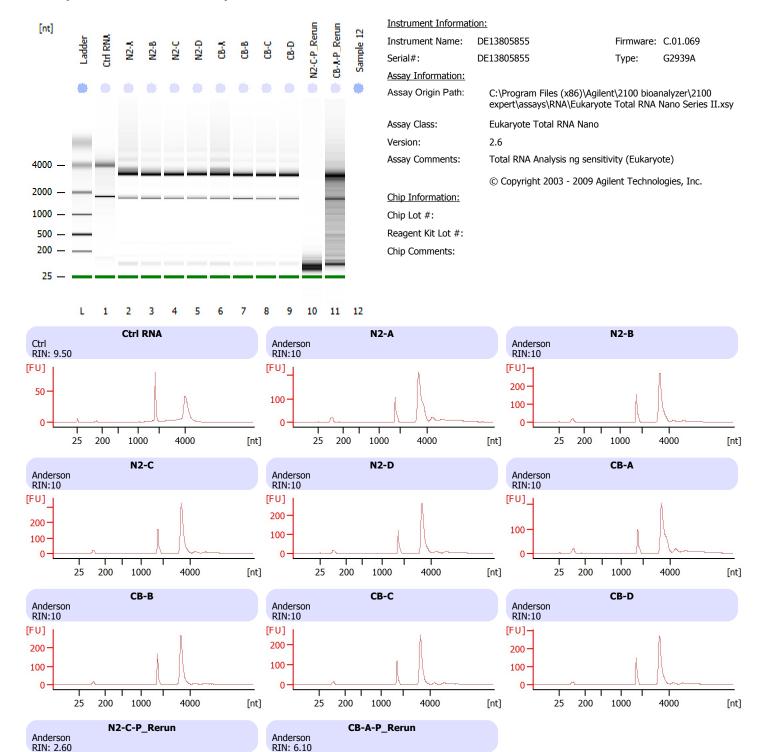
Assay Class: Eukaryote Total RNA Nano C:\...Eukaryote Total RNA Nano_DE13805855_2016-12-27_13-56-10.xad Created: 12/27/2016 1:56:09 PM Modified: 12/27/2016 2:22:28 PM

Electrophoresis File Run Summary



25 200

[FU]

10

1000

4000

[nt]

Printed: 12/27/2016 2:22:58 PM

[FU] 10

5

25 200

[nt]

4000

1000

Page 2 of

Printed: 12/27/2016 2:22:58 PM

Assay Class: Eukaryote Total RNA Nano C:\...Eukaryote Total RNA Nano_DE13805855_2016-12-27_13-56-10.xad Created: 12/27/2016 1:56:09 PM Modified: 12/27/2016 2:22:28 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Stat us	Result Label	Result Color
Ctrl RNA	Ctrl	~	RIN: 9.50	
N2-A	Anderson	~	RIN:10	
N2-B	Anderson	~	RIN:10	
N2-C	Anderson	~	RIN:10	
N2-D	Anderson	~	RIN:10	
CB-A	Anderson	~	RIN:10	
CB-B	Anderson	~	RIN:10	
CB-C	Anderson	~	RIN:10	
CB-D	Anderson	~	RIN:10	
N2-C-P_Rerun	Anderson	~	RIN: 2.60	
CB-A-P_Rerun	Anderson	~	RIN: 6.10	
Sample 12				
Ladder		~	All Other Samples	

Chip Lot # Reagent Kit Lot #

Chip Comments:

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Printed: 12/27/2016 2:22:58 PM

Assay Class: Eukaryote Total RNA Nano C:\...Eukaryote Total RNA Nano_DE13805855_2016-12-27_13-56-10.xad Created: 12/27/2016 1:56:09 PM Modified: 12/27/2016 2:22:28 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

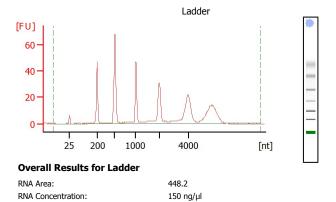
Page 4 of 6

Assay Class: Eukaryote Total RNA Nano Created: 12/27/2016 1:56:09 PM C:\...Éukaryote Total RNA Nano_DE13805855_2016-12-27_13-56-10.xad Data Path: Modified: 12/27/2016 2:22:28 PM

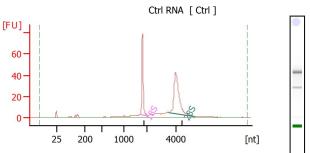
Electropherogram Summary

Result Flagging Color:

Result Flagging Label:



All Other Samples



Ctrl RNA Overall Results for sample 1:

RNA Area: 281.0 94 ng/µl RNA Concentration: rRNA Ratio [28s / 18s]: 1.6

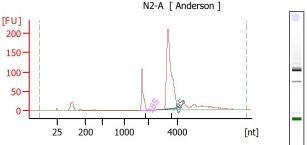
9.5 (B.02.08, Anomaly Threshold(s) manually adapted) RNA Integrity Number (RIN):

Result Flagging Color: RIN: 9.50 Result Flagging Label:

Fragment table for sample 1: Ctrl RNA

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

18S 23.1 1,733 2,032 65.0 28S 3,530 5,272 101.0 36.0





RNA Area: 944.0 RNA Concentration: 316 ng/µl rRNA Ratio [28s / 18s]: 4.3

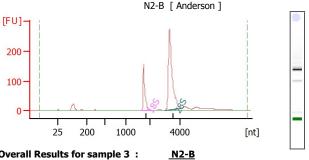
RNA Integrity Number (RIN): 10 (B.02.08, Anomaly Threshold(s) manually adapted) Result Flagging Color:

RIN:10 Result Flagging Label:

Fragment table for sample 2: N2-A

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
100	1 623	2 202	125.5	12.2

28S 3,036 4,135 533.9 56.6



Overall Results for sample 3:

RNA Area: 950.4 RNA Concentration: 318 ng/µl rRNA Ratio [28s / 18s]: 3.3

RNA Integrity Number (RIN): 10 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color: Result Flagging Label: RIN:10

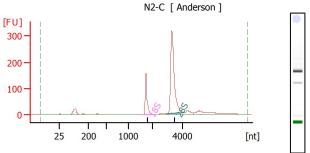
Fragment table for sample 3: <u>N2-B</u>

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	1,620	2,140	170.2	17.9
28S	3,018	4,108	556.2	58.5

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Assay Class: Eukaryote Total RNA Nano Created: 12/27/2016 1:56:09 PM C:\...Éukaryote Total RNA Nano_DE13805855_2016-12-27_13-56-10.xad Data Path: Modified: 12/27/2016 2:22:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 4: <u>N2-C</u>

RNA Area: 1,017.8 341 ng/µl RNA Concentration: rRNA Ratio [28s / 18s]:

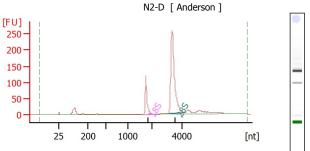
RNA Integrity Number (RIN): 10 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color: Result Flagging Label: RIN:10

Fragment table for sample 4: N2-C

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

18S 1,600 2,115 164.5 16.2 28S 2.944 4.095 603.0 59.3



Overall Results for sample 5: <u>N2-D</u>

RNA Area: 834.8 RNA Concentration: 279 ng/µl rRNA Ratio [28s / 18s]:

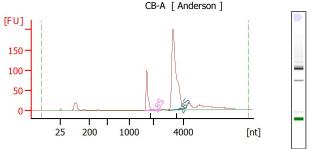
RNA Integrity Number (RIN): 10 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color: Result Flagging Label:

RIN:10 Fragment table for sample 5: <u>N2-D</u>

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

18S 1,602 2,102 124.9 15.0 285 2.999 4.106 502.8 60.2



Overall Results for sample 6: CB-A

825.1 RNA Area: RNA Concentration: 276 ng/µl rRNA Ratio [28s / 18s]: 4.4

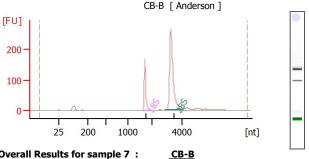
RNA Integrity Number (RIN): 10 (B.02.08, Anomaly Threshold(s) manually adapted) Result Flagging Color:

RIN:10 Result Flagging Label:

Fragment table for sample 6: CB-A

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

18S 1,635 2,270 108.2 13.1 28S 3,000 4,095 474.0 57.4



Overall Results for sample 7:

RNA Area: 812.4 RNA Concentration: 272 ng/µl rRNA Ratio [28s / 18s]: 3.2

RNA Integrity Number (RIN): 10 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color: RIN:10 Result Flagging Label:

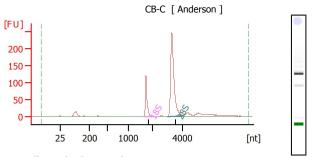
Fragment table for sample 7: CB-B

Name Start Size [nt] End Size [nt] Area % of total Area 18S 1,565 2,003 149.3 18.4 28S 2,916 4,095 480.1 59.1

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Assay Class: Eukaryote Total RNA Nano C:\...Eukaryote Total RNA Nano_DE13805855_2016-12-27_13-56-10.xad Created: 12/27/2016 1:56:09 PM Modified: 12/27/2016 2:22:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 8: <u>CB-C</u>
RNA Area: 716.0

RNA Concentration: 240 ng/µl rRNA Ratio [28s / 18s]: 3.7

RNA Integrity Number (RIN): 10 (B.02.08, Anomaly Threshold(s) manually adapted)

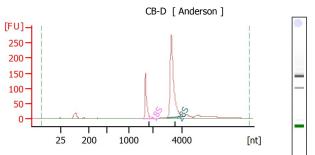
Result Flagging Color:
Result Flagging Label:
RIN:10

Fragment table for sample 8 : <u>CB-C</u>

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

 18S
 1,590
 2,072
 116.0
 16.2

 28S
 2,959
 4,042
 433.8
 60.6



Overall Results for sample 9: <u>CB-D</u>

RNA Area: 894.6 RNA Concentration: 299 ng/μl rRNA Ratio [28s / 18s]: 3.4

RNA Integrity Number (RIN): 10 (B.02.08, Anomaly Threshold(s) manually adapted)

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Result Flagging Color: Result Flagging Label:

Result Flagging Label: RIN:10

Fragment table for sample 9: CB-D

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

 18S
 1,595
 2,155
 153.5
 17.2

 28S
 2,960
 4,062
 526.8
 58.9