

# 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

430442

18  
h-index

676716

22  
g-index

24  
all docs

24  
docs citations

24  
times ranked

31865  
citing authors

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Ensembl 2018. Nucleic Acids Research, 2018, 46, D754-D761.  | 6.5 | 2,710     |
| 2  | Ensembl 2016. Nucleic Acids Research, 2016, 44, D710-D716.  | 6.5 | 1,372     |
| 3  | Ensembl 2021. Nucleic Acids Research, 2021, 49, D884-D891.  | 6.5 | 1,231     |
| 4  | Ensembl 2014. Nucleic Acids Research, 2014, 42, D749-D755.  | 6.5 | 1,211     |
| 5  | Ensembl 2015. Nucleic Acids Research, 2015, 43, D662-D669.  | 6.5 | 1,145     |
| 6  | Ensembl 2022. Nucleic Acids Research, 2022, 50, D988-D995.  | 6.5 | 1,103     |
| 7  | Ensembl 2020. Nucleic Acids Research, 2020, 48, D682-D688.  | 6.5 | 1,076     |
| 8  | Ensembl 2019. Nucleic Acids Research, 2019, 47, D745-D751.  | 6.5 | 879       |
| 9  | Ensembl 2017. Nucleic Acids Research, 2017, 45, D635-D642.  | 6.5 | 535       |
| 10 | Ensembl Genomes 2016: more genomes, more complexity. Nucleic Acids Research, 2016, 44, D574-D580.   | 6.5 | 530       |
| 11 | Ensembl Genomes 2018: an integrated omics infrastructure for non-vertebrate species. Nucleic Acids Research, 2018, 46, D802-D808.   | 6.5 | 489       |
| 12 | Ensembl Genomes 2020“enabling non-vertebrate genomic research. Nucleic Acids Research, 2020, 48, D689-D695.   | 6.5 | 416       |
| 13 | The International Genome Sample Resource (IGSR) collection of open human genomic variation resources. Nucleic Acids Research, 2020, 48, D941-D947.                                    | 6.5 | 221       |
| 14 | Ensembl Plants: Integrating Tools for Visualizing, Mining, and Analyzing Plant Genomics Data. Methods in Molecular Biology, 2016, 1374, 115-140.                                      | 0.4 | 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. | 6.5 | 215       |
| 16 | Ensembl Plants: Integrating Tools for Visualizing, Mining, and Analyzing Plant Genomic Data. Methods in Molecular Biology, 2017, 1533, 1-31.  | 0.4 | 189       |
| 17 | Ensembl Genomes 2022: an expanding genome resource for non-vertebrates. Nucleic Acids Research, 2022, 50, D996-D1003.   | 6.5 | 141       |
| 18 | Ensembl regulation resources. Database: the Journal of Biological Databases and Curation, 2016, 2016, bav119.   | 1.4 | 45        |

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 19 | Annotating and prioritizing genomic variants using the Ensembl Variant Effect Predictor—a tutorial. <i>Human Mutation</i> , 2022, 43, 986-997.  | 1.1 | 30        |
| 20 | The Ensembl Genome Browser: Strategies for Accessing Eukaryotic Genome Data. <i>Methods in Molecular Biology</i> , 2018, 1757, 115-139.   | 0.4 | 13        |
| 21 | The value of primary transcripts to the clinical and non-clinical genomics community: Survey results and roadmap for improvements. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2021, 9, e1786. | 0.6 | 5         |
| 22 | Parents in science. <i>Genome Biology</i> , 2018, 19, 180.  | 3.8 | 1         |