Curriculum Vitae

Name: Raja Jothi

Office address: National Institute of Environmental Health Sciences (NIEHS)

National Institutes of Health (NIH)

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Research Triangle Park, NC 27709

Telephone: 984-287-3696 Email: jothi@nih.gov

Web: http://www.niehs.nih.gov/research/atniehs/labs/escbl/pi/systemsbiology/

ACADEMIC & PROFESSIONAL POSITIONS

| 2015 - | Senior Investigator (with tenure) | National Institutes of Health (NIH), NIEHS, Epigenetics & Stem Cell Biology Laboratory, Research Triangle Park, NC. |
|-------------|-----------------------------------|--|
| 2009 - 2015 | Investigator (tenure-track) | National Institutes of Health (NIH), NIEHS, Epigenetics & Stem Cell Biology Laboratory, Research Triangle Park, NC |
| 2007 - 2009 | Research Fellow | National Institutes of Health (NIH), NHLBI, Laboratory of Molecular Immunology, Bethesda, MD (Mentor: Keji Zhao) |
| 2004 - 2007 | Research Associate | National Institutes of Health (NIH), NCBI/NLM, Computational Biology Branch, Bethesda, MD (Mentor: Teresa Przytycka) |
| 2001 - 2004 | Research Assistant | University of Texas at Dallas, Department of Computer Science, Richardson, TX |
| 2000 - 2001 | Software Engineer | Westwave Communications (acquired by Alcatel), Richardson, TX |
| 1998 - 1999 | Database Engineer | P'Four Software and Marketing Services, Chennai, India |

HONORS AND AWARDS

| 2019 | Paper of the Year Award, NIEHS, NIH |
|------|---|
| 2018 | Khairallah Lecture, University of Connecticut |
| 2018 | NIH Director's Seminar Series Lecture |
| 2017 | NIH Award of Merit |
| 2017 | Paper of the Year Award, NIEHS, NIH |
| 2016 | Ruth L. Kirschstein Mentoring Award, NIH |
| 2014 | Paper of the Year Award, NIEHS, NIH |
| 2009 | Early Career "Rising Star" Award, NIEHS, NIH |

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EDUCATION

| 2001 - 2004 | Ph.D., Computer Science , University of Texas at Dallas, Richardson, Texas, USA |
|-------------|--|
| 2000 | M.S., Computer Science, University of Texas at Dallas, Richardson, Texas, USA |
| 1994 - 1998 | B.E., Computer Science & Engineering, University of Madras, Chennai, India |

PROFESSIONAL SERVICES

NIEHS Institutional Service

| 2022 - | Co-Chair, Diversity Equity Inclusion & Accessibility Working Group, DIR, NIEHS | |
|-------------|--|--|
| 2022 | Chair, 2022 DIR Retreat Planning Committee | |
| 2020 - 2021 | President, Assembly of Scientists, NIEHS | |
| 2021 - 2022 | Search Committee for Tenure-Track Investigator in Biostatistics and Computational Biology Branch, NIEHS | |
| 2021 | Organizer, Epigenetics & Stem Cell Biology Laboratory Retreat, NIEHS | |
| 2020 | Search Committee for Assistant Scientific Director, DIR, NIEHS | |
| 2020 | Search Committee for the Chief of Comparative Medicine Branch, NIEHS | |
| 2020 | Subject Matter Expert, Search for Computer Scientist (Data Architect), Office of Data Science, NTP, NIEHS | |
| 2018 - 2019 | Chair, Search Committee for Staff Scientist, Stem Cell Biology, ESCBL, NIEHS | |
| 2018 - 2019 | Search Committee for Staff Scientist, Structural Cell Biology, GISBL, NIEHS | |
| 2018 | Selection Committee for the NIEHS Fellow of the Year Award | |
| 2018 | Search Committee for GISBL Structural Cell Biology Staff Scientist, NIEHS2017 - Elected Councilor, Assembly of Scientists, NIEHS, NIH | |
| 2017 - 2018 | Elected Councilor, Assembly of Scientists, NIEHS | |
| 2017 | Search Committee for Tenure-Track Investigator in Transcription, Epigenetics, or Chromatin Biology, NIEHS | |
| 2017 | Search Committee for Biostatistics Staff Scientist, NIEHS | |
| 2017 | Organizer, Epigenetics & Stem Cell Biology Laboratory Retreat, NIEHS | |
| 2016 - | Oversight Committee for the Viral Vector Core, NIEHS | |
| 2016 | Organizing Committee, Division of Intramural Research Retreat, NIEHS | |
| 2016 | Organizer, Epigenetics & Stem Cell Biology Laboratory Retreat, NIEHS | |
| 2015 - 2018 | IT Management Committee, NIEHS | |
| 2015 - 2017 | Committee on Promotions IV to review appointment, tier advancement, and renewal requests of Title 42 Staff Scientists, NIEHS, NIH | |
| 2013 - 2014 | Search Committee for Tenure-Track/Tenure-Eligible Investigator in Biostatistics/Computational Biology, NIEHS, NIH | |
| 2014 | Organizer, Laboratory of Molecular Carcinogenesis Retreat | |
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| 2013 - 2015 | Member, NIEHS Scientific Director's DIR General Council | |
|-------------|---|--|
| 2012 | Stem Cells Committee, NIEHS Cross-Divisional Strategic Plan Implementation | |
| 2012 | Epigenetics Committee, NIEHS Cross-Divisional Strategic Plan Implementation | |
| 2009 - 2013 | Review Committee, Next Generation Sequencing Projects, NIEHS, NIH | |
| 2009 - 2010 | Search Committee for Bioinformatics Staff Scientist, NIEHS, NIH | |

NIH Institutional Service

| 2021 - 2022 | Co-Chair, Search Committee for NIH Earl Stadtman Investigator in Chromosome Biology/Epigenetics/Transcription |
|-------------|---|
| 2020 - 2021 | Search Committee for NIH Earl Stadtman Investigator in Chromosome Biology/Epigenetics/Transcription |
| 2017 - 2018 | Search Committee for NIH Earl Stadtman Investigator in Chromosome Biology/Epigenetics/Transcription |
| 2017 | Review Panel, NIDDK/NIDDK Joint Fellowship Program |
| 2016 - 2017 | Search Committee for NIH Earl Stadtman Investigator in Chromosome Biology/Epigenetics/Transcription |
| 2015 - 2016 | Search Committee for NIH Earl Stadtman Investigator in Chromosome Biology/Epigenetics/Transcription |
| 2013 - 2014 | Search Committee for NIH Earl Stadtman Investigator in Stem Cells |
| 2014 | Review panel, NIH Fellows Award for Research Excellence |
| 2012 - 2013 | Search Committee for NIH Earl Stadtman Investigator in Stem Cells |
| 2013 | Review panel, NIH Fellows Award for Research Excellence |
| 2012 | Review panel, NIH Fellows Award for Research Excellence |
| 2011 | Review panel, NIH Fellows Award for Research Excellence |
| 2010 | Review panel, applications for the NIH National Graduate Student Research Festival |
| 2010 | Review panel, NIH Fellows Award for Research Excellence |

Conference/Symposium Organizing Committee

| 2017 | Organizing Committee - NIEHS/NIH Symposium on Epigenetics & Stem Cells, RTP, NC, June 1-2, 2017 |
|------|--|
| 2015 | Program Committee - International Conference on Bioinformatics and Biomedicine (BIBM), Washington DC, Nov 9-12, 2015. |
| 2014 | Program Committee - International Symposium on Network Enabled Health Informatics, Biomedicine and Bioinformatics, Beijing, China, Aug 18-19, 2014. |
| 2013 | Program Committee - Workshop on Epigenomics and Cell Function, ACM Conference on Bioinformatics, Computational Biology and Biomedical Informatics, Washington D.C., Sep 22-25, 2013. |

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| 2013 | Program Committee - International Symposium on Network Enabled Health Informatics, Biomedicine and Bioinformatics, Niagara Falls, Canada, Aug 25-28, 2013. |
|------|---|
| 2013 | Co-chair - NIEHS/NIH Symposium on Unlocking the Promise of Stem Cells, RTP, NC, April 11-12, 2013. |
| 2012 | Organizing Committee - Toxicology 2012, San Antonio, TX, Sep 17-19 2012. |
| 2012 | Program Committee - International Symposium on Network Enabled Health Informatics, Biomedicine and Bioinformatics Istanbul, Turkey, Aug 2012. |
| 2011 | Workshop Co-chair - IEEE International Conference on Computational Advances in Bio and medical Sciences, Orlando, FL, Feb 2011 |
| 2010 | Co-chair - Symposium on "Epigenetics, Chromatin, and Gene Regulation", NIH Research Festival, Bethesda, MD, Oct 2010. |
| 2010 | Program Committee - ACM International Conference on Bioinformatics and Computational Biology (ACM-BCB), Niagara Falls, NY, Aug 2-4, 2010 |
| 2010 | Program Committee - The 8th International Bioinformatics Workshop (IBW), Wuhan, China, June 4-6, 2010 |
| 2009 | Chair - Symposium on "Epigenetics, Chromatin, and Gene Regulation", NIH Research Festival, Bethesda, MD, Oct 2009. |
| 2009 | Program Committee - The IEEE International Conference on Bioinformatics and Biomedicine (BIBM), Washington DC, Nov 1-4, 2009 |
| 2008 | Vice-chair - The IEEE International Conference on Bioinformatics and Biomedicine (BIBM), Philadelphia, CA, Nov 7-9, 2008. |
| 2007 | Program Committee - 15th Annual International Conference on Intelligent Systems for Molecular Biology (ISMB) & 6th European Conference on Computational Biology (ECCB), Vienna, Austria, Jul 21-25, 2007. |
| 2007 | Program Committee - The IEEE International Conference on Bioinformatics and Biomedicine (BIBM), Silicon Valley, CA, Nov 2-4, 2007. |
| 2004 | Session Chair, 7th INFORMS Telecommunications Conference, Boca Raton, FL, 2004. |
| 2004 | Session Chair, 3rd IEEE International Conference on Networking, 2004. |
| 2003 | Session Chair, 15th IASTED International Conference on Parallel and Distributed Computing and Systems, Las Vegas, NV, 2003. |
| | |

Ad-hoc Reviewer

Grants Wellcome Trust

The Biotechnology and Biological Sciences Research Council (BBSRC) – UK

Journals Nature Genetics

Nature Communications

Nature Methods Nature Protocols

PNAS

Science Advances

Cell Reports Cell Systems

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Genome Research

Genome Biology

Nucleic Acids Research

eLife

Bioinformatics

Trends in Genetics

PLoS Genetics

PLoS Computational Biology

PLoS ONE

Molecular Biology and Evolution

Stem Cells and Development

Proteins

Oncotarget

Molecular Biosystems

BMC Bioinformatics

BMC Genomics

BMC Systems Biology

Toxicological Sciences

Toxicology and Applied Pharmacology

IEEE/ACM Transactions on Computational Biology and Bioinformatics

IEEE Transactions on NanoBioscience

Pattern Recognition Letters

Discrete and Computational Geometry

Networks

Journal of Graph Algorithms and Applications

Journal Editorial Board

2011 - Frontiers in Bioinformatics and Computational Biology

2010 - 2017 PLoS ONE

INVITED TALKS

Mar 2020 **Keystone Symposium on Systems Biology: Global Regulation of Gene Expression**, NY – Bivalent chromatin does not poise genes for rapid activation

Jan 2020 **Keystone Symposium on Chromatin and Epigenetics**, Keystone, CO – Bivalent chromatin does not poise genes for rapid activation

Nov 2019 **Society for Scientific Advancement**, 8th Annual Conference, Kingston, Jamaica – Transcriptional and epigenetic control of cell fate decisions

Nov 2019 **North Carolina State University**, NC – Uncovering unknown unknowns of gene regulation

Nov 2018 Wake Forest University, NC – Uncovering unknown unknowns of gene regulation

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- Oct 2018 Khairallah Lecture, University of Connecticut, CT Uncovering unknown unknowns of gene regulation
- Sep 2018 **CSHL Meeting on Meeting on Epigenetics & Chromatin**, NY Crm1 promotes the formation of broad H3K27me3 repressive domains at developmental genes (presenter: Conway AE)
- May 2018 **CSHL Meeting on Nuclear Organization & Function**, NY Crm1 promotes the formation of broad H3K27me3 repressive domains at developmental genes (presenter: Conway AE)Apr 2018 **NIH Director's Seminar Series Lecture**, NIH, MD Uncovering unknown unknowns of Gene Regulation
- Oct 2017 University of North Carolina, NC Uncovering unknown unknowns of Gene Regulation
- Oct 2017 NCBI Retreat Keynote, NLM, NIH, MD Uncovering unknown unknowns of Gene Regulation
- Sep 2017 **CSHL Meeting on Mechanisms of Eukaryotic Transcription**, NY Transcription at intragenic enhancers attenuate gene expression
- May 2016 **CSHL Meeting on Nuclear Organization & Function**, NY A non-canonical role for nuclear export receptor CRM1 in developmental gene regulation (presenter: Conway AE)
- May 2016 NAEHS Council Meeting, NIEHS, NIH, NC –Why not to look under the lamppost?
- Mar 2016 **Keystone Symposium on Chromatin and Epigenetics**, Whistler, Canada Transcription at intragenic enhancers attenuates gene expression.
- Mar 2015 **Symposium on Systems Biology of Stem Cells**, Oberstdorf, Germany– Gene networks controlling ES cell identity and homeostasis.
- Oct 2014 **Systems Biology Forum, NIH**, Bethesda, MD Gene networks controlling embryonic stem cell identity and homeostasis.
- Sep 2014 **NHGRI, NIH**, Bethesda, MD Gene networks controlling embryonic stem cell identity and homeostasis.
- Apr 2013 **Symposium on Unlocking the Promise of Stem Cells**, RTP, NC ES cell identity and homeostasis.
- Sep 2012 **Genomics Day Lecture, NIEHS, NIH**, RTP, NC Meta-analysis identifies determinants of embyronic stem cell identity.
- Aug 2012 **FASEB Meeting** on Biological Methylation: From DNA and Histones to Disease, Snowmass, CO Tet1-dependent 5hmC is required for maintaining pluripotency.
- Oct 2011 Laboratory of Toxicology and Pharmacology (LTP), NIEHS, NIH, RTP, NC Embryonic stem cells and gene regulation.
- Sep 2011 **Bertinoro Computational Biology Meeting** on Computational Methods in Functional Genomics, Bertinoro, Italy Embryonic stem cells and gene regulation.

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- Sep 2011 **Laboratory of Molecular Carcinogenesis (LMC), NIEHS, NIH**, RTP, NC esBAF facilitates pluripotency by conditioning the genome for LIF/STAT3 signaling and by regulating Polycomb function.
- Apr 2011 **University of Texas at Dallas,** Richardson, TX Embryonic stem cells and gene regulation.
- Mar 2011 **New York University,** New York, NY Embryonic stem cells and gene regulation.
- Feb 2011 **University of Michigan**, Ann Arbor, MI Embryonic stem cells and gene regulation.
- Oct 2010 **Duke University,** Institute for Genome Sciences & Policy, Durham, NC Embryonic stem cells and gene regulation
- Oct 2010 **Symposium on Epigenetics, Chromatin, and Gene Regulation**, Bethesda, MD esBAF conditions the pluripotent genome for LIF/STAT3 signaling by opposing polycomb
- Mar 2010 **Keystone Meeting** on Biomolecular Interaction Networks: Function and Disease, Quebec, Canada Genomic analysis reveals a tight link between transcription factor dynamics and regulatory network architecture.
- Feb 2010 **North Carolina State University**, Department of Environmental and Molecular Toxicology, Raleigh, NC Systems biology and epigenetics of gene regulation
- Dec 2009 **RECOMB Conference on Regulatory Genomics,** Cambridge, MA Genomic analysis reveals a tight link between transcription factor dynamics and regulatory network architecture.
- Nov 2009 **Laboratory of Signal Transduction (LST), NIEHS, NIH**, RTP, NC Seeing the forest for the trees: stories on Brg1 and CTCF.
- Nov 2009 **NIEHS Early Career Award Lecture, NIH**, RTP, NC Seeing the forest for the trees: a systems level understanding of differential cell-fate outcome.
- Oct 2009 **Indian Institute of Technology**, Chennai, India Systems biology and epigenetics of gene regulation.
- Oct 2009 **Center for Cellular and Molecular Biology (CCMB)**, Hyderabad, India Systems biology and epigenetics of gene regulation.
- Oct 2009 **Center for DNA Finger Printing and Diagnostics (CDFD)**, Hyderabad, India Systems biology and epigenetics of gene regulation.
- Oct 2009 **Indian Institute of Science**, Bangalore, India Systems biology and epigenetics of gene regulation.
- Sep 2009 **Biostatistics Branch, NIEHS, NIH**, RTP, NC Systems biology and epigenetics of gene regulation.
- Nov 2008 **RECOMB Conference on Regulatory Genomics,** Cambridge, MA Genome-wide identification of in vivo protein-DNA binding sites from ChIP-Seq data.
- Jun 2008 **NIEHS, NIH**, RTP, NC Chromatin modifications, gene expression, and regulatory networks.

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- May 2008 NCBI/NLM, NIH, Bethesda, MD Chromatin modifications, gene expression, and regulatory networks.
- Jan 2008 **Rutgers University**, Camden, NJ Regulatory proteins within a hierarchical framework have distinct dynamic properties.
- Aug 2007 **Department of Defense Biotechnology HPC Software Applications Institute (BHSAI)**, Fort Detrick, Frederick, MD Inferring protein and domain interactions using sequence co-evolution and combinatorial optimization approaches.
- May 2007 **NCBI/NLM, National Institutes of Health (NIH)**, Bethesda, MD Inferring molecular interactions using sequence co-evolution and co-inheritance: biases, strengths and weaknesses
- Apr 2007 **George Mason University**, Fairfax, VA Co-evolution (correlated mutations) as an indicator of protein and domain interactions.
- Mar 2007 **Philips Research**, Briarcliff Manor, NY– Co-evolution as an indicator of protein and domain interactions.
- Feb 2007 **University of Connecticut**, Storrs, CT Co-evolution as an indicator of protein and domain interactions.
- Oct 2006 **9th Annual Computational Genomics Conference**, Baltimore, MD Co-evolutionary analysis of domains in interacting proteins reveals insights into domain-domain interactions mediating protein-protein interactions.
- Sep 2005 **NCBI/NLM**, **National Institutes of Health (NIH)**, Bethesda, MD A new phylogenetic approach to delineate orthologous groups
- Jun 2005 International Conference on Intelligent Systems for Molecular Biology (ISMB), Detroit, MI –Predicting protein-protein interaction by searching evolutionary tree automorphism space.
- May 2005 **DIMACS Workshop on Biomolecular Networks: Topological Properties and Evolution**, Rutgers University, New Brunswick, NJ An Evolution-Based Clustering Method to Separate Orthologous Genes from Out-Paralogs.
- Dec 2004 **University of Maryland**, College Park, MD The effects of evolutionary tree topology on predicting protein interaction specificity.
- Dec 2004 **Georgetown University**, Washington D.C The effects of evolutionary tree topology on predicting protein interaction specificity.
- Apr 2004 **NCBI/NLM, National Institutes of Health (NIH)**, Bethesda, MD Protein folding in the Hydrophobic-Hydrophilic Model.
- Mar 2004 **7th INFORMS Telecommunications Conference**, Boca Raton, FL Survivable Network design: the capacitated minimum spanning network problem.
- Nov 2003 International Conference on Parallel and Distributed Computing and Systems, Marina del Rey, CA Design of local access networks.

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- May 2003 **DIMACS Workshop on Geometric Optimization,** Rutgers University, New Brunswick, NJ Leave no stones unturned: improved approximation algorithms for degree-bounded minimum spanning trees.
- Apr 2003 **University of Maryland,** College Park, MD Approximation algorithms for capacitated minimum spanning tree problem and its variants in network design.
- Jan 2003 **14th ACM-SIAM Symposium on Discrete Algorithms**, Baltimore, MD A 5/4-approximation algorithm for minimum 2-edge-connectivity.

MENTORING

Trainees

| Name Sailu Yellaboina* | Type Postdoctoral Fellow | Duration 2009 – 2011 | Current Position Associate Professor (with tenure), All India Institute of Medical Sciences, Hyderabad, India |
|---------------------------|---------------------------------------|-----------------------------|--|
| Johannes Freudenberg* | Postdoctoral Fellow | 2010 – 2011 | Director, Clincial Biomarker Analytics at GlaxoSmithKline, USA |
| Leelavati Narlikar* | Special Volunteer | 2009 – 2013 | Associate Professor and Deputy Chair, Indian Institute of Science Education and Research (IISER), Pune, India |
| Viju Mathew [†] | High School Student Summer Intern | 2010, 2011 | Software Engineer, Funding Circle US |
| Swati Ghosh* | Postdoctoral Fellow | 2010 – 2012 | Postdoctoral Fellow, University of Colorado, Denver |
| Pengyi Yang* | Postdoctoral Fellow | 2013 – 2015 | Associate Professor (with tenure), University of Sydney, Australia |
| Senthilkumar Cinghu* | Postdoctoral Fellow & Staff Scientist | 2011 – 2021 | Senior Scientist, Life Edit Therapeutics, NC, USA |
| Andrew Oldfield* | Postdoctoral Fellow | 2012 – 2016 | Staff Scientist (with tenure), CNRS, France |
| Rajneesh Pathania* | Postdoctoral Fellow | 2015 – 2019 | Veterinarian, Banfield Hospital, Greensboro, NC |
| Brian Deskin* | Postdoctoral Fellow | 2016 – 2019 | Assistant Professor, Tulane University, LA, USA |
| Wilfred Wong | Undergraduate Summer Intern | 2018 | Doctoral Student, Weill Cornell |
| Julie Dickerson | Undergraduate Summer Intern | 2018, 2019 | MD, PhD Student, Medical University of South Carolina |
| Amanda Conway* | Postdoctoral Fellow | 2013 – 2020 | Medical Writer & Consultant, |

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Dhirendra Kumar* Postdoctoral Fellow 2016 – 2020 Staff Scientist, NIEHS, USA

Megan Justice Postdoctoral Fellow 2021 –

AWARDS/HONORS TO TRAINEES

- Dhirendra Kumar, Ph.D., Postdoctoral Fellow (2016 2020)
 - ✓ NIH Fellows Award for Research Excellence (2019-2020)
 - ✓ Winner, "Big Picture, Small Talk" competition, NIEHS (2018)
- Brian Deskin, Ph.D., Postdoctoral Fellow (2016 2019)
 - ✓ NIH Fellows Award for Research Excellence (2017-2018)
- Rajneesh Pathania, Ph.D., Postdoctoral Fellow (2015 2019)
 - ✓ NIH Fellows Award for Research Excellence (2016-2017)
 - ✓ NIEHS Science Day Best Poster Award (2017)
- Amanda Conway, Ph.D., Postdoctoral Fellow (2013 2020)
 - ✓ Oral Presentation, CSHL Meeting on Chromatin & Epigenetics (2018)
 - ✓ Oral Presentation, CSHL Meeting on Nuclear Function & Organization (2018)
 - ✓ NIGMS PRAT Fellowship (2014 2017)
 - ✓ NIH Fellows Award for Research Excellence (2017-2018)
 - ✓ Oral Presentation, CSHL Meeting on Nuclear Function & Organization (2016)
 - ✓ NIH Fellows Award for Research Excellence (2015-2016)
- Pengyi Yang, Ph.D., Postdoctoral Fellow, Ph.D., (2013 present)
 - ✓ NIH Fellows Award for Research Excellence (2015-2016)
 - ✓ NIH Fellows Award for Research Excellence (2014-2015)
- Andrew Oldfield, Ph.D., Postdoctoral Fellow (2012 2016)
 - ✓ NIH Fellows Award for Research Excellence (2014-2015)
 - ✓ Intramural Research Paper of the Year (2014)
 - ✓ Best Scientific Presentation, LMC Retreat (2014)
 - ✓ Intramural Research Paper of the Month (Oct 2014)
- Senthilkumar Cinghu, Ph.D., Postdoctoral Fellow (2011 2016)
 - ✓ Intramural Research Paper of the Year (2017)
 - ✓ Intramural Research Paper of the Month (Dec 2017)
 - ✓ NIH Fellows Award for Research Excellence (2014-2015)
 - ✓ NIH Fellows Award for Research Excellence (2013-2014)
 - ✓ Intramural Research Paper of the Month (June 2014)
- Swati Ghosh, Ph.D., Postdoctoral Fellow (2010 2012)
 - ✓ NIH Fellows Award for Research Excellence (2012-2013)
- Johannes Freudenberg, Ph.D., Postdoctoral Fellow (2010 2011)
 - ✓ NIH Fellows Award for Research Excellence (2011-2012)

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^{*}Recipient, NIH Fellows Award for Research Excellence (FARE)

[†] High school student; contributing author on a manuscript published in Nucleic Acids Research (2013), >100 citations

- ✓ Invited Oral Presentation, NIH Research Festival, Bethesda, MD (2011)
- ✓ Intramural Research Paper of the Month (June 2014)
- Leelavati Narlikar, Ph.D., Special Volunteer (2009 2013)
 - ✓ NIH Fellows Award for Research Excellence (2011-2012)
 - ✓ Wellcome Trust/DBT Early Career Fellowship (2012 2016)
 - ✓ Ramanujan Fellowship (2010 2011)
- Sailu Yellaboina, Ph.D., Postdoctoral Fellow (2009 2011)
 - ✓ NIH Fellows Award for Research Excellence (2010-2011)
 - √ Young Investigator Fellowship, IndiaBioScience.org (2011)
 - ✓ Invited Oral Presentation, NIH Research Festival, Bethesda, MD (2010)

PUBLICATIONS

Peer-Reviewed Articles (in reverse chronological order)

- † Co-first author
- * Corresponding/Co-corresponding author
 - 1. Kumar D, Cinghu S, Oldfield AJ, Yang P, **Jothi R*** (2021). Decoding the function of bivalent chromatin in development and cancer. <u>Genome Research</u>, 31:2170-2184. (4 citations)
 - 2. Kim HJ, Osteil P, Humphrey SJ, Cinghu S, Oldfield AJ, Patrick E, Wilkie EE, Peng G, Suo S, **Jothi R**, Tam PPL, Yang P (2020). Transcriptional Network Dynamics During the Progression of Pluripotency Revealed by Integrative Statistical Learning. *Nucleic Acids Research*, 48(4):1828-1842. (8 citations)
 - 3. Oldfield AJ, Henriques T, Burkholder AB, Kumar D, Agirre E, Paulet D, Scruggs, BS, Lavender CA, Yang P, Bennett B, Adelman K, **Jothi R*** (2019). NF-Y controls fidelity of transcription initiation at gene promoters through maintenance of the nucleosome-depleted region. *Nature Communications*, 10(1):3072. (23 citations)
 - Paper of the Year, NIEHS Intramural Research (2019)
 - 4. Yang P[†], Humphrey SJ[†], Cinghu S[†], Pathania R, Oldfield AJ, Kumar D, Perera KD, Yang JYH, James DE, Mann M, **Jothi R*** (2019). Multi-omic profiling reveals dynamics of the phased progression of pluripotency. *Cell Systems*, 8(5):427-445. (>65 citations)
 - Recommended by Faculty of 1000 Biology
 - 5. Jeon K, Kumar D, Conway AE, Park K, **Jothi R**, Jetten AM (2019). GLIS3 Transcriptionally Activates WNT Genes to Promote Differentiation of Human Embryonic Stem Cells into Posterior Neural Progenitors. <u>Stem Cells</u>, 37(2):202-215. (14 citations)
 - 6. Cinghu S[†], Yang P[†], Kosak J, Conway AE, Kumar D, Oldfield AJ, Adelman K, **Jothi R*** (2017). Intragenic enhancers attenuate gene expression. *Molecular Cell*. 68(1):104-117. (>60 citations)
 - Commentary in Nature Reviews in Genetics, doi:10.1038/nrg.2017.90
 - Commentary in Nature Reviews in Molecular Cell Biology, doi:10.1038/nrm.2017.111

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- Paper of the Year, NIEHS Intramural Research (2017)
- Kang HS, Kumar D, Liao G, Lichti-Kaiser K, Gerrish K, Liao X-H, Refetoff S, Jothi R, Jetten AM (2017). GLIS3 is indispensable for TSH/TSHR-dependent thyroid hormone biosynthesis and follicular cell proliferation. <u>The Journal of Clinical Investigation</u>, 127(12):4326-4337 (>30 citations)
 - Paper of the Year, NIEHS Intramural Research (2017)
- 8. Zheng X, Yang P, Lackford B, Bennett B, Wang L, Li H, Miao Y, Fargo D, Jin Y, Williams CJ, **Jothi R**, Hu G (2016). CNOT3-dependent mRNA deadenylation safeguards the pluripotent state. <u>Stem Cell Reports</u>, 7(5):897-910. (>20 citations)
- 9. Yang P, Humphrey SJ, James DE, Yang YH, **Jothi R*** (2016). Positive-unlabeled ensemble learning for kinase substrate prediction from phosphoproteomics data. *Bioinformatics*, 32(2):252-9. (>30 citations)
- Minard AY, Tan S-X, Yang P, Fazakerley DJ, Domanova W, Parker BL, Humphrey SJ, Jothi R, Stockli J, James DE (2016). mTORC1 is a major regulatory node of the FGF21 signaling network. <u>Cell Reports</u>, 17(1):29-36. (>70 citations)
- 11. Yang P, Ellis P, Humphrey SJ, James DE, **Jothi R**, Yang YH (2016). KinasePA: Phosphoproteomics data annotation using hypothesis driven kinase perturbation analysis. *Proteomics*, 16(130:1868-71. (18 citations)
- 12. Yang P, Zheng X, Jayaswal, V, Hu G, Yang YH, **Jothi R*** (2015). Knowledge-based analysis for detecting key signaling events from time-series phosphoproteomics data. <u>PLoS Computational Biology</u> 11(8):e1004403. (23 citations)
- 13. Hoffman NJ, Parker BL, Chaudhuri R, Fisher-Wellman KH, Kleinert M, Humphrey SJ, Yang P, Holliday M, Trefely S, Fazakerley DJ, Stöckli J, Burchfield JG, Jensen TE, **Jothi R**, Kiens B, Wojtaszewski JF, Richter EA, James DE (2015). Global Phosphoproteomic Analysis of Human Skeletal Muscle Reveals a Network of Exercise-Regulated Kinases and AMPK Substrates. *Cell Metabolism*, 22(5):922-35 (>270 citations).
 - Recommended by Faculty of 1000 Biology
- 14. Pathania R, Ramachandran S, Elangovan S, Padia R, Yang P, Cinghu S, Veeranan-Karmegam R, Arjunan P, Gnana-Prakasam JP, Sadanand F, Pei L, Chang CS, Choi JH, Shi H, Manicassamy S, Prasad PD, Sharma S, Ganapathy V, **Jothi R**, Thangaraju M. (2015) DNMT1 is essential for mammary and cancer stem cell maintenance and tumorigenesis. *Nature Communications* 6:6910 (>190 citations).
 - Recommended by Faculty of 1000 Biology
- 15. Oldfield AJ[†], Yang P[†] ([†]Co-first authors), Conway AE, Cinghu S, Freudenberg JM, Yellaboina S, **Jothi R*** (2014). Histone-fold domain protein NF-Y promotes chromatin accessibility for cell type-specific master transcription factors. *Molecular Cell* 55(5):708-722 (>135 citations).
 - Paper of the Year, NIEHS Intramural Research (2014)
- 16. Takeda Y, Kang HS, Freudenberg JM, DeGraff LM, **Jothi R**, Jetten AM (2014) Retinoic Acid-Related Orphan Receptor γ (RORγ): A Novel Participant in the Diurnal Regulation of Hepatic Gluconeogenesis and Insulin Sensitivity. *PLoS Genetics* 10(5), e1004331 (>65 citations).

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- 17. Wang L, Du Y, Ward JM, Shimbo T, Lackford B, Zheng X, Miao Y, Zhou B, Fargo DC, **Jothi R**, Williams CJ, Wade PA, Hu G (2014) An essential role of INO80 in the core pluripotency transcription circuitry. *Cell Stem Cell* 14(5):575-91 (>145 citations).
- 18. Cinghu S[†], Yellaboina S[†] ([†]Co-first authors), Freudenberg JM, Ghosh S, Zheng X, Oldfield AJ, Lackford B, Zaykin DV, Hu G, **Jothi R*** (2014). Integrative framework for identification of key cell identity genes uncovers determinants of ES cell identity and homeostasis. *PNAS* 111(16):E1581-90 (>20 citations).
- 19. Li R, Mav D, Grimm S, **Jothi R**, Shah R, Wade PA (2014). Fine-tuning of epigenetic regulation with respect to promoter CpG content in a cell type-specific manner. <u>Epigenetics</u> 9(5):747-49 (14 citations).
- 20. Lackford B, Yao C, Charles G, Weng L, Zheng X, Choi E-A, Xie X, Wan J, Xing Y, Freudenberg JM, Yang P, **Jothi R**, Hu G, Shi Y (2014) Fip1 regulates mRNA alternative polyadenylation to promote stem cell self-renewal. *EMBO J* 33(8):878-89 (>135 citations).
- 21. Li L[†], Freudenberg JM[†] ([†]Co-first author), Cui K, Dean A, Zhao K, **Jothi R*** (***Co-corresponding author**), Love PE* (2013). Ldb1-nucleated transcription complexes function as primary mediators of global erythroid gene activation. <u>Blood</u> 121(22):4575-85 (>80 citations).
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