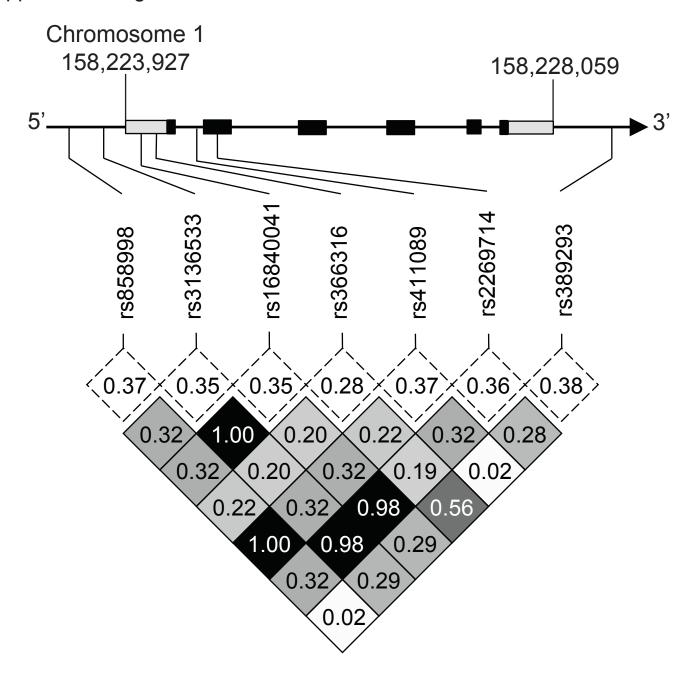
Seshadri et al. Supplemental Figure 1



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 \mp 10kB flanking regions in CHB/JPT HapMap populations.