

Eleanor M Maine

List of Publications by Year in descending order

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Version: 2024-02-01

28
papers

1,255
citations

623188

14
h-index

552369

26
g-index

29
all docs

29
docs citations

29
times ranked

1188
citing authors

#	ARTICLE	IF	CITATIONS
1	Screening by deep sequencing reveals mediators of microRNA tailing in <i>C. elegans</i> . <i>Nucleic Acids Research</i> , 2021, 49, 11167-11180.	6.5	16
2	Meiotic H3K9me2 distribution is influenced by the ALG-3 and ALG-4 pathway and by poly(U) polymerase activity. <i>MicroPublication Biology</i> , 2021, 2021, .	0.1	0
3	A DNA repair protein and histone methyltransferase interact to promote genome stability in the <i>Caenorhabditis elegans</i> germ line. <i>PLoS Genetics</i> , 2019, 15, e1007992.	1.5	19
4	The balance of poly(U) polymerase activity ensures germline identity, survival and development in <i>Caenorhabditis elegans</i> . <i>Development (Cambridge)</i> , 2018, 145, .	1.2	6
5	The Molecular Chaperone HSP90 Promotes Notch Signaling in the Germline of <i>Caenorhabditis elegans</i> . <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 1535-1544.	0.8	4
6	Regulated nuclear accumulation of a histone methyltransferase times the onset of heterochromatin formation in <i>C. elegans</i> embryos. <i>Science Advances</i> , 2018, 4, eaat6224.	4.7	55
7	Simplified detection of a point mutation in using tetra-primer ARMS-PCR. <i>MicroPublication Biology</i> , 2018, 2018, .	0.1	4
8	UBR-5, a Conserved HECT-Type E3 Ubiquitin Ligase, Negatively Regulates Notch-Type Signaling in <i>Caenorhabditis elegans</i> . <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 2125-2134.	0.8	15
9	Fine-scale belowground species associations in temperate grassland. <i>Molecular Ecology</i> , 2015, 24, 3206-3216.	2.0	29
10	Enrichment of H3K9me2 on Unsynapsed Chromatin in <i>Caenorhabditis elegans</i> Does Not Target de Novo Sites. <i>G3: Genes, Genomes, Genetics</i> , 2015, 5, 1865-1878.	0.8	12
11	Using model organisms in an undergraduate laboratory to link genotype, phenotype, and the environment. <i>Journal of Biological Education</i> , 2013, 47, 52-59.	0.8	3
12	Epigenetic Control of Germline Development. <i>Advances in Experimental Medicine and Biology</i> , 2013, 757, 373-403.	0.8	4
13	Cyclin E and CDK-2 regulate proliferative cell fate and cell cycle progression in the <i>C. elegans</i> germline. <i>Development (Cambridge)</i> , 2011, 138, 2223-2234.	1.2	142
14	Developmental Biology: Small RNAs Play Their Part. <i>Current Biology</i> , 2011, 21, R274-R276.	1.8	0
15	Meiotic silencing in <i>Caenorhabditis elegans</i> . <i>International Review of Cell and Molecular Biology</i> , 2010, 282, 91-134.	1.6	24
16	An RNA-Mediated Silencing Pathway Utilizes the Coordinated Synthesis of Two Distinct Populations of siRNA. <i>Molecular Cell</i> , 2010, 37, 593-595.	4.5	1
17	Regulation of Heterochromatin Assembly on Unpaired Chromosomes during <i>Caenorhabditis elegans</i> Meiosis by Components of a Small RNA-Mediated Pathway. <i>PLoS Genetics</i> , 2009, 5, e1000624.	1.5	82
18	Studying gene function in <i>Caenorhabditis elegans</i> using RNA-mediated interference. <i>Briefings in Functional Genomics & Proteomics</i> , 2008, 7, 184-194.	3.8	16

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19	The Bro1-Domain Protein, EGO-2, Promotes Notch Signaling in <i>Caenorhabditis elegans</i> . <i>Genetics</i> , 2007, 176, 2265-2277.	1.2	13
20	Eukaryotic translation initiation factor 5B activity regulates larval growth rate and germline development in <i>Caenorhabditis elegans</i> . <i>Genesis</i> , 2006, 44, 412-418.	0.8	8
21	EGO-1, a Putative RNA-Dependent RNA Polymerase, Is Required for Heterochromatin Assembly on Unpaired DNA during <i>C. elegans</i> Meiosis. <i>Current Biology</i> , 2005, 15, 1972-1978.	1.8	85
22	EGO-1, a Putative RNA-Directed RNA Polymerase, Promotes Germline Proliferation in Parallel With GLP-1/Notch Signaling and Regulates the Spatial Organization of Nuclear Pore Complexes and Germline P Granules in <i>Caenorhabditis elegans</i> . <i>Genetics</i> , 2005, 170, 1121-1132.	1.2	52
23	<i>Caenorhabditis elegans</i> atx-2 Promotes Germline Proliferation and the Oocyte Fate Sequence data from this article have been deposited with the EMBL/GenBank Data Libraries under accession no. AY571963.. <i>Genetics</i> , 2004, 168, 817-830.	1.2	43
24	RNAi As a Tool for Understanding Germline Development in <i>Caenorhabditis elegans</i> : Uses and Cautions. <i>Developmental Biology</i> , 2001, 239, 177-189.	0.9	23
25	EGO-1 is related to RNA-directed RNA polymerase and functions in germ-line development and RNA interference in <i>C. elegans</i> . <i>Current Biology</i> , 2000, 10, 169-178.	1.8	502
26	A conserved mechanism for post-transcriptional gene silencing?. <i>Genome Biology</i> , 2000, 1, reviews1018.1.	13.9	21
27	Carboxy-terminal truncation activates glp-1 protein to specify vulval fates in <i>Caenorhabditis elegans</i> . <i>Nature</i> , 1991, 352, 811-815.	13.7	65
28	Genetic control of cell communication in <i>C. elegans</i> development. <i>BioEssays</i> , 1990, 12, 265-271.	1.2	11