## Anushya Muruganujan

## 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.

18 19,519 19 11 h-index g-index citations papers 16.2 6.57 24,180 19 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
18	PANTHER: Making genome-scale phylogenetics accessible to all. <i>Protein Science</i> , <b>2021</b> ,	6.3	12
17	PANTHER version 16: a revised family classification, tree-based classification tool, enhancer regions and extensive API. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, D394-D403	20.1	248
16	PEREGRINE: A genome-wide prediction of enhancer to gene relationships supported by experimental evidence. <i>PLoS ONE</i> , <b>2020</b> , 15, e0243791	3.7	3
15	PhyloGenes: An online phylogenetics and functional genomics resource for plant gene function inference. <i>Plant Direct</i> , <b>2020</b> , 4, e00293	3.3	8
14	PEREGRINE: A genome-wide prediction of enhancer to gene relationships supported by experimental evidence <b>2020</b> , 15, e0243791		
13	PEREGRINE: A genome-wide prediction of enhancer to gene relationships supported by experimental evidence <b>2020</b> , 15, e0243791		
12	PEREGRINE: A genome-wide prediction of enhancer to gene relationships supported by experimental evidence <b>2020</b> , 15, e0243791		
11	PEREGRINE: A genome-wide prediction of enhancer to gene relationships supported by experimental evidence <b>2020</b> , 15, e0243791		
10	Protocol Update for large-scale genome and gene function analysis with the PANTHER classification system (v.14.0). <i>Nature Protocols</i> , <b>2019</b> , 14, 703-721	18.8	462
9	PANTHER version 14: more genomes, a new PANTHER GO-slim and improvements in enrichment analysis tools. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, D419-D426	20.1	1256
8	Ancestral Genomes: a resource for reconstructed ancestral genes and genomes across the tree of life. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, D271-D279	20.1	5
7	PANTHER version 11: expanded annotation data from Gene Ontology and Reactome pathways, and data analysis tool enhancements. <i>Nucleic Acids Research</i> , <b>2017</b> , 45, D183-D189	20.1	1470
6	PANTHER version 10: expanded protein families and functions, and analysis tools. <i>Nucleic Acids Research</i> , <b>2016</b> , 44, D336-42	20.1	656
5	Large-scale gene function analysis with the PANTHER classification system. <i>Nature Protocols</i> , <b>2013</b> , 8, 1551-66	18.8	1648
4	PANTHER version 7: improved phylogenetic trees, orthologs and collaboration with the Gene Ontology Consortium. <i>Nucleic Acids Research</i> , <b>2010</b> , 38, D204-10	20.1	45 <sup>1</sup>
3	The PANTHER database of protein families, subfamilies, functions and pathways. <i>Nucleic Acids Research</i> , <b>2005</b> , 33, D284-8	20.1	589
2	PANTHER: a library of protein families and subfamilies indexed by function. <i>Genome Research</i> , <b>2003</b> , 13, 2129-41	9.7	2102

The sequence of the human genome. *Science*, **2001**, 291, 1304-51

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