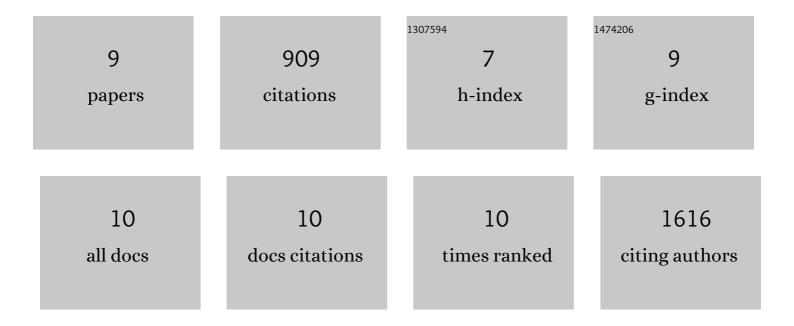
## Yannick Mahlich

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2738249/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	A large-scale evaluation of computational protein function prediction. Nature Methods, 2013, 10, 221-227.	19.0	789
2	Quantifying structural relationships of metal-binding sites suggests origins of biological electron transfer. Science Advances, 2022, 8, eabj3984.	10.3	24
3	HFSP: high speed homology-driven function annotation of proteins. Bioinformatics, 2018, 34, i304-i312.	4.1	22
4	fusionDB: assessing microbial diversity and environmental preferences via functional similarity networks. Nucleic Acids Research, 2018, 46, D535-D541.	14.5	21
5	Common sequence variants affect molecular function more than rare variants?. Scientific Reports, 2017, 7, 1608.	3.3	20
6	Predicted Molecular Effects of Sequence Variants Link to System Level of Disease. PLoS Computational Biology, 2016, 12, e1005047.	3.2	14
7	Fingerprinting cities: differentiating subway microbiome functionality. Biology Direct, 2019, 14, 19.	4.6	11
8	Computational Approaches for Unraveling the Effects of Variation in the Human Genome and Microbiome. Annual Review of Biomedical Data Science, 2020, 3, 411-432.	6.5	5
9	Low Diversity of Human Variation Despite Mostly Mild Functional Impact of De Novo Variants. Frontiers in Molecular Biosciences, 2021, 8, 635382.	3.5	2