lncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (45 / 1554) = (func/total lncRNA) ::: biotype = remove_antisense logRank, range = median +/- 2 ds 1.5 -1.0 factor(label) density 0.5 -0.0 -

0.6

logRank

8.0

1.0

0.4

0.2