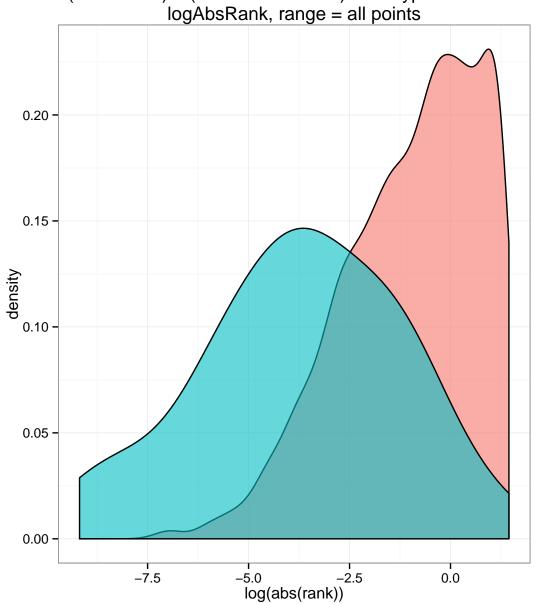
IncRNA group = allLncRNA ::: data cols = Inpa ( 30 / 2632 ) = (func/total IncRNA) ::: biotype = lincRNA logAbsRank, range = all points



factor(label)