IncRNA group = IDRlessthan0\_1 ::: data cols = bothPullDowns ( 69 / 2581 ) = (func/total lncRNA) ::: biotype = all\_biotypes logRank, range = median +/- 2 ds0 60 40 -20 factor(label) 2.0 -0 1 1.5 1.0 0.5 0.0 0.6 0.4 8.0 1.0 logRank