Supplementary figure 2. Linkage disequilibrium (LD) in *SREBF2*. Genotypes from the German sample controls were uploaded and analysed in Haploview. $R^2$ values for 2 markers comparison are displayed as numbers in losanges. The color scale displays the extent of LD with the GOLD standards (Abecasis and Cookson, 2000, GOLD--graphical overview of linkage disequilibrium, Bioinformatics, 16(2): 182-3.) as implemented in Haploview; with a gradient of color from red for strong LD to blue for no LD.