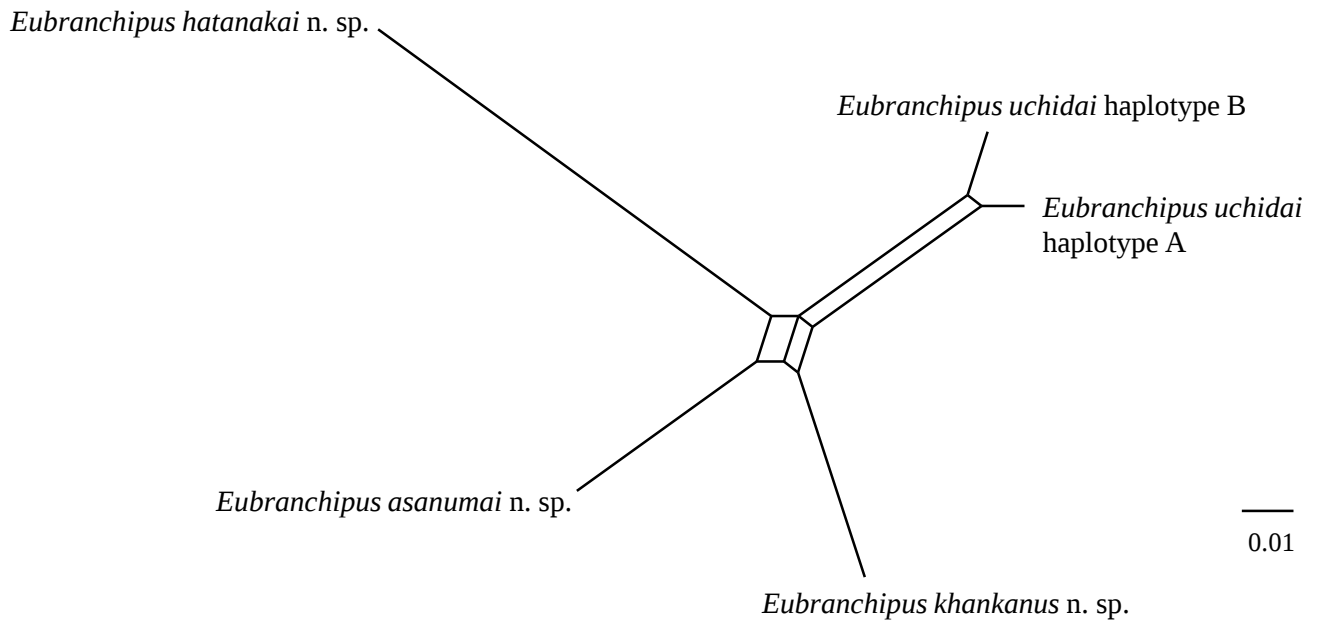


A



B

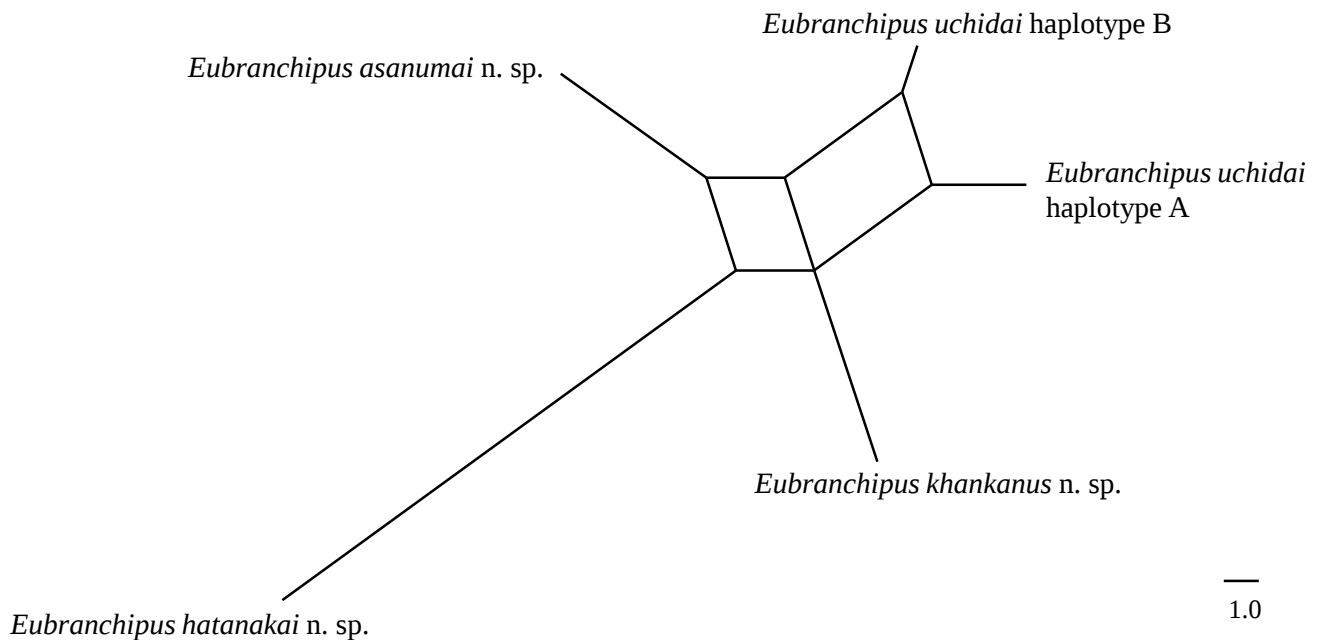


Fig. S1. Phylogenetic networks of Far East *Eubbranchipus* species based on COI (A) and 16S (B) data. Networks were reconstructed using the Neighbor-Net method (Bryant and Moulton 2004) with p-distance implemented in the SplitsTree4 software (Huson and Bryant 2006).