

# Karen J Yook

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4529767/publications.pdf>

Version: 2024-02-01

20  
papers

2,181  
citations

516710

16  
h-index

752698

20  
g-index

20  
all docs

20  
docs citations

20  
times ranked

3792  
citing authors

#	ARTICLE	IF	CITATIONS
1	WormBase in 2022â€”data, processes, and tools for analyzing <i>Caenorhabditis elegans</i> . <i>Genetics</i> , 2022, 220, .	2.9	128
2	Harmonizing model organism data in the Alliance of Genome Resources. <i>Genetics</i> , 2022, 220, .	2.9	52
3	WormBase: a modern Model Organism Information Resource. <i>Nucleic Acids Research</i> , 2020, 48, D762-D767.	14.5	213
4	Alliance of Genome Resources Portal: unified model organism research platform. <i>Nucleic Acids Research</i> , 2020, 48, D650-D658.	14.5	145
5	WormBase 2017: molting into a new stage. <i>Nucleic Acids Research</i> , 2018, 46, D869-D874.	14.5	172
6	Micropublication: incentivizing community curation and placing unpublished data into the public domain. <i>Database: the Journal of Biological Databases and Curation</i> , 2018, 2018, .	3.0	22
7	WormBase 2016: expanding to enable helminth genomic research. <i>Nucleic Acids Research</i> , 2016, 44, D774-D780.	14.5	329
8	WormBase 2014: new views of curated biology. <i>Nucleic Acids Research</i> , 2014, 42, D789-D793.	14.5	149
9	WormBase. <i>Worm</i> , 2012, 1, 15-21.	1.0	14
10	WormBase 2012: more genomes, more data, new website. <i>Nucleic Acids Research</i> , 2012, 40, D735-D741.	14.5	175
11	Toward an interactive article: integrating journals and biological databases. <i>BMC Bioinformatics</i> , 2011, 12, 175.	2.6	12
12	Worm Phenotype Ontology: Integrating phenotype data within and beyond the <i>C. elegans</i> community. <i>BMC Bioinformatics</i> , 2011, 12, 32.	2.6	62
13	WormBase: a comprehensive resource for nematode research. <i>Nucleic Acids Research</i> , 2010, 38, D463-D467.	14.5	325
14	Mos1 Mutagenesis Reveals a Diversity of Mechanisms Affecting Response of <i>Caenorhabditis elegans</i> to the Bacterial Pathogen <i>Microbacterium nematophilum</i> . <i>Genetics</i> , 2007, 175, 681-697.	2.9	71
15	WormBase 2007. <i>Nucleic Acids Research</i> , 2007, 36, D612-D617.	14.5	95
16	Genome characterization, analysis of virulence and transformation of <i>Microbacterium nematophilum</i> , a coryneform pathogen of the nematode <i>Caenorhabditis elegans</i> . <i>FEMS Microbiology Letters</i> , 2006, 264, 145-151.	1.8	20
17	Multiple Genes Affect Sensitivity of <i>Caenorhabditis elegans</i> to the Bacterial Pathogen <i>Microbacterium nematophilum</i> . <i>Genetics</i> , 2005, 171, 1033-1045.	2.9	108
18	Complementation. <i>WormBook</i> , 2005, , 1-17.	5.3	14

#	ARTICLE	IF	CITATIONS
19	Rules of Nonallelic Noncomplementation at the Synapse in <i>Caenorhabditis elegans</i> . <i>Genetics</i> , 2001, 158, 209-220.	2.9	45
20	Selective Loss of Sperm Bearing a Compound Chromosome in the <i>Drosophila</i> Female. <i>Genetics</i> , 1996, 143, 1629-1642.	2.9	30