lncRNA group = IDRlessthan0\_01 ::: data cols = bothPullDowns (11/109) = (func/total IncRNA) ::: biotype = lincRNAPageRank, range = median +/- 2 ds 0 8 6 -4 2 -2.0 factor(label) 0 1 1.5 -1.0 0.5 0.0 0.4 1.2 8.0 1.6 rank