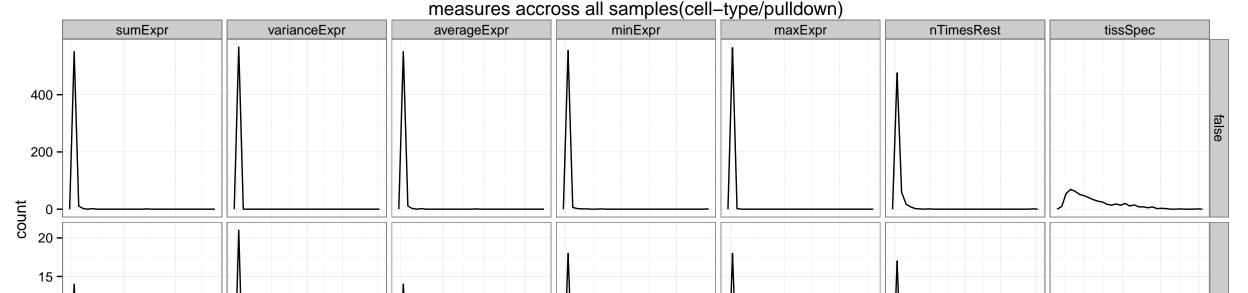
IncRNA group = IDRIessthan0\_1 ::: data cols = Ipa ( 22 / 588 ) = (func/total IncRNA) ::: biotype = processed\_transcript subset = 22 IncRNAs out of 588 total IncRNAs



0.2

value of facet label

0

100

200

25

50

75

100

0.25 0.50 0.75 1.00

10

5

100

200

1000 1500 20000.0

2.5

5.0

7.5

0.0