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.