



Figure S3. Enrichment of genic and non-genic SNPs for values of δ between the HGDP Yoruba and HapMap Tuscans. Enrichments of genic and nongenic SNPs are calculated within each δ value bin, with bootstrap confidence intervals, as in Figures S1 and S2 and main text. SNPs where the derived allele is fixed in the Yorubans and absent in the Tuscans are depicted on the right hand side of the plot (near 1); SNPs where the derived allele is fixed in the Tuscans and absent in the Yorubans are depicted on the left hand side of the plot (near -1).