MDS normalised, hc =2 con\_sp.hsa.canonical.isoforms\_junc.exons protein\_groups 30m\_vs\_ud 1h\_vs\_ud + 6h\_vs\_ud 0.2 × 1d\_vs\_ud Leading logFC dim 2 0.1 0.0 -0.1 -0.2 -0.2-0.10.0 0.1 0.2 0.3 Leading logFC dim 1