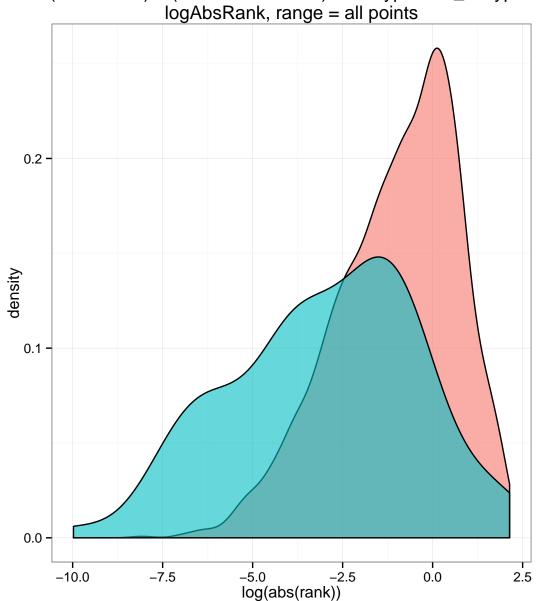
IncRNA group = IDRlessthan0_2 ::: data cols = Ipa (82 / 5045) = (func/total IncRNA) ::: biotype = all_biotypes logAbsRank, range = all points



factor(label)