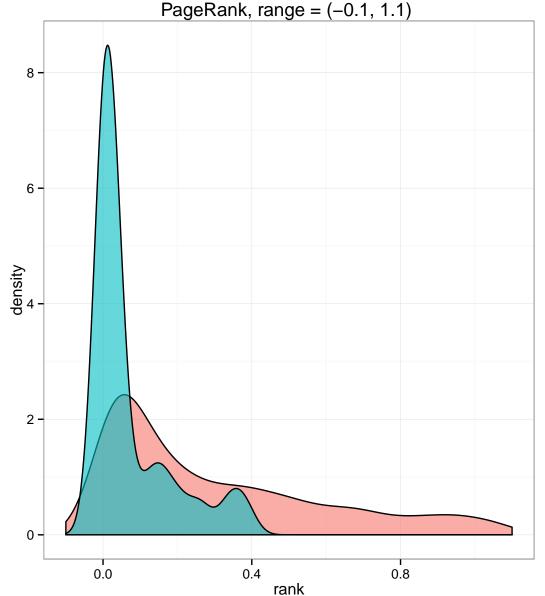
IncRNA group = allLncRNA ::: data cols = bothPullDowns (27 / 2546) = (func/total IncRNA) ::: biotype = lincRNA PageRank, range = (-0.1, 1.1)



factor(label)