

# Michael Han

## List of Publications by Year in descending order

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Version: 2024-02-01

28  
papers

7,133  
citations

257450

24  
h-index

501196

28  
g-index

29  
all docs

29  
docs citations

29  
times ranked

11950  
citing authors

#	ARTICLE	IF	CITATIONS
1	Ensembl Genomes 2022: an expanding genome resource for non-vertebrates. <i>Nucleic Acids Research</i> , 2022, 50, D996-D1003.	14.5	141
2	WormBase in 2022—data, processes, and tools for analyzing <i>Caenorhabditis elegans</i> . <i>Genetics</i> , 2022, 220, .	2.9	128
3	Harmonizing model organism data in the Alliance of Genome Resources. <i>Genetics</i> , 2022, 220, .	2.9	52
4	The Gene Ontology resource: enriching a GOLD mine. <i>Nucleic Acids Research</i> , 2021, 49, D325-D334.	14.5	2,416
5	WormBase: a modern Model Organism Information Resource. <i>Nucleic Acids Research</i> , 2020, 48, D762-D767.	14.5	213
6	Alliance of Genome Resources Portal: unified model organism research platform. <i>Nucleic Acids Research</i> , 2020, 48, D650-D658.	14.5	145
7	Ensembl Genomes 2020—enabling non-vertebrate genomic research. <i>Nucleic Acids Research</i> , 2020, 48, D689-D695.	14.5	416
8	Genomic and transcriptomic variation defines the chromosome-scale assembly of <i>Haemonchus contortus</i> , a model gastrointestinal worm. <i>Communications Biology</i> , 2020, 3, 656.	4.4	91
9	Nearly Complete Genome Sequence of <i>Brugia malayi</i> Strain FR3. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.6	13
10	Sex chromosome evolution in parasitic nematodes of humans. <i>Nature Communications</i> , 2020, 11, 1964.	12.8	38
11	Ensembl Genomes 2018: an integrated omics infrastructure for non-vertebrate species. <i>Nucleic Acids Research</i> , 2018, 46, D802-D808.	14.5	489
12	WormBase 2017: molting into a new stage. <i>Nucleic Acids Research</i> , 2018, 46, D869-D874.	14.5	172
13	Using WormBase: A Genome Biology Resource for <i>Caenorhabditis elegans</i> and Related Nematodes. <i>Methods in Molecular Biology</i> , 2018, 1757, 399-470.	0.9	28
14	Combining RNA-seq data and homology-based gene prediction for plants, animals and fungi. <i>BMC Bioinformatics</i> , 2018, 19, 189.	2.6	192
15	Ensembl Genomes 2016: more genomes, more complexity. <i>Nucleic Acids Research</i> , 2016, 44, D574-D580.	14.5	530
16	WormBase 2016: expanding to enable helminth genomic research. <i>Nucleic Acids Research</i> , 2016, 44, D774-D780.	14.5	329
17	WormBase 2014: new views of curated biology. <i>Nucleic Acids Research</i> , 2014, 42, D789-D793.	14.5	149
18	Ensembl Genomes 2013: scaling up access to genome-wide data. <i>Nucleic Acids Research</i> , 2014, 42, D546-D552.	14.5	205

#	ARTICLE	IF	CITATIONS
19	WormBase. <i>Worm</i> , 2012, 1, 15-21.	1.0	14
20	Ensembl Genomes: an integrative resource for genome-scale data from non-vertebrate species. <i>Nucleic Acids Research</i> , 2012, 40, D91-D97.	14.5	179
21	WormBase 2012: more genomes, more data, new website. <i>Nucleic Acids Research</i> , 2012, 40, D735-D741.	14.5	175
22	WormBase: a comprehensive resource for nematode research. <i>Nucleic Acids Research</i> , 2010, 38, D463-D467.	14.5	325
23	WormBase: new content and better access. <i>Nucleic Acids Research</i> , 2007, 35, D506-D510.	14.5	80
24	WormBook: the online review of <i>Caenorhabditis elegans</i> biology. <i>Nucleic Acids Research</i> , 2007, 35, D472-D475.	14.5	160
25	WormBase 2007. <i>Nucleic Acids Research</i> , 2007, 36, D612-D617.	14.5	95
26	The H-Invitational Database (H-InvDB), a comprehensive annotation resource for human genes and transcripts. <i>Nucleic Acids Research</i> , 2007, 36, D793-D799.	14.5	57
27	Integrative Annotation of 21,037 Human Genes Validated by Full-Length cDNA Clones. <i>PLoS Biology</i> , 2004, 2, e162.	5.6	290
28	CHRONIC HIP INSTABILITY AS A CAUSE OF AUTONOMIC DYSREFLEXIA: SUCCESSFUL MANAGEMENT BY RESECTION ARTHROPLASTY. <i>Journal of Bone and Joint Surgery - Series A</i> , 2003, 85, 126-128.	3.0	11