

Ryuji J Machida

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

19
papers

1,085
citations

11
h-index

25
g-index

25
ext. papers

1,462
ext. citations

4.9
avg, IF

4.37
L-index

#	Paper	IF	Citations
19	A new versatile primer set targeting a short fragment of the mitochondrial COI region for metabarcoding metazoan diversity: application for characterizing coral reef fish gut contents. <i>Frontiers in Zoology</i> , 2013 , 10, 34	2.8	582
18	Metazoan mitochondrial gene sequence reference datasets for taxonomic assignment of environmental samples. <i>Scientific Data</i> , 2017 , 4, 170027	8.2	90
17	GenBank is a reliable resource for 21st century biodiversity research. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 22651-22656	11.5	75
16	Large-scale gene rearrangements in the mitochondrial genomes of two calanoid copepods <i>Eucalanus bungii</i> and <i>Neocalanus cristatus</i> (Crustacea), with notes on new versatile primers for the srRNA and COI genes. <i>Gene</i> , 2004 , 332, 71-8	3.8	66
15	PCR primers for metazoan nuclear 18S and 28S ribosomal DNA sequences. <i>PLoS ONE</i> , 2012 , 7, e46180	3.7	57
14	Zooplankton diversity analysis through single-gene sequencing of a community sample. <i>BMC Genomics</i> , 2009 , 10, 438	4.5	54
13	Comparison of morphological and molecular traits for species identification and taxonomic grouping of oncaeid copepods. <i>Hydrobiologia</i> , 2011 , 666, 111-125	2.4	37
12	Dissimilarity of species and forms of planktonic <i>Neocalanus</i> copepods using mitochondrial COI, 12S, nuclear ITS, and 28S gene sequences. <i>PLoS ONE</i> , 2010 , 5, e10278	3.7	31
11	Genetic diversity and cryptic speciation of the deep sea chaetognath <i>Caecosagitta macrocephala</i> (Fowler, 1904). <i>Deep-Sea Research Part II: Topical Studies in Oceanography</i> , 2010 , 57, 2211-2219	2.3	28
10	MIDORI server: a webserver for taxonomic assignment of unknown metazoan mitochondrial-encoded sequences using a curated database. <i>Bioinformatics</i> , 2018 , 34, 3753-3754	7.2	26
9	PCR primers for metazoan mitochondrial 12S ribosomal DNA sequences. <i>PLoS ONE</i> , 2012 , 7, e35887	3.7	20
8	Occurrence of mitochondrial CO1 pseudogenes in <i>Neocalanus plumchrus</i> (Crustacea: Copepoda): Hybridization indicated by recombined nuclear mitochondrial pseudogenes. <i>PLoS ONE</i> , 2017 , 12, e0172710	3.7	7
7	Reply to Locatelli et al.: Evaluating species-level accuracy of GenBank metazoan sequences will require experts' effort in each group. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 32213-32214	11.5	2
6	Ways to mix multiple PCR amplicons into single 454 run for DNA barcoding. <i>Methods in Molecular Biology</i> , 2012 , 858, 355-61	1.4	2
5	To the light side: molecular diversity and morphology of stomatopod larvae and juveniles (Crustacea: Malacostraca: Stomatopoda) from crustose coralline algal reefs in Taiwan. <i>Marine Biodiversity</i> , 2021 , 51, 1	1.4	2
4	MIDORI2: A collection of quality controlled, preformatted, and regularly updated reference databases for taxonomic assignment of eukaryotic mitochondrial sequences. <i>Environmental DNA</i> ,	7.6	2
3	Nuclear and mitochondrial ribosomal ratio as an index of animal growth rate. <i>Limnology and Oceanography: Methods</i> , 2019 , 17, 575-584	2.6	1

- 2 Using metatranscriptomics to estimate the diversity and composition of zooplankton communities. *Molecular Ecology Resources*, **2021**, 8.4 1
- 1 Seabed mining could come at a high price for a unique fauna. *Molecular Ecology*, **2020**, 29, 4506-4509 5.7 0