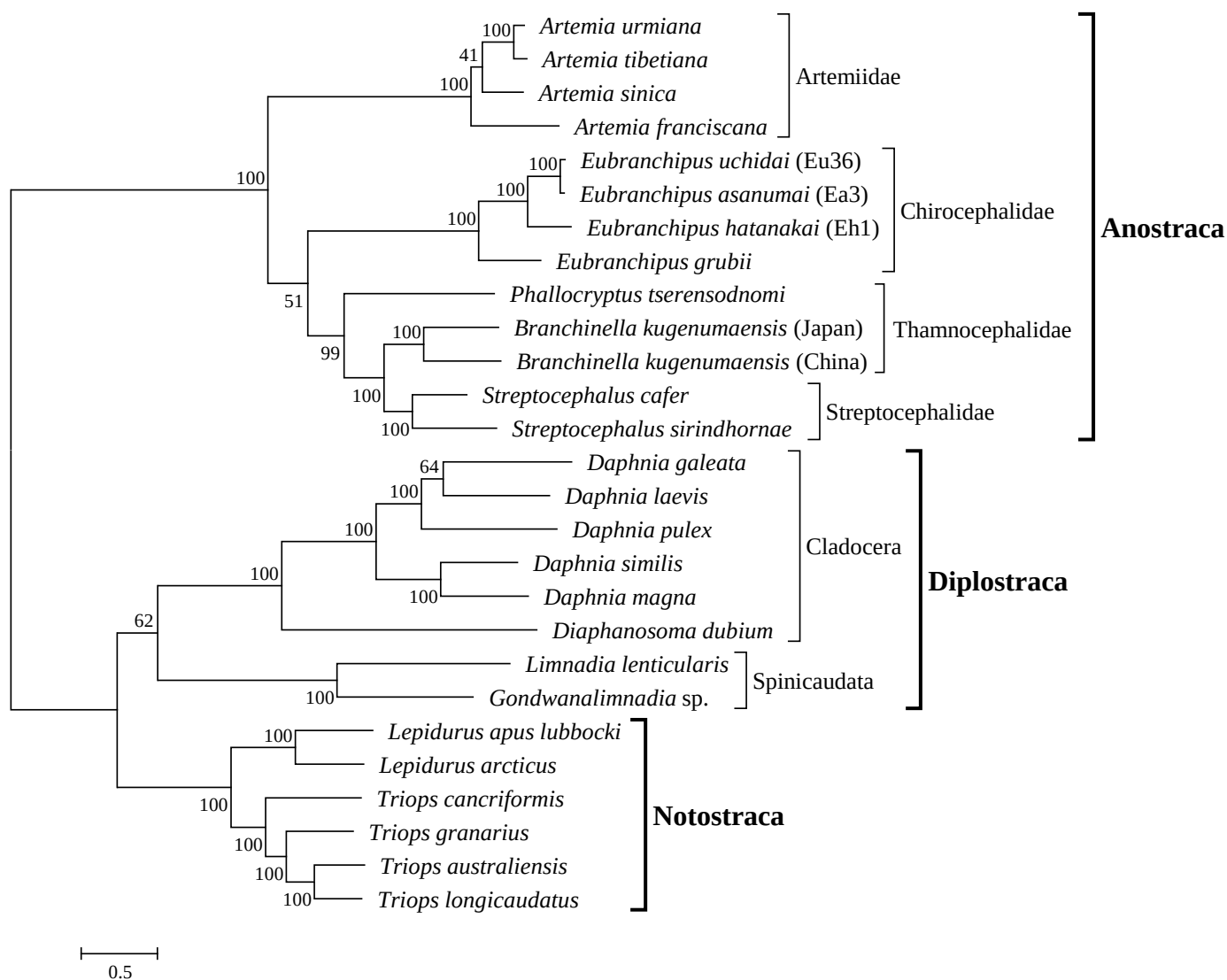




Supplementary Fig. S3 The maximum likelihood tree of Branchiopoda species based on the concatenated 13 protein-coding genes in the mitochondrial genome. A scale bar representing the number of nucleotide substitutions per site is shown. Values near the nodes represent the percent bootstrap probability.