

# Magali Ruffier

## 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

9,061  
citations

17  
h-index

19  
g-index

19  
ext. papers

13,336  
ext. citations

15.2  
avg, IF

4.82  
L-index

#	Paper	IF	Citations
19	Ensembl 2018. <i>Nucleic Acids Research</i> , <b>2018</b> , 46, D754-D761	20.1	1822
18	GENCODE reference annotation for the human and mouse genomes. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, D766-D773	20.1	1140
17	Ensembl 2016. <i>Nucleic Acids Research</i> , <b>2016</b> , 44, D710-6	20.1	1094
16	Ensembl 2014. <i>Nucleic Acids Research</i> , <b>2014</b> , 42, D749-55	20.1	1087
15	Ensembl 2015. <i>Nucleic Acids Research</i> , <b>2015</b> , 43, D662-9	20.1	1013
14	Ensembl 2020. <i>Nucleic Acids Research</i> , <b>2020</b> , 48, D682-D688	20.1	645
13	Ensembl 2019. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, D745-D751	20.1	554
12	The Ensembl gene annotation system. <i>Database: the Journal of Biological Databases and Curation</i> , <b>2016</b> , 2016,	5	537
11	Ensembl 2017. <i>Nucleic Acids Research</i> , <b>2017</b> , 45, D635-D642	20.1	404
10	Ensembl 2021. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, D884-D891	20.1	324
9	The Ensembl REST API: Ensembl Data for Any Language. <i>Bioinformatics</i> , <b>2015</b> , 31, 143-5	7.2	108
8	A comprehensive genomic history of extinct and living elephants. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2018</b> , 115, E2566-E2574	11.5	86
7	GENCODE 2021. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, D916-D923	20.1	82
6	Ensembl 2022. <i>Nucleic Acids Research</i> , <b>2021</b> ,	20.1	72
5	Ensembl core software resources: storage and programmatic access for DNA sequence and genome annotation. <i>Database: the Journal of Biological Databases and Curation</i> , <b>2017</b> , 2017,	5	35
4	Variant calling on the GRCh38 assembly with the data from phase three of the 1000 Genomes Project. <i>Wellcome Open Research</i> , <b>2019</b> , 4, 50	4.8	24
3	Variant calling on the GRCh38 assembly with the data from phase three of the 1000 Genomes Project. <i>Wellcome Open Research</i> , <b>2019</b> , 4, 50	4.8	19

- 2 Ensembl Genomes 2022: an expanding genome resource for non-vertebrates. *Nucleic Acids Research*, **2021**, 20.1 12
- 1 The Ensembl COVID-19 resource: ongoing integration of public SARS-CoV-2 data. *Nucleic Acids Research*, **2021**, 20.1 3