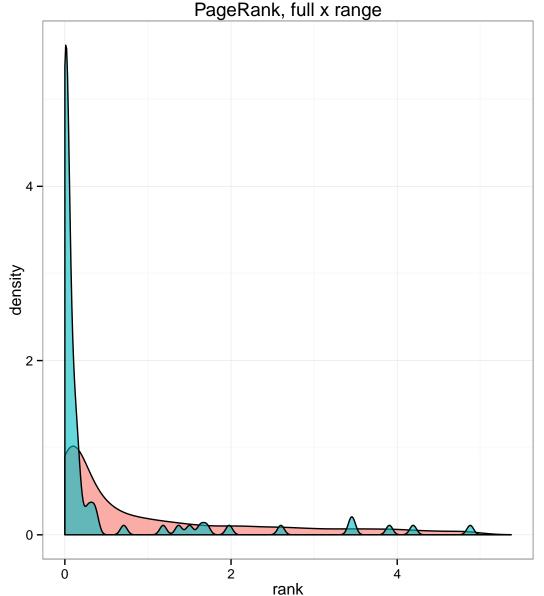
IncRNA group = allLncRNA ::: data cols = Ipa (90 / 6451) = (func/total IncRNA) ::: biotype = all_biotypes PageRank, full x range



factor(label)