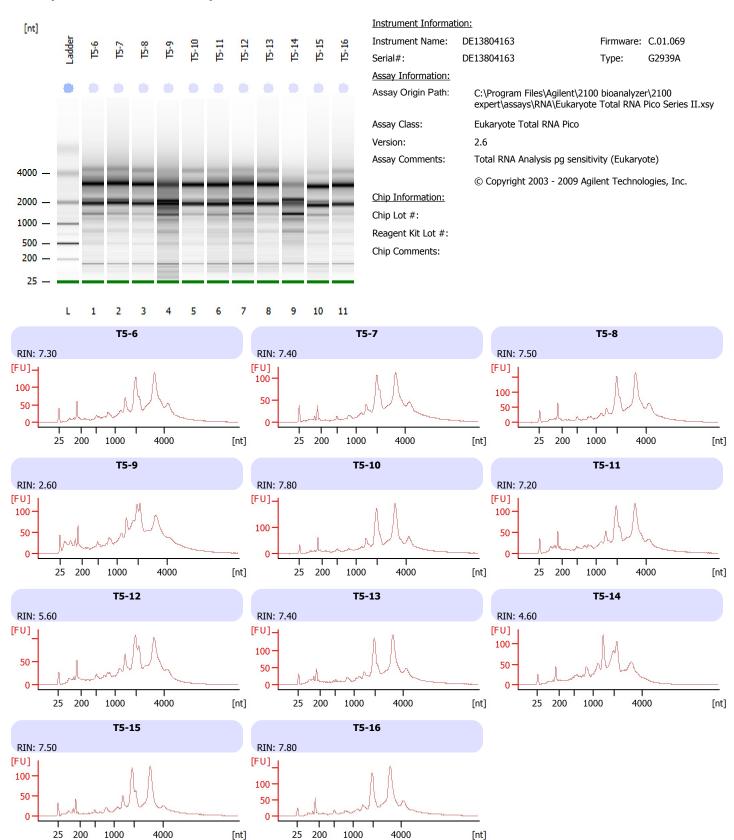
## **Electrophoresis File Run Summary**



Printed:

6/24/2016 12:24:34 PM

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Assay Class: Eukaryote Total RNA Pico C:\...Eukaryote Total RNA Pico\_DE13804163\_2016-06-23\_16-59-36.xad Created: 6/23/2016 4:59:35 PM Modified: 6/23/2016 5:59:56 PM

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

Sample Name	Sample Comment	Status	Result Label	Result Color
T5-6		~	RIN: 7.30	
T5-7		~	RIN: 7.40	
T5-8		~	RIN: 7.50	
T5-9		~	RIN: 2.60	
T5-10		~	RIN: 7.80	
T5-11		<b>~</b>	RIN: 7.20	
T5-12		~	RIN: 5.60	
T5-13		~	RIN: 7.40	
T5-14		<b>~</b>	RIN: 4.60	
T5-15		~	RIN: 7.50	
T5-16		~	RIN: 7.80	
Ladder		~	All Other Samples	

Chip Lot # Reagent Kit Lot #

**Chip Comments:** 

Page 3 of 7

Printed: 6/24/2016 12:24:34 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**

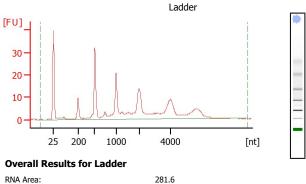
Filter Width [s]: 0.5 Polynomial Order: 4

#### Ladder

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

6/23/2016 4:59:35 PM Assay Class: Eukaryote Total RNA Pico Created: C:\...Éukaryote Total RNA Pico\_DE13804163\_2016-06-23\_16-59-36.xad Data Path: Modified: 6/23/2016 5:59:56 PM

#### **Electropherogram Summary**

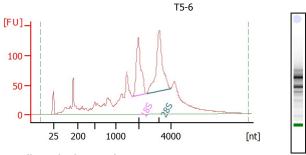


1,000 pg/µl

All Other Samples

RNA Concentration:

Result Flagging Color: Result Flagging Label:



Overall Results for sample 1: <u>T5-6</u>

RNA Area: 2,709.1 RNA Concentration: 9,619 pg/µl rRNA Ratio [28s / 18s]: 1.3

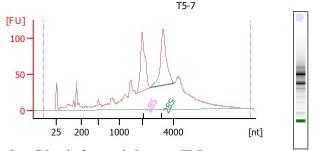
RNA Integrity Number (RIN): 7.3 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color: RIN: 7.30 Result Flagging Label:

Fragment table for sample 1: <u>T5-6</u>

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

18S 1,707 2,453 290.7 10.7 285 2.479 4,003 379.6 14.0



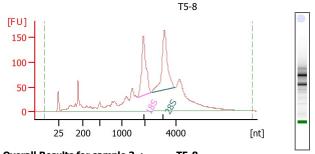
Overall Results for sample 2: <u>T5-7</u> RNA Area: 2,090.2

RNA Concentration: 7,422 pg/µl rRNA Ratio [28s / 18s]: 1.1 RNA Integrity Number (RIN): 7.4 (B.02.08) Result Flagging Color: RIN: 7.40 Result Flagging Label:

Fragment table for sample 2: <u>T5-7</u>

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

18S 1,737 2,480 238.8 11.4 28S 2,506 3,996 266.8 12.8



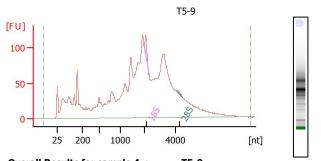
Overall Results for sample 3: T5-8

RNA Area: 2.736.5 RNA Concentration: 9,717 pg/µl rRNA Ratio [28s / 18s]: 1.2 RNA Integrity Number (RIN): 7.5 (B.02.08) Result Flagging Color: Result Flagging Label: RIN: 7.50

Fragment table for sample 3: <u>T5-8</u>

Name Start Size [nt] End Size [nt] Area % of total Area 18S 1,695 2,401 329.8 12.1 28S 2,427 3,911 385.3 14.1

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

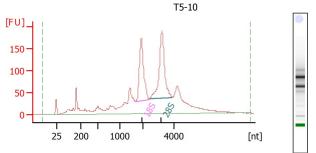


Result Flagging Label: RIN: 2.60

Fragment table for sample 4: T5-9

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

18S 2,019 2,309 33.6 1.0 28S 4,023 4,445 4.8 0.1



Overall Results for sample 5 : T5-10

RNA Area: 2,726.6 RNA Concentration: 9,682 pg/µl rRNA Ratio [28s / 18s]: 1.2

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

T5-10

Result Flagging Color:

Result Flagging Label:

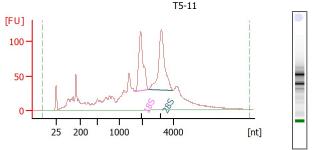
RIN: 7.80

Fragment table for sample 5:

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

 18S
 1,686
 2,414
 385.0
 14.1

 28S
 2,440
 3,871
 471.4
 17.3



Overall Results for sample 6 : T5-11

RNA Area: 2,262.8

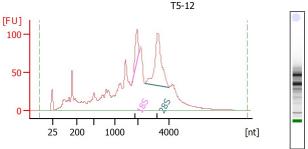
Result Flagging Label: RIN: 7.20

Fragment table for sample 6: T5-11

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

 18S
 1,704
 2,387
 245.5
 10.8

 28S
 2,414
 3,937
 303.4
 13.4



Overall Results for sample 7: T5-12
RNA Area: 2,437.7

RNA Concentration: 8,656 pg/µl
rRNA Ratio [28s / 18s]: 3.0
RNA Integrity Number (RIN): 5.6 (B.02.08)
Result Flagging Color:
Result Flagging Label: RIN: 5.60

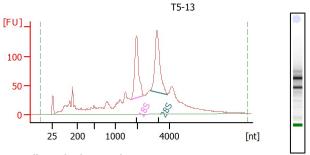
Fragment table for sample 7: <u>T5-12</u>

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

 18S
 1,760
 2,104
 85.9
 3.5

 28S
 2,479
 3,992
 261.3
 10.7

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



 Overall Results for sample 8 : T5-13

 RNA Area:
 2,366.7

 RNA Concentration:
 8,404 pg/µl

 rRNA Ratio [28s / 18s]:
 1.1

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

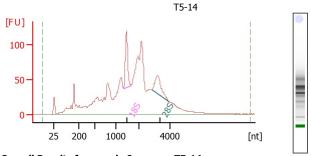
 Result Flagging Color:
 RIN: 7.40

Fragment table for sample 8: <u>T5-13</u>

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

 18S
 1,686
 2,385
 276.9
 11.7

 28S
 2,774
 3,888
 307.2
 13.0



 Coverall Results for sample 9 :
 T5-14

 RNA Area:
 2,333.7

 RNA Concentration:
 8,287 pg/µl

 rRNA Ratio [28s / 18s]:
 0.7

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

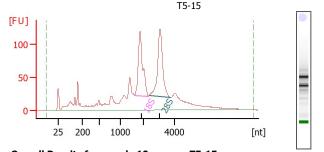
 Result Flagging Color:
 RIN: 4.60

Fragment table for sample 9: <u>T5-14</u>

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

 18S
 1,338
 1,661
 125.7
 5.4

 28S
 2,844
 3,941
 94.2
 4.0



Overall Results for sample 10 : T5-15

RNA Area: 1,937.2

RNA Concentration: 6,879 pg/μl rRNA Ratio [28s / 18s]: 1.1

RNA Integrity Number (RIN): 7.5 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

Result Flagging Label:

RIN: 7.50

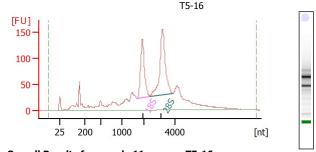
Fragment table for sample 10:

T5-15

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

 18S
 1,627
 2,357
 279.0
 14.4

 28S
 2,384
 3,773
 307.5
 15.9



 Coverall Results for sample 11
 T5-16

 RNA Area:
 2,257.2

 RNA Concentration:
 8,015 pg/µl

 rRNA Ratio [28s / 18s]:
 1.5

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

 Result Flagging Color:
 Essult Flagging Label:

 Result Flagging Label:
 RIN: 7.80

Fragment table for sample 11:  $\underline{\mathsf{T5-16}}$ 

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

 18S
 1,687
 2,342
 279.6
 12.4

 28S
 2,369
 3,835
 405.4
 18.0

