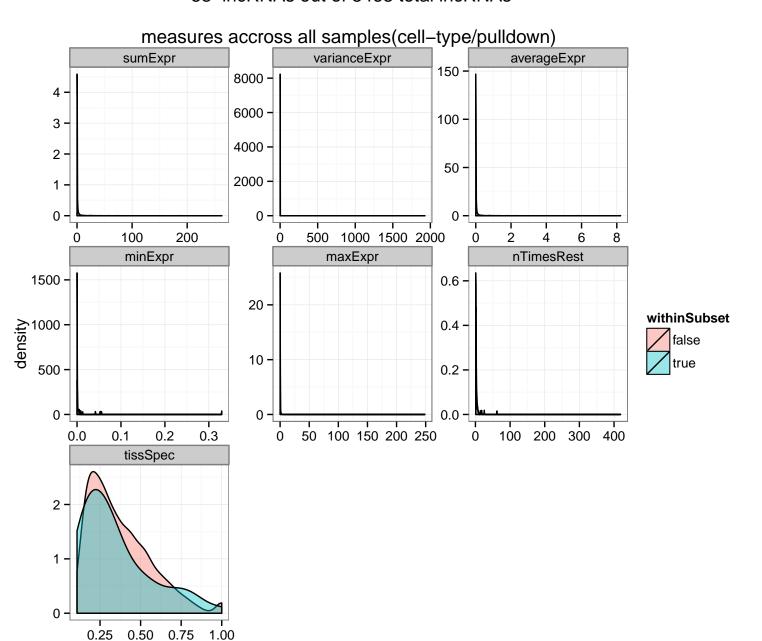# lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns
(53 / 3463) = (func/total lncRNA) ::: biotype = remove_antisense
subset =
53 lncRNAs out of 3463 total lncRNAs

measures accross all samples(cell−type/pulldown)