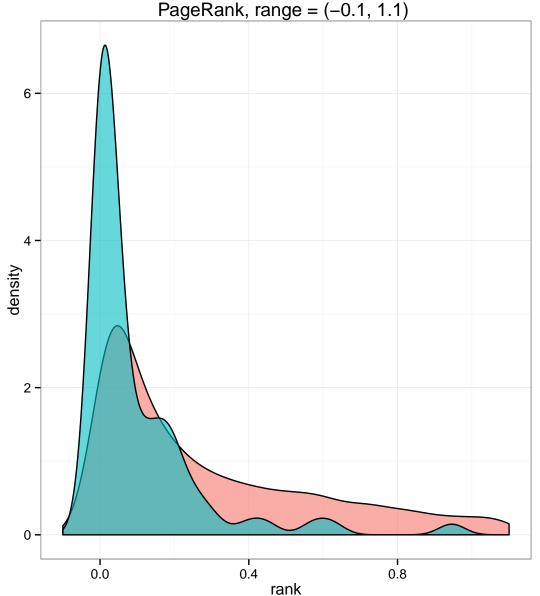
IncRNA group = allLncRNA ::: data cols = bothPullDowns (87 / 6104) = (func/total IncRNA) ::: biotype = all_biotypes PageRank, range = (-0.1, 1.1)



factor(label)