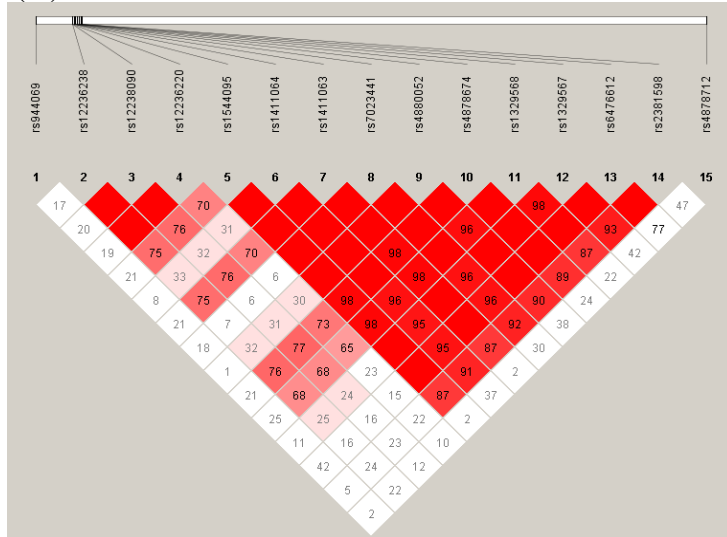


**Figure S8.** Linkage disequilibrium patterns in the 1000 Genomes AFR reference panel for the GWAS-implicated region spanning from *PAX5* to *FRMPD1* on chromosome 9. The patterns, according to (A)  $D'$  and (B)  $r^2$  values, are shown for the 15 GWAS-implicated SNPs tested for replication on chromosome 9. Darker shading from pink to red indicates higher  $D'$  values, and darker shading from grey to black indicates higher  $r^2$  values.

(A)



(B)

