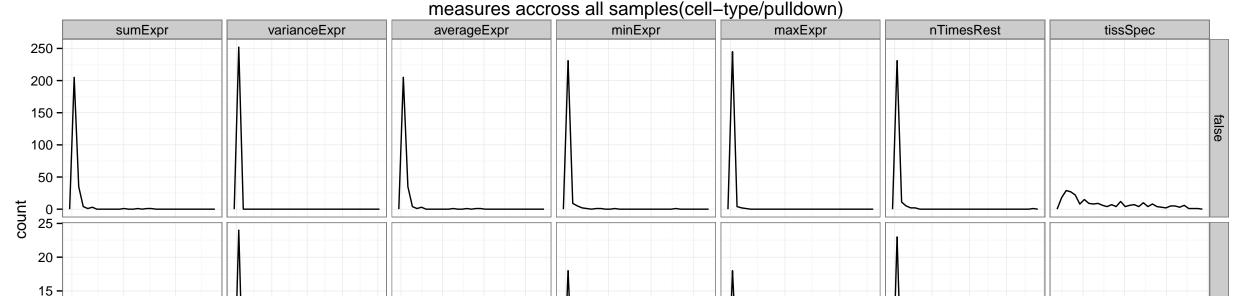
IncRNA group = IDRlessthan0_01 ::: data cols = bothPullDowns (25 / 277) = (func/total IncRNA) ::: biotype = all_biotypes subset = 25 IncRNAs out of 277 total IncRNAs



10

5

200

1000 1500 20000.0

2.5

5.0

7.5

0.0

0.1

0.2

value of facet label

0.3

0

100

200

100

200

300

400

0.25