

Henning Schiebenhoefer

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.

13 papers	90 citations	6 h-index	9 g-index
17 ext. papers	166 ext. citations	9.2 avg, IF	2.3 L-index

#	Paper	IF	Citations
13	Critical Assessment of MetaProteome Investigation (CAMPI): a multi-laboratory comparison of established workflows.. <i>Nature Communications</i> , 2021 , 12, 7305	17.4	8
12	MegaGO: A Fast Yet Powerful Approach to Assess Functional Gene Ontology Similarity across Meta-Omics Data Sets. <i>Journal of Proteome Research</i> , 2021 , 20, 2083-2088	5.6	6
11	Survey of metaproteomics software tools for functional microbiome analysis. <i>PLoS ONE</i> , 2020 , 15, e0241503	15.3	15
10	A complete and flexible workflow for metaproteomics data analysis based on MetaProteomeAnalyzer and Prophan. <i>Nature Protocols</i> , 2020 , 15, 3212-3239	18.8	14
9	Survey of metaproteomics software tools for functional microbiome analysis 2020 , 15, e0241503		
8	Survey of metaproteomics software tools for functional microbiome analysis 2020 , 15, e0241503		
7	Survey of metaproteomics software tools for functional microbiome analysis 2020 , 15, e0241503		
6	Survey of metaproteomics software tools for functional microbiome analysis 2020 , 15, e0241503		
5	Survey of metaproteomics software tools for functional microbiome analysis 2020 , 15, e0241503		
4	Survey of metaproteomics software tools for functional microbiome analysis 2020 , 15, e0241503		
3	Challenges and promise at the interface of metaproteomics and genomics: an overview of recent progress in metaproteogenomic data analysis. <i>Expert Review of Proteomics</i> , 2019 , 16, 375-390	4.2	33
2	Survey of metaproteomics software tools for functional microbiome analysis		1
1	Critical Assessment of Metaproteome Investigation (CAMPI): A Multi-Lab Comparison of Established Workflows		8