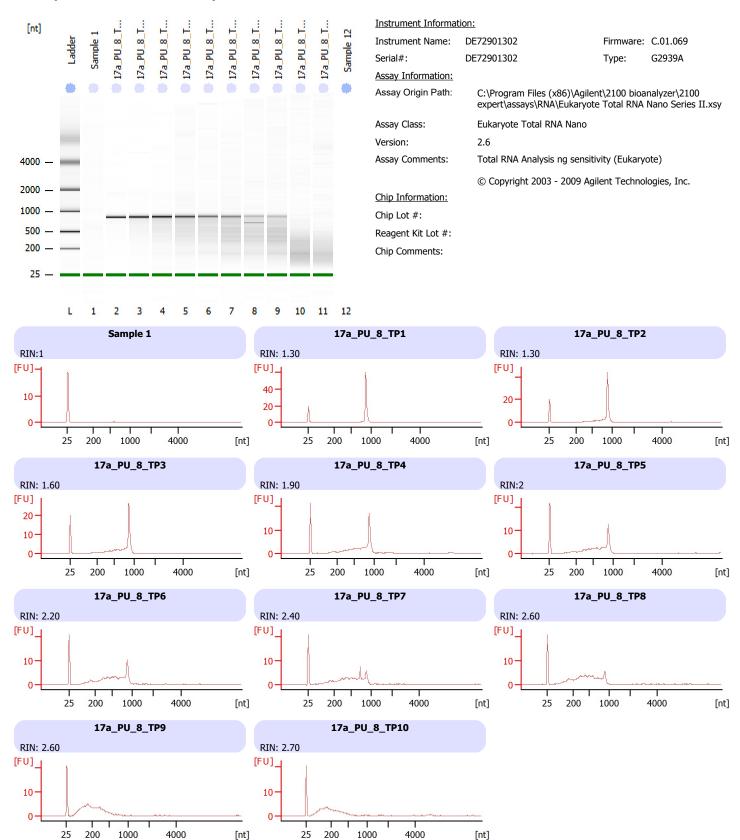
Assay Class: Eukaryote Total RNA Nano Created: 12/10/2020 12:35:05 PM Data Path: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_12-35-05.xad Created: 12/10/2020 3:02:51 PM

### **Electrophoresis File Run Summary**



Assay Class: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_12-35-05.xad Created: 12/10/2020 12:35:05 PM Modified: 12/10/2020 3:02:51 PM

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

Sample Name	Sample Comment	Stat us	Result Label	Result Color
Sample 1		~	RIN:1	
17a_PU_8_TP1		~	RIN: 1.30	
17a_PU_8_TP2		~	RIN: 1.30	
17a_PU_8_TP3		~	RIN: 1.60	
17a_PU_8_TP4		~	RIN: 1.90	
17a_PU_8_TP5		~	RIN:2	
17a_PU_8_TP6		~	RIN: 2.20	
17a_PU_8_TP7		~	RIN: 2.40	
17a_PU_8_TP8		~	RIN: 2.60	
17a_PU_8_TP9		~	RIN: 2.60	
17a_PU_8_TP10		~	RIN: 2.70	
Sample 12				
Ladder		~	All Other	
			Samples	

Chip Lot # Reagent Kit Lot #

**Chip Comments:** 

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_12-35-05.xad Created: 12/10/2020 12:35:05 PM Modified: 12/10/2020 3:02:51 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**

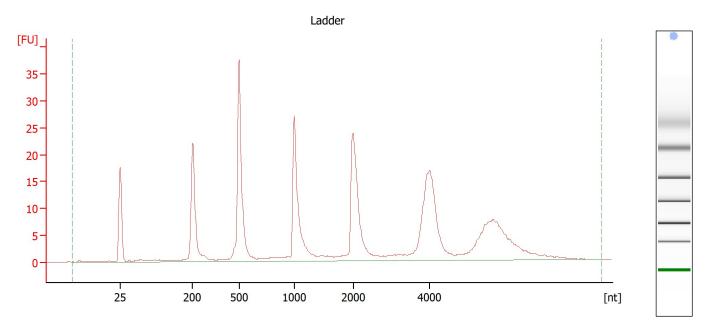
Filter Width [s]: 0.5 Polynomial Order: 4

#### 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\_12-35-05.xad Created: 12/10/2020 12:35:05 PM Modified: 12/10/2020 3:02:51 PM

### **Electropherogram Summary**

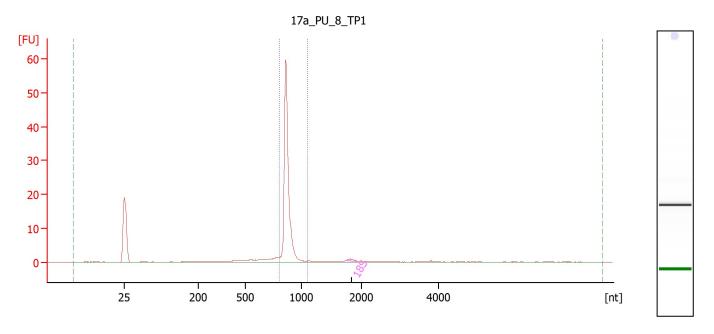


#### **Overall Results for Ladder**

RNA Area: 285.5 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: 1.30

71.6

Overall Results for sample 2: 17a PU 8 TP1

RNA Area: 93.6 RNA Concentration: 49 ng/µl

rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN): 1.3 (B.02.11,

Anomaly Threshold(s) manually adapted)

Fragment table for sample 2: <u>17a PU 8 TP1</u>

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

18S 1,736 1,933 0.7 0.8

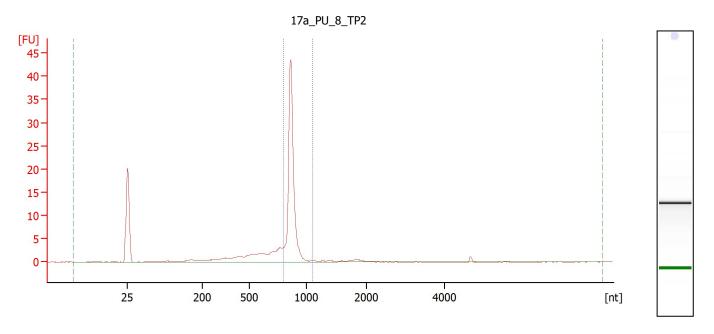
Region table for sample 2 : 17a PU 8 TP1

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

 Region 1
 800
 1,100
 71.6
 76

Assay Class: Created: 12/10/2020 12:35:05 PM Modified: 12/10/2020 3:02:51 PM Eukaryote Total RNA Nano Data Path: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_12-35-05.xad

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

62

RIN: 1.30

58.9

Colo

Overall Results for sample 3: 17a PU 8 TP2

RNA Area: 95.1 RNA Concentration: 50 ng/µl

rRNA Ratio [28s / 18s]: RNA Integrity Number (RIN): 1.3 (B.02.11,

0.0

Anomaly Threshold(s) manually adapted)

58.9

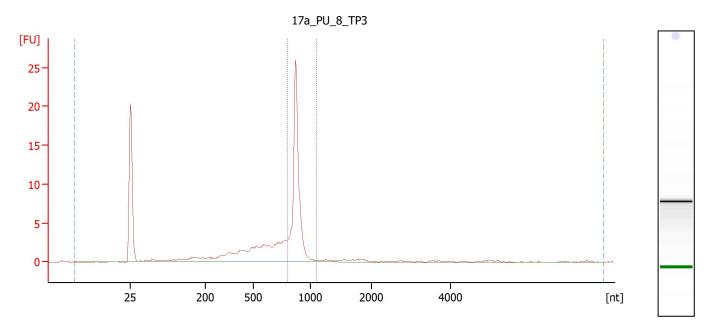
Region table for sample 3: 17a PU 8 TP2 Name From [nt] To [nt] Corr. Area % of Total 800

1,100

Region 1

Assay Class: Created: 12/10/2020 12:35:05 PM Modified: 12/10/2020 3:02:51 PM Eukaryote Total RNA Nano Data Path: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_12-35-05.xad

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 1.60

38.8

Overall Results for sample 4: 17a PU 8 TP3

RNA Area: 81.1 **RNA Concentration:** 43 ng/µl

rRNA Ratio [28s / 18s]:

0.0

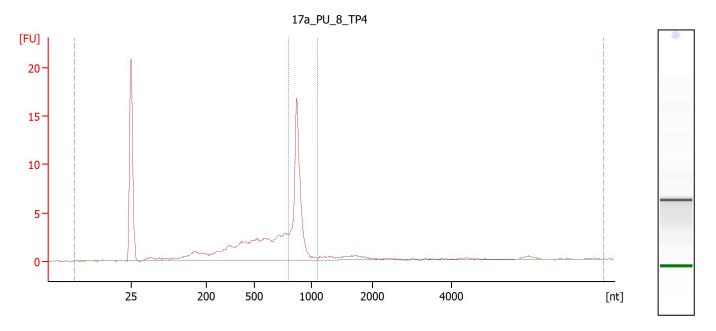
RNA Integrity Number (RIN): 1.6 (B.02.11,

Anomaly Threshold(s) manually adapted)

Region table for sample 4: 17a PU 8 TP3 Name From [nt] To [nt] Corr. Area

% of Total Colo Region 1 800 1,100 38.8 48

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



Overall Results for sample 5 : 17a PU 8 TP4

 RNA Area:
 77.9
 Result Flagging Color:

 RNA Concentration:
 41 ng/μl
 Result Flagging Label:
 RIN: 1.90

 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 27.9

RNA Integrity Number (RIN): 1.9 (B.02.11,
Anomaly Threshold(s)

Anomaly Infestiologs manually adapted)

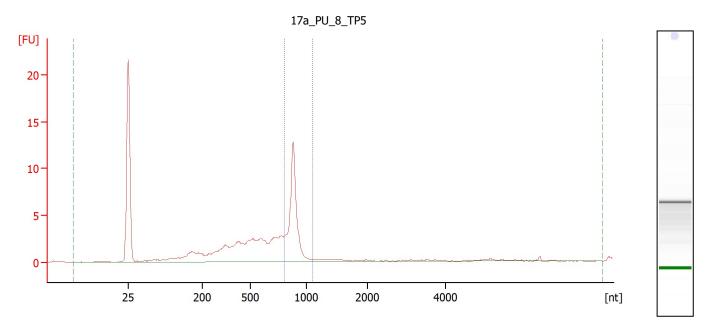
 Region table for sample 5:
 17a PU 8 TP4

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

 Region 1
 800
 1,100
 27.9
 36

Assay Class: Created: 12/10/2020 12:35:05 PM Modified: 12/10/2020 3:02:51 PM Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_12-35-05.xad Data Path:

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN:2

23.4

Overall Results for sample 6: 17a PU 8 TP5

RNA Area: 77.4 **RNA Concentration:** 41 ng/µl

rRNA Ratio [28s / 18s]: 0.0

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

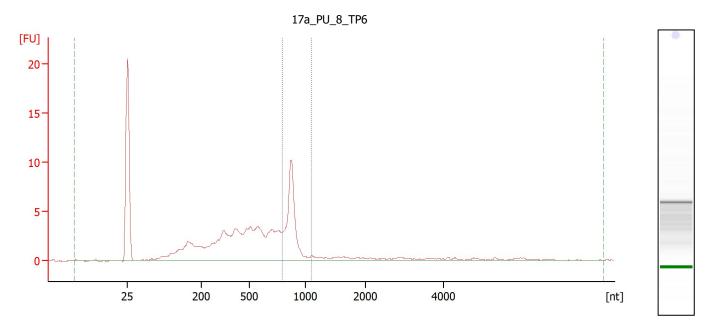
Threshold(s) manually

adapted)

Region table for sample 6: 17a PU 8 TP5

Name From [nt] To [nt] Corr. Area % of Total Colo Region 1 800 1,100 23.4 30

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



# Overall Results for sample 7: 17a PU 8 TP6

 RNA Area:
 101.9
 Result Flagging Color:

 RNA Concentration:
 54 ng/μl
 Result Flagging Label:
 RIN: 2.20

 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 20.8

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

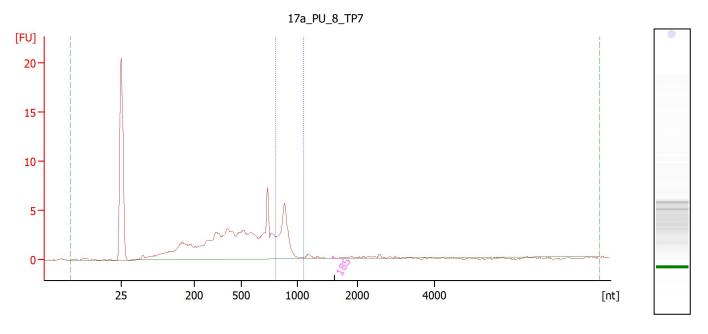
 Region table for sample 7: 17a PU 8 TP6

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

 Region 1
 800
 1,100
 20.8
 20

Assay Class: Created: 12/10/2020 12:35:05 PM Modified: 12/10/2020 3:02:51 PM Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_12-35-05.xad Data Path:

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 2.40

12.9

Overall Results for sample 8: 17a PU 8 TP7

RNA Area: 83.9 RNA Concentration: 44 ng/µl

rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN): 2.4 (B.02.11,

Anomaly Threshold(s) manually adapted)

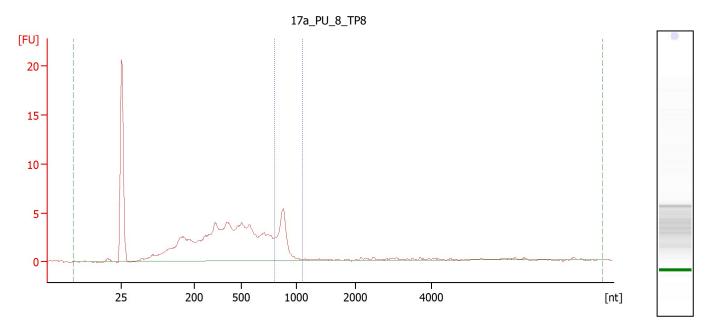
Fragment table for sample 8: 17a PU 8 TP7

Name Start Size [nt] End Size [nt] Area % of total Area 0.2

**18S** 1,555 1,681 0.1

Region table for sample 8: 17a PU 8 TP7 % of Total From [nt] Corr. Area Name To [nt] Colo Region 1 800 1,100 12.9 15

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



Overall Results for sample 9: 17a PU 8 TP8

RNA Area: 102.4 Result Flagging Color: RNA Concentration: 54 ng/µl Result Flagging Label: RIN: 2.60 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 12.9

RNA Integrity Number (RIN): 2.6 (B.02.11, Anomaly Threshold(s) manually adapted)

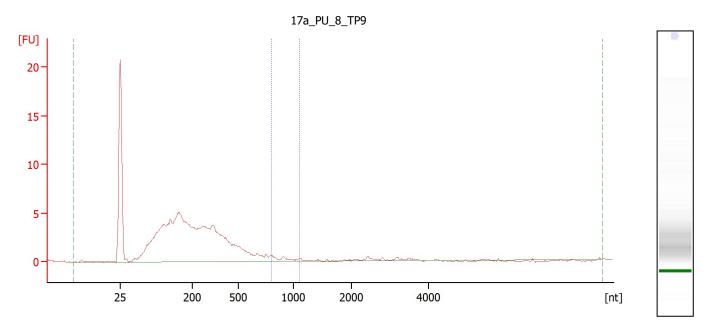
 Region table for sample 9:
 17a PU 8 TP8

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

 Region 1
 800
 1,100
 12.9
 13

Assay Class: Eukaryote Total RNA Nano Created: 12/10/2020 12:35:05 PM Data Path: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_12-35-05.xad Created: 12/10/2020 3:02:51 PM

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



RIN: 2.60

1.7

Overall Results for sample 10 : 17a PU 8 TP9

RNA Area: 107.1 Result Flagging Color: RNA Concentration: 56 ng/µl Result Flagging Label: rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1:

RNA Integrity Number (RIN): 2.6 (B.02.11,

Anomaly Threshold(s) manually adapted)

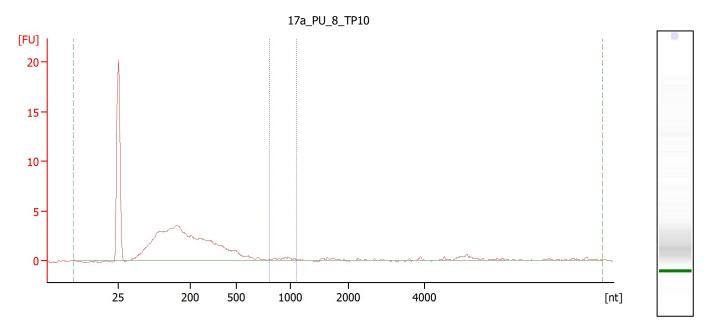
Region table for sample 10 : 17a PU 8 TP9

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

 Region 1
 800
 1,100
 1.7
 2

Assay Class: Created: 12/10/2020 12:35:05 PM Modified: 12/10/2020 3:02:51 PM Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_12-35-05.xad Data Path:

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



#### Overall Results for sample 11: 17a PU 8 TP10

RNA Area: 72.7 Result Flagging Color: RIN: 2.70 **RNA Concentration:** 38 ng/µl Result Flagging Label: Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 1.0

RNA Integrity Number (RIN): 2.7 (B.02.11, Anomaly Threshold(s) manually adapted)

Region table for sample 11: 17a PU 8 TP10 Name From [nt] To [nt] Corr. Area

% of Total Colo Region 1 800 1,100 1.0 1

Assay Class: Data Path: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_12-35-05.xad Created: 12/10/2020 12:35:05 PM Modified: 12/10/2020 3:02:51 PM **Gel Image** 17a\_PU\_8\_T... 17a PU 8 T... 17a\_PU\_8\_T... 17a\_PU\_8\_T... 17a PU 8 T... 17a PU 8 T... 17a\_PU\_8\_T... 17a PU 8 T... 17a PU 8 T... 17a PU 8 T... [nt] Sample 12 Sample 1 Ladder . 4000 -2000 -1000 500 200 25 3 5 6 8 9 10 11 12