ratios con_sp.hsa.canonical.isoforms protein_groups -0.28 timepoint 30m_vs_ud 1h_vs_ud PC1 (94.9%) 6h_vs_ud 1d_vs_ud Replicate ΙB IB2 ■ KR -0.30 -0.31 **-**-0.3 0.0 0.3 0.6 -0.6 PC2 (1.38%)

PCA loadings