lncRNA group = IDRlessthan0_01 ::: data cols = lpa (25 / 277) = (func/total lncRNA) ::: biotype = all_biotypes logRank, range = median +/- 2 ds 12.5 10.0 7.5 5.0 2.5 factor(label) 0.0 0 1 2.0 1.5 -1.0 0.5 0.0 0.4 0.6 8.0 1.0 logRank

count