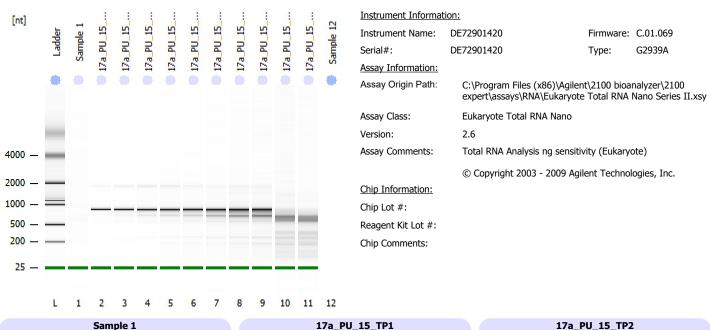
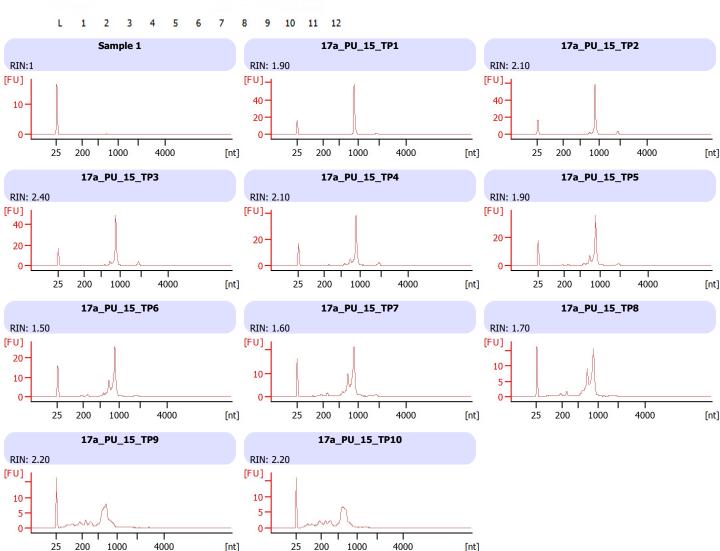
# **Electrophoresis File Run Summary**





1000

[nt]

25 200

25 200 1000

4000

[nt]

Printed: 12/10/2020 4:58:03 PM

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_16-13-55.xad Created: 12/10/2020 4:13:55 PM Modified: 12/10/2020 4:57:47 PM

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

Sample Name	Sample Comment	Stat Result Labous	el Result Color
Sample 1		RIN:1	
17a_PU_15_TP1		RIN: 1.90	
17a_PU_15_TP2		RIN: 2.10	
17a_PU_15_TP3		RIN: 2.40	
17a_PU_15_TP4		RIN: 2.10	
17a_PU_15_TP5		RIN: 1.90	
17a_PU_15_TP6		RIN: 1.50	
17a_PU_15_TP7		RIN: 1.60	
17a_PU_15_TP8		RIN: 1.70	
17a_PU_15_TP9		RIN: 2.20	
17a_PU_15_TP10		RIN: 2.20	
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\_16-13-55.xad Created: 12/10/2020 4:13:55 PM Modified: 12/10/2020 4:57:47 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**

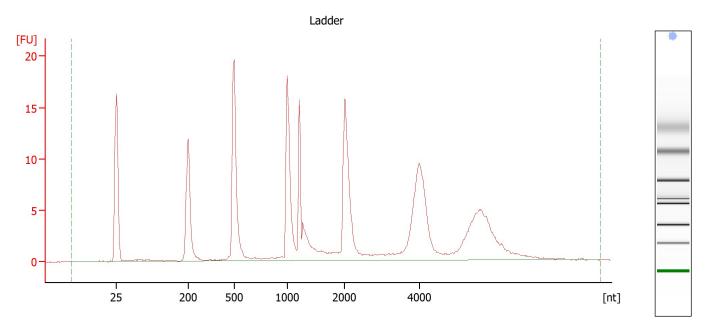
Filter Width [s]: 0.5 Polynomial Order: 4

#### 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\_16-13-55.xad Created: 12/10/2020 4:13:55 PM Modified: 12/10/2020 4:57:47 PM

# **Electropherogram Summary**



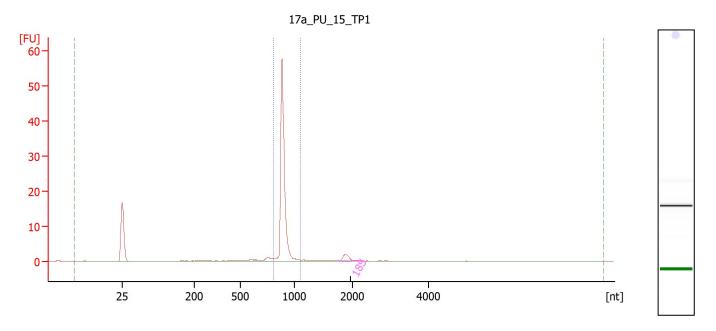
#### **Overall Results for Ladder**

RNA Area: 169.2 RNA Concentration: 150 ng/µl Result Flagging Color: Result Flagging Label:

All Other Samples

Assay Class: Eukaryote Total RNA Nano Created: 12/10/2020 4:13:55 PM Data Path: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_16-13-55.xad Modified: 12/10/2020 4:57:47 PM

# **Electropherogram Summary Continued ...**



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 1.90

59.4

Overall Results for sample 2 : 17a PU 15 TP1

RNA Area: 75.8 RNA Concentration: 67 ng/µl

rRNA Ratio [28s / 18s]: 0.0

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

Anomaly Threshold(s) manually adapted)

Fragment table for sample 2 : <u>17a\_PU\_15\_TP1</u>

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

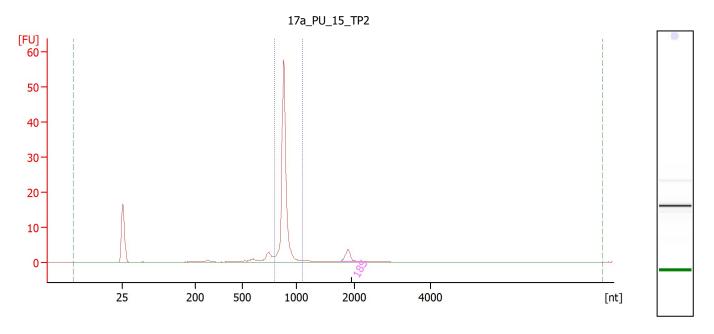
18S 1,727 2,332 2.9 3.9

Region table for sample 2: <u>17a PU 15 TP1</u>

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

 Region 1
 800
 1,100
 59.4
 78

# **Electropherogram Summary Continued ...**



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 2.10

66.6

Overall Results for sample 3: 17a PU 15 TP2

RNA Area: 87.2 RNA Concentration: 77 ng/µl

rRNA Ratio [28s / 18s]: 0.0

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

Anomaly Threshold(s) manually adapted)

Fragment table for sample 3: 17a PU 15 TP2

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

18S 1,749 2,193 4.3 4.9

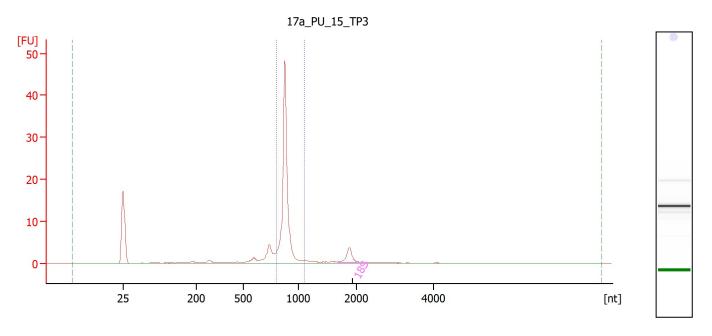
 Region table for sample 3:
 17a PU 15 TP2

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

 Region 1
 800
 1,100
 66.6
 76

Assay Class: 12/10/2020 4:13:55 PM Eukaryote Total RNA Nano Created: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_16-13-55.xad Modified: 12/10/2020 4:57:47 PM Data Path:

# **Electropherogram Summary Continued ...**



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 2.40

62.0

Overall Results for sample 4: 17a PU 15 TP3

RNA Area: 91.9 RNA Concentration: 81 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 4: 17a PU 15 TP3

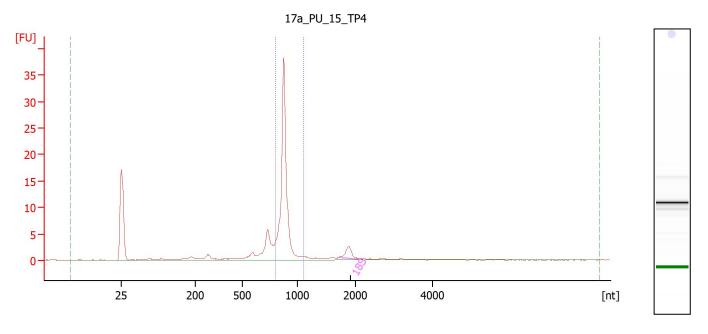
Name Start Size [nt] End Size [nt] Area % of total Area **18S** 1,662 2,280 4.9 5.3

Region table for sample 4: 17a PU 15 TP3

From [nt] Corr. Area % of Total Name To [nt] Colo Region 1 800 1,100 62.0 67

Assay Class: 12/10/2020 4:13:55 PM Eukaryote Total RNA Nano Created: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_16-13-55.xad Modified: 12/10/2020 4:57:47 PM Data Path:

# **Electropherogram Summary Continued ...**



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 2.10

54.6

Overall Results for sample 5: 17a PU 15 TP4

RNA Area: 91.6 RNA Concentration: 81 ng/µl

rRNA Ratio [28s / 18s]: 0.0

Anomaly Threshold(s)

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

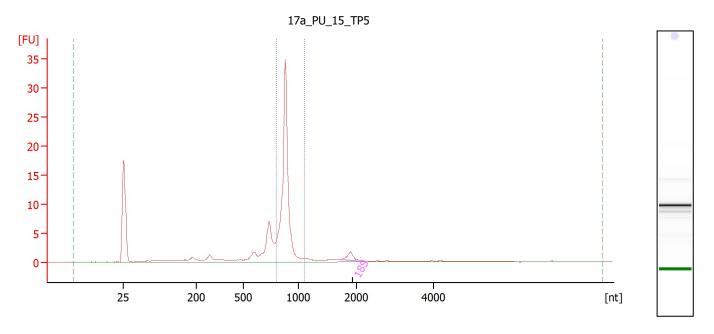
manually adapted)

Fragment table for sample 5: 17a PU 15 TP4

Name Start Size [nt] End Size [nt] Area % of total Area 2,114 **18S** 1,746 2.9 3.2

Region table for sample 5: 17a PU 15 TP4 % of Total From [nt] Name To [nt] Corr. Area Colo Region 1 800 1,100 54.6 60

# **Electropherogram Summary Continued ...**



Result Flagging Color:

Result Flagging Label:

Corr. Area 1:

RIN: 1.90

53.4

Overall Results for sample 6 : 17a PU 15 TP5

RNA Area: 94.4 RNA Concentration: 84 ng/µl

rRNA Ratio [28s / 18s]: 0.0

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

Anomaly Threshold(s) manually adapted)

Fragment table for sample 6 : <u>17a\_PU\_15\_TP5</u>

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

18S 1,679 2,293 2.5 2.6

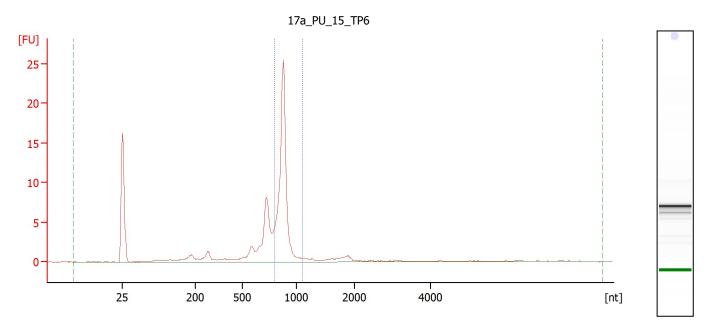
 Region table for sample 6:
 17a PU 15 TP5

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

 Region 1
 800
 1,100
 53.4
 56

Assay Class: Eukaryote Total RNA Nano 12/10/2020 4:13:55 PM Created: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_16-13-55.xad Data Path: Modified: 12/10/2020 4:57:47 PM

# **Electropherogram Summary Continued ...**



#### Overall Results for sample 7: 17a PU 15 TP6

RNA Area: 85.1 **RNA Concentration:** 75 ng/µl

rRNA Ratio [28s / 18s]:

0.0

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

Anomaly Threshold(s) manually adapted)

Result Flagging Color: Result Flagging Label:

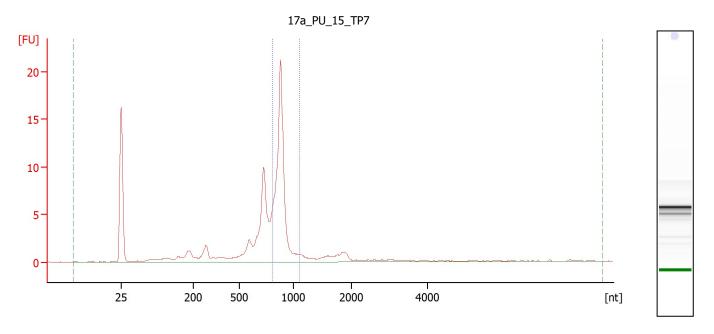
RIN: 1.50 Corr. Area 1: 43.8

Region table	for sample 7	:	<u>17a</u>	PU_15	TP6
Name	Fuere Feet	-	- F-47	C	A

% of Total Colo Name From [nt] To [nt] Corr. Area Region 1 800 1,100 43.8 51

Assay Class: Eukaryote Total RNA Nano 12/10/2020 4:13:55 PM Created: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_16-13-55.xad Data Path: Modified: 12/10/2020 4:57:47 PM

# **Electropherogram Summary Continued ...**



RIN: 1.60

Overall Results for sample 8: 17a PU 15 TP7

RNA Area: 92.3 Result Flagging Color: **RNA Concentration:** 82 ng/µl Result Flagging Label:

Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 41.6

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

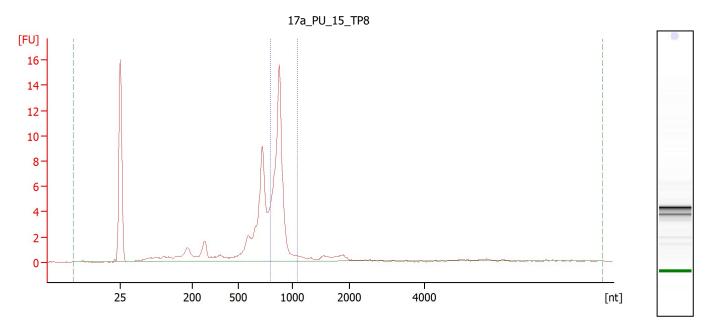
manually adapted)

Region table for sample 8: 17a PU 15 TP7

Name From [nt] To [nt] Corr. Area % of Total Colo Region 1 800 1,100 41.6 45

Assay Class: Eukaryote Total RNA Nano 12/10/2020 4:13:55 PM Created: C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_16-13-55.xad Data Path: Modified: 12/10/2020 4:57:47 PM

# **Electropherogram Summary Continued ...**



Overall Results for sample 9: 17a PU 15 TP8

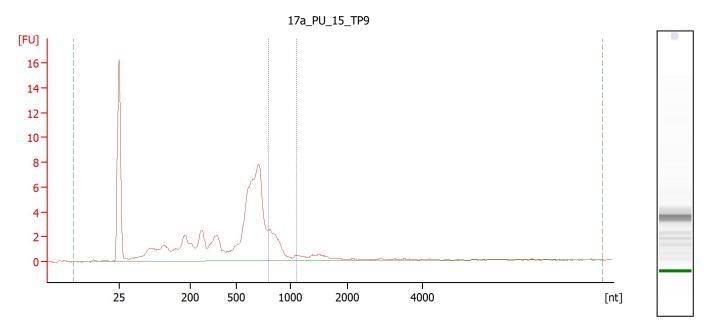
RNA Area: 75.6 Result Flagging Color: RNA Concentration: 67 ng/µl Result Flagging Label: RIN: 1.70 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 32.4

RNA Integrity Number (RIN): 1.7 (B.02.11, Anomaly Threshold(s)

manually adapted)

Region table for sample 9: 17a PU 15 TP8 Name From [nt] To [nt] Corr. Area % of Total Colo Region 1 800 1,100 32.4 43

# **Electropherogram Summary Continued ...**



# Overall Results for sample 10 : 17a PU 15 TP9

 RNA Area:
 90.0
 Result Flagging Color:

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

 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 7.8

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

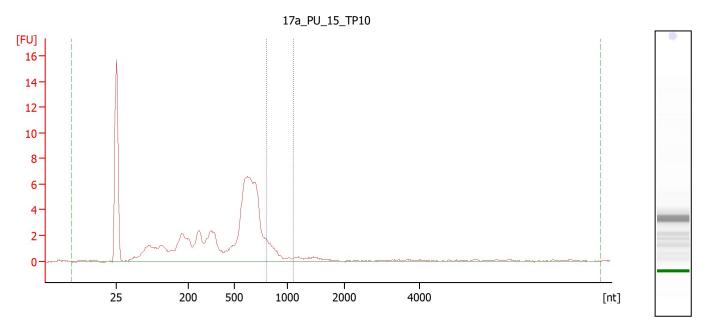
Anomaly Threshold(s) manually adapted)

Region table for sample 10 : 17a PU 15 TP9

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

 Region 1
 800
 1,100
 7.8
 9

# **Electropherogram Summary Continued ...**



# Overall Results for sample 11: 17a PU 15 TP10

RNA Area: 84.1 Result Flagging Color: RNA Concentration: 75 ng/µl Result Flagging Label: RIN: 2.20 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 4.7

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

Anomaly Threshold(s) manually adapted)

Region table for sample 11: 17a PU 15 TP10

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

 Region 1
 800
 1,100
 4.7
 6

Assay Class: Data Path: Eukaryote Total RNA Nano C:\...2020-12-10\Eukaryote Total RNA Nano\_2020-12-10\_16-13-55.xad Created: Modified: 12/10/2020 4:13:55 PM 12/10/2020 4:57:47 PM **Gel Image** 17a\_PU\_15\_... 17a PU 15 ... 17a PU 15 ... 17a\_PU\_15\_... 17a PU 15 ... 17a\_PU\_15\_... 17a PU 15 ... 17a\_PU\_15\_... 17a\_PU\_15\_... 17a PU 15 ... [nt] Sample 12 Ladder Sample 1 . 4000 2000 -1000 500 200 -2 3 5 6 10 11 12