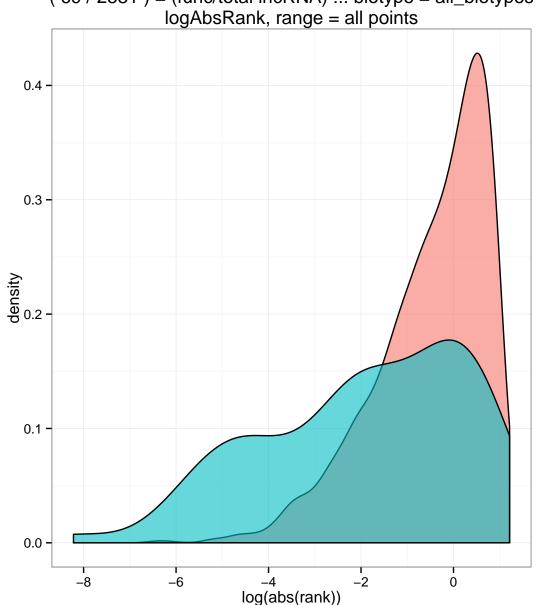
lncRNA group = IDRlessthan0\_1 ::: data cols = lpa (69 / 2581) = (func/total lncRNA) ::: biotype = all\_biotypes logAbsRank, range = all points



factor(label)