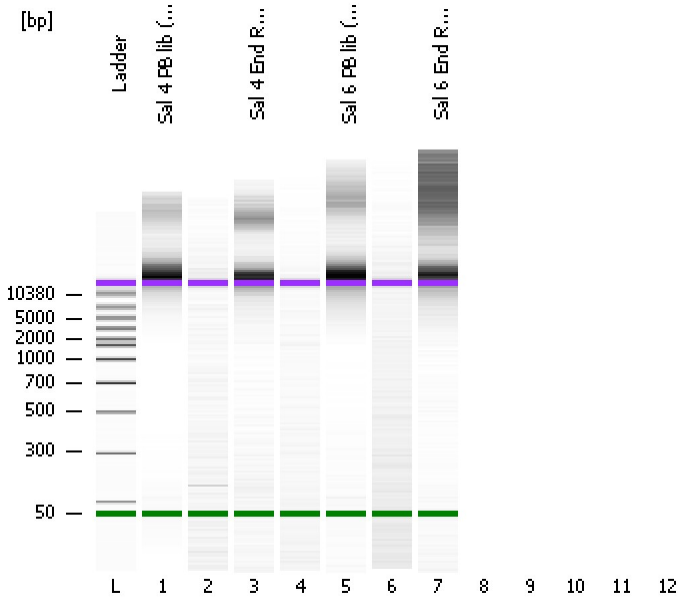


Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
Modified: 10/15/2015 11:33:20 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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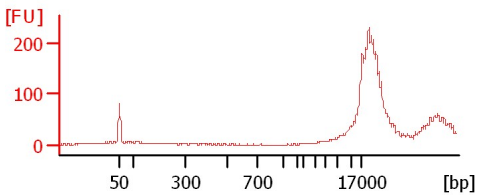
Chip Information:

Chip Lot #:

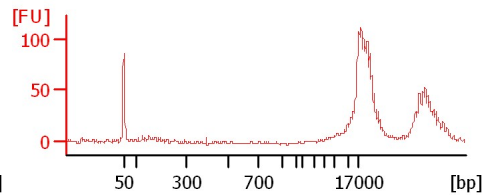
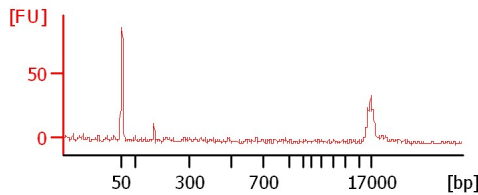
Reagent Kit Lot #:

Chip Comments:

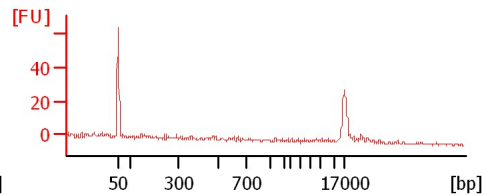
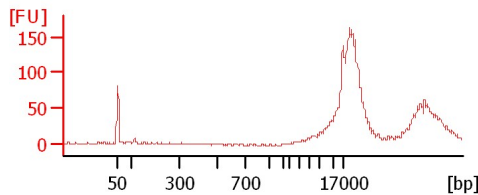
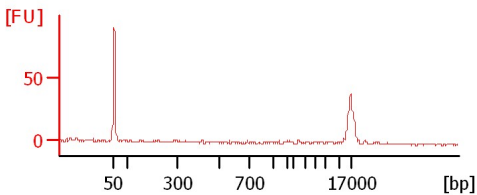
Sal 4 PB lib (1:2)



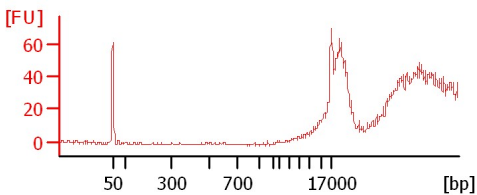
Sal 4 End Repair



Sal 6 PB lib (1:2)



Sal 6 End Repair



Assay Class: DNA 12000
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
 Modified: 10/15/2015 11:33:20 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Sal 4 PB lib (1:2)		<input type="checkbox"/>	✓			
Sal 4 End Repair		<input type="checkbox"/>	✓			
Sal 6 PB lib (1:2)		<input type="checkbox"/>	✓			
Sal 6 End Repair		<input type="checkbox"/>	✓			
		<input type="checkbox"/>				
		<input type="checkbox"/>				
		<input type="checkbox"/>				
		<input type="checkbox"/>				
Ladder		<input type="checkbox"/>	✓			
Chip Lot #				Reagent Kit Lot #		

Chip Comments :

Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
Modified: 10/15/2015 11:33:20 AM

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

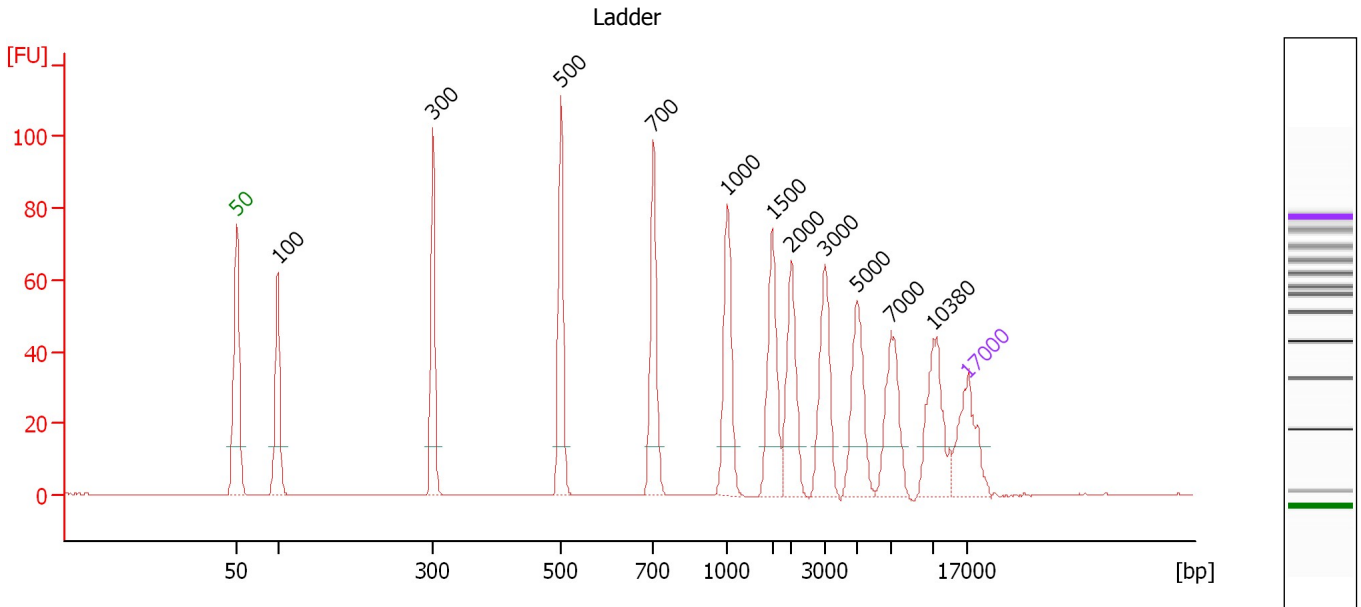
Ladder

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

Assay Class: DNA 12000
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
 Modified: 10/15/2015 11:33:20 AM

Electropherogram Summary



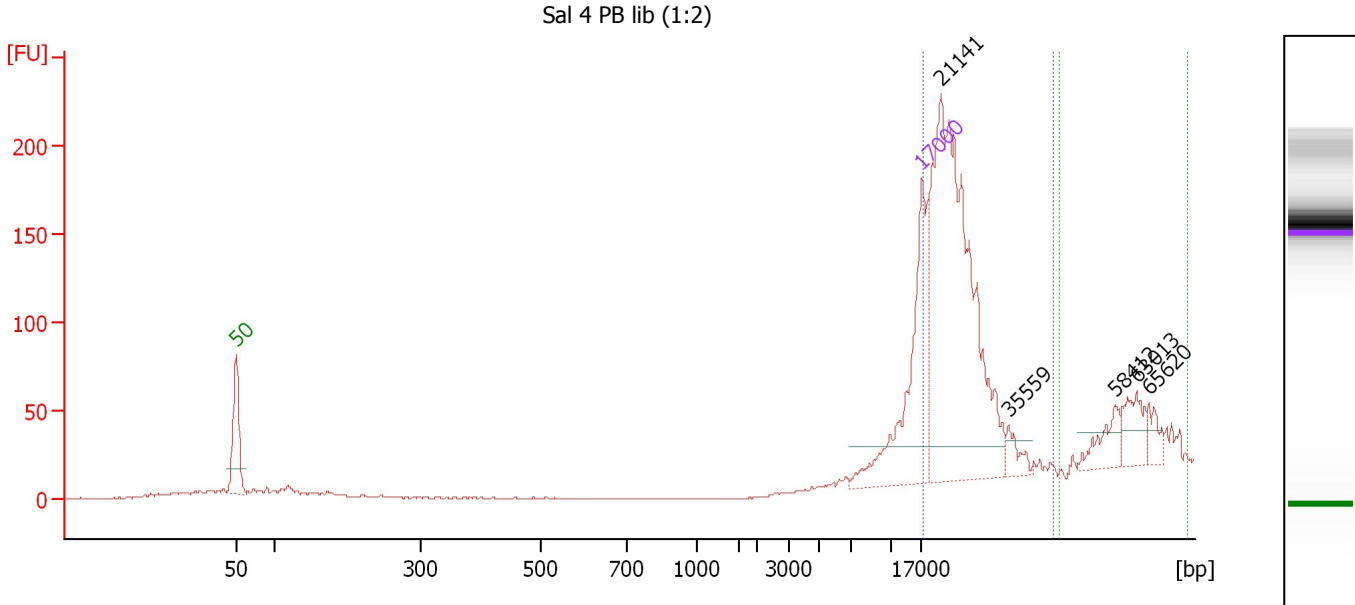
Peak table for Ladder

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

Assay Class: DNA 12000
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
 Modified: 10/15/2015 11:33:20 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : Sal 4 PB lib (1:2)

Number of peaks found: 0 Area 2: 161.4
 Area 1: 669.0

Peak table for sample 1 : Sal 4 PB lib (1:2)

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
3	21,141	0.00	0.0	
4	35,559	0.00	0.0	
5	58,412	0.00	0.0	
6	63,013	0.00	0.0	
7	65,620	0.00	0.0	

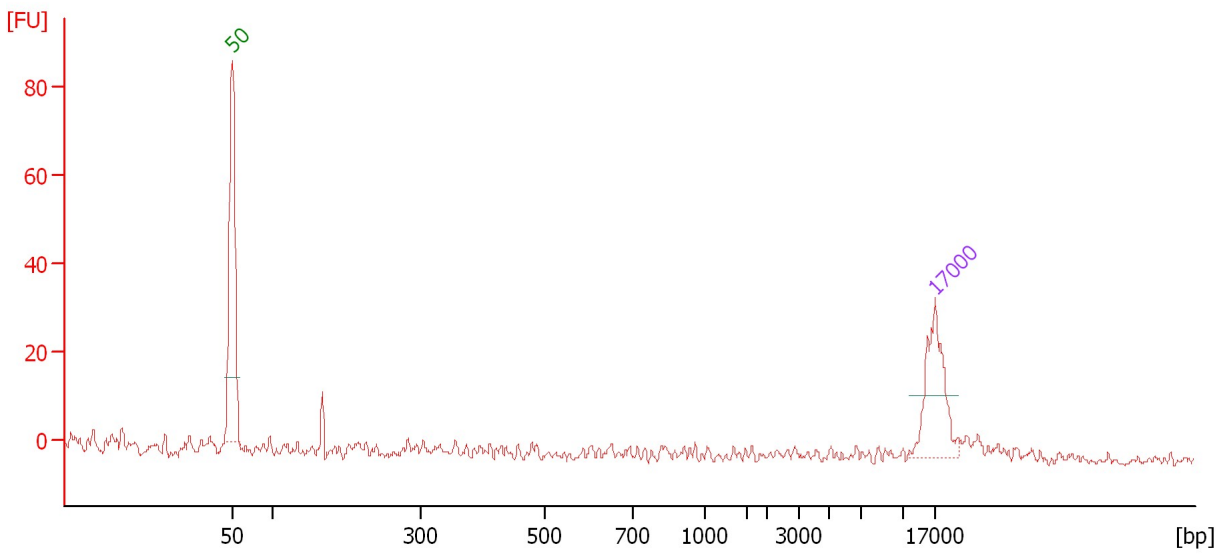
Region table for sample 1 : Sal 4 PB lib (1:2)

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
17,185	45,182	24,023	10.54	669.0	80	19.8	Blue
46,537	74,082	62,500	2.19	161.4	19	8.4	Green

Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
Modified: 10/15/2015 11:33:20 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 :

Number of peaks found: 0

Peak table for sample 2 :

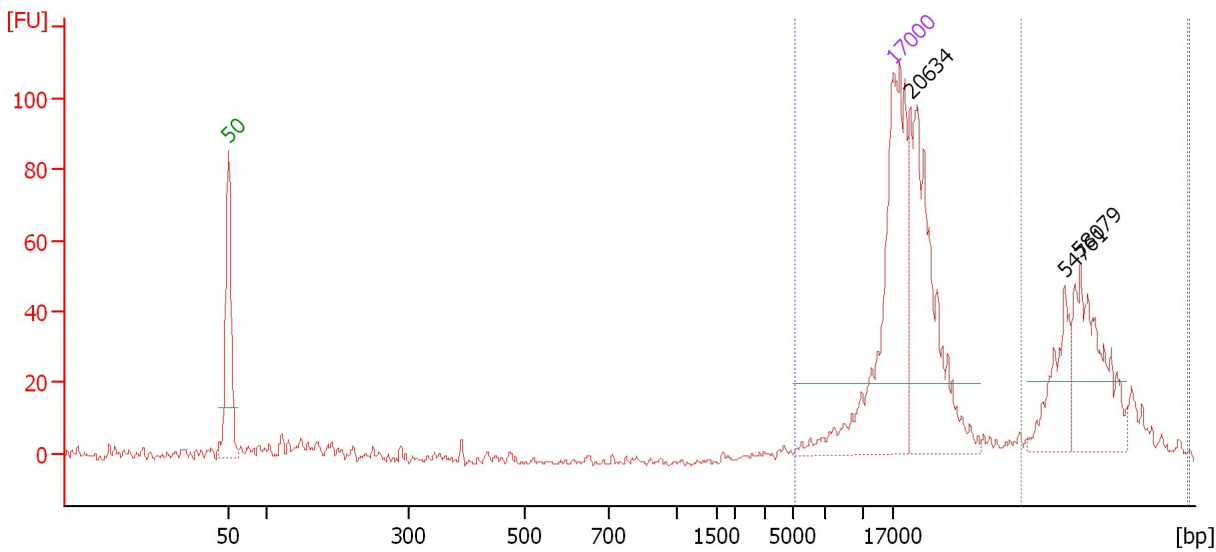
Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

Assay Class: DNA 12000
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
 Modified: 10/15/2015 11:33:20 AM

Electropherogram Summary Continued ...

Sal 4 End Repair



Overall Results for sample 3 : Sal 4 End Repair

Number of peaks found: 0 Area 2: 216.4
 Area 1: 423.1

Peak table for sample 3 : Sal 4 End Repair

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
3	20,634	0.00	0.0	
4	54,761	0.00	0.0	
5	58,079	0.00	0.0	

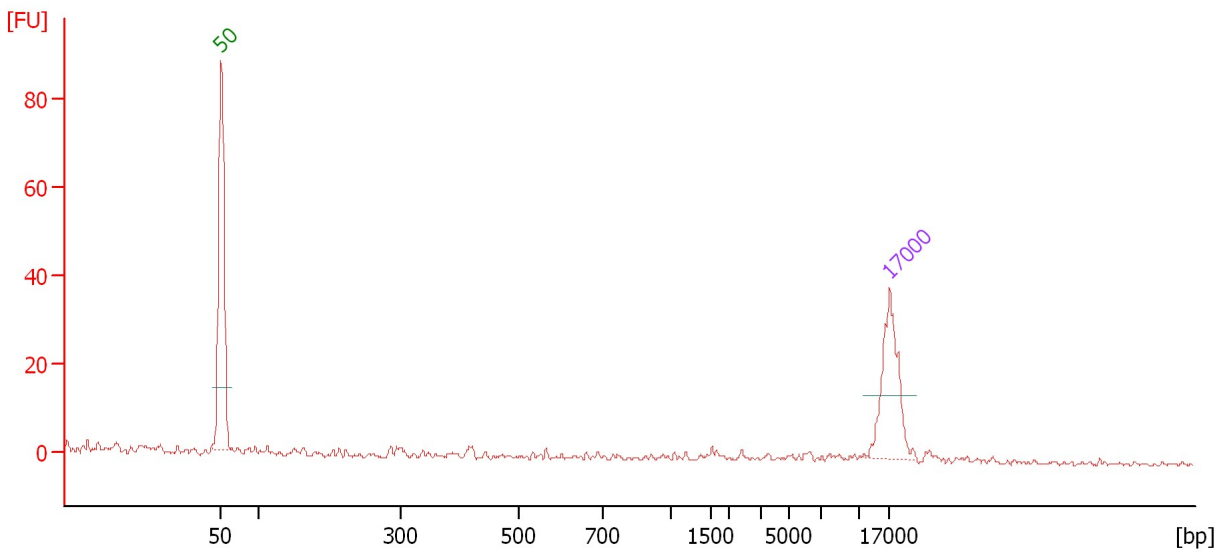
Region table for sample 3 : Sal 4 End Repair

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
5,120	82,420	33,450	7.12	423.1	89	59.3	Blue
44,929	81,952	59,838	3.39	216.4	46	11.2	Green

Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
Modified: 10/15/2015 11:33:20 AM

Electropherogram Summary Continued ...



Overall Results for sample 4 :

Number of peaks found: 0

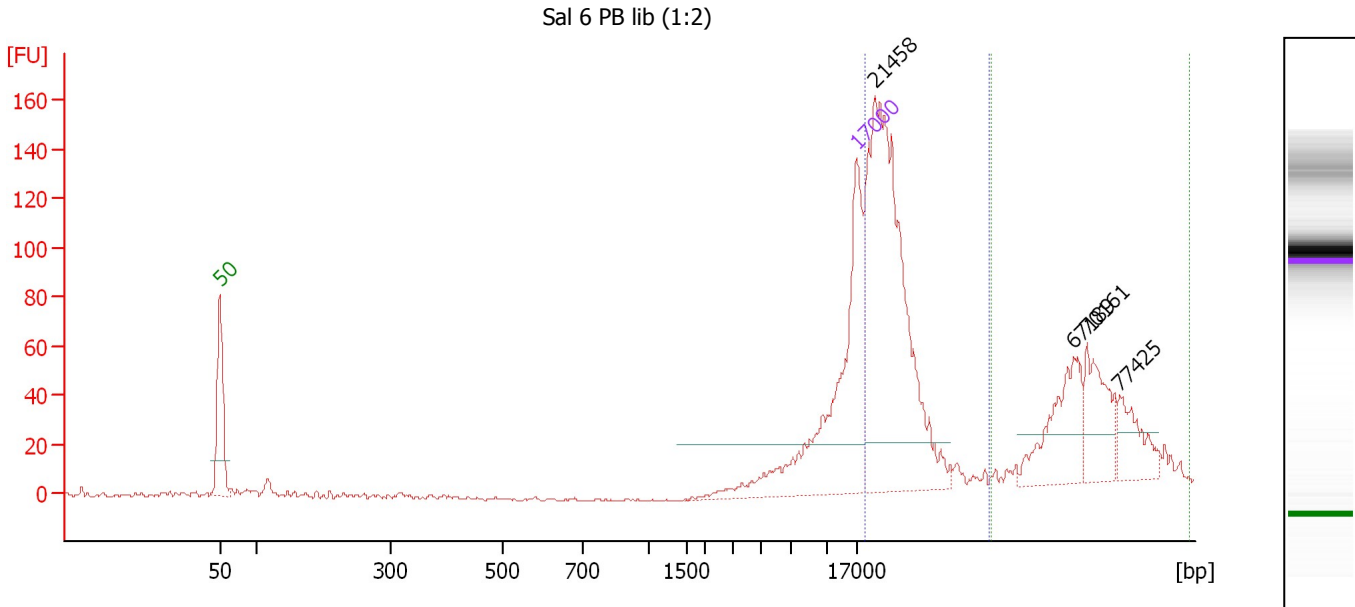
Peak table for sample 4 :

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:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
 Modified: 10/15/2015 11:33:20 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : Sal 6 PB lib (1:2)

Number of peaks found: 0 Area 2: 317.9
 Area 1: 472.5

Peak table for sample 5 : Sal 6 PB lib (1:2)

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
3	21,458	0.00	0.0	
4	67,189	0.00	0.0	
5	70,161	0.00	0.0	
6	77,425	0.00	0.0	

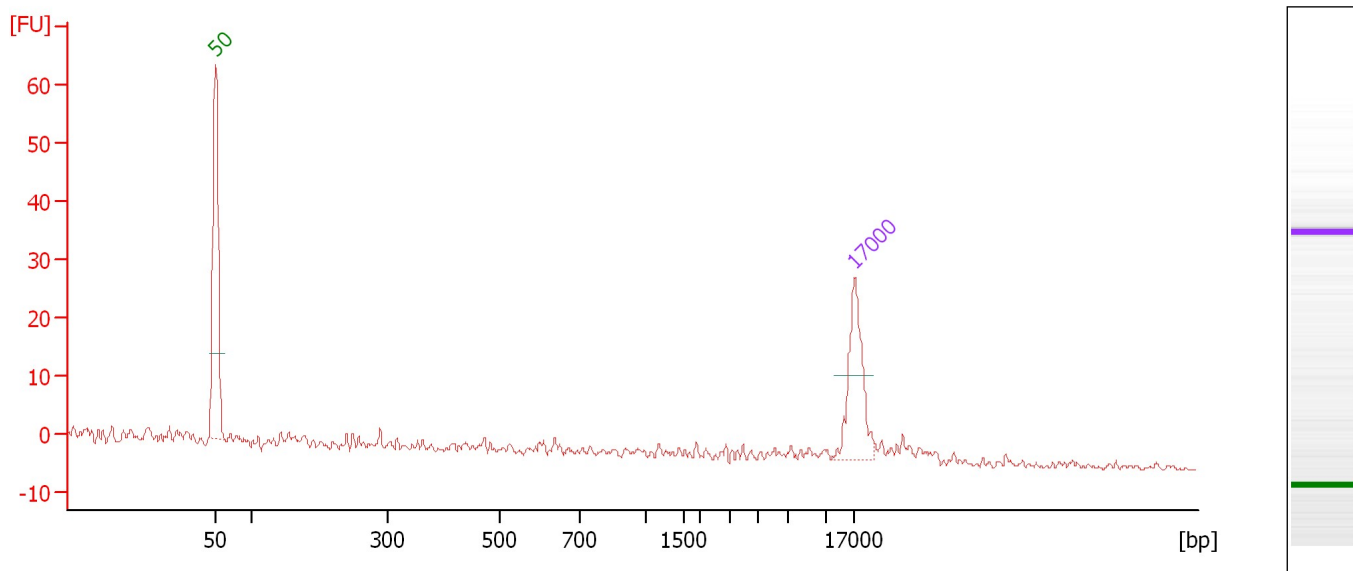
Region table for sample 5 : Sal 6 PB lib (1:2)

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
19,020	47,423	25,160	6.68	472.5	59	19.0	Blue
47,913	93,946	71,510	3.76	317.9	40	12.3	Green

Assay Class: DNA 12000
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
 Modified: 10/15/2015 11:33:20 AM

Electropherogram Summary Continued ...



Overall Results for sample 6 :

Number of peaks found: 0

Peak table for sample 6 :

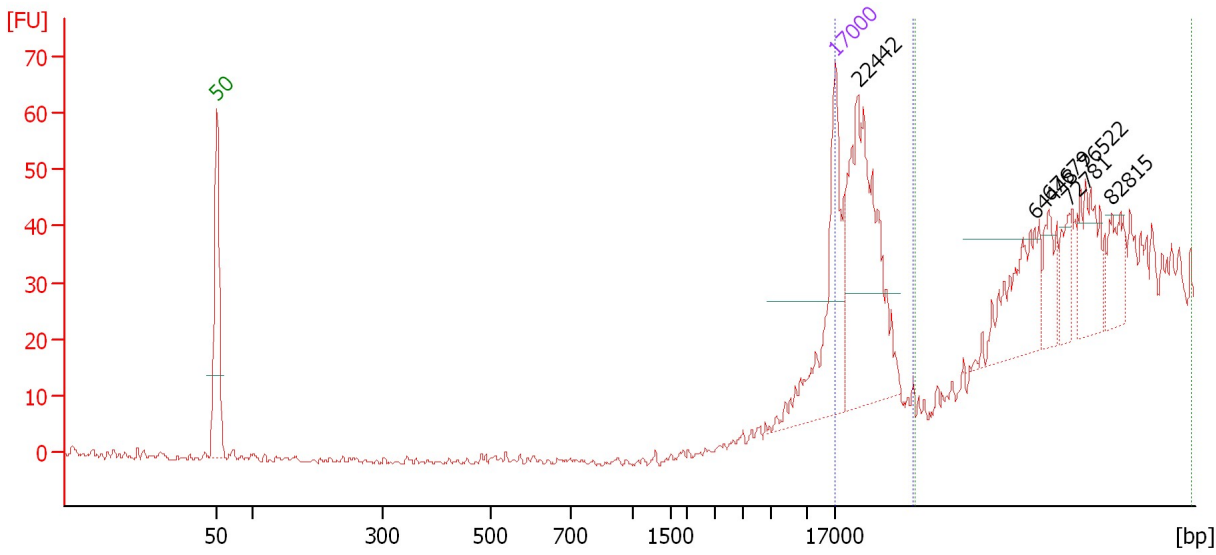
Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

Assay Class: DNA 12000
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
 Modified: 10/15/2015 11:33:20 AM

Electropherogram Summary Continued ...

Sal 6 End Repair



Overall Results for sample 7 : Sal 6 End Repair

Number of peaks found: 0 Area 2: 122.7
 Area 1: 79.9

Peak table for sample 7 : Sal 6 End Repair

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
3	22,442	0.00	0.0	
4	64,448	0.00	0.0	
5	67,679	0.00	0.0	
6	72,781	0.00	0.0	
7	76,522	0.00	0.0	
8	82,815	0.00	0.0	

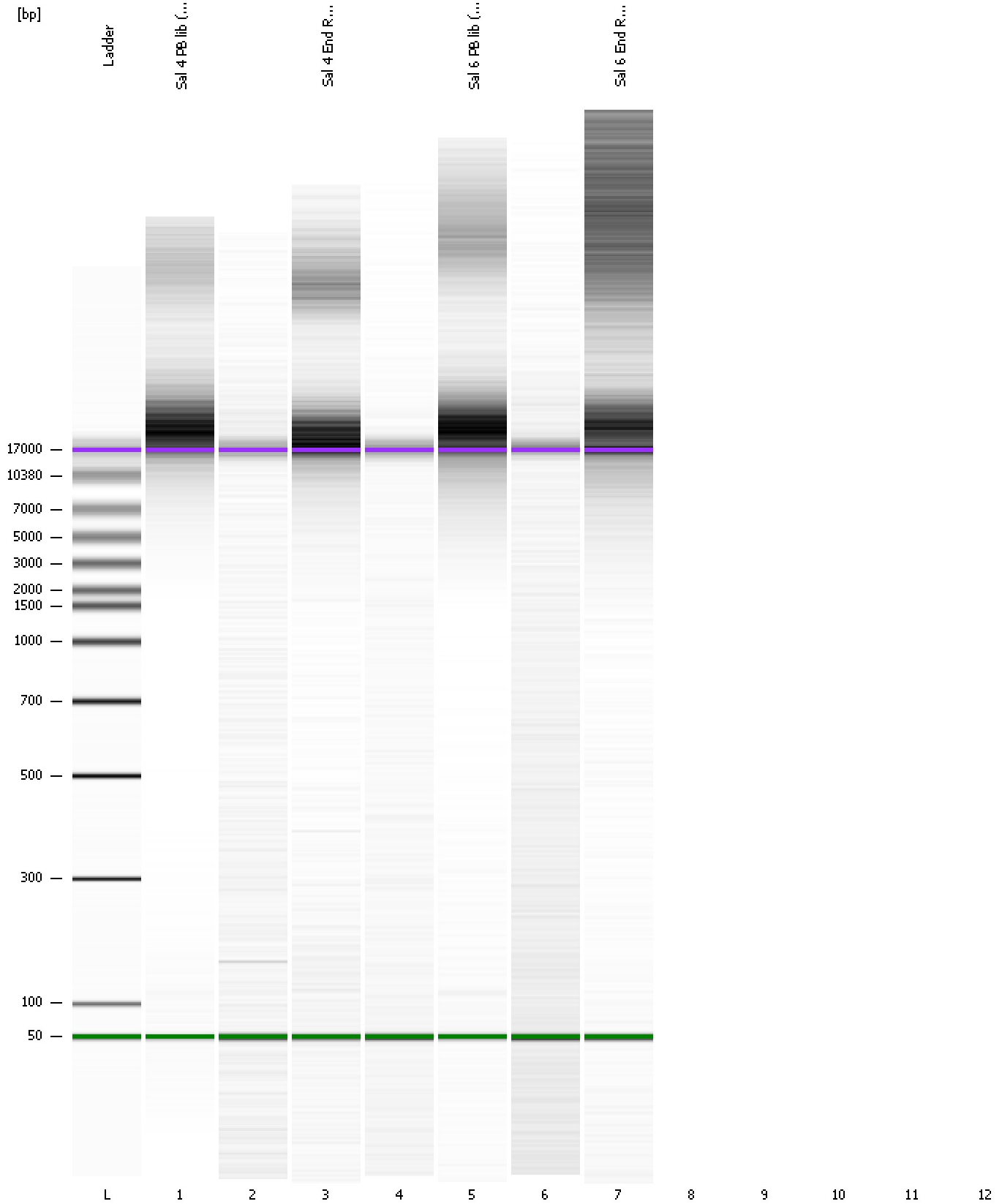
Region table for sample 7 : Sal 6 End Repair

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
17,120	35,280	22,371	4.18	79.9	39	14.5	Blue
35,785	101,862	76,200	5.22	122.7	60	13.6	Green

Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
Modified: 10/15/2015 11:33:20 AM

Gel Image

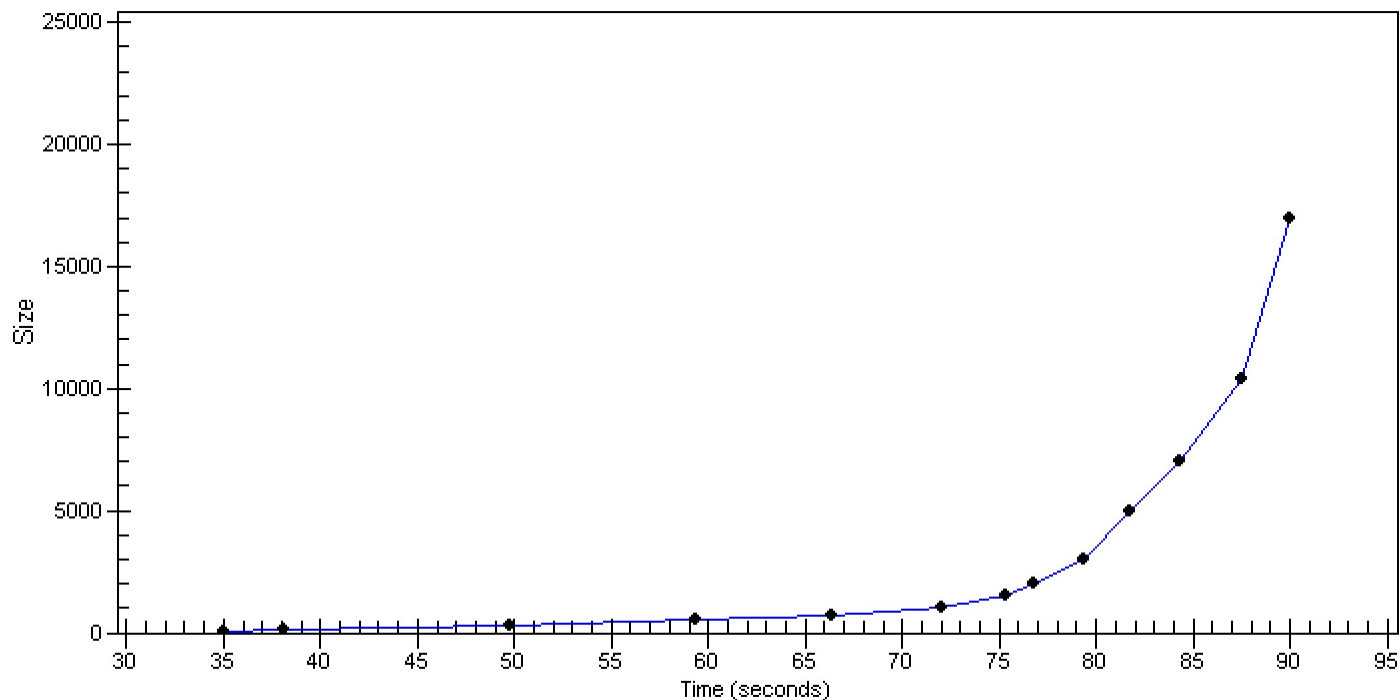


Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
Modified: 10/15/2015 11:33:20 AM

Curves

Standard Curve



Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
Modified: 10/15/2015 11:33:20 AM

Invalid Samples

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.

Assay Class: DNA 12000
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad

Created: 10/15/2015 11:01:13 AM
 Modified: 10/15/2015 11:33:20 AM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 8)		Instrument	Run		10/15/2015 11:25:36 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-10-15\2015-10-15_001.xad)		Instrument	Run		10/15/2015 11:01:18 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		10/15/2015 11:01:18 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		10/15/2015 11:01:18 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		10/15/2015 11:01:18 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		10/15/2015 11:01:18 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		10/15/2015 11:01:18 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		10/15/2015 11:01:18 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1