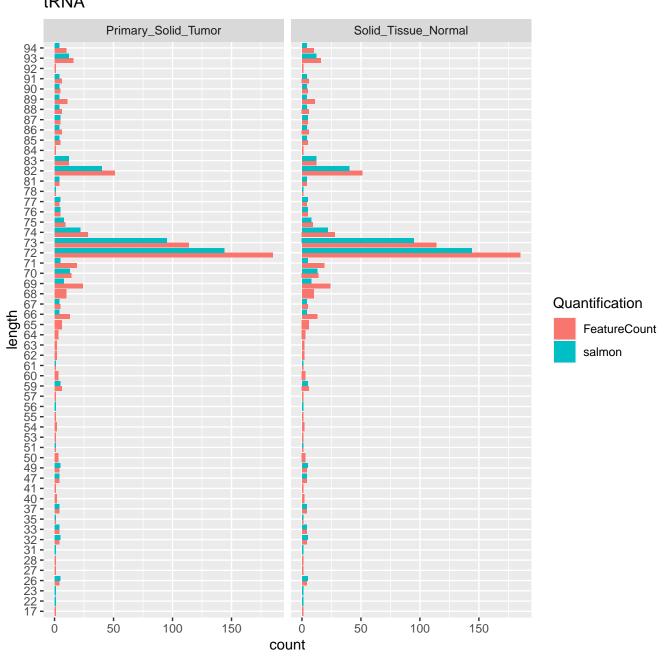
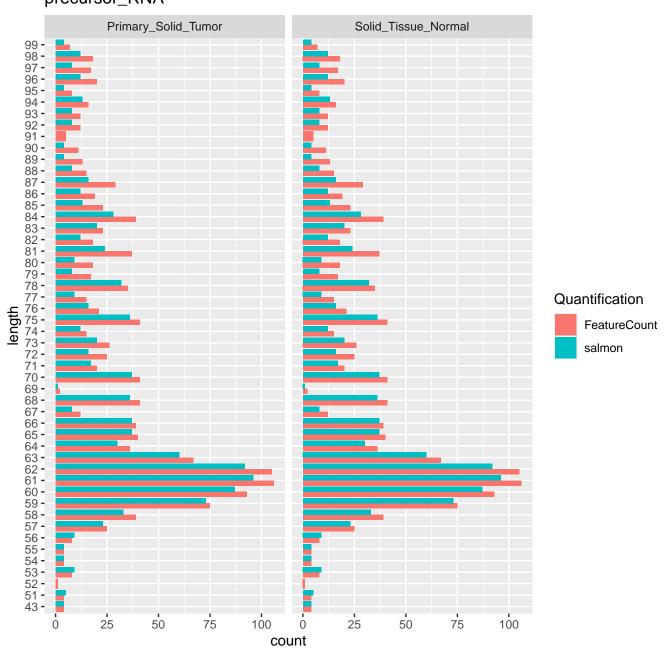
tRNA

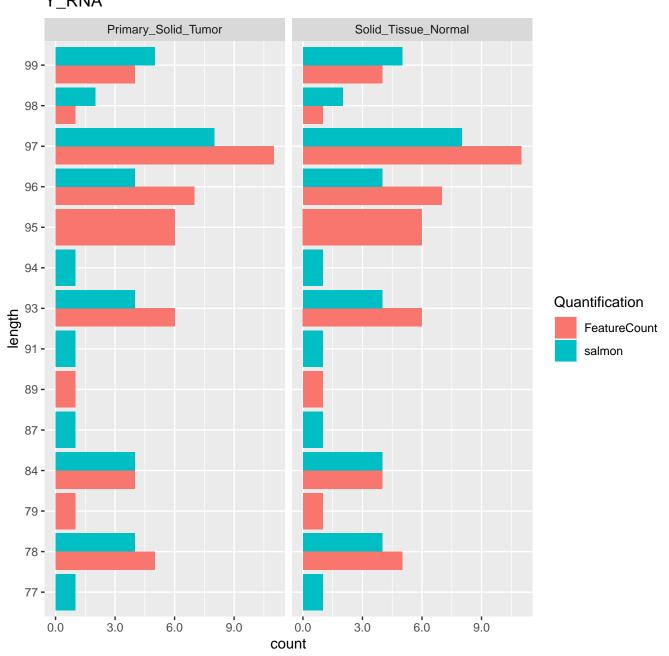


precursor_RNA



miRNA Primary_Solid_Tumor Solid_Tissue_Normal 27 -26 -25 **-**24 -23 -Quantification length 22 -FeatureCount salmon 21 -20 -19 -18 -17 -100 300 400 100 200 300 200 400 count

Y_RNA



snRNA Primary_Solid_Tumor Solid_Tissue_Normal 95 -70 -68 **-**67 -66 -Quantification length FeatureCount salmon 62 **-**61 -49 -43 -

4.0 0.0

count

1.0

2.0

3.0

4.0

35 **-**

0.0

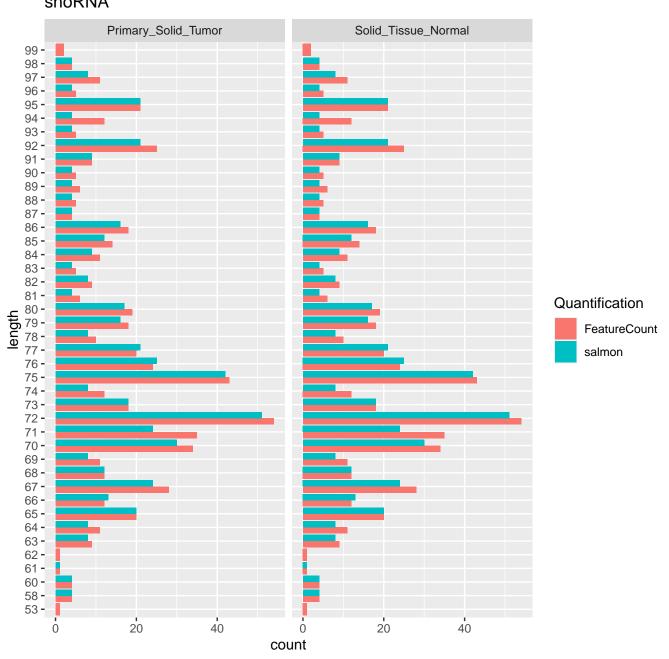
1.0

2.0

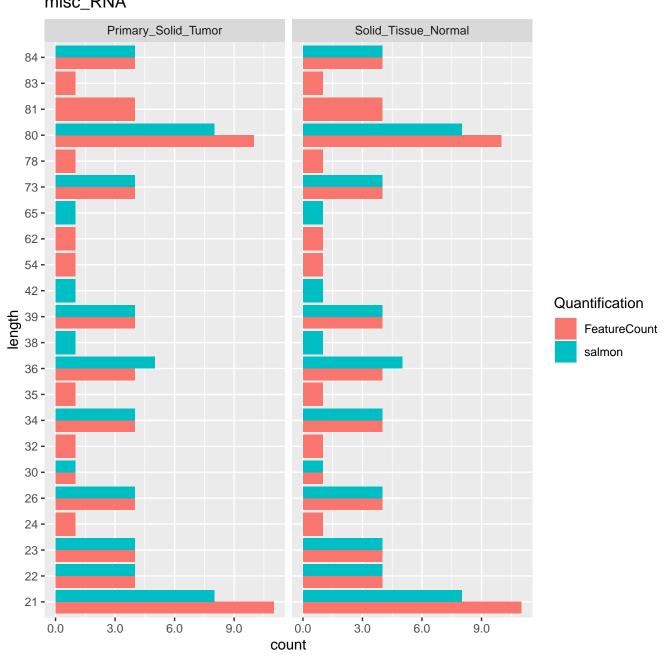
3.0

piRNA Primary_Solid_Tumor Solid_Tissue_Normal 32 -31 -30 -Quantification length FeatureCount salmon 28 -27 -26 -20 20 40 40 60 60 count

snoRNA

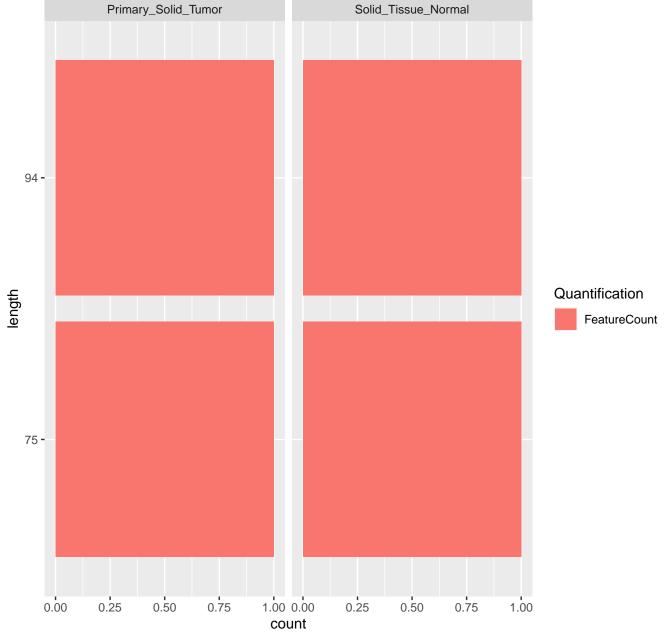


misc_RNA



other Primary_Solid_Tumor Solid_Tissue_Normal 81 -Quantification length FeatureCount salmon 25 -0.0 1.0 4.0 0.0 1.0 2.0 3.0 3.0 4.0 2.0 count

IncRNA



scRNA Primary_Solid_Tumor Solid_Tissue_Normal 93 -Quantification length FeatureCount salmon 71 -

0.25

1.00 0.00

count

0.00

0.25

0.75

0.50

0.50

0.75

1.00

rRNA Primary_Solid_Tumor Solid_Tissue_Normal 98 -95 -91 -Quantification length • 06 FeatureCount salmon 78 **-**62 **-**

4.0 0.0

count

3.0

2.0

1.0

2.0

3.0

4.0

51 **-**

0.0

1.0