

Suzanne E Mcgaugh

List of Publications by Year in descending order

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Version: 2024-02-01

31
papers

1,262
citations

516215

16
h-index

500791

28
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41
all docs

41
docs citations

41
times ranked

1742
citing authors

#	ARTICLE	IF	CITATIONS
1	Hybridization underlies localized trait evolution in cavefish. <i>IScience</i> , 2022, 25, 103778.	1.9	17
2	CaveCrawler: an interactive analysis suite for cavefish bioinformatics. <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	0.8	0
3	A chromosome-level genome of <i>Astyanax mexicanus</i> surface fish for comparing population-specific genetic differences contributing to trait evolution. <i>Nature Communications</i> , 2021, 12, 1447.	5.8	60
4	Whole-genome variation of transposable element insertions in a maize diversity panel. <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	0.8	16
5	Repeated evolution of circadian clock dysregulation in cavefish populations. <i>PLoS Genetics</i> , 2021, 17, e1009642.	1.5	29
6	Pleiotropic function of the <i>oca2</i> gene underlies the evolution of sleep loss and albinism in cavefish. <i>Current Biology</i> , 2021, 31, 3694-3701.e4.	1.8	30
7	The utility of genomic prediction models in evolutionary genetics. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2021, 288, 20210693.	1.2	13
8	It's time to stop sweeping recombination rate under the genome scan rug. <i>Molecular Ecology</i> , 2020, 29, 4249-4253.	2.0	14
9	Dark world rises: The emergence of cavefish as a model for the study of evolution, development, behavior, and disease. <i>Journal of Experimental Zoology Part B: Molecular and Developmental Evolution</i> , 2020, 334, 397-404.	0.6	31
10	Unique transcriptional signatures of sleep loss across independently evolved cavefish populations. <i>Journal of Experimental Zoology Part B: Molecular and Developmental Evolution</i> , 2020, 334, 497-510.	0.6	11
11	Using multiple reference genomes to identify and resolve annotation inconsistencies. <i>BMC Genomics</i> , 2020, 21, 281.	1.2	10
12	An Adult Brain Atlas Reveals Broad Neuroanatomical Changes in Independently Evolved Populations of Mexican Cavefish. <i>Frontiers in Neuroanatomy</i> , 2019, 13, 88.	0.9	36
13	Stable transgenesis in <i>Astyanax mexicanus</i> using the <i>Tol2</i> transposase system. <i>Developmental Dynamics</i> , 2019, 248, 679-687.	0.8	57
14	Contrasting Patterns of Rapid Molecular Evolution within the <i>p53</i> Network across Mammal and Sauropsid Lineages. <i>Genome Biology and Evolution</i> , 2019, 11, 629-643.	1.1	7
15	Nonrandom RNAseq gene expression associated with RNAlater and flash freezing storage methods. <i>Molecular Ecology Resources</i> , 2019, 19, 456-464.	2.2	31
16	A local duplication of the Melanocortin receptor 1 locus in <i>Astyanax</i> . <i>Genome</i> , 2018, 61, 254-265.	0.9	5
17	The limited role of differential fractionation in genome content variation and function in maize (<i>Zea mays</i> L.) inbred lines. <i>Plant Journal</i> , 2018, 93, 131-141.	2.8	42
18	The role of gene flow in rapid and repeated evolution of cave-related traits in Mexican tetra, <i>Astyanax mexicanus</i> . <i>Molecular Ecology</i> , 2018, 27, 4397-4416.	2.0	160

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19	Tandem Duplicate Genes in Maize Are Abundant and Date to Two Distinct Periods of Time. <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 3049-3058.	0.8	13
20	The evolution of a series of behavioral traits is associated with autism-risk genes in cavefish. <i>BMC Evolutionary Biology</i> , 2018, 18, 89.	3.2	29
21	Growth and stress response mechanisms underlying post-feeding regenerative organ growth in the Burmese python. <i>BMC Genomics</i> , 2017, 18, 338.	1.2	32
22	Righting ability in hatchling turtles does not predict survival during dispersal in the field. <i>Biological Journal of the Linnean Society</i> , 2016, , .	0.7	5
23	Mapping the Genetic Basis of Troglomorphy in <i>Astyanax</i> . , 2016, , 111-135.		11
24	Rapid molecular evolution across amniotes of the IIS/TOR network. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 7055-7060.	3.3	59
25	The cavefish genome reveals candidate genes for eye loss. <i>Nature Communications</i> , 2014, 5, 5307.	5.8	256
26	Population genetics of Blanding's turtle (<i>Emys blandingii</i>) in the midwestern United States. <i>Conservation Genetics</i> , 2014, 15, 61-73.	0.8	14
27	Recombination Modulates How Selection Affects Linked Sites in <i>Drosophila</i> . <i>PLoS Biology</i> , 2012, 10, e1001422.	2.6	104
28	Genomic impacts of chromosomal inversions in parapatric <i>Drosophila</i> species. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2012, 367, 422-429.	1.8	111
29	Comparative population genetics of aquatic turtles in the desert. <i>Conservation Genetics</i> , 2012, 13, 1561-1576.	0.8	14
30	Color Variation among Habitat Types in the Spiny Softshell Turtles (<i>Trionychidae: Apalone</i>) of Cuatrociénegas, Coahuila, Mexico. <i>Journal of Herpetology</i> , 2008, 42, 347-353.	0.2	23
31	Evidence for rapid phenotypic and behavioural shifts in a recently established cavefish population. <i>Biological Journal of the Linnean Society</i> , 0, , .	0.7	5