

# Sandip Chakraborty

## List of Publications by Year in Descending Order

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