

Andrea Giachetti

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3699141/publications.pdf>

Version: 2024-02-01

12
papers

792
citations

1040056

9
h-index

1281871

11
g-index

12
all docs

12
docs citations

12
times ranked

1177
citing authors

#	ARTICLE	IF	CITATIONS
1	Structural Biology in the Clouds: The WeNMR-EOSC Ecosystem. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 729513.	3.5	308
2	WeNMR: Structural Biology on the Grid. <i>Journal of Grid Computing</i> , 2012, 10, 743-767.	3.9	170
3	Paramagnetism-Based Restraints for Xplor-NIH. <i>Journal of Biomolecular NMR</i> , 2004, 28, 249-261.	2.8	119
4	Blind Testing of Routine, Fully Automated Determination of Protein Structures from NMR Data. <i>Structure</i> , 2012, 20, 227-236.	3.3	75
5	A Grid-enabled web portal for NMR structure refinement with AMBER. <i>Bioinformatics</i> , 2011, 27, 2384-2390.	4.1	55
6	Investigation of the Iron(II) Release Mechanism of Human H-Ferritin as a Function of pH. <i>Journal of Chemical Information and Modeling</i> , 2017, 57, 2112-2118.	5.4	22
7	An atomistic view of the YiiP structural changes upon zinc(II) binding. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2019, 1863, 1560-1567.	2.4	13
8	A protocol for the refinement of NMR structures using simultaneously pseudocontact shift restraints from multiple lanthanide ions. <i>Journal of Biomolecular NMR</i> , 2016, 66, 175-185.	2.8	10
9	Insights into the Dynamics of the Human Zinc Transporter ZnT8 by MD Simulations. <i>Journal of Chemical Information and Modeling</i> , 2021, 61, 901-912.	5.4	10
10	Molecular dynamics simulations of metalloproteins: A folding study of rubredoxin from <i>Pyrococcus furiosus</i> . <i>AIMS Biophysics</i> , 2018, 5, 77-96.	0.6	5
11	EGI federated platforms supporting accelerated computing. , 2017, , .		3
12	West-Life: A Virtual Research Environment for structural biology. <i>Journal of Structural Biology: X</i> , 2019, 1, 100006.	1.3	2