lncRNA group = allLncRNA ::: data cols = lpa
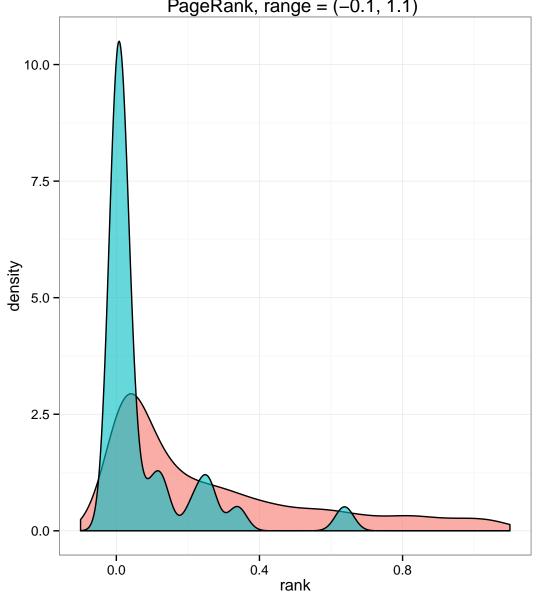(32 / 3068) = (func/total lncRNA) ::: biotype = lincRNA
PageRank, range = (−0.1, 1.1)