lncRNA group = IDRlessthan0_2 ::: data cols = lpa
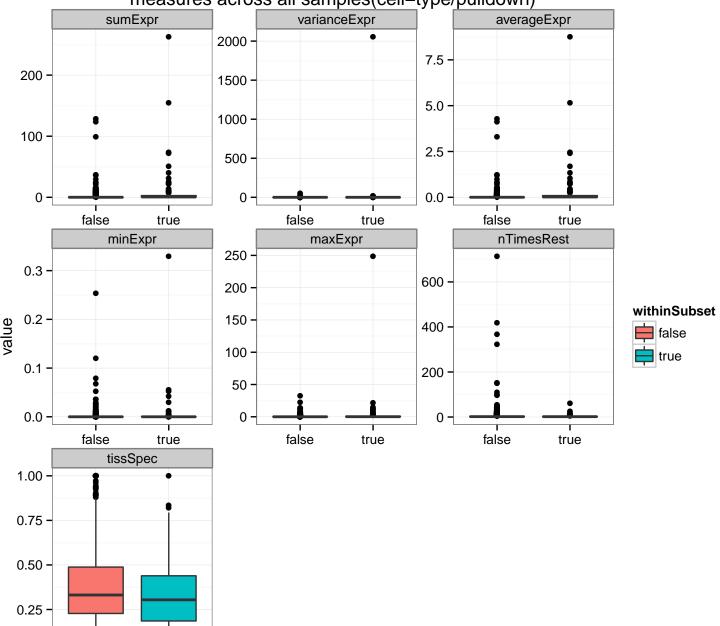(82 / 4574) = (func/total lncRNA) ::: biotype = all_biotypes
measures across all samples(cell−type/pulldown)