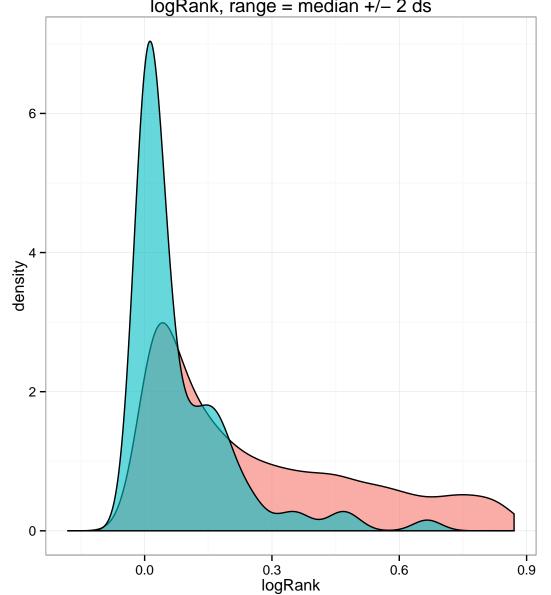
IncRNA group = allLncRNA ::: data cols = bothPullDowns (87 / 6104) = (func/total IncRNA) ::: biotype = all\_biotypes logRank, range = median +/- 2 ds



factor(label)