

Ken A. Dill

Louis and Beatrice Laufer Endowed Chair of Physical and Quantitative Biology
SUNY Distinguished Professor of Physics, Chemistry, and Applied Math and Statistics
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Stony Brook University, Stony Brook, NY 11794-5252
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Education

1978 – 1981 Postdoctoral Fellow, Chemistry, Stanford University. Advisor--Paul J. Flory
1971 – 1978 Ph.D. Biology Department, UC San Diego, La Jolla. Advisor--Bruno H. Zimm
1966 – 1971 S.B. and S.M., Mechanical Engineering Department, M.I.T.

Professional Experience

2012 - present SUNY Distinguished Professor, Physics and Chemistry, Stony Brook University
2010 – present Louis & Beatrice Laufer Endowed Chair of Physical and Quantitative Biology
2010 – present Director, Laufer Center for Physical & Quantitative Biology, Stony Brook University
2010 – present Professor of Chemistry & Physics, Stony Brook University
2010 Distinguished Professor, Pharmaceutical Chemistry & Biophysics, UC San Francisco
1989 – 2010 Professor, Pharmaceutical Chemistry & Biophysics, UC San Francisco
2001 – 2010 Associate Dean of Research, School of Pharmacy, UC San Francisco
1989 – 2007 Adjunct Professor, Pharmaceutics, University of Utah
1996 – present Faculty Biochemist, Lawrence Berkeley National Lab
1985 – 1989 Associate Professor, Pharmaceutical Chemistry, UC San Francisco
1985 – 1989 Associate Adjunct Professor, University of Utah
1982 – 1985 Assistant Professor, UC San Francisco
1981 – 1982 Assistant Professor, Chemistry, University of Florida, Gainesville

Other Appointments

2017 – present Affiliated Distinguished Professor, Applied Math and Statistics, Stony Brook University

Service to Professional Organizations

2016 Scientific and Academic Advisory Committee review of *Structural Biology*, Weizmann Institute of Science
2011 Gordon Conference on Stochastic Physics in Biology Founding Co-chair
2004 NIBIB Review Board on Intramural Activities
2003 – 2010 Bridging the Sciences Coalition, Founder and Co-Director, a coalition of 15 basic research societies, supporting deep innovation and the Life/Physical Sciences interface.
2003 – 2006 American Physical Society, Physics Policy Committee
2003 – 2006 American Physical Society, Executive Committee, Division of Biological Physics
2002 – 2005 Protein Society, Executive Committee
2001 – 2007 Biophysical Society, Public Affairs Committee, Chair

2001 – 2004	American Physical Society, Division of Biological Physics, Executive Committee
2001 – 2004	American Physical Society, Awards Committee
1998 – 2000	Biophysical Society, Joint Steering Committee for Public Policy
1998 – 1999	Biophysical Society, President
1997 – 2001	NIH BBCA study section
1996	UC San Francisco BWF Graduate Program in Quantitative Biology, Co-director
1996	NIH BBCA study section, ad hoc
1994 – 1995	UCSF Biophysics Program, Acting Director
1994	Protein Society, Nominating Committee, Chair
1993	Biophysical Society, Nominating Committee, Chair
1993	Gordon Conference on Proteins, Co-chair
1993	NIH BBCA study section, ad hoc
1992 – 1995	Biophysical Society, Council
1992	Protein Society, Nominating Committee
1992	Biophysical Society, Executive Board
1991	Biophysical Society, Nominating Committee
1990	AAAS Symposium on Protein Folding, Co-organizer

Service to Editorial/Advisory Boards

Current	Annual Review of Biophysics, Editor (2013-2022), Protein Engineering, Physical Biology, Multiscale Modeling & Simulation, Structure, Biopolymers, Biophysical Chemistry, Journal of Molecular Recognition.
Past	Journal of Chemical Physics (2014-2016), Annual Review of Biophysics, Associate Editor & Board (2009-2013), Protein Science, Annual Reviews of Physical Chemistry, Biophysical Journal, Theochem, Chemical Physics, Current Biology.

Honors

2019	Sackler Biophysics Prize (Tel Aviv University)
2019	Max Delbruck Prize for Biological Physics (American Physical Society)
2018	Dill 70th Festschrift (https://pubs.acs.org/toc/jpcbfbk/122/21)
2013	American Academy of Arts and Science
2012	Distinguished Professor Chemistry and Physics, State University of New York
2012	Emily Gray Award, Biophysical Society
2010	UCSF 53rd Faculty Research Lecturer
2008	National Academy of Sciences, elected Member
2007	Biophysical Society, Distinguished Service Award
2004	Institute of Physics, elected Fellow
2002	Biophysical Society, elected Fellow
1998	Protein Society, first Hans Neurath Award
1997	AAAS, elected Fellow
1991	American Physical Society, elected Fellow
1987	UCSF Academic Senate, Distinguished Teaching Award
1987	UCSF Joseph M. Long Foundation Prize for Excellence in Teaching
1985 – 1989	Pew Scholar

1979 – 1980 Damon Runyon-Walter Winchell Postdoctoral Fellowship
1971 – 1974 National Science Foundation Pre-doctoral Fellowship

Activities at Stony Brook University

Since Oct 1, 2010

Founding and current director of the Laufer Center for Physical & Quantitative Biology—a community of 19 faculty & 3 Fellows from Stony Brook University, Cold-Spring Harbor & Brookhaven National Labs

- Developed Physical Biology Grad Program in Physics, Chemistry, Applied Math & Statistics Departments
- Campuswide opening event, May 2012
- Hosted a national scientific meeting on protein folding, June 2012; northeast regional scientific meeting on protein design, June 2013; physical and quantitative biology retreats April 2015 & April 2017
- Established the Laufer Distinguished Lecture Series

Member of the Advisory Board of the Institute of Chemical Biology & Drug Discovery

Member of the Institute for Cell & Developmental Biology in the Department of Biochemistry & Cell Biology

Member of the Provost's committee on MOOCs

Member of the search committee for a senior faculty member in Biochemistry and Cell Biology

Textbooks

PROTEIN ACTIONS, Principles & Modeling, with co-authors Ivet Bahar, Robert Jernigan, Garland Science Publishing. First edition February 2017.

MOLECULAR DRIVING FORCES, with co-author Sarina Bromberg, Garland Science Publishing. First edition 2003; Second edition 2012. A textbook of physical chemistry used in more than 150 courses worldwide; it won the Emily Gray Award from the Biophysical Society 2012.

Special Lectures

- 2020 Biological Physics Public Lecture (UCLA, 2020)
- 2019 Bing & Esther Humphrey Lecture, Chemistry (U Vermont, 2019)
- 2019 Daniel Kivelson Lecture, Chemistry (UCLA, 2019)
- 2017 Zymeworks ZED talk (Vancouver, 2017)
- 2017 Greater Boston Theoretical Chemistry 3-lecture series (MIT, Harvard U, BU)
- 2016 Fred W & Gladys E Laird Lecture, Chemistry (U British Columbia)
- 2016 Gary K Acker Lecturer, Gibbs Conference (Carbondale, Ill)
- 2016 Cyril N Hinshelwood Six Lectures, Chemistry (Oxford U)
- 2016 International Symposium, "The Technological Revolution in Structural Biology: Impact on Biology and Chemistry (Weizmann Institute of Science)
- 2016 FW Laird Lecture, Chemistry, U British Columbia
- 2016 60th Annual Meeting, Biophysics, Los Angeles
- 2016 NC3 Award Lecture, Chemistry (University Nevada-Lincoln)
- 2015 Sackler Symposium (Yale University)
- 2014 Morris Visiting Fellow, Chemistry (Hamilton College, Clinton, NY)
- 2012 Sackler Symposium (Yale University)
- 2011 Joseph Priestley Lecture (Penn State University)
- 2011 Eminent Scholar Lecture, Chemistry (University of Arizona)
- 2010 Five-campus Lecture Series (University of Massachusetts, Amherst University, Smith College, Mount Holyoke College, Hampshire College)

- 2008 Gary Griffin Lecture (University of New Orleans)
- 2006 Engbretsons Lecture (North Dakota State University)
- 2004 Inaugural Harrison Shull Lecture, Chemistry (Indiana University)
- 2002 Nieuwland-Reilly Lectures, Chemistry (Notre Dame University)
- 2001 Meloche Lecture, Chemistry (University of Wisconsin)
- 1998 Norman Hascoe Lecture, Physics (University Connecticut)
- 1998 Harland G. Wood Lecture, Biochemistry (Case-Western Reserve University)
- 1998 Clayton Foundation Regents Lectures (University of Texas)
- 1997 Keynote Lecture, Texas Folders Meeting
- 1997 Moses Gomberg Lecture (University of Michigan)
- 1996 National Lecture, Biophysical Society
- 1996 Joseph F. Foster Lecture (Purdue University)
- 1996 C.B. Anfinsen Lecture, Johns Hopkins Folding meeting
- 1995 Warren L. McCabe Lecture (North Carolina State University)
- 1993 Jesse W. Beams Lectures (University of Virginia)
- 1993 Wesleyan Lectures (Wesleyan University)
- 1992 Merck Lecture (Purdue University)
- 1991 Dartmouth Lectures (Dartmouth College)
- 1989 Bayer/Mobay Lecture (University of Pittsburgh)

PhD Students

- 1986 – 1990 Linda De Young, Ph.D., Head, Product Development, NuMedii
- 1984 – 1991 Kevin Scott, Ph.D., AstraZeneca
- 1985 – 1991 Darwin O. V. Alonso, Ph.D., retired, University of Washington, Seattle
- 1986 – 1994 David P. Yee, Ph.D., Translational Sci Project Leader, Novartis Inst for BioMedical Research
- 1987 – 1994 Klaus M. Fiebig, Ph.D., Senior Director Strategic Programs, Ontario Bioscience Innovation Organization
- 1991 – 1996 Paul D. Thomas, Ph.D., Associate Professor, Norris Cancer Center, U Southern Cal
- 1991 – 1997 Kevin A. T. Silverstein, Ph.D., RISS Operations Manager and Scientific Lead, Supercomputing Institute, U Minnesota
- 1991 – 1997 Karen E. S. Tang, Ph.D., Biochem, Molec Bio & Biophysics Department, U Minnesota
- 1994 – 1997 David W. Miller, Ph.D., Open Eye Software
- 1994 – 1999 Kent Kirshenbaum, Ph.D. Professor of Chemistry at NYU
- 1994 – 2000 Rachel Brem, Ph.D., Associate Professor, Buck Institute for Aging, Marin, CA
- 1996 – 2000 Jack Schonbrun, Ph.D., Director Software Engineering - Analytics, Pandora
- 1998 – 2001 Noel Southall, Ph.D., NIH, National Center for Advancing Translational Sciences
- 1999 – 2006 John Chodera, Assistant Member, Memorial Sloan-Kettering Cancer Center, NYC
- 2001 – 2006 Vince Voelz, Associate Professor of Chemistry, Temple University
- 2001 – 2006 Byoung-Chul Lee, 23 & Me
- 2004 – 2006 Justin Bradford, Information Systems, Consultant, Genentech
- 2008 – 2013 Gabriel Rocklin, Senior Fellow, Department of Biochemistry, Univ of Washington
- 2008 – 2013 Charlie Kehoe, Software Engineer, Google
- 2009 – 2012 Jack Peterson, Co-founder and lead develop at Augur
- 2009 – 2013 Geoff Rollins, Decision Analytics, Dropbox
- 2010 – 2012 Relly Brandman, Product Manager, Google
- 2012 – 2016 Eliza Guseva, data scientist, Gartner, Inc., Stamford

2012 – 2016 Michael Hazoglou, postdoc, UC San Diego
 2012 – 2017 Mariola Szenk (join with Gabor Balazsi), Translational Partnerships Lead, nreference, Boston)
 2015 – 2016 Valentin Walther (MS student), now a Ph.D. student Aarhus University
 2016 – present Luca Agozzino, postdoc at Stony Brook University
 2016 – present Cong Liu, Stony Brook University
 2016 – present Corey Weistuch, Stony Brook University
 2017 – present Roy Nassar, Stony Brook University
 2019 – present Jonathan Pachter, Stony Brook University
 2020 – present Charles Kocher, Stony Brook University
 2020 – present Chris Fortran, Stony Brook University

Postdoctoral Associates and Junior Fellows

1981 – 1984 Robert S. Cantor, Ph.D., Professor, Dartmouth College
 1983 – 1986 Jeffrey A. Marqusee, Ph.D., Chief Scientist, Noblis, Inc, Falls Church, VA
 1985 – 1989 Kit Lau, Ph.D., Principal, TBS Analytics
 1986 – 1987 Ling-Chu Li, Ph.D.
 1987 – 1998 Hue Sun Chan, Ph.D., Canada Research Chair Professor,
 Department of Biochemistry, University of Toronto
 1989 – 1991 Gregg B Fields, Ph.D., Professor of Chemistry, Florida Atlantic University
 1989 – 1996 Sarina Bromberg, Ph.D., Biophysics writer, editor, illustrator
 1990 – 2000 Kaizhi Yue, Ph.D., Owner, Conformational Search Solutions, San Francisco
 1993 – 1995 Shaojian Sun, Ph.D.
 1994 – 1999 Shi-Jie Chen, Ph.D., Professor of Physics & Astronomy, University of Missouri
 1995 – 1997 Thomas Beutler, Ph.D., Swiss Banking System
 1996 – 1997 Annelise Barron, Ph.D., Associate Professor of Bioengineering, Stanford University
 1997 – 2000 Kenneth Foreman, Ph.D., Coferon, Inc.
 1999 – 2001 Keith Ball, Ph.D., Senior HPC Engineer, RedLine, Falls Church, VA
 1999 – 2001 Adam Lucas, Ph.D., Lecturer, Department of Statistics, Univ of California, Berkeley
 2000 – 2001 Nick Braun, Ph.D.
 2000 – 2002 Thomas Truskett Ph.D., Professor & Chair, Chemical Engineering, Univ of Texas, Austin
 2000 – 2003 Matteo Palassini, Ph.D., Professor of Physics, University of Barcelona, Spain
 2000 – 2003 Thomas Weikl, Ph.D., Group Leader, Max Planck Institute, Potsdam, Germany
 2000 – 2004 Chaok Seok, Professor, Chemistry, Seoul National University, Korea
 2002 – 2006 Kingshuk Ghosh, Ph.D., Associate Professor of Physics, University of Denver
 2003 – 2005 Ilya Chorny, Ph.D., Senior Manager of Product Marketing, Illumina, San Diego, CA
 2003 – 2006 Banu Ozkan, Ph.D., Associate Professor of Physics, Arizona State University
 2004 – 2005 Ke Fan, Ph.D.
 2004 – 2005 Huafeng Xu, Ph.D. DE Shaw Research Co.
 2004 – 2007 Guo-Hong (Albert) Wu, Ph.D.
 2004 – 2007 Julia Hockenmaier, Ph.D., Assoc Prof of Computer Sci, U Illinois, Urbana-Champaign
 2004 – 2008 David Mobley, Ph.D., Associate Professor, Chemistry & Pharm Sci, UC Irvine
 2005 – 2006 Bosco Ho, Ph.D., Scientific and web programmer
 2005 – 2007 M. Scott Shell, Ph.D., Associate Professor of Chemical Engineering, UC Santa Barbara
 2007 – 2010 Jeremy Schmit, Ph.D., Assistant Professor of Physics, Kansas State University
 2007 – 2013 Christopher Fennell, Ph.D., Associate Professor of Chemistry, Oklahoma State Univ
 2007 – 2014 Justin MacCallum, Ph.D., Assoc Prof of Chemistry, Canada Research Chair, Univ of Calgary
 2008 – 2013 Steve Presse, Ph.D., Assistant Professor of Physics, Arizona State University
 2011 – 2013 Daniel W. Farrell, Ph.D., Data Scientist, Facebook

2011 – 2014	Arijit Maitra, Ph.D., Assistant Professor, BML Munjal University, India
2011 – 2014	Libo Li, Ph.D., Associate Professor, South China University of Technology, Guangzhou
2011 – 2014	Arijit Roy, Ph.D., Scientist, TCS Innovation Labs, India
2014 – 2016	Joseph Morrone, Researcher, IBM, Thomas J. Watson Research Center, NY
2016 – 2017	Lane Votapka, Ph.D., Assistant Professor of Chemistry, Point Loma Nazarene University
2010 – present	Alberto Perez, Ph.D., Assistant Professor, Univ Florida, Gainesville
2012 – present	Adam De Graff, Ph.D., Sr Research Scientist, Methuselah Health UK LTD, Cambridge
2013 – present	Emiliano Brini, Ph.D., Laufer Junior Fellow, Stony Brook University
2013 – present	Mantu Santra, Ph.D., Indian Institute of Technology, Goa, India
2013 – present	Jason Wagoner, Ph.D., Laufer Junior Fellow, Stony Brook University
2013 – 2017	Li Guo, Ph.D. (joint with Ron Zuckermann), Jiangsu University, China
2016 – present	James Robertson, Ph.D., Scientist, Janssen Pharmaceutical, Philadelphia
2018 – present	Bhanita Sharma, Stony Brook University
2019 – present	Gregory Dignon, Stony Brook University
2019 – present	Sridip Parui, Stony Brook University
2020 –present	Luca Agozzino, Stony Brook University

Publications

333. C. Liu, E. Brini, A. Perez, K. A. Dill, Computing Ligands Bound to Proteins Using MELD-Accelerated MD. *Journal of Chemical Theory and Computation*, 10.1021/acs.jctc.0c00543 (2020).

332. L. Agozzino, G. Balázsi, J Wang, and K. A. Dill, How Do Cells Adapt? Stories Told in Landscapes. *Annual Review of Chemical and Biomolecular Engineering*, 11: 155-82 (2020).

331. D. Padhorny, K.A. Porter, M. Ignatov, A. Alekseenko, D. Beglov, S. Kotelnikov, R. Ashizawa, I. Desta, N. Alam Z. Sun, E. Brini, K.A. Dill, O. Schueler-Furman, S. Vajda, and D. Kozakov, ClusPro in rounds 38 to 45 of CAPRI: Toward combining template-based methods with free docking. *Proteins: Structure Function and Bioinformatic*, doi:10.1002/prot.25887 (2020).

330. L.R. Mujica-Parodi, A. Amgalan, S.F. Sultan, B. Antal, X. Sun, S. Skiena, A. Lithen, N. Adra, E.-M. Ratai, C. Weistuch, S.T. Govindarajan, H.H. Strey, K.A. Dill, S.M. Stufflebeam, R.L. Veech, and K. Clarke, Diet modulates brain network stability, a biomarker for brain aging, in young adults. *Proc Natl Acad Sci USA*, 117 (11): doi:10.1073/pnas.1913042117 (2020).

329. K. Ghosh, P.D. Dixit, L. Agozzino, K.A. Dill, The Maximum Caliber Variational Principle for Nonequilibria. *Annual Review of Physical Chemistry*, 71: doi: 10.1146/annurev-physchem-071119-040206 (2020).

328. M. Ignatov, C. Liu, A. Alekseenko, Z. Sun, D. Padhorny, S. Kotelnikov, A. Kazennov, I. Grebenkin, Y. Kholodov, I. Kolosvari, A. Perez, K.A. Dill, and D. Kozakov, Monte Carlo on the manifold and MD refinement for binding pose prediction of protein–ligand complexes: 2017 D3R Grand Challenge. *J Comput Aided Mol Des*, 33(1): 119-127 (2019).

327. A. Khramushin, O. Marcu, N. Alam, O. Shimony, D. Padhorny, E. Brini, K.A. Dill, S. Vajda, D. Kozakov, O. Schueler-Furman, Modeling beta-sheet peptide-protein interactions: Rosetta FlexPepDock in CAPRI rounds 38-45. *Proteins: Structure Function and Bioinformatic*, (2019).

326. M. Santra, K. A. Dill, A. M.R. de Graff, Proteostasis collapse is a driver of cell aging and death. *Proc Natl Acad Sci USA*, 116 (44): 22173-8 (2019).

325. J. Wagoner, K.A. Dill, Opposing pressures of speed and efficiency guide the evolution of molecular machines. *Molecular Biology and Evolution*, (2019).
324. L. Agozzino, K. A. Dill, Minimal constraints for maximum caliber analysis of dissipative steady-state systems. *Phys. Rev. E*, (2019).
323. J.C. Robertson, R. Nassar, C. Liu, E. Brini, K.A. Dill, A. Perez, NMR-assisted protein structure prediction with MELDxMD. *Proteins*, doi: 10.1002/prot.25788: (2019).
322. E. Brini, D. Kozakov, K.A. Dill, Predicting Protein Dimer Structures Using MELD × MD. *J. Chem. Theory Comput.*, (2019).
321. J. Wagoner, K.A. Dill, Mechanisms for achieving high speed and efficiency in biomolecular machines. *Proc Natl Acad Sci USA*, 116: 5902-07 (2019).
320. P.D. Dixit, K.A. Dill, Building Markov state models using optimal transport theory. *The Journal of Chemical Physics*, 150: 54105 (2019).
319. T. Urbic, K.A. Dill, Water Is a Cagey Liquid. *J. of the American Chemical Society*, DOI: 10.1021/jacs.8b08856: (2018).
318. L. Agozzino, K. A. Dill, Protein evolution speed depends on its stability and abundance and on chaperone concentrations. *Proc Natl Acad Sci USA*, 115 (37): 9092 (2018).
317. M. Santra, K. A. Dill, A. M.R. de Graff, How Do Chaperones Protect a Cell's Proteins from Oxidative Damage?. *Cell Systems*, 743-751 (2018).
316. K. A. Dill, Colleagues of Ken A. Dill. *The Journal of Physical Chemistry B*, 5267 (2018).
315. K. A. Dill, Autobiography of Ken A. Dill. *The Journal of Physical Chemistry B*, 5263 (2018).
314. B. Hribar-Lee, C. Seok, E. Coutsias, and M. Lukšič, Tribute to Ken A. Dill. *The Journal of Physical Chemistry B*, 5261 (2018).
313. A. Perez, F. Sittel, G. Stock, and K.A. Dill, MELD-Path Efficiently Computes Conformational Transitions, Including Multiple and Diverse Paths. *J. Chem. Theory Comput.*, 14: 2109 (2018).
312. P.D. Dixit, J. Wagoner, C. Weistuch, S. Pressé, K. Ghosh, K.A. Dill,, Perspective: Maximum Caliber is a General Variational Principle for Dynamical Systems. *J. Chem. Phys.*, 148 (1): 10901 (2018).
311. E Brini, CJ Fennell, M Fernandez-Serra, B Hribar-Lee, M Luksic, KA Dill, How water's properties are encoded in its molecular structure and energies. *Chemical Reviews*. (2017).
310. T Urbic, KA Dill, Analytical theory of the hydrophobic effect of solutes in water. *Physical Review E* 96 (3), 032101. (2017).
309. M Szenk, KA Dill, AMR de Graff, Why Do Fast-Growing Bacteria Enter Overflow Metabolism? Testing the Membrane Real Estate Hypothesis. *Cell Systems*. (2017).
308. A Perez, JA Morrone, KA Dill, Accelerating physical simulations of proteins by leveraging external knowledge. *Wiley Interdisciplinary Reviews: Computational Molecular Science*. (2017).
307. M Santra, DW Farrell, KA Dill, Bacterial proteostasis balances energy and chaperone utilization efficiently.

Proceedings of the National Academy of Sciences, 201620646. (2017).

306. KA Dill, MF Holovko, B Hribar-Lee, N Malikova, The scientific life of Vojko Vlachy. *Journal of Molecular Liquids* 228, 1-3. (2017)

305. I Bahar, RL Jernigan, KA Dill, *Protein Actions: Principles and Modeling*. Garland Science. (2017).

304. JA Wagoner, K Dill, Molecular Motors have Evolved to Optimize Thermodynamic Performance. *Biophysical Journal* 112 (3), 279a. (2017).

303. JA Morrone, A Perez, J MacCallum, KA Dill, Computed binding of peptides to proteins with MELD-accelerated molecular dynamics. *Journal of chemical theory and computation* 13 (2), 870-876. (2017).

302. JA Morrone, A Perez, Q Deng, SN Ha, MK Holloway, TK Sawyer, Bradley S Sherborne, Frank K Brown, Ken A Dill, Molecular simulations identify binding poses and approximate affinities of stapled α -helical peptides to MDM2 and MDMX. *Journal of chemical theory and computation* 13 (2), 863-869. (2017).

301. A. Perez, J.A. Morrone, E. Brini, J. L. MacCallum, K.A. Dill, Blind protein structure prediction using accelerated free-energy simulations. *Science Advance*, 2: e1601274 (2016).

300. E. Brini, S.S. Paranehewage, C.J. Fennell, K.A. Dill, Adapting the semi-explicit assembly solvation model for estimating water-cyclohexane partitioning with the SAMPL5 molecules. *Journal of Computer-Aided Molecular Design*, doi:10.1007/s10822-016-9961-9 (2016).

299. K. Ghosh, A.M.R. de Graff, L Sawle and K.A. Dill, Role of Proteome Physical Chemistry in Cell Behavior. *Journal of Physical Chemistry B*, web: (2016).

298. J.A. Wagoner and K.A. Dill, Molecular Motors: Power Strokes Outperform Brownian Ratchets. *Journal of Physical Chemistry B*, DOI: 10.1021/acs.jpcc.6b02776 (2016).

297. A. Maitra and K.A. Dill, Modeling the Overproduction of Ribosomes when Antibacterial Drugs Act on Cells. *Biophysical Journal*, 110: 743-748 (2016).

296. A. Perez, J.A. Morrone, C. Simmerling and K.A. Dill, Advances in free-energy-based simulations of protein folding and ligand binding. *Current Opinion in Structural Biology*, 36: 25-31 (2016).

295. A.M.R de Graff, M.J. Hazoglou, and K.A. Dill, Highly Charged Proteins: The Achilles' Heel of Aging Proteomes. *Structure*, 24: 1 (2016).

294. A. Perez, J. L. MacCallum, E.A. Coutsias, and K.A. Dill, Constraint methods that accelerate free-energy simulations of biomolecules. *Journal of Chemical Physics*, 143: 243143 (2015).

293. P.D. Dixit, A. Jain, G. Stock, and K.A. Dill, Inferring Transition Rates of Networks from Populations in Continuous-Time Markov Processes. *Journal of Chemical Theory and Computation*, DOI: 10.1021/acs.jctc.5b00537 (2015).

292. A. Perez, J. L. MacCallum, E. Brini, C. Simmerling, and K.A. Dill, Grid-Based Backbone Correction to the ff12SB Protein Force Field for Implicit-Solvent Simulations. *Journal of Chemical Theory and Computation*, DOI: 10.1021/acs.jctc.5b00662: (2015).

291. M.J. Hazoglou, V. Walther, P.D. Dixit and K.A. Dill, Communication: Maximum caliber is a general variational principle for nonequilibrium statistical mechanics. *Journal of Chemical Physics*, 143: 51104 (2015).

290. S. Presse, K. Ghosh, J. Lee and K.A. Dill, Reply to C. Tsallis' "Conceptual Inadequacy of the Shore and Johnson Axioms for Wide Classes of Complex Systems". *Entropy*, 17: 5043-5046 (2015).
289. A.Perez, J.L. MacCallum, K. A. Dill, Accelerating molecular simulations of proteins using Bayesian inference on weak information. *PNAS*, 112: 11846-11851 (2015).
288. J.L. MacCallum, A.Perez, K. A. Dill, Determining protein structures by combining semireliable data with atomistic physical models by Bayesian inference. *PNAS*, 112: 6985-6990 (2015) PMID: PMC4460504.
287. A. Maitra, K. A. Dill, Bacterial growth laws reflect the evolutionary importance of energy efficiency. *PNAS*, 112: 406-411 (2015).
286. M. Kastelica, Y.V. Kalyuzhnyib, B. Hribar-Lee, K.A. Dillc and V. Vlachy, Protein aggregation in salt solutions. *PNAS*, 112(10): 6766–6770 (2015).
285. L. Li, K.A. Dill, C.J. Fennell, Testing the semi-explicit assembly model of aqueous solvation in the SAMPL4 challenge. *J Comput Aided Mol Des.*, 28: 259-264 (2014).
284. L. Li, C. J. Fennell, and K. A. Dill, Small molecule solvation changes due to the presence of salt are governed by the cost of solvent cavity formation and dispersion. *Journal of Chemical Physics*, 141: 22D518 (2014).
283. M. Luksic, C. J. Fennell, K. A. Dill, Using Interpolation for Fast and Accurate Calculation of Ion-Ion Interactions. *Journal of Physical Chemistry B Article ASAP*, (2014).
282. G. J. Rocklin, D. L. Mobley, K. A. Dill, P. H. Hunenberger, Calculating the binding free energies of charged species based on explicit-solvent simulations employing lattice-sum methods: An accurate correction scheme for electrostatic finite-size effects. *Journal of Chemical Physics*, 139: 184103 (2013).
281. S. Pressé, K. Ghosh, J. Lee, K. A. Dill, Principles of maximum entropy and maximum caliber in statistical physics. *Rev. Mod. Phys.*, 85 (3): 1115-41 (2013).
280. S. Pressé, K.Ghosh, J.Lee, K. A. Dill, Nonadditive Entropies Yield Probability Distributions with Biases not Warranted by the Data. *Phys. Rev. Lett.*, 111: 180604 (2013).
279. S. Pressé, J.Lee, K. A. Dill, Extracting Conformational Memory from Single-Molecule Kinetic Data. *J. Phys. Chem. B*, 117 (2): 495-502 (2013).
278. A. Roy, A.Perez, K. A. Dill, J. MacCallum, Computing the Relative Stabilities and the Per-Residue Components in Protein Conformational Changes. *Structure*, 22: 168-175 (2013).
277. Rocklin GJ, Mobley DL, Dill KA, Calculating the sensitivity and robustness of binding free energy calculations to force field parameters, *J Chem Theory Comput*. 2013 Jul 9; 9(7):3072-3083
276. Rocklin GJ, Boyce SE, Fischer M, Fish I, Mobley DL, Shoichet BK, Dill KA, Blind Prediction of Charged Ligand Binding Affinities in a Model Binding Site. *J Mol Biol*. 2013 Jul 26. doi:pii: S0022-2836(13)00477-4. 10.1016/j.jmb.2013.07.030.
275. Dill KA, *Annu Rev Biophys*. 2013; 42. doi: 10.1146/annurev-bb-42-050113-100001. Biophysics. Introduction.
274. Rocklin GJ, Mobley DL, Dill KA, Separated topologies--a method for relative binding free energy

calculations using orientational restraints, *J Chem Phys.* 2013 Feb 28;138(8):085104. doi: 10.1063/1.4792251.

273. Kortkhonjia E, Brandman R, Zhou JZ, Voelz VA, Chorny I, Kabakoff B, Patapoff TW, Dill KA, Swartz TE, Probing antibody internal dynamics with fluorescence anisotropy and molecular dynamics simulations, *MAbs.* 2013 Mar-Apr;5(2):306-22. doi: 10.4161/mabs.23651. *Epub* 2013 Feb 8.

272. E. Kortkhonjia, R. Brandman, J. Z. Zhou, V. A. Voelz, I. Chorny, B. Kabakoff, T. W. Patapoff, K. A. Dill, T. E. Swartz, Probing antibody internal dynamics with fluorescence anisotropy and molecular dynamics simulations, *MAbs* 5, (2): 306-322 (2013).

271. K. A. Dill and J. L. MacCallum, The protein folding problem, 50 years on, *Science* 338, 1042-1046 (2012). (PDF) (Full Text Online) (podcast)

270. A. Perez, Z. Yang, I. Bahar, K. A. Dill, and J. L. MacCallum, FlexE: Using elastic network models to compare models of protein structure, *Journal of Chemical Theory and Computation* 8: 3985–3991 (2012). (PDF)

269. C. W. Kehoe, C. J. Fennell, and K. A. Dill, Testing the semi-explicit assembly solvation model in the SAMPL3 community blind test, *Journal of Computer Aided Molecular Design* 26: 563–568 (2012). (PDF)

268. C. J. Fennell, L. Li, and K. A. Dill, Simple liquid models with corrected dielectric constants, *Journal of Physical Chemistry B* 11: 6936–6944, (2012). (PDF)

267. G. L. Butterfoss, B. Yoo, J. N. Jaworski, I. Chorny, K. A. Dill, R. N. Zuckermann, R. Bonneau, K. Kirshenbaum and V. A. Voelz, De novo structure prediction and experimental characterization of folded peptoid oligomers. *Proceedings of the National Academy of Sciences USA*, 109:1 14320-14325 (2012). (PDF)

266. J. D. Schmit and K. A. Dill, Growth rates of protein crystals. *Journal of the American Chemical Society* 134:9, 3934-3937 (2012). (PDF)

265. G. J. Peterson, S. Pressé, K. S. Peterson and K. A. Dill, Simulated evolution of protein-protein interaction networks with realistic topology. *PLoS ONE* 7, e39052 (2012). (PDF)

264. Lukšič M, Urbic T, Hribar-Lee B, Dill KA. Simple model of hydrophobic hydration. *Journal of Chemical Physics B.* 116(21):6177-86 (2012) *Epub* 2012 May 21 PMID:22564051. (PDF)

263. Fennell CJ, Li L, Dill KA. Simple liquid models with corrected dielectric constants. *Journal of Chemical Physics B.* 116(23):6936-44 (2012) *Epub* 2012 Mar 21. PMID: 22397577. (PDF)

262. S. Pressé, K. Ghosh, J. Lee and K.A. Dill, The principles of Maximum Entropy and Maximum Caliber in statistical physics. Preprint.

261. H. Ge, S. Pressé, K. Ghosh, and K. A. Dill, Markov processes follow from the principle of maximum caliber, *Journal of Chemical Physics* 136, 064108 (2012) PMID: 22360170. (PDF)

260. K. A. Dill, America needs more deep innovation, *ASBMB today* (2012) (view online, PDF)

259. Voelz VA, Dill KA, Chorny I. Peptoid conformational free energy landscapes from implicit-solvent molecular simulations in AMBER. *Biopolymers.* 96(5):639-50 (2011) PMID: 21184487. (PDF)

258. K. A. Dill, K. Ghosh, and J. D. Schmit, Physical limits of cells and proteomes, *Proceedings of the National*

- Academy of Sciences USA* **108**: 17876-17882 (2011) PMID: 22006304. (PDF)
257. J. D. Schmit, S. Whitelam, and K. A. Dill. Electrostatics and aggregation: how charge can turn a crystal into a gel. *Journal of Chemical Physics B* **135**: 085103 (2011) PMID: 21895221. (PDF)
256. S. Pressé, K. Ghosh, and K. A. Dill. Modeling Stochastic Dynamics in Biochemical Systems with Feedback Using Maximum Caliber. *Journal of Physical Chemistry B* **115**: 6202--6212 (2011) PMID: 21524067. (PDF)
255. H. Ge, S. Pressé, K. Ghosh, and K. A. Dill, Markov processes follow from the principle of maximum caliber, *J. Chem. Phys.* **136**, 064108 (2012).
254. K. A. Dill, America needs more deep innovation, *ASBMB today*, January (2012)
253. K. A. Dill, K. Ghosh, and J. D. Schmit, Physical limits of cells and proteomes, *Proceedings of the National Academy of Sciences USA* **108**: 17876-17882 (2011).
252. C.J. Fennell, K.A. Dill, Physical Modeling of Aqueous Solvation. *Journal of Statistical Physics* **145**, 209--226 (2011).
251. J. D. Schmit, S. Whitelam, and K. A. Dill. Electrostatics and aggregation: how charge can turn a crystal into a gel. *Journal of Chemical Physics* **135**: 085103 (2011).
250. S. Pressé, K. Ghosh, and K. A. Dill. Modeling Stochastic Dynamics in Biochemical Systems with Feedback Using Maximum Caliber. *Journal of Physical Chemistry B* **115**: 6202--6212 (2011).
249. C. J. Fennell and K. A. Dill. Physical Modeling of Aqueous Solvation. *J Stat Phys*: 1--18 (2011).
248. C. J. Fennell, C. W. Kehoe, and K. A. Dill. Modeling aqueous solvation with semi-explicit assembly. *Proceedings of the National Academy of Sciences USA* **108** (8): 3234--3239 (2011).
247. J. Schmit, K. Ghosh and K. A. Dill. What drives amyloid molecules to assemble into oligomers and fibrils? *Biophysical Journal* **100** (2): 450--458 (2011).
246. K. Ghosh and K. A. Dill. Cellular proteomes have broad distributions of protein stability. *Biophysical Journal* **99** (12): 3996--4002 (2010).
245. B. Hribar-Lee, K. A. Dill, and V. Vlachy. Receptacle model of salting-in by tetramethylammonium ions. *Journal of Physical Chemistry B* **114** (46): 15085-15091 (2010).
244. S. Pressé, K. Ghosh, R. Phillips, and K. A. Dill. Dynamical fluctuations in biochemical reactions and cycles. *Physical Review E* **82** (3): 031905 1-6 (2010).
243. G. J. Peterson, S. Pressé and K. A. Dill. Nonuniversal power law scaling in the probability distribution of scientific citations. *Proceedings of the National Academy of Sciences USA* **107** (37): 16023--16027 (2010).
242. T. Urbic and K. A. Dill. A Statistical Mechanical Theory for a Two-dimensional Model of Water. *Journal of Physical Chemistry* **132** : 224507 (2010).
241. J. D. Schmit and K. A. Dill. The Stabilities of Protein Crystals. *Journal of Physical Chemistry B* **114** : 4020--4027 (2010).
240. C. J. Fennell, C. Kehoe, and K. A. Dill. Oil/Water Transfer Is Partly Driven by Molecular Shape, Not Just Size. *Journal of the American Chemical Society* **132** (1): 234--240 (2010).
239. E. Kortkhonjia, R. Brandman, I. Chorny, J. Kissman, T. Patapoff, B. Kabakoff, K. A. Dill, and T. Swartz. Solution Dynamics of Monoclonal Antibodies: Experimental and Computational Approach, *Biophysical*

- Journal* **98**, (Issue 3, Supplement 1): 443a (2010).
238. J. L. MacCallum, L. Hua, M. J. Schnieders, V. J. Pande, M. P. Jacobson, and K. A. Dill. Assessment of the protein-structure refinement category in CASP8. *Proteins* **77** (S9): 66--80 (2009).
237. A. Bizjak, T. Urbic, V. Vlachy, and K. A. Dill. Theory for the three-dimensional Mercedes-Benz model of water. *Journal of Physical Chemistry* **131**: 194504 1--7 (2009).
236. S. E. Boyce, D. L. Mobley, G. J. Rocklin, A. P. Graves, K. A. Dill, and B. K. Shoichet. Predicting ligand binding affinity with alchemical free energy methods in a polar model binding site. *Journal of Molecular Biology* **394** (4): 747--763 (2009).
235. C. J. Fennell, A. Bizjak, V. Vlachy, and K. A. Dill, S. Sarupria, S. Rajamani, and S. Garde. Additions and Corrections to Ion pairing in molecular simulations of aqueous alkali halide solutions. *Journal of Physical Chemistry B* **113**: 14837--14838 (2009).
234. M. Cruchok, V. Vlachy, and K. A. Dill. Explicit-water molecular dynamics study of a short-chain 3,3 ionene in solutions with sodium halides. *Journal of Physical Chemistry* **130**: 134903 (2009).
233. K. Ghosh and K. A. Dill. Computing protein stabilities from their chain lengths. *Proceedings of the National Academy of Sciences USA* **106** (26): 10649--10654 (2009).
232. D. Wu, K. Ghosh, M. Inamdar, H. J. Lee, S. Fraser, K. A. Dill, and R. Phillips. Trajectory approach to two-state kinetics of single particles on sculpted energy landscapes. *Physical Review Letters* **103** (5): 050603(1-4) (2009).
231. C. J. Fennell, A. Bizjak, V. Vlachy, and K. A. Dill. Ion pairing in molecular simulations of aqueous alkali halide solutions. *Journal of Physical Chemistry B* **113**: 6782--6791 (2009).
230. D. L. Mobley, C. L. Bayly, M. D. Cooper, M. R. Shirts, and K. A. Dill. Small molecule hydration free energies in explicit solvent: an extensive test of fixed-charge atomistic simulations. *Journal of Chemical Theory and Computation* **5**: 350--358 (2009).
229. M. S. Shell, S. B. Ozkan, V. Voelz, G. A. Wu, and K. A. Dill. Blind Test of Physics-Based Prediction of Protein Structures. *Biophysical Journal* **96**: 917--924 (2009).
228. K. Ghosh and K. A. Dill. Theory for Protein folding cooperativity: Helix Bundles. *Journal of the American Chemical Society* **131** (6): 2306--2312 (2009).
227. D. L. Mobley and K. A. Dill. Binding of Small-Molecule Ligands to Proteins: "What You See" Is Not Always "What You Get". *Structure* **17**: 489 -- 498 (2009).
226. B. Hribar-Lee, V. Vlachy, K. A. Dill. Modeling Hofmeister Effects. *Acta Chimica Slovenica* **56**: 196 -- 202 (2009).
225. V. A. Voelz, M. S. Shell, K. A. Dill. Predicting Peptide Structures in Native Proteins from Physical Simulations of Fragments. *PLOS Computational Biology* **5** (2): e1000281, 1--12 (2009).
224. M. S. Shell, R. Ritterson, K. A. Dill. A Test on Peptide Stability of AMBER Force Fields with Implicit Solvation. *Journal of Physical Chemistry B* **112**: 6878--6886 (2008).
223. G. Albert Wu, E. A. Coutsiias, K. A. Dill. Iterative Assembly of Helical Proteins by Optimal Hydrophobic Packing. *Structure* **16**: 1257--1266 (2008).

222. Byoung-Chul Lee, T.K. Chu, K. A. Dill, and R. N. Zuckermann. Biomimetic Nanostructures: Creating a high-affinity Zinc-binding site in a folded nonbiological polymer. *Journal of the American Chemical Society* **130** : 8847--8855 (2008).
221. Tjasa Urbic, Tomaz Urbic, F. Avbelj, and K. A. Dill. Molecular simulations find stable structures in fragments of Protein G. *Acta Chimica Slovenica* **55**: 385--395 (2008).
220. G. Stock, K Ghosh, and K. A. Dill. Maximum Caliber: A variational approach applied to two-state dynamics. *Journal of Physical Chemistry* **128** (19): 194102 (2008).
219. K. A. Dill, S. B. Ozkan, M. S. Shell, and T. R. Weikl. The protein folding problem. *Annual Review of Biophysics* **37**:289--316 (2008).
218. D. L. Mobley, A.E. Barber II, C. J. Fennell, and K. A. Dill. Charge asymmetries in hydration of polar solutes. *Journal of Physical Chemistry B* **112** : 2405--2414 (2008). ([doi](#))
217. D. L. Mobley, J. D. Chodera, and K. A. Dill. Treating entropy and conformational changes in implicit solvent simulations of small molecules. *Journal of Physical Chemistry B* **112**(3) : 938--946 (2008). ([doi](#))
216. K. Ghosh, S. B. Ozkan, and K. A. Dill. The ultimate speed limit to protein folding is conformational searching. *Journal of the American Chemical Society* **129**: 11920--11927 (2007).
215. T. Urbic, V Vlachy, Y. V. Kalyuzhnyi, and K. A. Dill. An improved thermodynamic perturbation theory for Mercedes-Benz water. *Journal of Chemical Physics* **127**: 174511 (1-4) (2007).
214. T. Urbic, V Vlachy, Y. V. Kalyuzhnyi, and K. A. Dill. Theory for the solvation of nonpolar solutes in water. *Journal of Chemical Physics* **127**: 174505 (1-9) (2007).
213. D. L. Mobley, J. D. Chodera, and K. A. Dill. Confine and Release: Obtaining Correct Binding Free Energies in the Presence of Protein Conformational Change. *Journal of Chemical Theory and Computation* **3** (4): 1231-1235 (2007).
212. A. Bizjak, T. Urbic, V. Vlachy, and K.A. Dill. The three-dimensional "Mercedes Benz" model of water. *Acta Chimica Slovenica* **54**: 532--537 (2007).
211. A. Lucas, L. Huang, A. Joshi and K.A. Dill. Statistical mechanics of helix bundles using a dynamic programming approach. *Journal of the American Chemical Society* **129**: 4272--4281 (2007).
210. D.L. Mobley, A.P. Graves, J.D. Chodera, A.C. McReynolds, B.K. Shoichet and K.A. Dill. Predicting absolute ligand binding free energies to a simple model site. *Journal of Molecular Biology* **371** (4): 1118--1134 (2007).
209. S. Banu Ozkan, G. Albert Wu, J.D. Chodera and K.A. Dill. Protein folding by zipping and assembly. *Proceedings of the National Academy of Sciences USA* **104** (29): 11987--11992 (2007).
208. K.A. Dill, S. Banu Ozkan, T.R. Weikl, J.D. Chodera and V.A. Voelz. The protein folding problem: when will it be solved?. *Current Opinion in Structural Biology* **17**: 342--346 (2007).
207. J.D. Chodera, N. Singhal, V.S. Pande, K.A. Dill, and W.C. Swope. Automatic discovery of metastable states for the construction of Markov models of macromolecular conformational dynamics. *Journal of Chemical Physics* **126**: 155101 (2007).
206. J.A. Bradford and K.A. Dill. Stochastic innovation as a mechanism by which catalysts might self-assemble into chemical reaction networks. *Proceedings of the National Academy of Sciences USA* **104** (24): 10098--10103 (2007).
205. K.A. Dill, A. Lucas, J. Hockenmaier, L. Huang, D. Chiang, and A. Joshi. Computational Linguistics: a new Curriculum Vitae, Ken A. Dill

- tool for exploring biopolymer structures and statistical mechanics. *Polymer* **48**: 4289--4300 (2007).
204. V.A. Voelz and K.A. Dill Exploring zipping and assembly as a protein folding principle. *Proteins: Structure Function and Bioinformatics* **66** (4):877--888 (2007).
203. E. Seitaridou, M.M. Inamdar, R. Phillips, K. Ghosh, and K.A. Dill Measuring flux distributions for diffusion in the small-numbers limit. *Journal of Physical Chemistry B* **111**: 2288--2292 (2007).
202. D.L. Mobley, E. Dumont, J.D. Chodera, and K.A. Dill Comparison of charge models for fixed-charge force fields: small-molecule hydration free energies in explicit solvent. *Journal of Physical Chemistry B* **111**: 2242-2254 (2007).
201. T.R. Weikl and K.A. Dill. Transition states in protein-folding kinetics: the structural interpretation of Phi values. *Journal of Molecular Biology* **365**: 1578--1586 (2007).
200. J. Hockenmaier, A.K. Joshi, K.A. Dill. Routes are trees: The parsing perspective on protein folding. *Proteins: Structure, Function, and Bioinformatics* **66**: 1--15 (2007).
199. J.D. Chodera, W.C. Swope, J.W. Pitera, C. Seok, and K.A. Dill. Use of the weighted histogram analysis method for the analysis of simulated and parallel tempering simulations. *Journal of Chemical Theory and Computation* **3**: 26--41 (2007).
198. J.D. Chodera, W.C. Swope, J.W. Pitera, K.A. Dill. Long-time protein folding dynamics from short-time molecular dynamics simulations. *Multiscale Modeling & Simulation* **5** (4): 1214--1226 (2006).
197. B. Hribar-Lee and K.A. Dill. Modeling Simple Alcohols in Two Dimensions. *Acta Chimica Slovenica* **53**: 257--263 (2006).
196. D.L. Mobley, J.D. Chodera, and K.A. Dill. On the use of orientational restraints and symmetry corrections in al-chemical free energy calculations. *Journal of Chemical Physics* **125** (8): 084902 (2006).
195. T. Urbic, V. Vlachy, and K.A. Dill. Confined water: a Mercedes-Benz model study. *Journal of Physical Chemistry B* **110** (10): 4963--70 (2006).
194. B.K. Ho and K.A. Dill. Correction: Folding very short peptides using molecular dynamics. *PLOS Computational Biology* **2** (5): 476--477 (2006).
193. B.K. Ho and K.A. Dill. Folding very short peptides using molecular dynamics. *PLOS Computational Biology* **2** (4):1--10 (2006).
192. K. Ghosh, K.A. Dill, M.M. Inamdar, E. Seitaridou, and R. Phillips. Teaching the principles of statistical dynamics. *American Journal of Physics* **74** (2): 123--133 (2006).
191. D. Chaing, Aravind K. Joshi, and K. A. Dill. A grammatical theory for the conformational changes of simple helix bundles. *Journal of Computational Biology* **13** : 21-42 (2006).
190. E. A. Coutsias, Chaok Seok, Michael J. Wester, and K. A. Dill. Resultants and Loop Closure. *International Journal of Quantum Chemistry* **106**: 176-189 (2006).
189. I. Chorny, K.A. Dill, M.P. Jacobson. Surfaces Affect Ion Pairing. *Journal of Physical Chemistry* **109** (50): 24056--60 (2005).
188. H. Xu and K.A. Dill. Water's Hydrogen Bonds in the Hydrophobic Effect: A Simple Model. *Journal of Physical Chemistry B* **109** (49): 23611--7 (2005).
187. Byoung-Chul Lee, Ronald N. Zuckermann, and K. A. Dill. Folding a Nonbiological Polymer into a Compact Multihelical Structure. *Journal of the American Chemical Society* **127**: 10999--11009 (2005).

186. Claudia Merlo, K. A. Dill, and Thomas R. Weikl. Φ Values in protein-folding kinetics have energetic and structural components. *Proceedings of the National Academy of Sciences USA* **102** (29): 10171--10175 (2005).
185. Sheila S. Jaswal, Stephanie M.E. Truhlar, K. A. Dill and D. A. Agard. Comprehensive Analysis of Protein Folding Activation Thermodynamics Reveals a Universal Behavior Violated by Kinetically Stable Proteases. *Journal of Molecular Biology* **347** (2): 355-366 (2005).
184. Bosco K. Ho, E. A. Coutsias, Chaok Seok, K. A. Dill. The flexibility in the Proline ring couples to the protein back- bone. *Protein Science* **14**: 1011-1018 (2005).
183. K. A. Dill, T. Truskett, V. Vlachy, B. Hribar-Lee. Modeling water, the hydrophobic effect, and ion solvation. *Annual Review of Biophysics & Biomolecular Structure* **34**: 173-199 (2005).
182. E. A. Coutsias, Chaok Seok, K. A. Dill. Using quaternions to calculate RMSD. *Journal of Computational Chemistry* **25**: 1849-1857 (2004).
181. T. Urbic, V. Vlachy, O. Pizio, K. A. Dill. Water-like fluid in the presence of Lennard-Jones obstacles: predictions of an associative replica Ornstein-Zernike theory. *Journal of Molecular Liquids* **112**: 71-80 (2004).
180. V. Vlachy, B. Hribar-Lee, Yu.V. Kalyuzhnyi, K. A. Dill. Short-range interactions: from simple ions to polyelectrolyte solutions. *Current Opinion in Colloid and Interface Science* **9**: 128-132 Elsevier (2004).
179. Thomas R. Weikl, Matteo Palassini, K. A. Dill. Cooperativity in 2-state Protein Folding Kinetics. *Protein Science* **13**(3): 822-829 (2004).
178. Thomas Horn, Byoung-Chul Lee, K. A. Dill, and Ronald N. Zuckermann. Incorporation of Chemoselective Functionalities into Peptoids via Solid-Phase Submonomer Synthesis. *Bioconjugate Chemistry* **15**: 428--435 (2004).
177. E.A. Coutsias, C. Seok, M.P. Jacobson, and K.A. Dill. A Kinematic View of Loop Closure. *Journal of Computational Chemistry* **25**: 510-528 (2004).
176. Lucas, A. and K.A. Dill. Statistical mechanics of pseudoknot polymers. *Journal of Chemical Physics* **119** (4): 2414-2421 (2003).
175. Wu, C.W., K. Kirschenbaum, T.J. Sanborn, J.A. Patch, K. Huang, K.A. Dill, R.N. Zuckermann, and A.E. Barron. Structural and Spectroscopic Studies of Peptoid Oligomers with α -Chiral Aliphatic Side Chains. *Journal of the American Chemical Society* **125**: 13525-13530 (2003).
174. Schonbrun, J., and K.A. Dill. Fast Protein Folding Kinetics. *Proceedings of the National Academy of Sciences USA* **100** (22): 12678-12682 (2003).
173. Weikl, T.R., and K.A. Dill. Folding Kinetics of Two-state Proteins: Effect of Circularization, Permutation, and Crosslinks. *Journal of Molecular Biology* **332** (4): 953-963 (2003).
172. Urbic, T., V. Vlachy, Y.V. Kalyuzhnyi, and K.A. Dill. Orientation-dependent Integral Equation Theory for a Two- dimensional Model of Water. *Journal of Chemical Physics* **118**: 5516 (2003).
171. Weikl, T.R., and K.A. Dill. Folding Rates and Low-entropy-loss Routes of Two-state Proteins. *Journal of Molecular Biology* **329** (3): 585-598 (2003).
170. Ozkan, S. Banu, K.A. Dill, and I. Bahar. Computing the Transition State Populations in Simple Protein Models. *Bio- polymers* **68** (1): 35-46 (2003).
169. Southall, N.T., and K.A. Dill. Potential of Mean Force Between Two Hydrophobic Solutes in Water.

Biophysical Chemistry **101--102**: 295-307 (2002).

168. Truskett, T.M., and K.A. Dill. A Simple Statistical Mechanical Model of Water. *Journal of Physical Chemistry B* **106**:11829-11842 (2002).

167. Seok, C., J.B. Rosen, J.D. Chodera, and K.A. Dill. MOPED: Method for Optimizing Physical Energy Parameters Using Decoys. *Journal of Computational Chemistry* **24**: 89-97 (2002).

166. Hribar, B., N.T. Southall, V. Vlachy, and K.A. Dill. How ions affect the structure of water. *Journal of the American Chemical Society* **124** (41): 12302-12311 (2002).

165. Ozkan, S.B., K.A. Dill, and I. Bahar. Fast-Folding Protein Kinetics, Hidden Intermediates, and the Sequential Stabilization Model. *Protein Science* **11**: 1958-1970 (2002).

164. Truskett, T.M., and K.A. Dill. Predicting Water's Phase Diagram and Liquid-State Anomalies. *Journal of Chemical Physics* **117**: 5101-5104 (2002).

163. Southall, N.T., K.A. Dill, and A.D.J. Haymet. A View of the Hydrophobic Effect. *Journal of Physical Chemistry B* **106**:521-533 (2002).

162. Urbic, T., V. Vlachy, Yu.V. Kalyuzhnyi, N.T. Southall, and K.A. Dill. A Two-Dimensional Model of Water: Solvation of Nonpolar Solutes. *Journal of Chemical Physics* **116**: 723-729 (2002).

161. Ball, K.D., B. Erman, and K.A. Dill. The Elastic Net Algorithm and Protein Structure Prediction. *Journal of Computational Chemistry* **23**: 77-83 (2002).

160. Erkip, A., B. Erman, C. Seok, and K.A. Dill. Parameter Optimization for the Gaussian model of protein folding. *Polymer* **43**: 495-501 (2002).

159. Kevin A. T. Silverstein, K. A. Dill, A.D.J. Haymet. Hydrophobicity in a simple model of water: Entropy penalty as a sum of competing terms via full, angular expansion. *Journal of Chemical Physics* **114** (4): 6303-6314 (2001).

158. Zhou, H.-X., and K.A. Dill. Stabilization of Proteins in Confined Spaces. *Biochemistry* **40**: 11289-11293 (2001).

157. Ozkan, S.B., I. Bahar, and K. A. Dill. Transition States and the Meaning of f-Values in Protein Folding Kinetics. *Nature Structural Biology* **8**: 765-769 (2001).

156. Kalyuzhnyi, Y.V., Vojko Vlachy, and K. A. Dill. Hydration of Simple Ions. Effect of the Charge Density. *Acta Chimica Slovenica* **48**: 309-316 (2001).

155. Phillips, A.T., J.B. Rosen, and K.A. Dill. Convex Global Underestimation for Molecular Structure and Prediction. In *Local To Global Optimization*, P.M. Pardalos, et al., eds. Kluwer Academic Publishers, 1-18, (2001).

154. Liang, Jie, and K. A. Dill. Are Proteins Well-Packed? *Biophysical Journal* **81**: 751-766 (2001).

153. Tostesen, Eivind, Shi-Jie Chen, and K. A. Dill. RNA Folding Transitions and Cooperativity. *Journal of Physical Chemistry B* **105**: 1618-1630 (2001).

152. Southall, N.T., and K.A. Dill. Response to "Comment on 'The Mechanism of Hydrophobic Solvation Depends on Solute Radius,' *Journal of Physical Chemistry B* 2000, 104, 1326" *Journal of Physical Chemistry B* **105**: 2082-2083 (2001).

151. Phillips, A.T., J.B. Rosen, and K.A. Dill. Energy Landscape Projections of Molecular Potential Functions. In *Nonconvex Optimization and Its Applications*. C.A. Floudas and P.M. Pardalos, eds. Kluwer Academic Publishers, 47-55 (2000).

150. Rosen, J.B., A.T. Phillips, S.Y. Oh, and K.A. Dill. A Method for Parameter Optimization in Computational Biology. *Biophysical Journal* **79**: 2818-2824 (2000).
149. Wang, T., J. Miller, N.S. Wingreen, C. Tang, and K.A. Dill. Symmetry and Designability for Lattice Protein Models. *Journal of Chemical Physics* **113**: 8329-8336 (2000).
148. Yue, K., and K.A. Dill. Constraint-Based Assembly of Tertiary Protein Structures from Secondary Structure Elements. *Protein Science* **9**: 1935-1946 (2000).
147. Silverstein, K.A.T., A.D.J. Haymet, and K.A. Dill. The Strength of Hydrogen Bonds in Liquid Water and Around Nonpolar Solutes. *Journal of the American Chemical Society* **122**: 8037-8041 (2000).
146. Brem, R., H.S. Chan, and K.A. Dill. Extracting microscopic energies from oil-phase solvation experiments. *Journal of Physical Chemistry B* **104**: 7471-7482 (2000).
145. Southall, N.T., and K.A. Dill. The Mechanism of Hydrophobic Solvation Depends on Solute Radius. *Journal of Physical Chemistry B* **104**: 1326-1331 (2000).
144. Chen, S.-J., and K.A. Dill. RNA Folding Energy Landscapes. *Proceedings of the National Academy of Sciences USA* **97**: 646-651 (2000).
143. Urbic, T., V. Vlachy, Yu.V. Kalyuzhnyi, N.T. Southall, and K.A. Dill. A Two-Dimensional Model of Water: Theory and Computer Simulations. *Journal of Chemical Physics* **112**: 2843-2848 (2000).
142. Erman, B., and K.A. Dill. Gaussian Model of Protein Folding. *Journal of Chemical Physics* **112**: 1050-1056 (2000).
141. Dill, K.A., J. Deisenhofer, G.R. Fleming, H. Frauenfelder, K. Gerwert, J.A. McCammon, and H. Michel. Group Report: What Do We Need to Know About Proteins and Nucleic Acids? In *Simplicity and Complexity in Proteins and Nucleic Acids*. Edited by H. Frauenfelder, J. Deisenhofer, and P.G. Wolynes. Dahlem University Press, 1999.
140. Young, M., K. Kirshenbaum, K.A. Dill, and S. Highsmith. Predicting Conformational Switches in Proteins. *Protein Science* **8**: 1752-1764 (1999).
139. Foreman, K.W., A.T. Phillips, J.B. Rosen, and K.A. Dill. Comparing Search Strategies for Finding Global Optima on Energy Landscapes. *Journal of Computational Chemistry* **20**: 1527-1532 (1999).
138. Tang, K.E.S., and K.A. Dill. How Experiments See Fluctuations of Native Proteins: Perspective from an Exact Model. *International Journal of Quantum Chemistry* **75**: 147-164 (1999).
137. Silverstein, K.A.T., A.D.J. Haymet, and K.A. Dill. Molecular Model of Hydrophobic Solvation. *Journal of Chemical Physics* **111**: 8000-8009 (1999).
136. Kirshenbaum, K., R.N. Zuckermann, and K.A. Dill. Designing Polymers that Mimic Biomolecules. *Current Opinion in Structural Biology* **9**: 530-535 (1999).
135. Dill, K.A. Strengthening Biomedicine's Roots. *Nature* **400**: 309-310 (1999).
134. Dill, K.A. Polymer Principles and Protein Folding. *Protein Science* **8**: 1166-1180 (1999).
133. Brem, R., and K.A. Dill. The Effect of Multiple Binding Modes on Empirical Modeling of Ligand Docking to Proteins. *Protein Science* **8**: 1134-1143 (1999).
132. Ishikawa K., K. Yue, and K.A. Dill. Predicting the Structures of 18 Peptides Using Geocore. *Protein Science* **8**: 716-721 (1999).

131. Tang, K.E.S. and K.A. Dill. Native Protein Fluctuations: The Conformational-Motion Temperature and the Inverse Correlation of Protein Flexibility with Protein Stability. *Journal of Biomolecular Structure & Dynamics* **16**: 397-411 (1998).
130. Silverstein, K.A.T., K.A. Dill, and A.D.J. Haymet. Hydrophobicity in a Simple Model of Water: Solvation and Hydro- gen Bond Energies. *Fluid Phase Equilibria* **150-151**: 83-90 (1998).
129. Chen, Shi-Jie and K.A. Dill. Theory for the Conformational Changes of Double-Stranded Chain Molecules. *Journal of Chemical Physics* **109**: 4602-4616 (1998).
128. DeVido, D.R., J.G. Dorsey, H.S. Chan, and K.A. Dill. Oil/Water Partitioning Has a Different Thermodynamic Signature When the Oil Solvent Chains Are Aligned Than When They Are Amorphous. *Journal of Physical Chemistry B* **102**:7272-7279 (1998).
127. Silverstein, K.A.T., A.D.J. Haymet, and K.A. Dill. A simple model of water and the hydrophobic effect. *Journal of the American Chemical Society* **120**: 3166-3175 (1998).
126. Armand, P., K. Kirshenbaum, R.A. Goldsmith, S. Farr-Jones, A.E. Barron, K.T.V. Truong, K.A. Dill, D.F. Mierke, F.E. Cohen, R.N. Zuckermann, and E.K. Bradley. NMR determination of the major solution conformation of a peptoid pen- tamer with chiral side chains. *Proceedings of the National Academy of Sciences USA* **95**: 4309-4314 (1998).
125. Kirshenbaum, K., A.E. Barron, R.A. Goldsmith, P. Armand, E.K. Bradley, K.T.V. Truong, K.A. Dill, F.E. Cohen, and R.N. Zuckermann. Sequence-specific polypeptoids: A diverse family of heteropolymers with stable secondary structure. *Proceedings of the National Academy of Sciences USA* **95**: 4303-4308 (1998).
124. Chan, H.S., and K.A. Dill. Protein Folding in the Landscape Perspective: Chevron Plots and Non-Arrhenius Kinetics. *Proteins: Structure, Function, and Genetics* **30**: 2-33 (1998).
123. Armand, P., K. Kirshenbaum, A. Falicov, R.L. Dunbrack, K.A. Dill, R.N. Zuckermann, and F.E. Cohen. Chiral N- substituted glycines can form stable helical conformations. *Folding & Design* **2**: 369-375 (1997).
122. Miller, D.W., and K.A. Dill. Ligand Binding to Proteins: The Binding Landscape Model. *Protein Science* **6**: 2166-2179 (1997).
121. Dill, K.A., J.B. Rosen, and A.T. Phillips. Protein Structure and Energy Landscape Dependence on Sequence Using a Continuous Energy Function. *Journal of Computational Biology* **4**: 227-239 (1997).
120. Chan, H.S., and K.A. Dill. Solvation: How to Obtain Microscopic Energies from Partitioning and Solvation Experiments. *Annual Review of Biophysics and Biomolecular Structure* **26**: 425-459 (1997).
119. Haymet, A.D.J., K.A.T. Silverstein, and K.A. Dill. Hydrophobicity Reinterpreted as "Minimization of the Entropy Penalty of Solvation." *Faraday Discussion* **103**: 117-124 (1997).
118. Dill, K.A., A.T. Phillips, and J.B. Rosen. Molecular Structure Prediction by Global Optimization. In *Developments in Global Optimization*, pp. 217-234. I.M. Bomze, et al., eds. The Netherlands: Kluwer Academic Publishers, 1997.
117. Dill, K.A. Additivity Principles in Biochemistry. Minireview for *The Journal of Biological Chemistry* **272**: 701-704 (1997).
116. Dill, K.A., and H.S. Chan. From Levinthal to Pathways to Funnels: The "New View" of Protein Folding Kinetics. *Nature Structural Biology* **4**: 10-19 (1997).
115. Dill, K.A., and A.R. Fersht. Editorial overview: Folding and Binding. *Current Opinion in Structural Biology* **6**: 1-2 (1996).

114. Dill, K.A. Great Organizing Principles. *Current Biology* **6**: 1204 (1996).
113. Thomas, P.D., and K.A. Dill. An Iterative Method for Extracting Energy-Like Quantities from Protein Structures. *Proceedings of the National Academy of Science USA* **93**: 11628-11633 (1996).
112. Stigter, D., and K.A. Dill. Binding of Ionic Ligands to Polyelectrolytes. *Biophysical Journal* **71**: 2064-2074 (1996).
111. Beutler, T.C., and K.A. Dill. A Fast Conformational Search Strategy for Finding Low Energy Structures of Model Proteins. *Protein Science* **5**: 2037-2043 (1996).
110. Wallace, D.G., and K.A. Dill. Treating Sequence Dependence of Protein Stability in a Mean-Field Model. *Biopolymers* **39**: 115-127 (1996).
109. Chen, S.-J., and K.A. Dill. Structures and Symmetries of Proteins: A Knot Theory Approach. *Journal of Chemical Physics* **104**: 5964-5973 (1996).
108. Thomas, P.D., and K.A. Dill. Statistical Potentials Extracted From Protein Structures: How Accurate Are They? *Journal of Molecular Biology* **257**: 457-469 (1996).
107. Chan, H.S., and K.A. Dill. Comparing Folding Codes for Proteins and Polymers. *Proteins: Structure, Function, and Genetics* **24**: 335-344 (1996).
106. Chan, H.S., and K.A. Dill. A Simple Model of Chaperonin-Mediated Protein Folding. *Proteins: Structure, Function, and Genetics* **24**: 345-351 (1996).
105. Yue, K., and K.A. Dill. Folding Proteins With a Simple Energy Function and Extensive Conformational Searching. *Protein Science* **5**: 254-261 (1996).
104. Sun, S., R. Brem, H.S. Chan, and K.A. Dill. Designing Amino Acid Sequences To Fold With Good Hydrophobic Cores. *Protein Engineering* **9**: 1205-1213 (1995).
103. Krukowski, A.E., H.S. Chan, and K.A. Dill. An Exact Lattice Model of Complex Solutions: Chemical Potentials Depend on Solute and Solvent Shape. *Journal of Chemical Physics* **103**: 10675-10688 (1995).
102. Sun, S., P.D. Thomas, and K.A. Dill. A Simple Protein Folding Algorithm Using a Binary Code and Secondary Structure Constraints. *Protein Engineering* **8**: 769-778 (1995).
101. Dill, K.A., H.S. Chan, and K. Yue. The Protein Folding Problem: Searching Conformations of Compact Chain Molecules. *Macromolecular Symposia* **98**: 615-617 (1995).
100. Chen, S.-J., and K.A. Dill. Statistical Thermodynamics of Double-Stranded Polymer Molecules. *Journal of Chemical Physics* **103**: 5802-5813 (1995).
99. Miller, D.W., and K.A. Dill. A Statistical Mechanical Model for Hydrogen Exchange in Globular Proteins. *Protein Science* **4**: 1860-1873 (1995).
98. Stigter, D., and K.A. Dill. Theory for Radii and Second Virial Coefficients of Polymers. II. Weakly Charged Polyelectrolytes. *Macromolecules* **28**: 5338-5346 (1995).
97. Stigter, D., and K.A. Dill. Theory for Radii and Second Virial Coefficients of Polymers. I. Highly Charged Polyelectrolytes. *Macromolecules* **28**: 5325-5337 (1995).
96. Chan, H.S., S. Bromberg, and K.A. Dill. Models of Cooperativity in Protein Folding. *Philosophical Transactions of the Royal Society, London B* **348**: 61-70 (1995).
95. Dill, K.A., S. Bromberg, K. Yue, K.M. Fiebig, D.P. Yee, P.D. Thomas, and H.S. Chan. Principles of Protein

- Folding - A Perspective From Simple Exact Models. *Protein Science* **4** (4): 561-602 (1995).
94. Yue, K. and K.A. Dill. Forces of Tertiary Structural Organization in Globular Proteins. *Proceedings of the National Academy of Sciences USA* **92**: 146-150 (1995).
93. Dill, K.A. and D. Stigter. Modeling Protein Stability as Heteropolymer Collapse. *Advances in Protein Chemistry* **46**:59-104 (1995).
92. Yue, K., K.M. Fiebig, P.D. Thomas, H.S. Chan, E.I. Shakhnovich, and K.A. Dill. A Test of Lattice Protein Folding Algorithms. *Proceedings of the National Academy of Sciences USA* **92**: 325-329 (1995).
91. Dill, K.A. and K.M. Fiebig. Hydrophobic Zippers: A Conformational Search Strategy for Proteins. In *Statistical Mechanics, Protein Structure, and Protein Substrate Interactions*, p. 109. S. Doniach, ed. New York: Plenum Press (1994).
90. Chan, H.S. and K.A. Dill. Solvation: Effects of Molecular Size and Shape. *Journal of Chemical Physics* **101**: 7007-7026 (1994).
89. Yee, D.P., H.S. Chan, T.F. Havel, and K.A. Dill. Does Compactness Induce Secondary Structure in Proteins? A Study of Poly-Alanine Chains Computed by Distance Geometry. *Journal of Molecular Biology* **241**: 557-573 (1994).
88. Chan, H.S., and K.A. Dill. Transition States and Folding Dynamics of Proteins and Heteropolymers. *Journal of Chemical Physics* **100**: 9238-9257 (1994).
87. Bromberg, S. and K.A. Dill. Side-Chain Entropy and Packing in Proteins. *Protein Science* **3**: 997-1009 (1994).
86. Lattman, E.E., K.M. Fiebig, and K.A. Dill. Modeling Compact Denatured States of Proteins. *Biochemistry* **33**: 6158-6166 (1994).
85. De Young, L.R., A.L. Fink, and K.A. Dill. Aggregation of Globular Proteins. *Accounts of Chemical Research* **26**: 614-620 (1993).
84. Chan, H.S. and K.A. Dill. Energy Landscapes and the Collapse Dynamics of Homopolymers. *Journal of Chemical Physics* **99**: 2116-2127 (1993).
83. Stigter, D. and K.A. Dill. Theory for Second Virial Coefficients of Short DNA. *Journal of Physical Chemistry* **97**:12995-12997 (1993).
82. Yue, K. and K.A. Dill. Sequence-Structure Relationships in Proteins and Copolymers. *Physical Review E* **48**: 2267-2278 (1993).
81. Thomas, P.D. and K.A. Dill. Local and Nonlocal Interactions in Globular Proteins, and Mechanisms of Alcohol Denaturation. *Protein Science* **2**: 2050-2065 (1993).
80. Baker, D., H.S. Chan, and K.A. Dill. Coordinate-Space Formulation of Polymer Lattice Cluster Theory. *Journal of Chemical Physics* **98**: 9951-9962 (1993).
79. Stigter, D. and K.A. Dill. Theory for Protein Solubilities. *Fluid Phase Equilibria* **82**: 237-249 (1993).
78. Yee, D.P. and K.A. Dill. Families and the Structural Relatedness Among Globular Proteins. *Protein Science* **2**: 884-899 (1993).
77. De Young, L.R., K.A. Dill, and A.L. Fink. Aggregation and Denaturation of Apomyoglobin in Aqueous Urea Solutions. *Biochemistry* **32**: 3877-3886 (1993).
76. Fiebig, K.M. and K.A. Dill. Protein Core Assembly Processes. *Journal of Chemical Physics* **98**: 3475-3487

(1993).

75. Dill, K.A. Folding Proteins: Finding a Needle in a Haystack. *Current Opinion in Structural Biology* **3**: 99-103 (1993).
74. Dill, K.A., K.M. Fiebig, and H.S. Chan. Cooperativity in Protein-Folding Kinetics. *Proceedings of the National Academy of Sciences USA* **90**: 1942-1946 (1993).
73. Chan, H.S. and K.A. Dill. The Protein Folding Problem. *Physics Today* **46**: 4-32 (1993). Reprinted in Japanese in Parity, *The Japanese Physical Science Magazine*, December 1993, pp. 24-35.
72. Stickle, D.F., L.G. Presta, K.A. Dill, and G.D. Rose. Hydrogen Bonding in Globular Proteins. *Journal of Molecular Biology* **226**: 1143-1159 (1992).
71. Fields, G.B., D.O.V. Alonso, D. Stigter, and K.A. Dill. Theory for the Aggregation of Proteins and Copolymers. *Journal of Physical Chemistry* **96**: 3974-3981 (1992).
70. Stigter, D., J. Mingins, and K.A. Dill. Phospholipid Interactions in Model Membrane Systems. II. Theory. *Biophysical Journal* **61**: 1616-1629 (1992).
69. Mingins, J., D. Stigter, and K.A. Dill. Phospholipid Interactions in Model Membrane Systems. I. Experiments on Monolayers. *Biophysical Journal* **61**: 1603-1615 (1992).
68. Cole, L.A., J.G. Dorsey, and K.A. Dill. Temperature Dependence of Retention in Reversed-Phase Liquid Chromatography: II. Mobile Phase Considerations. *Analytical Chemistry* **64**: 1324-1327 (1992).
67. Yue, K. and K.A. Dill. Inverse Protein Folding Problem: Designing Polymer Sequences. *Proceedings of the National Academy of Sciences USA* **89**: 4163-4167 (1992).
66. Shortle, D., H.S. Chan, and K.A. Dill. Modeling the Effects of Mutations on the Denatured States of Proteins. *Protein Science* **1**: 201-215 (1992).
65. Miller, R., C.A. Danko, M.J. Fasolka, A.C. Balazs, H.S. Chan, and K.A. Dill. Folding kinetics of proteins and copolymers. *Journal of Chemical Physics* **96**: 768-780 (1992).
64. Alonso, D.O.V., K.A. Dill, and D. Stigter. The Three States of Globular Proteins: Acid Denaturation. *Biopolymers* **31**: 1631-1649 (1991).
63. Kollman, P.A. and K.A. Dill. Decisions in Force Field Development: An Alternative to Those Described by Roterman et al. *Journal of Biomolecular Structure & Dynamics* **8**: 1103-1107 (1991).
62. Chan, H.S., M.R. Wattenbarger, D.F. Evans, V.A. Bloomfield, and K.A. Dill. Enhanced Structure in Polymers at Interfaces. *Journal of Chemical Physics* **94**: 8542-8557 (1991).
61. Naghizadeh, J. and K.A. Dill. Statistical Mechanics of Chain Molecules at Interfaces. *Macromolecules* **24**: 1768-1778 (1991).
60. Alonso, D.O.V. and K.A. Dill. Solvent Denaturation and Stabilization of Globular Proteins. *Biochemistry* **30**: 5974-5985 (1991).
59. Stigter, D., D.O.V. Alonso, and K.A. Dill. Protein Stability: Electrostatics and Compact Denatured States. *Proceedings of the National Academy of Sciences USA* **88**: 4176-4180 (1991).
58. Chan, H.S. and K.A. Dill. "Sequence Space Soup" of Proteins and Copolymers. *Journal of Chemical Physics* **95**: 3775-3787 (1991).

57. Dill, K.A. and D. Shortle. Denatured States of Proteins. *Annual Review of Biochemistry* **60**: 795-825 (1991).
56. Chan, H.S. and K.A. Dill. Polymer Principles in Protein Structure and Stability. *Annual Review of Biophysics and Biophysical Chemistry* **20**: 447-490 (1991).
55. Wattenbarger, M.R., H.S. Chan, D.F. Evans, V.A. Bloomfield, and K.A. Dill. Surface-Induced Enhancement of Internal Structure in Polymers and Proteins. *Journal of Chemical Physics* **93**: 8343 (1990).
54. Dill, K.A. The Meaning of Hydrophobicity. *Science* **250**: 297-298 (1990).
53. Chan, H.S. and K.A. Dill. Origins of Structure in Globular Proteins. *Proceedings of the National Academy of Sciences USA* **87**: 6388 (1990).
52. Dill, K.A. Dominant Forces in Protein Folding. *Biochemistry* **29**: 7133-7155 (1990).
51. Dill, K.A. Review of Lipid and Biopolymer Monolayers at Liquid Interfaces. *Journal of the American Chemical Society* **112**: 1299 (1990).
50. Chan, H.S. and K.A. Dill. The Effects of Internal Constraints on the Configurations of Chain Molecules. *Journal of Chemical Physics* **92**: 3118-3135 (1990).
49. Lau, K.F. and K.A. Dill. Theory for Protein Mutability and Biogenesis. *Proceedings of the National Academy of Sciences USA* **87**: 638-642 (1990).
48. Stigter, D. and K.A. Dill. Charge Effects on Folded and Unfolded Proteins. *Biochemistry* **29**: 1262-1271 (1990).
47. DeYoung, L.R. and K.A. Dill. Partitioning of Nonpolar Solutes into Bilayers and Amorphous n-Alkanes. *Journal of Physical Chemistry* **94**: 801-809 (1990).
46. Chan, H.S. and K.A. Dill. Compact Polymers. *Macromolecules* **22**: 4559-4573 (1989).
45. Stigter, D. and K.A. Dill. Free Energy of Electrical Double Layers: Entropy of Adsorbed Ions and the Binding Polynomial. *Journal of Physical Chemistry* **93**: 6737-6743 (1989).
44. Lau, K.F. and K.A. Dill. A Lattice Statistical Mechanics Model of the Conformational and Sequence Spaces of Proteins. *Macromolecules* **22**: 3986-3997 (1989).
43. Dorsey, J.G. and K.A. Dill. The Molecular Mechanism of Retention in Reversed-Phase Liquid Chromatography. *Chemical Reviews* **89**: 331-346 (1989).
42. Dill, K.A., D.O.V. Alonso, and K. Hutchinson. Thermal Stabilities of Globular Proteins. *Biochemistry* **28**: 5439-5449 (1989).
41. Chan, H.S. and K.A. Dill. Intra-Chain Loops in Polymers: Effects of Excluded Volume. *Journal of Chemical Physics* **90**: 492-509 (1989).
40. Ying, P.T., J.G. Dorsey, and K.A. Dill. Retention Mechanisms of Reversed Phase Liquid Chromatography: Determination of Solute-Solvent Interaction Free Energies. *Analytical Chemistry* **61**: 2540-2546 (1989).
39. Dill, K.A. and D.O.V. Alonso. Conformational Entropy and Protein Stability. In *Protein Structure and Protein Engineering* **39**, pp. 51-58. R. Huber, and E.L. Winnacker, eds. Colloquium-Mosbach der Gesellschaft für Biologische Chemie. Berlin: Springer-Verlag, 1988.
38. Dill, K.A., J. Naghizadeh, and J. A. Marqusee. Chain Molecules at High Densities at Interfaces. *Annual Review of Physical Chemistry* **39**: 425-461 (1988).
37. DeYoung, L. and K.A. Dill. Solute Partitioning into Lipid Bilayer Membranes. *Biochemistry* **27**: 5281-5289

(1988).

36. Dill, K.A. and D. Stigter. Lateral Interactions Among PC and PE Headgroups in Phospholipid Monolayers and Bilayers. *Biochemistry* **27**: 3446-3453 (1988).

35. Stigter, D. and K.A. Dill. Lateral Interactions Among Phospholipid Headgroups at the Heptane/Water Interface. *Langmuir* **4**: 200-209 (1988).

34. Dill, K.A. Protein Surgery. *Protein Engineering* **1**: 369-372 (1987).

33. Dill, K.A. The Mechanism of Solute Retention in Reversed Phase Liquid Chromatography. *Journal of Physical Chemistry* **91**(7): 1980-1988 (1987).

32. Dill, K.A. The Stabilities of Globular Proteins. In *Protein Engineering*, p. 187. D.L. Oxender and C.F. Fox, eds. Alan R. Liss, Inc., 1987.

31. Stigter, D. and K.A. Dill. Interactions in Dilute Monolayers of Long Chain Ions at the Interface Between n-Heptane and Aqueous Salt Solutions. *Langmuir* **2**: 791 (1986).

30. Marqusee, J.A. and K.A. Dill. Chain Configurations in Lamellar Semicrystalline Polymer Interphases. *Macromolecules* **19**: 2420 (1986).

29. Marqusee, J.A. and K.A. Dill. Solute Partitioning into Chain Molecule Interphases: Monolayers, Bilayer Membranes and Micelles. *Journal of Chemical Physics* **85**: 434 (1986).

28. Cantor, R.S. and K.A. Dill. Theory for the Equation of State of Phospholipid Monolayers. *Langmuir* **2**: 331 (1986).

27. Dill, K.A. The Chain Conformations in Membranes and Micelles. *Advances in Colloid and Interface Science* **26**: 99 (1986).

26. Dill, K.A. and R.S. Cantor. The Statistical Thermodynamics of Surfactant Aggregates. In *Physics of Amphiphiles: Micelles, Vesicles, and Microemulsions*, p. 376. V. Degiorgio, and M. Corti, eds. Amsterdam: North Holland, 1986.

25. Cantor, R.S. and K.A. Dill. Statistical Thermodynamic Theory for the Melting of n-Alkanes from Their Rotator Phases. *Macromolecules* **18**: 1875 (1985).

24. Dill, K.A. Comment: Molecular Conformations in Surfactant Micelles. *Nature* **313**: 603 (1985).

23. Dill, K.A. Theory for the Folding and Stability of Globular Proteins. *Biochemistry* **24**: 1501 (1985).

22. Chen, S.H. and K.A. Dill. Comment: Water in the Hydrocarbon Core of Micelles. *Nature* **314**: 385 (1985).

21. Marqusee, J.A., M. Warner, and K.A. Dill. Frequency Dependence of NMR Relaxation Rates in Bilayer Membranes. *Journal of Chemical Physics* **81**: 6404 (1984).

20. Dill, K.A., D.E. Koppel, R.S. Cantor, J.D. Dill, D. Bendedouch, and S.H. Chen. The Molecular Conformations in Surfactant Micelles. *Nature* **309**: 42-45 (1984).

19. Cantor, R.S. and K.A. Dill. Statistical Thermodynamics of Short Chain Molecule Interphases II. Configurational Properties of Amphiphilic Aggregates. *Macromolecules* **17**: 384-388 (1984).

18. Dill, K.A. and R.S. Cantor. Statistical Thermodynamics of Short Chain Molecule Interphases I. Theory. *Macromolecules* **17**: 380-384 (1984).

17. Flory, P.J., D.Y. Yoon, and K.A. Dill. The Interphase in Lamellar Semicrystalline Polymers. *Macromolecules* **17**: 862-868 (1984).

16. Dill, K.A. Molecular Organization in Amphiphilic Aggregates. In *Surfactants in Solution*. K.L. Mittal, ed. New York: Plenum Press, 1983.
15. Dill, K.A. Configurations of Amphiphilic Molecules in Micelles. *Journal of Physical Chemistry* **86**: 1498 (1982).
14. Tullis, R.H., K.A. Dill, and P.A. Price. Fluorescence and Kinetic Studies on the Divalent Metal Ion Induced Conformational Changes in DNaseA. *Journal of Biological Chemistry* **256**: 5656 (1981).
13. Dill, K.A. and P.J. Flory. Molecular Organization in Micelles and Vesicles. *Proceedings of the National Academy of Sciences USA* **78**: 676 (1981).
12. Dill, K.A. and P.J. Flory. Interphases of Chain Molecules: Monolayers and Lipid Bilayer Membranes. *Proceedings of the National Academy of Sciences USA* **77**: 3115 (1980).
11. Dill, K.A. Concentration Dependence of the Viscosity and Viscoelasticity of Polymer Solutions: Application of the Theory of Muthukumar and Freed. *Macromolecules* **13**: 620 (1980).
10. Dill, K.A. Dynamics of Polymer Solutions: IV. Shear-Stress Relaxation Experiments on Solutions of DNA from Bacteriophage T2. *Macromolecules* **13**: 438 (1980).
9. Troll, M., K.A. Dill, and B.H. Zimm. Dynamics of Polymer Solutions: III. An Instrument for Stress Relaxations on Dilute Solutions of Large Polymer Molecules. *Macromolecules* **13**: 436 (1980).
8. Dill, K.A. and B.H. Zimm. Dynamics of Polymer Solutions: II. The Determination of Molecular Weight Distribution by Viscoelasticity. *Macromolecules* **13**: 432 (1980).
7. Dill, K.A. and B.H. Zimm. Dynamics of Polymer Solutions I. Theory for an Instrument. *Macromolecules* **13**: 426 (1980).
6. Dill, K.A. and B.H. Zimm. Stress- and Strain-Relaxation Dynamics of Polymer Solutions: I. Theory for an Instrument. *Macromolecules* **13**: 426 (1980).
5. Dill, K.A. Chain Folding in Semicrystalline Polymers. *Faraday Discussion, Chemical Society* **68**: 104, 452 (1979).
4. Dill, K.A., and B.H. Zimm. A Rheological Separator for Very Large DNA Molecules. *Nucleic Acids Research* **7**: 735 (1979).
3. Dill, K.A. Theory for the Separation of Very Large DNA Molecules by Radial Migration. *Biophysical Chemistry* **10**: 327 (1979).
2. Dill, K. A. and R.H. Shafer. Radial Migration of DNA Molecules in Cylindrical Flow. III. Circles and the Effect of Non- Gaussian Polymer Statistics. *Biophysical Chemistry* **4**: 51 (1976).
1. Dill, K. A. and M. Troll. FET Programs Op Amp for Invertible Gain. *Electronics* **113**, Jan. 9 (1975).