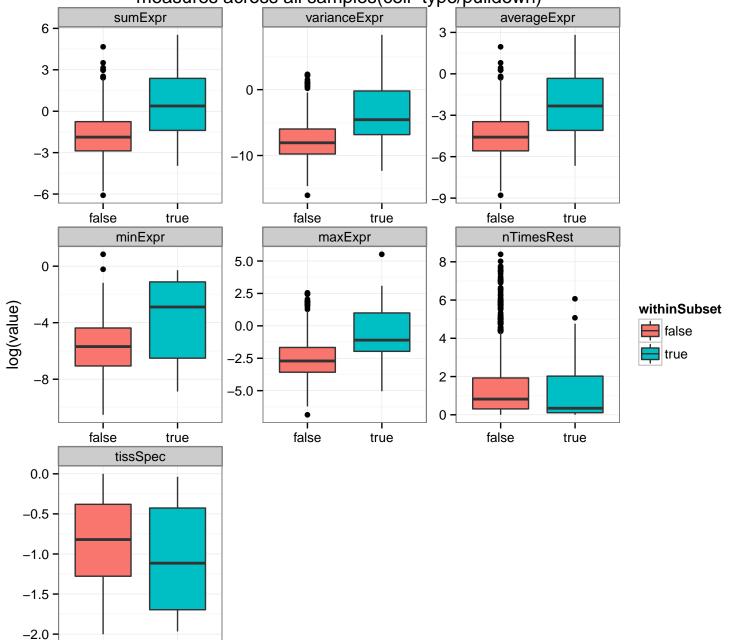lncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns
(42 / 1363) = (func/total lncRNA) ::: biotype = lincProcTrans
measures across all samples(cell−type/pulldown)