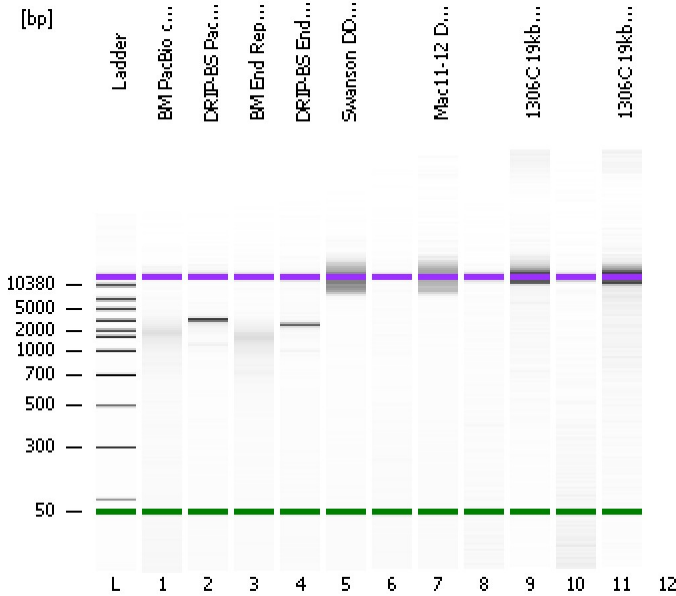


Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
Modified: 10/7/2015 11:22:25 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

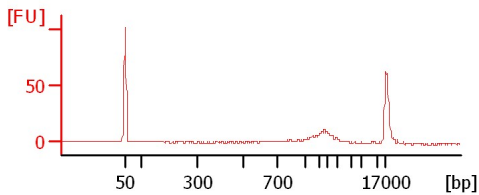
Assay Class: DNA 12000
Version: 2.4
Assay Comments: DNA Analysis 100 -12000 bp

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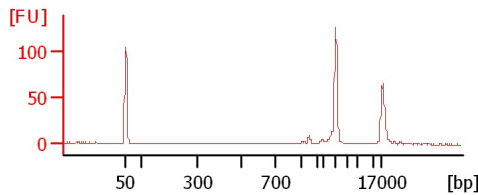
Chip Information:

Chip Lot #:
Reagent Kit Lot #:
Chip Comments:

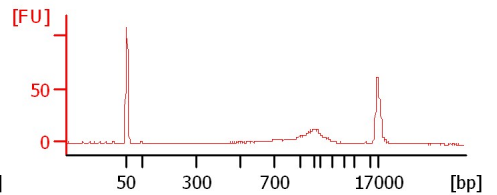
BM PacBio cDNA Lib (1:2)



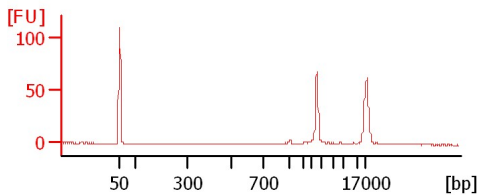
DRIP-BS PacBio amplicon lib (1:2)



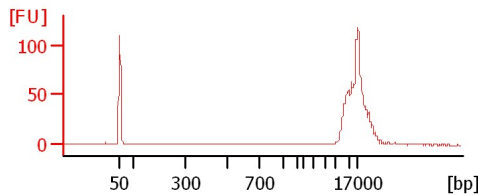
BM End Repair (1:3)



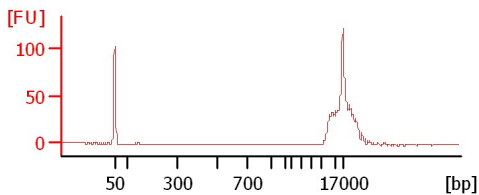
DRIP-BS End Repair (1:3)



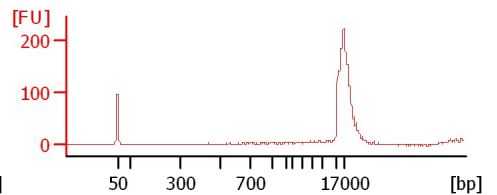
Swanson DDR lib



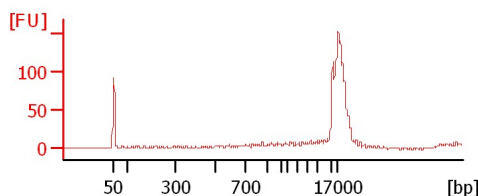
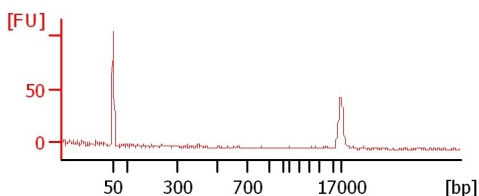
Mac11-12 DDR lib



1306C 19kbCut Lib (1:2) ?



1306C 19kbCut DDR Lib (1:2)



Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
Modified: 10/7/2015 11:22:25 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
BM PacBio cDNA Lib (1:2)		<input type="checkbox"/>	✓			
DRIP-BS PacBio amplicon lib (1:2)		<input type="checkbox"/>	✓			
BM End Repair (1:3)		<input type="checkbox"/>	✓			
DRIP-BS End Repair (1:3)		<input type="checkbox"/>	✓			
Swanson DDR lib		<input type="checkbox"/>	✓			
		<input type="checkbox"/>	✓			
Mac11-12 DDR lib		<input type="checkbox"/>	✓			
		<input type="checkbox"/>	✓			
1306C 19kbCut Lib (1:2) ?		<input type="checkbox"/>	✓			
		<input type="checkbox"/>	✓			
1306C 19kbCut DDR Lib (1:2)		<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\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
Modified: 10/7/2015 11:22:25 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

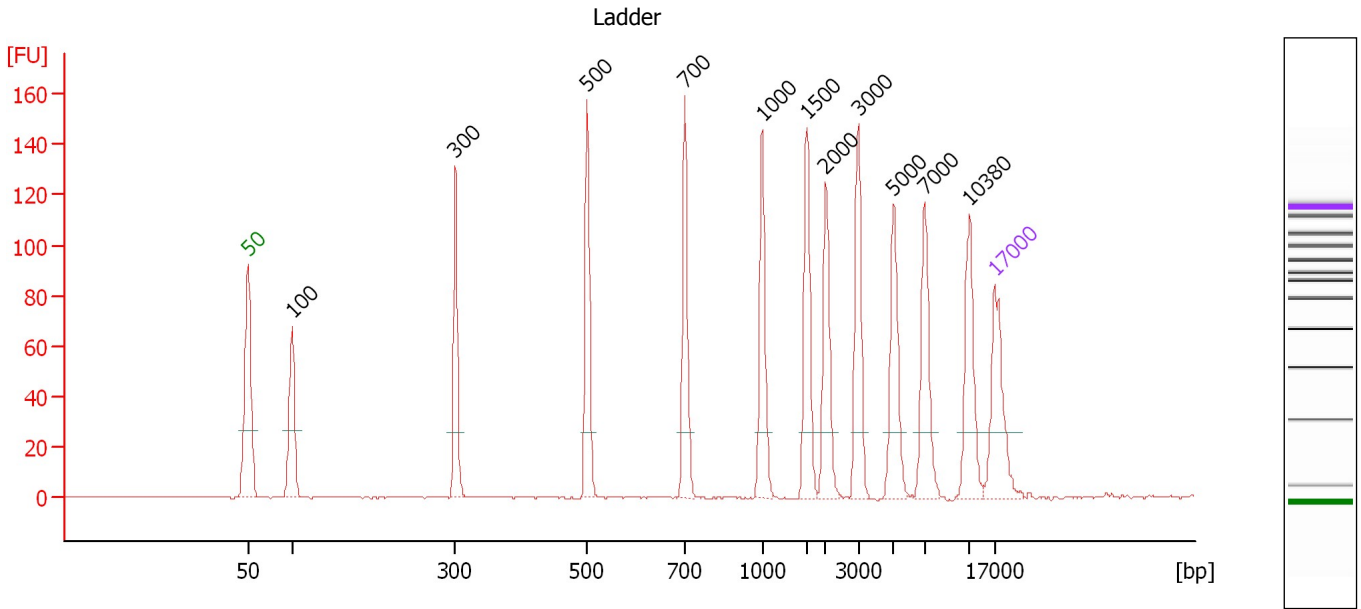
Ladder

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\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
 Modified: 10/7/2015 11:22:25 AM

Electropherogram Summary



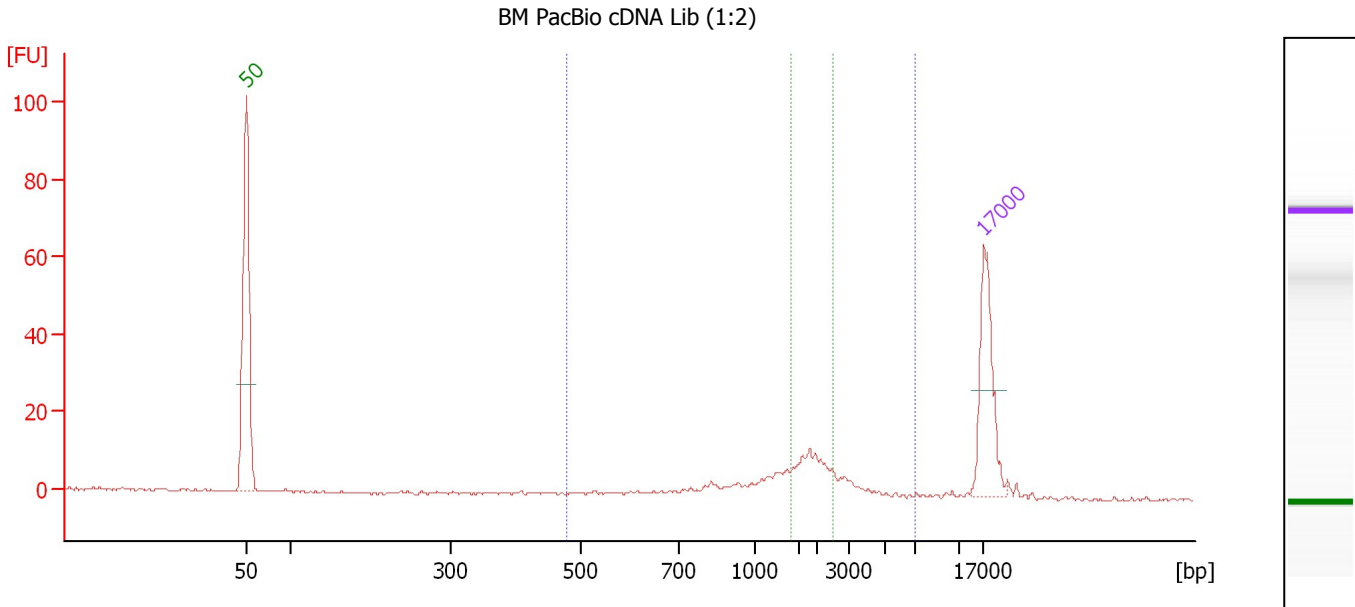
Peak table for Ladder

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\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
 Modified: 10/7/2015 11:22:25 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : BM PacBio cDNA Lib (1:2)

Number of peaks found: 0 Area 2: 26.7
 Area 1: 68.0

Peak table for sample 1 : BM PacBio cDNA 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

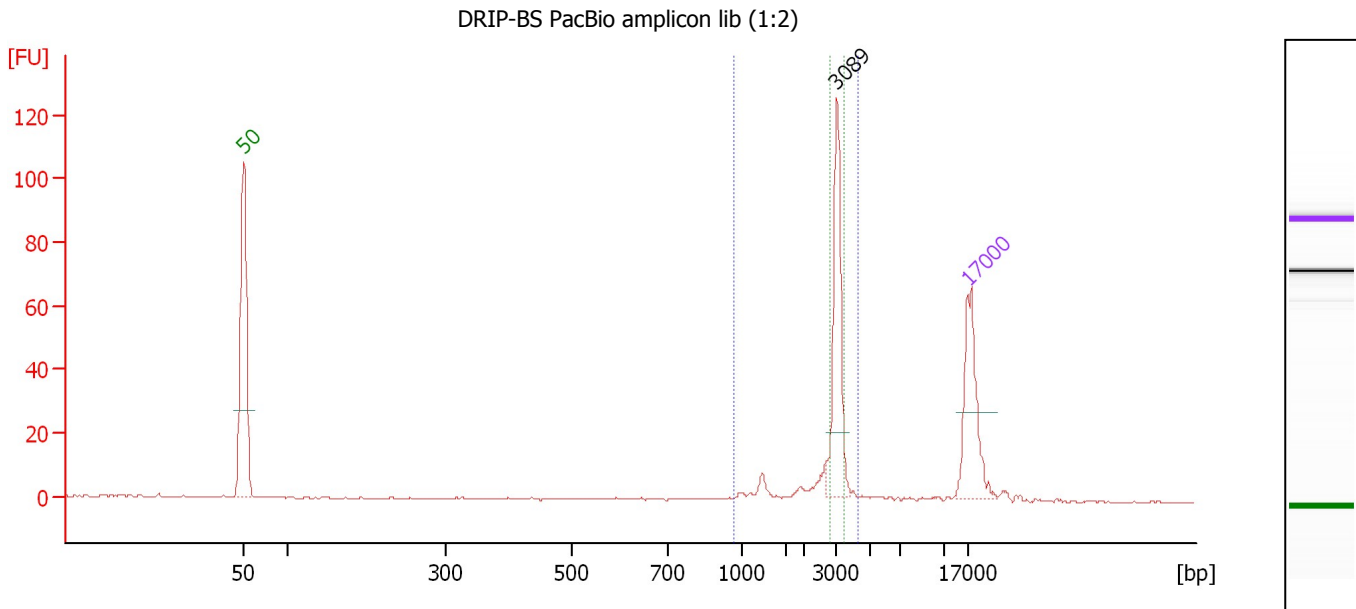
Region table for sample 1 : BM PacBio cDNA Lib (1:2)

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
476	7,022	1,775	5.11	68.0	85	57.0	Blue
1,426	2,514	1,876	1.95	26.7	34	16.4	Green

Assay Class: DNA 12000
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
 Modified: 10/7/2015 11:22:25 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 : DRIP-BS PacBio amplicon lib (1:2)

Number of peaks found: 1 Area 2: 72.1
 Area 1: 101.8

Peak table for sample 2 : DRIP-BS PacBio amplicon lib (1:2)

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	3,089	5.36	2.6	
3	17,000	4.20	0.4	Upper Marker

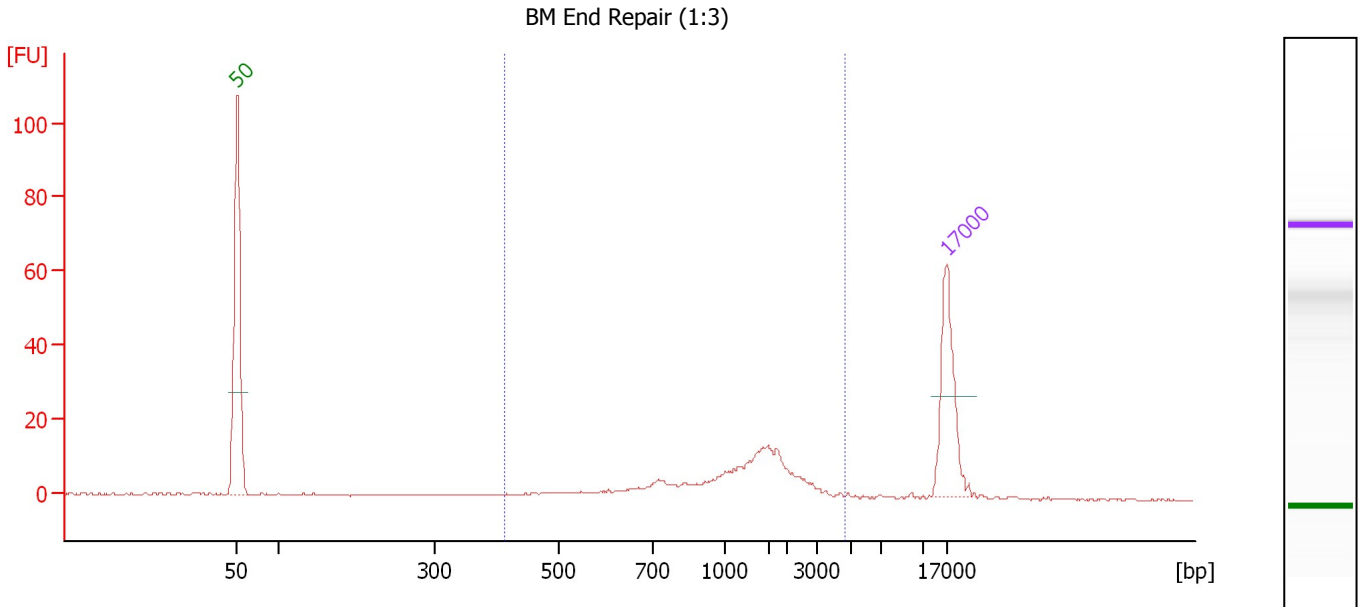
Region table for sample 2 : DRIP-BS PacBio amplicon lib (1:2)

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
968	4,298	2,861	7.16	101.8	84	22.2	Blue
2,826	3,492	3,110	5.01	72.1	59	5.0	Green

Assay Class: DNA 12000
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
 Modified: 10/7/2015 11:22:25 AM

Electropherogram Summary Continued ...



Overall Results for sample 3 : BM End Repair (1:3)

Number of peaks found: 0 Area 1: 93.9

Peak table for sample 3 : BM End Repair (1:3)

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

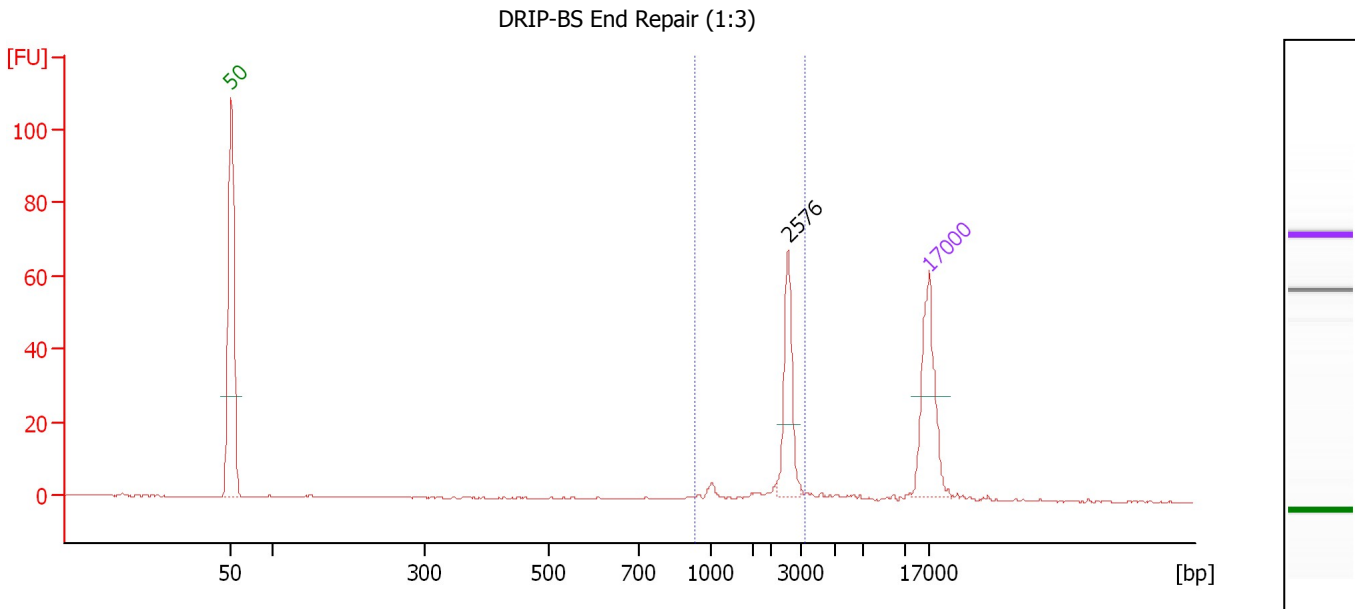
Region table for sample 3 : BM End Repair (1:3)

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
414	4,664	1,359	7.22	93.9	84	51.9	Blue

Assay Class: DNA 12000
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
 Modified: 10/7/2015 11:22:25 AM

Electropherogram Summary Continued ...



Overall Results for sample 4 : DRIP-BS End Repair (1:3)

Number of peaks found: 1 Area 1: 55.4

Peak table for sample 4 : DRIP-BS End Repair (1:3)

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	2,576	3.29	1.9	
3	17,000	4.20	0.4	Upper Marker

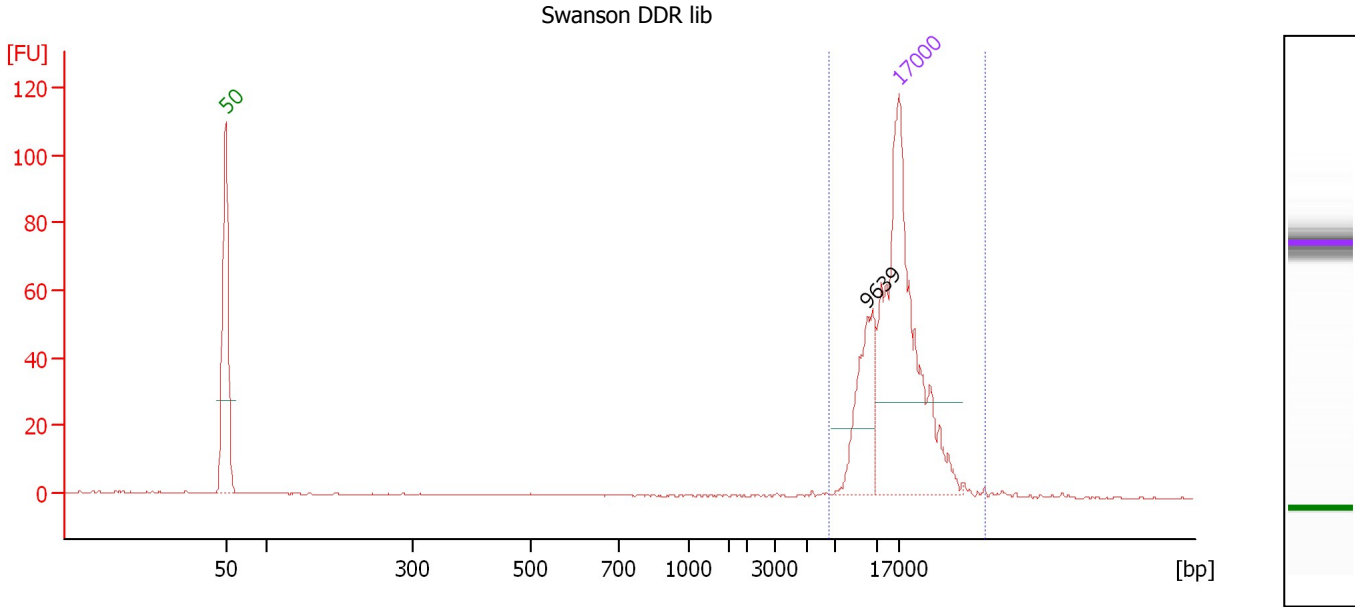
Region table for sample 4 : DRIP-BS End Repair (1:3)

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
928	3,297	2,406	4.16	55.4	74	20.0	Blue

Assay Class: DNA 12000
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
 Modified: 10/7/2015 11:22:25 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : Swanson DDR lib

Number of peaks found: 1 Area 1: 79.2

Peak table for sample 5 : Swanson DDR lib

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	9,639	1.24	0.2	
3	17,000	4.20	0.4	Upper Marker

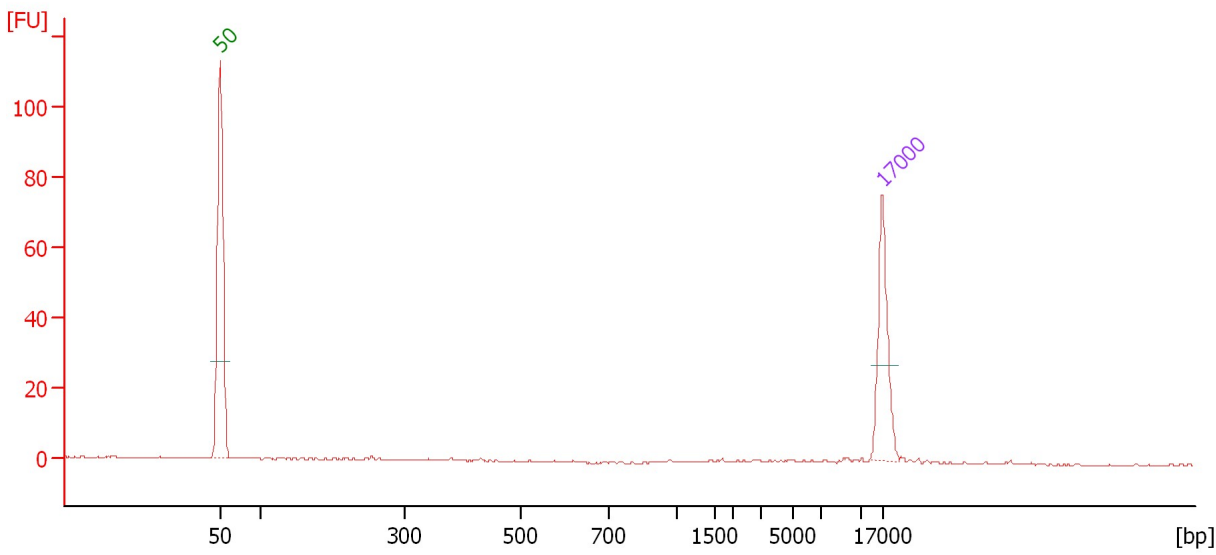
Region table for sample 5 : Swanson DDR lib

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
6,515	42,228	16,721	1.28	79.2	88	38.4	Blue

Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
Modified: 10/7/2015 11:22:25 AM

Electropherogram Summary Continued ...



Overall Results for sample 6 :

Number of peaks found: 0

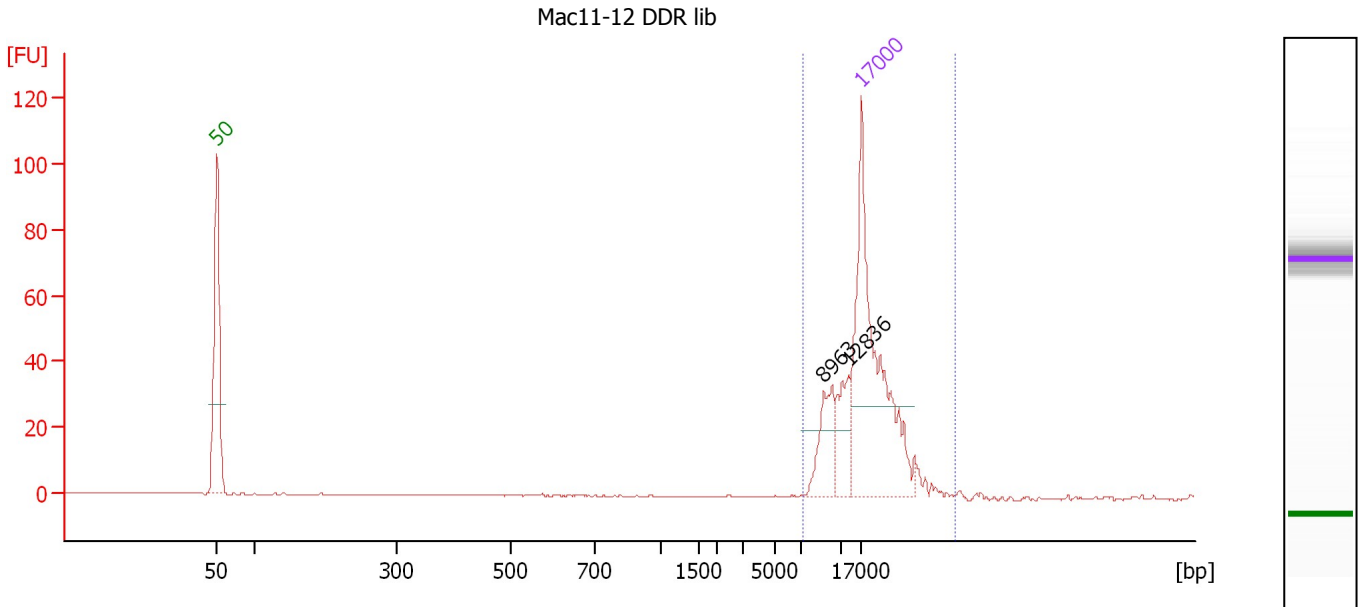
Peak table for sample 6 :

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\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
 Modified: 10/7/2015 11:22:25 AM

Electropherogram Summary Continued ...



Overall Results for sample 7 : Mac11-12 DDR lib

Number of peaks found: 2 Area 1: 78.4

Peak table for sample 7 : Mac11-12 DDR lib

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	8,963	0.90	0.2	
3	12,836	0.82	0.1	
4	17,000	4.20	0.4	Upper Marker

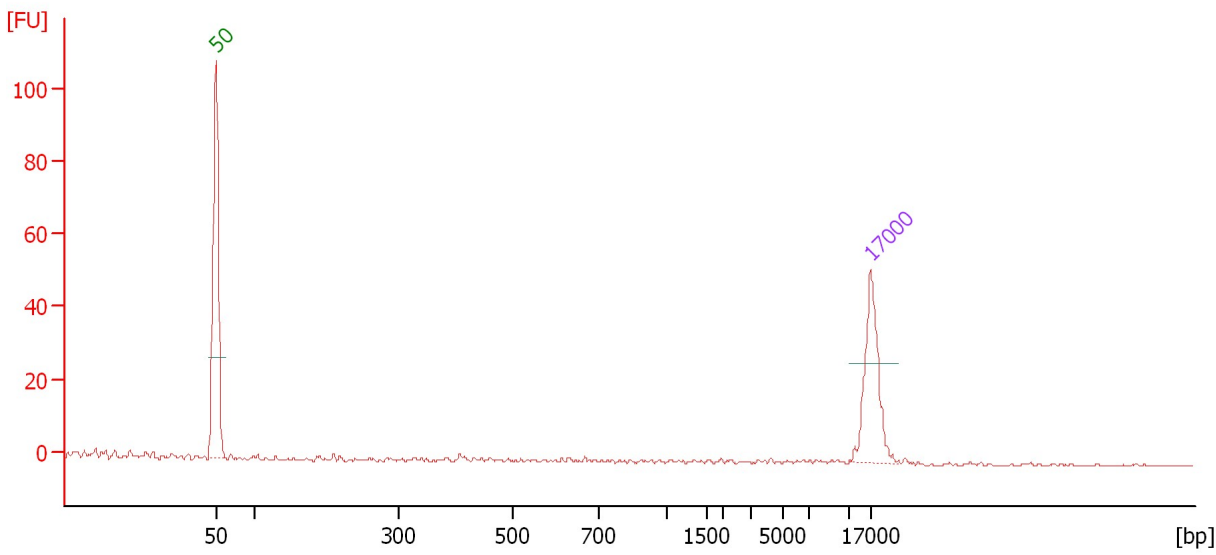
Region table for sample 7 : Mac11-12 DDR lib

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
7,190	45,476	18,068	1.79	78.4	99	37.5	Blue

Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
Modified: 10/7/2015 11:22:25 AM

Electropherogram Summary Continued ...



Overall Results for sample 8 :

Number of peaks found: 0

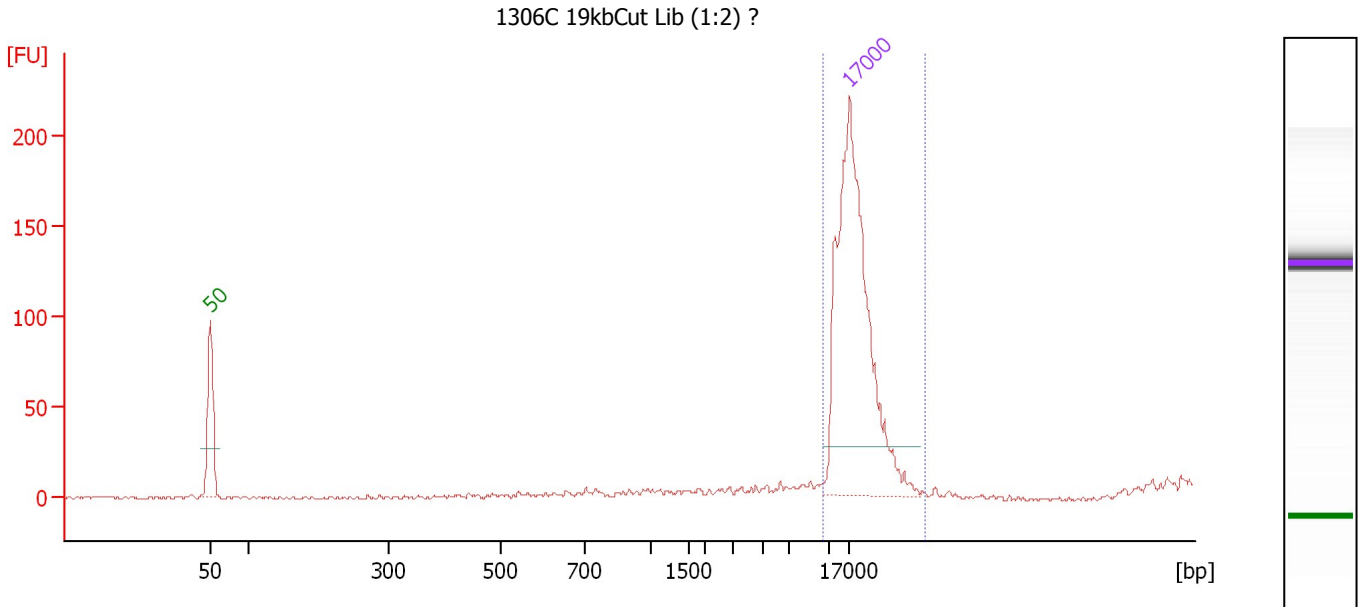
Peak table for sample 8 :

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\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
 Modified: 10/7/2015 11:22:25 AM

Electropherogram Summary Continued ...



Overall Results for sample 9 : 1306C 19kbCut Lib (1:2) ?

Number of peaks found: 0 Area 1: 0.0

Peak table for sample 9 : 1306C 19kbCut 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

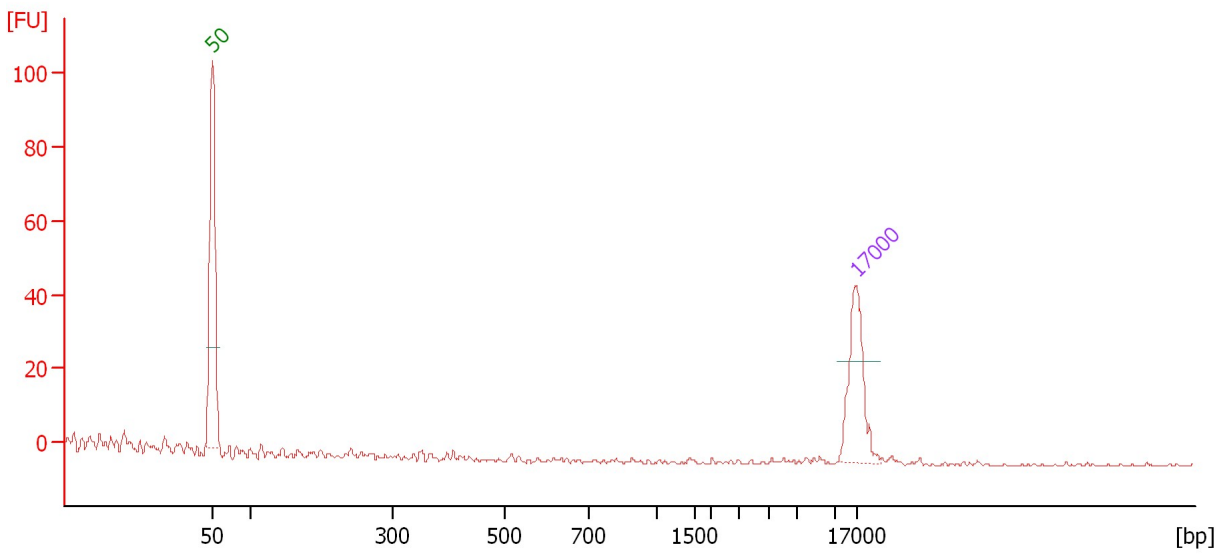
Region table for sample 9 : 1306C 19kbCut Lib (1:2) ?

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
9,984	40,172	18,563	0.00	0.0	0	25.7	Blue

Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
Modified: 10/7/2015 11:22:25 AM

Electropherogram Summary Continued ...



Overall Results for sample 10 :

Number of peaks found: 0

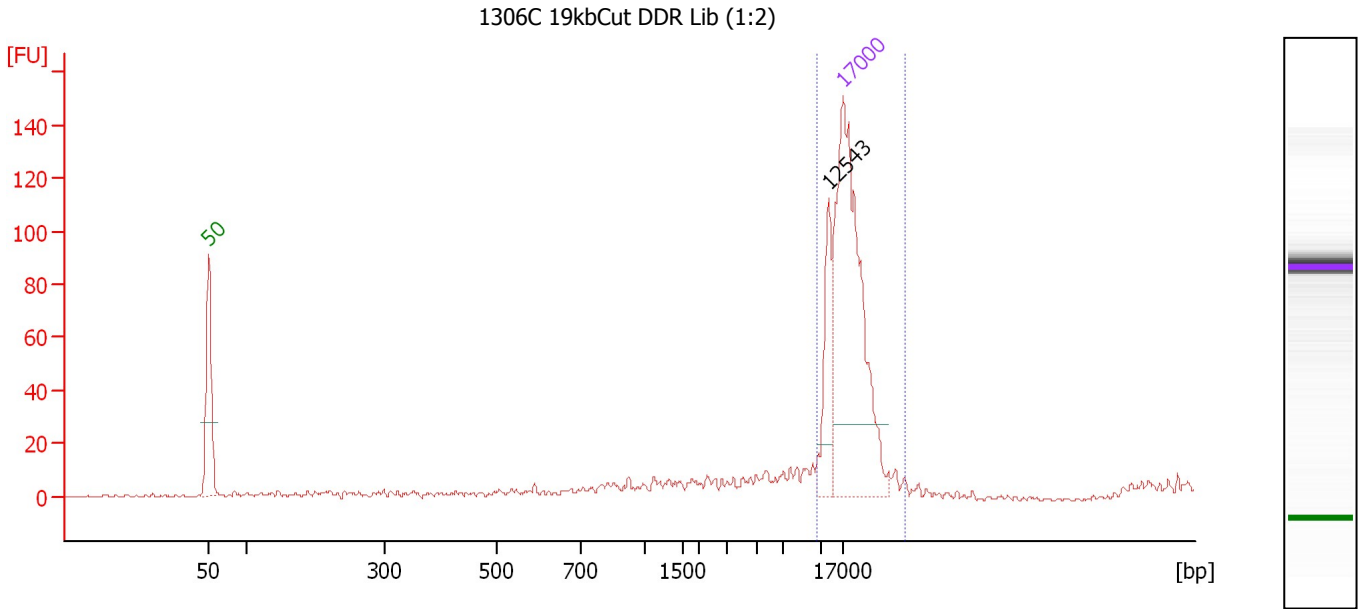
Peak table for sample 10 :

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\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
 Modified: 10/7/2015 11:22:25 AM

Electropherogram Summary Continued ...



Overall Results for sample 11 : 1306C 19kbCut DDR Lib (1:2)

Number of peaks found: 1 Area 1: 72.4

Peak table for sample 11 : 1306C 19kbCut DDR Lib (1:2)

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	12,543	1.02	0.1	
3	17,000	4.20	0.4	Upper Marker

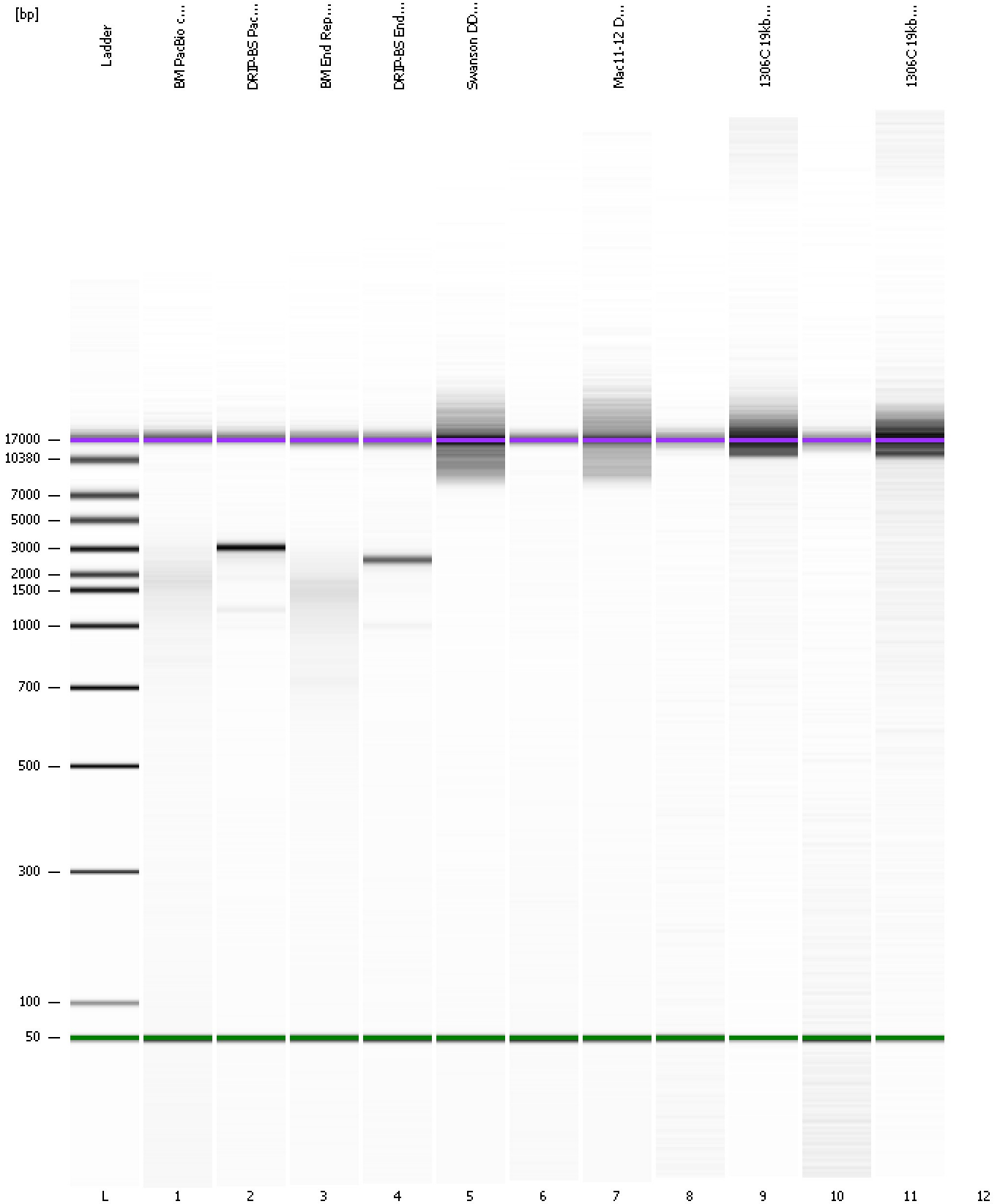
Region table for sample 11 : 1306C 19kbCut DDR Lib (1:2)

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
9,913	36,512	18,441	1.02	72.4	43	26.4	Blue

Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
Modified: 10/7/2015 11:22:25 AM

Gel Image

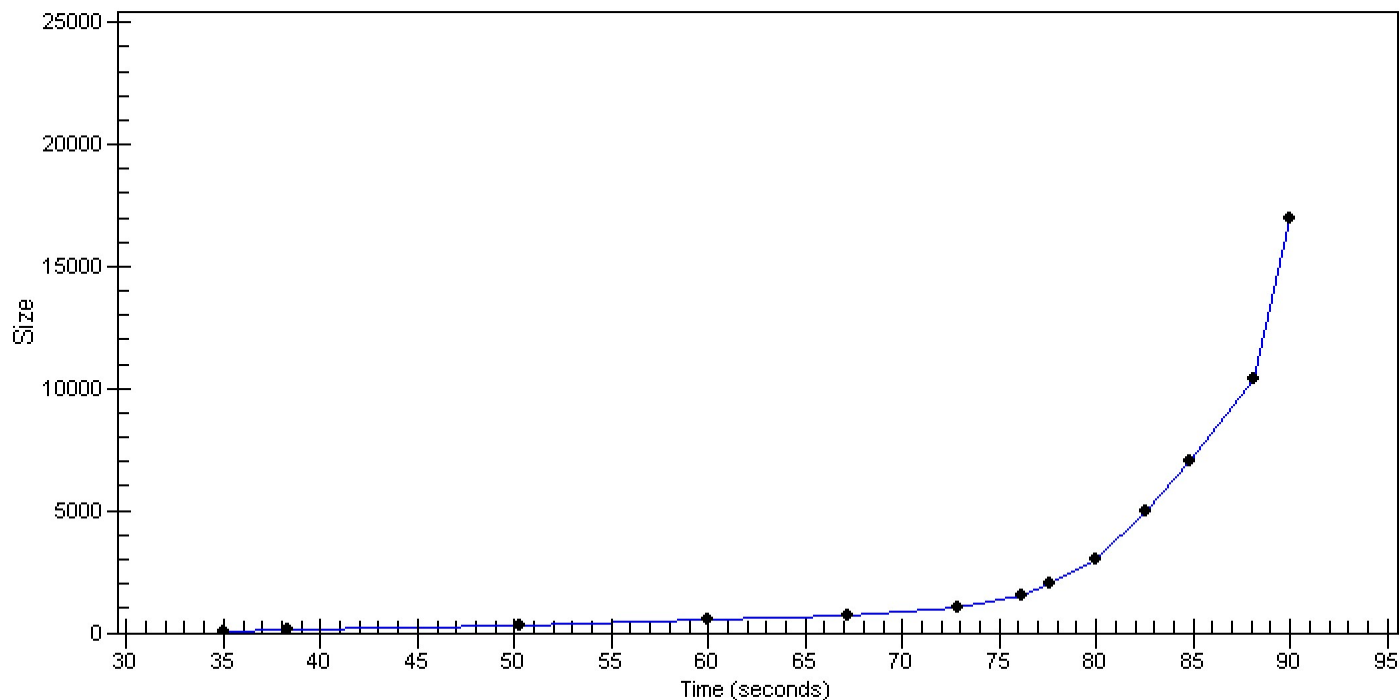


Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
Modified: 10/7/2015 11:22:25 AM

Curves

Standard Curve



Assay Class: DNA 12000
Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
Modified: 10/7/2015 11:22:25 AM

Invalid Samples

Sample 12 has not been run, no results available.

Assay Class: DNA 12000
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad

Created: 10/7/2015 10:38:18 AM
 Modified: 10/7/2015 11:22:25 AM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 12)		Instrument	Run		10/7/2015 11:11:13 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-10-07\2015-10-07_001.xad)		Instrument	Run		10/7/2015 10:38:23 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		10/7/2015 10:38:23 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		10/7/2015 10:38:23 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		10/7/2015 10:38:23 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		10/7/2015 10:38:23 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		10/7/2015 10:38:23 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		10/7/2015 10:38:23 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1