Shigeru Kitanishi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1451649/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Habitat niche separation of the nonnative rainbow trout and native masu salmon in the Atsuta River, Hokkaido, Japan. Fisheries Science, 2010, 76, 251-256.	1.6	20
2	Patterns of genetic diversity of mitochondrial DNA within captive populations of the endangered itasenpara bitterling: implications for a reintroduction program. Environmental Biology of Fishes, 2013, 96, 567-572.	1.0	18
3	Influences of habitat fragmentation by damming on the genetic structure of masu salmon populations in Hokkaido, Japan. Conservation Genetics, 2012, 13, 1017-1026.	1.5	16
4	Phylogeography of Opsariichthys platypus in Japan based on mitochondrial DNA sequences. Ichthyological Research, 2016, 63, 506-518.	0.8	16
5	Fine scale relationships between sex, life history, and dispersal of masu salmon. Ecology and Evolution, 2012, 2, 920-929.	1.9	13
6	Strong population genetic structure and its implications for the conservation and management of the endangered Itasenpara bitterling. Conservation Genetics, 2013, 14, 901-906.	1.5	9
7	Abiotic factors associated with the occurrence of introduced rainbow trout in the Atsuta River. Ichthyological Research, 2010, 57, 305-309.	0.8	8
8	Relationships between body size and secondary sexual characters, and sperm characters in male Dolly Varden char (<i>Salvelinus malma</i>). Ecology of Freshwater Fish, 2017, 26, 397-402.	1.4	8
9	Comparison of genetic structure between juvenile and adult masu salmon indicates relatively low reproductive success of dispersers. Environmental Biology of Fishes, 2015, 98, 405-411.	1.0	6
10	Dispersal patterns of anadromous and freshwater resident masu salmon at different spatial scales in mid-western Hokkaido, Japan. Ichthyological Research, 2017, 64, 111-115.	0.8	6
11	Shortâ€ŧerm temporal instability in fineâ€scale genetic structure of masu salmon. Freshwater Biology, 2017, 62, 1655-1664.	2.4	6
12	Spatial distribution of juvenile masu salmon (Oncorhynchus masou) with incongruent genotypic and phenotypic sex in Hokkaido, Japan. Environmental Biology of Fishes, 2012, 95, 399-405.	1.0	5
13	Hierarchical genetic structure of native masu salmon populations in Hokkaido, Japan. Environmental Biology of Fishes, 2018, 101, 699-710.	1.0	5
14	Genetic introgression between masu salmon <i>Oncorhynchus masou masou</i> and red spotted masu salmon <i>Oncorhynchus masou ishikawae</i> . Nippon Suisan Gakkaishi, 2017, 83, 400-402.	0.1	5
15	The effects of severe flooding on native masu salmon and nonnative rainbow trout in the Atsuta River, Hokkaido, Japan. Journal of Freshwater Ecology, 2015, 30, 589-596.	1.2	4
16	A simple SNP genotyping method reveals extreme invasions of non-native haplotypes in pale chub Opsariichthys platypus, a common cyprinid fish in Japan. PLoS ONE, 2018, 13, e0191731.	2.5	4
17	Genetic population structure of the eight-barbel loach Lefua echigonia in the Ise Bay region, a single paleo-river basin in central Honshu, Japan. Ichthyological Research, 2019, 66, 411-416.	0.8	3
18	Effect of parental phenotype on dispersal, growth and maturation of offspring in wild masu salmon (Oncorhynchus masou). Evolutionary Ecology, 2021, 35, 253-269.	1.2	3

#	Article	IF	CITATIONS
19	Isolation and characterisation of 23 microsatellite markers for marbled sole, Pleuronectes yokohamae. Conservation Genetics Resources, 2014, 6, 951-953.	0.8	1
20	Links between paternal life history and offspring metabolic rate and body size during early life in masu salmon Oncorhynchus masou. Ecology of Freshwater Fish, 2021, 30, 296-305.	1.4	1
21	Effect of paternal life-history form and egg size on offspring life-history traits in masu salmon Oncorhynchus masou. Hydrobiologia, 0, , .	2.0	1