Kam D. Dahlquist, Ph.D.

Associate Professor Department of Biology Loyola Marymount University 1 LMU Drive, MS 8220 Los Angeles, CA 90045-2659 Tel: 310-338-7697 Fax: 310-338-4479 kdahlquist@lmu.edu http://myweb.lmu.edu/kdahlqui/ Last modified: November 25, 2009

EDUCATION

Ph.D. University of California, Santa Cruz

March 2000

Molecular, Cellular, and Developmental Biology Program

Advisor: Joseph D. Puglisi, Ph.D.

Committee: Harry F. Noller, Ph.D., Manuel Ares, Jr., Ph.D. Thesis: Interaction of Translation Initiation Factor IF1

with the E. coli Ribosomal A Site

B.A. Pomona College, Claremont, California

May 1993

Biology, cum laude

University College, Oxford University, Oxford, England

Fall 1991

Study Abroad Program

Specialized tutorial in Philosophy of Science

POSITIONS HELD

Associate Professor
Assistant Professor
2009–present
2005–2009

Department of Biology, Loyola Marymount University, Los Angeles, California

Assistant Professor 2003–2005

Department of Biology, Vassar College, Poughkeepsie, New York

Postdoctoral Fellow 2000–2003

Gladstone Institute of Cardiovascular Disease, University of California, San Francisco

Adjunct Lecturer Spring 2000

Department of Biology, Santa Clara University, Santa Clara, California

Visiting Researcher 1997–2000

Department of Structural Biology, Stanford University, Stanford, California

Research Assistant 1994–1997

Department of Biology, University of California, Santa Cruz, California

GRANTS, FELLOWSHIPS, HONORS, AWARDS

NSF-DMS Mathematical Biology, MCB Genes and Genome Systems

2009-2012

\$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

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

Kadner-Pitts Research Grant

2007–2008

Department of Biology, Loyola Marymount University

\$10,000, Mapping Gene Regulatory Networks in Yeast using DNA Microarrays, Mathematical Modeling, and GenMAPP

Merck-AAAS Undergraduate Science Research Program

2006-2008

\$120,000 (\$60,000 plus \$60,000 matching funds from Loyola Marymount University)

Chemical and Biological Aspects of Pollution in the Ballona Wetlands

Principal Investigator: M. Catherine McElwain

Director and Co-Principal Investigator: Kam D. Dahlquist

Co-Principal Investigators: Rachel Adams, Lambert Doezema, John Dorsey, Philippa M. Drennan, Gary A. Kuleck, Jim Landry, Jeremy McCallum, David Moffet, Martin G. Ramirez, James Roe, and Carl R. Urbinati

Interdisciplinary Research Project with David Moffet and Carl R. Urbinati entitled Identifying Soil Bacteria and Biochemical Pathways in the Ballona Wetlands for the Bioremediation of Organic Pollutants

W.M. Keck Foundation

2007

\$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

Academic Technology Grant, Loyola Marymount University

2007

\$4,000, Introducing DNA Microarray Technology in the New Laboratory Course, Biology 478: Molecular Biology of the Genome

Dartmouth Faculty Summer Institute Travel Award and Stipend

August 2006

ELSI Reunion and Conference, Dartmouth University, Hanover, New Hampshire

| Summer Research Grant for New Faculty, Loyola Marymount University | 2006 |
|---|--------------|
| \$4,000, The Transcriptional and Proteomic Response to Cold Shock and Rec Saccharomyces cerevisiae | overy in |
| Merck-AAAS Undergraduate Science Research Program Principal Investigator: Richard B. Hemmes, Department of Biology, Vassar Interdisciplinary Research Project with Eric S. Eberhardt, Department of Ch. Vassar College entitled Examining the Molecular Details of Oxidative Stress Genome to the Proteome [I declined my share of the funding upon my move to Loyola Marymount University] | emistry, |
| Mellon Faculty Conversations Award, Vassar College \$2,000, Effective Grading: A Tool for Learning and Assessment | 2004–2005 |
| Dartmouth Faculty Summer Institute Travel Award and Stipend Ethical, Legal, and Social Implications of the Human Genome Project Dartmouth University, Hanover, New Hampshire | July 2004 |
| Sigma Xi, Full Membership Associate Membership | 2004 1992 |
| Carolyn Grant Endowment for Embodied Learning, Vassar College \$2,000, Sponsored a visit by Jean Couch to lead workshops on <i>Balanced Pos</i> for <i>Introduction to Biological Thought: The Human Genome</i> and the campus | |
| 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 West Coast Undergraduate Research Conference in the Biological Sciences | 1992 |

Summer 1993

RESEARCH EXPERIENCE

Associate Professor 2009–present

Department of Biology, Loyola Marymount University, Los Angeles, California *Current Research Projects:*

- Creation of MAPPs and Gene Databases for GenMAPP software;
 XMLPipeDB: A Reusable Open Source Tool Chain for Building Relational Databases from XML Sources
- The Global Transcriptional Response of *Saccharomyces cerevisiae* to Cold Shock and Recovery
- Mathematical Modeling of the Transcriptional Network Controlling the Environmental Stress Response in *Saccharomyces cerevisiae*
- Identifying Soil Bacterial and Biochemical Pathways in the Ballona Wetlands for the Bioremediation of Organic Pollutants

Research advisor for a total of 21 undergraduates and 1 Master's level student at LMU since 2005

Assistant Professor 2005–2009

Department of Biology, Loyola Marymount University, Los Angeles, California **Assistant Professor**2003–2005

Department of Biology, Vassar College, Poughkeepsie, New York *Research Projects:*

- Transcriptional and Proteomic Response of *Saccharomyces cerevisiae* to Cold Shock and Recovery
- Creation of MAPPs and Gene Databases for GenMAPP software

Research advisor for a total of 6 undergraduate students at Vassar 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 *Advisor:* Joseph D. Puglisi, Ph.D.

Research Assistant 1994–1997

Department of Biology, University of California, Santa Cruz

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

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

TEACHING EXPERIENCE

College Level

Associate Professor Assistant Professor

2009–present 2005–2009

Department of Biology, Loyola Marymount University

Biology 201: Cell Function

• Sophomore-level course in four-semester lower division curriculum for biology majors

Biology 275: Human Genetics

• Fulfills University core requirement for non-science majors

Biology 439: Molecular Biology Applications

- Intensive laboratory course in molecular biology
- Students performed semi-independent cloning project based on my dissertation research

Biology 398/Computer Science 398/Honors 398-01: Biological Databases

- Cross-listed and team taught with John David N. Dionisio, Ph.D., Department of Electrical Engineering and Computer Science
- Interdisciplinary student teams created GenMAPP Gene Databases for Plasmodium falciparum, Pseudomonas putida KT2440, Sinorhizobium meliloti, and Vibrio cholerae using XMLPipeDB open source software
- Students maintained online laboratory notebooks using MediaWiki software

Biology 398-03: Bioinformatics Laboratory

- Project-based computer laboratory using GenMAPP, MAPPFinder, and other bioinformatics software
- Projects included sequence and structural analysis of the gp120 protein of HIV and analysis of DNA microarray experiments
- Students maintained online laboratory notebooks using Google Sites.

Biology 478: Molecular Biology of the Genome

- Subject of 2007 LMU Academic Technology Grant
- Intensive laboratory course in molecular biology
- Students perform a DNA microarray experiment
- DNA microarray data analyzed with GenMAPP, MAPPFinder, and other bioinformatics software

Biology 498/Computer Science 698: Special Studies in Bioinformatics

• 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 developed XMLPipeDB software and Gene Database for GenMAPP using open source tools and the development environment of SourceForge

Biology 585: Issues in Biotechnology

- Seminar and capstone experience for biology majors
- Read, present, and discuss articles from the primary literature
- Present and discuss the ethical, legal, and social implications of biotechnology research and scientific misconduct

Honors 240: On the Nature of Things

- University core requirement for students in the Honors Program
- An examination of the history, philosophy, and nature of scientific discovery, theory, and practice
- Focus on recent advances in biology, specifically biotechnology and genomics, epistemology, and genetic determinism
- Seminar-based discussion course with presentations and final projects relating to the students' major field of study

Delivered Guest Lecture in the following courses

- BIOL 114: Biology for Engineers (March 2009)
- PHIL 666: Philosophy of Science (October 2008)
- CMSI 686: Database Systems (April 2007)
- MATH 298: Biomathematics (April 2007)
- MGMT 498: Technology Ventures (March 2006)
- CMSI 486: Introduction to Database Systems (October 2005)

Assistant Professor 2003–2005

Department of Biology, Vassar College

Introduction to Biological Thought: The Human Genome

- Students used MAPPFinder to analysis 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
- Evaluated highly on organization and meeting course goals

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

Adjunct Lecturer Spring 2000

Department of Biology, Santa Clara University *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
- Evaluated highly by students for stimulating independent thinking and for demonstrating an interest in them

Course Assistant Winter 1998

Department of Biology, Stanford University *Cell Biology*

• Led discussion of research articles

Teaching Assistant

Fall 1994

Department of Biology, U.C. Santa Cruz *Concepts in Biology*

• Lectured when professor was out of town

Teaching Assistant

Summer 1994

Howard Hughes Summer Institute, U.C. Santa Cruz *Molecular and Cell Biology Laboratory*

• Supervised semi-independent research projects on the cloning of *frequency* homologues in different species of fungi

K-12

Scientist Volunteer 2001–2002

Herbert Hoover Middle School and U.C. San Francisco Science and Health Education Partnership Triad Science Club

• Developed and led hands-on activities, including gel electrophoresis

Elective Teacher Fall 1994

Mission Hill Junior High School, U.C. Santa Cruz

Project SAME: Science and Math Equity

• Taught a girl-only elective class on building simple machines with the Lego-Logo system

SERVICE & PROFESSIONAL INVOLVEMENT

Internal

Loyola Marymount University

University-wide

Research Council 2009—present Scholarship of Teaching and Learning Working Group 2005—present Interviewer of candidates for Director of Sponsored Projects Office Summer 2008

| Frank R. Seaver College of Science and Engineering | |
|---|----------------------|
| Information Technology Committee | 2005, 2009–present |
| Search Committee for Presidential Professorship | |
| in Computational Biology | 2008–present |
| Search Committee for Presidential Professorship | |
| in Mathematical Biology | 2006–2008 |
| Department of Biology | |
| Coordinator for Biology 201: Cell Function | 2009–present |
| Search Committee for Vertebrate Physiologist | 2009–present |
| APRC Review, Student-centered goals subcommittee | 2006-present |
| Webmaster for Department web site | 2006-present |
| 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 | 2001 2000 |
| West Coast Biological Sciences Undergraduate Research Conference | • |
| Presentation and Poster Judge | 2008 |
| Member, Organizing Committee | 2007 |
| Poster Judge | 2006 |
| 1 Oster Judge | 2000 |
| Open Bioinformatics Foundation | |
| At-large Member, Board of Directors | 2008-present |
| Chair, Bioinformatics Open Source Conference (BOSC) | June 27–28, 2009 |
| Chair, Bioinformatics Open Source Conference (BOSC) | July 18–19, 2008 |
| International Society for Computational Biology | |
| Member, Education Committee | 2006-present |
| Scientific Publications | 1 |
| Associate Editor, International Journal of Computational Bioscienc | e 2009–present |
| Peer-reviewer | |
| CBE – Life Sciences Education 20 | 03, 2006, 2008, 2009 |
| Bioinformatics 20 | 2003, 2009 |
| EURASIP Journal on Advances in Signal Processing | 2009 |
| PLoS ONE | 2009 |
| Briefings in Functional Genomics and Proteomics | 2008 |
| Molecular and Cellular Proteomics | 2004 |
| Chapter Reviewer , Watson et al., <i>Recombinant DNA</i> , 3 rd edition | 2006 |
| complet 220 120 1102, 11 anson of any recommuna Divis, 5 cultion | 2000 |

Association for Women in Science (AWIS)

Chair, Programs Committee, Palo Alto Chapter

2001-2003

- 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 |

Memberships

| Open Bioinformatics Foundation | 2006–present |
|---|--------------|
| American Society for Cell Biology | 2003–present |
| International Society for Computational Biology | 2002–present |
| Association for Women in Science (AWIS) | 1998–present |
| American Association for the Advancement of Science | 1995–present |

PUBLICATIONS

Peer-reviewed Research

- Ogando, D.G., **Dahlquist, K.D.**, Kunchithapautham, K., Yu, N., Alizadeh, M., Li, J., Vollrath, D., LaVail, M.M., Rohrer, B., Danciger, M. (2009) Candidate genes for Chromosomes 6 and 10 quantitative trait loci for age-related retinal degeneration in mice. *Molecular Vision*, *submitted*.
- 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.
- 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.
- Segal, M.R., **Dahlquist, K.D.**, & Conklin, B.R. (2003) Regression Approaches for Microarray Data Analysis. *Journal of Computational Biology* **10**: 961-980.
- 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.
- **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.

- **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.
- 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.
- 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.

Reviews, Book Chapters, Conference Proceedings

- **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.

Software and Databases

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

Co-Principal Investigator with John David N. Dionisio, 2006–present

Availability: http://sourceforge.net/projects/xmlpipedb

Web site: http://xmlpipedb.cs.lmu.edu

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

Project Manager, 2000-2003

Metabolic Pathway MAPP Archive for Saccharomyces cerevisiae, 2005

Gene Database for Escherichia coli K12, 2006, 2009

Metabolic Pathway MAPP Archive for E. coli K12, 2008

Gene Database for Arabidopsis thaliana, 2007, 2009

Gene Database for Plasmodium falciparum, 2009

Gene Database for Vibrio cholerae, 2009

Availability: http://www.GenMAPP.org

PRESENTATIONS

External Talks

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

Pepperdine University

Malibu, California, February 2008

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

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

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

An Open Source Framework for Teaching Bioinformatics

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

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

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

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

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

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

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

Bay Area RNA Club

San Francisco, California, June 1996

Rites of Initiation: Decoding the role of IF1

Internal Talks

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)

Intelligent Systems for Molecular Biology

Stockholm, Sweden, June 2009

with, Alexandrea 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

<u>Student Presentations and Posters</u> (*indicates undergraduate student, **bold** indicates presenting author)

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?

Annual Meeting of the Society for Mathematical Biology

Vancouver, British Columbia, Canada, August 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?

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

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?

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

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*

The First Annual Undergraduate Research Symposium: Foundations for the Future Lovola 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

Kevin C. Entzminger*

Does Cin5p Regulate the Early Transcriptional Response to Cold Shock in Saccharomyces cerevisiae?

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

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

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

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

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*

Annual Meeting of the Society for Mathematical Biology

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

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?

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

Elizabeth M. Liu*, Olivia S. Sakhon*, Robert Hybki*, Kam D. Dahlquist

The Global Transcriptional Response of Saccharomyces cerevisiae to Cold Shock and Recovery

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?

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 Kam D. Dahlquist, Ph.D.

Curriculum Vitae

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

Elizabeth M. Liu*, Olivia S. Sakhon*, Robert Hybki*, Kam D. Dahlquist

The Global Transcriptional Response of Saccharomyces cerevisiae to Cold Shock and Recovery

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

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

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

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

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

Elizabeth M. Liu*, Olivia S. Sakhon*, Kam D. Dahlquist

The Global Transcriptional Response of Saccharomyces cerevisiae to Cold Shock and Recovery

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

Elizabeth M. Liu*, Olivia S. Sakhon*, Kam D. Dahlquist

The Global Transcriptional Response of Saccharomyces cerevisiae to Cold Shock and Recovery

2006–2007

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

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

Bellarmine 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

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

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

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

Nikoleta Tsvetanova*, Meredith Braymer*, Eric S. Eberhardt

Cold-Shock Response in Saccharomyces cerevisiae

PROFESSIONAL DEVELOPMENT WORKSHOPS ATTENDED

President's Institute May 2009 Loyola Marymount University, Los Angeles, California **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

Jesuit and Feminist Education: October 2006

Fall 2000

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

Microarray Academy
Genomics Core, Gladstone Institutes, San Francisco, California

Gladstone Institutes, San Francisco, California