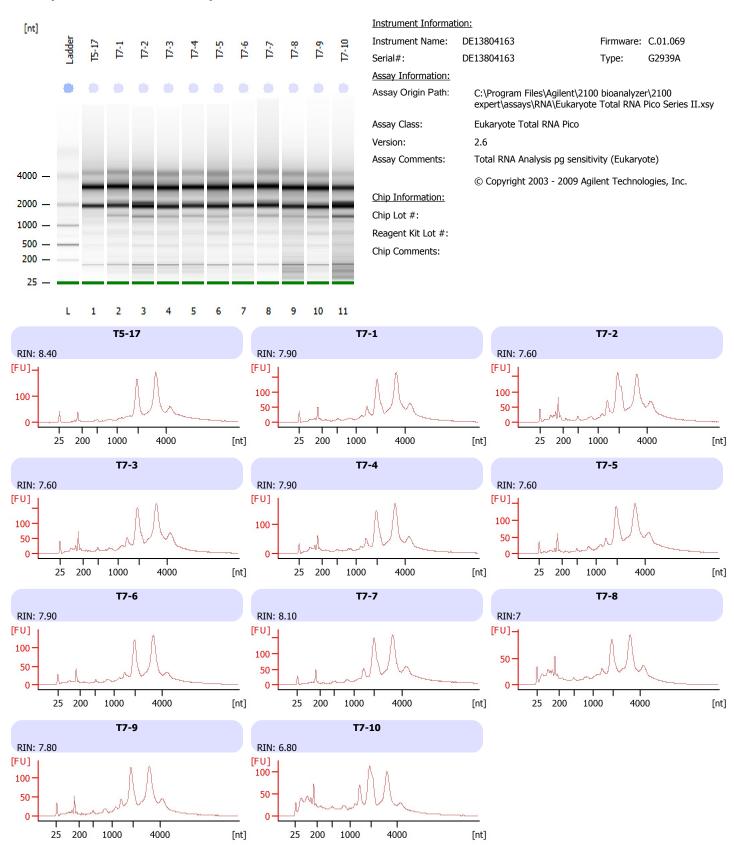
Electrophoresis File Run Summary



Printed:

6/24/2016 12:24:47 PM

Assay Class: Eukaryote Total RNA Pico C:\...Eukaryote Total RNA Pico_DE13804163_2016-06-23_17-33-16.xad Created: 6/23/2016 5:33:15 PM Modified: 6/23/2016 5:58:10 PM

Electrophoresis File Run Summary (Chip Summary)

| Sample Name | Sample Comment | Status | Result Label | Result Color |
|-------------|----------------|----------|-------------------|--------------|
| T5-17 | | ~ | RIN: 8.40 | |
| T7-1 | | ~ | RIN: 7.90 | |
| T7-2 | | ~ | RIN: 7.60 | |
| T7-3 | | ~ | RIN: 7.60 | |
| T7-4 | | ~ | RIN: 7.90 | |
| T7-5 | | ~ | RIN: 7.60 | |
| T7-6 | | ~ | RIN: 7.90 | |
| T7-7 | | ~ | RIN: 8.10 | |
| T7-8 | | ~ | RIN:7 | |
| T7-9 | | ~ | RIN: 7.80 | |
| T7-10 | | ~ | RIN: 6.80 | |
| Ladder | | ~ | All Other Samples | |

Chip Lot # Reagent Kit Lot #

Chip Comments:

Page 3 of 7

Printed: 6/24/2016 12:24:47 PM

Electrophoresis Assay Details

General Analysis Settings

Number of Available Sample and Ladder Wells (Max.): 12

Minimum Visible Range [s]: 17

Maximum Visible Range [s]: 70

Start Analysis Time Range [s]: 18

End Analysis Time Range [s]: 69

Ladder Concentration [pg/µl]: 1000

Lower Marker Concentration [pg/µl]: 0

Upper Marker Concentration [pg/µl]: 0

Used Lower Marker for Quantitation

Standard Curve Fit is Logarithmic

Show Data Aligned to Lower Marker

Integrator Settings

Integration Start Time [s]: 18 Integration End Time [s]: 69 Slope Threshold: 0.6 Height Threshold [FU]: 0.5 Area Threshold: 0.2 Width Threshold [s]: 0.3 Baseline Plateau [s]: 6

Filter Settings

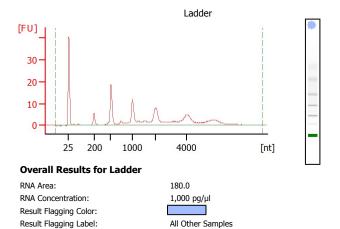
Filter Width [s]: 0.5 Polynomial Order: 4

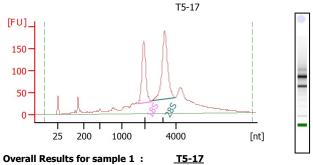
Ladder

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

Assay Class: Eukaryote Total RNA Pico C:\...Eukaryote Total RNA Pico_DE13804163_2016-06-23_17-33-16.xad Created: 6/23/2016 5:33:15 PM Modified: 6/23/2016 5:58:10 PM

Electropherogram Summary





RNA Area: 2,430.7

RNA Concentration: 13,503 pg/µl

rRNA Ratio [28s / 18s]: 1.4

RNA Integrity Number (RIN): 8.4 (B.02.08)

Result Flagging Color:

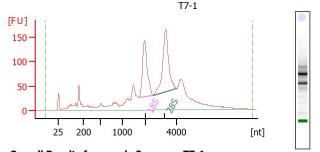
Result Flagging Label: RIN: 8.40

Fragment table for sample 1: <u>T5-17</u>

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

 18S
 1,684
 2,432
 354.3
 14.6

 28S
 2,459
 3,891
 507.6
 20.9



 T7-1

 RNA Area:
 2,594.4

 RNA Concentration:
 14,413 pg/μl

 rRNA Ratio [28s / 18s]:
 1.3

 RNA Integrity Number (RIN):
 7.9 (B.02.08)

 Result Flagging Color:
 RIN: 7.90

 Result Flagging Label:
 RIN: 7.90

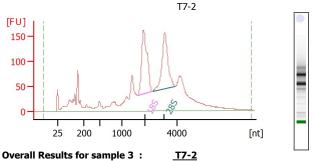
Fragment table for sample 2: <u>T7-1</u>

| Name | Start Size [nt] | Ena Size [nt] | Area | % of total Are |
|------|-----------------|---------------|-------|----------------|
| 400 | 4.740 | 0.400 | 040.4 | 40.4 |

436.7

16.8

4,003



RNA Area: 3,100.9
RNA Concentration: 17,226 pg/µl
rRNA Ratio [28s / 18s]: 0.9

RNA Integrity Number (RIN): 7.6 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

Result Flagging Label:

RIN: 7.60

Fragment table for sample 3: <u>T7-2</u>

Name Start Size [nt] End Size [nt] Area

| 18S | 1,671 | 2,433 | 444.1 | 14.3 |
|-----|-------|-------|-------|------|
| 28S | 2,459 | 3,896 | 381.6 | 12.3 |

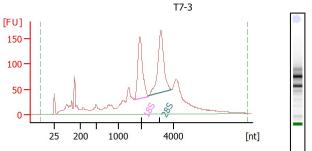
28S

2,513

% of total Area

Assay Class: Eukaryote Total RNA Pico Created: 6/23/2016 5:33:15 PM C:\...Éukaryote Total RNA Pico_DE13804163_2016-06-23_17-33-16.xad Data Path: Modified: 6/23/2016 5:58:10 PM

Electropherogram Summary Continued ...



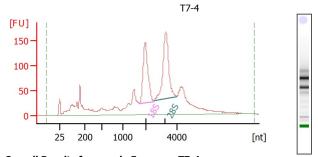
Overall Results for sample 4: <u>T7-3</u> RNA Area: 2,924.7 16,248 pg/µl RNA Concentration: rRNA Ratio [28s / 18s]: 1.2 7.6 (B.02.08) RNA Integrity Number (RIN): Result Flagging Color:

Result Flagging Label: Fragment table for sample 4: <u>T7-3</u>

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

RIN: 7.60

18S 1,671 2,392 367.0 12.5 28S 2,419 3.869 429.7 14.7

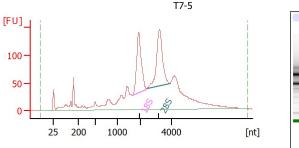


Overall Results for sample 5: <u>T7-4</u> RNA Area: 2,654.1 14,744 pg/µl RNA Concentration: rRNA Ratio [28s / 18s]: 1.2 7.9 (B.02.08) RNA Integrity Number (RIN): Result Flagging Color: Result Flagging Label: RIN: 7.90

Fragment table for sample 5: <u>T7-4</u>

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

18S 1,690 2,459 373.1 14.1 285 2.486 3.979 441.8 16.6



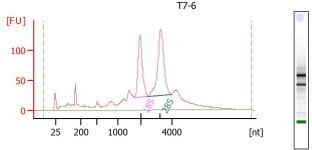
Overall Results for sample 6: <u>T7-5</u> 2,613.7 RNA Area: RNA Concentration: 14,520 pg/µl

rRNA Ratio [28s / 18s]: 1.1 7.6 (B.02.08) RNA Integrity Number (RIN): Result Flagging Color: RIN: 7.60

Result Flagging Label: Fragment table for sample 6: <u>T7-5</u>

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

18S 1,665 2,418 327.2 12.5 28S 2,445 3,921 357.6 13.7

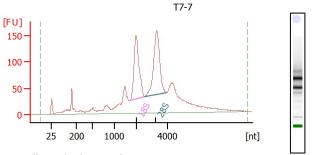


Overall Results for sample 7: T7-6 RNA Area: 2,066.7 RNA Concentration: 11,481 pg/µl rRNA Ratio [28s / 18s]: 1.4 RNA Integrity Number (RIN): 7.9 (B.02.08) Result Flagging Color: Result Flagging Label: RIN: 7.90

Fragment table for sample 7: <u>T7-6</u>

Name Start Size [nt] End Size [nt] Area % of total Area 18S 1,718 2,444 280.7 13.6 28S 2,472 3,998 391.9 19.0

Electropherogram Summary Continued ...



 Overall Results for sample 8 :
 T7-7

 RNA Area:
 2,549.2

 RNA Concentration:
 14,161 pg/µl

rRNA Ratio [28s / 18s]: 1.1

RNA Integrity Number (RIN): 8.1 (B.02.08, Anomaly Threshold(s) manually adapted)
Result Flagging Color:

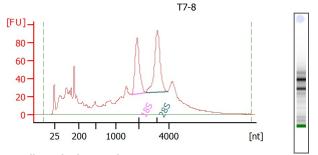
Result Flagging Label: RIN: 8.10

Fragment table for sample 8: T7-7

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

 18S
 1,697
 2,513
 391.4
 15.4

 28S
 2,540
 3,977
 432.7
 17.0

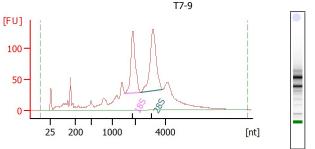


Fragment table for sample 9 : $\underline{T7-8}$

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

 18S
 1,665
 2,429
 185.0
 8.8

 28S
 2,457
 3,890
 230.1
 10.9



Overall Results for sample 10 : T7-9

RNA Area: 2,289.4 RNA Concentration: 12,718 pg/µl rRNA Ratio [28s / 18s]: 1.3

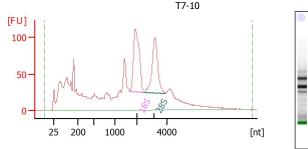
RNA Integrity Number (RIN): 7.8 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

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

 18S
 1,542
 2,359
 292.3
 12.8

 28S
 2,387
 3,823
 369.5
 16.1



Overall Results for sample 11 : T7-10

 RNA Area:
 2,905.8

 RNA Concentration:
 16,143 pg/μl

 rRNA Ratio [28s / 18s]:
 0.8

 RNA Integrity Number (RIN):
 6.8 (B.02.08)

 Result Flagging Color:
 RIN: 6.80

Fragment table for sample 11 : $\underline{\mathsf{T7-10}}$

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

 18S
 1,663
 2,456
 330.8
 11.4

 28S
 2,484
 3,898
 258.7
 8.9

