Sara Light

List of Publications by Year in descending order

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687363 1058476 1,078 14 13 14 citations h-index g-index papers 16 16 16 1631 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	What properties characterize the hub proteins of the protein-protein interaction network of Saccharomyces cerevisiae?. Genome Biology, 2006, 7, R45.	9.6	337
2	Domain Rearrangements in Protein Evolution. Journal of Molecular Biology, 2005, 353, 911-923.	4.2	190
3	Evolution after gene duplication: models, mechanisms, sequences, systems, and organisms. Journal of Experimental Zoology Part B: Molecular and Developmental Evolution, 2007, 308B, 58-73.	1.3	148
4	Protein Expansion Is Primarily due to Indels in Intrinsically Disordered Regions. Molecular Biology and Evolution, 2013, 30, 2645-2653.	8.9	75
5	Preferential attachment in the evolution of metabolic networks. BMC Genomics, 2005, 6, 159.	2.8	60
6	Network analysis of metabolic enzyme evolution in Escherichia coli. BMC Bioinformatics, 2004, 5, 15.	2.6	50
7	High GC content causes orphan proteins to be intrinsically disordered. PLoS Computational Biology, 2017, 13, e1005375.	3. 2	50
8	Nebulin: A Study of Protein Repeat Evolution. Journal of Molecular Biology, 2010, 402, 38-51.	4.2	47
9	Long indels are disordered: A study of disorder and indels in homologous eukaryotic proteins. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2013, 1834, 890-897.	2.3	30
10	Orphans and new gene origination, a structural and evolutionary perspective. Current Opinion in Structural Biology, 2014, 26, 73-83.	5.7	27
11	Quantitative assessment of the structural bias in protein–protein interaction assays. Proteomics, 2008, 8, 4657-4667.	2.2	22
12	The evolution of filamin – A protein domain repeat perspective. Journal of Structural Biology, 2012, 179, 289-298.	2.8	22
13	Massive parallel sequencing questions the pathogenic role of missense variants in dilated cardiomyopathy. International Journal of Cardiology, 2017, 228, 742-748.	1.7	16
14	Identification of a Non-Pentapeptide Region Associated with Rapid Mycobacterial Evolution. PLoS ONE, 2016, 11, e0154059.	2.5	1