Michael Strong, PhD

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FACULTY

I am an Assistant Professor in the Integrated Center for Genes, Environment, and Health, at **National Jewish Health**. Research interests include genomics, systems biology, structural informatics, and the development of methods to better examine the interplay among humans, human pathogens, and environmental factors. <u>http://www.nationaljewish.org/research/programs-depts/genetics-therapeutics/labs/michael-strong/index.aspx</u>

I am also a core faculty member of the Computational Bioscience Program at the **University of Colorado, Denver**, School of Medicine. <u>http://compbio.ucdenver.edu/pages/corefaculty.html</u>

POSTDOCTORAL

I conducted my postdoctoral work in **George Church**'s lab in the Department of Genetics at **Harvard Medical School**. Research interests include genomics, biochemistry, systems biology, and synthetic biology. (Recipient of a Jane Coffin Childs Foundation Research Grant). <u>http://arep.med.harvard.edu/~strong/</u>

EDUCATION

Ph.D. in Molecular Biology. Graduate Advisor – **David Eisenberg**, **University of California, Los Angeles**. 2005. My research focused on identifying the modular nature of genome-wide protein networks, and developing biochemical tools to identify protein-protein interactions. Research interests include genomics, protein biochemistry, molecular biology, bioinformatics, and structural biology. (recipient of NIH Cell and Molecular Biology Training Grant)

B.S. Microbiology, concentration in Genetic Engineering. University of California, Santa Barbara. (recipient of HHMI undergraduate research grant).

PUBLICATIONS

Translating basic science insight into public health action for MDR- and XDR-TB. Walter, Nicholas; **Strong, Michael**; Belknap, Robert; Ordway, Diane; Daley, Charles L; Chan, Ed; *Respirology*. 17:772-791 (2012).

Research, Collaboration, and Open Science using Web 2.0. Kevin Shee*, **Michael Strong***, Nicholas J. Guido, Robert Lue, George M. Church, Alain Viel. *Equal contribution, Journal of Microbiology and Biology Education, 11:2 p130-134, (2010).

Reducing Global Health Disparities through Research, Education, and International Collaboration. **Michael Strong**. *Center for Strategic and International Studies, Smart Global Health Policy Report p47-48* (2010).

Tuberculosis Drug Resistance Mutation Database . Andreas Sandgren, **Michael Strong**, Preetika Mutukrishnan, Brian K. Weiner, George M. Church, Megan B. Murray. *PLoS Medicine* 6:2 (2009).

Gem of an idea for a unique machine. **Michael Strong** and George Church. *Nature* 447:112 (2007).

The Protein Network as a Tool for Finding Novel Drug Targets. **Michael Strong** and David Eisenberg. Book Chapter in Systems Biological Approaches in Infectious Diseases. Edited by H.I.M Boshoff and C.E. Barry III, Birkhauser. Verlag Publishing. *Progress in Drug Research*, 64:191, 193-215 (2007). Functional linkages can reveal protein complexes for structure determination. Kim SM, Bowers PM, Pal D, **Strong M**, Terwilliger TC, Kaufmann M, Eisenberg D. *Structure* 15:1079-89 (2007).

Toward the Structural Genomics of Complexes: Crystal Structure of a PE/PPE protein complex from *Mycobacterium tuberculosis*.

Michael Strong, Michael Sawaya, Shuishu Wang, Martin Philips, Duilio Cascio, and David Eisenberg. *Proc. Natl. Acad. Sci. U. S. A.* 103:8060-8065 (2006).

Structural Proteomics and Computational Analysis of a Deadly Pathogen: Combating *M. tuberculosis* from Multiple Fronts. **Michael Strong** and Celia Goulding. Book Chapter (Chapter 15) in Microbial Proteomics: Functional Biology of Whole Organisms. Edited by Ian Humphrey-Smith and Michael Hecker. Wiley Publishing, Inc. *Methods Biochem Analysis* 49:245-69 (2006).

Unique Transcriptome Signature of *Mycobacterium tuberculosis* in Pulmonary Tuberculosis. Helmy Rachman, **Michael Strong**, Timo Ulrichs, Leander Grode, Johannes Schuchhardt, Hans Mollenkopf, George A. Kosmiadi, David Eisenberg, S.H.E. Kaufmann. *Infection and Immunity* 74:1233-1242. (2006).

Mycobacterium tuberculosis Gene Expression Profiling within the Context of Protein Networks. Helmy Rachman^{*}, **Michael Strong**^{*}, Ulrich Schaible, Johannes Schuchhardt, Kristine Hagens, Hans Mollenkopf, David Eisenberg, Stefan H.E. Kaufmann. *Equal Contribution. *Microbes and Infection.* 8:747-757 (2006).

A specific secretion system mediates PPE41 transport in pathogenic mycobacteria. Abdallah M. Abdallah, Theo Verboom, Fredericke Hannes, Mohamad Safi, **Michael Strong**, David Eisenberg, Rene Musters, Christina M.J.E. Vandenbroucke-Grauls, Ben J. Appelmelk, Joen Luirink4 and Wilbert Bitter. *Molecular Microbiology* 62:667-79 (2006).

Scientists and Societies: Community Outreach. **Michael Strong**. *Nature*. 434, 418. (2005).

Protein Nanomachines. **Michael Strong**. *PLoS Biology* Mar, 2:E73. (2004).

A Web-Based Comparative Genomics Tutorial for Investigating Microbial Genomes. **Michael Strong**, Duilio Cascio, and David Eisenberg. *ASM Microbiology Education Journal* 5: 30-35. (2004).

Genomes, Maps, and Modules: Navigating the *M. tuberculosis* Genome. **Michael Strong**. *UCLA Scientific Review* 1: 28-32. (2004).

Visualization and Interpretation of Protein Networks in *Mycobacterium tuberculosis* Based on Hierarchical Clustering of Genome-Wide Functional Linkage Maps.

Michael Strong, Thomas G. Graeber, Morgan Beeby, Matteo Pellegrini, Michael J. Thompson, Todd O. Yeates, and David Eisenberg. *Nucleic Acids Research*, 31: 7099-7109. (2003).

Inference of Protein Function and Protein Linkages in *M. tuberculosis* Based on Prokaryotic Genome Organization: A Combined Computational Approach.

Michael Strong, Parag Mallick, Matteo Pellegrini, Michael J. Thompson, and David Eisenberg. *Genome Biology* 4:R59.1-R59.16. (2003).

2'-Ribose-ferrocene oligonucleotides for electronic detection of nucleic acids. Yu CJ, Wang H, Wan Y, Yowanto H, Kim JC, Donilon LH, Tao C, **Strong M**, Chong Y. *Journal of Organic Chemistry* 66:2937-2942. (2001).

Uridine-Conjugated Ferrocene DNA Oligonucleotides: Unexpected Cyclization Reaction of the Uridine Base. Yu, C. J.; Yowanto, H.; Wan, Y.; Meade, T. J.; Chong, Y.; **Strong, M**.; Donilon, L. H.; Kayyem, J. F.; Gozin, M.; Blackburn, G. F. *Journal of the American Chemical Society*. 122: 6767-6768. (2000).

WORK EXPERIENCE

California Institute of Technology. I worked at Caltech with Suzanna Horvath on projects involving DNA and RNA synthesis applying phosphoramidite chemistry 1997-1998.

Clinical Microsensors (Motorola Biosciences). I worked with C.J. Yu on projects relating to the electronic detection of nucleic acids, for use in medical diagnostics. 1998-1999

TEACHING AND MENTORING EXPERIENCE

University of California, Los Angeles.

*Teaching Assistant, Microbiology 101L, Microbiology Laboratory Course. (2002 and 2003) *I created a web-based genomics tutorial for the UCLA Macromolecular Structure class M253 (Fall 2003) *I taught a database and web design tutorial for the UCLA Molecular Biology Institute (Fall 2003) *I mentored middle school and high school students once a week at UCLA for about two years (2005-2006)

Harvard University

*Teaching Fellow. I was a teaching fellow for the 2007 Harvard iGEM team (Summer 2007)

National Jewish Health

Volunteered to teach basic Genetics concepts to students at the Kunsberg School (Summer 2010).

University of Colorado, Denver

Faculty- CPBS 7711 - Bioinformatics 1 (2 lectures) Fall 2010, 2011

Faculty– Biological Science Program Journal Club (1 lecture) Fall 2010

Faculty CPBS 7712 – Bioinformatics (2 lectures) Spring 2011, 2012

AWARDS

CSIS Smart Global Health Essay Contest, 1st Place Essay (2010) Jane Coffin Childs Memorial Fund For Medical Research, Postdoctoral Fellowship (2006 – 2009) NIH National Research Service Award GM07185. (Cell and Molecular Biology Training Grant) (2003-2005) Paul D. Boyer Outstanding Teaching Award (2003) Amgen-MBI Dissertation Year Award (2004) UCLA Dissertation Year Fellowship (2004) Role Model Award (2004) Glen T. Seaborg Poster Award (2004) Howard Hughes Medical Institute Undergraduate Research Grant (1996) UCSB College of Letters and Science Award for Outstanding Research (1997) HHMI Award for Outstanding Undergraduate Research (1997)

TALKS

American Society for Microbiology Biodefense Conference (2003) 10th International Conference on Microbial Genomes (2002) Lake Arrowhead Biological Chemistry Retreat (2002) UCLA Bioinformatics Seminar Series (2004) UCLA Molecular Biology Interdepartmental Retreat (2003,2004) West Coast Protein Crystallography Conference (2005) MIT iGEM T3 (2007) National Jewish Health (2009) Harvard Medical School, Dept. of Genetics (2009) University of Colorado, Denver, Computational Bioscience Program Seminar (2010) Colorado State University, Front Range TB meeting (2010) Tuberculosis Trials Consortium meeting TBTC (2011) University of Colorado, Infectious Diseases Department Grand Rounds (2012)

POSTER PRESENTATIONS

Mycobacterium tuberculosis Keystone conference (2003,2005) TB Structural Genomics Consortium Retreat (2001, 2002, 2003) UCLA Bioinformatics Recruitment Program (2002). 24th Annual MBI Lake Arrowhead Conference (2001) NIH Protein Structure Initiative Workshop (2004) 12th International Conference on Microbial Genomes (2004) 1st Annual Duke Systems Biology Symposium (2006) Harvard Medical School Genetics Retreat (2007) New England TB Retreat at the Broad Institute (2007) Jane Coffin Childs Annual Symposium (2007, 2008) Pacific Symposium on Biocomputing (2011, 2012)

COMMITTEES/ORGANIZER

UCLA Faculty Executive Committee (2003-2004 school year) Graduate Student Representative Molecular Biology Interdepartmental Program (2003-2005) Graduate Student Representative Biological Sciences Council (2004-2005 school year) MBI Graduate Student Representative National Jewish Health, Integrated Center for Genes, Environment, and Health, Genetics Group Meeting Co-organizer (2010,2011)

University of Colorado, Denver, Computational Bioscience Program, Admissions Committee (2010-present) University of Colorado, Denver, Computational Bioscience Program, Preliminary Exam Chair (2011)

MEMBERSHIPS

American Society for Microbiology

University of Colorado, Center for Global Health, Affiliate

International Society for Computational Biology