MDS normalised, hc =2
con_sp.hsa.canonical.isoforms_trembl
protein_groups

Legend:
- 30m_vs_ud (○)
- 1h_vs_ud (△)
- 6h_vs_ud (+)
- 1d_vs_ud (×)

X-axis: Leading logFC dim 1
Y-axis: Leading logFC dim 2