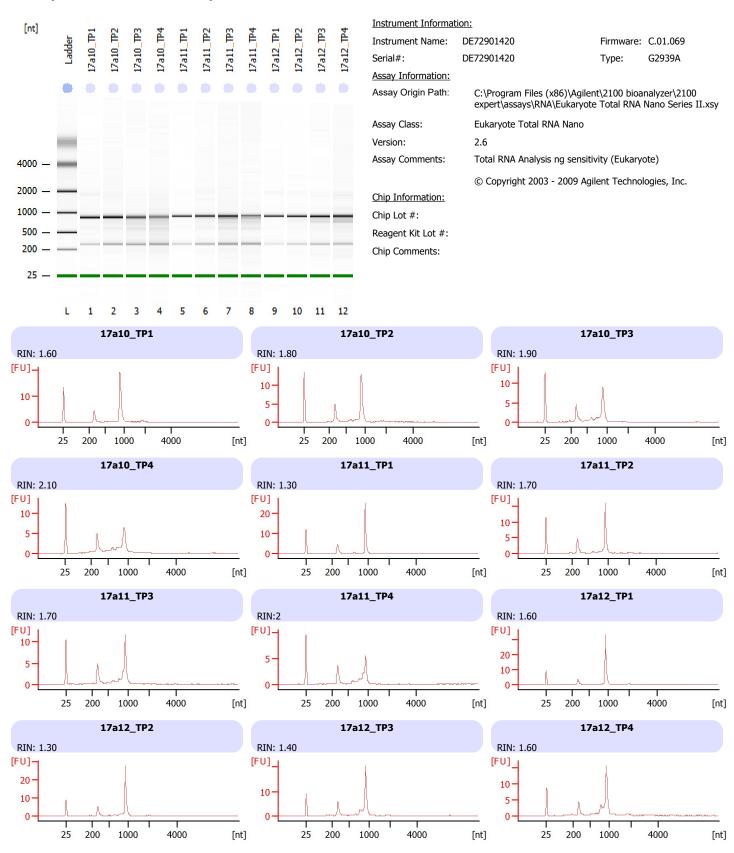
Electrophoresis File Run Summary



Assay Class: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02_15-24-46.xad Eukaryote Total RNA Nano_2020-12-02_15-24-46.xad C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02\Eukaryote Total RNA Na

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Stat Result Labe us	I Result Color
17a10_TP1		✓ RIN: 1.60	
17a10_TP2		✓ RIN: 1.80	
17a10_TP3		RIN: 1.90	
17a10_TP4		✓ RIN: 2.10	
17a11_TP1		✓ RIN: 1.30	
17a11_TP2		RIN: 1.70	
17a11_TP3		RIN: 1.70	
17a11_TP4		RIN:2	
17a12_TP1		✓ RIN: 1.60	
17a12_TP2		✓ RIN: 1.30	
17a12_TP3		✓ RIN: 1.40	
17a12_TP4		RIN: 1.60	
Ladder		All Other	
		Samples	

Chip Lot # Reagent Kit Lot #

Chip Comments:

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02_15-24-46.xad C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02_15-24-46.xad C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02_15-24-46.xad Modified: 12/2/2020 3:59:11 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

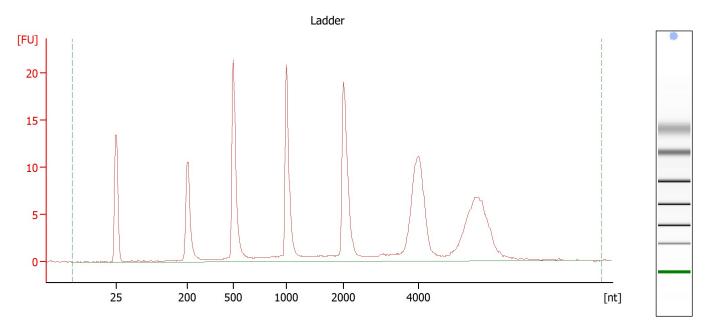
Ladder Peak	Size	
1	25	
2	200	
3	500	
4	1000	
5	2000	
6	4000	

Printed:

12/2/2020 3:59:32 PM

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02_15-24-46.xad Eukaryote Total RNA Nano_2020-12-02_15-24-46.xad C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02\Eukaryote Total RNA Na

Electropherogram Summary

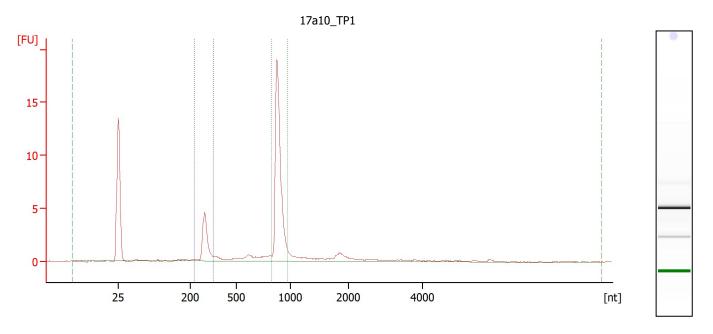


Overall Results for Ladder

RNA Area: 178.8 RNA Concentration: 150 ng/µl Result Flagging Color: Result Flagging Label:

All Other Samples

Electropherogram Summary Continued ...

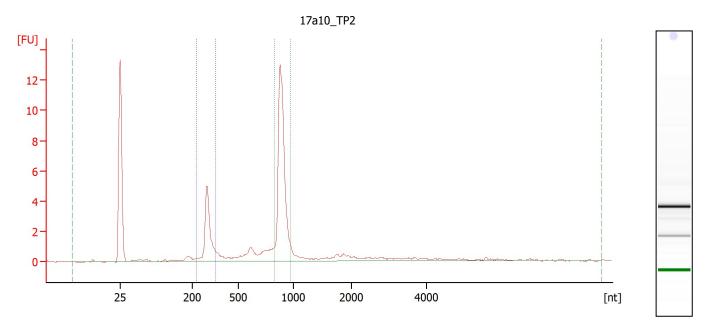


Overall Results for sample 1: 17a10 TP1

RNA Area: 51.3 Result Flagging Color: **RNA Concentration:** 43 ng/µl Result Flagging Label: RIN: 1.60 Corr. Area 1: 7.7 rRNA Ratio [28s / 18s]: 0.0 RNA Integrity Number (RIN): Corr. Area 2: 1.6 (B.02.11, 28.8 Anomaly Threshold(s)

Region table for sample 1: <u>17a10_TP1</u>					
Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	7.7	15	-
Region 2	825	975	28.8	56	

Electropherogram Summary Continued ...



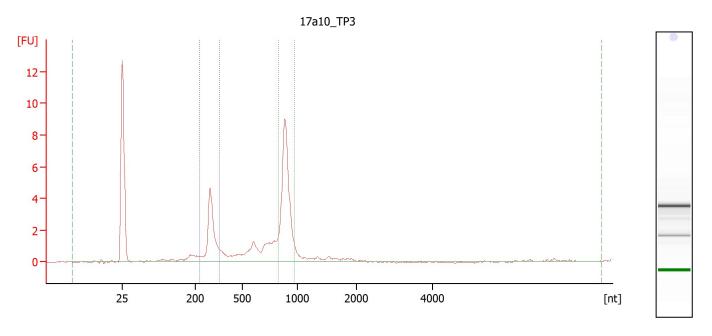
Overall Results for sample 2: <u>17a10_TP2</u>

RNA Area: 47.0 Result Flagging Color: **RNA Concentration:** 39 ng/µl Result Flagging Label: RIN: 1.80 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 8.8 RNA Integrity Number (RIN): Corr. Area 2: 1.8 (B.02.11, 22.3 Anomaly Threshold(s)

Region table for sample 2 : 17a10 TP2

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	8.8	19	
Region 2	825	975	22.3	47	

Electropherogram Summary Continued ...



40

Overall Results for sample 3: 17a10 TP3

 RNA Area:
 44.3
 Result Flagging Color:

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

 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 8.9

 RNA Integrity Number (RIN):
 1.9 (B.02.11,
 Corr. Area 2:
 17.9

Anomaly Threshold(s) manually adapted)

17.9

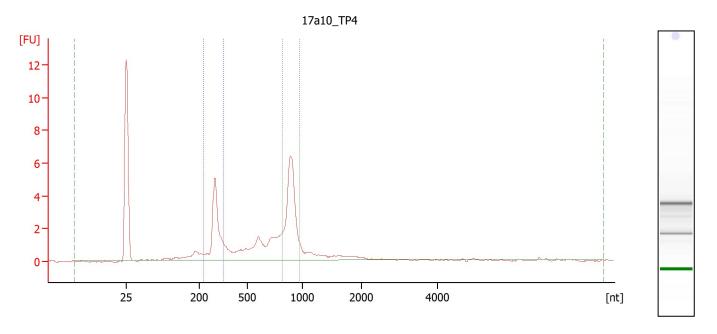
975

Region 2

825

Colo

Electropherogram Summary Continued ...



Overall Results for sample 4: 17a10 TP4

RNA Area: 47.4 Result Flagging Color:

RNA Concentration: 40 ng/µl Result Flagging Label: RIN: 2.10

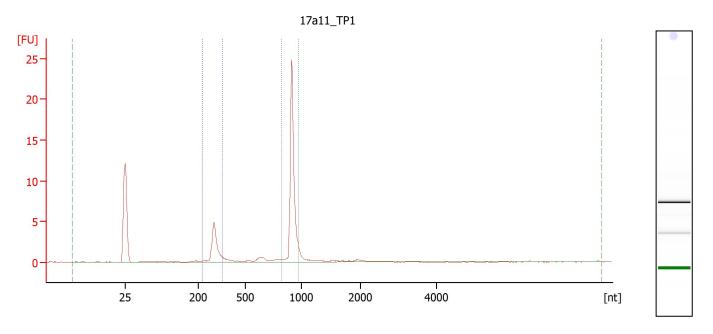
rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 10.0

RNA Integrity Number (RIN): 2.1 (B.02.11, Corr. Area 2: 14.3

Anomaly Threshold(s) manually adapted)

Region table for sample 4: <u>17a10_TP4</u>					
Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	10.0	21	
Region 2	825	975	14.3	30	

Electropherogram Summary Continued ...



Overall Results for sample 5 : 17a11 TP1

 RNA Area:
 40.4
 Result Flagging Color:

 RNA Concentration:
 34 ng/μl
 Result Flagging Label:
 RIN: 1.30

 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 8.1

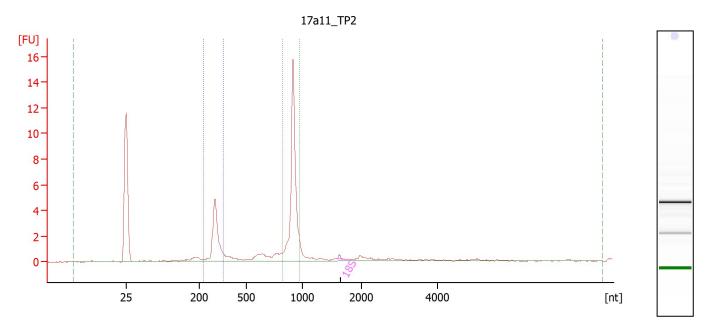
 RNA Integrity Number (RIN):
 1.3 (B.02.11,
 Corr. Area 2:
 24.6

Anomaly Threshold(s) manually adapted)

Region table for sample 5: 17a11 TP1 Name From [nt] To [nt] Corr. Area % of Total Region 1 225 350 8.1 20 Region 2 825 975 24.6 61

Colo

Electropherogram Summary Continued ...



Overall Results for sample 6: 17a11 TP2

RNA Area: 41.9 Result Flagging Color: RNA Concentration: 35 ng/ μ l Result Flagging Label: RIN: 1.70 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 8.5 RNA Integrity Number (RIN): 1.7 (B.02.11, Corr. Area 2: 19.2

Anomaly Threshold(s) manually adapted)

Fragment table for sample 6: 17a11 TP2

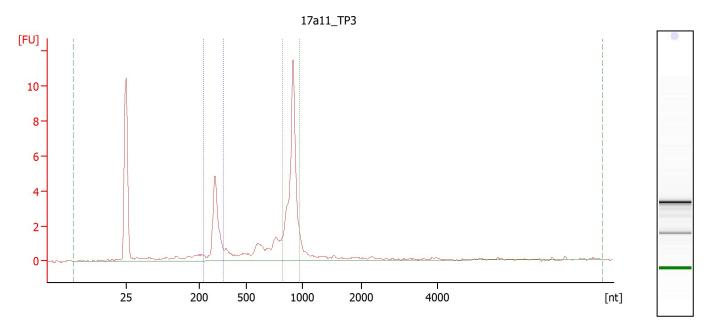
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,571
 1,751
 0.2
 0.5

Region table for sample 6: 17a11 TP2

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	8.5	20	
Region 2	825	975	19.2	46	

Electropherogram Summary Continued ...



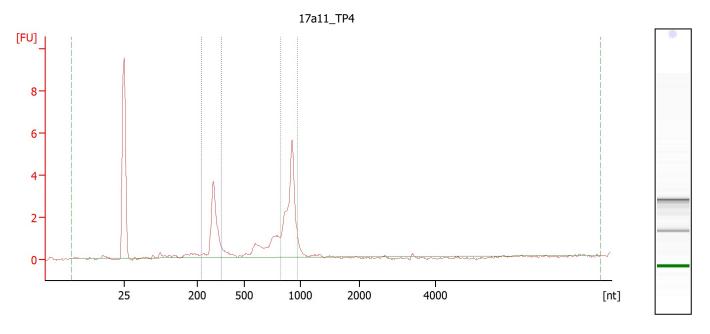
Overall Results for sample 7: 17a11 TP3

RNA Area: 49.8 Result Flagging Color: RIN: 1.70 **RNA Concentration:** 42 ng/µl Result Flagging Label: Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 9.1 RNA Integrity Number (RIN): Corr. Area 2: 1.7 (B.02.11, 18.8 Anomaly Threshold(s)

Region table for sample 7: 17a11 TP3

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	9.1	18	
Region 2	825	975	18.8	38	

Electropherogram Summary Continued ...



Overall Results for sample 8 : 17a11 TP4

RNA Area: 26.5 Result Flagging Color:

RNA Concentration: 22 ng/µl Result Flagging Label: RIN:2

rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 6.6

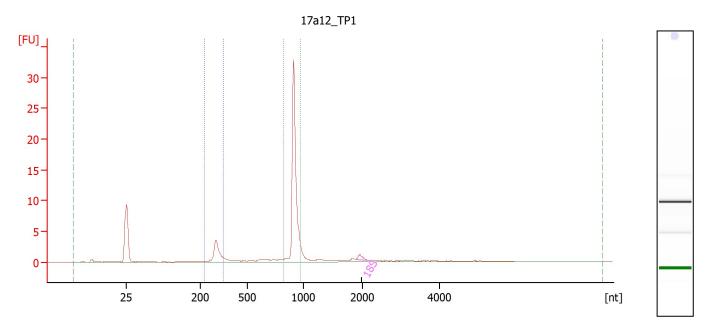
RNA Integrity Number (RIN): 2 (B.02.11, Anomaly Corr. Area 2: 10.6

Threshold(s) manually

adapted)

Region table for sample 8: <u>17a11_TP4</u>					
Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	6.6	25	
Region 2	825	975	10.6	40	

Electropherogram Summary Continued ...



Overall Results for sample 9: 17a12 TP1

RNA Area: 59.5 Result Flagging Color: RNA Concentration: 50 ng/ μ l Result Flagging Label: RIN: 1.60 rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 6.5 RNA Integrity Number (RIN): 1.6 (B.02.11, Corr. Area 2: 33.5

Anomaly Threshold(s) manually adapted)

Fragment table for sample 9: 17a12 TP1

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

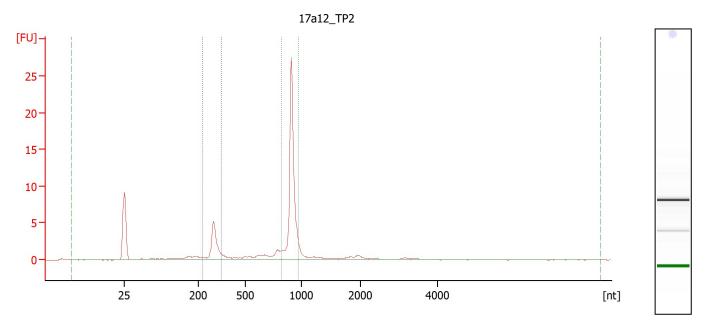
 18S
 1,879
 2,161
 0.9
 1.6

Region table for sample 9: 17a12 TP1

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	6.5	11	
Region 2	825	975	33.5	56	

Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02_15-24-46.xad Assay Class: Created: 12/2/2020 3:24:45 PM Data Path: Modified: 12/2/2020 3:59:11 PM

Electropherogram Summary Continued ...



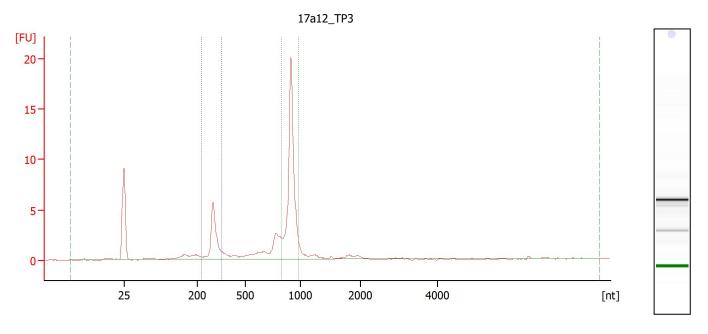
Overall Results for sample 10: 17a12_TP2

RNA Area: 60.6 Result Flagging Color: **RNA Concentration:** 51 ng/µl Result Flagging Label: RIN: 1.30 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 9.3 RNA Integrity Number (RIN): Corr. Area 2: 1.3 (B.02.11, 32.1 Anomaly Threshold(s)

Region table	for sample 10	: <u>17a12</u>	<u>TP2</u>
Name	From [nt]	To [nt]	Corr. Area

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo
					<u>r</u>
Region 1	225	350	9.3	15	
Region 2	825	975	32.1	53	

Electropherogram Summary Continued ...



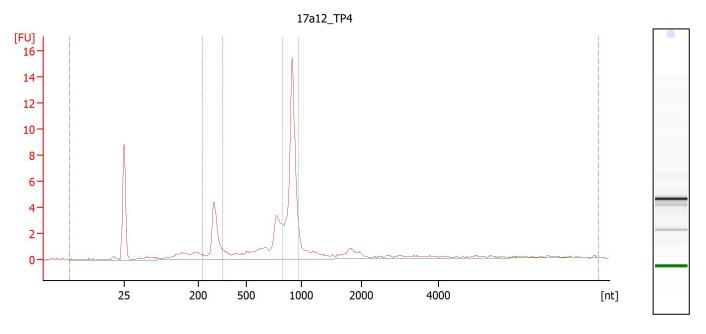
Overall Results for sample 11: 17a12 TP3

RNA Area: 61.3 Result Flagging Color: RIN: 1.40 **RNA Concentration:** 51 ng/µl Result Flagging Label: Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 10.1 RNA Integrity Number (RIN): Corr. Area 2: 1.4 (B.02.11, 29.1 Anomaly Threshold(s) manually adapted)

Region table for sample 11: 17a12 TP3

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	10.1	16	
Region 2	825	975	29.1	48	

Electropherogram Summary Continued ...



Overall Results for sample 12: 17a12 TP4

RNA Area: 65.1 Result Flagging Color: **RNA Concentration:** 55 ng/µl Result Flagging Label: RIN: 1.60 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 8.4 RNA Integrity Number (RIN): Corr. Area 2: 1.6 (B.02.11, 25.6 Anomaly Threshold(s)

manually adapted)

Region table for sample 12: 17a12 TP4

Name	From [nt]	To [nt]	Corr. Area	% of Total	Colo r
Region 1	225	350	8.4	13	
Region 2	825	975	25.6	39	

Assay Class: Data Path: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02_15-24-46.xad Created: Modified: 12/2/2020 3:24:45 PM 12/2/2020 3:59:11 PM **Gel Image** [nt] 17a10_TP2 17a10_TP3 17a10_TP4 17a11_TP3 17a11_TP4 17a12_TP2 17a12_TP3 17a12_TP4 17a11_TP1 17a10_TP1 17a12_TP1 Ladder ø • • • • ø • • ø ø • 4000 2000 1000 500 200 25 3 5 10 11 12