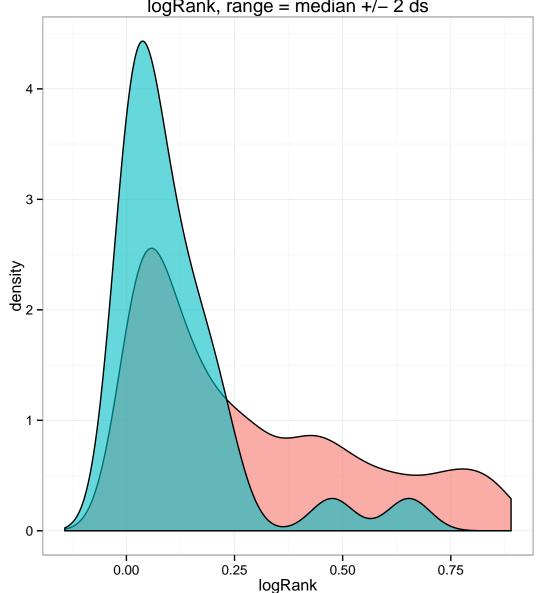
IncRNA group = allLncRNA ::: data cols = bothPullDowns (36 / 2243) = (func/total lncRNA) ::: biotype = antisense logRank, range = median +/- 2 ds



factor(label)