

TABLE S6 | Pairwise distances of the mitochondrial marker *mt-co1* between samples identified as *Hypanus guttatus*, with *H. berthallutzae* as an outgroup. Values in %.

%	<i>H. berthallutzae</i> GN18496	<i>H. aff. guttatus</i> GN13939	<i>H. aff. guttatus</i> GN13946	<i>H. guttatus</i> GN18434	<i>H. guttatus</i> GN18435	<i>H. guttatus</i> GN18436	<i>H. guttatus</i> GN18443
<i>H. aff. guttatus</i> GN13939	5,92						
<i>H. aff. guttatus</i> GN13946	5,53	0,38					
<i>H. guttatus</i> GN18434	6,11	1,72	1,34				
<i>H. guttatus</i> GN18435	5,92	1,53	1,15	0,19			
<i>H. guttatus</i> GN18436	5,92	1,53	1,15	0,19	0,00		
<i>H. guttatus</i> GN18443	5,92	1,53	1,15	0,19	0,00	0,00	
<i>H. guttatus</i> GN18451	5,92	1,53	1,15	0,19	0,00	0,00	0,00
<i>H. guttatus</i> GN18453	5,92	1,53	1,15	0,19	0,00	0,00	0,00
<i>H. guttatus</i> GN18454	5,92	1,53	1,15	0,19	0,00	0,00	0,00
<i>H. guttatus</i> GN18458	5,92	1,53	1,15	0,19	0,00	0,00	0,00
<i>H. guttatus</i> GN18459	5,92	1,53	1,15	0,19	0,00	0,00	0,00
<i>H. guttatus</i> GN18463	6,11	1,72	1,34	0,38	0,19	0,19	0,19
<i>H. guttatus</i> GN18466	5,92	1,53	1,15	0,19	0,00	0,00	0,00
<i>H. guttatus</i> GN18473	6,11	1,72	1,34	0,00	0,19	0,19	0,19
<i>H. guttatus</i> GN18474	6,11	1,72	1,34	0,00	0,19	0,19	0,19

→ **TABLE S6** | Continued in: <https://www.ni.bio.br/content/v22n1/1982-0224-2023-0046/supplementary/1982-0224-ni-22-01-e230046-s6.htm>



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