lncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (25 / 277) = (func/total lncRNA) ::: biotype = all_biotypes logRank, range = median +/- 2 ds 0 9 -6 3 -2.0 factor(label) 0 1 1.5 1.0 0.5 0.0 0.4 0.6 8.0 1.0 logRank