

Satoshi Yamamoto

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

1,629
citations

566801

15
h-index

580395

25
g-index

25
all docs

25
docs citations

25
times ranked

1443
citing authors

#	ARTICLE	IF	CITATIONS
1	Environmental DNA emission by two carangid fishes in single and mixed-species tanks. <i>Fisheries Science</i> , 2022, 88, 55-62.	0.7	4
2	Estimating fish population abundance by integrating quantitative data on environmental DNA and hydrodynamic modelling. <i>Molecular Ecology</i> , 2021, 30, 3057-3067.	2.0	50
3	Characterizing the spatial and temporal occurrence patterns of the endangered botiid loach <i>Parabotia curtus</i> by environmental DNA analysis using a newly developed species-specific primer set. <i>Ichthyological Research</i> , 2021, 68, 152-157.	0.5	7
4	An illustrated manual for environmental DNA research: Water sampling guidelines and experimental protocols. <i>Environmental DNA</i> , 2021, 3, 8-13.	3.1	102
5	Compilation of real-time PCR conditions toward the standardization of environmental DNA methods. <i>Ecological Research</i> , 2021, 36, 379-388.	0.7	14
6	eDNA as a tool for non-invasive monitoring of the fauna of a turbid, well-mixed system, the Elbe estuary in Germany. <i>PLoS ONE</i> , 2021, 16, e0250452.	1.1	12
7	Population abundance gradient of <i>Inurois punctigera</i> along altitude. <i>Entomological Science</i> , 2020, 23, 23-27.	0.3	1
8	Detection of herbivory: eDNA detection from feeding marks on leaves. <i>Environmental DNA</i> , 2020, 2, 627-634.	3.1	13
9	Sedimentary eDNA provides different information on timescale and fish species composition compared with aqueous eDNA. <i>Environmental DNA</i> , 2020, 2, 505-518.	3.1	77
10	Estimations of Riverine Distribution, Abundance, and Biomass of Anguillid Eels in Japan and Taiwan Using Environmental DNA Analysis. <i>Zoological Studies</i> , 2020, 59, e17.	0.3	9
11	Biomass-dependent emission of environmental DNA in jack mackerel <i>Trachurus japonicus</i> juveniles. <i>Journal of Fish Biology</i> , 2019, 95, 979-981.	0.7	18
12	Effect of water temperature and fish biomass on environmental DNA shedding, degradation, and size distribution. <i>Ecology and Evolution</i> , 2019, 9, 1135-1146.	0.8	183
13	Dispersion and degradation of environmental DNA from caged fish in a marine environment. <i>Fisheries Science</i> , 2019, 85, 327-337.	0.7	102
14	Comparing local and regional scale estimations of the diversity of stream fish using eDNA metabarcoding and conventional observation methods. <i>Freshwater Biology</i> , 2018, 63, 569-580.	1.2	88
15	A generalist herbivore requires a wide array of plant species to maintain its populations. <i>Biological Conservation</i> , 2018, 228, 167-174.	1.9	13
16	Environmental DNA metabarcoding reveals local fish communities in a species-rich coastal sea. <i>Scientific Reports</i> , 2017, 7, 40368.	1.6	348
17	Rapid degradation of longer DNA fragments enables the improved estimation of distribution and biomass using environmental DNA. <i>Molecular Ecology Resources</i> , 2017, 17, e25-e33.	2.2	113
18	Environmental DNA reflects spatial and temporal jellyfish distribution. <i>PLoS ONE</i> , 2017, 12, e0173073.	1.1	87

#	ARTICLE	IF	CITATIONS
19	Phylogenetic analysis of the winter geometrid genus <i>Inurois</i> reveals repeated reproductive season shifts. <i>Molecular Phylogenetics and Evolution</i> , 2016, 94, 47-54.	1.2	9
20	Environmental DNA as a "Snapshot" of Fish Distribution: A Case Study of Japanese Jack Mackerel in Maizuru Bay, Sea of Japan. <i>PLoS ONE</i> , 2016, 11, e0149786.	1.1	192
21	Spatial Segregation and Aggregation of Ectomycorrhizal and Root-Endophytic Fungi in the Seedlings of Two <i>Quercus</i> Species. <i>PLoS ONE</i> , 2014, 9, e96363.	1.1	32
22	Parallel allochronic divergence in a winter moth due to disruption of reproductive period by winter harshness. <i>Molecular Ecology</i> , 2012, 21, 174-183.	2.0	22
23	Incipient allochronic speciation by climatic disruption of the reproductive period. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2009, 276, 2711-2719.	1.2	51
24	Phylogeny of the Geometridae and the evolution of winter moths inferred from a simultaneous analysis of mitochondrial and nuclear genes. <i>Molecular Phylogenetics and Evolution</i> , 2007, 44, 711-723.	1.2	75