

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**
One of two founding members of the computational biology unit.
Introduced ATAC-seq, scRNA-seq and scATAC-seq to eGenesis.
Created an atlas of the pig kidney cortex
Used the above methods to generate a compendium of promoters that stably express genes ubiquitously or with high tissue-specificity at desired levels with low burstiness
- 7/2020 – **Visiting Scholar, Laboratory of Ilya Ruvinsky,
Northwestern University**
I collaborate with the Ruvinsky lab on RNA-seq analysis of *C. elegans* biology.
- 01/2019–11/2019 **Postdoctoral Associate, Laboratory of Eric J. Alm,
Massachusetts Institute of Technology**
I performed research on microbiome analysis. I terminated my postdoctoral research prematurely as a result of a personal loss, unrelated to my work at MIT.
- 11/2018–01/2019 **Postdoctoral Fellow, Labs of Paul W. Sternberg and Matt Thomson,
California Institute of Technology**
I used this short period to finish some projects, and developed an unpublished algorithm for analysis of CRISPR screens in mammalian cells, using transcriptomes as phenotypes for genetic analysis.

Research Publications

[†] denotes equal contributions.

Journal Articles

- 1 Basta, D. W., Angeles-Albores, D., Spero, M. A., Ciemniecki, J. A., & Newman, D. K. (2020). Heat-shock proteases promote survival of *Pseudomonas aeruginosa* during growth arrest. *Proceedings of the National Academy of Sciences of the United States of America*. doi:10.1073/pnas.1912082117

- 2 Duncan, E. M., Nowotarski, S. H., Guerrero-Hernández, C., Ross, E. J., D’Orazio, J. A., Clubes de Ciencia México, W. f. D. B., ... Alvarado, A. S. (2020). A new species of planarian flatworm from Mexico: *Girardia guanajuatensis*. *bioRxiv*. doi:10.1101/2020.07.01.183442. eprint: <https://www.biorxiv.org/content/early/2020/07/02/2020.07.01.183442.full.pdf>
- 3 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
- 4 †Angeles-Albores, D., †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
- 5 †Angeles-Albores, D., †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
- 6 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
- 7 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
- 8 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

μPublications

- 1 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 **21st 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

Scientific Talks (continued)

- Center for Environmental Microbial Interactions, California Institute of Technology
Genome-wide unbiased experimental genetics
- 2016 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

- 2020 eGenesis Leadership Award
- 2019 HHMI Hanna Gray Fellow Finalist
- 2015 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
- 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 Kyung Hoi Min, Caltech undergraduate, experimental and computational student, *California Institute of Technology*
- Summer 2018 Vladimir Molchanov, Saint Petersburg Bioinformatics Institute undergraduate, experimental student, *California Institute of Technology*
- Summer 2015 Tiffany Tsou, UCSB undergraduate, experimental student, *California Institute of Technology*
- 2014–2015 Isabelle Phinney, Polytechnic School, computational student, *California Institute of Technology*

Outreach

- 2020 **Invited speaker**, *Clubeando en casa, el Podcast* Clubes de Ciencia México
- 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

Toby Bloom

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Professor Eric J Alm

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Professor Paul W. Sternberg

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Assistant Research Professor

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