

HapMap Populations

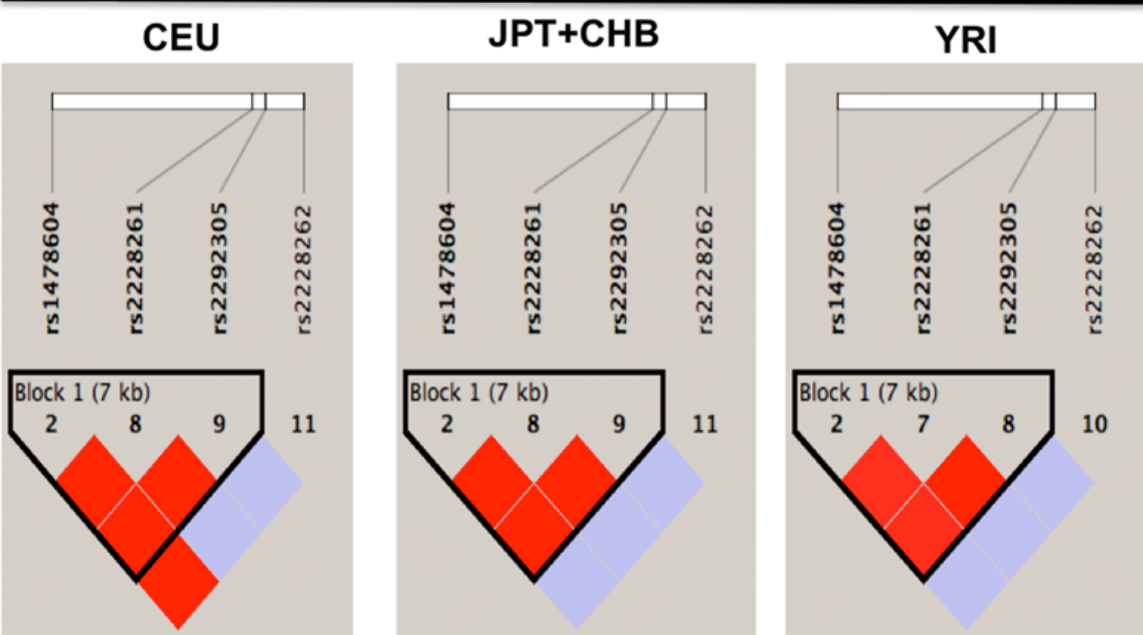


FIGURE 4. Linkage disequilibrium (LD) pattern of four analyzed single nucleotide polymorphisms (SNPs) in the study population resembled those detected in HapMap populations. The figure shows the D' -based pairwise LD map of 4 SNPs genotyped in HapMap populations. Darker shades of red indicate higher LD coefficient (D'). The same haplotype block was outlined in the three populations.