

Christopher M Baker

List of Publications by Year in descending order

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16
papers

1,390
citations

687363

13
h-index

940533

16
g-index

16
all docs

16
docs citations

16
times ranked

1967
citing authors

#	ARTICLE	IF	CITATIONS
1	Tautomer Standardization in Chemical Databases: Deriving Business Rules from Quantum Chemistry. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 3781-3791.	5.4	5
2	Polarizable force fields for molecular dynamics simulations of biomolecules. <i>Wiley Interdisciplinary Reviews: Computational Molecular Science</i> , 2015, 5, 241-254.	14.6	139
3	Insights into the binding of intrinsically disordered proteins from molecular dynamics simulation. <i>Wiley Interdisciplinary Reviews: Computational Molecular Science</i> , 2014, 4, 182-198.	14.6	56
4	Interplay between partner and ligand facilitates the folding and binding of an intrinsically disordered protein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 15420-15425.	7.1	144
5	Matching of Additive and Polarizable Force Fields for Multiscale Condensed Phase Simulations. <i>Journal of Chemical Theory and Computation</i> , 2013, 9, 2826-2837.	5.3	24
6	Optimization of the CHARMM Additive Force Field for DNA: Improved Treatment of the BI/BII Conformational Equilibrium. <i>Journal of Chemical Theory and Computation</i> , 2012, 8, 348-362.	5.3	464
7	Development of CHARMM Polarizable Force Field for Nucleic Acid Bases Based on the Classical Drude Oscillator Model. <i>Journal of Physical Chemistry B</i> , 2011, 115, 580-596.	2.6	121
8	Polarizability rescaling and atom-based Thole scaling in the CHARMM Drude polarizable force field for ethers. <i>Journal of Molecular Modeling</i> , 2010, 16, 567-576.	1.8	62
9	Accurate Calculation of Hydration Free Energies using Pair-Specific Lennard-Jones Parameters in the CHARMM Drude Polarizable Force Field. <i>Journal of Chemical Theory and Computation</i> , 2010, 6, 1181-1198.	5.3	131
10	The Effect of Solvation on Biomolecular Conformation: α -2-Amino-1-phenylethanol. <i>Journal of Physical Chemistry B</i> , 2007, 111, 9940-9954.	2.6	12
11	Modeling Aromatic Liquids: α -Toluene, Phenol, and Pyridine. <i>Journal of Chemical Theory and Computation</i> , 2007, 3, 530-548.	5.3	38
12	Role of aromatic amino acids in protein-nucleic acid recognition. <i>Biopolymers</i> , 2007, 85, 456-470.	2.4	90
13	The Structure of Liquid Benzene. <i>Journal of Chemical Theory and Computation</i> , 2006, 2, 947-955.	5.3	36
14	A solvent induced mechanism for conformational change. <i>Chemical Communications</i> , 2006, , 1387.	4.1	10
15	Structure, electronic circular dichroism and Raman optical activity in the gas phase and in solution: a computational and experimental investigation. <i>Physical Chemistry Chemical Physics</i> , 2005, 7, 1432.	2.8	39
16	Electronic Circular Dichroism Spectroscopy of 1-(R)-Phenylethanol: The α -Sector Rule Revisited and an Exploration of Solvent Effects. <i>Israel Journal of Chemistry</i> , 2004, 44, 27-36.	2.3	19