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List of Publications by Year in descending order

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Version: 2024-02-01

19
papers

442
citations

759190

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794568

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times ranked

582
citing authors

#	ARTICLE	IF	CITATIONS
1	Signatures of Divergence, Invasiveness, and Terrestrialization Revealed by Four Apple Snail Genomes. <i>Molecular Biology and Evolution</i> , 2019, 36, 1507-1520.	8.9	65
2	The Scaly-foot Snail genome and implications for the origins of biomineralised armour. <i>Nature Communications</i> , 2020, 11, 1657.	12.8	64
3	Host-Endosymbiont Genome Integration in a Deep-Sea Chemosymbiotic Clam. <i>Molecular Biology and Evolution</i> , 2021, 38, 502-518.	8.9	46
4	De novo transcriptome analysis of <i>Perna viridis</i> highlights tissue-specific patterns for environmental studies. <i>BMC Genomics</i> , 2014, 15, 804.	2.8	38
5	Genomic Signatures Supporting the Symbiosis and Formation of Chitinous Tube in the Deep-Sea Tubeworm <i>Paraescarpia echinospica</i> . <i>Molecular Biology and Evolution</i> , 2021, 38, 4116-4134.	8.9	37
6	Developmental toxicity and molecular responses of marine medaka (<i>Oryzias melastigma</i>) embryos to ciguatoxin P-CTX-1 exposure. <i>Aquatic Toxicology</i> , 2017, 185, 149-159.	4.0	27
7	Photosynthetic and transcriptional responses of the marine diatom <i>Thalassiosira pseudonana</i> to the combined effect of temperature stress and copper exposure. <i>Marine Pollution Bulletin</i> , 2017, 124, 938-945.	5.0	26
8	Molecular phylogeny and toxicity of harmful benthic dinoflagellates <i>Coolia</i> (Ostreopsidaceae). <i>Marine Pollution Bulletin</i> , 2017, 124, 878-889.	5.0	24
9	AmpuBase: a transcriptome database for eight species of apple snails (Gastropoda: Ampullariidae). <i>BMC Genomics</i> , 2018, 19, 179.	2.8	20
10	Multi-omic approach provides insights into osmoregulation and osmoconformation of the crab <i>Scylla paramamosain</i> . <i>Scientific Reports</i> , 2020, 10, 21771.	3.3	19
11	De novo transcriptome assembly of the marine gastropod <i>Reishia clavigera</i> for supporting toxic mechanism studies. <i>Aquatic Toxicology</i> , 2016, 178, 39-48.	4.0	15
12	De novo transcriptomic profile in the gonadal tissues of the intertidal whelk <i>Reishia clavigera</i> . <i>Marine Pollution Bulletin</i> , 2014, 85, 499-504.	5.0	14
13	Understanding the transition from water to land: Insights from multi-omic analyses of the perivitelline fluid of apple snail eggs. <i>Journal of Proteomics</i> , 2019, 194, 79-88.	2.4	11
14	Host-symbiont transcriptomic changes during natural bleaching and recovery in the leaf coral <i>Pavona decussata</i> . <i>Science of the Total Environment</i> , 2022, 806, 150656.	8.0	10
15	Genomic insights into the sessile life and biofouling of barnacles (Crustacea: Cirripedia). <i>Heliyon</i> , 2021, 7, e07291.	3.2	7
16	Transcriptomics reveal triphenyltin-induced molecular toxicity in the marine mussel <i>Perna viridis</i> . <i>Science of the Total Environment</i> , 2021, 790, 148040.	8.0	7
17	Egg perivitelline fluid proteome of a freshwater snail: Insight into the transition from aquatic to terrestrial egg deposition. <i>Rapid Communications in Mass Spectrometry</i> , 2020, 34, e8605.	1.5	5
18	Molecular analyses revealed three morphologically similar species of non-native apple snails and their patterns of distribution in freshwater wetlands of Hong Kong. <i>Diversity and Distributions</i> , 2022, 28, 97-111.	4.1	4

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19	Insights into the Influence of Natural Retinoic Acids on Imposex Induction in Female Marine Gastropods in the Coastal Environment. <i>Environmental Science and Technology Letters</i> , 2021, 8, 1002-1008.	8.7	3