lncRNA group = IDRlessthan0\_01 ::: data cols = bothPullDowns (11/109) = (func/total IncRNA) ::: biotype = lincRNAlogRank, range = median +/- 2 ds 6 4 2 factor(label) 0 count 0 1.00 -0.75 -0.50 0.25 0.00 0.6 0.4 8.0 1.0 logRank