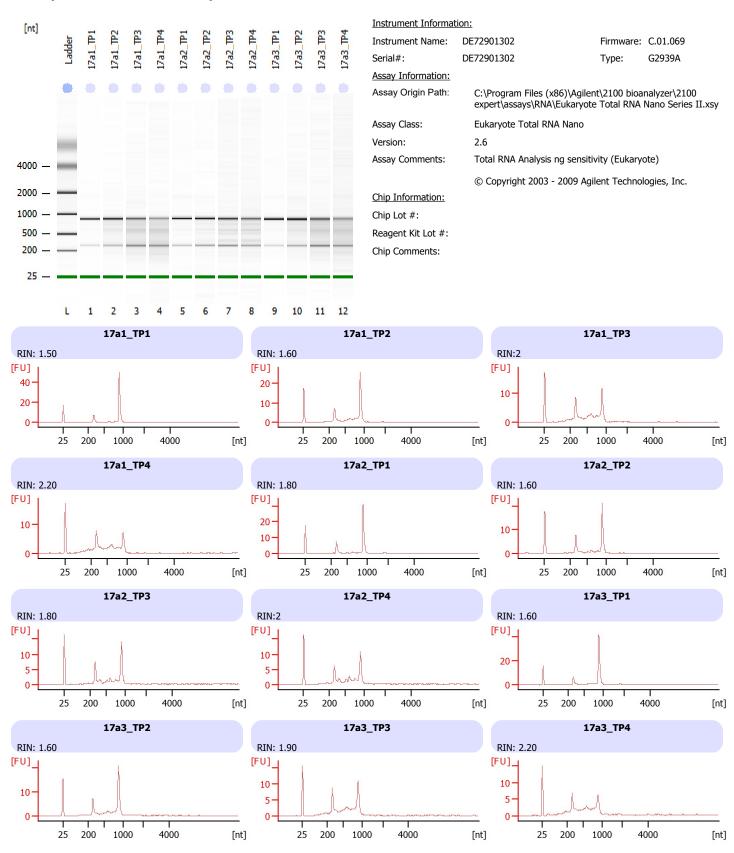
## **Electrophoresis File Run Summary**



Assay Class: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano\_2020-12-02\_14-31-30.xad Created: 12/2/2020 2:31:29 PM Modified: 12/2/2020 3:31:50 PM

## **Electrophoresis File Run Summary (Chip Summary)**

Sample Name	Sample Comment	Stat Result L us	abel Result Color
17a1_TP1		✓ RIN: 1.50	
17a1_TP2		✓ RIN: 1.60	
17a1_TP3		RIN:2	
17a1_TP4		✓ RIN: 2.20	
17a2_TP1		✓ RIN: 1.80	
17a2_TP2		✓ RIN: 1.60	
17a2_TP3		✓ RIN: 1.80	
17a2_TP4		RIN:2	
17a3_TP1		✓ RIN: 1.60	
17a3_TP2		✓ RIN: 1.60	
17a3_TP3		✓ RIN: 1.90	
17a3_TP4		✓ RIN: 2.20	
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

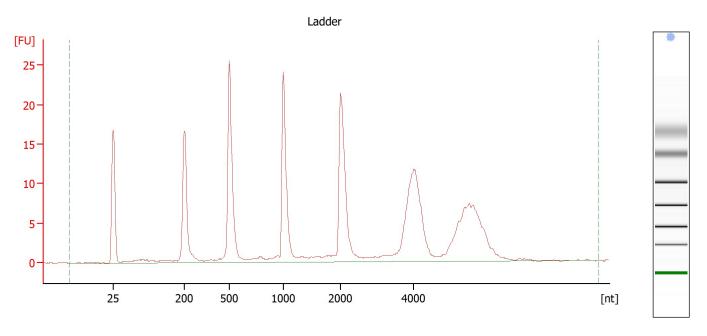
Ladder Peak	Size		
1	25		
2	200		
3	500		
4	1000		
5	2000		
6	4000		

Printed:

12/2/2020 3:32:10 PM

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano\_2020-12-02\_14-31-30.xad Created: 12/2/2020 2:31:29 PM Modified: 12/2/2020 3:31:50 PM

## **Electropherogram Summary**

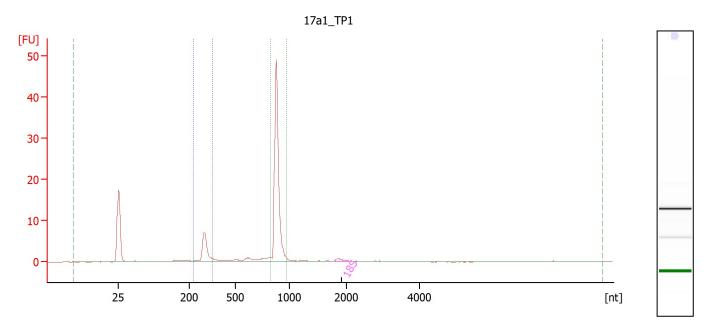


#### **Overall Results for Ladder**

RNA Area: 216.6 RNA Concentration: 150 ng/µl Result Flagging Color: Result Flagging Label:

All Other Samples

## **Electropherogram Summary Continued ...**



Overall Results for sample 1 : 17a1 TP1

RNA Area: 82.2 Result Flagging Color: RNA Concentration: 57 ng/µl Result Flagging Label: RIN: 1.50 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 11.5 RNA Integrity Number (RIN): 1.5 (B.02.11, Corr. Area 2: 54.2 Anomaly Threshold(s)

manually adapted)

Fragment table for sample 1 : <u>17a1\_TP1</u>

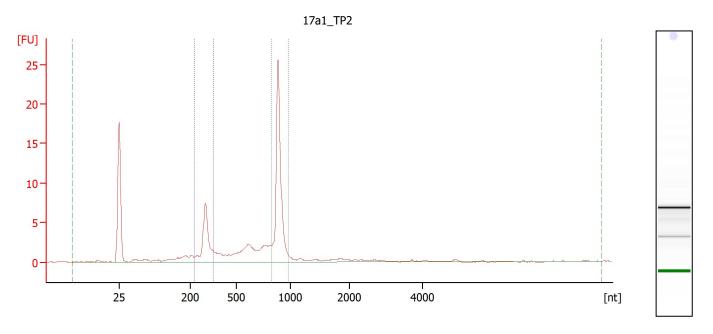
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,762
 2,090
 1.2
 1.5

Region table for sample 1: 17a1 TP1

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo	
					r	
Region 1	225	350	11.5	14		
Region 2	825	975	54.2	66		

## **Electropherogram Summary Continued ...**



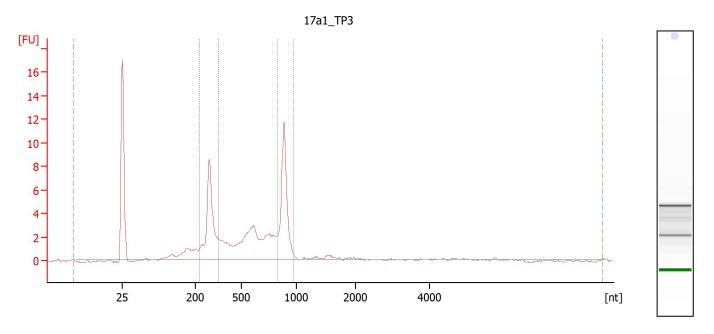
## Overall Results for sample 2: <u>17a1 TP2</u>

RNA Area: 84.9 Result Flagging Color: **RNA Concentration:** 59 ng/µl Result Flagging Label: RIN: 1.60 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 14.1 RNA Integrity Number (RIN): Corr. Area 2: 1.6 (B.02.11, 33.5 Anomaly Threshold(s)

Region table for sample 2: <u>17a1 TP2</u>

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	14.1	17	
Region 2	825	975	33.5	39	

## **Electropherogram Summary Continued ...**



Overall Results for sample 3: 17a1 TP3

RNA Area: 73.1 Result Flagging Color: **RNA Concentration:** 51 ng/µl Result Flagging Label: RIN:2 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 18.0 RNA Integrity Number (RIN): 2 (B.02.11, Anomaly Corr. Area 2: 18.1 Threshold(s) manually

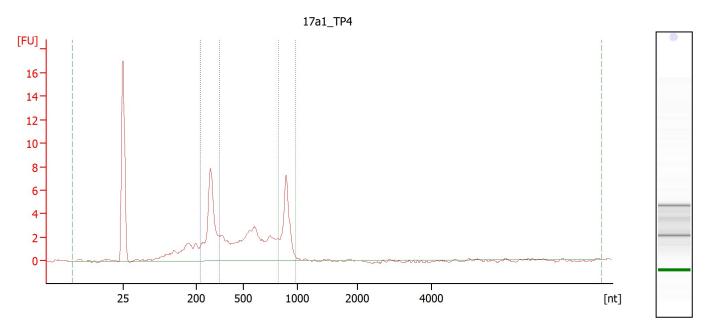
Region table for sample 3: 17a1 TP3

Name From [nt] To [nt] Cor

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	18.0	25	
Region 2	825	975	18.1	25	

adapted)

## **Electropherogram Summary Continued ...**



Overall Results for sample 4: <u>17a1 TP4</u>

RNA Area: 74.2 Result Flagging Color:

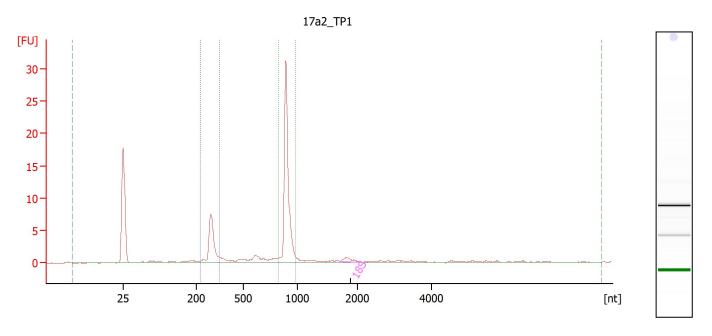
RNA Concentration: 51 ng/µl Result Flagging Label: RIN: 2.20
rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 18.4

RNA Integrity Number (RIN): 2.2 (B.02.11, Corr. Area 2: 12.5

Anomaly Threshold(s) manually adapted)

Region table for sample 4: <u>17a1 TP4</u>					
Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	18.4	25	
Region 2	825	975	12.5	17	

## **Electropherogram Summary Continued ...**



Overall Results for sample 5 : 17a2 TP1

 RNA Area:
 71.0
 Result Flagging Color:

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

 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 12.6

 RNA Integrity Number (RIN):
 1.8 (B.02.11,
 Corr. Area 2:
 34.9

Anomaly Threshold(s) manually adapted)

Fragment table for sample 5: 17a2 TP1

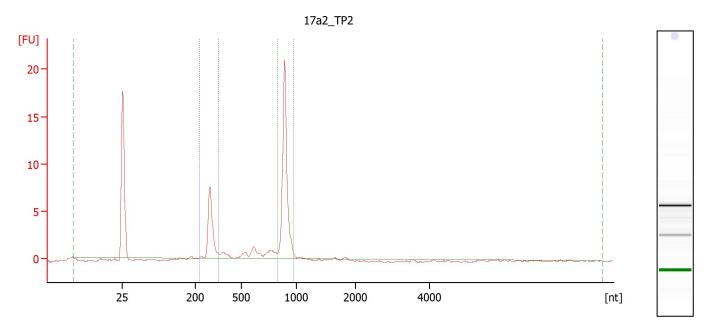
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,705
 2,146
 1.3
 1.8

Region table for sample 5: 17a2 TP1

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	12.6	18	
Region 2	825	975	34.9	49	

## **Electropherogram Summary Continued ...**



Overall Results for sample 6 : 17a2 TP2

RNA Area: 44.6 Result Flagging Color:

RNA Concentration: 31 ng/µl Result Flagging Label: RIN: 1.60

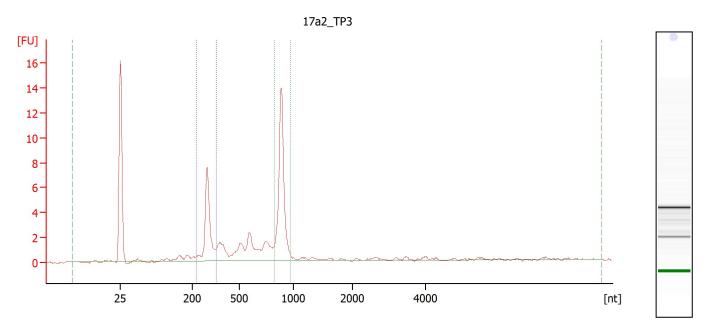
rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 11.3

RNA Integrity Number (RIN): 1.6 (B.02.11, Corr. Area 2: 25.3

Anomaly Threshold(s) manually adapted)

Region table for sample 6: <u>17a2_TP2</u>					
Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo
Region 1	225	350	11.3	25	
Region 2	825	975	25.3	57	

## **Electropherogram Summary Continued ...**



Overall Results for sample 7: 17a2 TP3

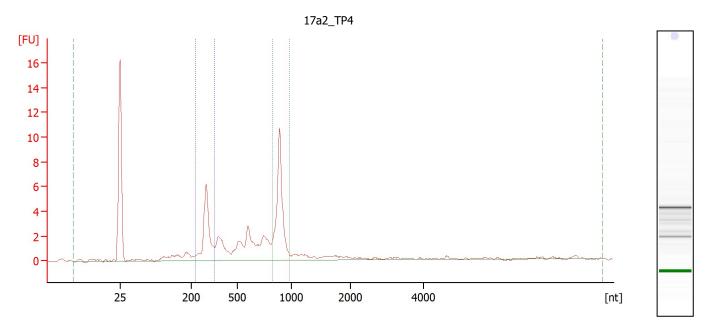
RNA Area: 56.0 Result Flagging Color: **RNA Concentration:** 39 ng/µl Result Flagging Label: RIN: 1.80 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 12.5 RNA Integrity Number (RIN): Corr. Area 2: 1.8 (B.02.11, 19.2 Anomaly Threshold(s)

Region table for sample 7: <u>17a2 TP3</u>

Name From [nt] To [nt] Corr.

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	12.5	22	
Region 2	825	975	19.2	34	

## **Electropherogram Summary Continued ...**



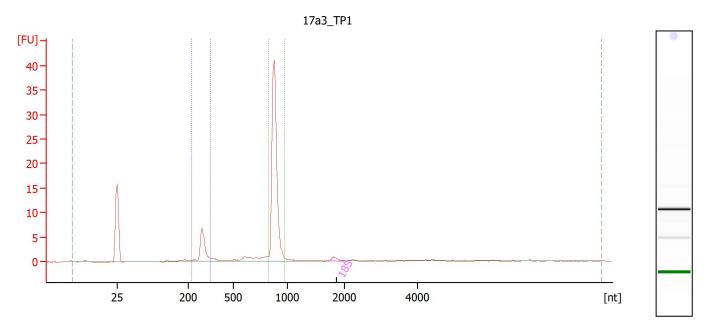
Overall Results for sample 8 : 17a2 TP4

RNA Area: 61.0 Result Flagging Color: **RNA Concentration:** 42 ng/µl Result Flagging Label: RIN:2 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 11.4 RNA Integrity Number (RIN): Corr. Area 2: 2 (B.02.11, Anomaly 16.1 Threshold(s) manually

adapted)

Region table for sample 8: <u>17a2_TP4</u>					
Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo
Region 1	225	350	11.4	19	
Region 2	825	975	16 1	26	

## **Electropherogram Summary Continued ...**



Overall Results for sample 9: 17a3 TP1

RNA Area: 75.7 Result Flagging Color: RNA Concentration: 52 ng/ $\mu$ l Result Flagging Label: RIN: 1.60 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 10.6 RNA Integrity Number (RIN): 1.6 (B.02.11, Corr. Area 2: 54.5

Anomaly Threshold(s) manually adapted)

Fragment table for sample 9 : <u>17a3\_TP1</u>

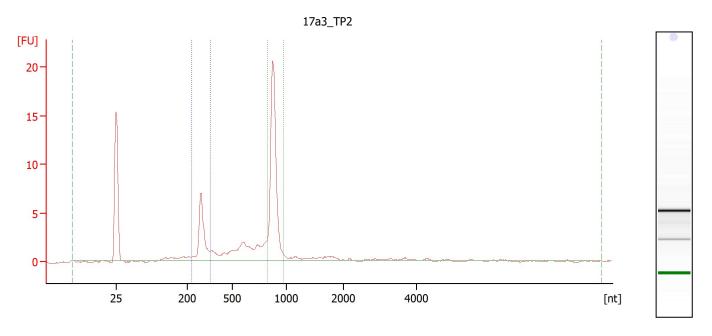
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,695
 2,016
 1.3
 1.7

Region table for sample 9: 17a3 TP1

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo	
					Г	
Region 1	225	350	10.6	14		
Region 2	825	975	54.5	72		

## **Electropherogram Summary Continued ...**



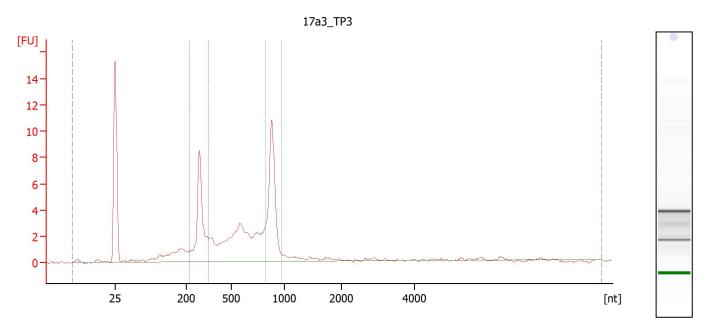
## Overall Results for sample 10 : 17a3 TP2

RNA Area: 67.5 Result Flagging Color: **RNA Concentration:** 47 ng/µl Result Flagging Label: RIN: 1.60 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 11.9 RNA Integrity Number (RIN): Corr. Area 2: 1.6 (B.02.11, 31.4 Anomaly Threshold(s)

Region table for sample 10: 17a3 TP2

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	11.9	18	
Region 2	825	975	31.4	46	

## **Electropherogram Summary Continued ...**



## Overall Results for sample 11: 17a3 TP3

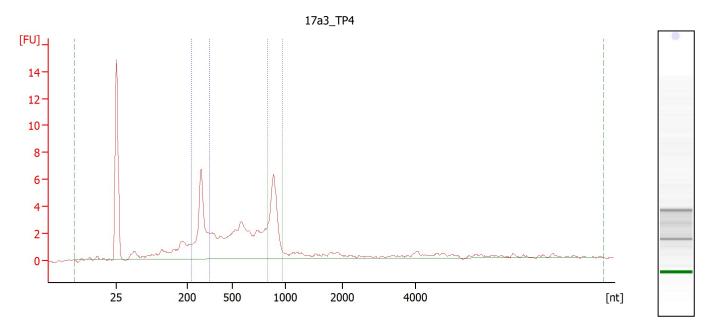
RNA Area: 82.7 Result Flagging Color: **RNA Concentration:** 57 ng/µl Result Flagging Label: RIN: 1.90 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 17.3 RNA Integrity Number (RIN): Corr. Area 2: 1.9 (B.02.11, 19.2 Anomaly Threshold(s)

Region table for sample 11 : 17a3 TP3

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	17.3	21	
Region 2	825	975	19.2	23	

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano\_2020-12-02\_14-31-30.xad Created: 12/2/2020 2:31:29 PM Modified: 12/2/2020 3:31:50 PM

## **Electropherogram Summary Continued ...**



# Overall Results for sample 12: 17a3 TP4

RNA Area: 82.8 Result Flagging Color: **RNA Concentration:** 57 ng/µl Result Flagging Label: RIN: 2.20 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 15.3 RNA Integrity Number (RIN): Corr. Area 2: 2.2 (B.02.11, 13.3 Anomaly Threshold(s)

Region table for sample 12: <u>17a3 TP4</u>

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	15.3	18	
Region 2	825	975	13.3	16	

Assay Class: Data Path: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano\_2020-12-02\_14-31-30.xad Created: Modified: 12/2/2020 2:31:29 PM 12/2/2020 3:31:50 PM **Gel Image** [nt] 17a2\_TP4 17a3\_TP4 17a1\_TP2 17a2\_TP1 17a3\_TP1 17a2\_TP3 17a3\_TP2 Ladder ۰ ۰ ٠ ۰ • ۰ ٠ • • ø 4000 2000 -1000 500 200 25 -2 3 5 10 11 12