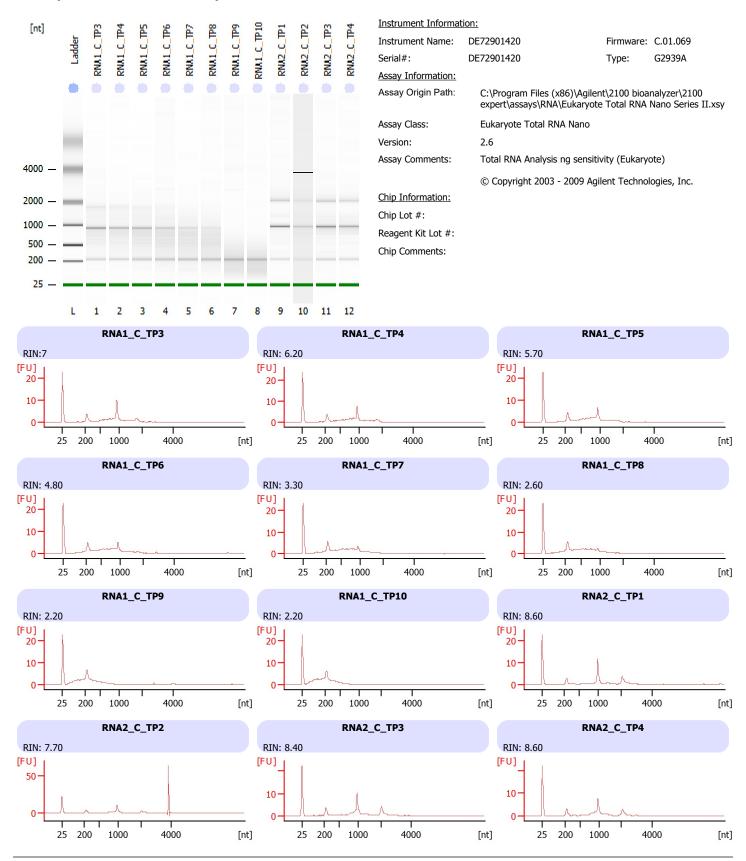
Printed: 10/21/2020 4:46:36 PM

Assay Class: Eukaryote Total RNA Nano C:\...2020-10-21\Eukaryote Total RNA Nano\_2020-10-21\_16-14-00.xad Created: 10/21/2020 4:13:59 PM Modified: 10/21/2020 4:46:22 PM

## **Electrophoresis File Run Summary**



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

Sample Name	Sample Comment	Stat us	Result Label	Result Color
RNA1_C_TP3		~	RIN:7	
RNA1_C_TP4		~	RIN: 6.20	
RNA1_C_TP5		~	RIN: 5.70	
RNA1_C_TP6		~	RIN: 4.80	
RNA1_C_TP7		~	RIN: 3.30	
RNA1_C_TP8		~	RIN: 2.60	
RNA1_C_TP9		~	RIN: 2.20	
RNA1_C_TP10		~	RIN: 2.20	
RNA2_C_TP1		~	RIN: 8.60	
RNA2_C_TP2		~	RIN: 7.70	
RNA2_C_TP3		~	RIN: 8.40	
RNA2_C_TP4		~	RIN: 8.60	
Ladder		~	All Other	
			Samples	

Chip Lot # Reagent Kit Lot #

**Chip Comments:** 

Printed: 10/21/2020 4:46:36 PM

Assay Class: Eukaryote Total RNA Nano C:\...2020-10-21\Eukaryote Total RNA Nano\_2020-10-21\_16-14-00.xad Created: 10/21/2020 4:13:59 PM Modified: 10/21/2020 4:46:22 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

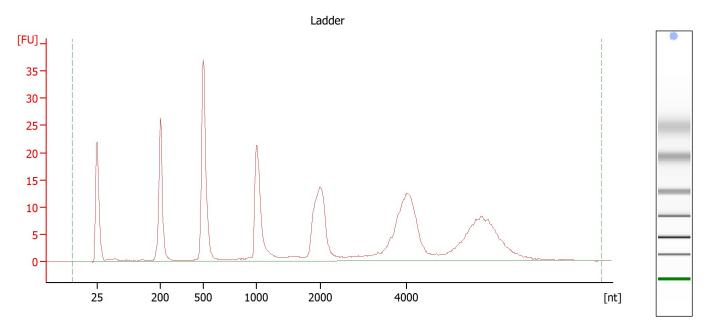
#### **Filter Settings**

Filter Width [s]: 0.5 Polynomial Order: 4

#### Ladder

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

# **Electropherogram Summary**

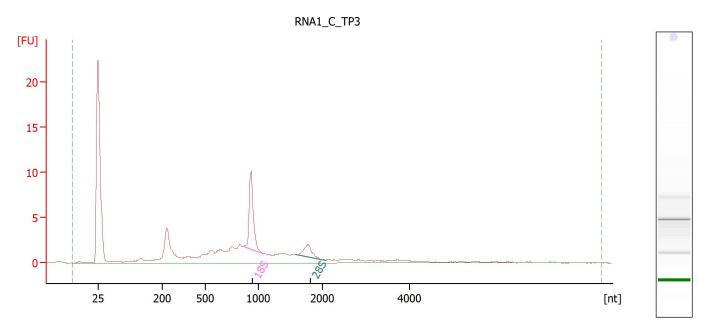


#### **Overall Results for Ladder**

RNA Area: 312.5 RNA Concentration: 150 ng/µl Result Flagging Color: Result Flagging Label:

All Other Samples

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



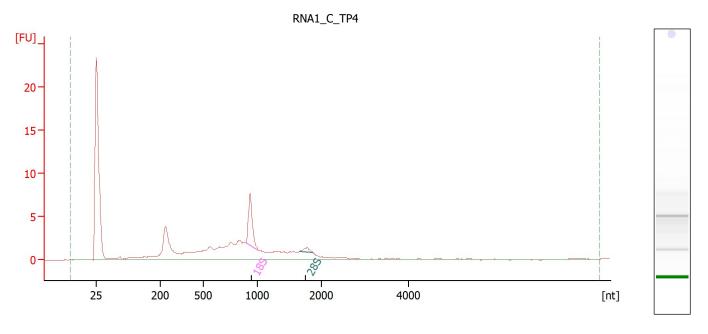
# Overall Results for sample 1: RNA1 C TP3

RNA Area: 75.0 RNA Integrity Number (RIN): 7 (B.02.10)
RNA Concentration: 36 ng/µl Result Flagging Color: RNA Ratio [28s / 18s]: 0.3 Result Flagging Label: RIN:7

#### Fragment table for sample 1: RNA1 C TP3

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	853	1,055	10.1	13.5
28S	1,582	2,074	3.0	4.0

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



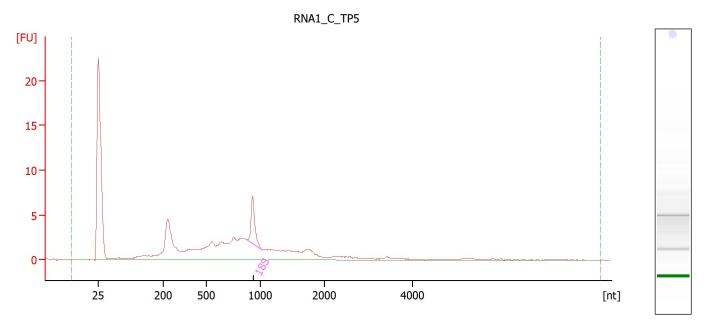
# Overall Results for sample 2: RNA1 C TP4

RNA Area: 67.8 RNA Integrity Number (RIN): 6.2 (B.02.10)
RNA Concentration: 33 ng/µl Result Flagging Color: RNA Ratio [28s / 18s]: 0.1 Result Flagging Label: RIN: 6.20

#### Fragment table for sample 2: RNA1 C TP4

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	890	1,016	6.7	9.9
28S	1,650	1,859	0.8	1.1

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



Overall Results for sample 3: RNA1 C TP5

RNA Area: 84.5 RNA Integrity Number (RIN): 5.7 (B.02.10)
RNA Concentration: 41 ng/µl Result Flagging Color:

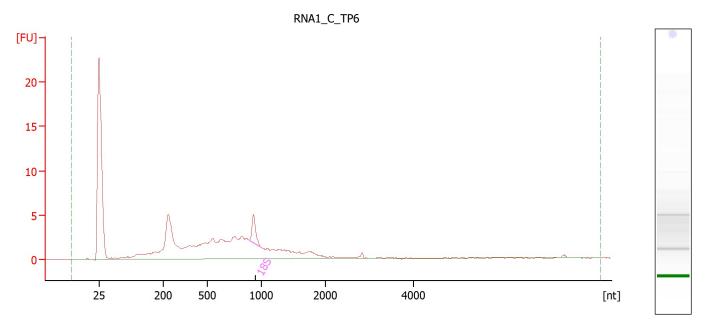
RNA Concentration: 41 ng/µl Result Flagging Color: rRNA Ratio [28s / 18s]: 0.0 Result Flagging Label: RIN: 5.70

Fragment table for sample 3: RNA1 C TP5

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

18S 869 1,021 6.2 7.4

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



Overall Results for sample 4: RNA1 C TP6

RNA Area: 83.4 RNA Integrity Number (RIN): 4.8 (B.02.10)

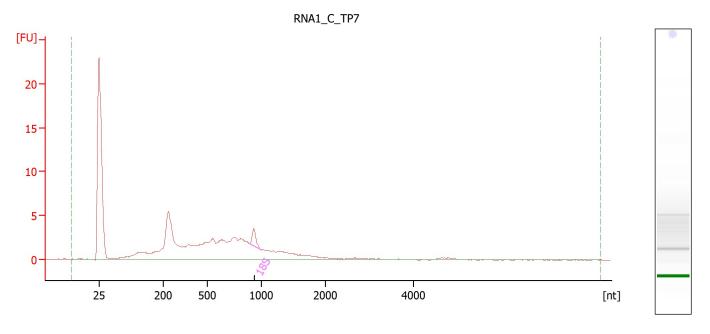
RNA Concentration: 40 ng/µl Result Flagging Color: rRNA Ratio [28s / 18s]: 0.0 Result Flagging Label: RIN: 4.80

Fragment table for sample 4: RNA1 C TP6

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

18S 890 1,007 3.5 4.2

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



Overall Results for sample 5 : RNA1 C TP7

RNA Area: 82.5 RNA Integrity Number (RIN): 3.3 (B.02.10)
RNA Concentration: 40 ng/µl Result Flagging Color:

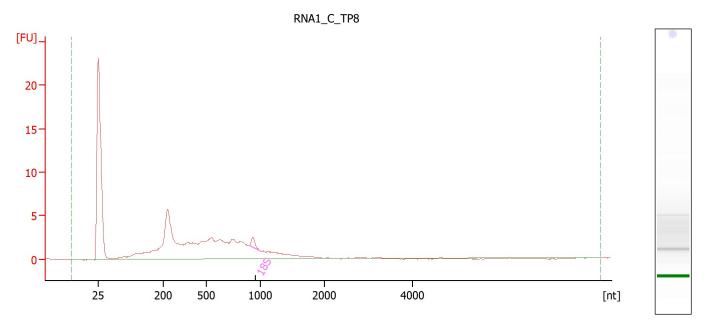
RNA Concentration: 40 ng/µl Result Flagging Color: rRNA Ratio [28s / 18s]: 0.0 Result Flagging Label: RIN: 3.30

Fragment table for sample 5: RNA1 C TP7

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

18S 890 994 2.2 2.7

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



Overall Results for sample 6 : RNA1 C TP8

RNA Area: 77.5 RNA Integrity Number (RIN): 2.6 (B.02.10)
RNA Concentration: 37 ng/µl Result Flagging Color:

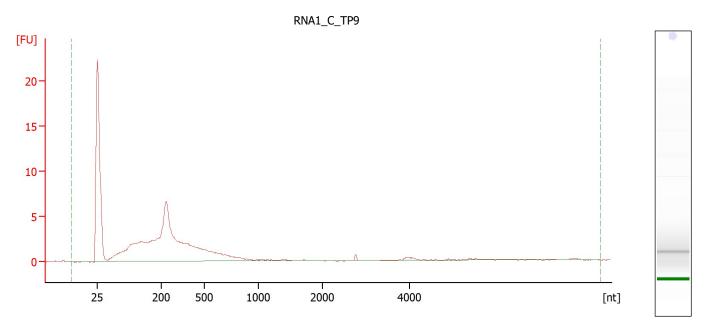
RNA Concentration: 37 ng/µl Result Flagging Color: RNA Ratio [28s / 18s]: 0.0 Result Flagging Label: RIN: 2.60

Fragment table for sample 6: RNA1 C TP8

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

18S 899 1,013 1.2 1.5

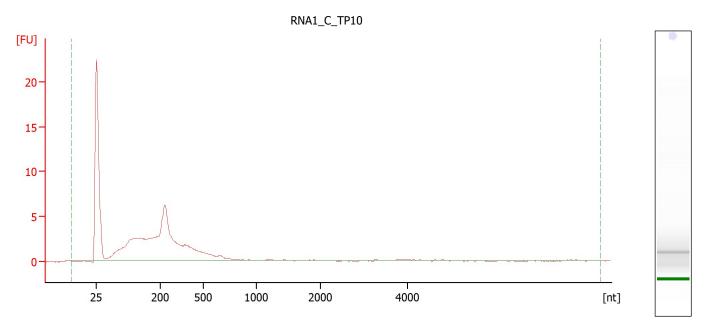
## **Electropherogram Summary Continued ...**



# Overall Results for sample 7: RNA1 C TP9

RNA Area: 80.1 RNA Integrity Number (RIN): 2.2 (B.02.10)
RNA Concentration: 38 ng/µl Result Flagging Color: RNA Ratio [28s / 18s]: 0.0 Result Flagging Label: RIN: 2.20

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



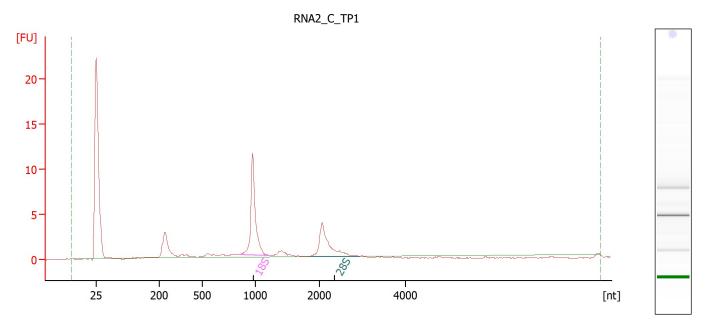
# Overall Results for sample 8: RNA1 C TP10

RNA Area: 76.6 RNA Integrity Number (RIN): 2.2 (B.02.10)
RNA Concentration: 37 ng/µl Result Flagging Color:

rRNA Ratio [28s / 18s]: 0.0 Result Flagging Label: RIN: 2.20

Assay Class: Eukaryote Total RNA Nano Created: 10/21/2020 4:13:59 PM Data Path: C:\...2020-10-21\Eukaryote Total RNA Nano\_2020-10-21\_16-14-00.xad Modified: 10/21/2020 4:46:22 PM

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



Overall Results for sample 9: RNA2 C TP1

RNA Area: 40.0

RNA Concentration: 19 ng/µl

rRNA Ratio [28s / 18s]: 0.6

RNA Integrity Number (RIN):

8.6 (B.02.10, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

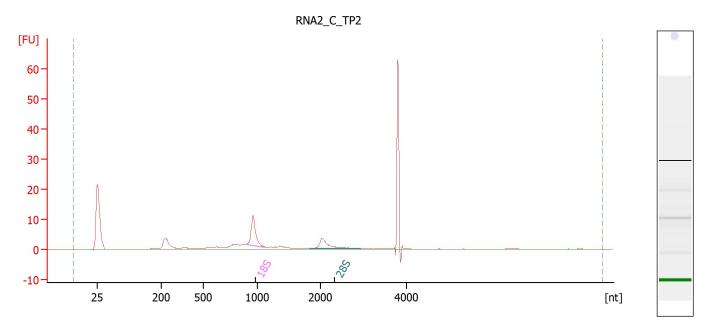
Result Flagging Label: RIN: 8.60

Fragment table for sample 9: RNA2 C TP1

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	849	1,201	15.7	39.1
28S	1,848	2,925	8.8	21.9

Assay Class: Eukaryote Total RNA Nano Created: 10/21/2020 4:13:59 PM C:\...2020-10-21\Eukaryote Total RNA Nano\_2020-10-21\_16-14-00.xad Data Path: Modified: 10/21/2020 4:46:22 PM

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



#### Overall Results for sample 10: RNA2 C TP2

RNA Area: 88.1

**RNA Concentration:** 42 ng/µl

rRNA Ratio [28s / 18s]: 0.6 RNA Integrity Number (RIN):

7.7 (B.02.10, Anomaly Threshold(s) manually adapted)

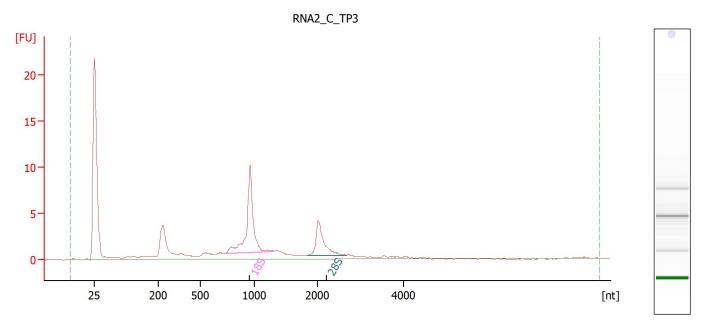
Result Flagging Color:

Result Flagging Label: RIN: 7.70

Fragment table for sample 10: RNA2 C TP2

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	888	1,140	13.2	14.9
28S	1,822	2,939	8.0	9.1

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



# Overall Results for sample 11: RNA2 C TP3

RNA Area: 65.2

RNA Concentration: 31 ng/µl

rRNA Ratio [28s / 18s]: 0.4

RNA Integrity Number (RIN):

Result Flagging Color:

Result Flagging Label:

8.4 (B.02.10,

Anomaly Threshold(s) manually adapted)

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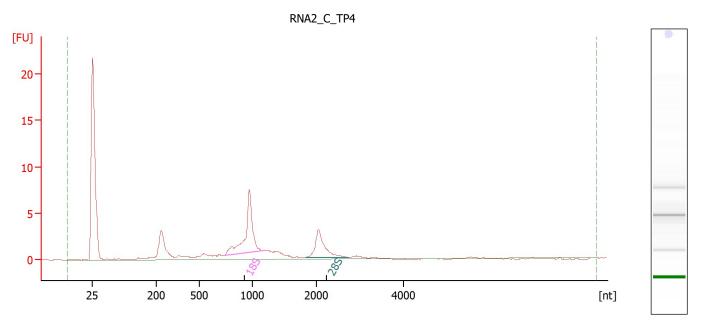
RIN: 8.40

Fragment table for sample 11: RNA2 C TP3

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	722	1,325	17.6	27.1
285	1 839	2 680	7 2	11 1

Assay Class: Eukaryote Total RNA Nano Created: 10/21/2020 4:13:59 PM Data Path: C:\...2020-10-21\Eukaryote Total RNA Nano\_2020-10-21\_16-14-00.xad Modified: 10/21/2020 4:46:22 PM

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



# Overall Results for sample 12: RNA2 C TP4

RNA Area: 49.8 RNA Concentration: 24 ng/µl

rRNA Ratio [28s / 18s]: 0.5

RNA Integrity Number (RIN):

8.6 (B.02.10, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

Result Flagging Label:

RIN: 8.60

Fragment table for sample 12: RNA2 C TP4

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	737	1,178	14.3	28.7
28S	1,824	2,727	6.7	13.5

Assay Class: Data Path: Eukaryote Total RNA Nano C:\...2020-10-21\Eukaryote Total RNA Nano\_2020-10-21\_16-14-00.xad Created: Modified: 10/21/2020 4:13:59 PM 10/21/2020 4:46:22 PM **Gel Image** RNA1\_C\_TP10 [nt] RNA1\_C\_TP9 RNA1\_C\_TP3 RNA1 C TP4 RNA1\_C\_TP6 RNA1\_C\_TP7 RNA1\_C\_TP8 RNA2 C TP2 RNA2\_C\_TP3 RNA2\_C\_TP4 RNAZ C TP1 Ladder ٠ • • ٠ • 4000 -2000 -1000 500 200 25 2 1 3 5 8 9 10 11 12