

Julie Ahringer

List of Publications by Year in descending order

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83
papers

22,421
citations

44444

50
h-index

75989

78
g-index

96
all docs

96
docs citations

96
times ranked

20736
citing authors

#	ARTICLE	IF	CITATIONS
1	Systematic functional analysis of the <i>Caenorhabditis elegans</i> genome using RNAi. <i>Nature</i> , 2003, 421, 231-237.	13.7	3,343
2	Genes that act downstream of DAF-16 to influence the lifespan of <i>Caenorhabditis elegans</i> . <i>Nature</i> , 2003, 424, 277-283.	13.7	1,998
3	Functional genomic analysis of <i>C. elegans</i> chromosome I by systematic RNA interference. <i>Nature</i> , 2000, 408, 325-330.	13.7	1,655
4	Genome-wide RNAi screening in <i>Caenorhabditis elegans</i> . <i>Methods</i> , 2003, 30, 313-321.	1.9	1,244
5	Rates of Behavior and Aging Specified by Mitochondrial Function During Development. <i>Science</i> , 2002, 298, 2398-2401.	6.0	974
6	Genome-wide RNAi analysis of <i>Caenorhabditis elegans</i> fat regulatory genes. <i>Nature</i> , 2003, 421, 268-272.	13.7	940
7	Effectiveness of specific RNA-mediated interference through ingested double-stranded RNA in <i>Caenorhabditis elegans</i> . <i>Genome Biology</i> , 2000, 2, research0002.1.	13.9	918
8	Integrative Analysis of the <i>Caenorhabditis elegans</i> Genome by the modENCODE Project. <i>Science</i> , 2010, 330, 1775-1787.	6.0	912
9	A systematic RNAi screen identifies a critical role for mitochondria in <i>C. elegans</i> longevity. <i>Nature Genetics</i> , 2003, 33, 40-48.	9.4	900
10	Genetic Analysis of Tissue Aging in <i>Caenorhabditis elegans</i> : A Role for Heat-Shock Factor and Bacterial Proliferation. <i>Genetics</i> , 2002, 161, 1101-1112.	1.2	718
11	Differential chromatin marking of introns and expressed exons by H3K36me3. <i>Nature Genetics</i> , 2009, 41, 376-381.	9.4	592
12	Genome-Wide RNAi of <i>C. elegans</i> Using the Hypersensitive rrf-3 Strain Reveals Novel Gene Functions. <i>PLoS Biology</i> , 2003, 1, e12.	2.6	545
13	Loss of the Putative RNA-Directed RNA Polymerase RRF-3 Makes <i>C. elegans</i> Hypersensitive to RNAi. <i>Current Biology</i> , 2002, 12, 1317-1319.	1.8	529
14	Cell Polarity in Eggs and Epithelia: Parallels and Diversity. <i>Cell</i> , 2010, 141, 757-774.	13.5	430
15	The art and design of genetic screens: RNA interference. <i>Nature Reviews Genetics</i> , 2008, 9, 554-566.	7.7	413
16	The <i>C. elegans</i> Hook Protein, ZYG-12, Mediates the Essential Attachment between the Centrosome and Nucleus. <i>Cell</i> , 2003, 115, 825-836.	13.5	380
17	An assessment of histone-modification antibody quality. <i>Nature Structural and Molecular Biology</i> , 2011, 18, 91-93.	3.6	369
18	Comparative analysis of metazoan chromatin organization. <i>Nature</i> , 2014, 512, 449-452.	13.7	363

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19	NuRD and SIN3. Trends in Genetics, 2000, 16, 351-356.	2.9	356
20	Broad chromosomal domains of histone modification patterns in <i>C. elegans</i> . Genome Research, 2011, 21, 227-236.	2.4	256
21	Distinct roles for G1 and G2 in regulating spindle position and orientation in <i>Caenorhabditis elegans</i> embryos. Nature Cell Biology, 2001, 3, 297-300.	4.6	252
22	Control of the sperm-to-oocyte switch in <i>Caenorhabditis elegans</i> hermaphrodites by the fem-3 3' untranslated region. Nature, 1991, 349, 346-348.	13.7	237
23	Asymmetrically Distributed <i>C. elegans</i> Homologs of AGS3/PINS Control Spindle Position in the Early Embryo. Current Biology, 2003, 13, 1029-1037.	1.8	229
24	CDC-42 controls early cell polarity and spindle orientation in <i>C. elegans</i> . Current Biology, 2001, 11, 482-488.	1.8	205
25	Identification of genes that protect the <i>C. elegans</i> genome against mutations by genome-wide RNAi. Genes and Development, 2003, 17, 443-448.	2.7	196
26	A Genome-Wide Screen Identifies 27 Genes Involved in Transposon Silencing in <i>C. elegans</i> . Current Biology, 2003, 13, 1311-1316.	1.8	180
27	Reverse genetics. WormBook, 2006, , .	5.3	173
28	aPKC Cycles between Functionally Distinct PAR Protein Assemblies to Drive Cell Polarity. Developmental Cell, 2017, 42, 400-415.e9.	3.1	162
29	G Proteins Are Required for Spatial Orientation of Early Cell Cleavages in <i>C. elegans</i> Embryos. Cell, 1996, 86, 619-629.	13.5	153
30	NURD-complex genes antagonise Ras-induced vulval development in <i>Caenorhabditis elegans</i> . Current Biology, 2000, 10, 223-226.	1.8	146
31	Systematic bias in high-throughput sequencing data and its correction by BEADS. Nucleic Acids Research, 2011, 39, e103-e103.	6.5	137
32	Control of cell polarity and mitotic spindle positioning in animal cells. Current Opinion in Cell Biology, 2003, 15, 73-81.	2.6	132
33	SeqPlots - Interactive software for exploratory data analyses, pattern discovery and visualization in genomics. Wellcome Open Research, 2016, 1, 14.	0.9	132
34	Chromatin regulation and sumoylation in the inhibition of Ras-induced vulval development in <i>Caenorhabditis elegans</i> . EMBO Journal, 2005, 24, 2613-2623.	3.5	119
35	TAC-1, a Regulator of Microtubule Length in the <i>C. elegans</i> Embryo. Current Biology, 2003, 13, 1499-1505.	1.8	111
36	Comparative assessment of fluorescent proteins for in vivo imaging in an animal model system. Molecular Biology of the Cell, 2016, 27, 3385-3394.	0.9	108

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37	MosSCI and Gateway Compatible Plasmid Toolkit for Constitutive and Inducible Expression of Transgenes in the <i>C. elegans</i> Germline. PLoS ONE, 2011, 6, e20082.	1.1	94
38	The landscape of RNA polymerase II transcription initiation in <i>C. elegans</i> reveals promoter and enhancer architectures. Genome Research, 2013, 23, 1339-1347.	2.4	89
39	A team of heterochromatin factors collaborates with small RNA pathways to combat repetitive elements and germline stress. ELife, 2017, 6, .	2.8	87
40	PAR proteins direct asymmetry of the cell cycle regulators Polo-like kinase and Cdc25. Journal of Cell Biology, 2008, 180, 877-885.	2.3	84
41	Systematic genetic interaction screens uncover cell polarity regulators and functional redundancy. Nature Cell Biology, 2013, 15, 103-112.	4.6	84
42	Repressive Chromatin in <i>Caenorhabditis elegans</i> : Establishment, Composition, and Function. Genetics, 2018, 208, 491-511.	1.2	82
43	Microtubules are involved in anterior-posterior axis formation in <i>C. elegans</i> embryos. Journal of Cell Biology, 2007, 179, 397-402.	2.3	76
44	A Casein Kinase 1 and PAR Proteins Regulate Asymmetry of a PIP2 Synthesis Enzyme for Asymmetric Spindle Positioning. Developmental Cell, 2008, 15, 198-208.	3.1	76
45	Chromatin accessibility dynamics across <i>C. elegans</i> development and ageing. ELife, 2018, 7, .	2.8	76
46	Stable <i>Caenorhabditis elegans</i> chromatin domains separate broadly expressed and developmentally regulated genes. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E7020-E7029.	3.3	73
47	PRDE-1 is a nuclear factor essential for the biogenesis of Ruby motif-dependent piRNAs in <i>C. elegans</i> . Genes and Development, 2014, 28, 783-796.	2.7	72
48	H4K20me1 Contributes to Downregulation of X-Linked Genes for <i>C. elegans</i> Dosage Compensation. PLoS Genetics, 2012, 8, e1002933.	1.5	70
49	Involvement of fatty acid pathways and cortical interaction of the pronuclear complex in <i>Caenorhabditis elegans</i> embryonic polarity. BMC Developmental Biology, 2003, 3, 8.	2.1	68
50	Similar requirements for CDC-42 and the PAR-3/PAR-6/PKC-3 complex in diverse cell types. Developmental Biology, 2007, 305, 347-357.	0.9	61
51	Extreme HOT regions are CpG-dense promoters in <i>C. elegans</i> and humans. Genome Research, 2014, 24, 1138-1146.	2.4	57
52	Genome-wide RNAi screens in <i>Caenorhabditis elegans</i> : impact on cancer research. Oncogene, 2004, 23, 8340-8345.	2.6	56
53	Physical and functional interaction between SET1/COMPASS complex component CFP-1 and a Sin3S HDAC complex in <i>C. elegans</i> . Nucleic Acids Research, 2019, 47, 11164-11180.	6.5	54
54	Repression by the 3' UTR <i>offm-3</i> , a sex-determining gene, relies on a ubiquitous <i>mog</i> -dependent control in <i>Caenorhabditis elegans</i> . EMBO Journal, 1998, 17, 6337-6347.	3.5	53

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55	The DREAM complex promotes gene body H2A.Z for target repression. <i>Genes and Development</i> , 2015, 29, 495-500.	2.7	53
56	Asymmetry of Early Endosome Distribution in <i>C. elegans</i> Embryos. <i>PLoS ONE</i> , 2007, 2, e493.	1.1	50
57	Duplication and Retention Biases of Essential and Non-Essential Genes Revealed by Systematic Knockdown Analyses. <i>PLoS Genetics</i> , 2013, 9, e1003330.	1.5	48
58	Morphogenetic degeneracies in the actomyosin cortex. <i>ELife</i> , 2018, 7, .	2.8	41
59	A regulatory network of T-box genes and the even-skippedhomologue <i>vab-7</i> controls patterning and morphogenesis in <i>C. elegans</i> . <i>Development (Cambridge)</i> , 2004, 131, 2373-2385.	1.2	40
60	Axis determination in <i>C. elegans</i> : initiating and transducingpolarity. <i>Current Opinion in Genetics and Development</i> , 2001, 11, 367-373.	1.5	39
61	An Alternative STAT Signaling Pathway Acts in Viral Immunity in <i>Caenorhabditis elegans</i> . <i>MBio</i> , 2017, 8, .	1.8	38
62	Comparative Epigenomics Reveals that RNA Polymerase II Pausing and Chromatin Domain Organization Control Nematode piRNA Biogenesis. <i>Developmental Cell</i> , 2019, 48, 793-810.e6.	3.1	37
63	Turn to the worm!. <i>Current Opinion in Genetics and Development</i> , 1997, 7, 410-415.	1.5	36
64	The USTC co-opts an ancient machinery to drive piRNA transcription in <i>C. elegans</i> . <i>Genes and Development</i> , 2019, 33, 90-102.	2.7	34
65	Distinctive regulatory architectures of germline-active and somatic genes in <i>C. elegans</i> . <i>Genome Research</i> , 2020, 30, 1752-1765.	2.4	32
66	Genome organization at different scales: nature, formation and function. <i>Current Opinion in Cell Biology</i> , 2018, 52, 145-153.	2.6	23
67	CDK phosphorylation of SLD-2 is required for replication initiation and germline development in <i>C. elegans</i> . <i>Journal of Cell Biology</i> , 2014, 204, 507-522.	2.3	21
68	The <i>Caenorhabditis elegans</i> homolog of the Evi1 proto-oncogene, <i>egl-43</i> , coordinates G1 cell cycle arrest with pro-invasive gene expression during anchor cell invasion. <i>PLoS Genetics</i> , 2020, 16, e1008470.	1.5	19
69	Broad Chromatin Domains: An Important Facet of Genome Regulation. <i>BioEssays</i> , 2017, 39, 1700124.	1.2	18
70	Identification of the <i>C. elegans</i> anaphase promoting complex subunit <i>Cdc26</i> by phenotypic profiling and functional rescue in yeast. <i>BMC Developmental Biology</i> , 2007, 7, 19.	2.1	11
71	DREAM represses distinct targets by cooperating with different THAP domain proteins. <i>Cell Reports</i> , 2021, 37, 109835.	2.9	6
72	Accessible Region Conformation Capture (ARC-C) gives high-resolution insights into genome architecture and regulation. <i>Genome Research</i> , 2022, 32, 357-366.	2.4	6

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73	periodicDNA: an R/Bioconductor package to investigate k-mer periodicity in DNA. F1000Research, 2021, 10, 141.	0.8	5
74	Determination of Octadecylamine in Water by Quantitative High Performance thin Layer Chromatography. Journal of Liquid Chromatography and Related Technologies, 1984, 7, 2743-2749.	0.9	3
75	The caenorhabditis elegans CDT-2 ubiquitin ligase is required for attenuation of EGFR signalling in vulva precursor cells. BMC Developmental Biology, 2010, 10, 109.	2.1	3
76	Playing Ping Pong with Pins: Cortical and Microtubule-Induced Polarity. Cell, 2005, 123, 1184-1186.	13.5	2
77	Generating fragment density plots in R/Bioconductor with VplotR. Journal of Open Source Software, 2021, 6, 3009.	2.0	2
78	Title is missing!. , 2020, 16, e1008470.		0
79	Title is missing!. , 2020, 16, e1008470.		0
80	Title is missing!. , 2020, 16, e1008470.		0
81	Title is missing!. , 2020, 16, e1008470.		0
82	Title is missing!. , 2020, 16, e1008470.		0
83	Title is missing!. , 2020, 16, e1008470.		0