

Differential gene flow patterns for two commercially exploited species of shark  
(*Galeorhinus galeus*) and common smoothhound (*Mustelus*)  
south of the west coast of South Africa

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Molecular species identification and population genetics of chondrichthyans in South Africa: current challenges, priorities and progress. <i>African Zoology</i> , 2015, 50, 205-217.	0.2	17
2	Phylogeography of the pelagic fish <i>Seriola lalandi</i> at different scales: confirmation of inter-ocean population structure and evaluation of southern African genetic diversity. <i>African Journal of Marine Science</i> , 2016, 38, 513-524.	0.4	7
3	Species identification and comparative population genetics of four coastal houndsharks based on novel NGSâ€mined microsatellites. <i>Ecology and Evolution</i> , 2017, 7, 1462-1486.	0.8	24
4	Molecular research on the systematically challenging smoothhound shark genus <i>Mustelus</i> : a synthesis of the past 30 years. <i>African Journal of Marine Science</i> , 2017, 39, 373-387.	0.4	4
5	The importance of considering genetic diversity in shark and ray conservation policies. <i>Conservation Genetics</i> , 2018, 19, 501-525.	0.8	71
6	Strong genetic isolation despite wide distribution in a commercially exploited coastal shark. <i>Hydrobiologia</i> , 2019, 838, 121-137.	1.0	6
7	Genomic resources for the spotted ragged-tooth shark <i>Carcharias taurus</i> . <i>African Journal of Marine Science</i> , 2019, 41, 115-118.	0.4	2
8	Spatio-temporal genetic variation of juvenile smooth hammerhead sharks in South Africa. <i>Marine Biology Research</i> , 2019, 15, 568-579.	0.3	7
9	Brown banded bamboo shark ( <i>Chiloscyllium punctatum</i> ) shows high genetic diversity and differentiation in Malaysian waters. <i>Scientific Reports</i> , 2021, 11, 14874.	1.6	3
10	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
11	Site fidelity and shallow genetic structure in the common smoothâ€hound shark <i>Mustelus mustelus</i> confirmed by tagâ€recapture and genetic data. <i>Journal of Fish Biology</i> , 2021, , .	0.7	4
12	Distribution and population structure of the smoothâ€hound shark, <i>Mustelus mustelus</i> (Linnaeus, 1758), across an oceanic archipelago: Combining several data sources to promote conservation. <i>Ecology and Evolution</i> , 2022, 12, .	0.8	2
13	Sharks Do Not Always Grow Slowly: Tagging Data Reveal a Different Pattern of Growth, Longevity and Maturity for Threatened Smooth-Hounds in the Central Mediterranean Sea. <i>Journal of Marine Science and Engineering</i> , 2022, 10, 1647.	1.2	3