Sequence	Accession	Sequence	Sequence	Fold Chang	P-value	Ratio	Intensity1	Intensity2
A_23_P21	NM_00196	Homo sapi	EGR1,TIS8	464.5436	0	464.5436	681.5943	321201.4
A_23_P10	NM_00525	v-fos FBJ r	FOS,c-fos	236.9381	0	236.9381	519.5927	122112
A_32_P87	NM_00058	interleukin	IL8,K60,NA	162.3819	0	162.3819	502.8458	83156.53
A_23_P34	NM_00402	Homo sapi	ATF3	158.9427	0	158.9427	1071.668	177636.8
A_23_P46	NM_00039	Homo sapi	EGR2,CMT	156.2562	0	156.2562	356.1077	56206.38
A_23_P20	NM_00222	Homo sapi	JUN,AP1	121.5397	0	121.5397	4182.786	510720.4
A_23_P38	NM_00196	Homo sapi	EGR4,NGF	104.9958	0	104.9958	1217.094	132433.8
A_23_P12	NM_00213	Homo sapi	NR4A1,HM	77.882	0	77.882	3539.875	280217.1
A_24_P33	NM_00402	activating t	ATF3	58.08273	3.53E-38	58.08273	41.68084	2393.278
A_24_P18	NM_00209	GRO3 onc	CXCL3,GR	47.99849	0	47.99849	476.8029	22994.58
A_24_P23	NM_01567	growth arre	GADD45B,	45.09765	0	45.09765	606.0646	27774.48
A_23_P31	NM_00208	Homo sapi	CXCL2,GR	41.35002	0	41.35002	4121.537	170681.5
A_23_P21	NM_00408	Homo sapi	DDIT3,CH	39.06985	0	39.06985	1133.613	44969.34
A_32_P74	BC052560	Hypothetic	LOC38776	37.30019	2.62E-43	37.30019	581.8859	22638.88
A_23_P15	NM_00490	Homo sapi	IER2,ETR1	36.48785	0	36.48785	13957.83	517561.8
A_23_P11	NM_00215	Homo sapi	HSPA6	32.43428	0	32.43428	741.1545	24432.38
A_32_P36	NM_00490	immediate	IER2,ETR1	30.90698	0	30.90698	22857.05	710043.6
A_23_P11	NM_00441	Homo sapi	DUSP1,HV	28.07072	0	28.07072	2456.328	70420.39
A_23_P71	NM_00151	Homo sapi	CXCL1,GR	27.80392	0	27.80392	5433.419	153650.5
A_23_P43	BC015544	Homo sapi	histone ger	27.75549	0	27.75549	313.1942	8793.668
A_23_P13	NM_00297	Homo sapi	SAT,DC21	26.02123	0	26.02123	2720.347	71749.4
A_23_P39	NM_00340	Homo sapi	ZFP36,TTF	25.94412	5.70E-34	25.94412	3028.552	81090.66
A_23_P21	NM_03141	Homo sapi	NFKBIZ,IK	25.46993	1.19E-35	25.46993	335.2256	9016.4
A_24_P77	NM_00096	prostaglan	PTGS2,CC	25.36269	0	25.36269	293.793	7606.039
A_24_P30	NM_00154	Homo sapi	IFIT2,G10F	24.56284	0	24.56284	106.4769	2627.928
A_23_P37	NM_00297	spermidine	SAT,DC21	22.1608	0	22.1608	1210.49	27265.21
A_23_P72	NM_00057	Homo sapi	IL1A,IL1,IL	19.51928	7.71E-10	19.51928	75.8701	1220.448
A_23_P39	NM_17320	Homo sapi	NR4A3,CH	18.70332	4.91E-32	18.70332	445.8285	8484.807
A_24_P25	NM_00096	prostaglan	PTGS2,CC	18.52013	0	18.52013	486.8897	9213.003
A_24_P25	NM_00208	GRO2 onc	CXCL2,GR	17.66887	0	17.66887	2174.97	39539.17
A_23_P70	NM_00423	Homo sapi	CD83,BL1	16.55241	0	16.55241	987.2628	16662.44
A_23_P92	NM_02504	Homo sapi	ARF7,FLJ2	16.29337	0	16.29337	237.8635	3947.584
A_23_P211468		Unknown	A_23_P21	15.96048	0	15.96048	388.6905	6245.389
A_23_P21	NM_00353	Homo sapi	HIST1H3D	15.7847	0	15.7847	282.6393	4513.019
A_24_P37	NM_00441	dual specif	DUSP2,PA	15.68921	0	15.68921	1395.635	21980.24
A_23_P36	NM_01519	activity-reg	ARC,KIAA	15.63	0	15.63	961.8306	15352.11
A_24_P683829		Unknown	THC18075	14.6666	0	14.6666	317.7976	4687.551
A_23_P24	NM_00154	Homo sapi	IFIT2,G10F	14.51711	0	14.51711	1572.804	23249.09
A_24_P28	NM_08065	Homo sapi	RSAD2,cig	14.25458	7.96E-17	14.25458	39.76581	530.5926
A_23_P11	NM_01627	Homo sapi	KLF2,LKLF	13.8936	0	13.8936	1200.643	16909.42
A_23_P13	NM_00618	nuclear rec	NR4A2,NC	13.01016	0	13.01016	164.7286	2166.61
A_23_P21	NM_00443	Homo sapi	EGR3,PILC	12.99403	8.31E-09	12.99403	95.04533	1007.735
A_24_P18	NM_00720	dual specif	DUSP10,M	12.74372	3.01E-10	12.74372	49.32599	514.5877
A_24_P93	BC015987	Homo sapi	BC015987	12.3526	1.77E-37	12.3526	124.5665	1488.519
A_23_P10	NM_02052	Homo sapi	NFKBIA,IK	12.26182	0	12.26182	11169.13	137027.3
A_24_P91	NM_01326	myosin reg	MYLIP,MIF	12.06917	0	12.06917	233.8587	2828.718
A_23_P52	NM_00154	Homo sapi	IFIT1,G10F	12.04725	0	12.04725	923.3189	11136.3
A_32_P99804		Unknown	THC18224	12.02354	0	12.02354	1204.014	15147.22
A_32_P97	BC024020	Transmem	VMP1	12.01745	0	12.01745	607.1211	7344.472
A_23_P42	NM_00673	Homo sapi	FOSB,G0S	11.93662	0	11.93662	824.1993	10043.69
A_23_P10	NM_00441	Homo sapi	DUSP2,PA	11.93552	0	11.93552	8969.526	107383.1

A_24_P93	NM_00552	hairy (Dros	HES1,HHL	11.35458	0	11.35458	2160.043	25123.87
A_24_P710681		Unknown	THC19808	11.34807	9.58E-43	11.34807	253.5288	2887.107
A_23_P42	NM_00007	Homo sapi	CDKN1C,E	11.21564	0	11.21564	549.5429	6158.189
A_23_P54	NM_01468	Homo sapi	HERPUD1	11.11378	0	11.11378	18124.81	202438.1
A_23_P15	NM_00441	dual specif	DUSP5,DU	11.03976	0	11.03976	1063.02	11731.05
A_23_P63	NM_00130	Homo sapi	KLF6,GBF,	10.96973	0	10.96973	924.2804	10225.22
A_23_P32	NM_17351	Homo sapi	PNLDC1,F	10.78517	1.16E-14	10.78517	96.5481	906.3457
A_23_P42	NM_00389	Homo sapi	IER3,DIF2,	10.67289	0	10.67289	5886.044	62958.27
A_23_P48	NM_00222	Homo sapi	JUNB	10.29243	0	10.29243	10670.19	112626.5
A_23_P35	NM_00354	Homo sapi	HIST1H4F,	10.23536	0	10.23536	1020.196	10567.66
A_24_P26	NM_00416	Ras-relate	RRAD,RAD	10.1569	0	10.1569	191.1504	1920.825
A_23_P88	NM_00416	Homo sapi	RRAD,RAD	9.97642	0	9.97642	1210.104	12130.42
A_24_P31	NM_02196	myeloid ce	MCL1,TM,I	9.75612	0	9.75612	524.7914	5140.129
A_23_P90	NM_00614	Homo sapi	DNAJB1,H	9.64393	0	9.64393	6298.114	60879.16
A_23_P46	NM_01894	Gene 33\/M	ERRFI1,MI	9.35995	0	9.35995	1676.655	15720.33
A_24_P917810		Unknown	A_24_P91	9.13533	2.85E-07	9.13533	39.37141	316.4672
A_24_P174353		Unknown	A_24_P17	9.08989	1.23E-09	9.08989	44.45643	374.2371
A_23_P43	NM_00354	Homo sapi	HIST2H4,H	8.92372	0	8.92372	2456.528	22205.75
A_23_P12	NM_00112	Homo sapi	ADM,AM	8.91206	0	8.91206	4673.381	42253
A_23_P32	NM_00354	Homo sapi	HIST1H4H	8.86125	0	8.86125	1111.486	9980.064
A_23_P17	NM_00459	small induc	CCL20,CK	8.82099	1.62E-24	8.82099	285.1448	2509.052
A_23_P251002		Unknown	A_23_P25	8.81629	0	8.81629	445.4096	3977.39
A_23_P20	NM_00395	Homo sapi	SOCS3,CIS	8.50512	0	8.50512	4079.549	34649.61
A_23_P17	NM_00552	Homo sapi	HES1,HHL	8.45728	0	8.45728	1707.213	14668.29
A_23_P21	NM_01337	CDK4-bind	SERTAD1,	8.3825	0	8.3825	3397.287	28587.31
A_32_P103776		Unknown	A_32_P10	8.32713	0	8.32713	2459.157	20793.13
A_24_P14	U87836	Splicing fa	SFRS10	8.12896	0	8.12896	2709.605	22123.79
A_23_P15	NM_13931	Homo sapi	ANGPTL4,	8.109	5.48E-07	8.109	94.84409	619.7779
A_24_P68	NM_00534	Heat shock	HSPA1A	8.10726	0	8.10726	5668.797	46085.52
A_32_P15	AK092260	CDNA FLJ	DUSP5P,F	7.8167	1.09E-40	7.8167	1563.199	12584.53
A_24_P20	NM_00349	H4 histone	HIST1H4I,I	7.81527	0	7.81527	953.3478	7461.55
A_24_P11	NM_02407	hypothetica	SECISBP2	7.80134	0	7.80134	3629.11	28561.64
A_23_P31	NM_01326	myosin reg	MYLIP,MIF	7.7159	7.89E-31	7.7159	143.7228	1076.878
A_23_P12	NM_01860	DnaJ (Hsp	DNAJA4,M	7.70861	3.62E-15	7.70861	130.3933	903.8911
A_23_P98	NM_00116	Homo sapi	BIRC3,AIP	7.64587	8.37E-25	7.64587	125.6679	915.6813
A_24_P92	BC020539	KIAA1961	KIAA1961	7.46167	0	7.46167	417.1499	3096.572
A_23_P74	AF277181	PNAS-123	LOC85028	7.35637	0	7.35637	7695.625	57009.2
A_32_P22	NM_00351	H2A histon	HIST1H2A	7.28088	0	7.28088	656.6816	4778.458
A_24_P25	NM_00209	GRO3 onc	CXCL3,GR	6.98948	3.44E-42	6.98948	6305.701	45033.62
A_24_P34	NM_01608	ras-related	RASD1,AG	6.96353	0.00001	6.96353	67.07541	368.6379
A_32_P181548		C40201 art	THC19339	6.77726	0.00009	6.77726	44.61544	252.0516
A_24_P74	AK022379	Beta-2-mic	B2M	6.6557	1.95E-22	6.6557	183.7753	1163.667
A_32_P26	AV756170	Transcribe	USP9Y,DF	6.5602	0.00001	6.5602	262.8691	1589.609
A_23_P13	NM_02030	cyclin L ani	CCNL1,BM	6.41669	0	6.41669	1531.347	9897.773
A_23_P93	NM_00353	Homo sapi	HIST1H3B	6.41595	0	6.41595	1967.36	12816.81
A_23_P35	NM_00154	interferon-i	IFIT3,IRG2	6.40724	1.26E-44	6.40724	323.374	2056.807
A_24_P12	NM_00534	Heat shock	HSPA1A	6.33146	0	6.33146	11397.93	73137.46
A_24_P33	NM_02030	cyclin L ani	CCNL1,BM	6.00312	0	6.00312	1181.457	7121.404
A_24_P39	NM_00703	DnaJ (Hsp	DNAJB4,D	5.99426	0	5.99426	1323.94	7982.229
A_23_P51	NM_00703	Homo sapi	DNAJB4,D	5.99225	0	5.99225	822.4202	4929.594
A_23_P13	NM_00194	Homo sapi	DUSP6,MK	5.92358	1.55E-39	5.92358	695.0916	4160.371
A_23_P13	BC000297	3-hydroxy-	HMGCS1	5.90343	0	5.90343	429.7402	2525.849

A_32_P213948	BC003099	THC18819	5.79633	8.28E-29	5.79633	244.1073	1376.902
A_23_P13	NM_00373	Homo sapi	OASL,TRIF	5.75634	2.26E-34	5.75634	4023.981	23565.55
A_24_P23	NM_18172	Ankyrin rep	Lrp2bp	5.74048	2.66E-44	5.74048	466.5822	2678.075
A_32_P201976	Unknown	A_32_P20	5.72029	2.41E-14	5.72029	94.19211	522.7108
A_24_P69	NM_00130	core promo	KLF6,GBF	5.64819	2.13E-39	5.64819	397.5803	2211.654
A_32_P16	AL833005	MRNA; cDI	AL833005	5.5553	0	5.5553	1124.037	6287.571
A_23_P11	NM_01608	Homo sapi	RASD1,AG	5.48457	1.40E-21	5.48457	3354.916	19384.26
A_23_P43	NM_01826	hypothetica	H2AFJ,MG	5.37903	1.77E-10	5.37903	131.204	621.0766
A_23_P81	NM_00510	interferon-s	G1P2,UCR	5.37583	0	5.37583	6876.134	36969.75
A_23_P47	NM_00428	Homo sapi	BAG3,BIS,	5.28842	0	5.28842	1783.822	9528.66
A_23_P65	NM_00552	hairy (Dros	HES1,HHL	5.28658	1.62E-06	5.28658	60.44539	304.8661
A_23_P12	NM_02499	Homo sapi	ATF7IP2,M	5.26996	2.81E-26	5.26996	217.2426	1107.095
A_23_P23	NM_00192	Homo sapi	GADD45A,	5.26836	0	5.26836	3274.258	17508.13
A_32_P123743	U12390 be	THC19144	5.21996	0	5.21996	4336.169	22912.82
A_32_P30898	Unknown	A_32_P30	5.19913	2.08E-32	5.19913	541.5852	2812.856
A_23_P32	NM_02507	MCP-1 trea	FLJ23231	5.19566	6.88E-25	5.19566	1869.29	9741.75
A_23_P88	NM_00077	cytochrome	CYP3A5,C	5.12382	0.00008	5.12382	60.00282	267.4317
A_23_P86	NM_01654	Homo sapi	IER5,SBBI	5.09868	0	5.09868	20863.89	106768.5
A_23_P52	NM_00428	BCL2-asso	BAG3,BIS,	5.06273	0	5.06273	9770.443	50043.71
A_23_P20	NM_02112	Homo sapi	PMAIP1,AI	5.05238	0	5.05238	1524.47	7715.395
A_24_P15	NM_00629	tumor necr	TNFAIP3,A	4.97659	7.24E-42	4.97659	1199.285	6036.177
A_32_P16	AL833005	MRNA; cDI	AL833005	4.95211	0	4.95211	983.0056	4913.746
A_23_P16	AK128645	Syndecan I	SDCBP	4.9444	4.84E-19	4.9444	228.8995	1078.874
A_23_P32	NM_00543	Homo sapi	FOSL1,FR	4.8947	2.16E-08	4.8947	94.49641	422.1796
A_23_P59	NM_02105	Homo sapi	HIST1H2A	4.88995	0	4.88995	3700.308	18142.4
A_32_P13	BE937815	Transcribe	BE937815	4.87473	2.34E-15	4.87473	379.1601	1917.574
A_23_P70	NM_00552	Heat shock	HSPA1L	4.76158	0	4.76158	2412.558	11500.48
A_24_P17	NR_00145	Homo sapi	LOC85391	4.75976	3.26E-22	4.75976	159.525	747.7806
A_23_P73096	Unknown	A_23_P73	4.69189	1.67E-28	4.69189	486.6971	2269.769
A_23_P88	NM_00664	Homo sapi	HSPH1,HS	4.69029	0	4.69029	20894.13	99585.45
A_32_P174121	Unknown	A_32_P17	4.65316	7.37E-14	4.65316	1511.273	7349.998
A_23_P16	NM_00486	Homo sapi	GDF15,PD	4.62501	0	4.62501	23876.68	110433.2
A_24_P40	NM_02090	pleckstrin h	PLEKHA4,	4.62043	3.32E-07	4.62043	101.4837	414.5394
A_32_P208654	A47582 B-c	THC19033	4.61204	0.00016	4.61204	78.56785	299.2471
A_23_P32	NM_00538	Homo sapi	NFIL3,E4B	4.60154	2.94E-43	4.60154	1412.05	6548.019
A_24_P17	BC066345	AT rich inte	ARID5B	4.57986	1.84E-28	4.57986	299.1644	1343.189
A_23_P21	NM_00183	clusterin (c	CLU,CLI,A	4.56806	0	4.56806	2070.37	9465.635
A_23_P76	BC018929	Pleckstrin h	PHLDA1	4.47372	0	4.47372	4132.374	18491.22
A_23_P71	NM_00060	Homo sapi	IL6,HGF,H	4.46971	2.82E-33	4.46971	1760.164	7970.87
A_23_P35	NM_02509	KIAA1453	USP36,DU	4.39496	1.24E-14	4.39496	521.1262	2290.327
A_32_P10403	ALU1_HUM	THC18991	4.35158	1.81E-43	4.35158	503.3888	2156.017
A_24_P32	NM_00442	dual specif	DUSP8,HB	4.34092	1.09E-22	4.34092	362.8618	1530.577
A_23_P47	NM_00331	Homo sapi	PHLDA2,IF	4.32227	1.16E-37	4.32227	17631.58	77872.91
A_32_P105940	Unknown	A_32_P10	4.30642	3.47E-35	4.30642	595.0411	2548.217
A_23_P90	NM_01433	Homo sapi	PPP1R15A	4.3022	4.41E-29	4.3022	1246.641	5433.411
A_23_P30	NM_00455	Homo sapi	NFKBIE,IK	4.28163	3.11E-39	4.28163	815.6448	3503.521
A_23_P51	NM_00720	dual specif	DUSP10,M	4.277	2.35E-11	4.277	208.1509	814.4408
A_23_P93	NM_00344	Homo sapi	ZNF165,LD	4.24399	5.13E-28	4.24399	390.5721	1633.525
A_23_P32	AK002200	SMC4 stru	SMC4L1	4.14953	5.71E-22	4.14953	230.231	979.4745
A_24_P39	NM_19821	Homo sapi	ING1,p33,p	4.10393	0	4.10393	720.5878	2953.288
A_23_P89	NM_00261	Homo sapi	PER1,PER	4.05971	8.18E-41	4.05971	504.3368	2029.146
A_23_P12	NM_00531	H1 histone	HIST1H1C	4.05843	0	4.05843	12734.23	51878.46

A_32_P10	AI744472	BTG family	BTG2	4.05519	9.16E-13	4.05519	163.6628	629.2361
A_23_P11	NM_00534	Heat shock	HSPA1A	4.04691	0	4.04691	172800.1	702079
A_23_P29	NM_00531	Homo sapi	H1F0,H10,	4.03901	1.39E-14	4.03901	427.5831	1684.946
A_24_P50	AK056455	Poliovirus r	PVRL1	4.03001	1.53E-42	4.03001	4918.849	19922.48
A_23_P32	NM_13839	hypothetica	LOC93082	4.02581	9.23E-20	4.02581	436.3677	1714.294
A_32_P171488		Unknown	A_32_P17	3.97911	3.96E-08	3.97911	262.3674	938.3945
A_24_P38	NM_18259	hypothetica	FLJ39575,	3.96497	2.88E-40	3.96497	570.5965	2252.661
A_24_P33	NM_02196	myeloid ce	MCL1,TM,I	3.96193	0	3.96193	1329.266	5276.206
A_23_P12	NM_00285	Homo sapi	PTX3,TSG	3.94303	7.40E-12	3.94303	2192.208	8841.14
A_23_P11	NM_03235	hypothetica	MGC1074	3.93231	2.58E-20	3.93231	283.534	1082.943
A_24_P35	NM_19826	similar to s	FLJ11021	3.92778	0	3.92778	1493.178	5872.436
A_23_P42	NM_02106	Homo sapi	HIST1H2A	3.92148	0	3.92148	10754.99	42359.07
A_23_P10	NM_00351	Homo sapi	HIST2H2A	3.9183	0	3.9183	15827.85	62558.7
A_23_P36	BC001694	Ubiquitin-c	UBE2B	3.90416	1.38E-06	3.90416	82.30389	321.7947
A_23_P32	NM_00220	Homo sapi	ISG20,HEN	3.88852	7.83E-27	3.88852	3026.019	12047
A_24_P11	NM_01894	Gene 33/M	ERRFI1,MI	3.87884	0.00057	3.87884	64.50916	235.6653
A_24_P33	NM_02196	myeloid ce	MCL1,TM,I	3.8675	9.76E-38	3.8675	27840.7	110030.7
A_23_P170713		Unknown	A_23_P17	3.83188	0	3.83188	6832.036	26458.34
A_23_P21	NM_00195	Homo sapi	EDN1,ET1	3.80939	9.58E-26	3.80939	440.796	1654.783
A_32_P173177		AF537132	THC19115	3.79704	8.41E-12	3.79704	173.7941	624.3632
A_23_P51	NM_00407	Homo sapi	PLK3,CNK	3.7949	0	3.7949	1512.402	5751.701
A_32_P100365		Unknown	A_32_P10	3.79201	4.14E-16	3.79201	269.4356	972.6516
A_24_P11	NM_00052	low density	LDLR,FH,F	3.77968	6.99E-34	3.77968	5639.842	21622.21
A_23_P14	NM_03344	Homo sapi	HIST3H2A	3.77963	0	3.77963	5965.465	22656.26
A_23_P52	NM_01234	Homo sapi	BAMBI,NM	3.77844	0	3.77844	2045.98	7778.202
A_23_P21	NM_00607	Homo sapi	CITED2,MI	3.74387	0	3.74387	569.6959	2126.059
A_23_P25	NM_00526	Homo sapi	GEM,KIR,N	3.70127	1.32E-14	3.70127	412.2376	1473.107
A_32_P124728		Unknown	THC18931	3.69451	2.00E-19	3.69451	293.9509	1055.086
A_24_P36	AK092260	CDNA FLJ	DUSP5P,F	3.68614	2.51E-21	3.68614	1861.277	7008.959
A_23_P36	NM_00535	Homo sapi	JUND	3.65094	3.12E-31	3.65094	22075.72	81776.47
A_24_P26	NM_02100	Homo sapi	UBC,HMG	3.63293	0	3.63293	198451.2	720713.2
A_24_P32	NM_00702	decidual pr	C10orf10,F	3.62264	1.90E-08	3.62264	129.756	447.303
A_24_P10	NM_02281	microtubule	MAP1LC3E	3.61827	0	3.61827	1011.444	3666.238
A_32_P78613		Unknown	A_32_P78	3.61532	3.33E-08	3.61532	131.0064	513.0694
A_24_P29	AK124426	CDNA clon	AK124426	3.60343	8.18E-24	3.60343	243.4294	876.3939
A_24_P91	NM_01644	hypothetica	LOC51233	3.59844	2.33E-39	3.59844	2006.346	7236.761
A_23_P32	NM_02100	Homo sapi	UBC,HMG	3.58796	0	3.58796	202416.1	727933.1
A_23_P33	NM_00735	pleckstrin h	PHLDA1,P	3.58672	5.87E-36	3.58672	566.5662	2013.213
A_23_P16	NM_02106	H2A histon	HIST1H2A	3.57399	0	3.57399	4990.316	17860.39
A_24_P68	NM_17506	Homo sapi	HIST2H2A	3.56394	0	3.56394	7155.11	25680.14
A_24_P932822		B40201 art	THC19032	3.56055	9.04E-06	3.56055	106.5113	422.3822
A_23_P11	NM_00292	Homo sapi	RGS2,G0S	3.55997	0	3.55997	7149.135	25446.68
A_24_P18	NM_00534	heat shock	HSPA5,BIF	3.55538	0	3.55538	4663.342	16542.22
A_23_P76	NM_00123	serine (or c	SERPINH1	3.55342	0	3.55342	1379.782	4902.614
A_23_P16	BC017379	Homo sapi	HIST1H2A	3.53178	0	3.53178	995.6473	3514.075
A_23_P16	NM_00351	H2B histon	HIST1H2B	3.52561	0	3.52561	1388.655	4909.695
A_23_P30	NM_02196	Homo sapi	HIST1H4J,	3.5181	1.86E-30	3.5181	3185.136	11305.83
A_24_P86	NM_00351	H2A histon	HIST1H2A	3.51161	0	3.51161	6439.935	22636.46
A_23_P30	NM_00351	Homo sapi	HIST2H2A	3.5108	3.99E-38	3.5108	1865.342	6635.011
A_24_P12	NM_00670	growth arre	GADD45G	3.48909	0.00009	3.48909	598.8401	1935.978
A_23_P71	NM_00182	Homo sapi	CKS2,CKS	3.48699	2.24E-33	3.48699	52657.91	187142.9
A_24_P33	NM_01763	hypothetica	FLJ20035,	3.46931	4.63E-14	3.46931	229.1555	762.093

A_24_P26	NM_02411	hypothetica	MGC5306	3.46871	1.03E-38	3.46871	3170.913	11189.05
A_24_P54	BC068044	Histone 2,	HIST2H2A	3.46474	0	3.46474	3330.916	11557.55
A_24_P39	NM_02106	H2A histon	HIST1H2A	3.46119	3.24E-38	3.46119	2339.672	8205.212
A_23_P37	NM_02012	Homo sapi	TUFT1	3.45972	1.44E-21	3.45972	1752.037	6131.297
A_24_P14	NM_02106	H2B histon	HIST1H2B	3.44611	0	3.44611	1385.774	4780.402
A_23_P16	NM_02053	Homo sapi	OSM,MGC	3.44165	2.04E-34	3.44165	876.1311	3014.212
A_23_P25	NM_00165	Homo sapi	AREG,AR,	3.43585	4.61E-19	3.43585	922.7894	3181.108
A_24_P87	NM_00351	Homo sapi	HIST2H2A	3.414	0	3.414	8894.885	30441.83
A_23_P22	NM_00554	Homo sapi	INSIG1,CL	3.36977	0	3.36977	1699.317	5736.158
A_24_P192586		DJA1_HUM	THC19000	3.36574	0	3.36574	7911.749	26702.16
A_23_P42	NM_00353	Homo sapi	HIST1H3G	3.36391	0	3.36391	3784.704	12771.71
A_24_P15	NM_00352	H2B histon	HIST2H2B	3.35777	0	3.35777	2628.79	8846.459
A_24_P55	NM_02105	H2B histon	HIST1H2B	3.35037	0	3.35037	1213.889	4067.523
A_24_P15	NM_00329	tumor rejec	TRA1,ECG	3.33317	0	3.33317	8986.49	29848.95
A_23_P93	NM_00352	Homo sapi	HIST1H2B	3.32759	0	3.32759	1423.247	4738.24
A_24_P516728		Unknown	A_24_P51	3.32722	0	3.32722	10520.67	34923.7
A_24_P29	U60873	Trophoblas	TncRNA	3.30248	1.10E-13	3.30248	293.1792	924.9707
A_24_P29	AF187554	Homo sapi	sperm anti	3.29624	8.34E-12	3.29624	9221.86	31032.49
A_24_P12	NM_18178	Homo sapi	NCOA7,ES	3.289	9.14E-43	3.289	3436.824	11413.98
A_24_P545200		Unknown	A_24_P54	3.27953	0	3.27953	17861.2	58639.24
A_32_P23	AL110141	Growth arr	GAS5	3.27829	0	3.27829	16639.94	54306.2
A_24_P33	NM_01909	hypothetica	C1orf183,F	3.27673	1.11E-08	3.27673	5278.425	17243.38
A_24_P17	NM_00353	H3 histone	HIST1H3B	3.27665	0	3.27665	13024.2	42654.05
A_23_P41	NM_00143	Homo sapi	EREG,ER	3.2682	1.58E-13	3.2682	260.3858	816.6914
A_32_P16	NM_03326	winged hel	FOXQ1,HF	3.2618	3.85E-26	3.2618	1421.075	4703.94
A_23_P40	NM_00352	Homo sapi	HIST1H2B	3.24768	0	3.24768	3542.222	11544.06
A_23_P96	NM_00602	H1 histone	H1FX,H1X	3.23441	0	3.23441	1104.588	3578.552
A_24_P16361		Unknown	A_24_P16	3.23327	3.60E-10	3.23327	256.8799	774.0521
A_24_P401090		Unknown	ENST0000	3.23305	0	3.23305	1342.082	4340.282
A_23_P10	NM_00443	Homo sapi	ELF3,ERT,	3.21992	2.28E-12	3.21992	206.1873	685.9478
A_24_P24	NM_00222	jun B proto	JUNB	3.21603	5.35E-11	3.21603	1035.444	3235.788
A_24_P67552		Unknown	A_24_P67	3.20673	0	3.20673	1695.671	5462.837
A_24_P82	NM_00329	tropomyosi	TPM4	3.20634	3.15E-26	3.20634	2002.269	6505.968
A_24_P21	NM_00351	H2A histon	HIST1H2A	3.20058	0	3.20058	8368.483	26812.78
A_23_P34	NM_00525	Homo sapi	FOSL2,FR	3.19485	9.87E-06	3.19485	173.2246	484.0889
A_24_P37	BC036402	MAX dimer	MAD	3.19338	6.23E-22	3.19338	561.0252	1771.314
A_23_P55	NM_00650	Homo sapi	RELB,I-RE	3.18124	0	3.18124	1700.503	5407.974
A_23_P13	NM_00112	Homo sapi	ADFP,ADF	3.18051	6.42E-34	3.18051	417.2078	1323.655
A_24_P53	AK057151	CAMP resp	CREB5	3.17163	3.68E-15	3.17163	378.9346	1162.745
A_23_P86	NM_15278	hypothetica	MGC40405	3.15518	1.68E-43	3.15518	1802.2	5713.99
A_23_P13	NM_00598	Homo sapi	SNAI1,SNA	3.15339	0	3.15339	1879.185	5940.421
A_23_P25	NM_00442	Homo sapi	EFNA1,B6	3.15179	1.37E-41	3.15179	3195.942	10120.89
A_23_P36	NM_02123	selenoprot	SELK,HSP	3.14344	2.80E-45	3.14344	8090.519	25549.31
A_24_P37	NM_00352	H2B histon	HIST1H2B	3.14023	0	3.14023	3893.729	12269.85
A_23_P11	NM_01632	cyclin-E bir	HERC5,CE	3.12862	1.43E-06	3.12862	192.4828	533.7332
A_23_P30	NM_00352	Homo sapi	HIST1H2B	3.1252	0	3.1252	5237.894	16333.17
A_23_P11	NM_00352	H2B histon	HIST1H2B	3.12115	0	3.12115	3920.864	12334.04
A_23_P46	NM_00155	cysteine-ric	CYR61,CC	3.09704	7.64E-13	3.09704	243.4703	726.3488
A_23_P33	NM_17505	Homo sapi	HIST3H2B	3.09589	0	3.09589	2546.984	7888.862
A_23_P20	NM_00052	low density	LDLR,FH,F	3.09548	6.18E-12	3.09548	316.6636	931.0571
A_23_P39	AL833005	MRNA; cDI	AL833005	3.09408	0	3.09408	3288.664	10196.88
A_24_P12	NM_20703	Homo sapi	C1orf63,NF	3.08464	4.01E-06	3.08464	160.1499	446.8889

A_24_P88	NM_00346	secretogra	SCG2,CHC	3.07574	0.00077	3.07574	99.2647	275.9213
A_23_P42	NM_00352	Homo sapi	HIST1H2B	3.0621	0	3.0621	7460.503	22810.65
A_23_P62	NM_00676	Homo sapi	BTG2,PC3	3.05974	1.71E-20	3.05974	4866.411	15189.14
A_24_P11	NM_01223	RING1 and	RYBP,AAF	3.05921	1.83E-36	3.05921	1105.554	3399.037
A_24_P45	NM_02105	Homo sapi	HIST2H3C	3.04345	0	3.04345	5742.793	17545.38
A_24_P95	AK129833	Tissue fact	TFPI2	3.03904	8.05E-25	3.03904	465.6697	1396.137
A_23_P13	NM_00063	Homo sapi	SOD2,IPO	3.03706	0	3.03706	7147.02	21702.28
A_23_P11	NM_02105	Homo sapi	HIST2H3C	3.02602	0	3.02602	14151.2	42780.25
A_24_P27	NM_00354	H4 histone	HIST1H4G	3.01453	7.47E-08	3.01453	184.4369	521.2712
A_23_P14	NM_00216	Homo sapi	ID2,ID2A,II	3.0084	2.20E-18	3.0084	3078.774	9479.532
A_24_P98	NM_00534	heat shock	HSPA5,BIF	3.00798	1.55E-37	3.00798	2025.276	6123.201
A_23_P33	NM_18249	hypothetica	ZFAND2A	3.00654	0	3.00654	2210.512	6647.19
A_24_P75948		Unknown	A_24_P75	3.00552	0	3.00552	5388.696	16196.75
A_23_P67	NM_00381	Homo sapi	TNFSF9,4-	3.00403	2.91E-10	3.00403	925.3169	2666.276
A_23_P40	NM_01744	Homo sapi	H2BFS	3.0026	2.28E-35	3.0026	989.1964	2974.625
A_32_P20	BC038747	Transmem	TMAP1	2.99809	0.00001	2.99809	150.233	539.2077
A_32_P73143		ALU7_HUM	THC19625	2.99719	6.12E-09	2.99719	200.2556	569.0714
A_23_P81	NM_08059	Homo sapi	HIST1H2A	2.98903	0	2.98903	8230.31	24662.89
A_24_P923654		Unknown	A_24_P92	2.97835	0.00003	2.97835	151.039	405.8309
A_24_P13	NM_00688	CD86 antig	CD86,B70,	2.96671	5.62E-23	2.96671	835.4627	2475.417
A_23_P11	NM_02106	Homo sapi	HIST1H2B	2.96228	0	2.96228	2444.326	7252.578
A_24_P324465		H3 histone	HIST2H3C	2.94984	0	2.94984	22476.02	66540.19
A_24_P27	NM_02080	Homo sapi	ARRDC3,K	2.94844	1.40E-22	2.94844	716.7957	2103.706
A_24_P63	AK095492	3-hydroxy-	HMGCS1	2.94812	0	2.94812	1726.249	5091.665
A_23_P11	NM_00244	Homo sapi	MSX1,HO	2.94626	1.40E-27	2.94626	486.2615	1417.167
A_23_P35	NM_03145	sestrin 2	SESN2,HIS	2.93905	4.21E-16	2.93905	549.2378	1566.587
A_24_P24	NM_15278	hypothetica	MGC40405	2.93197	0	2.93197	946.0016	2774.243
A_23_P12	NM_00196	Homo sapi	EIF4A2,DD	2.93158	7.39E-41	2.93158	12853.3	38059.55
A_23_P21	NM_00292	Homo sapi	RGS16,RG	2.93149	4.43E-42	2.93149	1295.434	3801.048
A_23_P30	NM_00351	Homo sapi	HIST2H2A	2.92724	3.64E-21	2.92724	1291.334	3796.332
A_32_P91773		Unknown	A_32_P91	2.92442	3.74E-14	2.92442	384.6516	1167.049
A_24_P96	NM_00153	DnaJ (Hsp	DNAJA1,D	2.91956	3.78E-41	2.91956	19108.3	55497.51
A_32_P60687		Unknown	A_32_P60	2.91918	5.67E-07	2.91918	1804.208	5532.886
A_24_P16	NM_00108	cytochrome	CYP4F2,C	2.91429	1.91E-11	2.91429	604.616	1733.942
A_32_P67680		ENPL_HUM	THC18153	2.91261	1.37E-16	2.91261	769.9921	2296.492
A_24_P916853		Unknown	A_24_P91	2.90581	2.13E-22	2.90581	830.4506	2401.554
A_23_P21	NM_00246	Homo sapi	MYC,c-My	2.89768	4.16E-33	2.89768	30685.67	89845.36
A_24_P50509		Unknown	A_24_P50	2.89031	0	2.89031	5417.199	15715.09
A_23_P40	NM_00235	Homo sapi	MXD1,MAD	2.87746	3.58E-09	2.87746	298.8021	803.9074
A_23_P51	NM_00404	Homo sapi	RHOB,ARH	2.85507	8.67E-25	2.85507	811.1011	2296.613
A_23_P80	NM_00351	Homo sapi	HIST1H2B	2.85482	0	2.85482	8802.009	25128.91
A_32_P450879		Unknown	THC19005	2.84893	0	2.84893	846.9572	2410.496
A_23_P94	BC007651	Homo sapi	MGC16037	2.8392	4.68E-34	2.8392	1002.644	2852.812
A_24_P67	BM689022	HLAT1-3TM	IMAA	2.82033	9.28E-15	2.82033	574.9774	1580.935
A_23_P14	NM_00329	tropomyosi	TPM4	2.81883	0	2.81883	2918.259	8245.037
A_24_P30	NM_02106	H2A histon	HIST1H2A	2.80511	0	2.80511	6717.484	18874.65
A_32_P222571		Unknown	THC18595	2.80152	2.49E-23	2.80152	4466.208	12677.29
A_24_P832831		AF247142	THC19660	2.79935	0.00034	2.79935	140.5665	467.1533
A_23_P13	NM_17335	Homo sapi	SNF1LK,M	2.79832	1.26E-12	2.79832	534.0779	1456.565
A_32_P41267		Unknown	THC18482	2.79269	2.03E-07	2.79269	242.5175	625.5042
A_24_P358305		Unknown	A_24_P358	2.79102	1.19E-16	2.79102	918.7808	2561.404
A_32_P52	AK026328	Hypothetic	FLJ22675	2.77735	0	2.77735	2646.721	7352.077

A_32_P75	BG400725	60246411C	RNU2,U2	2.76915	7.91E-16	2.76915	1345.832	3677.362
A_32_P38	BC039555	CDNA clon	BC039555	2.76317	0	2.76317	26572.78	73461.55
A_23_P25	NM_01959	Homo sapi	LRRC8A,L	2.75345	1.83E-36	2.75345	5885.955	16354.36
A_24_P93	NM_00353	H3 histone	HIST1H3I,I	2.75126	7.01E-45	2.75126	4035.33	11167.07
A_23_P60	NM_00153	Homo sapi	DNAJA1,D	2.74801	0	2.74801	31173.8	85542.68
A_32_P10	BQ013066	Vestigial lik	VGLL4	2.74078	4.31E-16	2.74078	454.2018	1216.368
A_23_P15	NM_00298	small induc	CCL5,SISc	2.74029	4.90E-35	2.74029	571.8109	1560.615
A_23_P13	NM_00731	Homo sapi	ABL2,ARG	2.73405	3.28E-32	2.73405	528.9199	1449.23
A_24_P16	BC032480	Similar to F	HSUP1	2.73111	7.29E-36	2.73111	49532.2	136638.3
A_23_P25	NM_00532	Homo sapi	HIST1H1B	2.72526	2.07E-07	2.72526	159.4358	439.8021
A_24_P12	NM_18178	Homo sapi	NCOA7,ES	2.72352	3.08E-25	2.72352	1022.803	2790.18
A_23_P59	NM_00352	Homo sapi	HIST1H2B	2.71255	0	2.71255	6594.347	17915.23
A_23_P16	NM_01651	Pipecolic a	PIPOX	2.70563	0.00021	2.70563	2596.835	6353.631
A_32_P69	NM_00216	inhibitor of	ID2,ID2A,II	2.70416	7.79E-35	2.70416	2855.272	7781.099
A_24_P56	AK027191	Trophoblas	TncRNA	2.6867	0	2.6867	2571.429	6924.199
A_23_P10	NM_00688	Homo sapi	ZFP36L2,E	2.68597	8.19E-29	2.68597	4447.44	12105.12
A_23_P51	NM_02043	Homo sapi	GJA12,Cx4	2.67978	0	2.67978	2843.049	7611.169
A_23_P36	NM_00352	Homo sapi	HIST1H2B	2.66952	0	2.66952	7945.025	21219.13
A_23_P16	NM_00541	Homo sapi	SLC20A1,F	2.66806	6.17E-44	2.66806	18756.77	50297.04
A_23_P34	NM_17506	Homo sapi	HIST2H2A	2.66802	5.59E-16	2.66802	379.7031	1032.497
A_24_P25	NM_02519	phosphopr	TRIB1,C8F	2.66577	6.75E-32	2.66577	3707.324	9954.867
A_24_P92	BG182941	Transcribe	BG182941	2.66415	4.40E-36	2.66415	6720.743	17958.71
A_24_P38	AK074675	Protein kin	PRKAG2	2.66388	0.00001	2.66388	135.7487	365.9009
A_23_P147404		Unknown	A_23_P14	2.65236	0	2.65236	3396.674	8993.57
A_23_P53	NM_02301	similar to s	FLJ11021	2.65184	2.55E-33	2.65184	9889.369	26320.55
A_32_P12	BC066906	Similar to H	LOC44009	2.65165	0	2.65165	67010.03	177538
A_32_P80016		Unknown	A_32_P80	2.64739	6.83E-09	2.64739	370.5994	925.6787
A_23_P13	NM_00353	Homo sapi	HIST1H3C	2.64364	0	2.64364	25073.49	66226.08
A_24_P35	BC007651	Homo sapi	MGC16037	2.62318	1.14E-33	2.62318	1306.329	3435.503
A_24_P82	NM_00532	H3 histone	H3F3B,H3.	2.62232	0	2.62232	35720.41	93387.95
A_32_P142802		Unknown	THC19486	2.61882	2.05E-13	2.61882	4213.442	11413.02
A_32_P10	AK129982	Homo sapi	AK129982	2.61702	5.01E-13	2.61702	412.4269	1046.246
A_24_P17	XM_29327	Homo sapi	XM_29327	2.6125	1.29E-40	2.6125	4277.929	11231.04
A_23_P35	NM_01959	hypothetica	KIAA1434,	2.60951	2.06E-16	2.60951	4064.518	10787.51
A_23_P40	AF258593	Hypothetic	LOC20117	2.60855	7.78E-19	2.60855	8923.285	23334.69
A_23_P14	NM_01550	TCDD-indu	TIPARP	2.6078	2.87E-17	2.6078	2295.52	6072.758
A_23_P34	NM_02031	Homo sapi	C1orf63,NF	2.60688	7.21E-12	2.60688	1016.961	2645.128
A_24_P34	NM_00244	msh (Dros	MSX1,HO	2.60512	3.18E-21	2.60512	4291.423	11338.91
A_32_P65472		U12390 be	THC18924	2.60053	1.25E-08	2.60053	272.4087	672.3582
A_23_P30	NM_00354	Homo sapi	HIST1H4K	2.59759	3.05E-24	2.59759	3131.044	8239.689
A_23_P71	NM_18159	Homo sapi	UPP1,UP,U	2.57801	2.15E-07	2.57801	373.8755	896.9851
A_23_P38	NM_01441	Homo sapi	BBC3,JFY	2.57736	0	2.57736	2513.459	6478.322
A_24_P73	AL832183	Hypothetic	LOC28445	2.57723	1.34E-29	2.57723	1637.567	4218.483
A_23_P36	NM_00351	H2A histon	HIST1H2A	2.57658	5.86E-14	2.57658	287.1195	730.4623
A_23_P54	BC051317	Homo sapi	RBBP6	2.56992	2.20E-11	2.56992	386.6505	955.4377
A_24_P14	NM_01647	hypothetica	C3orf19,FL	2.56926	1.46E-22	2.56926	578.3086	1472.812
A_23_P36	NM_01509	Homo sapi	CAMTA2,K	2.56496	1.33E-39	2.56496	604.857	1548.216
A_23_P39	NM_15375	Homo sapi	C21orf81	2.56483	0.00001	2.56483	222.7842	519.5514
A_24_P88	NM_02100	Homo sapi	UBC,HMG	2.54935	0	2.54935	283920.2	723539.8
A_24_P92	AK096990	Homo sapi	CHCHD5,C	2.54868	0	2.54868	4234.411	10799.4
A_23_P20	NM_01539	Homo sapi	ZNF10,KO	2.5471	8.48E-10	2.5471	384.3978	934.0952
A_23_P31	AK128423	Cold induci	CIRBP	2.54305	1.01E-41	2.54305	880.3613	2236.393

A_23_P20	NM_02476	hypothetica	WDR78,FL	2.54109	3.50E-12	2.54109	896.6664	2271.222
A_23_P42	NM_02172	nuclear rec	NR1D1,EA	2.54013	2.48E-19	2.54013	2057.369	5268.138
A_24_P25	NM_00407	CDC-like k	CLK1,CLK	2.52713	8.25E-07	2.52713	270.1523	631.1002
A_23_P35	NM_02099	Homo sapi	NEUROG3	2.51822	3.61E-11	2.51822	7827.543	20385.57
A_24_P31	AK094323	Nuclear tra	NFYC	2.51811	3.34E-15	2.51811	7642.242	19600.97
A_23_P26	NM_00329	Homo sapi	TRA1,ECG	2.51731	1.32E-29	2.51731	105917.2	267964.1
A_23_P15	NM_01621	mannosida	MAN1B1,M	2.51608	1.32E-25	2.51608	787.97	1969.978
A_23_P21	NM_00525	FOS-like a	FOSL2,FR	2.51596	1.39E-22	2.51596	1074.17	2705.301
A_32_P19	AW939148	Leiomodin	LMOD1	2.51114	9.06E-11	2.51114	606.3365	1484.907
A_24_P37	AJ001898	Glial cell d	GDNF	2.51008	4.02E-20	2.51008	18236.11	46297.2
A_32_P22	NM_02281	microtubul	MAP1LC3E	2.50522	0	2.50522	2780.76	6968.847
A_23_P14	NM_01567	Homo sapi	GADD45B,	2.50388	9.05E-16	2.50388	2744.628	6900.042
A_24_P932253		Unknown	A_24_P93	2.49894	1.54E-10	2.49894	8252.315	20662.02
A_32_P79115		Unknown	A_32_P79	2.49723	0	2.49723	2867.342	7155.977
A_24_P27	NM_17221	Homo sapi	CAMKK2,C	2.48682	1.54E-15	2.48682	8110.683	20549.31
A_24_P29	NM_15320	Heat shock	HSPA8	2.48281	0	2.48281	7462.404	18508.43
A_23_P61	BC017083	Pim-3 onc	PIM3	2.47215	3.96E-32	2.47215	14466.25	35977.09
A_24_P324284		Unknown	A_24_P32	2.46917	7.04E-11	2.46917	10107.47	25196.89
A_32_P20	BC035247	Homo sapi	BC035247	2.4662	0.00005	2.4662	170.7468	395.2826
A_24_P33	AK024229	Hypothetic	FLJ14167	2.46588	7.23E-20	2.46588	702.1281	1717.162
A_24_P11	NM_03312	Homo sapi	SCRT2	2.46579	6.73E-20	2.46579	23079.79	57540.02
A_32_P225647		CTF1_MO	THC18926	2.46437	1.39E-12	2.46437	4515.791	11819.1
A_24_P57	BX089650	Hypothetic	BX089650	2.46226	9.41E-13	2.46226	5808.797	14483.25
A_23_P43	NM_17588	hypothetica	LOC22217	2.46101	2.37E-18	2.46101	2613.656	6417.805
A_24_P37	NM_00403	interferon r	IRF7,IRF7	2.45558	9.62E-32	2.45558	1218.714	2989.141
A_23_P34	NM_00432	BCL2-asso	BAX,Bax z	2.45373	8.94E-28	2.45373	7428.259	18320.12
A_23_P15	NM_00532	Homo sapi	H3F3B,H3.	2.45256	0	2.45256	68518.91	167919.8
A_23_P43	NM_02518	Homo sapi	KIAA1539,	2.45195	4.54E-18	2.45195	4586.2	11314.83
A_24_P15	XM_37300	similar to H	LOC39156	2.44617	4.40E-19	2.44617	548.422	1357.026
A_24_P58	XM_37327	similar to m	LOC39228	2.44558	5.37E-12	2.44558	715.1187	1722.632
A_23_P57836		Unknown	A_23_P57	2.44444	2.00E-15	2.44444	2367.339	5906.698
A_23_P22	NM_01327	Homo sapi	PCSK1N,S	2.44441	3.36E-15	2.44441	13652.11	34211.25
A_24_P95	NM_01504	KIAA0625	ALS4,AOA	2.44381	9.70E-31	2.44381	2498.62	6147.89
A_32_P38821		Unknown	A_32_P38	2.44318	0.00029	2.44318	123.2019	303.5442
A_23_P79	NM_00532	Homo sapi	HIST1H1E	2.44022	5.21E-23	2.44022	3317.858	8195.401
A_23_P79	NM_00057	Homo sapi	IL1B,IL-1,I	2.43999	1.56E-09	2.43999	1143.558	2720.698
A_23_P20	NM_01860	Homo sapi	DNAJA4,M	2.43475	2.67E-11	2.43475	2706.578	6715.877
A_23_P29	NM_00262	phosphofru	PFKL,PFK	2.43331	6.93E-15	2.43331	22356.71	55783.6
A_24_P53	NM_20734	hypothetica	LOC28473	2.43148	3.54E-11	2.43148	3197.826	8012.804
A_32_P201107		Unknown	A_32_P20	2.42669	1.04E-14	2.42669	8442.07	20777.19
A_32_P74	BC069026	Sp5 transc	SP5	2.41861	1.57E-16	2.41861	18449.32	44984.37
A_24_P34	NM_00623	POU doma	POU3F3,B	2.41749	3.38E-22	2.41749	15316.58	37295.74
A_23_P35	NM_17844	tuberoinfun	TIP39	2.41673	1.06E-15	2.41673	6667.761	16551.02
A_23_P10	NM_01905	Homo sapi	DDIT4,Dig	2.41512	0	2.41512	38138.08	92128.66
A_32_P86180		Unknown	A_32_P86	2.41508	2.35E-14	2.41508	11542.3	28424.27
A_24_P14	NM_02210	G-protein s	GPSM3	2.41184	1.72E-17	2.41184	19325.54	46853.91
A_24_P36	NM_17078	Homo sapi	PPP1R11,I	2.41055	7.18E-16	2.41055	27336.12	66699.78
A_32_P45	NM_20342	hypothetica	LOC38919	2.40641	1.18E-10	2.40641	11426.62	27822.85
A_24_P56	NM_00345	zinc finger	ZNF205,ZN	2.39984	1.49E-18	2.39984	13336.39	31896.53
A_32_P35	NM_02100	deformed e	DEAF1,SP	2.39895	2.17E-14	2.39895	9954.967	24361.8
A_24_P35	NM_03230	Hydroxyac	HAGHL	2.39693	3.66E-08	2.39693	2199.811	5361.87
A_24_P84	XM_37322	similar to H	LOC39215	2.39669	3.67E-18	2.39669	12770.76	30935.35

A_32_P88317	BA299N6.3	ENST0000	2.39388	4.65E-15	2.39388	27313.36	66654.2
A_23_P48	XR_00022	RPL13-2 p	LOC28334	2.39016	3.40E-13	2.39016	521.707	1222.23
A_24_P110101	Unknown	ENST0000	2.38949	3.44E-17	2.38949	23585.51	56962.44
A_23_P27	NM_01895	Homo sapi	UBB,MGC	2.38946	2.62E-39	2.38946	224145.8	538299.3
A_23_P12	NM_00117	Homo sapi	ARG2	2.38374	7.46E-08	2.38374	353.4327	796.2729
A_32_P11	NM_00196	eukaryotic	EIF4A2,DD	2.3833	4.02E-19	2.3833	1600.611	3810.729
A_24_P15	XM_29107	hypothetica	LOC90113	2.3831	1.29E-10	2.3831	6390.251	15706.25
A_23_P103951	Unknown	A_23_P10	2.38052	7.25E-27	2.38052	6602.098	15817.78
A_32_P17	AK128457	Similar to L	AK128457	2.37924	3.17E-17	2.37924	19508.95	46884.03
A_24_P37	NM_02204	solute carri	SLC26A1,E	2.37901	1.07E-10	2.37901	52448.92	128533.3
A_23_P15	NM_02451	leucine-rich	LRRC2	2.37761	4.57E-08	2.37761	298.8281	679.2417
A_24_P18	XM_37692	Homo sapi	XM_37692	2.37504	2.19E-18	2.37504	14131.08	33885.12
A_32_P9931	ROU_HUM	THC18914	2.37422	2.43E-09	2.37422	302.5657	697.1708
A_24_P332953	Unknown	A_24_P33	2.37326	1.12E-30	2.37326	2676.602	6373.859
A_24_P12	NM_01506	Homo sapi	TRIM35,HU	2.37319	5.22E-11	2.37319	16425.21	40186.59
A_24_P716405	FXM1_HUI	THC19576	2.37044	1.45E-10	2.37044	6541.567	15690.87
A_24_P845649	Unknown	A_24_P84	2.36639	1.16E-10	2.36639	12028.5	29169.66
A_23_P34	NM_01989	Homo sapi	ERO1LB	2.36639	7.81E-06	2.36639	217.9644	559.3491
A_24_P54	BG620990	Hypothetica	BG620990	2.36506	2.59E-13	2.36506	28963.14	70203.58
A_23_P96	NM_02156	glutamate r	GRIN1,NR	2.36444	4.48E-10	2.36444	18355.27	44168.75
A_23_P12	NM_02520	PH domain	pp9099,PP	2.36333	2.34E-16	2.36333	524.8771	1221.667
A_23_P32	NM_00292	Homo sapi	RGS16,RG	2.3616	5.26E-26	2.3616	4584.614	10896.23
A_24_P24	XM_37107	Homo sapi	XM_37107	2.36102	4.43E-11	2.36102	20315.4	48614.82
A_24_P26	NM_03096	sprouty (Dr	SPRY4	2.35723	4.24E-07	2.35723	201.4902	465.3776
A_24_P11	NM_01895	ubiquitin B	UBB,MGC	2.35445	5.45E-40	2.35445	307110.4	722568.9
A_32_P21	BC030211	Hypothetica	LOC49725	2.35383	2.31E-10	2.35383	2581.276	6016.134
A_23_P25	NM_00510	Homo sapi	BRD2,NAT	2.35296	1.24E-16	2.35296	477.6278	1132.148
A_24_P30	NM_00510	bromodom	BRD2,NAT	2.35223	1.14E-20	2.35223	24105.66	56825.52
A_24_P27	NM_00622	protein (pe	PIN1L	2.34958	1.81E-13	2.34958	3809.381	9142.814
A_24_P41	NM_17494	Homo sapi	ZNF575,FL	2.34862	1.65E-09	2.34862	29147.63	69551.66
A_32_P203959	Unknown	A_32_P20	2.34759	6.30E-09	2.34759	41132.46	100374.7
A_24_P20	NM_00132	C-terminal	CTBP1,MG	2.34496	5.87E-13	2.34496	10192.48	24261.7
A_23_P34	BX647822	FOS-like a	FOSL2	2.34473	1.47E-16	2.34473	1106.635	2587.674
A_24_P11	NM_15251	Homo sapi	NM_15251	2.34467	8.63E-17	2.34467	3590.989	8390.184
A_24_P12	NM_20158	Homo sapi	MAFA,hMa	2.3444	8.55E-13	2.3444	18167.71	42750.61
A_24_P36	NM_02472	hypothetica	ACBD4,FL	2.34338	2.26E-07	2.34338	18847.19	45428.42
A_23_P14	BC047030	Uridine phc	UPP1	2.34309	9.30E-42	2.34309	3002.552	7054.849
A_23_P12	NM_03302	AXIN1 up-r	AXUD1,UF	2.3417	2.15E-07	2.3417	415.3676	915.8219
A_23_P33	NM_00353	Homo sapi	HIST1H3H	2.33935	1.66E-24	2.33935	1512.266	3543.769
A_32_P50	NM_20732	nuclear loc	NLF1	2.33874	1.34E-08	2.33874	25628.97	62980.88
A_23_P21	NM_00194	Homo sapi	HBEGF,DT	2.33831	2.34E-13	2.33831	725.437	1666.799
A_24_P32	NM_01617	NCX protei	TLX2,Enx,I	2.33742	1.15E-14	2.33742	5203.611	12308
A_32_P139894	ALU7_HUM	THC19992	2.33639	1.26E-27	2.33639	1018.735	2374.496
A_24_P86	NM_00154	heat shock	HSPB1,CM	2.33084	4.91E-17	2.33084	31350.25	74256.88
A_32_P44	BU602485	Transcribe	BU602485	2.33032	6.24E-07	2.33032	204.2488	472.1972
A_23_P43	NM_00353	Homo sapi	HIST1H4A	2.32994	1.41E-08	2.32994	711.757	1612.248
A_32_P76	NM_00154	heat shock	HSPB1,CM	2.32764	1.32E-16	2.32764	42255.27	99969.5
A_24_P60	AK056793	Hypothetica	LOC14569	2.32632	1.65E-09	2.32632	14390.83	34295.88
A_23_P97	NM_00585	Homo sapi	IRX5,IRX-2	2.32555	7.96E-20	2.32555	1383.004	3227.126
A_24_P25	NM_01436	NK-2 (Dros	NKX2-8,NK	2.32286	1.70E-14	2.32286	17242.66	40321.25
A_32_P122907	AY211916	THC18595	2.32131	1.73E-10	2.32131	26471.39	62755.94
A_24_P39	AK025430	Homo sapi	FLJ21777	2.32079	3.87E-14	2.32079	7840.694	18374.14

A_32_P206175	PTNI_HUM	THC19327	2.31949	2.23E-11	2.31949	11461.98	27401.33
A_23_P20	NM_00528	Homo sapi	GPR7,MG	2.31825	1.27E-12	2.31825	12159.11	28512.07
A_23_P56	NM_00446	Homo sapi	FAP,FAPA	2.31435	0.00002	2.31435	217.9679	473.5095
A_23_P92	NM_15300	Homo sapi	GPR156,P	2.31337	2.81E-15	2.31337	21795.87	51002.79
A_24_P87	NM_01515	KIAA0071	RCOR1,R	2.31332	0.00072	2.31332	128.0499	293.3349
A_24_P878992	Unknown	A_24_P87	2.31123	5.83E-16	2.31123	22885.2	53203.91
A_24_P17	NM_01343	gamma-glu	GGT1,GGT	2.31116	6.24E-16	2.31116	10440.22	24120.16
A_24_P93	XM_37947	hypothetica	LOC40131	2.30885	3.86E-11	2.30885	292.8846	669.4074
A_24_P92	AF113674	Homo sapi	PRO0398	2.3087	0.00017	2.3087	176.9778	384.4957
A_24_P928250	Unknown	A_24_P92	2.3074	1.70E-26	2.3074	19382.09	44894.85
A_24_P75	NM_19945	testis/prost	TEPP,MG	2.30678	3.44E-07	2.30678	1418.073	3217.978
A_32_P19	BU191598	Solute carr	SLC19A1	2.30396	1.81E-10	2.30396	13933.65	32889.38
A_24_P91	X89741	H.sapiens	CHRNA4	2.30352	3.08E-12	2.30352	5890.765	13939.35
A_32_P233870	Unknown	A_32_P23	2.30314	1.11E-10	2.30314	9770.813	22618.28
A_24_P58	NM_00311	Homo sapi	SP3,SPR-2	2.303	2.29E-12	2.303	7184.328	16694.21
A_24_P20	NM_02433	Homo sapi	IRX3,IRX-1	2.30207	4.44E-12	2.30207	37502.6	88799.63
A_32_P37943	Unknown	A_32_P37	2.30203	1.24E-10	2.30203	8851.226	20942.05
A_23_P152570	Unknown	THC18494	2.30171	3.62E-24	2.30171	1529.958	3540.136
A_23_P10	NM_19839	c-Maf-indu	CMIP,KIA	2.30157	1.24E-08	2.30157	68800.1	165296.3
A_32_P15887	Unknown	A_32_P15	2.30133	8.10E-09	2.30133	48346.59	115068.6
A_24_P22	NM_00351	H2A histon	HIST1H2A	2.29914	0	2.29914	1664.383	3827.065
A_23_P10	NM_00693	solute carri	SLC5A3,S	2.29369	3.47E-10	2.29369	313.8502	712.3265
A_23_P12	NM_00540	Homo sapi	CXCL11,I	2.29129	3.20E-07	2.29129	212.9097	480.3013
A_32_P85	AK074191	Zinc finger	MGC1571	2.29107	1.28E-13	2.29107	8269.049	19041.02
A_24_P31	AB015330	Hypothetic	DKFZp761	2.28944	3.00E-08	2.28944	5917.715	13593.76
A_32_P13	BC020341	Homo sapi	ACK1	2.289	1.95E-11	2.289	9860.991	22840.36
A_24_P745960	Unknown	A_24_P74	2.288	1.06E-09	2.288	30355.23	70653.98
A_32_P160254	121509 cut	THC19154	2.28792	1.37E-11	2.28792	31589.84	74112.32
A_24_P39	AK026328	Hypothetic	FLJ22675	2.28762	5.56E-22	2.28762	805.9166	1855.302
A_23_P34	NM_13090	Homo sapi	C15orf16,C	2.28688	4.09E-14	2.28688	1820.828	4225.295
A_24_P27	NM_00600	zinc finger	ZA20D2,Z	2.28578	5.63E-22	2.28578	699.4105	1591.954
A_24_P30	BC034752	Chromosor	C9orf62	2.28555	2.34E-12	2.28555	67944.29	157320
A_23_P25	NM_00154	Homo sapi	HSPB1,CM	2.28315	4.26E-16	2.28315	31334.19	73124.39
A_24_P18	NM_03091	likely ortho	LBH,DKFZ	2.28192	5.91E-13	2.28192	32210.08	74423.89
A_23_P70	NM_00337	vascular er	VEGF,VPF	2.28026	2.80E-13	2.28026	499.4006	1120.465
A_23_P16	NM_01245	Homo sapi	SYNGR4,M	2.28017	1.61E-10	2.28017	14310.95	33354.97
A_24_P91	BC037430	CDNA clon	BC037430	2.27997	3.25E-21	2.27997	1019.939	2322.79
A_24_P230195	Unknown	A_24_P23	2.27667	2.39E-12	2.27667	43744.84	101695.9
A_24_P52	NM_00380	harakiri, B	HRK,DP5,I	2.27655	1.47E-16	2.27655	12720.83	29372.28
A_23_P10	NM_00049	Homo sapi	AVP,VP,A	2.27455	2.60E-12	2.27455	1913.172	4377.95
A_24_P551530	AP003962	THC19123	2.27195	3.56E-10	2.27195	21042.08	48984.28
A_24_P43	NM_01802	hypothetica	FRMD4A,F	2.27142	3.45E-10	2.27142	24259.8	56032.88
A_24_P16	NM_00108	cytochrom	CYP4F2,C	2.27074	3.92E-08	2.27074	8761.166	20836.42
A_32_P233278	S14984 gly	THC19005	2.27044	1.17E-10	2.27044	35393.81	82547.2
A_23_P13	NM_17335	Homo sapi	SNF1LK,M	2.27005	1.64E-31	2.27005	4687.283	10758.33
A_24_P461664	Unknown	A_24_P46	2.26997	8.09E-08	2.26997	51040.55	119850.3
A_24_P780709	Unknown	A_24_P78	2.26918	2.21E-16	2.26918	67321.66	153927
A_24_P478362	seven tran	NP511100	2.26802	2.33E-11	2.26802	15567.04	36258.89
A_24_P21	NM_01211	CCR4-like	CCRN4L,N	2.26652	9.90E-11	2.26652	675.9774	1500.38
A_23_P21	BX538238	Metastasis	MALAT1	2.26621	3.86E-09	2.26621	966.2733	2265.636
A_32_P196247	AF195969	THC19600	2.26482	1.25E-13	2.26482	46081.5	105955.9
A_23_P136753	AF343666	THC18988	2.26338	2.67E-09	2.26338	32287.86	75538.12

A_23_P62	NM_01233	nasopharyr	CCDC19,N	2.26154	4.82E-07	2.26154	304.8424	655.428
A_24_P10	NM_00450	isopenteny	IDI1,IPP1,I	2.25771	5.30E-38	2.25771	4255.183	9646.582
A_24_P23	NM_17792	Homo sapi	H2AFJ,MG	2.25537	3.11E-23	2.25537	33126.37	75231.03
A_23_P34	NM_00072	Homo sapi	CACNA1E	2.2539	1.21E-06	2.2539	42284.81	100832.3
A_24_P65	NM_01458	SRY (sex c	SOX8,MG	2.24873	1.13E-10	2.24873	23903.5	54859.91
A_24_P14	NM_01327	granin-like	PCSK1N,S	2.248	3.45E-10	2.248	40414.8	92239.36
A_24_P635355		Unknown	A_24_P63	2.24658	3.90E-13	2.24658	1198.072	2682.896
A_32_P116219		Unknown	A_32_P11	2.2464	3.37E-09	2.2464	19603.88	44665.75
A_23_P16	NM_02520	hypothetica	PP2447,LF	2.24422	8.16E-13	2.24422	14240.13	32127.94
A_24_P26	NM_00616	neurogenin	NEUROG1	2.24261	1.44E-11	2.24261	29439.79	66656.98
A_23_P16	NM_00565	Homo sapi	KLF10,EGI	2.23914	0	2.23914	1583.748	3545.565
A_24_P10	NM_02526	CAT56 pro	PRR3,CAT	2.23904	6.07E-18	2.23904	4256.846	9627.816
A_24_P849245		Unknown	THC18695	2.2387	4.65E-19	2.2387	1823.496	4102.604
A_23_P50	NM_13349	Homo sapi	SYN1,SYN	2.23716	1.50E-09	2.23716	43871.91	100311.5
A_24_P927245		Unknown	A_24_P92	2.23705	8.29E-14	2.23705	24251.02	54298.81
A_24_P30	NM_02472	MICAL-like	MICAL-L2	2.23496	9.18E-33	2.23496	3099.84	6952.08
A_23_P84	NM_02156	Homo sapi	GRIN1,NR	2.23433	1.05E-11	2.23433	23032.41	52667.91
A_24_P10	NM_15333	Homo sapi	SCARF2,N	2.23335	2.86E-14	2.23335	30260.48	67990.4
A_24_P770494		Unknown	A_24_P77	2.23306	4.06E-09	2.23306	6097.29	14265.11
A_24_P94	NM_00578	Tyrosine ki	TNK2	2.23114	2.11E-09	2.23114	8735.623	20110.36
A_23_P20	NM_01431	Homo sapi	DDX58,RI	2.23103	1.21E-19	2.23103	671.3577	1482.764
A_23_P41	NM_00519	Homo sapi	CEBPB,LA	2.23102	5.05E-19	2.23102	25690.23	58040.99
A_24_P471099		Unknown	A_24_P47	2.23087	6.85E-09	2.23087	23456.99	53193.89
A_24_P21	NM_20733	Homo sapi	ZNF467,E	2.23079	2.09E-10	2.23079	60567.52	137310.5
A_32_P216602		ALU1_HUM	THC19686	2.23076	7.75E-13	2.23076	49117.98	111054.1
A_32_P33	AF390550	Zinc finger	ZNF434	2.22984	2.12E-08	2.22984	337.0291	725.3428
A_24_P68	BC043212	Hypothetic	BC043212	2.22945	1.08E-17	2.22945	502.1794	1120.954
A_23_P31	NM_00519	Homo sapi	CEBPD,CE	2.22872	1.20E-31	2.22872	6160.069	13777.92
A_23_P25	NM_14498	Homo sapi	CDH24,CD	2.22775	1.01E-09	2.22775	54433.55	123089.3
A_24_P91	AF343007	Homo sapi	FOXC1,AR	2.22612	5.93E-19	2.22612	3927.256	8798.369
A_24_P15	AK091796	Hypothetic	MGC13045	2.22608	2.24E-08	2.22608	11092.22	25658.84
A_32_P11	BE300110	CDNA clon	UBB,MGC	2.22419	9.81E-45	2.22419	3131.419	6964.229
A_23_P10	AB037863	KIAA1442	KIAA1442	2.22365	2.93E-12	2.22365	7654.28	17118.51
A_24_P82	NM_00525	forkhead b	FOXC2,MF	2.22323	7.35E-12	2.22323	33699.96	75645.34
A_24_P11	NM_00598	SRY (sex c	SOX1	2.22313	1.49E-11	2.22313	41167.95	92189.22
A_32_P78	NM_00496	high-mobili	HMGN1,HI	2.22195	2.20E-29	2.22195	6566.595	14684.68
A_24_P25	NM_00588	adenomato	APC2,APC	2.22069	3.15E-09	2.22069	14390.91	32364.85
A_24_P37	NM_01654	immediate	IER5,SBBI	2.22065	5.54E-10	2.22065	85116.48	192464
A_23_P35	X94553	H.sapiens	HFKH4	2.2175	2.32E-08	2.2175	26977.52	62544.04
A_32_P78	NM_14527	similar to h	LOC14632	2.21601	2.13E-12	2.21601	18065.64	40746.69
A_23_P36	NM_02467	hypothetica	C1orf113,F	2.21576	5.26E-11	2.21576	22005.07	50095.63
A_24_P23	BC004366	Hypothetic	MGC10334	2.21539	1.77E-10	2.21539	22416.93	50057.68
A_23_P20	NM_02118	Homo sapi	IGSF4B,BI	2.21041	4.06E-11	2.21041	3918.342	8717.533
A_32_P65793		Unknown	THC18106	2.20905	2.36E-09	2.20905	98548.52	225100.8
A_23_P54	NM_03127	Testis expr	TEX14	2.20727	6.83E-06	2.20727	198.0648	433.7889
A_24_P10	NM_14517	Homo sapi	C21orf93	2.20655	5.61E-19	2.20655	7502.098	16529.09
A_24_P778928		WASL_BO	THC19380	2.20472	8.64E-10	2.20472	28873.35	65130.68
A_24_P46	NM_02245	likely ortho	SOX17,FL	2.20366	1.51E-10	2.20366	39282.23	87320.32
A_24_P87	BC043357	Discs, larg	DLGAP1	2.20361	5.64E-07	2.20361	248.3996	556.3915
A_24_P36	NM_14576	Homo sapi	GFRA4	2.2035	9.26E-13	2.2035	12244.95	27378.84
A_23_P37	BC004262	Chromosor	C19orf29	2.20349	4.09E-08	2.20349	8511.482	19571.86
A_24_P21	NM_00181	Homo sapi	CENPB	2.2018	3.69E-16	2.2018	3319.04	7260.771

A_24_P929946		ORF [NP11	NP111687	2.20159	7.52E-09	2.20159	7066.536	16052.63
A_32_P19	AK027019	Hypothetic	MGC45731	2.20158	0.00079	2.20158	151.2135	348.1298
A_23_P80	AL834499	Hypothetic	LOC90113	2.19817	1.74E-11	2.19817	6621.675	14652.12
A_24_P25	NM_02194	Chromosor	C6orf85	2.1972	4.24E-08	2.1972	12330.83	27901.87
A_24_P15	AK098038	Ubiquitin-li	UHRF1	2.1971	1.75E-09	2.1971	46472.59	104273.9
A_23_P14	NM_01218	Homo sapi	FOXB1,FK	2.19678	1.85E-10	2.19678	21390.61	47460.24
A_24_P15	XM_37160	Homo sapi	ENST0000	2.19626	1.95E-08	2.19626	9514.127	21766.66
A_23_P10	NM_00585	Homo sapi	CDK2AP2,	2.19596	7.23E-32	2.19596	871.3274	1911.318
A_23_P15	NM_03280	Homo sapi	ZNF206,ZS	2.19505	3.19E-09	2.19505	59132.36	132603
A_24_P28	AF490258	KIAA1545	KIAA1545	2.19425	6.42E-09	2.19425	55336.22	126354.1
A_32_P13	NM_01233	methionine	MSRA	2.1928	5.38E-15	2.1928	6503.083	14337.9
A_24_P92	NM_20330	Homo sapi	RKHD1,TIN	2.19073	2.10E-11	2.19073	90750.73	203786.3
A_23_P57	NM_13893	Homo sapi	BCL6,BCL	2.19007	2.57E-13	2.19007	662.4238	1431.324
A_23_P32	NM_03257	Homo sapi	GLIS2	2.18975	1.10E-11	2.18975	3783.528	8253.561
A_24_P57898		BA305P22	ENST0000	2.18692	1.82E-12	2.18692	37543.38	83371.84
A_24_P21	NM_00436	CCAAT/en	CEBPA,CE	2.18644	1.59E-08	2.18644	68921.49	157206.2
A_23_P38	NM_00535	Homo sapi	LIPE,HSL,I	2.18525	1.92E-25	2.18525	4175.011	9109.696
A_23_P10	NM_03255	Homo sapi	GPR54,AX	2.18498	1.40E-12	2.18498	16165.2	35450.88
A_24_P78	XM_29421	similar to R	LOC34635	2.18198	5.35E-10	2.18198	52676.59	116810
A_24_P38	NM_13345	RAVER1	RAVER1,K	2.18083	1.45E-08	2.18083	58517.13	129262.3
A_24_P38	NM_00612	mannosida	MAN2A2,M	2.17947	9.60E-14	2.17947	16122.56	35393.94
A_23_P69	NM_08081	Homo sapi	GPR78	2.17947	2.55E-13	2.17947	6139.677	13481.08
A_23_P12	NM_13844	Homo sapi	SLITL2,VA	2.17909	1.03E-08	2.17909	1001.798	2156.222
A_23_P36	NM_17851	hypothetica	LOC28348	2.17787	3.42E-12	2.17787	1502.9	3252.024
A_24_P29	NM_00600	zinc finger	ZA20D2,ZN	2.17646	3.50E-19	2.17646	3248.747	7120.91
A_24_P13	BC009233	Hypothetic	LOC92659	2.1759	1.39E-09	2.1759	39768.2	87520
A_32_P33294		Unknown	A_32_P33	2.17572	1.30E-10	2.17572	35223.4	77149.78
A_24_P39	AK092271	Homo sapi	AK092271	2.17534	3.71E-09	2.17534	95639.55	212767.2
A_24_P83	NM_20740	FLJ45717	FLJ45717	2.17409	3.89E-10	2.17409	35997.25	78678.89
A_23_P30	AF320053	V-myc mye	MYCN	2.17289	3.46E-09	2.17289	15796.7	35217.43
A_24_P22	NM_02283	hypothetica	FLJ22833,	2.17153	0.00005	2.17153	200.7658	454.5627
A_23_P16	NM_00242	Homo sapi	MMP1,CLC	2.17144	0.00031	2.17144	204.678	521.9109
A_23_P13	U06936	D site of all	DBP	2.17132	3.43E-12	2.17132	7233.258	15531.56
A_24_P401663		Unknown	A_24_P40	2.16886	1.70E-06	2.16886	72832.55	163442.2
A_32_P105865		Unknown	A_32_P10	2.16885	7.42E-07	2.16885	90823.95	203221.8
A_24_P16	NM_05283	pannexin 2	PANX2,hP	2.16847	6.31E-11	2.16847	4092.457	8993.366
A_32_P80	AK096255	CDNA FLJ	AK096255	2.16814	4.22E-07	2.16814	543.4111	1229.202
A_32_P17	NM_01849	G protein-c	LGR4,GPR	2.16746	3.25E-08	2.16746	682.0275	1433.833
A_24_P760708		Unknown	THC18699	2.16736	1.25E-12	2.16736	15458.29	33654.98
A_24_P161581		Unknown	A_24_P16	2.16687	1.71E-08	2.16687	14635.75	32479.87
A_23_P54	NM_15381	Homo sapi	ZFPM1,FO	2.16651	5.78E-12	2.16651	80309.24	175996.2
A_32_P101073		Unknown	A_32_P10	2.16526	3.39E-09	2.16526	86036.22	190324.3
A_24_P791968		AF000298	THC18503	2.16388	1.47E-09	2.16388	78715.23	172970.9
A_24_P247169		Unknown	A_24_P24	2.16315	1.96E-07	2.16315	58221.24	130887.6
A_32_P21	NM_00329	tropomyosi	TPM4	2.16293	1.01E-08	2.16293	3503.843	7734.726
A_32_P44073		Unknown	A_32_P44	2.1605	1.72E-16	2.1605	2730.456	5923.724
A_24_P23	BC033110	Hydroxyste	HSD17B1	2.15819	1.62E-12	2.15819	23927.5	52676.47
A_24_P27	NM_00494	Homo sapi	DOCK3,MC	2.15782	2.60E-08	2.15782	20173.78	43996.94
A_24_P41	NM_02495	hypothetica	C20orf98,F	2.1568	1.07E-09	2.1568	12131.63	26797.32
A_23_P43	AF193612	Lunatic frin	LFNG	2.15677	6.74E-09	2.15677	7478.292	16567.52
A_24_P32	AF161351	HSPC088	HSPC088	2.1549	4.38E-08	2.1549	57939.7	126726.1
A_23_P11	NM_08087	Homo sapi	SLC34A3,N	2.15303	7.60E-10	2.15303	32074.27	69425.27

A_23_P78	NM_00556	Homo sapi	LHX1,LIM1	2.15292	2.87E-10	2.15292	9798.729	21012.29
A_24_P87	NM_01664	mesenchyr	LOC51333	2.15233	3.82E-12	2.15233	1406.396	3058.791
A_24_P39	NM_02109	calcium ch	CACNA1I,	2.15228	3.57E-10	2.15228	110764.6	245848.6
A_32_P703		Unknown	A_32_P70	2.15203	3.10E-08	2.15203	290.4263	615.5013
A_24_P12	NM_03318	keratin ass	KRTAP2-4	2.15099	4.63E-09	2.15099	23472.84	51007.2
A_24_P63	BC036435	Homo sapi	BC036435	2.15077	3.01E-10	2.15077	12667.63	27552.5
A_24_P22	AK095945	Hypothetic	MGC12458	2.14931	4.06E-12	2.14931	31394.07	68383.48
A_24_P21	NM_01910	tenascin XI	TNXB,XB,	2.14872	7.71E-08	2.14872	57287.88	122936.4
A_23_P41	NM_03319	Homo sapi	HSPB9	2.14731	4.04E-13	2.14731	4349.104	9368.938
A_23_P12	NM_00219	Homo sapi	INHBA,EDI	2.14487	4.71E-13	2.14487	500.0403	1054.616
A_24_P40	BC025310	Leucine ric	LRFN1	2.14448	6.89E-11	2.14448	1405.331	3022.182
A_23_P14	NM_03251	Homo sapi	PARD6G,F	2.14408	1.45E-07	2.14408	88488.66	194831.3
A_32_P20	BC063316	Homo sapi	FBXL17	2.14404	2.14E-07	2.14404	125878.4	277412.6
A_23_P16	NM_00407	Homo sapi	CLK1,CLK	2.1421	1.97E-16	2.1421	911.973	1941.252
A_24_P59	BC009749	TatD DNas	TATDN2	2.14087	2.78E-18	2.14087	3474.444	7400.155
A_24_P61	NM_00182	creatine kir	CKB,B-CK	2.14061	7.45E-07	2.14061	103805	229390.1
A_32_P10	NM_00600	zinc finger	ZA20D2,ZN	2.13902	9.43E-09	2.13902	850.3091	1793.648
A_23_P78	NM_01239	Homo sapi	PIP5K1C,K	2.13825	2.98E-11	2.13825	35700.92	77847.56
A_24_P38	NM_18249	hypothetica	LRP5L,DKI	2.13824	1.59E-08	2.13824	404.4229	837.587
A_24_P533990		Unknown	A_24_P53	2.13775	1.07E-07	2.13775	153129.4	335842.4
A_24_P40	NM_00520	Homo sapi	COX6A2,C	2.13749	9.56E-10	2.13749	104860.3	229798.5
A_23_P20	NM_02255	Homo sapi	GH2,GHL,	2.13711	5.59E-11	2.13711	9005.344	19586.47
A_24_P29	NM_20748	FLJ34870	FLJ34870	2.13674	1.45E-07	2.13674	385.0963	789.8936
A_23_P18	NM_03325	CaM-KII inl	CAMK2N2	2.13452	1.02E-07	2.13452	3292.796	7077.532
A_23_P53	M10036	Human trio	TPI1	2.13365	1.49E-13	2.13365	5189.19	11294.01
A_24_P92	Y10152	Corticotrop	CRHR2	2.13347	3.32E-08	2.13347	126829	280266.1
A_24_P40	NM_00728	synaptopo	SYNPO,KI	2.13147	6.57E-10	2.13147	37594.53	81946.92
A_23_P42	NM_15221	Homo sapi	GJC1,GJA	2.13057	3.77E-08	2.13057	10268.12	21899.66
A_24_P34	NM_15247	Homo sapi	RNF165	2.12856	5.73E-07	2.12856	180649.6	399546.4
A_24_P753760		153353 tro	THC18313	2.12669	3.21E-11	2.12669	1977.786	4152.569
A_23_P41	NM_00427	Homo sapi	HOMER1,	2.12609	2.39E-21	2.12609	2394.341	5120.826
A_23_P46	NM_02520	Homo sapi	EPC1,DKF	2.12541	9.62E-10	2.12541	531.6108	1098.473
A_24_P20	BC037281	Hypothetic	LOC43995	2.12341	8.93E-09	2.12341	116908.8	252601.8
A_24_P37	NM_19924	Homo sapi	GPR150,P	2.12327	1.32E-06	2.12327	181923.2	396331
A_24_P94	AF220656	Pleckstrin	PHLDA1	2.12313	3.06E-08	2.12313	581.7231	1192.905
A_23_P48	NM_03094	Homo sapi	AMN,PRO	2.12067	1.49E-11	2.12067	17434.8	37229.62
A_32_P85711		Unknown	A_32_P85	2.1195	2.01E-08	2.1195	101176	222598.3
A_23_P25	NM_01841	Homo sapi	SOX18,HL	2.1177	4.48E-10	2.1177	14302.98	31260.48
A_24_P30	NM_00247	myogenic f	MYOD1,PU	2.11613	8.27E-14	2.11613	4974.084	10549.11
A_32_P17	NM_19430	hypothetica	DKFZp434	2.11519	2.96E-09	2.11519	84355.23	181205.3
A_24_P47	AK093171	Hypothetic	AK093171	2.11434	3.70E-10	2.11434	61006.3	131641.5
A_32_P106925		Unknown	THC18879	2.11385	2.10E-12	2.11385	8721.877	18322.1
A_32_P200820		Unknown	A_32_P20	2.11188	2.23E-08	2.11188	13405.46	28706.75
A_24_P27	BC036731	Zinc finger	ZBTB24	2.11186	7.31E-07	2.11186	265.0565	550.9315
A_24_P16	NM_01754	hypothetica	H41	2.11123	3.23E-39	2.11123	7800.949	16511.94
A_23_P78	NM_02472	Homo sapi	MYH14,DF	2.11104	8.48E-10	2.11104	42363.2	90440.13
A_32_P49	BM720538	Transcribe	BM720538	2.11088	3.82E-07	2.11088	321.4457	655.5963
A_24_P11	NM_00563	SRY (sex	SOX3,MR	2.11037	2.76E-07	2.11037	275215.3	601968.8
A_24_P91	NM_00532	H3 histone	H3F3B,H3	2.1098	0	2.1098	14736.58	31086.82
A_24_P78	AK095849	RNA, U22	RNU22	2.10931	1.14E-17	2.10931	7066.766	15093.03
A_24_P11	AF318368	Pp8961	pp8961	2.10887	4.49E-10	2.10887	389.814	842.4902
A_24_P49	AL050210	Metastasis	MALAT1	2.10847	2.18E-06	2.10847	1240.849	2548.878

A_24_P10	BC028232	Homo sapi	BC028232	2.10753	4.37E-07	2.10753	114222.9	249417.5
A_23_P21	NM_15277	hypothetica	MGC33212	2.10721	2.14E-14	2.10721	623.4822	1300.969
A_23_P39	NM_00145	Homo sapi	FOXC1,AR	2.10681	3.56E-19	2.10681	14623.61	31118.17
A_23_P33	NM_00357	Homo sapi	UTF1	2.10645	2.39E-13	2.10645	9883.387	20906.55
A_32_P71	BC035731	Similar to C	BC035731	2.10302	1.03E-06	2.10302	161357.2	350381.8
A_23_P39	NM_01440	GPI-ancho	LYPD3,C4	2.10178	2.12E-10	2.10178	16506.54	35289.63
A_23_P15	NM_00200	Homo sapi	FCN1,FCN	2.10167	9.59E-17	2.10167	4719.99	9945.201
A_23_P21	NM_02212	basic-helix-	NPAS3,MC	2.10156	1.74E-07	2.10156	162963.9	354321.2
A_23_P41	NM_00354	H4 histone	HIST1H4E	2.10132	1.20E-17	2.10132	1252.899	2617.258
A_23_P39	NM_00442	dual specif	DUSP8,HB	2.10106	1.01E-32	2.10106	1802.641	3787.997
A_24_P19	NM_03306	keratin ass	KRTAP4-1	2.10063	2.07E-08	2.10063	105907.7	226680
A_23_P43	NM_03208	protocadhe	PCDHGA7	2.10057	2.07E-10	2.10057	11717.51	24925.4
A_23_P43	NM_03306	Homo sapi	ATCAY,CL	2.10005	8.15E-11	2.10005	20434.48	43311.16
A_32_P10	BX090092	KIAA0963	KIAA0963	2.09962	8.08E-14	2.09962	5031.704	10567.71
A_24_P914308		AC007061	THC18485	2.09883	0.00007	2.09883	356.195	691.3527
A_23_P57	NM_02018	Homo sapi	TMEPAI,ST	2.09689	1.63E-11	2.09689	2314.909	4883.502
A_24_P84	AL531532	Full-length	KIAA1924	2.09684	7.71E-10	2.09684	67267.31	145174.3
A_32_P33	NM_17357	Homo sapi	C11orf35,M	2.09606	7.29E-15	2.09606	927.0292	1921.288
A_24_P25	BC036529	Enhancer c	EPC1	2.09508	2.31E-11	2.09508	424.7656	878.596
A_23_P15	NM_20337	similar to R	LOC38911	2.09502	7.34E-08	2.09502	4687	9464.393
A_24_P12	NM_03312	naked cutic	NKD2	2.09444	1.40E-07	2.09444	81843.63	176860.9
A_23_P21	NM_00086	Homo sapi	HRH1,H1-I	2.09267	8.49E-06	2.09267	312.4011	696.3266
A_24_P20	NM_03056	hypothetica	PRR7,MGC	2.09179	9.30E-08	2.09179	188857.7	405355.8
A_23_P97	NM_20737	Homo sapi	GPR153,P	2.09127	6.67E-08	2.09127	140601.5	303506.5
A_24_P862251		Unknown	A_24_P862	2.09081	2.07E-13	2.09081	4850.078	10201.19
A_24_P87	AF001540	Metastasis	MALAT1	2.09051	3.96E-21	2.09051	983.1205	2045.156
A_23_P79	NM_02017	Signal pep	SPPL2B	2.08932	8.24E-08	2.08932	136580.8	295571.6
A_23_P15	NM_00083	Homo sapi	GRIN2D,EI	2.08862	7.18E-08	2.08862	137871.3	295196
A_24_P30	AK075298	Hypothetic	LOC34809	2.08815	2.43E-06	2.08815	9963.075	20824.13
A_32_P101441		Unknown	THC19509	2.08799	2.25E-25	2.08799	1338.79	2794.212
A_24_P14	AK025837	Hypothetic	FLJ22184	2.08722	1.72E-07	2.08722	189681.9	409602.9
A_23_P40	NM_01541	Homo sapi	C20orf28,D	2.08701	1.60E-09	2.08701	13154.28	27498.81
A_24_P76288		Unknown	A_24_P762	2.08672	1.49E-10	2.08672	19353.48	40745.39
A_23_P33	XM_37485	similar to d	LOC39983	2.08621	1.46E-07	2.08621	158156.3	345235.1
A_23_P156609		Unknown	A_23_P156	2.08548	8.43E-06	2.08548	574.481	1136.781
A_23_P62	AB011539	EGF-like-d	EGFL3	2.08435	4.60E-11	2.08435	2356.555	4903.142
A_32_P12	BC024745	Homo sapi	LOC39988	2.08212	7.12E-08	2.08212	680.2843	1366.266
A_23_P29	NM_00180	Homo sapi	CEBPE,CF	2.08174	6.43E-08	2.08174	663.4675	1340.817
A_24_P31	NM_00523	v-ets avian	ETS2	2.08078	2.12E-10	2.08078	430.5418	883.9409
A_23_P16	NM_00704	Homo sapi	EMILIN1,g	2.07827	2.09E-11	2.07827	6514.923	13646.42
A_32_P20	NM_13346	Homo sapi	CITED4	2.07795	6.93E-25	2.07795	5020.807	10480.52
A_24_P94	NM_01755	filamin-binc	FBLIM1,CA	2.07653	4.72E-10	2.07653	3263.032	6633.871
A_32_P33	NM_18257	Homo sapi	C19orf19,F	2.07566	5.05E-11	2.07566	3963.746	8332.157
A_24_P26	NM_20747	Homo sapi	NM_20747	2.07554	9.25E-09	2.07554	58014.52	122764.3
A_24_P17	NM_03191	BTE-bindin	KLF16,DRI	2.07337	1.39E-06	2.07337	155805.1	332533.6
A_24_P85	BU556542	CDNA clon	BU556542	2.07194	3.27E-09	2.07194	6646.813	13848.83
A_32_P11425		Unknown	A_32_P114	2.07001	2.47E-09	2.07001	68371.28	142194.2
A_24_P25	BC001892	NPD014 pr	NPD014	2.06955	7.00E-26	2.06955	1103.844	2282.461
A_23_P13	NM_00544	Homo sapi	FAIM3,TOS	2.06856	1.64E-08	2.06856	2146.414	4487.298
A_32_P188143		AP005936	THC18482	2.06786	3.62E-08	2.06786	178712.8	377234.6
A_23_P14	NM_01894	Homo sapi	UTS2R,UT	2.06728	6.59E-08	2.06728	78045.07	165390.9
A_24_P12	BC018035	Homo sapi	BC018035	2.06625	4.10E-18	2.06625	21111.25	43861.96

A_23_P41	NM_00318	Homo sapi	TAF9,CIP,T	2.06611	9.03E-28	2.06611	16386.81	34045.49
A_23_P21	NM_00549	Homo sapi	DNAJB6,M	2.06608	4.35E-31	2.06608	5403.984	11203.47
A_24_P27	NM_00600	zinc finger	ZA20D2,ZN	2.06602	1.27E-15	2.06602	4366.821	9057.215
A_23_P14	NM_00516	Homo sapi	RND3,ARH	2.06594	1.69E-16	2.06594	3760.874	7793.2
A_23_P81	NM_13084	dendritic ce	C5orf20,DC	2.06503	3.67E-07	2.06503	1728.538	3489.332
A_23_P31	NM_19846	FLJ37183	FLJ37183	2.06427	2.93E-11	2.06427	28373.27	59027.56
A_24_P76	XM_37977	KIAA1856	KIAA1856	2.06404	1.02E-06	2.06404	207092.9	438956.8
A_24_P583549		AE003463	THC18184	2.06324	5.85E-08	2.06324	47293.52	99599.95
A_24_P16	NM_01406	SPR1 prote	PSORS1C	2.06265	3.26E-08	2.06265	44023.52	92150.25
A_24_P20	NM_00652	wingless-ty	WNT6	2.06247	2.53E-07	2.06247	215901.6	457266.8
A_24_P93	D31828	Human mR	T cell recep	2.06198	1.56E-06	2.06198	441.4508	866.3052
A_24_P7785		Unknown	A_24_P77	2.05953	2.62E-06	2.05953	122648.6	258392.5
A_32_P95	BX090181	Transcribe	BX090181	2.05767	7.68E-09	2.05767	13386.46	27667.68
A_23_P20	NM_03210	sema dom	SEMA6B,S	2.05752	4.97E-12	2.05752	3346.313	6911.705
A_32_P21	XM_37947	hypothetica	LOC40131	2.05675	1.30E-10	2.05675	2606.942	5397.226
A_32_P230059		Unknown	A_32_P23	2.05659	2.10E-07	2.05659	118778.9	248853.8
A_23_P11	NM_00016	Homo sapi	GIPR,MGC	2.05503	4.57E-28	2.05503	4158.134	8539.729
A_32_P142013		BC041700	THC19765	2.05345	0	2.05345	3523.249	7234.383
A_23_P61	NM_01745	Homo sapi	BAIAP2,BA	2.05187	9.93E-19	2.05187	1739.138	3558.766
A_24_P922101		Unknown	A_24_P92	2.0513	3.99E-29	2.0513	90889.45	186716.2
A_24_P21	AK057884	Componen	COG3	2.0471	9.01E-09	2.0471	155163	323960.3
A_23_P33	BC022076	Hypothetic	LOC25578	2.04687	5.56E-07	2.04687	208921.3	443091.4
A_24_P16	BC036762	Hypothetic	MGC46336	2.04579	1.54E-07	2.04579	10983.96	22979.98
A_32_P64	H64096	Transcribe	HSPA8,LA	2.04465	1.26E-06	2.04465	5883.073	12139.44
A_24_P87	NM_00366	apolipoprot	APOL1,AP	2.04155	3.68E-12	2.04155	25655.63	52931.87
A_32_P234913		Unknown	A_32_P23	2.04155	1.48E-07	2.04155	187184.1	390143.6
A_24_P32	AK024930	Homo sapi	FLJ21277	2.04118	4.00E-06	2.04118	166648.4	351180.1
A_24_P28	NM_15237	Fc recepto	FLJ31052	2.04063	6.35E-07	2.04063	169268.8	355638
A_23_P32	NM_00423	Homo sapi	KLF4,EZF,	2.03945	7.58E-17	2.03945	1128.348	2285.357
A_32_P90	NM_00318	TATA box	TAF9,CIP,T	2.03907	6.36E-43	2.03907	9746.442	19885.39
A_32_P17	AK128192	Hypothetic	AK128192	2.03905	3.22E-10	2.03905	20552.77	42273.77
A_24_P494658		AY026037	THC18351	2.03836	1.98E-10	2.03836	45376.36	93076.73
A_23_P15	NM_08061	Dual specif	DUSP15	2.03805	1.00E-13	2.03805	10218.96	20846.46
A_32_P9707		Unknown	A_32_P97	2.03776	2.13E-07	2.03776	8562.931	17218.95
A_24_P83	XM_37247	Homo sapi	XM_37247	2.03746	6.63E-19	2.03746	1341.813	2731.358
A_32_P61074		Unknown	A_32_P61	2.037	6.04E-06	2.037	267153.3	559397.2
A_24_P56	BM023388	Nuclear loc	NLF2	2.03671	1.87E-11	2.03671	2940.715	6035.53
A_24_P21	AF318321	Casein kin	CSNK1D	2.03623	1.71E-09	2.03623	512.7196	1018.657
A_24_P38	NM_01660	chromoson	FAM53C,C	2.0356	2.45E-12	2.0356	2674.651	5513.862
A_24_P733083		Unknown	A_24_P73	2.03541	1.48E-13	2.03541	1066.064	2161.344
A_24_P76	NM_00671	RNA-bindir	RNPS1,E5	2.03458	1.48E-07	2.03458	3972.688	8373.396
A_24_P887768		AJ004940	THC18437	2.03426	0.0003	2.03426	6345.146	13239.82
A_24_P40	NM_20745	FLJ35934	FLJ35934	2.03364	0.00002	2.03364	122615.1	262829.2
A_23_P10	NM_00299	Homo sapi	SDC4,SYN	2.03299	4.93E-14	2.03299	4099.636	8344.412
A_24_P10	NM_01536	proline rich	PRR5,PP6	2.03294	4.40E-08	2.03294	164060.8	340945.3
A_24_P834210		Unknown	A_24_P83	2.03247	4.45E-17	2.03247	4017.608	8128.582
A_24_P42107		Unknown	A_24_P42	2.03076	1.72E-06	2.03076	188217.8	394307.8
A_24_P752279		Unknown	A_24_P75	2.03014	2.25E-08	2.03014	34060.41	70420.52
A_32_P22	NM_01544	Ras-induce	RIS1,p40B	2.02908	3.76E-09	2.02908	203000.7	418730.2
A_32_P22	NM_00549	DnaJ (Hsp	DNAJB6,M	2.0281	1.86E-39	2.0281	10060.91	20451.71
A_23_P21	NM_08064	Homo sapi	TBX1,DGS	2.02749	5.29E-14	2.02749	850.8411	1707.352
A_24_P50	AK096179	Hypothetic	LOC28595	2.02705	2.91E-33	2.02705	7612.877	15457.72

A_23_P13	NM_00043	Homo sapi	NCF2,NOX	2.02251	0.00098	2.02251	178.572	350.1783
A_24_P17	NM_00261	peroxisome	PEX10,NA	2.02249	4.47E-14	2.02249	2168.818	4433.355
A_23_P98	NM_01223	Homo sapi	SIRT1,SIR	2.02126	7.44E-08	2.02126	1010.938	2029.068
A_23_P30	NM_17843	Homo sapi	LCE5A,LEI	2.02035	1.87E-06	2.02035	287431.4	592824.9
A_23_P25	NM_00523	Homo sapi	ETS2	2.01927	1.35E-32	2.01927	10669.45	21593.3
A_24_P27	NM_02481	hypothetica	BAALC,FL	2.01863	2.74E-07	2.01863	188300.2	389146.7
A_24_P37	NM_02493	hypothetica	FLJ12056	2.01709	4.86E-10	2.01709	8255.803	16893.48
A_23_P13	NM_19907	Homo sapi	C21orf58	2.01614	1.37E-07	2.01614	13546.86	27811.59
A_24_P932958		Unknown	A_24_P93	2.01398	0.00014	2.01398	1332.386	2533.772
A_23_P11	NM_01427	Homo sapi	ADAMTS7	2.01367	5.17E-08	2.01367	105672.9	216461.1
A_32_P8724		Unknown	THC18930	2.01346	4.41E-07	2.01346	106538.5	219014.2
A_23_P87	NM_00173	B-cell trans	BTG1	2.01327	7.75E-14	2.01327	1810.416	3654.642
A_24_P25	NM_03289	hypothetica	MGC14376	2.01309	1.35E-06	2.01309	18804.44	38603.28
A_23_P16	NM_00384	Homo sapi	TNFRSF1	2.01273	5.03E-07	2.01273	442.1876	853.2638
A_24_P44	NM_14505	Homo sapi	CDC42EP	2.01265	1.11E-06	2.01265	271011.9	568198.3
A_24_P29	NM_01894	cytochrome	CYCS,CYC	2.01234	0.00008	2.01234	267.3682	510.0447
A_24_P26	AL831836	Hypothetic	LOC28368	2.01141	1.45E-23	2.01141	1092.487	2192.787
A_24_P83	XM_29393	similar to R	LOC34571	2.01118	1.62E-14	2.01118	3336.313	6779.055
A_23_P25	NM_01334	replication	RPA4,HSU	2.01086	0.00001	2.01086	329.4243	630.6915
A_23_P11	NM_00624	Homo sapi	PRB3,G1,F	2.01066	1.41E-09	2.01066	13338.8	27374.07
A_23_P99	NM_03194	oculospani	TSPAN10,	2.00926	3.59E-07	2.00926	140044.5	290755.6
A_23_P52	NM_03095	likely ortho	NUAK2,SN	2.00832	2.27E-25	2.00832	1573.27	3153.462
A_23_P41	AB046797	Hypothetic	MGC40405	2.00549	1.31E-23	2.00549	1874.208	3761.086
A_32_P152999		Unknown	A_32_P15	2.00514	1.21E-07	2.00514	201171.7	416467.7
A_23_P10	NM_03258	Homo sapi	HES7	2.00315	5.10E-06	2.00315	262141.3	537819.1
A_23_P15	NM_00422	Interleukin	NK4	2.00315	4.39E-11	2.00315	1873.051	3775.858
A_23_P39	NM_00558	Homo sapi	LU,AU,BC	2.00251	9.82E-09	2.00251	40605.54	81815.86
A_24_P60	BM701491	Hypothetic	BM701491	2.00155	7.01E-11	2.00155	1575.164	3159.548
A_23_P14	NM_00681	Homo sapi	PNRC1,B4	2.00105	4.71E-08	2.00105	2247.635	4540.783
A_23_P21	NM_03256	esophagus	ECG2,ECF	1.99984	8.80E-14	1.99984	8255.525	16579.54
A_32_P59	NM_17361	hypothetica	FLJ35785	1.99964	0.00011	1.99964	275.0848	518.3904
A_24_P689119		Unknown	A_24_P68	1.99921	5.61E-06	1.99921	141399.8	293506.3
A_23_P25	NM_00361	Homo sapi	GALR3	1.99819	6.04E-08	1.99819	152136.6	310874.3
A_23_P26	NM_00396	Homo sapi	RHBDL1,R	1.99804	7.31E-06	1.99804	94287.23	191799.5
A_23_P10	BC050568	Meteorin, g	METRNL	1.99643	4.89E-14	1.99643	3934.017	7866.956
A_24_P853410		Unknown	A_24_P85	1.99584	3.84E-08	1.99584	17711.67	35943.85
A_23_P15	NM_00080	gastrin	GAST,GAS	1.99434	1.14E-10	1.99434	945.0817	1863.239
A_24_P36	BC028099	Homo sapi	BC028099	1.99398	4.07E-08	1.99398	47280.78	96000.05
A_24_P23	NM_17231	Homo sapi	KCNG1,K1	1.99362	2.63E-12	1.99362	6854.624	13673.96
A_24_P33	AK056484	LOC44251	LOC34028	1.99318	8.31E-14	1.99318	21615.37	43127.76
A_32_P10	AF130102	Homo sapi	PRO0522	1.99314	2.36E-12	1.99314	6729.67	13210.7
A_24_P24	BC014556	Hypothetic	FLJ35390	1.99093	3.15E-09	1.99093	49797	101237
A_24_P81	AK092421	Hypothetic	MGC3234	1.98896	1.55E-07	1.98896	61391.15	123819.5
A_23_P10	NM_00346	Chemokine	CXCR4	1.98832	3.12E-08	1.98832	681.3066	1311.761
A_24_P20	NM_01635	iroquois ho	IRX4	1.98644	1.77E-07	1.98644	10226.08	20584.05
A_32_P22	BF512678	Transcribe	PTPN4,PT	1.98387	2.96E-16	1.98387	18323.63	36600.77
A_24_P930551		AY080897	THC19769	1.98218	1.73E-13	1.98218	249614.5	498571.3
A_24_P73	NM_01837	hypothetica	FBXL8,FBI	1.97923	4.66E-21	1.97923	1572.808	3096.504
A_24_P89	BC016022	Similar to a	BC016022	1.97851	0.00087	1.97851	202.0861	382.0973
A_24_P823768		Unknown	A_24_P82	1.9773	7.28E-31	1.9773	1529.529	3025.102
A_24_P15	AB018318	ProSAPiP2	ProSAPiP2	1.97702	9.58E-08	1.97702	1626.959	3265.943
A_24_P25	NM_05716	Homo sapi	ARTN,EVN	1.97614	3.84E-11	1.97614	4639.054	9286.38

A_23_P19	NM_17504	Homo sapi	PILRB,FDF	1.97575	3.67E-20	1.97575	5829.139	11605.31
A_23_P26	NM_13426	Homo sapi	RORA,ROI	1.97543	0.00004	1.97543	229.8191	459.3054
A_32_P232327		Unknown	A_32_P23	1.97541	9.85E-13	1.97541	514.4067	1025.521
A_23_P13	NM_01615	Homo sapi	MMP17,MT	1.97525	4.19E-08	1.97525	42122.49	85192.29
A_24_P711809		Unknown	A_24_P71	1.97496	2.57E-07	1.97496	3138.209	6329.677
A_24_P63	NM_00624	protein phc	PPP1R2,IF	1.97342	1.67E-15	1.97342	660.306	1293.596
A_24_P24	NM_00468	Metastasis	MTA1	1.97283	9.05E-10	1.97283	22855.54	45285.54
A_24_P48	XM_37269	Homo sapi	XM_37269	1.97236	1.07E-07	1.97236	51672.51	102363.3
A_24_P10	NM_00447	forkhead b	FOXD1,FK	1.97227	2.58E-13	1.97227	3099.698	6110.299
A_23_P15	XM_37059	Homo sapi	LOC38773	1.97042	0.00004	1.97042	249884.3	503428.6
A_24_P90	NM_17361	hypothetica	FLJ36701	1.96989	0.00002	1.96989	228547	462712.4
A_23_P36	AF332224	Chromosor	CYorf15A	1.96968	5.29E-07	1.96968	349.9101	674.3265
A_32_P163215		Unknown	A_32_P16	1.96896	9.83E-13	1.96896	8822.449	17420.5
A_24_P71	XM_37288	Homo sapi	XM_37288	1.96895	3.17E-06	1.96895	172502.5	348777.3
A_32_P77759		ALU5_HUM	THC19702	1.96732	1.82E-07	1.96732	71499.21	142030.3
A_32_P52713		Unknown	THC19332	1.96631	0.00002	1.96631	291357.9	596634.4
A_24_P48	NM_00374	JAK bindin	SOCS1,JA	1.96594	6.96E-11	1.96594	597.2815	1152.306
A_23_P10	NM_03317	Homo sapi	DUX4	1.96557	2.85E-06	1.96557	284479.9	569326.6
A_32_P78	BM994509	Superoxide	SOD2	1.9648	1.07E-06	1.9648	426.6294	875.4474
A_23_P46	NM_01218	Homo sapi	FOXD3,HF	1.96461	1.00E-37	1.96461	3427.267	6732.412
A_24_P247273		Unknown	ENST0000	1.96357	7.32E-17	1.96357	10212.7	20070.95
A_24_P30	NM_01827	hypothetica	FLJ10925,	1.96246	8.64E-09	1.96246	12664.97	25109.4
A_23_P41	NM_01763	hypothetica	FLJ20035,	1.96222	4.15E-11	1.96222	774.0557	1499.764
A_24_P22	NM_20337	similar to z	LOC14780	1.96183	4.76E-16	1.96183	1890.334	3707.772
A_24_P31	NM_02212	MAWD bin	MAWBP,FI	1.96159	5.38E-09	1.96159	1096.499	2133.728
A_24_P315581		Unknown	A_24_P31	1.96105	7.68E-09	1.96105	56619.41	112595.6
A_23_P40	BC033237	Hypothetic	MGC45922	1.9595	1.41E-06	1.9595	51594.1	102681.4
A_24_P620521		Unknown	THC19990	1.95814	8.98E-16	1.95814	7737.576	15367.15
A_24_P417922		double hon	DUX1	1.95767	3.80E-06	1.95767	262441.1	524542.1
A_24_P11	AK027006	Trinucleotic	TNRC9	1.95682	8.56E-13	1.95682	8650.102	16949.51
A_23_P12	NM_00584	Homo sapi	SPRY2,hS	1.95654	4.00E-06	1.95654	486.0705	907.0031
A_23_P25	NM_00587	Homo sapi	RGS19,GA	1.95592	3.43E-13	1.95592	762.7934	1516.088
A_23_P88	NM_00420	synaptogyr	SYNGR3,M	1.9548	5.51E-10	1.9548	1204.44	2333.643
A_24_P26	AK092264	Ankyrin rep	ANKRD11	1.95374	3.19E-15	1.95374	1980.956	3868.827
A_24_P17	AB095932	KIAA2012	LOC33980	1.95356	3.94E-06	1.95356	79633.15	161405.8
A_24_P936836		putative AT	NP285481	1.95305	1.76E-29	1.95305	1169.587	2285.283
A_23_P25	NM_17745	Homo sapi	LYNX1,SLI	1.95304	1.25E-07	1.95304	107770.3	213180.3
A_23_P34	NM_00036	Homo sapi	TNNT2,CM	1.95174	5.47E-11	1.95174	566.3872	1087.745
A_24_P36	NM_00198	even-skipp	EVX1	1.95119	6.69E-08	1.95119	30728.12	61080.58
A_32_P44	AK091220	Homo sapi	HIST3H2B	1.95	0.00004	1.95	277.726	522.9025
A_32_P71864		Unknown	A_32_P71	1.95	2.96E-06	1.95	317231.4	635787.6
A_24_P23	NM_03326	winged hel	FOXQ1,HF	1.94842	3.42E-08	1.94842	166719.8	328997.7
A_23_P77	NM_02281	Homo sapi	MAP1LC3E	1.94594	3.52E-09	1.94594	1548.391	3017.178
A_32_P203585		Unknown	A_32_P20	1.94421	1.35E-06	1.94421	171706.2	341554.5
A_24_P349788		Unknown	ENST0000	1.94385	1.18E-06	1.94385	239346.4	473919.1
A_23_P21	NM_01611	NEDD8 ulti	NYREN18,	1.94187	3.32E-10	1.94187	3689.008	7276.128
A_24_P23	NM_17335	Homo sapi	SNF1LK,M	1.94139	2.19E-07	1.94139	164479.8	324338.5
A_23_P56	NM_00333	ubiquitin A-	UBA52,CE	1.93781	5.44E-32	1.93781	35441.68	68496.79
A_23_P15	NM_03318	Homo sapi	KRTAP2-4	1.93747	1.16E-09	1.93747	80260.61	157776.3
A_24_P654066		Unknown	A_24_P65	1.93412	3.29E-12	1.93412	5954.738	11430.11
A_24_P565503		Unknown	A_24_P56	1.93326	1.78E-13	1.93326	3309.712	6407.607
A_23_P16	NM_00233	low density	LRP3	1.93293	2.67E-07	1.93293	54160.46	106157.7

A_32_P101195		Unknown	A_32_P10	1.93279	1.82E-06	1.93279	268801.6	535722.1
A_24_P48	BM913108	Vacuolar p	VPS52	1.93231	2.89E-09	1.93231	13891.68	26903.88
A_32_P44628		Unknown	A_32_P44	1.93221	1.04E-08	1.93221	987.4341	1946.723
A_23_P16	NM_01660	Homo sapi	FAM53C,C	1.93122	3.21E-12	1.93122	1310.464	2520.266
A_24_P82	BX538238	Metastasis	MALAT1	1.93114	5.78E-16	1.93114	40955.2	78515.8
A_24_P91	X90978	Runt-relate	RUNX1	1.93094	5.69E-07	1.93094	6535.919	12748.21
A_32_P233211		Unknown	A_32_P23	1.93052	4.90E-06	1.93052	239202.2	472984.8
A_32_P70341		Unknown	THC18270	1.93019	1.96E-06	1.93019	318298.8	628253.8
A_24_P92	NM_18404	Homo sapi	ALDOA,AL	1.93017	8.61E-06	1.93017	247953.5	491384.6
A_32_P73	AK095260	Hypothetic	FLJ10769	1.92949	3.59E-08	1.92949	13641.17	26632.71
A_24_P21	NM_00516	aristaless (	PHOX2A,A	1.92653	1.66E-06	1.92653	224269.7	443375.1
A_24_P34	NM_00245	mucin 2, in	MUC2,MLF	1.9265	2.83E-11	1.9265	14566.23	28134.29
A_24_P118142		Unknown	A_24_P11	1.92572	7.29E-18	1.92572	33861.04	64994.81
A_32_P6233		Unknown	A_32_P62	1.92457	3.07E-11	1.92457	593.443	1125.026
A_24_P932736		Unknown	THC19585	1.92446	6.96E-07	1.92446	542.1746	1008.233
A_23_P78	NM_00047	Homo sapi	AMH,MIF,M	1.92182	2.87E-13	1.92182	876.8028	1663.981
A_24_P88	BC041970	Homo sapi	C9orf122	1.92144	5.85E-12	1.92144	2701.247	5137.168
A_32_P28	NM_00329	tropomyosi	TPM4	1.92094	1.29E-08	1.92094	8349.059	16197.8
A_32_P14	AW269579	Heat shock	HSPB1	1.9203	1.58E-12	1.9203	1228.732	2375.102
A_23_P15	NM_02211	Homo sapi	TSPYL2,CI	1.91974	2.82E-09	1.91974	19219.84	37384.35
A_23_P13	BX640843	Hypothetic	BX640843	1.91782	3.32E-36	1.91782	6471.599	12421.87
A_23_P11	NM_01907	Homo sapi	ING3,ING2	1.91552	3.09E-31	1.91552	1614.615	3091.008
A_24_P31	BC022826	Homo sapi	BC022826	1.91409	1.88E-07	1.91409	46757.85	90949.75
A_23_P15	NM_18178	Homo sapi	NCOA7,ES	1.91398	4.44E-07	1.91398	1934.401	3617.264
A_24_P36	NM_00397	sperm ass	SPAG9,HS	1.91126	1.20E-35	1.91126	1850.242	3534.963
A_32_P10	BX364765	Similar to D	BX364765	1.91048	0.00014	1.91048	541.7977	1112.797
A_23_P16	NM_01620	Homo sapi	AMOTL2,L	1.91047	5.42E-16	1.91047	2070.498	3965.967
A_24_P24	BC031319	Homo sapi	BC031319	1.91039	5.72E-09	1.91039	1171.082	2200.901
A_23_P14	AF131778	Discs, larg	DLGAP3	1.90978	4.51E-06	1.90978	134805.5	263704.9
A_24_P14	NM_00534	heat shock	HSPCA,HS	1.90833	3.32E-32	1.90833	137703	262227.3
A_24_P35	NM_05290	Homo sapi	SLC25A25	1.90789	3.57E-10	1.90789	2228.685	4261.329
A_24_P76210		TVMVRR p	THC19532	1.90636	1.40E-15	1.90636	6142.183	11657.69
A_24_P21	BC029861	Vacuolar p	VPS35	1.90575	1.73E-12	1.90575	2774.836	5296.75
A_24_P37	NM_01959	Homo sapi	LRRC8A,L	1.90409	7.61E-08	1.90409	425.7585	792.9012
A_23_P96	NM_00142	Homo sapi	ELF4,MEF	1.9038	0.00001	1.9038	274.1237	535.4807
A_23_P11	AB002301	Microtubule	MAST4	1.90238	2.13E-16	1.90238	663.0689	1256.144
A_24_P22	AK097425	Acyl-Coen	ACAD10	1.90232	5.77E-12	1.90232	7917.506	15020.98
A_24_P23	NM_01345	variable ch	VCX,VCX1	1.90162	0.00038	1.90162	298.3538	532.4934
A_24_P10	NM_14586	Homo sapi	KLK3,APS	1.90119	0.00014	1.90119	271118	526123.8
A_24_P39	NM_00196	eukaryotic	EIF5,EIF-5	1.89953	1.82E-17	1.89953	1372.586	2607.994
A_32_P10	BF084890	Trophoblas	TncRNA	1.89918	6.51E-06	1.89918	388.6965	772.221
A_23_P46	AK026882	Tripartite m	TRIM46	1.89898	1.05E-15	1.89898	2736.984	5168.192
A_32_P213543		Unknown	A_32_P21	1.89831	4.04E-28	1.89831	1277.68	2420.644
A_23_P86	NM_00656	Homo sapi	LBX1,HPX	1.89822	4.20E-07	1.89822	334313.3	647141.8
A_32_P14	AK023798	Hypothetic	LOC92249	1.89693	2.43E-07	1.89693	106391.6	202358.1
A_32_P170368		Unknown	A_32_P17	1.89662	0.00001	1.89662	344893.8	666571.8
A_23_P82	NM_00310	SRY (sex c	SOX4,EVI1	1.89572	2.01E-11	1.89572	607.3948	1163.61
A_24_P38	NM_20343	Homo sapi	IER5L,MG	1.89389	1.09E-16	1.89389	6907.356	13194.51
A_24_P32178		Unknown	ENST0000	1.8904	0.00001	1.8904	160872.4	314491.3
A_23_P20	NM_00395	Homo sapi	MAP3K14,	1.8903	1.65E-26	1.8903	1078.325	2037.708
A_32_P153916		AF254569	THC18279	1.88853	0.00008	1.88853	327.8767	588.7896
A_24_P35	NM_01494	KIAA0872	MON1B,HS	1.88831	1.58E-10	1.88831	41461.57	78433.1

A_23_P56	NM_20731	similar to a	LOC11271	1.88594	1.48E-07	1.88594	2853.074	5318.45
A_24_P62	NM_01784	hypothetica	CXorf17,O	1.88568	1.24E-10	1.88568	7608.715	14439.69
A_24_P641742		Unknown	A_24_P64	1.88528	2.16E-07	1.88528	390120.9	735488.6
A_24_P21	NM_00353	H3 histone	HIST1H3D	1.88513	1.43E-27	1.88513	15526.91	29360.37
A_23_P14	NM_00095	Homo sapi	PTGER4,E	1.88456	8.02E-06	1.88456	358.6108	700.4753
A_32_P14	BC043547	XK-related	UNQ2754	1.88226	1.10E-07	1.88226	22145.57	42433.41
A_32_P107797		Unknown	A_32_P10	1.88172	5.46E-14	1.88172	1001.437	1874.631
A_24_P18	NM_02040	protocadhe	PCDH9	1.88072	1.28E-13	1.88072	608.508	1144.306
A_23_P83	NM_01416	Homo sapi	MRPL18,H	1.88033	1.07E-40	1.88033	19882.82	37370.64
A_32_P38	NM_17491	Homo sapi	TCTE3	1.88004	1.06E-08	1.88004	14065.38	26873.54
A_24_P39	NM_13856	protein 7 tr	XTP7,MGC	1.87943	7.49E-07	1.87943	100799.1	191124.2
A_24_P307104		Unknown	A_24_P30	1.87898	3.39E-06	1.87898	602.6427	1085.259
A_23_P15	NM_00271	protein phc	PPP1R10,I	1.87898	1.20E-11	1.87898	4226.188	8008.48
A_32_P76566		Unknown	THC19641	1.8784	5.25E-06	1.8784	97896.36	186117.3
A_24_P38	NM_00091	oxytocin, p	OXT,OT,O	1.87837	4.57E-08	1.87837	10713.95	20351.02
A_24_P72	NM_00671	RNA-bindir	RNPS1,E5	1.87702	5.99E-07	1.87702	592.6044	1082.265
A_32_P11	NM_14588	Homo sapi	LRDD,PIDI	1.87645	3.14E-13	1.87645	1716.465	3218.305
A_24_P58	NM_02481	hypothetica	FLJ22955	1.87263	1.01E-10	1.87263	5467.828	10238.58
A_24_P82	BC016872	Centrosom	Cep72	1.87189	3.81E-13	1.87189	3724.088	6978.974
A_32_P112100		Unknown	A_32_P11	1.87132	1.92E-10	1.87132	358584.8	677023.3
A_23_P21	NM_00043	Homo sapi	PCSK1,PC	1.87115	2.76E-10	1.87115	586.5388	1083.713
A_23_P25	NM_00655	Homo sapi	KHDRBS3	1.86965	1.27E-15	1.86965	3282.911	6099.693
A_23_P10	NM_00145	Homo sapi	FHL2,DRA	1.86943	1.57E-10	1.86943	3450.474	6513.054
A_23_P74	NM_00394	selenium b	SELENBP	1.86916	1.97E-06	1.86916	371.702	676.4204
A_23_P15	NM_03319	Homo sapi	UCN2,SRF	1.86887	2.90E-07	1.86887	81065.39	155403.2
A_23_P16	NM_00045	Serine/thre	STK11	1.86788	2.06E-18	1.86788	4084.965	7620.793
A_23_P123645		Unknown	THC19529	1.86754	0.00006	1.86754	292.9155	532.049
A_32_P69456		Unknown	THC19372	1.86688	0.00008	1.86688	209180	409992.8
A_24_P93	BC010732	Hypothetic	LOC40426	1.86594	9.01E-17	1.86594	2976.655	5530.586
A_23_P42	NM_17851	hypothetica	LOC28384	1.86571	3.63E-13	1.86571	869.3141	1605.014
A_24_P33	AK091771	Centaurin,	CENTG3	1.86551	0.00003	1.86551	202197.6	383251.7
A_24_P24	NM_20742	Homo sapi	LOC39985	1.86312	3.33E-10	1.86312	16300.62	30683.98
A_24_P93	L24123	Homo sapi	NRF1	1.86262	7.02E-08	1.86262	310287.9	586959
A_23_P43	AK055561	Endoplasm	ERN1	1.86194	9.05E-07	1.86194	817.2515	1489.183
A_24_P59	BC063888	Homo sapi	BC063888	1.86125	4.04E-14	1.86125	4917.882	9144.322
A_24_P28	AL136789	Chromosor	DKFZp434	1.86044	3.83E-08	1.86044	12447.04	23477.23
A_23_P84	M94173	Calcium ch	CACNA1B	1.85912	5.22E-21	1.85912	3440.92	6377.936
A_24_P74	NM_15229	hypothetica	C1orf158,M	1.85834	4.00E-09	1.85834	14902.17	27965.36
A_23_P40	NM_00528	Homo sapi	GPR8	1.85593	2.37E-08	1.85593	1647.62	3042.34
A_23_P21	NM_00118	Homo sapi	BACH1	1.85583	5.10E-08	1.85583	604.2861	1096.665
A_23_P39	NM_00353	H4 histone	HIST1H4D	1.85417	4.54E-40	1.85417	15952.59	29594.23
A_23_P50	NM_03238	Homo sapi	C5orf4,FLJ	1.85319	2.77E-09	1.85319	2238.453	4141.156
A_23_P35	BC009198	RRN3 RNA	RRN3	1.85314	3.04E-06	1.85314	349.7683	638.596
A_23_P82	NM_01658	Homo sapi	NRN1,NRN	1.85281	1.71E-06	1.85281	15213.38	28816.82
A_24_P37	AK093505	CDNA FLJ	CXorf18,FL	1.85278	0.00009	1.85278	557.7608	978.7855
A_24_P828125		Unknown	A_24_P82	1.85132	9.88E-08	1.85132	56349.94	106325.6
A_24_P256155		Unknown	A_24_P25	1.84818	1.75E-07	1.84818	34674.02	64633.03
A_24_P23	NR_00004	Homo sapi	RNU12	1.84814	9.76E-10	1.84814	1552.544	2857.042
A_23_P38	NM_05824	Homo sapi	DNAJB6,M	1.84715	6.44E-08	1.84715	581.6012	1051.609
A_24_P668572		ALU5_HUM	THC19090	1.84665	0.00005	1.84665	285.2285	517.7147
A_24_P31	NM_01571	glioma tum	GLTSCR1	1.8461	6.58E-11	1.8461	2399.843	4425.769
A_24_P938465		Unknown	A_24_P93	1.84464	2.01E-12	1.84464	1577.315	2914.631

A_23_P58	NM_05300	TIGA1	TIGA1,MG	1.84455	3.89E-37	1.84455	24921.94	45904.75
A_32_P83	NM_14525	LOC12673	C1orf96,FL	1.84391	1.51E-17	1.84391	823.6077	1522.041
A_23_P71	NM_00130	Homo sapi	CLDN3,RV	1.84306	2.23E-08	1.84306	890.0569	1606.656
A_24_P41	AB014771	RasGEF d	RASGEF1I	1.84247	8.45E-06	1.84247	11263.52	21096.54
A_32_P162520		Unknown	A_32_P16	1.84163	3.81E-11	1.84163	2075.888	3848.062
A_23_P16	NM_01837	Receptor-i	RIF1	1.841	7.02E-07	1.841	577.8873	1034.068
A_24_P161655		Unknown	ENST0000	1.8408	1.16E-25	1.8408	8434.996	15589.17
A_32_P16	BG547692	Transcribe	BG547692	1.83989	3.10E-13	1.83989	3922.413	7214.488
A_23_P20	NM_00065	Homo sapi	BCL2,Bcl-2	1.83852	1.21E-07	1.83852	32238.67	59894.92
A_24_P41	NM_19427	Homo sapi	C14orf43,L	1.83841	5.47E-12	1.83841	9270.004	17195.94
A_24_P30	AK027130	Chromosor	C19orf33	1.83693	4.99E-08	1.83693	5237.688	9744.912
A_32_P138366		ALU4_HUM	THC19596	1.83534	4.96E-06	1.83534	20987.02	39014.68
A_24_P24	AL833358	DnaJ (Hsp	DNAJB12	1.83397	2.68E-12	1.83397	1048.134	1911.073
A_24_P20	NM_02472	MICAL-like	MICAL-L2	1.83359	5.66E-12	1.83359	1256.875	2294.491
A_23_P39	NM_00652	Homo sapi	TFPI2,PP5	1.83275	1.04E-21	1.83275	13579.61	25005.7
A_24_P58	NM_00381	tumor necr	TNFSF9,4-	1.83273	7.27E-27	1.83273	2328.742	4263.979
A_23_P53	NM_00173	Homo sapi	KLF5,CKLF	1.8327	1.26E-14	1.8327	2860.422	5297.715
A_23_P32	NM_00579	Homo sapi	DHRS2,HE	1.83266	1.33E-16	1.83266	893.1002	1643.975
A_32_P138586		Unknown	A_32_P13	1.83234	5.87E-08	1.83234	334949.9	627562.6
A_32_P73491		Unknown	A_32_P73	1.83175	6.38E-11	1.83175	3384.554	6215.213
A_23_P36	NM_00628	Homo sapi	THY1,CD9	1.83118	9.34E-08	1.83118	33763.88	61993.88
A_32_P13	XM_37947	Homo sapi	XM_37947	1.83005	4.04E-07	1.83005	7310.892	13615.86
A_24_P91	BC002724	Homo sapi	SNX5	1.82909	8.35E-18	1.82909	6290.82	11580.85
A_23_P11	NM_15233	hypothetica	MGC26885	1.82832	7.15E-07	1.82832	62582.73	115587.4
A_23_P80	NM_00108	Homo sapi	AADAC,DA	1.82811	4.02E-14	1.82811	1027.613	1889.115
A_24_P48	NM_01811	hypothetica	FLJ10490	1.82759	4.58E-07	1.82759	357783.8	663821.5
A_23_P39	NM_00346	Homo sapi	FZD5,HFZ	1.82588	2.36E-08	1.82588	1708.138	3132.538
A_23_P38	NM_15334	Homo sapi	C6orf141,M	1.82506	1.54E-19	1.82506	970.2642	1768.172
A_23_P17	NM_00546	Homo sapi	MAFB,KRM	1.82336	8.91E-14	1.82336	931.3635	1681.927
A_24_P13	AK096715	Annexin A8	ANXA8	1.8222	1.86E-06	1.8222	87919.48	162690.3
A_24_P58	BC041467	Hypothetic	LOC33921	1.82143	5.75E-06	1.82143	341.4186	612.5193
A_24_P38	NM_17368	hypothetica	MGC33584	1.82121	1.08E-10	1.82121	15847.81	29144.82
A_24_P29	NM_15260	Homo sapi	ZNF579,FL	1.82037	4.88E-06	1.82037	332250.2	621921.9
A_32_P83	NM_20143	Homo sapi	RTN3,HAP	1.81983	0.00015	1.81983	331203.3	619075.4
A_23_P33	NM_02432	hypothetica	MGC11256	1.81968	2.57E-09	1.81968	19958.41	36714.32
A_23_P13	NM_00142	Homo sapi	EN2	1.81926	3.55E-06	1.81926	143839.9	267641.1
A_24_P893130		AP003451	THC19578	1.817	4.12E-12	1.817	2604.054	4720.941
A_23_P10	NM_00239	mouse dou	MDM4,MD	1.81695	3.55E-08	1.81695	605.2246	1077.598
A_32_P19	NM_00534	heat shock	HSPCA,HS	1.81577	3.06E-15	1.81577	32877.53	59204.19
A_32_P82	AW002637	BTB (POZ)	BTBD14B	1.81541	3.02E-08	1.81541	20391.53	37290.14
A_32_P154079		AB028926	THC18921	1.81494	0.00001	1.81494	1441.131	2658.354
A_24_P35	NM_14501	Homo sapi	MRGPRF,I	1.81152	1.25E-07	1.81152	41038.65	74363.2
A_23_P14	NM_02117	Homo sapi	HES4	1.81112	1.04E-13	1.81112	8345.254	15255.77
A_23_P37	NM_17835	Homo sapi	LCE1D,LE	1.81086	0.00051	1.81086	313241.7	578059.1
A_24_P63	NM_00549	DnaJ (Hsp	DNAJB6,M	1.8102	1.46E-16	1.8102	2068.253	3753.974
A_32_P99715		AJ004940	THC18434	1.80951	8.09E-28	1.80951	1876.358	3396.437
A_24_P25	AF132202	Homo sapi	PRO1859	1.80716	0.00015	1.80716	488.7451	940.4607
A_23_P11	NM_18180	Homo sapi	BMP8A	1.80502	6.92E-07	1.80502	364576.9	667205.1
A_24_P92	AK127966	CDNA FLJ	AK127966	1.80322	1.19E-06	1.80322	588.5356	1030.174
A_23_P11	NM_00560	Homo sapi	NKG7,GIG	1.80297	1.02E-07	1.80297	1298.565	2312.543
A_24_P17	NM_00054	Ryanodine	RYR1	1.80286	0.00002	1.80286	393.6302	686.6953
A_23_P11	NM_00392	Homo sapi	BCL10,CL	1.80258	1.14E-11	1.80258	866.7514	1551.66

A_24_P281025	Unknown	ENST0000	1.80052	0.00001	1.80052	107177.1	195995.5
A_24_P152793	Unknown	ENST0000	1.80031	1.11E-21	1.80031	16513.77	29835.15
A_32_P12	NM_17506	Homo sapi	WBSCR19	1.79725	1.49E-09	1.79725	1317.241	2351.371
A_24_P10	NM_00541	solute carri	SLC20A1,F	1.79712	1.54E-10	1.79712	695.9806	1237.708
A_24_P91	L79986	Homo sapi	nuclear aut	1.79538	5.48E-09	1.79538	15742.69	28515.57
A_23_P43	NM_00052	keratin 14 (	KRT14,K14	1.79482	4.95E-10	1.79482	8817.992	15910.55
A_32_P90178	Unknown	A_32_P90	1.7943	0.00001	1.7943	157543.2	286552.6
A_32_P23	NM_02451	hypothetica	FAM65A,F	1.79394	1.70E-06	1.79394	15262.34	27624.61
A_23_P10	NM_01497	Homo sapi	MAST1,SA	1.79374	6.60E-20	1.79374	1600.677	2865.243
A_24_P72	XM_20893	similar to R	LOC28395	1.79307	1.86E-06	1.79307	8229.512	14971.38
A_32_P14	AF086205	Tuftelin 1	TUFT1	1.79274	0.00001	1.79274	308.9338	560.6491
A_24_P39	NM_00043	sialidase 1	NEU1,NEU	1.79256	6.63E-15	1.79256	1354.322	2419.765
A_24_P92	M14087	Lectin, gala	LGALS1	1.79255	3.97E-14	1.79255	3375.872	6077.052
A_32_P12	AI792678	qp41c06.y	AI792678	1.7923	0.00001	1.7923	10374.69	18848.39
A_24_P783679	Unknown	A_24_P78	1.79183	5.03E-10	1.79183	59543.37	105571.9
A_32_P99533	Unknown	A_32_P99	1.79165	2.16E-06	1.79165	970.4775	1670.453
A_24_P23	NM_00380	tumor necr	TNFSF14,I	1.78933	2.20E-08	1.78933	468.1459	826.934
A_23_P70	NM_00354	Homo sapi	HIST1H4L,	1.78833	1.72E-23	1.78833	52035.63	92920.84
A_24_P20	NM_17060	Homo sapi	MLL3,HALI	1.78668	1.19E-06	1.78668	374833.9	677440.4
A_24_P17	AF534825	Triggering	TREML4	1.78519	5.25E-06	1.78519	333.4379	591.2447
A_24_P23	NM_02109	peptide YY	PYY2	1.78506	5.56E-12	1.78506	10035.05	17976.29
A_23_P11	NM_00413	Homo sapi	GZMB,HLF	1.78496	6.46E-16	1.78496	863.0785	1523.352
A_24_P621434	HS7C_HUI	THC18728	1.7842	5.29E-24	1.7842	28657.39	51130.71
A_23_P84	NM_02407	hypothetica	SECISBP2	1.78261	8.57E-06	1.78261	411.2127	754.7745
A_32_P93036	Unknown	A_32_P93	1.78244	2.55E-11	1.78244	4981.196	8864.67
A_23_P55	NM_14590	Homo sapi	RIOK3,SUI	1.78099	6.89E-06	1.78099	513.324	884.4565
A_23_P50	NM_00414	Homo sapi	NDUFB7,B	1.77941	8.92E-09	1.77941	14602.32	26320.2
A_23_P14	NM_00667	Homo sapi	ARID5A,MI	1.77906	0.00003	1.77906	296.3632	522.2767
A_24_P16	NM_00354	H4 histone	HIST1H4B	1.77861	3.47E-24	1.77861	56635.13	100434.7
A_24_P28	AK095583	Similar to F	AK095583	1.77859	1.79E-06	1.77859	58465.97	106609.2
A_23_P13	NM_01791	Homo sapi	C19orf24,F	1.77845	4.35E-16	1.77845	4608.107	8184.485
A_24_P26	NM_00367	basic helix-	BHLHB2,D	1.77844	1.36E-09	1.77844	1176.044	2082.207
A_32_P15	AK125992	Similar to h	LOC44244	1.77706	8.21E-11	1.77706	6257.771	11102.83
A_24_P925559	Unknown	A_24_P92	1.77655	4.73E-08	1.77655	516.6631	901.2695
A_23_P88	NM_00506	Homo sapi	SEL1L,IBD	1.77655	3.36E-08	1.77655	1132.563	1999.907
A_23_P13	NM_17072	Homo sapi	KCNJ14,IR	1.77547	5.14E-11	1.77547	787.5029	1390.802
A_23_P16	NM_00543	FOS-like a	FOSL1,FR	1.77528	7.40E-08	1.77528	915.3016	1596.14
A_24_P57	NM_03058	collagen, ty	COL18A1,I	1.77513	2.63E-06	1.77513	760.2151	1315.435
A_23_P11	BX648200	DKFZP434	DKFZP434	1.77443	2.41E-09	1.77443	13801.32	24646.23
A_32_P10	BC056606	Similar to V	MGC5735	1.77429	2.58E-10	1.77429	4474.67	7987.868
A_24_P24	NM_00194	dentatorub	ATN1,B37,	1.77423	0.00001	1.77423	54874.46	98414.16
A_32_P90	BC013284	KIAA1545	KIAA1545	1.77356	0.00004	1.77356	351518.6	639857.4
A_23_P34	NM_00121	chromoson	C16orf3	1.77177	3.03E-06	1.77177	34521.77	62332.7
A_23_P16	NM_00534	heat shock	HSPCA,HS	1.77156	5.27E-14	1.77156	129602.4	228521.8
A_24_P41890	Glucose 6-	ENST0000	1.77153	8.46E-33	1.77153	5740.434	10163.88
A_23_P50	AB027121	Chromosor	C18orf12	1.77147	9.23E-08	1.77147	16285.8	28402.75
A_24_P23	NM_00674	Src-like-ad	SLA,SLA1,	1.77121	3.94E-08	1.77121	3290.603	5855.317
A_23_P88	NM_01506	Homo sapi	ZNF423,O	1.77114	2.74E-07	1.77114	442.7972	767.6469
A_24_P26	NM_00532	H1 histone	HIST1H1D	1.77046	4.07E-17	1.77046	7124.49	12702.14
A_24_P13	AF116713	Inter-alpha	ITIH1	1.7703	9.49E-06	1.7703	2396.084	4172.386
A_23_P15	NM_05285	Homo sapi	CREB3L1,	1.77015	0.00002	1.77015	22813.34	40719.16
A_23_P12	NM_13928	Homo sapi	ORMDL3	1.77	6.47E-07	1.77	22680.27	40862.17

A_32_P14	AL122093	MRNA; cDI	AL122093	1.76928	5.87E-22	1.76928	6190.172	10918.77
A_24_P16	NM_18262	Hypothetic	FLJ36766	1.76914	0.00007	1.76914	74573.14	135299.1
A_23_P47546		Unknown	A_23_P47	1.76884	0.0006	1.76884	298.4403	504.4498
A_23_P26	NM_01509	PI-3-kinase	SMG1	1.76875	2.85E-09	1.76875	1820.57	3241.305
A_24_P492865		Unknown	A_24_P49	1.76854	6.47E-16	1.76854	1124.124	1979.923
A_24_P101879		Unknown	A_24_P10	1.76853	2.06E-09	1.76853	23524.82	41899.19
A_24_P127517		Unknown	ENST0000	1.7674	6.73E-09	1.7674	3958.063	6982
A_32_P231880		Unknown	A_32_P23	1.76722	0.00001	1.76722	388102.7	691965.9
A_24_P70	AK125680	ARHGEF5	FLJ43692	1.76653	0.0005	1.76653	277.8416	474.967
A_23_P14	AK027130	Chromosor	C19orf33	1.76651	0.00081	1.76651	417.3324	687.2471
A_23_P32	NM_00439	Homo sapi	DDX10,HR	1.76632	7.99E-09	1.76632	9738.824	17490.54
A_23_P14	NM_08059	Homo sapi	HIST1H2B	1.76552	1.84E-17	1.76552	13904	24713.71
A_23_P10	NM_00178	Homo sapi	CD79A,IG	1.76385	1.10E-06	1.76385	31558.43	56235.19
A_24_P40	NM_02515	hypothetic	FLJ21477	1.76338	0.00003	1.76338	127114.6	224806.8
A_24_P30	NM_01497	KIAA1049	KIAA1049,	1.76333	2.12E-17	1.76333	5041.971	8844.782
A_23_P16	NM_03141	Homo sapi	GABARAP	1.76332	8.57E-11	1.76332	682.5947	1194.471
A_23_P14	NM_00511	Homo sapi	GFPT2,GF	1.76236	1.80E-11	1.76236	669.939	1174.65
A_24_P349783		Unknown	ENST0000	1.76061	9.49E-12	1.76061	6607.833	11634.08
A_24_P68	BX640843	Hypothetic	BX640843	1.75789	1.22E-19	1.75789	9623.644	17010.74
A_24_P40	NM_01742	cat eye syr	CECR1,ID	1.75769	7.84E-10	1.75769	10844.39	19068.23
A_24_P273214		Unknown	A_24_P27	1.75703	0.00057	1.75703	178736.5	319337
A_23_P49	NM_01663	Homo sapi	TNFRSF12	1.75693	3.02E-17	1.75693	6711.157	11748.55
A_23_P38	NM_00555	Homo sapi	KRT7,K7,C	1.75656	5.97E-17	1.75656	8033.324	14031.33
A_32_P23	NM_18107	88-kDa gol	GOLGA8A	1.7536	0.00033	1.7536	466.1008	770.4462
A_24_P14	NM_02513	hypothetic	OPA3,MG	1.7524	1.69E-10	1.7524	5872.451	10322.66
A_23_P12	NM_01694	Chromosor	C6orf48	1.75222	3.10E-23	1.75222	34487.8	60249.76
A_32_P74243		Unknown	A_32_P74	1.74952	1.65E-08	1.74952	514.5147	888.4696
A_23_P53	NM_00557	Homo sapi	LMO2,TTG	1.74883	6.70E-12	1.74883	280.7569	497.1944
A_23_P67	NM_13833	hypothetic	SBBI54	1.74676	4.57E-16	1.74676	5185.506	9035.489
A_24_P19	NM_18256	hypothetic	MGC29814	1.74648	2.70E-10	1.74648	3904.188	6882.943
A_24_P341603		Unknown	ENST0000	1.74553	5.08E-06	1.74553	76458.03	135174.7
A_32_P56	BE068542	MR1-BT03	BE068542	1.74539	0.00066	1.74539	333271.3	590975
A_24_P630916		ALU6_HUM	THC19031	1.74452	5.56E-06	1.74452	803.2905	1432.856
A_24_P366495		Unknown	ENST0000	1.7439	8.98E-13	1.7439	7515.464	13224.45
A_24_P39	BC044311	Hypothetic	LOC33934	1.74374	0.00003	1.74374	144369.2	255980
A_23_P41	NM_17562	Homo sapi	MT1J,MT1	1.74371	8.36E-16	1.74371	4926.275	8598.969
A_24_P714750		Unknown	A_24_P71	1.74339	5.37E-10	1.74339	19980.3	35071.23
A_24_P58	AY358924	KIAA1201	KIAA1201	1.74323	4.36E-08	1.74323	1056.771	1824.741
A_24_P418176		p15 protein	NP106131	1.7423	5.20E-12	1.7423	2128.474	3711.735
A_23_P20	NM_01776	Homo sapi	CASZ1,ZN	1.73971	9.70E-09	1.73971	457.0502	795.5704
A_32_P13	XM_08693	Homo sapi	ENST0000	1.73938	2.28E-35	1.73938	4480.78	7792.341
A_23_P10	NM_01865	golgin-like	GOLGA6,G	1.73901	3.46E-09	1.73901	540.9395	936.8613
A_24_P655888		ALU8_HUM	THC19533	1.73882	0.00077	1.73882	238.3788	411.5982
A_23_P30	NM_00424	Jumonji do	JMJD1C	1.7381	0.00029	1.7381	335.3301	561.6624
A_24_P153002		Unknown	A_24_P15	1.73792	0.00033	1.73792	258139.1	455600.5
A_32_P56890		AP002913	THC18401	1.7378	6.10E-11	1.7378	3167.751	5511.876
A_23_P12	NM_00190	Homo sapi	CTH,MGC	1.73692	5.86E-12	1.73692	3849.326	6752.691
A_24_P14	AK055956	FLJ31394	FLJ31394	1.73662	1.62E-15	1.73662	838.157	1448.717
A_32_P44	NM_15247	hypothetic	FLJ32214	1.73636	7.23E-07	1.73636	5068.734	8919.275
A_23_P37	NM_00057	Homo sapi	DAF,CR,T	1.73472	4.54E-07	1.73472	2670.001	4683.93
A_23_P30	AK094772	Homo sapi	AK094772	1.73455	0.00001	1.73455	374.2622	636.7108
A_24_P643376		AL935260	THC18081	1.73432	1.35E-11	1.73432	6699.58	11458.53

A_23_P58	NM_00598	Homo sapi	S100P,MIC	1.73355	1.63E-17	1.73355	105441.4	184031.8
A_24_P341616		Unknown	A_24_P34	1.73325	5.11E-08	1.73325	16424.21	28558.27
A_32_P20	NM_18172	Homo sapi	EFHA2	1.73221	5.98E-09	1.73221	534.929	924.7618
A_24_P92	AF159295	Homo sapi	AF159295	1.73047	3.28E-14	1.73047	391395.8	680170.4
A_24_P21	NM_03194	oculospani	TSPAN10,	1.72989	3.54E-11	1.72989	11257.21	19462.44
A_23_P30	NM_02124	Homo sapi	PRM3	1.72812	1.73E-16	1.72812	2115.379	3615.486
A_23_P10	NM_15247	Homo sapi	ZNF565,FL	1.72802	5.28E-19	1.72802	1007.628	1741.266
A_23_P56	NM_00566	Homo sapi	RNF103,KI	1.72761	7.23E-06	1.72761	1057.743	1807.446
A_23_P15	NM_00469	Homo sapi	SLC16A6,M	1.72739	0.00012	1.72739	337.04	568.1677
A_24_P50601		Unknown	ENST0000	1.7269	5.80E-06	1.7269	58177.5	101252.1
A_23_P25	NM_00084	Homo sapi	GSTA3,GT	1.72655	0.00011	1.72655	386.8362	694.65
A_23_P37	NM_00072	Homo sapi	CACNB3,C	1.72639	1.49E-07	1.72639	14411.67	24942.58
A_32_P22	BC047717	Homo sapi	BC047717	1.72629	4.27E-09	1.72629	7523.727	12977.53
A_23_P14	NM_01660	Homo sapi	CCR10,GP	1.72585	1.73E-16	1.72585	1124.322	1930.676
A_24_P264416		Unknown	A_24_P26	1.72582	1.05E-26	1.72582	9605.574	16610.11
A_23_P48	NM_18290	Homo sapi	DHRS2,HE	1.72554	1.02E-09	1.72554	949.3535	1621.361
A_24_P931711		Unknown	A_24_P93	1.72516	2.71E-10	1.72516	3837.72	6639.48
A_32_P156776		Unknown	A_32_P15	1.72481	0.0001	1.72481	21230.27	37822.39
A_23_P46	NM_00517	Homo sapi	BCL3,BCL	1.72331	5.40E-25	1.72331	2594.084	4469.369
A_32_P159939		Unknown	THC19593	1.7233	1.01E-06	1.7233	671.1334	1130.488
A_24_P11	NM_00130	claudin 4	CLDN4,CP	1.72173	2.93E-09	1.72173	710.0301	1234.943
A_24_P31	NM_01656	BM88 antig	BM88,MGC	1.72161	2.58E-10	1.72161	33610.81	58037.24
A_23_P20	NM_01663	ADP-ribosy	LOC51326	1.72141	4.51E-06	1.72141	471.6593	794.9623
A_23_P60	NM_13902	Homo sapi	MAPK15,E	1.72078	4.29E-07	1.72078	603.6622	1013.666
A_24_P37	NM_13838	hypothetica	LOC92154	1.72045	2.43E-09	1.72045	16233.7	27920.79
A_32_P11	BC063309	Golgi autoa	LOC39053	1.72024	0.00011	1.72024	388.9814	646.8189
A_32_P33	NM_01337	Homo sapi	ZDHHC8,Z	1.71808	6.04E-10	1.71808	6745.624	11630.28
A_23_P33	NM_02214	Homo sapi	ELSPBP1,	1.71803	0.00001	1.71803	517.5953	858.345
A_32_P113533		Unknown	A_32_P11	1.71709	2.52E-35	1.71709	7619.345	13083.56
A_24_P11	BC004792	Homo sapi	BC004792	1.71707	6.98E-15	1.71707	1818.03	3114.528
A_24_P290068		Unknown	A_24_P29	1.71653	9.27E-08	1.71653	23242.07	39992.25
A_32_P160200		AF312770	THC19589	1.71569	0.00018	1.71569	368519.6	647478.9
A_23_P21	AF272774	Coagulatio	F7	1.71562	1.09E-10	1.71562	17279.26	29635.96
A_24_P18	BM472535	KIAA0826	KIAA0826	1.71517	4.13E-06	1.71517	386.7158	662.2233
A_32_P82	NM_17856	Homo sapi	AGBL3,FL	1.71371	8.47E-06	1.71371	5916.224	10138.68
A_23_P90	NM_00716	Homo sapi	SF3A2,PR	1.71216	0.00005	1.71216	125002.6	217065.2
A_24_P32	AB040899	Homo sapi	KIAA1466	1.71187	7.16E-11	1.71187	1298.098	2197.97
A_24_P92	NM_00538	Natural kill	NKTR	1.71148	3.12E-20	1.71148	3182.139	5456.61
A_24_P12	NM_00595	metallothio	MT1X,MT1	1.71119	2.15E-34	1.71119	4429.966	7580.865
A_24_P10	NM_00508	X-box bind	XBP1,XBP	1.71108	0.00003	1.71108	363.4069	612.4688
A_24_P34	NM_18258	Homo sapi	CYB561D1	1.70977	0.0001	1.70977	10189.2	17785.96
A_32_P85	NM_17361	hypothetica	FLJ35785	1.70964	0.00002	1.70964	355.4567	607.7465
A_24_P33	NM_00458	Rab geran	RABGGTB	1.70931	2.43E-35	1.70931	5478.331	9364.287
A_24_P68	NM_02100	Homo sapi	UBC,HMG	1.70913	4.09E-15	1.70913	420157.2	718731.1
A_24_P41	NM_00542	tumor prote	TP73,P73	1.70896	5.38E-08	1.70896	811.9373	1362.385
A_32_P61	AK055160	CDNA FLJ	AK055160	1.70726	4.31E-17	1.70726	1315.979	2239.477
A_24_P33	NM_18257	hypothetica	THEM5,FL	1.70644	0.00002	1.70644	65670.05	112426.6
A_24_P25	AK090467	Hypothetic	LOC28454	1.70585	0.00023	1.70585	193061.8	339719.1
A_23_P38	NM_01656	Homo sapi	TBX3,UMS	1.70457	0.0007	1.70457	402.7252	648.3118
A_24_P375962		Unknown	A_24_P37	1.70446	4.72E-09	1.70446	562.2013	952.4361
A_32_P39	BC030567	Deleted in	1-Dec	1.70358	0.00072	1.70358	444.3903	713.2407
A_23_P13	NM_01212	cysteine ar	CHORDC1	1.70357	1.56E-22	1.70357	4696.575	8020.739

A_24_P82	NM_19918	Homo sapi	NPM1,B23	1.70241	6.34E-08	1.70241	28225.61	48341.02
A_23_P30	NM_00595	Homo sapi	MT1X,MT1	1.70136	1.42E-30	1.70136	4656.068	7921.789
A_23_P74	NM_01571	putative lyr	G0S2,RP1	1.70124	1.46E-09	1.70124	10499.42	18092.81
A_32_P177685		Unknown	A_32_P17	1.7006	0.00006	1.7006	565.1531	924.7791
A_24_P35	AK074630	PTK2B pro	PTK2B	1.69992	0.00077	1.69992	571.2262	916.1901
A_24_P36	NM_00595	metallothio	MT2A,MT2	1.69796	8.28E-14	1.69796	20139.34	34497.41
A_24_P29	NM_18257	hypothetica	FLJ36070,	1.69761	3.93E-17	1.69761	4219.743	7169.746
A_24_P94	AB014525	Amyotroph	ALS4	1.6975	0.00002	1.6975	1363.92	2313.338
A_24_P38	BC030084	MRNA; cDI	BC030084	1.69638	0.00006	1.69638	4041.485	6937.065
A_24_P19	NM_02431	hypothetica	PLEKHF1,	1.69624	3.36E-09	1.69624	3047.995	5207.248
A_24_P93321		Unknown	A_24_P93	1.69524	1.05E-18	1.69524	2705.482	4594.476
A_24_P20	NM_17854	hypothetica	LOC33948	1.69472	4.30E-15	1.69472	1241.655	2094.679
A_23_P30	NM_01223	Homo sapi	RYBP,AAF	1.69442	1.83E-09	1.69442	1567.974	2652.046
A_32_P13	BC042539	Homo sapi	LOC44242	1.69395	1.67E-07	1.69395	6576.597	11224.28
A_32_P65934		Unknown	A_32_P65	1.69263	6.40E-10	1.69263	1969.878	3304.94
A_23_P14	NM_03130	Homo sapi	SCRT1,SC	1.69133	7.08E-11	1.69133	518.8932	877.7007
A_24_P45	AK126267	Chromosor	C9orf111	1.69032	1.63E-19	1.69032	1445.582	2436.977
A_24_P879135		Unknown	A_24_P87	1.69031	1.97E-07	1.69031	1658.449	2750.533
A_23_P21	NM_03348	Homo sapi	CDC2L1,p	1.69015	2.90E-25	1.69015	7848.321	13289.18
A_24_P493005		Unknown	A_24_P49	1.68987	0.00003	1.68987	886.8971	1461.852
A_23_P15	NM_15268	hypothetica	MGC9712,	1.68935	6.84E-15	1.68935	816.7074	1381.302
A_24_P25	NM_17569	Homo sapi	SSX2,SSX	1.68931	2.49E-07	1.68931	2127.896	3591.013
A_23_P25	NM_01763	collaborate	CARF,FLJ	1.68927	2.30E-09	1.68927	1730.166	2922.222
A_24_P40	NM_18279	Homo sapi	PBEF1,PB	1.68831	2.40E-26	1.68831	5877.438	9921.219
A_24_P30	NM_01910	tenascin XI	TNXB,XB,T	1.68781	2.79E-06	1.68781	445.6738	763.6249
A_24_P92	AK055915	CDC42 eff	CDC42EP	1.68769	0.00009	1.68769	352.4959	585.351
A_24_P38	NM_02487	hypothetica	FLJ12748	1.68549	4.95E-12	1.68549	3941.395	6616.901
A_23_P50	NM_01393	Homo sapi	OR10H2	1.68471	1.07E-08	1.68471	17310.85	29764.69
A_24_P18	BC011805	Homo sapi	EDEM1,ED	1.68435	6.80E-08	1.68435	18495.5	31223.29
A_24_P792130		AF071172	THC19993	1.68435	4.10E-06	1.68435	583.2106	962.3237
A_23_P30	NM_02012	Homo sapi	SPHK2	1.6842	5.88E-09	1.6842	8291.163	14049.86
A_24_P77	BM687208	Transcribe	BM687208	1.68336	2.66E-07	1.68336	22002.52	37383.78
A_24_P187539		Unknown	A_24_P18	1.68199	2.76E-07	1.68199	410.1383	690.8187
A_24_P91	AI337101	qx88c05.x1	AI337101	1.68031	2.86E-09	1.68031	6154.772	10288.15
A_24_P655735		Unknown	A_24_P65	1.67896	4.42E-09	1.67896	673.2156	1143.376
A_23_P25	NM_08086	SPRY dom	SPSB4,SS	1.67789	0.00003	1.67789	72834.09	125342.9
A_23_P68	NM_02216	Homo sapi	IFIH1,Hlcd	1.6772	6.99E-07	1.6772	745.5955	1218.574
A_23_P20	NM_03277	Homo sapi	ZNF503,M	1.67664	2.13E-09	1.67664	1258.27	2095.861
A_23_P12	NM_00508	Homo sapi	XBP1,XBP	1.67643	3.71E-06	1.67643	13692.88	23141.87
A_23_P31	NM_00335	UDP-gluco	UGCG,GC	1.67587	1.32E-06	1.67587	512.9683	848.0814
A_24_P21	AK095727	CDNA FLJ	AK095727	1.67565	0.00002	1.67565	2523.279	4277.566
A_24_P170309		Unknown	A_24_P17	1.67537	1.58E-24	1.67537	2864.983	4807.143
A_32_P46	NM_02449	Homo sapi	C14orf4,IR	1.67472	4.55E-18	1.67472	2114.972	3555.863
A_23_P46	NM_00155	Homo sapi	CYR61,CC	1.67422	1.10E-08	1.67422	1261.304	2091.123
A_24_P36	NM_17586	Homo sapi	ARID1B,6A	1.67356	4.60E-10	1.67356	8612.063	14472.31
A_23_P17	NM_03281	Homo sapi	ZNF341	1.67287	4.49E-26	1.67287	2830.941	4730.086
A_32_P207789		Unknown	A_32_P20	1.67231	1.96E-09	1.67231	1103.013	1829.343
A_23_P33	NM_01908	Homo sapi	FBXL19,Fb	1.6712	0.00016	1.6712	11863.4	19557.16
A_24_P79	NM_03325	G-protein g	GNG8	1.67076	1.35E-06	1.67076	1471.177	2429.594
A_23_P10	NM_03220	hypothetica	FLJ21742	1.67035	4.07E-10	1.67035	7850.135	13081.41
A_23_P11902		Unknown	A_23_P11	1.6701	4.46E-13	1.6701	1246.465	2067.771
A_23_P24	NM_00105	Homo sapi	TACR2,SK	1.66943	3.63E-11	1.66943	4148.706	6912.497

A_23_P46	NM_03344	elastase 2/	ELA2A,ELA	1.66927	2.65E-09	1.66927	1609.426	2611.47
A_23_P32	NM_01446	RNA-bindir	CSDC2,PIF	1.66926	2.16E-06	1.66926	12020.92	20097.53
A_23_P11	NM_00704	Homo sapi	KATNA1	1.66873	9.35E-06	1.66873	2811.781	4759.218
A_23_P20	NM_02076	CASK inter	CASKIN1	1.66866	8.96E-08	1.66866	424712.1	710422.6
A_32_P54	BX104097	TBC1 dom	TBC1D14	1.66799	1.04E-07	1.66799	5350.321	8684.485
A_24_P25	NM_00650	myelin-ass	MOBP,MG	1.66736	6.52E-09	1.66736	3087.144	5082.984
A_23_P11	NM_01239	Homo sapi	SPDEF,PD	1.66687	1.06E-07	1.66687	2187.141	3628.939
A_23_P31	NM_01778	Homo sapi	CYP2W1,M	1.66562	1.07E-08	1.66562	2492.434	4172.398
A_23_P11	NM_00192	Homo sapi	DF,ADN	1.66549	3.25E-25	1.66549	13854.69	23034.86
A_24_P30	NM_03208	protocadhe	PCDHGA7	1.665	2.32E-18	1.665	1317.452	2187.358
A_24_P357836		Unknown	ENST0000	1.66487	3.69E-17	1.66487	4241.514	7043.78
A_24_P14	NM_00706	CDC42 eff	CDC42EP	1.66438	0.00001	1.66438	23675.11	39517.04
A_24_P39	NM_13873	Homo sapi	GNAO1,GN	1.66357	1.52E-21	1.66357	1540.504	2558.367
A_24_P40	AK091259	CDNA clon	AK091259	1.6634	0.00001	1.6634	34390.53	58028.12
A_23_P170719		Unknown	A_23_P17	1.66314	0.0001	1.66314	403.7459	652.5491
A_32_P76	NM_00215	Heat shock	HSPD1	1.66255	1.05E-16	1.66255	4600.931	7613.054
A_23_P58	AK127224	Nedd4 binc	N4BP3	1.66205	0.00004	1.66205	462.266	744.2363
A_24_P68	BX647769	Ankyrin rep	UNQ2430	1.66201	1.68E-20	1.66201	1769.125	2939.871
A_24_P35	NM_02095	Homo sapi	KIAA1618	1.66159	4.29E-12	1.66159	1971.351	3278.173
A_23_P36	NM_01395	neuregulin	NRG1,GGF	1.66133	0.00001	1.66133	516.8299	879.5745
A_23_P41	NM_03209	protocadhe	PCDHGB2	1.66092	2.72E-07	1.66092	375.2596	626.6127
A_24_P493646		Unknown	A_24_P49	1.66072	3.66E-07	1.66072	6586.88	11055.44
A_23_P12	NM_17073	Brain-deriv	BDNF	1.65901	0.00077	1.65901	295.9307	482.879
A_23_P40	NM_17213	Homo sapi	IL28A,IFNL	1.65875	2.83E-09	1.65875	1403.173	2289.929
A_23_P14	NM_00193	Homo sapi	MPP3,DLG	1.65769	2.66E-06	1.65769	843.9365	1375.98
A_24_P35	NM_00409	ephrin-B2	EFNB2,HT	1.65726	8.39E-07	1.65726	3025.598	5019.344
A_32_P126647		Unknown	THC19266	1.65551	0.00062	1.65551	389.7847	617.543
A_24_P40	NM_02255	Major histo	HLA-DRB1	1.65484	1.46E-12	1.65484	1095.371	1821.447
A_23_P16	NM_18279	Homo sapi	NPM2,MG	1.65434	4.30E-07	1.65434	443.6526	730.1873
A_32_P49	BC036909	Hypothetic	LOC28488	1.65351	3.23E-28	1.65351	3931.505	6502.712
A_24_P27	NM_00547	hyperpolari	HCN4	1.65338	9.38E-06	1.65338	419238.2	696954.6
A_23_P71	NM_01447	serine prot	SPINK4,PE	1.65281	1.41E-12	1.65281	2610.012	4323.498
A_23_P47	NM_00647	PCAF assc	TAF6L,PAF	1.65267	7.66E-09	1.65267	560.0465	920.3391
A_23_P68	NM_01598	Homo sapi	ANGPT4,A	1.65212	2.82E-10	1.65212	1144.058	1860.189
A_24_P238769		Unknown	A_24_P23	1.65123	8.89E-08	1.65123	13786.22	22758.63
A_23_P42	X97261	Metallothio	MT1L	1.6511	2.59E-10	1.6511	4820.486	8025.845
A_23_P67	NM_00712	nuclear rec	NR1H2,NE	1.65095	2.20E-11	1.65095	6973.656	11548.05
A_24_P35	AY462868	Protein exp	POTE2	1.65032	0.00039	1.65032	331.5079	535.3763
A_24_P15	AK021860	Homo sapi	FBXO17,F	1.65004	0.00002	1.65004	60849.87	101605
A_32_P12	BC031698	Homo sapi	BC031698	1.64814	1.99E-09	1.64814	1354.914	2227.812
A_24_P23	NM_00463	RAB7, mer	RAB7,PSN	1.6459	3.35E-13	1.6459	2921.269	4816.824
A_32_P187975		U39779 be	THC19107	1.64573	0.00002	1.64573	1497.912	2500.028
A_23_P13	NM_01324	Homo sapi	CLCF1,CL	1.64508	3.90E-07	1.64508	592.415	952.9006
A_23_P13	NM_01400	Homo sapi	ZNF297B,Z	1.64419	3.13E-10	1.64419	1040.306	1722.808
A_32_P43911		Unknown	THC20047	1.6434	0.00003	1.6434	446391.8	733599.8
A_32_P9107		Unknown	A_32_P91	1.64323	5.91E-19	1.64323	1534.153	2515.2
A_24_P40	NM_00040	excision re	ERCC2,EM	1.643	1.13E-06	1.643	12833.13	20951.46
A_24_P741378		Unknown	THC18086	1.64281	2.11E-22	1.64281	1751.061	2878.068
A_24_P31	NM_00588	p21(CDKN	PAK4	1.64266	6.15E-10	1.64266	2790.704	4647.885
A_24_P27	NM_20140	deubiquitin	DUB3	1.64263	1.92E-10	1.64263	3540.065	5822.652
A_32_P73	XM_37903	hypothetica	LOC40089	1.64234	1.86E-11	1.64234	783.8201	1284.769
A_32_P12	BQ935463	AGENCOL	BQ935463	1.64205	8.59E-10	1.64205	807.8522	1338.163

A_23_P78	NM_00705	Homo sapi	SFRS16,C	1.6415	9.39E-11	1.6415	4572.328	7552.167
A_23_P14	NM_00334	ubiquitin-co	UBE2H,UB	1.63981	5.10E-09	1.63981	1450.333	2365.565
A_23_P25	NM_00155	Homo sapi	IFRD1,PC4	1.63971	1.04E-16	1.63971	4505.761	7415.334
A_24_P160868		Unknown	ENST0000	1.63895	3.61E-14	1.63895	877.3138	1437.497
A_23_P34	NM_00525	Homo sapi	FOSL2,FR	1.63893	4.65E-09	1.63893	4078.263	6691.713
A_23_P369966		Unknown	THC20015	1.63818	3.51E-07	1.63818	787.8403	1329.068
A_24_P41	NM_19807	golgi autoa	FLJ40113,	1.63748	7.87E-22	1.63748	2897.573	4739.034
A_24_P22	NM_00549	RAN bindir	RANBP9,R	1.63728	0.0002	1.63728	347.1503	558.5787
A_32_P185089		Unknown	A_32_P18	1.63707	2.40E-08	1.63707	11578.34	18867.42
A_24_P68	BU738278	Transcribe	BU738278	1.6367	0.00002	1.6367	1014.241	1625.396
A_24_P41	NM_20330	Homo sapi	FBXO33,F	1.63624	0.00009	1.63624	906.4568	1454.659
A_24_P84268		Unknown	A_24_P84	1.63451	3.77E-06	1.63451	2736.669	4484.035
A_23_P99	NM_00492	Homo sapi	ZFP36L1,E	1.63434	1.62E-07	1.63434	3546.384	5837.304
A_23_P14	NM_01757	G protein-c	MKNK2,M	1.63416	5.49E-11	1.63416	1445.73	2354.7
A_24_P78	NM_03226	hypothetica	DKFZp434	1.63374	4.61E-10	1.63374	727.1415	1184.8
A_24_P11	AB075851	Similar to j	KIAA1971	1.63366	1.02E-09	1.63366	1340.311	2188.723
A_23_P53	NM_00055	Homo sapi	HBG1,HB	1.6333	0.00054	1.6333	297.2546	481.4501
A_23_P98	NM_03093	unc93 (C.e	UNC93B1,	1.63314	7.17E-09	1.63314	1329.003	2152.737
A_24_P81	BC041882	Activating t	ATF7IP2	1.63296	6.61E-06	1.63296	499.8285	805.444
A_23_P20	NM_00454	Homo sapi	NDUFA3,B	1.63142	2.18E-06	1.63142	21436.77	35409.49
A_32_P15	NM_01226	heparan su	HS2ST1,F	1.63098	0.00004	1.63098	449721.9	733488.3
A_23_P27	NM_00333	Homo sapi	TYROBP,D	1.62991	1.30E-13	1.62991	3785.99	6160.77
A_24_P51	NM_17490	Homo sapi	C3orf6,YM	1.62967	4.78E-07	1.62967	1442.771	2342.082
A_23_P35	NM_00180	Homo sapi	CEBPG,G	1.62962	4.28E-11	1.62962	4096.925	6720.336
A_23_P25	NM_03062	Homo sapi	CPEB4,KI	1.62947	1.00E-10	1.62947	1322.74	2150.339
A_23_P33	NM_05708	Homo sapi	KRT3,K3,C	1.62929	3.68E-07	1.62929	9326.375	15181.86
A_32_P79	NM_00574	pre-B-cell c	PBEF1,PB	1.62918	8.13E-28	1.62918	17152.42	27953.07
A_24_P20	NM_00322	transcriptio	TFAP2B,C	1.62872	0.00042	1.62872	506.0872	787.9854
A_24_P24	NM_17069	Homo sapi	dJ222E13.	1.62836	1.00E-06	1.62836	11341.4	18590.27
A_32_P218823		A8A2_MO	THC19684	1.62771	3.33E-13	1.62771	2286.342	3649.726
A_32_P57	CD365380	V-ets eryth	ETS2	1.62768	0.00011	1.62768	1086.242	1812.985
A_32_P93807		Unknown	A_32_P93	1.62717	5.61E-06	1.62717	22018.93	36252.73
A_24_P834066		Unknown	A_24_P83	1.62699	5.75E-09	1.62699	973.3373	1573.375
A_23_P10	NM_17583	Homo sapi	SMOX,PA	1.62699	3.00E-06	1.62699	12219.17	20226.82
A_23_P11	NM_19916	CXYorf1-re	FLJ25222	1.6265	3.05E-11	1.6265	38533.14	63051.47
A_24_P35	NM_00722	Homo sapi	NXPH4,NF	1.62618	2.46E-06	1.62618	16123.64	26322.96
A_23_P94	NM_01339	Homo sapi	TMEM2	1.62368	2.28E-07	1.62368	1622.738	2632.516
A_32_P162797		Unknown	THC19156	1.62345	2.66E-07	1.62345	2048.031	3311.155
A_24_P10	NM_00198	envoplakin	EVPL,EVP	1.62339	2.96E-09	1.62339	15099.05	24528.96
A_23_P27	NM_00227	Homo sapi	KRT15,K1	1.62323	1.54E-13	1.62323	5001.707	8110.983
A_23_P56868		Unknown	A_23_P56	1.62296	1.02E-08	1.62296	5394.604	8727.854
A_24_P18	NM_00218	interleukin	IL1RAP,IL1	1.62292	0.00004	1.62292	405.0483	650.7882
A_23_P32	NM_03149	protocadhe	PCDHA3,F	1.62289	1.14E-08	1.62289	602.2553	982.6481
A_23_P21	NM_00354	H4 histone	HIST1H4C	1.62271	7.90E-24	1.62271	195678.3	317272.7
A_32_P63562		U16359 nit	THC18591	1.62215	3.77E-06	1.62215	896.8109	1430.33
A_23_P21	NM_00363	Homo sapi	ENC1,NRF	1.62197	0.0003	1.62197	362.0602	578.618
A_23_P14	XM_37274	similar to p	LOC39097	1.62178	4.31E-06	1.62178	2389.444	3895.625
A_23_P25	BC007034	Metallothio	MT2A	1.6215	4.69E-14	1.6215	20887.12	34079.67
A_23_P11	NM_00524	Fibroblast g	FGF3	1.6214	0.00002	1.6214	53945.09	88341.88
A_32_P86	AB040943	KIAA1510	KIAA1510	1.62133	5.04E-19	1.62133	3408.365	5486.648
A_24_P28	NM_00659	heat shock	HSPA8,LA	1.62122	6.03E-19	1.62122	236182.1	384291.8
A_24_P35	NM_03058	UDP-Gal:b	B4GALT2	1.62065	5.47E-14	1.62065	6937.006	11262.99

A_32_P12	BF592593	Glutamate	GRIN2D	1.62041	1.91E-10	1.62041	856.3475	1381.906
A_23_P50	NM_13077	osteoclast-	OSCAR,PI	1.61964	0.00003	1.61964	897.1752	1423.668
A_23_P60	NM_00114	Homo sapi	ALOX15B	1.61917	1.87E-08	1.61917	1093.977	1742.916
A_24_P10	NM_08082	Homo sapi	C20orf144,	1.61899	1.13E-10	1.61899	1743.222	2811.834
A_23_P11	NM_02479	Homo sapi	ABHD9,FL	1.61805	6.38E-07	1.61805	32023.4	51998.61
A_23_P25	NM_01232	Homo sapi	DNAJB9,M	1.61721	8.92E-07	1.61721	2812.747	4557.026
A_24_P13	U94353	Radical frin	RFNG	1.61719	1.07E-20	1.61719	2691.401	4348.757
A_24_P18	NM_00357	latent trans	LTBP4,LTE	1.61697	6.26E-15	1.61697	2375.52	3831.159
A_23_P24	NM_00659	heat shock	HSPA8,LA	1.616	1.78E-16	1.616	239485	388679.4
A_23_P39	NM_20330	Homo sapi	RKHD1,TIN	1.61587	5.89E-08	1.61587	5508.917	9001.326
A_23_P67	NM_03078	Homo sapi	RSHL1,RS	1.61578	6.13E-10	1.61578	3711.51	5969.153
A_32_P15	BC067760	Mitogen-ac	LOC40783	1.61566	3.14E-08	1.61566	12331.5	20112.82
A_24_P12	NM_00515	actin, alpha	ACTC	1.61537	4.51E-14	1.61537	7015.968	11372.53
A_23_P38	NM_15262	Homo sapi	GLIS3,ZNF	1.61521	2.90E-12	1.61521	1739.82	2829.098
A_23_P32	NM_00075	carnitine ac	CRAT,CAT	1.61389	2.89E-11	1.61389	613.6523	985.5291
A_24_P29	NM_00061	CD4 antige	CD4,CD4m	1.61387	2.79E-06	1.61387	72400.21	118338
A_24_P93	BC039241	LOC12673	LOC12673	1.61334	7.53E-09	1.61334	2836.981	4565.963
A_23_P12	NM_00327	Homo sapi	TM7SF2,A	1.61279	7.03E-12	1.61279	5282.531	8514.356
A_24_P73	BI041703	Cisplatin re	CRR9	1.61251	1.94E-07	1.61251	16119.55	26064.51
A_24_P638608		Unknown	A_24_P63	1.61245	4.71E-09	1.61245	691.0018	1106.766
A_32_P13	NM_00659	heat shock	HSPA8,LA	1.61234	2.00E-23	1.61234	225092.5	363383.6
A_23_P20	NM_02099	chorionic s	CSH2,CSE	1.61213	3.17E-06	1.61213	662.043	1047.069
A_32_P186018		Unknown	THC19167	1.61201	0.00065	1.61201	44127.44	71068.31
A_24_P48	NM_19847	Homo sapi	FLJ46111	1.61189	5.87E-27	1.61189	5955.429	9602.879
A_23_P37	NM_00708	Homo sapi	SOX21,SO	1.60981	1.35E-13	1.60981	4305.369	6915.062
A_23_P53	NM_05410	H-rev107-li	HRLP5	1.60949	1.33E-08	1.60949	6152.696	9909.395
A_24_P14	NM_00194	diphtheria t	HBEGF,DT	1.60934	9.30E-10	1.60934	785.2679	1269.953
A_24_P869138		Unknown	A_24_P86	1.60875	5.24E-09	1.60875	8946.186	14414.04
A_24_P32	AF432209	KIAA1641	KIAA1641	1.6082	0.00041	1.6082	514.6995	795.8579
A_24_P84	AY090769	Ribosomal	AY090769	1.60787	1.53E-24	1.60787	5023.883	8083.225
A_24_P35	NM_00603	histone de	HDAC4,HD	1.60734	0.00092	1.60734	347.584	545.0361
A_23_P22	NM_01552	Homo sapi	TBC1D10E	1.60718	0.00014	1.60718	82504.34	134190
A_24_P13	NM_00155	interferon-r	IFRD1,PC	1.60553	2.41E-11	1.60553	4949.912	7999.229
A_24_P264413		Unknown	A_24_P26	1.60543	0.00044	1.60543	467.0018	779.8841
A_32_P83	BC031954	Zinc finger	ZEC	1.60469	2.12E-07	1.60469	2295.095	3665.357
A_24_P306614		Unknown	A_24_P30	1.60403	1.88E-21	1.60403	4060.252	6514.135
A_24_P90	BQ674642	Transcribe	BQ674642	1.60287	5.08E-16	1.60287	1354.966	2165.114
A_24_P75	AK025685	Homo sapi	CREB3L2,	1.60276	5.56E-08	1.60276	4309.497	6937.099
A_24_P17	NM_00008	ceroid-lipof	CLN3,BTS	1.60232	4.46E-09	1.60232	614.6458	987.8247
A_24_P16	NM_02157	breakpoint	BCR,ALL,	1.60214	4.09E-06	1.60214	684.3336	1074.946
A_24_P34	NM_00660	unactive pr	PTGES3,P	1.60182	2.07E-11	1.60182	5917.924	9537.393
A_24_P31	NM_00219	insulinoma	INSM1,IA1	1.60095	4.03E-21	1.60095	3437.133	5493.365
A_24_P39	NM_03077	tumor stror	ZBP1,DLM	1.60004	7.27E-06	1.60004	532.6826	838.7671
A_23_P89	NM_00365	Homo sapi	CBX4,PC2	1.59911	4.66E-11	1.59911	2259.324	3611.174
A_24_P56	BC035751	Homo sapi	TAFA5	1.59805	0.00004	1.59805	18970.18	30344.79
A_24_P40	NM_17504	Homo sapi	PILRB,FDF	1.59804	7.13E-15	1.59804	1185.169	1894.43
A_23_P11	NM_00437	Homo sapi	CRABP1,R	1.59755	0.00001	1.59755	8175.556	13083.09
A_24_P307075		Unknown	A_24_P30	1.59738	3.46E-10	1.59738	4942.479	7872.434
A_32_P75563		Unknown	THC19151	1.59618	9.03E-08	1.59618	10847.53	17257.8
A_23_P37	NM_15239	Homo sapi	AHSA2,Hc	1.59572	2.89E-09	1.59572	2121.677	3391.361
A_24_P93	AY358179	Brain acyl-	BACH	1.59534	0.00037	1.59534	484.7366	745.2273
A_24_P24	NM_14530	hypothetica	LOC22007	1.59503	1.40E-07	1.59503	15648.54	24996.41

A_23_P34	NM_00264	Homo sapi	PIM1,PIM	1.59468	4.03E-06	1.59468	1188.313	1872.498
A_23_P30	NM_00574	Homo sapi	PBEF1,PB	1.59231	9.61E-07	1.59231	9090	14716.92
A_23_P10	NM_02044	Homo sapi	NAV1,POM	1.5916	0.00019	1.5916	1016.531	1584.469
A_24_P36	NM_00106	tumor necr	TNFRSF1A	1.59072	1.28E-12	1.59072	3560.465	5663.13
A_32_P214611		Unknown	THC19045	1.59068	2.55E-10	1.59068	23362.8	37211.93
A_23_P78	NM_00705	suppressor	SFRS16,C	1.59059	0.00004	1.59059	505.6394	792.0248
A_24_P126731		Unknown	ENST0000	1.59039	8.18E-21	1.59039	3404.369	5416.128
A_24_P93	Z25424	H.sapiens	Z25424	1.59028	2.60E-11	1.59028	2115.031	3344.383
A_24_P21	AY063455	D21S2091	D21S2091	1.59025	1.09E-06	1.59025	777.454	1218.206
A_24_P814096		AE003449	THC18772	1.58957	2.74E-08	1.58957	4959.56	7950.826
A_23_P85	NM_00458	Homo sapi	RABGGTB	1.58935	1.17E-12	1.58935	22394.64	35830.61
A_23_P14	NM_00368	Homo sapi	PDLIM4,RI	1.58906	1.91E-06	1.58906	15283.14	24321.71
A_32_P108567		Unknown	A_32_P10	1.58834	5.05E-08	1.58834	966.3857	1515.773
A_23_P12	M26004	Compleme	CR2	1.58779	5.70E-10	1.58779	5227.984	8285.713
A_24_P19	AB029038	Homo sapi	KIAA1115	1.58746	2.04E-06	1.58746	16746.52	26608.98
A_24_P35	NM_00462	wingless-ty	WNT11,HW	1.58579	7.62E-15	1.58579	2031.629	3198.093
A_24_P705167		Unknown	A_24_P70	1.5856	3.33E-21	1.5856	1503.442	2383.323
A_24_P32	NM_00059	insulin-like	IGFBP3,IB	1.58529	0.00077	1.58529	1035.336	1556.222
A_23_P10	NM_01232	Homo sapi	MAFF,U-M	1.58466	2.35E-08	1.58466	654.7323	1030.594
A_23_P27	NM_01589	Homo sapi	ZBTB7A,LI	1.58437	6.92E-13	1.58437	4714.626	7466.38
A_24_P28	NM_00193	Homo sapi	DLAT,DLT	1.58413	4.25E-06	1.58413	8611.068	13535.04
A_23_P13	NM_00342	Homo sapi	ZNF42,MZ	1.58393	6.48E-12	1.58393	1281.862	2021.996
A_23_P43	BC022822	Immunoglo	IGLC2	1.58355	5.92E-09	1.58355	2384.537	3693.914
A_32_P57	NM_00631	ubiquitin sp	USP15,UN	1.5827	2.52E-17	1.5827	2625.642	4153.609
A_24_P12	AK023580	Chromosor	C9orf88	1.58143	0.0001	1.58143	463552.6	733074.7
A_32_P3564		Unknown	A_32_P35	1.58128	1.30E-07	1.58128	6984.841	10945.13
A_32_P79152		Unknown	THC19758	1.58093	9.23E-07	1.58093	31770.77	50248.16
A_23_P15	NM_00490	Homo sapi	CREB5,CR	1.58081	3.84E-06	1.58081	643.4247	994.6454
A_32_P21	AK124803	Clone 236	AK124803	1.58062	0.00009	1.58062	613.5835	941.7241
A_23_P25	NM_00048	Homo sapi	ARSA,MLD	1.57996	5.48E-12	1.57996	4437.089	7005.593
A_23_P412927		Unknown	A_23_P41	1.57914	2.85E-06	1.57914	1157.352	1809.103
A_23_P10	NM_00595	Homo sapi	MT2A,MT2	1.57796	2.06E-11	1.57796	29350.29	46796.5
A_24_P289504		Unknown	A_24_P28	1.57792	5.12E-06	1.57792	707.3319	1101.407
A_23_P87	NM_02392	Homo sapi	C1QDC1,E	1.57726	1.16E-07	1.57726	630.9345	992.4415
A_24_P290114		Unknown	A_24_P29	1.57701	3.91E-07	1.57701	46570.28	73898.7
A_23_P39	NM_03352	Homo sapi	C19orf33,II	1.57473	3.55E-23	1.57473	5409.519	8514.627
A_23_P20	NM_03352	Homo sapi	C19orf33,II	1.57472	1.13E-20	1.57472	14791.56	23326.04
A_32_P74	U21556	Similar to b	LOC16699	1.57454	0.00002	1.57454	423.8806	672.7955
A_24_P21	NM_00173	Kruppel-lik	KLF5,CKLI	1.57345	7.45E-08	1.57345	917.0327	1423.005
A_24_P592591		PBEF_HUI	THC18675	1.57307	3.46E-12	1.57307	12607.51	19962.73
A_23_P22	NM_03262	Homo sapi	BEX2,BEX	1.57265	1.38E-07	1.57265	2350.568	3699.467
A_24_P91	BX647673	Golgin-67	GOLGIN-6	1.57236	0.0001	1.57236	390.5519	610.4203
A_24_P29	BC032478	Mitogen-ac	MAP2K3	1.57212	1.46E-10	1.57212	9141.006	14442.63
A_23_P58	NM_03232	hypothetica	MGC4618	1.57125	0.00018	1.57125	172586.9	274625.7
A_23_P25	NM_02461	Homo sapi	HSPBAP1,	1.57119	7.00E-06	1.57119	610.5908	951.219
A_24_P897062		Unknown	A_24_P89	1.57077	0.00003	1.57077	382.5044	603.0366
A_23_P39	NM_00627	related RA	RRAS	1.57073	7.59E-09	1.57073	6172.577	9690.276
A_24_P26	NM_00529	G protein-c	GPR20	1.57052	9.78E-09	1.57052	3822.523	5977.886
A_24_P88	XM_38010	hypothetica	LOC40250	1.5704	0.00039	1.5704	1035.936	1597.189
A_24_P732565		Unknown	THC19153	1.56956	1.76E-07	1.56956	768.9283	1218.532
A_24_P81	BC033983	Homo sapi	BC033983	1.56894	0.00002	1.56894	739.6289	1132.014
A_23_P55	NM_00273	Homo sapi	PRKCA,PK	1.5685	0.00002	1.5685	653.2958	1000.886

A_23_P20	NM_00217	Homo sapi	IFNA10,MC	1.56786	6.85E-09	1.56786	702.3593	1096.335
A_24_P64	BC052374	Hypothetic	LOC25303	1.56706	1.22E-11	1.56706	1005.046	1579.748
A_23_P15	NM_02410	hypothetica	MGC2654,	1.56685	0.00002	1.56685	5097.267	8064.17
A_23_P42	NM_13862	Homo sapi	BCL2L11,E	1.56668	0.00003	1.56668	626.5971	959.5272
A_24_P41	NM_14517	NSE1	FAM84A,N	1.56575	4.49E-08	1.56575	8777.51	13840.68
A_23_P35	AB051487	Dual specif	DUSP16	1.56544	8.73E-11	1.56544	1568.731	2452.805
A_23_P36	NM_20733	hypothetica	ERICH1,H	1.56473	0.00039	1.56473	367.9315	567.7494
A_23_P60	NM_00595	Homo sapi	MT1G,MT1	1.56314	3.18E-11	1.56314	6772.875	10639.4
A_24_P12	NM_00471	hepatocyte	HGS,HRS,	1.56313	0.00003	1.56313	8203.063	12691.6
A_23_P49	NM_00680	KDEL (Lys	KDELR1,E	1.56264	3.43E-12	1.56264	6366.806	9954.48
A_23_P34	NM_00598	Homo sapi	SPRR1A,S	1.56241	2.77E-12	1.56241	24604.83	38451.16
A_23_P37	NM_00055	Homo sapi	HBA1,CD3	1.56205	8.58E-12	1.56205	1529.663	2380.942
A_32_P48	AK058133	Hypothetic	FLJ25404	1.56203	1.24E-11	1.56203	1252.989	1945.103
A_23_P14	NM_00573	ADP-ribosy	ARL4,ARL	1.5619	4.56E-08	1.5619	1000.7	1553.766
A_24_P91	NM_03283	hypothetica	FLJ14775	1.56186	2.86E-12	1.56186	2965.599	4630.224
A_24_P28	BC015531	Homo sapi	CA8,CALS	1.56149	0.00002	1.56149	752.2747	1142.767
A_24_P93	NM_00201	fragile histi	FHIT,FRA3	1.5611	0.00005	1.5611	453630.4	709585.3
A_23_P12	NM_01334	G protein-c	GPR132,G	1.56105	0.00002	1.56105	1219.575	1888.059
A_23_P41	NM_01208	Homo sapi	ELL2	1.561	6.37E-09	1.561	2731.722	4286.21
A_32_P23	NM_13344	Homo sapi	CTGLF1,M	1.56089	1.28E-07	1.56089	1463.352	2271.138
A_32_P23	NM_15278	hypothetica	MGC40405	1.55926	0.00023	1.55926	614.647	924.0468
A_24_P67	NM_19933	hypothetica	LOC37476	1.55879	8.51E-06	1.55879	764.4487	1165.844
A_23_P90	NM_01589	Homo sapi	PIAS4,PIA	1.55877	1.56E-09	1.55877	3429.463	5334.387
A_23_P15	NM_00335	Homo sapi	SCGB1A1,	1.5587	2.92E-09	1.5587	2927.944	4522.81
A_23_P34	NM_02031	Homo sapi	MNT,ROX,	1.55864	3.52E-10	1.55864	4465.506	6970.632
A_23_P13	NM_00134	diacylglyce	DGKQ,DA	1.55821	1.53E-15	1.55821	1732.293	2694.936
A_24_P10	NM_00364	diacylglyce	DGKZ,DA	1.5576	0.00027	1.5576	973.854	1483.451
A_24_P52	BC031320	LOC44030	BC031320	1.5575	3.58E-13	1.5575	5526.648	8639.12
A_24_P392505		Unknown	A_24_P39	1.55704	1.59E-09	1.55704	2488.649	3861.201
A_24_P67	AF307332	Meningiom	MGEA5	1.55666	1.40E-07	1.55666	1830.137	2854.188
A_32_P52	NM_03320	hypothetica	BC002942	1.55613	1.06E-08	1.55613	4345.407	6749.739
A_24_P20	U28831	Protein imr	LOC40098	1.55609	4.12E-11	1.55609	1667.693	2595.75
A_23_P20	NM_00228	Homo sapi	KRTHB5,H	1.55543	3.27E-07	1.55543	5191.833	8090.236
A_24_P15	NM_02273	Lipid phosp	LPPR2	1.55517	2.72E-22	1.55517	1551.489	2412.786
A_24_P67	AK074570	Similar to k	LOC37525	1.55474	2.97E-17	1.55474	1877.578	2918.969
A_24_P55	BC038432	Retinoic ac	RARA	1.55425	0.00003	1.55425	841.4758	1282.156
A_32_P74	BM930959	Transcribe	BM930959	1.55403	0.00044	1.55403	1701.935	2501.513
A_24_P57	AL137705	Glutamate	GRIK5	1.55374	0.00017	1.55374	45195.19	71177.01
A_24_P25	NM_17239	Homo sapi	NFATC1,N	1.55337	4.86E-08	1.55337	876.9186	1349.536
A_24_P20	NM_00374	insulin rece	IRS2	1.55326	1.57E-10	1.55326	5005.206	7767.307
A_24_P22939		Unknown	A_24_P22	1.55301	4.99E-11	1.55301	2997.026	4651.232
A_24_P91	L20941	Ferritin, he	FTH1	1.55287	0.00005	1.55287	92633.57	145484.3
A_32_P80295		Unknown	THC18796	1.55276	8.15E-07	1.55276	1891.872	2954.047
A_24_P35	AK055956	FLJ31394	FLJ31394	1.55268	7.50E-07	1.55268	1316.755	2063.824
A_23_P21	NM_01615	Homo sapi	TAOK2,PS	1.55259	5.57E-06	1.55259	20545.63	31774.21
A_32_P75867		ALU1_HUM	THC18592	1.55176	2.98E-06	1.55176	732.6758	1122.88
A_32_P83	AF318368	Pp8961	pp8961	1.55172	1.18E-12	1.55172	1164.221	1811.345
A_23_P13	NM_01656	E2IG2 prot	CHCHD8,E	1.55073	0.00003	1.55073	79934.27	126371.5
A_24_P400690		Unknown	ENST0000	1.54991	1.17E-06	1.54991	562.3363	867.5579
A_24_P780393		Unknown	A_24_P78	1.54976	0.00052	1.54976	463280	717786.6
A_24_P24	NM_18261	hypothetica	FAM70B,M	1.54915	3.24E-12	1.54915	3257.527	5040.226
A_24_P479065		Unknown	A_24_P47	1.54913	1.88E-10	1.54913	1357.165	2096.721

A_24_P92	AF321778	Toll interac	TOLLIP	1.54901	2.67E-24	1.54901	5756.259	8917.965
A_23_P16	NM_01494	KIAA0854	ZHX2,KIA	1.54896	3.55E-09	1.54896	2481.77	3842.752
A_24_P18	NM_00057	decay acce	DAF,CR,T	1.54833	5.06E-08	1.54833	3964.845	6194.767
A_23_P10	NM_00588	adenomato	APC2,APC	1.54831	0.00026	1.54831	408.5776	623.418
A_23_P34	NM_00151	Homo sapi	GTF2B,TF	1.54728	4.88E-08	1.54728	2367.952	3673.594
A_23_P37	AK056767	V-maf mus	MAFK	1.54708	4.23E-11	1.54708	3069.546	4746.675
A_23_P25	NM_00495	Homo sapi	EBI2	1.54704	0.00003	1.54704	414.2369	639.098
A_32_P219126		Unknown	A_32_P21	1.54671	2.29E-14	1.54671	1822.176	2821.381
A_32_P15	BU730134	Chromobo	CBX6	1.54635	0.00081	1.54635	476.8104	765.0341
A_23_P78	NM_01624	Homo sapi	DHRS10,re	1.54564	4.83E-10	1.54564	1708.857	2624.74
A_24_P19	NM_14777	Homo sapi	SNX15,HS	1.54563	0.00012	1.54563	422.5394	645.4655
A_23_P33	NM_00448	Homo sapi	GOLGB1,G	1.54546	1.02E-08	1.54546	3038.7	4726.912
A_32_P165823		Unknown	A_32_P16	1.54535	0.00039	1.54535	810.2188	1295.614
A_23_P47	NM_13839	Membrane	MARCH-IX	1.54441	4.97E-08	1.54441	8621.373	13317.38
A_23_P10	AF357221	DNM1DN8	DNM1DN8	1.54327	3.02E-07	1.54327	464146.3	716907.1
A_23_P32	NM_14504	hypothetica	MGC20983	1.54308	0.00054	1.54308	461.5691	691.4353
A_24_P751082		CLS1_HU	THC19430	1.54303	0.00003	1.54303	910.9089	1375.414
A_23_P25	NM_01335	Homo sapi	PKN3,RP1	1.54303	2.37E-06	1.54303	534.4856	821.6454
A_23_P11	NM_00601	Homo sapi	ST8SIA2,S	1.54261	6.33E-09	1.54261	3471.927	5351.386
A_32_P95	NM_17213	Homo sapi	WFDC10B	1.54258	4.57E-13	1.54258	16324.2	25191.91
A_23_P85	NM_00568	G protein-c	GPR52,MG	1.54257	1.96E-06	1.54257	873.5183	1326.172
A_24_P152278		Unknown	ENST0000	1.5422	2.03E-06	1.5422	2156.579	3341.733
A_32_P14	BG540029	Transcribe	BG540029	1.54156	1.42E-09	1.54156	8325.572	12848.24
A_24_P34	BC031698	Homo sapi	ENST0000	1.54139	0.00016	1.54139	877.0365	1325.514
A_23_P12	NM_08073	Homo sapi	C20orf141,	1.54136	0.00002	1.54136	20957.94	32042.21
A_24_P52	NM_00321	thymine-D	TDG	1.54093	9.26E-08	1.54093	6006.597	9325.775
A_23_P13	NM_00139	Homo sapi	DUSP4,TY	1.54049	1.06E-11	1.54049	6584.033	10165.17
A_24_P24	BC014322	MRNA, en	BC014322	1.54036	6.62E-07	1.54036	5950.58	9186.746
A_24_P37	NM_00318	synaptotag	SYT5	1.53938	1.08E-17	1.53938	5684.877	8750.041
A_23_P78	NM_02030	Homo sapi	SLC17A7,E	1.53931	7.40E-08	1.53931	10158.37	15639.89
A_23_P42	BC060318	Homo sapi	BC060318	1.53874	3.63E-09	1.53874	2826.208	4328.851
A_23_P43	NM_17360	Homo sapi	C14orf80,M	1.53837	8.29E-12	1.53837	1289.679	1988.329
A_32_P86	AK131056	Chymotryp	CTRB1	1.53831	0.00002	1.53831	61294.55	95332.41
A_24_P890956		Unknown	A_24_P89	1.53773	2.38E-12	1.53773	8311.379	12777.13
A_23_P86	M32220	DR1-assoc	DRAP1	1.53714	0.00001	1.53714	457.2601	698.7235
A_23_P20	NM_17561	Homo sapi	MT1E,MT1	1.53651	1.24E-09	1.53651	2496.023	3844.63
A_24_P26	AK096981	CDNA FLJ	AK096981	1.53646	2.80E-19	1.53646	4859.718	7500.646
A_32_P49	XM_37204	similar to a	LOC38965	1.53565	3.44E-23	1.53565	17227.23	26448.91
A_24_P287526		Unknown	A_24_P28	1.5355	7.31E-06	1.5355	3596.777	5530.153
A_24_P550411		Unknown	A_24_P55	1.53544	0.00008	1.53544	933.3114	1402.136
A_32_P29130		Unknown	A_32_P29	1.53533	9.89E-09	1.53533	4314.159	6667.176
A_32_P30320		Unknown	A_32_P30	1.53524	5.17E-11	1.53524	1085.806	1661.401
A_23_P20	NM_17496	Homo sapi	ST3GAL3,	1.53456	7.92E-13	1.53456	1336.522	2046.703
A_32_P25838		Unknown	A_32_P25	1.53361	2.17E-08	1.53361	6562.449	10058.1
A_32_P22	NM_13344	Homo sapi	CTGLF1,M	1.53352	3.66E-06	1.53352	1405.612	2146.485
A_24_P27	NM_18301	Homo sapi	CREM,ICE	1.53314	0.00001	1.53314	711.6546	1076.378
A_24_P332471		Unknown	A_24_P33	1.53297	5.94E-22	1.53297	27755.66	42583.2
A_23_P13	NM_02206	Homo sapi	FAM38B,H	1.53275	4.62E-09	1.53275	798.1617	1215.79
A_24_P10	NM_00164	apolipoprot	APOC1	1.53258	2.34E-20	1.53258	11422.22	17514.97
A_24_P93	S75708	RNA, U17	CHC1	1.53242	4.53E-20	1.53242	3144.916	4822.231
A_23_P16	NM_18342	TSC22 dor	TGFB1I4	1.53228	2.66E-08	1.53228	991.0723	1528.148
A_23_P20	NM_03286	Homo sapi	TNS4,CTE	1.53218	0.00007	1.53218	1606.732	2445.377

A_24_P28	NM_00079	dopamine r	DRD2,D2R	1.53167	1.63E-09	1.53167	4874.975	7440.18
A_23_P67	S49953	V-myc mye	MYCNOS	1.53098	2.08E-08	1.53098	757.7056	1162.738
A_23_P32	NM_02283	hypothetica	FLJ22833,	1.53077	7.16E-14	1.53077	2238.983	3429.125
A_24_P23	NM_00132	cystatin SA	CST2,MGC	1.53065	8.13E-08	1.53065	2177.146	3311.625
A_23_P15	NM_03216	Homo sapi	ZNF394,FL	1.53064	1.52E-10	1.53064	1433.243	2186.609
A_23_P17	NM_00574	Homo sapi	DNAL4,PIC	1.5292	2.22E-16	1.5292	1634.185	2496.267
A_23_P30	NM_14523	LOC90353	LOC90353	1.52894	0.00023	1.52894	366.76	561.0118
A_24_P479364		Unknown	THC19576	1.52675	4.70E-10	1.52675	1177.766	1810.803
A_32_P17	NM_00016	glycerol kin	GK,GKD	1.52673	1.84E-06	1.52673	1020.041	1538.541
A_32_P41	NM_05289	Homo sapi	MBD6,KIA	1.5266	2.12E-17	1.5266	4320.873	6601.603
A_24_P21	NM_00384	tumor necr	TNFRSF1C	1.52643	1.65E-18	1.52643	3528.461	5385.891
A_23_P43	BC042539	Homo sapi	LOC44242	1.52628	1.33E-12	1.52628	1255.461	1911.467
A_23_P51	NM_00592	Homo sapi	MEF2D	1.52599	0.00002	1.52599	12534.66	19239.88
A_24_P34	NM_03232	Inhibitor of	ING5	1.52599	0.00005	1.52599	417.4103	635.915
A_32_P12344		Unknown	A_32_P12	1.52527	0.00005	1.52527	8262.012	12516.56
A_32_P209163		Unknown	THC18595	1.52506	0.00002	1.52506	670.1461	1006.466
A_23_P79	NM_01647	Homo sapi	C20orf111,	1.52487	2.91E-16	1.52487	3740.778	5712.084
A_23_P50	NM_00113	Homo sapi	AGRP,ART	1.52464	0.00035	1.52464	388.0868	587.395
A_24_P10	NM_01502	KIAA1037	WDTC1,KI	1.52445	0.00042	1.52445	6784.006	10448.83
A_23_P40	NM_17846	Homo sapi	PTPNS1L2	1.52406	0.0002	1.52406	420.9847	647.2894
A_23_P36	NM_01441	Homo sapi	TCL6,TNG	1.52349	0.00005	1.52349	8210.235	12522.44
A_23_P76	NM_00263	Homo sapi	PGF,PLGF	1.52341	6.60E-07	1.52341	1416.603	2102.787
A_23_P36	NM_00054	Homo sapi	TCF1,HNF	1.52335	2.58E-06	1.52335	12777.82	19283.7
A_23_P17	NM_01656	BM88 antig	BM88,MGC	1.52122	6.06E-15	1.52122	1976.398	3003.124
A_32_P41	NM_00262	prefoldin 1	PFDN1,PD	1.5212	3.41E-07	1.5212	823.1996	1243.761
A_24_P891899		Unknown	A_24_P89	1.52102	0.00002	1.52102	8426.145	12684.5
A_23_P40	XM_37118	hypothetica	LOC38855	1.52057	1.68E-10	1.52057	2433.52	3692.831
A_24_P10	AK130825	Isolate A8	AK130825	1.52028	0.00019	1.52028	709.8619	1048.014
A_24_P17	NM_00042	loricrin	LOR	1.51997	0.00071	1.51997	383.7261	599.048
A_32_P139634		Unknown	THC18390	1.51973	0.00023	1.51973	75382.32	115871
A_24_P49	XM_29185	similar to E	LOC34109	1.51895	1.91E-07	1.51895	14804.13	22432.79
A_24_P75	U80753	Trinucleotid	TNRC18	1.51876	4.56E-08	1.51876	10088.39	15282.13
A_23_P34	BC032412	Hypothetica	MGC40574	1.51814	1.44E-06	1.51814	1572.501	2327.009
A_32_P172002		Unknown	A_32_P17	1.51757	0.00005	1.51757	8988.76	13712.33
A_24_P161525		Unknown	ENST0000	1.51708	5.89E-12	1.51708	5663.044	8573.588
A_24_P13	AK074570	Similar to K	LOC37525	1.51675	0.00001	1.51675	549.6205	834.449
A_23_P41	NM_14529	similar to R	LOC19733	1.51604	0.00003	1.51604	3639.673	5557.18
A_24_P15	NM_15278	hypothetica	MGC40405	1.51537	0.00002	1.51537	3359.612	5142.628
A_23_P38	NM_02264	chorionic s	CSH1,PL,C	1.51533	0.00009	1.51533	561.9226	834.1285
A_23_P40	NM_00109	acrosin	ACR	1.51515	0.00009	1.51515	549.7402	817.7796
A_24_P743672		U5 snRNP-	NP106139	1.51505	5.27E-15	1.51505	4404.051	6667.779
A_23_P35	NM_03330	Caspase 4	CASP4	1.51486	0.00002	1.51486	21411.78	32843.64
A_23_P68	NM_01416	HSPC072	HSPC072	1.51342	2.34E-08	1.51342	942.2997	1417.437
A_23_P90	NM_01441	Bcl-2 bindi	BBC3,JFY	1.5134	0.0004	1.5134	483983	732459.1
A_23_P37	NM_00594	Homo sapi	MT1B,MT1	1.51325	5.40E-09	1.51325	7773.611	11818.04
A_23_P51	NM_01636	myo-inosit	ISYNA1	1.51303	3.21E-06	1.51303	626.0284	955.1046
A_24_P92	NM_13902	Homo sapi	ADAMTS1	1.51294	1.18E-18	1.51294	2347.529	3551.761
A_23_P66	NM_02470	Homo sapi	ZNF668,FL	1.51276	0.00001	1.51276	3918.013	5913.476
A_23_P21	NM_00138	Homo sapi	DSCAM,CH	1.51275	0.00011	1.51275	324.1606	487.4882
A_23_P20	NM_00595	metallothio	MT1G,MT1	1.51269	9.80E-06	1.51269	2234.178	3393.652
A_23_P14	NM_01223	Homo sapi	PTRF,FKS	1.51257	0.00023	1.51257	394.1435	598.4196
A_23_P11	NM_00325	Homo sapi	ICAM5,TLN	1.51169	0.00002	1.51169	664.3452	989.8636

A_23_P84	NM_01894	homeo box	HMX1,H6	1.51167	3.86E-09	1.51167	928.752	1409.64
A_24_P349896		Unknown	A_24_P34	1.51109	5.49E-13	1.51109	4847.407	7317.933
A_23_P33	NM_17850	Homo sapi	DTX3,RNF	1.5104	1.54E-10	1.5104	2281.641	3442.177
A_32_P67	BC027978	Interleukin	IL6	1.51014	2.51E-06	1.51014	17540.35	26790.89
A_23_P84016		Unknown	A_23_P84	1.5099	3.50E-08	1.5099	4310.601	6547.173
A_23_P17	NM_00645	Homo sapi	PDLIM5,L9	1.50984	0.00044	1.50984	1445.586	2131.323
A_24_P13	BC021276	Immunoglo	IGHD	1.50977	5.00E-08	1.50977	5123.595	7740.291
A_23_P37	NM_01470	KIAA0256	KIAA0256	1.50973	5.60E-06	1.50973	602.8745	901.9828
A_23_P38	NM_03076	Homo sapi	WNT4,WN	1.50923	1.17E-09	1.50923	1069.856	1608.263
A_24_P936820		Unknown	A_24_P93	1.50911	6.92E-14	1.50911	1262.511	1908.037
A_23_P25	NM_01394	Homo sapi	CLIC4,H1,I	1.50855	1.84E-08	1.50855	1019.154	1544.276
A_24_P41530		Unknown	A_24_P41	1.50678	1.12E-10	1.50678	2094.384	3154.563
A_24_P17	NM_00735	heat shock	HSPCB,HS	1.50665	2.29E-08	1.50665	5409.916	8100.637
A_23_P45	NM_00446	Homo sapi	FIGF,VEGI	1.50628	1.96E-06	1.50628	991.9857	1477.478
A_23_P12	NM_02235	Homo sapi	DPEP3	1.50624	1.02E-07	1.50624	8422.313	12700.49
A_23_P10	NM_00383	Homo sapi	MATN4,HE	1.50613	3.40E-08	1.50613	798.8942	1206.931
A_24_P79	NM_00261	platelet fac	PF4,CXCL	1.50585	2.83E-08	1.50585	1032.37	1544.084
A_24_P30	NM_01207	Conserved	C16orf35,C	1.50555	6.41E-16	1.50555	1893.552	2848.799
A_23_P30	NM_00060	fibroblast g	FGFR1,H2	1.50519	0.00007	1.50519	688.9221	1018.667
A_23_P25	NM_00706	Homo sapi	TBC1D8,A	1.50518	5.13E-10	1.50518	1601.346	2407.723
A_23_P42	NM_00374	Homo sapi	SOCS1,JA	1.50463	5.47E-11	1.50463	5215.226	7836.328
A_32_P11	NM_00213	heterogene	HNRPA1,H	1.50462	7.91E-09	1.50462	5939.235	8896.55
A_32_P49	AL137566	MRNA; cDI	AL137566	1.5046	0.00013	1.5046	772.5139	1134.26
A_23_P25	NM_14520	similar to C	NAPRT1,P	1.50441	2.85E-17	1.50441	2073.301	3119.219
A_24_P290214		Unknown	A_24_P29	1.50327	7.99E-06	1.50327	691.0663	1050.647
A_23_P16	NM_00265	plasminoge	PLAUR,CD	1.5032	0.00081	1.5032	437.545	667.4746
A_24_P82	NM_13842	hypothetica	LOC11317	1.5027	1.10E-08	1.5027	737.2479	1105.244
A_23_P46	NM_00164	Homo sapi	APOC1	1.50259	9.44E-10	1.50259	13422.29	20283.49
A_23_P20	NM_17792	Homo sapi	H2AFJ,MG	1.50231	3.65E-12	1.50231	15724.58	23677.63
A_24_P26	NM_00660	unactive pr	PTGES3,P	1.50188	0.00007	1.50188	2396.766	3634.376
A_32_P122494		Unknown	A_32_P12	1.50135	0.00003	1.50135	678.4012	1001.751
A_24_P153342		Unknown	ENST0000	1.50132	0.00017	1.50132	793.8079	1155.557
A_24_P24	NM_00188	activating t	ATF2,HB1	1.50096	0.00012	1.50096	491.804	743.9901
A_32_P42	BC011998	Homo sapi	BC011998	1.5001	0.00001	1.5001	3362.908	5052.916
A_23_P42	NM_00491	protein dis	PDIA4,ERI	1.4999	0.00001	1.4999	6810.345	10335.15
A_23_P89	NM_00125	Homo sapi	CDK3	1.49897	6.39E-08	1.49897	759.4857	1136.701
A_24_P32930		Unknown	A_24_P32	1.49894	4.38E-15	1.49894	2836.55	4253.771
A_24_P229438		Unknown	A_24_P22	1.49856	1.04E-15	1.49856	3193.541	4782.359
A_24_P910381		Unknown	A_24_P91	1.49849	8.07E-13	1.49849	1213.681	1821.287
A_23_P59	NM_00038	cyclin-depe	CDKN1A,F	1.49738	2.74E-09	1.49738	7088.604	10642.55
A_24_P54	BC035669	Homo sapi	SPIRE2	1.49729	2.82E-10	1.49729	1610.597	2397.354
A_24_P764866		Unknown	A_24_P76	1.49703	1.72E-09	1.49703	2944.49	4399.322
A_23_P93	NM_15332	Homo sapi	SLC22A7,N	1.49688	1.56E-15	1.49688	1846.22	2761.791
A_32_P53713		CLN3_HUI	THC19788	1.49642	1.02E-09	1.49642	3028.889	4527.184
A_24_P20	NM_03299	Williams-B	WBSCR14	1.49639	0.00049	1.49639	26768.84	40117.79
A_24_P27	NM_01867	mesoderm	MESP1,IR	1.49577	1.67E-13	1.49577	1671.52	2498.29
A_24_P20	NM_00413	potassium	KCNMB1,S	1.49556	1.06E-09	1.49556	8360.332	12516.58
A_23_P42	NM_02011	hypothetica	ZC3HAV1,	1.49538	0.00002	1.49538	3999.021	6032.782
A_23_P30	NM_00399	Homo sapi	NFKB1,KB	1.49484	4.79E-06	1.49484	3872.989	5808.275
A_23_P41	NM_00725	Homo sapi	B4GALT7,	1.49447	1.36E-11	1.49447	2341.471	3494.141
A_32_P16	NM_00455	nuclease s	YBX1,YB1	1.49426	1.95E-18	1.49426	5388.319	8059.403
A_23_P41	NM_18277	Homo sapi	ALS2CL,R	1.494	0.00046	1.494	489.9619	723.2025

A_24_P31	AK026491	Homo sapi	RPL12	1.4936	8.70E-12	1.4936	31880.63	47411.96
A_23_P12	NM_00440	Homo sapi	DFFA,DFF	1.49352	1.11E-06	1.49352	5848.698	8553.121
A_23_P35	NM_17074	Homo sapi	HIST1H2A	1.49346	0.00064	1.49346	439.265	669.0635
A_24_P856273		Unknown	A_24_P85	1.49339	1.88E-09	1.49339	12658.03	18811.54
A_24_P12	NM_02054	choline ace	CHAT,CMS	1.49332	4.51E-11	1.49332	1650.123	2469.341
A_23_P329062		AF506835	THC18912	1.49322	0.00003	1.49322	13522.55	19969.14
A_32_P83795		BC032539	THC18158	1.49299	5.80E-07	1.49299	3791.906	5659.998
A_23_P16	NM_01801	Hypothetic	FLJ10154	1.49237	2.33E-07	1.49237	1304.4	1940.562
A_23_P34	NM_17843	Homo sapi	LCE3E,LEI	1.49237	4.41E-06	1.49237	45319.45	67946.27
A_24_P25	AL390214	Triple funct	TRIO	1.49232	0.00054	1.49232	741.593	1078.758
A_23_P25	NM_02522	Homo sapi	BPIL1,RYS	1.49168	3.31E-06	1.49168	3366.033	4974.09
A_24_P23	AF177925	Mucin 4, tra	MUC4	1.49026	2.69E-06	1.49026	21736.33	32606.27
A_23_P15	AB014541	Apoptosis-a	AATK	1.49013	2.43E-09	1.49013	828.4787	1232.551
A_23_P26	NM_01246	Homo sapi	TPSG1,TM	1.49008	0.00002	1.49008	479068.9	714441.9
A_32_P100464		Unknown	A_32_P10	1.48993	4.49E-10	1.48993	1686.66	2495.231
A_32_P123255		Unknown	THC19253	1.48976	3.50E-08	1.48976	3221.206	4780.641
A_24_P30	NM_14778	Homo sapi	CTSB,APP	1.48976	4.13E-09	1.48976	744.1728	1103.089
A_24_P298894		Unknown	ENST0000	1.48928	2.55E-18	1.48928	8557.691	12736.7
A_24_P35	NM_00130	claudin 3	CLDN3,RV	1.48909	2.82E-09	1.48909	5146.023	7657.99
A_23_P35	NM_15326	hypothetica	MGC33486	1.48876	1.61E-06	1.48876	2629.99	3935.188
A_24_P10	XM_37757	similar to 6	LOC40194	1.48859	8.65E-13	1.48859	13025.01	19322.61
A_23_P124699		Unknown	A_23_P12	1.4884	0.00001	1.4884	749.8395	1097.7
A_23_P13	NM_03142	retbindin	RTBDN	1.48812	0.00007	1.48812	11837.88	17631.8
A_24_P23	BC024297	Homo sapi	JRK,JH8,F	1.48742	3.26E-12	1.48742	1113.578	1650.959
A_24_P48	NM_02075	Homo sapi	CASKIN2,F	1.48709	1.37E-08	1.48709	3964.443	5892.334
A_32_P12	NM_19828	hypothetica	LOC34911	1.48666	2.19E-08	1.48666	8146.672	12118.71
A_24_P350252		Unknown	A_24_P35	1.48636	1.34E-11	1.48636	1499.152	2213.834
A_32_P90	BC038466	Hypothetic	LOC40111	1.48559	1.43E-19	1.48559	12437.71	18479.8
A_32_P75	BC034617	KIAA0565	KIAA0565	1.48545	3.86E-06	1.48545	2492.985	3681.401
A_24_P63	AK126509	Centaurin,	CENTB1	1.48493	2.65E-11	1.48493	1768.194	2617.422
A_24_P65	BG519703	602578841	BG519703	1.48474	3.25E-11	1.48474	1167.063	1730.333
A_23_P14	NM_00167	Homo sapi	ASNS,TS1	1.48469	4.07E-12	1.48469	39823.5	58875.98
A_24_P65	XM_37255	DNM1DN3	DNM1DN3	1.48442	0.00039	1.48442	604.0871	873.9926
A_23_P11	NM_00239	Homo sapi	MDK,MK,N	1.48353	3.76E-08	1.48353	10959.63	16331.31
A_24_P32	NM_00322	trefoil facto	TFF1,pS2,	1.4829	1.75E-12	1.4829	9213.729	13691.49
A_24_P86	AK022745	Hypothetic	RNU71B,U	1.48284	0.00079	1.48284	13231.04	20083.46
A_24_P409910		Unknown	A_24_P40	1.48234	9.61E-17	1.48234	2577.153	3820.439
A_23_P21	NM_00488	Homo sapi	CXCL14,K	1.48067	0.00019	1.48067	461.0602	677.3861
A_24_P929650		Unknown	A_24_P92	1.48052	2.19E-16	1.48052	4752.563	7040.938
A_23_P36	NM_15244	Homo sapi	FBXL14,Fb	1.48048	1.77E-07	1.48048	1504.973	2211.385
A_24_P170365		Unknown	ENST0000	1.48004	9.37E-08	1.48004	1645.236	2394.054
A_23_P49	NM_00450	homeo box	HOXB7,HC	1.47993	1.66E-11	1.47993	2913.115	4291.704
A_24_P12	NM_00166	ADP-ribosy	ARF4L,AR	1.47775	5.64E-06	1.47775	649.5427	965.7786
A_24_P35	NM_02454	hypothetica	C10orf76,F	1.47723	2.36E-08	1.47723	1218.302	1791.74
A_24_P91	AY129015	Amylase, a	AMY2B	1.4772	9.49E-15	1.4772	2669.303	3943.698
A_23_P75	NM_00239	solute carri	SLC3A2,4F	1.47716	0.00092	1.47716	3313.621	4894.746
A_23_P47	NM_02197	Homo sapi	ST14,HAI,I	1.47707	5.55E-17	1.47707	2521.359	3723.228
A_23_P27	AB075836	Zinc finger	ZNF418	1.47695	1.52E-07	1.47695	4077.879	5929.261
A_24_P27	NM_00086	histamine r	HRH1,H1-F	1.47628	1.28E-08	1.47628	1065.657	1570.687
A_23_P40	NM_02124	Homo sapi	CDH22,C2	1.47624	0.00049	1.47624	62046.38	92923.27
A_32_P23	BC037911	Homo sapi	BC037911	1.47607	0.00002	1.47607	7263.558	10547.59
A_23_P16	NM_01337	Homo sapi	VPREB3,8	1.47557	8.53E-09	1.47557	2690.055	3909.992

A_24_P184	BC042057	Papilin, pro	PAPLN	1.47542	0.00025	1.47542	531.7245	769.6349
A_23_P31	NM_00110	Homo sapi	ACTB	1.4754	3.00E-12	1.4754	68061.18	100712.6
A_23_P89	NM_01906	Homo sapi	SDK2,FLJ1	1.47524	0.00002	1.47524	30784.08	45608.25
A_24_P358534		Unknown	A_24_P358	1.4748	0.00045	1.4748	3101.527	4369.375
A_32_P20	NM_17357	hypothetica	LOC25531	1.47479	1.66E-06	1.47479	15143.83	22445.31
A_32_P171984		Unknown	A_32_P17	1.4747	0.00005	1.4747	486078.4	717016.3
A_24_P752362		Unknown	A_24_P752	1.47444	1.75E-13	1.47444	2838.742	4187.217
A_32_P184937		Unknown	A_32_P184	1.47437	1.06E-15	1.47437	5061.351	7467.999
A_23_P41	NM_01599	Homo sapi	KLF13,BTE	1.47426	2.23E-06	1.47426	848.3641	1264.832
A_23_P52	NM_00679	Homo sapi	EHD1,PAS	1.47397	4.38E-13	1.47397	4657.859	6868.541
A_23_P46	NM_02505	hypothetica	TRIM46,FL	1.47347	0.00009	1.47347	474.8942	688.8631
A_24_P46	XM_29281	Homo sapi	XM_29281	1.47311	1.42E-10	1.47311	6191.244	9137.494
A_23_P264	NM_00051	Homo sapi	HBA2,HBA	1.473	7.71E-06	1.473	1004.919	1458.554
A_23_P31	NM_15334	Ataxin 7-lik	ATXN7L2	1.47271	5.69E-13	1.47271	4113.099	6087.941
A_23_P13	NM_00436	Homo sapi	COL6A3,D	1.47221	8.18E-08	1.47221	10613.23	15616.95
A_32_P1291		Unknown	A_32_P129	1.4716	0.00092	1.4716	497474.9	732081.7
A_23_P33	AK054935	Homo sapi	TTC9B,FL	1.4715	4.14E-06	1.4715	3878.118	5704.833
A_24_P29	NM_01897	amino acid	SLC38A2,A	1.47056	0.00014	1.47056	5483.922	8204.644
A_24_P274	NM_02506	hypothetica	ZNF703,ZN	1.47037	1.34E-07	1.47037	688.0234	1009.933
A_24_P934	BQ422089	Heat shock	HSPH1	1.47035	7.48E-07	1.47035	909.1077	1329.169
A_23_P91	NM_00036	Homo sapi	THBD,TM,	1.47034	8.59E-07	1.47034	2340.616	3440.493
A_32_P139	BF510844	Transcribe	BF510844	1.46995	0.00003	1.46995	29642.48	43587.82
A_23_P86	NM_02283	von Willebr	VWA1,WA	1.46961	1.55E-07	1.46961	12464.92	18315.95
A_23_P84	NM_02101	synovial sa	SSX5,MGC	1.46956	0.00004	1.46956	608.7474	887.4612
A_23_P37	NM_20745	FLJ35767	FLJ35767	1.46941	0.00003	1.46941	636.6091	927.8612
A_23_P114	NM_02477	Homo sapi	LONRF3,R	1.46864	0.00007	1.46864	2252.442	3321.362
A_23_P13	NM_00216	Homo sapi	ID3,HEIR-1	1.46863	1.95E-12	1.46863	6319.663	9310.252
A_23_P129	NM_03093	Homo sapi	TMEM49,V	1.46749	1.75E-07	1.46749	4326.497	6363.457
A_32_P71	BU664361	Similar to h	LOC28545	1.46719	0.00021	1.46719	24819.82	36692.96
A_23_P20	NM_00051	Homo sapi	GH1,GH,G	1.46696	6.08E-06	1.46696	638.452	929.8477
A_23_P35	NM_14580	Homo sapi	ABTB2,DK	1.46693	9.28E-10	1.46693	2522.156	3696.449
A_24_P281514		Unknown	ENST0000	1.46666	1.06E-10	1.46666	1098.16	1609.298
A_24_P11	NM_15289	Homo sapi	C20orf161,	1.46628	0.0004	1.46628	757.0569	1082.02
A_24_P524452		Unknown	A_24_P524	1.4662	9.51E-18	1.4662	3019.335	4427.036
A_24_P41	AJ291670	DMRT-like	DMRTC1	1.46602	0.00091	1.46602	661.4442	934.6552
A_23_P20	NM_01903	Homo sapi	RHOF,RIF,	1.46599	4.75E-08	1.46599	1553.448	2271.582
A_23_P21	NM_01897	Homo sapi	SLC38A2,A	1.46561	0.00009	1.46561	618.502	898.3257
A_24_P17	NM_00450	heterogene	HNRPU,SA	1.4654	0.00004	1.4654	601.2579	884.4336
A_24_P7494		Unknown	ENST0000	1.46524	6.14E-10	1.46524	2189.733	3203.996
A_23_P68	NM_00322	Homo sapi	TFF1,pS2,	1.46502	0.00002	1.46502	19087.48	28396.84
A_23_P37	NM_01780	Homo sapi	LIME1,LIM	1.465	0.00002	1.465	2644.75	3830.512
A_23_P80	NM_15288	Homo sapi	CHST13,C	1.46483	5.72E-06	1.46483	9855.625	14477.53
A_24_P39	NM_00089	leukotriene	LTC4S,MG	1.4643	5.32E-07	1.4643	908.1644	1317.719
A_23_P10	BC002811	Homo sapi	BC002811	1.46412	4.36E-07	1.46412	23909.95	35152.57
A_32_P74778		Unknown	A_32_P74	1.46363	2.28E-13	1.46363	1846.249	2699.678
A_24_P624	AK074149	Phospholip	PLCL4	1.46334	0.00008	1.46334	1048.419	1506.307
A_23_P96	NM_00016	Homo sapi	GK,GKD	1.46326	0.0001	1.46326	717.1469	1035.067
A_24_P31	BC018095	DR1-assoc	DRAP1	1.46288	6.42E-06	1.46288	8790.404	12902.56
A_23_P35	NM_00113	Homo sapi	AFG3L1,AI	1.46257	2.03E-08	1.46257	3571.198	5245.502
A_23_P56	NM_00577	Homo sapi	DHRS9,RD	1.46251	0.0007	1.46251	565.301	810.0109
A_24_P938583		Unknown	A_24_P938	1.46214	2.55E-07	1.46214	1209.995	1778.479
A_32_P25	AB021742	Neurogenic	NEUROD2	1.46149	1.48E-06	1.46149	682.5188	1003.007

A_23_P32	AK075506	Midnolin	MIDN	1.46066	1.89E-06	1.46066	1087.782	1576.864
A_23_P13	NM_00161	Homo sapi	ACP2	1.46063	7.69E-06	1.46063	674.2971	979.0818
A_24_P91	AK128640	Solute carr	SLC11A2	1.46011	2.78E-06	1.46011	3774.203	5486.698
A_32_P175462		Unknown	A_32_P17	1.45988	1.62E-14	1.45988	1826.618	2664.868
A_23_P35	NM_00076	Homo sapi	CYP3A7,C	1.4597	7.66E-09	1.4597	1023.785	1488.094
A_24_P31	NM_01423	vesicle-ass	VAMP2,SY	1.45951	0.00005	1.45951	4949.994	7288.519
A_32_P64	X05126	Calponin 2	CNN2	1.45936	7.91E-06	1.45936	882.2944	1274.834
A_32_P85	AB015282	Dual-specil	DYRK1A	1.45929	0.00039	1.45929	608.3953	908.0408
A_24_P91	AB065089	Ribosomal	RPS2	1.45918	7.51E-17	1.45918	2989.185	4362.878
A_32_P220881		Unknown	A_32_P22	1.45899	0.00003	1.45899	10283.34	15005.71
A_23_P11	NM_02193	invasion inl	IIP45,FLJ1	1.45896	0.00001	1.45896	555.2618	810.1701
A_23_P14	NM_01612	putative rin	RNF12,RL	1.45867	0.0004	1.45867	1133.679	1630.687
A_32_P72622		Unknown	A_32_P72	1.45756	4.42E-09	1.45756	2825.41	4138.564
A_23_P14	NM_00519	CCAAT/en	CEBPB,LA	1.45743	4.19E-06	1.45743	5864.684	8537.873
A_23_P85	NM_08074	Homo sapi	SRCRB4D	1.45727	5.08E-18	1.45727	4400.903	6414.612
A_23_P8812		Unknown	A_23_P88	1.45713	0.00031	1.45713	1944.45	2832.383
A_24_P16	NM_00677	ring finger	TRIM10,RN	1.4571	0.00004	1.4571	12411.71	17677.44
A_32_P196411		BC008039	THC18667	1.45637	3.67E-07	1.45637	9924.803	14459.49
A_24_P94	AB095940	Family with	FAM22A	1.45582	0.00014	1.45582	1940.275	2799.365
A_32_P194962		Unknown	A_32_P19	1.45579	5.63E-17	1.45579	8074.601	11761.82
A_24_P20	NM_18255	Homo sapi	CNIH2,Cni	1.45538	0.0003	1.45538	1874.859	2681.24
A_32_P19101		Unknown	THC18500	1.45536	0.00003	1.45536	2358.323	3363.329
A_24_P33	NM_15235	hypothetica	MGC40157	1.45535	3.70E-12	1.45535	4146.26	6025.104
A_23_P51	NM_15226	similar to tF	PARS2,MG	1.45512	2.14E-11	1.45512	1449.213	2106.386
A_23_P10	NM_18261	hypothetica	MGC6155(	1.45509	0.00001	1.45509	789.256	1169.18
A_24_P17	NM_18301	Homo sapi	CREM,ICE	1.45497	7.36E-06	1.45497	2626.723	3844.898
A_23_P21	NM_02115	Homo sapi	TRIB3,NIP	1.45458	1.50E-10	1.45458	32703.15	47764.55
A_24_P62	AK000180	Hypothetic	FLJ22679	1.45445	0.00003	1.45445	22016.21	32042.89
A_24_P475115		Unknown	A_24_P47	1.4544	3.19E-09	1.4544	5902.656	8610.48
A_24_P17	NM_00048	aquaporin	AQP2,AQF	1.45423	6.20E-06	1.45423	39593.3	57717.74
A_23_P12	NM_02429	Homo sapi	C20orf149,	1.45355	1.25E-07	1.45355	30171.63	44018.34
A_24_P89	XM_37766	Homo sapi	XM_37766	1.45341	1.93E-08	1.45341	1676.705	2423.728
A_23_P50	NM_03255	Homo sapi	IL1F10,FK	1.4529	2.51E-12	1.4529	1803.201	2616.101
A_24_P99	NM_00131	casein kina	CSNK1G2,	1.45253	9.91E-06	1.45253	9565.877	13966.08
A_32_P36	NM_01780	hypothetica	RASIP1,RA	1.45234	0.00054	1.45234	537.668	798.2502
A_23_P16	NM_00447	Homo sapi	FKBP2,PP	1.45234	1.19E-10	1.45234	18906.79	27365.78
A_24_P358474		Unknown	A_24_P35	1.45214	1.88E-12	1.45214	2285.821	3318.961
A_32_P153361		Unknown	A_32_P15	1.45193	8.03E-06	1.45193	1007.722	1452.136
A_24_P896381		Unknown	THC18044	1.45192	5.54E-09	1.45192	3840.202	5560.715
A_23_P40	NM_14500	Homo sapi	SUSD3,MC	1.45149	2.51E-08	1.45149	3058.831	4428.832
A_24_P31	NM_02523	HZFw1 pro	RNF39,HZ	1.4511	1.30E-06	1.4511	1059.334	1524.512
A_23_P12	NM_00180	carboxyl es	CEL,BAL,F	1.45018	8.41E-09	1.45018	1358.232	1962.206
A_23_P63	NM_13191	Homo sapi	LOC15023	1.45016	0.00001	1.45016	5361.607	7709.592
A_32_P133402		Unknown	A_32_P13	1.44997	2.13E-08	1.44997	1959.936	2832.173
A_24_P32	NM_13836	hypothetica	LOC90799	1.44977	0.00087	1.44977	548.0109	786.3101
A_24_P22	NM_15323	Homo sapi	LIX1,C5orf	1.44913	7.83E-07	1.44913	713.7427	1029.671
A_24_P32	X74954	Mucin 5, su	MUC5B	1.44796	1.86E-10	1.44796	10166.99	14703.34
A_32_P14	AK127789	Hypothetic	MGC21881	1.44792	2.97E-10	1.44792	1960.979	2847.007
A_23_P13	NM_02256	Homo sapi	NYX,CLRF	1.44742	0.00005	1.44742	10097.03	14475.92
A_24_P32	NM_13084	Homo sapi	AMOTL1,J	1.44738	0.00014	1.44738	499.0813	727.6183
A_23_P31	NM_00608	tubulin, bet	TUBB2	1.44627	6.53E-19	1.44627	16737.77	24209.57
A_24_P73	U01925	Human BT	RPL36A,L4	1.44525	1.37E-10	1.44525	2815.042	4070.652

A_24_P92	AJ008188	Isolate RA	VHDJH	1.44515	0.00028	1.44515	521.546	762.5659
A_24_P29	NM_02280	uncoupling	UCP3,SLC	1.44425	5.74E-11	1.44425	5249.786	7578.871
A_23_P31	NM_00529	Homo sapi	GPR25	1.44407	0.00005	1.44407	3541.018	5089.653
A_23_P25	NM_00302	SHB (Src h	SHB	1.44387	4.04E-08	1.44387	873.0811	1259.105
A_24_P19	NM_02430	matrix met	MMP28,M	1.44385	1.35E-08	1.44385	11071.63	15991.76
A_23_P71	NM_00600	Homo sapi	ZA20D2,Z	1.44313	2.86E-13	1.44313	7793.624	11223.01
A_24_P12	NM_00337	VGF nerve	VGF	1.44249	1.73E-07	1.44249	2030.975	2909.807
A_24_P31	NM_00591	MADS box	MEF2B,RS	1.44229	8.40E-06	1.44229	20676.55	29857.41
A_32_P99	NM_00719	Wnt inhibit	WIF1,WIF-	1.44224	0.00014	1.44224	769.0429	1089.861
A_23_P117480		olfactory re	NP389850	1.44172	1.73E-09	1.44172	1124.709	1625.789
A_24_P12	NM_00713	Homo sapi	ZNF79,pT	1.44148	0.00002	1.44148	710.8374	1017.201
A_32_P40611		6PGL_HU	THC18780	1.44137	0.00014	1.44137	836.894	1225.406
A_23_P11	NM_00019	Homo sapi	SHH,HHG	1.44128	0.00009	1.44128	641.0773	936.473
A_32_P11	XM_37149	Homo sapi	XM_37149	1.44061	0.00073	1.44061	1054.986	1508.509
A_24_P38	BC020899	RNA termi	RCL1	1.44045	1.94E-10	1.44045	1883.218	2706.105
A_23_P14	NM_00009	collagen, t	COL7A1,E	1.44032	2.22E-09	1.44032	1336.267	1920.584
A_23_P77	NM_20341	similar to R	LOC92162	1.44005	1.04E-07	1.44005	1905.553	2727.552
A_24_P375573		Unknown	ENST0000	1.43938	1.57E-10	1.43938	4526.919	6504.369
A_23_P10	NM_02192	Homo sapi	SCT	1.43936	2.63E-08	1.43936	1200.677	1738.25
A_23_P21	NM_00592	Homo sapi	MFI2,MTF	1.43932	0.00012	1.43932	845.8959	1192.963
A_23_P15	NM_01662	Arginine/se	MGC1219	1.43812	0.00002	1.43812	750.1774	1073.079
A_23_P30	AF211119	Cyclin-dep	CDKN2A	1.43791	3.47E-12	1.43791	2046.499	2937.248
A_24_P98948		ALU8_HU	THC18915	1.43786	1.14E-08	1.43786	994.9989	1428.823
A_23_P11	NM_14458	Homo sapi	COMTD1,F	1.4372	5.11E-08	1.4372	9504.926	13640.14
A_24_P81	NM_17798	tubulin, bet	TUBB8,FL	1.43663	5.42E-14	1.43663	5579.352	8007.589
A_23_P40	NM_17834	Homo sapi	LCE1A,LE	1.43613	0.00056	1.43613	100617.6	145267.3
A_32_P74	AK095895	Cytochrom	CYP4V2	1.43476	5.44E-14	1.43476	2026.154	2906.24
A_32_P139021		ALU7_HU	THC19532	1.43467	7.80E-08	1.43467	33132.87	47686.55
A_32_P30187		Unknown	A_32_P30	1.4345	0.00037	1.4345	60819.2	87767.58
A_23_P16	NM_00049	Homo sapi	CYP11B1,I	1.43411	5.96E-06	1.43411	11543.49	16478.88
A_32_P95	NM_15234	hypothetic	FLJ30656	1.43409	0.00004	1.43409	2571.537	3690.378
A_23_P16	NM_01910	protein F2	F25965,M	1.43401	1.52E-09	1.43401	6348.243	9091.85
A_32_P15	NM_20683	Homo sapi	CTXN1,CT	1.43352	1.96E-11	1.43352	1221.092	1752.052
A_23_P50	NM_01760	Homo sapi	PPP1R12	1.43338	1.55E-10	1.43338	1809.79	2590.351
A_23_P98	NM_00241	mitogen-ac	MAP3K11,	1.43316	1.56E-08	1.43316	2144.06	3084.353
A_24_P230100		Unknown	A_24_P23	1.43293	1.99E-12	1.43293	3047.462	4371.615
A_24_P57	BC040420	Cortexin 1	CTXN1	1.43232	1.43E-08	1.43232	1078.479	1550.465
A_24_P67	NM_15226	Homo sapi	TPM3,TRK	1.43231	0.00006	1.43231	1387.143	1980.202
A_24_P35	NM_03143	hypothetic	C7orf21,SE	1.43197	5.49E-12	1.43197	6903.45	9879.499
A_32_P41	NM_01787	hypothetic	ASB6,MG	1.43184	9.58E-06	1.43184	701.6713	1002.485
A_24_P358215		Unknown	A_24_P35	1.43155	1.39E-07	1.43155	1224.007	1748.238
A_24_P17	NM_00499	methionine	MARS,ME	1.43131	0.00044	1.43131	9206.221	13261.4
A_32_P49	BC020336	TAO kinas	RPS2	1.43081	3.85E-09	1.43081	1222.534	1750.831
A_32_P224498		Unknown	A_32_P22	1.43064	1.58E-08	1.43064	946.7948	1353.31
A_23_P64	NM_00300	Homo sapi	SELPLG,C	1.43062	4.36E-08	1.43062	3147.697	4432.5
A_24_P99	NM_00492	butyrate re	ZFP36L1,E	1.43041	3.74E-17	1.43041	4230.248	6051.225
A_32_P193952		Unknown	A_32_P19	1.43035	0.00002	1.43035	641.5915	910.0491
A_23_P34	NM_00727	Homo sapi	CTRC,CLC	1.43021	1.03E-07	1.43021	2345.474	3334.351
A_23_P41	NM_15246	Homo sapi	EVER2,EV	1.43004	3.89E-09	1.43004	907.03	1296.725
A_24_P194347		ALU1_HU	THC19160	1.42974	9.43E-09	1.42974	1889.499	2710.117
A_24_P101480		Unknown	ENST0000	1.42927	7.25E-11	1.42927	7061.547	10072.96
A_32_P45	NM_18264	Homo sapi	VGLL2,VG	1.42871	2.50E-08	1.42871	1178.612	1690.646

A_32_P76526		Unknown	A_32_P76	1.42851	7.07E-07	1.42851	1389.822	1999.261
A_23_P49	AB047079	Homo sapi	IRE1b	1.42848	0.00035	1.42848	5110.136	7333.545
A_23_P11	NM_02282	Homo sapi	PORCN,PF	1.42847	3.47E-08	1.42847	9617.955	13738.53
A_24_P76142		Unknown	A_24_P76	1.42828	1.77E-15	1.42828	3671.563	5242.746
A_23_P84	NM_00607	Homo sapi	CACNG2	1.42802	9.64E-06	1.42802	619.9758	880.26
A_23_P78	NM_02163	Homo sapi	ZNF350,ZF	1.42783	0.0008	1.42783	794.8394	1111.982
A_24_P71	AK127768	MRNA; cDI	AK127768	1.42738	0.00028	1.42738	61602.41	87999.48
A_23_P13	NM_00644	Homo sapi	DRAP1,NC	1.42736	0.00005	1.42736	13408.87	19342.21
A_23_P43	NM_00143	Homo sapi	EPAS1,HL	1.42735	0.00035	1.42735	623.4681	902.905
A_24_P51	NM_00213	heterogene	HNRPA1,H	1.42702	1.91E-07	1.42702	6102.257	8653.734
A_24_P931244		Unknown	A_24_P93	1.4268	0.00014	1.4268	581.8298	839.6169
A_24_P24	NM_00448	golgi autoa	GOLGB1,G	1.42617	3.52E-06	1.42617	1116.368	1584.412
A_24_P92	AK000422	CDNA FLJ	AK000422	1.42516	0.00002	1.42516	621.3948	887.6982
A_24_P93	AK097259	Docking pr	DOK3	1.42511	1.28E-06	1.42511	617.2766	885.7507
A_24_P66	BC014173	KIAA0226	KIAA0226	1.42491	0.0006	1.42491	522.673	734.7413
A_23_P34	NM_01804	Homo sapi	GPATC2,F	1.42478	7.02E-06	1.42478	795.1188	1130.011
A_23_P49	NM_00616	Homo sapi	NEUROD2	1.4246	5.94E-09	1.4246	1258.641	1786.071
A_24_P20	NM_13332	Homo sapi	DEDD2,FL	1.42434	0.00021	1.42434	2843.928	3872.004
A_24_P48	NM_03362	JM11 prote	JM11	1.42429	2.06E-09	1.42429	5139.209	7335.145
A_23_P27	NM_01527	ubiquitin-pr	NEDD4L,R	1.4241	0.00008	1.4241	969.7315	1362.59
A_32_P21	NM_13344	Homo sapi	CTGLF1,M	1.42405	0.00017	1.42405	657.4959	949.0892
A_24_P21	NM_01765	Abelson he	AHI1	1.42378	0.00001	1.42378	2843.232	4035.104
A_23_P13	NM_02405	hypothetica	MGC5356	1.42378	0.00007	1.42378	5510.687	7815.542
A_23_P34	NM_17488	Myc-induce	mimitin,B1	1.42371	2.33E-09	1.42371	6575.588	9326.461
A_23_P65	AK098698	Thioredoxi	TXNRD1	1.4232	0.00045	1.4232	967.1355	1356.323
A_23_P21	NM_05300	Homo sapi	GNB1L,GY	1.42284	2.77E-08	1.42284	6352.305	9028.33
A_23_P42	NM_03321	tripartite m	TRIM14,KI	1.42276	1.83E-15	1.42276	2369.974	3372.284
A_23_P37	NM_00692	Homo sapi	SFRS5,HR	1.42251	6.32E-13	1.42251	12737.47	18091.07
A_23_P30	AB051487	Dual specif	DUSP16	1.42235	0.00012	1.42235	855.871	1232.313
A_24_P802562		Unknown	A_24_P80	1.42224	0.00008	1.42224	916.1008	1289.269
A_24_P730256		Unknown	A_24_P73	1.42222	5.08E-11	1.42222	98116.09	139423.5
A_23_P108534		Unknown	A_23_P10	1.42175	4.41E-11	1.42175	1763.537	2507.851
A_23_P16	NM_00253	Homo sapi	ODC1	1.42175	1.60E-12	1.42175	43242.49	61518.26
A_23_P11	NM_14511	Homo sapi	MAP2K3,M	1.42058	0.00075	1.42058	556.0125	799.2161
A_23_P16	NM_00004	apolipoprot	APOE,AD2	1.42057	3.56E-07	1.42057	2698.224	3832.424
A_23_P39	NM_00627	Homo sapi	SFRS7,9G	1.4204	3.82E-14	1.4204	27697.36	39392.7
A_32_P22	BC020462	Ribosomal	RPS9	1.41982	0.00006	1.41982	3806.778	5403.761
A_32_P28480		Seven tran	ENST0000	1.4197	2.48E-11	1.4197	2025.921	2873.329
A_32_P102575		RL35_HUM	THC19457	1.41932	5.88E-10	1.41932	4990.409	7074.298
A_23_P56	NM_13077	Homo sapi	CNTNAP5,	1.41914	0.00018	1.41914	1545.799	2164.964
A_32_P194740		Unknown	A_32_P19	1.41888	0.00006	1.41888	506.7589	718.7191
A_24_P28	NM_03334	tripartite m	TRIM7,GN	1.41856	2.04E-08	1.41856	1589.878	2264.669
A_23_P32	AL136788	Homo sapi	MGC16471	1.41829	0.00003	1.41829	732.621	1029.7
A_24_P61	NM_00224	potassium	KCNMA1,S	1.41822	1.31E-11	1.41822	2959.515	4190.856
A_23_P20	NM_02257	Homo sapi	CSHL1,CS	1.41808	1.94E-08	1.41808	2395.2	3385.038
A_32_P11	BF368083	Cyclin H	CCNH	1.41772	2.12E-06	1.41772	76512.17	108932.8
A_23_P20	NM_01425	Homo sapi	CLEC4M,C	1.4175	6.36E-06	1.4175	10827.34	15308.56
A_24_P36	NM_02006	GLUT4 enh	SLC2A4RC	1.41714	1.64E-12	1.41714	1946.372	2756.322
A_32_P88905		Unknown	A_32_P88	1.41655	8.93E-07	1.41655	1752.466	2475.455
A_23_P49	NM_00208	Granulin	GRN	1.41644	0.00004	1.41644	43241.71	61596.72
A_23_P21	NM_19920	ubiquitin-co	Kua-UEV,U	1.41635	3.14E-12	1.41635	5032.97	7137.155
A_32_P10	AK091337	CDNA FLJ	AK091337	1.41615	0.00001	1.41615	1210.147	1699.277

A_23_P35	NM_00106	Homo sapi	TBXA2R,T	1.41607	8.75E-06	1.41607	6605.972	9363.307
A_23_P81	AF022375	Vascular e	VEGF	1.41599	0.00001	1.41599	2079.842	2876.054
A_23_P90	NM_18291	TIR domai	TICAM1,TI	1.41587	2.08E-06	1.41587	2364.907	3342.36
A_23_P36	BC011940	CDNA clon	BC011940	1.41572	0.00006	1.41572	2881.427	4068.091
A_23_P43	L78790	Integrin, be	ITGB2	1.41572	3.52E-08	1.41572	1143.666	1610.224
A_23_P14	NM_01592	liver-specif	LISCH7,LS	1.41516	9.02E-06	1.41516	1062.405	1516.678
A_23_P11	NM_00150	Homo sapi	CXCR3,GF	1.41495	0.00016	1.41495	1854.903	2602.407
A_23_P32	NM_17822	Homo sapi	IQGAP3,M	1.41471	1.53E-09	1.41471	3959.474	5604.632
A_24_P14	NM_17355	Homo sapi	FGD2,ZFY	1.41451	4.02E-07	1.41451	926.0777	1303.642
A_24_P49800		Unknown	A_24_P49	1.41434	6.80E-17	1.41434	20303.38	28713.67
A_24_P45	AK123446	CDNA FLJ	AK123446	1.41411	0.00082	1.41411	1676.764	2335.998
A_32_P132422		Unknown	A_32_P13	1.41407	2.14E-09	1.41407	506338.7	716342.1
A_32_P13	AK000028	Hypothetic	FLJ20021	1.41377	7.92E-11	1.41377	4448.862	6287.105
A_23_P21	NM_02311	fibroblast g	FGFR1,H2	1.41373	0.00097	1.41373	593.2744	818.1829
A_23_P89	NM_02215	Homo sapi	FN3K	1.41346	2.21E-10	1.41346	1567.086	2211.729
A_24_P91	BC009479	Homo sapi	TTLL3	1.41285	0.00016	1.41285	10758.74	15238.25
A_23_P59	NM_00008	Homo sapi	CLCN1,CL	1.41264	6.56E-07	1.41264	5506.547	7755.199
A_32_P37	BC008580	Homo sapi	BC008580	1.4117	0.00004	1.4117	650.8867	914.8389
A_24_P840868		Unknown	A_24_P84	1.41158	1.76E-06	1.41158	70917.78	100909.4
A_23_P36	NM_00117	ATP-bindin	ABCC6,AR	1.4115	3.25E-10	1.4115	1433.895	2020.056
A_24_P27	NM_01906	chromoson	EML4,C2o	1.41149	2.72E-15	1.41149	7266.857	10262.69
A_24_P15	NM_00374	insulin rece	IRS2	1.41145	8.96E-07	1.41145	3037.961	4309.886
A_24_P884506		Unknown	A_24_P88	1.41086	9.29E-13	1.41086	1433.984	2024.584
A_32_P180610		Unknown	THC18394	1.41067	0.00054	1.41067	491.9288	688.9285
A_24_P15	NM_14501	hypothetica	FLJ30707	1.41057	0.00004	1.41057	12001.28	17045.33
A_23_P16	NM_03225	hypothetica	TBC1D3,P	1.41041	0.00064	1.41041	1199.349	1669.09
A_24_P23	NM_00411	Homo sapi	FKHL18,FI	1.41022	0.00018	1.41022	2787.537	3831.16
A_24_P91	NM_03283	hypothetica	FLJ14775	1.40978	0.00007	1.40978	2135.352	2995.213
A_24_P31	BC040620	Thrombos	THSD6	1.40971	0.00026	1.40971	715.8654	1025.648
A_23_P34	NM_13844	hypothetica	LOC11509	1.40947	8.78E-15	1.40947	6527.688	9199.913
A_23_P16	NM_00595	Homo sapi	MT1H,MT1	1.40934	0.00007	1.40934	9900.49	13994.29
A_24_P37	NM_00097	ribosomal	RPL12	1.40919	2.13E-15	1.40919	56648.28	79771.17
A_23_P35	NM_02196	Homo sapi	TCL1A,TC	1.40891	5.63E-07	1.40891	2749.025	3856.781
A_23_P11	NM_00272	Homo sapi	PRB4,Po	1.40803	5.36E-16	1.40803	6186.879	8710.523
A_23_P90	NM_14896	Homo sapi	OTOS,OTC	1.40755	7.59E-08	1.40755	3220.85	4518.543
A_32_P111492		ALU5_HUM	THC18966	1.40611	0.00012	1.40611	652.3842	908.4447
A_23_P14	NM_01757	G protein-c	MKNK2,MI	1.40605	0.00074	1.40605	28400.16	40400.29
A_32_P26034		ALU5_HUM	THC20049	1.40576	1.94E-06	1.40576	4708.436	6626.048
A_32_P10	BC034930	Homo sapi	BC034930	1.40563	1.25E-06	1.40563	983.2163	1390.403
A_24_P585004		Unknown	A_24_P58	1.40561	7.41E-12	1.40561	2418.156	3396.298
A_24_P17	NM_00435	cell divisior	CDC34,UB	1.40542	4.65E-11	1.40542	1487.814	2090.098
A_32_P85	BC008632	Chromosor	C6orf176	1.4051	1.22E-06	1.4051	5624.193	7955.153
A_23_P34	NM_02109	Calcium ch	CACNA1H	1.40493	2.98E-13	1.40493	5005.959	7026.984
A_24_P28	AL050020	DKFZP564	DKFZP564	1.40454	2.35E-07	1.40454	10222.56	14376.9
A_24_P92	L02922	Human tro	tropomyosi	1.40438	3.13E-07	1.40438	1101.969	1551.108
A_24_P482124		Unknown	THC18096	1.40432	3.60E-07	1.40432	987.7697	1391.073
A_23_P79	NM_00163	Homo sapi	ALPP,ALP	1.40421	6.54E-07	1.40421	3263.643	4569.038
A_24_P19	NM_05305	Homo sapi	CCND1,BC	1.40413	3.47E-14	1.40413	3936.037	5527.108
A_24_P15	AF370409	Solute carr	SLC30A2	1.40405	8.72E-09	1.40405	1814.214	2556.091
A_24_P25	XM_37331	similar to ri	LOC39238	1.40397	1.10E-08	1.40397	10812.98	15212.62
A_32_P14	BC015064	Transduce	TOB1	1.40381	1.60E-06	1.40381	1209.917	1681.59
A_23_P14	NM_01893	Homo sapi	PCDHB13,	1.40365	0.00002	1.40365	8672.359	12146.16

A_32_P15706		Unknown	A_32_P15	1.40363	2.14E-07	1.40363	6748.482	9461.662
A_23_P14	NM_01780	nuclear RN	NXF2,TAP	1.40323	4.96E-07	1.40323	1375.444	1919.741
A_23_P48	NM_00553	Homo sapi	IFI27,P27,I	1.40279	3.67E-09	1.40279	1007.335	1417.015
A_23_P37	AF155102	Chromosor	C19orf29	1.40279	0.00002	1.40279	1340.95	1860.022
A_24_P92	NM_19929	Tyrosine h	TH	1.40241	0.00001	1.40241	5435.166	7614.906
A_23_P20	NM_01221	integrin, al	ITGA11,Hs	1.40204	0.00019	1.40204	776.9609	1098.064
A_24_P21	NM_18263	Homo sapi	SLC6A18,X	1.40182	6.70E-09	1.40182	3971.734	5558.234
A_24_P76	BC009051	Homo sapi	BC009051	1.40132	0.00003	1.40132	1160.343	1663.601
A_23_P80	NM_00651	SET domai	SETMAR	1.40121	0.00088	1.40121	2011.132	2822.031
A_23_P10	NM_18253	hypothetica	MGC29671	1.40091	4.83E-06	1.40091	1917.918	2669.468
A_24_P11	AK074050	Homo sapi	FLJ00121	1.40085	0.00082	1.40085	520.3765	736.6411
A_24_P28	NM_17817	Homo sapi	TREML1,T	1.40076	0.00022	1.40076	509.4796	708.2738
A_23_P25	NM_01761	Homo sapi	RNPC3,RN	1.40076	2.89E-06	1.40076	5637.883	7823.578
A_24_P34	AK001784	Homo sapi	FLJ10922	1.40051	0.00024	1.40051	1457.269	2022.802
A_32_P49035		Unknown	A_32_P49	1.40002	7.52E-08	1.40002	2415.433	3378.003
A_32_P64	BC033996	Homo sapi	ENST0000	1.39961	0.00006	1.39961	5047.28	7026.49
A_23_P15	AK021957	Collagen, t	COL27A1	1.39885	0.00003	1.39885	2093.156	2907.879
A_24_P247576		Unknown	ENST0000	1.39876	4.24E-10	1.39876	3139.671	4386.692
A_24_P890995		Unknown	A_24_P89	1.3975	8.69E-11	1.3975	1689.614	2361.298
A_24_P42	AK056449	Myozenin 3	MYOZ3	1.39732	0.00004	1.39732	1458.619	2021.216
A_23_P10	NM_15231	glycosyltra	GYLTL1B,I	1.39715	1.04E-09	1.39715	4458.379	6212.98
A_24_P34	BC012882	Hypothetic	LOC15035	1.39673	1.02E-08	1.39673	1287.288	1800.203
A_23_P20	NM_01400	vinculin	VCL	1.39613	0.00034	1.39613	1070.083	1474.581
A_32_P18	BX641020	AT rich inte	ARID5B	1.39588	0.00062	1.39588	2621.719	3633.997
A_24_P10	AK094893	Fucosyltra	FUT8	1.39521	0.00014	1.39521	1055.017	1455.512
A_24_P709949		Unknown	A_24_P70	1.3949	0.00037	1.3949	703.9734	996.1863
A_24_P11	NM_00122	caspase 9,	CASP9,MC	1.39484	9.77E-08	1.39484	3526.8	4933.533
A_23_P19	NM_00338	Homo sapi	VIP,PHM2	1.39477	0.00004	1.39477	650.5791	901.4764
A_24_P92	NM_01404	DKFZP564	LRP10,MS	1.3946	1.35E-09	1.3946	4559.594	6351.77
A_23_P60009		Annexin A1	ANXA13	1.39435	0.00001	1.39435	647.218	900.8067
A_23_P44	NM_01848	developme	DDEF1,PA	1.39433	0.00028	1.39433	657.2638	930.8915
A_24_P13	NM_00269	POU doma	POU3F1,O	1.39414	1.83E-06	1.39414	1109.39	1565.203
A_32_P118686		AF273048	THC18261	1.39378	0.00004	1.39378	2738.238	3815.913
A_23_P41	NM_00363	contactin a	CNTNAP1,	1.39366	8.24E-10	1.39366	2329.747	3241.051
A_23_P14	NM_00095	Homo sapi	PTGDS,PD	1.39341	5.88E-10	1.39341	4835.757	6727.914
A_23_P12	NM_01489	Homo sapi	PDAP1,PA	1.39329	8.56E-07	1.39329	2223.101	3089.198
A_32_P190802		ALU1_HUM	THC19316	1.39268	0.0004	1.39268	534.2174	741.139
A_24_P35	NM_00211	major histo	HLA-DOA,I	1.39256	0.00002	1.39256	4764.046	6757.041
A_24_P418386		Unknown	ENST0000	1.39216	0.00005	1.39216	1076.419	1485.989
A_24_P28	NM_01433	peptidylpro	PPIL2,CYC	1.39214	0.00017	1.39214	795.9075	1121.625
A_24_P578445		Unknown	A_24_P57	1.39188	5.79E-15	1.39188	6338.944	8823.83
A_23_P37	NM_14517	Homo sapi	ATOH7,Ma	1.39161	1.09E-07	1.39161	1197.272	1671.917
A_23_P14	NM_00002	Homo sapi	ADRB2,BA	1.39118	0.00034	1.39118	784.5447	1076.442
A_24_P37	NM_20738	FLJ41287	FLJ41287	1.39077	1.73E-06	1.39077	1080.668	1507.144
A_24_P615207		Unknown	THC19467	1.39039	9.69E-07	1.39039	1482.347	2046.809
A_23_P35	NM_00395	Homo sapi	SOCS3,CI	1.38995	0.0001	1.38995	1844.709	2508.973
A_23_P59	NM_00109	amiloride b	ABP1,ABP	1.3899	2.52E-10	1.3899	1102.271	1534.963
A_32_P150950		RS28_HUM	THC18329	1.3898	1.84E-09	1.3898	3179.997	4418.552
A_23_P82	NM_00392	Homo sapi	FOXH1,FA	1.38974	1.45E-09	1.38974	2063.688	2865.199
A_23_P30	NM_13836	hypothetica	LOC90799	1.38969	0.00012	1.38969	3744.94	5253.497
A_24_P35	AK091483	Hypothetic	LOC33977	1.38965	0.00018	1.38965	748.5122	1039.907
A_23_P50	NM_17237	Homo sapi	IL4I1,FIG1	1.38964	0.00097	1.38964	706.1663	964.5216

A_23_P16	NM_03215	PRAM-1 pr	PRAM1,PF	1.38928	0.00023	1.38928	3331.521	4642.459
A_32_P98227		C40201 art	THC18603	1.38917	9.48E-09	1.38917	1684.573	2333.093
A_23_P17	AL096750	5-oxoprolin	OPLAH	1.38878	0.00002	1.38878	761.5534	1055.479
A_32_P125219		Unknown	A_32_P12	1.38868	0.00004	1.38868	1088.433	1494.181
A_32_P10	XM_05992	similar to N	LOC13782	1.38847	0.00002	1.38847	983.3302	1354.657
A_23_P16	NM_00522	Homo sapi	ARID3A,DI	1.38815	0.00013	1.38815	1004.203	1376.337
A_23_P21	NM_18148	Homo sapi	ZGPAT,ZC	1.38805	0.00003	1.38805	5858.92	8185.018
A_24_P90	BQ188387	Hypothetic	MGC2747	1.38778	0.00062	1.38778	573.7562	792.9292
A_24_P401069		Unknown	A_24_P40	1.38757	5.38E-08	1.38757	1326.448	1846.293
A_23_P21	D13814	Angiotensi	AGTR1	1.38749	0.00071	1.38749	560.452	787.3024
A_23_P38	NM_01527	ubiquitin-pr	NEDD4L,R	1.38741	0.00094	1.38741	706.0165	966.5263
A_23_P35	M34432	Pvt1 oncog	PVT1	1.38739	5.84E-08	1.38739	39703.84	54846.32
A_24_P341006		Unknown	A_24_P34	1.38687	0.0004	1.38687	4575.393	6356.859
A_23_P31	NM_03143	Homo sapi	C7orf21,SE	1.38682	0.00007	1.38682	6237.899	8671.325
A_32_P20	AW168145	Transcribe	TUBB2	1.38626	0.00002	1.38626	1655.457	2286.864
A_23_P14	NM_00545	Homo sapi	PDLIM7	1.38616	0.00033	1.38616	991.4448	1352.499
A_23_P54	NM_00555	Homo sapi	KIFC3,DKF	1.38604	0.00008	1.38604	1300.254	1819.178
A_32_P217346		Unknown	A_32_P21	1.38563	1.80E-10	1.38563	2440.483	3389.051
A_23_P13	NM_00217	insulin-like	IGFBP6,IB	1.38552	1.67E-09	1.38552	2569.457	3556.481
A_23_P14	NM_08074	Homo sapi	C20orf52,M	1.3851	0.00007	1.3851	38800.42	54156.68
A_23_P10	NM_05821	Homo sapi	EXOSC6,p	1.38503	2.33E-08	1.38503	4167.5	5753.051
A_23_P15	NM_00514	Homo sapi	GNG7,FLJ	1.38501	0.00002	1.38501	3152.101	4359.271
A_23_P123234		Unknown	A_23_P12	1.38487	3.00E-06	1.38487	439.7997	608.2134
A_23_P85	NM_00508	Homo sapi	U2AF1L2,U	1.38479	4.19E-11	1.38479	2443.693	3381.025
A_32_P197611		Unknown	THC18426	1.38445	2.60E-11	1.38445	20734.79	28732.97
A_24_P81	AK074088	Tumor prot	TP53I5	1.3844	0.001	1.3844	624.8877	879.9793
A_32_P12	NM_03098	small GTP-	RAB1B	1.38392	2.31E-06	1.38392	9488.932	13179.49
A_23_P74	NM_03321	Zinc finger	MGC12466	1.38345	1.28E-07	1.38345	1309.341	1816.673
A_24_P38	NM_17506	Homo sapi	RASGEF1	1.38297	3.68E-07	1.38297	1178.163	1613.279
A_32_P13	AB051464	Kelch-like	KLHL15	1.38252	0.00003	1.38252	1076.602	1481.263
A_23_P21	NM_02210	G-protein s	GPSM3	1.38168	5.24E-12	1.38168	6515.354	8992.307
A_24_P59	BX109136	BX109136	BX109136	1.38159	7.49E-06	1.38159	774.9858	1072.26
A_24_P47	U96394	Ig lambda	U96394	1.38154	8.17E-08	1.38154	3316.003	4598.182
A_32_P34480		Unknown	A_32_P34	1.38118	5.69E-10	1.38118	3692.515	5098.899
A_23_P16	NM_00134	Homo sapi	DAPK3,ZIF	1.38056	3.12E-06	1.38056	1166.524	1615.665
A_24_P920081		Unknown	A_24_P92	1.38051	0.0002	1.38051	1794.922	2402.519
A_32_P170104		Unknown	A_32_P17	1.37978	3.55E-10	1.37978	1469.823	2028.474
A_23_P14	NM_02173	Homo sapi	AVPI1,VIP	1.37959	7.05E-09	1.37959	11865.36	16418.13
A_24_P40	XM_37231	Homo sapi	XM_37231	1.37952	2.44E-09	1.37952	42049.19	57859.97
A_24_P14	NM_00352	H2B histon	HIST2H2B	1.37901	0.00016	1.37901	1607.843	2198.692
A_23_P10	NM_00630	glucosidas	GCS1	1.37901	0.0005	1.37901	31847.66	44222.42
A_23_P91	NM_00241	Homo sapi	MIF,GIF,GI	1.37883	2.02E-10	1.37883	159247.7	220132.1
A_23_P53	NM_00258	Homo sapi	PAWR,PAI	1.37848	8.91E-11	1.37848	5280.915	7289.8
A_23_P34	NM_17074	Homo sapi	C11orf31,S	1.37846	1.35E-09	1.37846	29392.6	40674.88
A_23_P10	NM_02235	potassium	KCNK15,K	1.37838	0.00047	1.37838	21513.67	29466.41
A_24_P37	NM_00197	eukaryotic	EIF5A,uOF	1.37838	1.98E-06	1.37838	3704.724	5136.776
A_23_P20	NM_01480	KIAA0435	PCNXL2,F	1.37819	2.93E-06	1.37819	2066.484	2843.055
A_24_P565496		Unknown	A_24_P56	1.37797	0.00002	1.37797	1313.716	1784.911
A_23_P10	NM_03081	hypothetica	MGC10471	1.37792	3.58E-06	1.37792	7478.68	10330.45
A_32_P79	BG771581	602719909	BG771581	1.37741	0.00094	1.37741	3576.648	5062.22
A_23_P21	NM_00712	Utrophin (h	UTRN	1.37726	0.00004	1.37726	627.4771	861.3934
A_32_P175667		Unknown	THC18384	1.37718	4.70E-07	1.37718	1586.955	2199.432

A_32_P177725		Unknown	THC19357	1.37711	1.19E-07	1.37711	3317.576	4571.498
A_23_P79	NM_00639	nucleolar p	NOL5A,NC	1.37674	2.90E-08	1.37674	7156.152	9819.798
A_23_P11	NM_00299	Homo sapi	CXCL5,SC	1.3767	7.57E-06	1.3767	2228.772	3073.448
A_24_P30	BC056606	Similar to V	MGC5735	1.37658	7.01E-07	1.37658	917.652	1261.05
A_23_P36	NM_00533	Homo sapi	HBZ	1.37578	9.69E-06	1.37578	1136.619	1550.192
A_23_P63	NM_00614	stratifin	SFN	1.37578	0.00024	1.37578	1189.751	1653.269
A_24_P31	BC005365	Homo sapi	MAP2K7,M	1.3755	2.08E-07	1.3755	1011.384	1387.144
A_24_P850069		Unknown	A_24_P85	1.37521	1.93E-07	1.37521	2902.129	3978.303
A_32_P6062		Unknown	A_32_P60	1.37493	0.00002	1.37493	2638.265	3538.727
A_32_P44139		Unknown	A_32_P44	1.37483	0.00002	1.37483	1992.243	2744.663
A_32_P36	AK125884	Similar to A	AK125884	1.37467	3.33E-07	1.37467	999.1123	1370.656
A_24_P255786		Unknown	A_24_P25	1.37461	1.08E-09	1.37461	17011.81	23359.48
A_24_P94	BG259015	Transcribe	BG259015	1.3746	0.00002	1.3746	1146.276	1588.601
A_24_P13	NM_02303	calcium ch	CACNA1A	1.37456	0.00011	1.37456	2480.465	3392.636
A_23_P39	NM_00438	Homo sapi	NKX2-5,CS	1.37435	4.17E-09	1.37435	6898.927	9464.951
A_32_P157775		AF193342	THC19565	1.3736	7.62E-12	1.3736	2004.048	2752.946
A_23_P14	XM_17299	similar to H	LOC25580	1.37359	2.97E-09	1.37359	3050.656	4180.479
A_23_P91	NM_00306	Homo sapi	SLPI,ALP,I	1.37357	8.45E-06	1.37357	37057.21	51222.03
A_24_P33217		Unknown	A_24_P33	1.37303	3.62E-06	1.37303	10959.25	14919.86
A_32_P11	AW378392	Transcribe	RPL39	1.37296	0.00035	1.37296	6074.24	8309.281
A_23_P30	NM_13829	Homo sapi	PTCRA,PT	1.37294	0.00001	1.37294	2308.292	3148.579
A_32_P17	NM_20685	Homo sapi	MGC8714	1.37293	3.63E-07	1.37293	1298.926	1788.027
A_23_P24	NM_00405	Homo sapi	C3AR1,AZ	1.37286	8.29E-09	1.37286	3276.081	4494.885
A_23_P40	BC021238	Hypothetic	MGC1976	1.37216	0.00001	1.37216	929.011	1276.291
A_24_P74	NM_01501	Amine oxid	AOF2	1.37191	0.00008	1.37191	775.4688	1056.824
A_24_P91	U62539	Human B c	PAX5	1.37181	2.23E-07	1.37181	1277.399	1745.505
A_23_P42	NM_03341	hypothetic	ARMC6,M	1.37152	0.00008	1.37152	17922.34	24637.49
A_23_P90	NM_02246	Homo sapi	CHST8	1.37094	3.20E-06	1.37094	4768.338	6499.424
A_32_P42537		Unknown	THC18710	1.37034	0.00038	1.37034	654.9264	888.7376
A_23_P99	NM_19821	Homo sapi	ING1,p33,p	1.3701	0.00002	1.3701	1120.457	1548.644
A_23_P21	AF336133	Homo sapi	CECR2	1.36993	1.40E-08	1.36993	5204.71	7110.343
A_23_P16	NM_01223	Homo sapi	POMZP3,F	1.36955	5.84E-08	1.36955	4318.255	5887.386
A_23_P10	NM_00221	Homo sapi	ITIH4,H4P,	1.36945	7.72E-06	1.36945	1068.98	1455.632
A_23_P25	NM_00664	Homo sapi	NOXA1,p5	1.36941	4.66E-09	1.36941	1295.436	1774.761
A_23_P82	NM_00460	Homo sapi	STX1A,ST	1.3694	0.00043	1.3694	2896.119	3938.587
A_32_P14	AK090602	Hypothetic	LOC28314	1.36928	0.00018	1.36928	873.546	1209.822
A_32_P21	AK125817	Hypothetic	LOC28406	1.36909	0.00026	1.36909	895.7559	1243.581
A_23_P38	NM_01838	hypothetic	FLJ11286	1.36862	0.00048	1.36862	3666.527	5040.642
A_23_P13	NM_00601	Homo sapi	ARMET,AF	1.36821	1.78E-07	1.36821	83657.27	114763.2
A_23_P16	AK128435	Cofactor re	CRSP7	1.3679	0.00099	1.3679	1521.873	2064.176
A_32_P93	XM_37112	Homo sapi	XM_37112	1.36789	0.00011	1.36789	5144.729	7006.727
A_23_P11	NM_00096	Homo sapi	RARG,RAF	1.36734	0.00006	1.36734	1768.995	2395.281
A_23_P54	NM_08067	dynein ligh	Dlc2,MGC	1.36693	5.15E-07	1.36693	4063.522	5578.12
A_23_P12	NM_00255	Homo sapi	P2RX3,P2	1.36689	0.00032	1.36689	3062.793	4179.397
A_23_P75	NM_02122	Homo sapi	ARHGAP2	1.36677	3.54E-06	1.36677	1334.895	1830.617
A_23_P78	NM_02021	carcinoemb	CEACAM1	1.36666	2.94E-06	1.36666	21383.07	29148.38
A_23_P109224		SP4L_HUM	THC18919	1.36633	0.00008	1.36633	980.4137	1318.915
A_24_P298886		Unknown	A_24_P29	1.36612	6.54E-13	1.36612	4916.781	6714.086
A_32_P47	NM_14565	Homo sapi	GSH1,Gsh	1.36581	3.35E-12	1.36581	2469.553	3373.125
A_23_P42	NM_03200	protocadhe	PCDHGA2	1.3656	0.00008	1.3656	2821.709	3840.171
A_23_P11	NM_00633	Homo sapi	LRRN5,GA	1.36559	0.00002	1.36559	9756.736	13348.93
A_23_P21	NM_02063	ankyrin rep	RIPK4,DIK	1.36544	1.08E-07	1.36544	1848.487	2512.705

A_24_P28	NM_02126	Homo sapi	ZFYVE1,D	1.36525	0.0001	1.36525	1047.449	1415.62
A_24_P56	XM_20889	Homo sapi	XM_20889	1.36508	0.00003	1.36508	2417.659	3283.459
A_24_P67	AF035028	HRV Fab 0	AF035028	1.36506	0.00014	1.36506	757.1433	1042.692
A_24_P91	NM_20337	hypothetica	LOC28491	1.3649	0.00008	1.3649	947.1643	1304.361
A_23_P11	NM_00102	Homo sapi	RPS19,DB	1.36477	1.35E-13	1.36477	66950.76	91400.74
A_32_P21	BM473515	LOC44073	BM473515	1.36462	0.00003	1.36462	2987.968	4099.343
A_23_P12	NM_00201	FK506-bin	FKBP4,HB	1.36427	0.00063	1.36427	10493.67	14427.02
A_23_P43	NM_00140	Homo sapi	EFNB3,EF	1.36411	0.00002	1.36411	7994.114	10890.97
A_23_P32	NM_02504	Homo sapi	C15orf20,F	1.36378	2.58E-06	1.36378	907.7412	1226.975
A_24_P678056		Unknown	A_24_P67	1.36321	2.40E-10	1.36321	2754.819	3757.941
A_23_P26	AK026675	Phosphatid	PEMT	1.36292	0.00008	1.36292	8641.862	11795.03
A_24_P110541		Unknown	A_24_P11	1.36284	6.58E-10	1.36284	42574.29	58149.63
A_23_P70	Z33459	Major histo	HLA-C	1.36246	0.00002	1.36246	1944.118	2653.386
A_23_P40	NM_15222	Homo sapi	CSNK1E,H	1.36228	0.00005	1.36228	11324.27	15545.58
A_24_P27	NM_00553	interferon,	IFI27,P27,I	1.36177	0.00009	1.36177	808.454	1097.169
A_23_P11	NM_00633	Homo sapi	HMG20B,S	1.36122	1.13E-07	1.36122	1302.588	1774.769
A_23_P10	NM_03285	Homo sapi	CORO6,FL	1.36095	3.44E-08	1.36095	4540.021	6186.607
A_32_P206751		Unknown	A_32_P20	1.36087	2.59E-06	1.36087	4966.382	6767.197
A_23_P67	NM_00070	Homo sapi	BCKDHA,M	1.36053	0.00012	1.36053	14978.15	20399.23
A_32_P196142		Unknown	THC19333	1.36028	0.00009	1.36028	1324.985	1773.916
A_24_P10	NM_01767	hypothetica	BCAS3,MA	1.36015	3.21E-07	1.36015	1564.555	2136.004
A_24_P69	BC014776	Hypothetic	MGC4677	1.3599	0.00008	1.3599	7079.807	9726.033
A_24_P887805		Unknown	A_24_P88	1.35973	1.06E-10	1.35973	1805.712	2459.959
A_23_P70	NM_00060	Homo sapi	NOS3,eNC	1.35969	3.93E-08	1.35969	3507.66	4754.09
A_24_P89	NM_00344	zinc finger	TRIM26,AF	1.35968	1.04E-06	1.35968	4574.651	6233.835
A_24_P37	XM_37271	Homo sapi	XM_37271	1.35962	7.12E-07	1.35962	38113.44	52022.84
A_32_P120534		Unknown	A_32_P12	1.35937	1.36E-10	1.35937	5201.983	7079.659
A_23_P38	NM_17379	Hypothetic	LOC12897	1.35936	4.86E-06	1.35936	1376.407	1867.575
A_24_P65	AK125790	Hypothetic	LOC40135	1.35906	7.62E-06	1.35906	8100.993	10980.63
A_24_P212314		Unknown	A_24_P21	1.35846	1.19E-06	1.35846	12353.92	16829.81
A_24_P52	BI910970	Similar to K	BI910970	1.35845	3.99E-07	1.35845	2433.743	3306.07
A_23_P21	NM_00497	Homo sapi	JARID2,JM	1.35837	0.00011	1.35837	2549.412	3441.883
A_23_P13	NM_00291	renin-bindi	RENBP,RE	1.35816	8.38E-07	1.35816	1175.048	1589.459
A_23_P38	BC022076	Hypothetic	LOC25578	1.35774	0.00071	1.35774	1581.38	2121.594
A_23_P13	NM_02468	Homo sapi	TBC1D17,I	1.35724	0.00032	1.35724	2503.495	3383.416
A_23_P42	AB051475	KIAA1688	KIAA1688	1.35714	0.00083	1.35714	709.6149	951.1066
A_23_P13	NM_03091	semaphori	SEMA6C,S	1.3569	0.00005	1.3569	2092.426	2819.792
A_24_P161393		Unknown	A_24_P16	1.3566	5.17E-12	1.3566	22224.82	30164.89
A_23_P31	BC034629	Hypothetic	MGC4271	1.35643	1.69E-07	1.35643	1782.141	2423.715
A_24_P22	AK025047	SID1 trans	SIDT1	1.35638	0.00017	1.35638	4394.114	5977.838
A_24_P35	NM_01456	Homo sapi	OTX1,FLJ3	1.35631	0.00021	1.35631	1780.661	2392.47
A_23_P31	NM_14462	Homo sapi	DCST2,FL	1.35629	5.24E-09	1.35629	4631.404	6264.943
A_24_P80	BC041359	Hypothetic	MGC2477	1.35623	0.00005	1.35623	1716.748	2322.713
A_23_P11	NM_00422	Homo sapi	PSCD2,AR	1.35598	0.00002	1.35598	1112.859	1498.787
A_24_P771173		Unknown	A_24_P77	1.35591	5.90E-06	1.35591	3926.247	5308.353
A_24_P41	AL133632	Similar to h	DKFZp434	1.35573	0.00019	1.35573	21278.07	28789.21
A_24_P92	AK021842	FLJ25476	FLJ25476	1.35564	5.98E-08	1.35564	11841.48	16040.17
A_24_P11	AK130207	CDNA FLJ	ENST0000	1.35543	2.40E-11	1.35543	16763.71	22698.22
A_23_P213350		Unknown	A_23_P21	1.35517	0.00073	1.35517	808.0635	1109.955
A_23_P82	NM_02465	hypothetica	PRKRIP1,(	1.35471	0.00005	1.35471	1163.754	1586.347
A_24_P26	NM_01443	NADPH-de	NDOR1,NF	1.3544	0.00023	1.3544	12154.57	16556.54
A_23_P16	NM_02104	Homo sapi	DHH,HHG-	1.35368	5.32E-08	1.35368	3442.872	4663.66

A_23_P105732		Unknown	A_23_P10	1.35356	5.92E-07	1.35356	27568.56	37315.88
A_23_P36	BC009220	LOC44116	C6orf160,M	1.35334	2.40E-07	1.35334	69005.42	93823.25
A_23_P12	NM_00642	Homo sapi	SLC34A2,M	1.35332	0.00013	1.35332	2219.583	2993.285
A_32_P132337		Unknown	A_32_P13	1.35313	3.06E-11	1.35313	24298.2	32900.5
A_32_P22	BM994413	Thymosin,	TMSB10	1.35299	9.96E-07	1.35299	7810.757	10557.45
A_23_P10	NM_13917	MDAC1	MDAC1	1.35299	0.00003	1.35299	1369.616	1872.479
A_23_P29	NM_00407	crystallin, b	CRYBB3,C	1.35292	0.00001	1.35292	5548.039	7480.862
A_32_P22	NM_05819	Homo sapi	C21orf70,F	1.35286	0.00001	1.35286	3915.311	5326.098
A_32_P170847		AP003829	THC18060	1.35263	1.98E-08	1.35263	2173.529	2935.538
A_23_P29	AK022953	Hypothetica	FLJ35725	1.35243	0.00035	1.35243	2687.938	3613.602
A_23_P90	NM_13839	Homo sapi	C19orf32,D	1.35243	3.91E-10	1.35243	4852.564	6578.632
A_24_P26	NM_00325	thyroid hor	THRA,AR7	1.35231	1.53E-09	1.35231	2033.084	2747.15
A_24_P33	NM_02112	polymerase	POLR2L,R	1.35208	7.66E-06	1.35208	26566.54	36203.45
A_23_P13	NM_01807	hypothetica	FLJ10374	1.3519	0.00017	1.3519	1119.979	1501.244
A_32_P192389		Unknown	A_32_P19	1.35166	0.00002	1.35166	1064.988	1448.395
A_23_P80	AK097004	Homo sapi	SFI1,PISD,	1.35163	5.03E-10	1.35163	1991.016	2691.111
A_24_P55	NM_05300	odd-skippe	OSR2,FLJ	1.35149	6.47E-06	1.35149	831.6392	1121.781
A_23_P34	NM_05819	Homo sapi	TRIM29,AT	1.35136	3.82E-07	1.35136	1837.544	2474.98
A_32_P70	BC015772	Homo sapi	BC015772	1.35134	0.00029	1.35134	2818.666	3791.817
A_24_P10	BC013171	Melanoma	MAGEA12	1.35132	2.71E-09	1.35132	3763.779	5087.476
A_24_P15	AK096229	CDNA FLJ	AK096229	1.35113	0.00013	1.35113	18113.7	24495.36
A_24_P35	NM_01516	Homo sapi	ARL6IP,AII	1.35103	2.38E-10	1.35103	25751.51	34729.5
A_32_P83	NM_20738	FLJ31222	FLJ31222	1.35092	0.00002	1.35092	3247.452	4373.008
A_32_P28	XM_05858	Homo sapi	MGC90512	1.35092	4.70E-08	1.35092	4888.417	6599.776
A_23_P13	NM_02517	Homo sapi	DEPDC2,D	1.35089	1.12E-06	1.35089	1727.091	2323.002
A_23_P37	NM_14471	Homo sapi	CCDC12,F	1.3504	5.11E-08	1.3504	7381.918	9982.347
A_32_P124125		Unknown	A_32_P12	1.35037	7.00E-10	1.35037	4762.755	6428.359
A_32_P141338		Unknown	A_32_P14	1.35018	0.00007	1.35018	1818.654	2468.836
A_32_P20	BU561469	Transcribe	BU561469	1.35017	0.00023	1.35017	1681.574	2255.146
A_23_P25	NM_00074	cholinergic	CHRNA2	1.34993	0.00047	1.34993	616.2267	827.1737
A_23_P15	NM_01541	ribosomal	RPL36,DK	1.34972	4.17E-12	1.34972	46502.84	62729.13
A_23_P15	NM_00201	Homo sapi	FOXO1A,F	1.34957	0.00027	1.34957	1066.298	1426.673
A_23_P76	NM_00724	Homo sapi	PRR4,LPR	1.34953	0.00037	1.34953	4123.305	5559.401
A_24_P30	NM_02115	protein kina	TRIB3,NIP	1.34942	0.0003	1.34942	981.9168	1340.881
A_24_P88	NM_00596	Homo sapi	MUC6	1.34933	3.52E-06	1.34933	11955.03	16018.51
A_32_P12	NM_01416	HSPC128	HSPC128,	1.34933	0.00014	1.34933	3683.662	5012.602
A_23_P13	NM_00626	Homo sapi	PROP1	1.34906	0.00032	1.34906	24315.29	32652.66
A_24_P93	AY358802	Homo sapi	VLGN1945	1.3488	2.98E-06	1.3488	2120.909	2840.388
A_32_P11	AF195420	Odz, odd C	ODZ3	1.34877	0.0003	1.34877	631.6459	852.3936
A_24_P27	NM_01238	p8 protein	P8,COM1	1.3486	0.00056	1.3486	2520.993	3411.175
A_32_P148058		Sequence	THC19810	1.34852	0.00005	1.34852	671.3044	905.0497
A_24_P11	NM_20749	Homo sapi	FLJ40092	1.34805	0.00014	1.34805	1419.885	1897.714
A_23_P46	NM_01607	Homo sapi	BOLA1,CG	1.34761	0.00032	1.34761	5424.648	7365.204
A_24_P13	NM_00210	glycophorin	GYPC,GE,	1.34758	4.42E-08	1.34758	1342.382	1807.729
A_23_P34	NM_08086	SPRY dom	SPSB3,SS	1.34708	8.72E-07	1.34708	4087.534	5509.766
A_24_P91	M14755	Homo sapi	ABL1	1.34698	0.00009	1.34698	15034.96	20192.15
A_23_P40	NM_00150	Homo sapi	GNRH2,Gr	1.34692	2.44E-06	1.34692	6642.178	8941.217
A_23_P40	NM_00113	Homo sapi	AFG3L1,AI	1.34681	3.62E-06	1.34681	1649.375	2218.104
A_24_P10	BC022822	Immunoglo	IGLC2	1.34674	0.00005	1.34674	25770.96	34860.11
A_24_P52168		Unknown	A_24_P52	1.34662	3.01E-06	1.34662	3210.262	4320.898
A_24_P127442		Unknown	A_24_P12	1.34648	5.72E-06	1.34648	116681.1	157845.3
A_24_P63	BC008632	Chromosor	C6orf176	1.34591	0.00019	1.34591	2773.844	3751.405

A_32_P20294		Unknown	A_32_P20	1.34564	0.00016	1.34564	12584.58	16947.5
A_24_P25	NM_00608	tubulin, bet	TUBB3,MC	1.34555	1.96E-12	1.34555	23697.16	31893.01
A_24_P72	AK123446	CDNA FLJ	AK123446	1.34494	0.00001	1.34494	6858.69	9150.898
A_24_P10	AK026491	Homo sapi	RPL12	1.34443	1.05E-10	1.34443	54348.23	73030.98
A_24_P11	NM_01451	Homo sapi	KIR3DL1	1.34427	1.22E-08	1.34427	1861.953	2507.178
A_23_P24	NM_17358	Homo sapi	RCOR2	1.34424	0.00003	1.34424	738.4552	992.8643
A_23_P35	NM_03208	Homo sapi	PCDHGA8	1.34398	6.93E-06	1.34398	3631.831	4874.667
A_32_P58	AF070632	Clone 244	AF070632	1.34374	6.44E-07	1.34374	6284.105	8438.819
A_23_P67	NM_01796	hypothetica	FLJ20850,	1.34371	9.85E-07	1.34371	5508.634	7423.456
A_23_P16	NM_00460	syntaxin 1	STX1A,ST	1.34362	0.00001	1.34362	2547.62	3424.877
A_24_P6903		Unknown	ENST0000	1.34359	0.00005	1.34359	10524.94	14136.24
A_24_P26	BC020884	Homo sapi	HIST4H4	1.34302	0.00009	1.34302	694.7687	931.0059
A_23_P61	NM_00659	Homo sapi	SLC12A7,	1.34297	1.86E-07	1.34297	12525.16	16888.29
A_32_P14	BC053367	Opposite s	GATS	1.34289	0.00002	1.34289	918.3658	1243.157
A_23_P28	NM_00208	glutathione	GPX4,MC	1.34286	8.62E-07	1.34286	5916.693	7931.664
A_24_P30	NM_02069	Transmem	TMCC3	1.34252	1.02E-10	1.34252	3174.807	4263.525
A_23_P10	NM_08067	Homo sapi	SPAG4L,T	1.34245	0.00075	1.34245	1183.619	1574.353
A_23_P32	NM_01337	Homo sapi	DPP7,QPF	1.34233	8.90E-10	1.34233	2827.899	3795.581
A_24_P54	NM_15240	hypothetica	FLJ39370	1.34191	0.00005	1.34191	2653.133	3564.199
A_23_P20	NM_18301	Homo sapi	CREM,ICE	1.34189	0.00055	1.34189	1987.816	2672.99
A_23_P20	NM_01231	Homo sapi	KIN,BTCD,	1.3418	0.00031	1.3418	821.3348	1107.562
A_23_P24	NM_01693	Homo sapi	EFEMP2,U	1.34178	7.56E-06	1.34178	1049.89	1404.444
A_32_P217853		Unknown	THC18711	1.3416	0.00007	1.3416	949.9617	1264.717
A_23_P15	NM_00071	Calcium ch	CACNA1B	1.34146	2.14E-06	1.34146	6613.021	8864.537
A_24_P31	XM_37298	Homo sapi	ENST0000	1.34116	1.94E-06	1.34116	1390.251	1872.743
A_24_P35	NM_00225	Killer cell ir	KIR2DL4	1.34112	0.00002	1.34112	1749.965	2329.125
A_23_P66	XM_37094	Homo sapi	LOC38822	1.34089	1.43E-07	1.34089	2114.31	2827.36
A_23_P96	NM_01615	LW-1	LW-1	1.34076	9.53E-09	1.34076	1539.689	2065.771
A_23_P21	NM_00263	Homo sapi	PI3,ESI,W	1.34027	0.00004	1.34027	7154.012	9600.555
A_24_P33	NM_01664	HDCMA18	HDCMA18	1.34017	3.34E-06	1.34017	3055.077	4108.7
A_32_P23	NR_00143	Homo sapi	HLA-H	1.34015	2.10E-06	1.34015	2981.905	3988.209
A_24_P57	NM_03260	CREB/ATF	CREB3L3,	1.34003	0.00007	1.34003	1751.593	2335.559
A_24_P340891		Unknown	A_24_P34	1.33982	8.07E-07	1.33982	33639.35	45245.53
A_24_P37	AK090421	RGD, leuci	RLTPR	1.33979	0.00008	1.33979	14187	19037.23
A_32_P33	BC007632	RIM bindin	KIAA0318	1.33959	3.89E-09	1.33959	2906.866	3890.946
A_24_P161173		Unknown	A_24_P16	1.33932	3.83E-07	1.33932	3687.968	4939.196
A_32_P87582		Unknown	THC18127	1.33931	0.00003	1.33931	2626.941	3489.732
A_24_P51	AL834347	Hydroxy-de	HSD3B7	1.33921	2.17E-06	1.33921	1503.825	2019.954
A_23_P56	BC019266	Dystrophia	DMWD	1.33919	0.00016	1.33919	31021.66	41629.09
A_24_P27	NM_05287	hypothetica	MGC4677,	1.33901	3.28E-06	1.33901	7668.768	10333.1
A_23_P15	NM_00419	acetylserot	ASMTL,AS	1.33896	0.00056	1.33896	773.3607	1047.858
A_24_P21	NM_01902	hypothetica	FLJ20010	1.33835	0.00001	1.33835	4757.233	6373.372
A_24_P117803		BA161I19.	ENST0000	1.33829	2.66E-07	1.33829	2828.909	3787.571
A_24_P17	AK092134	Hypothetic	AK092134	1.33824	6.90E-06	1.33824	3846.331	5139.929
A_32_P15	BC020336	TAO kinas	RPS2	1.3381	0.00001	1.3381	1423.783	1900.812
A_23_P78	NM_01645	protein kin	PRKD2,PK	1.33793	5.49E-11	1.33793	3110.9	4160.813
A_23_P25	NM_00259	procollager	PCOLCE,F	1.33778	2.08E-09	1.33778	2415.429	3234.191
A_24_P36	NM_03288	KIAA1870	COL27A1,	1.33764	0.00024	1.33764	8158.883	10915.16
A_23_P42	NM_03189	Homo sapi	CACNG8	1.33751	0.00094	1.33751	1133.516	1542.907
A_23_P35	NM_18159	Homo sapi	UPP1,UP,	1.33743	1.20E-09	1.33743	2043.099	2729.377
A_23_P10	AB007914	Ciliary root	CROCC	1.33741	0.00005	1.33741	930.1954	1250.183
A_24_P13	NM_00102	ribosomal	RPS19,DB	1.33687	3.75E-06	1.33687	155738.3	208783.5

A_24_P36	NM_19807	golgi autoa	FLJ40113,	1.33682	0.0007	1.33682	955.7565	1289.509
A_24_P255874		Unknown	ENST0000	1.33638	2.63E-10	1.33638	7537.361	10067
A_23_P35	NM_00558	Homo sapi	MEF2A,AD	1.33635	2.18E-07	1.33635	1348.291	1799.241
A_24_P910050		ALU7_HUM	THC19460	1.33628	0.00012	1.33628	3181.741	4242.423
A_23_P50	AK075226	Zinc finger	ZNF331	1.33573	0.00003	1.33573	1654.491	2213.298
A_23_P11	NM_19916	CXYorf1-re	FLJ25222	1.33535	1.28E-11	1.33535	6902.198	9219.264
A_23_P16	NM_02410	hypothetica	MLPH,In,M	1.33534	0.00073	1.33534	5829.655	7836.529
A_23_P83	NM_01540	Homo sapi	DFNB31,W	1.335	2.62E-06	1.335	1630.491	2185.765
A_23_P40	NM_00218	Homo sapi	IL7R,CD12	1.33467	0.0002	1.33467	1112.461	1472.11
A_24_P938263		Unknown	A_24_P93	1.33467	0.00001	1.33467	909.5143	1206.251
A_24_P34	NM_20342	hypothetica	LOC38815	1.33448	1.45E-06	1.33448	3037.357	4043.565
A_23_P34	AB051464	Kelch-like	KLHL15	1.33412	0.00014	1.33412	968.2141	1284.629
A_23_P24	NM_00265	Homo sapi	PLAU,ATF	1.33409	0.00074	1.33409	2676.265	3554.778
A_23_P15	NM_03057	START do	STARD5	1.33407	5.40E-10	1.33407	1913.527	2554.986
A_24_P18	NM_02085	Homo sapi	VPS18,KIA	1.33385	7.01E-06	1.33385	7055.496	9400.296
A_23_P38	NM_14499	Homo sapi	LRRC45,M	1.33383	0.00003	1.33383	1678.22	2246.758
A_32_P21794		Unknown	A_32_P21	1.33378	4.95E-07	1.33378	3326.678	4434.397
A_23_P83	AF086417	X-prolyl am	XPNPEP2	1.3335	4.37E-06	1.3335	2450.177	3255.375
A_24_P18	AB016902	Myeloid/lyn	MLLT4	1.33302	3.25E-07	1.33302	4860.129	6462.823
A_24_P84	AK055809	Hypothetic	LOC14870	1.33285	2.59E-11	1.33285	6412.301	8546.07
A_24_P642328		Unknown	A_24_P64	1.3325	3.31E-07	1.3325	2511.975	3346.925
A_24_P896373		Unknown	THC18044	1.33207	5.11E-06	1.33207	3132.729	4169.911
A_23_P15	NM_03276	Homo sapi	NM_03276	1.33205	0.00004	1.33205	2462.798	3276.783
A_24_P17	M13930	V-myc mye	MYC	1.33144	0.0002	1.33144	6468.776	8513.246
A_32_P2177		Unknown	THC19021	1.33118	1.11E-06	1.33118	1423.616	1898.693
A_24_P96	NM_18148	Homo sapi	ZGPAT,ZC	1.33083	5.34E-07	1.33083	1811.163	2413.333
A_24_P18	NM_01584	methyl-Cp	MBD1,RFT	1.33074	1.89E-06	1.33074	1218.648	1617.024
A_23_P55	NM_00166	ADP-ribosy	ARF4L,AR	1.33062	5.51E-08	1.33062	3037.62	4044.857
A_24_P69	XM_37170	hypothetica	LOC38922	1.33051	0.00003	1.33051	1017.823	1360.657
A_23_P20	NM_01661	Homo sapi	KCNK4,TR	1.33045	2.19E-06	1.33045	4030.856	5368.78
A_24_P921980		Unknown	A_24_P92	1.33017	0.00001	1.33017	6118.732	8106.514
A_32_P17	BC020462	Ribosomal	RPS9	1.33005	0.00039	1.33005	3724.454	4979.596
A_23_P31	NM_17384	Homo sapi	SPACA3,L	1.33	1.94E-07	1.33	2176.499	2890.661
A_23_P26	NM_01442	copine VII	CPNE7,MC	1.32995	0.00001	1.32995	4001.842	5330.792
A_23_P30	NM_00551	Major histo	HLA-E	1.32968	0.00003	1.32968	1606.787	2137.531
A_24_P17	AJ422142	Unc-93 hor	UNC93B1	1.32922	0.00001	1.32922	3761.697	4984.912
A_24_P34	NM_00541	sine oculis	SIX3,HPE2	1.32903	0.00094	1.32903	916.273	1200.755
A_32_P12	NM_00688	ATP synth	ATP5E,AT	1.32852	2.83E-06	1.32852	13859.79	18480
A_23_P61	NM_05819	Homo sapi	C21orf70,F	1.32844	3.55E-08	1.32844	3899.564	5188.137
A_24_P51	BC031250	Hypothetic	BC031250	1.32841	0.00007	1.32841	1777.177	2352.394
A_24_P118382		Unknown	A_24_P11	1.32808	0.00015	1.32808	48960.06	65526.66
A_32_P14	NM_18252	Homo sapi	C22orf15,N	1.32789	1.91E-07	1.32789	3146.498	4174.566
A_24_P16	U28759	Major histo	HLA-B	1.32781	0.00042	1.32781	3326.684	4397.044
A_23_P93	NM_00667	solute carri	SLC22A7,N	1.32768	0.00018	1.32768	780.8067	1039.375
A_24_P393087		Unknown	ENST0000	1.32743	2.70E-06	1.32743	5393.903	7133.746
A_23_P10	NM_03260	Homo sapi	CREB3L3,	1.32731	0.00004	1.32731	903.5172	1205.79
A_24_P93	AK074614	Insulin-like	IGF2	1.32698	0.00002	1.32698	7611.776	10051.15
A_24_P91	M74509	Human en	M74509	1.32695	0.00084	1.32695	19317.12	25722.5
A_23_P33	NM_14896	Homo sapi	OXER1,GF	1.32691	8.27E-09	1.32691	6997.009	9276.536
A_23_P42	NM_02196	Homo sapi	TEAD1,TC	1.32663	0.00001	1.32663	1111.982	1479.916
A_24_P23	NM_02018	myelin prot	C14orf162,	1.32637	2.88E-07	1.32637	5309.664	7031.369
A_24_P230074		Unknown	ENST0000	1.32609	0.00011	1.32609	927.2213	1222.473

A_23_P41	NM_00482	Homo sapi	NCR2,LY9	1.32606	7.93E-07	1.32606	1607.048	2123.032
A_32_P96970		Unknown	A_32_P96	1.326	0.00025	1.326	4464.85	5890.156
A_23_P75	NM_01327	Homo sapi	C11orf9,KI	1.3259	9.79E-10	1.3259	8841.076	11718.5
A_23_P31	NM_01234	Homo sapi	NTSR2,NT	1.32585	0.00026	1.32585	721.5034	958.3872
A_23_P12	NM_08061	Homo sapi	WFDC3,W	1.32583	6.45E-07	1.32583	4200.526	5571.421
A_24_P477610		Unknown	A_24_P47	1.32546	3.94E-09	1.32546	4267.174	5661.951
A_24_P73	AI939539	Transcribe	AI939539	1.32539	0.00075	1.32539	3429.887	4520.43
A_24_P418712		Unknown	A_24_P41	1.32532	4.98E-09	1.32532	78993.22	104872.1
A_32_P18	NM_00608	tubulin, bet	TUBB2	1.32524	0.00089	1.32524	203977.7	274173.6
A_24_P101439		olfactory re	NP186050	1.32502	8.04E-07	1.32502	1606.179	2123.583
A_23_P11	NM_03284	Homo sapi	FIBCD1,FL	1.32495	3.45E-06	1.32495	1193.063	1576.432
A_23_P33	NM_17214	Homo sapi	IL29,IFNL1	1.32449	0.00024	1.32449	1087.115	1453.698
A_23_P31	NM_18324	hypothetica	LOC28625	1.3244	1.05E-06	1.3244	3655.916	4848.705
A_24_P90	NM_02492	esterase 3	FLJ21736,	1.32427	0.0007	1.32427	655.0729	856.2909
A_24_P25	NM_02172	nuclear rec	NR1D1,EA	1.324	3.45E-06	1.324	1469.956	1938.319
A_32_P66684		Unknown	THC19704	1.32394	7.59E-06	1.32394	1581.159	2101.236
A_24_P36	NM_01617	hypothetica	LOC51149	1.32377	1.14E-09	1.32377	3230.233	4273.263
A_32_P11	NM_13844	Homo sapi	C6orf150	1.32333	0.00009	1.32333	1299.351	1706.405
A_23_P37	NM_00071	calcium ch	CACNA1C	1.32322	0.00026	1.32322	12168.43	16047.01
A_24_P773012		CO1A_HUI	THC18706	1.32307	1.06E-06	1.32307	3357.556	4443.953
A_23_P37	NM_15322	Homo sapi	CILP2,CLIF	1.32304	3.97E-09	1.32304	2465.813	3265.695
A_24_P39	AW302767	xr55h07.x1	AW302767	1.323	0.00024	1.323	1384.912	1817.345
A_23_P11	NM_01755	Homo sapi	KIAA1193	1.32282	0.00004	1.32282	8824.152	11674.36
A_24_P152753		Unknown	A_24_P15	1.32278	0.00007	1.32278	68843.21	91683.91
A_23_P16	NM_02182	hypothetica	PPCDC,MI	1.32269	0.00004	1.32269	1198.235	1580.323
A_24_P28	NM_01665	heptacellul	DACT1,DP	1.32267	0.00068	1.32267	775.0388	1034.689
A_24_P27	NM_20585	PNPK6288	LOC38985	1.32241	0.0001	1.32241	1263.742	1688.752
A_24_P91	AV647560	Transcribe	AV647560	1.32211	0.00011	1.32211	8411.093	11010.86
A_24_P83	NM_00084	glutamate r	GRM5,mG	1.32198	1.62E-08	1.32198	1731.977	2288.658
A_24_P24	AK094167	UDP-gluco	UGP2	1.32194	2.58E-06	1.32194	6969.294	9179.264
A_23_P13	NM_02487	Homo sapi	DOK3,FLJ	1.32148	0.00063	1.32148	1024.476	1336.929
A_32_P16848		RNP2_HUI	THC19660	1.32134	3.28E-07	1.32134	4179.317	5529.041
A_24_P37	NM_00455	nuclease s	YBX1,YB1	1.32132	2.33E-06	1.32132	11234.18	14766.43
A_24_P18	AF099011	Homo sapi	EHD1,PAS	1.32063	0.00003	1.32063	1145.387	1503.71
A_24_P57	NM_15332	Homo sapi	ALDH16A1	1.32057	8.38E-07	1.32057	1300.33	1726.21
A_32_P64	NM_15260	Chromosor	FLJ35382	1.32011	6.41E-06	1.32011	1089.275	1437.551
A_24_P18	NM_02034	ankyrin rep	ANKRD2,A	1.31986	3.60E-07	1.31986	20581.64	27233.1
A_23_P97	NM_00122	Homo sapi	CASP9,MC	1.31959	4.84E-07	1.31959	2551.798	3367.165
A_23_P40	NM_00211	Major histo	HLA-A	1.31918	7.23E-06	1.31918	8228.578	10814.03
A_23_P13	NM_00537	Homo sapi	MYCL2	1.31883	5.64E-07	1.31883	6085.309	8005.112
A_24_P39	AW302767	xr55h07.x1	AW302767	1.31862	0.00002	1.31862	68422.3	90629.95
A_24_P34	NM_00230	lectin, gala	LGALS7,G	1.3185	0.00005	1.3185	856.6144	1130.021
A_23_P18	NM_15262	hypothetica	FLJ35954	1.31813	8.51E-09	1.31813	1975.527	2605.128
A_23_P12	BX281566	Hypothetic	LOC44095	1.31797	0.00005	1.31797	1120.355	1468.313
A_24_P92	AY358167	AT rich inte	ARID5A	1.31777	6.95E-07	1.31777	1463.604	1924.795
A_24_P23	AF177925	Mucin 4, tr	MUC4	1.31768	5.48E-06	1.31768	19393.96	25545.59
A_23_P46	NM_02486	Hypothetic	FLJ14050	1.3175	0.0009	1.3175	697.4402	915.6775
A_23_P78	NM_01770	Homo sapi	FBXL12,Fb	1.31738	0.00003	1.31738	1603.401	2102.978
A_24_P51	NM_02011	hypothetica	LARS,LEU	1.31735	4.16E-11	1.31735	7263.276	9568.314
A_24_P340886		Unknown	A_24_P34	1.31732	3.16E-06	1.31732	42499.23	56211.7
A_24_P82	NM_00735	heat shock	HSPCB,HS	1.31711	0.00066	1.31711	4125.883	5473.411
A_23_P20	NM_00008	collagen, ty	COL1A1,O	1.31702	0.00011	1.31702	918.3738	1214.994

A_24_P34	NM_00313	Serum res	SRF	1.31674	4.62E-06	1.31674	7558.726	9966.15
A_32_P19	NM_00555	Homo sapi	LAMA1,LA	1.31644	1.99E-06	1.31644	2572.459	3369.886
A_32_P20	AF085962	Full length	AF085962	1.3162	3.42E-06	1.3162	2875.406	3772.747
A_32_P92	AF346307	Chromosor	F379	1.31613	1.47E-09	1.31613	4220.605	5553.444
A_23_P92	NM_00246	myeloid dif	MYD88	1.3157	0.0002	1.3157	818.2907	1082.123
A_24_P24	NM_14575	Homo sapi	PHLDB2,LI	1.31545	0.00074	1.31545	1864.002	2436.523
A_24_P28	BC008094	Retinal G p	RGR	1.31536	5.10E-06	1.31536	2415.813	3168.722
A_32_P86	XM_37338	Homo sapi	XM_37338	1.31483	0.00002	1.31483	2188.684	2869.224
A_23_P35	NM_17855	hypothetica	MGC44505	1.31464	0.00001	1.31464	976.3333	1285.504
A_23_P41	NM_15240	Homo sapi	GRPEL2,F	1.31462	0.0005	1.31462	3723.005	4920.902
A_24_P81	AK097447	FLJ44082 p	FLJ44082	1.31429	1.93E-08	1.31429	3217.661	4226.033
A_23_P30	NM_00605	Homo sapi	TNIP1,VAN	1.31404	4.15E-08	1.31404	3367.464	4425.628
A_24_P93	BC033528	Homo sapi	BC033528	1.31395	0.00061	1.31395	13390.71	17603.16
A_24_P518908		Unknown	A_24_P51	1.31387	0.0003	1.31387	904.1757	1181.502
A_23_P31	NM_01895	major histo	HLA-F,HLA	1.31359	2.74E-09	1.31359	9107.779	11959.71
A_23_P72	NM_17803	Homo sapi	CYP4X1,M	1.31355	2.47E-08	1.31355	3661.373	4803.901
A_24_P35	XM_37331	Homo sapi	XM_37331	1.31352	6.41E-07	1.31352	13999.73	18429.55
A_23_P11	AK125431	Hypothetica	DJ159A19.	1.3129	3.40E-08	1.3129	3632.778	4760.646
A_23_P27	NM_00162	Homo sapi	ALOX5AP,	1.31282	0.00007	1.31282	2057.578	2698.703
A_23_P38	NM_01233	histone ace	MYST4,qkl	1.31223	7.68E-07	1.31223	1370.978	1798.145
A_23_P31	NM_00409	Homo sapi	EMX1	1.31216	0.00003	1.31216	13118.88	17249.82
A_23_P44	NM_13896	Homo sapi	MSI2,MSI2	1.31209	0.00003	1.31209	1582.894	2072.341
A_32_P20	AL137476	Hypothetica	LOC14983	1.3118	7.81E-07	1.3118	2800.814	3663.035
A_32_P90987		G3P2_HUM	THC19647	1.31178	7.16E-08	1.31178	8074.985	10587.21
A_24_P341606		Seven tran	ENST0000	1.31175	3.45E-10	1.31175	7117.319	9330.733
A_32_P223488		Unknown	THC18173	1.31174	8.75E-07	1.31174	5505.673	7211.753
A_24_P46	NM_01647	scotin	SCOTIN	1.31126	7.68E-07	1.31126	6959.001	9113.86
A_24_P290046		Unknown	A_24_P29	1.31111	0.00061	1.31111	940.3283	1244.367
A_23_P36	NM_02291	vasculin	GPBP1,GF	1.31071	0.00027	1.31071	5287.607	6973.707
A_32_P23960		Unknown	A_32_P23	1.31069	7.25E-09	1.31069	10191.9	13343.12
A_23_P40	NM_17587	hypothetica	ANKRD43	1.31056	0.00018	1.31056	5445.726	7097.071
A_24_P91	AF287958	Major histo	HLA-A	1.31054	3.08E-06	1.31054	6398.373	8367.738
A_23_P31	NM_17853	Homo sapi	FLJ38379	1.3104	1.64E-07	1.3104	1888.866	2472.793
A_23_P14	NM_02020	Homo sapi	SHD	1.31034	0.00005	1.31034	4764.751	6239.152
A_23_P93	NM_13845	Homo sapi	C6orf68,M	1.31027	2.75E-08	1.31027	3122.202	4105.889
A_23_P97	BX641020	AT rich inte	ARID5B	1.31026	0.00063	1.31026	1015.312	1314.373
A_24_P306527		Unknown	A_24_P30	1.30987	0.00004	1.30987	68720.01	90505.8
A_23_P10	NM_00261	Homo sapi	SERPINF1	1.30971	0.00002	1.30971	938.4539	1229.482
A_23_P59	NM_01393	Homo sapi	OR11A1,O	1.3097	0.00002	1.3097	6081.18	7916.235
A_23_P12	NM_01838	hypothetica	FLJ11292	1.30969	3.24E-06	1.30969	1488.046	1947.65
A_32_P199244		Unknown	A_32_P19	1.30927	5.20E-10	1.30927	3556.867	4656.526
A_23_P78	NM_00063	Homo sapi	VTN,V75,V	1.30917	0.00008	1.30917	1531.13	1995.477
A_24_P324224		Unknown	A_24_P32	1.30902	1.60E-06	1.30902	93296.02	122573.7
A_23_P61	AK123975	Hypothetica	MGC27165	1.30899	0.00007	1.30899	1002.047	1308.779
A_23_P56	NM_00215	Homo sapi	HSPE1,CP	1.30889	0.00002	1.30889	263453.7	346774
A_24_P27	NM_00017	glycoprotei	GP1BA,BS	1.30875	0.00016	1.30875	1931.346	2513.834
A_23_P38	NM_13349	Homo sapi	ASAH3,AC	1.3085	0.00014	1.3085	1863.359	2395.843
A_32_P168727		Unknown	A_32_P16	1.30849	7.20E-06	1.30849	2078.949	2710.691
A_24_P135461		Unknown	ENST0000	1.30847	0.00012	1.30847	16801.78	21655.59
A_24_P49	NM_03145	selenoprot	SELO	1.30821	0.00005	1.30821	1140.982	1499.151
A_24_P926484		Unknown	A_24_P92	1.30802	1.79E-07	1.30802	6393.167	8331.096
A_32_P52	NM_17573	hypothetica	LOC20124	1.30799	0.00003	1.30799	1478.434	1939.61

A_32_P121755	Unknown	THC19145	1.30774	0.00013	1.30774	3768.147	4922.674
A_24_P92	AF230200	Zinc finger	ZNF24	1.30771	0.00012	1.30771	2016.219	2607.287
A_23_P49	NM_13917	testis nucle	LOC16193	1.30768	0.00012	1.30768	1695.245	2234.666
A_24_P94	NM_15260	hypothetica	C1orf177,F	1.30763	1.78E-06	1.30763	2051.773	2690.153
A_24_P15	NM_02065	junctophilin	JPH3,JP3,	1.30755	3.09E-06	1.30755	2107.763	2760.765
A_23_P33	NM_00163	alkaline ph	ALPI,IAP	1.30727	0.00002	1.30727	1109.792	1461.648
A_24_P82	NM_03237	hypothetica	EEF1D,EF	1.3071	0.00002	1.3071	70220.48	91310.59
A_24_P306994	Unknown	A_24_P30	1.30705	0.00098	1.30705	2877.528	3791.623
A_24_P606663	Unknown	A_24_P60	1.30698	0.00004	1.30698	75872.61	99818
A_32_P141938	Unknown	THC18959	1.30681	0.00002	1.30681	1221.554	1592.972
A_23_P50	NM_17822	Killer cell ir	KIR2DS4	1.30678	0.00002	1.30678	1563.457	2034.722
A_23_P48	NM_08074	Homo sapi	RNF36,Trif	1.30659	0.00019	1.30659	9352.338	12184.93
A_23_P14	NM_00277	Homo sapi	PRTN3,ME	1.30651	0.00031	1.30651	935.3694	1213.165
A_23_P21293	Unknown	A_23_P21	1.30642	0.00087	1.30642	1081.476	1427.924
A_32_P42	U12206	Human clo	unknown	1.30613	0.0009	1.30613	1114.67	1473.243
A_32_P97	BC043279	Homo sapi	BC043279	1.30587	0.00059	1.30587	702.8508	915.8319
A_23_P49	NM_00533	Homo sapi	HBQ1	1.30571	0.00007	1.30571	3751.698	4885.924
A_23_P39	NM_02500	Thrombosp	TSRC1	1.30556	0.00031	1.30556	779.5845	1013.393
A_24_P29	NM_00463	sema dom	SEMA3B,S	1.30539	8.21E-08	1.30539	3185.626	4160.558
A_24_P540555	Unknown	A_24_P54	1.30539	6.42E-09	1.30539	2667.542	3489.41
A_24_P47	BM720284	A kinase (F	AKAP10	1.30498	0.00047	1.30498	17369.54	22701.72
A_24_P511776	Unknown	A_24_P51	1.30484	0.00025	1.30484	1073.035	1411.677
A_32_P14	NM_00698	nuclear por	NPIP	1.30472	0.00023	1.30472	29500.54	38651.61
A_32_P17	D86974	KIAA0220-	KIAA0220	1.30456	0.00049	1.30456	38248.46	50040.17
A_32_P514599	CH60_HUM	THC19561	1.30441	0.00065	1.30441	19028.88	24607.04
A_23_P66	NM_01330	Homo sapi	ZDHHC1,Z	1.30426	0.00027	1.30426	966.1907	1250.863
A_24_P31	XM_29204	similar to ri	LOC34141	1.30389	1.17E-09	1.30389	115131.3	150203.1
A_24_P41	NM_00671	mannosida	MAN2C1,M	1.30361	0.00038	1.30361	836.4316	1084.691
A_23_P125109	Unknown	A_23_P12	1.30325	1.71E-06	1.30325	12063.04	15666.74
A_23_P12	NM_00364	Homo sapi	DGKZ,DAC	1.30271	2.08E-09	1.30271	2595.848	3381.385
A_23_P50	NM_15262	Homo sapi	ZNF92,TF1	1.30266	0.00042	1.30266	1470.836	1910.92
A_24_P20	NM_02407	hypothetica	TMEM38A	1.30255	5.55E-06	1.30255	8610.23	11116.94
A_23_P27	NM_19924	multidrug r	MGC1317	1.30253	2.17E-09	1.30253	9869.508	12849.74
A_24_P93	NM_00062	Nitric oxide	NOS1	1.30242	2.01E-09	1.30242	3182.123	4138.612
A_24_P29	M28207	Major histo	HLA-C	1.30197	0.00006	1.30197	35070.88	45925.45
A_23_P12	NM_02121	MUM2 prot	TRAPPC1,	1.3015	9.29E-07	1.3015	4085.311	5308.153
A_23_P20	NM_01209	Homo sapi	ANGPTL2,	1.30118	0.00001	1.30118	1499.813	1946.091
A_32_P20	AJ291676	Myotubular	MTMR9	1.301	0.00056	1.301	16396.41	21337.53
A_24_P91	XM_37661	Homo sapi	XM_37661	1.30081	0.00021	1.30081	5486.883	7120.942
A_23_P93	NM_01754	Homo sapi	EPDR1,UC	1.30077	1.70E-06	1.30077	1556.138	2027.904
A_24_P48	BX647217	Heterogen	HNRPA3	1.30076	1.82E-06	1.30076	2107.948	2741.132
A_23_P16	NM_00306	Homo sapi	SNAI2,SLU	1.30044	0.00073	1.30044	3083.46	4027.901
A_23_P66	AF090930	Forkhead b	FOXK2	1.30034	0.00018	1.30034	6286.908	8140.206
A_23_P86	NM_00732	Homo sapi	DMBT1,GF	1.30031	0.00003	1.30031	1698.774	2205.129
A_24_P23	AF289554	PERQ ami	PERQ1	1.30015	0.00082	1.30015	1929.737	2469.467
A_23_P16	NM_00230	Homo sapi	LGALS1,G	1.30007	1.44E-06	1.30007	30198.41	39313.2
A_23_P31	NM_03237	hypothetica	EEF1D,EF	1.29986	0.00019	1.29986	45540.2	58794.62
A_23_P34	NM_00714	Polycomb (	PCGF2	1.29976	0.00009	1.29976	3190.685	4129.638
A_24_P13	NM_00674	protein phc	PPP1R1A	1.29947	0.00008	1.29947	913.8173	1189.19
A_24_P83	NM_15345	Homo sapi	C21orf86	1.29944	0.00078	1.29944	1386.557	1816.959
A_23_P15	NM_00376	Homo sapi	STX11	1.29936	0.00003	1.29936	1281.233	1668.792
A_23_P41	NM_02194	Homo sapi	SMAP1,SM	1.29922	9.29E-07	1.29922	1775.154	2293.96

A_23_P39	AK021860	Homo sapi	FBXO17,F	1.29921	0.00002	1.29921	11236.39	14552.61
A_24_P74	NM_18261	G protein-c	GPR144	1.29919	8.53E-10	1.29919	4655.867	6050.141
A_24_P64	XM_37755	similar to e	LOC40193	1.29905	4.07E-10	1.29905	10412.6	13523.52
A_23_P16	NM_00574	Homo sapi	TOB1,TOB	1.29854	5.12E-08	1.29854	4325.49	5609.633
A_23_P14	AK097070	Transmem	TM6SF2	1.29841	0.00011	1.29841	1021.685	1333.555
A_23_P13	NM_00067	Homo sapi	ADRA1D,D	1.29804	0.00045	1.29804	13654.04	17651.31
A_24_P60	NM_00066	acetylcholi	ACHE,YT	1.29791	1.51E-06	1.29791	1971.495	2555.154
A_24_P76	NM_01230	f-box and l	FBXL7,FBI	1.29786	0.00006	1.29786	830.6896	1077.136
A_24_P30	NM_00710	ATP synth	ATP5I,ATF	1.29778	0.00004	1.29778	25773.51	33368.45
A_23_P51	NM_00156	Homo sapi	TNFRSF9,	1.29775	0.00042	1.29775	1208.666	1560.849
A_23_P35	AK090500	M4-50 mRI	AK090500	1.29768	8.56E-06	1.29768	18479.5	23994.57
A_32_P65	NM_03237	hypothetic	EEF1D,EF	1.29762	0.0003	1.29762	3165.803	4092.755
A_23_P20	NM_00371	Homo sapi	NPFF,FMR	1.2976	0.00001	1.2976	2366.835	3079.332
A_32_P16	BM981501	Similar to 6	BM981501	1.29735	0.00035	1.29735	5544.809	7165.362
A_23_P30	NM_00008	cholinergic	CHRNE,AC	1.29732	0.00008	1.29732	1735.027	2271.618
A_24_P55	NM_15282	Homo sapi	TES,TESS	1.29726	6.48E-08	1.29726	2169.283	2815.034
A_23_P36	NM_13870	beta-1,3-N	B3Gn-T6,M	1.29721	1.81E-06	1.29721	4197.343	5427.244
A_23_P34	BC009979	G-protein s	GPSM1	1.29707	0.00031	1.29707	1034.826	1351.801
A_24_P74	BC047425	Hypothetic	LOC34802	1.29693	5.66E-06	1.29693	1257.847	1628.812
A_23_P35	NM_14575	Homo sapi	KIFC2	1.29689	3.32E-08	1.29689	2145.282	2781.6
A_23_P25	NM_00050	Homo sapi	CYP21A2,	1.29689	0.00003	1.29689	1132.824	1466.402
A_24_P16	BX640625	Hypothetic	MGC27165	1.29669	3.96E-06	1.29669	1284.676	1664.654
A_24_P28	AK057161	MADS box	MEF2B	1.29665	0.00002	1.29665	22074.91	28558.34
A_23_P336854		DJ999L4.1	RPL12L3	1.29651	1.52E-06	1.29651	111164.1	144720.8
A_24_P11	NM_15347	Homo sapi	CSAG1,CS	1.2964	0.00043	1.2964	2763.634	3567.462
A_32_P211603		Unknown	THC18222	1.29564	0.00003	1.29564	1772.551	2285.542
A_23_P31	AF090884	Unc-13 hor	UNC13B	1.29561	5.08E-08	1.29561	3462.845	4482.747
A_23_P75	NM_02230	Homo sapi	HRH2,H2R	1.29546	6.39E-06	1.29546	4301.181	5560.352
A_24_P31	NM_00133	cardiotroph	CTF1,CT1,	1.29542	0.00021	1.29542	2301.09	2971.854
A_23_P16	NM_02160	Homo sapi	FXYD2,HC	1.29539	0.00007	1.29539	2043.715	2637.979
A_23_P43	M76744	Carcinoem	CEACAM1	1.29529	6.28E-06	1.29529	1785.355	2308.588
A_23_P161968		Unknown	ENST0000	1.29527	0.00047	1.29527	3202.122	4132.547
A_24_P13	NM_19851	Homo sapi	ITIH5L,UN	1.29523	0.00021	1.29523	1233.88	1574.584
A_32_P224992		Unknown	A_32_P22	1.29494	0.00084	1.29494	1214.069	1559.93
A_24_P698698		Unknown	A_24_P69	1.29477	0.00013	1.29477	1531.518	1992.173
A_23_P33	NM_08078	Homo sapi	ACPT	1.29467	1.60E-06	1.29467	2720.533	3517.817
A_24_P94	NM_00254	oxoglutarat	OGDH,E1k	1.29455	0.00027	1.29455	12729.02	16490.67
A_23_P20	NM_00240	mitogen-ac	MAP3K3,M	1.2943	0.00003	1.2943	11150.69	14431.25
A_24_P76169		Unknown	A_24_P76	1.29405	0.00004	1.29405	19068.78	24579.73
A_24_P31	NM_13090	Homo sapi	C15orf16,C	1.29367	0.00007	1.29367	1550.179	2015.299
A_24_P11	NM_00551	Major histo	HLA-B	1.2934	6.69E-09	1.2934	11426.33	14773.28
A_24_P38	BC022830	Homo sapi	BC022830	1.29339	0.00023	1.29339	5511.85	7143.014
A_32_P20	AK024303	Homo sapi	AK024303	1.29303	9.14E-08	1.29303	2268.003	2933.71
A_24_P861442		RS15_HUM	THC19122	1.29299	0.00018	1.29299	122283.2	159057
A_23_P63	NM_00507	Homo sapi	CNTN2,AX	1.29282	9.09E-06	1.29282	1835.53	2366.833
A_23_P36	NM_14891	Abhydrolas	WBSCR21	1.29271	0.00032	1.29271	1300.477	1689.636
A_24_P100695		Lipin 3 (Lip	LPIN3	1.2927	0.00005	1.2927	912.7898	1179.066
A_32_P25	NM_19890	Homo sapi	GABRG2,C	1.29254	0.0001	1.29254	4798.124	6194.114
A_23_P13	NM_02064	Homo sapi	C11orf16	1.29245	2.32E-06	1.29245	6147.089	7930.011
A_23_P17	NM_01786	Hypothetic	FLJ20531	1.29235	0.00018	1.29235	1487.185	1928.38
A_23_P62	NM_00325	tissue inhib	TIMP1,EPA	1.29229	0.00039	1.29229	14036.2	18263.69
A_23_P50	NM_13863	Homo sapi	BCL2L12,M	1.2922	0.00004	1.2922	8700.746	11267.89

A_23_P64	NM_00192	Homo sapi	DCN,PG4(	1.29209	0.00001	1.29209	1605.604	2063.667
A_24_P30	NM_00452	lethal giant	LLGL2,HG	1.29183	5.97E-08	1.29183	1953.766	2524.049
A_24_P22	NM_18256	hypothetica	FLJ40319	1.29138	0.00019	1.29138	4772.629	6145.85
A_23_P15	NM_00431	Homo sapi	ARRB2,AR	1.29128	1.32E-06	1.29128	3742.365	4831.612
A_24_P221375		Unknown	A_24_P22	1.29116	3.73E-06	1.29116	140906.5	182523.5
A_32_P22	AK098506	Homo sapi	LOC28411	1.29109	0.00009	1.29109	1502.933	1929.959
A_32_P67	BC045163	Homo sapi	BC045163	1.29109	0.00059	1.29109	1100.004	1412.275
A_23_P91	NM_00195	Homo sapi	ECGF1,TP	1.29078	3.18E-07	1.29078	2312.785	2985.584
A_32_P72	NM_01450	ubiquitin ca	UBE2S,E2	1.29067	8.25E-08	1.29067	10169.69	13105.04
A_23_P35	AK057602	Ribosomal	RPL12	1.29066	3.81E-08	1.29066	204531.3	264358.1
A_24_P34	XM_37181	Homo sapi	XM_37181	1.29056	1.09E-08	1.29056	115632.2	149402.6
A_23_P21	AB002373	RUN and S	RUSC2	1.29055	1.65E-08	1.29055	4380.55	5672.289
A_23_P11	NM_00428	Homo sapi	H6PD,GDH	1.2902	0.00007	1.2902	6962.837	8964.625
A_24_P69	BC066359	CDNA clon	LOC55423	1.29014	1.52E-06	1.29014	1577.685	2036.266
A_23_P85	NM_02458	hypothetica	TMEM53,F	1.28996	1.45E-08	1.28996	2728.974	3519.329
A_24_P272403		Unknown	A_24_P27	1.28993	0.00008	1.28993	26497.34	34059.5
A_23_P64	NM_00356	Homo sapi	PLA2G6,G	1.28958	0.00007	1.28958	9123.052	11751.91
A_23_P10	AK000707	Hepatoma-	HDGF2	1.28955	0.00004	1.28955	7906.457	10168.89
A_24_P10	BC038434	Forkhead b	KIAA0415	1.28947	0.00016	1.28947	1216.675	1578.821
A_23_P14	NM_00455	Homo sapi	NFATC4,N	1.28936	0.0004	1.28936	1715.503	2192.635
A_24_P101742		Unknown	ENST0000	1.28927	1.37E-06	1.28927	4280.167	5523.989
A_23_P33	NM_01899	Homo sapi	CXorf9,SL	1.28903	6.17E-06	1.28903	1169.234	1507.593
A_24_P34	AK090466	Dolichyl-di	DDOST	1.28877	0.0007	1.28877	2902.209	3731.614
A_24_P383199		13.2 kDa p	ENST0000	1.28874	0.00006	1.28874	1389.128	1785.463
A_23_P44	NM_03305	Homo sapi	KRTAP4-1	1.28865	0.00023	1.28865	5444.641	6977.472
A_24_P16	NM_02493	hypothetica	FLJ12056	1.28852	0.00004	1.28852	1205.107	1558.742
A_23_P41	NM_00304	Homo sapi	SLC5A2,S	1.28842	0.00016	1.28842	12254.11	15679.81
A_24_P78	XM_29445	hypothetica	LOC34691	1.28828	0.00034	1.28828	7509.773	9658.016
A_24_P109973		Unknown	A_24_P10	1.28825	0.00098	1.28825	122601.1	159281.8
A_24_P59	AK093400	A disintegri	ADAM6	1.28796	0.00017	1.28796	1855.05	2377.594
A_23_P29	NM_14526	CG9886-lik	GLYCTK	1.28728	3.21E-06	1.28728	3853.67	5014.819
A_32_P36767		Unknown	A_32_P36	1.28718	0.00049	1.28718	872.6558	1137.448
A_23_P88	AK057899	Homo sapi	MCF2L,DB	1.28714	6.06E-06	1.28714	3898.361	5009.174
A_23_P33	AX721087	Sequence	AX721087	1.28689	0.00003	1.28689	3854.854	4949.945
A_23_P91	NM_00303	Homo sapi	SLC1A4,S	1.28685	0.00011	1.28685	1250.378	1603.14
A_24_P83	AK024371	Homo sapi	AK024371	1.28679	0.0004	1.28679	6281.523	8059.297
A_32_P92	BC052596	Hypothetic	LOC38856	1.2865	0.00018	1.2865	2712.782	3481.011
A_23_P50	NM_00178	cell divisior	CDC2L1,p	1.28643	2.45E-08	1.28643	2572.701	3309.554
A_23_P25	NM_01564	Homo sapi	C1QTNF5,	1.28619	0.00003	1.28619	1938.288	2506.163
A_32_P30	AB046827	WDFY fam	WDFY4	1.28612	0.00049	1.28612	824.7396	1062.99
A_32_P67837		Unknown	THC18189	1.28597	0.00002	1.28597	3642.228	4657.775
A_23_P37	NM_03226	hypothetica	C21orf56,M	1.28597	7.80E-07	1.28597	1165.905	1496.971
A_24_P79	XM_37990	Homo sapi	MUC12	1.28594	0.00002	1.28594	1003.32	1289.452
A_23_P15	AF338192	Homo sapi	FKSG63	1.28594	0.00001	1.28594	3628.868	4662.746
A_24_P35	NM_00200	fibroblast g	FGF4,HST	1.28594	0.00008	1.28594	1144.864	1479.206
A_32_P40	AF119863	Maternally	MEG3	1.2856	0.00047	1.2856	19666.79	25253.79
A_32_P19	AA150251	Protein kin	PRKCE	1.28526	3.56E-06	1.28526	1844.128	2375.351
A_24_P366566		Unknown	ENST0000	1.28513	0.00068	1.28513	11986.96	15370.38
A_23_P12	NM_00551	Major histo	HLA-B	1.28493	0.00001	1.28493	4909.077	6291.866
A_24_P92722		Unknown	A_24_P92	1.28478	0.00038	1.28478	10451.85	13548.26
A_32_P221437		D90774 Re	THC19540	1.28477	0.00043	1.28477	2658.209	3398.488
A_32_P22	NM_14460	hypothetica	MGC33367	1.28467	0.00019	1.28467	1391.602	1780.755

A_23_P98	NM_01748	Homo sapi	UBQLN3,T	1.28467	0.00082	1.28467	1142.332	1443.265
A_24_P80	NM_00561	ribosomal p	RPS14,EM	1.28413	0.00007	1.28413	49172.34	62901.89
A_23_P12	NM_00078	cytochrome	CYP11A1,	1.28401	0.00002	1.28401	3354.216	4294.584
A_23_P19	NM_01432	Homo sapi	HEBP2,PP	1.28389	3.02E-06	1.28389	7766.924	9942.563
A_24_P24	BC029274	CXYorf1 pr	CXYORF1	1.28382	1.61E-07	1.28382	7429.177	9530.678
A_24_P38	NM_00299	small induc	CX3CL1,N	1.28345	0.00011	1.28345	2937.108	3758.11
A_23_P65	NM_01585	Homo sapi	GTF2A1,TI	1.28327	0.00007	1.28327	981.3499	1259.875
A_23_P50	AK129550	Similar to F	LOC39094	1.28303	0.00098	1.28303	1117.077	1422.34
A_32_P175683		Olfactory re	ENST0000	1.28269	8.54E-08	1.28269	2706.028	3471.146
A_23_P15	NM_00190	Homo sapi	CTRL,CTR	1.28255	0.00006	1.28255	2808.262	3605.407
A_23_P11	NM_01657	8D6 antige	CD320,8D	1.28251	1.29E-07	1.28251	10102.09	12968.61
A_23_P29	NM_01570	CGI-96 pro	CGI-96,BK	1.28234	0.00014	1.28234	1213.671	1552.021
A_24_P204664		Unknown	A_24_P20	1.28232	0.00003	1.28232	19123.45	24599.37
A_23_P16	NM_02495	Homo sapi	UBTD1,FL	1.282	0.00048	1.282	765.0732	981.4144
A_24_P73	AY010112	Hypothetic	FLJ14082	1.28196	0.00079	1.28196	758.6871	968.5082
A_24_P92	BC017854	Epithelial m	EMP1	1.28181	5.45E-07	1.28181	2995.169	3830.238
A_24_P450870		RL39_HUM	THC18772	1.28157	0.00077	1.28157	5834.128	7482.682
A_23_P35	NM_13842	likely ortho	GRCC10	1.28143	1.38E-07	1.28143	12751.7	16361.12
A_24_P926849		Unknown	A_24_P92	1.281	4.45E-06	1.281	3369.635	4310.13
A_23_P20	AK093036	Zinc finger	ZNF577	1.28077	0.0008	1.28077	1784.677	2275.883
A_23_P14	NM_01431	Homo sapi	RASD2,Rh	1.28049	0.00002	1.28049	1010.179	1294.186
A_24_P10	NM_01343	nuclear fac	NFKBIL2,I	1.28045	3.75E-06	1.28045	2974.87	3804.236
A_32_P21	NM_14526	Homo sapi	SPATA18,I	1.2804	2.92E-06	1.2804	2060.25	2642.318
A_23_P51	NM_01584	Elastase 2I	ELA2B	1.2803	9.74E-08	1.2803	2979.991	3822.598
A_23_P20	NM_01775	hypothetica	FLJ20309	1.28003	5.37E-07	1.28003	5789.308	7404.455
A_23_P16	NM_07848	Homo sapi	SLC36A1,I	1.27993	0.00076	1.27993	1029.856	1305.093
A_32_P98	NM_00503	polymerase	POLRMT,A	1.27976	0.00003	1.27976	4811.554	6154.656
A_24_P789033		Unknown	THC18364	1.27971	0.00018	1.27971	5140.839	6547.792
A_24_P34	BC002886	Hypothetic	FAM27A,F	1.27949	0.00034	1.27949	5650.22	7219.88
A_32_P43914		Unknown	THC18378	1.27935	7.05E-08	1.27935	2502.173	3200.774
A_23_P27	NM_00157	Homo sapi	IRF3	1.27932	0.00096	1.27932	993.5707	1265.065
A_23_P16	NM_03065	NG3 protei	EGFL8,NG	1.27921	0.00003	1.27921	4829.794	6158.322
A_23_P37	AF426265	Chromosor	C21orf87	1.27915	1.64E-06	1.27915	4648.697	5938.167
A_24_P27	NM_00299	small induc	CXCL5,SC	1.27914	0.00087	1.27914	19216.29	24589.04
A_23_P14	NM_00598	Homo sapi	SPRR2A	1.27897	3.39E-06	1.27897	10647.7	13626.95
A_23_P30	NM_17315	Homo sapi	NR4A1,HM	1.27883	0.001	1.27883	4524.271	5795.55
A_23_P38	NM_13845	Homo sapi	TXNL6,RD	1.27882	0.00054	1.27882	3204.776	4082.221
A_24_P19	NM_00306	secretory le	SLPI,ALP,I	1.27881	5.71E-06	1.27881	6343.978	8112.841
A_24_P755169		Unknown	A_24_P75	1.2787	3.63E-06	1.2787	8085.677	10296.79
A_32_P10	AX721252	Sequence	AX721252	1.27856	4.34E-07	1.27856	50645.7	64871.23
A_32_P86905		Unknown	THC18081	1.27839	0.00004	1.27839	2292.042	2943.775
A_32_P11	AF086098	Hypothetic	LOC25740	1.27831	0.00005	1.27831	5820.774	7420.943
A_23_P79	AY305861	Ladybird h	LBX2	1.27823	0.00003	1.27823	1629.534	2090.831
A_32_P49	NM_19898	Homo sapi	LENG9	1.27812	0.00012	1.27812	2098.723	2642.365
A_32_P13081		Unknown	A_32_P13	1.27801	0.00043	1.27801	1315.219	1684.895
A_23_P25	NM_18264	Homo sapi	DLC1,HP,A	1.27798	0.00071	1.27798	1823.334	2321.831
A_23_P12	NM_00170	Homo sapi	BAI1	1.27788	0.00001	1.27788	3655.16	4669.406
A_23_P16	NM_01772	hypothetica	TRMT1,FL	1.27779	0.00018	1.27779	4081.098	5242.796
A_32_P60632		Unknown	A_32_P60	1.27777	0.00004	1.27777	1766.652	2252.052
A_24_P30	NM_03237	hypothetica	EEF1D,EF	1.27764	0.00008	1.27764	87236.75	110889.8
A_23_P82	NM_00321	TEA domai	TEAD3	1.27756	9.71E-09	1.27756	3718.389	4750.658
A_24_P32	AJ011409	Chromosor	C21orf104	1.27722	0.00023	1.27722	7972.528	10180.06

A_23_P36	NM_01656	Homo sapi	RLN3R1,S	1.2772	2.95E-06	1.2772	2065.065	2641.661
A_23_P50	NM_02161	Homo sapi	KCNN2,SK	1.27677	0.00056	1.27677	1622.979	2060.896
A_24_P86	AK092921	M4-50 mRI	AK092921	1.27651	0.00003	1.27651	13783.91	17670.27
A_23_P11	AL136837	Hypothetic	DKFZp434	1.27642	0.00007	1.27642	3654.125	4663.339
A_23_P12	NM_02101	Homo sapi	KCNJ12,IR	1.27627	0.00002	1.27627	1096.593	1400.604
A_23_P16	NM_03121	Chromosor	C19orf27	1.27624	0.00008	1.27624	1830.704	2342.311
A_23_P15	NM_00419	acetylserot	ASMTL,AS	1.27558	3.50E-06	1.27558	1693.52	2170.629
A_32_P64	BC062632	Hypothetic	LOC40059	1.2755	0.00097	1.2755	19856.23	25268.55
A_32_P198601		Unknown	A_32_P19	1.27548	0.00002	1.27548	4604.241	5845.517
A_23_P52	NM_18184	Homo sapi	NUDT8,FL	1.27545	1.79E-08	1.27545	4011.63	5116.842
A_24_P93	AY358191	LOC44052	NINP6167	1.27527	0.00001	1.27527	2913.832	3711.042
A_24_P31	BX248300	Rearrange	BX248300	1.27516	0.00033	1.27516	2720.386	3456.557
A_24_P26	NM_19853	short-chain	SCDR10	1.2751	0.00003	1.2751	2413.297	3065.362
A_24_P22	NM_00502	paired-like	PITX3,PTX	1.27505	1.34E-07	1.27505	2996.711	3830.698
A_23_P30	NM_02209	dJ467N11.	ASCC3,RN	1.27478	0.00047	1.27478	1262.383	1600.857
A_24_P59	NM_01410	PRO2000 p	ATAD2,MG	1.2745	0.00005	1.2745	1186.614	1509.127
A_23_P42	NM_17557	Homo sapi	WFIKKN2,	1.27443	0.00061	1.27443	1288.863	1629.518
A_23_P71	NM_20334	MSFL2541	UNQ2541	1.27433	0.00089	1.27433	980.1105	1241.003
A_24_P69	AK023769	Zinc finger	ZNF552	1.27414	0.00043	1.27414	1027.375	1314.513
A_23_P10	NM_00736	Homo sapi	GPR161,R	1.27385	1.57E-07	1.27385	4621.429	5885.387
A_32_P20	AF075027	Full length	AF075027	1.27373	0.00002	1.27373	4227.374	5390.338
A_24_P22	NM_00213	high-mobili	HMGA1,HM	1.2735	0.00093	1.2735	1254.101	1612.454
A_23_P32	NM_00523	Homo sapi	ETS1,ETS	1.27327	0.00095	1.27327	2370.386	3000.934
A_24_P16	NM_00168	ATP synth	ATP5D	1.27313	7.10E-06	1.27313	6803.858	8647.385
A_23_P38	NM_03237	hypothetica	EEF1D,EF	1.2731	0.00006	1.2731	28737.12	36386.83
A_32_P100076		Unknown	A_32_P10	1.27303	0.00006	1.27303	6105.245	7760.047
A_23_P13	NM_08091	Asialoglyco	ASGR2	1.273	0.00005	1.273	1723.057	2171.259
A_24_P21	NM_03229	hypothetica	GON4,GOI	1.27276	0.00031	1.27276	1372.054	1741.814
A_24_P72	NM_02502	hypothetica	FLJ14107	1.27274	0.00019	1.27274	1125.04	1428.505
A_24_P48	NM_19424	Homo sapi	HNRPA3,F	1.27274	0.00001	1.27274	3052.386	3878.754
A_24_P31	NM_00212	HLA-G hist	HLA-G,MH	1.27241	0.00001	1.27241	33478.23	42653.62
A_23_P20	NM_03236	Hypothetic	MGC1311	1.27224	0.00037	1.27224	18517.92	23649.73
A_23_P13	NM_03346	Homo sapi	MELL1,NL	1.27217	0.00002	1.27217	1699.952	2174.343
A_23_P21	NM_00089	Homo sapi	LHB,CGB4	1.27217	0.00072	1.27217	10703.43	13572.78
A_23_P14	NM_00280	proteasom	PSMB10,L	1.27204	3.62E-07	1.27204	13967.3	17741.89
A_24_P35	AL110245	Chromosor	C6orf85	1.27202	5.86E-07	1.27202	9012.529	11455.16
A_23_P55	NM_00076	Homo sapi	CYP2A13,	1.27199	1.18E-07	1.27199	3298.244	4193.208
A_23_P21	NM_05296	mas-relate	MAS1L,MF	1.27172	0.00003	1.27172	2089.608	2668.808
A_24_P40	BU603529	AGENCOL	BU603529	1.27169	0.00007	1.27169	1300.788	1660.462
A_24_P456043		Unknown	A_24_P45	1.27146	0.00034	1.27146	1373.376	1735.359
A_24_P69	AK125299	Similar to c	AK125299	1.27146	0.00024	1.27146	1273.464	1612.227
A_23_P13	NM_05304	Homo sapi	UCN3,SCF	1.27145	9.21E-06	1.27145	3452.261	4364.851
A_32_P180265		Unknown	THC18073	1.27101	0.00011	1.27101	1126.747	1430.802
A_32_P38	AK098534	Homo sapi	AK098534	1.27074	0.00005	1.27074	2933.71	3716.248
A_32_P12	XM_37076	similar to ri	LOC38799	1.27057	0.00006	1.27057	185334.1	236512
A_23_P25	AK001481	Vacuolar p	VPS13D	1.2705	0.00003	1.2705	1076.116	1367.559
A_24_P77	BC034005	Hypothetic	LOC38849	1.2704	1.12E-07	1.2704	6626.202	8407.752
A_23_P42	NM_00059	Compleme	C4A	1.27039	5.36E-08	1.27039	3108.32	3948.209
A_23_P10	NM_00618	Homo sapi	NTN2L	1.27008	0.00029	1.27008	1266.417	1616.943
A_23_P16	NM_03279	Homo sapi	ZNF499,FL	1.27006	0.00002	1.27006	3966.128	5036.58
A_23_P29	NM_00593	Homo sapi	AFF1,AF4,	1.27003	0.00005	1.27003	1417.338	1791.443
A_23_P101025		Unknown	A_23_P10	1.27002	0.00015	1.27002	1832.736	2315.787

A_32_P128960		Unknown	A_32_P12	1.26986	0.00013	1.26986	1090.139	1387.578
A_23_P43	NM_00613	Homo sapi	CD7,GP40	1.26985	0.0001	1.26985	1722.724	2201.352
A_23_P74	NM_03201	Homo sapi	STK40,SH	1.2694	0.00091	1.2694	1875.257	2375.1
A_23_P41	NM_17071	Homo sapi	RASSF1,1	1.26938	8.04E-06	1.26938	2568.895	3259.227
A_32_P42666		Unknown	THC19592	1.26928	2.91E-06	1.26928	6324.344	8018.789
A_23_P69	XM_11603	GABA-C re	GABRR3,D	1.2691	1.06E-06	1.2691	2029.105	2571.915
A_24_P39	AK124741	Similar to r	AK124741	1.269	1.94E-06	1.269	38318.57	48501.25
A_23_P37	BC000632	S100 calci	S100A13	1.26882	0.00003	1.26882	22968.92	29318.93
A_23_P33	NM_17350	Homo sapi	C16orf52	1.26848	0.00003	1.26848	1668.041	2112.564
A_32_P181822		Unknown	A_32_P18	1.26834	1.14E-06	1.26834	1281.416	1626.756
A_23_P23	NM_02108	Homo sapi	DAB1	1.26802	0.00022	1.26802	5053.674	6392.346
A_23_P32	NM_00238	Homo sapi	MC1R,MSH	1.26795	0.00002	1.26795	1662.99	2106.44
A_32_P18	NM_01450	ubiquitin ca	UBE2S,E2	1.26795	3.20E-06	1.26795	13970.26	17785.72
A_24_P23	NM_01450	putative Ra	RABGEF1,	1.26766	0.00038	1.26766	1278.385	1616.959
A_24_P109962		Unknown	A_24_P10	1.26765	1.71E-06	1.26765	19940.93	25236.52
A_32_P19	T12588	Transcribe	T12588	1.26731	2.41E-07	1.26731	19059.88	24177.88
A_24_P91	NM_17490	Homo sapi	C3orf6,YM	1.26729	0.00035	1.26729	1739.061	2213.557
A_24_P38	NM_00523	v-ets avian	ETS2	1.26721	0.00004	1.26721	1244.642	1575.543
A_24_P37	NM_00211	Major histo	HLA-A	1.26708	0.00062	1.26708	1737.992	2208.991
A_24_P409560		Unknown	A_24_P40	1.26704	0.00014	1.26704	8740.182	11126.68
A_23_P35	AK092792	Hypothetic	LOC38983	1.26702	0.00001	1.26702	1683.009	2131.517
A_24_P850172		Unknown	A_24_P85	1.26683	1.54E-07	1.26683	24549.95	31073.6
A_23_P82	NM_00321	TEA domai	TEAD3	1.26672	0.00012	1.26672	9696.03	12178.68
A_23_P14	NM_02181	Homo sapi	LMAN1L,E	1.26658	7.15E-07	1.26658	4129.595	5219.563
A_23_P33	XM_37740	Similar to p	ENST0000	1.26655	0.00003	1.26655	2875.897	3638.063
A_32_P226869		RS25_HUM	THC18783	1.26633	7.93E-09	1.26633	10670.74	13512.39
A_23_P37	BC035647	M4-50 mRI	BC035647	1.26597	1.06E-07	1.26597	5991.383	7588.854
A_24_P28	NM_00225	Killer cell ir	KIR2DL4	1.26579	0.00018	1.26579	991.316	1258.159
A_23_P103315		Unknown	A_23_P10	1.26574	0.00008	1.26574	111417.3	141411.4
A_23_P14	NM_03295	Homo sapi	WBSCR14	1.26572	0.00003	1.26572	2162.26	2729.999
A_24_P678826		Unknown	THC18725	1.26552	0.0002	1.26552	3724.694	4709.493
A_23_P20	AB011085	KIAA0513	KIAA0513	1.26545	3.94E-06	1.26545	1473.191	1863.153
A_23_P27	NM_01226	Homo sapi	PLD3,HU-K	1.26536	0.00059	1.26536	1101.427	1387.853
A_23_P58	NM_18188	Homo sapi	UBE2D3,U	1.26531	0.00014	1.26531	5342.142	6781.458
A_32_P49	NM_17488	Myc-induce	mimitin,B1	1.26515	0.00005	1.26515	8901.494	11306.74
A_23_P71	NM_03076	Homo sapi	AKNA,KIA	1.26492	0.00055	1.26492	1133.867	1443.838
A_32_P31	NM_19912	Homo sapi	GGTL4,MG	1.26476	1.11E-07	1.26476	4177.54	5282.717
A_23_P56	NM_01443	Homo sapi	IL1F7,FIL1	1.26466	0.00046	1.26466	951.7826	1199.308
A_24_P18	NM_01663	SRp25 nuc	ARL6IP4,S	1.26465	0.00004	1.26465	6713.799	8455.466
A_32_P21	BX091616	Transcribe	BX091616	1.26464	0.00009	1.26464	2527.477	3189.879
A_24_P94	NM_19808	hypothetica	LOC28400	1.26464	0.00013	1.26464	3869.819	4879.307
A_23_P25	NM_00628	Homo sapi	SSR4,TRA	1.26459	4.03E-06	1.26459	19306.82	24359.03
A_23_P36	NM_13837	hypothetica	MGC1604	1.26443	0.00029	1.26443	2344.934	2952.797
A_32_P234570		Unknown	THC19462	1.26433	0.00043	1.26433	26347.09	33401.91
A_23_P10	NM_00136	deiodinase	DIO3,D3,5	1.26431	0.00099	1.26431	938.9314	1199.877
A_23_P36	NM_17488	Myc-induce	mimitin,B1	1.26419	8.35E-06	1.26419	10849.91	13762.21
A_23_P33	NM_15248	Homo sapi	C19orf23,M	1.26414	0.0005	1.26414	1907.643	2399.829
A_23_P25	NM_01648	Homo sapi	C9orf156,N	1.26404	0.00026	1.26404	1119.176	1416.681
A_24_P25	AK055463	Homo sapi	FLJ30901	1.26368	1.26E-07	1.26368	5363.033	6764.109
A_24_P15	NM_01866	papillomav	ZNF395,PE	1.26351	0.00069	1.26351	1964.482	2500.406
A_23_P42	NM_13363	Homo sapi	RHOV,Chp	1.26329	0.00002	1.26329	1444.255	1825.468
A_24_P41	BC020891	HLA-G hist	HLA-G	1.26305	0.00004	1.26305	13231.99	16735.6

A_23_P96	NM_03264	hypothetica	C1orf90,M	1.26301	8.31E-06	1.26301	1547.323	1955.728
A_23_P25	NM_03251	Homo sapi	BOK,BOKL	1.26292	0.00076	1.26292	942.1642	1192.521
A_32_P66	BM981501	Similar to 6	BM981501	1.26284	0.00013	1.26284	6714.179	8460.93
A_23_P33	NM_14517	Homo sapi	GPHB5,GF	1.26278	0.00002	1.26278	2602.97	3295.142
A_23_P53	NM_01672	Homo sapi	FOLR1,FB	1.2627	5.31E-08	1.2627	4654.349	5878.138
A_32_P160471		Unknown	A_32_P16	1.26256	0.00014	1.26256	3437.07	4333.422
A_24_P480722		Unknown	A_24_P48	1.26221	0.00076	1.26221	15767.29	19870.96
A_24_P64	NM_03277	hypothetica	MGC16291	1.26217	0.00028	1.26217	1035.736	1304.736
A_23_P21	NM_00097	Homo sapi	RPL12	1.26204	1.51E-06	1.26204	311279.6	393757.4
A_24_P92	XM_06053	similar to ri	LOC12754	1.26203	0.00002	1.26203	28864.79	36521.82
A_24_P392863		Unknown	ENST0000	1.26193	5.83E-07	1.26193	4060.261	5128.718
A_23_P38	NM_02453	Homo sapi	CORO7,CF	1.2619	0.00061	1.2619	3762.962	4797.981
A_23_P94	NM_00070	Homo sapi	ANXA1,AN	1.26188	0.00003	1.26188	67880.61	85776.61
A_23_P11	NM_00313	Serum resp	SRF	1.26152	0.00075	1.26152	1432.784	1784.13
A_23_P36	NM_03351	Homo sapi	LIMS3	1.26137	0.00083	1.26137	1036.979	1312.806
A_23_P16	NM_00010	Homo sapi	CYBA	1.26136	5.19E-08	1.26136	8269.203	10427.67
A_24_P22	NM_00080	gamma-am	GABRA1,E	1.26135	0.00035	1.26135	1237.815	1569.415
A_32_P23	BX105253	Transcribe	BX105253	1.26105	0.00005	1.26105	6000.083	7549.517
A_24_P27	NM_17304	Homo sapi	IL18BP,IL1	1.2607	0.00008	1.2607	4103.19	5151.893
A_24_P15	NM_00322	ADP-ribosy	ARFRP1,A	1.26066	0.00007	1.26066	1104.929	1392.556
A_24_P92	AF130065	Clone FLB	PRO2158	1.26057	0.0005	1.26057	899.6311	1126.884
A_23_P20	AK027798	Transmem	MSP	1.26029	0.00024	1.26029	983.9847	1241.221
A_23_P20	NM_00272	Homo sapi	PPY,PNP	1.25996	9.42E-06	1.25996	2317.606	2900.334
A_23_P60	NM_15320	Homo sapi	EDF1,MBF	1.25994	6.00E-07	1.25994	30769.12	38705.15
A_24_P27	AK000313	Homo sapi	FLJ20306	1.2598	0.0001	1.2598	1674.488	2100.634
A_24_P91	NM_13921	homeodom	HOP,OB1,	1.25978	0.00002	1.25978	1138.431	1439.847
A_23_P40	AK026312	Hypothetic	FLJ22659	1.25959	0.00044	1.25959	1235.042	1547.842
A_24_P85	NM_00484	basement	C1orf38,IC	1.25951	0.00005	1.25951	1243.211	1562.612
A_24_P10	NM_03278	Homo sapi	RSPO3,PV	1.25944	0.0001	1.25944	1665.151	2094.308
A_24_P22	NM_00227	keratin 13	KRT13,K1	1.2594	0.00016	1.2594	1069.561	1346.039
A_23_P10	NM_01745	microfibrilla	MFAP2,MA	1.25936	0.00041	1.25936	2265.336	2837.082
A_23_P41	NM_17688	Homo sapi	TAS2R44,	1.25933	0.00017	1.25933	5034.792	6321.523
A_23_P36	NM_00525	Homo sapi	FOXL1,FK	1.25843	0.00001	1.25843	5198.889	6557.641
A_23_P117912		AY059636	THC20037	1.2582	0.00014	1.2582	1640.696	2067.767
A_24_P38	XM_29332	BA370B6.1	ENST0000	1.25817	0.00001	1.25817	2293.56	2886.813
A_24_P88	AK094353	Hypothetic	FLJ37034	1.2581	0.0002	1.2581	1956.239	2449.649
A_24_P41551		Unknown	A_24_P41	1.25776	0.00003	1.25776	83246.36	104955.6
A_24_P60	L38562	Anti-c-erbB	IGL@,IGL	1.25756	0.00003	1.25756	1594.887	2008.911
A_23_P25	NM_00346	zyxin	ZYX,ESP-2	1.25754	0.00031	1.25754	1239.631	1553.863
A_23_P20	NM_02400	Homo sapi	GJB3,CX3	1.25738	0.00012	1.25738	3988.771	5031.377
A_32_P12	BC037580	Pseudoger	BC037580	1.25724	8.43E-06	1.25724	2321.013	2915.998
A_23_P50	NM_00168	Homo sapi	ATP5D	1.25719	0.00006	1.25719	5170.271	6484.393
A_23_P20	NM_02042	CTL2 prote	SLC44A2,	1.25714	0.00064	1.25714	7128.905	8950.184
A_23_P37	NM_17843	Homo sapi	LCE2D,LE	1.25706	0.00047	1.25706	21289.14	26749.4
A_24_P289884		Unknown	A_24_P28	1.25698	0.00046	1.25698	36921.42	46586.77
A_24_P93	AF311912	Secreted fr	SFRP2	1.2569	0.0006	1.2569	1393.804	1761.872
A_23_P54	NM_01654	Homo sapi	GNG13,h2	1.25683	0.00043	1.25683	6659.278	8347.138
A_23_P11	NM_00506	Homo sapi	RPL3L	1.2566	0.00016	1.2566	2129.568	2665.376
A_23_P43	NM_00312	sialophorin	SPN,LSN,	1.25649	0.00075	1.25649	7148.542	8969.152
A_32_P21	AK128072	Malate deh	MDH2	1.25635	0.00003	1.25635	1426.675	1791.079
A_23_P14	BC004960	LOC44086	MGC10955	1.25625	0.00003	1.25625	1714.724	2151.447
A_24_P267686		27228 men	THC19301	1.25609	0.00011	1.25609	5328.405	6657.325

A_24_P20	NM_00172	basigin (OI	BSG,M6,O	1.25571	0.00003	1.25571	2252.562	2828.545
A_23_P24	NM_00035	Homo sapi	ABCC8,HI,	1.2556	0.00001	1.2556	1019.149	1280.926
A_24_P272590		R3HU15 ri	THC20072	1.25522	4.33E-06	1.25522	153266.5	192763.5
A_32_P20	AK124956	CDNA FLJ	AK124956	1.25509	0.00085	1.25509	1287.516	1623.412
A_32_P13	AK001432	LOC44003	AK001432	1.25504	0.00011	1.25504	2908.261	3666.364
A_23_P12	NM_00710	ATP synth	ATP5I,ATF	1.25503	0.00016	1.25503	30888.13	38539.34
A_23_P54	NM_01435	Homo sapi	RAB26,V4	1.25455	0.00007	1.25455	1338.905	1676.164
A_24_P703830		GL6S_CAF	THC18921	1.25444	0.00026	1.25444	1175.081	1468.796
A_32_P149625		Unknown	THC19143	1.25418	0.00011	1.25418	1696.749	2123.736
A_32_P105730		Unknown	THC19096	1.25398	0.00006	1.25398	5158.432	6430.515
A_24_P27	NM_03330	hypothetic	MGC4655	1.25352	1.21E-07	1.25352	3688.959	4624.576
A_24_P315769		Unknown	A_24_P31	1.25343	1.78E-07	1.25343	66807.28	83765.3
A_23_P12	NM_03282	hypothetic	FLJ14721	1.2534	0.0001	1.2534	1405.318	1758.75
A_24_P30	NM_00033	solute carri	SLC12A3,T	1.25339	0.00036	1.25339	730.1663	915.4276
A_23_P18	NM_15254	hypothetic	FLJ35725,	1.25319	0.00016	1.25319	1381.177	1725.578
A_23_P20	NM_13084	Homo sapi	SLC39A4,A	1.25316	4.58E-06	1.25316	4517.075	5656.082
A_23_P25	NM_00308	Homo sapi	SNCG,SR,	1.25283	0.00006	1.25283	1853.297	2324.82
A_23_P30	NM_14520	Homo sapi	CABP4	1.25275	0.00026	1.25275	2523.532	3144.415
A_24_P36	NM_00478	solute carri	SLC9A3R2	1.25275	0.00089	1.25275	3042.674	3798.325
A_24_P76	AJ296345	Keratin ass	KRTAP2-1	1.25241	5.56E-08	1.25241	7215.292	9034.751
A_32_P225173		Unknown	THC18827	1.25216	0.00075	1.25216	2082.955	2596.802
A_23_P71	NM_20692	apical early	MAMDC4,A	1.25212	0.00015	1.25212	1259.989	1575.891
A_23_P25	NM_01848	hepatocell	HCA112,G	1.25192	0.00014	1.25192	1182.206	1490.05
A_23_P11	M21963	Major histo	HLA-C	1.25179	0.00005	1.25179	51808.91	65124.05
A_24_P36	NM_00422	Homo sapi	ZNF213,CI	1.25166	0.00044	1.25166	11208.94	13995.53
A_32_P152348		Unknown	A_32_P15	1.25149	0.00024	1.25149	2949.108	3697.178
A_24_P22	NM_03143	SOX7 tran	SOX7,MG	1.25148	0.00009	1.25148	2152.994	2705.681
A_23_P67	NM_01961	Immunity-r	R30953_1	1.25143	0.00086	1.25143	1093.936	1363.737
A_23_P90	AF200344	Napsin B a	NAPSB	1.25139	1.68E-06	1.25139	1965.087	2458.029
A_23_P10	NM_03244	Homo sapi	FBN3,KIA	1.25133	0.00001	1.25133	3368.044	4207.974
A_23_P19	NM_14568	Homo sapi	APBB1,RIF	1.2513	0.00004	1.2513	3170.993	3955.001
A_23_P60	AF152351	Homo sapi	TIMM13a	1.25112	1.13E-06	1.25112	8501.375	10646.57
A_23_P59	NM_03192	Homo sapi	RSHL2,RS	1.25099	0.00079	1.25099	1293.656	1611.137
A_23_P82	NM_00605	Homo sapi	LAMC3,DK	1.25085	4.63E-07	1.25085	3728.836	4661.984
A_24_P76	NM_01343	gamma-glu	GGT1,GGT	1.25075	2.39E-06	1.25075	4003.217	5004.083
A_24_P144254		Unknown	A_24_P14	1.25072	8.87E-07	1.25072	14296.11	17898.17
A_23_P13	NM_00348	Myeloid/ly	MLL2	1.2507	0.00086	1.2507	1187.209	1493.517
A_24_P31	AK093659	Bone morp	BMP8A	1.25064	0.00064	1.25064	807.7703	1008.124
A_23_P10	NM_00241	Homo sapi	MGMT	1.25051	6.48E-06	1.25051	2519.902	3151.891
A_24_P375227		F20887_1,	ENST0000	1.25042	0.00029	1.25042	1085.141	1355.031
A_23_P10	NM_02521	Homo sapi	WNT10A,F	1.25029	0.00002	1.25029	1021.245	1276.714
A_23_P58	NM_01617	hypothetic	LOC51149	1.25028	0.00068	1.25028	879.291	1102.423
A_24_P23	NM_00407	chloride ch	CLCNKA,h	1.25022	0.00005	1.25022	1978.693	2469.031
A_24_P99	NM_02471	hypothetic	ISOC2	1.25011	0.00009	1.25011	2377.531	2976.889
A_23_P66	NM_01458	Homo sapi	SOX8,MG	1.25002	0.00077	1.25002	9814.467	12231.16
A_23_P16	NM_01768	Homo sapi	VMD2L1,F	1.24997	0.00003	1.24997	2163.134	2698.46
A_24_P59	NM_08083	Homo sapi	GGTL4,MG	1.24991	2.50E-06	1.24991	2208.96	2762.708
A_24_P22	NM_00101	ribosomal	RPS15,RIC	1.2498	0.00006	1.2498	189132.1	237103.5
A_24_P35	NM_01500	SMART/HI	SPEN,MIN	1.24956	0.00062	1.24956	1550.031	1953.296
A_24_P86	AK095081	Hypothetic	LOC28317	1.2493	0.00028	1.2493	4739.234	5907.38
A_24_P926115		Unknown	A_24_P92	1.24929	0.00025	1.24929	8966.885	11181.88
A_24_P118341		Unknown	A_24_P11	1.24922	0.00016	1.24922	2344.981	2927.359

A_23_P97	NM_05305	LYST-inter	LIP8	1.24911	0.00004	1.24911	7555.722	9421.337
A_24_P289753		Unknown	A_24_P28	1.24907	0.00005	1.24907	7341.263	9145.855
A_32_P11	AW901958	Protein kin	PRKCB1	1.24886	0.00003	1.24886	1697.263	2118.203
A_32_P111507		Unknown	A_32_P11	1.24855	2.24E-07	1.24855	21557.19	26929.31
A_23_P14	AJ457067	Keratin ass	KRTAP19-	1.24842	0.00002	1.24842	6191.099	7717.951
A_23_P15	BC050733	ATP-bindin	ABCC6	1.24839	0.00064	1.24839	3372.984	4197.563
A_24_P31	NM_03229	hypothetica	FLYWCH1	1.24837	0.00032	1.24837	1074.832	1341.01
A_32_P19	NM_01899	hypothetica	KCTD5,FL	1.24836	0.0007	1.24836	3798.4	4743.304
A_23_P10	NM_03141	MAP/micro	MARK4,M	1.24826	0.0001	1.24826	1223.362	1526.999
A_24_P384469		Unknown	A_24_P38	1.24824	0.00057	1.24824	4452.034	5542.337
A_23_P15	NM_03234	Homo sapi	C6orf125,M	1.24824	6.33E-07	1.24824	9505.806	11855.3
A_24_P195272		Putative G-	ENST0000	1.2482	4.05E-06	1.2482	2415.712	3014.639
A_23_P15	NM_00684	hepatitis de	DIPA	1.24809	0.00002	1.24809	3768.464	4700.229
A_23_P94	NM_17503	Homo sapi	ST6GALNA	1.24797	0.00005	1.24797	3108.468	3874.435
A_23_P49	NM_00680	Homo sapi	KDELR1,E	1.24786	0.00014	1.24786	5338.434	6651.434
A_32_P16	NM_00149	GDNF fam	GFRA2,NT	1.24746	0.00082	1.24746	3421.342	4252.542
A_23_P31	NM_03237	hypothetica	EEF1D,EF	1.24732	5.98E-08	1.24732	140390.8	175099
A_32_P109002		Unknown	THC18082	1.24719	0.00022	1.24719	876.878	1091.377
A_23_P25	NM_01450	putative Ra	RABGEF1,	1.24715	0.00065	1.24715	1250.353	1563.072
A_23_P30	NM_13905	Homo sapi	ARX,ISSX,	1.24694	7.32E-06	1.24694	2411.869	3004.047
A_23_P20	NM_00101	Homo sapi	RPS15,RIC	1.24658	5.72E-06	1.24658	172696.7	215814.1
A_24_P91	M55405	Mucin 3A, i	MUC3A	1.24642	0.00002	1.24642	7586.115	9437.79
A_32_P47107		Unknown	A_32_P47	1.2464	9.57E-06	1.2464	7470.218	9305.725
A_23_P31	NM_00710	Homo sapi	TCEB2,SII	1.24639	0.00001	1.24639	62365.09	77547.14
A_24_P83	M30474	Gamma-gl	GGT2	1.24589	0.0001	1.24589	2342.993	2916.167
A_23_P50	NM_14464	Homo sapi	IL17RE,FL	1.24578	0.00006	1.24578	6023.109	7494.109
A_24_P26	NM_02107	latent trans	LTBP3,LTE	1.24562	0.00002	1.24562	2712.4	3376.029
A_24_P736638		Unknown	A_24_P73	1.24553	0.00004	1.24553	20932.87	26179.79
A_23_P20	NM_00696	zinc finger	ZNF24,KO	1.24492	0.00003	1.24492	13433.62	16757.71
A_23_P15	NM_03251	Homo sapi	MAP1LC3A	1.24476	0.0001	1.24476	2414.783	2999.513
A_24_P33	NM_02037	organic cat	SLC22A17	1.24474	2.08E-07	1.24474	2998.586	3732.857
A_32_P2470		ALU7_HUM	THC19937	1.2447	0.00024	1.2447	1757.549	2195.245
A_23_P12	NM_02466	Homo sapi	NPEPL1,F	1.24456	0.00001	1.24456	10517.2	13093.84
A_32_P40250		Unknown	THC19464	1.24434	0.0001	1.24434	1859.954	2309.973
A_24_P10	NM_00455	nuclease s	YBX1,YB1	1.24423	2.88E-06	1.24423	14446.25	17948.66
A_23_P66	NM_00002	Homo sapi	SGCA,A2,A	1.24421	7.27E-06	1.24421	1628.545	2021.036
A_24_P836315		ALU1_HUM	THC18921	1.24418	0.0007	1.24418	1877.728	2321.705
A_24_P70	AF085886	FLJ46675 p	FLJ46675	1.24371	0.00005	1.24371	19374.2	24025.41
A_24_P52	XM_37808	Homo sapi	ENST0000	1.24365	7.19E-07	1.24365	3100.2	3856.058
A_24_P573834		PHB_MOU	THC19266	1.24359	4.79E-06	1.24359	3231.05	4014.055
A_24_P761490		Unknown	A_24_P76	1.24352	1.83E-06	1.24352	6933.849	8631.269
A_32_P116113		Unknown	A_32_P11	1.24346	0.00006	1.24346	1449.288	1800.684
A_24_P27	NM_00074	cholinergic	CHRNB1,A	1.24346	0.00087	1.24346	1212.008	1500.12
A_23_P14	NM_01895	Homo sapi	HLA-F,HLA	1.24345	9.36E-06	1.24345	3943.729	4900.577
A_23_P24	NM_03212	hypothetica	DKFZP566	1.24323	0.00033	1.24323	6335.095	7868.676
A_24_P74	NM_17562	Homo sapi	MT1J,MT1	1.24307	0.00003	1.24307	1812.703	2249.437
A_24_P39	BC038715	Purkinje ce	PCP2	1.24303	0.00034	1.24303	1821.99	2271.651
A_24_P10	NM_01638	natural kille	CD244,2B4	1.24296	8.26E-06	1.24296	1969.6	2450.265
A_23_P75	NM_13080	Homo sapi	MEN1,MEA	1.24295	0.00047	1.24295	4860.332	6036.702
A_24_P37	NM_15230	Homo sapi	C14orf172,	1.24288	0.00083	1.24288	3203.253	3971.945
A_23_P10	NM_00497	Homo sapi	KCNC3,KV	1.24268	0.00005	1.24268	2662.041	3313.25
A_24_P84	AK090418	Hypothetic	LOC34919	1.24246	0.00006	1.24246	1859.027	2316.051

A_23_P54	AF119842	Homo sapi	PRO1048	1.24229	4.28E-06	1.24229	2941.016	3650.474
A_24_P25	AK092418	Fc fragmer	FCGR1A	1.24219	0.00039	1.24219	2759.417	3408.009
A_23_P14	NM_02477	hypothetica	FLJ13798	1.24214	0.0007	1.24214	3721.246	4623.209
A_24_P204484		Unknown	A_24_P20	1.24211	0.00067	1.24211	13752.78	16991.4
A_24_P23	BC063888	Homo sapi	BC063888	1.2421	9.52E-06	1.2421	37020.09	45910.34
A_23_P15	NM_02451	pre-mRNA	WDR25,M	1.24187	0.00005	1.24187	3540.753	4390.234
A_32_P27	NM_00127	chromogra	CHGA,CG	1.24173	0.00002	1.24173	4480.017	5555.546
A_24_P32	BC010562	Homo sapi	MGC1541	1.24165	0.0001	1.24165	3806.639	4696.172
A_24_P14	NM_03147	hypothetica	INHBE,MG	1.24156	0.00072	1.24156	1143.532	1427.04
A_24_P187626		Unknown	A_24_P18	1.24156	7.87E-06	1.24156	32329.74	40108.43
A_23_P39	AK124717	Purkinje ce	PCP2	1.24153	0.00003	1.24153	12068.37	15023.99
A_24_P97	NM_01332	cytokine in	CISH,CIS,	1.2415	0.00002	1.2415	1584.269	1960.536
A_24_P178300		unknown [	ENST0000	1.24145	0.00014	1.24145	5513.312	6822.19
A_23_P31	NM_14500	hypothetica	TSNARE1,	1.24139	2.95E-06	1.24139	2905.731	3601.342
A_24_P18	NM_01467	KIAA0685	SAPS2,SA	1.24138	0.00039	1.24138	1082.285	1338.816
A_24_P79	NM_20659	Homo sapi	ESRRG,E	1.24134	0.00004	1.24134	1693.172	2097.519
A_23_P12	NM_01758	Homo sapi	C9orf7,D9	1.24115	0.00008	1.24115	3465.441	4293.292
A_23_P20	AK126752	Ubiquitin s	USP31	1.241	0.00052	1.241	1909.059	2360.341
A_23_P37	NM_00560	nitrilase 1	NIT1,MGC	1.24099	3.50E-06	1.24099	2158.333	2677.712
A_23_P14	NM_01223	Homo sapi	SIRT2,SIR	1.24078	0.00006	1.24078	5094.289	6310.71
A_23_P47	NM_13076	Homo sapi	GPHA2,G	1.24061	0.00094	1.24061	2303.155	2849.725
A_24_P11	BC063512	Homo sapi	LOC34818	1.24042	0.00006	1.24042	4895.295	6086.854
A_23_P23	NM_01484	Homo sapi	SV2A,SV2	1.24008	0.00022	1.24008	6341.306	7854.685
A_23_P38	NM_01617	Homo sapi	TLX2,Enx,	1.2399	0.00037	1.2399	789.4705	978.1428
A_24_P366768		Unknown	A_24_P36	1.23989	0.00026	1.23989	4220.642	5247.19
A_23_P10	NM_00230	lectin, gala	LGALS7,G	1.23922	0.00073	1.23922	817.8846	1016.516
A_24_P14	AF318337	LOC44013	pp13759	1.23908	0.00006	1.23908	5856.002	7266.981
A_23_P41	NM_00170	Homo sapi	OPN1SW,	1.23895	0.00069	1.23895	3402.979	4144.552
A_23_P16	NM_00725	Homo sapi	PNKP,PNK	1.23876	0.00002	1.23876	2766.332	3420.325
A_23_P15	NM_00556	lectin, gala	LGALS3BP	1.23855	0.00001	1.23855	4944.883	6127.3
A_24_P484357		Unknown	A_24_P48	1.23853	8.17E-06	1.23853	2413.927	2993.06
A_32_P11501		Unknown	A_32_P11	1.23841	0.00026	1.23841	1348.194	1664.724
A_24_P75	AY312160	Mucin 6, g	MUC6	1.23832	0.00053	1.23832	2996.147	3702.594
A_24_P33	NM_00300	secreted a	SECTM1,K	1.2381	0.00022	1.2381	6594.667	8135.638
A_32_P46112		Unknown	A_32_P46	1.23808	0.00006	1.23808	5788.792	7158.653
A_23_P23	NM_01211	Homo sapi	CBX5,HP1	1.23802	0.00068	1.23802	1813.438	2241.896
A_24_P14	NM_01749	seb4D	RNPC1,SE	1.23798	0.00009	1.23798	2037.915	2519.754
A_24_P83	BC037244	Similar to u	BC037244	1.23789	0.00062	1.23789	3114.552	3872.966
A_24_P179033		Unknown	A_24_P17	1.23774	8.78E-06	1.23774	37382.01	46192.78
A_23_P25	NM_00688	Homo sapi	ATP5E,AT	1.2376	3.42E-06	1.2376	91310.03	113165.4
A_24_P20	AF174045	Ig rearrang	IgH	1.23751	0.00006	1.23751	2662.858	3288.562
A_23_P16	NM_00306	Homo sapi	SLIT1,MEC	1.2373	0.00019	1.2373	2126.051	2637.287
A_32_P217523		R5HU32 ri	THC19766	1.23723	0.001	1.23723	27249.35	33883.68
A_23_P23	NM_00029	Homo sapi	PKP1,B6P	1.23721	0.00048	1.23721	1309.391	1625.358
A_23_P39	NM_00057	Homo sapi	SLC11A1,	1.2372	0.00002	1.2372	3873.342	4785.909
A_24_P18	NM_01417	HSPC141	PHPT1,PH	1.23667	0.0001	1.23667	20726.62	25716.32
A_23_P38	NM_17849	hypothetica	FLJ23657	1.23625	0.00088	1.23625	1078.729	1336.871
A_24_P41	NM_00073	cholinergic	CHRM1,M	1.23622	0.00005	1.23622	2749.969	3408.353
A_23_P35	NM_00285	poliovirus r	PVRL2,HV	1.23612	0.0005	1.23612	875.7435	1078.611
A_32_P18	BM678546	Activating t	ATF4	1.23607	6.99E-06	1.23607	2209.959	2730.863
A_23_P13	NM_00531	granzyme	GZMM,ME	1.23598	3.29E-06	1.23598	5167.573	6410.321
A_23_P16	NM_02216	Homo sapi	LIN7B,VEL	1.23576	0.00015	1.23576	3665.609	4525.159

A_24_P184	NM_14530	Homo sapi	LRRC51	1.23565	0.00009	1.23565	4163.753	5150.355
A_23_P71	NM_00047	Homo sapi	TWIST1,S	1.23542	4.73E-07	1.23542	4062.367	5018.757
A_32_P52	BC011002	Hypothetic	LOC11323	1.23529	0.00088	1.23529	7300.497	9011.953
A_23_P20	NM_00193	Homo sapi	DPT,TRAM	1.235	0.00083	1.235	7684.032	9367.828
A_24_P570930		Unknown	A_24_P57	1.23422	9.72E-06	1.23422	1679.673	2073.205
A_23_P25	NM_00365	cold shock	CSDA,DBF	1.23376	0.00001	1.23376	5739.169	7078.286
A_24_P18	NM_02277	hypothetica	FLJ12681,	1.23366	0.00008	1.23366	4308.033	5304.265
A_24_P28	NM_02277	hypothetica	RABL5,FL	1.23339	0.00004	1.23339	3187.019	3928.477
A_23_P16	NM_02405	cross-imm	PCIA1,MG	1.23333	0.00005	1.23333	1914.442	2359.27
A_24_P34	NM_14569	Homo sapi	BRF1,BRF	1.23328	0.00021	1.23328	2551.537	3128.351
A_23_P50	NM_00129	Homo sapi	CLPTM1	1.2331	0.00032	1.2331	1230.086	1519.053
A_32_P21	BC042649	Homo sapi	BC042649	1.23304	0.00008	1.23304	3515.733	4327.755
A_23_P14	NM_00099	ribosomal	RPL37A,M	1.23296	1.18E-06	1.23296	257488.3	317705.1
A_23_P50	NM_17797	Homo sapi	CHRD	1.23285	9.26E-06	1.23285	2762.927	3408.344
A_23_P97	NM_00356	Homo sapi	BCAR3,NS	1.23283	0.00065	1.23283	4090.059	5048.319
A_24_P204084		S55917 rib	THC18551	1.23281	0.00001	1.23281	68334.97	84088.95
A_24_P50057		Unknown	ENST0000	1.23267	0.00002	1.23267	2639.562	3253.467
A_24_P666795		Unknown	A_24_P66	1.23251	0.00062	1.23251	1892.416	2317.493
A_32_P39	U79275	Hypothetic	HSU79275	1.23237	0.00035	1.23237	5550.824	6834.695
A_32_P94354		Unknown	A_32_P94	1.23236	7.14E-06	1.23236	2236.857	2755.719
A_32_P72	NM_00720	ribosomal	RPL35	1.23226	1.73E-06	1.23226	37767.42	46501.92
A_32_P217561		Unknown	THC18525	1.23199	0.00002	1.23199	1719.314	2111.629
A_23_P33	NM_02112	Homo sapi	POLR2L,R	1.23199	2.43E-06	1.23199	101046.3	124368.6
A_23_P15	BC012609	Serine (or	SERPINB2	1.23194	0.00084	1.23194	3150.665	3815.272
A_24_P11	NM_01550	adaptin-ea	NECAP1	1.23188	0.00002	1.23188	5288.993	6515.458
A_24_P22	NM_00689	carcinoemb	CEACAM7	1.23159	0.00021	1.23159	1559.401	1916.273
A_24_P25	NM_02486	Hypothetic	FLJ14050	1.23157	4.29E-06	1.23157	3667.782	4517.495
A_23_P35	NM_00024	Homo sapi	CIITA,C2T	1.23146	0.00013	1.23146	4910.91	6039.016
A_24_P350008		Unknown	A_24_P35	1.23143	0.00001	1.23143	28997.78	35639.99
A_24_P383901		Unknown	A_24_P38	1.23136	2.79E-07	1.23136	26934.83	33166.7
A_32_P83	AF128541	Polymeras	POLE	1.2313	1.05E-06	1.2313	3907.889	4811.202
A_23_P21	NM_00207	guanine nu	GNAZ	1.2312	3.41E-06	1.2312	7889.621	9709.115
A_23_P85	NM_02164	Fc fragmer	FCGR2A	1.23083	0.00058	1.23083	1383.146	1709.912
A_24_P25	NM_02432	hypothetica	EFHD2,MC	1.23066	0.00067	1.23066	1523.72	1873.812
A_23_P16	NM_01983	Homo sapi	LTB4R2,BI	1.2305	0.0005	1.2305	3479.967	4274.074
A_24_P392910		Unknown	A_24_P39	1.23048	0.00003	1.23048	1748.074	2146.279
A_23_P25	NM_02163	apoptosis i	FKSG2	1.23016	0.00032	1.23016	20208.54	24886.46
A_24_P297798		Unknown	ENST0000	1.23008	6.06E-07	1.23008	24800.33	30508.02
A_23_P43	J04536	Sialophorin	SPN	1.23006	0.00032	1.23006	1622.117	1989.172
A_24_P30	NM_00021	Janus kina	JAK3,JAKL	1.23003	0.0003	1.23003	7995.535	9793.949
A_23_P42	AB011133	Microtubul	MAST3	1.23	0.00057	1.23	1569.99	1922.597
A_23_P338113		DJ377H14.	ENST0000	1.22927	0.00038	1.22927	5434.05	6667.71
A_23_P40	NM_02393	hypothetica	MGC2752,	1.22926	0.0001	1.22926	3131.272	3846.386
A_23_P20	NM_02454	hypothetica	C1orf166,F	1.22926	0.00006	1.22926	5113.495	6283.054
A_24_P15	AK027150	Homo sapi	FLJ23497	1.22896	0.00043	1.22896	1387.683	1694.822
A_23_P11	BC050088	Similar to	LOC13050	1.22894	0.00042	1.22894	2873.545	3534.234
A_24_P60680		Olfactory r	ENST0000	1.22891	0.00013	1.22891	2225.853	2729.205
A_24_P28	XM_06363	similar to 6	LOC12339	1.22878	0.00007	1.22878	18185.62	22301.73
A_23_P83	NM_00273	Homo sapi	PRKAR1B,	1.22878	0.00032	1.22878	1232.165	1515.128
A_23_P11	NM_14462	hypothetica	MGC1729	1.22872	0.00074	1.22872	987.0443	1212.969
A_23_P22	NM_01771	hypothetica	PGPEP1,P	1.22814	0.00086	1.22814	1410.913	1739.707
A_23_P32	AF116619	Homo sapi	PRO1051	1.22811	0.00045	1.22811	2897.995	3556.427

A_32_P17	BC032064	Hypothetic	LOC38828	1.22808	0.00047	1.22808	3185.698	3931.48
A_23_P27	NM_01833	Homo sapi	ZNF444,E	1.22781	9.33E-07	1.22781	11261.88	13832.22
A_23_P31	NM_14719	Homo sapi	GLIS1,FLJ	1.22775	0.00095	1.22775	3253.99	3974.388
A_24_P41	NM_01753	Homo sapi	DNAH3,DL	1.22774	0.00052	1.22774	1938.507	2369.152
A_24_P178654		Unknown	A_24_P17	1.22767	0.00001	1.22767	19426.93	23887.56
A_32_P53603		Unknown	THC18273	1.22756	1.23E-06	1.22756	5268.135	6464.595
A_23_P43	NM_17358	Hypothetic	MGC4584	1.22728	3.46E-06	1.22728	20762.39	25471.25
A_23_P17	NM_00223	Homo sapi	CD82,R2,4	1.22708	0.00003	1.22708	2186.183	2684.433
A_23_P68	NM_02110	thymosin,	TMSB10,M	1.22705	0.00002	1.22705	20298.32	24925.03
A_23_P12	NM_02065	Homo sapi	PELI1,pelli	1.22688	0.00068	1.22688	1527.952	1869.564
A_23_P28	NM_03262	Homo sapi	GTPBP3,M	1.22687	0.00083	1.22687	1023.568	1256.124
A_23_P10	NM_17225	Homo sapi	MRPL54	1.22674	0.00018	1.22674	5177.351	6337.714
A_24_P22	AK096634	Tara-like p	HRIHFB21	1.2266	0.00019	1.2266	2909.891	3542.347
A_23_P40	NM_01331	Homo sapi	YPEL1,FK	1.22643	0.00034	1.22643	2077.713	2538.335
A_24_P384369		Unknown	ENST0000	1.22635	4.51E-06	1.22635	83812.21	102873.1
A_24_P62	AF007191	SIB 276 int	MUC3	1.22617	0.00014	1.22617	2032.878	2489.288
A_23_P99	NM_00033	Homo sapi	SLC12A1,E	1.22593	0.00005	1.22593	2425.709	2971.142
A_23_P26	NM_00684	Homo sapi	SDS,SDH	1.22576	0.00037	1.22576	2033.495	2483.578
A_24_P338151		Unknown	A_24_P33	1.22568	0.0007	1.22568	2393.278	2919.501
A_32_P45153		Unknown	THC19699	1.22546	0.00024	1.22546	6631.5	8108.732
A_24_P24002		Unknown	A_24_P24	1.22532	0.0002	1.22532	1531.726	1878.197
A_24_P18	NM_00242	matrix met	MMP19,M	1.22525	4.46E-06	1.22525	3301.035	4042.56
A_32_P124538		Unknown	THC18725	1.22519	0.00033	1.22519	1330.451	1633.327
A_23_P44	NM_00132	Homo sapi	CSRP2,CF	1.2251	0.00001	1.2251	2920.883	3578.621
A_23_P51	NM_01222	mutY (E. c	MUTYH,M	1.22493	0.00003	1.22493	3154.29	3864.294
A_24_P31	XM_17153	Homo sapi	MRGPRE,I	1.22462	0.0001	1.22462	3819.979	4690.526
A_24_P238819		Unknown	A_24_P23	1.22457	0.0002	1.22457	2834.779	3462.428
A_32_P143596		Unknown	A_32_P14	1.22456	1.27E-06	1.22456	5177.54	6340.306
A_24_P66	XM_37231	similar to R	LOC38995	1.22447	0.00088	1.22447	1469.276	1805.54
A_24_P510502		Unknown	A_24_P51	1.22418	0.00054	1.22418	7022.06	8570.775
A_23_P27	NM_03248	Homo sapi	CNFN,PLA	1.22408	0.00078	1.22408	18827.54	23084.49
A_23_P30	NM_01817	Nedd4 binc	N4BP2,B3I	1.22408	0.00019	1.22408	2018.485	2475.982
A_23_P12	NM_00135	Homo sapi	DAXX,DAF	1.22396	0.00006	1.22396	2016.725	2471.906
A_24_P33	BX640978	Pecanex-lil	PCNXL3	1.22391	0.00029	1.22391	1630.245	1990.926
A_23_P93	AK098134	Similar to (	AK098134	1.22376	0.0001	1.22376	4818.321	5895.141
A_24_P28	BC053367	Opposite s	GATS	1.22374	0.00067	1.22374	1370.891	1684.125
A_32_P98	NM_00442	dual specif	DUSP8,HE	1.22355	0.00044	1.22355	2863.536	3497.861
A_24_P332441		Unknown	A_24_P33	1.22344	0.00003	1.22344	41304.72	50626.79
A_24_P22	NM_14472	Homo sapi	ZMAT2,FL	1.22327	0.00014	1.22327	1955.459	2392.014
A_23_P32	NM_01212	Cip1-intera	CIZ1,LSFR	1.22282	0.00093	1.22282	1514.875	1858.574
A_32_P14	BQ027568	Apolipopro	APOA2	1.22258	0.00018	1.22258	3441.734	4193.481
A_23_P65	NM_00569	COX17 (ye	COX17,MC	1.22241	7.26E-06	1.22241	24218.82	29628.2
A_32_P182246		Unknown	A_32_P18	1.22131	0.00002	1.22131	3640.627	4443.629
A_24_P332780		Unknown	A_24_P33	1.22081	0.00002	1.22081	4031.898	4924.883
A_23_P95499		RL1X_MO	THC18444	1.22062	0.00001	1.22062	30560.92	37353.02
A_23_P41	NM_02257	putative le	GPR135,H	1.22062	0.00053	1.22062	3146.758	3828.343
A_24_P17	AK092577	CDNA FLJ	FLJ35258	1.2206	0.00077	1.2206	2470.542	2997.642
A_32_P165993		Unknown	A_32_P16	1.2202	0.00061	1.2202	1548.798	1894.36
A_24_P25	NM_00301	Surfactant,	SFTPC	1.22018	0.00065	1.22018	4400.663	5364.358
A_24_P93	AF007193	Mucin (MU	MUC3	1.21991	0.00011	1.21991	1441.416	1759.802
A_24_P41	NM_03145	hypothetic	MGC2560	1.21984	0.00021	1.21984	2974.455	3622.15
A_24_P27	NM_02245	hypothetic	FBS1,FBS	1.21951	0.00067	1.21951	12168.15	14807.5

A_23_P76	NM_01397	Homo sapi	BNIP1,NIP	1.21948	0.00049	1.21948	1308.719	1594.031
A_23_P92	AB051544	Hypothetic	FLJ10707	1.21914	0.00001	1.21914	4219.63	5143.582
A_24_P499481		Unknown	A_24_P49	1.21905	0.00052	1.21905	16465.72	20072.81
A_24_P34	NM_15269	hypothetic	FLJ38377	1.21884	0.00028	1.21884	2943.431	3579.769
A_23_P10	NM_00227	Homo sapi	KRTHA1,H	1.21847	0.00002	1.21847	5028.088	6117.242
A_32_P199083		Unknown	THC18483	1.21842	0.00036	1.21842	1444.166	1757.18
A_24_P42071		Unknown	ENST0000	1.21781	0.00013	1.21781	13460.53	16439.38
A_24_P15	NM_01910	tenascin XI	TNXB,XB,T	1.21777	0.00065	1.21777	2672.318	3251.913
A_32_P18	BC044608	Hypothetic	LOC44124	1.21759	0.00001	1.21759	2453.257	2986.533
A_23_P47	NM_01663	Homo sapi	ARL6IP4,S	1.21742	0.00028	1.21742	12328.41	14982.55
A_23_P20	NM_00552	Homo sapi	DNAJC4,H	1.21741	0.00073	1.21741	3646.311	4435.15
A_24_P83	NM_01492	NACHT, le	NALP1	1.21721	8.10E-06	1.21721	2835.394	3450.595
A_24_P45	AY007126	Hypothetic	LOC33922	1.21716	2.32E-06	1.21716	5863.378	7136.436
A_24_P187365		Unknown	A_24_P18	1.21705	0.00018	1.21705	5991.403	7271.424
A_23_P69	NM_00206	Homo sapi	GLRX,GRX	1.21701	1.69E-06	1.21701	8631.264	10504.96
A_23_P37	NM_15278	Hypothetic	MGC25181	1.21692	0.00006	1.21692	3296.558	4011.942
A_23_P42	NM_00289	Homo sapi	RBMS2,SC	1.2169	0.00023	1.2169	2017.259	2459.369
A_24_P84	AL122109	Hypothetic	LOC90835	1.21684	0.00031	1.21684	2157.386	2621.537
A_32_P79719		Unknown	A_32_P79	1.21682	0.00014	1.21682	9236.677	11254.1
A_23_P30	NM_00541	v-src avian	SRC,ASV,S	1.21659	7.52E-06	1.21659	4900.624	5956.808
A_24_P17	NM_00337	vascular er	VEGF,VPF	1.21653	0.0006	1.21653	1367.703	1659.916
A_23_P71	NM_05709	Homo sapi	CYP3A43	1.21651	0.00028	1.21651	4831.154	5869.816
A_23_P10	NM_00027	Homo sapi	NPHP1,NP	1.2159	0.00004	1.2159	2286.332	2779.024
A_32_P216809		AF054182	THC19530	1.21583	0.00023	1.21583	1983.99	2409.101
A_24_P93	AY358246	Chromosor	C6orf12	1.2158	0.00047	1.2158	6830.933	8297.381
A_23_P12	NM_02301	Homo sapi	C2orf26	1.21563	0.00053	1.21563	1813.677	2203.745
A_24_P19	NM_01533	Homo sapi	ASXL1,KIA	1.2156	0.0007	1.2156	2202.902	2668.672
A_23_P45	NM_00662	FUS-intera	FUSIP1,NS	1.21558	0.00013	1.21558	1891.02	2298.746
A_23_P13	NM_17586	Homo sapi	MAGEA6,M	1.21543	2.94E-09	1.21543	1323.114	1608.407
A_32_P111098		Unknown	A_32_P11	1.21503	0.00008	1.21503	2586.099	3137.096
A_32_P81	NM_18177	Homo sapi	SLC36A3,F	1.21499	0.00043	1.21499	1875.009	2274.104
A_24_P110082		Unknown	A_24_P11	1.2146	0.0004	1.2146	21886	26491.87
A_23_P38	NM_00375	Homo sapi	EIF3S4,EIF	1.21451	0.00066	1.21451	14877.63	18111.95
A_24_P22	NM_00302	SH3-doma	SH3BP2,C	1.21448	0.00009	1.21448	10235.13	12416.11
A_24_P32	NM_00551	Major histo	HLA-E	1.21441	4.56E-06	1.21441	21523.11	26135.89
A_23_P25	NM_15238	Homo sapi	BBS5	1.21425	0.00047	1.21425	10989.85	13380.87
A_23_P21	NM_02522	Homo sapi	ADAM33,F	1.21416	0.00011	1.21416	2967.393	3609.106
A_23_P20	NM_01606	mitochondr	Magmas,C	1.2139	9.22E-06	1.2139	5453.825	6619.986
A_24_P34	NM_13834	hypothetic	KIAA2013,	1.21378	0.00031	1.21378	6676.783	8094.974
A_24_P25	NM_00213	heterogene	HNRPA1,H	1.21377	0.00027	1.21377	11029.2	13349.3
A_23_P34	NM_01587	Homo sapi	ZNF593,ZT	1.21337	0.00017	1.21337	7640.755	9253.601
A_23_P61	NM_02481	Homo sapi	RABEP2,F	1.2132	0.00003	1.2132	2101.79	2550.042
A_23_P26	NM_03225	Homo sapi	WDR24,JF	1.21306	0.00002	1.21306	2236.822	2713.117
A_23_P37	NM_00348	solute carri	SLC7A5,E	1.21291	0.00015	1.21291	2087.157	2527.964
A_24_P587993		Unknown	A_24_P58	1.2128	0.00064	1.2128	3375.881	4100.34
A_24_P333052		Unknown	A_24_P33	1.2127	0.00008	1.2127	6385.031	7750.046
A_32_P22	AK055783	Hepatocyte	HNF4A	1.21249	0.00012	1.21249	3782.365	4588.179
A_32_P34	NM_02021	hypothetic	LOC56964	1.21248	0.00006	1.21248	4118.863	4989.714
A_32_P13	BX110985	Transcribe	BX110985	1.21242	0.00001	1.21242	2452.501	2972.97
A_32_P184304		Unknown	A_32_P18	1.21222	0.00015	1.21222	1593.686	1931.198
A_23_P16	NM_03147	hypothetic	FAM57B,F	1.21215	0.00013	1.21215	1401.933	1695.595
A_23_P213468		Unknown	A_23_P21	1.21212	0.00003	1.21212	5317.18	6449.405

A_24_P65	BC031069	Similar to F	MGC3369	1.21202	0.00097	1.21202	4759.728	5743.517
A_32_P42	NM_17827	eEF1A2 bi	DKFZp434	1.21197	0.00008	1.21197	2720.423	3298.89
A_23_P36	NM_17856	Homo sapi	NM_17856	1.21194	0.00003	1.21194	1966.398	2380.304
A_23_P80	NM_00622	Homo sapi	PLCD1	1.21171	0.00005	1.21171	2473.87	2999.07
A_24_P33	NM_00091	oxytocin re	OXTR,OT-	1.21162	0.00088	1.21162	3009.11	3665.058
A_23_P10	NM_00030	Homo sapi	PMP22,DS	1.21135	0.00067	1.21135	3603.538	4367.404
A_23_P10	BC031654	Family with	FAM53B	1.21121	0.00011	1.21121	3034.206	3669.662
A_24_P34	NM_01496	OGT(O-Gl	OIP106	1.2112	0.00039	1.2112	1916.325	2316.382
A_23_P12	XM_29185	Olfactory r	OR5F1	1.2108	0.00044	1.2108	2737.585	3284.29
A_23_P89	NM_02275	Homo sapi	ZNF574,FL	1.21072	0.00084	1.21072	8463.816	10205.45
A_24_P41	NM_08066	similar to R	MGC1960	1.21061	0.00014	1.21061	4914.923	5950.111
A_24_P85	AB051538	KIAA1751	KIAA1751	1.21038	0.00003	1.21038	2970.792	3596.219
A_24_P34	NM_00430	activin A re	ACVR1B,A	1.21034	0.00005	1.21034	6145.939	7485.882
A_23_P10	NM_17740	Homo sapi	SYT2,SytII	1.21007	0.00006	1.21007	1962.707	2375.349
A_24_P736617		Unknown	A_24_P73	1.20989	0.00008	1.20989	76717.31	93024.25
A_24_P17	NM_14719	Ankyrin re	FLJ35740	1.20976	0.00034	1.20976	1977.507	2386.286
A_32_P78	NM_00099	ribosomal	RPL37A,M	1.20956	0.00003	1.20956	221124.2	267821.6
A_23_P83	NM_01417	Homo sapi	PHPT1,PH	1.20946	0.00007	1.20946	23467.52	28440.59
A_32_P14	AK057319	CDNA FLJ	AK057319	1.20906	0.00042	1.20906	1505.315	1816.895
A_23_P13	NM_00537	Homo sapi	MPL,MPL	1.20902	0.00073	1.20902	3694.895	4452.864
A_24_P86	NM_00073	CD3E anti	CD3E,T3E	1.20892	0.0004	1.20892	3819.547	4634.662
A_23_P12	BC002872	KIAA0515	KIAA0515	1.20889	0.00004	1.20889	2150.999	2598.469
A_23_P50	NM_00275	Homo sapi	MAPK11,P	1.20887	0.0002	1.20887	1860.534	2248.13
A_24_P332461		Unknown	A_24_P33	1.2088	0.00005	1.2088	4186.293	5060.853
A_24_P12	NM_00230	leukemia i	LIF,CDF,H	1.2087	0.00095	1.2087	2493.428	3007.5
A_24_P922624		AF106680	THC19347	1.20845	0.00057	1.20845	1543.763	1868.972
A_23_P40	NM_01232	Homo sapi	MAPRE1,E	1.20843	0.00013	1.20843	2045.556	2471.876
A_23_P86	NM_02454	Homo sapi	C10orf76,F	1.20833	0.00016	1.20833	4107.627	5001.624
A_24_P872838		Unknown	A_24_P87	1.20789	0.00034	1.20789	1984.53	2386.763
A_24_P91	Z34282	Mucin 5, s	MUC5AC	1.20786	0.00064	1.20786	3813.876	4566.577
A_24_P20	NM_02498	hypothetic	FLJ12331	1.20755	0.00008	1.20755	53123.37	64111.84
A_32_P21	BC012946	Rho-guani	RGNEF	1.20755	0.00084	1.20755	1991.316	2399.821
A_23_P11	NM_00140	ephrin-A2	EFNA2,EL	1.20754	0.00069	1.20754	1618.703	1960.479
A_23_P31	AK024335	Intracellula	IPLA2(GA	1.20647	0.00028	1.20647	1706.237	2058.084
A_32_P40229		Unknown	THC19630	1.20611	0.00002	1.20611	3408.496	4112.155
A_23_P72	NM_17817	high densit	LOC33832	1.20592	0.00082	1.20592	2566.997	3104.589
A_24_P47	BX118521	BX118521	BX118521	1.20567	0.00038	1.20567	10216.14	12320.85
A_24_P102043		Unknown	ENST0000	1.2056	0.00005	1.2056	9685.098	11663.74
A_24_P393141		RL1X_HU	THC18124	1.20553	0.00029	1.20553	22799.21	27575.52
A_23_P32	NM_17755	Homo sapi	GPR109A,	1.20543	0.00044	1.20543	2724.226	3291.523
A_24_P45	NM_05294	guanylate	GBP4,Mpa	1.2051	0.00098	1.2051	4361.949	5246.779
A_23_P75	NM_01900	Homo sapi	TOLLIP,IL-	1.20457	0.00041	1.20457	2599.829	3137.67
A_23_P42	NM_00529	Homo sapi	GPR31	1.20432	0.00011	1.20432	1959.483	2361.649
A_23_P21	NM_00236	Homo sapi	MAGEB1,D	1.20427	0.00091	1.20427	1775.308	2134.273
A_32_P24	BC012486	Homo sapi	KRTAP2-1	1.20422	0.00001	1.20422	3148.763	3792.188
A_32_P13	AI693937	Similar to C	AI693937	1.20414	0.00005	1.20414	3526.608	4243.675
A_24_P811014		AF481879	THC18073	1.20408	0.0002	1.20408	3730.654	4466.805
A_24_P21	NM_14500	Homo sapi	YPEL4,FL	1.20405	0.00027	1.20405	1454.275	1750.807
A_24_P34	NM_00212	Major histo	HLA-DRB1	1.20399	0.00068	1.20399	4345.128	5243.352
A_23_P13	NM_02500	Thrombos	TSRC1	1.20337	0.00002	1.20337	3458.407	4163.893
A_32_P214340		AF271261	THC19223	1.20334	0.00004	1.20334	4067.38	4896.371
A_23_P10	NM_02402	Homo sapi	UNKL,ZC3	1.20331	0.00075	1.20331	1829.9	2199.966

A_23_P32	NM_18257	Homo sapi	ZSCAN1,M	1.20308	0.00001	1.20308	4026.095	4843.817
A_23_P27	NM_02431	Homo sapi	LILRP2,CD	1.20305	0.00039	1.20305	1546.652	1859.338
A_32_P96	NM_00329	tumor prote	TPT1,HRF	1.20302	0.00001	1.20302	109324.1	131533.1
A_23_P143676		Unknown	A_23_P14	1.20298	0.00021	1.20298	2307.543	2773.429
A_32_P19	NM_00098	ribosomal p	RPL18A	1.20282	0.00002	1.20282	52405.26	63079.02
A_23_P11	NM_00145	Homo sapi	FOXF1,FK	1.20221	0.00084	1.20221	1974.444	2365.083
A_24_P26	XM_06142	similar to S	LOC11935	1.2017	0.00014	1.2017	38674.38	46403.38
A_24_P34	NM_00686	leukocyte i	LILRA1,LIF	1.20169	0.0001	1.20169	3457.371	4151.177
A_24_P32	BC067760	Mitogen-ac	LOC40783	1.20134	0.00047	1.20134	1569.463	1886.369
A_24_P41	NM_08066	similar to R	MGC19604	1.20101	0.00019	1.20101	9172.719	11032.62
A_32_P38782		Unknown	THC19271	1.20099	0.00009	1.20099	3977.072	4787.13
A_32_P22	AW974787	Chromosor	C2orf10	1.20089	0.00013	1.20089	4339.943	5207.583
A_23_P87	BC025767	Rhophilin,	RHPN1	1.20077	0.0007	1.20077	8346.949	10025.66
A_24_P21	NM_00099	ribosomal p	RPL31,MG	1.20065	0.00073	1.20065	394613.6	475553.3
A_24_P10	AB088847	BPA-1 mR	BPA-1	1.20036	0.0004	1.20036	1690.016	2029.771
A_23_P58	NM_00240	mannosyl (	MGAT1,MC	1.20022	0.00077	1.20022	8006.778	9595.405
A_32_P15	AL713776	Hypothetic	DKFZp434	1.2001	0.00036	1.2001	1613.143	1935.367
A_23_P70	NM_00427	Homo sapi	LY86,MD-1	1.20004	0.00004	1.20004	3946.269	4740.097
A_24_P15	AB058778	Hypothetic	KIAA1875	1.19985	0.00094	1.19985	4403.302	5266.142
A_23_P39	NM_17590	hypothetica	LOC28429	1.19974	0.0002	1.19974	1665.921	1998.788
A_24_P10	NM_01228	potassium	KCNE1L,K	1.19936	0.00022	1.19936	1954.522	2346.582
A_24_P21	U05597	Solute carr	SLC4A3	1.19935	0.0009	1.19935	2104.115	2528.518
A_24_P74	NM_17754	Homo sapi	SNRPD2,S	1.19904	0.00014	1.19904	43976.89	52640.88
A_23_P27	NM_20732	hypothetica	LOC14765	1.1985	0.00016	1.1985	4203.118	5016.287
A_23_P13	BC021178	Chromosor	C20orf141	1.19848	0.00098	1.19848	7504.002	8974.543
A_32_P94	BC042589	Homo sapi	BC042589	1.19835	0.0002	1.19835	4039.333	4834.666
A_24_P19	NM_00482	potassium	KCNK6,TC	1.19824	0.00061	1.19824	3402.282	4084.795
A_24_P581618		Unknown	A_24_P58	1.19783	0.00023	1.19783	2349.326	2817.212
A_32_P105217		AF280545	THC18797	1.19783	0.00006	1.19783	2778.931	3327.688
A_23_P90	NM_00098	Homo sapi	RPL18A	1.19763	0.00003	1.19763	70347.15	84192.66
A_32_P751535		Z49148 rib	THC18781	1.1976	0.00003	1.1976	14829.47	17765.22
A_32_P10	BG191459	Transcribe	BG191459	1.19747	0.00052	1.19747	2962.569	3575.212
A_23_P43	NM_05819	Homo sapi	CDKN2A,A	1.19686	0.00016	1.19686	7503.443	8966.274
A_23_P41	NM_13896	Homo sapi	GPR73,ZA	1.19671	0.00022	1.19671	2770.645	3321.458
A_24_P93	BC009009	Phospholip	PLCB2	1.1966	0.00011	1.1966	4085.81	4886.134
A_23_P13	NM_00245	Homo sapi	NUDT1,MT	1.19649	0.00005	1.19649	7072.715	8457.115
A_23_P75192		Unknown	A_23_P75	1.19618	0.00026	1.19618	3856.613	4620.934
A_24_P22	AK094477	CDNA FLJ	AK094477	1.19583	0.00096	1.19583	8273.918	9787.938
A_24_P418019		Unknown	A_24_P41	1.19579	0.00052	1.19579	15490.11	18472.55
A_24_P20	NM_00333	ubiquitin-co	UBE2B,HR	1.19565	0.00012	1.19565	2713.012	3243.871
A_24_P29	NM_01432	nesca prote	RUSC1,NE	1.19561	0.00009	1.19561	4329.573	5177.521
A_23_P425492		B53A_HUM	THC18559	1.19555	0.00004	1.19555	3867.521	4623.188
A_23_P31	NM_17835	Homo sapi	LCE4A,LEI	1.19551	0.00008	1.19551	3700.916	4422.131
A_24_P28	NM_00059	CD14 antig	CD14	1.19548	0.00038	1.19548	3573.032	4265.983
A_23_P384857		DJ408B20.	ENST0000	1.1954	0.00017	1.1954	4500.104	5382.612
A_24_P39	AK093583	Hemicentin	DKFZp434	1.19524	0.00019	1.19524	4176.359	4990.164
A_24_P41	NM_01827	hypothetica	FLJ10922	1.19448	0.00017	1.19448	4180.156	5020.29
A_24_P50157		Unknown	A_24_P50	1.1938	0.00002	1.1938	82272.52	98230.32
A_24_P281304		Unknown	A_24_P28	1.19333	0.00017	1.19333	25552.16	30446.93
A_23_P25	NM_00099	Homo sapi	RPL29,HIF	1.19294	0.00005	1.19294	35990.39	42903.7
A_32_P69386		Unknown	ENST0000	1.19284	0.00054	1.19284	1787.459	2128.671
A_24_P65	AF147791	Mucin 11	MUC11	1.19274	0.00055	1.19274	2723.748	3245.323

A_23_P81	NM_00390	sequestosc	SQSTM1,p	1.19247	0.00002	1.19247	78546.64	93643.23
A_23_P21	AK075554	Elastin (su	ELN	1.19246	0.00017	1.19246	2316.099	2763.657
A_23_P41	NM_00416	Homo sapi	PYY,PYY1	1.19219	0.00027	1.19219	1794.052	2139.312
A_32_P61884		Unknown	A_32_P61	1.19218	0.00004	1.19218	4109.141	4898.725
A_24_P19	XM_07101	similar to b	LOC13865	1.1919	0.0008	1.1919	1416.981	1688.787
A_23_P39277		Unknown	A_23_P39	1.19143	0.00005	1.19143	7405.029	8816.445
A_24_P110601		Unknown	A_24_P11	1.19115	0.00012	1.19115	10173.76	12113.59
A_24_P31	NM_03278	hypothetica	ZC3H10,FI	1.191	0.00019	1.191	1930.14	2297.485
A_24_P53	AL080233	FLJ16124	FLJ16124	1.19076	0.00004	1.19076	4960.164	5903.561
A_23_P50	NM_00363	potassium	KCNAB2,A	1.19072	0.00023	1.19072	2508.285	2984.364
A_24_P11	XM_37185	Homo sapi	XM_37185	1.19032	0.00014	1.19032	53063.62	63212.09
A_23_P75	NM_00397	Homo sapi	AIP,ARA9,	1.19013	0.00037	1.19013	6778.368	8067.852
A_24_P75	NM_02526	G5C protei	LY6G5C,G	1.18984	0.00066	1.18984	5410.82	6432.73
A_23_P47	XM_37066	Homo sapi	OR8G3P	1.18981	0.00083	1.18981	1818.044	2159.199
A_24_P15	AK097703	Hypothetic	LOC92482	1.18975	0.00011	1.18975	2610.108	3100.633
A_32_P52	NM_15370	receptor tra	RTP1,MGC	1.18947	0.00017	1.18947	2534.095	3014.565
A_24_P36	NM_19854	Homo sapi	MGC35434	1.18932	0.00014	1.18932	2713.732	3229.05
A_32_P40744		Unknown	A_32_P40	1.18919	0.00076	1.18919	2839.87	3369.725
A_23_P82461		I46880 T-c	THC19741	1.18912	0.00006	1.18912	4787.151	5684.726
A_24_P26	NM_15375	Homo sapi	C21orf88	1.18897	0.00027	1.18897	5881.412	6990.166
A_24_P22	BC042161	Trafficking	TRAPPC5	1.18882	0.00023	1.18882	9210.833	10953.77
A_24_P24	NM_00577	small EDR	SERF2,FA	1.18734	0.00016	1.18734	73147.17	86931.69
A_23_P18	NM_00577	Homo sapi	RBM5,G15	1.18705	0.00005	1.18705	5032.437	5973.899
A_23_P97	NM_00389	Homo sapi	LDB1,NLI,(	1.18687	0.0009	1.18687	2423.247	2871.161
A_32_P193079		Unknown	A_32_P19	1.18641	0.00002	1.18641	39544.8	46913.48
A_24_P24	NM_18160	Homo sapi	KRTAP13-	1.18549	0.00035	1.18549	4938.319	5848.538
A_24_P52	AF117899	Fucosyltrar	FUT1	1.18539	0.00006	1.18539	4877.479	5783.414
A_32_P52	NM_17857	Homo sapi	RTN4RL2,	1.1852	0.00041	1.1852	7990.646	9443.271
A_32_P11	BC039095	Homo sapi	BC039095	1.18507	0.00005	1.18507	6736.434	7983.934
A_24_P35	NM_00098	ribosomal	RPL21,L21	1.1849	0.00013	1.1849	28906.66	34285.19
A_24_P204165		Unknown	A_24_P20	1.18401	0.00009	1.18401	10582.42	12532.17
A_24_P790909		Unknown	A_24_P79	1.18391	0.00076	1.18391	3689.389	4371.863
A_23_P21	AF028825	Discs, larg	DLG4	1.18379	0.00043	1.18379	3277.656	3899.473
A_23_P51	NM_00328	Homo sapi	TNNI1,TNN	1.18376	0.00046	1.18376	3015.709	3566.06
A_23_P42	NM_00619	Homo sapi	PAX9	1.18321	0.00082	1.18321	2743.48	3239.519
A_32_P21	AK094447	LOC44210	AK094447	1.18267	0.00055	1.18267	2030.025	2408.084
A_23_P35	NM_05292	Homo sapi	PNMA5,KI	1.18244	0.00084	1.18244	8998.393	10623.46
A_32_P3408		Unknown	THC19697	1.18138	0.00073	1.18138	3312.508	3894.178
A_23_P81	NM_00183	Homo sapi	CLPS	1.18128	0.00041	1.18128	2718.567	3214.686
A_24_P409521		Unknown	ENST0000	1.18106	0.00017	1.18106	4308.308	5087.29
A_32_P49	BX111592	Transcribe	BX111592	1.17966	0.0001	1.17966	7506.152	8854.043
A_24_P629163		Unknown	A_24_P62	1.17943	0.00038	1.17943	2210.635	2608.409
A_24_P932046		Unknown	THC19079	1.17873	0.00073	1.17873	3410.104	4014.13
A_23_P13	NM_00414	Homo sapi	LLGL1,DL(	1.17847	0.00066	1.17847	2253.852	2654.194
A_23_P44	NM_15326	Homo sapi	PLCXD2,F	1.1784	0.00015	1.1784	5315.279	6267.441
A_32_P20	AV759013	Transcribe	RPL21,L21	1.17778	0.00032	1.17778	42856.29	50537.01
A_32_P88	NM_01331	hypothetica	YPEL1,FK	1.17722	0.00098	1.17722	1716.459	2020.897
A_23_P41	NM_01239	Submaxilla	PROL5	1.17692	0.00024	1.17692	4508.83	5312.46
A_32_P21	AV759013	Transcribe	RPL21,L21	1.17689	0.00029	1.17689	41413.51	48796.28
A_23_P111766		Unknown	A_23_P11	1.17671	0.00043	1.17671	3541.166	4165.192
A_24_P24142		Unknown	A_24_P24	1.17607	0.00068	1.17607	65144.7	76503.34
A_32_P75888		AF078857	THC19577	1.17581	0.00078	1.17581	7327.17	8602.105

A_23_P77	AB007945	KIAA0476	KIAA0476	1.17413	0.00073	1.17413	4775.609	5606.446
A_23_P10	NM_20711	Homo sapi	C14orf68,H	1.17396	0.00055	1.17396	2541.067	2985.213
A_23_P14	NM_03320	Homo sapi	ZNF101,HZ	1.1735	0.00082	1.1735	2388.525	2802.423
A_23_P15	NM_01798	hypothetica	GPR172B,	1.1731	0.00098	1.1731	2632.424	3091.155
A_23_P43	NM_03354	mitofusin 1	MFN1,hfzo	1.17192	0.00037	1.17192	2887.772	3382.662
A_24_P27	NM_01242	ribosomal p	RPL13A	1.17157	0.00012	1.17157	117705.2	137936
A_24_P10	NM_17488	Homo sapi	CRB3,MG	1.17149	0.00017	1.17149	4212.225	4933.523
A_23_P39	NM_00312	Spi-B trans	SPIB,SPI-E	1.17102	0.00063	1.17102	2797.872	3274.691
A_24_P85	BC008927	Zinc finger,	ZFYVE26	1.17065	0.00037	1.17065	2671.473	3124.566
A_24_P26	NM_14524	Homo sapi	FAM14B	1.16956	0.0006	1.16956	6107.706	7147.13
A_23_P27	NM_00300	Homo sapi	SEPW1,se	1.16897	0.00032	1.16897	7898.607	9234.822
A_32_P53	AF060511	Homo sapi	My016 prot	1.16893	0.00042	1.16893	55364.16	64654.73
A_23_P39	NM_01571	Homo sapi	GLTSCR2,	1.16637	0.00024	1.16637	59109.2	68913.69
A_32_P35	NM_00464	poly(A)-bin	PABPN1,C	1.16622	0.00081	1.16622	7388.411	8608.487
A_32_P16	AV759013	Transcribe	RPL21,L21	1.16575	0.00014	1.16575	61872.66	72128.92
A_23_P20	NM_00308	small nucle	SNAPC4,S	1.16443	0.00091	1.16443	5782.486	6739.717
A_23_P25	NM_01659	Homo sapi	FKBP11,FI	1.16442	0.00043	1.16442	6174.49	7187.991
A_23_P15	NM_00537	Homo sapi	MOS,MSV,	1.16371	0.00034	1.16371	8252.713	9605.89
A_23_P15	BC035682	Hemoglobi	HBZ	1.16296	0.00061	1.16296	2371.554	2760.809
A_24_P392082		Unknown	ENST0000	1.16291	0.00047	1.16291	21031.29	24445.79
A_32_P32	NM_03240	protocadhe	PCDHGC5	1.16272	0.00078	1.16272	3061.652	3558.203
A_23_P42	NM_00629	Homo sapi	TNFAIP2,E	1.16247	0.00026	1.16247	35952.44	41787.98
A_24_P383934		Unknown	A_24_P38	1.16169	0.00051	1.16169	4316.79	5016.337
A_23_P15	NM_02455	Homo sapi	LASS4,Trh	1.16162	0.00068	1.16162	1853.985	2154.087
A_32_P234996		ALU2_HUM	THC19909	1.1615	0.00027	1.1615	7730.991	8980.643
A_23_P50	NM_00440	Homo sapi	DMPK,DM,	1.16144	0.00068	1.16144	3253.76	3777.953
A_23_P12	BC036120	F-box and	FBXL15	1.16094	0.00081	1.16094	7685.577	8914.53
A_32_P16	NM_15364	Homo sapi	TPM3,TRK	1.16091	0.00064	1.16091	5597.835	6495.58
A_23_P51	NM_01528	KIAA0467	KIAA0467	1.16024	0.00078	1.16024	3475.282	4028.792
A_24_P169855		Unknown	A_24_P16	1.15941	0.00066	1.15941	35808.02	41485.14
A_32_P21	BX115137	Transcribe	BX115137	1.15928	0.00097	1.15928	3649.527	4225.268
A_23_P14	NM_00427	A kinase (P	AKAP6,AD	1.15925	0.00048	1.15925	3650.953	4232.67
A_32_P192586		Unknown	A_32_P19	1.15766	0.00079	1.15766	6436.17	7444.166
A_23_P22	NM_00394	Homo sapi	HAP1,HLP	1.15738	0.00053	1.15738	6938.127	8027.984
A_24_P12	NM_01568	hypothetica	C2orf24,C	1.15694	0.00094	1.15694	4395.119	5081.7
A_32_P11	AK026131	CDNA: FL	AK026131	1.15675	0.00034	1.15675	6534.94	7558.513
A_23_P33	NM_02391	Homo sapi	TAS2R9,TZ	1.15622	0.00094	1.15622	4136.949	4775.782
A_23_P71	NM_00254	Homo sapi	TNFRSF11	1.15362	0.00038	1.15362	664.5137	766.3854
A_24_P392271		Unknown	A_24_P39	1.15296	0.0008	1.15296	73379.73	84574.63
A_24_P12	NM_00248	NADH deh	NDUFA2,B	1.14828	0.00097	1.14828	12646.19	14516.64
A_24_P41	XM_37167	Homo sapi	XM_37167	1.14561	0.00091	1.14561	26765.22	30669.47
A_23_P25	NM_00715	Homo sapi	ZNF195,DI	1.14522	0.00006	1.14522	1835.921	2105.492
A_32_P10	NM_02274	hypothetica	SMYD3,ZM	-1.15295	0.00076	0.86734	18765.56	16283.06
A_23_P21	NM_01424	Homo sapi	RBX1,ROC	-1.15336	0.00067	0.86703	9917.566	8601.105
A_23_P50	NM_02110	Homo sapi	MRPS12,R	-1.15554	0.0008	0.86539	44440.67	38478.23
A_24_P20	NM_00164	N-acylamin	APEH,APH	-1.15673	0.00036	0.86451	9319.636	8056.993
A_32_P15	NM_14571	Homo sapi	ATXN2L,A	-1.15781	0.00029	0.8637	16519.6	14267.88
A_23_P94	NM_01231	Homo sapi	LETM1	-1.15879	0.00083	0.86297	3272.001	2823.61
A_23_P28	NM_00376	Homo sapi	VAMP8,ED	-1.1597	0.00075	0.86229	21903.47	18869.82
A_23_P13	NM_14572	Homo sapi	MRPL24,M	-1.16013	0.00091	0.86197	19487.3	16814.06
A_32_P20	XM_03464	similar to ri	LOC15834	-1.16451	0.00024	0.85873	23995.54	20610.98
A_24_P358337		Unknown	A_24_P35	-1.1648	0.00087	0.85851	10728.75	9199.705

A_23_P354	NM_01846	Homo sapi	C10orf70,M	-1.16654	0.00017	0.85724	13522.83	11590.95
A_24_P250	NM_00459	small nucle	SNRPA	-1.16743	0.00026	0.85658	8023.077	6872.771
A_23_P154	NM_00045	Superoxide	SOD1	-1.1694	0.00046	0.85514	142823.5	121983.8
A_32_P865	AF038185	Clone 2370	AF038185	-1.17017	0.00097	0.85458	2576.343	2200.952
A_32_P160	XM_37067	similar to h	LOC38783	-1.17058	0.00025	0.85428	4732.551	4043.996
A_23_P204	NM_05327	Homo sapi	RPLP0,P0,	-1.17115	0.0003	0.85386	258685	221065.3
A_23_P428	NM_03201	Homo sapi	MRPS24,H	-1.17158	0.00036	0.85355	64547.57	55140.66
A_23_P52	NM_01624	Homo sapi	CYB5R1,B	-1.17174	0.0009	0.85343	4486.688	3830.094
A_23_P85	NM_01323	CD24 antig	CD24,CD2	-1.17193	0.00072	0.85329	3287.289	2802.536
A_32_P180	NM_15364	Homo sapi	TPM3,TRK	-1.17199	0.00055	0.85325	6721.905	5737.769
A_24_P465	BC058822	Similar to 4	LOC20401	-1.17287	0.00032	0.85261	31497.38	26876.7
A_23_P80	NM_03283	Homo sapi	DIRC2,RC	-1.17456	0.00051	0.85139	2573.81	2191.345
A_23_P415	NM_01778	hypothetica	FLJ20364	-1.17503	0.00001	0.85104	2907.848	2488.238
A_23_P20	NM_01662	Homo sapi	WAC,Wwp	-1.1758	0.00093	0.85048	8757.214	7462.929
A_32_P75	BC034247	F-box prote	FBXO10	-1.17663	0.0002	0.84989	31412.94	26720.66
A_23_P13	NM_00675	Homo sapi	TCEA1,SII	-1.17664	0.00026	0.84988	8900.295	7555.333
A_23_P85	NM_00375	Homo sapi	EIF3S2,TR	-1.17726	0.00067	0.84943	19467.76	16572.93
A_23_P14	NM_01557	Homo sapi	TNRC15,G	-1.17816	0.00022	0.84878	3765.855	3196.317
A_23_P12	NM_00281	Homo sapi	PSMD4,AF	-1.17875	0.00082	0.84836	10679.6	9037.518
A_23_P16	NM_00393	Homo sapi	WASF1,W	-1.17938	0.0008	0.8479	3764.164	3193.469
A_24_P754	NM_00538	nucleolin	NCL,C23,F	-1.18034	0.00045	0.84722	12385.01	10510.02
A_23_P150	NM_03263	Homo sapi	SKP2,FBL	-1.18077	0.00034	0.84691	2600.469	2202.872
A_24_P400981		Unknown	A_24_P40	-1.18084	0.00039	0.84686	2702.671	2289.432
A_23_P904	NM_02432	hypothetica	MGC10433	-1.18204	0.00073	0.846	6433.274	5434.226
A_23_P92	NM_00676	interferon-r	IFRD2,SM	-1.18228	0.00005	0.84582	10325.77	8733.438
A_24_P32	AB051452	Polymeras	POLR3H	-1.18252	0.00068	0.84565	5410.823	4576.83
A_23_P148	NM_00313	Homo sapi	SRP14,AL	-1.18292	0.00027	0.84537	89317.52	75371.69
A_23_P875	NM_01418	Homo sapi	ORMDL2,M	-1.18332	0.00009	0.84508	13163.18	11130.02
A_23_P125	NM_00100	Homo sapi	RPS4X,CC	-1.18358	0.00011	0.84489	230624.5	195028.6
A_23_P20	NM_01649	HSPC244	MGC13379	-1.18424	0.00073	0.84442	2483.247	2095.834
A_23_P11	NM_00309	Homo sapi	SNRPF,SM	-1.18444	0.00035	0.84428	83748.03	70850.31
A_24_P630	AK022529	Hypothetic	LOC13007	-1.18532	0.00055	0.84365	2175.103	1835.245
A_23_P68	NM_08066	similar to R	MGC15407	-1.18538	0.00048	0.84361	6185.688	5226.997
A_23_P115	NM_00125	tumor necr	TNFSF7,C	-1.18555	0.00075	0.84349	4016.891	3389.793
A_23_P13	NM_17067	Homo sapi	SKP1A,OC	-1.18571	0.00019	0.84338	22414.8	18930.11
A_24_P67	NM_00144	fatty acid b	FABP5,EF	-1.18624	0.00006	0.843	19836.45	16722.86
A_23_P874	NM_02072	Homo sapi	FAM62B,C	-1.18698	0.00018	0.84247	3908.729	3291.815
A_32_P230	BC007205	Homo sapi	BC007205	-1.18803	0.00049	0.84173	3494.964	2941.983
A_23_P285	NM_00646	associated	STAMBP,A	-1.18867	0.00093	0.84127	2318.147	1951.679
A_32_P15	XM_37176	Homo sapi	XM_37176	-1.18906	0.00009	0.841	47833.06	40194.12
A_23_P698	NM_02297	small EDR	SERF1B,F	-1.18941	0.00068	0.84075	13076.27	11022.99
A_23_P11	NM_00484	Homo sapi	ATG5,ASP	-1.19015	0.00017	0.84023	2735.186	2298.283
A_24_P273284		Unknown	ENST0000	-1.19028	0.00016	0.84014	3195.59	2684.677
A_23_P218	NM_14888	Homo sapi	MRPL10,R	-1.19039	0.00004	0.84006	6979.01	5862.847
A_23_P984	NM_00019	Homo sapi	HMBS,UPS	-1.19126	0.00015	0.83945	21354.33	17934.79
A_23_P21	NM_00241	Homo sapi	CD99,MIC	-1.19127	0.00001	0.83944	19695.96	16533.52
A_23_P20	NM_00287	Homo sapi	RAD51C,R	-1.19135	0.0004	0.83938	5003.901	4195.077
A_23_P145	NM_00405	Bystin-like	BYSL	-1.1914	0.00053	0.83935	4436.828	3720.095
A_23_P410	AL122036	Three prim	TREX2	-1.19257	0.00043	0.83852	2653.006	2224.74
A_24_P294	NM_00346	protein tyr	PTP4A1,H	-1.19349	0.00002	0.83788	7776.021	6515.882
A_24_P152983		Unknown	A_24_P15	-1.19365	0.00064	0.83777	50971.36	42783.28
A_24_P194	NM_19882	Ras-relate	RAC1	-1.19415	0.00018	0.83742	6222.611	5213.736

A_24_P28	NM_08091	Homo sapi	DGUOK,dC	-1.19421	0.00011	0.83737	5098.389	4271.932
A_23_P88	NM_00074	Homo sapi	CHRNA5	-1.19451	0.0004	0.83716	4455.271	3727.144
A_23_P21	NM_00094	Homo sapi	PON3	-1.19474	0.00004	0.837	5427.214	4542.113
A_23_P27	NM_15246	Homo sapi	C17orf32,M	-1.19476	0.00023	0.83699	2405.22	2015.476
A_23_P33	BC013792	Homo sapi	SSH1,SSH	-1.19549	0.00037	0.83648	1915.001	1601.602
A_23_P90	NM_01823	dudulin 2	STEAP3,2,	-1.19587	0.00019	0.83621	2264.915	1894.306
A_23_P92	NM_00481	Homo sapi	TJP2,ZO2,	-1.19615	0.00034	0.83602	2357.271	1969.715
A_32_P12	NM_01323	px19-like p	PX19,PRE	-1.19617	0.00006	0.836	18994.63	15867.19
A_24_P23	BC000821	Hypothetic	LOC14960	-1.19636	0.00076	0.83587	5507.992	4610.53
A_23_P88	NM_01610	Homo sapi	SCFD1,RA	-1.19637	0.00069	0.83586	5853.381	4900.059
A_23_P13	NM_00300	Homo sapi	SDHD,PGL	-1.19673	0.00002	0.83561	8341.094	6969.649
A_23_P12	NM_00290	Homo sapi	RDBP,RD,	-1.19674	0.00011	0.8356	9307.047	7787.918
A_24_P10	NM_00639	nucleolar p	NOL5A,NC	-1.19721	0.00002	0.83527	52643.11	43987.92
A_32_P148824		Unknown	A_32_P14	-1.19738	0.001	0.83516	3052.999	2551.021
A_23_P20	NM_00092	Homo sapi	PDHB,PHE	-1.19744	0.00019	0.83511	6170.837	5151.349
A_23_P68	NM_08082	Homo sapi	C20orf108,	-1.19766	0.00007	0.83496	3345.399	2793.102
A_23_P14	NM_00300	Homo sapi	SDHB,IP,S	-1.19782	0.00005	0.83485	11780.67	9826.149
A_23_P10	NM_03247	Homo sapi	MRPS6,RF	-1.19803	0.00038	0.8347	26475.58	22150.06
A_23_P75	NM_00455	NADH deh	NDUFS3	-1.19808	0.00008	0.83467	9974.427	8314.306
A_23_P25	AK025749	Chromosor	C14orf35	-1.19839	0.0005	0.83445	2794.538	2327.897
A_23_P97	NM_01810	Homo sapi	LRRC8D,L	-1.1986	0.00039	0.83431	2505.824	2091.467
A_23_P50	NM_02406	hypothetica	MGC2749	-1.19885	0.00008	0.83413	14900.43	12442.22
A_23_P15	NM_00562	Homo sapi	SLC6A8,C	-1.19896	0.00047	0.83406	2856.677	2382.658
A_23_P20	NM_00473	Homo sapi	MTA2,PID,	-1.19911	0.00088	0.83395	1899.03	1585.942
A_23_P25	NM_00181	Homo sapi	CENPE,KIF	-1.19934	0.00005	0.83379	3373.294	2815.721
A_24_P127181		Unknown	A_24_P12	-1.19934	0.00001	0.83379	87027.13	72538.33
A_32_P21	XM_37910	similar to C	LOC40096	-1.19953	0.00024	0.83366	4564.06	3804.874
A_23_P16	NM_01241	Homo sapi	QPCT,QC,	-1.19977	0.00014	0.83349	4553.435	3794.551
A_23_P15	NM_01751	Homo sapi	ENOSF1,R	-1.20026	0.00095	0.83315	2308.392	1921.747
A_23_P50	NM_00014	Ferritin, ligl	FTL	-1.20101	0.00015	0.83263	419743.8	349109.6
A_24_P340976		Unknown	A_24_P34	-1.20101	0.00007	0.83263	51676.3	43055.63
A_32_P186821		Unknown	THC18595	-1.20101	0.00096	0.83263	3693.498	3077.149
A_23_P56	NM_02053	Homo sapi	RTN4,ASY	-1.20173	0.00015	0.83213	10483.37	8729.354
A_23_P96	NM_01612	Homo sapi	C1orf41,PF	-1.20217	0.00069	0.83183	7490.891	6244.029
A_24_P400751		Unknown	A_24_P40	-1.20255	0.00069	0.83157	17667.79	14733.46
A_32_P23	NM_00671	activated R	SUB1,P15,	-1.20267	0.00033	0.83148	33046.69	27553.82
A_23_P21	NM_00047	adenylate k	AK1	-1.20297	0.00017	0.83127	3252.547	2702.817
A_24_P30	NM_00053	propionyl C	PCCB,DKF	-1.20315	0.00077	0.83115	2082.21	1731.341
A_24_P31	NM_01475	mastermin	MAML1,Ma	-1.20395	0.00006	0.8306	6474.486	5369.526
A_24_P25	NM_00294	ribosomal p	RPL15,EC	-1.20414	0.00018	0.83047	148046.5	123249.6
A_23_P99	NM_00531	Homo sapi	PDIA3,P58	-1.20485	0.00004	0.82998	10214.62	8483.664
A_23_P14	NM_01772	hypothetica	FLJ20245	-1.20531	0.00062	0.82966	5773.426	4796.455
A_23_P17	NM_19825	Homo sapi	E2F6,E2F-	-1.20592	0.00024	0.82924	4150.837	3440.784
A_23_P65	NM_00281	Homo sapi	PSME2,PA	-1.2062	9.26E-06	0.82905	64539.35	53490.07
A_23_P15	NM_00039	ATP-bindin	ABCC2,DJ	-1.20661	0.00005	0.82877	6821.085	5655.27
A_32_P88	NM_03299	general tra	GTF2I,DIW	-1.20698	0.00063	0.82851	3052.893	2528.432
A_23_P92	NM_01610	Homo sapi	SARA2,CM	-1.20783	0.00017	0.82793	6151.446	5097.332
A_23_P35	NM_02236	popeye pro	POPDC3,F	-1.20795	0.0001	0.82785	3362.606	2783.377
A_24_P608790		UCRH_HU	THC18041	-1.20833	0.00012	0.82759	53567.56	44427.41
A_23_P13	NM_00336	Homo sapi	UQCRC1,D	-1.2085	0.00087	0.82747	12033.49	9942.789
A_23_P25	NM_08062	Homo sapi	BRI3BP,KC	-1.20853	0.00011	0.82745	8669.011	7184.803
A_24_P30	NM_02463	hypothetica	MAK10,FL	-1.20864	0.00039	0.82738	5435.626	4510.345

A_23_P67	NM_02507	Homo sapi	UXS1,FLJ2	-1.20877	0.00045	0.82729	2853.704	2358.743
A_23_P12	NM_01429	Homo sapi	TDRD7,KI	-1.20879	0.00068	0.82728	4513.311	3737.957
A_23_P14	NM_03148	Homo sapi	GRWD1,G	-1.20883	0.00013	0.82725	3254.427	2691.296
A_23_P38	NM_00194	Homo sapi	E2F3,E2F-	-1.20958	0.00026	0.82673	5188.646	4293.337
A_32_P24	AW847262	Transducin	TBL1XR1	-1.20989	0.00021	0.82652	2566.969	2124.555
A_23_P53	NM_00488	Homo sapi	TP53I3,PI	-1.20996	0.00034	0.82648	2440.313	2016.985
A_23_P15	NM_00097	Homo sapi	RPL6,TXR	-1.21044	0.00032	0.82614	133735	110150.7
A_23_P21	NM_01230	Homo sapi	FBXW11,H	-1.21051	0.00088	0.8261	6812.883	5628.331
A_24_P814444		Unknown	THC18375	-1.21112	0.00004	0.82569	2714.592	2241.105
A_23_P30	NM_14717	A kinase (F	AKAP9	-1.21114	0.00094	0.82567	1890.169	1558.828
A_23_P37	BC006267	Asparagina	ASRGL1	-1.21125	0.00093	0.82559	2082.86	1718.317
A_23_P14	NM_00580	Homo sapi	PRDX2,PR	-1.21144	0.00019	0.82546	8626.776	7132.4
A_23_P79	NM_03263	seven tran	NIFIE14,M	-1.21147	5.58E-06	0.82544	15473.25	12768.73
A_23_P49	NM_00125	Homo sapi	CDC6,CD	-1.2123	7.01E-06	0.82488	4977.779	4106.174
A_23_P11	NM_01422	Homo sapi	MTCP1,C6	-1.21253	0.00009	0.82472	3763.45	3103.21
A_23_P21	NM_01604	Homo sapi	C20orf45,d	-1.21289	0.00001	0.82448	4634.86	3821.84
A_24_P87	NM_14530	similar to C	FAM18B2,	-1.21324	0.00053	0.82424	2905.922	2395.721
A_23_P71	NM_00135	Homo sapi	DECR1,DE	-1.21376	1.81E-06	0.82389	15542.53	12805.38
A_24_P37	AL442077	Homo sapi	DKFZp667	-1.21407	0.00058	0.82367	1676.611	1379.948
A_23_P16	NM_00460	Homo sapi	TBCA	-1.21418	0.00001	0.8236	88257.33	72636.24
A_24_P13	NM_01496	KIAA0952	BTBD3,KI	-1.2149	0.00035	0.82311	2711.338	2233.315
A_24_P23	NM_00066	ribosomal p	RPL9,FLJ2	-1.21524	0.00046	0.82288	82116.28	67821.86
A_23_P54	NM_00169	Homo sapi	ATP6V0C,	-1.21552	0.00002	0.8227	54383.62	44791.8
A_23_P36	NM_00269	Homo sapi	POLR2G,R	-1.21555	0.00047	0.82268	13041.13	10733.25
A_24_P47	NM_01844	TBP-intera	CAND1,TI	-1.21561	0.00041	0.82263	7441.549	6123.882
A_24_P25	NM_01849	COBW-like	CBWD1	-1.2159	0.00001	0.82244	4106.596	3377.105
A_32_P18	NM_14580	Homo sapi	ZNF511,M	-1.2159	0.00083	0.82244	4974.024	4097.53
A_24_P272713		Unknown	A_24_P27	-1.21597	0.00086	0.82239	3624.542	2981.88
A_23_P10	NM_00387	Homo sapi	TMEM11,F	-1.21598	8.75E-06	0.82238	6594.498	5424.097
A_23_P13	NM_00417	Homo sapi	SNRPD3,S	-1.21612	0.00004	0.82229	32956.91	27153.35
A_24_P14	NM_00138	dihydropyri	DPYSL3,D	-1.21622	0.00077	0.82222	7761.363	6367.737
A_24_P15	NM_00500	NADH deh	NDUFB9,B	-1.21639	0.0007	0.82211	49859.86	40827.86
A_23_P10	NM_02107	Homo sapi	NMT1,NMT	-1.21642	0.00009	0.82209	5024.861	4133.834
A_24_P82	NM_01804	mago-nash	FLJ10292,	-1.21668	0.00098	0.82191	5112.827	4213.74
A_23_P16	NM_00037	Homo sapi	VDR,NR1I	-1.21682	0.00061	0.82181	4351.882	3583.377
A_23_P46	NM_01599	Homo sapi	ADIPOR1,I	-1.21686	0.00002	0.82179	4998.557	4102.745
A_24_P73	NM_00356	unc-51 (C.	ULK1,ATG	-1.21696	0.00024	0.82172	2278.168	1874.681
A_24_P212764		Unknown	A_24_P21	-1.217	8.97E-06	0.82169	7828.684	6431.811
A_23_P43	NM_00657	Homo sapi	DCTN6,WS	-1.21735	0.00031	0.82146	5224.974	4284.636
A_23_P15	NM_02410	hypothetica	MGC2747	-1.21741	0.00025	0.82142	2870.578	2355.897
A_23_P42	NM_00669	Homo sapi	CPSF4,NA	-1.21763	0.00004	0.82126	8264.877	6784.319
A_24_P28	NM_03291	hypothetica	FAM86B1,	-1.21764	0.00025	0.82126	4438.404	3650.595
A_23_P63	NM_03365	Homo sapi	DAP3,DAP	-1.21789	0.00043	0.82109	44592.85	36741.91
A_23_P79	NM_01323	px19-like p	PX19,PRE	-1.21796	5.61E-06	0.82104	29393.75	24151.03
A_23_P98	NM_01596	Homo sapi	CUTC,CGI	-1.21811	0.00025	0.82094	3611.538	2961.548
A_23_P13	NM_03317	Homo sapi	BAT4,G5,D	-1.21828	0.00026	0.82083	4358.956	3583.874
A_23_P43	NM_01642	Homo sapi	ZNF219,ZF	-1.21844	1.79E-06	0.82072	15003.54	12315.98
A_32_P4532		Unknown	A_32_P45	-1.21851	0.00055	0.82067	164675.9	135556.4
A_32_P21	NM_00321	thymine-D	TDG	-1.21864	0.00031	0.82059	9765.402	8024.075
A_23_P56	BC015995	General tra	GTF3C3	-1.21873	0.00015	0.82053	2819.441	2310.509
A_24_P17	NM_05718	Homo sapi	VPS29,DC	-1.21885	0.00054	0.82045	7959.997	6554.149
A_23_P31	NM_01543	Homo sapi	CCDC28A,	-1.21887	0.00048	0.82044	3361.209	2758.011

A_23_P48637		Unknown	A_23_P48	-1.21888	5.35E-09	0.82043	2364.784	1939.419
A_32_P49	XM_37615	similar to ri	LOC40101	-1.21888	0.00003	0.82043	74773.54	61414.63
A_24_P161914		Unknown	A_24_P16	-1.21898	2.83E-06	0.82035	39245.06	32198.79
A_23_P15	NM_08088	Homo sapi	DBN1,D0S	-1.21901	0.00008	0.82034	2594.08	2128.538
A_23_P14	NM_01448	Homo sapi	ZNF544	-1.2194	0.00052	0.82007	1877.004	1540.227
A_23_P60	NM_00331	Homo sapi	DNAJC7,T	-1.21966	0.00003	0.8199	4703.067	3854.678
A_24_P16	NM_00101	ribosomal p	RPS7	-1.2199	0.00004	0.81974	97331.72	79616.84
A_23_P98	NM_01545	Homo sapi	ZDHHC5,Z	-1.22013	0.0007	0.81958	1443.083	1181.164
A_23_P32	NM_13896	Homo sapi	RPS4Y2	-1.22048	0.00027	0.81935	44717.75	36572.95
A_23_P88	NM_13837	Homo sapi	TTC5	-1.22082	0.00066	0.81912	1567.667	1285.853
A_23_P81	NM_00313	Homo sapi	SRP19	-1.22091	0.00021	0.81906	8635.671	7055.237
A_23_P47	NM_00601	Homo sapi	CLPP	-1.22115	0.00003	0.8189	16491.79	13521.86
A_23_P20	NM_00325	Homo sapi	THRA,AR7	-1.22131	0.00022	0.81879	2732.593	2239.048
A_23_P21	NM_01848	uncharacte	HT007	-1.22145	0.00002	0.8187	5229.811	4283.168
A_24_P29	XM_37098	Homo sapi	ENST0000	-1.22218	0.0001	0.81821	35742.39	29183.04
A_23_P35	NM_00654	calcium/cal	CAMKK2,C	-1.22245	0.00017	0.81803	6817.939	5578.574
A_23_P60	NM_01520	KIAA0460	KIAA0460	-1.22252	0.0001	0.81798	2672.24	2184.438
A_24_P15	NM_01595	thioredoxin	TXNDC14,	-1.22257	0.00029	0.81795	4456.305	3641.86
A_23_P83	NM_00426	15 kDa sel	15-Sep	-1.22295	0.00089	0.81769	18083.96	14844.69
A_23_P15	NM_01221	Homo sapi	LTB4DH,M	-1.22342	0.00001	0.81738	18409.21	15027.27
A_23_P11	AB029039	RNA bindir	RBM16	-1.22354	0.00081	0.8173	1749.099	1428.054
A_24_P85	CB962310	GRB2-ass	GAB1	-1.22366	0.00038	0.81722	38525.78	31396.3
A_24_P12	NM_00676	tyrosine 3-r	YWHAE,M	-1.22382	0.00081	0.81711	1468.49	1200.143
A_23_P38	NM_02074	Homo sapi	AARSL,KIA	-1.22402	0.0002	0.81698	1602.655	1304.614
A_23_P10	NM_00393	Adaptor-rel	AP3D1	-1.22406	7.54E-06	0.81695	30198.82	24704.34
A_23_P85	NM_00064	interleukin	IL13RA2,IL	-1.22414	0.00027	0.8169	38503.96	31538.4
A_24_P298179		Unknown	A_24_P298	-1.22426	0.00063	0.81682	7726.934	6293.743
A_23_P43	NM_00249	Homo sapi	NDUFB6,C	-1.22427	0.00005	0.81681	13173.03	10777.41
A_23_P86171		FXD2_HUM	THC18578	-1.22451	0.00077	0.81666	1812.843	1488.915
A_23_P35	NM_05401	Homo sapi	FUSIP1,NS	-1.22531	0.00089	0.81612	2540.015	2072.387
A_32_P22	NM_00551	high-mobili	HMGN2,HI	-1.22533	0.00001	0.81611	41936.98	34272.13
A_23_P81	NM_00346	Homo sapi	PTP4A1,HI	-1.22572	0.00006	0.81585	38186.71	31236.46
A_24_P21	NM_01908	hypothetica	PAF1,PD2	-1.22586	0.00058	0.81576	2300.031	1878.598
A_23_P10	NM_13828	Homo sapi	NUP35,MF	-1.22628	0.00004	0.81548	5322.083	4340.173
A_32_P28	NM_00309	small nucle	SNRPA1	-1.2264	1.39E-06	0.8154	10774.74	8789.398
A_23_P11	NM_01406	Homo sapi	MCTS1,MC	-1.22649	0.00005	0.81534	20161.29	16460.66
A_23_P20	NM_05284	Homo sapi	WDR34,MC	-1.22654	0.00005	0.8153	5040.299	4108.852
A_23_P87	NM_02206	mitochondr	MRPL17,R	-1.22671	5.24E-06	0.81519	9937.855	8109.693
A_23_P36	NM_00311	spastic par	SPG7,CAF	-1.22674	4.60E-06	0.81517	4552.972	3711.522
A_23_P16	NM_02477	Zinc finger,	ZMYM1	-1.22693	0.00026	0.81504	1623.655	1323.908
A_24_P34	BC032431	Hypothetic	LOC38920	-1.22698	0.00014	0.81501	2104.147	1713.753
A_23_P37	NM_00632	Homo sapi	TIMM23,TI	-1.22708	0.00043	0.81495	7229.255	5912.11
A_23_P20	NM_00430	Rho GDP c	ARHGDIA,	-1.22739	0.00001	0.81473	3124.277	2545.512
A_23_P11	NM_01553	Homo sapi	ZZZ3,DKF	-1.22783	1.75E-10	0.81445	881.3157	717.3027
A_23_P82	NM_02116	Homo sapi	GATAD1,C	-1.22801	0.0009	0.81433	1340.085	1092.362
A_24_P884915		Unknown	A_24_P884	-1.22812	0.00016	0.81426	20202.61	16501.01
A_32_P21	NM_01845	centrosom	CENPJ,LA	-1.22819	0.00018	0.8142	2190.676	1782.386
A_23_P38	NM_00297	Homo sapi	ATXN2,AT	-1.22834	0.00021	0.81411	6884.542	5610.679
A_32_P99	NM_01625	high-gluco	YTHDF2,H	-1.22856	0.00067	0.81396	8333.584	6807.875
A_23_P34	NM_14472	Homo sapi	ZMAT2,FL	-1.22882	0.00041	0.81379	6034.774	4909.049
A_24_P26	XM_37710	similar to 4	LOC40163	-1.22928	0.00006	0.81348	46936.78	38299.79
A_23_P15	NM_00500	Homo sapi	NDUFB9,B	-1.22929	0.00018	0.81348	63686.25	51958.05

A_24_P15	AK098605	Formin 2	FMN2	-1.22938	0.00006	0.81342	101703.3	82879.3
A_23_P59	NM_00309	Homo sapi	SNRPC,FL	-1.22943	0.00003	0.81338	25620.12	20877.27
A_23_P10	NM_00693	Homo sapi	SNRPD1,S	-1.22951	0.00001	0.81334	58385.38	47417.33
A_23_P16	NM_00517	ATP syntha	ATP5G1,A	-1.22977	5.80E-07	0.81316	21847.59	17761.41
A_23_P12	NM_02255	Homo sapi	XRCC4	-1.22979	0.00083	0.81314	1411.597	1149.678
A_23_P14	NM_13880	similar to m	LOC13414	-1.22997	0.00004	0.81303	3159.912	2568.744
A_23_P21	NM_01665	Homo sapi	KDELR3,E	-1.23029	0.00015	0.81282	2530.837	2054.507
A_23_P11	NM_18164	Homo sapi	CKLF,C32,	-1.23053	0.00005	0.81266	7331.442	5957.762
A_24_P101694		Unknown	A_24_P10	-1.23057	0.00006	0.81263	37220.9	30137.48
A_23_P16	NM_00684	Homo sapi	ILVBL,AHA	-1.23095	0.00003	0.81238	3738.984	3040.256
A_24_P10	NM_00393	suppressio	ST13,HIP,I	-1.23141	0.0003	0.81208	1595.982	1297.552
A_23_P16	NM_00338	Homo sapi	VIM,FLJ36	-1.23143	0.00004	0.81206	83507.84	67677.52
A_23_P12	NM_00309	Homo sapi	SNRPE,SM	-1.23168	1.10E-06	0.8119	25464.63	20688.98
A_23_P11	NM_13928	Homo sapi	WDR36,TA	-1.23173	0.00057	0.81187	1617.561	1314.124
A_24_P186986		Unknown	ENST0000	-1.23207	6.33E-07	0.81165	21161.65	17181.07
A_23_P35	AL833070	SH3 multip	SH3MD1	-1.23223	0.00059	0.81154	1295.615	1051.304
A_23_P32	NM_01633	Homo sapi	SRRM2,30	-1.23264	0.00006	0.81127	16395.31	13326.17
A_23_P79	NM_01610	Homo sapi	FKBP7,FK	-1.23276	0.00046	0.81119	1460.409	1184.75
A_23_P21	NM_00612	iduronate 2	IDS,MPS2,	-1.23319	0.00001	0.81091	3237.28	2625.134
A_23_P13	NM_03265	Homo sapi	DHX37,DD	-1.23335	0.00016	0.8108	3908.507	3169.918
A_23_P20	NM_02429	leukocyte r	LENG4,BB	-1.23343	3.58E-07	0.81075	10790.92	8748.289
A_24_P21	NM_03067	SEC13 (S.	SEC13L1,S	-1.23346	0.00016	0.81073	3314.807	2686.206
A_23_P15	NM_01989	Homo sapi	C3orf4,HSI	-1.23357	0.00003	0.81066	5902.145	4785.607
A_23_P58072		Unknown	A_23_P58	-1.23362	0.00067	0.81062	69119.19	56072.05
A_24_P45	BX451504	Transcribe	BX451504	-1.2338	0.00069	0.8105	3029.56	2466.487
A_23_P97	NM_02215	Rag C prot	RRAGC,G	-1.23427	0.00003	0.8102	3533.274	2864.202
A_23_P84	NM_00341	Homo sapi	ZNF9,DM2	-1.23443	9.16E-07	0.81009	13956.36	11308.12
A_23_P21	NM_01527	Homo sapi	TRIM2,RNI	-1.23446	7.01E-06	0.81007	2460.812	1991.924
A_23_P58	NM_01594	Homo sapi	PELO,CGI-	-1.23466	1.06E-06	0.80994	11185.75	9055.031
A_23_P67	NM_00174	Homo sapi	CAPNS1,3	-1.23522	0.00071	0.80958	4789.85	3890.056
A_23_P35	NM_17490	hypothetica	MGC2390	-1.23545	5.91E-06	0.80942	3978.063	3221.38
A_24_P69	NM_08092	hypothetica	LOC96610	-1.23562	0.00026	0.80931	2390.792	1935.188
A_23_P77	NM_01902	protein arg	PRMT7,FL	-1.23582	0.00075	0.80918	6449.541	5240.638
A_23_P25	NM_00664	Homo sapi	SDCCAG3	-1.23587	0.00023	0.80914	8553.151	6935.453
A_23_P42	NM_00477	Homo sapi	PPARBP,F	-1.23606	9.39E-06	0.80902	6134.635	4957.401
A_23_P96	NM_00630	Homo sapi	SRPX,ETX	-1.23612	0.00004	0.80898	3783.459	3060.177
A_23_P11	NM_00138	Homo sapi	DPH1,DPH	-1.23625	3.32E-06	0.8089	12818.28	10380.36
A_23_P66	U18197	ATP citrate	ACLY	-1.23634	0.00002	0.80884	2958.161	2392.331
A_23_P376449		Unknown	A_23_P37	-1.23635	0.00014	0.80883	5029.758	4077.684
A_23_P93	NM_00543	Homo sapi	ARHGEF5	-1.23644	0.00055	0.80877	1661.466	1345.901
A_23_P32	NM_13861	Homo sapi	DHX30,DD	-1.23649	0.00027	0.80874	3103.252	2508.996
A_32_P16	CD101793	Chromosor	C6orf209	-1.23659	0.00048	0.80867	7014.473	5656.509
A_23_P13	NM_01603	Homo sapi	MRPS2,CC	-1.23667	8.92E-06	0.80862	17441.52	14102.29
A_24_P93	NM_01434	mitochondr	MTCH2,HS	-1.23679	0.00082	0.80855	1403.968	1137.664
A_24_P84	NM_00290	RD RNA-bi	RDBP,RD,	-1.23684	0.00026	0.80851	24732.43	20069.66
A_23_P40	NM_01225	Homo sapi	RRAS2,TC	-1.23705	0.00084	0.80838	1535.135	1237.895
A_23_P28	NM_00283	Homo sapi	PTPRA,LR	-1.23714	0.00035	0.80831	2114.424	1710.677
A_23_P15	NM_01613	Homo sapi	CHCHD2,C	-1.23731	0.00003	0.8082	80338.11	65089.22
A_32_P66625		ALU2_HUM	THC19452	-1.2377	0.00051	0.80795	2391.492	1928.477
A_24_P35	NM_00229	laminin rec	RPSA,LRP	-1.23788	8.61E-07	0.80783	69210.27	55958.81
A_23_P11	NM_01752	Homo sapi	LEPROT,L	-1.2383	6.68E-07	0.80756	5773.959	4662.699
A_32_P86	NM_00472	Ac-like tran	ZBED1,AL	-1.23834	0.00006	0.80753	3120.351	2517.232

A_24_P850187		Unknown	A_24_P85(	-1.23909	8.02E-07	0.80704	54365.06	43907.67
A_32_P11	NM_00660	unactive pr	PTGES3,P	-1.2391	0.00003	0.80704	5398.009	4363.45
A_23_P56	NM_01815	Homo sapi	SLC4A1AF	-1.23916	0.00003	0.807	3500.435	2824.163
A_24_P29	NM_03096	hypothetica	TMEM14B	-1.23952	0.00007	0.80676	11344.75	9179.055
A_24_P333112		Unknown	A_24_P33	-1.23978	0.00018	0.80659	69054.09	56019.61
A_23_P21	NM_01592	Homo sapi	C6orf82,CU	-1.23984	0.00025	0.80655	20365.81	16474.32
A_24_P418106		Unknown	A_24_P41	-1.23992	0.00005	0.8065	24250.55	19568.5
A_23_P75	NM_00362	Homo sapi	PPFIA1,LIF	-1.23996	0.00012	0.80647	1594.732	1285.923
A_23_P25	NM_00570	Homo sapi	RNUT1,KP	-1.23996	0.00001	0.80648	7505.027	6062.609
A_23_P12	NM_01425	Homo sapi	SLC25A15	-1.23999	0.00006	0.80646	5955.118	4809.377
A_23_P39	AK098781	Homo sapi	AK098781	-1.24004	0.00087	0.80642	1633.899	1312.186
A_23_P15	NM_01215	Homo sapi	EPB41L1,4	-1.24024	0.00096	0.8063	3770.182	3045.521
A_23_P14	NM_01416	Homo sapi	METTL5,H	-1.24044	0.00076	0.80617	8687.159	7025.447
A_23_P15	NM_01552	small fragm	REXO2,RF	-1.24053	0.00061	0.80611	13277.9	10731.16
A_32_P233727		Unknown	A_32_P23	-1.24056	0.00005	0.80609	3476.65	2802.679
A_23_P21	NM_01410	Homo sapi	ATAD2,MG	-1.24062	0.00039	0.80605	1520.231	1229.175
A_23_P10	AK026518	Myosin hea	MYOHD1	-1.24066	0.00027	0.80602	3479.588	2805.624
A_23_P99	AK074057	Hypothetica	FLJ10357	-1.24072	0.00005	0.80598	4442.975	3574.756
A_23_P10	NM_03144	Homo sapi	C18orf21,X	-1.241	0.0001	0.8058	2385.322	1922.459
A_23_P59	NM_03282	Homo sapi	SLC35B4,Y	-1.24113	0.00063	0.80572	1315.208	1058.552
A_24_P34	NM_00229	laminin rec	RPSA,LRP	-1.24119	0.00003	0.80568	43002.16	34608.47
A_24_P789842		Unknown	A_24_P78	-1.24141	5.90E-06	0.80554	52895.91	42668.71
A_24_P57	AK057167	Hypothetica	AK057167	-1.24147	0.00023	0.8055	1790.369	1443.257
A_23_P44	NM_01428	pM5 protei	NOMO1,PI	-1.24184	0.00088	0.80525	1250.215	1007.947
A_23_P23	NM_00725	Homo sapi	VPS45A,H	-1.24192	0.00009	0.8052	3149.331	2540.214
A_23_P21	NM_01339	Homo sapi	C6orf49,OE	-1.24204	0.00053	0.80513	16784.85	13562.99
A_24_P264549		Unknown	A_24_P26	-1.24205	1.36E-07	0.80512	63426.49	51059.11
A_23_P91	NM_03277	Homo sapi	MINA,MDI(	-1.24205	0.00013	0.80512	1625.446	1308.512
A_24_P255654		Unknown	A_24_P25	-1.24233	0.00001	0.80494	22017.77	17751.84
A_23_P16	AK026132	PHD finger	PHF17	-1.24233	0.00002	0.80494	3083.689	2481.979
A_24_P24	NM_18150	Exosome c	EXOSC8	-1.24234	1.08E-07	0.80493	13069.9	10520.63
A_23_P14	AL161962	Zinc finger,	ZDHHC9	-1.24314	4.43E-06	0.80441	3513.012	2825.489
A_24_P30	XM_06088	similar to p	LOC12819	-1.24329	0.00007	0.80432	24669.02	19885.46
A_23_P10	XM_06750	Homo sapi	ENST0000	-1.24335	0.00054	0.80428	27921.26	22521.87
A_24_P26	NM_08092	hypothetica	LOC96610	-1.24337	0.00021	0.80426	2695.11	2168.065
A_24_P119094		Unknown	A_24_P11	-1.24341	0.00029	0.80424	16135.83	12935.6
A_23_P13	NM_02407	Homo sapi	TSEN34,LE	-1.24363	0.00044	0.8041	10411.39	8376.613
A_24_P229616		Unknown	ENST0000	-1.24367	0.00001	0.80407	25080.41	20197.21
A_23_P10	D63478	Ubiquitin a	UBAP2L	-1.24374	0.00043	0.80403	2330.938	1876.511
A_23_P30	NM_00415	Homo sapi	PPP2CB,P	-1.24427	0.00022	0.80368	2536.93	2044.589
A_23_P10	NM_01470	Homo sapi	RAB11FIP	-1.24435	0.00017	0.80363	1728.32	1387.604
A_23_P91	NM_00375	Homo sapi	EIF3S7,MC	-1.24445	9.92E-07	0.80357	25519.14	20520.09
A_32_P38	NM_00256	poly(A)-bin	PABPC1,P	-1.2446	3.05E-06	0.80347	70878.72	57014.98
A_23_P25	NM_01838	hypothetica	FLJ11305,	-1.24463	4.46E-06	0.80345	3276.854	2631.89
A_24_P101722		Unknown	ENST0000	-1.24501	1.75E-06	0.80321	26812.87	21562.19
A_23_P31	NM_00314	single-strar	SSBP1,SS	-1.24507	1.43E-06	0.80317	25649.38	20577.78
A_23_P69	NM_00160	acetyl-Coe	ACAA1,AC	-1.2455	0.00009	0.80289	3916.254	3141.243
A_32_P20	NM_00182	CDC28 pro	CKS1B,CK	-1.24581	0.00007	0.80269	9194.685	7408.505
A_23_P76	NM_15232	Homo sapi	ADSSL1,F	-1.24591	0.00061	0.80263	2700.327	2164.756
A_23_P14	NM_03202	Homo sapi	TM2D1,BB	-1.24644	0.00036	0.80229	4940.829	3976.986
A_23_P20	NM_08042	Homo sapi	PTPN2,PT	-1.24648	0.00009	0.80226	2752.239	2203.5
A_24_P196024		Unknown	ENST0000	-1.24674	0.0003	0.80209	24652.02	19828.53

A_23_P97	NM_02215	Homo sapi	RRAGC,G	-1.2471	0.00003	0.80186	1992.21	1597.089
A_24_P15	XM_06289	similar to p	LOC12198	-1.24759	0.00058	0.80155	25763.85	20681.39
A_24_P34	NM_20343	Peptidylprc	PPIA	-1.24797	0.00025	0.8013	42109.02	33811.7
A_24_P384411		Unknown	A_24_P38	-1.24798	0.00024	0.80129	51241.38	41218.52
A_23_P15	NM_00379	Homo sapi	CTNNAL1,	-1.2485	0.00019	0.80096	11645.46	9340.557
A_24_P78	NM_00229	laminin rec	RPSA,LRP	-1.24867	6.03E-06	0.80085	79896.48	64131.9
A_23_P14	NM_01863	Homo sapi	WSB2,SBA	-1.24878	0.0003	0.80078	1497.932	1197.824
A_23_P55	NM_01764	hypothetica	FTSJ3,EP	-1.24887	0.00006	0.80072	5261.448	4201.053
A_24_P36	XM_29296	similar to p	LOC34417	-1.24896	0.00005	0.80067	33759.39	27085.52
A_23_P45	NM_01773	FLJ20259	FLJ20259	-1.24896	0.00034	0.80067	2641.261	2117.373
A_23_P22	NM_01600	Homo sapi	SH3GLB1,	-1.2493	0.00044	0.80045	3317.111	2657.932
A_32_P92	NM_03238	hypothetica	COG8,DO	-1.24933	4.89E-07	0.80043	4493.271	3597.01
A_23_P14	NM_00586	Homo sapi	OPRS1,SF	-1.24942	4.80E-06	0.80037	7541.389	6040.863
A_24_P28	NM_00140	eukaryotic	EEF1G,EF	-1.24943	0.00014	0.80036	84330.2	67192.94
A_24_P153324		Unknown	A_24_P15	-1.24949	0.00006	0.80033	54330.44	43622.9
A_23_P11	NM_01429	Homo sapi	TRAM1,TR	-1.24996	0.00056	0.80003	10938.82	8749.619
A_23_P13	NM_01600	WD repeat	CGI-48	-1.25002	4.08E-06	0.79999	5797.129	4632.922
A_23_P72	NM_00038	Homo sapi	SLC25A20	-1.25038	0.00093	0.79976	1256.859	1002.215
A_23_P10	NM_01458	Homo sapi	ERO1L,ER	-1.25039	0.00005	0.79975	5335.024	4265.679
A_23_P63	NM_03233	Homo sapi	C10orf58,M	-1.2504	0.00009	0.79974	5765.947	4603.7
A_24_P37	NM_01894	cytochrome	CYCS,CY	-1.25054	0.00006	0.79965	25465.28	20317.22
A_23_P70	NM_01813	Homo sapi	C6orf139,F	-1.25058	0.00005	0.79963	2099.697	1681.023
A_23_P45	NM_00719	Homo sapi	PROSC	-1.25078	0.00059	0.7995	1315.358	1050.309
A_32_P98511		Unknown	A_32_P98	-1.25094	2.23E-06	0.7994	27803.65	22189.23
A_23_P13	NM_01806	Homo sapi	FANCL,PC	-1.25121	0.00012	0.79923	10622.43	8515.981
A_32_P22275		Unknown	A_32_P22	-1.25176	0.00036	0.79887	2495.441	1993.856
A_23_P14	NM_00521	Homo sapi	DDOST,OS	-1.25176	0.00089	0.79888	8331.748	6673.77
A_23_P15	NM_00119	Homo sapi	BID,MGC1	-1.25181	0.00004	0.79885	1733.885	1384.287
A_32_P39	AK055214	CDNA FLJ	AK055214	-1.25194	0.00017	0.79876	1910.871	1529.557
A_24_P417996		Unknown	A_24_P41	-1.25229	4.56E-07	0.79854	5249.702	4193.45
A_32_P14	NM_05402	Homo sapi	CNOT7,CA	-1.25246	0.00006	0.79843	1839.098	1469.959
A_23_P12	NM_00682	Homo sapi	RAB40B,R	-1.25252	0.00009	0.79839	1558.907	1243.3
A_24_P87	BC061590	GTF2I repe	GTF2IRD2	-1.25258	0.0006	0.79835	1571.228	1250.872
A_32_P16	AK021744	CDNA FLJ	AK021744	-1.25259	0.00064	0.79835	2274.254	1810.606
A_24_P12	NM_00278	proteasom	PSMA4,HC	-1.25268	3.80E-06	0.79829	17476.46	13916.91
A_23_P67	NM_03249	Homo sapi	JAGN1,GL	-1.25278	0.00004	0.79822	10464.89	8381.904
A_23_P50	NM_00269	Homo sapi	POLD1,PC	-1.25304	6.75E-06	0.79806	2879.927	2297.695
A_23_P46	NM_00287	RAB13, me	RAB13,GI	-1.25312	0.00099	0.79801	1738.509	1383.891
A_23_P78	NM_00143	fibrillarin	FBL,FIB,FL	-1.25327	0.00001	0.79791	29645.84	23580.1
A_24_P392690		Unknown	A_24_P39	-1.25366	0.00002	0.79767	6824.175	5431.431
A_23_P25	NM_01643	glycolipid t	GLTP	-1.25378	2.14E-06	0.79759	2631.375	2099.788
A_24_P87	NM_01797	mitochondr	MRPL20,L	-1.25381	0.00006	0.79757	23592.3	18715.88
A_24_P914208		Unknown	A_24_P91	-1.25404	0.00065	0.79742	5341.754	4235.308
A_23_P71	NM_01794	Homo sapi	NOL8,Nop	-1.25416	3.45E-06	0.79735	8566.855	6815.454
A_23_P92	NM_00434	Homo sapi	CASP3,CP	-1.25428	0.00062	0.79727	7381.493	5910.885
A_23_P91	NM_01338	prolactin re	PREB,SEC	-1.25433	6.71E-06	0.79724	3730.877	2974.441
A_23_P13	NM_00693	Homo sapi	SKP1A,OC	-1.25437	0.00028	0.79721	61691.06	49409.36
A_32_P84	BC041995	Homo sapi	BC041995	-1.25461	0.00048	0.79706	1094.742	872.5042
A_23_P13	NM_00467	Homo sapi	ASH2L,AS	-1.25493	0.00032	0.79686	2490.059	1979.629
A_23_P63	NM_00649	lectin, gala	LGALS8,G	-1.25498	0.00053	0.79683	1376.299	1095.591
A_24_P93	AK097197	Hepatocyte	HGS	-1.25501	0.0001	0.79681	1506.047	1199.859
A_24_P75308		Unknown	A_24_P75	-1.2552	1.31E-06	0.79668	33792.93	26971.32

A_23_P13	NM_00156	Homo sapi	IL13RA1,N	-1.25526	0.0001	0.79665	3477.746	2770.68
A_23_P20	NM_00015	glucosidas	GBA,GCB,	-1.2553	0.00092	0.79662	1372.752	1097.401
A_23_P30	NM_00123	Homo sapi	CCNH,CAI	-1.25548	0.00091	0.79651	4751.427	3805.723
A_23_P35	NM_00037	Homo sapi	UROS	-1.25562	0.00024	0.79642	1893.066	1503.76
A_24_P178415		Unknown	A_24_P17	-1.25565	3.42E-07	0.7964	45038.23	35902.1
A_23_P99	NM_01837	hypothetica	FLJ11259	-1.25573	1.51E-06	0.79635	2712.492	2160.015
A_23_P21	NM_00237	Homo sapi	MAP4,MG	-1.25575	0.00003	0.79634	3334.446	2657.331
A_24_P39	XM_11639	similar to p	LOC20222	-1.25607	0.00007	0.79613	29916.51	23845.77
A_23_P21	NM_01545	Homo sapi	THUMPD3	-1.25618	0.00072	0.79606	1716.554	1361.804
A_23_P65	NM_19852	Similar to F	HDDC3,MC	-1.25635	3.23E-06	0.79596	3029.09	2410.917
A_23_P10	NM_01406	nin one bin	NOB1P,AF	-1.25641	0.00013	0.79592	3869.833	3084.364
A_24_P912871		Unknown	A_24_P91	-1.25646	0.00021	0.79588	10521.77	8411.612
A_23_P25	X78926	Similar to Z	HZF3	-1.25659	5.98E-06	0.7958	3087.719	2457.809
A_32_P98	NM_00425	cytochrome	COX5A,VA	-1.2566	0.00016	0.7958	35830.92	28633.13
A_23_P65	NM_01656	Homo sapi	BFAR,BAR	-1.2568	4.46E-06	0.79567	2454.384	1953.403
A_23_P54	NM_00629	Homo sapi	TYRO3,BY	-1.25748	0.00041	0.79524	2607.486	2080.21
A_23_P37	NM_01428	Homo sapi	DNAJC8,S	-1.25763	0.00002	0.79515	6458.136	5151.239
A_24_P36	NM_00578	ATP bindin	TXNDC9,A	-1.25779	0.00008	0.79505	1871.997	1487.028
A_23_P54	NM_03247	Homo sapi	MRPL38,R	-1.25802	0.00024	0.7949	15704.53	12411.08
A_23_P21	NM_01608	palladin	KIAA0992,	-1.25827	2.81E-06	0.79474	4733.636	3765.938
A_23_P21	NM_01639	TH1-like (D	TH1L,TH1,	-1.25832	7.33E-07	0.79471	4425.187	3516.373
A_23_P21	D50683	Homo sapi	TGF-betaII	-1.25837	5.72E-06	0.79468	6125.411	4871.353
A_24_P187154		Unknown	ENST0000	-1.25837	0.00008	0.79468	7177.674	5716.732
A_24_P341476		Unknown	A_24_P34	-1.25865	0.00056	0.7945	31519.25	25191.13
A_24_P10	NM_01645	hypothetica	LOC51234	-1.25865	0.00017	0.7945	11651.14	9225.789
A_23_P14	NM_18277	Homo sapi	RPS3A	-1.25865	0.00002	0.7945	153904.4	122425.8
A_23_P76	NM_01883	13kDa diffe	NDUFA12,	-1.25873	3.70E-08	0.79445	20052.17	15926.16
A_23_P11	NM_08039	Homo sapi	PTP4A2,HI	-1.25875	0.00003	0.79444	14652.06	11691.88
A_23_P40	NM_03144	hypothetica	MGC4268	-1.25889	3.98E-06	0.79435	4090.128	3248.542
A_24_P777185		Unknown	A_24_P77	-1.25894	0.00069	0.79432	2794.546	2219.9
A_23_P25	BC040043	Steroid rec	SRA1	-1.25969	0.00002	0.79385	3338.783	2646.678
A_23_P11	NM_00012	Homo sapi	ERCC5,XF	-1.25971	0.00037	0.79383	2493.995	1985.312
A_24_P87	AK093346	CDNA FLJ	AK093346	-1.25972	1.56E-06	0.79383	3142.673	2494.324
A_24_P28	AK074029	Hypothetic	FLJ20255	-1.25998	0.0007	0.79367	1243.927	985.8296
A_23_P11	NM_00056	glutathione	GSTM1,MU	-1.26002	0.00002	0.79364	5265.334	4180.593
A_32_P145243		Unknown	A_32_P14	-1.26033	0.00018	0.79344	1758.739	1398.616
A_24_P80	NM_00190	catenin (ca	CTNNA1,C	-1.26039	0.00021	0.7934	2060.328	1642.715
A_32_P19	NM_15227	hypothetica	LONRF1,R	-1.26041	0.00002	0.79339	1630.605	1294.05
A_24_P15	NM_00559	nascent-po	NACA,MG	-1.26047	0.00031	0.79336	97371.28	77630.17
A_24_P33	BC006174	F-box prote	FBXO42	-1.261	0.0001	0.79302	1453.123	1151.736
A_23_P36	NM_13326	WIRE prote	WIRE,WIC	-1.26109	0.00014	0.79296	2824.977	2241.757
A_23_P21	NM_00408	Homo sapi	TIMM8A,D	-1.26148	0.00004	0.79272	6177.867	4902.647
A_23_P94	NM_00191	Homo sapi	CTSL,MEF	-1.26153	0.00007	0.79269	33381.63	26324.42
A_32_P27	NM_00195	E2F transc	E2F6,E2F-	-1.26176	0.00006	0.79254	1950.85	1544.093
A_23_P50	NM_07847	Homo sapi	TM2D2,BL	-1.262	3.99E-06	0.79239	3804.733	3014.976
A_23_P20	NM_17506	hypothetica	LOC28436	-1.26218	0.00017	0.79228	3609.695	2864.793
A_32_P19	BX537704	Amyotroph	ALS2CR13	-1.26228	0.00002	0.79222	2783.961	2208.64
A_24_P127051		Unknown	A_24_P12	-1.26235	0.00006	0.79217	256801.5	204280.5
A_23_P34	NM_00412	Homo sapi	GTF2F2,B	-1.26239	0.00054	0.79215	4091.908	3254.244
A_24_P27	NM_00449	general tra	GTF2A2,TI	-1.2626	1.19E-06	0.79202	14599.26	11590.37
A_24_P84	BC032295	Ribosomal	RPL22	-1.26262	0.0001	0.79201	2011.096	1591.021
A_23_P37	NM_01600	Homo sapi	TRNT1,CC	-1.26281	0.00093	0.79188	1262.321	1001.992

A_24_P754	XM_37793	similar to p	LOC40225	-1.26286	0.00044	0.79186	27269.53	21744.52
A_23_P165	NM_00115	Homo sapi	ANXA4,AN	-1.26289	0.00052	0.79183	5987.915	4754.851
A_24_P367	XM_37203	Homo sapi	XM_37203	-1.26302	9.57E-08	0.79175	103846.6	82310.01
A_23_P457	XM_37278	similar to p	LOC39106	-1.26303	1.82E-07	0.79175	50746.98	40212.13
A_32_P140	NM_00229	laminin rec	RPSA,LRP	-1.26307	4.22E-07	0.79172	48235.42	38238.02
A_24_P873	BC039021	CDNA clon	BC039021	-1.26317	0.00013	0.79166	2142.621	1691.374
A_24_P254437		D86074 ins	THC19409	-1.26342	0.00024	0.7915	3156.252	2506.141
A_23_P21	NM_00169	Homo sapi	ATP6V1A,I	-1.26358	0.00057	0.7914	2631.203	2087.229
A_24_P384539		Unknown	ENST0000	-1.26384	5.68E-06	0.79124	112622.5	89329.38
A_24_P195	AB014563	KIAA0663	KIAA0663	-1.26388	9.25E-08	0.79121	4468.083	3535.427
A_23_P160	NM_00257	Homo sapi	PAFAH1B3	-1.2641	0.00009	0.79108	1600.549	1264.905
A_23_P138	NM_00405	Homo sapi	BNIP3,NIP	-1.26416	0.00003	0.79104	17939.95	14239.89
A_23_P42	NM_00094	Homo sapi	PPIC,CYP	-1.26417	1.13E-07	0.79103	5394.955	4267.01
A_24_P920	AF233453	Protein kin	PRKCBP1	-1.26429	0.00007	0.79096	1955.075	1549.298
A_23_P138	NM_19929	Homo sapi	APITD1,M	-1.26432	0.00005	0.79094	2922.659	2310.24
A_24_P280	NM_00533	Huntingtin	HIP2	-1.26438	0.00003	0.7909	20595.57	16221.21
A_23_P200	Z24727	Tropomyos	TPM1	-1.2645	0.00002	0.79083	1295.889	1025.616
A_24_P27	NM_00381	Homo sapi	CDS2	-1.26452	0.00049	0.79081	1592.604	1255.081
A_23_P160	NM_01606	Homo sapi	GLRX2,GF	-1.26458	4.17E-08	0.79078	10257.98	8114.413
A_24_P384	NM_02403	hypothetica	MGC2803	-1.2646	0.00055	0.79076	2057.497	1622.696
A_24_P349756		Unknown	A_24_P34	-1.26495	1.14E-07	0.79055	16230.32	12817.51
A_23_P125	NM_01897	Homo sapi	TERF2IP,F	-1.26497	0.00064	0.79053	3862.125	3049.169
A_23_P134	NM_00561	Ras homol	RHEB,RHE	-1.26498	0.00057	0.79052	14121.21	11216.21
A_23_P125	NM_17574	Homo sapi	RHOC,ARI	-1.26506	0.00012	0.79048	7809.354	6196.95
A_24_P390	NM_00376	splicing fac	SFRS9,SR	-1.26532	0.00003	0.79031	11850.22	9356.719
A_23_P335	NM_01751	Homo sapi	SLC35F2,H	-1.26534	0.00008	0.7903	1834.423	1453.224
A_24_P341427		Unknown	A_24_P34	-1.26535	2.80E-06	0.79029	46559.97	36891.7
A_23_P882	NM_01604	CGI-112 pr	C14orf122,	-1.26567	0.00045	0.7901	3419.888	2695.479
A_24_P305	NM_03248	novel MCM	MCM8,MG	-1.26568	0.00065	0.79009	1347.65	1066.312
A_23_P144	NM_00671	Homo sapi	COPS8,CC	-1.2657	3.96E-07	0.79008	3236.722	2557.229
A_23_P200	NM_00640	Acidic (leu	ANP32B	-1.26574	4.78E-06	0.79005	7869.714	6200.865
A_23_P104	NM_00608	Homo sapi	BPNT1,PIF	-1.26583	1.42E-06	0.79	3423.305	2705.247
A_23_P213	NM_02459	Homo sapi	PANK3,FL	-1.2661	0.00042	0.78982	1899.999	1496.039
A_23_P13	NM_02012	potassium	KCMF1,FI	-1.2663	2.50E-07	0.7897	10753.3	8497.337
A_23_P128	NM_00280	Homo sapi	PSMC6,P4	-1.26637	0.00023	0.78966	10047.74	7967.736
A_23_P337	NM_00694	Homo sapi	SRP72	-1.26688	0.00002	0.78934	13146.42	10333.53
A_23_P104	NM_08059	Homo sapi	OSBPL1A,	-1.26689	0.00008	0.78933	1882.162	1488.13
A_23_P109	NM_08092	hypothetica	LOC96610	-1.26694	0.00032	0.7893	1956.566	1541.522
A_24_P772	NM_20343	Peptidylpro	PPIA	-1.26695	4.00E-08	0.7893	48606.84	38350.69
A_24_P175	NM_00332	thioredoxin	TXN,TRX,N	-1.2674	1.90E-07	0.78902	133684.8	105334.3
A_23_P630	NM_00496	Homo sapi	IDE,INSUL	-1.26757	1.24E-06	0.78891	3391.104	2674.42
A_23_P405	NM_02092	Homo sapi	BCOR,MA	-1.26766	0.00034	0.78886	1108.705	875.4407
A_23_P628	NM_00268	Homo sapi	EXOSC10,	-1.26769	0.00028	0.78884	3793.248	2996.039
A_24_P135515		Unknown	A_24_P13	-1.26782	4.99E-06	0.78876	35756.13	28226.07
A_23_P622	NM_02452	NF-kappaE	NKAP,FLJ2	-1.26782	0.00005	0.78876	2785.272	2197.182
A_23_P154	NM_00271	Homo sapi	PPP1R7,S	-1.26816	2.59E-06	0.78854	4412.069	3475.733
A_23_P26	NM_01799	Homo sapi	DET1,FLJ1	-1.26837	0.00075	0.78842	2145.375	1687.585
A_24_P849	NM_00680	protein tra	SEC61B	-1.26851	5.52E-08	0.78833	41206.95	32514.58
A_23_P218	NM_00200	Homo sapi	FGF2,BFG	-1.26896	0.00099	0.78805	2138.588	1695.261
A_24_P360	NM_00672	Homo sapi	LRBA,BGL	-1.26897	0.0009	0.78804	1830.51	1450.37
A_24_P810	BC044637	Hypothetic	LOC34892	-1.26917	0.00006	0.78792	5174.931	4079.271
A_24_P255663		Unknown	A_24_P25	-1.26918	0.0001	0.78791	40166.02	31777.9

A_23_P13	NM_00140	Homo sapi	EEF1G,EF	-1.2692	3.39E-06	0.7879	119083.5	93684.12
A_24_P692030		Unknown	A_24_P69	-1.26941	0.00011	0.78777	4679.539	3682.45
A_24_P38	NM_00578	Homo sapi	THOC4,AL	-1.27017	0.00058	0.78729	2587.812	2036.313
A_24_P152792		Unknown	A_24_P15	-1.27023	0.00001	0.78726	3987.042	3139.704
A_23_P77	NM_00436	Homo sapi	CETN3,CE	-1.27033	0.00018	0.7872	1917.159	1507.484
A_23_P21	NM_03217	Homo sapi	RNUXA,PH	-1.27038	0.00002	0.78716	5450.406	4284.669
A_24_P375599		Unknown	A_24_P37	-1.27058	1.56E-06	0.78704	161643.1	127397.8
A_23_P20	NM_00398	Homo sapi	PRC1,MG	-1.2706	9.69E-06	0.78703	9701.187	7642.651
A_23_P31	NM_15264	DCP1 deca	DCP1B	-1.2709	0.00095	0.78684	2329.01	1826.802
A_32_P16	AK075484	Homo sapi	AK075484	-1.27093	4.10E-07	0.78682	3194.062	2513.952
A_23_P15	BC017032	Glucosami	GCNT3	-1.27095	0.00004	0.78681	2007.655	1576.778
A_24_P26	NM_17823	Peptidylpro	COAS2	-1.27095	0.00067	0.78681	28269.86	22333.82
A_24_P27	NM_01811	hypothetica	TMEM38B,	-1.27104	0.00011	0.78676	1933.86	1518.843
A_23_P87	NM_00517	Homo sapi	ATP5G2	-1.27126	3.83E-09	0.78662	71960.97	56604.24
A_24_P24	NM_20343	Peptidylpro	PPIA	-1.2713	1.87E-07	0.7866	33118.12	26056.61
A_24_P55	NM_00466	hect domai	HERC2,jdf	-1.2713	7.01E-06	0.7866	1911.455	1503.699
A_32_P10	AF527552	B melanom	BAGE	-1.27168	0.00065	0.78636	8008.276	6313.467
A_24_P15821		Unknown	A_24_P15	-1.2718	0.00002	0.78629	36794.95	28969.44
A_23_P15	NM_00708	Homo sapi	NUDT6,gfg	-1.27191	0.00029	0.78622	1530.445	1205.852
A_23_P25	NM_00100	Homo sapi	RPS4Y1,R	-1.27199	0.00008	0.78617	60487.07	47306.67
A_24_P32895		Unknown	A_24_P32	-1.27204	4.18E-06	0.78614	45879.09	36144.64
A_23_P16	NM_01604	splicing fac	SF3B14,P1	-1.2724	0.00005	0.78592	16540.23	13007.32
A_23_P52	NM_01488	metalloprot	PITRM1,M	-1.27244	0.00006	0.78589	1422.535	1118.239
A_32_P15	NM_00256	poly(A)-bin	PABPC1,P	-1.27274	4.65E-06	0.78571	47690.35	37584.63
A_23_P95	NM_00681	chromoson	LMAN2,GF	-1.27318	0.00077	0.78543	4751.888	3727.354
A_23_P39	NM_00685	putative nu	RY1	-1.27318	0.00016	0.78544	5646.726	4446.365
A_23_P78	NM_00143	Homo sapi	FBL,FIB,FL	-1.27339	1.27E-06	0.78531	77489.16	60927.43
A_23_P13	NM_00141	Homo sapi	EIF4A1,DD	-1.27343	3.84E-06	0.78528	62979.85	49320.66
A_23_P25	NM_02454	Homo sapi	C13orf7,FL	-1.27344	0.00003	0.78527	3771.072	2971.624
A_23_P37	AL050279	Family with	KIAA0592	-1.27355	0.00073	0.78521	3785.42	2976.523
A_24_P19	NM_00316	suppressor	SUPT4H1,	-1.27361	0.00022	0.78517	2127.962	1677.849
A_23_P70	NM_00118	aldehyde d	ALDH7A1,	-1.27384	0.00001	0.78503	4827.167	3796.48
A_23_P69	NM_00683	Homo sapi	CCNI,CYI,	-1.27391	2.17E-06	0.78499	11103.82	8734.193
A_32_P92	NM_00681	stress-indu	STIP1,HOF	-1.27405	0.00046	0.7849	1469.435	1150.302
A_32_P22	NM_01806	hypothetica	RPRC1,PA	-1.27406	0.00047	0.78489	6706.361	5230.085
A_23_P59	NM_02117	U6 snRNA-	LSM2,G7b	-1.27421	0.00001	0.7848	1623.794	1274.196
A_24_P24	AK074960	Homo sapi	AK074960	-1.27439	0.00012	0.78469	4913.409	3862.919
A_23_P74	NM_00715	Upstream (	UNR	-1.27469	8.91E-09	0.7845	9719.514	7626.619
A_23_P21	NM_00301	SET transl	SET,2PP2	-1.27473	0.00009	0.78448	3168.218	2488.246
A_24_P32	NM_00737	bromodom	BRD3,ORF	-1.275	0.00039	0.78432	1082.19	849.6443
A_23_P41	AF151783	Chromosor	C9orf88	-1.27518	0.00015	0.78421	20775.33	16276.46
A_24_P28	NM_00582	adaptor-rel	AP3S2,AP	-1.2753	0.00004	0.78413	2202.128	1724.635
A_23_P52	NM_00483	Homo sapi	GGPS1,GG	-1.2754	0.0001	0.78407	2904.287	2278.344
A_23_P41	NM_00458	Homo sapi	RPS6KA3,	-1.27548	0.00037	0.78402	1320.982	1033.09
A_32_P71	NM_00280	proteasom	PSMC6,P4	-1.27551	0.00031	0.784	17403.19	13755.71
A_23_P16	NM_00325	Homo sapi	TIAL1,TCB	-1.27553	0.00005	0.78399	2938.849	2303.056
A_24_P56	NM_17067	Homo sapi	SKP1A,OC	-1.27559	0.0002	0.78395	3881.47	3053.887
A_23_P16	NM_01835	hypothetica	FLJ11184	-1.27576	9.47E-07	0.78385	3154.926	2473.09
A_23_P25	NM_02113	Homo sapi	PPIA,CYP	-1.27622	0.00014	0.78357	67749.05	53311.51
A_23_P12	NM_02245	Homo sapi	RAB3IP,RA	-1.27634	4.02E-06	0.78349	3830.411	3001.528
A_23_P57	NM_00651	Homo sapi	SLC2A1,G	-1.27643	0.00004	0.78344	3535.068	2772.613
A_23_P88	NM_00003	aldolase A,	ALDOA,AL	-1.27686	1.15E-07	0.78317	62663.99	49157.28

A_23_P46	NM_03263	differential	PSRC1,DD	-1.27691	0.00074	0.78314	1331.33	1037.037
A_23_P11	NM_03194	Homo sapi	CENTG3,A	-1.27705	5.63E-07	0.78305	8164.895	6403.162
A_23_P17	NM_02465	Homo sapi	MORC4,ZC	-1.27719	0.00004	0.78297	2727.667	2134.742
A_32_P23	NM_03340	Homo sapi	DOCK7,KI	-1.27773	0.00064	0.78264	1663.188	1294.276
A_24_P35	AF161353	Chromosor	C9orf126	-1.27777	0.00018	0.78262	1591.053	1241.404
A_24_P865672		Unknown	A_24_P86	-1.27784	0.00001	0.78257	6404.324	5012.413
A_23_P33	NM_00269	Homo sapi	POLG,PEC	-1.27788	0.00014	0.78255	1959.443	1537.571
A_23_P33	NM_02409	XTP3-trans	XTP3TPA,	-1.27802	2.19E-09	0.78246	11380.03	8904.341
A_24_P76	NM_03121	LAT1-3TM	LAT1-3TM	-1.27808	2.71E-07	0.78242	6312.646	4946.023
A_24_P101503		Unknown	A_24_P10	-1.27811	0.0005	0.78241	2140.284	1679.912
A_23_P12	NM_00331	tetratricope	TTC3,DCR	-1.27822	0.00062	0.78234	1293.008	1019.493
A_23_P84	NM_00683	Homo sapi	COPS6,MC	-1.27848	2.05E-08	0.78218	31285.28	24449.38
A_23_P24	NM_01804	mago-nash	FLJ10292,	-1.2785	0.00025	0.78217	9460.877	7446.878
A_32_P21	NM_00097	ribosomal p	RPL6,TXR	-1.2785	3.69E-06	0.78216	119552.3	93572.18
A_24_P46	NM_17078	Homo sapi	ZNRD1,Rp	-1.27877	9.94E-09	0.782	7641.374	5977.292
A_24_P18	NM_01988	glycogen s	GSK3A,DK	-1.27901	0.00009	0.78185	8219.892	6453.192
A_24_P10	NM_19885	Homo sapi	TRIM50C	-1.27912	0.00007	0.78179	1733.078	1361.61
A_23_P84	NM_01596	Homo sapi	THAP4,CG	-1.27916	0.00002	0.78176	6162.384	4824.849
A_23_P12	NM_01463	Homo sapi	SH2BP1,C	-1.2793	3.90E-08	0.78168	3924.709	3067.984
A_23_P81	NM_01836	Homo sapi	LMBRD1,C	-1.27942	5.06E-06	0.7816	4026.166	3145.507
A_23_P35	NM_00269	Homo sapi	POLR2C,R	-1.27951	9.93E-06	0.78155	3987.901	3120.682
A_23_P28	NM_01334	GTP-bindir	PTD004	-1.27951	0.00013	0.78155	4198.435	3284.11
A_23_P34	NM_01329	Homo sapi	SAC3D1,S	-1.27955	0.00001	0.78152	6497.407	5063.841
A_23_P59	NM_02163	hypothetica	GPBP1L1,	-1.27973	2.46E-07	0.78142	2718.577	2124.314
A_24_P683734		Unknown	ENST0000	-1.27979	7.22E-09	0.78138	16626.88	13000.16
A_24_P603890		Unknown	A_24_P60	-1.27981	2.81E-07	0.78137	20208.22	15759.37
A_23_P25	NM_00236	Homo sapi	MAGEB2,D	-1.27986	8.92E-07	0.78134	4539.601	3546.458
A_23_P11	NM_01219	Homo sapi	EIF2C1,Q9	-1.27986	0.00045	0.78134	2181.053	1735.908
A_23_P43	NM_17370	Homo sapi	NM_17370	-1.27991	0.00033	0.78131	5865.804	4604.717
A_23_P10	NM_00120	Homo sapi	NM_00120	-1.27995	0.00003	0.78128	24467.33	19168.76
A_23_P10	NM_01645	hypothetica	LOC51234	-1.28008	4.62E-09	0.7812	10159.79	7938.726
A_24_P79	NM_00280	proteasom	PSMC6,P4	-1.28009	0.00005	0.78119	6345.563	4968.303
A_23_P28	NM_14473	hypothetica	PRO1853	-1.28018	2.08E-08	0.78114	3976.677	3106.313
A_23_P13	NM_19899	Homo sapi	KCTD1,C1	-1.28031	0.00022	0.78106	2163.554	1686.694
A_24_P12	AK092123	Solute carr	SLC41A1	-1.2808	0.00001	0.78076	3342.081	2607.843
A_23_P45	NM_00682	hypothetica	IFI44L,GS3	-1.28088	0.00005	0.78071	2169.903	1690.175
A_24_P36	XM_37481	Homo sapi	XM_37481	-1.2809	0.00019	0.7807	74028.27	58111.78
A_23_P25	AK074565	KIAA0310	KIAA0310	-1.28102	0.00003	0.78063	1712.752	1336.553
A_32_P96	NM_17823	Peptidylpro	COAS2	-1.28115	5.40E-07	0.78055	50052.65	39151.69
A_23_P27	NM_01804	Homo sapi	PLEKHJ1,	-1.28129	1.57E-08	0.78047	9638.943	7520.044
A_23_P80	NM_00640	Homo sapi	PROCR,C	-1.2814	7.51E-08	0.7804	8447.985	6585.108
A_24_P35	NM_01401	HSPC009 p	HSPC009	-1.28147	2.29E-06	0.78035	6757.432	5279.422
A_24_P39	NM_00141	eukaryotic	EIF4A1,DD	-1.28166	1.27E-06	0.78024	41341.61	32172.81
A_23_P10	NM_00276	Procollagen	PCOLN3	-1.28175	3.03E-06	0.78018	3397.722	2649.756
A_23_P21	NM_00474	cell cycle p	TBRG4,CF	-1.28183	0.00002	0.78014	4640.624	3611.436
A_23_P21	NM_01648	Homo sapi	PAIP2,MG	-1.28202	4.70E-09	0.78002	10424.09	8132.933
A_23_P91	NM_00288	Homo sapi	RANBP1,M	-1.28214	0.00004	0.77995	65195.64	51005.61
A_24_P20	NM_00001	acetyl-Coe	ACAT1,T2,	-1.28216	0.00067	0.77993	1500.27	1174.764
A_23_P21	NM_00396	Homo sapi	UBE1C,UB	-1.28219	0.00052	0.77991	1422.254	1116.034
A_23_P15	NM_00632	Homo sapi	RBM14,SIF	-1.28238	4.68E-08	0.7798	9923.494	7738.465
A_24_P21	NM_00340	YY1 transc	YY1,DELT	-1.28275	0.00014	0.77958	3609.565	2816.934
A_24_P110591		Unknown	A_24_P11	-1.28287	2.08E-06	0.7795	10880.23	8502.434

A_23_P15	NM_00279	Homo sapi	PSMB6,Y,I	-1.28313	5.23E-08	0.77935	55944.25	43537.39
A_23_P32	NM_13842	hypothetica	LOC11317	-1.28345	0.00013	0.77915	3384.573	2639.12
A_24_P17	NM_00166	ras homolo	RHOA,ARH	-1.28348	0.00044	0.77913	3066.471	2390.693
A_24_P38	NM_00280	proteasom	PSMC6,P4	-1.28353	1.38E-07	0.7791	18783.56	14605.39
A_24_P84808		Unknown	A_24_P84	-1.28373	5.44E-06	0.77898	76813.71	59940.08
A_24_P12	NM_03290	phosphatid	PIG-Y	-1.284	8.31E-07	0.77882	11082.13	8625.386
A_23_P34	NM_01554	Homo sapi	RPAP1,MG	-1.28417	9.40E-07	0.77871	5651.991	4405.753
A_23_P89	NM_00014	Ferrochela	FECH	-1.28447	5.50E-06	0.77853	3626.48	2823.982
A_23_P11	NM_01609	Homo sapi	BRP44L,C	-1.28462	4.14E-06	0.77844	8691.66	6782.594
A_23_P14	NM_02244	Homo sapi	TPK1,HTP	-1.28474	0.00011	0.77837	1847.051	1444.429
A_23_P32	NM_00243	Homo sapi	MNAT1,MA	-1.28483	0.00093	0.77831	3651.875	2854.773
A_23_P86	NM_00514	Homo sapi	SART1,Ara	-1.28494	0.00001	0.77825	1642.249	1277.354
A_24_P27	NM_00657	novel RGD	DCTN6,WS	-1.28497	0.00001	0.77823	3154.981	2453.677
A_24_P25	NM_03223	Ubiquitin s	USP48	-1.28506	0.00026	0.77818	2398.251	1860.667
A_23_P20	NM_01777	Homo sapi	DEPDC1,D	-1.28513	0.00009	0.77813	2663.676	2070.797
A_24_P82	NM_00619	proliferatio	PA2G4,EB	-1.28533	0.00006	0.77801	8737.691	6815.674
A_23_P10	NM_02501	Homo sapi	TUBA4,FL	-1.28544	0.00002	0.77794	26593.03	20603.02
A_23_P63	NM_00431	Homo sapi	ARL3,ARF	-1.28548	0.00014	0.77792	10569.33	8264.621
A_23_P13	NM_01553	DNA polym	DNAPTP6,	-1.28557	0.00003	0.77786	6577.258	5127.749
A_23_P15	NM_02466	hypothetica	VPS37B,FI	-1.28571	0.00067	0.77778	1294.15	1001.825
A_23_P36	NM_01327	Homo sapi	ANKRD11,	-1.28605	6.06E-07	0.77757	6125.864	4757.166
A_23_P59	NM_01344	Homo sapi	MKRN1,RN	-1.28605	2.40E-06	0.77757	8976.197	6988.363
A_32_P20	AB002374	KIAA0376	KIAA0376	-1.28624	0.00046	0.77746	2544.786	1970.851
A_23_P25	NM_01832	Homo sapi	BXDC2,BR	-1.28624	9.74E-08	0.77746	10616.19	8264.57
A_24_P33607		Unknown	A_24_P33	-1.2863	4.64E-07	0.77742	99321.7	77379.95
A_23_P62	NM_01497	Homo sapi	KIFAP3,KA	-1.28643	0.00017	0.77734	4314.055	3356.263
A_24_P20	BC015780	KIAA0323	KIAA0323	-1.28644	0.00019	0.77734	1309.529	1015.633
A_24_P370484		Unknown	A_24_P37	-1.28661	0.00062	0.77723	1466.148	1144.721
A_23_P14	NM_00717	protein-O-r	POMT1,RT	-1.28662	9.82E-06	0.77723	3645.22	2830.445
A_32_P68	BC036448	Hypothetic	FLJ13305	-1.2868	0.00063	0.77712	1573.274	1232.362
A_23_P42	AL080183	Similar to K	LOC38768	-1.28688	0.0001	0.77707	2393.046	1858.79
A_32_P88	NM_03226	hypothetica	AE2,DKFZ	-1.28691	0.00016	0.77705	1453.724	1133.203
A_23_P75	NM_00629	Homo sapi	TSG101,TS	-1.28695	0.00003	0.77703	12121.62	9447.754
A_24_P118271		Unknown	A_24_P11	-1.28703	0.00041	0.77698	5799.887	4515.557
A_23_P16	NM_02182	hypothetica	MDS025,M	-1.28717	0.00007	0.7769	3719.601	2893.025
A_24_P135242		similar to 6	ENST0000	-1.28802	6.18E-08	0.77638	133121.3	103463.9
A_23_P71	NM_00683	Homo sapi	COPS5,CS	-1.28841	0.00021	0.77615	11595.57	9054.408
A_23_P40	NM_01815	hypothetica	SLC25A36	-1.28842	4.39E-06	0.77615	2513.593	1950.904
A_24_P854913		AF096870	THC18692	-1.28843	0.00019	0.77614	1476.975	1141.978
A_32_P10	NM_00120	basic trans	BTF3,NAC	-1.2885	9.40E-07	0.7761	22254.56	17306.96
A_32_P83	XM_20843	similar to T	LOC28649	-1.28855	3.12E-06	0.77606	2664.249	2066.631
A_23_P41	AK000282	Formin bin	FNBP1L	-1.28856	0.00093	0.77606	1596.729	1232.411
A_32_P10	NM_02123	TERA prot	FAM60A,L	-1.28878	0.00097	0.77593	1294.788	999.0314
A_23_P25	NM_03294	Homo sapi	RECQL,Re	-1.28879	0.00001	0.77592	1768.895	1373.64
A_23_P42	NM_14501	hypothetica	FLJ25416,	-1.28881	5.92E-06	0.77591	1774.522	1377.952
A_23_P10	NM_01554	Homo sapi	LRIG1,LIG	-1.28902	0.00012	0.77578	2248.188	1750.183
A_32_P64990		Unknown	A_32_P64	-1.28928	0.00048	0.77563	1139.596	881.5409
A_23_P34	NM_01630	pp21 homo	WBP5	-1.28928	0.00019	0.77563	5380.062	4193.639
A_23_P59	NM_00144	fatty acid b	FABP5,EF	-1.28936	3.04E-09	0.77558	43706.02	33888.87
A_23_P15	NM_03233	Homo sapi	CAPNS2,M	-1.28938	0.00039	0.77556	1230.205	951.8648
A_23_P258108		Unknown	A_23_P25	-1.28945	0.00025	0.77552	188186.9	146934
A_24_P367109		Unknown	A_24_P36	-1.28946	0.00002	0.77552	2346.83	1820.011

A_24_P195714	Unknown	A_24_P19	-1.28953	2.37E-07	0.77547	18122.47	14030.17
A_23_P31	NM_00328	Homo sapi	TPD52L1,D	-1.28985	0.00016	0.77529	12279.46	9546.504
A_23_P12	NM_02233	Homo sapi	C11orf24,D	-1.29001	1.24E-06	0.77519	7361.228	5699.833
A_23_P15	NM_00458	Homo sapi	SCO1,SCO	-1.29016	5.41E-08	0.7751	4306.973	3336.777
A_24_P55	NM_01606	Chromosor	C6orf74	-1.29024	0.00002	0.77505	4359.813	3372.73
A_23_P10	NM_02018	DC13 prote	DC13,2310	-1.29028	0.00001	0.77503	9710.63	7541.033
A_32_P25809	Unknown	A_32_P25	-1.29031	2.23E-09	0.77501	111485.6	86351.05
A_32_P21	NM_00229	laminin rec	RPSA,LRP	-1.29034	3.25E-06	0.77499	90495.45	70350.57
A_23_P35	NM_00408	Homo sapi	TIMM8A,D	-1.29043	2.12E-06	0.77494	2030.293	1573.626
A_24_P37	NM_18279	Homo sapi	TMSL4	-1.29058	2.74E-08	0.77484	23399.17	18112.14
A_23_P14	NM_00375	eukaryotic	EIF3S5,eIF	-1.29064	7.19E-07	0.77481	3894.644	3013.848
A_32_P12	NM_01476	KIAA0174	KIAA0174,	-1.29065	8.80E-07	0.7748	6642.87	5153.799
A_24_P16071	Unknown	A_24_P16	-1.29089	9.26E-07	0.77466	2202.22	1705.41
A_32_P18	NM_02393	hypothetica	FUNDC2,D	-1.2909	7.91E-06	0.77465	5883.262	4567.63
A_23_P74	NM_00660	nuclear dis	NUDC,HNU	-1.29091	0.00001	0.77465	6969.883	5375.934
A_23_P11	NM_01541	Homo sapi	BRP44,MG	-1.29115	4.11E-06	0.7745	15614.58	12126.63
A_24_P478940	Unknown	A_24_P47	-1.29118	0.0002	0.77449	104665.5	81621.13
A_23_P88	NM_00005	Homo sapi	BLM,BS,RI	-1.29124	0.00001	0.77445	3440.553	2664.476
A_23_P95	NM_00276	Homo sapi	PRPS1,PR	-1.29127	0.00002	0.77443	10933.48	8483.482
A_23_P34	NM_02307	hypothetica	C1orf163,F	-1.29127	3.26E-06	0.77443	2200.385	1703.222
A_23_P50	NM_03238	hypothetica	COG8,DOI	-1.29131	0.00005	0.77441	3451.667	2675.529
A_23_P13	XM_06088	similar to p	LOC12819	-1.29145	0.00006	0.77432	42820.67	33278.98
A_32_P26	AB046820	KIAA1600	KIAA1600	-1.29163	2.15E-06	0.77422	3807.626	2947.618
A_23_P87	NM_01418	Homo sapi	COMMD9,	-1.29168	0.00097	0.77419	1691.971	1301.336
A_24_P19	XM_37298	Homo sapi	XM_37298	-1.29183	0.00059	0.7741	52876.5	40880.38
A_23_P38	NM_01818	Homo sapi	ARL10C,G	-1.29185	0.00046	0.77408	7279.674	5632.791
A_24_P34	NM_01761	Homo sapi	RNPC3,RN	-1.29188	0.00003	0.77406	1575.764	1222.704
A_23_P61	NM_02412	hypothetica	MGC4825	-1.29195	1.67E-07	0.77402	6874.996	5317.959
A_23_P79	NM_17078	Homo sapi	MKKS,KMS	-1.29196	0.00006	0.77402	1926.162	1489.636
A_24_P341376	Unknown	A_24_P34	-1.29223	2.65E-06	0.77386	43575.19	33817.15
A_24_P247464	Unknown	A_24_P24	-1.2923	4.40E-06	0.77381	1634.863	1264.424
A_23_P20	AB043587	Acyl-Coenz	ACBD3	-1.29243	0.00003	0.77373	3885.973	3011.088
A_23_P15	NM_02213	golgi phosp	GOLPH3,G	-1.29247	5.71E-07	0.77371	2485.634	1922.808
A_24_P587938	Unknown	A_24_P58	-1.29279	0.00018	0.77352	112972.9	86817.65
A_24_P75	XM_37330	Homo sapi	XM_37330	-1.29287	2.93E-06	0.77347	2131.2	1648.886
A_24_P33	NM_02017	Nicalin hon	NCLN	-1.29292	0.00002	0.77344	9351.499	7260.664
A_24_P316074	Unknown	A_24_P31	-1.293	0.00001	0.77339	182919.3	142003.9
A_32_P80	BC063128	C-type lect	OCIL	-1.29311	0.00002	0.77333	16933.82	13004.35
A_23_P29	NM_00175	carbonyl re	CBR1,CBR	-1.29326	0.00084	0.77324	7408.659	5729.636
A_32_P15	NM_00097	ribosomal p	RPL7,huml	-1.29332	2.80E-06	0.77321	188693.9	146322.5
A_23_P44	NM_00425	Homo sapi	COX5A,VA	-1.29333	2.82E-08	0.7732	22867.99	17706.19
A_24_P127042	Unknown	A_24_P12	-1.29356	0.00035	0.77306	5506.52	4269.73
A_23_P11	NM_00576	Homo sapi	CEBPZ,CE	-1.29356	0.00009	0.77306	7329.43	5694.495
A_24_P916228	Unknown	A_24_P91	-1.29366	3.40E-07	0.773	4477.032	3457.319
A_23_P70	NM_00002	Homo sapi	AGA,GA,A	-1.29368	0.0005	0.77299	2041.684	1573.732
A_23_P12	NM_01499	Homo sapi	RAB21,KIA	-1.29373	0.00036	0.77296	7789.663	6042.087
A_23_P48	XM_29208	similar to p	LOC34145	-1.29397	0.00002	0.77281	52483.42	40715.74
A_23_P17	NM_03235	hypothetica	MGC12981	-1.294	0.00024	0.7728	4594.17	3553.054
A_24_P323778	Unknown	A_24_P32	-1.29404	3.92E-06	0.77277	51771.52	40131.09
A_32_P83	NM_08062	Homo sapi	BRI3BP,KC	-1.29428	1.47E-06	0.77263	2313.745	1788.115
A_32_P53	BC040307	CDNA clon	BC040307	-1.29428	0.0005	0.77263	1222.741	949.1898
A_32_P11	NM_00334	ubiquitin-co	UBE2N,UB	-1.29433	0.00095	0.7726	11921.57	9276.302

A_23_P51	NM_00288	RAP1A, m	RAP1A,RA	-1.29445	0.00008	0.77253	1787.316	1377.939
A_24_P23	XM_37744	similar to p	LOC40185	-1.29455	2.07E-06	0.77247	58366.25	45243.16
A_24_P530977		Unknown	A_24_P53	-1.29476	7.35E-06	0.77234	38165.47	29375.82
A_23_P12	NM_00038	Homo sapi	ALDH3A2,	-1.29483	0.00036	0.7723	1763.174	1370.247
A_24_P20	NM_17823	Peptidylpro	COAS2	-1.29487	9.25E-07	0.77228	53654.38	41570.41
A_24_P72	NM_00309	small nucle	SNRPB2,M	-1.29504	9.57E-07	0.77218	3644.042	2812.844
A_24_P22746		Unknown	THC18471	-1.2951	0.00029	0.77214	1137.229	882.4555
A_23_P50	NM_01549	Homo sapi	ANKRD25,	-1.29533	9.80E-07	0.772	8768.34	6770.411
A_24_P24890		Unknown	A_24_P24	-1.29536	8.57E-06	0.77199	5196.36	4006.319
A_23_P12	NM_00386	Homo sapi	SAP30	-1.29537	0.0004	0.77198	4662.392	3597.589
A_24_P34	NM_00192	dCMP dea	DCTD,MG	-1.29542	0.00031	0.77195	2068.054	1599.5
A_23_P13	NM_00366	Homo sapi	PIR	-1.29562	0.00004	0.77183	23979.83	18529.55
A_23_P11	NM_00280	Homo sapi	PSMB9,LM	-1.29578	2.26E-06	0.77174	4791.963	3690.453
A_32_P43	NM_15382	Proteasom	PSMD4	-1.29586	4.80E-08	0.77169	38227.52	29452.84
A_23_P13	NM_00680	Homo sapi	SEC61B	-1.29603	9.18E-08	0.77159	54849.82	42391.99
A_24_P25	X64990	Olfactory r	OR5L2	-1.29604	0.00008	0.77158	3597.729	2766.504
A_23_P25	NM_13841	Homo sapi	CKLFSF7,I	-1.29634	1.07E-06	0.77141	4643.484	3584.177
A_24_P11	NM_00200	farnesyl di	FDPS,FPS	-1.29637	0.00002	0.77139	25031.12	19399.09
A_23_P33	NM_00675	syntrophin,	SNTB2,SN	-1.29637	0.00021	0.77138	2596.562	2003.001
A_23_P26	NM_00637	membrane	VAT1,VAT	-1.29639	4.04E-06	0.77137	13208.84	10212.43
A_23_P25	NM_01324	Homo sapi	NXT1,P15,	-1.29671	0.00023	0.77118	13406.15	10404.63
A_32_P11	NM_00012	electron-tra	ETFA,EMA	-1.29676	2.35E-06	0.77116	2946.979	2271.51
A_23_P10	NM_00191	Homo sapi	CYB5,MCE	-1.29681	1.96E-06	0.77112	12987.82	10049.85
A_32_P26443		AF010144	THC19676	-1.29682	4.17E-06	0.77112	1853.226	1428.045
A_24_P174824		ALU4_HUM	THC19983	-1.29683	5.21E-07	0.77111	2392.18	1844.574
A_24_P10	NM_14586	Homo sapi	ANXA11,A	-1.29692	0.00003	0.77106	1389.443	1069.9
A_23_P11	D14041	Homo sapi	pKBF-2	-1.29704	0.00025	0.77099	2787.57	2188.307
A_23_P12	NM_02196	Homo sapi	NAP1L2,BI	-1.29716	0.00007	0.77092	1919.815	1475.893
A_23_P25	NM_01648	hypothetica	C6orf203,F	-1.29719	0.00041	0.7709	1263.156	972.1613
A_23_P36	NM_00681	coated ves	TMED2,P2	-1.29747	1.26E-07	0.77073	4890.678	3770.438
A_23_P16	NM_03184	heterogene	HNRPU,SA	-1.29755	1.20E-07	0.77069	36487.13	28177.95
A_23_P14	NM_01215	Homo sapi	FBXL3,FBI	-1.29758	0.00009	0.77067	1506.927	1157.505
A_23_P50	NM_00318	Homo sapi	MAP3K7,T	-1.29769	0.00009	0.7706	2026.989	1560.926
A_23_P211436		S3A1_HUM	THC19887	-1.29775	0.00016	0.77056	4725.479	3630.447
A_23_P10	NM_01812	hypothetica	FLJ10534,I	-1.29802	0.00009	0.7704	2564.24	1977.336
A_23_P24	AK023921	Tetratricop	TTC12	-1.29812	8.91E-07	0.77034	2259	1738.992
A_23_P10	NM_00279	Homo sapi	PSMB3,HC	-1.29852	8.76E-07	0.77011	10103.06	7794.087
A_23_P69	NM_00096	ribosomal p	RPL4,HRP	-1.29873	0.00002	0.76999	64526.11	49918.34
A_23_P20	NM_01220	Homo sapi	GTF3C4,TI	-1.2989	4.84E-08	0.76988	3225.9	2484.956
A_24_P23	NM_02274	hypothetica	MOSC1,FL	-1.29899	0.00017	0.76983	1412.672	1091.879
A_23_P12	NM_00513	Homo sapi	ANXA3,AN	-1.29905	0.0001	0.76979	3411.511	2630.412
A_23_P21	NM_01822	hypothetica	FLJ10803	-1.29943	1.69E-06	0.76957	2887.809	2225.558
A_23_P21	NM_03271	Homo sapi	C14orf151,	-1.29962	0.00005	0.76946	2442.217	1878.55
A_23_P36	NM_01809	Homo sapi	NETO2,NE	-1.29964	0.00089	0.76944	6862.829	5300.4
A_24_P218509		Unknown	THC19441	-1.30001	0.00002	0.76922	1672.389	1290.665
A_23_P93	NM_00005	Homo sapi	BCKDHB,E	-1.30006	0.0009	0.7692	1311.479	1002.15
A_23_P36	NM_00365	Homo sapi	AGPS,ADA	-1.30016	0.00003	0.76914	1851.757	1421.517
A_32_P54	NM_00600	ubiquinol-c	UQCRH	-1.30033	0.00006	0.76904	110064	85035.06
A_23_P25	NM_00533	Homo sapi	HCCS,CCH	-1.30065	1.54E-07	0.76884	6451.917	4953.469
A_32_P69	NM_19427	protein con	MOPT,BLC	-1.30129	7.62E-06	0.76847	3753.976	2888.979
A_23_P40	NM_00415	Ornithine d	OAZ1	-1.30135	0.00023	0.76843	111870.6	86623.35
A_23_P15	NM_01224	Homo sapi	SEPHS1,S	-1.30138	0.00017	0.76842	8254.671	6357.665

A_23_P42	NM_00428	Homo sapi	NFE2L3,NI	-1.30153	0.00018	0.76833	1791.135	1371.718
A_24_P34	XM_37106	similar to ri	LOC38840	-1.30155	9.36E-10	0.76832	132390.1	101763.5
A_23_P43	NM_01504	Homo sapi	KIAA0090,	-1.30161	0.00002	0.76828	1874.443	1437.201
A_23_P92	NM_01839	Homo sapi	ABHD10,F	-1.30162	0.00063	0.76827	1652.924	1279.555
A_24_P40	BG206642	Similar to F	BG206642	-1.30168	0.00071	0.76824	1402.916	1084.546
A_24_P76	NM_00140	eukaryotic	EEF1A1,EI	-1.3018	4.34E-07	0.76817	211905.2	163055.9
A_23_P62	NM_01868	Homo sapi	KIAA1166,	-1.30186	0.00015	0.76813	1600.015	1225.061
A_23_P16	NM_14564	Homo sapi	MRPL35	-1.30188	0.00003	0.76812	5634.006	4337.181
A_23_P25	NM_13841	Homo sapi	FAM54A,D	-1.3019	0.00035	0.76811	1222.105	934.6263
A_32_P99	NM_00227	karyopheri	TNPO1,MI	-1.30193	8.63E-07	0.76809	16504.65	12681.98
A_23_P21	NM_02015	Homo sapi	SMARCAD	-1.302	3.45E-14	0.76805	1235.475	951.277
A_23_P37	AF030162	Translocas	TIMM23	-1.30208	0.00011	0.768	5542.721	4272.803
A_23_P35	NM_01532	KIAA0409	KIAA0409	-1.30212	3.34E-06	0.76798	10041.87	7728.615
A_23_P20	NM_05293	pro-oncosis	PORIMIN,I	-1.30216	0.00014	0.76795	2027.264	1562.757
A_32_P140262		Unknown	A_32_P14	-1.30218	0.00002	0.76794	4261.193	3274.29
A_23_P11	NM_13924	methionyl-t	MTFMT,FM	-1.30225	0.0001	0.7679	1216.187	936.5559
A_23_P15	NM_02162	Homo sapi	SCPEP1,R	-1.30242	0.00001	0.7678	3504.21	2693.348
A_23_P16	NM_01567	Homo sapi	TRUB2,CL	-1.30264	0.00008	0.76767	4876.696	3752.852
A_23_P13	NM_00272	Homo sapi	PREP,PE,I	-1.30316	0.00008	0.76737	3736.398	2865.183
A_23_P15	NM_00533	Huntingtin	HIP2	-1.30321	0.00011	0.76733	2061.738	1577.196
A_23_P70	AK000420	Homo sapi	FLJ20413	-1.30343	0.00009	0.76721	2902.19	2220.941
A_24_P707543		Unknown	A_24_P70	-1.3036	0.00033	0.7671	88457.67	68220.05
A_23_P15	NM_00737	Homo sapi	DDX42,RH	-1.30383	7.45E-06	0.76697	3042.694	2336.379
A_23_P36	NM_00214	Homo sapi	HOXB5,HC	-1.30421	0.00002	0.76675	1680.084	1291.958
A_24_P367191		Unknown	A_24_P36	-1.30439	2.12E-08	0.76664	99345.14	76290.94
A_24_P59	XM_37244	similar to e	LOC39028	-1.30447	2.10E-08	0.7666	9777.865	7501.408
A_23_P38	AK025703	HCV F-trar	LOC40115	-1.30447	0.00081	0.76659	1869.681	1454.22
A_23_P416477		Unknown	ENST0000	-1.30477	1.35E-06	0.76642	2393.033	1832.47
A_23_P30	NM_03334	Homo sapi	TRIM7,GN	-1.30483	7.04E-07	0.76638	3353.955	2572.201
A_23_P38	AK055693	Zinc finger	FLJ35036	-1.30486	0.00002	0.76636	3122.861	2395.145
A_23_P43	NM_00030	Homo sapi	PMM2,CD	-1.30488	4.16E-06	0.76635	2493.087	1911.215
A_23_P43	NM_00175	calpastatin	CAST,BS-	-1.30524	0.00059	0.76614	8065.615	6190.991
A_24_P28	NM_17586	Homo sapi	ELYS	-1.30536	0.00023	0.76607	2533.102	1950.607
A_23_P61	NM_03298	Sec23 (S.	SEC23B	-1.30538	0.00001	0.76606	1678.235	1282.458
A_23_P27	NM_00293	ribonucleas	RNASEH1	-1.30568	1.01E-07	0.76589	3630.334	2782.675
A_23_P51	NM_15333	Homo sapi	PUSL1,FL	-1.30568	0.0001	0.76588	5062.417	3890.438
A_24_P27	NM_01871	calreticulin	CDCA7L,R	-1.3057	0.00008	0.76587	5655.277	4325.077
A_24_P13	NM_00429	ring finger	RNF14,AR	-1.30576	0.00005	0.76584	1585.272	1220.008
A_32_P14	NM_01595	CGI-27 pro	C2orf4,CG	-1.30584	0.00055	0.76579	1689.231	1284.522
A_23_P16	AB002324	Zinc finger	ZNF629	-1.30585	5.28E-06	0.76578	6493.989	4980.935
A_23_P13	NM_00421	Homo sapi	FIBP,FGFI	-1.30613	8.79E-09	0.76562	3965.209	3036.207
A_24_P38	NM_01631	polymeras	POLR3K,C	-1.30624	2.88E-08	0.76556	14223.51	10879.63
A_23_P15	NM_01461	KIAA0182	KIAA0182	-1.30626	0.00028	0.76554	1622.974	1235.818
A_24_P41	NM_01630	60S riboso	C15orf15,L	-1.30629	0.00015	0.76553	2424.637	1855.925
A_24_P92	BC042363	Activating t	ATF7	-1.30644	0.00001	0.76544	1439.5	1102.474
A_24_P50	NM_00229	laminin rec	RPSA,LRP	-1.30659	3.99E-07	0.76535	189376.9	145176.5
A_23_P53	NM_03327	Ku70-bindi	KUB3	-1.30678	3.51E-07	0.76524	2289.96	1753.225
A_32_P84	NM_17881	Homo sapi	CKLFSF4	-1.30686	9.12E-06	0.76519	6969.1	5365.303
A_24_P42	NM_01632	RAB14, me	RAB14,FB	-1.30688	0.00002	0.76518	2334.959	1785.813
A_24_P67748		Unknown	A_24_P67	-1.30688	0.00048	0.76518	1835.375	1396.094
A_23_P66	NM_02016	Homo sapi	LXN,ECI,T	-1.30702	2.69E-06	0.7651	6721.485	5156.735
A_32_P22	NM_00096	ribosomal	RPL5,MST	-1.30718	1.74E-07	0.76501	158834.3	121844.6

A_23_P25	NM_02457	Homo sapi	C6orf211,F	-1.30719	5.71E-07	0.765	2537.976	1942.391
A_32_P88	BC016882	Chromosor	C22orf9	-1.30722	2.43E-07	0.76498	7994.11	6116.959
A_23_P21	NM_01224	sirtuin (sile	SIRT5,SIR	-1.30728	8.64E-06	0.76495	2023.233	1544.519
A_24_P16	NM_03121	LAT1-3TM	LAT1-3TM	-1.30731	6.01E-06	0.76493	4380.285	3353.264
A_23_P25	NM_02462	Homo sapi	MLF1IP,KL	-1.30736	0.001	0.7649	4785.007	3669.833
A_23_P86	NM_02076	retinoblast	RBAF600,I	-1.3074	2.46E-06	0.76488	2397.704	1835.47
A_24_P914088		Unknown	A_24_P91	-1.30751	0.00006	0.76481	47586.45	36604.99
A_23_P21	AK094968	Interferon,	IFI16	-1.30767	4.89E-06	0.76472	5995.484	4587.404
A_23_P21	NM_00630	Homo sapi	SMC1L1,S	-1.3077	3.52E-07	0.7647	3906.472	2987.9
A_23_P70	NM_00066	Homo sapi	RPL9,FLJ2	-1.30772	1.91E-06	0.76469	239258.1	183301
A_23_P21	NM_00229	lysosomal-	LAMP2,LA	-1.30781	0.00003	0.76464	1646.865	1254.77
A_24_P340866		Unknown	A_24_P34	-1.308	7.21E-06	0.76453	2190.334	1678.996
A_23_P25	NM_00078	Homo sapi	CYP51A1,I	-1.30824	0.00023	0.76439	1315.261	1002.729
A_23_P24	NM_00202	Homo sapi	FNTA,FPT	-1.30829	0.0009	0.76435	4487.428	3454.629
A_32_P14	BC046178	Chromosor	C6orf173	-1.30836	4.34E-06	0.76432	7386.418	5665.479
A_23_P68	NM_00587	translation	GC20	-1.30845	0.00023	0.76426	9473.932	7185.787
A_23_P11	NM_03215	MBLR prot	PCGF6,ME	-1.30847	0.00004	0.76425	1924.051	1473.782
A_24_P11	NM_00043	platelet-act	PAFAH1B1	-1.30852	0.0009	0.76422	2350.884	1800.087
A_23_P16	AL390151	Zinc finger	ZBTB7	-1.30879	0.00004	0.76406	2850.254	2170.545
A_23_P14	NM_02457	hypothetica	FLJ11712	-1.30895	9.26E-08	0.76397	2583.591	1973.456
A_23_P91	NM_14471	Homo sapi	FNDC6,MC	-1.30898	1.90E-07	0.76395	2655.621	2029.005
A_24_P21	NM_02094	KIAA1609	KIAA1609,	-1.30925	0.00057	0.76379	1543.486	1168.835
A_23_P16	NM_03230	Homo sapi	HSDL2,C9	-1.30935	0.00002	0.76374	2602.112	1991.96
A_32_P61	NM_17823	Peptidylprc	COAS2	-1.30956	3.62E-09	0.76361	75516.13	57719.95
A_24_P35	BC022779	V-akt murir	AKT2	-1.30974	0.00001	0.76351	14189.88	10870.83
A_23_P35	NM_13834	hypothetica	KIAA2013,	-1.30978	1.70E-09	0.76349	4095.695	3126.407
A_23_P54	NM_02461	Homo sapi	NARG2	-1.30997	7.43E-06	0.76337	2515.472	1919.176
A_24_P12	NM_01606	mitochondr	MRPS16,R	-1.31004	1.92E-07	0.76334	12448.04	9475.219
A_23_P73	NM_02515	Homo sapi	RUFY1,ZF	-1.31008	0.00015	0.76331	4779.284	3653.519
A_23_P93	NM_00353	Homo sapi	HIST1H3J,	-1.31012	4.23E-11	0.76329	25592.77	19533.8
A_32_P89	NM_18258	Homo sapi	NM_18258	-1.3102	2.20E-06	0.76324	1849.678	1411.603
A_24_P37	NM_00449	hepatoma-	HDGF,HM	-1.31021	8.51E-06	0.76323	18229.43	13959.53
A_32_P15	NM_02078	Homo sapi	ZNF398,P5	-1.31022	1.42E-07	0.76323	2783.062	2123.24
A_24_P20	XM_29431	similar to H	LOC34652	-1.31035	7.12E-07	0.76316	72748.02	55720.41
A_23_P13	NM_01838	hypothetica	FLJ11301	-1.31049	0.00006	0.76307	1396.729	1064.194
A_23_P68	NM_02194	Homo sapi	NM_02194	-1.31056	0.00007	0.76303	4218.49	3232.544
A_32_P44	NM_15329	Homo sapi	FAM10A4	-1.31065	0.00066	0.76298	6395.799	4887.515
A_23_P63	NM_00720	Homo sapi	DDX20,DP	-1.31069	0.00004	0.76296	2738.628	2091.628
A_23_P70	AK027125	Hypothetic	ENST0000	-1.31092	0.00007	0.76282	2106.75	1615.162
A_24_P786257		Unknown	A_24_P78	-1.31101	0.00006	0.76277	1116.051	851.9383
A_32_P10	NM_00257	p21 (CDKN	PAK2,PAK	-1.31106	0.00019	0.76274	1911.537	1462.88
A_23_P47	AK000854	Vacuolar p	VPS37C	-1.31148	0.00001	0.7625	2431.473	1851.745
A_32_P32315		Unknown	A_32_P32	-1.31153	0.00075	0.76247	1388.417	1066.648
A_23_P25	NM_00018	Homo sapi	HGD,AKU,	-1.31159	3.88E-07	0.76243	8686.843	6640.632
A_23_P14432		Unknown	A_23_P14	-1.31168	0.00037	0.76238	1573.584	1192.331
A_23_P13	NM_00501	Homo sapi	NUCB2,NE	-1.31188	0.00003	0.76227	2820.855	2148.704
A_23_P42	NM_01846	Homo sapi	C3orf10,MI	-1.31206	0.00003	0.76216	30314.89	23236.57
A_24_P73	BC013276	Homo sapi	BC013276	-1.31213	0.00008	0.76212	1106.038	843.9183
A_23_P66	NM_00125	Homo sapi	CDC27,AP	-1.31213	0.00002	0.76212	2679.93	2044.634
A_23_P50	NM_15318	Homo sapi	ANKRD15,	-1.31215	0.00001	0.76211	3143.832	2385.164
A_23_P12	NM_01900	Spindlin fa	SPIN2	-1.31217	1.81E-06	0.7621	1975.519	1507.16
A_24_P16	NM_00682	peroxisoma	ACOT2,Mt	-1.31221	0.00003	0.76207	8957.011	6860.803

A_23_P48	NM_00474	Homo sapi	CCPG1,CF	-1.31246	4.28E-12	0.76193	4028.3	3072.181
A_23_P16	BC001097	Eukaryotic	EIF4B	-1.31259	3.59E-07	0.76185	25200.95	19253.37
A_32_P20	NM_01609	ribosomal	RPL26L1,F	-1.31267	0.00013	0.7618	17742.57	13506.42
A_23_P47	NM_00315	Homo sapi	STAT6,ST	-1.31269	9.77E-06	0.7618	3945.905	3005.038
A_23_P77	NM_02451	Homo sapi	C16orf53,M	-1.31281	0.00001	0.76173	21902.61	16768.8
A_23_P63	NM_00037	uroporphyr	UROD,PC	-1.31282	0.00001	0.76172	8194.544	6210.043
A_24_P859431		Unknown	A_24_P85	-1.31292	3.56E-08	0.76166	2910.707	2216.733
A_23_P43	AF358454	Homo sapi	AF358454	-1.31299	1.63E-10	0.76162	5956.706	4536.925
A_23_P12	NM_01500	SMART/HD	SPEN,MIN	-1.31333	3.91E-06	0.76143	1724.974	1310.24
A_23_P20	NM_00480	Homo sapi	POLR2D,R	-1.31351	2.98E-08	0.76132	3509.343	2671.264
A_32_P13	U63541	Tuftelin inte	TFIP11	-1.31366	2.27E-07	0.76123	1870.715	1424.34
A_24_P11	NM_03329	T-cell activ	MRFAP1,F	-1.31389	5.74E-06	0.7611	2375.443	1807.075
A_32_P16	XM_37171	similar to a	LOC38924	-1.31395	4.77E-07	0.76106	153165.8	116908.1
A_23_P25	NM_01664	mitochondr	MRPS30,P	-1.31418	0.00014	0.76093	2049.971	1552.461
A_23_P12	NM_00454	Homo sapi	NDUFA10,	-1.31426	0.00002	0.76089	2131.602	1620.185
A_24_P247014		Unknown	A_24_P24	-1.31459	1.87E-08	0.7607	136105.2	103757.2
A_24_P22	AK092040	Zinc finger,	ZCSL2	-1.31499	0.00008	0.76046	1130.119	860.7948
A_23_P23	NM_17380	Homo sapi	BLOC1S2,	-1.31506	1.63E-09	0.76042	9187.129	6980.765
A_23_P31	NM_00691	Homo sapi	RIT1,RIT,R	-1.31526	0.00011	0.7603	4982.666	3802.382
A_23_P14	NM_01570	Homo sapi	HIRIP5,Nfu	-1.31534	0.00001	0.76026	6429.438	4914.857
A_23_P53	NM_01646	Homo sapi	C14orf112,	-1.31542	1.59E-06	0.76021	18708.01	14160.21
A_23_P15	NM_00588	Homo sapi	DLEU1,BC	-1.31561	0.00019	0.76011	8543.554	6514.665
A_23_P55	NM_01531	pumilio (Dr	PUM2,PUM	-1.31591	0.00001	0.75993	1456.887	1104.679
A_23_P20	NM_03228	Homo sapi	PRPF38A,	-1.31596	1.02E-07	0.7599	2665.818	2029.053
A_32_P10	NM_15225	leucine-rich	MGC45866	-1.31617	4.29E-10	0.75978	5948.581	4521.892
A_24_P57	AL831944	Homo sapi	AL831944	-1.31626	0.00012	0.75973	1623.095	1240.537
A_32_P68	NM_00117	ADP-ribosy	ARL1,ARF	-1.31648	3.89E-06	0.7596	3260.805	2474.443
A_24_P92	NM_00681	chromoson	LMAN2,GF	-1.31658	0.0002	0.75955	3851.26	2907.87
A_23_P15	AK054969	Ubiquitin p	UBR1	-1.31674	0.0002	0.75945	2251.814	1707.821
A_23_P89	NM_00376	beclin 1 (co	BECN1,AT	-1.3168	9.90E-08	0.75942	4447.016	3377.64
A_23_P89	NM_01496	Homo sapi	EPN2,EHB	-1.31693	0.00057	0.75934	3504.732	2657.499
A_23_P13	NM_00473	eukaryotic	ETF1,ERF,	-1.31697	1.06E-06	0.75932	3960.032	3009.546
A_23_P20	NM_00631	Homo sapi	NCOR2,SM	-1.31718	2.01E-07	0.7592	21652.36	16384.06
A_23_P20	NM_01682	Homo sapi	ADD3,ADD	-1.31721	0.00084	0.75918	2299.697	1730.677
A_32_P2213		Unknown	A_32_P22	-1.31761	0.00003	0.75895	1533.095	1167.446
A_23_P21	NM_01221	Homo sapi	ILF3,DRBF	-1.31763	6.96E-08	0.75894	6837.479	5194.855
A_23_P31	NM_13879	senescenc	LEO1,RDL	-1.31764	0.00003	0.75893	5440.479	4143.831
A_23_P82	NM_01241	Homo sapi	RAD54B	-1.31768	6.48E-14	0.75891	2018.533	1536.623
A_24_P92771		Unknown	A_24_P92	-1.31777	6.83E-06	0.75886	224694.9	171604.4
A_23_P15	NM_00629	X-ray repai	XRCC1,RC	-1.318	1.86E-06	0.75872	3213.156	2434.405
A_23_P20	NM_01593	Homo sapi	KIAA0859,	-1.31802	1.75E-06	0.75871	8877.057	6741.627
A_23_P31	AL080183	Similar to k	LOC38768	-1.3182	4.08E-07	0.75861	2344.857	1778.717
A_23_P21	NM_00163	Homo sapi	AMD1,AMD	-1.31822	5.69E-10	0.7586	8797.155	6674.034
A_23_P14	AF178574	MSTP146	MST146	-1.31842	8.42E-06	0.75848	1935.951	1473.174
A_24_P25	NM_01653	apoptosis-r	FLJ39616	-1.31848	0.00017	0.75845	1397.564	1054.808
A_23_P12	NM_00483	Homo sapi	GSTO1,P2	-1.31856	3.09E-09	0.7584	24472.89	18533.75
A_23_P21	NM_01772	leucine rich	LRRFIP2,H	-1.31867	0.00011	0.75834	1286.217	977.9694
A_23_P29	NM_00607	TRK-fused	TFG,TF6,T	-1.31869	1.13E-06	0.75833	11189.94	8507.345
A_23_P19	NM_00654	IGF-II mRN	IMP-3,KOC	-1.31874	2.28E-07	0.7583	3789.813	2874.633
A_23_P39	NM_18247	Pyruvate ki	PKM2	-1.31878	5.06E-10	0.75828	16343.93	12383.11
A_23_P20	NM_01503	Androgen-i	APRIN	-1.31888	0.00098	0.75822	836.7734	633.4877
A_23_P15233		Unknown	A_23_P15	-1.31917	0.0007	0.75805	2589.137	1956.235

A_24_P66	AK092674	Chromosor	C14orf35	-1.31917	0.00003	0.75805	1174.325	890.9507
A_32_P23	AK074151	KIAA1608	KIAA1608	-1.31919	0.00093	0.75804	1352.813	1015.949
A_32_P223173		MY5B_HUI	THC18204	-1.31951	0.00015	0.75786	1072.499	816.6003
A_24_P17	NM_02429	hypothetica	DERL1,DE	-1.31965	0.00034	0.75778	2690.408	2037.663
A_23_P32	NM_00561	Homo sapi	RBBP4,RB	-1.31966	3.02E-10	0.75777	9002.703	6821.419
A_23_P13	NM_00527	Homo sapi	GLUD1,GD	-1.31984	1.68E-08	0.75767	3172.98	2403.699
A_24_P93	AB020706	Adaptor-rel	AP2A2	-1.31992	0.00002	0.75762	1742.505	1324.48
A_23_P16	NM_00559	Homo sapi	NACA,MG	-1.32005	2.30E-07	0.75755	165080.9	125390
A_24_P927537		ALU1_HUN	THC19022	-1.3203	1.35E-06	0.7574	2893.4	2194.702
A_23_P97	AB046826	Hypothetic	FLJ20203	-1.32034	1.65E-06	0.75738	2393.946	1815.752
A_24_P91	NM_00084	glutathione	GSTM3,GS	-1.32035	6.00E-09	0.75738	24681.27	18723.54
A_23_P43	NM_17351	hypothetica	FLJ33814,	-1.32038	0.0004	0.75736	1977.959	1489.678
A_23_P69	BC032300	Similar to F	LOC49385	-1.32062	0.00021	0.75722	5187	3931.128
A_23_P24	NM_01417	thymocyte	THY28,MY	-1.32096	0.00001	0.75703	1414.494	1072.726
A_24_P37	NM_00215	Homo sapi	HSPA4,RY	-1.32101	3.69E-06	0.757	54622.36	41511.91
A_23_P20	NM_00002	presenilin 1	PSEN1,AD	-1.32107	0.00084	0.75696	1261.86	945.7651
A_24_P11	NM_01404	DKFZP564	CHMP2B,C	-1.3211	0.00002	0.75695	1369.416	1036.267
A_23_P99	NM_00268	Homo sapi	PNN,DRS,	-1.32122	0.00058	0.75688	7981.369	6083.826
A_23_P17	NM_01226	Homo sapi	C22orf5,FM	-1.32149	0.0004	0.75672	4264.104	3224.428
A_24_P41	AF318364	Protein phc	PPP1R14E	-1.32165	5.35E-11	0.75663	19265.02	14569.77
A_23_P13	NM_01642	Homo sapi	RNF141,ZI	-1.32166	4.13E-06	0.75662	1960.291	1480.883
A_23_P11	NM_13870	Homo sapi	NDNL2,HC	-1.32179	0.00002	0.75655	4889.552	3700.997
A_23_P14	NM_00101	Homo sapi	RPS7	-1.32221	0.00001	0.75631	402416.9	305814.8
A_23_P57393		Unknown	THC18373	-1.32223	1.11E-06	0.7563	2121.727	1600.498
A_23_P11	AB067472	Valyl-tRNA	VARS2L	-1.32225	5.92E-08	0.75629	2349.796	1777.335
A_23_P90	NM_01857	amyotrophi	ALS2CR2,	-1.32239	1.91E-06	0.75621	2231.848	1684.9
A_24_P29	NM_01813	SHQ1 hom	FLJ10539	-1.32244	0.00041	0.75618	1042.784	783.0726
A_23_P52	NM_01210	Homo sapi	BACE1,AS	-1.32245	2.11E-07	0.75617	3612.386	2738.915
A_24_P37	BC020242	Protein tyr	PLIP	-1.3225	5.56E-06	0.75614	6735.387	5078.404
A_23_P32	NM_14458	Zinc finger,	ZFYVE27	-1.32256	5.62E-06	0.75611	5706.167	4328.582
A_24_P37	AK026976	Hypothetic	FLJ23323	-1.32261	0.00008	0.75608	2477.355	1867.642
A_23_P20	NM_19879	Homo sapi	MAP4K5,K	-1.32281	0.00094	0.75597	888.9922	672.6577
A_23_P51	NM_01784	tRNA seler	SECP43,P	-1.32298	0.00033	0.75587	1257.416	947.3628
A_32_P125820		Unknown	THC18366	-1.32301	0.00003	0.75585	1473.879	1110.171
A_24_P28	NM_01814	mitochondr	MRPS10,M	-1.32304	0.00059	0.75584	1053.592	796.947
A_23_P31	NM_00682	peroxisom	ACOT2,Mt	-1.32328	0.00003	0.7557	7919.603	5994.729
A_23_P74	NM_01431	small mem	TMEM50A	-1.32338	0.00009	0.75564	11256.42	8545.994
A_32_P13	AK128391	Similar to H	LOC40123	-1.32347	0.00022	0.75559	1544.105	1161.103
A_23_P14	NM_00468	Homo sapi	EIF1AY	-1.32356	3.44E-07	0.75554	30753.29	23309.71
A_23_P206598		AY184957	THC19654	-1.32357	1.25E-09	0.75553	3978.313	3005.821
A_23_P13	NM_01754	Homo sapi	POGK,BAS	-1.32367	0.00003	0.75548	1593.323	1209.874
A_23_P47	NM_02047	Homo sapi	YIF1A,54T	-1.32396	7.35E-06	0.75531	11025.45	8307.8
A_32_P93	XM_37272	Homo sapi	NUDT19,R	-1.32442	5.05E-07	0.75505	1668.258	1258.721
A_23_P37	NM_20153	Homo sapi	NDRG2,SY	-1.32459	9.37E-07	0.75495	1906.195	1440.386
A_23_P12	NM_00314	Homo sapi	SSR2,TLA	-1.325	5.24E-08	0.75472	54479.24	41234.96
A_24_P42	BC000845	Homo sapi	BC000845	-1.3251	4.55E-07	0.75466	95298.71	72100.14
A_24_P10	NM_00600	ubiquinol-c	UQCRH	-1.32515	1.59E-08	0.75463	79800.86	60344.21
A_24_P38	NM_00583	Rab9 effec	RABEPK,p	-1.32525	1.02E-09	0.75458	5793.175	4373.431
A_23_P14	NM_15368	Homo sapi	DSCR5,DC	-1.32526	0.00005	0.75457	2497.842	1878.361
A_24_P14	NM_02110	thymosin, b	TMSB4X,F	-1.32561	5.62E-12	0.75437	67764.33	51122.3
A_23_P68	NM_03248	Homo sapi	MCM8,MG	-1.32601	7.00E-08	0.75414	2540.76	1915.939
A_23_P10	NM_02524	Homo sapi	TMEM22,M	-1.32602	0.00023	0.75414	3222.175	2414.814

A_24_P764595		Unknown	A_24_P76	-1.32618	4.11E-11	0.75405	37012.35	27923.84
A_24_P37	NM_00710	solute carri	SLC22A18	-1.32631	2.47E-11	0.75397	12255.66	9240.272
A_23_P69	NM_03271	hypothetica	MGC11324	-1.32638	6.51E-10	0.75393	5913.897	4456.208
A_24_P186354		Unknown	ENST0000	-1.32639	1.71E-07	0.75393	9496.295	7178.753
A_24_P36	BF976161	C-type lect	OCIL	-1.32659	1.49E-06	0.75381	45960.41	34469.38
A_24_P83	NM_19918	Homo sapi	NPM1,B23	-1.32659	0.00011	0.75381	30052.04	22401.63
A_23_P14	NM_00169	Homo sapi	ATP5O,AT	-1.32663	5.87E-08	0.75379	48808.8	36839.45
A_23_P19	NM_01589	Homo sapi	GMNN,Ge	-1.3267	0.00002	0.75375	12515.16	9455.975
A_24_P679796		Unknown	A_24_P67	-1.32676	6.48E-12	0.75372	16878.26	12722.21
A_32_P22014		Unknown	A_32_P22	-1.32679	0.00077	0.7537	1679.282	1276.484
A_23_P79	NM_02067	Homo sapi	RAB22A,M	-1.3268	0.00001	0.75369	2578.723	1942.752
A_23_P56	AB028964	Forkhead b	FOXJ3	-1.32704	5.18E-06	0.75356	3264.014	2459.072
A_24_P65	NM_01824	hypothetica	C20orf44,E	-1.32718	6.97E-07	0.75348	2022.662	1525.257
A_23_P46	NM_01333	Homo sapi	NME7	-1.32718	0.00002	0.75348	1741.333	1306.926
A_24_P839254		ALU8_HUM	THC19993	-1.32733	0.00035	0.75339	3451.679	2599.045
A_23_P21	NM_01561	WIPI49-like	WIPI-2	-1.32736	3.74E-06	0.75338	3036.81	2280.142
A_23_P17	NM_08082	Homo sapi	HARS2,DU	-1.32782	3.26E-06	0.75311	9555.447	7224.888
A_23_P73	NM_01808	Homo sapi	TMEM48,N	-1.32782	0.00001	0.75311	4223.416	3196.947
A_23_P39	NM_00307	SWI/SNF r	SMARCA4	-1.32794	0.00002	0.75304	2907.543	2188.01
A_23_P11	NM_01414	nudix (nucl	NUDT5,YS	-1.32804	5.05E-07	0.75299	8528.887	6443.265
A_32_P75	NM_13880	hypothetica	MGC16471	-1.32805	0.00001	0.75298	5559.712	4186.575
A_32_P14	NM_19929	alpha-NAC	ANAC,MG	-1.32805	1.94E-08	0.75298	113523.4	85696.13
A_24_P30	NM_00483	glutathione	GSTO1,P2	-1.32813	5.94E-06	0.75294	16601.87	12422.8
A_32_P157671		Unknown	A_32_P15	-1.32819	0.00096	0.7529	9412.425	7111.719
A_23_P15	NM_01593	CGI-04 pro	YARS2,CG	-1.32851	0.00006	0.75272	2374.995	1788.324
A_23_P42	NM_18183	Homo sapi	ORC3L,LA	-1.32859	3.21E-06	0.75268	2672.12	2022.133
A_23_P41	NM_03258	Homo sapi	CARD6,CII	-1.32879	0.00004	0.75256	1656.02	1240.22
A_23_P51	NM_00599	Homo sapi	TSNAX,TR	-1.32883	0.00005	0.75254	2244.346	1690.731
A_23_P136392		Unknown	A_23_P13	-1.32886	0.00002	0.75253	1928.684	1447.077
A_32_P33	NM_00534	histidine tri	HINT1,HIN	-1.32889	7.86E-06	0.75251	68525.65	51206.2
A_23_P42	NM_18264	Homo sapi	FALZ,BPTI	-1.32891	0.00062	0.7525	1024.819	765.2107
A_23_P13	NM_00544	Homo sapi	CSPG6,BA	-1.32955	1.19E-06	0.75213	7602.939	5724.766
A_32_P31355		Unknown	A_32_P31	-1.32955	0.00002	0.75214	1412.874	1063.299
A_24_P61	NM_02019	GK001 pro	GK001	-1.32961	0.00038	0.7521	991.1022	745.0856
A_24_P37	NM_00215	Homo sapi	HSPA4,RY	-1.33034	0.00005	0.75169	8291.152	6268.332
A_23_P38	NM_00477	Homo sapi	ZNHIT3,TF	-1.33047	1.54E-06	0.75161	22092.04	16651.41
A_23_P56	NM_03260	Homo sapi	MCEE	-1.33054	7.36E-08	0.75157	2402.919	1806.722
A_23_P53	NM_01425	Homo sapi	TMEM4,MS	-1.33058	1.58E-07	0.75155	64338.45	48489.78
A_24_P92	BC033998	Hypothetic	LOC12451	-1.33063	0.00006	0.75153	1355.929	1021.116
A_24_P18	AL713644	Neurofilam	NEFL	-1.3308	4.10E-07	0.75143	27203.46	20506.06
A_32_P63	NM_00680	splicing fac	SF3A3,PR	-1.33093	8.38E-11	0.75135	5495.283	4128.348
A_23_P33	NM_01828	mitochondr	IMP3,BRM	-1.33105	1.10E-07	0.75129	11873.39	8886.72
A_23_P12	NM_00084	Homo sapi	GSTM3,GS	-1.33139	0.00013	0.7511	74458.89	55860.4
A_32_P81	NM_17490	hypothetica	MGC23909	-1.33153	4.81E-09	0.75102	4898.654	3674.46
A_24_P28	AF272382	Myeloid/lyn	MLL	-1.33156	0.00006	0.751	4917.52	3710.057
A_23_P36	NM_00640	Homo sapi	DCTN2,RB	-1.3316	0.00016	0.75098	5223.604	3930.283
A_23_P39	NM_02110	thymosin, b	TMSB4X,F	-1.3316	2.76E-12	0.75097	72286.86	54289.11
A_24_P31	BC040210	Aldo-keto r	AKR1C1	-1.33178	0.00001	0.75088	4611.333	3453.804
A_23_P49	NM_00309	Homo sapi	SNX1,SNX	-1.33223	3.32E-06	0.75062	1565.567	1172.967
A_23_P25	NM_03214	Homo sapi	FBXO30,F	-1.33237	0.00055	0.75054	2906.637	2179.197
A_23_P14	NM_01595	CGI-30 pro	DPH5,CGI	-1.33261	0.00012	0.75041	4103.063	3068.733
A_24_P37	BC008564	Protein pho	PPP1R9B	-1.33265	3.11E-10	0.75038	3275.969	2458.084

A_23_P25	NM_00708	Homo sapi	LSM6,YDR	-1.33269	2.12E-07	0.75036	9406.271	7074.805
A_23_P25	NM_00294	Homo sapi	RPA3,REP	-1.33273	0.00025	0.75034	13408.56	10074.49
A_23_P93	NM_00279	Homo sapi	PSMB7,Z	-1.33273	1.74E-06	0.75034	41731.83	31148.77
A_23_P95	NM_01803	Homo sapi	WDR6,FLJ	-1.33276	5.57E-10	0.75032	7086.15	5312.576
A_23_P13	NM_00141	eukaryotic	EIF2S3,EIF	-1.33294	3.48E-06	0.75022	2483.28	1862.353
A_23_P13	AK001178	Solute carr	SLC6A15	-1.33313	0.00004	0.75011	1193.188	895.6892
A_24_P30	NM_01242	retinoic aci	IFIT5,RI58	-1.33322	0.00012	0.75007	2760.646	2073.783
A_23_P15	NM_19427	HESB like	HBLD1	-1.33328	0.00003	0.75003	5315.653	3978.236
A_23_P15	NM_15241	hypothetica	DKFZp762	-1.3333	0.00006	0.75002	1228.199	918.3591
A_23_P16	NM_17071	Homo sapi	DAZAP1,M	-1.33365	0.00035	0.74982	6882.024	5198.521
A_23_P84	NM_00652	Homo sapi	TFE3,RCC	-1.33367	0.00005	0.74981	1454.692	1099.663
A_24_P349489		Unknown	A_24_P34	-1.33376	6.09E-06	0.74976	16077.94	12095.54
A_23_P21	NM_00394	Homo sapi	ATP6V0E,I	-1.33379	1.52E-07	0.74974	10699.41	8026.084
A_23_P10	NM_02006	Homo sapi	SLC2A4RC	-1.33388	0.00004	0.74969	20657.6	15558.3
A_23_P15	NM_00450	HUS1 (S. p	HUS1	-1.33394	0.00068	0.74966	971.0085	726.9354
A_23_P12	NM_03321	hypothetica	LOC90355	-1.33395	0.00003	0.74965	3098.674	2321.313
A_23_P20	AF283776	Activating t	ATF2	-1.33405	0.00048	0.7496	1218.002	906.9135
A_23_P13	BC005991	Ubiquitin sp	USP45	-1.33438	0.00046	0.74941	906.1899	678.0731
A_23_P34	NM_01844	Homo sapi	PAG1,CBF	-1.33446	2.29E-06	0.74937	1666.325	1250.5
A_23_P33	NM_01836	meiosis-sp	MNS1,FLJ	-1.33463	0.00005	0.74927	1197.296	895.3565
A_32_P10	NM_01482	KIAA0801	DDX46,MG	-1.33474	4.36E-08	0.74921	2643.504	1982.036
A_23_P12	NM_02481	hypothetica	FLJ22955	-1.33478	3.23E-08	0.74919	2228.601	1668.276
A_23_P25	NM_01832	hypothetica	BXDC2,BR	-1.33518	0.00025	0.74896	7679.196	5795.965
A_23_P63	NM_02280	Homo sapi	CTBP2	-1.33528	2.19E-10	0.7489	23118.28	17293.35
A_24_P20	NM_03310	prostein pr	SLC45A3,F	-1.33535	0.0004	0.74887	1141.275	858.7253
A_24_P170186		Unknown	ENST0000	-1.33537	8.21E-10	0.74886	25481.79	19115.03
A_23_P16	NM_00484	Homo sapi	EIF4E2,4E	-1.33543	0.00002	0.74882	17578.66	13219.25
A_23_P25	NM_02018	DC12 prote	DC12,MGC	-1.33546	0.00036	0.74881	2405.161	1818.264
A_23_P14	NM_01516	homolog of	RRS1,KIA	-1.33566	4.13E-08	0.7487	10204.12	7618.318
A_24_P86	AF318345	Similar to C	LOC39981	-1.33593	3.30E-07	0.74854	1961.865	1468.19
A_24_P43	NM_01847	TPA regula	TPARL,TM	-1.33614	0.00008	0.74842	2464.731	1840.987
A_23_P13	NM_01902	RNA-bindir	FLJ20273	-1.33615	0.00029	0.74842	3070.785	2293.359
A_24_P16	NM_15364	Homo sapi	TPM3,TRK	-1.33621	0.00033	0.74839	1475.396	1098.675
A_23_P20	NM_01507	Homo sapi	KIF1B,KLP	-1.33626	0.00091	0.74836	1136.353	860.5243
A_23_P53	NM_00187	Homo sapi	CPS1	-1.33654	1.15E-06	0.7482	7272.114	5467.663
A_24_P30	AF092095	HIV-1 Tat i	HTATIP2	-1.33668	0.00061	0.74812	855.6596	639.7529
A_24_P36	NM_01557	Homo sapi	FAM61A,C	-1.33681	4.42E-09	0.74805	4456.206	3335.819
A_24_P67268		Unknown	A_24_P67	-1.33691	3.83E-09	0.74799	5538.827	4136.652
A_23_P57	NM_00350	Homo sapi	CDC45L,C	-1.33692	0.0001	0.74799	7656.599	5729.982
A_23_P97	NM_01797	Homo sapi	MRPL20,L	-1.33692	0.00002	0.74799	26763.86	20170.57
A_23_P96	NM_00705	Homo sapi	FAF1,hFAF	-1.3371	0.0009	0.74789	4038.558	3026.513
A_23_P76	NM_00279	proteasom	PSMB4,HN	-1.33711	0.00012	0.74788	61693.58	46108.31
A_23_P12	BC003353	Hypothetic	MGC5309	-1.33716	0.00001	0.74785	2506.625	1874.77
A_24_P22	NM_00440	DNA fragm	DFFA,DFF	-1.3372	0.00001	0.74783	1505.347	1126.489
A_24_P39	NM_00138	diptheria to	DPH2,DPH	-1.33769	0.00002	0.74756	2140.077	1603.85
A_24_P41	NM_20735	similar to F	MGC7210	-1.33777	0.00003	0.74751	2894.46	2158.999
A_23_P11	NM_01228	Homo sapi	KEAP1,INr	-1.33817	0.00003	0.74729	4607.836	3457.572
A_23_P10	NM_14568	Homo sapi	MAP4K4,H	-1.33831	4.17E-08	0.74721	1999.006	1495.406
A_24_P349869		Unknown	A_24_P34	-1.33832	1.22E-09	0.7472	24442.34	18230.13
A_24_P19	NM_03221	Homo sapi	ANKRD17,	-1.33839	0.00033	0.74717	1176.979	884.8688
A_24_P41	NM_00094	protein phc	PPP3CA,C	-1.33859	0.00011	0.74705	3007.738	2249.931
A_23_P20	NM_00645	Homo sapi	PAIP1,MG	-1.33891	8.25E-11	0.74688	5533.401	4132.861

A_23_P11	NM_00640	Homo sapi	HBXIP,XIP	-1.33905	3.35E-10	0.7468	22426.39	16767.75
A_23_P20	BX648862	Zinc finger	ZNF161	-1.33915	0.00003	0.74674	1353.36	1012.596
A_24_P10	NM_01836	LIN-7 prote	LIN7C,VEL	-1.33926	0.00016	0.74668	1121.265	842.242
A_23_P34	NM_03223	hypothetica	FLJ22789	-1.3393	0.00069	0.74666	1160.094	857.9844
A_23_P64	AB029016	Trinucleoti	TNRC6B	-1.33956	0.00076	0.74651	886.4542	668.2662
A_24_P33	NM_02014	putatative 2	LOC56902	-1.33961	0.00003	0.74649	2748.925	2063.75
A_24_P31	NM_00335	ubiquitin-lik	SUMO1,GI	-1.3397	8.91E-06	0.74644	1890.291	1413.346
A_23_P25	AK056630	Zinc finger	ZKSCAN1	-1.3399	6.34E-06	0.74632	1708.926	1280.472
A_23_P18	NM_00721	Homo sapi	PDCD10,C	-1.34003	1.38E-06	0.74625	7196.294	5353.187
A_23_P21	NM_02032	Homo sapi	RARSL,DA	-1.34014	1.78E-06	0.74619	6338.064	4741.115
A_23_P54	NM_03283	Homo sapi	CIRH1A,N	-1.3402	4.02E-06	0.74616	3016.956	2249.53
A_24_P60	NM_00680	splicing fac	SF3A3,PR	-1.34021	0.00003	0.74615	8959.449	6719.384
A_23_P10	NM_00281	Homo sapi	PSMD7,P4	-1.34057	6.07E-06	0.74595	24799.98	18562.69
A_23_P30	NM_00128	chloride int	CLIC1,G6,	-1.34093	2.73E-07	0.74575	15740.58	11790.88
A_23_P20	NM_01455	Homo sapi	SENP1,Su	-1.34104	7.62E-07	0.74569	1627.695	1214.867
A_32_P36	NM_00117	ADP-ribosy	ARL1,ARF	-1.34126	5.63E-09	0.74557	2952.563	2201.721
A_23_P10	NM_01519	Homo sapi	DNAJC9,JI	-1.34138	0.00011	0.7455	10083.75	7518.99
A_32_P106246		Unknown	THC19718	-1.3414	0.00089	0.74549	1585.295	1193.277
A_23_P42	NM_17360	Homo sapi	C15orf21,D	-1.34164	0.0005	0.74536	134263.1	100425.7
A_24_P35	NM_00626	RAN bindir	RANBP2,T	-1.34165	0.00011	0.74535	1169.399	869.5018
A_23_P25	NM_00229	Homo sapi	RPSA,LRP	-1.34167	1.98E-11	0.74534	213352.6	158903.2
A_32_P15	AL117661	TAF3 RNA	TAF3	-1.34173	0.00081	0.74531	1320.163	993.7428
A_23_P93	NM_18271	Homo sapi	SYPL1,SY	-1.34185	5.18E-10	0.74524	11557.29	8613.853
A_23_P25	NM_00207	Homo sapi	GOLGA4,G	-1.34187	7.45E-09	0.74523	2261.169	1684.98
A_24_P31	NM_08063	Homo sapi	UPF3B,UP	-1.34191	0.00069	0.74521	2714.057	2025.932
A_23_P69	NM_00128	Homo sapi	AP3S1,CL	-1.34203	0.00002	0.74514	13554.63	10145.63
A_23_P19	NM_01403	ASF1 anti-s	ASF1A	-1.34214	1.41E-08	0.74508	3183.449	2372.099
A_23_P15	NM_05285	Homo sapi	RFT1	-1.34224	0.00012	0.74502	4287.811	3197.547
A_23_P63	NM_03211	Homo sapi	MRPL43,bl	-1.3423	1.98E-08	0.74499	16140.16	12050.72
A_24_P16	NM_15251	Homo sapi	NM_15251	-1.34235	0.00016	0.74496	4609.679	3427.118
A_32_P22	BC067856	Succinate (	SDHAL2	-1.34236	4.63E-08	0.74495	2691.823	2004.473
A_23_P12	NM_00663	methylene	MTHFD2,N	-1.34241	1.80E-07	0.74493	26267.98	19641.3
A_24_P12	NM_00175	caveolin 1,	CAV1,CAV	-1.3427	0.00064	0.74477	3486.202	2605.75
A_24_P53	NM_00012	electron-tra	ETFA,EMA	-1.34296	2.48E-09	0.74462	13320.48	9891.961
A_23_P94	NM_02521	G kinase a	GKAP1	-1.34314	0.00081	0.74452	998.9404	749.3865
A_23_P13	NM_01773	hypothetica	FLJ20272	-1.34322	4.54E-07	0.74448	7818.165	5839.231
A_23_P83	NM_01603	CGI-90 pro	FAM82B,C	-1.34328	0.00022	0.74444	1965.501	1455.782
A_23_P35	NM_03096	Homo sapi	TRIM56,RN	-1.34341	0.00018	0.74437	1565.479	1177.144
A_23_P39	AK025800	Nucleopori	NUPL1	-1.34344	2.28E-06	0.74436	1927.319	1432.635
A_23_P12	NM_01557	Homo sapi	AUTS2,KIA	-1.34366	9.35E-07	0.74424	1583.857	1176.031
A_23_P63	NM_01242	Homo sapi	IFIT5,RI58	-1.34371	0.0004	0.74421	1285.23	946.116
A_23_P88	BC013943	Hypothetic	MGC24381	-1.34386	2.25E-06	0.74412	3777.037	2813.648
A_23_P15	NM_00279	proteasom	PSMB1,HC	-1.34399	4.61E-11	0.74405	25630.52	19094.99
A_23_P14	BX648086	Coronin, a	CORO2A	-1.34413	3.38E-07	0.74397	1828.325	1361.266
A_23_P31	NM_00463	Homo sapi	MAPKAPK	-1.34442	0.00013	0.74381	2988.116	2229.985
A_23_P20	NM_02467	Homo sapi	SUV39H2,	-1.34448	6.61E-07	0.74378	2652.498	1974.27
A_24_P410246		Unknown	A_24_P41	-1.34454	4.28E-10	0.74375	5634.733	4193.178
A_23_P21	NM_01542	Homo sapi	WDR51A,D	-1.34461	0.00002	0.74371	2305.969	1710.117
A_23_P76	NM_01344	Homo sapi	BAZ1A,AC	-1.34467	2.52E-11	0.74368	2301.135	1715.446
A_23_P97	NM_01222	Homo sapi	MSRB2,CE	-1.34467	1.09E-06	0.74367	5773.012	4306.728
A_23_P14	NM_01805	hypothetica	XKR8,XRG	-1.34493	0.00012	0.74353	3540.789	2672.779
A_23_P13	BC012022	Protein pho	PPP2CB	-1.34503	4.62E-06	0.74348	8891.065	6651.798

A_23_P20	AK000796	Chromosor	C14orf129	-1.34527	0.00063	0.74335	5035.886	3775.029
A_23_P97	NM_01808	hypothetica	IPO9,Imp9	-1.34529	3.46E-07	0.74334	1691.542	1258.936
A_24_P18	NM_00115	solute carri	SLC25A5,T	-1.3453	8.52E-11	0.74333	37488.88	27903.57
A_23_P14	NM_01603	Homo sapi	C14orf166,	-1.34553	7.31E-13	0.7432	45209.33	33593
A_23_P41	NM_13844	Homo sapi	CCDC5,HE	-1.34557	7.34E-07	0.74318	5338.791	3975.149
A_32_P15	NM_01781	hypothetica	CHCHD3,F	-1.34589	0.00002	0.743	2724.553	2025.23
A_23_P20	NM_00227	RAN bindir	RANBP5	-1.34599	1.37E-10	0.74295	3392.239	2520.147
A_23_P13	NM_01554	Homo sapi	PTCD1,KIA	-1.34606	6.22E-06	0.74291	7640.13	5692.539
A_23_P65	NM_01217	Homo sapi	FBXO22,F	-1.34641	1.30E-07	0.74272	5898.454	4372.188
A_23_P25	NM_01775	hypothetica	FLJ20297,	-1.34642	0.00002	0.74271	19832.91	14764.96
A_32_P16	NM_14564	Homo sapi	MRPL35	-1.34684	0.00002	0.74248	5381.319	4011.542
A_32_P41	NM_00551	high-mobili	HMGN2,HI	-1.34691	0.00014	0.74244	141274.5	105738.8
A_24_P20	NM_20338	Homo sapi	ACSL5,AC	-1.34693	0.00043	0.74243	885.2744	656.7354
A_23_P86	BC015167	KIAA0319-	FLJ13840	-1.34699	5.13E-06	0.7424	3217.923	2409.686
A_23_P14	NM_01241	Homo sapi	H2AFV,H2	-1.34706	2.99E-13	0.74236	60255.13	44731.95
A_24_P384239		Unknown	A_24_P38	-1.34707	0.0001	0.74235	8179.758	6070.076
A_23_P14	NM_01639	hypothetica	HSPC111,	-1.34749	1.01E-09	0.74212	61296.87	45566.52
A_24_P269315		ILEU_HUM	THC19557	-1.3475	2.99E-08	0.74211	2214.398	1642.618
A_23_P21	NM_03089	hypothetica	ZNF323,ZN	-1.3476	0.00002	0.74206	1394.598	1030.934
A_23_P25	NM_03127	Homo sapi	AGXT2L1	-1.3476	0.00003	0.74206	2392.946	1768.11
A_23_P49	NM_02523	recombinat	WDR61,RE	-1.34761	0.00021	0.74205	5607.653	4195.829
A_23_P35	NM_00428	Homo sapi	BAG2,BAG	-1.34771	0.00015	0.742	8844.736	6604.01
A_32_P19	NM_17588	Homo sapi	PRPS1L1,	-1.3478	0.0001	0.74195	1318.066	973.7134
A_24_P73	AK093246	Ribosomal	RPL13	-1.348	0.00022	0.74184	1159.734	854.3882
A_24_P39	NM_00375	eukaryotic	EIF3S5,eIF	-1.34801	2.16E-11	0.74183	12601.42	9340.275
A_23_P13	NM_00290	reticulocalb	RCN1,RCN	-1.34803	0.00014	0.74182	3008.28	2235.527
A_23_P13	NM_01640	hypothetica	HSPC138,	-1.34824	0.00004	0.74171	7816.409	5806.821
A_23_P30	NM_19850	HGS_RE4	UNQ1912,	-1.34827	0.00009	0.74169	2187.479	1618.603
A_23_P25	NM_00118	Homo sapi	ATP6AP1,	-1.34829	7.83E-11	0.74168	9368.88	6944.476
A_32_P57	NM_00280	proteasom	PSMC1,S4	-1.34837	8.05E-09	0.74164	41117.71	30570.52
A_23_P66	NM_01814	hypothetica	FLJ10587	-1.34875	0.00016	0.74143	1657.033	1219.975
A_23_P10	NM_14573	Homo sapi	OSBPL6,O	-1.3489	5.60E-06	0.74135	1431.209	1062.758
A_23_P71	NM_01609	Homo sapi	GOLGA7,G	-1.34893	2.57E-07	0.74133	5040.708	3744.372
A_24_P177795		Unknown	ENST0000	-1.34895	7.31E-07	0.74132	314562.3	233864
A_23_P12	NM_03270	tubulin alpl	TUBA6,MG	-1.34895	1.33E-11	0.74132	24182.25	17906.4
A_23_P88	NM_01825	related to C	RC74,RC-7	-1.3491	0.00092	0.74124	2905.18	2152.57
A_24_P39	NM_01811	hypothetica	SDAD1,FL	-1.34935	0.00007	0.7411	1542.791	1149.152
A_23_P39	NM_02023	x 003 prote	TNFSF5IP	-1.34944	2.78E-09	0.74105	2385.212	1767.585
A_23_P46	NM_00126	chromoson	RCC1,CHC	-1.34945	3.17E-06	0.74104	1358.363	1006.792
A_32_P24	AL360137	Hypothetic	LOC12372	-1.3495	1.41E-07	0.74101	5895.917	4379.586
A_24_P38	NM_01842	RNA polym	RRN3,TIFI	-1.34967	0.00052	0.74092	1646.318	1210.431
A_32_P15	NM_00390	eukaryotic	EIF2S2,EIF	-1.34979	0.00021	0.74086	6444.523	4726.446
A_23_P60	NM_00243	mannose p	MPI,PMI,P	-1.34984	0.00001	0.74083	1662.888	1227.232
A_24_P127121		Unknown	A_24_P12	-1.34994	0.00011	0.74078	17578.17	13086.96
A_23_P10	NM_15286	Homo sapi	ARPC2,AR	-1.35004	2.72E-08	0.74072	27536.73	20331
A_32_P173744		Unknown	A_32_P17	-1.35078	2.79E-06	0.74032	21405.05	15916.44
A_23_P36	NM_00069	aldehyde d	ALDH2,AL	-1.35087	2.63E-09	0.74026	4455.957	3300.494
A_23_P96	NM_03222	hypothetica	RP13-360E	-1.35087	4.53E-06	0.74026	4598.023	3405.824
A_24_P22	NM_01862	presenilins	PSARL,PA	-1.35118	0.00026	0.7401	2966.51	2200.427
A_23_P20	NM_14528	similar to h	LOC15119	-1.35158	0.00038	0.73987	2373.743	1753.168
A_23_P92	AK025789	Homo sapi	RFWD2,CC	-1.35159	0.00008	0.73987	3889.999	2871.128
A_23_P15	NM_00052	Homo sapi	HEXB,ENC	-1.35161	7.04E-06	0.73986	6018.333	4474.008

A_23_P20	NM_00281	Homo sapi	PSMD11,S	-1.35172	4.90E-10	0.7398	9691.102	7177.002
A_24_P586660		Unknown	A_24_P58	-1.35178	1.42E-08	0.73976	9416.73	6981.282
A_23_P42	NM_19815	Homo sapi	MITF,WS2	-1.35189	0.00028	0.7397	888.5131	658.8349
A_23_P12	NM_00547	Homo sapi	ABI1,E3B1	-1.35192	0.00021	0.73969	2831.194	2095.374
A_24_P263803		Unknown	ENST0000	-1.35201	3.12E-08	0.73964	2307.246	1704.446
A_23_P20	AK001522	Protein-kin	PRKRIR	-1.3521	1.74E-07	0.73959	6316.8	4673.893
A_23_P28797		DJ470L14.	ARPC3B	-1.35211	7.31E-11	0.73959	14944.44	11043.47
A_24_P298228		Unknown	A_24_P29	-1.35215	9.06E-06	0.73956	22988.05	17057.43
A_23_P11	NM_01792	Homo sapi	C14orf119,	-1.35217	7.21E-09	0.73955	4963.482	3673.61
A_24_P418590		Unknown	A_24_P41	-1.35233	0.00006	0.73946	1397.022	1027.687
A_32_P48	BM726874	Homo sapi	BM726874	-1.35238	0.00029	0.73943	1190.335	885.3796
A_32_P14	NM_18247	Pyruvate ki	PKM2	-1.35241	4.82E-08	0.73942	6656.979	4916.538
A_23_P25	NM_01614	Homo sapi	COMMD1C	-1.35242	4.45E-10	0.73941	3465.138	2561.324
A_23_P77	NM_03249	hypothetica	HH114,MG	-1.35244	0.00015	0.7394	988.5408	733.9216
A_24_P22	NM_00619	proliferatio	PA2G4,EB	-1.35252	9.79E-06	0.73936	13400.77	9841.992
A_24_P101493		Unknown	A_24_P10	-1.35263	3.86E-07	0.7393	2586.152	1912.835
A_23_P13	NM_00427	Homo sapi	PMPCB,MI	-1.35274	1.65E-08	0.73924	20976.43	15548.7
A_24_P22	NM_00452	kinesin-like	KIF11,EG5	-1.3528	1.56E-07	0.73921	2484.174	1831.748
A_23_P40	AK027615	Widely-inte	WIZ	-1.35289	0.00072	0.73916	3886.555	2876.821
A_23_P120644		Unknown	A_23_P12	-1.35302	8.99E-07	0.73908	6872.565	5096.667
A_23_P11	NM_00611	HS1 bindin	HAX1,HS1	-1.35305	1.68E-07	0.73907	14079.11	10386.28
A_24_P46	NM_01786	Homo sapi	PIGX,FLJ2	-1.35308	3.44E-06	0.73905	2748.747	2029.256
A_24_P42	NM_00486	secretory c	SCAMP1,S	-1.35308	0.00058	0.73905	1249.469	933.0978
A_23_P15	NM_03237	hypothetica	MGC4251	-1.35309	0.00039	0.73905	1285.178	957.4802
A_23_P73	BC000968	T-complex-	TCTE1L	-1.35327	0.00001	0.73895	6655.675	4917.128
A_23_P35	M90656	Glutamate-	GCLC	-1.35341	3.73E-09	0.73888	4432.887	3277.211
A_32_P95	NM_00221	Homo sapi	ITGB1,CD2	-1.35346	0.00017	0.73885	20972.47	15547.65
A_23_P14	NM_01828	Homo sapi	ARHGAP1	-1.35356	0.00005	0.73879	1388.774	1021.444
A_24_P17	NM_02019	GK001 pro	GK001	-1.35364	3.78E-07	0.73875	6131.436	4532.571
A_24_P213144		Unknown	A_24_P21	-1.35369	0.00053	0.73872	4214.78	3134.423
A_23_P78	NM_03129	hypothetica	MGC2963	-1.35415	3.70E-06	0.73847	15083.17	11108.38
A_24_P38	NM_17223	Homo sapi	SYVN1,HR	-1.35436	0.00067	0.73836	2144.272	1565.496
A_24_P18	NM_00115	solute carri	SLC25A5,T	-1.35454	1.04E-12	0.73826	44238.91	32676
A_23_P57	NM_00635	Homo sapi	SLC25A17	-1.35457	1.40E-10	0.73824	3232.534	2386.664
A_23_P72	NM_00676	Homo sapi	YWHAE,M	-1.35463	3.00E-10	0.73821	7739.445	5710.009
A_23_P82	AB067484	Hypothetic	FLJ20485	-1.35464	4.72E-08	0.7382	8806.922	6499.341
A_23_P88	NM_00565	Homo sapi	TP53BP1,p	-1.35466	6.52E-06	0.73819	1797.503	1330.9
A_23_P92	NM_00106	Homo sapi	TKT,TKT1	-1.35489	0.00024	0.73807	119780.7	87423.72
A_23_P98	NM_02121	transposon	LOC58486	-1.35493	5.49E-09	0.73805	4827.156	3566.386
A_23_P10	NM_00070	ATPase, N	ATP1A1,M	-1.35495	7.74E-06	0.73804	8127.112	5974.241
A_23_P25	NM_17507	Homo sapi	APTX,AOA	-1.35497	6.25E-06	0.73802	1577.263	1161.93
A_23_P37	NM_08039	Homo sapi	PTP4A2,H	-1.35501	2.44E-12	0.738	51113.39	37695.45
A_24_P587803		Unknown	A_24_P58	-1.35552	5.77E-07	0.73772	71092.19	52567.83
A_24_P651129		Unknown	A_24_P65	-1.35582	0.00091	0.73756	4048.209	3005.774
A_24_P144556		Unknown	A_24_P14	-1.35583	0.0001	0.73756	1724.851	1279.462
A_24_P38	NM_00500	non-histon	NHP2L1,N	-1.35642	5.17E-06	0.73724	3980.461	2927.779
A_24_P17	NM_01799	hypothetica	FLJ10081,	-1.35654	0.00083	0.73717	1347.313	979.7805
A_23_P63	NM_03211	Homo sapi	MRPL43,b	-1.35657	2.16E-10	0.73715	7173.446	5294.26
A_23_P20	NM_00208	Homo sapi	GPC1	-1.35681	1.19E-12	0.73702	24122.58	17786.34
A_23_P11	NM_00412	hypothetica	LOC55289	-1.35684	0.0001	0.73701	3362.715	2482.568
A_23_P26	NM_01844	selenoprot	SELS,VIMI	-1.35692	3.13E-08	0.73696	12788.33	9452.545
A_24_P13	NM_02182	hypothetica	MDS025,M	-1.35707	0.00052	0.73688	1050.662	769.027

A_23_P14	NM_00519	Homo sapi	CCNC	-1.35721	0.00003	0.7368	1463.086	1082.524
A_23_P20	NM_01418	HSPC163	CNIH4,HSI	-1.35773	1.89E-12	0.73652	30770.32	22679.58
A_23_P11	NM_00723	calcium bin	CHP,SLC9	-1.35794	1.60E-07	0.73641	46461.64	34281.68
A_32_P11	NM_00203	Ferritin, he	FTH1	-1.35807	0.00004	0.73634	351715.2	260123.6
A_23_P74	NM_00726	Homo sapi	PARK7,DJ	-1.35836	5.09E-12	0.73618	48577.59	35731.13
A_24_P25	NM_02410	hypothetica	MGC2747	-1.35849	1.73E-07	0.73611	2419.559	1779.554
A_23_P20	NM_00302	Homo sapi	SHC1,SHC	-1.35887	8.51E-10	0.73591	9067.795	6676.015
A_23_P20	NM_01221	Homo sapi	MGEA5,ME	-1.35898	1.17E-09	0.73584	8690.799	6386.287
A_24_P418786		Unknown	ENST0000	-1.35904	0.00002	0.73581	10122.54	7484.026
A_23_P50	NM_00221	Homo sapi	ITGAV,CD5	-1.35927	1.40E-07	0.73569	2124.875	1561.941
A_23_P10	NM_00692	stromal cel	SDF2	-1.3594	0.00038	0.73562	1080.245	798.5877
A_23_P36	NM_00662	Homo sapi	AHCYL1,D	-1.35941	4.16E-13	0.73561	18235.43	13416.62
A_23_P10	NM_05296	Homo sapi	C1orf19,se	-1.35949	2.32E-06	0.73557	2730.929	2006.968
A_23_P14	NM_00396	ubiquitin-co	UBE2M,UE	-1.3595	1.10E-06	0.73557	2195.454	1620.894
A_23_P25	NM_00357	Homo sapi	STK24,MS	-1.35952	3.66E-09	0.73555	3208.709	2358.154
A_23_P37	NM_01224	SKI-interac	SNW1,Bx4	-1.35953	3.03E-07	0.73555	5780.876	4261.228
A_23_P25	NM_00372	Homo sapi	RTCD1,RP	-1.35956	3.37E-08	0.73553	6100.294	4501.587
A_24_P91	BC058041	Ribosomal	RPL23A	-1.3596	0.00009	0.73551	4467.628	3296.602
A_23_P25	NM_01603	Homo sapi	TTC15,CG	-1.35961	1.52E-06	0.7355	1597.583	1177.06
A_32_P51524		A40201 art	THC19422	-1.35985	0.00008	0.73538	1024.016	753.0726
A_32_P21	M83234	Human nu	NSEP	-1.35987	0.00003	0.73536	22339.5	16473.93
A_24_P24	NM_13904	Homo sapi	SMARCA3	-1.35991	0.00023	0.73534	1734.163	1281.562
A_32_P104263		AJ318213	THC19216	-1.36	1.09E-07	0.73529	2882.143	2122.497
A_23_P99	NM_01591	homolog of	POP5,RPP	-1.3602	6.52E-08	0.73518	3707.126	2721.172
A_23_P52	NM_00319	Homo sapi	TBCE,HRD	-1.36021	6.46E-12	0.73518	12247.24	9008.48
A_23_P38	NM_01589	B cell RAG	GALNAC4	-1.36021	0.0007	0.73518	1154.945	837.2694
A_32_P42236		Unknown	THC18311	-1.3603	6.27E-06	0.73513	5506.078	4049.558
A_23_P15	NM_00334	Homo sapi	UBE2I,P18	-1.36043	3.66E-09	0.73506	21214.19	15613.76
A_32_P25	BX648347	Vacuolar p	VPS41	-1.36043	0.00067	0.73506	798.6265	583.8713
A_24_P62	NM_00044	presenilin 2	PSEN2,AD	-1.36047	0.00034	0.73504	1557.146	1136.204
A_24_P20	NM_01405	likely ortho	HIGD1A,H	-1.36049	2.52E-11	0.73503	11602.47	8515.077
A_24_P40	NM_00496	high-mobili	HMGN1,HI	-1.36071	0.0002	0.73491	24871.1	18424.4
A_23_P29	NM_00371	vesicle doc	VDP,TAP,F	-1.36072	7.85E-13	0.7349	5027.712	3694.877
A_24_P46	BC046178	Chromosor	C6orf173	-1.36079	7.17E-08	0.73487	4714.726	3467.291
A_24_P77	AK055214	CDNA FLJ	AK055214	-1.36096	0.00005	0.73478	974.8949	715.8621
A_23_P29	NM_03067	Homo sapi	SEC13L1,S	-1.36136	5.67E-06	0.73456	52448.3	38695.28
A_23_P11	NM_00395	Homo sapi	MPZL1,PZ	-1.36141	1.30E-07	0.73454	3803.442	2794.198
A_23_P35	NM_03081	hypothetica	MGC11335	-1.36149	0.00003	0.73449	2496.785	1828.81
A_24_P18	NM_00285	paxillin	PXN	-1.36153	0.00032	0.73447	14279.63	10563.1
A_24_P34	NM_01765	hypothetica	ZNF562,FL	-1.36155	0.00033	0.73446	870.8983	637.7253
A_23_P11	NM_00514	DnaJ (Hsp	DNAJA3,T	-1.36174	1.18E-07	0.73436	12646.65	9320.731
A_24_P36	AK023023	WD repeat	WDR26	-1.36185	0.00036	0.7343	1498.777	1091.819
A_23_P55	NM_02515	hypothetica	TARSL1,FI	-1.36186	6.56E-07	0.73429	2560.38	1879.683
A_23_P60	NM_00254	Homo sapi	ODF2,ODF	-1.36201	2.06E-06	0.73421	1531.636	1122.104
A_23_P16	NM_00335	UDP-gluco	UGDH,GDI	-1.36203	0.00004	0.7342	3274.844	2402.625
A_23_P146736		SYQ_HUM	THC19213	-1.36222	1.80E-11	0.7341	8463.43	6208.106
A_23_P10	NM_13046	Homo sapi	D4ST1,HD	-1.36225	2.81E-07	0.73408	3989.758	2924.103
A_23_P39	NM_00023	Homo sapi	LIG1	-1.36228	4.42E-09	0.73407	3571.779	2622.429
A_24_P230009		Unknown	A_24_P23	-1.36235	5.21E-08	0.73403	1810.162	1329.214
A_23_P39	NM_00704	Homo sapi	BTN3A2,B	-1.36243	0.00066	0.73398	960.4979	712.9331
A_23_P21	NM_00189	Homo sapi	CSNK1A1,	-1.36244	0.00034	0.73398	6163.541	4537.576
A_23_P20	NM_13043	Homo sapi	DPP8,MST	-1.36247	0.00002	0.73396	2338.96	1718.177

A_23_P13	NM_01744	polymeras	POLE3,p1	-1.36249	6.50E-11	0.73395	10947.07	8037.907
A_24_P10	NM_00142	enolase 1,	ENO1,NNE	-1.36267	1.25E-07	0.73385	44755.05	32695.66
A_23_P50	NM_13364	Homo sapi	SURF5,ME	-1.36284	0.00001	0.73376	7580.623	5598.511
A_23_P33	NM_00575	Ras-GTPa	G3BP,HDH	-1.36321	1.45E-06	0.73356	7106.608	5228.429
A_23_P10	BC018664	Chromosor	C2orf29	-1.36324	0.0001	0.73355	1279.45	932.2987
A_24_P25	NM_00189	casein kina	CSNK1A1,	-1.36331	7.78E-09	0.73351	3228.919	2370.148
A_23_P77	NM_02461	fructosamir	FN3KRP,F	-1.36332	3.02E-08	0.7335	4160.19	3054.977
A_23_P50	NM_02198	ubiquitin-co	UBE2V1,C	-1.36341	0.00001	0.73346	5574.798	4099.936
A_23_P46	D86960	Lysophosp	LPGAT1	-1.36348	1.05E-08	0.73342	2421.109	1775.624
A_24_P34	NM_00280	proteasom	PSMC1,S4	-1.36355	2.49E-09	0.73338	44053.14	32393.49
A_23_P25	NM_00536	Homo sapi	MAGEA12,	-1.36358	0.00016	0.73336	2447.008	1790.841
A_24_P681218		EF11_HUM	THC19753	-1.36387	5.50E-06	0.73321	443987.3	327175.7
A_23_P201998		AF320996	THC19433	-1.36389	0.00011	0.7332	5936.809	4371.702
A_23_P32	NM_02459	Homo sapi	RHBDL6,R	-1.36452	0.00001	0.73286	1964.074	1432.978
A_24_P96	NM_00647	ATP synth	ATP5L,ATI	-1.36454	0.00007	0.73285	1426.26	1050.655
A_23_P12	NM_14523	Homo sapi	C14orf143,	-1.36465	9.49E-07	0.73279	2423.296	1777.805
A_23_P20	NM_03223	Ubiquitin s	USP48	-1.36481	0.00012	0.7327	3301.448	2416.727
A_23_P11	NM_01797	Zwilch	ZWILCH,K	-1.36485	3.67E-08	0.73268	2623.331	1921.962
A_23_P30	NM_00280	Homo sapi	PSMC2,S7	-1.36497	5.83E-10	0.73262	13315.22	9729.15
A_24_P10	NM_01906	hypothetica	GNL3L,FL	-1.36497	0.00001	0.73262	9316.982	6862.561
A_23_P31	NM_01658	Homo sapi	CBX3,HEC	-1.36529	7.50E-07	0.73244	72796.98	53610.27
A_23_P10	NM_00487	Homo sapi	TMEM59,C	-1.36539	5.05E-08	0.73239	9289.114	6817.265
A_23_P53	NM_00616	Homo sapi	NFYB,HAF	-1.36549	4.20E-10	0.73234	2873.307	2103.967
A_24_P58607		Unknown	A_24_P58	-1.36562	1.16E-06	0.73227	2220.348	1620.858
A_32_P11	NM_01475	KIAA0187	BMS1L,KIA	-1.36572	0.00031	0.73222	821.426	601.7399
A_24_P84719		Unknown	ENST0000	-1.36595	0.00096	0.73209	906.6122	656.3966
A_23_P13	NM_03076	Homo sapi	NSBP1	-1.36596	1.70E-07	0.73209	5124.782	3776.259
A_23_P70	NM_01248	Homo sapi	HMMR,IHA	-1.36619	0.00023	0.73196	5630.64	4136.202
A_23_P200815		U91538 ve	THC19312	-1.3662	1.39E-06	0.73196	1468.824	1080.977
A_23_P25	NM_13907	hypothetica	FLJ13614,	-1.36639	0.00099	0.73185	758.8807	551.276
A_24_P24982		Unknown	A_24_P24	-1.36658	1.01E-08	0.73176	14535.95	10670.46
A_32_P62	AK091784	Homo sapi	AK091784	-1.36661	0.0009	0.73174	773.6537	565.6235
A_24_P41	NM_00578	Not56 (D. r	ALG3,CDG	-1.36661	3.53E-11	0.73174	10036.23	7345.978
A_23_P21	NM_00371	Homo sapi	CDC2L5,C	-1.36679	0.00004	0.73164	1609.794	1171.654
A_23_P29	NM_00623	Homo sapi	POLR2H,R	-1.36685	4.08E-11	0.73161	16663.93	12203.04
A_23_P40	NM_01909	Homo sapi	C20orf155,	-1.36714	2.99E-08	0.73145	8122.108	5936.51
A_32_P86	BC032913	Hypothetic	LOC38902	-1.36717	4.25E-08	0.73144	1691.326	1236.861
A_24_P94	AK074645	Apolipopro	APOL6	-1.36725	0.00027	0.7314	1073.206	780.183
A_32_P15	NM_02406	hypothetica	PDCL3,VIA	-1.36729	2.72E-11	0.73138	3801.008	2779.624
A_23_P21	NM_01233	nuclear pre	NARF,FLJ	-1.36742	1.50E-06	0.7313	1452.183	1060.45
A_24_P56	NM_14477	Homo sapi	MBNL2,ME	-1.3675	7.91E-06	0.73126	1577.005	1148.656
A_23_P26	AB014572	KIAA0672	KIAA0672	-1.3676	0.00005	0.73121	1323.674	961.8471
A_23_P29	NM_00730	Synuclein,	SNCA	-1.36769	1.63E-08	0.73116	4759.322	3480.846
A_23_P34	NM_01492	KIAA0971	KIAA0971	-1.36776	3.15E-11	0.73112	3532.434	2582.437
A_23_P25	NM_01627	selenoprot	SELT	-1.36779	7.81E-09	0.73111	7130.354	5222.574
A_32_P97	NM_00280	proteasom	PSMC1,S4	-1.36801	1.96E-11	0.73099	11708.22	8548.189
A_23_P20	NM_00686	Homo sapi	RAB35,RA	-1.36805	4.82E-06	0.73097	6087.451	4463.053
A_23_P13	NM_02196	Homo sapi	ZNF148,BE	-1.36814	3.60E-07	0.73092	3489.4	2553.658
A_24_P94	AL137342	UDP glyco	UGT8	-1.36828	0.00093	0.73084	743.0189	541.7366
A_32_P19460		AP003240	THC19590	-1.36852	5.58E-07	0.73072	2471.738	1801.409
A_24_P16	AK096509	ADP-ribosy	ARFIP1	-1.36876	7.85E-08	0.73059	3176.381	2315.589
A_23_P20	NM_03057	outer mitoc	CYB5-M,O	-1.36901	0.00001	0.73045	1093.811	799.3909

A_23_P38	NM_00179	Homo sapi	CDH2,CDH	-1.3691	0.00006	0.7304	1854.115	1363.403
A_24_P84	XM_29392	similar to p	LOC34564	-1.36912	1.90E-07	0.7304	6474.669	4737.394
A_23_P14	NM_00592	Homo sapi	MFAP1	-1.36923	1.07E-07	0.73034	5639.037	4120.452
A_24_P71	NM_01241	glutaminyl-	QPCT,QC,	-1.36924	0.00024	0.73033	6373.839	4678.896
A_32_P20	NM_14571	Homo sapi	TIGD2	-1.36928	0.00092	0.73031	954.9381	684.9097
A_24_P28	NM_00417	solute carri	SLC1A3,E	-1.36933	0.00019	0.73029	1370.03	1009.352
A_23_P11	NM_00692	stromal cel	SDF2	-1.36939	3.20E-11	0.73025	3928.863	2869.017
A_24_P81	NM_02103	RAP2A, m	RAP2A,KR	-1.36958	0.00043	0.73015	1297.237	936.6182
A_23_P80	NM_02024	Homo sapi	KIF15,HKL	-1.37	9.58E-07	0.72993	2402.24	1751.016
A_24_P64	NM_02489	hypothetica	FLJ11783	-1.37008	0.00031	0.72988	1162.029	838.6778
A_24_P58647		Unknown	A_24_P58	-1.37043	1.06E-10	0.7297	9370.338	6844.104
A_24_P12	NM_02020	Nit protein	NIT2	-1.37046	0.0005	0.72968	9896.785	7282.678
A_23_P27	NM_03234	hypothetica	MGC13096	-1.37062	0.00009	0.7296	4392.321	3211.46
A_24_P864777		SMP1_HUI	THC20041	-1.37065	5.74E-09	0.72958	3170.282	2312.35
A_23_P16	NM_00553	interferon,	IFI16,PYHI	-1.37084	6.03E-09	0.72948	3699.015	2699.136
A_32_P29	NM_00608	IK cytokine	IK,RED,CS	-1.37094	1.53E-12	0.72943	7298.082	5322.432
A_32_P11	XM_37526	similar to H	LOC40045	-1.37116	3.15E-13	0.72931	35888.18	26158.06
A_23_P25	NM_17217	Homo sapi	MRPL42,M	-1.37117	8.47E-09	0.7293	5776.642	4208.139
A_24_P30	NM_05292	Homo sapi	SMYD4,ZM	-1.37126	0.00089	0.72926	909.7128	670.4044
A_23_P38	NM_02182	hypothetica	FLJ13149,	-1.37155	3.64E-11	0.7291	8572.233	6258.963
A_24_P40	BC034962	Hypothetic	15E1.2	-1.37164	0.00003	0.72906	1154.8	845.0477
A_23_P37	AK055367	Chromosor	C14orf126	-1.37166	0.00009	0.72904	1537.601	1129.07
A_24_P21	NM_00320	nuclear fac	NFE2L1,NI	-1.37219	1.45E-08	0.72876	4083.428	2980.073
A_24_P58	NM_00203	Ferritin, he	FTH1	-1.3722	3.58E-06	0.72876	473963.6	347548
A_23_P16	NM_02115	hypothetica	TXNDC13,	-1.37241	0.00007	0.72865	964.955	705.6584
A_24_P409440		Unknown	A_24_P40	-1.37251	6.48E-10	0.72859	7140.285	5193.404
A_32_P16	NM_01910	beta-1,4 m	ALG1,HMT	-1.37263	6.47E-09	0.72853	3537.159	2575.598
A_24_P324644		Unknown	A_24_P32	-1.37278	0.00002	0.72845	1339.182	978.6121
A_24_P32	NM_02523	hypthetical	WDR23,GL	-1.37299	0.0006	0.72834	2846.888	2081.783
A_23_P35	NM_00410	Homo sapi	FDX1,ADX	-1.37318	4.24E-07	0.72824	6935.405	5063.318
A_23_P35	AB037838	Inhibitor of	IBTK	-1.3734	0.00002	0.72812	5000.557	3657.031
A_23_P20	AB007925	SLIT-ROB	SRGAP2	-1.3735	5.17E-08	0.72807	2572.49	1871.969
A_23_P99	NM_01785	timeless-in	FLJ20516	-1.37353	0.00014	0.72805	15805.79	11563.48
A_23_P34	NM_00311	Homo sapi	SP100,FLJ	-1.37354	2.54E-06	0.72805	93266.36	68489.13
A_24_P40	XM_37336	similar to c	LOC39253	-1.37411	6.34E-10	0.72774	3384.156	2463.039
A_24_P33	NM_01476	KIAA0174	KIAA0174,	-1.37412	7.92E-09	0.72774	2530.875	1841.545
A_23_P76	NM_15272	Homo sapi	EFHA1,DC	-1.37421	3.16E-09	0.72769	5593.042	4074.766
A_23_P97	NM_05283	secretory c	SCAMP3,C	-1.37441	8.97E-07	0.72758	3785.017	2755.89
A_23_P17	NM_00429	Homo sapi	ABCB7,AB	-1.37446	8.15E-08	0.72756	3029.273	2203.129
A_32_P95	NM_01819	hypothetica	FLJ10719,	-1.37459	5.78E-07	0.72749	4170.737	3043.659
A_24_P34	L34035	Malic enzy	ME1	-1.3746	0.00001	0.72748	3229.695	2349.423
A_23_P16	NM_01243	Homo sapi	SNAPAP,S	-1.3746	1.54E-09	0.72749	4602.111	3346.025
A_24_P32	XM_06790	similar to T	LOC13255	-1.37465	0.00006	0.72746	39011.16	28552.31
A_24_P13	NM_02125	RAB18, m	RAB18,RA	-1.37467	2.73E-09	0.72745	2079.904	1512.234
A_23_P16	NM_00368	KH-type sp	KHSRP	-1.37469	1.47E-08	0.72744	27525.72	20062.36
A_23_P39	NM_01508	Homo sapi	SPG20,SP	-1.37472	0.00016	0.72742	927.9276	677.7453
A_32_P99	NM_00450	heat shock	HSF2,MGC	-1.37472	0.0001	0.72742	1416.381	1041.762
A_23_P32	NM_02119	Homo sapi	SQRDL,CC	-1.37482	4.10E-07	0.72737	9806.277	7177.332
A_23_P32	NM_01788	Homo sapi	C9orf95,NF	-1.37495	0.00001	0.7273	1689.664	1224.583
A_23_P72	NM_00379	Homo sapi	SNX4	-1.3752	2.48E-11	0.72717	2806.403	2040.884
A_32_P11	AK092667	FLJ35348	FLJ35348	-1.37544	3.14E-09	0.72704	4754.319	3458.254
A_23_P98042		Unknown	A_23_P98	-1.37564	1.97E-06	0.72693	13957.7	10193.35

A_23_P69	NM_15228	Homo sapi	ZFP62,ZET	-1.37572	1.04E-06	0.72689	4229.743	3073.685
A_23_P21	NM_01640	Homo sapi	C20orf43,C	-1.37576	2.13E-07	0.72687	4580.527	3332.287
A_24_P24	NM_00561	RE1-silenc	REST,XBR	-1.37581	0.00005	0.72684	1594.094	1149.996
A_23_P13	NM_00600	tubulin, alp	TUBA3,FL	-1.376	3.68E-10	0.72675	27692.91	20074.48
A_23_P10	NM_00587	splicing fac	SF3A1,PR	-1.37609	1.31E-09	0.7267	6477.796	4713.63
A_24_P31	NM_01552	small fragm	REXO2,RF	-1.3763	6.18E-07	0.72658	3174.173	2307.363
A_32_P16	AK094623	Programme	PDCD6	-1.37633	0.00023	0.72657	983.1975	720.731
A_24_P27	NM_00427	TRF-proxin	TRFP,PRC	-1.37642	0.00012	0.72652	884.2643	641.171
A_23_P11	NM_00112	Homo sapi	ADK,AK	-1.37648	1.09E-06	0.72649	4404.834	3202.685
A_23_P21	NM_00539	Homo sapi	PODXL,PC	-1.37648	2.10E-07	0.72649	8305.028	6057.194
A_23_P42	NM_00683	Homo sapi	RAB32	-1.37666	1.88E-07	0.72639	4265.111	3106.186
A_23_P21	NM_00220	integrin, alp	ITGAE,CD	-1.3768	1.79E-06	0.72632	5869.108	4273.101
A_32_P18	AB032963	ATPase, C	ATP8B2	-1.37691	4.51E-08	0.72626	3037.178	2202.913
A_24_P27	AB032986	Homo sapi	KIAA1160	-1.37697	0.00001	0.72623	1408.808	1018.516
A_24_P28	NM_01329	TBP-1 inte	TBPIP,GT1	-1.37703	0.00003	0.7262	1729.421	1250.96
A_23_P11	NM_01646	Homo sapi	TMEM14C	-1.37714	7.83E-14	0.72614	25522.07	18541.04
A_23_P36	NM_00531	Homo sapi	GTF2H1,B	-1.37722	6.16E-07	0.7261	2691.409	1952.176
A_23_P10	NM_01481	Homo sapi	PSMD6,S1	-1.37727	1.11E-12	0.72607	10329.55	7506.479
A_24_P34	NM_01500	SMART/HD	SPEN,MIN	-1.37732	9.19E-08	0.72605	1686.196	1228.474
A_24_P80	NM_02406	hypothetica	PDCL3,VIA	-1.37744	0.00064	0.72599	8115.449	5940.564
A_24_P32	NM_01343	ubiquilin 1	UBQLN1,D	-1.3775	3.09E-09	0.72595	4504.601	3266.904
A_23_P69	NM_01844	30 kDa pro	LOC55831	-1.3777	0.00013	0.72585	2230.711	1623.762
A_23_P25	NM_00166	ADP-ribosy	ARF5	-1.37771	0.00006	0.72584	1598.927	1167.155
A_23_P41	NM_01781	hypothetica	LYAR,FLJ2	-1.37772	7.03E-07	0.72584	3601.037	2616.623
A_23_P15	NM_00727	Prohibitin 2	REA	-1.37794	4.31E-09	0.72572	60922.91	44315.98
A_23_P13	NM_00404	Homo sapi	ATP6V0B,A	-1.37799	1.90E-07	0.72569	11401.3	8283.273
A_23_P92	NM_03233	hypothetica	MGC2408	-1.37804	9.02E-12	0.72567	6116.266	4440.071
A_23_P37	NM_00121	Homo sapi	CA12,CAX	-1.37815	0.0002	0.72561	15293.62	11196.43
A_23_P40	NM_01239	Homo sapi	PFAS,PUR	-1.3784	0.00012	0.72548	928.1831	671.9571
A_23_P40	NM_00309	small nucle	SNRPB2,M	-1.37846	5.83E-06	0.72544	12997.39	9487.227
A_32_P31	NM_01243	splicing fac	SF3B1,PR	-1.37853	1.67E-09	0.72541	9683.639	7041.483
A_23_P88	NM_00361	Homo sapi	SIP1,GEM	-1.37863	6.63E-09	0.72536	3588.912	2606.052
A_23_P20	NM_00638	Homo sapi	APPBP2,P	-1.37869	0.00004	0.72532	1503.316	1091.164
A_23_P75	NM_00061	Homo sapi	CD59,MIN	-1.37873	2.82E-13	0.72531	5745.877	4166.802
A_24_P56	NM_01598	cytokine re	CRLF3,CR	-1.37879	3.26E-07	0.72527	2284.148	1653.794
A_23_P32	NM_14516	Homo sapi	SFT2D1,pF	-1.37888	1.95E-09	0.72522	11883.89	8596.73
A_24_P255473		Unknown	A_24_P25	-1.37898	1.89E-13	0.72517	10150.76	7358.805
A_23_P11	NM_03120	hypothetica	FAHD1,YIS	-1.37898	4.08E-06	0.72517	1706.45	1240.982
A_23_P52	NM_01402	Homo sapi	DCPS,HSL	-1.37902	7.90E-08	0.72515	4830.243	3493.882
A_24_P409661		Unknown	A_24_P40	-1.37905	5.32E-10	0.72514	8776.742	6362.036
A_23_P58	NM_00719	Homo sapi	ANXA10,A	-1.37912	7.06E-14	0.7251	55683.41	40397.29
A_24_P35	NM_01827	hypothetica	TMEM19,F	-1.37915	0.00075	0.72508	1647.976	1210.028
A_23_P29	NM_14463	Homo sapi	TMEM42,M	-1.3793	1.04E-06	0.725	8174.478	5932.095
A_24_P600622		Unknown	A_24_P60	-1.3795	0.00024	0.7249	1511.244	1106.65
A_32_P15	NM_01527	Homo sapi	KIAA1033	-1.37956	2.97E-10	0.72487	3938.781	2853.389
A_24_P37	NM_02273	nuclear ubi	NUCKS1,J	-1.3797	0.00005	0.7248	15663	11265.88
A_23_P15	NM_01810	Homo sapi	ZDHHC4,Z	-1.38009	1.74E-08	0.72459	10953.5	7954.352
A_24_P14	NM_01534	hypothetica	DULLARD,	-1.38011	0.00012	0.72458	1235.964	903.621
A_23_P19	D79992	Mediator of	MDC1	-1.38018	4.54E-07	0.72454	1341.209	970.1475
A_23_P52	NM_00564	Homo sapi	TARBP1,T	-1.38019	0.00025	0.72454	1854.566	1335.948
A_23_P59	NM_02018	Homo sapi	ACN9,DC1	-1.38035	6.79E-06	0.72445	6038.738	4380.679
A_23_P25	NM_01343	Homo sapi	UBQLN1,D	-1.38035	4.91E-07	0.72445	2060.366	1494.276

A_23_P34	NM_00334	Homo sapi	UBE2N,UB	-1.38057	0.00062	0.72434	6753.697	4932.136
A_23_P45	NM_01785	hypothetica	C1orf109,F	-1.38064	2.11E-07	0.7243	2167.667	1585.069
A_23_P42	AK097877	Homo sapi	AK097877	-1.38093	4.30E-07	0.72415	2701.902	1951.403
A_23_P14	NM_03229	Homo sapi	ANKRD32,	-1.38102	9.83E-08	0.7241	2125.58	1536.713
A_23_P20	NM_01591	endoplasm	TXNDC12,	-1.38113	0.00029	0.72404	1167.308	836.151
A_23_P16	NM_02465	Homo sapi	IPO4,Imp4	-1.38154	3.75E-07	0.72383	4469.252	3235.762
A_23_P39	NM_05284	hypothetica	MGC2025	-1.38155	3.00E-06	0.72383	5319.94	3866.808
A_23_P21	NM_01783	Homo sapi	OCIAD1,O	-1.38162	0.00002	0.72379	9968.386	7262.579
A_24_P401670		Unknown	ENST0000	-1.38164	2.64E-10	0.72378	31366.56	22734.56
A_32_P21	NM_01833	hypothetica	RFK,RIFK,	-1.38168	9.86E-06	0.72375	8037.38	5828.59
A_23_P58	NM_01898	Homo sapi	NOLA1,GA	-1.38175	1.64E-06	0.72372	19489.84	14158.7
A_24_P144149		Unknown	ENST0000	-1.38189	7.90E-07	0.72364	2343.533	1693.156
A_32_P43	NM_02407	hypothetica	DDX54,DP	-1.38201	9.88E-06	0.72358	31784.81	23091.35
A_24_P33	NM_14507	Homo sapi	HTRA2,OM	-1.3822	9.73E-08	0.72349	2109.653	1530.13
A_24_P21	NM_03288	U7 snRNP-	LSM10,MS	-1.38253	0.00034	0.72331	1902.675	1386.111
A_24_P16	NM_01538	hypothetica	DJ328E19.	-1.38269	0.00013	0.72323	1986.576	1430.67
A_32_P11723		Unknown	ENST0000	-1.38331	8.31E-13	0.7229	37002.27	26704.01
A_24_P53	NM_00130	carboxypep	CPD	-1.38338	3.06E-06	0.72287	1604.238	1156.699
A_23_P38	NM_01812	Homo sapi	ELAC2,EL	-1.3836	0.00001	0.72275	15292.9	11131.37
A_24_P14	NM_03263	hypothetica	PAPOLA,P	-1.38362	0.00017	0.72274	4806.06	3481.331
A_24_P16	XM_37138	AG1 protei	AG1	-1.38366	1.99E-07	0.72272	10133.77	7285.649
A_23_P42	NM_00565	Homo sapi	TCF20,AR	-1.38368	1.84E-06	0.72271	1267.536	915.3274
A_23_P15	NM_01232	Homo sapi	LSM4,YER	-1.38391	4.69E-06	0.72259	13700.57	9969.313
A_24_P15	NM_00204	glycyl-tRNA	GARS,CM	-1.38398	3.31E-08	0.72255	3861.483	2794.903
A_23_P15	NM_08064	Homo sapi	APEX1,AP	-1.38418	4.80E-07	0.72245	1561.977	1131.54
A_32_P13	AB014563	KIAA0663	KIAA0663	-1.38431	4.20E-06	0.72238	2387.13	1729.49
A_32_P41	NM_00337	voltage-dep	VDAC2,FL	-1.38438	3.61E-06	0.72234	36340.45	26327.98
A_24_P11	NM_01431	host cell fa	KLHDC2,L	-1.3845	0.00006	0.72228	2427.754	1747.815
A_23_P29	NM_00501	Homo sapi	OXA1L	-1.38452	6.25E-06	0.72227	3248.447	2342.267
A_23_P25	NM_02278	hypothetica	FLJ13910,	-1.38468	4.30E-07	0.72219	1440.212	1042.123
A_23_P63	NM_01544	DKFZP586	C1orf43,NI	-1.38477	1.31E-11	0.72214	13260.65	9583.092
A_23_P21	AL831922	DKFZP586	DLG1	-1.38555	0.00024	0.72173	1481.425	1083.909
A_23_P12	NM_01328	Homo sapi	RBM15B,H	-1.38566	0.00043	0.72168	4491.907	3268.066
A_32_P78	NM_00457	phosphose	PSPH,PSF	-1.3857	2.04E-08	0.72165	1677.469	1211.653
A_24_P32	NM_01561	WIPI49-like	WIPI-2	-1.38586	1.70E-10	0.72157	11757.68	8494.904
A_32_P16	NM_00496	high-mobili	HMGN1,HI	-1.38606	1.18E-14	0.72147	38503.09	27770.84
A_23_P34	NM_01651	putative se	SCLY,SCL	-1.38619	1.04E-06	0.7214	2139.629	1539.238
A_23_P11	NM_01417	Homo sapi	UBE2T,HS	-1.38622	6.17E-07	0.72139	22724.07	16511.6
A_23_P49	NM_00692	splicing fac	SFRS1,AS	-1.3864	1.01E-07	0.72129	2345.892	1697.061
A_23_P11	NM_03328	Homo sapi	TBL1Y,TBI	-1.38649	6.77E-07	0.72125	1497.49	1078.053
A_23_P38	NM_01629	heat shock	TRAP1,HS	-1.38659	4.71E-08	0.7212	2623.22	1891.99
A_24_P64	NM_01475	KIAA0187	BMS1L,KIA	-1.38678	0.00014	0.72109	1074.25	781.6502
A_24_P75399		Unknown	A_24_P75	-1.38699	0.00016	0.72099	52769.13	38358.79
A_23_P36	NM_15236	EID-2-like i	EID-3,FLJ3	-1.38706	0.0005	0.72095	759.4291	545.5017
A_23_P35	NM_01524	Homo sapi	MGRN1,GI	-1.38713	0.00053	0.72091	6296.23	4584.78
A_24_P40	NM_01465	KIAA0446	KIAA0446,	-1.38743	4.43E-10	0.72076	3436.943	2475.775
A_32_P28	AL713754	Hypothetic	DKFZp667	-1.38758	0.00023	0.72068	965.6083	689.5936
A_32_P83	NM_18337	hypothetica	LOC20003	-1.38767	3.73E-10	0.72063	18576.89	13357.5
A_24_P39	NM_15364	Homo sapi	TPM3,TRK	-1.38775	9.30E-10	0.72059	17845.75	12886.63
A_23_P25	NM_02210	Homo sapi	C6orf79,FL	-1.38785	0.00012	0.72054	1654.752	1182.601
A_23_P13	NM_02207	hypothetica	MANBAL	-1.38786	2.54E-10	0.72053	2846.401	2051.58
A_32_P123		ADT2_HUI	THC19056	-1.38787	5.65E-13	0.72053	42120.73	30372.66

A_23_P14	NM_00682	transmemb	TMED10,T	-1.3882	7.32E-09	0.72036	13843.11	9995.089
A_23_P21	AK092776	Hypothetic	MGC16471	-1.38823	2.51E-08	0.72034	3374.064	2431.364
A_23_P70	NM_01883	Homo sapi	MATR3,MC	-1.38823	1.16E-07	0.72034	5368.987	3873.242
A_24_P303193		Unknown	THC18193	-1.3883	0.00093	0.7203	726.2028	525.5406
A_24_P39	NM_01902	Homo sapi	TMCO1,PC	-1.38836	2.61E-13	0.72027	13912.25	10020.03
A_32_P11	AB029025	KIAA1102	KIAA1102	-1.38856	5.21E-10	0.72017	7420.434	5345.349
A_23_P35	NM_13838	hypothetic	LOC92305	-1.38856	0.00004	0.72017	1525.918	1090.69
A_23_P10	NM_00409	Homo sapi	ECHS1,SC	-1.38866	0.00003	0.72012	30959.86	22118.16
A_23_P32	NM_15320	hypothetic	MGC35048	-1.38873	1.70E-06	0.72008	1100.959	794.0101
A_24_P178154		Unknown	ENST0000	-1.38875	2.01E-13	0.72007	16339.19	11770.47
A_24_P39	NM_18174	Homo sapi	ORC5L,OF	-1.38891	0.00098	0.71999	963.6047	703.2854
A_24_P24	NM_02163	kelch-like p	KLHL12,DI	-1.38895	0.00004	0.71997	1422.777	1015.994
A_24_P34	NM_03226	hypothetic	AE2,DKFZ	-1.38904	5.43E-10	0.71992	14254.01	10227.86
A_23_P13	NM_00418	Homo sapi	UCHL1,PA	-1.3892	0.00002	0.71984	192404.8	139737.3
A_23_P13	NM_00263	Homo sapi	PHB	-1.3894	6.38E-13	0.71973	5195.558	3740.503
A_24_P152845		Unknown	A_24_P15	-1.38949	0.00004	0.71969	18976.45	13704.39
A_23_P61	NM_00445	Homo sapi	ETFDH,MA	-1.38958	1.55E-12	0.71964	3729.662	2684.323
A_23_P16	NM_01595	PTD013 pr	RWDD1,P	-1.38962	0.00003	0.71962	6594.686	4763.513
A_23_P13	NM_00679	Homo sapi	AFG3L2	-1.38964	0.00034	0.71961	1002.342	728.1965
A_23_P15	NM_00029	Homo sapi	PHKA2,PH	-1.38978	0.00014	0.71954	1013.651	737.5422
A_23_P11	NM_00666	Homo sapi	PGRMC1,M	-1.38981	6.05E-06	0.71952	7706.045	5571.814
A_23_P13	NM_00698	Homo sapi	MAGED1,M	-1.38999	2.26E-12	0.71943	8822.747	6350.253
A_24_P32	NM_01602	CGI-79 pro	RBMX2,CC	-1.39009	6.08E-07	0.71938	1962.428	1417.529
A_23_P10	NM_01648	Hypothetic	LOC51249	-1.39021	0.00019	0.71931	838.9248	606.1825
A_23_P13	NM_03232	K+ channe	BTBD10,G	-1.39034	0.00039	0.71925	927.1495	676.782
A_32_P45	NM_03075	t-complex	TCP1,CCT	-1.39071	3.03E-10	0.71906	6173.285	4450.782
A_24_P11	NM_00116	baculoviral	BIRC2,API	-1.39081	0.00008	0.719	1107.679	791.5356
A_24_P12	NM_13907	Homo sapi	MAPKAPK	-1.39085	2.28E-06	0.71899	5677.903	4100.767
A_32_P11	NM_01234	F-box only	FBXO9,FB	-1.39099	3.21E-09	0.71891	1699.703	1221.729
A_23_P15	NM_01436	Homo sapi	HIBCH,HIE	-1.39124	3.98E-08	0.71878	11301.58	8126.097
A_23_P11	NM_00368	Homo sapi	AKR7A2,A	-1.39129	0.00007	0.71876	5670.874	4090.535
A_23_P12	NM_01214	Homo sapi	DTYMK,CD	-1.39154	2.52E-06	0.71863	33361.11	24103.16
A_24_P33	NM_00676	tyrosine 3-r	YWHAE,M	-1.39173	0.00083	0.71853	991.2701	702.7521
A_23_P14	NM_03329	T-cell activ	MRFAP1,F	-1.39173	4.65E-13	0.71853	7559.284	5432.211
A_32_P194095		ALU5_HUM	THC20037	-1.39204	0.00061	0.71837	1912.047	1388.28
A_24_P57	NM_13917	Homo sapi	SLC39A11	-1.39227	0.00009	0.71825	971.7698	698.9824
A_23_P14	NM_00378	Homo sapi	B4GALT2,I	-1.39242	5.05E-10	0.71817	2068.098	1485.221
A_23_P13	NM_02107	Homo sapi	NDUFV2	-1.39249	1.42E-10	0.71814	17877.88	12799.57
A_24_P14	XM_06574	Homo sapi	ENST0000	-1.39276	0.00001	0.718	1418.063	1022.906
A_24_P38	NM_14478	Homo sapi	DEGS1,ML	-1.39285	0.00006	0.71795	2515.232	1810.188
A_23_P51	NM_00452	Homo sapi	MGST3,GS	-1.39308	1.13E-12	0.71783	53771.59	38658.75
A_23_P68	NM_03211	hypothetic	WDR54,FL	-1.39322	3.58E-09	0.71776	2913.648	2091.813
A_23_P11	NM_00201	Homo sapi	FKBP3,PP	-1.39346	0.00014	0.71764	15341.07	11107.38
A_23_P14	NM_00412	Homo sapi	GTF2F2,B	-1.3936	0.00042	0.71757	3735.198	2688.036
A_23_P10	NM_00031	Homo sapi	PRNP,CJD	-1.39362	0.00045	0.71756	13753.81	9993.506
A_23_P15	NM_01834	Homo sapi	RSAD1,FL	-1.39371	7.87E-08	0.71751	2027.738	1457.078
A_24_P37	NM_18279	Homo sapi	TMSL4	-1.39387	5.50E-07	0.71743	184556.8	131499.5
A_32_P19	NM_00182	CDC28 pro	CKS1B,CK	-1.394	3.08E-06	0.71736	81700.45	58852.76
A_23_P20	NM_00556	Homo sapi	STMN1,La	-1.39402	5.50E-10	0.71735	17373.27	12475.57
A_23_P38	AY359048	Thioredoxi	TXNDC4	-1.39405	3.00E-10	0.71733	2746.226	1968.53
A_23_P12	NM_00224	Homo sapi	KCNK1,DF	-1.39431	0.00048	0.7172	2473.069	1764.864
A_24_P584992		Unknown	A_24_P58	-1.3945	4.29E-09	0.7171	1720.524	1233.429

A_24_P88	NM_00471	cytochrome	COX7A2L,	-1.39461	7.64E-09	0.71705	5330.538	3809.409
A_23_P60	NM_00038	Homo sapi	XPA,XP1,X	-1.39474	0.00009	0.71698	1255.107	908.364
A_24_P28	AL080183	Similar to K	LOC38768	-1.39475	4.68E-08	0.71697	1545.537	1108.099
A_23_P34	NM_01779	Homo sapi	RPP25,FL	-1.39481	5.20E-12	0.71694	5590.258	4010.547
A_23_P71	NM_00071	Homo sapi	BLVRA,BL	-1.39485	4.87E-13	0.71693	19059.03	13653.08
A_24_P54	AK026938	Hypothetic	LOC38983	-1.39493	2.15E-06	0.71688	2233.35	1592.901
A_24_P75	AL049461	MRNA; cDI	ENST0000	-1.39496	0.00002	0.71687	1271.25	907.2763
A_23_P41	NM_01838	Homo sapi	WDR33,W	-1.39497	1.65E-08	0.71686	2013.474	1440.381
A_32_P14	AI204246	Transmem	TMED4	-1.39499	0.00049	0.71685	2033.185	1472.583
A_23_P41	NM_02122	DC2 protei	DC2	-1.39527	6.80E-11	0.71671	11807.24	8474.102
A_24_P23	NM_01795	hypothetic	FLJ20297,	-1.39533	0.00001	0.71667	10088.84	7268.631
A_32_P46700		Unknown	A_32_P46	-1.39533	0.00051	0.71668	1180.726	856.6936
A_24_P101072		Unknown	ENST0000	-1.39548	2.04E-13	0.7166	15396.41	11037.36
A_23_P34	NM_15225	leucine-rich	MGC45866	-1.39585	0.00014	0.71641	849.2474	605.9343
A_23_P32593		Unknown	A_23_P32	-1.39602	9.50E-11	0.71632	3197.692	2290.83
A_24_P34	BC004103	GA binding	GABPB2	-1.39609	0.00026	0.71629	833.3853	600.1007
A_24_P749042		Unknown	A_24_P74	-1.39609	3.70E-11	0.71629	6149.546	4408.122
A_24_P32	NM_01607	CGI-141 pr	GOLT1B,G	-1.39634	0.00021	0.71616	970.6139	689.8806
A_23_P14	NM_01598	Homo sapi	UCHL5,UC	-1.39639	9.44E-12	0.71613	11171.26	8026.99
A_24_P98	NM_00735	guanine nu	GNA12,RM	-1.3966	8.00E-10	0.71603	2413.804	1728.869
A_24_P78	BC041856	LOC44015	BC041856	-1.39665	3.08E-06	0.716	1840.339	1323.055
A_23_P45	NM_00329	Homo sapi	TPR	-1.39674	2.07E-12	0.71595	6226.998	4462.905
A_24_P46	NM_00274	protein kina	PRKCI,PK	-1.39692	0.00003	0.71586	1558.758	1125.892
A_24_P20	NM_03057	hypothetic	NDFIP1,N	-1.39707	1.42E-09	0.71578	1873.804	1340.647
A_24_P94	NM_01510	Homo sapi	PHF8,ZNF	-1.3972	7.52E-08	0.71572	3894.43	2784.901
A_24_P80	AJ004914	Tmp21-II ,	Tmp21-II	-1.39738	3.43E-08	0.71562	1868.426	1338.354
A_24_P94	AB002330	U2-associa	SR140	-1.39762	2.31E-07	0.7155	1314.084	941.0087
A_32_P13	NM_00258	poly(A)-spe	PARN,DAN	-1.39787	2.95E-07	0.71537	1871.885	1341.605
A_23_P11	NM_01215	Homo sapi	F8A1,F8A,	-1.39788	3.93E-08	0.71537	7934.887	5673.541
A_24_P13	NM_18337	hypothetic	LOC20003	-1.39801	3.89E-07	0.7153	13338.05	9480.078
A_24_P42	NM_01233	acyl-Coenz	ACATE2,C	-1.39845	1.27E-06	0.71508	2427.631	1739.091
A_23_P34	AL833885	Egl nine ho	EGLN1	-1.3985	1.04E-07	0.71505	4171.545	2988.479
A_24_P37	NM_01471	CP110 pro	CP110,KIA	-1.39852	0.0002	0.71504	1691.571	1230.144
A_23_P13	NM_01450	malonyl-Co	MT,MCT,M	-1.39867	0.00002	0.71496	1616.168	1164.067
A_24_P58	NM_15364	Homo sapi	TPM3,TRK	-1.39871	6.76E-07	0.71495	7854.591	5649.791
A_32_P84	NM_02247	germ cell-le	GMCL1L,G	-1.3989	7.53E-06	0.71485	1361.263	967.6814
A_24_P31	NM_00574	protein dis	PDIA6,P5,I	-1.39897	6.98E-06	0.71481	5782.913	4145.233
A_24_P31	AF447582	Casein kina	CSNK1A1	-1.39905	4.45E-10	0.71477	3267.85	2336.729
A_24_P33429		Unknown	ENST0000	-1.39907	2.71E-13	0.71476	28489.79	20348.11
A_23_P68	NM_00619	Homo sapi	PCBP1,HN	-1.39914	0.00009	0.71472	99806.77	71672.25
A_23_P11	NM_00588	Membrane	MARCH-VI	-1.39914	1.97E-08	0.71473	2840.466	2034.846
A_24_P58	NM_01910	beta-1,4 m	ALG1,HMT	-1.3992	1.92E-08	0.7147	2748.85	1961.849
A_23_P44	NM_01612	Homo sapi	COPG,COI	-1.39921	1.29E-06	0.71469	1332.427	950.1326
A_23_P69	NM_01844	30 kDa pro	LOC55831	-1.39928	3.20E-07	0.71465	17652.43	12650.75
A_23_P25	NM_00162	Homo sapi	AKR1B1,A	-1.39934	7.83E-15	0.71462	83733.59	59877.38
A_23_P61	NM_01233	Homo sapi	MSRA	-1.39953	0.00004	0.71453	1342.523	951.648
A_23_P47	NM_02093	Homo sapi	DDX55,KIA	-1.39978	0.00023	0.7144	1625.454	1150.771
A_23_P20	NM_00025	Homo sapi	MTR	-1.3999	4.79E-15	0.71433	7402.634	5288.087
A_23_P10	NM_01439	PTD010 p	GHITM,DE	-1.39997	2.22E-10	0.7143	14971.71	10660.24
A_24_P23	NM_05296	Homo sapi	C1orf24,NI	-1.40004	0.00009	0.71426	1079.488	778.0045
A_23_P11	NM_00004	Homo sapi	AR,KD,AIS	-1.40015	2.09E-09	0.71421	3114.92	2224.69
A_24_P127425		Unknown	ENST0000	-1.40023	3.02E-07	0.71417	2243.91	1605.182

A_23_P88	NM_14460	Homo sapi	CKLFSF3,I	-1.40025	4.12E-07	0.71416	4089.435	2911.98
A_23_P41	NM_00645	Homo sapi	PAICS,AIR	-1.40045	6.41E-07	0.71406	29493.69	21092.33
A_23_P31	NM_01336	Homo sapi	ANAPC4,A	-1.40049	1.70E-07	0.71404	2852.318	2034.357
A_23_P12	NM_01764	Homo sapi	TRIT1,IPT,	-1.40065	1.35E-07	0.71395	1659.709	1182.667
A_24_P12	NM_02075	Homo sapi	SERINC1,	-1.4008	5.17E-10	0.71388	2025.287	1446.125
A_24_P41	NM_00196	eukaryotic	EIF5,EIF-5	-1.40117	0.00004	0.71369	22376.47	16092.54
A_24_P255303		Unknown	A_24_P25	-1.40128	1.61E-13	0.71363	12591.23	8973.111
A_24_P34	NM_00196	eukaryotic	EIF4E,CBF	-1.40156	7.14E-06	0.71349	1285.953	924.9358
A_32_P17	NM_01603	CGI-97 pro	SBDS,SDS	-1.40168	4.25E-13	0.71343	3791.329	2705.189
A_23_P20	NM_03232	Homo sapi	C1orf57,FL	-1.40177	4.08E-11	0.71338	2403.589	1714.677
A_32_P211752		Unknown	THC18206	-1.40199	3.27E-06	0.71327	1409.994	1001.574
A_23_P70	NM_01603	CGI-97 pro	SBDS,SDS	-1.40199	2.60E-08	0.71327	3808.169	2712.072
A_23_P14	NM_14890	Homo sapi	OSBPL9,O	-1.40209	1.78E-09	0.71322	2999.458	2138.782
A_23_P13	NM_02072	neurolysin	NLN,AGTB	-1.40242	3.54E-09	0.71305	1973.296	1405.799
A_23_P13	NM_01814	Homo sapi	RNMTL1,H	-1.40245	7.78E-07	0.71304	4740.56	3386.856
A_23_P40	NM_15227	Homo sapi	FAM58A,M	-1.40257	1.69E-11	0.71298	11475.65	8198.789
A_24_P24806		Unknown	A_24_P24	-1.40257	0.00028	0.71298	1850.604	1325.39
A_24_P28	AB002387	Myosin VI	MYO6	-1.40258	0.00026	0.71297	753.8912	540.1737
A_24_P84112		Unknown	A_24_P84	-1.403	0.00002	0.71276	20167.67	14356.93
A_23_P45	NM_00313	signal reco	SRP9,ALU	-1.4034	3.94E-09	0.71255	7142.402	5110.699
A_24_P24	NM_02461	NOD9 prot	NOD9,FLJ	-1.40342	3.89E-07	0.71254	1830.227	1310.875
A_23_P72	NM_02520	Homo sapi	MED28,EG	-1.40347	8.87E-06	0.71252	2032.623	1443.658
A_24_P39	NM_02064	chromoson	TMEM9B,C	-1.40391	0.00009	0.7123	970.801	696.6092
A_32_P38	NM_15381	Chromosor	C10orf46	-1.40417	0.00003	0.71216	1431.09	1011.448
A_32_P58	NM_15381	Homo sapi	PHF13,PH	-1.40457	0.00053	0.71196	3091.876	2222.715
A_23_P33	NM_18170	hypothetica	LOC14423	-1.40463	9.95E-13	0.71193	2633.674	1875.249
A_23_P21	NM_01774	Homo sapi	ANKHD1,M	-1.40465	0.00002	0.71192	1459.672	1033.676
A_23_P31	NM_00511	Homo sapi	GOLGA5,F	-1.40473	3.62E-07	0.71188	1920.503	1362.78
A_32_P63	NM_00531	glucose re	PDIA3,P58	-1.40482	5.58E-09	0.71183	10973.34	7780.491
A_24_P93	AK095167	CDNA FLJ	AK095167	-1.40504	0.00032	0.71173	1031.965	743.6699
A_24_P50	AK074291	Homeodon	HIPK2	-1.40509	2.70E-06	0.7117	1631.23	1155.877
A_32_P91156		AF030694	THC18524	-1.4051	0.0001	0.71169	988.0883	708.4916
A_32_P20454		ALU8_HUM	THC18449	-1.40529	5.16E-09	0.7116	1773.42	1262.714
A_23_P11	NM_00241	Homo sapi	MGST2,GS	-1.40531	3.79E-08	0.71159	6129.531	4368.387
A_24_P15	NM_01846	Homo sapi	C3orf10,MI	-1.40553	4.61E-08	0.71148	3143.361	2239.357
A_23_P41	NM_00164	N-acylamin	APEH,APH	-1.40555	0.00002	0.71147	1153.578	826.6775
A_32_P12	BE018697	Similar to F	BE018697	-1.40556	6.89E-08	0.71146	1728.514	1231.287
A_23_P11	NM_00619	proliferatio	PA2G4,EB	-1.40558	1.13E-12	0.71145	24530.46	17428.12
A_24_P56	AF086032	Protein pho	PPP1R12A	-1.40626	1.17E-06	0.71111	2283.242	1619.099
A_24_P16	AK001770	Poly (ADP-	PARP14	-1.40631	0.00052	0.71108	917.9565	662.6704
A_23_P73	NM_00058	Homo sapi	SPP1,OPN	-1.40631	3.09E-15	0.71108	206073.8	146733.1
A_23_P83	NM_01884	Homo sapi	KLHL9,RP	-1.40637	7.66E-10	0.71105	3365.446	2392.343
A_24_P14	NM_01777	Homo sapi	PXK,FLJ20	-1.40663	0.00008	0.71092	1183.426	846.3554
A_23_P36	NM_02463	Sin3A asso	SAP30L,FL	-1.40674	3.79E-14	0.71086	3939.261	2800.256
A_23_P92	NM_00222	Homo sapi	ITPR1,IP3R	-1.40689	0.00001	0.71079	2386.021	1692.114
A_23_P50	NM_01531	Homo sapi	ARHGEF1	-1.40692	0.00015	0.71077	1793.828	1261.962
A_32_P56434		Unknown	A_32_P56	-1.407	0.0006	0.71073	2507.452	1780.569
A_24_P31	NM_01500	Homo sapi	EXOSC7,p	-1.40727	0.00002	0.7106	5868.661	4202.033
A_23_P10	NM_00039	Homo sapi	COL5A2,M	-1.40769	3.51E-07	0.71039	2265.231	1613.039
A_23_P73	NM_02409	Homo sapi	ELOVL6,F	-1.40776	0.00095	0.71035	1402.589	1022.225
A_32_P12	AB011542	EGF-like-d	EGFL5	-1.40792	0.00005	0.71027	884.3033	628.4979
A_32_P17	NM_01693	polymerase	POLA	-1.40793	2.44E-14	0.71026	4705.187	3341.667

A_23_P20	NM_00383	Homo sapi	FPGT,GFP	-1.40801	0.00042	0.71022	762.4599	537.8134
A_23_P20	NM_01593	CGI-01 pro	KIAA0859,	-1.40803	0.00018	0.71021	1126.174	791.4664
A_23_P12	NM_00700	Homo sapi	NUDT21,C	-1.40808	1.95E-15	0.71019	13052.66	9267.59
A_24_P24	AF060925	Hematolog	HN1	-1.40828	5.24E-09	0.71009	1874.453	1329.967
A_23_P68	NM_00312	sepiapterin	SPR	-1.40833	3.80E-12	0.71006	4758.194	3378.198
A_32_P84	BX640902	KIAA0888	KIAA0888	-1.40833	2.83E-06	0.71006	1117.113	795.0023
A_24_P25	NM_20332	Homo sapi	SLC23A2,N	-1.40839	0.00011	0.71003	888.7677	633.8623
A_23_P12	S72869	Coiled-coil	CCDC6	-1.4085	0.00003	0.70998	1750.074	1256.373
A_23_P87	NM_03284	hypothetica	FLJ14827	-1.40866	9.37E-09	0.70989	5931.031	4203.69
A_23_P20	NM_00169	Homo sapi	AUH	-1.40884	3.21E-07	0.70981	2222.655	1573.97
A_32_P160186		Unknown	A_32_P16	-1.40893	3.60E-06	0.70976	1668.33	1190.084
A_23_P15	NM_00181	Homo sapi	CTSC,HMS	-1.40907	6.22E-08	0.70969	2317.368	1654.02
A_23_P59	NM_00549	Homo sapi	RANBP9,R	-1.40908	1.06E-08	0.70968	4118.321	2923.985
A_23_P21	NM_01593	CGI-09 pro	CGI-09,MG	-1.40913	2.27E-11	0.70966	3134.995	2225.197
A_24_P28	AB002303	Zinc finger,	ZFYVE16	-1.40913	0.00081	0.70966	732.1992	523.5762
A_24_P33	NM_14520	Homo sapi	CSNK1A1L	-1.40914	0.00078	0.70965	2065.314	1456.366
A_24_P28	NM_03235	hypothetica	MGC10744	-1.40963	2.25E-08	0.70941	4019.383	2854.104
A_23_P85	NM_02283	Homo sapi	ARMCX5,F	-1.41023	0.00064	0.70911	2464.83	1742.354
A_24_P83	AK026647	CDNA: FL	AK026647	-1.41027	0.00001	0.70908	3550.907	2515.455
A_23_P75	NM_00337	Homo sapi	VDAC2,FL	-1.41085	1.80E-07	0.70879	55258.97	39336.01
A_24_P40	NM_03092	integral me	ITM2C,E25	-1.41094	0.00023	0.70875	3053.084	2159.334
A_24_P28	NM_00280	proteasom	PSMD3,S3	-1.41115	1.48E-07	0.70864	3398.916	2406.622
A_23_P25	NM_01407	Homo sapi	NCOA6,NF	-1.41122	5.60E-09	0.70861	2196.844	1555.923
A_32_P76060		Unknown	THC18807	-1.41146	8.69E-06	0.70848	1617.852	1140.992
A_24_P79	AK000866	Family with	FAM36A	-1.41146	0.00014	0.70849	870.6564	618.1128
A_23_P59	AB002329	Sema dom	SEMA3E	-1.41158	0.00089	0.70843	833.5049	582.1807
A_23_P69	NM_00524	Homo sapi	FAT,ME5,F	-1.41159	6.08E-09	0.70842	3488.482	2468.961
A_24_P13	NM_01840	hypothetica	HSA27219	-1.41159	3.04E-14	0.70842	9237.166	6538.893
A_23_P85	NM_00662	phosphogl	PHGDH,PD	-1.41196	3.47E-14	0.70823	4905.949	3473.987
A_23_P71	NM_01829	Homo sapi	PGM2,MST	-1.41241	8.01E-10	0.70801	5003.527	3542.979
A_24_P264004		Unknown	A_24_P26	-1.41266	0.00003	0.70788	1523.375	1094.118
A_32_P202708		Unknown	A_32_P20	-1.41266	0.00013	0.70788	930.2346	662.2626
A_32_P67	AB058701	L(3)mbt-lik	L3MBTL3	-1.41269	0.00052	0.70787	796.5387	569.6308
A_23_P10	AK024487	Zinc finger	ZNF598	-1.41271	0.00015	0.70786	18332.81	13069.04
A_24_P560909		ALU1_HUM	THC19132	-1.41283	4.13E-12	0.7078	3017.021	2134.492
A_23_P68	NM_00372	Homo sapi	DSCR2,C2	-1.41299	1.64E-15	0.70772	27555.38	19492.35
A_23_P10	NM_01640	Ufm1-conju	UFC1,HSP	-1.41324	4.40E-09	0.70759	4508.094	3181.675
A_24_P14	NM_00472	DEAD (Asp	DDX21	-1.41337	2.36E-09	0.70753	14706.13	10422.62
A_23_P48	NM_00028	Homo sapi	PCCA	-1.41343	0.00014	0.7075	1131.549	813.17
A_24_P37	AK055071	Phosphatid	PIGK	-1.41354	4.60E-07	0.70744	1279.452	904.2461
A_23_P25	NM_01793	Hypothetica	FLJ20699	-1.41361	0.00005	0.70741	1492.964	1048.626
A_24_P59239		Unknown	ENST0000	-1.41365	1.93E-11	0.70739	27547.58	19541.46
A_23_P31	NM_01864	Homo sapi	CHST12,C	-1.41398	0.00001	0.70723	1256.321	882.202
A_23_P86	NM_00279	Homo sapi	PSMA5,PS	-1.41398	2.31E-10	0.70723	30022.01	21293.3
A_24_P17	AK022463	Homo sapi	FLJ12649,	-1.4141	0.00021	0.70716	1110.899	776.2667
A_23_P10	NM_02102	Homo sapi	CFHL3,FH	-1.4141	0.00068	0.70716	728.1184	518.4577
A_23_P58	NM_00090	Homo sapi	NQO2,QR2	-1.41416	1.42E-07	0.70713	8282.454	5813.734
A_23_P93	NM_01610	Homo sapi	AIG1,AIG-1	-1.41432	0.0001	0.70705	1648.749	1155.599
A_24_P33	NM_02508	hypothetica	FLJ21128	-1.41435	3.89E-06	0.70704	1391.37	990.3341
A_23_P38	NM_02079	Homo sapi	TAOK1,PS	-1.41438	4.72E-06	0.70702	1188.846	843.532
A_23_P13	NM_03298	Homo sapi	TBL2,WBS	-1.41444	4.11E-16	0.707	7767.648	5492.262
A_23_P10	NM_01610	Homo sapi	NAT5,ARD	-1.41466	3.45E-14	0.70688	14461.7	10235.05

A_23_P75	NM_03281	Homo sapi	ATAD1,AF	-1.41469	2.36E-12	0.70687	3819.541	2700.94
A_24_P19	NM_01900	Spindlin fa	SPIN2	-1.4148	3.38E-07	0.70682	1488.372	1048.538
A_24_P23	NM_00279	proteasom	PSMB1,HC	-1.41513	2.59E-11	0.70665	70635.38	49968.57
A_24_P567349		ALU8_HUM	THC18703	-1.41542	0.00076	0.70651	765.5662	534.0464
A_24_P25	NM_01847	uncharacte	ACTR10,A	-1.41546	1.49E-10	0.70649	3254.916	2298.327
A_24_P12	NM_01822	parvin, alp	PARVA,MX	-1.41558	7.08E-06	0.70643	1425.578	1000.206
A_23_P39	NM_00193	Homo sapi	DR1,NC2,M	-1.41567	5.21E-09	0.70638	3271.985	2310.889
A_24_P39	XM_37296	Homo sapi	ENST0000	-1.41582	7.99E-12	0.70631	9376.757	6643.42
A_32_P67	BC067908	CDNA clon	BC067908	-1.41586	0.0002	0.70629	1204.037	842.0392
A_24_P29	NM_17380	Homo sapi	BLOC1S2,	-1.41615	7.88E-08	0.70614	10165.29	7215.002
A_24_P91	NM_00682	transmemb	TMED10,T	-1.41617	3.80E-11	0.70613	5198.936	3670.204
A_24_P18	NM_14505	hypothetica	RP11-311F	-1.41622	2.91E-08	0.7061	1704.662	1206.347
A_23_P24	NM_00001	Homo sapi	ACAT1,T2,	-1.41633	2.24E-06	0.70605	8219.546	5855.307
A_23_P42	NM_01806	Homo sapi	C6orf166,F	-1.41644	3.97E-15	0.70599	6939.369	4898.35
A_23_P16	NM_00126	cyclin-depe	CDK9,TAK	-1.41652	5.06E-12	0.70596	2861.478	2018.519
A_24_P460195		AY173017	THC18890	-1.41654	0.00002	0.70595	1566.809	1114.37
A_24_P38	NM_01414	HSPC274	C20orf30,H	-1.41674	4.26E-06	0.70585	7790.734	5519.517
A_32_P10	NM_20731	similar to a	LOC11271	-1.41676	0.00004	0.70584	36751.75	25670.59
A_23_P92	NM_02526	Homo sapi	TSEN2,SE	-1.41693	1.59E-09	0.70575	4067.264	2872.816
A_23_P12	NM_13906	Homo sapi	SMARCC2	-1.41698	1.87E-15	0.70573	13357.78	9430.964
A_23_P20	NM_00271	Homo sapi	PPP2R5C,	-1.41707	2.54E-10	0.70568	3351.192	2364.13
A_23_P13	NM_01404	Homo sapi	ACAD9,AC	-1.4172	5.52E-06	0.70562	7715.519	5486.333
A_23_P27	NM_00727	Homo sapi	GABARAP	-1.41728	1.65E-08	0.70558	3242.762	2285.319
A_23_P82	NM_01829	Homo sapi	QRSL1,Ga	-1.41736	0.00009	0.70554	1607.881	1146.894
A_23_P11	NM_13046	jun dimeriz	JDP2,JUNI	-1.41742	4.99E-08	0.70551	7101.642	5024.635
A_23_P11	NM_00043	Homo sapi	PAFAH1B1	-1.41763	1.84E-09	0.7054	5613.654	3967.498
A_24_P19	AF301222	NY-REN-4	NY-REN-4	-1.41768	0.00004	0.70538	4485.889	3179.369
A_32_P17	AK123297	Zinc finger	ZNF37B	-1.41809	0.00005	0.70518	1472.99	1051.07
A_24_P157720		Unknown	A_24_P157	-1.41814	0.00003	0.70515	1181.148	843.5722
A_24_P66932		Unknown	ENST0000	-1.41825	0.00003	0.7051	1709.629	1195.855
A_24_P272886		Unknown	ENST0000	-1.4183	1.72E-08	0.70507	2883.528	2035.15
A_32_P75	NM_15227	Homo sapi	FAM58A,M	-1.41883	5.91E-07	0.70481	6649.634	4706.293
A_32_P14	NM_00693	SMT3 (sup	SUMO2,HS	-1.41895	1.39E-11	0.70475	24108.86	17013.23
A_24_P33	D31888	REST core	RCOR1	-1.41898	3.07E-07	0.70473	2264.51	1593.593
A_24_P67	NM_00583	serine/argi	SRRM1,16	-1.41917	0.00003	0.70464	1122.343	784.5369
A_23_P11	L13689	Polycomb	PCGF4	-1.41932	4.42E-10	0.70456	3607.337	2541.103
A_24_P83	NM_00327	transmemb	TSPAN6,T	-1.41934	0.00006	0.70455	1408.067	1000.485
A_23_P10	NM_00464	Chromosor	C21orf33	-1.41957	1.92E-15	0.70444	6100.372	4298.442
A_23_P25	NM_01791	hypothetica	FLJ20647	-1.4198	0.00002	0.70433	1643.283	1165.281
A_23_P44	NM_01325	Homo sapi	TBK1,NAK	-1.41992	2.15E-06	0.70426	3940.567	2780.826
A_32_P43	NM_00447	FSHD regi	FRG1,FSG	-1.41999	0.00056	0.70423	12326.12	8709.732
A_23_P21	NM_00191	Homo sapi	CYB561	-1.42004	0.00065	0.70421	1924.839	1367.94
A_23_P69	NM_02110	Homo sapi	PLSCR1,M	-1.42012	3.80E-11	0.70417	2599.297	1832.631
A_23_P34	NM_00669	jumping tra	JTB,PAR,h	-1.42015	9.95E-11	0.70415	29106.2	20553.71
A_23_P11	BC008430	Family with	FAM18B	-1.42019	3.94E-10	0.70413	4351.729	3068.766
A_24_P80	NM_03299	general tra	GTF2I,DIW	-1.42021	0.00041	0.70412	767.3137	537.8379
A_23_P42	NM_20346	Homo sapi	MRFAP1L1	-1.42029	1.46E-09	0.70408	2701.63	1901.965
A_23_P10	NM_02037	Homo sapi	AVEN,PDC	-1.42032	1.46E-07	0.70406	6216.869	4389.623
A_23_P61	NM_01684	neuronal S	SHC3,NSH	-1.42053	1.04E-06	0.70396	2137.946	1500.151
A_32_P15	NM_00441	desmoplak	DSP,DPI,D	-1.42054	5.35E-08	0.70396	1606.571	1133.323
A_23_P77	NM_00208	Homo sapi	GRB2,ASH	-1.42065	6.52E-07	0.7039	8086.463	5683.741
A_24_P20	NM_02099	B-cell CLL/	BCL7A,BC	-1.42066	3.07E-08	0.7039	1475.072	1037.024

A_23_P78	NM_03271	Homo sapi	MAFG,MG	-1.42073	5.32E-10	0.70386	5244.472	3697.872
A_23_P15	NM_00554	Homo sapi	KARS,KAF	-1.42082	0.00057	0.70382	1129.933	807.9731
A_23_P12	NM_00642	Homo sapi	MRPL28,p	-1.42088	1.06E-11	0.70379	8216.184	5770.879
A_23_P93	NM_00591	Homo sapi	MCM7,MC	-1.42096	7.15E-09	0.70375	54824.6	38692.69
A_23_P77	NM_00375	Homo sapi	EIF3S8,eIF	-1.42109	3.17E-15	0.70368	11603.76	8168.889
A_32_P47	NM_00534	histidine tri	HINT1,HIN	-1.42137	1.93E-09	0.70355	176946.2	125029.5
A_23_P49	NM_00276	Homo sapi	PRPSAP2,	-1.4214	2.18E-09	0.70353	2811.94	1976.409
A_23_P14	NM_00350	Homo sapi	CDC7,Hsk	-1.42157	7.04E-07	0.70345	2322.557	1637.882
A_23_P20	NM_00572	Homo sapi	PPIF,CYP3	-1.42165	1.21E-12	0.70341	76342.77	53803.81
A_23_P15	NM_01811	Homo sapi	POLR3E,S	-1.42166	3.68E-08	0.7034	2680.357	1887.595
A_23_P21	AK025603	HSPC159	HSPC159	-1.42173	1.86E-10	0.70337	1989.614	1397.993
A_24_P272653		Unknown	A_24_P27	-1.42173	3.86E-07	0.70337	2719.616	1908.787
A_23_P25	NM_01609	HSPC038	ZNF706,HS	-1.42179	1.13E-07	0.70334	17603.93	12423.38
A_23_P45	NM_00182	Homo sapi	CKS1B,CK	-1.42201	3.49E-11	0.70323	92539.92	65202.66
A_23_P58	NM_00123	Homo sapi	CCNA2,CC	-1.42201	8.95E-23	0.70323	2103.394	1483.138
A_23_P17	NM_00657	Homo sapi	EBP,CPX,(	-1.4221	6.91E-16	0.70319	28172.34	19801.48
A_24_P19	NM_19868	Homo sapi	RAB15	-1.42216	0.0001	0.70316	897.3986	627.2092
A_24_P29	NM_01531	pumilio (Dr	PUM2,PUM	-1.42217	5.34E-08	0.70315	2062.367	1448.121
A_24_P913005		AF124394	THC18472	-1.42219	0.00009	0.70314	1064.197	740.6867
A_32_P15	BX537645	Hypothetic	LOC40139	-1.4222	0.00001	0.70314	2044.654	1446.352
A_23_P25	NM_01208	Homo sapi	HDHD1A,G	-1.4222	8.53E-06	0.70313	1721.331	1221.555
A_23_P35	NM_15290	Homo sapi	NEDD1,FL	-1.42226	0.0002	0.70311	779.3382	550.9744
A_23_P325411		Unknown	A_23_P32	-1.4223	6.31E-09	0.70309	3011.594	2116.573
A_23_P10	NM_00606	Homo sapi	COX4NB,N	-1.42239	0.0001	0.70304	14300.48	10122.93
A_32_P48	NM_03250	Homo sapi	FAM11A,C	-1.42271	3.83E-07	0.70288	2984.687	2095.018
A_24_P22	NM_00439	dystroglyca	DAG1,A3a	-1.42273	1.21E-06	0.70287	1342.421	947.3183
A_32_P18	NM_01564	PAI-1 mRN	SERBP1,C	-1.42305	3.26E-08	0.70272	22627.25	15791.87
A_24_P17	NM_01206	dynamin 1-	DNM1L,DL	-1.42307	0.0001	0.70271	1178.863	821.0968
A_24_P30	NM_01899	Homo sapi	RIN2,RAS	-1.42321	0.00002	0.70264	1529.124	1065.111
A_32_P82963		Unknown	A_32_P82	-1.42339	4.04E-13	0.70255	2480.44	1742.723
A_23_P81	NM_01903	Homo sapi	DHX29,DD	-1.42345	1.95E-06	0.70252	1784.309	1268.138
A_23_P21	NM_00021	Homo sapi	ITGA6,CD4	-1.42348	2.85E-07	0.7025	3739.429	2626.497
A_24_P22	NM_02274	Homo sapi	ATPAF1,A	-1.42353	0.00017	0.70248	1191.292	826.1755
A_23_P15	NM_00468	Homo sapi	NMI	-1.42355	3.21E-12	0.70247	3753.238	2645.236
A_24_P49	BC037535	CDNA clon	BC037535	-1.42373	1.05E-07	0.70238	3480.811	2436.401
A_23_P25	NM_00187	Homo sapi	CPE	-1.42382	0.00023	0.70234	1787.258	1243.124
A_23_P18	NM_00291	Homo sapi	RFC1,A1,F	-1.42422	0.00002	0.70214	1983.472	1405.063
A_23_P25	NM_00708	Homo sapi	WDHD1,AI	-1.42427	3.73E-15	0.70211	2084.754	1466.115
A_23_P38	NM_02044	KIAA1191	KIAA1191,	-1.42435	0.00002	0.70208	976.6161	686.5432
A_23_P30	AL161991	Vasculin	DKFZp761	-1.42454	4.86E-06	0.70198	1174.179	819.9884
A_23_P20	NM_00550	Homo sapi	SCARB1,C	-1.42456	1.18E-06	0.70197	93304.48	65932.71
A_23_P35	AB051523	Hypothetic	FLJ20294	-1.42466	0.00005	0.70192	1497.942	1060.349
A_24_P33	AB011115	KIAA0543	KIAA0543	-1.42469	0.00006	0.7019	809.704	565.5752
A_23_P71	NM_02452	hypothetica	GCC1,GC	-1.42471	0.00002	0.7019	955.5757	671.2058
A_24_P12	NM_00700	cleavage a	CPSF6,CF	-1.42478	1.10E-10	0.70186	3584.236	2517.805
A_23_P43	NM_01662	Homo sapi	FAM49B,L	-1.42493	4.30E-08	0.70179	4960.185	3484.453
A_23_P23	NM_08039	Homo sapi	PTP4A2,H	-1.42506	3.68E-08	0.70172	3283.12	2304.009
A_32_P10	NM_00659	SnRNP as	USP39,SA	-1.42531	9.19E-14	0.7016	22121.48	15514.55
A_32_P16	NM_13845	hypothetica	LOC11614	-1.42538	0.00085	0.70157	1109.449	793.5997
A_23_P38	NM_00593	MHC class	MICB	-1.42542	0.00085	0.70155	958.2108	659.3656
A_23_P59	NM_02193	Rad50-inte	RINT-1,RIN	-1.42547	5.88E-09	0.70152	2207.329	1556.627
A_23_P14	NM_02469	Homo sapi	ZFAND1,F	-1.42548	1.38E-10	0.70152	2823.439	1978.073

A_23_P20	NM_01611	Homo sapi	TRPV2,VR	-1.42564	1.62E-11	0.70144	4890.906	3432.729
A_24_P41	NM_01840	Homo sapi	LAPTM4B,	-1.42574	2.31E-10	0.70139	2036.092	1429.095
A_32_P182070		AF131778	THC19336	-1.42595	2.49E-07	0.70129	2953.869	2068.163
A_23_P31	NM_01626	GIOT-3 for	ZNF12,KO	-1.42608	1.28E-08	0.70122	1662.054	1164.909
A_23_P21	NM_03143	Homo sapi	UCK1,URK	-1.42612	0.00006	0.7012	1161.271	821.9475
A_24_P66	NM_00313	signal reco	SRP9,ALU	-1.42638	7.92E-08	0.70108	7339.229	5175.459
A_32_P11	NM_01441	Siah-intera	CACYBP,S	-1.42653	1.93E-14	0.701	24876.78	17428.48
A_23_P20	NM_17574	Homo sapi	C14orf130,	-1.42674	2.82E-08	0.7009	1444.131	1010.979
A_23_P25	X66610	H.sapiens	enolase	-1.42677	0.00043	0.70088	2695.393	1887.735
A_23_P16	NM_19896	Homo sapi	AES,GRG,	-1.42688	2.67E-07	0.70083	4236.047	2976.875
A_23_P57	NM_01780	Homo sapi	CKLFSF6,I	-1.42689	8.29E-10	0.70083	10269.25	7216.925
A_23_P16	NM_15342	Homo sapi	PITX2,RS,	-1.42692	9.88E-08	0.70081	1237.249	867.243
A_32_P64928		Unknown	A_32_P64	-1.42694	1.69E-07	0.7008	2768.27	1940.261
A_24_P31	NM_00503	polo (Dros	PLK1,PLK,	-1.42714	0.00004	0.7007	18305.13	12873.45
A_23_P60	NM_01467	KIAA0103	KIAA0103	-1.42725	7.84E-07	0.70065	2070.742	1446.643
A_23_P15	NM_01847	Homo sapi	ACTR10,A	-1.42748	7.97E-13	0.70053	5585.723	3914.982
A_23_P20	NM_00611	Homo sapi	VAV3	-1.42754	2.08E-10	0.7005	3228.683	2263.037
A_23_P43	AL117511	Chromosor	C14orf125	-1.42755	0.00015	0.7005	1519.441	1051.903
A_23_P39	NM_01513	Myosin phc	M-RIP	-1.42795	5.00E-07	0.7003	1935.2	1358.899
A_24_P32	AF306679	Establishm	ESCO2	-1.42796	0.00076	0.7003	1107.516	758.431
A_23_P64	NM_00686	putative G	RABL4,RA	-1.42796	7.03E-07	0.7003	1215.387	848.2958
A_24_P58	NM_03270	tubulin alp	TUBA6,MG	-1.42798	1.07E-08	0.70029	30924.46	21549.18
A_24_P625683		Unknown	A_24_P62	-1.42798	7.68E-13	0.70029	8967.117	6290.865
A_23_P35	NM_18261	hypothetica	FLJ34283,	-1.42807	6.67E-10	0.70025	9606.766	6753.096
A_24_P38	NM_00271	protein phc	PPP1CC,P	-1.42823	5.13E-08	0.70017	6762.316	4754.03
A_32_P182395		Unknown	A_32_P18	-1.42838	6.43E-07	0.7001	1293.593	903.4435
A_23_P25	NM_01223	SREBP cle	SCAP	-1.42857	3.62E-14	0.7	5573.386	3903.241
A_24_P21	AB018326	PHD finger	PHF14	-1.42858	0.00001	0.7	3161.216	2215.74
A_23_P50	NM_14586	Homo sapi	EDARADD	-1.42862	1.91E-10	0.69998	28322.01	19762.82
A_23_P44	NM_00675	transaldola	TALDO1,T	-1.42866	7.16E-08	0.69996	30736.03	21392.09
A_23_P13	NM_02079	Arylacetam	KIAA1363	-1.42887	6.40E-12	0.69986	2908.909	2037.401
A_23_P16	NM_01327	Homo sapi	ANKRD11,	-1.42911	3.02E-09	0.69973	1752.142	1227.52
A_32_P16	NM_01539	Homo sapi	ANAPC13,	-1.42922	6.48E-12	0.69968	2852.289	1995.645
A_24_P33115		Unknown	A_24_P33	-1.42929	0.00069	0.69965	8151.745	5685.225
A_32_P11	BC040651	Isopenteny	IDI2	-1.4294	5.18E-10	0.69959	2319.036	1620.455
A_23_P15	NM_02233	Homo sapi	ITGB1BP1	-1.42965	1.40E-11	0.69947	12606.91	8803.557
A_23_P50	D31886	RAB3 GTP	ZRANB3	-1.42976	2.75E-08	0.69942	1594.928	1113.676
A_24_P315405		Unknown	A_24_P31	-1.42997	1.73E-10	0.69932	19688.57	13814.47
A_23_P30	NM_17681	Homo sapi	CHMP4B,S	-1.43024	0.00002	0.69918	4218.458	2952.735
A_23_P15	NM_01621	Homo sapi	C3orf18,G	-1.43026	0.00035	0.69918	732.2144	513.5256
A_23_P13	NM_01214	Homo sapi	DTYMK,CD	-1.43037	8.52E-12	0.69912	5901.936	4136.386
A_24_P85	NM_01223	sirtuin (sile	SIRT3,SIR	-1.4304	8.12E-12	0.69911	2266.855	1585.168
A_23_P69	NM_17804	Homo sapi	ACTL6A,A	-1.43043	9.90E-09	0.69909	10416.76	7286.39
A_23_P23	NM_01428	Homo sapi	ITGB3BP,N	-1.43045	1.26E-22	0.69908	2421.53	1693.505
A_23_P25	NM_00337	Homo sapi	VBP1,PFD	-1.43054	2.25E-07	0.69904	2184.463	1523.464
A_32_P151557		Unknown	A_32_P15	-1.4307	2.72E-09	0.69896	2135.146	1491.292
A_23_P10	NM_02475	Homo sapi	AGMAT,FL	-1.43113	0.00004	0.69875	1936.318	1345.456
A_24_P11	NM_02045	Homo sapi	C13orf1,CL	-1.43116	0.00038	0.69874	1168.219	803.7809
A_23_P169766		AF208291	THC18182	-1.43153	6.15E-06	0.69855	4226.679	2959.558
A_24_P62	NM_08092	hypothetica	LOC96610	-1.43155	0.00069	0.69854	806.2206	571.6422
A_23_P62	NM_02210	Homo sapi	MRPS14,M	-1.43176	1.45E-09	0.69844	11328.03	7931.428
A_23_P11	NM_01810	Homo sapi	PAPD1,mtI	-1.43189	0.00005	0.69838	1706.756	1179.018

A_23_P54	NM_01693	Homo sapi	UBN1,VT,\	-1.4319	3.62E-10	0.69837	2464.048	1721.856
A_32_P57702		Unknown	A_32_P57	-1.43199	1.61E-16	0.69833	6296.369	4396.232
A_23_P25	NM_00457	Homo sapi	PSPH,PSF	-1.43208	7.54E-11	0.69828	3049.736	2129.513
A_23_P13	NM_01789	hypothetica	KS,FLJ206	-1.43225	1.37E-07	0.6982	4892.666	3433.857
A_24_P13	AK022759	Ubiquitin sp	USP42	-1.43226	8.59E-11	0.6982	2125.933	1484.113
A_23_P26	NM_01601	Homo sapi	NDUFAF1,	-1.43275	1.76E-17	0.69796	7478.9	5220.017
A_23_P70	NM_01515	Homo sapi	PHF3,KIA	-1.43292	5.68E-07	0.69788	1498.639	1041.061
A_23_P83	NM_02307	hypothetica	FLJ13855,	-1.43299	0.00001	0.69784	9930.53	6985.313
A_23_P20	NM_00460	Homo sapi	TROVE2,S	-1.43319	0.00025	0.69775	1476.406	1044.075
A_23_P34	NM_01406	Homo sapi	MAGEH1,A	-1.43333	2.49E-07	0.69768	5084.46	3558.157
A_23_P69	NM_00166	ras homolo	RHOA,ARH	-1.43361	6.54E-15	0.69754	16041.93	11201.16
A_23_P12	NM_01866	Homo sapi	PXMP2,PM	-1.43372	5.71E-08	0.69749	5740.081	4003.217
A_24_P48	NM_03311	MADP-1 pr	MADP-1,M	-1.43389	0.00024	0.6974	1597.717	1101.767
A_23_P15	NM_01820	Homo sapi	CKAP2,LB	-1.43415	1.99E-08	0.69728	2685.856	1872.589
A_23_P83	NM_00712	Homo sapi	VCP,p97,T	-1.43422	2.96E-10	0.69724	42724.48	29840.91
A_23_P91	NM_01444	stress-assc	SERP1,RA	-1.43428	2.03E-10	0.69721	8218.766	5749.967
A_24_P48	NM_00543	v-yes-1 Ya	YES1,Yes,	-1.43439	9.26E-13	0.69716	3334.208	2324.968
A_23_P25	NM_00392	Homo sapi	HERC1,p5	-1.43439	0.00017	0.69716	861.6432	595.726
A_24_P32	NM_00650	melanoma	MCAM,CD	-1.43464	5.42E-08	0.69704	1626.676	1135.767
A_24_P40	BC030093	Myelin bas	MBP	-1.43474	0.00075	0.69699	1060.806	724.2213
A_23_P43	NM_03229	hypothetica	CAMKK1,C	-1.43498	4.54E-06	0.69687	1278.36	888.0247
A_23_P19	NM_01559	Homo sapi	PGM3,AGN	-1.43513	0.00023	0.6968	1691.44	1195.121
A_24_P32708		Unknown	A_24_P32	-1.43518	1.29E-09	0.69677	5360.68	3738.161
A_23_P35	NM_00249	Homo sapi	NEK2,NLK	-1.43539	0.00066	0.69667	1069.981	729.3013
A_23_P12	NM_01604	Exosome c	EXOSC3	-1.43542	5.86E-10	0.69666	7965.393	5565.223
A_24_P59	BC026736	RB-associa	RBAK	-1.43547	5.68E-07	0.69664	3489.722	2450.218
A_24_P18	NM_03270	tubulin alpl	TUBA6,MG	-1.43585	0.00001	0.69645	303484	213429.7
A_23_P23	NM_03230	Homo sapi	POLR3GL,	-1.43628	0.00003	0.69624	5799.831	4072.666
A_24_P88	NM_01650	putative S1	PS1D,pNO	-1.43631	2.01E-06	0.69623	3044.64	2114.161
A_23_P81	NM_00651	Homo sapi	TCTEL1,C	-1.43655	1.77E-18	0.69611	26975.28	18777.21
A_24_P3704		ALU7_HUM	THC18162	-1.43657	2.75E-07	0.6961	1311.461	916.7657
A_24_P39	NM_01908	hypothetica	FLJ10287,	-1.43661	0.00078	0.69609	718.0231	505.5195
A_23_P13	NM_00360	Homo sapi	STK6,AIK,A	-1.43716	1.29E-15	0.69582	4067.168	2829.886
A_23_P10	NM_00581	Homo sapi	PRKD3,EP	-1.43723	0.00031	0.69578	1152.955	788.8622
A_23_P12	NM_19903	Homo sapi	KLHL5	-1.4373	0.00061	0.69575	1060.723	744.9611
A_23_P124823		SPC4_HUM	THC19785	-1.43734	4.63E-14	0.69573	8238.35	5734.27
A_24_P332326		Unknown	A_24_P33	-1.43788	4.13E-08	0.69547	50085.38	34981.27
A_24_P41	NM_02454	hypothetica	LRRIQ2,FL	-1.43799	2.29E-06	0.69542	1337.043	934.1824
A_23_P93	NM_00670	Homo sapi	NUDT3,DIF	-1.43805	0.00004	0.69539	1114.563	781.7968
A_24_P17	NM_03131	heterogene	HNRPC,C1	-1.43816	3.86E-17	0.69533	9848.863	6849.072
A_23_P51	NM_00726	Homo sapi	STXBP3,P	-1.43835	1.69E-09	0.69524	2618.327	1820.982
A_24_P418517		Unknown	ENST0000	-1.43846	6.60E-14	0.69519	18012.36	12518.82
A_23_P12	NM_00138	Homo sapi	DPH2,DPH	-1.43863	2.32E-10	0.6951	3851.937	2674.913
A_23_P33	NM_02275	S100P bin	S100PBPR	-1.43882	8.30E-06	0.69501	2098.542	1447.225
A_23_P15	NM_00393	Homo sapi	SCAP2,PR	-1.43928	6.45E-07	0.69479	5148.833	3590.235
A_24_P25	NM_00031	phosphata	PTEN,BZS	-1.43941	0.00005	0.69473	1262.136	864.3829
A_24_P83	NM_00566	Wolf-Hirsch	WHSC2,NE	-1.43947	0.00002	0.6947	1004.302	693.2458
A_23_P21	NM_00017	Homo sapi	NR3C1,GR	-1.43969	0.00084	0.6946	3758.895	2598.822
A_23_P10	NM_00295	ribophorin I	RPN2,RIBI	-1.43974	1.28E-10	0.69457	2009.985	1395.791
A_23_P14	NM_01231	Homo sapi	KIF4A,KIF4	-1.44026	0.00001	0.69432	1924.582	1346.846
A_23_P44	NM_00637	Homo sapi	CRTAP,CA	-1.44033	0.00003	0.69428	1735.908	1197.162
A_24_P36	NM_01339	FtsJ homol	FTSJ2,FJH	-1.4404	0.00003	0.69425	954.2163	664.0384

A_23_P13	NM_13877	hypothetica	LOC91137	-1.44046	9.73E-10	0.69422	1786.323	1239.381
A_24_P383834		Unknown	A_24_P38	-1.44094	1.78E-08	0.69399	6390.66	4447.093
A_23_P21	NM_01401	Homo sapi	SACM1L,S	-1.441	0.00032	0.69396	944.6233	669.9191
A_24_P18	NM_01598	ubiquitin C	UCHL5,UC	-1.44117	1.49E-08	0.69388	4563.429	3176.922
A_23_P95	NM_14508	Homo sapi	NSMCE1,N	-1.44122	4.43E-14	0.69386	3325.323	2307.359
A_23_P15	NM_00436	Homo sapi	CDH7,CDH	-1.4413	6.75E-06	0.69382	1353.292	932.4581
A_24_P91	AK026497	Solute carr	SLC25A29	-1.44145	1.53E-15	0.69375	8214.17	5696.37
A_23_P150827		JE0272 lov	THC18357	-1.44148	5.34E-06	0.69373	1312.004	917.2471
A_23_P16	NM_00657	Homo sapi	RRAGA,RA	-1.44154	6.61E-14	0.6937	7267.884	5044.078
A_23_P10	NM_01631	Homo sapi	WBP11,NF	-1.44155	2.88E-12	0.6937	2251.016	1560.834
A_24_P90	D28589	KIAA0114	KIAA0114	-1.44159	1.50E-08	0.69368	3398.431	2353.409
A_23_P14	NM_01793	Homo sapi	PHIP,ndrp,	-1.44175	2.58E-07	0.6936	5132.056	3555.997
A_23_P11	NM_01742	Homo sapi	COQ3,bA9	-1.44206	9.37E-06	0.69345	1227.639	858.3079
A_24_P20	NM_02274	hypothetica	FLJ22558	-1.44208	0.00008	0.69345	2591.465	1795.987
A_24_P79	NM_01430	SEC11-like	SEC11L1	-1.44231	2.77E-13	0.69333	35378.82	24463.05
A_23_P15	NM_01621	Homo sapi	TRIP4,HsT	-1.44255	1.66E-09	0.69322	2325.683	1610.849
A_23_P54	NM_02182	Homo sapi	NIF3L1,CA	-1.44256	2.12E-15	0.69321	4878.655	3381.697
A_32_P54	BX537645	Hypothetic	LOC40139	-1.44266	1.17E-14	0.69316	9462.594	6563.663
A_24_P27	NM_01902	Homo sapi	TMCO1,PC	-1.44268	4.35E-12	0.69316	9782.165	6769.049
A_23_P89	NM_00365	Homo sapi	COPS3,SG	-1.44291	5.89E-07	0.69304	15467.33	10806.97
A_23_P26	NM_02475	modulator	FLJ13213,	-1.44293	1.37E-11	0.69304	2560.566	1775.463
A_23_P32	NM_00013	Homo sapi	FANCC,FA	-1.44299	9.12E-08	0.69301	1565.707	1092.914
A_23_P12	NM_03250	Homo sapi	ACSS1,AC	-1.4432	0.00056	0.69291	1656.662	1170.604
A_32_P32	NM_01567	neurobeacl	NBEA,BCL	-1.44331	0.00053	0.69285	765.9598	537.5408
A_32_P19608		AF247786	THC18468	-1.44339	0.00009	0.69282	898.9527	616.4388
A_23_P11	NM_00621	Homo sapi	PHYH,PAH	-1.44343	0.00015	0.69279	1063.354	726.923
A_23_P15	NM_00461	topoisomer	TOP3A,TO	-1.44356	0.0007	0.69273	758.0933	533.6999
A_23_P25	NM_03120	Homo sapi	LAS1L,FLJ	-1.44362	1.80E-06	0.6927	24653.28	17136.46
A_23_P37	NM_01810	Homo sapi	CDCA8,BC	-1.44369	1.18E-14	0.69267	8857.184	6137.445
A_24_P33	NM_19848	Homo sapi	RPL7L1,M	-1.4439	1.31E-10	0.69257	2088.933	1447.77
A_24_P18	NM_18250	hypothetica	MTERFD2,	-1.44408	4.59E-07	0.69248	1932.602	1333.146
A_23_P10	NM_01535	Homo sapi	SUZ12,CH	-1.44416	0.00002	0.69244	6742.717	4691.533
A_24_P40	NM_00247	myosin, he	MYH9,MHA	-1.44436	1.35E-15	0.69235	3661.181	2533.677
A_24_P126851		Unknown	A_24_P12	-1.44438	1.13E-06	0.69234	1392.965	959.6832
A_24_P234871		Unknown	A_24_P23	-1.44441	4.18E-07	0.69232	2144.327	1477.996
A_32_P16	NM_00337	voltage-de	VDAC1,PC	-1.44449	0.00003	0.69228	2021.055	1415.462
A_24_P60	AL832758	SNRPN up	SNRPN	-1.44485	2.58E-07	0.69212	1889.247	1301.149
A_23_P34	NM_00286	RAB2, mer	RAB2,RAB	-1.44486	2.49E-06	0.69211	3104.518	2144.206
A_23_P39	NM_02121	Homo sapi	C9orf80,HS	-1.44497	2.58E-09	0.69206	1950.255	1351.196
A_23_P10	NM_01765	aftiphilin pr	AFTIPHILII	-1.445	0.00007	0.69204	990.5335	680.3994
A_23_P10	NM_00504	PWP2 (per	PWP2H,PV	-1.44512	2.84E-08	0.69198	1329.352	921.282
A_24_P25	NM_03283	hypothetica	CIRH1A,NA	-1.44512	0.00008	0.69199	3143.071	2188.898
A_23_P58	NM_00034	survival of	SMN1,SMN	-1.44526	2.22E-13	0.69192	36701.27	25396.23
A_23_P16	NM_17686	Homo sapi	PSME3,Ki,	-1.44539	1.25E-08	0.69186	2473.685	1714.272
A_24_P46	NM_00487	etoposide-i	EI24,PIG8,	-1.44553	4.47E-07	0.69179	3714.01	2570.226
A_32_P86	BX537972	Hypothetic	LOC11606	-1.44579	0.00016	0.69166	1721.998	1210.129
A_32_P13	NM_02507	Ngg1 intera	NIF3L1BP1	-1.44582	2.58E-08	0.69165	14000.38	9738.299
A_23_P42	NM_03244	ligand-dep	MLR2,LCC	-1.44588	0.00027	0.69162	1087.355	738.7683
A_23_P11	NM_02442	Homo sapi	PLA2G4A,	-1.44595	1.31E-12	0.69158	3466.932	2402.059
A_23_P35	NM_02103	Homo sapi	MBNL1,EX	-1.44599	1.00E-13	0.69157	9540.947	6609.341
A_23_P32	NM_03190	SVH protei	SVH,MGC3	-1.44606	1.22E-07	0.69153	4672.714	3240.832
A_24_P281853		Unknown	A_24_P28	-1.44606	6.27E-08	0.69154	2736.551	1891.064

A_23_P30	NM_02245	Homo sapi	LMBR1,AC	-1.44612	2.98E-06	0.69151	1079.09	748.7301
A_32_P28	NM_00675	transaldola	TALDO1,T	-1.44616	0.00004	0.69149	51360.55	35816.73
A_23_P87810		Y15491 fer	THC18913	-1.44622	1.94E-07	0.69146	1204.219	839.2523
A_24_P32	NM_15371	Homo sapi	NUP62,p62	-1.4463	0.00004	0.69142	15147.54	10521.79
A_23_P41	NM_14526	similar to R	LOC13395	-1.44635	2.24E-08	0.69139	2085.998	1445.172
A_23_P20	NM_00165	ADP-ribosy	ARF1	-1.44645	0.00087	0.69135	685.3796	478.759
A_23_P50	NM_01421	Homo sapi	IMPA2	-1.44647	5.50E-12	0.69134	22909.96	15866.7
A_24_P16	NM_00571	actin relate	ARPC3,AR	-1.44653	8.26E-12	0.69131	58277.28	40618.72
A_24_P38	NM_00112	adaptor-rel	AP1G1,AD	-1.44665	6.30E-10	0.69125	4402.466	3041.556
A_23_P25	NM_00680	Homo sapi	HNRPA0,h	-1.4467	5.15E-07	0.69123	7327.145	5042.082
A_23_P34	NM_03142	Homo sapi	MRPL9,L9	-1.44673	2.88E-09	0.69121	26383.57	18321.41
A_23_P38	NM_01812	elaC (E. co	ELAC2,EL	-1.44691	2.62E-07	0.69113	30894.89	21499.32
A_23_P39	NM_00307	Homo sapi	SMARCA4	-1.44694	2.80E-11	0.69111	15173.56	10525.46
A_24_P39	NM_00166	ADP-ribosy	ARF5	-1.4472	0.00093	0.69099	1850.004	1290.056
A_23_P30	NM_13913	Homo sapi	NUP98,AD	-1.44732	6.34E-08	0.69093	1544.151	1069.698
A_23_P96	NM_00213	Homo sapi	RBMX,RNI	-1.44732	0.00001	0.69093	11880.96	8173.646
A_23_P96	NM_00614	Homo sapi	DNCLI2,LI	-1.4474	1.06E-17	0.69089	6755.763	4667.093
A_23_P11	NM_03250	Homo sapi	RBM13	-1.44745	1.04E-07	0.69087	2983.085	2056.465
A_23_P19	NM_03126	Homo sapi	HNRPAB,A	-1.44757	0.00003	0.69081	2994.47	2063.499
A_24_P12	NM_00280	proteasom	PSMD1,S1	-1.44787	3.02E-10	0.69067	2917.634	2014.305
A_23_P16	NM_18275	Homo sapi	MCM10,CN	-1.4481	1.33E-08	0.69056	2058.004	1416.007
A_32_P14201		Unknown	ENST0000	-1.44822	0.00078	0.6905	736.7374	516.7466
A_23_P16	NM_00548	Homo sapi	PARP2,AD	-1.44826	4.59E-07	0.69048	3583.918	2485.385
A_23_P14	NM_00455	Homo sapi	NDUFS2	-1.44861	8.59E-06	0.69032	2077.191	1439.654
A_32_P18	NM_01326	bromodom	BRD7,BP7	-1.44874	6.93E-10	0.69026	10836.51	7445.515
A_23_P21	NM_03330	Homo sapi	VPS13A,C	-1.44875	0.00003	0.69025	941.0205	646.3325
A_24_P18	NM_02037	chromoson	C12orf4	-1.44899	0.00086	0.69014	646.6322	447.759
A_24_P93	AK057835	CDNA FLJ	AK057835	-1.44923	0.00075	0.69002	886.2788	624.7828
A_23_P11	NM_01601	Homo sapi	LCMT1,LC	-1.44923	1.45E-14	0.69002	3583.789	2473.421
A_24_P12	NM_00466	RAB11A, n	RAB11A,Y	-1.44954	1.27E-06	0.68987	4240.467	2926.69
A_23_P16	NM_00121	Homo sapi	BUB1B,SS	-1.45003	5.86E-08	0.68964	2526.181	1751.019
A_24_P21	AB002377	Ankyrin rep	ANKRD28	-1.4502	0.00014	0.68956	945.7105	643.1738
A_24_P410117		Unknown	ENST0000	-1.45043	6.49E-17	0.68945	14603.86	10063.58
A_23_P11	NM_00640	Homo sapi	PRDX4,AC	-1.45054	6.60E-10	0.6894	58403.56	40097.33
A_24_P682601		Unknown	A_24_P68	-1.45069	5.56E-09	0.68933	1954.382	1349.435
A_23_P72	NM_02019	Homo sapi	MRPS22,G	-1.45087	1.06E-10	0.68924	11211.69	7734.713
A_23_P67	NM_00403	Homo sapi	ADCY3,AC	-1.4509	9.40E-10	0.68923	7963.789	5491.837
A_23_P25	NM_18231	Homo sapi	COVA1,AF	-1.45091	3.02E-06	0.68922	1448.981	1004.242
A_23_P10	NM_01900	hypothetica	FLJ20232,	-1.4512	0.00014	0.68909	1357.47	954.0642
A_24_P141804		Unknown	THC18908	-1.45154	4.37E-08	0.68893	3513.792	2424.285
A_23_P29	NM_05296	Homo sapi	RPL39L,RI	-1.4516	1.06E-14	0.6889	3261.979	2247.267
A_23_P36	NM_00237	Homo sapi	MAN2A1,M	-1.45162	0.00009	0.68889	1355.617	918.8569
A_32_P23	NM_01243	sec22 hom	SEC22L2,S	-1.45176	0.00002	0.68882	1427.37	974.1093
A_24_P98	NM_03312	Homo sapi	ANKRD13,	-1.45181	0.00005	0.68879	1041.334	708.6732
A_23_P94	NM_00349	Homo sapi	TRRAP,TR	-1.45284	1.72E-06	0.6883	1972.564	1351.344
A_23_P64	BC014125	Splicing fac	SF3B2	-1.45286	5.30E-16	0.6883	4848.522	3338.442
A_23_P69	NM_02277	Homo sapi	OSBPL11,	-1.45287	9.05E-15	0.68829	2994.662	2061.381
A_23_P30	NM_15242	hypothetica	FLJ39827,	-1.45297	0.00021	0.68825	862.3499	585.4136
A_23_P21	NM_02067	Homo sapi	ABHD6	-1.453	7.52E-08	0.68823	2851.624	1966.771
A_23_P10	NM_01607	Homo sapi	MRPS23,C	-1.45308	8.64E-12	0.6882	5172.043	3569.462
A_23_P69	NM_00166	Homo sapi	RHOA,ARI	-1.45309	1.38E-09	0.68819	33528.11	23157.45
A_32_P10	NM_00153	heterogene	HNRPL,hn	-1.45324	0.00062	0.68812	6610.656	4572

A_23_P82394		Unknown	THC18444	-1.45339	0.00056	0.68804	663.8344	454.3796
A_23_P38	NM_01430	SEC11-like	SEC11L1	-1.45353	6.50E-16	0.68798	44617.13	30738.04
A_32_P182135		Unknown	A_32_P18	-1.45364	0.00047	0.68793	733.1056	498.1935
A_23_P16	NM_02407	Homo sapi	DDX54,DP	-1.45376	7.79E-11	0.68787	4242.93	2926.564
A_24_P384196		Unknown	A_24_P38	-1.45393	0.00012	0.68779	834.37	582.3275
A_23_P29	NM_14564	Homo sapi	APOL3,CG	-1.45395	0.00068	0.68778	754.0746	526.5349
A_24_P47	AK057359	Hypothetic	LOC14634	-1.45408	3.28E-09	0.68772	1795.157	1239.834
A_23_P43	NM_00273	Homo sapi	PRKAR1A,	-1.45409	8.68E-08	0.68772	3123.729	2151.256
A_24_P28	AK000364	Chromodo	CHD7	-1.45443	2.55E-07	0.68756	1249.392	857.8839
A_23_P72	NM_00381	Homo sapi	ADAM9,MC	-1.4545	0.00004	0.68752	1588.886	1082.931
A_32_P60	NM_03232	Inhibitor of	ING5	-1.45474	1.02E-06	0.68741	1912.959	1310.228
A_23_P31	NM_00591	malate deh	MDH2,MDI	-1.45507	2.72E-10	0.68725	16996.45	11735.6
A_24_P26	NM_13843	hypothetica	C8orf40	-1.45535	0.00013	0.68712	854.2859	591.0579
A_32_P151648		Unknown	A_32_P15	-1.45549	0.00032	0.68705	2551.057	1766.908
A_23_P40	NM_13840	Homo sapi	C6orf51,bA	-1.45561	4.45E-08	0.687	27504.82	18961.16
A_23_P11	NM_03194	IFP38	IFP38	-1.45567	1.47E-15	0.68697	3443.116	2365.327
A_23_P18	NM_02481	Homo sapi	UBE1DC1,	-1.45585	0.00013	0.68689	1662.072	1127.35
A_23_P10	AK057777	Hypothetic	MGC4767	-1.45596	1.15E-10	0.68683	4512.336	3102.307
A_23_P98	NM_00578	Homo sapi	HRMT1L3,	-1.45597	6.88E-09	0.68683	1408.382	967.8732
A_24_P59	NM_02198	ubiquitin-c	UBE2V1,C	-1.4561	0.00016	0.68677	2134.245	1451.688
A_23_P35	NM_01812	hypothetica	C10orf6	-1.4561	0.0003	0.68677	688.7832	474.6625
A_24_P549927		Unknown	A_24_P54	-1.45621	0.00053	0.68672	928.6869	625.7509
A_24_P11	NM_00226	karyopheri	KPNA4,QIF	-1.45626	1.92E-07	0.68669	6630.498	4570.468
A_24_P80	NM_01436	protein tyr	PTPN18,BI	-1.45642	0.00013	0.68662	1271.76	860.138
A_24_P255914		Unknown	A_24_P25	-1.45668	6.07E-08	0.68649	1368.771	941.4404
A_23_P252125		TRI6_HUM	THC18150	-1.45675	0.00076	0.68646	13578.73	9420.322
A_23_P39	NM_00510	Homo sapi	FEZ2,HUM	-1.45676	7.65E-11	0.68646	3654.471	2508
A_23_P10	NM_01401	Homo sapi	HSPC009	-1.45698	1.61E-17	0.68635	16405.69	11266.81
A_24_P25	NM_02121	hypothetica	C20orf77,d	-1.45722	2.73E-07	0.68624	2762.478	1903.729
A_24_P35	AK129956	Nuclear fac	NFIC	-1.45731	0.00012	0.6862	5746.946	3960.478
A_32_P62	BC039145	Hypothetic	LOC28346	-1.4574	5.50E-09	0.68615	1486.255	1021.202
A_23_P91	NM_00375	eukaryotic	EIF3S7,MC	-1.45741	1.13E-06	0.68615	58007.56	40081.62
A_23_P33	NM_00660	Homo sapi	STAG2,SA	-1.45764	1.04E-07	0.68604	2777.125	1900.094
A_23_P34	NM_14506	Homo sapi	C13orf3,RA	-1.45769	9.96E-11	0.68602	3560.39	2442.13
A_23_P41	NM_03066	Homo sapi	RAI1,SMS,	-1.45775	1.05E-06	0.68599	2984.894	2048.231
A_23_P14	NM_08079	Homo sapi	MRPL39,L	-1.45785	5.56E-14	0.68594	7227.229	4963
A_23_P11	NM_01473	Homo sapi	KIAA0101,	-1.45799	5.14E-09	0.68588	68870.4	47443.37
A_23_P39	NM_14531	Homo sapi	RASGEF1A	-1.45809	0.00004	0.68583	1260.7	855.1485
A_23_P10	NM_00142	Homo sapi	EMP2,XMF	-1.4581	1.22E-13	0.68583	7808.317	5357.547
A_24_P35	NM_14463	hypothetica	MGC21688	-1.45813	2.29E-12	0.68581	2879.412	1974.288
A_23_P96	NM_00498	melanoma	MAGEA1,M	-1.45819	1.61E-12	0.68578	12935	8888.948
A_23_P20	NM_01483	chromoson	C1orf16,ES	-1.45862	3.59E-11	0.68558	1817.07	1246.632
A_23_P52	NM_03236	Homo sapi	ACBD6,MC	-1.45877	3.30E-09	0.68551	9315.969	6416.268
A_23_P15	NM_00314	Homo sapi	SSB,LARP	-1.45902	1.24E-09	0.68539	43918.54	30241.44
A_23_P43	NM_00115	Homo sapi	SLC25A5,T	-1.45926	8.46E-13	0.68528	33302.61	22744.25
A_23_P43	NM_03236	Homo sapi	DNAJC14,I	-1.45935	0.00009	0.68524	1213.727	818.9315
A_23_P14	NM_00721	Homo sapi	RNF139,R	-1.45938	8.20E-12	0.68522	2227.936	1527.018
A_24_P31	NM_01449	nuclear pro	ZNF638,ZF	-1.45941	0.00083	0.68521	620.8178	420.7239
A_23_P16	AK000318	Leucine zip	MTB	-1.45942	3.33E-07	0.6852	3273.374	2244.477
A_23_P58	NM_03091	Homo sapi	FIP1L1,Rh	-1.45942	0.00003	0.68521	7641.619	5249.963
A_23_P99	L40392	RNA bindir	RBM25	-1.4597	5.17E-09	0.68507	2555.089	1755.393
A_23_P25	NM_01905	Homo sapi	MRPL50,M	-1.45979	1.04E-07	0.68503	8511.078	5847.343

A_23_P41	NM_00132	Homo sapi	CTBP1,MG	-1.4598	5.64E-13	0.68503	9138.861	6263.247
A_24_P56	NM_01657	GMPR2 for	GMPR2,M	-1.46002	0.00003	0.68492	9632.584	6626.211
A_23_P69	NM_01614	Homo sapi	DNCLI1	-1.46021	1.61E-06	0.68483	1210.96	825.2637
A_23_P14	NM_01663	Homo sapi	SPG21,MA	-1.46043	1.27E-17	0.68473	14254.99	9761.129
A_23_P35	NM_00141	Homo sapi	EIF1AX,EII	-1.46066	4.78E-09	0.68462	26314.48	18117.48
A_24_P67	NM_00036	thiopurine S	TPMT	-1.46077	0.00005	0.68457	868.9951	598.7228
A_24_P26	NM_00011	dystonia 1,	TOR1A,DC	-1.4608	2.16E-06	0.68456	1460.107	993.6431
A_24_P35	NM_00321	testis enha	TEGT,BI-1	-1.46087	0.00001	0.68452	31256.07	21386.58
A_24_P18	NM_01824	hypothetica	TMEM30A	-1.46093	0.00007	0.6845	877.5674	604.9168
A_23_P13	NM_02515	hypothetica	WDR71,PA	-1.46097	0.0004	0.68448	628.8364	432.8479
A_23_P13	NM_00178	Homo sapi	CD63,MLA	-1.46111	1.09E-07	0.68441	91492.08	62282.51
A_23_P90	NM_02291	Homo sapi	MRPL44,F	-1.46125	4.88E-08	0.68435	3104.223	2120.721
A_24_P57	NM_01849	COBW-like	CBWD1	-1.46146	0.00005	0.68425	1062.613	718.97
A_32_P50	NM_03115	heterogene	HNRPA1,H	-1.4615	1.04E-12	0.68423	59654.63	40726.54
A_23_P85	NM_00703	RER1 reter	RER1	-1.46153	1.39E-14	0.68422	5950.404	4069.283
A_24_P11	NM_00670	similar to S	TXNL4A,D	-1.46176	3.08E-10	0.68411	7365.34	5058.829
A_23_P34	NM_00311	Homo sapi	SP3,SPR-2	-1.46176	1.41E-09	0.68411	3593.286	2452.883
A_23_P25	NM_00295	Homo sapi	RSN,CLIP,	-1.46178	2.32E-10	0.6841	4480.11	3064.486
A_23_P20	NM_00719	Homo sapi	SEC23IP,F	-1.46191	3.08E-07	0.68404	5222.674	3583.356
A_24_P25	NM_02183	chromoson	PEO1,PEC	-1.46201	0.00017	0.68399	7584.927	5200.551
A_24_P57	NM_00068	S-adenosy	AHCY,SAH	-1.46202	0.00033	0.68399	1013.136	708.2072
A_23_P13	NM_00670	Homo sapi	TXNL4A,D	-1.46234	4.92E-15	0.68384	25524.55	17423.54
A_24_P195454		Unknown	A_24_P19	-1.46239	3.53E-07	0.68381	1226.406	837.3552
A_24_P14	NM_01474	KIAA0255	TM9SF4,K	-1.46243	4.73E-09	0.68379	2839.771	1940.245
A_23_P35	NM_14586	Homo sapi	ANXA11,A	-1.46256	2.33E-16	0.68373	9850.236	6728.289
A_32_P18	NM_13877	Homo sapi	MRRF,RRF	-1.46307	1.19E-09	0.6835	1800.373	1231.932
A_23_P13	NM_00533	Homo sapi	HDLBP,HB	-1.46314	0.00002	0.68346	2703.548	1845.131
A_24_P273014		Unknown	ENST0000	-1.46326	6.78E-07	0.6834	1872.06	1284.273
A_32_P76	NM_01609	HSPC025	EIF3S6IP,E	-1.46328	8.90E-11	0.6834	5445.969	3710.76
A_23_P70	NM_00630	JTV1 gene	JTV1,P38,I	-1.46337	8.07E-18	0.68335	11040.03	7539.23
A_24_P36	NM_00454	NADH deh	NDUFC2,B	-1.46359	0.00022	0.68325	691.3829	473.8537
A_23_P50	NM_03184	heterogene	HNRPU,SA	-1.4638	2.56E-14	0.68315	15396.44	10538.7
A_24_P34	NM_17202	Homo sapi	POM121,M	-1.46386	3.99E-10	0.68312	1960.472	1340.448
A_23_P25	AB018256	Homo sapi	KIAA0713	-1.46402	0.00025	0.68305	1073.92	721.3486
A_24_P31	NM_00613	capping pr	CAPZA1,C	-1.4642	3.87E-19	0.68297	15483.28	10571.53
A_23_P72	NM_00560	Homo sapi	PTK2,FAK	-1.46426	7.66E-06	0.68294	1060.385	734.7512
A_23_P14	NM_00553	Homo sapi	IDH3A	-1.46444	5.19E-08	0.68286	1657.646	1126.535
A_23_P26	AK026008	WD-repeat	HAN11	-1.46459	0.00024	0.68278	2986.313	2063.225
A_32_P11	NM_01593	CGI-07 pro	NMD3,CGI	-1.46467	7.23E-06	0.68275	5615.952	3854.049
A_24_P66	AK055641	Chromobo	CBX5	-1.46469	3.76E-17	0.68274	4713.712	3218.323
A_24_P39	NM_00437	COX11 (ye	COX11,CC	-1.46499	9.90E-06	0.6826	1863.875	1261.913
A_23_P37	NM_01823	hypothetica	FLJ10826	-1.46507	1.70E-07	0.68256	4517.465	3095.608
A_24_P41	AK075087	G protein-c	GPR126	-1.46526	0.00041	0.68247	1074.763	716.8101
A_23_P10	NM_00206	Homo sapi	GCLM,GLC	-1.46536	2.23E-08	0.68243	15945.92	10967.98
A_23_P63	NM_00193	Homo sapi	DR1,NC2,N	-1.46538	0.00024	0.68242	821.7714	553.4491
A_23_P20	NM_00164	Homo sapi	APLP2,APL	-1.46545	5.87E-06	0.68238	8520.732	5837.54
A_23_P32	NM_00239	Homo sapi	MEF2C	-1.4655	0.00008	0.68236	1209.893	812.9411
A_23_P12	NM_03098	hypothetica	ISG20L2,F	-1.46562	6.97E-13	0.68231	2323.997	1585.094
A_24_P36	NM_00135	DEAD/H (A	DHX9,LKP	-1.4657	1.80E-11	0.68227	2867.482	1956.604
A_24_P14	NM_00637	cartilage as	CRTAP,CA	-1.4657	8.31E-06	0.68227	1151.766	779.7809
A_23_P15	NM_08064	Homo sapi	APEX1,AP	-1.46573	2.69E-06	0.68226	6905.308	4709.454
A_23_P34	NM_00012	Homo sapi	EPHX1,ME	-1.46587	2.68E-06	0.68219	2519.524	1716.36

A_23_P53	NM_03257	dpy-30-like	LOC84661	-1.46614	3.50E-09	0.68206	14844.41	10139.83
A_23_P91	NM_01904	putative pr	NBLA0005	-1.46617	3.20E-06	0.68205	4286.05	2925.096
A_23_P21	NM_01826	Homo sapi	ASXL2,AS	-1.46619	6.03E-08	0.68204	3828.579	2608.57
A_23_P14	NM_01438	Homo sapi	ATP2C1,H	-1.46622	4.83E-15	0.68203	4729.975	3226.956
A_23_P91	NM_00476	Homo sapi	COPB2,be	-1.4663	4.36E-19	0.68199	12250.39	8354.951
A_23_P86	NM_00272	Homo sapi	PRG1,PPG	-1.46632	1.87E-14	0.68198	32172.06	21906.18
A_24_P37	NM_01822	chromoson	MOBK1B,M	-1.46646	4.02E-07	0.68191	1624.705	1103.219
A_24_P39	NM_00660	putative ho	PHTF1,PH	-1.46656	0.00015	0.68187	705.9982	488.4474
A_23_P35	NM_01902	Homo sapi	TMCO1,PC	-1.4668	3.56E-18	0.68176	13694.93	9330.687
A_24_P341489		Unknown	A_24_P34	-1.46703	0.00008	0.68165	896.6173	615.4457
A_24_P13	NM_00386	carboxyles	CES2,iCE,	-1.46711	1.36E-11	0.68161	9395.137	6420.388
A_23_P12	NM_00213	Homo sapi	HMOX1,HC	-1.46714	9.56E-08	0.6816	96553.52	65353.4
A_24_P13	NM_01438	ATPase, C	ATP2C1,H	-1.46715	4.65E-07	0.68159	1136.719	770.0495
A_23_P11	NM_00663	Homo sapi	RPP40,RN	-1.46744	3.27E-13	0.68146	4836.619	3297.787
A_23_P12	NM_00447	FSHD regi	FRG1,FSG	-1.46772	7.16E-07	0.68133	10216.27	7005.456
A_23_P11	AB007963	KIAA0494	KIAA0494	-1.46784	8.48E-08	0.68127	1644.035	1115.697
A_24_P28	NM_00538	protein-L-is	PCMT1	-1.46808	2.69E-13	0.68116	2673.218	1821.038
A_23_P16	NM_00541	Homo sapi	SHMT2,GL	-1.46822	2.24E-14	0.6811	24411.68	16648.04
A_23_P31	NM_17812	Homo sapi	CXorf40A,E	-1.46828	1.41E-18	0.68107	5988.94	4079.233
A_24_P26	NM_00135	2,4-dienoyl	DECR1,DE	-1.4684	3.32E-13	0.68101	2831.363	1928.265
A_23_P16	NM_02274	Homo sapi	HIPK2,PRC	-1.46854	2.10E-11	0.68095	3758.631	2560.694
A_23_P65	NM_01641	Homo sapi	C14orf87,C	-1.46858	1.85E-16	0.68093	20389.34	13891.94
A_24_P18	NM_03246	aspartate b	ASPH,BAH	-1.46859	2.19E-14	0.68092	7864.13	5362.822
A_23_P13	NM_00608	Homo sapi	IK,RED,CS	-1.46869	1.15E-17	0.68088	12972.49	8840.565
A_24_P13	NM_01438	ATPase, C	ATP2C1,H	-1.4687	0.00058	0.68087	1516.568	1010.176
A_24_P41	NM_00532	L-3-hydrox	HADHSC,S	-1.46871	6.81E-08	0.68087	2174.668	1484.669
A_23_P41	NM_00489	Homo sapi	SEC22L1,E	-1.46877	6.18E-09	0.68084	2955.693	2011.166
A_23_P21	NM_01216	Homo sapi	FBXL4,FBL	-1.46886	0.00018	0.6808	967.7125	649.0179
A_32_P24	NM_01534	Homo sapi	ZFYVE26,F	-1.46891	4.66E-07	0.68078	1425.828	976.2001
A_24_P17	NM_01504	KIAA0676	KIAA0676	-1.46915	0.00004	0.68066	4366.948	2976.918
A_23_P14	NM_03201	Chromosor	C9orf5	-1.4692	1.87E-11	0.68064	4699.366	3193.154
A_24_P98	NM_00398	microtubule	MAP7,EMA	-1.46942	1.25E-09	0.68054	1580.757	1072.704
A_32_P10	NM_01564	PAI-1 mRN	SERBP1,C	-1.46946	0.00005	0.68052	8006.641	5456.744
A_23_P30	NM_01869	Homo sapi	ERBB2IP,L	-1.46956	0.00002	0.68047	2373.938	1605.578
A_32_P22	NM_14569	Homo sapi	YWHAZ,KC	-1.46979	2.13E-07	0.68037	3903.515	2661.992
A_23_P11	NM_03149	peroxisoma	LONPL,LO	-1.46979	0.00001	0.68037	3882.977	2645.655
A_23_P93	NM_01910	homeo box	HOXA5,HC	-1.46992	0.00003	0.68031	949.4989	638.8071
A_32_P86009		Unknown	A_32_P86	-1.47019	1.51E-11	0.68019	2270.333	1545.158
A_24_P46	NM_03254	short coilec	SCOC,SCC	-1.47035	4.39E-06	0.68011	1648.267	1130.392
A_23_P21	AF111113	Kelch-like 7	KLHL7	-1.47059	3.66E-08	0.68	2656.557	1807.468
A_23_P16	NM_13901	signal pept	SPPL3,IMP	-1.47064	8.59E-06	0.67997	2939.716	1991.787
A_24_P25	NM_00165	ADP-ribosy	ARF3	-1.47085	3.28E-13	0.67988	5752.265	3913.468
A_24_P80	NM_00543	BENE prot	MALL,BEN	-1.47104	8.59E-08	0.67979	3646.638	2481.23
A_32_P65061		XPF_HUM	THC18473	-1.4711	0.00058	0.67977	830.4548	575.7543
A_23_P91	NM_01694	chromoson	C21orf6,GL	-1.47119	0.00006	0.67972	853.7765	585.8945
A_32_P21	AK023660	CDNA FLJ	AK023660	-1.47121	1.67E-07	0.67971	1151.993	784.5735
A_23_P54	NM_00375	Homo sapi	EIF3S1,eIF	-1.47132	3.24E-06	0.67966	19619.37	13454.99
A_23_P13	BC006795	Hypothetic	MGC5370	-1.47134	0.00024	0.67965	756.7056	517.8245
A_23_P76	NM_00610	Homo sapi	SKB1,JBP	-1.4716	1.14E-11	0.67953	8952.648	6065.751
A_23_P13	NM_01649	Homo sapi	MRPL37,M	-1.47164	1.45E-07	0.67951	8570.586	5855.338
A_24_P323916		Unknown	A_24_P32	-1.47175	0.00005	0.67946	1229.792	846.4998
A_23_P12	NM_00409	Homo sapi	DUSP3,VH	-1.47179	3.22E-09	0.67945	2204.321	1500.666

A_24_P25	NM_00633	C2f protein	C2F	-1.47197	9.81E-17	0.67936	7876.903	5345.565
A_23_P21	NM_13892	Homo sapi	SON,SON	-1.47202	7.46E-14	0.67934	5322.628	3616.035
A_23_P39	NM_00683	Homo sapi	IMMT,HMF	-1.47205	1.78E-07	0.67933	21724.71	14771.63
A_24_P73	NM_00411	flap structu	FEN1,MF1	-1.47209	6.94E-08	0.67931	1956.463	1325.001
A_23_P15	NM_01883	Homo sapi	SULF2,HS	-1.4721	6.48E-12	0.6793	2637.715	1790.737
A_23_P15	NM_00432	BCS1 (yea	BCS1L,BC	-1.4721	1.06E-14	0.6793	2796.176	1898.863
A_24_P23	NM_13839	Homo sapi	HNRPLL,S	-1.47221	1.58E-06	0.67925	1759.067	1185.656
A_24_P34	NM_15252	Homo sapi	SLC16A14	-1.47227	2.67E-07	0.67922	1217.269	829.0984
A_23_P20	NM_01833	Homo sapi	KIAA1598,	-1.47232	3.21E-08	0.6792	2223.715	1507.743
A_24_P30	NM_01433	CREBBP/E	CRI1,EID1	-1.47239	1.39E-14	0.67917	16325.35	11109.63
A_24_P27	NM_15282	Homo sapi	SNX3,SDP	-1.47264	9.69E-07	0.67905	4071.291	2761.971
A_32_P10	NM_00452	minichrom	MCM2,BM	-1.47274	3.23E-13	0.67901	7599.27	5165.993
A_23_P23	NM_00681	Homo sapi	MLLT11,AF	-1.47287	7.75E-08	0.67895	22029.36	14981.02
A_23_P20	NM_02245	Homo sapi	NOC3L,AD	-1.47294	1.72E-08	0.67891	3366.679	2298.286
A_23_P83	NM_00068	Homo sapi	ALDH1A1,	-1.47308	1.51E-09	0.67885	13727.29	9356.352
A_23_P40	AL049246	Hypothetic	FLJ10618	-1.47331	0.00039	0.67874	1321.365	878.2163
A_23_P65	NM_01430	Homo sapi	TGDS,TDF	-1.4735	0.00006	0.67866	2100.405	1409.786
A_32_P16	NM_14459	Homo sapi	TTC8,BBS	-1.4736	4.21E-07	0.67861	2401.486	1623.436
A_23_P48	NM_01398	Homo sapi	DIO2,SelY	-1.47367	7.59E-06	0.67858	3955.316	2710.229
A_23_P16	NM_02112	pyrophosp	PPA1,PP,F	-1.47384	4.04E-18	0.6785	31960.39	21702.7
A_23_P15	NM_01840	hypothetic	HSA27219	-1.47402	5.70E-10	0.67842	31087.04	21137.34
A_24_P37	NM_02524	hypothetic	TMEM22,M	-1.4741	5.49E-06	0.67838	1005.357	685.6564
A_32_P17	AY007094	IK cytokine	IK	-1.4741	1.63E-11	0.67838	12937.96	8808.377
A_23_P14	NM_00519	Homo sapi	CHES1,FC	-1.47417	3.14E-08	0.67835	3952.632	2685.886
A_23_P15	NM_01403	Homo sapi	BZW2,MST	-1.4742	1.21E-20	0.67833	25922.31	17582.57
A_24_P29	BC014203	Hypothetic	MGC45871	-1.47421	1.58E-08	0.67833	2792.175	1902.098
A_23_P42	NM_15282	Homo sapi	SNX3,SDP	-1.47435	7.99E-18	0.67826	12328.35	8367.996
A_23_P15	NM_00035	Homo sapi	TGFBI,CSI	-1.47438	1.75E-06	0.67825	9558.365	6524.678
A_23_P20	NM_02121	HCF-bindir	ZF	-1.47438	5.58E-11	0.67825	2228.47	1514.429
A_23_P44	NM_00201	Homo sapi	FLII,FLI,FL	-1.4746	5.95E-17	0.67815	4000.14	2712.335
A_23_P30	NM_03082	Homo sapi	PLA2G12A	-1.47463	1.67E-07	0.67814	2256.775	1526.432
A_23_P70	NM_01814	mitochondr	MRPS10,M	-1.47469	3.88E-12	0.67811	2456.895	1665.435
A_23_P15	AL117509	KIAA1666	KIAA1666	-1.47479	0.00001	0.67806	1000.534	684.5141
A_23_P11	NM_01609	DNA replic	Pfs2,HSPC	-1.47481	1.75E-11	0.67805	3328.872	2258.858
A_23_P21	NM_01819	Homo sapi	TMLHE,TM	-1.47486	1.72E-06	0.67803	999.6545	678.5496
A_23_P15	NM_14498	hypothetic	MGC23401	-1.47508	9.63E-08	0.67793	1242.012	844.8867
A_23_P12	NM_02475	hypothetic	PTCD2,FL	-1.47523	0.00015	0.67786	616.5838	416.6692
A_23_P42	NM_00239	Homo sapi	ME1,MES,	-1.47531	3.50E-18	0.67782	15794.75	10699.33
A_24_P324084		Unknown	A_24_P32	-1.4755	4.76E-13	0.67773	4928.88	3341.621
A_23_P16	NM_02064	Homo sapi	TMEM9B,C	-1.47556	9.00E-09	0.67771	13700.3	9261.199
A_23_P20	NM_02518	lin-10 prote	LIN10,lin-1	-1.47557	4.60E-13	0.6777	1960.054	1328.592
A_23_P51	NM_01908	hypothetic	FLJ10287,	-1.47563	0.00066	0.67767	762.6448	508.6171
A_23_P29	NM_03292	hypothetic	MGC13159	-1.4759	0.00024	0.67755	842.7692	579.9827
A_23_P14	NM_00063	Homo sapi	GSR,MGC	-1.47631	1.45E-10	0.67736	4840.816	3273.207
A_23_P45	BC017665	Chromosor	CXorf39	-1.47632	4.17E-06	0.67736	5314.54	3624.405
A_23_P13	NM_00583	Homo sapi	HRSP12,P	-1.47675	8.01E-08	0.67716	2991.76	2021.658
A_23_P12	NM_00367	Homo sapi	DEGS1,ML	-1.47694	8.34E-10	0.67708	4251.027	2877.441
A_24_P497843		Unknown	THC18702	-1.477	7.75E-07	0.67705	1637.52	1118.239
A_23_P50	NM_01819	Homo sapi	ZFP64,MG	-1.47702	2.25E-09	0.67704	2136.134	1442.813
A_24_P10	NM_14457	Homo sapi	C14orf32,M	-1.47703	2.98E-16	0.67703	5275.241	3574.02
A_23_P12	NM_02248	hypothetic	FLJ21657,	-1.47707	0.00003	0.67702	906.8597	610.4462
A_23_P20	NM_18323	Homo sapi	GSTO2,bA	-1.47708	1.32E-07	0.67701	2317.892	1573.419

A_23_P30	NM_18183	Homo sapi	UBE2D2,U	-1.47747	1.83E-13	0.67683	4664.823	3160.843
A_23_P34	NM_00361	Homo sapi	RAE1,MIG	-1.47753	1.70E-10	0.6768	2843.798	1930.286
A_32_P34	AK125119	Chromosor	C6orf68	-1.47759	1.11E-10	0.67678	1748.987	1182.647
A_23_P10	NM_02522	hypothetica	PRO2730,I	-1.4776	2.10E-06	0.67678	1865.59	1268.575
A_32_P42	BX538293	Homo sapi	BX538293	-1.47763	0.00002	0.67676	2076.222	1412.284
A_23_P20	NM_02098	Homo sapi	ANK3,ANK	-1.47765	1.75E-08	0.67675	3301.518	2237.364
A_23_P74	NM_01871	RCC1-like	RCC2,TD-(	-1.4778	2.42E-07	0.67668	2277.803	1558.423
A_32_P11	BC024301	Sorting nex	SNX12	-1.47783	1.05E-08	0.67667	9185.641	6226.12
A_23_P53	NM_00230	Homo sapi	LDHB,TRG	-1.47784	9.68E-21	0.67666	164581.7	111385.1
A_23_P71	NM_00446	Homo sapi	FDFT1,SS,	-1.47798	5.50E-19	0.6766	5488.918	3713.685
A_23_P94	NM_01479	Homo sapi	MELK,HPK	-1.47813	1.37E-12	0.67653	3021.284	2044.646
A_23_P11	NM_00016	GLI-Kruppe	GLI3,PHS,	-1.47819	3.79E-08	0.6765	1198.471	808.4513
A_23_P74	NM_00568	Homo sapi	TAF1A,SL1	-1.47837	0.00014	0.67642	1762.587	1178.427
A_23_P51	NM_02466	Hypothetic	FLJ11838	-1.47838	1.92E-07	0.67642	5871.115	3982.025
A_23_P86	S63912	Heterogen	HNRPA3P	-1.47858	3.44E-09	0.67632	2821.455	1915.773
A_23_P25	NM_01780	hypothetica	FLJ20397	-1.47869	1.92E-17	0.67627	3904.837	2640.435
A_23_P11	NM_18280	Arsenate re	ARS2	-1.4788	0.00001	0.67623	14931.46	10204.71
A_24_P21	NM_00235	mannose-6	M6PR,SMF	-1.47935	0.00011	0.67597	724.6904	488.6221
A_23_P20	NM_01602	DORA reve	DREV1,DF	-1.47967	0.00028	0.67583	913.6458	633.6616
A_32_P145241		Unknown	A_32_P14	-1.47969	1.12E-13	0.67582	4146.719	2802.498
A_23_P13	NM_01475	KIAA0097	CKAP5,TO	-1.47984	4.84E-11	0.67575	2058.265	1389.248
A_23_P14	NM_00637	Homo sapi	VTI1B,VTI1	-1.48002	2.12E-16	0.67566	13898.51	9380
A_23_P77	NM_00375	eukaryotic	EIF3S8,eIF	-1.48029	6.48E-06	0.67554	97705.67	66486.73
A_23_P11	NM_00196	eukaryotic	EIF4E,CBF	-1.48037	4.11E-06	0.67551	2083.505	1397.04
A_23_P84	NM_00178	Septin 7	7-Sep	-1.48045	2.90E-08	0.67547	3905.947	2643.942
A_23_P30	AK098569	Hypothetic	DKFZp667	-1.48062	1.59E-06	0.67539	1013.679	683.3993
A_23_P33	NM_03079	T-cell imm	CDA08,TIF	-1.48106	1.80E-16	0.67519	3362.676	2270.37
A_24_P12	NM_00270	phosphorib	PPAT,GPA	-1.48121	1.39E-08	0.67512	3168.339	2141.212
A_24_P409471		Unknown	A_24_P40	-1.48124	2.32E-11	0.67511	2331.634	1574.487
A_23_P99	NM_13847	hypothetica	MGC5987	-1.48125	2.67E-06	0.67511	3015.474	2037.011
A_23_P62	NM_01600	CGI-49 pro	CGI-49	-1.48135	8.71E-18	0.67506	12879.1	8700.234
A_23_P18	NM_01658	Chromosor	C3orf1	-1.48151	1.06E-11	0.67499	27372.68	18414.65
A_24_P91	AK025925	WD-repeat	HAN11	-1.48162	9.57E-10	0.67494	1909.737	1292.949
A_24_P928765		Unknown	A_24_P92	-1.48168	1.58E-08	0.67491	1675.474	1135.129
A_23_P25	NM_00202	Homo sapi	FNTB,FPT	-1.4819	0.0002	0.67481	1018.687	702.2581
A_23_P15	NM_00578	Homo sapi	TXNDC9,A	-1.48208	2.42E-08	0.67473	6376.416	4304.247
A_24_P19	NM_00568	ATP-bindin	ABCB6,AB	-1.48222	4.69E-06	0.67466	2294.617	1556.016
A_23_P13	NM_02452	hypothetica	FLJ22555,I	-1.48228	5.86E-11	0.67464	3413.545	2302.564
A_23_P63	NM_03299	Homo sapi	ZWINT,KN	-1.4823	4.68E-12	0.67463	10192.86	6892.796
A_23_P25	NM_00281	Homo sapi	PSMD10,p	-1.48255	1.55E-12	0.67452	7672.561	5168.361
A_32_P13	NM_01564	PAI-1 mRN	SERBP1,C	-1.4826	2.91E-16	0.67449	51386.7	34704.42
A_23_P43	NM_17886	source of ir	SIMP,FLJ9	-1.48269	0.00004	0.67445	807.6496	542.7827
A_24_P38	NM_00654	thioredoxin	TXNL2,PIC	-1.48283	1.88E-17	0.67439	17200.44	11592.24
A_24_P47211		AY211929	THC19758	-1.48325	0.00032	0.67419	1051.027	691.7783
A_24_P37	NM_02118	hypothetica	RAP2C	-1.48331	0.00003	0.67417	699.3831	474.8778
A_32_P22	BC065723	SUMO1 ps	LOC39125	-1.4834	3.54E-08	0.67413	2221.733	1491.689
A_23_P64	NM_02193	likely ortho	RIC8A,RIC	-1.48352	1.44E-12	0.67407	16091.2	10872.55
A_24_P87	AK055112	CDNA FLJ	AK055112	-1.48363	6.69E-06	0.67402	978.2178	663.0475
A_32_P20	NM_01414	HSPC274	C20orf30,H	-1.48369	5.88E-11	0.674	6908.885	4654.981
A_23_P17	NM_00068	Homo sapi	AHCY,SAH	-1.4837	2.86E-14	0.67399	45381.89	30655.73
A_23_P13	NM_00333	ubiquitin-a	UBE1,A1S	-1.48377	7.93E-09	0.67396	1923.326	1299.375
A_24_P20	NM_14569	Homo sapi	YWHAZ,K(	-1.48377	1.32E-06	0.67396	5454.567	3673.475

A_23_P63	NM_01418	Homo sapi	SSU72,HS	-1.48407	2.40E-20	0.67382	9667.555	6514.985
A_23_P21	NM_00534	Homo sapi	HMGB3,HM	-1.48433	1.70E-20	0.6737	12187.64	8209.73
A_23_P12	NM_01801	Homo sapi	ESRRBL1,	-1.4844	3.03E-07	0.67367	4658.607	3148.55
A_23_P14	NM_00235	Homo sapi	LYN,JTK8	-1.48444	2.91E-08	0.67366	3102.518	2086.814
A_23_P53	NM_01605	CGI-116 pr	CGI-116	-1.48471	1.86E-14	0.67353	2663.291	1793.924
A_24_P779258		AB040450	THC19434	-1.48479	2.22E-14	0.6735	2786.121	1876.322
A_24_P13	NM_15232	Homo sapi	PPIL5,LRR	-1.4848	3.33E-12	0.67349	2864.657	1929.766
A_23_P20	NM_01542	Homo sapi	AASDHPP	-1.48496	2.66E-09	0.67342	4458.688	2998.202
A_23_P40	NM_18269	Homo sapi	SRPK2,SF	-1.48498	0.00033	0.67341	707.6547	472.5504
A_32_P22	BG283145	Hypothetic	LOC28602	-1.48503	0.00033	0.67339	958.8457	661.2997
A_23_P46	NM_00262	Homo sapi	PFKP,PFK	-1.48517	4.70E-18	0.67332	7698.358	5183.006
A_23_P42	NM_02405	Homo sapi	C7orf24,M	-1.48523	1.28E-17	0.6733	19310.35	12986.32
A_23_P21	NM_00017	Homo sapi	GSS,GSHS	-1.48557	9.60E-07	0.67314	4102.902	2751.09
A_24_P32	NM_00079	dihydrofola	DHFR	-1.48616	7.02E-12	0.67287	2173.763	1463.081
A_23_P99	NM_03249	PNAS-127	C14orf142,	-1.48623	5.43E-07	0.67284	1048.361	704.8131
A_23_P26	NM_01564	RAP1B, m	RAP1B,K-F	-1.48645	4.77E-21	0.67274	25294.37	17019.77
A_23_P21	AK094187	Kruppel-lik	LOC37706	-1.48661	0.00009	0.67267	832.6227	566.365
A_32_P12	AK002014	Chromosor	C6orf70	-1.48681	2.18E-07	0.67258	1407.812	952.5303
A_24_P89	NM_00210	histidyl-tRN	HARS,HRS	-1.48735	1.66E-10	0.67233	3677.117	2471.176
A_24_P13	AK026078	ELK3, ETS	ELK3	-1.48737	6.39E-12	0.67233	3505.227	2357.318
A_23_P97	NM_01809	adaptin-ea	NECAP2,F	-1.48748	5.76E-12	0.67228	3021.59	2030.555
A_24_P22	AB028987	Chromosor	C19orf7	-1.48781	2.92E-13	0.67213	2965.535	1995.2
A_23_P87	NM_00633	C2f protein	C2F	-1.48795	8.23E-17	0.67207	26630.15	17922.22
A_23_P50	NM_02450	Homo sapi	LRFN3,MG	-1.488	7.78E-08	0.67204	1625.404	1087.2
A_24_P80	NM_01784	hypothetica	FLJ20507,	-1.48831	3.34E-10	0.6719	2205.93	1479.041
A_24_P20	NM_13337	splicing fac	YT521,KIA	-1.4885	0.0004	0.67182	1618.776	1066.957
A_32_P155512		Unknown	A_32_P15	-1.48856	1.97E-10	0.67179	1657.357	1114.095
A_23_P64	NM_00641	Homo sapi	HTATIP2,C	-1.48893	7.76E-12	0.67162	14038.52	9461.908
A_23_P99	NM_00683	Homo sapi	METAP2,p	-1.48945	1.00E-09	0.67139	29955.29	20180.38
A_24_P22	NM_01232	U6 snRNA	LSM4,YER	-1.4895	5.51E-12	0.67137	2731.744	1834.128
A_23_P21	NM_00465	Homo sapi	USP9X,DF	-1.48968	1.22E-14	0.67129	13236.6	8899.134
A_24_P42	NM_00153	heterogene	HNRPL,hn	-1.48986	1.30E-06	0.6712	3404.84	2283.641
A_32_P13	BC015814	KIAA1799	KIAA1799	-1.48989	3.97E-06	0.67119	1161.214	773.4875
A_32_P22	NM_00156	eukaryotic	EIF3S6,IN	-1.48992	0.00002	0.67118	1375.833	934.3485
A_23_P76	NM_14505	hypothetica	MGC7036,	-1.48997	1.89E-11	0.67115	4223.531	2837.587
A_32_P149416		Unknown	A_32_P14	-1.49024	2.66E-06	0.67103	2114.512	1408.5
A_23_P98	NM_00490	Homo sapi	PRDX6,PR	-1.4903	3.24E-06	0.671	112711.9	76344.73
A_23_P40	NM_15335	Homo sapi	FBXL16,Fb	-1.49036	0.00013	0.67098	1036.529	681.9294
A_32_P73	NM_01868	CMP-N-ac	CMAS	-1.49049	0.00001	0.67092	1351.761	897.2194
A_23_P21	NM_03313	Homo sapi	CNP,CNP1	-1.49056	7.51E-06	0.67089	18586.94	12582.1
A_23_P13	NM_02183	Homo sapi	PEO1,PEC	-1.49061	0.00015	0.67087	870.0911	594.6423
A_23_P26	NM_00487	mannose-F	MPDU1,SL	-1.49067	8.13E-11	0.67084	3373.427	2259.685
A_23_P14	NM_01826	membrane	MTCBP-1,	-1.4909	4.84E-11	0.67074	3721.926	2499.271
A_23_P36	NM_15246	Homo sapi	EME1,MMS	-1.49105	0.00097	0.67067	1665.544	1143.389
A_24_P31	NM_18251	Spindle pol	Spc24	-1.49106	0.0004	0.67066	808.2678	553.8682
A_23_P57	NM_03307	Homo sapi	CECR5	-1.49111	1.62E-09	0.67064	3930.256	2634.468
A_23_P77	NM_01660	Homo sapi	JMJD1B,5	-1.49121	5.43E-09	0.6706	2624.984	1757.569
A_32_P22	BC009506	Putative M	MGC3794	-1.49129	4.56E-06	0.67056	3664.578	2455.095
A_23_P74	NM_01441	Homo sapi	CACYBP,S	-1.4913	2.30E-21	0.67055	48943.61	32810.55
A_23_P20	NM_01775	all-trans-13	RetSat,FLJ	-1.49173	0.00032	0.67036	791.3345	539.4688
A_23_P40	NM_02019	Homo sapi	C2orf33,GI	-1.49189	2.48E-07	0.67029	91117.57	61458.19
A_24_P83	XM_08836	Homo sapi	SEC11L2,S	-1.49194	3.41E-10	0.67027	3530.601	2364.141

A_24_P92	NM_03115	heterogene	HNRPA1,H	-1.4924	5.37E-13	0.67006	55002.73	36682.04
A_23_P43	NM_00256	purinergic r	P2RX5,P2	-1.49243	0.00001	0.67005	960.9457	639.1896
A_23_P13	NM_01773	GPI7 prote	GPI7,PRO	-1.49252	6.37E-17	0.67001	3335.506	2234.865
A_24_P12	NM_00590	microtubule	MAP1B,MA	-1.49254	3.42E-09	0.67	2861.45	1917.15
A_23_P42	NM_14463	Homo sapi	CHCHD4,F	-1.49272	3.16E-07	0.66992	4963.465	3343.271
A_23_P30	NM_13928	T-cell activ	TA-PP2C,D	-1.49272	4.14E-08	0.66992	1461.437	976.3138
A_24_P47	NM_00632	RAN, mem	RAN,TC4,A	-1.49272	7.38E-09	0.66992	42090.3	27962.71
A_24_P25	NM_00416	RAB1A, m	RAB1A	-1.49281	0.00002	0.66988	1875.664	1263.558
A_24_P14	NM_00538	protein-L-is	PCMT1	-1.49284	9.50E-15	0.66987	3737.886	2504.678
A_23_P12	NM_02037	Homo sapi	C12orf5	-1.49294	4.73E-11	0.66982	3787.139	2535.338
A_23_P11	AK025895	Hypothetic	FLJ32731	-1.49299	3.96E-08	0.6698	1763.49	1173.67
A_23_P42	NM_01405	Homo sapi	TMEM14A	-1.49311	1.51E-11	0.66974	7079.561	4752.502
A_23_P16	NM_00152	Homo sapi	GTF3C2,K	-1.49312	1.90E-10	0.66974	4723.605	3165.294
A_23_P17	NM_01493	Homo sapi	SEC31L1,A	-1.49315	1.69E-06	0.66973	2243.489	1491.636
A_23_P51	AK096246	Enabled ho	ENAH	-1.49317	8.47E-16	0.66971	11674.17	7805.675
A_23_P15	NM_01805	homolog of	ARHGAP1	-1.49369	2.34E-14	0.66948	2497.337	1672.164
A_23_P67	NM_02457	UDP-N-ace	GALNT14	-1.49388	3.19E-08	0.6694	10849.29	7309.908
A_23_P13	NM_02407	Homo sapi	ALG8,MGC	-1.49392	7.53E-13	0.66938	5906.234	3944.534
A_23_P35	NM_00486	Homo sapi	LITAF,PIG	-1.49439	1.60E-08	0.66917	4470.986	2992.28
A_23_P49	NM_02019	GK001 pro	GK001	-1.49455	2.38E-12	0.6691	9088.391	6086.997
A_24_P27	NM_01493	yeast Sec3	SEC31L1,A	-1.49462	0.00014	0.66906	1225.948	804.3189
A_23_P98	AK002085	Hypothetic	LOC14443	-1.49488	2.60E-10	0.66895	1618.46	1083.821
A_23_P73	NM_17678	Homo sapi	MTMR1	-1.49516	9.40E-07	0.66882	1107.295	743.0504
A_23_P13	NM_02038	Homo sapi	C6orf210,b	-1.49519	0.0002	0.66881	796.3446	523.3852
A_32_P13	NM_17202	Homo sapi	POM121,M	-1.49521	5.13E-10	0.6688	26291.18	17610.63
A_32_P195924		Unknown	THC19928	-1.49541	1.38E-07	0.66871	1845.311	1228.53
A_23_P31	NM_00030	Homo sapi	PON2	-1.49544	1.07E-09	0.6687	1822.76	1223.065
A_23_P50	NM_03301	Homo sapi	TRIM4,RNI	-1.49553	1.30E-19	0.66866	5305.967	3548.055
A_23_P41	AY423045	Vitamin K e	VKORC1L	-1.49561	4.43E-09	0.66862	2631.626	1758.949
A_23_P21	NM_01428	Homo sapi	FREQ,FLU	-1.49576	6.23E-10	0.66856	3108.177	2073.708
A_23_P15	NM_00678	Homo sapi	RALBP1,R	-1.49579	1.29E-06	0.66854	1106.21	737.2221
A_24_P18	AB037807	Ankyrin rep	ANKIB1	-1.496	2.62E-06	0.66845	867.4944	579.7097
A_24_P20	NM_01814	mitochondr	MRPS10,M	-1.49627	1.43E-09	0.66833	4160.78	2781.527
A_24_P13	NM_00409	eukaryotic	EIF2S1,EIF	-1.49635	2.57E-08	0.66829	5414.871	3610.789
A_23_P28	NM_01608	apoptosis r	C2orf28,p1	-1.49675	7.76E-13	0.66811	6251.189	4188.233
A_23_P10	NM_01625	Homo sapi	YTHDF2,H	-1.49675	4.21E-12	0.66811	10039.5	6719.58
A_24_P10	BC013255	Hypothetic	MGC10485	-1.49686	0.00047	0.66807	1146.961	796.3855
A_23_P50	NM_02196	Homo sapi	ABR,MDB	-1.4969	1.02E-16	0.66805	4680.531	3128.014
A_23_P40	NM_00698	transcriptio	MTERF	-1.49693	0.00004	0.66803	1353.361	890.005
A_23_P16	NM_01781	Homo sapi	RG9MTD1	-1.49694	1.54E-27	0.66803	10453.12	7004.026
A_23_P30	NM_00427	Homo sapi	CRSP9,ME	-1.49695	1.53E-06	0.66802	1890.789	1258.397
A_23_P92	NM_02466	Homo sapi	ANKHD1,M	-1.49699	0.00007	0.66801	1718.782	1130.429
A_24_P11	NM_00234	lanosterol s	LSS,OSC	-1.49713	5.68E-06	0.66794	1060.141	715.7612
A_24_P40	NM_01501	Homo sapi	USP33,VD	-1.49714	3.94E-06	0.66794	890.0691	593.6615
A_23_P55	NM_00067	Homo sapi	ADORA2B	-1.49765	2.31E-09	0.66771	2333.091	1553.871
A_23_P14	NM_14565	Homo sapi	RAD52B,R	-1.49767	0.00088	0.6677	555.0504	370.565
A_23_P32	NM_01516	Peptidase r	PMPCA	-1.49786	1.32E-11	0.66762	7035.593	4693.316
A_23_P20	NM_00333	Homo sapi	TXNRD1,T	-1.4983	2.90E-07	0.66742	3126.717	2084.093
A_24_P384200		Unknown	A_24_P38	-1.49848	1.25E-19	0.66735	5779.313	3856.086
A_23_P95	NM_01809	hypothetica	ECT2,FLJ1	-1.49869	3.05E-10	0.66725	8477.496	5669.957
A_23_P68	NM_01430	hypothetica	HSPC117,	-1.4987	1.66E-12	0.66725	13120.95	8786.147
A_23_P24	NM_00173	Compleme	C1S	-1.4992	1.00E-09	0.66702	1470.489	978.6146

A_23_P20	NM_00275	Homo sapi	MAP2K1,M	-1.49944	2.10E-09	0.66691	2811.244	1872.544
A_23_P58	NM_00373	Homo sapi	EIF4EBP3,	-1.49964	0.00044	0.66683	685.176	463.3404
A_23_P44	NM_02290	hypothetica	FLJ12442	-1.49975	0.00045	0.66678	593.1102	392.5953
A_24_P18	NM_01840	Homo sapi	LAPTM4B,	-1.49977	2.98E-15	0.66677	8589.319	5735.619
A_24_P195724		Unknown	ENST0000	-1.49984	8.05E-15	0.66674	50929.88	33894.3
A_23_P60	NM_01786	hypothetica	FLJ20534	-1.49986	1.58E-14	0.66673	5406.154	3610.919
A_23_P30	NM_15226	Homo sapi	RPUSD2,C	-1.50012	0.00001	0.66661	6893.834	4666.983
A_23_P14	NM_01434	Homo sapi	MTCH1,PS	-1.50025	3.10E-14	0.66656	21927.52	14653.53
A_32_P13	NM_00566	Homo sapi	C5orf18,DF	-1.50046	6.10E-15	0.66646	9327.799	6233.872
A_23_P14	NM_00174	Homo sapi	CANX,CNX	-1.50046	2.58E-09	0.66646	15124.94	10125.61
A_23_P15	NM_03228	Homo sapi	ZCCHC9,D	-1.50066	1.00E-08	0.66638	3283.181	2190.594
A_23_P34	NM_02275	Homo sapi	SFXN1,FL	-1.50081	2.19E-08	0.66631	2479.636	1653.636
A_24_P12	NM_02029	aldo-keto r	AKR1B10,I	-1.50094	1.99E-13	0.66625	74278.27	49621.27
A_24_P16	BC047111	Nuclear rec	NCOA2	-1.50103	0.00009	0.66621	881.7838	576.2326
A_23_P25	NM_00209	Homo sapi	GTF3A,AP	-1.50108	1.14E-19	0.66619	13016.79	8666.548
A_32_P69	NM_20734	Homo sapi	NM_20734	-1.5011	1.38E-08	0.66618	1281.031	852.0535
A_23_P99	NM_00398	Homo sapi	SLC7A7,LF	-1.50117	1.20E-09	0.66615	2175.716	1446.213
A_23_P40	NM_00714	Homo sapi	ZNF161,DE	-1.50118	2.26E-06	0.66614	1244.813	821.95
A_24_P68	AB002449	Hypothetic	AB002449	-1.50119	0.00068	0.66614	667.5425	453.2832
A_32_P17	XM_37988	similar to H	LOC40256	-1.5012	1.44E-13	0.66614	63927.63	42687.48
A_23_P86	NM_15374	Homo sapi	PIGC,GPI2	-1.50125	8.35E-08	0.66611	1371.992	918.0073
A_24_P37	NM_01522	retinoblast	RAP140,se	-1.50138	0.00002	0.66605	1508.334	988.2824
A_24_P11	NM_00437	COX15 (ye	COX15	-1.5014	0.00041	0.66604	644.5593	426.7134
A_23_P41	AB023148	PH domain	KIAA0931	-1.5014	8.67E-06	0.66605	842.0515	560.6934
A_24_P82	NM_01868	zinc finger	ZNF313,RI	-1.50153	4.97E-11	0.66599	2168.141	1442.862
A_23_P16	NM_01630	Homo sapi	SS18L2,KI	-1.50153	1.37E-11	0.66599	11339.61	7576.021
A_23_P54	NM_01780	hypothetica	OSGEP,G	-1.50157	1.21E-08	0.66597	2675.867	1772.25
A_23_P89	NM_00614	LIM and SH	LASP1,ML	-1.50194	1.86E-10	0.6658	7964.972	5306.666
A_23_P12	NM_03229	hypothetica	DCUN1D5,	-1.50218	4.77E-08	0.6657	12769.53	8517.705
A_24_P38	NM_00214	heterogene	HNRPK,CS	-1.50256	6.74E-12	0.66553	21128.36	13995.75
A_23_P14	NM_03308	Homo sapi	ALG2,CDG	-1.50267	1.61E-09	0.66548	1850.014	1227.767
A_23_P11	NM_00701	Homo sapi	DDX52,RC	-1.50309	4.82E-10	0.6653	1487.56	991.7579
A_23_P37	NM_03330	hypothetica	MGC4655	-1.50314	1.41E-14	0.66527	4596.47	3057.901
A_24_P24	NM_00538	protein-L-is	PCMT1	-1.50318	6.32E-10	0.66526	4174.71	2766.046
A_32_P13	BC043383	Homo sapi	BC043383	-1.50331	5.63E-06	0.6652	1217.285	799.7145
A_23_P66	NM_02275	Family with	C18orf11	-1.50365	0.00016	0.66505	1119.391	728.3214
A_23_P21	NM_01867	smooth mu	SMAP-1,FI	-1.50366	2.45E-12	0.66504	4049.384	2694.884
A_23_P22	NM_00563	Homo sapi	SYBL1,VA	-1.50372	4.78E-18	0.66502	3218.265	2140.889
A_23_P20	NM_03285	Homo sapi	MUM1,MU	-1.50378	2.07E-09	0.66499	1313.821	871.7057
A_32_P50	NM_00459	spermine s	SMS,SRS,	-1.50383	2.63E-11	0.66497	1541.864	1025.063
A_24_P38	NM_17321	Homo sapi	ST6GAL1,	-1.50384	0.00001	0.66496	913.6942	612.078
A_23_P10	NM_00377	UDP-Gal:b	B4GALT3,I	-1.50387	4.12E-09	0.66495	2868.264	1907.771
A_23_P94	NM_01217	Homo sapi	FBXO25,F	-1.50408	0.00007	0.66486	945.4807	620.5006
A_23_P14	NM_00302	Homo sapi	SH3BGRL,	-1.50457	6.66E-14	0.66464	1821.371	1211.124
A_23_P50	NM_02479	Homo sapi	FAM57A,C	-1.50468	1.94E-08	0.66459	1997.899	1335.766
A_24_P27	NM_00178	CD63 antig	CD63,MLA	-1.50519	0.00004	0.66437	63021.51	40939.82
A_23_P40	AB033045	Homo sapi	KIAA1219	-1.50542	6.19E-07	0.66426	1092.545	730.7263
A_23_P34	NM_01502	Homo sapi	POP1,KIA	-1.50547	4.76E-12	0.66424	2606.967	1734.877
A_23_P75	NM_03279	Homo sapi	RPUSD4,F	-1.50565	6.98E-07	0.66416	3107.822	2053.834
A_23_P39	NM_01223	Homo sapi	PTRF,FKS	-1.50575	6.75E-11	0.66412	28445.59	18982.46
A_23_P21	NM_02245	hypothetica	NSD1,STC	-1.5059	0.0001	0.66405	688.0894	456.6265
A_24_P19	NM_02466	Hypothetic	FLJ11838	-1.50626	1.59E-06	0.6639	1266.84	847.2345

A_23_P40	NM_00245	Homo sapi	MTIF2	-1.5069	1.47E-09	0.66362	5669.103	3781.825
A_23_P25	NM_01793	hypothetica	FLJ20718	-1.50692	4.50E-08	0.66361	8234.684	5487.501
A_24_P19	NM_00285	peroxisom	PEX19,PX	-1.50722	9.78E-08	0.66347	2319.481	1527.572
A_23_P14	NM_01437	Homo sapi	REPIN1,A	-1.50725	8.78E-06	0.66346	13246.35	8834.822
A_23_P13	NM_01847	TPA regula	TPARL,TM	-1.50726	1.12E-06	0.66346	4950.552	3288.007
A_24_P38	NM_13083	Homo sapi	UBE3A,AS	-1.50742	4.36E-11	0.66338	5120.987	3399.885
A_23_P76	NM_00310	Homo sapi	SNX2,TRG	-1.50748	9.97E-06	0.66336	10117.13	6788.7
A_23_P25	AL832848	Tetratricop	TTC7B	-1.50756	1.58E-11	0.66332	2445.249	1620.983
A_23_P33	NM_01439	staufen (D	STAU2,39	-1.50762	0.00053	0.6633	623.2198	410.1334
A_23_P80	NM_02487	hypothetica	FLJ12748	-1.50773	2.48E-06	0.66325	2040.321	1371.577
A_23_P87	NM_03259	Homo sapi	FBXL10,C	-1.50811	0.00017	0.66308	508.8047	338.2802
A_32_P20	NM_02031	hypothetica	CIAPIN1,P	-1.50838	1.78E-07	0.66296	2044.698	1351.894
A_24_P34	NM_03302	tripartite m	TRIM33,PT	-1.50843	0.00002	0.66294	1626.693	1064.761
A_24_P727868		VAPA_HU	THC18449	-1.50846	3.61E-13	0.66293	6389.487	4244.56
A_23_P14	NM_00432	Homo sapi	BMPR1A,A	-1.50872	0.00017	0.66281	765.8809	498.8221
A_24_P10	NM_18169	Homo sapi	PPP2R1B,	-1.50892	1.71E-07	0.66272	1248.102	831.9401
A_24_P16	NM_00088	IMP (inosi	IMPDH2,IM	-1.50924	5.54E-15	0.66258	4404.231	2913.122
A_23_P10	NM_01847	Homo sapi	FAM45B,H	-1.50928	0.00009	0.66257	7748.957	5191.102
A_24_P57	AK025344	Homo sapi	AK025344	-1.50937	1.74E-09	0.66253	3786.262	2507.974
A_23_P14	NM_03202	Homo sapi	FUCA2,MG	-1.50942	9.69E-12	0.66251	3109.164	2060.1
A_32_P56415		Unknown	A_32_P56	-1.50952	1.10E-06	0.66246	926.0858	615.0892
A_23_P64	NM_00572	Homo sapi	TSFM,EF-	-1.50953	1.02E-15	0.66246	20684.48	13732.28
A_23_P50	NM_00340	Homo sapi	YWHAB,H	-1.50969	1.13E-21	0.66239	60211.09	39895.11
A_23_P37	NM_01598	Homo sapi	UBE2D4,H	-1.50972	0.00004	0.66237	916.7634	617.307
A_23_P81	NM_01328	Homo sapi	MAT2B,TG	-1.51042	6.73E-06	0.66207	2584.712	1741.805
A_32_P18	NM_00252	nucleophos	NPM1,B23	-1.51048	2.27E-12	0.66204	60701.99	40156.7
A_24_P17	BC011136	Transformi	TGFBRAP	-1.5106	0.00005	0.66199	892.6393	600.0607
A_23_P14	NM_00500	Homo sapi	NDUFAB1,	-1.51085	1.42E-22	0.66188	34773.63	23011.13
A_32_P23	NM_17587	Homo sapi	EXOC8,EX	-1.51094	2.51E-07	0.66184	1573.782	1036.167
A_24_P54	NM_00550	CD36 antig	SCARB2,C	-1.51097	0.00025	0.66183	703.1339	460.7562
A_23_P16	NM_17060	Homo sapi	MLX,MAD	-1.51105	7.65E-06	0.66179	1342.709	898.3831
A_23_P15	NM_00125	Homo sapi	CDC5L,hC	-1.51119	2.13E-07	0.66173	2565.267	1693.614
A_23_P43	NM_01422	Homo sapi	NDUFA8,P	-1.51124	5.66E-12	0.66171	51575.37	34150.84
A_24_P38	NM_13079	Nucleolar	NOL6	-1.51148	2.44E-11	0.6616	3524.58	2327.447
A_23_P24	NM_01463	Homo sapi	MICAL2,KI	-1.5116	2.74E-06	0.66155	1330.823	870.0826
A_24_P17	NM_00306	solute carri	SLC22A5,	-1.51177	5.73E-18	0.66148	2530.449	1673.944
A_23_P74	NM_17845	Hypothetic	MGC5428	-1.5121	1.15E-06	0.66133	1497.949	999.624
A_24_P40	NM_00641	solute carri	SLC35A1,	-1.51221	1.43E-06	0.66129	1583.812	1037.89
A_23_P54	NM_01433	CREBBP/E	CRI1,EID1	-1.51244	2.77E-11	0.66118	25692.77	17104.35
A_23_P90	NM_01478	Homo sapi	RB1CC1,C	-1.51255	1.87E-10	0.66113	1729.138	1141.371
A_23_P10	NM_00493	Homo sapi	DDX1,DBF	-1.51274	8.57E-13	0.66105	43743.76	29029.02
A_32_P234275		Unknown	A_32_P23	-1.513	0.00047	0.66094	839.4481	541.6969
A_32_P22	AB014574	KIAA0674	KIAA0674	-1.51311	3.80E-07	0.66089	1285.13	845.3101
A_23_P25	NM_01781	hypothetica	CHCHD3,F	-1.51345	2.68E-10	0.66074	4236.213	2806.764
A_23_P37	NM_13879	Homo sapi	OACT2,FL	-1.51353	0.00006	0.66071	773.8427	508.8869
A_23_P66	NM_01801	Homo sapi	MED9,ME	-1.51366	0.00012	0.66065	1555.741	1043.901
A_23_P21	NM_00547	Homo sapi	GNE,NM,D	-1.51371	0.00003	0.66063	2453.511	1616.026
A_23_P83	NM_00307	Homo sapi	SMARCC1	-1.5139	6.71E-09	0.66054	2442.556	1606.663
A_32_P49	NM_00252	nucleophos	NPM1,B23	-1.51406	4.31E-20	0.66047	109797.9	72473.09
A_23_P66	BC047010	Similar to T	LOC40050	-1.5141	2.35E-08	0.66046	2869.184	1886.769
A_23_P60	NM_01811	Homo sapi	TMEM38B	-1.51434	2.62E-12	0.66035	3121.9	2062.731
A_24_P14	NM_01558	DKFZP586	POLDIP2,F	-1.51445	3.95E-13	0.66031	9227.679	6095.357

A_23_P79	NM_00018	Homo sapi	HADHB,HA	-1.51445	1.01E-11	0.6603	13417.69	8893.779
A_23_P35	NM_01333	Homo sapi	ALG6	-1.51451	1.05E-08	0.66028	1526.938	1012.556
A_23_P11	NM_00693	Homo sapi	SUMO2,HS	-1.51479	9.06E-11	0.66016	26558.79	17539.41
A_23_P12	NM_02120	E-1 enzym	MASA,MST	-1.51498	8.15E-12	0.66008	3017.391	1990.564
A_23_P36	NM_01480	KIAA0528	KIAA0528	-1.51503	2.93E-08	0.66005	1923.773	1265.028
A_24_P36	BC030112	Homeodon	HIPK3	-1.51516	1.52E-07	0.65999	2291.752	1505.559
A_23_P12	NM_00282	Homo sapi	QSCN6,Q6	-1.51538	3.44E-16	0.6599	5169.893	3408.178
A_23_P72	AF157326	TBP-intera	TIP120A	-1.51553	0.00025	0.65984	584.4745	385.7924
A_23_P36	NM_01984	Homo sapi	SLC10A3,F	-1.51583	1.16E-11	0.6597	3293.82	2170.76
A_23_P99	NM_00345	zinc finger	ZNF198,FI	-1.5159	0.00001	0.65967	1267.976	849.8676
A_23_P92	NM_01742	Homo sapi	NUP54,MG	-1.51591	3.84E-14	0.65967	4461.487	2939.438
A_23_P30	NM_14459	Homo sapi	C10orf32,A	-1.51615	4.05E-15	0.65956	2323.787	1532.727
A_23_P93	NM_02029	Homo sapi	AKR1B10,I	-1.51668	2.68E-09	0.65934	100805.8	66780.03
A_23_P36	NM_14526	Homo sapi	NUDCD2	-1.51701	3.78E-06	0.65919	11657.39	7692.771
A_32_P192615		Unknown	THC18582	-1.51718	7.55E-12	0.65912	3308.643	2176.987
A_24_P20	NM_14459	Homo sapi	C10orf32,A	-1.51718	3.70E-06	0.65912	1629.709	1064.029
A_24_P12	NM_01241	rab3 GTPa	RAB3GAP2	-1.51722	1.06E-08	0.6591	5363.318	3550.299
A_23_P82	NM_00041	Homo sapi	IFNGR1,CI	-1.51738	4.65E-17	0.65903	4300.406	2833.541
A_24_P94	AB023198	Phosphatid	PIP5K3	-1.5174	0.00021	0.65902	729.0263	474.7862
A_24_P27	AL713674	Male sterili	MLSTD2	-1.51745	0.00007	0.659	924.8303	598.1101
A_23_P15	NM_00717	Homo sapi	PRSS23,S	-1.51751	3.79E-17	0.65898	23209.75	15315.61
A_23_P38	NM_03298	caspase 2,	CASP2,ICH	-1.51752	0.00033	0.65897	1346.607	868.4246
A_23_P84	NM_00266	Pleiotropic	PLRG1	-1.5178	7.94E-11	0.65885	3875.363	2556.576
A_23_P76	NM_15348	Homo sapi	NUP155,N	-1.51805	5.97E-07	0.65874	2180.3	1447.1
A_24_P49	NM_01786	hypothetica	FLJ20534	-1.51806	2.59E-07	0.65873	1320.645	877.387
A_23_P11	NM_00576	Homo sapi	ATP6AP2,I	-1.51827	8.59E-15	0.65864	3479.987	2291.539
A_24_P26	NM_13325	Homo sapi	LRPPRC,L	-1.51831	4.38E-09	0.65863	6984.611	4625.386
A_23_P37	NM_00678	Homo sapi	RALBP1,R	-1.51852	5.03E-20	0.65854	5842.644	3846.885
A_23_P10	NM_18292	Homo sapi	KTN1,CG1	-1.51878	0.00001	0.65842	17422.24	11569.76
A_23_P12	NM_03196	Homo sapi	CCNB1,CC	-1.51905	1.68E-14	0.65831	13590.04	8935.478
A_24_P84	NM_03256	emopamil b	EBPL,EBR	-1.5198	5.02E-10	0.65798	4614.198	3039.634
A_23_P12	NM_00129	Homo sapi	CLNS1A,C	-1.51985	3.21E-12	0.65796	24260.84	16013.03
A_23_P74	D50923	KIAA0133	KIAA0133	-1.51992	1.50E-10	0.65793	2683.864	1761.474
A_24_P30	NM_00459	spermine s	SMS,SRS,	-1.52034	7.99E-15	0.65775	13723.76	9043.533
A_24_P39	NM_20749	Homo sapi	FLJ44896	-1.52061	0.0001	0.65763	695.5249	455.1336
A_24_P10	NM_13927	Homo sapi	MCFD2,F5	-1.52064	0.00054	0.65762	563.0396	370.6643
A_24_P69	NM_00208	glutamic-o	GOT2,FLJ4	-1.5207	0.00013	0.65759	1151.851	740.7233
A_24_P35	NM_00448	UDP-N-ace	GALNT2,G	-1.52075	0.00042	0.65757	2192.565	1422.887
A_24_P39	AK095244	Cytochrom	CYB561	-1.52095	1.25E-19	0.65749	7092.513	4665.529
A_24_P37	NM_01510	Homo sapi	PLXND1,P	-1.52115	4.35E-09	0.6574	1175.851	772.3038
A_23_P44	NM_00094	Homo sapi	PRIM2A,p5	-1.52123	0.00063	0.65736	1064.98	677.1578
A_32_P43	NM_00574	accessory	BCAP31,C	-1.52125	4.33E-13	0.65736	51078.41	33462.75
A_23_P13	NM_01605	CGI-119 pr	TMBIM4,S	-1.52151	1.25E-16	0.65724	3043.008	2000.079
A_23_P66	NM_00227	Homo sapi	KRT19,K19	-1.52164	5.86E-16	0.65718	64436.18	42236.91
A_23_P35	NM_01829	Homo sapi	CWF19L1,	-1.52181	2.52E-09	0.65711	1967.795	1296.758
A_24_P76	AK056691	Germ cell a	GSG2	-1.52189	5.98E-10	0.65708	4091.746	2684.148
A_24_P12	AK021904	Hypothetica	RPL7L1	-1.52192	2.56E-08	0.65706	1735.735	1145.979
A_24_P17	NM_00624	protein pho	PPP1R2,IF	-1.52212	1.18E-07	0.65698	2831.834	1860.331
A_23_P20	BC039050	ADP-ribosy	ARL6IP2	-1.52213	1.55E-10	0.65697	6680.472	4397.772
A_24_P39	NM_00698	melanoma	MAGED1,N	-1.52219	1.69E-06	0.65695	1639.25	1066.656
A_23_P11	NM_02236	Homo sapi	PJA1,RNF	-1.52227	1.50E-07	0.65692	1898.668	1238.678
A_23_P37	NM_02406	Homo sapi	SPATA5L1	-1.52249	0.00011	0.65682	5179.856	3385.352

A_23_P10	NM_03282	hypothetica	FLJ14668	-1.5225	3.83E-09	0.65681	15668.3	10313.14
A_23_P21	NM_00601	SWI/SNF r	ARID1A,B1	-1.5226	0.00029	0.65677	1069.532	721.421
A_24_P20	NM_00156	eukaryotic	EIF3S6,IN	-1.52273	9.61E-24	0.65672	36643.38	24066.48
A_23_P38	NM_01824	hypothetica	CCDC25,F	-1.52309	3.83E-12	0.65656	2740.3	1799.209
A_23_P40	BC029424	Similar to F	LOC49386	-1.52311	4.17E-19	0.65655	4201.742	2759.177
A_24_P633686		C560_HUM	THC19557	-1.52339	0.00002	0.65643	6289.575	4097.907
A_24_P170283		Unknown	ENST0000	-1.5235	0.00009	0.65638	1127.441	756.7347
A_23_P58	NM_00406	Homo sapi	CCNG1,CC	-1.52366	8.52E-13	0.65632	2892.801	1894.694
A_23_P37	NM_00728	Homo sapi	OIP5,5730	-1.52397	2.42E-09	0.65618	9337.949	6162.778
A_23_P13	NM_00435	CD81 antig	CD81,S5.7	-1.52405	8.48E-16	0.65615	10933.38	7180.957
A_23_P21	NM_01534	leucyl-tRNA	LARS2,LE	-1.52432	1.80E-11	0.65603	3060.99	2007.646
A_23_P50	NM_00539	PHD finger	PHF2,GRC	-1.52458	0.00001	0.65592	2258.474	1472.417
A_24_P72	NM_01532	KIAA0828	KIAA0828	-1.52466	7.51E-07	0.65588	1202.899	781.3366
A_32_P131797		Unknown	A_32_P13	-1.52494	9.40E-06	0.65576	1247.666	832.0447
A_24_P20	AB033038	KIAA1212	KIAA1212	-1.52495	0.00022	0.65576	835.782	537.0049
A_23_P20	NM_01481	centrosome	CAP350,gr	-1.52502	1.44E-11	0.65573	1598.884	1047.484
A_24_P27	NM_00599	tubulin-spe	TBCD,KIA	-1.52514	2.22E-16	0.65568	11738.79	7705.292
A_23_P10	NM_02089	Homo sapi	ZBTB4,KA	-1.52534	5.49E-09	0.65559	1552.75	1020.737
A_23_P87	NM_00198	Homo sapi	ESD	-1.52537	8.13E-08	0.65558	4472.821	2949.25
A_23_P41	NM_01534	Homo sapi	BRRN1,CA	-1.52544	1.73E-09	0.65555	2331.031	1532.074
A_23_P60	NM_00566	Homo sapi	VDAC3,HD	-1.52552	2.15E-22	0.65551	8084.739	5300.383
A_24_P30	NM_05293	pro-oncosis	PORIMIN,H	-1.52571	7.07E-08	0.65543	9195.803	6070.098
A_23_P59	NM_01601	CGI-74 pro	LUC7L2,C	-1.52572	9.60E-07	0.65543	2341.556	1526.967
A_23_P31	NM_00616	Homo sapi	NFYB,HAF	-1.52594	8.63E-09	0.65533	3387.411	2222.933
A_23_P20	NM_02462	Homo sapi	SMC6L1,S	-1.52596	2.28E-10	0.65532	1561.495	1027.993
A_23_P38	NM_00416	RAB1A, m	RAB1A	-1.52616	1.95E-09	0.65524	4047.464	2637.3
A_24_P39	NM_01323	like mouse	ATXN10,E	-1.52622	3.43E-07	0.65521	2945.845	1932.567
A_23_P92	BC033524	RNA bindir	RBM27	-1.52627	8.25E-09	0.65519	1783.349	1163.52
A_23_P17	NM_01414	HSPC274	C20orf30,H	-1.52642	1.05E-07	0.65513	3329.937	2181.991
A_24_P21	XM_37661	Homo sapi	ENST0000	-1.52644	1.19E-16	0.65512	9946.967	6525.747
A_24_P33	NM_00693	SMT3 (sup	SUMO2,HS	-1.52651	2.33E-10	0.65509	35622	23424.57
A_23_P14	NM_00655	GAP-assoc	KHDRBS1	-1.5267	2.96E-10	0.65501	4491.186	2953.398
A_23_P13	NM_00687	Homo sapi	ELF2,EU32	-1.5268	0.00012	0.65496	1136.125	760.7062
A_24_P238286		Unknown	A_24_P23	-1.5271	0.00045	0.65484	772.7772	520.7396
A_32_P17	NM_00290	reticulocalb	RCN2,E6B	-1.52731	8.26E-11	0.65474	11219.52	7392.41
A_23_P72	NM_00020	Homo sapi	IF,FI,factor	-1.52731	0.00009	0.65475	1348.062	866.0616
A_24_P56	NM_00134	defender a	DAD1	-1.52733	3.45E-09	0.65474	14882.59	9811.06
A_23_P24	NM_01475	Homo sapi	BMS1L,KIA	-1.52747	1.58E-10	0.65468	2847.609	1863.622
A_23_P11	NM_00127	Homo sapi	CHEK1,CH	-1.52765	4.68E-09	0.6546	2295.071	1494.867
A_24_P19	NM_01340	fatty acid d	FADS1,D5	-1.52771	5.76E-11	0.65458	14465.29	9508.882
A_32_P16	NM_02217	Williams-B	WBSCR1,E	-1.52816	2.21E-14	0.65438	29517.95	19291.12
A_23_P31	NM_00430	Homo sapi	ARHGAP1	-1.52818	2.31E-06	0.65437	9208.486	6049.578
A_23_P31	NM_01533	Zinc finger,	ZDHHC17	-1.5284	0.00016	0.65428	1229.748	786.8733
A_23_P25	NM_02162	Homo sapi	PRDM13,F	-1.52848	3.62E-08	0.65425	2001.824	1332.264
A_32_P32	NM_01764	hypothetica	FAM29A,R	-1.52864	1.37E-08	0.65417	1576.72	1026.39
A_23_P44	NM_01407	Homo sapi	MRPL13,L	-1.52876	1.76E-11	0.65412	13675.44	8985.388
A_23_P65	NM_01630	60S riboso	C15orf15,L	-1.52877	5.19E-16	0.65412	10242.95	6716.522
A_23_P53	NM_03057	hypothetica	MGC10993	-1.5288	2.22E-12	0.65411	2561.631	1675.937
A_23_P13	NM_00500	Homo sapi	NDUFS1,C	-1.52886	4.04E-19	0.65408	6618.777	4329.401
A_32_P124580		Unknown	A_32_P12	-1.52899	6.85E-11	0.65402	1664.857	1087.17
A_23_P35	NM_15230	hypothetica	MGC24665	-1.52902	0.00049	0.65401	2929.475	1974.539
A_24_P170384		Unknown	A_24_P17	-1.52919	0.00062	0.65394	675.9771	432.7593

A_23_P39	AL080064	SEC24 rela	SEC24A	-1.52936	1.63E-06	0.65387	1137.634	737.8145
A_23_P75	NM_01233	Homo sapi	MYO10,KI	-1.52942	0.00004	0.65384	2756.957	1798.448
A_23_P72	NM_00420	Homo sapi	PDCD8,AI	-1.52961	3.09E-14	0.65376	15548.74	10167.11
A_23_P12	NM_00121	Homo sapi	CALU	-1.52977	1.31E-06	0.65369	2112.098	1379.605
A_23_P14	NM_01872	Homo sapi	MYO5C,m	-1.53005	0.00011	0.65357	682.8355	443.4371
A_23_P10	NM_02208	hypothetica	HPS4,LE,K	-1.53006	1.30E-10	0.65357	2148.243	1399.709
A_23_P88	NM_02015	Homo sapi	C15orf24,H	-1.53024	1.00E-17	0.65349	39351.09	25697.89
A_23_P42	NM_19848	Homo sapi	RPL7L1,M	-1.5304	4.88E-08	0.65342	3061.952	1993.649
A_24_P94	AB051476	B double p	BDP1	-1.53055	0.00064	0.65336	973.21	615.0992
A_23_P98	AB020674	Mlx interac	MONDOA	-1.53076	2.33E-09	0.65327	3308.822	2157.868
A_23_P15	NM_02234	chromoson	HCAP-G,C	-1.53083	6.60E-12	0.65324	6666.495	4363.832
A_23_P12	NM_02275	hypothetica	C6orf106,M	-1.53086	1.27E-22	0.65323	6994.45	4568.435
A_24_P475814		ALU8_HUM	THC18082	-1.53087	8.45E-10	0.65322	1901.362	1244.496
A_23_P21	NM_01622	Homo sapi	SNX9,SDP	-1.53087	7.69E-06	0.65322	2030.102	1311.275
A_23_P94	NM_01820	Homo sapi	TBC1D13,I	-1.53109	5.20E-07	0.65313	13103.45	8597.745
A_24_P24	NM_00018	hydroxyacy	HADHA,GE	-1.53124	1.47E-11	0.65307	2762.499	1803.471
A_23_P37	NM_00358	Homo sapi	DUSP11,P	-1.5313	9.68E-08	0.65304	1056.549	689.5079
A_24_P13	AK090397	Hypothetic	MGC15875	-1.53143	0.0001	0.65298	1180.218	792.2905
A_23_P28	NM_02119	Homo sapi	CTDSP1,S	-1.53144	6.03E-23	0.65298	8531.838	5570.201
A_24_P26	NM_01775	hypothetica	FLJ20297,	-1.53145	2.58E-17	0.65297	4530.371	2956.928
A_32_P30	NM_00660	unactive pr	PTGES3,P	-1.53192	1.22E-21	0.65277	109210.2	71357.89
A_24_P27	NM_00573	ARP1 (acti	ACTR1A,A	-1.53204	2.43E-12	0.65273	1828.133	1194.239
A_23_P15	NM_00710	signal sequ	SSR3,TRA	-1.53211	1.45E-07	0.65269	5594.072	3675.9
A_32_P56	NM_01471	KIAA0738	KIAA0738	-1.5324	2.96E-06	0.65257	1149.629	741.7987
A_23_P14	NM_00704	Homo sapi	FGFR1OP,	-1.53249	0.00003	0.65253	1550.529	997.3458
A_23_P21	NM_02248	PC3-96 pro	ATG3,APG	-1.5325	4.03E-16	0.65253	4594.679	3002.517
A_23_P68	NM_00414	Homo sapi	DRG1,NED	-1.53272	1.03E-11	0.65244	4674.49	3041.271
A_23_P65	NM_00013	Homo sapi	FBN1,FBN	-1.53288	0.00001	0.65237	1185.308	785.8734
A_23_P20	NM_00461	Homo sapi	TRAF5,RN	-1.53289	3.69E-08	0.65236	1131.556	741.9623
A_24_P39	NM_17307	Homo sapi	RUNDC1,D	-1.53289	0.00007	0.65236	887.3422	568.5352
A_24_P26	AK055669	Transportir	TNPO1	-1.5332	6.90E-16	0.65223	4354.579	2843.899
A_23_P37	NM_00735	MLN51 pro	CASC3,BT	-1.53332	3.45E-13	0.65218	21194.39	13871.56
A_23_P20	NM_01492	Homo sapi	KIAA0971	-1.53336	0.00037	0.65216	793.0991	530.9819
A_23_P14	NM_00132	Homo sapi	CSTF2,Cst	-1.53346	1.61E-15	0.65212	4450.207	2900.598
A_23_P15	NM_02277	hypothetica	FLJ13912	-1.53358	2.33E-06	0.65207	5218.191	3399.423
A_24_P19	NM_00487	mannose-F	MPDU1,SL	-1.5338	0.00035	0.65198	2032.59	1330.495
A_23_P63	NM_03075	nuclear rec	NRBF2,CC	-1.53454	0.00005	0.65166	694.874	453.6468
A_23_P74	NM_14569	Homo sapi	CDCA1,NU	-1.5347	7.32E-19	0.65159	4028.947	2624.552
A_24_P79	NM_00510	RNA bindir	RBM8A,Y1	-1.53471	4.51E-17	0.65159	2534.142	1651.175
A_24_P36	BC029755	Synaptogy	SYNGR2	-1.53475	3.10E-15	0.65157	59600.91	38709.16
A_23_P11	NM_00345	zinc finger	ZNF207,DI	-1.53481	6.85E-11	0.65155	7816.67	5107.391
A_23_P34	AK094181	Tetratricop	TTC15	-1.53532	0.00015	0.65133	698.0527	462.0396
A_32_P16	BC041925	Solute carr	SLC7A11	-1.5354	1.95E-18	0.6513	5226.104	3402.729
A_24_P324506		Unknown	A_24_P32	-1.53543	1.77E-20	0.65128	6048.112	3940.526
A_23_P11	NM_01544	Homo sapi	CNOT10,F	-1.53557	4.81E-12	0.65123	3792.907	2471.415
A_32_P175557		Unknown	A_32_P17	-1.53563	7.68E-07	0.6512	2119.235	1385.264
A_23_P38	NM_00052	Homo sapi	HOXA13,H	-1.53565	0.00063	0.65119	553.0432	358.4496
A_32_P17	NM_01627	transducer	TOB2,TOB	-1.5358	1.16E-06	0.65113	2900.042	1878.851
A_23_P50	NM_03285	AUT (S. ce	ATG4C,AP	-1.5363	1.24E-08	0.65091	1497.132	979.345
A_23_P10	NM_00642	Chaperonir	CCT7	-1.53638	2.34E-11	0.65088	37158.84	24180.8
A_24_P36	NM_01467	basic leuci	BZW1,orf,E	-1.5367	9.16E-10	0.65075	6971.061	4554.355
A_24_P204257		Unknown	A_24_P20	-1.53692	6.47E-13	0.65065	4117.861	2674.672

A_23_P82	NM_00148	glioblastom	GBAS,NIP	-1.53715	0.00001	0.65056	815.5668	529.3187
A_24_P10	NM_00210	H3 histone	H3F3A,H3	-1.53756	7.23E-10	0.65038	158119.4	102347.8
A_23_P88	NM_00287	Homo sapi	RAD51,RE	-1.53765	1.86E-11	0.65034	3897.176	2538.338
A_23_P25	NM_00015	Homo sapi	GALC	-1.53782	2.66E-14	0.65027	2041.455	1326.477
A_23_P25	NM_00262	Homo sapi	PFN2,PFL,	-1.53786	5.19E-10	0.65025	1491.34	967.9321
A_23_P20	NM_01504	Homo sapi	ALS2CR3,	-1.53788	0.00016	0.65025	707.9035	469.5971
A_24_P29	AK125387	Collagen, t	COL4A3BP	-1.53801	0.00006	0.65019	724.2154	473.1461
A_23_P14	NM_01547	Homo sapi	C18orf10,L	-1.53813	2.11E-18	0.65014	4490.244	2918.852
A_32_P20	NM_18170	Quiescin Q	QSCN6L1	-1.53817	9.23E-07	0.65012	1106.71	714.5547
A_32_P46	BC035136	Hypothetic	DKFZp434	-1.5389	1.31E-08	0.64981	1791.19	1184.826
A_23_P43	AL161973	Myeloid/lyn	MLLT4	-1.53899	0.00073	0.64978	791.812	497.7847
A_32_P15	NM_20741	Similar to F	MGC57827	-1.53915	1.63E-06	0.64971	1630.543	1068.558
A_32_P27	NM_01563	Chromosor	C10orf61	-1.5393	4.42E-15	0.64965	4714.902	3066.959
A_23_P33	NM_05816	Homo sapi	TRIM6,RN	-1.53958	0.00011	0.64953	738.6937	474.0284
A_23_P21	NM_03278	carbonic re	CBR4,FLJ	-1.5396	1.54E-07	0.64952	1324.291	866.4595
A_24_P32	NM_00036	tuberous s	TSC1,LAM	-1.53973	0.00002	0.64946	912.6378	583.1545
A_24_P13	NM_01461	C9orf10 pr	C9orf10,D	-1.53994	5.73E-19	0.64937	2939.621	1908.861
A_32_P20	AK093982	CDNA FLJ	AK093982	-1.54021	0.0004	0.64926	646.9841	428.072
A_23_P92	NM_00586	Homo sapi	SDCCAG1	-1.54027	1.88E-10	0.64924	2488.16	1613.159
A_23_P76	NM_00600	Homo sapi	UCHL3	-1.54033	3.39E-11	0.64921	5881.518	3827.913
A_23_P20	NM_00010	Homo sapi	CYP1B1,C	-1.54034	9.83E-16	0.64921	2448.645	1590.016
A_23_P65	NM_02513	hypothetica	FLJ21439,	-1.54036	4.79E-07	0.6492	1772.837	1159.501
A_32_P16	AK096786	TAK1-bindi	TAB3	-1.5404	0.0001	0.64918	710.0868	456.9829
A_23_P67	NM_00320	Homo sapi	TCF3,E2A,	-1.54056	1.99E-16	0.64912	3863.995	2504.948
A_23_P11	NM_00336	Homo sapi	UQCRC2	-1.54058	4.47E-17	0.64911	23155.91	15064.52
A_23_P37	NM_03080	hypothetica	DKFZP564	-1.54089	6.85E-15	0.64898	2823.816	1834.761
A_23_P11	NM_01484	KIAA0196	KIAA0196	-1.54091	5.47E-08	0.64897	4266.259	2798.533
A_24_P673209		Unknown	A_24_P67	-1.54132	1.16E-16	0.6488	7313.055	4733.032
A_23_P74	NM_01512	Mid-1-relat	CLCC1,MC	-1.54139	1.37E-07	0.64877	1214.664	782.9419
A_23_P15	AL832489	KIAA1970	KIAA1970	-1.54154	0.00006	0.6487	1064.611	676.1163
A_24_P39	NM_00178	cell divisior	CDC25A,C	-1.54166	5.76E-12	0.64865	1700.629	1104.238
A_24_P339869		Z295_HUM	THC18809	-1.54175	0.00004	0.64862	843.4417	539.5542
A_23_P45	NM_00428	chromodon	CHD1L,AL	-1.54178	1.16E-17	0.6486	2946.451	1911
A_23_P33	NM_15241	hypothetica	C8orf38,M	-1.54183	0.00004	0.64858	820.7203	538.0978
A_23_P13	NM_03228	Leucine zip	LDOC1L	-1.54191	2.47E-08	0.64854	2500.635	1615.254
A_23_P25	NM_02401	Homo sapi	MTRR,MSI	-1.54191	0.00018	0.64855	1210.172	763.5872
A_24_P19	NM_13039	Homo sapi	WRNIP1,W	-1.54223	0.00003	0.64841	971.4808	643.3154
A_23_P10	NM_00515	Homo sapi	USP10,UB	-1.54262	9.24E-12	0.64825	2614.681	1693.211
A_23_P11	NM_00019	Homo sapi	HPRT1,HP	-1.54266	1.15E-08	0.64823	13793.68	8991.263
A_23_P25	NM_01786	hypothetica	TMEM70,F	-1.54287	1.27E-19	0.64814	9518.428	6174.661
A_23_P21	NM_01230	Homo sapi	AP2A2,HIF	-1.54299	4.73E-11	0.64809	2092.29	1354.296
A_23_P30	NM_17812	Homo sapi	CXorf40A,I	-1.54299	2.62E-15	0.64809	3669.967	2378.261
A_32_P102252		Unknown	A_32_P10	-1.54303	1.69E-16	0.64808	5292.491	3431.384
A_23_P10	NM_02110	Homo sapi	NFS1,NIFS	-1.54311	4.10E-15	0.64804	3080.948	1997.179
A_23_P25	NM_00573	Homo sapi	KIF20A,MK	-1.54333	5.45E-15	0.64795	2144.798	1391.697
A_24_P24	AK025794	PI-3-kinase	SMG1	-1.54337	0.00001	0.64793	1469.985	939.8803
A_32_P97	NM_14569	Homo sapi	YWHAZ,K	-1.54345	2.62E-10	0.6479	5334.71	3438.131
A_23_P31	AF356877	Acetyl-Coe	ACAT2	-1.54361	3.26E-19	0.64783	6039.999	3916.049
A_23_P12	NM_00297	Homo sapi	SCP2,NLT	-1.54365	1.72E-13	0.64781	3707.402	2403.128
A_24_P91	BC004338	Uroporphyr	UROS	-1.54379	3.24E-09	0.64775	1985.553	1281.143
A_24_P23	NM_00468	eukaryotic	EIF1AY	-1.54393	0.00025	0.6477	1265.983	796.7861
A_24_P64	NM_00390	copine III	CPNE3,CP	-1.54455	7.28E-07	0.64744	867.6835	568.4063

A_23_P20	NM_02207	FLJ22794	FLJ22794,	-1.54458	5.49E-19	0.64743	5531.927	3584.513
A_23_P13	NM_00115	Homo sapi	SLC25A5,	-1.54523	5.70E-14	0.64715	92438.1	60006.46
A_24_P32	NM_00470	protein-kin	PRKRIR,D	-1.54539	8.84E-10	0.64709	1290.398	835.1714
A_23_P33	NM_00307	SWI/SNF r	SMARCE1	-1.5454	7.06E-06	0.64708	906.7907	580.3297
A_24_P31	NM_00515	ubiquitin sp	USP14,TG	-1.5454	4.52E-17	0.64708	3496.594	2263.451
A_24_P20	NM_01837	hypothetica	SYNJ2BP,	-1.54543	0.00037	0.64707	550.9468	356.5706
A_24_P23	NM_00686	RAB31, me	RAB31,Ral	-1.54545	4.30E-07	0.64706	962.5634	623.0161
A_23_P87	NM_00471	Homo sapi	SFRS2IP,S	-1.54546	2.44E-17	0.64705	2518.627	1629.718
A_32_P89	Z24727	Tropomyos	TPM1	-1.54572	2.01E-06	0.64695	2262.535	1456.759
A_23_P21	NM_14778	Homo sapi	CTSB,APP	-1.54637	5.87E-06	0.64668	5365.827	3482.776
A_23_P12	NM_00206	Homo sapi	GNAI1,Gi	-1.54638	0.00016	0.64667	598.8582	387.867
A_23_P41	NM_00132	C-terminal	CTBP1,MG	-1.54653	5.75E-09	0.64661	1469.476	942.9146
A_23_P16	NM_00638	protein with	CHERP,DA	-1.54684	1.97E-10	0.64648	6215.668	4012.125
A_23_P16	NM_01457	Homo sapi	BRD1,BRL	-1.54724	1.14E-11	0.64631	1929.143	1244.764
A_23_P35	NM_00381	Homo sapi	ADAM23,M	-1.54737	4.46E-10	0.64626	1952.814	1257.359
A_23_P48	NM_02405	Homo sapi	NUP37,p37	-1.54771	4.63E-14	0.64612	7924.905	5138.863
A_23_P14	NM_00590	Homo sapi	SMAD5,Dw	-1.54786	0.00003	0.64606	684.1728	441.7781
A_24_P32	NM_00620	phosphodie	PDE4A,PD	-1.54795	7.08E-09	0.64602	1353.884	878.6323
A_23_P42	NM_01898	Homo sapi	SSH1,SSH	-1.54799	0.00005	0.646	777.0599	497.1364
A_23_P16	NM_00334	Homo sapi	UBE2L3,E2	-1.548	7.09E-14	0.64599	4778.402	3086.77
A_32_P89	NM_00310	sorbitol de	SORD,SOI	-1.54806	5.22E-12	0.64597	3668.57	2370.612
A_23_P25	NM_00295	ribophorin	RPN1,RBP	-1.54838	9.44E-19	0.64584	21895.13	14114.42
A_23_P11	NM_01649	Homo sapi	TBC1D7,F	-1.54849	1.18E-14	0.64579	2938.227	1898.727
A_23_P13	NM_00264	Homo sapi	PIGF,MGC	-1.54856	7.89E-17	0.64576	3079.307	1988.74
A_23_P26	NM_01542	DKFZP564	C16orf51,D	-1.54862	1.62E-09	0.64573	1768.574	1145.912
A_23_P13	NM_00219	interferon r	IRF2,IRF-2	-1.54901	0.00007	0.64557	1027.124	646.1943
A_23_P24	NM_01646	hypothetica	HSPC196	-1.54904	4.02E-07	0.64556	1044.774	677.4667
A_23_P40	NM_00676	Homo sapi	MYST3,MO	-1.54921	3.25E-08	0.64549	1604.008	1040.403
A_23_P20	NM_00090	diaphorase	NQO1,DTD	-1.54943	4.36E-09	0.6454	5555.032	3599.773
A_23_P15	NM_13333	Homo sapi	RAD17,CC	-1.54945	6.66E-11	0.64539	2362.548	1522.8
A_23_P13	NM_01508	Homo sapi	MRPS27,M	-1.54952	4.21E-07	0.64536	3820.785	2462.232
A_23_P25	NM_00577	Homo sapi	LHFPL2,KI	-1.54957	6.32E-10	0.64534	2305.123	1482.776
A_23_P20	NM_00320	Homo sapi	TFAM,TCF	-1.54978	2.71E-11	0.64525	8127.006	5266.456
A_23_P21	NM_00111	Homo sapi	ADD1,ADD	-1.54994	7.93E-09	0.64519	7405.016	4795.14
A_23_P43	AY037156	Hypothetic	SB145	-1.55026	0.00004	0.64505	869.5125	551.2459
A_24_P11	NM_03204	hypothetica	DKFZP564	-1.55048	1.45E-07	0.64496	1332.839	867.3608
A_23_P72	NM_01792	nuclear pro	NPL4,FLJ2	-1.55081	1.63E-10	0.64483	2422.793	1565.151
A_32_P18	NM_00675	transaldola	TALDO1,T	-1.55086	1.69E-08	0.6448	94041.1	61060.73
A_24_P27	NM_00737	TAR DNA	TARDBP,T	-1.55103	6.51E-18	0.64473	2940.175	1895.494
A_23_P21	NM_00036	Homo sapi	TPMT	-1.55119	0.00001	0.64467	1118.29	711.0254
A_32_P84	AK096674	Chromosor	C14orf32	-1.55133	4.20E-11	0.64461	1534.695	990.2373
A_23_P10	NM_01207	Homo sapi	C16orf35,C	-1.55138	8.15E-12	0.64459	8718.13	5637.843
A_23_P35	NM_01543	Homo sapi	RCHY1,AR	-1.55164	9.18E-17	0.64448	4601.73	2967.558
A_23_P86	NM_13333	Fer-1-like 3	FER1L3	-1.55166	1.06E-16	0.64447	7087.176	4574.944
A_24_P30	NM_07846	Homo sapi	BCCIP,TOI	-1.55188	1.13E-07	0.64438	6858.134	4444.048
A_24_P69	NM_00429	ring finger	RNF14,AR	-1.55205	0.00003	0.64431	1237.409	781.6311
A_23_P31	NM_00385	cyclin K	CCNK,CPI	-1.55221	2.08E-07	0.64424	1177.818	753.6406
A_23_P36	NM_01476	Homo sapi	SCRN1,SE	-1.5524	2.18E-06	0.64416	784.2065	508.3731
A_32_P27	AB033062	Kinesin fan	KIF26A	-1.55247	1.72E-07	0.64413	1916.778	1226.059
A_23_P95	NM_03229	hypothetica	SLC37A3,M	-1.55256	4.60E-09	0.6441	2066.817	1325.109
A_23_P94	NM_01417	Homo sapi	MRPL15,R	-1.5528	1.60E-25	0.644	11322.63	7291.802
A_23_P12	NM_02023	Homo sapi	BBX,MDSC	-1.55331	1.30E-19	0.64379	5431.777	3499.712

A_24_P587882		Unknown	A_24_P58	-1.55337	5.58E-07	0.64376	6043.074	3931.419
A_23_P86	NM_00382	Homo sapi	FADD,GIG	-1.5534	8.97E-10	0.64375	8613.429	5557.271
A_23_P10	NM_02275	sarcoma a	C1orf149,F	-1.5534	6.31E-08	0.64375	2687.489	1724.724
A_23_P14	NM_01220	Golgi appa	GLG1,CFR	-1.5534	3.06E-08	0.64375	1765.319	1145.084
A_23_P43	NM_00156	Homo sapi	EIF3S6,IN	-1.55346	1.59E-24	0.64372	44050.48	28367.78
A_23_P91	NM_00369	Homo sapi	PRKRA,RA	-1.55351	3.86E-09	0.6437	2487.744	1593.902
A_23_P14	NM_03079	Homo sapi	FBXO38,M	-1.55369	0.0001	0.64363	875.6884	552.7886
A_32_P24	AL832427	Fanconi an	FANCD2	-1.5537	0.00012	0.64363	4047.591	2617.982
A_24_P93	NM_00717	protease, s	PRSS23,S	-1.55377	2.96E-10	0.6436	4497.687	2903.9
A_23_P70	NM_00353	Homo sapi	HIST1H3E	-1.5538	2.78E-14	0.64358	22167.5	14288.56
A_23_P37	NM_01822	Homo sapi	C14orf108,	-1.55389	1.18E-06	0.64355	1324.416	843.8801
A_24_P48	NM_00300	secreted a	SECTM1,K	-1.55397	0.0001	0.64351	1013.144	635.9418
A_23_P28	NM_00462	Homo sapi	TSN,TRSL	-1.55409	3.77E-09	0.64346	2427.177	1557.43
A_23_P63	NM_18174	Homo sapi	LASS2,L3,	-1.55427	4.71E-07	0.64339	5175.104	3338.281
A_24_P18	NM_00271	protein pho	PPP2CA,R	-1.55434	5.35E-18	0.64336	5341.071	3441.507
A_23_P21	NM_02475	Homo sapi	EHMT1,GL	-1.55435	4.70E-09	0.64336	1500.973	971.1648
A_23_P13	NM_03231	Homo sapi	C2orf7,PAI	-1.55453	3.98E-09	0.64328	2056.513	1327.464
A_23_P56	NM_02477	Homo sapi	GEMIN6,F	-1.55499	2.66E-08	0.64309	15792.18	10205.09
A_23_P26	NM_01415	Homo sapi	ZC3H7A,Z	-1.55573	2.91E-17	0.64278	3252.726	2089.975
A_32_P29	NM_15364	Homo sapi	TPM3,TRK	-1.55573	4.62E-20	0.64279	30865.61	19869.11
A_24_P14	NM_02107	N-myristoy	NMT1,NMT	-1.55589	5.83E-07	0.64272	8057.818	5206.75
A_23_P21	NM_00471	cytochrome	COX7A2L,	-1.55592	4.55E-19	0.64271	21767.5	14019.8
A_23_P77	NM_00310	Homo sapi	SORD,SOI	-1.55619	1.41E-09	0.64259	4753.614	3063.64
A_23_P20	NM_00357	Homo sapi	VAPA,VAP	-1.5562	5.02E-11	0.64259	2090.229	1347.083
A_23_P96	NM_01773	O-linked m	POMGNT1	-1.5562	1.96E-06	0.64259	879.6843	569.2593
A_23_P40	NM_00687	Homo sapi	DSTN,ADF	-1.55629	3.47E-12	0.64255	24244.89	15540.17
A_32_P69	AK023633	Leucyl/cyst	LNPEP	-1.5563	3.94E-07	0.64255	1372.243	891.4341
A_23_P16	NM_01528	Homo sapi	WDR7,TRA	-1.55638	0.00002	0.64252	1148.953	752.6882
A_32_P14	NM_00693	SMT3 (sup	SUMO2,HS	-1.55647	1.70E-08	0.64248	24986.59	16119.44
A_23_P21	NM_03238	Homo sapi	GFM2,EFG	-1.55651	1.82E-09	0.64246	6749.74	4348.298
A_23_P14	NM_02040	Homo sapi	NUP107,N	-1.55663	1.02E-08	0.64241	12909.72	8322.846
A_23_P21	NM_02211	cutaneous	C13orf10,F	-1.55673	5.91E-10	0.64237	1238.772	795.8602
A_24_P19	NM_19948	Homo sapi	C20orf24,F	-1.55677	2.54E-13	0.64236	56966.21	36733.57
A_23_P12	NM_02469	Homo sapi	ECHDC3,F	-1.55681	0.00067	0.64234	4708.466	3030.699
A_23_P14	NM_00259	programme	PDCD2,RF	-1.5569	1.78E-11	0.6423	2466.09	1582.528
A_23_P20	NM_00238	Homo sapi	MCP,TLX,	-1.557	3.47E-07	0.64226	879.8273	563.6451
A_23_P14	NM_01742	Homo sapi	GALNT7,G	-1.55702	2.02E-11	0.64225	2307.869	1486.458
A_23_P11	NM_03235	Homo sapi	MRPL45,M	-1.55731	3.89E-13	0.64213	15725.88	10149.65
A_23_P40	NM_00409	E2F transc	E2F2,E2F-	-1.55739	1.12E-17	0.6421	3403.558	2185.045
A_24_P17	AL133117	THO comp	THOC2	-1.55743	0.0001	0.64208	3000.446	1920.581
A_24_P14	AY007122	WAS prote	WASF2	-1.55746	2.32E-12	0.64207	6687.774	4308.197
A_23_P25	NM_03146	Homo sapi	CD99L2,M	-1.55748	1.73E-18	0.64206	19398.47	12484.97
A_23_P13	NM_03131	carboxype	CPVL,HVL	-1.55755	0.00001	0.64203	1136.175	743.478
A_23_P39	Z24727	Tropomyos	TPM1	-1.5578	2.68E-10	0.64193	4384.723	2821.519
A_23_P32	NM_03316	Homo sapi	SURF4,ER	-1.55816	7.90E-21	0.64178	4811.982	3089.46
A_23_P15	NM_00600	Homo sapi	TUBA1,FL	-1.55854	6.31E-20	0.64163	133158.7	85292.84
A_24_P22	NM_02234	protein kin	NJMU-R1	-1.55858	4.98E-08	0.64161	3154.035	2021.66
A_24_P13	NM_00636	dendritic ce	hfl-B5,GA1	-1.5592	4.03E-17	0.64136	42629.23	27411.14
A_23_P25	NM_08065	similar to R	MGC13017	-1.55923	6.44E-15	0.64134	3422.617	2192.939
A_24_P94	NM_00470	rabaptin-5	RABEP1,R	-1.55936	0.0009	0.64129	718.3479	444.6888
A_23_P20	NM_02513	melanoma/	ARMC9,AF	-1.55938	3.06E-09	0.64128	1129.374	726.0786
A_24_P84	NM_00331	tetratricope	TTC3,DCR	-1.55949	7.49E-09	0.64124	2072.696	1334.23

A_23_P17	BC059410	Hypothetic	LOC28514	-1.55961	1.06E-24	0.64119	9364.528	6004.688
A_24_P246963		Unknown	ENST0000	-1.5597	0.00001	0.64115	902.163	586.4298
A_23_P29	NM_01404	Signal pep	SPCS1	-1.55973	1.83E-12	0.64114	16213.81	10450.88
A_24_P81	NM_00675	transaldola	TALDO1,T	-1.55991	9.34E-14	0.64106	63441.79	40472.98
A_23_P16	NM_00599	Homo sapi	CCT3,CCT	-1.55995	2.99E-09	0.64105	91185.75	58000.05
A_23_P59	NM_00734	Homo sapi	PAXIP1,PT	-1.55997	2.81E-15	0.64104	3394.417	2175.022
A_32_P53338		Unknown	A_32_P53	-1.56004	4.36E-07	0.64101	910.7087	586.9279
A_23_P84	NM_00094	Homo sapi	PPIC,CYP	-1.56011	2.03E-07	0.64098	5704.247	3675.467
A_23_P25	NM_01622	Homo sapi	DCTN4	-1.56018	2.43E-06	0.64095	3044.614	1940.355
A_24_P25	NM_00128	adaptor-rel	AP2B1,AD	-1.56021	0.00073	0.64094	596.1284	391.2424
A_24_P87	AK054998	Pleckstrin	PLEKHB2	-1.56023	8.46E-16	0.64093	7589.194	4872.431
A_24_P22	NM_00460	Sjogren sy	TROVE2,S	-1.56025	1.60E-09	0.64092	5791.764	3730.303
A_23_P25	NM_14891	Homo sapi	PSMB8,LM	-1.56063	0.00077	0.64077	1985.79	1285.617
A_23_P10	D13627	Chaperonir	CCT8	-1.56105	3.34E-17	0.6406	17955.94	11528.49
A_32_P170444		Unknown	THC18467	-1.56122	1.14E-17	0.64053	3398.099	2177.294
A_24_P28	NM_03075	makorin, ri	MKRN4	-1.56131	0.00033	0.64049	595.8608	376.7524
A_23_P13	NM_00177	Homo sapi	CD58,LFA	-1.56157	9.17E-13	0.64038	2253.519	1445.653
A_23_P36	NM_00314	structure s	SSRP1,FA	-1.56201	3.94E-11	0.6402	21446.43	13808.54
A_23_P19	NM_17814	Solute carr	SLC35B2	-1.56202	1.63E-16	0.6402	5301.361	3399.188
A_24_P711050		Unknown	A_24_P71	-1.56211	1.40E-09	0.64016	3183.275	2034.674
A_23_P16	NM_02491	Homo sapi	C20orf172,	-1.56222	1.92E-11	0.64011	2470.628	1580.671
A_23_P13	NM_01767	Homo sapi	C20orf42,U	-1.56222	1.28E-10	0.64012	4675.358	2993.962
A_24_P22	NM_01232	microtubul	MAPRE1,E	-1.56234	2.43E-13	0.64006	4535.42	2905.652
A_23_P12	NM_00106	Topoisome	TOP2B	-1.56242	4.57E-10	0.64003	17207.52	11019.46
A_23_P15	NM_01505	MYC bindi	MYCBP2	-1.56243	6.14E-14	0.64003	2728.146	1747.924
A_24_P39	AK098354	BS 3076	AK098354	-1.56273	1.55E-06	0.63991	977.7783	631.4777
A_23_P21	NM_00229	Homo sapi	LAMB2,LA	-1.56276	1.62E-23	0.6399	4472.53	2861.66
A_23_P13	NM_00178	Homo sapi	CDC2,CDK	-1.56283	2.16E-09	0.63987	13026.17	8365.813
A_23_P25	NM_01234	Homo sapi	FBXO9,FB	-1.56316	1.92E-11	0.63973	4905.801	3126.88
A_32_P19	NM_00213	RNA bindir	RBMX,RNI	-1.5634	1.09E-12	0.63963	3975.17	2544.823
A_23_P11	NM_00477	CCR4-NOT	CNOT8,CA	-1.56364	4.89E-14	0.63953	2574.196	1648.935
A_23_P89	NM_00576	tripartite m	TRIM28,KA	-1.56368	4.19E-21	0.63952	6470.674	4135.196
A_23_P14	NM_00579	M-phase pl	MPHOSPH	-1.56385	0.00021	0.63945	947.0643	588.0295
A_23_P13	NM_01466	enthoprotin	ENTH,EPN	-1.56401	5.89E-13	0.63938	4346.243	2784.622
A_23_P14	BC030119	Chromosor	C14orf109	-1.56414	9.92E-10	0.63933	1964.326	1259.672
A_23_P18	NM_03280	glycosyltra	AGO61,FL	-1.56424	9.67E-26	0.63929	9404.749	6012.529
A_24_P247616		Unknown	A_24_P24	-1.5645	0.00001	0.63918	1576.901	994.5824
A_23_P16	NM_00367	Homo sapi	DENR,DRF	-1.56469	0.00004	0.6391	4671.026	2987.887
A_23_P12	NM_02301	Homo sapi	C1orf60,FL	-1.56499	1.10E-06	0.63898	1232.595	776.3421
A_23_P14	NM_13900	Homo sapi	HFE,HH,HI	-1.56504	0.00025	0.63896	651.3822	409.5043
A_23_P15	NM_00227	Homo sapi	TNPO1,MI	-1.56512	1.23E-12	0.63893	4144.81	2638.82
A_23_P25	NM_00451	Homo sapi	ILF2,NF45	-1.56537	6.43E-09	0.63883	44790.93	28615.26
A_23_P18	NM_01832	phosphatid	PI4K2B,PI	-1.56577	8.41E-07	0.63866	1186.204	748.3002
A_23_P20	NM_01768	Homo sapi	VPS13C	-1.56592	7.98E-07	0.6386	1460.027	922.464
A_23_P15	NM_01814	hypothetica	FAM82C,p	-1.56593	4.64E-14	0.6386	6572.944	4197.075
A_23_P16	NM_00688	calmodulin	CALM1,CA	-1.56602	2.26E-07	0.63856	11339.46	7295.105
A_32_P20	AL713796	Hypothetic	DKFZp667	-1.56605	0.00002	0.63855	1202.357	752.5748
A_24_P936911		Unknown	A_24_P93	-1.56609	0.00081	0.63853	659.6336	408.3313
A_23_P13	NM_14456	Homo sapi	RPIA,RPI	-1.56644	5.32E-15	0.63839	7414.087	4739.359
A_24_P409750		Unknown	A_24_P40	-1.56708	5.72E-07	0.63813	1159.064	733.5352
A_24_P22	BC064492	Ubiquitin-c	UBE2I	-1.56723	1.79E-08	0.63807	1106.789	703.9337
A_32_P93	NM_13836	Homo sapi	FAM44B	-1.56725	3.73E-11	0.63806	4472.989	2864.208

A_24_P118813		Unknown	A_24_P11	-1.56788	0.00009	0.6378	2102.097	1351.167
A_23_P11	NM_01604	Homo sapi	DERL2,FL	-1.56841	2.33E-21	0.63759	3985.649	2540.954
A_24_P26	NM_01447	dimethylad	HSA9761	-1.56844	4.31E-14	0.63758	2035.821	1298.535
A_23_P55	NM_00442	Homo sapi	DVL2	-1.56858	4.14E-06	0.63752	1413.425	887.1761
A_23_P20	NM_00237	mago-nash	MAGOH	-1.56864	5.68E-07	0.6375	3938.047	2479.953
A_24_P32	AK023593	Peroxisom	PEX3	-1.56874	3.16E-08	0.63745	1965.581	1246.171
A_23_P10	NM_02109	Homo sapi	SLC5A6,SI	-1.56874	5.99E-07	0.63746	966.5918	611.2431
A_24_P34	AB082526	NIMA (nev	NEK9	-1.56877	3.14E-18	0.63744	3826.408	2438.875
A_24_P820302		Unknown	A_24_P82	-1.56907	1.17E-09	0.63732	1590.519	1008.169
A_23_P89	NM_01521	Homo sapi	EXOC7,EX	-1.56908	7.96E-19	0.63732	4118.855	2622.592
A_32_P19	NM_03115	heterogene	HNRPA1,H	-1.56911	4.10E-19	0.6373	113390.8	72407.62
A_24_P28619		AF010144	THC19528	-1.56913	3.46E-08	0.63729	1498.837	948.5158
A_24_P29	NM_01836	hypothetica	AGPAT5,L	-1.56923	1.21E-07	0.63725	1147.217	736.449
A_32_P159176		Unknown	THC19964	-1.56937	0.00001	0.6372	731.6121	464.3329
A_23_P77	NM_00466	RAB11A, n	RAB11A,Y	-1.5697	0.00019	0.63706	643.9681	417.4119
A_23_P12	NM_00512	Homo sapi	NUP153,N	-1.56974	8.43E-21	0.63705	2946.785	1877.512
A_23_P16	NM_00299	Homo sapi	SDC1,SDC	-1.56978	3.09E-12	0.63703	8004.234	5104.21
A_24_P18	NM_01542	aminoadip	AASDHPP	-1.56979	5.10E-08	0.63703	2450.198	1558.052
A_32_P17	BC017675	BTB (POZ)	HSPC063	-1.5698	8.19E-07	0.63702	975.9026	618.5398
A_24_P40	NM_00208	growth fact	GRB2,ASH	-1.56987	0.00002	0.637	1124.72	738.1779
A_24_P15	NM_00135	aldo-keto r	AKR1C1,C	-1.5699	9.06E-09	0.63698	11424.23	7202.571
A_24_P42	NM_00280	proteasom	PSMD2,S2	-1.57043	5.18E-06	0.63677	3382.009	2138.029
A_32_P13	BC013077	Membrane	MPP6	-1.57055	2.66E-06	0.63672	1362.55	853.1928
A_24_P11	NM_01553	Glutamate	GRINL1A	-1.57072	0.00009	0.63665	932.8709	579.4549
A_24_P32	AB014733	Golgi mem	SMAP-5	-1.57101	0.00014	0.63653	879.0531	544.9938
A_24_P204204		Unknown	A_24_P20	-1.57111	0.00006	0.63649	1298.175	847.7175
A_23_P39	NM_01646	Homo sapi	CXXC5,HS	-1.57115	1.56E-24	0.63647	7617.749	4847.317
A_23_P46	NM_15237	hypothetica	C1orf131,D	-1.57119	3.54E-08	0.63646	4069.149	2594.449
A_24_P65	AY007106	Keratin ass	KRTAP4-7	-1.5714	4.56E-10	0.63637	2106.335	1333.85
A_23_P15	NM_00541	Homo sapi	SHMT2,GL	-1.57159	3.00E-20	0.6363	7241.359	4615.066
A_23_P21	NM_00737	Homo sapi	BRD3,ORF	-1.57174	4.11E-06	0.63624	1972.254	1265.437
A_32_P44	NM_03076	nucleosom	NSBP1	-1.57205	0.0007	0.63611	990.2073	604.8436
A_24_P45	NM_03115	heterogene	HNRPA1,H	-1.57217	3.60E-27	0.63606	70508.45	44845.05
A_23_P12	NM_01813	hypothetica	FLJ10560	-1.57228	0.00048	0.63602	534.5659	336.4124
A_32_P22	NM_13928	Homo sapi	CDC26,C9	-1.57239	9.22E-15	0.63598	9046.883	5772.202
A_32_P14	AL530645	Prenylcyste	PCYOX1	-1.57257	0.00018	0.6359	1172.272	766.6386
A_24_P19	NM_15230	Homo sapi	DDX52,RC	-1.57273	6.80E-12	0.63584	4068.862	2576.528
A_23_P42	NM_03231	hypothetica	YIPF4,FinC	-1.57296	0.00001	0.63574	1328.702	830.9149
A_23_P15	NM_01662	archaemet	AMZ2	-1.5734	1.36E-15	0.63556	8704.846	5548.417
A_23_P50	NM_00610	Homo sapi	KNTC2,HE	-1.57387	1.61E-17	0.63538	3901.213	2479.946
A_24_P22	NM_00103	ribonucleot	RRM2,R2,I	-1.57403	1.30E-13	0.63531	4664.382	2963.089
A_32_P14	NM_03131	heterogene	HNRPC,C1	-1.57415	1.66E-18	0.63526	20899.14	13252.39
A_23_P16	XM_37482	Homo sapi	PDCD11	-1.57417	8.33E-12	0.63526	9523.623	6069.475
A_24_P34	NM_17686	Homo sapi	PSME3,Ki,	-1.57419	2.61E-11	0.63525	1387.619	878.9987
A_23_P10	NM_00340	Homo sapi	YWHAH,Y	-1.57421	4.98E-14	0.63524	2078.039	1321.163
A_24_P76	AK024037	Homo sapi	AK024037	-1.57448	0.00018	0.63513	4850.369	3076.169
A_24_P891265		Unknown	A_24_P89	-1.57458	7.14E-15	0.63509	5549.211	3528.231
A_23_P50	NM_13916	Homo sapi	SMCR7,MC	-1.57464	0.00017	0.63506	697.9208	453.5658
A_23_P12	NM_02272	Homo sapi	FANCF,FA	-1.57466	4.85E-06	0.63506	777.7362	496.4959
A_24_P610945		Unknown	ENST0000	-1.57478	3.34E-09	0.63501	2076.369	1324.309
A_23_P14	NM_00333	Homo sapi	UBE2A,UB	-1.57494	3.04E-08	0.63495	13607.97	8686.341
A_24_P24	NM_01828	hypothetica	NUDT15,M	-1.57509	5.34E-06	0.63489	806.895	514.2732

A_23_P37	NM_17356	Homo sapi	C6orf69,M	-1.57537	1.46E-07	0.63477	859.6947	545.1571
A_23_P30	NM_02402	hypothetica	MGC3265	-1.57565	6.17E-11	0.63466	1814.139	1156.272
A_24_P40	NM_00330	Homo sapi	TRIP6,OIP	-1.57573	3.18E-16	0.63463	12824.24	8165.516
A_23_P80	NM_00680	Homo sapi	BTG3,ANA	-1.57617	1.44E-09	0.63445	5586.731	3565.089
A_32_P85813		Unknown	A_32_P85	-1.57618	0.00046	0.63444	897.9082	548.683
A_32_P11	NM_01553	Glutamate	GRINL1A	-1.57619	0.00001	0.63444	1194.727	743.2325
A_24_P15	NM_00736	non-POU-c	NONO,P54	-1.57642	6.87E-16	0.63435	12846.51	8169.31
A_24_P13	NM_00694	stress 70 p	STCH	-1.57672	2.46E-08	0.63423	1475.82	942.1967
A_24_P41	NM_17202	Homo sapi	POM121,M	-1.57686	1.69E-12	0.63417	1655.889	1050.817
A_23_P74	NM_01775	Homo sapi	NSUN2,TR	-1.57698	1.04E-15	0.63412	11529.43	7302.563
A_23_P43	NM_01594	CGI-12 pro	MTERFD1,	-1.577	8.59E-13	0.63411	2245.544	1420.209
A_23_P20	NM_01961	WDR45-lik	WDR45L	-1.57709	2.40E-08	0.63408	11492.87	7307.703
A_23_P29	AL096748	Armadillo r	ARMC8	-1.57735	0.00062	0.63397	677.1993	415.7861
A_32_P17635		JC7219 nu	THC19529	-1.57735	6.28E-07	0.63398	1371.342	879.5752
A_23_P37	NM_00656	cell growth	CGRRF1,C	-1.57754	1.78E-09	0.6339	1384.392	874.4219
A_32_P80	NM_03115	heterogene	HNRPA1,H	-1.57796	1.21E-27	0.63373	141122	89436.82
A_23_P20	NM_17798	Homo sapi	PPM1G,PF	-1.57807	1.01E-19	0.63369	4560.687	2886.686
A_23_P98	NM_00031	Homo sapi	PTEN,BZS	-1.57818	6.07E-12	0.63364	1480.277	938.1661
A_23_P35	NM_13333	Fer-1-like 3	FER1L3	-1.57827	1.36E-23	0.6336	6947.4	4404.355
A_23_P16	NM_01560	lamina-ass	TOR1AIP1	-1.57841	1.37E-06	0.63355	845.339	537.9214
A_23_P15	NM_00640	Homo sapi	TM9SF1,M	-1.57842	7.72E-09	0.63354	2157.658	1373.461
A_32_P37	AK056809	CDNA FLJ	LOC11606	-1.57849	8.75E-14	0.63352	2351.401	1492.673
A_23_P40	D86971	KIAA0217	KIAA0217	-1.57878	2.85E-14	0.6334	3580.703	2268.446
A_23_P12	BC030623	Homo sapi	BC030623	-1.57883	7.11E-21	0.63338	1313.949	833.0765
A_23_P25	NM_00573	Homo sapi	RAD50,hR	-1.57889	4.52E-24	0.63336	2161.227	1372.864
A_24_P31	NM_14450	Homo sapi	F11R,JAM	-1.57922	1.86E-12	0.63322	5037.461	3189.89
A_23_P60	NM_01337	Homo sapi	PDCD6IP,A	-1.57942	6.37E-10	0.63315	8193.574	5196.442
A_23_P84	NM_02208	Homo sapi	GALNT11,	-1.57958	2.20E-14	0.63308	4744.028	3004.594
A_24_P669216		Unknown	THC18199	-1.57973	5.48E-07	0.63302	1032.665	663.921
A_23_P21	NM_13083	Homo sapi	OPA1,NPG	-1.57981	4.05E-07	0.63299	2110.364	1349.545
A_32_P71	NM_00456	phosphatid	PIGH,GPI-	-1.57995	6.31E-08	0.63293	1074.828	678.0225
A_23_P77	NM_02247	Homo sapi	ZFP106,SH	-1.57996	4.38E-16	0.63293	3473.542	2200.074
A_23_P12	NM_01416	Homo sapi	MED4,VDF	-1.58004	5.26E-08	0.6329	7604.732	4849.854
A_24_P22	NM_01549	DKFZP434	KIAA1429,	-1.58029	4.93E-14	0.63279	4813.741	3041.609
A_24_P30	AK000126	GTPase ac	DKFZP434	-1.58033	0.00057	0.63278	562.2138	360.1216
A_23_P41	NM_01517	Homo sapi	BAT2D1,X	-1.58063	3.02E-20	0.63266	4797.74	3033.21
A_24_P17	NM_19428	DnaJ homo	DNAJA5	-1.58064	0.00003	0.63266	1084.353	672.3994
A_23_P31	NM_00038	Homo sapi	BLMH,BH,	-1.58086	7.03E-13	0.63257	4600.393	2904.738
A_23_P15	NM_02214	leucine zip	FKSG14,P	-1.58106	7.87E-12	0.63249	2533.76	1601.436
A_24_P11	AK022628	Forkhead b	KIAA0415	-1.58113	4.48E-08	0.63246	2049.088	1287.856
A_23_P30	NM_00419	Homo sapi	P4HA2,4-P	-1.5813	3.66E-09	0.63239	4039.919	2554.913
A_23_P10	NM_01909	pleckstrin h	PLEKHA3,	-1.58159	8.02E-07	0.63228	1647.017	1029.893
A_23_P69	NM_01802	Homo sapi	YEATS2,F	-1.58205	6.58E-10	0.63209	3304.504	2090.612
A_23_P66	NM_00261	3-phosphoi	PDPK1	-1.58211	2.03E-09	0.63207	1643.663	1034.428
A_23_P25	NM_01234	F-box only	FBXO9,FB	-1.58263	1.26E-17	0.63186	8355.499	5275.157
A_24_P170103		Unknown	A_24_P17	-1.58272	6.00E-19	0.63182	96962.41	61363.63
A_24_P14	AK130583	Poly(rC) bi	PCBP2	-1.58288	4.71E-12	0.63176	3088.198	1950.505
A_23_P70	U83115	Absent in n	AIM1	-1.58291	1.14E-13	0.63175	1977.137	1247.224
A_23_P12	NM_02453	hypothetica	FLJ12118,	-1.58293	5.11E-26	0.63174	19227.54	12150.66
A_23_P25	NM_00028	Homo sapi	PDHA1,PD	-1.58322	2.69E-18	0.63162	25500.79	16076.93
A_24_P54	AK026687	Signal seq	SSR1	-1.58332	3.61E-06	0.63158	1238.933	770.0815
A_32_P19	NM_01599	Kruppel-lik	KLF13,BTE	-1.58333	4.20E-19	0.63158	4380.804	2767.336

A_23_P12	NM_03270	tubulin alpl	TUBA6,MG	-1.58335	8.36E-13	0.63157	132124.6	83009.05
A_23_P72	NM_00422	Homo sapi	CRSP2,CS	-1.58342	9.66E-07	0.63154	1017.187	636.1835
A_23_P57	NM_01609	Homo sapi	EIF3S6IP,E	-1.58345	1.08E-11	0.63153	1946.962	1226.584
A_23_P10	NM_08066	Homo sapi	CDCA5,MC	-1.5835	2.23E-08	0.63151	5183.535	3273.254
A_24_P35	BX648218	Additional s	ASXL2	-1.58357	0.00005	0.63148	768.0509	494.639
A_23_P24	AK000367	Chromosor	C10orf18	-1.58366	0.00002	0.63145	2606.746	1633.217
A_23_P74	NM_01338	Homo sapi	SLC25A24	-1.58376	1.41E-07	0.63141	4385.436	2787.171
A_23_P59	NM_01635	solute carri	SLCO4A1,	-1.58393	1.91E-09	0.63134	2179.308	1373.936
A_24_P19	NM_00165	ADP-ribosy	ARF1	-1.58393	1.50E-13	0.63134	50043.97	31728.38
A_23_P54	NM_01610	comparativ	CGI-37,KD	-1.58402	1.08E-14	0.63131	5125.113	3241.293
A_32_P27327		Unknown	A_32_P27	-1.58424	1.76E-08	0.63122	6358.712	4025.021
A_23_P16	NM_00735	low density	COG2,LDL	-1.58465	9.11E-11	0.63105	4822.444	3047.852
A_32_P12	NM_17351	hypothetica	FLJ90709,	-1.5847	3.63E-08	0.63104	1232.407	783.0922
A_24_P192840		ALU4_HUM	THC19457	-1.58477	0.00022	0.63101	580.2195	369.4891
A_23_P23	NM_00252	Homo sapi	NRD1,hNR	-1.58508	3.64E-17	0.63088	5969.428	3768.92
A_23_P38	NM_15230	Homo sapi	C20orf158,	-1.58512	5.57E-18	0.63087	3582.267	2260.494
A_23_P54	NM_03287	Homo sapi	JUB,Ajuba,	-1.58523	2.67E-07	0.63082	2337.056	1461.962
A_23_P15	NM_01434	Homo sapi	FJX1,FLJ2	-1.58552	9.20E-08	0.63071	2065.301	1289.574
A_23_P77	NM_03280	signal pept	SPPL2A,IM	-1.58553	7.36E-16	0.6307	3731.152	2359.405
A_23_P13	NM_00395	Homo sapi	SLC25A14	-1.58557	3.81E-09	0.63069	2580.051	1630.304
A_23_P21	NM_02492	Homo sapi	NUP210,G	-1.5856	1.08E-12	0.63068	1409.033	888.9475
A_24_P43	NM_01799	hypothetica	C9orf40,FL	-1.58575	2.12E-09	0.63062	1977.605	1241.402
A_23_P31	NM_15267	Homo sapi	SLC10A4,M	-1.58577	0.00007	0.63061	935.2665	573.0195
A_23_P59	NM_18172	hypothetica	FLJ12760	-1.58578	0.0006	0.63061	1260.439	821.1779
A_23_P14	NM_00275	Homo sapi	EIF2AK2,P	-1.58586	3.39E-18	0.63057	3740.458	2358.389
A_32_P23	NM_17308	Homo sapi	LIN9,TGS,	-1.58604	1.94E-22	0.6305	4711.141	2970.685
A_24_P12	NM_01466	enthoprotin	ENTH,EPN	-1.58606	1.58E-14	0.63049	2665.49	1674.33
A_23_P15	NM_01563	DKFZP586	EIF2B4,EIF	-1.58609	0.00007	0.63048	1011.428	654.4723
A_24_P97	NM_00091	peptidylgly	PAM,PAL,P	-1.58627	9.25E-11	0.63041	1325.762	834.3864
A_23_P16	AB018353	Unc-84 hor	UNC84A	-1.58666	6.05E-07	0.63025	1708.396	1061.214
A_23_P39	NM_01570	Homo sapi	C2orf30,CL	-1.58669	1.60E-08	0.63024	2062.041	1304.289
A_23_P12	NM_00363	Homo sapi	NIPSNAP1	-1.5867	0.00002	0.63024	901.6763	560.7424
A_23_P34	NM_00538	Homo sapi	PDCL,PhLP	-1.5869	0.00071	0.63016	1209.878	732.8532
A_23_P25	NM_00294	ribosomal p	RPL15,EC4	-1.58736	1.47E-13	0.62998	3451.554	2173.441
A_23_P35	NM_01564	PAI-1 mRN	SERBP1,C	-1.58747	1.08E-10	0.62993	30058.83	19053.09
A_24_P70	AL832246	Heterogene	HNRPA3	-1.5875	8.55E-14	0.62992	2753.153	1733.903
A_23_P90	NM_02090	PAN2 prote	RDH14,PA	-1.58758	1.96E-06	0.62989	3133.418	1963.453
A_23_P41	NM_01220	Homo sapi	HARSL,HC	-1.58764	1.58E-08	0.62986	2038.235	1288.1
A_32_P19	AK097068	Hypothetica	AK097068	-1.58792	0.00002	0.62975	817.8113	509.3635
A_23_P83	NM_00204	Homo sapi	GAS1	-1.58795	2.27E-08	0.62974	2482.995	1573.026
A_23_P78	NM_03283	hypothetica	FLJ14775	-1.58802	5.32E-15	0.62971	4526.396	2853.63
A_23_P34	NM_15325	Zinc finger,	FLJ25952	-1.58804	4.04E-11	0.62971	6397.579	4039.961
A_23_P25	NM_00347	DEK oncog	DEK	-1.58806	4.70E-09	0.6297	7302.348	4624.493
A_23_P28	NM_03190	mitochondr	MRPS5,MR	-1.58807	1.32E-10	0.6297	2420.487	1524.31
A_23_P11	NM_00153	Homo sapi	HNRPL,hn	-1.58841	1.90E-09	0.62956	4229.995	2648.785
A_24_P24	NM_00637	vesicle-ass	VTI1B,VTI1	-1.58899	6.02E-11	0.62933	1622.813	1023.498
A_23_P12	NM_02514	Homo sapi	MAK3,MAK	-1.58902	2.13E-06	0.62932	5704.943	3611.145
A_23_P32	NM_01532	Est1p-like p	EST1B,SM	-1.58911	6.60E-09	0.62928	2362.644	1490.671
A_23_P58	NM_00190	Homo sapi	CTNNA1,C	-1.58917	8.84E-09	0.62926	8703.675	5500.46
A_24_P22	NM_17743	Homo sapi	DICER1,Di	-1.58929	9.69E-09	0.62921	1207.901	765.3178
A_23_P17	NM_00279	Homo sapi	PSMB2,HC	-1.58933	2.71E-20	0.6292	6015.225	3786.134
A_23_P21	NM_03353	Homo sapi	FBXL5,FBL	-1.58933	3.91E-06	0.6292	1768.187	1097.617

A_24_P18	NM_02468	hypothetica	FLJ23560	-1.58935	1.56E-08	0.62919	950.2549	598.7898
A_23_P25	NM_01602	CGI-82 pro	RDH11,MD	-1.58947	6.03E-18	0.62914	4661.876	2936.685
A_32_P23	AY007155	Hypothetic	LOC43994	-1.58958	1.01E-08	0.6291	1153.521	722.5331
A_23_P11	NM_00409	Homo sapi	EIF4EBP2,	-1.58963	1.70E-06	0.62908	5746.742	3625.015
A_23_P13	NM_00464	Homo sapi	CDK2AP1,	-1.58981	1.38E-16	0.62901	16746.96	10556.97
A_23_P16	NM_00568	Homo sapi	FARSLB,F	-1.58992	5.08E-14	0.62896	23352.8	14734.67
A_32_P45	NM_00662	FUS-intera	FUSIP1,NS	-1.59028	6.73E-13	0.62882	5167.465	3250.483
A_24_P38	NM_03126	heterogene	HNRPK,CS	-1.59053	6.24E-10	0.62872	7752.69	4840.894
A_32_P68	NM_01822	homolog of	SMU1,BWI	-1.59075	1.38E-06	0.62864	800.3549	503.6979
A_23_P15	NM_01436	3-hydroxyis	HIBCH,HIE	-1.59081	6.81E-18	0.62861	2809.666	1767.262
A_24_P33	BC007880	KIAA1970	KIAA1970	-1.59089	9.11E-17	0.62858	6558.61	4117.395
A_24_P10	NM_14717	Homo sapi	HS6ST2	-1.591	1.34E-08	0.62854	6495.932	4091.074
A_23_P40	AB018280	Chromosor	C14orf92	-1.59109	9.96E-07	0.6285	1436.713	891.0195
A_24_P39	NM_01544	DKFZP586	C1orf43,NI	-1.59114	1.27E-07	0.62848	3137.235	1961.292
A_23_P20	NM_00436	Homo sapi	CDH1,UVC	-1.59117	2.55E-06	0.62847	914.2181	566.7937
A_23_P12	NM_03256	Homo sapi	EBPL,EBR	-1.59126	1.54E-07	0.62843	32584.39	20571.21
A_23_P13	NM_00626	Homo sapi	RANBP2,T	-1.59134	5.92E-06	0.6284	5658.944	3574.152
A_24_P42	NM_00685	KDEL (Lys	KDELR2,E	-1.59135	9.48E-15	0.6284	5378.219	3378.056
A_23_P29	NM_01217	Homo sapi	FBXO7,FB	-1.59157	5.83E-16	0.62831	5330.464	3354.009
A_24_P33	NM_01900	hypothetica	FLJ20232,	-1.59166	6.18E-09	0.62827	2691.705	1700.579
A_24_P12	NM_01633	zinc finger	ZNF639,ZA	-1.5917	4.58E-09	0.62826	1441.005	909.6316
A_23_P62	NM_00167	Homo sapi	ATP1B1,A	-1.59203	6.46E-28	0.62813	16915.98	10626.02
A_23_P38	NM_17886	Homo sapi	SFXN4,BC	-1.59208	2.47E-08	0.62811	3569.316	2237.666
A_24_P68	BC044933	Hypothetic	LOC14690	-1.59242	4.12E-10	0.62797	1867.965	1177.3
A_23_P10	NM_00510	Homo sapi	RBM8A,Y1	-1.59246	1.88E-21	0.62796	72954.92	45887.73
A_23_P14	NM_08065	Homo sapi	ATP6V1E2	-1.59306	0.00004	0.62772	697.5256	433.6265
A_24_P18	AF447582	Casein kin	CSNK1A1	-1.59325	3.33E-06	0.62765	943.4659	584.7589
A_23_P40	AK025697	F-box prote	FBXO45	-1.59333	4.40E-06	0.62761	943.3374	600.8297
A_23_P10	NM_00682	Homo sapi	EBNA1BP2	-1.59347	2.62E-25	0.62756	69739.34	43789.17
A_24_P53	BC041849	Homo sapi	ROCK1	-1.59394	8.58E-07	0.62738	823.052	516.7827
A_23_P15	NM_01524	Ankyrin rep	ANKS1	-1.59395	5.49E-12	0.62737	1449.939	910.9813
A_23_P97	NM_00135	Homo sapi	DHX9,LKP	-1.59411	3.76E-16	0.62731	28779.08	18080.99
A_23_P13	NM_01657	PTX1 prote	PTX1	-1.59412	2.42E-31	0.62731	2475.502	1550.946
A_23_P14	NM_01654	Golgi phos	GOLPH2	-1.59434	7.75E-14	0.62722	2324.523	1460.636
A_23_P55	NM_00424	U5 snRNP-	EFTUD2,S	-1.59443	2.63E-14	0.62718	17229.38	10837.24
A_32_P27	NM_17363	hypothetica	MGC8902,	-1.59443	7.60E-07	0.62718	12288.94	7698.316
A_24_P32	NM_00486	suppressor	VPS4B,MI	-1.59469	1.07E-14	0.62708	2482.342	1552.365
A_23_P43	NM_15370	hypothetica	MGC33648	-1.59483	0.00001	0.62703	867.6085	534.3423
A_23_P10	NM_00451	integrin-link	ILK,P59,DK	-1.59489	0.00004	0.627	690.3697	438.2083
A_24_P39	NM_00474	cell cycle p	CCPG1,CF	-1.59499	0.00061	0.62696	994.1077	597.4018
A_23_P21	NM_00459	Homo sapi	SFRS10,TI	-1.59517	6.61E-15	0.62689	26521.96	16688.28
A_23_P73	AV650911	Ariadne ho	ARIH2	-1.5952	9.23E-08	0.62688	1397.874	866.8669
A_23_P18	NM_01590	Homo sapi	LAP3,LAP,	-1.59529	4.04E-08	0.62685	4381.067	2755.834
A_24_P25	BC034616	Hypothetic	LOC40158	-1.59545	2.13E-06	0.62678	1133.665	700.9919
A_23_P40	NM_17858	Homo sapi	HM13,H13	-1.5962	2.34E-13	0.62649	1881.865	1176.343
A_24_P10	NM_00572	ARP3 (acti	ACTR3,AR	-1.59647	2.79E-10	0.62638	2695.358	1688.553
A_23_P19	NM_00313	Homo sapi	SRPK1,SF	-1.59647	5.56E-10	0.62638	15963.38	10057.89
A_24_P94	AB046821	Sarcolemm	SLMAP	-1.5965	8.04E-12	0.62637	2225.83	1392.393
A_23_P70	BC038232	KIAA1549	KIAA1549	-1.59678	0.00007	0.62626	1473.495	902.0199
A_23_P45	NM_00390	Homo sapi	FUBP1,FB	-1.59702	6.53E-07	0.62617	6404.71	4001.905
A_32_P30	NM_00137	Homo sapi	DNCI2,IC2	-1.59703	5.28E-08	0.62616	2301.947	1447.601
A_32_P22	NM_00613	capping pr	CAPZA2,C	-1.59711	4.06E-08	0.62613	2846.433	1779.375

A_23_P15	NM_00617	Homo sapi	NSF,SKD2	-1.59725	6.11E-23	0.62608	5195.756	3253.567
A_23_P30	NM_00624	Homo sapi	PPP2R5E	-1.59763	1.05E-10	0.62593	3301.602	2060.941
A_23_P92	NM_00392	Homo sapi	MBD4,MEI	-1.59797	0.00001	0.62579	1510.868	926.3937
A_23_P63	NM_00679	Homo sapi	PRDX3,AC	-1.59809	2.20E-23	0.62575	10448.86	6543.841
A_24_P36	NM_01490	KIAA0941	RAB11FIP	-1.59847	0.00002	0.6256	1061.766	649.9741
A_32_P42	NM_03115	heterogene	HNRPA1,H	-1.59855	1.67E-18	0.62557	92371.45	57670.77
A_32_P6274		Unknown	ENST0000	-1.59879	1.16E-08	0.62547	1163.204	723.865
A_24_P25	NM_00605	LanC (bact	LANCL1,p	-1.59879	6.12E-10	0.62547	1380.281	867.0627
A_23_P20	NM_00648	Homo sapi	DYRK2	-1.59889	7.03E-14	0.62543	2024.195	1270.394
A_23_P26	NM_00280	proteasom	PSMD3,S3	-1.59905	2.56E-09	0.62537	19333.61	12182.81
A_23_P78	NM_03121	Homo sapi	SEH1L,Sel	-1.59909	3.13E-17	0.62535	3171.415	1983.747
A_23_P13	NM_01605	Homo sapi	PPIL1,CYF	-1.5993	1.89E-10	0.62528	5223.183	3272.645
A_23_P20	NM_00280	proteasom	PSMD1,S1	-1.59947	7.34E-17	0.62521	19754.91	12383.31
A_32_P10	NM_01488	predicted o	FAM3C,GS	-1.59956	2.45E-10	0.62517	1229.476	769.5884
A_23_P11	NM_02162	Homo sapi	PLEKHA1,	-1.59961	2.65E-10	0.62515	1769.345	1102.968
A_23_P25	NM_00279	Homo sapi	PSMA6,IO	-1.59972	1.01E-23	0.62511	49513.77	30964.04
A_23_P80	NM_00270	Homo sapi	PPAT,GPA	-1.59995	6.68E-12	0.62502	1367.392	854.5563
A_32_P54	NM_20346	Homo sapi	LASS6	-1.60004	1.12E-12	0.62498	2683.472	1676.162
A_24_P23	NM_00307	SWI/SNF r	SMARCD1	-1.60035	3.17E-24	0.62486	5273.303	3295.386
A_24_P33	NM_03212	implantatio	DKFZp564	-1.60037	1.26E-10	0.62486	9609.959	6019.571
A_32_P23	NM_00195	E2F transc	E2F6,E2F-	-1.60042	2.02E-07	0.62484	871.3644	542.176
A_24_P35	NM_01405	mitochondr	MRPL42,M	-1.60053	2.05E-22	0.62479	4953.562	3092.945
A_24_P13	NM_02237	ATP-deper	TOR3A,AD	-1.6007	0.00001	0.62473	1799.522	1141.25
A_23_P37	NM_01819	hypothetica	FLJ10719,	-1.60073	1.16E-09	0.62472	4269.92	2674.997
A_23_P11	NM_17347	Homo sapi	TMEM32	-1.60077	9.54E-12	0.6247	8954.018	5586.996
A_32_P13	NM_20700	Homo sapi	FAM45A	-1.60081	3.67E-12	0.62468	1801.247	1122.456
A_23_P57	NM_01745	Homo sapi	STAU	-1.60087	3.33E-13	0.62466	3453.266	2154.587
A_24_P60	NM_00029	phosphogl	PGK1,PGK	-1.60097	3.41E-16	0.62462	10370.46	6453.258
A_23_P21	NM_01779	KIAA1797	KIAA1797	-1.60099	2.88E-13	0.62461	2559.39	1597.415
A_23_P16	NM_00358	Homo sapi	DHX16,DB	-1.6012	5.20E-11	0.62453	1200.279	749.6278
A_32_P71437		ALU5_HUM	THC19106	-1.6014	5.39E-19	0.62445	14832.38	9269.528
A_23_P11	NM_02474	Homo sapi	HPS6,FLJ2	-1.60179	7.54E-15	0.6243	8818.855	5519.826
A_23_P63	NM_00517	ATP synth	ATP5C1,A	-1.60181	3.78E-22	0.62429	4213.769	2630.696
A_23_P81	NM_02283	Homo sapi	USP46,FL	-1.60198	0.00006	0.62423	893.2035	574.5344
A_23_P30	NM_00267	promyeloc	PML,MYL,I	-1.60239	0.00015	0.62407	565.7335	354.3791
A_23_P20	NM_01607	CGI-146 pr	C1orf121,F	-1.6024	7.98E-25	0.62406	5115.007	3191.796
A_24_P53	NM_00206	glutamate-	GLUL,GS,(	-1.60246	0.00057	0.62404	666.6729	401.8001
A_23_P43	NM_00376	Homo sapi	BECN1,AT	-1.60254	9.07E-12	0.62401	1622.986	1015.863
A_23_P12	NM_01905	hypothetica	FAM35A,M	-1.60287	4.34E-08	0.62388	1754.905	1087.212
A_23_P12	NM_03284	hypothetica	FLJ14803	-1.60287	8.63E-21	0.62388	5201.499	3246.103
A_24_P54	NM_00282	prothymosi	PTMA,TMS	-1.60298	1.80E-19	0.62384	151650.1	94824.66
A_24_P38	NM_00631	ring finger	PCGF3,RN	-1.60313	9.37E-07	0.62378	2691.715	1665.433
A_24_P187448		Unknown	A_24_P18	-1.60326	6.78E-06	0.62373	864.016	546.3519
A_24_P11	NM_03144	Homo sapi	C19orf12,M	-1.60332	5.08E-15	0.6237	2743.141	1709.663
A_24_P93	U68494	Solute carr	SLC30A1	-1.60364	3.20E-19	0.62358	3823.903	2386.406
A_23_P30	NM_00613	Homo sapi	CAPZA2,C	-1.60383	2.57E-08	0.62351	3230.378	2016.638
A_23_P38	NM_02483	Homo sapi	ZNF403,DI	-1.60412	4.90E-08	0.62339	2454.258	1538.918
A_32_P18	NM_00659	Homo sapi	POLD3,P6	-1.60419	9.69E-13	0.62337	2295.056	1425.28
A_23_P74	NM_00319	Homo sapi	TCEB3,SII	-1.6042	1.71E-09	0.62337	1887.006	1169.725
A_23_P34	NM_00565	Homo sapi	TFCP2,CP	-1.60429	2.97E-06	0.62333	1959.521	1209.555
A_24_P40	NM_01665	crooked ne	CRNKL1,C	-1.60437	0.00019	0.6233	770.9977	466.2198
A_32_P11	AB058761	Zinc finger	ZNF469	-1.60463	4.84E-14	0.6232	2705.327	1683.568

A_23_P31	AB037851	KIAA1430	KIAA1430	-1.6047	1.38E-15	0.62317	2340.823	1461.269
A_23_P16	NM_02099	creatine kir	CKMT1B,C	-1.60498	0.00073	0.62306	761.3447	495.945
A_23_P89	NM_01466	Homo sapi	NUP93,KIA	-1.60504	1.03E-18	0.62304	2953.258	1840.154
A_23_P35	BC003519	Homo sapi	FLJ34077,	-1.6051	0.00016	0.62302	1549.51	941.9775
A_24_P41	AK093442	Zinc finger	ZNF596	-1.6052	2.94E-17	0.62297	6338.872	3939.676
A_23_P75	NM_01619	Homo sapi	MPHOSPH	-1.6054	0.00003	0.6229	937.0288	597.8008
A_23_P25	NM_01602	Homo sapi	LACTB2,C	-1.6055	1.69E-10	0.62286	1761.956	1091.736
A_24_P92	NM_03313	caldesmon	CALD1,CD	-1.60556	1.43E-11	0.62284	3709.308	2309.302
A_23_P40	AB002365	KIAA0367	KIAA0367	-1.60572	0.00011	0.62277	632.1281	389.107
A_23_P16	NM_01450	down-regul	DRIM	-1.60578	6.87E-16	0.62275	1991.391	1238.901
A_32_P17	BM701156	Transcribe	BM701156	-1.6059	6.09E-06	0.6227	779.3273	490.8455
A_23_P20	NM_07847	Homo sapi	COX15	-1.606	2.70E-20	0.62267	3984.705	2482.114
A_23_P89	NM_01521	Homo sapi	EXOC7,EX	-1.60605	2.23E-06	0.62265	7123.385	4458.856
A_23_P59	NM_00564	Homo sapi	TAF11,TAF	-1.60653	2.47E-08	0.62246	2093.83	1297.255
A_24_P47	NM_00134	defender a	DAD1	-1.60657	4.33E-13	0.62244	30060.62	18789.64
A_32_P20	NM_01341	adenylate k	AK2,ADK2	-1.60673	4.83E-17	0.62238	3755.954	2337.196
A_24_P30	NM_14521	Homo sapi	MRPL30,R	-1.60702	5.34E-07	0.62227	6368.521	3953.224
A_24_P94	NM_03237	hypothetica	PCGF5,RN	-1.60732	4.30E-06	0.62215	800.0254	503.4678
A_23_P13	NM_00141	eukaryotic	EIF1AX,EIF	-1.60732	1.27E-08	0.62215	7881.28	4948.952
A_24_P10	NM_00674	sex comb c	SCML1	-1.60749	0.00002	0.62209	767.1179	484.3485
A_23_P13	NM_01460	Homo sapi	PIK3R4,p1	-1.60754	1.75E-09	0.62207	956.9172	594.6152
A_23_P28	NM_01825	Homo sapi	WDR12,YT	-1.60783	1.16E-16	0.62196	6226.81	3863.895
A_23_P31	NM_03284	hypothetica	C8orf76,MC	-1.60842	2.18E-12	0.62173	2221.702	1384.018
A_24_P94	NM_01812	hypothetica	FLJ10534,	-1.60843	3.95E-20	0.62172	4423.596	2753.409
A_23_P33	BC012766	Endoplasm	KIAA1181	-1.60856	5.60E-08	0.62167	7822.391	4886.974
A_23_P20	NM_00231	Homo sapi	ABLIM1,AE	-1.6086	1.06E-18	0.62166	12296.42	7663.327
A_23_P11	NM_01823	Homo sapi	CCAR1,CA	-1.60883	3.91E-12	0.62157	6484.716	4004.236
A_32_P22	NM_03245	Homo sapi	PHF6,MGC	-1.60911	6.02E-24	0.62146	3435.098	2134.674
A_23_P90	NM_01590	neuroblast	NAG,DKFZ	-1.60919	0.00032	0.62143	871.0463	564.399
A_32_P50	BQ888129	Hypothetic	BQ888129	-1.6092	2.86E-06	0.62143	1156.144	706.0736
A_23_P37	NM_00279	Homo sapi	PSMB5,X,M	-1.60922	1.25E-19	0.62142	53661.77	33255.31
A_24_P54847		Unknown	THC18817	-1.60925	0.00007	0.62141	614.4359	377.806
A_24_P39	NM_00525	cyclin G as	GAK,MGC	-1.60941	6.60E-16	0.62134	2264.554	1406.605
A_24_P29	AB046844	G protein-c	GPR107	-1.60982	2.99E-13	0.62119	1662.557	1033.785
A_23_P16	NM_00568	phenylalan	FARSLB,F	-1.60983	1.27E-12	0.62118	1541.624	959.5823
A_24_P33	NM_01220	heterogene	HNRPH3,2	-1.60992	1.82E-10	0.62115	3312.769	2058.277
A_24_P409410		Unknown	A_24_P40	-1.61	7.55E-10	0.62112	2928.667	1820.078
A_24_P12	NM_20350	Ras-GTPa	G3BP2	-1.61007	7.28E-10	0.62109	1116.134	694.6449
A_23_P25	NM_15274	Homo sapi	C7orf27,MC	-1.61024	6.19E-10	0.62103	14114.14	8810.904
A_23_P21	NM_00252	Homo sapi	NPM1,B23	-1.61024	8.63E-15	0.62103	111291.8	68822.31
A_23_P15	NM_00636	dendritic ce	hfl-B5,GA1	-1.61199	5.91E-29	0.62035	55961.73	34722.29
A_23_P55	NM_00447	flotillin 2	FLOT2,ES	-1.61213	0.00004	0.6203	2718.877	1700.006
A_23_P14	NM_01764	Homo sapi	FAM29A,R	-1.61224	0.00005	0.62026	824.8884	501.4764
A_23_P36	NM_17371	Homo sapi	NM_17371	-1.61228	1.14E-27	0.62024	16927.04	10496.68
A_23_P21	NM_01647	scotin	SCOTIN	-1.61231	6.03E-07	0.62023	56383.24	35348.43
A_23_P14	NM_01776	Homo sapi	LRRC40,F	-1.61257	2.94E-06	0.62013	1361.677	829.1675
A_23_P17	NM_02266	Homo sapi	ANAPC1,A	-1.61296	1.33E-16	0.61998	5844.679	3620.878
A_23_P10	NM_00134	Homo sapi	DAD1	-1.61307	1.29E-13	0.61994	31703.61	19773.06
A_23_P34	D87716	WD repeat	WDR43	-1.61324	6.32E-06	0.61987	3466.94	2154.254
A_24_P68	AK123675	Lix1 homol	MGC4671	-1.61328	4.84E-09	0.61985	932.2814	580.3984
A_23_P12	NM_00445	Homo sapi	ERH,DROE	-1.61343	2.78E-15	0.6198	27052.46	16729.02
A_23_P13	NM_00210	histidyl-tRN	HARS,HRS	-1.61358	6.08E-08	0.61974	5411.984	3347.195

A_24_P713185		Unknown	A_24_P71	-1.61368	4.61E-08	0.6197	3364.793	2090.407
A_23_P15	NM_13921	Homo sapi	TAF15,RBI	-1.61368	1.27E-07	0.6197	19129.44	11788.25
A_32_P18	NM_00226	karyopheri	KPNB1,IMI	-1.6137	5.44E-08	0.61969	46740.29	29085.51
A_23_P58	NM_02409	hypothetica	MGC5297,	-1.61396	1.42E-12	0.61959	2564.269	1590.463
A_23_P34	NM_00548	Homo sapi	PIGK,GPI8	-1.61421	8.45E-17	0.6195	2139.913	1327.029
A_32_P215691		Unknown	A_32_P21	-1.6145	5.33E-09	0.61939	1087.317	672.3057
A_24_P12	NM_01818	hypothetica	SBNO1,Sn	-1.61461	2.14E-09	0.61935	2885.494	1788.407
A_23_P86	NM_00294	Homo sapi	RPA2,REP	-1.61478	1.18E-22	0.61928	5238.316	3245.539
A_24_P24	NM_00545	zinc finger	ZNF265,ZI	-1.61536	5.36E-20	0.61906	4346.306	2692.592
A_23_P21	NM_03349	hexokinase	HK1,HKI,H	-1.61591	5.52E-19	0.61885	2783.209	1722.523
A_23_P12	NM_13855	Homo sapi	PPP1R8,A	-1.61594	5.55E-14	0.61883	3222.984	1995.074
A_23_P14	NM_00114	Homo sapi	AMFR,GP7	-1.61601	3.09E-09	0.61881	2976.344	1826.927
A_32_P70	AK124072	KDEL (Lys	KDELR2	-1.61605	7.06E-06	0.61879	1285.747	811.9349
A_23_P21	NM_01812	Homo sapi	TMEM33,F	-1.61637	1.46E-19	0.61867	5528.775	3413.972
A_32_P8666		Unknown	ENST0000	-1.61652	5.80E-12	0.61861	7724.79	4747.044
A_23_P20	NM_00690	Homo sapi	MYO9A,my	-1.61657	4.73E-06	0.61859	726.4902	446.7683
A_32_P21	AL049250	Homo sapi	DKFZp547	-1.61661	1.39E-10	0.61858	1221.644	755.3539
A_23_P71	NM_00462	Homo sapi	FANCG,FA	-1.61661	1.42E-24	0.61858	5496.154	3400.26
A_24_P74	AJ420454	Hypothetic	LOC28633	-1.6169	1.22E-06	0.61847	918.8075	577.2998
A_24_P27	AK123385	Speckle-ty	SPOP	-1.6169	3.74E-15	0.61847	2284.282	1410.848
A_24_P206848		ALU1_HUM	THC18433	-1.61707	5.94E-11	0.6184	2340.428	1452.321
A_23_P50	NM_01761	Homo sapi	DONSON,I	-1.61735	0.00005	0.6183	4563.651	2831.118
A_23_P58	NM_01447	dimethylad	HSA9761	-1.61739	5.34E-13	0.61828	7586.668	4713.92
A_24_P37	NM_17316	Homo sapi	NFATC3,N	-1.61739	1.84E-08	0.61828	1014.242	625.3828
A_23_P79	NM_01607	Homo sapi	VPS24,NE	-1.6174	1.67E-17	0.61828	4091.338	2530.258
A_24_P10	NM_01247	tyrosine 3-r	YWHAG,14	-1.61749	1.57E-27	0.61824	69992.46	43294.51
A_23_P79	NM_00384	succinate-(	SUCLG1,S	-1.61752	1.34E-21	0.61823	5896.816	3650.491
A_24_P62	NM_01821	Chromosor	C20orf31	-1.61777	6.77E-12	0.61813	1746.394	1083.304
A_23_P10	NM_01884	Homo sapi	C20orf24,F	-1.61818	4.60E-21	0.61798	125383.3	77616.05
A_23_P15	NM_02035	PEST-cont	PCNP,DKF	-1.61873	2.89E-26	0.61777	10636.68	6566.178
A_23_P31	BC007792	YLP motif (	YLPM1	-1.6192	1.19E-10	0.61759	2199.261	1363.77
A_32_P58606		Unknown	THC19254	-1.61955	3.93E-06	0.61746	777.7333	485.4423
A_23_P44	NM_01784	Homo sapi	COMMD8,I	-1.61985	5.95E-09	0.61734	5425.173	3348.926
A_23_P80	NM_00081	Homo sapi	GART,AIR	-1.61987	1.16E-10	0.61733	5885.586	3644.364
A_23_P38	NM_00589	Homo sapi	ACAT2	-1.6199	9.32E-12	0.61732	1593.074	987.1494
A_32_P10	NM_01341	adenylate k	AK3L1,AK3	-1.62007	0.00003	0.61726	9376.463	5818.352
A_23_P21	NM_00437	Homo sapi	COL12A1,(	-1.62009	9.09E-14	0.61725	3458.034	2132.648
A_23_P40	NM_00591	Homo sapi	MAT2A,MA	-1.62024	1.67E-16	0.61719	13002.41	8047.646
A_23_P11	NM_20330	Homo sapi	C9orf132,E	-1.62078	1.55E-08	0.61699	10262.6	6382.571
A_24_P66	AK023696	Hypothetic	LOC40150	-1.62085	0.00008	0.61696	982.1169	624.6351
A_23_P92	NM_00041	Homo sapi	HSD17B4,I	-1.62123	4.00E-12	0.61681	4043.08	2500.445
A_24_P34	NM_02114	X-ray repai	XRCC5,KU	-1.62125	9.85E-15	0.61681	6658.064	4117.672
A_32_P49	NM_02474	disrupted i	ALG9,DIBD	-1.62134	0.00001	0.61677	1313.098	792.9303
A_23_P11	NM_01794	Homo sapi	FBXO34,F	-1.6216	1.98E-08	0.61667	1183.436	724.6252
A_23_P10	AB014592	KIAA0692	KIAA0692	-1.62186	2.69E-06	0.61658	782.4937	480.7432
A_32_P43	NM_00279	Homo sapi	PSMA6,IO	-1.62188	4.91E-25	0.61657	76900.28	47470.16
A_23_P48	NM_00286	Homo sapi	PYGL	-1.62192	7.53E-10	0.61655	5264.405	3256.871
A_23_P31	NM_01244	six transme	STEAP1,S	-1.6225	2.39E-09	0.61633	2639.75	1622.533
A_23_P12	NM_00271	protein pho	PPP2CA,R	-1.62264	9.20E-15	0.61628	6705.466	4147.944
A_23_P13	NM_00641	ribonuclea	RPP30,TS	-1.62286	2.92E-10	0.6162	1758.242	1078.158
A_23_P20	NM_00485	Homo sapi	ROCK2,KI	-1.62292	2.13E-06	0.61617	807.1787	499.1939
A_23_P47	NM_01614	Homo sapi	HSD17B12	-1.62311	8.16E-10	0.6161	8781.235	5449.593

A_23_P85	NM_01332	Homo sapi	SNX10,MG	-1.62312	1.06E-17	0.6161	3990.902	2458.536
A_23_P41	NM_00405	Homo sapi	BCL2L2,BC	-1.62319	4.83E-14	0.61607	1842.846	1136.968
A_23_P73	NM_00244	moesin	MSN	-1.62373	3.01E-08	0.61586	1837.951	1122.821
A_23_P10	NM_00248	Homo sapi	NDN,HsT1	-1.62388	5.63E-13	0.61581	1470.042	906.2372
A_23_P41	NM_01835	hypothetica	FLJ11200	-1.62423	6.26E-12	0.61568	1293.998	795.3206
A_24_P31	NM_00735	low density	COG2,LDL	-1.62448	5.24E-06	0.61558	683.6085	419.9556
A_24_P726336		Unknown	THC19678	-1.62513	0.00019	0.61534	577.4258	351.2767
A_24_P92	NM_00185	solute carri	SLC31A1,C	-1.62529	1.54E-06	0.61528	764.6296	469.7388
A_23_P20	NM_00193	Homo sapi	DLAT,DLT	-1.62533	5.28E-13	0.61526	2988.103	1832.009
A_23_P16	AK091324	Hypothetic	LOC21985	-1.62534	2.39E-12	0.61525	2556.657	1571.623
A_23_P20	NM_00626	Homo sapi	RAD21,HR	-1.62538	4.07E-10	0.61524	8646.831	5342.163
A_24_P37	NM_01562	CGI-43 pro	C7orf28A,C	-1.62544	2.24E-11	0.61522	8744.921	5408.806
A_23_P11	NM_01773	hypothetica	THUMPD1	-1.62552	3.45E-15	0.61519	2758.999	1696.974
A_23_P30	NM_00041	Homo sapi	HLCS,HCS	-1.62579	4.65E-06	0.61509	711.9459	435.8749
A_23_P11	NM_01602	Homo sapi	DHRS7,ret	-1.62606	1.10E-12	0.61498	2247.923	1385.147
A_24_P531074		Unknown	A_24_P53	-1.62651	2.70E-11	0.61481	2450.927	1503.063
A_32_P217428		Unknown	A_32_P21	-1.62666	0.00047	0.61476	954.1199	612.1422
A_23_P21	NM_01451	Homo sapi	UBP1,LBP	-1.6267	0.00004	0.61474	6065.337	3756.371
A_24_P93	BC034716	NEDD8 ulti	NYREN18	-1.62675	0.00002	0.61472	788.6563	476.0966
A_23_P42	NM_01761	downstrear	DONSON,I	-1.6269	0.00002	0.61467	1173.882	703.2502
A_23_P83	NM_02483	Homo sapi	ZNF672,FL	-1.62723	1.80E-15	0.61454	1926.868	1185.105
A_23_P66	NM_00037	Homo sapi	UMPS,OPI	-1.62725	5.41E-21	0.61453	9974.699	6144.066
A_24_P29	NM_13080	hypothetica	LOC13361	-1.62731	1.00E-09	0.61451	1336.636	815.499
A_32_P19	NM_18178	hypothetica	DPY19L4	-1.6274	8.77E-08	0.61448	944.8699	579.0768
A_23_P16	NM_20738	FLJ43339	FLJ43339,	-1.62747	6.33E-14	0.61445	2077.498	1279.334
A_23_P11	NM_00412	hypothetica	LOC55289	-1.62788	8.56E-07	0.6143	11791.88	7263.677
A_23_P37	AK027341	KIAA1856	KIAA1856	-1.62828	7.44E-07	0.61414	4314.78	2633.686
A_24_P289818		Unknown	A_24_P28	-1.62832	6.51E-07	0.61413	973.6029	591.6293
A_23_P73	NM_00244	Homo sapi	MSN	-1.62843	1.04E-16	0.61409	10940.79	6743.278
A_23_P11	NM_15333	Homo sapi	TCEAL8,M	-1.62846	6.87E-14	0.61408	6394.168	3937.411
A_24_P37	NM_00587	splicing fac	SF3A1,PR	-1.62852	2.95E-10	0.61405	1108.238	682.2337
A_24_P64	AK025909	CDNA: FL	AK025909	-1.62856	0.00012	0.61404	573.7554	351.2115
A_23_P10	NM_01514	Homo sapi	ZCCHC14,	-1.62857	4.36E-13	0.61404	1729.509	1059.853
A_24_P34	NM_01463	chondrocyt	MTFR1,CH	-1.62859	3.43E-08	0.61403	1045.384	647.144
A_23_P15	NM_03232	Homo sapi	ZDHHC16,	-1.62876	1.02E-09	0.61396	1789.74	1105.939
A_23_P13	NM_00256	prostatic bi	PBP,HCNF	-1.62876	5.72E-10	0.61396	66327.62	40963.88
A_23_P25	NM_02114	PC4 and S	PSIP1,p52	-1.62883	8.92E-11	0.61394	1475.716	902.1634
A_24_P715924		ALU5_HUM	THC19985	-1.62928	3.01E-07	0.61377	1576.136	955.2605
A_32_P14	AK091132	Sestrin 3	SESN3	-1.62933	4.28E-12	0.61375	4998.73	3076.035
A_23_P14	NM_00577	Homo sapi	CNIH,CNIL	-1.62934	1.28E-23	0.61375	19349.84	11887.99
A_23_P86	NM_00375	Homo sapi	EIF3S10,E	-1.62954	4.05E-18	0.61367	47624.42	29328.22
A_23_P20	NM_00686	RAB35, me	RAB35,RA	-1.62957	1.39E-12	0.61366	9085.121	5593.719
A_23_P25	NM_01826	Homo sapi	WDR41,MS	-1.62994	4.10E-17	0.61352	2712.424	1663.525
A_23_P88	NM_15381	Homo sapi	SLC38A6,M	-1.62999	1.63E-18	0.6135	5245.739	3211.939
A_23_P14	NM_00660	Homo sapi	TCFL5,CH	-1.63001	3.06E-08	0.61349	2264.878	1376.485
A_23_P15	NM_01488	predicted o	FAM3C,GS	-1.63005	0.00003	0.61348	1007.641	601.51
A_23_P24	NM_00255	Homo sapi	OSBP,OSE	-1.63072	8.40E-07	0.61323	818.8063	500.1938
A_32_P199725		Unknown	A_32_P19	-1.63089	7.60E-10	0.61316	6178.33	3812.568
A_23_P25	NM_01404	Homo sapi	UNC50,UR	-1.63099	3.25E-28	0.61313	6519.039	3996.458
A_23_P47	NM_01818	Homo sapi	DHX32,DD	-1.63103	0.00003	0.61311	787.3934	493.6679
A_23_P29	NM_02068	Homo sapi	C3orf14,HT	-1.63164	2.02E-27	0.61288	7239.114	4434.724
A_24_P40	NM_00437	COX11 (ye	COX11,CC	-1.63165	2.41E-18	0.61288	2225.944	1363.701

A_23_P11	NM_02237	Homo sapi	TOR3A,AD	-1.63174	2.77E-21	0.61284	13951.4	8563.324
A_23_P11	NM_00595	Homo sapi	MTHFD1,M	-1.63207	4.27E-09	0.61272	9626.783	5905.681
A_23_P25	NM_00289	Homo sapi	RBBP8,RIN	-1.63208	2.03E-08	0.61272	3938.347	2418.787
A_23_P14	AB014595	Cullin 4B	CUL4B	-1.63243	3.00E-14	0.61258	1906.458	1169.583
A_23_P38	NM_01491	Homo sapi	WHSC1,NS	-1.63244	9.87E-11	0.61258	1393.466	849.4124
A_24_P68	NM_02084	Homo sapi	NM_02084	-1.63268	1.85E-07	0.61249	1107.842	672.0134
A_32_P63	NM_00301	mitogen-ac	MAP2K4,JI	-1.63285	4.60E-23	0.61243	3273.336	2004.568
A_23_P10	AK000900	Hypothetic	FLJ10038	-1.63322	1.66E-12	0.61229	1697.827	1041.805
A_23_P20	NM_00112	Homo sapi	ADSS,ADE	-1.63327	9.81E-14	0.61227	6134.24	3770.948
A_23_P34	NM_00278	Homo sapi	PSMA2,MU	-1.63362	5.58E-17	0.61214	22301.21	13715.42
A_24_P96	NM_03228	Leucine zip	LDOC1L	-1.63363	0.00021	0.61213	844.4997	497.3394
A_32_P19	AK094044	CDNA FLJ	AK094044	-1.63368	0.00009	0.61212	619.1428	371.4811
A_24_P29	NM_01643	hepatocellu	PHF20,NZI	-1.63379	2.01E-06	0.61207	844.6005	513.119
A_24_P94	NM_19806	Homo sapi	GNPNAT1	-1.63381	1.15E-17	0.61207	3034.423	1856.742
A_23_P10	NM_13841	hypothetica	KTI12,TOT	-1.63382	0.00002	0.61206	729.2769	440.9546
A_23_P41	NM_14460	hypothetica	FLJ31795	-1.63391	8.84E-24	0.61203	3317.159	2030.245
A_23_P38	NM_02024	Tubby like	TULP4	-1.63403	0.0002	0.61198	1181.656	696.7554
A_23_P11	NM_02494	Homo sapi	C9orf76,RN	-1.63425	1.43E-26	0.6119	5408.188	3309.053
A_23_P31	NM_01514	Homo sapi	GPD1L,KIA	-1.6344	7.03E-09	0.61185	2152.293	1307.08
A_23_P25	NM_00135	Homo sapi	AKR1C1,C	-1.63459	1.93E-30	0.61177	174978.7	107066
A_32_P14	BU608659	RAS p21 p	RASA2	-1.63482	0.00048	0.61169	4255.69	2576.566
A_24_P91	AK024986	Phosphata	PTEN	-1.63485	1.71E-10	0.61168	1258.129	772.3856
A_23_P34	NM_01430	Homo sapi	NIFUN,ISC	-1.63489	5.27E-17	0.61166	13792.81	8461.631
A_23_P37	NM_03223	hypothetica	FLJ22875	-1.63499	5.92E-17	0.61162	35855.6	22022.44
A_23_P50	NM_15363	Homo sapi	HOXA3,HC	-1.63508	6.52E-09	0.61159	2636.439	1609.243
A_24_P79	NM_00637	NS1-assoc	SYNCRIP,	-1.63522	3.98E-13	0.61154	18348.69	11130.93
A_23_P16	NM_03195	Homo sapi	KCTD10,U	-1.63524	1.50E-08	0.61153	1760.622	1068.005
A_23_P17	NM_01765	Homo sapi	DYM,DMC	-1.63546	6.64E-18	0.61145	2582.466	1579.107
A_23_P21	AK000619	Two pore s	TPCN1	-1.63559	2.56E-08	0.6114	5767.874	3529.502
A_32_P15	NM_01910	beta-1,4 m	ALG1,HMT	-1.63562	4.62E-11	0.61139	1952.374	1199.774
A_23_P30	NM_02047	Homo sapi	GALNT1,G	-1.6357	7.16E-10	0.61136	1527.054	928.2763
A_23_P18	NM_00482	Homo sapi	ABCG2,MF	-1.63583	7.63E-21	0.61131	13159.1	8037.571
A_24_P76	BC035146	Clone A9A	BC035146	-1.63624	3.04E-12	0.61116	2141.459	1303.854
A_23_P16	NM_00333	ubiquitin-co	UBE2B,HR	-1.63695	0.00003	0.61089	629.2195	381.2485
A_23_P20	NM_01624	Homo sapi	AKAP11,PI	-1.63709	2.34E-06	0.61084	1578.205	984.9123
A_24_P250614		S3A1_HUM	THC19887	-1.63711	9.42E-14	0.61083	1760.03	1073.341
A_23_P11	NM_00226	Homo sapi	KPNB1,IMI	-1.63719	1.52E-26	0.6108	9053.352	5525.319
A_23_P11	NM_00196	eukaryotic	EIF4E,CBF	-1.63725	1.89E-09	0.61078	9086.799	5586.165
A_23_P15	NM_14897	Homo sapi	PSMA1,NU	-1.63753	3.23E-17	0.61068	35833.76	21967.17
A_24_P165595		ARL5_HUM	THC18801	-1.63755	1.72E-07	0.61067	1519.108	938.5295
A_23_P21	NM_00590	Homo sapi	SMAD1,BS	-1.6379	4.76E-12	0.61054	1744.706	1058.883
A_23_P24	NM_01425	Homo sapi	TMEM5,HF	-1.63792	2.31E-24	0.61053	10287.54	6287.877
A_24_P36	AB046784	Chromodo	CHD8	-1.63804	0.00009	0.61049	2025.105	1256.452
A_23_P10	NM_00591	Homo sapi	MDH1,MDI	-1.63807	3.94E-24	0.61047	58693.77	35762.03
A_24_P20	NM_00044	regulatory	RFX5	-1.6382	7.45E-14	0.61042	1464	894.2052
A_23_P25	NM_02182	Homo sapi	MRPS35,M	-1.63844	7.36E-24	0.61034	14571.19	8903.957
A_23_P68	NM_00167	Homo sapi	ATP1B3,A	-1.63854	1.69E-18	0.6103	48690.66	29603.91
A_24_P55391		Unknown	A_24_P55	-1.63868	6.83E-13	0.61025	2248.185	1374.891
A_32_P77	S63912	Heterogen	HNRPA3P	-1.63914	7.81E-11	0.61008	30478.91	18405.7
A_24_P63	BC039118	Homo sapi	STX6	-1.63923	4.44E-09	0.61004	1194.216	723.4832
A_24_P31	AB011132	Aquarius h	AQR	-1.63986	0.00002	0.60981	1648.977	987.6945
A_24_P50	NM_01341	adenylate k	AK2,ADK2	-1.63987	2.19E-10	0.60981	5909.789	3615.085

A_24_P71	AK023423	Kynurenine	KMO	-1.64005	1.82E-17	0.60974	2289.531	1396.351
A_24_P37	NM_00655	SNARE pr	YKT6	-1.64023	7.46E-11	0.60967	2753.4	1681.539
A_23_P31	NM_00327	Homo sapi	TMEM1,EH	-1.64024	0.00018	0.60967	774.9283	455.4285
A_23_P42	NM_02192	hypothetica	FLJ21820	-1.64042	3.00E-07	0.6096	951.5658	587.59
A_23_P15	NM_01795	FLJ20758	FLJ20758	-1.64059	1.38E-27	0.60954	8049.496	4908.902
A_24_P616082		Unknown	A_24_P61	-1.64083	1.83E-07	0.60945	930.0529	571.6462
A_23_P30	NM_01554	Dystonin	DST	-1.64095	0.0002	0.6094	504.2472	306.3023
A_23_P12	NM_00182	Homo sapi	CLCN3,CL	-1.64107	4.18E-07	0.60936	951.8256	575.506
A_23_P86	NM_01233	c-myc bind	MYCBP,AM	-1.6412	2.85E-06	0.60931	1200.554	719.3713
A_23_P16	NM_02206	Homo sapi	C14orf133,	-1.64122	1.42E-06	0.6093	1176.736	703.9691
A_24_P213175		Unknown	A_24_P21	-1.6413	5.39E-11	0.60927	7538.688	4592.092
A_24_P41	NM_00646	trans-golgi	TGOLN2,T	-1.64133	3.11E-28	0.60926	11193.87	6820.613
A_24_P19	NM_01449	golgi phosp	GOLPH4,P	-1.64142	0.00004	0.60923	1029.772	609.7339
A_32_P21	AF130091	Potassium	KCTD9	-1.64168	4.99E-06	0.60913	713.343	434.9258
A_24_P14	NM_00599	translin-ass	TSNAX,TR	-1.64185	2.11E-07	0.60907	2533.842	1538.175
A_32_P172608		Unknown	A_32_P17	-1.64294	0.00012	0.60866	16373.86	9804.396
A_24_P28	NM_00368	calcium/cal	CASK,LIN2	-1.64308	0.00012	0.60861	747.5471	441.6408
A_24_P16	NM_13931	Homo sapi	YME1L1,F	-1.64327	2.51E-12	0.60854	1794.721	1088.614
A_23_P92	NM_02019	Homo sapi	C2orf33,GL	-1.64329	3.45E-10	0.60854	9850.028	5997.677
A_23_P15	NM_01514	Homo sapi	METAP1,K	-1.64342	4.50E-26	0.60849	6502.88	3955.766
A_24_P24	NM_01594	CGI-12 pro	MTERFD1,	-1.64345	3.67E-09	0.60848	1539.361	930.0203
A_23_P331087		Unknown	A_23_P33	-1.64347	2.35E-08	0.60847	3449.374	2093.51
A_23_P36	NM_02291	Homo sapi	VPS33A,FI	-1.6435	3.26E-09	0.60846	1967.915	1205.051
A_24_P23	AB020660	KIAA0853	KIAA0853	-1.64369	0.00018	0.60839	719.4535	423.191
A_24_P11	AL050139	Hypothetic	FLJ13910	-1.6441	2.20E-10	0.60824	3055.886	1852.522
A_23_P30	NM_14530	similar to R	LOC20245	-1.64423	3.91E-18	0.60819	4980.572	3033.861
A_32_P89	AF110908	TNF recep	TRAF3	-1.64432	6.64E-14	0.60815	2332.877	1415.048
A_23_P12	NM_00631	aminopepti	NPEPPS,P	-1.64443	4.95E-31	0.60811	16467.62	10012.41
A_24_P30	NM_00349	stannin	SNN	-1.64474	0.00008	0.608	881.9981	555.8723
A_23_P10	NM_00572	Homo sapi	ACTR3,AR	-1.64488	1.54E-17	0.60795	6288.215	3825.459
A_23_P11	NM_00352	Homo sapi	HIST1H3A	-1.64527	4.41E-24	0.6078	25514.68	15527.08
A_23_P13	NM_00373	aldo-keto r	AKR1C3,D	-1.64528	3.33E-18	0.6078	110620	67535.21
A_23_P38	NM_02113	Homo sapi	TNFAIP1,E	-1.64534	6.57E-22	0.60778	2513.418	1527.582
A_32_P22	NM_00213	RNA bindir	RBMX,RNI	-1.6455	3.49E-08	0.60772	4289.598	2608.876
A_23_P10	NM_00509	zinc finger	ZNF262,M	-1.64566	0.00035	0.60766	505.6251	303.6669
A_23_P33	NM_02249	SDS3 prot	SDS3,FLJ	-1.64614	3.98E-10	0.60748	1297.629	785.1242
A_23_P58	NM_02456	hypothetica	FLJ14054	-1.6468	9.76E-15	0.60724	2386.552	1453.013
A_24_P19	NM_00721	SEC63 pro	SEC63,ER	-1.64684	9.32E-15	0.60722	1728.342	1049.158
A_23_P40	NM_01634	Centromer	CENPF	-1.64694	9.53E-15	0.60719	13221.37	8027.108
A_23_P38	NM_00357	VAMP (ves	VAPA,VAP	-1.647	4.48E-12	0.60717	7122.447	4344.46
A_24_P33	NM_02245	hypothetica	XPO4,FLJ	-1.64723	0.00025	0.60708	555.2942	331.5353
A_23_P119794		Unknown	THC18076	-1.64741	0.0001	0.60702	893.4854	561.8853
A_23_P70	NM_15229	Homo sapi	TARS,ThrF	-1.64749	1.00E-08	0.60699	28510.28	17443.73
A_24_P25	NM_00029	phosphogl	PGK1,PGK	-1.64764	0.00011	0.60693	2306.452	1385.306
A_23_P28	NM_01795	hypothetica	PLEKHB2,	-1.64768	0.00003	0.60691	926.6981	578.3355
A_24_P45	AK024653	Hypothetic	LOC90246	-1.6486	1.09E-06	0.60658	944.4507	580.5863
A_23_P14	NM_00259	Homo sapi	PDCD2,RF	-1.64922	3.03E-13	0.60635	3890.06	2363.119
A_24_P14	NM_02278	likely ortho	ARV1	-1.64946	1.54E-06	0.60626	1097.564	655.5682
A_23_P85	NM_01837	hypothetica	FLJ11273	-1.6498	4.63E-13	0.60614	3327.7	2018.985
A_24_P28	NM_01511	KIAA0399	ZZEF1,ZZ	-1.65014	1.12E-06	0.60601	3096.09	1868.785
A_24_P16	NM_01829	hypothetica	CWF19L1,	-1.6503	2.40E-11	0.60595	1390.712	849.9185
A_24_P71	NM_01467	basic leuci	BZW1,orf,E	-1.65035	1.44E-15	0.60593	14597.19	8862.666

A_23_P37	AK125140	Homo sapi	ARG99,OL	-1.65048	0.00001	0.60588	717.9404	431.1152
A_23_P16	BC028424	Rififylin	RFFL	-1.6505	1.73E-07	0.60588	3117.912	1886.948
A_32_P12820		143Z_HUM	THC19778	-1.65059	3.42E-12	0.60584	6141.222	3717.954
A_24_P779605		Unknown	THC19984	-1.65063	3.82E-13	0.60583	2925.449	1768.731
A_24_P96	NM_01634	Centromer	CENPF	-1.65082	0.00066	0.60576	506.9328	301.1445
A_24_P32	NM_01327	nasopharyn	ANKRD11,	-1.65102	7.83E-16	0.60568	2266.552	1373.293
A_24_P32	D28476	Thyroid ho	TRIP12	-1.65106	0.00004	0.60567	1032.841	607.3013
A_32_P10	AL832591	Serine/thre	STYX	-1.65108	1.93E-21	0.60566	3261.365	1975.339
A_23_P21	NM_00500	Homo sapi	NDUFA5,B	-1.65118	1.03E-20	0.60563	9704.543	5857.637
A_23_P10	NM_02024	Homo sapi	CHPT1,CP	-1.65126	6.04E-18	0.6056	3916.612	2370.765
A_32_P187875		Unknown	A_32_P18	-1.65158	1.23E-21	0.60548	9632.188	5818.797
A_24_P48	AF086257	ST3 beta-g	SIAT4B	-1.65183	1.68E-08	0.60539	902.6105	544.0244
A_23_P57	NM_00544	Homo sapi	CHAF1B,C	-1.65185	2.06E-38	0.60538	3809.156	2309.101
A_23_P47	NM_01402	Homo sapi	WDR37,FL	-1.65188	0.00042	0.60537	730.4811	422.7723
A_32_P11	AK025431	Hypothetic	LOC25398	-1.65204	1.18E-08	0.60531	2374.063	1429.196
A_23_P11	NM_00116	Homo sapi	BIRC5,API	-1.65213	1.33E-17	0.60528	27589.52	16713.54
A_23_P63	NM_01811	WD40 repe	BRWD2,DI	-1.65227	4.43E-11	0.60523	1457.388	886.8114
A_24_P25	NM_02115	hypothetica	TXNDC13,	-1.65245	6.48E-11	0.60516	1999.561	1205
A_32_P103464		Unknown	A_32_P10	-1.65278	1.43E-17	0.60504	2677.673	1619.797
A_23_P21	NM_02212	Homo sapi	PERP,THV	-1.65283	2.04E-14	0.60502	8518.168	5162.214
A_32_P20	AB032986	Homo sapi	KIAA1160	-1.65287	9.02E-06	0.60501	1034.49	610.4497
A_23_P13	NM_00387	Homo sapi	GGH,GH	-1.65353	4.28E-14	0.60477	5714.454	3469.875
A_24_P69	NM_03298	Sec23 (S.	SEC23B	-1.65362	5.14E-13	0.60473	2080.27	1254.511
A_32_P39	AF256215	Homo sapi	cycle-like f	-1.65422	0.00017	0.60452	614.8119	380.8998
A_23_P41	NM_01426	Homo sapi	MAPRE2,E	-1.65422	5.33E-06	0.60451	700.7576	427.8157
A_23_P76	AF173389	Early endo	EEA1	-1.65443	5.52E-08	0.60444	2034.282	1220.558
A_23_P10	NM_02404	Homo sapi	DDX50,GU	-1.65458	7.70E-21	0.60438	8991.041	5444.75
A_23_P14	NM_00278	Homo sapi	PSMA3,HC	-1.65463	2.91E-20	0.60436	30878.59	18587.71
A_23_P25	NM_00287	Homo sapi	RAD23B,P	-1.65465	6.94E-09	0.60436	28204.09	17143.21
A_24_P34	NM_00679	mortality fa	MORF4,CS	-1.65475	2.38E-15	0.60432	4397.053	2651.029
A_23_P17	NM_02019	Homo sapi	SMYD2,HS	-1.65486	1.63E-07	0.60428	6604.801	4012.963
A_23_P25	BC013629	WNK lysine	WNK1	-1.65521	1.86E-15	0.60415	2886.466	1745.733
A_23_P20	NM_02451	Homo sapi	CYP2R1,M	-1.65603	1.75E-09	0.60385	1387.157	845.489
A_23_P55	NM_00128	Homo sapi	AP2B1,AD	-1.65604	1.64E-12	0.60385	1320.007	797.1115
A_32_P137745		Unknown	A_32_P13	-1.65604	0.00046	0.60385	1251.149	784.4977
A_32_P15	AF517524	Heterogen	HNRPA3	-1.65628	9.90E-20	0.60376	19840.8	12014.92
A_24_P22	NM_01538	likely ortho	PREI3,2C4	-1.65655	2.63E-11	0.60366	1899.822	1142.792
A_23_P21	NM_02011	hypothetica	LARS,LEU	-1.65656	8.71E-09	0.60366	1632.045	973.4978
A_23_P15	NM_00137	Homo sapi	DNCI2,IC2	-1.65656	2.54E-09	0.60366	4127.372	2499.953
A_23_P10	NM_02038	Cancer sus	AF15Q14	-1.65678	0.00002	0.60358	755.9504	448.2071
A_32_P42	AB011164	Family with	KIAA0592	-1.65753	2.00E-12	0.60331	4743.088	2852.897
A_23_P38	NM_05285	Homo sapi	CCDC16,M	-1.65758	7.13E-07	0.60329	1809.822	1077.58
A_24_P69	AK021884	Hypothetic	LOC44043	-1.65778	0.00011	0.60321	645.1212	380.7188
A_23_P50	NM_00122	Homo sapi	CASP6,MC	-1.65784	8.59E-19	0.60319	1963.778	1184.761
A_23_P67	NM_00633	nuclear DN	C1D,SUNC	-1.65785	1.16E-22	0.60319	3401.878	2051.897
A_24_P12	AK091674	Hypothetic	FLJ13291	-1.6581	4.40E-09	0.6031	1469.046	878.0083
A_24_P135801		Unknown	A_24_P13	-1.65829	7.36E-13	0.60303	24996.03	15007.75
A_23_P25	NM_01486	chromoson	CNAP1,KI	-1.6583	0.00003	0.60303	2322.085	1423.721
A_24_P90	NM_01682	adducin 3 (	ADD3,ADD	-1.6583	3.59E-13	0.60303	2996.548	1805.003
A_32_P16	AF161398	Chromosor	C6orf115	-1.65848	1.47E-09	0.60296	5939.174	3593.244
A_23_P32	NM_05282	Homo sapi	WDR5,BIG	-1.65853	3.12E-09	0.60294	1105.147	670.1173
A_23_P14	NM_00312	Homo sapi	SQLE	-1.65868	6.81E-08	0.60289	2591.224	1548.457

A_23_P21	NM_01810	Homo sapi	EFHC1,EJ	-1.65874	0.00072	0.60287	508.4205	317.3458
A_23_P50	NM_01510	Homo sapi	POGZ,SUH	-1.65874	3.32E-09	0.60287	1910.178	1142.416
A_24_P13	NM_03281	hypothetica	FLJ14627	-1.65875	3.11E-19	0.60286	3628.895	2188.856
A_23_P16	NM_00597	Homo sapi	RNF6,DKF	-1.65939	1.07E-08	0.60263	1776.16	1060.879
A_23_P25	NM_15381	Homo sapi	SNX14,RG	-1.65974	7.22E-09	0.60251	3068.601	1839.183
A_23_P10	NM_00253	ornithine d	OAZ2	-1.65994	5.79E-11	0.60243	3173.522	1908.238
A_23_P50	NM_01828	Homo sapi	PHF10,XA	-1.66034	2.63E-10	0.60229	5739.815	3462.222
A_23_P11	NM_14525	Homo sapi	STYX	-1.66077	6.13E-08	0.60213	901.7853	541.3897
A_32_P10	AK127572	Full-length	AK127572	-1.66129	1.84E-10	0.60194	1762.749	1058.818
A_23_P57	NM_02086	Homo sapi	DHX36,DD	-1.66173	1.25E-24	0.60178	4456.84	2681.475
A_24_P11	NM_00662	S-adenosy	AHCYL1,D	-1.66177	6.25E-13	0.60177	1810.403	1086.091
A_23_P25	NM_01814	Homo sapi	FAIM,FAIM	-1.66204	1.29E-20	0.60167	7021.864	4212.743
A_23_P12	NM_01646	Homo sapi	ORMDL1,D	-1.66271	2.02E-15	0.60143	7500.579	4495.255
A_24_P91	NM_05287	transcriptio	BTBD14B,	-1.66296	3.34E-12	0.60134	4545.012	2733.463
A_23_P39	NM_00674	Homo sapi	PSKH1	-1.6632	1.85E-14	0.60125	1873.491	1123.235
A_23_P21	NM_03364	Homo sapi	COL4A6,M	-1.66322	1.71E-11	0.60124	6396.864	3862.169
A_24_P40	NM_01458	ERO1 (S. c	ERO1L,ER	-1.66351	0.00015	0.60114	544.1235	330.6376
A_23_P47	NM_01533	Homo sapi	THRAP2,M	-1.66355	7.34E-07	0.60113	1794.301	1086.298
A_24_P38	NM_00069	aldehyde d	ALDH9A1,	-1.66371	3.76E-21	0.60107	5259.364	3156.496
A_24_P332532		Unknown	ENST0000	-1.66371	2.62E-22	0.60107	29336.51	17587
A_23_P38	NM_00089	Homo sapi	LTA4H	-1.66373	1.41E-26	0.60106	19332.66	11635.66
A_23_P40	AB040896	KIAA1463	KIAA1463	-1.66385	0.00005	0.60101	855.2934	498.8298
A_24_P13	NM_00320	transcriptio	TFAM,TCF	-1.66425	2.67E-12	0.60087	1957.682	1171.814
A_24_P49	BM926530	CDNA clon	BM926530	-1.66432	1.92E-07	0.60084	1106.021	674.2065
A_23_P36	NM_14579	Homo sapi	MGST1,MC	-1.66441	1.35E-26	0.60081	45184.59	27177.72
A_32_P13	NM_01741	ubiquitin sp	USP18,ISG	-1.66465	1.83E-18	0.60073	2695.693	1618.916
A_23_P28	NM_00456	Homo sapi	PIGH,GPI-	-1.66491	4.80E-12	0.60063	1206.99	726.0555
A_23_P16	NM_02249	Homo sapi	ACTR6,AR	-1.66559	1.68E-10	0.60039	4255.755	2549.842
A_24_P53	BC013405	Similar to F	LOC44014	-1.66571	1.03E-15	0.60034	2171.738	1302.511
A_23_P13	NM_00286	Homo sapi	ALDH18A1	-1.66588	2.61E-20	0.60028	5891.624	3537.449
A_24_P29	CD703575	EST20102	CD703575	-1.66592	1.99E-06	0.60027	997.5336	612.0022
A_32_P62	NM_00510	RNA bindir	RBM8A,Y1	-1.66596	1.82E-14	0.60026	1872.688	1121.789
A_23_P83	NM_00270	Homo sapi	PPP1CB,P	-1.66603	1.41E-07	0.60023	1457.397	861.8206
A_23_P25	NM_17881	LYRIC/3D3	MTDH,3D3	-1.66629	1.49E-30	0.60014	1954.582	1171.597
A_23_P60	NM_03263	hypothetica	PIGO,MGC	-1.66635	4.20E-09	0.60011	1198.151	724.5487
A_23_P14	NM_02407	Homo sapi	ZFYVE21,I	-1.66653	5.93E-28	0.60005	7019.187	4210.157
A_24_P21	NM_00690	inorganic p	PPA2,HSP	-1.66663	3.11E-28	0.60001	8720.859	5229.272
A_23_P17	NM_00726	Homo sapi	XAB1,MBD	-1.6668	8.60E-18	0.59995	6122.762	3676.819
A_23_P10987		Unknown	A_23_P10	-1.66723	0.00002	0.5998	620.7647	370.6592
A_23_P11	NM_01959	heterogene	HNRPH2,F	-1.66787	3.01E-16	0.59957	6171.925	3706.891
A_23_P70	AK058186	Hypothetic	LOC22171	-1.66791	0.00026	0.59955	783.2178	449.5626
A_23_P20	NM_02523	Homo sapi	BTBD1,C1	-1.66845	2.07E-09	0.59936	5973.521	3609.064
A_23_P56	NM_00168	ATP synth	ATP5G3,M	-1.66857	3.47E-20	0.59932	140989.7	84847.75
A_24_P92	AK025735	Thyroid ho	THRAP3	-1.66869	8.60E-09	0.59927	5139.382	3064.157
A_32_P23795		Unknown	A_32_P23	-1.66871	1.02E-26	0.59927	4848.335	2903.643
A_23_P99	NM_00602	Homo sapi	PNMA1,MA	-1.66875	9.39E-19	0.59925	10323.25	6198.097
A_23_P35	NM_00313	Homo sapi	SRI,SCN	-1.66877	9.54E-23	0.59924	28638.91	17200.32
A_23_P42	NM_00583	nuclear do	NDP52,MG	-1.66879	2.48E-07	0.59924	963.2794	585.4942
A_24_P83	NM_19884	Homo sapi	C9orf10OS	-1.66893	2.25E-11	0.59919	10046.28	6033.723
A_24_P28	NM_00249	NADH deh	NDUFB5,S	-1.66916	3.48E-11	0.5991	1449.591	872.8513
A_23_P25	NM_01504	KIAA0676	KIAA0676	-1.66939	2.63E-09	0.59902	4481.529	2683.865
A_24_P32	NM_00313	signal reco	SRPR,DP,	-1.6696	3.53E-08	0.59894	9836.717	5924.47

A_32_P148407		Unknown	A_32_P14	-1.66963	2.55E-21	0.59894	4320.967	2588.428
A_24_P67	AK023329	CDNA FLJ	AK023329	-1.66993	0.00002	0.59883	627.3925	372.2216
A_23_P14	NM_01520	SCC-112 p	SCC-112	-1.67039	2.29E-26	0.59866	4049.561	2424.408
A_32_P42	NM_19809	Homo sapi	C7orf28B,N	-1.67069	3.56E-16	0.59855	12280.52	7368.963
A_24_P23	NM_00477	UDP-Gal:b	B4GALT5,	-1.67081	0.00008	0.59851	1160.296	719.8097
A_23_P12	NM_01648	hypothetica	PPHLN1,H	-1.67086	6.08E-29	0.59849	5438.09	3253.815
A_23_P17	S63912	Heterogen	HNRPA3P	-1.67091	1.01E-24	0.59848	40730.39	24428.13
A_23_P38	NM_00144	Homo sapi	EXTL3,RE	-1.6711	4.58E-09	0.59841	1155.617	698.8095
A_32_P13	NM_00651	ret finger p	RFP,RNF7	-1.67114	6.98E-18	0.5984	5622.2	3370.017
A_23_P34	NM_00723	Homo sapi	XPOT,XPC	-1.67176	1.13E-10	0.59817	4956.398	2971.809
A_24_P791829		AK122494	THC18321	-1.67184	2.32E-08	0.59814	3721.026	2217.265
A_23_P64	NM_20127	Homo sapi	MTMR2,CN	-1.67187	3.27E-13	0.59813	2229.935	1326.891
A_24_P23	NM_00572	ARP3 (acti	ACTR3,AR	-1.67197	6.62E-22	0.5981	3793.063	2266.245
A_32_P160793		Unknown	A_32_P16	-1.67203	5.43E-10	0.59807	4628.308	2772.291
A_32_P10	BF911038	SAR1a ger	SARA1	-1.67215	7.99E-24	0.59803	22845.09	13694.95
A_32_P16	NM_00313	signal reco	SRP9,ALU	-1.67228	5.42E-11	0.59799	14412.55	8640.764
A_24_P30	NM_17508	Homo sapi	GART,AIRS	-1.67229	1.10E-07	0.59798	1113.432	658.0135
A_24_P23	NM_00566	voltage-de	VDAC3,HD	-1.6723	3.81E-18	0.59798	3672.22	2192.319
A_23_P14	D28476	Thyroid ho	TRIP12	-1.67293	1.23E-15	0.59775	5843.464	3506.195
A_23_P13	NM_01834	Homo sapi	RIOK2,FLJ	-1.67305	7.13E-12	0.59771	4377.977	2635.581
A_23_P36	NM_00681	coated ves	TMED2,P2	-1.67335	1.80E-14	0.5976	1910.587	1140.456
A_24_P39	D87447	KIAA0258	KIAA0258	-1.67359	1.61E-07	0.59752	1083.371	657.5369
A_23_P34	NM_01607	hepatoma-	HDGFRP3	-1.6739	2.93E-07	0.59741	2115.082	1245.493
A_23_P20	NM_03269	Homo sapi	MGC16385	-1.67399	2.15E-06	0.59738	848.047	499.2051
A_23_P68	NM_00385	Homo sapi	DPM1,MPD	-1.67404	2.62E-13	0.59736	18561.86	11134.51
A_23_P35	NM_01597	Homo sapi	SNX7,MGC	-1.67404	1.07E-10	0.59736	3670.028	2195.697
A_23_P35	NM_02506	Homo sapi	BXDC5,RF	-1.6741	1.36E-16	0.59734	3605.631	2153.763
A_23_P96	NM_00001	acyl-Coenz	ACADM,M	-1.67426	6.44E-15	0.59728	2404.098	1434.808
A_24_P23	D86962	Growth fac	GRB10	-1.67454	0.00007	0.59718	997.9529	574.2345
A_23_P26	NM_02484	Homo sapi	PCNT1,PC	-1.67476	2.90E-13	0.5971	5524.604	3307.84
A_23_P15	NM_00578	Homo sapi	THOC4,AL	-1.67478	2.71E-10	0.59709	28680.81	17256.18
A_32_P50	NM_00632	RAN, mem	RAN,TC4,A	-1.67489	6.20E-10	0.59705	332587.2	199995.9
A_23_P31	NM_01900	Homo sapi	XRN1,SEP	-1.67498	0.00005	0.59702	635.9907	374.1509
A_23_P40	NM_17371	Homo sapi	NM_17371	-1.67516	4.25E-20	0.59696	5813.659	3474.058
A_24_P22	NM_00213	heterogene	HNRPD,P3	-1.67527	5.34E-14	0.59692	9319.693	5581.209
A_23_P30	NM_03226	hypothetica	AE2,DKFZ	-1.67561	8.96E-07	0.5968	2085.042	1227.605
A_23_P98	NM_00027	Homo sapi	OAT,HOGA	-1.67576	2.29E-17	0.59674	5294.499	3161.99
A_24_P10	NM_20128	Homo sapi	MUTED,MU	-1.67623	0.00004	0.59658	1583.435	927.8805
A_24_P14	AB033029	Ubiquitin s	USP31	-1.67671	0.00009	0.59641	551.4194	333.1721
A_24_P83	AK002036	CDNA FLJ	AK002036	-1.6769	1.05E-22	0.59634	3405.663	2030.809
A_23_P20	NM_02469	Homo sapi	RSNL2,FL	-1.67752	4.20E-10	0.59612	2792.15	1662.73
A_32_P48066		ALU1_HUM	THC18811	-1.67771	7.84E-08	0.59605	950.0941	567.9667
A_24_P22	AK057676	RNA bindir	RBM18	-1.67791	2.85E-12	0.59598	1412.46	839.1825
A_32_P12	AK096908	CDNA FLJ	AK096908	-1.67838	1.00E-07	0.59581	1269.414	746.3793
A_24_P31	NM_00662	cyclic AMP	ARPP-19,A	-1.67845	3.07E-07	0.59579	1022.379	601.7793
A_23_P97	NM_02021	Homo sapi	RNPEP,DK	-1.67853	1.22E-07	0.59576	1736.316	1019.435
A_24_P19	NM_01457	hypothetica	MAC30	-1.67862	4.57E-24	0.59573	3149.488	1875.928
A_23_P18	NM_00274	Homo sapi	PRKCI,PKC	-1.67881	3.85E-10	0.59566	1178.533	700.1709
A_23_P38	NM_02109	solute carri	SLC5A6,SI	-1.67882	0.00098	0.59566	853.108	539.5263
A_23_P25	NM_03232	hypothetica	RNF135,L1	-1.67932	1.01E-12	0.59548	2707.317	1613.629
A_32_P18	NM_18170	hypothetica	LOC90624	-1.67951	0.00011	0.59541	628.8601	383.3932
A_23_P13	AL832747	Guanine n	GNAQ	-1.67974	1.11E-15	0.59533	3791.038	2255.677

A_23_P20	NM_02003	Homo sapi	ACCN2,AS	-1.68	8.88E-06	0.59524	654.4407	392.9821
A_23_P12	NM_00433	Homo sapi	BUB1,BUB	-1.6805	6.19E-15	0.59506	1875.844	1114.095
A_23_P94	NM_03330	chorea aca	VPS13A,C	-1.68055	4.44E-09	0.59504	843.0212	502.3709
A_23_P70	NM_02214	Homo sapi	MAGEF1,M	-1.68104	1.75E-19	0.59487	3745.657	2229.087
A_32_P61	AB040901	KIAA1468	KIAA1468	-1.68105	0.00004	0.59487	596.3617	351.1426
A_32_P36	BX436400	BX436400	BX436400	-1.68156	1.08E-17	0.59469	2414.88	1436.189
A_23_P45	NM_03338	Collagen, t	COL4A5	-1.68161	7.35E-08	0.59467	2774.625	1639.413
A_24_P38	NM_00468	myotubular	MTMR4,ZF	-1.68164	3.58E-09	0.59466	3165.703	1875.151
A_24_P17	NM_01325	muskelin 1	MKLN1,FL	-1.68198	0.00068	0.59454	531.2054	327.3834
A_24_P48	BC062438	Succinate	SDHAL2	-1.68222	1.33E-11	0.59445	2663.271	1578.132
A_24_P32	NM_13317	Homo sapi	ELMO2,CE	-1.68239	2.60E-07	0.59439	938.4545	551.2783
A_24_P229756		Unknown	ENST0000	-1.68257	1.12E-22	0.59433	44157.81	26181.56
A_24_P40	NM_01594	CGI-20 pro	C7orf20,C	-1.68259	1.61E-10	0.59432	2350.788	1393.8
A_24_P94	NM_00622	phosphotid	PITPNA,PI	-1.68265	1.17E-13	0.5943	7240.33	4311.377
A_24_P11	NM_17074	Homo sapi	ALDH5A1,	-1.68298	8.23E-07	0.59419	847.0356	510.7531
A_32_P42	BC018676	Homo sapi	BC018676	-1.683	4.84E-07	0.59418	1850.653	1085.182
A_23_P14	NM_00473	Homo sapi	SLC33A1,A	-1.68317	0.0001	0.59412	553.3021	331.308
A_23_P50	NM_01777	Homo sapi	TTC19,FLJ	-1.68322	3.92E-24	0.5941	3780.016	2244.602
A_23_P21	NM_00472	Ac-like tran	ZBED1,AL	-1.68373	0.00002	0.59392	735.8982	428.6196
A_23_P15	NM_03212	Homo sapi	HDHD2,31	-1.6841	1.29E-26	0.59379	3249.275	1929.792
A_23_P77	NM_00485	Homo sapi	PIGB,MGC	-1.68414	7.68E-06	0.59377	1057.309	610.6995
A_23_P52	NM_01813	Homo sapi	ASPM,ASP	-1.6842	1.85E-26	0.59375	7516.41	4466.659
A_23_P10	NM_13931	Homo sapi	YME1L1,F	-1.68426	1.32E-21	0.59373	6449.897	3832.09
A_24_P680548		Unknown	A_24_P68	-1.68438	2.74E-31	0.59369	51170.8	30402.22
A_23_P19	NM_00685	Homo sapi	KDELR2,E	-1.68451	8.07E-10	0.59365	11097.69	6631.254
A_32_P33	BX538015	LOC44107	DKFZp686	-1.68454	4.58E-16	0.59363	7074.777	4212.444
A_23_P14	NM_01906	Homo sapi	EML4,C2o	-1.68485	1.11E-09	0.59352	6873.377	4090.049
A_24_P30	NM_02012	SM-11044	SMBP,EP7	-1.68485	3.00E-12	0.59352	4001.536	2369.114
A_24_P58	NM_01910	beta-1,4 m	ALG1,HMT	-1.685	1.62E-10	0.59347	2102.346	1239.74
A_23_P87	NM_01791	hypothetica	FLJ20641	-1.68643	1.71E-11	0.59297	2683.73	1583.831
A_23_P13	NM_03130	Homo sapi	PUS3,FKS	-1.68716	3.12E-13	0.59271	2467.039	1460.131
A_23_P20	BX648591	Contactin 1	CNTN1	-1.68742	4.53E-31	0.59262	8622.235	5110.586
A_23_P10	NM_00599	Homo sapi	TBCD,KIA	-1.68745	0.00033	0.59261	640.846	364.8514
A_23_P12	NM_14526	homolog of	DNAJC19,	-1.68747	0.00005	0.5926	966.9994	594.9579
A_24_P26	AK001723	Nedd4 fam	NDFIP2	-1.68749	5.31E-11	0.5926	1224.055	721.8954
A_24_P27	NM_01820	vacuolar pr	VPS35,ME	-1.6881	3.24E-08	0.59238	18362.82	10959.57
A_23_P91	NM_13838	hypothetica	MGC15763	-1.68843	5.25E-12	0.59227	1860.266	1105.212
A_24_P41	NM_15303	hypothetica	FLJ32065,	-1.689	6.77E-15	0.59207	5564.862	3298.732
A_23_P13	NM_15230	Homo sapi	MRFAP1L	-1.6896	1.19E-14	0.59186	1675.49	988.0015
A_23_P94	AB018254	KIAA0711	KIAA0711	-1.68966	9.96E-13	0.59183	2400.042	1422.264
A_23_P52	NM_00452	Homo sapi	KIF11,EG5	-1.68968	2.42E-10	0.59183	2022.779	1189.898
A_24_P94	NM_01824	hypothetica	TMEM34,F	-1.68973	4.57E-13	0.59181	1471.406	867.548
A_24_P94	NM_00079	dihydrofola	DHFR	-1.6899	5.42E-11	0.59175	10455.02	6225.902
A_23_P48	NM_00282	Homo sapi	PTK9,A6,M	-1.69006	1.52E-13	0.59169	11863.71	7043.052
A_24_P92	BC015224	Adaptor-rel	AP4E1	-1.69017	1.04E-16	0.59166	3644.717	2157.59
A_23_P89	NM_00646	Homo sapi	SPAG5,MA	-1.69036	7.75E-10	0.59159	7009.71	4174.186
A_24_P19	NM_01494	KIAA0911	CLSTN1,C	-1.69069	3.02E-06	0.59147	1306.802	755.2564
A_24_P37	NM_02522	hypothetica	PRO2730,I	-1.69092	2.27E-23	0.5914	3233.125	1911.825
A_24_P471121		Unknown	A_24_P47	-1.69093	6.44E-07	0.59139	855.2897	499.7427
A_23_P97	NM_02512	hypothetica	C10orf57,F	-1.69104	6.71E-07	0.59135	1009.598	584.8387
A_23_P20	NM_03185	Neighbor o	M17S2	-1.69109	8.84E-12	0.59134	3909.996	2312.519
A_24_P19	NM_07847	Homo sapi	TM2D3,BL	-1.69113	0.00003	0.59132	730.6093	444.18

A_24_P354	NM_01906	phosphatid	MTMR12,3	-1.6915	4.10E-06	0.59119	649.536	385.4369
A_24_P945	AB033058	Discs, larg	DLG3	-1.69154	5.23E-06	0.59118	1007.505	580.9694
A_23_P408	NM_03238	Homo sapi	HPS3,SUT	-1.69155	2.38E-33	0.59117	629.239	371.2974
A_23_P71	NM_00652	Homo sapi	SLBP,HBP	-1.69165	9.69E-10	0.59114	11057.27	6568.7
A_23_P105	NM_00141	Homo sapi	EIF2B1,EIF	-1.69168	6.36E-15	0.59113	10484.4	6214.122
A_23_P40	AB018268	DDHD dom	DDHD2	-1.69178	1.03E-07	0.5911	766.6682	453.4461
A_23_P99	NM_01649	Homo sapi	MRPL51,C	-1.69215	5.56E-34	0.59097	24824.88	14674.24
A_24_P63	NM_00295	ribophorin	RPN1,RBF	-1.69223	8.23E-10	0.59094	1273.136	757.439
A_32_P164203		Unknown	THC18909	-1.69242	1.42E-09	0.59087	1385.815	825.7672
A_24_P28	NM_05285	hypothetica	ADCK2,AA	-1.69244	4.14E-10	0.59086	1439.472	856.7144
A_23_P254	NM_00452	Homo sapi	KIF2,HK2	-1.69247	3.47E-09	0.59085	2769.826	1627.713
A_32_P34404		Unknown	THC19462	-1.69347	2.04E-14	0.5905	1315.238	776.0828
A_32_P824	NM_01327	nasopharyn	ANKRD11,	-1.6935	5.56E-13	0.59049	4346.419	2572.02
A_32_P718	AK124776	CDNA FLJ	AK124776	-1.69392	1.35E-13	0.59035	2065.454	1224.455
A_32_P125568		Unknown	THC18118	-1.69421	0.00027	0.59025	457.9766	269.1844
A_32_P30785		Unknown	A_32_P30	-1.69423	0.00056	0.59024	1179.675	724.9404
A_23_P74	NM_01432	Homo sapi	OPN3,ECF	-1.69428	1.51E-06	0.59022	1086.226	627.4227
A_23_P10	NM_01479	Homo sapi	LCMT2,MC	-1.6943	4.41E-10	0.59021	2365.471	1387.664
A_24_P149	NM_00477	P311 prote	C5orf13,P3	-1.69453	8.33E-06	0.59013	740.1139	446.1113
A_23_P44	NM_01609	Homo sapi	IER3IP1,H	-1.69483	2.19E-10	0.59003	6218.746	3686.846
A_24_P22	NM_00135	aldo-keto r	AKR1C1,C	-1.69492	4.57E-23	0.59	186025.1	109984.9
A_23_P149	NM_13839	Homo sapi	C1orf37	-1.69503	1.37E-10	0.58996	5171.038	3034.502
A_24_P57	XM_37490	Homo sapi	XM_37490	-1.6954	0.00025	0.58983	517.1194	300.8207
A_23_P61	AB002370	KIAA0372	KIAA0372	-1.69566	8.79E-25	0.58974	2861.973	1693.203
A_23_P21	NM_03217	Src-associ	UTP15,FL	-1.69573	7.15E-08	0.58972	1492.68	892.5626
A_23_P86	NM_00403	annexin A7	ANXA7,SN	-1.69593	3.43E-16	0.58965	5313.25	3137.744
A_23_P70	NM_00697	Homo sapi	SLC39A7,K	-1.69614	0.00008	0.58957	827.6468	507.3427
A_24_P12	NM_01399	discoidin d	DDR1,CAK	-1.69635	1.54E-11	0.5895	4368.668	2580.661
A_23_P13	NM_00585	Homo sapi	ZMPSTE24	-1.69639	2.11E-20	0.58949	17732.56	10443.63
A_32_P58999		VAPA_HUI	THC18449	-1.69641	3.89E-19	0.58948	4933.935	2910.143
A_23_P10	NM_14516	Homo sapi	PIGM,MGC	-1.69682	7.81E-09	0.58934	1053.741	628.4282
A_23_P204	NM_02012	Homo sapi	MDM1	-1.69685	3.64E-07	0.58933	867.7092	506.0959
A_23_P75	NM_01488	Homo sapi	DCLRE1A,	-1.69732	5.28E-09	0.58916	1872.438	1094.636
A_23_P33	NM_01592	Androgen-i	APRIN	-1.69752	0.00058	0.58909	677.8125	378.3255
A_23_P57	NM_01336	Homo sapi	PCOLCE2,	-1.69767	2.18E-17	0.58904	5970.126	3522.065
A_23_P43	AL832783	Leishmano	LMLN	-1.69799	5.95E-06	0.58893	648.5151	379.6739
A_23_P38	NM_02076	raptor	raptor,KIA	-1.69844	3.05E-11	0.58878	2820.715	1657.199
A_24_P170203		Unknown	A_24_P17	-1.69848	4.73E-17	0.58876	2232.958	1313.398
A_23_P55	AB023141	Zinc finger	ZNF652	-1.69863	1.25E-15	0.58871	1692.19	997.9225
A_24_P234	NM_01660	ALEX3 pro	ARMCX3,A	-1.69878	1.96E-07	0.58866	1045.377	607.3644
A_23_P20	NM_01546	Homo sapi	NIPSNAP3	-1.69889	2.22E-16	0.58862	2716.716	1600.093
A_23_P89	NM_00611	acetyl-Coe	ACAA2,DS	-1.69913	4.15E-13	0.58854	3673.698	2162.489
A_32_P65371		Unknown	A_32_P65	-1.69928	1.66E-09	0.58848	1200.042	715.1433
A_23_P43	NM_00641	Homo sapi	SPTLC1,H	-1.69941	1.48E-13	0.58844	3282.917	1932.678
A_23_P92	NM_00094	Homo sapi	PPP3CA,C	-1.69941	1.62E-10	0.58844	4565.167	2693.838
A_23_P21	NM_01323	Homo sapi	PDCD6,AL	-1.69945	3.02E-15	0.58843	29272.01	17218.85
A_24_P36	NM_00111	adenosine	ADAR,DSH	-1.69951	4.95E-19	0.5884	7785.603	4586.51
A_24_P39	AL834278	Ubiquitin s	USP30	-1.69975	0.0002	0.58832	484.5876	290.9843
A_23_P154	NM_01246	Homo sapi	C20orf14,T	-1.69979	8.11E-13	0.58831	6374.315	3717.806
A_24_P19	NM_00245	methylthioa	MTAP,MSA	-1.70114	2.26E-08	0.58784	1148.81	667.3989
A_24_P234	NM_01786	hypothetica	C1orf56,FL	-1.70158	8.89E-17	0.58769	5796.309	3414.392
A_23_P364	NM_01816	Homo sapi	C12orf11,F	-1.70166	1.37E-06	0.58766	2712.704	1579.253

A_23_P63	BC036782	Homo sapi	CUGBP1,C	-1.70207	2.45E-18	0.58752	3934.651	2309.505
A_24_P17	NM_03194	c-Myc targ	CDCA7,JP	-1.70208	1.84E-08	0.58752	807.8692	474.8615
A_23_P14	NM_15375	Homo sapi	NAP1L5,D	-1.70213	0.0008	0.5875	643.5308	400.2428
A_23_P11	NM_01805	hypothetica	FLJ10324,	-1.70225	0.00001	0.58746	828.9408	474.4399
A_23_P99	NM_01743	Homo sapi	CPSF2,CP	-1.70231	5.87E-18	0.58744	1640.732	964.9221
A_23_P13	NM_00500	NADH deh	NDUFS1,C	-1.70237	2.91E-14	0.58742	5107.533	3010.091
A_24_P37	NM_00549	SUMO-1 a	UBA2,SAE	-1.70256	2.92E-30	0.58735	4273.493	2509.622
A_24_P92	NM_00638	DEAD/H (A	DDX17,P7	-1.70258	5.13E-09	0.58734	1593.611	927.2575
A_23_P20	AK025457	Golgi appa	GLG1	-1.7033	6.76E-25	0.5871	4095.053	2402.865
A_32_P15	NM_00313	signal reco	SRP9,ALU	-1.70333	1.05E-12	0.58709	7856.991	4563.666
A_23_P14	NM_13844	Homo sapi	C7orf30	-1.70389	8.41E-25	0.58689	6421.708	3774.423
A_24_P88	AK094413	MRNA full	AK094413	-1.70394	8.34E-06	0.58688	1285.484	785.1368
A_24_P17	AL136922	ClpX casei	CLPX	-1.70405	1.79E-12	0.58684	1210.663	712.1179
A_23_P45	NM_03305	Homo sapi	HIAT1,DKF	-1.70426	2.99E-11	0.58676	1366.224	806.0905
A_32_P11	NM_19899	N-acyl-pho	NAPE-PLD	-1.70479	5.81E-06	0.58658	1268.022	765.8047
A_23_P16	NM_14471	Homo sapi	KLHL23,M	-1.70513	0.00001	0.58647	761.3264	458.7083
A_23_P39	AK127721	KIAA0329	KIAA0329	-1.70525	0.00009	0.58643	1023.708	574.9788
A_24_P95	NM_17823	Homo sapi	TUSC3,N3	-1.70565	1.23E-22	0.58629	3116.594	1827.28
A_23_P50	BC001528	Guanine n	GNA11	-1.70572	3.16E-06	0.58626	9236.117	5452.255
A_24_P98	NM_00028	phosphofru	PFKM,MG	-1.70589	1.65E-13	0.58621	10160.43	5975.413
A_23_P347100		Unknown	A_23_P34	-1.70622	1.17E-25	0.58609	36651.39	21531.86
A_23_P33	NM_00286	Homo sapi	PYGB,MG	-1.70674	2.91E-15	0.58591	2912.603	1708.353
A_23_P31	NM_01512	Homo sapi	ALMS1,AL	-1.70682	7.59E-12	0.58589	2541.365	1483.983
A_23_P36	NM_00366	CGG triplet	CGGBP1,C	-1.70684	1.79E-08	0.58588	1358.961	786.6185
A_24_P82	BC001196	Heparan s	HS6ST1	-1.70686	9.89E-13	0.58587	3447.519	2018.591
A_24_P134319		Unknown	THC19678	-1.70717	1.23E-15	0.58577	1857.225	1089.343
A_24_P675386		Unknown	A_24_P67	-1.70748	8.88E-10	0.58566	1497.625	869.4643
A_23_P75	NM_00489	vacuolar p	VPS26A,H	-1.70751	4.53E-16	0.58565	4625.638	2717.104
A_32_P11	AK092180	Homo sapi	LOC44124	-1.7078	4.20E-11	0.58555	13599	7974.438
A_23_P21	NM_00411	Homo sapi	FGF13,FG	-1.708	1.36E-16	0.58548	3566.198	2083.241
A_23_P88	AY359110	Mesoderm	MESDC2	-1.70804	1.83E-25	0.58547	4653.988	2724.05
A_32_P60	NM_00300	succinate d	SDHD,PGL	-1.70806	2.56E-16	0.58546	2809.464	1647.466
A_23_P57	AF231919	Chromosor	C21orf108	-1.70814	6.86E-10	0.58543	838.3368	490.8824
A_24_P11	NM_00583	serine/argi	SRRM1,16	-1.70825	3.01E-13	0.58539	3593.691	2100.659
A_24_P64	NM_02409	hypothetica	ASB8,MGC	-1.70848	0.00089	0.58532	423.4185	249.2476
A_24_P268210		GNDS_HU	THC18772	-1.70875	3.50E-07	0.58522	742.1975	433.7834
A_23_P39	NM_03142	Homo sapi	C9orf58,IB	-1.7091	1.39E-12	0.5851	1314.589	767.7543
A_23_P13	AF220046	Protein ph	PPP2R2D	-1.70912	1.58E-17	0.5851	2187.854	1278.893
A_23_P13	NM_00358	neutral sph	NSMAF,FA	-1.70921	3.50E-19	0.58507	3231.536	1891.386
A_23_P11	NM_00336	Homo sapi	USP1,UBP	-1.70926	8.03E-19	0.58505	12971.92	7625.703
A_24_P290188		Unknown	ENST0000	-1.70959	7.53E-15	0.58494	1328.002	777.3055
A_23_P32	NM_00472	Homo sapi	BUB3,BUB	-1.70966	5.93E-25	0.58491	4501.79	2635.912
A_24_P71	NM_01614	dynein ligh	DNCLI1	-1.70967	6.12E-10	0.58491	1353.317	796.4557
A_24_P91	BC039896	ROD1 reg	ROD1	-1.70984	7.42E-12	0.58485	2994.399	1766.267
A_32_P35	AB023164	KIAA0947	KIAA0947	-1.71018	6.37E-07	0.58473	1041.166	594.722
A_23_P21	NM_01783	hypothetica	C9orf6,CG	-1.71026	6.84E-14	0.58471	2694.417	1570.69
A_32_P52	NM_01826	membrane	MTCBP-1,	-1.71043	3.13E-11	0.58465	2885.925	1686.758
A_24_P18	NM_00252	nucleophos	NPM1,B23	-1.7105	3.75E-23	0.58463	377140.5	219806.7
A_32_P214011		Unknown	A_32_P21	-1.7106	8.78E-07	0.58459	981.3516	586.1111
A_24_P245246		U85245 ph	THC18059	-1.71093	2.35E-11	0.58448	2992.78	1742.421
A_23_P21	BC048298	Transducin	TBL1XR1	-1.71128	3.30E-08	0.58436	1170.458	693.9336
A_32_P84	NM_00637	NS1-assoc	SYNCRIP,	-1.71158	7.69E-30	0.58425	6855.397	4003.061

A_23_P50	NM_00416	Homo sapi	SHMT1,SH	-1.71164	8.00E-20	0.58424	2550.933	1488.623
A_23_P58	NM_18183	Homo sapi	UBE2D2,U	-1.71167	1.72E-14	0.58422	3681.313	2150.062
A_23_P16	NM_01475	Homo sapi	PTDSS1,P	-1.7117	9.19E-33	0.58422	29057.52	16984.88
A_32_P23	NM_01812	hypothetica	ARMC1,Ar	-1.7118	1.06E-22	0.58418	2704.318	1579.359
A_23_P25	NM_00023	Homo sapi	SGCB,A3b	-1.71193	2.75E-08	0.58414	1723.107	997.3544
A_23_P20	D21262	Nucleolar a	NOLC1	-1.71204	4.99E-26	0.5841	4762.868	2784.934
A_23_P17	NM_00062	Homo sapi	IFNAR1,AV	-1.71219	0.00009	0.58405	545.7337	323.1581
A_23_P94	NM_01570	Homo sapi	C2orf25,CL	-1.71233	4.46E-23	0.584	11938.25	6949.904
A_23_P37	AB011090	MAX gene	MGA	-1.7124	1.27E-09	0.58397	1622.272	937.7292
A_23_P11	NM_00635	Homo sapi	UBE2E3,U	-1.71251	7.32E-32	0.58394	8513.629	4971.286
A_24_P11	NM_01602	DORA reve	DREV1,DR	-1.71269	2.51E-20	0.58388	3229.326	1885.162
A_23_P16	NM_02273	Homo sapi	MFSD1,FL	-1.71275	2.35E-11	0.58386	5789.854	3400.316
A_24_P92	NM_02121	transposon	LOC58486	-1.71305	1.05E-11	0.58375	1618.143	938.1469
A_23_P16	NM_17745	Homo sapi	PAQR3	-1.71329	6.60E-10	0.58367	1035.145	606.5586
A_23_P20	NM_00179	Homo sapi	CDC42,G2	-1.71356	4.51E-16	0.58358	11237.57	6568.278
A_24_P26	NM_01447	dimethylad	HSA9761	-1.71365	2.18E-14	0.58355	4419.017	2585.714
A_24_P786172		Unknown	THC18812	-1.7139	6.44E-06	0.58346	593.8996	346.231
A_24_P29	NM_20735	hypothetica	LOC33944	-1.7139	4.73E-10	0.58347	5965.128	3482.364
A_23_P41	NM_01843	Homo sapi	RNF130,G	-1.71421	1.86E-06	0.58336	4967.362	2913.114
A_23_P38	NM_00603	Homo sapi	CDC42BPE	-1.71466	1.35E-08	0.5832	4431.359	2597.117
A_23_P83	NM_03255	hypothetica	HIATL1,FL	-1.71483	2.23E-09	0.58315	2323.141	1356.898
A_23_P20	NM_00290	Homo sapi	RCN1,RCN	-1.71497	6.65E-18	0.5831	7579.292	4429.256
A_23_P55	NM_01546	DKFZP586	NOL11,DK	-1.715	4.93E-28	0.58309	3854.112	2247.287
A_24_P674924		ALU8_HUM	THC19984	-1.71522	5.99E-11	0.58301	1879.065	1088.478
A_24_P41	NM_00028	peroxisoma	PEX12	-1.71646	3.02E-07	0.58259	807.8517	466.0004
A_23_P25	NM_00624	Homo sapi	PPP2R5A	-1.71663	9.66E-23	0.58254	4857.193	2832.571
A_23_P91	NM_00235	Homo sapi	TACSTD1,	-1.71688	5.79E-10	0.58245	2074.14	1200.506
A_32_P19	AB002348	KIAA0350	KIAA0350	-1.71703	7.64E-07	0.5824	721.7778	423.2177
A_23_P13	NM_03290	phosphatid	PIG-Y	-1.7172	1.84E-11	0.58234	9328.944	5461.383
A_24_P77	NM_13363	Homo sapi	ROBO1,SA	-1.71755	1.19E-06	0.58222	2765.717	1598.023
A_23_P41	AF289571	Hypothetic	LOC28337	-1.71778	2.82E-10	0.58215	1458.637	842.7124
A_24_P34	AK023152	Hypothetic	FLJ10099	-1.71822	4.30E-14	0.582	3337.532	1941.546
A_24_P19	NM_01748	telomeric r	TERF1,TR	-1.71909	0.00008	0.5817	607.3902	361.8535
A_23_P41	NM_02471	Homo sapi	C5orf14,UN	-1.71937	5.18E-21	0.58161	5671.812	3305.388
A_24_P15	NM_17308	SNF2 histo	SHPRH	-1.71954	0.00002	0.58155	589.3181	343.2681
A_23_P92	NM_00621	Homo sapi	PIK3CA,PI	-1.71958	0.00056	0.58154	501.1517	301.941
A_32_P79	NM_00178	Septin 7	7-Sep	-1.71997	2.59E-10	0.58141	4901.862	2836.246
A_32_P87	NM_00591	methionine	MAT2A,MA	-1.72031	8.76E-13	0.58129	55972.47	32610.6
A_32_P14	AF419616	AG1 protei	AG1	-1.72049	0.00015	0.58123	1149.118	635.8176
A_23_P11	NM_01505	SWAP-70 p	SWAP70,H	-1.72066	1.87E-08	0.58117	984.0122	578.2192
A_23_P34	NM_01752	low density	LRP8,APO	-1.72081	1.24E-12	0.58112	1308.479	762.1141
A_32_P15	AK127081	Ubiquitin s	USP54	-1.72105	8.06E-09	0.58104	902.6093	523.9625
A_23_P33	NM_00168	Homo sapi	ATP5B,ATI	-1.72154	3.83E-11	0.58087	23430.12	13449.02
A_23_P11	NM_02410	methylosor	WDR77,ME	-1.72179	1.34E-21	0.58079	2302.432	1337.806
A_24_P36	NM_00477	UDP-Gal:b	B4GALT5,(	-1.7218	6.96E-16	0.58079	2890.197	1680.584
A_32_P21	BC022429	Hypothetic	LOC90624	-1.72182	5.57E-08	0.58078	2720.562	1567.142
A_23_P21	NM_00637	Homo sapi	SYNCRIP,p	-1.72185	9.84E-14	0.58077	6798.08	3963.607
A_24_P911648		Unknown	A_24_P91	-1.72211	2.45E-20	0.58068	2590.635	1504.946
A_23_P15	NM_00176	Homo sapi	CCT6A,CC	-1.72245	3.16E-12	0.58057	15471.09	9051.832
A_24_P89	NM_00345	zinc finger	ZNF189	-1.72266	0.0002	0.5805	480.5087	275.9689
A_24_P15	BX647930	Guanine n	GNA13	-1.72296	6.76E-07	0.5804	691.5125	399.9775
A_23_P10	NM_01476	Homo sapi	MRPL19,R	-1.72317	6.06E-19	0.58033	21210.44	12361.9

A_23_P38	NM_00413	glycogenin	GYG	-1.72318	1.20E-15	0.58032	3796.744	2207.225
A_23_P90	NM_00662	Homo sapi	POP4,RPF	-1.72319	1.75E-30	0.58032	6175.818	3586.13
A_32_P71	AL049987	FLJ40092	FLJ40092	-1.72323	2.90E-14	0.58031	1520.594	880.0129
A_23_P12	NM_01614	PTD009 pr	TRAPPC4,	-1.72354	9.83E-14	0.5802	9463.094	5515.647
A_24_P94	AF074331	Homo sapi	PAPSS2	-1.7236	3.77E-10	0.58018	1481.386	851.978
A_23_P96	AK074719	FLJ20105	FLJ20105	-1.72388	2.23E-06	0.58009	943.0518	533.4253
A_23_P21	NM_01836	Homo sapi	AGPAT5,L	-1.72402	0.00015	0.58004	640.6439	386.3225
A_24_P14	NM_01401	dexametha	DEXI,MYLI	-1.72434	2.78E-09	0.57993	7427.174	4353.249
A_23_P74	NM_00357	Homo sapi	RAD54L,H	-1.72467	4.32E-10	0.57982	1861.291	1072.532
A_24_P13	NM_03220	tumor endo	ANTXR1,A	-1.72476	1.74E-12	0.57979	1279.598	744.2526
A_24_P20	NM_02064	twisted gas	TWSG1,TS	-1.72499	0.00018	0.57971	716.3756	396.517
A_23_P15	NM_03193	Homo sapi	RAB34,RA	-1.72508	3.88E-15	0.57968	69736.78	40626.05
A_23_P31	NM_17371	Homo sapi	NM_17371	-1.72535	1.33E-23	0.57959	8774.635	5091.409
A_23_P66	NM_01572	Homo sapi	GEMIN4,H	-1.72546	6.94E-08	0.57956	5040.115	2933.277
A_24_P94	AK022820	Family with	FAM33A	-1.72547	1.41E-07	0.57955	1055.391	624.4075
A_23_P35	BX247964	Methyltrans	METTL3	-1.72589	0.00015	0.57941	618.9458	372.3619
A_23_P11	AB058729	KIAA1826	KIAA1826	-1.72609	1.47E-08	0.57934	949.8032	546.4334
A_24_P39	NM_00679	MORF-rela	MORF4L1,	-1.72658	4.86E-13	0.57918	47124.48	27457.43
A_23_P20	NM_00290	radixin	RDX	-1.72673	5.91E-14	0.57913	2475.375	1427.427
A_24_P594293		Unknown	A_24_P59	-1.72687	2.27E-21	0.57908	8986.715	5199.406
A_23_P32	NM_01654	Mst3 and S	MASK,RP6	-1.7269	3.85E-30	0.57907	4701.763	2721.855
A_23_P13	NM_00682	Homo sapi	YWHAQ,1	-1.72695	3.20E-16	0.57906	25398.5	14777.63
A_23_P16	NM_02464	Homo sapi	C14orf169,	-1.72743	1.26E-28	0.5789	5487.733	3178.349
A_24_P97	NM_01780	hypothetica	CKLFSF6,I	-1.72773	7.06E-13	0.5788	2849.069	1649.411
A_23_P38	AB014511	ATPase, C	ATP9A	-1.72826	3.95E-10	0.57862	1805.896	1034.233
A_24_P26	NM_17845	Hypothetic	MGC5428	-1.72844	3.44E-08	0.57856	1018.25	595.6559
A_32_P17	BC059174	TNF recep	TRAF3IP1	-1.72865	7.84E-10	0.57849	1728.083	989.2014
A_24_P19	NM_01812	hypothetica	TMEM33,F	-1.72913	0.00031	0.57833	518.843	292.9585
A_32_P145443		Unknown	A_32_P14	-1.72922	0.00058	0.5783	1022.791	623.9232
A_23_P16	NM_00307	Homo sapi	SMARCE1	-1.72922	3.37E-19	0.57829	8137.893	4712.177
A_23_P38	NM_00644	Homo sapi	PRPF8,PR	-1.72975	1.00E-13	0.57812	5639.985	3253.191
A_23_P12	NM_00135	DEAD/H (A	DDX3X,DB	-1.72997	2.09E-10	0.57805	7789.345	4535.073
A_24_P50	BC031327	Homo sapi	BC031327	-1.7302	0.00006	0.57797	761.0015	423.5076
A_23_P12	NM_00633	Homo sapi	TIMM17A,	-1.73036	2.34E-16	0.57792	47529.87	27568.57
A_23_P33	NM_00615	Homo sapi	NEDD4,KI	-1.73085	0.00001	0.57775	1464.474	821.9957
A_24_P35	NM_02116	hypothetica	GATAD1,C	-1.73086	4.86E-10	0.57775	1669.787	955.4764
A_23_P13	NM_00295	ribophorin	RPN1,RBF	-1.73098	9.80E-06	0.57771	6416.995	3728.577
A_24_P281468		Unknown	ENST0000	-1.73109	4.73E-12	0.57767	2833.108	1633.507
A_23_P20	NM_00610	cisplatin re	CROP,LUC	-1.73152	0.00003	0.57753	927.3585	557.9976
A_23_P69	NM_03145	Homo sapi	PARP9,BA	-1.73209	0.00037	0.57734	620.3256	376.6086
A_32_P22	AL832747	Guanine nu	GNAQ	-1.73218	3.37E-11	0.57731	4022.416	2323.249
A_24_P99	NM_00227	RAN bindir	RANBP5	-1.73281	2.57E-18	0.5771	1813.648	1046.852
A_32_P59	NM_00300	succinate d	SDHD,PGL	-1.73326	4.79E-18	0.57695	2248.966	1299.004
A_23_P14	NM_01826	membrane	MTCBP-1,	-1.73338	6.87E-21	0.57691	3097.48	1783.802
A_23_P21	NM_01475	Signal pep	SPCS2	-1.73359	1.53E-12	0.57684	7232.93	4143.288
A_23_P20	NM_02281	Homo sapi	PER2,FAS	-1.73364	6.83E-08	0.57682	795.1007	460.5994
A_32_P125558		Unknown	A_32_P12	-1.73372	2.07E-11	0.57679	3074.248	1799.482
A_24_P39	AF262027	Homo sapi	eIF-5A2	-1.7339	5.13E-12	0.57673	1961.694	1125.406
A_24_P31	AK090730	Solute carr	SLC35A2	-1.73429	8.24E-12	0.5766	2307.88	1337.014
A_23_P20	AK027500	Leucine ric	LRRN6A	-1.73449	6.26E-14	0.57654	4282.486	2468.837
A_23_P45	NM_01230	Homo sapi	EPB41L3,4	-1.73455	2.08E-13	0.57652	1582.036	906.4446
A_24_P27	NM_02470	hypothetica	ASB13,FL	-1.73455	6.72E-26	0.57652	3982.531	2296.481

A_24_P49	BC013389	LOC44015	BC013389	-1.73467	0.00026	0.57648	1212.444	666.2763
A_24_P850336		Unknown	A_24_P85	-1.73491	5.78E-22	0.5764	36243.41	20901.78
A_32_P20	BG178211	Mesenchyr	BG178211	-1.73505	2.11E-20	0.57635	2821.292	1624.506
A_24_P7380		Unknown	ENST0000	-1.73562	6.32E-28	0.57616	15127.67	8704.311
A_23_P36	NM_13877	hypothetica	LOC93081	-1.7359	3.38E-16	0.57607	5726.822	3314.723
A_23_P58	NM_03137	Homo sapi	HNRPD,P3	-1.73656	1.34E-15	0.57585	44017.55	25432.15
A_24_P139761		Unknown	A_24_P13	-1.73668	0.00005	0.57581	920.7371	553.4219
A_24_P64	BX538015	LOC44107	DKFZp686	-1.73685	2.64E-20	0.57575	3992.728	2297.114
A_23_P21	NM_00209	Homo sapi	GRSF1	-1.7374	3.38E-13	0.57557	5050.979	2920.082
A_23_P12	NM_00137	Homo sapi	DNCH1,p2	-1.73794	2.45E-08	0.57539	13550.99	7784.17
A_23_P50	NM_01749	WD repeat	WDR1,AIP	-1.73808	6.23E-11	0.57535	2993.158	1716.68
A_32_P16	NM_00506	stearoyl-Co	SCD,SCD1	-1.73816	2.24E-23	0.57532	14588.22	8386.161
A_23_P18	NM_00436	Homo sapi	CLGN	-1.73829	1.48E-19	0.57528	2089.819	1203.233
A_24_P24	NM_01560	Homo sapi	WDR21A,V	-1.73853	4.13E-10	0.5752	2512.048	1438.696
A_23_P94	NM_00219	aconitase 1	ACO1,IRP	-1.73879	4.39E-12	0.57511	1475.797	854.3663
A_23_P25	NM_00503	Homo sapi	PLS3,T-PL	-1.73881	1.57E-23	0.57511	5767.503	3314.602
A_23_P12	NM_01436	Homo sapi	SACS,ARS	-1.73885	1.77E-24	0.57509	4423.548	2545.681
A_24_P10	XM_11662	Homo sapi	ENST0000	-1.73893	1.03E-26	0.57507	4284.5	2465.224
A_24_P36	NM_02468	hypothetica	C17orf42,F	-1.73906	2.85E-25	0.57502	2861.731	1645.551
A_24_P73	NM_00168	ATPase, C	ATP2A2,D	-1.73925	4.60E-22	0.57496	2485.388	1428.241
A_23_P12	NM_00542	tumor prote	TP53BP2,5	-1.73947	4.13E-11	0.57489	1239.436	716.8316
A_23_P11	NM_00728	Homo sapi	GABARAP	-1.7397	6.04E-28	0.57481	8210.993	4712.216
A_23_P48	NM_00682	transmemb	CKAP4,p63	-1.73988	1.89E-20	0.57475	6596.326	3800.895
A_23_P15	AL834362	GTPase ac	GARNL1	-1.73989	1.78E-10	0.57475	1224.101	699.6185
A_24_P19	AL713633	Nudix (nuc	NUDT1	-1.74001	5.94E-10	0.57471	1424.405	808.6996
A_23_P16	NM_01613	Homo sapi	RAB10	-1.74017	2.64E-16	0.57466	7810.644	4499.353
A_24_P14	BC007568	capicua (D	CIC	-1.74042	9.00E-12	0.57457	1252.852	715.5629
A_24_P34	NM_01817	hypothetica	GOLPH3L,	-1.74071	8.61E-06	0.57448	1325.228	739.5568
A_23_P31	NM_13828	Homo sapi	C14orf147,	-1.74073	1.42E-19	0.57447	15038.47	8614.328
A_23_P14	NM_01482	Homo sapi	TOMM70A	-1.74077	1.14E-06	0.57446	1138.174	638.1543
A_24_P27	NM_00248	Nijmegen b	NBN,ATV,N	-1.7411	7.09E-10	0.57435	1113.758	635.1075
A_24_P58759		Unknown	A_24_P58	-1.74136	5.22E-27	0.57426	21704.2	12483.79
A_23_P12	NM_00263	Homo sapi	PGD,6PGD	-1.74161	2.91E-14	0.57418	3915.808	2253.761
A_24_P22	XM_29219	similar to b	LOC34165	-1.74261	9.03E-20	0.57385	17102.22	9844.498
A_23_P12	NM_01219	Homo sapi	FXC1,Tim9	-1.74302	8.26E-12	0.57372	1363.013	778.61
A_24_P30	BC002759	Hypothetica	FLJ20489	-1.74304	1.92E-21	0.57371	8632.527	4952.553
A_24_P889103		BC015704	THC18458	-1.74308	0.0007	0.5737	423.8046	240.907
A_23_P16	NM_00736	Homo sapi	NID2	-1.7432	0.00001	0.57366	1811.435	1018.223
A_23_P80	NM_00318	Homo sapi	TAF4,TAF2	-1.74324	5.20E-10	0.57364	1301.398	738.3615
A_23_P15	NM_00611	Homo sapi	PECI,DRS	-1.74398	6.30E-16	0.5734	11007.49	6354.209
A_23_P37	BC001774	Platelet-act	PAFAH1B2	-1.74428	2.94E-26	0.5733	3739.256	2142.765
A_23_P24	NM_01908	Homo sapi	CCNJ,bA6	-1.74445	7.17E-11	0.57325	1160.847	660.5743
A_23_P53	NM_00392	Homo sapi	TIMELESS	-1.74455	1.45E-24	0.57321	10449.15	5999.348
A_24_P41	AB014599	Bicaudal D	BICD2	-1.74471	2.89E-15	0.57316	2270.945	1305.464
A_23_P41	NM_02215	Homo sapi	SLC39A8,E	-1.74477	1.90E-12	0.57314	9855.439	5684.831
A_24_P79	U69195	Transcribe	LOC51580	-1.74482	9.19E-12	0.57313	1070.41	612.98
A_23_P15	NM_01814	hypothetica	C8orf35,FL	-1.74485	6.68E-16	0.57312	3764.225	2162.22
A_23_P12	NM_02408	Homo sapi	ASB9,MGC	-1.74549	1.86E-06	0.57291	1615.359	906.2255
A_24_P568645		Unknown	A_24_P56	-1.74559	2.05E-23	0.57287	2286.396	1311.224
A_23_P99	NM_14534	Homo sapi	PDCD4,H7	-1.74578	8.90E-06	0.57281	798.3446	444.3247
A_23_P37	AB040942	KIAA1509	KIAA1509	-1.74658	0.00057	0.57255	509.6505	281.6898
A_24_P15	NM_01754	hypothetica	H41	-1.74684	1.23E-30	0.57246	4428.474	2535.917

A_23_P21	NM_19882	Ras-relate	RAC1	-1.74729	7.40E-36	0.57231	12309.38	7046.862
A_23_P15	NM_01594	Homo sapi	ATP6V1H,	-1.74744	1.13E-08	0.57226	3276.408	1859.93
A_24_P41	NM_00210	H2A histon	H2AFZ,H2	-1.74747	2.18E-36	0.57226	88759.38	50806.18
A_23_P39	NM_02245	Ring finger	RFWD2	-1.74782	1.93E-17	0.57214	4482.578	2563.31
A_24_P41	NM_01589	pre-mRNA	CDC40,EH	-1.74791	2.04E-08	0.57211	1860.866	1051.233
A_32_P30	BM989848	Transcribe	BM989848	-1.74817	2.50E-15	0.57203	3476.582	1988.332
A_23_P20	NM_00229	Homo sapi	LAMC1,LA	-1.7485	8.39E-18	0.57192	4099.217	2348.918
A_23_P14	NM_00364	Homo sapi	SLC27A2,	-1.74899	5.18E-15	0.57176	3561.733	2034.213
A_23_P21	NM_02433	Chromosor	CXorf53	-1.749	1.20E-29	0.57175	8184.701	4685.903
A_23_P45	NM_01450	Homo sapi	HTATSF1,	-1.74928	1.59E-35	0.57166	15927.12	9110.098
A_32_P21	NM_20339	Homo sapi	E2F7	-1.7502	1.14E-20	0.57136	4261.89	2440.029
A_23_P37	NM_00121	Homo sapi	C1QBP,p3	-1.75024	7.85E-25	0.57135	147551	84500.58
A_23_P10	NM_01644	Homo sapi	DTL,RAMF	-1.75028	6.60E-24	0.57134	3365.225	1931.058
A_24_P35	AK054681	Chromosor	C22orf13	-1.75036	1.74E-21	0.57131	2795.794	1597.577
A_24_P72	AF317228	Spindlin	SPIN	-1.75065	6.35E-18	0.57122	2172.971	1240.719
A_24_P16	AK025511	ATPase, C	ATP11A	-1.75109	1.58E-08	0.57107	1214.281	682.2289
A_23_P12	NM_00340	Homo sapi	YWHAB,H	-1.75142	6.02E-13	0.57097	17269.59	9916.287
A_23_P14	NM_00337	Homo sapi	VDAC1,PC	-1.75216	1.36E-36	0.57072	49452.48	28235.77
A_23_P39	NM_00706	Homo sapi	PWP1,IEF-	-1.7527	2.00E-16	0.57055	28756.99	16426.73
A_24_P94	NM_00630	SMC1 (stru	SMC1L1,S	-1.75295	5.09E-07	0.57047	702.8953	398.5705
A_24_P32	NM_00438	CREB bind	CREBBP,C	-1.75313	5.76E-09	0.57041	1192.653	670.528
A_24_P94	D13642	Splicing fac	SF3B3	-1.75323	6.04E-06	0.57038	1116.665	616.1476
A_23_P40	NM_00340	Homo sapi	XPO1,CRM	-1.75334	2.66E-13	0.57034	9334.506	5351.141
A_23_P15	NM_01333	Homo sapi	ALG5,bA4	-1.75344	3.73E-15	0.57031	12278.83	7025.162
A_23_P14	NM_00572	Homo sapi	TSPAN3,T	-1.75346	1.07E-15	0.5703	7247.312	4155.703
A_24_P40	NM_01466	KIAA0140	FAM53B,K	-1.75374	1.89E-19	0.57021	2002.296	1142.492
A_23_P17	NM_00131	chromoson	CSE1L,CA	-1.75375	1.00E-23	0.57021	19239.54	11004.22
A_24_P26	NM_18275	Homo sapi	WDR72,FL	-1.75383	0.00026	0.57018	515.5098	287.7701
A_23_P88	NM_01475	Homo sapi	DLG7,DLG	-1.75393	8.28E-34	0.57015	4619.902	2633.955
A_23_P21	NM_01759	rotavirus X	ZC3H7B,R	-1.75407	7.75E-06	0.5701	1455.088	807.9667
A_23_P14	NM_00636	Homo sapi	SEC23A,M	-1.75426	4.78E-17	0.57004	9231.281	5285.936
A_24_P55	AK094583	Hypothetic	LOC33974	-1.7543	0.00001	0.57003	670.111	373.6357
A_23_P60	NM_00307	SWI/SNF r	SMARCA2	-1.75448	0.0003	0.56997	497.2122	277.4301
A_23_P53	NM_20161	IKK interac	IKIP,FLJ31	-1.75451	2.71E-08	0.56996	1999.438	1128.639
A_23_P15	NM_00392	Homo sapi	MBD2,DM	-1.75491	4.52E-11	0.56983	5382.685	3059.559
A_32_P181107		JC5193 nu	THC19381	-1.75497	6.28E-06	0.56981	830.2985	486.5567
A_24_P34	NM_00048	alpha thala	ATRX,SHS	-1.75518	5.78E-09	0.56974	1023.656	577.7763
A_23_P65	NM_00470	Homo sapi	CCNB2,Hs	-1.75527	7.14E-14	0.56971	12806.82	7309.883
A_23_P11	NM_00473	solute carri	SLC16A7,M	-1.75597	0.00033	0.56949	471.5252	264.5497
A_23_P59	NM_01404	Homo sapi	MRPS18B,	-1.75609	1.48E-13	0.56945	20430.68	11703.59
A_23_P85	NM_01423	Homo sapi	GNPAT,DA	-1.7567	2.74E-26	0.56925	4402.895	2508.714
A_23_P21	NM_00569	Homo sapi	ABCF2,AB	-1.75682	4.92E-15	0.56921	1449.134	826.7192
A_32_P21	AK025695	Transmem	TMED8	-1.75726	8.13E-15	0.56907	2510.96	1423.578
A_23_P41	NM_15226	hypothetica	MGC17943	-1.75762	9.66E-15	0.56895	6916.157	3952.019
A_24_P46093		Unknown	THC18076	-1.75763	1.50E-09	0.56895	3690.707	2073.503
A_23_P33	NM_00466	Homo sapi	HERC2,jdf	-1.75778	4.82E-15	0.5689	1538.664	873.0399
A_23_P11	NM_02237	ATP-deper	TOR3A,AD	-1.75798	2.93E-36	0.56884	37036.34	21062.07
A_23_P80	NM_02289	Homo sapi	ACTR8,AR	-1.75803	0.00067	0.56882	403.6235	231.0927
A_23_P65	NM_03281	hypothetica	FLJ14624,	-1.75831	6.20E-13	0.56873	1208.308	685.6496
A_24_P49	AK091503	Ribosomal	RPS6KB1	-1.75841	1.60E-06	0.5687	882.6818	490.7269
A_24_P157053		AD17_HUM	THC18575	-1.75855	5.58E-12	0.56865	2102.982	1200.61
A_24_P49	NM_18165	hypothetica	LOC28401	-1.75887	0.00021	0.56855	483.4354	271.598

A_32_P97	BX640888	Glypican 6	GPC6	-1.75889	6.34E-08	0.56854	1455.666	811.7885
A_32_P10856		Unknown	THC18928	-1.7589	7.17E-13	0.56854	4401.646	2497.578
A_23_P84	NM_03217	hypothetica	TOMM40L	-1.75925	1.18E-13	0.56842	2689.758	1525.275
A_24_P33	NM_00121	carbonic ar	CA12,CAX	-1.75997	2.98E-11	0.56819	1616.452	912.7804
A_23_P13	NM_00091	Homo sapi	OXTR,OT-	-1.7602	0.00008	0.56812	500.6335	285.7477
A_23_P63	NM_00658	chromoson	C1orf2,CO	-1.76115	4.59E-15	0.56781	8339.775	4755.974
A_23_P45	NM_00582	heterogene	HNRPR,HN	-1.76115	1.84E-23	0.56781	10779.18	6135.14
A_24_P41	NM_18275	Homo sapi	MCM10,CN	-1.76161	1.43E-08	0.56766	1419.172	794.3999
A_23_P15	NM_01865	bladder ca	SLC35E3,E	-1.76197	6.72E-10	0.56755	975.9957	550.3934
A_23_P20	NM_01623	Homo sapi	ANAPC5,A	-1.76252	3.48E-16	0.56737	33276.21	18910.71
A_23_P15	NM_01210	Homo sapi	ARL2BP,B	-1.76304	1.61E-13	0.5672	9524.795	5387.682
A_24_P71	NM_17371	Homo sapi	NM_17371	-1.76358	6.65E-11	0.56703	5996.947	3401.09
A_23_P12	NM_01822	Homo sapi	SMU1,BWI	-1.76365	0.00018	0.567	543.1642	317.9122
A_24_P91	NM_00652	t-complex-	TCTE1L,T	-1.76384	4.48E-17	0.56695	2774.717	1573.656
A_32_P42	AA521057	Transcribe	GGA2,VEA	-1.76416	4.36E-07	0.56684	687.6851	388.8738
A_24_P10	NM_00539	protein kin	PRKAB2,M	-1.76436	6.15E-25	0.56678	2248.26	1274.747
A_23_P26	AB037809	Zinc finger	ZNF319	-1.76454	0.00001	0.56672	800.7349	438.7878
A_23_P13	NM_00289	Homo sapi	RBM4,LAR	-1.76475	2.49E-27	0.56665	2476.862	1403.512
A_23_P32	NM_17379	Homo sapi	PDZK8,FL	-1.76481	0.00085	0.56663	526.6205	315.3473
A_23_P40	NM_01624	Homo sapi	DHRS8,PA	-1.76546	1.30E-13	0.56642	1249.963	708.8103
A_23_P16	NM_00556	Lysosomal	LAMP1	-1.76562	6.43E-25	0.56637	40839.05	23168.35
A_24_P94	AL833978	HP1-BP74	HP1-BP74	-1.76579	9.20E-12	0.56632	3210.677	1814.235
A_24_P41	NM_01504	Golgi-asso	GGA2,VEA	-1.76607	7.08E-10	0.56623	936.5692	528.9216
A_24_P59	BC019938	Hypothetic	LOC33926	-1.76624	9.60E-17	0.56617	1578.081	894.5795
A_24_P19	NM_00721	ring finger	RNF2,BAP	-1.76657	0.00026	0.56607	752.7492	401.3226
A_23_P20	NM_00655	Homo sapi	KHDRBS1	-1.76659	3.18E-16	0.56606	14820.19	8425.509
A_23_P10	NM_01794	Homo sapi	NM_01794	-1.76663	1.10E-14	0.56605	7666.863	4335.179
A_23_P40	NM_00282	Homo sapi	PTPN3,PT	-1.76668	0.00008	0.56604	716.0243	424.3102
A_24_P85	AK092312	Zinc finger,	ZFYVE20	-1.76676	9.11E-13	0.56601	1525.462	857.5915
A_23_P18	NM_00720	mitochondr	MRPL3,MF	-1.76712	8.48E-37	0.56589	11516.47	6515.032
A_23_P24	NM_00547	potassium	KCNE3,HC	-1.76723	0.00016	0.56586	506.3297	292.1833
A_23_P78	NM_00238	Homo sapi	MCM3,HC	-1.76774	6.07E-24	0.5657	19990.53	11339.52
A_23_P11	NM_00175	Homo sapi	CBFB,PEB	-1.76776	3.04E-11	0.56569	5627.711	3182.825
A_23_P30	D86979	KIAA0226	KIAA0226	-1.76826	1.99E-10	0.56553	1418.632	793.9353
A_23_P15	NM_00002	adenylosuc	ADSL	-1.76839	8.56E-19	0.56549	3126.557	1768.029
A_24_P57	AF086448	RNA U, sm	PHAX	-1.76863	1.30E-17	0.56541	2800.636	1582.356
A_23_P40	NM_00394	Homo sapi	USP13,ISC	-1.76869	5.09E-11	0.56539	1159.775	651.1606
A_23_P53	NM_00379	Homo sapi	EED,HEED	-1.76891	1.70E-14	0.56532	6287.643	3575.216
A_32_P54	NM_00176	chaperonin	CCT6A,CC	-1.76896	5.47E-18	0.5653	14495.59	8193.291
A_23_P25	NM_01844	Homo sapi	CAND1,TIF	-1.76927	3.62E-07	0.5652	1080.097	627.8541
A_32_P48	NM_20339	similar to R	RBM12B,M	-1.76941	1.89E-06	0.56516	828.7866	458.4981
A_23_P36	NM_03067	amino acid	SLC38A1,A	-1.7696	2.63E-27	0.5651	7681.505	4338.752
A_23_P16	NM_01612	Homo sapi	KCTD3,MC	-1.76967	1.28E-19	0.56508	12571.77	7121.929
A_23_P11	NM_01813	Homo sapi	C10orf3,FL	-1.76978	5.15E-34	0.56504	4884.76	2760.456
A_23_P30	NM_01482	KIAA0801	DDX46,MC	-1.76981	1.36E-24	0.56503	2547.359	1438.804
A_23_P21	NM_01871	testis speci	TSGA14,C	-1.77004	1.79E-08	0.56496	829.4908	465.2809
A_23_P10	BC000525	Glutamic-o	GOT2	-1.77054	3.73E-26	0.5648	9508.183	5379.246
A_23_P60	NM_00644	Homo sapi	SMC2L1,C	-1.77097	1.85E-08	0.56466	829.1439	464.8582
A_32_P21	NM_03216	hypothetica	WDR75,FL	-1.77108	4.14E-27	0.56463	3309.372	1868.261
A_23_P12	NM_00146	thyroid aut	XRCC6,ML	-1.77109	8.86E-18	0.56462	108454	61533.66
A_24_P94	NM_00657	guanine nu	GNA13,G1	-1.77132	0.00005	0.56455	1511.859	822.5463
A_23_P11	NM_01630	protein x 0	ATPBD1C,	-1.77137	1.48E-15	0.56454	4149.979	2352.545

A_24_P141736		AMP2_HUI	THC18455	-1.7721	7.16E-06	0.5643	911.2162	498.1266
A_32_P203300		IF4E_HUM	THC18793	-1.77236	5.71E-08	0.56422	871.053	498.9323
A_24_P19	NM_00325	Tight juncti	TJP1	-1.77239	3.75E-24	0.56421	3122.582	1762.251
A_24_P118953		Unknown	A_24_P11	-1.7725	0.00002	0.56418	577.7199	322.1968
A_23_P48	NM_02405	hypothetica	MGC5576	-1.77253	1.41E-19	0.56416	24931.14	14141.38
A_24_P92	BC032537	Hypothetic	LOC63929	-1.77272	1.51E-06	0.5641	731.3131	405.7311
A_24_P174063		Unknown	A_24_P17	-1.77287	0.00014	0.56406	982.8868	524.5774
A_23_P16	NM_00149	Homo sapi	GDI2,RAB	-1.7732	2.24E-17	0.56395	5486.036	3073.568
A_23_P56	NM_01557	Homo sapi	FAM61A,C	-1.77336	6.32E-14	0.5639	2873.035	1619.752
A_24_P34	NM_00484	class I cyto	IL27RA,CF	-1.77397	8.10E-06	0.56371	628.2145	347.607
A_24_P816844		ALU1_HUM	THC19686	-1.77444	2.07E-06	0.56356	1087.057	593.4672
A_23_P33	NM_00727	Homo sapi	FKBP9,FK	-1.77488	3.87E-29	0.56342	3712.028	2090.162
A_24_P24	NM_01328	methionine	MAT2B,TG	-1.77488	5.20E-20	0.56342	9635.407	5449.038
A_24_P91	NM_01869	hypothetica	NXT2,P15-	-1.77512	2.99E-23	0.56334	4194.813	2364.123
A_32_P44	NM_00556	lactate deh	LDHA,LDH	-1.77521	6.15E-28	0.56331	6873.971	3863.813
A_23_P13	NM_15293	Homo sapi	GLT8D1,A	-1.77618	1.73E-16	0.56301	2722.322	1528.171
A_23_P68	AL137442	Chromosor	C20orf177	-1.77675	1.13E-08	0.56283	3232.583	1815.71
A_23_P47	NM_00556	Homo sapi	LDHA,LDH	-1.77695	3.51E-37	0.56276	39055.48	21995.04
A_23_P11	NM_00211	Homo sapi	CFHL1,CF	-1.77725	1.74E-29	0.56267	4446.817	2501.175
A_24_P84	NM_00411	flap structu	FEN1,MF1	-1.77752	1.12E-20	0.56258	2121.283	1195.715
A_23_P20	NM_00025	5-methyltet	MTR	-1.77782	0.00001	0.56249	1035.137	603.1587
A_23_P40	NM_01846	uncharacte	HT008,KIA	-1.77791	3.30E-16	0.56246	1711.574	959.0023
A_23_P12	BC029424	Similar to F	LOC49386	-1.77825	3.60E-40	0.56235	3999.031	2244.676
A_24_P24	NM_01226	heparan su	HS2ST1,FI	-1.77849	0.0001	0.56227	495.7317	277.5273
A_32_P63	NM_00043	3-oxoacid (	OXCT1,OX	-1.77862	1.74E-13	0.56223	1496.944	837.2987
A_32_P20	AK021443	CDNA FLJ	AK021443	-1.77873	1.15E-06	0.5622	690.3483	383.871
A_24_P11	NM_02466	nuclear rec	TBL1XR1,(	-1.77874	1.98E-10	0.56219	2938.724	1647.547
A_24_P19	NM_02046	KIAA1387	SMEK2,KI	-1.77875	4.02E-06	0.56219	1235.699	673.1374
A_23_P20	NM_03281	hypothetica	FLJ14627	-1.77903	7.33E-10	0.5621	1102.234	614.3195
A_23_P29	NM_00534	Homo sapi	RBPSUH,c	-1.77908	2.98E-16	0.56209	3145.231	1764.084
A_23_P11	NM_00257	Homo sapi	PRDX1,PA	-1.77921	1.82E-06	0.56205	115671.2	65012.53
A_24_P96	AF008915	Ecotropic v	EVI5	-1.77921	0.00003	0.56205	604.8179	332.1705
A_32_P96	AB020684	KIAA0877	KIAA0877	-1.7794	1.90E-15	0.56199	1252.27	704.1301
A_23_P12	NM_00029	Homo sapi	PGK1,PGK	-1.77959	1.99E-14	0.56193	107328.1	60613.31
A_23_P12	NM_03286	Homo sapi	NUDCD1,(	-1.77962	1.05E-28	0.56192	2106.876	1183.806
A_23_P11	NM_00334	ubiquitin-co	UBE2N,UB	-1.7799	0.00004	0.56183	684.1176	399.1772
A_32_P184746		Unknown	THC19927	-1.78001	0.00075	0.56179	409.8868	232.9655
A_23_P42	NM_01555	Homo sapi	SS18L1,CF	-1.78031	0.00017	0.5617	629.2618	337.7781
A_24_P65	AK094066	Nuclear fac	NFIB	-1.78046	7.54E-22	0.56165	1923.688	1081.233
A_23_P15	NM_02216	Homo sapi	XYLT2,XT	-1.78069	6.00E-17	0.56158	3775.676	2117.175
A_23_P62	NM_01815	Homo sapi	NUDT11,A	-1.78099	8.24E-10	0.56148	1190.49	660.0902
A_23_P17	NM_02458	hypothetica	SCRN3,SE	-1.781	0.00003	0.56148	779.031	421.2864
A_23_P13	NM_01437	Homo sapi	RNF11,CG	-1.78135	5.83E-12	0.56137	3033.247	1700.09
A_24_P98	NM_03322	PC4 and S	PSIP1,p52	-1.7817	9.40E-11	0.56126	2867.3	1600.439
A_32_P14	AK026351	Protein kin	PRKAR2A	-1.78173	1.06E-11	0.56125	12197.69	6888.194
A_24_P31	NM_13845	Homo sapi	C6orf68,M	-1.78216	4.46E-15	0.56112	1845.923	1030.951
A_24_P32	NM_00521	diaphanous	DIAPH1,DI	-1.78235	3.05E-11	0.56106	25716.02	14521.03
A_24_P19	NM_00540	v-ral simiar	RALA,RAL	-1.78243	7.97E-09	0.56103	1496.053	827.8139
A_23_P94	NM_00209	general tra	GTF2E2,FI	-1.78258	6.20E-23	0.56098	2176.49	1221.346
A_24_P90	NM_00546	heterogene	HNRPDL,H	-1.7831	4.80E-21	0.56082	4354.976	2443.377
A_24_P350196		OK/SW-CL	ENST0000	-1.78313	3.09E-22	0.56081	12950.47	7276.171
A_23_P16	NM_00032	Homo sapi	QDPR,DHI	-1.7832	2.26E-12	0.56079	3054.477	1711.639

A_23_P25	NM_00094	Homo sapi	PRIM1,p49	-1.78323	1.74E-43	0.56078	4861.358	2729.208
A_23_P64	NM_00481	Homo sapi	DDX23,prp	-1.78349	3.34E-39	0.5607	8456.338	4741.102
A_24_P13	NM_00114	autocrine n	AMFR,GP7	-1.78373	9.10E-20	0.56062	1399.264	784.9171
A_24_P77	NM_18278	Homo sapi	PAIP1,MG	-1.78422	9.25E-15	0.56047	5552.817	3116.986
A_23_P81	NM_03057	hypothetica	NDFIP1,N4	-1.78429	3.26E-21	0.56045	4618.108	2592.835
A_23_P12	NM_19941	Homo sapi	PRCP,PCP	-1.78496	5.12E-28	0.56024	4115.513	2307.519
A_24_P20	U69127	Far upstrea	FUBP3	-1.78502	9.74E-16	0.56022	3890.555	2183.208
A_32_P10	AB046816	KIAA1596	KIAA1596	-1.78507	3.12E-13	0.5602	2224.619	1238.584
A_24_P39	NM_19856	FLJ44216	FLJ44216,	-1.78536	8.97E-13	0.56011	1080.229	605.12
A_24_P22	AF111845	3-phosphoi	PDPK1	-1.78551	7.19E-11	0.56006	1315.053	730.4323
A_24_P410686		A42286 EF	THC18334	-1.78553	4.70E-27	0.56006	9616.655	5372.705
A_23_P33	NM_01546	Homo sapi	GEMIN5,D	-1.78558	6.36E-11	0.56004	1370.102	761.0943
A_23_P20	NM_02440	Homo sapi	NOTCH2,h	-1.78558	1.79E-13	0.56004	5284.321	2950.246
A_32_P18	NM_00543	DNA segm	CCDC6,H4	-1.78625	1.54E-15	0.55983	15541.98	8761.152
A_23_P22	D00022	Homo sapi	F1 beta su	-1.7866	2.24E-12	0.55972	184235.8	103660
A_23_P211899		Unknown	THC19455	-1.7871	1.52E-14	0.55957	2371.392	1320.196
A_23_P55	NM_00538	Homo sapi	NCL,C23,F	-1.78754	3.98E-14	0.55943	134423.5	75709.69
A_24_P21	AF130080	Nuclear ub	NUCKS	-1.78775	9.27E-06	0.55936	661.178	378.7496
A_23_P10	NM_02494	NEFA-inter	NIP30,CDA	-1.78814	2.92E-13	0.55924	1276.406	711.678
A_24_P19	NM_00682	tyrosine 3-r	YWHAQ,1	-1.78908	5.61E-26	0.55895	58005.02	32538.15
A_24_P312381		Unknown	A_24_P31	-1.7896	1.10E-06	0.55878	2855.753	1598.564
A_24_P31	NM_14530	similar to R	LOC20245	-1.78968	9.41E-14	0.55876	4936.04	2758.949
A_32_P86	AB032986	Homo sapi	KIAA1160	-1.7897	0.00071	0.55875	733.1622	378.3269
A_24_P10	NM_02459	hypothetica	FLJ13154	-1.78977	2.62E-13	0.55873	14054.21	7889.192
A_23_P20	NM_02020	Homo sapi	PRTFDC1,	-1.78984	2.92E-16	0.55871	1546.361	861.6904
A_23_P11	NM_01630	Homo sapi	CMPK,UM	-1.78986	2.19E-09	0.5587	2421.129	1346.783
A_23_P32	NM_08072	Homo sapi	SRXN1,Np	-1.78993	4.10E-39	0.55868	32253.14	18015.52
A_24_P89	D79986	BCL2-asso	BCLAF1	-1.79019	2.77E-26	0.5586	8323.648	4661.538
A_23_P23	NM_00210	Homo sapi	H3F3A,H3	-1.79047	1.08E-28	0.55851	111812.2	62286.35
A_24_P97	AB033076	Likely hom	KIDINS22	-1.79084	9.17E-14	0.5584	1496.407	831.8786
A_23_P35	NM_00024	met proto-c	MET,HGFF	-1.79106	8.09E-18	0.55833	6739.557	3786.734
A_23_P11	NM_02066	Homo sapi	MRS2L,HF	-1.79109	2.03E-15	0.55832	1593.067	886.6754
A_23_P13	NM_00136	Homo sapi	DKC1,DKC	-1.79136	5.12E-32	0.55824	19026.21	10627.38
A_23_P81	NM_00573	conserved	CTDSP2,C	-1.79137	6.03E-29	0.55823	4521.95	2521.725
A_23_P38	NM_02429	hypothetica	PHF23,MG	-1.79145	5.63E-15	0.55821	2314.97	1290.656
A_24_P12	AB006624	KIAA0286	KIAA0286	-1.79215	4.79E-10	0.55799	1140.374	625.9579
A_23_P19	NM_01832	hypothetica	C6orf64,FL	-1.7922	0.00033	0.55797	438.7728	242.7793
A_23_P37	NM_00572	Homo sapi	ACTR2,AR	-1.79246	1.16E-34	0.55789	5976.444	3335.95
A_24_P66	NM_00660	stromal ant	STAG2,SA	-1.7925	1.53E-13	0.55788	2970.319	1651.071
A_23_P25	NM_00601	Homo sapi	CD164,MG	-1.79252	9.64E-26	0.55787	2378.567	1326.964
A_23_P23	NM_00174	Homo sapi	CAPN2,mC	-1.79255	6.80E-12	0.55786	49487.05	27805.89
A_23_P22	NM_00207	Homo sapi	GNB1	-1.7926	9.29E-16	0.55785	2898.189	1617.124
A_24_P12	NM_17070	Homo sapi	ICMT,PCM	-1.79293	3.23E-08	0.55775	1859.896	1022.008
A_24_P91	NM_00301	SET transl	SET,2PP2	-1.79297	7.01E-10	0.55773	19055.75	10752.27
A_23_P12	NM_00146	thyroid aut	XRCC6,ML	-1.793	7.22E-27	0.55772	4206.405	2343.308
A_24_P81298		Unknown	THC19103	-1.79322	0.00027	0.55766	461.3265	261.9609
A_24_P83	NM_00456	serine (or c	SERPINB6	-1.7935	1.64E-08	0.55757	1044.743	589.7795
A_23_P13	NM_00118	ataxia telar	ATR,FRP1	-1.79426	3.83E-34	0.55733	4122.942	2297.998
A_23_P12	NM_00226	Homo sapi	KPNA2,QIP	-1.79444	1.28E-25	0.55728	51049.67	28485.4
A_32_P55	AK096908	CDNA FLJ	AK096908	-1.79445	5.68E-11	0.55727	1823.738	1024.368
A_24_P31	NM_00632	spindle pol	TUBGCP3	-1.79447	6.11E-07	0.55727	1153.052	625.7537
A_24_P30	NM_00359	cullin 1	CUL1	-1.79486	9.39E-20	0.55715	2488.223	1385.871

A_32_P61	NM_00600	ubiquinol-c	UQCRFS1	-1.7949	3.23E-15	0.55714	22446.81	12590.98
A_32_P15	NM_00682	tyrosine 3-r	YWHAQ,1	-1.79509	2.68E-16	0.55707	47043.91	26194.9
A_23_P12	NM_01405	feline leuke	FLVCR	-1.79515	5.00E-24	0.55706	2239.532	1247.909
A_23_P33	NM_15336	Homo sapi	C10orf56,F	-1.7967	3.10E-06	0.55658	827.5579	449.0453
A_23_P39	NM_15249	Homo sapi	MANEAL,F	-1.79697	1.58E-09	0.55649	916.9092	513.4892
A_23_P43	NM_01763	Homo sapi	KCTD9,FL	-1.79719	1.15E-25	0.55642	4144.181	2308.717
A_23_P96	NM_02071	Homo sapi	LRRC47,K	-1.79743	1.05E-23	0.55635	3416.173	1900.874
A_32_P23	NM_00556	lactate deh	LDHA,LDH	-1.79821	9.58E-31	0.55611	71867.44	39886.12
A_23_P34	NM_01612	COP9 (con	COPS4,MC	-1.79869	1.32E-21	0.55596	5954.932	3313.638
A_24_P246591		NADP+-sp	ENST0000	-1.79899	1.04E-25	0.55587	4566.726	2536.706
A_24_P21	NM_15326	hypothetica	FLJ36812,	-1.79905	2.50E-06	0.55585	862.6142	466.1122
A_23_P19	AB028946	IQ motif co	IQCE	-1.7991	1.54E-12	0.55583	2508.572	1389.037
A_23_P35	NM_01868	Homo sapi	ANLN,Scra	-1.79933	1.47E-07	0.55576	2167.698	1192.978
A_24_P15	NM_02015	SAR1 prot	SARA1,SA	-1.7996	2.26E-12	0.55568	4650.624	2583.593
A_23_P48	NM_18178	SMILE pro	SMILE,FLJ	-1.80016	0.00004	0.5555	602.6033	344.7948
A_23_P81	NM_02019	HTGN29 p	C5orf15,KC	-1.80055	1.03E-10	0.55539	3562.174	1982.584
A_32_P14	NM_00693	SMT3 (sup	SUMO2,HS	-1.80097	1.87E-26	0.55526	26749.56	14880.31
A_32_P11181		P2G4_HUM	THC18164	-1.80172	8.42E-22	0.55502	18715.64	10408.32
A_32_P20	AK090739	CDNA FLJ	AK090739	-1.802	4.03E-21	0.55494	6299.501	3499.517
A_32_P77977		A40201 art	THC19995	-1.80215	4.91E-09	0.55489	806.6281	444.2312
A_23_P39	BC009229	Hypothetic	LOC90120	-1.80231	1.99E-07	0.55484	2554.854	1427.342
A_23_P25	NM_01243	Homo sapi	SEC22L2,S	-1.80261	7.63E-21	0.55475	1881.224	1043.07
A_23_P13	NM_01401	Homo sapi	ASCC3L1,	-1.80315	1.83E-34	0.55459	11330.57	6275.092
A_23_P87	NM_00310	Homo sapi	SORL1,LR	-1.80329	1.00E-07	0.55454	1046.014	594.8118
A_24_P12	NM_03223	Ubiquitin sp	USP48	-1.80341	3.15E-09	0.5545	1163.093	635.5367
A_23_P16	NM_01785	Homo sapi	PARP16,C	-1.80386	2.06E-06	0.55437	802.1107	433.8077
A_24_P28	NM_01754	KIAA1513	POGK,BAS	-1.80434	8.49E-12	0.55422	2593.873	1434.799
A_23_P25	AB058721	E1A bindin	EP400	-1.80517	4.45E-07	0.55396	751.6749	409.2629
A_24_P39	NM_01513	KIAA0252	KIAA0252	-1.80525	5.25E-18	0.55394	2242.104	1244.316
A_23_P13	NM_00210	H2A histon	H2AFZ,H2	-1.80529	9.48E-24	0.55393	74903.07	41613.76
A_23_P48	NM_00111	Homo sapi	ADAM10,k	-1.80534	8.70E-06	0.55391	674.6743	384.1149
A_24_P28	NM_14456	similar to R	ANGEL2,F	-1.80577	6.85E-26	0.55378	3408.935	1886.7
A_24_P45	NM_02044	Hypothetic	DJ462O23	-1.80597	0.00002	0.55372	884.4479	469.0536
A_23_P12	NM_02015	Homo sapi	STARD7,G	-1.80614	1.49E-22	0.55367	15769.09	8749.546
A_24_P29	D87454	KIAA0265	KIAA0265	-1.80702	1.31E-06	0.5534	817.1008	467.7047
A_24_P106166		Unknown	A_24_P10	-1.80739	0.00028	0.55328	396.5906	223.6118
A_23_P26	NM_02460	hypothetica	FLJ21908	-1.80753	1.32E-36	0.55324	2502.414	1387.434
A_24_P10	NM_01843	hypothetica	IMPACT,M	-1.80775	0.00033	0.55317	445.0718	243.9414
A_24_P418780		Unknown	A_24_P41	-1.80777	6.09E-26	0.55317	3683.908	2036.189
A_23_P91	NM_00473	Homo sapi	VAPB,ALS	-1.80793	1.34E-18	0.55312	1546.309	854.1675
A_23_P99	AL832695	Homo sapi	DKFZp313	-1.80796	4.82E-07	0.55311	693.6861	388.0172
A_24_P23	AB007963	KIAA0494	KIAA0494	-1.80796	3.59E-16	0.55311	1508.801	838.5278
A_23_P21	NM_01534	Homo sapi	LEPROTL1	-1.80815	1.18E-21	0.55305	2862.089	1583.865
A_32_P20	AK091492	Hypothetic	AK091492	-1.80839	8.49E-07	0.55298	779.0866	440.0087
A_32_P22	NM_00693	SMT3 (sup	SUMO2,HS	-1.80857	2.66E-26	0.55292	25041.41	13881.57
A_23_P16	NM_01545	KIAA1194	CNOT6,CC	-1.80902	3.42E-31	0.55279	3023.49	1671.339
A_24_P92	AK056260	LOC44075	AK056260	-1.80917	0.00002	0.55274	994.7535	573.6031
A_23_P10	NM_05296	Homo sapi	C1orf19,se	-1.80918	3.64E-29	0.55274	4012.993	2217.296
A_23_P34	NM_00014	Homo sapi	FH,MCL,LF	-1.80937	2.71E-27	0.55268	9550.816	5263.824
A_32_P132276		Unknown	A_32_P13	-1.80955	2.06E-10	0.55262	2668.487	1475.706
A_24_P607107		Unknown	A_24_P60	-1.81	7.41E-09	0.55249	3569.711	1946.25
A_32_P12	AI420375	Signal seq	SSR3	-1.8103	0.00017	0.55239	2781.538	1518.499

A_23_P20	NM_00032	Homo sapi	RB1,RB,O	-1.81083	2.34E-13	0.55223	3562.785	1973.156
A_24_P12	AL050097	DKFZP586	DLG1	-1.81133	0.00015	0.55208	506.572	273.467
A_23_P14	AF292100	Homo sapi	RP42 prote	-1.81157	1.77E-08	0.55201	1274.044	687.5387
A_23_P11	NM_01818	hypothetica	C1orf112,F	-1.81164	2.62E-20	0.55199	5077.112	2808.367
A_23_P13	NM_05283	Homo sapi	C6orf192,d	-1.81192	2.04E-29	0.5519	4731.051	2610.394
A_23_P21	NM_02300	Homo sapi	BAZ1B,WS	-1.81214	4.91E-16	0.55183	2764.999	1523.271
A_23_P20	NM_01595	likely ortho	MMRP19,A	-1.81232	0.00003	0.55178	624.1973	354.9801
A_23_P50	NM_00107	Homo sapi	TYMS,TS,T	-1.81251	2.69E-36	0.55172	17518.64	9655.025
A_23_P50	NM_01749	WD repeat	WDR1,AIP	-1.81253	2.66E-16	0.55172	4120.901	2274.158
A_23_P14	NM_01514	KIAA0153	KIAA0153	-1.81256	1.77E-23	0.5517	12197.66	6704.127
A_23_P13	NM_01466	Homo sapi	VGLL4,VG	-1.81268	5.48E-08	0.55167	742.583	412.6514
A_23_P95	NM_02468	hypothetica	FLJ23560	-1.81297	8.16E-06	0.55158	952.5121	546.2849
A_23_P12	NM_02044	Homo sapi	ACTR3B,A	-1.81347	5.16E-31	0.55143	3675.301	2027.259
A_32_P38	NM_18297	Homo sapi	RIMS4,RIM	-1.81458	1.56E-14	0.55109	2669.788	1474.009
A_24_P16	NM_01754	Homo sapi	EPDR1,UC	-1.81459	4.62E-31	0.55109	11789.34	6492.011
A_23_P92	NM_17680	Homo sapi	MOCS2,MI	-1.81497	1.55E-20	0.55097	1981.384	1091.316
A_24_P40	AK098759	Homo sapi	LOC55420	-1.81509	0.00002	0.55094	499.2241	273.6948
A_23_P13	NM_03148	Homo sapi	MARVELD	-1.8154	3.93E-33	0.55084	14397.28	7939.89
A_24_P30	NM_01239	phosphotid	PITPNB,VI	-1.81614	2.81E-31	0.55062	5649.57	3113.227
A_24_P98	NM_01332	sorting nex	SNX10,MG	-1.81623	1.26E-11	0.55059	1312.207	729.9092
A_23_P15	AB051551	KIAA1764	KIAA1764	-1.81649	0.00004	0.55051	536.5477	289.8094
A_23_P65	NM_13337	hypothetica	MGC4562,	-1.81695	8.58E-22	0.55037	2987.948	1644.846
A_23_P69	NM_00288	Homo sapi	RARS,ArgI	-1.81698	1.97E-30	0.55036	55324.13	30495.06
A_23_P42	AB028986	Ubiquitin sp	USP22	-1.81717	6.30E-16	0.55031	2168.243	1190.249
A_24_P10	NM_03286	Homo sapi	NUDCD1,C	-1.81751	0.00099	0.5502	454.4145	239.2738
A_32_P11	AF130048	Hypothetica	LOC28614	-1.81778	0.00015	0.55012	553.0786	293.2951
A_32_P16462		ALU6_HUM	THC19454	-1.81788	1.60E-06	0.55009	673.5231	377.1691
A_32_P43	NM_01511	KIAA0276	DCUN1D4	-1.81799	4.34E-07	0.55006	766.5778	415.6599
A_24_P76	AK123083	LYRIC/3D3	LYRIC	-1.81804	4.39E-09	0.55004	837.6205	464.0745
A_23_P21	NM_08054	Homo sapi	SLC44A1,C	-1.81869	3.74E-22	0.54985	15360.04	8471.194
A_24_P94	D80006	Homo sapi	KIAA0184	-1.81929	2.90E-06	0.54967	808.3046	432.0382
A_23_P41	NM_02523	nucleotide	COASY,NE	-1.81945	3.25E-27	0.54962	2962.098	1628.344
A_24_P29	NM_00680	chromobox	CBX1,CBX	-1.81985	1.00E-31	0.5495	9468.672	5204.995
A_23_P37	NM_01560	Homo sapi	WDR21A,W	-1.82049	7.53E-24	0.5493	2983.928	1638.487
A_24_P17	NM_01500	SMART/HE	SPEN,MIN	-1.82055	0.00011	0.54928	492.6895	265.0479
A_24_P41	NM_00481	U5 snRNP-	WDR57,SF	-1.82055	2.52E-06	0.54929	600.566	325.0459
A_32_P123204		Unknown	A_32_P12	-1.82088	0.00023	0.54918	1310.091	751.0259
A_23_P24	NM_00007	Homo sapi	CDK4,CMM	-1.82102	1.22E-19	0.54914	7689.075	4234.604
A_23_P41	AL834325	GTPase ac	DKFZP434	-1.82107	5.07E-09	0.54913	1296.356	723.98
A_24_P94	NM_01764	hypothetica	FAM29A,R	-1.82205	2.94E-14	0.54883	1719.933	938.1657
A_23_P35	NM_02468	likely ortho	E2F8,FLJ2	-1.82234	2.01E-07	0.54874	851.3628	457.5466
A_24_P41	BC014384	CDNA FLJ	BC014384	-1.82244	2.07E-22	0.54871	4084.996	2241.843
A_24_P23	NM_00614	dynein, cyt	DNCLI2,LI	-1.82263	3.13E-19	0.54866	6062.939	3336.136
A_23_P14	NM_01547	Homo sapi	C18orf10,L	-1.82265	2.53E-14	0.54865	1368.74	746.521
A_24_P93	AK021980	Transcripti	TCF4	-1.82276	0.00005	0.54862	697.3539	365.5918
A_24_P13	NM_00078	cytochrome	CYP51A1,I	-1.82331	1.73E-12	0.54845	5441.321	2994.18
A_24_P32	NM_00210	H3 histone	H3F3A,H3I	-1.82463	1.79E-27	0.54806	118805.3	65047.75
A_23_P80	NM_00462	Homo sapi	WRB,CHD	-1.82523	2.61E-11	0.54788	2419.199	1333.155
A_24_P36	NM_00320	Homo sapi	TCF3,E2A,	-1.82534	1.11E-15	0.54784	4964.967	2721.411
A_23_P51	NM_03063	Homo sapi	ZNF436,ZN	-1.8254	0.00068	0.54782	442.5761	252.7669
A_23_P16	NM_03299	Homo sapi	GTF2I,DIW	-1.8256	1.76E-15	0.54777	15474.37	8516.589
A_23_P10	NM_03188	Homo sapi	BBS2,BBS	-1.82578	2.69E-13	0.54771	3943.116	2160.515

A_32_P82	NR_00145	Homo sapi	C14orf62	-1.82604	4.81E-14	0.54763	1648.922	897.6112
A_23_P92	NM_13080	hypothetica	LOC13361	-1.82653	2.65E-11	0.54749	1279.753	707.6608
A_23_P31	NM_01898	Homo sapi	DNAJC10,	-1.82704	1.62E-10	0.54733	1737.359	940.8155
A_24_P80	NM_00315	serine/thre	NEK4,NRK	-1.82735	2.67E-15	0.54724	1541.7	838.6901
A_23_P21	NM_01590	Homo sapi	EIF5B,IF2,	-1.82746	1.19E-15	0.54721	1602.121	881.6164
A_32_P38228		Unknown	ENST0000	-1.82757	1.60E-08	0.54717	4214.806	2309.319
A_23_P12	NM_01490	KIAA0941	RAB11FIP	-1.82764	0.00034	0.54715	469.5929	249.8359
A_24_P68	BC041926	Homo sapi	BC041926	-1.82792	0.00002	0.54707	606.7515	324.1523
A_24_P39	NM_17369	Homo sapi	ATP11C,A	-1.82813	9.57E-11	0.54701	1600.893	866.9734
A_24_P14	NM_01792	hypothetica	PEX26,FL	-1.82857	1.19E-23	0.54688	4698.118	2573.509
A_24_P17	BC044624	Pre-B-cell l	PBX1	-1.82871	6.89E-22	0.54683	10721.95	5892.471
A_24_P92	AK001838	82-kD FMF	182-FIP	-1.82884	1.06E-16	0.54679	5324.058	2913.535
A_23_P10	AK026575	Heat shock	HSBP1	-1.82902	5.32E-14	0.54674	2024.682	1112.719
A_23_P43	NM_18261	hypothetica	FLJ34283,	-1.82906	4.56E-14	0.54673	1303.772	715.5541
A_23_P25	NM_01341	Homo sapi	IARS,ILRS	-1.82939	3.04E-20	0.54663	10573.22	5799.264
A_23_P76	NM_03075	Homo sapi	TXNDC,TM	-1.82982	1.54E-44	0.5465	16405.57	8965.679
A_24_P64	BC063684	Kinesin fan	KIF1B	-1.83079	2.69E-06	0.54621	936.4917	494.6978
A_23_P14	NM_00370	Homo sapi	SLC25A12	-1.83088	6.00E-24	0.54619	2801.371	1529.612
A_23_P25	NM_01604	Isochorism	CGI-111	-1.83115	5.20E-20	0.5461	4797.053	2612.652
A_23_P18	NM_00588	macrophag	MAEA,EMI	-1.83167	0.00004	0.54595	726.2125	378.8512
A_23_P39	AK055921	KIAA1935	KIAA1935	-1.83236	8.06E-08	0.54574	964.1862	515.9481
A_23_P91	NM_02463	Homo sapi	QTRTD1,F	-1.83274	5.83E-20	0.54563	2515.856	1371.49
A_24_P34	BC000147	Malic enzy	ME2	-1.83306	2.69E-08	0.54554	1024.502	548.5236
A_23_P15	NM_00276	Homo sapi	PRPSAP1,	-1.83344	5.13E-19	0.54542	2957.854	1610.136
A_23_P70	NM_03192	Homo sapi	REPS1,RA	-1.83445	5.75E-13	0.54512	4629.026	2529.795
A_23_P34	NM_02121	Homo sapi	ZNF71,EZI	-1.83463	1.82E-07	0.54507	863.0812	460.7262
A_23_P25	NM_00641	Homo sapi	AGPAT1,G	-1.83465	1.74E-27	0.54506	2413.182	1314.66
A_23_P21	AL834469	Enoyl Coe	ECHDC1	-1.83467	2.23E-20	0.54506	5601.283	3058.706
A_23_P50	NM_15273	Homo sapi	HOXA9,HC	-1.83486	1.33E-15	0.545	1310.111	716.0089
A_23_P41	NM_20743	FLJ40142	FLJ40142	-1.83511	5.80E-07	0.54493	1172.395	620.0006
A_32_P95	NM_01460	Homo sapi	UBXD2,UB	-1.83562	7.99E-08	0.54478	716.3388	390.424
A_23_P20	D43947	KIAA0100	KIAA0100	-1.83568	4.60E-39	0.54476	5463.893	2976.852
A_24_P36	NM_17858	Homo sapi	PPP2R5C,	-1.83576	5.23E-10	0.54473	1183.061	637.9689
A_24_P10	AK055664	WD repeat	WDR36	-1.83582	7.84E-08	0.54472	808.0554	448.6109
A_24_P39	NM_00169	ATPase, H	ATP6V1A,I	-1.83593	1.52E-12	0.54468	10248.87	5591.816
A_24_P18	AK023725	N-ethylmal	NAPG	-1.83629	1.10E-16	0.54458	1410.843	766.5186
A_23_P41	NM_20350	Ras-GTPa	G3BP2	-1.83654	1.55E-08	0.5445	871.609	480.6351
A_23_P33	BC032468	SH3 domai	EVE1	-1.83661	5.09E-10	0.54448	2117.136	1145.065
A_23_P15	NM_19840	Homo sapi	PTPLB	-1.83713	1.60E-13	0.54433	2112.811	1145.73
A_23_P31	NM_00518	Homo sapi	PCGF4,BM	-1.83723	1.04E-11	0.5443	1101.401	596.3263
A_23_P46	BX648514	Hypothetic	MGC29875	-1.83823	2.52E-08	0.544	2558.98	1380.68
A_24_P18	NM_01850	hypothetica	PRO1855	-1.83847	4.09E-12	0.54393	13749.04	7517.139
A_23_P12	NM_03333	Homo sapi	CASP7,MC	-1.83869	1.49E-06	0.54387	4914.441	2671.865
A_23_P38	AB002374	KIAA0376	KIAA0376	-1.83907	8.47E-08	0.54375	1060.807	564.4937
A_23_P50	NM_02053	Homo sapi	CSRP2BP,	-1.83957	2.86E-15	0.5436	2342.429	1268.52
A_23_P13	NM_02278	Homo sapi	ARV1	-1.84001	5.35E-34	0.54348	6055.774	3292.775
A_23_P16	NM_00596	Homo sapi	HNRPM,H	-1.84048	4.50E-19	0.54334	52473.51	28606.37
A_32_P31	BC035691	Glutathione	GSR	-1.84057	4.19E-14	0.54331	1754.141	947.4451
A_23_P13	NM_01323	putative rib	RNASEN,F	-1.84066	9.08E-13	0.54328	10854.88	5931.153
A_32_P16	NM_20683	Homo sapi	MORF4L1,	-1.84071	3.57E-25	0.54327	10935	5956.761
A_23_P34	NM_01444	gamma tub	76P,GCP4	-1.84077	4.25E-11	0.54325	880.376	478.3356
A_23_P21	NM_00387	Homo sapi	GMPS	-1.84079	3.41E-35	0.54325	11881.63	6447.378

A_24_P31	NM_00659	API5-like 1	API5,AAC-	-1.84101	5.62E-11	0.54318	1010.315	554.5105
A_32_P18	NM_01327	Rac GTPa	RACGAP1	-1.84118	1.88E-12	0.54313	7609.529	4158.723
A_32_P35	NM_01820	vacuolar pr	VPS35,ME	-1.84155	3.47E-09	0.54302	4433.157	2412.276
A_24_P32	NM_14506	Homo sapi	C18orf24,M	-1.84156	0.00039	0.54302	624.6162	364.2078
A_23_P66	NM_01572	gem (nucle	GEMIN4,H	-1.84179	8.92E-27	0.54295	7937.077	4322.789
A_23_P76	NM_00605	reticulon 3	RTN3,HAP	-1.84183	4.35E-07	0.54294	1360.959	718.9613
A_24_P90	NM_01849	G protein-c	LGR4,GPR	-1.84204	2.91E-11	0.54288	4276.804	2339.151
A_32_P14	CA314936	Transcribe	CA314936	-1.8421	1.76E-11	0.54286	1038.169	561.402
A_23_P13	NM_00040	Homo sapi	EXT2,SOT	-1.84237	6.20E-18	0.54278	1648.917	897.6105
A_32_P177953		Unknown	THC19679	-1.84267	2.71E-24	0.54269	12597.17	6858.442
A_24_P33	NM_03092	Homo sapi	NIPA2,MG	-1.84282	1.79E-09	0.54265	1397.416	772.075
A_23_P14	NM_01481	Homo sapi	KIAA0649,	-1.84285	9.81E-13	0.54264	1761.532	962.4965
A_24_P94	NM_01494	spastic par	SPAST,FS	-1.84345	5.57E-12	0.54246	999.1996	541.507
A_23_P15	NM_00600	Homo sapi	UQCRFS1	-1.8437	6.07E-32	0.54239	31562.82	17132.65
A_23_P21	NM_01432	Homo sapi	ZNF278,ZS	-1.84395	0.00034	0.54232	411.8804	222.9893
A_23_P54	AB037759	Eukaryotic	EIF2AK4	-1.84438	7.29E-11	0.54219	1500.587	823.3185
A_23_P21	NM_05402	Homo sapi	CNOT7,CA	-1.84543	2.34E-16	0.54188	1498.447	814.025
A_23_P11	NM_01605	CGI-120 pr	COPZ1,CC	-1.84581	4.79E-13	0.54177	1754.318	956.5681
A_24_P26	NM_00433	chromoson	C8orf1,hT4	-1.84613	6.74E-07	0.54167	824.1048	434.6636
A_23_P56	NM_17755	Homo sapi	ACP1,HAA	-1.84613	7.40E-21	0.54167	4401.475	2384.518
A_24_P31	AB007952	F-box prote	FBXO28	-1.84643	3.82E-09	0.54158	825.4962	443.6401
A_32_P33	NM_20747	FLJ27365	FLJ27365	-1.84657	1.19E-06	0.54155	640.4174	343.5043
A_23_P94	NM_00229	Homo sapi	LAMB1	-1.84669	1.26E-13	0.54151	10417.19	5669.734
A_24_P94	D80001	KIAA0179	KIAA0179	-1.84692	4.69E-11	0.54144	2257.833	1212.946
A_24_P14	NM_01752	leptin rece	LEPROT,L	-1.84697	9.20E-12	0.54143	1069.973	581.2949
A_32_P111852		Unknown	THC19984	-1.84719	1.57E-07	0.54136	2050.797	1094.448
A_23_P25	AK056178	Chromosor	C20orf121	-1.84722	6.99E-19	0.54135	4029.029	2181.443
A_23_P11	NM_01534	Homo sapi	PPWD1,KI	-1.84744	1.45E-26	0.54129	3413.552	1847.356
A_24_P13	NM_00574	accessory	BCAP31,C	-1.84769	1.08E-26	0.54122	5366.402	2897.123
A_23_P38	AK090846	TWIST nei	TWISTNB	-1.84819	3.52E-11	0.54107	2007.005	1075.402
A_23_P71	NM_00521	Homo sapi	DEFB1,BD	-1.8492	9.24E-11	0.54078	2102.026	1131.396
A_23_P60	NM_01617	Ubiquitin a	UBADC1	-1.84922	7.88E-30	0.54077	5862.244	3165.945
A_23_P54	NM_02124	Homo sapi	SNX6,TFA	-1.84958	1.24E-34	0.54066	4432.165	2395.194
A_24_P16	NM_00581	protein kin	PRKD3,EP	-1.84992	3.59E-08	0.54056	757.06	405.5201
A_23_P30	NM_15234	hypothetica	FLJ30656	-1.8505	3.28E-12	0.54039	1860.699	993.5629
A_24_P33	NM_00348	solute carri	SLC7A5,E	-1.85052	4.94E-18	0.54039	11598.18	6287.41
A_23_P26	NM_01908	limkain b1	LKAP,KIA	-1.85072	1.05E-16	0.54033	2148.723	1153.576
A_24_P55	AK057088	AF034176	AK057088	-1.85101	4.55E-17	0.54024	2508.77	1352.226
A_23_P36	NM_00461	transmemb	TSPAN8,C	-1.85136	3.21E-41	0.54014	15074.01	8147.406
A_24_P203976		Unknown	A_24_P20	-1.85152	3.20E-06	0.5401	674.8015	373.8055
A_32_P55414		ALU6_HUM	THC19988	-1.85215	1.62E-06	0.53991	1224.072	682.0066
A_23_P70	NM_01234	Homo sapi	NNT	-1.85248	2.60E-06	0.53982	657.1924	348.5167
A_23_P49	NM_00347	Homo sapi	USP7,TEF	-1.85313	4.77E-11	0.53963	5812.523	3164.698
A_24_P38	NM_02199	integral me	ITM2B,BRI	-1.85335	6.25E-10	0.53956	2157.066	1153.118
A_23_P33	NM_00039	Homo sapi	COL5A2,M	-1.85363	2.29E-10	0.53948	5477.38	2955.081
A_24_P10	NM_14572	Homo sapi	DMN,SYN,	-1.8537	5.15E-32	0.53946	2662.094	1436.055
A_23_P37	NM_00591	Homo sapi	MCM4,CD	-1.85582	4.35E-08	0.53885	3368.072	1792.837
A_23_P42	NM_00028	Homo sapi	PEX6,PAF	-1.85701	7.71E-14	0.5385	1364.891	742.8337
A_23_P79	NM_00681	proteasom	PSMF1,PI3	-1.85788	1.05E-20	0.53825	2243.463	1205.11
A_23_P11	NM_01495	Homo sapi	BAHD1,KI	-1.85797	5.75E-32	0.53822	3804.831	2048.535
A_23_P22	NM_00635	Homo sapi	SLC9A6,N	-1.85819	1.34E-09	0.53816	1464.696	776.243
A_23_P10	NM_00643	Homo sapi	CCT4,SRB	-1.85887	3.37E-12	0.53796	184054.7	99754.38

A_23_P32	NM_00174	Homo sapi	CALM2,PH	-1.8591	4.05E-12	0.53789	103888.5	56241.04
A_32_P17	NM_01335	CCR4-NOT	CNOT7,CA	-1.85994	1.90E-34	0.53765	6436.522	3457.379
A_23_P20	NM_00271	Homo sapi	PPP1CC,P	-1.86045	8.93E-18	0.53751	22132.67	11924.22
A_24_P410256		Unknown	A_24_P41	-1.86056	7.44E-24	0.53747	7480.167	4033.213
A_24_P35	NM_00210	H2A histon	H2AFZ,H2	-1.86068	1.20E-35	0.53744	133487.3	71858.48
A_23_P25	NM_00192	Homo sapi	DDB1,XPE	-1.86085	3.22E-12	0.53739	6512.149	3464.419
A_23_P23	NM_01443	Homo sapi	SLC39A1,Z	-1.86117	3.65E-10	0.5373	2296.967	1240.665
A_32_P13	NM_13927	Homo sapi	AKAP1,AK	-1.86123	9.92E-19	0.53728	8613.903	4649.228
A_24_P12	NM_00457	RAB4A, m	RAB4A	-1.86138	4.60E-16	0.53724	1932.532	1035.19
A_23_P37	NM_01535	Homo sapi	POFUT1,F	-1.86188	0.00022	0.53709	558.8727	284.9605
A_24_P94	NM_00677	Homo sapi	QKI,QK,Ho	-1.86265	1.59E-06	0.53687	1074.457	556.9841
A_24_P58	BC039246	Homo sapi	BC039246	-1.86352	7.26E-29	0.53662	3155.818	1694.801
A_23_P16	NM_02290	hypothetica	SLC30A5,Z	-1.8637	1.03E-06	0.53657	1148.805	595.106
A_24_P81	NM_01432	coronin, ac	CORO1C,H	-1.86484	6.26E-19	0.53624	3661.738	1963.775
A_23_P38	NM_17801	Homo sapi	TUBB,M40	-1.86526	7.31E-32	0.53612	3282.578	1759.797
A_23_P69	NM_18276	Homo sapi	SUMF1,FG	-1.86655	3.00E-09	0.53575	3041.637	1651.365
A_24_P79	AK056910	Hypothetic	FLJ11000	-1.86878	2.08E-16	0.53511	1538.393	820.6023
A_24_P12	NM_00608	3'(2'), 5'-bis	BPNT1,PIF	-1.86892	0.0001	0.53507	1218.46	614.5116
A_32_P61	AK096154	MRNA fron	AK096154	-1.869	0.00017	0.53504	498.1802	277.2466
A_32_P123629		ALU5_HUM	THC20038	-1.86976	8.69E-25	0.53483	2877.03	1540.278
A_23_P14	NM_13833	Homo sapi	GNPDA2,S	-1.8702	1.67E-13	0.5347	954.9533	510.7348
A_23_P24	NM_02467	hypothetica	FLJ23441	-1.87021	3.58E-11	0.5347	938.6265	498.6074
A_23_P21	NM_00190	Homo sapi	CTPS	-1.87034	2.13E-15	0.53466	9846.154	5312.055
A_24_P12	NM_00135	DEAD/H (A	DDX3X,DB	-1.87049	5.05E-20	0.53462	2593.49	1391.881
A_23_P11	NM_01819	Homo sapi	RCBTB1,G	-1.87064	3.01E-08	0.53458	751.2843	398.5424
A_23_P20	NM_00729	Homo sapi	BRCA1,IRI	-1.87156	4.48E-20	0.53431	3241.698	1742.997
A_24_P38754		Unknown	THC18917	-1.8719	9.34E-07	0.53422	646.1163	351.0644
A_23_P10	AK098212	Protein BA	FLJ10359	-1.87249	1.98E-18	0.53405	8143.328	4358.634
A_32_P27706		L10908 Gc	THC19450	-1.87277	7.57E-15	0.53397	1666.258	896.2838
A_24_P41	AL117555	SEC6-like	SEC6L1	-1.87305	3.35E-07	0.53389	1797.519	938.5876
A_24_P70	BC022928	Homo sapi	BC022928	-1.87431	1.34E-29	0.53353	17611.43	9406.721
A_23_P40	NM_13843	Homo sapi	GPRASP2	-1.87441	5.34E-12	0.5335	1238.35	653.9106
A_24_P40	NM_00168	ATPase, C	ATP2B4,M	-1.87454	6.83E-26	0.53347	2980.6	1589.258
A_23_P39	AB020684	KIAA0877	KIAA0877	-1.87503	6.77E-09	0.53332	1013.676	529.5259
A_24_P144303		Unknown	A_24_P14	-1.87538	4.87E-08	0.53322	1112.655	608.2298
A_24_P35	NM_00728	GABA(A) r	GABARAP	-1.87541	4.92E-25	0.53322	2753.125	1467.765
A_23_P20	NM_02513	hypothetica	C13orf23,F	-1.87716	6.91E-13	0.53272	1074.771	573.6365
A_32_P107752		Unknown	THC19025	-1.87741	7.28E-09	0.53265	952.3795	498.2936
A_23_P94	NM_03142	Homo sapi	C9orf58,IB	-1.87749	4.65E-23	0.53263	3365.317	1790.964
A_23_P93	NM_01602	Homo sapi	TFB1M,CG	-1.87909	2.76E-10	0.53217	1347.381	727.6232
A_24_P24	AK098497	THO comp	THOC1	-1.87944	2.02E-09	0.53207	1070.043	561.6607
A_32_P103131		ALU1_HUM	THC20038	-1.88025	6.42E-26	0.53185	3949.55	2099.304
A_23_P36	NM_17355	Sterile alph	SAMD6	-1.88115	2.18E-37	0.53159	4040.105	2147.675
A_24_P33	AB032966	Tetratricop	TTC7A	-1.88125	2.35E-08	0.53156	1153.684	600.3387
A_24_P91	AB062488	OK/SW-cl.9	OK/SW-cl.9	-1.88209	0.00009	0.53132	439.0568	234.2262
A_23_P16	NM_00532	Homo sapi	HADHSC,S	-1.8823	6.38E-16	0.53127	9577.699	5092.891
A_24_P521544		Unknown	A_24_P52	-1.88272	1.25E-13	0.53115	1549.099	815.9917
A_32_P47	AK025166	Chromosor	C10orf12	-1.88303	2.08E-16	0.53106	2112.791	1131.989
A_23_P13	NM_01651	Homo sapi	VPS54,HC	-1.88307	5.24E-17	0.53105	1609.113	851.12
A_24_P46	AK095803	Malate deh	MDH2	-1.88381	2.74E-14	0.53084	1379.344	728.6981
A_24_P15	NM_00736	nuclear ca	NCBP2,NIF	-1.8839	1.50E-23	0.53081	4248.577	2257.253
A_24_P25	BC012895	Solute carr	SLC7A1	-1.88419	2.40E-09	0.53073	2892.07	1534.565

A_24_P47	NM_02409	hypothetica	MGC5508	-1.88451	5.63E-13	0.53064	7268.107	3863.858
A_24_P72	NM_00640	actin relate	ARPC1A,A	-1.8846	1.70E-31	0.53062	11435.61	6074.004
A_32_P18	AF277175	KIAA0265	KIAA0265	-1.88485	2.74E-35	0.53055	4588.46	2433.472
A_23_P59	NM_00059	Homo sapi	TAP1,APT	-1.88534	1.80E-11	0.53041	5849.39	3136.055
A_23_P69	NM_00632	SEC24 (S.	SEC24B,S	-1.88552	0.00006	0.53036	672.7049	337.923
A_23_P92	NM_00235	Homo sapi	MAD2L1,M	-1.8857	2.79E-15	0.53031	4998.591	2639.948
A_24_P39	NM_01597	neuronal c	TAF9L,DN	-1.88605	8.21E-08	0.53021	676.8046	357.0548
A_24_P40	NM_00407	cryptochro	CRY1,PHL	-1.88649	0.00078	0.53009	380.9377	205.4751
A_23_P70	NM_03201	Homo sapi	STARD3NI	-1.88671	4.74E-23	0.53002	6194.819	3292.124
A_23_P65	NM_02191	Homo sapi	CFL2	-1.88713	5.85E-08	0.5299	2415.817	1267.857
A_24_P19	NM_00291	regulator o	RENT1,UP	-1.88725	3.67E-07	0.52987	5700.36	3023.641
A_23_P21	NM_01333	protein trar	SEC61A1,	-1.88754	2.21E-12	0.52979	2234.26	1188.03
A_23_P65	NM_01327	Homo sapi	RACGAP1	-1.88766	1.71E-14	0.52976	7101.385	3749.166
A_23_P83	NM_00586	Homo sapi	NET1,NET	-1.88827	8.49E-30	0.52959	4193.625	2223.026
A_23_P21	NM_00267	plastin 1 (I	PLS1,I-PL	-1.88838	0.00005	0.52955	551.7249	303.5466
A_23_P29	NM_00190	catenin (ca	CTNNB1,C	-1.88897	9.57E-42	0.52939	15294.79	8103.898
A_23_P72	NM_03288	hypothetica	MGC11308	-1.889	4.86E-14	0.52938	4235.357	2227.068
A_23_P14	BC021127	KIAA0368	KIAA0368	-1.88927	3.78E-42	0.5293	7038.288	3725.677
A_24_P40	NM_00632	progestero	PGRMC2,I	-1.88927	8.76E-25	0.52931	11270.31	5979.051
A_24_P23	BC009463	CDNA FLJ	BC009463	-1.88953	5.30E-20	0.52923	2493.675	1322.128
A_23_P39	NM_08067	Homo sapi	KCNE4,MI	-1.88963	4.38E-15	0.5292	3563.889	1884.831
A_24_P90	BC064610	Selenopho	SEPHS1	-1.88993	3.77E-06	0.52912	1157.407	587.0508
A_24_P20	NM_01433	solute carri	SLC7A11,>	-1.89017	9.52E-11	0.52905	1114.337	594.1737
A_23_P21	NM_17306	Homo sapi	CAST,BS-1	-1.89037	1.46E-28	0.529	6390.33	3376.871
A_23_P25	NM_00675	Homo sapi	UGP2,UDF	-1.89044	1.37E-14	0.52898	7037.827	3731.104
A_24_P10	NM_00029	polycystic	PKD2,PKD	-1.89045	0.00019	0.52897	508.1085	281.9985
A_32_P127248		Unknown	THC18193	-1.891	1.35E-12	0.52882	1015.323	538.7333
A_23_P13	NM_02033	Homo sapi	RAI17,Zim	-1.89137	5.09E-16	0.52872	6053.421	3211.959
A_32_P61	NM_00283	protein tyr	PTPN11,C	-1.89164	1.11E-21	0.52864	11427.39	6058.783
A_23_P13	NM_01242	Homo sapi	SF3B3,RS	-1.89168	1.83E-23	0.52863	2769.25	1467.116
A_32_P67	BX647764	Keratin ass	KRTAP4-7	-1.89201	2.03E-17	0.52854	5246.015	2777.261
A_24_P16	AB007891	Homo sapi	KIAA0431	-1.89218	6.58E-09	0.52849	801.48	418.5487
A_23_P12	NM_01523	likely ortho	CSTF2T,C	-1.89235	0.00002	0.52844	633.2527	348.6393
A_23_P54	NM_01538	Homo sapi	HECTD1,K	-1.8929	1.57E-29	0.52829	5730.387	3030.648
A_23_P40	NM_01476	Homo sapi	DAZAP2,M	-1.89313	2.07E-36	0.52823	23527.48	12414.55
A_24_P40	NM_00721	SEC63 pro	SEC63,ER	-1.89317	1.01E-07	0.52821	673.3602	357.1018
A_24_P191357		Unknown	THC19455	-1.8932	0.00041	0.52821	394.2004	211.9332
A_23_P34	NM_13847	Homo sapi	CDAN1,CD	-1.89336	0.00091	0.52816	495.569	280.6039
A_23_P28	NM_02462	hypothetica	FLJ21901,	-1.89337	1.60E-11	0.52816	2269.295	1205.143
A_23_P21	BC036001	Dehydroge	DHRS8	-1.89353	4.64E-26	0.52811	3148.892	1663.182
A_24_P76	BC048969	TSPY-like	TSPYL1	-1.89446	1.93E-19	0.52786	3840.067	2030.782
A_23_P14	NM_01460	Homo sapi	CYFIP1,SH	-1.89451	1.32E-08	0.52784	2254.318	1174.081
A_23_P32	NM_00327	Homo sapi	TMPO,TP,	-1.89457	3.82E-16	0.52782	1699.634	893.1051
A_32_P84	AK026668	U2AF hom	UHMK1	-1.89489	5.17E-14	0.52774	13006.9	6872.133
A_24_P21	NM_01789	hypothetica	C20orf11,T	-1.89522	1.21E-29	0.52764	4732.578	2496.53
A_24_P49	AB051517	Hypothetic	FLJ13456	-1.89544	0.00009	0.52758	564.3066	312.8948
A_23_P12	NM_02275	chromoson	C9orf12,IN	-1.89646	5.98E-11	0.5273	2607.719	1373.067
A_24_P29	NM_01461	expressed	ETEA,KIA	-1.89703	6.46E-08	0.52714	2198.441	1137.127
A_23_P80	NM_02120	Homo sapi	SRPRB,AF	-1.89732	2.96E-16	0.52706	6389.355	3373.598
A_24_P93	AF305817	Syntaxin 1	STX16	-1.89739	3.98E-15	0.52704	5861.268	3109.367
A_23_P36	AL390079	Hypothetic	LOC58489	-1.89743	1.86E-08	0.52703	746.3898	395.8655
A_23_P14	NM_01866	Homo sapi	ZNF395,PE	-1.89784	2.62E-09	0.52691	865.4024	450.4453

A_24_P41	NM_00180	centromere	CENPA	-1.89834	6.28E-08	0.52678	1328.865	681.9475
A_24_P12	NM_00513	DiGeorge s	DGCR2,ID	-1.89836	4.51E-23	0.52677	3031.751	1596.715
A_23_P25	NM_00727	Homo sapi	TUSC2,PA	-1.89851	8.50E-39	0.52673	6833.735	3599.281
A_24_P88	NM_15262	DCP2 deca	DCP2	-1.89866	1.21E-09	0.52669	1184.169	613.645
A_24_P122403		Unknown	THC18315	-1.89942	1.26E-17	0.52648	1269.214	666.197
A_23_P15	NM_00719	chromatin-s	SUPT16H,	-1.89963	4.10E-12	0.52642	2108.442	1099.944
A_32_P12	BC039468	Nucleopori	NUP50	-1.89966	8.72E-26	0.52641	2739.99	1445.255
A_24_P32	NM_01808	hypothetica	POLR3B,R	-1.90023	7.50E-16	0.52625	1228.869	646.2568
A_24_P32	BX538061	Chromosor	C9orf41	-1.9005	3.65E-15	0.52618	3067.912	1612.554
A_23_P33	NM_01590	neuroblast	NAG,DKFZ	-1.90174	3.96E-13	0.52583	1026.595	540.3533
A_24_P38	NM_03094	HESB like	HBLD2	-1.90184	0.00021	0.52581	434.5444	233.4742
A_23_P38	NM_01536	Homo sapi	R3HDM1,F	-1.90212	2.11E-08	0.52573	1456.552	747.3014
A_24_P37	NM_00478	vesicle-ass	VAMP3,CE	-1.90216	0.0002	0.52572	482.8682	244.7551
A_24_P73920		IMB1_HUM	THC18332	-1.90226	5.84E-10	0.52569	902.4766	468.4807
A_23_P36	NM_00132	Homo sapi	CSTF1,Cst	-1.90267	2.74E-27	0.52558	4503.981	2367.779
A_23_P19	NM_00633	lysophosph	LYPLA1,LF	-1.90322	1.28E-13	0.52542	2833.423	1487.189
A_32_P60	NM_00551	homeo box	HLXB9,HB	-1.90389	4.87E-12	0.52524	1563.062	811.3236
A_24_P13	NM_00487	BCL2-asso	BAG5,BAG	-1.90558	1.74E-24	0.52477	1983.235	1039.658
A_23_P38	NM_15333	Homo sapi	KCTD6,MC	-1.90742	5.27E-10	0.52427	953.2284	494.5474
A_32_P50	NM_00080	FK506-bin	FKBP1A,F	-1.90814	5.76E-25	0.52407	2035.136	1068.4
A_23_P88	NM_02100	Homo sapi	NR2F2,AR	-1.90839	3.14E-34	0.524	4391.714	2301.407
A_24_P61	NM_00358	cullin 4A	CUL4A	-1.90892	1.42E-10	0.52386	3890.229	2033.702
A_23_P10	NM_00025	Homo sapi	MSH2,FCC	-1.90935	6.41E-15	0.52374	7050.332	3709.376
A_24_P18	NM_02452	hypothetica	FLJ20986	-1.90943	1.61E-35	0.52372	4698.941	2458.383
A_24_P108281		Unknown	A_24_P108	-1.9095	3.16E-07	0.5237	622.7915	327.115
A_32_P51	D86978	Nucleopori	NUP205	-1.90979	5.32E-17	0.52362	6883.589	3611.354
A_32_P45	NM_00589	isocitrate d	IDH1,IDH,I	-1.90992	2.79E-35	0.52358	17407.97	9119.307
A_24_P35	NM_01659	DHHC1 pro	ZDHHC3,G	-1.91024	2.60E-16	0.5235	1729.138	900.8498
A_24_P14	NM_18271	Homo sapi	SYPL1,SYI	-1.9112	2.06E-14	0.52323	1104.972	578.9329
A_23_P41	NM_00669	bladder ca	BLCAP,BC	-1.91125	2.75E-20	0.52322	2008.641	1053.875
A_24_P88	NM_00603	histone de	HDAC4,HD	-1.91157	5.65E-08	0.52313	1536.922	787.6627
A_23_P68	NM_00404	Homo sapi	ATIC,PURI	-1.91174	9.93E-31	0.52308	8515.819	4463.672
A_24_P36	NM_02016	hypothetica	DHX33,DD	-1.91202	4.23E-16	0.52301	2141.336	1124.016
A_23_P63	NM_13916	Homo sapi	TRUB1	-1.91309	5.48E-12	0.52272	3302.61	1727.155
A_23_P41	AB040955	KIAA1522	KIAA1522	-1.91313	3.22E-31	0.5227	3539.311	1849.583
A_23_P74	NM_01871	RCC1-like	RCC2,TD-	-1.91326	2.07E-18	0.52267	10312.7	5349.545
A_23_P16	NM_01553	Homo sapi	GORASP2	-1.91385	5.34E-20	0.52251	3285.19	1716.811
A_24_P97	NM_03286	hypothetica	PRPF38A,	-1.91419	5.88E-17	0.52242	2481.624	1290.359
A_23_P37	NM_00679	MORF-rela	MORF4L1,	-1.91543	1.69E-40	0.52208	4868.78	2542.551
A_23_P35	NM_00063	Homo sapi	BCL2,Bcl-2	-1.91553	0.00023	0.52205	496.1915	273.307
A_24_P17	NM_02206	likely ortho	E2-230K,F	-1.91562	4.05E-26	0.52203	6504.642	3403.275
A_32_P86	BC035036	Aminoacyl	ACY1L2	-1.91588	1.36E-19	0.52195	1674.997	871.3096
A_23_P25	NM_00416	RAB1A, m	RAB1A	-1.91629	3.04E-20	0.52184	3034.056	1582.479
A_23_P11	NM_00188	Homo sapi	CRYZ	-1.91672	8.93E-14	0.52173	1611.999	852.9627
A_23_P25	NM_01207	Homo sapi	CCT5,CCT	-1.91801	0	0.52137	124007.4	64630.51
A_23_P11	NM_00010	Homo sapi	DLD,E3,LA	-1.91902	9.08E-21	0.5211	16705.12	8738.99
A_23_P12	NM_00608	tubulin, alp	K-ALPHA-	-1.91923	4.19E-13	0.52104	390968.6	204941.9
A_24_P24	NM_00404	ATP synth	ATP5A1,O	-1.92023	1.92E-39	0.52077	30520.32	15874.92
A_24_P51	NM_00715	Upstream	UNR	-1.92058	1.05E-29	0.52068	5458.575	2834.597
A_23_P10	NM_00017	Homo sapi	MSH6,GTE	-1.92069	1.64E-32	0.52065	26122.86	13618.54
A_24_P21	BC001097	Eukaryotic	EIF4B	-1.92118	7.41E-30	0.52051	3182.324	1657.558
A_23_P51	NM_02067	Homo sapi	SPBC25,A	-1.92138	2.26E-11	0.52046	6255.51	3276.642

A_24_P69	NM_03078	mitochondr	SLC25A32	-1.9214	1.71E-23	0.52045	1957.563	1015.805
A_23_P11	NM_00289	Homo sapi	RBBP7,Rb	-1.92152	5.61E-45	0.52042	42305.09	22030.06
A_24_P49	BC020516	Interferon r	IRF2BP2	-1.9221	1.00E-08	0.52026	770.9928	398.1175
A_24_P92	AL137543	Tropomodu	TMOD3	-1.92218	2.47E-10	0.52024	1279.706	678.3491
A_23_P59	NM_01779	Homo sapi	YTHDF1,C	-1.92255	1.76E-19	0.52014	12451.12	6502.712
A_23_P10	NM_00714	zinc finger	ZNF146,O	-1.92274	4.08E-10	0.52009	1036.63	531.6359
A_23_P46	NM_00168	Homo sapi	ATP5F1,M	-1.92444	1.99E-27	0.51963	18520.64	9656.064
A_23_P36	NM_15323	Homo sapi	CRI2,EID2	-1.92485	2.54E-17	0.51952	3629.706	1886.564
A_23_P12	NM_00632	Homo sapi	RAN,TC4,A	-1.92516	1.13E-07	0.51944	1148.337	613.7732
A_24_P75994		Unknown	A_24_P75	-1.92638	6.38E-07	0.51911	1631.187	828.7448
A_23_P253701		AF516696	THC20051	-1.92649	4.35E-08	0.51908	628.3407	327.7878
A_23_P20	AB011134	Glycine-, g	KIAA0562	-1.92778	5.53E-09	0.51873	1074.242	543.5186
A_23_P15	NM_00253	Homo sapi	NUP88,MG	-1.92799	9.84E-19	0.51867	4521.847	2352.192
A_23_P25	NM_00363	Homo sapi	PEX3,TRG	-1.92888	4.95E-09	0.51844	1018.093	518.5532
A_23_P10	NM_00141	Homo sapi	EIF4G2,p9	-1.92897	1.72E-20	0.51841	28226.37	14658.4
A_23_P11	NM_01754	NF-kappaE	NKRF	-1.9292	5.37E-38	0.51835	3080.952	1596.908
A_23_P82	NM_00619	Homo sapi	PCM1,PTC	-1.92927	4.17E-15	0.51833	1373.24	707.0681
A_32_P10	BC043426	FRBZ1 pro	FRBZ1	-1.9296	1.75E-15	0.51824	1659.367	855.6571
A_23_P16	NM_00293	ring finger	RNF4,SNU	-1.92966	4.26E-25	0.51823	2973.733	1539.099
A_24_P12	D31762	Translocati	TRAM2	-1.92978	0.00003	0.51819	585.3841	316.8637
A_23_P11	NM_00496	Homo sapi	HDAC1,HD	-1.93047	2.10E-38	0.51801	8969.271	4641.771
A_23_P16	NM_01431	Trans-pren	TPRT	-1.93126	2.72E-23	0.5178	2082.819	1082.167
A_24_P31	BX647782	Nuclear tra	NUTF2	-1.93133	2.70E-10	0.51778	863.9595	450.2263
A_24_P10	NM_20699	CCR4-NOT	CNOT1	-1.93178	0.00011	0.51766	442.373	226.3437
A_24_P91	M12996	Human glu	G6PD	-1.93292	1.91E-09	0.51735	4157.961	2145.494
A_32_P51	AF086045	Full length	AF086045	-1.93299	0.00098	0.51733	426.7341	208.9513
A_23_P98	NM_14564	Homo sapi	SLC15A4,F	-1.93367	1.07E-27	0.51715	3437.042	1777.397
A_32_P115258		Unknown	THC19690	-1.93376	4.28E-07	0.51713	579.7264	300.0718
A_23_P10	NM_02452	Homo sapi	TTC13,FLJ	-1.93377	5.04E-10	0.51712	867.2645	443.3433
A_24_P28	NM_00279	proteasom	PSMB2,HC	-1.93416	8.80E-36	0.51702	9819.916	5084.155
A_23_P14	NM_07847	Homo sapi	TM2D3,BL	-1.93471	3.68E-26	0.51687	10521.47	5455.304
A_23_P66	NM_01522	retinoblast	RAP140,se	-1.93515	0.0003	0.51676	490.6635	268.8755
A_24_P12	BC048298	Transducin	TBL1XR1	-1.93584	1.10E-16	0.51657	3634.219	1881.845
A_23_P10	NM_00693	Homo sapi	SUMO3,SM	-1.9359	5.05E-21	0.51655	2170.642	1123.525
A_23_P43	NM_00451	interleukin	ILF3,DRBF	-1.93656	2.94E-26	0.51638	2257.475	1163.088
A_23_P10	BC000907	Vang-like 1	VANGL1	-1.93674	5.68E-10	0.51633	1048.125	551.8397
A_32_P93	AK025522	Similar to F	LOC44014	-1.93707	2.32E-09	0.51624	2268.487	1161.466
A_23_P17	NM_02194	FLJ12716	FLJ12716	-1.93746	5.77E-06	0.51614	879.9427	432.0378
A_24_P671875		AB015066	THC19206	-1.93757	1.38E-24	0.51611	2599.996	1340.249
A_23_P19	NM_00682	Homo sapi	ASCC3,RN	-1.93822	8.71E-36	0.51594	3399.666	1754.281
A_23_P43	D83781	Nucleopori	NUP160	-1.93845	9.51E-11	0.51588	2590.247	1333.094
A_23_P21	NM_00551	Homo sapi	GTF2E1,FI	-1.93882	5.28E-17	0.51578	2773.876	1434.535
A_23_P79	BC005078	Hypothetic	FLJ10996	-1.93907	6.01E-06	0.51571	853.6844	461.9227
A_24_P92	NM_19913	hypothetica	LOC13414	-1.93973	0.00004	0.51553	490.6381	259.3076
A_23_P43	NM_01612	Homo sapi	COPS4,MC	-1.93981	5.71E-25	0.51552	4241.234	2189.701
A_23_P11	NM_00643	ADP-ribosy	PARP4,PH	-1.94032	2.51E-10	0.51538	2734.389	1405.322
A_24_P17	AB082527	Acyl-Coenz	ACBD5	-1.94097	5.19E-06	0.51521	711.713	351.9819
A_32_P23	AK124281	CDNA FLJ	AK124281	-1.9413	4.74E-16	0.51512	1756.594	909.4788
A_24_P88	NM_00384	peroxisom	PEX11B,P	-1.94152	3.36E-14	0.51506	1434.76	732.2336
A_24_P17	NM_00412	glia matura	GMFB,GM	-1.94153	8.82E-15	0.51506	2600.861	1333.793
A_23_P81	AJ420574	Malic enzy	ME1	-1.94239	3.38E-22	0.51483	3333.104	1715.891
A_23_P20	NM_00029	Homo sapi	PHKB	-1.94297	1.24E-18	0.51468	5243.813	2711.854

A_24_P39	NM_17585	Homo sapi	NM_17585	-1.94307	0.00065	0.51465	421.6595	207.7824
A_23_P45	NM_03336	Homo sapi	KRAS,KRA	-1.94352	0.00061	0.51453	523.1256	248.5842
A_24_P23	AB033004	CD2 antige	CD2BP2	-1.94353	0.00002	0.51453	510.3011	257.8466
A_32_P14243	AE016951	THC20045		-1.94395	7.32E-18	0.51442	1529.317	782.9599
A_23_P14	NM_15294	developme	DRB1	-1.94451	2.33E-26	0.51427	2296.162	1184.447
A_23_P16	NM_14472	hypothetica	FLJ31951,	-1.94497	1.22E-27	0.51415	2312.818	1191.787
A_23_P25	NM_00078	Homo sapi	DCK,MGC	-1.94613	3.19E-15	0.51384	1300.578	662.7124
A_24_P94	NM_00632	ariadne (D	ARIH2,ARI	-1.94615	2.20E-14	0.51384	9331.053	4823.488
A_23_P57	NM_01416	Homo sapi	MKRN2,RN	-1.94687	5.15E-18	0.51365	1906.887	976.6477
A_23_P70	NM_17689	Homo sapi	PPAP2A,P	-1.94825	1.30E-19	0.51328	2450.917	1262.549
A_24_P97	NM_01662	hypothetica	MRPL35	-1.94834	1.02E-26	0.51326	6179.292	3177.931
A_24_P12	NM_01866	papillomav	ZNF395,PE	-1.94863	1.29E-17	0.51318	1954.359	996.1169
A_23_P10	NM_01595	thioredoxin	TXNDC14,	-1.94888	1.89E-30	0.51312	4884.227	2502.979
A_23_P21	NM_01521	KIAA0433	KIAA0433	-1.94928	0	0.51301	2581.196	1327.854
A_24_P29	NM_01814	hypothetica	FAM82C,p	-1.94932	0.00004	0.513	473.2181	247.9037
A_32_P39	AL390079	Hypothetic	LOC58489	-1.94957	1.20E-24	0.51293	5518.62	2822.816
A_23_P47	NM_01655	seven tran	TM7SF3	-1.94969	7.47E-37	0.5129	5361.099	2748.164
A_23_P32	NM_01884	Homo sapi	KLHL7,KLH	-1.94984	0.00043	0.51286	439.5693	215.7376
A_24_P28	NM_01604	CGI-107 pr	C20orf45,d	-1.94991	1.42E-12	0.51284	5106.96	2619.669
A_23_P16	NM_14564	DEAH (Asp	DHX57	-1.95058	0.00003	0.51267	481.5627	242.8496
A_24_P373844	Unknown	THC19679		-1.95064	0.00004	0.51265	465.4072	243.2588
A_23_P21	AF063592	Chromosor	C7orf2	-1.95071	4.77E-07	0.51263	947.5485	468.3693
A_24_P166789	ALU6_HUM	THC19528		-1.95085	5.55E-24	0.5126	1548.613	792.7921
A_23_P38	NM_01563	Homo sapi	KIAA1279,	-1.9514	2.01E-16	0.51245	1452.407	749.5115
A_23_P10	NM_01613	Homo sapi	INSIG2,MG	-1.95151	0.00072	0.51242	400.9001	214.9832
A_23_P32	NM_00208	Homo sapi	GOT2,FLJ4	-1.95158	1.97E-32	0.51241	3945.853	2020.848
A_23_P21	NM_01833	Homo sapi	RFK,RIFK,	-1.95178	2.33E-10	0.51235	1904.562	965.5233
A_32_P15	NM_00300	succinate d	SDHD,PGL	-1.95206	1.26E-33	0.51228	4340.079	2224.368
A_23_P11	NM_00448	Homo sapi	GCSH,GCI	-1.95256	8.94E-27	0.51215	17775.18	9140.884
A_24_P92	AK130101	Peptidylpro	PPIA	-1.95284	5.18E-07	0.51207	751.7792	374.2184
A_23_P14	NM_01502	KIAA0251	KIAA0251,	-1.95365	1.04E-23	0.51186	2995.025	1532.74
A_23_P21	NM_01661	hypothetica	DKFZp434	-1.95405	0.00025	0.51176	383.4257	197.7002
A_24_P20	AL050007	Homo sapi	NUP50,NP	-1.95502	0.00002	0.5115	689.6119	372.6397
A_32_P69	AK025631	Polymeras	POLH	-1.95582	0.00075	0.51129	372.3831	195.7436
A_24_P28	NM_03322	purine-rich	PURB,PUF	-1.95612	4.68E-20	0.51122	4200.166	2148.205
A_23_P13	NM_05304	hypothetica	MGC14327	-1.95644	8.85E-40	0.51113	3650.823	1866.13
A_23_P28	NM_01474	Homo sapi	SNX17,KIA	-1.95745	1.35E-14	0.51087	1091.94	560.4865
A_23_P12	NM_01639	p53-inducib	P53CSV,W	-1.95773	1.73E-23	0.5108	3382.649	1727.851
A_24_P227121	Unknown	THC19677		-1.95777	0.00004	0.51078	474.9184	248.366
A_23_P21	NM_00454	Homo sapi	NDUFC2,B	-1.95799	4.35E-15	0.51073	47618.79	24529.54
A_32_P23	NM_00737	TAR DNA b	TARDBP,T	-1.95844	1.20E-15	0.51061	34496.53	17801.21
A_23_P69	NM_00177	Homo sapi	CD47,IAP,	-1.95856	1.87E-41	0.51058	5380.141	2745.716
A_23_P36	NM_13907	delta-notch	DNER	-1.95918	8.00E-10	0.51042	1520.781	762.0907
A_23_P20	AK027166	Lung canc	HLC-8	-1.95971	3.66E-17	0.51028	1476.491	749.5327
A_32_P37	AK096908	CDNA FLJ	AK096908	-1.95979	8.42E-18	0.51026	2083.64	1065.637
A_23_P53	NM_13920	Homo sapi	NAP1L1,NI	-1.9598	2.69E-28	0.51025	25231.68	12933.48
A_24_P73	NM_01415	HSPC065 p	HSPC065,	-1.96035	0.00008	0.51011	433.0604	224.9501
A_23_P16	NM_01209	adaptor pro	APPL,APP	-1.96143	7.57E-15	0.50983	1940.49	996.334
A_23_P16	NM_00611	Homo sapi	PRAME,MA	-1.96176	1.12E-24	0.50975	2342.997	1193.28
A_24_P12	NM_00474	discs, large	DLG5,PDL	-1.96185	7.86E-35	0.50972	3130.125	1595.432
A_23_P96	NM_00690	Homo sapi	PRKDC,HY	-1.96316	1.60E-25	0.50938	2500.005	1270.915
A_24_P90	BX640898	THUMP do	THUMPD1	-1.96405	0.00001	0.50915	1186.977	627.7233

A_23_P43	BC039468	Nucleopori	NUP50	-1.96485	6.08E-12	0.50894	3133.892	1586.328
A_23_P10	NM_01900	hypothetica	FLJ20232,	-1.96497	8.90E-31	0.50891	6785.486	3458.904
A_24_P18	NM_00347	protein tyro	PTP4A2,H	-1.96518	6.23E-20	0.50886	2088.65	1059.903
A_23_P15	NM_01431	integral inn	LEMD3,MA	-1.96519	8.30E-07	0.50886	785.1935	415.2617
A_24_P29	NM_02053	Homo sapi	C20orf3,BS	-1.96603	5.39E-07	0.50864	622.106	312.215
A_23_P40	NM_15240	Homo sapi	GRPEL2,F	-1.96606	1.91E-09	0.50863	1933.882	966.1592
A_32_P22	AK021705	Hypothetic	LOC13988	-1.96663	1.48E-07	0.50849	1430.636	704.0258
A_32_P20	NM_00682	tyrosine 3-r	YWHAQ,1(	-1.96678	1.39E-21	0.50845	45613.86	23310.89
A_23_P14	NM_00038	Homo sapi	BLMH,BH,	-1.96734	4.32E-35	0.5083	11790.36	5979.98
A_23_P165819		Unknown	A_23_P16	-1.96735	2.65E-32	0.5083	162706	82924.45
A_23_P20	NM_00347	Homo sapi	CUL5,VAC	-1.9679	0.00001	0.50815	734.7335	353.2716
A_24_P55	NM_00016	gap junctio	GJA1,ODD	-1.96795	3.50E-15	0.50814	1706.187	861.4295
A_23_P15	NM_03310	Homo sapi	PNPT1,OL	-1.96821	6.20E-28	0.50808	6192.844	3156.683
A_24_P92	BC019824	Homo sapi	BC019824	-1.96905	6.25E-12	0.50786	1352.191	677.1108
A_23_P11	NM_02453	hypothetica	FLJ12118,	-1.96917	0.00032	0.50783	426.7257	225.9659
A_24_P192338		AF167441	THC18072	-1.96983	1.69E-09	0.50766	863.3955	446.0859
A_24_P20	NM_00645	multifunctic	PAICS,AIR	-1.9699	1.56E-43	0.50764	8167.323	4145.681
A_24_P20	NM_00420	vesicle traf	SEC22L3,S	-1.9702	2.79E-24	0.50756	2137.896	1087.206
A_24_P89	NM_00078	deoxycytidi	DCK,MGC	-1.97099	4.64E-11	0.50736	783.5008	397.285
A_23_P33	BC033450	KIAA1423	KIAA1423	-1.97101	1.30E-08	0.50735	758.403	380.6232
A_24_P11	BM930591	Transcribe	BM930591	-1.97129	0.00007	0.50728	619.5475	295.0344
A_24_P41	NM_00111	adenylate c	ADCY7,KIA	-1.97133	2.59E-19	0.50727	1406.422	713.911
A_24_P39	NM_13842	H63 breast	CASC4,H6	-1.97228	1.75E-19	0.50703	3838.088	1942.592
A_23_P51	BC004501	Hypothetic	MGC33867	-1.97252	2.85E-16	0.50697	6044.507	3078.732
A_24_P56	AK024346	Peroxisom	PPARGC1	-1.97271	2.98E-10	0.50692	932.3085	466.0937
A_24_P82	NM_01868	zinc finger	ZNF313,RI	-1.97311	2.52E-25	0.50681	7612.965	3867.667
A_24_P19	NM_13895	Homo sapi	VANGL1,L	-1.97386	1.53E-35	0.50662	2648.347	1342.047
A_23_P46	NM_02236	MMS19 (M	MMS19L,M	-1.97476	9.74E-27	0.50639	3222.184	1632.335
A_23_P33	NM_00282	Homo sapi	PTPN1,PT	-1.97564	4.17E-13	0.50617	1496.753	766.5602
A_32_P47285		ALU7_HUM	THC18713	-1.97577	0.00012	0.50613	479.5934	233.622
A_24_P81	NM_00406	cyclin-depe	CDKN1B,K	-1.97637	7.63E-09	0.50598	1147.317	594.5122
A_24_P32	NM_02194	FLJ12716	FLJ12716	-1.97639	2.22E-06	0.50597	586.6877	289.4128
A_32_P67	NM_00416	succinate c	SDHA,FP,S	-1.9765	2.74E-20	0.50594	5100.548	2589.904
A_23_P16	NM_00364	Homo sapi	IKBKAP,FD	-1.9766	2.68E-25	0.50592	3683.999	1863.573
A_24_P22	NM_03092	hypothetica	ANP32E,LA	-1.97702	2.86E-20	0.50581	4256.109	2153.252
A_23_P20	NM_00645	Homo sapi	SPFH1,KE	-1.97727	9.11E-18	0.50575	6519.933	3304.769
A_23_P14	NM_01434	Homo sapi	ZNF318,TZ	-1.9773	3.08E-10	0.50574	785.3355	394.6791
A_24_P35	NM_01837	hypothetica	FLJ11259	-1.97735	3.91E-09	0.50573	1837.316	917.2576
A_32_P20	BC037983	Hypothetic	AMMECR1	-1.97742	1.12E-16	0.50571	1415.391	711.9025
A_23_P12	NM_02408	Homo sapi	KDELC1,E	-1.97769	0	0.50564	1211.951	613.4861
A_23_P50	NM_00604	Homo sapi	RBM12,SW	-1.97827	1.65E-09	0.50549	1223.701	605.6965
A_24_P17	NM_00664	WAS prote	WASF3,SC	-1.97878	2.49E-07	0.50536	1001.454	488.3577
A_32_P11	NM_15267	hypothetica	FLJ34969	-1.97885	2.06E-41	0.50534	3979.843	2011.445
A_32_P82	AK021751	Nucleosom	NAP1L1	-1.97967	4.38E-22	0.50513	4821.024	2436.117
A_23_P12	NM_02512	Homo sapi	RNF34,RF	-1.97973	7.68E-17	0.50512	2982.19	1503.408
A_24_P37	NM_00193	Homo sapi	DLAT,DLT	-1.97978	8.04E-10	0.50511	1595.098	790.4356
A_23_P40	NM_20345	Homo sapi	CAMSAP1	-1.98072	6.15E-21	0.50487	1612.786	818.2775
A_23_P25	NM_01844	TBP-intera	CAND1,TIF	-1.98087	1.36E-17	0.50483	1692.352	858.6334
A_24_P30	AK055441	O-sialoglyc	OSGEPL1	-1.98122	0.00005	0.50474	531.6535	257.324
A_23_P42	NM_01451	Homo sapi	ZNF232,ZS	-1.9814	0.00008	0.50469	592.6549	318.9808
A_24_P29	NM_01542	aminoadipa	AASDHPP	-1.98242	2.92E-14	0.50443	6703.476	3397.377
A_23_P15	NM_00375	eukaryotic	EIF3S9,PR	-1.98328	0	0.50421	22224.19	11205.8

A_23_P42	NM_01518	KIAA0746	KIAA0746,	-1.98395	1.12E-35	0.50404	4453.284	2245.287
A_24_P94	AB037796	Programm	PDCD6IP	-1.9842	7.83E-06	0.50398	478.4255	240.5972
A_24_P26	AK024263	Solute carr	SLC38A1	-1.98469	4.04E-25	0.50386	17562.62	8896.918
A_24_P221485		Unknown	A_24_P22	-1.98559	9.72E-09	0.50363	815.5795	419.6242
A_24_P84711		Unknown	A_24_P84	-1.98628	1.23E-13	0.50345	1213.114	605.368
A_23_P25	NM_00360	Homo sapi	SMARCA5	-1.98643	1.09E-11	0.50342	1335.397	660.8353
A_23_P35	NM_00566	Homo sapi	C5orf18,DI	-1.98675	0	0.50334	7865.085	3959.542
A_24_P63	AL049321	Interleukin-	IRAK1BP1	-1.98709	0.00005	0.50325	420.7448	211.3828
A_32_P14	AL831990	Hypothetic	LOC13788	-1.98739	0.00012	0.50317	626.864	292.1119
A_24_P94	AB032988	Hypothetic	DJ971N18	-1.98745	7.49E-15	0.50316	2305.775	1155.187
A_24_P39	NM_01867	smooth mu	SMAP-1,FI	-1.98749	0.0006	0.50315	2404.749	1212.243
A_24_P93	AK090752	Casein kin	CSNK2A2	-1.9879	1.91E-30	0.50304	6621.513	3323.094
A_23_P41	NM_01610	Homo sapi	ZFR	-1.98908	1.13E-40	0.50275	4513.326	2268.353
A_23_P84	NM_00631	Homo sapi	NPEPPS,F	-1.98997	3.14E-15	0.50252	1473.305	734.635
A_32_P12	NM_00310	sorbitol de	SORD,SOI	-1.99063	1.18E-27	0.50235	5732.12	2888.422
A_23_P13	NM_14503	Homo sapi	FAM55C,M	-1.99158	1.65E-14	0.50211	2250.483	1148.588
A_23_P46	AK126751	NHL repea	NHLRC2	-1.99193	2.21E-13	0.50203	899.9687	453.9589
A_23_P17	NM_02515	Nucleotide	NUBPL	-1.99202	0.00009	0.502	546.5543	291.6539
A_23_P69	NM_01819	Homo sapi	LEPREL1,I	-1.99234	6.83E-28	0.50192	2885.691	1449.031
A_32_P72	BX537987	Tripartite m	TRIM59	-1.99308	7.18E-18	0.50174	1514.629	757.6058
A_23_P88	NM_00390	Homo sapi	CDC16,AP	-1.99317	1.34E-17	0.50171	1944.834	979.2167
A_23_P38	NM_00503	peptidylpro	PPID,CYPI	-1.99318	7.26E-17	0.50171	6806.238	3432.023
A_23_P14	NM_00379	membrane	MBTPS1,S	-1.9934	1.71E-41	0.50165	3277.366	1643.975
A_24_P64	AK002197	Kelch dom	KIAA1340	-1.99345	1.61E-15	0.50164	1270.931	634.0071
A_24_P35	AB011087	KIAA0515	KIAA0515	-1.99351	4.92E-29	0.50163	3791.871	1900.933
A_24_P92	U31733	Hypothetic	hprt	-1.9936	5.55E-33	0.5016	4383.151	2197.934
A_23_P72	AK095357	ELMO dom	ELMOD2	-1.99367	6.12E-14	0.50159	1846.761	915.9998
A_23_P99	NM_00212	high-mobili	HMGB1,HM	-1.99579	3.61E-15	0.50105	16185.33	8166.509
A_23_P18	NM_00699	polymerase	POLS,POL	-1.99668	8.81E-25	0.50083	2080.388	1043.371
A_23_P21	NM_02451	Homo sapi	FYCO1,RU	-1.99724	1.86E-06	0.50069	717.7915	345.5385
A_23_P33	NM_15301	hypothetica	FLJ30596,	-1.99724	4.99E-08	0.50069	958.3194	465.1274
A_23_P40	NM_03238	Hermansky	HPS3,SUT	-1.99727	2.35E-12	0.50068	1318.183	651.428
A_24_P19	NM_01323	Ste-20 rela	STK39,DC	-1.99815	0.00063	0.50046	361.3866	185.8101
A_24_P39	NM_01476	translocase	TOMM20,M	-1.99821	8.78E-27	0.50045	4767.625	2391.108
A_23_P39	NM_02092	Ninein (GS	NIN	-1.99853	7.09E-06	0.50037	614.138	320.8317
A_24_P36	AF119896	KIAA0738	KIAA0738	-1.99872	1.25E-21	0.50032	2000.089	997.5206
A_24_P39	NM_00270	protein pho	PPP1CB,P	-1.99911	1.53E-17	0.50022	6142.093	3083.234
A_24_P200848		Unknown	THC18696	-1.99921	0.00019	0.5002	475.7177	226.4106
A_32_P10	BX641009	Spermatog	SPATA5	-1.99969	0.00002	0.50008	459.3431	228.9291
A_23_P21	NM_00670	Homo sapi	GLO1,GLY	-2.00041	3.13E-09	0.4999	784.0877	395.6533
A_24_P23	NM_00103	ribonucleot	RRM2,R2,I	-2.00049	1.12E-22	0.49988	5194.925	2601.07
A_24_P23	NM_00005	branched c	BCKDHB,E	-2.00062	0.00002	0.49985	765.7692	359.2032
A_23_P97	NM_00647	Homo sapi	TXNIP,THI	-2.00122	1.21E-23	0.49969	1805.877	898.8942
A_23_P33	NM_05716	Homo sapi	GIT2,CAT-	-2.0021	1.49E-11	0.49948	2393.638	1218.686
A_23_P15	NM_00437	COX11 (ye	COX11,CC	-2.00275	1.03E-16	0.49931	7588.376	3809.651
A_23_P14	NM_01813	Homo sapi	C14orf104,	-2.00347	1.75E-21	0.49913	3510.271	1751.363
A_24_P53	NM_13932	Homo sapi	ATRN,MG	-2.00391	1.67E-06	0.49902	679.7806	352.2449
A_32_P51	AK022869	Ribosomal	RPS6KA3	-2.00404	3.20E-20	0.49899	7061.136	3544.766
A_23_P83	AK022617	IGF-II mRN	IMP-1	-2.00496	1.36E-34	0.49876	1993.952	998.642
A_23_P204170		U09086 th	THC18110	-2.00624	7.39E-15	0.49845	2375.606	1176.733
A_24_P56	NM_01910	beta-1,4 m	ALG1,HMT	-2.0063	5.12E-11	0.49843	1410.74	688.9081
A_23_P87	NM_00683	ATP/GTP-b	HEAB,CLP	-2.0068	4.97E-15	0.49831	1067.2	530.9326

A_24_P47	AL117621	ADP-ribosy	ARF6	-2.00684	9.57E-19	0.49829	1567.359	776.6838
A_23_P25	NM_00671	activator of	ASK,DBF4	-2.00734	2.83E-20	0.49817	2037.517	1022.746
A_23_P99	NM_01639	butyrate-in	HSPC121,	-2.00942	5.13E-43	0.49766	23960.48	11946.17
A_23_P15	NM_15269	Homo sapi	C1GALT1	-2.00967	2.80E-45	0.49759	5135.968	2555.112
A_23_P17	NM_00327	Homo sapi	TSPAN6,T	-2.01094	2.78E-29	0.49728	2969.852	1477.042
A_23_P37	NM_01491	Homo sapi	CHSY1,KI	-2.01109	6.11E-34	0.49724	4379.935	2175.621
A_23_P78	NM_01608	CGI-150 pr	C17orf25,H	-2.0112	1.31E-22	0.49721	5714.581	2842.063
A_23_P40	NM_00737	Homo sapi	TARDBP,T	-2.0117	1.91E-08	0.49709	878.7212	425.1347
A_23_P21	NM_01632	RAB14, me	RAB14,FB	-2.01188	8.98E-11	0.49705	1745.803	882.294
A_23_P35	NM_01786	Homo sapi	PIGX,FLJ2	-2.01255	1.05E-06	0.49688	567.3984	278.3853
A_24_P25	BC022429	Hypothetic	LOC90624	-2.01297	9.32E-09	0.49678	1286.071	620.8154
A_23_P17	NM_00440	neural prec	SEPT2,DIF	-2.01415	2.80E-45	0.49649	6981.416	3468.503
A_32_P192480		Unknown	THC18692	-2.01439	7.63E-08	0.49643	1545.807	743.7972
A_32_P60	NM_00410	coagulatior	F2RL2,PAI	-2.01594	3.32E-18	0.49605	1902.141	938.9692
A_23_P13	NM_01645	coatomer p	COPB,DKF	-2.01734	2.31E-42	0.4957	3913.291	1940.188
A_32_P45	NM_00262	Phosphogl	PGAM1	-2.01818	1.14E-37	0.4955	19379.33	9594.438
A_23_P15	NM_01824	Homo sapi	NEIL3,FPG	-2.01834	4.28E-07	0.49546	750.0456	358.9876
A_24_P20	NM_03056	hypothetica	C6orf148,M	-2.01886	0.00007	0.49533	484.3002	231.1637
A_23_P98	NM_00132	Homo sapi	CSTF3,CS	-2.01924	8.57E-18	0.49524	4626.47	2278.636
A_23_P20	NM_00478	ubiquitinati	UBE4A,E4	-2.01932	1.46E-30	0.49522	3548.796	1757.13
A_24_P127462		Unknown	A_24_P12	-2.01984	0	0.49509	12035.7	5959.24
A_23_P90	NM_00591	Homo sapi	MCM6,Mis	-2.01999	1.56E-13	0.49505	12463.2	6203.825
A_32_P35	AK092508	Methyl-Cp	MBD3	-2.02012	2.22E-07	0.49502	861.4459	411.4827
A_23_P15	BC010144	Likely ortho	LARP	-2.02045	6.35E-36	0.49494	24249.48	12030.83
A_23_P58	AK054622	RGM doma	RGMB	-2.02131	2.39E-12	0.49473	860.0046	425.7502
A_23_P14	NM_00410	eyes abser	EYA4,CMD	-2.02188	0.00069	0.49459	593.6495	327.5664
A_24_P55	NM_14580	Homo sapi	MTPN,V-1,	-2.02246	1.80E-39	0.49445	4684.302	2313.74
A_23_P42	NM_00295	Homo sapi	RXRA,NR2	-2.02278	2.80E-23	0.49437	4680.645	2312.529
A_24_P71938		Unknown	THC18079	-2.02335	3.87E-20	0.49423	1934.832	952.3538
A_23_P10	NM_01490	Homo sapi	TMEM15,K	-2.02393	1.44E-15	0.49409	2139.448	1051.75
A_32_P17	AK091332	CDNA FLJ	AK091332	-2.02412	7.67E-26	0.49404	3114.419	1536.743
A_24_P94	AB051511	Selenoprot	SELI	-2.02456	1.09E-24	0.49393	4319.44	2133.573
A_32_P23	AF132203	Stearoyl-C	SCD	-2.02548	7.41E-31	0.49371	4108.881	2026.829
A_24_P30	L40403	YLP motif	YLPM1	-2.02576	0.00001	0.49364	615.7565	290.251
A_24_P926993		unnamed p	NP181386	-2.02671	4.36E-09	0.49341	782.3521	379.8255
A_23_P15	NM_02499	Homo sapi	GFM1,EFG	-2.02675	7.13E-24	0.4934	3242.58	1593.101
A_23_P41	NM_01403	DKFZP586	DKFZP586	-2.02694	1.76E-22	0.49336	2624.119	1292.482
A_23_P37	NM_17743	Homo sapi	DICER1,Di	-2.02811	2.42E-31	0.49307	1982.103	977.4878
A_32_P112623		BC032035	THC18073	-2.02832	5.72E-11	0.49302	792.6007	390.9905
A_23_P95	NM_18157	Homo sapi	RFC5,RFC	-2.02873	2.46E-27	0.49292	8519.134	4214.941
A_24_P57	NM_01503	Homo sapi	ZHX3,TIX1	-2.02896	5.41E-07	0.49286	810.6516	417.0577
A_24_P40	NM_03121	sec13-like	SEH1L,Sel	-2.02908	7.17E-17	0.49283	3135.375	1542.897
A_23_P43	NM_14457	Homo sapi	C16orf34,L	-2.02925	0	0.49279	14293.71	7047.032
A_23_P15	NM_02077	Homo sapi	TBC1D14,I	-2.02946	2.44E-22	0.49274	1758.657	868.7343
A_23_P43	NM_01799	Homo sapi	C9orf40,FL	-2.03157	4.31E-09	0.49223	1028.638	493.5545
A_23_P20	NM_00271	protein pho	PPP1CC,P	-2.03198	0	0.49213	5446.713	2680.537
A_23_P21	NM_00323	Homo sapi	TFRC,TFR	-2.0328	0	0.49193	20358.64	10021.95
A_23_P98	NM_00589	Homo sapi	M11S1,GP	-2.03283	4.08E-22	0.49192	5668.008	2772.994
A_23_P99	NM_00283	Homo sapi	PTPN11,C	-2.03306	2.61E-22	0.49187	6789.953	3342.107
A_32_P235159		AF117065	THC19025	-2.03346	2.64E-13	0.49177	963.1455	476.2451
A_32_P95914		Unknown	THC19244	-2.03451	3.04E-06	0.49152	721.4478	372.4296
A_23_P32	NM_02023	Homo sapi	CDC42SE	-2.03468	1.00E-23	0.49148	2837.92	1392.62

A_23_P12	NM_01487	Homo sapi	MFN2,HSG	-2.03487	1.61E-20	0.49143	17102.67	8454.374
A_23_P76	AK096242	Dynactin 2	DCTN2	-2.03501	5.09E-13	0.4914	2543.757	1241.262
A_23_P35	AB020684	KIAA0877	KIAA0877	-2.03506	4.20E-12	0.49139	1318.542	638.1602
A_23_P16	NM_00293	ring finger	RNF4,SNU	-2.03586	5.89E-26	0.49119	5568.866	2738.768
A_24_P25	NM_00605	Down synd	DSCR3,DC	-2.03602	2.57E-09	0.49115	792.8147	383.8226
A_23_P11	NM_00485	Clathrin, he	CLTC	-2.03648	7.54E-35	0.49104	21830.43	10725.42
A_32_P36046		ALU8_HUM	THC18442	-2.03757	5.25E-07	0.49078	537.2845	263.0236
A_32_P22	BC045174	CDNA FLJ	BC045174	-2.03768	0.00002	0.49075	530.1108	249.9692
A_23_P87	NM_00103	Homo sapi	RRM1,R1,I	-2.03793	4.01E-25	0.4907	20348.35	10023.97
A_24_P37	NM_01774	hypothetica	ZDHHC7,Z	-2.03834	3.99E-25	0.4906	2932.776	1438.584
A_32_P14	NM_00643	ADP-ribosy	PARP4,PH	-2.03855	5.40E-13	0.49055	822.1212	403.4261
A_23_P61	NM_00040	galactosida	GLB1	-2.03881	0	0.49048	8778.451	4306.082
A_24_P15	NM_01231	LIV-1 prote	SLC39A6,I	-2.03973	2.33E-12	0.49026	891.7347	434.8378
A_23_P73	NM_20584	Homo sapi	NCKAP1,H	-2.04038	2.08E-13	0.4901	2325.442	1132.285
A_32_P204205		Unknown	THC19329	-2.04064	0.00002	0.49004	451.9474	219.7038
A_23_P15	AB014537	Zinc finger,	ZBED4	-2.04227	0.00025	0.48965	450.687	234.4326
A_23_P13	NM_01478	Homo sapi	ARHGAP1	-2.04263	8.50E-06	0.48956	1204.924	622.9415
A_24_P38	NM_01642	cisplatin re	CROP,LUC	-2.04291	1.53E-24	0.4895	7155.956	3518.811
A_23_P54	NM_00568	ATP-bindin	ABCB6,AB	-2.04323	9.63E-41	0.48942	6399.159	3127.983
A_32_P25	NM_00104	solute carri	SLC12A2,E	-2.04468	0.00014	0.48907	1378.646	703.7279
A_23_P97	NM_00023	lipase A, ly	LIPA,LAL,C	-2.04503	0	0.48899	6921.343	3385.979
A_32_P64461		Unknown	ENST0000	-2.04531	0.00047	0.48892	414.851	216.3237
A_23_P77	NM_00390	Homo sapi	APPBP1,H	-2.04676	2.71E-36	0.48858	33920.36	16609.09
A_24_P13	NM_02072	Homo sapi	FAM62B,C	-2.04732	4.68E-19	0.48844	1402.364	687.2748
A_23_P82	NM_01425	solute carri	SLC25A13	-2.04756	5.30E-18	0.48839	1896.058	921.2857
A_23_P74	AB020714	KIAA0907	KIAA0907	-2.04944	4.48E-25	0.48794	1861.24	906.2324
A_24_P47	BM550834	Transcribe	BM550834	-2.05036	5.47E-11	0.48772	764.4845	371.9067
A_23_P38	NM_00141	Homo sapi	ELAVL1,HU	-2.05038	4.16E-06	0.48772	611.5895	287.6874
A_24_P21	NM_01818	hypothetica	DCUN1D2,	-2.05198	0.00058	0.48733	373.9158	176.6254
A_24_P349580		Unknown	A_24_P34	-2.05269	4.44E-10	0.48717	1124.869	535.8795
A_23_P20	NM_00443	Homo sapi	EML1,EMA	-2.05292	1.04E-12	0.48711	1140.97	562.5866
A_23_P13	NM_03322	Homo sapi	PURB,PUF	-2.0533	1.02E-11	0.48702	990.6513	475.0982
A_24_P15	NM_00719	chromatin-s	SUPT16H,	-2.05362	5.36E-20	0.48695	3117.723	1516.912
A_24_P306921		Unknown	A_24_P30	-2.05378	1.02E-38	0.48691	3946.329	1922.072
A_24_P827096		Unknown	THC18321	-2.05487	4.16E-31	0.48665	2436.916	1187.433
A_23_P25	NM_02409	defective ir	DCC1,MGC	-2.05549	3.92E-29	0.4865	906.2749	442.3604
A_24_P33	D86960	Lysophosp	LPGAT1	-2.05603	4.42E-15	0.48637	1725.539	846.8225
A_24_P14	NM_01486	KIAA0475	FAM20B	-2.0588	7.57E-30	0.48572	12556.38	6115.28
A_23_P72	AK000681	Hypothetic	FLJ20674	-2.05918	9.00E-06	0.48563	496.2869	235.6589
A_23_P33	BC024231	Hypothetic	MGC27345	-2.05948	3.91E-14	0.48556	1393.103	668.2334
A_23_P22	NM_03266	Homo sapi	FUT10,MG	-2.06034	0.00004	0.48536	484.5423	245.6571
A_23_P20	NM_00589	Homo sapi	GOLGA3,M	-2.06036	4.67E-32	0.48535	8031.474	3903.302
A_23_P12	NM_01813	Homo sapi	HRMT1L6,	-2.06143	1.45E-13	0.4851	2644.686	1277.403
A_23_P39	NM_14460	hypothetica	FLJ31153	-2.06225	4.64E-21	0.48491	13760.27	6677.056
A_24_P595877		ALU1_HUM	THC18221	-2.06244	0.00073	0.48486	315.1743	153.3738
A_23_P13	AL132665	BCL2/aden	BNIP3L	-2.06373	7.49E-08	0.48456	680.3719	321.9872
A_24_P77	AK093903	HBV preS1	PS1TP4	-2.06424	1.86E-07	0.48444	698.096	348.2758
A_24_P17	AL834364	Hypothetic	DKFZp762	-2.06512	2.39E-07	0.48423	666.3955	314.4635
A_23_P50	NM_03239	Homo sapi	MKI67IP,N	-2.0661	8.99E-40	0.484	4670.776	2260.06
A_23_P37	AL834325	GTPase ac	DKFZP434	-2.06672	7.22E-10	0.48386	778.8877	372.0465
A_23_P31	NM_13364	sterile alph	ZAK,MLTK	-2.06703	5.26E-16	0.48379	1671.751	814.4842
A_24_P91	AK023165	SMILE prot	SMILE	-2.06725	2.60E-08	0.48374	2318.213	1105.082

A_24_P16	NM_00581	protein kina	PRKD3,EP	-2.06726	3.21E-14	0.48373	1254.12	612.9668
A_24_P599496		Unknown	THC20035	-2.06784	8.55E-13	0.4836	1164.978	556.5496
A_24_P35	AK057071	Homo sapi	AK057071	-2.06792	9.45E-16	0.48358	32816.6	15971.72
A_23_P21	NM_02290	Homo sapi	SLC30A5,Z	-2.06908	4.84E-17	0.48331	2560.856	1234.403
A_23_P36	NM_01222	myosin IB	MYO1B,my	-2.06926	4.33E-13	0.48326	1164.253	554.7483
A_24_P61	AL833755	Chromosor	C6orf89	-2.0694	7.30E-34	0.48323	3496.473	1689.981
A_23_P65	NM_00240	Homo sapi	MGAT2,GN	-2.06948	4.94E-20	0.48321	6193.119	2996.265
A_24_P56	AL832817	Dual-specif	DYRK2	-2.06967	3.36E-14	0.48317	1133.21	541.7379
A_24_P20	NM_00046	ubiquitin pr	UBE3A,AS	-2.06986	2.25E-13	0.48312	964.1836	462.7123
A_23_P18	NM_00067	Homo sapi	ADH5,FDH	-2.07128	0.00002	0.48279	754.9767	339.149
A_24_P17	XM_37226	similar to D	LOC38990	-2.07216	0.00051	0.48259	348.6399	170.5483
A_23_P14	NM_01812	hypothetica	DARS2,FL	-2.07263	3.29E-27	0.48248	2017.59	976.8882
A_23_P42	NM_00717	Homo sapi	CIT,CRIK,S	-2.07326	3.68E-36	0.48233	2790.115	1345.315
A_32_P228254		Unknown	A_32_P228	-2.07326	1.99E-08	0.48233	2388.929	1153.406
A_23_P62	NM_00031	Homo sapi	PPT1,PPT	-2.07371	6.87E-27	0.48223	11458.29	5543.533
A_23_P12	NM_00384	Homo sapi	PEX11B,PI	-2.07408	7.31E-10	0.48214	887.3667	440.3904
A_32_P11	AL137938	PX domain	PXK	-2.07459	1.67E-29	0.48202	3745.518	1806.834
A_24_P27	AF284094	BRI3 bindir	BRI3BP	-2.07535	4.65E-14	0.48185	1270.617	605.3449
A_24_P29	NM_00712	valosin-cor	VCP,p97,T	-2.07674	9.68E-38	0.48152	5426.03	2606.876
A_24_P33	NM_00517	ATPase, H	ATP6V0A1	-2.07803	2.26E-36	0.48122	3411.326	1641.308
A_23_P97	BC007072	Hypothetic	MGC12538	-2.07829	3.10E-07	0.48116	793.0558	366.8606
A_24_P17	AY029066	Homo sapi	HN1	-2.08001	2.53E-09	0.48077	546217	263620.4
A_23_P44235		AF098788	THC18430	-2.0811	6.79E-33	0.48051	2576.764	1237.414
A_32_P19	AK056809	CDNA FLJ	LOC11606	-2.08171	2.62E-16	0.48037	1568.188	759.594
A_23_P20	NM_03283	Homo sapi	ALG10,KC	-2.08252	0.00017	0.48019	377.8138	182.8492
A_32_P32	BC036626	SUMO1/se	SENP5	-2.08347	2.88E-22	0.47997	2082.851	995.4083
A_24_P23	NM_03091	hypothetica	SNX27,MY	-2.08571	3.26E-22	0.47945	1527.636	734.9852
A_24_P28	BX537360	Cleavage a	CPSF5	-2.08639	1.29E-15	0.4793	2896.633	1381.405
A_23_P39	NM_15268	hypothetica	RWDD4A,I	-2.08835	1.67E-13	0.47885	1532.558	724.6447
A_32_P66	BQ926066	AGENCOL	BQ926066	-2.08849	0.00008	0.47882	591.0867	262.1307
A_24_P21	BC028940	Sterol O-ac	SOAT1	-2.08869	2.81E-14	0.47877	2296.334	1089.628
A_24_P28	NM_02431	MGC4170	GNPTAB,I	-2.08944	1.09E-12	0.4786	1506.373	732.1993
A_24_P21	NM_00437	COX11 (ye	COX11,CC	-2.08965	3.46E-24	0.47855	3816.709	1821.87
A_23_P21	NM_00721	Homo sapi	SEC63,ER	-2.08969	1.64E-29	0.47854	2541.375	1217.751
A_23_P35	NM_00272	Homo sapi	PPP6C	-2.08989	1.64E-31	0.47849	3967.764	1900.158
A_23_P93	NM_01445	Homo sapi	SESN1,PA	-2.09254	1.53E-06	0.47789	490.4329	235.0381
A_24_P12	NM_03137	heterogene	HNRPD,P3	-2.09276	5.24E-13	0.47784	963.0239	455.4251
A_32_P35	NM_01837	hypothetica	FLJ11273	-2.09308	1.53E-10	0.47776	3260.585	1546.127
A_24_P40	NM_00270	protein pho	PPP1CB,P	-2.09424	5.51E-22	0.4775	3074.073	1468.265
A_24_P38	NM_01817	hypothetica	P15RS,FL	-2.09456	7.36E-19	0.47743	2109.13	1001.621
A_32_P21	NM_00079	dihydrofola	DHFR	-2.09502	4.04E-31	0.47732	4947.14	2367.079
A_24_P92	AK022030	Sjogren sy	SSA2	-2.09605	9.25E-14	0.47709	7303.789	3504.404
A_24_P14	XM_37167	similar to p	LOC38917	-2.09794	0.00011	0.47666	441.9216	221.1115
A_32_P37	AK098081	Exportin, tF	XPOT	-2.0985	1.00E-38	0.47653	7034.472	3354.805
A_32_P232237		Unknown	THC18718	-2.09948	0.00003	0.47631	649.4692	287.2314
A_32_P11	AK094088	Aldehyde d	FLJ38508	-2.10029	1.32E-10	0.47612	941.2394	456.6938
A_24_P33	NM_00717	unr-interac	STRAP,MA	-2.10122	3.50E-17	0.47591	8725.006	4153.267
A_23_P21	NM_01478	Homo sapi	TRIM14,KI	-2.10148	2.83E-22	0.47585	1268.313	604.5252
A_24_P33	NM_00588	macrophag	MAEA,EMI	-2.1016	6.71E-38	0.47583	6725.229	3201.793
A_24_P39	NM_03195	MSTP028	KCTD10,U	-2.10307	0.00012	0.4755	486.4216	246.7683
A_24_P918875		Unknown	THC20024	-2.10453	0	0.47517	37857.38	17988.36
A_24_P23	AB040958	BTB (POZ)	BTBD7	-2.10513	5.28E-26	0.47503	1784.318	845.6123

A_23_P14	NM_00379	membrane	MBTPS1,S	-2.10544	4.98E-15	0.47496	3787.091	1794.606
A_23_P96	NM_02071	Homo sapi	LRRC47,K	-2.10603	3.75E-14	0.47483	2299.41	1080.797
A_32_P45	AF037219	Ubiquitin p	UBE3A	-2.10605	0.00001	0.47482	677.9485	301.8124
A_23_P11	NM_01835	acheron	LARP6,FL	-2.10703	3.37E-19	0.4746	1441.495	680.499
A_23_P37	NM_01476	24-dehydro	DHCR24,K	-2.10721	7.87E-20	0.47456	1900.28	898.3053
A_23_P37	NM_01623	Homo sapi	ANAPC5,A	-2.10751	3.24E-38	0.47449	4071.711	1929.619
A_24_P28	NM_00156	interleukin	IL13RA1,N	-2.10782	1.54E-20	0.47442	9208.01	4393.54
A_23_P28	NM_00259	Homo sapi	PCNA,MG	-2.10785	2.18E-36	0.47442	38612.28	18282
A_23_P33	NM_18275	hypothetica	FAM79A,F	-2.10908	0.00067	0.47414	373.4529	187.3482
A_32_P44	NM_01411	PRO0149 p	PRO0149	-2.10945	6.51E-16	0.47406	2726.827	1289.422
A_24_P89	NM_03316	Homo sapi	SURF4,ER	-2.11045	6.23E-12	0.47383	12553.33	5974.694
A_24_P37	NM_01642	C3HC4-like	RNF141,ZI	-2.11127	2.42E-07	0.47365	915.7534	414.5106
A_24_P69	AK092810	CDNA FLJ	AK092810	-2.11382	5.48E-20	0.47308	7654.133	3641.701
A_23_P39	NM_15326	hypothetica	FLJ38101,	-2.11412	3.42E-19	0.47301	1174.14	555.6605
A_23_P29	NM_01538	CGI-51 pro	SAMM50,C	-2.11589	9.44E-22	0.47261	7861.559	3734.902
A_32_P92	AK023269	Homo sapi	LYCAT,AL	-2.11756	0	0.47224	4552.17	2149.25
A_24_P36	NM_00272	protein pho	PPP6C	-2.11758	2.36E-09	0.47224	1004.79	487.5962
A_24_P75	AL832683	Protein pre	PTAR1	-2.11915	9.19E-21	0.47189	2936.088	1393.289
A_24_P27	NM_00307	SWI/SNF r	SMARCA3	-2.12017	3.68E-19	0.47166	2066.607	971.0562
A_24_P92	BX648462	Insulin-deg	IDE	-2.12043	3.15E-12	0.4716	2152.461	1002.369
A_23_P36	AK001563	Sno, strawl	SBNO1	-2.12154	4.40E-13	0.47136	843.2678	399.4882
A_24_P17	NM_02043	Homo sapi	PHTF2,MG	-2.12211	1.12E-07	0.47123	829.5415	375.9879
A_23_P35	AL832919	1-acylglyce	AGPAT3	-2.12264	1.67E-20	0.47111	2488.311	1174.941
A_23_P99	NM_00212	Homo sapi	HMGB1,HM	-2.12271	0	0.4711	5601.883	2638.922
A_32_P18	AK095472	Hypothetic	DKFZp762	-2.12365	1.46E-06	0.47089	553.2289	267.5026
A_32_P18	NM_00082	gamma-glu	GGCX,VKC	-2.12699	3.71E-10	0.47015	840.1161	403.8515
A_24_P524562		Unknown	A_24_P52	-2.12791	3.23E-11	0.46994	19100.56	8980.511
A_24_P17	AB007891	Homo sapi	KIAA0431	-2.12891	3.10E-06	0.46972	611.6975	274.7778
A_23_P97	NM_00444	Homo sapi	EPRS,EAR	-2.13049	9.59E-21	0.46938	15781.95	7452.174
A_24_P38	NM_00138	dihydropyri	DPYSL2,D	-2.13144	1.80E-30	0.46917	2158.822	1014.366
A_24_P27	NM_03228	hypothetica	MGC3207	-2.13328	8.71E-08	0.46876	818.8855	368.3997
A_32_P40	NM_00644	U5 snRNP-	PRPF8,PR	-2.13345	1.43E-29	0.46872	7986.646	3752.923
A_24_P115621		ALU8_HUM	THC19007	-2.13594	1.19E-11	0.46818	2907.823	1345.019
A_23_P20	NM_01790	Homo sapi	TMCO3,C1	-2.13741	3.31E-21	0.46786	2421.523	1143.049
A_23_P105753		PWP1_HU	THC18059	-2.13795	1.12E-22	0.46774	2140.088	996.7472
A_32_P166356		Unknown	THC19174	-2.13901	1.74E-16	0.46751	1241.009	575.378
A_24_P26	AK095775	Pleckstrin h	PLEKHA1	-2.13925	1.17E-20	0.46745	2069.281	969.6263
A_23_P14	BC032221	KIAA0391	KIAA0391	-2.13964	1.07E-12	0.46737	1284.401	589.3557
A_23_P80	NM_03228	Homo sapi	FYTTD1,D	-2.14012	1.49E-23	0.46726	1654.965	776.6644
A_24_P198820		Unknown	THC18573	-2.1406	5.77E-06	0.46716	496.2196	226.9837
A_32_P96	NM_02474	hypothetica	SHCBP1,F	-2.14148	6.94E-32	0.46697	1768.878	825.3772
A_23_P61	NM_00040	Homo sapi	GLB1	-2.14303	3.09E-18	0.46663	1666.696	771.8967
A_24_P20	NM_02519	chromoson	C1orf22	-2.14422	3.19E-29	0.46637	2946.652	1371.805
A_23_P42	NM_19806	Homo sapi	GNPNAT1,	-2.14716	2.94E-24	0.46573	3862.235	1790.509
A_23_P25	NM_01863	Homo sapi	WSB2,SBA	-2.14795	4.96E-18	0.46556	10231.52	4790.848
A_23_P12	NM_01233	acyl-Coenz	ACATE2,C	-2.14806	0	0.46554	5680.54	2643.939
A_24_P27	AL117568	Huntingtin	HIP2	-2.14955	2.42E-28	0.46521	2934.475	1365.97
A_23_P47	NM_00618	Homo sapi	OAS3,p10	-2.15337	2.78E-11	0.46439	1004.922	458.259
A_23_P20	NM_19833	Homo sapi	GANAB,G2	-2.15339	3.91E-29	0.46438	5508.943	2547.319
A_24_P854758		Unknown	THC19657	-2.15344	6.43E-34	0.46437	5160.321	2393.365
A_23_P21	NM_07862	Homo sapi	MSL3L1,DI	-2.15401	2.20E-13	0.46425	1205.908	550.5621
A_23_P20	NM_02483	Homo sapi	C10orf119,	-2.15606	1.91E-36	0.46381	3003.717	1393.689

A_24_P221285		Unknown	A_24_P22	-2.15626	7.91E-11	0.46377	1340.489	606.7466
A_32_P49	XM_37174	Homo sapi	XM_37174	-2.15687	1.51E-06	0.46363	651.6552	317.4019
A_23_P20	NM_01328	Homo sapi	UHRF1,Np	-2.15697	2.59E-30	0.46361	5546.915	2571.352
A_23_P72	NM_03288	hypothetica	MGC11308	-2.16079	5.03E-09	0.46279	1245.492	596.1603
A_23_P35	NM_00138	dihydropyri	DPYSL2,D	-2.16115	2.33E-22	0.46272	6361.49	2954.762
A_23_P33	AK095888	CD109 ant	CD109	-2.16188	9.43E-37	0.46256	2593.664	1198.302
A_23_P67	NM_03273	hypothetica	LMNB2,LM	-2.16207	6.45E-22	0.46252	4256.491	1967.382
A_32_P17	NM_01504	Homo sapi	KIAA0261,	-2.16309	3.73E-24	0.4623	3469.22	1600.082
A_23_P56	NM_00677	Homo sapi	DDX18,MrI	-2.16333	8.84E-42	0.46225	2451.209	1137.119
A_23_P21	NM_00339	Homo sapi	WNT5A,hW	-2.16514	3.60E-12	0.46186	2293.667	1047.075
A_24_P25	U90143	Butyrophilin	BTN3A2	-2.16575	6.77E-06	0.46173	568.9909	250.0876
A_24_P93	NM_00550	branched c	BCAT1,BC	-2.16627	3.20E-28	0.46162	3164.886	1460.094
A_23_P11	NM_00379	Homo sapi	C19orf2,RN	-2.16649	6.11E-17	0.46158	2390.883	1098.254
A_24_P39	NM_00166	aryl hydroc	ARNT,HIF1	-2.16722	0.00011	0.46142	417.9269	185.6792
A_23_P11	NM_01603	comparativ	UTP11L,C	-2.16731	7.27E-30	0.4614	6014.342	2782.381
A_24_P64	NM_00680	adaptor-rel	AP3M2,P4	-2.16936	5.61E-06	0.46097	535.1089	257.8752
A_23_P59	NM_00359	Homo sapi	CUL1	-2.16949	1.38E-38	0.46094	3323.389	1532.405
A_24_P33	NM_00117	ADP-ribosy	ARL1,ARF	-2.17071	1.97E-21	0.46068	3646.386	1678.801
A_24_P11	NM_00202	Homo sapi	FN1,FN,CI	-2.17164	3.46E-06	0.46048	598.5871	262.7618
A_24_P22	AK096415	Kelch-like 1	KLHL11	-2.17165	9.30E-12	0.46048	1474.615	692.1941
A_32_P11	BC000226	Ubiquitin sp	USP47	-2.17217	2.55E-11	0.46037	725.6421	332.3933
A_23_P41	AK027663	Stanniocalc	STC2	-2.17344	1.21E-11	0.4601	880.7595	399.9473
A_23_P25	NM_00662	Homo sapi	HBS1L,ER	-2.17379	3.43E-37	0.46003	3521.473	1623.038
A_32_P101264		AB023583	THC19580	-2.17455	7.99E-11	0.45987	853.3228	384.5913
A_23_P82	NM_01217	Homo sapi	FBXO5,EM	-2.17595	3.08E-15	0.45957	2835.638	1290.922
A_23_P37	AL833455	Membrane	MNAB	-2.17786	1.38E-24	0.45917	2184.091	1006.316
A_23_P14	NM_00149	Homo sapi	GCLC,GCS	-2.17837	6.52E-29	0.45906	4007.351	1842.002
A_23_P64	NM_01434	Homo sapi	MTCH2,HS	-2.17953	0	0.45882	16754.17	7701.876
A_23_P25	NM_00568	Homo sapi	ABCC5,MF	-2.18074	3.17E-17	0.45856	1582.233	718.1852
A_24_P38	NM_00234	lanosterol s	LSS,OSC	-2.18198	4.77E-10	0.4583	1468.935	652.1936
A_23_P42	NM_00131	mitogen-ac	MAPK14,R	-2.18219	1.34E-19	0.45826	3442.106	1577.607
A_23_P60	NM_02122	Zinc finger	ZNF462	-2.18241	1.89E-14	0.45821	1493.179	673.9807
A_23_P11	NM_01344	Homo sapi	UBQLN2,D	-2.1825	2.77E-35	0.45819	2090.444	958.8834
A_23_P32	D86985	KIAA0232	KIAA0232	-2.18309	2.95E-09	0.45807	773.78	345.6406
A_32_P22	NM_00589	membrane	M11S1,GP	-2.18343	9.03E-07	0.45799	578.8115	274.6464
A_24_P13	AL359062	Collagen, t	COL8A1	-2.18543	2.85E-06	0.45758	524.9096	248.8873
A_23_P15	NM_00087	insulin-like	IGF2R,MPI	-2.18576	9.80E-09	0.45751	1704.737	755.5736
A_23_P94	NM_01450	Homo sapi	TOR1B,DC	-2.18592	9.97E-07	0.45747	593.931	282.8696
A_23_P25	NM_03194	Homo sapi	CDCA7,JP	-2.186	4.83E-27	0.45746	5881.354	2684.406
A_32_P20	BC048193	Hypothetic	MGC22265	-2.18742	0.00007	0.45716	422.3107	201.4594
A_23_P65	NM_01802	hypothetica	FLJ10213	-2.18948	2.25E-09	0.45673	654.1034	295.6661
A_23_P18	NM_00291	Homo sapi	RFC4,A1,F	-2.1898	7.15E-44	0.45666	13625.06	6201.63
A_23_P33	BC036414	PAI-1 mRN	PAI-RBP1	-2.19185	0	0.45624	4805.225	2190.858
A_23_P76	NM_00500	Homo sapi	NDUFA9,N	-2.19246	1.39E-39	0.45611	5948.832	2705.965
A_24_P35	NM_00043	platelet-act	PAFAH1B1	-2.19325	9.73E-29	0.45594	3518.322	1606.329
A_24_P18	AL133052	Chromosor	C1orf37	-2.19395	5.46E-11	0.4558	1243.688	553.3507
A_24_P36	NM_01594	CGI-11 pro	ATP6V1H,	-2.19418	0.00003	0.45575	502.7327	216.885
A_23_P10	NM_00643	Homo sapi	CCT2,CCT	-2.19482	1.18E-40	0.45562	94899.03	43412.07
A_32_P96692		Unknown	THC18781	-2.19556	0.0001	0.45547	364.9706	167.5132
A_32_P11	AK025056	Hypothetic	FLJ13456	-2.19581	6.39E-20	0.45541	1325.842	601.1307
A_23_P38	AL833405	Transmem	TMEM16F	-2.19684	8.59E-22	0.4552	3634.731	1654.632
A_23_P28	NM_01620	cleavage a	CPSF3,CP	-2.19692	3.11E-28	0.45518	2701.519	1230.525

A_23_P12	NM_00152	Histone de	HDAC2	-2.1973	2.02E-21	0.4551	9925.5	4538.466
A_23_P57	NM_03224	Plexin A1	PLXNA1	-2.19844	9.62E-27	0.45487	4058.393	1841.782
A_32_P22	BX350880	Transcribe	BX350880	-2.19866	7.30E-07	0.45482	11465.11	4988.348
A_24_P67	BC016329	Homo sapi	BC016329	-2.19998	0.00014	0.45455	364.7861	169.7961
A_24_P94	NM_01461	C9orf10 pr	C9orf10,DN	-2.20016	2.55E-30	0.45451	7127.16	3248.137
A_32_P71	NM_01526	KIAA0056	hCAP-D3,K	-2.20031	7.44E-17	0.45448	2435.531	1093.125
A_23_P77	NM_00281	Homo sapi	PSMD12,p	-2.20065	0	0.45441	10952.51	4980.938
A_23_P31	NM_01547	DKFZP564	DKFZP564	-2.20078	2.70E-24	0.45438	2081.466	949.9038
A_23_P57	D86972	TatD DNas	TATDN2	-2.20199	7.40E-16	0.45414	2128.122	958.2744
A_23_P61	NM_00265	Homo sapi	PLAGL2	-2.20238	3.40E-12	0.45405	1128.735	502.7699
A_23_P90	NM_01471	Homo sapi	LAPTM4A,	-2.20668	5.74E-21	0.45317	21964.73	9839.236
A_23_P14	NM_13364	Homo sapi	MTO1,CGI	-2.2075	4.72E-14	0.453	1006.627	461.0289
A_24_P31	NM_00688	calmodulin	CALM1,CA	-2.20757	1.30E-16	0.45299	1166.31	532.5974
A_23_P20	NM_01333	hypoxia-inc	HIG2	-2.20759	1.18E-38	0.45298	4199.478	1901.156
A_23_P15	NM_00332	Homo sapi	HIRA,TUP1	-2.20877	2.63E-09	0.45274	2675.743	1227.615
A_23_P25	NM_00557	Homo sapi	LMNB1,LM	-2.21002	2.56E-25	0.45248	2887.477	1307.769
A_23_P30	NM_03336	v-Ki-ras2 K	KRAS,KRA	-2.21035	0.00063	0.45242	319.2347	147.0997
A_32_P12	BI013473	Transcribe	BI013473	-2.21098	1.03E-34	0.45229	1880.05	850.5543
A_23_P21	NM_00504	Homo sapi	PRKX,PKX	-2.21132	5.29E-09	0.45222	655.8971	302.098
A_24_P38	NM_01229	Ras-GTPa	G3BP2	-2.21238	9.70E-27	0.452	10662.39	4827.217
A_32_P202913		Unknown	A_32_P202	-2.2146	1.72E-08	0.45155	3090.595	1382.248
A_24_P41	NM_15368	hypothetica	FLJ38973	-2.21523	8.47E-12	0.45142	1134.825	501.5034
A_23_P38	NM_01647	Homo sapi	C14orf100,	-2.21616	0	0.45123	5734.966	2587.938
A_23_P37	AK128205	DKFZP434	DKFZP434	-2.21729	1.52E-42	0.451	2291.641	1033.446
A_23_P40	NM_02121	Homo sapi	PCTP,STA	-2.21856	6.89E-16	0.45074	2683.387	1204.716
A_23_P33	NM_15236	Homo sapi	ZNF573,FL	-2.22191	6.28E-13	0.45006	1293.52	570.6613
A_24_P11	NM_00276	phosphorib	PRPS2,PR	-2.22313	5.68E-23	0.44982	2307.829	1033.84
A_24_P50	AK022936	CDNA FLJ	AK022936	-2.22338	2.13E-11	0.44977	1441.176	631.4843
A_23_P25	NM_14498	hypothetica	ALG14,MG	-2.22673	3.34E-27	0.44909	1458.394	655.8988
A_24_P713729		ALU8_HUM	THC19913	-2.22684	1.84E-20	0.44907	1955.177	872.4237
A_23_P50	NM_01231	Homo sapi	SLC39A6,L	-2.22727	3.14E-27	0.44898	1398.473	627.0228
A_23_P36	NM_03210	Splicing fa	SRP46	-2.22834	1.09E-06	0.44876	710.0281	300.7111
A_24_P30	AK024670	GATA zinc	p66alpha	-2.22866	9.97E-16	0.4487	982.9198	440.1688
A_23_P63	NM_01629	Homo sapi	HSPA14,H	-2.23121	5.83E-43	0.44819	6058.936	2720.846
A_32_P13	NM_01774	hypothetica	ZCCHC2,F	-2.23276	3.47E-34	0.44788	1850.328	829.4728
A_24_P33	NM_00710	ubiquitin-lik	UBL3,HCG	-2.23357	1.38E-08	0.44771	1474.765	637.7006
A_23_P14	NM_01818	hypothetica	FLJ10700	-2.23656	1.79E-14	0.44712	1544.78	677.1608
A_24_P77	AW135367	Pre-B-cell l	PBX1	-2.23789	0.0008	0.44685	533.4606	275.9674
A_24_P23	NM_00141	eukaryotic	EIF1AX,EIF	-2.23813	2.80E-45	0.4468	4060.439	1814.718
A_23_P32	NM_00027	Homo sapi	OCRL,LOC	-2.23835	4.26E-30	0.44676	3676.359	1641.605
A_24_P70	AL832246	Heterogen	HNRPA3	-2.23949	7.01E-36	0.44653	1924.726	859.0169
A_32_P80	AK021715	DEAD (Asp	DDX6	-2.24101	6.04E-37	0.44623	2168.89	967.5115
A_24_P50	BC031860	LOC12449	LOC12449	-2.24167	6.22E-17	0.4461	2047.797	904.1301
A_32_P94	BC068588	CDNA clon	BC068588	-2.24473	0.00006	0.44549	368.1737	164.8937
A_23_P31	NM_19888	Homo sapi	NUP43,p42	-2.24542	3.85E-33	0.44535	3495.353	1554.594
A_23_P15	NM_02520	hypothetica	FLJ21945,	-2.24701	4.85E-07	0.44504	504.1009	223.3083
A_32_P51	AK095525	Heterogen	HNRPU	-2.24729	1.50E-24	0.44498	1407.818	624.4968
A_24_P52	AK091744	CDNA FLJ	AK091744	-2.24853	9.96E-13	0.44474	825.1461	365.3355
A_24_P66	BC008245	Hypothetic	LOC40002	-2.2486	0.00005	0.44472	393.5746	179.107
A_24_P93	BC038505	BCL2-asso	BAG4	-2.25089	4.25E-09	0.44427	596.9694	264.1541
A_23_P41	NM_00294	Homo sapi	ABCE1,RL	-2.25456	6.56E-11	0.44355	2008.589	903.9285
A_23_P13	NM_00717	Homo sapi	STRAP,MA	-2.25524	6.07E-25	0.44341	45501.04	20291.07

A_23_P42	BC018658	Chromosor	C10orf74	-2.25528	1.49E-25	0.4434	1696.358	754.1646
A_24_P26	NM_00553	inositol pol	INPP5A,5F	-2.25867	2.00E-06	0.44274	564.3179	262.3605
A_23_P20	NM_00002	Homo sapi	AGL,GDE	-2.25884	4.37E-11	0.4427	858.8366	372.8292
A_32_P22	AB007976	KIAA0507	KIAA0507	-2.25926	9.80E-09	0.44262	624.7609	273.1266
A_23_P83	NM_02111	Homo sapi	RECK,ST1	-2.2651	0.00003	0.44148	490.8277	204.4668
A_23_P25	NM_00560	Homo sapi	LGMN,AEF	-2.26541	0.00016	0.44142	858.4068	436.5809
A_23_P10	NM_17070	Homo sapi	ICMT,PCM	-2.26681	6.22E-33	0.44115	2650.18	1167.231
A_23_P37	NM_03342	Homo sapi	KIAA1737	-2.26841	1.20E-22	0.44084	1867.364	818.4911
A_32_P12	NM_01508	Homo sapi	SPG20,SP	-2.27013	0.00099	0.4405	365.5977	177.3149
A_24_P32	BC015689	Collagen, t	COL9A3	-2.27185	6.04E-17	0.44017	1910.908	832.1786
A_23_P10	NM_00453	Homo sapi	NARS,ASN	-2.27258	8.97E-23	0.44003	9912.211	4361.472
A_23_P34	NM_01492	MAK-relate	ICK,MRK,L	-2.27286	1.51E-15	0.43997	1356.654	588.2722
A_23_P11	NM_01536	Homo sapi	SKIV2L2,K	-2.273	0	0.43995	7618.907	3351.408
A_24_P930597		Unknown	A_24_P93	-2.27619	3.34E-31	0.43933	14469.39	6381.054
A_32_P38	BC015133	CDNA clon	BC015133	-2.27684	1.77E-06	0.4392	476.8987	214.339
A_24_P91	BX108121	Transcribe	BX108121	-2.27948	0.00006	0.4387	557.1612	268.4768
A_24_P94	AL833922	PAP assoc	PAPD5	-2.2805	1.14E-08	0.4385	548.8527	240.4773
A_23_P39	NM_01335	CCR4-NOT	CNOT7,CA	-2.28233	1.55E-42	0.43815	3383.559	1482.116
A_23_P40	NM_01239	Homo sapi	PFAS,PUR	-2.284	8.84E-32	0.43783	5172.685	2268.798
A_32_P105281		Unknown	A_32_P10	-2.28406	1.41E-19	0.43782	1498.201	651.3148
A_32_P12	NM_00281	proteasom	PSMD12,p	-2.28526	1.17E-37	0.43759	8855.971	3875.089
A_23_P50	NM_00410	Homo sapi	EFEMP1,D	-2.28614	6.76E-13	0.43742	1181.156	505.7193
A_23_P25	NM_01595	2-deoxyrib	DERA	-2.28622	4.67E-20	0.4374	9244.027	4048.544
A_32_P23	NM_19826	Homo sapi	HIPK1,Mya	-2.28667	9.14E-17	0.43732	2167.865	941.0209
A_23_P30	NM_03336	v-Ki-ras2 K	KRAS,KRA	-2.28698	1.35E-14	0.43726	1588.293	684.3687
A_23_P25	NM_03213	Homo sapi	KBTBD7,D	-2.28852	0.00044	0.43696	402.1628	161.7413
A_23_P11	NM_01220	Homo sapi	HNRPH3,2	-2.29018	0	0.43665	4297.054	1875.198
A_23_P33	NM_00364	Homo sapi	HAT1	-2.29108	7.37E-24	0.43647	10796.86	4744.733
A_24_P27	NM_01653	apoptosis-r	FLJ39616	-2.293	7.64E-11	0.43611	1548.781	689.3062
A_32_P23	BC026302	Membrane	M11S1	-2.29738	0	0.43528	13708.39	5964.893
A_23_P11	AK129920	Son of sev	SOS2	-2.29874	0.00038	0.43502	428.4083	169.4541
A_24_P92	AK058024	Zinc finger	ZNF161	-2.29879	4.21E-28	0.43501	1634.999	713.0548
A_23_P16	NM_01844	PC326 pro	IQWD1,NR	-2.30096	1.31E-11	0.4346	818.0916	350.3019
A_32_P15	AK022268	Transducin	TBL1XR1	-2.30136	1.40E-10	0.43453	584.6604	254.6847
A_24_P94	AK091153	AHNAK nu	MGC5395	-2.30283	0	0.43425	5328.863	2314.415
A_23_P33	NM_18337	Homo sapi	ARRDC4,F	-2.30393	0.00004	0.43404	401.2842	180.5777
A_24_P15	NM_01810	hypothetica	C14orf130,	-2.30424	2.03E-33	0.43398	1917.708	830.6442
A_23_P53	NM_03362	Homo sapi	FBXO21,F	-2.30458	0	0.43392	7043.401	3056.46
A_24_P38	NM_00284	protein tyr	PTPRF,LA	-2.30682	3.68E-42	0.4335	6241.588	2709.749
A_23_P25	NM_01804	hypothetica	AGGF1,VG	-2.31722	5.65E-18	0.43155	1162.437	504.8729
A_23_P20	NM_03234	Homo sapi	C9orf125,M	-2.31765	5.47E-24	0.43147	5249.763	2271.622
A_24_P94	NM_18098	intimal thic	ITR	-2.31818	4.91E-10	0.43137	619.297	269.9969
A_23_P43	BC029662	Homo sapi	C20orf142,	-2.31824	7.32E-21	0.43136	1307.855	567.5596
A_23_P32	BC064969	Acidic (leu	ANP32A	-2.31861	0	0.43129	4180.804	1804.161
A_24_P88	AK055625	Polycomb	PCGF5	-2.32005	6.36E-22	0.43102	1964.088	839.3385
A_23_P25	NM_03096	Homo sapi	RNF146,d	-2.32103	0.00001	0.43084	462.915	210.0438
A_23_P12	NM_01823	Homo sapi	NUP133,hI	-2.32265	1.85E-23	0.43054	2944.751	1268.025
A_23_P50	NM_00704	E1B-55kDa	HNRPUL1,	-2.32321	1.07E-21	0.43044	4269.782	1839.191
A_24_P13	XM_29302	similar to U	LOC34438	-2.3252	6.87E-30	0.43007	2739.67	1176.575
A_23_P35	NM_15243	Homo sapi	ZNF664,ZF	-2.32703	1.69E-29	0.42973	6409.908	2766.531
A_24_P24	NM_00462	translin	TSN,TRSL	-2.33026	0	0.42914	3612.492	1548.846
A_24_P64	AK124953	Similar to h	AK124953	-2.33112	0.00011	0.42898	359.6511	158.195

A_23_P47	NM_02456	Homo sapi	MPZL1,PZ	-2.33295	1.25E-29	0.42864	5061.008	2165.489
A_23_P37	NM_13344	Homo sapi	GPT2,ALT	-2.33309	2.74E-13	0.42862	1130.288	494.4052
A_23_P12	NM_01602	Anterior ph	APH-1A	-2.33374	2.05E-07	0.4285	1303.284	583.5755
A_32_P10	AK130366	Similar to h	AK130366	-2.33642	0.00018	0.428	338.1025	142.7777
A_23_P73	NM_00031	protein S (a	PROS1,PS	-2.33726	0.00004	0.42785	425.6691	192.096
A_24_P74	NM_00340	zinc finger	ZFP161,ZF	-2.33872	3.72E-10	0.42758	657.9575	278.7512
A_24_P35	NM_13083	Homo sapi	OPA1,NPG	-2.34063	2.93E-16	0.42723	1223.791	514.4332
A_32_P12	BC037740	CDNA clon	BC037740	-2.34309	2.10E-24	0.42679	4968.034	2104.14
A_23_P15	NM_00240	Homo sapi	MEST,PEG	-2.34318	2.45E-09	0.42677	925.8558	378.8561
A_24_P76	AB007954	KIAA0485	KIAA0485	-2.34464	1.72E-09	0.42651	960.682	426.8094
A_32_P111266		PEBB_HUI	THC19625	-2.34574	0.00001	0.4263	397.6939	170.7745
A_24_P922858		ALU5_HUM	THC18195	-2.34674	0.0001	0.42612	363.6838	159.8693
A_24_P38	NM_00350	frizzled (Dr	FZD1	-2.34742	3.34E-16	0.426	1109.159	466.0928
A_32_P35	AA482505	Solute carr	SLC7A5	-2.34761	0.00015	0.42596	838.6826	400.796
A_24_P40	AF131831	Full length	CDC42SE2	-2.34881	0	0.42575	3820.087	1626.457
A_23_P12	NM_01234	Homo sapi	GTPBP4,N	-2.34905	3.80E-28	0.4257	13346.09	5721.405
A_24_P37	NM_02191	cofilin 2 (m	CFL2	-2.35244	7.00E-24	0.42509	2544.709	1079.2
A_32_P105397		AB030032	THC19913	-2.35482	7.92E-07	0.42466	526.5125	216.3068
A_23_P23	NM_00678	Homo sapi	WDR3	-2.35615	3.01E-22	0.42442	6265.733	2667.963
A_24_P36	NM_00500	NADH deh	NDUFA9,N	-2.35634	3.85E-21	0.42439	4146.512	1740.667
A_23_P25	NM_00588	Homo sapi	SLC25A3,F	-2.35798	2.55E-42	0.42409	116926	49681.96
A_24_P26	AL833350	THAP dom	THAP6	-2.35876	1.02E-10	0.42395	614.4422	260.0406
A_24_P13	NM_19840	Homo sapi	PTPLB	-2.36125	2.75E-24	0.4235	2989.927	1265.882
A_23_P54	NM_01404	Homo sapi	MKL2,MRT	-2.36146	1.46E-18	0.42347	1813.293	760.6171
A_23_P74	NM_01806	Homo sapi	IARS2,FLJ	-2.36471	0	0.42289	15480.22	6548.928
A_24_P90	AF321617	Homo sapi	unknown	-2.36472	2.80E-30	0.42288	2337.48	985.3966
A_24_P34	NM_19429	spindle ass	SASS6,SA	-2.3663	8.19E-07	0.4226	461.0266	193.0017
A_24_P706752		ALU1_HUM	THC18774	-2.36684	3.59E-10	0.4225	649.9331	277.0635
A_23_P23	NM_00368	Homo sapi	EXO1,HEX	-2.36774	5.50E-32	0.42234	4752.234	2019.762
A_23_P35	AB046773	KIAA1553	KIAA1553	-2.36938	1.34E-37	0.42205	2123.259	896.825
A_32_P10	NM_14526	similar to C	LOC13739	-2.37194	1.10E-13	0.4216	1048.599	434.212
A_24_P12	NM_00678	WD repeat	WDR3	-2.37292	2.75E-22	0.42142	3783.342	1592.274
A_24_P24	NM_01628	adenylate k	AK3,AK6,F	-2.37863	0.00018	0.42041	358.1019	158.0348
A_23_P58	AY033606	Golgi asso	GOPC	-2.38118	0	0.41996	2889.136	1213.16
A_24_P37	NM_00163	S-adenosy	AMD1,AMI	-2.38134	0	0.41993	18744.83	7884.004
A_24_P31	NM_00677	Kaiso	ZBTB33,ZN	-2.3835	3.63E-14	0.41955	1188.731	488.6392
A_23_P16	NM_02208	Homo sapi	TXNDC5,E	-2.38449	0	0.41938	6955.352	2918.15
A_32_P183022		ALU1_HUM	THC18818	-2.38484	8.76E-08	0.41932	494.6105	206.6359
A_24_P13	AL831922	DKFZP586	DLG1	-2.39159	1.10E-16	0.41813	1742.89	719.2476
A_32_P20	NM_00210	H3 histone	H3F3A,H3I	-2.40296	2.80E-18	0.41615	88830.7	37218.31
A_24_P30	NM_00602	D123 gene	C10orf7,D1	-2.405	4.95E-35	0.4158	5972.879	2488.413
A_24_P14	NM_18341	Homo sapi	KIF1B,KLP	-2.40774	0.00006	0.41533	345.8098	144.0985
A_23_P10	AB020637	KIAA0830	KIAA0830	-2.40816	2.13E-22	0.41526	1511.943	631.9548
A_32_P62	NM_01849	PDZ-bindin	PBK,SPK,T	-2.41771	1.02E-39	0.41361	7575.827	3140.384
A_23_P21	NM_00638	DEAD/H (A	DDX17,P7	-2.41967	4.68E-06	0.41328	542.056	241.6613
A_32_P95	BX538303	Phosphorib	PAICS	-2.41984	0.0009	0.41325	1131.429	523.025
A_23_P20	NM_01901	Homo sapi	POLR1B,R	-2.4212	4.42E-27	0.41302	1571.069	652.3489
A_24_P15	NM_02074	Homo sapi	ANKFY1,A	-2.42337	4.54E-10	0.41265	618.7357	258.668
A_32_P8806		ALU4_HUM	THC19993	-2.42342	5.16E-26	0.41264	1382.486	573.7868
A_24_P18	NM_17370	Homo sapi	NM_17370	-2.42453	0	0.41245	6152.067	2536.624
A_24_P671842		PPOL_HUI	THC19431	-2.42792	0	0.41188	5784.796	2381.99
A_24_P27	NM_01682	v-crk avian	CRK,CRKI	-2.42799	0.00034	0.41186	850.3626	403.1872

A_23_P99	NM_01653	apoptosis-r	FLJ39616	-2.4287	2.33E-14	0.41174	1065.493	431.5071
A_23_P39	NM_02113	Homo sapi	RNASEL,H	-2.43322	7.74E-10	0.41098	804.4534	318.5627
A_24_P93	AB011092	Adenylate	ADCY9	-2.43521	4.36E-13	0.41064	936.6199	376.5368
A_23_P36	NM_00457	RAB4A, m	RAB4A	-2.44812	3.68E-42	0.40848	2553.082	1043.111
A_23_P25	NM_05817	Homo sapi	PSAT1,PS	-2.45419	1.69E-42	0.40747	20798.3	8498.723
A_23_P34	NM_17346	Homo sapi	MOBKL1A	-2.45883	7.91E-09	0.4067	712.3594	278.0877
A_24_P245108		Unknown	THC19435	-2.45919	5.06E-06	0.40664	532.993	232.7046
A_23_P69	NM_00135	Homo sapi	DHX15,DB	-2.4605	0	0.40642	6023.252	2449.812
A_32_P179005		ALU8_HUM	THC18442	-2.46884	1.34E-08	0.40505	551.1801	227.9925
A_24_P39	NM_00429	guanine de	GDA,CYPI	-2.46929	6.47E-15	0.40498	1250.044	495.6631
A_24_P59	NM_00210	H3 histone	H3F3A,H3I	-2.47108	0	0.40468	50946.24	20638.64
A_24_P20	NM_15238	Homo sapi	BBS5	-2.47117	0.00001	0.40467	385.749	154.0606
A_24_P592060		ALU6_HUM	THC19531	-2.47154	3.06E-07	0.40461	470.1422	188.2665
A_23_P63	NM_01329	Homo sapi	GPSM2,LG	-2.47187	0.00001	0.40455	509.5558	222.0413
A_24_P40	NM_02483	hypothetica	FLJ12443,	-2.47565	3.46E-08	0.40393	710.562	302.8785
A_23_P20	NM_00062	Homo sapi	SERPINA5	-2.47586	6.50E-06	0.4039	558.442	273.94
A_23_P61674		AF212224	THC19644	-2.48146	0.00002	0.40299	477.2497	179.5256
A_23_P32	NM_17382	mitochondr	FLJ38991	-2.48378	2.37E-26	0.40261	1245.501	501.6666
A_32_P18	AY007110	Acidic (leu	ANP32A	-2.49044	1.71E-41	0.40154	4241.358	1703.594
A_24_P87	AK022110	Similar to F	LOC49386	-2.49695	1.20E-20	0.40049	1178.726	478.0975
A_32_P10	NM_00294	ATP-bindin	ABCE1,RL	-2.50593	3.67E-29	0.39905	2625.891	1052.207
A_32_P73	BX119435	Transcribe	BX119435	-2.50909	0.00009	0.39855	359.0241	137.9915
A_23_P45	NM_02236	Homo sapi	TFB2M,Hk	-2.51959	1.92E-30	0.39689	10476.7	4172.372
A_23_P31	NM_01329	transforme	TRA2A,HS	-2.52072	2.41E-16	0.39671	875.9865	348.553
A_24_P22	NM_02091	Homo sapi	GPAM,KIA	-2.52689	5.92E-16	0.39574	2104.665	822.3821
A_32_P16	AK056476	Trinucleoti	TNRC8	-2.52726	2.80E-18	0.39569	1373.566	550.9534
A_24_P93	NM_00520	v-crk avian	CRKL	-2.53051	4.51E-06	0.39518	463.7701	174.8448
A_24_P925664		ALU1_HUM	THC18071	-2.53275	1.39E-09	0.39483	546.0513	215.2856
A_32_P82179		Unknown	THC18098	-2.53617	2.05E-30	0.3943	1677.66	665.2471
A_23_P43	NM_01560	Homo sapi	C10orf137,	-2.53732	9.63E-28	0.39412	1643.947	643.2756
A_24_P25	NM_00359	cullin 2	CUL2	-2.53852	4.52E-10	0.39393	625.4963	242.4329
A_32_P99	BM709498	Transcribe	BM709498	-2.53982	1.37E-09	0.39373	668.4417	254.803
A_24_P25	AB051504	SET domai	SET7	-2.54016	2.42E-10	0.39368	932.025	350.9678
A_24_P78	U25029	Nuclear re	NR3C1	-2.54269	3.05E-08	0.39328	509.8519	197.8331
A_24_P19	NM_03321	tripartite m	TRIM14,KI	-2.54466	1.63E-13	0.39298	1176.697	448.7946
A_23_P13	NM_01984	Homo sapi	EIF4ENIF1	-2.54604	1.84E-25	0.39277	1224.156	482.3286
A_24_P16	NM_00125	cyclin-depe	CDK6,PLS	-2.54959	8.55E-32	0.39222	3366.729	1318.827
A_24_P40481		Unknown	THC19923	-2.54967	9.04E-09	0.39221	546.8871	217.8698
A_32_P33213		Unknown	THC18645	-2.55039	9.54E-16	0.3921	771.218	301.1112
A_24_P17	AK027667	Homo sapi	FLJ14761	-2.5515	5.41E-22	0.39193	1515.176	586.7499
A_23_P87	BC004815	Hypothetic	MGC5139	-2.55754	1.01E-32	0.391	2361.438	927.3075
A_24_P204334		Unknown	A_24_P20	-2.56519	9.69E-21	0.38983	14805.42	5788.453
A_24_P80571		Unknown	A_24_P80	-2.566	9.09E-11	0.38971	589.0399	230.9453
A_24_P21	NM_00480	transmemb	TM9SF2,P	-2.56673	0	0.3896	4456.087	1735.027
A_24_P28	NM_00238	membrane	MCP,TLX,	-2.56907	7.48E-17	0.38925	1722.352	656.8876
A_32_P121226		Unknown	A_32_P12	-2.58463	6.01E-07	0.3869	605.7581	277.3202
A_24_P94	AB007896	Prolyl endo	KIAA0436	-2.58486	0	0.38687	3029.263	1169.948
A_23_P16	AF090927	Homo sapi	PRO0457	-2.58648	8.75E-06	0.38663	609.8297	261.5685
A_32_P127209		ALU4_HUM	THC18351	-2.59987	3.71E-22	0.38464	1722.056	655.3817
A_24_P926450		ALU8_HUM	THC18475	-2.60993	4.65E-13	0.38315	1098.034	408.6063
A_24_P94	AB037811	Family with	FLJ11280	-2.61354	0.00002	0.38262	355.9675	137.8123
A_23_P34	AB011166	SMC5 stru	SMC5L1	-2.62368	1.10E-30	0.38114	2145.528	812.7488

A_23_P29	NM_00588	Homo sapi	MAEA,EMI	-2.62575	0	0.38084	13563.6	5182.335
A_32_P112401		Unknown	THC19528	-2.62845	0.00051	0.38045	299.1385	120.1502
A_24_P20	AY033611	Homo sapi	AY033611	-2.63886	1.73E-30	0.37895	79976.44	30436.75
A_32_P69	AK023933	BTB (POZ)	HSPC063	-2.64148	1.13E-23	0.37858	1087.901	410.463
A_32_P70	NM_02513	hypothetica	C13orf23,F	-2.64231	1.30E-22	0.37846	2142.782	801.0812
A_23_P25	NM_01641	Homo sapi	NF2,ACN,S	-2.6427	1.79E-26	0.3784	6169.737	2340.933
A_23_P20	NM_00141	ELAV (emb	ELAVL1,HU	-2.64277	2.58E-32	0.37839	2967.524	1118.203
A_32_P48526		Unknown	THC19038	-2.64561	5.92E-06	0.37798	424.3151	168.9397
A_24_P72	BX538231	BTB (POZ)	BTBD7	-2.64866	3.76E-20	0.37755	1471.995	547.221
A_32_P50655		ALU5_HUM	THC18078	-2.6566	3.94E-18	0.37642	900.6237	337.544
A_24_P20	NM_01503	Formin bin	FNBP1	-2.65682	6.97E-08	0.37639	837.5107	291.4368
A_23_P10	NM_02053	Homo sapi	C20orf3,BS	-2.6587	0	0.37612	19147.65	7200.759
A_23_P40	NM_17370	Homo sapi	NM_17370	-2.65915	0	0.37606	31134.35	11688.7
A_23_P33	NM_17370	Homo sapi	NM_17370	-2.66659	2.92E-28	0.37501	26814.09	10131.71
A_23_P37	NM_17497	Homo sapi	CCNK	-2.66683	1.74E-27	0.37498	1535.517	572.4511
A_23_P35	NM_03290	hypothetica	RBM17,SP	-2.66692	0.00056	0.37496	311.2765	126.9391
A_23_P25	NM_00248	Homo sapi	NCBP1,NC	-2.66866	1.09E-40	0.37472	3583.597	1341.954
A_32_P171921		AB017004	THC18654	-2.66971	1.15E-06	0.37457	933.6983	394.8911
A_32_P10	AW135423	UI-H-BI1-a	AW135423	-2.67643	0.00014	0.37363	318.5832	115.9774
A_24_P552987		Unknown	A_24_P55	-2.67997	3.66E-23	0.37314	1223.883	461.1751
A_24_P46	AK056550	Syncoilin, i	SYNCOILII	-2.68378	9.18E-13	0.37261	980.8677	378.306
A_32_P157481		ALU7_HUM	THC18916	-2.68573	4.13E-08	0.37234	593.413	210.269
A_23_P32	NM_19907	Homo sapi	MDFIC,HIC	-2.69444	8.40E-09	0.37114	1147.719	447.7347
A_24_P55	M28016	Human mit	cytochrome	-2.69516	5.09E-43	0.37104	39587.9	14737.36
A_23_P20	NM_00229	Homo sapi	LBR,PHA,L	-2.69523	2.00E-23	0.37103	31068.73	11617.92
A_23_P30	AK096403	Epithelial m	EMP2	-2.70996	1.77E-19	0.36901	1246.644	467.5732
A_23_P12	NM_00225	potassium	KCNS3,KV	-2.71249	0.00018	0.36867	413.3311	135.0296
A_23_P11	NM_00106	Homo sapi	TOP2A,TO	-2.71875	0	0.36782	3535.078	1299.366
A_24_P25	NM_00640	vitamin A r	ARL6IP5,J	-2.72186	6.28E-35	0.3674	7893.284	2898.104
A_32_P13991		Unknown	THC18823	-2.72363	1.13E-10	0.36716	639.3283	229.3373
A_24_P24	NM_00648	dual-specif	DYRK2	-2.7365	7.43E-08	0.36543	485.2574	181.8698
A_23_P40	NM_00434	carbamoyl-	CAD	-2.73729	1.61E-35	0.36533	1823	663.2151
A_24_P20	AK074041	DnaJ (Hsp	DNAJC5	-2.74428	2.38E-14	0.36439	1167.569	438.2711
A_32_P133038		Unknown	THC18645	-2.76869	8.14E-10	0.36118	601.832	224.9117
A_23_P115902		ALU2_HUM	THC19761	-2.76893	6.70E-13	0.36115	793.6057	279.3271
A_32_P31	NM_00555	Homo sapi	LAMA1,LA	-2.77583	1.09E-27	0.36025	1371.623	491.6881
A_23_P14	NM_00093	Homo sapi	POLR2B,R	-2.78295	3.01E-08	0.35933	3774.021	1371.555
A_24_P29	NM_01221	tripartite m	TRIM32,HT	-2.8041	0	0.35662	1925.774	686.9573
A_23_P12	NM_03273	Homo sapi	RTN4IP1,N	-2.81195	0	0.35562	3451.425	1226.602
A_24_P902728		BC001731	THC18350	-2.81403	2.55E-08	0.35536	498.1659	182.1618
A_23_P13	NM_00466	Homo sapi	CDC23,AP	-2.8151	6.90E-36	0.35523	2382.693	841.3092
A_24_P93	AK123264	CDNA FLJ	AK123264	-2.81851	1.50E-09	0.3548	694.7312	234.2281
A_23_P30	NM_01532	Homo sapi	KIAA0776,	-2.82616	1.12E-31	0.35384	1834.506	645.971
A_32_P23	NM_01208	ATP-bindin	ABCB10,M	-2.83348	1.37E-10	0.35292	595.7543	215.7499
A_23_P15	M37712	Homo sapi	p58/GTA p	-2.83688	1.31E-25	0.3525	3683.993	1300.347
A_23_P78	BC000892	Chromosor	C18orf55	-2.8473	0	0.35121	7805.876	2738.986
A_23_P87	NM_01808	hypothetica	POLR3B,R	-2.84756	1.89E-28	0.35118	1895.705	658.8768
A_23_P33	BC036414	PAI-1 mRN	PAI-RBP1	-2.84767	1.77E-08	0.35116	554.3749	204.5202
A_23_P60	NM_03230	Homo sapi	C9orf64,M	-2.85244	0.00002	0.35058	1186.496	469.4175
A_23_P16	NM_00640	Homo sapi	ARL6IP5,J	-2.86418	0	0.34914	5930.394	2068.303
A_24_P30	AK021848	Homo sapi	FLJ11786	-2.86511	4.03E-12	0.34903	614.6672	216.7746
A_32_P18	NM_02162	guanine nu	GNB4	-2.8652	8.72E-13	0.34902	779.5585	265.3714

A_24_P79	XR_00018	Homo sapi	XR_00018	-2.8843	3.84E-13	0.3467	708.2279	240.0552
A_32_P20	BC058829	HLA compl	HCG12	-2.89111	1.72E-39	0.34589	2041.202	703.4596
A_24_P817223		AF194537	THC18585	-2.90266	5.08E-10	0.34451	582.4623	195.0796
A_24_P11	NM_02272	DiGeorge s	DGCR8,Gy	-2.90455	2.26E-16	0.34429	1068.159	377.2719
A_24_P63	AK022223	Hypothetic	FLJ22955	-2.92963	8.97E-09	0.34134	718.7213	262.9111
A_23_P10	NM_15271	Homo sapi	ITM1,TMC	-2.95633	0	0.33826	4584.864	1550.422
A_23_P43	NM_01457	Homo sapi	HEYL	-2.95999	9.39E-06	0.33784	729.1639	298.8361
A_32_P23	BX108897	BX108897	BX108897	-2.96197	0.00004	0.33761	1020.221	388.5218
A_23_P40	NM_17380	hypothetica	MGC5055	-2.97872	4.76E-06	0.33571	756.5419	314.9604
A_24_P26	AJ420430	Full length	AJ420430	-3.00459	3.59E-39	0.33282	2053.807	681.0625
A_23_P33	NM_15344	Homo sapi	GALGT2,C	-3.0253	6.22E-09	0.33055	659.152	259.9594
A_24_P92	BC020784	Similar to F	MGC1428	-3.02949	0	0.33009	3214.666	1061.095
A_32_P14	BC062473	Microsoma	MGST1	-3.03137	0	0.32988	5733.393	1892.882
A_23_P86	NM_00687	UDP-GlcN	B3GNT6,iC	-3.0433	5.37E-11	0.32859	880.4952	272.7964
A_24_P48	AF086017	Hypothetic	LOC28614	-3.05298	1.79E-06	0.32755	401.1266	138.2053
A_24_P93	AL133624	Ropporin, r	ROPN1	-3.0551	4.62E-08	0.32732	393.0709	142.4871
A_23_P149938		Unknown	THC19475	-3.0782	1.67E-11	0.32487	691.3126	235.0989
A_23_P20	NM_01208	Homo sapi	ABCB10,M	-3.07871	3.37E-35	0.32481	2542.437	820.744
A_23_P97	NM_00277	Homo sapi	HTRA1,L5	-3.08206	0	0.32446	8727.062	2839.269
A_32_P45	BX648283	KIAA1171	KIAA1171	-3.11609	2.26E-13	0.32092	853.2781	285.9731
A_24_P915079		Unknown	A_24_P91	-3.13167	4.85E-08	0.31932	748.1934	282.7574
A_24_P94	AB002331	Death assc	DATF1	-3.14027	3.10E-06	0.31844	416.0485	142.6951
A_23_P10	NM_02307	Chromosor	C16orf28	-3.14502	1.01E-39	0.31796	2747.706	872.0143
A_24_P59	BC013074	Mitochondr	MCART1	-3.1728	3.71E-09	0.31518	557.5934	185.86
A_23_P51	NM_01900	ETAA16 pr	ETAA16	-3.18574	2.70E-14	0.3139	787.7813	238.9482
A_24_P37	NM_02283	hypothetica	USP46,FL	-3.2198	0.00031	0.31058	255.6534	80.2039
A_24_P97785		Unknown	THC18911	-3.22432	7.52E-14	0.31014	591.2699	183.5171
A_23_P24	NM_02038	Homo sapi	XPNPEP1,	-3.26724	2.60E-08	0.30607	818.5407	287.1261
A_24_P59	AK094159	CDNA FLJ	AK094159	-3.29766	3.16E-16	0.30325	843.8577	248.6548
A_23_P40	D86973	GCN1 gen	GCN1L1	-3.30466	0	0.3026	18117.43	5492.113
A_23_P21	NM_01542	Gene mod	DKFZP564	-3.3072	0	0.30237	9320.514	2820.637
A_23_P33	NM_00087	Homo sapi	IGF2R,MP	-3.33146	1.54E-25	0.30017	2545.511	756.0175
A_24_P22	AF271776	Homo sapi	DC48	-3.35561	0	0.29801	7857.763	2334.04
A_23_P31	AB028999	KIAA1076	KIAA1076	-3.36786	0.00002	0.29692	344.3181	97.0251
A_24_P31	NM_19827	SET and M	SMYD1	-3.38034	2.34E-09	0.29583	665.9824	240.7633
A_32_P47	CA311129	Transcribe	CA311129	-3.41144	1.78E-09	0.29313	487.4576	169.0136
A_32_P22	BG436185	602508593	BG436185	-3.42299	0	0.29214	9861.699	2881.529
A_24_P32	NM_01766	hypothetica	TRPM6,HS	-3.44295	7.50E-09	0.29045	424.2977	155.6791
A_24_P93	AK092744	Homo sapi	AK092744	-3.44968	2.54E-09	0.28988	525.9974	183.2107
A_23_P21	NM_00503	Peroxisom	PPARA	-3.47566	1.66E-13	0.28771	731.2187	223.098
A_24_P11	AF289610	Clone pp9	unknown	-3.49294	0.00003	0.28629	506.6717	166.8974
A_23_P33	NM_17370	Homo sapi	NM_17370	-3.51508	3.69E-23	0.28449	154296.1	43986.61
A_24_P33	BC015876	Homo sapi	BC015876	-3.51623	5.56E-10	0.2844	439.4179	139.07
A_23_P56	NM_03241	esophagea	ECRG4	-3.53834	3.16E-11	0.28262	514.2326	159.6897
A_32_P34	AK026418	FLJ39739	FLJ39739	-3.54527	5.19E-33	0.28207	1189.624	339.2987
A_32_P15	NM_20737	FLJ45187	FLJ45187,	-3.56561	0.00003	0.28046	312.0877	93.37494
A_24_P852099		Unknown	A_24_P85	-3.5874	0	0.27875	106706.7	29763.41
A_24_P25040		Unknown	A_24_P25	-3.59648	9.65E-38	0.27805	15527.69	4248.493
A_23_P11	NM_00161	Homo sapi	PARP1,PA	-3.59892	0	0.27786	24320.75	6765.235
A_24_P34	NM_20737	Homo sapi	SERPINA1	-3.59915	8.04E-08	0.27784	536.1224	194.032
A_24_P861875		Unknown	THC19471	-3.61146	4.51E-11	0.2769	663.5716	218.7068
A_24_P92	NM_00020	I factor (co	IF,FI,factor	-3.65465	3.24E-10	0.27362	530.5601	168.0904

A_23_P13	NM_19913	hypothetica	LOC13414	-3.66628	5.58E-08	0.27276	526.6715	130.7824
A_32_P115277		ALU7_HUM	THC18362	-3.67922	3.97E-13	0.2718	634.2994	180.1054
A_23_P30	NM_17370	Homo sapi	NM_17370	-3.68265	0	0.27154	175628.2	47899.04
A_23_P34	NM_01741	frizzled (Dr	FZD3,Fz-3	-3.72647	0.00042	0.26835	249.446	64.23846
A_24_P16	NM_14890	Homo sapi	FOXP2,SP	-3.72696	2.29E-08	0.26832	838.2018	279.836
A_24_P55	AF391805	Homo sapi	unknown	-3.78021	0	0.26454	59217.42	15705.45
A_23_P49	NM_00692	Homo sapi	SFRS1,AS	-3.78684	0	0.26407	13367.58	3526.235
A_24_P94	AK024870	Dual-specif	DYRK2	-3.82359	1.57E-09	0.26153	473.8078	119.4877
A_23_P32	NM_02504	hypothetica	C15orf20,F	-3.86058	0	0.25903	1303.525	337.46
A_24_P534243		Unknown	A_24_P53	-3.86389	1.70E-13	0.25881	812.7372	241.5016
A_23_P10	NM_00294	Homo sapi	RPA1,HSS	-3.88268	0	0.25755	18380.55	4753.601
A_32_P75	BF063696	CDNA FLJ	BF063696	-4.01531	1.25E-12	0.24905	541.2093	149.517
A_23_P37	NM_02516	LAP (leucir	LRRC1,LA	-4.0447	1.24E-14	0.24724	840.6993	232.141
A_24_P88	AK127661	Hypothetic	MGC5509	-4.04974	1.39E-16	0.24693	922.7072	234.6905
A_23_P15	NM_00480	Homo sapi	TM9SF2,P	-4.09069	0	0.24446	7106.325	1739.552
A_32_P35	BC034591	Hypothetic	LOC28464	-4.18915	2.19E-11	0.23871	722.3524	217.38
A_32_P180958		ALU8_HUM	THC19992	-4.23196	1.27E-39	0.2363	3675.766	866.7258
A_24_P92	S81524	RC1=NADI	RC1	-4.26036	0	0.23472	11460.69	2677.543
A_24_P61	BC000206	Homo sapi	BC000206	-4.27122	0	0.23413	14762.32	3438.174
A_24_P92	AK096975	Protein tyr	PTPN3	-4.29004	3.51E-06	0.2331	348.4244	75.46396
A_24_P20	AK094466	TRNA splic	SEN54L	-4.31405	1.16E-12	0.2318	539.2351	121.9188
A_23_P11	NM_00281	proteasom	PSMD12,p	-4.34065	0	0.23038	1842.156	424.4357
A_23_P36	NM_17371	Homo sapi	NM_17371	-4.37448	0	0.2286	66876.19	15343.2
A_24_P74	NM_01417	HSPC154	C18orf55,H	-4.39389	0.00004	0.22759	340.9675	66.78474
A_24_P94	NM_01629	prenylcyste	PCYOX1,P	-4.90144	1.64E-26	0.20402	1188.241	236.2896
A_32_P12	BE877371	Clone pp9	BE877371	-5.67013	9.13E-07	0.17636	412.5563	86.35796
A_23_P31	NM_17371	Homo sapi	NM_17371	-6.13583	0	0.16298	6923.415	1124.564