raw con_sp.hsa.canonical.isoforms peptides X 30m_vs_ud 1h_vs_ud 1.0 + 6h_vs_ud \times 1d_vs_ud 0.5 Leading logFC dim 2 0.0 -0.5-1.0 -1.5 -1.0 -0.5 0.0 0.5 1.0 Leading logFC dim 1