lncRNA group = IDRlessthan0_1 ::: data cols = Inpa (69 / 2833) = (func/total lncRNA) ::: biotype = all_biotypes logRank, range = median +/- 2 ds 2.0 -1.5 factor(label) density 0.1 0.5 -0.0 0.50 0.00 0.25 0.75

logRank