MDS normalised, hc =2 sp.hsa.canonical.isoforms\_junc.exons protein\_groups 30m\_vs\_ud 1h\_vs\_ud 6h\_vs\_ud  $\times$  1d\_vs\_ud 0.5 Leading logFC dim 2 0.0 -0.5 $\times \triangle$ -1.0 -1.5 -1.5 -1.0 -0.50.0 0.5 1.0 Leading logFC dim 1