

Assay Class: DNA 12000  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-23\2016-05-23\_001.xad

Created: 5/23/2016 11:08:57 AM  
Modified: 5/23/2016 11:35:28 AM

**Electrophoresis File Run Summary**

Instrument Information:

Instrument Name: DE13701086      Firmware: C.01.069  
Serial#: DE13701086              Type: G2938B

Assay Information:

Assay Origin Path: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\assays\dsDNA\DNA 12000 Series II.xsy

Assay Class: DNA 12000

Version: 2.4

Assay Comments: DNA Analysis 100 -12000 bp

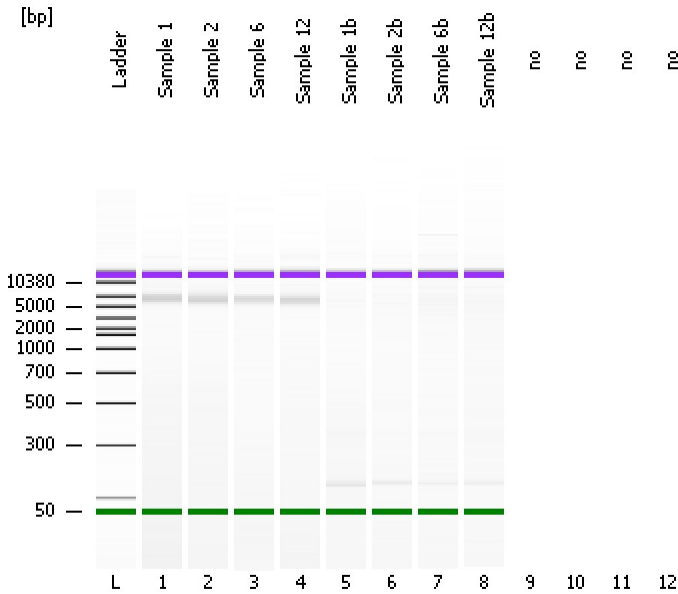
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Chip Information:

Chip Lot #:

Reagent Kit Lot #:

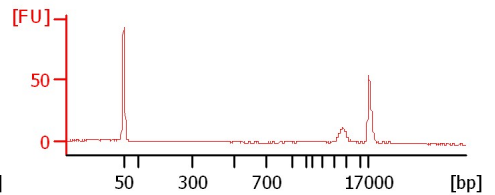
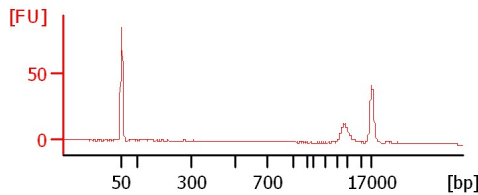
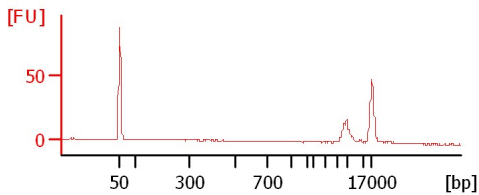
Chip Comments:



**Sample 1**

**Sample 2**

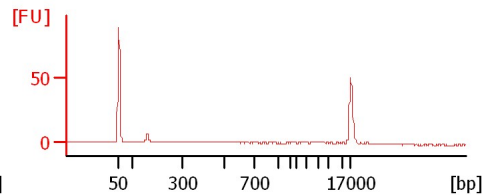
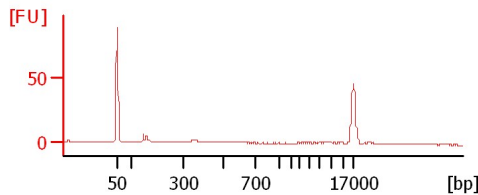
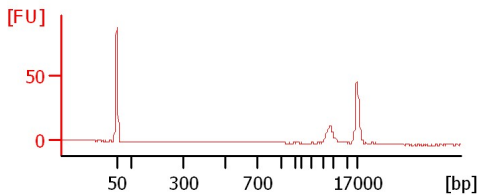
**Sample 6**



**Sample 12**

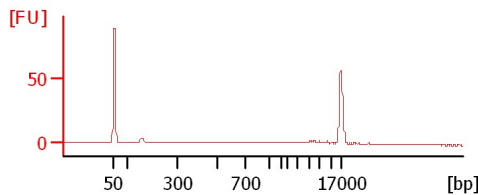
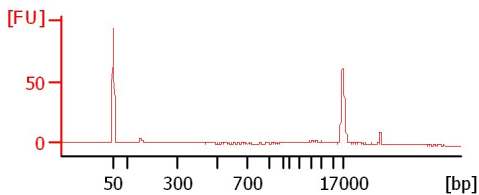
**Sample 1b**

**Sample 2b**



**Sample 6b**

**Sample 12b**



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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Sample 1		<input type="checkbox"/>	✓			
Sample 2		<input type="checkbox"/>	✓			
Sample 6		<input type="checkbox"/>	✓			
Sample 12		<input type="checkbox"/>	✓			
Sample 1b		<input type="checkbox"/>	✓			
Sample 2b		<input type="checkbox"/>	✓			
Sample 6b		<input type="checkbox"/>	✓			
Sample 12b		<input type="checkbox"/>	✓			
no		<input type="checkbox"/>				
no		<input type="checkbox"/>				
no		<input type="checkbox"/>				
no		<input type="checkbox"/>				
Ladder		<input type="checkbox"/>	✓			

**Chip Lot #**

**Reagent Kit Lot #**

**Chip Comments :**

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**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/μl] : 44  
Uses Standard Area for Ladder Fragments  
Lower Marker Concentration [ng/μl] : 8.3  
Upper Marker Concentration [ng/μl] : 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

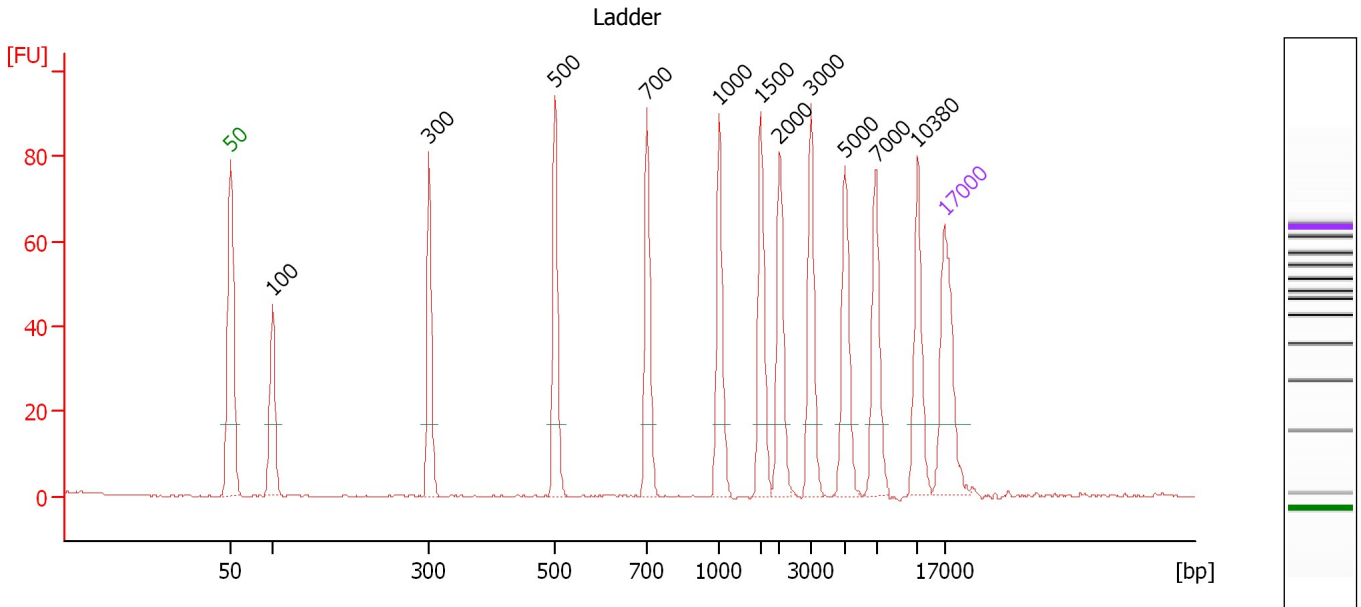
**Ladder**

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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**Electropherogram Summary**



**Peak table for Ladder**

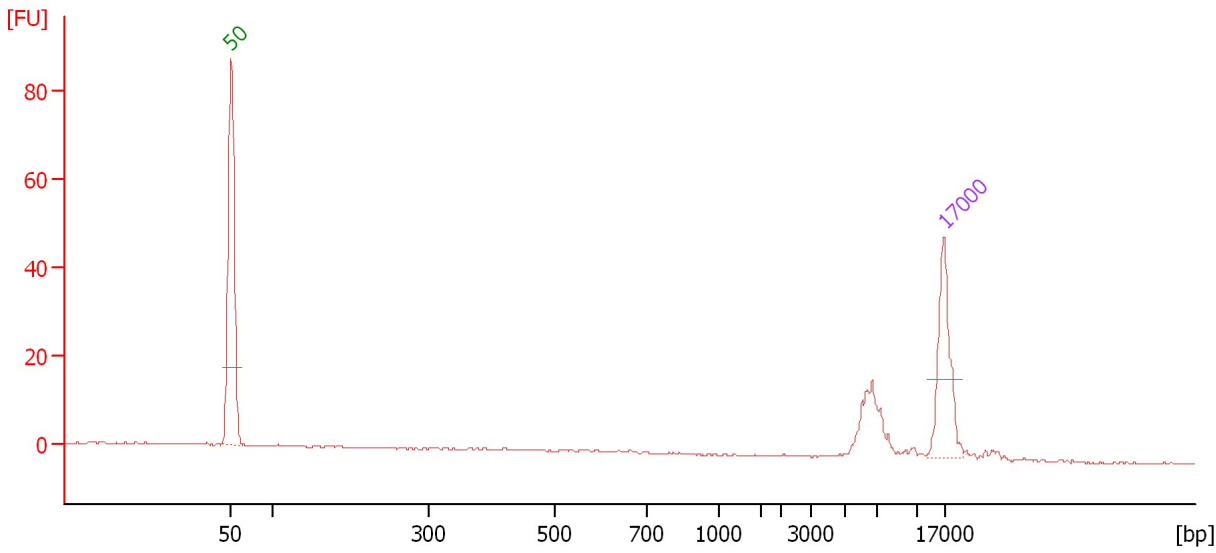
Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	100	4.00	60.6	Ladder Peak
3	300	4.00	20.2	Ladder Peak
4	500	4.00	12.1	Ladder Peak
5	700	4.00	8.7	Ladder Peak
6	1,000	4.00	6.1	Ladder Peak
7	1,500	4.00	4.0	Ladder Peak
8	2,000	4.00	3.0	Ladder Peak
9	3,000	4.00	2.0	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

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**Electropherogram Summary Continued ...**

Sample 1



**Overall Results for sample 1 : Sample 1**

Number of peaks found: 0

**Peak table for sample 1 : Sample 1**

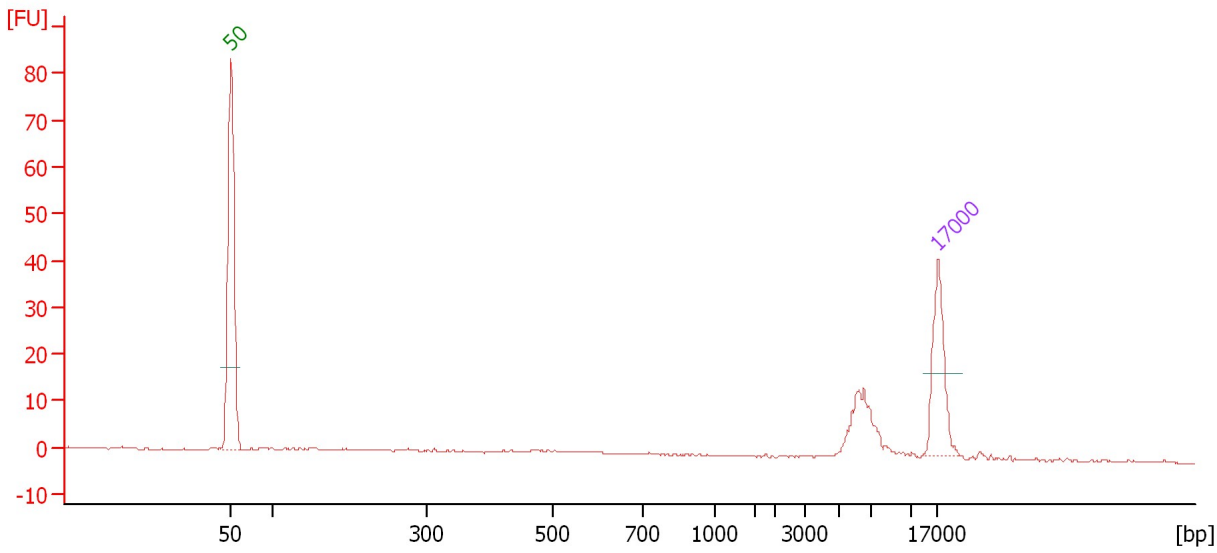
Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

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**Electropherogram Summary Continued ...**

Sample 2



**Overall Results for sample 2 : Sample 2**

Number of peaks found: 0

**Peak table for sample 2 : Sample 2**

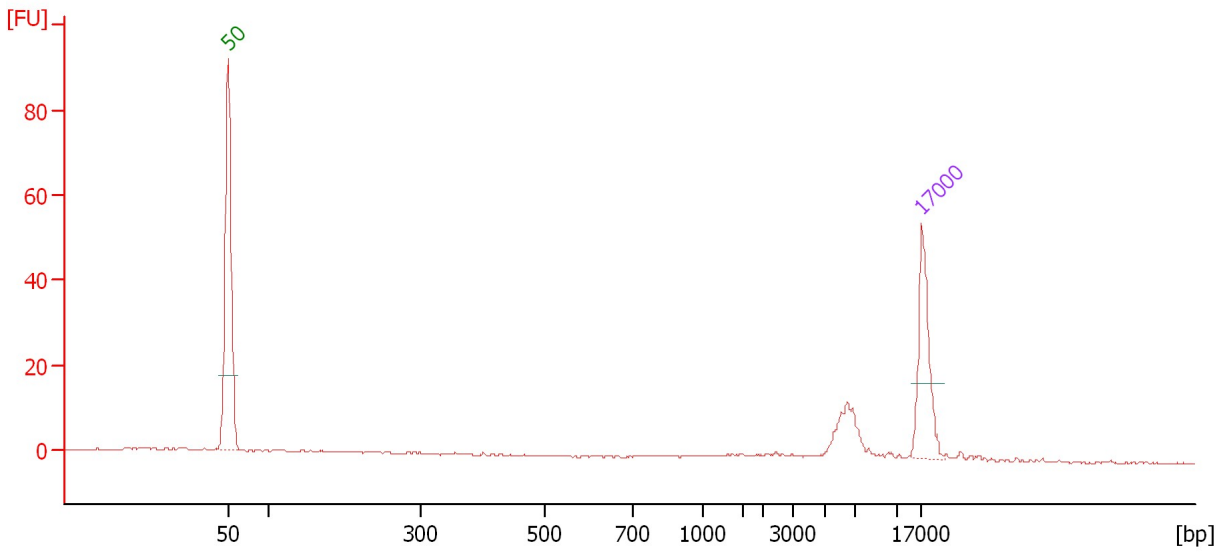
Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

Assay Class: DNA 12000  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-23\2016-05-23\_001.xad

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**Electropherogram Summary Continued ...**

Sample 6



**Overall Results for sample 3 : Sample 6**

Number of peaks found: 0

**Peak table for sample 3 : Sample 6**

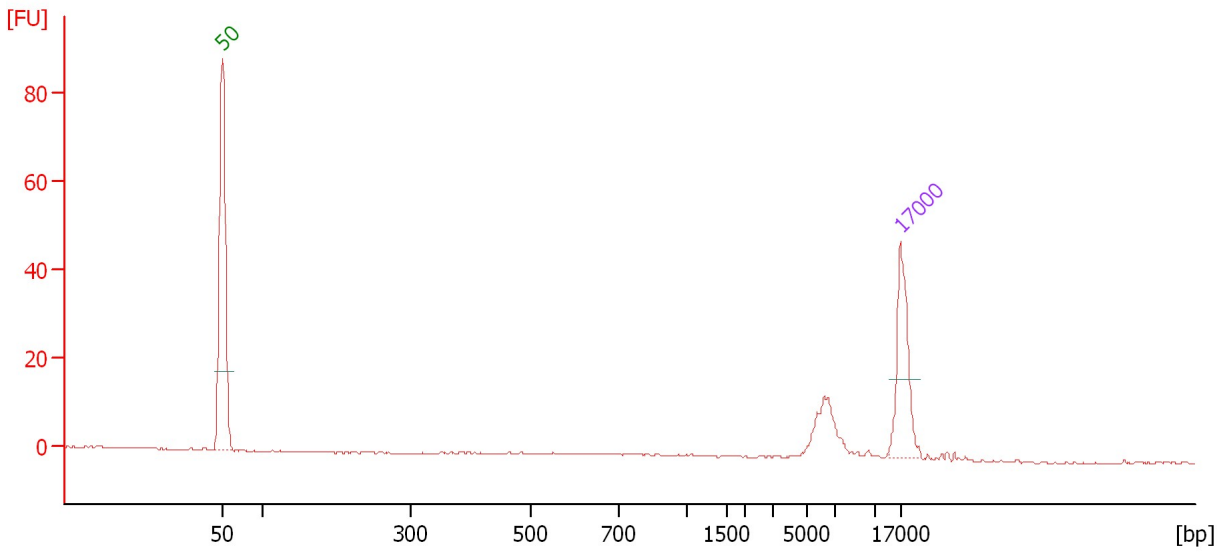
Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

Assay Class: DNA 12000  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-23\2016-05-23\_001.xad

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**Electropherogram Summary Continued ...**

Sample 12



**Overall Results for sample 4 : Sample 12**

Number of peaks found: 0

**Peak table for sample 4 : Sample 12**

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

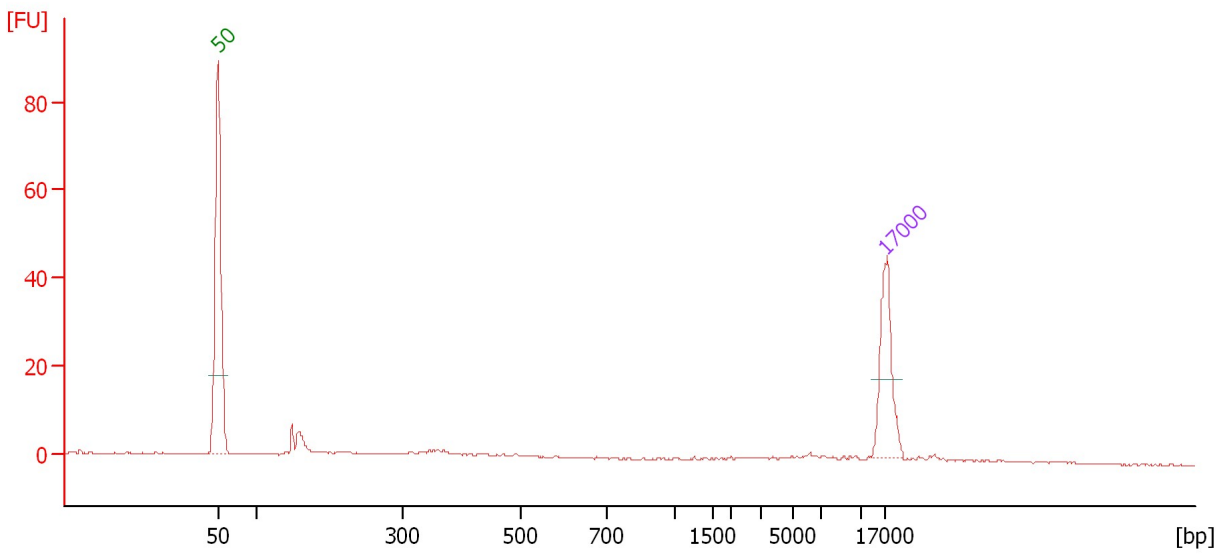


Assay Class: DNA 12000  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-23\2016-05-23\_001.xad

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**Electropherogram Summary Continued ...**

Sample 1b



**Overall Results for sample 5 : Sample 1b**

Number of peaks found: 0

**Peak table for sample 5 : Sample 1b**

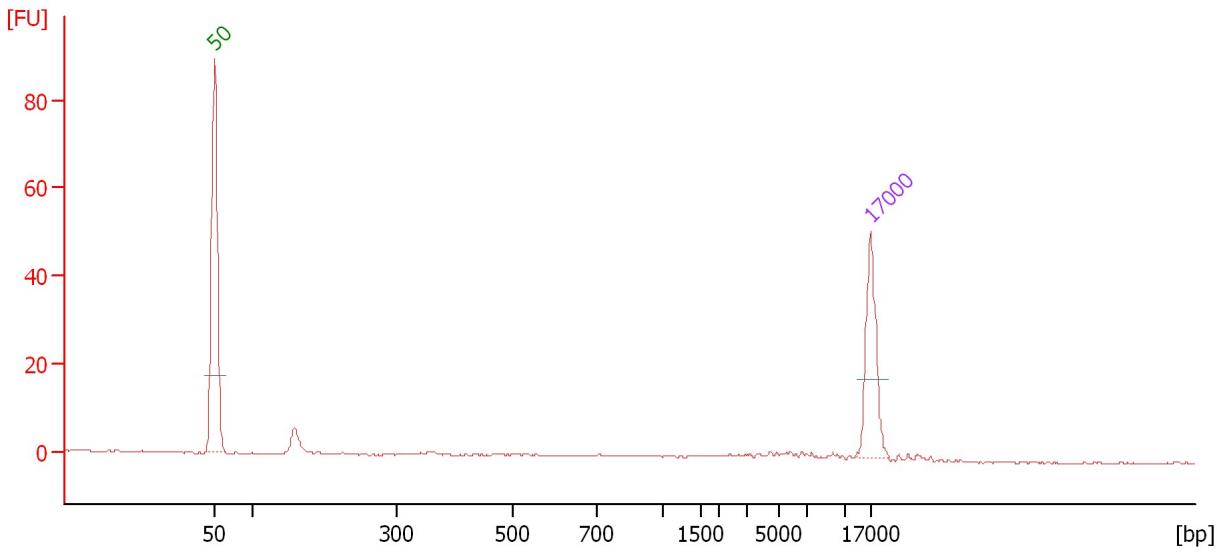
Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

Assay Class: DNA 12000  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-23\2016-05-23\_001.xad

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**Electropherogram Summary Continued ...**

Sample 2b



**Overall Results for sample 6 : Sample 2b**

Number of peaks found: 0

**Peak table for sample 6 : Sample 2b**

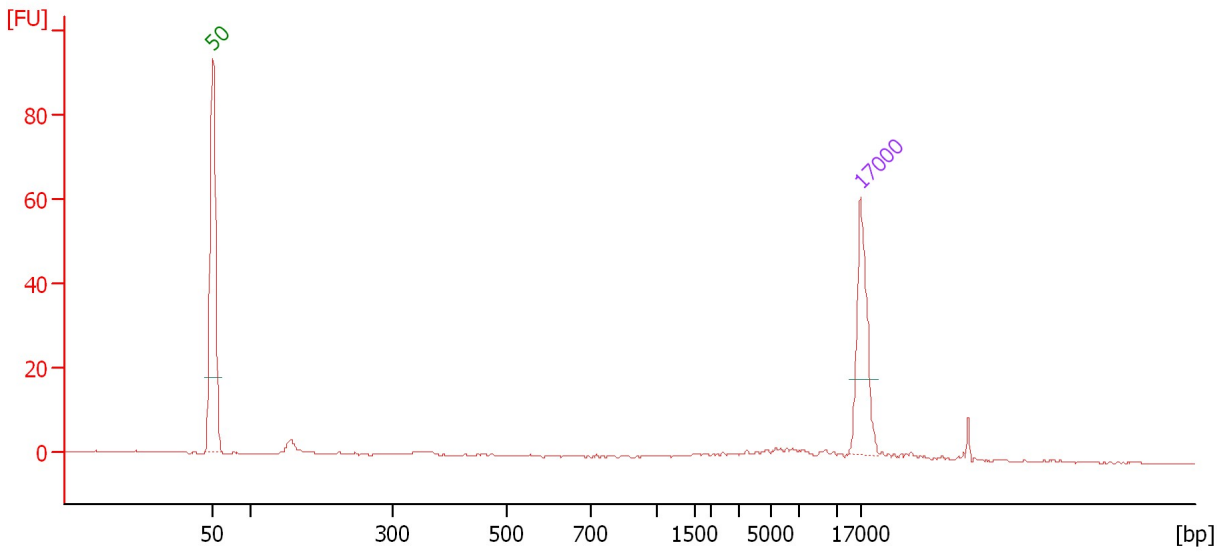
Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

Assay Class: DNA 12000  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-23\2016-05-23\_001.xad

Created: 5/23/2016 11:08:57 AM  
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**Electropherogram Summary Continued ...**

Sample 6b



**Overall Results for sample 7 : Sample 6b**

Number of peaks found: 0

**Peak table for sample 7 : Sample 6b**

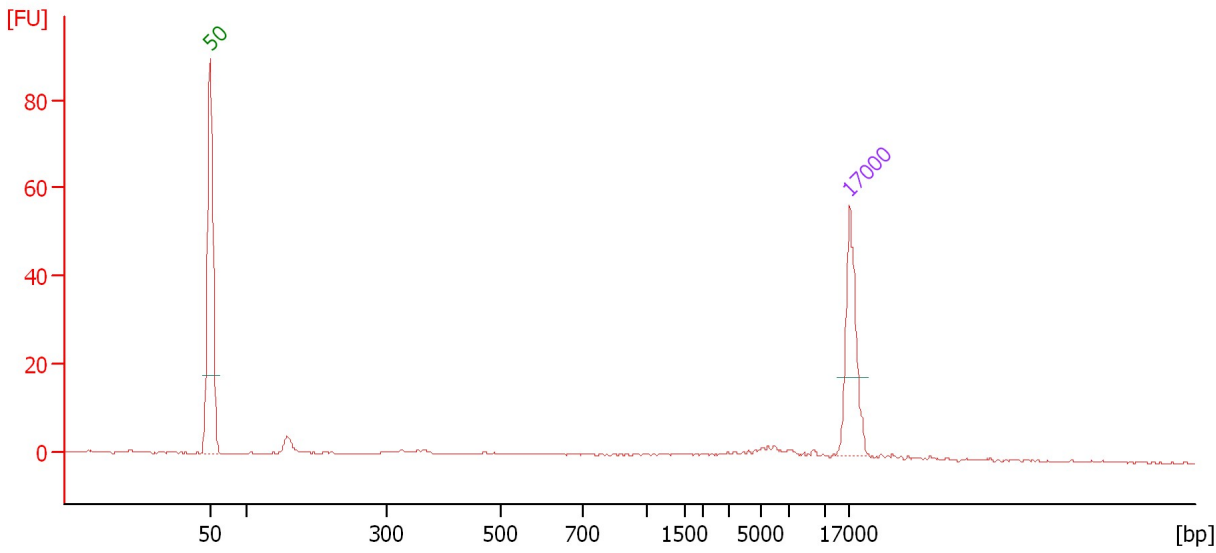
Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

Assay Class: DNA 12000  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-23\2016-05-23\_001.xad

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**Electropherogram Summary Continued ...**

Sample 12b



**Overall Results for sample 8 : Sample 12b**

Number of peaks found: 0

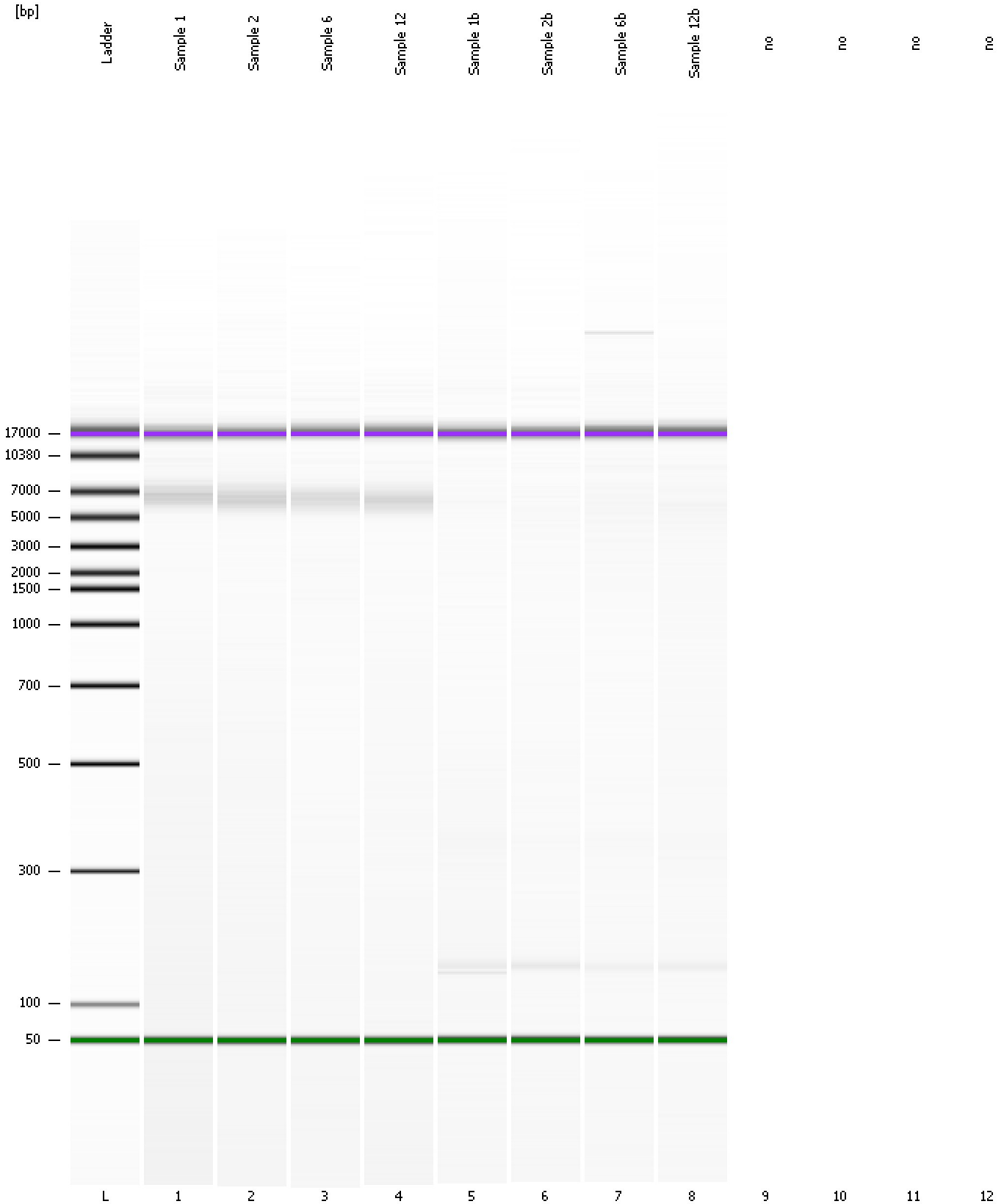
**Peak table for sample 8 : Sample 12b**

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

Assay Class: DNA 12000  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-23\2016-05-23\_001.xad

Created: 5/23/2016 11:08:57 AM  
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**Gel Image**



Assay Class: DNA 12000  
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**Invalid Samples**

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.

Assay Class: DNA 12000  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-23\2016-05-23\_001.xad

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**Run Logbook**

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 9)		Instrument	Run		5/23/2016 11:35:28 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-05-23\2016-05-23_001.xad)		Instrument	Run		5/23/2016 11:09:03 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		5/23/2016 11:09:03 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		5/23/2016 11:09:03 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		5/23/2016 11:09:03 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		5/23/2016 11:09:03 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		5/23/2016 11:09:03 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		5/23/2016 11:09:03 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1