Figure S1 Phylogenetic tree of Tanaidacea sequenced in this work constructed with Bayesian method with 32 BUSCO genes. The tree was constructed in Phylobayes under the GTR+G model. Posterior node support values are reported at each node.

Table S1 Collection information for specimens used in this study. KK, Keiichi Kakui

Supplementary Text Analysis of opsin in tanaidaceans.