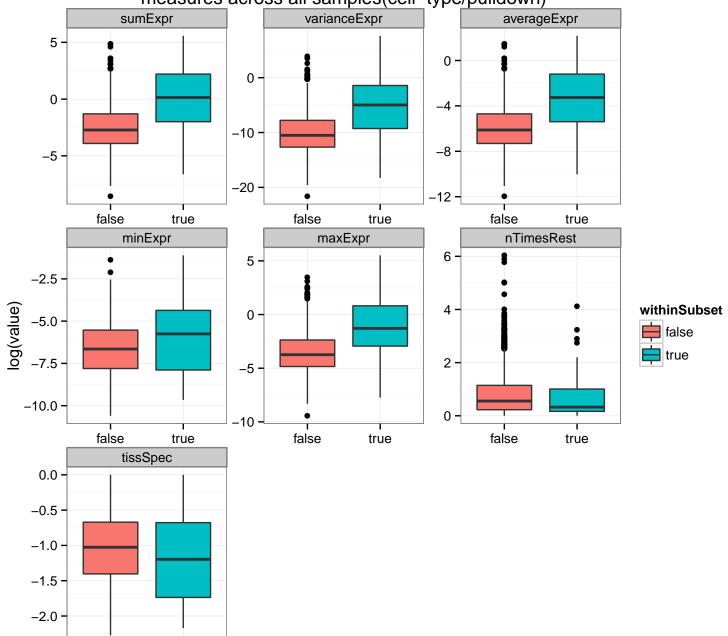lncRNA group = IDRlessthan0_2 ::: data cols = lpa
(51 / 2714) = (func/total lncRNA) ::: biotype = remove_antisense
measures across all samples(cell−type/pulldown)

value of facet label