IncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (23 / 901) = (func/total IncRNA) ::: biotype = antisense logRank, range = median +/- 2 ds 0 25 -20 15 -10 -5 · 2.0 factor(label) 0 1 1.5 -1.0 0.5 0.0 0.4 0.6 8.0 1.0 logRank