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 0.1 -0.1

0.0

Leading logFC dim 1

0.1

0.2

Leading logFC dim 2

-0.3

-0.2

-0.1