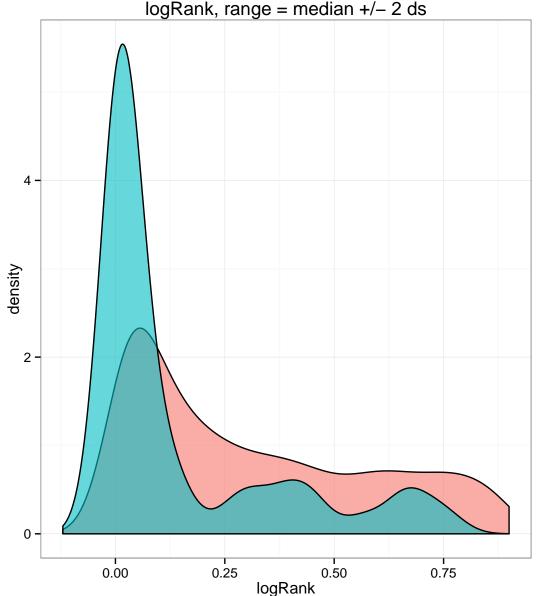
IncRNA group = IDRlessthan0_2 ::: data cols = Ipa (50 / 2684) = (func/total IncRNA) ::: biotype = remove_antisense logRank, range = median +/- 2 ds



factor(label)