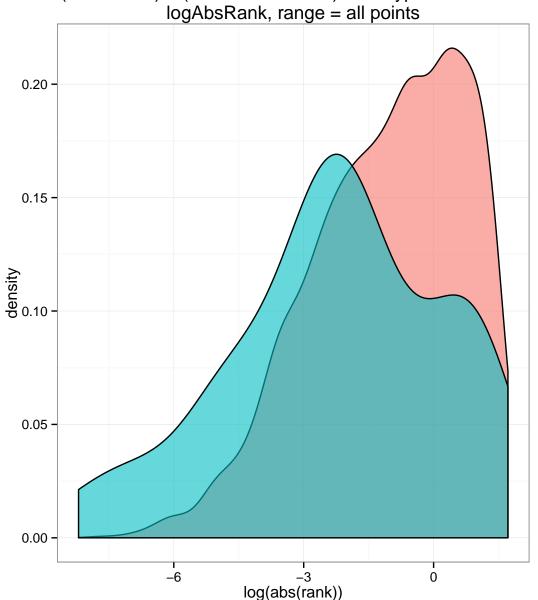
IncRNA group = allLncRNA ::: data cols = bothPullDowns (36 / 2243) = (func/total IncRNA) ::: biotype = antisense logAbsRank, range = all points



factor(label)