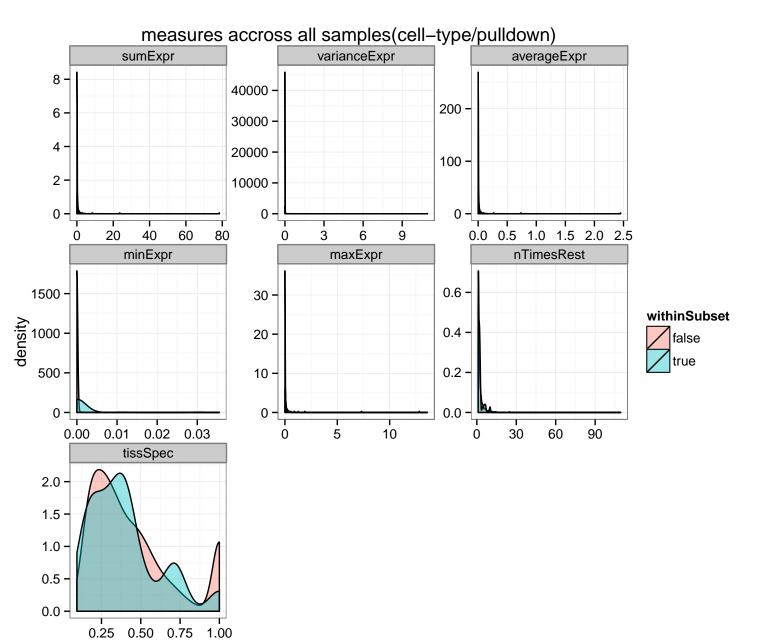# lncRNA group = allLncRNA ::: data cols = bothPullDowns
(38 / 2538) = (func/total lncRNA) ::: biotype = antisense
subset =
38 lncRNAs out of 2538 total lncRNAs

measures accross all samples(cell−type/pulldown)