

# Kim D Pruitt

## List of Publications by Citations

**Source:** <https://exaly.com/author-pdf/11336006/kim-d-pruitt-publications-by-citations.pdf>

**Version:** 2024-04-25

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.

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	Reference sequence (RefSeq) database at NCBI: current status, taxonomic expansion, and functional annotation. <i>Nucleic Acids Research</i> , <b>2016</b> , 44, D733-45	20.1	2530
66	NCBI prokaryotic genome annotation pipeline. <i>Nucleic Acids Research</i> , <b>2016</b> , 44, 6614-24	20.1	2376
65	NCBI reference sequences (RefSeq): a curated non-redundant sequence database of genomes, transcripts and proteins. <i>Nucleic Acids Research</i> , <b>2007</b> , 35, D61-5	20.1	2095
64	NCBI Reference Sequence (RefSeq): a curated non-redundant sequence database of genomes, transcripts and proteins. <i>Nucleic Acids Research</i> , <b>2005</b> , 33, D501-4	20.1	1186
63	NCBI Reference Sequences (RefSeq): current status, new features and genome annotation policy. <i>Nucleic Acids Research</i> , <b>2012</b> , 40, D130-5	20.1	876
62	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , <b>2018</b> , 46, D8-D13	20.1	746
61	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , <b>2009</b> , 37, D5-15	20.1	714
60	RefSeq: an update on mammalian reference sequences. <i>Nucleic Acids Research</i> , <b>2014</b> , 42, D756-63	20.1	713
59	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , <b>2007</b> , 35, D5-12	20.1	693
58	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , <b>2008</b> , 36, D13-21	20.1	631
57	NCBI Reference Sequences: current status, policy and new initiatives. <i>Nucleic Acids Research</i> , <b>2009</b> , 37, D32-6	20.1	619
56	Entrez Gene: gene-centered information at NCBI. <i>Nucleic Acids Research</i> , <b>2011</b> , 39, D52-7	20.1	546
55	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , <b>2011</b> , 39, D38-51	20.1	510
54	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , <b>2012</b> , 40, D13-25	20.1	458
53	Entrez Gene: gene-centered information at NCBI. <i>Nucleic Acids Research</i> , <b>2007</b> , 35, D26-31	20.1	452
52	RefSeq: an update on prokaryotic genome annotation and curation. <i>Nucleic Acids Research</i> , <b>2018</b> , 46, D851-D860	20.1	428
51	The consensus coding sequence (CCDS) project: Identifying a common protein-coding gene set for the human and mouse genomes. <i>Genome Research</i> , <b>2009</b> , 19, 1316-23	9.7	415

50	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , <b>2006</b> , 34, D173-80	20.1	394
49	Evaluation of GRCh38 and de novo haploid genome assemblies demonstrates the enduring quality of the reference assembly. <i>Genome Research</i> , <b>2017</b> , 27, 849-864	9.7	365
48	GenBank. <i>Nucleic Acids Research</i> , <b>2018</b> , 46, D41-D47	20.1	351
47	Gene: a gene-centered information resource at NCBI. <i>Nucleic Acids Research</i> , <b>2015</b> , 43, D36-42	20.1	345
46	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , <b>2005</b> , 33, D39-45	20.1	343
45	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, D23-D28	20.1	301
44	Data management and analysis for gene expression arrays. <i>Nature Genetics</i> , <b>1998</b> , 20, 19-23	36.3	266
43	GenBank. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, D94-D99	20.1	217
42	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , <b>2020</b> , 48, D9-D16	20.1	216
41	Introducing RefSeq and LocusLink: curated human genome resources at the NCBI. <i>Trends in Genetics</i> , <b>2000</b> , 16, 44-7	8.5	204
40	Human immunodeficiency virus type 1, human protein interaction database at NCBI. <i>Nucleic Acids Research</i> , <b>2009</b> , 37, D417-22	20.1	195
39	BioProject and BioSample databases at NCBI: facilitating capture and organization of metadata. <i>Nucleic Acids Research</i> , <b>2012</b> , 40, D57-63	20.1	180
38	Assembly: a resource for assembled genomes at NCBI. <i>Nucleic Acids Research</i> , <b>2016</b> , 44, D73-80	20.1	151
37	A New Chicken Genome Assembly Provides Insight into Avian Genome Structure. <i>G3: Genes, Genomes, Genetics</i> , <b>2017</b> , 7, 109-117	3.2	143
36	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, D10-D17	20.1	143
35	GenBank. <i>Nucleic Acids Research</i> , <b>2020</b> , 48, D84-D86	20.1	130
34	NCBI Reference Sequence project: update and current status. <i>Nucleic Acids Research</i> , <b>2003</b> , 31, 34-7	20.1	128
33	Current status and new features of the Consensus Coding Sequence database. <i>Nucleic Acids Research</i> , <b>2014</b> , 42, D865-72	20.1	122

32	Cataloguing the HIV type 1 human protein interaction network. <i>AIDS Research and Human Retroviruses</i> , <b>2008</b> , 24, 1497-502	1.6	103
31	The completion of the Mammalian Gene Collection (MGC). <i>Genome Research</i> , <b>2009</b> , 19, 2324-33	9.7	98
30	RNAcentral: an international database of ncRNA sequences. <i>Nucleic Acids Research</i> , <b>2015</b> , 43, D123-9	20.1	89
29	Cytochrome oxidase subunit II sequences in <i>Petunia</i> mitochondria: two intron-containing genes and an intron-less pseudogene associated with cytoplasmic male sterility. <i>Current Genetics</i> , <b>1989</b> , 16, 281-91	2.9	68
28	Transcription of the <i>Petunia</i> mitochondrial CMS-associated Pcf locus in male sterile and fertility-restored lines. <i>Molecular Genetics and Genomics</i> , <b>1991</b> , 227, 348-55		62
27	HIV-1, human interaction database: current status and new features. <i>Nucleic Acids Research</i> , <b>2015</b> , 43, D566-70	20.1	61
26	Clonal polymorphism and high heterozygosity in the celibate genome of the Amazon molly. <i>Nature Ecology and Evolution</i> , <b>2018</b> , 2, 669-679	12.3	60
25	GenBank. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, D92-D96	20.1	59
24	Database resources of the national center for biotechnology information. <i>Nucleic Acids Research</i> , <b>2021</b> ,	20.1	55
23	RNAcentral: A vision for an international database of RNA sequences. <i>Rna</i> , <b>2011</b> , 17, 1941-6	5.8	54
22	Meeting report: GenBank microbial genomic taxonomy workshop (12-13 May, 2015). <i>Standards in Genomic Sciences</i> , <b>2016</b> , 11,		51
21	Consensus coding sequence (CCDS) database: a standardized set of human and mouse protein-coding regions supported by expert curation. <i>Nucleic Acids Research</i> , <b>2018</b> , 46, D221-D228	20.1	46
20	Solving the Problem: Genome Annotation Standards before the Data Deluge. <i>Standards in Genomic Sciences</i> , <b>2011</b> , 5, 168-93		46
19	Tracking and coordinating an international curation effort for the CCDS Project. <i>Database: the Journal of Biological Databases and Curation</i> , <b>2012</b> , 2012, bas008	5	44
18	What everybody should know about the rat genome and its online resources. <i>Nature Genetics</i> , <b>2008</b> , 40, 523-7	36.3	37
17	Genome assembly and transcriptome resource for river buffalo, <i>Bubalus bubalis</i> (2n = 50). <i>GigaScience</i> , <b>2017</b> , 6, 1-6	7.6	27
16	Editing of rps3/rpl16 transcripts creates a premature truncation of the rpl16 open reading frame. <i>Current Genetics</i> , <b>1993</b> , 23, 472-6	2.9	25
15	RefSeq curation and annotation of stop codon recoding in vertebrates. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, 594-606	20.1	21

14	GenBank. <i>Nucleic Acids Research</i> , <b>2021</b> ,	20.1	20
13	Splicing of the <i>Petunia</i> cytochrome oxidase subunit II intron. <i>Current Genetics</i> , <b>1991</b> , 19, 191-7	2.9	19
12	Evaluation of GRCh38 and de novo haploid genome assemblies demonstrates the enduring quality of the reference assembly		13
11	A curated catalog of canine and equine keratin genes. <i>PLoS ONE</i> , <b>2017</b> , 12, e0180359	3.7	12
10	Comparison of RefSeq protein-coding regions in human and vertebrate genomes. <i>BMC Genomics</i> , <b>2013</b> , 14, 654	4.5	11
9	Mouse genome annotation by the RefSeq project. <i>Mammalian Genome</i> , <b>2015</b> , 26, 379-90	3.2	8
8	WebWise: navigating the Human Genome Project. <i>Genome Research</i> , <b>1997</b> , 7, 1038-9	9.7	8
7	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> , <b>2015</b> , 2015, bav063	5	5
6	RefSeq curation and annotation of antizyme and antizyme inhibitor genes in vertebrates. <i>Nucleic Acids Research</i> , <b>2015</b> , 43, 7270-9	20.1	4
5	Standards recommendations for the Earth BioGenome Project.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2022</b> , 119,	11.5	4
4	Cytoplasmic Male Sterility in <i>Petunia</i> <b>1991</b> , 383-399		4
3	WebWise: guide to the University of Washington Genome Center web site. <i>Genome Research</i> , <b>1998</b> , 8, 572-5	9.7	0
2	Accessing the human genome. <i>Current Protocols in Human Genetics</i> , <b>2002</b> , Chapter 6, Unit 6.9	3.2	
1	WebWise: guide to McDermott Center for Human Growth and Development at University of Texas Southwest web site. <i>Genome Research</i> , <b>1998</b> , 8, 422-6	9.7	