lncRNA group = allLncRNA ::: data cols = lpa
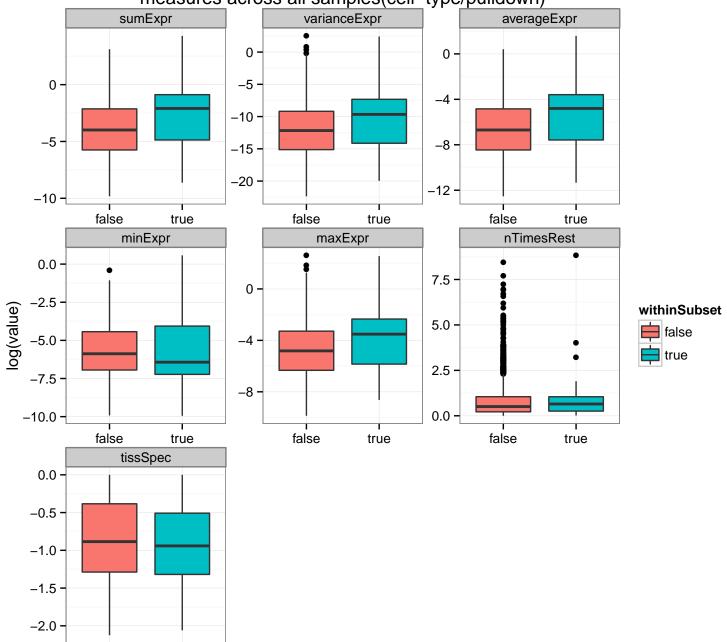(38 / 2567) = (func/total lncRNA) ::: biotype = antisense
measures across all samples(cell–type/pulldown)