

Table 2 Sites where *P. homarus* were sampled, number of individuals per sampling location (Na) and summary of genetic variability statistics; number of haplotypes (Nh), number of polymorphic sites (Np), haplotype diversity (Hd) and nucleotide diversity (Nd) in two mtDNA gene fragments.

Locus	Locality	Code	Na	Nh	Np	(Hd)	(Nd)
COI	Godawaya	GOD	6	4	19	0.866 7 +/- 0.1291	0.016 2 +/- 0.0101
	Hikkaduwa	HIK	8	8	20	1.000 0 +/- 0.0625	0.013 5 +/- 0.0081
	Kirinda	KIR	7	7	15	1.000 0 +/- 0.0764	0.011 4 +/- 0.0071
	Weligama	WEL	6	5	15	0.933 3 +/- 0.1217	0.011 4 +/- 0.0073
	Total		27	20	31		
CytB	Godawaya	GOD	10	9	43	0.977 8 +/- 0.0540	0.015 6 +/- 0.0087
	Hikkaduwa	HIK	9	9	37	1.000 0 +/- 0.0524	0.015 6 +/- 0.0088
	Kirinda	KIR	9	9	44	1.000 0 +/- 0.0524	0.014 9 +/- 0.0084
	Weligama	WEL	10	9	32	0.977 8 +/- 0.0540	0.010 6 +/- 0.0061
	Total		38	34	77		