lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (48 / 2413) = (func/total lncRNA) ::: biotype = lincProcTrans logRank, range = median +/- 2 ds 0 200 -150 100 50 factor(label) 0 0 1 20 10

0.50

logRank

0.75

0

0.00

0.25