lncRNA group = allLncRNA ::: data cols = lnpa
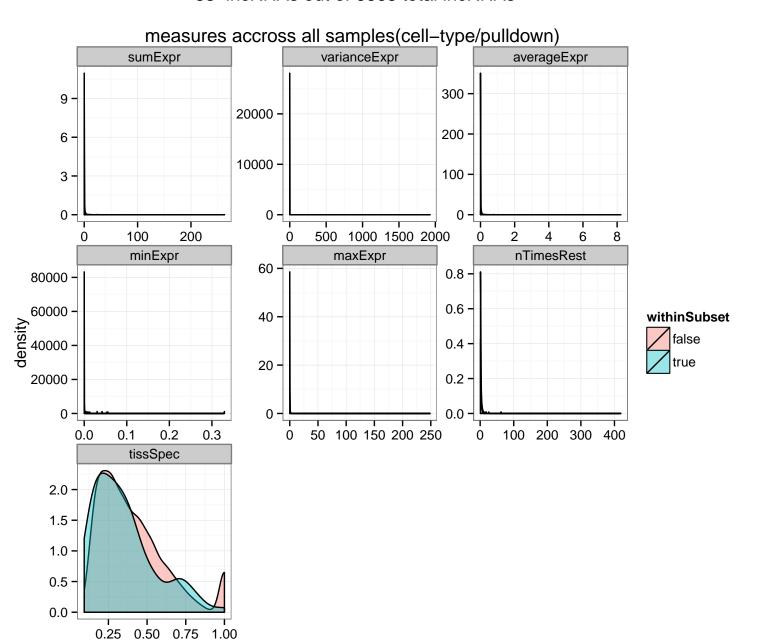(88 / 6668) = (func/total lncRNA) ::: biotype = all_biotypes
subset =
88 lncRNAs out of 6668 total lncRNAs

measures accross all samples(cell−type/pulldown)