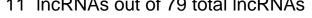
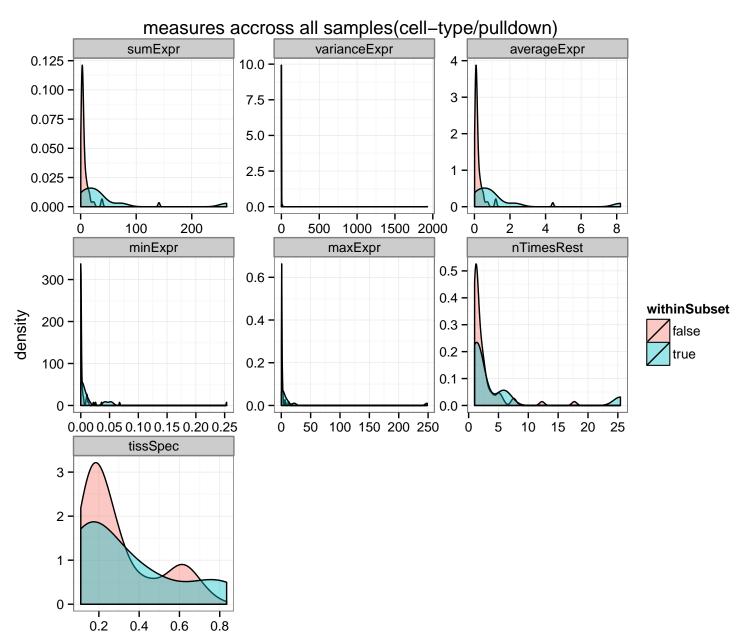
IncRNA group = IDRlessthan0_01 ::: data cols = lpa (11 / 79) = (func/total lncRNA) ::: biotype = processed_transcript subset = IncRNAs out of 79 total IncRNAs





value of facet label