

Global population structure of the tope (<i>Galeorhinus</i> mitochondrial control region sequence data

Molecular Ecology

18, 545-552

DOI: [10.1111/j.1365-294x.2008.04047.x](https://doi.org/10.1111/j.1365-294x.2008.04047.x)

Citation Report

#	ARTICLE	IF	CITATIONS
1	Tracking the fin trade: genetic stock identification in western Atlantic scalloped hammerhead sharks <i>Sphyrna lewini</i> . <i>Endangered Species Research</i> , 2009, 9, 221-228.	1.2	67
2	DNA Forensic Applications in Shark Management and Conservation. <i>Marine Biology</i> , 2010, , 593-610.	0.1	6
3	Editorial and retrospective 2010. <i>Molecular Ecology</i> , 2010, 19, 1-22.	2.0	11
4	Global population structure of the spiny dogfish <i>Squalus acanthias</i> , a temperate shark with an antitropical distribution. <i>Molecular Ecology</i> , 2010, 19, 1651-1662.	2.0	71
5	Is multiple mating beneficial or unavoidable? Low multiple paternity and genetic diversity in the shortspine spurdog <i>Squalus mitsukurii</i> . <i>Marine Ecology - Progress Series</i> , 2010, 403, 255-267.	0.9	63
6	Phylogeography of the copper shark (<i>Carcharhinus brachyurus</i>) in the southern hemisphere: implications for the conservation of a coastal apex predator. <i>Marine and Freshwater Research</i> , 2011, 62, 861.	0.7	40
7	Phylogeography and conservation of the bull shark (<i>Carcharhinus leucas</i>) inferred from mitochondrial and microsatellite DNA. <i>Conservation Genetics</i> , 2011, 12, 371-382.	0.8	106
8	Phylogeography and genetic population structure of Caribbean sharpnose shark <i>Rhizoprionodon porosus</i> . <i>Reviews in Fish Biology and Fisheries</i> , 2011, 21, 799-814.	2.4	19
9	A DNA Sequence-Based Approach To the Identification of Shark and Ray Species and Its Implications for Global Elasmobranch Diversity and Parasitology. <i>Bulletin of the American Museum of Natural History</i> , 2012, 367, 1-262.	1.2	352
10	Population Structure and Phylogeography of the Short-Tailed Stingray, <i>Dasyatis brevicaudata</i> (Hutton) Tj ETQq1 1 0.784314 r _g BT /Over	1.0	32
11	Inclusion of South American samples reveals new population structuring of the blacktip shark (<i>Carcharhinus limbatus</i>) in the western Atlantic. <i>Genetics and Molecular Biology</i> , 2012, 35, 752-760.	0.6	21
12	Molecular markers: progress and prospects for understanding reproductive ecology in elasmobranchs. <i>Journal of Fish Biology</i> , 2012, 80, 1120-1140.	0.7	55
13	Oceanic dispersal in a sedentary reef shark (<i>Triaenodon obesus</i>): genetic evidence for extensive connectivity without a pelagic larval stage. <i>Journal of Biogeography</i> , 2012, 39, 1144-1156.	1.4	50
14	A review of the application of molecular genetics for fisheries management and conservation of sharks and rays. <i>Journal of Fish Biology</i> , 2012, 80, 1789-1843.	0.7	190
15	Retrospective coalescent methods and the reconstruction of metapopulation histories in the sea. <i>Evolutionary Ecology</i> , 2012, 26, 291-315.	0.5	29
16	Effects of species biology on the historical demography of sharks and their implications for likely consequences of contemporary climate change. <i>Conservation Genetics</i> , 2013, 14, 125-144.	0.8	30
17	Multiple substitutions and reduced genetic variability in sharks. <i>Biochemical Systematics and Ecology</i> , 2013, 49, 21-29.	0.6	16
18	Historical demography and genetic differentiation inferred from the mitochondrial DNA of the silky shark (<i>Carcharhinus falciformis</i>) in the Pacific Ocean. <i>Fisheries Research</i> , 2013, 147, 36-46.	0.9	24

#	ARTICLE	IF	CITATIONS
19	Population Expansion and Genetic Structure in <i>Carcharhinus brevipinna</i> in the Southern Indo-Pacific. PLoS ONE, 2013, 8, e75169.	1.1	21
20	Extensive genetic population structure in the Indo-Pacific West Pacific spot-tail shark, <i>Carcharhinus sorrah</i> . Bulletin of Marine Science, 2014, 90, 427-454.	0.4	23
21	Frequency of multiple paternity varies between two populations of brown smoothhound shark, <i>Mustelus henlei</i> . Marine Biology, 2014, 161, 797-804.	0.7	20
22	Strong Population Structure and Shallow Mitochondrial Phylogeny in the Banded Guitarfish, <i>Zapteryx exasperata</i> (Jordan y Gilbert, 1880), from the Northern Mexican Pacific. Journal of Heredity, 2014, 105, 91-100.	1.0	25
23	The effect of biogeographic and phylogeographic barriers on gene flow in the brown smoothhound shark, <i>Mustelus henlei</i> , in the northeastern Pacific. Ecology and Evolution, 2015, 5, 1585-1600.	0.8	27
24	Microsatellite loci confirm a lack of population connectivity among globally distributed populations of the tope shark <i>Galeorhinus galeus</i> (Triakidae). Journal of Fish Biology, 2015, 87, 371-385.	0.7	14
25	A Tale of Three Tails: Cryptic Speciation in a Globally Distributed Marine Fish of the Genus <i>Seriola</i> . Copeia, 2015, 103, 357-368.	1.4	55
26	Genetic analysis of stock structure of blue shark (<i>Prionace glauca</i>) in the north Pacific ocean. Fisheries Research, 2015, 172, 181-189.	0.9	22
27	Demographic history and the South Pacific dispersal barrier for school shark (<i>Galeorhinus galeus</i>) inferred by mitochondrial DNA and microsatellite DNA mark. Fisheries Research, 2015, 167, 132-142.	0.9	15
28	Identification and Distribution of Morphologically Conserved Smoothhound Sharks in the Northern Gulf of Mexico. Transactions of the American Fisheries Society, 2015, 144, 1301-1310.	0.6	10
29	Molecular species identification and population genetics of chondrichthyans in South Africa: current challenges, priorities and progress. African Zoology, 2015, 50, 205-217.	0.2	17
30	Phylogeography and population structure of the red stingray, <i>Dasyatis akajei</i> inferred by mitochondrial control region. Mitochondrial DNA, 2015, 26, 505-513.	0.6	12
31	Global population genetic dynamics of a highly migratory, apex predator shark. Molecular Ecology, 2016, 25, 5312-5329.	2.0	51
32	Species identification and comparative population genetics of four coastal houndsharks based on novel NGS-mined microsatellites. Ecology and Evolution, 2017, 7, 1462-1486.	0.8	24
33	Review of Current Conservation Genetic Analyses of Northeast Pacific Sharks. Advances in Marine Biology, 2017, 77, 79-110.	0.7	12
34	Molecular research on the systemically challenging smoothhound shark genus <i>Mustelus</i> : a synthesis of the past 30 years. African Journal of Marine Science, 2017, 39, 373-387.	0.4	4
35	The importance of considering genetic diversity in shark and ray conservation policies. Conservation Genetics, 2018, 19, 501-525.	0.8	71
36	Strong trans-Pacific break and local conservation units in the Galapagos shark (<i>Carcharhinus</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10	1.2	37

#	ARTICLE	IF	CITATIONS
37	Effects of the Pleistocene on the mitochondrial population genetic structure and demographic history of the silky shark (<i>Carcharhinus falciformis</i>) in the western Atlantic Ocean. <i>Reviews in Fish Biology and Fisheries</i> , 2018, 28, 213-227.	2.4	30
38	Genetic diversity and connectivity of the megamouth shark (<i>Megachasma pelagios</i>). <i>PeerJ</i> , 2018, 6, e4432.	0.9	8
39	Population Connectivity of the Highly Migratory Shortfin Mako (<i>Isurus oxyrinchus Rafinesque 1810</i>) and Implications for Management in the Southern Hemisphere. <i>Frontiers in Ecology and Evolution</i> , 2018, 6, .	1.1	31
40	Weak population structure of the Spottail shark <i>Carcharhinus sorrah</i> and the Blacktip shark <i>C. limbatus</i> along the coasts of the Arabian Peninsula, Pakistan, and South Africa. <i>Ecology and Evolution</i> , 2018, 8, 9536-9549.	0.8	7
41	Conservation genetics of elasmobranchs of the Mexican Pacific Coast, trends and perspectives. <i>Advances in Marine Biology</i> , 2019, 83, 115-157.	0.7	5
42	Population structure, connectivity, and demographic history of an apex marine predator, the bull shark <i>Carcharhinus leucas</i> . <i>Ecology and Evolution</i> , 2019, 9, 12980-13000.	0.8	18
43	Study on population genetics of <i>Sillago aeolus</i> (Perciformes: Sillaginidae) in the Coast of China. Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2019, 30, 825-834.	0.7	3
44	Strong genetic isolation despite wide distribution in a commercially exploited coastal shark. <i>Hydrobiologia</i> , 2019, 838, 121-137.	1.0	6
45	Genetic population structure and demography of an apex predator, the tiger shark <i>Galeocerdo cuvier</i> . <i>Ecology and Evolution</i> , 2019, 9, 5551-5571.	0.8	22
46	Population genetic divergence as consequence of past range expansion of the smooth hammerhead shark <i>Sphyrna zygaena</i> . <i>Hydrobiologia</i> , 2019, 837, 31-46.	1.0	6
47	Novel multimarker comparisons address the genetic population structure of silvertip sharks (<i>Carcharhinus albimarginatus</i>). <i>Marine and Freshwater Research</i> , 2019, 70, 1007.	0.7	11
48	Phylogeography of eagle rays of the genus <i>Aetobatus</i> : <i>Aetobatus narinari</i> is restricted to the continental western Atlantic Ocean. <i>Hydrobiologia</i> , 2019, 836, 169-183.	1.0	15
49	Genetic stock structure of New Zealand fish and the use of genomics in fisheries management: an overview and outlook. <i>New Zealand Journal of Zoology</i> , 2021, 48, 1-31.	0.6	29
50	SkateBase, an elasmobranch genome project and collection of molecular resources for chondrichthyan fishes. <i>F1000Research</i> , 2014, 3, 191.	0.8	61
51	Population genetics of Southern Hemisphere tope shark (<i>Galeorhinus galeus</i>): Intercontinental divergence and constrained gene flow at different geographical scales. <i>PLoS ONE</i> , 2017, 12, e0184481.	1.1	22
52	Global phylogeography of the dusky shark <i>Carcharhinus obscurus</i> : implications for fisheries management and monitoring the shark fin trade. <i>Endangered Species Research</i> , 2011, 14, 13-22.	1.2	32
53	When two oceans meet: regional population genetics of an exploited coastal shark, <i>Mustelus mustelus</i> . <i>Marine Ecology - Progress Series</i> , 2016, 544, 183-196.	0.9	16
54	Genetics of Sharks, Skates, and Rays. , 2012, , 503-520.		4

#	ARTICLE	IF	CITATIONS
---	---------	----	-----------