Case-control association test for CSD chr1 chr2 chr3 chr4 chr5 chr6 10.0 -7.5 -5.0 -2.5 -10.0 --log10 p-value 7.5 -5.0 -2.5 -0.0 -10.0 -7.5 -5.0 -2.5 -0.0e + 0.00e + 0.00e

Genomic position