lncRNA group = IDRlessthan0_1 ::: data cols = lpa (23 / 945) = (func/total IncRNA) ::: biotype = lincRNA logRank, range = median +/- 2 ds 30 20 10 factor(label) 0 0 1 1.00 -0.75 -0.50 0.25 0.00 0.6 0.4 8.0 1.0 0.2 logRank

count