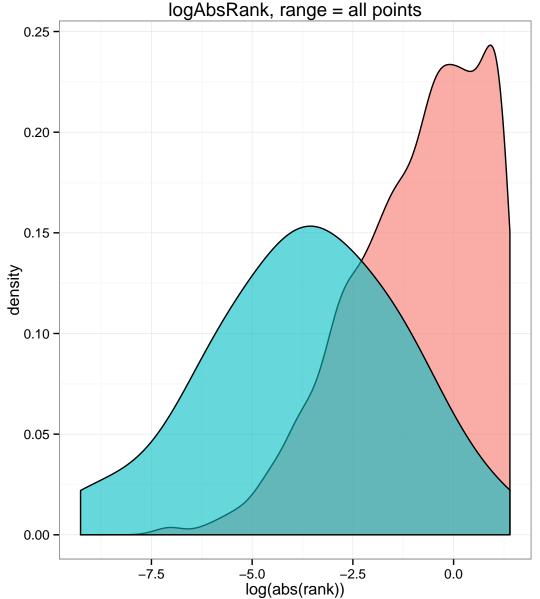
IncRNA group = allLncRNA ::: data cols = Inpa (27 / 2336) = (func/total IncRNA) ::: biotype = lincRNA logAbsRank, range = all points



factor(label)