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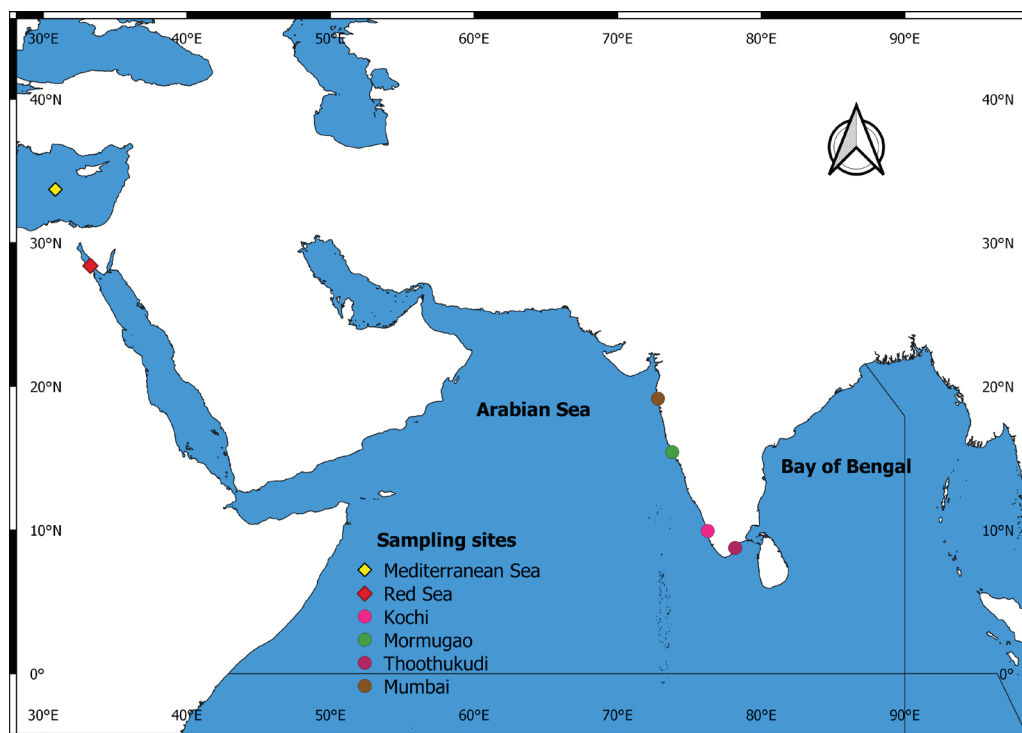
## Characterization of genetic diversity of an invasive Lessepsian Migrant, *Nemipterus randalli* Russell, 1986 from its native and non-native regions using mitochondrial DNA marker

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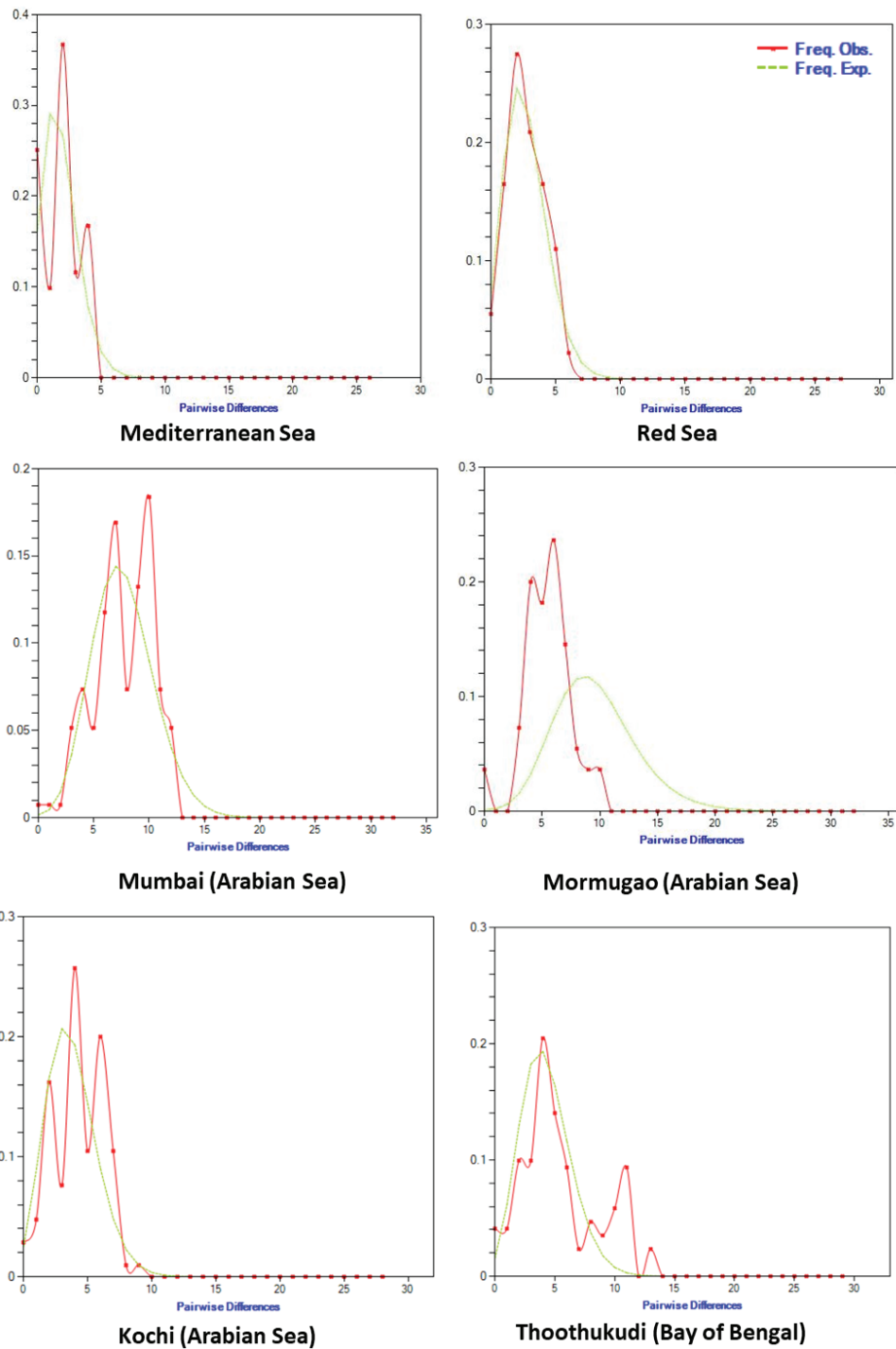
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**Table S1.** Pairwise  $\Phi_{ST}$  between geographical samples of *Nemipterus randalli* collected from different locations (\* Statistical significance,  $P < 0.05$ ).

Geographical locations	Mediterranean Sea	Red Sea	Mumbai (Arabian Sea)	Mormugao (Arabian Sea)	Kochi (Arabian Sea)	Thoothukudi (Bay of Bengal)
<b>Mediterranean Sea</b>						
<b>Red Sea</b>	0.12*					
<b>Mumbai (Arabian Sea)</b>	0.65*	0.56*				
<b>Mormugao (Arabian Sea)</b>	0.65*	0.57*	0.02			
<b>Kochi (Arabian Sea)</b>	0.64*	0.56*	0.03	0.001		
<b>Thoothukudi (Bay of Bengal)</b>	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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