

Table 2 Information of nine *Babylonia areolata* duplicated loci. The number of individuals analyzed (n), allele size across multi-locus range in base pairs (as) are presented

Locus	GenBank acc. no.	Repeat motif [§]	Primer sequences (5'-3')	Ta(°C)	MgCl ₂ (mM)	as	n
<i>HUBA02*</i>	FJ594999	(AC) _N (CT) _N	F: taggtgcaggccatcagag R: gtgaaacggcctcaatgtcaag	62	1.5	275-444	32
<i>HUBA04</i>	FJ595001	(GT) _N	F: gatcagcattgaaagactgaaag R: cggaacttgcaacatttaaac	60	1.5	191-311	32
<i>HUBA05</i>	FJ595002	(GT) _n (GA) _N	F: aagctcgttcaactcagctc R: gatcaatccgtgtctttctg	62	1.5	271-451	32
<i>HUBA06*</i>	FJ595003	(CA) _N (TCA) _N	F: atgccacaatgacaatcaac R: actggatgatgcgatgacaac	50	1.5	404-551	32
<i>HUBA11*</i>	FJ595008	(GT) _N (CT) _n	F: gaaactcaaacgacaccaaac R: gaaagagacgaagggtggaag	61	1.5	286-414	32
<i>HUBA12</i>	FJ595009	(GACA) _N (GA) _n	F: agtacgggttctctttgctc R: agcctcggaatctgattactg	60	1.5	277-425	30
<i>HUBA14</i>	FJ595011	(AC) _N	F: tgccaaattgtgtcaactg R: ctccgtcaacagtgaccttg	60	1.5	195-285	31
<i>HUBA16*</i>	FJ595013	(AC) _N	F: tgcctaattgggtgcaactg R: ctgctttcggtcacttttg	60	2.0	166-278	29
<i>HUBA19</i>	FJ595016	(TG) _n	F: tttctgcacatgaagaaccac R: aaaaacctccaccctttacc	53	2.0	242-508	32