Figure S7. P-values for the 5 most extreme 100 kb genomic windows according to the XP-EHH YRI 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.