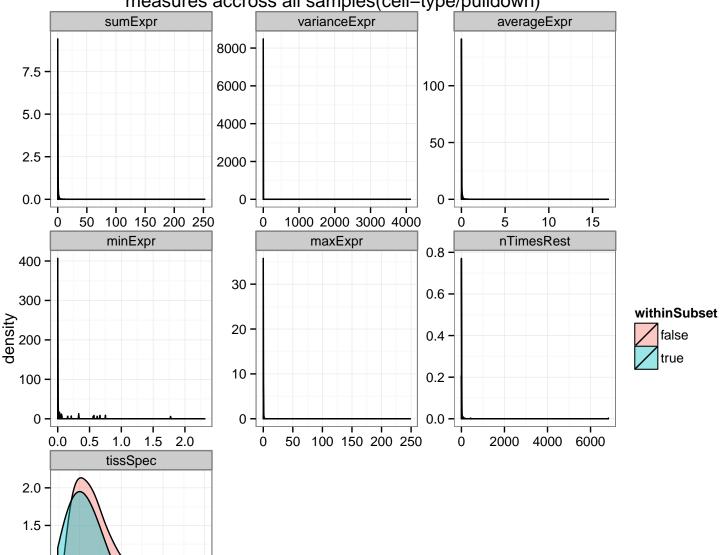lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns
(82 / 4643) = (func/total lncRNA) ::: biotype = all_biotypes
measures accross all samples(cell−type/pulldown)