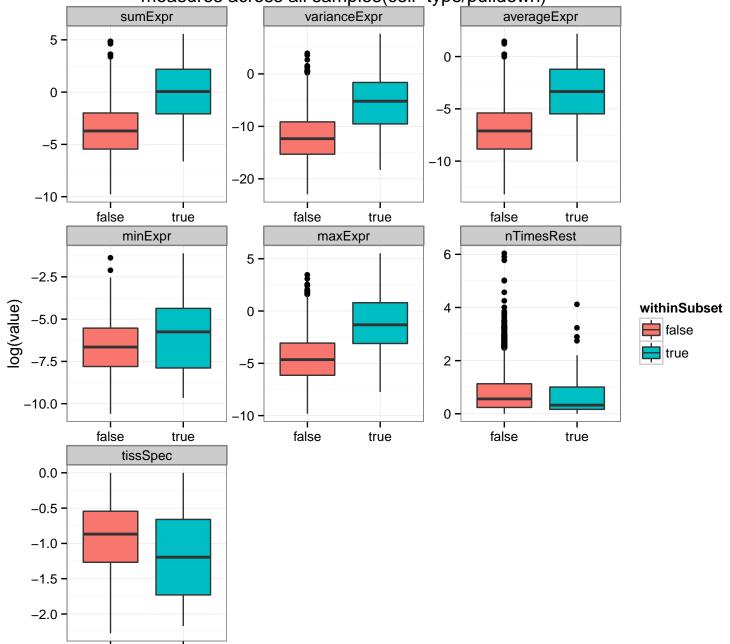
IncRNA group = allLncRNA ::: data cols = lpa (52 / 3884) = (func/total IncRNA) ::: biotype = remove_antisense measures across all samples(cell_type/pulldown)



value of facet label

false

true