Chris J Brauer

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/8602436/publications.pdf

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19	522	840776 11	839539 18
papers	citations	h-index	g-index

25	25	25	596
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Longitudinal monitoring of neutral and adaptive genomic diversity in a reintroduction. Conservation Biology, 2022, 36, .	4.7	6
2	Seascape genomics of coastal bottlenose dolphins along strong gradients of temperature and salinity. Molecular Ecology, 2022, 31, 2223-2241.	3.9	14
3	Variation in intraspecific demography drives localised concordance but species-wide discordance in response to past climatic change. Bmc Ecology and Evolution, 2022, 22, 35.	1.6	2
4	Fish out of water: Genomic insights into persistence of rainbowfish populations in the desert. Evolution; International Journal of Organic Evolution, 2022, 76, 171-183.	2.3	10
5	The roles of aridification and sea level changes in the diversification and persistence of freshwater fish lineages. Molecular Ecology, 2021, 30, 4866-4883.	3.9	10
6	Latitudinal variation in climateâ€associated genes imperils range edge populations. Molecular Ecology, 2020, 29, 4337-4349.	3.9	12
7	Recent and rapid anthropogenic habitat fragmentation increases extinction risk for freshwater biodiversity. Evolutionary Applications, 2020, 13, 2857-2869.	3.1	43
8	Adaptation of plasticity to projected maximum temperatures and across climatically defined bioregions. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 17112-17121.	7.1	44
9	Ecological disturbance influences adaptive divergence despite high gene flow in golden perch (Macquaria ambigua): Implications for management and resilience to climate change. Molecular Ecology, 2018, 27, 196-215.	3.9	24
10	Phylogenomic history of enigmatic pygmy perches: implications for biogeography, taxonomy and conservation. Royal Society Open Science, 2018, 5, 172125.	2.4	17
11	On the roles of landscape heterogeneity and environmental variation in determining population genomic structure in a dendritic system. Molecular Ecology, 2018, 27, 3484-3497.	3.9	52
12	Comparative ecological transcriptomics and the contribution of gene expression to the evolutionary potential of a threatened fish. Molecular Ecology, 2017, 26, 6841-6856.	3.9	30
13	<scp>swinger</scp>: a userâ€friendly computer program to establish captive breeding groups that minimize relatedness without pedigree information. Molecular Ecology Resources, 2017, 17, 278-287.	4.8	15
14	Range-wide fragmentation in a threatened fish associated with post-European settlement modification in the Murrayâ€“Darling Basin, Australia. Conservation Genetics, 2016, 17, 1377-1391.	1.5	29
15	Riverscape genomics of a threatened fish across a hydroclimatically heterogeneous river basin. Molecular Ecology, 2016, 25, 5093-5113.	3.9	91
16	Multi-generational evaluation of genetic diversity and parentage in captive southern pygmy perch (Nannoperca australis). Conservation Genetics, 2016, 17, 1469-1473.	1.5	9
17	A novel holistic framework for geneticâ€based captiveâ€breeding and reintroduction programs. Conservation Biology, 2016, 30, 1060-1069.	4.7	75
18	Catchment-Scale Conservation Units Identified for the Threatened Yarra Pygmy Perch (Nannoperca) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	2.5	29

#	Article	IF	Citations
19	Fisheries genomics of snapper (Chrysophrys auratus) along the west Australian coast. Evolutionary Applications, 0, , .	3.1	6