

Martin Ian Taylor

List of Publications by Year in descending order

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53
papers

2,069
citations

346980

22
h-index

274796

44
g-index

54
all docs

54
docs citations

54
times ranked

3924
citing authors

#	ARTICLE	IF	CITATIONS
1	Transposable element annotation in non-model species: The benefits of species-specific repeat libraries using semi-automated EDTA and DeepTE de novo pipelines. <i>Molecular Ecology Resources</i> , 2022, 22, 823-833.	2.2	10
2	Satyrization in <i>Drosophila</i> fruitflies. <i>Journal of Evolutionary Biology</i> , 2021, 34, 319-330.	0.8	3
3	No severe genetic bottleneck in a rapidly range-expanding bumblebee pollinator. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2021, 288, 20202639.	1.2	4
4	Contemporary evolution of the innate immune receptor gene <i>TLR3</i> in an isolated vertebrate population. <i>Molecular Ecology</i> , 2021, 30, 2528-2542.	2.0	15
5	Ageing European lobsters (<i>Homarus gammarus</i>) using DNA methylation of evolutionarily conserved ribosomal DNA. <i>Evolutionary Applications</i> , 2021, 14, 2305-2318.	1.5	11
6	Domestication-induced reduction in eye size revealed in multiple common garden experiments: The case of Atlantic salmon (<i>Salmo salar</i> L.). <i>Evolutionary Applications</i> , 2021, 14, 2319-2332.	1.5	4
7	Hematocrit, age, and survival in a wild vertebrate population. <i>Ecology and Evolution</i> , 2021, 11, 214-226.	0.8	12
8	Comparative genetic stock structure in three species of commercially exploited Indo-Malay Carangidae (Teleostei, Perciformes). <i>Journal of Fish Biology</i> , 2020, 96, 337-349.	0.7	15
9	Disentangling the effects of sex, life history and genetic background in Atlantic salmon: growth, heart and liver under common garden conditions. <i>Royal Society Open Science</i> , 2020, 7, 200811.	1.1	4
10	Help or hindrance? The evolutionary impact of whole-genome duplication on immunogenetic diversity and parasite load. <i>Ecology and Evolution</i> , 2020, 10, 13949-13956.	0.8	0
11	Pollution control can help mitigate future climate change impact on European grayling in the UK. <i>Diversity and Distributions</i> , 2020, 26, 517-532.	1.9	4
12	The sustainable use and exploitation of fishes. <i>Journal of Fish Biology</i> , 2019, 94, 833-836.	0.7	2
13	Inclusion of jellyfish in 30+ years of Ecopath with Ecosim models. <i>ICES Journal of Marine Science</i> , 2019, 76, 1941-1950.	1.2	19
14	OSF-Builder: A New Tool for Constructing and Representing Evolutionary Histories Involving Introgression. <i>Systematic Biology</i> , 2019, 68, 717-729.	2.7	2
15	Evolutionary drivers of kype size in Atlantic salmon (<i>Salmo salar</i>): domestication, age and genetics. <i>Royal Society Open Science</i> , 2019, 6, 190021.	1.1	16
16	Where are they all from? " sources and sustainability in the ornamental freshwater fish trade. <i>Journal of Fish Biology</i> , 2019, 94, 909-916.	0.7	75
17	Cryptic diets of forage fish: jellyfish consumption observed in the Celtic Sea and western English Channel. <i>Journal of Fish Biology</i> , 2019, 94, 1026-1032.	0.7	6
18	How quantitative is metabarcoding: A meta-analytical approach. <i>Molecular Ecology</i> , 2019, 28, 420-430.	2.0	274

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19	Whole genome duplication and transposable element proliferation drive genome expansion in <i>Corydoradinae</i> catfishes. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2018, 285, 20172732.	1.2	32
20	Neutral variation does not predict immunogenetic variation in the European grayling (<i>Thymallus</i>). <i>Evolutionary Ecology</i> , 2017, 31, 101-111.	2.0	3
21	The oddity effect drives prey choice but not necessarily attack time. <i>Ethology</i> , 2018, 124, 496-503.	0.5	15
22	Evolution: Fangtastic Venoms Underpin Parasitic Mimicry. <i>Current Biology</i> , 2017, 27, R295-R298.	1.8	2
23	Jellyfish on the menu: mtDNA assay reveals scyphozoan predation in the Irish Sea. <i>Royal Society Open Science</i> , 2017, 4, 171421.	1.1	25
24	A common garden design reveals population-specific variability in potential impacts of hybridization between populations of farmed and wild Atlantic salmon (<i>Salmo salar</i> L.). <i>Evolutionary Applications</i> , 2016, 9, 435-449.	1.5	23
25	Does density influence relative growth performance of farm, wild and F ₁ hybrid Atlantic salmon in semi-natural and hatchery common garden conditions?. <i>Royal Society Open Science</i> , 2016, 3, 160152.	1.1	10
26	Plasticity in growth of farmed and wild Atlantic salmon: is the increased growth rate of farmed salmon caused by evolutionary adaptations to the commercial diet?. <i>BMC Evolutionary Biology</i> , 2016, 16, 264.	3.2	34
27	Combination of genetics and spatial modelling highlights the sensitivity of cod (<i>Gadus morhua</i>) population diversity in the North Sea to distributions of fishing. <i>ICES Journal of Marine Science</i> , 2014, 71, 794-807.	1.2	45
28	Development of microsatellite markers from 454 transcriptome derived sequences for the scallop <i>Pecten maximus</i> . <i>Conservation Genetics Resources</i> , 2013, 5, 663-666.	0.4	5
29	Potential effects of stock enhancement with hatchery-reared seed on genetic diversity and effective population size. <i>Canadian Journal of Fisheries and Aquatic Sciences</i> , 2013, 70, 330-338.	0.7	17
30	Genomics in marine monitoring: New opportunities for assessing marine health status. <i>Marine Pollution Bulletin</i> , 2013, 74, 19-31.	2.3	196
31	Experimental harvesting of fish populations drives genetically based shifts in body size and maturation. <i>Frontiers in Ecology and the Environment</i> , 2013, 11, 181-187.	1.9	93
32	A genomic island linked to ecotype divergence in Atlantic cod. <i>Molecular Ecology</i> , 2013, 22, 2653-2667.	2.0	137
33	Spawning of Kattegat cod (<i>Gadus morhua</i>)—Mapping spatial distribution by egg surveys. <i>Fisheries Research</i> , 2013, 147, 63-71.	0.9	10
34	Can Long-Range PCR Be Used to Amplify Genetically Divergent Mitochondrial Genomes for Comparative Phylogenetics? A Case Study within Spiders (Arthropoda: Araneae). <i>PLoS ONE</i> , 2013, 8, e62404.	1.1	14
35	DNA Barcoding Reveals Cryptic Diversity within Commercially Exploited Indo-Malay Carangidae (Teleostei: Perciformes). <i>PLoS ONE</i> , 2012, 7, e49623.	1.1	74
36	Environmental selection on transcriptome-derived SNPs in a high gene flow marine fish, the Atlantic herring (<i>Clupea harengus</i>). <i>Molecular Ecology</i> , 2012, 21, 3686-3703.	2.0	205

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37	Novel Tools for Conservation Genomics: Comparing Two High-Throughput Approaches for SNP Discovery in the Transcriptome of the European Hake. PLoS ONE, 2011, 6, e28008.	1.1	59
38	Competition and phylogeny determine community structure in M ¹ / ₄ llerian co-mimics. Nature, 2011, 469, 84-88.	13.7	105
39	Substantial genetic structure among stocked and native populations of the European grayling (<i>Thymallus thymallus</i> , Salmonidae) in the United Kingdom. Conservation Genetics, 2011, 12, 731-744.	0.8	19
40	A TaqMan real-time PCR based assay targeting plaice (<i>Pleuronectes platessa</i> L.) DNA to detect predation by the brown shrimp (<i>Crangon crangon</i> L.) and the shore crab (<i>Carcinus maenas</i> L.)—Assay development and validation. Journal of Experimental Marine Biology and Ecology, 2010, 391, 178-189.	0.7	33
41	Development and Application of Microsatellites in <i>Carcinus maenas</i> : Genetic Differentiation between Northern and Central Portuguese Populations. PLoS ONE, 2009, 4, e7268.	1.1	20
42	Development of 30 microsatellite markers for dab (<i>Limanda limanda</i> L.): a key UK marine biomonitoring species. Molecular Ecology Resources, 2009, 9, 951-955.	2.2	6
43	Isolation and characterization of 28 new microsatellite markers for European flounder (<i>Platichthys flesus</i> L.). Molecular Ecology Resources, 2009, 9, 1065-1068.	2.2	2
44	Mapping the spawning grounds of North Sea cod (<i>Gadus morhua</i>) by direct and indirect means. Proceedings of the Royal Society B: Biological Sciences, 2008, 275, 1543-1548.	1.2	66
45	Isolation and characterization of polymorphic microsatellite loci in the gudgeon, <i>Gobio gobio</i> (Cyprinidae). Molecular Ecology Notes, 2006, 6, 387-389.	1.7	3
46	Characterization of microsatellite loci in <i>Brachyderes rugatus</i> , the Canary Islands pine weevil (Coleoptera: Curculionidae). Molecular Ecology Notes, 2006, 6, 820-822.	1.7	4
47	Benefits and costs to mussels from ejecting bitterling embryos: a test of the evolutionary equilibrium hypothesis. Animal Behaviour, 2005, 70, 31-37.	0.8	31
48	Variation in habitat preference and population structure among three species of the Lake Malawi cichlid genus <i>Protomelas</i> . Molecular Ecology, 2004, 13, 2691-2697.	2.0	26
49	Title is missing!. Conservation Genetics, 2003, 4, 129-140.	0.8	29
50	Evidence for genetic monogamy and female-biased dispersal in the biparental mouthbrooding cichlid <i>Eretmodus cyanostictus</i> from Lake Tanganyika. Molecular Ecology, 2003, 12, 3173-3177.	2.0	53
51	Species-specific TaqMan probes for simultaneous identification of (<i>Gadus morhua</i> L.), haddock (<i>Melanogrammus aeglefinus</i> L.) and whiting (<i>Merlangius merlangus</i> L.). Molecular Ecology Notes, 2002, 2, 599-601.	1.7	98
52	Characterization of tetranucleotide microsatellite loci in a Lake Victorian, haplochromine cichlid fish: a <i>Pundamilia pundamilia</i> x <i>Pundamilia nyererei</i> hybrid. Molecular Ecology Notes, 2002, 2, 443-445.	1.7	42
53	Sexual selection, parasites and bower height skew in a bower-building cichlid fish. Animal Behaviour, 1998, 56, 379-384.	0.8	57