SUPPLEMENTAL FIGURE 3
Haploblock partitioning across each genomic region. The SNPs in the locus are proportionally spaced in a bar across the top and named according to ID number for supplemental Table 2 for each genomic region. Each population is given a line with the name on the left (Chinese breeds are named with prefix C and European breeds are named with prefix E. If inside a haploblock a SNP is below the MAF (minor allele frequency) threshold or fails the Hardy-Weinberg test, the corresponding part of the segment is not colored.

Target 1