IncRNA group = IDRlessthan0\_1 ::: data cols = bothPullDowns (16 / 231) = (func/total lncRNA) ::: biotype = processed\_transcript PageRank, full range 10 -5 factor(label) 0 count 0 1 6 -4 -2 -0 -1.0 1.5 0.0 0.5 rank