lncRNA group = allLncRNA ::: data cols = lnpa
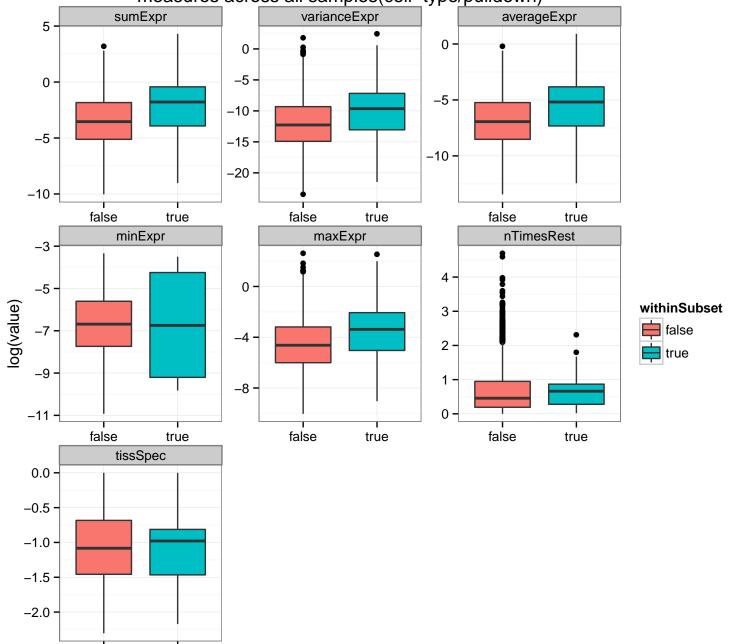(36 / 2219) = (func/total lncRNA) ::: biotype = antisense
measures across all samples(cell−type/pulldown)