

Eva Jimenez-guri

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

968
citations

516215

16
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610482

24
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28
all docs

28
docs citations

28
times ranked

999
citing authors

#	ARTICLE	IF	CITATIONS
1	Developmental toxicity of plastic leachates on the sea urchin <i>Paracentrotus lividus</i> . <i>Environmental Pollution</i> , 2021, 269, 115744.	3.7	38
2	Transgenerational effects on development following microplastic exposure in <i>Drosophila melanogaster</i> . <i>PeerJ</i> , 2021, 9, e11369.	0.9	20
3	<i>tarsal-less</i> is expressed as a gap gene but has no gap gene phenotype in the moth midge <i>Clogmia albipunctata</i> . <i>Royal Society Open Science</i> , 2018, 5, 180458.	1.1	4
4	A damped oscillator imposes temporal order on posterior gap gene expression in <i>Drosophila</i> . <i>PLoS Biology</i> , 2018, 16, e2003174.	2.6	65
5	Non-canonical dorsoventral patterning in the moth midge <i>Clogmia albipunctata</i> . <i>EvoDevo</i> , 2017, 8, 20.	1.3	7
6	Gap Gene Regulatory Dynamics Evolve along a Genotype Network. <i>Molecular Biology and Evolution</i> , 2016, 33, 1293-1307.	3.5	55
7	High-resolution gene expression data from blastoderm embryos of the scuttle fly <i>Megaselia abdita</i> . <i>Scientific Data</i> , 2015, 2, 150005.	2.4	5
8	SuperFly: a comparative database for quantified spatio-temporal gene expression patterns in early dipteran embryos. <i>Nucleic Acids Research</i> , 2015, 43, D751-D755.	6.5	16
9	Maternal Co-ordinate Gene Regulation and Axis Polarity in the Scuttle Fly <i>Megaselia abdita</i> . <i>PLoS Genetics</i> , 2015, 11, e1005042.	1.5	14
10	Quantitative system drift compensates for altered maternal inputs to the gap gene network of the scuttle fly <i>Megaselia abdita</i> . <i>ELife</i> , 2015, 4, .	2.8	68
11	Rearing the scuttle fly <i>Megaselia scalaris</i> (Diptera: Phoridae) on industrial compounds: implications on size and lifespan. <i>PeerJ</i> , 2015, 3, e1085.	0.9	4
12	A Staging Scheme for the Development of the Scuttle Fly <i>Megaselia abdita</i> . <i>PLoS ONE</i> , 2014, 9, e84421.	1.1	26
13	A Staging Scheme for the Development of the Moth Midge <i>Clogmia albipunctata</i> . <i>PLoS ONE</i> , 2014, 9, e84422.	1.1	28
14	A quantitative atlas of Even-skipped and Hunchback expression in <i>Clogmia albipunctata</i> (Diptera: Phoridae). <i>Development</i> , 2014, 141, 1075-1085.	1.3	57
15	Comparative transcriptomics of early dipteran development. <i>BMC Genomics</i> , 2013, 14, 123.	1.2	41
16	Evolution and expression of BMP genes in flies. <i>Development Genes and Evolution</i> , 2013, 223, 335-340.	0.4	12
17	An ancient mechanism of hindbrain patterning has been conserved in vertebrate evolution. <i>Evolution & Development</i> , 2011, 13, 38-46.	1.1	17
18	Clonal Analysis in Mice Underlines the Importance of Rhombomeric Boundaries in Cell Movement Restriction during Hindbrain Segmentation. <i>PLoS ONE</i> , 2010, 5, e10112.	1.1	37

#	ARTICLE	IF	CITATIONS
19	Origin and evolution of a myxozoan worm. <i>Integrative and Comparative Biology</i> , 2007, 47, 752-758.	0.9	19
20	<i>Buddenbrockia</i> Is a Cnidarian Worm. <i>Science</i> , 2007, 317, 116-118.	6.0	151
21	High nucleotide divergence in a dimorphic parasite with disparate hosts. <i>Zootaxa</i> , 2007, 1636, 59-68.	0.2	3
22	Hox and ParaHox genes in Nemertodermatida, a basal bilaterian clade. <i>International Journal of Developmental Biology</i> , 2006, 50, 675-679.	0.3	40
23	The Hox gene complement of acoel flatworms, a basal bilaterian clade. <i>Evolution & Development</i> , 2004, 6, 154-163.	1.1	103
24	Expression of the 22 nucleotide let-7 heterochronic RNA throughout the Metazoa: a role in life history evolution?. <i>Evolution & Development</i> , 2003, 5, 372-378.	1.1	130