lncRNA group = IDRlessthan0_1 ::: data cols = lpa
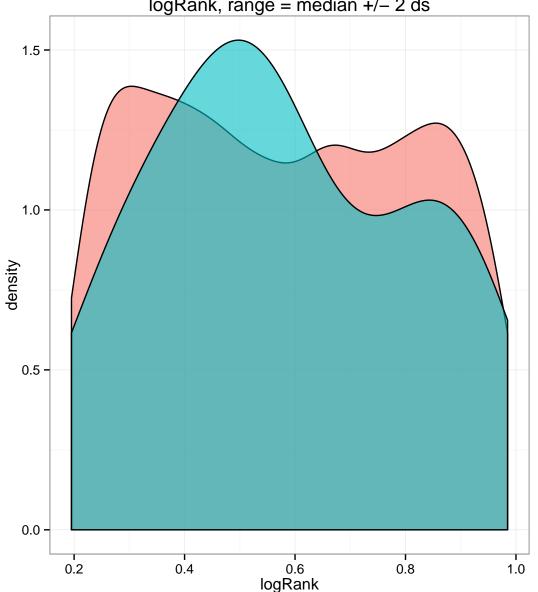(69 / 2856) = (func/total lncRNA) ::: biotype = all_biotypes
logRank, range = median +/− 2 ds