l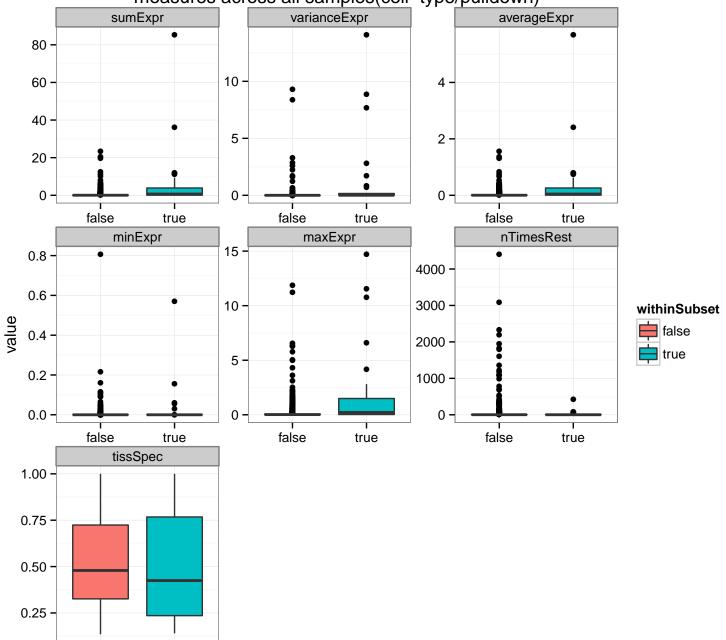ncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (31 / 2082) = (func/total lncRNA) ::: biotype = lincRNA measures across all samples(cell−type/pulldown)