lncRNA group = allLncRNA ::: data cols = lnpa
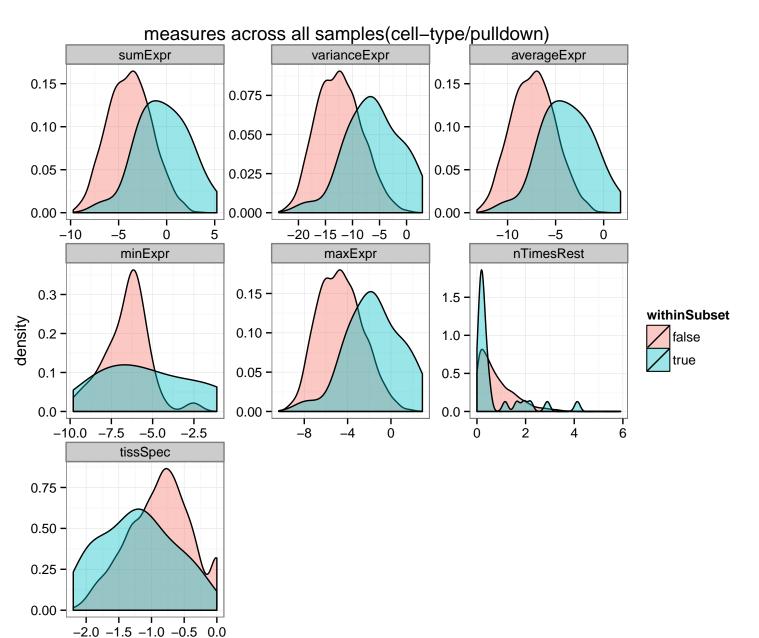(27 / 2505) = (func/total lncRNA) ::: biotype = lincRNA
subset =
27 lncRNAs out of 2505 total lncRNAs

measures across all samples(cell−type/pulldown)