

ABGD Group (P = 0.01-0.08)	Putative species	N	Intra-specific		Inter-specific		Nearest neighbor (min <i>p</i> -distance)
			min	max	min	max	
Nan03	<i>Campylaspis costata</i>	1	0.00	0.00	0.17	0.41	Nan05
Nan05	<i>Campylaspis horrida</i>	1	0.00	0.00	0.17	0.40	Nan03, Nan11
Nan11	<i>Campylaspis undata</i>	1	0.00	0.00	0.17	0.40	Nan05
Nan04	<i>Campylaspis globosa</i>	1	0.00	0.00	0.28	0.39	Nan05, Nan10
Nan10	<i>Campylaspis sulcata</i>	1	0.00	0.00	0.25	0.41	Nan07
Nan07	<i>Campylaspis rubicunda</i>	5	0.00	0.01	0.25	0.45	N06
	<i>Atlantocuma</i> sp.	1	0.00	0.00	0.31	0.41	B04, B05
Bod06	<i>Iphinoe serrata</i>	1	0.00	0.00	0.29	0.42	Bod05-A, -B
Bod05-A	<i>Cyclaspis longicaudata</i> (seq2)	1	0.00	0.00	0.08	0.43	Bod05-B
	<i>Cyclaspis</i> sp.	1	0.00	0.00	0.14	0.44	Bod05-B
Bod05-B	<i>Cyclaspis longicaudata</i> (ICE1-Bod001/ ICE1-Bod002)	2	0.00	0.00	0.08	0.45	Bod05-A
Bod03	Bodotriidae sp. 1	1	0.00	0.00	0.23	0.42	Bod04
Bod04	<i>Bathycuma brevirostre</i>	1	0.00	0.00	0.23	0.41	Bod03

N = Number of sequences