lncRNA group = IDRlessthan0\_01 ::: data cols = bothPullDowns (21/146) = (func/total IncRNA) ::: biotype = lincProcTrans logRank, range = median +/- 2 ds 0 7.5 5.0 2.5 count factor(label) 0 1 2.0 -1.5 -1.0 0.5 0.0 -0.8 0.4 0.6 1.0 logRank