lncRNA group = IDRlessthan0_01 ::: data cols = lnpa
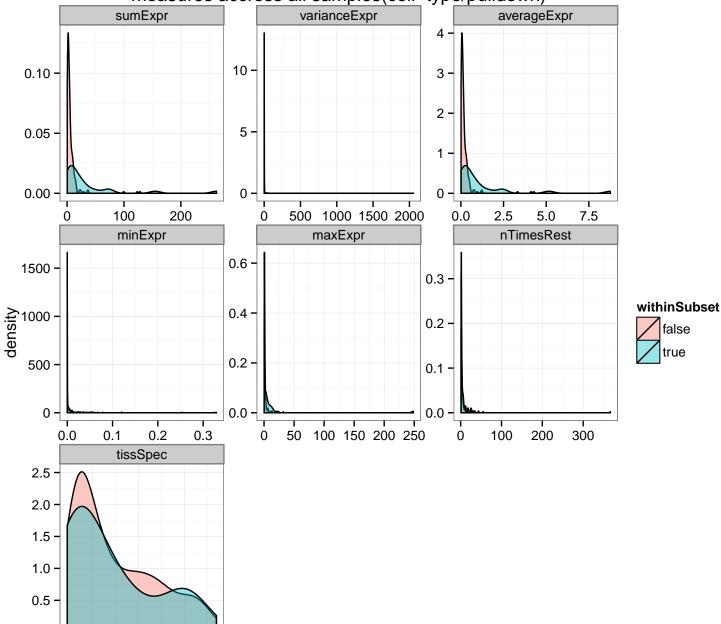(25 / 242) = (func/total lncRNA) ::: biotype = all_biotypes
measures accross all samples(cell−type/pulldown)