Kentaro Inoue

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Phylogenetic and morphometric analyses reveal ecophenotypic plasticity in freshwater mussels <i><scp>O</scp>bovaria jacksoniana</i> and <i><scp>V</scp>illosa arkansasensis</i> (<scp>B</scp> ivalvia: <scp>U</scp> nionidae). Ecology and Evolution, 2013, 3, 2670-2683.	0.8	62
2	Predicting the effects of climate change on population connectivity and genetic diversity of an imperiled freshwater mussel, <i>Cumberlandia monodonta</i> (Bivalvia: Margaritiferidae), in riverine systems. Global Change Biology, 2017, 23, 94-107.	4.2	48
3	Phylogeographic and population genetic analyses reveal Pleistocene isolation followed by high gene flow in a wide ranging, but endangered, freshwater mussel. Heredity, 2014, 112, 282-290.	1.2	47
4	Integrative taxonomy reveals a new species of freshwater mussel, <i>Potamilus streckersoni</i> sp. nov. (Bivalvia: Unionidae): implications for conservation and management. Systematics and Biodiversity, 2019, 17, 331-348.	0.5	34
5	Molecular phylogenetics and morphological variation reveal recent speciation in freshwater mussels of the genera <i>Arcidens</i> and <i>Arkansia</i> (Bivalvia: Unionidae). Biological Journal of the Linnean Society, 2014, 112, 535-545.	0.7	33
6	Joint species models reveal the effects of environment on community assemblage of freshwater mussels and fishes in European rivers. Diversity and Distributions, 2017, 23, 284-296.	1.9	33
7	Past climate change drives current genetic structure of an endangered freshwater mussel species. Molecular Ecology, 2015, 24, 1910-1926.	2.0	32
8	Molecular and morphometric analyses reveal cryptic diversity within freshwater mussels (Bivalvia:) Tj ETQq0 0 0 r Society, 2018, 124, 261-277.	gBT /Over 0.7	ock 10 Tf 50 30
9	Effects of land use on population presence and genetic structure of an amphibian in an agricultural landscape. Landscape Ecology, 2017, 32, 147-162.	1.9	26
10	Longâ€ŧerm markâ€andâ€recapture study of a freshwater mussel reveals patterns of habitat use and an association between survival and river discharge. Freshwater Biology, 2014, 59, 1872-1883.	1.2	23
11	A comprehensive approach uncovers hidden diversity in freshwater mussels (Bivalvia: Unionidae) with the description of a novel species. Cladistics, 2020, 36, 88-113.	1.5	23
12	The Pleurobemini (Bivalvia : Unionida) revisited: molecular species delineation using a mitochondrial DNA gene reveals multiple conspecifics and undescribed species. Invertebrate Systematics, 2018, 32, 689.	0.5	21
13	Isolation drives increased diversification rates in freshwater amphipods. Molecular Phylogenetics and Evolution, 2018, 127, 746-757.	1.2	17
14	High genetic diversity and low differentiation in North American Margaritifera margaritifera (Bivalvia: Unionida: Margaritiferidae). Biological Journal of the Linnean Society, 2018, 123, 850-863.	0.7	16
15	A spatially explicit approach to prioritize protection areas for endangered freshwater mussels. Aquatic Conservation: Marine and Freshwater Ecosystems, 2019, 29, 12-23.	0.9	16
16	A semi-arid river in distress: Contributing factors and recovery solutions for three imperiled freshwater mussels (Family Unionidae) endemic to the Rio Grande basin in North America. Science of the Total Environment, 2018, 631-632, 733-744.	3.9	15
17	Quantifying morphological and genetic variation of sympatric populations to guide conservation of endangered, microâ€endemic springsnails. Aquatic Conservation: Marine and Freshwater Ecosystems, 2014, 24, 536-545.	0.9	14
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Use of species delimitation approaches to assess biodiversity in freshwater planaria (Platyhelminthes,) Tj ETQq0 0 0 rgBT /Overlock 10 Ti 18 209-218.

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19	A new species of freshwater mussel in the genus Popenaias Frierson, 1927, from the Gulf coastal rivers of central Mexico (Bivalvia: Unionida: Unionidae)Âwith comments on the genus. Zootaxa, 2020, 4816, zootaxa.4816.4.3.	0.2	6
20	Development and characterization of 20 polymorphic microsatellite markers for the Texas hornshell, Popenaias popeii (Bivalvia: Unionidae), through next-generation sequencing. Conservation Genetics Resources, 2013, 5, 195-198.	0.4	5
21	<i>Identification of microsatellite loci and examination of genetic structure for the endangered springsnails</i> Juturnia kosteri <i>and</i> Pyrgulopsis roswellensis <i>in the Chihuahuan Desert</i> . Aquatic Conservation: Marine and Freshwater Ecosystems, 2016, 26, 715-723.	0.9	5
22	Misidentification of sex for Lampsilis teres, Yellow Sandshell, and its implications for mussel conservation and wildlife management. PLoS ONE, 2018, 13, e0197107.	1.1	5
23	Identification and characterization of 12 microsatellite loci for Physa in the Chihuahuan Desert. Conservation Genetics Resources, 2014, 6, 769-771.	0.4	3
24	Genetic structuring in the Pyramid Elimia, Elimia potosiensis (Gastropoda, Pleuroceridae), with implications for pleurocerid conservation. Zoosystematics and Evolution, 2017, 93, 437-449.	0.4	3
25	Isolation and characterization of 17 polymorphic microsatellite loci in the spectaclecase, Cumberlandia monodonta (Bivalvia: Margaritiferidae). Conservation Genetics Resources, 2011, 3, 57-60.	0.4	2
26	A comprehensive status, phylogenetic, and anatomical review of <scp><i>Stagnicola caperata</i></scp> (Say, 1829) in the southâ€west United States. Aquatic Conservation: Marine and Freshwater Ecosystems, 2018, 28, 527-534.	0.9	2
27	Range-wide Microsatellite Analysis of the Genetic Population Structure of Prairie Voles (Microtus) Tj ETQq1 1 0.78	34314 rgB 0.2	T 10verlock
28	New microsatellite markers for Ellipse, Venustaconcha ellipsiformis (Bivalvia: Unionidae), with notes on optimal sample size and cross-species amplification. Molecular Biology Reports, 2021, 48, 3037-3045.	1.0	1
29	STATUS OF FRESHWATER MUSSELS (UNIONIDAE) OF THE RÃO CONCHOS BASIN, CHIHUAHUA, MEXICO. Southwestern Naturalist, 2020, 64, .	0.1	0