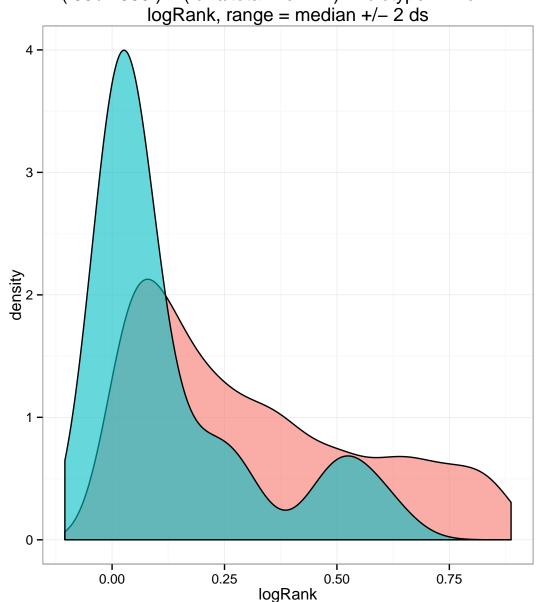
IncRNA group = IDRlessthan0_2 ::: data cols = Inpa (30 / 1980) = (func/total IncRNA) ::: biotype = lincRNA logRank, range = median +/- 2 ds



factor(label)