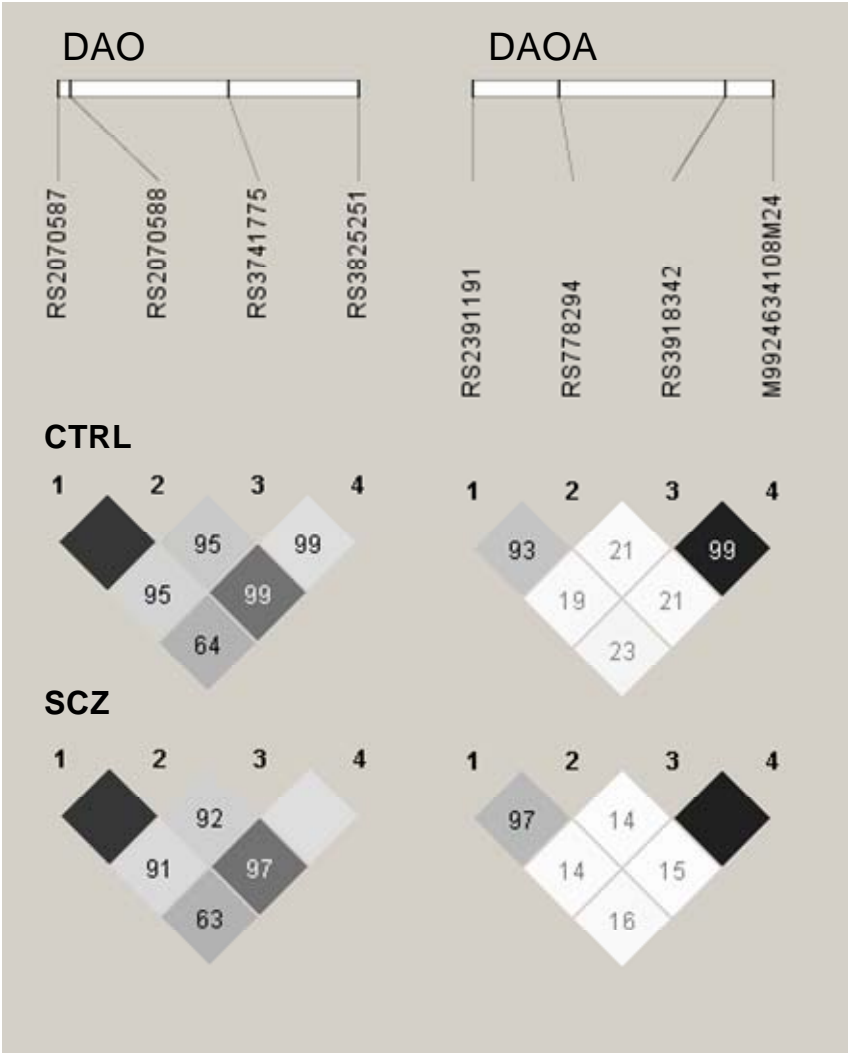
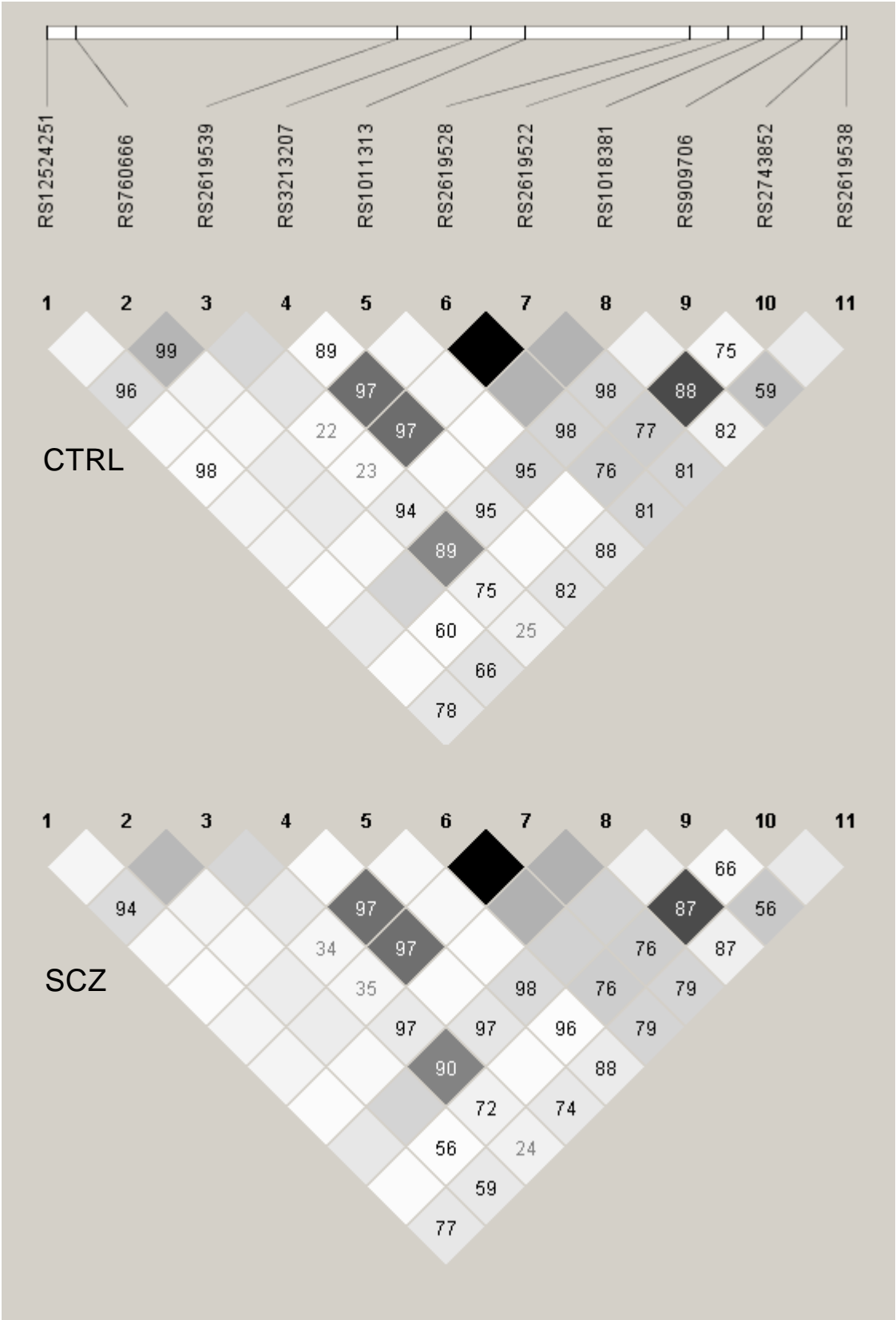


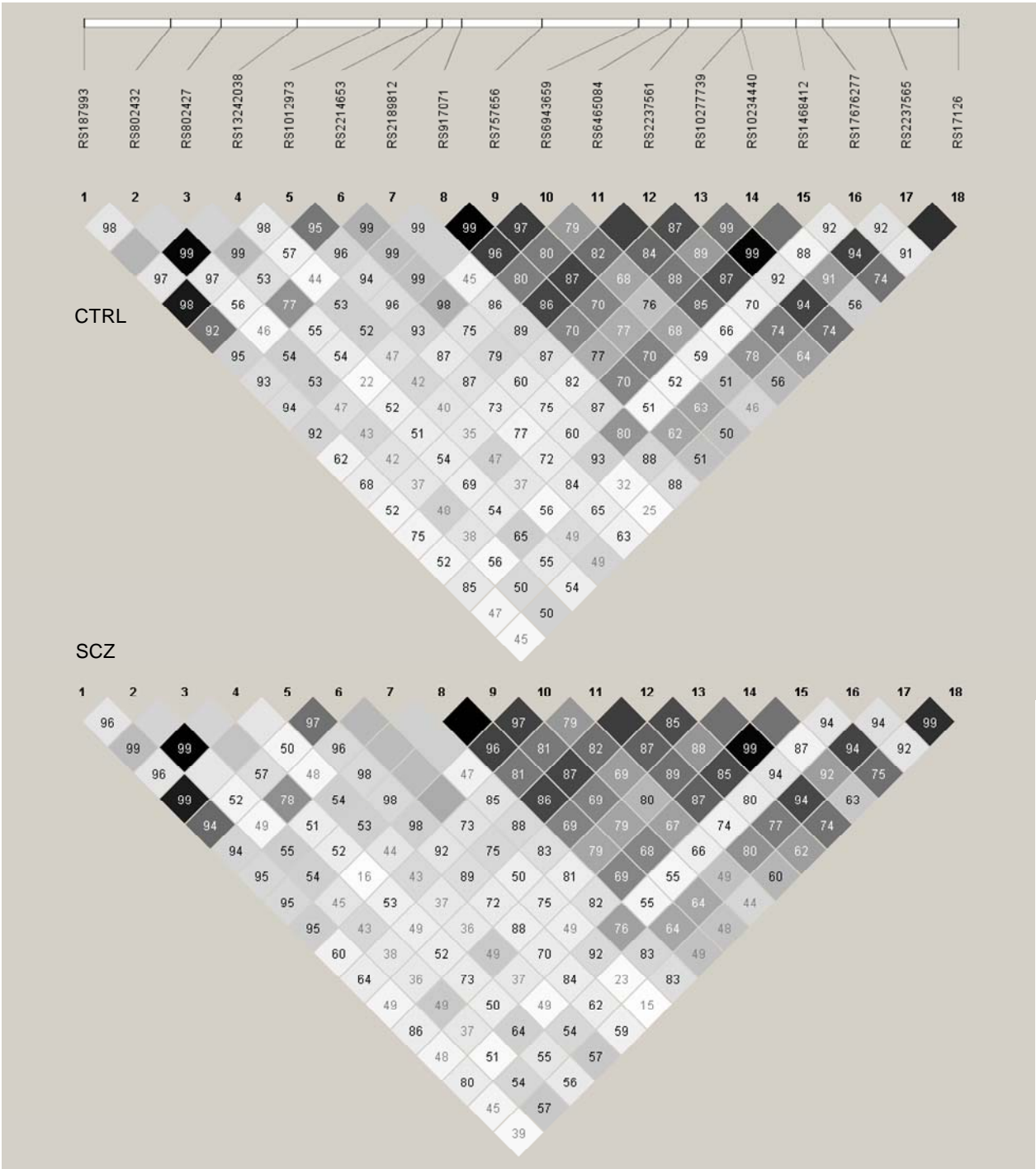
Supplementary figure 1a.



Supplementary figure 1b.



Supplementary figure 1d.



Supplementary **Figure 1a**. Linkage disequilibrium (LD) structure of SNPs genotyped in the D-amino-acid oxidase (*DAO*) and the D-amino-acid oxidase activator (*DAOA*) genes. LD structure was similar for unaffected (CTRL) and affected (SCZ) individuals. D' (numbers) and R^2 (shadings) are given. Squares lacking numbers indicate $D' = 1$.

Supplementary **Figure 1b**. Linkage disequilibrium (LD) structure of SNPs genotyped in dystrobrevin-binding protein 1 gene (*DTNBP1*). LD structure was similar for unaffected (CTRL) and affected (SCZ) individuals. D' (numbers) and R^2 (shadings) are given. Squares lacking numbers indicate $D' = 1$.

Supplementary **Figure 1c**. Linkage disequilibrium (LD) structure of SNPs genotyped in the neuregulin 1 gene (*NRG1*). LD structure was similar for unaffected (CTRL) and affected (SCZ) individuals. D' (numbers) and R^2 (shadings) are given. Squares lacking numbers indicate $D' = 1$.

Supplementary **figure 1d**. Linkage disequilibrium (LD) structure of SNPs genotyped in metabotropic glutamate receptor 3 gene (*GRM3*). LD structure was similar for unaffected (CTRL) and affected (SCZ) individuals. D' (numbers) and R^2 (shadings) are given. Squares lacking numbers indicate $D' = 1$.