MDS normalised, hc =2 sp.hsa.canonical.isoforms\_junc.exons peptides 30m\_vs\_ud 0.4 1h\_vs\_ud 6h\_vs\_ud  $\times$  1d\_vs\_ud 0.2 Leading logFC dim 2 0.0 -0.2 -0.4 -0.4 -0.2 0.0 0.2 0.4 Leading logFC dim 1