lncRNA group = IDRlessthan0_2 ::: data cols = lpa
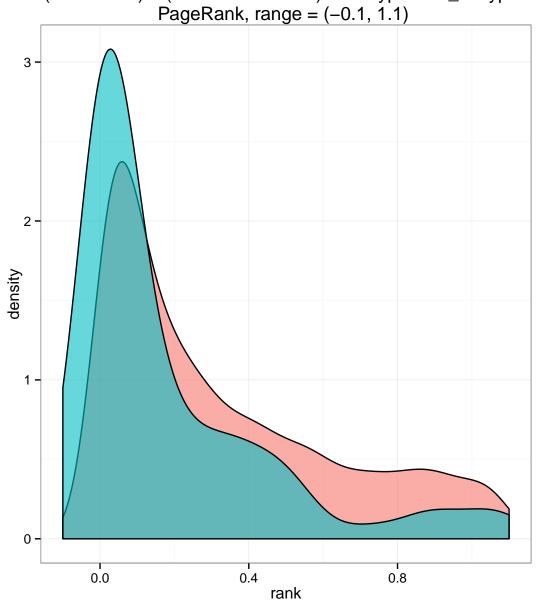(82 / 5045) = (func/total lncRNA) ::: biotype = all_biotypes
PageRank, range = (−0.1, 1.1)