## Anushya Muruganujan

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1038269/publications.pdf

Version: 2024-02-01

759233 1058476 26,749 18 12 14 citations h-index g-index papers 19 19 19 42755 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	The Sequence of the Human Genome. Science, 2001, 291, 1304-1351.	12.6	12,623
2	PANTHER: A Library of Protein Families and Subfamilies Indexed by Function. Genome Research, 2003, 13, 2129-2141.	5.5	2,703
3	PANTHER version 14: more genomes, a new PANTHER GO-slim and improvements in enrichment analysis tools. Nucleic Acids Research, 2019, 47, D419-D426.	14.5	2,455
4	Large-scale gene function analysis with the PANTHER classification system. Nature Protocols, 2013, 8, 1551-1566.	12.0	2,276
5	PANTHER version 11: expanded annotation data from Gene Ontology and Reactome pathways, and data analysis tool enhancements. Nucleic Acids Research, 2017, 45, D183-D189.	14.5	2,113
6	Protocol Update for large-scale genome and gene function analysis with the PANTHER classification system (v.14.0). Nature Protocols, 2019, 14, 703-721.	12.0	1,041
7	PANTHER version 16: a revised family classification, tree-based classification tool, enhancer regions and extensive API. Nucleic Acids Research, 2021, 49, D394-D403.	14.5	990
8	PANTHER version 10: expanded protein families and functions, and analysis tools. Nucleic Acids Research, 2016, 44, D336-D342.	14.5	790
9	The PANTHER database of protein families, subfamilies, functions and pathways. Nucleic Acids Research, 2004, 33, D284-D288.	14.5	686
10	PANTHER version 7: improved phylogenetic trees, orthologs and collaboration with the Gene Ontology Consortium. Nucleic Acids Research, 2010, 38, D204-D210.	14.5	553
11	<scp>PANTHER</scp> : Making genomeâ€scale phylogenetics accessible to all. Protein Science, 2022, 31, 8-22.	7.6	467
12	PhyloGenes: An online phylogenetics and functional genomics resource for plant gene function inference. Plant Direct, 2020, 4, e00293.	1.9	23
13	Ancestral Genomes: a resource for reconstructed ancestral genes and genomes across the tree of life. Nucleic Acids Research, 2019, 47, D271-D279.	14.5	17
14	PEREGRINE: A genome-wide prediction of enhancer to gene relationships supported by experimental evidence. PLoS ONE, 2020, 15, e0243791.	2.5	12
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17	Title is missing!. , 2020, 15, e0243791.		O
18	Title is missing!. , 2020, 15, e0243791.		0