

Kam D. Dahlquist, Ph.D.
Professor of Biology
Loyola Marymount University

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EDUCATION

Ph.D.	University of California, Santa Cruz Molecular, Cellular, and Developmental Biology Program <i>Advisor:</i> Joseph D. Puglisi, Ph.D. <i>Committee:</i> Harry F. Noller, Ph.D., Manuel Ares, Jr., Ph.D. <i>Thesis:</i> <i>Interaction of Translation Initiation Factor IF1</i> <i>with the E. coli Ribosomal A Site</i>	March 2000
B.A.	Pomona College, Claremont, California Biology, <i>cum laude</i>	May 1993
	University College, Oxford University, Oxford, England Study Abroad Program <i>Specialized tutorial in Philosophy of Science</i>	Fall 1991

POSITIONS HELD

Chair , Department of Biology, Loyola Marymount University, Los Angeles, California	2020–2023
Professor , Department of Biology, LMU	2017–present
Affiliate Faculty Bioethics Institute, LMU	2013–2015
William F. McLaughlin Chair of Biology , LMU	2010–2012
Associate Professor , Department of Biology, LMU	2009–2017
Assistant Professor , Department of Biology, LMU	2005–2009
Assistant Professor Department of Biology, Vassar College, Poughkeepsie, New York	2003–2005
Postdoctoral Fellow Gladstone Institute of Cardiovascular Disease, University of California, San Francisco	2000–2003
Adjunct Lecturer Department of Biology, Santa Clara University, Santa Clara, California	Spring 2000
Visiting Researcher Department of Structural Biology, Stanford University, Stanford, California	1997–2000
Research Assistant Department of Biology, University of California, Santa Cruz, California	1994–1997

GRANTS, FELLOWSHIPS, HONORS, AWARDS

- Kadner-Pitts Research Grant** 2025–2026
\$15,000, *Next-generation Sequencing Methods for Understanding Gene Regulatory Networks in Yeast*
- NIH National Institute of General Medical Sciences R15** 2019–2022
\$323,775, *Aim32p: a Novel Multi-faceted Protein in Mitochondrial Biogenesis*
Principal Investigator: Deepa Dabir; *Key Personnel:* **Kam D. Dahlquist**, bioinformatics consultant for Specific Aim 3, *Characterizing the Aim32p interactome*
- ASBMB Travel Grant** 2020
\$500, to present at the American Society for Biochemistry and Molecular Biology Annual Meeting, April 4–8, 2020, San Diego California (award rescinded when meeting was cancelled due to the COVID-19 pandemic)
- Kadner-Pitts Research Grant** 2017–2018
\$11,600, *Experimental, Mathematical Modeling, and Data Visualization Methods to Investigate the Properties of the Gene Regulatory Network Controlling the Cold Shock Response in Budding Yeast*
- ASBMB Travel Grant** 2016
\$500, to present at the American Society for Biochemistry and Molecular Biology Annual Meeting, April 2–5, 2016, San Diego California
- National Institute for Mathematical and Biological Synthesis (NIMBioS)** 2015–2017
Working Group, *Unpacking the Black Box: Teaching Quantitative Biology*, Invitation-only, collaborative group with face-to-face meetings held at the University of Tennessee, Knoxville and an online collaboration between meetings
- Elizabeth and Michael Rudinica Endowed Prize for Student-Faculty Research** 2015
Seaver College of Science and Engineering, Loyola Marymount University
- Kadner-Pitts Research Grant** 2015–2016
Department of Biology, Loyola Marymount University
\$13,400, *Extending and Refining the Mathematical Model of the Gene Regulatory Network Controlling the Cold Shock Response in Budding Yeast*
- ASBMB Thematic Best Poster Award in Systems Biology** 2012
\$500, for poster presented at the American Society for Biochemistry and Molecular Biology Annual Meeting, April 20–24, 2012, San Diego, California
- ASBMB and NSF Travel Grant** 2012
\$2,250, to present at the American Society for Biochemistry and Molecular Biology Annual Meeting, April 20–24, 2012, San Diego California
- NSF-DMS Mathematical Biology, MCB Genes and Genome Systems** 2009–2015
\$246,123, *Collaborative Research and RUI: Stochastic Dynamic Network Models of Gene Regulation under Environmental Stress*
Principal Investigator: **Kam D. Dahlquist**; *Co-Principal Investigator:* Ben G. Fitzpatrick
- Loyola Marymount University Center for Teaching Excellence Travel Grant, \$740** 2009
- NSF-UBM (Interdisciplinary Training for Undergraduates in Biological and Mathematical Sciences)** 2007–2010
\$240,000, *Analysis of Stress in Biological Systems*
Principal Investigator: Ben G. Fitzpatrick; *Co-Principal Investigators:* Wendy J. Binder, Erika Camacho, **Kam D. Dahlquist**, Gary A. Kuleck; *Faculty Associates:* Philippa M. Drennan, Martin G. Ramirez, *Interdisciplinary Research Project* with Ben G. Fitzpatrick entitled, *Modeling Gene Expression Networks in Saccharomyces cerevisiae*
- W.M. Keck Foundation** 2007–2010
\$300,000, *Equipment for the Molecular Analysis and Imaging Laboratory*
Principal Investigator: Gary A. Kuleck; *Co-Principal Investigators:* **Kam D. Dahlquist**, David Moffet, Martin G. Ramirez, Carl R. Urbinati

Kadner-Pitts Research Grant	2007–2008
Department of Biology, Loyola Marymount University	
\$10,000, <i>Mapping Gene Regulatory Networks in Yeast using DNA Microarrays, Mathematical Modeling, and GenMAPP</i>	
Merck-AAAS Undergraduate Science Research Program	2006–2008
\$120,000 (\$60,000 plus \$60,000 matching funds from Loyola Marymount University)	
<i>Chemical and Biological Aspects of Pollution in the Ballona Wetlands</i>	
<i>Principal Investigator:</i> M. Catherine McElwain; <i>Director and Co-Principal Investigator:</i> Kam D. Dahlquist ; <i>Co-Principal Investigators:</i> Rachel Adams, Lambert Doezeza, John Dorsey, Philippa M. Drennan, Gary A. Kuleck, Jim Landry, Jeremy McCallum, David Moffet, Martin G. Ramirez, James Roe, and Carl R. Urbinati, <i>Interdisciplinary Research Project</i> with David Moffet and Carl R. Urbinati entitled <i>Identifying Soil Bacteria and Biochemical Pathways in the Ballona Wetlands for the Bioremediation of Organic Pollutants</i>	
Academic Technology Grant, Loyola Marymount University	2007
\$4,000, <i>Introducing DNA Microarray Technology in the New Laboratory Course, Biology 478: Molecular Biology of the Genome</i>	
Dartmouth Faculty Summer Institute Travel Award and Stipend	August 2006
<i>ELSI Reunion and Conference, Dartmouth University, Hanover, New Hampshire</i>	
Summer Research Grant for New Faculty, Loyola Marymount University	2006
\$4,000, <i>The Transcriptional and Proteomic Response to Cold Shock and Recovery in Saccharomyces cerevisiae</i>	
Merck-AAAS Undergraduate Science Research Program	2005
<i>Principal Investigator:</i> Richard B. Hemmes, Department of Biology, Vassar College	
<i>Interdisciplinary Research Project</i> with Eric S. Eberhardt, Department of Chemistry, Vassar College entitled <i>Examining the Molecular Details of Oxidative Stress from the Genome to the Proteome</i> [I declined my share of the funding upon my move to Loyola Marymount University]	
Mellon Faculty Conversations Award, Vassar College	2004–2005
\$2,000, <i>Effective Grading: A Tool for Learning and Assessment</i>	
Dartmouth Faculty Summer Institute Travel Award and Stipend	July 2004
<i>Ethical, Legal, and Social Implications of the Human Genome Project</i>	
Dartmouth University, Hanover, New Hampshire	
Sigma Xi, Full Membership	2004
Associate Membership	1992
Carolyn Grant Endowment for Embodied Learning, Vassar College	2004
\$2,000, Sponsored a visit by Jean Couch to lead workshops on <i>Balanced Posture</i> for <i>Introduction to Biological Thought: The Human Genome</i> and the campus community	
GAANN Fellowship, U.C. Santa Cruz	1995–1997
Phi Beta Kappa	1993
Vaile Prize in Biology, Pomona College	1993
Senior Service Award, Pomona College	1993
Eda May Haskell Library Prize, Pomona College	1993
Best Seminar in Plant or Microbial Biology	1992
West Coast Biological Sciences Undergraduate Research Conference	

RESEARCH EXPERIENCE

Professor	2017–present
Department of Biology, Loyola Marymount University, Los Angeles, California	
<i>Current Research Projects:</i>	
<ul style="list-style-type: none"> Determining the gene regulatory network controlling the global transcriptional response of budding yeast, <i>Saccharomyces cerevisiae</i>, to cold shock and recovery (2003–present) Modeling the dynamics of this gene regulatory network through the development of the 	

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- GRNmap software (2006–present)
 - Visualizing the results of the dynamical network model through the development of the GRNsight software (2014–present)
 - Detection of SNP genotypes in humans that confer lactase persistence (2019–present)
Research advisor for 43 undergraduates and 1 Master's level student from the 2017–2018 academic year to the present.
 - Associate Professor** 2009–2017
Department of Biology, Loyola Marymount University, Los Angeles, California
 - In addition to the projects above, creation of GenMAPP-compatible Gene Databases using the XMLPipeDB software suite for the analysis of published microarray data (2006–2016).
Research advisor for 42 undergraduates from 2009–2017.
 - Assistant Professor** 2005–2009
Department of Biology, Loyola Marymount University, Los Angeles, California
 - In addition to the projects noted above, identifying soil bacteria and biochemical pathways in the Ballona Wetlands for the bioremediation of organic pollutants (2006–2008)
Research advisor for 8 undergraduates and 1 Master's level student from 2005–2008.
 - Assistant Professor** 2003–2005
Department of Biology, Vassar College, Poughkeepsie, New York
 - Transcriptional and Proteomic Response of *Saccharomyces cerevisiae* to Cold Shock and Recovery
 - Creation of MAPPs, Gene Databases, and Documentation for GenMAPP software
Research advisor for a total of 6 undergraduate students from 2003–2005.
 - Postdoctoral Fellow** 2000–2003
Gladstone Institute of Cardiovascular Disease, University of California, San Francisco
Advisor: Bruce R. Conklin, M.D.; *Topic:* Pathway-based analysis of microarray data; project management, design, testing, and documentation of the GenMAPP software
 - Visiting Researcher** 1997–2000
Department of Structural Biology, Stanford University, Stanford, California
Dissertation Advisor: Joseph D. Puglisi, Ph.D.
 - Research Assistant** 1994–1997
Department of Biology, University of California, Santa Cruz
Dissertation Advisor: Joseph D. Puglisi, Ph.D.
 - Rotation Student** 1993–1994
Department of Biology, University of California, Santa Cruz
Advisor: Jack K. Okamuro, Ph.D., *Topic:* Identification of additional members of the *Apetala2* family in *Arabidopsis thaliana*; *Advisor:* Jerry F. Feldman, Ph.D., *Topic:* Mapping of the *Period2* locus in *Neurospora crassa*
 - Undergraduate Researcher** Summer 1993
Howard Hughes Summer Institute, University of California, Santa Cruz
Advisor: Jane Silverthorne, Ph.D., *Topic:* Characterization of phytochrome genes in *Ginkgo biloba*
 - Undergraduate Researcher** 1991–1992
Department of Biology, Pomona College, Claremont, California
Advisor: David W. Becker, Ph.D., *Topic:* Effect of heat stress on photosynthesis in a high-temperature strain of the green alga, *Chlorella pyrenoidosa*
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TEACHING EXPERIENCE

College Level

- Department of Biology, Loyola Marymount University** 2005–present
Biology 112: General Biology II Laboratory (2021)

- Freshman lab required for biology and environmental science majors, post-baccalaureate students, and pre-medical students from other majors
 - Team-taught with Dr. Carl Urbinati, Professor Tatiana Kuzmenko, and twelve undergraduate Teaching Assistants
 - Successfully adapted a bioinformatics lab curriculum for online learning in Spring 2021
- Biology 190: Freshman Biology Seminar* (2023–2024)
- 0-credit orientation course required of all biology majors in Fall of their first year
 - Focus on study skills, academic advising, and student research
 - Concomitant with organizing Biology Department Seminars for the academic year
- Biology 201: Cell Function* (2005–2011, 2013–2015, 2017, 2019, 2021–2024)
- Sophomore-level requirement in four-semester lower division curriculum for biology, biochemistry, and environmental science majors
 - Course coordinator 2009–2011, 2013–2015, 2017, 2019, 2023
- Biology 275: Human Genetics* (2006)
- Fulfills University core requirement for non-science majors
- Biology 367/Computer Science 367: Biological Databases* (2008–2010, 2013, 2015, 2017, 2019, 2024)
- Cross-listed and team taught with John David N. Dionisio, Ph.D., Department of Electrical Engineering and Computer Science (through 2017)
 - Interdisciplinary student teams create yeast gene expression databases for incorporation into the GRNsight open source software for visualizing models of gene regulatory networks
 - Most recent course website:
https://xmlpipedb.cs.lmu.edu/biodb/spring2024/index.php/Main_Page
- Biology 368: Bioinformatics Laboratory* (2008, 2010–2011, 2014, 2016, 2020)
- Projects include sequence and structural analysis of the gp120 protein of HIV, spike protein of SARS-CoV-2, and analysis of DNA microarray experiments
 - Most recent course website: <http://www.openwetware.org/wiki/BIOL368/F20>
- Biology 388/Mathematics 388: Biomathematical Modeling* (2011, 2013, 2015, 2017, 2019, 2025)
- Cross-listed and team taught with Ben G. Fitzpatrick, Ph.D., Department of Mathematics
 - Students create mathematical models of nitrogen metabolism and use the GRNmap and GRNsight software for modeling gene regulatory networks in budding yeast
 - Course website: <https://openwetware.org/wiki/BIOL388/S25>
- Biology 439: Molecular Biology Applications* (2006–2007)
- Intensive laboratory course in molecular biology
 - Students performed semi-independent cloning project based on my dissertation research
- Biology 478: Molecular Biology of the Genome* (2007–2010, 2013–2020, 2025)
- Subject of 2007 LMU Academic Technology Grant
 - Intensive laboratory course in molecular biology; student-performed DNA microarray experiments contribute to dataset deposited in NCBI Gene Expression Omnibus database
- Biology 479: Molecular Mechanisms of Disease* (2024)
- Students investigated a gene-disease relationship through reviewing the literature and using bioinformatics tools to understand the properties of the gene and protein
- Biology 498/Computer Science 698: Special Studies in Bioinformatics* (2006)
- Master's-level course cross-listed with Computer Science, team taught with John David N. Dionisio, Ph.D., Department of Electrical Engineering and Computer Science
 - Project-based course initiated development of the XMLPipeDB software suite using open source tools and the SourceForge development environment
- Biology 585: Issues in Biotechnology* (2007, 2014, 2016, 2018, 2022–2023)
- Seminar and capstone experience for biology majors

- Read, present, and discuss articles from the primary biotechnology literature, followed by discussion of the ethical, legal, and social implications

Foundations First Year Seminar 1000: Sleep! Your Hidden Superpower (2023)

- University core requirement for first year students, focused on writing

Honors 240: On the Nature of Things (2009)

- University core requirement for students in the Honors Program
- An examination of the history, philosophy, and nature of scientific discovery, theory, and practice, focusing on recent advances in biotechnology and genomics, epistemology, and genetic determinism

Honors 2200: Nature of Science, Technology, and Mathematics (2024)

- University core requirement for students in the Honors Program
- Focus on human genetics, the human genome, biotechnology, and the ethical, legal, and social implications

Delivered Guest Lecture in the following courses

- BIOL 216: Medical Microbiology (Fall 2023)
- BIOL 475: Evolution (Fall 2023)
- FFYS 1000: Principles of Scientific Reasoning (Fall 2023)
- BIOE 630: Genetic Medicine (Fall 2009)
- BIOL 114: Biology for Engineers (Spring 2009)
- PHIL 666: Philosophy of Science (Fall 2008)
- CMSI 686: Database Systems (Spring 2007)
- MATH 298: Biomathematics (Spring 2007)
- CMSI 598/698: Open Source Software Development Workshop (Summer 2006)
- MGMT 498: Technology Ventures (Spring 2006)
- CMSI 486: Introduction to Database Systems (Fall 2005)

Department of Biology, Vassar College (Assistant Professor)

2003–2005

Introduction to Biological Thought: The Human Genome

- Students used MAPPFinder to analyze a publicly available cancer microarray dataset
- Students learned scientific writing step-by-step, culminating in a final draft of a review of a primary research article about a gene involved in cancer

Principles of Genetics

- Students used GenMAPP to draw a biochemical pathway and analyze microarray data related to their “wet” lab work
- Emphasized the “practical” aspects of successful scientific research through special exercise in teamwork
- Genetics and Society presentations, papers, and discussions taught students about the ethical implications of genetics research

Bioinformatics

- Project-based computer laboratory using GenMAPP, MAPPFinder, and other bioinformatics software
- Students designed web sites to showcase their work
- Emphasized presentation skills and reading primary scientific literature

Department of Biology, Santa Clara University (Adjunct Lecturer)

Spring 2000

Molecular Biology

- Taught lecture and lab to 20 upper-division, biology majors; was solely responsible for course content
- Developed lab exercise based on thesis research where students cloned different mutations in 16S rRNA into an *E. coli* expression vector and analyzed the phenotype of the mutant cells
- Developed bioinformatics lab exercise based on tools publicly available on the web

Department of Biology, Stanford University (Course Assistant)	Winter 1998
<i>Cell Biology</i> , led discussion of research articles	
Department of Biology, U.C. Santa Cruz (Teaching Assistant)	Fall 1994
<i>Concepts in Biology</i> , lectured when professor was out of town	
Howard Hughes Summer Institute, U.C. Santa Cruz (Teaching Assistant)	Summer 1994
<i>Molecular and Cell Biology Laboratory</i> , supervised semi-independent research projects on the cloning of <i>frequency</i> homologues in different species of fungi	
K-12	
Herbert Hoover Middle School and U.C. San Francisco (Scientist Volunteer)	2001–2002
<i>Science and Health Education Partnership Triad Science Club</i>	
<ul style="list-style-type: none"> Developed and led hands-on activities, including gel electrophoresis 	
Mission Hill Junior High School, U.C. Santa Cruz (Elective Teacher)	Fall 1994
<i>Project SAME: Science and Math Equity</i>	
<ul style="list-style-type: none"> Taught a girl-only elective class on building simple machines with the Lego-Logo system 	

PUBLICATIONS

Peer-reviewed Research (*indicates undergraduate co-author)

- Eaton, C.D., Callendar, H.L., **Dahlquist, K.D.**, LaMar, M.D., Ledder, G., Schugart, R.C. (2019) A “Rule of Five” Framework for Models and Modeling to Unify Mathematicians and Biologists and Improve Student Learning, *PRIMUS: Problems, Resources, and Issues in Mathematics Undergraduate Studies*, published online 12 March 2019. DOI: 10.1080/10511970.2018.1489318.
- Dahlquist, K.D.**, Dionisio, J.D.N., Libeskand-Hadas, R, Bargagliotti, A.E. (2018) Breaking Boundaries in Computing in Undergraduate Courses *Journal of Research in STEM Education* **4**: 81-100.
- Dahlquist, K.D.**, Dionisio, J.D.N., Fitzpatrick, B.G., Anguiano, N.A.*, Varshneya, A.*, Southwick, B.J.*, Samdarshi, M.* (2016) GRNsight: a web application and service for visualizing models of small- to medium-scale gene regulatory networks. *PeerJ Computer Science* **2**:e85. DOI: 10.7717/peerj-cs.85.
- Dahlquist, K.D.**, Fitzpatrick, B.G., Camacho, E.T., Entzminger, S.D.*, and Wanner, N.C.* (2015) Parameter Estimation for Gene Regulatory Networks from Microarray Data: Cold Shock Response in *Saccharomyces cerevisiae*. *Bulletin of Mathematical Biology*, **77**: 1457-1492, published online September 29, 2015. DOI: 10.1007/s11538-015-0092-6.
- Demir, E., Cary, M.P., Paley, S., Fukuda, K., Lemer, C., Vastrik, I., Wu, G., D’Eustachio, P., Schaefer, C., Luciano, J., Schacherer, F., Martinez-Flores, I., Hu, Z., Jimenez-Jacinto, V., Joshi-Tope, G., Kandasamy, K., Lopez-Fuentes, A.C., Mi, H., Pichler, E., Rodchenkov, I., Splendiani, A., Tkachev, S., Zucker, J., Gopinath, G., Rajasimha, H., Ramakrishnan, R., Shah, I., Syed, M., Anwar, N., Babur, O., Blinov, M., Brauner, E., Corwin, D., Donaldson, S., Gibbons, F., Goldberg, R., Hornbeck, P., Luna, A., Murray-Rust, P., Neumann, E., Reubenacker, O., Samwald, M., van Iersel, M., Wimalaratne, S., Allen, K., Braun, B., Whirl-Carrillo, M., Cheung, K.H., **Dahlquist, K.**, Finney, A., Gillespie, M., Glass, E., Gong, L., Haw, R., Honig, M., Hubaut, O., Kane, D., Krupa, S., Kutmon, M., Leonard, J., Marks, D., Merberg, D., Petri, V., Pico, A., Ravenscroft, D., Ren, L., Shah, N., Sunshine, M., Tang, R., Whaley, R., Letovsky, S., Buetow, K.H., Rzhetsky, A., Schachter, V., Sobral, B.S., Dogrusoz, U., McWeeney, S., Aladjem, M., Birney, E., Collado-Vides, J., Goto, S., Hucka, M., Le Novère, N., Maltsev, N., Pandey, A., Thomas, P., Wingender, E., Karp, P.D., Sander, C., and Bader, G.D. (2010) The BioPAX Community Standard for Pathway Data Sharing. *Nature Biotechnology* **28**: 935-942. DOI: 10.1038/nbt.1666
- Ogando, D.G., **Dahlquist, K.D.**, Alizadeh, M., Kunchithapautham, K., Li, J., Yu, N., LaVail, M.M., Rohrer, B., Vollrath, D., and Danciger, M. (2010) Candidate Genes for Chromosomes 6 and 10 Quantitative Trait Loci for Age-related Retinal Degeneration in Mice. *Molecular Vision* **16**: 1004-1018.
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- Dionisio, J.D.N. and **Dahlquist, K.D.** (2008) Improving the Computer Science in Bioinformatics Through Open Source Pedagogy *ACM SIGCSE Bulletin* **40**: 115-119. DOI: 10.1145/1383602.1383648.
- Salomonis, N., Hanspers, K., Zambon, A.C., Vranizan, K., Lawlor, S.C., **Dahlquist, K.D.**, Doniger, S.W., Stuart, J., Conklin, B.R., & Pico, A.R. (2007) GenMAPP 2: New Features and Resources for Pathway Analysis. *BMC Bioinformatics* **8**: 217. DOI: 10.1186/1471-2105-8-217.
- Segal, M.R., **Dahlquist, K.D.**, & Conklin, B.R. (2003) Regression Approaches for Microarray Data Analysis. *Journal of Computational Biology* **10**: 961-980. DOI: 10.1089/106652703322756177.
- Doniger, S.W., Salomonis, N., **Dahlquist, K.D.**, Vranizan, K., Lawlor, S.C., & Conklin, B.R. (2003) MAPPFinder: Using Gene Ontology and GenMAPP to Create a Global Gene-Expression Profile from Microarray Data. *Genome Biology* **4**: R7. DOI: 10.1186/gb-2003-4-1-r7.
- Dahlquist, K.D.**, Salomonis, N., Vranizan, K., Lawlor, S.C., & Conklin, B.R. (2002) GenMAPP, A New Tool for Viewing and Analyzing Microarray Data on Biological Pathways. *Nature Genetics* **31**: 19-20. DOI: 10.1038/ng0502-19.
- Dahlquist, K.D.** & Puglisi, J.D. (2000) Interaction of Translation Initiation Factor IF1 with the *E. coli* Ribosomal A site. *Journal of Molecular Biology* **299**: 1-15. DOI: 10.1006/jmbi.2000.3672.
- Recht, M.I., Douthewaite, S., **Dahlquist, K.D.**, & Puglisi, J.D. (1999) Effect of Mutations in the A site of 16S rRNA on Aminoglycoside Antibiotic-Ribosome Interaction. *Journal of Molecular Biology* **286**: 33-43. DOI: 10.1006/jmbi.1998.2446.
- Recht, M.I., Fourmy, D., Blanchard, S.C., **Dahlquist, K.D.**, & Puglisi, J.D. (1996) RNA Sequence Determinants for Aminoglycoside Binding to an A-site rRNA Model Oligonucleotide. *Journal of Molecular Biology* **262**: 421-436. DOI: 10.1006/jmbi.1996.0526.

Reviews, Book Chapters, Conference Proceedings

- Dahlquist, K.D.**, editor (2010) Proceedings of the 11th Annual Bioinformatics Open Source Conference (BOSC) 2010. *BMC Bioinformatics* **11**(Suppl 12): S1-S13.
- Dahlquist, K.D.** (2004) Using GenMAPP and MAPPFinder to View Microarray Data on Biological Pathways and Identify Global Trends in the Data. In *Current Protocols in Bioinformatics* (Baxevanis, A.D., Davison, D.B., Page, R., Stein, L., Stormo, G., eds.), John Wiley & Sons, Inc., New York, N.Y., pp. 7.5.1-7.5.26.
- Puglisi, J.D., Blanchard, S.C., **Dahlquist, K.D.**, Eason, R.G., Fourmy, D., Lynch, S.R., Recht, M.I., & Yoshizawa, S. (1999) Aminoglycoside Antibiotics and Decoding. In *The Ribosome: Structure, Function, Antibiotics, and Cellular Interactions* (Garrett, R.A., Douthewaite, S.R., Liljas, A., Matheson, A.T., Moore, P.B., & Noller, H.F., eds.), pp. 419-429. ASM Press, Washington, D.C.
- Dahlquist, K.** & Puglisi, J.D. (1995) Investigating the Structure and Function of Translation Initiation Factor 1. *Nucleic Acids Symposium Series* **33**: 170-171.

Preprints

- Dahlquist, K.D.**, Aikens, M.L., Dauer, J.T., Donovan, S.S., Eaton, C.D., Highlander, H.C., Jenkins, K.P., Jungck, J.R., LaMar, M.D., Ledder, G., Mayes, R.L., Schugart, R.C. (2017) An Invitation to Modeling: Building a Community with Shared Explicit Practices, *PeerJ Preprints* 5:e3215v1 <https://doi.org/10.7287/peerj.preprints.3215v1>.

Software, Databases, and Datasets (*indicates undergraduate co-author)

NCBI Gene Expression Omnibus Series GSE83656

Dahlquist K.D., Abdulla, H.*, Arnell, A.J.*, Arsan, C.*, Baker, J.M.*, Carson, R.M.*, Citti, W.T.*, De Las Casas, S.E.*, Ellis, L.G.*, Entzminger, K.C.*, Entzminger, S.D.*, Fitzpatrick, B.G., Flores, S.P.*, Harmon, N.S.*, Hennessy, K.P.*, Herman, A.F.*, Hong, M.V.*, King, H.L.*, Kubeck, L.N.*, La-Anyane, O.M.*, Land, D.L.*, Leon Guerrero, M.J.*, Liu, E.M.*, Luu, M.D.*, McGee, K.P.*,

Mejia, M.R.*, Melone, S.N.*, Pepe, N.T.*, Rodriguez, K.R.*, Rohacz, N.A.*, Rovetti, R.J., Sakhon, O.S.*, Sampana, J.T.*, Sherbina, K.*, Terada, L.H.*, Vega, A.J.*, Wavrin, A.J.*, Wyllie, K.W.*, Zapata, B.B.* (2016) Global transcriptional response of wild type and transcription factor deletion strains of *Saccharomyces cerevisiae* to the environmental stress of cold shock and subsequent recovery. Dataset of 137 DNA microarray hybridizations performed by undergraduate students as part of independent research and the course Biology 478: Molecular Biology of the Genome from 2006 to 2016. A manuscript describing this dataset is in preparation.

GRNmap (Gene Regulatory Network Modeling and Parameter Estimation)

Co-Principal Investigator with Ben G. Fitzpatrick and in collaboration with John David N. Dionisio and undergraduate research students, 2014–present

Availability (Open Source BSD license): <http://kdahlquist.github.io/GRNmap/index.html>,
<https://github.com/kdahlquist/GRNmap/>

GRNsight (Web Application for Visualizing Small-Scale Models of Gene Regulatory and Protein-Protein Interaction Networks)

Co-Principal Investigator with John David N. Dionisio and Ben G. Fitzpatrick and in collaboration with undergraduate research students, 2014–present

Availability (Open Source BSD license): <http://dondi.github.io/GRNsight/index.html>,
<https://github.com/dondi/GRNsight>

XMLPipeDB (A Reusable, Open Source Tool Chain for Building Relational Databases from XML Sources) and Gene Databases for 19 species:

Arabidopsis thaliana, 2007, 2009; *Bordetella pertussis*, 2015; *Burkholderia cenocepacia*, 2015; *Chlamydia trachomatis*, 2013; *Escherichia coli* K12, 2006, 2009; *Helicobacter pylori*, 2011; *Leishmania infantum*, 2014; *Leishmania major*, 2014; *Mycobacterium smegmatis*, 2011; *Mycobacterium tuberculosis* H37Rv, 2010; *Plasmodium falciparum*, 2009; *Pseudomonas aeruginosa* PAO1, 2010; *Salmonella typhimurium*, 2011; *Shewanella oneidensis*, 2015; *Shigella flexneri*, 2015; *Sinorhizobium meliloti*, 2013; *Staphylococcus aureus* MRSA 252, 2010, *Streptococcus pneumoniae*, 2013; and *Vibrio cholerae*, 2009, 2010, 2016. Co-Principal Investigator with John David N. Dionisio, and in collaboration with 16 undergraduate research students, 1 Master's student, and students in the Biology/Computer Science 367: Biological Databases courses, 2006–present; Availability (Open Source LGPL license): <http://xmllpipedb.cs.lmu.edu>,
<https://github.com/lmu-bioinformatics/xmllpipedb>

GenMAPP (Gene Map Annotator and Pathway Profiler) 1.0 and 2.0

Project Manager, 2000–2003

Metabolic Pathway MAPP Archive for *Saccharomyces cerevisiae*, 2005; for *E. coli* K12, 2008

Availability: <http://www.GenMAPP.org>, <https://github.com/GenMAPP/GenMAPP>

PRESENTATIONS

Invited Talks

Gladstone Institutes Trainee to Tenure Track Series

San Francisco, California (virtual), December 2025

Panelist, *The Faculty Search Committee's Perspective*

Pomona College Department of Biology Seminar Series

Claremont, California, September 2025

From the lac operon to the spaghettiome, what can we learn from modeling gene regulatory networks on a medium scale?

Microscopy & Modeling Group Meeting

University of California, Los Angeles, December 2018

Dynamical Systems Modeling and Visualization of Yeast Cold Shock Gene Regulatory Networks: a Progress Report

Quantitative and Computational Biosciences Retreat

Malibu, California, September 2018

*Mathematical Modeling of Small GRNs Controlling the Cold Shock Response in Saccharomyces cerevisiae***BioQUEST Summer Workshop 2018, Wicked Problems: Investigating Real World Problems in the Biology Classroom**

Harvey Mudd College, Claremont, California, June 2018 (with Carrie Diaz Eaton)

*An Invitation to Modeling: Exploring the process of science through the process of modeling***Quantitative and Computational Biosciences Seminar**

University of California, Los Angeles, March 2018

*Dynamical Systems Modeling and Visualization of Gene Regulatory Networks: What Can We Learn from Networks on the "Medium" Scale?***BioQUEST / HHMI / CaseNet Summer Workshop 2017, Making Meaning Through Modeling: Problem Solving in Biology**

Michigan State University, East Lansing, Michigan, July 2017, (with Carrie Diaz Eaton, M. Drew LaMar, and Glenn Ledder)

*An Invitation to Modeling: Exploring the process of science through the process of modeling***Breaking the Boundaries in STEM Education Research Conference**

Loyola Marymount University, Los Angeles, California, April 2017

*A Framework for Models and Modeling to Unify Mathematicians and Biologists and Improve Student Learning***National Center for Ecological Analysis and Synthesis**

Santa Barbara, California, March 2017

*GRNmap and GRNsight: Open Source Software for Dynamical Systems Modeling and Visualization of Medium-Scale Gene Regulatory Networks***BioQUEST / HHMI / CaseNet Summer Workshop 2015, Count the Ways: Engaging Students in Quantitative Biology Applications**

Harvey Mudd College, Claremont, California, June 2015

*Open Science, Open Data, Open Source Projects for Undergraduate Research Experiences***SCELC (Statewide California Electronic Library Consortium) Colloquium**

Loyola Marymount University, February 2015

Panelist: *In the Open: the Future of Open Access Publishing and Libraries*Talk: *Open Access Publishing: A PUI Faculty Perspective***Chapman University**

Orange, California, May 2012

*Brrrr--How Do Yeast Cope When It's Cold Outside? Using DNA Microarrays and Mathematical Modeling to Understand Gene Regulatory Networks in Saccharomyces cerevisiae***Harvey Mudd College**

Claremont, California, April 2012

*Brrrr--How Do Yeast Cope When It's Cold Outside? Using DNA Microarrays and Mathematical Modeling to Understand Gene Regulatory Networks in Saccharomyces cerevisiae***Mount Saint Mary's College**

Los Angeles, California, March 2012

*Teaching and Learning Bioinformatics***Career Day in Fields of Science, Institute for Integrative Genome Biology, University of California, Riverside**

Riverside, California, May 2011

*Career Envy: The Road to a Successful PUI Position***Graduate Student Career Workshop, University of California, Los Angeles**

Los Angeles, California, February 2011

*Career Envy: The Road to a Successful PUI Position***Postdoctoral Scholars Association Career Workshop, University of California, Irvine**

Irvine, California, November 2010

Career Envy: The Road to a Successful PUI Position

Beyond Bio2010 Symposium: Celebration and Opportunities, National Academy of Sciences

Washington, D.C., May 2010 (with John David N. Dionisio)

An Open Source, Open Science Pedagogy for Computational Biology

Young Women in Computing and CREST, New Mexico State University

Las Cruces, New Mexico, February 2010

It's a Good Time to Be a Computational Biologist! and Bioinformatics Workshop

Pepperdine University

Malibu, California, February 2008

Guest lecture in Molecular Biology course: *MAPPFinder Analysis of Prostate Cancer*

Microarray Data

MCD Biology Department, University of California, Los Angeles

Los Angeles, California, May 2007

Mapping the Gene Regulatory Networks in Yeast that Control the Environmental Stress Response to Cold Temperatures

Gladstone Institute of Cardiovascular Disease

San Francisco, California, October 2006, joint seminar with John David N. Dionisio

XMLPipeDB: A Reusable, Open Source Tool Chain for Building Relational Databases from XML Sources

Bioinformatics Special Interest Group, California Institute of Technology

Pasadena, California, July 2006

Mapping Gene Regulatory Networks in Yeast using DNA Microarrays, Proteomics, and GenMAPP

Careers in Science Panel Discussion and Dinner, Claremont Colleges

Claremont, California, July 2006

Panelist

Natural Science Division, Pepperdine University

Malibu, California, October 2005

Mapping Gene Regulatory Networks in Yeast using DNA Microarrays, Proteomics, and GenMAPP

Department of Biological Sciences, Central Connecticut State University

New Britain, Connecticut, November 2004

Mapping Gene Regulatory Networks in Yeast using DNA Microarrays, Proteomics, and GenMAPP

BioQUEST Curriculum Consortium Summer Workshop 2004: Systems Biology Education

Beloit College, Beloit, Wisconsin, June 2004

GenMAPP and MAPPFinder for Systems Biology Education

Association for Laboratory Automation, smallTalk2003

San Jose, California, July 2003

GenMAPP and MAPPFinder: Tools for Viewing and Analyzing Microarray Data on Biological Pathways

W. Henry Feinstone Symposium, University of Memphis

Memphis, Tennessee, June 2003

Tutorial: GenMAPP and MAPPFinder, Tools for Viewing and Analyzing Microarray Data on using Biological Pathways and Gene Ontology

Seminar: Analysis of Microarray Data from a Mouse Model of Dilated Cardiomyopathy, New Insights from GenMAPP

Department of Plant Biology, The Carnegie Institution of Washington

Stanford, California, May 2003

GenMAPP and MAPPFinder: Tools for Viewing and Analyzing Microarray Data using Biological Pathways and Gene Ontology

Possibilities and Pitfalls of Mining DNA Microarray Data: from Mice to Men, University of Wyoming

Laramie, Wyoming, February 2003

*Tutorial: GenMAPP and MAPPFinder, Tools for Viewing and Analyzing Microarray Data on Biological Pathways**Seminar: Analysis of Microarray Data from a Mouse Model of Dilated Cardiomyopathy, New Insights from GenMAPP***Advanced Topics in Microarray Analysis, National Institutes of Health**

Bethesda, Maryland, January 2003

*GenMAPP and MAPPFinder, Tools for Viewing and Analyzing Microarray Data on Biological Pathways***Lillehei Heart Institute, University of Minnesota**

Minneapolis, Minnesota, October 2002

*Tutorial: GenMAPP and MAPPFinder, Tools for Viewing and Analyzing Microarray Data on Biological Pathways**Seminar: Analysis of Microarray Data from a Mouse Model of Dilated Cardiomyopathy, New Insights from GenMAPP***NIH-NHLBI Programs for Genomic Applications, External Scientific Panel Review**

Bethesda, Maryland, June 2001

*GenMAPP Enriches the BayGenomics Gene Trap Resource***Iconix Pharmaceuticals**

Mountain View, California, June 2001

*GenMAPP: A New Tool for the Functional Mapping of Microarray Data***Department of Neurosciences, University of New Mexico Health Sciences Center**

Albuquerque, New Mexico, October 2000

*Defining the Genomic Responses to G Protein Signals by Engineering Receptors and G Proteins in Transgenic Mice***National Center for Genome Resources**

Santa Fe, New Mexico, October 2000

*Defining the Genomic Responses to G Protein Signals by Engineering Receptors and G Proteins in Transgenic Mice***University of California, Berkeley, History of Science Graduate Student Workshop**

Berkeley, California, January 1997

*Panelist: The Relevance of History of Science to Practicing Scientists***Contributed Talks****Southern California Systems Biology Conference**

University of California, Irvine, March 2024

*Dynamical systems modeling and visualization of yeast cold shock gene regulatory networks***Joint Bioinformatics Community Conference and Bioinformatics Open Source Conference**

Virtual, July 2020

*New features for GRNsight: a web application for visualizing models of small- to medium-scale gene regulatory networks***American Society for Biochemistry and Molecular Biology Webinar: Constructing Authentic Learning Experiences** (in lieu of Annual Meeting that was cancelled due to the COVID-19 pandemic)

Virtual, June 2020

*An invitation to modeling*Abstract published in *The FASEB Journal*, <https://doi.org/10.1096/fasebj.2020.34.s1.07048>**Bioinformatics Open Source Conference (BOSC)**Orlando, Florida, July 2016; Slides in *F1000 Research* DOI: 10.7490/f1000research.1112534.1

GRNmap and GRNsight: open source software for dynamical systems modeling and visualization of medium-scale gene regulatory networks

American Society for Biochemistry and Molecular Biology Annual Meeting

San Diego, California, April 2016; published abstract in *The FASEB Journal* 30(1) Supplement

GRNmap and GRNsight: open source software for dynamical systems modeling and visualization of medium-scale gene regulatory networks

Fifth Annual Southern California Systems Biology Conference

University of California, Irvine, January 2015

GRNmap and GRNsight: Open Source Software for Dynamical Systems Modeling and Visualization of Medium-Scale Gene Regulatory Networks

American Society for Biochemistry and Molecular Biology Annual Meeting

San Diego, California, April 2012; published abstract in *The FASEB Journal* 26(1) Supplement

*Regulatory Dynamics of the Transcriptional Network Controlling the Cold Shock Response in *Saccharomyces cerevisiae**

Bioinformatics Open Source Conference (BOSC)

Stockholm, Sweden, June 2009

XMLPipeDB: A Reusable, Open Source Tool Chain for Building Relational Databases from XML Sources

Yeast Genetics and Molecular Biology Meeting

Toronto, Ontario, Canada, July 2008

*Mathematical Modeling of the Transcriptional Regulatory Network Controlling the Cold Shock Response in *Saccharomyces cerevisiae**

8th BioPathways Meeting

Vienna, Austria, July 2007

*Mathematical Modeling of the Transcriptional Network Controlling the Environmental Stress Response in *Saccharomyces cerevisiae**

Bioinformatics Open Source Conference (BOSC)

Vienna, Austria, July 2007 (two talks)

XMLPipeDB: A Reusable, Open Source Tool Chain for Building Relational Databases from XML Sources; An Open Source Framework for Teaching Bioinformatics

ELSI Reunion and Conference, Dartmouth University

Hanover, New Hampshire, August 2006

Discussion of Ethical, Legal, and Social Implications of Biological Research Incorporated into Courses in Genetics, Molecular Biology Applications, and a Seminar on Issues in Biotechnology

Bioinformatics Open Source Conference (BOSC)

Fortaleza, Brazil, August 2006

XMLPipeDB: A Reusable, Open Source Tool Chain for Building Relational Databases from XML Sources

The Fifth BioPathways Consortium Meeting, Intelligent Systems for Molecular Biology

Brisbane, Queensland, Australia, June 2003

GenMAPP and MAPPFinder 2.0: Tools for the Organization, Display, and Exchange of Pathway Information

The Fourth BioPathways Consortium Meeting, Intelligent Systems for Molecular Biology

Edmonton, Alberta, Canada, August 2002

GenMAPP and Gene Ontology: Tools for the Organization, Display and Exchange of Pathway Information

Physiological Genomics of Cardiovascular Disease: from Technology to Physiology

San Francisco, California, February 2002

GenMAPP: A New Tool for Viewing and Analyzing Microarray Data on Biological Pathways

Bay Area Bioinformatics Discussion Group

Stanford, California, January 2002

GenMAPP: A New Tool for Viewing and Analyzing Microarray Data on Biological Pathways

Bay Area RNA Club

San Francisco, California, June 1996

*Rites of Initiation: Decoding the role of IF1***Internal Talks****Department of Biology Seminar, Loyola Marymount University**

Los Angeles, California, April 2025

*Expanding the Narrative in STEM: Herman Branson***Department of Biology Seminar, Loyola Marymount University**

Los Angeles, California, September 2018

*From the lac operon to the spaghettiome, what can we learn from modeling gene regulatory networks on a medium scale?***Frank R. Seaver College Professorial Lecture, Loyola Marymount University**

Los Angeles, California, February 2018

*The Process is the Product: Systems Biology within an Open Science Ecosystem***Department of Biology Seminar, Loyola Marymount University**

Los Angeles, California, September 2016

*GRNmap and GRNsight: Using the power of genomics, mathematics, and open source visualization software to understand gene regulatory networks in yeast***Department of Biology Seminar, Loyola Marymount University**

Los Angeles, California, March 2013, with Dr. John David N. Dionisio

*XMLPipeDB: Teaming up to Analyze Data from Pathogenic Microorganisms***Department of Biology Seminar, Loyola Marymount University**

Los Angeles, California, October 2012

*Brrrr--How Do Yeast Cope When It's Cold Outside? Using DNA Microarrays and Mathematical Modeling to Understand Gene Regulatory Networks in *Saccharomyces cerevisiae****Friday Faculty Colloquium Series, Loyola Marymount University**

Los Angeles, California, February 2010

*The Genome is the New Soul***Biology/Bioethics Movie Night, Loyola Marymount University**

Los Angeles, California, October 2009

The Biology of Cancer, followed by a screening of the film *Wit***Junior Faculty Seminar, Loyola Marymount University**

Los Angeles, California, February 2009

*The Genome is the New Soul***Center for Teaching Excellence, Loyola Marymount University**

Los Angeles, California, October 2008 (with John David N. Dionisio)

*Create. Share. Learn. Using Google Sites and MediaWiki***President's Day Forum, Loyola Marymount University**

Los Angeles, California, March 2008

*The \$1000 Genome***Department of Biology, Loyola Marymount University, Kadner-Pitts Research Grant Talk**

Los Angeles, California, March 2008

*Brrrr—How Do Yeast Cope When It's Cold Outside? Using DNA Microarrays and Mathematical Modeling to Understand Gene Regulatory Networks in Yeast***Center for Teaching Excellence, Loyola Marymount University**

Los Angeles, California, March 2008

How Do You Teach "Research"? Incorporating DNA Microarray Technology into an Upper-division Biology Laboratory Course

Parent's Weekend, Loyola Marymount University

Los Angeles, California, February 2008

*How Close are We to GATTACA?***Center for Teaching Excellence, Loyola Marymount University**

Los Angeles, California, November 2007

*Panelist, Explorations of Faith and the Intellectual Life***President's Day Forum, Loyola Marymount University**

Los Angeles, California, March 2007

*How Close are We to GATTACA?***Science Seminar and Film Series, Loyola Marymount University**

Los Angeles, California, organized by LMU undergraduate Morgan Henry '07, November 2006

Our Post-genomic Future, accompanied by screening of *GATTACA***Junior Faculty Seminar Series, Loyola Marymount University**

Los Angeles, California, joint seminar with John David N. Dionisio, November 2006,

*Collaborating Early and Often: Bringing Biology and Computer Science Together Through an Open Source Culture***President's Day Forum, Loyola Marymount University**

Los Angeles, California, March 2006

*The Human Genome and Beyond***Women's Studies Brown Bag Lunch, Loyola Marymount University**

Los Angeles, California, November 2006

*Jesuit and Feminist Education: Transformative Discourses for Teaching & Learning Conference Report***Department of Mathematics, Loyola Marymount University**

Los Angeles, California, October 2005

*What is Bioinformatics?***Women's Studies Program First Friday, Vassar College**

Poughkeepsie, New York, October 2004

*The Ethical, Legal, and Social Implications of the Human Genome Project: Feminist Reflections (with Mary Shanley, Department of Political Science, Vassar College)***Vassar College Orientation Week Faculty Research Talks**

Poughkeepsie, New York, September 2004

*Matthew Vassar Enters the Genomics Era: DNA Microarrays, Proteomics, and Bioinformatics in Yeast***Gladstone Institute of Cardiovascular Disease Scientists Meeting**

San Francisco, California, May 2003

*GenMAPP 2.0 and Beyond...Connecting Scientists and Science Education in the Genomics Era***Gladstone Institute of Cardiovascular Disease Scientists Meeting**

San Francisco, California, May 2002

*Analysis of Microarray Data from a Mouse Model of Dilated Cardiomyopathy: New Insights from GenMAPP***U.C. San Francisco, Pharmaceutical Sciences and Pharmacogenomics Program Retreat**

Marshall, California, November 2001

*GenMAPP: A New Tool for Viewing and Analyzing Microarray Data on Biological Pathways***The J. David Gladstone Institutes Joint Scientific Retreat**

Pacific Grove, California, May 2001

*GenMAPP: A New Tool for the Functional Mapping of Microarray Data***Gladstone Institute of Neurological Disease Weekly Seminar**

San Francisco, California, November 2000

GenMAPP: A New Tool for the Functional Mapping of Microarray Data

Stanford University, Structural Biology Department Retreat

Pacific Grove, California, November 1998

*Interactions between Initiation Factor 1 and the E. coli ribosome***Stanford University, Molecular Biophysics Club**

Stanford, California, February 1998

*Interactions of Translation Initiation Factor 1 with the Ribosomal A site***U.C. Santa Cruz, MCD Biology Seminar**

Santa Cruz, California, May 1996

*Investigating the Structure and Function of Translation Initiation Factor 1 in E. coli***U.C. Santa Cruz, RNA Club**

Santa Cruz, California, December 1994

*Investigating the Structure and Function of Translation Initiation Factor 1 in E. coli***External Posters** (*indicates undergraduate co-author, **indicates Master's student co-author)**QCBio Retreat**

University of California, Los Angeles, September 22, 2025 (with Ben G. Fitzpatrick, John David N. Dionisio, Ngoc K. Tran*, Nikki C. Chun*, Cecilia J. Zaragoza*, A'Kaia L. Phelps*, Amelie T. Dinh*, Milka Y. Zekarias*)

*Sensitivity analysis of GRNmap and new features for GRNsight: open source software for dynamical systems modeling and visualization of small-scale gene regulatory networks***Intelligent Systems for Molecular Biology (ISMB)/European Conference for Computational Biology (ECCB)**

Liverpool, England, July 20-24, 2025 (with Ben G. Fitzpatrick, John David N. Dionisio, Ngoc K. Tran*, Nikki C. Chun*, Cecilia J. Zaragoza*, A'Kaia L. Phelps*, Amelie T. Dinh*, Milka Y. Zekarias*)

*Sensitivity analysis of GRNmap and new features for GRNsight: open source software for dynamical systems modeling and visualization of small-scale gene regulatory networks***The Allied Genetics Conference (TAGC)**

National Harbor, Maryland, March 6-10, 2024 (with Onariaginoso O. Igbinedion**, Ahmad R. Mersaghian*, Ngan Tran*, Deepa V. Dabir, John David N. Dionisio)

*Extending the GRNsight application for visualizing small-to-medium gene regulatory networks to incorporate physical protein-protein interaction data from SGD***Yeast Genetics Meeting**

University of California, Los Angeles, August 17-21, 2022 (with Onariaginoso O. Igbinedion**, Ahmad R. Mersaghian*, Sarron A. Tadesse*, John David N. Dionisio)

*Integration of SGD regulatory and expression data into the GRNmap and GRNsight applications for modeling and visualizing small-to-medium gene regulatory networks***Yeast Genetics Meeting**

Stanford University, Stanford, California, August 22-26, 2018 (with Ben G. Fitzpatrick, Brandon J. Klein*, Margaret J. O'Neil*, Lauren M. Kelly*)

*Mathematical modeling of small gene regulatory networks reveals key regulators and network properties important for controlling the early response to cold shock in Saccharomyces cerevisiae***Bioinformatics Open Source Conference (BOSC) and Intelligent Systems for Molecular Biology (ISMB)**, poster in *F1000 Research* DOI: 10.7490/f1000research.1112518.1

Orlando, Florida, July 8-12, 2016 (with Ben G. Fitzpatrick, John David N. Dionisio, Nicole A. Anguiano*, Juan S. Carrillo*, Tessa A. Morris*, Anindita Varshneya*, Natalie E. Williams*, K. Grace Johnson*, Trixie Anne M. Roque*, Kristen M. Horstmann*, Mihir Samdarshi*, Chukwuemeka E. Azinge*, Brandon J. Klein*, Margaret J. O'Neil*)

GRNmap and GRNsight: open source software for dynamical systems modeling and visualization of medium-scale gene regulatory networks

American Society for Biochemistry and Molecular Biology Annual Meeting

San Diego, California, April 2-5, 2016 (with Ben G. Fitzpatrick, John David N. Dionisio, Nicole A. Anguiano*, Juan S. Carrillo*, Kristen M. Horstmann*, Kayla C. Jackson*, K. Grace Johnson*, Tessa A. Morris*, Trixie Anne M. Roque*, Mihir Samdarshi*, and Anindita Varshneya*, Natalie E. Williams*), published abstract in *The FASEB Journal* 30(1) Supplement

GRNmap and GRNsight: open source software for dynamical systems modeling and visualization of medium-scale gene regulatory networks

Yeast Genetics and Molecular Biology Meeting

University of Washington, Seattle, Washington, July 29-August 3, 2014 (with Ben G. Fitzpatrick, Cybele Arsan*, Wesley T. Citti*, Kevin C. Entzminger*, Andrew F. Herman*, Lauren N. Kubeck*, Stephanie D. Kuelbs*, Heather King*, Elizabeth M. Liu*, Matthew Mejia*, Kenny R. Rodriguez*, Nicholas A. Rohacz*, Olivia S. Sakhon*, Katrina Sherbina*, Alondra J. Vega*)

*Cin5, Gln3, Hmo1, and Zap1 Contribute to the Gene Regulatory Network Controlling the Cold Shock Response in *Saccharomyces cerevisiae**

American Society for Biochemistry and Molecular Biology Annual Meeting

San Diego, California, April 26-30, 2014 (with Nicolette Harmon*, Chidinma Amakiri*, Katrina Sherbina*, Nicholas A. Rohacz*, and Ben G. Fitzpatrick), published abstract in *The FASEB Journal* 28(1) Supplement

*Comparative genomics of the response to cold shock in *Saccharomyces paradoxus* and *Saccharomyces cerevisiae**

American Society for Biochemistry and Molecular Biology Annual Meeting

San Diego, California, April 20-24, 2012 (with Ben G. Fitzpatrick, Nicholas A. Rohacz*, Katrina Sherbina*), published abstract in *The FASEB Journal* 26(1) Supplement

*Regulatory Dynamics of the Transcriptional Network Controlling the Cold Shock Response in *Saccharomyces cerevisiae**

I received the ASBMB Thematic Best Poster Award in Systems Biology for this poster.

Systems Biology: Global Regulation of Gene Expression

Cold Spring Harbor Laboratory, Cold Spring Harbor, New York, March 20-24, 2012 (with Ben G. Fitzpatrick, Nicholas A. Rohacz*, Katrina Sherbina*)

*Regulatory Dynamics of the Transcriptional Network Controlling the Cold Shock Response in *Saccharomyces cerevisiae**

Southern California Systems Biology Conference

University of California, Irvine, January 29-30, 2011 (with Alondra J. Vega*, Ben G. Fitzpatrick)

*Mathematical Modeling of the Gene Regulatory Network Controlling the Cold Shock Response in *Saccharomyces cerevisiae**

Yeast Genetics and Molecular Biology Meeting

Vancouver, British Columbia, Canada, July-August 2010 (with Alondra J. Vega*, Ben G. Fitzpatrick)

*Mathematical Modeling of the Gene Regulatory Network Controlling the Cold Shock Response in *Saccharomyces cerevisiae**

Intelligent Systems for Molecular Biology

Boston, Massachusetts, July 2010 (with Alondra J. Vega*, Stephanie D. Kuelbs, Ben G. Fitzpatrick)

*Mathematical Modeling of the Gene Regulatory Network Controlling the Cold Shock Response in *Saccharomyces cerevisiae**

American Society for Cell Biology Annual Meeting

San Diego, California, December 2009 (with John David N. Dionisio)

Fostering Interdisciplinary Teamwork in an Undergraduate Biological Databases Course

Intelligent Systems for Molecular Biology

Stockholm, Sweden, June 2009 (with, Alexandra Alphonso*, Derek Smith*, Chad Villaflores*, John David N. Dionisio)

XMLPipeDB: A Reusable, Open Source Tool Chain for Building Relational Databases from XML Sources

First RECOMB Satellite Conference on Bioinformatics Education

San Diego, California, March 2009 (with John David N. Dionisio)

Fostering Interdisciplinary Teamwork in an Undergraduate Biological Databases Course

Yeast Genetics and Molecular Biology Meeting

Toronto, Ontario, Canada, July 2008 (with Stephanie D. Kuelbs*, Kevin C. Entzminger*, Kenny R. Rodriguez*, Ben G. Fitzpatrick)

*Mathematical Modeling of the Transcriptional Regulatory Network Controlling the Cold Shock Response in *Saccharomyces cerevisiae**

Intelligent Systems for Molecular Biology

Toronto, Ontario, Canada, July 2008 (with Stephanie D. Kuelbs*, Kevin C. Entzminger*, Kenny R. Rodriguez*, Ben G. Fitzpatrick)

*Mathematical Modeling of the Transcriptional Regulatory Network Controlling the Cold Shock Response in *Saccharomyces cerevisiae**

International Conference on Systems Biology

Long Beach, California, October 2007 (with Stephanie Kuelbs*, Nathan C. Wanner*, Ben G. Fitzpatrick, and Erika Camacho)

*Mathematical Modeling of the Transcriptional Network Controlling the Environmental Stress Response in *Saccharomyces cerevisiae**

Intelligent Systems for Molecular Biology

Vienna, Austria, July 2007 (with Nathan C. Wanner* and Erika Camacho)

*Mathematical Modeling of the Transcriptional Network Controlling the Environmental Stress Response in *Saccharomyces cerevisiae**

San Diego Systems Biology Symposium: Systems to Synthesis

Salk Institute, La Jolla, California, January 2007 (with Jeffrey Nicholas** and John David N. Dionisio)

XMLPipeDB: A Reusable, Open Source Tool Chain for Building Relational Databases from XML Sources

American Society for Cell Biology Annual Meeting

San Diego, California, December 2006 (with Wesley T. Citti*, Matthew Mejia*, Eric S. Eberhardt)
*The Transcriptional and Proteomic Response to Cold Shock and Recovery in *Saccharomyces cerevisiae**

Intelligent Systems for Molecular Biology

Fortaleza, Brazil, August 2006 (with, Joey Barrett**, Joe Boyle**, Adam Carasso**, David Hoffman**, Babak Naffas**, Jeffrey Nicholas**, Roberto Ruiz**, Scott Spicer**, John David N. Dionisio)

XMLPipeDB: A Reusable, Open Source Tool Chain for Building Relational Databases from XML Sources

Intelligent Systems for Molecular Biology

Glasgow, Scotland, United Kingdom, August 2004

GenMAPP and MAPPFinder 2.0: Tools for Viewing and Analyzing Genomic Data Using Gene Ontology and Biological Pathways

Intelligent Systems for Molecular Biology

Brisbane, Queensland, Australia, June 2003

GenMAPP and MAPPFinder 2.0: Tools for Viewing and Analyzing Genomic and Proteomic Data Using Gene Ontology and Biological Pathways

Intelligent Systems for Molecular Biology

Edmonton, Alberta, Canada, August 2002

GenMAPP: A Tool for Viewing and Analyzing Microarray Data on Biological Pathways

Physiological Genomics of Cardiovascular Disease: from Technology to Physiology

San Francisco, California, February 2002

GenMAPP: A New Tool for Viewing and Analyzing Microarray Data on Biological Pathways

The Third International Meeting on Microarray Data Standards, Annotations, Ontologies and Databases

Stanford, California, March 2001

*GenMAPP: A New Approach for the Functional Mapping of Microarray Data***The Ribosome: Structure, Function, Antibiotics, and Cellular Interactions**

Helsingør, Denmark, June 1999

*Interactions of Translation Initiation Factor 1 with the Ribosomal A site***RNA Society Meeting**

Madison, Wisconsin, May 1998

*Interactions of Translation Initiation Factor 1 with the Ribosomal A site***RNA Structure Meeting**

Santa Cruz, California, June 1997

*Interactions of Translation Initiation Factor 1 (IF1) with the Ribosomal A site***RNA Society Meeting**

Banff, Alberta, Canada, May 1997

*Interactions of Translation Initiation Factor 1 with the Ribosomal A site***Keystone Symposium: RNA-Protein Interactions**

Taos, New Mexico, February 1997

*Interactions of Translation Initiation Factor 1 (IF1) with the Ribosomal A site***RNA Society Meeting**

Madison, Wisconsin, May 1996

*Translation Initiation Factor 1 (IF1) is an A-site Ribosomal RNA Binding Protein***Symposium on RNA Biology I: RNA-Protein Interactions**

Research Triangle Park, North Carolina, October 1995

*Investigating the Structure and Function of Translation Initiation Factor 1 in Escherichia coli***Frontiers in Translation**

Victoria, British Columbia, Canada, May 1995

*Investigating the Structure and Function of Translation Initiation Factor 1 in E. coli***Sigma Xi Forum: Scientists, Educators, and National Standards: Action at the Local Level**

Atlanta, Georgia, April 1994

*Science Mentor Program at Mission Hill Junior High School***Internal Posters****Center for Teaching Excellence Scholarship of Teaching and Learning Showcase Week**

Loyola Marymount University, Los Angeles, California, September 2009

*Fostering Interdisciplinary Teamwork in an Undergraduate Biological Databases Course***Center for Teaching Excellence Scholarship of Teaching and Learning Showcase Week**

Loyola Marymount University, Los Angeles, California, September 2008

*How Do You Teach "Research"? Incorporating DNA Microarray Technology into an Upper-division Biology Laboratory Course***Teaching with Technology Forum**

Vassar College, Poughkeepsie, New York, April 2004

*GenMAPP: Connecting Students to Cutting-edge Genomics and Bioinformatics Research***The J. David Gladstone Institutes Joint Scientific Retreat**

Pacific Grove, California, May 2003

*GenMAPP and MAPPFinder 2.0***U.C. San Francisco, Pharmaceutical Sciences and Pharmacogenomics Program Retreat**

Marshall, California, November 2002

GenMAPP: A Tool for Viewing and Analyzing Microarray Data on Biological Pathways

The J. David Gladstone Institutes Joint Scientific Retreat

Pacific Grove, California, May 2002

*Analysis of Microarray Data from Mouse Models of Dilated and Hypertrophic Cardiomyopathy: New Insights from GenMAPP***U.C. San Francisco, Cardiovascular Research Institute Retreat**

Tahoe City, California, November 2001

*GenMAPP: A New Tool for Viewing and Analyzing Microarray Data on Biological Pathways***U.C. San Francisco, Biomedical Sciences Program Retreat**

Tahoe City, California, November 2000

*GenMAPP: A New Approach for the Functional Mapping of Microarray Data***U.C. San Francisco, Tetrad Retreat**

Tahoe City, California, September 2000

*GenMAPP: A New Approach for the Functional Mapping of Microarray Data***Student Presentations and Posters** (bold indicates presenting author)**Annual Biomedical Research Conference for Minoritized Scientists**

San Antonio, Texas, November 19-22, 2025

Nikki C. Chun, Ben G. Fitzpatrick, and Kam D. Dahlquist, *Accuracy of parameter estimation for a simple gene regulatory network model is sensitive to the number of parameters estimated and the magnitude and direction of regulatory relationships* (poster)**Southern California Conference for Undergraduate Research (SCCUR)**

Cal-State University, Channel Islands, Camarillo, California, November 22, 2025

Ngoc K. Tran, Milka Y. Zekarias, Cecilia J. Zaragoza, A'Kaia L. Phelps, Amelie T. Dinh, John David N. Dionisio, Kam D. Dahlquist, *GRNsight v7: a web application for visualizing gene regulatory networks and protein-protein interactions* (talk)**West Coast Biological Sciences Undergraduate Research Conference**

California State University San Marcos, April 12, 2025

Nikki C. Chun, Ben G. Fitzpatrick, and Kam D. Dahlquist, *Accuracy of parameter estimation for a simple gene regulatory network model is sensitive to the number of parameters estimated and the magnitude and direction of regulatory relationships* (poster)**17th Annual Loyola Marymount University Undergraduate Research Symposium**

March 22, 2025

Nikki C. Chun, Ben G. Fitzpatrick, and Kam D. Dahlquist, *Accuracy of parameter estimation for a simple gene regulatory network model is sensitive to the number of parameters estimated and the magnitude and direction of regulatory relationships* (poster)**Ngoc K. Tran, Cecilia J. Zaragoza, A'Kaia L. Phelps, Amelie T. Dinh, and Milka Y. Zekarias**, John David N. Dionisio, Kam D. Dahlquist, *Database, Display, and DevOps Improvements for GRNsight 7.2, a Web Application for Visualizing Gene Regulatory and Protein-Protein Interaction Network Models* (poster)**National Conference for Undergraduate Research (NCUR)**

Long Beach California, April 2024

Catherine I. Channell, Odo O. Okwuosa, **Naomi Mesfin**, Nathanael Wudineh, Kam D. Dahlquist, *Determining the Genotype Frequency of SNP rs4988235 That Confers Lactase Persistence in an LMU Population* (talk)**Southern California Systems Biology Conference**

University of California, Irvine, March 2024

Nikki C. Chun, Ben G. Fitzpatrick, and Kam D. Dahlquist, *Parameter sensitivity analysis of GRNmap, a dynamical systems model of gene regulatory networks* (poster)**16th Annual Loyola Marymount University Undergraduate Research Conference**

March 2024

Catherine Channell, Odoaba Okwuosa, Naomi Mesfin, Kam D. Dahlquist: *Determining the genotype frequency of SNP rs4988235 that confers lactase persistence in an LMU population* (talk)

Nikki C. Chun, Ben G. Fitzpatrick, and Kam D. Dahlquist, *Parameter sensitivity analysis of GRNmap, a dynamical systems model of gene regulatory networks* (poster)

Ngoc K. Tran, Cecilia J. Zaragoza, A'Kaia Phelps, John David N. Dionisio, Kam D. Dahlquist, *Propagating Protein-Protein Interaction Network Support into GRNsight 7.0, a Web Application for Visualizing Gene Regulatory Network Models* (poster)

West Coast Biological Sciences Undergraduate Research Conference

Loyola Marymount University, April 2023

Catherine Channell, Odoaba Okwuosa, Kam D. Dahlquist, *Isolating DNA from human hair to determine the frequency of the lactase persistence at SNP rs4988235* (poster)

15th Annual Loyola Marymount University Undergraduate Research Conference

March 2023

Catherine Channell, Odoaba Okwuosa, Kam D. Dahlquist, *Determining the genotype frequency of SNP rs4988235 that confers lactase persistence in an LMU population.* (poster)

Ahmad Mersaghian, Sarron Tadesse, Ngan N. Tran, John David N. Dionisio, Kam D. Dahlquist, *Improved Functionality of GRNsight 6.0: a Web Application for Visualizing Gene Regulatory Network Models* (poster)

West Coast Biological Sciences Undergraduate Research Conference

Point Loma Nazarene University, April 2022

Catherine Channell, Odoaba Okwuosa, Kam D. Dahlquist, *Isolating DNA from human hair to determine the frequency of the lactase persistence at SNP rs4988235* (poster)

Ahmad Mersaghian, Sarron Tadesse, John David N. Dionisio, Kam D. Dahlquist, *A New Gene Expression Dataset for GRNsight: a Web Application for Visualizing Gene Network Models* (poster)

Claire Kosewic, Erykah Walton, Elizabeth Haugan, Kam D. Dahlquist, *Worldwide frequencies of SNPs conferring lactase persistence in humans* (talk)

14th Annual Loyola Marymount University Undergraduate Research Conference

March, 2022

Catherine Channell, Odoaba Okwuosa, Kam D. Dahlquist, *Isolating DNA from human hair to determine the frequency of the lactase persistence at SNP rs4988235* (poster)

Ahmad Mersaghian, Sarron Tadesse, John David N. Dionisio, Kam D. Dahlquist, *A New Gene Expression Dataset for GRNsight: a Web Application for Visualizing Gene Network Models* (poster)

Onariaginosa O. Igbinedion, John David N. Dionisio, Kam D. Dahlquist, *New Databases and Export Features for GRNsight: a Web Application for Visualizing Models of GRNs* (poster)

Claire Kosewic, Erykah Walton, Elizabeth Haugan, Kam D. Dahlquist, *Worldwide frequencies of SNPs conferring lactase persistence in humans* (talk)

American Society for Biochemistry and Molecular Biology Annual Meeting

Virtual, April 2021

Madeleine B. King^{*^}, Annika G. Dinulos^{*}, Jessica A. DaMota^{*}, Erykah I. Walton^{*}, Kam D. Dahlquist, *A bioinformatics approach to investigating the structural and functional consequences of SNPs in TMPRSS2 for COVID-19 infection* (poster)

Abstract published in *The FASEB Journal*, <https://doi.org/10.1096/fasebj.2021.35.S1.04753>

[^]**Madeleine B. King won Honorable Mention in the Undergraduate Poster Competition**

13th Annual Loyola Marymount University Undergraduate Research Conference

Virtual, March 2021

Madeleine B. King^{*}, Annika G. Dinulos^{*}, Jessica A. DaMota^{*}, and Kam D. Dahlquist, *Bioinformatic analysis of critical residues for binding of SARS-CoV-2 spike protein to host TMPRSS2* (talk)

Erykah I. Walton*, **Odoba O. Okwuosa***, Kam D. Dahlquist, *Bioinformatic analysis of coding SNPs of the human TMPRSS2 protease: implications for SARS-CoV-2* (poster)

Onariaginos O. Igbiniedion*, **Ian M. Green***, **Ahmad R. Mersaghian***, John David N. Dionisio, Kam D. Dahlquist, *More Robust Testing of Data and UI for GRNsight: a Web Application for Visualizing Models of GRNs* (poster)

Claire M. Kosewic*, **Catherine I. Channell***, Kam D. Dahlquist, *Bioinformatics data mining to determine the lactase persistence trait frequency in world populations* (poster)

American Society for Biochemistry and Molecular Biology Annual Meeting

Scheduled for April 2020, cancelled due to the COVID-19 pandemic

Susanne M. Carpenter*, **Alyssa N. Weisblatt***, Kam D. Dahlquist, *Development of RFLP and qPCR Assays to Detect the Human C/T-13910 SNP, which Is Associated with Lactase Persistence* (poster)

Abstract published in *The FASEB Journal*, <https://doi.org/10.1096/fasebj.2020.34.s1.07025>

Alice K. Finton*, Lauren M. Kelly*, Ben G. Fitzpatrick, Kam D. Dahlquist, *Modeling the Dynamics of a Family of Small Gene Regulatory Networks in which Single Edges Have Been Deleted Indicates Regulation by Hmo1, Msn2, and Cin5 Are Important for Controlling the Cold Shock Response in *Saccharomyces cerevisiae** (poster)

Abstract published in *The FASEB Journal*, <https://doi.org/10.1096/fasebj.2020.34.s1.06538>

12th Annual Loyola Marymount University Undergraduate Research Conference

Virtual, April 2020

Alice K. Finton*, Ben G. Fitzpatrick, Kam D. Dahlquist, *Modeling of Gene Regulatory Network Dynamics Reveals Important Regulatory Relationships that Control the Cold Shock Response in *Saccharomyces cerevisiae** (talk)

Alexia M. Filler*, **Kevin B. Patterson***, John David N. Dionisio, Kam D. Dahlquist, *An Expression Database for Node Coloring and Expansion of Species for Gene Pages for GRNsight* (talk)

West Coast Biological Sciences Undergraduate Research Conference

University of San Diego, April 2019

Genesis J. Cruz*, Kam D. Dahlquist, *Cultural Context and Global Health Implications of the Lactase Persistence SNP in Humans* (poster)

11th Annual Undergraduate Research Conference

Loyola Marymount University, March 2019

Lauren M. Kelly*, Ben G. Fitzpatrick, Kam D. Dahlquist, *Modeling of Gene Regulatory Network Dynamics Predicts which Regulatory Relationships are Important for Controlling the Cold Shock Response in *Saccharomyces cerevisiae** (talk)

Mihir Samdarshi*, John David N. Dionisio, Kam D. Dahlquist, *New Layouts, Data Types, and Architecture for GRNsight 3: a Web Application for Visualizing Gene Regulatory Networks* (talk)

Genesis J. Cruz*, Kam D. Dahlquist, *Cultural Context and Global Health Implications of the Lactase Persistence SNP in Humans* (poster)

8th Annual Southern California Systems Biology Conference

University of California, Irvine, February 2019

Lauren M. Kelly*, Margaret J. O'Neil, Ben G. Fitzpatrick, Kam D. Dahlquist, *Modeling of Gene Regulatory Network Dynamics Predicts which Regulatory Relationships are Important for Controlling the Cold Shock Response in *Saccharomyces cerevisiae** (poster)

Mihir Samdarshi*, John L. Lopez, John David N. Dionisio, Kam D. Dahlquist, *New Layouts, Data Types, and Architecture for GRNsight 3: a Web Application for Visualizing Gene Regulatory Networks* (poster)

2018 Beta Beta Beta Biological Honor Society's Pacific District Convention

Concordia University, Irvine, California, March 2018

Lauren M. Kelly*, Margaret J. O'Neil, Ben G. Fitzpatrick, Kam D. Dahlquist, *Modeling of Gene Regulatory Network Dynamics Predicts which Regulatory Relationships are Important for Controlling the Cold Shock Response in Saccharomyces cerevisiae* (poster)

Brandon J. Klein*, Ben G. Fitzpatrick, Kam D. Dahlquist, *Mathematical Modeling of Six Database-Derived Gene Regulatory Networks Identifies Key Regulators and Network Properties Controlling the Early Response to Cold Shock in Saccharomyces cerevisiae* (talk)

Brandon was awarded second place for the Frank G. Brooks Award for Excellence in Student Research for this talk.

Margaret J. O'Neil*, Ben G. Fitzpatrick, Kam D. Dahlquist, *Using Graph Statistics to Investigate the Properties of a Gene Regulatory Network that May Control the Cold Shock Response in Saccharomyces cerevisiae* (talk)

Mihir Samdarshi*, John David N. Dionisio, Kam D. Dahlquist, *Data Comparison Features and Development Tool Improvements for GRNsight: a Web App for Visualizing Gene Regulatory Networks* (poster)

Mihir was awarded Honorable Mention for the John C. Johnson Award for Excellence in Student Research for a posters presentation.

Nika Vafadari*, **Katherine D. Scheker***, Kam D. Dahlquist, *Identifying Regulatory Transcription Factors that Control Gene Expression Changes Due to Cold Shock in Saccharomyces cerevisiae* (talk)

10th Annual Undergraduate Research Conference

Loyola Marymount University, March 2018

Eileen J. Choe*, John David N. Dionisio, Kam D. Dahlquist, *Extending the Visualization Capabilities of GRNsight: a Web Application for Visualizing Models of Gene Regulatory Networks* (talk)

Lauren M. Kelly*, Margaret J. O'Neil, Ben G. Fitzpatrick, Kam D. Dahlquist, *Modeling of Gene Regulatory Network Dynamics Predicts which Regulatory Relationships are Important for Controlling the Cold Shock Response in Saccharomyces cerevisiae* (poster)

Brandon J. Klein*, Ben G. Fitzpatrick, Kam D. Dahlquist, *Mathematical Modeling of Six Database-Derived Gene Regulatory Networks Identifies Key Regulators and Network Properties Controlling the Early Response to Cold Shock in Saccharomyces cerevisiae* (talk)

Margaret J. O'Neil*, Ben G. Fitzpatrick, Kam D. Dahlquist, *Using Graph Statistics to Investigate the Properties of a Gene Regulatory Network that May Control the Cold Shock Response in Saccharomyces cerevisiae* (talk)

Mihir Samdarshi*, John David N. Dionisio, Kam D. Dahlquist, *Data Comparison Features and Development Tool Improvements for GRNsight: a Web App for Visualizing Gene Regulatory Networks* (poster)

Yeon-Soo Shin*, John David N. Dionisio, Kam D. Dahlquist, *New Graph Layouts for GRNsight: a Web Application for Visualizing Models of Gene Regulatory Networks* (talk)

Justin Kyle T. Torres*, **John L. Lopez***, Ben G. Fitzpatrick, John David N. Dionisio, Kam D. Dahlquist, *Paying Off Our Technical Debt for GRNmap, a Gene Regulatory Network Modeling Application* (poster)

Nika Vafadari*, **Katherine D. Scheker***, Kam D. Dahlquist, *Identifying Regulatory Transcription Factors that Control Gene Expression Changes Due to Cold Shock in Saccharomyces cerevisiae* (talk)

Southern California Conference for Undergraduate Research

California State Polytechnic University, Pomona, November 2017

Chukwuemeka E Azinge*, **Justin Kyle T. Torres***, John David N. Dionisio, Ben G. Fitzpatrick, Kam D Dahlquist, *Restructuring the Data Architecture of GRNmap, a Gene Regulatory Network Modeling Application* (poster)

Eileen J. Choe*, **Yeon-Soo Shin***, **Edward B. Bachoura***, John David N. Dionisio, Kam D Dahlquist, *GRNsight v2: a Web Application for Visualizing Models of Gene Regulatory Networks*, (talk)

Yeon-Soo Shin*, Eileen J. Choe*, Edward B. Bachoura*, Ben G. Fitzpatrick, John David N. Dionisio, Kam D Dahlquist, *Improved Visual Performance and Enhanced Test Files for Different File Formats for GRNsight: a Web Application for Visualizing Models of Gene Regulatory Networks*, (poster)

WE17: Society for Women Engineers (SWE) Collegiate Competition

Austin, Texas, October 2017

Eileen J. Choe*, Nicole A. Anguiano*, Anindita Varshneya*, Mihir Samdarshi*, Yeon-Soo Shin*, Edward B. Bachoura*, John David N. Dionisio, and Kam D. Dahlquist, *GRNsight v2: a web application for visualizing models of small gene regulatory networks* (talk)

9th Annual Undergraduate Research Symposium

Loyola Marymount University, March 2017

Nicole A. Anguiano*, **Anindita Varshneya***, John David N. Dionisio, Kam D. Dahlquist, *Design and Layout Improvement to GRNsight v2.0: a Web Application and Service for Visualizing Small- to Medium-Scale Gene Regulatory Networks* (talk)

Monica V. Hong*, Kam D. Dahlquist, *The transcription factors Hap4 and Swi4 contribute to the regulation of the transcriptional response to cold shock in Saccharomyces cerevisiae* (talk)

Kristen M. Horstmann*, Ben G. Fitzpatrick, and Kam D. Dahlquist, *Systems modeling and statistical analysis allows comparison in the response to cold shock* (talk) in *Saccharomyces cerevisiae* between Hap4 and randomly generated networks

Brandon J. Klein*, Natalie E. Williams*, Ben G. Fitzpatrick, and Kam D. Dahlquist, *Dynamical systems modeling of six related small gene regulatory networks suggest that the transcription factors Cin5, Gln3, Hmo1, and Yhp1 play a role in controlling the cold shock response in Saccharomyces cerevisiae* (poster)

Margaret J. O'Neil*, Ben G. Fitzpatrick, Kam D. Dahlquist, *Using Graph Statistics to Investigate the Properties of Six Candidate Gene Regulatory Networks for Controlling the Cold Shock Response in Saccharomyces cerevisiae* (poster)

Trixie Anne M. Roque*, **Chukwuemeka E. Azinge***, **Justin Kyle T. Torres***, John David N. Dionisio, Ben G. Fitzpatrick, Kam D. Dahlquist, *Restructuring the Data Architecture of GRNmap, a Gene Regulatory Network Modeling Application* (poster)

Mihir Samdarshi*, **Yeon-Soo Shin***, **Edward B. Bachoura***, **Eileen J. Choe***, Nicole A. Anguiano*, Anindita Varshneya*, John David N. Dionisio, Kam D. Dahlquist, *Improved data interoperability for GRNsight: a web application for visualizing models of gene regulatory networks* (poster)

Nika Vafadari*, Katherine D. Scheker*, Kam D. Dahlquist, *Targeted reverse genetic screen in Saccharomyces cerevisiae identifies transcription factor deletion strains that are impaired for growth at cold temperatures* (poster)

Natalie E. Williams*, Ben G. Fitzpatrick, Kam D. Dahlquist, *Comparison of the regulatory dynamics of related small gene regulatory networks that control the cold shock response in Saccharomyces cerevisiae* (talk)

7th Annual Southern California Systems Biology Conference

University of California, Irvine, January 2017

Monica V. Hong*, Kevin W. Wyllie*, Kevin P. McGee*, Kam D. Dahlquist, *The transcription factors Hap4 and Swi4 contribute to the regulation of the transcriptional response to cold shock in Saccharomyces cerevisiae* (poster)

Kristen M. Horstmann*, **Margaret J. O'Neil***, Ben G. Fitzpatrick, Kam D. Dahlquist, *Dynamical systems modeling and gene regulatory network structure analysis reveals Hap4's role in regulating the response to cold shock in Saccharomyces cerevisiae* (poster)

Anindita Varshneya*, Mihir Samdarshi*, Nicole A. Anguiano*, Yeon-Soo Shin*, John David N. Dionisio, and Kam D. Dahlquist, *New features improve GRNsight: a web application and service for visualizing models of small- to medium-scale gene regulatory networks* (poster)

Natalie E. Williams*, **Brandon J. Klein***, Ben G. Fitzpatrick, and Kam D. Dahlquist, *Dynamical systems modeling of six related small gene regulatory networks suggest that the*

transcription factors Cin5, Hmo1, Msn2, and Yhp1 play a role in controlling the cold shock response in Saccharomyces cerevisiae, (poster)

American Society for Biochemistry and Molecular Biology Annual Meeting

San Diego, California, April 2016; published abstracts in *The FASEB Journal* 30(1) Supplement

K. Grace Johnson*, Natalie E. Williams*, Ben G. Fitzpatrick, and Kam D. Dahlquist, *Modeling the Dynamics of a 21-gene, 50-edge Gene Regulatory Network Controlling the Transcriptional Response to Cold Shock in Saccharomyces cerevisiae using GRNmap* (poster)

Tessa A. Morris*, **Kristen M. Horstmann***, **Kayla C. Jackson***, Ben G. Fitzpatrick, and Kam D. Dahlquist, *Mathematical Modeling Shows that Gln3 and Zap1 Affects the Dynamics of the Gene Regulatory Network Controlling the Cold Shock Response in Saccharomyces cerevisiae* (poster)

Anindita Varshneya*, **Mihir Samdarshi***, Kam D. Dahlquist, John David N. Dionisio, and Ben G. Fitzpatrick, *Test-driven development improves GRNsight: a web application for visualizing models of gene regulatory networks* (poster)

Kevin W. Wyllie*, **Kevin P. McGee***, **Monica V. Hong***, Kam D. Dahlquist, *The Transcription Factors Swi4 and Hap4 Contribute to the Regulation of the Transcriptional Response to Cold Shock in Saccharomyces cerevisiae* (poster)

8th Annual Undergraduate Research Symposium

Loyola Marymount University, March 2016

Juan S. Carrillo Quinche*, **Trixie Anne M. Roque***, Kam D. Dahlquist, and John David N. Dionisio, *Usability Improvements to GRNmap: Software for Gene Regulatory Network Modeling and Parameter Estimation* (talk)

Kristen M. Horstmann*, Tessa A. Morris*, Brandon J. Klein*, Kam D. Dahlquist, and Ben G. Fitzpatrick, *Mathematical Modeling Reveals Zap1's Role in the Gene Regulatory Network that Controls the Response to Cold Shock in Saccharomyces cerevisiae* (poster)

K. Grace Johnson*, **Margaret J. O'Neil***, Kam D. Dahlquist, and Ben G. Fitzpatrick, *Evaluating Hap4's Role in the Gene Regulatory Network that Controls the Response to Cold Shock in Saccharomyces cerevisiae using GRNmap* (poster)

Tessa A. Morris*, Kam D. Dahlquist, Ben G. Fitzpatrick, *Mathematical Modeling Shows that Gln3 Affects the Dynamics of the Gene Regulatory Network Controlling the Cold Shock Response in Saccharomyces cerevisiae* (talk)

Anindita Varshneya*, **Mihir Samdarshi***, Kam D. Dahlquist, John David N. Dionisio, and Ben G. Fitzpatrick, *Test-driven development improves GRNsight: a web application for visualizing models of gene regulatory networks* (poster)

Kevin W. Wyllie*, **Monica V. Hong***, Kam D. Dahlquist, *The Transcription Factors Swi4 and Hap4 Contribute to the Regulation of the Transcriptional Response to Cold Shock in Saccharomyces cerevisiae* (poster)

Society for the Advancement of Chicanos and Native Americans in Science National Conference

Washington, D.C., October 2015

Trixie Anne M. Roque*, Tessa A. Morris*, Kam D. Dahlquist, John David N. Dionisio, and Ben G. Fitzpatrick, *Test-Driven Development and Functionality Improvements to GRNmap, a Gene Regulatory Network Modeling Application* (poster)

West Coast Biological Sciences Undergraduate Research Conference

Point Loma Nazarene University, San Diego, California, April 2015

Nicole Anguiano*, **Anindita Varshneya***, Kam D. Dahlquist, John David N. Dionisio, Ben G. Fitzpatrick, *Improvements to GRNsight: a Web Application for Visualizing Models of Gene Regulatory Networks* (poster)

Monica Hong*, **Kevin Wyllie***, Kam D. Dahlquist, *The Transcription Factor Swi4 Contributes to the Regulation of the Transcriptional Response to Cold Shock in Saccharomyces cerevisiae* (poster)

Natalie Williams*, **K. Grace Johnson***, Kam D. Dahlquist, Ben G. Fitzpatrick, *Comparing the Dynamics of the Cold Shock Gene Regulatory Network in Yeast with a Random Network* (poster)

7th Annual Undergraduate Research Symposium

Loyola Marymount University, March 2015

Nicole Anguiano*, **Anindita Varshneya***, Kam D. Dahlquist, John David N. Dionisio, Ben G. Fitzpatrick, *Improvements to GRNsight: a Web Application for Visualizing Models of Gene Regulatory Networks* (poster)**Juan Carrillo***, **Trixie Anne Roque***, Kam D. Dahlquist, Ben G. Fitzpatrick, *Software refactoring and Usability Enhancement for GRNmap, a Gene Regulatory Network Modeling Application* (poster)**Monica Hong***, **Kevin Wyllie***, Kam D. Dahlquist, *The Transcription Factor Swi4 Contributes to the Regulation of the Transcriptional Response to Cold Shock in Saccharomyces cerevisiae* (poster)**Natalie Williams***, **K. Grace Johnson***, Kam D. Dahlquist, Ben G. Fitzpatrick, *Comparing the Dynamics of the Cold Shock Gene Regulatory Network in Yeast with a Random Network* (poster)**5th Annual Southern California Systems Biology Conference**

University of California, Irvine, January 2015

Nicole Anguiano*, **Anindita Varshneya***, Kam D. Dahlquist, John David N. Dionisio, Ben G. Fitzpatrick, *Improvements to GRNsight: a Web Application for Visualizing Models of Gene Regulatory Networks* (poster)**Southern California Conference for Undergraduate Research**

California State University, Fullerton, November 2014

Nicole Anguiano*, **Anindita Varshneya***, Kam D. Dahlquist, John David N. Dionisio, Ben G. Fitzpatrick, *Improvements to GRNsight: a Web Application for Visualizing Models of Gene Regulatory Networks* (poster)**Sarah Patno***, Kam D. Dahlquist, John David N. Dionisio, *Constructing a Combined Gene Database for Staphylococcus aureus strains MRSA252 and COL for the Analysis of Microarray Data* (poster)**Mitchell Petredis***, Kam D. Dahlquist, John David N. Dionisio, *Gene Database Construction and GenMAPP Analysis of Sinorhizobium meliloti Microarray Data Comparing Salt and Sucrose Stress* (talk)**Society for the Advancement of Chicanos and Native Americans in Science National Conference**

Los Angeles, California, October 2014

Juan S. Carrillo*, **Katrina Sherbina***, Kam D. Dahlquist, Ben G. Fitzpatrick, *Software Refactoring and Usability Enhancement for GRNmap, a Gene Regulatory Network Modeling Application* (poster)**Beta Beta Beta Pacific District Convention**

Chapman University, Orange, California, April 2014

Kevin McGee*, Kam D. Dahlquist, John David N. Dionisio, *Generating a New Gene Database for Leishmania major and Leishmania infantum for Analyzing Microarray Data* (poster)**Mitchell Petredis***, Kam D. Dahlquist, John David N. Dionisio, *Gene Database Construction and GenMAPP Analysis of Sinorhizobium meliloti Microarray Data Comparing Salt and Sucrose Stress* (poster)**Andrew Pita***, Kam D. Dahlquist, John David N. Dionisio, *Constructing a GenMAPP-compatible Gene Database for Streptococcus pneumoniae to perform pathway analysis on microarray data comparing biofilm versus planktonic forms* (talk)**6th Annual Undergraduate Research Symposium**

Loyola Marymount University, March 2014

Kevin McGee*, Kam D. Dahlquist, John David N. Dionisio, *Pathway Analysis of Leishmania major Promastigote and Amastigote Stages with GenMAPP and MAPPFinder* (poster)**Mitchell Petredis***, Kam D. Dahlquist, John David N. Dionisio, *Gene Database Construction and GenMAPP Analysis of Sinorhizobium meliloti Microarray Data Comparing Salt and Sucrose Stress* (poster)

Andrew Pita*, Kam D. Dahlquist, John David N. Dionisio, *Constructing a GenMAPP-compatible Gene Database for Streptococcus pneumoniae to perform pathway analysis on microarray data comparing biofilm versus planktonic forms* (talk)

Britain Southwick*, **Nicole Anguiano***, Kam D. Dahlquist, John David N. Dionisio, Ben G. Fitzpatrick, *GRNsight: a Web Application for Visualizing Models of Gene Regulatory Networks* (talk)

Joint Mathematics Meetings

Baltimore, Maryland, January 2014

Katrina Sherbina*, Kam D. Dahlquist, Ben G. Fitzpatrick, *Dynamical Systems Modeling of the Cold Shock Response in Saccharomyces cerevisiae* (poster)

Katrina was given an "Outstanding Presentation" Award by the Mathematical Association of America for this poster.

Beta Beta Beta Pacific District Convention

Azusa Pacific University, Azusa, California, April 2013

Katrina Sherbina*, Kam D. Dahlquist, Ben G. Fitzpatrick, *Dynamical Systems Modeling of the Cold Shock Response in Saccharomyces cerevisiae* (talk).

Katrina was awarded first place for the Frank G. Brooks Award for Excellence in Student Research for this talk.

Nicholas A. Rohacz*, Kam D. Dahlquist, Ben G. Fitzpatrick. *Continuous Time Markov Chain Models of Gene Regulation Regulatory Networks under the Environmental Stress of Cold Shock in Saccharomyces cerevisiae* (talk).

Nicholas was awarded second place for the Frank G. Brooks Award for Excellence in Student Research for this talk.

5th Annual Undergraduate Research Symposium

Loyola Marymount University, March 2013

Nicolette Harmon*, Chidinma Amakiri*, Nicholas A. Rohacz*, Katrina Sherbina*, Kam D. Dahlquist, Ben G. Fitzpatrick, *A wild species of budding yeast, Saccharomyces paradoxus, is more resistant to cold temperature stress than the domesticated species, Saccharomyces cerevisiae* (talk)

Nicholas A. Rohacz*, Katrina Sherbina*, Kam D. Dahlquist, Ben G. Fitzpatrick, *Continuous Time Markov Chain Models of Gene Regulation Regulatory Networks under the Environmental Stress of Cold Shock in Saccharomyces cerevisiae* (talk)

Katrina Sherbina*, Nicholas A. Rohacz*, Kam D. Dahlquist, Ben G. Fitzpatrick, *Dynamical Systems Modeling of the Cold Shock Response in Saccharomyces cerevisiae* (talk)

Southern California Conference for Undergraduate Research

California State University, Channel Islands, Camarillo, California, November 2012

Nicolette Harmon*, Chidinma Amakiri*, Nicholas A. Rohacz*, Katrina Sherbina*, Kam D. Dahlquist, Ben G. Fitzpatrick, *A wild species of budding yeast, Saccharomyces paradoxus, is more resistant to cold temperature stress than the domesticated species, Saccharomyces cerevisiae* (poster)

Katrina Sherbina*, Nicholas A. Rohacz*, Kam D. Dahlquist, Ben G. Fitzpatrick, *Dynamical Systems Modeling of the Cold Shock Response in Saccharomyces cerevisiae* (poster)

Society for Mathematical Biology Annual Meeting

Knoxville, Tennessee, July 2012

Katrina Sherbina*, Nicholas A. Rohacz*, Kam D. Dahlquist, Ben G. Fitzpatrick, *Dynamical Systems Modeling of the Cold Shock Response in Saccharomyces cerevisiae* (poster)

Nicholas A. Rohacz*, Katrina Sherbina*, Kam D. Dahlquist, Ben G. Fitzpatrick, *Continuous Time Markov Chain Models of Gene Regulation Regulatory Networks under the Environmental Stress of Cold Shock in Saccharomyces cerevisiae* (poster)

West Coast Biological Sciences Undergraduate Research Conference

Loyola Marymount University, April 2012

Nicholas Rohacz*, **Katrina Sherbina***, Kam D. Dahlquist, Ben G. Fitzpatrick, *Mathematical Analysis of Gene Regulation in Saccharomyces cerevisiae in Response to Cold Shock* (poster)

Andrew Herman*, Kam D. Dahlquist

Saccharomyces cerevisiae Responds to Cold Shock by Changing the Expression of Genes

Involved in Nitrogen Metabolism (poster)

4th Annual Undergraduate Research Symposium

Loyola Marymount University, March 2012

Nicholas Rohacz*, **Katrina Sherbina***, Kam D. Dahlquist, Ben G. Fitzpatrick, *Mathematical Analysis of Gene Regulation in Saccharomyces cerevisiae in Response to Cold Shock* (poster)

Andrew Herman*, Kam D. Dahlquist

Saccharomyces cerevisiae Responds to Cold Shock by Changing the Expression of Genes

Involved in Nitrogen Metabolism (poster)

2nd Annual Southern California Systems Biology Conference

University of California, Irvine, January 2012

Nicholas Rohacz*, **Katrina Sherbina***, Kam D. Dahlquist, Ben G. Fitzpatrick, *Mathematical Analysis of Gene Regulation in Saccharomyces cerevisiae in Response to Cold Shock* (poster)

Southern California Conference for Undergraduate Research

Mt. San Antonio College, Walnut, California, November 2011

Cybele Arsan*, Kam D. Dahlquist, *The Hmo1 Transcription Factor Regulates the Expression of Ribosome Biogenesis Genes during Cold Shock and Recovery in Saccharomyces cerevisiae* (talk)

Richard Brous*, Kam D. Dahlquist, John David N. Dionisio

Implementing Multiple Species Export in XMLPipeDB's GenMAPP Builder (talk)

Andrew Herman*, Kam D. Dahlquist, *Saccharomyces cerevisiae Responds to Cold Shock by Changing the Expression of Genes Involved in Nitrogen Metabolism* (poster)

Nicholas Rohacz*, **Katrina Sherbina***, Kam D. Dahlquist, Ben G. Fitzpatrick, *Mathematical Analysis of Gene Regulation in Saccharomyces cerevisiae in Response to Cold Shock* (poster)

Beta Beta Pacific District Convention

Azusa Pacific University, Azusa Pacific, California, April 2011

Cybele Arsan*, Andrew F. Herman*, Alondra J. Vega*, Lauren N. Kubeck*, Kam D. Dahlquist. *The Hmo1 transcription factor regulates the expression of ribosome biogenesis genes during cold shock and recovery in Saccharomyces cerevisiae.* (poster). **Cybele was given the second place**

John C. Johnson Award for Excellence in Student Research for posters in Microbiology.

Andrew F. Herman*, Alondra J. Vega*, Lauren N. Kubeck*, Kenny R. Rodriguez*, Kam D. Dahlquist, *Saccharomyces cerevisiae Responds to Cold Shock by Changing the Expression of Genes Involved in Nitrogen Metabolism* (poster).

Andrew was given the second place John C. Johnson Award for Excellence in Student Research for posters in Physiology or Molecular Biology.

Kelly C. Parks*, Kam D. Dahlquist, John David N. Dionisio.

Using XMLPipeDB to Create a GenMAPP-compatible Gene Database for the Analysis of DNA Microarray Data for Staphylococcus aureus MRSA252 (talk)

3rd Annual Undergraduate Research Symposium

Loyola Marymount University, March 2011

Cybele Arsan*, Andrew F. Herman*, Alondra J. Vega*, Lauren N. Kubeck*, Kam D. Dahlquist, *The Hmo1 transcription factor regulates the expression of ribosome biogenesis genes during cold shock and recovery in Saccharomyces cerevisiae.* (poster).

Andrew F. Herman*, Alondra J. Vega*, Lauren N. Kubeck*, Kenny R. Rodriguez*, Kam D. Dahlquist, *Saccharomyces cerevisiae Responds to Cold Shock by Changing the Expression of Genes Involved in Nitrogen Metabolism* (poster).

Kelly C. Parks*, Kam D. Dahlquist, John David N. Dionisio, *Using XMLPipeDB to Create a GenMAPP-compatible Gene Database for the Analysis of DNA Microarray Data for Staphylococcus aureus MRSA252* (talk)

Don B. Murphy*, Kam D. Dahlquist, John David N. Dionisio, *Implementing Support for Multiple Species in XMLPipeDB's GenMAPP Builder* (poster)

Southern California Conference for Undergraduate Research

Pepperdine University, Malibu, California, November 2010

Andrew Herman*, Alondra J. Vega*, Lauren N. Kubeck*, Kenny R. Rodriguez*, Kam D. Dahlquist, *Saccharomyces cerevisiae Responds to Cold Shock by Changing the Expression of Genes Involved in Nitrogen Metabolism* (talk)

Kelly C. Parks*, Kam D. Dahlquist, John David N. Dionisio, *Using XMLPipeDB to Create a GenMAPP-compatible Gene Database for the Analysis of DNA Microarray Data for Staphylococcus aureus MRSA252* (talk)

Don B. Murphy*, Kam D. Dahlquist, John David N. Dionisio, *Implementing Support for Multiple Species in XMLPipeDB's GenMAPP Builder* (poster)

Society for the Advancement of Chicanos and Native Americans in Science National Conference

Anaheim, California, October 2010

Alondra J. Vega*, Andrew F. Herman*, Lauren N. Kubeck*, Kam D. Dahlquist, and Ben G. Fitzpatrick, *Mathematical Modeling of the Gene Regulatory Network Controlling the Cold Shock Response in Saccharomyces cerevisiae* (poster)

Kevin Paiz-Ramirez*, Kam D. Dahlquist, John David N. Dionisio, *Using XMLPipeDB to Create a GenMAPP-compatible Gene Database for the Analysis of DNA Microarray Data for Mycobacterium tuberculosis* (poster)

Experimental Biology 2010

Anaheim, California, April 2010, published abstracts in *The FASEB Journal* 24(1) Supplement

Kristen Buckmelter*, Bianca Infanzon*, Elizabeth M. Liu*, Olivia S. Sakhon*, Kenny R. Rodriguez*, Wesley T. Citti*, *Saccharomyces cerevisiae responds to cold shock by inducing the transcription of genes required for zinc ion homeostasis* (poster)

Bianca Infanzon*, Kristen Buckmelter*, Elizabeth M. Liu*, Olivia S. Sakhon*, Kenny R. Rodriguez*, Wesley T. Citti*, Kam D. Dahlquist, *Saccharomyces cerevisiae responds to cold shock by inducing the transcription of ribosome biogenesis genes* (poster)

Lauren N. Kubeck*, **Andrew F. Herman***, Kenny R. Rodriguez*, Kevin C. Entzminger*, Stephanie D. Kuelbs*, Kristine B. Hubbard*, Kam D. Dahlquist, *Phenotypic and Functional Genomic Analysis of Heat and Cold Stress in Transcription Factor Deletion Strains of Saccharomyces cerevisiae* (poster)

Bernadette Pak*, Don Murphy*, Kam D. Dahlquist, John David N. Dionisio, *Extending XMLPipeDB with GO Slim to Update the GenMAPP-compatible Gene Database for Budding Yeast, Saccharomyces cerevisiae, for the Analysis of DNA Microarray Data* (poster)

Kelly C. Parks*, Andrew J. Hirning*, **Kelia McDonald***, John David N. Dionisio, Kam D. Dahlquist, *Extending XMLPipeDB to Create a GenMAPP-compatible Gene Databases for the Analysis of DNA Microarray Data from human pathogens* (poster)

Stephen Speicher*, Kam D. Dahlquist, *Gene Ontology Term Enrichment Analysis of Gene Expression Changes Observed in the TRAMP Mouse Model of Prostate Cancer upon Treatment with Green Tea Catechins* (poster)

Alondra J. Vega*, Stephanie D. Kuelbs*, Ben G. Fitzpatrick, Kam D. Dahlquist, *Mathematical Modeling of the Gene Regulatory Network Controlling the Cold Shock Response in Saccharomyces cerevisiae* (talk)

Alondra J. Vega was awarded an NSF/ASBMB Travel Fellowship so that she could make this platform presentation

2nd Annual Undergraduate Research Symposium

Loyola Marymount University, March 2010

Kristen Buckmelter*, **Bianca Infanzon***, *Saccharomyces cerevisiae responds to cold shock by inducing the transcription of genes required for ribosome biogenesis and zinc ion homeostasis* (talk)

Lauren N. Kubeck*, **Andrew F. Herman***, Kam D. Dahlquist, *Phenotypic and Functional*

Genomic Analysis of Heat and Cold Stresses in Transcription Factor Deletion Strains of Saccharomyces cerevisiae" (poster)

Kelia McDonald*, Kam D. Dahlquist, John David N. Dionisio, *Using XMLPipeDB to Create a GenMAPP-compatible Gene Database for Pseudomonas aeruginosa for the Analysis of DNA Microarray Data* (poster)

Bernadette Pak*, **Don Murphy***, Kam D. Dahlquist, John David N. Dionisio, *Extending XMLPipeDB with GO Slim to Update the GenMAPP-compatible Gene Database for Budding Yeast, Saccharomyces cerevisiae, for the Analysis of DNA Microarray Data* (poster)

Kelly Parks*, Kam D. Dahlquist, John David N. Dionisio, *Using XMLPipeDB to Create a GenMAPP-compatible Gene Database for the Analysis of DNA Microarray Data for Staphylococcus aureus MRSA252* (poster)

Alondra J. Vega*, *Mathematical Modeling of the Gene Regulatory Network Controlling the Cold Shock Response in Saccharomyces cerevisiae* (talk)

Stephen Speicher*, *Gene Ontology Term Enrichment Analysis of Gene Expression Changes Observed in the TRAMP Mouse Model of Prostate Cancer upon Treatment with Green Tea Catechins* (talk)

Stephen Speicher won a Sigma Xi award for this presentation

Society for the Advancement of Chicanos and Native Americans in Science National Conference
Dallas, TX, October 2009

Kenny R. Rodriguez*, Kevin C. Entzminger*, Stephanie D. Kuelbs*, Kam D. Dahlquist, *Does Cin5p Regulate the Early Transcriptional Response to Cold Shock in Saccharomyces cerevisiae?* (poster)

Society for Mathematical Biology Annual Meeting
Vancouver, British Columbia, Canada, July 2009

Kenny R. Rodriguez*, Kevin C. Entzminger*, Stephanie D. Kuelbs*, Kam D. Dahlquist, *Does Cin5p Regulate the Early Transcriptional Response to Cold Shock in Saccharomyces cerevisiae?* (poster)

West Coast Biological Sciences Undergraduate Research Conference
Point Loma Nazarene University, San Diego, California, April 2009

Kara Taylor*, Wesley T. Citti*, Jeffrey D. McGowan*, Kam D. Dahlquist, Carl R. Urbinati, *Characterizing Soil Microbial Diversity in the Ballona Wetlands* (talk)

Kevin C. Entzminger*, **Kenny R. Rodriguez***, Stephanie D. Kuelbs*, Kam D. Dahlquist, *Does Cin5p Regulate the Early Transcriptional Response to Cold Shock in Saccharomyces cerevisiae?* (talk)

Alexandrea Alphonso*, **Chad Villaflores***, Derek Smith*, Kam D. Dahlquist, John David N. Dionisio, *Extending XMLPipeDB to Create GenMAPP-compatible Gene Databases for Plants and Microorganisms for the Analysis of DNA Microarray Data* (talk)

Kristine B. Hubbard*, Kenny R. Rodriguez, Stephanie D. Kuelbs, Kam D. Dahlquist, *Phenotypic and Functional Genomic Analysis of Heat and Cold Stresses in Transcription Factor Deletion Strains of Saccharomyces cerevisiae* (poster)

First Annual Undergraduate Research Symposium: Foundations for the Future
Loyola Marymount University, March 2009

Kara Taylor*, Wesley T. Citti*, Jeffrey D. McGowan*, Kam D. Dahlquist, Carl R. Urbinati, *Characterizing Soil Microbial Diversity in the Ballona Wetlands* (talk)

Kevin C. Entzminger*, *Does Cin5p Regulate the Early Transcriptional Response to Cold Shock in Saccharomyces cerevisiae?* (talk)

Kevin C. Entzminger won a Sigma Xi award for this presentation

Alexandrea Alphonso*, **Chad Villaflores***, Derek Smith*, Kam D. Dahlquist, John David N. Dionisio, *Extending XMLPipeDB to Create GenMAPP-compatible Gene Databases for Plants and Microorganisms for the Analysis of DNA Microarray Data* (poster)

Kenny R. Rodriguez*, Stephanie D. Kuelbs*, Kam D. Dahlquist, *Phenotypic and Functional Genomic Analysis of Heat and Cold Stresses in Transcription Factor Deletion Strains of*

Saccharomyces cerevisiae” (poster)

Stephanie D. Kuelbs*, *Mathematical Modeling of the Transcriptional Network Controlling the Cold Shock Response in Saccharomyces cerevisiae* (talk)

First RECOMB Satellite Conference on Bioinformatics Education

San Diego, California, March 2009

Alexandrea Alphonso*, **Chad Villaflores***, Derek Smith*, Kam D. Dahlquist, John David N. Dionisio, *Extending XMLPipeDB to Create GenMAPP-compatible Gene Databases for Plants and Microorganisms for the Analysis of DNA Microarray Data* (poster)

Sigma Xi Annual Meeting

Washington, D.C., November 2008

Kara Taylor*, Wesley T. Citti*, Jeffrey D. McGowan*, Kam D. Dahlquist, Carl R. Urbinati, *Identifying Soil Bacterial and Biochemical Pathways in the Ballona Wetlands* (poster)

Society for the Advancement of Chicanos and Native Americans in Science National Conference

Salt Lake City, Utah, October 2008

Kenny R. Rodriguez*, Kevin C. Entzminger*, Stephanie D. Kuelbs*, Kam D. Dahlquist, *Phenotypic and Functional Genomic Analysis of Heat and Cold Stress in Transcription Factor Deletion Strains of Saccharomyces cerevisiae* (poster)

Society for Mathematical Biology Annual Meeting

Toronto, Ontario, Canada, August 2008

Stephanie D. Kuelbs*, Kevin C. Entzminger*, Kenny R. Rodriguez*, Ben G. Fitzpatrick, Kam D. Dahlquist, *Mathematical Modeling of the Transcriptional Network Controlling the Cold Shock Response in Saccharomyces cerevisiae* (poster)

Yeast Genetics and Molecular Biology

Toronto, Ontario, Canada, July 2008

Kevin C. Entzminger*, Kenny R. Rodriguez*, Stephanie D. Kuelbs*, Kam D. Dahlquist, *Does Cin5p Regulate the Early Transcriptional Response to Cold Shock in Saccharomyces cerevisiae?* (poster)

West Coast Biological Sciences Undergraduate Research Conference

Point Loma Nazarene University, San Diego, California, April 2008

Wesley T. Citti*, **Jeffrey D. McGowan***, Kam D. Dahlquist, Carl R. Urbinati, *Identification and Diversity Analysis of Soil Bacteria in the Ballona Wetlands* (talk)

Elizabeth M. Liu*, **Olivia S. Sakhon***, **Robert Hybki***, Kam D. Dahlquist, *The Global Transcriptional Response of Saccharomyces cerevisiae to Cold Shock and Recovery* (talk)

Kenny R. Rodriguez*, **Kevin C. Entzminger***, **Stephanie D. Kuelbs***, Kam D. Dahlquist, *Does the Transcription Factor CIN5 Regulate the Transcriptional Response to Cold Shock in Saccharomyces cerevisiae?* (poster)

Pacific Coast Undergraduate Math Conference

Loyola Marymount University, Los Angeles, California, April 2008

Stephanie D. Kuelbs*, *Mathematical Modeling of the Transcriptional Network Controlling the Cold Shock Response in Saccharomyces cerevisiae* (talk)

Southern California Conference for Undergraduate Research

California State University, Los Angeles, November 2007

Wesley T. Citti*, **Jeffrey D. McGowan***, Kam D. Dahlquist, Carl R. Urbinati, *Identifying Soil Bacteria and Biochemical Pathways for Bioremediation in Ballona Wetlands* (poster)

Elizabeth M. Liu*, **Olivia S. Sakhon***, **Robert Hybki***, Kam D. Dahlquist, *The Global Transcriptional Response of Saccharomyces cerevisiae to Cold Shock and Recovery* (poster)

Kevin C. Entzminger*, **Stephanie D. Kuelbs***, **Kenny R. Rodriguez***, Kam D. Dahlquist, *Mathematical Modeling and Biological Analysis of the Transcriptional Response to Cold Shock in Saccharomyces cerevisiae* (poster)

Interdisciplinary Student Research Symposium

Loyola Marymount University, Los Angeles, California, October 2007

Wesley T. Citti*, **Jeffrey D. McGowan***, Kam D. Dahlquist, Carl R. Urbinati, *Identifying Soil*

Bacteria and Biochemical Pathways for Bioremediation in Ballona Wetlands (poster)

Kevin C. Entzminger*, **Stephanie D. Kuelbs***, **Kenny R. Rodriguez***, Kam D. Dahlquist, *Mathematical Modeling and Biological Analysis of the Transcriptional Response to Cold Shock in Saccharomyces cerevisiae* (poster)

Annual Meeting of the Society for Mathematical Biology

San Jose, California, August 2007

Nathan C. Wanner*, Erika Camacho, Kam D. Dahlquist, *Mathematical Modeling of the Transcriptional Network Controlling the Environmental Stress Response in Saccharomyces cerevisiae* (poster)

West Coast Biological Sciences Undergraduate Research Conference

Loyola Marymount University, Los Angeles, California, April 2007

Wesley T. Citti*, Kam D. Dahlquist, Carl R. Urbinati, *Identifying Bacteria and Biochemical Pathways for Bioremediation in Ballona Wetlands* (poster)

Elizabeth M. Liu*, **Olivia S. Sakhon***, Kam D. Dahlquist, *The Global Transcriptional Response of Saccharomyces cerevisiae to Cold Shock and Recovery* (poster)

Sigma Xi Induction Ceremony and Poster Session

Loyola Marymount University, Los Angeles, California, April 2007

Wesley T. Citti*, Kam D. Dahlquist, Carl R. Urbinati, *Identifying Bacteria and Biochemical Pathways for Bioremediation in Ballona Wetlands* (poster)

Elizabeth M. Liu*, **Olivia S. Sakhon***, Kam D. Dahlquist, *The Global Transcriptional Response of Saccharomyces cerevisiae to Cold Shock and Recovery* (poster)

San Diego Systems Biology Symposium: Systems to Synthesis

Salk Institute, La Jolla, California, January 2007

Nathan C. Wanner*, Erika Camacho, Kam D. Dahlquist, *Mathematical Modeling of the Transcriptional Network Controlling the Environmental Stress Response in Saccharomyces cerevisiae* (poster)

Nathan C. Wanner won the third place poster prize at this symposium.

Southern California Conference for Undergraduate Research

Occidental College, Los Angeles, California, November 2006

Wesley T. Citti*, Kam D. Dahlquist, Carl R. Urbinati, *Identifying Bacteria and Biochemical Pathways for Bioremediation in Ballona Wetlands* (poster)

Bellarmino Forum on Environmental Responsibility

Loyola Marymount University, Los Angeles, California, November 2006

Wesley T. Citti*, Kam D. Dahlquist, Carl R. Urbinati, *Identifying Bacteria and Biochemical Pathways for Bioremediation in Ballona Wetlands* (poster)

West Coast Biological Sciences Undergraduate Research Conference

Point Loma Nazarene University, San Diego, California, April 2006

Wesley T. Citti*, Heather King*, and Kam D. Dahlquist, *The Transcriptional Response of Saccharomyces cerevisiae to Cold Shock and Recovery* (poster)

Wesley T. Citti won a poster award at this conference.

2004 Undergraduate Research Summer Institute Symposium

Vassar College, Poughkeepsie, New York, September 2004

Meredith Braymer*, Eric S. Eberhardt, Kam D. Dahlquist, *Global Changes in Gene Expression during Cold Shock and Recovery in Saccharomyces cerevisiae* (poster)

Jessica Heckman* and Kam D. Dahlquist, *New Resources for GenMAPP 2.0: A New Gene Database and Pathway MAPPs for the Comparison of Changes in Gene Expression due to Environmental Stresses in Saccharomyces cerevisiae and Escherichia coli* (poster)

Nikoleta Tsvetanova*, Meredith Braymer*, Eric S. Eberhardt, *Cold-Shock Response in Saccharomyces cerevisiae* (poster)

SERVICE & PROFESSIONAL INVOLVEMENT

Internal**Loyola Marymount University****University-wide**

Honors Program Faculty Fellow	2014–2020
Mission Day Planning Committee	2015–2016
Search Committee for Dean of the Seaver College of Science and Engineering	2014–2015
Library Committee	2013–2015
LMU Undergraduate Research Symposium Session Chair	2013, 2014, 2017, 2018, 2022, 2023
Performed Assessment of LMU's Oral Communication Learning Outcome	2013
Advisory Committee on Undergraduate Research	2013
Digital Scholarship Repository Project Team	2010–2011
High Performance Computing Task Force	2010–2011
Research Council	2009–2015
Valedictorian Committee	2009, 2011
Scholarship of Teaching and Learning Brown Bag Group	2005–2011
Interviewer of candidates for Director of Sponsored Projects Office	Summer 2008

Frank R. Seaver College of Science and Engineering

Health Professions Advising Committee	2022–2024
Chair	2024
Seaver Leadership Council	2020–2023
Seaver Rank & Tenure Standards Normalization Committee	2020–2021
College Equipment and Computing Committee	2019–2020
College Curriculum Committee	2017–2019
Breaking the Boundaries in STEM Education Research Conference	April 2017
Computational Thinking Thread Co-Chair	
4-Unit Task Force	2015–2016
Prioritization Committee	2013–2014
Pre-tenure Faculty Guidance Committee	2010–2011
Information Technology Committee	2005, 2009–2010
Search Committee for Presidential Professorship in Computational Biology	2008–2010
Search Committee for Presidential Professorship in Mathematical Biology	2006–2008

Department of Biology

Seminar Coordinator	2023–2025
Chair	2020–2023
Curriculum Committee, Chair	2017–2020
Search Committee for Animal Physiologist, Chair	2017–2018
TriBeta Biology Honor Society Advisor	2016–present
Laboratory Safety Committee, Chair	2010, 2016–2020
Reviewer, Kadner-Pitts Research Grants	2013, 2016, 2018
4-unit Curriculum Model Task Force	2014–2015, 2017–2019
Search Committee for Biochemist/Cell Physiologist	2010–2011
Faculty mentor	2009–2020
Search Committee for Vertebrate Physiologist	2009–2010
APRC Review Committee	2006–2011
Chair	2011
Webmaster for Department website	2006–2012
Review of Faculty Research Funds subcommittee	2006–2008
Sensitive Equipment subcommittee	Spring 2006

Vassar College

Women in Science and Mathematics Faculty Adviser	2003–2005
Career Development Office Advisory Committee	2003–2005
Carolyn Grant Endowment Committee for Embodied Learning	2004–2005
Biology Department Curriculum Committee	2004–2005
Biology Department Community Committee	2004–2005
Women's Studies Steering Committee	2004–2005

External

Argo Navis School, Playa Vista, California	2025
Led Cahuita, Costa Rica Field Guide Project for middle school students	
Centro Educativo Complementaria Cahuita, Costa Rica	2025
Taught PCR for DNA barcoding to middle school students	
Project Scientist Summer STEM Academy for Girls, Superstar (Guest Speaker)	2019
West Coast Biological Sciences Undergraduate Research Conference	
Chair, LMU Organizing Committee	2020
Member, LMU Organizing Committee	2007
Oral Session Chair	2019
Presentation or Poster Judge	2006, 2008, 2015, 2019
Beta Beta Beta Pacific District Convention	2018
Oral Session Judge	
Southern California Conference for Undergraduate Research	
Abstract Reviewer and Session Chair	2014
Intel International Science and Engineering Fair	
Sigma Xi Special Awards Judge, Los Angeles, California	2011
Open Bioinformatics Foundation	
Abstract Reviewer, Bioinformatics Open Source Conference (Boston)	2011–2014
Codefest Host, Loyola Marymount University	2012
At-large Member, Board of Directors	2008–2010
Chair, Bioinformatics Open Source Conference (Boston)	July 9–10, 2010
Chair, Bioinformatics Open Source Conference (Stockholm)	June 27–28, 2009
Chair, Bioinformatics Open Source Conference (Toronto)	July 18–19, 2008
International Society for Computational Biology	
Member, Education Committee	2006–2015
Genome Consortium for Active Teaching (GCAT)	
Alternate scanning center for DNA microarrays	2010–2015
Grants and Publishing	
Review Panel, National Science Foundation	2009, 2011, 2014, 2015, 2022
Peer-reviewer	
<i>Bulletin of Mathematical Biology</i>	2023, 2025
<i>Database</i>	2023
<i>Applied Microbiology and Biotechnology</i>	2022
<i>International Journal of STEM Education</i>	2019
<i>GigaScience</i>	2018
<i>Journal of Research in STEM Education</i>	2018
<i>PLoS ONE</i>	2009, 2017
<i>PLoS Computational Biology</i>	2016
<i>Reinvention: an International Journal of Undergraduate Research</i>	2015–2016
<i>Nucleic Acids Research</i>	2015
<i>Journal of Computational Science Education</i>	2011
<i>CBE – Life Sciences Education</i>	2003, 2006, 2008, 2009
<i>Bioinformatics</i>	2003, 2009
<i>EURASIP Journal on Advances in Signal Processing</i>	2009
<i>Briefings in Functional Genomics and Proteomics</i>	2008

<i>Molecular and Cellular Proteomics</i>	2004
Chapter Reviewer , Watson et al., <i>Recombinant DNA</i> , 3 rd edition	2006
Association for Women in Science (AWIS)	
Chair, Programs Committee, Palo Alto Chapter	2001–2003
<ul style="list-style-type: none"> Organized and led monthly chapter meetings attended by 50-75 people Invited speakers (women scientists, career development) 	
Postdoctoral Women Peer-mentoring Group , U.C. San Francisco	2001–2003
Alumni Volunteer Admissions Interviewer , Pomona College	1995–1998, 2001
Phoenix II Seminars , San Jose, California	
Graduate, Leadership Program	1994
Staff volunteer for courses and exit interviews	1994–1995
<u>Memberships</u>	
Genetics Society of America	2024–present
American Society for Biochemistry and Molecular Biology	2009–present
Open Bioinformatics Foundation	2006–present
American Society for Cell Biology	2003–2015
International Society for Computational Biology	2002–2016, 2025
Association for Women in Science (AWIS)	1998–2015
American Association for the Advancement of Science	1995–2017

PROFESSIONAL DEVELOPMENT WORKSHOPS ATTENDED

UCLA QCBio Collaboratory Interactive Workshops	2025
Intro to Python, UNIX Command Line I, Using NGS Analysis Tools, Library Prep for NGS, Protein Structure with AlphaFold, RNA-seq I Analysis, RNA-seq II Analysis, Intro to R and Data Visualization, Intro to Modern Statistics with R	
Seaver College HHMI STEM Equity Faculty Learning Community	2023–2024
Loyola Marymount University eFaculty Training for Online Courses	May 2020
BioQUEST Summer Workshop 2018	June 2018
Wicked Problems: Investigating Real World Problems in the Biology Classroom	
Harvey Mudd College, Claremont, California, June 2018 (with Carrie Diaz Eaton)	
BioQUEST / HHMI / CaseNet Summer Workshop 2017	July 2017
Making Meaning Through Modeling: Problem Solving in Biology	
Michigan State University, East Lansing, Michigan	
Breaking the Boundaries in STEM Education Research Conference	April 2017
Loyola Marymount University, Los Angeles, California	
GCAT-SEEK Workshop	June-July 2016
California State University at Los Angeles, Los Angeles, California	
BioQUEST / HHMI / CaseNet Summer Workshop 2015	June 2015
Count the Ways: Engaging Students in Quantitative Biology Applications	
Harvey Mudd College, Claremont, California,	
Loyola Marymount University President's Institute	May 2009, May 2013
BioQUEST Curriculum Consortium Summer Workshop 2011	June 2011
Undergraduate Biology in the 21st Century , Beloit College, Beloit, Wisconsin	
Peer Evaluation of Teaching Workshop	May 2011
Center for Teaching Excellence, Loyola Marymount University, Los Angeles, California	
BioQUEST Curriculum Consortium Summer Workshop 2009	June 2009
Green Architecture – Green Curriculum , Beloit College, Beloit, Wisconsin	
BioQUEST Curriculum Consortium Summer Workshop 2007:	June 2007
Exploratory Evolution Education , Beloit College, Beloit, Wisconsin	
Women in bioScience Conference	May 2007
Association for Women in Science, San Diego, California	

Pedagogy Workshop for Second-year Faculty Loyola Marymount University, Los Angeles, California	2006–2007
Jesuit and Feminist Education: Transformative Discourses for Teaching & Learning Conference Fairfield University, Fairfield, Connecticut	October 2006
Collegium: A Colloquy on Faith and Intellectual Life St. John's University, Collegeville, Minnesota	June 2006
BioQUEST Curriculum Consortium Summer Workshop 2005: Investigating Interdisciplinary Interactions , Beloit College, Beloit, Wisconsin (attended with Erika Camacho who was then in the Department of Mathematics at LMU)	June 2005
The Embodied Voice Faculty Workshop Vassar College, Poughkeepsie, New York	Spring 2005
Dartmouth Faculty Summer Institute Ethical, Legal, and Social Implications of the Human Genome Project Dartmouth University, Hanover, New Hampshire	July 2004
BEDROCK Workshop–Bioinformatics in Biology Education: Working with Sequence, Structure, and Function Cornell Theory Center, Ithaca, New York	October 2003
Analysis of Regulatory Sequences Controlling Expression of Biological Networks; Extracting Biological Information from System-scale Protein Interactome Data Intelligent Systems for Molecular Biology Tutorials, Brisbane, Queensland, Australia	June 2003
Strategies in Gender Equitable Teaching U.C. Berkeley Extension, Berkeley, California	2001–2002
Beginning Dreamweaver 4 Ciber Training Center, San Francisco, California	April 2002
Advanced Microsoft Access 97 New Horizons Computer Learning Centers, Inc., San Francisco, California	August 2001
Biostatistics 183: Introduction to Statistical Analysis U.C. San Francisco, San Francisco, California	Fall 2001
Art of Lecturing Gladstone Institutes, San Francisco, California	Summer 2001
Scientific Writing Gladstone Institutes, San Francisco, California	Spring 2001
Gladstone Genomics Core Microarray Academy	Fall 2000