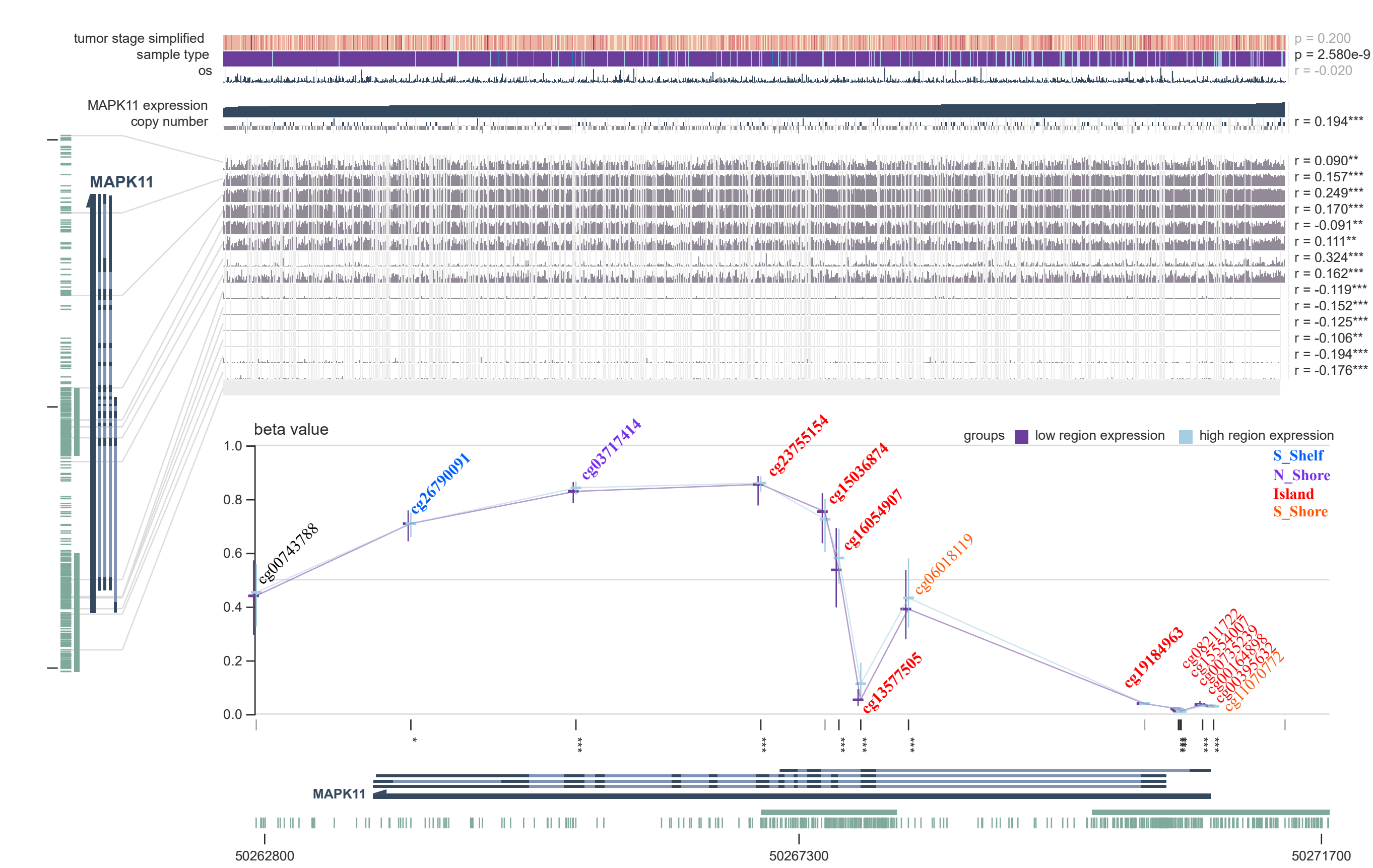
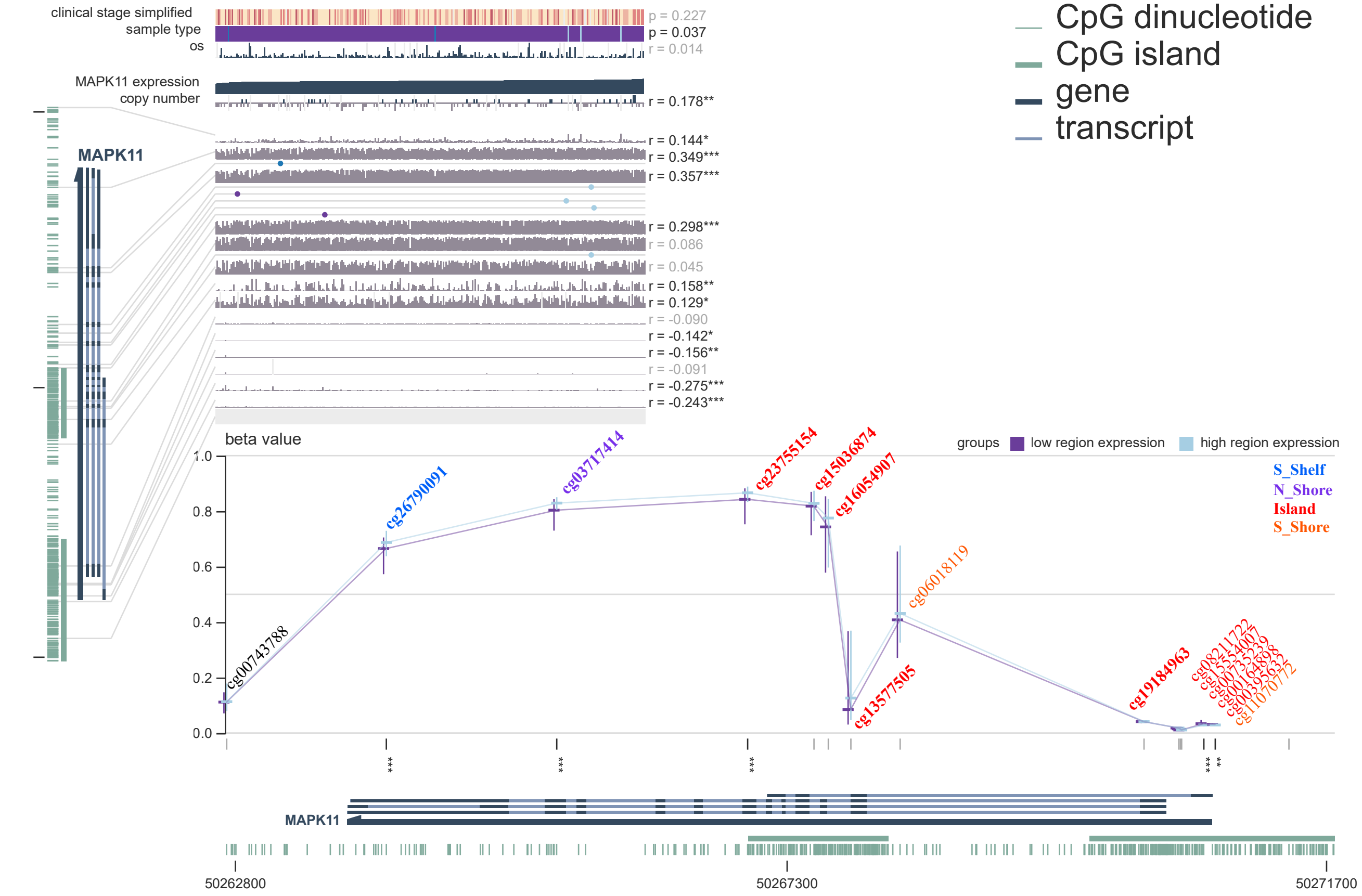


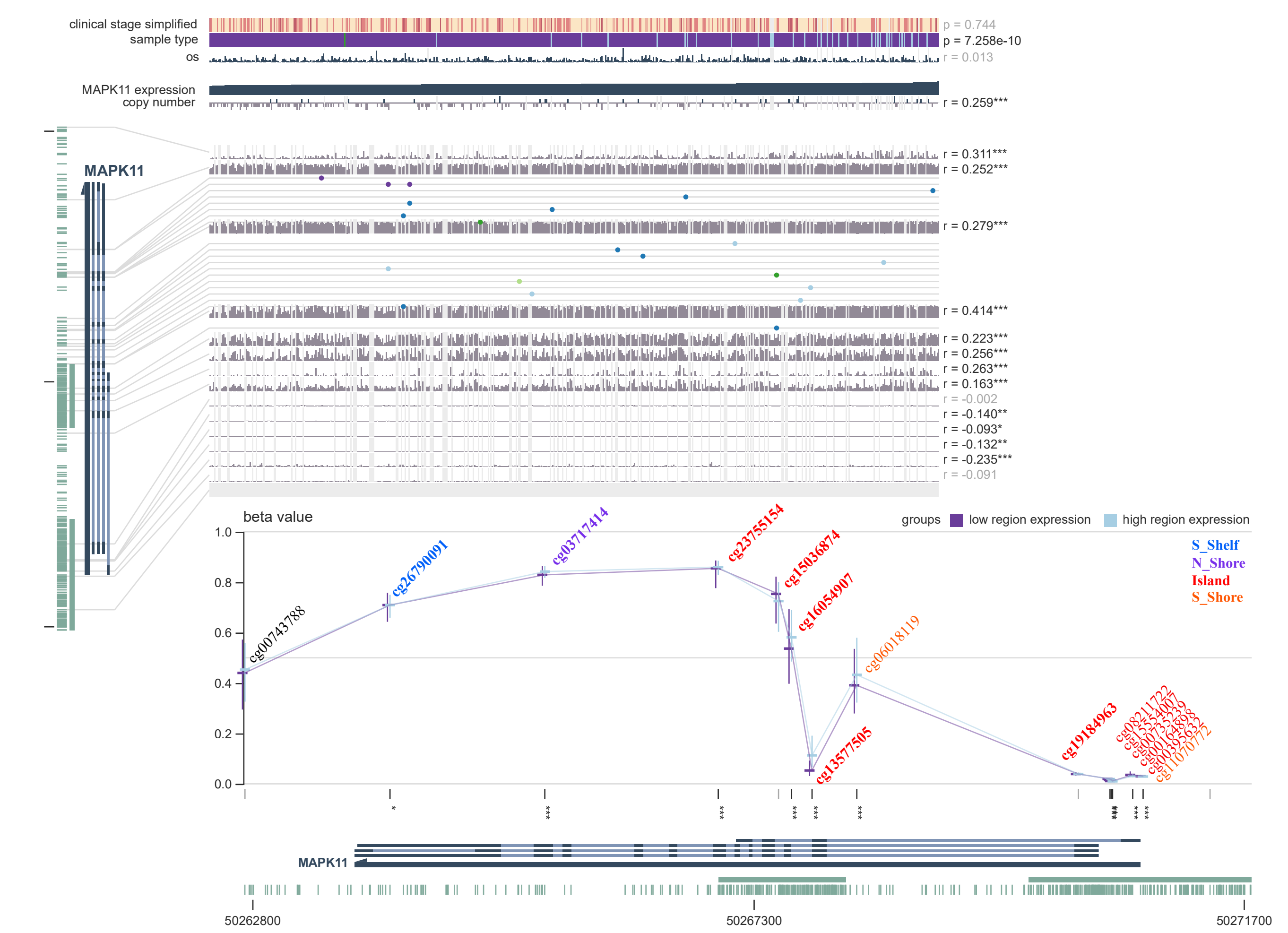
BRCA



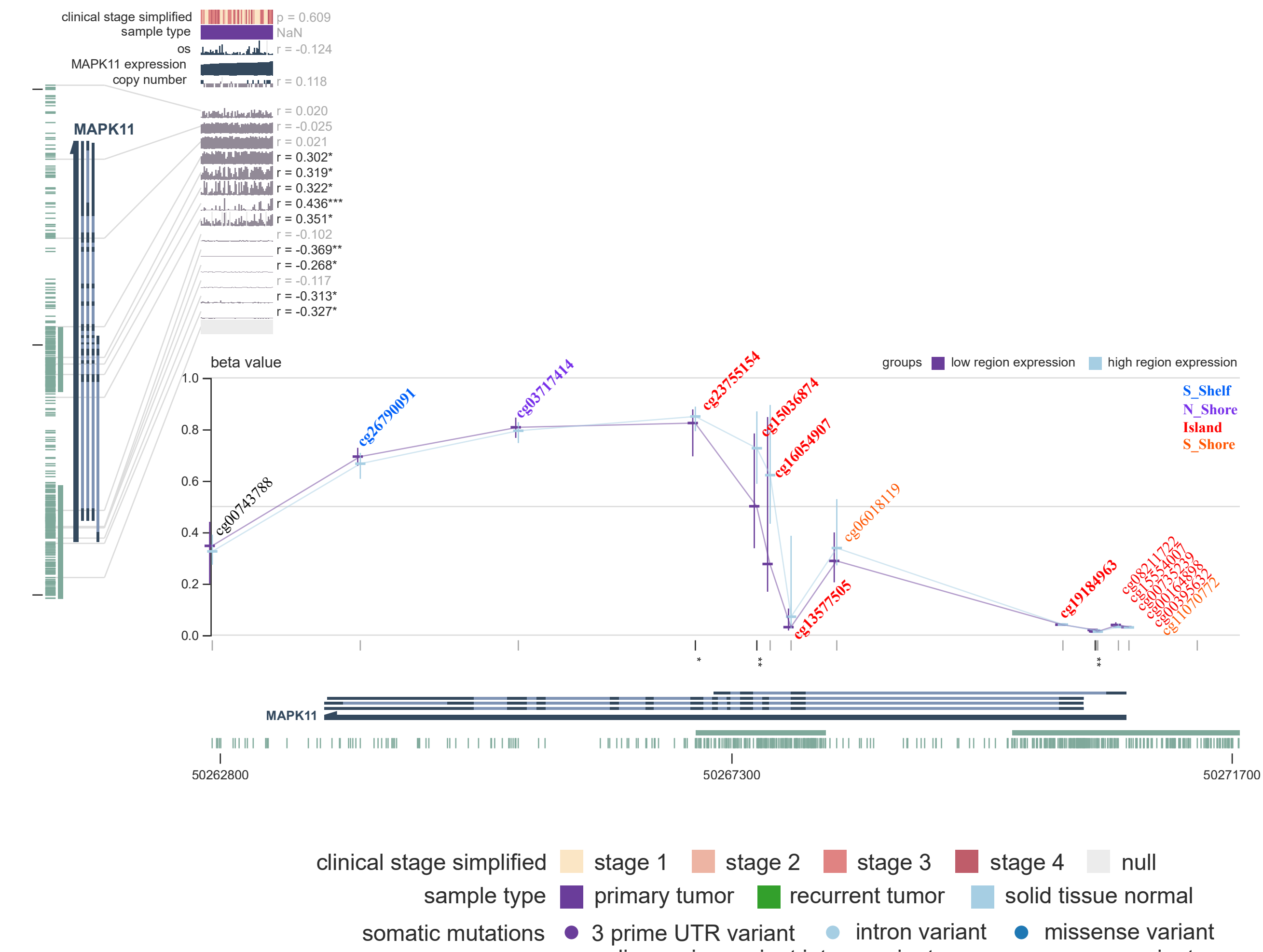
CESC



UCEC



UCS



clinical stage simplified stage 1 stage 2 stage 3 stage 4 null
sample type primary tumor recurrent tumor solid tissue normal
somatic mutations 3 prime UTR variant intron variant missense variant
splice region variant;intron variant synonymous variant
copy number -2: homozygous deletion -1: single copy deletion 0: diploid normal
+1: low-level amplification +2: high-level amplification
statistics p >= 0.05 * p < 0.05 ** p < 0.01 *** p < 0.001