raw con_sp.hsa.canonical.isoforms_junc.exons peptides 0.8 30m_vs_ud 1h_vs_ud 9.0 + 6h_vs_ud \times 1d_vs_ud 0.4 Leading logFC dim 2 0.2 \times 0.0 -0.2 -0.4 -0.50.0 0.5 Leading logFC dim 1