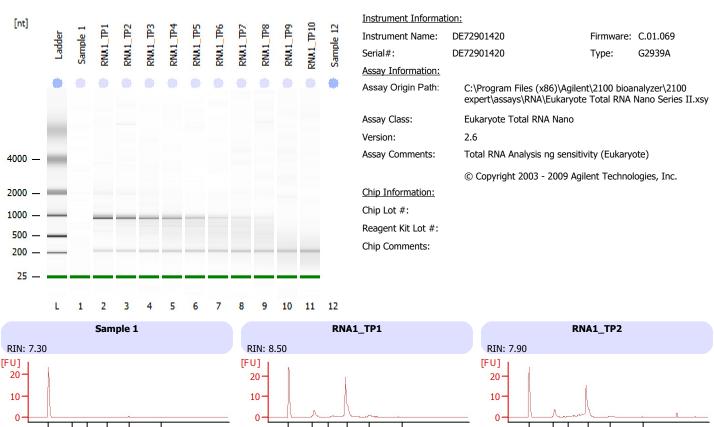
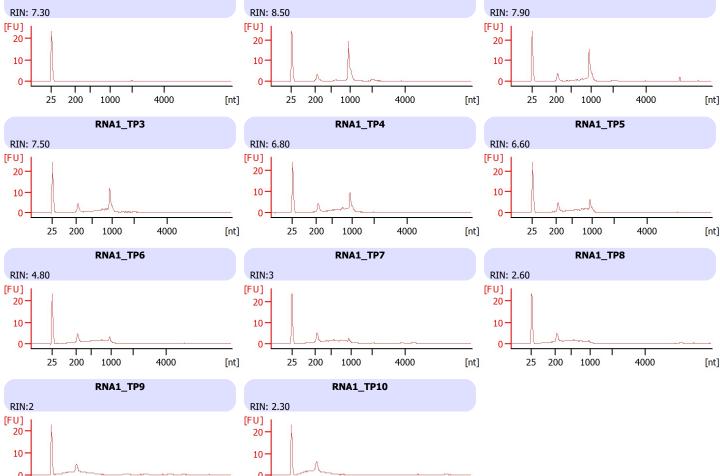
## **Electrophoresis File Run Summary**





200

1000

4000

[nt]

25 200

25

1000

4000

[nt]

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

Sample Name	Sample Comment	Stat Resu	ult Label	Result Color
Sample 1		RIN:	7.30	
RNA1_TP1		RIN:	8.50	
RNA1_TP2		RIN:	7.90	
RNA1_TP3		RIN:	7.50	
RNA1_TP4		RIN:	6.80	
RNA1_TP5		RIN:	6.60	
RNA1_TP6		RIN:	4.80	
RNA1_TP7		RIN:	3	
RNA1_TP8		RIN:	2.60	
RNA1_TP9		RIN:	2	
RNA1_TP10		RIN:	2.30	
Sample 12		All O	ther	
•		Samı	ples	
Ladder		All O	ther	
		Sam	ples	

Chip Lot # Reagent Kit Lot #

**Chip Comments:** 

## **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

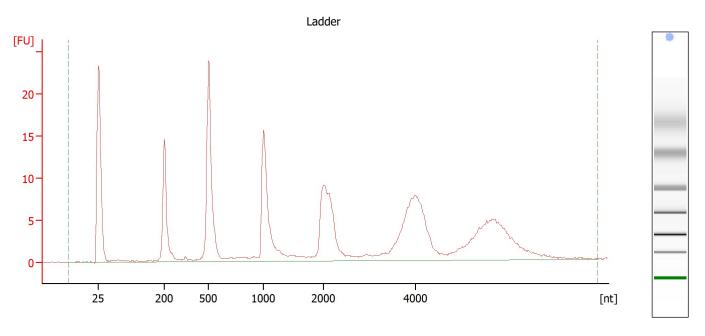
#### **Filter Settings**

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**

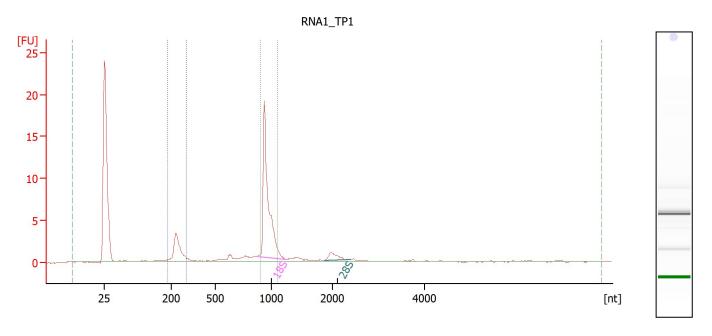


#### **Overall Results for Ladder**

RNA Area: 206.5 RNA Concentration: 150 ng/µl Result Flagging Color: Result Flagging Label:

All Other Samples

# **Electropherogram Summary Continued ...**



## Overall Results for sample 2: RNA1\_TP1

RNA Area: 53.6 Result Flagging Color: RNA Concentration: 39 ng/µl Result Flagging Label: RIN: 8.50 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.1 8.2 RNA Integrity Number (RIN): Corr. Area 2: 8.5 (B.02.10, 27.9 Anomaly Threshold(s)

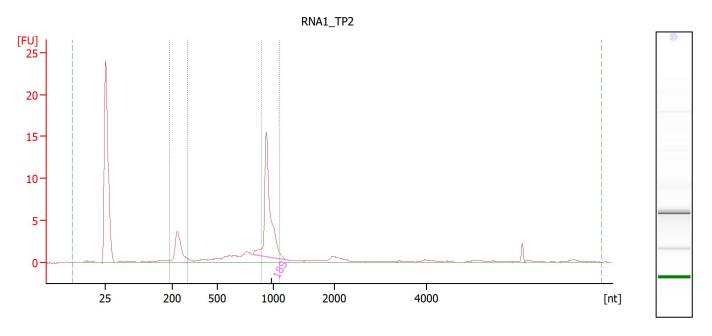
manually adapted)

Fragment table for sample 2: RNA1 TP1

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	866	1,247	26.9	50.3
28S	1,879	2,408	2.0	3.8

Region table for sample 2: RNA1 TP1 % of Total Name From [nt] To [nt] Corr. Area Colo Region 1 190 300 8.2 15 Region 2 900 1,100 27.9 52

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



Overall Results for sample 3: RNA1\_TP2

RNA Area: 56.1 Result Flagging Color: RNA Concentration: 41 ng/ $\mu$ l Result Flagging Label: RIN: 7.90 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 8.3 RNA Integrity Number (RIN): 7.9 (B.02.10, Corr. Area 2: 24.1

Anomaly Threshold(s) manually adapted)

Fragment table for sample 3: RNA1\_TP2

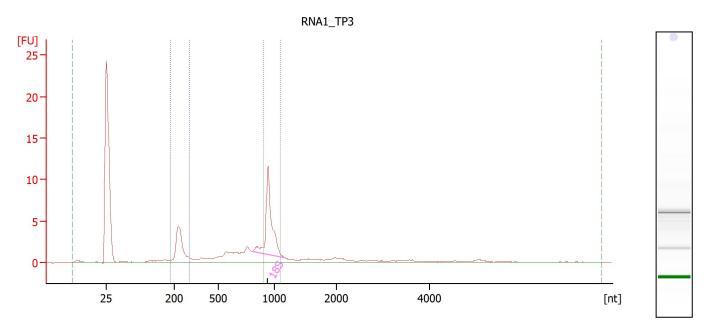
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 812
 1,278
 23.5
 41.9

Region table for sample 3: RNA1 TP2

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	190	300	8.3	15	
Region 2	900	1,100	24.1	43	

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



Overall Results for sample 4: RNA1\_TP3

RNA Area: 60.0 Result Flagging Color:

RNA Concentration: 44 ng/µl Result Flagging Label: RIN: 7.50

rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 10.3

RNA Integrity Number (RIN): 7.5 (B.02.10, Corr. Area 2: 18.8

Anomaly Threshold(s) manually adapted)

Fragment table for sample 4: RNA1\_TP3

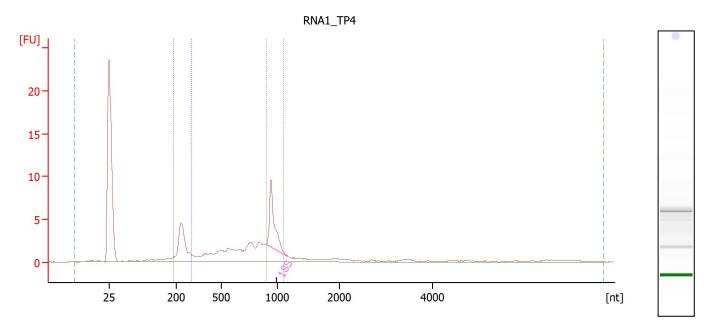
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 801
 1,143
 16.7
 27.9

Region table for sample 4: RNA1 TP3

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	190	300	10.3	17	
Region 2	900	1,100	18.8	31	

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



Overall Results for sample 5: RNA1 TP4

 RNA Area:
 72.5
 Result Flagging Color:

 RNA Concentration:
 53 ng/μl
 Result Flagging Label:
 RIN: 6.80

 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 11.7

 RNA Integrity Number (RIN):
 6.8 (B.02.10)
 Corr. Area 2:
 17.3

Fragment table for sample 5: RNA1 TP4

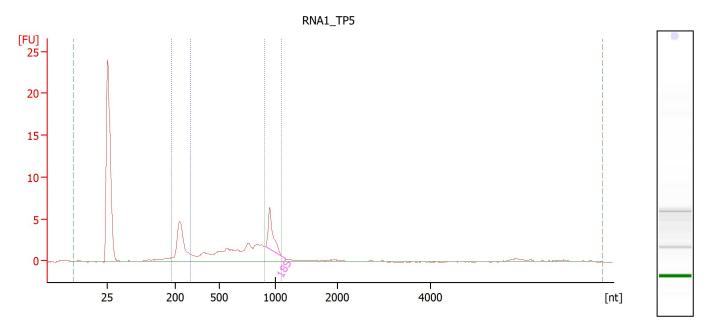
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 903
 1,162
 10.6
 14.6

Region table for sample 5 : RNA1\_TP4

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	190	300	11.7	16	
Region 2	900	1,100	17.3	24	

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



Overall Results for sample 6: RNA1\_TP5

RNA Area: 58.4 Result Flagging Color: RNA Concentration: 42 ng/ $\mu$ l Result Flagging Label: RIN: 6.60 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 12.1 RNA Integrity Number (RIN): 6.6 (B.02.10) Corr. Area 2: 12.0

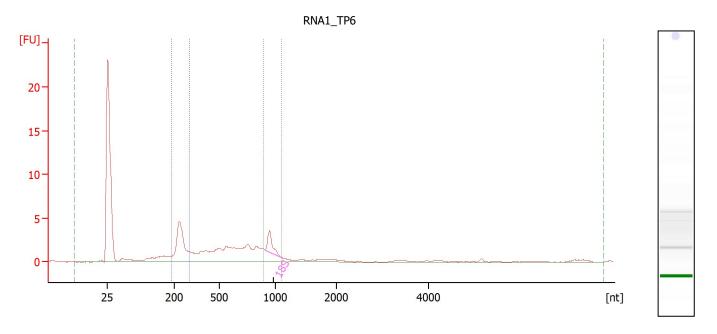
Fragment table for sample 6: RNA1 TP5

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

18S 906 1,151 6.9 11.8

RNA1\_TP5 Region table for sample 6: Name From [nt] To [nt] Corr. Area % of Total Colo 300 Region 1 190 12.1 21 Region 2 900 1,100 12.0 21

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



# Overall Results for sample 7: RNA1 TP6

RNA Area: 61.7 Result Flagging Color: RNA Concentration: 45 ng/ $\mu$ l Result Flagging Label: RIN: 4.80 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 12.7 RNA Integrity Number (RIN): 4.8 (B.02.10) Corr. Area 2: 7.5

Fragment table for sample 7: RNA1 TP6

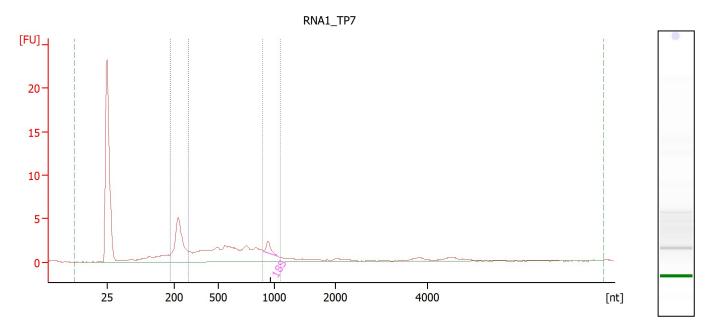
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 913
 1,112
 3.1
 5.0

Region table for sample 7: RNA1 TP6

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	190	300	12.7	21	
Region 2	900	1,100	7.5	12	

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



Overall Results for sample 8: RNA1 TP7

RNA Area: 65.6 Result Flagging Color: RNA Concentration: 48 ng/µl Result Flagging Label: RIN:3 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 14.7 RNA Integrity Number (RIN): 3 (B.02.10) Corr. Area 2: 5.4

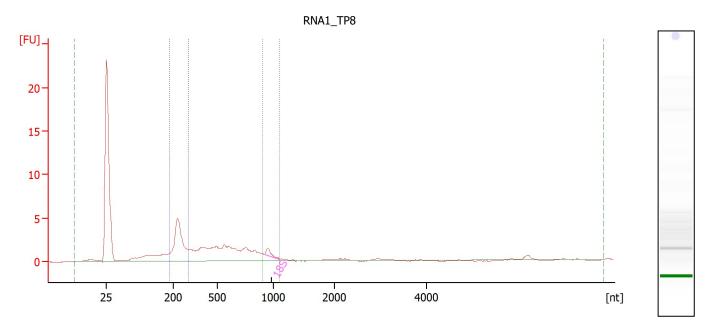
Fragment table for sample 8 : RNA1 TP7

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

18S 906 1,065 1.7 2.6

Region table for sample 8: RNA1\_TP7 Name From [nt] To [nt] Corr. Area % of Total Colo Region 1 190 300 14.7 22 Region 2 900 1,100 5.4

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



Overall Results for sample 9: RNA1 TP8

RNA Area: 61.2 Result Flagging Color: RNA Concentration: 44 ng/µl Result Flagging Label: RIN: 2.60 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 15.0 RNA Integrity Number (RIN): 2.6 (B.02.10) Corr. Area 2: 3.5

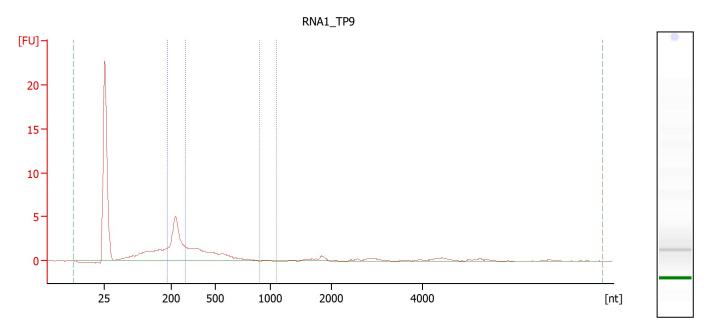
Fragment table for sample 9: RNA1 TP8

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

18S 919 1,107 1.1 1.8

Region table for sample 9: RNA1\_TP8 Name From [nt] To [nt] Corr. Area % of Total Colo 300 Region 1 190 15.0 25 Region 2 900 1,100 3.5

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

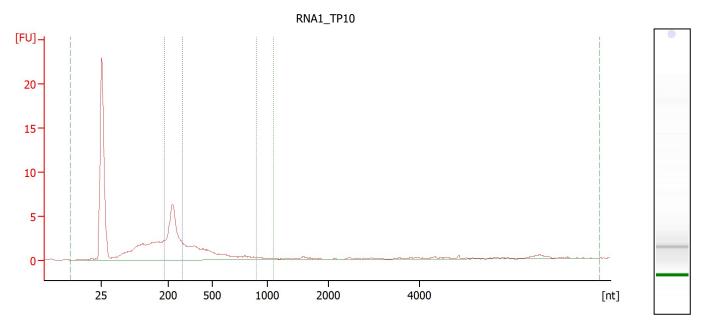


# Overall Results for sample 10 : RNA1 TP9

RNA Area: 52.3 Result Flagging Color: RNA Concentration: 38 ng/µl Result Flagging Label: RIN:2 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 16.7 RNA Integrity Number (RIN): 2 (B.02.10) Corr. Area 2: 0.1

Region table for sample 10 : RNA1 TP9						
Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo	
Region 1	190	300	16.7	32	•	
Region 2	900	1,100	0.1	0		

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



# Overall Results for sample 11: RNA1 TP10

RNA Area:	67.8	Result Flagging Color:	
RNA Concentration:	49 ng/µl	Result Flagging Label:	RIN: 2.30
rRNA Ratio [28s / 18s]:	0.0	Corr. Area 1:	21.6
RNA Integrity Number (RIN):	2.3 (B.02.10)	Corr. Area 2:	0.6

Region table for sample 11: RNA1 TP10						
Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo	
Region 1	190	300	21.6	32	•	
Region 2	900	1,100	0.6	1		

Assay Class: Data Path: Eukaryote Total RNA Nano C:\...2020-10-07\Eukaryote Total RNA Nano\_2020-10-07\_13-56-03.xad Created: Modified: 10/7/2020 1:56:02 PM 10/7/2020 2:30:29 PM **Gel Image** [nt] RNA1\_TP10 Sample 12 RNA1\_TP3 RNA1\_TP6 RNA1\_TP9 RNA1\_TP1 Sample 1 Ladder . ٠ • • • ø • • 4000 — 2000 -1000 500 200 25 1 3 5 10 11 12