

Carlos Garca Girn

List of Publications by Year in Descending Order

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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.

14
papers

8,822
citations

13
h-index

15
g-index

15
ext. papers

12,959
ext. citations

17.3
avg, IF

4.72
L-index

#	Paper	IF	Citations
14	Ensembl 2022. <i>Nucleic Acids Research</i> , 2021 ,	20.1	72
13	Ensembl 2021. <i>Nucleic Acids Research</i> , 2021 , 49, D884-D891	20.1	324
12	GENCODE 2021. <i>Nucleic Acids Research</i> , 2021 , 49, D916-D923	20.1	82
11	K-mer counting and curated libraries drive efficient annotation of repeats in plant genomes. <i>Plant Genome</i> , 2021 , 14, e20143	4.4	2
10	Ensembl 2020. <i>Nucleic Acids Research</i> , 2020 , 48, D682-D688	20.1	645
9	GENCODE reference annotation for the human and mouse genomes. <i>Nucleic Acids Research</i> , 2019 , 47, D766-D773	20.1	1140
8	Ensembl 2019. <i>Nucleic Acids Research</i> , 2019 , 47, D745-D751	20.1	554
7	Ensembl 2018. <i>Nucleic Acids Research</i> , 2018 , 46, D754-D761	20.1	1822
6	Consensus coding sequence (CCDS) database: a standardized set of human and mouse protein-coding regions supported by expert curation. <i>Nucleic Acids Research</i> , 2018 , 46, D221-D228	20.1	46
5	Ensembl 2017. <i>Nucleic Acids Research</i> , 2017 , 45, D635-D642	20.1	404
4	Ensembl 2016. <i>Nucleic Acids Research</i> , 2016 , 44, D710-6	20.1	1094
3	The Ensembl gene annotation system. <i>Database: the Journal of Biological Databases and Curation</i> , 2016 , 2016,	5	537
2	Ensembl 2015. <i>Nucleic Acids Research</i> , 2015 , 43, D662-9	20.1	1013
1	Ensembl 2014. <i>Nucleic Acids Research</i> , 2014 , 42, D749-55	20.1	1087