lncRNA group = IDRlessthan0\_1 ::: data cols = Inpa ( 69 / 2832 ) = (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