lncRNA group = allLncRNA ::: data cols = lnpa
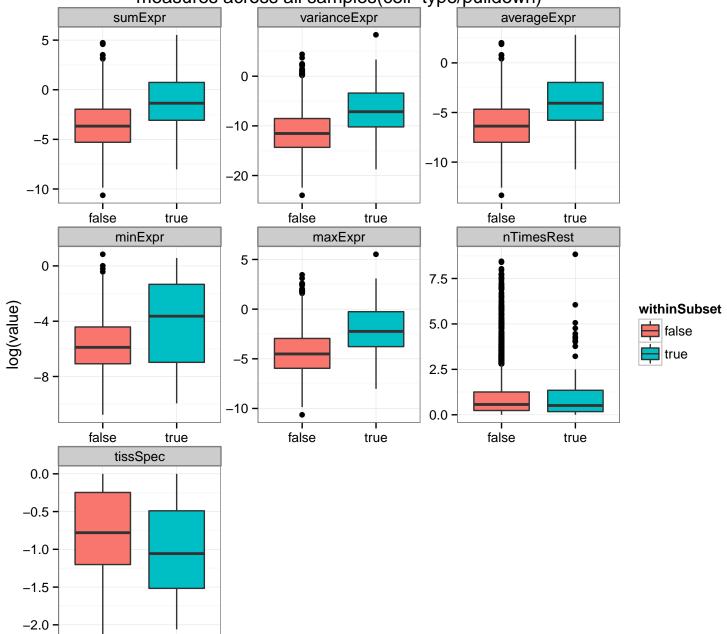(87 / 6034) = (func/total lncRNA) ::: biotype = all_biotypes
measures across all samples(cell−type/pulldown)