Table S2 K2P distances of mtCOI sequences between genus *Halistemma* from Genbank and this study. Standard error estimates are shown above the diagonal with italics.

mtCOI	1	2	3	4	5	6	7	8	9
1. *Halistemma rubrum* | AY937372		*0.015*	*0.015*	*0.015*	*0.015*	*0.015*	*0.015*	*0.015*	*0.015*
2. *Halistemma* sp. 1 | GQ119989	0.122		*0.002*	*0.015*	*0.015*	*0.015*	*0.015*	*0.015*	*0.015*
3. *Halistemma* sp. 1 | GQ119991	0.122	0.003		*0.015*	*0.015*	*0.015*	*0.015*	*0.015*	*0.015*
4. *Halistemma isabu* **sp. nov.** | Nectophore 1 | 11R | MT933179	0.120	0.122	0.120		*0.000*	*0.000*	*0.000*	*0.000*	*0.002*
5. *Halistemma isabu* **sp. nov.** | Nectophore 2 | 11R | MT933180	0.120	0.122	0.120	0.000		*0.000*	*0.000*	*0.000*	*0.002*
6. *Halistemma isabu* **sp. nov.** | Nectophore 3 | 11R | MT933181	0.120	0.122	0.120	0.000	0.000		*0.000*	*0.000*	*0.002*
7. *Halistemma isabu* **sp. nov.** | Nectophore 4 | Eddy1 | MT933183	0.120	0.122	0.120	0.000	0.000	0.000		*0.000*	*0.002*
8. *Halistemma isabu* **sp. nov.** | Bract 1 | 11R | MT933178	0.120	0.122	0.120	0.000	0.000	0.000	0.000		*0.002*
9. *Halistemma isabu* **sp. nov.** | Bract 2 | Eddy1 | MT933182	0.118	0.124	0.122	0.002	0.002	0.002	0.002	0.002	