lncRNA group = allLncRNA ::: data cols = lnpa
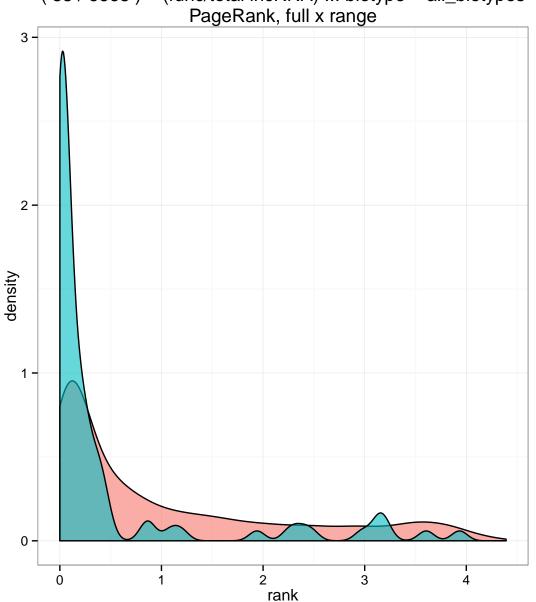(88 / 6668) = (func/total lncRNA) ::: biotype = all_biotypes
PageRank, full x range