

Peter T Euclide

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

278
citations

1040018

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h-index

996954

15
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22
all docs

22
docs citations

22
times ranked

518
citing authors

#	ARTICLE	IF	CITATIONS
1	Characterization of a Yâ€specific duplication/insertion of the antiâ€Mullerian hormone type II receptor gene based on a chromosomeâ€scale genome assembly of yellow perch, <i>Perca flavescens</i>. <i>Molecular Ecology Resources</i> , 2020, 20, 531-543.	4.8	76
2	Building a global genomics observatory: Using GEOME (the Genomic Observatories Metadatabase) to expedite and improve deposition and retrieval of genetic data and metadata for biodiversity research. <i>Molecular Ecology Resources</i> , 2020, 20, 1458-1469.	4.8	32
3	Trends in <i>Mysis diluviana</i> abundance in the Great Lakes, 2006â€2016. <i>Journal of Great Lakes Research</i> , 2018, 44, 590-599.	1.9	20
4	Mixedâ€stock analysis using Rapture genotyping to evaluate stockâ€specific exploitation of a walleye population despite weak genetic structure. <i>Evolutionary Applications</i> , 2021, 14, 1403-1420.	3.1	19
5	Partial diel vertical migration in an omnivorous macroinvertebrate, <i>Mysis diluviana</i> . <i>Hydrobiologia</i> , 2017, 787, 387-396.	2.0	18
6	<sc>RAD</sc>â€Seq Refines Previous Estimates of Genetic Structure in Lake Erie Walleye. <i>Transactions of the American Fisheries Society</i> , 2020, 149, 159-173.	1.4	17
7	Environmental DNA metabarcoding as a tool for biodiversity assessment and monitoring: reconstructing established fish communities of northâ€temperate lakes and rivers. <i>Diversity and Distributions</i> , 2021, 27, 1966-1980.	4.1	17
8	Attack of the PCR clones: Rates of clonality have little effect on RADâ€seq genotype calls. <i>Molecular Ecology Resources</i> , 2020, 20, 66-78.	4.8	16
9	The ghosts of propagation past: haplotype information clarifies the relative influence of stocking history and phylogeographic processes on contemporary population structure of walleye (<i>Sander</i> Tj ETQq1 1 0:3:84314 rgBT /Ove		
10	Genetic Structure of Smallmouth Bass in the Lake Michigan and Upper Mississippi River Drainages Relates to Habitat, Distance, and Drainage Boundaries. <i>Transactions of the American Fisheries Society</i> , 2020, 149, 383-397.	1.4	8
11	Further evidence from common garden rearing experiments of heritable traits separating lean and siscowet lake charr (<i>Salvelinus namaycush</i>) ecotypes. <i>Molecular Ecology</i> , 2022, 31, 3432-3450.	3.9	7
12	Genetic versus demographic stock structure of rainbow smelt in a large fragmented lake. <i>Journal of Great Lakes Research</i> , 2020, 46, 622-632.	1.9	6
13	A chromosomal inversion may facilitate adaptation despite periodic gene flow in a freshwater fish. <i>Ecology and Evolution</i> , 2022, 12, e8898.	1.9	6
14	Lack of genetic population structure of slimy sculpin in a large, fragmented lake. <i>Ecology of Freshwater Fish</i> , 2018, 27, 699-709.	1.4	5
15	Genetic diversity and structure of lake whitefish (<i>Coregonus clupeaformis</i>) 100â€years after closure of the commercial fishery. <i>Journal of Great Lakes Research</i> , 2019, 45, 1310-1319.	1.9	4
16	Role of drainage and barriers in the genetic structuring of a tessellated darter population. <i>Conservation Genetics</i> , 2018, 19, 1379-1392.	1.5	3
17	A Synthesis of the Biology and Ecology of Sculpin Species in the Laurentian Great Lakes and Implications for the Adaptive Capacity of the Benthic Ecosystem. <i>Reviews in Fisheries Science and Aquaculture</i> , 2021, 29, 96-121.	9.1	3
18	Genome-wide genetic diversity may help identify fine-scale genetic structure among lake whitefish spawning groups in Lake Erie. <i>Journal of Great Lakes Research</i> , 2022, 48, 1298-1305.	1.9	2

#	ARTICLE	IF	CITATIONS
19	Using Genomic Data to Guide Walleye Management in the Great Lakes. , 2021, , 115-139.		1
20	Effects of gut content on $\delta^{15}\text{N}$, $\delta^{13}\text{C}$ and C:N of the macroinvertebrate <i>Mysis diluviana</i> . Journal of Great Lakes Research, 2015, 41, 926-929.	1.9	0