



Supplementary Figure 1. Linkage disequilibrium plots of CD1A SNPs in HapMap populations. CD1A consists of six exons (■) and two untranslated regions (▒) on chromosome 1. Shown are minor allele frequencies (dotted box) and R² coefficients (shaded box) for SNPs spanning CD1A ± 10kB flanking regions in CHB/JPT HapMap populations.