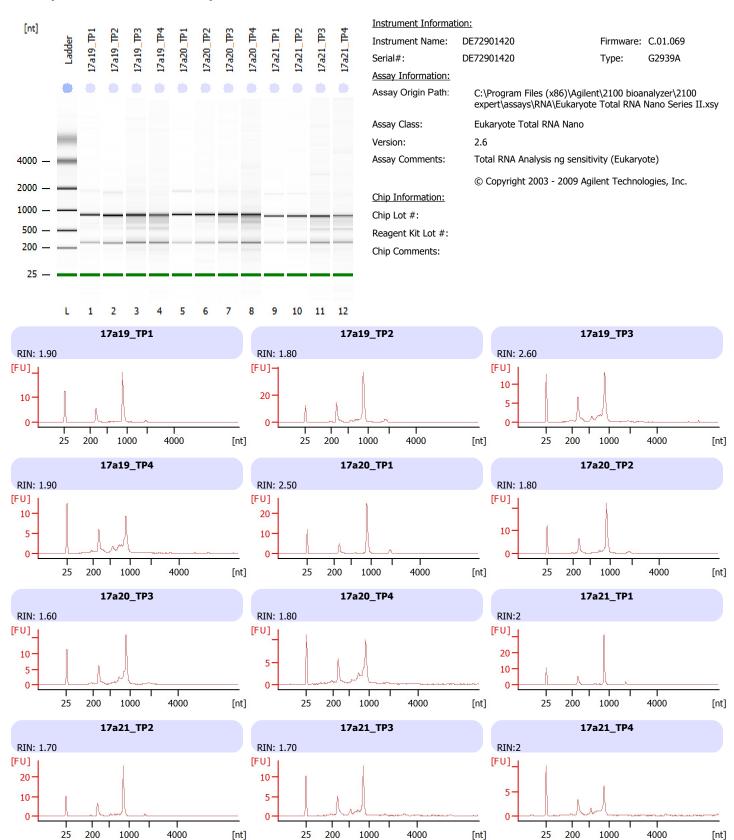
Electrophoresis File Run Summary



Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Stat Result Labe	el Result Color
17a19_TP1		✓ RIN: 1.90	
17a19_TP2		✓ RIN: 1.80	
17a19_TP3		RIN: 2.60	
17a19_TP4		RIN: 1.90	
17a20_TP1		RIN: 2.50	
17a20_TP2		RIN: 1.80	
17a20_TP3		RIN: 1.60	
17a20_TP4		RIN: 1.80	
17a21_TP1		RIN:2	
17a21_TP2		RIN: 1.70	
17a21_TP3		RIN: 1.70	
17a21_TP4		RIN:2	
Ladder		All Other	
		Samples	

Chip Lot # Reagent Kit Lot #

Chip Comments:

Electrophoresis Assay Details

General Analysis Settings

Number of Available Sample and Ladder Wells (Max.): 13

Minimum Visible Range [s]: 17

Maximum Visible Range [s]: 70

Start Analysis Time Range [s]: 19

End Analysis Time Range [s]: 69

Ladder Concentration [ng/µl]: 150

Lower Marker Concentration [ng/µl]: 0

Upper Marker Concentration [ng/µl]: 0

Used Lower Marker for Quantitation

Standard Curve Fit is Logarithmic

Show Data Aligned to Lower Marker

Integrator Settings

Integration Start Time [s]: 19
Integration End Time [s]: 69
Slope Threshold: 0.6
Height Threshold [FU]: 0.5
Area Threshold: 0.2
Width Threshold [s]: 0.5
Baseline Plateau [s]: 6

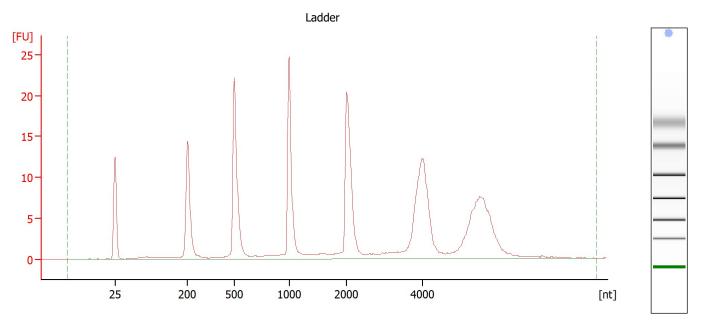
Filter Settings

Filter Width [s]: 0.5 Polynomial Order: 4

Ladder

Ladder Peak	Size
1	25
2	200
3	500
4	1000
5	2000
6	4000

Electropherogram Summary

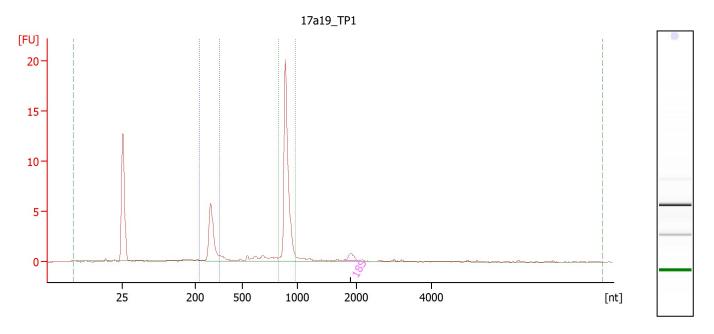


Overall Results for Ladder

RNA Area: 196.3 RNA Concentration: 150 ng/µl Result Flagging Color: Result Flagging Label:

All Other Samples

Electropherogram Summary Continued ...



Corr. Area 2:

23.8

Overall Results for sample 1: 17a19 TP1

RNA Integrity Number (RIN):

RNA Area: 43.1 Result Flagging Color:

RNA Concentration: 33 ng/µl Result Flagging Label: RIN: 1.90
rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 9.1

1.9 (B.02.11, Anomaly Threshold(s) manually adapted)

Fragment table for sample 1 : 17a19 TP1

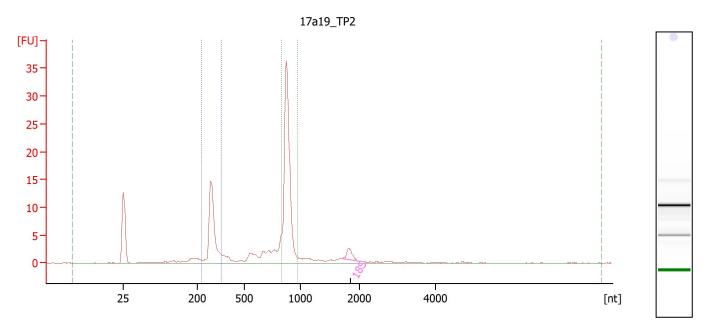
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,763
 2,060
 1.0
 2.2

Region table for sample 1: 17a19 TP1

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	9.1	21	
Region 2	825	975	23.8	55	

Electropherogram Summary Continued ...



Overall Results for sample 2: 17a19 TP2

 RNA Area:
 121.3
 Result Flagging Color:

 RNA Concentration:
 93 ng/μl
 Result Flagging Label:
 RIN: 1.80

 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 25.7

 RNA Integrity Number (RIN):
 1.8 (B.02.11,
 Corr. Area 2:
 54.6

Anomaly Threshold(s) manually adapted)

Fragment table for sample 2: 17a19 TP2

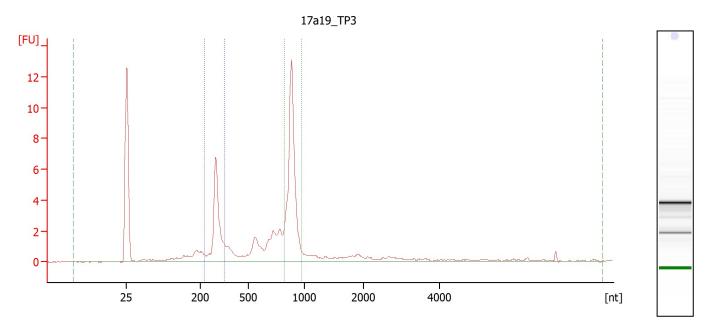
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,695
 2,037
 2.8
 2.3

Region table for sample 2: 17a19 TP2

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo
					r
Region 1	225	350	25.7	21	
Region 2	825	975	54.6	45	

Electropherogram Summary Continued ...



Overall Results for sample 3: 17a19 TP3

RNA Area: 63.1 Result Flagging Color:

RNA Concentration: 48 ng/µl Result Flagging Label: RIN: 2.60

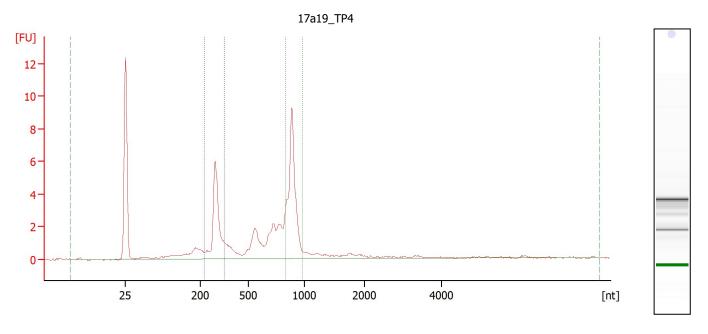
rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 12.1

RNA Integrity Number (RIN): 2.6 (B.02.11, Corr. Area 2: 21.6

Anomaly Threshold(s) manually adapted)

Region table for sample 3: <u>17a19_TP3</u>					
Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	12.1	19	
Region 2	825	975	21.6	34	

Electropherogram Summary Continued ...



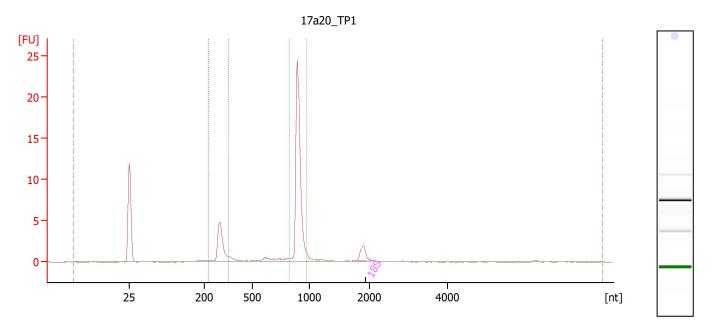
Overall Results for sample 4: 17a19 TP4

RNA Area: 53.8 Result Flagging Color: RNA Concentration: 41 ng/µl Result Flagging Label: RIN: 1.90 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 11.0 RNA Integrity Number (RIN): 1.9 (B.02.11, Corr. Area 2: 16.7

Anomaly Threshold(s) manually adapted)

Region table for sample 4: <u>17a19 TP4</u>					
Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	11.0	20	
Region 2	825	975	16.7	31	

Electropherogram Summary Continued ...



Overall Results for sample 5 : 17a20 TP1

 RNA Area:
 48.3
 Result Flagging Color:

 RNA Concentration:
 37 ng/μl
 Result Flagging Label:
 RIN: 2.50

 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 8.3

 RNA Integrity Number (RIN):
 2.5 (B.02.11,
 Corr. Area 2:
 28.6

Anomaly Threshold(s) manually adapted)

Fragment table for sample 5: 17a20 TP1

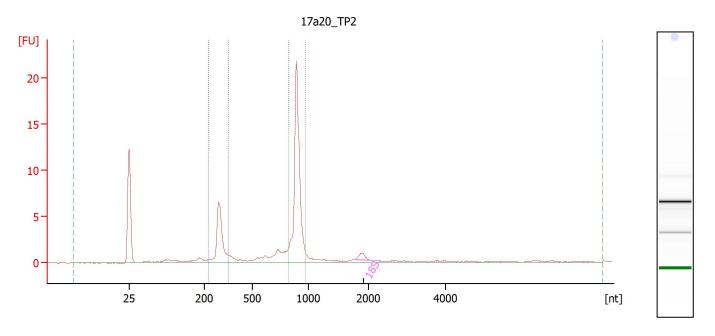
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,762
 2,199
 2.6
 5.4

Region table for sample 5: 17a20 TP1

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	8.3	17	
Region 2	825	975	28.6	59	

Electropherogram Summary Continued ...



Overall Results for sample 6: 17a20 TP2

RNA Area: 62.0 Result Flagging Color:

RNA Concentration: 47 ng/µl Result Flagging Label: RIN: 1.80

rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 11.2

RNA Integrity Number (RIN): 1.8 (B.02.11, Corr. Area 2: 29.5

Anomaly Threshold(s) manually adapted)

Fragment table for sample 6 : <u>17a20_TP2</u>

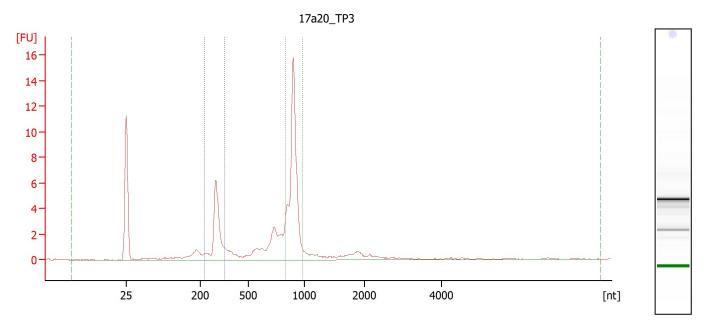
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,770
 2,129
 1.1
 1.8

Region table for sample 6: 17a20 TP2

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	11.2	18	
Region 2	825	975	29.5	48	

Electropherogram Summary Continued ...

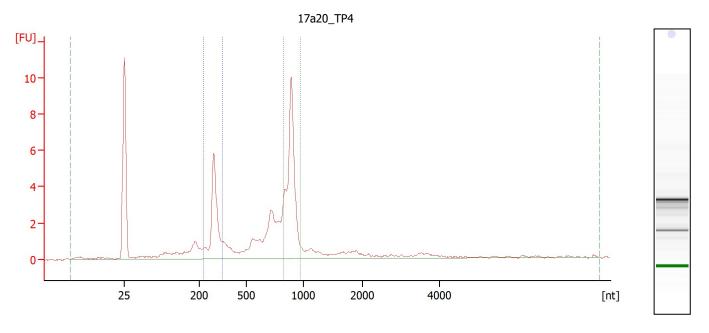


Overall Results for sample 7: 17a20 TP3

RNA Area: 64.1 Result Flagging Color: **RNA Concentration:** 49 ng/µl Result Flagging Label: RIN: 1.60 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 11.3 RNA Integrity Number (RIN): Corr. Area 2: 1.6 (B.02.11, 25.9 Anomaly Threshold(s) manually adapted)

Region table for sample 7: <u>17a20_TP3</u>						
Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r	
Region 1	225	350	11.3	18		
Region 2	825	975	25.9	40		

Electropherogram Summary Continued ...



Overall Results for sample 8: 17a20_TP4

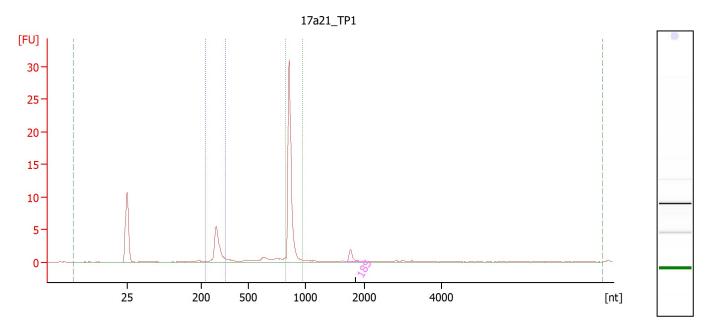
RNA Area: 62.3 Result Flagging Color: **RNA Concentration:** 48 ng/µl Result Flagging Label: RIN: 1.80 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 11.0 RNA Integrity Number (RIN): Corr. Area 2: 1.8 (B.02.11, 18.8 Anomaly Threshold(s)

Region table for sample 8: 17a20 TP4

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	11.0	18	
Region 2	825	975	18.8	30	

manually adapted)

Electropherogram Summary Continued ...



Overall Results for sample 9: 17a21 TP1

RNA Area: 51.6 Result Flagging Color:

RNA Concentration: 39 ng/µl Result Flagging Label: RIN:2

rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 9.2

RNA Integrity Number (RIN): 2 (B.02.11, Anomaly Corr. Area 2: 27.3

Threshold(s) manually adapted)

Fragment table for sample 9: 17a21 TP1

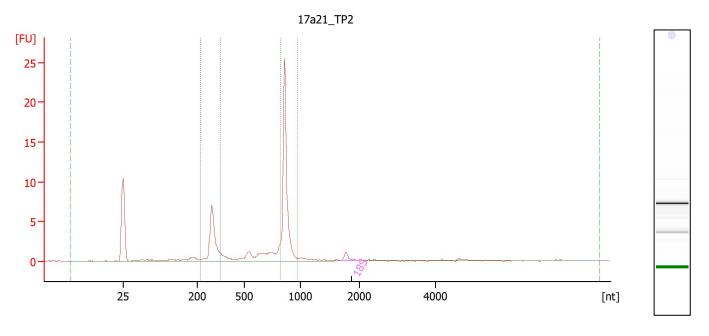
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,692
 2,028
 1.7
 3.2

Region table for sample 9: 17a21 TP1

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	9.2	18	
Region 2	825	975	27.3	53	

Electropherogram Summary Continued ...



Overall Results for sample 10 : 17a21 TP2

RNA Area: 57.3 Result Flagging Color:

RNA Concentration: 44 ng/µl Result Flagging Label: RIN: 1.70

rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 12.0

RNA Integrity Number (RIN): 1.7 (B.02.11, Corr. Area 2: 26.3

Anomaly Threshold(s) manually adapted)

Fragment table for sample 10: 17a21 TP2

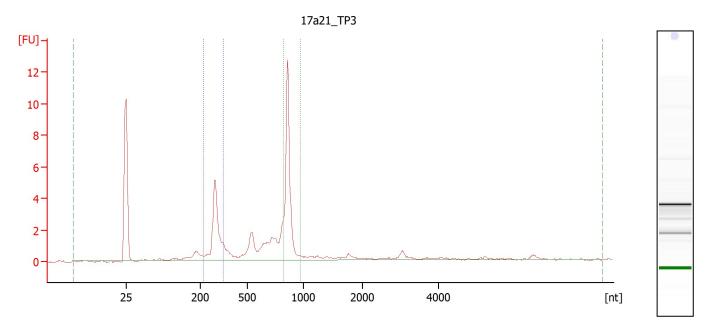
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,680
 2,081
 1.2
 2.1

Region table for sample 10: 17a21 TP2

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	12.0	21	
Region 2	825	975	26.3	46	

Electropherogram Summary Continued ...



32

Overall Results for sample 11: 17a21 TP3

RNA Area: 45.9 Result Flagging Color: **RNA Concentration:** 35 ng/µl Result Flagging Label: RIN: 1.70 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 9.5 RNA Integrity Number (RIN): Corr. Area 2: 1.7 (B.02.11, 14.7 Anomaly Threshold(s)

manually adapted)

14.7

Region table for sample 11: <u>17a21_TP3</u>					
Name	From [nt]	To [nt]	Corr. Area	% of Total	
Region 1	225	350	9.5	21	

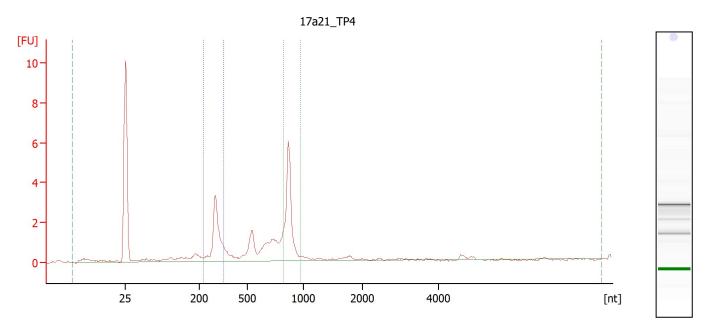
975

Region 2

825

Colo

Electropherogram Summary Continued ...



Overall Results for sample 12: 17a21 TP4

RNA Area:	31.4	Result Flagging Color:	
RNA Concentration:	24 ng/µl	Result Flagging Label:	RIN:2
rRNA Ratio [28s / 18s]:	0.0	Corr. Area 1:	6.4
RNA Integrity Number (RIN):	2 (B.02.11, Anomaly Threshold(s) manually	Corr. Area 2:	8.4

Region table for sample 12: 17a21 TP4 Name From [nt] To [nt] % of Total Corr. Area Colo Region 1 225 350 6.4 20 Region 2 825 975 8.4 27

adapted)

Assay Class: Data Path: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02_16-14-33.xad Created: Modified: 12/2/2020 4:14:32 PM 12/2/2020 5:06:42 PM **Gel Image** [nt] 17a19_TP2 17a19_TP3 17a19_TP4 17a20_TP1 ĕ 17a20_TP3 17a20_TP4 17a21_TP2 17a21_TP3 17a21_TP4 17a19_TP1 17a21_TP1 Ladder 17a20 ø • • • • ø • • ø ø • 4000 2000 1000 -500 200 25 3 1 5 8 10 11 12