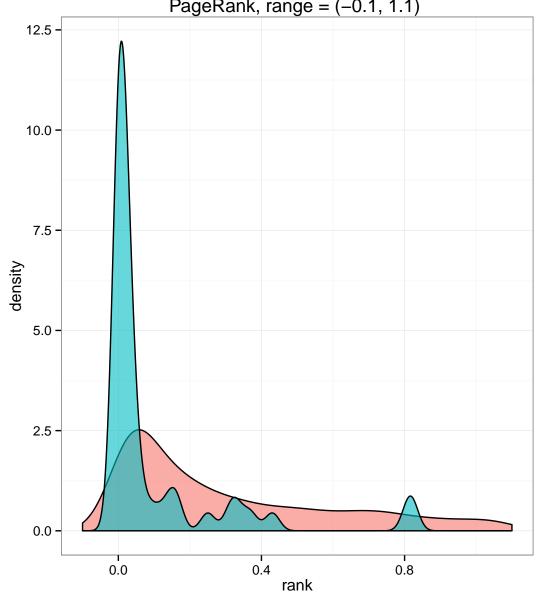
IncRNA group = allLncRNA ::: data cols = Inpa (47 / 3175) = (func/total IncRNA) ::: biotype = lincProcTrans PageRank, range = (-0.1, 1.1)



factor(label)