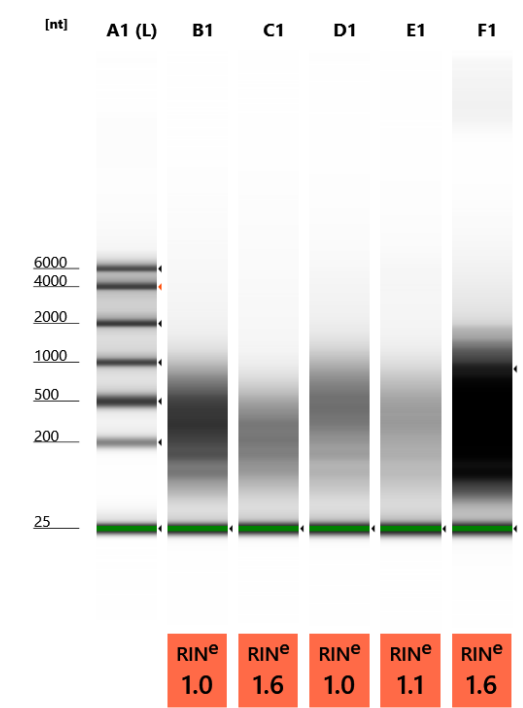


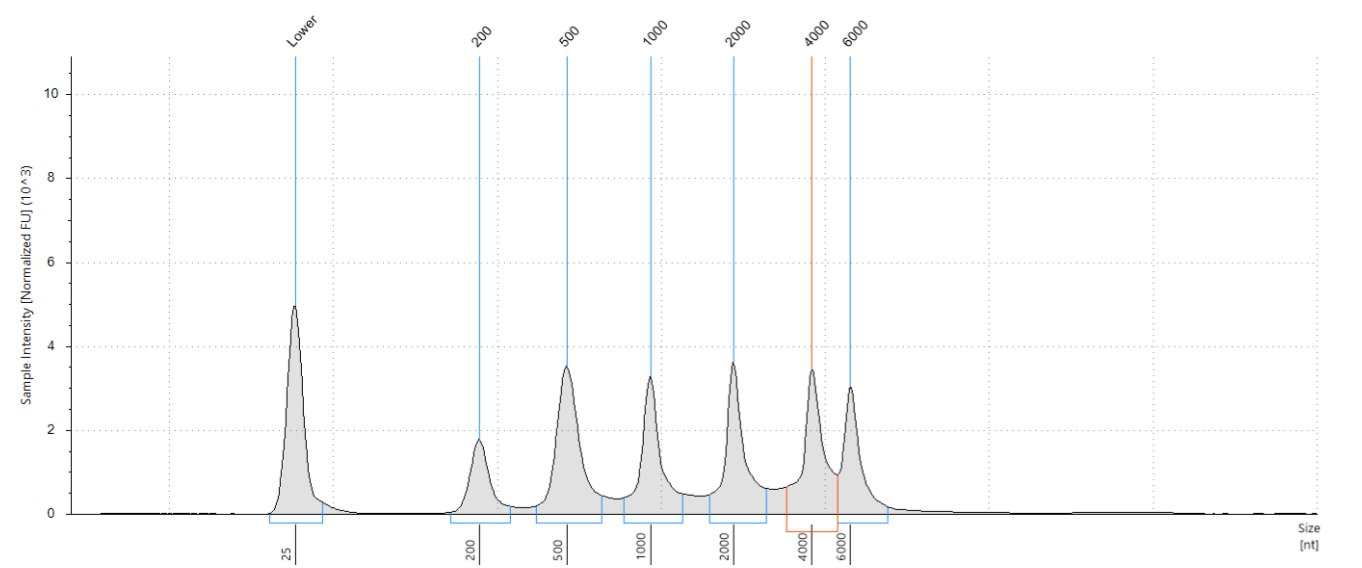
Filename: 2018-12-20 - 15.03.08.RNA**Sample Info**

Well	RINe	28S/18S (Area)	Conc. [ng/µl]	Sample Description	Alert	Observations
A1	-	-	112	Ladder		Ladder
B1	1.0	-	165	Mytilus RNA 1		
C1	1.6	-	80.0	Mytilus RNA 2		
D1	1.0	-	96.0	Mytilus RNA 3		
E1	1.1	-	60.8	Mytilus RNA 4		
F1	1.6	-	411	Mytilus RNA 5		



Default image (Contrast 100%)

A1: Ladder

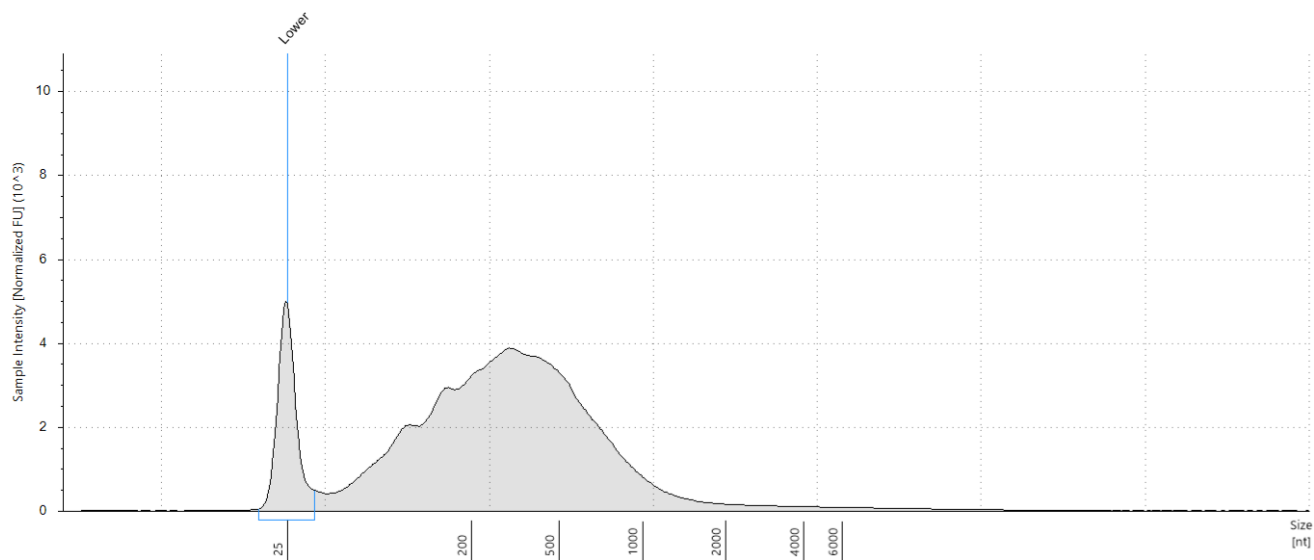


Sample Table

Well	RINe	28S/18S (Area)	Conc. [ng/μl]	Sample Description	Alert	Observations
A1	-	-	112	Ladder		Ladder

Peak Table

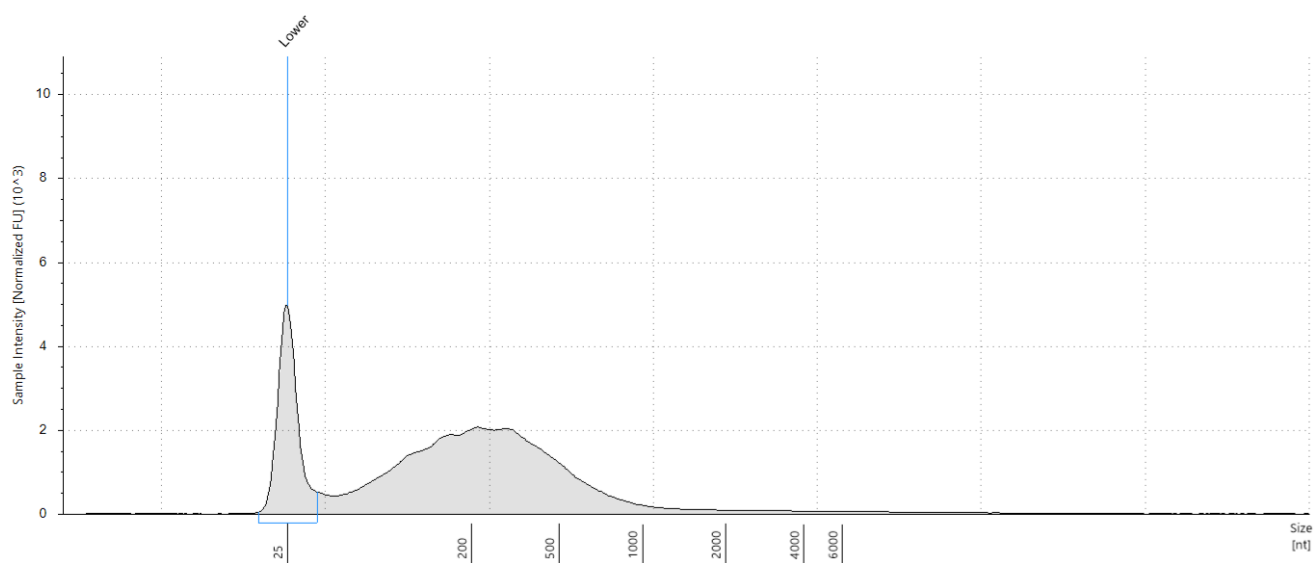
Size [nt]	Calibrated Conc. [ng/μl]	Assigned Conc. [ng/μl]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	40.0	40.0	4710	-		Lower Marker
200	9.61	-	141	9.55		
500	22.4	-	132	22.27		
1000	17.1	-	50.4	17.02		
2000	18.5	-	27.3	18.44		
4000	18.9	-	13.9	18.79		
6000	14.0	-	6.87	13.94		

B1: Mytilus RNA 1**Sample Table**

Well	RINe	28S/18S (Area)	Conc. [ng/μl]	Sample Description	Alert	Observations
B1	1.0	-	165	Mytilus RNA 1		

Peak Table

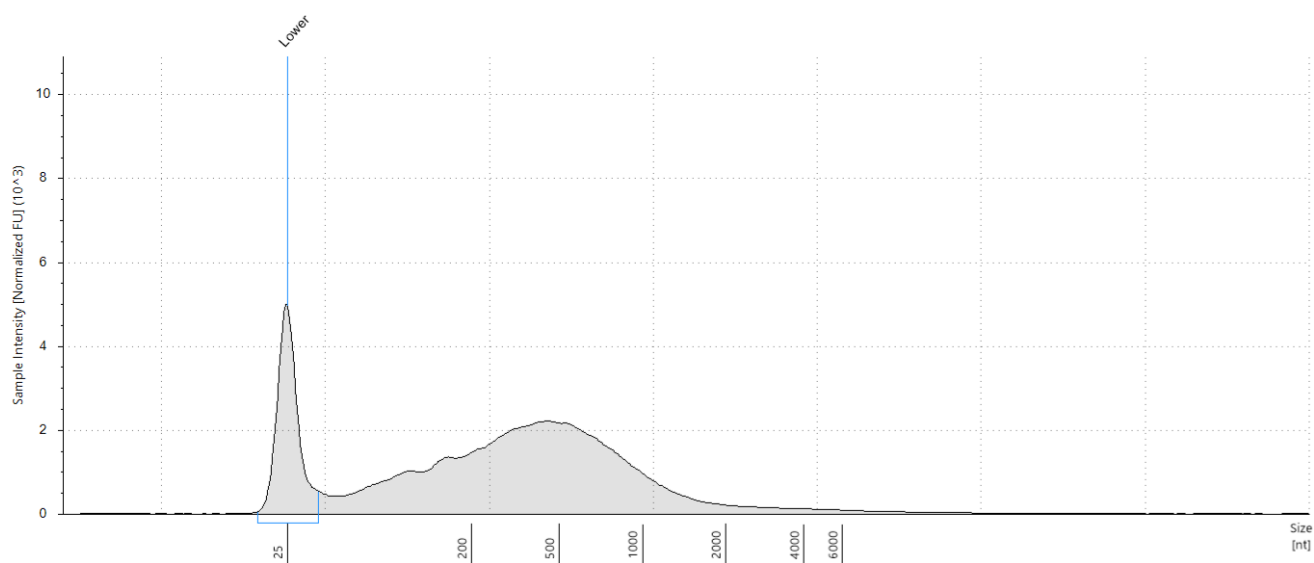
Size [nt]	Calibrated Conc. [ng/μl]	Assigned Conc. [ng/μl]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	40.0	40.0	4710	-		Lower Marker

C1: Mytilus RNA 2**Sample Table**

Well	RINe	28S/18S (Area)	Conc. [ng/μl]	Sample Description	Alert	Observations
C1	1.6	-	80.0	Mytilus RNA 2		

Peak Table

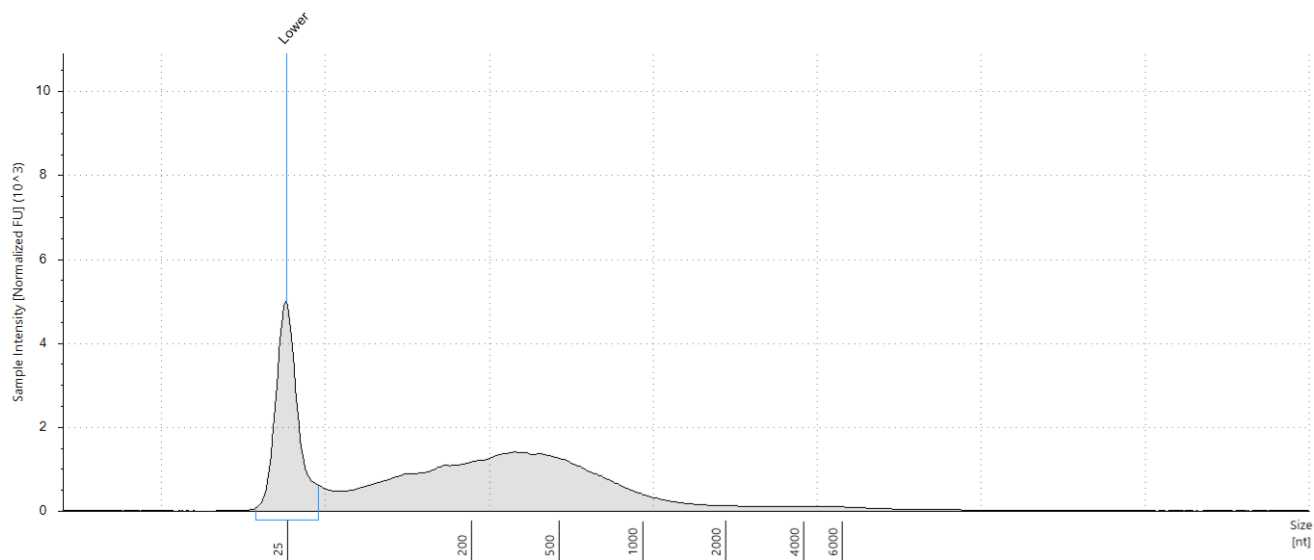
Size [nt]	Calibrated Conc. [ng/μl]	Assigned Conc. [ng/μl]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	40.0	40.0	4710	-		Lower Marker

D1: Mytilus RNA 3**Sample Table**

Well	RINe	28S/18S (Area)	Conc. [ng/μl]	Sample Description	Alert	Observations
D1	1.0	-	96.0	Mytilus RNA 3		

Peak Table

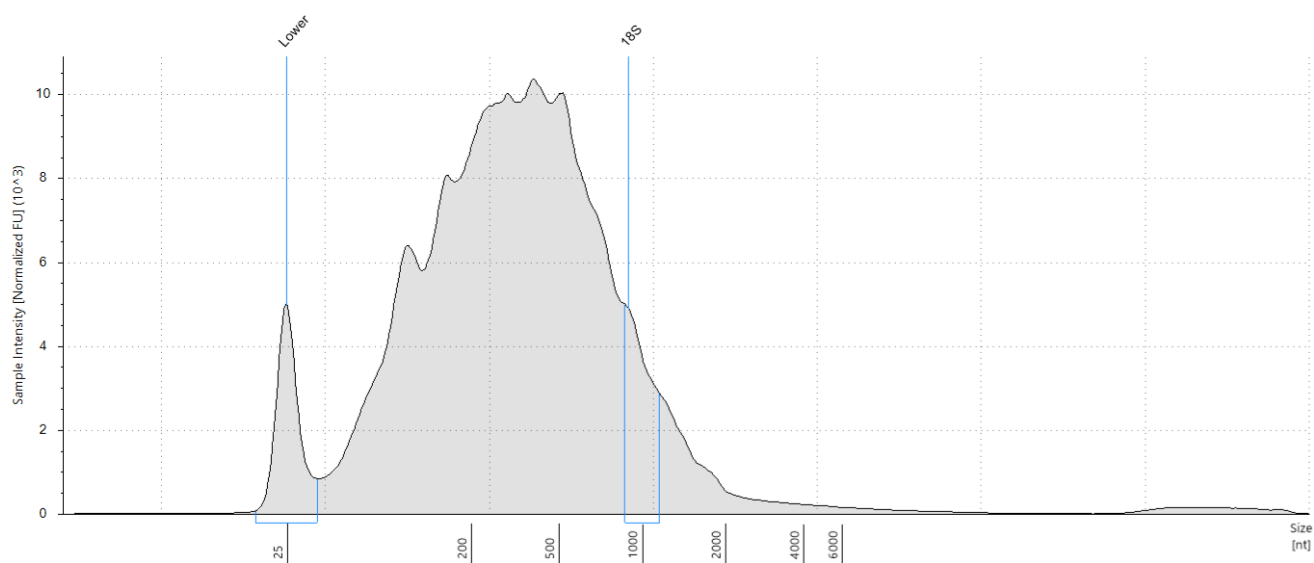
Size [nt]	Calibrated Conc. [ng/μl]	Assigned Conc. [ng/μl]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	40.0	40.0	4710	-		Lower Marker

E1: Mytilus RNA 4**Sample Table**

Well	RINe	28S/18S (Area)	Conc. [ng/μl]	Sample Description	Alert	Observations
E1	1.1	-	60.8	Mytilus RNA 4		

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

F1: Mytilus RNA 5**Sample Table**

Well	RINe	28S/18S (Area)	Conc. [ng/μl]	Sample Description	Alert	Observations
F1	1.6	-	411	Mytilus RNA 5		

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
892	24.2	-	79.8	100.00		18S