

Jaydeep P Bardhan

List of Publications by Year in descending order

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35
papers

624
citations

706676

14
h-index

651938

25
g-index

35
all docs

35
docs citations

35
times ranked

544
citing authors

#	ARTICLE	IF	CITATIONS
1	Guest Editor's Introduction: DOE Computational Science Graduate Fellowship Research Showcase. Computing in Science and Engineering, 2021, 23, 5-8.	1.2	1
2	High-performance transformation of protein structure representation from internal to Cartesian coordinates. Journal of Computational Chemistry, 2020, 41, 2104-2114.	1.5	3
3	Solvation thermodynamics of neutral and charged solutes using the solvation-layer interface condition continuum dielectric model. International Journal of Quantum Chemistry, 2019, 119, e25771.	1.0	5
4	Predicting solvation free energies and thermodynamics in polar solvents and mixtures using a solvation-layer interface condition. Journal of Chemical Physics, 2017, 146, .	1.2	14
5	Extending the Solvation-Layer Interface Condition Continuum Electrostatic Model to a Linearized Poisson-Boltzmann Solvent. Journal of Chemical Theory and Computation, 2017, 13, 2897-2914.	2.3	7
6	Effects of Catalytic Action and Ligand Binding on Conformational Ensembles of Adenylate Kinase. Biochemistry, 2017, 56, 4559-4567.	1.2	13
7	Theory of wavelet-based coarse-graining hierarchies for molecular dynamics. Physical Review E, 2017, 96, 013301.	0.8	1
8	Comparative Assessment of Nonlocal Continuum Solvent Models Exhibiting Overscreening. Computational and Mathematical Biophysics, 2017, 5, 40-57.	0.6	2
9	A framework to optimize protein structure from solution scattering using ADMM and an elastic subdomain network. , 2016, , .		1
10	Generalising the mean spherical approximation as a multiscale, nonlinear boundary condition at the solute-solvent interface. Molecular Physics, 2016, 114, 2558-2567.	0.8	6
11	Nonlocal Electrostatics in Spherical Geometries Using Eigenfunction Expansions of Boundary-Integral Operators. Computational and Mathematical Biophysics, 2015, 3, 1-22.	0.6	2
12	Constrained Maximum Likelihood Estimation of Relative Abundances of Protein Conformation in a Heterogeneous Mixture From Small Angle X-Ray Scattering Intensity Measurements. IEEE Transactions on Signal Processing, 2015, 63, 5383-5394.	3.2	10
13	Diffusion Wavelet Decomposition for Coarse-Graining of Polymer Chains. Materials Research Society Symposia Proceedings, 2015, 1753, 24.	0.1	0
14	Accelerating protein coordinate conversion using GPUs. , 2014, , .		1
15	Communication: Modeling charge-sign asymmetric solvation free energies with nonlinear boundary conditions. Journal of Chemical Physics, 2014, 141, 131103.	1.2	23
16	A biomolecular electrostatics solver using Python, GPUs and boundary elements that can handle solvent-filled cavities and Stern layers. Computer Physics Communications, 2014, 185, 720-729.	3.0	33
17	Gradient models in molecular biophysics: progress, challenges, opportunities. Journal of the Mechanical Behavior of Materials, 2013, 22, 169-184.	0.7	11
18	Analysis of fast boundary-integral approximations for modeling electrostatic contributions of molecular binding. Computational and Mathematical Biophysics, 2013, 1, 124-150.	0.6	2

#	ARTICLE	IF	CITATIONS
19	Computational science and re-discovery: open-source implementation of ellipsoidal harmonics for problems in potential theory. <i>Computational Science & Discovery</i> , 2012, 5, 014006.	1.5	7
20	Comparison of three-dimensional Poisson solution methods for particle-based simulation and inhomogeneous dielectrics. <i>Physical Review E</i> , 2012, 86, 011912.	0.8	22
21	Biomolecular electrostaticsâ€”I want your solvation (model). <i>Computational Science & Discovery</i> , 2012, 5, 013001.	1.5	30
22	Affine-response model of molecular solvation of ions: Accurate predictions of asymmetric charging free energies. <i>Journal of Chemical Physics</i> , 2012, 137, 124101.	1.2	40
23	https://doi.org/10.1109/MDT.2012.2213176 IEEE Design and Test of	1.4	0
24	Biomolecular electrostatics using a fast multipole BEM on up to 512 gpus and a billion unknowns. <i>Computer Physics Communications</i> , 2011, 182, 1272-1283.	3.0	73
25	Efficiently accounting for ion correlations in electrokinetic nanofluidic devices using density functional theory. <i>Journal of Colloid and Interface Science</i> , 2011, 359, 520-529.	5.0	45
26	Nonlocal continuum electrostatic theory predicts surprisingly small energetic penalties for charge burial in proteins. <i>Journal of Chemical Physics</i> , 2011, 135, 104113.	1.2	23
27	Mathematical analysis of the boundary-integral based electrostatics estimation approximation for molecular solvation: Exact results for spherical inclusions. <i>Journal of Chemical Physics</i> , 2011, 135, 124107.	1.2	10
28	Discretization of the induced-charge boundary integral equation. <i>Physical Review E</i> , 2009, 80, 011906.	0.8	24
29	Bounding the electrostatic free energies associated with linear continuum models of molecular solvation. <i>Journal of Chemical Physics</i> , 2009, 130, 104108.	1.2	14
30	Accurate solution of multi-region continuum biomolecule electrostatic problems using the linearized Poissonâ€”Boltzmann equation with curved boundary elements. <i>Journal of Computational Chemistry</i> , 2009, 30, 132-153.	1.5	80
31	â€œReverse-Schurâ€ Approach to Optimization with Linear PDE Constraints: Application to Biomolecule Analysis and Design. <i>Journal of Chemical Theory and Computation</i> , 2009, 5, 3260-3278.	2.3	8
32	Numerical solution of boundary-integral equations for molecular electrostatics. <i>Journal of Chemical Physics</i> , 2009, 130, 094102.	1.2	41
33	Interpreting the Coulomb-field approximation for generalized-Born electrostatics using boundary-integral equation theory. <i>Journal of Chemical Physics</i> , 2008, 129, 144105.	1.2	27
34	Numerical integration techniques for curved-element discretizations of molecule-solvent interfaces. <i>Journal of Chemical Physics</i> , 2007, 127, 014701.	1.2	33
35	An Accurate Surface Formulation for Biomolecule Electrostatics in Non-Ionic Solutions. , 2005, 2005, 7591-5.		12