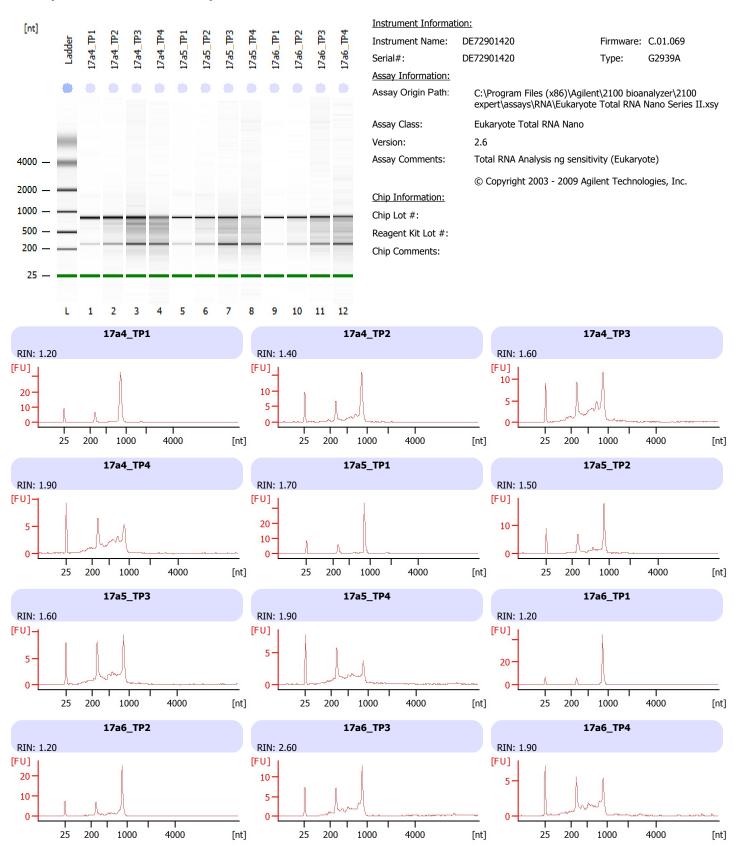
Electrophoresis File Run Summary



Assay Class: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02_14-38-16.xad Created: 12/2/2020 2:38:15 PM Modified: 12/2/2020 3:36:05 PM

Electrophoresis File Run Summary (Chip Summary)

| Sample Name | Sample Comment | Stat Result Labe | Result Color |
|-------------|-------------------|---|--------------|
| 17a4_TP1 | | RIN: 1.20 | |
| 17a4_TP2 | | ✓ RIN: 1.40 | |
| 17a4_TP3 | | ✓ RIN: 1.60 | |
| 17a4_TP4 | | RIN: 1.90 | |
| 17a5_TP1 | | RIN: 1.70 | |
| 17a5_TP2 | | ✓ RIN: 1.50 | |
| 17a5_TP3 | | ✓ RIN: 1.60 | |
| 17a5_TP4 | | ✓ RIN: 1.90 | |
| 17a6_TP1 | | ✓ RIN: 1.20 | |
| 17a6_TP2 | | RIN: 1.20 | |
| 17a6_TP3 | | RIN: 2.60 | |
| 17a6_TP4 | | ✓ RIN: 1.90 | |
| Ladder | | All Other Samples | |

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

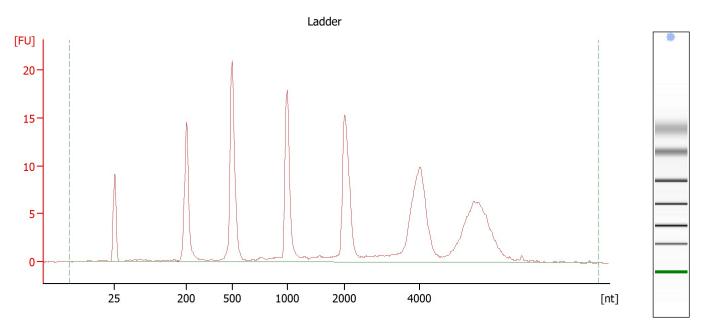
Filter Width [s]: 0.5 Polynomial Order: 4

Ladder

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

Assay Class: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02_14-38-16.xad Created: 12/2/2020 2:38:15 PM Modified: 12/2/2020 3:36:05 PM

Electropherogram Summary

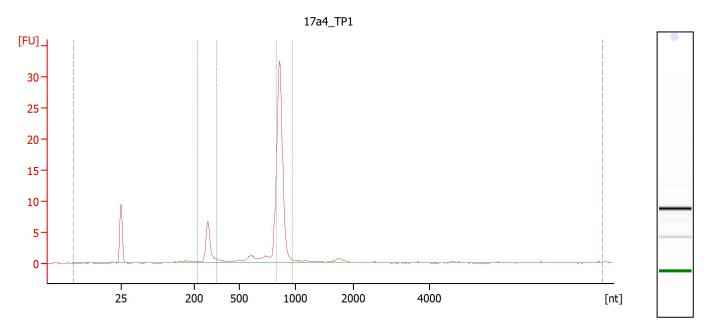


Overall Results for Ladder

RNA Area: 188.4 RNA Concentration: 150 ng/µl Result Flagging Color: Result Flagging Label:

All Other Samples

Electropherogram Summary Continued ...

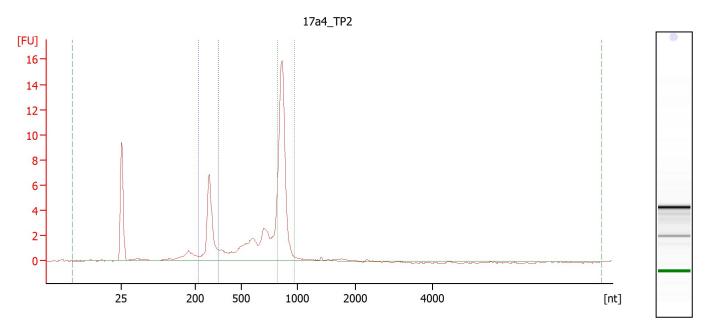


Overall Results for sample 1 : 17a4 TP1

RNA Area: 79.7 Result Flagging Color: **RNA Concentration:** 63 ng/µl Result Flagging Label: RIN: 1.20 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 10.6 RNA Integrity Number (RIN): Corr. Area 2: 1.2 (B.02.11, 51.1 Anomaly Threshold(s)

| Region table for sample 1: <u>17a4_TP1</u> | | | | | |
|--|-----------|---------|------------|------------|-----------|
| Name | From [nt] | To [nt] | Corr. Area | % of Total | Colo r |
| Region 1 | 225 | 350 | 10.6 | 13 | |
| Region 2 | 825 | 975 | 51.1 | 64 | |

Electropherogram Summary Continued ...



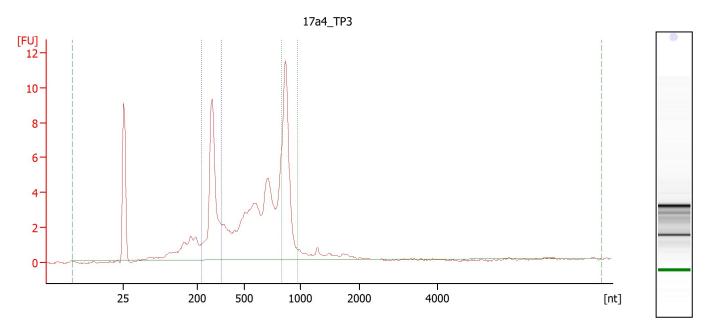
Overall Results for sample 2: <u>17a4_TP2</u>

RNA Area: 68.6 Result Flagging Color: **RNA Concentration:** 55 ng/µl Result Flagging Label: RIN: 1.40 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 12.4 RNA Integrity Number (RIN): Corr. Area 2: 1.4 (B.02.11, 27.1 Anomaly Threshold(s) manually adapted)

Region table for sample 2 : 17a4 TP2

| Name | From [nt] | To [nt] | Corr. Area | % of Total | Colo r |
|----------|-----------|---------|------------|------------|-----------|
| Region 1 | 225 | 350 | 12.4 | 18 | |
| Region 2 | 825 | 975 | 27.1 | 39 | |

Electropherogram Summary Continued ...



Overall Results for sample 3: 17a4 TP3

 RNA Area:
 98.6
 Result Flagging Color:

 RNA Concentration:
 78 ng/μl
 Result Flagging Label:
 RIN: 1.60

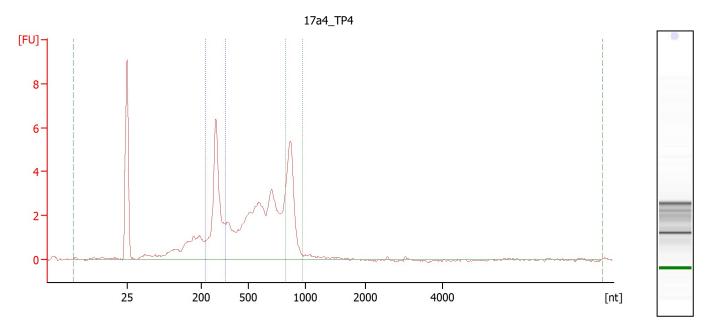
 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 19.6

 RNA Integrity Number (RIN):
 1.6 (B.02.11, Approach Threshold(s))
 Corr. Area 2:
 21.8

Anomaly Threshold(s) manually adapted)

| Region table for sample 3: <u>17a4_TP3</u> | | | | | |
|--|-----------|---------|------------|------------|------|
| Name | From [nt] | To [nt] | Corr. Area | % of Total | Colo |
| Region 1 | 225 | 350 | 19.6 | 20 | |
| Region 2 | 825 | 975 | 21.8 | 22 | |

Electropherogram Summary Continued ...



Overall Results for sample 4: <u>17a4_TP4</u>

RNA Area: 67.6 Result Flagging Color: **RNA Concentration:** 54 ng/µl Result Flagging Label: RIN: 1.90 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 14.1 RNA Integrity Number (RIN): Corr. Area 2: 1.9 (B.02.11, 10.8 Anomaly Threshold(s)

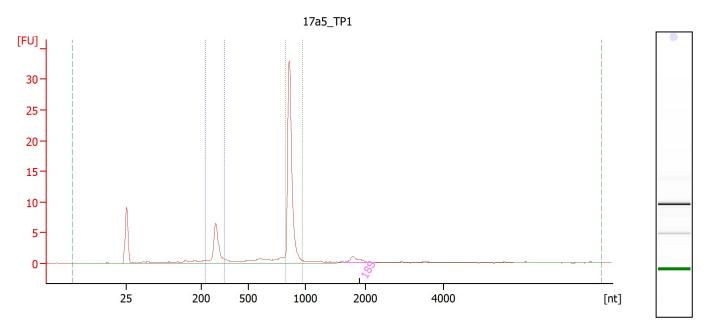
manually adapted)

Region table for sample 4: <u>17a4 TP4</u>

Name From [nt] To [nt] Corr.

| Name | From [nt] | To [nt] | Corr. Area | % of Total | Colo r |
|----------|-----------|---------|------------|------------|-----------|
| Region 1 | 225 | 350 | 14.1 | 21 | |
| Region 2 | 825 | 975 | 10.8 | 16 | |

Electropherogram Summary Continued ...



Overall Results for sample 5 : 17a5 TP1

RNA Area: 65.8 Result Flagging Color: RNA Concentration: 52 ng/µl Result Flagging Label: RIN: 1.70 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 11.2 RNA Integrity Number (RIN): 1.7 (B.02.11, Corr. Area 2: 36.7 Anomaly Threshold(s)

manually adapted)

Fragment table for sample 5 : <u>17a5_TP1</u>

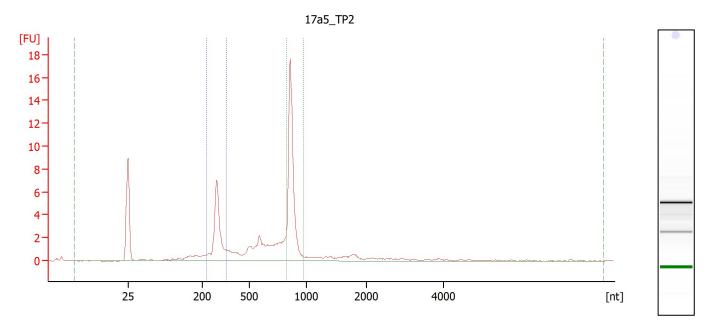
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,656
 2,240
 1.9
 2.9

Region table for sample 5: 17a5 TP1

| Name | From [nt] | To [nt] | Corr. Area | % of Total | Colo r |
|----------|-----------|---------|------------|------------|-----------|
| Region 1 | 225 | 350 | 11.2 | 17 | |
| Region 2 | 825 | 975 | 36.7 | 56 | |

Electropherogram Summary Continued ...



Overall Results for sample 6 : 17a5 TP2

 RNA Area:
 64.7
 Result Flagging Color:

 RNA Concentration:
 51 ng/µl
 Result Flagging Label:
 RIN: 1.50

 rRNA Ratio [28s / 18s]:
 0.0
 Corr. Area 1:
 12.8

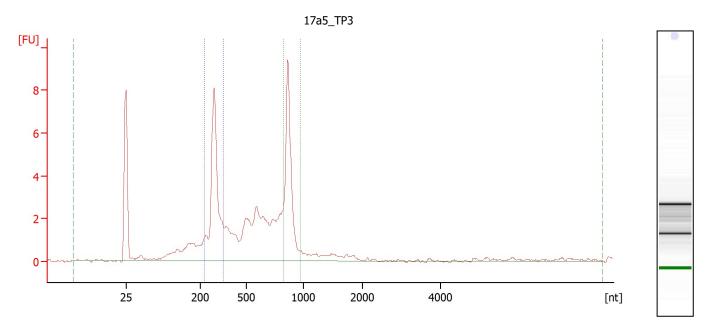
 RNA Integrity Number (RIN):
 1.5 (B.02.11,
 Corr. Area 2:
 22.5

Anomaly Threshold(s) manually adapted)

| Region table for sample 6: <u>17a5_TP2</u> | | | | | |
|--|-----------|---------|------------|------------|------|
| Name | From [nt] | To [nt] | Corr. Area | % of Total | Colo |
| Region 1 | 225 | 350 | 12.8 | 20 | |
| Region 2 | 825 | 975 | 22.5 | 35 | |

Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02_14-38-16.xad Assay Class: Created: 12/2/2020 2:38:15 PM Data Path: Modified: 12/2/2020 3:36:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 7: 17a5_TP3

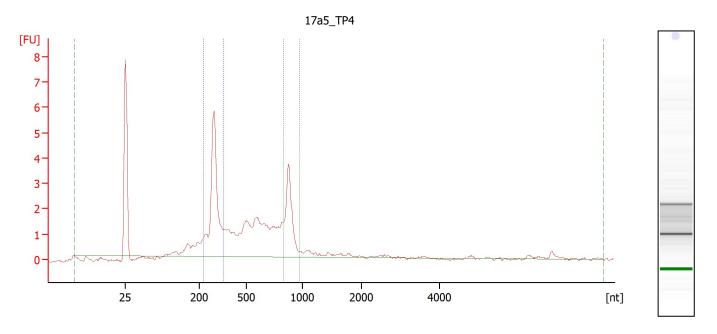
RNA Area: 71.6 Result Flagging Color: **RNA Concentration:** 57 ng/µl Result Flagging Label: RIN: 1.60 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 17.4 RNA Integrity Number (RIN): Corr. Area 2: 1.6 (B.02.11, 14.4 Anomaly Threshold(s)

manually adapted)

| Region table for sample 7: <u>17a5_TP3</u> | | | | | | |
|--|-----------|---------|------------|------------|--|--|
| Name | From [nt] | To [nt] | Corr. Area | % of Total | | |
| Region 1 | 225 | 350 | 17.4 | 24 | | |
| Region 2 | 825 | 975 | 14.4 | 20 | | |

Colo

Electropherogram Summary Continued ...

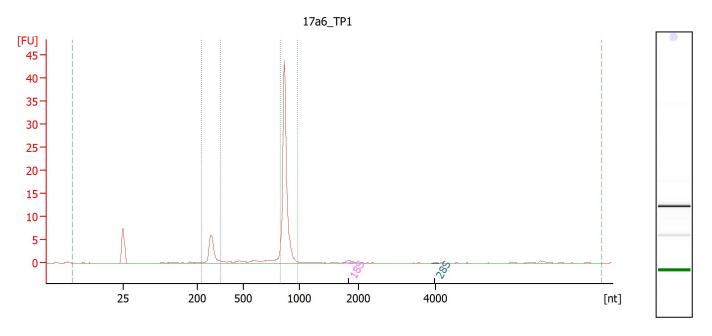


Overall Results for sample 8 : 17a5 TP4

RNA Area: 42.5 Result Flagging Color: **RNA Concentration:** 34 ng/µl Result Flagging Label: RIN: 1.90 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 11.9 RNA Integrity Number (RIN): Corr. Area 2: 1.9 (B.02.11, 6.5 Anomaly Threshold(s)

| Region table for sample 8: <u>17a5_TP4</u> | | | | | | |
|--|-----------|---------|------------|------------|-----------|--|
| Name | From [nt] | To [nt] | Corr. Area | % of Total | Colo r | |
| Region 1 | 225 | 350 | 11.9 | 28 | | |
| Region 2 | 825 | 975 | 6.5 | 15 | | |

Electropherogram Summary Continued ...



Overall Results for sample 9: 17a6 TP1

RNA Area: 69.8 Result Flagging Color: RNA Concentration: 56 ng/ μ l Result Flagging Label: RIN: 1.20 rRNA Ratio [28s / 18s]: 0.2 Corr. Area 1: 9.9 RNA Integrity Number (RIN): 1.2 (B.02.11, Corr. Area 2: 48.8

Anomaly Threshold(s) manually adapted)

Fragment table for sample 9: 17a6 TP1

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

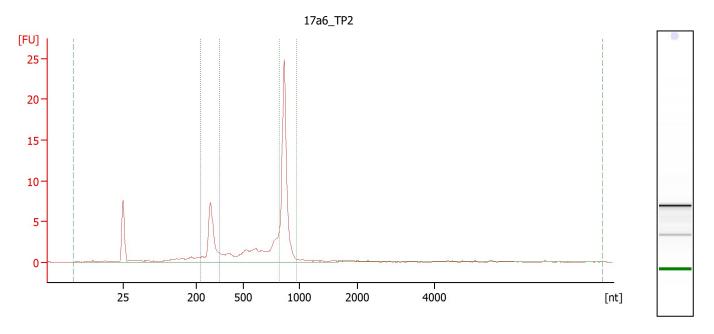
 18S
 1,701
 1,936
 0.7
 1.0

 28S
 3,885
 4,079
 0.1
 0.2

Region table for sample 9: 17a6 TP1

| 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1 | | | | | |
|--|-----------|---------|------------|------------|------|
| Name | From [nt] | To [nt] | Corr. Area | % of Total | Colo |
| Region 1 | 225 | 350 | 9.9 | 14 | r |
| Region 2 | 825 | 975 | 48.8 | 70 | |

Electropherogram Summary Continued ...



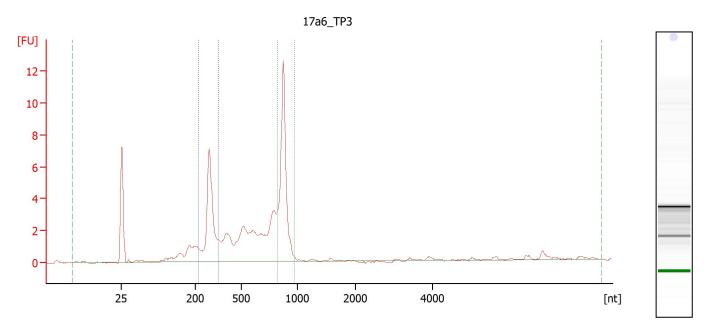
Overall Results for sample 10 : 17a6 TP2

RNA Area: 78.2 Result Flagging Color: **RNA Concentration:** 62 ng/µl Result Flagging Label: RIN: 1.20 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 14.0 RNA Integrity Number (RIN): Corr. Area 2: 1.2 (B.02.11, 31.8 Anomaly Threshold(s)

Region table for sample 10 : 17a6 TP2

| Name | From [nt] | To [nt] | Corr. Area | % of Total | Colo r |
|----------|-----------|---------|------------|------------|-----------|
| Region 1 | 225 | 350 | 14.0 | 18 | |
| Region 2 | 825 | 975 | 31.8 | 41 | |

Electropherogram Summary Continued ...



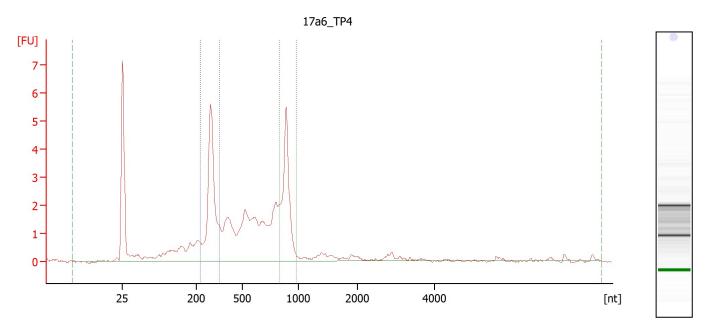
Overall Results for sample 11: 17a6 TP3

RNA Area: 69.2 Result Flagging Color: **RNA Concentration:** 55 ng/µl Result Flagging Label: RIN: 2.60 Corr. Area 1: rRNA Ratio [28s / 18s]: 0.0 14.2 RNA Integrity Number (RIN): Corr. Area 2: 2.6 (B.02.11, 18.5 Anomaly Threshold(s)

Region table for sample 11: 17a6 TP3

| Name | From [nt] | To [nt] | Corr. Area | % of Total | Colo r |
|----------|-----------|---------|------------|------------|-----------|
| Region 1 | 225 | 350 | 14.2 | 21 | |
| Region 2 | 825 | 975 | 18.5 | 27 | |

Electropherogram Summary Continued ...



Overall Results for sample 12: 17a6 TP4

RNA Area: 54.3 Result Flagging Color:

RNA Concentration: 43 ng/µl Result Flagging Label: RIN: 1.90

rRNA Ratio [28s / 18s]: 0.0 Corr. Area 1: 11.9

RNA Integrity Number (RIN): 1.9 (B.02.11, Corr. Area 2: 9.7

Anomaly Threshold(s) manually adapted)

| Region table | for sample 12 : | 17a6 TP4 |
|--------------|-----------------|----------|
| | | |

| Name | From [nt] | To [nt] | Corr. Area | % of Total | Colo |
|----------|-----------|---------|------------|------------|------|
| Region 1 | 225 | 350 | 11.9 | 22 | |
| Region 2 | 825 | 975 | 9.7 | 18 | |

Assay Class: Data Path: Eukaryote Total RNA Nano C:\...2020-12-02\Eukaryote Total RNA Nano_2020-12-02_14-38-16.xad Created: Modified: 12/2/2020 2:38:15 PM 12/2/2020 3:36:05 PM **Gel Image** [nt] 17a6_TP4 17a4_TP2 17a5_TP3 17a6_TP1 17a6_TP2 Ladder • ۰ ۰ • ٠ ٠ • ۰ • 4000 -2000 -1000 500 200 25 -2 3 5 1 6 10 11 12