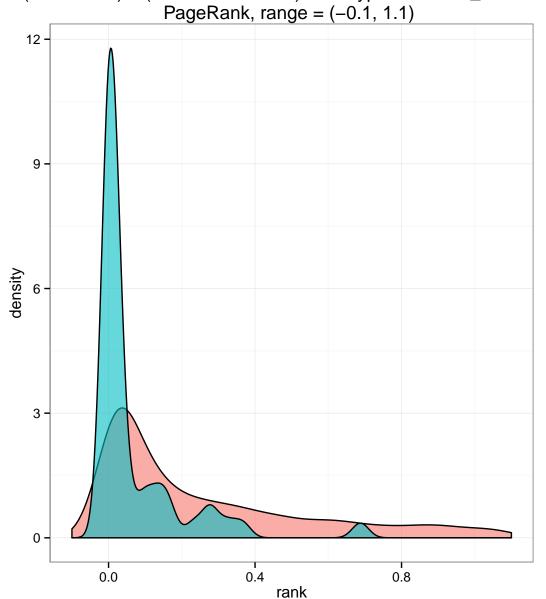
IncRNA group = allLncRNA ::: data cols = lpa (52 / 3884) = (func/total lncRNA) ::: biotype = remove_antisense PageRank, range = (-0.1, 1.1)



factor(label)