



Figure 1-figure supplement 7: Diversity of populations along chromosomes measured based on 10 kb non-overlapping windows. Nucleotide diversity (π) distribution of 10 kb windows in population Highland (a) and 7 Lowland (b). (c) Nucleotide diversity ratios ($\pi_{\text{Lowland}} / \pi_{\text{Highland}}$). (d) Pairwise genome-wide fixation index (F_{ST}) between Highland and Lowland. The broken horizontal line represents the top 1% threshold.