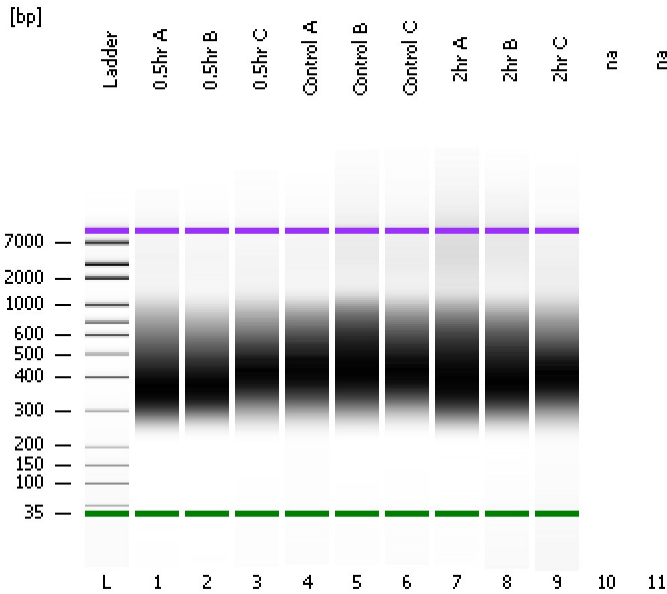


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
Modified: 3/2/2017 3:38:29 PM

### Electrophoresis File Run Summary



#### Instrument Information:

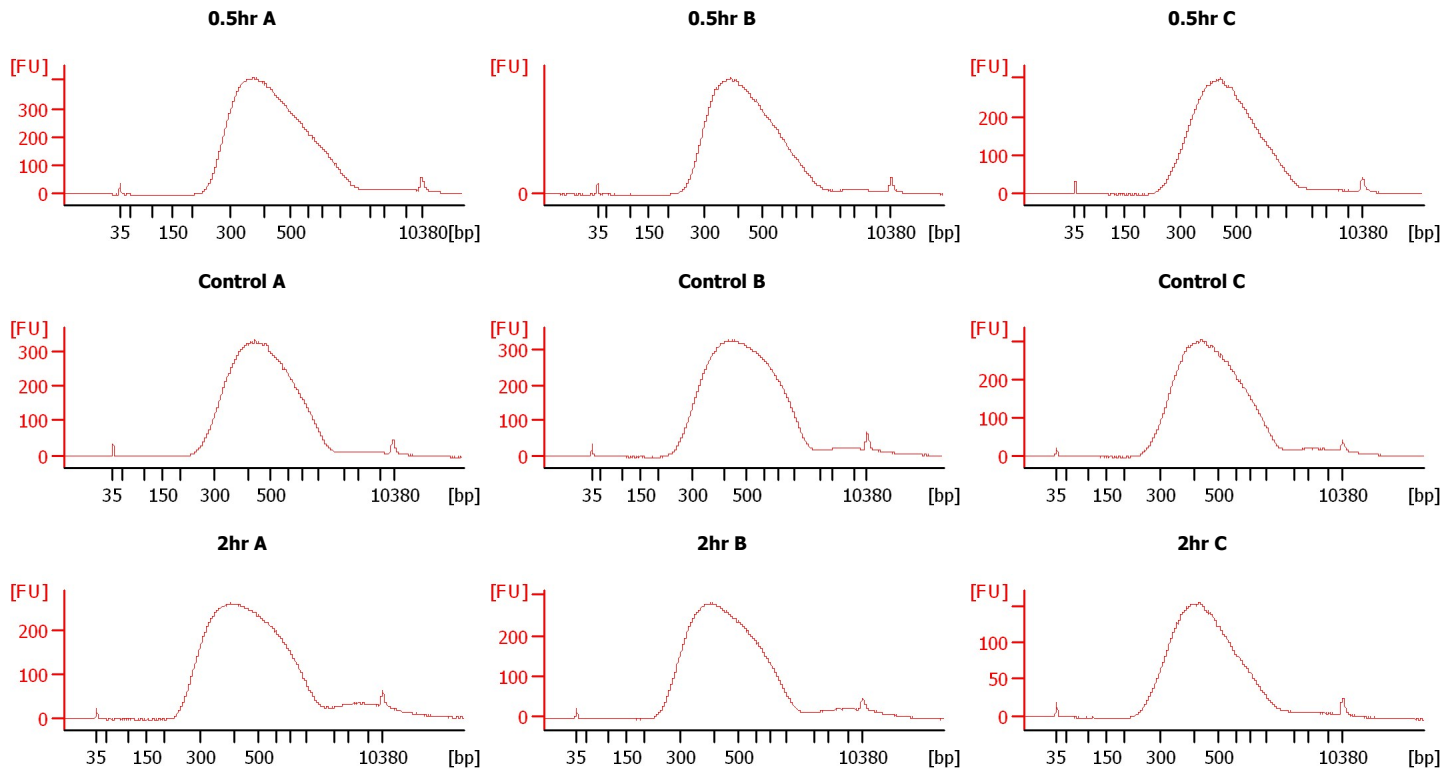
Instrument Name: DE13701086      Firmware: C.01.069  
Serial#: DE13701086      Type: G2938B

#### 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:



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
 Modified: 3/2/2017 3:38:29 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
0.5hr A		<input type="checkbox"/>	✓			
0.5hr B		<input type="checkbox"/>	✓			
0.5hr C		<input type="checkbox"/>	✓			
Control A		<input type="checkbox"/>	✓			
Control B		<input type="checkbox"/>	✓			
Control C		<input type="checkbox"/>	✓			
2hr A		<input type="checkbox"/>	✓			
2hr B		<input type="checkbox"/>	✓			
2hr C		<input type="checkbox"/>	✓			
na		<input type="checkbox"/>				
na		<input type="checkbox"/>				
Ladder		<input type="checkbox"/>	✓			

**Chip Lot #**

**Reagent Kit Lot #**

**Chip Comments :**

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
Modified: 3/2/2017 3:38:29 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

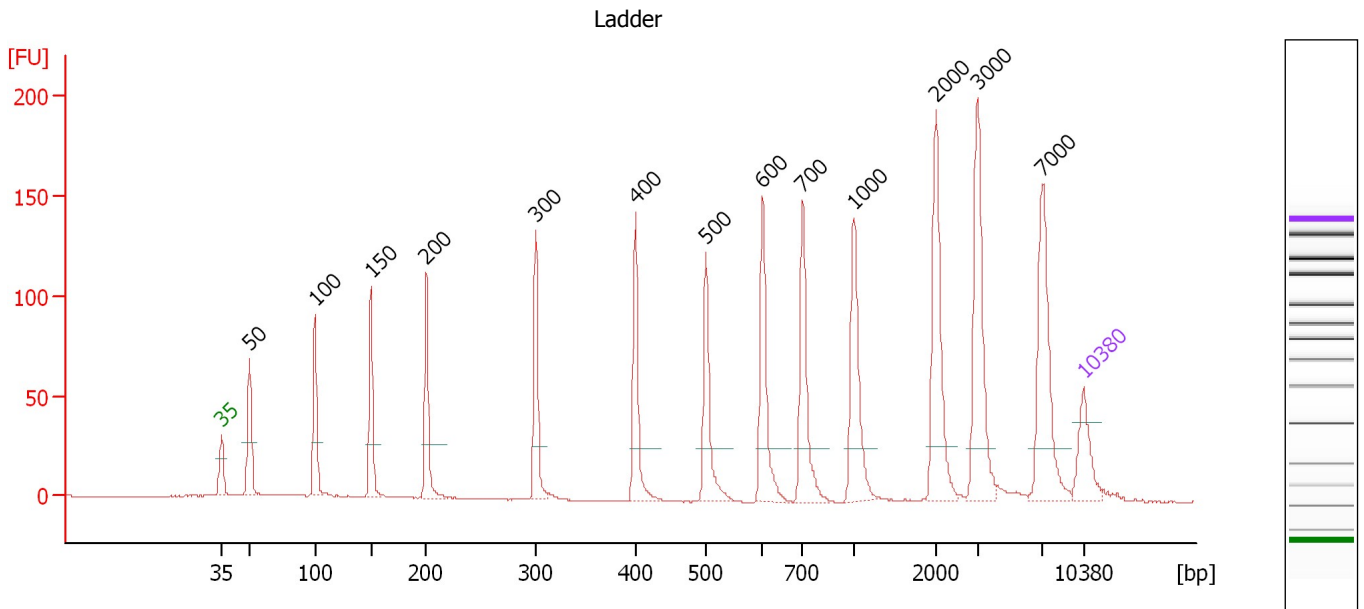
**Ladder**

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:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
 Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.2

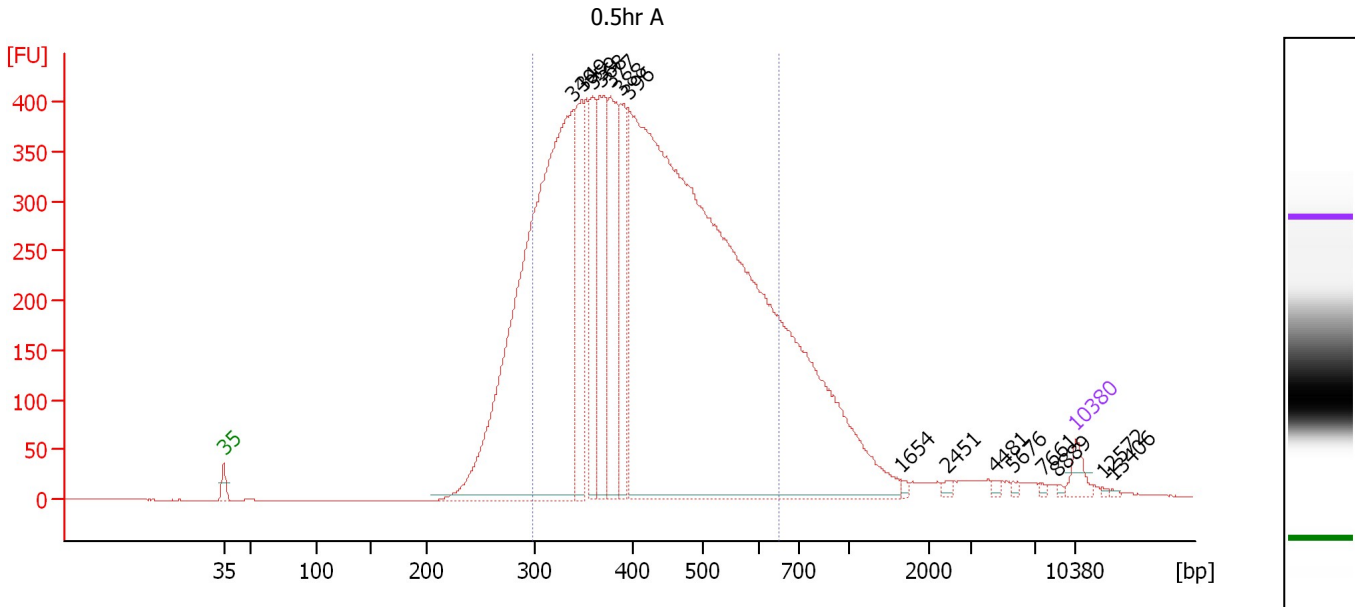
**Peak table for Ladder**

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	45.25
3	100	150.00	2,272.7	Ladder Peak	50.58
4	150	150.00	1,515.2	Ladder Peak	55.12
5	200	150.00	1,136.4	Ladder Peak	59.62
6	300	150.00	757.6	Ladder Peak	68.50
7	400	150.00	568.2	Ladder Peak	76.59
8	500	150.00	454.5	Ladder Peak	82.35
9	600	150.00	378.8	Ladder Peak	86.94
10	700	150.00	324.7	Ladder Peak	90.19
11	1,000	150.00	227.3	Ladder Peak	94.34
12	2,000	150.00	113.6	Ladder Peak	100.97
13	3,000	150.00	75.8	Ladder Peak	104.34
14	7,000	150.00	32.5	Ladder Peak	109.62
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
 Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : 0.5hr A**

Number of peaks found: 15                      Corr. Area 1: 8,666.2  
 Noise: 0.2

**Peak table for sample 1 : 0.5hr A**

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	340	5,033.71	22,431.9		71.74
3	349	798.58	3,465.0		72.48
4	359	571.63	2,413.0		73.27
5	368	907.15	3,739.1		73.97
6	377	879.60	3,531.9		74.76
7	388	605.91	2,368.4		75.59
8	396	10,332.48	39,559.9		76.25
9	1,654	15.91	14.6		98.68
10	2,451	19.32	11.9		102.49
11	4,481	14.39	4.9		106.30
12	5,676	12.13	3.2		107.87
13	7,661	8.90	1.8		110.28
14	8,889	7.33	1.2		111.51
15	10,380	75.00	10.9	Upper Marker	113.00
16	12,572	0.00	0.0		115.19
17	13,406	0.00	0.0		116.02

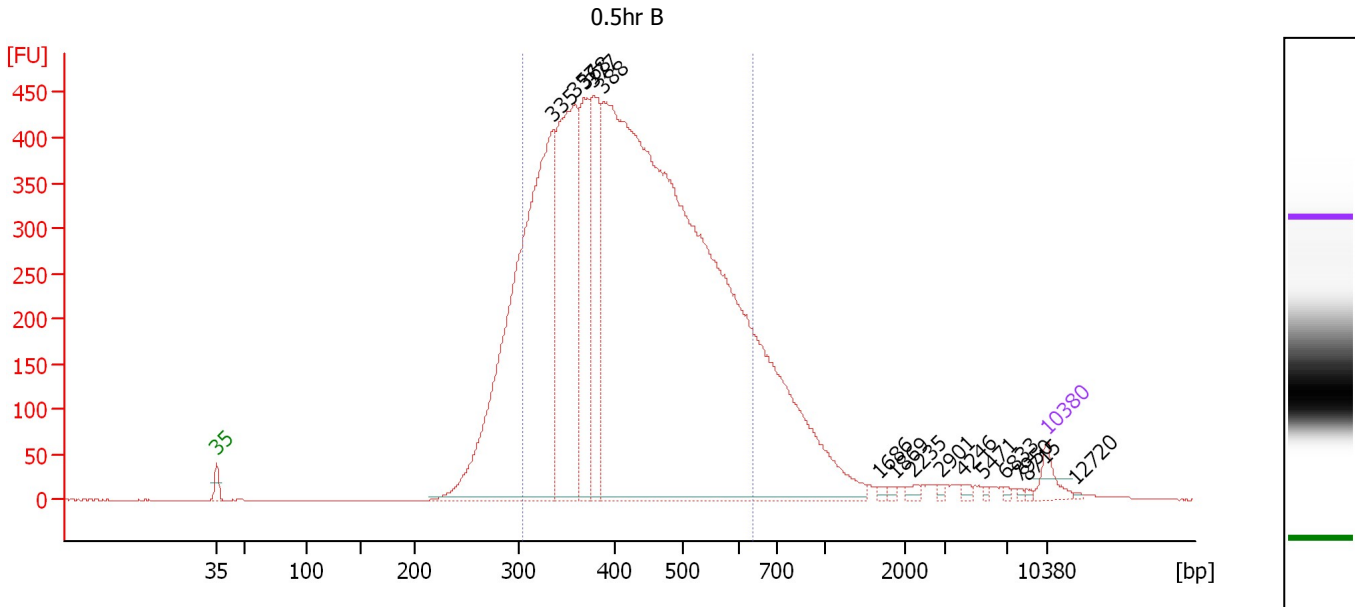
**Region table for sample 1 : 0.5hr A**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
299	653	428	8,666.2	55,927.4	14,936.94	79	21.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
 Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : 0.5hr B**

Number of peaks found: 15                      Corr. Area 1: 9,047.4  
 Noise: 0.4

**Peak table for sample 2 : 0.5hr B**

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	335	3,573.12	16,138.3		71.37
3	357	1,692.60	7,181.1		73.12
4	368	815.27	3,359.7		73.98
5	377	803.16	3,226.9		74.74
6	388	9,416.54	36,803.4		75.59
7	1,686	12.31	11.1		98.88
8	1,869	14.40	11.7		100.10
9	2,235	19.87	13.5		101.76
10	2,901	11.66	6.1		104.01
11	4,246	15.06	5.4		105.99
12	5,471	8.41	2.3		107.61
13	6,833	7.68	1.7		109.40
14	7,950	7.76	1.5		110.57
15	8,715	6.92	1.2		111.34
16	10,380	75.00	10.9	Upper Marker	113.00
17	12,720	0.00	0.0		115.34

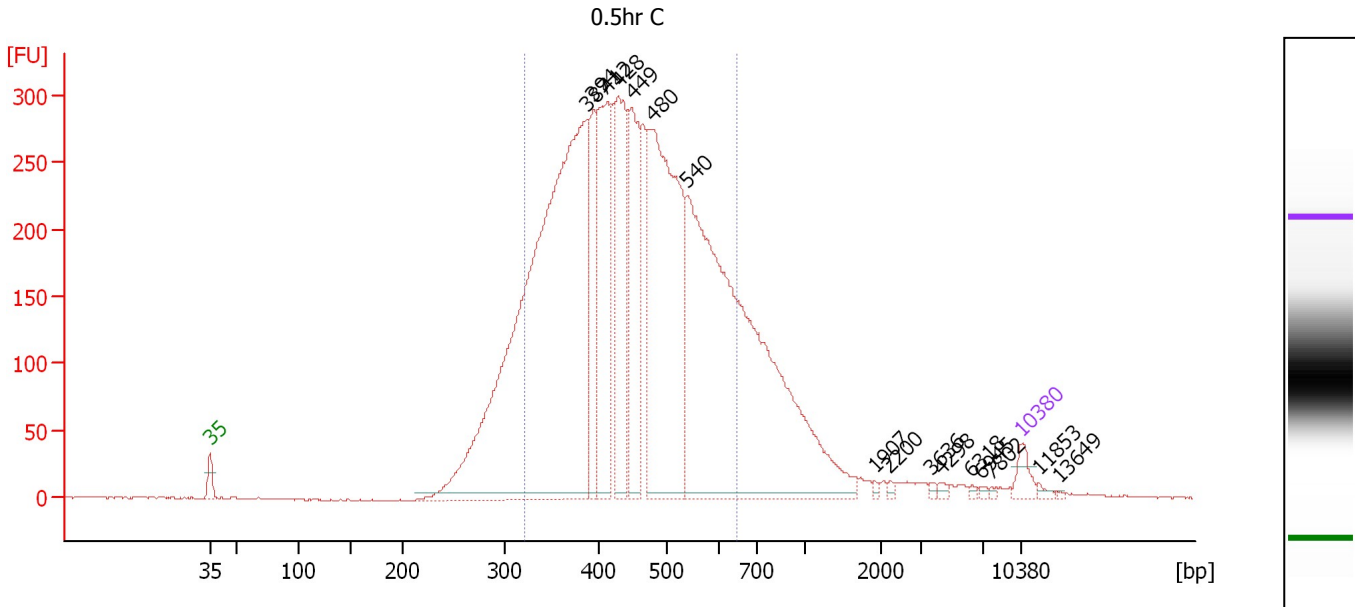
**Region table for sample 2 : 0.5hr B**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
303	640	428	9,047.4	46,885.2	12,594.83	80	20.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
 Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : 0.5hr C**

Number of peaks found: 16                      Corr. Area 1: 5,643.3  
 Noise: 0.5

**Peak table for sample 3 : 0.5hr C**

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	387	5,526.48	21,653.6		75.52
3	394	615.40	2,369.5		76.07
4	412	1,187.65	4,362.8		77.31
5	428	1,019.51	3,605.5		78.23
6	449	889.48	3,000.2		79.43
7	480	2,334.42	7,363.3		81.22
8	540	4,235.21	11,872.4		84.21
9	1,907	12.89	10.2		100.35
10	2,200	12.74	8.8		101.64
11	3,636	15.18	6.3		105.18
12	4,298	16.52	5.8		106.06
13	6,318	9.72	2.3		108.72
14	6,945	12.32	2.7		109.55
15	7,802	8.78	1.7		110.42
16	10,380	75.00	10.9	Upper Marker	113.00
17	11,853	0.00	0.0		114.47
18	13,649	0.00	0.0		116.27

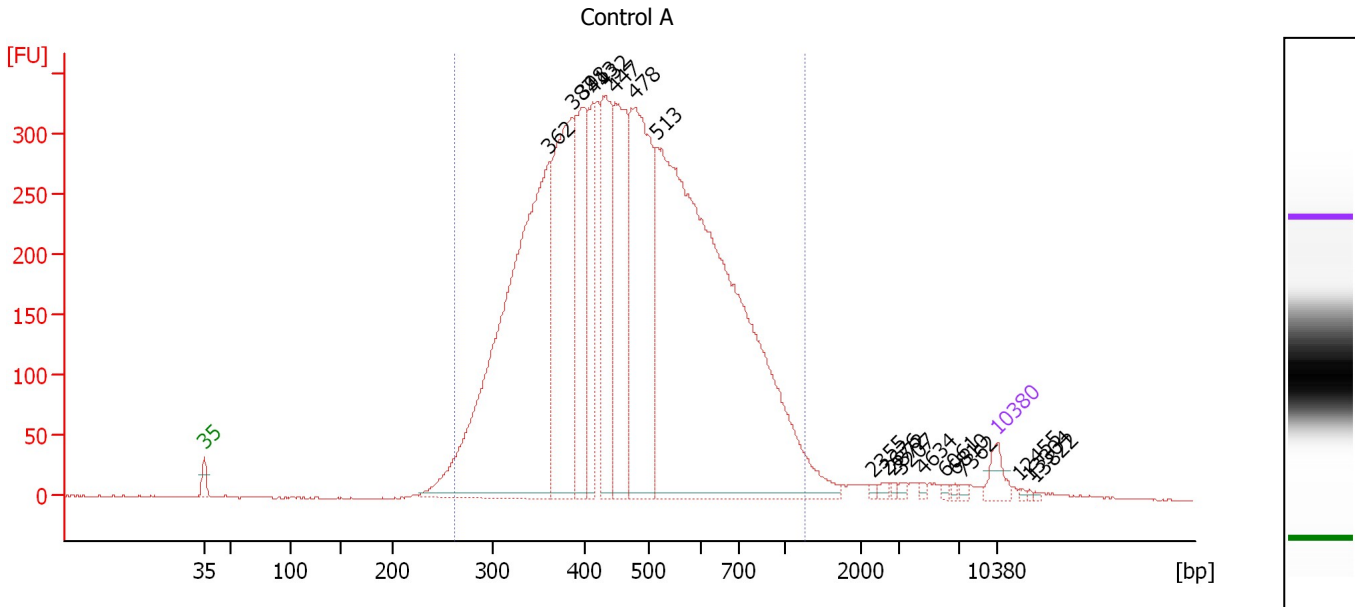
**Region table for sample 3 : 0.5hr C**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
321	649	451	5,643.3	46,037.5	13,079.44	77	18.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
 Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : Control A**

Number of peaks found: 19                      Corr. Area 1: 8,276.2  
 Noise: 0.7

**Peak table for sample 4 : Control A**

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	362	3,483.06	14,597.8		73.48
3	387	1,461.02	5,726.9		75.50
4	398	843.50	3,209.7		76.45
5	413	527.14	1,934.0		77.34
6	432	843.45	2,959.6		78.42
7	447	1,022.31	3,462.5		79.32
8	478	1,534.46	4,867.8		81.06
9	513	5,064.76	14,959.5		82.95
10	2,355	10.08	6.5		102.17
11	2,676	18.79	10.6		103.25
12	2,872	10.48	5.5		103.91
13	3,207	13.55	6.4		104.62
14	4,634	10.35	3.4		106.50
15	6,061	11.94	3.0		108.38
16	6,810	10.45	2.3		109.37
17	7,362	12.37	2.5		109.99
18	10,380	75.00	10.9	Upper Marker	113.00
19	12,455	0.00	0.0		115.07
20	13,304	0.00	0.0		115.92
21	13,822	0.00	0.0		116.44

**Region table for sample 4 : Control A**

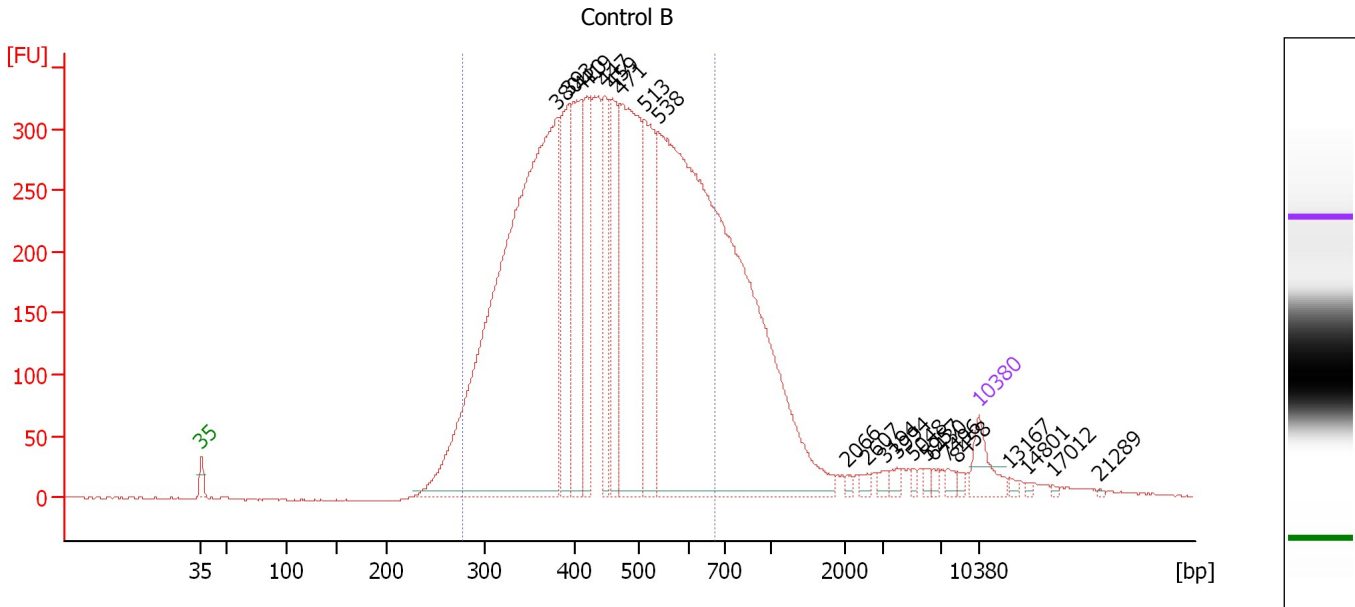
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
263	1,262	493	8,276.2	50,777.7	14,625.21	96	33.0



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
 Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : Control B**

Number of peaks found: 22                      Corr. Area 1: 7,616.9  
 Noise: 0.7

**Peak table for sample 5 : Control B**

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	380	2,866.82	11,430.2		74.98
3	393	436.92	1,684.2		76.03
4	410	523.85	1,934.7		77.18
5	419	306.63	1,109.9		77.66
6	447	299.56	1,015.5		79.30
7	459	315.42	1,040.1		80.02
8	471	872.71	2,806.6		80.69
9	513	487.57	1,440.2		82.95
10	538	3,293.96	9,275.5		84.10
11	2,066	8.98	6.6		101.19
12	2,607	13.36	7.8		103.01
13	3,194	17.17	8.1		104.60
14	3,994	17.07	6.5		105.65
15	5,048	11.37	3.4		107.05
16	5,957	12.22	3.1		108.25
17	6,430	12.66	3.0		108.87
18	7,496	17.44	3.5		110.12
19	8,458	10.02	1.8		111.08
20	10,380	75.00	10.9	Upper Marker	113.00
21	13,167	0.00	0.0		115.78
22	14,801	0.00	0.0		117.42
23	17,012	0.00	0.0		119.63
24	21,289	0.00	0.0		123.90

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary Continued ...**

... Region table for sample 5 :

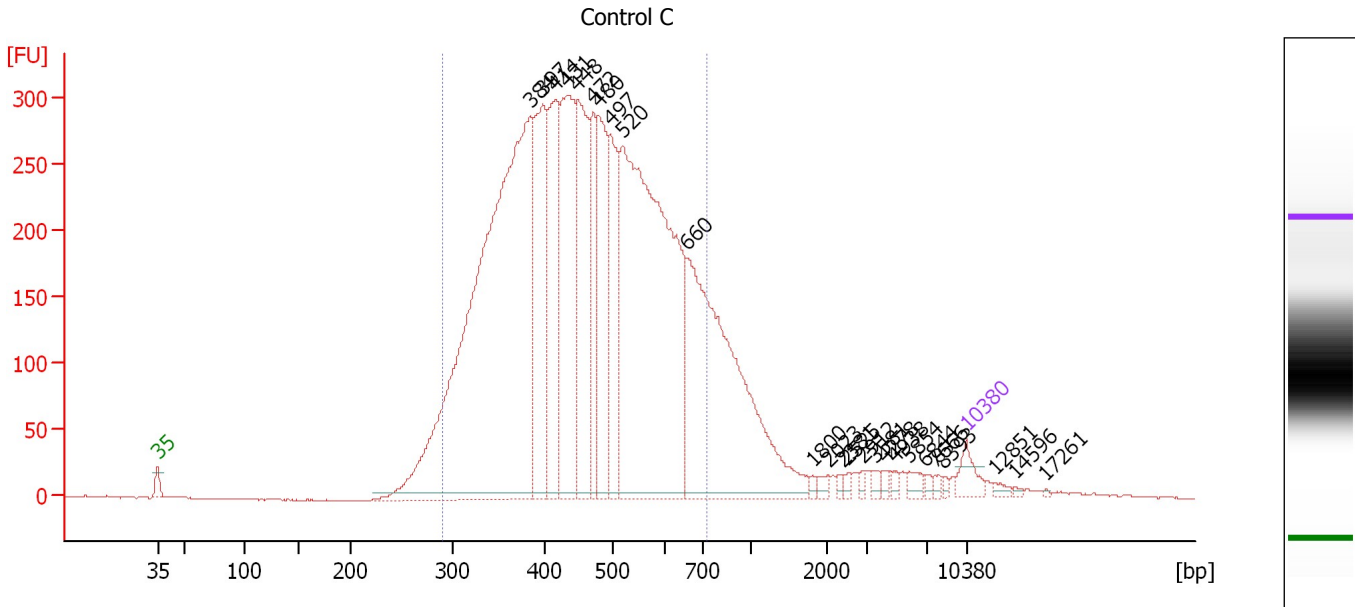
**Control B**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
277	668	451	7,616.9	30,212.5	8,385.54	81	22.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
 Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : Control C**

Number of peaks found: 25                      Corr. Area 1: 6,584.1  
 Noise: 0.6

**Peak table for sample 6 : Control C**

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	384	4,255.16	16,794.4		75.29
3	397	891.09	3,400.4		76.35
4	414	784.36	2,868.2		77.42
5	431	1,032.87	3,629.6		78.39
6	448	855.54	2,893.6		79.36
7	472	428.77	1,377.8		80.71
8	480	642.93	2,029.7		81.20
9	497	572.32	1,745.7		82.16
10	520	2,933.75	8,545.3		83.28
11	660	1,869.88	4,291.7		88.89
12	1,800	15.09	12.7		99.64
13	2,023	26.82	20.1		101.04
14	2,381	14.97	9.5		102.25
15	2,525	17.07	10.2		102.74
16	2,912	17.19	8.9		104.04
17	3,581	23.03	9.7		105.11
18	4,278	21.32	7.6		106.03
19	4,938	19.31	5.9		106.90
20	5,854	36.60	9.5		108.11
21	6,844	15.86	3.5		109.42
22	7,666	17.86	3.5		110.29
23	8,393	12.60	2.3		111.02
24	10,380	75.00	10.9	Upper Marker	113.00
25	12,851	0.00	0.0		115.47


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary Continued ...****... Peak table for sample 6 : Control C**

Peak	Size [bp]	Conc. [pg/ $\mu$ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	14,596	0.00	0.0		117.21
27	17,261	0.00	0.0		119.87

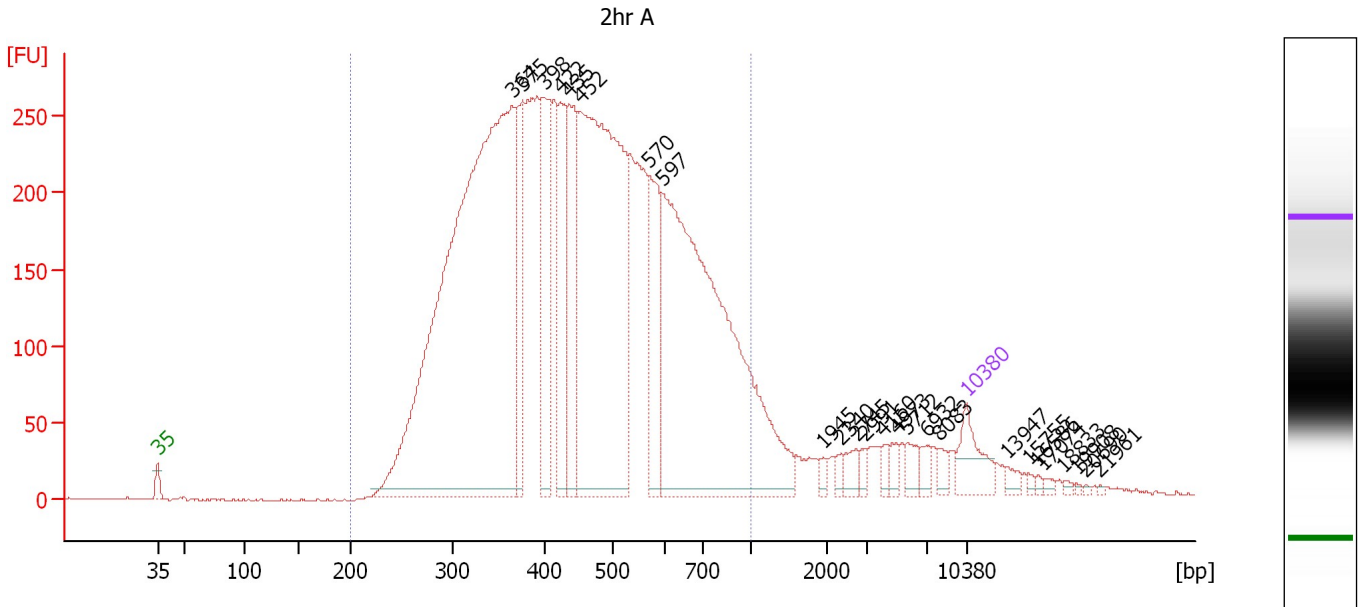
**Region table for sample 6 : Control C**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ $\mu$ l]	% of Total	Size distribution in CV [%]
291	726	463	6,584.1	44,527.7	 12,713.87	85	22.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
 Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : 2hr A**

Number of peaks found: 25                      Corr. Area 1: 7,409.6  
 Noise: 0.8

**Peak table for sample 7 : 2hr A**

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	364	2,407.57	10,029.4		73.66
3	375	240.89	972.9		74.58
4	398	315.17	1,199.6		76.44
5	422	288.17	1,034.9		77.85
6	435	267.32	931.9		78.59
7	452	1,317.70	4,421.3		79.56
8	570	249.71	663.7		85.57
9	597	1,356.52	3,444.7		86.79
10	1,945	12.44	9.7		100.60
11	2,340	14.37	9.3		102.11
12	2,745	27.35	15.1		103.48
13	2,991	15.91	8.1		104.31
14	4,160	16.06	5.9		105.87
15	4,973	19.76	6.0		106.95
16	5,712	29.48	7.8		107.92
17	6,932	20.90	4.6		109.53
18	8,083	18.87	3.5		110.71
19	10,380	75.00	10.9	Upper Marker	113.00
20	13,947	0.00	0.0		116.56
21	15,755	0.00	0.0		118.37
22	16,586	0.00	0.0		119.20
23	17,074	0.00	0.0		119.69
24	18,833	0.00	0.0		121.44
25	19,908	0.00	0.0		122.52


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary Continued ...****... Peak table for sample 7 : 2hr A**

Peak	Size [bp]	Conc. [pg/ $\mu$ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	20,690	0.00	0.0		123.30
27	21,961	0.00	0.0		124.57

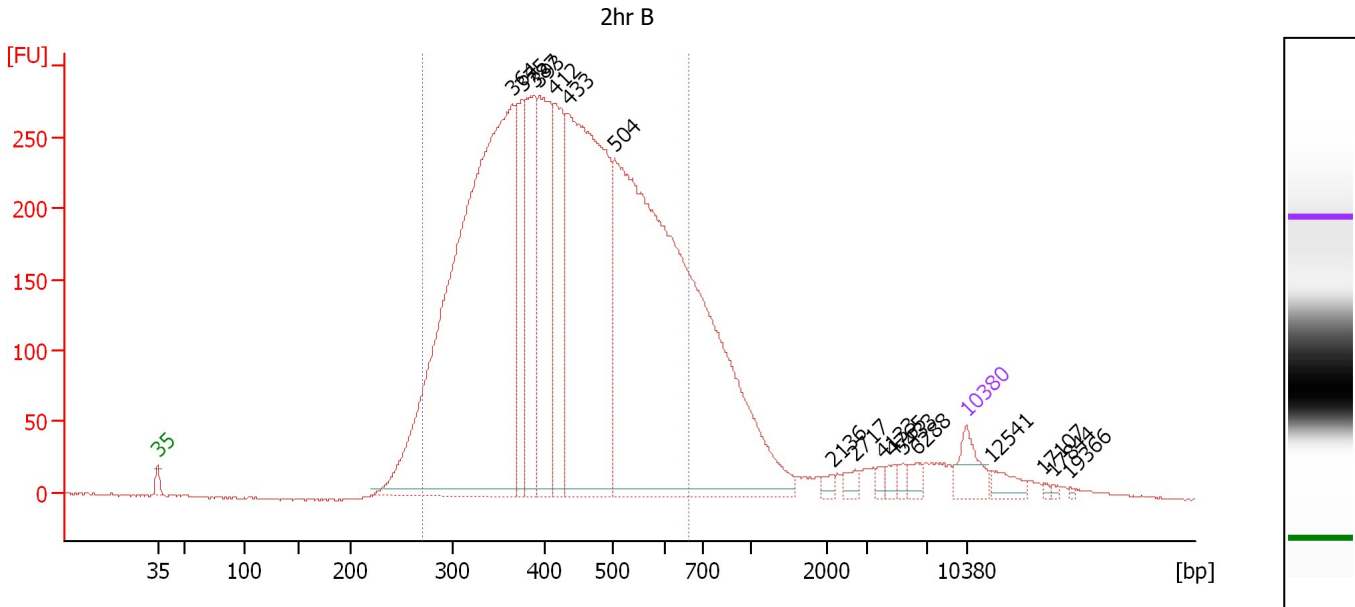
**Region table for sample 7 : 2hr A**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ $\mu$ l]	% of Total	Size distribution in CV [%]
200	1,000	472	7,409.6	27,740.0	 7,569.88	91	32.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
 Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : 2hr B**

Number of peaks found: 17                      Corr. Area 1: 6,500.1  
 Noise: 0.7

**Peak table for sample 8 : 2hr B**

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	364	2,811.85	11,692.8		73.71
3	375	352.86	1,424.7		74.59
4	387	477.82	1,868.9		75.57
5	393	596.70	2,297.9		76.06
6	412	428.05	1,573.8		77.29
7	433	1,551.45	5,424.0		78.52
8	504	2,913.90	8,758.7		82.54
9	2,136	15.73	11.2		101.42
10	2,717	22.10	12.3		103.39
11	4,133	17.25	6.3		105.84
12	4,765	18.08	5.7		106.67
13	5,433	18.63	5.2		107.56
14	6,288	24.60	5.9		108.68
15	10,380	75.00	10.9	Upper Marker	113.00
16	12,541	0.00	0.0		115.16
17	17,107	0.00	0.0		119.72
18	17,844	0.00	0.0		120.46
19	19,366	0.00	0.0		121.98

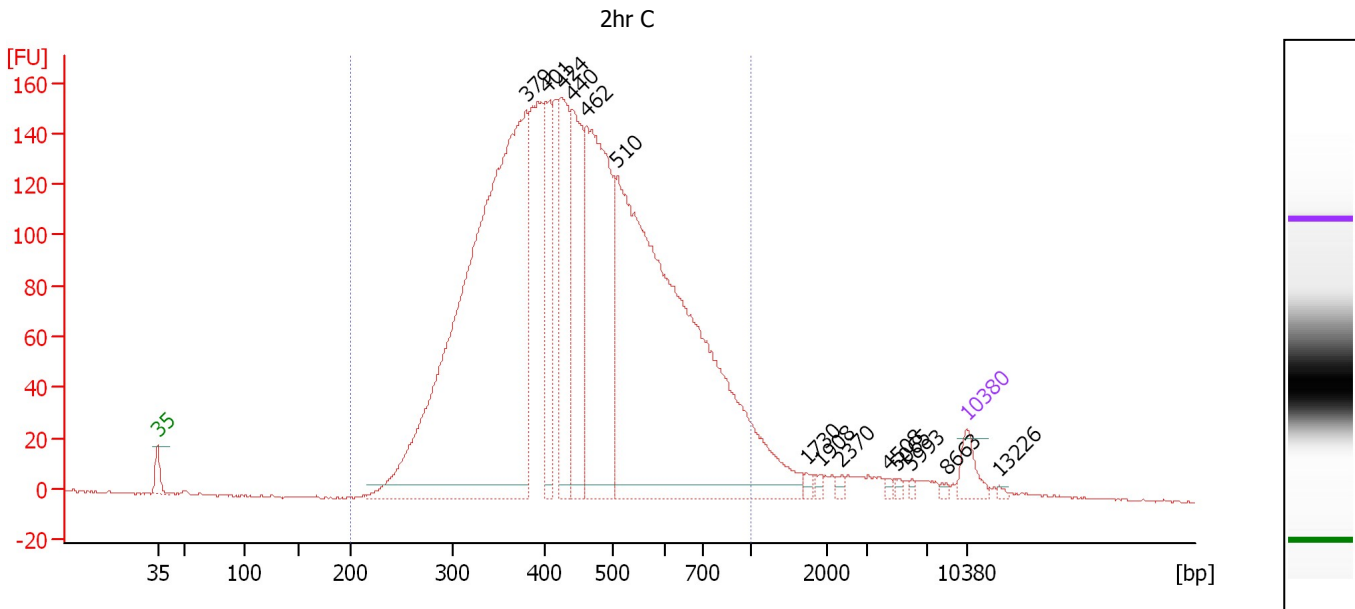
**Region table for sample 8 : 2hr B**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
271	660	433	6,500.1	29,857.8	7,947.36	83	22.6

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
 Modified: 3/2/2017 3:38:29 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : 2hr C**

Number of peaks found: 14                      Corr. Area 1: 3,661.3  
 Noise: 0.4

**Peak table for sample 9 : 2hr C**

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	379	3,832.78	15,337.7		74.86
3	401	405.40	1,533.3		76.63
4	424	653.80	2,333.9		78.00
5	440	619.06	2,132.9		78.88
6	462	1,347.63	4,420.6		80.16
7	510	3,372.93	10,023.0		82.80
8	1,730	16.99	14.9		99.18
9	1,908	16.61	13.2		100.35
10	2,370	14.70	9.4		102.22
11	4,508	10.32	3.5		106.33
12	5,065	10.43	3.1		107.07
13	5,993	9.50	2.4		108.29
14	8,663	9.58	1.7		111.28
15	10,380	75.00	10.9	Upper Marker	113.00
16	13,226	0.00	0.0		115.84

**Region table for sample 9 : 2hr C**

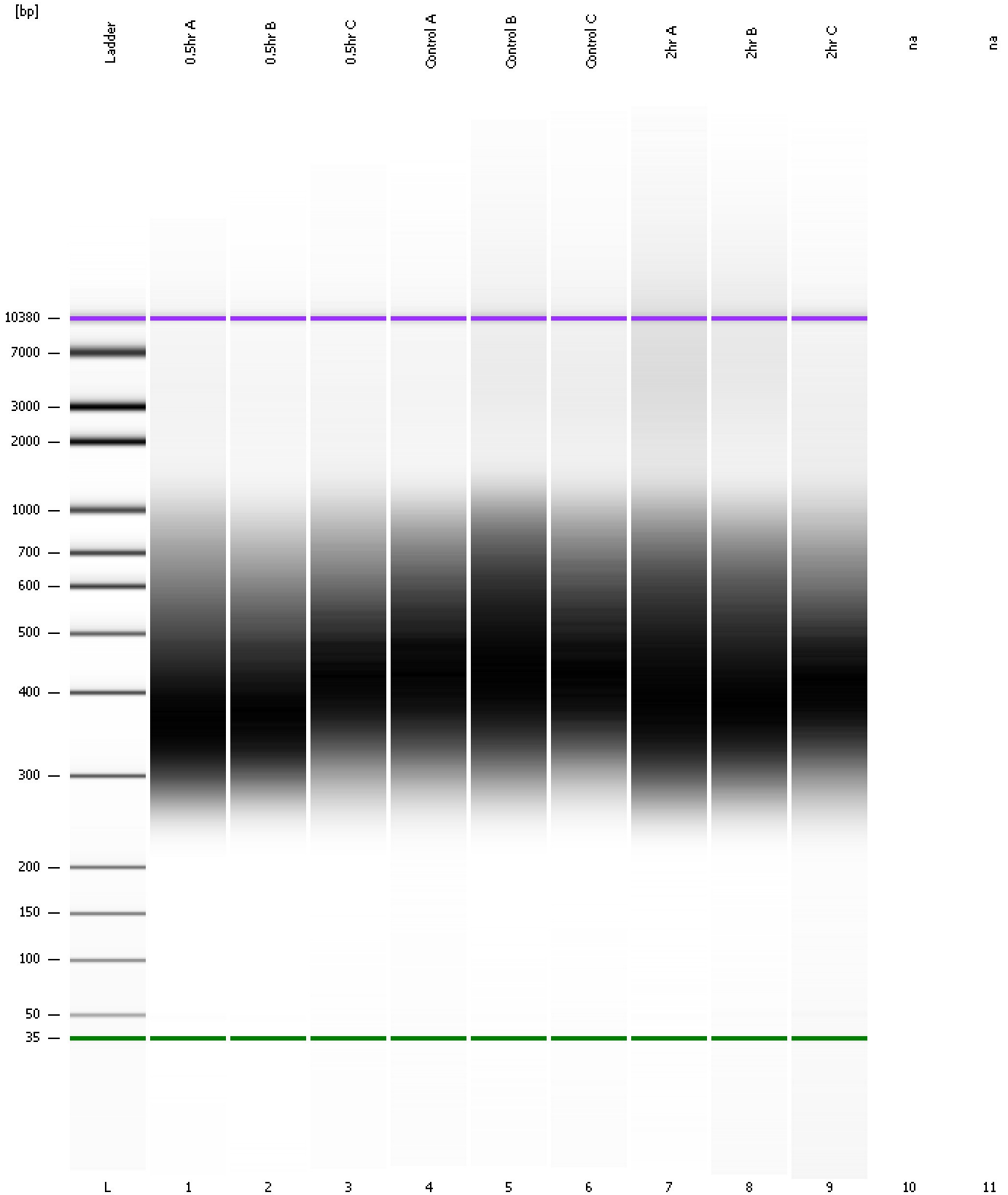
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	465	3,661.3	39,983.3	11,001.55	95	30.3



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
Modified: 3/2/2017 3:38:29 PM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
Modified: 3/2/2017 3:38:29 PM

**Invalid Samples**

Sample 10 has not been run, no results available.

Sample 11 has not been run, no results available.

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-03-02\2017-03-02\_002.xad

Created: 3/2/2017 3:02:49 PM  
 Modified: 3/2/2017 3:38:29 PM

**Run Logbook**

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 10)		Instrument	Run		3/2/2017 3:38:25 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2017-03-02\2017-03-02_002.xad)		Instrument	Run		3/2/2017 3:02:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		3/2/2017 3:02:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		3/2/2017 3:02:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		3/2/2017 3:02:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		3/2/2017 3:02:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		3/2/2017 3:02:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		3/2/2017 3:02:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1