

Mark Alber

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Fields: Mathematical and Computational Biology, Multi-scale Modeling, Blood Clotting, Bacterial Biofilms and Swarming, Cancer Invasion, Early Development

Education:

- 1990 - Ph.D. in Mathematics, *University of Pennsylvania, Philadelphia, PA*
advisor: J.E. Marsden, Caltech
1983 - Honors M.S. in Applied Mathematics, *Department of Applied Mathematics, Moscow Institute of Technology*

Positions Held:

- 2016 - present Distinguished Professor, Department of Mathematics,
University of California, Riverside (UCR)
2017 - present Director, Center for Quantitative Modeling in Biology, UCR
2017 - present Cooperating Faculty, Department of Internal Medicine,
Division of Clinical Sciences, School of Medicine, UCR
2017 - present Cooperating Faculty, Department of Bioengineering, UCR
2017 - present Member of the Institute for Integrative Genome Biology, UCR
2017 - present Member of the Faculty Advisory Board to the Office of Research and
Economic Development, UCR
2017 - present Member of the UCR Bioengineering Interdepartmental Graduate Program
2010 - present Adjunct Professor of Medicine, Indiana University School of Medicine
2009 - 2016 Vincent J. Duncan Family Professor of Applied Mathematics,
University of Notre Dame (UND)
2011 - 2015 Director of Graduate Studies, ACMS Department, University of Notre Dame
2003 - 2016 Director, Interdisciplinary Center for the Study of Biocomplexity
2003 - 2016 Concurrent Professor of Physics, UND
2011 - 2016 Concurrent Professor of Computer Science and Engineering, UND
2012 - 2016 Program Co-Leader of the Computational Modeling, Drug Development
and Delivery (CoMD3) Program, Harper Cancer Research Institute,
IU School of Medicine and University of Notre Dame
2006 - 2009 Notre Dame University Professor of Applied Mathematics
2001 - 2006 Full Professor of Mathematics, University of Notre Dame
1996 - 2001 Tenured Associate Professor, University of Notre Dame
2000 - 2001 On Sabbatical leave at Stanford University, CA
March-June 2001 Senior Fellow, Institute for Pure and Applied Mathematics, UCLA, CA
October 2000 Institute for Mathematics and its Applications (IMA), Minneapolis
1990 - 1996 Tenure Track Assistant Professor, University of Notre Dame

Summer 1995, 1996	Basic Research Institute in the Mathematical Sciences (BRIMS) Hewlett-Packard Research Lab., Bristol, UK
Summer 1994	CNLS, Los Alamos Natl. Lab., NM
1993 - 1994	On Sabbatical leave at UC Berkeley, CA; Institute for Advanced Study, Princeton, NJ; Mathematical Sciences Research Institute, Berkeley, CA
June 1993	The Fields Institute for Research in Mathematical Sciences, Canada
Summer 1990	Instructor, University of Pennsylvania

Honors and Service:

2011	Elected Fellow, American Association for the Advancement of Science (AAAS)
2016 - present	Distinguished Professor, Department of Mathematics, UC Riverside
2009 - 2016	Vincent J. Duncan Family Professor of Applied Mathematics
2006 - 2009	Notre Dame University Professor of Applied Mathematics
2016	Keynote Speaker, 8th International Bio-fluid Mechanics Symposia, Caltech
2013	Distinguished Lecture in Applied Mathematics, U of Massachusetts, Amherst
2013 - present	Deputy Editor, PLoS Computational Biology
2010 - present	Associate Editor, Bulletin of Mathematical Biology
2010 - present	Member of the Editorial Board, Biophysical Journal
2010-present	Review Editor of Frontiers in Computational Physiology and Medicine
2011 - 2013	Member of the Editorial Board, PLoS Computational Biology
2008 - 2012	Member of the Editorial Board, Journal of Statistical Physics
2017	Member of the IRNIA Computational Biology Review Panel (Paris, France)
2010-2017	Member of 5 NSF Panels and 12 NIH panels including 2017 NSF CAREER Review Panel, 2017 NSF National Scientific Centers Review Panel, 2017 NIH National Institute of Biomedical Imaging and Bioengineering Special Emphasis Panel ZEB1 OSR-C (J2) MSM Program Review Panel (two times), Modeling and Analysis of Biological Systems (MABS) Study Section, Special Emphasis Panel ZRG1 IMSTJ 40, SBIR/STTR Study Section IMST-K(14)B, Special Emphasis Panel ZRG1 BST-N (51), 2014 Big Data Panels at NIH and NSF.
2009 - 2015	Field Editor for Applications in Biosciences and Medicine of the Encyclopedia of Applied and Computational Mathematics (Springer)
2010-present	Member of the Committee of the American Mathematical Society on Human Rights of Mathematicians
2013 - 2016	Representative of the American Mathematical Society to the AAAS Human Rights Coalition (HRC) and member of the Council of the HRC
June 23, 2009	Member of the External Scientific Review Board of the NIH Center for Complex Biological Systems, University of California, Irvine
2001, 2004, 2008	Member of NSF and NIH Panels

Research Support:

DMS-1762063, Joint NSF DMS/NIH NIGMS Initiative to Support Research at the Interface of the Biological and Mathematical Sciences, Combined Modeling and Experimental Study of the Mechanisms of Growth Patterns in Stem Cell Homeostasis in Plants. 7/1/2018-6/30/22, \$1,100,000, PI: Alber. This interdisciplinary research program aims to understand

how mechanical connections among cells and chemical signals between them collaborate to control the growth, self-organization and differentiation of stem cells during plant growth. Mathematical and computer models will allow researchers to perform virtual experiments that are currently impossible in the lab. Coupled with live imaging experiments and new image analysis methods, these experiments will yield insights into biological mechanisms governing organ formation in plants and animals and development of cancer in epithelial cell layers of the colon. UC Riverside is a Hispanic-serving institution with very diverse student population and with many students being first in their families to attend college.

NIH grant U01 HL116330 (Predictive Multiscale Models for Biomedical, Biological, Behavioral, Environmental and Clinical Research Interagency Initiative). Multiscale modeling and empirical study of a mechanism limiting blood clot growth. 07/26/2014 - 07/30/2019, \$3,550,000, PI: Alber. Collaborative project with University of Pennsylvania School of Medicine and Argonne National Laboratory. Thrombosis is a major cause of death in the developed world and results from the growth of thrombi (blood clots forming within blood vessels) that restricts blood flow to vital organs. The project integrates multiscale modeling and experiments to examine novel hypothesis related to the role of fibrin networks in processes halting thrombus growth. This will help physicians to estimate risk of thrombotic disease for an individual patient by identifying critical values of parameters of processes regulating thrombogenesis.

NIH grant 1R01GM095959. Combined multiscale modeling and experimental study of bacterial swarming, 04/01/2012 - 12/31/2016, \$1,149,272, PI: Alber. Most infections are the result of surface-attached biofilm communities of bacteria that colonize host surfaces. *Pseudomonas aeruginosa* is an opportunistic pathogen responsible for both acute and persistent infections in susceptible individuals, as exemplified by those for burn victims and people with cystic fibrosis. A key aspect of these infections is the formation of bacterial swarms, which are surface-associated, socially organized communities of cells. Because identification of single cell behavior within groups is extremely difficult experimentally, we will use multiscale models to perform predictive simulations describing complex bacterial interactions that potentially control swarming. This combined multiscale modeling and laboratory study of bacterial behavior on surfaces will provide new critical information needed for the eradication, prevention and treatment of the *P. aeruginosa* infections.

NIH grant 1R01GM100470 (Joint NSF DMS/NIH NIGMS Initiative to Support Research at the Interface of the Biological and Mathematical Sciences). Study of the interplay of motility mechanisms during swarming of *Myxococcus xanthus*, 06/01/11 - 06/01/15, \$779,565, PI: Alber. The main goal of this interdisciplinary project is to combine simulations using new three-dimensional multiscale modeling environment and specifically designed experiments to study basic coordination events of *M. xanthus* swarming, which is essential to understanding how millions of bacteria function in real environments. A collaboration has been established between Drs. Alber (PI), Xu, Chen, Shrout (Notre Dame), Dr. Aronson from the Argonne National Laboratory (ANL) and Dr. Kaiser from Stanford University to achieve these goals.

NSF Grant CBET-1403887 (CBET - BIOTECH, BIOCHEM & BIOMASS ENG), Decoding organ-level intercellular signaling in an active, regulated microenvironment, 05/01/14 - 04/30/17, \$600,000, PI: Zartman, Co-PIs: Alber and Hoelzle. The proposed study aims

to investigate the hypothesis that intercellular calcium waves encode information on the size, differentiation state and overall physiology of epithelia. To test this hypothesis, new biophysical methods will be combined with computer simulations to investigate intercellular calcium signaling in intact epithelial organs within a controlled microenvironment.

NSF grant MCB-1244593. BioMaPS: Experimental and Computational Studies of Microtubule Dynamics and Regulation by Binding Proteins, 2017 07/01/13 - 06/30/17, \$667,346, PI: Goodson, Co-PI: Alber; The long-term goal of this collaborative project is to develop a comprehensive understanding of MT dynamics and its regulation by MT binding proteins through a coordinated program of multiscale computational modeling and experiment. The specific goals of this renewal are to 1) use combined experiment and modeling to establish fundamental principles for how groups of MTBinding Proteins (MTBPs) work together; 2) define the relationship between the behavior of the bulk polymer and that of the individual MTs, and 3) develop freely available software packages to allow students and researchers at remote sites to use our models by implementing them on their own computers or running them on the web.

NSF grant MCB 0951264, Cellular Organization, Computational and Experimental Studies of Microtubule Dynamics and Regulation by Binding proteins, 03/15/10 - 03/15/13, \$640,058, PI: Goodson, Co-PI: Alber. The long-term goal of this project was to develop a predictive and quantitative understanding of the MT cytoskeleton and its regulation by MTBPs, which will impact fields ranging from systems biology to nanotechnology. The flexible model and tutorials developed through this project allows researchers to develop and test specific hypotheses about the mechanisms of dynamic instability and MTBP action, which will in turn help design and direct future experiments.

Gerber Foundation. Development of technology to assess bruises. 09/15/10-/09/15/13, \$90,000, PI: Alber. The main goal of this project is to use novel spectrophotometry and predictive 3D multiscale modeling to generate specific hypothesis on the age of bruises.

NSF Grant DMS-0800612 (Joint NSF DMS/NIH NIGMS Initiative to Support Research at the Interface of the Biological and Mathematical Sciences), Integrating Multiscale Modeling and in vivo Experiments for Studying Blood Clot Development, 09/01/08 - 09/01/11, \$864,000, PI: Alber. The overall goal of this project was to develop 3-dimensional multiscale mathematical models and a computational toolkit for simulating thrombus formation. These models were validated with specifically designed experiments to test predictions of thrombus development, its structure and stability. Moreover, the development of models served as a generator of new hypotheses that were tested in experiments in vivo and micro fluidic devices.

NSF grant BCS 0826958, DHB. Longitudinal Analysis and Modeling of Large-Scale Social Networks, 10/01/08 - 10/01/11, \$699,770, Co-PI: Alber. The data generated by digital communication technologies was used to (1) test/validate existing social network theories about the mechanisms underlying network dynamics by developing quantitative high fidelity temporal stochastic models of human behavior within social networks; (2) produce a data-driven, dynamic network modeling suite with predictive capabilities.

NSF Grant DMS-0719895, AMS-SS: Multiscale stochastic model of myxobacteria dynamics, 08/15/07-07/31/2010, \$222,000, PI: Alber; A large amount of available, reliable experimental data on single wild type and mutant cell behavior and interactions between cells was used as building blocks for developing comprehensive mathematical and computational multi-dimensional hybrid model of bacterial swarming based on short and long range interactions. To bring different scale levels of description together new mathematical techniques and tools were used including elements of stochastic analysis, nonlinear analysis and kinetic theory.

NSF Grant CCF 0622940, CompBio: Simulation of self-emerging properties of coupled biochemical and cellular networks in social behavior of Myxobacteria, Biology and Information Technology, 09/01/06-08/31/09, \$300,000, Co-PI:Alber. The project bridged the molecular, subcellular, cellular, and macroscopic level models by introducing consistent mathematical and computational interfaces among them. The general approach was demonstrated by developing a multi-scale computational model of formation of Myxobacteria fruiting bodies based on short range (C-signaling) interactions, differentiation and motility regulation, and by analyzing the effect of genetic mutations on macroscopic development.

DOE: Northwest Indiana Computational Grid (NWICG), Modeling Microtubule Dynamics, 09/01/08 - 09/01/09, \$50,000, PI: Alber. The focus of this project was on 1) the development of the detailed microscale model of several microtubules formation and its parallelized computational implementation; 2) development of multi-scale model of venous thrombosis.

NIH 1 RO1 GM065420: Supplement for the Study of Complex Biological Systems, 08/01/05-07/31/08, \$320,000, Co-PI: Alber. This grant supported collaboration between cell biologist Holly Goodson and applied mathematician Mark Alber to use a coupled program of iterative computational modeling and experiment to investigate key aspects of dynamic microtubule systems.

NIH 1 RO1 GM76692-01: Interagency Opportunities in Multiscale Modeling in Biomedical, Biological and Behavioral NSF 04.6071, (collaboration with IU Bloomington and Medical School at Kansas University) 09/01/05-08/31/08, \$800,000, Co-PI: Alber. The investigators developed an open-source comprehensive Tissue-Simulation-Toolkit (TST) using multi-scale mathematical modelling and computer simulation techniques to investigate the mechanism of segmentation during somite and limb formation.

NSF Grant 0344647, Division of Integrative Biology and Neuroscience, BIO Directorate and DMS Applied Mathematics, Testing a Developmental Mechanism by an Integrated Empirical-Computational Approach, (collaboration with Medical College of New York) 04/15/04-04/15/07, \$360,000, Co-PI: Alber. The objective of this project was to define and test an activator-inhibitor based mechanism for skeletal pattern formation in vertebrate limb mesenchymal cell cultures.

NSF MRI Grant 0420980, Acquisition of the Notre Dame Biocomplexity Computer Cluster, 08/01/04-07/31/07, PI: Alber. This grant supported the purchase of a Beowulf type Biocomplexity Cluster (62 dual processors nodes with 64-bit architecture) to provide computer infrastructure for The Notre Dame Interdisciplinary Center for the Study of Biocomplexity

(ICSB) to meet its broad research and educational goals.

NSF Grant 0083653, BIOCOMPLEXITY Multiscale Simulation of Avian Limb Development (collaboration with IU Bloomington, PI on the Notre Dame subcontract: Alber) 9/1/00-8/31/05, \$3,000,000. The goal the project was to develop mathematical and computational models of avian limb structural development based on physical processes including cell adhesion, chemotaxis and haptotaxis, mitosis, and cell differentiation. A unified, object-oriented, three-dimensional computational framework was introduced which included composite discrete and continuous modeling and computation. It was used for simulating chondrogenesis in the avian limb. The computer package is available for uploading at: <https://simtk.org/home/compuccell3d>

REU Supplement, NSF Biocomplexity Grant 0083653, 06/01/03-09/01/07, \$25,000, PI.

NSF DMS05-17864, 09/01/05-08/31/06, \$25,000, PI; DOE grant \$15,000, PI; Indiana 21st Century Research and Technology Fund, \$5,000; Workshop on Applications of Methods of Stochastic Systems and Statistical Physics

Burroughs Wellcome Fund, Biocomplexity V Workshop, August 2003, \$5,000, PI

Publications:

Edited Special Issues of Journals and Books:

1. Special Issue of the Bulletin of Mathematical Biology, "Multi-scale Modeling of Tissue Growth and Shape", Mark Alber, Christophe Godin, Philip Maini, Roeland Merks, Eric Mjolsness, Editors (under preparation).
2. Mark Alber, Editor for the Section on Mathematical and Computational Methods in Biosciences and Medicine of the Encyclopedia of Applied and Computational Mathematics, *Springer*, 2016, 1676 p. In 2 volumes.
3. Special Issue of the Bulletin of Mathematical Biology, "Biomedical Modeling", M.Alber, P.Maini, G. Niebur, Editors, Volume 75, Issue 8, August 2013. ISSN: 0092-8240.
4. Special Issue on Applications to Biology of the Journal of Statistical Physics, Mark Alber, Ray Goldstein, Erwin Frey, Editors, *Springer*, Volume 128, Issue 1-2, July 2007.
5. Special Issue on Multiscale Modeling in Biology, SIAM Journal: Multiscale Modeling and Simulation, Mark Alber, Thomas Hou, James A. Glazier, Yi Jiang, Editors, *SIAM*, Volume 3, Number 2, 2005.
6. Special Issue of the Journal: Biofilms, Clay Fuqua, James A. Glazier, Yves Brun and Mark S. Alber, Editors, *Cambridge University Press*, Volume 1, Number 4, 2004.
7. Alber, M.S. , B. Hu and J. Rosenthal, Editors, Current and future directions in applied mathematics [1997]. Papers from the symposium held at the University of Notre Dame, Notre Dame, IN, April 1996. *Birkhuser Boston, Inc., Boston, MA* x+261.

Papers in Peer-reviewed Journals:

Mathematical and Computational Biology and Medicine, Biomaging and Experimental Approaches

8. Shixin Xu, Mark Alber and Zhiliang Xu [2017], *Three-phase Model of Visco-elastic Incompressible Fluid Flow and its Computational Implementation*, *Communications in Computational Physics* (to appear).
9. Shixin Xu, Zhiliang Xu, Oleg Kim, Rustem I. Litvinov, John W. Weisel and Mark Alber [2017], Model Predictions of Deformation, Embolization, and Permeability of Partially Obstructive Blood Clots under Variable Shear Flow, *Journal of the Royal Society Interface* 14: 20170441. <http://dx.doi.org/10.1098/rsif.2017.0441>
10. Oleg V. Kim, Rustem I. Litvinov, Mark S. Alber and John W. Weisel [2017], Quantitative Structural Mechanobiology of Platelet-Driven Blood Clot Contraction, *Nature Communications* 8: 1274. <https://www.nature.com/articles/s41467-017-00885-x.pdf> (authors for correspondence: J.W. Weisel and M. Alber).
11. Peter Hk, Rustem I. Litvinov, Oleg V. Kim, Shixin Xu, Zhiliang Xu, Joel S. Bennett, Mark S. Alber and John W. Weisel [2017], Strong Binding of Platelet Integrin IIb3 to Fibrin Clots: Potential Target to Destabilize Thrombi, *Scientific Reports* 7: 13001 (published by Nature) DOI:10.1038/s41598-017-12615-w
12. Y Klymenko, O Kim, E Loughran, J Yang, R Lombard, M Alber and MS Stack [2017], Cadherin composition and multicellular aggregate invasion in organotypic models of epithelial ovarian cancer intraperitoneal metastasis, *Oncogene* Oct 19;36(42):5840-5851. doi: 10.1038/onc.2017.171 (published by Springer Nature, authors for correspondence: M.S. Stack and M. Alber).
13. Peter Hook, Teresa Brito-Robinson, Oleg Kim, Cofy, Narciso, Holly V. Goodson, John W. Weisel, Mark S. Alber and Jeremiah J. Zartman, Whole Blood Clot Optical Clearing for Nondestructive 3D Imaging and Quatitative Analysis, *Biomedical Optics Express*, Volume 8, Issue 8, Page 3671. (authors for correspondence: J.Zartman, J.W. Weisel and M. Alber).
14. Ali Nematbakhsh, Wenzhao Sun, Pavel A. Brodskiy, Aboutaleb Amiri, Cody Narciso, Zhiliang Xu, Jeremiah J. Zartman, Mark Alber [2017], Multi-scale computational study of the mechanical regulation of cell mitotic rounding in epithelia, *PLoS Computational Biology*, 13(5): e1005533. <https://doi.org/10.1371/journal.pcbi.1005533>
15. Aranda R. Duan, Erin M. Jonasson, Emily O. Alberico, Chunlei Li, Jared P. Scripture, Rachel A. Miller, Mark S. Alber and Holly V. Goodson [2017], Interactions between Tau and different conformations of tubulin: Implications for Tau function and mechanism, *Journal of Molecular Biology* 429(9):1424-1438. doi: 10.1016/j.jmb.2017.03.018.
16. O.V. Kim, R.I. Litvinov, J. Chen, D.Z. Chen, J.W. Weisel and M.S. Alber [2017], Compression-induced structural and mechanical changes of fibrin-collagen composites, *Matrix Biol.* 2017 Jul;60-61:141-156. doi: 10.1016/j.matbio.2016.10.007. Epub 2016 Oct 15.

17. Aboutaleb Amiri, Cameron Harvey, Amy Buchmann, Scott Christley, Joshua D. Shrout, Igor S. Aranson, Mark Alber [2017], Reversals and collisions optimize protein exchange in bacterial swarms, *Physical Review E* 95, 032408.
18. Martina Buka and Mark Alber[2017], Multi-component model of intramural hematoma, *Journal of Biomechanics* 50, 4249.
19. Kim O, Liang X, Litvinov RI, Weisel JW, Alber MS, Purohit PK. [2016], Foam-like compression behavior of fibrin networks, *Biomech Model Mechanobiol.* 15:213228.
20. Chen, J., M.S. Alber, and D.Z. Chen, A Hybrid Approach for Segmentation and Tracking of Myxococcus xanthus Swarms [2016], *IEEE Transactions on Medical Imaging* 35(9):2074-84.
21. Giordano Tierra, Juan P. Pavissich, Robert Nerenberg, Zhiliang Xu and Mark S. Alber [2015], Multicomponent model of deformation and detachment of a biofilm under fluid flow, *J. R. Soc. Interface* 12: 20150045.
22. Morales-Soto, N., Anyan, M. E., Mattingly, A. E., Madukoma, C. S., Harvey, C. W., Alber, M., Dziel, E., Kearns, D. B., and J.D. Shrout [2015], Preparation, Imaging, and Quantification of Bacterial Surface Motility Assays. *Journal of Visualized Experiments*, (98), e52338, doi:10.3791/52338.
23. Morgen E. Anyan, Aboutaleb Amiri, Cameron W. Harvey, Giordano Tierra, Nydia Morales-Soto, Callan M. Driscoll, Mark S. Alber, Joshua D. Shrout [2014], Type IV Pili Interactions Promote Intercellular Association and Moderate Swarming of *Pseudomonas aeruginosa*, *Proc. Natl. Acad. Sci. USA* vol. 111, no. 50, 18013-18018 (authors for correspondence: J. Shrout and M. Alber).
24. Wu Z, Xu Z, Kim O, Alber M. [2014], Three-dimensional multi-scale model of deformable platelets adhesion to vessel wall in blood flow. *Philosophical Transactions of the Royal Society A* 372: 20130380.
25. Chunlei Li, Jun Li, Holly V. Goodson and Mark S Alber [2014], Microtubule Dynamics Instability: the Role of Cracks between Protofilaments, *Soft Matter* 10, 2069-2080.
26. Amy Buchmann, Mark Alber and Jeremiah J. Zartman [2014], The mechanical feedback hypothesis of organ growth regulation, Modelling developmental signalling, *Seminars in Cell and Developmental Biology* 35, 7381.
27. J. Chen, C.W. Harvey, M.S. Alber, and D.Z. Chen [2014], A Matching Model Based on Earth Mover's Distance for Tracking Myxococcus xanthus, *Med Image Comput Assist Interv.* 17(Pt 2):113-20.
28. Oleg Kim, John McMurdy, Gregory Jay, Collin Lines, Gregory Crawford, and Mark Alber [2014], Combined Reflectance Spectroscopy and Stochastic Modeling Approach for Noninvasive Hemoglobin Determination via Palpebral Conjunctiva, *Physiological Reports*, Vol. 2, 1, e00192.
29. Oleg V. Kim, Rustem I. Litvinov, John W. Weisel and Mark S. Alber [2014], Structural basis for the nonlinear mechanics of fibrin networks under compression, *Biomaterials* 35, 6739–6749.

30. Cameron W. Harvey, Chinedu S. Madukoma, Shant Mahserejian, Mark S. Alber and Joshua D. Shrout [2014], Cell Division Resets Polarity and Motility for the Bacterium *Myxococcus xanthus*, *Journal of Bacteriology* 196, 22, 3853-3861.
31. Gupta, K., Li, C., Duan, A.R., Alberic, E.O., Kim, O.V., Alber M.S., and Goodson, H. V. [2013], A mechanism for the catastrophe-promoting activity of the microtubule destabilizer Op18/stathmin, *Proc. Natl. Acad. Sci. USA*, 2013 Dec 17;110(51):20449-54. doi: 10.1073/pnas.1309958110.
32. Oleg V. Kim, Zhiliang Xu, Elliot D. Rosen and Mark S. Alber [2013], Fibrin Networks Regulate Protein Transport during Thrombus Development, *PLoS Computational Biology* 9 (6), e1003095.
33. Cameron Harvey, Mark Alber, Lev Tsimring, Igor Aronson [2013], Continuum modeling of clustering of myxobacteria, *New Journal of Physics* 15, 035029.
34. Yong-Tao Zhang, Mark S. Alber and Stuart A. Newman [2013], Mathematical Modeling of Vertebrate Limb Development, *Mathematical Biosciences* 243, 117 (one of 5 most downloaded papers in 2013).
35. Cameron W Harvey, Huijing Du, Zhiliang Xu, Dale Kaiser, Igor Aranson, Mark Alber [2012], Interconnected Cavernous Structure of Bacterial Fruiting Bodies, *PLoS Computational Biology* 8 (12), e1002850.
36. Constance L. Slaboch, Mark S. Alber, Elliot D. Rosen, Timothy C. Ovaert [2012], Mechano-rheological properties of the murine thrombus determined via nanoindentation and finite element modeling, *Journal of the Mechanical Behavior of Biomedical Materials* 10, 75–86.
37. Huijing Du, Zhiliang Xu, Morgen Anyan, Oleg Kim, W. Matthew Leevy, Joshua D. Shrout and Mark Alber [2012], High density waves of the bacterium *Pseudomonas aeruginosa* in propagating swarms result in efficient colonization of surfaces, *Biophysical Journal* 103(3), 601-609.
38. Richard Gejji, Pavel Lushnikov and Mark Alber [2012], Macroscopic model of self-propelled bacteria swarming with regular reversals, *Physical Review E* 85, 021903 (highlighted in Faculty of 1000).
39. Richard Gejji, Bogdan Kazmierczak and Mark Alber [2012], Classification and Stability of Global Inhomogeneous Solutions of a Macroscopic Model of Cell Motion, *Mathematical Biosciences*, 238(1) 21–31.
40. Gennady Margolin, Ivan V. Gregoretto, Trevor M. Ciskovski, Chunlei Li, Wei Shi, Mark S. Alber and Holly V. Goodson [2012], The Mechanisms of Microtubule Catastrophe and Rescue: Implications from analysis of a dimer-scale computational model, *Molecular Biology of the Cell*, 23:4 642–656 (highlighted in Faculty of 1000).
41. Oleg Kim, John McMurdy, Collin Lines, Susan Duffy, Gregory Crawford and Mark Alber [2012], Reflectance spectrometry of normal and bruised human skins: Experiments and modeling, *Physiological Measurement* 33, 159-175.
42. Zhiliang Xu, Oleg Kim, Malgorzata Kamocka, Elliot Rosen and Mark Alber, Multi-scale Models of Thrombogenesis [2012], *Wiley Interdiscip Rev Syst Biol Med.* 4(3):237-46.

43. Christopher R. Sweet, Santanu Chatterjee, Zhiliang Xu, Katharine Bisordi, Elliot D. Rosen and Mark Alber [2011], Modeling Platelet-Blood Flow Interaction Using Sub-cellular Element Langevin Method, *Journal of the Royal Society Interface* 8 (65), 1760-71.
44. Yilin Wu, Yi Jiang, A. Dale Kaiser, Mark Alber [2011], Self-organization in bacterial swarming: Lessons from Myxobacteria, *Physical Biology* 8 (5), 055003.
45. Huijing Du, Zhiliang Xu, Joshua D. Shrouf and Mark Alber [2011], Multiscale Modeling of *Pseudomonas aeruginosa* Swarming, *Mathematical Models and Methods in Applied Sciences*, Vol. 21, Suppl. 939-954.
46. Cameron W. Harvey, Faruck Morcos, Christopher R. Sweet, Dale Kaiser, Santanu Chatterjee, Xiaomin Lu, Danny Chen and Mark Alber [2011], Study of elastic collisions of *M. xanthus* in swarms, *Physical Biology* 8, 026016.
47. Eungjun Kim, Oleg V. Kim, Kellie R. Machlus, Xiaomin Liu, Timur Kupaev, a Joshua Lioi, Alisa S. Wolberg, Danny Z. Chen, Elliot D. Rosen, Zhiliang Xu, and Mark Alber [2011], Correlation between fibrin network structure and mechanical properties: an experimental and computational analysis, *Soft Matter* 7, 4983.
48. Gennady Margolin, Holly V. Goodson, and Mark S. Alber [2011], Mean-field study of the role of lateral cracks in microtubule dynamics, *Physical Review E* 83, 041905.
49. Xu, Z., M.M. Kamocka, M.S. Alber, and E.D. Rosen [2011], Computational Approaches to Studying Thrombus Development, *Arterioscler Thromb Vasc Biol* 31, 500-505.
50. Jianfeng Zhu, Yong-Tao Zhang, Mark S. Alber and Stuart A. Newman [2010], Bare bones pattern formation: a core regulatory network in varying geometries reproduces major features of vertebrate limb development and evolution, *PLoS ONE* (5): e10892.
51. Xu, Z., J. Lioi, J. Mu, X. Liu, M.M. Kamocka, E.D. Rosen, D.Z. Chen and M.S. Alber [2010], A Multiscale Model of Venous Thrombus Formation with Surface-Mediated Control of Blood Coagulation Cascade, *Biophysical Journal* 98, 9, 1723–1732.
52. Morcos, Faruck, Marcin Sikora, Mark Alber, Dale Kaiser, and Jesus A. Izaguirre [2010], Belief Propagation Estimation of Protein and Domain Interactions using the Sum-Product Algorithm, *IEEE Transactions on Information Theory, Special Issue on Molecular Biology* 56, 2, 742–755.
53. Mu, J., X. Liu, M.M. Kamocka, Z. Xu, M.S. Alber, and E.D. Rosen, D.Z. Chen [2010], Segmentation, Reconstruction, and Analysis of Blood Thrombi in 2-Photon Microscopy Images, *EURASIP Journal on Advances in Signal Processing* Vol. 2010, Article ID 147216, 8 pages. doi:10.1155/2010/147216.
54. Kamocka, M.M., J. Mu, X. Liu, N. Chen, A. Zollman, B. Sturonas-Brown, K. Dunn, Z. Xu, D.Z. Chen, M.S. Alber and E.D. Rosen [2010], 2-Photon Intravital Imaging of Thrombus Development In Vivo, *Journal of Biomedical Optics* 15, 1, 016020.
55. Zhenyu Shi, Nan Chen, Yanan Du, Ali Khademhosseini and Mark Alber [2009], Stochastic model of self-assembly of cell-laden hydrogels, *Phys. Rev. E* **80** 061901.

56. Alber, M., Gejji, R., B. Kazmierczak [2009], Existence of Global Solutions of a Macroscopic Model of Cellular Motion in a Chemotactic Field, *Applied Mathematics Letters* **22** 1645-1648.
57. Zhu J., Zhang, Y., Newman, S.A., M.Alber [2009], A finite element model based on discontinuous Galerkin methods on moving grids for vertebrate limb pattern formation, *Mathematical Modeling of Natural Phenomena* **4** 4, 131–148.
58. Zhu, J., Y.-T. Zhang, S.A. Newman and M. Alber [2009], Application of discontinuous Galerkin methods for reaction-diffusion systems in developmental biology, *Journal of Scientific Computing* **40** 391-418.
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Results obtained by the Alber's group were featured in the following newspaper articles and reviews:

"UC Riverside math researchers find key to stopping ovarian cancer from spreading" By MARK MUCKENFUSS — mmuckenfuss@scng.com — The Press-Enterprise PUBLISHED: August 9, 2017 at 6:01 am — UPDATED: August 13, 2017 at 1:34 pm
<http://www.pe.com/2017/08/09/ucr-math-researchers-find-key-to-stopping-ovarian-cancer-from-spreading/>

Group's results on contraction of blood clots were featured on the front page of the NSF and by the NSF news.

"Researchers Look Inside Dangerous Blood Clots with Optical Clearing Technique", Press Release by the Optical Society of America, 17 July 2017

How blood clots shrink, *Microscopy and Analysis*, Editorial, November 10, 2017:
<https://microscopy-analysis.com/editorials/editorial-listings/how-blood-clots-shrink>

'Reversing helps bacterial swarms to spread' by Heidi Ledford, Nature News, January 20th, 2009, doi:10.1038/news.2009.43:
<http://www.nature.com/news/2009/090120/full/news.2009.43.html>

'Bacteria periodically reverse direction', The Times of India, 21 Jan 2009, 1249 hrs IST, ANI: <http://timesofindia.indiatimes.com/articleshow/msid-4010918,prtpage-1.cms>

July 1, 2013 cover story in The Scientist mentions Alber group's results on social behavior of *Myxococcus xanthus* and includes an interview with the former student Yilin Wu: <http://www.the-scientist.com/?articles.view/articleNo/36101/title/Crowd-Control/>

Lushnikov, P.P., Chen, N., and M.S. Alber [2008], Macroscopic dynamics of biological cells interacting via chemotaxis and direct contact, *Phys. Rev. E*, 78, 061904, was highlighted in the Faculty of 1000 Biology and Medicine: <http://f1000.com/prime>.

Gennady Margolin, Ivan V. Gregoretto, Trevor M. Ciskovski, Chunlei Li, Wei Shi, Mark S. Alber and Holly V. Goodson [2012], The Mechanisms of Microtubule Catastrophe and Rescue: Implications from analysis of a dimer-scale computational model, *Molecular Biology of the Cell*, 23:4 642–656, was highlighted in the Faculty of 1000 Biology and Medicine: <http://f1000.com/prime>

Group's results on modeling bacterial collective motion were reviewed in: Tamas Vicsek and Anna Zafeiris, Collective motion, *Physics Reports* 517 (2012) 71-140.

Optical Coherence Tomography News featured the groups' paper on collaborative project with Argonne National laboratory "Cameron W Harvey, Huijing Du, Zhiliang Xu, Dale Kaiser, Igor Aranson, Mark Alber [2012], Interconnected Cavernous Structure of Bacterial Fruiting Bodies, *PLoS Computational Biology* 8 (12), e1002850": <http://www.octnews.org/>
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Recent Invited Talks:

June 6, 2018, Conference on Multiscale Problems in Materials and Biology, The Fields Institute for Research in Mathematical Sciences, University of Toronto, Canada

April 22, 2018, NSF Biofilm Mechanical Properties Workshop, University of Notre Dame

January 12, 2018, Coupled Multi-scale Modeling and Experimental Study of Blood Clot Contraction and Deformation, SSH Forum on Simulation and Modeling in Healthcare, Los Angeles Convention Center

December 1, 2017, Mathematics Colloquium, UC Merced

October 20, 2017, Invited Talk, Retreat of the Institute for Integrative Genome Biology, UC Riverside

October 10, 2017, Colloquium, Universite Paris Diderot, Paris, France

July 18, 2017, Invited Talk, Annual Meeting of the Society for Mathematical Biology, Salt

Lake City, UT

May 3, 2017, Claremont Colleges Mathematics Colloquium, Claremont, CA

April 24, 2017, Mathematical Biology Colloquium, University of California, Davis, CA

February 1, 2017, Bioengineering Colloquium, University of California, San Diego

January 28, 2017, Invited Speaker at the 7th Annual Southern California Systems Biology Conference, University of California, Irvine

January 25th, 2017, Applied Mathematics Colloquium, Stanford University, Palo Alto, CA

September 4, 2016, Invited Talk, Workshop: Computing a Tissue, Modeling Multicellular Systems, 15th European Conference on Computational Biology, World Forum, The Hague, Netherlands

August 4th, 2016, Invited Talk, Mechbio Symposium: Putting together the Cell Mechanome, UC San Diego

February 13, 2016, Keynote Speaker at the 8th International Bio-fluid Mechanics Symposia, Caltech, Pasadena, CA:

November 14, 2015, Invited Talk, AMS Eastern Regional Meeting, Special Session "Multi-scale Methods in Cell and Developmental Biology", Rutgers University, Newark, NJ

September 8, 2015, Invited Talk, Meeting of the Interagency Modeling and Analysis Group (IMAG), Multiscale Modeling (MSM) Consortium, NIH, Bethesda, MD

September 2, 2015, Invited Talk, Computational Fluid Dynamics (CFD) in Medicine and Biology II An ECI Conference Series, Albufeira, Portugal

July 28, 2015, Invited Talk, 13th US National Congress on Computational Mechanics, San Diego, CA

May 8, 2015, Plenary Lecture, Workshop on Interdisciplinary Mathematics, Penn State University, PA

October 22, 2014, Applied Mathematics Colloquium, Duke University, Durham, NC

October 22, 2014, Applied Mathematics Colloquium, Stanford University, Palo Alto, CA

May 12, 2014, Invited Talk, 111th Statistical Mechanics Conference, Rutgers University, Piscataway, NJ

December 10, 2013, Invited Talk, SIAM Conference on Analysis of Partial Differential Equations, Special Session on Applications of PDEs to Biology, Lake Buena Vista, Florida

December 6, 2013, Mathematics Colloquium, IUPUI, IN

October 26, 2013, Invited Talk, Meeting of the Editorial Board of the PLoS Computational Biology, Washington DC.

October 24, 2013, Distinguished Lecture in Applied Mathematics, University of Massachusetts, Amherst, MA

October 1, 2013, Invited Talk, Seminar of the Center for Theoretical Biological Physics, Rice University, Houston, TX

May 6, 2013, Colloquium, Northeastern University, Boston, MA

March 26, 2013, Colloquium, North Carolina State University, Raleigh, NC

February 12, 2013, Colloquium, Dartmouth College, Hanover, NH

November 5, 2012, Invited Talk, Workshop on Collective Motion in Biological Systems: from Data to Models, Center for Interdisciplinary Research (ZiF), Bielefeld University, Germany

October 15, 2012, Invited Talk, Stowers Institute for Medical Research, Kansas City, Missouri

September 9, 2012, Colloquium, Institute for Medicine and Engineering (IME), University of Pennsylvania, Philadelphia, PA

August 8, 2012, Invited Talk, SIAM Meeting on Life Sciences, San Diego, CA

July 29, 2012, Invited Talk, Myxo 2012: 39th International Conference on the Biology of the Myxobacteria, July 29 - August 1, 2012, Chicago, IL

May 1, 2012, Invited Talk, 3rd International Conference on Engineering Frontiers in Pediatric and Congenital Heart Disease, Stanford University, Palo Alto, CA

April 17, 2012, Colloquium, Department of Mathematics, University of South Alabama, Mobile, AL

April 16, 2012, Colloquium, Department of Mathematics, University of South Carolina, Columbia, SC

April 12, 2012, Colloquium, Harper Cancer Research Institute, Indiana University School of Medicine, South Bend, IN

March 1, 2012, Invited Talk, Invited Session: Physical Mechanisms of Collective Microbial Dynamics, American Physical Society March Meeting 2012, Boston, MA

February 9, 2012, Colloquium, Department of Applied Mathematics, University of Washington, Seattle, WA

November 4, 2011, Invited Talk, Department of Mathematics, Duke University, NC

October 13, 2011, Invited Talk, 48th Annual Technical Conference of Society of Engineering

Sciences, Northwestern University, Evanston, Illinois.

April 12, 2011, Invited Talk, Principles and Theory of Self-Assembly, 7th Annual Conference on Foundations of Nanoscience: Self-assembled Architecture and Devices (FNANO 2011), Snowbird Cliff Lodge, Snowbird, Utah.

February 24, 2011, Mathematical Physics Colloquium, Rutgers University, New Brunswick, NJ.

February 2, 2011, Colloquium, Department of Mechanical Engineering, UC San Diego, CA.

September 14, 2010, Invited Talk, The Hot Topics Workshop "Medical Device-biological Interactions at the Material-tissue Interfaces", Institute for Mathematics and Its Applications (IMA), University of Minnesota, Minneapolis, MN.

June 25, 2010, Applied Mathematics Colloquium, Imperial College, London, UK.

June 21, 2010, Invited Talk, OCCAM Conference on Modelling at different scales in biology, St Anne's College, Oxford University, UK.

May 10, 2010, Invited Talk, 103rd Statistical Mechanics Conference, Rutgers University, NJ.

April 17, 2010, Plenary Talk, SIAM Great Lakes Conference: Modeling and Numerical PDEs in Mathematical Biology, University of Michigan-Dearborn, MI.

March 8, 2010, Applied Mathematics Colloquium, Department of Mathematics, Massachusetts Institute of Technology, Boston, MA

January 22, 2010, Mathematics Colloquium, Case Western University, Cleveland, OH

October 27, 2009, Workshop on Self-Organization and Multi-Scale Mathematical Modeling of Active Biological Systems, NSF Statistical and Applied Mathematical Sciences Institute (SAMSI), Research Triangle Park, NC

October 12, 2009, Conference on Agent-Based Complex Systems, NSF Institute for Pure and Applied Mathematics (IPAM), UCLA, Los Angeles, CA

September 26, 2009, Symposium on Engineered and Natural Complex Systems-Modeling, Simulation and Analysis, 2009 IEEE Toronto International Conference - Science and Technology for Humanity, Canada

August 30, 2009, Tutorial Lecture, 2009-10 Program on Stochastic Dynamics, NSF Statistical and Applied Mathematical Sciences Institute (SAMSI), Research Triangle Park, NC

August 24, 2009, Plenary Talk, Workshop, NSF Mathematical Biology Institute, Columbus, OH

July 23, 2009, Invited Talk, Center for Mathematical Biology, Oxford University, UK

July 22, Invited Talk, Conference on Cardiac Physiome Project, Newton Institute, Cambridge University, UK

April 10, 2009, Applied Mathematics Colloquium, Department of Mathematics, Stanford University, Palo Alto, CA

April 3, 2009, Colloquium, Department of Mathematics, New Jersey Institute of Technology, Newark, NJ

April 2, 2009, Department of Mathematics, Rutgers University, Piscataway, NJ

March 11, 2009, Department of Mechanical Engineering, University of California, Santa Barbara, CA

November 17, 2008, Department of Control Theory and Dynamical Systems, Caltech, Pasadena, CA

November 14, 2008, Department of Mathematics, University of California, Irvine, CA

October 14, 2008, Pattern Formation and Development in Colonial Organisms, NSF Mathematical Biology Institute, Ohio State University, Columbus, OH

August 6, 2008, Mini symposium on Networks Structure and Dynamics, SIAM Meeting in Life Sciences, Montreal, Canada

July 31, 2008, Mini symposium on Multiscale Modeling in Biology, Annual Meeting of the Society for Mathematical Biology Conference, Toronto, Canada

July 25, 2008, NSF EMT Workshop, Princeton University, NJ

29 February 2008, Department of Systems Biology, Harvard Medical School, Boston

December 10, 2007, Workshop on Biomechanics and Chemotaxis, Johann Radon Institute for Computational and Applied Mathematics (RICAM), Linz, Austria

November 12, 2007, Conference on Microfluids, NSF Mathematical Biosciences Institute, Columbus, Ohio

October 5, 2007, Special Session on Networks, American Mathematical Society regional meeting, De Paul University, Chicago

July 24, 2007, 6th European Conference on Computational Biology (ECCB), Vienna, Austria

June 1, 2007, Plenary Invited Talk, Indy Midwest Regional Bioinformatics Conference, IUPUI Conference Center, Indianapolis, IN

May 21, 2007, Conference on Mathematical Issues in Stochastic Approaches for Multiscale Modeling, Mathematical Sciences Research Institute, Berkeley, CA

March 16, 2007, Applied Mathematics and Computational Science (AMCS) Colloquium, University of Pennsylvania, Philadelphia

March 12, 2007, Department of Genetics, Indiana University Medical School, Indianapolis

University Service:

2017 - current, Director, Interdisciplinary Center for Quantitative Modeling in Biology, UC Riverside

2017 - current, Member of the Advisory Board to the Office of Research and Economic Development, UC Riverside

2003 - 2016: Director, Interdisciplinary Center for the Study of Biocomplexity, University of Notre Dame

2012 - 2013: Member of the Talent Subcommittee of the University of Notre Dame Corporate Governance Committee

2008-2010, Member of the University of Notre Dame Committee on Statistics

2005-2009: Member of the Faculty Advisory Committee of the Center for Research Computing, University of Notre Dame

2006-2007: Member of the University of Notre Dame Graduate Council

2006 - 2010: Member of the Board of the Center for Applied Mathematics, University of Notre Dame

2005-2006: Member of the University of Notre Dame Council for Academic Technologies

Postdoctoral Associates:

University of California, Riverside:

2015-current, Ali Nematbakhsh, Oleg Kim

2015-2017, Shixin Xu,

University of Notre Dame:

2014-2016, Peter Hook

2014-2015, Ling Xu, Ava Mauro, Nydia Morales Soto

2012-2014, Ziheng Wu, computational biology

2013-2014, Cameron Harvey, computational biology

2012-2013, Giordano Tierra Chica, computational and mathematical biology

2010-2013, Oleg Kim, computational and mathematical biology

2011-2014, Nydia Morales Soto, microbiology

2011-2012, Xiaomin Liu, image reconstruction and analysis

2009-2010, Chris Sweet, computational and mathematical biology

2009-2010, EunJung Kim, computational and mathematical biology

2005-2008, Nan Chen, computational and mathematical biology. He is currently a Computational Scientist at University of Texas, M.D. Anderson Cancer Center, Houston, TX.

2005-2008, Gennady Margolin, computational and mathematical biology. He is currently a Bioinformatics Scientist at National Institutes of Health, Bethesda, MD.

2005-2006, Pavel Lushnikov, mathematical biology. He is currently Full Professor, Department of Mathematics, University of New Mexico.

2004-2006, Olga Sozinova, mathematical biology

2001-2004, Rajiv Chaturvedi, mathematical biology

2003-2004, Bogdan Kazmierczak, mathematical biology

2001-2002, Xinan Zhang, mathematical biology

1995-1997, Gregory Luther, NSF Math. Sci. Postdoctoral Industrial Research Fellowship

Graduate Students:

University of California, Riverside:

2017-present, Charles Holderman

2015-present, Samuel Britton

2013-present, Mikahl Banwarth-Kuhn

2013-present, Kevin Tsai

University of Notre Dame:

2012-17, Francesco Pancaldi (Applied Mathematics)

2012-17, Shant Mahserejian (Applied Mathematics)

2012-17, Aboutaleb Amiri (Physics)

2010-2016, Timur Kupaev (Applied Mathematics)

2011-2015, Amy Buchmann, Wenzhao Sun (Applied Mathematics)

2010-2014, Chunlei Li (Applied Mathematics) (research department at Amazon)

2010-2015, Cameron Harvey (Physics) (Director of Academics at Visible Music College)

2008-2013, Joshua Lioi (Applied Mathematics) (postdoc associate, U Arizona)

2008-2013, Huijing Du (Applied Mathematics) (postdoc associate, UC Irvine)

2007-2012, Yuan Liu (Applied Mathematics) (postdoc associate, Michigan State U)

2005-2012, Tanya Salyers (Applied Mathematics) (currently at Amazon)

2005-2010, Richard Gejji (Applied Mathematics) (currently at the Department of Defense)

2005-2009, Jianfeng Zhu (Applied Mathematics) (analyst, Mendoza College of Business, University of Notre Dame)

2005-2009, Yilin Wu (Physics) (Assistant Professor, Chinese University of Hong Kong))

2005-2008, Scott Christley (Computer Science) (currently research scientist, University of Chicago)

2003-2008, Matt Rissler (Mathematics) (Assistant Professor of Mathematics, Loras College, Iowa)

2003-2008 Ivan Gregoretti (Biochemistry) (currently at Cell Signaling Technology, Boston, MA)

2000-2004, Maria Byrne (Mathematics) (Associate Professor of Mathematics, University of South Alabama)

1995-2000, Charles Miller (Mathematics) (Twitter, Computer Security Department)

Master's Students (separate from Ph.D. Program), University of Notre Dame:

2012 - 2013, Tomas Collins, ESTEEM Professional Science and Engineering Master's (PSEM) Program

2012 - 2013, Natasha Flores, ESTEEM Professional Science and Engineering Master's (PSEM) Program

2011-2012, Stephen Gaudet, ESTEEM Professional Science and Engineering Master's (PSEM) Program

2008-2010, Faruck Morcos (also received Ph.D. in Computer Science and Engineering)

2007-2008, Zhenyu Shi; 2005-2006, Fang Qi; 2004-2005, Xuelian Zhu; 2002-2004, Mihaela Vajiac; 1999-2001, Olga Vasillieva

1998-99, Andrea Bieberich

Undergraduate Students, REU Program:

Summer 2017, Andrew Wittaker, Plant biology project

Summer 2011, John Riley, NSF REU, Myxobacteria project

Summer 2008, Jose Garcia (McNair Scholar), David Sheehan (NSF REU Nano/Bio Summer School)

Summer 2007, Fernando Monjarez (McNair Scholar, now in a Ph.D. program in physics at Texas Tech University), Jeston Edwin Greenwood

Summer 2006, Anudha Mittal (NSF REU Nano/Bio Summer School)

2005-2006: James Boyle

2003-2004: Adam Willis

Summer 2004, Michael Bell

High School Teachers: Summers 2008-2015

Helene Dauerty, Physics teacher, Central High School, Elkhart, Indiana

Tom Finke, Lead Mathematics and Science Teacher, Associate Head of School, Trinity School at Greenlawn, South Bend, Indiana

Mike Sinclair, Physics and mathematics instructor, Kalamazoo Area Mathematics and Science Center, Kalamazoo, Michigan

Professional Societies:

Society for Mathematical Biology

Society for Industrial and Applied Mathematics (SIAM)

American Association for the Advancement of Science (AAAS)

Biophysical Society

Organization of Conferences:

Co-organizer, November 15-16, 2017, Conference on "Multi-scale systems biology methods for studying biomedical processes in patients under stress or with chronic or acute diseases, UC Riverside: <http://icqmb.ucr.edu/>

Co-organizer, November 4, 2017, Special Session: Stochastic and Multi-scale Models in Mathematical Biology, Analysis and Simulations, at the AMS Fall Western Sectional Meet-

ing, University of California, Riverside, CA.

Co-organizer, Interdisciplinary Workshop on Multi-scale Modeling of Complex System in Developmental & Plant Biology, University of California, Riverside, December 15, 2016

Co-organizer, Workshop 2: Modelling of Tissue Growth and Form, NSF Mathematical Biology Institute, Columbus, OH, March 6-10, 2017

Co-organizer, Symposium on the Multi-scale Study of Cancer, Annual Meeting of the American Association for Advancement of Science (AAAS), February 14-18, 2013, Boston.

Co-organizer, Workshop on the Physics of Bacterial Communities, June 11th - 12th, 2012, sponsored by the Argonne National Laboratory and University of Notre Dame, University of Notre Dame Chicago Commons.

Organizer, Workshop on Interdisciplinary Biomedical Research, July 18-19, 2011, University of Notre Dame London Center, UK.

Organizer, Workshop on Interdisciplinary Biomedical Research, March 1-3, 2009, University of Notre Dame

Organizer, Focus Group Meeting: Multiscale Methods in Biology, NSF Mathematical Biology Institute, Ohio State University, Columbus, OH, November 2-4, 2008

Organizer, Special Session on Multiscale Methods in Biological Modeling: Hybrid Systems and Coarsening Methods, SIAM Meeting in Life Sciences, Montreal, Canada, August 6, 2008

Co-organizer, Workshop on Interdisciplinary Biomedical Research between University of Notre Dame and Indiana University School of Medicine, University of Notre Dame, April 10 - 11, 2008

Organizer, Special Session on Mathematical Biology: Modeling Cancer, AMS Meeting, University of Notre Dame, April 8 - 9, 2006

Co-organizer, Workshop on Stochastic Modelling, Notre Dame, March 24-36, 2006

Co-organizer, Special Session on Mathematical Biology, American Mathematical Society Meeting, Notre Dame, April 7-9, 2006

Co-organizer, Workshop on Applications of Methods of Stochastic Systems and Statistical Physics in Biology, University of Notre Dame, October 28-30, 2005

Member of the Scientific Committee, Managing Complexity - Systems Biology, 7th World Congress of Chemical Engineering (WCCE 2005), Glasgow, Scotland, July 4-7, 2005.

Organizer, Biocomplexity Workshop VIII: Applications of Methods of Stochastic Systems and Statistical Physics to Biology, University of Notre Dame, October 28-30, 2005.

Co-organizer, Special Session on Mathematical Biology, American Mathematical Society

Meeting, University of Notre Dame, April 8-9, 2006.

Co-organizer, Biocomplexity Workshop VII: Unravelling the Function and Kinetics of Biochemical Networks: from Experiments to Systems Biology, IU Bloomington, May 9 - 11, 2005.

Co-organizer, Biocomplexity Workshop VI: Complex Behavior in Unicellular Organisms, IU Bloomington, May 12 - 16, 2004.

Organizer, Biocomplexity Workshop V: Multiscale Modeling in Biology, University of Notre Dame, August 14 - 17, 2003.

Co-organizer, Biocomplexity Workshop IV: Regenerative Biology and Medicine, University of Indiana, Bloomington, May 15 - 18, 2003.

Co-organizer, Biocomplexity Workshop III: The Role of Tissue Mechanics in Biological Responses to Mechanical Loading, University of Notre Dame, November 8-10, 2002.

Member of a Program Committee of the 17th International Symposium on Mathematical Theory of Networks and Systems, University of Notre Dame, August 12-16, 2002.

Co-organizer of a mini-symposium on applications to biology.

Co-organizer, Notre Dame Workshop on Modeling Cytoskeleton and Cell Motility, University of Notre Dame, November 10-11, 2001.

Co-organizer of 2 invited Mini-symposiums on Applications of Nonlinear Dynamical Systems in Biology, Fifth SIAM Conference on Control and its Applications held jointly with the 2001 SIAM Annual Meeting, San Diego, July 11-14, 2001.

Co-organizer, Conference on Nonlinear Problems in Applied Mathematics, University of Notre Dame, April 7, 2000.

Co-organizer, Special session on Nonlinear Waves and Integrable Systems, AMS Meeting, University of Notre Dame, April 8-9, 2000.

Newly Developed Courses:

Graduate Course: Multi-scale modeling in biology and physics

Graduate Course: Topics in Applied Mathematics, Mathematical Biology

Cross listed Course: Nonlinear Dynamical Systems

Cross listed Course: Mathematical and Computational Modeling in Biology and Physics

Undergraduate Seminar on Interdisciplinary Biological Research: Mathematical and Computational Modeling in Biology

Miscellaneous: Married, two sons