lncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns
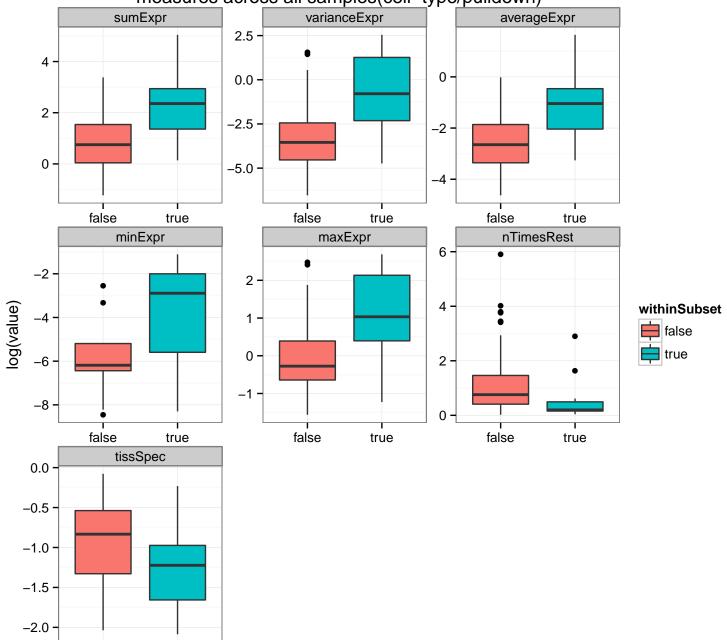(11 / 109) = (func/total lncRNA) ::: biotype = lincRNA
measures across all samples(cell−type/pulldown)