lncRNA group = allLncRNA ::: data cols = lnpa
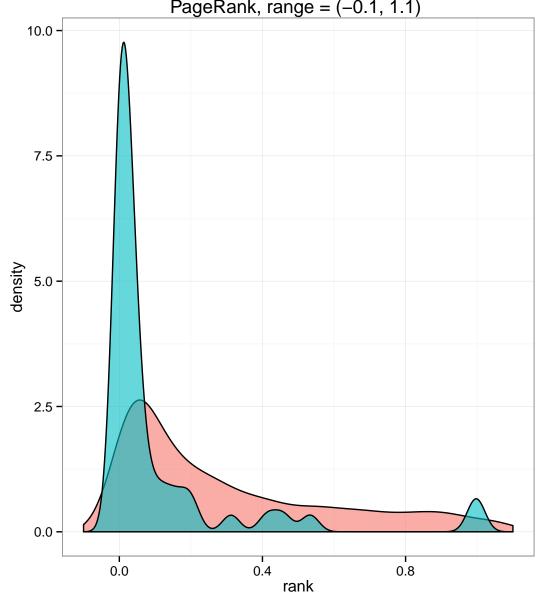(52 / 4230) = (func/total lncRNA) ::: biotype = remove_antisense
PageRank, range = (−0.1, 1.1)