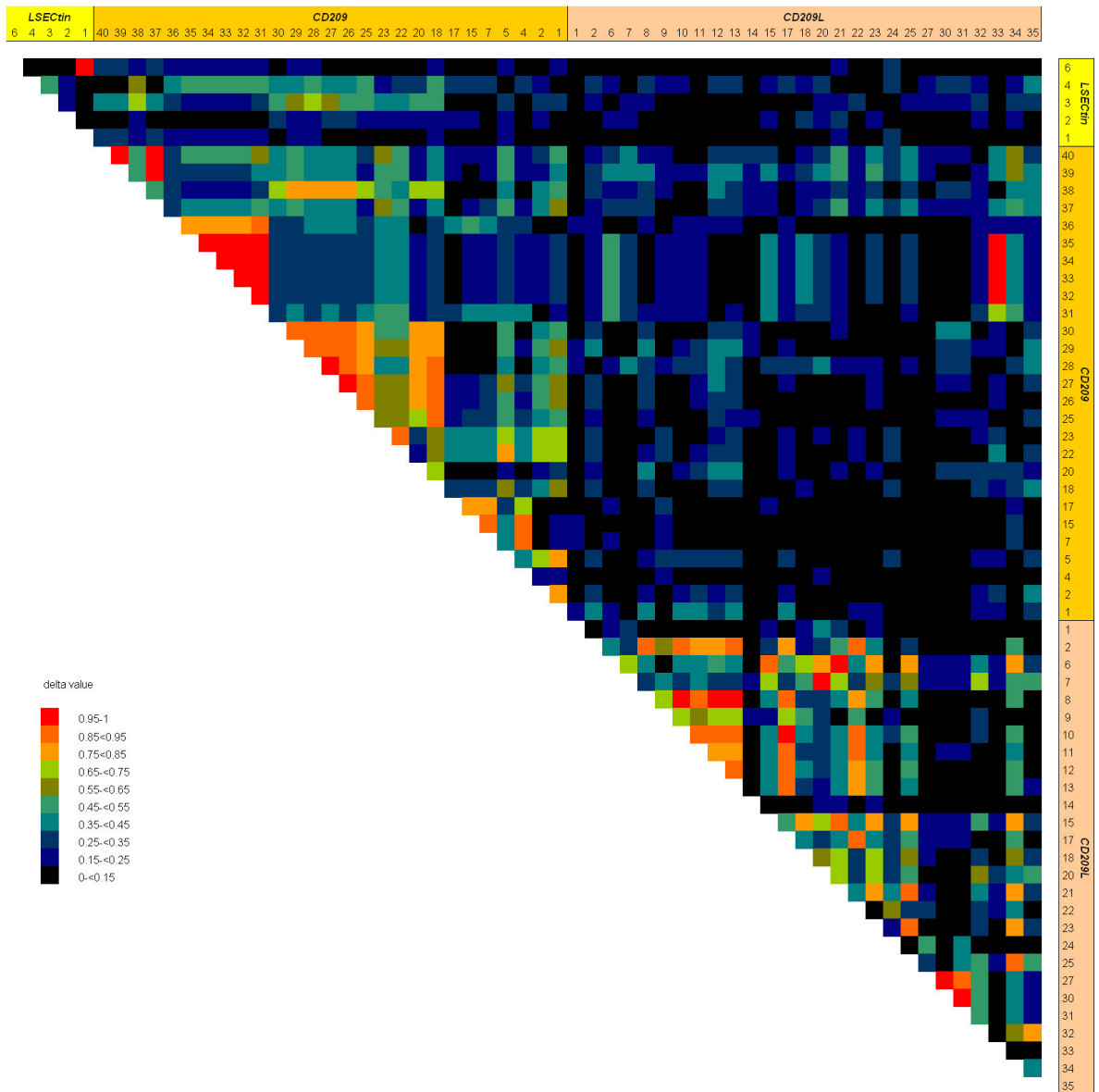


Supplementary Figure 1. Linkage disequilibrium between SNPs in *CD209* and flanking genes *CLEC4M* (*CD209L*) and *CLEC4G* (*LSECtin*)



Notes:

Computation of Δ -values was restricted to SNPs with rare allele frequency ≥ 0.02 . Polymorphism DCSIGN2-ex4RPT (code 9) is multiallelic, and the Δ -value was computed using the most frequent allele and all other alleles grouped. None of the other alleles showed significant LD with SNPs located in *CD209*.