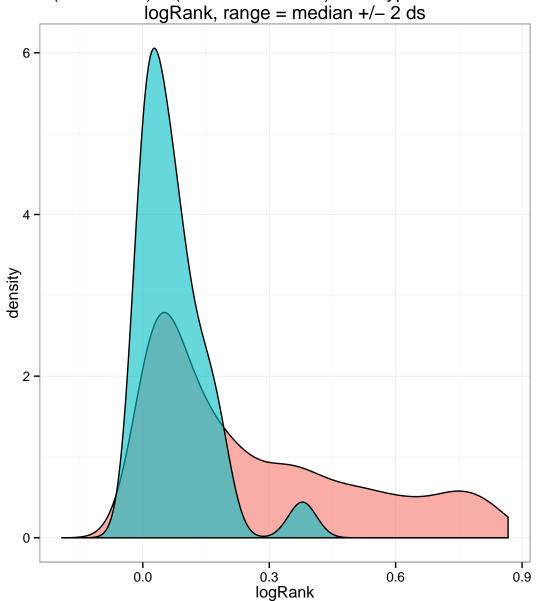
IncRNA group = allLncRNA ::: data cols = Ipa (36 / 2193) = (func/total IncRNA) ::: biotype = antisense logRank, range = median +/- 2 ds



factor(label)