lncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (9/81) = (func/total lncRNA) ::: biotype = lincRNA PageRank, range = median +/- 2 ds 5 · 4 3 -2 · 1 2.0 factor(label) 0 1 1.5 -1.0 0.5 -0.0 -1.2 0.4 8.0 1.6 rank