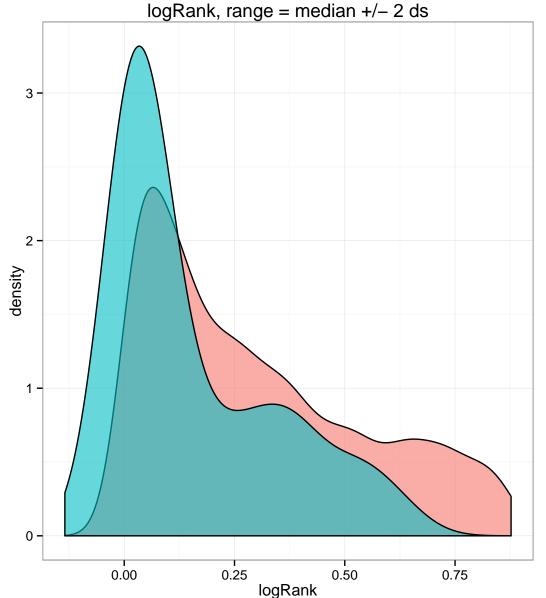
IncRNA group = IDRlessthan0_2 ::: data cols = Inpa (81 / 4501) = (func/total IncRNA) ::: biotype = all_biotypes logRank, range = median +/- 2 ds



factor(label)