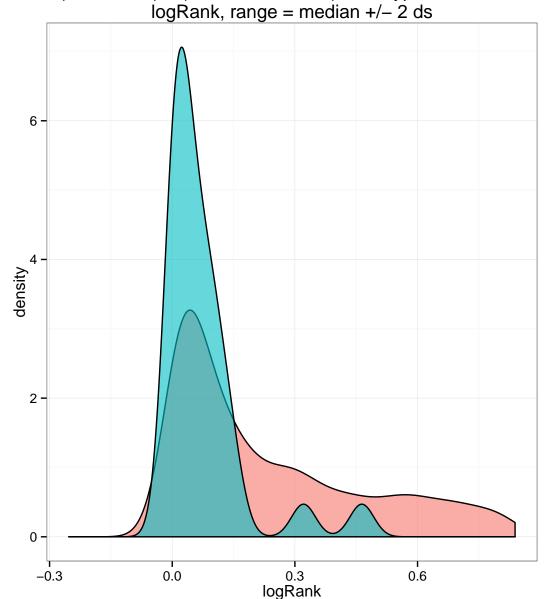
IncRNA group = allLncRNA ::: data cols = bothPullDowns (39 / 2750) = (func/total IncRNA) ::: biotype = antisense logRank, range = median +/- 2 ds



factor(label)