PCA loadings logFC sp.hsa.canonical.isoforms_junc.exons protein_groups 0.4 timepoint 30m 0.2 -1h PC1 (39.8%) 6h 1d Replicate ΙB IB2 KR 0.0 -0.2 0.0 0.6 -0.2 0.2 0.4 PC2 (21.6%)