5/31/2019 11:42:41 AM

5/31/2019 12:23:47 PM

Created:

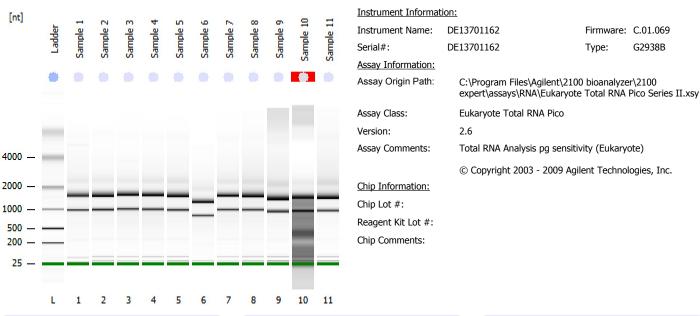
Modified:

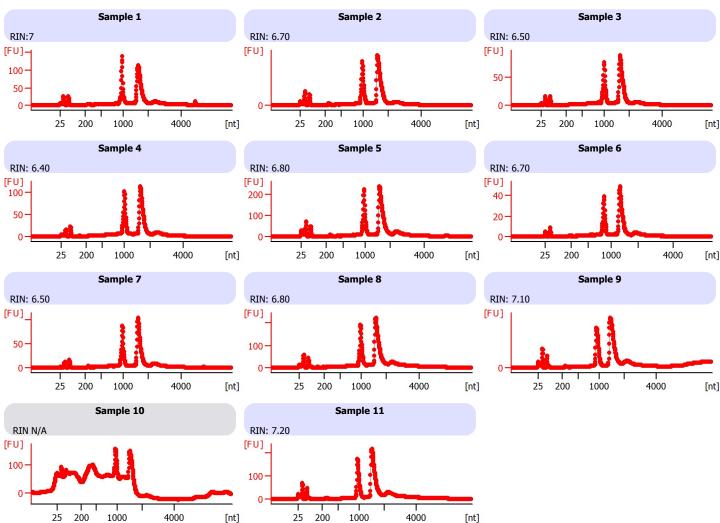
Printed:

5/31/2019 12:24:08 PM

Assay Class: Eukaryote Total RNA Pico
Data Path: Eukaryote Total RNA Pico
C:\...uri\Desktop\Brent\2019-05-31_ribopure half samples 6-12.xad

Electrophoresis File Run Summary





Assay Class: Eukaryote Total RNA Pico C:\...uri\Desktop\Brent\2019-05-31_ribopure half samples 6-12.xad Created: 5/31/2019 11:42:41 AM Modified: 5/31/2019 12:23:47 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Statı s	u Result Label	Result Color
Sample 1		~	RIN:7	
Sample 2		~	RIN: 6.70	
Sample 3		~	RIN: 6.50	
Sample 4		~	RIN: 6.40	
Sample 5		~	RIN: 6.80	
Sample 6		~	RIN: 6.70	
Sample 7		~	RIN: 6.50	
Sample 8		~	RIN: 6.80	
Sample 9		~	RIN: 7.10	
Sample 10		~	RIN N/A	
Sample 11		~	RIN: 7.20	
Ladder		~	All Other Samples	

Chip Lot # Reagent Kit Lot #

Chip Comments:

8

Assay Class: Eukaryote Total RNA Pico C:\...uri\Desktop\Brent\2019-05-31_ribopure half samples 6-12.xad Cc:\...uri\Desktop\Brent\2019-05-31_ribopure half samples 6-12.xad

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

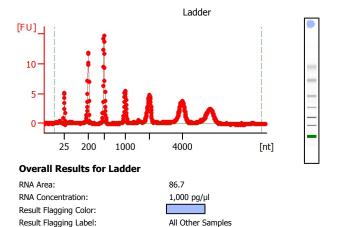
Ladder

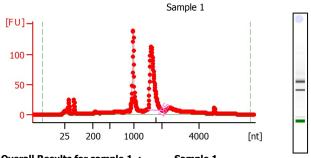
Ladder Peak	Size
1	25
2	200
3	500
4	1000
5	2000
6	4000

8

Assay Class: Eukaryote Total RNA Pico C:\...uri\Desktop\Brent\2019-05-31_ribopure half samples 6-12.xad Cc:\...uri\Desktop\Brent\2019-05-31_ribopure half samples 6-12.xad

Electropherogram Summary





 Overall Results for sample 1
 Sample 1

 RNA Area:
 864.1

 RNA Concentration:
 9,972 pg/µl

 rRNA Ratio [28s / 18s]:
 0.0

RNA Integrity Number (RIN): 7 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

Result Flagging Label: RIN:7

Fragment table for sample 1 : Sample 1

Sample 3

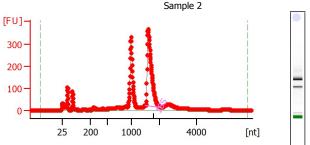
4000

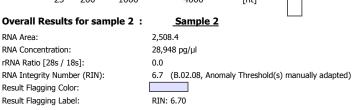
Sample 3

[nt]

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

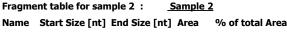
18S 1,489 2,127 276.8 32.0



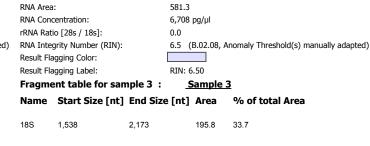


864.4

34.5



2,127



1000

18S

1,504

[FU] 80

60

40

20

25

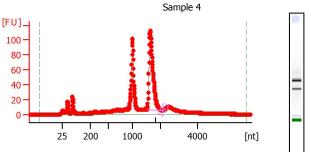
Overall Results for sample 3:

200

Page 5 of 8

Assay Class: Eukaryote Total RNA Pico C:\...uri\Desktop\Brent\2019-05-31_ribopure half samples 6-12.xad Cc:\...uri\Desktop\Brent\2019-05-31_ribopure half samples 6-12.xad

Electropherogram Summary Continued ...



Overall Results for sample 4: Sample 4

RNA Concentration: 9,250 pg/µl rRNA Ratio [28s / 18s]: 0.0

RNA Area:

RNA Integrity Number (RIN): 6.4 (B.02.08, Anomaly Threshold(s) manually adapted)

Sample 4

801.6

Result Flagging Color:

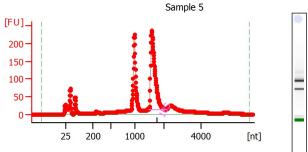
Result Flagging Label:

RIN: 6.40

Fragment table for sample 4:

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

18S 1,519 2,152 252.1 31.4



Overall Results for sample 5 : Sample 5

RNA Area: 1,663.4
RNA Concentration: 19,196 pg/µl
rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN): 6.8 (B.02.08, Anomaly Threshold(s) manually adapted)

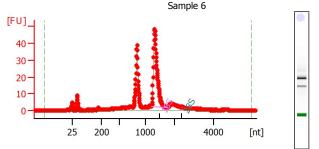
Result Flagging Color:

Result Flagging Label: RIN: 6.80

Fragment table for sample 5 : <u>Sample 5</u>

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

18S 1,490 2,119 572.3 34.4



Overall Results for sample 6 : Sample 6

RNA Area: 278.9
RNA Concentration: 3,218 pg/µl
rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN): 6.7 (B.02.08, Anomaly Threshold(s) manually adapted)
Result Flagging Color:

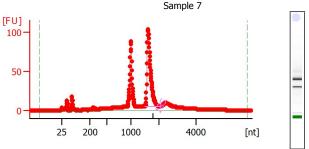
Result Flagging Label: RIN: 6.70

Fragment table for sample 6: Sample 6

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

 18S
 1,207
 1,750
 101.5
 36.4

 28S
 2,335
 2,432
 0.1
 0.0



Overall Results for sample 7 : Sample 7

RNA Area: 631.2
RNA Concentration: 7,284 pg/µl
rRNA Ratio [28s / 18s]: 0.0
RNA Integrity Number (RIN): 6.5 (B.02.1

RNA Integrity Number (RIN): 6.5 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

Printed: 5/31/2019 12:24:08 PM

Result Flagging Label: RIN: 6.50

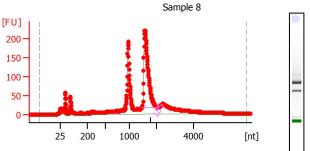
Fragment table for sample 7 : Sample 7

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

 18S
 1,504
 2,092
 210.1
 33.3

Assay Class: Eukaryote Total RNA Pico Created: 5/31/2019 11:42:41 AM C:\...uri\Desktop\Brent\2019-05-31_ribopure half samples 6-12.xad 5/31/2019 12:23:47 PM Data Path: Modified:

Electropherogram Summary Continued ...



Sample 8 Overall Results for sample 8:

RNA Area: 1.738.7 20,065 pg/µl RNA Concentration: rRNA Ratio [28s / 18s]:

RNA Integrity Number (RIN): 6.8 (B.02.08, Anomaly Threshold(s) manually adapted) Result Flagging Color:

534.0

30.7

Result Flagging Label: RIN: 6.80

18S

1,488

Fragment table for sample 8: Sample 8 Start Size [nt] End Size [nt] Area % of total Area Name

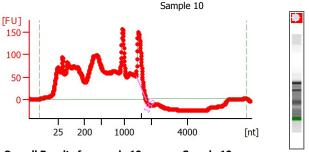
2,112

Sample 9 [FU] 400 300 200 100 25 200 1000 4000 [nt]

Overall Results for sample 9: Sample 9 RNA Area: 2,327.9 26,864 pg/µl RNA Concentration: rRNA Ratio [28s / 18s]: 0.0 RNA Integrity Number (RIN): 7.1 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color: RIN: 7.10 Result Flagging Label:

Fragment table for sample 9: Sample 9 Name Start Size [nt] End Size [nt] Area % of total Area 1,173.8 1,309 1,902 50.4



Overall Results for sample 10: Sample 10 RNA Area: 4,120.5

RNA Concentration: 47,551 pg/µl rRNA Ratio [28s / 18s]: 0.0

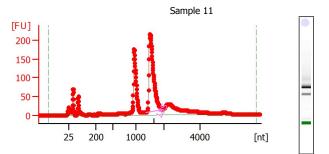
RNA Integrity Number (RIN): N/A (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color: RIN N/A Result Flagging Label:

Fragment table for sample 10: Sample 10

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

18S 1,422 1,804 273.8 6.6



Overall Results for sample 11: Sample 11 RNA Area: 1,637.2 RNA Concentration: 18,893 pg/µl 0.0

rRNA Ratio [28s / 18s]: RNA Integrity Number (RIN): 7.2 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color: RIN: 7.20 Result Flagging Label:

Fragment table for sample 11: Sample 11

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

18S 1,363 1,972 566.2 34.6

Eukaryote Total RNA Pico C:\...uri\Desktop\Brent\2019-05-31_ribopure half samples 6-12.xad Created: Modified: 5/31/2019 11:42:41 AM 5/31/2019 12:23:47 PM Assay Class: Data Path: **Gel Image** [nt] Sample 10 Sample 11 Sample 9 ٠ ٠ • ۰ ۰ ۰ ۰ ٠ ٠ 4000 2000 -1000 -500 200 25 — 2 1 3 5 6 9 10 11

Assay Class: Eukaryote Total RNA Pico C:\...uri\Desktop\Brent\2019-05-31_ribopure half samples 6-12.xad Created: 5/31/2019 11:42:41 AM Modified: 5/31/2019 12:23:47 PM

Run Logbook

Description Run ended on port 1 (Number of wells acquired: 12)	Number	Source Instrument	Category Run	Sub Category	Time 5/31/2019 12:05:32 PM	Time Zone (GMT05:00) Central Standard Time	User BD Accuri	Host BDAccuri-PC
Run started on port 1 (File: C:\Program Files\Agilent\21 00 bioanalyzer\210 0 expert\Data\20: 9-05-31\2100 expert_Eukaryo e Total RNA Pico_DE137011 62_2019-05-31_11-42-41.xad)	1 t	Instrument	Run		5/31/2019 11:42:46 AM	(GMT05:00) Central Standard Time	BD Accuri	BDAccuri-PC
Product Number: G2938B	r	Instrument	Run		5/31/2019 11:42:46 AM	(GMT05:00) Central Standard Time	BD Accuri	BDAccuri-PC
Name :		Instrument	Run		5/31/2019 11:42:46 AM	(GMT05:00) Central Standard Time	BD Accuri	BDAccuri-PC
Vendor : Agilent Technologies	t	Instrument	Run		5/31/2019 11:42:46 AM	(GMT05:00) Central Standard Time	BD Accuri	BDAccuri-PC
Serial# : DE13701162		Instrument	Run		5/31/2019 11:42:46 AM	(GMT05:00) Central Standard Time	BD Accuri	BDAccuri-PC
Firmware : C.01.069		Instrument	Run		5/31/2019 11:42:46 AM	(GMT05:00) Central Standard Time	BD Accuri	BDAccuri-PC
Cartridge : Electrode		Instrument	Run		5/31/2019 11:42:46 AM	(GMT05:00) Central Standard Time	BD Accuri	BDAccuri-PC