

Peter Belmann

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3014943/publications.pdf>

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11
papers

1,096
citations

1307594

7
h-index

1281871

11
g-index

13
all docs

13
docs citations

13
times ranked

2415
citing authors

#	ARTICLE	IF	CITATIONS
1	Multiple Occurrences of a 168-Nucleotide Deletion in SARS-CoV-2 ORF8, Unnoticed by Standard Amplicon Sequencing and Variant Calling Pipelines. <i>Viruses</i> , 2021, 13, 1870.	3.3	7
2	Assessing taxonomic metagenome profilers with OPAL. <i>Genome Biology</i> , 2019, 20, 51.	8.8	67
3	CAMISIM: simulating metagenomes and microbial communities. <i>Microbiome</i> , 2019, 7, 17.	11.1	117
4	de.NBI Cloud federation through ELIXIR AAI. <i>F1000Research</i> , 2019, 8, 842.	1.6	13
5	The de.NBI / ELIXIR-DE training platform - Bioinformatics training in Germany and across Europe within ELIXIR. <i>F1000Research</i> , 2019, 8, 1877.	1.6	1
6	The de.NBI / ELIXIR-DE training platform - Bioinformatics training in Germany and across Europe within ELIXIR. <i>F1000Research</i> , 2019, 8, 1877.	1.6	3
7	AMBER: Assessment of Metagenome BinnerERs. <i>GigaScience</i> , 2018, 7, .	6.4	72
8	Common ELIXIR Service for Researcher Authentication and Authorisation. <i>F1000Research</i> , 2018, 7, 1199.	1.6	23
9	Critical Assessment of Metagenome Interpretation – a benchmark of metagenomics software. <i>Nature Methods</i> , 2017, 14, 1063-1071.	19.0	635
10	Deeply sequenced metagenome and metatranscriptome of a biogas-producing microbial community from an agricultural production-scale biogas plant. <i>GigaScience</i> , 2015, 4, 33.	6.4	68
11	Bioboxes: standardised containers for interchangeable bioinformatics software. <i>GigaScience</i> , 2015, 4, 47.	6.4	84