

Gregory A Voth

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536
papers

36,352
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96
h-index

164
g-index

596
ext. papers

39,725
ext. citations

5.3
avg, IF

7.85
L-index

#	Paper	IF	Citations
536	A multiscale coarse-graining method for biomolecular systems. <i>Journal of Physical Chemistry B</i> , 2005 , 109, 2469-73	3.4	879
535	Unique spatial heterogeneity in ionic liquids. <i>Journal of the American Chemical Society</i> , 2005 , 127, 12192-6	3.4	853
534	Flexible simple point-charge water model with improved liquid-state properties. <i>Journal of Chemical Physics</i> , 2006 , 124, 024503	3.9	742
533	On the Structure and Dynamics of Ionic Liquids. <i>Journal of Physical Chemistry B</i> , 2004 , 108, 1744-1752	3.4	599
532	The multiscale coarse-graining method. I. A rigorous bridge between atomistic and coarse-grained models. <i>Journal of Chemical Physics</i> , 2008 , 128, 244114	3.9	550
531	The computer simulation of proton transport in water. <i>Journal of Chemical Physics</i> , 1999 , 111, 9361-9381	3.9	515
530	Multiscale coarse graining of liquid-state systems. <i>Journal of Chemical Physics</i> , 2005 , 123, 134105	3.9	467
529	Rigorous formulation of quantum transition state theory and its dynamical corrections. <i>Journal of Chemical Physics</i> , 1989 , 91, 7749-7760	3.9	448
528	The formulation of quantum statistical mechanics based on the Feynman path centroid density. II. Dynamical properties. <i>Journal of Chemical Physics</i> , 1994 , 100, 5106-5117	3.9	399
527	Molecular dynamics simulation of nanostructural organization in ionic liquid/water mixtures. <i>Journal of Physical Chemistry B</i> , 2007 , 111, 4812-8	3.4	395
526	Multistate Empirical Valence Bond Model for Proton Transport in Water. <i>Journal of Physical Chemistry B</i> , 1998 , 102, 5547-5551	3.4	376
525	Coarse-graining methods for computational biology. <i>Annual Review of Biophysics</i> , 2013 , 42, 73-93	21.1	373
524	Molecular Dynamics Simulation of Ionic Liquids: The Effect of Electronic Polarizability. <i>Journal of Physical Chemistry B</i> , 2004 , 108, 11877-11881	3.4	367
523	Computer simulation of proton solvation and transport in aqueous and biomolecular systems. <i>Accounts of Chemical Research</i> , 2006 , 39, 143-50	24.3	365
522	Tail aggregation and domain diffusion in ionic liquids. <i>Journal of Physical Chemistry B</i> , 2006 , 110, 18601-8	3.4	365
521	Ab initio molecular dynamics: Propagating the density matrix with Gaussian orbitals. III. Comparison with Born-Oppenheimer dynamics. <i>Journal of Chemical Physics</i> , 2002 , 117, 8694-8704	3.9	361
520	Multiscale modeling of biomolecular systems: in serial and in parallel. <i>Current Opinion in Structural Biology</i> , 2007 , 17, 192-8	8.1	359

519	Effective force fields for condensed phase systems from ab initio molecular dynamics simulation: a new method for force-matching. <i>Journal of Chemical Physics</i> , 2004 , 120, 10896-913	3.9	349
518	Ab initio molecular dynamics: Propagating the density matrix with Gaussian orbitals. II. Generalizations based on mass-weighting, idempotency, energy conservation and choice of initial conditions. <i>Journal of Chemical Physics</i> , 2001 , 115, 10291	3.9	320
517	The quantum dynamics of an excess proton in water. <i>Journal of Chemical Physics</i> , 1996 , 104, 2056-2069	3.9	314
516	Understanding ionic liquids through atomistic and coarse-grained molecular dynamics simulations. <i>Accounts of Chemical Research</i> , 2007 , 40, 1193-9	24.3	285
515	The multiscale coarse-graining method. II. Numerical implementation for coarse-grained molecular models. <i>Journal of Chemical Physics</i> , 2008 , 128, 244115	3.9	284
514	The formulation of quantum statistical mechanics based on the Feynman path centroid density. I. Equilibrium properties. <i>Journal of Chemical Physics</i> , 1994 , 100, 5093-5105	3.9	273
513	The formulation of quantum statistical mechanics based on the Feynman path centroid density. IV. Algorithms for centroid molecular dynamics. <i>Journal of Chemical Physics</i> , 1994 , 101, 6168-6183	3.9	268
512	A second generation multistate empirical valence bond model for proton transport in aqueous systems. <i>Journal of Chemical Physics</i> , 2002 , 117, 5839-5849	3.9	266
511	Special pair dance and partner selection: elementary steps in proton transport in liquid water. <i>Journal of Physical Chemistry B</i> , 2008 , 112, 9456-66	3.4	263
510	Proton solvation and transport in aqueous and biomolecular systems: insights from computer simulations. <i>Journal of Physical Chemistry B</i> , 2007 , 111, 4300-14	3.4	260
509	Ionic liquids. <i>Accounts of Chemical Research</i> , 2007 , 40, 1077-8	24.3	243
508	The Hydrated Proton at the Water Liquid/Vapor Interface. <i>Journal of Physical Chemistry B</i> , 2004 , 108, 14804-14806	3.4	237
507	A derivation of centroid molecular dynamics and other approximate time evolution methods for path integral centroid variables. <i>Journal of Chemical Physics</i> , 1999 , 111, 2371-2384	3.9	230
506	Further developments in the local-orbital density-functional-theory tight-binding method. <i>Physical Review B</i> , 2001 , 64,	3.3	222
505	The curious case of the hydrated proton. <i>Accounts of Chemical Research</i> , 2012 , 45, 101-9	24.3	215
504	Mixed atomistic and coarse-grained molecular dynamics: simulation of a membrane-bound ion channel. <i>Journal of Physical Chemistry B</i> , 2006 , 110, 15045-8	3.4	213
503	A bond-order analysis of the mechanism for hydrated proton mobility in liquid water. <i>Journal of Chemical Physics</i> , 2005 , 122, 14506	3.9	212
502	The formulation of quantum statistical mechanics based on the Feynman path centroid density. III. Phase space formalism and analysis of centroid molecular dynamics. <i>Journal of Chemical Physics</i> , 1994 , 101, 6157-6167	3.9	209

501	A new perspective on quantum time correlation functions. <i>Journal of Chemical Physics</i> , 1993 , 99, 10070-10073	3.9	204
500	An improved multistate empirical valence bond model for aqueous proton solvation and transport. <i>Journal of Physical Chemistry B</i> , 2008 , 112, 467-82	3.4	203
499	Direct observation of Bin/amphiphysin/Rvs (BAR) domain-induced membrane curvature by means of molecular dynamics simulations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 15068-72	11.5	196
498	The vibrational spectrum of the hydrated proton: Comparison of experiment, simulation, and normal mode analysis. <i>Journal of Chemical Physics</i> , 2002 , 116, 737-746	3.9	189
497	Structural basis of membrane bending by the N-BAR protein endophilin. <i>Cell</i> , 2012 , 149, 137-45	56.2	188
496	The properties of water: insights from quantum simulations. <i>Journal of Physical Chemistry B</i> , 2009 , 113, 5702-19	3.4	187
495	Structure of the liquid-vacuum interface of room-temperature ionic liquids: a molecular dynamics study. <i>Journal of Physical Chemistry B</i> , 2006 , 110, 1800-6	3.4	185
494	When Physics Takes Over: BAR Proteins and Membrane Curvature. <i>Trends in Cell Biology</i> , 2015 , 25, 780-793	12.3	183
493	Well-tempered metadynamics converges asymptotically. <i>Physical Review Letters</i> , 2014 , 112, 240602	7.4	182
492	Path integral centroid variables and the formulation of their exact real time dynamics. <i>Journal of Chemical Physics</i> , 1999 , 111, 2357-2370	3.9	178
491	A quantum model for water: Equilibrium and dynamical properties. <i>Journal of Chemical Physics</i> , 1997 , 106, 2400-2410	3.9	176
490	The formation and dynamics of proton wires in channel environments. <i>Biophysical Journal</i> , 2001 , 80, 1691-702	7.02	169
489	An accurate and simple quantum model for liquid water. <i>Journal of Chemical Physics</i> , 2006 , 125, 184507	3.9	165
488	Molecular dynamics simulations of imidazolium-based ionic liquid/water mixtures: Alkyl side chain length and anion effects. <i>Fluid Phase Equilibria</i> , 2010 , 294, 148-156	2.5	164
487	Coarse-grained modeling of the actin filament derived from atomistic-scale simulations. <i>Biophysical Journal</i> , 2006 , 90, 1572-82	2.9	161
486	Allostery of actin filaments: molecular dynamics simulations and coarse-grained analysis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 13111-6	11.5	161
485	The Mechanism of Hydrated Proton Transport in Water. <i>Journal of the American Chemical Society</i> , 2000 , 122, 12027-12028	16.4	161
484	Coarse-grained peptide modeling using a systematic multiscale approach. <i>Biophysical Journal</i> , 2007 , 92, 4289-303	2.9	160

483	Multiscale coarse-graining and structural correlations: connections to liquid-state theory. <i>Journal of Physical Chemistry B</i> , 2007 , 111, 4116-27	3.4	160
482	Modeling real dynamics in the coarse-grained representation of condensed phase systems. <i>Journal of Chemical Physics</i> , 2006 , 125, 151101	3.9	160
481	Transition State Dynamics and Relaxation Processes in Solutions: A Frontier of Physical Chemistry. <i>The Journal of Physical Chemistry</i> , 1996 , 100, 13034-13049		159
480	Multiscale coarse-graining of ionic liquids. <i>Journal of Physical Chemistry B</i> , 2006 , 110, 3564-75	3.4	158
479	Characterization of the solvation and transport of the hydrated proton in the perfluorosulfonic acid membrane nafion. <i>Journal of Physical Chemistry B</i> , 2006 , 110, 18594-600	3.4	156
478	Hydroxide Solvation and Transport in Anion Exchange Membranes. <i>Journal of the American Chemical Society</i> , 2016 , 138, 991-1000	16.4	155
477	Mechanism of membrane curvature sensing by amphipathic helix containing proteins. <i>Biophysical Journal</i> , 2011 , 100, 1271-9	2.9	147
476	Quantum effects in liquid water from an ab initio-based polarizable force field. <i>Journal of Chemical Physics</i> , 2007 , 127, 074506	3.9	142
475	The theory of electron transfer reactions: what may be missing?. <i>Journal of the American Chemical Society</i> , 2003 , 125, 7470-8	16.4	138
474	Proton Transfer in the Enzyme Carbonic Anhydrase: An ab Initio Study. <i>Journal of the American Chemical Society</i> , 1998 , 120, 4006-4014	16.4	138
473	The mechanism of proton exclusion in aquaporin channels. <i>Proteins: Structure, Function and Bioinformatics</i> , 2004 , 55, 223-8	4.2	136
472	The 2018 biomembrane curvature and remodeling roadmap. <i>Journal Physics D: Applied Physics</i> , 2018 , 51,	3	133
471	Path-Integral Centroid Methods in Quantum Statistical Mechanics and Dynamics. <i>Advances in Chemical Physics</i> , 2007 , 135-218		133
470	Multiscale Coarse-Graining of Mixed Phospholipid/Cholesterol Bilayers. <i>Journal of Chemical Theory and Computation</i> , 2006 , 2, 637-48	6.4	133
469	Mechanism of fast proton transport along one-dimensional water chains confined in carbon nanotubes. <i>Journal of the American Chemical Society</i> , 2010 , 132, 11395-7	16.4	130
468	Friction Mediates Scission of Tubular Membranes Scaffolded by BAR Proteins. <i>Cell</i> , 2017 , 170, 172-184.e36.2	36.2	128
467	A systematic methodology for defining coarse-grained sites in large biomolecules. <i>Biophysical Journal</i> , 2008 , 95, 5073-83	2.9	124
466	Atomistic Modeling of the Electrode/Electrolyte Interface in Li-Ion Energy Storage Systems: Electrolyte Structuring. <i>Journal of Physical Chemistry C</i> , 2013 , 117, 3747-3761	3.8	122

465	Probing selected morphological models of hydrated Nafion using large-scale molecular dynamics simulations. <i>Journal of Physical Chemistry B</i> , 2010 , 114, 3205-18	3.4	121
464	Coarse-grained free energy functions for studying protein conformational changes: a double-well network model. <i>Biophysical Journal</i> , 2007 , 93, 3860-71	2.9	120
463	Linear aggregation of proteins on the membrane as a prelude to membrane remodeling. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 20396-401	11.5	119
462	Membrane tension controls the assembly of curvature-generating proteins. <i>Nature Communications</i> , 2015 , 6, 7219	17.4	119
461	A comparative study of imaginary time path integral based methods for quantum dynamics. <i>Journal of Chemical Physics</i> , 2006 , 124, 154103	3.9	119
460	Excess proton solvation and delocalization in a hydrophilic pocket of the proton conducting polymer membrane nafion. <i>Journal of Physical Chemistry B</i> , 2005 , 109, 3727-30	3.4	118
459	Hybrid Ab-Initio/Empirical Molecular Dynamics: Combining the ONIOM Scheme with the Atom-Centered Density Matrix Propagation (ADMP) Approach. <i>Journal of Physical Chemistry B</i> , 2004 , 108, 4210-4220	3.4	115
458	The Theory of Ultra-Coarse-Graining. 1. General Principles. <i>Journal of Chemical Theory and Computation</i> , 2013 , 9, 2466-80	6.4	114
457	Applications of higher order composite factorization schemes in imaginary time path integral simulations. <i>Journal of Chemical Physics</i> , 2001 , 115, 7832-7842	3.9	113
456	Systematic multiscale parameterization of heterogeneous elastic network models of proteins. <i>Biophysical Journal</i> , 2008 , 95, 4183-92	2.9	112
455	Infrared spectroscopy and hydrogen-bond dynamics of liquid water from centroid molecular dynamics with an ab initio-based force field. <i>Journal of Physical Chemistry B</i> , 2009 , 113, 13118-30	3.4	111
454	Proton solvation and transport in hydrated nafion. <i>Journal of Physical Chemistry B</i> , 2011 , 115, 5903-12	3.4	108
453	The formulation of quantum statistical mechanics based on the Feynman path centroid density. V. Quantum instantaneous normal mode theory of liquids. <i>Journal of Chemical Physics</i> , 1994 , 101, 6184-6192	3.9	107
452	Charge delocalization in proton channels, I: the aquaporin channels and proton blockage. <i>Biophysical Journal</i> , 2007 , 92, 46-60	2.9	106
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450	Calculation of solvent free energies for heterogeneous electron transfer at the water-metal interface: Classical versus quantum behavior. <i>Journal of Chemical Physics</i> , 1995 , 102, 529-539	3.9	102
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448	CarParrinello molecular dynamics simulation of liquid water: New results. <i>Journal of Chemical Physics</i> , 2002 , 116, 10372-10376	3.9	101

447	A role for a specific cholesterol interaction in stabilizing the Apo configuration of the human A(2A) adenosine receptor. <i>Structure</i> , 2009 , 17, 1660-1668	5.2	100
446	A Multiscale Coarse-Graining Study of the Liquid/Vacuum Interface of Room-Temperature Ionic Liquids with Alkyl Substituents of Different Lengths. <i>Journal of Physical Chemistry C</i> , 2008 , 112, 1132-1139	3.8	100
445	Mechanism and Determinants of Amphipathic Helix-Containing Protein Targeting to Lipid Droplets. <i>Developmental Cell</i> , 2018 , 44, 73-86.e4	10.2	98
444	Proton transport behavior through the influenza A M2 channel: insights from molecular simulation. <i>Biophysical Journal</i> , 2007 , 93, 3470-9	2.9	98
443	Systematic coarse-graining of nanoparticle interactions in molecular dynamics simulation. <i>Journal of Physical Chemistry B</i> , 2005 , 109, 17019-24	3.4	98
442	Nucleotide-dependent conformational states of actin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 12723-8	11.5	97
441	Effective force coarse-graining. <i>Physical Chemistry Chemical Physics</i> , 2009 , 11, 2002-15	3.6	97
440	Bridging microscopic and mesoscopic simulations of lipid bilayers. <i>Biophysical Journal</i> , 2002 , 83, 3357-70	2.9	96
439	Efficient, Regularized, and Scalable Algorithms for Multiscale Coarse-Graining. <i>Journal of Chemical Theory and Computation</i> , 2010 , 6, 954-65	6.4	95
438	The multiscale coarse-graining method. IV. Transferring coarse-grained potentials between temperatures. <i>Journal of Chemical Physics</i> , 2009 , 131, 024103	3.9	95
437	Hydrated excess proton at water-hydrophobic interfaces. <i>Journal of Physical Chemistry B</i> , 2009 , 113, 4017-30	3.4	95
436	Exact exchange in ab initio molecular dynamics: An efficient plane-wave based algorithm. <i>Journal of Chemical Physics</i> , 1998 , 108, 4697-4700	3.9	94
435	A Failure of Continuum Theory: Temperature Dependence of the Solvent Reorganization Energy of Electron Transfer in Highly Polar Solvents. <i>Journal of Physical Chemistry B</i> , 1999 , 103, 9130-9140	3.4	94
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433	The multiscale coarse-graining method. VI. Implementation of three-body coarse-grained potentials. <i>Journal of Chemical Physics</i> , 2010 , 132, 164107	3.9	92
432	Systematic coarse-graining of a multicomponent lipid bilayer. <i>Journal of Physical Chemistry B</i> , 2009 , 113, 1501-10	3.4	92
431	How curvature-generating proteins build scaffolds on membrane nanotubes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 11226-11231	11.5	91
430	A centroid molecular dynamics study of liquid para-hydrogen and ortho-deuterium. <i>Journal of Chemical Physics</i> , 2004 , 121, 6412-22	3.9	89

429	A Feynman path centroid dynamics approach for the computation of time correlation functions involving nonlinear operators. <i>Journal of Chemical Physics</i> , 2000 , 113, 919-929	3.9	88
428	Ab initio molecular-dynamics simulation of aqueous proton solvation and transport revisited. <i>Journal of Chemical Physics</i> , 2005 , 123, 044505	3.9	86
427	Molecular dynamics simulation of proton transport near the surface of a phospholipid membrane. <i>Biophysical Journal</i> , 2002 , 82, 1460-8	2.9	86
426	Coarse-grained simulation reveals key features of HIV-1 capsid self-assembly. <i>Nature Communications</i> , 2016 , 7, 11568	17.4	85
425	Membrane remodeling from N-BAR domain interactions: insights from multi-scale simulation. <i>Biophysical Journal</i> , 2007 , 92, 3595-602	2.9	85
424	Factors influencing local membrane curvature induction by N-BAR domains as revealed by molecular dynamics simulations. <i>Biophysical Journal</i> , 2008 , 95, 1866-76	2.9	84
423	Coarse-graining of multiprotein assemblies. <i>Current Opinion in Structural Biology</i> , 2012 , 22, 144-50	8.1	83
422	Ligand-dependent activation and deactivation of the human adenosine A(2A) receptor. <i>Journal of the American Chemical Society</i> , 2013 , 135, 8749-59	16.4	83
421	Actin filament remodeling by actin depolymerization factor/cofilin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 7299-304	11.5	81
420	Simple reversible molecular dynamics algorithms for NosFloover chain dynamics. <i>Journal of Chemical Physics</i> , 1997 , 107, 9514-9526	3.9	81
419	Modeling the free energy surfaces of electron transfer in condensed phases. <i>Journal of Chemical Physics</i> , 2000 , 113, 5413	3.9	81
418	Quantum-mechanical reaction rate constants from centroid molecular dynamics simulations. <i>Journal of Chemical Physics</i> , 2001 , 115, 9209-9222	3.9	81
417	Ab initio centroid molecular dynamics: a fully quantum method for condensed-phase dynamics simulations. <i>Chemical Physics Letters</i> , 1999 , 300, 93-98	2.5	80
416	Peptide folding using multiscale coarse-grained models. <i>Journal of Physical Chemistry B</i> , 2008 , 112, 13079-90	3.4	79
415	Propensity of Hydrated Excess Protons and Hydroxide Anions for the Air-Water Interface. <i>Journal of the American Chemical Society</i> , 2015 , 137, 12610-6	16.4	78
414	Multiscale coarse-graining of monosaccharides. <i>Journal of Physical Chemistry B</i> , 2007 , 111, 11566-75	3.4	78
413	Modeling Protein-Lipid Interactions during Viral Assembly of SARS-CoV-2. <i>Biophysical Journal</i> , 2021 , 120, 49a	2.9	78
412	Molecular Dynamics Simulations of Proton Transport in 3M and Nafion Perfluorosulfonic Acid Membranes. <i>Journal of Physical Chemistry C</i> , 2013 , 117, 8079-8091	3.8	77

411	Computer simulation of explicit proton translocation in cytochrome c oxidase: the D-pathway. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 6795-800	11.5	77
410	A unified framework for quantum activated rate processes. I. General theory. <i>Journal of Chemical Physics</i> , 1996 , 105, 6856-6870	3.9	77
409	Solvent-free lipid bilayer model using multiscale coarse-graining. <i>Journal of Physical Chemistry B</i> , 2009 , 113, 4443-55	3.4	76
408	A novel method for simulating quantum dissipative systems. <i>Journal of Chemical Physics</i> , 1996 , 104, 4189-4197	3.4	76
407	A path integral study of electronic polarization and nonlinear coupling effects in condensed phase proton transfer reactions. <i>Journal of Chemical Physics</i> , 1994 , 100, 3039-3047	3.9	76
406	Coarse-grained modeling of the self-association of therapeutic monoclonal antibodies. <i>Journal of Physical Chemistry B</i> , 2012 , 116, 8045-57	3.4	75
405	A linear-scaling self-consistent generalization of the multistate empirical valence bond method for multiple excess protons in aqueous systems. <i>Journal of Chemical Physics</i> , 2005 , 122, 144105	3.9	75
404	Mechanisms of passive ion permeation through lipid bilayers: insights from simulations. <i>Journal of Physical Chemistry B</i> , 2006 , 110, 21327-37	3.4	75
403	Structure and dynamics of the actin filament. <i>Journal of Molecular Biology</i> , 2010 , 396, 252-63	6.5	74
402	Kinetics of proton migration in liquid water. <i>Journal of Physical Chemistry B</i> , 2010 , 114, 333-9	3.4	74
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400	Ab initio molecular dynamics simulation of the Ag(111)-water interface. <i>Journal of Chemical Physics</i> , 2001 , 115, 7196-7206	3.9	74
399	Classical and Quantum Simulation of Electron Transfer Through a Polypeptide. <i>Journal of Physical Chemistry B</i> , 1999 , 103, 7367-7382	3.4	74
398	Understanding the role of amphipathic helices in N-BAR domain driven membrane remodeling. <i>Biophysical Journal</i> , 2013 , 104, 404-11	2.9	73
397	Nanostructural organization in acetonitrile/ionic liquid mixtures: molecular dynamics simulations and optical Kerr effect spectroscopy. <i>ChemPhysChem</i> , 2012 , 13, 1687-700	3.2	73
396	Ab Initio Calculations of Reactive Pathways for β -Octahydro-1,3,5,7-tetranitro-1,3,5,7-tetrazocine (β HMX). <i>Journal of Physical Chemistry A</i> , 2000 , 104, 11384-11389	2.8	73
395	Semiclassical approximations to quantum dynamical time correlation functions. <i>Journal of Chemical Physics</i> , 1996 , 104, 273-285	3.9	73
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392	On the representability problem and the physical meaning of coarse-grained models. <i>Journal of Chemical Physics</i> , 2016 , 145, 044108	3.9	72
391	A Multi-State Empirical Valence Bond Model for Weak Acid Dissociation in Aqueous Solution□ <i>Journal of Physical Chemistry A</i> , 2001 , 105, 2814-2823	2.8	71
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388	Ab initio molecular dynamics simulation of the Cu(110)□water interface. <i>Journal of Chemical Physics</i> , 2001 , 114, 3248-3257	3.9	69
387	Protein-mediated transformation of lipid vesicles into tubular networks. <i>Biophysical Journal</i> , 2013 , 105, 711-9	2.9	68
386	Role of protein interactions in defining HIV-1 viral capsid shape and stability: a coarse-grained analysis. <i>Biophysical Journal</i> , 2010 , 98, 18-26	2.9	68
385	Infrared Spectrum of the Hydrated Proton in Water. <i>Journal of Physical Chemistry Letters</i> , 2011 , 2, 81-6	6.4	68
384	New insights into BAR domain-induced membrane remodeling. <i>Biophysical Journal</i> , 2009 , 97, 1616-25	2.9	68
383	Origins of proton transport behavior from selectivity domain mutations of the aquaporin-1 channel. <i>Biophysical Journal</i> , 2006 , 90, L73-5	2.9	68
382	The role of amino acid sequence in the self-association of therapeutic monoclonal antibodies: insights from coarse-grained modeling. <i>Journal of Physical Chemistry B</i> , 2013 , 117, 1269-79	3.4	67
381	Transferable Coarse-Grained Models for Ionic Liquids. <i>Journal of Chemical Theory and Computation</i> , 2009 , 5, 1091-8	6.4	66
380	A coarse-grained model for double-helix molecules in solution: spontaneous helix formation and equilibrium properties. <i>Journal of Chemical Physics</i> , 2005 , 122, 124906	3.9	65
379	Semiclassical molecular dynamics computation of spontaneous light emission in the condensed phase: Resonance Raman spectra. <i>Journal of Chemical Physics</i> , 2001 , 114, 7130-7143	3.9	65
378	A theory for the activated barrier crossing rate constant in systems influenced by space and time dependent friction. <i>Journal of Chemical Physics</i> , 1994 , 101, 7811-7822	3.9	65
377	Dynamic force matching: A method for constructing dynamical coarse-grained models with realistic time dependence. <i>Journal of Chemical Physics</i> , 2015 , 142, 154104	3.9	64
376	Effective force field for liquid hydrogen fluoride from ab initio molecular dynamics simulation using the force-matching method. <i>Journal of Physical Chemistry B</i> , 2005 , 109, 6573-86	3.4	64

375	Efficient and Minimal Method to Bias Molecular Simulations with Experimental Data. <i>Journal of Chemical Theory and Computation</i> , 2014 , 10, 3023-30	6.4	63
374	Proton Transport Mechanism of Perfluorosulfonic Acid Membranes. <i>Journal of Physical Chemistry C</i> , 2014 , 118, 17436-17445	3.8	63
373	Multiscale computer simulation of the immature HIV-1 virion. <i>Biophysical Journal</i> , 2010 , 99, 2757-65	2.9	63
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371	Storage of an excess proton in the hydrogen-bonded network of the d-pathway of cytochrome C oxidase: identification of a protonated water cluster. <i>Journal of the American Chemical Society</i> , 2007 , 129, 2910-3	16.4	63
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