

OBseries consensus matrix for som of 572 isoforms, 324 RBPs, 1984 lncRNAs, 1 snRNAs, 78 TFcomplexs (any qvalue0.01 dPSI greaterthan 0.2), sweep from 3x4 to 7x11

variable

cocluster_percentage

1

gene_isoform_ID