lncRNA group = IDRlessthan0_01 ::: data cols = lpa
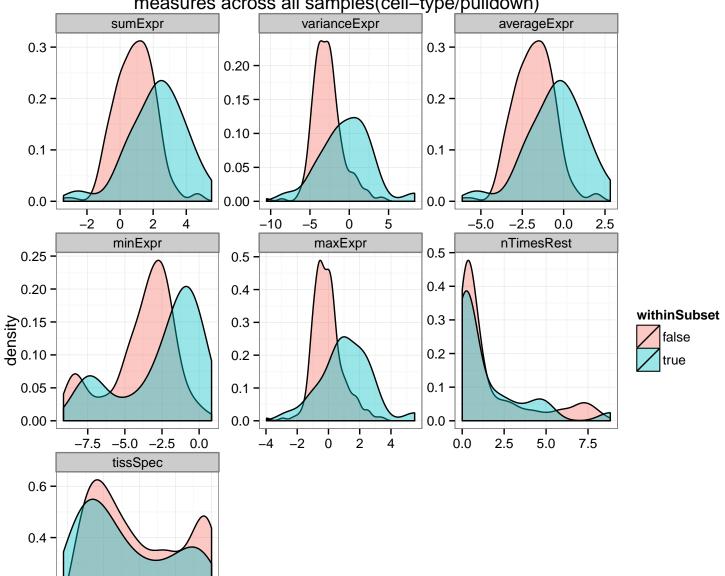(25 / 243) = (func/total lncRNA) ::: biotype = all_biotypes
measures across all samples(cell−type/pulldown)