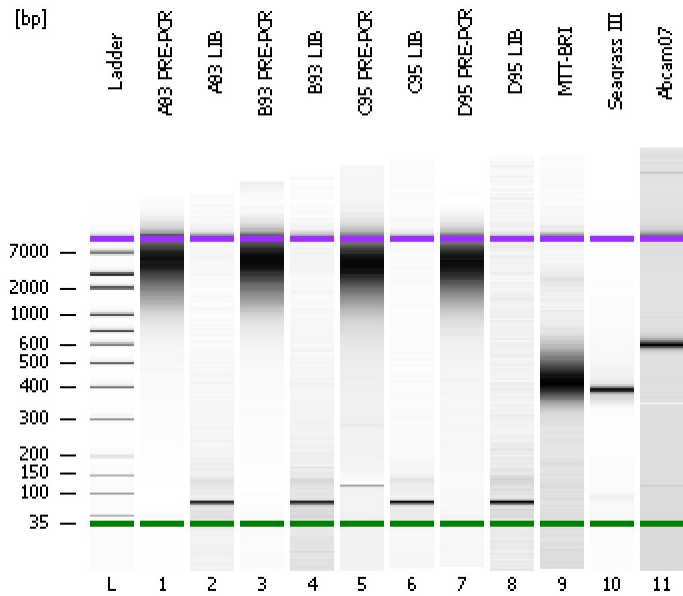


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
Modified: 2/11/2015 3:38:34 PM

**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\High Sensitivity DNA.xsy  
Assay Class: High Sensitivity DNA Assay  
Version: 1.03  
Assay Comments: Copyright © 2003-2010 Agilent Technologies

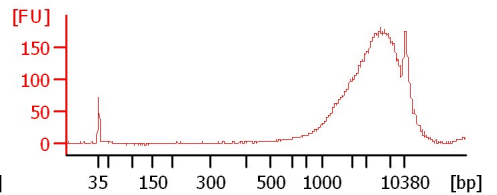
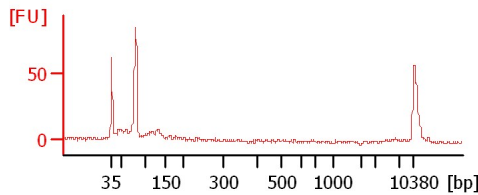
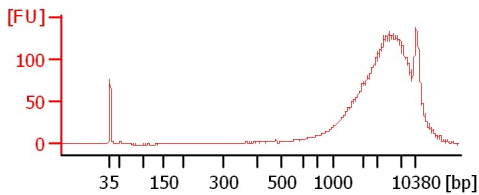
Chip Information:

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

**A93 PRE-PCR**

**A93 LIB**

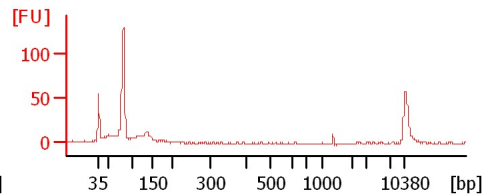
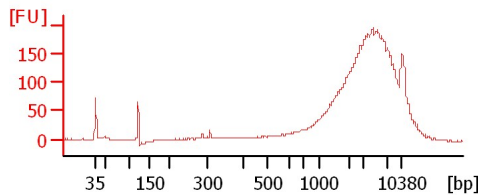
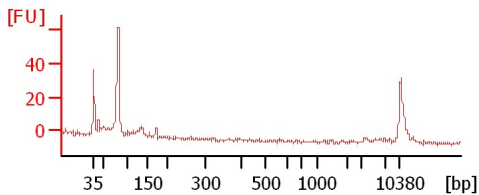
**B93 PRE-PCR**



**B93 LIB**

**C95 PRE-PCR**

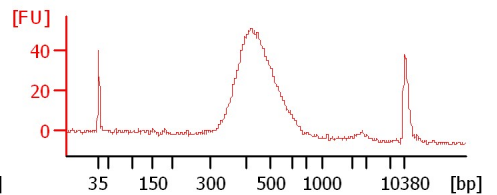
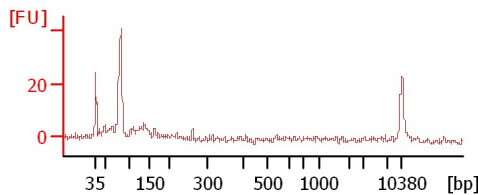
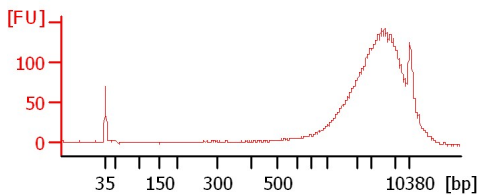
**C95 LIB**



**D95 PRE-PCR**

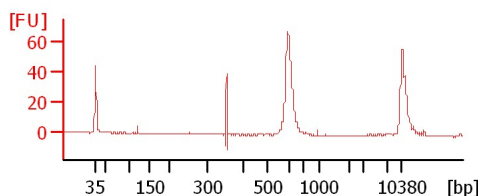
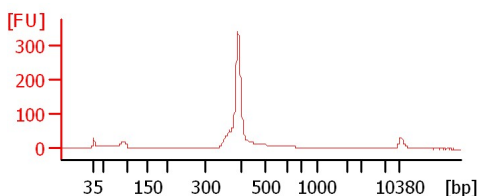
**D95 LIB**

**MTT-BRI**



**Seagrass III**

**Abcam07**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
A93 PRE-PCR		<input type="checkbox"/>	✓			
A93 LIB		<input type="checkbox"/>	✓			
B93 PRE-PCR		<input type="checkbox"/>	✓			
B93 LIB		<input type="checkbox"/>	✓			
C95 PRE-PCR		<input type="checkbox"/>	✓			
C95 LIB		<input type="checkbox"/>	✓			
D95 PRE-PCR		<input type="checkbox"/>	✓			
D95 LIB		<input type="checkbox"/>	✓			
MTT-BRI		<input type="checkbox"/>	✓			
Seagrass III		<input type="checkbox"/>	✓			
Abcam07		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

**Chip Lot #**

**Reagent Kit Lot #**

**Chip Comments :**

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
Modified: 2/11/2015 3:38:34 PM

**Electrophoresis Assay Details**

**General Analysis Settings**

Number of Available Sample and Ladder Wells (Max.) : 12  
Minimum Visible Range [s] : 32  
Maximum Visible Range [s] : 138  
Start Analysis Time Range [s] : 33  
End Analysis Time Range [s] : 137.5  
Ladder Concentration [pg/μl] : 1950  
Uses Standard Area for Ladder Fragments  
Lower Marker Concentration [pg/μl] : 125  
Upper Marker Concentration [pg/μl] : 75  
Used Upper Marker for Quantitation  
Standard Curve Fit is Point to Point  
Show Data Aligned to Lower and Upper Marker

**Integrator Settings**

Integration Start Time [s] : 33.05  
Integration End Time [s] : 137  
Slope Threshold : 0.8  
Height Threshold [FU] : 5  
Area Threshold : 0.1  
Width Threshold [s] : 0.6  
Baseline Plateau [s] : 0.5

**Filter Settings**

Filter Width [s] : 0.5  
Polynomial Order : 4

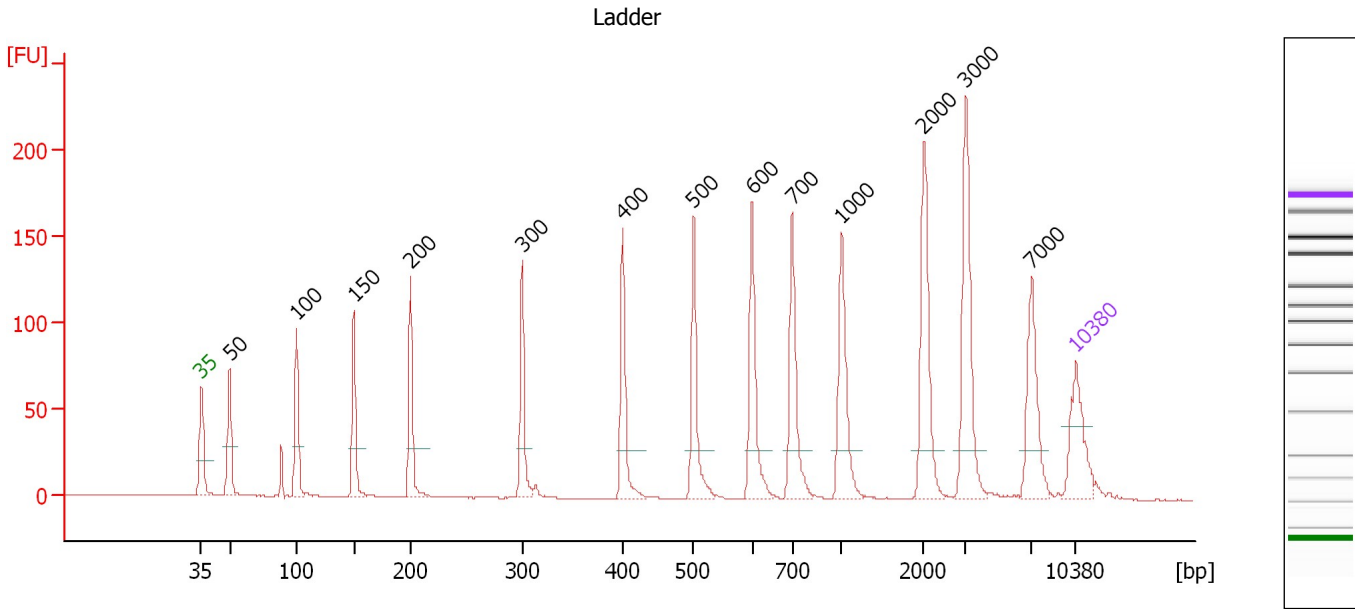
**Ladder**

Ladder Peak	Size	Area
1	35	160
2	50	210
3	100	208
4	150	221
5	200	242
6	300	270
7	400	305
8	500	306
9	600	336
10	700	321
11	1000	366
12	2000	413
13	3000	411
14	7000	400
15	10380	214

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.2

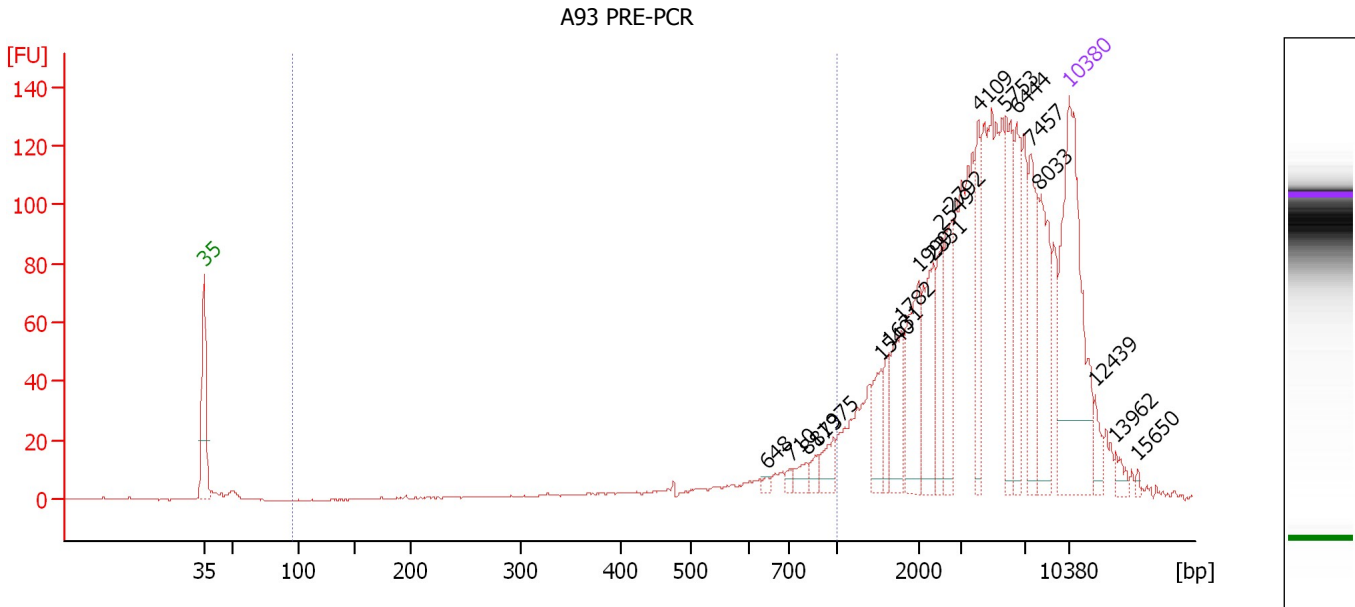
**Peak table for Ladder**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	50	150.00	4,545.5	Ladder Peak	45.26
3	100	150.00	2,272.7	Ladder Peak	50.64
4	150	150.00	1,515.2	Ladder Peak	55.21
5	200	150.00	1,136.4	Ladder Peak	59.73
6	300	150.00	757.6	Ladder Peak	68.70
7	400	150.00	568.2	Ladder Peak	76.72
8	500	150.00	454.5	Ladder Peak	82.40
9	600	150.00	378.8	Ladder Peak	87.09
10	700	150.00	324.7	Ladder Peak	90.29
11	1,000	150.00	227.3	Ladder Peak	94.26
12	2,000	150.00	113.6	Ladder Peak	100.84
13	3,000	150.00	75.8	Ladder Peak	104.21
14	7,000	150.00	32.5	Ladder Peak	109.46
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : A93 PRE-PCR**

Number of peaks found: 20                      Corr. Area 1: 153.8  
 Noise: 0.2

**Peak table for sample 1 : A93 PRE-PCR**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	648	1.69	4.0		88.62
3	710	2.08	4.4		90.43
4	811	5.05	9.4		91.77
5	873	4.24	7.4		92.59
6	975	9.36	14.6		93.92
7	1,540	13.69	13.5		97.81
8	1,631	9.46	8.8		98.41
9	1,782	20.87	17.7		99.41
10	1,999	31.65	24.0		100.83
11	2,331	27.29	17.7		101.95
12	2,549	16.98	10.1		102.69
13	2,792	24.94	13.5		103.51
14	4,109	22.86	8.4		105.66
15	5,753	30.54	8.0		107.82
16	6,444	26.06	6.1		108.73
17	7,457	28.79	5.8		109.94
18	8,033	32.54	6.1		110.54
19	10,380	75.00	10.9	Upper Marker	113.00
20	12,439	0.00	0.0		115.16
21	13,962	0.00	0.0		116.75
22	15,650	0.00	0.0		118.52


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary Continued ...**

... Region table for sample 1 :

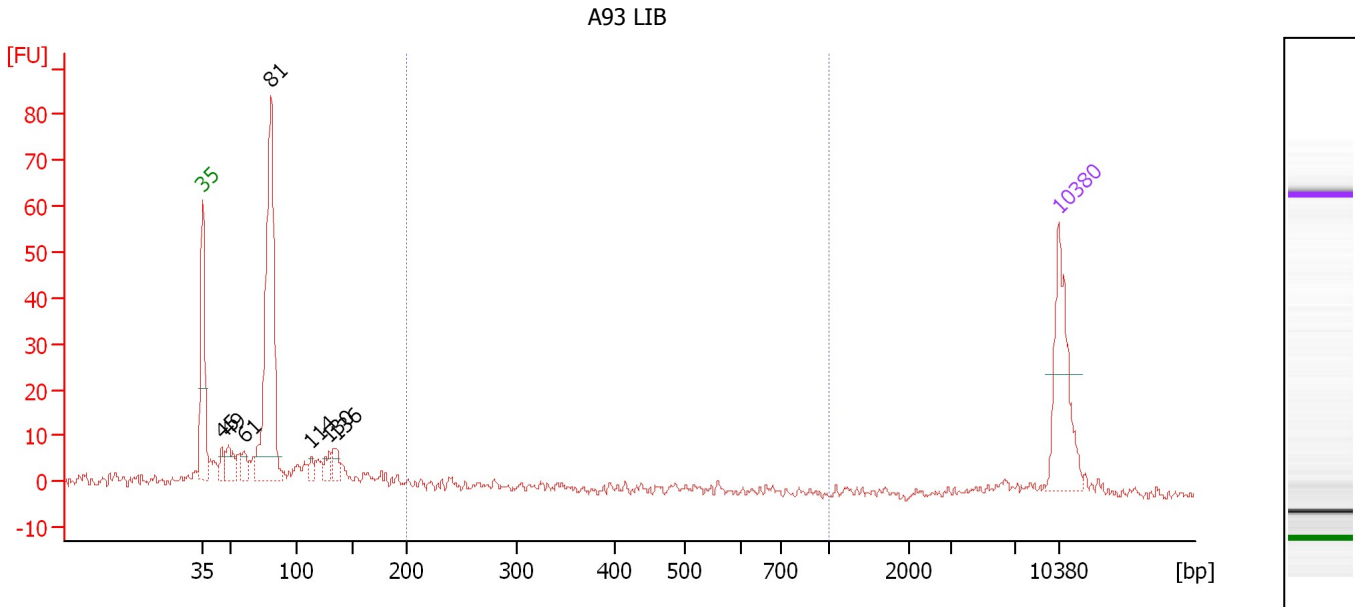
**A93 PRE-PCR**

<b>From [bp]</b>	<b>Average Size [bp]</b>	<b>To [bp]</b>	<b>Corr. Area</b>	<b>% of Total</b>	<b>Size distribution in CV [%]</b>	<b>Conc. [pg/<math>\mu</math>l]</b>	<b>Molarity [pmol/l]</b>	<b>Co lor</b>
95	672	1,000	153.8	9	31.0	66.27	182.6	

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : A93 LIB**

Number of peaks found: 7                      Corr. Area 1: 25.2  
 Noise: 1.1

**Peak table for sample 2 : A93 LIB**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	45	17.10	570.6		44.57
3	49	30.84	950.4		45.14
4	61	18.94	473.5		46.40
5	81	340.24	6,342.8		48.63
6	114	11.44	152.4		51.90
7	130	17.19	200.3		53.38
8	136	19.97	222.1		53.95
9	10,380	75.00	10.9	Upper Marker	113.00

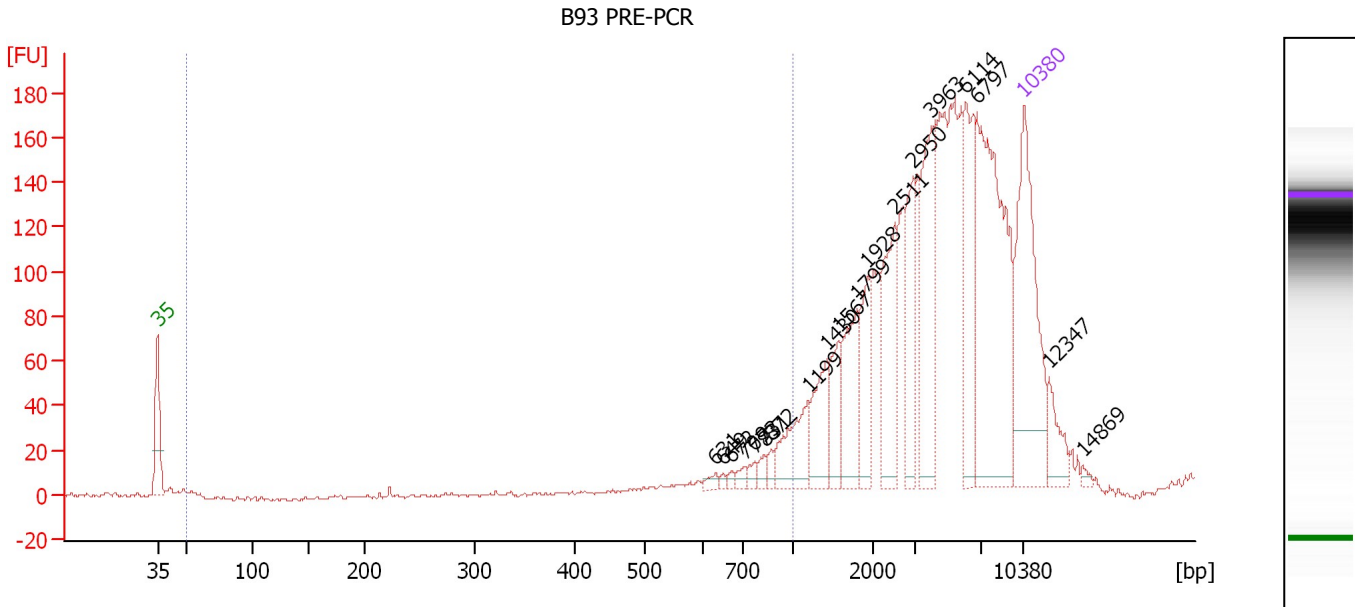
**Region table for sample 2 : A93 LIB**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	374	1,000	25.2	7	40.3	43.69	216.7	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : B93 PRE-PCR**

Number of peaks found: 19                      Corr. Area 1: 83.7  
 Noise: 0.7

**Peak table for sample 3 : B93 PRE-PCR**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	631	3.13	7.5		88.08
3	649	1.64	3.8		88.67
4	672	2.09	4.7		89.38
5	709	3.64	7.8		90.41
6	773	3.72	7.3		91.27
7	831	4.43	8.1		92.03
8	872	3.76	6.5		92.56
9	1,199	28.05	35.5		95.57
10	1,430	26.23	27.8		97.09
11	1,567	16.99	16.4		97.99
12	1,799	32.15	27.1		99.51
13	1,928	26.00	20.4		100.36
14	2,511	40.71	24.6		102.56
15	2,950	30.87	15.9		104.04
16	3,963	55.03	21.0		105.47
17	6,114	42.52	10.5		108.29
18	6,797	120.65	26.9		109.19
19	10,380	75.00	10.9	Upper Marker	113.00
20	12,347	0.00	0.0		115.06
21	14,869	0.00	0.0		117.71

**Region table for sample 3 : B93 PRE-PCR**

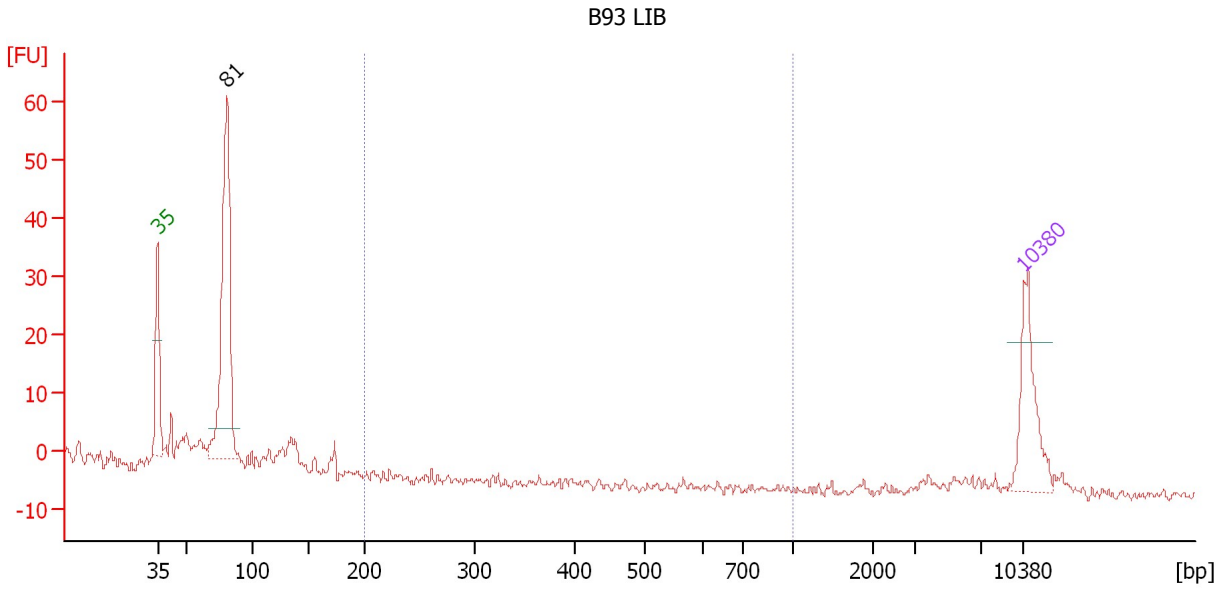
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
50	827	1,000	83.7	4	15.3	29.28	56.9	Blue



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : B93 LIB**

Number of peaks found: 1                      Corr. Area 1: 0.0  
 Noise: 0.8

**Peak table for sample 4 : B93 LIB**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	81	360.75	6,752.4		48.59
3	10,380	75.00	10.9	Upper Marker	113.00

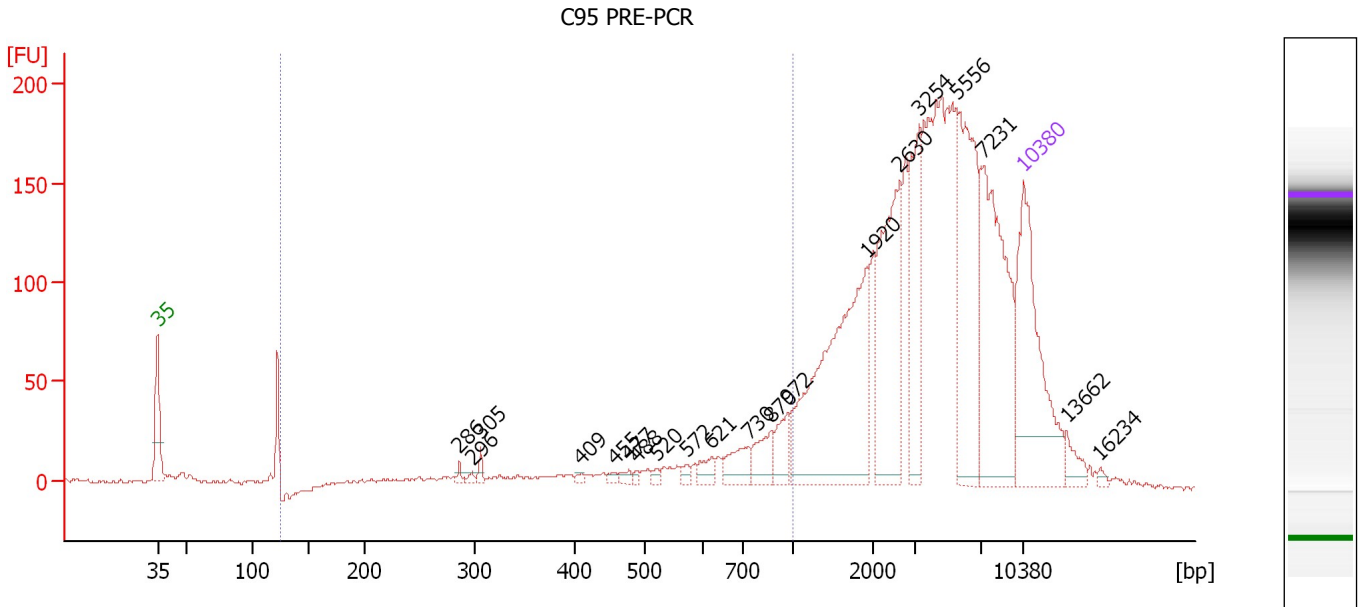
**Region table for sample 4 : B93 LIB**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	0	1,000	0.0	0	0.0	0.00	0.0	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : C95 PRE-PCR**

Number of peaks found: 20                      Corr. Area 1: 324.9  
 Noise: 0.9

**Peak table for sample 5 : C95 PRE-PCR**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	286	2.25	11.9		67.40
3	296	2.11	10.8		68.38
4	305	3.44	17.1		69.12
5	409	1.86	6.9		77.21
6	455	2.05	6.8		79.86
7	477	3.03	9.6		81.07
8	488	1.69	5.3		81.72
9	520	2.21	6.4		83.35
10	572	3.08	8.2		85.76
11	621	6.91	16.9		87.76
12	730	14.54	30.2		90.69
13	870	15.02	26.1		92.55
14	972	14.06	21.9		93.90
15	1,920	125.42	99.0		100.31
16	2,630	78.36	45.1		102.96
17	3,254	46.97	21.9		104.54
18	5,556	89.38	24.4		107.56
19	7,231	97.19	20.4		109.70
20	10,380	75.00	10.9	Upper Marker	113.00
21	13,662	0.00	0.0		116.44
22	16,234	0.00	0.0		119.14

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
Modified: 2/11/2015 3:38:34 PM

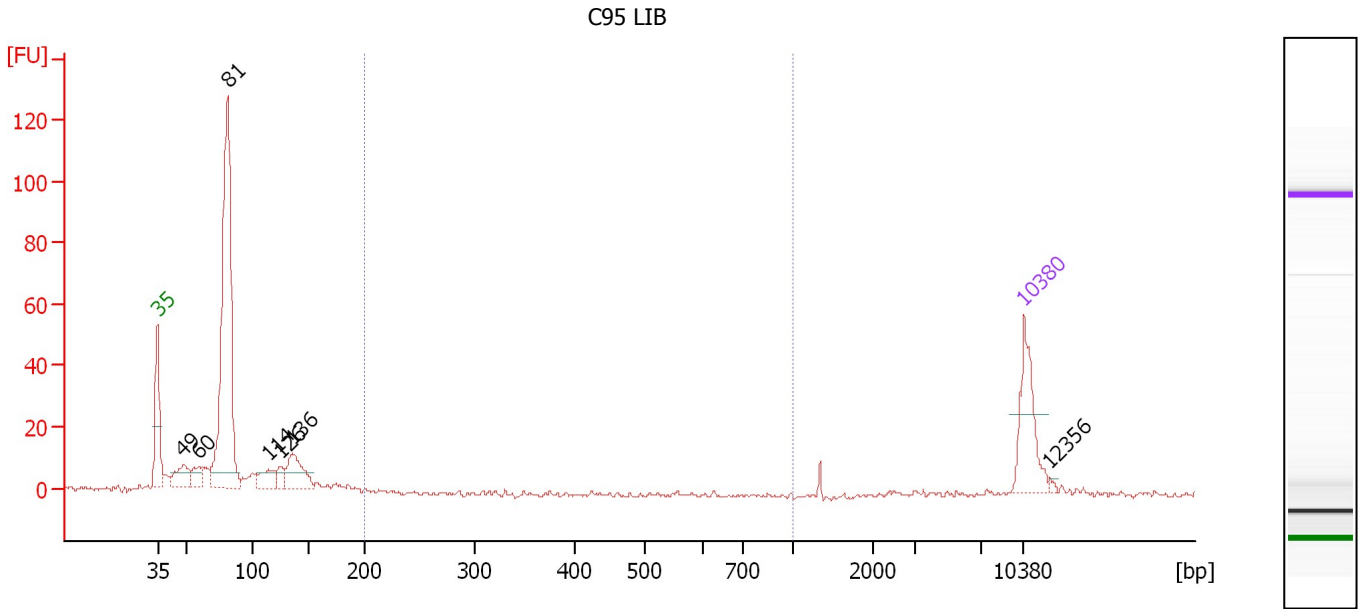
**Electropherogram Summary Continued ...****... Region table for sample 5 :      C95 PRE-PCR**

<b>From [bp]</b>	<b>Average Size [bp]</b>	<b>To [bp]</b>	<b>Corr. Area</b>	<b>% of Total</b>	<b>Size distribution in CV [%]</b>	<b>Conc. [pg/μl]</b>	<b>Molarity [pmol/l]</b>	<b>Co lor</b>
125	639	1,000	324.9	12	34.1	114.88	342.7	■

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : C95 LIB**

Number of peaks found: 7                      Corr. Area 1: 0.3  
 Noise: 0.8

**Peak table for sample 6 : C95 LIB**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	49	52.87	1,643.9		45.07
3	60	31.54	801.8		46.30
4	81	575.35	10,698.0		48.65
5	114	38.64	512.3		51.94
6	126	21.50	259.3		52.98
7	136	74.78	833.6		53.92
8	10,380	75.00	10.9	Upper Marker	113.00
9	12,356	0.00	0.0		115.07

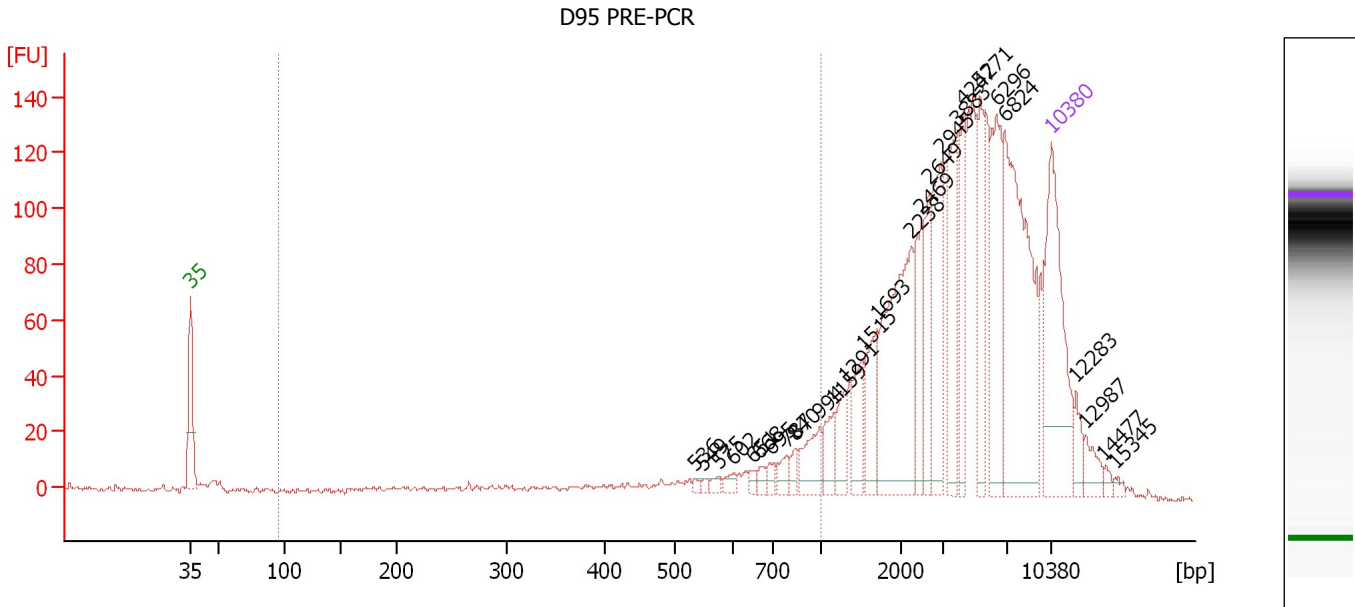
**Region table for sample 6 : C95 LIB**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	421	1,000	0.3	0	40.6	0.55	2.8	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : D95 PRE-PCR**

Number of peaks found: 27                      Corr. Area 1: 203.6  
 Noise: 0.8

**Peak table for sample 7 : D95 PRE-PCR**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	536	2.04	5.8		84.07
3	549	1.90	5.2		84.68
4	575	3.53	9.3		85.94
5	602	4.24	10.7		87.15
6	651	3.56	8.3		88.71
7	668	4.81	10.9		89.28
8	695	4.28	9.3		90.14
9	787	7.59	14.6		91.44
10	840	5.58	10.1		92.14
11	994	20.14	30.7		94.18
12	1,159	14.16	18.5		95.30
13	1,291	13.84	16.3		96.17
14	1,515	20.20	20.2		97.65
15	1,693	26.70	23.9		98.82
16	2,238	95.17	64.4		101.64
17	2,469	31.07	19.1		102.42
18	2,649	26.75	15.3		103.02
19	2,945	43.43	22.3		104.02
20	3,883	49.56	19.3		105.37
21	4,247	33.52	12.0		105.84
22	5,271	34.65	10.0		107.19
23	6,296	56.17	13.5		108.53
24	6,824	125.53	27.9		109.23
25	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad


Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

### Electropherogram Summary Continued ...

#### ... Peak table for sample 7 : D95 PRE-PCR

Peak	Size [bp]	Conc. [pg/ $\mu$ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	12,283	0.00	0.0		115.00
27	12,987	0.00	0.0		115.73
28	14,477	0.00	0.0		117.29
29	15,345	0.00	0.0		118.20

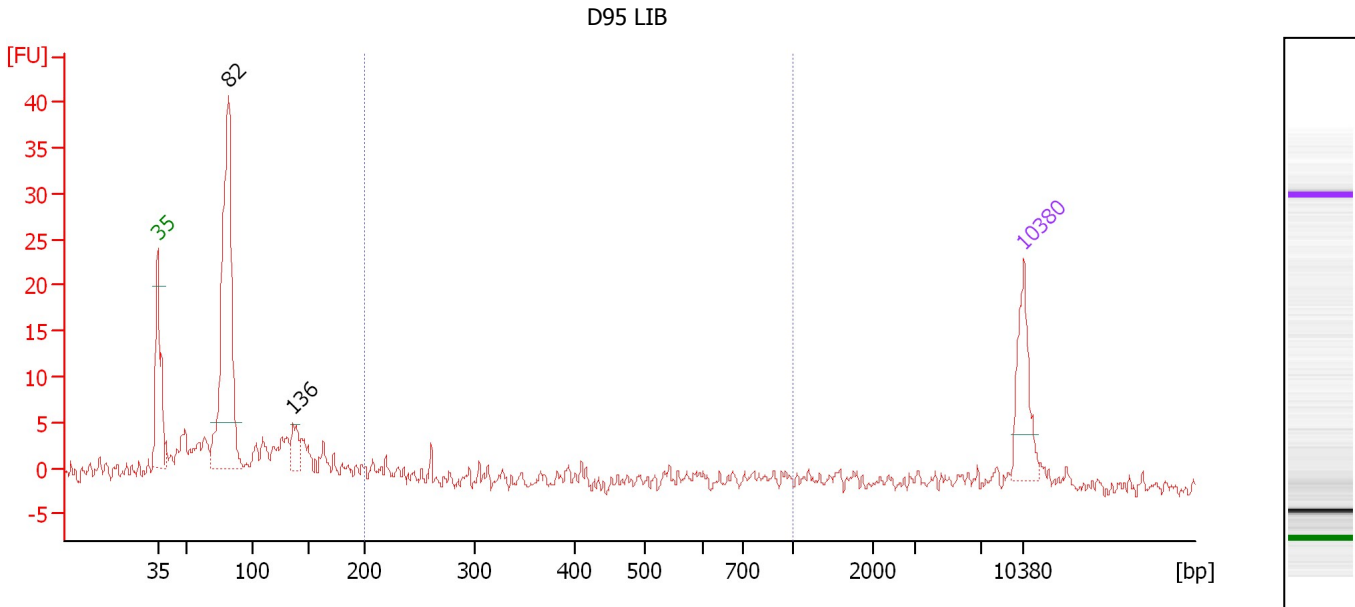
#### Region table for sample 7 : D95 PRE-PCR

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/ $\mu$ l]	Molarity [pmol/l]	Color
95	642	1,000	203.6	11	34.9	110.08	347.8	

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : D95 LIB**

Number of peaks found: 2                      Corr. Area 1: 26.2  
 Noise: 1.0

**Peak table for sample 8 : D95 LIB**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	82	541.56	9,977.6		48.73
3	136	43.98	489.8		53.93
4	10,380	75.00	10.9	Upper Marker	113.00

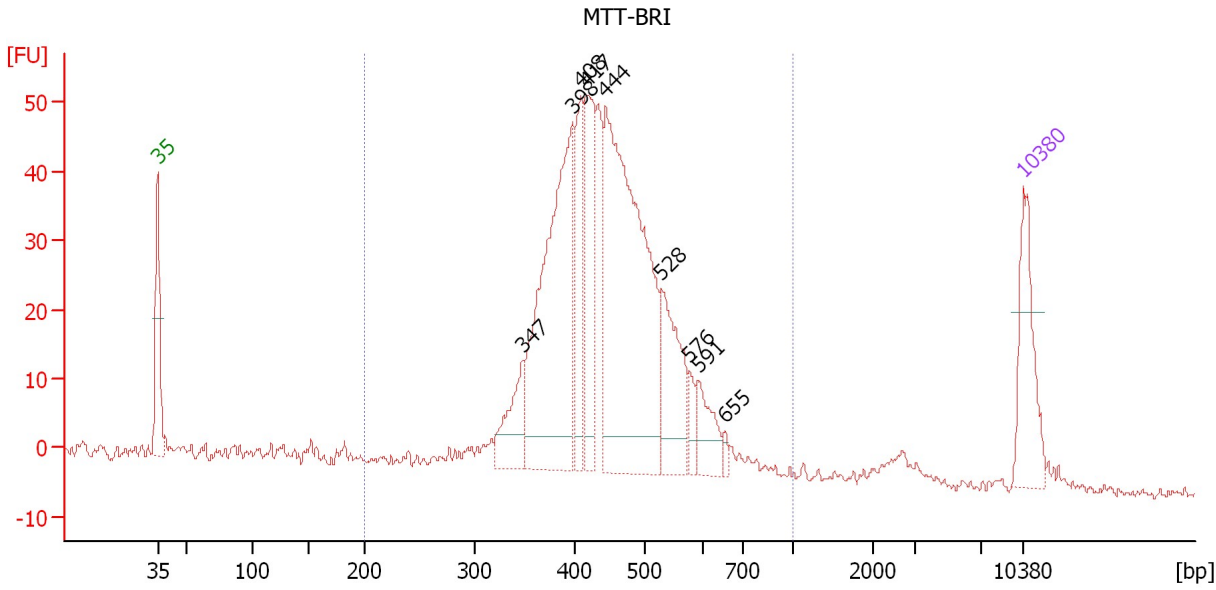
**Region table for sample 8 : D95 LIB**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	509	1,000	26.2	12	44.8	127.35	528.7	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : MTT-BRI**

Number of peaks found: 9                      Corr. Area 1: 737.9  
 Noise: 0.6

**Peak table for sample 9 : MTT-BRI**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	347	72.33	315.9		72.46
3	398	361.42	1,375.5		76.57
4	408	88.29	327.7		77.19
5	417	131.27	477.3		77.67
6	444	484.13	1,653.6		79.19
7	528	109.69	314.5		83.73
8	576	19.61	51.6		85.97
9	591	45.73	117.1		86.69
10	655	7.39	17.1		88.84
11	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 9 : MTT-BRI**

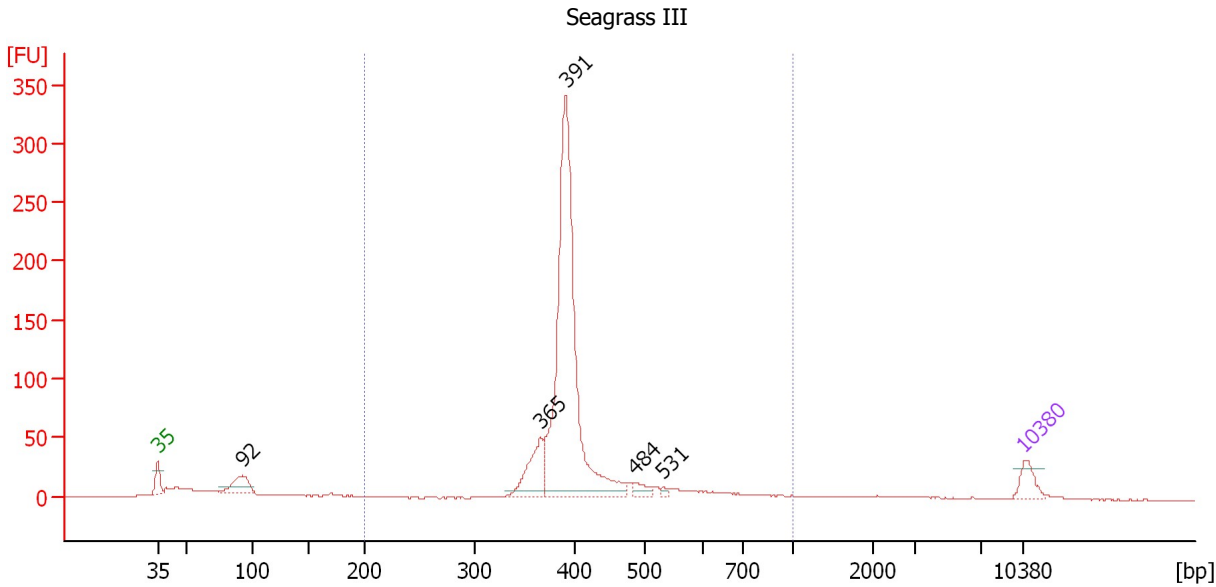
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	449	1,000	737.9	92	18.9	1,514.00	5,336.5	Blue



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : Seagrass III**

Number of peaks found: 5                      Corr. Area 1: 976.8  
 Noise: 0.2

**Peak table for sample 10 : Seagrass III**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	92	174.57	2,860.3		49.83
3	365	282.04	1,171.0		73.91
4	391	2,076.82	8,049.8		75.99
5	484	59.91	187.4		81.51
6	531	16.29	46.5		83.84
7	10,380	75.00	10.9	Upper Marker	113.00

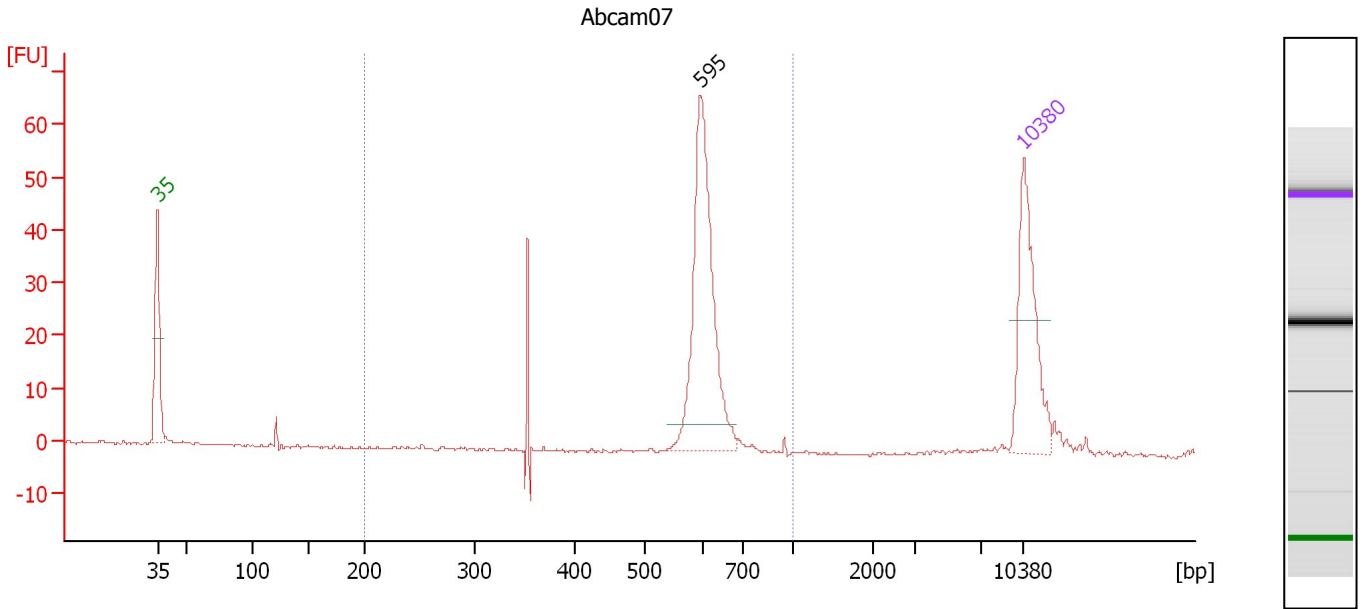
**Region table for sample 10 : Seagrass III**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	415	1,000	976.8	82	19.9	2,689.95	10,146.2	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : Abcam07**

Number of peaks found: 1                      Corr. Area 1: 150.9  
 Noise: 0.3

**Peak table for sample 11 : Abcam07**

Pea k	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	595	178.61	455.0		86.85
3	10,380	75.00	10.9	Upper Marker	113.00

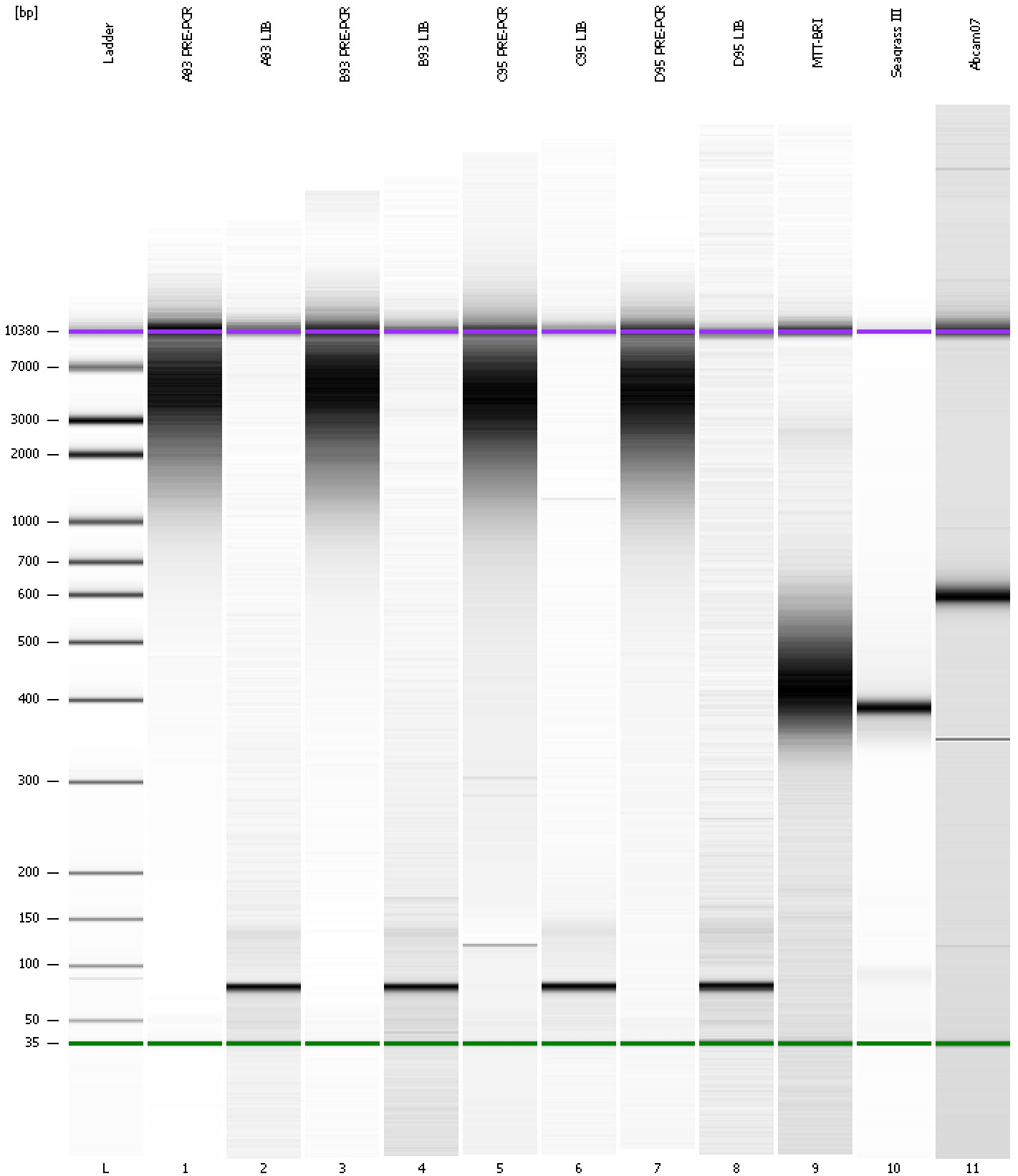
**Region table for sample 11 : Abcam07**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	592	1,000	150.9	87	11.7	195.77	517.6	Blue

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
Modified: 2/11/2015 3:38:34 PM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-11\2015-02-11\_002.xad

Created: 2/11/2015 2:57:10 PM  
 Modified: 2/11/2015 3:38:34 PM

**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		2/11/2015 3:38:29 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-02-11\2015-02-11_002.xad)		Instrument	Run		2/11/2015 2:57:15 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		2/11/2015 2:57:15 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		2/11/2015 2:57:15 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		2/11/2015 2:57:15 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		2/11/2015 2:57:15 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		2/11/2015 2:57:15 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		2/11/2015 2:57:15 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1