

# Alberto M Dos Santos

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5761808/publications.pdf>

Version: 2024-02-01

8  
papers

127  
citations

1163117

8  
h-index

1588992

8  
g-index

8  
all docs

8  
docs citations

8  
times ranked

202  
citing authors

#	ARTICLE	IF	CITATIONS
1	Experimental study and computational modelling of cruzain cysteine protease inhibition by dipeptidyl nitriles. <i>Physical Chemistry Chemical Physics</i> , 2018, 20, 24317-24328.	2.8	38
2	Evaluating QM/MM Free Energy Surfaces for Ranking Cysteine Protease Covalent Inhibitors. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 880-889.	5.4	19
3	Structural Analysis of Viral Infectivity Factor of HIV Type 1 and Its Interaction with A3G, EloC and EloB. <i>PLoS ONE</i> , 2014, 9, e89116.	2.5	18
4	Computational study of conformational changes in human 3-hydroxy-3-methylglutaryl coenzyme reductase induced by substrate binding. <i>Journal of Biomolecular Structure and Dynamics</i> , 2019, 37, 4374-4383.	3.5	13
5	Computed insight into a peptide inhibitor preventing the induced fit mechanism of MurA enzyme from <i>Pseudomonas aeruginosa</i> . <i>Chemical Biology and Drug Design</i> , 2017, 89, 599-607.	3.2	10
6	Unraveling the conformational dynamics of glycerol 3-phosphate dehydrogenase, a nicotinamide adenine dinucleotide-dependent enzyme of <i>Leishmania mexicana</i> . <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 2044-2055.	3.5	10
7	Targeting Peptidyl-prolyl Cis-trans Isomerase NIMA-interacting 1: A Structure-based Virtual Screening Approach to Find Novel Inhibitors. <i>Current Computer-Aided Drug Design</i> , 2020, 16, 605-617.	1.2	10
8	Unraveling the Addition–Elimination Mechanism of EPSP Synthase through Computer Modeling. <i>Journal of Physical Chemistry B</i> , 2017, 121, 8626-8637.	2.6	9