raw con_sp.hsa.canonical.isoforms_junc.exons protein_groups o ud_1_24 1.5 △ ud_30_6 30m 1h 1.0 6h ∇ 1d Leading logFC dim 2 0.5 0.0 -0.5 -1.0 -1.5 0 2 3 Leading logFC dim 1