IncRNA group = IDRlessthan0\_1 ::: data cols = bothPullDowns (23 / 901) = (func/total IncRNA) ::: biotype = antisense logRank, range = median +/- 2 ds 2.0 -1.5 factor(label) density 0.0 0.5 -0.0 0.4 0.6 8.0 1.0 logRank