

ABGD Group (P = 0.01-0.1)	Putative species	N	Intra-specific		Inter-specific		Nearest neighbor (min <i>p</i> -distance)
			min	max	min	max	
Dia01	<i>Diastylis cornuta</i>	2	0.01	0.01	0.15	0.33	Dia09
Dia03	<i>Diastylis goodsiri</i>	1	NA	NA	0.15	0.32	Dia07, Dia06
Dia04	<i>Diastylis laevis</i>	1	NA	NA	0.17	0.35	Dia01, D08
Dia05	<i>Diastylis lucifera</i>	1	NA	NA	0.23	0.31	Dia03, Dia06
Dia06	<i>Diastylis polaris</i>	8	0.00	0.00	0.15	0.33	Dia03, Dia07
Dia07	<i>Diastylis rathkei</i>	2	0.04	0.04	0.15	0.34	Dia03, Dia06
Dia08	<i>Diastylis spinulosa</i>	1	NA	NA	0.20	0.34	Dia07
Dia09	<i>Diastylis tumida</i>	2	0.00	0.00	0.15	0.32	Dia01
Dia11	<i>Diastylodes biplicatus</i>	1	NA	NA	0.18	0.37	Dia16-B
Dia12	<i>Diastylodes serratus</i>	3	0.00	0.00	0.18	0.37	Dia11
	<i>Diastylopsis</i> sp.	1	NA	NA	0.26	0.31	D17
Dia16-B	<i>Leptostylis</i> cf. <i>longimana</i> / <i>Leptostylis ampullacea</i>	14	0.00	0.01	0.21	0.33	Dia17
Dia10	<i>Diastylodes atlanticus</i>	2	0.00	0.00	0.23	0.32	Dia08
Dia15	<i>Leptostylis borealis</i>	1	NA	NA	0.24	0.37	Dia16-B, Dia17
Dia16-A	<i>Leptostylis longimana</i> (seq52/seq53)	2	0.00	0.00	0.26	0.33	Dia16-B
Dia17	<i>Leptostylis</i> sp. 1	1	NA	NA	0.21	0.33	Dia16-B
Pse01	<i>Petalosarsia declivis</i>	1	NA	NA	0.29	0.34	Dia08, Dia09

N = Number of sequences