

Table 2 Haplotype ( $h$ ) and nucleotide ( $\pi$ ) diversities (384 bp) of *E. imbricata* aggregations in the Wider Caribbean. Values between square brackets represent adult individuals

Aggregations	Genetic diversities		N
	$h \pm SD$	$\pi \pm SD$	
JR90	0.800 $\pm$ 0.078	0.0628 $\pm$ 0.0328	15
	[0.857 $\pm$ 0.050]	[0.0662 $\pm$ 0.0335]	25
JRNA04	0.714 $\pm$ 0.081	0.0493 $\pm$ 0.0252	24
JRNA05	0.733 $\pm$ 0.053	0.0532 $\pm$ 0.0271	25
JRNA04-06	0.775 $\pm$ 0.029	0.0529 $\pm$ 0.0262	54
JRA $\delta$ 04-06	[0.790 $\pm$ 0.059]	[0.0574 $\pm$ 0.0292]	25
JRA $\delta$ 04-06	[0.754 $\pm$ 0.042]	[0.0510 $\pm$ 0.0260]	26
JRA $\delta$ 04-06	[0.768 $\pm$ 0.032]	[0.0550 $\pm$ 0.0274]	51
DL90	0.814 $\pm$ 0.062	0.0604 $\pm$ 0.0309	21
	[0.245 $\pm$ 0.113]	[0.0262 $\pm$ 0.0138]	23
IJ90	0.699 $\pm$ 0.033	0.0584 $\pm$ 0.0288	83
	[0.826 $\pm$ 0.040]	[0.0575 $\pm$ 0.0290]	31
RL90	0.605 $\pm$ 0.111	0.0123 $\pm$ 0.0070	21
DO90 <sup>1</sup>	0.669 $\pm$ 0.034	0.0102 $\pm$ 0.0057	90
BA90 <sup>1</sup>	0.740 $\pm$ 0.024	0.0093 $\pm$ 0.0053	78
USVI90 <sup>1</sup>	0.750 $\pm$ 0.036	0.0108 $\pm$ 0.0060	68
	0.724 $\pm$ 0.033	0.0596 $\pm$ 0.0293	93
MI90	[0.833 $\pm$ 0.081]	[0.0725 $\pm$ 0.0382]	13
MIRes00	0.676 $\pm$ 0.050	0.0521 $\pm$ 0.0260	51
MI $\delta$ 00	[0.677 $\pm$ 0.060]	[0.0346 $\pm$ 0.0175]	54
CI00	0.666 $\pm$ 0.036	0.0559 $\pm$ 0.0276	89
FKW00	0.584 $\pm$ 0.052	0.0424 $\pm$ 0.0213	47
FPB00	0.490 $\pm$ 0.056	0.0268 $\pm$ 0.0137	91
ToL00	0.7721 $\pm$ 0.057	0.0620 $\pm$ 0.0321	17
ToW00	0.6003 $\pm$ 0.065	0.0479 $\pm$ 0.0240	49
TnC00	0.733 $\pm$ 0.033	0.0541 $\pm$ 0.0272	36

Note: JR: Jardines del Rey, Cuba; DL: Doce Leguas, Cuba; IJ: Isla de la Juventud, Cuba; RL: Rio Lagartos, Yucatan, Mexico; DO: south of Dominican Republic; BA: Bahamas; USVI: Buck Island, United States Virgin Islands; MI: Mona Island, Puerto Rico; CI: Bloody Bay and west of Grand Cayman, Cayman Islands; FKW: Key West National Wildlife Refuge, Florida, USA; FPB: Palm Beach County, Florida, USA; ToL: Tobago leeward; ToW: Tobago windward; TnC: Turks and Caicos; Ad: adult individuals, N: number of individuals with haplotypes, NA: non-adult individuals, Res: platform resident juveniles from Velez-Zuazo et al. (2008), SD: standard deviation. 00: decade of 2000, 04-06: total samples of the present study, 90: decade of 1990 from D áz-Fernández (1999); <sup>1</sup>values extracted from Bowen et al. (2007);  $\delta$ : males;  $\delta$ : females