

Michael R Jones

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2654219/publications.pdf>

Version: 2024-02-01

11
papers

137
citations

1307366

7
h-index

1372474

10
g-index

11
all docs

11
docs citations

11
times ranked

168
citing authors

#	ARTICLE	IF	CITATIONS
1	Obtaining QM/MM binding free energies in the SAMPL8 drugs of abuse challenge: indirect approaches. <i>Journal of Computer-Aided Molecular Design</i> , 2022, 36, 263-277.	1.3	4
2	A replica exchange umbrella sampling (REUS) approach to predict host-guest binding free energies in SAMPL8 challenge. <i>Journal of Computer-Aided Molecular Design</i> , 2021, 35, 667-677.	1.3	5
3	Quantum chemical predictions of water-octanol partition coefficients applied to the SAMPL6 logP blind challenge. <i>Journal of Computer-Aided Molecular Design</i> , 2020, 34, 485-493.	1.3	13
4	Multi-phase Boltzmann weighting: accounting for local inhomogeneity in molecular simulations of water-octanol partition coefficients in the SAMPL6 challenge. <i>Journal of Computer-Aided Molecular Design</i> , 2020, 34, 471-483.	1.3	7
5	SAMPL6 logP challenge: machine learning and quantum mechanical approaches. <i>Journal of Computer-Aided Molecular Design</i> , 2020, 34, 495-510.	1.3	11
6	Impact of intracellular ionic strength on dimer binding in the NF- κ B Inducing kinase. <i>Journal of Structural Biology</i> , 2018, 202, 183-190.	1.3	0
7	SAMPL6 host-guest challenge: binding free energies via a multistep approach. <i>Journal of Computer-Aided Molecular Design</i> , 2018, 32, 1097-1115.	1.3	16
8	Prediction of CB[8] host-guest binding free energies in SAMPL6 using the double-decoupling method. <i>Journal of Computer-Aided Molecular Design</i> , 2018, 32, 1059-1073.	1.3	13
9	Absolute and relative pKa predictions via a DFT approach applied to the SAMPL6 blind challenge. <i>Journal of Computer-Aided Molecular Design</i> , 2018, 32, 1179-1189.	1.3	25
10	Partition coefficients for the SAMPL5 challenge using transfer free energies. <i>Journal of Computer-Aided Molecular Design</i> , 2016, 30, 1129-1138.	1.3	19
11	Calculating distribution coefficients based on multi-scale free energy simulations: an evaluation of MM and QM/MM explicit solvent simulations of water-cyclohexane transfer in the SAMPL5 challenge. <i>Journal of Computer-Aided Molecular Design</i> , 2016, 30, 989-1006.	1.3	24