## **Emily Pritchard**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7552441/publications.pdf

Version: 2024-02-01

22 papers 13,774 citations

430874 18 h-index 677142 22 g-index

24 all docs

24 docs citations

times ranked

24

31865 citing authors

#	Article	IF	CITATIONS
1	Ensembl 2018. Nucleic Acids Research, 2018, 46, D754-D761.	14.5	2,710
2	Ensembl 2016. Nucleic Acids Research, 2016, 44, D710-D716.	14.5	1,372
3	Ensembl 2021. Nucleic Acids Research, 2021, 49, D884-D891.	14.5	1,231
4	Ensembl 2014. Nucleic Acids Research, 2014, 42, D749-D755.	14.5	1,211
5	Ensembl 2015. Nucleic Acids Research, 2015, 43, D662-D669.	14.5	1,145
6	Ensembl 2022. Nucleic Acids Research, 2022, 50, D988-D995.	14.5	1,103
7	Ensembl 2020. Nucleic Acids Research, 2020, 48, D682-D688.	14.5	1,076
8	Ensembl 2019. Nucleic Acids Research, 2019, 47, D745-D751.	14.5	879
9	Ensembl 2017. Nucleic Acids Research, 2017, 45, D635-D642.	14.5	535
10	Ensembl Genomes 2016: more genomes, more complexity. Nucleic Acids Research, 2016, 44, D574-D580.	14.5	530
11	Ensembl Genomes 2018: an integrated omics infrastructure for non-vertebrate species. Nucleic Acids Research, 2018, 46, D802-D808.	14.5	489
12	Ensembl Genomes 2020â€"enabling non-vertebrate genomic research. Nucleic Acids Research, 2020, 48, D689-D695.	14.5	416
13	The International Genome Sample Resource (IGSR) collection of open human genomic variation resources. Nucleic Acids Research, 2020, 48, D941-D947.	14.5	221
14	Ensembl Plants: Integrating Tools for Visualizing, Mining, and Analyzing Plant Genomics Data. Methods in Molecular Biology, 2016, 1374, 115-140.	0.9	216
15	The international Genome sample resource (IGSR): A worldwide collection of genome variation incorporating the 1000 Genomes Project data. Nucleic Acids Research, 2017, 45, D854-D859.	14.5	215
16	Ensembl Plants: Integrating Tools for Visualizing, Mining, and Analyzing Plant Genomic Data. Methods in Molecular Biology, 2017, 1533, 1-31.	0.9	189
17	Ensembl Genomes 2022: an expanding genome resource for non-vertebrates. Nucleic Acids Research, 2022, 50, D996-D1003.	14.5	141
18	Ensembl regulation resources. Database: the Journal of Biological Databases and Curation, 2016, 2016, bav119.	3.0	45

#	Article	IF	CITATIONS
19	Annotating and prioritizing genomic variants using the Ensembl Variant Effect Predictor—A tutorial. Human Mutation, 2022, 43, 986-997.	2.5	30
20	The Ensembl Genome Browser: Strategies for Accessing Eukaryotic Genome Data. Methods in Molecular Biology, 2018, 1757, 115-139.	0.9	13
21	The value of primary transcripts to the clinical and nonâ€clinical genomics community: Survey results and roadmap for improvements. Molecular Genetics & Denomic Medicine, 2021, 9, e1786.	1.2	5
22	Parents in science. Genome Biology, 2018, 19, 180.	8.8	1