

Simon C Potter

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.

10 papers	11,177 citations	10 h-index	10 g-index
10 ext. papers	16,523 ext. citations	19.4 avg, IF	6.28 L-index

#	Paper	IF	Citations
10	MGnify: the microbiome analysis resource in 2020. <i>Nucleic Acids Research</i> , 2020 , 48, D570-D578	20.1	127
9	The EMBL-EBI search and sequence analysis tools APIs in 2019. <i>Nucleic Acids Research</i> , 2019 , 47, W636-W641	20.1	2132
8	The Pfam protein families database in 2019. <i>Nucleic Acids Research</i> , 2019 , 47, D427-D432	20.1	2298
7	InterPro in 2019: improving coverage, classification and access to protein sequence annotations. <i>Nucleic Acids Research</i> , 2019 , 47, D351-D360	20.1	835
6	Ensembl Genomes 2018: an integrated omics infrastructure for non-vertebrate species. <i>Nucleic Acids Research</i> , 2018 , 46, D802-D808	20.1	330
5	EBI Metagenomics in 2017: enriching the analysis of microbial communities, from sequence reads to assemblies. <i>Nucleic Acids Research</i> , 2018 , 46, D726-D735	20.1	130
4	HMMER web server: 2018 update. <i>Nucleic Acids Research</i> , 2018 , 46, W200-W204	20.1	596
3	InterPro in 2017-beyond protein family and domain annotations. <i>Nucleic Acids Research</i> , 2017 , 45, D190-D199	20.1	970
2	The Pfam protein families database: towards a more sustainable future. <i>Nucleic Acids Research</i> , 2016 , 44, D279-85	20.1	3678
1	EBI metagenomics in 2016--an expanding and evolving resource for the analysis and archiving of metagenomic data. <i>Nucleic Acids Research</i> , 2016 , 44, D595-603	20.1	81