lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (30 / 1701) = (func/total IncRNA) ::: biotype = antisense logRank, range = median +/- 2 ds 100 -75 50 25 factor(label) 0 count 0 1 8 -6 -4 2 0 0.50 0.25 0.75 0.00 logRank