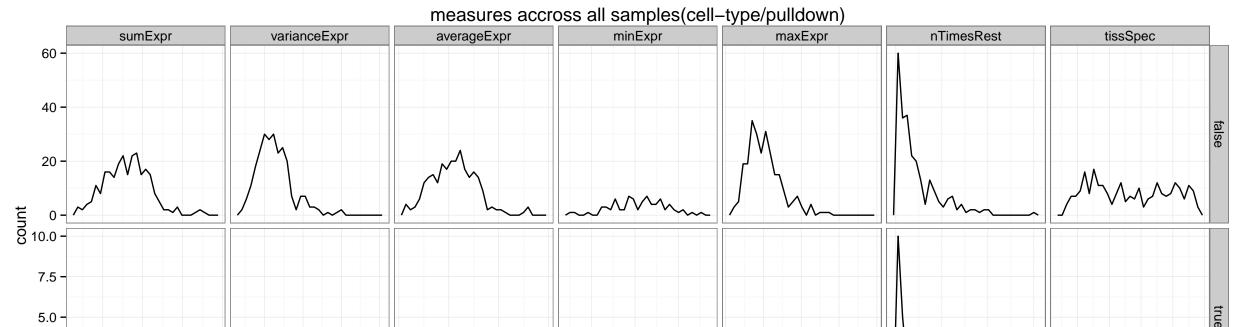lncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns
(25 / 277) = (func/total lncRNA) ::: biotype = all_biotypes
subset =
25 lncRNAs out of 277 total lncRNAs

measures accross all samples(cell−type/pulldown)