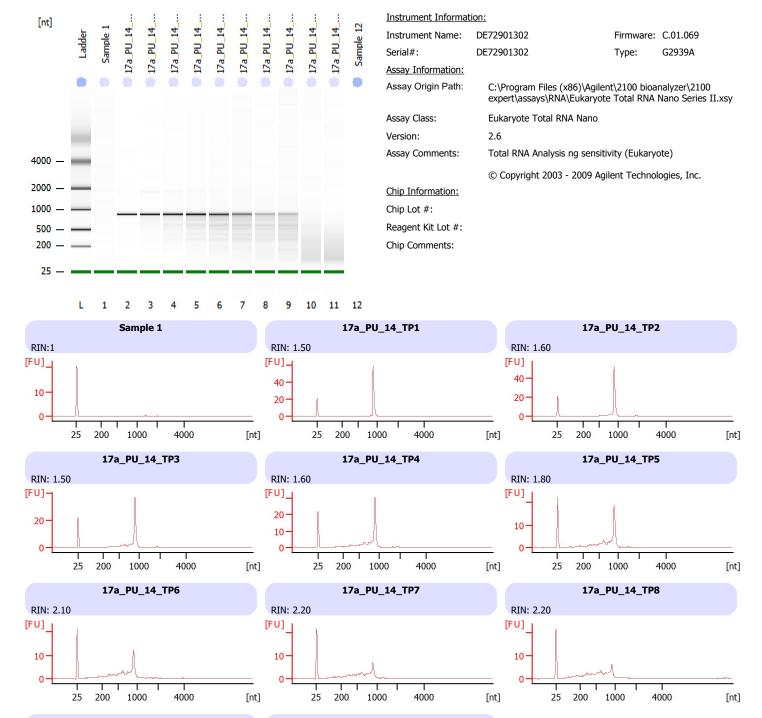
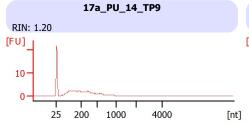
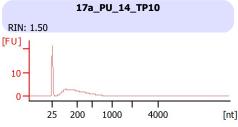
Assay Class: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad Created: 12/10/2020 3:52:31 PM Modified: 12/10/2020 4:23:30 PM

### **Electrophoresis File Run Summary**







Assay Class: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\Eukaryote

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

Sample Name	Sample Comment	Stat Result Labous	el Result Color
Sample 1		RIN:1	
17a_PU_14_TP1		RIN: 1.50	
17a_PU_14_TP2		RIN: 1.60	
17a_PU_14_TP3		RIN: 1.50	
17a_PU_14_TP4		RIN: 1.60	
17a_PU_14_TP5		RIN: 1.80	
17a_PU_14_TP6		RIN: 2.10	
17a_PU_14_TP7		RIN: 2.20	
17a_PU_14_TP8		RIN: 2.20	
17a_PU_14_TP9		RIN: 1.20	
17a_PU_14_TP10		RIN: 1.50	
Sample 12			
Ladder		All Other	
		Samples	

Chip Lot # Reagent Kit Lot #

**Chip Comments:** 

Printed: 12/10/2020 4:23:49 PM

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad Created: 12/10/2020 3:52:31 PM Modified: 12/10/2020 4:23:30 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**

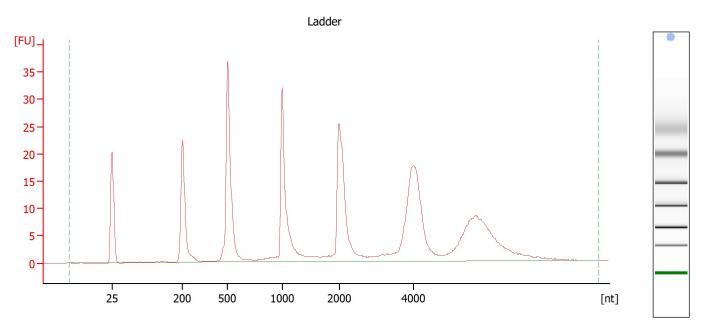
Filter Width [s]: 0.5 Polynomial Order: 4

#### Ladder

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

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\Eukaryote

### **Electropherogram Summary**

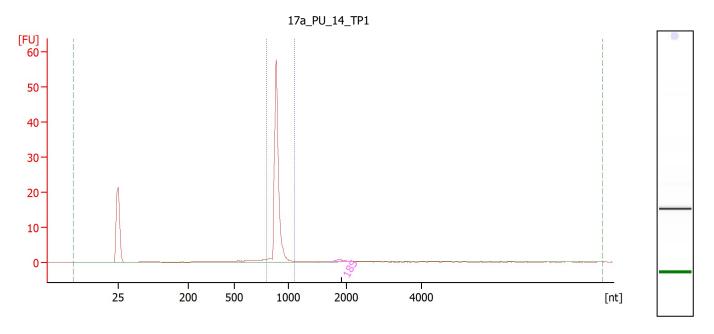


#### **Overall Results for Ladder**

RNA Area: 312.4 RNA Concentration: 150 ng/µl Result Flagging Color: Result Flagging Label:

All Other Samples

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 1.50

64.9

Overall Results for sample 2 : 17a PU 14 TP1

RNA Area: 77.6 RNA Concentration: 37 ng/µl

rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN): 1.5 (B.02.11,

Anomaly Threshold(s) manually adapted)

Fragment table for sample 2 : 17a PU 14 TP1

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

18S 1,724 2,175 1.2 1.6

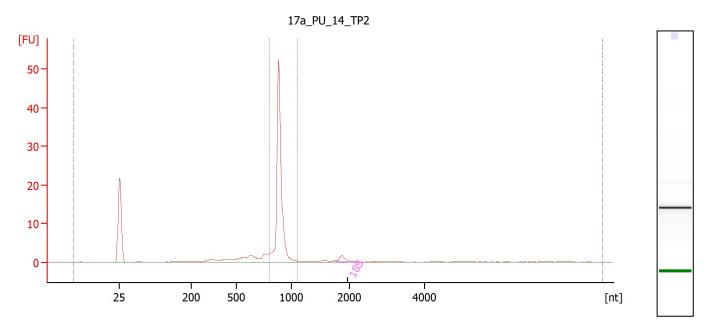
Region table for sample 2: <u>17a PU 14 TP1</u>

 Name
 From [nt]
 To [nt]
 Corr. Area
 % of Total
 Color

 Region 1
 800
 1,100
 64.9
 84

Assay Class: 12/10/2020 3:52:31 PM Eukaryote Total RNA Nano Created: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad Modified: 12/10/2020 4:23:30 PM Data Path:

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 1.60

66.5

Overall Results for sample 3: 17a PU 14 TP2

RNA Area: 96.4 RNA Concentration: 46 ng/µl

rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN): 1.6 (B.02.11, Anomaly Threshold(s)

manually adapted)

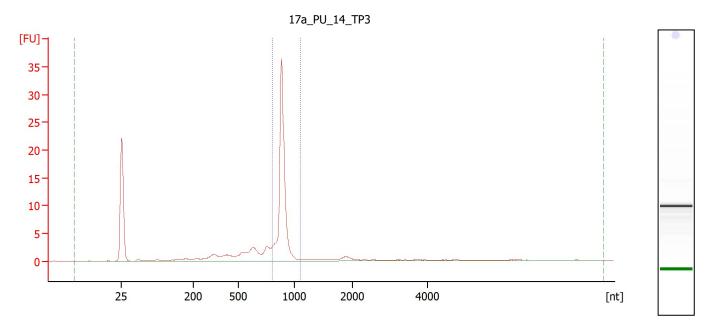
Fragment table for sample 3: 17a PU 14 TP2

Name Start Size [nt] End Size [nt] **Area** % of total Area 2.1

**18S** 1,765 2,278 2.0

Region table for sample 3: 17a PU 14 TP2 From [nt] Corr. Area % of Total Name To [nt] Colo Region 1 800 1,100 66.5 69

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



Overall Results for sample 4: 17a PU 14 TP3

RNA Area: 85.7
RNA Concentration: 41 ng/µl

0.0

Result Flagging Color: Result Flagging Label:

Corr. Area 1:

RIN: 1.50 50.4

RNA Integrity Number (RIN): 1.5

rRNA Ratio [28s / 18s]:

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

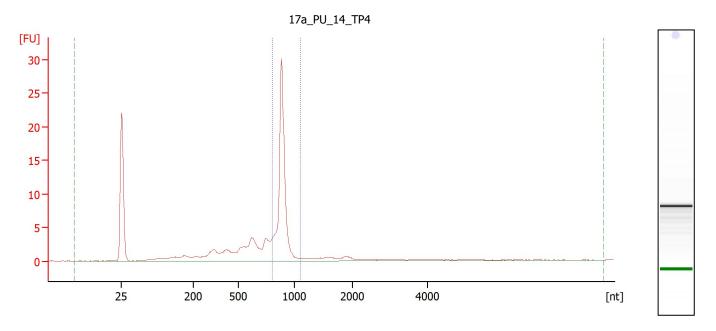
Region table for sample 4: 17a PU 14 TP3

Name From [nt] To [nt] Corr. Are:

 Name
 From [nt]
 To [nt]
 Corr. Area
 % of Total
 Color

 Region 1
 800
 1,100
 50.4
 59

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



Overall Results for sample 5 : 17a PU 14 TP4

 RNA Area:
 104.0
 Result Flagging Color:

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

 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 47.1

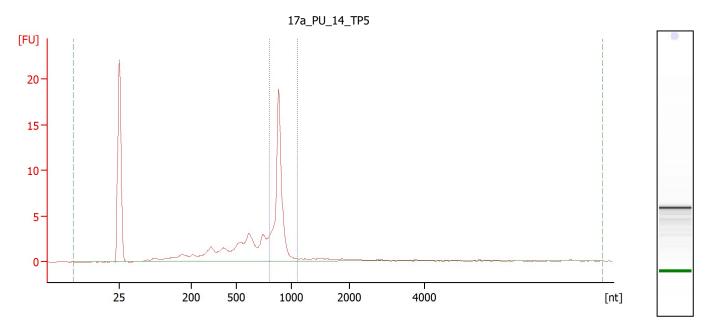
RNA Integrity Number (RIN): 1.6 (B.02.11, Anomaly Threshold(s) manually adapted)

 Region table for sample 5:
 17a PU 14 TP4

 Name
 From [nt]
 To [nt]
 Corr. Area
 % of Total
 Color

 Region 1
 800
 1,100
 47.1
 45

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



RIN: 1.80

Overall Results for sample 6 : 17a PU 14 TP5

RNA Area: 77.3 Result Flagging Color: RNA Concentration: 37 ng/µl Result Flagging Label:

rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 31.9

RNA Integrity Number (RIN): 1.8 (B.02.11, Anomaly Threshold(s)

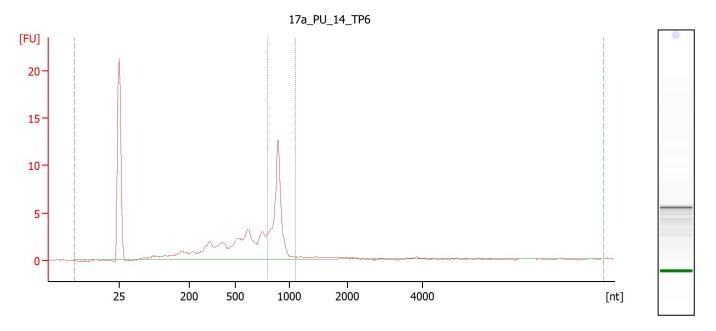
manually adapted)

 Region table for sample 6:
 17a
 PU
 14
 TP5

 Name
 From [nt]
 To [nt]
 Corr. Area
 % of Total
 Color

 Region 1
 800
 1,100
 31.9
 41

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



# Overall Results for sample 7: 17a PU 14 TP6

RNA Area: 69.7 Result Flagging Color: RNA Concentration: 33 ng/µl Result Flagging Label: RIN: 2.10 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 24.2

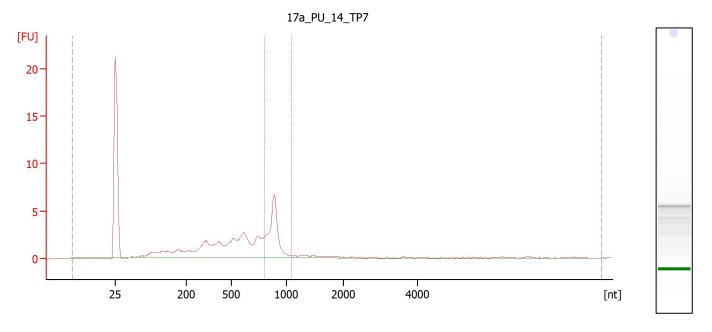
RNA Integrity Number (RIN): 2.1 (B.02.11, Anomaly Threshold(s) manually adapted)

 Region table for sample 7:
 17a PU 14 TP6

 Name
 From [nt]
 To [nt]
 Corr. Area
 % of Total
 Color

 Region 1
 800
 1,100
 24.2
 35

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



Overall Results for sample 8 : 17a PU 14 TP7

RNA Area: 61.8 Result Flagging Color: RNA Concentration: 30 ng/µl Result Flagging Label: RIN: 2.20

rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 15.5

RNA Integrity Number (RIN): 2.2 (B.02.11, Anomaly Threshold(s) manually adapted)

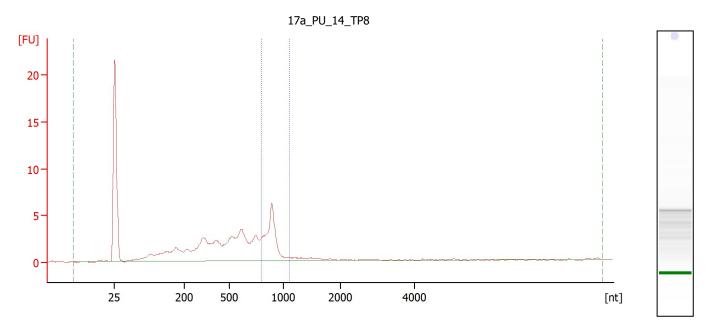
 Region table for sample 8 : 17a PU 14 TP7

 Name
 From [nt]
 To [nt]
 Corr. Area
 % of Total
 Color

 Region 1
 800
 1,100
 15.5
 25

Assay Class: Eukaryote Total RNA Nano 12/10/2020 3:52:31 PM Created: 12/10/2020 4:23:30 PM C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad Data Path: Modified:

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



Overall Results for sample 9: 17a PU 14 TP8

RNA Area: 78.3 Result Flagging Color: **RNA Concentration:** 38 ng/µl Result Flagging Label: RIN: 2.20 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 15.7

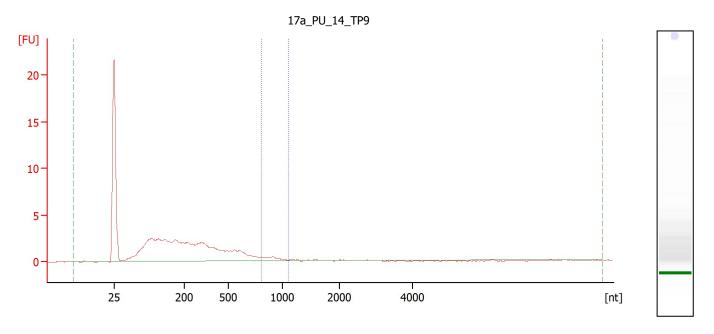
RNA Integrity Number (RIN): 2.2 (B.02.11, Anomaly Threshold(s)

manually adapted)

Region table for sample 9: 17a PU 14 TP8 Name From [nt] To [nt] Corr. Area % of Total Colo Region 1 800 1,100 15.7 20

Assay Class: Eukaryote Total RNA Nano 12/10/2020 3:52:31 PM Created: 12/10/2020 4:23:30 PM Data Path: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad Modified:

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



#### Overall Results for sample 10: 17a PU 14 TP9

RNA Area: 65.0 Result Flagging Color: RNA Concentration: 31 ng/µl Result Flagging Label: RIN: 1.20 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 1.9

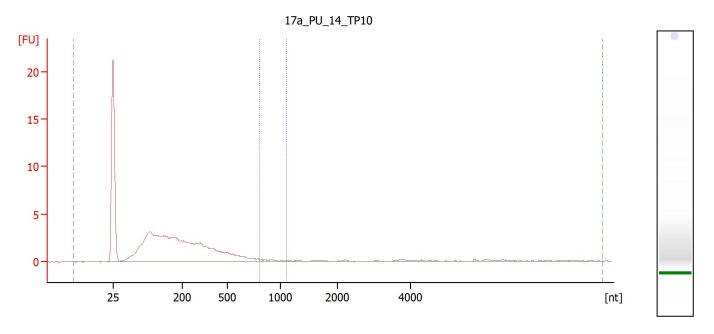
RNA Integrity Number (RIN): 1.2 (B.02.11, Anomaly Threshold(s) manually adapted)

Region table for sample 10: 17a PU 14 TP9 Name From [nt] To [nt] Corr. Area

% of Total Colo Region 1 800 1,100 1.9 3

Assay Class: Eukaryote Total RNA Nano Created: 12/10/2020 3:52:31 PM 12/10/2020 4:23:30 PM C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad Modified: Data Path:

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



#### Overall Results for sample 11: 17a PU 14 TP10

RNA Area: 69.8 Result Flagging Color: **RNA Concentration:** 34 ng/µl Result Flagging Label: RIN: 1.50 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 8.0

RNA Integrity Number (RIN): 1.5 (B.02.11, Anomaly Threshold(s)

manually adapted)

#### Region table for sample 11: 17a PU 14 TP10

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo
					r
Region 1	800	1.100	0.8	1	

Printed: 12/10/2020 4:23:49 PM

Assay Class: Data Path: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-52-31.xad Created: Modified: 12/10/2020 3:52:31 PM 12/10/2020 4:23:30 PM **Gel Image** 17a\_PU\_14\_... 17a\_PU\_14\_... 17a\_PU\_14\_... 17a\_PU\_14\_... 17a PU 14 ... 17a\_PU\_14\_... 17a PU 14 ... 17a\_PU\_14\_... 17a\_PU\_14\_... 17a PU\_14 ... [nt] Sample 12 Ladder Sample 1 . 4000 -2000 -1000 500 200 25 3 5 6 8 10 11 12