

Freed Ahmad

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6699586/publications.pdf>

Version: 2024-02-01

12
papers

343
citations

1163117

8
h-index

1199594

12
g-index

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all docs

12
docs citations

12
times ranked

888
citing authors

#	ARTICLE	IF	CITATIONS
1	The Glanville fritillary genome retains an ancient karyotype and reveals selective chromosomal fusions in Lepidoptera. <i>Nature Communications</i> , 2014, 5, 4737.	12.8	196
2	Highly Continuous Genome Assembly of Eurasian Perch (<i>Perca fluviatilis</i>) Using Linked-Read Sequencing. <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 3737-3743.	1.8	42
3	Less is more: extreme genome complexity reduction with ddRAD using Ion Torrent semiconductor technology. <i>Molecular Ecology Resources</i> , 2015, 15, 1145-1152.	4.8	22
4	Invasion genomics: genotyping-by-sequencing approach reveals regional genetic structure and signatures of temporal selection in an introduced mud crab. <i>Marine Biology</i> , 2017, 164, 1.	1.5	13
5	Comparative High-Density Linkage Mapping Reveals Conserved Genome Structure but Variation in Levels of Heterochiasmy and Location of Recombination Cold Spots in the Common Frog. <i>G3: Genes, Genomes, Genetics</i> , 2017, 7, 637-645.	1.8	12
6	Association mapping reveals candidate loci for resistance and anaemic response to an emerging temperature-driven parasitic disease in a wild salmonid fish. <i>Molecular Ecology</i> , 2018, 27, 1385-1401.	3.9	11
7	The strength and form of natural selection on transcript abundance in the wild. <i>Molecular Ecology</i> , 2021, 30, 2724-2737.	3.9	11
8	Double-restriction-site-associated DNA (dRAD) approach for fast microsatellite marker development in Eurasian perch (<i>Perca fluviatilis</i> L.). <i>Conservation Genetics Resources</i> , 2014, 6, 183-184.	0.8	10
9	Know your enemy – transcriptome of myxozoan <i>Tetracapsuloides bryosalmonae</i> reveals potential drug targets against proliferative kidney disease in salmonids. <i>Parasitology</i> , 2021, 148, 726-739.	1.5	9
10	Humic-acid-driven escape from eye parasites revealed by RNA-seq and target-specific metabarcoding. <i>Parasites and Vectors</i> , 2020, 13, 433.	2.5	7
11	Mapping of quantitative trait loci for life history traits segregating within common frog populations. <i>Heredity</i> , 2019, 122, 800-808.	2.6	5
12	Tonsillar microbial diversity, abundance, and interrelations in atopic and non-atopic individuals. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2020, 75, 2133-2135.	5.7	5