

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	
Lam01	<i>Alamprops augustinensis</i>	1	NA	NA	0.26	0.32	Lam04, Lam09
Lam02	<i>Chalarostylis elegans</i>	2	0.00	0.00	0.18	0.31	Lam08
Lam04	<i>Hemilamprops assimilis</i>	1	NA	NA	0.13	0.34	Lam09
Lam05-A	<i>Hemilamprops cristatus</i> (seq81/ seq82/ ICE1- Lam018)	3	0.00	0.02	0.13	0.31	Lam08
Lam05-B	<i>Hemilamprops cf. cristatus</i> (ICE1-Lam002)	1	NA	NA	0.20	0.32	Lam13
Lam06	<i>Hemilamprops aff. diversus</i>	3	0.00	0.00	0.27	0.32	Lam02, Lam08
Lam07	<i>Hemilamprops pterini</i>	2	0.00	0.00	0.13	0.28	Lam05, Lam06
Lam08	<i>Hemilamprops roseus</i>	1	NA	NA	0.13	0.32	Lam04
Lam10	<i>Hemilamprops</i> sp. 2	1	NA	NA	0.24	0.30	Lam12
Lam11	<i>Hemilamprops uniplicatus</i>	2	0.00	0.00	0.17	0.30	Lam08, Lam05
Lam12	<i>Mesolamprops denticulatus</i>	1	NA	NA	0.20	0.31	Lam06
Lam13	<i>Platysympus typicus</i>	4	0.00	0.01	0.25	0.31	Lam05, Cer01
Cer01	<i>Cimmerius reticulatus</i>	1	NA	NA	0.25	0.33	Lam14

N = Number of sequences