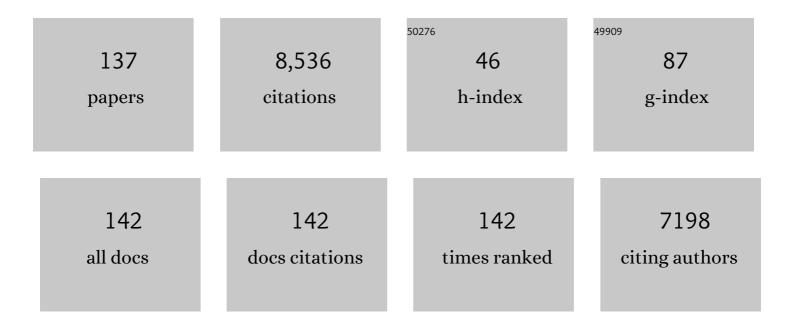
Daniel E Ruzzante

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	ENVIRONMENTAL CORRELATES OF POPULATION DIFFERENTIATION IN ATLANTIC HERRING. Evolution; International Journal of Organic Evolution, 2005, 59, 2656-2668.	2.3	537
2	Genetic estimates of contemporary effective population size: what can they tell us about the importance of genetic stochasticity for wild population persistence?. Molecular Ecology, 2008, 17, 3428-3447.	3.9	487
3	Understanding and Estimating Effective Population Size for Practical Application in Marine Species Management. Conservation Biology, 2011, 25, 438-449.	4.7	270
4	Wholeâ€genome sequencing approaches for conservation biology: Advantages, limitations and practical recommendations. Molecular Ecology, 2017, 26, 5369-5406.	3.9	249
5	Power for detecting genetic divergence: differences between statistical methods and marker loci. Molecular Ecology, 2006, 15, 2031-2045.	3.9	215
6	Genetic differentiation between inshore and offshore Atlantic cod (<i>Gadus morhua</i>) off Newfoundland: microsatellite DNA variation and antifreeze level. Canadian Journal of Fisheries and Aquatic Sciences, 1996, 53, 634-645.	1.4	214
7	Evidence of a hybridâ€zone in Atlantic cod (Gadus morhua) in the Baltic and the Danish Belt Sea revealed by individual admixture analysis. Molecular Ecology, 2003, 12, 1497-1508.	3.9	206
8	Parallel adaptive evolution of Atlantic cod on both sides of the Atlantic Ocean in response to temperature. Proceedings of the Royal Society B: Biological Sciences, 2010, 277, 3725-3734.	2.6	206
9	Biocomplexity in a highly migratory pelagic marine fish, Atlantic herring. Proceedings of the Royal Society B: Biological Sciences, 2006, 273, 1459-1464.	2.6	205
10	Microsatellite polymorphism and the population structure of Atlantic cod (Gadus morhua) in the northwest Atlantic. Canadian Journal of Fisheries and Aquatic Sciences, 1996, 53, 2706-2721.	1.4	194
11	Marine landscapes and population genetic structure of herring (Clupea harengus L.) in the Baltic Sea. Molecular Ecology, 2005, 14, 3219-3234.	3.9	192
12	Poleward bound: biological impacts of Southern Hemisphere glaciation. Trends in Ecology and Evolution, 2012, 27, 462-471.	8.7	186
13	A comparison of several measures of genetic distance and population structure with microsatellite data: bias and sampling variance. Canadian Journal of Fisheries and Aquatic Sciences, 1998, 55, 1-14.	1.4	184
14	Domestication effects on aggressive and schooling behavior in fish. Aquaculture, 1994, 120, 1-24.	3.5	183
15	A nuclear DNA basis for shelf- and bank-scale population structure in northwest Atlantic cod (Gadus) Tj ETQq1	1 0.784314 3.9	rg $_{183}^{ m BT}/ m Overloo$
16	DISENTANGLING THE EFFECTS OF EVOLUTIONARY, DEMOGRAPHIC, AND ENVIRONMENTAL FACTORS INFLUENCING GENETIC STRUCTURE OF NATURAL POPULATIONS: ATLANTIC HERRING AS A CASE STUDY. Evolution; International Journal of Organic Evolution, 2009, 63, 2939-2951.	2.3	183
17	Long-term effective population sizes, temporal stability of genetic composition and potential for local adaptation in anadromous brown trout (Salmo trutta) populations. Molecular Ecology, 2002, 11, 2523-2535.	3.9	156
18	Spatial and temporal variation in the genetic composition of a larval cod (<i>Gadus morhua</i>) aggregation: cohort contribution and genetic stability. Canadian Journal of Fisheries and Aquatic Sciences, 1996, 53, 2695-2705.	1.4	141

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19	Genomic islands of divergence and their consequences for the resolution of spatial structure in an exploited marine fish. Evolutionary Applications, 2013, 6, 450-461.	3.1	136
20	Climate control on ancestral population dynamics: insight from Patagonian fish phylogeography. Molecular Ecology, 2008, 17, 2234-2244.	3.9	134
21	Title is missing!. Conservation Genetics, 2001, 2, 257-269.	1.5	131
22	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 May 2009–31 July 2009. Molecular Ecology Resources, 2009, 9, 1460-1466.	4.8	128
23	Detecting population structure in a high gene-flow species, Atlantic herring (Clupea harengus): direct, simultaneous evaluation of neutral vs putatively selected loci. Heredity, 2011, 106, 270-280.	2.6	126
24	A review of the evidence for genetic structure of cod (Gadus morhua) populations in the NW Atlantic and population affinities of larval cod off Newfoundland and the Gulf of St. Lawrence. Fisheries Research, 1999, 43, 79-97.	1.7	123
25	Microsatellite and mitochondrial DNA polymorphism reveals life-history dependent interbreeding between hatchery and wild brown trout (Salmo trutta L.). Molecular Ecology, 2000, 9, 583-594.	3.9	123
26	Across the southern Andes on fin: glacial refugia, drainage reversals and a secondary contact zone revealed by the phylogeographical signal of <i>Galaxias platei</i> in Patagonia. Molecular Ecology, 2008, 17, 5049-5061.	3.9	117
27	The genetic structure of Atlantic cod (Gadus morhua) around Iceland: insight from microsatellites, the Pan I locus, and tagging experiments. Canadian Journal of Fisheries and Aquatic Sciences, 2006, 63, 2660-2674.	1.4	116
28	Phylogeography of the Percichthyidae (Pisces) in Patagonia: roles of orogeny, glaciation, and volcanism. Molecular Ecology, 2006, 15, 2949-2968.	3.9	108
29	Population structure and gene flow reversals in Atlantic salmon (<i>Salmo salar</i>) over contemporary and longâ€term temporal scales: effects of population size and life history. Molecular Ecology, 2007, 16, 4504-4522.	3.9	107
30	Bay-scale population structure in coastal Atlantic cod in Labrador and Newfoundland, Canada. Journal of Fish Biology, 2000, 56, 431-447.	1.6	98
31	Parallel adaptive evolution of geographically distant herring populations on both sides of the North Atlantic Ocean. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E3452-E3461.	7.1	87
32	Microsatellite loci reveal highly significant genetic differentiation among Atlantic salmon (Salmo) Tj ETQq0 0 0 r	gBT_/Overl	lock 10 Tf 50 2
33	MIXED-STOCK ANALYSIS OF ATLANTIC COD NEAR THE GULF OF ST. LAWRENCE BASED ON MICROSATELLITE DNA. , 2000, 10, 1090-1109.		78
34	BROWN TROUT (SALMO TRUTTA) STOCKING IMPACT ASSESSMENT USING MICROSATELLITE DNA MARKERS. , 2001, 11, 148-160.		75
35	Stocking impact and migration pattern in an anadromous brown trout (Salmo trutta) complex: where have all the stocked spawning sea trout gone?. Molecular Ecology, 2004, 13, 1433-1445.	3.9	73
36	Interaction of landscape and life history attributes on genetic diversity, neutral divergence and gene flow in a pristine community of salmonids. Molecular Ecology, 2009, 18, 4854-4869.	3.9	68

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37	Distribution of individual inbreeding coefficients, relatedness and influence of stocking on native anadromous brown trout (Salmo trutta) population structure. Molecular Ecology, 2001, 10, 2107-2128.	3.9	67
38	North Sea herring population structure revealed by microsatellite analysis. Marine Ecology - Progress Series, 2005, 303, 245-257.	1.9	67
39	Genetic monitoring of supportive breeding in brown trout (<i>Salmo trutta</i> L.), using microsatellite DNA markers. Canadian Journal of Fisheries and Aquatic Sciences, 2000, 57, 2130-2139.	1.4	65
40	Surviving historical Patagonian landscapes and climate: molecular insights from Galaxias maculatus. BMC Evolutionary Biology, 2010, 10, 67.	3.2	61
41	Changes in the distribution of native fishes in response to introduced species and other anthropogenic effects. Global Ecology and Biogeography, 2010, 19, 697-710.	5.8	61
42	Scorched mussels (BIVALVIA: MYTILIDAE: BRACHIDONTINAE) from the temperate coasts of South America: Phylogenetic relationships, trans-Pacific connections and the footprints of Quaternary glaciations. Molecular Phylogenetics and Evolution, 2015, 82, 60-74.	2.7	60
43	Phylogeography and phenotypic diversification in the Patagonian fish Percichthys trucha: the roles of Quaternary glacial cycles and natural selection. Biological Journal of the Linnean Society, 2011, 103, 514-529.	1.6	58
44	Historical changes in genotypic frequencies at the <i>Pantophysin</i> locus in Atlantic cod (<i>Gadus) Tj ETQq0 C 2011, 4, 562-573.</i>	0 o rgBT /C 3.1	overlock 10 Tf 58
45	Native and introduced fish species richness in <scp>C</scp> hilean <scp>P</scp> atagonian lakes: inferences on invasion mechanisms using salmonidâ€free lakes. Diversity and Distributions, 2012, 18, 1153-1165.	4.1	54
46	Genetic differentiation between inshore and offshore Atlantic cod (<i>Gadus morhua</i>) off Newfoundland: a test and evidence of temporal stability. Canadian Journal of Fisheries and Aquatic Sciences, 1997, 54, 2700-2708.	1.4	53
47	Resource polymorphism in a Patagonian fish Percichthys trucha (Percichthyidae): phenotypic evidence for interlake pattern variation. Biological Journal of the Linnean Society, 2003, 78, 497-515.	1.6	53
48	Age Structure, Changing Demography and Effective Population Size in Atlantic Salmon (<i>Salmo) Tj ETQq0 0 0</i>	rgBT/Ovei	rlo <u>ck</u> 10 Tf 50
49	Effective number of breeders, effective population size and their relationship with census size in an iteroparous species, <i>Salvelinus fontinalis</i> . Proceedings of the Royal Society B: Biological Sciences, 2016, 283, 20152601.	2.6	47
50	Title is missing!. Conservation Genetics, 2001, 2, 309-324.	1.5	46
51	Latitudinal and ecological correlates of morphological variation in Galaxias platei (Pisces,) Tj ETQq1 1 0.784314 r	gBT /Over	lock 10 Tf 50
52	Mature male parr contribution to the effective size of an anadromous <scp>A</scp> tlantic salmon (<i><scp>S</scp>almo salar</i>) population over 30Âyears. Molecular Ecology, 2013, 22, 2394-2407.	3.9	43
53	RAPID BEHAVIORAL CHANGES IN MEDAKA (<i>ORYZIAS LATIPES</i>) CAUSED BY SELECTION FOR COMPETITIVE AND NONCOMPETITIVE GROWTH. Evolution; International Journal of Organic Evolution, 1991, 45, 1936-1946.	2.3	42
54	Trophic polymorphism, habitat and diet segregation in Percichthys trucha (Pisces: Percichthyidae) in the Andes. Biological Journal of the Linnean Society, 1998, 65, 191-214.	1.6	42

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55	Validation of closeâ€kin mark–recapture (CKMR) methods for estimating population abundance. Methods in Ecology and Evolution, 2019, 10, 1445-1453.	5.2	42
56	Genetic divergence between sympatric Arctic charr <i>Salvelinus alpinus </i> morphs in Gander Lake, Newfoundland: roles of migration, mutation and unequal effective population sizes. Journal of Fish Biology, 2008, 73, 2040-2057.	1.6	38
57	Climateâ€induced changes to the ancestral population size of two Patagonian galaxiids: the influence of glacial cycling. Molecular Ecology, 2011, 20, 5280-5294.	3.9	37
58	Long Distance Linkage Disequilibrium and Limited Hybridization Suggest Cryptic Speciation in Atlantic Cod. PLoS ONE, 2014, 9, e106380.	2.5	37
59	Genomic evidence of past and future climate-linked loss in a migratory Arctic fish. Nature Climate Change, 2021, 11, 158-165.	18.8	36
60	Environmental correlates of population differentiation in Atlantic herring. Evolution; International Journal of Organic Evolution, 2005, 59, 2656-68.	2.3	36
61	Rapid Behavioral Changes in Medaka (Oryzias latipes) Caused by Selection for Competitive and Noncompetitive Growth. Evolution; International Journal of Organic Evolution, 1991, 45, 1936.	2.3	35
62	Scaling of the variance and the quantification of resource monopolization. Behavioral Ecology, 1996, 7, 199-207.	2.2	35
63	Genetic and environmental correlates of morphological variation in a marine fish: the case of Baltic Sea herring (<i>Clupea harengus</i>). Canadian Journal of Fisheries and Aquatic Sciences, 2008, 65, 389-400.	1.4	35
64	Long term diet differences between morphs in trophically polymorphic Percichthys trucha (Pisces:) Tj ETQq0 0 0 69, 599-616.) rgBT /Ove 1.6	erlock 10 Tf 5(34
65	Contemporary effective population and metapopulation size (<i>N</i> _e and) Tj ETQq1 1 0.784314 differing in gene flow and its asymmetries. Ecology and Evolution, 2013, 3, 569-580.	ł rgBT /Ovo 1.9	erlock 10 Tf 50 34
66	Body shape variation of three species of Percichthys in relation to their coexistence in the Limay River basin, in northern Patagonia. Environmental Biology of Fishes, 1998, 53, 143-153.	1.0	33
67	Movement of a South American perch Percichthys trucha in a mountain Patagonian lake during spawning and prespawning periods. Journal of Fish Biology, 2007, 70, 215-230.	1.6	33
68	Estuarine survival and migratory behaviour of Atlantic salmon <i>Salmo salar</i> smolts. Journal of Fish Biology, 2012, 81, 1626-1645.	1.6	33
69	The effects of diadromy and its loss on genomic divergence: The case of amphidromous <i>Galaxias maculatus</i> populations. Molecular Ecology, 2019, 28, 5217-5231.	3.9	32
70	Resolving fineâ€scale population structure and fishery exploitation using sequenced microsatellites in a northern fish. Evolutionary Applications, 2020, 13, 1055-1068.	3.1	32
71	A temporal perspective on population structure and gene flow in Atlantic salmon (Salmo salar) in Newfoundland, Canada. Canadian Journal of Fisheries and Aquatic Sciences, 2010, 67, 225-242.	1.4	31
72	Correlates of estuarine survival of Atlantic salmon postsmolts from the Southern Upland, Nova Scotia, Canada. Canadian Journal of Fisheries and Aquatic Sciences, 2013, 70, 452-460.	1.4	31

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73	Evolution of Social Behavior in a Resource-Rich, Structured Environment: Selection Experiments with Medaka (Oryzias latipes). Evolution; International Journal of Organic Evolution, 1993, 47, 456.	2.3	30
74	Distinctness, phylogenetic relations and biogeography of intertidal mussels (<i>Brachidontes</i> ,) Tj ETQq0 C United Kingdom, 2013, 93, 1843-1855.	0 rgBT /Ove 0.8	erlock 10 Tf 50 30
75	Evidence supporting panmixia in Greenland halibut (<i>Reinhardtius hippoglossoides</i>) in the Northwest Atlantic. Canadian Journal of Fisheries and Aquatic Sciences, 2014, 71, 763-774.	1.4	29
76	Trophic interference by <i>Salmo trutta</i> on <i>Aplochiton zebra</i> and <i>Aplochiton taeniatus</i> in southern Patagonian lakes. Journal of Fish Biology, 2013, 82, 430-443.	1.6	28
77	Demographic and genetic factors shaping contemporary metapopulation effective size and its empirical estimation in salmonid fish. Heredity, 2011, 107, 444-455.	2.6	27
78	Echoes of a distant time: effects of historical processes on contemporary genetic patterns in <i>Galaxias platei</i> in Patagonia. Molecular Ecology, 2015, 24, 4112-4128.	3.9	25
79	Humanâ€induced habitat fragmentation effects on connectivity, diversity, and population persistence of an endemic fish, <i>Percilia irwini</i> , in the BiobÃo River basin (Chile). Evolutionary Applications, 2020, 13, 794-807.	3.1	24
80	Genetic mixed-stock analysis of Atlantic herring populations in a mixed feeding area. Marine Ecology - Progress Series, 2011, 442, 187-199.	1.9	24
81	EVOLUTION OF SOCIAL BEHAVIOR IN A RESOURCE-RICH, STRUCTURED ENVIRONMENT: SELECTION EXPERIMENTS WITH MEDAKA (<i>ORYZIAS LATIPES</i>). Evolution; International Journal of Organic Evolution, 1993, 47, 456-470.	2.3	23
82	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 December 2011 – 31 January 2012. Molecular Ecology Resources, 2012, 12, 570-572.	4.8	23
83	Genetic and phenotypic variation among <i>Galaxias maculatus</i> populations reflects contrasting landscape effects between northern and southern Patagonia. Freshwater Biology, 2013, 58, 36-49.	2.4	23
84	<scp>H</scp> ierarchical population structure and genetic diversity of lake trout (<i><scp>S</scp>alvelinus namaycush</i>) in a dendritic system in <scp>N</scp> orthern <scp>L</scp> abrador. Freshwater Biology, 2013, 58, 1903-1917.	2.4	23
85	Palaeogeography and palaeoclimatology of Patagonia: effects on biodiversity. Biological Journal of the Linnean Society, 2011, 103, 221-228.	1.6	22
86	Functional genetic diversity in an exploited marine species and its relevance to fisheries management. Proceedings of the Royal Society B: Biological Sciences, 2021, 288, 20202398.	2.6	22
87	Chromosome level reference of Atlantic halibut <i>Hippoglossushippoglossus</i> provides insight into the evolution of sexual determination systems. Molecular Ecology Resources, 2021, 21, 1686-1696.	4.8	21
88	Genetic divergence among and within Arctic char (<i>Salvelinus alpinus</i>) populations inhabiting landlocked and sea-accessible sites in Labrador, Canada. Canadian Journal of Fisheries and Aquatic Sciences, 2018, 75, 1256-1269.	1.4	20
89	Behavioural and growth responses to the intensity of intraspecific social interaction among medaka, Oryzias latipes (Temminck and Schlegel) (Pisces, Cyprinodontidae). Journal of Fish Biology, 1990, 37, 663-673.	1.6	19
90	Nuclear and mitochondrial markers reveal distinctiveness of a small population of bottlenose whales (Hyperoodon ampullatus) in the western North Atlantic. Molecular Ecology, 2006, 15, 3115-3129.	3.9	19

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91	Biocomplexity in a demersal exploited fish, white hake (<i>Urophycis tenuis</i>): depth-related structure and inadequacy of current management approaches. Canadian Journal of Fisheries and Aquatic Sciences, 2012, 69, 415-429.	1.4	19
92	Past, present, and future of a freshwater fish metapopulation in a threatened landscape. Conservation Biology, 2018, 32, 849-859.	4.7	19
93	Extensive secondary contact among three glacial lineages of Arctic Char (<i>Salvelinus alpinus</i>) in Labrador and Newfoundland. Ecology and Evolution, 2019, 9, 2031-2045.	1.9	19
94	Genomic basis of the loss of diadromy in Galaxias maculatus : Insights from reciprocal transplant experiments. Molecular Ecology, 2020, 29, 4857-4870.	3.9	19
95	Scorched mussels (<i>Brachidontes</i> spp., Bivalvia: Mytilidae) from the tropical and warmâ€ŧemperate southwestern Atlantic: the role of the Amazon River in their speciation. Ecology and Evolution, 2016, 6, 1778-1798.	1.9	18
96	A putative structural variant and environmental variation associated with genomic divergence across the Northwest Atlantic in Atlantic Halibut. ICES Journal of Marine Science, 2021, 78, 2371-2384.	2.5	18
97	The composition of adult overwintering and juvenile aggregations of Atlantic cod (<i>Gadus) Tj ETQq1 1 0.7843 Journal of Fisheries and Aquatic Sciences, 2012, 69, 307-320.</i>	14 rgBT /C 1.4	overlock 10 Th 17
98	Limited genetic parallelism underlies recent, repeated incipient speciation in geographically proximate populations of an Arctic fish (<i>Salvelinus alpinus</i>). Molecular Ecology, 2020, 29, 4280-4294.	3.9	17
99	A portrait of a sucker using landscape genetics: how colonization and life history undermine the idealized dendritic metapopulation. Molecular Ecology, 2016, 25, 4126-4145.	3.9	16
100	Investigating Diadromy in Fishes and Its Loss in an -Omics Era. IScience, 2020, 23, 101837.	4.1	16
101	Temporal stability and assignment power of adaptively divergent genomic regions between herring (<i>Clupea harengus</i>) seasonal spawning aggregations. Ecology and Evolution, 2019, 9, 500-510.	1.9	14
102	Multiple drainage reversal episodes and glacial refugia in a Patagonian fish revealed by sequenced microsatellites. Proceedings of the Royal Society B: Biological Sciences, 2020, 287, 20200468.	2.6	14
103	Population abundance in arctic grayling using genetics and closeâ€kin markâ€recapture. Ecology and Evolution, 2021, 11, 4763-4773.	1.9	14
104	Genomic basis of deepâ€water adaptation in Arctic Charr (<i>Salvelinus alpinus</i>) morphs. Molecular Ecology, 2021, 30, 4415-4432.	3.9	13
105	Genetic Causes and Consequences of Sympatric Morph Divergence in Salmonidae: A Search for Mechanisms. Annual Review of Animal Biosciences, 2022, 10, 81-106.	7.4	13
106	Long term diet differences between morphs in trophically polymorphic Percichthys trucha (Pisces:) Tj ETQq0 0 0 69, 599-616.	rgBT /Ove 1.6	rlock 10 Tf 50 12
107	Genetic and phenotypic differentiation among Galaxias maculatus populations in a Patagonian postglacial lake system. Biological Journal of the Linnean Society, 2012, 107, 368-382.	1.6	12
108	Genomic tools for management and conservation of Atlantic cod in a coastal marine protected area. Canadian Journal of Fisheries and Aquatic Sciences, 2018, 75, 1915-1925.	1.4	11

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109	Invasive species and postglacial colonization: their effects on the genetic diversity of a Patagonian fish. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20182567.	2.6	11
110	Young-of-the-year Coho Salmon Oncorhynchus kisutch recruit in fresh waters of remote Patagonian fjords in southern Chile (51°S). Biological Invasions, 2017, 19, 1127-1136.	2.4	10
111	ENVIRONMENTAL CORRELATES OF POPULATION DIFFERENTIATION IN ATLANTIC HERRING. Evolution; International Journal of Organic Evolution, 2005, 59, 2656.	2.3	8
112	Insights on the history of the scorched mussel Brachidontes rodriguezii (Bivalvia: Mytilidae) in the Southwest Atlantic: a geometric morphometrics perspective. Historical Biology, 2018, 30, 564-572.	1.4	8
113	Isolation and characterization of eight microsatellite loci for white hake (Urophycis tenuis). Molecular Ecology Notes, 2006, 6, 924-926.	1.7	7
114	Finding the "Conservation―in Conservation Genetics—Progress in Latin America: Table 1 Journal of Heredity, 2015, 106, 423-427.	2.4	6
115	Bay-scale population structure in coastal Atlantic cod in Labrador and Newfoundland, Canada. Journal of Fish Biology, 2000, 56, 431-447.	1.6	6
116	Ten novel microsatellite loci characterized for a remarkably widespread fish: Galaxias maculatus (Galaxiidae). Molecular Ecology Resources, 2009, 9, 1503-1505.	4.8	5
117	Development of 17 novel microsatellite markers for the longnose sucker (Catostomus catostomus) and successful cross-specific amplification of 14 previously developed markers from congeneric species. Conservation Genetics Resources, 2014, 6, 329-332.	0.8	5
118	Landscape, colonization, and life history: their effects on genetic diversity in four sympatric species inhabiting a dendritic system. Canadian Journal of Fisheries and Aquatic Sciences, 2019, 76, 2288-2302.	1.4	5
119	The Genomic Consistency of the Loss of Anadromy in an Arctic Fish (<i>Salvelinus alpinus</i>). American Naturalist, 2022, 199, 617-635.	2.1	5
120	Development and characterization of 36 novel microsatellite markers for lake chub (Couesius) Tj ETQq0 0 0 rgBT	- /Qvgrlock	2 10 Tf 50 302
121	Historical and Contemporary Diversity of Galaxiids in South America: Biogeographic and Phylogenetic Perspectives. Diversity, 2020, 12, 304.	1.7	4
122	Juvenile salmon presence effects on the diet of native Puye Galaxias maculatus in lakes and estuaries of Patagonian fjords. Biological Invasions, 2022, 24, 81-92.	2.4	4
123	Development of 26 novel microsatellite makers for the round whitefish (Prosopium cylindraceum) and successful polymorphic cross-specific amplification of seven previously developed salmonid markers. Conservation Genetics Resources, 2014, 6, 1023-1026.	0.8	3
124	Crustacean zooplankton assemblages in inland waters of southern Patagonia (Alacalufes National) Tj ETQqO 0 0	rgBT_/Ove	rloçk 10 Tf 50
125	High prevalence of multiple paternity in the deep-sea shrimp Acanthephyra pelagica. Marine Biology, 2016, 163, 1.	1.5	3

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127	Connectivity, diversity, and hybridization between two endemic fish species (Percilia spp.) in a complex temperate landscape. Conservation Genetics, 2022, 23, 23-33.	1.5	3
128	Characterization of a major histocompatibility class II <i>A </i> gene (<i>Clhaâ€DAA</i>) with an embedded microsatellite marker in Atlantic herring (<i>Clupea harengus </i> L.). Journal of Fish Biology, 2008, 73, 367-381.	1.6	2
129	Isolation and characterization of 13 microsatellite loci for <i>Percichthys trucha</i> (Percichthyidae). Molecular Ecology Resources, 2008, 8, 907-909.	4.8	2
130	Development and design of five multi-locus microsatellite PCR panels for population genetic surveys of white hake (Urophycis tenuis) in the Northwest Atlantic. Conservation Genetics Resources, 2010, 2, 45-49.	0.8	2
131	Isolation and characterization of 26 novel microsatellite loci in the deep-sea shrimp Acanthephyra pelagica. Conservation Genetics Resources, 2014, 6, 731-733.	0.8	2
132	Development and characterization of 15 novel microsatellite markers for the freshwater fish Galaxias platei. Conservation Genetics Resources, 2014, 6, 899-901.	0.8	2
133	Behaviour, growth, and intensity of social interaction among fish. Canadian Journal of Statistics, 1992, 20, 221-227.	0.9	1
	Development of 13 polymorphic microsatellite loci for the Greenland Halibut (Reinhardtius) Ti ETOo0 0 0 rgBT/C	verlock 1) Tf 50 462 To

134 Development of 13 polymorphic microsatellite loci for the Greenland Halibut (Reinhardtius) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 462 To

135	Common inheritance. Nature, 1993, 365, 688-688.	27.8	0
136	Seventeen novel microsatellite markers for the southern hemisphere fish Galaxias maculatus. Conservation Genetics Resources, 2014, 6, 925-927.	0.8	0
137	Panmixia and incomplete lineage sorting can give similar mtDNA signals: response to comment by Carr and Davidson (2015). Canadian Journal of Fisheries and Aquatic Sciences, 2015, 72, 1122-1123.	1.4	0