

Adam Frankish

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/4470639/adam-frankish-publications-by-year.pdf>

Version: 2024-04-26

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

31
papers

9,282
citations

20
h-index

37
g-index

37
ext. papers

13,910
ext. citations

17.7
avg, IF

5.07
L-index

#	Paper	IF	Citations
31	Non-coding regulatory elements: potential roles in disease and the case of epilepsy. <i>Neuropathology and Applied Neurobiology</i> , 2021 ,	5.2	2
30	Ensembl 2022. <i>Nucleic Acids Research</i> , 2021 ,	20.1	72
29	Ensembl 2021. <i>Nucleic Acids Research</i> , 2021 , 49, D884-D891	20.1	324
28	GENCODE 2021. <i>Nucleic Acids Research</i> , 2021 , 49, D916-D923	20.1	82
27	The Ensembl COVID-19 resource: ongoing integration of public SARS-CoV-2 data. <i>Nucleic Acids Research</i> , 2021 ,	20.1	3
26	A spatially resolved brain region- and cell type-specific isoform atlas of the postnatal mouse brain. <i>Nature Communications</i> , 2021 , 12, 463	17.4	27
25	The value of primary transcripts to the clinical and non-clinical genomics community: Survey results and roadmap for improvements. <i>Molecular Genetics & Genomic Medicine</i> , 2021 , e1786	2.3	2
24	Progress, Challenges, and Surprises in Annotating the Human Genome. <i>Annual Review of Genomics and Human Genetics</i> , 2020 , 21, 55-79	9.7	6
23	Expert curation of the human and mouse olfactory receptor gene repertoires identifies conserved coding regions split across two exons. <i>BMC Genomics</i> , 2020 , 21, 196	4.5	10
22	Ensembl 2020. <i>Nucleic Acids Research</i> , 2020 , 48, D682-D688	20.1	645
21	Transcriptional activity and strain-specific history of mouse pseudogenes. <i>Nature Communications</i> , 2020 , 11, 3695	17.4	8
20	Perspectives on ENCODE. <i>Nature</i> , 2020 , 583, 693-698	50.4	61
19	Integrative transcriptomic analysis suggests new autoregulatory splicing events coupled with nonsense-mediated mRNA decay. <i>Nucleic Acids Research</i> , 2019 , 47, 5293-5306	20.1	22
18	Getting the Entire Message: Progress in Isoform Sequencing. <i>Frontiers in Genetics</i> , 2019 , 10, 709	4.5	16
17	Discovery of high-confidence human protein-coding genes and exons by whole-genome PhyloCSF helps elucidate 118 GWAS loci. <i>Genome Research</i> , 2019 , 29, 2073-2087	9.7	26
16	Re-annotation of 191 developmental and epileptic encephalopathy-associated genes unmasks de novo variants in. <i>Npj Genomic Medicine</i> , 2019 , 4, 31	6.2	12
15	GENCODE reference annotation for the human and mouse genomes. <i>Nucleic Acids Research</i> , 2019 , 47, D766-D773	20.1	1140

14	Ensembl 2019. <i>Nucleic Acids Research</i> , 2019 , 47, D745-D751	20.1	554
13	Ensembl 2018. <i>Nucleic Acids Research</i> , 2018 , 46, D754-D761	20.1	1822
12	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
11	Genome-wide association study: Exploring the genetic basis for responsiveness to ketogenic dietary therapies for drug-resistant epilepsy. <i>Epilepsia</i> , 2018 , 59, 1557-1566	6.4	6
10	Sixteen diverse laboratory mouse reference genomes define strain-specific haplotypes and novel functional loci. <i>Nature Genetics</i> , 2018 , 50, 1574-1583	36.3	91
9	Towards a complete map of the human long non-coding RNA transcriptome. <i>Nature Reviews Genetics</i> , 2018 , 19, 535-548	30.1	248
8	High-throughput annotation of full-length long noncoding RNAs with capture long-read sequencing. <i>Nature Genetics</i> , 2017 , 49, 1731-1740	36.3	140
7	Comparative analysis of pseudogenes across three phyla. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 13361-6	11.5	54
6	The GENCODE pseudogene resource. <i>Genome Biology</i> , 2012 , 13, R51	18.3	232
5	GENCODE: the reference human genome annotation for The ENCODE Project. <i>Genome Research</i> , 2012 , 22, 1760-74	9.7	3142
4	Combining RT-PCR-seq and RNA-seq to catalog all genic elements encoded in the human genome. <i>Genome Research</i> , 2012 , 22, 1698-710	9.7	44
3	GENCODE: producing a reference annotation for ENCODE. <i>Genome Biology</i> , 2006 , 7 Suppl 1, S4.1-9	18.3	434
2	High-throughput annotation of full-length long noncoding RNAs with Capture Long-Read Sequencing		3
1	A community-driven roadmap to advance research on translated open reading frames detected by Ribo-seq		4