



Supplementary Figure 3. Haplotype structure $\pm 15\text{Kb}$ around rs368234815 for three population per continent.

Columns represent SNPs with a derived allele frequency $> 5\%$ in at least one population ($n=99$ SNPs), with the ancestral allele in white, and the derived allele in black. Horizontal lines represent the haplotypes they fall in, as inferred with SHAPEIT by the 1000 Genomes consortium [Mcvean 2012]. Haplotypes were sorted based on rs368234815 (red arrow) and SNPs in perfect LD with it in CHS (black arrows); see also Table 2 and Figure 4. The bar on the left-hand side of each plot indicates haplotypes carrying the TT allele (red) or the ΔG allele (blue).