Figure S10: For each continent, the difference in heterozygosity between high and low-latitude populations. Red line represents the 99.5 % confidence interval for the difference in \( \hat{\pi} \). Yellow dots indicate windows outside in the 99.5% confidence interval that also contain an outlier SNP in the 1% tail. For North America, the distribution of yellow dots has a p-value of $1.2e^{-7}$, under the expectation that they are binomially distributed with frequency equal to the proportion of windows in one extreme over the other. In Australia, the p-value is 0.0019.