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Genetic differentiation and phylogeography of Mediterranean-North Eastern Atlantic blue shark (*Prionace glauca*, L. 1758) using mitochondrial DNA: panmixia or complex stock structure?

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ABSTRACT
Background: The blue shark (*Prionace glauca*, Linnaeus 1758) is one of the most abundant migratory shark inhabiting all the coastal waters of the world, including the Mediterranean Sea. Its genetic structure has not been confirmed at basin and inter-basin scales. For tagging programs in the Atlantic Ocean failed to find evidence of migration of blue sharks between the Mediterranean and the adjacent Atlantic, despite the narrow region of their range, although the high connectivity within the Mediterranean basin, to date no genetic study on Mediterranean blue shark was carried out, which constitutes a significant knowledge gap, considering that this population is classified as "critically endangered", under the same status category.
Methods: Blue shark phylogeography and demography in the Mediterranean Sea and North Atlantic-Oceanic Ocean were inferred using two mitochondrial genes (12S and

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