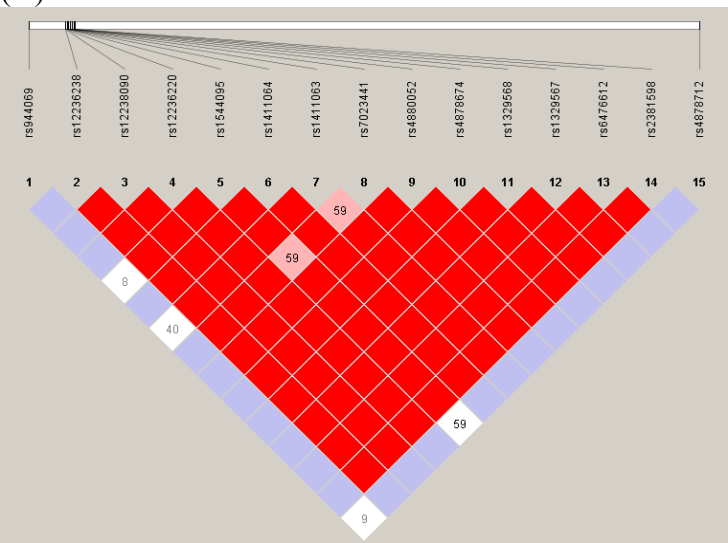
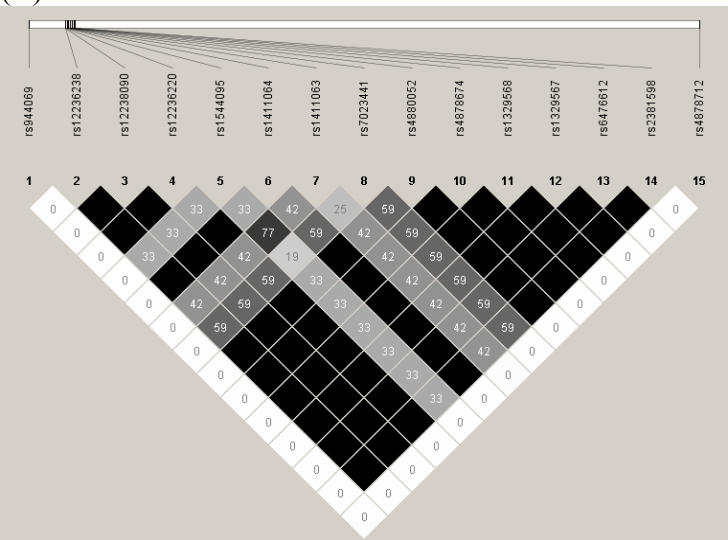
Figure S9. Linkage disequilibrium patterns in the 1000 Genomes EUR reference panel for the GWAS-implicated region spanning from *PAX5* to *FRMPD1* on chromosome 9. The patterns, according to (A) D' and (B) r^2 values, are shown for the 15 GWAS-implicated SNPs tested for replication on chromosome 9. Darker shading from pink to red indicates higher D' values, and darker shading from grey to black indicates higher r^2 values. Purple coloring indicates high D' values but lower levels of statistical significance.

(A)



(B)