

Fanchi Meng

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10437635/publications.pdf>

Version: 2024-02-01

10
papers

525
citations

1039880

9
h-index

1372474

10
g-index

10
all docs

10
docs citations

10
times ranked

518
citing authors

#	ARTICLE	IF	CITATIONS
1	Functional and structural characterization of osteocytic MLO-Y4 cell proteins encoded by genes differentially expressed in response to mechanical signals in vitro. <i>Scientific Reports</i> , 2018, 8, 6716.	1.6	11
2	High-throughput prediction of disordered moonlighting regions in protein sequences. <i>Proteins: Structure, Function and Bioinformatics</i> , 2018, 86, 1097-1110.	1.5	29
3	Comprehensive review of methods for prediction of intrinsic disorder and its molecular functions. <i>Cellular and Molecular Life Sciences</i> , 2017, 74, 3069-3090.	2.4	153
4	Computational Prediction of Intrinsic Disorder in Proteins. <i>Current Protocols in Protein Science</i> , 2017, 88, 2.16.1-2.16.14.	2.8	49
5	fDETECT webserver: fast predictor of propensity for protein production, purification, and crystallization. <i>BMC Bioinformatics</i> , 2017, 18, 580.	1.2	22
6	Compartmentalization and Functionality of Nuclear Disorder: Intrinsic Disorder and Protein-Protein Interactions in Intra-Nuclear Compartments. <i>International Journal of Molecular Sciences</i> , 2016, 17, 24.	1.8	94
7	Autophagy-related intrinsically disordered proteins in intra-nuclear compartments. <i>Molecular BioSystems</i> , 2016, 12, 2798-2817.	2.9	27
8	DFLpred: High-throughput prediction of disordered flexible linker regions in protein sequences. <i>Bioinformatics</i> , 2016, 32, i341-i350.	1.8	72
9	Computational Prediction of Protein Secondary Structure from Sequence. <i>Current Protocols in Protein Science</i> , 2016, 86, 2.3.1-2.3.10.	2.8	10
10	Unstructural biology of the dengue virus proteins. <i>FEBS Journal</i> , 2015, 282, 3368-3394.	2.2	58