## Edoardo Sarti

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8338133/publications.pdf

Version: 2024-02-01

11 papers	101 citations	1478505 6 h-index	9 g-index
16	16	16	139
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	AlignMe: an update of the web server for alignment of membrane protein sequences. Nucleic Acids Research, 2022, 50, W29-W35.	14.5	1
2	Refining pairwise sequence alignments of membrane proteins by the incorporation of anchors. PLoS ONE, 2021, 16, e0239881.	2.5	2
3	MemSTATS: A Benchmark Set of Membrane Protein Symmetries and Pseudosymmetries. Journal of Molecular Biology, 2020, 432, 597-604.	4.2	12
4	FilterDCA: Interpretable supervised contact prediction using inter-domain coevolution. PLoS Computational Biology, 2020, 16, e1007621.	3.2	8
5	EncoMPASS: an online database for analyzing structure and symmetry in membrane proteins. Nucleic Acids Research, 2019, 47, D315-D321.	14.5	17
6	Analyzing the Structure and Symmetry of Membrane Proteins through the Systematic Online Database EncoMPASS. Biophysical Journal, 2018, 114, 190a.	0.5	0
7	Toward a unified scoring function for native state discrimination and drug-binding pocket recognition. Physical Chemistry Chemical Physics, 2018, 20, 17148-17155.	2.8	5
8	Protein-protein structure prediction by scoring molecular dynamics trajectories of putative poses. Proteins: Structure, Function and Bioinformatics, 2016, 84, 1312-1320.	2.6	11
9	Native fold and docking pose discrimination by the same residueâ€based scoring function. Proteins: Structure, Function and Bioinformatics, 2015, 83, 621-630.	2.6	17
10	BACHSCORE. A tool for evaluating efficiently and reliably the quality of large sets of protein structures. Computer Physics Communications, 2013, 184, 2860-2865.	7.5	20
11	Photoluminescence Study of Low Thermal Budget Ill–V Nanostructures on Silicon by Droplet Epitaxy. Nanoscale Research Letters, 2010, 5, 1650-1653.	5.7	6