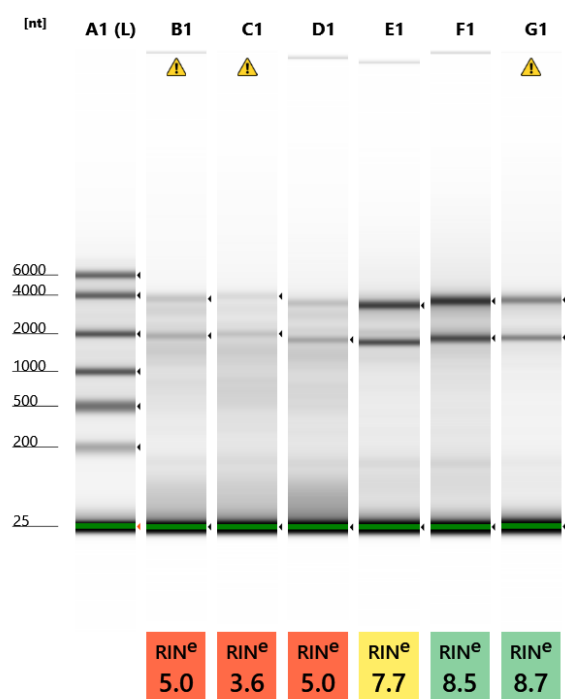


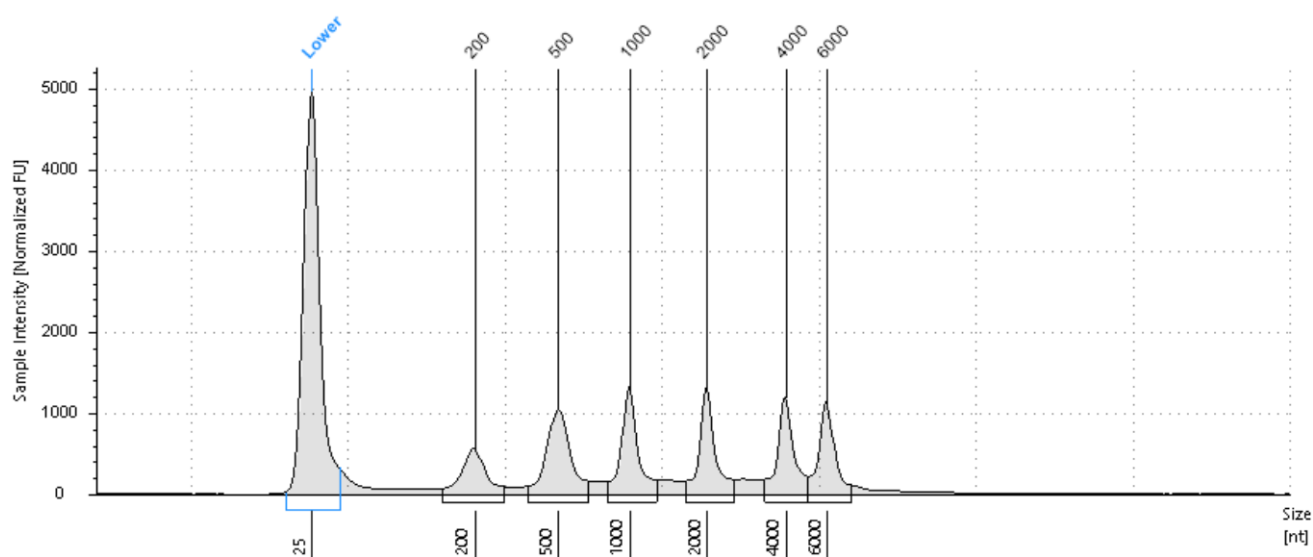
Filename: 2021-02-25 - 16.51.24.RNA



Default image (Contrast 100%)

### Sample Info

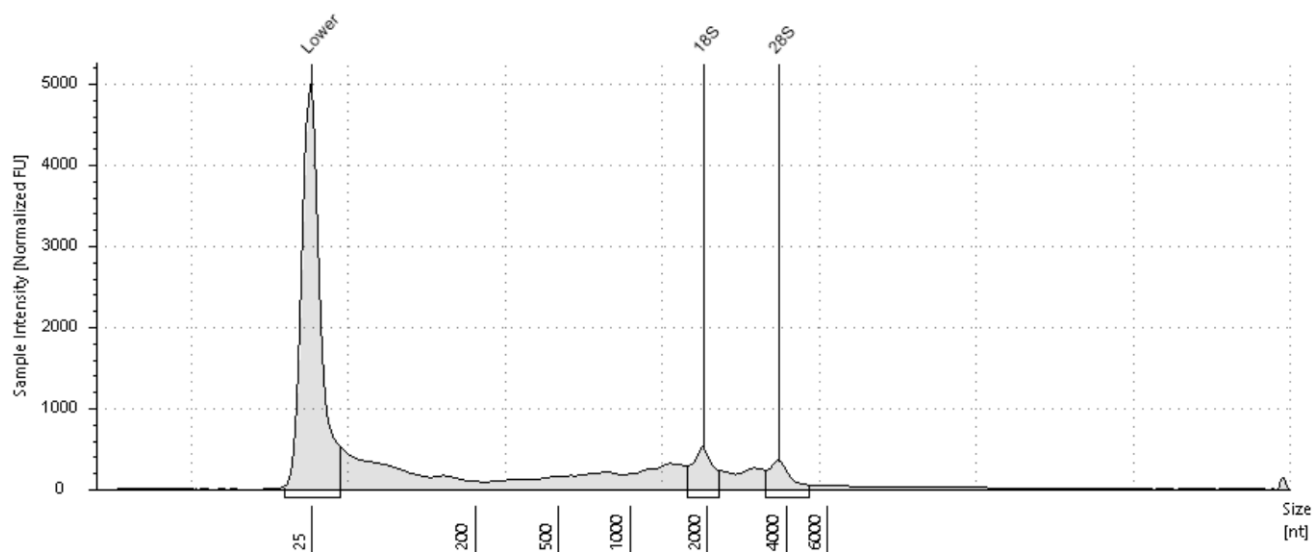
Well	RIN <sup>e</sup>	28S/18S (Area)	Conc. [ng/μl]	Sample Description	Alert	Observations
A1	-	-	40.8	Ladder		Ladder
B1	5.0	0.8	23.6	M209 10-30-2019 (13)	⚠	Sample concentration outside functional range for RIN <sup>e</sup> and the assay
C1	3.6	0.7	23.8	M4 10-30-2019 (14)	⚠	Sample concentration outside functional range for RIN <sup>e</sup> and the assay
D1	5.0	-	27.5	M202 10-30-2019 (15)		The upper ribosomal fragment has degraded
E1	7.7	-	27.0	M217 12-4-2019 (16)		The upper ribosomal fragment has degraded
F1	8.5	1.4	32.9	M212 12-4-2019 (17)		
G1	8.7	1.3	16.3	M4 12-4-2019 (18)	⚠	Sample concentration outside functional range for RIN <sup>e</sup> and the assay

**A1: Ladder****Sample Table**

Well	RINe	28S/18S (Area)	Conc. [ng/ul]	Sample Description	Alert	Observations
A1	-	-	40.8	Ladder		Ladder

**Peak Table**

Size [nt]	Calibrated Conc. [ng/ul]	Assigned Conc. [ng/ul]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	40.0	40.0	4710	-		Lower Marker
200	3.88	-	57.1	11.57		
500	7.01	-	41.3	20.90		
1000	6.08	-	17.9	18.13		
2000	5.92	-	8.71	17.65		
4000	5.59	-	4.11	16.66		
6000	5.07	-	2.48	15.10		

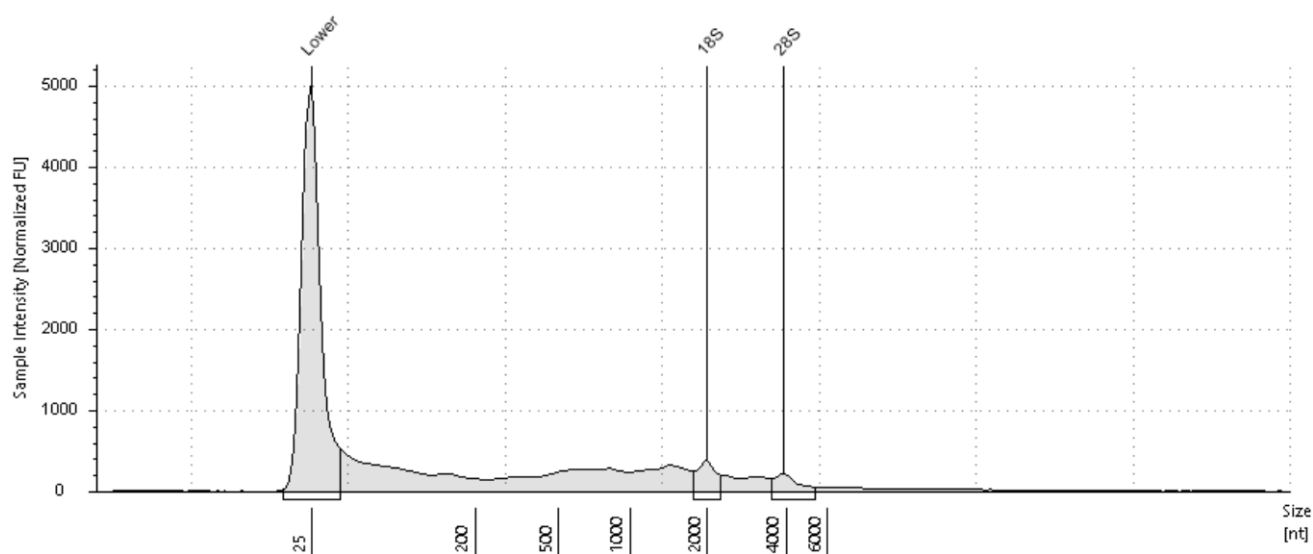
**B1: M209 10-30-2019 (13)****Sample Table**

Well	RINe	28S/18S (Area)	Conc. [ng/μl]	Sample Description	Alert	Observations
B1	5.0	0.8	23.6	M209 10-30-2019 (13)	⚠	Sample concentration outside functional range for RINe and the assay

**Peak Table**

Size [nt]	Calibrated Conc. [ng/μl]	Assigned Conc. [ng/μl]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	40.0	40.0	4710	-		Lower Marker
1949	2.55	-	3.84	56.88		18S
3779	1.93	-	1.50	43.12		28S

## C1: M4 10-30-2019 (14)

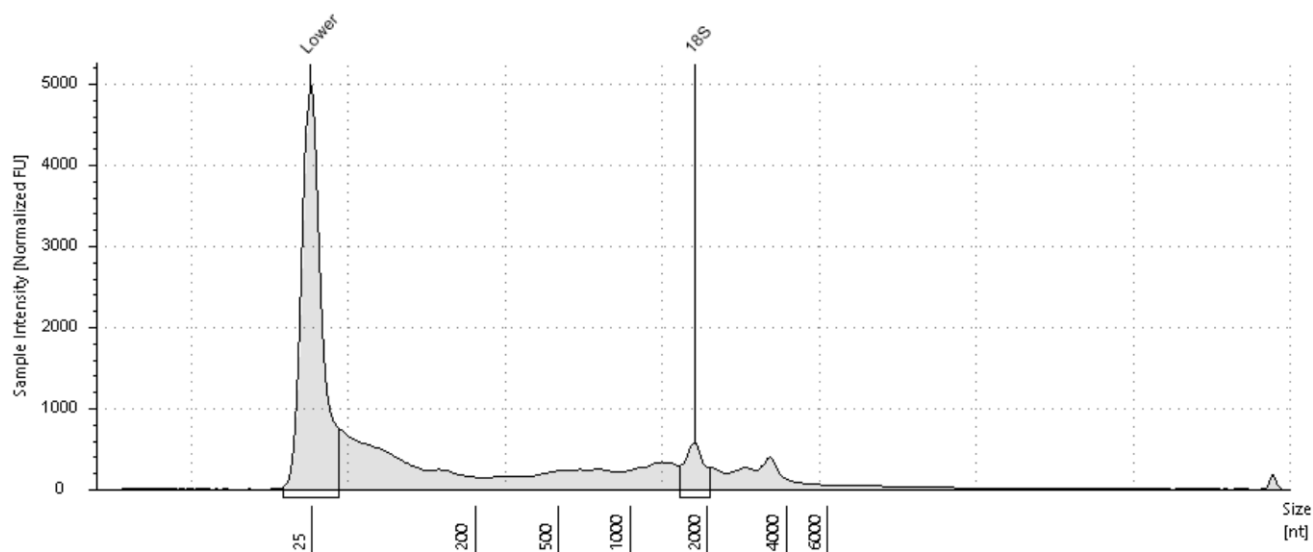


Sample Table

Well	RINe	28S/18S (Area)	Conc. [ng/ul]	Sample Description	Alert	Observations
C1	3.6	0.7	23.8	M4 10-30-2019 (14)	⚠	Sample concentration outside functional range for RINe and the assay

Peak Table

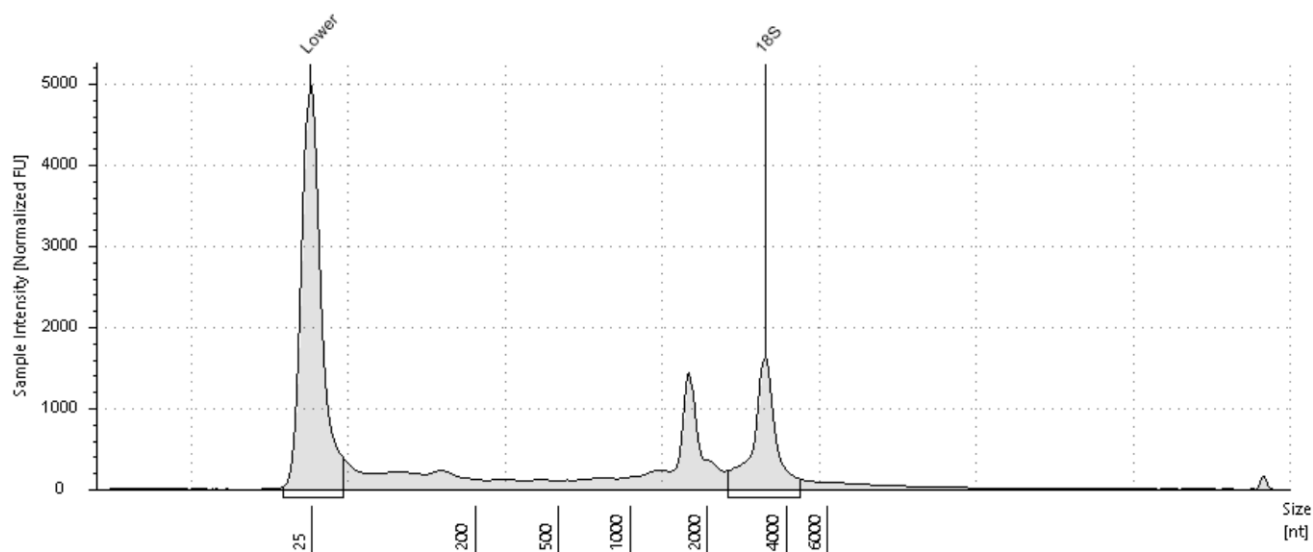
Size [nt]	Calibrated Conc. [ng/ul]	Assigned Conc. [ng/ul]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	40.0	40.0	4710	-		Lower Marker
2016	1.68	-	2.44	57.91		18S
3955	1.22	-	0.906	42.09		28S

**D1: M202 10-30-2019 (15)****Sample Table**

Well	RINe	28S/18S (Area)	Conc. [ng/μl]	Sample Description	Alert	Observations
D1	5.0	-	27.5	M202 10-30-2019 (15)		The upper ribosomal fragment has degraded

**Peak Table**

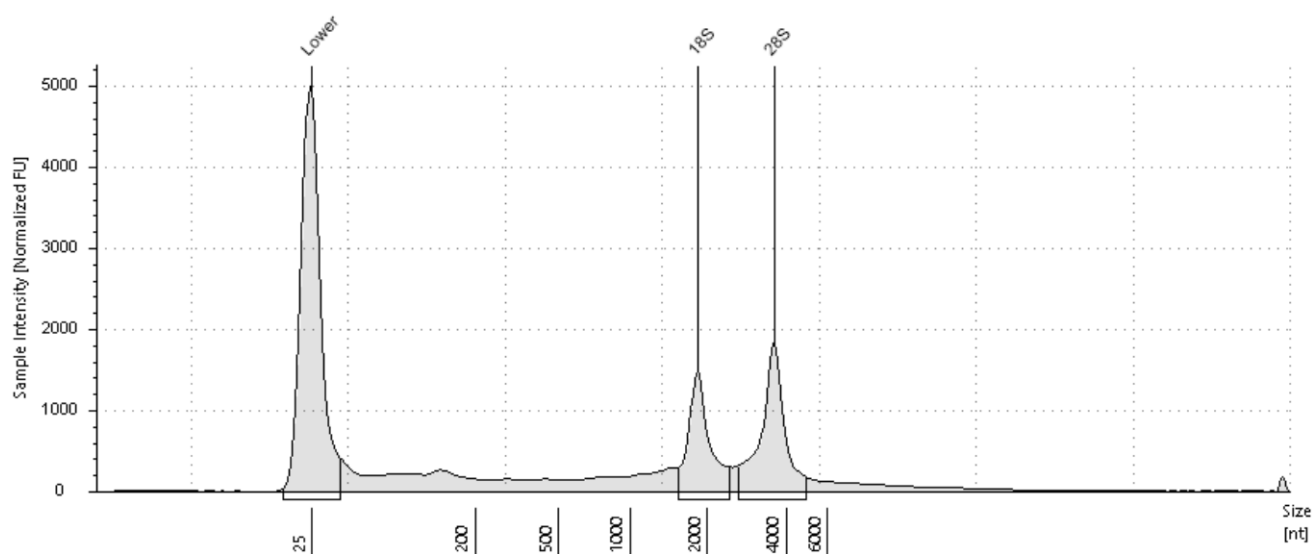
Size [nt]	Calibrated Conc. [ng/μl]	Assigned Conc. [ng/μl]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	40.0	40.0	4710	-		Lower Marker
1818	2.46	-	3.98	100.00		18S

**E1: M217 12-4-2019 (16)****Sample Table**

Well	RINe	28S/18S (Area)	Conc. [ng/μl]	Sample Description	Alert	Observations
E1	7.7	-	27.0	M217 12-4-2019 (16)		The upper ribosomal fragment has degraded

**Peak Table**

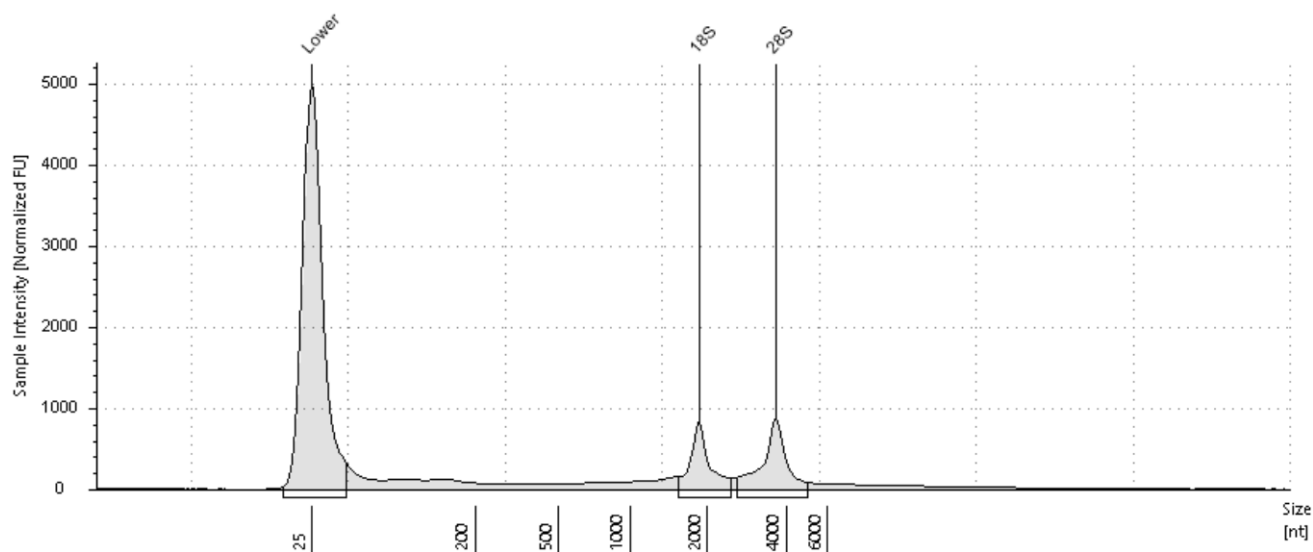
Size [nt]	Calibrated Conc. [ng/μl]	Assigned Conc. [ng/μl]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	40.0	40.0	4710	-		Lower Marker
3353	8.10	-	7.11	100.00		18S

**F1: M212 12-4-2019 (17)****Sample Table**

Well	RINe	28S/18S (Area)	Conc. [ng/ul]	Sample Description	Alert	Observations
F1	8.5	1.4	32.9	M212 12-4-2019 (17)		

**Peak Table**

Size [nt]	Calibrated Conc. [ng/ul]	Assigned Conc. [ng/ul]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	40.0	40.0	4710	-		Lower Marker
1863	6.98	-	11.0	41.56		18S
3627	9.82	-	7.96	58.44		28S

**G1: M4 12-4-2019 (18)****Sample Table**

Well	RINe	28S/18S (Area)	Conc. [ng/ul]	Sample Description	Alert	Observations
G1	8.7	1.3	16.3	M4 12-4-2019 (18)	⚠	Sample concentration outside functional range for RINe and the assay

**Peak Table**

Size [nt]	Calibrated Conc. [ng/ul]	Assigned Conc. [ng/ul]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	40.0	40.0	4710	-		Lower Marker
1879	3.39	-	5.31	43.08		18S
3697	4.49	-	3.57	56.92		28S