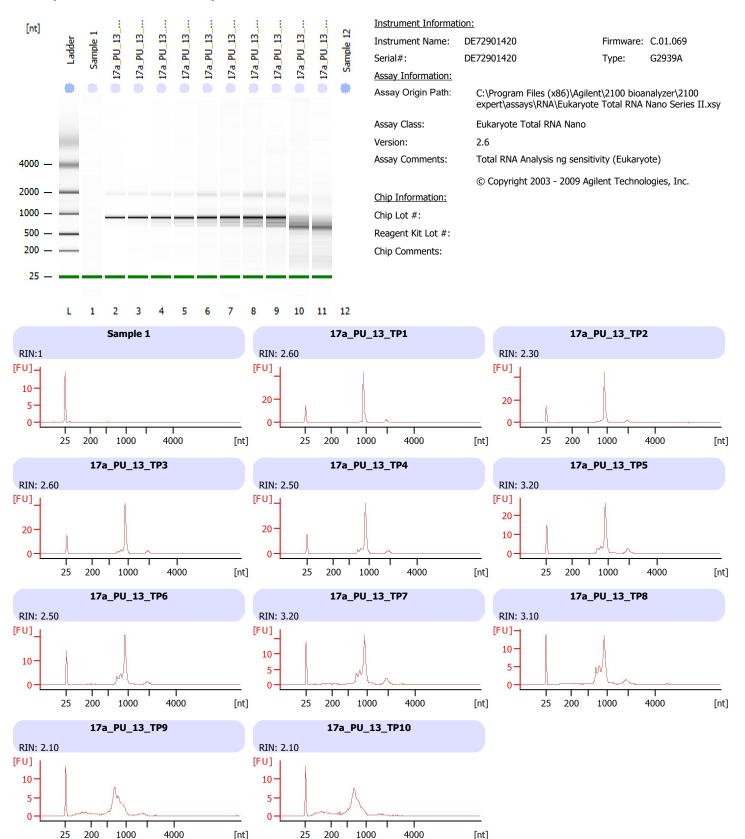
### **Electrophoresis File Run Summary**



1000

[nt]

25 200

25 200 1000

4000

[nt]

Printed: 12/10/2020 4:20:57 PM

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad Created: 12/10/2020 3:49:27 PM Modified: 12/10/2020 4:20:36 PM

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

| Sample Name    | Sample<br>Comment | Stat<br>us | t Result Label       | Result Color |
|----------------|-------------------|------------|----------------------|--------------|
| Sample 1       |                   | ~          | RIN:1                |              |
| 17a_PU_13_TP1  |                   | ~          | RIN: 2.60            |              |
| 17a_PU_13_TP2  |                   | <b>V</b>   | RIN: 2.30            |              |
| 17a_PU_13_TP3  |                   | ~          | RIN: 2.60            |              |
| 17a_PU_13_TP4  |                   | ~          | RIN: 2.50            |              |
| 17a_PU_13_TP5  |                   | <b>V</b>   | RIN: 3.20            |              |
| 17a_PU_13_TP6  |                   | ~          | RIN: 2.50            |              |
| 17a_PU_13_TP7  |                   | ~          | RIN: 3.20            |              |
| 17a_PU_13_TP8  |                   | ~          | RIN: 3.10            |              |
| 17a_PU_13_TP9  |                   | ~          | RIN: 2.10            |              |
| 17a_PU_13_TP10 |                   | ~          | RIN: 2.10            |              |
| Sample 12      |                   |            |                      |              |
| Ladder         |                   | ~          | All Other<br>Samples |              |

Chip Lot # Reagent Kit Lot #

**Chip Comments:** 

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Assay Class: Eukaryote Total RNA Nano Created: 12/10/2020 3:49:27 PM Data Path: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad Created: 12/10/2020 4:20:36 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**

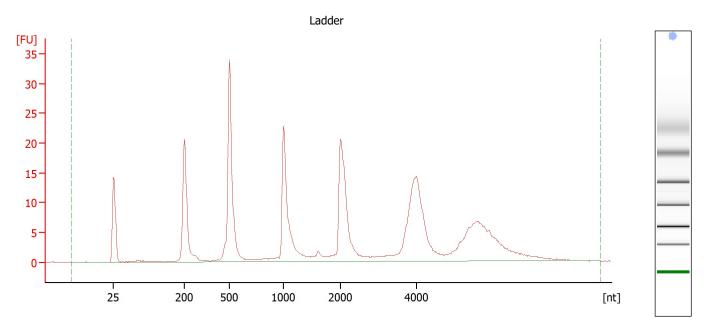
Filter Width [s]: 0.5 Polynomial Order: 4

#### 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\_15-49-28.xad Created: 12/10/2020 3:49:27 PM Modified: 12/10/2020 4:20:36 PM

# **Electropherogram Summary**

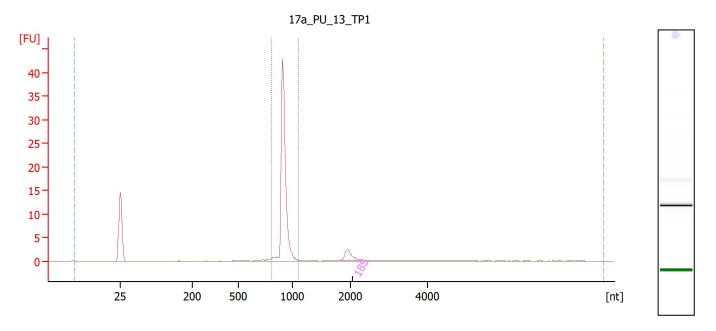


#### **Overall Results for Ladder**

RNA Area: 270.3 RNA Concentration: 150 ng/µl Result Flagging Color: Result Flagging Label:

All Other Samples

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



Result Flagging Color:

Overall Results for sample 2 : 17a PU 13 TP1

RNA Area: 59.5

RNA Concentration: 33 ng/ $\mu$ l Result Flagging Label: RIN: 2.60 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 50.2

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

Anomaly Threshold(s) manually adapted)

Fragment table for sample 2: 17a PU 13 TP1

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

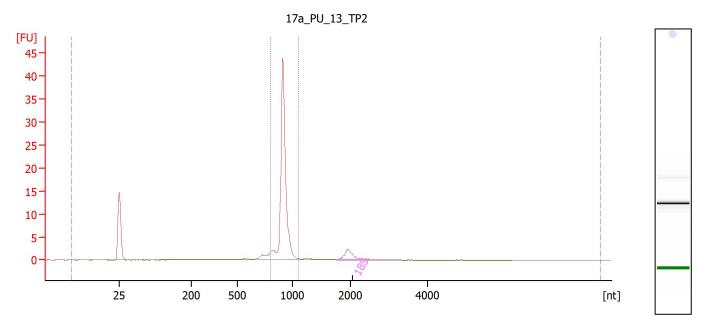
18S 1,739 2,498 4.4 7.4

 Region table for sample 2 : 17a PU 13 TP1

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

 Region 1
 800
 1,100
 50.2
 84

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 2.30

58.4

Overall Results for sample 3: 17a PU 13 TP2

RNA Area: 74.4 RNA Concentration: 41 ng/µl

rRNA Ratio [28s / 18s]: 0.0

Fragment table for sample 3:

RNA Integrity Number (RIN): 2.3 (B.02.11,

Anomaly Threshold(s) manually adapted)

17a PU 13 TP2

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

18S 1,761 2,457 4.1 5.6

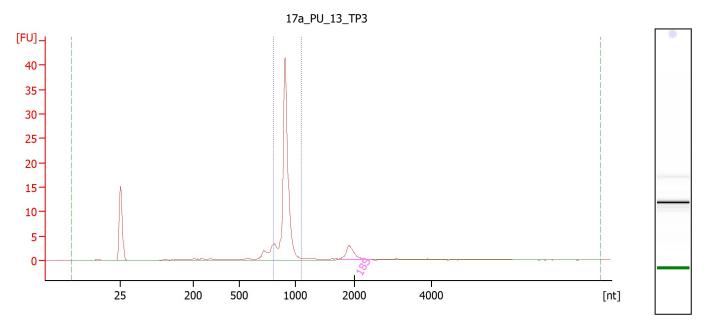
Region table for sample 3: 17a PU 13 TP2

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

 Region 1
 800
 1,100
 58.4
 78

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad Modified: 12/10/2020 4:20:36 PM

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 2.60

60.5

Overall Results for sample 4: 17a PU 13 TP3

RNA Area: 79.1 RNA Concentration: 44 ng/µl

rRNA Ratio [28s / 18s]: 0.0

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

Anomaly Threshold(s) manually adapted)

Fragment table for sample 4: <u>17a\_PU\_13\_TP3</u>

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

18S 1,738 2,458 5.5 6.9

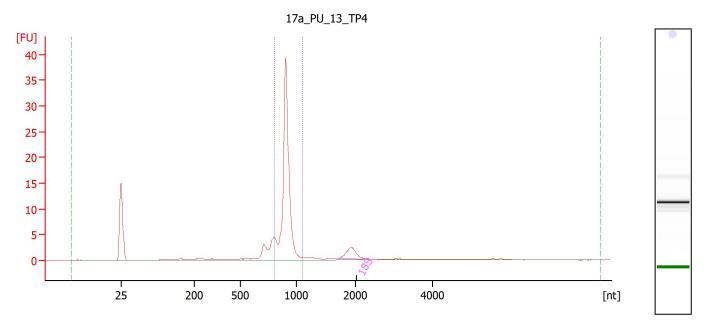
Region table for sample 4: <u>17a PU 13 TP3</u>

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

 Region 1
 800
 1,100
 60.5
 77

Assay Class: Eukaryote Total RNA Nano 12/10/2020 3:49:27 PM Created: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad Modified: 12/10/2020 4:20:36 PM Data Path:

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 2.50

62.7

Overall Results for sample 5: 17a PU 13 TP4

RNA Area: 90.2 RNA Concentration: 50 ng/µl

rRNA Ratio [28s / 18s]:

Anomaly Threshold(s)

RNA Integrity Number (RIN): 2.5 (B.02.11,

manually adapted)

Fragment table for sample 5: 17a PU 13 TP4

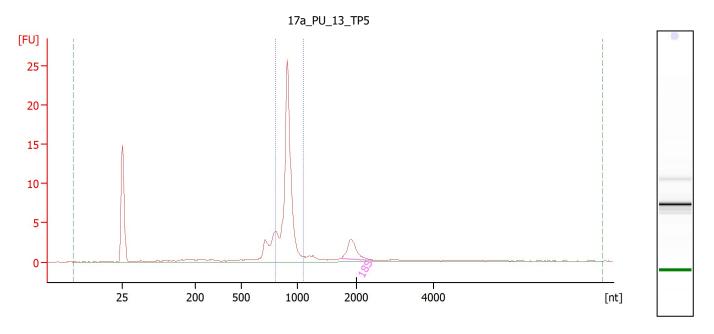
Name Start Size [nt] End Size [nt] **Area** % of total Area **18S** 1,676 2,555 5.5 6.1

0.0

Region table for sample 5: 17a PU 13 TP4 From [nt] % of Total Name To [nt] Corr. Area Colo Region 1 800 1,100 62.7 69

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad Modified: 12/10/2020 4:20:36 PM

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 3.20

47.5

Overall Results for sample 6 : 17a PU 13 TP5

RNA Area: 76.9 RNA Concentration: 43 ng/µl

rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN): 3.2 (B.02.11,

Anomaly Threshold(s) manually adapted)

Fragment table for sample 6 : <u>17a\_PU\_13\_TP5</u>

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

 18S
 1,729
 2,444
 5.7
 7.4

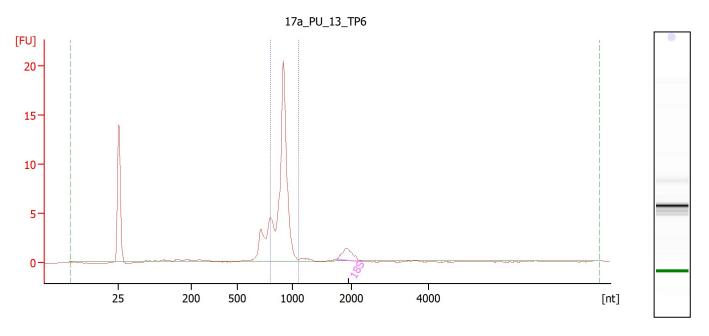
Region table for sample 6: 17a PU 13 TP5

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

 Region 1
 800
 1,100
 47.5
 62

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad Modified: 12/10/2020 4:20:36 PM

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 2.50

41.4

Overall Results for sample 7: 17a PU 13 TP6

RNA Area: 58.1 RNA Concentration: 32 ng/µl

rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN): 2.5 (B.02.11,

Anomaly Threshold(s) manually adapted)

Fragment table for sample 7: <u>17a\_PU\_13\_TP6</u>

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

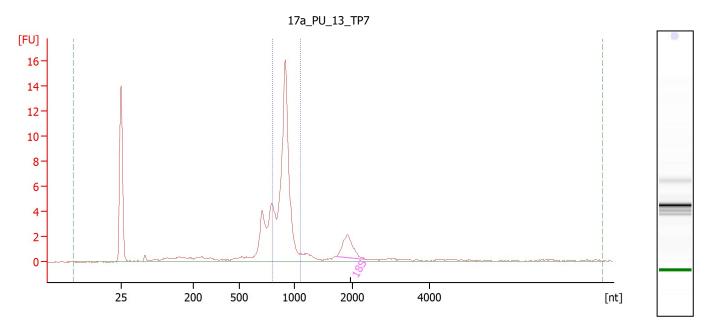
 18S
 1,726
 2,305
 3.2
 5.6

Region table for sample 7: 17a PU 13 TP6

Name From [nt] To [nt] Corr. Area % of Total Colo r
Region 1 800 1,100 41.4 71

Assay Class: Eukaryote Total RNA Nano 12/10/2020 3:49:27 PM Created: 12/10/2020 4:20:36 PM C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad Modified: Data Path:

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 3.20

37.1

Overall Results for sample 8: 17a PU 13 TP7

RNA Area: 69.9 RNA Concentration: 39 ng/µl

rRNA Ratio [28s / 18s]:

0.0

RNA Integrity Number (RIN): 3.2 (B.02.11,

Anomaly Threshold(s) manually adapted)

Fragment table for sample 8: 17a PU 13 TP7

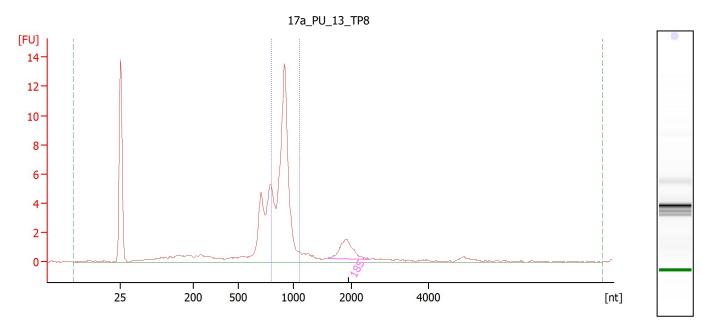
Name Start Size [nt] End Size [nt] **Area** % of total Area **18S** 1,706 2,346 4.4 6.4

Region table for sample 8: 17a PU 13 TP7

% of Total From [nt] Corr. Area Name To [nt] Colo Region 1 800 1,100 37.1 53

Assay Class: Eukaryote Total RNA Nano 12/10/2020 3:49:27 PM Created: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad Modified: 12/10/2020 4:20:36 PM Data Path:

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



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 3.10

35.4

Overall Results for sample 9: 17a PU 13 TP8

RNA Area: 76.6 RNA Concentration: 42 ng/µl

rRNA Ratio [28s / 18s]:

Anomaly Threshold(s)

RNA Integrity Number (RIN): 3.1 (B.02.11,

manually adapted)

Fragment table for sample 9: 17a PU 13 TP8

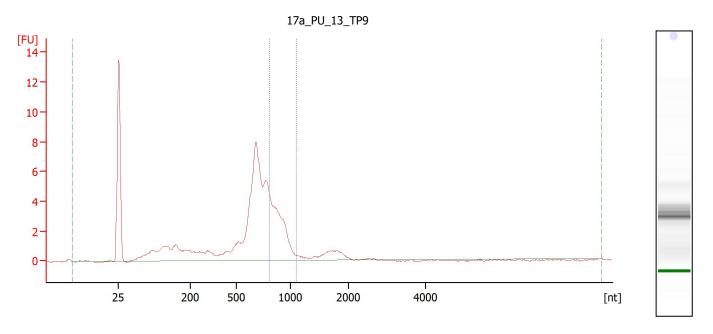
Name Start Size [nt] End Size [nt] **Area** % of total Area 5.6

0.0

**18S** 1,576 2,471 4.3

17a PU 13 TP8 Region table for sample 9: % of Total From [nt] Name To [nt] Corr. Area Colo Region 1 800 1,100 35.4 46

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



# Overall Results for sample 10 : 17a PU 13 TP9

RNA Area: 75.0 Result Flagging Color: RNA Concentration: 42 ng/μl Result Flagging Label: RIN: 2.10 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 16.4

RNA Integrity Number (RIN): 2.1 (B.02.11,

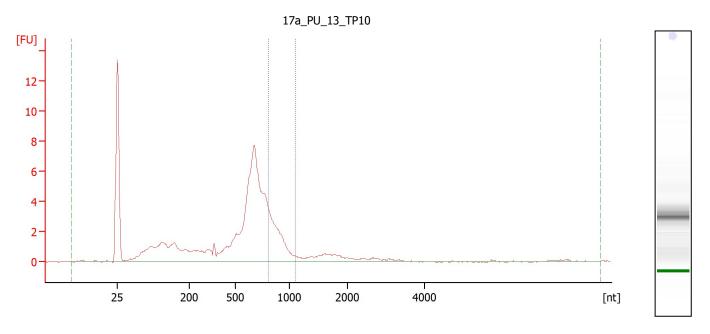
Anomaly Threshold(s) manually adapted)

Region table for sample 10 : 17a PU 13 TP9

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

 Region 1
 800
 1,100
 16.4
 22

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



# Overall Results for sample 11: 17a PU 13 TP10

 RNA Area:
 80.8
 Result Flagging Color:

 RNA Concentration:
 45 ng/μl
 Result Flagging Label:
 RIN: 2.10

 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 11.4

RNA Integrity Number (RIN): 2.1 (B.02.11, Anomaly Thresh

Anomaly Threshold(s) manually adapted)

Region table for sample 11 : 17a PU 13 TP10

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

 Region 1
 800
 1,100
 11.4
 14

Printed: 12/10/2020 4:20:57 PM

Assay Class: Data Path: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_15-49-28.xad Created: Modified: 12/10/2020 3:49:27 PM 12/10/2020 4:20:36 PM **Gel Image** 17a\_PU\_13\_... 17a PU 13 ... 17a PU 13 ... 17a\_PU\_13\_... 17a\_PU\_13\_... 17a\_PU\_13\_... 17a PU 13 ... 17a\_PU\_13\_... 17a PU\_13 ... 17a PU\_13 .... [nt] Sample 12 Ladder Sample 1 . ø 4000 -2000 -1000 500 200 25 -3 5 8 10 11 12