lncRNA group = IDRlessthan0\_1 ::: data cols = bothPullDowns (42 / 1363) = (func/total lncRNA) ::: biotype = lincProcTrans PageRank, range = median  $\pm -2$  ds 0 60 -40 -20 factor(label) 0 count 0 1 5 -4 -3 -2 -1 · 0 -8.0 1.2 1.6 0.0 0.4 rank