lncRNA group = IDRlessthan0\_1 ::: data cols = lpa ( 23 / 939 ) = (func/total IncRNA) ::: biotype = lincRNA logRank, range = median +/- 2 ds 30 20 10 factor(label) 0 0 1.00 -0.75 0.50 0.25 0.00 0.6 0.4 0.2 8.0 1.0 logRank

count