

Figure S4

Comparison of admixture time among Japanese clusters. The admixture time was estimated using the 1-reference ALDER test (Table 1). The squares and whiskers represent the estimate and SE of the admixture time for single clusters. A diamond represents the meta-analysis of a group of clusters; the center and width of the diamond indicate the combined estimate and SE. The decimal beside the diamond is the P -value for heterogeneity of admixture times among the clusters. There was no heterogeneity among the clusters in Hondo or among the clusters in Ryukyu ($P_{\text{het}} > 0.05$). However, when all Japanese clusters were combined, there was significant heterogeneity in admixture time ($P_{\text{het}} < 0.05$ when the reference population is C284 or C291).

