

Daniel Trzesniak

List of Publications by Year in descending order

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14
papers

1,820
citations

758635

12
h-index

1058022

14
g-index

15
all docs

15
docs citations

15
times ranked

2130
citing authors

#	ARTICLE	IF	CITATIONS
1	On using a too large integration time step in molecular dynamics simulations of coarse-grained molecular models. <i>Physical Chemistry Chemical Physics</i> , 2009, 11, 1934-41.	1.3	76
2	A Comparison of Methods to Compute the Potential of Mean Force. <i>ChemPhysChem</i> , 2007, 8, 162-169.	1.0	243
3	Comparison of Thermodynamic Properties of Coarse-Grained and Atomic-Level Simulation Models. <i>ChemPhysChem</i> , 2007, 8, 452-461.	1.0	102
4	Enthalpy-Entropy Compensation in the Effects of Urea on Hydrophobic Interactions. <i>Journal of Physical Chemistry B</i> , 2006, 110, 12852-12855.	1.2	32
5	Protein under pressure: Molecular dynamics simulation of the arc repressor. <i>Proteins: Structure, Function and Bioinformatics</i> , 2006, 65, 136-144.	1.5	31
6	Catalytic mechanism of cyclophilin as observed in molecular dynamics simulations: Pathway prediction and reconciliation of X-ray crystallographic and NMR solution data. <i>Protein Science</i> , 2006, 15, 2544-2551.	3.1	29
7	Pathway dependence of the efficiency of calculating free energy and entropy of solute-solute association in water. <i>Chemical Physics</i> , 2006, 330, 410-416.	0.9	4
8	Simulation of an all- ^{13}C -icosapeptide containing the 20 proteinogenic side chains: Effect of temperature, pH, counterions, solvent, and force field on helix stability. <i>Biopolymers</i> , 2006, 83, 636-645.	1.2	5
9	Biomolecular Modeling: Goals, Problems, Perspectives. <i>Angewandte Chemie - International Edition</i> , 2006, 45, 4064-4092.	7.2	503
10	The GROMOS software for biomolecular simulation: GROMOS05. <i>Journal of Computational Chemistry</i> , 2005, 26, 1719-1751.	1.5	592
11	Energy-Entropy Compensation in the Transfer of Nonpolar Solutes from Water to Cosolvent/Water Mixtures. <i>ChemPhysChem</i> , 2005, 6, 1010-1010.	1.0	0
12	Interpreting NMR Data for ^{15}N -Peptides Using Molecular Dynamics Simulations. <i>Journal of the American Chemical Society</i> , 2005, 127, 14320-14329.	6.6	39
13	Energy-Entropy Compensation in the Transfer of Nonpolar Solutes from Water to Cosolvent/Water Mixtures. <i>ChemPhysChem</i> , 2004, 5, 144-147.	1.0	49
14	Computer simulation studies on the solvation of aliphatic hydrocarbons in 6.9 M aqueous urea solution. <i>Physical Chemistry Chemical Physics</i> , 2004, 6, 697.	1.3	84