

2013-10-31_004.xad					Page	2 of 10
Assay Class: DNA 12000 Data Path: C:\ents and Electrophoresis File Run S	Created: Modified:		3 3:16:59 PM 3 3:38:07 PM			
Sample Name	Sample Comment	Rest. Digest	Status Observation	Resi	ılt Label	Result Color
Pp1290 gTube 5200 CaCal35 gTube 5200			×.			
Tim1 gTube 5200 Tim2 gTube 5200			×			
		ğ	•			
		Ā				
Ladder		$\Box$	~			
Chip Lot #			Reagent Kit Lot #			

Chip Comments :

## Electrophoresis Assay Details

#### **General Analysis Settings**

Number of Available Sample and Ladder Wells (Max.) : 13 Minimum Visible Range [s] : 20 Maximum Visible Range [s] : 99 Start Analysis Time Range [s] : 20 End Analysis Time Range [s] : 98.95 Ladder Concentration  $[ng/\mu I]$  : 44 Uses Standard Area for Ladder Fragments Lower Marker Concentration  $[ng/\mu I]$  : 8.3 Upper Marker Concentration  $[ng/\mu I]$  : 4.2 Used Upper Marker for Quantitation This is a Qualitative Assay Only Standard Curve Fit is Point to Point Show Data Aligned to Lower and Upper Marker

#### **Integrator Settings**

Integration Start Time [s] : 20 Integration End Time [s] : 98.95 Slope Threshold : 0.8 Height Threshold [FU] : 20 Area Threshold : 0.1 Width Threshold [s] : 0.5 Baseline Plateau [s] : 0.5

#### **Filter Settings**

Filter Width [s] : 0.5 Polynomial Order : 4

#### Ladder

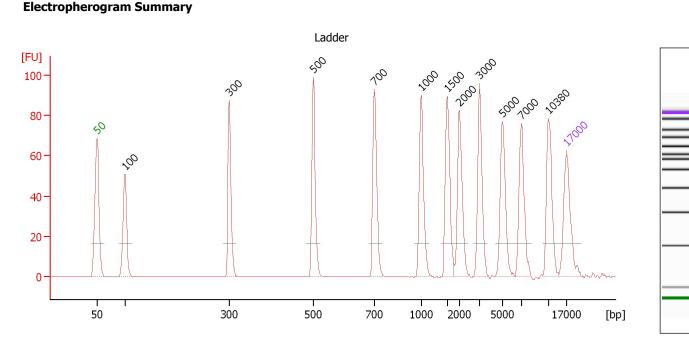
Ladder Peak	Size	Area
1	50	120
2	100	48
3	300	61
4	500	77
5	700	81
6	1000	86
7	1500	92
8	2000	92
9	3000	96
10	5000	100
11	7000	98
12	10380	102
13	17000	110

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Created:

Modified:

## Assay Class: DNA 12000 Data Path: C:\...ents and Settings\Bioanalyzer\2013-10-31\2013-10-31\_004.xad



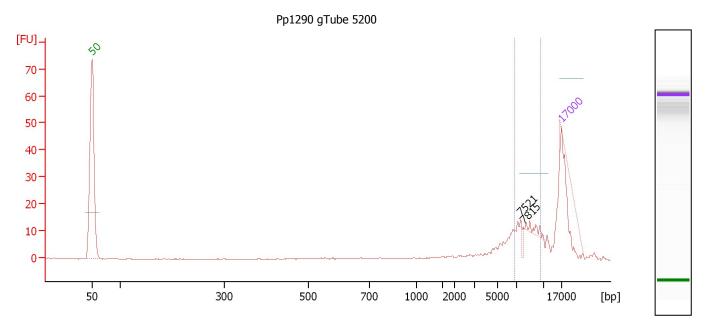
## Peak table for Ladder

Peak	Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	100	4.00	60.6	Ladder Peak
3	300	4.00	20.2	Ladder Peak
4	500	4.00	12.1	Ladder Peak
5	700	4.00	8.7	Ladder Peak
6	1,000	4.00	6.1	Ladder Peak
7	1,500	4.00	4.0	Ladder Peak
8	2,000	4.00	3.0	Ladder Peak
9	3,000	4.00	2.0	Ladder Peak
10	5,000	4.00	1.2	Ladder Peak
11	7,000	4.00	0.9	Ladder Peak
12	10,380	4.00	0.6	Ladder Peak
13	17,000	4.20	0.4	Upper Marker

# Assay Class:DNA 12000Created:Data Path:C:\...ents and Settings\Bioanalyzer\2013-10-31\2013-10-31\_004.xadModified:

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Electropherogram Summary Continued ...



Overall Results for sample 1 :	Pp1290 aTube 5200
Overall Results for sample 1:	Pp1290 grube 5200

Number of peaks found:	2	Area 1:	34.1

Peak	Peak table for sample 1 : Pp1290 gTube 5200										
Peak		Size [b	p] Conc.	[ng/µl]	Molari	ty [nmol/l]	Observa	tions			
1	-	50	8.30		251.5		Lower Ma	arker			
2		7,521	118.70		23.9						
3		7,815	42.18		8.2						
4	6	17,000	4.20		0.4		Upper Ma	arker			
Regio	on tab	le for sa	mple 1 : <u>F</u>	2 <u>p1290 gTu</u>	<u>be 5200</u>						
From	[bp]	To [bp]	Average Size	Conc. [ng	J∕µl] Area	Col Size distribut	ion in CV	% of Total			
			[bp]			<u>or [%]</u>					
6,814		10,066	8,332	889.18	34.1	11.4		38			

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Assay Class: Data Path: <b>Electrophe</b>		nd Settings\Bioanalyze mary Continued	r\2013-10-31\2013-10-31_(	Created: 10/31/2013 3:16:59 PM 004.xad Modified: 10/31/2013 3:38:07 PM
			CaCal35 gTube 5200	
[FU] 70 - 60 - 50 - 40 - 30 - 20 - 10 -	\$			TOSO I
	50	300	500 700 1	.000 2000 5000 17000 [bp]
Number of pe	aults for sampl aks found: for sample 2 :	1	Area 1:	93.1
Peak	Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2 3	10,877 17,000	2.47 4.20	0.3 0.4	Upper Marker
Region tabl	le for sample 2	2 : <u>CaCal35 gT</u>	<u>ube 5200</u>	

Region ta	ble for sa	ample 2 :	CaCal35 gTu	<u>be 5200</u>		
From [bp] To [bp] Average Size		Conc. [ng/µl] Area		Col Size distribution in CV	% of Total	
		[bp]			<u>or [%]</u>	
5,001	32,612	14,619	2.88	93.1	35.2	89

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Assay Cla Data Pat	lass: DNA 12000 th: C:\ents and Settings\Bioanalyzer\2013-10-31\2013-10- pherogram Summary Continued	31_004.xad Created: Modified:	10/31/2013 3:16:59 PM 10/31/2013 3:38:07 PM
LIECUU			
	Tim1 gTube 5200		
[FU]-]	4 <sup>0</sup>		
70 -		0	
60 -		-1 <sup>00</sup>	
50 -		- ()	
40-		Str	
30 -			
20 -			
10-			
0		Min	
	50 300 500 700	1000 2000 5000 17000 [b	p]

#### Overall Results for sample 3 : Tim1 gTube 5200

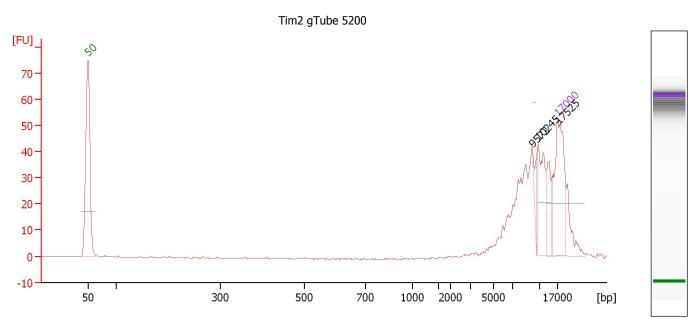
1

Number of peaks found:

Peak table for sample 3 :		for sample 3 :	<u>Tim1 gTube 5200</u>	<u>)</u>	
Peak		Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1		50	8.30	251.5	Lower Marker
2		9,812	4.00	0.6	
3		17,000	4.20	0.4	Upper Marker

#### 10/31/2013 3:16:59 PM 10/31/2013 3:38:07 PM Assay Class: DNA 12000 Created: C:\...ents and Settings\Bioanalyzer\2013-10-31\2013-10-31\_004.xad Modified: Data Path:

**Electropherogram Summary Continued ...** 



#### **Overall Results for sample 4 :** Tim2 gTube 5200 2

Number of peaks found:

Peak table for sample 4 :		for sample 4 :	Tim2 gTube 5200	<u>)</u>	
Peak		Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1		50	8.30	251.5	Lower Marker
2		9,572	0.31	0.0	
3		10,245	2.85	0.4	
4	B	17,000	4.20	0.4	Upper Marker
5		17,525	0.00	0.0	

Assay Clas Data Path <b>Gel Ima</b> g		A 12000 ents and	Settings\I	Bioanalyze	er\2013-10	)-31\2013-	10-31_004	.xad		Created Modified	: 10/31/ : 10/31/	2013 3:16:59 PM 2013 3:38:07 PM
[bp]	Ladder	Pp1290 qTu	Galss afu	Tim1 qTube	Tim2 qTube							
17000 — 10380 — 7000 — 5000 — 3000 — 2000 — 1500 —												
1500 — 1000 — 700 —	_											
500 —												
300 —												
100 — 50 —	_											
	L	1	2	3	4	5	6	7	8	9 10	11	12

Sample 5 has not been run, no results available. Sample 6 has not been run, no results available. Sample 7 has not been run, no results available. Sample 8 has not been run, no results available. Sample 9 has not been run, no results available. Sample 10 has not been run, no results available. Sample 11 has not been run, no results available. Sample 12 has not been run, no results available. Created: 10/31/2013 3:16:59 PM Modified: 10/31/2013 3:38:07 PM