

**S1 Table. Mean genetic divergences (= Kimura-2-parameter distances) for the cytochrome oxidase *C* subunit 1 (*COI*) nucleotide sequences within species in the order Calanoida.**

<b>Species</b>	<b>Average</b>	<b>S. E.</b>
<i>Acartia erythrea</i>	0.00	0.000
<i>Acartia steueri</i>	0.00	0.002
<i>Acartia tsuensis</i>	0.00	0.002
<i>Sinocalanus tenellus</i>	0.01	0.003
<i>Heliodyptomus kikuchii</i>	-	-
<i>Neodyptomus schmackeri</i>	0.01	0.004
<i>Sinodyptomus sarsi</i>	-	-
<i>Acanthodyptomus pacificus</i>	0.03	0.007
<i>Calanus sinicus</i>	0.00	0.002
<i>Paracalanus parvus</i>	0.17	0.017
<i>Pseudodyptomus inopinus</i>	0.01	0.004
<i>Pseudodyptomus marinus</i>	0.01	0.004
<i>Pseudodyptomus nihonkaiensis</i>	0.01	0.003
<i>Eurytemora affinis</i>	-	-
<i>Eurytemora pacifica</i>	0.03	0.006
<i>Temora turbinata</i>	0.00	0.001