



Figure S1. Enrichment of genic and non-genic SNPs for values of $|\delta|$ between pairs of populations. Enrichments of genic and non-genic SNPs are calculated within each $|\delta|$ value bin (width 0.1). (A) Enrichments calculated separately for each pair of populations among the 11 populations (55 pairs in total). Each line indicates a unique population pair. (B) Enrichments of genic and non-genic SNPs averaged over population pairs -- i.e., an average of the lines in panel (A), for genic and non-genic enrichments separately. (C) Summary of panel (A). Black line indicates the number of population pairs with a defined value of $|\delta|$ for each bin (i.e., the number of population pairs where there exists at least one SNP in the given $|\delta|$ bin). Red line indicates the number of population pairs where there appears to be an enrichment of genic SNPs (enrichment value > 1) in the given $|\delta|$ bin.