Figure S8

Chromosome: 2L, putative site: 4824431, detected by $iHS$ (-4.03)

Chromosome: 2L, putative site: 11036917, detected by $iHS$ (-3.82)

Chromosome: 2L, putative site: 1147263, detected by $nS_L$ (-3.77)

Chromosome: 2L, putative site: 12375980, detected by $nS_L$ (-3.46)

Chromosome: 2R, putative site: 5708056, detected by $iHS$ (-3.33)
Chromosome: 2R, putative site: 10665005, detected by \( iHS \) (-3.86)

Chromosome: 2R, putative site: 18089697, detected by \( nS_L \) (-2.80)

Chromosome: 2R, putative site: 5548485, detected by \( nS_L \) (-2.42)
Chromosome: 3L, putative site: 5966477, detected by *iHS* (-4.18)

Chromosome: 3L, putative site: 14431036, detected by *iHS* (-4.12)

Chromosome: 3L, putative site: 11654544, detected by *nS*$_L$ (-3.59)

Chromosome: 3L, putative site: 2998073, detected by *nS*$_L$ (-3.82)
Chromosome: 3R, putative site: 18973529, detected by $iHS (-3.49)$

Chromosome: 3R, putative site: 8567660, detected by $iHS (-3.70)$

Chromosome: 3R, putative site: 20876949, detected by $nS_L (-3.49)$

Chromosome: 3R, putative site: 19839035, detected by $nS_L (-3.17)$
Chromosome: X, putative site: 18636156, detected by $iHS$ (-5.23)

Chromosome: X, putative site: 19093150, detected by $iHS$ (-5.80)

Chromosome: X, putative site: 17124249, detected by $nSL$ (-4.25)

Chromosome: X, putative site: 14975977, detected by $nSL$ (-4.16)

**Figure S8 legend:** Polymorphism patterns of genome areas surrounding the putative site under selection detected exclusively by $iHS$ or $nSL$ method. For each chromosome arm, top two candidate loci with strongest signals by each method, however not significant by the other tests, are shown.