

Iratxe Montes

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

284
citations

933447

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docs citations

18
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575
citing authors

#	ARTICLE	IF	CITATIONS
1	New Nuclear SNP Markers Unravel the Genetic Structure and Effective Population Size of Albacore Tuna (<i>Thunnus alalunga</i>). <i>PLoS ONE</i> , 2015, 10, e0128247.	2.5	43
2	SNP Discovery in European Anchovy (<i>Engraulis encrasicolus</i> , L) by High-Throughput Transcriptome and Genome Sequencing. <i>PLoS ONE</i> , 2013, 8, e70051.	2.5	38
3	Authoritative subspecies diagnosis tool for European honey bees based on ancestry informative SNPs. <i>BMC Genomics</i> , 2021, 22, 101.	2.8	34
4	Worldwide genetic structure of albacore <i>Thunnus alalunga</i> revealed by microsatellite DNA markers. <i>Marine Ecology - Progress Series</i> , 2012, 471, 183-191.	1.9	29
5	SP110 as a novel susceptibility gene for <i>Mycobacterium avium</i> subspecies paratuberculosis infection in cattle. <i>Journal of Dairy Science</i> , 2010, 93, 5950-5958.	3.4	25
6	Insights on the drivers of genetic divergence in the European anchovy. <i>Scientific Reports</i> , 2017, 7, 4180.	3.3	17
7	No loss of genetic diversity in the exploited and recently collapsed population of Bay of Biscay anchovy (<i>Engraulis encrasicolus</i> , L.). <i>Marine Biology</i> , 2016, 163, 1.	1.5	14
8	Transcriptome analysis deciphers evolutionary mechanisms underlying genetic differentiation between coastal and offshore anchovy populations in the Bay of Biscay. <i>Marine Biology</i> , 2016, 163, 1.	1.5	14
9	Genetic population structure of anchovy (<i>Engraulis encrasicolus</i>) in North-western Europe and variability in the seasonal distribution of the stocks. <i>Fisheries Research</i> , 2020, 229, 105619.	1.7	14
10	Connectivity and population structure of albacore tuna across southeast Atlantic and southwest Indian Oceans inferred from multidisciplinary methodology. <i>Scientific Reports</i> , 2020, 10, 15657.	3.3	13
11	High resolution SNPs selection in <i>Engraulis encrasicolus</i> through Taqman OpenArray. <i>Fisheries Research</i> , 2016, 177, 31-38.	1.7	9
12	Genetic structure of Iranian indigenous sheep breeds: insights for conservation. <i>Tropical Animal Health and Production</i> , 2020, 52, 2283-2290.	1.4	9
13	Hygroregulation, a key ability for eusocial insects: Native Western European honeybees as a case study. <i>PLoS ONE</i> , 2019, 14, e0200048.	2.5	7
14	Genomic Resources Notes Accepted 1 June 2015 - 31 July 2015. <i>Molecular Ecology Resources</i> , 2015, 15, 1510-1512.	4.8	6
15	Reduced Single Nucleotide Polymorphism Panels for Assigning Atlantic Albacore and Bay of Biscay Anchovy Individuals to Their Geographic Origin: Toward Sustainable Fishery Management. <i>Journal of Agricultural and Food Chemistry</i> , 2017, 65, 4351-4358.	5.2	5
16	Development of gene-associated single nucleotide polymorphisms for Japanese anchovy <i>Engraulis japonicus</i> through cross-species amplification. <i>Fisheries Science</i> , 2018, 84, 1-7.	1.6	3
17	A novel transcriptome-derived SNPs array for tench (<i>Tinca tinca</i> L.). <i>PLoS ONE</i> , 2019, 14, e0213992.	2.5	3
18	Transcriptomic dataset for <i>Sardina pilchardus</i> : Assembly, annotation, and expression of nine tissues. <i>Data in Brief</i> , 2021, 39, 107583.	1.0	1