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Printed: 3/8/2012 12:37:13 PM

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Assay Class: Data Path:	Prokaryote Total RNA Nano C:\tings\Bioanalyzer\2012-03-08\2012-03-08_005_ErinLynchHS.xad	Created: Modified:	3/8/2012 12:00:13 PM 3/8/2012 12:37:02 PM		
Electrophoresis File Run Summary (Chip Summary)					

Sample Name	Sample Comment Sta	us Result Labe	I Result Color
A0 undepleted	~		
A1 undepleted	~		
A2 undepleted	~		
A3 undepleted	~		
A4 undepleted	~		
B0 undepleted	~		
B1 undepleted	~		
B2 undepleted	~		
B3 undepleted	~		
B4 undepleted	~		
B0 depleted	~		
B1 depleted	~		
Ladder	~		
Chip Lot #		Reage	nt Kit Lot #

**Chip Comments :** 

#### **Electrophoresis Assay Details**

#### **General Analysis Settings**

Number of Available Sample and Ladder Wells (Max.): 13

Minimum Visible Range [s] : 17Maximum Visible Range [s] : 70Start Analysis Time Range [s] : 19End Analysis Time Range [s] : 69Ladder Concentration  $[ng/\mu I] : 150$ Lower Marker Concentration  $[ng/\mu I] : 0$ Upper Marker Concentration  $[ng/\mu I] : 0$ 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

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2012-03-08_005_ErinLynchHS.xad				
Assay Class: Data Path: Electropher	Prokaryote Total RNA Nano C:\tings\Bioanalyzer\2012-03-08\2012-03-08_005_ErinLynchHS.xad ogram Summary	Created: Modified:	3/8/2012 12:00:13 PM 3/8/2012 12:37:02 PM	
	Ladder			
[FU] _	1			
20 -				
15-				
10-				
5-		$\wedge$		



## **Overall Results for Ladder**

RNA Area: 188.2 RNA Concentration: 150 ng/µl



#### Overall Results for sample 1 : A0 undepleted

RNA Area:	1,146.8	RNA Integrity Number (RIN):	10 (B.02.08,
RNA Concentration:	914 ng/µl		Anomaly Threshold(s) manually adapted)
rRNA Ratio [23s / 16s]:	2.0		

Fragment table for sample 1 : <u>A0 undepleted</u>					
Name	Start Size [nt]	End Size [nt]	Area	% of total Area	
16S	1,380	1,545	297.0	25.9	
23S	2,468	2,957	605.5	52.8	

2012-03-08_	2012-03-08_005_ErinLynchHS.xad			
Assay Class: Data Path:	Prokaryote Total RNA Nano C:\tings\Bioanalyzer\2012-03-08\2012-03-08_005_	ErinLynchHS.xad	Created: Modified:	3/8/2012 12:00:13 PM 3/8/2012 12:37:02 PM
Electrophere	ogram Summary Continued			
	A1 undepleted			
[FU]				
450 -		l		
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530

4000

[nt]

2000

# Overall Results for sample 2 : <u>A1 undepleted</u>

1 25

300 -250 -200 -150 -100 -50 -0 -

RNA Area:	767.3	RNA Integrity Number (RIN):	10 (B.02.08,
RNA Concentration:	612 ng/µl		Anomaly Threshold(s) manually adapted)
rRNA Ratio [23s / 16s]:	2.1		

1000

Fragment table for sample 2 : <u>A1 undepleted</u>					
Name	Start Size [nt]	End Size [nt]	Area	% of total Area	
16S	1,394	1,558	196.8	25.6	
23S	2,500	3,121	413.8	53.9	

200

500



# Overall Results for sample 3 : <u>A2 undepleted</u>

RNA Area:	977.3	RNA Integrity Number (RIN):	10	(B.02.08,
RNA Concentration:	779 ng/µl		Anor	naly Threshold(s) ually adapted)
rRNA Ratio [23s / 16s]:	1.9			

Fragment table for sample 3 : <u>A2 undepleted</u>					
Name	Start Size [nt]	End Size [nt]	Area	% of total Area	
16S	1,390	1,553	254.3	26.0	
23S	2,488	2,949	488.8	50.0	



## Overall Results for sample 4 : <u>A3 undepleted</u>

RNA Area:	1,030.5	RNA Integrity Number (RIN):	10 (B.02.08,
RNA Concentration:	821 ng/µl		Anomaly Threshold(s) manually adapted)
rRNA Ratio [23s / 16s]:	2.0		

Fragment table for sample 4 : <u>A3 undep</u>				
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
16S	1,379	1,535	267.0	25.9
23S	2,455	2,884	523.3	50.8

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Assay Data	ay Class: Prokaryote Total RNA Nano Created: a Path: C:\tings\Bioanalyzer\2012-03-08\2012-03-08_005_ErinLynchHS.xad Modified:		3/8/2012 12:0 3/8/2012 12:3	0:13 PM 7:02 PM						
Elec	tropher	ogram Summa	ary Contin	ued						
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		25	200	500	í 1000	2000	4000	[ni		

#### A4 undepleted Overall Results for sample 5 :

RNA Area:	1,084.5	RNA Integrity Number (RIN):	4.9	(B.02.08,
RNA Concentration:	864 ng/µl		Anor man	maly Threshold(s) ually adapted)
rRNA Ratio [23s / 16s]:	0.3			

Fragment	leted			
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
16S	1,302	1,530	77.3	7.1
23S	2,389	2,764	25.8	2.4

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Assay Class: Data Path:	Assay Class: Prokaryote Total RNA Nano Created: 3/   Data Path: C:\tings\Bioanalyzer\2012-03-08\2012-03-08_005_ErinLynchHS.xad Modified: 3/   Electropherogram Summary Continued Modified: 3/				
Electropher					
	B0 undepleted				
[FU]					
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120-					
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0					

2000

4000

[nt]

# Overall Results for sample 6 : <u>B0 undepleted</u>

1 25

RNA Area:	1,361.3	RNA Integrity Number (RIN):	5.6 (B.02.08,	6 (B.02.08,
RNA Concentration:	1,085 ng/µl		Anomaly Threshold(s manually adapted)	5)
rRNA Ratio [23s / 16s]:	0.6			

1000

Fragmen	t table for sample			
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
16S	1,295	1,517	115.1	8.5
23S	2,356	2,757	74.3	5.5

1 200 500



# Overall Results for sample 7 : <u>B1 undepleted</u>

RNA Area:	1,755.9	RNA Integrity Number (RIN):	10 (B.02.08,
RNA Concentration:	1,399 ng/µl		Anomaly Threshold(s) manually adapted)
rRNA Ratio [23s / 16s]:	1.8		

Fragme	nt table for sampl			
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
16S	1,336	1,503	479.5	27.3
23S	2,380	2,808	862.2	49.1

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Assay Class: Data Path:	Prokaryote Total RNA Nano C:\tings\Bioanalyzer\2012-03-08\2012-03-08_005_ErinLynchHS.xad	3/8/2012 12:00:13 PM 3/8/2012 12:37:02 PM	
Electropher	ogram Summary Continued		
	B2 undepleted		
[FU] 500 -			
0			

2000

4000

[nt]

# Overall Results for sample 8 : <u>B2 undepleted</u>

200

500

1 25

RNA Area:	1,820.1	RNA Integrity Number (RIN):	10 (B.02.08,
RNA Concentration:	1,451 ng/µl		Anomaly Threshold(s) manually adapted)
rRNA Ratio [23s / 16s]:	1.8		

1000

Fragmer	nt table for sample			
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
16S	1,321	1,490	504.2	27.7
23S	2,346	2,766	883.9	48.6

Created: 3/8/2012 12:00:13 PM Modified: 3/8/2012 12:37:02 PM

# Electropherogram Summary Continued ...



## Overall Results for sample 9 : <u>B3 undepleted</u>

RNA Area:	1,146.3	RNA Integrity Number (RIN):	10 (B.02.08,
RNA Concentration:	914 ng/µl		Anomaly Threshold(s) manually adapted)
rRNA Ratio [23s / 16s]:	1.7		

Fragme	nt table for sampl	e9: <u>B3 und</u>	<u>epleted</u>	
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
16S	1,282	1,485	304.5	26.6
23S	2,335	2,747	513.7	44.8

Created: 3/8/2012 12:00:13 PM Modified: 3/8/2012 12:37:02 PM

# Electropherogram Summary Continued ...



## Overall Results for sample 10 : <u>B4 undepleted</u>

RNA Area:	1,135.5	RNA Integrity Number (RIN):	9.9 (B.02.08,
RNA Concentration:	905 ng/µl		Anomaly I hreshold(s) manually adapted)
rRNA Ratio [23s / 16s]:	1.5		······································

441.5

38.9

Fragment table for sample 10 : <u>B4 undepleted</u>					
Name	Start Size [nt]	End Size [nt]	Area	% of total Area	
16S	1,260	1,464	288.4	25.4	

2,727

23S

2,265

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# Electropherogram Summary Continued ...



## Overall Results for sample 11 : <u>B0 depleted</u>

RNA Area:	308.4	RNA Integrity Number (RIN):	8.7 (	B.02.08,
RNA Concentration:	246 ng/µl		Anom	aly Threshold(s) ally adapted)
rRNA Ratio [23s / 16s]:	3.0			

Fragme	nt table for sample	e11 :	B0 dep	leted	
Name	Start Size [nt]	End Siz	e [nt]	Area	%

1,493

2,730

	-	
16S		1,279
23S		2,301

Area 42.5 129.1

% of total Area 13.8 41.9

Created: 3/8/2012 12:00:13 PM Modified: 3/8/2012 12:37:02 PM

# Electropherogram Summary Continued ...



## Overall Results for sample 12 : <u>B1 depleted</u>

RNA Area:	280.8	RNA Integrity Number (RIN):	8.3 (B.02.08,
RNA Concentration:	224 ng/µl		Anomaly Threshold(s) manually adapted)
rRNA Ratio [23s / 16s]:	5.9		······································

## Fragment table for sample 12 : <u>B1 depleted</u>

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
16S	1,322	1,486	30.5	10.8
23S	2,325	2,741	179.1	63.8

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[nt]	Ladder	A0 undepleted	Ai undepleted	A2 undepleted	A3 undepleted	A4 undepleted	B0 undepleted	B1 undepleted	B2 undepleted	B3 undepleted	B4 undepleted	B0 depleted	B1 depleted
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### Assay Class: Data Path: Prokaryote Total RNA Nano C:\...tings\Bioanalyzer\2012-03-08\2012-03-08\_005\_ErinLynchHS.xad

Created: Modified:

3/8/2012 12:00:13 PM 3/8/2012 12:37:02 PM

#### **Run Logbook**

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 13)		Instrument	Run		3/8/2012 12:24:01 PM	(GMT08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanal yzer\2012-03-08 \2012-03-08_00 5.xad)	3	Instrument	Run		3/8/2012 12:00:18 PM	(GMT08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		3/8/2012 12:00:18 PM	(GMT08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		3/8/2012 12:00:18 PM	(GMT08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		3/8/2012 12:00:18 PM	(GMT08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		3/8/2012 12:00:18 PM	(GMT08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		3/8/2012 12:00:18 PM	(GMT08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		3/8/2012 12:00:18 PM	(GMT08:00) Pacific Standard Time	UC Davis	D8XSMGH1