

Patrick Berrebi

List of Publications by Year in descending order

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128
papers

3,954
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134
all docs

134
docs citations

134
times ranked

2917
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#	ARTICLE	IF	CITATIONS
1	Spatial heterogeneity in the Mediterranean Biodiversity Hotspot affects barcoding accuracy of its freshwater fishes. <i>Molecular Ecology Resources</i> , 2014, 14, 1210-1221.	4.8	224
2	Phylogeny and Biogeography of the Family Cyprinidae in the Middle East Inferred from Cytochrome b DNA " Evolutionary Significance of This Region. <i>Molecular Phylogenetics and Evolution</i> , 2002, 22, 91-100.	2.7	173
3	Genetic structure of the flounders <i>Platichthys flesus</i> and <i>P. stellatus</i> at different geographic scales. <i>Marine Biology</i> , 1997, 129, 233-246.	1.5	145
4	Molecular Phylogeny of North Mediterranean Freshwater Barbs (Genus <i>Barbus</i> : Cyprinidae) Inferred from Cytochrome b Sequences: Biogeographic and Systematic Implications. <i>Molecular Phylogenetics and Evolution</i> , 2000, 14, 165-179.	2.7	123
5	Phylogeography of the barbel (<i>Barbus barbus</i>) assessed by mitochondrial DNA variation. <i>Molecular Ecology</i> , 2001, 10, 2177-2185.	3.9	116
6	Extreme genetic differentiation among the remnant populations of marble trout (<i>Salmo</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 542 T	3.9	101
7	The genetic diversity of native, stocked and hybrid populations of marble trout in the Soca river, Slovenia. <i>Heredity</i> , 2000, 85, 277-287.	2.6	95
8	Allozyme variation in turbot (<i>Psetta maxima</i>) and brill (<i>Scophthalmus rhombus</i>) (Osteichthyes,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 46	1.6	83
9	Microsatellite polymorphism and genetic impact of restocking in Mediterranean brown trout (<i>Salmo</i>) Tj ETQq1 1 0.784314 rgBT /Ove	2.6	83
10	Multiple origins of polyploidy in the phylogeny of southern African barbs (Cyprinidae) as inferred from mtDNA markers. <i>Heredity</i> , 2002, 88, 466-473.	2.6	77
11	Unexpected Genetic Diversity among and within Populations of the Toxic Dinoflagellate <i>Alexandrium catenella</i> as Revealed by Nuclear Microsatellite Markers. <i>Applied and Environmental Microbiology</i> , 2009, 75, 2037-2045.	3.1	73
12	Rapid radiation of the Mediterranean <i>Luciobarbus</i> species (Cyprinidae) after the Messinian salinity crisis of the Mediterranean Sea, inferred from mitochondrial phylogenetic analysis. <i>Biological Journal of the Linnean Society</i> , 2003, 80, 207-222.	1.6	70
13	Genetic introgression between wild and stocked salmonids and the prospects for using molecular markers in population rehabilitation: the case of the Adriatic grayling (<i>Thymallus thymallus</i> L. 1785). <i>Heredity</i> , 2004, 93, 273-282.	2.6	65
14	Phylogenetic relationships of hexaploid large-sized barbs (genus <i>Labeobarbus</i> , Cyprinidae) based on mtDNA data. <i>Molecular Phylogenetics and Evolution</i> , 2010, 56, 851-856.	2.7	58
15	Speciation of the genus <i>Barbus</i> in the north Mediterranean basin: Recent advances from biochemical genetics. <i>Biological Conservation</i> , 1995, 72, 237-249.	4.1	57
16	Distribution of non-native trout in Slovenia and their introgression with native trout populations as observed through microsatellite DNA analysis. <i>Biological Conservation</i> , 2005, 123, 381-388.	4.1	56
17	Oceanic larval life of <i>La R�union �bichiques�</i> , amphidromous gobiid post-larvae. <i>Marine Ecology - Progress Series</i> , 2007, 333, 303-308.	1.9	52
18	Stocking impact and allozyme diversity in brown trout from Mediterranean southern France. <i>Journal of Fish Biology</i> , 2000, 56, 949-960.	1.6	51

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19	Ubiquity of <i>Sicyopterus lagocephalus</i> (Teleostei: Gobioidae) and phylogeography of the genus <i>Sicyopterus</i> in the Indo-Pacific area inferred from mitochondrial cytochrome b gene. <i>Molecular Phylogenetics and Evolution</i> , 2005, 37, 721-732.	2.7	51
20	Hybridization mechanisms between the endangered marble trout (<i>Salmo marmoratus</i>) and the brown trout (<i>Salmo trutta</i>) as revealed by in-stream experiments. <i>Biological Conservation</i> , 2007, 136, 602-611.	4.1	50
21	Nuclear and mitochondrial DNA sequence data reveal the evolutionary history of <i>Barbus</i> (Cyprinidae) in the ancient lake systems of the Balkans. <i>Molecular Phylogenetics and Evolution</i> , 2010, 55, 488-500.	2.7	49
22	Evolution of Polyploidy and Functional Diploidization in Sturgeons: Microsatellite Analysis in 10 Sturgeon Species. <i>Journal of Heredity</i> , 2014, 105, 521-531.	2.4	47
23	Relative ages of present populations of <i>Barbus barbus</i> and <i>Barbus meridionalis</i> (Cyprinidae) in southern France: preliminary considerations. <i>Aquatic Living Resources</i> , 1990, 3, 253-263.	1.2	45
24	Spanish barbel hybridization detected using enzymatic markers: <i>Barbus meridionalis</i> Risso – <i>Barbus haasi</i> Mertens (Osteichthyes, Cyprinidae). <i>Aquatic Living Resources</i> , 1990, 3, 295-303.	1.2	45
25	Biogeography and pattern of gene flow among <i>Barbus</i> species (Teleostei: Cyprinidae) inhabiting the Italian Peninsula and neighbouring Adriatic drainages as revealed by allozyme and mitochondrial sequence data. <i>Biological Journal of the Linnean Society</i> , 2002, 75, 83-99.	1.6	42
26	Phylogeographic structure and demographic patterns of brown trout in North-West Africa. <i>Molecular Phylogenetics and Evolution</i> , 2011, 61, 203-211.	2.7	41
27	Nuclear Markers of Danube Sturgeons Hybridization. <i>International Journal of Molecular Sciences</i> , 2011, 12, 6796-6809.	4.1	40
28	Genetic architecture of trout from Albania as revealed by mtDNA control region variation. <i>Genetics Selection Evolution</i> , 2009, 41, 22.	3.0	38
29	A Dense Brown Trout (<i>Salmo trutta</i>) Linkage Map Reveals Recent Chromosomal Rearrangements in the <i>Salmo</i> Genus and the Impact of Selection on Linked Neutral Diversity. <i>G3: Genes, Genomes, Genetics</i> , 2017, 7, 1365-1376.	1.8	38
30	Genomic consequences of a recent three-way admixture in supplemented wild brown trout populations revealed by local ancestry tracts. <i>Molecular Ecology</i> , 2018, 27, 3466-3483.	3.9	38
31	Natural hybridization of two species of tetraploid barbels: <i>Barbus meridionalis</i> and <i>Barbus barbus</i> (Osteichthyes, Cyprinidae) in southern France. <i>Biological Journal of the Linnean Society</i> , 1993, 48, 319-333.	1.6	37
32	Genetic structure of brown trout (<i>Salmo trutta</i> , L.) populations from south-western France: data from mitochondrial control region variability. <i>Molecular Ecology</i> , 2001, 10, 1551-1561.	3.9	36
33	Primers for EPIC amplification of intron sequences for fish and other vertebrate population genetic studies. <i>BioTechniques</i> , 2003, 35, 676-682.	1.8	36
34	Genetic subdivision and biogeography of the Danubian rheophilic barb <i>Barbus petenyi</i> inferred from phylogenetic analysis of mitochondrial DNA variation. <i>Molecular Phylogenetics and Evolution</i> , 2002, 24, 10-18.	2.7	35
35	Intron polymorphism (EPIC-PCR) reveals phylogeographic structure of <i>Zacco platypus</i> in China: a possible target for aquaculture development. <i>Heredity</i> , 2005, 94, 589-598.	2.6	35
36	Assessment of cryptic species diversity within blooms and cyst bank of the <i>Alexandrium tamarense</i> complex (Dinophyceae) in a Mediterranean lagoon facilitated by semi-multiplex PCR. <i>Journal of Plankton Research</i> , 2011, 33, 405-414.	1.8	35

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37	Brown trout (<i>Salmo trutta</i> L.) high genetic diversity around the Tyrrhenian Sea as revealed by nuclear and mitochondrial markers. <i>Hydrobiologia</i> , 2019, 826, 209-231.	2.0	35
38	Do self-fertilization and genetic drift promote a very low genetic variability in the allotetraploid <i>Bulinus truncatus</i> (Gastropoda: Planorbidae) populations?. <i>Genetical Research</i> , 1993, 62, 89-100.	0.9	34
39	<i>Salmo macrostigma</i> (Teleostei, Salmonidae): Nothing more than a brown trout (<i>S.</i>) Tj ETQq1 1 0.784314 rgBT /Overlo	1.6	34
40	A genomic perspective on an old question: <i>Salmo</i> trouts or <i>Salmo trutta</i> (Teleostei: Salmonidae)?. <i>Molecular Phylogenetics and Evolution</i> , 2021, 162, 107204.	2.7	33
41	Diploid and tetraploid African <i>Barbus</i> (Osteichthyes, Cyprinidae): on the coding of differential gene expression. <i>Aquatic Living Resources</i> , 1990, 3, 313-323.	1.2	33
42	Two lineages, diploid and tetraploid, demonstrated in African species <i>Barbus</i> (Osteichthyes, Cyprinidae). <i>Aquatic Living Resources</i> , 1990, 3, 305-311.	1.2	32
43	Morphological variation in hybrids between <i>Salmo marmoratus</i> , and alien <i>Salmo</i> species in the Volarja stream, Soca River basin, Slovenia. <i>Journal of Fish Biology</i> , 2000, 57, 1199-1212.	1.6	32
44	Historical presence of the sturgeon <i>Acipenser sturio</i> in the Rhône basin determined by the analysis of ancient DNA cytochrome b sequences. <i>Conservation Genetics</i> , 2009, 10, 217-224.	1.5	32
45	Genetic diversity and population structure of domestic brown trout (<i>Salmo trutta</i>) in France. <i>Aquaculture</i> , 2016, 462, 1-9.	3.5	32
46	Natural and artificial secondary contact in brown trout (<i>Salmo trutta</i> , L.) in the French western Pyrenees assessed by allozymes and microsatellites. <i>Heredity</i> , 2002, 89, 171-183.	2.6	31
47	Population structure and systematics of <i>Opsariichthys bidens</i> (Osteichthyes: Cyprinidae) in south-east China using a new nuclear marker: the introns (EPIC-PCR). <i>Biological Journal of the Linnean Society</i> , 2006, 87, 155-166.	1.6	31
48	Microgeographic genetic isolation in chub (<i>Cyprinidae: Squalius cephalus</i>) population of the Durance River: estimating fragmentation by dams. <i>Ecology of Freshwater Fish</i> , 2010, 19, 267-278.	1.4	31
49	Genetic divergence among morphotypes of Lake Tana (Ethiopia) barbs. <i>Biological Journal of the Linnean Society</i> , 1998, 64, 369-384.	1.6	29
50	Genome-wide nucleotide diversity of hatchery-reared Atlantic and Mediterranean strains of brown trout (<i>Salmo trutta</i>) compared to wild Mediterranean populations. <i>Journal of Fish Biology</i> , 2016, 89, 2717-2734.	1.6	29
51	Microsatellites and artificial neural networks: tools for the discrimination between natural and hatchery brown trout (<i>Salmo trutta</i> , L.) in Atlantic populations. <i>Ecological Modelling</i> , 1999, 120, 313-324.	2.5	27
52	Evolutionary history of a widespread Indo-Pacific goby: The role of Pleistocene sea-level changes on demographic contraction/expansion dynamics. <i>Molecular Phylogenetics and Evolution</i> , 2012, 62, 566-572.	2.7	27
53	Rare and asymmetrical hybridization of the endemic <i>Barbus carpathicus</i> with its widespread congener <i>Barbus barbus</i> . <i>Journal of Fish Biology</i> , 2009, 74, 418-436.	1.6	26
54	Effects of current and historic habitat fragmentation on the genetic structure of the sand goby <i>Pomatoschistus minutus</i> (Osteichthys, Gobiidae). <i>Biological Journal of the Linnean Society</i> , 2011, 102, 175-198.	1.6	26

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55	Differences between nuclear and mitochondrial introgressions of brown trout populations from a restocked main river and its unstocked tributary. <i>Biological Journal of the Linnean Society</i> , 1998, 63, 379-392.	1.6	25
56	Gene flow and genetic structure of <i>Sicyopterus lagocephalus</i> in the south-western Indian Ocean, assessed by intron-length polymorphism. <i>Journal of Experimental Marine Biology and Ecology</i> , 2007, 349, 223-234.	1.5	25
57	Development of a large SNPs resource and a low-density SNP array for brown trout (<i>Salmo trutta</i>) population genetics. <i>BMC Genomics</i> , 2019, 20, 582.	2.8	25
58	Asymmetrical introgression in a freshwater fish hybrid zone as revealed by a morphological index of hybridization. <i>Biological Journal of the Linnean Society</i> , 1999, 67, 57-72.	1.6	24
59	Present and past genetic connectivity of the Indo-Pacific tropical eel <i>Anguilla bicolor</i> . <i>Journal of Biogeography</i> , 2012, 39, 408-420.	3.0	24
60	Species boundaries in the <i>Himantura uarnak</i> species complex (Myliobatiformes: Dasyatidae). <i>Molecular Phylogenetics and Evolution</i> , 2013, 66, 429-435.	2.7	24
61	Experimental hybridization of <i>Barbus barbus</i> and <i>Barbus meridionalis</i> : physiological, morphological, and genetic aspects. <i>Aquatic Living Resources</i> , 1990, 3, 325-332.	1.2	24
62	Interpopulation relationships in two species of Antarctic fish <i>Notothenia rossii</i> and <i>Champscephalus gunnari</i> from the Kerguelen Islands: an allozyme study. <i>Antarctic Science</i> , 1995, 7, 351-356.	0.9	23
63	<i>Anguilla marmorata</i> larval migration plasticity as revealed by otolith microstructural analysis. <i>Canadian Journal of Fisheries and Aquatic Sciences</i> , 2008, 65, 2127-2137.	1.4	23
64	Palaeogenetics of western French sturgeons spotlights the relationships between <i>Acipenser sturio</i> and <i>Acipenser oxyrinchus</i> . <i>Journal of Biogeography</i> , 2013, 40, 382-393.	3.0	22
65	Enzymatic survey of four populations of <i>Atherina boyeri</i> based on electrophoresis and the occurrence of a microsporidiosis. <i>Journal of Fish Biology</i> , 1980, 16, 149-157.	1.6	21
66	Population genetic structure of blue-spotted maskray <i>Neotrygon kuhlii</i> and two other Indo-West Pacific stingray species (Myliobatiformes: Dasyatidae), inferred from size-polymorphic intron markers. <i>Journal of Experimental Marine Biology and Ecology</i> , 2012, 438, 32-40.	1.5	21
67	<i>Himantura tutul</i> sp. nov. (Myliobatoidei: Dasyatidae), a new ocellated whipray from the tropical Indo-West Pacific, described from its cytochrome-oxidase I gene sequence. <i>Comptes Rendus - Biologies</i> , 2013, 336, 82-92.	0.2	21
68	Three brown trout <i>Salmo trutta</i> lineages in Corsica described through allozyme variation. <i>Journal of Fish Biology</i> , 2015, 86, 60-73.	1.6	21
69	Ocean currents drive secondary contact between <i>Anguilla marmorata</i> populations in the Indian Ocean. <i>Marine Ecology - Progress Series</i> , 2009, 379, 267-278.	1.9	20
70	Does polyploidy lead to fewer and shorter microsatellites in <i>Barbus</i> (Teleostei: Cyprinidae)? <i>Molecular Ecology</i> , 1997, 6, 169-178.	3.9	19
71	Differential distribution of the two cryptic species, <i>Pomatoschistus microps</i> and <i>P. marmoratus</i> , in the lagoons of southern France, with an emphasis on the genetic organisation of <i>P. microps</i> . <i>Estuarine, Coastal and Shelf Science</i> , 2005, 65, 708-716.	2.1	19
72	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 April 2013–31 May 2013. <i>Molecular Ecology Resources</i> , 2013, 13, 966-968.	4.8	19

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73	Shape plasticity in response to water velocity in the freshwater blenny <i>Salaria fluviatilis</i> . <i>Journal of Fish Biology</i> , 2016, 88, 1191-1203.	1.6	19
74	Genetic monitoring for the successful re-stocking of a critically endangered diadromous fish with low diversity. <i>Biological Conservation</i> , 2018, 221, 91-102.	4.1	19
75	Intégration génomique et repeuplements chez la truite commune du versant méditerranéen. <i>Knowledge and Management of Aquatic Ecosystems: an International Journal on Aquatic Ecosystems</i> , 1997, , 309-322.	0.4	18
76	Selection footprint at the first intron of the <i>Prl</i> gene in natural populations of the flathead mullet (<i>Mugil cephalus</i> , L. 1758). <i>Journal of Experimental Marine Biology and Ecology</i> , 2010, 387, 60-67.	1.5	18
77	Phylogeography of the European sturgeon (<i>Acipenser sturio</i>): A critically endangered species. <i>Molecular Phylogenetics and Evolution</i> , 2016, 94, 346-357.	2.7	18
78	An isozyme study of the natural cyprinid hybrid <i>Alburnus alburnus</i> x <i>Rutilus rubilio</i> in Greece. <i>Journal of Fish Biology</i> , 1989, 34, 307-313.	1.6	16
79	Allozymes, mtDNA and microsatellites study introgression in a stocked trout population in France. <i>Reviews in Fish Biology and Fisheries</i> , 2000, 10, 281-292.	4.9	16
80	Genetic Diversity and Conservation of the Prespa Trout in the Balkans. <i>International Journal of Molecular Sciences</i> , 2013, 14, 23454-23470.	4.1	16
81	Biodiversity and distribution of leptocephali west of the Mascarene Plateau in the southwestern Indian Ocean. <i>Progress in Oceanography</i> , 2015, 137, 84-102.	3.2	16
82	Geographic structure evidenced in the toxic dinoflagellate <i>Alexandrium pacificum</i> Litaker (A.) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 387. <i>Marine Pollution Bulletin</i> , 2015, 98, 95-105.	5.0	16
83	A behavioural study of hybridization between <i>Barbus barbus</i> and <i>Barbus meridionalis</i> . <i>Journal of Fish Biology</i> , 1994, 45, 447-451.	1.6	15
84	The ecology of sexual dimorphism in size and shape of the freshwater blenny <i>Salaria fluviatilis</i> . <i>Environmental Epigenetics</i> , 2018, 64, 183-191.	1.8	15
85	Genetic diversity of domestic brown trout stocks in Europe. <i>Aquaculture</i> , 2021, 544, 737043.	3.5	15
86	Parasites as biological markers: evolutionary relationships in the heterospecific combination of helminths (monogeneans) and teleosts (Gadidae). <i>Biological Journal of the Linnean Society</i> , 1992, 47, 173-182.	1.6	14
87	Distinguishing species of European sturgeons <i>Acipenser</i> spp. using microsatellite allele sequences. <i>Journal of Fish Biology</i> , 2011, 78, 208-226.	1.6	14
88	New Light on the Evolutionary History of the Common Goby (<i>Pomatoschistus microps</i>) with an Emphasis on Colonization Processes in the Mediterranean Sea. <i>PLoS ONE</i> , 2014, 9, e91576.	2.5	14
89	Natural and domestic introgressions in the marble trout population of Soča River (Slovenia). <i>Hydrobiologia</i> , 2017, 785, 277-291.	2.0	14
90	Spatially varying natural selection in a fish hybrid zone. <i>Journal of Fish Biology</i> , 2002, 61, 696-711.	1.6	13

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91	Unsuspected intraspecific variability in the toxin production, growth and morphology of the dinoflagellate <i>Alexandrium pacificum</i> R.W. Litaker (Group IV) blooming in a South Western Mediterranean marine ecosystem, Annaba Bay (Algeria). <i>Toxicon</i> , 2020, 180, 79-88.	1.6	13
92	The role of the south-western Alps as a unidirectional corridor for Mediterranean brown trout (<i>Salmo trutta</i> complex) lineages. <i>Biological Journal of the Linnean Society</i> , 2020, 131, 909-926.	1.6	12
93	Morphologic and genetic characterisation of Corsican and Sardinian trout with comments on <i>Salmo</i> taxonomy. <i>Knowledge and Management of Aquatic Ecosystems</i> , 2020, , 21.	1.1	12
94	Ecological and genetic differentiation of <i>Barbus callensis</i> populations in Tunisia. <i>Journal of Fish Biology</i> , 1995, 47, 850-864.	1.6	11
95	Microsatellite markers and management of brown trout <i>Salmo trutta fario</i> populations in southwestern France. <i>Genetics Selection Evolution</i> , 1998, 30, 1.	3.0	11
96	Autosomal differences between males and females in hybrid zones: a first report from <i>Barbus barbus</i> and <i>Barbus meridionalis</i> (Cyprinidae). <i>Heredity</i> , 2004, 93, 128-134.	2.6	11
97	Population genetic structure of <i>Cichla pleiozona</i> (Perciformes: Cichlidae) in the Upper Madera basin (Bolivian Amazon): Sex-biased dispersal?. <i>Molecular Phylogenetics and Evolution</i> , 2010, 57, 1334-1340.	2.7	11
98	Development of compound microsatellite markers in the toxic dinoflagellate <i>Alexandrium catenella</i> (Dinophyceae). <i>Plankton and Benthos Research</i> , 2007, 2, 128-133.	0.6	11
99	Parentage assignment in the critically endangered European sturgeon (<i>Acipenser sturio</i>) based on a novel microsatellite multiplex assay: a valuable resource for restocking, monitoring and conservation programs. <i>Conservation Genetics Resources</i> , 2016, 8, 313-322.	0.8	9
100	Esterases of the flounder (<i>Platichthys flesus</i> , Pleuronectidae, Teleostei): Development of an identification protocol using starch gel electrophoresis and characterization of loci. <i>Experientia</i> , 1990, 46, 863-867.	1.2	8
101	Biodiversity and biogeography in heterospecific teleostean (<i>Gadidae</i>) "copepod (<i>Lernaecocera</i>) associations. <i>Canadian Journal of Zoology</i> , 1993, 71, 1639-1645.	1.0	8
102	Development of a key using morphological characters to distinguish south-western Indian Ocean anguillid glass eels. <i>Journal of Fish Biology</i> , 2009, 74, 2171-2177.	1.6	8
103	Absence of spatial genetic structure in common dentex (<i>Dentex dentex</i> Linnaeus, 1758) in the Mediterranean Sea as evidenced by nuclear and mitochondrial molecular markers. <i>PLoS ONE</i> , 2018, 13, e0203866.	2.5	8
104	Cytogenetic and molecular genetic data. <i>Italian Journal of Zoology</i> , 1998, 65, 15-20.	0.6	7
105	Isolation of microsatellite loci and cross-species amplifications in three gobiid fish of the genus <i>Pomatoschistus</i> . <i>Molecular Ecology Notes</i> , 2006, 6, 724-727.	1.7	7
106	Isolation of ten microsatellite markers using a pyrosequencing procedure and cross-priming in the <i>Salaria</i> genus. <i>Conservation Genetics Resources</i> , 2012, 4, 151-154.	0.8	7
107	Morphological variation in the freshwater blenny <i>Salaria fluviatilis</i> from Corsican rivers: adaptive divergence, phenotypic plasticity or both?. <i>Journal of Fish Biology</i> , 2014, 84, 31-44.	1.6	7
108	Detecting anthropogenic effects on a vulnerable species, the freshwater blenny (<i>Salaria fluviatilis</i>): The importance of considering key ecological variables. <i>Ecological Indicators</i> , 2014, 36, 386-391.	6.3	7

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109	Genetic evidence of recent migration among isolated-by-sea populations of the freshwater blenny (<i>Salaria fluviatilis</i>). <i>Conservation Genetics</i> , 2016, 17, 389-399.	1.5	7
110	Sturgeon Recovery Plan in the Rhône River (France): Preliminary Results on Species Determination and Habitat Suitability. , 2009, , 403-421.		7
111	Genetic structure of a vulnerable species, the freshwater blenny (<i>Salaria fluviatilis</i>). <i>Conservation Genetics</i> , 2015, 16, 571-581.	1.5	6
112	Diversity and biogeography of Mediterranean freshwater blennies (Blenniidae, <i>Salaria</i>). <i>Diversity and Distributions</i> , 2021, 27, 1832-1847.	4.1	6
113	La osmoregulaci3n como factor potencial de la distribuci3n diferencial de dos especies cr3pticas de g3bido, <i>Pomatoschistus microps</i> y <i>P. marmoratus</i>, en las lagunas mediterr3neas francesas. <i>Scientia Marina</i> , 2008, 72, .	0.6	6
114	Polymorphic microsatellite loci in the widespread amphidromous goby <i>Sicyopterus lagocephalus</i> and cross-genus amplification among <i>Sicydiinae</i>. <i>Molecular Ecology Resources</i> , 2009, 9, 607-609.	4.8	5
115	Single and joint gene segregation in intraspecific hybrids of brown trout (<i>Salmo trutta</i> L.) lineages. <i>Aquaculture</i> , 2000, 186, 1-12.	3.5	4
116	Parentage Analysis with Few Contributing Breeders: Validation and Improvement. <i>Journal of Heredity</i> , 2008, 99, 323-334.	2.4	4
117	Isolation of 12 microsatellite markers following a pyrosequencing procedure and cross-priming in two invasive cryptic species, <i>Alexandrium catenella</i> (group IV) and <i>A. tamarense</i> (group III) (Dinophyceae). <i>Marine Pollution Bulletin</i> , 2014, 83, 302-305.	5.0	4
118	Microsatellite diversity of a critically endangered sturgeon, <i>Acipenser sturio</i> L. 1758, assessed from museum and archaeological tissue remains. <i>Journal of Biogeography</i> , 2018, 45, 1043-1053.	3.0	4
119	Brown trout in Japan ~ introduction history, distribution and genetic structure. <i>Knowledge and Management of Aquatic Ecosystems</i> , 2020, , 18.	1.1	4
120	Delimiting species by reproductive isolation: the genetic structure of epigean and hypogean <i>Trichomycterus</i> spp. (Teleostei, Siluriformes) in the restricted area of Torotoro (Upper Amazon,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 29		
121	Evolutionary process of a tetranucleotide microsatellite locus in <i>Acipenseriformes</i> . <i>Journal of Genetics</i> , 2011, 90, 217-227.	0.7	3
122	Population genetic structure of the tropical eel <i>Anguilla bicolor</i> in Indonesian waters based on microsatellite markers. <i>Folia Zoologica</i> , 2015, 64, 87-96.	0.9	3
123	Restoring marble trout genes in the SoÅa River (Slovenia). <i>Conservation Genetics</i> , 2022, 23, 481-494.	1.5	3
124	Genetic diversity of <i>Pomatoschistus microps</i> (Perciformes: Gobiidae) in ecologically differentiated estuarine systems. <i>Folia Zoologica</i> , 2012, 61, 106-117.	0.9	1
125	Genetic evidence of unisexual reproduction in the Moroccan hexaploid barbel <i>Labeobarbus fritschi</i> . <i>Folia Zoologica</i> , 2013, 62, 257-263.	0.9	1
126	Distribution and hybridization of two sedentary gobies (<i>Pomatoschistus microps</i> and <i>Pomatoschistus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 T	0.6	1

#	ARTICLE	IF	CITATIONS
127	Evidence of unidirectional gene flow in a fragmented population of <i>Salmo trutta</i> L.. Scientific Reports, 2021, 11, 23417.	3.3	1
128	The "Barbus" model (cyprinidae). A focus of multidisciplinary european interest. Aquatic Living Resources, 1990, 3, 251-252.	1.2	0