

# Jack C H Ip

## List of Publications by Year in descending order

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Version: 2024-02-01

19  
papers

442  
citations

759233

12  
h-index

794594

19  
g-index

19  
all docs

19  
docs citations

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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

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