lncRNA group = IDRlessthan0_2 ::: data cols = lpa
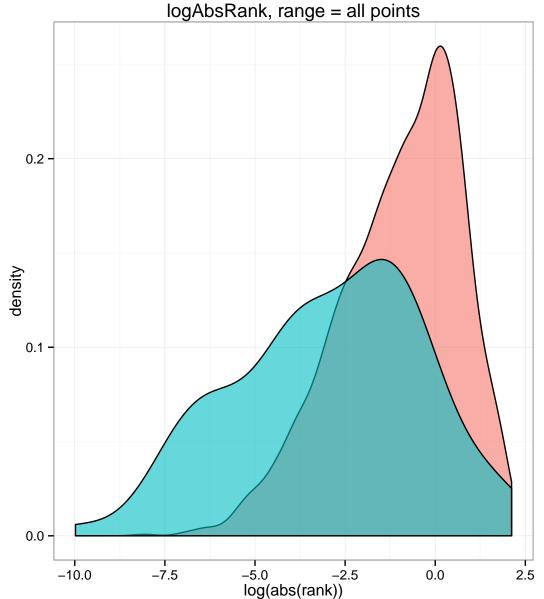(83 / 5087) = (func/total lncRNA) ::: biotype = all_biotypes
logAbsRank, range = all points