lncRNA group = IDRlessthan0_2 ::: data cols = lpa
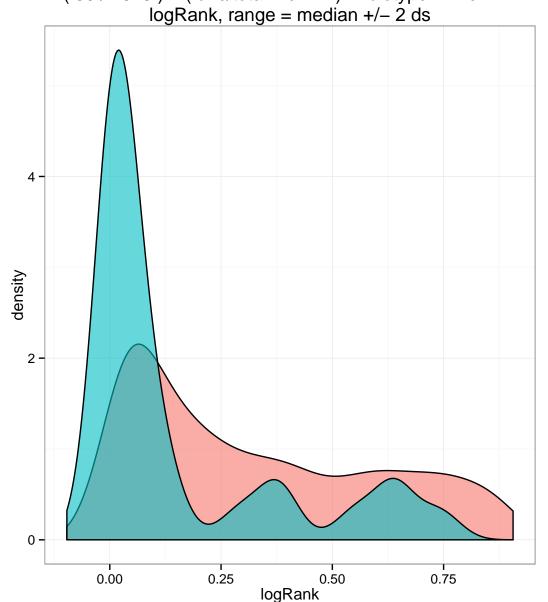(30 / 2025) = (func/total lncRNA) ::: biotype = lincRNA
logRank, range = median +/− 2 ds