Sergei V Krivov

List of Publications by Citations

Source: https://exaly.com/author-pdf/4067945/sergei-v-krivov-publications-by-citations.pdf

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

21 830 12 23 g-index

23 900 5.6 4.82 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
21	Hidden complexity of free energy surfaces for peptide (protein) folding. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 14766-70	11.5	328
20	Diffusive reaction dynamics on invariant free energy profiles. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 13841-6	11.5	102
19	One-dimensional barrier-preserving free-energy projections of a beta-sheet miniprotein: new insights into the folding process. <i>Journal of Physical Chemistry B</i> , 2008 , 112, 8701-14	3.4	75
18	The free energy landscape analysis of protein (FIP35) folding dynamics. <i>Journal of Physical Chemistry B</i> , 2011 , 115, 12315-24	3.4	48
17	On Reaction Coordinate Optimality. Journal of Chemical Theory and Computation, 2013, 9, 135-46	6.4	43
16	Is protein folding sub-diffusive?. PLoS Computational Biology, 2010, 6, e1000921	5	41
15	Potential energy surfaces and conformational transitions in biomolecules: a successive confinement approach applied to a solvated tetrapeptide. <i>Physical Review Letters</i> , 2002 , 88, 038101	7.4	31
14	Numerical construction of the p(fold) (committor) reaction coordinate for a Markov process. Journal of Physical Chemistry B, 2011 , 115, 11382-8	3.4	27
13	Optimal reaction coordinates. <i>Wiley Interdisciplinary Reviews: Computational Molecular Science</i> , 2016 , 6, 748-763	7.9	21
12	Nonparametric variational optimization of reaction coordinates. <i>Journal of Chemical Physics</i> , 2015 , 143, 184108	3.9	18
11	Exploratory search during directed navigation in and larva. ELife, 2017, 6,	8.9	16
10	High-Resolution Free-Energy Landscape Analysis of Ehelical Protein Folding: HP35 and Its Double Mutant. <i>Journal of Chemical Theory and Computation</i> , 2013 , 9, 5257-5266	6.4	13
9	Protein Folding Free Energy Landscape along the Committor - the Optimal Folding Coordinate. <i>Journal of Chemical Theory and Computation</i> , 2018 , 14, 3418-3427	6.4	12
8	Robust Estimation of Diffusion-Optimized Ensembles for Enhanced Sampling. <i>Journal of Chemical Theory and Computation</i> , 2014 , 10, 543-53	6.4	11
7	Optimal dimensionality reduction of complex dynamics: the chess game as diffusion on a free-energy landscape. <i>Physical Review E</i> , 2011 , 84, 011135	2.4	11
6	Fep1d: a script for the analysis of reaction coordinates. <i>Journal of Computational Chemistry</i> , 2015 , 36, 878-82	3.5	8
5	Optimal reaction coordinate as a biomarker for the dynamics of recovery from kidney transplant. <i>PLoS Computational Biology</i> , 2014 , 10, e1003685	5	8

LIST OF PUBLICATIONS

4	Method to describe stochastic dynamics using an optimal coordinate. <i>Physical Review E</i> , 2013 , 88, 06213 1 .	·4	6
3	High-resolution free energy landscape analysis of protein folding. <i>Biochemical Society Transactions</i> , 2015 , 43, 157-61	.1	4
2	Blind Analysis of Molecular Dynamics. <i>Journal of Chemical Theory and Computation</i> , 2021 , 17, 2725-2736 6.	-4	4
1	Nonparametric Analysis of Nonequilibrium Simulations. <i>Journal of Chemical Theory and Computation</i> , 2021 , 17, 5466-5481	·4	3