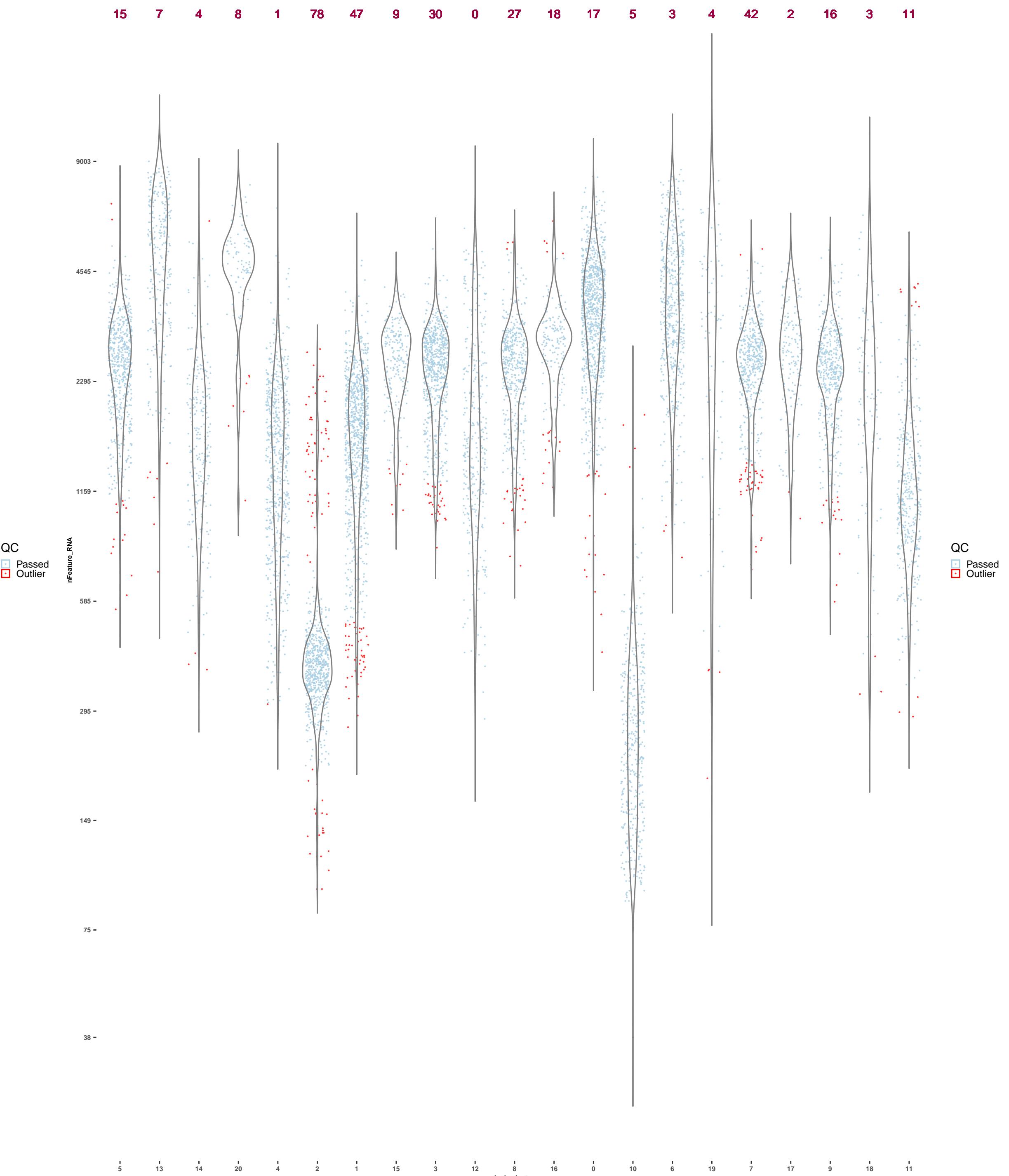
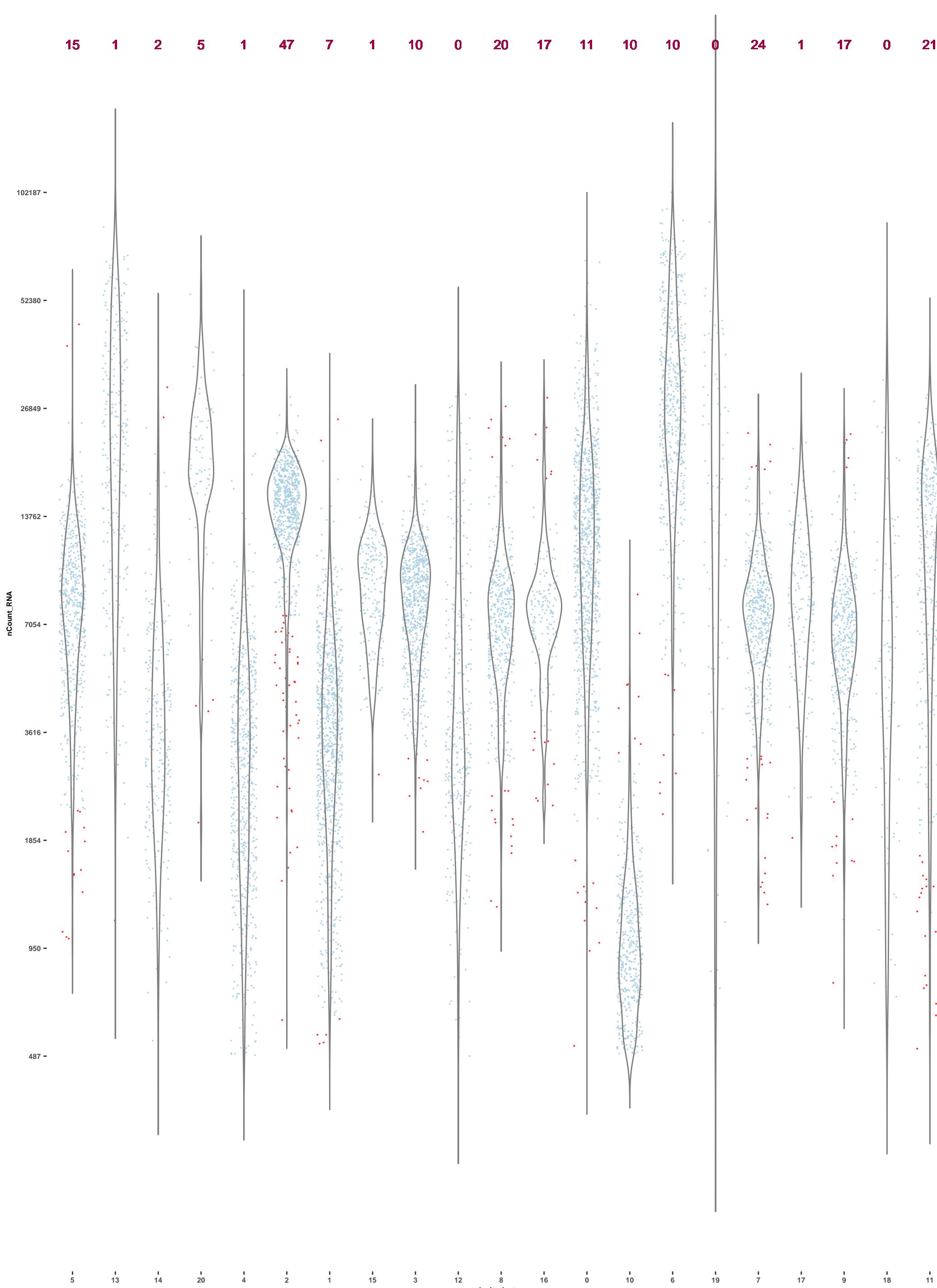


S1_D1_cell(log+1): -3MAD > outliers > +3MAD

S1_D1_cell(log+1): -3MAD > outliers > +3MAD



QC

Passed

Outlier

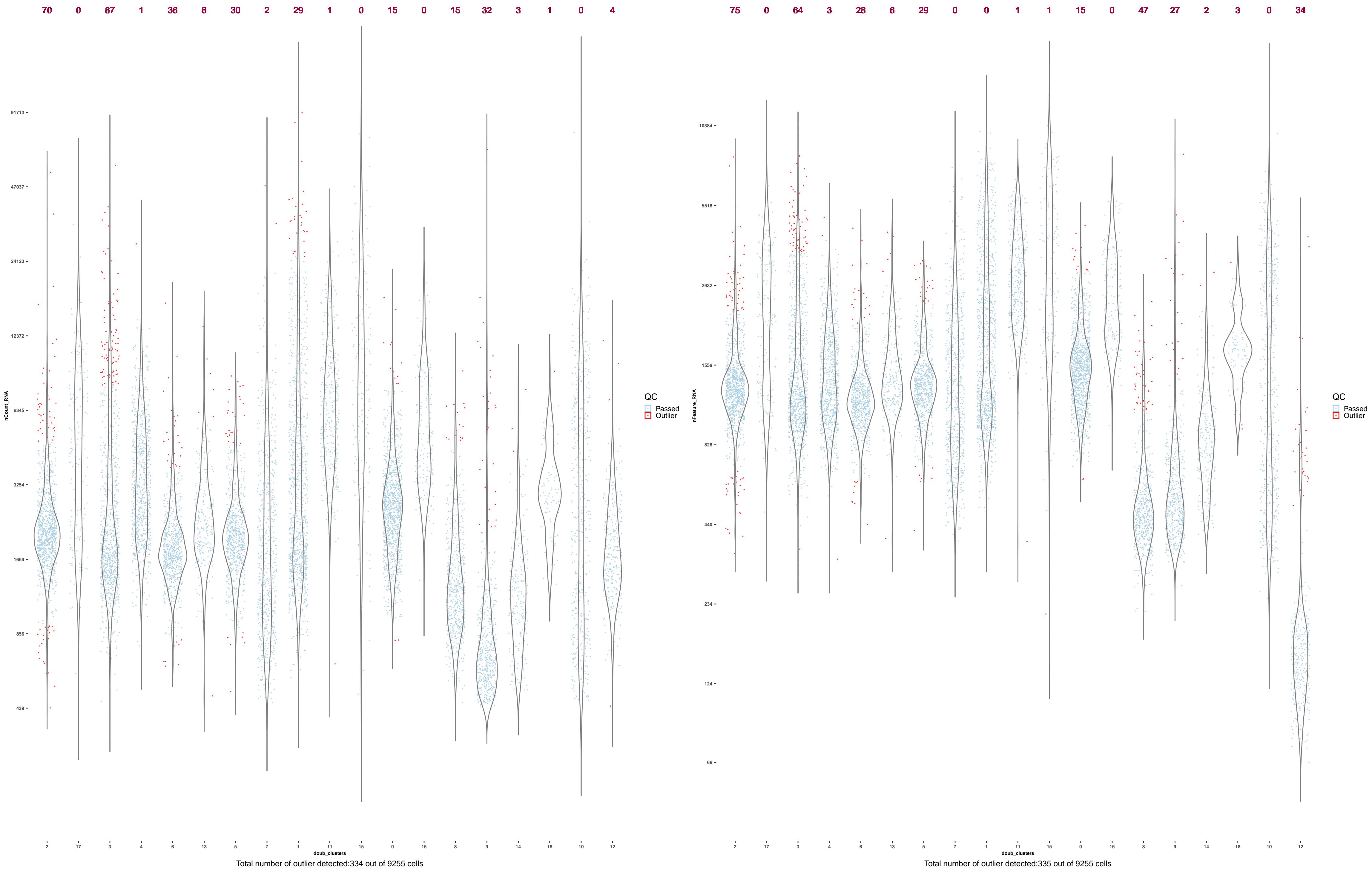
QC

Passed

Outlier

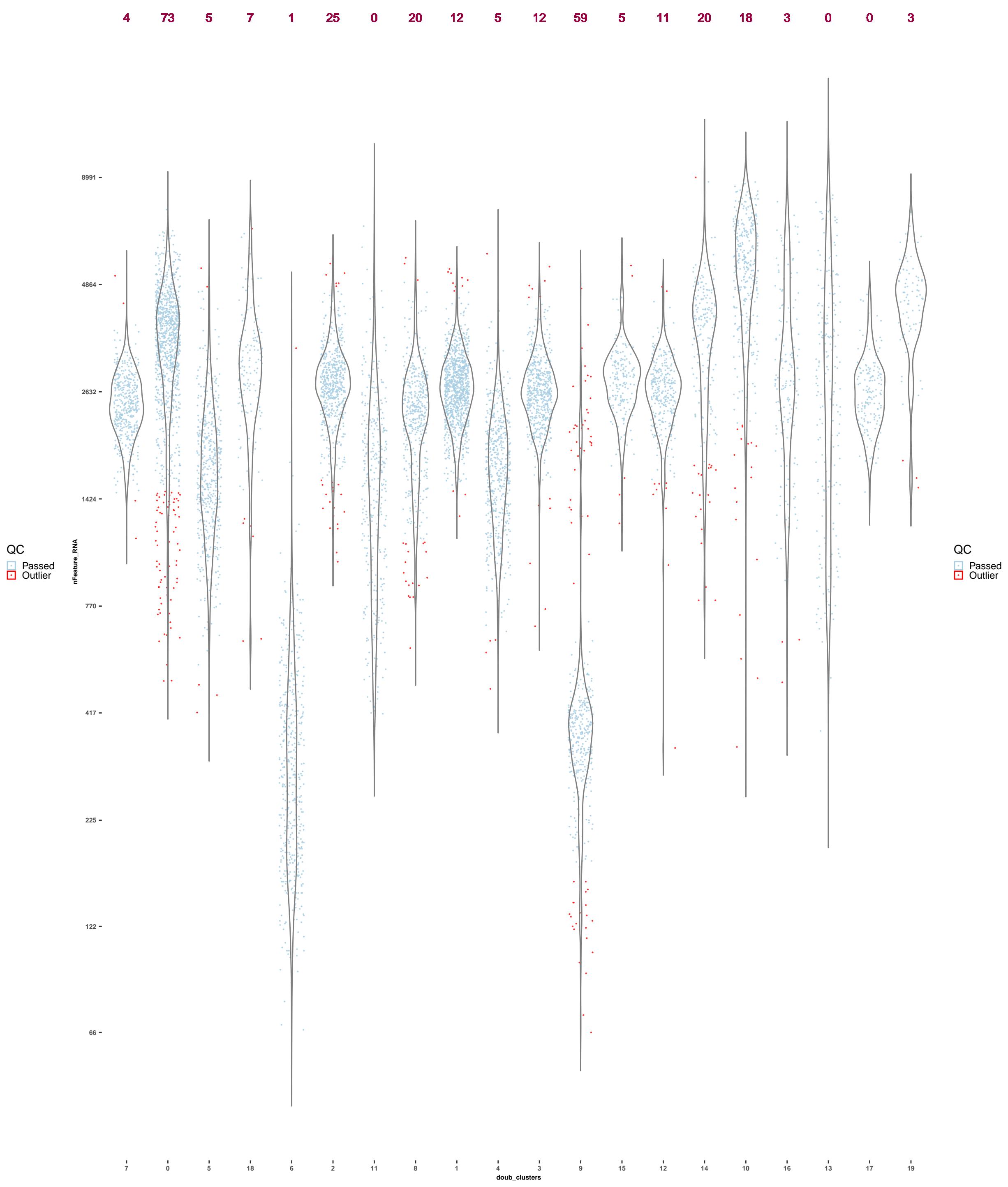
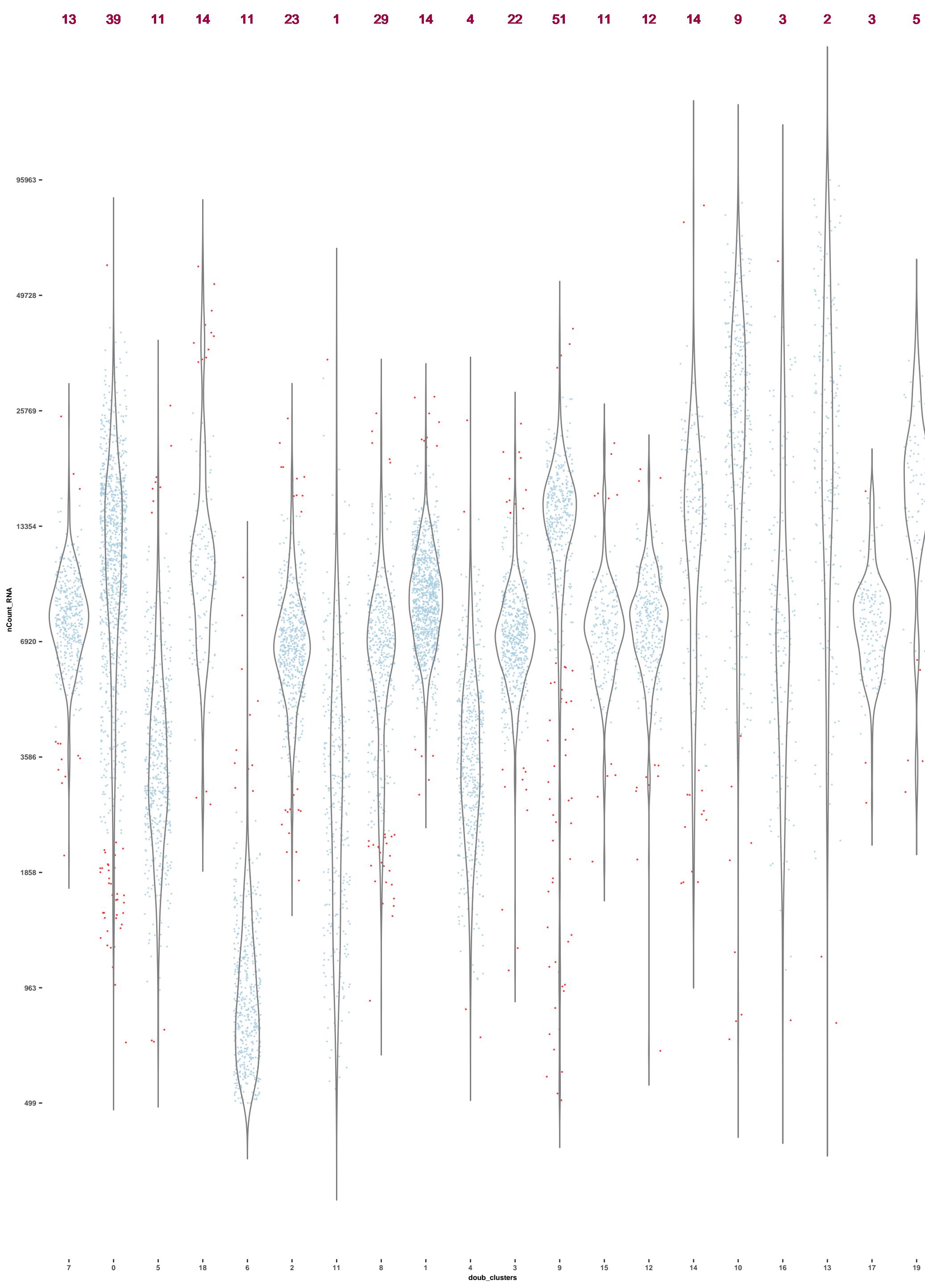
S1_D1_Nuclei(log+1): -3MAD > outliers > +3MAD

S1_D1_Nuclei(log+1): -3MAD > outliers > +3MAD



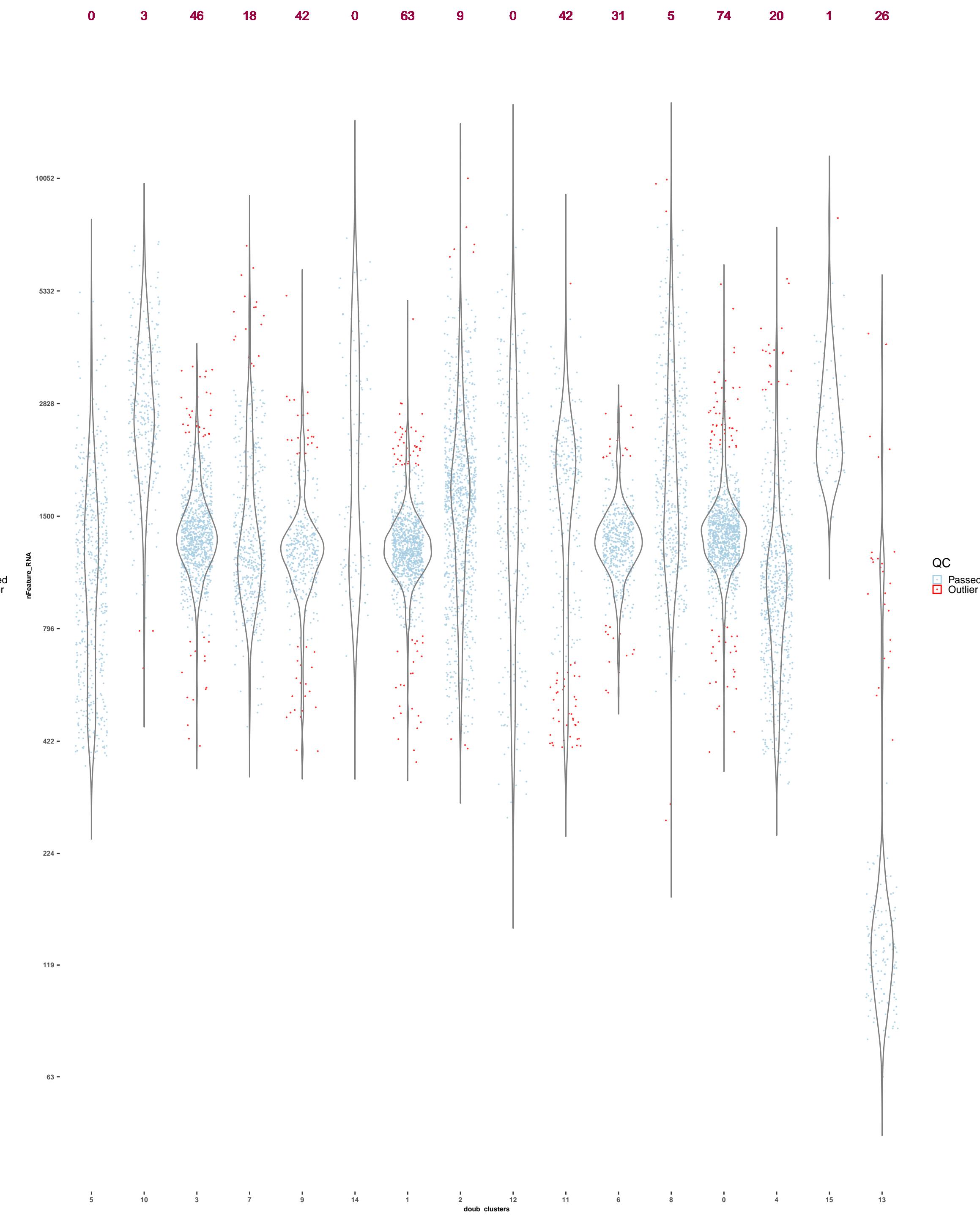
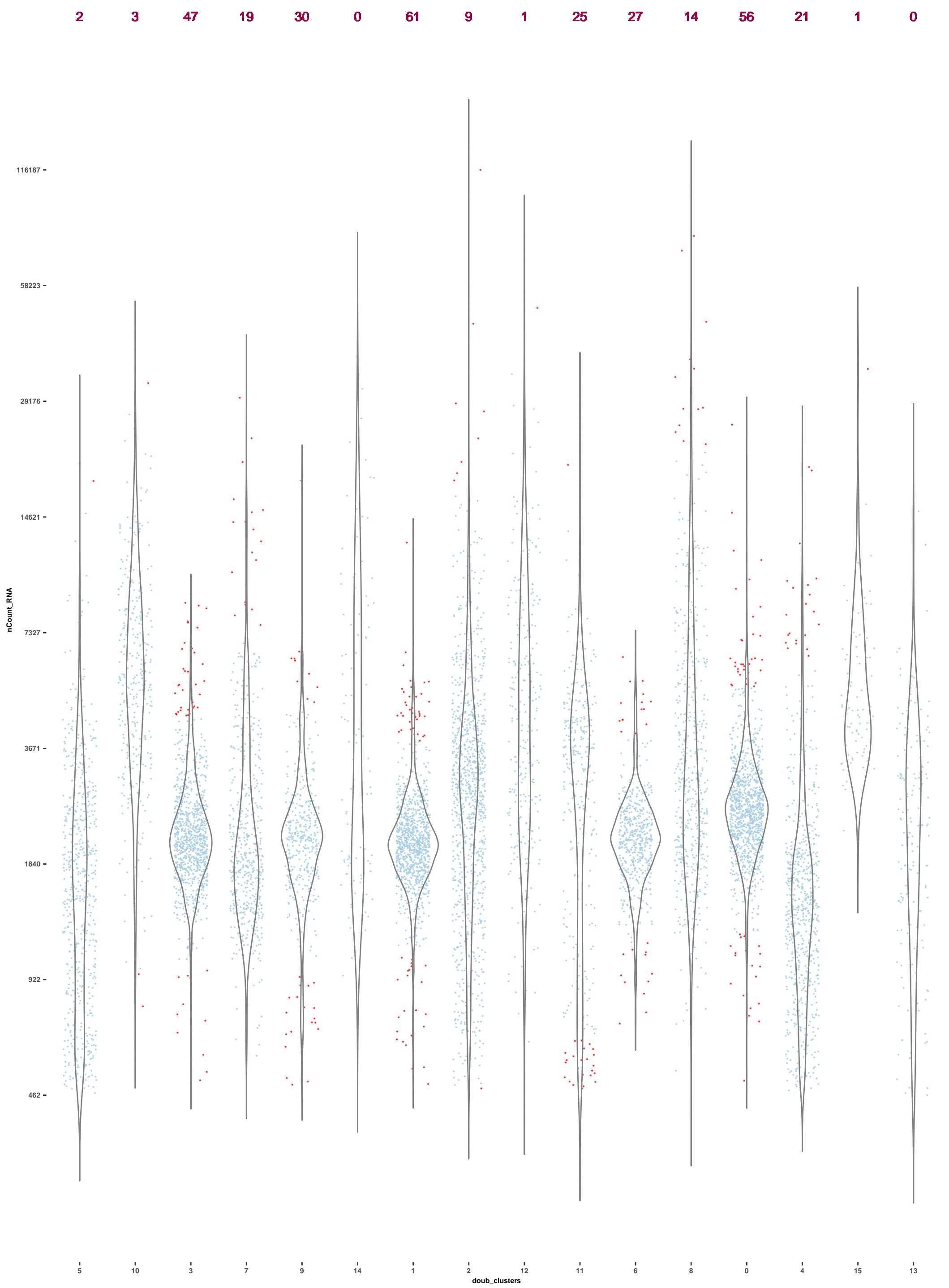
S1_D2_cell(log+1): -3MAD > outliers > +3MAD

S1_D2_cell(log+1): -3MAD > outliers > +3MAD



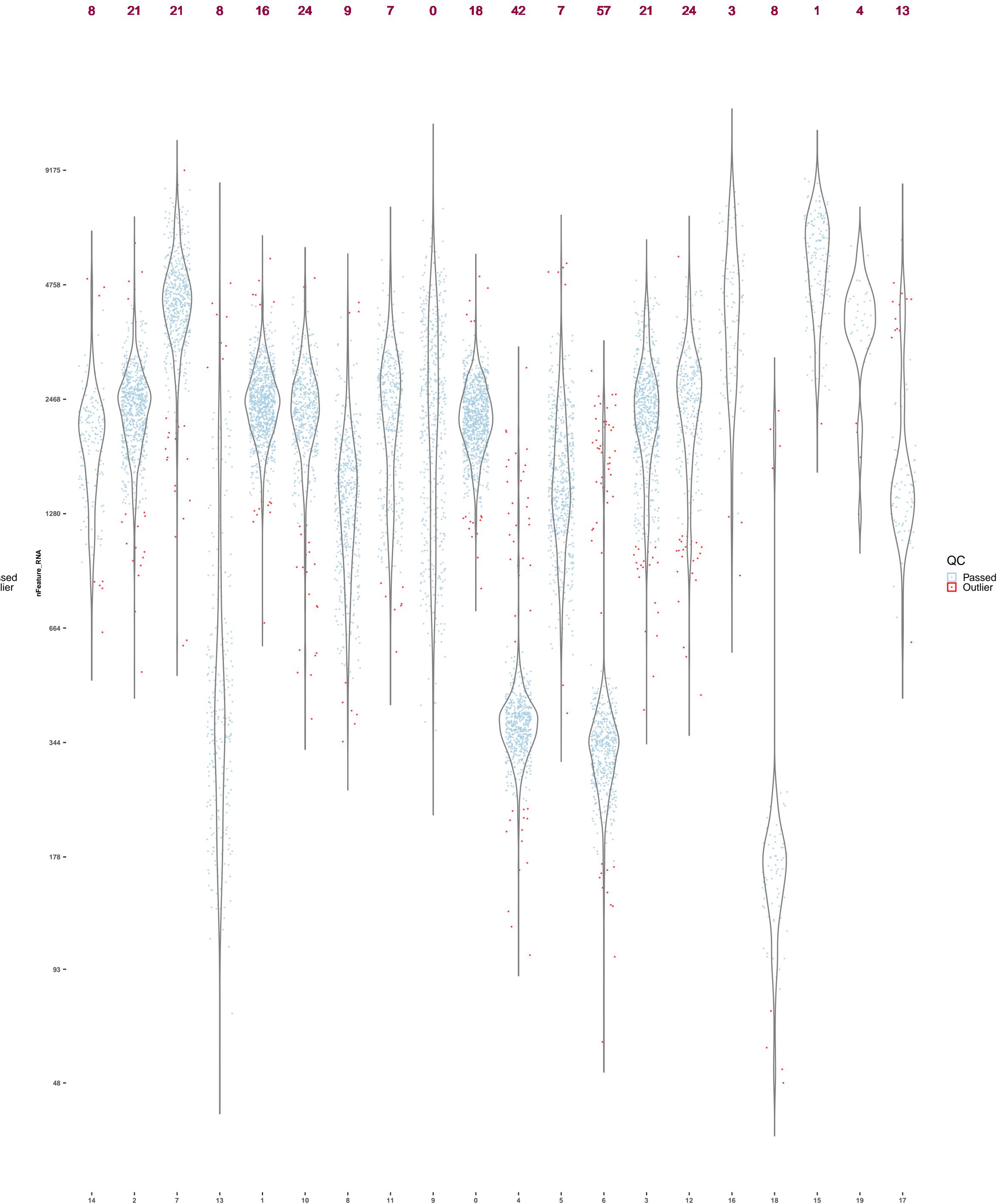
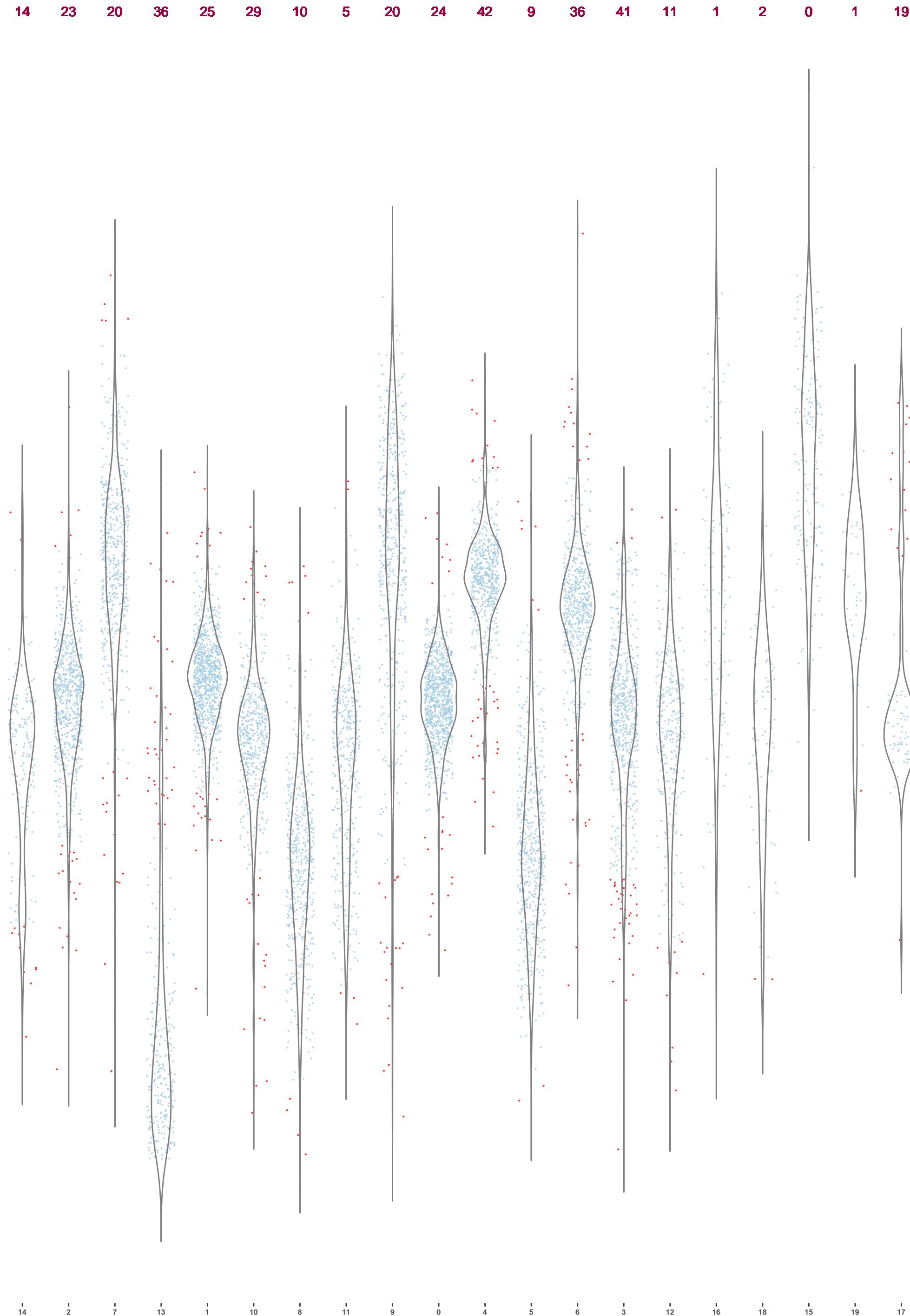
S1_D2_Nuclei(log+1): -3MAD > outliers > +3MAD

S1_D2_Nuclei(log+1): -3MAD > outliers > +3MAD



S1_D3_cell(log+1): -3MAD > outliers > +3MAD

S1_D3_cell(log+1): -3MAD > outliers > +3MAD

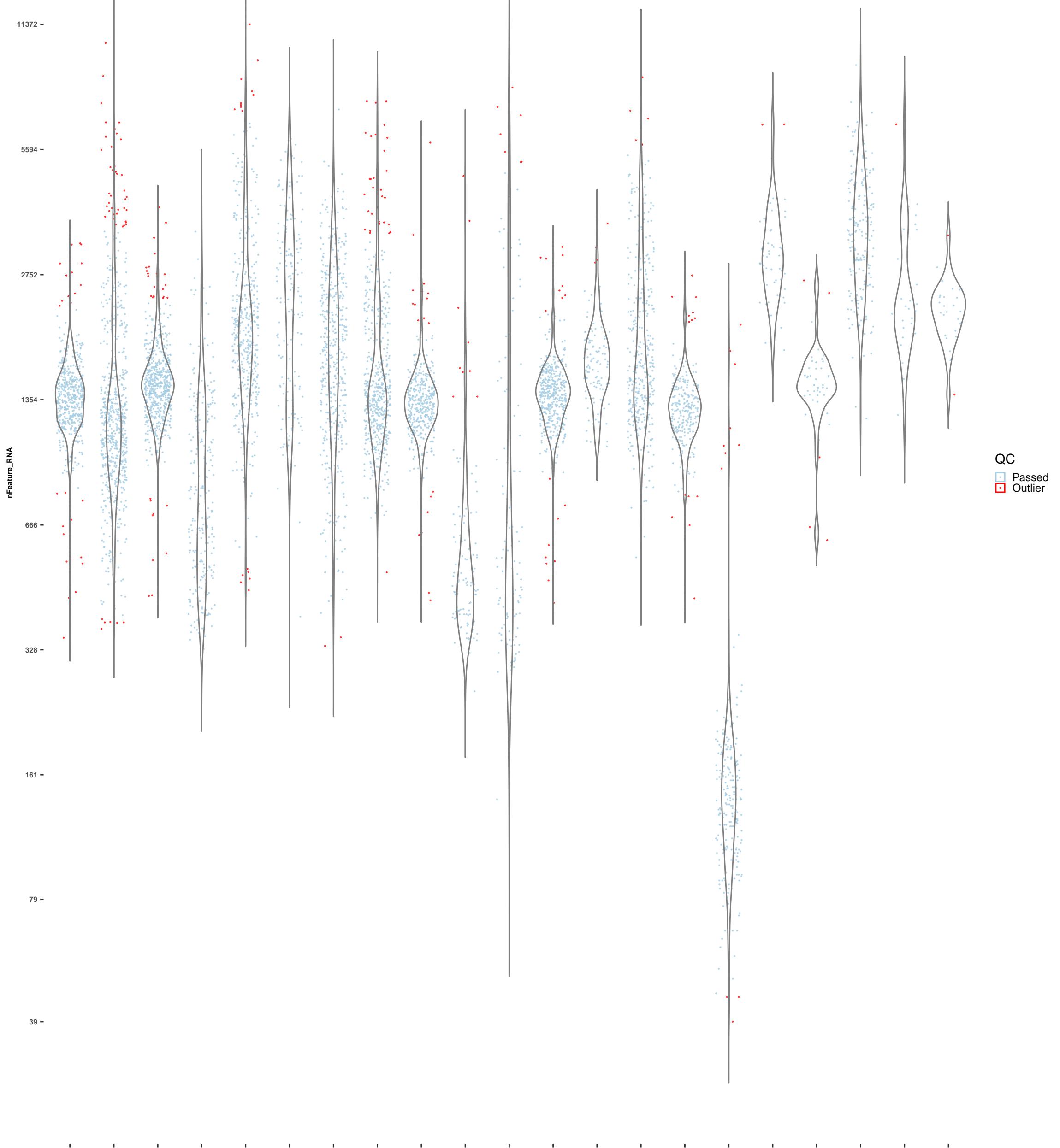
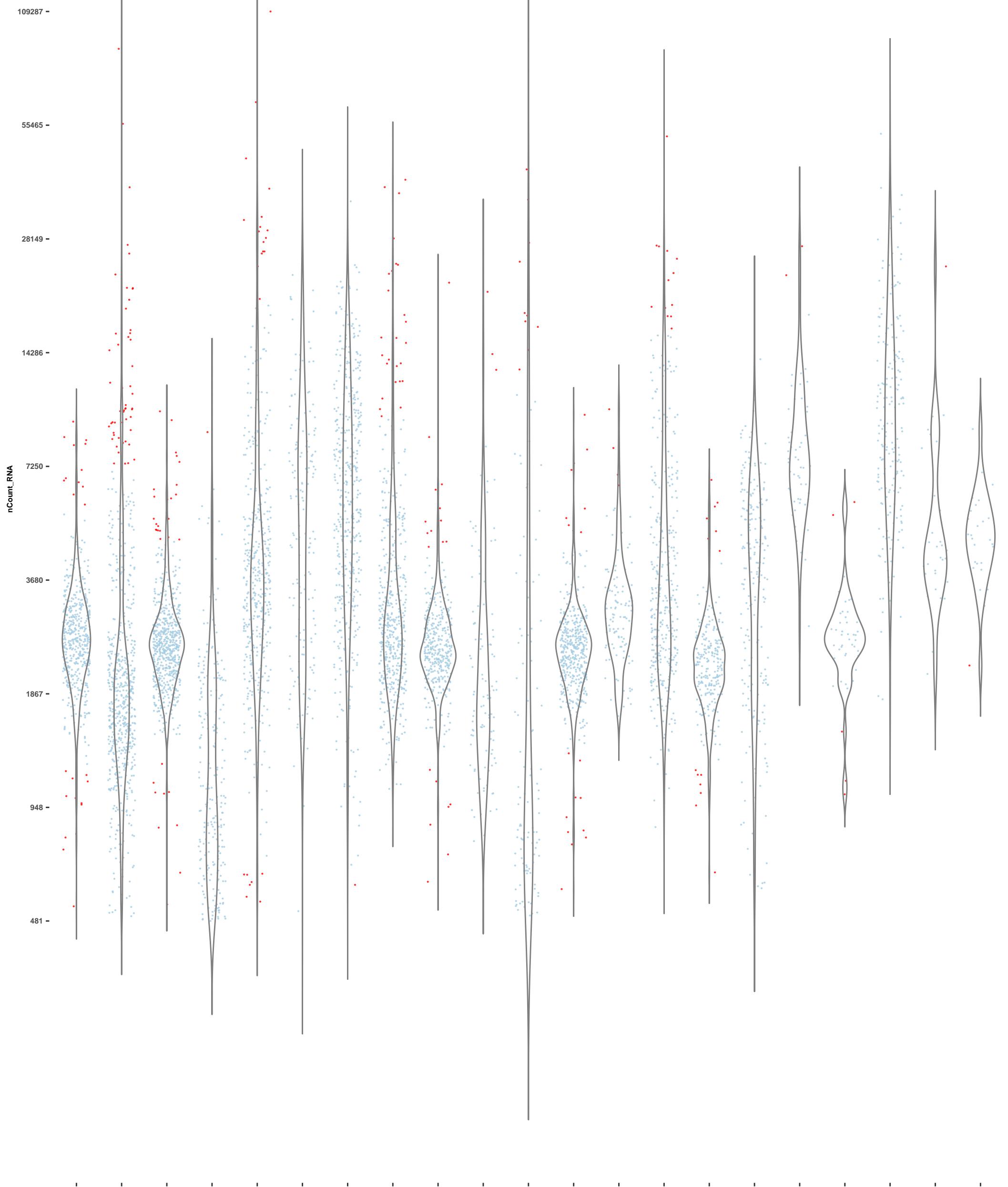


S1_D3_Nuclei(log+1): -3MAD > outliers > +3MAD

S1_D3_Nuclei(log+1): -3MAD > outliers > +3MAD

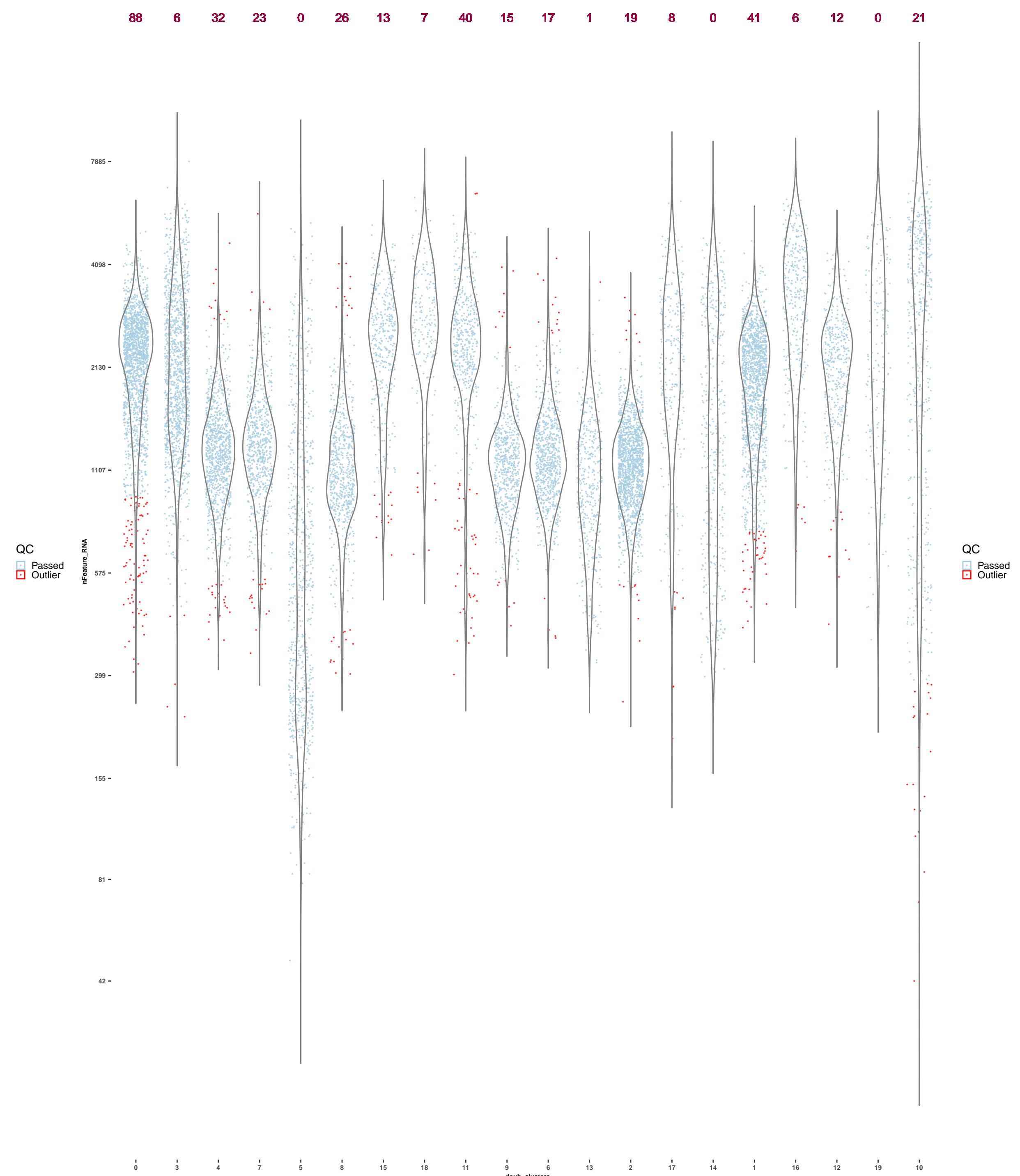
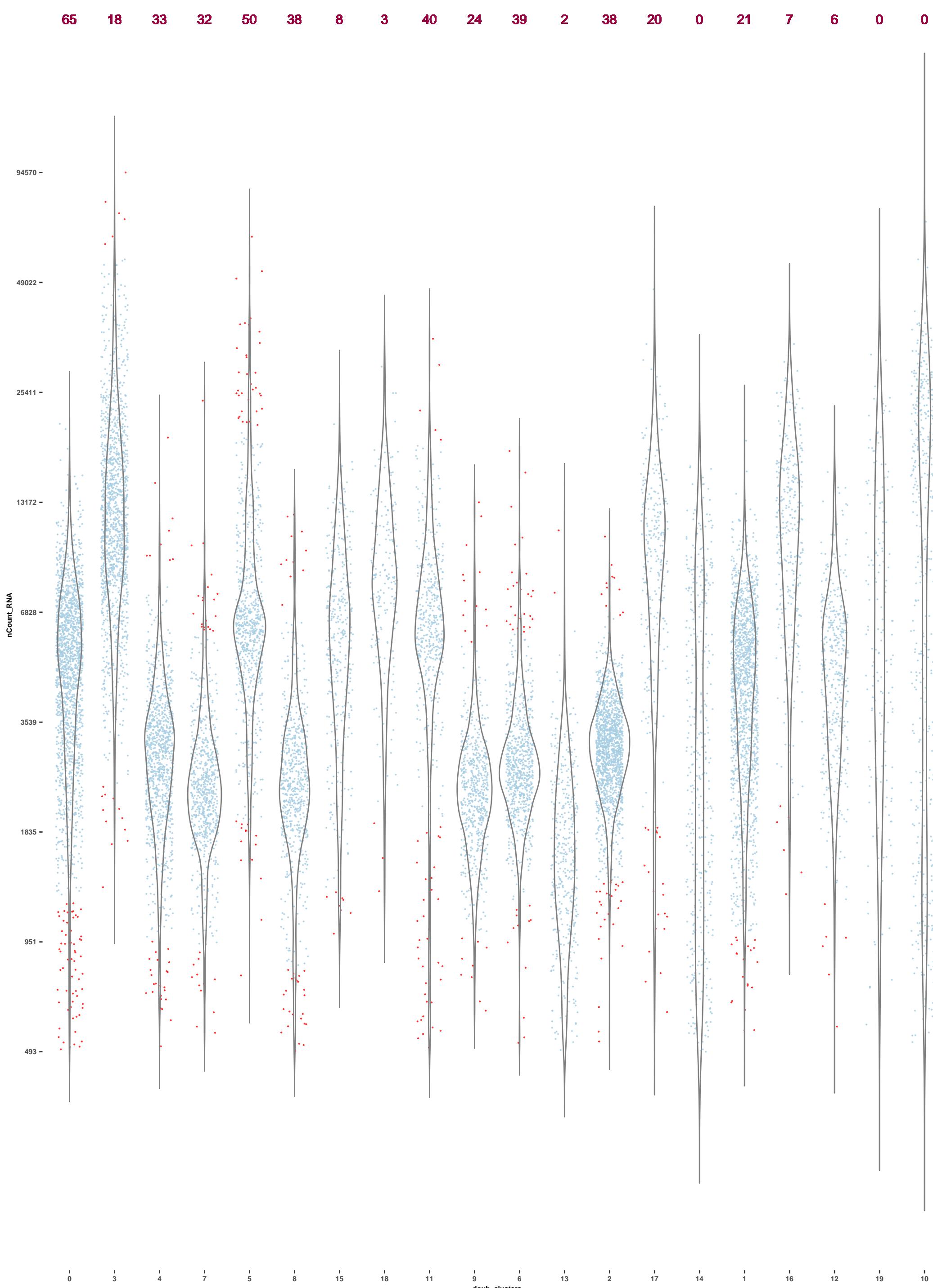
24 52 32 1 23 0 1 27 19 3 10 19 4 13 14 0 2 5 0 1 1

27 46 31 0 16 0 2 34 20 9 7 19 4 6 14 12 2 5 0 1 2



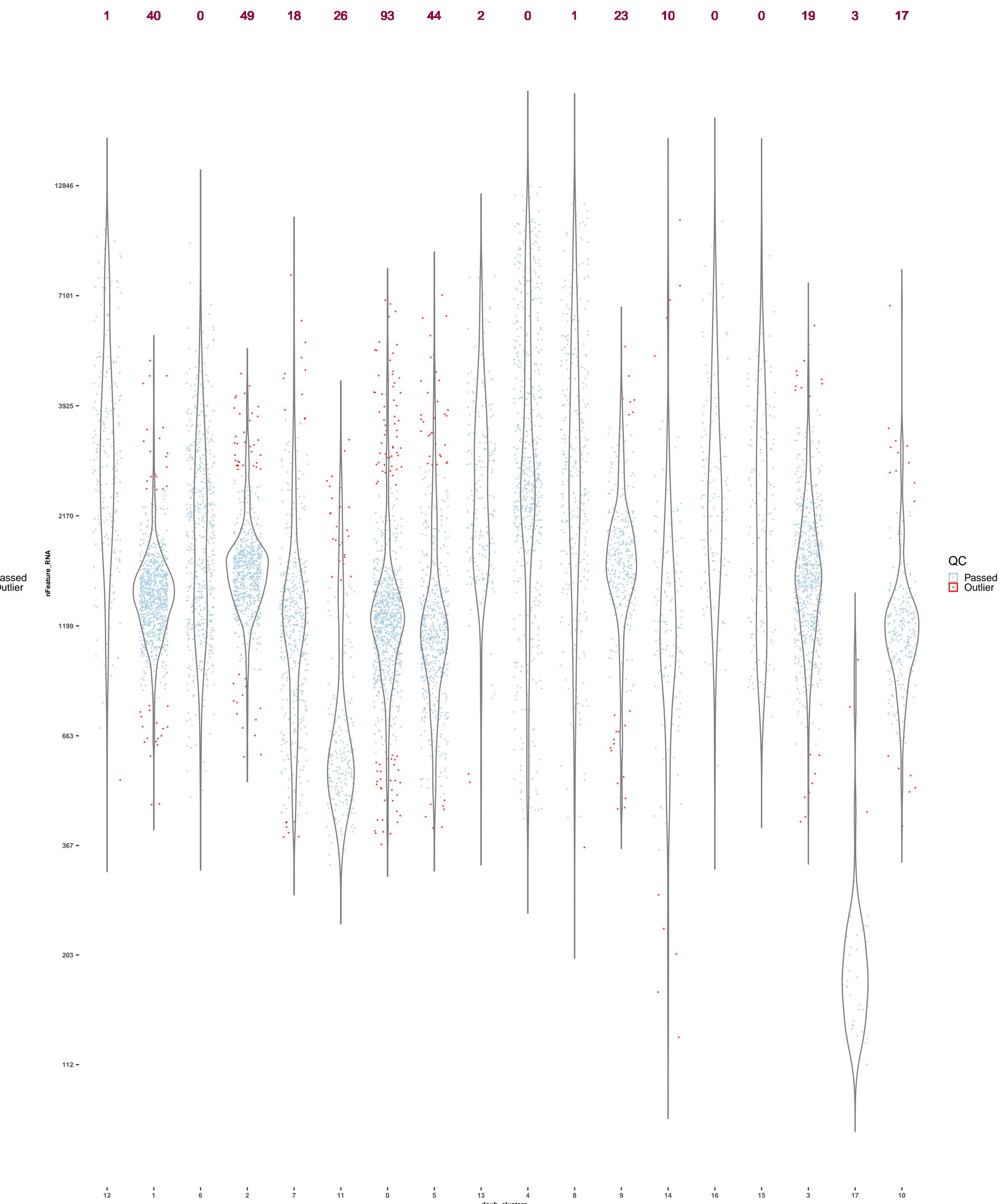
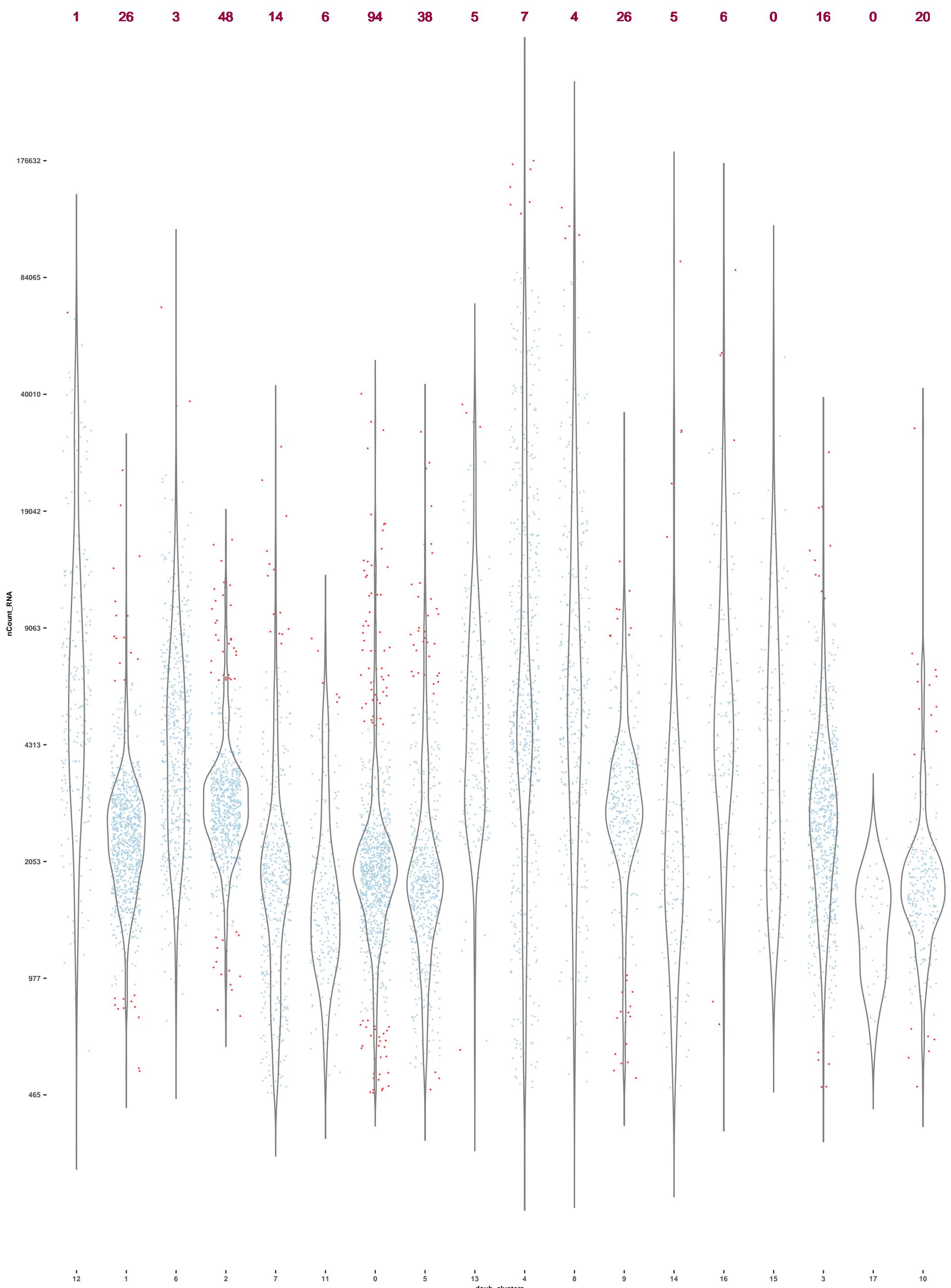
S2_D1_cell(log+1): -3MAD > outliers > +3MAD

S2_D1_cell(log+1): -3MAD > outliers > +3MAD



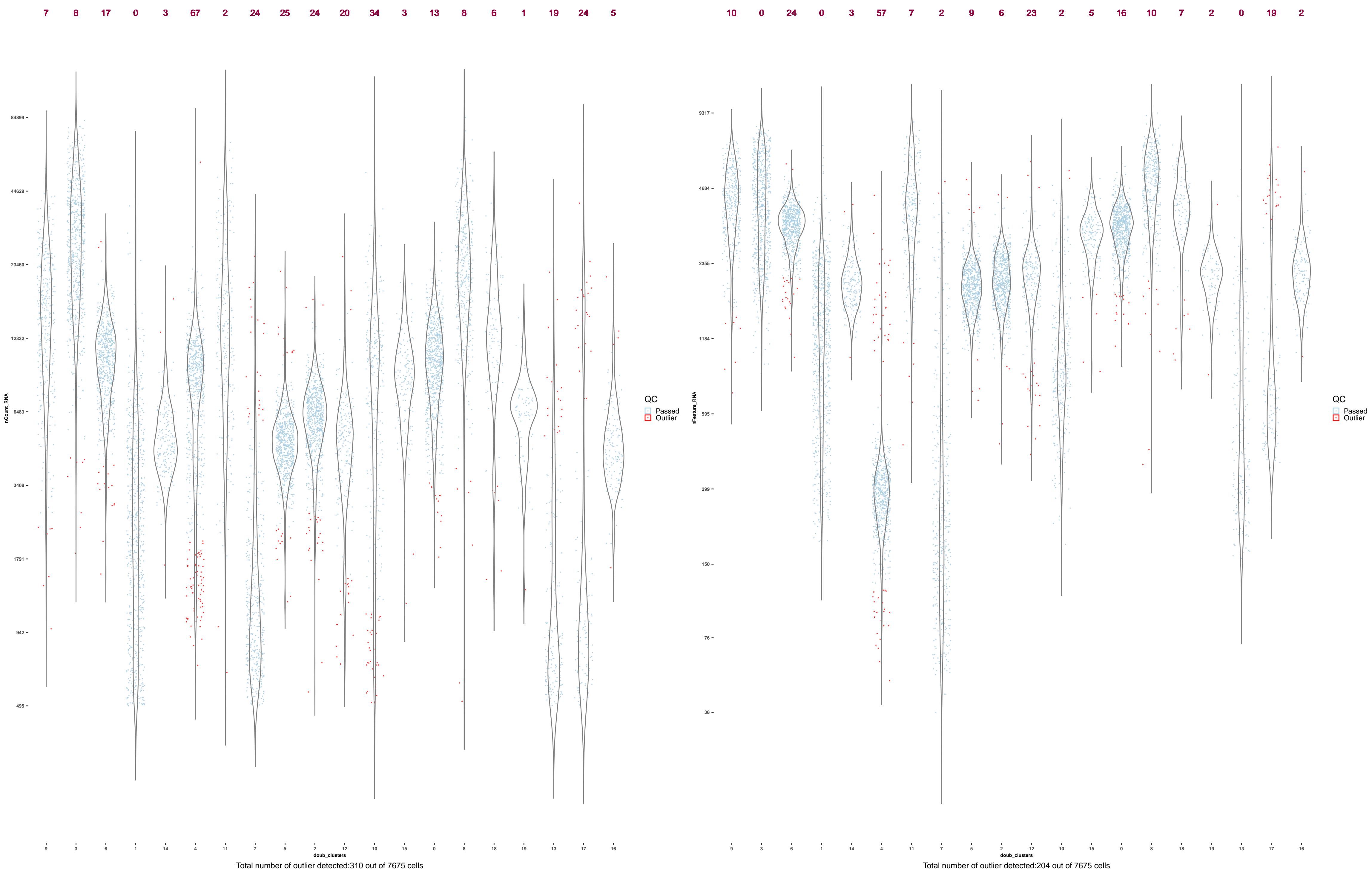
S2_D1_Nuclei(log+1): -3MAD > outliers > +3MAD

S2_D1_Nuclei(log+1): -3MAD > outliers > +3MAD



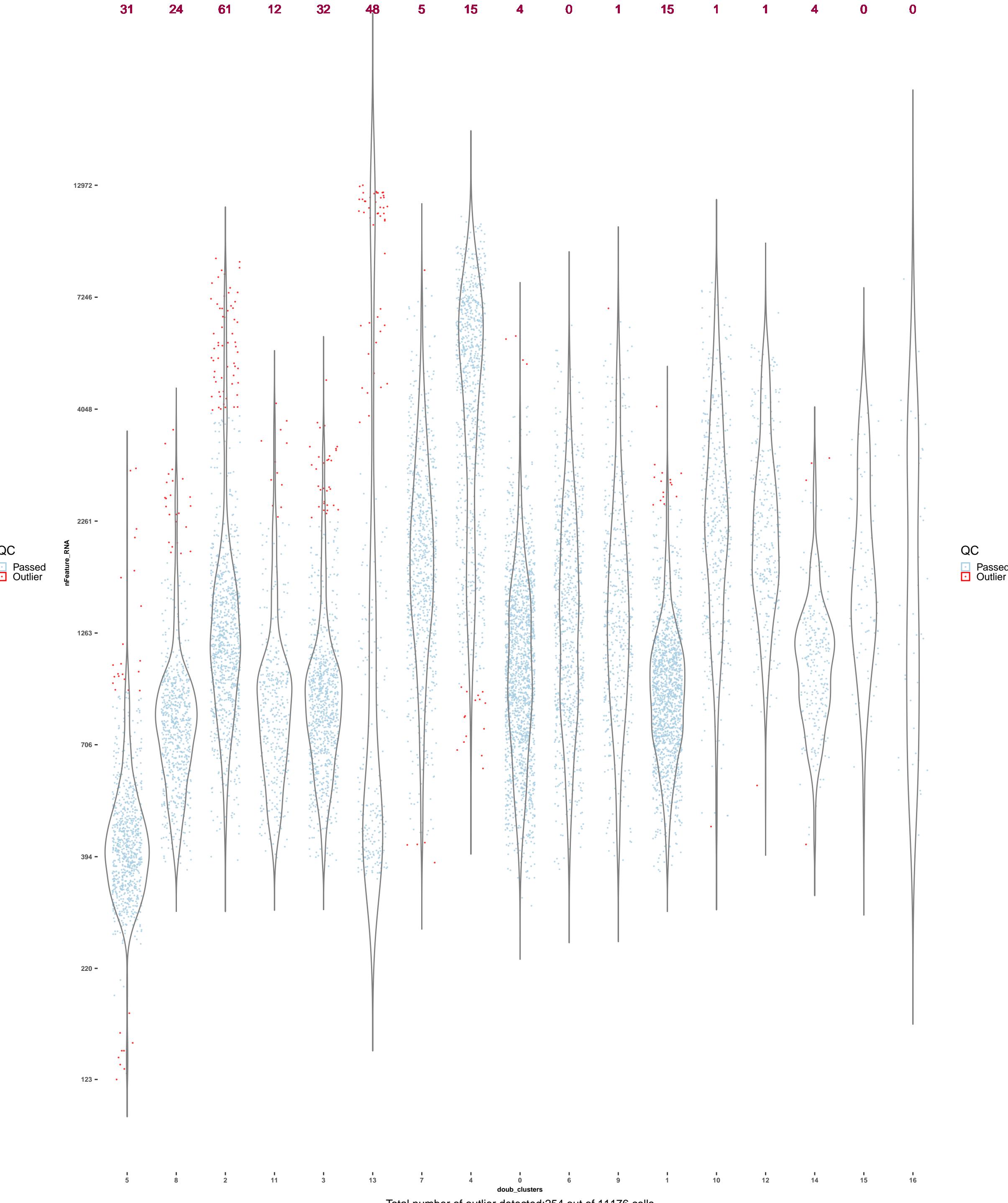
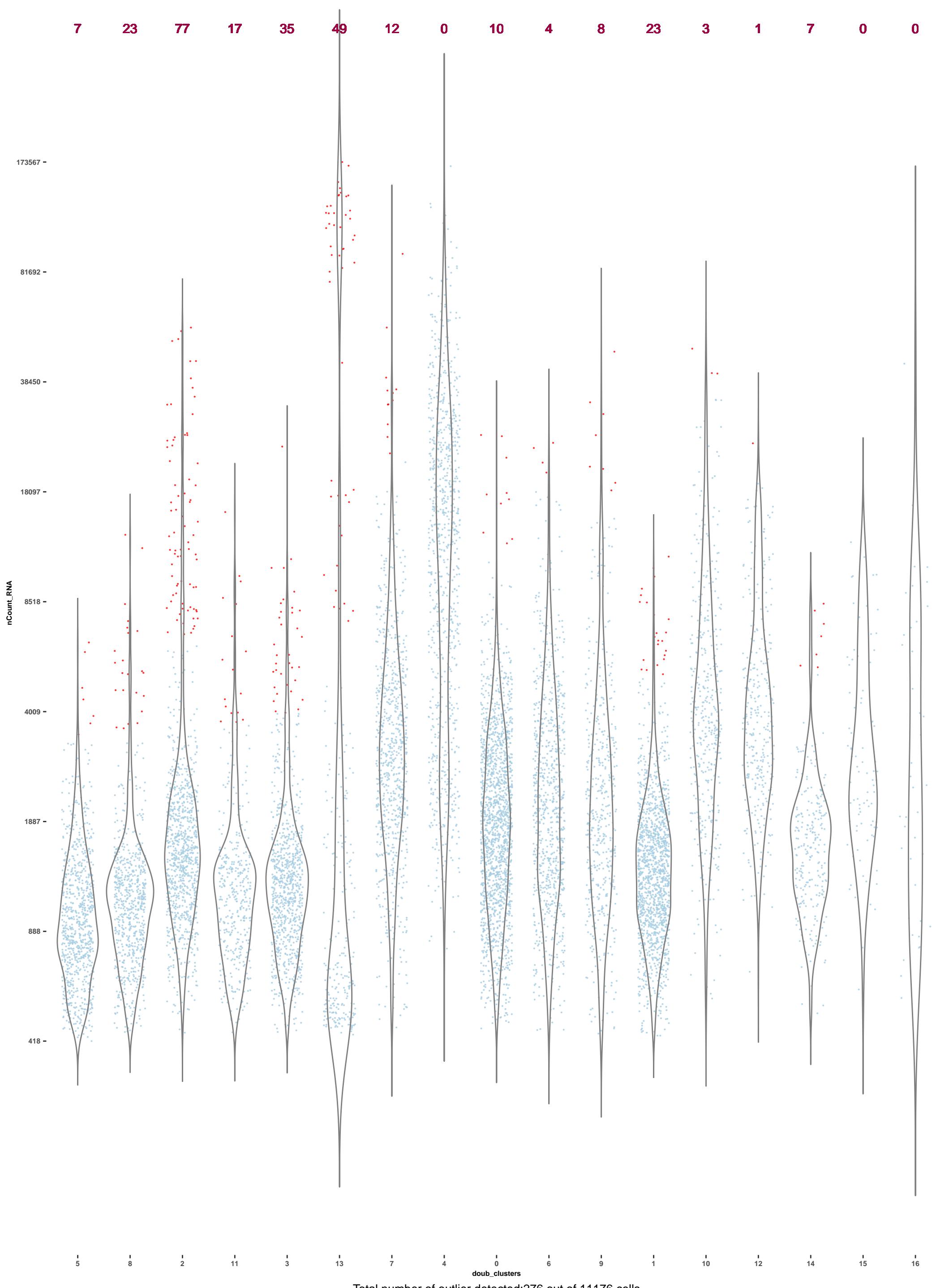
S2_D4_cell(log+1): -3MAD > outliers > +3MAD

S2_D4_cell(log+1): -3MAD > outliers > +3MAD



S2_D4_Nuclei(log+1): -3MAD > outliers > +3MAD

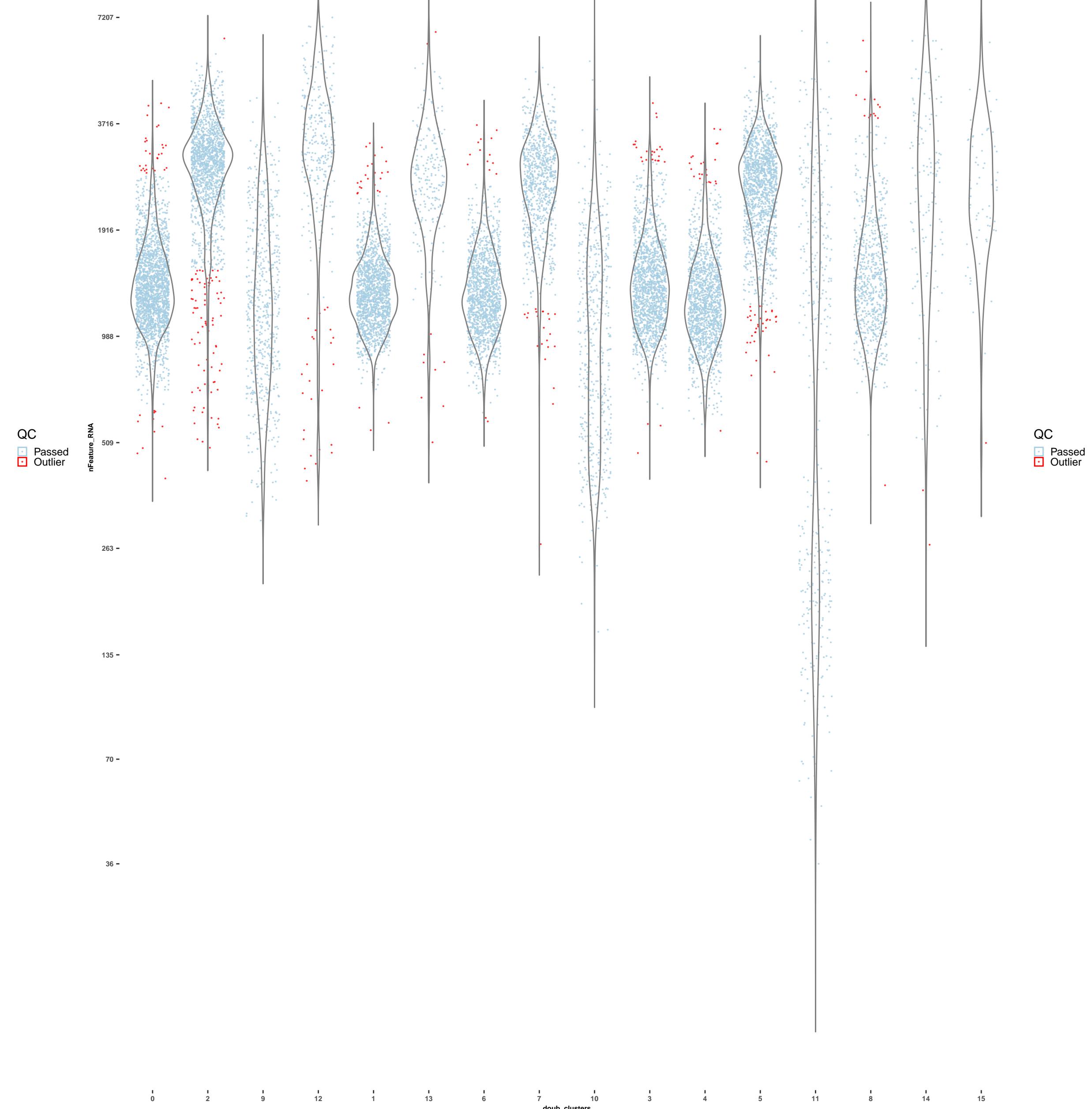
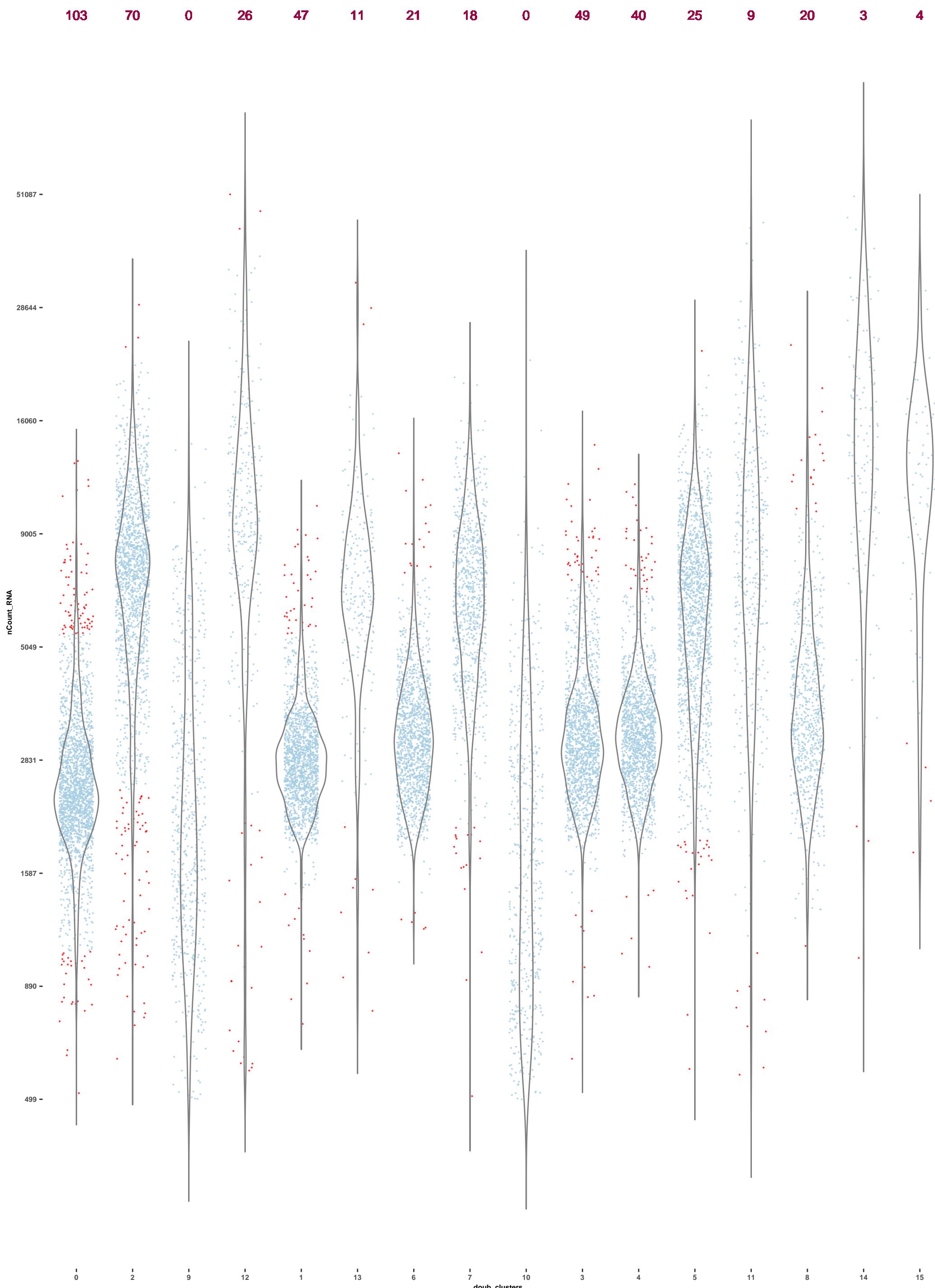
S2_D4_Nuclei(log+1): -3MAD > outliers > +3MAD



QC
Passed
Outlier

S2_D5_cell(log+1): -3MAD > outliers > +3MAD

S2_D5_cell(log+1): -3MAD > outliers > +3MAD



QC

Passed

Outlier

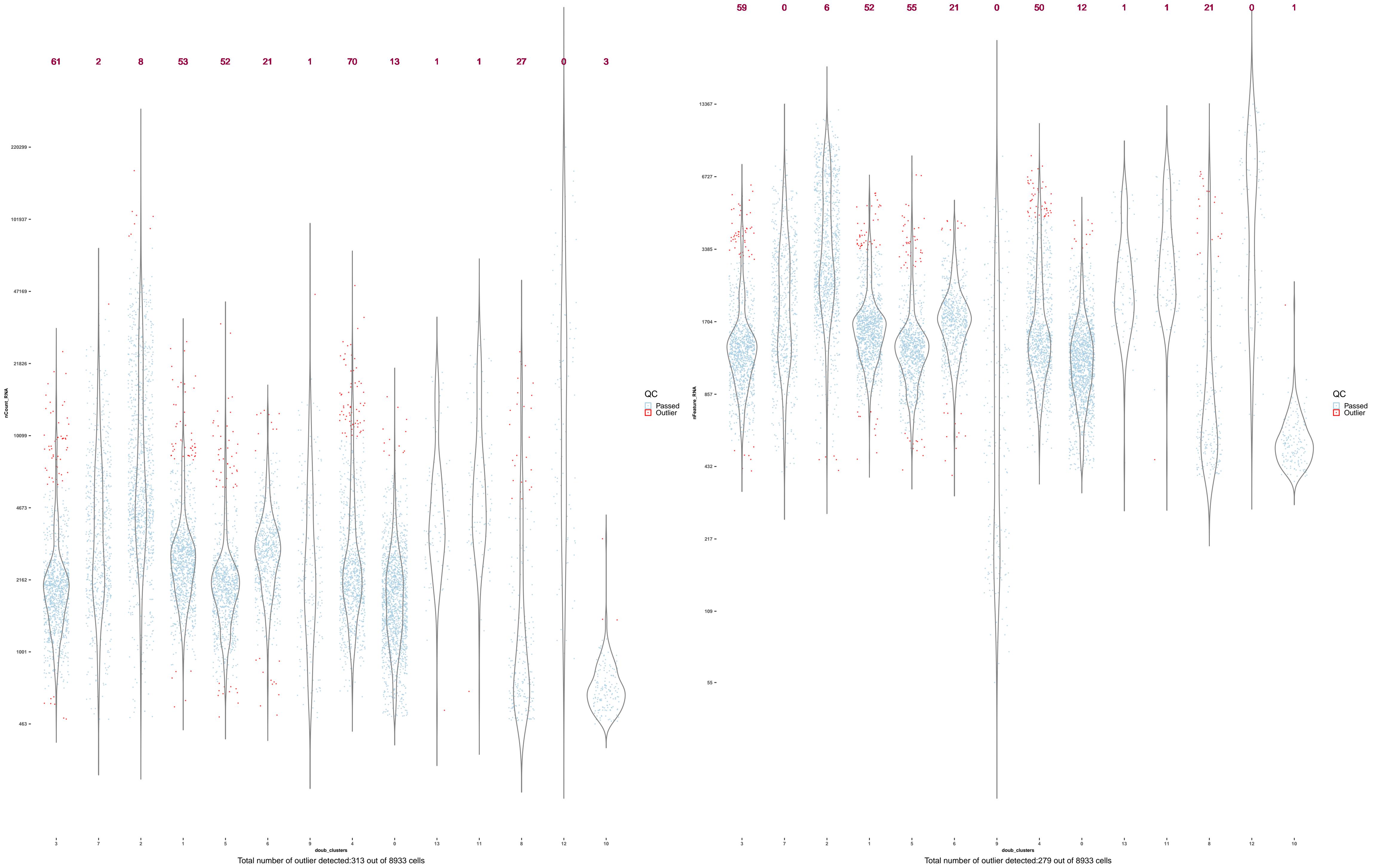
QC

Passed

Outlier

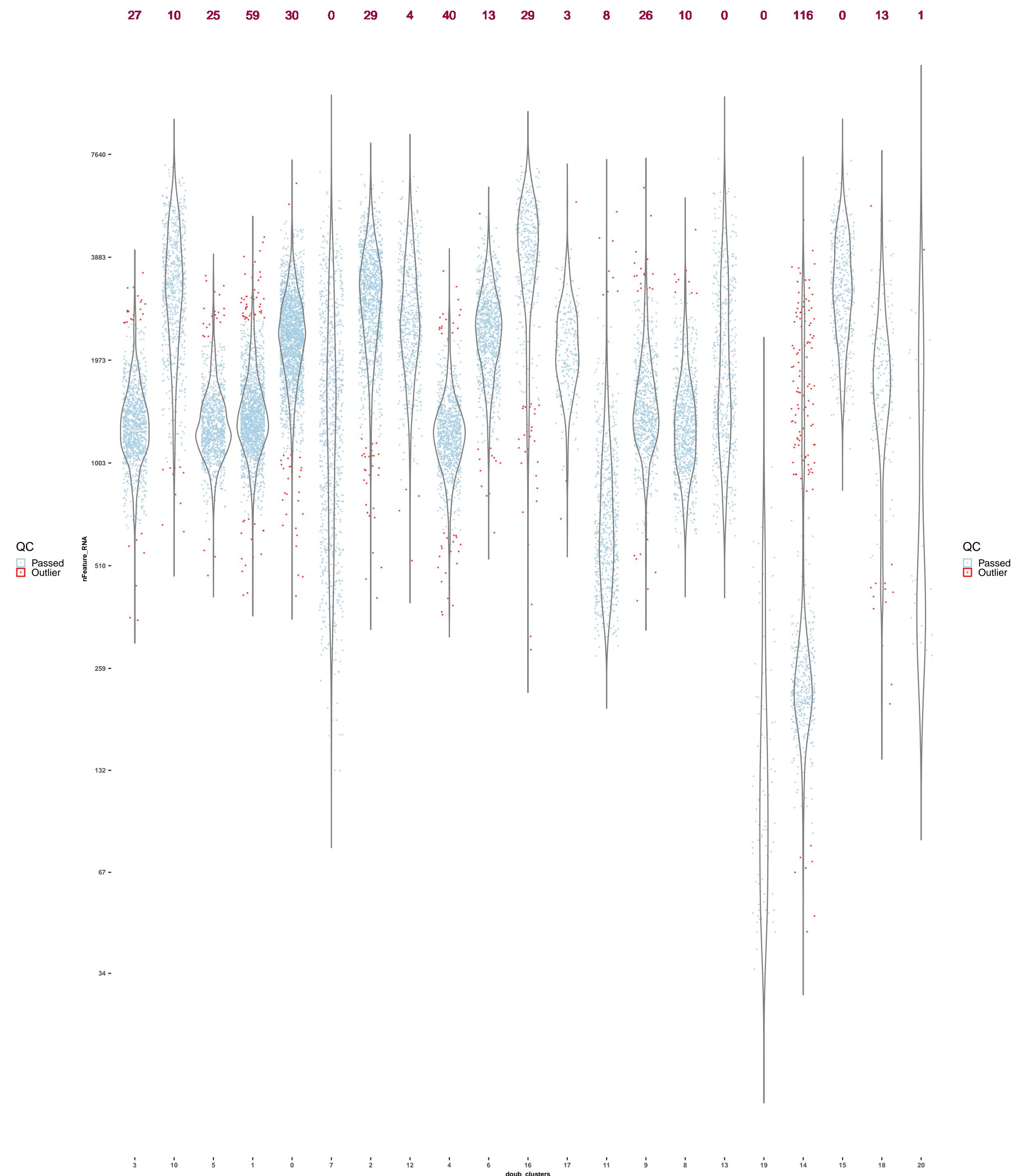
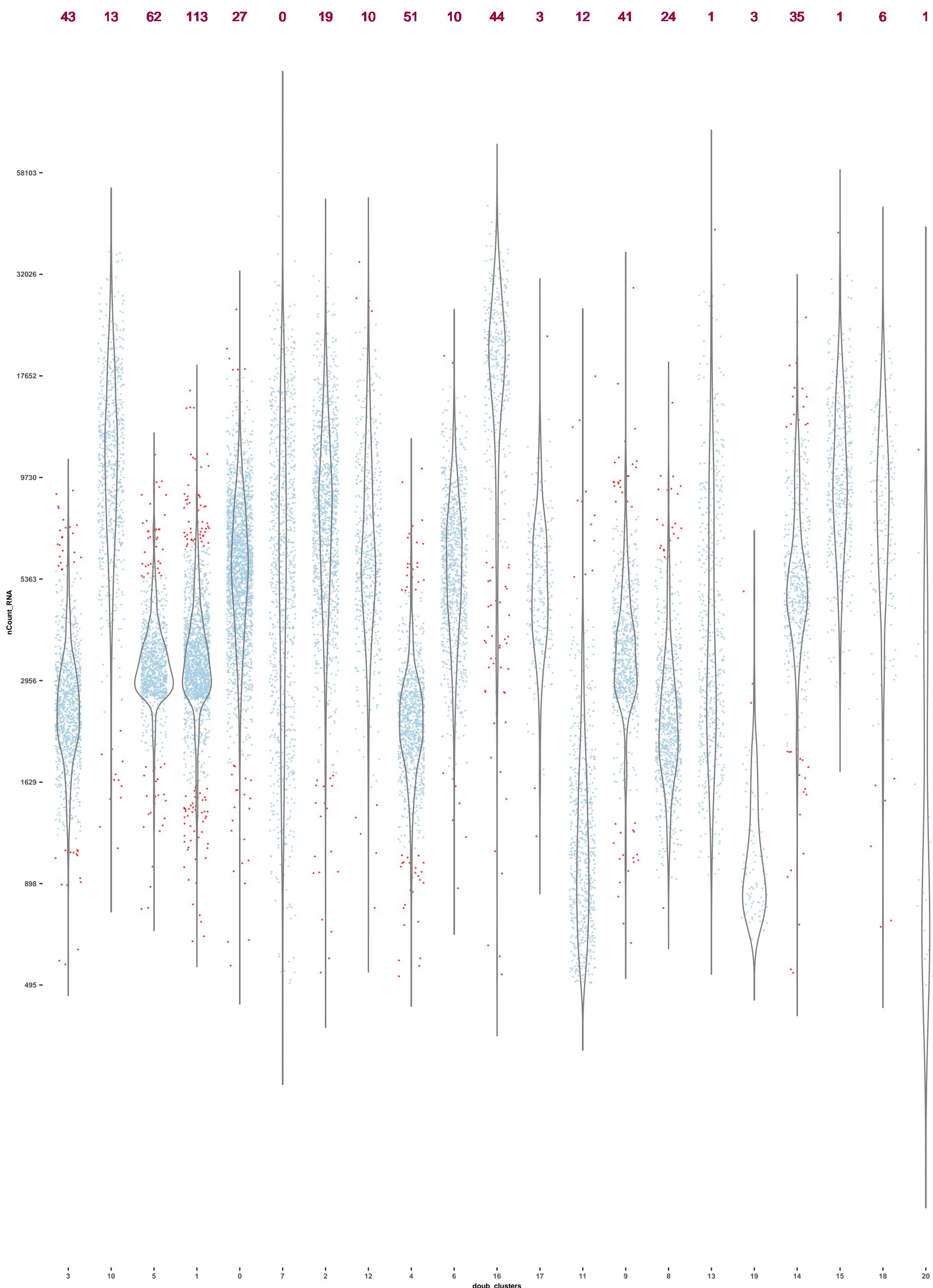
S2_D5_Nuclei(log+1): -3MAD > outliers > +3MAD

S2_D5_Nuclei(log+1): -3MAD > outliers > +3MAD



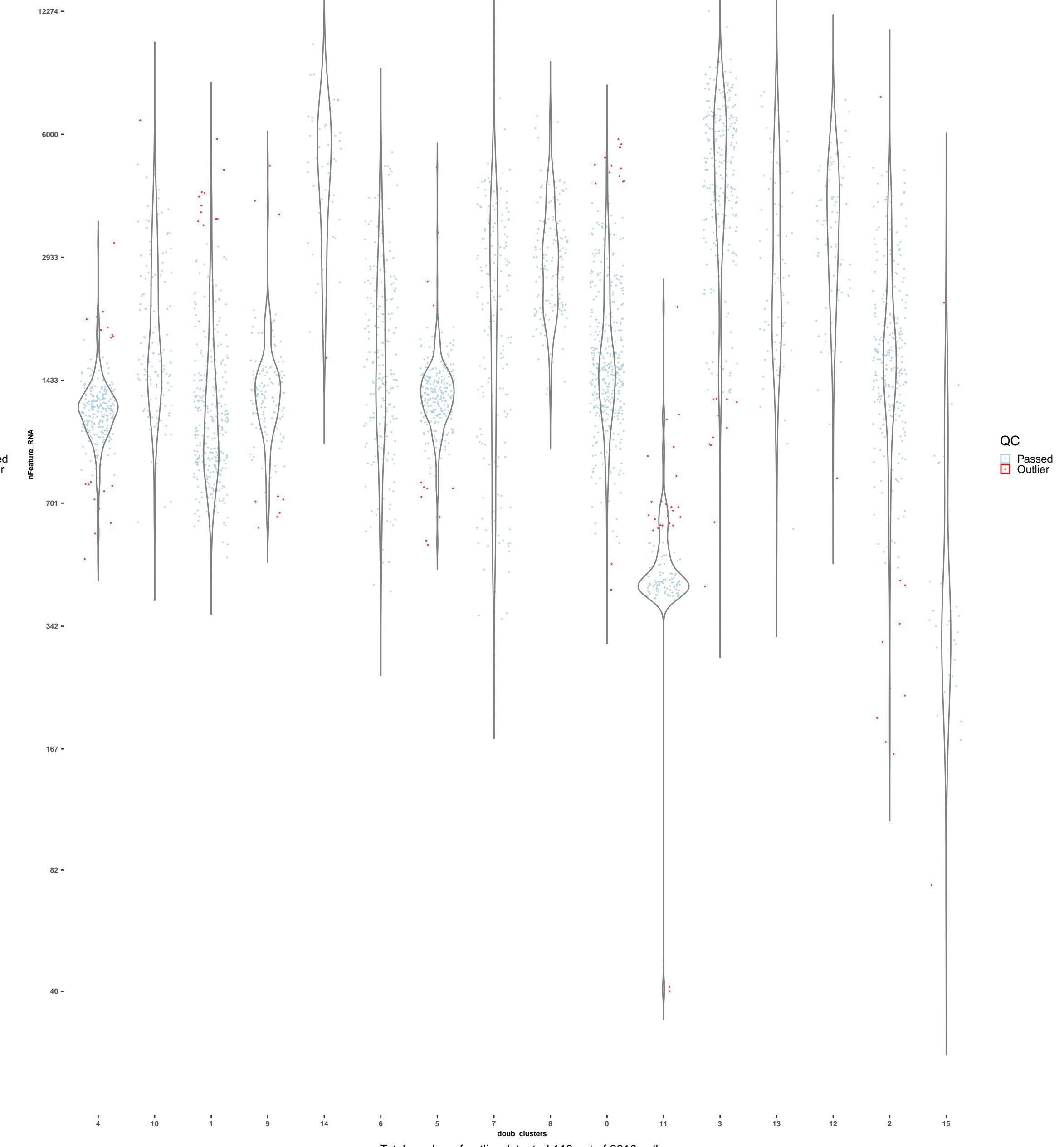
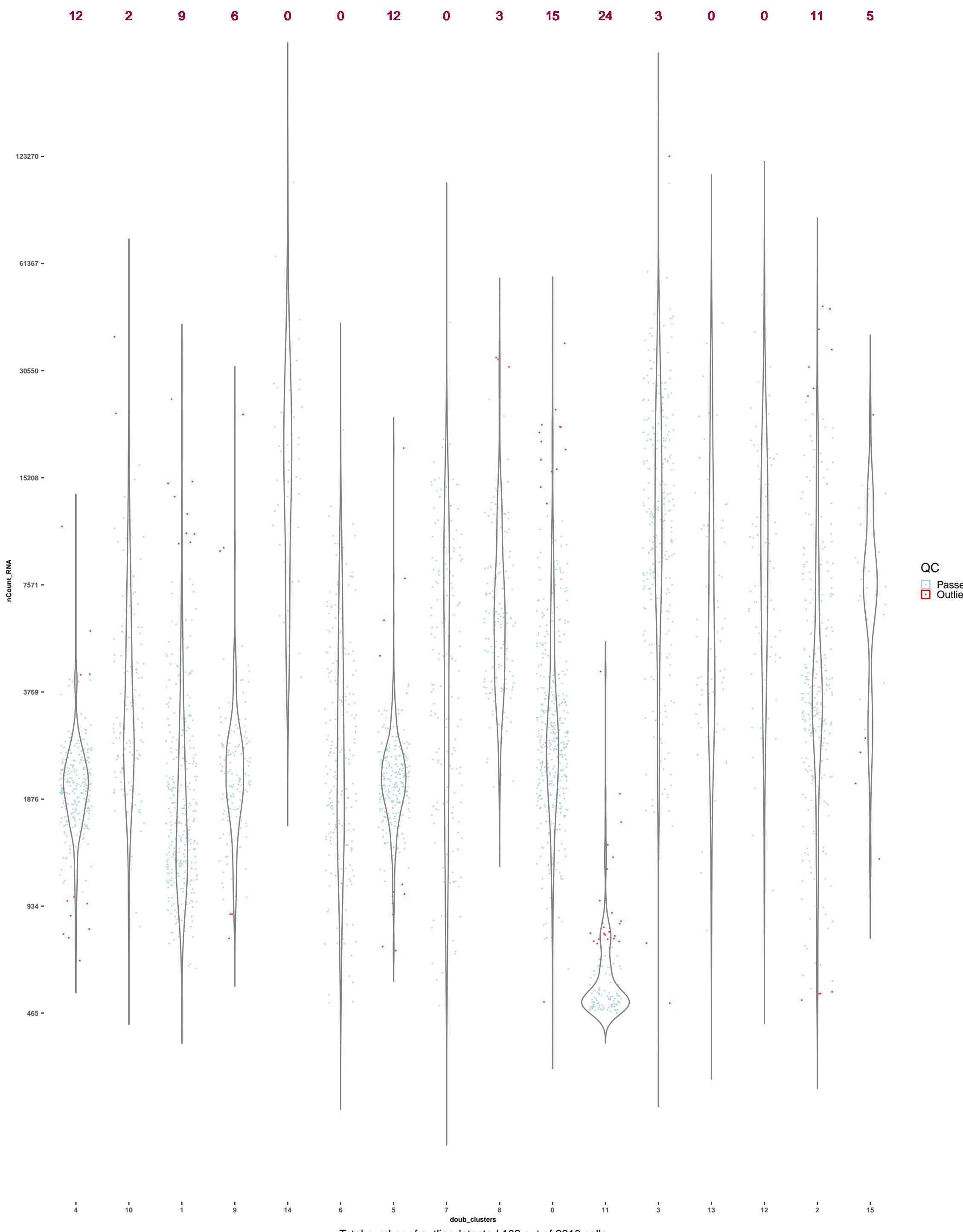
S3_D6_cell(log+1): -3MAD > outliers > +3MAD

S3_D6_cell(log+1): -3MAD > outliers > +3MAD



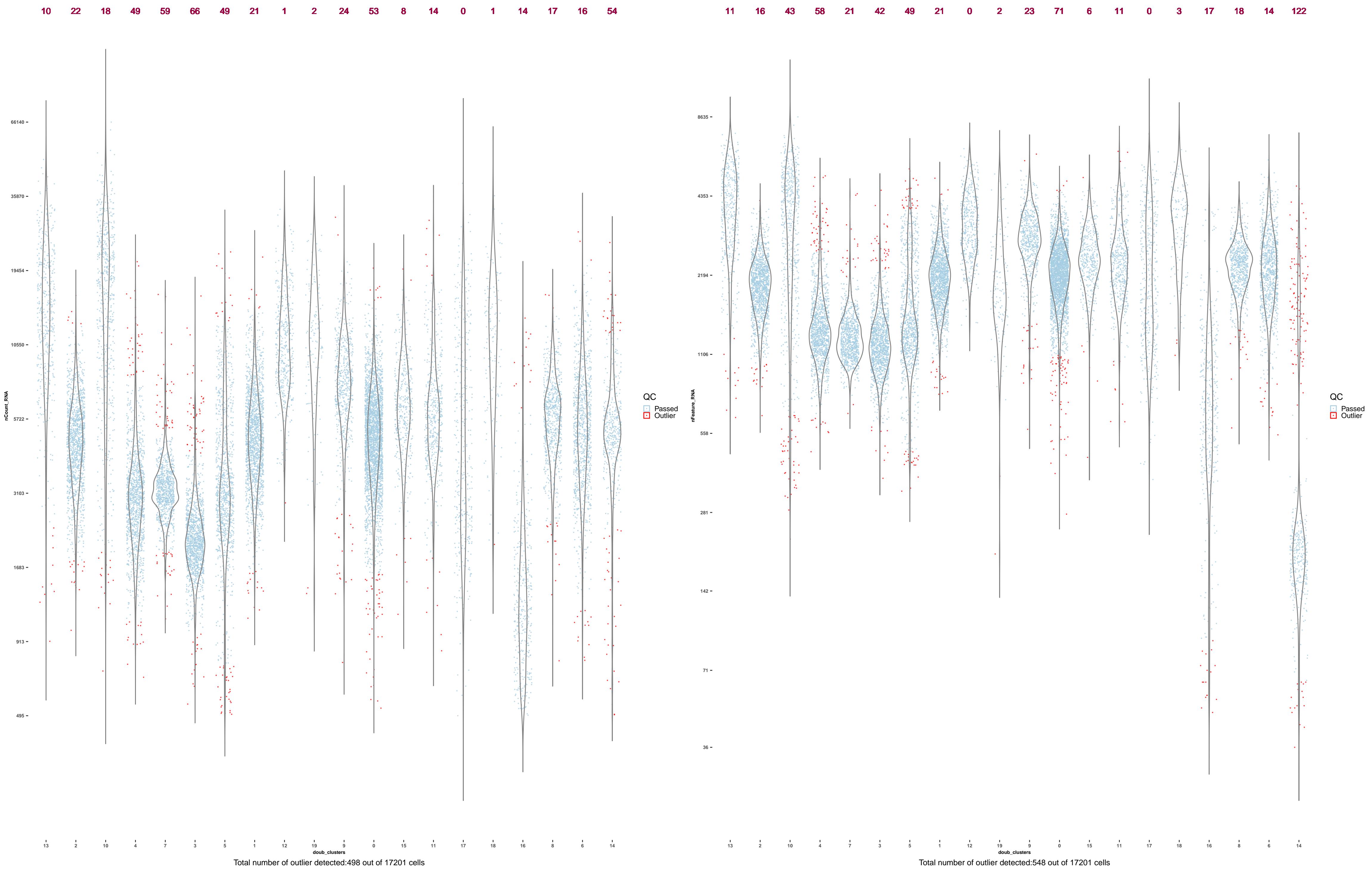
S3_D6_Nuclei(log+1): -3MAD > outliers > +3MAD

S3_D6_Nuclei(log+1): -3MAD > outliers > +3MAD



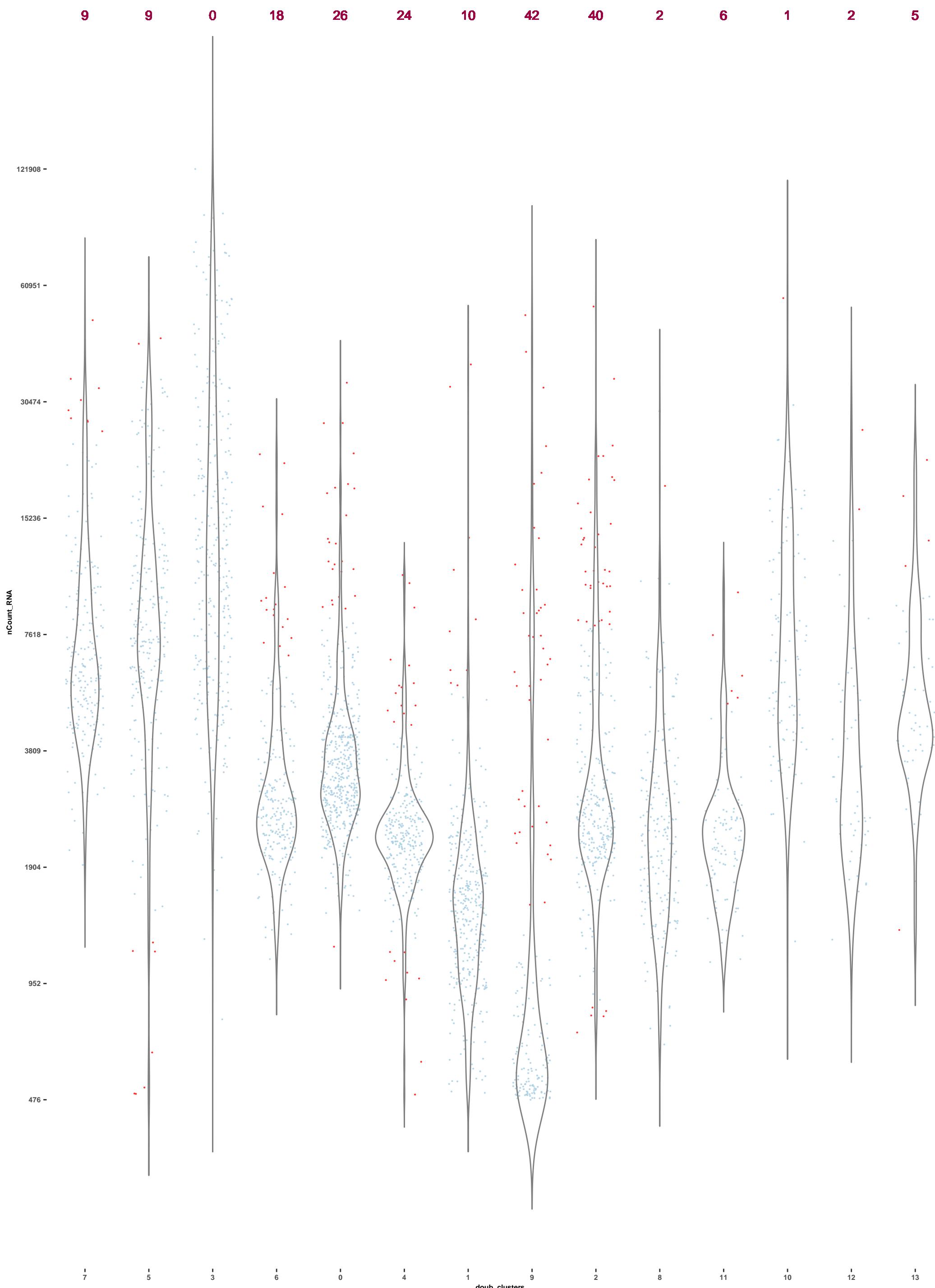
S3_D7_cell(log+1): -3MAD > outliers > +3MAD

S3_D7_cell(log+1): -3MAD > outliers > +3MAD



S3_D7_Nuclei(log+1): -3MAD > outliers > +3MAD

S3_D7_Nuclei(log+1): -3MAD > outliers > +3MAD



QC

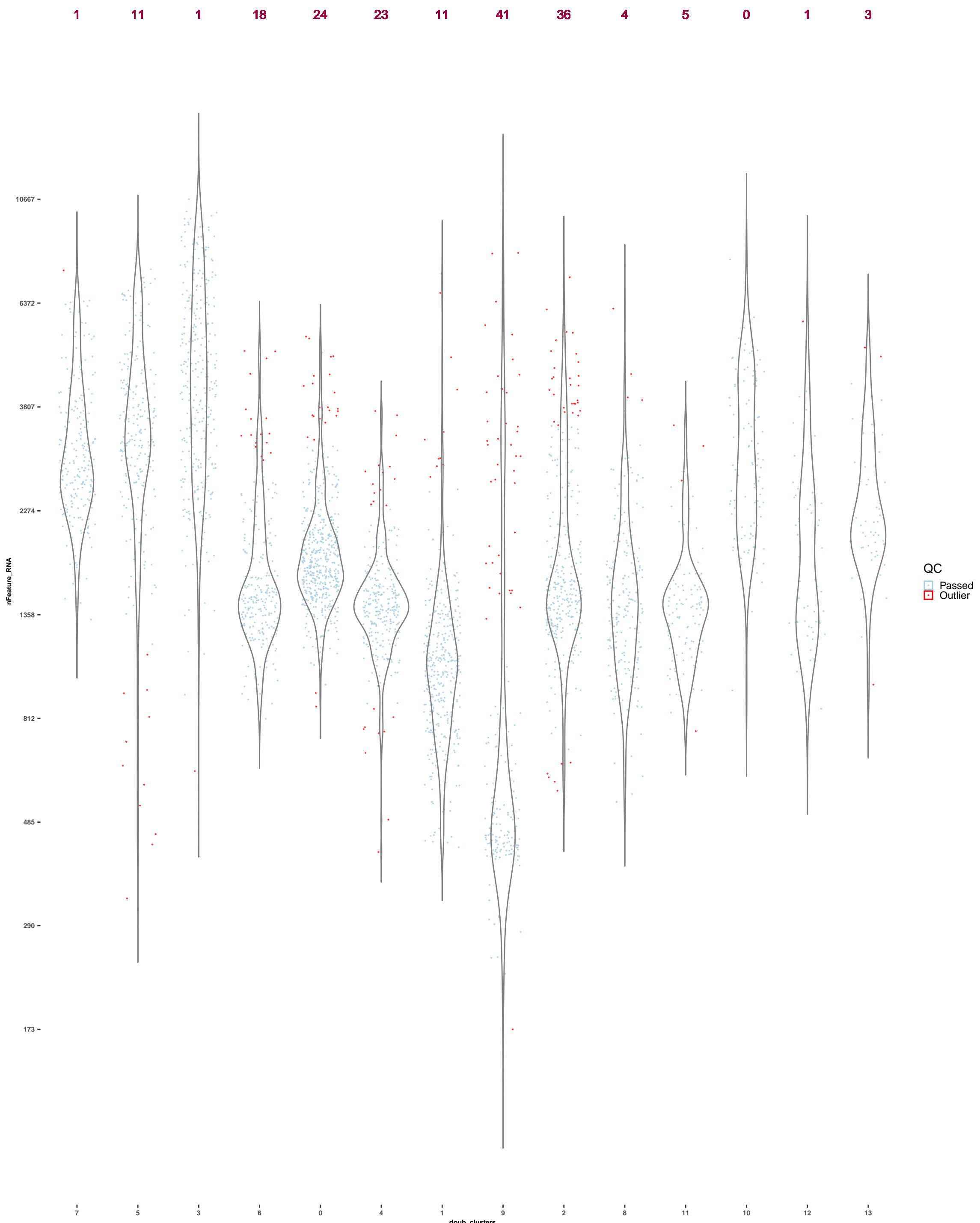
Passed

Outlier

QC

Passed

Outlier

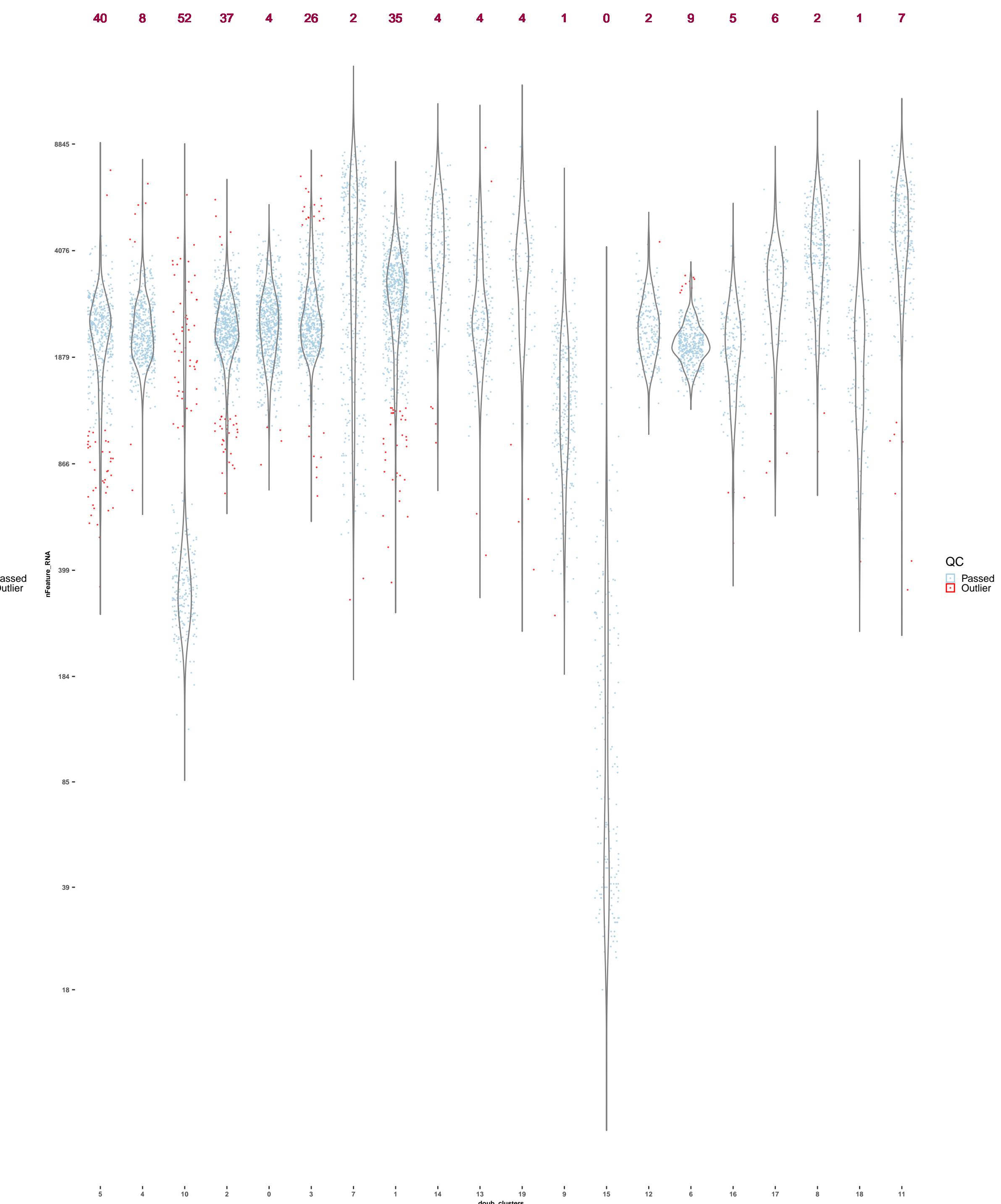
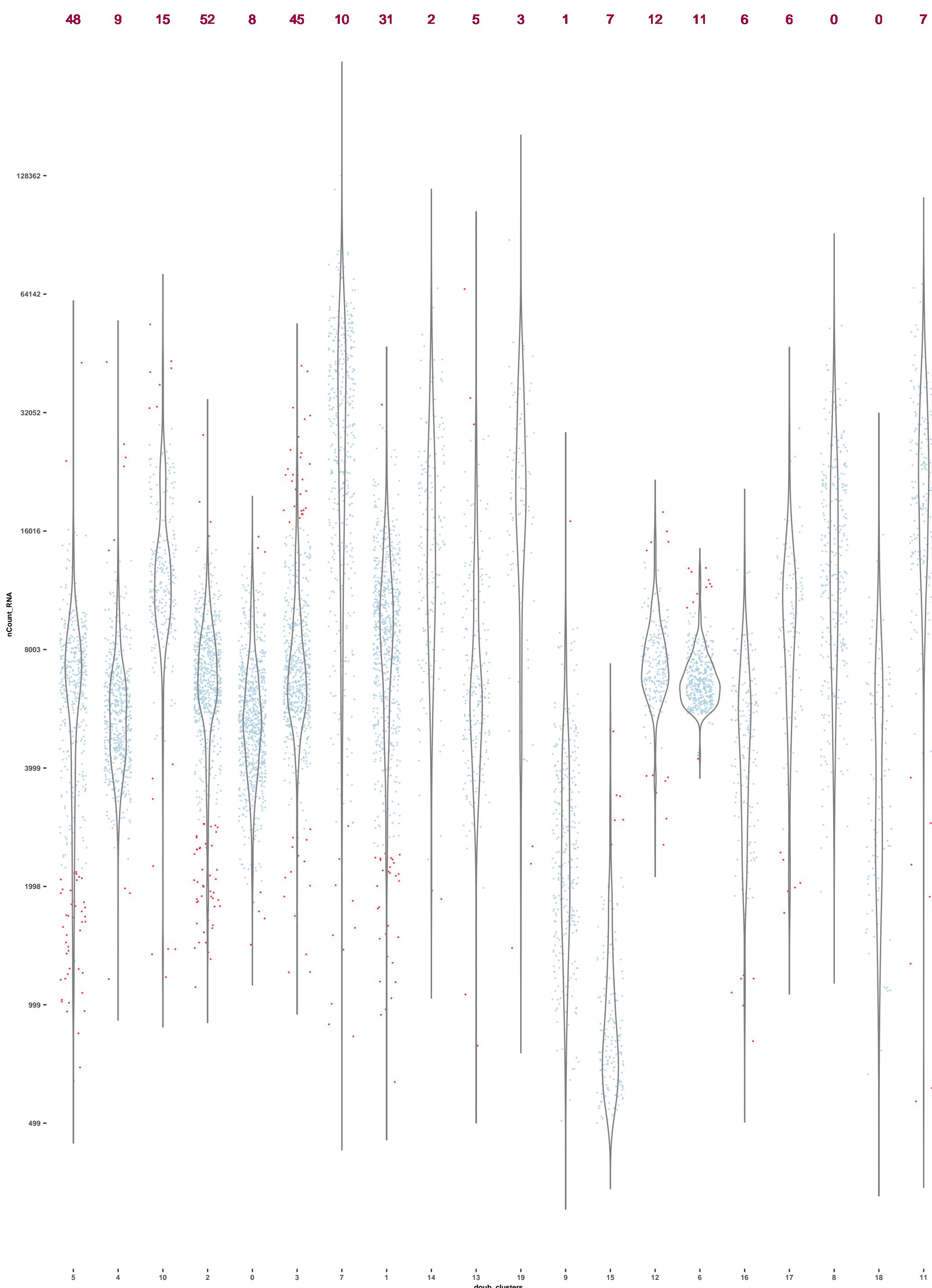


Total number of outlier detected: 194 out of 3041 cells

Total number of outlier detected: 179 out of 3041 cells

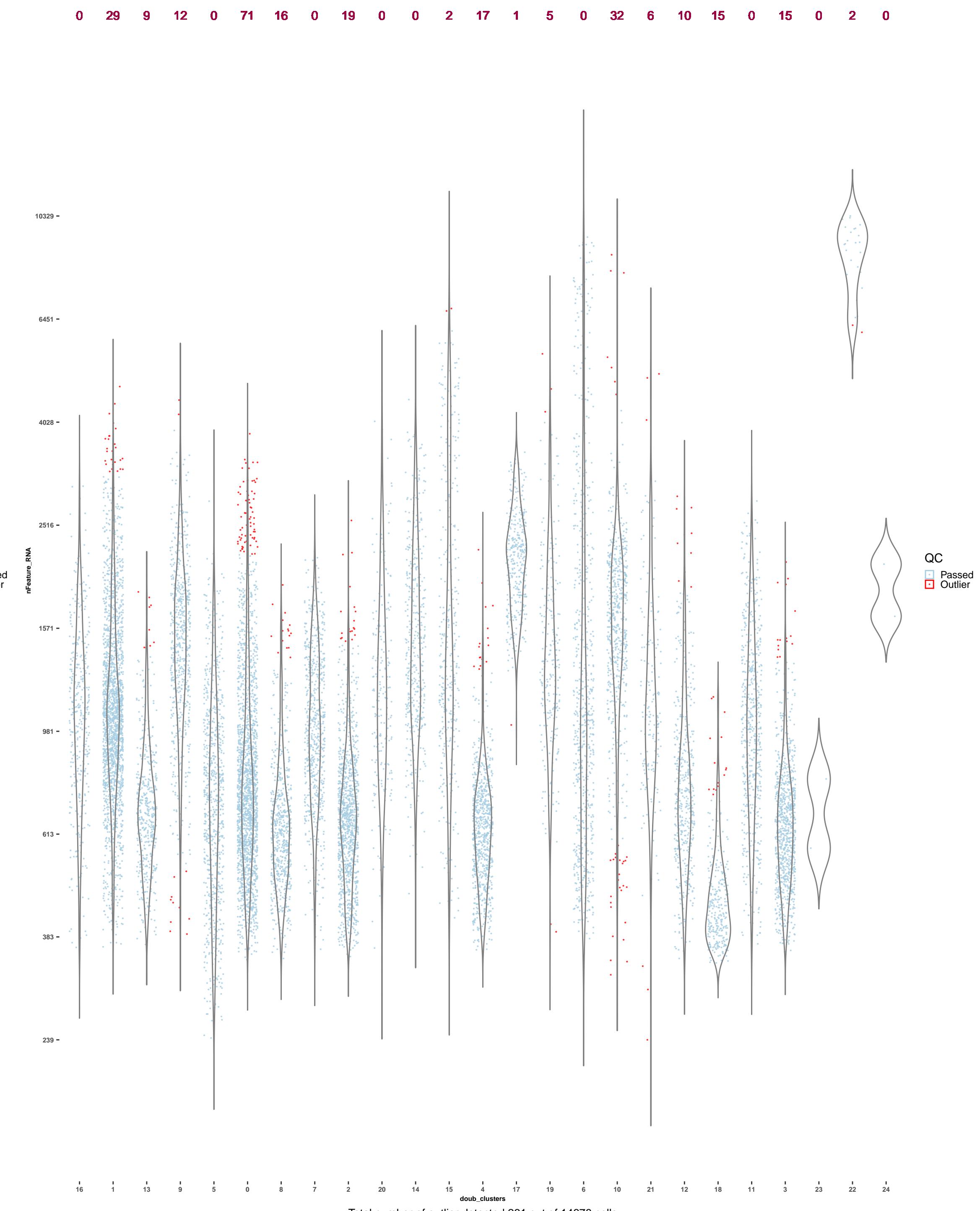
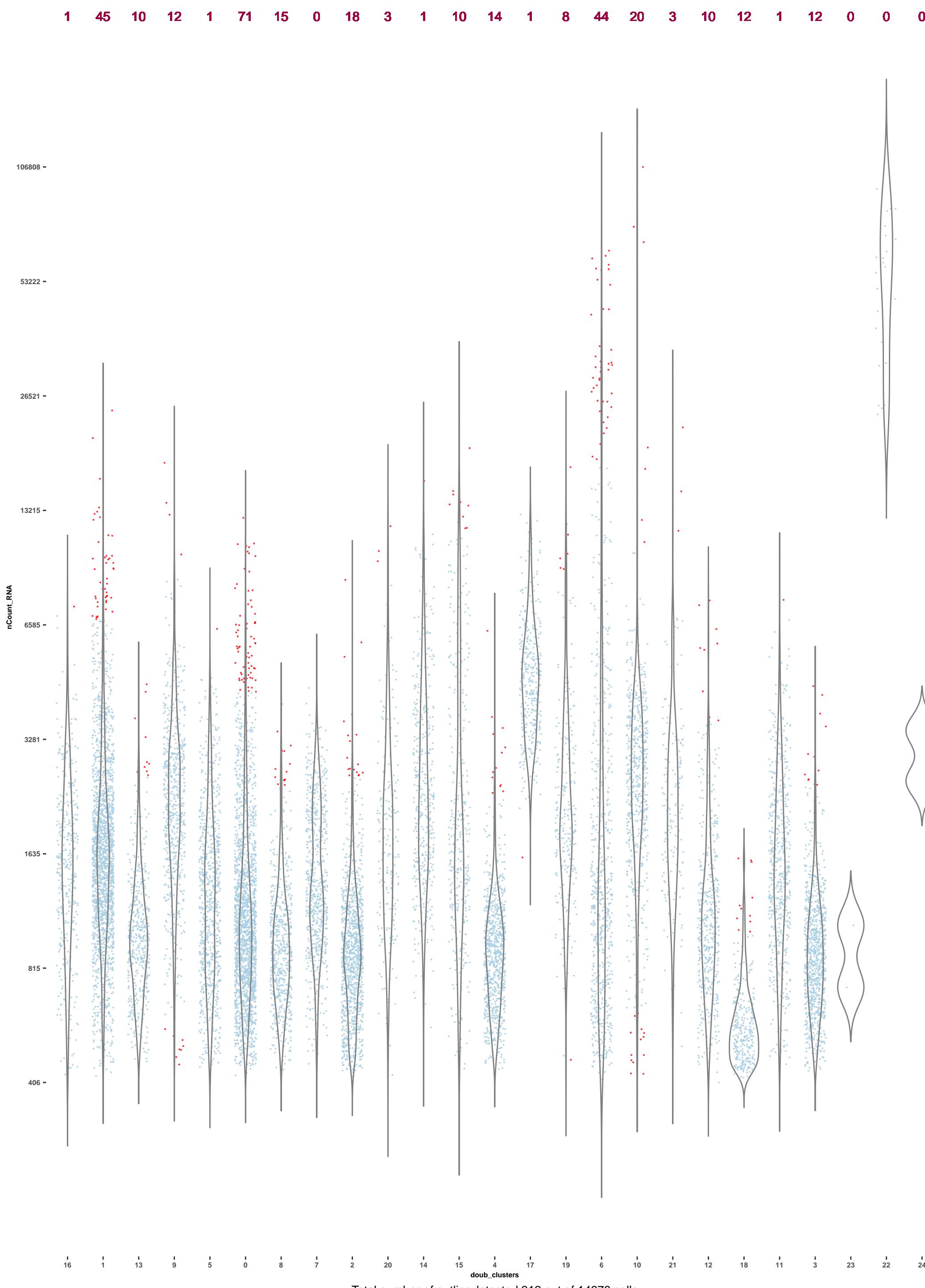
S4_D1_cell(log+1): -3MAD > outliers > +3MAD

S4_D1_cell(log+1): -3MAD > outliers > +3MAD



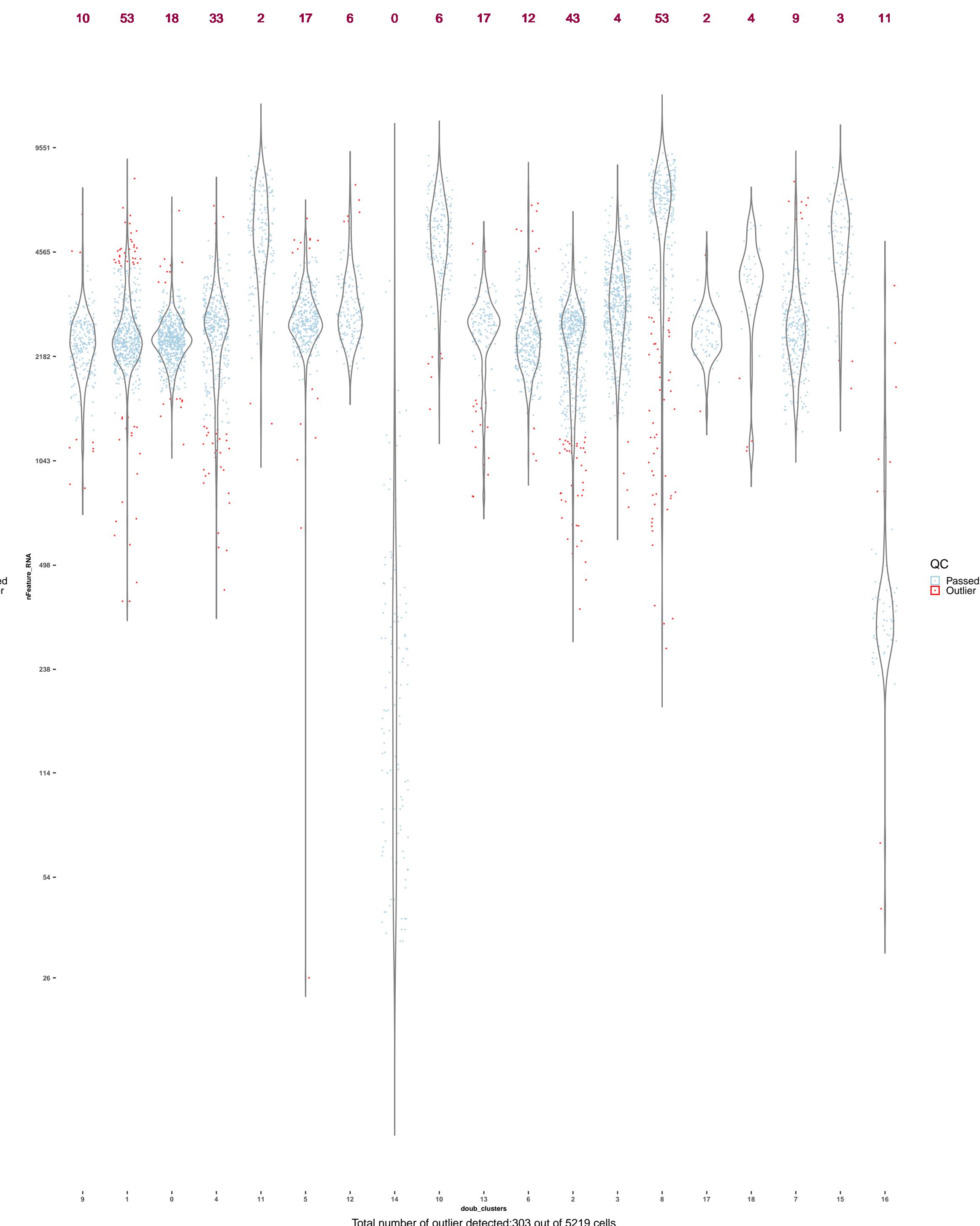
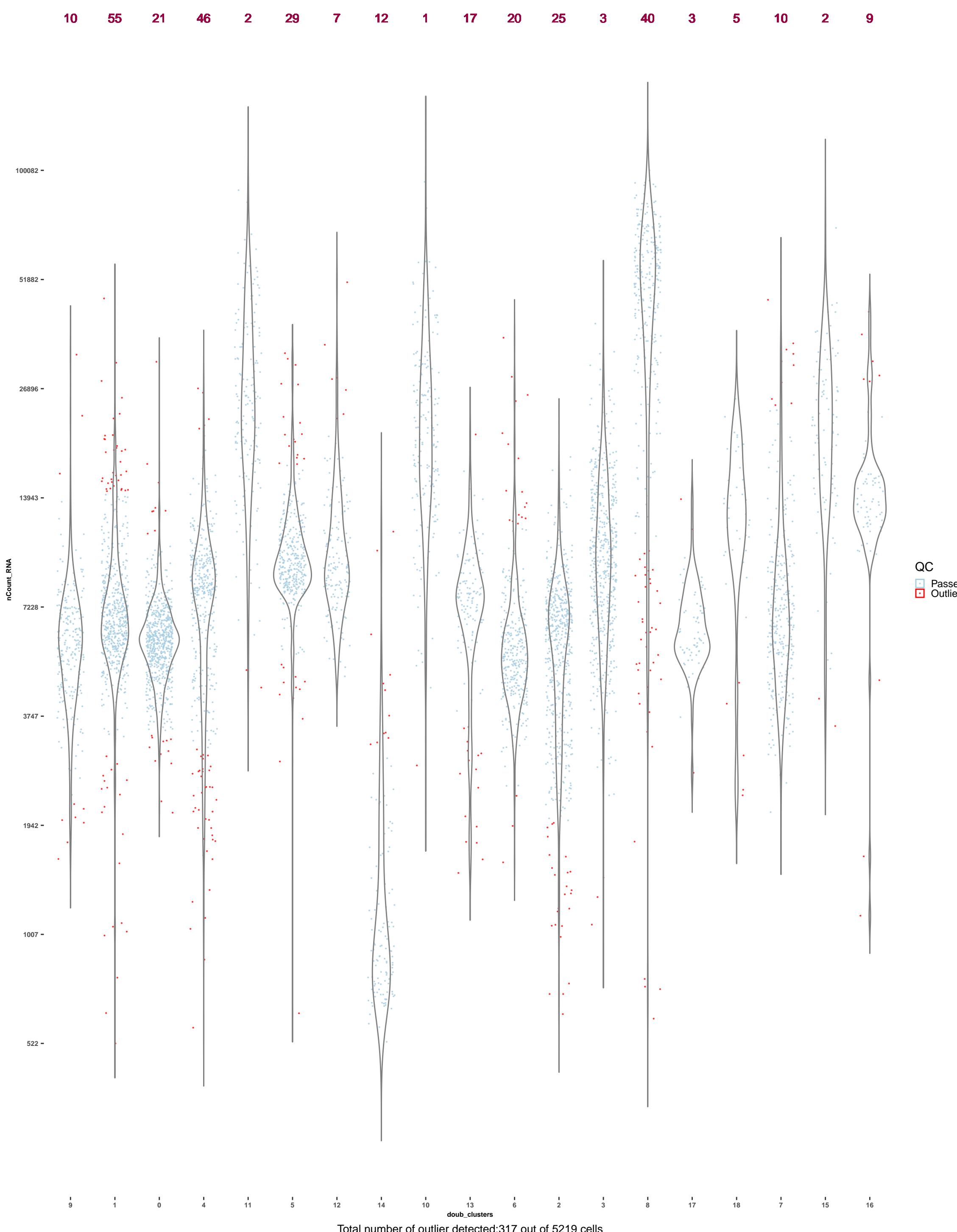
S4_D1_Nuclei(log+1): -3MAD > outliers > +3MAD

S4_D1_Nuclei(log+1): -3MAD > outliers > +3MAD



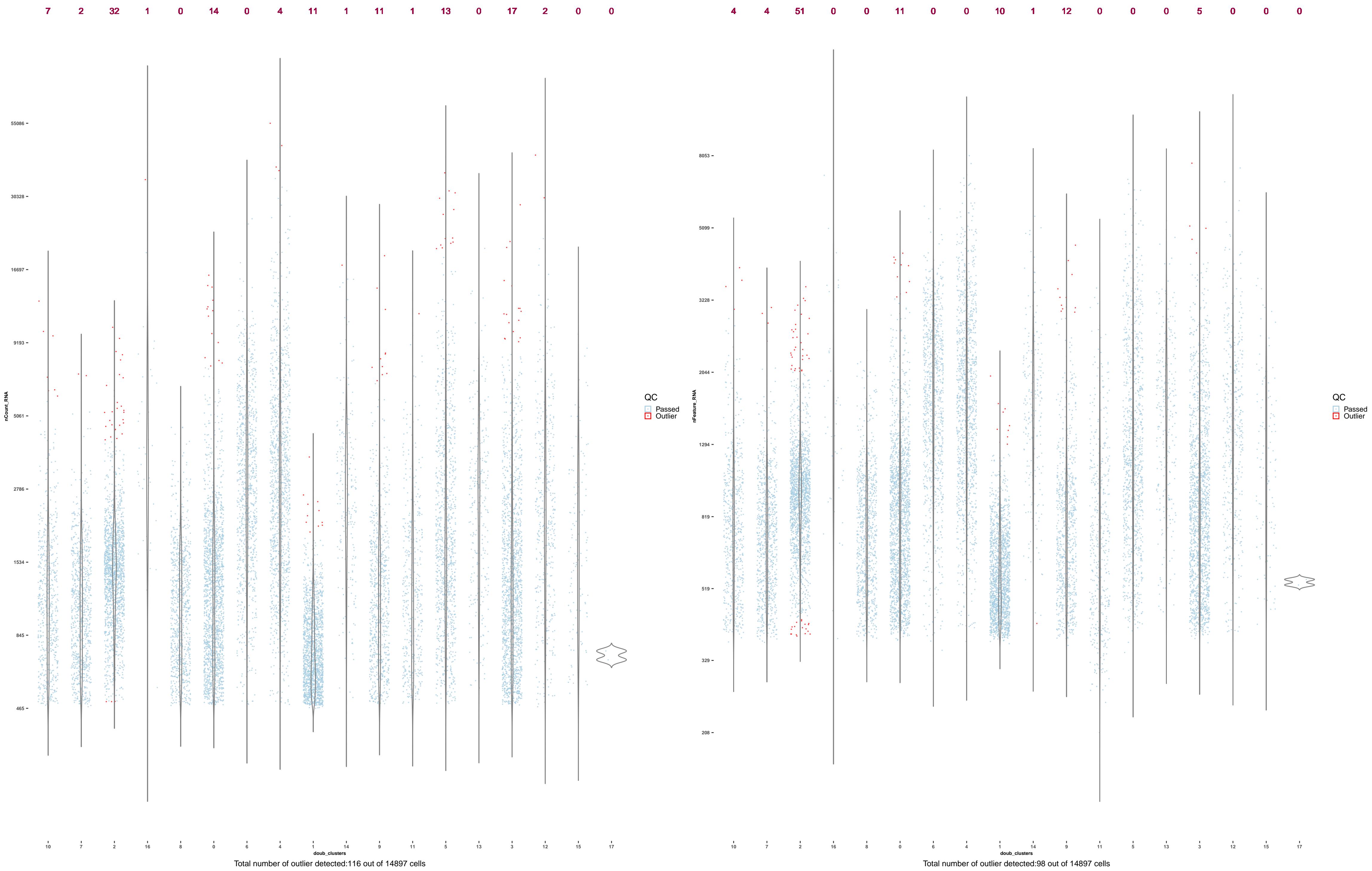
S4_D8_cell(log+1): -3MAD > outliers > +3MAD

S4_D8_cell(log+1): -3MAD > outliers > +3MAD



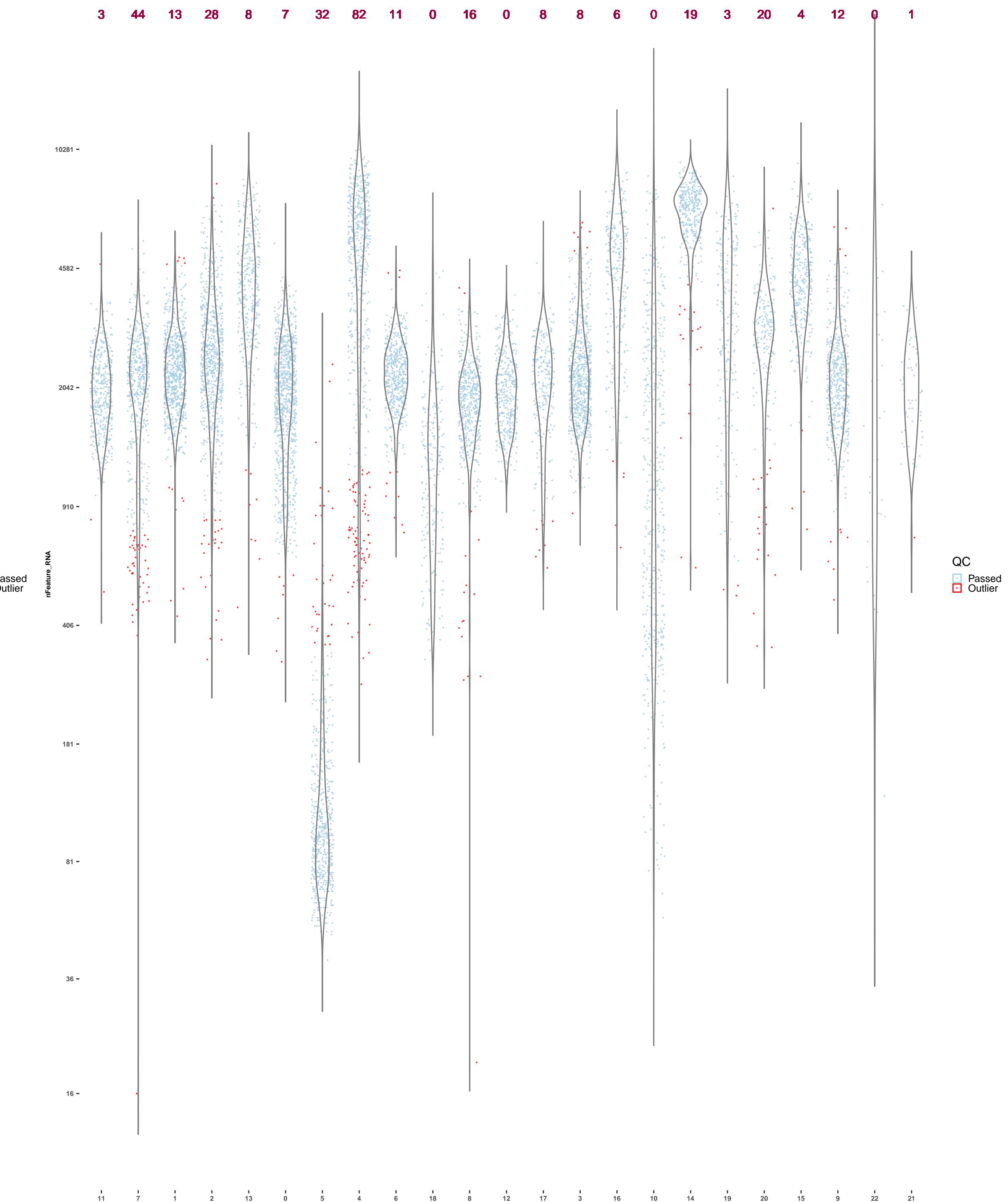
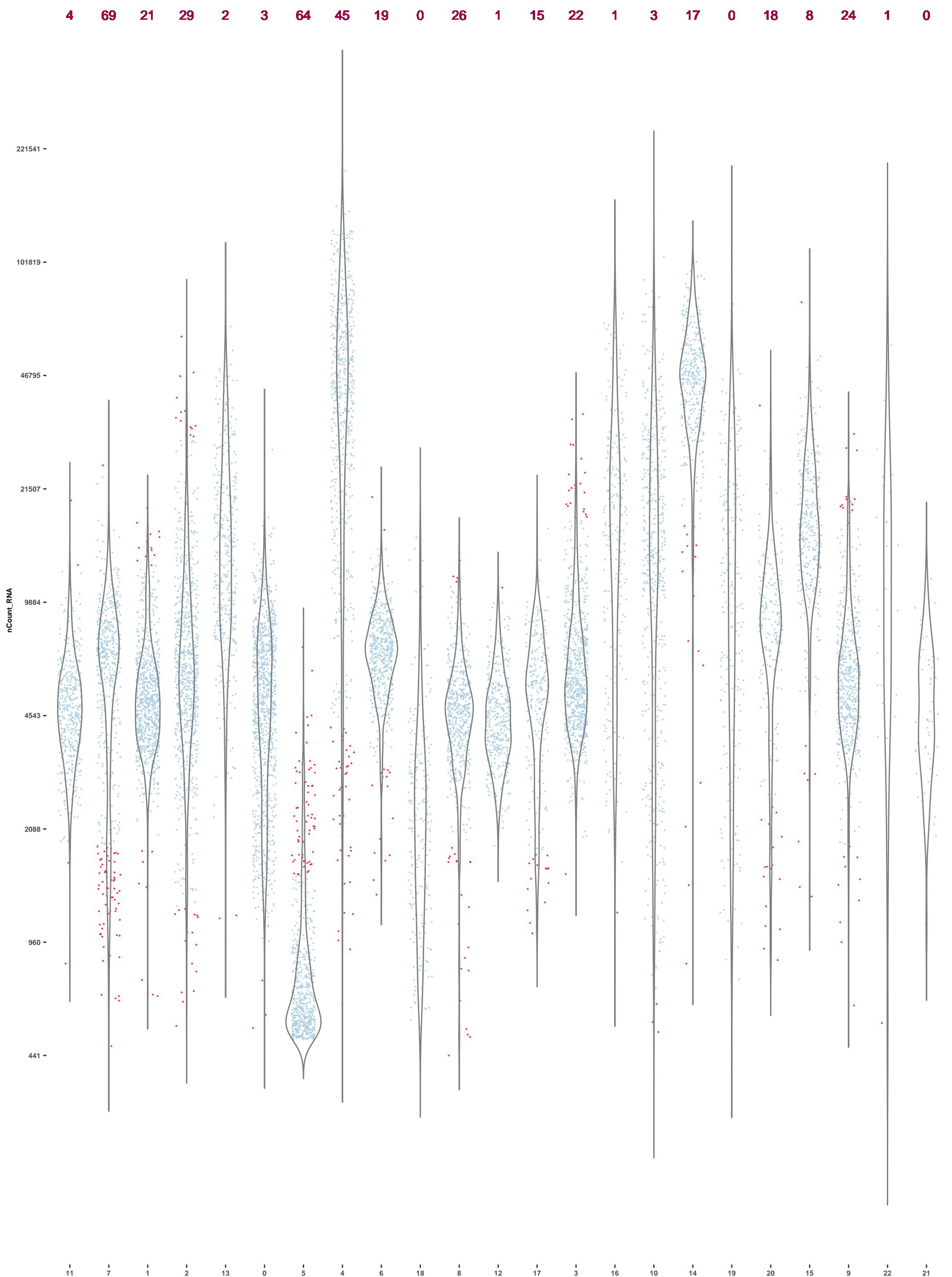
S4_D8_Nuclei(log+1): -3MAD > outliers > +3MAD

S4_D8_Nuclei(log+1): -3MAD > outliers > +3MAD

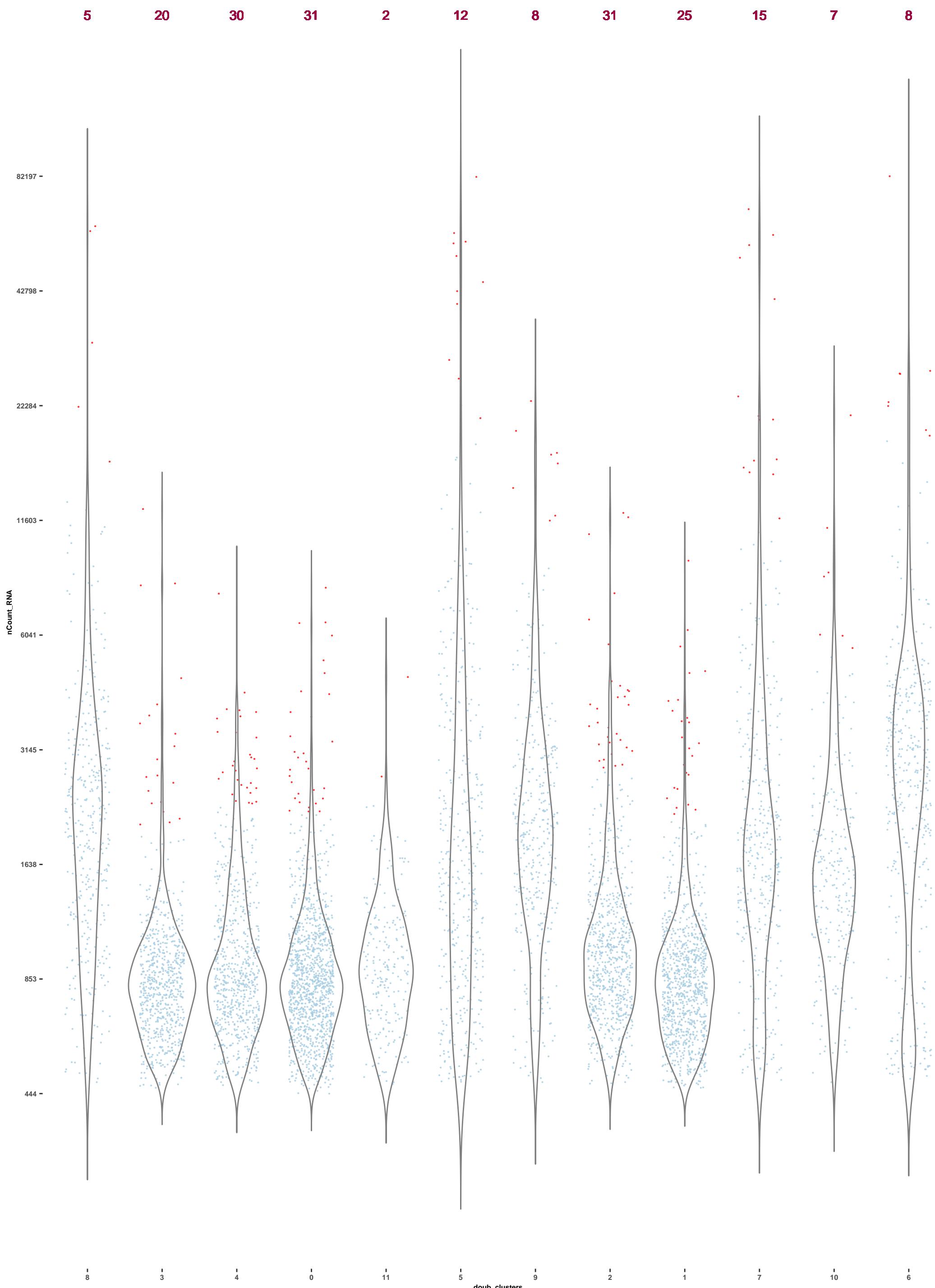


S4_D9_cell(log+1): -3MAD > outliers > +3MAD

S4_D9_cell(log+1): -3MAD > outliers > +3MAD



S4_D9_Nuclei(log+1): -3MAD > outliers > +3MAD



S4_D9_Nuclei(log+1): -3MAD > outliers > +3MAD

