lncRNA group = allLncRNA ::: data cols = lnpa
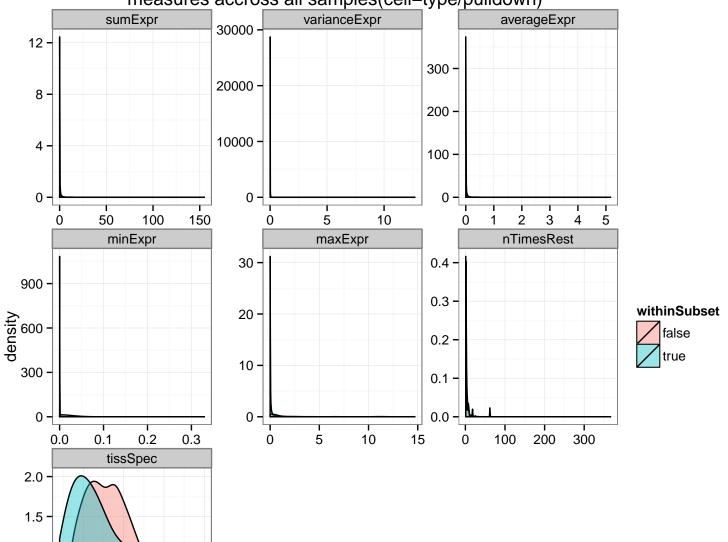( 30 / 2803 ) = (func/total lncRNA) ::: biotype = lincRNA
measures accross all samples(cell−type/pulldown)