lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns
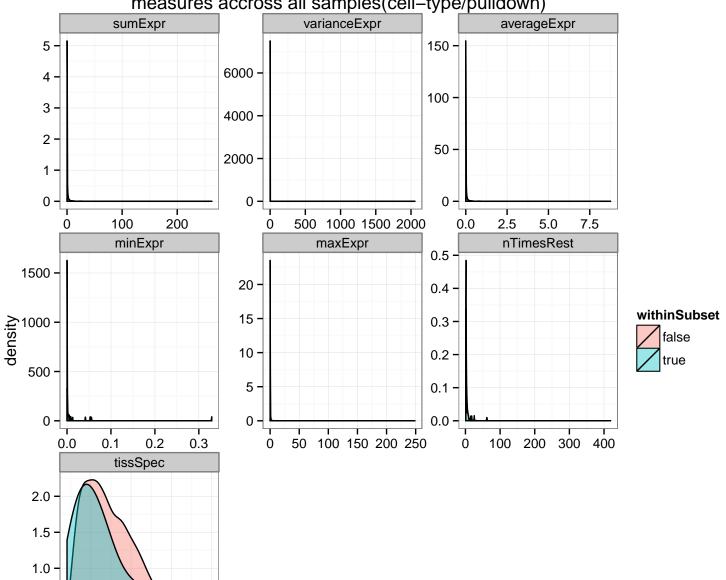(51 / 2754) = (func/total lncRNA) ::: biotype = remove_antisense
measures accross all samples(cell−type/pulldown)