lncRNA group = IDRlessthan0_2 ::: data cols = Inpa (82 / 5010) = (func/total lncRNA) ::: biotype = all_biotypes logRank, range = median +/- 2 ds 0 400 300 200 -100 factor(label) 0 0 1 20 10 0

0.50

0.75

0.25

logRank

0.00