lncRNA group = IDRlessthan0_2 ::: data cols = lpa
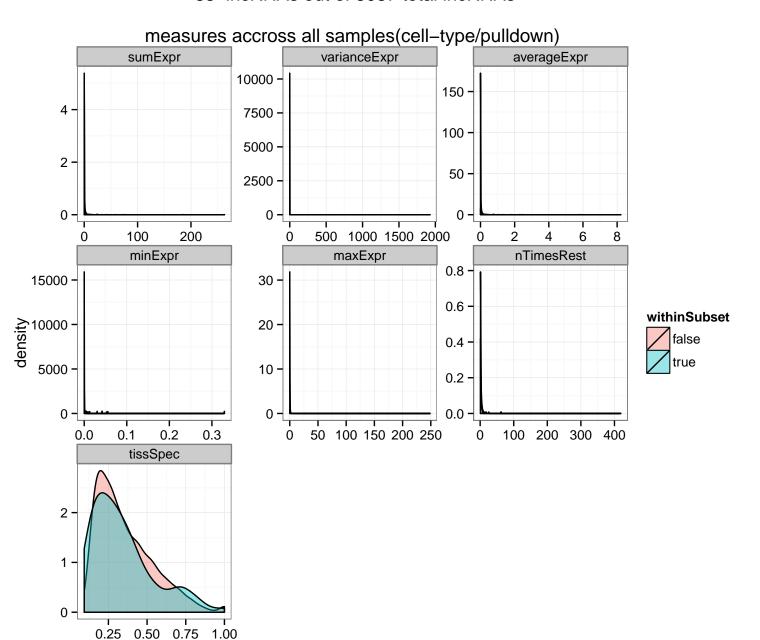(83 / 5087) = (func/total lncRNA) ::: biotype = all_biotypes
subset =
83 lncRNAs out of 5087 total lncRNAs

measures accross all samples(cell−type/pulldown)