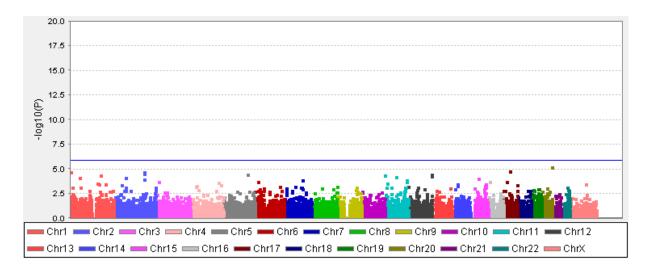
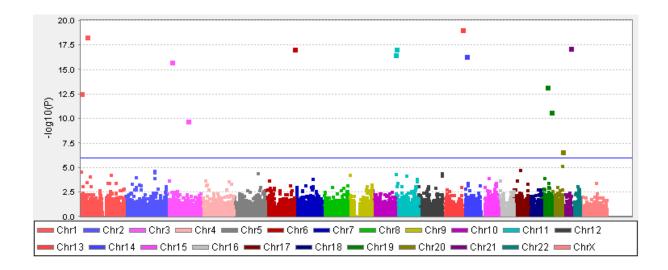
## S2 Figure. Manhattan plots.



**Figure A.** Manhattan plot of the Fisher's exact test association p-values for the single variant analyses with MAF  $\geq$ 1.0%. Chromosomes are shown on the x-axis. On the y-axis, the  $-\log 10$  of the p-value is shown. The blue line represents the threshold for significance after Bonferroni correction for 36,032 variants (1.39\*10<sup>-6</sup>, which corresponds to a value of 5.86). Variants above this line are considered statistically significant.



**Figure B.** Manhattan plot of the Fisher's exact test association p-values for the single variant analyses with MAF ≥0.4%. Chromosomes are shown on the x-axis. On the y-axis, the  $-\log 10$  of the p-value is shown. The blue line represents the threshold for significance after Bonferroni correction for 43,653 variants (1.15\*10<sup>-6</sup>, which corresponds to a value of 5.94). Variants above this line are considered statistically significant.