Additional file 2: Figure S1: Neighbor-joining tree of mtDNA haplotypes from CR data.
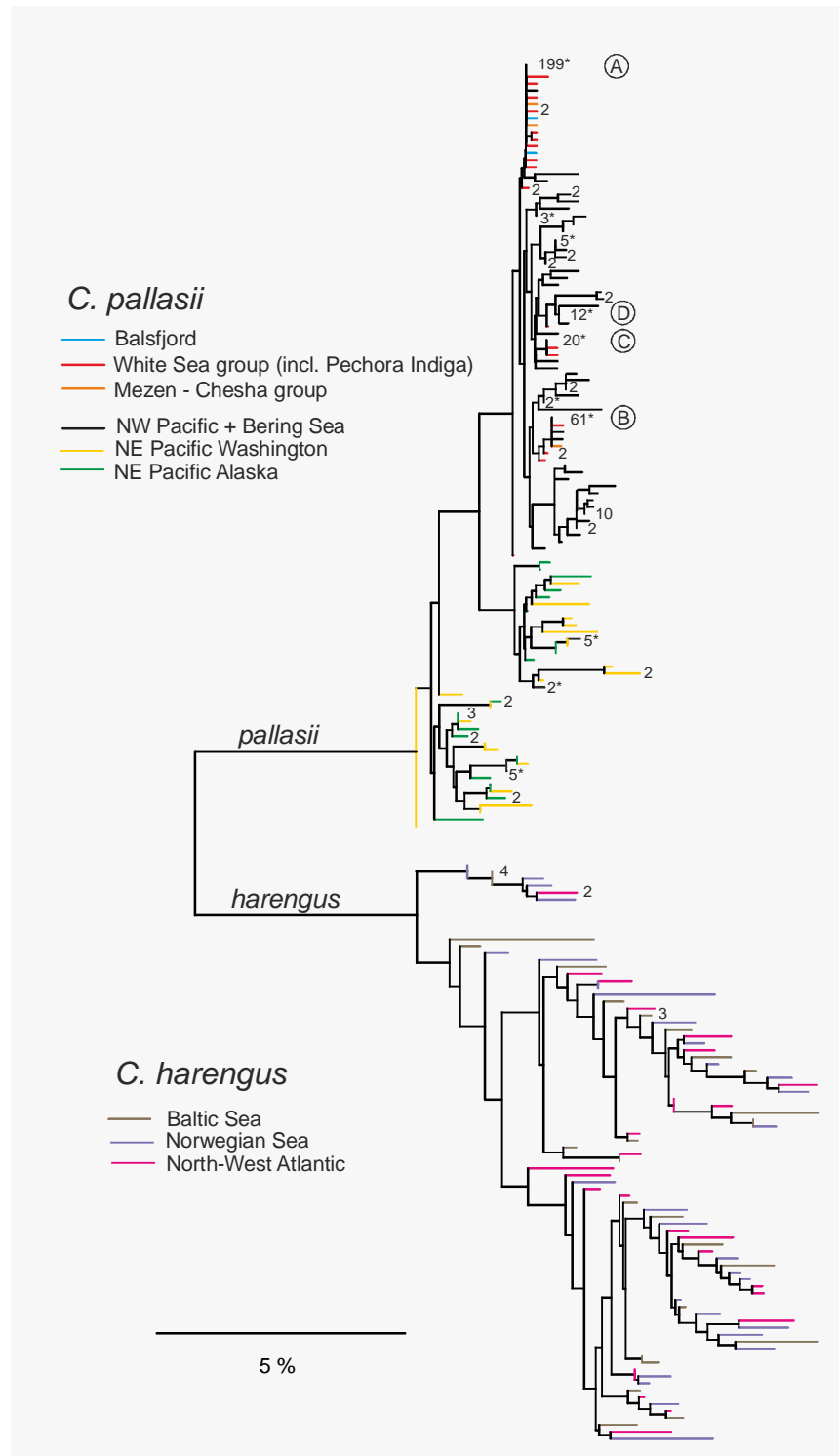

Figure S1: Neighbor-joining tree of mtDNA haplotypes from CR data. The tree is based on GTR+I+Γ model distances, and is comparable to the figures for *cyt-b* and concatenated data presented in Figure 2 of the paper itself. The numbers are observed frequencies of haplotypes found in multiple individuals; haplotypes shared between localities are indicated with asterisks. Color codes refer to population groups as in Figure 1. The core haplotypes A-D from Figure 3 are indicated.