

Mailie L Gall

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

843
citations

1040056

9
h-index

1058476

14
g-index

18
all docs

18
docs citations

18
times ranked

1923
citing authors

#	ARTICLE	IF	CITATIONS
1	Staying in place and moving in space: contrasting larval thermal sensitivity explains distributional changes of sympatric sea urchin species to habitat warming. <i>Global Change Biology</i> , 2022, , .	9.5	9
2	Co-infection with SARS-CoV-2 Omicron and Delta variants revealed by genomic surveillance. <i>Nature Communications</i> , 2022, 13, 2745.	12.8	64
3	SARS-CoV-2 Within-Host and in vitro Genomic Variability and Sub-Genomic RNA Levels Indicate Differences in Viral Expression Between Clinical Cohorts and in vitro Culture. <i>Frontiers in Microbiology</i> , 2022, 13, .	3.5	4
4	Effects of marine heatwave conditions across the metamorphic transition to the juvenile sea urchin (<i>Heliocidaris erythrogramma</i>). <i>Marine Pollution Bulletin</i> , 2021, 163, 111914.	5.0	13
5	Tracking the international spread of SARS-CoV-2 lineages B.1.1.7 and B.1.351/501Y-V2. <i>Wellcome Open Research</i> , 2021, 6, 121.	1.8	115
6	Tracking the international spread of SARS-CoV-2 lineages B.1.1.7 and B.1.351/501Y-V2 with grinch. <i>Wellcome Open Research</i> , 2021, 6, 121.	1.8	129
7	Documenting elimination of co-circulating COVID-19 clusters using genomics in New South Wales, Australia. <i>BMC Research Notes</i> , 2021, 14, 415.	1.4	4
8	Added Value of Genomic Surveillance of Virulence Factors in Shiga Toxin-Producing <i>Escherichia coli</i> in New South Wales, Australia. <i>Frontiers in Microbiology</i> , 2021, 12, 713724.	3.5	2
9	Revealing COVID-19 transmission in Australia by SARS-CoV-2 genome sequencing and agent-based modeling. <i>Nature Medicine</i> , 2020, 26, 1398-1404.	30.7	283
10	An emergent clade of SARS-CoV-2 linked to returned travellers from Iran. <i>Virus Evolution</i> , 2020, 6, veaa027.	4.9	119
11	Diet-induced shifts in the crown-of-thorns (<i>Acanthaster</i> sp.) larval microbiome. <i>Marine Biology</i> , 2018, 165, 1.	1.5	28
12	From pole to pole: the potential for the Arctic seastar <i>Asterias amurensis</i> to invade a warming Southern Ocean. <i>Global Change Biology</i> , 2016, 22, 3874-3887.	9.5	35
13	Characterisation of thirteen polymorphic microsatellite markers for the red sea urchin <i>Heliocidaris tuberculata</i> (Lamarck, 1816) developed using a 454-sequencing approach. <i>Conservation Genetics Resources</i> , 2014, 6, 237-239.	0.8	0
14	Differential tolerance to copper, but no evidence of population-level genetic differences in a widely-dispersing native barnacle. <i>Ecotoxicology</i> , 2013, 22, 929-937.	2.4	7
15	A biomonitor as a measure of an ecologically-significant fraction of metals in an industrialized harbour. <i>Journal of Environmental Monitoring</i> , 2012, 14, 830.	2.1	12