lncRNA group = allLncRNA ::: data cols = bothPullDowns
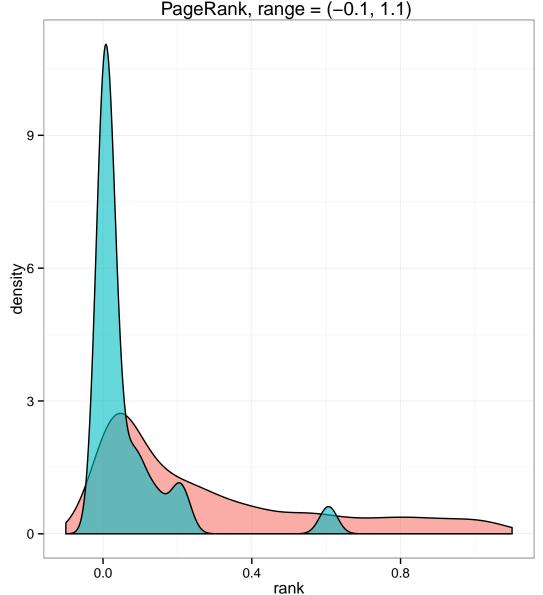(28 / 2998) = (func/total lncRNA) ::: biotype = lincRNA
PageRank, range = (−0.1, 1.1)