

Nathan Andrew Baker

Curriculum Vitae

PO Box 999, MSID K7-20
Richland, WA 99354
☎ +1 509 375 3997
✉ nathan.baker@pnnl.gov
📄 <https://goo.gl/UeODNL>

Nathan A. Baker, Ph.D. is a Laboratory Fellow in the Applied Statistics and Computational Modeling Group at Pacific Northwest National Laboratory (PNNL). His research focuses on the development of new algorithms and mathematical methods in biophysics, nanotechnology, and informatics. Current projects include new computational methods for modeling solvation in biomolecular systems (<http://www.poissonboltzmann.org/>), mathematical methods for mesoscale materials modeling (<http://www.pnnl.gov/computing/cm4/>), and development of new methods for signature discovery (<http://signatures.pnnl.gov>).

Experience

- 2015–present **Visiting Professor**, *Division of Applied Mathematics, Brown University*, Providence, RI.
- 2012–present **Laboratory Fellow**, *Computational and Statistical Analytics Division, Pacific Northwest National Laboratory*, Richland, WA.
- 2013–2015 **Technical Group Manager**, *Applied Statistics and Computational Modeling Group, Pacific Northwest National Laboratory*, Richland, WA.
- 2010–2012 **Chief Scientist**, *Computational and Statistical Analytics Division, Pacific Northwest National Laboratory*, Richland, WA.
- 2006–2010 **Associate Professor (tenured)**, *Department of Biochemistry and Molecular Biophysics, Washington University in St Louis*, St Louis, MO.
- 2002–2006 **Assistant Professor**, *Department of Biochemistry and Molecular Biophysics, Washington University in St Louis*, St Louis, MO.

Education

- 2001–2002 **Postdoctoral Researcher**, *Department of Chemistry and Biochemistry, University of California San Diego*, La Jolla, CA.
- 1997–2001 **PhD, Physical Chemistry**, *University of California San Diego*, La Jolla, CA.
- 1993–1997 **BS, Chemistry**, *University of Iowa*, Iowa City, IA, Honors and highest distinction.

Selected honors and awards

- 2012 **Fellow**, *American Association for the Advancement of Science*.
- 2010 **Cancer Biomedical Informatics Grid (caBIG) Connecting Collaborators Award**, *National Cancer Institute*.
- 2007 **Hewlett-Packard Junior Faculty Excellence Award**, *American Chemical Society*.

2004 **Research Fellowship**, *Alfred P Sloan Foundation*.

Service

Selected extramural service

- 2014–2017 **Associate Editor**, *Biophysical Journal*, Biophysical Society.
- 2015–2016 **External Advisory Board Member**, *Visual Analytics for sense-making in Criminal Intelligence analysis (VALCRI) Project*, European Commission.
- 2014–present **Editorial Board member**, *Scientific Data*, Nature Publishing Group.
- 2012–2016 **Member**, *Macromolecular Structure and Function D Study Section*, National Institutes of Health.
- 2015 **Advisory board member**, *Data Management and Analysis Advisory Board*, Novozymes.
- 2012–2014 **Co-Chair**, *Nanotechnology Databases and Ontologies, US-EU Communities of Research*, OSTP National Nanotechnology Coordinating Office.
- 2011–2014 **Editor-in-Chief**, *Computational Science and Discovery*, Institute of Physics.
- 2009–2014 **Chair**, *E56.01 Nanotechnology Subcommittee on Informatics and Terminology*, ASTM.
- 2013 **Reviewer**, *Progress in Research on Environmental, Health, and Safety Aspects of Engineered Nanomaterials*, National Academy of Sciences.
- 2010–2013 **Advisory committee member**, *Nanomaterial Registry*, RTI International & National Institutes of Health.
- 2009–2013 **Working group lead**, *Nanotechnology Working Group, caBIG Integrated Cancer Research Workspace*, National Cancer Institute.
- 2012 **Panelist**, *National Academies Board on Environmental Studies and Toxicology Research Progress on Environmental, Health, and Safety Aspects of Nanotechnology Workshop*, National Academies of Science.
- 2012 **Member**, *Program Committee, caBIG Annual Meeting*, National Cancer Institute.

Selected intramural service

- 2014–present **Chair**, *PNNL Institutional Computing Steering Committee*, Pacific Northwest National Laboratory.
- 2005–2010 **Director**, *Siteman Center for Cancer Nanotechnology Excellence Biocomputing Core*, Washington University in St Louis.
- 2007–2010 **Director**, *Computational and Molecular Biophysics graduate program*, Washington University in St Louis.

Selected funding

- 2012–2017 **PNNL lead**, *Collaboratory on Mathematics for Mesoscopic Modeling of Materials (FWP 63024)*, DOE ASCR.

- 2004–2017 **PI**, *APBS: Nanoscale biomolecular electrostatics software (R01 GM069702)*, NIH NIGMS.
- 2012–2016 **co-PI**, *DNA-DNA interactions with atomic detail (R01 GM099450)*, NIH NIGMS.
- 2011–2014 **co-I**, *Mechanism of oxysterol activation of membrane cholesterol (R01 HL067773)*, NIH NHLBI.
- 2012–2013 **PI**, *ISA-TAB curation of electrostatic data*, OpenEye Software.
- 2010–2013 **co-I**, *Characterization/bioinformatics-modeling of nanoparticle-complement interactions (U01 NS073457)*, NIH NINDS.
- 2009–2013 **co-PI**, *Collaborative research: Geometric flow approach to implicit solvation modeling (R01 GM090208-01)*, NIH NIGMS.
- 2004–2013 **co-I**, *National Biomedical Computation Resource (P41 RR0860516)*, NIH NCRR.
- 2008–2011 **DBP PI**, *Cancer Nanotechnology Knowledgebase for Nanoparticle Analysis and Design (U54 HG004028)*, NIH NHGRI.
- 2008–2011 **PI**, *caBIG Integrative Cancer Research Workspace (GS-35F-0306J)*, NIH NCI subcontract.
- 2007–2010 **PI**, *caNanoLab Data Submission Support (N01-CN-12400)*, NIH NCI subcontract.
- 2005–2010 **Core PI**, *The Siteman Cancer Center Nanotechnology Excellence at Washington University Biocomputing Core (U54 CA11934205)*.
- 2008–2010 **co-I**, *New Inhibitors of Acetylcholinesterase that Block Inactivation by Organophosphates (HDTRA1-08-C-0015)*, DoD DTRA.

Selected publications

- Dowling, Chase P., Sylvia Johnson, Elizabeth Jurrus, and Nathan A. Baker (2015). “An ISA-Tab specification for protein titration data exchange”. In: *arXiv*, p. 1511.06431. URL: <http://arxiv.org/abs/1511.06431>.
- Harper, Bryan, Dennis Thomas, Satish Chikkagoudar, Nathan Baker, Kaizhi Tang, Alejandro Heredia-Langner, Roberto Lins, and Stacey Harper (2015). “Comparative hazard analysis and toxicological modeling of diverse nanomaterials using the embryonic zebrafish (EZ) metric of toxicity”. In: *Journal of Nanoparticle Research* 17.6, pp. 1–12. DOI: 10.1007/s11051-015-3051-0. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC4454819/>.
- Hogan, Emilie, Kyle Monson, and Nathan A Baker (2015). “Energy Minimization of Discrete Protein Titration State Models Using Graph Theory”. In: *arXiv preprint arXiv:1507.07021*. URL: <http://arxiv.org/abs/1507.07021>.
- Lei, Huan, Xiu Yang, Bin Zheng, Guang Lin, and Nathan A Baker (2015). “Constructing Surrogate Models of Complex Systems with Enhanced Sparsity: Quantifying the Influence of Conformational Uncertainty in Biomolecular Solvation”. In: *Multiscale Modeling & Simulation* 13, pp. 1327–1353. DOI: 10.1137/140981587. URL: <http://arxiv.org/abs/1408.5629>.

- Pan, Wenxiao, Michael Daily, and Nathan A Baker (2015). "Numerical calculation of protein-ligand binding rates through solution of the Smoluchowski equation using smoothed particle hydrodynamics". In: *BMC Biophysics* 8.1, p. 7. DOI: 10.1186/s13628-015-0021-y. URL: <http://www.biomedcentral.com/2046-1682/8/7/abstract>.
- Sushko, Maria L, Dennis G Thomas, Suzette A Pabit, Lois Pollack, Alexey V Onufriev, and Nathan A Baker (2015). "The role of correlation and solvation in ion interactions with B-DNA". In: *arXiv preprint arXiv:1506.07951*. URL: <http://arxiv.org/abs/1506.07951>.
- Bielska, Agata A, Brett N Olsen, Sarah E Gale, Laurel Mydock-McGrane, Kathiresan Krishnan, Nathan Andrew Baker, Paul H Schlesinger, Douglas F Covey, and Daniel S Ory (2014). "Side-Chain Oxysterols Modulate Cholesterol Accessibility through Membrane Remodeling". In: *Biochemistry* 53.18, pp. 3042–3051. DOI: 10.1021/bi5000096. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC4020583/>.
- Daily, Michael D, Brett N Olsen, Paul H Schlesinger, Daniel S Ory, and Nathan A Baker (2014). "Improved coarse-grained modeling of cholesterol-containing lipid bilayers". In: *Journal of Chemical Theory and Computation* 10.5, pp. 2137–2150. DOI: 10.1021/ct401028g. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC4044725/>.
- Gosink, Luke J, Emilie A Hogan, Trenton C Pulsipher, and Nathan A Baker (2014). "Bayesian model aggregation for ensemble-based estimates of protein pKa values". In: *Proteins: Structure, Function, and Bioinformatics* 82.3, pp. 354–363. DOI: 10.1002/prot.24390. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3946329/>.
- Pham, Christine TN et al. (2014). "Application of a hemolysis assay for analysis of complement activation by perfluorocarbon nanoparticles". In: *Nanomedicine: Nanotechnology, Biology and Medicine* 10.3, pp. 651–660. DOI: 10.1016/j.nano.2013.10.012. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3966962/>.
- Richardson, C et al. (2014). *Materials Frontiers to Empower Quantum Computing*. Technical report. Los Alamos National Laboratory.
- Thomas, Dennis G, Satish Chikkagoudar, Alejandro Heredia-Langner, et al. (2014). "Physicochemical signatures of nanoparticle-dependent complement activation". In: *Computational Science & Discovery* 7.1, p. 015003. DOI: 10.1088/1749-4699/7/1/015003. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC4169987/>.
- Tolokh, Igor S, Suzette A Pabit, Andrea M Katz, Yujie Chen, Aleksander Drozdetski, Nathan Baker, Lois Pollack, and Alexey V Onufriev (2014). "Why double-stranded RNA resists condensation". In: *Nucleic Acids Research* 42.16, pp. 10823–31. DOI: 10.1093/nar/gku756. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC4176364/>.
- Wei, Guo Wei and Nathan A Baker (2014). "Differential geometry-based solvation and electrolyte transport models for biomolecular modeling: a review". In: *arXiv preprint arXiv:1412.0176*. URL: <http://arxiv.org/abs/1412.0176>.
- Baker, Nathan A, Juli D Klemm, Stacey L Harper, Sharon Gaheen, Mervi Heiskanen, Philippe Rocca-Serra, and Susanna-Assunta Sansone (2013). "Standardizing data". In: *Nature Nanotechnology*

- 8.2, pp. 73–74. DOI: 10.1038/nnano.2013.12. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC4054689/>.
- Baker, Nathan et al. (2013). “Research towards a systematic signature discovery process”. In: *Intelligence and Security Informatics (ISI), 2013 IEEE International Conference on*. IEEE, pp. 301–308. DOI: 10.1109/isi.2013.6578848. URL: <http://ieeexplore.ieee.org/lpdocs/epic03/wrapper.htm?arnumber=6578848>.
- Chakraborty, Sandeep, Basuthkar J Rao, Nathan Baker, and Bjarni Asgeirsson (2013). “Structural phylogeny by profile extraction and multiple superimposition using electrostatic congruence as a discriminator”. In: *Intrinsically Disordered Proteins* 1.1, e25463. DOI: 10.4161/idp.25463. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC4212511/>.
- Daily, Michael D, Jaehun Chun, Alejandro Heredia-Langner, Guowei Wei, and Nathan A Baker (2013). “Origin of parameter degeneracy and molecular shape relationships in geometric-flow calculations of solvation free energies”. In: *The Journal of Chemical Physics* 139.20, p. 204108. DOI: 10.1063/1.4832900. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3862591/>.
- Harper, Stacey L et al. (2013). “Nanoinformatics workshop report: current resources, community needs and the proposal of a collaborative framework for data sharing and information integration”. In: *Computational Science & Discovery* 6.1, p. 014008. DOI: 10.1088/1749-4699/6/1/014008. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3895330/>.
- Olsen, Brett N, Agata A Bielska, Tiffany Lee, Michael D Daily, Douglas F Covey, Paul H Schlesinger, Nathan A Baker, and Daniel S Ory (2013). “The structural basis of cholesterol accessibility in membranes”. In: *Biophysical Journal* 105.8, pp. 1838–1847. DOI: 10.1016/j.bpj.2013.08.042. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3797575/>.
- Thomas, Dennis G, Jaehun Chun, Zhan Chen, Guowei Wei, and Nathan A Baker (2013). “Parameterization of a geometric flow implicit solvation model”. In: *Journal of Computational Chemistry* 34.8, pp. 687–695. DOI: 10.1002/jcc.23181. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3578971/>.
- Thomas, Dennis G, Sharon Gaheen, et al. (2013). “ISA-TAB-Nano: a specification for sharing nano-material research data in spreadsheet-based format”. In: *BMC Biotechnology* 13.1, p. 2. DOI: 10.1186/1472-6750-13-2. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3598649/>.
- Chen, Zhan, Shan Zhao, Jaehun Chun, Dennis G Thomas, Nathan A Baker, Peter W Bates, and GW Wei (2012). “Variational approach for nonpolar solvation analysis”. In: *The Journal of Chemical Physics* 137.8, p. 084101. DOI: 10.1002/jcc.23181. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3578971/>.
- Jacob, Ferosh, Jeff Gray, Adam Wynne, Yan Liu, and Nathan Baker (2012). “Domain-specific languages for composing signature discovery workflows”. In: *Proceedings of the 2012 workshop on Domain-specific modeling*. ACM, pp. 61–64. DOI: 10.1145/2420918.2420934. URL: <http://dx.doi.org/10.1145/2420918.2420934>.
- Konecny, Robert, Nathan A Baker, and J Andrew McCammon (2012). “iAPBS: a programming interface to the adaptive Poisson–Boltzmann solver”. In: *Computational Science & Discovery* 5.1,

- p. 015005. DOI: 10.1088/1749-4699/5/1/015005. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3419494/>.
- Lee, Sun-Joo, Paul H Schlesinger, Samuel A Wickline, Gregory M Lanza, and Nathan A Baker (2012). "Simulation of fusion-mediated nanoemulsion interactions with model lipid bilayers". In: *Soft Matter* 8.26, pp. 7024–7035. DOI: 10.1039/C2SM25847A. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3375911/>.
- Maojo, Victor et al. (2012). "Nanoinformatics: developing new computing applications for nanomedicine". In: *Computing* 94.6, pp. 521–539. DOI: 10.1007/s00607-012-0191-2. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3430140/>.
- Olsen, Brett N, Paul H Schlesinger, Daniel S Ory, and Nathan A Baker (2012). "Side-chain oxysterols: from cells to membranes to molecules". In: *Biochimica et Biophysica Acta (BBA)-Biomembranes* 1818.2, pp. 330–336. DOI: 10.1016/j.bbamem.2011.06.014. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3197895/>.
- Ren, Pengyu, Jaehun Chun, Dennis G Thomas, Michael J Schnieders, Marcelo Marucho, Jiajing Zhang, and Nathan A Baker (2012). "Biomolecular electrostatics and solvation: a computational perspective". In: *Quarterly Reviews of Biophysics* 45.04, pp. 427–491. DOI: 10.1017/S003358351200011X. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3533255/>.
- Thomas, Dennis G, Satish Chikkagoudar, Alan R Chappell, Nathan Baker, et al. (2012). "Annotating the structure and components of a nanoparticle formulation using computable string expressions". In: *Bioinformatics and Biomedicine Workshops (BIBMW), 2012 IEEE International Conference on*. IEEE, pp. 889–894. DOI: 10.1109/BIBMW.2012.6470259. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3922075/>.
- Alexov, Emil et al. (2011). "Progress in the prediction of pKa values in proteins". In: *Proteins: structure, function, and bioinformatics* 79.12, pp. 3260–3275. DOI: 10.1002/prot.23189. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3243943/>.
- Carstensen, Tommy, Damien Farrell, Yong Huang, Nathan A Baker, and Jens Erik Nielsen (2011). "On the development of protein pKa calculation algorithms". In: *Proteins: Structure, Function, and Bioinformatics* 79.12, pp. 3287–3298. DOI: 10.1002/prot.23091. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3193850/>.
- Chen, Zhan, Nathan A Baker, and GW Wei (2011). "Differential geometry based solvation model II: Lagrangian formulation". In: *Journal of Mathematical Biology* 63.6, pp. 1139–1200. DOI: 10.1007/s00285-011-0402-z. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3113640/>.
- Hossain, KSM Tozammel, Chris Bailey-Kellogg, Alan M Friedman, Michael J Bradley, Nathan Baker, and Naren Ramakrishnan (2011). "Using physicochemical properties of amino acids to induce graphical models of residue couplings". In: *BIOKDD '11*, Article 3. DOI: 10.1145/2003351.2003354. URL: <http://dl.acm.org/citation.cfm?id=2003351.2003354>.
- Lee, Sun-Joo, Paul H Schlesinger, Samuel A Wickline, Gregory M Lanza, and Nathan A Baker (2011). "Interaction of melittin peptides with perfluorocarbon nanoemulsion particles". In: *The*

- Journal of Physical Chemistry B* 115.51, pp. 15271–15279. DOI: 10.1021/jp209543c. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3245340/>.
- Olsen, Brett N, Paul H Schlesinger, Daniel S Ory, and Nathan A Baker (2011). “25-Hydroxycholesterol increases the availability of cholesterol in phospholipid membranes”. In: *Biophysical Journal* 100.4, pp. 948–956. DOI: 10.1016/j.bpj.2010.12.3728. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3037553/>.
- Thomas, Dennis G, Fred Klaessig, et al. (2011). “Informatics and standards for nanomedicine technology”. In: *Wiley Interdisciplinary Reviews: Nanomedicine and Nanobiotechnology* 3, pp. 511–532. DOI: 10.1002/wnan.152. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3189420/>.
- Unni, Samir, Yong Huang, Robert M Hanson, Malcolm Tobias, Sriram Krishnan, Wilfred W Li, Jens E Nielsen, and Nathan A Baker (2011). “Web servers and services for electrostatics calculations with APBS and PDB2PQR”. In: *Journal of Computational Chemistry* 32.7, pp. 1488–1491. DOI: 10.1002/jcc.21720. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3062090/>.
- Callenberg, Keith M, Om P Choudhary, Gabriel L de Forest, David W Gohara, Nathan A Baker, and Michael Grabe (2010). “APBSmem: a graphical interface for electrostatic calculations at the membrane”. In: *PloS ONE* 5.9, e12722. DOI: 10.1371/journal.pone.0012722. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2947494/>.
- Chen, Zhan, Nathan A Baker, and Guo-Wei Wei (2010). “Differential geometry based solvation model I: Eulerian formulation”. In: *Journal of Computational Physics* 229.22, pp. 8231–8258. DOI: 10.1016/j.jcp.2010.06.036. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2951687/>.
- Lee, Sun-Joo, Brett Olsen, Paul H Schlesinger, and Nathan A Baker (2010). “Characterization of perfluorooctylbromide-based nanoemulsion particles using atomistic molecular dynamics simulations”. In: *The Journal of Physical Chemistry B* 114.31, pp. 10086–10096. DOI: 10.1021/jp103228c. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2917838/>.
- Thomas, Dennis G, Rohit V Pappu, and Nathan A Baker (2010). “NanoParticle Ontology for cancer nanotechnology research”. In: *Journal of Biomedical Informatics* 44.1, pp. 59–74. DOI: 10.1016/j.jbi.2010.03.001. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3042056/>.
- Chen, Alan A, Marcelo Marucho, Nathan A Baker, and Rohit V Pappu (2009). “Simulations of RNA interactions with monovalent ions”. In: *Methods in Enzymology* 469, pp. 411–432. DOI: 10.1016/S0076-6879(09)69020-0. URL: <http://www.ncbi.nlm.nih.gov/pubmed/20946801>.
- Olsen, Brett N, Paul H Schlesinger, and Nathan A Baker (2009). “Perturbations of membrane structure by cholesterol and cholesterol derivatives are determined by sterol orientation”. In: *Journal of the American Chemical Society* 131.13, pp. 4854–4865. DOI: 10.1021/ja8095224. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2675926/>.
- Silva, Jonathan R, Hua Pan, Dick Wu, Ali Nekouzadeh, Keith F Decker, Jianmin Cui, Nathan A Baker, David Sept, and Yoram Rudy (2009). “A multiscale model linking ion-channel molecular dynamics and electrostatics to the cardiac action potential”. In: *Proceedings of the National Academy of Sciences* 106.27, pp. 11102–11106. DOI: 10.1073/pnas.0904505106. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2700153/>.

- Thomas, Dennis G, Rohit V Pappu, Nathan Baker, et al. (2009). "Ontologies for cancer nanotechnology research". In: *Engineering in Medicine and Biology Society, 2009. EMBC 2009. Annual International Conference of the IEEE*. IEEE, pp. 4158–4161. DOI: 10.1109/IEMBS.2009.5333941. URL: <http://www.ncbi.nlm.nih.gov/pubmed/19964619>.
- Bradley, Michael J, Peter T Chivers, and Nathan A Baker (2008). "Molecular dynamics simulation of the Escherichia coli NikR protein: equilibrium conformational fluctuations reveal interdomain allosteric communication pathways". In: *Journal of Molecular Biology* 378.5, pp. 1155–1173. DOI: 10.1016/j.jmb.2008.03.010. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2478562/>.
- Dong, Feng, Brett Olsen, and Nathan A Baker (2008). "Computational methods for biomolecular electrostatics". In: *Methods in Cell Biology* 84, pp. 843–870. DOI: 10.1016/S0091-679X(07)84026-X. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2423940/>.
- Dong, Feng, Jason A Wagoner, and Nathan A Baker (2008). "Assessing the performance of implicit solvation models at a nucleic acid surface". In: *Physical Chemistry Chemical Physics* 10.32, pp. 4889–4902. DOI: 10.1039/b807384h. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2538626/>.
- Lee, Sun-Joo, Yuhua Song, and Nathan A Baker (2008). "Molecular dynamics simulations of asymmetric NaCl and KCl solutions separated by phosphatidylcholine bilayers: potential drops and structural changes induced by strong Na⁺-lipid interactions and finite size effects". In: *Biophysical Journal* 94.9, pp. 3565–3576. DOI: 10.1529/biophysj.107.116335. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2292386/>.
- Cerutti, David S, Nathan A Baker, and J Andrew McCammon (2007). "Solvent reaction field potential inside an uncharged globular protein: A bridge between implicit and explicit solvent models?" In: *The Journal of Chemical Physics* 127, p. 155101. DOI: 10.1063/1.2771171. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2556216/>.
- Cheng, Yuhui et al. (2007). "Finite element analysis of the time-dependent Smoluchowski equation for acetylcholinesterase reaction rate calculations". In: *Biophysical Journal* 92.10, pp. 3397–3406. DOI: 10.1529/biophysj.106.102533. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC1853150/>.
- Dolinsky, Todd J, Paul Czodrowski, Hui Li, Jens E Nielsen, Jan H Jensen, Gerhard Klebe, and Nathan A Baker (2007). "PDB2PQR: expanding and upgrading automated preparation of biomolecular structures for molecular simulations". In: *Nucleic Acids Research* 35.suppl 2, W522–W525. DOI: 10.1093/nar/gkm276. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC1933214/>.
- Schnieders, Michael J, Nathan A Baker, Pengyu Ren, and Jay W Ponder (2007). "Polarizable atomic multipole solutes in a Poisson-Boltzmann continuum". In: *The Journal of Chemical Physics* 126, p. 124114. DOI: 10.1063/1.2714528. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2430168/>.
- Swanson, Jessica MJ, Jason A Wagoner, Nathan A Baker, and J Andrew McCammon (2007). "Optimizing the Poisson dielectric boundary with explicit solvent forces and energies: lessons learned

- with atom-centered dielectric functions". In: *Journal of Chemical Theory and Computation* 3.1, pp. 170–183. DOI: 10.1021/ct600216k. URL: <http://dx.doi.org/10.1021/ct600216k>.
- Baker, Nathan A, Donald Bashford, and David A Case (2006). "Implicit solvent electrostatics in biomolecular simulation". In: *New Algorithms for Macromolecular Simulation*. Springer Berlin Heidelberg, pp. 263–295. DOI: 10.1007/3-540-31618-3. URL: <http://www.springerlink.com/index/10.1007/3-540-31618-3>.
- Konecny, Robert, Joanna Trylska, Florence Tama, Deqiang Zhang, Nathan A Baker, Charles L Brooks, and J Andrew McCammon (2006). "Electrostatic properties of cowpea chlorotic mottle virus and cucumber mosaic virus capsids". In: *Biopolymers* 82.2, pp. 106–120. DOI: 10.1002/bip.20409. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2440512/>.
- Wagoner, Jason A and Nathan A Baker (2006). "Assessing implicit models for nonpolar mean solvation forces: the importance of dispersion and volume terms". In: *Proceedings of the National Academy of Sciences* 103.22, pp. 8331–8336. DOI: 10.1073/pnas.0600118103. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC1482494/>.
- Wong, C Jason, Rachel L Rice, Nathan A Baker, Tao Ju, and Timothy M Lohman (2006). "Probing 3'-ssDNA Loop Formation in E. coli RecBCD/RecBC-DNA Complexes Using Non-natural DNA: A Model for 'Chi' Recognition Complexes". In: *Journal of Molecular Biology* 362.1, pp. 26–43. DOI: 10.1016/j.jmb.2006.07.016. URL: <http://www.sciencedirect.com/science/article/pii/S0022283606008795>.
- Zhang, Xiaoyu, Chandrajit L Bajaj, Bongjune Kwon, Todd J Dolinsky, Jens E Nielsen, and Nathan A Baker (2006). "Application of new multi-resolution methods for the comparison of biomolecular electrostatic properties in the absence of global structural similarity". In: *Multiscale Modeling & Simulation* 5.4, p. 1196. DOI: 10.1137/050647670. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2561295/>.
- Baker, Nathan A (2005a). "Biomolecular applications of Poisson-Boltzmann methods". In: *Reviews in Computational Chemistry* 21, p. 349. DOI: 10.1002/0471720895. URL: <http://doi.wiley.com/10.1002/0471720895>.
- (2005b). "Improving implicit solvent simulations: a Poisson-centric view". In: *Current Opinion in Structural Biology* 15.2, pp. 137–143. DOI: 10.1016/j.sbi.2005.02.001. URL: <http://www.sciencedirect.com/science/article/pii/S0959440X05000448>.
- Showalter, Scott A, Nathan A Baker, Changguo Tang, and Kathleen B Hall (2005). "Iron responsive element RNA flexibility described by NMR and isotropic reorientational eigenmode dynamics". In: *Journal of Biomolecular NMR* 32.3, pp. 179–193. DOI: 10.1007/s10858-005-7948-2. URL: <http://www.ncbi.nlm.nih.gov/pubmed/16132819>.
- Song, Yuhua, Victor Guallar, and Nathan A Baker (2005). "Molecular dynamics simulations of salicylate effects on the micro-and mesoscopic properties of a dipalmitoylphosphatidylcholine bilayer". In: *Biochemistry* 44.41, pp. 13425–13438. DOI: 10.1021/bi0506829. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2435121/>.

- Zhang, Deqiang, Jason Suen, et al. (2005). "Tetrameric mouse acetylcholinesterase: continuum diffusion rate calculations by solving the steady-state Smoluchowski equation using finite element methods". In: *Biophysical Journal* 88.3, pp. 1659–1665. DOI: 10.1529/biophysj.104.053850. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC1305222/>.
- Dolinsky, Todd J, PMJ Burgers, Kevin Karplus, and Nathan A Baker (2004). "SPrCY: comparison of structural predictions in the *Saccharomyces cerevisiae* genome". In: *Bioinformatics* 20.14, pp. 2312–2314. DOI: 10.1093/bioinformatics/bth223. URL: <http://bioinformatics.oxfordjournals.org/content/20/14/2312.short>.
- Dolinsky, Todd J, Jens E Nielsen, J Andrew McCammon, and Nathan A Baker (2004). "PDB2PQR: an automated pipeline for the setup of Poisson-Boltzmann electrostatics calculations". In: *Nucleic Acids Research* 32.suppl 2, W665–W667. DOI: 10.1093/nar/gkh381. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC441519/>.
- Song, Yuhua, Yongjie Zhang, Chandrajit L Bajaj, and Nathan A Baker (2004). "Continuum diffusion reaction rate calculations of wild-type and mutant mouse acetylcholinesterase: adaptive finite element analysis". In: *Biophysical Journal* 87.3, pp. 1558–1566. DOI: 10.1529/biophysj.104.041517. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC1304562/>.
- Song, Yuhua, Yongjie Zhang, Tongye Shen, Chandrajit L Bajaj, J Andrew McCammon, and Nathan A Baker (2004). "Finite element solution of the steady-state Smoluchowski equation for rate constant calculations". In: *Biophysical Journal* 86.4, pp. 2017–2029. DOI: 10.1016/S0006-3495(04)74263-0. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC1304055/>.
- Vitalis, Andreas, Nathan A Baker, and J Andrew McCammon (2004). "ISIM: a program for grand canonical Monte Carlo simulations of the ionic environment of biomolecules". In: *Molecular Simulation* 30.1, pp. 45–61. DOI: 10.1080/08927020310001597862. URL: <http://www.tandfonline.com/doi/abs/10.1080/08927020310001597862#.VKSAbyvF8T0>.
- Wagoner, Jason and Nathan A Baker (2004). "Solvation forces on biomolecular structures: a comparison of explicit solvent and Poisson-Boltzmann models". In: *Journal of Computational Chemistry* 25.13, pp. 1623–1629. DOI: 10.1002/jcc.20089. URL: <http://www.ncbi.nlm.nih.gov/pubmed/15264256>.
- Zhang, Deqiang, Robert Konecny, Nathan A Baker, and J Andrew McCammon (2004). "Electrostatic interaction between RNA and protein capsid in cowpea chlorotic mottle virus simulated by a coarse-grain RNA model and a Monte Carlo approach". In: *Biopolymers* 75.4, pp. 325–337. DOI: 10.1002/bip.20120. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2426774/>.
- Baker, Nathan A (2003). "Poisson-Boltzmann methods for biomolecular electrostatics". In: *Methods in Enzymology* 383, pp. 94–118. DOI: 10.1016/S0076-6879(04)83005-2. URL: <http://www.sciencedirect.com/science/article/pii/S0076687904830052>.
- Sept, David, Nathan A Baker, and J Andrew McCammon (2003). "The physical basis of microtubule structure and stability". In: *Protein Science* 12.10, pp. 2257–2261. DOI: 10.1110/ps.03187503. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2366909>.

- Tai, Kaihsu, Stephen D Bond, Hugh R MacMillan, Nathan Andrew Baker, Michael Jay Holst, and J Andrew McCammon (2003). "Finite element simulations of acetylcholine diffusion in neuromuscular junctions". In: *Biophysical Journal* 84.4, pp. 2234–2241. DOI: 10.1016/S0006-3495(03)75029-2. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC1302790/>.
- Baker, Nathan A, Kaihsu Tai, Richard Henchman, David Sept, Adrian Elcock, Michael Holst, and J Andrew McCammon (2002). "Mathematics and molecular neurobiology". In: *Computational Methods for Macromolecules: Challenges and Applications*. Springer Berlin Heidelberg, pp. 31–60. DOI: 10.1007/978-3-642-56080-4. URL: <http://www.springerlink.com/index/10.1007/978-3-642-56080-4>.
- Lin, Jung-Hsin, Nathan A Baker, and J Andrew McCammon (2002). "Bridging implicit and explicit solvent approaches for membrane electrostatics". In: *Biophysical Journal* 83.3, pp. 1374–1379. DOI: 10.1016/S0006-3495(02)73908-8. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC1302236/>.
- Ma, Chiansan, Nathan A Baker, Simpson Joseph, and J Andrew McCammon (2002). "Binding of aminoglycoside antibiotics to the small ribosomal subunit: a continuum electrostatics investigation". In: *Journal of the American Chemical Society* 124.7, pp. 1438–1442. DOI: 10.1021/ja016830+. URL: <http://pubs.acs.org/doi/abs/10.1021/ja016830+>.
- Baker, Nathan A, David Sept, Michael J Holst, and James Andrew McCammon (2001). "The adaptive multilevel finite element solution of the Poisson-Boltzmann equation on massively parallel computers". In: *IBM Journal of Research and Development* 45.3.4, pp. 427–438. DOI: 10.1147/rd.453.0427. URL: <http://ieeexplore.ieee.org/lpdocs/epic03/wrapper.htm?arnumber=5389050>.
- Baker, Nathan A, David Sept, Simpson Joseph, Michael J Holst, and J Andrew McCammon (2001). "Electrostatics of nanosystems: application to microtubules and the ribosome". In: *Proceedings of the National Academy of Sciences* 98.18, pp. 10037–10041. DOI: 10.1073/pnas.181342398. URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC56910/>.
- Baker, N, M Holst, and Feng Wang (2000). "Adaptive multilevel finite element solution of the Poisson-Boltzmann equation II. Refinement at solvent-accessible surfaces in biomolecular systems". In: *Journal of Computational Chemistry* 21.15, pp. 1343–1352. DOI: 10.1002/1096-987X(20001130)21:15<1343::AID-JCC2>3.0.CO;2-K. URL: [http://dx.doi.org/10.1002/1096-987X\(20001130\)21:15%3C1343::AID-JCC2%3E3.0.CO;2-K](http://dx.doi.org/10.1002/1096-987X(20001130)21:15%3C1343::AID-JCC2%3E3.0.CO;2-K).
- Holst, Michael, N Baker, and Feng Wang (2000). "Adaptive multilevel finite element solution of the Poisson-Boltzmann equation I. Algorithms and examples". In: *Journal of Computational Chemistry* 21.15, pp. 1319–1342. DOI: 10.1002/1096-987X(20001130)21:15<1319::AID-JCC1>3.0.CO;2-8. URL: [http://dx.doi.org/10.1002/1096-987X\(20001130\)21:15%3C1319::AID-JCC1%3E3.0.CO;2-8](http://dx.doi.org/10.1002/1096-987X(20001130)21:15%3C1319::AID-JCC1%3E3.0.CO;2-8).
- Quinn, Daniel M, Shawn R Feaster, Haridasan K Nair, Nathan A Baker, Zoran Radic, and Palmer Taylor (2000). "Delineation and decomposition of energies involved in quaternary ammonium binding in the active site of acetylcholinesterase". In: *Journal of the American Chemical Society* 122.13, pp. 2975–2980. DOI: 10.1021/ja9933588. URL: <http://dx.doi.org/10.1021/ja9933588>.

- Baker, Nathan A, Volkhard Helms, and J Andrew McCammon (1999). "Dynamical properties of fasciculin-2". In: *Proteins: Structure, Function, and Bioinformatics* 36.4, pp. 447–453. URL: <http://www.ncbi.nlm.nih.gov/pubmed/10450086>.
- Baker, Nathan A, Philippe H Hunenberger, and J Andrew McCammon (1999). "Polarization around an ion in a dielectric continuum with truncated electrostatic interactions". In: *The Journal of Chemical Physics* 110.22, pp. 10679–10692. DOI: 10.1063/1.479013. URL: <http://scitation.aip.org/content/aip/journal/jcp/110/22/10.1063/1.479013>.
- Baker, Nathan A and J Andrew McCammon (1999). "Non-Boltzmann rate distributions in stochastically gated reactions". In: *The Journal of Physical Chemistry B* 103.4, pp. 615–617. DOI: 10.1021/jp984151o. URL: <http://pubs.acs.org/doi/abs/10.1021/jp984151o>.
- Malany, Siobhan, Nathan Baker, Michelle Verweyst, Rohit Medhekar, Daniel M Quinn, Baruch Velan, Chanoch Kronman, and Avigdor Shafferman (1999). "Theoretical and experimental investigations of electrostatic effects on acetylcholinesterase catalysis and inhibition". In: *Chemico-Biological Interactions* 119, pp. 99–110. DOI: 10.1016/S0009-2797(99)00018-6. URL: <http://www.sciencedirect.com/science/article/pii/S0009279799000186>.
- Quinn, Daniel M, Rohit Medhekar, and N Baker (1999). "Ester hydrolysis". In: *Enzymes, Enzyme Mechanisms, Proteins, and Aspects of NO Chemistry*, pp. 101–137. URL: http://scholar.google.com/citations?view_op=view_citation&hl=en&user=L9dwKyUAAAAJ&pagesize=80&sortby=title&citation_for_view=L9dwKyUAAAAJ:Wp0gIr-vW9MC.
- Medhekar, Rohit, Nathan Baker, William Kearney, Kenneth Sando, and Daniel M. Quinn (1998). "Low-barrier hydrogen bond in the catalytic triad of serine esterases". In: *Structure and function of cholinesterases and related proteins*, p. 233. DOI: 10.1007/978-1-4899-1540-5. URL: <http://link.springer.com/10.1007/978-1-4899-1540-5>.
- Feaster, Shawn R, Keun Lee, Nathan Baker, David Y Hui, and Daniel M Quinn (1996). "Molecular recognition by cholesterol esterase of active site ligands: structure-reactivity effects for inhibition by aryl carbamates and subsequent carbamylenzyme turnover". In: *Biochemistry* 35.51, pp. 16723–16734. DOI: 10.1021/bi961677v. URL: <http://dx.doi.org/10.1021/bi961677v>.