

Geraldo Rodrigues Sartori

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

90
citations

1937685

4
h-index

1372567

10
g-index

14
all docs

14
docs citations

14
times ranked

209
citing authors

#	ARTICLE	IF	CITATIONS
1	Hydrogen Bond Basicity Prediction for Medicinal Chemistry Design. <i>Journal of Medicinal Chemistry</i> , 2016, 59, 4278-4288.	6.4	48
2	Small-angle X-ray scattering and structural modeling of full-length: cellobiohydrolase I from <i>Trichoderma harzianum</i> . <i>Cellulose</i> , 2013, 20, 1573-1585.	4.9	10
3	Integration of methods in cheminformatics and biocalorimetry for the design of trypanosomatid enzyme inhibitors. <i>Future Medicinal Chemistry</i> , 2014, 6, 17-33.	2.3	10
4	Ligand-induced conformational selection predicts the selectivity of cysteine protease inhibitors. <i>PLoS ONE</i> , 2019, 14, e0222055.	2.5	6
5	Highly predictive hologram QSAR models of nitrile-containing cruzain inhibitors. <i>Journal of Biomolecular Structure and Dynamics</i> , 2017, 35, 3232-3249.	3.5	4
6	Development of New Potential Inhibitors of α 1 Integrins through In Silico Methods—Screening and Computational Validation. <i>Life</i> , 2022, 12, 932.	2.4	4
7	Polymorphisms in plastoquinol oxidase (PTOX) from <i>Arabidopsis</i> accessions indicate SNP-induced structural variants associated with altitude and rainfall. <i>Journal of Bioenergetics and Biomembranes</i> , 2019, 51, 151-164.	2.3	3
8	Computationally-obtained structural insights into the molecular interactions between Pidilizumab and binding partners DLL1 and PD-1. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, , 1-13.	3.5	2
9	Investigation of Unprecedented Sites and Proposition of New Ligands for Programmed Cell Death Protein I through Molecular Dynamics with Probes and Virtual Screening. <i>Journal of Chemical Information and Modeling</i> , 2022, 62, 1236-1248.	5.4	2
10	Guidelines To Predict Binding Poses of Antibody—Integrin Complexes. <i>ACS Omega</i> , 2020, 5, 16379-16385.	3.5	1
11	Ligand-induced conformational selection predicts the selectivity of cysteine protease inhibitors. , 2019, 14, e0222055.		0
12	Ligand-induced conformational selection predicts the selectivity of cysteine protease inhibitors. , 2019, 14, e0222055.		0
13	Ligand-induced conformational selection predicts the selectivity of cysteine protease inhibitors. , 2019, 14, e0222055.		0
14	Ligand-induced conformational selection predicts the selectivity of cysteine protease inhibitors. , 2019, 14, e0222055.		0