# **David Angeles-Albores**

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### **Education**

2013–2018 **Ph.D.**, Biochemistry and Molecular Biophysics,

California Institute of Technology Defense Date: 18 September, 2018 Degree Awarded: 31 October, 2018

2009–2013 B.A., cum laude, Biology

Cornell University

## **Appointments**

11/2019 - Computational Biologist II, eGenesis

01/2019–11/2019 Postdoctoral Associate, Laboratory of Eric J. Alm,

Massachusetts of Technology

11/2018-01/2019 **Postdoctoral Fellow**, Labs of Paul W. Sternberg and Matt Thomson,

California Institute of Technology

#### Research Publications

#### In Preparation, Press or Revision

- Angeles-Albores, D., Min, K.-H., Lee, R. Y., Chan, J., & Sternberg, P. W. (2019). Alaska: Automated rna-seq analysis software for wormbase. *In preparation*.
- Basta, D., Angeles-Albores, M. A., Spero, Ciemniecki, J. A., David, & Newman, D. K. (2019). Heat shock proteases delay cell death during *Pseudomonas aeruginosa* growth arrest. *Proceedings of the National Academy of Sciences.* In press.

#### **Journal Articles**

- Angeles-Albores, D., & Sternberg, P. W. (2018). Using Transcriptomes as Mutant Phenotypes Reveals Functional Regions of a Mediator Subunit in *Caenorhabditis elegans*. *Genetics*, genetics.301133.2018. doi:10.1534/genetics.118.301133
- <sup>†</sup>Angeles-Albores, D., <sup>†</sup>Puckett Robinson, Williams, B. A., Wold, B. J., & Sternberg, P. W. (2018). Reconstructing a metazoan genetic pathway with transcriptome-wide epistasis measurements. *Proceedings of the National Academy of Sciences*, 201712387. doi:10.1073/pnas.1712387115
- <sup>†</sup>**Angeles-Albores**, **D.**, <sup>†</sup>Leighton, D. H. W., Tsou, T., Khaw, T. H., Antoshechkin, I., & Sternberg, P. W. (2017). The *Caenorhabditis elegans* Female-Like State: Decoupling the Transcriptomic Effects of Aging and Sperm Status. *G3 (Bethesda, Md.)* 7(9), 2969–2977. doi:10.1534/g3.117.300080

<sup>†</sup> denotes equal contributions.

- Angeles-Albores, D., N. Lee, R. Y., Chan, J., & Sternberg, P. W. (2016). Tissue enrichment analysis for *C. elegans* genomics. *BMC Bioinformatics*, 17(1), 366. doi:10.1186/s12859-016-1229-9
- Albores-Saavedra, J., Chable-Montero, F., **Angeles-Albores**, **D.**, Schwartz, **A.**, Klimstra, D. S., & Henson, D. E. (2011). Early Gallbladder Carcinoma. *American Journal of Clinical Pathology*, 135(4), 637–642. doi:10.1309/AJCPFRKCFEDLV03Y
- Albores-Saavedra, J., Schwartz, A. M., Henson, D. E., Kostun, L., Hart, A., **Angeles-Albores**, **D.**, & Chablé-Montero, F. (2011). Cutaneous angiosarcoma. Analysis of 434 cases from the surveillance, epidemiology, and end results program, 1973-2007. *Annals of Diagnostic Pathology*, 15(2), 93-97. doi:10.1016/j.anndiagpath.2010.07.012

#### $\mu$ Publications

Angeles-Albores, D., N. Lee, R. Y., Chan, J., & Sternberg, P. W. (2018). Two new functions in the WormBase Enrichment Suite. *microPublication Biology*. doi:10.17912/W25Q2N

### Scientific Talks

2019 Probabilistic Modeling in Genomics

Genetics is an active learning algorithm for causal reconstruction of biological networks

Hanna H Gray Semifinalist Symposium

Phenotypes, epistasis, and probability theory

ASBMB Special Symposium: Evolution and Core Processes in Gene Expression Transcriptomes as phenotypes

2018 Bay Area Worm Meeting

Allelic series analyses using transcriptomic phenotypes

2017 21<sup>st</sup> C. elegans International Meeting, WormBase: Tools, Content and Community Annotation, Workshop

Gene Set Analysis tool for Gene Ontology (GO), Phenotype, and Tissue Enrichment

Annual Departmental Retreat, California Institute of Technology

Genome-wide, unbiased experimental genetics

Biochemistry and Molecular Biophysics Seminar Series, California Institute of Technology

Transcriptomic Genetics: A new way to use RNA-sequencing data

**Center for Environmental Microbial Interactions**, California Institute of Technology Genome-wide unbiased experimental genetics

Annual Biochemistry and Molecular Biophysics Program Retreat, California Institute of Technology

Reconstruction of a genetic pathway using whole-organism expression profiles

Graduate Biology Seminar, California Institute of Technology

Transcriptome-wide epistasis in mRNA expression profiles

#### **Awards**

- 2019 HHMI Hanna Gray Fellow Finalist
- Florence C. Rose and S. Meryl Rose Endowed Scholarship for attendance to the Embryology course at the Marine Biological Laboratory
- 2014 Amgen Graduate Student Fellowship
- 2012 EXROP Capstone Award

# **Awards (continued)**

2011 EXROP HHMI Summer Fellowship

# **Scientific Courses**

2015 Embryology, Marine Biological Laboratory at Woods Hole

# **Teaching and Mentoring Experience**

## **Teaching**

2017–2018	Systems Genetics, Teaching Assistant, California Institute of Technology
2016	Introduction to Biology, Teaching Assistant, California Institute of Technology
2015	Order of Magnitude Biology, Teaching Assistant, California Institute of Technology
2014	Advanced Experimental Methods in Bioorganic Chemistry, Teaching Assistant, California Institute of Technology

## Mentoring

2016–2019	<b>Kyung Hoi Min</b> , Caltech undergraduate, experimental and computational student, <i>California Institute of Technology</i>
Summer 2018	<b>Vladimir Molchanov</b> , Saint Petersburg Bioinformatics Institute undergraduate, experimental student, <i>California Institute of Technology</i>
Summer 2015	<b>Tiffany Tsou</b> , UCSB undergraduate, experimental student, <i>California Institute of Technology</i>
2014–2015	<b>Isabelle Phinney</b> , Polytechnic School, computational student, <i>California Institute of Technology</i>

## Outreach

2017	Speaker, What is Clubes de Ciencias Mexico?, SACNAS
2016	Co-instructor, De Planarias y Derivas, Clubes de Ciencia México, Guanajuato
	Co-instructor, De Genes y Animales, Clubes de Ciencia México, Ensenada
2015	Student selection committee, Clubes de Ciencia México
2014	Guest instructor, Biología a través de los números, Clubes de Ciencia México, Ensenada

# **Scientific Societies**

2014–Present	Genetics Society of America
2015–Present	Society for Developmental Biology

## References

### Professor Paul W. Sternberg

California Institute of Technology, MC 156-29, Pasadena CA 91125

pws@caltech.edu

### **Assistant Professor Matthew Thomson**

California Institute of Technology, 216–76, Pasadena CA 91125

### Professor Dianne K. Newman

California Institute of Technology, MC 147-75, Pasadena CA 91125

### **Assistant Research Professor** Erich M. Schwarz

Cornell University, Biotechnology 351, Cornell University, Ithaca, NY 14853-2703