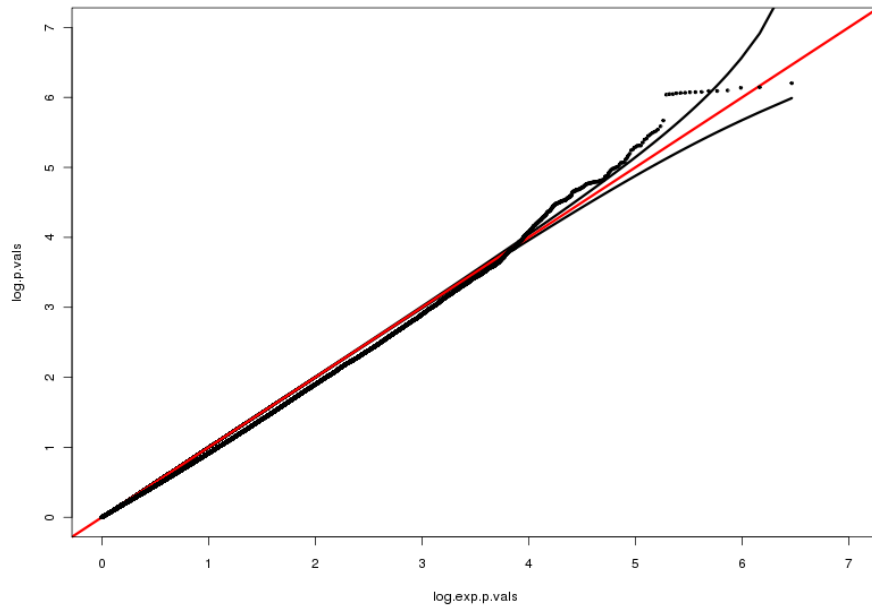


ESM Fig. 3. Q-Q plot for combined genome-wide association results in a total of 684 T2D patients and 955 controls based on the 2,925,090 analyzed SNPs. The black lines represent the upper and lower boundaries of the 95% confidence bands.

Original



Genomic control

