MDS normalised, hc =2 con_sp.hsa.canonical.isoforms_junc.exons protein_groups o ud_1_24 △ ud_30_6 30m 1h 6h √ 1d 0.5 0.0 -0.5 -1.0 -1.0 -0.50.0 0.5

Leading logFC dim 1

Leading logFC dim 2