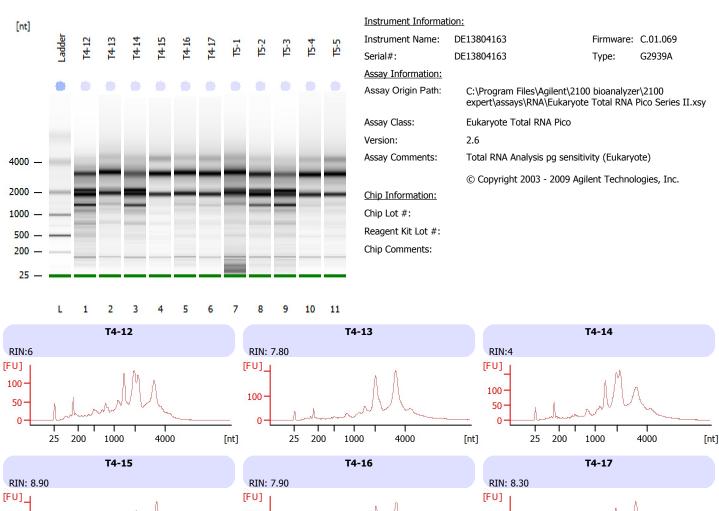
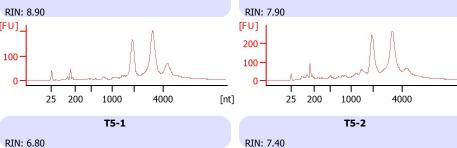
## **Electrophoresis File Run Summary**

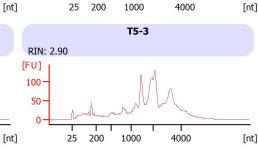




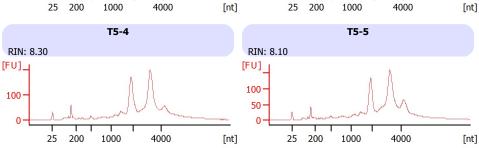
[FU].

100

50



100



[FU]

50

Printed: 6/24/2016 12:24:20 PM

Assay Class: Eukaryote Total RNA Pico C:\...Eukaryote Total RNA Pico\_DE13804163\_2016-06-23\_16-24-04.xad Created: 6/23/2016 4:24:04 PM Modified: 6/23/2016 6:02:43 PM

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

Sample Name	Sample Comment	Status	Result Label	Result Color
T4-12		~	RIN:6	
T4-13		~	RIN: 7.80	
T4-14		<b>~</b>	RIN:4	
T4-15		~	RIN: 8.90	
T4-16		~	RIN: 7.90	
T4-17		<b>~</b>	RIN: 8.30	
T5-1		~	RIN: 6.80	
T5-2		~	RIN: 7.40	
T5-3		<b>~</b>	RIN: 2.90	
T5-4		~	RIN: 8.30	
T5-5		<b>~</b>	RIN: 8.10	
Ladder		<b>~</b>	All Other Samples	

Chip Lot # Reagent Kit Lot #

**Chip Comments:** 

Page 3 of 7

Printed: 6/24/2016 12:24:20 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**

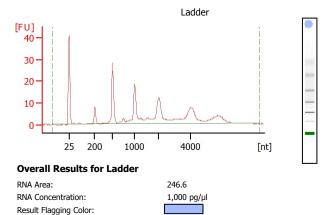
Filter Width [s]: 0.5 Polynomial Order: 4

#### Ladder

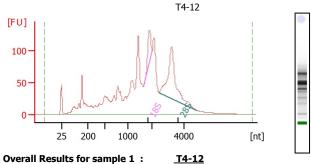
Ladder Peak	Size
1	25
2	200
3	500
4	1000
5	2000
6	4000

#### **Electropherogram Summary**

Result Flagging Label:



All Other Samples



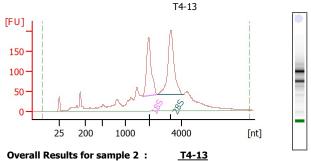
RNA Area: 2,817.2
RNA Concentration: 11,424 pg/µl
rRNA Ratio [28s / 18s]: 3.5
RNA Integrity Number (RIN): 6 (B.02.08)
Result Flagging Color:
Result Flagging Label: RIN:6

Fragment table for sample 1 : T4-12

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

 18S
 1,703
 2,040
 98.0
 3.5

 28S
 2,470
 4,805
 344.4
 12.2



RNA Area: 2,994.9

RNA Concentration: 12,144 pg/µl

RNA Ratio [28s / 18s]: 1.3

RNA Integrity Number (RIN): 7.8 (B.02.08)

Result Flagging Color:

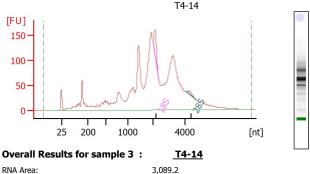
Result Flagging Label: RIN: 7.80

Fragment table for sample 2 : <u>T4-13</u>

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

18S	1,738	2,509	413.3	13.8
200	2.549	4 166	545.6	10.2

% of total Area



RNA Area: 3,089.2
RNA Concentration: 12,527 pg/µl
rRNA Ratio [28s / 18s]: 0.2
RNA Integrity Number (RIN): 4 (B.02.08)
Result Flagging Color:
Result Flagging Label: RIN:4

Fragment table for sample 3: T4-14

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

Itallic	Start Size [iit]	Liid Size [iit]	Aica	70 OI total Alea
18S	2,032	2,393	80.9	2.6
28S	4,071	4,678	14.7	0.5

% of total Area

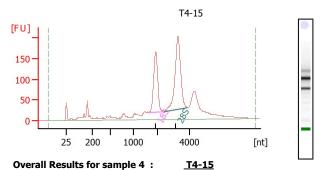
Page 5 of 7

Assay Class: Eukaryote Total RNA Pico C:\...Eukaryote Total RNA Pico\_DE13804163\_2016-06-23\_16-24-04.xad C:\...Eukaryote Total RNA Pico\_DE13804163\_2016-06-23\_16-24-04.xad C:\...Eukaryote Total RNA Pico\_DE13804163\_2016-06-23\_16-24-04.xad Modified: 6/23/2016 6:02:43 PM

285

2.484

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



RNA Area: 2,203.8

RNA Concentration: 8,937 pg/μl

rRNA Ratio [28s / 18s]: 1.5

RNA Integrity Number (RIN): 8.9 (B.02.08)

Result Flagging Color:

Result Flagging Label: RIN: 8.90

Fragment table for sample 4: <u>T4-15</u>

18S

28S

1,642

2,420

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

369.1

558.5

16.7

25.3

2,394

3.823

[FU] 250 200 150 100 50 25 200 1000 4000 [nt] Overall Results for sample 5: T4-16 RNA Area: 3,805.1 15,430 pg/µl RNA Concentration: rRNA Ratio [28s / 18s]: 1.4

T4-16

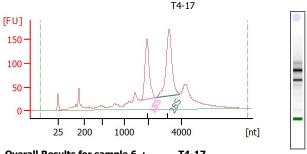
 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

 18S
 1,716
 2,458
 540.6
 14.2

740.0

19.4

4.039



 Overall Results for sample 6 :
 T4-17

 RNA Area:
 2,210.1

 RNA Concentration:
 8,962 pg/µl

 rRNA Ratio [28s / 18s]:
 1.3

RNA Integrity Number (RIN):

Result Flagging Color:

Result Flagging Label:

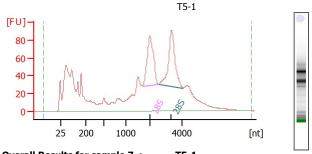
RIN: 8.30

Fragment table for sample 6 : <u>T4-17</u>

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

 18S
 1,665
 2,445
 344.2
 15.6

 28S
 2,471
 3,894
 458.0
 20.7



 Coverall Results for sample 7 :
 T5-1

 RNA Area:
 2,414.4

 RNA Concentration:
 9,790 pg/µl

 rRNA Ratio [28s / 18s]:
 1.1

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

 Result Flagging Color:
 RIN: 6.80

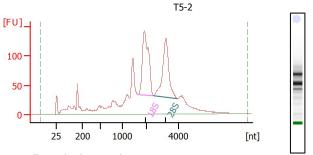
Fragment table for sample 7: <u>T5-1</u>

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

 18S
 1,692
 2,509
 210.2
 8.7

 28S
 2,535
 4,065
 226.5
 9.4

# **Electropherogram Summary Continued ...**



**Overall Results for sample 8 :** T5-2 RNA Area: 2,533.6

RNA Concentration: 10,274 pg/μl rRNA Ratio [28s / 18s]: 0.8

RNA Integrity Number (RIN): 7.4 (B.02.08, Anomaly Threshold(s) manually adapted)
Result Flagging Color:

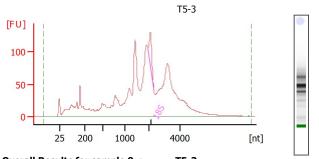
Result Flagging Label: RIN: 7.40

Fragment table for sample 8: T5-2

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

 18S
 1,673
 2,444
 423.1
 16.7

 28S
 2,470
 3,950
 350.4
 13.8



 Overall Results for sample 9 :
 T5-3

 RNA Area:
 2,579.4

 RNA Concentration:
 10,459 pg/µl

 rRNA Ratio [28s / 18s]:
 0.0

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

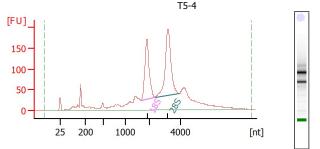
 Result Flagging Color:
 RIN: 2.90

 Result Flagging Label:
 RIN: 2.90

Fragment table for sample 9: <u>T5-3</u>

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

18S 1,988 2,351 75.5 2.9



Overall Results for sample 10 : T5-4

RNA Area: 2,586.7

RNA Concentration: 10,489 pg/µl

rRNA Ratio [28s / 18s]: 1.3

RNA Integrity Number (RIN): 8.3 (B.02.08)

Result Flagging Color:

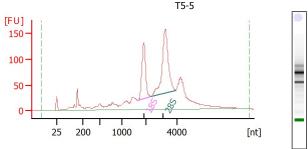
Result Flagging Label: RIN: 8.30

Fragment table for sample 10 : T5-4

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

 18S
 1,660
 2,351
 389.4
 15.1

 28S
 2,377
 3,893
 516.3
 20.0



Overall Results for sample 11: T5-5

RNA Area: 2,129.0

RNA Concentration: 8,633 pg/µl

rRNA Ratio [28s / 18s]: 1.6

RNA Integrity Number (RIN): 8.1 (B.02.08)

Result Flagging Color:

Result Flagging Label: RIN: 8.10

Fragment table for sample 11: <u>T5-5</u>

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

 18S
 1,675
 2,324
 260.9
 12.3

 28S
 2,350
 3,886
 428.8
 20.1

