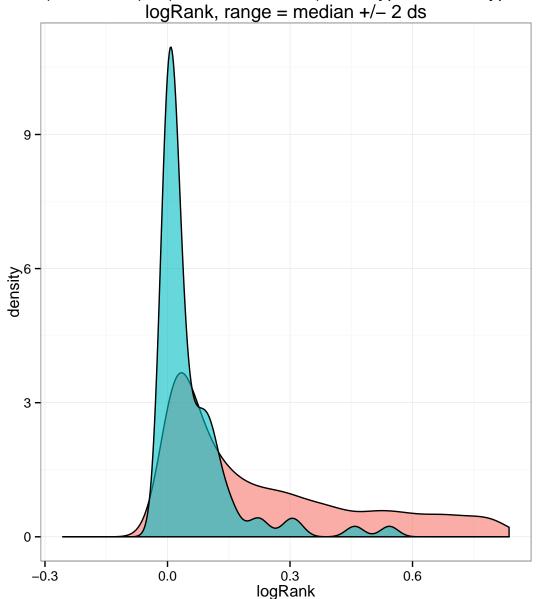
IncRNA group = allLncRNA ::: data cols = bothPullDowns (92 / 7622) = (func/total IncRNA) ::: biotype = all\_biotypes logRank\_range = median +/- 2 ds



factor(label)