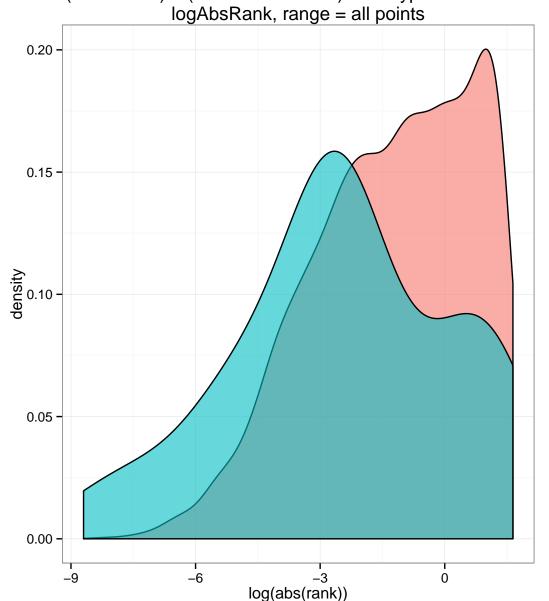
IncRNA group = allLncRNA ::: data cols = lpa ( 38 / 2567 ) = (func/total IncRNA) ::: biotype = antisense logAbsRank, range = all points



factor(label)