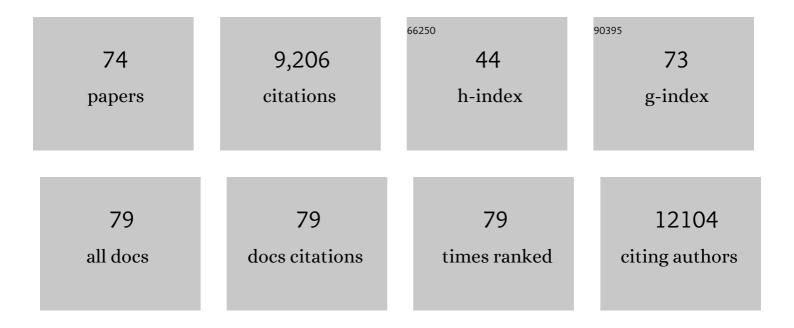
## Gary R Carvalho

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10481043/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Environmental DNA provides higher resolution assessment of riverine biodiversity and ecosystem function via spatio-temporal nestedness and turnover partitioning. Communications Biology, 2021, 4, 512.	2.0	30
2	Comparative genetic stock structure in three species of commercially exploited Indoâ€Malay Carangidae (Teleosteii, Perciformes). Journal of Fish Biology, 2020, 96, 337-349.	0.7	15
3	Executing multi-taxa eDNA ecological assessment via traditional metrics and interactive networks. Science of the Total Environment, 2020, 729, 138801.	3.9	51
4	Performance of amplicon and shotgun sequencing for accurate biomass estimation in invertebrate community samples. Molecular Ecology Resources, 2018, 18, 1020-1034.	2.2	104
5	Stepping stones to isolation: Impacts of a changing climate on the connectivity of fragmented fish populations. Evolutionary Applications, 2018, 11, 978-994.	1.5	18
6	Acidity promotes degradation of multi-species environmental DNA in lotic mesocosms. Communications Biology, 2018, 1, 4.	2.0	219
7	Annual time-series analysis of aqueous eDNA reveals ecologically relevant dynamics of lake ecosystem biodiversity. Nature Communications, 2017, 8, 14087.	5.8	229
8	Recommendations for developing and applying genetic tools to assess and manage biological invasions in marine ecosystems. Marine Policy, 2017, 85, 54-64.	1.5	74
9	Plio-Pleistocene phylogeography of the Southeast Asian Blue Panchax killifish, Aplocheilus panchax. PLoS ONE, 2017, 12, e0179557.	1.1	22
10	No loss of genetic diversity in the exploited and recently collapsed population of Bay of Biscay anchovy (Engraulis encrasicolus, L.). Marine Biology, 2016, 163, 1.	0.7	14
11	Populationâ€level consequences for wild fish exposed to sublethal concentrations of chemicals – a critical review. Fish and Fisheries, 2016, 17, 545-566.	2.7	119
12	Gene-associated markers can assign origin in a weakly structured fish, Atlantic herring. ICES Journal of Marine Science, 2015, 72, 1790-1801.	1.2	50
13	Oceanography and life history predict contrasting genetic population structure in two <scp>A</scp> ntarctic fish species. Evolutionary Applications, 2015, 8, 486-509.	1.5	46
14	Fish Product Mislabelling: Failings of Traceability in the Production Chain and Implications for Illegal, Unreported and Unregulated (IUU) Fishing. PLoS ONE, 2014, 9, e98691.	1.1	128
15	Combination of genetics and spatial modelling highlights the sensitivity of cod (Gadus morhua) population diversity in the North Sea to distributions of fishing. ICES Journal of Marine Science, 2014, 71, 794-807.	1.2	45
16	A reliable DNA barcode reference library for the identification of the North European shelf fish fauna. Molecular Ecology Resources, 2014, 14, 1060-1071.	2.2	93
17	Environmental DNA for wildlife biology and biodiversity monitoring. Trends in Ecology and Evolution, 2014, 29, 358-367.	4.2	920
18	Outlier <scp>SNP</scp> markers reveal fineâ€scale genetic structuring across <scp>E</scp> uropean hake populations ( <i><scp>M</scp>erluccius merluccius</i> ). Molecular Ecology, 2014, 23, 118-135.	2.0	171

GARY R CARVALHO

#	Article	IF	CITATIONS
19	Metagenetic analysis of patterns of distribution and diversity of marine meiobenthic eukaryotes. Global Ecology and Biogeography, 2014, 23, 1293-1302.	2.7	96
20	Borneo and Indochina are Major Evolutionary Hotspots for Southeast Asian Biodiversity. Systematic Biology, 2014, 63, 879-901.	2.7	283
21	Investigating the molecular systematic relationships amongst selected <i><scp>P</scp>lesionika</i> ( <scp>D</scp> ecapoda: <scp>P</scp> andalidae) from the <scp>N</scp> ortheast <scp>A</scp> tlantic and <scp>M</scp> editerranean <scp>S</scp> ea. Marine Ecology, 2013, 34, 157-170.	0.4	14
22	Paleo-Drainage Basin Connectivity Predicts Evolutionary Relationships across Three Southeast Asian Biodiversity Hotspots. Systematic Biology, 2013, 62, 398-410.	2.7	78
23	Experimental harvesting of fish populations drives genetically based shifts in body size and maturation. Frontiers in Ecology and the Environment, 2013, 11, 181-187.	1.9	93
24	A genomic island linked to ecotype divergence in <scp>A</scp> tlantic cod. Molecular Ecology, 2013, 22, 2653-2667.	2.0	137
25	Population genetics provides new insights into biomarker prevalence in dab ( <i><scp>L</scp>imanda) Tj ETQq1 891-909.</i>	1 0.78431 1.5	.4 rgBT /Over 5
26	SNP Discovery in European Anchovy (Engraulis encrasicolus, L) by High-Throughput Transcriptome and Genome Sequencing. PLoS ONE, 2013, 8, e70051.	1.1	38
27	Gene-associated markers provide tools for tackling illegal fishing and false eco-certification. Nature Communications, 2012, 3, 851.	5.8	199
28	DNA Barcoding Reveals Cryptic Diversity within Commercially Exploited Indo-Malay Carangidae (Teleosteii: Perciformes). PLoS ONE, 2012, 7, e49623.	1.1	74
29	A Ranking System for Reference Libraries of DNA Barcodes: Application to Marine Fish Species from Portugal. PLoS ONE, 2012, 7, e35858.	1.1	89
30	SNP Discovery Using Next Generation Transcriptomic Sequencing in Atlantic Herring (Clupea) Tj ETQq0 0 0 rgBT	/Oyerlock	10, Țf 50 302
31	Environmental selection on transcriptomeâ€derived SNPs in a high gene flow marine fish, the Atlantic herring ( <i>Clupea harengus</i> ). Molecular Ecology, 2012, 21, 3686-3703.	2.0	205
32	Biogeography of the Indo-Australian Archipelago. Annual Review of Ecology, Evolution, and Systematics, 2011, 42, 205-226.	3.8	400
33	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
34	Anonymous nuclear markers for halfbeak fishes of the genus Hemirhamphodon. Conservation Genetics Resources, 2011, 3, 155-157.	0.4	2
35	Systematic and Evolutionary Insights Derived from mtDNA COI Barcode Diversity in the Decapoda (Crustacea: Malacostraca). PLoS ONE, 2011, 6, e19449.	1.1	139
36	Multigene Molecular Systematics Confirm Species Status of Morphologically Convergent Pagurus Hermit Crabs. PLoS ONE, 2011, 6, e28233.	1.1	22

GARY R CARVALHO

#	Article	IF	CITATIONS
37	Anonymous nuclear markers for SouthEast Asian halfbeak fishes (Dermogenys). Conservation Genetics Resources, 2010, 2, 325-327.	0.4	4
38	New insights into molecular evolution: prospects from the Barcode of Life Initiative (BOLI). Theory in Biosciences, 2010, 129, 149-157.	0.6	22
39	Second-generation environmental sequencing unmasks marine metazoan biodiversity. Nature Communications, 2010, 1, 98.	5.8	321
40	Genomic signatures of local directional selection in a high gene flow marine organism; the Atlantic cod (Gadus morhua). BMC Evolutionary Biology, 2009, 9, 276.	3.2	198
41	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.	1.1	183
42	Molecular phylogeny supports division of the â€~cosmopolitan' taxon Celleporella (Bryozoa;) Tj ETQq0 0 0 rg	gBT_/Overlo	ock 10 Tf 50 5
43	Genetic homogeneity among breeding grounds and nursery areas of an exploited Lake Malawi cichlid fish. Freshwater Biology, 2008, 53, 1823-1831.	1.2	10
44	Paradigm shifts in marine fisheries genetics: ugly hypotheses slain by beautiful facts. Fish and Fisheries, 2008, 9, 333-362.	2.7	492
45	The Barcode of Life Initiative: synopsis and prospective societal impacts of DNA barcoding of Fish. Genomics Society and Policy, 2007, 3, 1.	0.2	61
46	Age of Cichlids: New Dates for Ancient Lake Fish Radiations. Molecular Biology and Evolution, 2007, 24, 1269-1282.	3.5	268
47	Mating trials validate the use of DNA barcoding to reveal cryptic speciation of a marine bryozoan taxon. Proceedings of the Royal Society B: Biological Sciences, 2007, 274, 199-207.	1.2	96
48	Validation of the barcoding gene COI for use in forensic genetic species identification. Forensic Science International, 2007, 173, 1-6.	1.3	278
49	Concordance of allozyme and microsatellite differentiation in a marine fish, but evidence of selection at a microsatellite locus. Molecular Ecology, 2007, 16, 1135-1147.	2.0	68
50	Mitochondrial DNA phylogeography and mating compatibility reveal marked genetic structuring and speciation in the NE Atlantic bryozoan Celleporella hyalina. Molecular Ecology, 2007, 16, 2173-2188.	2.0	49
51	Association between Growth andPan I*Genotype within Atlantic Cod Full-Sibling Families. Transactions of the American Fisheries Society, 2006, 135, 241-250.	0.6	33
52	Power for detecting genetic divergence: differences between statistical methods and marker loci. Molecular Ecology, 2006, 15, 2031-2045.	2.0	215
53	Biocomplexity in a highly migratory pelagic marine fish, Atlantic herring. Proceedings of the Royal Society B: Biological Sciences, 2006, 273, 1459-1464.	1.2	205
54	ENVIRONMENTAL CORRELATES OF POPULATION DIFFERENTIATION IN ATLANTIC HERRING. Evolution; International Journal of Organic Evolution, 2005, 59, 2656-2668.	1.1	537

#	Article	IF	CITATIONS
55	Environmental correlates of population differentiation in Atlantic herring. Evolution; International Journal of Organic Evolution, 2005, 59, 2656-68.	1.1	36

## Temporal analysis of archived samples indicates marked genetic changes in declining North Sea cod () Tj ETQq0 0 0 rgBT /Overlock 10 Tr 206

57	Loss of microsatellite diversity and low effective population size in an overexploited population of New Zealand snapper (Pagrus auratus). Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 11742-11747.	3.3	441
58	SPECIATION IN ANCIENT CRYPTIC SPECIES COMPLEXES: EVIDENCE FROM THE MOLECULAR PHYLOGENY OF BRACHIONUS PLICATILIS (ROTIFERA). Evolution; International Journal of Organic Evolution, 2002, 56, 1431-1444.	1.1	331
59	Title is missing!. Journal of Paleolimnology, 2002, 27, 481-486.	0.8	10
60	Phylogeography and regional endemism of a passively dispersing zooplankter: mitochondrial DNA variation in rotifer resting egg banks. Proceedings of the Royal Society B: Biological Sciences, 2000, 267, 2189-2197.	1.2	134
61	Molecular markers and the species concept: New techniques to resolve old disputes?. Reviews in Fish Biology and Fisheries, 1999, 9, 379-382.	2.4	14
62	Molecular Genetic Analysis of Atlanto-Scandian Herring (Clupea Harengus) Populations Using Allozymes and Mitochondrial Dna Markers. Journal of the Marine Biological Association of the United Kingdom, 1998, 78, 269-283.	0.4	42
63	Advances in the molecular analysis of fish population structure. Italian Journal of Zoology, 1998, 65, 21-33.	0.6	74
64	Sex Ratio Variation in the Intertidal Isopod, Jaera Albifrons. Journal of the Marine Biological Association of the United Kingdom, 1996, 76, 825-828.	0.4	6
65	Artificial introductions, evolutionary change and population differentiation in Trinidadian guppies (Poecilia reticulata: Poeciliidae). Biological Journal of the Linnean Society, 1996, 57, 219-234.	0.7	37
66	Detection of High Levels of Genetic Relatedness in Rock-Populations of an Intertidal Isopod Using DNA Fingerprinting. Journal of the Marine Biological Association of the United Kingdom, 1995, 75, 967-976.	0.4	6
67	Genetic impacts of fish introductions: a perspective on African lakes. , 1995, , 457-493.		8
68	Evolution of adaptive variation in antipredator behaviour. Marine and Freshwater Behaviour and Physiology, 1993, 23, 29-44.	0.9	43
69	Resting eggs of lake- Daphnia I. Distribution, abundance and hatching of eggs collected from various depths in lake sediments. Freshwater Biology, 1989, 22, 459-470.	1.2	125
70	Resting eggs of lake-Daphania II. In situ observations on the hatching of eggs and their contribution to population and community structure. Freshwater Biology, 1989, 22, 471-478.	1.2	86
71	Haemoglobin synthesis in Daphnia magna Straus (Crustacea: Cladocera): ecological differentiation between neighbouring populations. Freshwater Biology, 1984, 14, 501-506.	1.2	16
72	The effect of food availability, female culture-density and photoperiod on ephippia production in Daphnia magna Straus (Crustacea: Cladocera). Freshwater Biology, 1983, 13, 37-46.	1.2	167

#	Article	IF	CITATIONS
73	Patterns of self compatibility, inbreeding depression, outcrossing, and sex allocation in a marine bryozoan suggest the predominating influence of sperm competition. Biological Journal of the Linnean Society, 0, 98, 519-531.	0.7	11

Molecular biogeography and phylogeography of the freshwater fauna of the Indo-Australian Archipelago. , 0, , 316-347.

5