



Fig. 1. Genetic divergence of the common Japanese intertidal goby *Chaenogobius annularis* inferred from mitochondrial DNA data. In total, 169 unique haplotypes were obtained, and phylogenetic trees showed two genetically distinct lineages: the Pacific Ocean and the Sea of Japan lineages, whose divergence was estimated to have occurred in the early Pleistocene, related to the paleoenvironmental history of the Sea of Japan (Hirase *et al.*, 2012).