

Kim D Pruitt

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.

67
papers

21,311
citations

46
h-index

70
g-index

70
ext. papers

28,163
ext. citations

15.5
avg, IF

6.61
L-index

#	Paper	IF	Citations
67	Standards recommendations for the Earth BioGenome Project.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022 , 119,	11.5	4
66	Database resources of the national center for biotechnology information. <i>Nucleic Acids Research</i> , 2021 ,	20.1	55
65	GenBank. <i>Nucleic Acids Research</i> , 2021 ,	20.1	20
64	GenBank. <i>Nucleic Acids Research</i> , 2021 , 49, D92-D96	20.1	59
63	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , 2021 , 49, D10-D17	20.1	143
62	GenBank. <i>Nucleic Acids Research</i> , 2020 , 48, D84-D86	20.1	130
61	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , 2020 , 48, D9-D16	20.1	216
60	RefSeq curation and annotation of stop codon recoding in vertebrates. <i>Nucleic Acids Research</i> , 2019 , 47, 594-606	20.1	21
59	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , 2019 , 47, D23-D28	20.1	301
58	GenBank. <i>Nucleic Acids Research</i> , 2019 , 47, D94-D99	20.1	217
57	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , 2018 , 46, D8-D13	20.1	746
56	Clonal polymorphism and high heterozygosity in the celibate genome of the Amazon molly. <i>Nature Ecology and Evolution</i> , 2018 , 2, 669-679	12.3	60
55	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
54	GenBank. <i>Nucleic Acids Research</i> , 2018 , 46, D41-D47	20.1	351
53	RefSeq: an update on prokaryotic genome annotation and curation. <i>Nucleic Acids Research</i> , 2018 , 46, D851-D860	20.1	428
52	Evaluation of GRCh38 and de novo haploid genome assemblies demonstrates the enduring quality of the reference assembly. <i>Genome Research</i> , 2017 , 27, 849-864	9.7	365
51	A New Chicken Genome Assembly Provides Insight into Avian Genome Structure. <i>G3: Genes, Genomes, Genetics</i> , 2017 , 7, 109-117	3.2	143

50	A curated catalog of canine and equine keratin genes. <i>PLoS ONE</i> , 2017 , 12, e0180359	3.7	12
49	Genome assembly and transcriptome resource for river buffalo, <i>Bubalus bubalis</i> (2n = 50). <i>GigaScience</i> , 2017 , 6, 1-6	7.6	27
48	Reference sequence (RefSeq) database at NCBI: current status, taxonomic expansion, and functional annotation. <i>Nucleic Acids Research</i> , 2016 , 44, D733-45	20.1	2530
47	NCBI prokaryotic genome annotation pipeline. <i>Nucleic Acids Research</i> , 2016 , 44, 6614-24	20.1	2376
46	Meeting report: GenBank microbial genomic taxonomy workshop (12-13 May, 2015). <i>Standards in Genomic Sciences</i> , 2016 , 11,		51
45	Assembly: a resource for assembled genomes at NCBI. <i>Nucleic Acids Research</i> , 2016 , 44, D73-80	20.1	151
44	RNAcentral: an international database of ncRNA sequences. <i>Nucleic Acids Research</i> , 2015 , 43, D123-9	20.1	89
43	Mouse genome annotation by the RefSeq project. <i>Mammalian Genome</i> , 2015 , 26, 379-90	3.2	8
42	Creating a specialist protein resource network: a meeting report for the protein bioinformatics and community resources retreat. <i>Database: the Journal of Biological Databases and Curation</i> , 2015 , 2015, bav063	5	5
41	Gene: a gene-centered information resource at NCBI. <i>Nucleic Acids Research</i> , 2015 , 43, D36-42	20.1	345
40	RefSeq curation and annotation of antizyme and antizyme inhibitor genes in vertebrates. <i>Nucleic Acids Research</i> , 2015 , 43, 7270-9	20.1	4
39	HIV-1, human interaction database: current status and new features. <i>Nucleic Acids Research</i> , 2015 , 43, D566-70	20.1	61
38	Current status and new features of the Consensus Coding Sequence database. <i>Nucleic Acids Research</i> , 2014 , 42, D865-72	20.1	122
37	RefSeq: an update on mammalian reference sequences. <i>Nucleic Acids Research</i> , 2014 , 42, D756-63	20.1	713
36	Comparison of RefSeq protein-coding regions in human and vertebrate genomes. <i>BMC Genomics</i> , 2013 , 14, 654	4.5	11
35	Tracking and coordinating an international curation effort for the CCDS Project. <i>Database: the Journal of Biological Databases and Curation</i> , 2012 , 2012, bas008	5	44
34	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , 2012 , 40, D13-25	20.1	458
33	NCBI Reference Sequences (RefSeq): current status, new features and genome annotation policy. <i>Nucleic Acids Research</i> , 2012 , 40, D130-5	20.1	876

32	BioProject and BioSample databases at NCBI: facilitating capture and organization of metadata. <i>Nucleic Acids Research</i> , 2012 , 40, D57-63	20.1	180
31	Solving the Problem: Genome Annotation Standards before the Data Deluge. <i>Standards in Genomic Sciences</i> , 2011 , 5, 168-93		46
30	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , 2011 , 39, D38-51	20.1	510
29	Entrez Gene: gene-centered information at NCBI. <i>Nucleic Acids Research</i> , 2011 , 39, D52-7	20.1	546
28	RNAcentral: A vision for an international database of RNA sequences. <i>Rna</i> , 2011 , 17, 1941-6	5.8	54
27	The consensus coding sequence (CCDS) project: Identifying a common protein-coding gene set for the human and mouse genomes. <i>Genome Research</i> , 2009 , 19, 1316-23	9.7	415
26	The completion of the Mammalian Gene Collection (MGC). <i>Genome Research</i> , 2009 , 19, 2324-33	9.7	98
25	Human immunodeficiency virus type 1, human protein interaction database at NCBI. <i>Nucleic Acids Research</i> , 2009 , 37, D417-22	20.1	195
24	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , 2009 , 37, D5-15	20.1	714
23	NCBI Reference Sequences: current status, policy and new initiatives. <i>Nucleic Acids Research</i> , 2009 , 37, D32-6	20.1	619
22	What everybody should know about the rat genome and its online resources. <i>Nature Genetics</i> , 2008 , 40, 523-7	36.3	37
21	Cataloguing the HIV type 1 human protein interaction network. <i>AIDS Research and Human Retroviruses</i> , 2008 , 24, 1497-502	1.6	103
20	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , 2008 , 36, D13-21	20.1	631
19	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , 2007 , 35, D5-12	20.1	693
18	NCBI reference sequences (RefSeq): a curated non-redundant sequence database of genomes, transcripts and proteins. <i>Nucleic Acids Research</i> , 2007 , 35, D61-5	20.1	2095
17	Entrez Gene: gene-centered information at NCBI. <i>Nucleic Acids Research</i> , 2007 , 35, D26-31	20.1	452
16	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , 2006 , 34, D173-80	20.1	394
15	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , 2005 , 33, D39-45	20.1	343

14	NCBI Reference Sequence (RefSeq): a curated non-redundant sequence database of genomes, transcripts and proteins. <i>Nucleic Acids Research</i> , 2005 , 33, D501-4	20.1	1186
13	NCBI Reference Sequence project: update and current status. <i>Nucleic Acids Research</i> , 2003 , 31, 34-7	20.1	128
12	Accessing the human genome. <i>Current Protocols in Human Genetics</i> , 2002 , Chapter 6, Unit 6.9	3.2	
11	Introducing RefSeq and LocusLink: curated human genome resources at the NCBI. <i>Trends in Genetics</i> , 2000 , 16, 44-7	8.5	204
10	Data management and analysis for gene expression arrays. <i>Nature Genetics</i> , 1998 , 20, 19-23	36.3	266
9	WebWise: guide to the University of Washington Genome Center web site. <i>Genome Research</i> , 1998 , 8, 572-5	9.7	0
8	WebWise: guide to McDermott Center for Human Growth and Development at University of Texas Southwest web site. <i>Genome Research</i> , 1998 , 8, 422-6	9.7	
7	WebWise: navigating the Human Genome Project. <i>Genome Research</i> , 1997 , 7, 1038-9	9.7	8
6	Editing of rps3/rpl16 transcripts creates a premature truncation of the rpl16 open reading frame. <i>Current Genetics</i> , 1993 , 23, 472-6	2.9	25
5	Transcription of the Petunia mitochondrial CMS-associated Pcf locus in male sterile and fertility-restored lines. <i>Molecular Genetics and Genomics</i> , 1991 , 227, 348-55		62
4	Splicing of the Petunia cytochrome oxidase subunit II intron. <i>Current Genetics</i> , 1991 , 19, 191-7	2.9	19
3	Cytoplasmic Male Sterility in Petunia 1991 , 383-399		4
2	Cytochrome oxidase subunit II sequences in Petunia mitochondria: two intron-containing genes and an intron-less pseudogene associated with cytoplasmic male sterility. <i>Current Genetics</i> , 1989 , 16, 281-91	2.9	68
1	Evaluation of GRCh38 and de novo haploid genome assemblies demonstrates the enduring quality of the reference assembly		13