

S3 Table. Mean genetic divergences for the cytochrome oxidase *c* subunit 1 (*COI*) nucleotide sequences (Kimura-2-parameter [K2P] distances) of within-species among Cyclopoida.

Species	Average	S. E.
<i>Cyclops kikuchii</i>	-	-
<i>Diacyclops bicuspidatus</i>	-	-
<i>Macrocyclops albidus</i>	16.88	0.018
<i>Megacyclops viridis</i>	1.18	0.004
<i>Mesocyclops pehpeiensis</i>	0.52	0.003
<i>Mesocyclops dissimilis</i>	-	-
<i>Acnthocyclops vernalis</i>	-	-
<i>Apocyclops borneoensis</i>	-	-
<i>Halicyclops itohi</i>	-	-
<i>Paracyclops fimbriatus</i>	-	-
<i>Tropocyclops setulifer</i>	-	-
<i>Bonnierilla curvicaudata</i>	1.29	0.004
<i>Doropygus rigidus</i>	-	-
<i>Lonchidiopsis hartmeyeri</i>	-	-
<i>Pachypygus curvatus</i>	-	-
<i>Oithona similis</i>	1.41	0.005
<i>Oithona davisae</i>	-	-