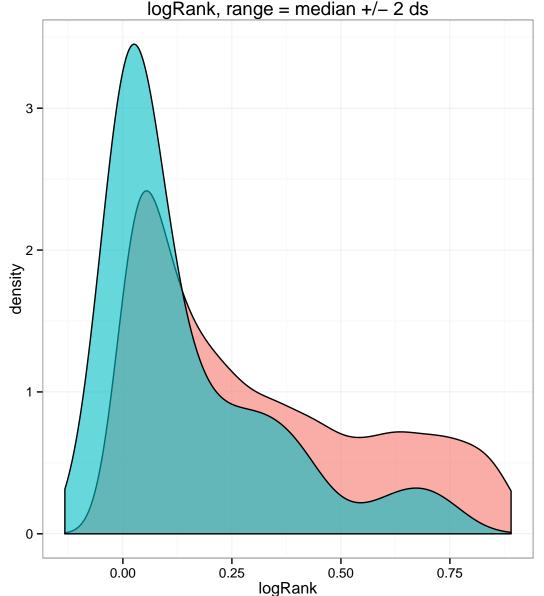
IncRNA group = IDRlessthan0_2 ::: data cols = Ipa (82 / 5045) = (func/total IncRNA) ::: biotype = all_biotypes logRank, range = median +/- 2 ds



factor(label)