IncRNA group = IDRlessthan0\_1 ::: data cols = bothPullDowns (22 / 587) = (func/total IncRNA) ::: biotype = processed\_transcript PageRank, full range 30 -20 -10 factor(label) 0 count 0 1 6 -4 · 2 -

1.0

rank

1.5

0 -

0.0

0.5