

Genetic regulatory variation and environmental response

David L. Aylor

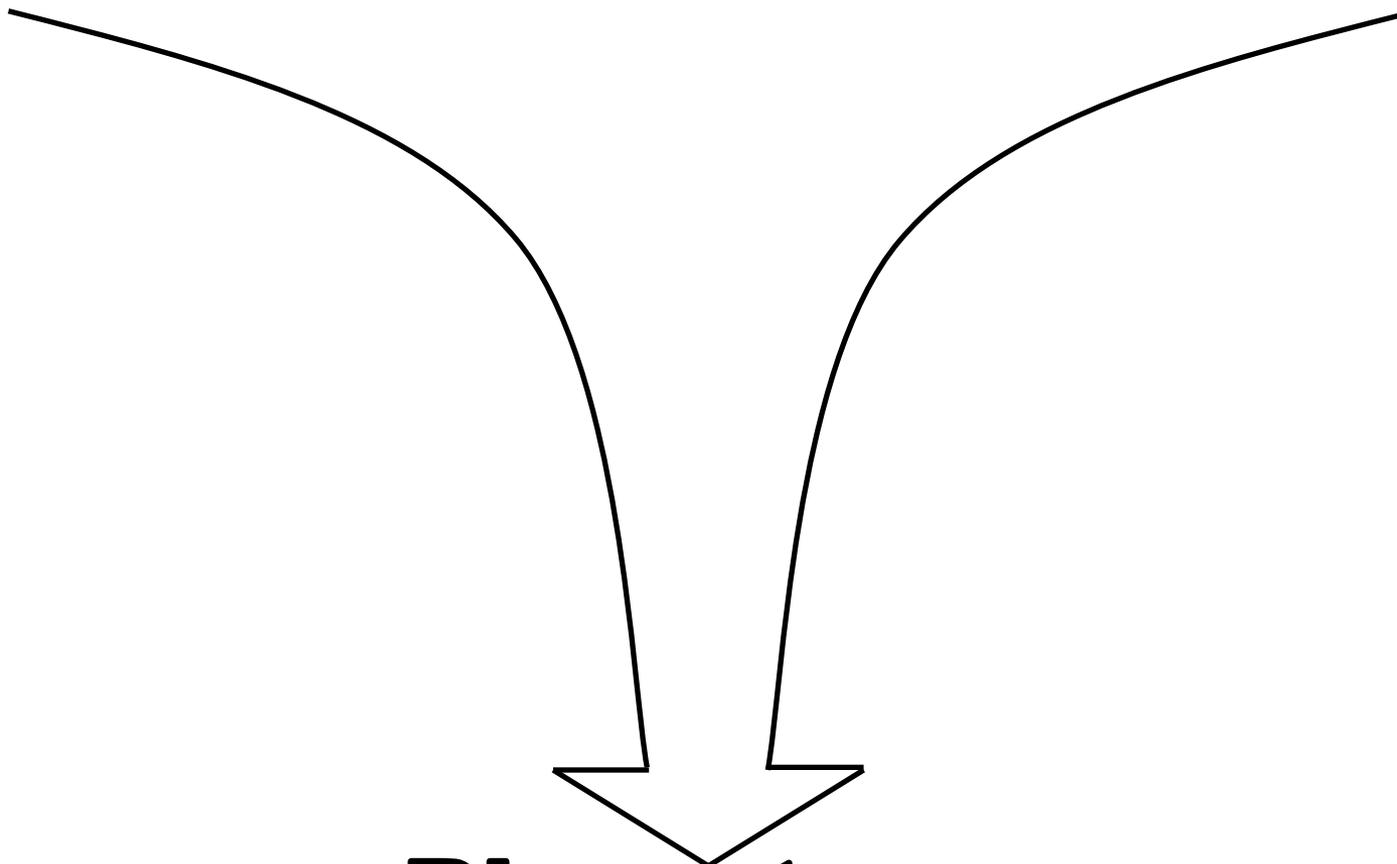
Population-Based Rodent Resources for
Environmental Health Sciences

NIEHS

March 19, 2015

Genotype

Environment



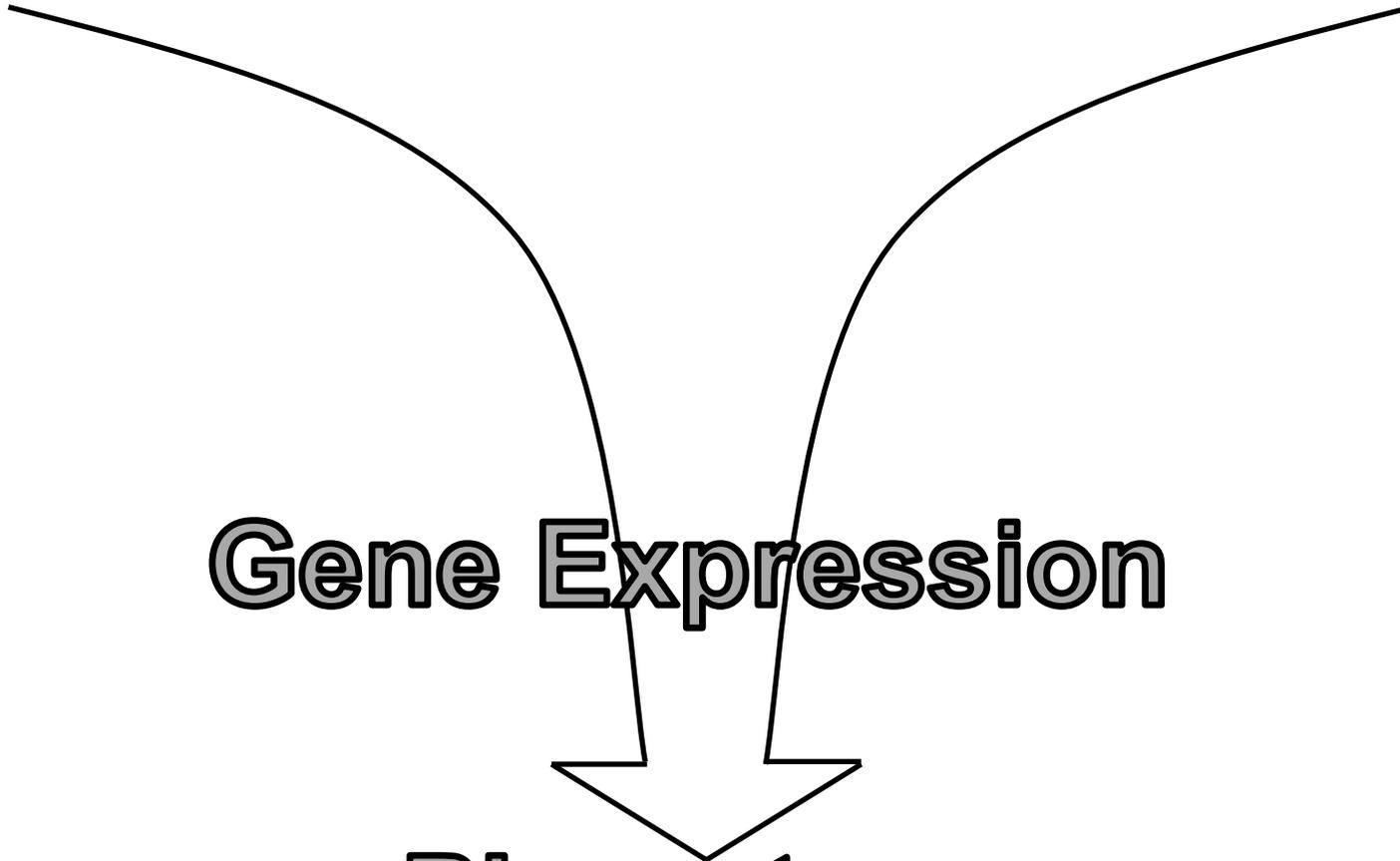
Phenotype

Genotype

Environment

Gene Expression

Phenotype



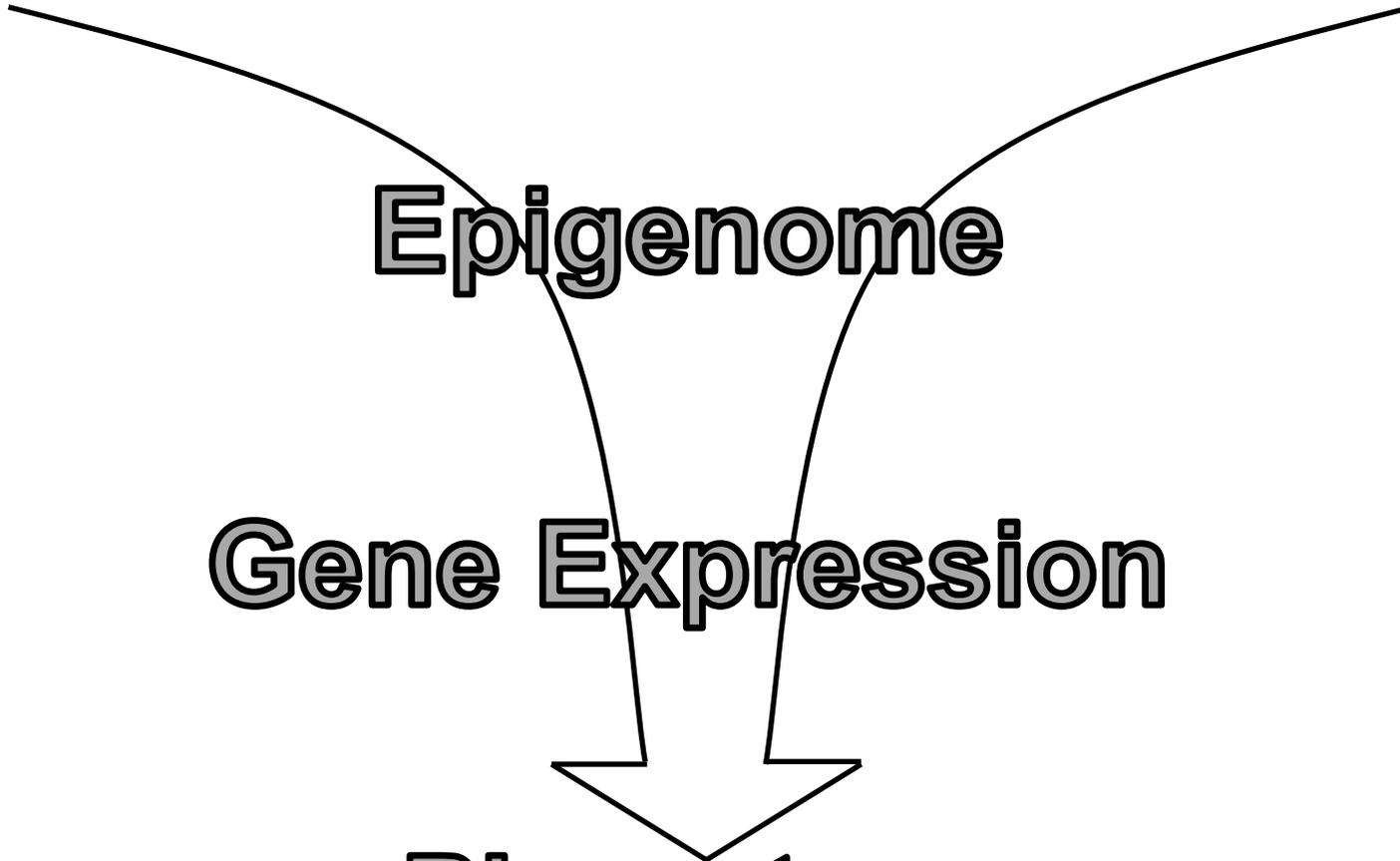
Genotype

Environment

Epigenome

Gene Expression

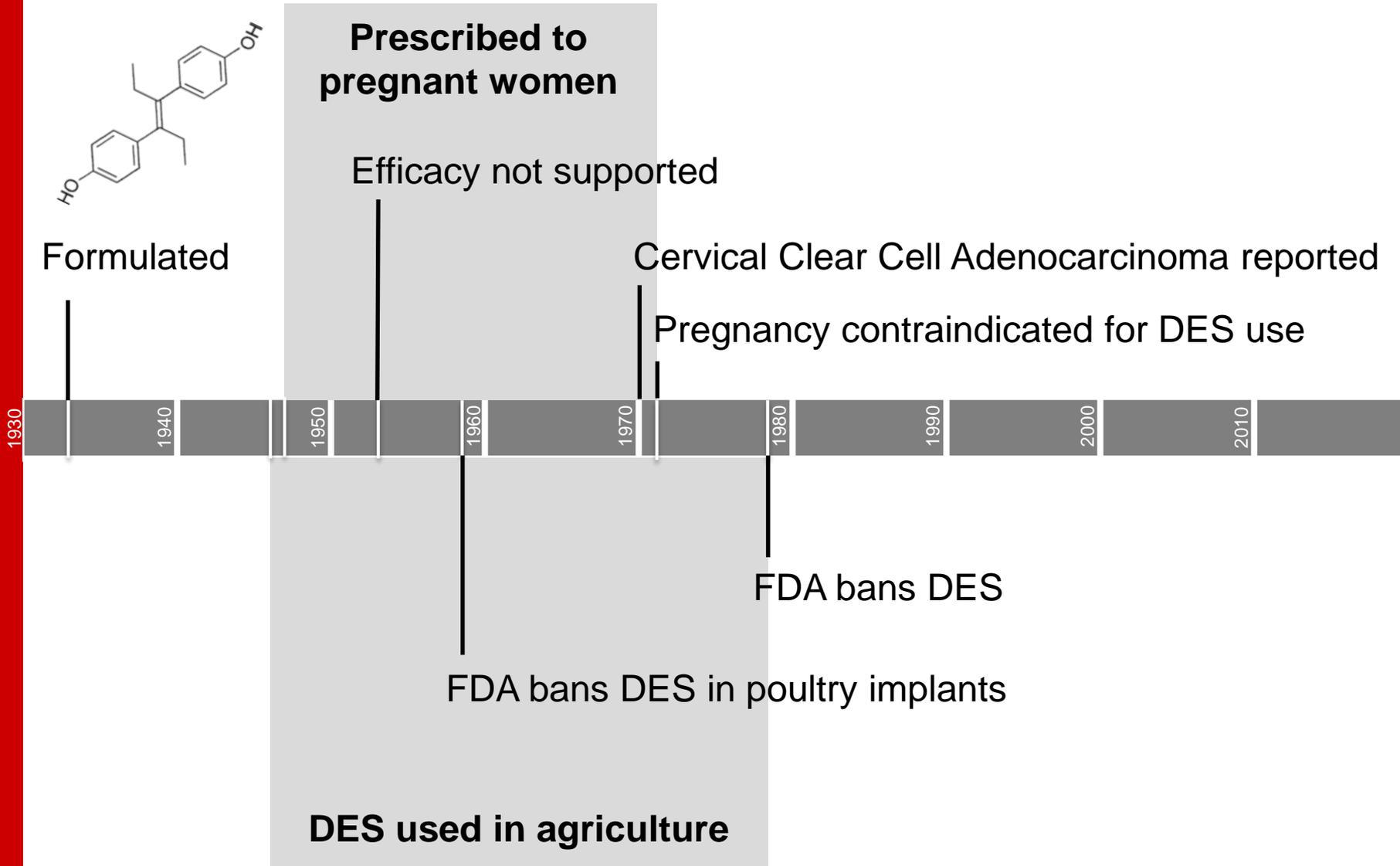
Phenotype



Outline

1. Differential susceptibility to diethylstilbestrol (DES)
1. eQTL meta-analysis in the CC
2. Epigenome profiling in embryonic fibroblasts

History of diethylstilbestrol (DES)



DES health effects

DES daughters

- Reproductive structural abnormalities
- Increased infertility and pregnancy complications
- Early puberty and menopause
- Breast cancer
- Cervical clear cell adenocarcinoma (CCAC)

DES sons

- Reproductive structural abnormalities
- Decreased sperm count
- Testicular cancer?

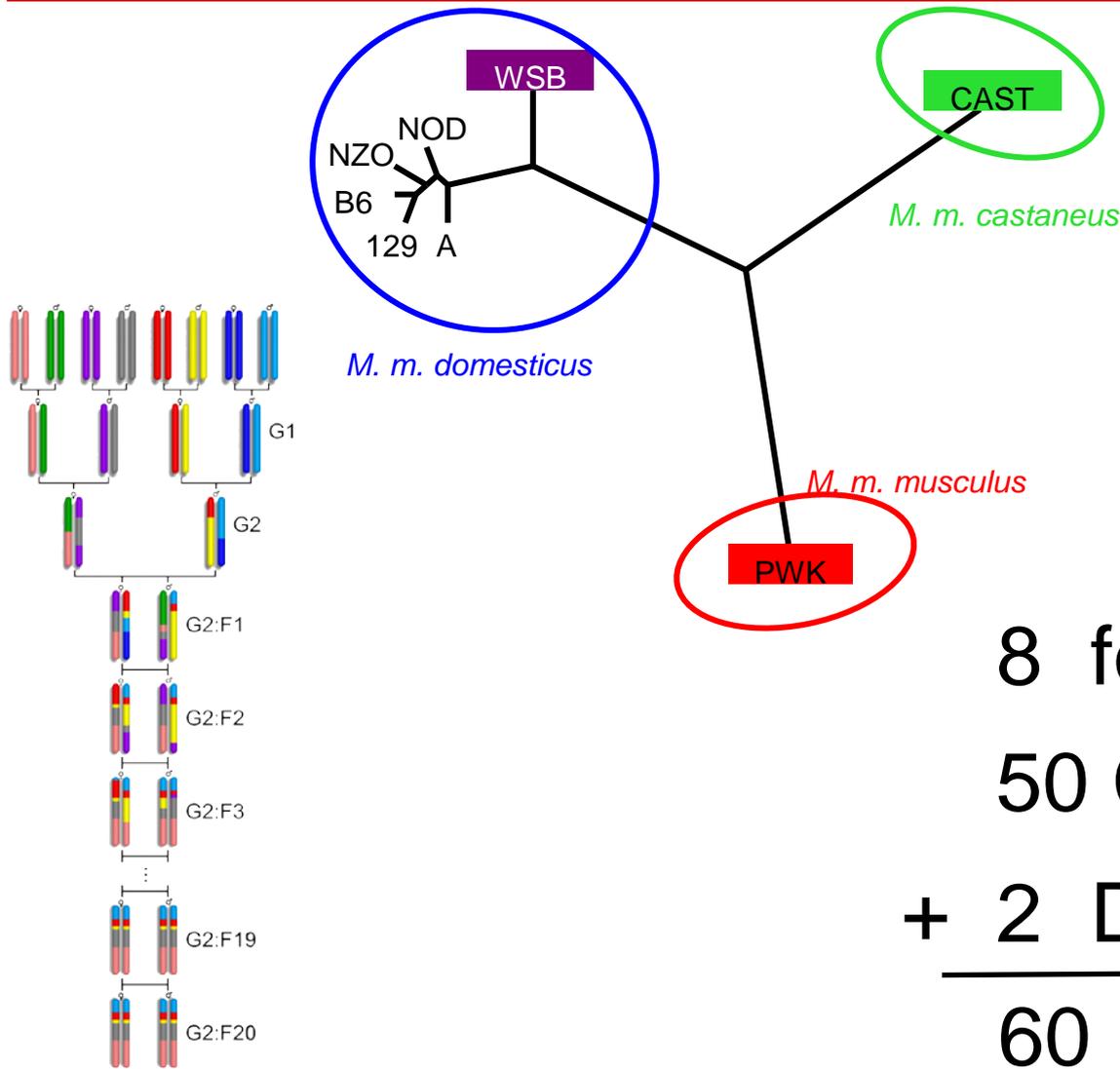
DES female mice

- Uterine cancer
- Uterine dysplasia
- Oviduct malformation
- Uterine gene expression changes

DES male mice

- Decreased sperm count
- Testicular gene expression changes

Strain selection



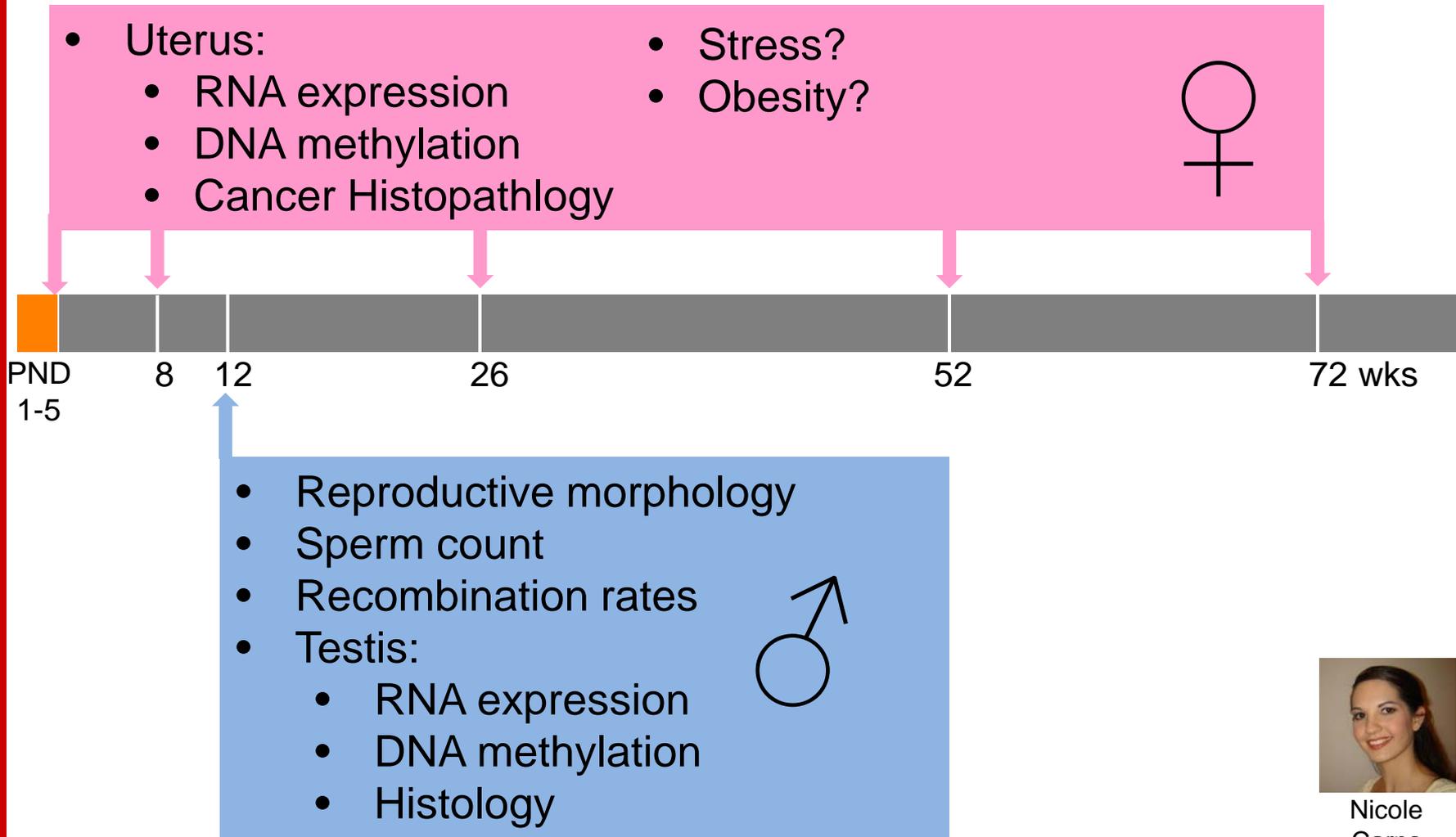
8 founders

50 CC

+ 2 DBA2 & FVB

60

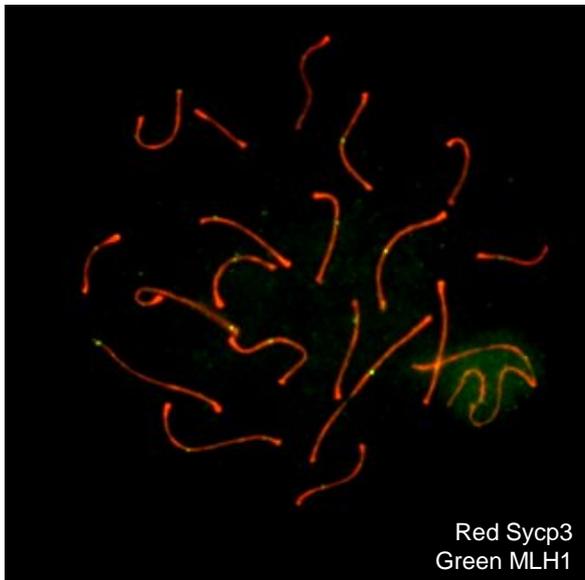
DES exposure and phenotyping



Nicole Carns

Effects of DES exposure in males

- “Curvy” vas deferens
- Reduced gonadal fat pad
- Recombination rate variation?



 PLOS | GENETICS

RESEARCH ARTICLE

Estrogenic Exposure Alters the Spermatogonial Stem Cells in the Developing Testis, Permanently Reducing Crossover Levels in the Adult

Lisa A. Vrooman, Jon M. Oatley, Jodi E. Griswold, Terry J. Hassold, Patricia A. Hunt*

School of Molecular Biosciences, Center for Reproductive Biology, Washington State University, Pullman, Washington, United States of America

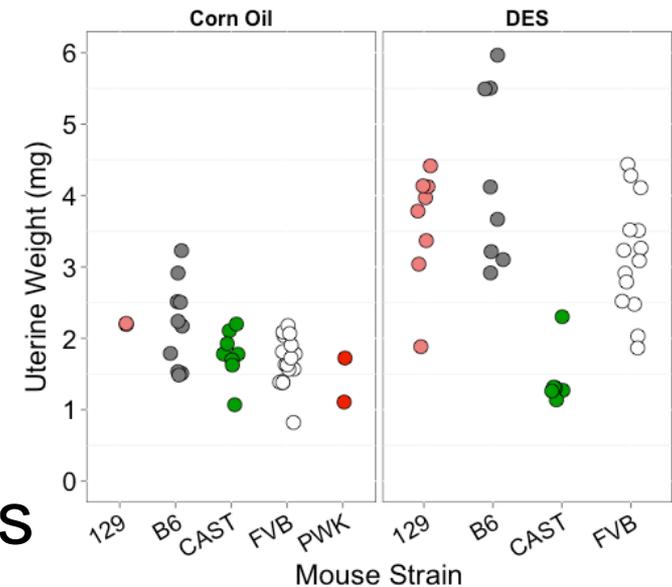
* pathunt@vetmed.wsu.edu



Carlee
Hemphill

Effects of DES exposure in females

- Anxiety in dams (FVB)
- Vascularized uterus (CAST)
- Increased uterine size
- Gene expression changes



RNA-sequencing: Acute effects of DES exposure on 5 day old uterus

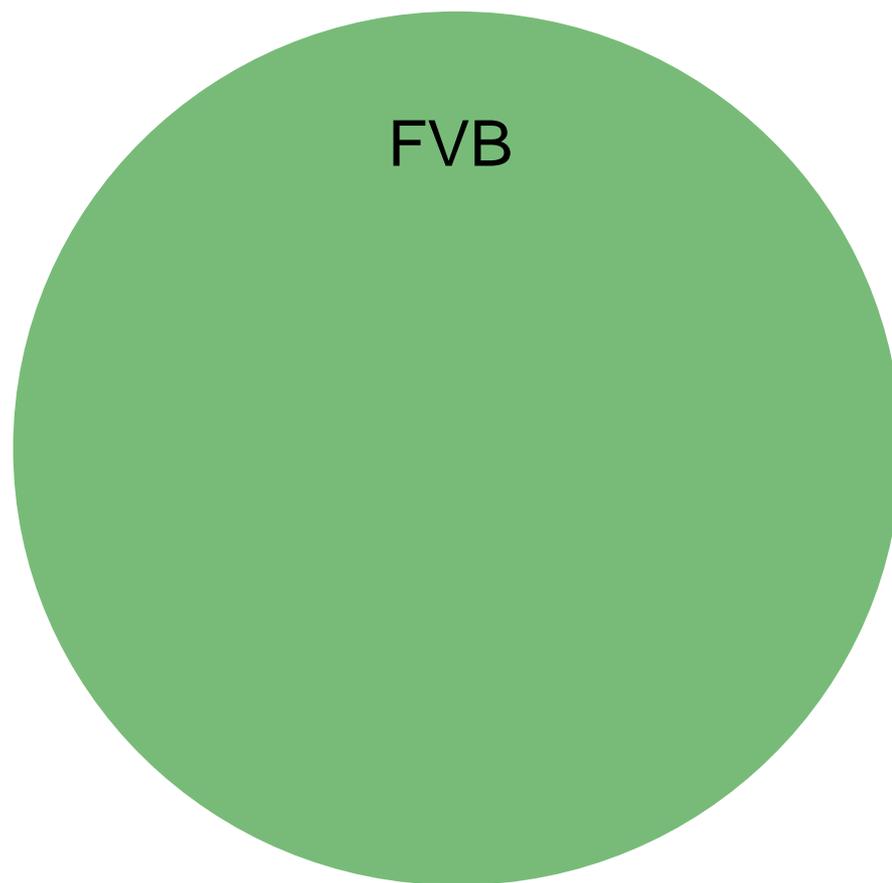
2	X	2	X	6
TREATMENTS		STRAINS		REPLICATES
DES		B6		
Vehicle		FVB		

- One lane HiSeq 100bp SE reads $\sim 11 \times 10^6$ reads/sample
- Aligned to B6 and FVB reference genomes
- 11320 genes average expression ≥ 100 reads/sample



Thomas
Konneker

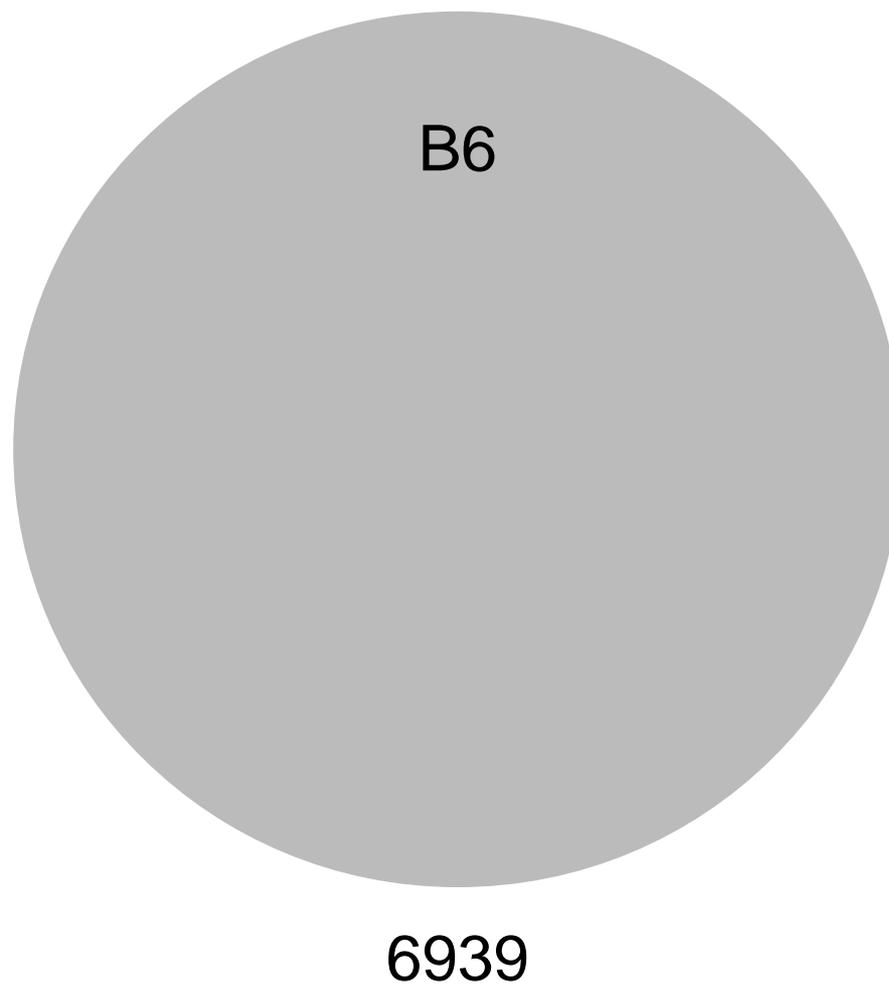
RNA-sequencing – DES v. Vehicle



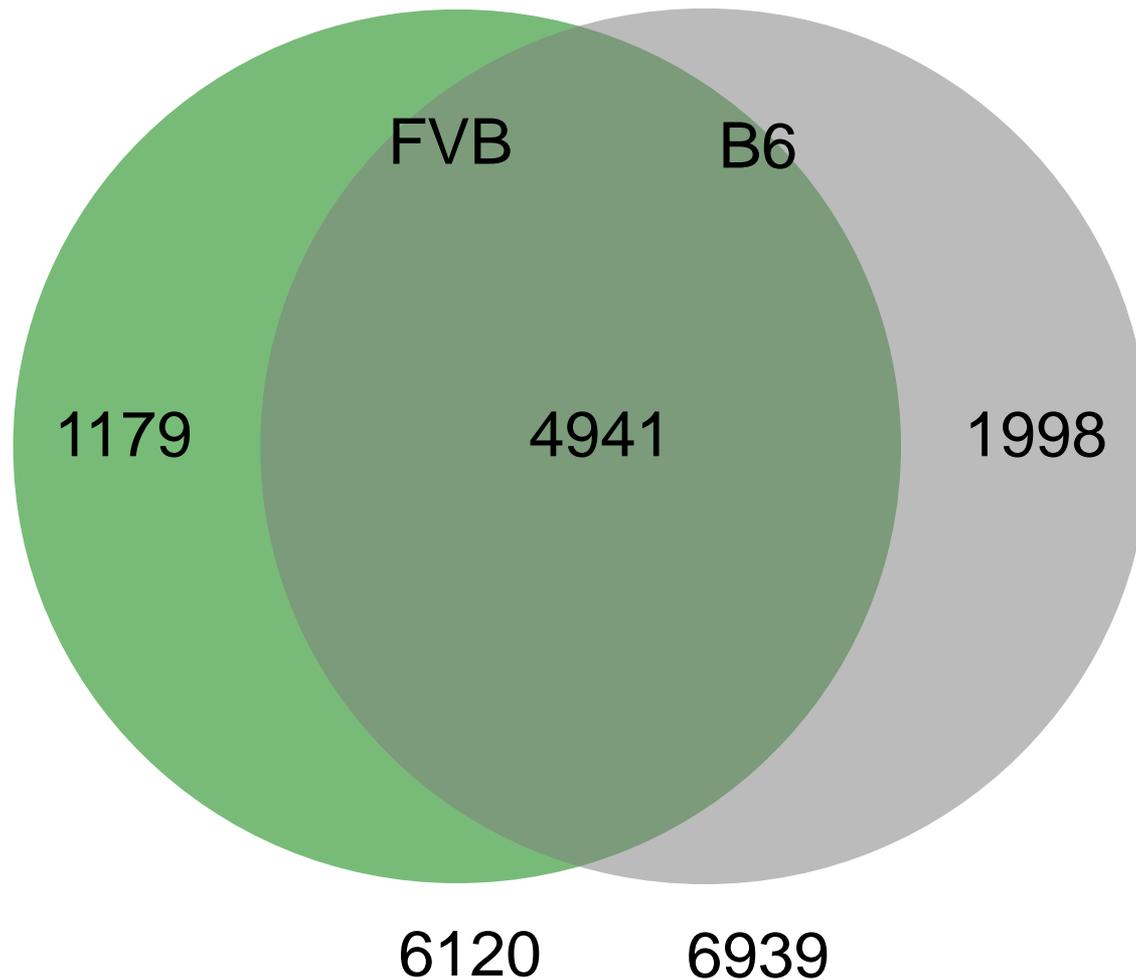
FVB

6120

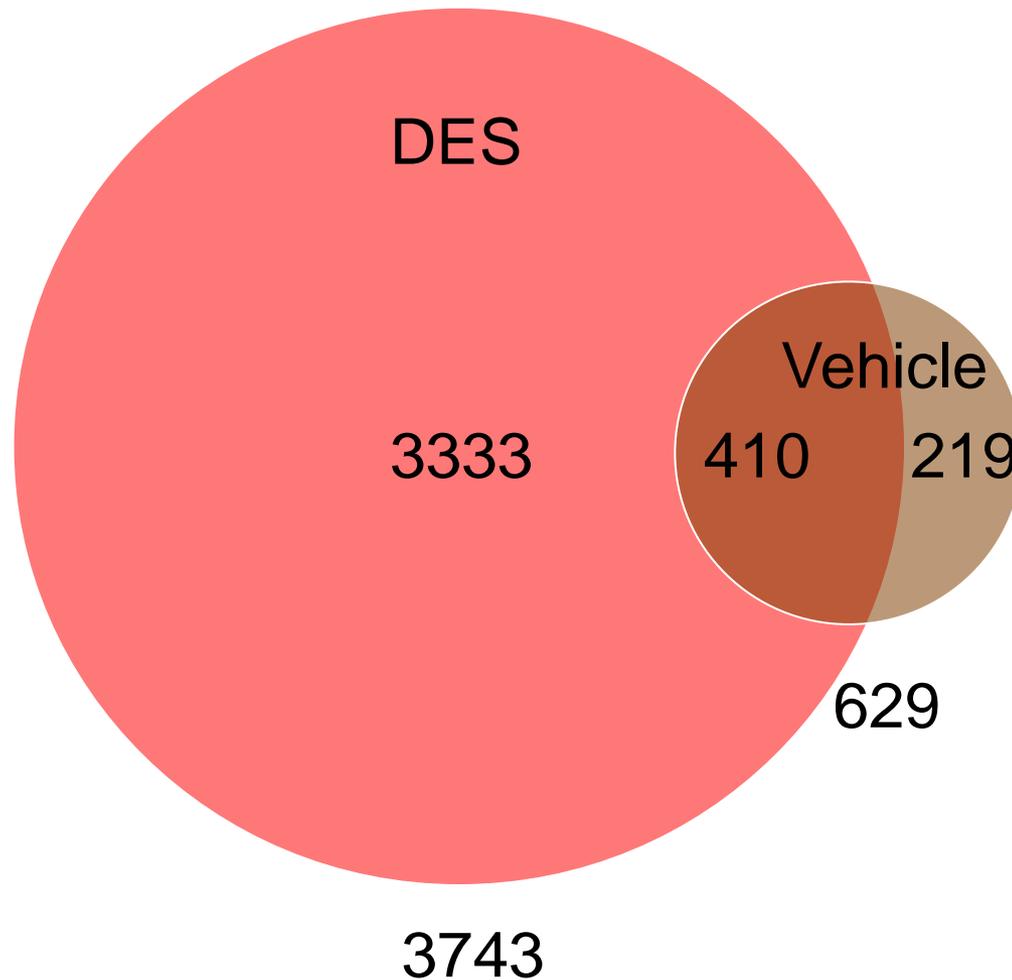
RNA-sequencing – DES v. Vehicle



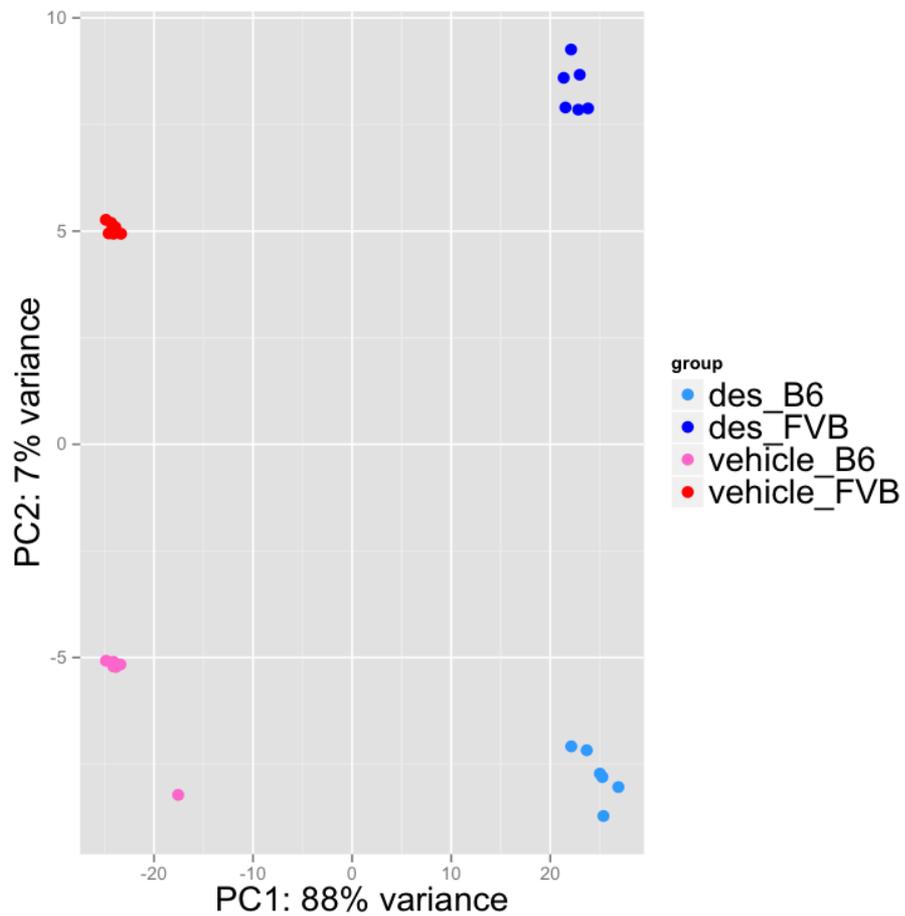
RNA-sequencing – DES v. Vehicle



RNA-sequencing – FVB v. B6



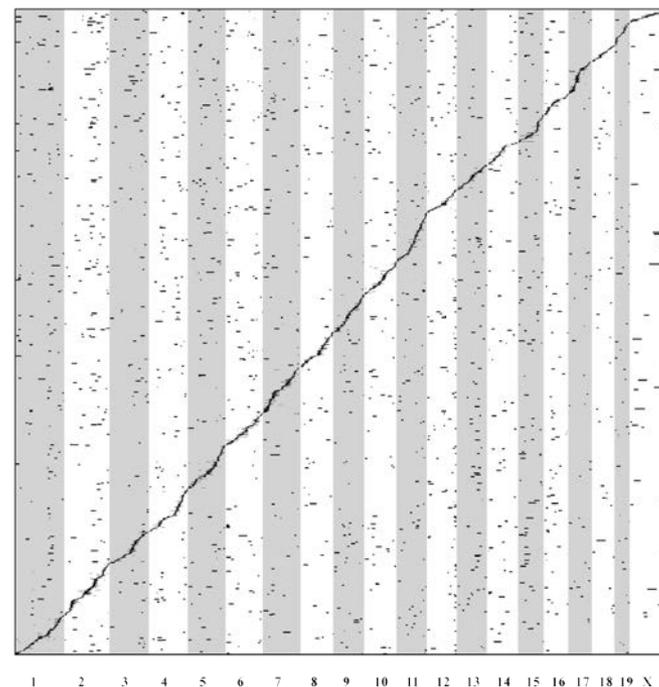
GxE in the DES-exposed transcriptome



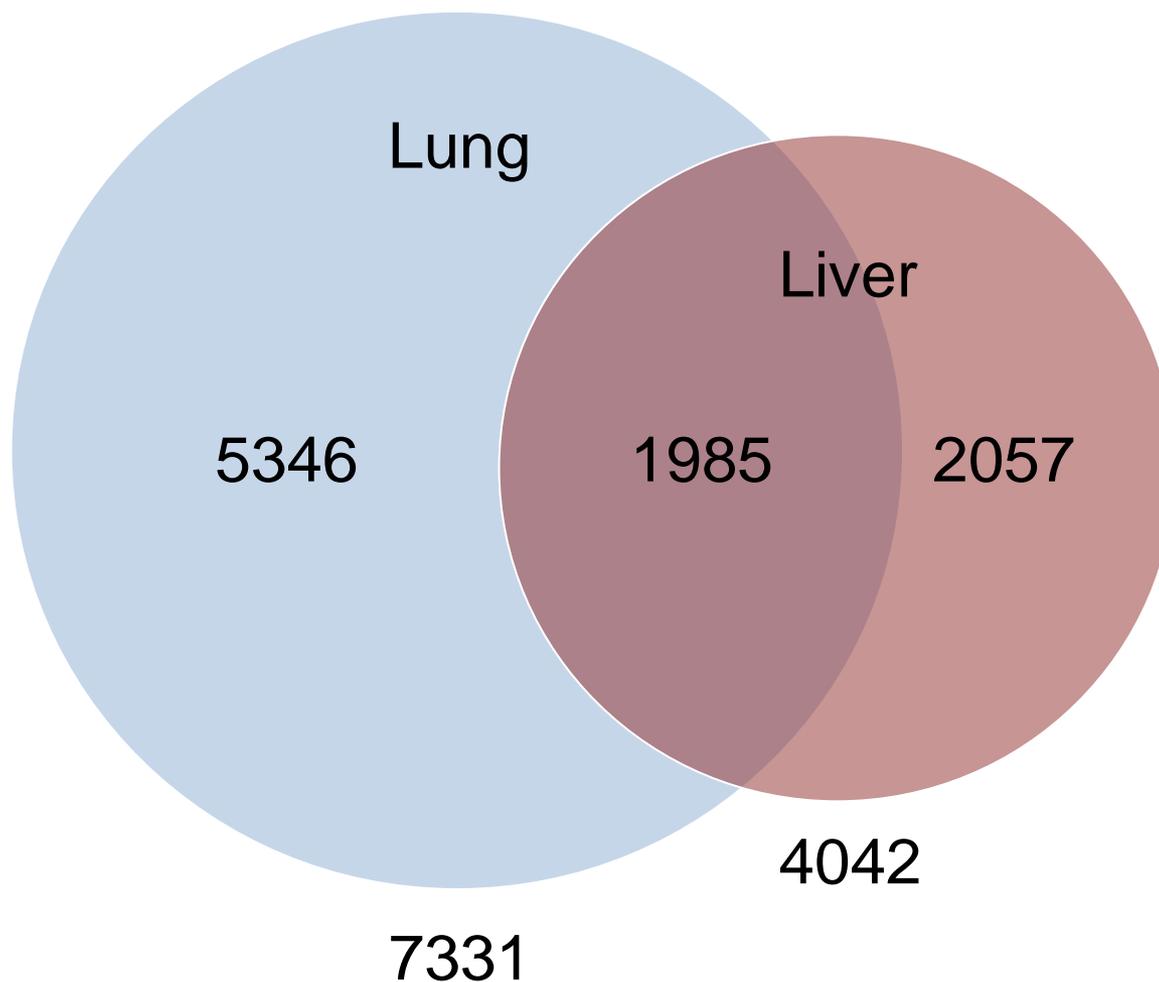
CC gene expression QTL

Study	Year	Local eQTL	Distant eQTL
pre-CC liver	2011	4042	1881
pre-CC lung (asthma)	2013	7331	2523

75-87% of eQTL are local

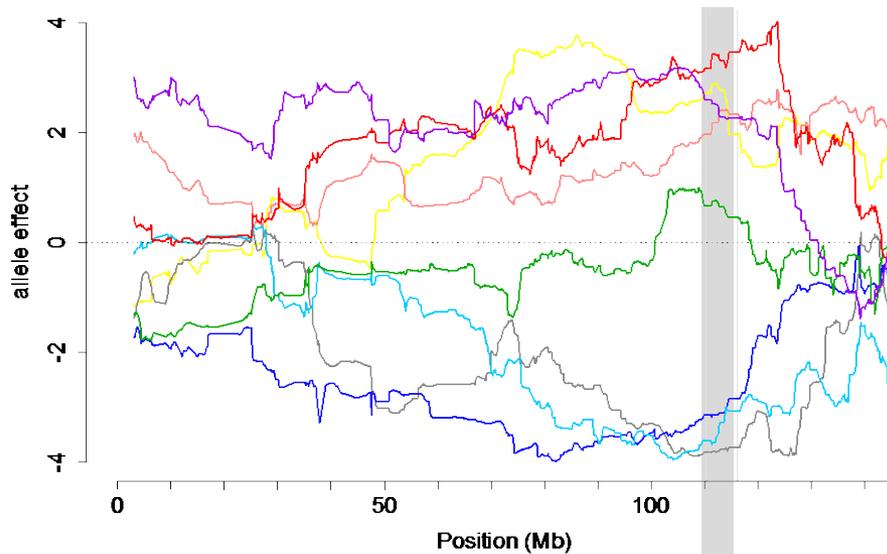


Local eQTL shared in liver and lung



Sam
Widmayer

Allele effect patterns

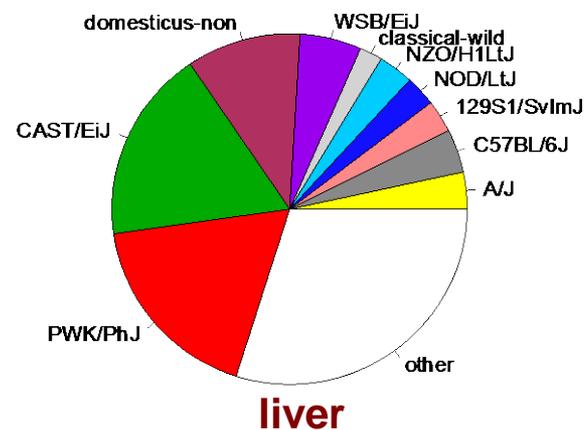
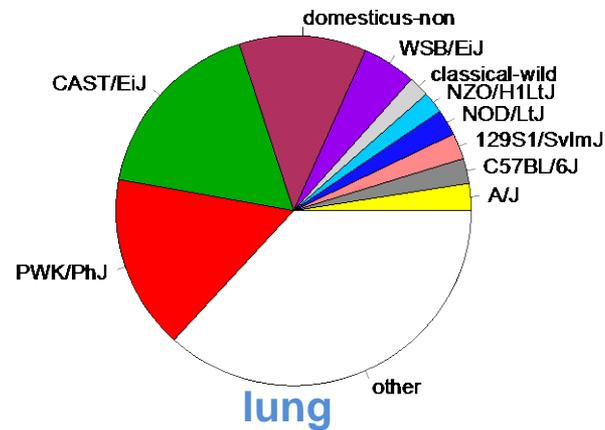
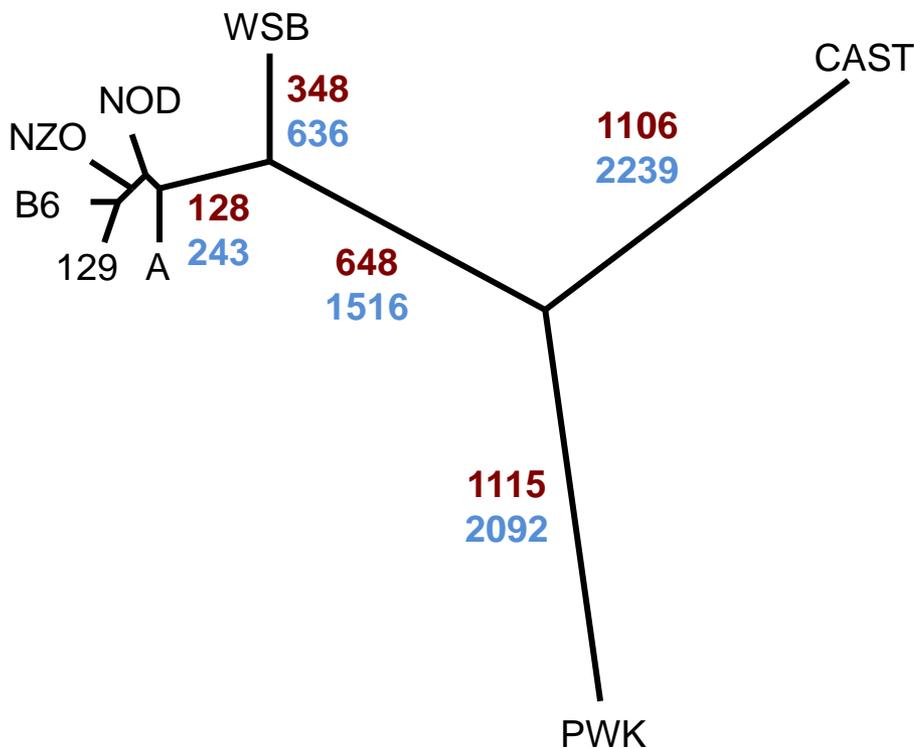


254 possible patterns
 $2^8 - 2$ null patterns

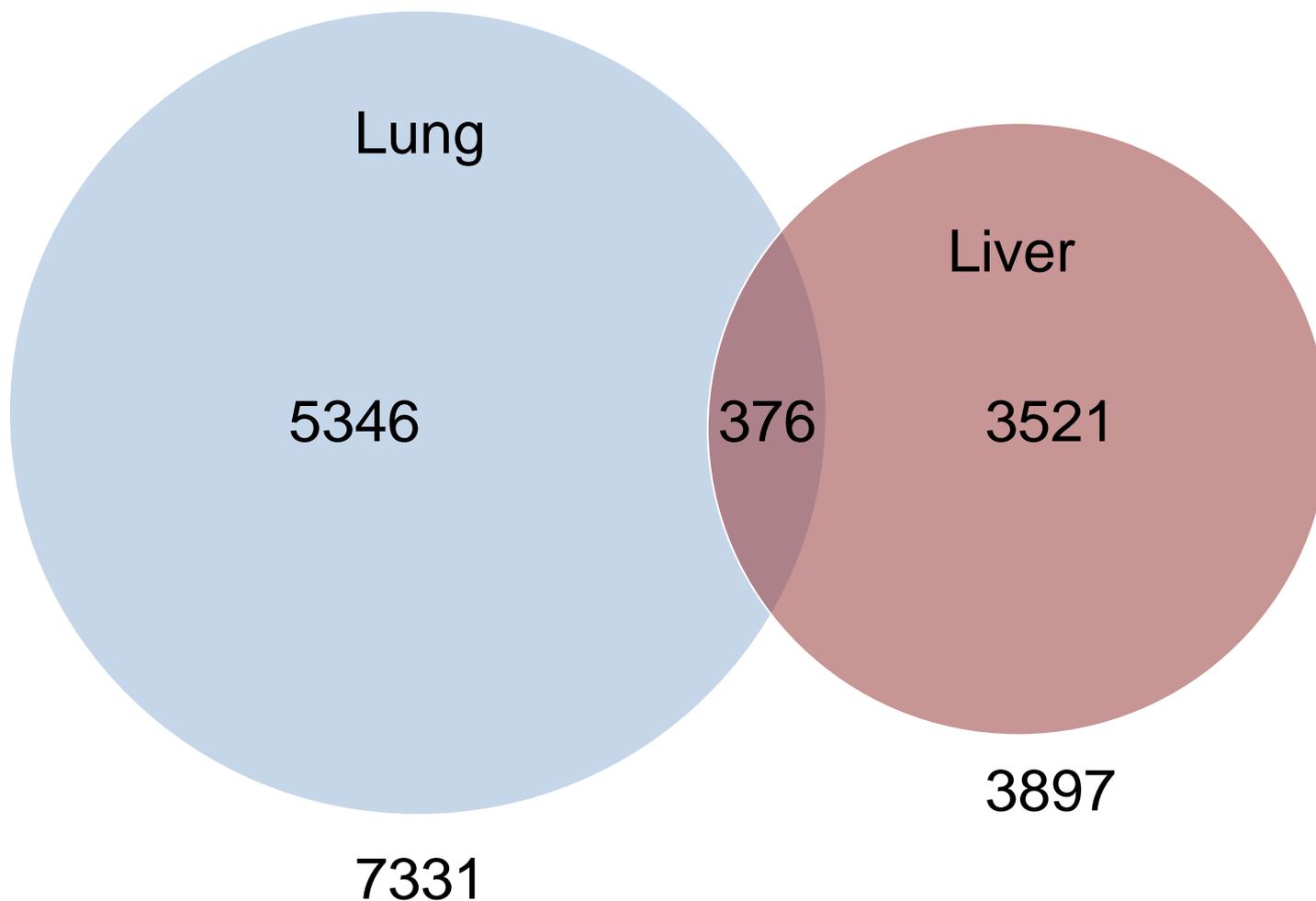
	A	B6	129S1	NOD	NZO	CAST	PWK	WSB
High	1	1	1	1	1	1	1	1
Low	0	0	0	0	0	0	0	0
	1	0	1	0	0	1	1	1

- A/J
- C57BL/6J
- 129S1/SvImJ
- NOD/LtJ
- NZO/HILtJ
- CAST/EiJ
- PWK/PhJ
- WSB/EiJ

Allele effects reflect strain relationships



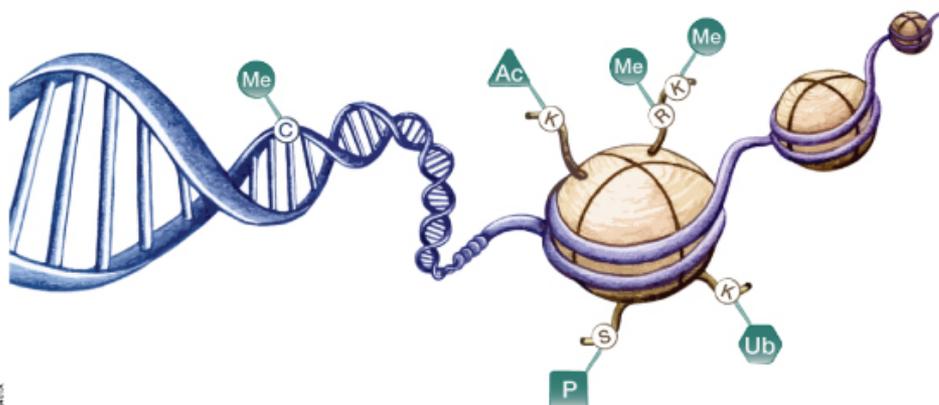
Shared eQTL driven by same alleles



State of eQTL knowledge – 2015

- Abundant
- Local and *cis*-acting
- Allele effects agree between CC, DO, and founder strains
- Proportional to genetic distance
- Shared across tissues

Open chromatin profiling of embryonic fibroblasts



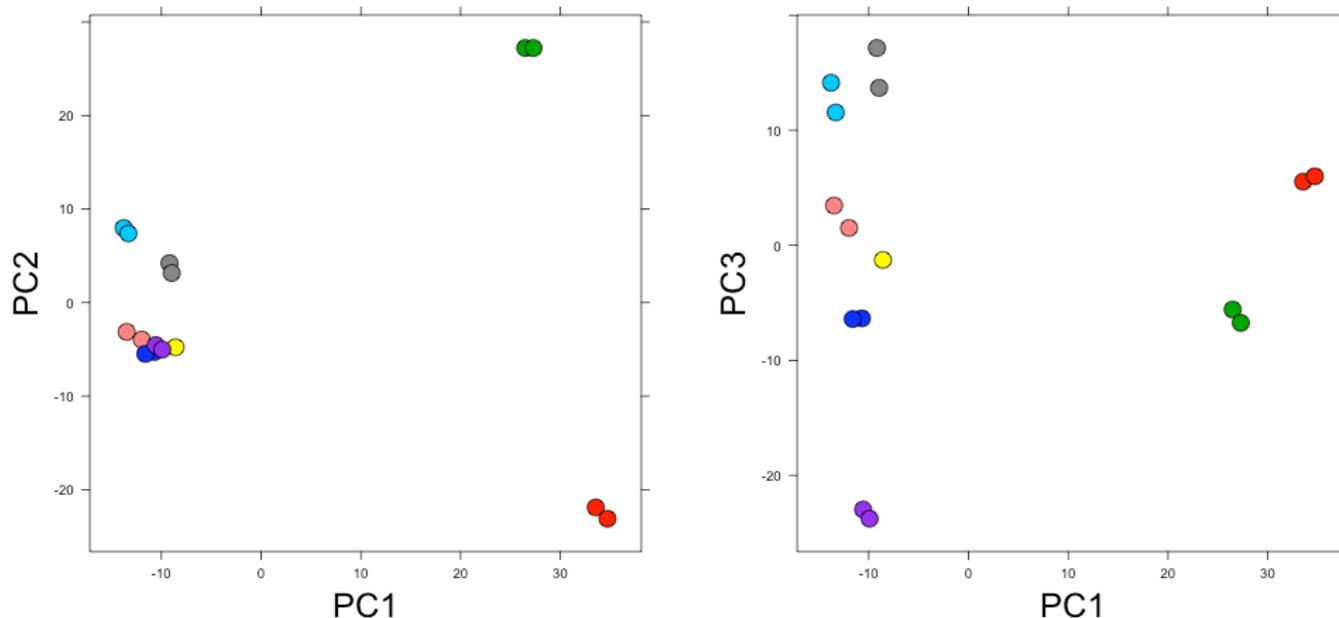
- Embryonic fibroblasts (MEFs) from 8 inbred strains
- Two biological replicates per strain
- 124K-141K peaks/strain
- Average peak size is 169bp
- 249,534 DNaseI hypersensitive sites



Greg
Crawford

Thomas
Konneker

Open chromatin differs by strain



- 51% of DHS peaks vary between strains
- PCA shows genetic diversity is proportional to functional diversity

A/J

C57BL/6J

129S1/SvImJ

NOD/LtJ

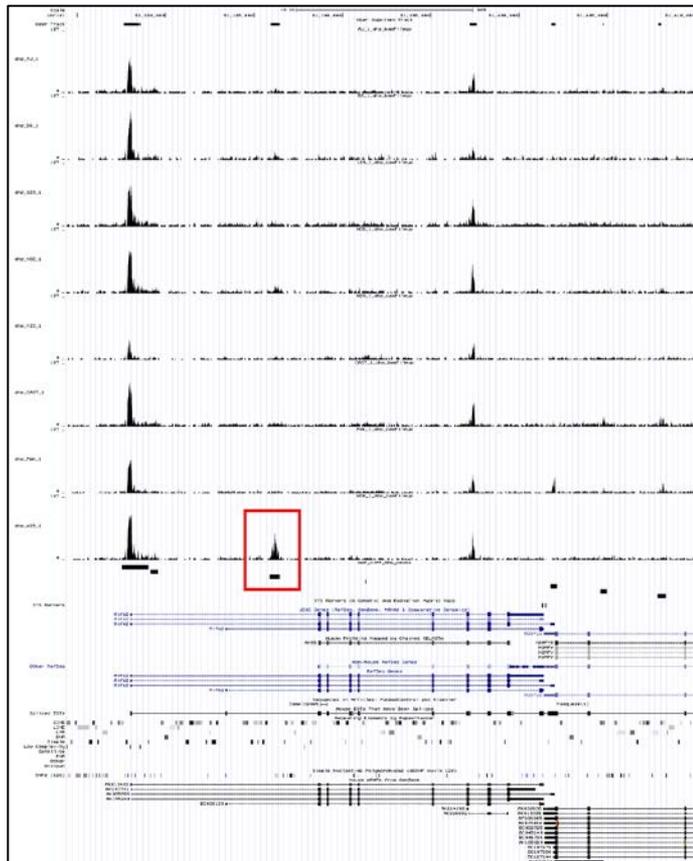
NZO/HILtJ

CAST/EiJ

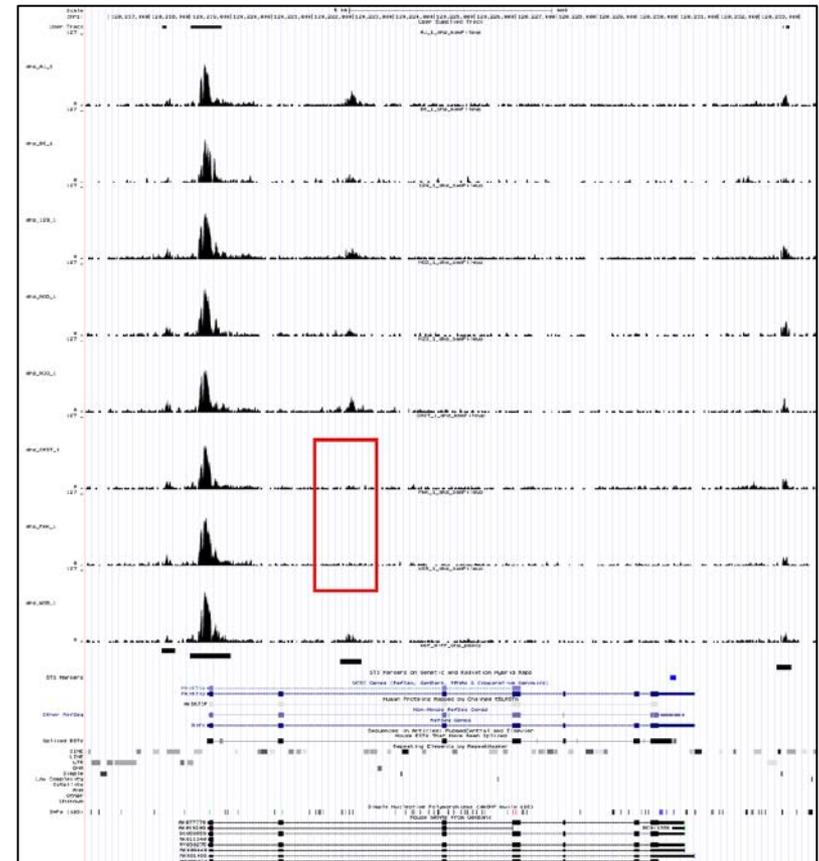
PWK/PhJ

WSB/EiJ

Genetic control of gene regulation



Aifm2



Mki67ip

- A/J
- C57BL/6J
- 129S1/SvImJ
- NOD/LtJ
- NZO/HILtJ
- CAST/EiJ
- PWK/PhJ
- WSB/EiJ

The vast GxE landscape

Genetic Variants 4×10^7

Genes 2.2×10^5

Cell types 2×10^2

Tox21 compounds 1×10^5

1.6×10^{20}

Multiple approaches

- Experimental and computational
- In vivo and ex vivo
- Single compound, many compounds, no compound

Thank you

Aylor Lab



Nicole Carns



Thomas Konneker



Tiffany Garbutt



Kevin Gillespie



Sam Widmayer



Carlee Hemphill

NC STATE UNIVERSITY

Duke

Greg Crawford



UNC

Samir Kelada



NCSU

Dahlia Nielson



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