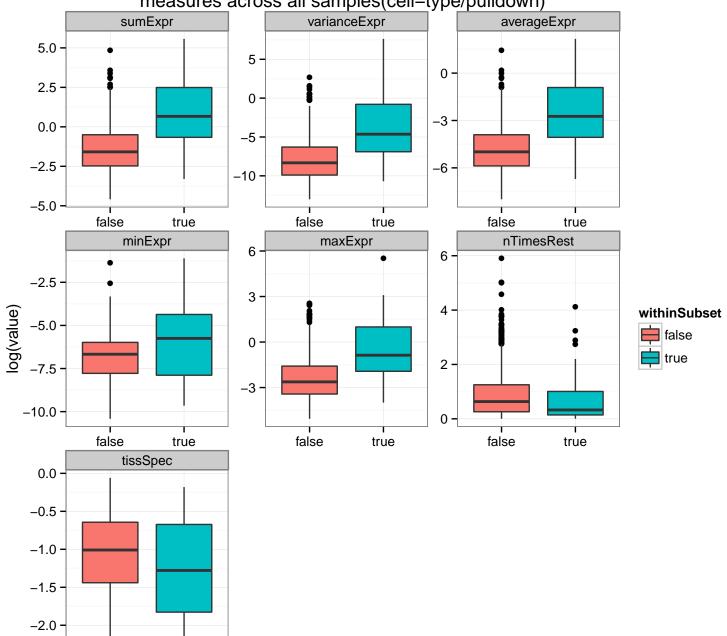lncRNA group = IDRlessthan0_1 ::: data cols = lnpa
(42 / 1350) = (func/total lncRNA) ::: biotype = lincProcTrans
measures across all samples(cell−type/pulldown)