

Sandip Chakraborty

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

11
papers

87
citations

6
h-index

9
g-index

11
ext. papers

109
ext. citations

3.7
avg, IF

2.66
L-index

#	Paper	IF	Citations
11	Position Matters: Network Centrality Considerably Impacts Rates of Protein Evolution in the Human Protein-Protein Interaction Network. <i>Genome Biology and Evolution</i> , 2017 , 9, 1742-1756	3.9	24
10	Positive Selection and Centrality in the Yeast and Fly Protein-Protein Interaction Networks. <i>BioMed Research International</i> , 2016 , 2016, 4658506	3	11
9	Protein complex forming ability is favored over the features of interacting partners in determining the evolutionary rates of proteins in the yeast protein-protein interaction networks. <i>BMC Systems Biology</i> , 2010 , 4, 155	3.5	10
8	Evolutionary rate heterogeneity of core and attachment proteins in yeast protein complexes. <i>Genome Biology and Evolution</i> , 2013 , 5, 1366-75	3.9	9
7	Deciphering the cause of evolutionary variance within intrinsically disordered regions in human proteins. <i>Journal of Biomolecular Structure and Dynamics</i> , 2017 , 35, 233-249	3.6	7
6	Complex-forming proteins escape the robust regulations of miRNA in human. <i>FEBS Letters</i> , 2013 , 587, 2284-7	3.8	7
5	Protein intrinsic disorder negatively associates with gene age in different eukaryotic lineages. <i>Molecular BioSystems</i> , 2017 , 13, 2044-2055		6
4	Exploring the evolutionary rate differences between human disease and non-disease genes. <i>Genomics</i> , 2016 , 108, 18-24	4.3	5
3	GC-made protein disorder sheds new light on vertebrate evolution. <i>Genomics</i> , 2014 , 104, 530-7	4.3	4
2	Chaperone client proteins evolve slower than non-client proteins. <i>Functional and Integrative Genomics</i> , 2020 , 20, 621-631	3.8	2
1	Insights into Eukaryotic Interacting Protein Evolution 2011 , 51-70		2