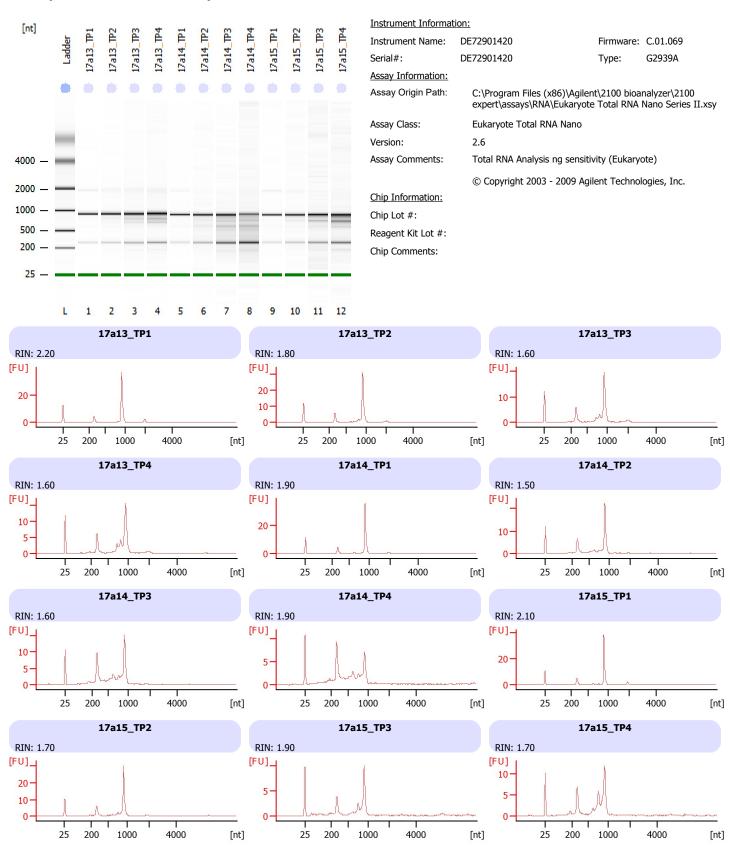
Assay Class: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano\_2020-12-02\_15-48-34.xad Created: 12/2/2020 3:48:34 PM Modified: 12/2/2020 4:25:46 PM

## **Electrophoresis File Run Summary**



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

Sample Name	Sample Comment	Stat Result Labe us	Result Color
17a13_TP1		RIN: 2.20	
17a13_TP2		RIN: 1.80	
17a13_TP3		RIN: 1.60	
17a13_TP4		RIN: 1.60	
17a14_TP1		RIN: 1.90	
17a14_TP2		RIN: 1.50	
17a14_TP3		RIN: 1.60	
17a14_TP4		RIN: 1.90	
17a15_TP1		RIN: 2.10	
17a15_TP2		RIN: 1.70	
17a15_TP3		RIN: 1.90	
17a15_TP4		RIN: 1.70	
Ladder		<ul><li>All Other Samples</li></ul>	

Chip Lot # Reagent Kit Lot #

**Chip Comments:** 

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano\_2020-12-02\_15-48-34.xad Created: 12/2/2020 3:48:34 PM Modified: 12/2/2020 4:25:46 PM

#### **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

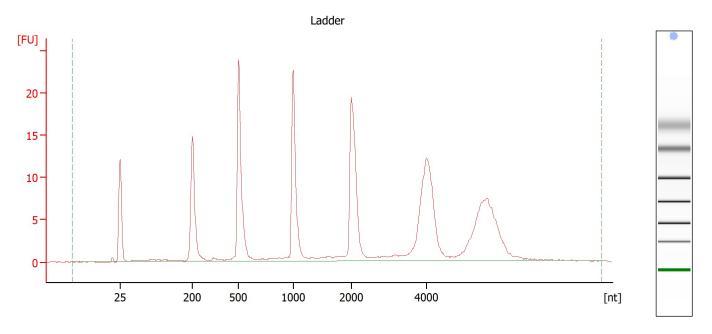
#### **Filter Settings**

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**

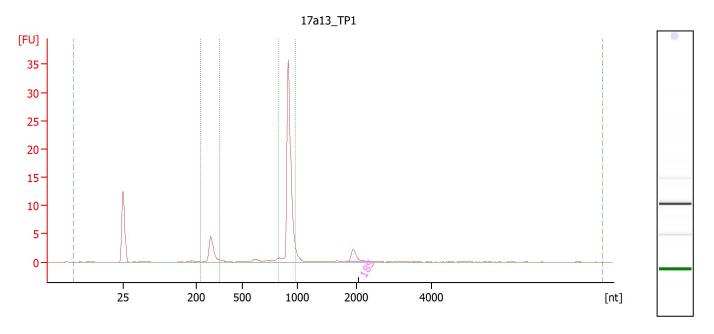


#### **Overall Results for Ladder**

RNA Area: 191.1 RNA Concentration: 150 ng/µl Result Flagging Color: Result Flagging Label:

All Other Samples

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



Overall Results for sample 1: 17a13 TP1

RNA Area: 60.9 Result Flagging Color: RNA Concentration: 48 ng/µl Result Flagging Label: RIN: 2.20 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 7.3 RNA Integrity Number (RIN): 2.2 (B.02.11, Corr. Area 2: 38.9 Anomaly Threshold(s)

manually adapted)

Fragment table for sample 1 : <u>17a13\_TP1</u>

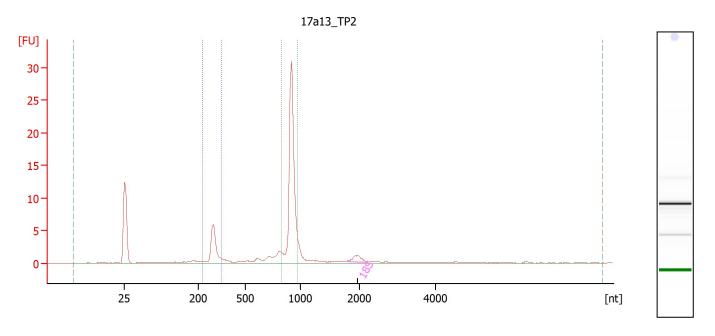
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,794
 2,430
 2.9
 4.8

Region table for sample 1: 17a13 TP1

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	7.3	12	
Region 2	825	975	38.9	64	

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



Overall Results for sample 2: <u>17a13\_TP2</u>

RNA Area: 77.4 Result Flagging Color: RNA Concentration: 61 ng/µl Result Flagging Label: RIN: 1.80 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 10.3 RNA Integrity Number (RIN): 1.8 (B.02.11, Corr. Area 2: 39.5 Anomaly Threshold(s)

manually adapted)

Fragment table for sample 2: <u>17a13\_TP2</u>

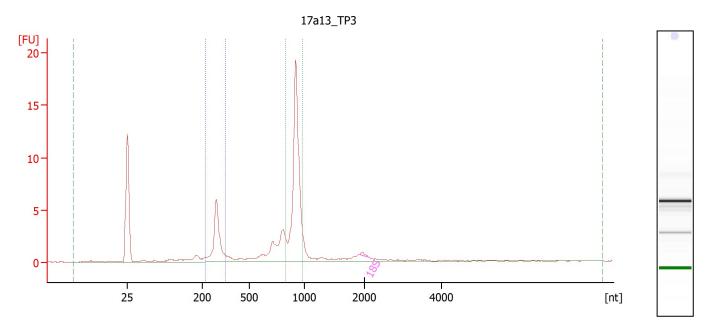
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,770
 2,234
 1.9
 2.4

Region table for sample 2: 17a13 TP2

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	10.3	13	
Region 2	825	975	39.5	51	

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



Overall Results for sample 3: 17a13 TP3

RNA Area: 65.4 Result Flagging Color: RNA Concentration: 51 ng/ $\mu$ l Result Flagging Label: RIN: 1.60 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 10.7 RNA Integrity Number (RIN): 1.6 (B.02.11, Corr. Area 2: 29.3

Anomaly Threshold(s) manually adapted)

Fragment table for sample 3: <u>17a13\_TP3</u>

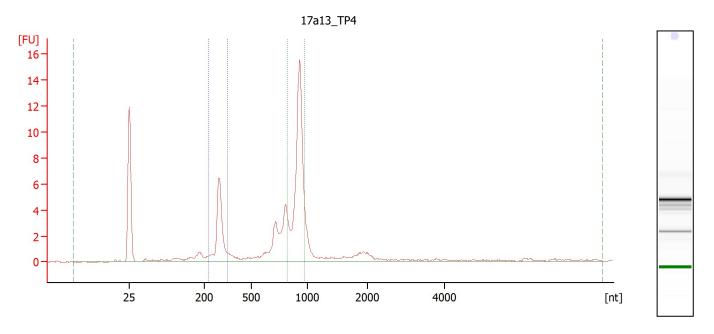
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,911
 2,127
 0.3
 0.5

Region table for sample 3: 17a13 TP3

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	10.7	16	
Region 2	825	975	29.3	45	

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

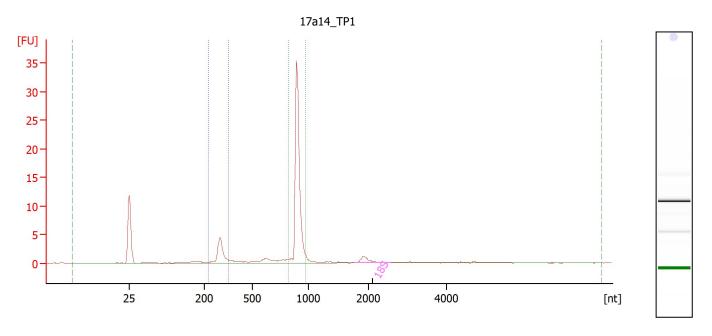


Overall Results for sample 4: 17a13 TP4

RNA Area: 70.9 Result Flagging Color: RNA Concentration: 56 ng/µl Result Flagging Label: RIN: 1.60 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 11.2 RNA Integrity Number (RIN): 1.6 (B.02.11, Corr. Area 2: 27.8 Anomaly Threshold(s)

Region table for sample 4: 17a13 TP4 Name From [nt] To [nt] Corr. Area % of Total Colo Region 1 225 350 11.2 16 Region 2 825 975 27.8 39

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



Overall Results for sample 5 : 17a14 TP1

RNA Area: 59.0 Result Flagging Color: RNA Concentration: 46 ng/µl Result Flagging Label: RIN: 1.90 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 7.7 RNA Integrity Number (RIN): 1.9 (B.02.11, Corr. Area 2: 37.5 Anomaly Threshold(s)

manually adapted)

Fragment table for sample 5: 17a14 TP1

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

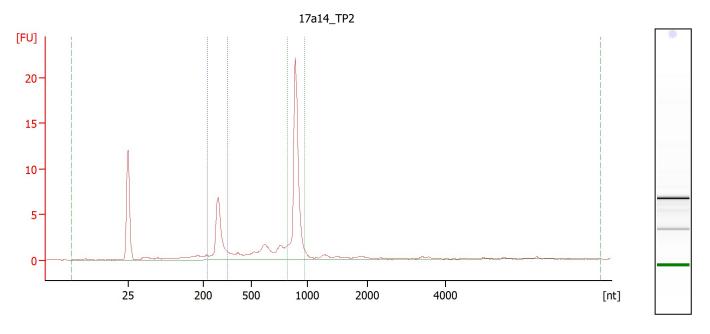
 18S
 1,757
 2,560
 2.3
 3.8

Region table for sample 5: 17a14 TP1

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	7.7	13	
Region 2	825	975	37.5	64	

Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano\_2020-12-02\_15-48-34.xad 12/2/2020 3:48:34 PM 12/2/2020 4:25:46 PM Assay Class: Created: Data Path: Modified:

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



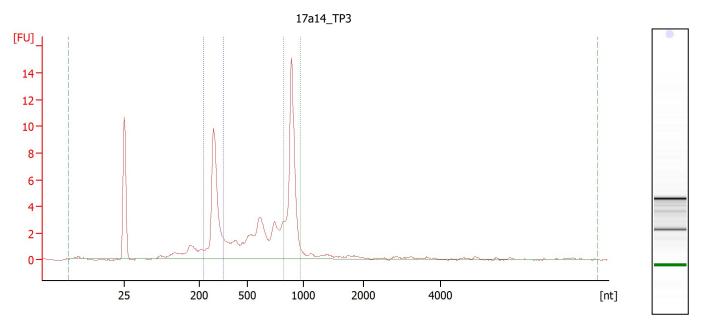
Overall Results for sample 6: 17a14\_TP2

RNA Area: 66.2 Result Flagging Color: **RNA Concentration:** 52 ng/µl Result Flagging Label: RIN: 1.50 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 12.4 RNA Integrity Number (RIN): Corr. Area 2: 1.5 (B.02.11, 27.8 Anomaly Threshold(s)

Region table for sample 6: 17a14 TP2

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	12.4	19	
Region 2	825	975	27.8	42	

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

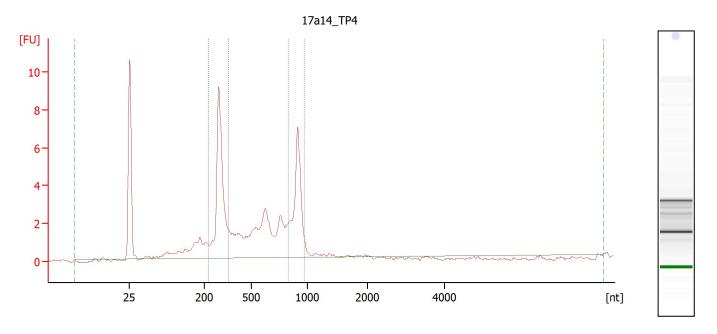


# Overall Results for sample 7: 17a14 TP3

RNA Area: 83.9 Result Flagging Color: **RNA Concentration:** 66 ng/µl Result Flagging Label: RIN: 1.60 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 19.4 RNA Integrity Number (RIN): Corr. Area 2: 1.6 (B.02.11, 23.5 Anomaly Threshold(s)

Region tab	le for sample 7	: <u>17a1</u>	<u>4_TP3</u>		
Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo
Region 1	225	350	19.4	23	•
Region 2	825	975	23.5	28	

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



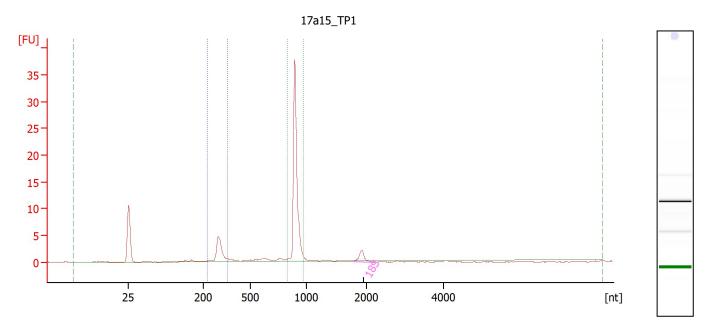
Overall Results for sample 8 : 17a14 TP4

RNA Area: 62.8 Result Flagging Color: **RNA Concentration:** 49 ng/µl Result Flagging Label: RIN: 1.90 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 18.0 RNA Integrity Number (RIN): Corr. Area 2: 1.9 (B.02.11, 12.4 Anomaly Threshold(s)

Region table for sample 8 : <u>17a14 TP4</u>

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	18.0	29	
Region 2	825	975	12.4	20	

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



Overall Results for sample 9: <u>17a15\_TP1</u>

RNA Area: 53.9 Result Flagging Color: RNA Concentration: 42 ng/ $\mu$ l Result Flagging Label: RIN: 2.10 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 8.0 RNA Integrity Number (RIN): 2.1 (B.02.11, Corr. Area 2: 36.7

Anomaly Threshold(s) manually adapted)

Fragment table for sample 9: 17a15 TP1

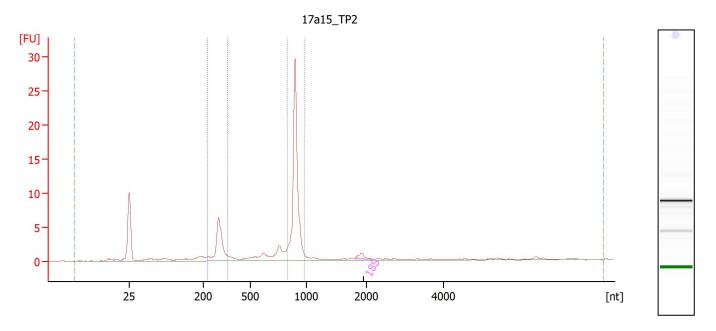
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,726
 2,254
 2.7
 5.0

Region table for sample 9: 17a15 TP1

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo
Region 1	225	350	8.0	15	
Region 2	825	975	36.7	68	

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



Overall Results for sample 10 : 17a15 TP2

RNA Area: 77.5 Result Flagging Color: RNA Concentration: 61 ng/µl Result Flagging Label: RIN: 1.70 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 12.0 RNA Integrity Number (RIN): 1.7 (B.02.11, Corr. Area 2: 36.2 Anomaly Threshold(s)

manually adapted)

Fragment table for sample 10 : 17a15 TP2

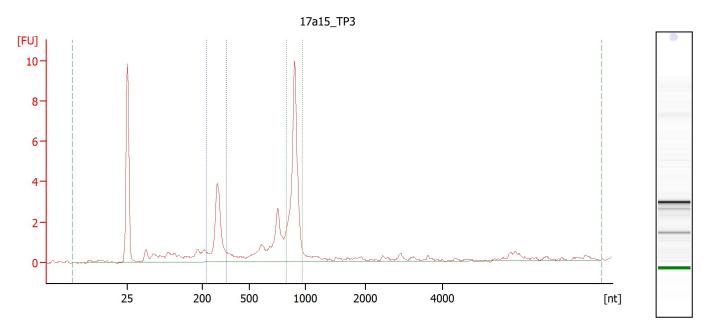
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,778
 2,170
 1.4
 1.8

Region table for sample 10: 17a15 TP2

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	12.0	16	
Region 2	825	975	36.2	47	

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



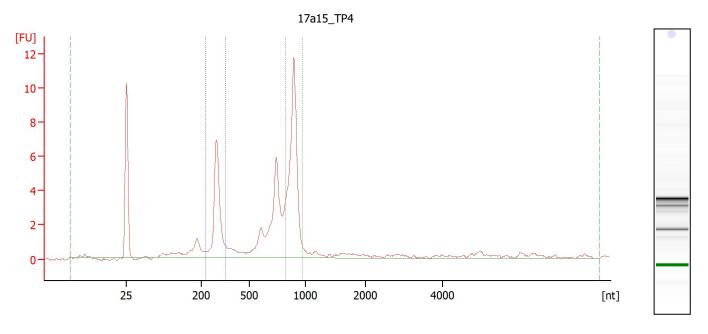
# Overall Results for sample 11: 17a15 TP3

RNA Area: 48.8 Result Flagging Color: **RNA Concentration:** 38 ng/µl Result Flagging Label: RIN: 1.90 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 8.0 RNA Integrity Number (RIN): Corr. Area 2: 1.9 (B.02.11, 15.6 Anomaly Threshold(s)

Region t	table for sa	mple 11	:	<u>17a15</u>	<u>TP3</u>
	_			_	_

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	8.0	16	
Region 2	825	975	15.6	32	

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



# Overall Results for sample 12: 17a15 TP4

RNA Area: 68.9 Result Flagging Color: **RNA Concentration:** 54 ng/µl Result Flagging Label: RIN: 1.70 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 12.6 RNA Integrity Number (RIN): Corr. Area 2: 1.7 (B.02.11, 21.8 Anomaly Threshold(s)

Region table for sample 12: <u>17a15\_TP4</u>

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	12.6	18	
Region 2	825	975	21.8	32	

Assay Class: Data Path: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano\_2020-12-02\_15-48-34.xad Created: Modified: 12/2/2020 3:48:34 PM 12/2/2020 4:25:46 PM **Gel Image** [nt] 17a13\_TP2 17a13\_TP3 17a13\_TP4 17a14\_TP3 17a14\_TP4 17a15\_TP2 17a15\_TP3 17a15\_TP4 17a14\_TP1 17a13\_TP1 17a15\_TP1 Ladder ø • • • • ø • • ø ø ø ۰ 4000 -2000 1000 -500 200 25 1 3 5 10 11 12