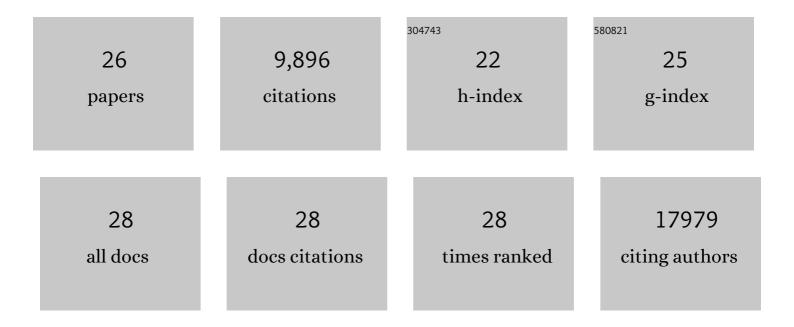
## J Rodney Brister

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3472783/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	The Sequence Read Archive: a decade more of explosive growth. Nucleic Acids Research, 2022, 50, D387-D390.	14.5	128
2	Database resources of the national center for biotechnology information. Nucleic Acids Research, 2022, 50, D20-D26.	14.5	887
3	Bacterial Viruses Subcommittee and Archaeal Viruses Subcommittee of the ICTV: update of taxonomy changes in 2021. Archives of Virology, 2021, 166, 3239-3244.	2.1	24
4	STAT: a fast, scalable, MinHash-based k-mer tool to assess Sequence Read Archive next-generation sequence submissions. Genome Biology, 2021, 22, 270.	8.8	45
5	Perspective on taxonomic classification of uncultivated viruses. Current Opinion in Virology, 2021, 51, 207-215.	5.4	31
6	Analysis of Spounaviruses as a Case Study for the Overdue Reclassification of Tailed Phages. Systematic Biology, 2020, 69, 110-123.	5.6	89
7	Database resources of the National Center for Biotechnology Information. Nucleic Acids Research, 2020, 48, D9-D16.	14.5	381
8	VADR: validation and annotation of virus sequence submissions to GenBank. BMC Bioinformatics, 2020, 21, 211.	2.6	46
9	Taxonomic assignment of uncultivated prokaryotic virus genomes is enabled by gene-sharing networks. Nature Biotechnology, 2019, 37, 632-639.	17.5	569
10	The Phylogeny and Pathogenesis of Sacbrood Virus (SBV) Infection in European Honey Bees, Apis mellifera. Viruses, 2019, 11, 61.	3.3	28
11	Minimum Information about an Uncultivated Virus Genome (MIUViG). Nature Biotechnology, 2019, 37, 29-37.	17.5	414
12	Strengthening the Interaction of the Virology Community with the International Committee on Taxonomy of Viruses (ICTV) by Linking Virus Names and Their Abbreviations to Virus Species. Systematic Biology, 2019, 68, 828-839.	5.6	11
13	Database resources of the National Center for Biotechnology Information. Nucleic Acids Research, 2019, 47, D23-D28.	14.5	502
14	Taxonomy of prokaryotic viruses: 2017 update from the ICTV Bacterial and Archaeal Viruses Subcommittee. Archives of Virology, 2018, 163, 1125-1129.	2.1	172
15	Bacteriophage Taxonomy: An Evolving Discipline. Methods in Molecular Biology, 2018, 1693, 57-71.	0.9	34
16	Virus taxonomy in the age of metagenomics. Nature Reviews Microbiology, 2017, 15, 161-168.	28.6	590
17	Taxonomy of prokaryotic viruses: 2016 update from the ICTV bacterial and archaeal viruses subcommittee. Archives of Virology, 2017, 162, 1153-1157.	2.1	57
18	Virus Variation Resource – improved response to emergent viral outbreaks. Nucleic Acids Research, 2017, 45, D482-D490.	14.5	342

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#	Article	IF	CITATIONS
19	Reference sequence (RefSeq) database at NCBI: current status, taxonomic expansion, and functional annotation. Nucleic Acids Research, 2016, 44, D733-D745.	14.5	4,739
20	HIV-1, human interaction database: current status and new features. Nucleic Acids Research, 2015, 43, D566-D570.	14.5	79
21	NCBI Viral Genomes Resource. Nucleic Acids Research, 2015, 43, D571-D577.	14.5	541
22	Filovirus RefSeq Entries: Evaluation and Selection of Filovirus Type Variants, Type Sequences, and Names. Viruses, 2014, 6, 3663-3682.	3.3	49
23	Nomenclature- and Database-Compatible Names for the Two Ebola Virus Variants that Emerged in Guinea and the Democratic Republic of the Congo in 2014. Viruses, 2014, 6, 4760-4799.	3.3	83
24	Virus Variation Resource—recent updates and future directions. Nucleic Acids Research, 2014, 42, D660-D665.	14.5	43
25	Microbial virus genome annotation—Mustering the troops to fight the sequence onslaught. Virology, 2012, 434, 175-180.	2.4	12
26	Quantifying the immunological distinctiveness of emerging SARS-CoV-2 variants in the context of prior regional herd exposure. , 0, , .		0

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