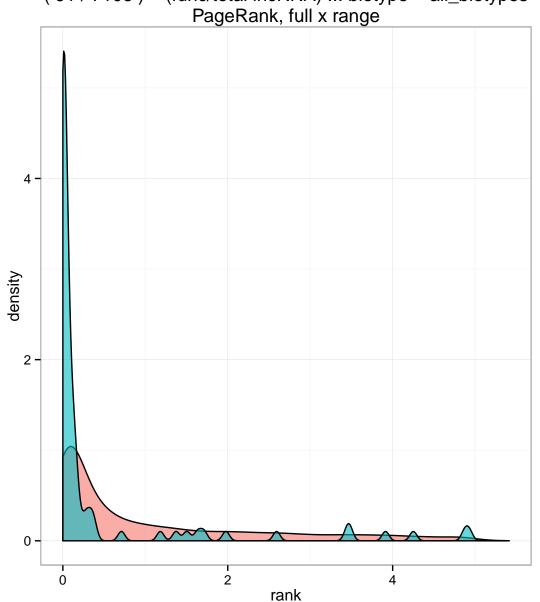
IncRNA group = allLncRNA ::: data cols = lpa ( 91 / 7106 ) = (func/total IncRNA) ::: biotype = all\_biotypes



factor(label)