lncRNA group = allLncRNA ::: data cols = lpa
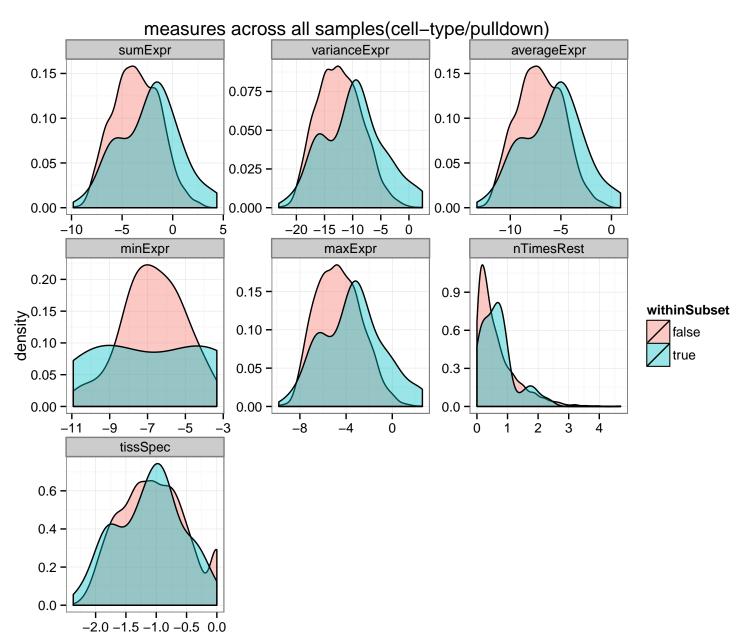(37 / 2359) = (func/total lncRNA) ::: biotype = antisense
subset =
37 lncRNAs out of 2359 total lncRNAs