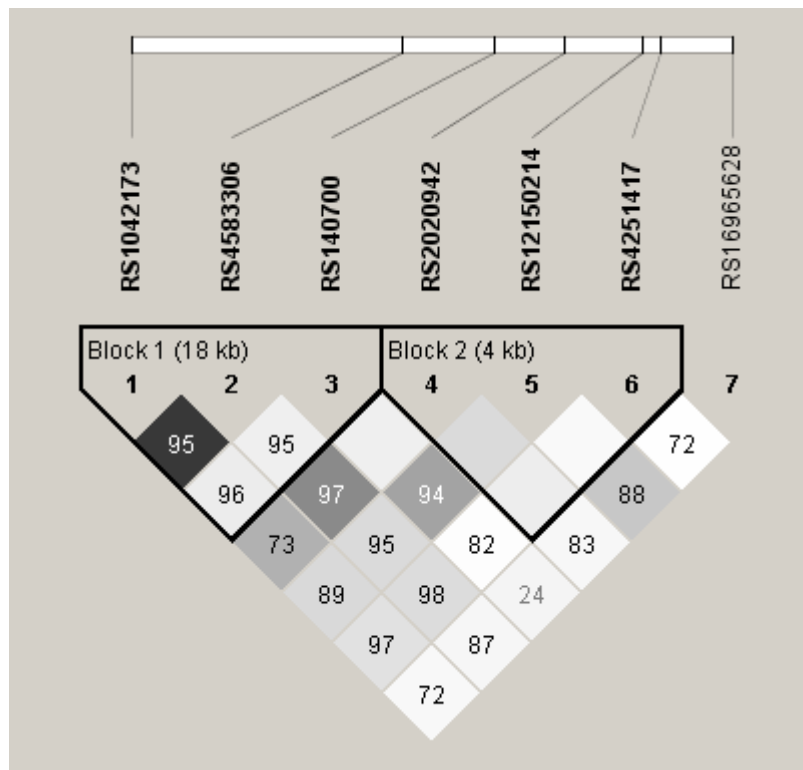


Figure S1. Linkage disequilibrium (LD) structure in the three Scandinavian control samples of seven serotonin transporter gene (*SLC6A4*) single nucleotide polymorphisms.



The figure shows pair-wise LD among the seven *SLC6A4* SNPs, as calculated by the Haploview 4.0 software. High D' values with strong linkage disequilibrium are indicated by dark color. Gray and white colors indicate weak linkages with low D' values. The genetic positions of the SNPs in the *SLC6A4* gene are indicated in the white bar above the LD-plot.