Case-control association test for CSD chr1 chr2 chr3 chr4 chr5 chr6 10.0 -7.5 --log10 p-value 5.0 -2.5 -

0.0e+600e+05e+07e+00 5e+06 1e+070.0e+600e+060e+05e+000e+060e+07.5e+0725006000076000000e+001e+062e+06e+06

Genomic position