Case-control associaiton test for CSD chr1 chr2 chr3 chr4 chr5 chr6 6 --log2 p-value 2 -1e+07 0.0e+06.0e+06.0e+07.5e+07.0e+005.0e+061.0e+071.5e+070 250006000007600000 0e+001e+062e+063e+06 0.0e+0**5**.0e+0**6**.0e+0**7**.5e+0**7**0e+00

Genomic position