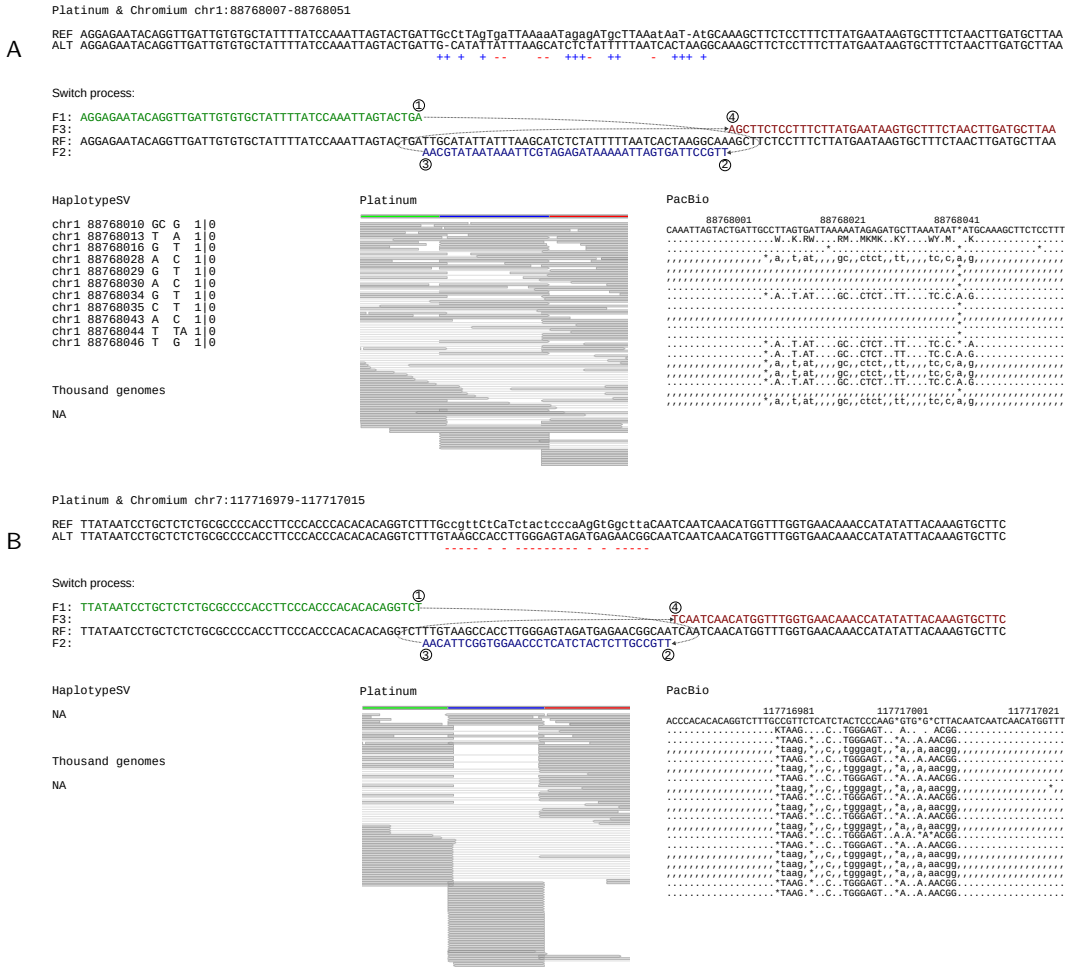

Fig S10. TSM candidate loci missing from HaplotypeSV data. (**A**) A partially called locus. Called and missing changes are indicated with plus (blue) and minus (red) signs. (**B**) A locus completely missing from HaplotypeSV data. (**C**) A locus that is missing despite being in full in the 1000 Genomes (1kG) data. The plots show the two alleles and their TSM solution (top); HaplotypeSV and 1kG variants (left); NA12878 Platinum (middle) and PacBio (right) data for the given locus. 1kG allele frequencies are shown for all and European samples. Alignment gaps cannot be visualised with the Gviz package and in (C) the Platinum data are shown in text format.