

# Fabian Schreiber

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11304538/publications.pdf>

Version: 2024-02-01

11  
papers

2,177  
citations

1040056

9  
h-index

1372567

10  
g-index

11  
all docs

11  
docs citations

11  
times ranked

4153  
citing authors

#	ARTICLE	IF	CITATIONS
1	HMMER web server: 2015 update. <i>Nucleic Acids Research</i> , 2015, 43, W30-W38.	14.5	849
2	Phylogenomics Revives Traditional Views on Deep Animal Relationships. <i>Current Biology</i> , 2009, 19, 706-712.	3.9	611
3	Deep metazoan phylogeny: When different genes tell different stories. <i>Molecular Phylogenetics and Evolution</i> , 2013, 67, 223-233.	2.7	242
4	Standardized benchmarking in the quest for orthologs. <i>Nature Methods</i> , 2016, 13, 425-430.	19.0	198
5	TreeFam v9: a new website, more species and orthology-on-the-fly. <i>Nucleic Acids Research</i> , 2014, 42, D922-D925.	14.5	125
6	Letter to the Editor: SeqXML and OrthoXML: standards for sequence and orthology information. <i>Briefings in Bioinformatics</i> , 2011, 12, 485-488.	6.5	51
7	Hieranoid: Hierarchical Orthology Inference. <i>Journal of Molecular Biology</i> , 2013, 425, 2072-2081.	4.2	38
8	Gearing up to handle the mosaic nature of life in the quest for orthologs. <i>Bioinformatics</i> , 2018, 34, 323-329.	4.1	36
9	OrthoSelect: a protocol for selecting orthologous groups in phylogenomics. <i>BMC Bioinformatics</i> , 2009, 10, 219.	2.6	23
10	OrthoSelect: a web server for selecting orthologous gene alignments from EST sequences. <i>Nucleic Acids Research</i> , 2009, 37, W185-W188.	14.5	3
11	treeWidget: a BioJS component to visualise phylogenetic trees. <i>F1000Research</i> , 0, 3, 49.	1.6	1