lncRNA group = IDRlessthan0_01 ::: data cols = lpa
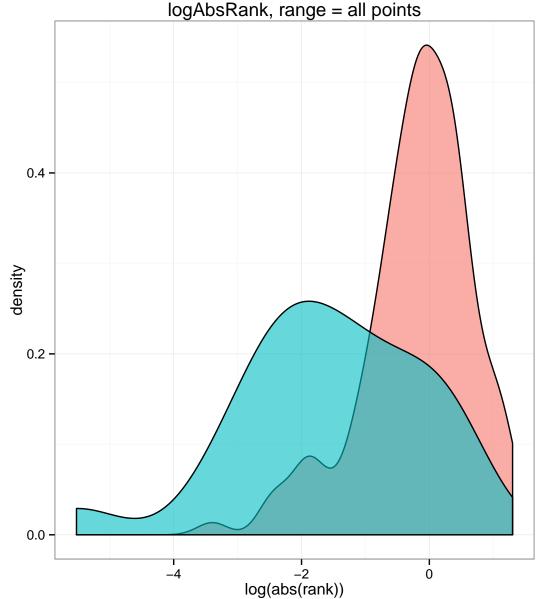(21 / 146) = (func/total lncRNA) ::: biotype = lincProcTrans
logAbsRank, range = all points