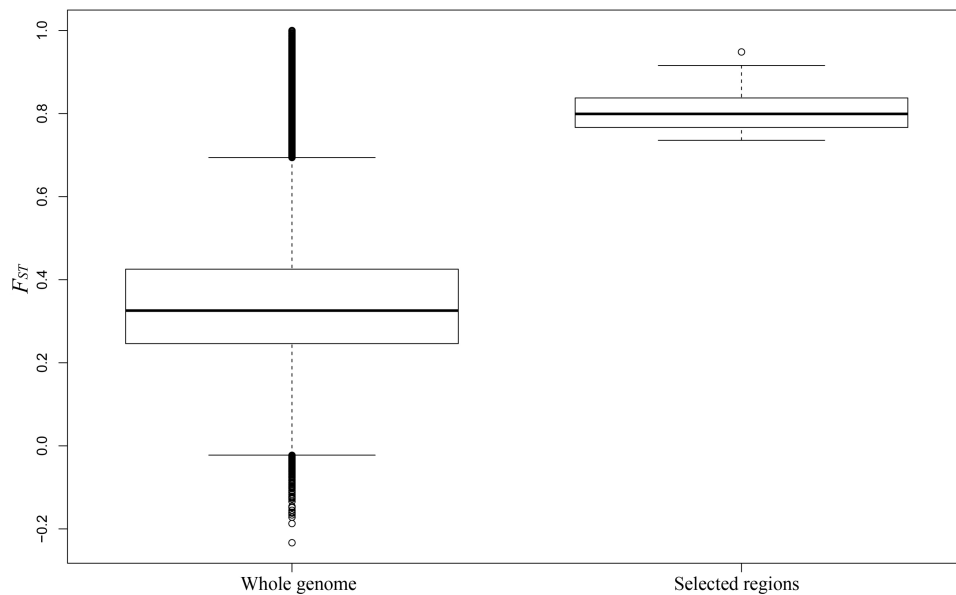
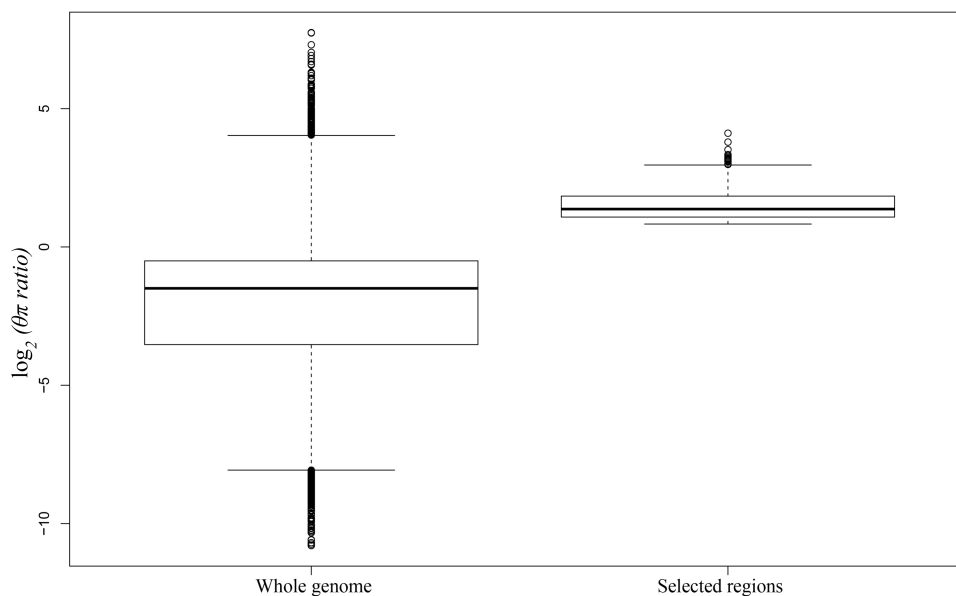


A**B**

Additional file 15: Fig S1. Differences in $Z(F_{ST})$ and $\theta\pi$ ratio values between the selected regions and the whole-genome scale for Tarim red deer (TRD). (A) Comparison between $Z(F_{ST})$ values for genomic regions that have undergone selective sweeps and $Z(F_{ST})$ values at the whole-genome scale in TRD. The upper, middle (within the box), and lower boundary lines for the boxes represent 25%, 50% (median value), and 75% of the $Z(F_{ST})$ and $\theta\pi$ ratio values, respectively. (B) Comparison between $\theta\pi$ ratio values for genomic regions that have undergone selective sweeps and $\theta\pi$ ratio values at the whole-genome scale.