lncRNA group = IDRlessthan0\_01 ::: data cols = bothPullDowns (20 / 212) = (func/total IncRNA) ::: biotype = remove\_antisense logRank, range = median +/- 2 ds12.5 10.0 7.5 5.0 2.5 factor(label) 0.0 0 1 2.0 1.5 -1.0 0.5 0.0 0.6 0.4 0.8 1.0 logRank

count