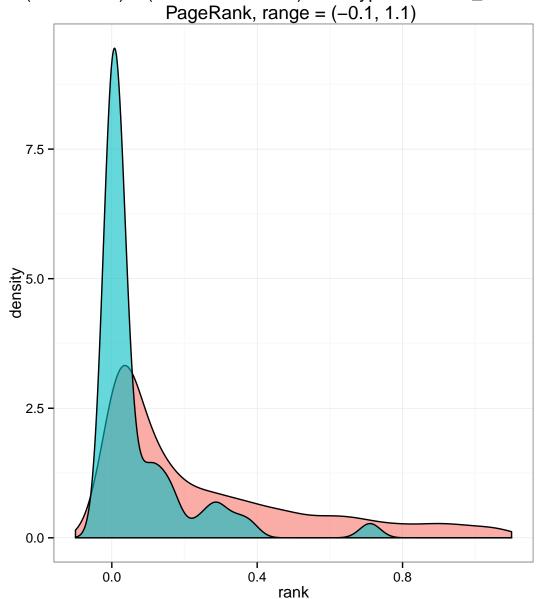
IncRNA group = allLncRNA ::: data cols = lpa (54 / 4747) = (func/total lncRNA) ::: biotype = remove_antisense PageRank, range = (-0.1, 1.1)



factor(label)