



Figure S8. P-values for the 5 most extreme 100 kb genomic windows according to the XP-EHH KHB statistic in each population. Red indicates $p < 0.01$, orange indicates $p < 0.05$, yellow indicates $p < 0.10$, and white indicates $p > 0.10$. "Genes" column lists the genes located within the indicated windows.