raw con_sp.hsa.canonical.isoforms_junc.exons protein_groups 3e+09 o ud_1_24 △ ud_30_6 30m 1h 2e+09 6h ∇ 1d Leading logFC dim 2 1e+09 0e+00 -1e+09 -6e+09 -4e+09 -2e+09 0e+00 Leading logFC dim 1