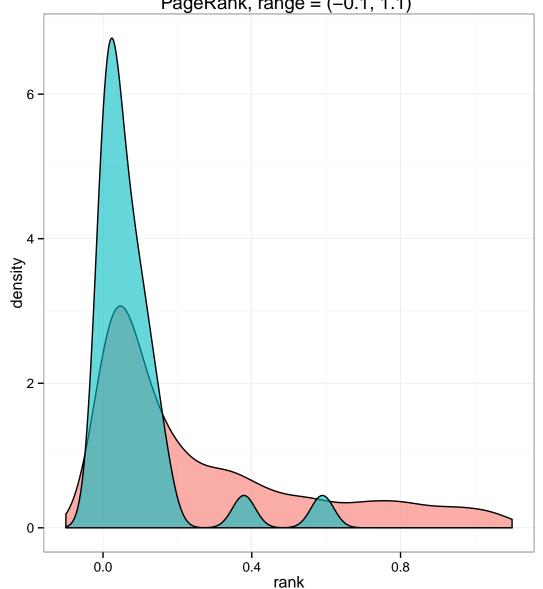
IncRNA group = allLncRNA ::: data cols = bothPullDowns (39 / 2750) = (func/total IncRNA) ::: biotype = antisense PageRank, range = (-0.1, 1.1)



factor(label)