

Table S4 K2P distances of 18S rRNA sequences between genus *Halitemma* from Genbank and this study. Standard error estimates are shown above the diagonal with italics.

18S rRNA	1	2	3	4	5	6
1. <i>Halitemma rubrum</i> AY937323		<i>0.000</i>	<i>0.000</i>	<i>0.001</i>	<i>0.001</i>	<i>0.001</i>
2. <i>Halitemma rubrum</i> AY937325	0.000		<i>0.000</i>	<i>0.001</i>	<i>0.001</i>	<i>0.001</i>
3. <i>Halitemma rubrum</i> AY937358	0.000	0.000		<i>0.001</i>	<i>0.001</i>	<i>0.001</i>
4. <i>Halitemma isabu</i> sp. nov. Nectophore 1 11R MT935668	0.002	0.002	0.002		<i>0.000</i>	<i>0.000</i>
5. <i>Halitemma isabu</i> sp. nov. Bract 1 11R MT935667	0.002	0.002	0.002	0.000		<i>0.000</i>
6. <i>Halitemma isabu</i> sp. nov. Bract 2 Eddy1 MT935669	0.002	0.002	0.002	0.000	0.000	