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Characterization of genetic diversity of an invasive Lessepsian Migrant, *Nemipterus randalli* Russell, 1986 from its native and non-native regions using mitochondrial DNA marker

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Table S1. Pairwise Φ_{ST} between geographical samples of *Nemipterus randalli* collected from different locations (* Statistical significance, $P < 0.05$).

Geographical locations	Mediterranean Sea	Red Sea	Mumbai (Arabian Sea)	Mormugao (Arabian Sea)	Kochi (Arabian Sea)	Thoothukudi (Bay of Bengal)
Mediterranean Sea						
Red Sea	0.12*					
Mumbai (Arabian Sea)	0.65*	0.56*				
Mormugao (Arabian Sea)	0.65*	0.57*	0.02			
Kochi (Arabian Sea)	0.64*	0.56*	0.03	0.001		
Thoothukudi (Bay of Bengal)	0.61*	0.50*	0.04	0.05	0.08	

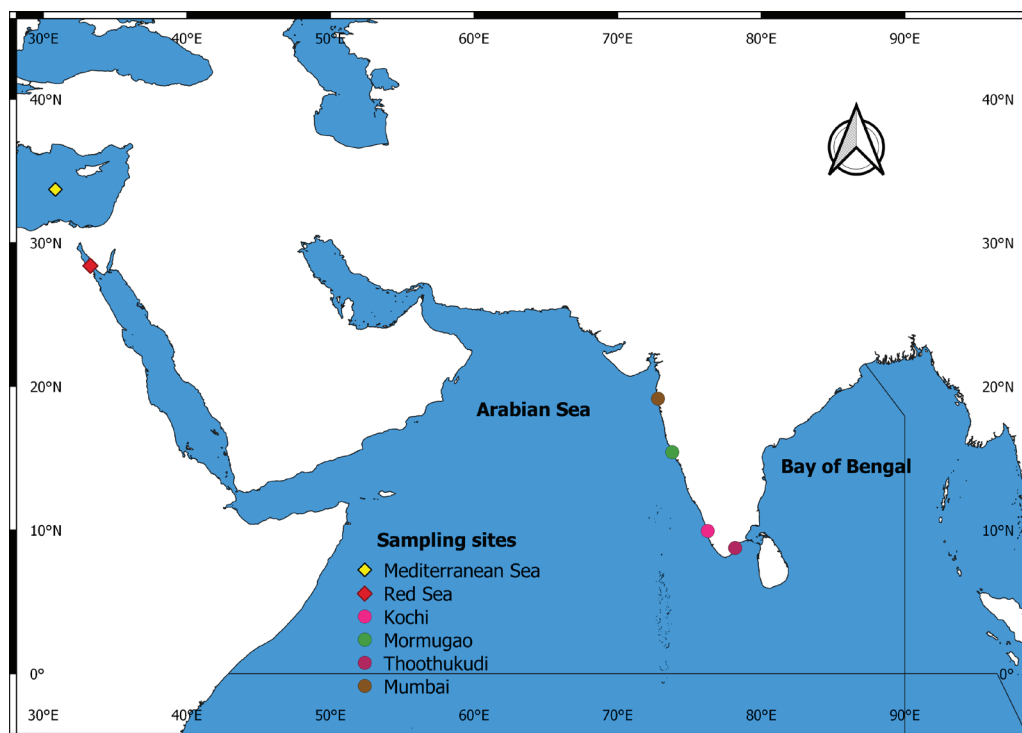


Fig. S1: Sampling locations used in the present study. *Diamond indicates the samples from Tikochinski *et al.*, 2019, circles indicate samples from the present study.

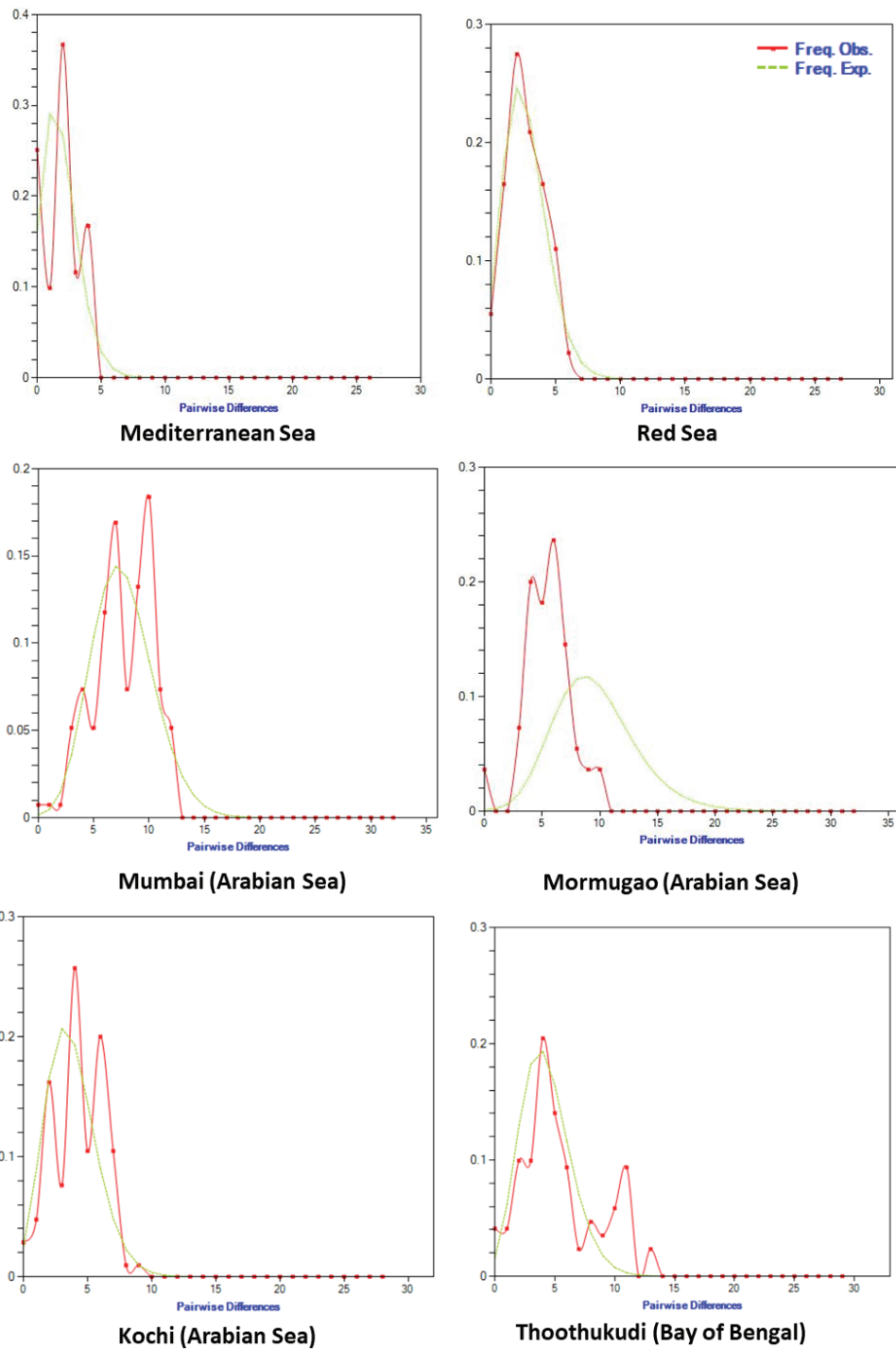


Fig. S2: Mismatch distribution curves for *Nemipterus randalli* from sampling locations.

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