Alexey Aleksandrov

List of Publications by Citations

Source: https://exaly.com/author-pdf/6295684/alexey-aleksandrov-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.

26 426 13 20 h-index g-index citations papers 6.6 3.86 27 535 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
26	Molecular dynamics simulations show that conformational selection governs the binding preferences of imatinib for several tyrosine kinases. <i>Journal of Biological Chemistry</i> , 2010 , 285, 13807-	15 ^{5.4}	65
25	Alchemical free energy simulations for biological complexes: powerful but temperamental. <i>Journal of Molecular Recognition</i> , 2010 , 23, 117-27	2.6	46
24	Additive CHARMM force field for naturally occurring modified ribonucleotides. <i>Journal of Computational Chemistry</i> , 2016 , 37, 896-912	3.5	36
23	Protonation patterns in tetracycline:tet repressor recognition: simulations and experiments. <i>ChemBioChem</i> , 2007 , 8, 675-85	3.8	32
22	Tet repressor induction by tetracycline: a molecular dynamics, continuum electrostatics, and crystallographic study. <i>Journal of Molecular Biology</i> , 2008 , 378, 898-912	6.5	29
21	Predicting the acid/base behavior of proteins: a constant-pH Monte Carlo approach with generalized born solvent. <i>Journal of Physical Chemistry B</i> , 2010 , 114, 10634-48	3.4	28
20	Mechanism and dynamics of fatty acid photodecarboxylase. <i>Science</i> , 2021 , 372,	33.3	28
19	Molecular dynamics simulations of the 30S ribosomal subunit reveal a preferred tetracycline binding site. <i>Journal of the American Chemical Society</i> , 2008 , 130, 1114-5	16.4	24
18	Mechanism of activation of elongation factor Tu by ribosome: catalytic histidine activates GTP by protonation. <i>Rna</i> , 2013 , 19, 1218-25	5.8	18
17	Identification of a second GTP-bound magnesium ion in archaeal initiation factor 2. <i>Nucleic Acids Research</i> , 2015 , 43, 2946-57	20.1	18
16	A hybrid elastic band string algorithm for studies of enzymatic reactions. <i>Physical Chemistry Chemical Physics</i> , 2012 , 14, 12544-53	3.6	18
15	Combining the polarizable Drude force field with a continuum electrostatic Poisson-Boltzmann implicit solvation model. <i>Journal of Computational Chemistry</i> , 2018 , 39, 1707-1719	3.5	13
14	Tetracycline-tet repressor binding specificity: insights from experiments and simulations. <i>Biophysical Journal</i> , 2009 , 97, 2829-38	2.9	13
13	The mechanism of citryl-coenzyme A formation catalyzed by citrate synthase. <i>Journal of Physical Chemistry B</i> , 2014 , 118, 4505-13	3.4	11
12	A Molecular Mechanics Model for Flavins. <i>Journal of Computational Chemistry</i> , 2019 , 40, 2834-2842	3.5	9
11	p Calculations with the Polarizable Drude Force Field and Poisson-Boltzmann Solvation Model. <i>Journal of Chemical Theory and Computation</i> , 2020 , 16, 4655-4668	6.4	7
10	Cyclization Reaction Catalyzed by Cyclodipeptide Synthases Relies on a Conserved Tyrosine Residue. <i>Scientific Reports</i> , 2018 , 8, 7031	4.9	6

LIST OF PUBLICATIONS

9	Advances and challenges in drug design against tuberculosis: application of in silico approaches. <i>Expert Opinion on Drug Discovery</i> , 2019 , 14, 35-46	6.2	6
8	Additive CHARMM36 Force Field for Nonstandard Amino Acids. <i>Journal of Chemical Theory and Computation</i> , 2021 , 17, 3554-3570	6.4	5
7	Aminoacetylation Reaction Catalyzed by Leucyl-tRNA Synthetase Operates via a Self-Assisted Mechanism Using a Conserved Residue and the Aminoacyl Substrate. <i>Journal of Physical Chemistry B</i> , 2016 , 120, 4388-98	3.4	5
6	Electrostatic free energies in translational GTPases: Classic allostery and the rest. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2015 , 1850, 1006-1016	4	4
5	Characterization of Light-Induced, Short-Lived Interacting Radicals in the Active Site of Flavoprotein Ferredoxin-NADP Oxidoreductase. <i>Journal of the American Chemical Society</i> , 2021 , 143, 2757-2768	16.4	2
4	An atomistic model for simulations of nilotinib and nilotinib/kinase binding. <i>Theoretical Chemistry Accounts</i> , 2011 , 129, 747-756	1.9	1
3	Mechanism of Naphthoquinone Selectivity of Thymidylate Synthase ThyX. <i>Biophysical Journal</i> , 2020 , 119, 2508-2516	2.9	1
2	Photochemical processes in flavo-enzymes as a probe for active site dynamics: TrmFO of Thermus thermophilus. <i>Photochemical and Photobiological Sciences</i> , 2021 , 20, 663-670	4.2	0
1	Cyclodipeptide Synthases of the NYH Subfamily Recognize tRNA Using an EHelix Enriched with Positive Residues. <i>Biochemistry</i> , 2021 , 60, 64-76	3.2	