

CURRICULUM VITAE

January 2025

Thomas E. Cheatham, III

<http://scholar.google.com/citations?user=fCEXKK4AAAAJ&hl=en>

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Education and Work Experience

- Professor** 7/14 to present
Department of Medicinal Chemistry, Director of Graduate Studies (FY 2014-2017)
College of Pharmacy, University of Utah
- Director, Center for High Performance Computing** 7/14 to present
University of Utah
- Associate Professor (with tenure) and Director of Graduate Studies (DOG)** 7/11 to 7/14
Department of Medicinal Chemistry
College of Pharmacy, University of Utah
- Associate Professor (with tenure)** 7/09 to 7/11
- Assistant Professor** 7/02 to 7/09
Department of Medicinal Chemistry (60%)
Department of Pharmaceutics and Pharmaceutical Chemistry (40%)
College of Pharmacy, University of Utah
- Adjunct Professor** 7/19 to 7/24
- Adjunct Associate Professor** 7/09 to 7/19
- Adjunct Assistant Professor** 7/00 to 7/09
Department of Biomedical Engineering (formerly Dept. of Bioengineering)
University of Utah
- Research Assistant Professor, Department of Medicinal Chemistry, U Utah** 1/00 to 7/02
- National Research Council Research Associate** 3/97 to 12/99
Laboratory of Biophysical Chemistry, NHLBI
National Institutes of Health
- Advisor: Bernard R. Brooks**
- PhD in Pharmaceutical Chemistry** 9/90 to 3/97
Department of Pharmaceutical Chemistry
University of California at San Francisco
- Advisor: Peter A. Kollman**
- Programmer/Analyst** 6/88 to 8/90
Aiken Computation Laboratory, Division of Applied Sciences
Harvard University
- Middlebury College, Middlebury Vermont** **March 1989**
- B.A. Chemistry (honors)**
Thesis: The Structure of Function of the Active Oxygen Intermediate of Cytochrome P450
- B.A. Mathematics and Computer Science**
Senior Project: The Fourier Transform and Applications in Digital Signal Processing
- Minor: Personality and Social Psychology**

Active Grants

- **NSF**
9/01/23-8/31/26

CICI: TCR: Prompt, Reliable, and Safe Security Update for Cyberinfrastructure
 PI: Xu, J; Co-PI: Cheatham

This OAC-2319880 award for \$1,198,133 is for development of software patching in HPC environments.
- **NSF / Internet2**
5/15/21-10/31/24

CI CoE: Demo Pilot: Advancing Research Computing and Data: Strategic Tools, Practices, and Professional Development
 PI: Brunson, Co-PIs: Cheatham, Yockel, Schmitz, and Mizumoto.

This ~\$1.5M NSF award for research computing and data professionalization, workforce development, and development of an RCD resource and career center.
- **NIH**
1/01/21-12/31/25

A Leukemia Cell-Specific Coiled-Coil Protein for Treatment of Chronic Myeloid Leukemia
 PI: Lim, CS.

This \$1,143,750 NIH award (direct costs) is to develop a protein inhibitor against Bcr-Abl, the causative agent of chronic myeloid leukemia (CML) and 30% of acute lymphoblastic leukemia (ALL), that circumvents current problems with Bcr-Abl targeted therapy. Effort: 0.5 calendar month/yr
- **NSF / University of Texas at Austin**
10/01/19-9/30/24

Portable applications driven approach to scalability on Frontera and future exascale systems
 PI: Berzins, sub-contract to NSF OAC-1818253 *Operations & maintenance for the endless frontier*, PI: Stanzone, Co-PIs: Panda, Minyard and Ghattas.

This is a sub-contract for Frontera support, ~\$250K over 5 years split by Berzins & Cheatham.
- **National Institutes of Health, R01-GM-081411**
9/23/19-8/30/24

Biomolecular simulation for the end-stage refinement of nucleic acid structure
 PI: Cheatham

Core NIH grant for assessment, validation and improvement of force fields for nucleic acids and experimental and computational ensembles of model RNA systems.
- **National Science Foundation, OAC-1919667**
10/01/19-9/30/24

MRI: Development of ACCORD, a Community Cyberinstrument for Broadening Access to Research on Sensitive Data
 PI: Ron Hutchins, Co-PIs: Sosonkina, Crawford, Midkiff, and Cheatham.

This \$2.5M equipment grant is for a compute infrastructure for U of Virginia for research on protected or sensitive data and Cheatham serves as a consultant.

Completed Grants (Utah)

- **National Science Foundation, OAC-1659425**
4/01/17-3/31/21

CC Cyber Team: Creating a Community of Regional Data and Workflow Cyberinfrastructure Facilitators*
 PI: Hauser, Co-PIs: Burns, Williams, Siegel, and Cheatham.

This is a collaborative ~\$1.5M grant for data facilitation in the region with Colorado State U and U Colorado Boulder as partners.

- **National Science Foundation, ACI-1443054** 10/01/14-9/30/19
CIF21 DIBBS: Middleware and high performance analytics libraries for scalable data science.
 PI: Fox, Co-PIs: Wang, Qiu, Jha, Marathe; Cheatham is significant personnel.
 This is a data infrastructure building blocks award for the development of SPIDAL (spidal.org) and middleware. In no-cost-extension.
- **National Science Foundation, ACI-1515572** 9/01/15-7/31/19
PRAC – Ensembles of molecular dynamics engines for assessing force fields, conformational change, and free energies of proteins and nucleic acids.
 PI: Cheatham, Co-PIs: Simmerling (Stony Brook U), Roitberg (UFI), and Case (Rutgers)
 This travel award allowed requests for allocations on Blue Waters and the team was awarded 12 million node hours on Blue Waters per year for three years. Currently in no cost extension.
- **National Institutes of Health, S10OD021644** 4/01/17-3/30/18
From genomics to natural language processing: A protected environment for research computing in the health sciences.
 PI: Cheatham.
 This is a major equipment grant to replacement of the protected environment for restricted data at Utah.
- **National Science Foundation, ACI-1341935** 3/01/14-8/31/18
Advanced Cyberinfrastructure – Research and Educational Facilitation: Campus-based computational research support.
 PI: Bottum (Clemson), Co-PIs: Jacobs (Hawaii), Wilson (Wisconsin), Cuff (Harvard), Dougherty (USC).
 This award supporting facilitators at six universities and advocated for the need for research computing and data facilitation on campuses; <http://aci-ref.org>. Cheatham PI of subcontract to U of Utah and PI Chair 2016-2018.
- **National Science Foundation, CNS-1338155** 8/30/13-6/30/17
MRI: Development of Apt, A Testbed Instrument with Adaptable Profiles for Network and Computational Science.
 PI: Ricci, Co-PIs: Facelli, Cheatham, Eide, van der Merwe
 This major research equipment grant is for the bare metal testbed hardware Apt. In no cost extension.
- **NSF Cyberinfrastructure Partnership / TeraGrid / XSEDE** 10/01/16-9/30/17
XRAC/LRAC MCA01S027: Insight into biomolecular structure, dynamics, interactions and energetics from simulation.
 PI: Cheatham, Computer time award: ~7M core hours awarded in 2017, award since 2002.
- **National Science Foundation, ACI-1521728** 4/01/15-3/31/17
RAPID: Optimizing experimental approaches to Ebola membrane fusion inhibitor peptide design through high-throughput biomolecular simulation workflows on Blue Waters. PI: Cheatham
 Computational protein design using simulation of Ebola membrane fusion inhibitor peptides.
- **National Science Foundation, ACI-1341034** 10/01/13-9/30/16
CC-NIE Integration: Science slices converting network research innovation into enhanced capability for computational science and engineering at the University of Utah.
 PI: Cheatham, Co-PIs: Bolton, van der Merwe, Ricci
 Development of a research computing and data science DMZ to bypass campus firewalls for large scale data transfers.

- **National Science Foundation, CHE-1266307** 10/01/13-9/30/16
CDS&E: Tools to facilitate deeper data analysis, exploration, management, and sharing of ensembles of molecular dynamics trajectory data.
 PI: Cheatham

Development of software and tools for annotating, analyzing, managing and sharing of ensembles of molecular dynamics simulation data.
- **National Institutes of Health, R01 GM098102** 9/30/11-8/31/16
RNA-ligand interactions: simulation and experiment.
 M-PIs: Cheatham, Kathleen Hall (Wash U, contact), Carlos Simmerling (Stony Brook).
 Combined experiment and theory to understand RNA and RNA-ligand interactions.
- **National Science Foundation, OCI-1440031** 9/01/14-8/31/16
PRAC – Hierarchical molecular dynamics sampling for assessing pathways and free energies of RNA catalysis, ligand binding, and conformational change.
 PI: Cheatham, Co-PIs: Simmerling (Stony Brook U), Roitberg (UFI), Case (Rutgers), and York (Rutgers)

This travel award allowed requests for allocations on Blue Waters and the team was awarded 7 million node hours on Blue Waters for half of 2014-2015.
- **National Science Foundation, OCI-1036208** 2/01/11-9/30/15
PRAC – Hierarchical molecular dynamics sampling for assessing pathways and free energies of RNA catalysis, ligand binding, and conformational change.
 PI: Cheatham, Co-PIs: Simmerling (Stony Brook U), Roitberg (UFI), and York (Rutgers)

This travel award allowed requests for allocations on Blue Waters and the team was awarded 14 million node hours on Blue Waters in 2013-2014.
- **National Institutes of Health, R01 GM074249** 1/01/11-12/31/14
P450-mediated dehydrogenation mechanisms.
 PI: Gary Yost, Co-PIs: Cheatham, Reilly
- **National Institutes of Health, R01 GM081411** 2/01/08-1/31/14
Biomolecular simulation for the end-stage refinement of nucleic acid structure.
 PI: Cheatham. No cost extension 2013-2014.
- **Pittsburgh Supercomputing Center, PSC12038P** 11/01/12-7/31/13
Converging simulations of a DNA duplex and explorations of a DNA minicircle on the microsecond timescale using MD on Anton
 PI: Cheatham, Computer time award: 100,000 hours on DE Shaw's Anton machine.
- **National Institutes of Health, R01 GM079383** 9/28/07-8/31/12
AMBER force field consortium: A coherent biomolecular simulation platform.
 PI: Yong Duan (UC Davis)
 Co-PIs: Cheatham, Carlos Simmerling (Stony Brook), Ray Luo (UC Irvine), Piotr Cieplak (Burnham Inst), Junmei Wang (Incisive, Inc.).
- **Pittsburgh Supercomputing Center, PSCA00033P** 4/01/11-12/31/11
Molecular dynamics of DNA and protein-DNA complexes: A proposal for obtaining microsecond trajectories using Anton.
 PI: Cheatham, Computer time award: ~50K node hours awarded in 2011 on DE Shaw's Anton machine at PSC supported by an NIH Go grant. This will support the ABC consortium.

- **Pittsburgh Supercomputing Center, PSCA00067P** 4/01/11-12/31/11
Development and testing of improved fixed-charge force fields for proteins
 PI: Case, Co-PIs: Cheatham, Simmerling, Merz. Computer time award: ~50K node hours awarded in 2011 on DE Shaw's Anton machine at PSC supported by an NIH Go grant.
- **DARPA, BAA-09-29** 4/01/10-3/31/11
Modeling and optimization of hemostatic peptide sequences for SAIC Wound Stasis System, sub-contract to Rapid hemostasis via a wound-targeting nanodelivery system.
 Sub-contract PI: Darrell Davis, Co-PI: Cheatham
- **Office of Naval Research, N00014-05-1-0457** 4/01/05-9/30/08
A new research tool for the computer simulation of chemical dynamics in complex systems.
 PI: Greg Voth (Chemistry); Co-PIs: Cheatham (10%), Dave Case (Scripps), Bill Miller (Berkeley) and Bernie Schlegel (Wayne State).
- **University of Utah, PID2502088.** 6/1/05-5/31/06
Funding Incentive Seed Grant: Integration of computer-aided drug-design tools with biomolecular simulation: Flexibility and scoring through iterative molecular dynamics and docking simulations on computational grids
 PI: Cheatham (\$30,500)
- **ACS-Petroleum Research Fund.** 1/1/05-10/1/05
Summer school on computation, simulation, and theory in chemistry, biological chemistry and materials chemistry
 PI: Jack Simons (Chemistry), Co-PIs: Cheatham, T. Seideman (Northwestern U), K. Jordon (U Pitt), E. Carter (UCLA), J.E. Shea (UCSB), J. Tully (Yale), E. Heller (Harvard), M. Head-Gordon (Berkeley), J. Doll (Brown), and B. Garrison (Penn State)
 Travel funds for students and teachers to Park City, Utah.
- **National Science Foundation, CHE-0326027** 9/01/03-8/31/07
ITR: Development of a Web-based Grid-computing Environment for Research and Education in Computational Science and Engineering
 PI: Thanh Truong (Chemistry), Co-PIs: Cheatham (10%), Julio Facelli (CHPC), James Lewis (WVU)
- **National Science Foundation, CHE-0218739.** 9/1/02-8/31/05
ITR: Multiscale simulation of Biomolecular Assemblies on a Computational Grid
 PI: Voth (Chemistry), Co-PI's: Cheatham, Ayton (Chemistry).
 < 1 mo. Salary/year for 1 year.
- **National Institutes of Health, 1 R21 EB002880** 9/23/03-8/31/05
Molecular docking and imaging devices for drug delivery
 PI: Y. Bruce Yu (Pharmaceutics), Collaborator: Cheatham, 5% effort.
- **National Institutes of Health, 1 S10 RR017214-01** 10/1/03-9/30/05
High End Instrumentation Program-- Proteins to Populations: A metacluster for bioinformatics
 PI: Facelli, Co-PIs: L. Canon-Albright (Medical Informatics), Cheatham, DM Grant (Chemistry), GA Voth (Chemistry), JA Weiss (Bioengineering), RB Weiss (Genetics).
 \$1,531,008 + \$500,000 UofU match; equipment only (no overhead).
- **University of Utah, College of Pharmacy** 7/1/01-6/30/02
Research Support Grant: Bioinformatics of Bio-molecular Simulation
 PI: Cheatham (~\$5,000)

Fellowships, Honors and Awards

- Gilda Loew Lectureship, International Society of Quantum Biology and Pharmacology President's Meeting, Innsbruck 2022
- University of Utah College of Pharmacy P2 Teacher of the year, 2013.
- Nominated for the College of Pharmacy, University of Utah, Teaching Award (2006-2012).
- Plenary speaker, eScience 2009, Oxford, U.K.
- Keynote speaker, TeraGrid 09, 2009
- Keynote speaker, Mercury conference at Hamilton College, 2009
- Chosen to present a Telluride Town Talk on the Science of Recreational Drugs, 2008
- Recipient of "Hewlett-Packard Outstanding Junior Faculty Award" from the American Chemical Society COMP division for outstanding work in computational chemistry, 2007.
- Member, NSF CyberInfrastructure User Advisory Committee, 2006-2007.
- Associate Editor, *Molecular Modeling and Computational Chemistry Results*, MMCC, Inc. (Editor: D. Busath, BYU), 2004-2005; Assistant Editor 2000-2004.
- National Research Council Research Associate, March 1997 to December 2000.
- National Science Foundation Travel Award, September 1997 (CECAM).
- Finalist, Computerworld-Smithsonian Awards in the category of Science, 1998.
- DNA simulation research was featured in the 1996 annual report and the 1998 PSC multimedia collection, see <http://www.psc.edu/science/Kollman/kollman.html>
- UCSF Chancellor's Graduate Research Fellow, September 1995 to June 1996.
- UCSF Pharmacy School, Frank Goyan Award for Excellence in Achievement in Phys. Chemistry, 1995.
- UCSF Dept. Pharmaceutical Chemistry Award for Outstanding Service, 1994-1995.

Professional Service / Review Panels

International Society of Quantum Biology and Pharmacology Councilor (2022-2026)
 ACCESS Researcher Advisory Committee (2023-)
 Delta Advisory Committee – NSF computational infrastructure at NCSA (2021-)
 Frontera at Texas Advanced Supercomputing Center Advisory Committee (2019-)
 Campus Research Computing Consortium (CaRCC), Chair (2016-), Council Chair (2017-2019), Acting Chair (2019-)
 Center for Computational Engineering and Sciences, UNICAMP, Brazil, Intl Advisory Board (2018-)
 International HPC Summer School, Organizing Committee & Presenter & Mentor (2012-, except 2023)
 NIH NIGMS MSFA study section (ad hoc) (October 2024)
 NSF Office of Advanced Cyberinfrastructure Review Panel (Fall 2023)
 National Academy of Sciences, Anton Molecular Dynamics Review Panel (September 2023)
 Utah Education Network Advisory Council (2015-2025)
 Natl. Academies Comm. on Proposal Evaluation for Allocation of Supercomputing Time on Anton (2013, 2014, 2023)
 NSF XSEDE User Advisory Committee (2015-2022)
 NSF Office of Advanced Cyberinfrastructure Review Panel (Fall 2022)
 NSF XSEDE Science Advisory Board (2016-2020)
 NSF XSEDE User Requirements Evaluation and Prioritization Committee (2016-2020)
 XSEDE Campus Champions Leadership Team (2020-2022)
 Internet2 E-CAS "Exploring Cloud for Acceleration of Science" Advisory Board (2018-2021)
 NIH NIGMS BCMB Member Conflict Panel (March 2021)
 NIH NIGMS BCMB-A (51) RM 18-099 NIH Transformational Research Awards Review (March 2020)
 Rocky Mountain Advanced Computing Consortium (RMACC), Vice-Chair (2017-2018), Board Member (2018-2020)
 Practice and Experience in Research Computing on Campuses (PEARC) Meeting Steering Committee (2016-2020)
 NSF XSEDE Science Advisory Board (2016-2020)
 NSF XSEDE User Requirements Evaluation and Prioritization Committee (2016-2020)
 Blue Waters Science and Engineering Team Advisory Committee – SETAC (2014-2019)
 NIH NIGMS MSFD Computational Biophysics Study Section (October 2019)

NSF Office of Advanced Cyberinfrastructure Review Panel (Spring 2019)
 NIH NIGMS BCMB Special Emphasis Panel (June 2018)
 Advanced Cyberinfrastructure Research and Educator Facilitator (ACI-REF), PI Chair (2016-2018)
 Advanced Research Computing on Campuses (ARCC) @ PEARC17, Chair (2017)
 Supercomputing 14/15/16 Education Program Executive Committee (2014-2016)
 Scientific Advisory Committee of the Swedish National Infrastructure for Computing (SNIC) (10/2013-10/2016)
 External Thesis Examiner and Opponent, KTH Royal Institute of Technology, Stockholm Sweden (4/2016)
 NSF XSEDE User Advisory Committee, Chair (2012-4/2015)
 NSF XSEDE Science Advisory Board (2012-4/2015)
 NSF XSEDE Senior Management Team (2012-4/2015)
 PRACE Supercomputing reviews (2015, 2013, 2012)
 NIH ZRG1 MSFD Study Section (charter member 7/11-6/15)
 XSEDE14 Science Track Chair (2013-2014)
 External PhD Thesis Examiner and Opponent, IRB Barcelona, Spain (10/14)
 NSF Review Panel, CHE (3/14)
 Great Lakes Consortium for Petascale Computation Blue Waters Allocations Review Panel (1/14)
 NIH ZRG1 MID-B NIAID Study Section (10/13)
 Referee for CECAM workshops (2013)
 NSF Review Panel OCI/CHE (11/12, 5/13)
 Supercomputing 12 Program Committee (2011-2012)
 Swiss Supercomputing Center proposal reviews (2011, 2012)
 NSF Site Visit Team, Engineering/CI (3/12)
 NIH ZRG F04 Fellowship: Chemistry, Biochemistry, Biophysics, and Bioengineering F30-F33 Study Section (3/12)
 National Institute of Computational Sciences User Advisory Committee (2010-2011)
 External PhD Thesis examiner and Opponent, University of Bergen, Norway (4/11)
 NIH NCI intramural lab site review (3/11)
 NSF Teragrid Science Advisory Board (6/07-7/11, Chair 7/10-7/11)
 NSF Review Panels (10/10, 5/10)
 University of Toronto, visit as external thesis examiner (9/10)
 NIH ZRG1 MSFD Study Section (Computational Biophysics) (ad hoc 2/09, 6/09, 6/10)
 AAAS review, University of Southern Florida neuroscience seed grants (2/10)
 NIH ZRG1 BCMB-P (58) RRFA-09-003: Challenge Grants Panel 7 mail reviews (6/09)
 NIH ZRG1 BCMB-B 90 (Special Topics in Biological Sciences) (ad hoc 7/08)
 TeraGrid Resource Allocations Committee/LRAC/MRAC, NSF Review Panels for computer time, (3/04-6/08, 4x/year), Chair 6/07, 9/07, 12/07, 3/08, 6/08.
 External PhD Thesis Examiner and Opponent, Karolinska Institute, Stockholm Sweden (2/08)
 External PhD Thesis Examiner and Opponent, Scuola Internazionale Superiore di Studi Avanzati, Trieste, Italy (1/08)
 NIH ZRG1 MSFD Study Section (Computational Biophysics) (ad hoc 10/07)
 NSF Cyberinfrastructure User Advisory Committee (6/06-6/07)
 NIH ZRG1 F04B-A (20) Study Section (Biophysics Fellowships) (ad hoc 3/06)
 NIH ZRG1 BCMB-Q (02) Study Section (Computational Biophysics) (ad hoc 10/05, 3/06)
 NSF Chemistry Division, mail review (10/05, 3/06, 8/06, 9/06, 8/07)
 Ad hoc reviewer for the University of Utah Funding Incentive Seed Grant, (10/04, 3/05)
 Universitat Innsbruck, Austria; External PhD thesis review (2004, 2005)
 NIH Special Emphasis Panel ZAI1 AR-M (M1), Cooperative Research for Development of Vaccines, Adjuvants, Therapeutics, Immunotherapeutics and Diagnostics for Biodefense and SARs (ad hoc 2/04)
 Cooperative Grants Program of the U.S. Civilian Res. Dev. Foundation, mail reviewer, (12/03)
 NIH BCCA Study Section, (ad hoc 2/03)
 Fonds zur Forderung der Wissenschaftlichen Forschung (Austria), mail reviewer, (3/02)
 Alliance Allocations Board/NRAC (NSF/Computer time) (3/02-9/04, 4x/year)
 DOE Site review team, Environmental Health Sciences Laboratory, PNNL (11/01)

Publications

- (1) DA Pearlman, DA Case, JW Caldwell, WS Ross, TE Cheatham, III, S DeBolt, DM Ferguson, GL Seibel & PA Kollman. "AMBER, a package of computer programs applying molecular mechanics, normal mode analysis, molecular dynamics and free energy calculations to simulate the structural and energetic properties of molecules." *Comp. Phys. Comm.* 91, 1-41 (1995).
- (2) TE Cheatham, III, JL Miller, T Fox, TA Darden & PA Kollman. "Molecular dynamics simulations on solvated biomolecular systems: The particle mesh Ewald method leads to stable trajectories of DNA, RNA and proteins." *J. Amer. Chem. Soc.* 117, 4193-4194 (1995).
- (3) TE Cheatham, III & PA Kollman. "Observation of the A-DNA to B-DNA transition during unrestrained molecular dynamics simulations in aqueous solution." *J. Mol. Biol.* 259, 434-444. (1996).
- (4) TE Cheatham, III & PA Kollman. "Molecular dynamics simulations highlight the structural differences among DNA:DNA, RNA:RNA and DNA:RNA hybrid duplexes." *J. Amer. Chem. Soc.* 119, 4805-4825 (1997).
- (5) P Cieplak, TE Cheatham, III & PA Kollman. "Molecular dynamics simulations find that 3' phosphoramidate modified DNA duplexes undergo a B to A transition and normal DNA duplexes an A to B transition." *J. Amer. Chem. Soc.* 119, 6722-6730 (1997).
- (6) T Spector, TE Cheatham, III & PA Kollman. "Unrestrained molecular dynamics of photodamaged DNA in aqueous solution." *J. Amer. Chem. Soc.* 119, 7095-7104 (1997).
- (7) TE Cheatham, III, MF Crowley, T Fox & PA Kollman. "A molecular level picture of the stabilization of A-DNA in mixed ethanol-water solutions." *Proc. Nat. Acad. Sci.* 94, 9626-9630 (1997).
- (8) TE Cheatham, III & PA Kollman. "Insight into the stabilization of A-DNA by specific ion association: Spontaneous B-DNA to A-DNA transitions observed in molecular dynamics simulations of d[ACCCGCGGGT]₂ in the presence of hexaamminecobalt(III)." *Structure* 5, 1297-1311 (1997).
- (9) MF Crowley, TA Darden, TE Cheatham, III & D Deerfield. "Adventures in improving the scaling and accuracy of a parallel molecular dynamics program." *J. Supercomputing.* 11, 255-278 (1997).
- (10) TA Darden, LG Pedersen, AY Toukmaji, MF Crowley & TE Cheatham, III. "Particle-mesh based methods for fast Ewald summation in molecular dynamics simulations". *Proceedings of the Eighth SIAM Conference on Parallel Processing for Scientific Computing*. Eds: M. Heath *et al.*, (Minn, MN). March (1997).
- (11) TE Cheatham, III, JL Miller, TI Spector, P Cieplak & PA Kollman. "Molecular dynamics simulations on nucleic acid systems using the Cornell *et al.* force field and particle mesh Ewald electrostatics." in *Modeling and Structure Determination of Nucleic Acids*. Eds: NB Leontis & J Santa Lucia, Jr (ACS Press: Washington, DC) p. 285-303 (1998).
- (12) SC Harvey, RK-Z Tan & TE Cheatham, III. "The flying ice cube: Velocity rescaling in molecular dynamics leads to violation of energy equipartition." *J. Comp. Chem.* 19, 726-740 (1998).
- (13) S Bogusz, TE Cheatham, III & BR Brooks. "Removal of pressure and free energy artifacts in charged periodic systems via net charged corrections to the Ewald potential." *J. Chem. Phys.* 108, 7070-7084 (1998).
- (14) TE Cheatham, III & PA Kollman. "Molecular dynamics simulation of nucleic acids in solution: How sensitive are the results to small perturbations in the force field and environment?" in *Structure, Motion, Interactions and Expression of Biological Macromolecules*. Eds: R.H. Sarma & M.H. Sarma (Adenine Press: Albany, NY) p. 99-116 (1998).
- (15) TE Cheatham, III & BR Brooks. "Recent advances in molecular dynamics simulation towards realistic representation of biomolecules in solution". *Theor. Chem. Acc.* 99, 279-288 (1998).

- (16) M Hodoscek, EM Billings, TE Cheatham, III & BR Brooks. "High performance computing in biophysics: Recent experiences and developments of CHARMM." *Proceedings of the International Symposium on Supercomputing: New Horizons of Computational Science*. (Kluwer Academic) (1998).
- (17) JL Miller, TE Cheatham, III, & PA Kollman. "Simulation of Nucleic Acid Structure." in *Oxford Handbook of Nucleic Acid Structure*. Ed: S. Neidle (Oxford University Press) p.95-115 (1999).
- (18) J Srinivasan, TE Cheatham, III, P Cieplak, PA Kollman & DA Case. "Continuum solvent studies of the stability of DNA, RNA and phosphoramidate-DNA helices". *J. Amer. Chem. Soc.* 120, 9401-9409 (1998).
- (19) PA Kollman, DA Pearlman, DA Case, JW Caldwell, WS Ross, TE Cheatham, III, S DeBolt, DM Ferguson & G Seibel. "AMBER." in *Encyclopedia of Computational Chemistry* (Wiley-Interscience: NY) (1998).
- (20) MF Crowley, TA Darden, TE Cheatham, III & D Deerfield. "Fine- and coarse-grain parallel AMBER and particle mesh Ewald on MPP's" in *Parallel Computing for Industrial and Scientific Applications*, Eds: J Jenness (Morgan-Kaufmann: NY) (1999).
- (21) TE Cheatham, III, J Srinivasan, DA Case & PA Kollman. "Molecular dynamics and continuum solvent studies of the stability of polyG-polyC and polyA-polyT DNA duplexes in solution." *J. Biomol. Struct. Dyn.* 16, 265-280 (1998).
- (22) TE Cheatham, III, BR Brooks & PA Kollman. "Molecular modeling of nucleic acid structure" in *Current Protocols in Nucleic Acid Chemistry*. (Wiley: New York) 7.5.1-7.5.13 (1999).
- (23) DE Konerding, TE Cheatham, III, PA Kollman & TL James. "Unrestrained and restrained PME MD on NMR-derived DNA decamer and dodecamer sequences." *J. Biomol. NMR.* 13, 119-131 (1999).
- (24) TE Cheatham, III, P Cieplak & PA Kollman. "A modified version of the Cornell *et al.* force field with improved sugar pucker phases and helical repeat." *J. Biomol. Struct. Dyn.* 16, 845-862 (1999).
- (25) TE Cheatham, III & PA Kollman. "Molecular dynamics simulations of nucleic acids". *Ann. Rev. Phys. Chem.* 51, 435-471 (2000).
- (26) PA Kollman, I Massova, C Reyes, B Kuhn, S Huo, L Chong, M Lee, T Lee, Y Duan, W Wang, O Donini, P Cieplak, J Srinivasan, DA Case & TE Cheatham, III. "Calculating structures and free energies of complex molecules: Combining molecular mechanics and continuum methods." *Acc. Chem. Res.* 33, 889-897 (2000).
- (27) TE Cheatham, III, BR Brooks & PA Kollman. "Molecular modeling of nucleic acid structure: Energy and sampling" in *Current Protocols in Nucleic Acid Chemistry*. (Wiley: New York) 7.8.1-7.8.21 (2001).
- (28) TE Cheatham, III, BR Brooks & PA Kollman. "Molecular modeling of nucleic acid structure: Electrostatics and solvation" in *Current Protocols in Nucleic Acid Chemistry*. (Wiley: New York) 7.9.1-7.9.22 (2001).
- (29) TE Cheatham, III, BR Brooks & PA Kollman. "Molecular modeling of nucleic acid structure: Setup and analysis" in *Current Protocols in Nucleic Acid Chemistry*. (Wiley: New York) 7.10.1-7.10.18 (2001).
- (30) TE Cheatham, III and MA Young. "Molecular dynamics simulations of nucleic acids: Successes, limitations and promise." *Biopolymers Nuc. Acid Sci.* 56, 232-256 (2001).
- (31) F Lankas, TE Cheatham, III, P Hobza, J Langowski, N Spackova, and J Sponer. "Critical effect of the N2 amino group on structure, dynamics and elasticity of DNA polypurine tracts" *Biophys. J.* 82, 2592-2609 (2002).
- (32) JP Lewis, J Pikus, TE Cheatham, III, EB Starikov, H Wang, J Tomfohr, and OF Sankey. "A comparison of electronic states in periodic and aperiodic poly(dA)-poly(dT) DNA." *Phys. Stat. Sol. (b)* 233, 90-100 (2002).

- (33) N Spackova, TE Cheatham, III, F Ryjacek, F Lankas, L van Meervelt, P Hobza, and J Sponer. "Molecular dynamics simulations and thermodynamics analysis of DNA-drug complexes. Minor groove binding between 4'-6-diamino-2-phenylindole and DNA duplexes in solution." *J. Amer. Chem. Soc.* 125, 1759-1769 (2003).
- (34) R Stefl, TE Cheatham, III, N Spackova, E Fadrna, I Berger, J Koca, and J Sponer. "Formation pathways of a guanine-quadruplex DNA revealed by molecular dynamics and thermodynamical analysis of the substates." *Biophys. J.* 85, 1787-1804 (2003). PMID: PMC1303352
- (35) JP Lewis, TE Cheatham, III, H Wang, E Starikow, and OF Sankey. "Dynamically amorphous character of electronic states in poly(dA)-poly(dT) DNA." *J. Phys. Chem. B* 107, 2581-2587 (2003).
- (36) F Lankas, J Sponer, J Langowski & TE Cheatham, III. "DNA base-pair step deformability inferred from molecular dynamics simulation" *Biophys. J.* 85, 2872-2883 (2003). PMID: PMC1303568
- (37) E Fadrna, N Spackova, R Stefl, J Koca, TE Cheatham, III, and J Sponer. "Molecular dynamics simulations of guanine quadruplex loops: Advances and force field limitations." *Biophys. J.* 87, 227-242 (2004). PMID: PMC1304345.
- (38) F Lankas, J Sponer, J Langowski, and TE Cheatham, III, "DNA deformability at the base pair level." *J. Amer. Chem. Soc.* 126, 4124-4125 (2004).
- (39) DL Beveridge, G Barreiro, KS Byun, DA Case, TE Cheatham III, SB Dixit, E Giudice, F Lankas, R Lavery, J Maddocks, R Osman, H Sklenar, G Stoll, KM Thayer, P Varnai, and MA Young "Molecular dynamics simulations of the 136 unique tetranucleotide sequences of DNA oligonucleotides. I. Research design, informatics, and results on d(CpG) steps." *Biophys. J.* 87, 3799-3813 (2004). PMID: PMC1304892
- (40) TE Cheatham, III "Simulation and modeling of nucleic acid structure, dynamics and interactions" *Curr. Opin. Struct. Biol.* 14, 360-367 (2004).
- (41) TE Cheatham, III "Molecular modeling and atomistic simulation of nucleic acids" in *Ann. Reports Comp. Chem.*, Ed: Spellmeyer, D., Vol. 1, pp. 75-89 (Elsevier) (2005).
- (42) DA Case, TE Cheatham, III, TA Darden, H Gohlke, R Luo, KM Merz, Jr., A Onufriev, C Simmerling, B Wang, and R Woods. "The AMBER biomolecular simulation programs" *J. Comp. Chem.* 26, 1668-1688 (2005). PMID: PMC1989667
- (43) SB Dixit, DL Beveridge, DA Case, TE Cheatham III, E Giudice, F Lankas, R Lavery, J Maddocks, R Osman, H Sklenar, KM Thayer, and P Varnai "Molecular dynamics simulations of the 136 unique tetranucleotide sequences of DNA oligonucleotides. II. Sequence context effects on the dynamical structures of the 10 unique dinucleotide steps." *Biophys. J.* 89, 3721-3740 (2005). PMID: PMC1366942.
- (44) TE Cheatham, III and DA Case, "Using AMBER to simulate DNA and RNA" in *Computational Studies of RNA and DNA* (Springer, The Netherlands, Editors: J. Sponer, F. Lankas) p 45-72 (2006)
- (45) N Spackova, TE Cheatham, III, and J Sponer, "Molecular dynamics simulations of nucleic acids" in *Computational Studies of RNA and RNA* (Springer, The Netherlands, Editors: J. Sponer, F. Lankas) p 301-326 (2006).
- (46) TN Truong, M Nayak, H Huynh, T Cook, P Marajan, LT Tran, J Bharath, S Jain, HB Pham, N Nguyen, Y Kim, S Choe, TE Cheatham, III, and J Facelli, "Computational Science and Engineering Online (CSE-Online): A Cyber-Infrastructure for scientific computing", *J. Chem. Info. Mod.* 46, 971-984 (2006).
- (47) F Paesani, W Zhang, DA Case, TE Cheatham, III, and GA Voth. "An accurate and simple quantum model for liquid water." *J. Chem. Phys.* 125, 184507 (2006).
- (48) A Perez, I Marchan, D Svozil, J Sponer, TE Cheatham, III, CA Loughton, M Orozco. "Refinement of the AMBER force field for nucleic acids. Improving the description of α/γ conformers". *Biophys. J.* 92, 3817-3829 (2007). PMID: PMC1868997.

- (49) A Wierzbicki, P Dalal, TE Cheatham, III, JE Knickelbein, ADJ Haymet, and JD Madura. "Antifreeze proteins at the ice/water interface: Three calculated discriminating properties for orientation of type I proteins." *Biophys. J.* 93, 1442-1451 (2007). PMC1948032.
- (50) J Shao, SW Tanner, N Thompson, TE Cheatham, III. "Clustering molecular dynamics trajectories: I. Characterizing the performance of different clustering algorithms." *J. Chem. Theory Comp.* 3, 2312-2334 (2007).
- (51) P Auffinger, TE Cheatham, III, and AC Vaiana. "Spontaneous formation of KCl aggregates in biomolecular simulations: a force field issue?" *J. Chem. Theory Comp.* 3, 1851-1859 (2007).
- (52) In Suk Joung, and TE Cheatham, III. "Determination of alkali and halide monovalent ion parameters for use in explicitly solvated biomolecular simulations" *J. Phys. Chem. B* 112, 9020-9041 (2008). PMC2652252.
- (53) D Svozil, JE Spomer, I Marchan, A Perez, TE Cheatham, III, J Luque, M Orozco, and J Spomer. "Geometrical and electronic structure variability of the sugar-phosphate backbone in nucleic acids." *J. Phys. Chem. B* 112, 8188-8197 (2008).
- (54) SS Pendley, YB Yu, and TE Cheatham, III. "Molecular dynamics guided study of salt bridge length dependence in both fluorinated and non-fluorinated parallel dimeric coiled-coil proteins." *Proteins* 74, 612-629 (2009) [DOI: 10.1002/prot.22177]. PMC2692595.
- (55) H Wang, TE Cheatham, III, PM Gannett, and J Lewis. "Differential electronic states observed during the A-B DNA duplex conformational transitions." *Soft Matter* 5, 685-690 (2009).
- (56) TS Han, M-M Zhang, A Walewska, P Gruszczynski, CR Robertson, TE Cheatham, III, D Yoshikami, B M Olivera, and G Bulaj. "Structurally-minimized μ -conotoxin analogs as sodium channel blockers: Implications for designing conopeptide-based therapeutics." *Chem. Med. Chem.* 4, 406-414 (2009). NIHMSID256909.
- (57) In Suk Joung, and TE Cheatham, III. "Molecular dynamics simulations of the dynamic and energetic properties of alkali and halide ions using water-model specific ion parameters." *J. Phys. Chem. B* 113, 13279-13290 (2009). PMC2755304.
- (58) E Fadrna, N Spackova, J Sarzynska, J Koca, M Orozco, TE Cheatham, III, T Kulinski, J Spomer. "Single stranded loops of quadruplex DNA as key benchmark for testing nucleic acids force field" *J. Chem. Theory Comp.* 5, 2514-2530 (2009).
- (59) In Suk Joung, Ö Persil Çetinkol, NV Hud, and TE Cheatham, III. "Molecular dynamics simulations and coupled nucleotide substitution experiments indicate the nature of A•A base pairing and a putative structure of the coralyne-induced homo-adenine duplex." *Nuc. Acids Res.* 37, 7715-7727 (2009). PMC2794157.
- (60) T Murai, CR Orton, K Shahrokh, TE Cheatham, III, RM Ward, and GS Yost. "CYP3A5-specific mechanism-based inactivation by fluticasone propionate, an inhaled glucocorticoid." *Processings of the 16th international conference on cytochrome P450*. (Medimond s.r.l., Bologna Italy) ISBN 978-88-7587-524-4, pg. 15-18 (2009).
- (61) R Lavery, K Zakrzewska, DL Beveridge, TC Bishop, DA Case, TE Cheatham, III, S Dixit, B Jayaram, F Lankas, C Laughton, JH Maddocks, A Michon, R Osman, M Orozco, A Perez, T Singh, N Spackova, and J Spomer. "A systematic molecular dynamics study of nearest-neighbor effects on base pair and base pair step conformations and fluctuations in B-DNA". *Nuc. Acids Res.* 38, 299-313 (2010). PMC2800215.
- (62) T Truong, H Freedman, L Le, TE Cheatham, III, J Tuszynski, and L Huynh. "Explicitly-solvated ligand contribution to continuum solvation models for binding free energies: Selectivity of theophylline binding to an RNA aptamer." *J. Phys. Chem. B* 114, 2227-2237 (2010). NIHMSID173947.

- (63) RB Paulsen, PP Seth, EE Swayze, RH Griffey, JJ Skalicky, TE Cheatham, III, and DR Davis. "Inhibitor induced structure change in the HCV IRES domain IIa RNA." *Proc. Natl. Acad. Sci.* 107, 7263-7268 (2010). PMC2867761.
- (64) CD Moore, K Shahrokh, SF Sontum, TE Cheatham, III, and GS Yost. "Improved Cyp3A4 molecular models accurately predict Phe215 requirement for raloxifene dehydrogenation selectivity." *Biochemistry* 49, 9011-9019 (2010). PMC2958526.
- (65) KF Wong, JL Sonnerberg, F Paesani, T Yamamoto, J Vanicek, J Zhang, HB Schlegl, DA Case, TE Cheatham, III, WH Miller, and GA Voth. "Proton transfer studied using a combined ab initio reactive potential energy surface with quantum path integral methodology." *J. Chem. Theory Comp.* 6, 2566-2580 (2010). PMC2992356.
- (66) P Banas, D Hollas, M Zgarbova, P Jurecka, M Orozco, TE Cheatham, III, J Sponer, and M Otyepka. "Performance of molecular mechanics force fields for RNA simulations. Stability of UUCG and GNRA hairpins." *J. Chem. Theory Comp.* 6, 3836-3849 (2010). NIHMS345258.
- (67) R DeMille, TE Cheatham, III, and V Molinero. "A coarse-grained model of DNA with explicit solvation by water and ions." *J. Phys. Chem. B* 115, 132-142 (2011). PMC3019136.
- (68) X Cang, J Sponer, and TE Cheatham, III. "Explaining the varied glycosidic conformational, G-tract length and sequence preferences for anti-parallel G-quadruplexes." *Nuc. Acids Res.* 39, 4499-4512 (2011). PMC3105399.
- (69) KC Thomas, M Ethirajan, K Shahrokh, H Sun, J Lee, TE Cheatham, III, GS Yost, and CA Reilly. "Structure activity relationship of capsaicin analogues and TRPV₁-mediated human lung epithelial cell toxicity." *J. Pharm. Exp. Ther.* 337, 400-410 (2011). PMC3083109.
- (70) X Wei, NM Henriksen, JJ Skalicky, MK Harper, TE Cheatham, III, CM Ireland, and RM Van Wagoner. "Araiosamines A-D: Tris-bromoindeole cyclic guanidine alkaloids from the marine sponge *Clathria* (*Thalysias*) *araiosa*." *J. Org. Chem.* 76, 5515-5523 (2011). PMC3188435.
- (71) AS Dixon, SS Pendley, BJ Bruno, DW Woessner, AA Shimpi, TE Cheatham, III, and CS Lim. "Disruption of Bcr-Abl coiled coil oligomerization by design." *J. Biol. Chem.* 286, 27751-27760 (2011). PMC3149365.
- (72) M Zgarbova, M Otyepka, J Sponer, A Mladek, P Banas, TE Cheatham, III, and P Jurecka. "Refinement of the Cornell *et al.* nucleic acid force field based on reference quantum chemical calculations of torsion profiles of the glycosidic torsion." *J. Chem. Theory Comp.* 7, 2886-2902 (2011). DOI: 10.1021/ct200162x. PMC3171997.
- (73) X Cang, J Sponer, and TE Cheatham, III. "Insight into G-DNA structural polymorphism and folding from sequence and loop connectivity through free energy analysis." *J. Amer. Chem. Soc.* 133, 14270-14279 (2011). DOI: 10.1021/ja107805r. PMC3168932.
- (74) K. Shahrokh, A Orendt, GS Yost, and TE Cheatham, III. "Quantum mechanically derived AMBER-compatible heme parameters for various states of the cytochrome P450 catalytic cycle." *J. Comp. Chem.* 33, 119-133 (2012). PMC3242737.
- (75) AS Dixon, GD Miller, BJ Bruno, JE Constance, DW Woessner, TP Fidler, JC Robertson, TE Cheatham, III, and CS Lim. "Improved coiled-coil design enhances interaction with Bcr-Abl and induces apoptosis." *Mol. Pharm.* 9, 187-195 (2012). PMC3083109.
- (76) Z Lin, M Flores, I Forteza, NM Henriksen, GP Concepcion, G Rosenberg, MG Haygood, BM Olivera, AR Light, TE Cheatham, III and EW Schmidt. "Totopotensamides, polyketide-cyclic peptide hybrids from a mollusk-associated bacterium *Streptomyces* sp." *J. Nat. Prod.* 75, 644-649 (2012). PMC3338888.

- (77) K Shahrokh, TE Cheatham, III, and GS Yost. "Conformational dynamics of CYP3A4 demonstrate the important role of Arg212 coupled with the opening of ingress, egress and solvent channels to dehydrogenation of 4-hydroxy-tamoxifen". *Biochem. Biophys. Acta.* 1820, 1605-1617 (2012). PMC3404218.
- (78) DL Beveridge, TE Cheatham, III, and M Mezei. "The ABCs of molecular dynamics simulations on B-DNA, circa 2012". *J. Bioscience* 37, 379-397 (2012).
- (79) J Sponer, X Cang, and TE Cheatham, III. "Molecular dynamics simulations of G-DNA and perspectives on the simulation of nucleic acid structures." *Methods* 57, 25-39 (2012). PMC3775459.
- (80) M Krepl, M Zgarbova, P. Stadlbauer, M Otyepka, P Banas, J Koca, TE Cheatham, III, P Jurecka, and J Sponer. "Reference simulations of noncanonical nucleic acids with different X variants of the AMBER force field: Quadruplex DNA, quadruplex RNA, and Z-DNA. *J. Chem. Theory Comp.* 8, 2506-2520 (2012). PMC3506181.
- (81) NM Henriksen, DR Davis, and TE Cheatham, III. "Molecular dynamics re-refinement of two different small RNA loop structures using the original NMR data suggest a common structure." *J. Biomol. NMR* 53, 321-339 (2012). PMC3405240.
- (82) JC Thibault, JC Facelli, and TE Cheatham, III. "iBIOMES: Managing and sharing biomolecular simulation data in a distributed environment." *J. Chem. Inf. Model.* 53, 725-736 (2013).
- (83) NM Henriksen, DR Roe, and TE Cheatham, III. "Reliable oligonucleotide conformational ensemble generation in explicit solvent for force field assessment using reservoir replica exchange molecular dynamics simulation" *J. Phys. Chem. B* 117, 4014-4027 (2013). PMC3775460.
- (84) M Zgarbova, F Javier Luque, J Sponer, TE Cheatham, III, M Otyepka, and P Jurecka, "Toward improved description of DNA backbone: Revisiting epsilon and zeta torsion force field parameters." *J. Chem. Theory Comp.* 9, 2339-2354 (2013). PMC3775469.
- (85) P Stadlbauer, M Krepl, TE Cheatham, III, J Koca and J Sponer. "Structural dynamics of possible late-stage intermediates in folding quadruplex DNA studied by molecular simulations." *Nuc. Acids Res.* 41, 7128-7143 (2013). PMC3737530.
- (86) J Sponer, A Mladek, N Spackova, X Cang, TE Cheatham, III and S Grimme. "Relative stability of different DNA guanine quadruplex stem topologies derived using large-scale quantum-chemical computations." *J. Amer. Chem. Soc.* 135, 9785-9796 (2013). PMC3775466.
- (87) DR Roe and TE Cheatham, III. "PTRAJ and CPPTRAJ: Software for processing and analysis of molecular dynamics trajectory data." *J. Chem. Theory Comp.* 9, 3084-3095 (2013).
- (88) TE Cheatham, III and DA Case. "Twenty-five years of nucleic acid simulations." *Biopolymers* 99, 969-977 (2013).
- (89) R Galindo-Murillo, C Bergonzo, and TE Cheatham, III. "Molecular modeling of nucleic acid structure." *Current Protocols Nucleic Acid Chemistry* 54: 7.5.1-7.5.13 (2013).
- (90) C Bergonzo, R Galindo-Murillo, and TE Cheatham, III. "Molecular modeling of nucleic acid structure: Energy and Sampling." *Current Protocols Nucleic Acid Chemistry* 54: 7.8.1-7.8.21 (2013).
- (91) C Bergonzo, R Galindo-Murillo, and TE Cheatham, III. "Molecular modeling of nucleic acid structure: Electrostatics and solvation" *Current Protocols Nucleic Acid Chemistry* 54: 7.9.1-7.9.27 (2013).
- (92) NN Ashton, DR Roe, RB Weiss, TE Cheatham, III, and RJ Stewart. "Self-tensioning aquatic caddisfly silk: Ca²⁺-dependent structure, strength, and load cycle hysteresis." *Biomacromolecules* 14, 3668-3681 (2013).

- (93) C Bergonzo, N Henriksen, DR Roe, J Swails, AE Roitberg, and TE Cheatham, III. "Multi-dimensional replica exchange molecular dynamics yields a converged ensemble of an RNA tetranucleotide." *J. Chem. Theory Comp.* 10, 492-499 (2014).
- (94) R Galindo-Murillo, C Bergonzo, and TE Cheatham, III. "Molecular modeling of nucleic acid structure: Setup and analysis." *Current Protocols Nucleic Acid Chemistry* 56: 7.10.1-7.10.21 (2014).
- (95) JC Thibault, DR Roe, JC Facelli, and TE Cheatham, III. "Data model, dictionaries, and desiderata for biomolecular simulation data indexing and sharing." *J. Cheminformatics* 6 (2014). doi:10.1186/1758-2946-6-4
- (96) DR Roe, C Bergonzo, and TE Cheatham, III. "Evaluation of enhanced sampling provided by accelerated molecular dynamics with Hamiltonian replica exchange methods." *J. Phys. Chem. B* 118, 3542-3552 (2014).
- (97) R Galindo-Murillo and TE Cheatham, III. "DNA-binding dynamics and energetics of Co, Ni, and Cu metallopeptides." *Chem. Med. Chem.* 9, 1252-1259 (2014).
- (98) J Thibault, TE Cheatham, III, and JC Facelli. "iBIOMES Lite: Summarizing biomolecular simulation data in limited settings." *J. Chem. Info Model.* 54, 1810-1819 (2014).
- (99) NM Henriksen, Hamed Hayatshahi, DR Davis, and TE Cheatham, III. "Structural and energetic analysis of 2-aminobenzimidazole inhibitors in complex with the hepatitis C virus IRES RNA using molecular dynamics simulations." *J. Chem. Info Model.* 54, 1758-1772 (2014).
- (100) A Okal, S Cornillie, SJ Matissek, KJ Matissek, TE Cheatham, III and CS Lim. "A re-engineered p53 chimera with enhanced homo-oligomerization that maintains tumor suppressor activity." *Mol. Pharm.* 11, 2442-2452 (2014).
- (101) P Stadlbauer, L Trantírek, TE Cheatham, J Koca and J Sponer. "Triplex intermediates in folding of human telomeric quadruplexes probed by microsecond-scale molecular dynamics simulations." *Biochimie* 105: 22-35 (2014) doi:10.1016/j.biochi.2014.07.009.
- (102) Z Lin, M Koch, M Abdel Aziz, D Tianero, R Galindo, TE Cheatham, III, L Barrows, C Reilly, and E Schmidt. "Oxazin A, A Pseudodimeric Natural Product of Mixed Bio-synthetic Origin from a Filamentous Fungus." *Organic Letters* 16, 4774-4777 (2014) doi: 10.1021/ol502227x.
- (103) R Galindo-Murillo, DR Roe, and TE Cheatham, III. "On the absence of intrahelical DNA dynamics on the μ s to ms timescale." *Nature Commun.* 5:5152 (2014) doi: 10.1038/ncomms6152.
- (104) M Pasi, JH Maddocks, D Beveridge, TC Bishop, DA Case, TE Cheatham, III, PD Dans, B Jayaram, F Lankas, C Laughton, J Mitchell, R Osman, M Orozco, A Perez, D Petkeviciute, N Spackova, J Sponer, K Zakrzewska, and R Lavery. " μ ABC: A systematic microsecond molecular dynamics study of tetranucleotide sequence effects in B-DNA." *Nuc. Acids Res.* 42, 12272-12283 (2014).
- (105) R Galindo-Murillo, DR Roe, and TE Cheatham, III. "Convergence and reproducibility in molecular dynamics simulations of the DNA duplex d(GCACGAACGAACGAACGC)." *Biochimica Biophys. Acta* 1850, 1041-1058 (2015) doi: 10.1016/j.bbagen.2014.09.007.
- (106) TE Cheatham, III and DR Roe. "The impact of heterogeneous computing on workflows for biomolecular simulation and analysis." *Computing in Science and Engineering* 17:2, 30-39 (2015).
- (107) R Galindo-Murillo, JC Garcia-Ramos, L Ruiz-Azuara, TE Cheatham, III, and F Cortes-Guzman. "Intercalation processes of copper complexes in DNA." *Nuc. Acids Res.* 43, 5364-5376 (2015).
- (108) C Bergonzo, N Henriksen, DR Roe, and TE Cheatham, III. "Highly sampled tetranucleotide and tetraloop motifs enable evaluation of common RNA force fields." *RNA* 29, 1578-1590 (2015).
- (109) AC Simmonett, FC Pickard IV, Y Shao, TE Cheatham, III and BR Brooks. "Efficient treatment of induced dipoles." *J. Chem. Phys.* 143, 074115 (2015).

- (110) C Bergonzo and TE Cheatham, III. "Improved force field parameters lead to a better description of RNA structure" *J. Chem. Theory Comp.* 11, 3969-3972 (2015).
- (111) C Bergonzo, KB Hall, and TE Cheatham, III. "Stem-loop V of Varkud satellite RNA exhibits characteristics of the Mg^{2+} bound structure in the presence of monovalent ions." *J. Phys. Chem. B* 119, 12355-12364 (2015). PMC4634716
- (112) R Galindo-Murillo and TE Cheatham, III. "Using information about DNA structure and dynamics from experiment and simulation to give insight into genome-wide association studies." Chapter 4 in *Translation Cardiometabolic Genomic Medicine*, edited by A Rodriguez-Oquendo, p 81-98 (2015).
- (113) JC Thibault, DR Roe, K Eilbeck, TE Cheatham, III, and JC Facelli. "Development of an informatics infrastructure for data exchange of biomolecular simulations: Architecture, data models, and ontology." *SAR and QSAR in Environmental Research* DOI: 10.1080/1062936X.2015.1076515 (2015).
- (114) JC Robertson and TE Cheatham, III. "DNA backbone BI/BII distribution and dynamics in E2 protein-bound environment determined by molecular dynamics simulation." *J. Phys. Chem. B* 119, 14111-14119 (2015).
- (115) M Zgarbova, J Sponer, M Otyepka, TE Cheatham, III, R Galindo-Murillo, and P Jurecka. "Refinement of the sugar-phosphate backbone torsion beta for the AMBER force fields improves the description of Z-DNA and B-DNA." *J. Chem. Theory Comp.* 11, 5723-5736 (2015).
- (116) R Galindo-Murillo, DR Davis, and TE Cheatham, III. "Probing the influence of hypermodified residues within the tRNA₃^{Lys} anticodon stem loop interacting with the A-loop primer sequence from HIV-1." *Biochimica Biophys. Acta* 1860, 607-617 (2016).
- (117) JT Waters, X-J Lu, R Galindo-Murillo, JC Gumbart, HD Kim, TE Cheatham, III and SC Harvey. "Transitions of double-stranded DNA between the A- and B- forms." *J. Phys. Chem. B* 120, 8449-8456 (2016). DOI: 10.1021/acs.jpcc.6b02155
- (118) C Bergonzo, KB Hall, and TE Cheatham, III. "Divalent Ion Dependent Conformational Changes in an RNA Stem-Loop Observed by Molecular Dynamics." *J. Chem. Theory Comp.* 12, 3382-3389 (2016).
- (119) R Galindo-Murillo, JC Robertson, M Zgarbova, J Šponer, M Otyepka, P Jurečka, and TE Cheatham III. "Assessing the current state of Amber force field modifications for DNA." *J. Chem. Theory Comp.* 12, 4114-4127 (2016). DOI: 10.1021/acs.jctc.6b00186
- (120) Z Heidari, DR Roe, R Galindo-Murillo, JB Ghasemi, and TE Cheatham III. "Using Wavelet Analysis To Assist in Identification of Significant Events in Molecular Dynamics Simulations." *J. Chem. Inf. Model.* 56, 1282-1291 (2016).
- (121) Y Hao, E Pierce, D Roe, M Morita, JA McIntosh, V Agarwal, TE Cheatham III, EW Schmidt, and SK Nair. "Molecular basis of broad-substrate selectivity of a peptide prenyltransferase." *PNAS* 113, 14037-14042 (2016).
- (122) HS Hayatshahi, DR Roe, R Galindo-Murillo, KB Hall, and TE Cheatham, III. "Computational Assessment of Potassium and Magnesium Ion Binding to a Buried Pocket in the GTPase-Associating Center RNA." *J. Phys. Chem. B* 121, 451-462 (2017).
- (123) M Zgarbova, P Jurecka, F Lankas, Filip; TE Cheatham, III, J Sponer, M Otyepka. "Influence of BII Backbone Substates on DNA Twist: A Unified View and Comparison of Simulation and Experiment for all 136 Distinct Tetranucleotide Sequences." *J. Chem. Info. Model.* 57, 275-287 (2017).
- (124) Y Wang, BJ Bruno, S Cornillie, JM Nogueira, D Chen, TE Cheatham, III, CS Lim, and D Chou. "Application of thiol-yne/thiol-ene reactions for peptide and protein macrocyclizations." *Chemistry* 23: 7087-7092 (2017).

- (125) C Bergonzo and TE Cheatham, III. "Mg²⁺ binding promotes SLV as a scaffold in Varkud Satellite Ribozyme SLI-SLV kissing loop junction." *Biophys. J.* 113, 313-320 (2017) DOI:10.1016/j.bpj.2017.06.008.
- (126) R Galindo-Murillo and TE Cheatham III. "Computational DNA binding studies of (-)-epigallocatechin-3-gallate." *J. Biomol. Struct. Dyn.* 36, 3311-3323 (2017). DOI:10.1080/07391102.2017.1389306.
- (127) M Xu, R Galindo-Murillo, TE Cheatham III, and RM Franzini RM. "Dissociative reactions of benzonorbornadienes with tetrazines: scope of leaving groups and mechanistic insights." *Org. Biomol. Chem.* 15, 9855-9865 (2017). DOI:10.1039/c7ob02191g.
- (128) HS Hayatshahi, C Bergonzo, and TE Cheatham III. "Investigating the ion dependence of the first unfolding step of GTPase-associating center ribosomal RNA." *J. Biomol. Struct. Dyn.* 36, 243-253 (2018) DOI:10.1080/07391102.2016.1274272.
- (129) HS Hayatshahi, NM Henriksen, and TE Cheatham III. "Consensus conformations of dinucleotide monophosphates described with well-converged molecular dynamics simulations." *J. Chem. Theory Comp.* 14, 1456-1470 (2018). DOI:10.1021/acs.jctc.7b00581.
- (130) SP Cornillie, BJ Bruno, CS Lim, and TE Cheatham III. "Computational modeling of stapled peptides toward a treatment strategy for CML and broader implications in the design of lengthy peptide therapeutics." *J. Phys. Chem. B* 122, 3864-3875 (2018). doi: 10.1021/acs.jpcc.8b01014.
- (131) IM Chrisman, M Nemetcheck, I Mitchell de Vera, J Shang, Z Heidari, Y Long, H Reyes-Caballero, R Galindo-Murillo, T Cheatham, A Blayo, Y Shin, J Fuhrmann, P Griffin, T Kamenecka, D Kojetin, and T Hughes "Defining a conformational ensemble that directs activation of PPAR γ ." *Nature Comm.* 9, 1794 (2018) doi: 10.1038/s41467-018-04176-x.
- (132) I Paraskevagos, A Luckow, M Khoshlessan, G Chantzialexiou, TE Cheatham, III, O Beckstein, GC Fox, and S. Jha. "Task-parallel analysis of molecular dynamics trajectories." ACM, ICPP 2018 Proceedings of the 47th International Conference on Parallel Processing, Eugene OR, August 13-16, 2018, article 49. doi:10.1145/3225058.3225128
- (133) DR Roe and TE Cheatham, III "Parallelization of CPPTRAJ enables large scale analysis of molecular dynamics trajectory data." *J. Comp. Chem.* 39, 2110-2117 (2018) doi:10.1002/jcc.25382.
- (134) R Galindo-Murillo, TE Cheatham III, and RC Hopkins "Exploring potentially alternative non-canonical DNA duplex structures through simulation." *J. Biomol. Struct. Dyn.* 37, 2201-2210 (2019) doi: 10.1080/07391102.2018.1483839
- (135) J Ma, A Culich, T Battelle, D Brunson, T Cheatham, J Goodhue, C Hill, L Passin, J Smith, and S Yockel. "On launching Ask.Cl, a Q & A platform for research computing using StackExchange and Discouse" In *SC18 Proceedings* (2018). DOI: <https://doi.org/10.22369/issn.2153-4136/12/2/9>
- (136) AJ Monaghan, B Milash, T Magle, T Hauser, TE Cheatham, PJ Burns, S Knuth, AM Orendt, S White, D Paschal, HJ Siegel, T Kaiser, AM Johnson, and B Yeager. "Improving regional cyberinfrastructure services through collaboration: Cyberteam for the Rocky Mountain Advanced Computing Consortium." In *Practice and Experience in Advanced Research Computing (PEARC' 19)*, July 28-August 1, 2019, Chicago, IL, USA. ACM, New York, NY, USA, 8 pages. <https://doi.org/10.1145/3332186.3332232>.
- (137) CA Stewart, DY Hancock, J Wernert, T Furlani, D Lifka, A Sill, N Berente, DF McMullen, T Cheatham, A Apon, R Payne, and SD Slavin. "Assessment of non-financial returns cyberinfrastructure: A survey of current methods." In *JARC '19 Proceedings of the Humans in the Loop: Enabling and Facilitating Research on Cloud Computing*, July 2019, 2, 1-10 (2019). pages. <https://doi.org/10.1145/3355738.3355749>.

- (138) CA Stewart, DY Hancock, J Wernert, T Furlani, D Lifka, A Sill, N Berente, DF McMullen, T Cheatham, A Apon, R Payne, and SD Slavin. 2019. "Assessment of financial returns on investments in cyberinfrastructure facilities: A survey of current methods." In Practice and Experience in Advanced Research Computing (PEARC '19), July 28-August 1, 2019, Chicago, IL, USA. ACM, New York, NY, USA, 8 pages. <https://doi.org/10.1145/3332186.3332228>.
- (139) R Galindo-Murillo and TE Cheatham, III. "Lessons learned in atomistic simulation of double-stranded DNA: Solvation and salt concerns." *Living J. Comp. Mol. Sci.* <https://doi.org/10.33011/livecoms.1.2.9974> (2019).
- (140) G Fox, O Beckstein, J Qui, D Crandall, G von Laszewski, J Paden, S Jha, F Wang, M Marathe, A Vullikanti, and T Cheatham. "Contributions to High-Performance Big Data Computing" In Future Trends of HPC in a Disruptive Scenario IOS Press, Volume 34 of Advances in Parallel Computing (2019).
- (141) J Kessler, G Kang, Z Qin, H Kang, F Whitby, T Cheatham, C Hill, Y Li, and SM Yu. "Peptoid residues make diverse hyperstable collagen triple helices". *ChemRxiv*8/9/2020. <https://doi.org/10.26434/chemrxiv.12921995.v1>
- (142) A Roy, D Roe, M Hall, T Cheatham. "Polyhedral compilation support for C++ features: A case study with CPPTRAJ". In: Rauchwerger, L. (eds) Languages and Compilers for Parallel Computing. LCPC 2017. Lecture Notes in Computer Science(), vol 11403. Springer, Cham. https://doi.org/10.1007/978-3-030-35225-7_3
- (143) S Broude Geva, D Brunson, T Cheatham III, J Deaton, J Griffioen, CW Hillegas, DM Jennewein, G Krovitz, T Magle, P Schmitz, K Tomko, and JC Wilgenbusch. "Fostering Collaboration Among Organizations in the Research Computing and Data Ecosystem". In Practice and Experience in Advanced Research Computing (PEARC '20), July 26–30, 2020, Portland, OR, USA. ACM, New York, NY, USA, 13 pages (2020). <https://doi.org/10.1145/3311790.3396645>
- (144) P Schmitz, C Mizumoto, J Hicks, D Brunson, G Krovitz, JR Bottum, J Cutcher-Gershenfeld, K Wetzel, and T Cheatham III. "A Research Computing and Data Capabilities Model for Strategic Decision-Making". In Practice and Experience in Advanced Research Computing (PEARC '20), July 26–30, 2020, Portland, OR, USA. ACM, New York, NY, USA, 12 pages (2020). <https://doi.org/10.1145/3311790.3396643>
- (145) R Galindo-Murillo, L Winkler, JC Garcia-Ramos, L Ruiz-Azuara, F Cortez-Guzman, and TE Cheatham III. "Ancillary Ligand in Ternary Cu^{II} Complexes Guides Binding Selectivity Toward Minor-Groove DNA." *J. Phys. Chem. B* 124, 11648-11658 (2020). <https://doi.org/10.1021/acs.jpcc.0c09296>
- (146) J Ma, T Battelle, K Bulekova, A Culich, J Goodhue, J Pessin, V Sochat, D Brunson, T Cheatham, S Najafi, C Hill, A Del Maestro, B Segee, R Zottola, S Valcourt, Z Braiterman, R Singh, R Thoelen, and J Smith. "Ask.Cyberinfrastructure.org: Creating a Platform for Self-Service Learning and Collaboration in the Rapidly Changing Environment of Research Computing" *J. Comp. Sci. Education* (2021). DOI: <https://doi.org/10.22369/issn.2153-4136/12/2/9>
- (147) R Galindo-Murillo and TE Cheatham, "Ethidium bromide interactions with DNA: an exploration of a classic DNA-ligand complex with unbiased molecular dynamics simulations." *Nucleic Acids Res.* 49, 3735-3747 (2021). <https://doi.org/10.1093/nar/gkab143>
- (148) P Schmitz, S Yockell, C Mizumoto, T Cheatham III, and Dana Brunson, "Advancing the Workforce that Supports Computationally and Data Intensive Research". *Computing in Sci. & Eng.* Sept.-Oct. pp. 19-27, vol 23 (2021). <https://doi.org/10.1109/MCSE.2021.309842>
- (149) JL Kessler, G Kang, Z Qin, H Kang, FG Whitby, TE Cheatham III, CP Hill, Y Li, and SM Yu, "Peptoid Residues Make Diverse, Hyperstable Collagen Triple-Helices". *J. Amer. Chem. Soc.* 143, 29, 10910-10919 (2021). <https://doi.org/10.1021/jacs.1c00708>

- (150) G da Rosa, L Grille, V Calzada, K Ahmad, JP Arcon, F Battistini, G Bayarri, T Bishop, P Carloni, T Cheatham III, R Collepardo-Guevara, J Czub, JR Espinosa, R Gallino-Murillo, SA Harris, A Hospital, C Laughton, JH Maddocks, A Noy, M Orozco, M Pasi, A Perez, D Petkeviciute-Gerlach, R Sharma, R Sun, and PD Dans. "Sequence-dependent structural properties of B-DNA: what have we learned in 40 years?" *Biophysical Reviews* 13, 995-1005 (2021). <https://doi.org/10.1007/s12551-021-00893-8>
- (151) T Rajasekaran, GC Freestone, R Galindo-Murillo, B Lugato, L Rico, JC Salinas, H Gaus, MT Migawa, EE Swayze, TE Cheatham, III, S Hanessian, and PP Seth, "Backbone hydrocarbon-constrained nucleic acids modulate hybridization kinetics for RNA". *J. Amer. Chem. Soc.* 144, 1941-1950 (2022). <https://doi.org/10.1021/jacs.1c12323>
- (152) R Galindo-Murillo*, L Winkler*, J Ma, F Hanelli, A Fleming, C Burrows, and T Cheatham "Riboflavin Stabilizes Abasic, Oxidized G-Quadruplex Structures". *Biochemistry* 61, 265-275 (2022). <https://doi.org/10.1021/acs.biochem.1c00598>
- (153) OD Love, MCP Lima, CH Clark, S Cornillie, SM Roalstad, TE Cheatham III. "Evaluating the accuracy of the AMBER protein force fields in modeling dihydrofolate reductase structures: Misbalance in the conformational arrangements of the flexible loop domains". *J. Biomol. Struct. Dyn.* (2022). <https://doi.org/10.1080/07391102.2022.2098823>
- (154) R Galindo-Murillo and TE Cheatham III. "Transient Hoogsteen Base Pairs Observed in Unbiased Molecular Dynamics Simulations of DNA" *J. Phys. Chem. Lett* 13, 6283-6287 (2022). <https://doi.org/10.1021/acs.jpcclett.2c01348>
- (155) T Hacker, P Smith, D Brunson, L Arafune, T Cheatham, and E Deelman. "Building the research innovation workforce: Challenges and recommendations from a virtual workshop to advance the research computing community." In Practice and Experience in Advanced Research Computing (PEARC '22), July 10–14, 2022, Boston, MA, USA. ACM, New York, NY, USA 7 Pages. <https://doi.org/10.1145/3491418.3530288>
- (156) T Rajasekaran, G Freestone, R Galindo-Murillo, B Lugato, H Gaus, M Migawa, E Swayze, TE Cheatham, III, P Seth and S Hanessian. "Systematic investigation of tether length and phosphorous configuration in backbone constrained macrocyclic nucleic acids to modulate binding kinetics for RNA". *J. Organic Chem.* 88, 3599-3614 (2023). <https://doi.org/10.1021/acs.joc.2c02796>
- (157) L Winkler, R Galindo-Murillo and TE Cheatham, III. "The structures and dynamics of DNA mini-dumbbells are force field dependent". *J. Chem. Theory Comp.* 19, 2198-2212 (2023). <https://doi.org/10.1021/acs.jctc.3c00130>
- (158) L Winkler and TE Cheatham, III. "Benchmarking the Drude polarizable force field using the r(GACC) tetranucleotide." *J. Chem. Info. Modeling* 63, 2505-2511 (2023). <https://doi.org/10.1021/acs.jcim.3c00250>
- (159) O Love, R Galindo-Murillo, M Zgarbová, J Šponer, P Jurečka and TE Cheatham, III. "Assessing the Current State of Amber Force Field Modifications for DNA—2023 Edition." *J. Chem. Theory Comput.* 19, 4299-4307 (2023). <https://doi.org/10.1021/acs.jctc.3c00233>
- (160) J Cutcher-Gershenfeld, T Middelkoop, D Brunson, T Cheatham, J Fosso Tande, D Jennewein, T Battelle, J Ma, LA Michael, H Neeman and P Schmitz. "Professionalization of Research Computing and Data: An Expanded Agenda" *PEARC '23: Practice and Experience in Advanced Research Computing* 129-136 (2023). Won Best Paper in the Workforce Development Track.
- (161) DA Case, HM Aktulga, K Belfon, DS Cerutti, GA Cisneros, VWD Cruzeiro, N Forouzes, TJ Giese, AW Gotz, H Gohlke, S Izadi, K Kasavajhala, MC Kaymak, E King, T Kurtzman, TS Lee, P Li, J Liu, T Luchko, R Luo, M Manathinga, MR Machado, HM Nguyen, KA O'Hearn, A Onufriev, F Pan, S Pantano, R Qi, A Rahnamoun, A Risheh, S Schott-Verdugo, A Shajan, J Swails, J Wang, H Wei, X Wu, S Zhang, S Zhao, Q Zu, TE Cheatham, III, DR Roe, A Roitberg, C Simmerling, DM York, MC

- Nagan and KM Merz, Jr. "AmberTools". *J. Chem Info Modeling* 63, 6183-6191 (2023)
<https://doi.org/10.1021/acs.jcim.3c01153>.
- (162) L Winkler, R Galindo-Murillo, and TE Cheatham, III. "Assessment of A- to B- DNA Transitions Utilizing the Drude Polarizable Force Field" *J. Chem. Theory Comp.* 19: 8955-8966 (2023)
<https://doi.org/10.1021/acs.jctc.3c01002>.
- (163) WK Olson, JH Maddocks, PD Dans, TE Cheatham III, S Harris, C Laughton, M Orozco and L Pollack. "An open call for contributions to a special issue of *Biophysical Reviews* focused on multiscale simulations of DNA from electrons to nucleosomes." *Biophys. Rev.* 15, 1901-1902 (2023)
<https://doi.org/10.1007/s12551-023-0118>.
- (164) O Love, L Winkler, and TE Cheatham, III. "van der Waals parameter scanning with Amber nucleic acid force fields: Revisiting means to better capture the RNA/DNA structure through MD." *J. Chem. Theory Comput.* 20, 625-643 (2024) <https://doi.org/10.1021/acs.jctc.3c01164>.
- (165) DT Hansen, NJ Rueb, ND Levinzon, TE Cheatham III, R Gaston Jr., K Tanvir Ahmed, S Osburn-Staker, JE Cox, GB Dudley, and AM Barrios. "The mechanism of covalent inhibition of LAR phosphatase by illudalic acid." *Bioorg. Med. Chem. Lett.* 104:129740 (2024)
<https://doi.org/10.1016/j.bmcl.2024.129740>.
- (166) JH Maddocks, PD Dans, TE Cheatham III, S Harris, C Laughton, M Orozco, L Pollack and WK Olson. "Special issue: Multiscale simulations of DNA from electrons to nucleosomes." *Biophys. Rev.* 16, 259-262 (2024) <https://doi.org/10.1007/s12551-024-01204-7>.
- (167) MCP Lima, BD Hornsby, CS Lim, and TE Cheatham III. "Molecular modeling of single- and double-hydrocarbon stapled coiled-coil inhibitors against Bcr-Abl: Toward a treatment strategy for CML." *J. Phys. Chem. B.* 128,6476-6491 (2024) <https://doi.org/10.1021/acs.jpcc.4c02699>.
- (168) O Love, R Galindo-Murillo, D Roe, P Dans, TE Cheatham, III and C Bergonzo. "modXNA: A modular approach to parameterization of modified nucleic acids for use with Amber force fields." *J. Chem. Theory Comp.* 20, 9354-9393 (2024) <https://doi.org/10.1021/acs.jctc.4c01164>.

Abstracts:

MA Munger, FS Albright, JE Buskupiak, DK Blumenthal, TE Cheatham, III, FM Creekmore. "Utility of the foreign graduate equivalency examination to assess pharmacy curricula". 107th annual meeting AACP (2006).

JE Buskupiak, DK Blumenthal, TE Cheatham, III, FS Albright, MA Munger. "An employer survey as a pharmacy curriculum assessment tool". 108th annual meeting AACP (2007).

Recent technical training and teaching experience

- **PharmD Recitation**, Pharm 5151: Spring 2018, 2019, 2020, 2022, 2023
 2 credits; 26 contact hours, Co-Course Master
 1 credit; 15 contact hours, Co-Course Master Spring 2023
- **Organic Medicinal Chemistry**, MDCH 5210: 1st half Fall semester
 2 credits; 16-20 contact hours, 2002-2015, Course Master.
- **Principles of Medicinal Chemistry**, MDCHEM 6990: ½ Spring semester.
 2 credits, ~3-6 contact hours, Spring 2006-2013.
- **Mechanisms and Kinetics of Drug Degradation & Stabilization / Biotechnology**, PHCEU 7040
 Fall 2002-2011. 8-12 hours in 2002-2011, 4 hours in 2013.
- **Molecular Modeling and Biomolecular Simulation from a Pharmaceutical Perspective**, PHCEU/MDCHEM 7095: 2nd half Spring semester.

- 2 credits, ~30 contact hours, Spring 2004, 2005, 2006, 2011, 2012, 2013; Course Master.
- **Introduction to Bioinformatics**, MDINF 6600.
 - 3 lectures on structure prediction, Spring 2002-2006.
- **RNA/DNA as a Drug Target Journal Club**, PHCEU 5660.
 - 1 credit/15 hours, Fall 2004, Course Master.
- **Fundamentals of Pharmaceutical Sciences**, PHSCI 7113.
 - 2 lectures on drug development, 1 on solutions; Fall 2004-present.

Selected Presentations and Workshops (2014-present)

- University of Indiana School of Medicine, Biochemistry Seminar, September 2024 [invited]
- International HPC Summer School, Kobe, Japan, July 2024 [invited]
- PEARC24, Providence, RI, July 2024 [invited]
- ISQBP Meeting, May 2024
- ACS National Meeting, New Orleans, March 2023 [invited]
- AMBER Developers Meeting, March 2023 [invited]
- SHEKATE Workshop, University of Utah, November 2023 [invited]
- Molecular Dynamics Database Workshop, Oxford University, October 2023 [invited]
- U of Utah Bioscience Symposium, September 2023 [invited]
- ACS National Meeting, San Francisco, August 2023 [invited]
- RNA Dynamics, Telluride, July 2023 [invited]
- PEARC23, Portland, July 2023 [invited]
- ABC Meeting, Ascona, April 2023 [invited]
- CASC Spring 2023 meeting, Arlington, March 2023 [invited]
- AMBER Developers Meeting, March 2023 [invited]
- UNICAMP, Center for Computational Science and Engineering, Retreat, December 2022 [invited]
- CASC Fall 2022 meeting, Crystal City, VA, October 2022 [invited]
- ACS National Meeting, Chicago, August 2022 [invited]
- Gordon Research Conference, Computational Chemistry, Spain, July 2022
- ISQBP Presidents meeting, Austria, July 2022 [invited]
- RCD-Nexus day @ PEARC22, July 2022
- International HPC summer school, Greece, June 2022 [invited]
- Supercomputing21, St. Louis, November 2021
- EMBO Virtual Workshop on Advances and Challenges in Biomolecular Simulations, October 2021 [invited]
- PEARC21, July 2021 [invited]
- ISQBP meeting, June 2021 [invited]
- AAU/APLU Workshop on Accelerating Public Access to Research Data, May 2021 [invited]
- CASC Spring 2021 meeting, April 2021 [invited]
- CASC Fall 2020 meeting, October 2020 [invited]
- AAU/APLU Conference on Accelerating Public Access to Research Data, October 2020 [invited]
- Building the research innovation workforce, August 2020 [invited, steering committee]
- PEARC20, July 2020 [invited to multiple workshops, events and paper presentations]
- CASC Spring 2020 meeting, March 2020 [invited]
- AAU/APLU Conference on Accelerating Public Access to Research Data, February 2020 [invited]
- AMBER Developers Meeting, Safety Harbor FL, February 2020 [invited]
- Westnet CIO meeting, January 2020 [invited]
- Texas Advanced Supercomputing Center Frontera Futures Workshop, January 2020 [invited]
- UNICAMP, Center for Computational Science and Engineering, Retreat, November 2019 [invited]

- CaRCC Sustainability workshop, DFW, October 2019 [organizer]
- Research Computing and Data Maturity model workshop, EDUCAUSE, October 2019 [organizer]
- CASC meeting, Arlington VA, September 2019 [invited]
- National Research Platform Workshop, Minneapolis MN, September 2019 [organizing committee]
- NSF CI for Large Facilities Workshop, Arlington VA, September 2019 [organizing committee]
- TACC Frontera Dedication, Austin TX, September 2019 [invited]
- RMACC Systems Administrator meet-up, Tempe AZ, August 2019
- PEARC19, Chicago IL, July 2019 [workshop organizer, BoF, Town Hall, keynote, panels]
- International HPC Summer School, Kobe Japan, July 2019 [invited]
- NSF CI Coordination Workshop, Arlington VA, June 2019 [facilitator/organizer]
- BioExcel/PRACE HPC for Life Sciences, Stockholm Sweden, June 2019 [invited]
- Blue Waters Symposia, Sun River OR, June 2019 [invited]
- RMACC, U Colorado Boulder, May 2019
- Sealy Symposium, Galveston TX, May 2019 [invited]
- CaRCC Ecosystem of Research Computing and Data Workshop, St. Louis, April 2019 [organizer]
- ACS National Meeting, Orlando FL, April 2019 [1 invited, 1 contributed]
- AMBER Developers Meeting, Safety Harbor FL, March 2019 [invited]
- CASC, Arlington VA, March 2019 [invited]
- Theory-Software Workshop, New Orleans LA, February 2019 [invited]
- Internet2 Research Computing Maturity Model Workshop, December 2018 [invited]
- University of Missouri, State of Research Computing at Utah, November 2018 [invited]
- APLU-AUU Workshop on Accelerating Public Access to Research Data, October 2018 [invited]
- CASC, Arlington VA, October 2018 [invited]
- Georgia State University, Department of Chemistry, September 2018 [invited]
- RMACC, U Colorado Boulder, August 2018
- National Research Platform, U Montana Bozemann, August 2018
- PEARC18, Pittsburgh, PA, July 2018
- International HPC Summer School, Ostrava Czech Republic, July 2018 [invited]
- ISQBP, Barcelona Spain, June 2018 [invited]
- Blue Waters Symposia, Sun River OR, June 2018 [invited]
- CECAM on multiscale modeling of epigenetics, Laussane Switzerland, May 2018 [invited]
- RMACC System Administrators meet-up, Arizona State U, April 2018
- Sustainable Software Institute Workshop, Berkeley CA, April 2018 [invited]
- Open Science Grid all-hands meeting, University of Utah, March 2018 [invited]
- CASC, Arlington VA, March 2018 [invited]
- CI Professionalization Workshop, Arlington VA, March 2018 [invited]
- CENIC, Monterey CA, March 2018 [invited]
- AMBER Developers Meeting, Florida, February 2018 [invited]
- WestNet, Tucson AZ, January 2018 [invited]
- CASC, Arlington VA, October 2017 [invited]
- 7th iCatse International Conference on IT Convergence and Security, Seoul Korea, September 2017 [invited]
- Lucian Symposium, St Edwards College, Austin TX, September 2017 [invited]
- National Research Platform workshop, Bozeman MT, August 2017 [invited]
- RNA Dynamics @ Telluride, July 2017
- ARCC Tutorial and CaRC @ PEARC17. New Orleans, July 2017 [invited]
- International HPC Summer School, Boulder Co., June 2017 [invited]
- DE Shaw Research, New York, NY, June 2017 [invited]
- Utah Education and Telehealth Network Tech Summit, Salt Lake City, UT, June 2017 [invited]

- MolSSI workshop, Blacksburg VA, June 2017 [invited]
- DDN Best Practices in Data, Boston MA, May 2017 [invited]
- Blue Waters Symposia, Sun River OR, May 2017 [invited]
- NSF workshop in Big Data in Chemistry, Arlington VA, April 2017 [invited]
- PRACE workshop on Gromacs, AMBER and NAMD, Stockholm Sweden, April 2017 [invited]
- CI Practitioners workshop, Washington DC, March 2017 [invited]
- NSF Workshop on Integrating Computation into Biology and Chemistry, New Orleans, March 2017 [invited]
- OSG All Hands, San Diego CA, March 2017 [invited]
- NSF SI2 PI Meeting, Arlington VA, February 2017 [invited]
- AMBER Developers Meeting, Athens GA, February 2017 [invited]
- ACI-REF face to face meeting, Miami FL, February 2017 [invited]
- Westnet CIO meeting, Tempe AZ, January 2017 [invited]
- NSF Cloud workshop, Arlington VA, December 2016 [invited]
- SIGUCCS. Denver, Panel, November 2016 [invited]
- Pacific Rim Platform – ACI-REF joint meeting at Cal-IT2, October 2016 [invited]
- RMACC workshop, Colorado State U, August 2016 [invited]
- International HPC Summer School, Ljubljana, Slovenia, June 2016 [invited]
- ISQBP President's meeting, Bergen, Norway, June 2016 [invited]
- Blue Waters Symposia, Sun River, OR, June 2016 [invited]
- RNA: Structure, Dynamics and Function, SISSA, Trieste, Italy, May 2016 [invited]
- Stockholm University, Lindahl Lab, Stockholm, April 2016 [invited]
- Arizona State University, Computational Physics Seminar, April 2016 [invited]
- Advanced Research Computing on Campuses, UIUC, Urbana, March 2016 [invited]
- AMBER Developers Meeting, San Diego, March 2016 [invited]
- ACS National Meeting, San Diego, March 2016 [invited]
- RNA Structure Prediction workshop, Punta Cana, Dominican Republic, December 2015 [invited]
- SaintCon Utah's IT security conference, Weber State, October 2015 [invited]
- Mexico National Chemistry Congress, Queretaro, October 2015 [keynote]
- RMACC workshop, Boulder Colorado, August 2015 [invited]
- Zing Computational Biological Chemistry, Cairnes Australia, August 2015 [invited]
- RNA Dynamics, Telluride, July 2015 [invited]
- AMBER workshop at University College London, July 2015 [invited]
- International HPC Summer School, Toronto, June 2015 [invited]
- 19th Conversation, University of Albany, NY, June 2015 [invited]
- Blue Waters Symposium, Sunriver, OR, May 2015 [invited]
- ACS National Meeting, David Case Awards Symposia, Denver, CO, March 2015 [invited]
- Advanced Research Computing on Campuses, Researcher Perspectives Panel, Clemson, March 2015 [invited]
- AMBER developers meeting, University of Florida, February 2015
- National Academy of Sciences Future of NSF HPC testimony, Mountain View, December 2014 [invited]
- AMBER workshop at UNAM, Mexico City, October 2014 [organizer]
- CHANGES 2014, Chinese Academy of Sciences, Beijing, September 2014 [invited]
- XXVI IUPAP Conference on Computational Physics, Boston, August 2014 [invited]
- XSEDE14, Atlanta, July 2014 [Science Track Chair, Presenter]
- ISQBP Presidents Meeting, Telluride, June 2014 [President, Organizer]
- International HPC Summer School, Budapest, June 2014
- Sandia National Labs, Biomaterials Science, May 2014
- Blue Waters Symposia, U Illinois, May 2014
- CECAM Biomolecules in unnatural conditions, Stuttgart, March 2014

- American Physical Society National Meeting, Denver, March 2014
- St. Louis University, Department of Chemistry, January 2014
- AMBER Developers Meeting, Stony Brook U, January 2014

Committee appointments (Utah)

Utah Education Network Education Advisory Council (2016-)
 University of Utah, Senate Advisory Committee on Information Technology (2016-2025)
 University of Utah, Responsible AI Community of Practice (2023-)
 University of Utah, Data Infrastructure & Training Committee (2023-)
 University of Utah, DELPHI Infrastructure committee (2022-)
 University of Utah, University Academic Center and Institute Coordinating Committee (2021-2024)
 University of Utah, Academic Senate Executive Committee (2019-2022, 2024-)
 University of Utah, Academic Senate (2019-2022, 2024-)
 University of Utah, Senate Advisory Committee on Information Technology (2016-2019, 2019-2022)
 University of Utah, Senate Advisory Committee on Strategic Planning (2020-2022)
 University of Utah, Director of Graduate Studies Medicinal Chemistry (2011-2017)
 University of Utah, Graduate Council Review Committee, Dept. of Atmospheric Sciences (2017)
 University of Utah, Senate Ad Hoc Faculty Information Technology Committee (2015-2016)
 University of Utah, Information Technology Research Portfolio Committee, Chair (2013-2015)
 University of Utah, Operational Information Technology Committee (2013-2015)
 University of Utah, Academic Senate (2012-2015)
 University of Utah, Personnel and Elections Committee (2012-2015)
 University of Utah, CyberInfrastructure Advisory Committee (2006-2007)
 University of Utah, Bioinformatics Advisory Committee (2002-2004)
 University of Utah Information Technology Leadership Team (2015-)
 U Utah, Health Sciences Center Information Technology Projects Approval (2001-2003)
 University of Utah, Cyberinfrastructure Council (2007-2012)
 University of Utah Information Technology Council (2001-2012)
 Center for High Performance Computing, User Advisory Committee (2001-, ad hoc 2015-)
 College of Pharmacy, Curriculum Committee (2019-)
 College of Pharmacy, Mentoring Committee (2012-)
 College of Pharmacy, Outcomes Assessment Committee (2004-)
 College of Pharmacy, Computer Committee (2000-present, Chair 2007-2014, ad hoc 2014-)
 College of Pharmacy, PharmD Seminar Committee (2009-2010)
 Medicinal Chemistry, Seminar Coordinator (2003-2004, 2010-2011)

Committee appointments (External)

Community of Communities Working Group (for RCD) (2019-)
 Campus Research Computing Consortium, <https://carcc.org> – CARCC – Acting Chair, current activities include: Chairs Leadership Team, Logistics Working Group, Capabilities Model Working Group, Engagement Working Group, Staff and Student Workforce Development Interest Groups; Annual Retreat (2016-)
 Coalition of Academic Scientific Computation, University of Utah Representative (2014-)

Campus Champion (formerly XSEDE CC, CC Leadership Team 2020-2022) (2013-)
AMBER Executive Committee – Biomolecular Simulation Software Suite (2002-)