lncRNA group = allLncRNA ::: data cols = lpa
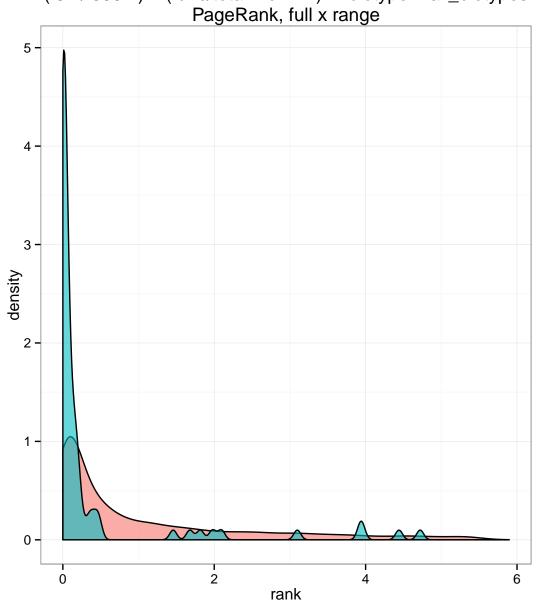(87 / 5951) = (func/total lncRNA) ::: biotype = all_biotypes
PageRank, full x range