







































**Table 1. Kimura 2 parameter (K2P) intraspecific genetic distance values based on COI sequences and standard error values (d-intra-genetic distance, e-standard error).**

	d	e
<i>Brachionus plicatilis</i>	0.041	0.012
<i>Brachionus angularis</i>	0.066	0.016
<i>Brachionus calyciflorus</i>	0.044	0.015
<i>Keratella tropica</i>	0.044	0.015
<i>Asplanchna brightwelli</i>	0.012	0.006