IncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (24 / 1026) = (func/total IncRNA) ::: biotype = antisense logRank, range = median +/- 2 ds 2.0 1.5 factor(label) density 0.1 0.5 -0.0 0.4 0.6 8.0 1.0

logRank