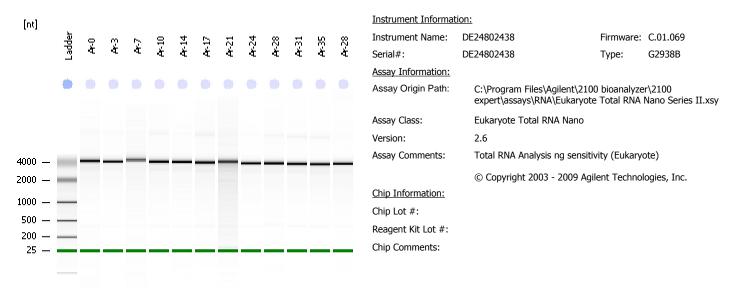
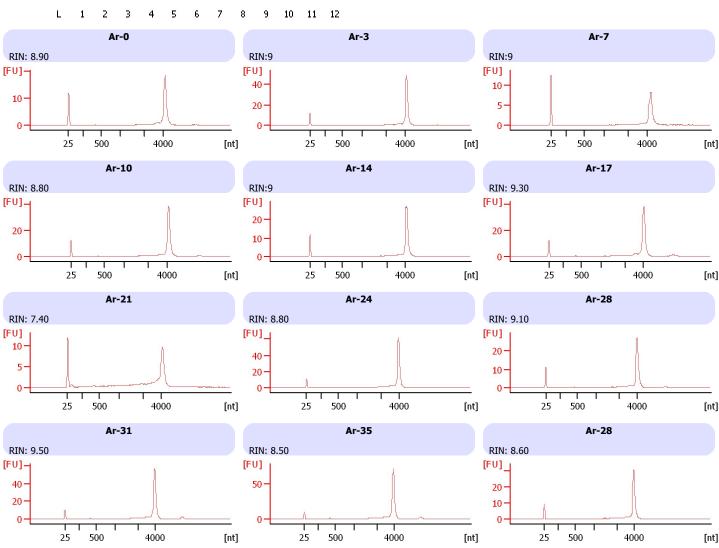
# **Electrophoresis File Run Summary**





Page 2 of 18

Assay Class: Eukaryote Total RNA Nano C:\...040513\_RNA\_Nano\_Tot\_Euk\_Stanton-Geddes-RNA\_Sample\_001\_2.xad C:\...040513\_RNA\_Sample\_001\_2.xad C:\...040513\_RNA\_Sample\_001\_2.xad C:\...040513\_RNA\_Sample\_00

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

Sample Name	Sample Comment	Status	Result Label	Result Color
Ar-0		~	RIN: 8.90	
Ar-3		~	RIN:9	
Ar-7		<b>~</b>	RIN:9	
Ar-10		~	RIN: 8.80	
Ar-14		~	RIN:9	
Ar-17		~	RIN: 9.30	
Ar-21		~	RIN: 7.40	
Ar-24		~	RIN: 8.80	
Ar-28		~	RIN: 9.10	
Ar-31		~	RIN: 9.50	
Ar-35		~	RIN: 8.50	
Ar-28		~	RIN: 8.60	

Chip Lot # Reagent Kit Lot #

**Chip Comments:** 

Page 3 of 18

Assay Class: Eukaryote Total RNA Nano Created: 4/5/2013 4:03:27 PM Data Path: C:\...040513\_RNA\_Nano\_Tot\_Euk\_Stanton-Geddes-RNA\_Sample\_001\_2.xad Modified: 4/5/2013 4:32:39 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

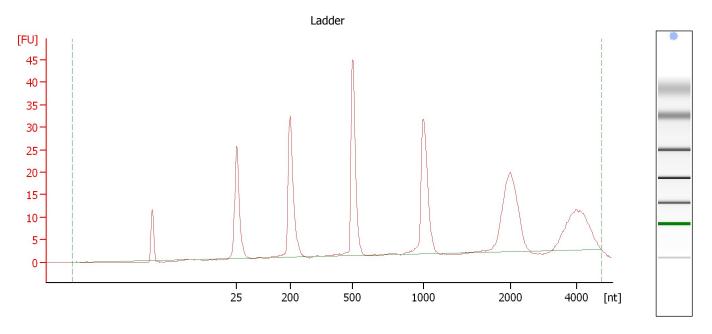
Ladder Peak	Size
1	25
2	200
3	500
4	1000
5	2000
6	4000

Printed:

4/5/2013 4:34:24 PM

Assay Class: Eukaryote Total RNA Nano C:\...040513\_RNA\_Nano\_Tot\_Euk\_Stanton-Geddes-RNA\_Sample\_001\_2.xad C:\...040513\_RNA\_Sample\_001\_2.xad C:\...040513\_RNA\_Sample\_001\_2.xad C:\...040513\_RNA\_Sample\_00

# **Electropherogram Summary**



#### **Overall Results for Ladder**

RNA Area: 227.8 RNA Concentration: 150 ng/µl Result Flagging Color: Result Flagging Label:

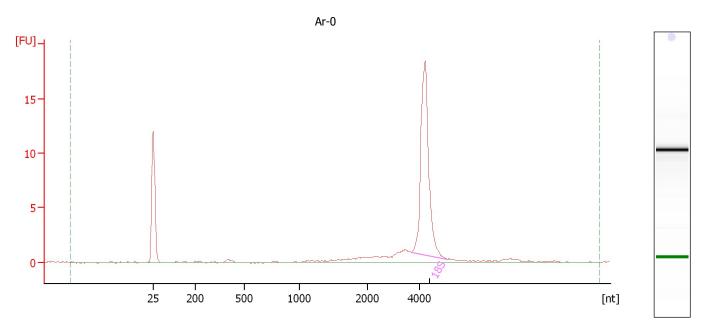
All Other Samples

Assay Class: Eukaryote Total RNA Nano

Data Path: C:\...040513\_RNA\_Nano\_Tot\_Euk\_Stanton-Geddes-RNA\_Sample\_001\_2.xad

Created: Modified: 4/5/2013 4:03:27 PM 4/5/2013 4:32:39 PM

# **Electropherogram Summary Continued ...**



Overall Results for sample 1 : Ar-0

RNA Area: 44.2

rRNA Ratio [28s / 18s]: 0.0

RNA Concentration:

RNA Integrity Number (RIN):

8.9 (B.02.07, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

Result Flagging Label:

RIN: 8.90

Fragment table for sample 1: <u>Ar-0</u>

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

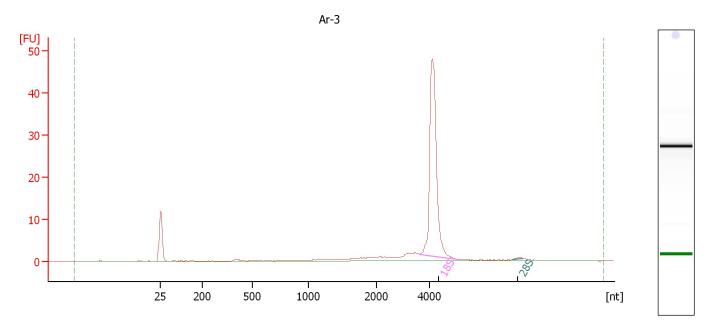
29 ng/µl

18S 3,725 5,087 28.7 65.0

Page 6 of 18

Assay Class: 4/5/2013 4:03:27 PM 4/5/2013 4:32:39 PM Eukaryote Total RNA Nano Created: Data Path: C:\...040513\_RNA\_Nano\_Tot\_Euk\_Stanton-Geddes-RNA\_Sample\_001\_2.xad Modified:

# **Electropherogram Summary Continued ...**



#### Overall Results for sample 2: <u>Ar-3</u>

RNA Area: 97.6

**RNA Concentration:** 64 ng/µl

rRNA Ratio [28s / 18s]: 0.0 RNA Integrity Number (RIN):

9 (B.02.07, Anomaly Threshold(s) manually

adapted)

Result Flagging Color:

Result Flagging Label:

RIN:9

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

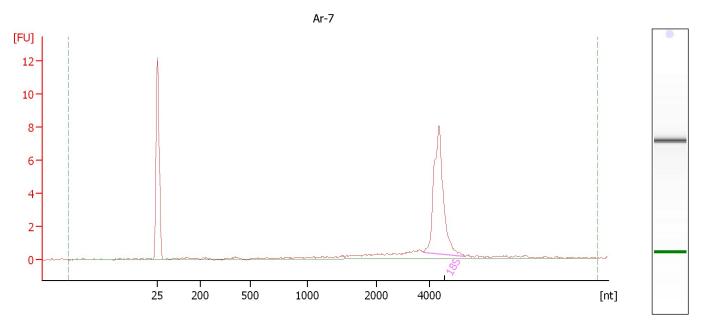
Start Size [nt] End Size [nt] % of total Area Name Area **18S** 66.3

3,637 5,085 64.7 **28S** 7,166 7,575 0.3 0.3 Assay Class: Eukaryote Total RNA Nano

Data Path: C:\...040513\_RNA\_Nano\_Tot\_Euk\_Stanton-Geddes-RNA\_Sample\_001\_2.xad

Created: Modified: 4/5/2013 4:03:27 PM 4/5/2013 4:32:39 PM

# **Electropherogram Summary Continued ...**



Overall Results for sample 3: <u>Ar-7</u>

RNA Area: 25.5 RNA Concentration: 17 ng/µl

rRNA Ratio [28s / 18s]: 0.0 RNA Integrity Number (RIN):

9 (B.02.07, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

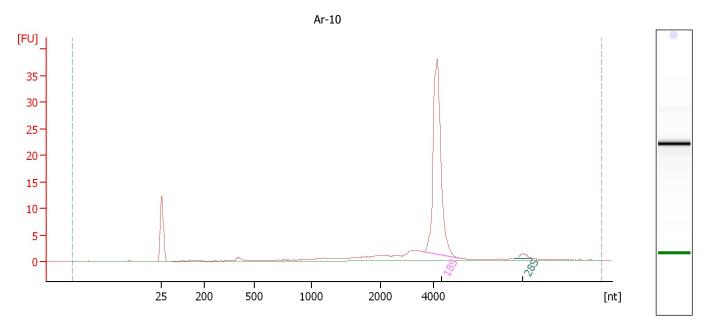
Result Flagging Label: RIN:9

Fragment table for sample 3: <u>Ar-7</u>

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

**18S** 3,772 5,367 14.6 57.1 Assay Class: Eukaryote Total RNA Nano C:\...040513\_RNA\_Nano\_Tot\_Euk\_Stanton-Geddes-RNA\_Sample\_001\_2.xad C:\...040513\_RNA\_Sample\_001\_2.xad C:\...040513\_RNA\_Sample\_001\_2.xad C:\...040513\_RNA\_Sample\_00

# **Electropherogram Summary Continued ...**



# Overall Results for sample 4: <u>Ar-10</u>

RNA Area: 99.2 RNA Concentration: 65 ng/µl

rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN):

8.8 (B.02.07, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

Result Flagging Label:

RIN: 8.80

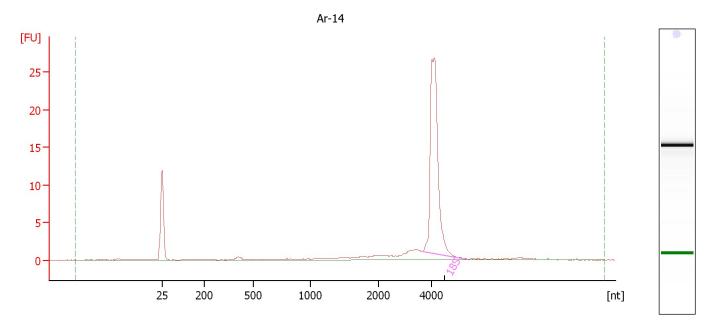
Fragment table for sample 4: <u>Ar-10</u>

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	3,591	5,044	58.8	59.3
285	7 082	7 708	1.1	1 1

Page 9 of 18

Assay Class: 4/5/2013 4:03:27 PM 4/5/2013 4:32:39 PM Eukaryote Total RNA Nano Created: Data Path: C:\...040513\_RNA\_Nano\_Tot\_Euk\_Stanton-Geddes-RNA\_Sample\_001\_2.xad Modified:

**Electropherogram Summary Continued ...** 



Overall Results for sample 5: <u>Ar-14</u>

RNA Area: 63.5

RNA Concentration: 42 ng/µl

rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN):

Result Flagging Label:

Result Flagging Color:

9 (B.02.07, Anomaly Threshold(s) manually

adapted)

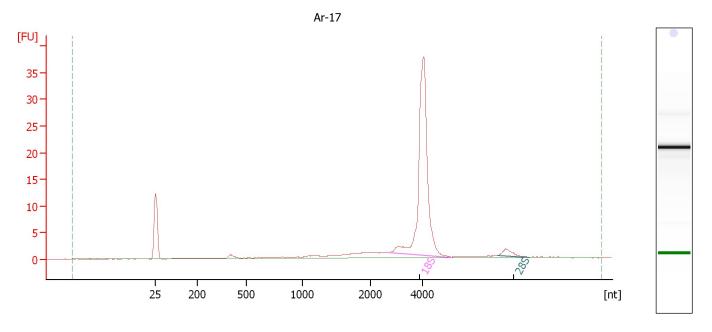
RIN:9

Fragment table for sample 5: Ar-14

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

**18S** 3,660 5,370 41.1 64.8

# **Electropherogram Summary Continued ...**



Overall Results for sample 6: <u>Ar-17</u>

RNA Area: 98.2 65 ng/µl

RNA Concentration: rRNA Ratio [28s / 18s]: 0.0 RNA Integrity Number (RIN):

9.3 (B.02.07, Anomaly Threshold(s) manually adapted)

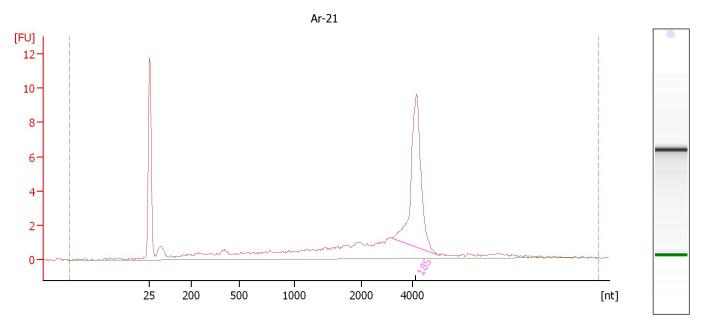
Result Flagging Color: Result Flagging Label:

RIN: 9.30

Fragment table for sample 6: Ar-17

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	2,677	5,148	66.7	68.0
28S	6,919	8,154	2.3	2.3

# **Electropherogram Summary Continued ...**



Overall Results for sample 7: <u>Ar-21</u>

RNA Area: 58.0

RNA Concentration: 38 ng/µl rRNA Ratio [28s / 18s]: 0.0

Result Flagging Color:

Result Flagging Label:

RNA Integrity Number (RIN):

7.4 (B.02.07,

RIN: 7.40

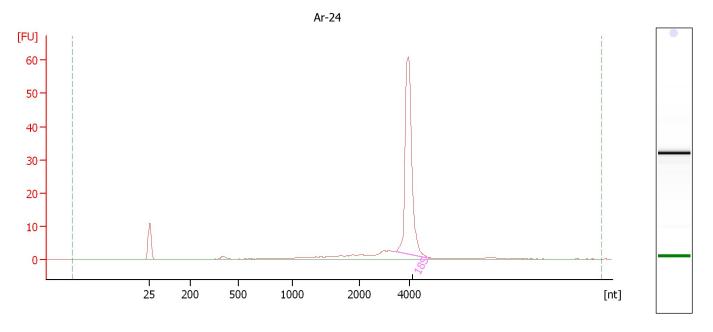
Anomaly Threshold(s) manually adapted)

Fragment table for sample 7: <u>Ar-21</u>

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

**18S** 3,214 5,046 18.4 31.8

# **Electropherogram Summary Continued ...**



Overall Results for sample 8 : <u>Ar-24</u>

RNA Area: 123.8

RNA Concentration: 82 ng/µl

rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN):

8.8 (B.02.07, Anomaly Threshold(s) manually adapted)

manually adapted

Result Flagging Color:

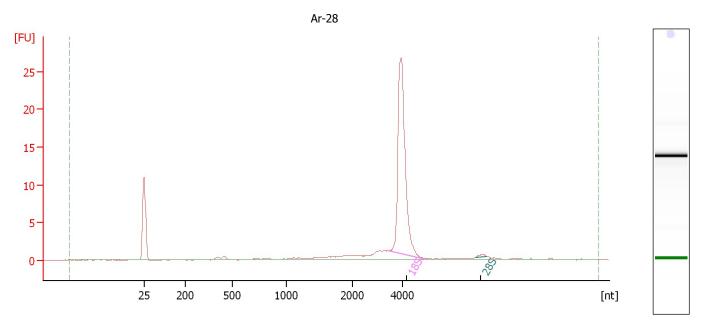
Result Flagging Label: RIN: 8.80

Fragment table for sample 8: <u>Ar-24</u>

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

18S 3,447 4,836 84.1 67.9

# **Electropherogram Summary Continued ...**



Overall Results for sample 9: Ar-28

RNA Area: 61.2 RNA Concentration: 40 ng/µl

rRNA Ratio [28s / 18s]: 0.0

9.1 (B.02.07, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

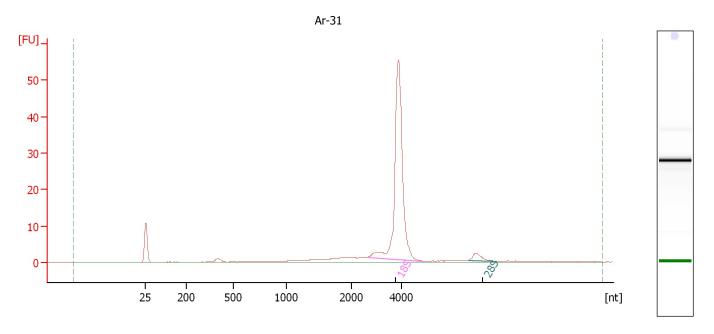
RNA Integrity Number (RIN):

Result Flagging Label: RIN: 9.10

Fragment table for sample 9: <u>Ar-28</u>

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	3,414	4,962	41.1	67.3
28S	6,892	7,338	0.4	0.7

# **Electropherogram Summary Continued ...**



#### Overall Results for sample 10: <u>Ar-31</u>

RNA Area: 132.8 RNA Concentration: 87 ng/µl

rRNA Ratio [28s / 18s]: 0.0 RNA Integrity Number (RIN):

9.5 (B.02.07, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

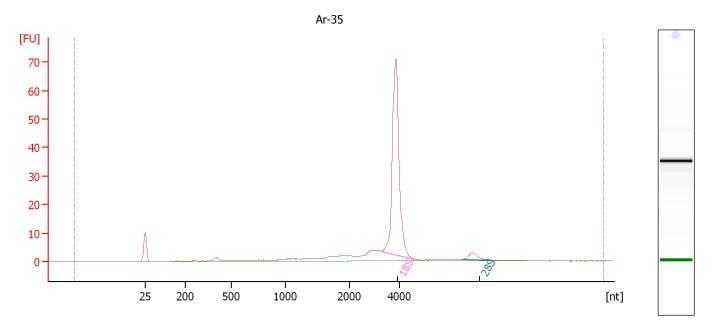
Result Flagging Label:

RIN: 9.50

Fragment table for sample 10: <u>Ar-31</u>

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	2,628	4,898	91.9	69.2
285	6 696	7.810	3 4	2.5

# **Electropherogram Summary Continued ...**



#### Overall Results for sample 11: <u>Ar-35</u>

RNA Area: 171.1

RNA Concentration: 113 ng/µl

rRNA Ratio [28s / 18s]: 0.0 RNA Integrity Number (RIN):

total Area

8.5 (B.02.07, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

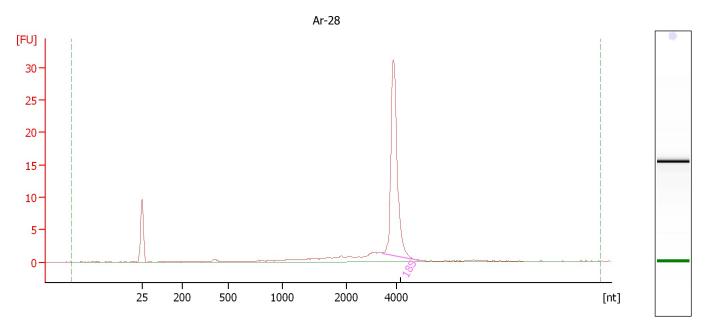
Result Flagging Label:

RIN: 8.50

Fragment table for sample 11: <u>Ar-35</u>

Name	Start Size [nt]	End Size [nt]	Area	% of
18S	3,260	4,617	101.4	59.2
285	6 556	7 934	4 2	2.4

# **Electropherogram Summary Continued ...**



Overall Results for sample 12: Ar-28

RNA Area: 65.5 RNA Concentration: 43 ng/µl

rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN):

8.6 (B.02.07, Anomaly Threshold(s) manually adapted)

RIN: 8.60

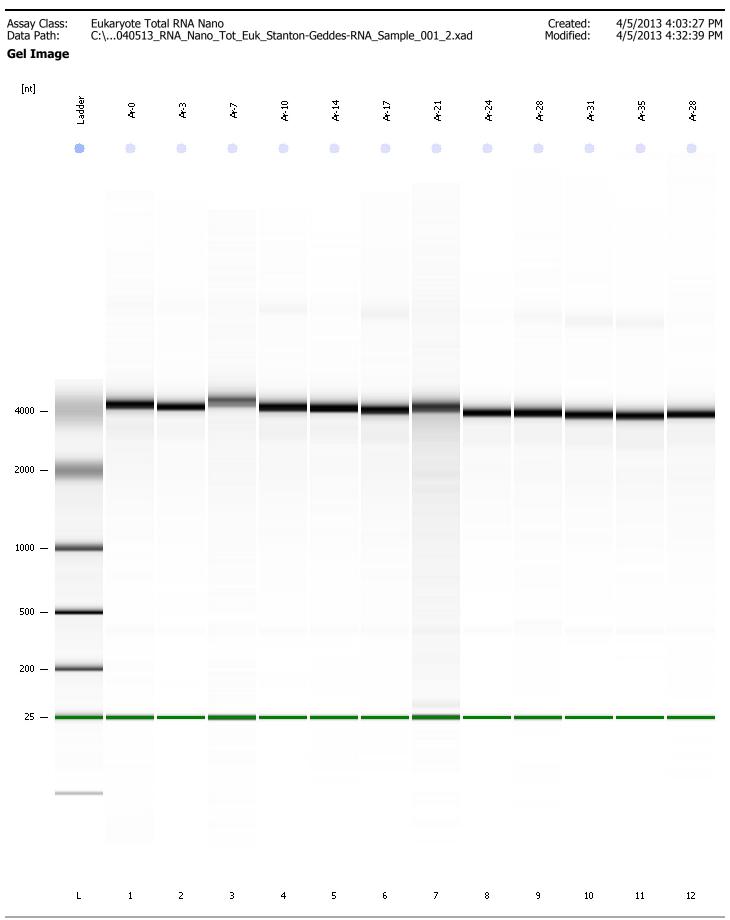
Result Flagging Color:

Result Flagging Label:

Fragment table for sample 12: <u>Ar-28</u>

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

18S 3,433 4,940 42.2 64.5



#### **Run Logbook**

<b>Description</b> Run ended on port 1 (Number of wells acquired: 13)	Number	<b>Source</b> Instrument	<b>Category</b> Run	Sub Category	<b>Time Stamp</b> 4/5/2013 4:26:34 PM	<b>Time Zone</b> (GMT06:00) Central Standard Time	<b>User</b> BMGC	Host BMGC-AAC36D7 D03
Run started on port 1 (File: C:\Program Files\Agilent\21 00 bioanalyzer\210 0 expert\Data\20: 3-04-05\040513 _Eukaryote Total RNA Nano_DE24802 438_2013-04-0! _16-03-27.xad)	1 3	Instrument	Run		4/5/2013 4:03:33 PM	(GMT06:00) Central Standard Time	BMGC	BMGC-AAC36D7 D03
Product Number: G2938B	r	Instrument	Run		4/5/2013 4:03:33 PM	(GMT06:00) Central Standard Time	BMGC	BMGC-AAC36D7 D03
Name :		Instrument	Run		4/5/2013 4:03:33 PM	(GMT06:00) Central Standard Time	BMGC	BMGC-AAC36D7 D03
Vendor : Agilent Technologies	t	Instrument	Run		4/5/2013 4:03:33 PM	(GMT06:00) Central Standard Time	BMGC	BMGC-AAC36D7 D03
Serial# : DE24802438		Instrument	Run		4/5/2013 4:03:33 PM	(GMT06:00) Central Standard Time	BMGC	BMGC-AAC36D7 D03
Firmware : C.01.069		Instrument	Run		4/5/2013 4:03:33 PM	(GMT06:00) Central Standard Time	BMGC	BMGC-AAC36D7 D03
Cartridge : Electrode		Instrument	Run		4/5/2013 4:03:33 PM	(GMT06:00) Central Standard Time	BMGC	BMGC-AAC36D7 D03