IncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (69 / 2573) = (func/total lncRNA) ::: biotype = all_biotypes logRank, range = median +/- 2 ds 0 60 40 -20 factor(label) 2.0 -0 1 1.5 -1.0 0.5 -0.0 -0.4 0.6 8.0 1.0 logRank