

Jai Ram Rideout

List of Publications by Citations

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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.

19
papers

7,333
citations

16
h-index

31
g-index

31
ext. papers

12,961
ext. citations

12.3
avg, IF

5.21
L-index

#	Paper	IF	Citations
19	Reproducible, interactive, scalable and extensible microbiome data science using QIIME 2. <i>Nature Biotechnology</i> , 2019 , 37, 852-857	44.5	4050
18	Optimizing taxonomic classification of marker-gene amplicon sequences with QIIME 2's q2-feature-classifier plugin. <i>Microbiome</i> , 2018 , 6, 90	16.6	1227
17	The Biological Observation Matrix (BIOM) format or: how I learned to stop worrying and love the ome-ome. <i>GigaScience</i> , 2012 , 1, 7	7.6	484
16	Subsampled open-reference clustering creates consistent, comprehensive OTU definitions and scales to billions of sequences. <i>PeerJ</i> , 2014 , 2, e545	3.1	395
15	Qiita: rapid, web-enabled microbiome meta-analysis. <i>Nature Methods</i> , 2018 , 15, 796-798	21.6	231
14	The interpersonal and intrapersonal diversity of human-associated microbiota in key body sites. <i>Journal of Allergy and Clinical Immunology</i> , 2012 , 129, 1204-8	11.5	207
13	QIIME 2: Reproducible, interactive, scalable, and extensible microbiome data science	138	
12	q2-longitudinal: Longitudinal and Paired-Sample Analyses of Microbiome Data. <i>MSystems</i> , 2018 , 3,	7.6	104
11	Stability of operational taxonomic units: an important but neglected property for analyzing microbial diversity. <i>Microbiome</i> , 2015 , 3, 20	16.6	88
10	QIIME 2: Reproducible, interactive, scalable, and extensible microbiome data science 2018 ,	78	
9	mockrobiota: a Public Resource for Microbiome Bioinformatics Benchmarking. <i>MSystems</i> , 2016 , 1,	7.6	55
8	Gut microbial and short-chain fatty acid profiles in adults with chronic constipation before and after treatment with lubiprostone. <i>Anaerobe</i> , 2015 , 33, 33-41	2.8	37
7	QIIME 2: Reproducible, interactive, scalable, and extensible microbiome data science	36	
6	Ghost-tree: creating hybrid-gene phylogenetic trees for diversity analyses. <i>Microbiome</i> , 2016 , 4, 11	16.6	30
5	Keemei: cloud-based validation of tabular bioinformatics file formats in Google Sheets. <i>GigaScience</i> , 2016 , 5, 27	7.6	16
4	A standardized, extensible framework for optimizing classification improves marker-gene taxonomic assignments	8	
3	cual-id: Globally Unique, Correctable, and Human-Friendly Sample Identifiers for Comparative Omics Studies. <i>MSystems</i> , 2016 , 1,	7.6	4

LIST OF PUBLICATIONS

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|---|---|---|
| 2 | A standardized, extensible framework for optimizing classification improves marker-gene taxonomic assignments | 2 |
| 1 | A standardized, extensible framework for optimizing classification improves marker-gene taxonomic assignments | 2 |