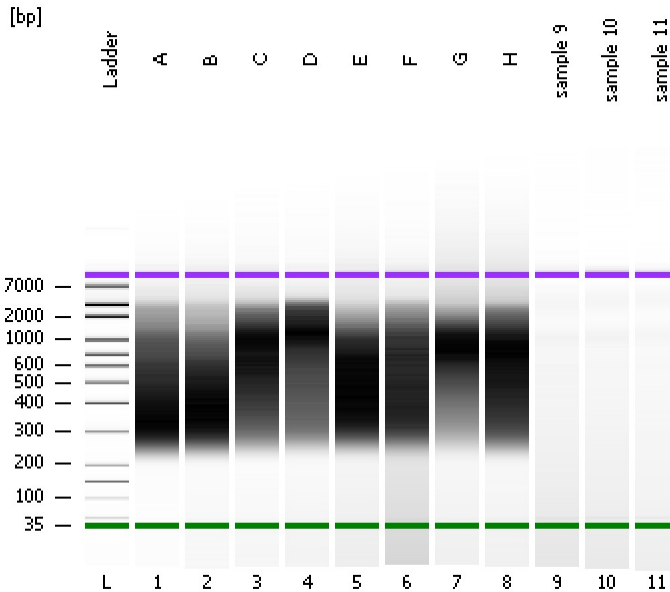


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
Modified: 7/26/2016 12:35:05 PM

Electrophoresis File Run Summary



Instrument Information:

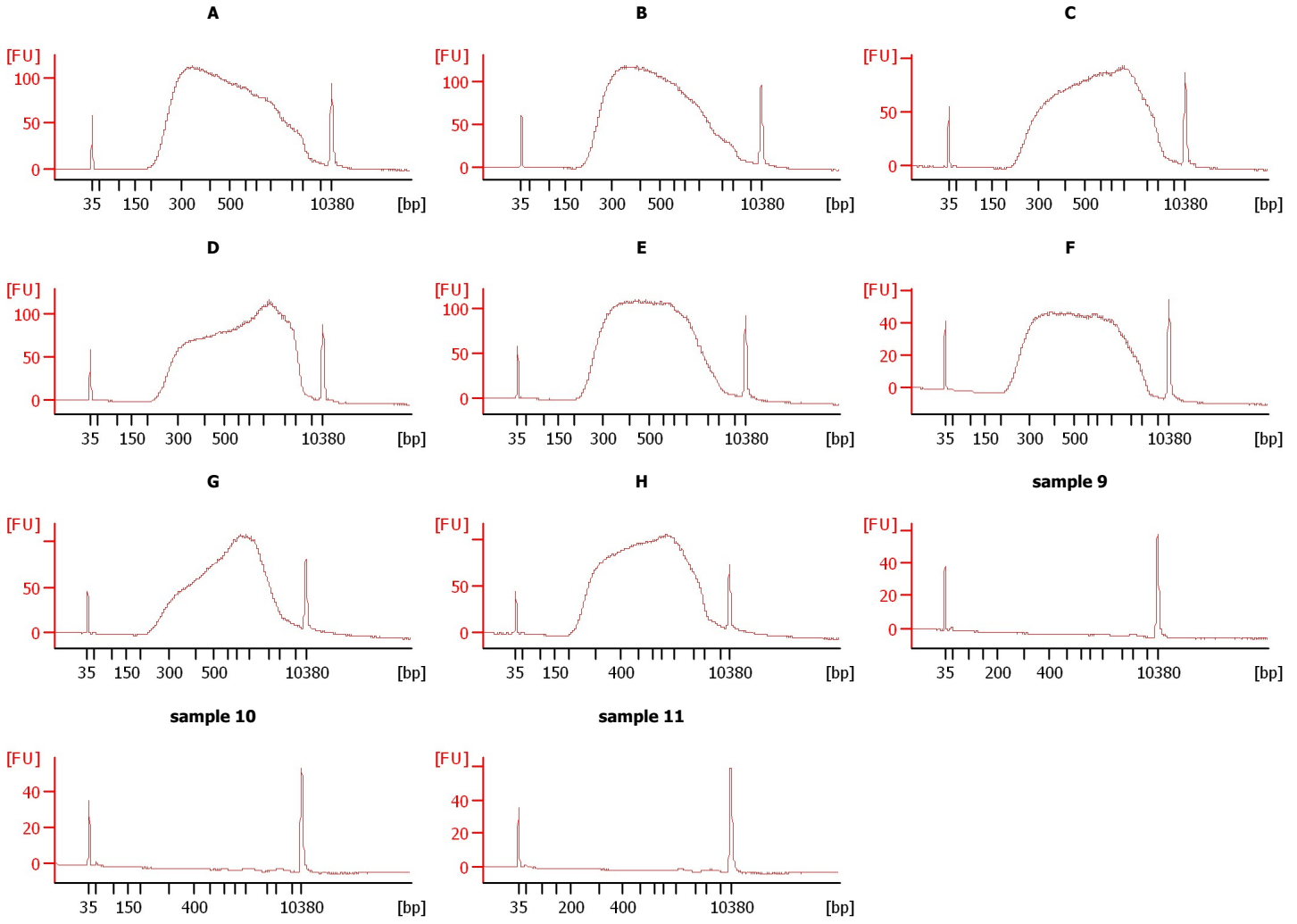
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
Modified: 7/26/2016 12:35:05 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
A		<input type="checkbox"/>	✓			
B		<input type="checkbox"/>	✓			
C		<input type="checkbox"/>	✓			
D		<input type="checkbox"/>	✓			
E		<input type="checkbox"/>	✓			
F		<input type="checkbox"/>	✓			
G		<input type="checkbox"/>	✓			
H		<input type="checkbox"/>	✓			
sample 9		<input type="checkbox"/>	✓			
sample 10		<input type="checkbox"/>	✓			
sample 11		<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\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
Modified: 7/26/2016 12:35:05 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

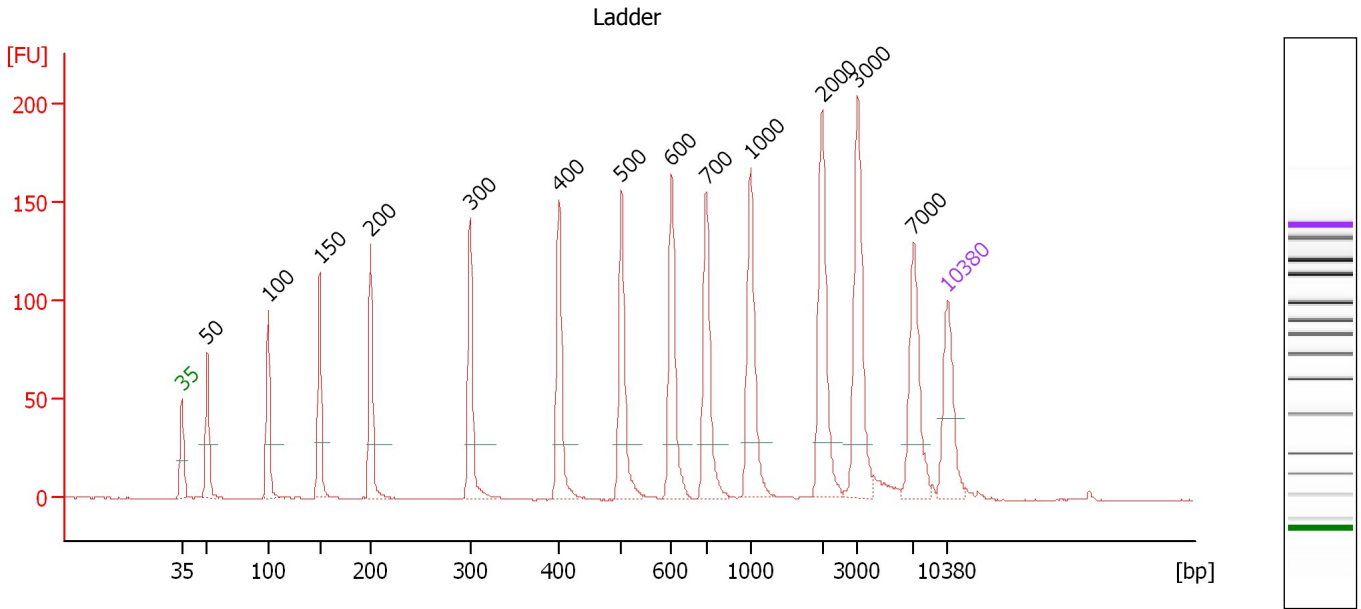
Ladder

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\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

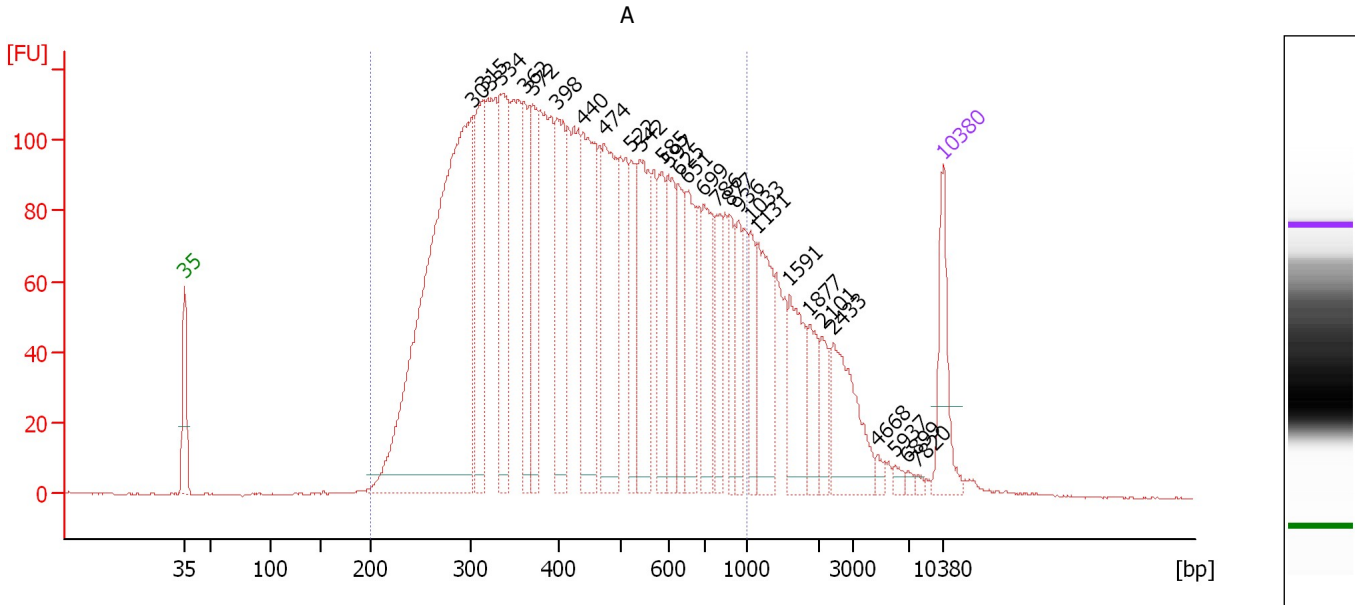
Peak table for Ladder

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	45.34
3	100	150.00	2,272.7	Ladder Peak	50.89
4	150	150.00	1,515.2	Ladder Peak	55.62
5	200	150.00	1,136.4	Ladder Peak	60.24
6	300	150.00	757.6	Ladder Peak	69.35
7	400	150.00	568.2	Ladder Peak	77.49
8	500	150.00	454.5	Ladder Peak	83.19
9	600	150.00	378.8	Ladder Peak	87.77
10	700	150.00	324.7	Ladder Peak	90.93
11	1,000	150.00	227.3	Ladder Peak	94.98
12	2,000	150.00	113.6	Ladder Peak	101.50
13	3,000	150.00	75.8	Ladder Peak	104.77
14	7,000	150.00	32.5	Ladder Peak	109.88
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 1 : A

Number of peaks found: 28 Corr. Area 1: 3,850.9
 Noise: 0.2

Peak table for sample 1 : 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	303	1,104.24	5,529.4		69.56
3	315	211.03	1,016.1		70.55
4	334	238.64	1,082.5		72.12
5	362	149.64	626.6		74.38
6	372	135.52	552.7		75.17
7	398	192.96	734.4		77.34
8	440	250.14	860.4		79.80
9	474	243.17	777.1		81.71
10	522	97.89	284.4		84.17
11	542	187.89	525.3		85.11
12	585	98.12	254.2		87.08
13	597	112.83	286.5		87.62
14	625	92.32	223.9		88.55
15	651	111.58	259.6		89.39
16	699	124.15	269.0		90.91
17	786	65.51	126.3		92.09
18	877	61.08	105.5		93.32
19	936	57.89	93.7		94.11
20	1,033	61.36	90.0		95.19
21	1,131	125.22	167.7		95.83
22	1,591	93.15	88.7		98.83
23	1,877	52.94	42.7		100.70
24	2,101	36.13	26.1		101.83
25	2,433	112.22	69.9		102.92

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
 Modified: 7/26/2016 12:35:05 PM

Electropherogram Summary Continued ...

... Peak table for sample 1 : A

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	4,668	8.59	2.8		106.90
27	5,937	7.85	2.0		108.52
28	6,899	4.64	1.0		109.75
29	7,820	3.86	0.7		110.64
30	10,380	75.00	10.9	Upper Marker	113.00

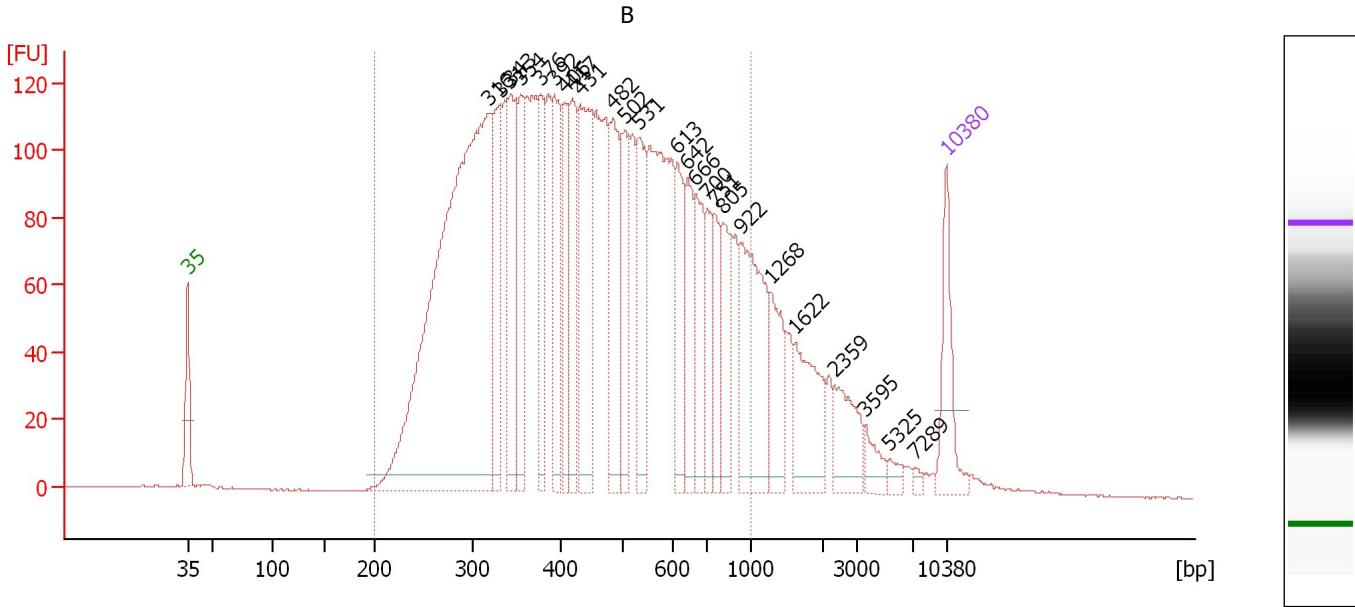
Region table for sample 1 : A

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	464	3,850.9	20,209.4	5,157.39	86	38.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
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Electropherogram Summary Continued ...



Overall Results for sample 2 : B

Number of peaks found: 25 Corr. Area 1: 3,958.7
 Noise: 0.2

Peak table for sample 2 : 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	316	1,203.55	5,766.2		70.68
3	331	162.21	743.0		71.86
4	343	192.51	850.8		72.84
5	354	129.00	552.6		73.72
6	376	122.66	494.2		75.54
7	392	162.29	627.6		76.82
8	406	113.89	425.5		77.80
9	417	130.31	473.8		78.44
10	431	237.13	834.5		79.23
11	482	155.52	488.6		82.18
12	502	120.18	363.1		83.26
13	531	146.69	418.9		84.59
14	613	104.29	257.8		88.18
15	642	100.24	236.4		89.11
16	666	111.18	253.1		89.85
17	700	71.81	155.5		90.93
18	751	77.53	156.5		91.62
19	805	81.96	154.2		92.35
20	922	202.34	332.5		93.93
21	1,268	77.72	92.8		96.73
22	1,622	107.96	100.8		99.04
23	2,359	68.24	43.8		102.68
24	3,595	24.72	10.4		105.53
25	5,325	11.40	3.2		107.74

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

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Electropherogram Summary Continued ...

... Peak table for sample 2 : B

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	7,289	5.18	1.1		110.15
27	10,380	75.00	10.9	Upper Marker	113.00

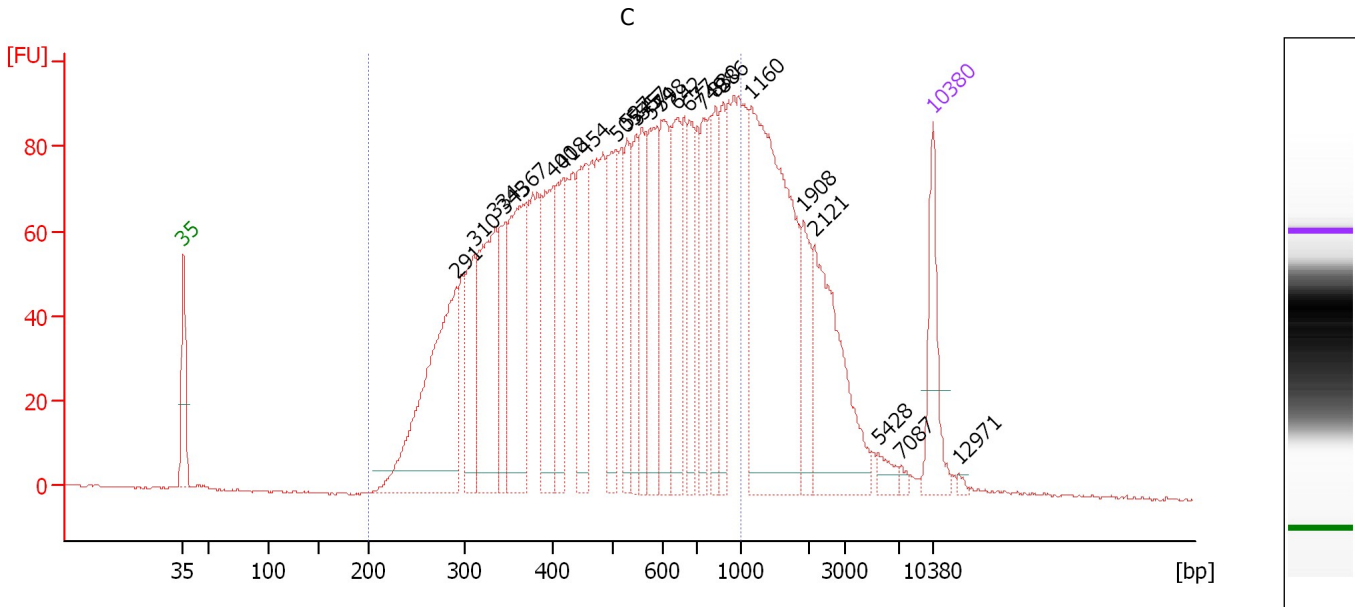
Region table for sample 2 : B

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
200	1,000	467	3,958.7	18,765.2	4,871.72	88	36.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 3 : C

Number of peaks found: 25 Corr. Area 1: 2,681.5
 Noise: 0.3

Peak table for sample 3 : 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	291	395.08	2,060.4		68.49
3	310	125.67	615.0		70.13
4	334	248.01	1,124.8		72.13
5	345	115.85	508.7		73.02
6	367	225.98	934.2		74.76
7	400	160.76	609.6		77.45
8	418	123.89	449.4		78.50
9	454	159.52	532.9		80.54
10	505	132.87	398.5		83.43
11	527	85.83	246.8		84.42
12	545	91.16	253.2		85.27
13	557	91.45	248.6		85.82
14	574	147.73	390.1		86.56
15	598	132.58	336.1		87.66
16	642	133.35	314.6		89.10
17	677	88.80	198.8		90.20
18	749	80.57	163.0		91.59
19	830	78.92	144.0		92.69
20	886	83.25	142.4		93.43
21	1,160	455.55	594.9		96.02
22	1,908	72.74	57.8		100.90
23	2,121	188.67	134.8		101.90
24	5,428	15.60	4.4		107.87
25	7,087	5.44	1.2		109.96

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
 Modified: 7/26/2016 12:35:05 PM

Electropherogram Summary Continued ...

... Peak table for sample 3 : C

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	10,380	75.00	10.9	Upper Marker	113.00
27	12,971	0.00	0.0		115.39

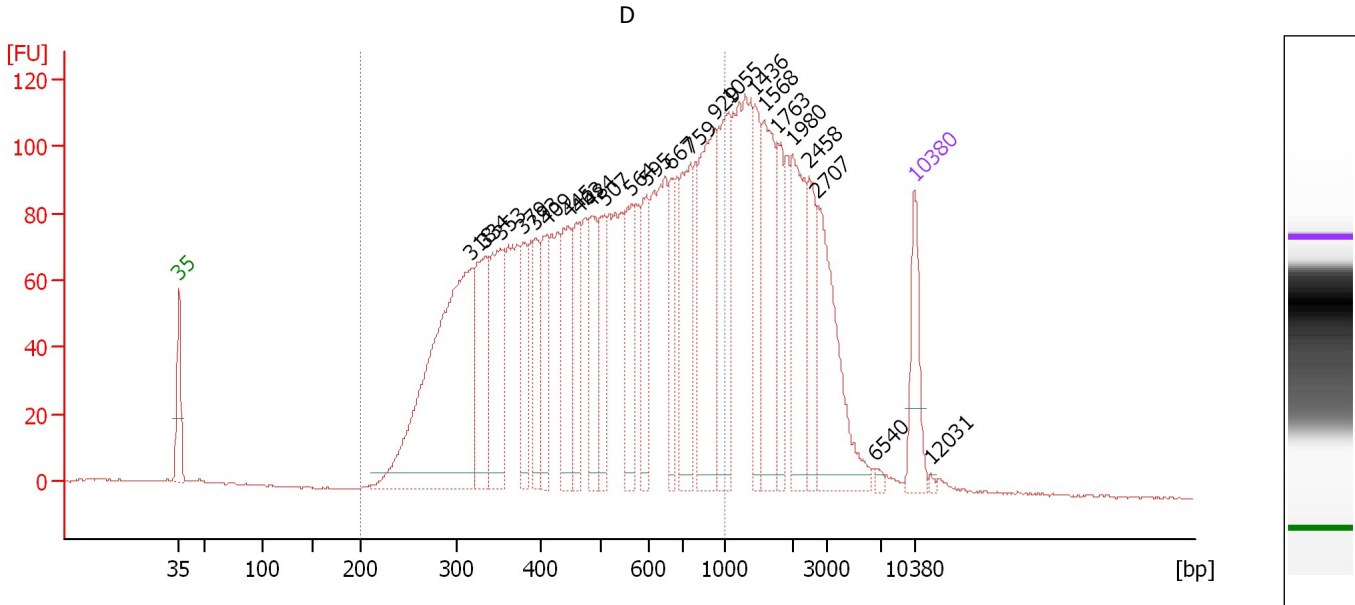
Region table for sample 3 : C

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
200	1,000	521	2,681.5	13,094.5	3,752.92	78	36.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
 Modified: 7/26/2016 12:35:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : D

Number of peaks found: 24 Corr. Area 1: 2,822.7
 Noise: 0.2

Peak table for sample 4 : D

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	318	690.03	3,287.8		70.82
3	334	201.54	913.7		72.14
4	353	199.70	857.3		73.66
5	379	104.73	419.2		75.74
6	393	109.76	423.3		76.91
7	409	115.54	427.6		78.03
8	445	143.21	487.6		80.06
9	463	109.95	359.9		81.07
10	484	124.36	389.1		82.29
11	507	108.55	324.4		83.51
12	564	129.41	348.0		86.10
13	595	108.02	275.3		87.52
14	667	94.96	215.6		89.90
15	759	169.37	338.0		91.73
16	929	225.96	368.7		94.02
17	1,055	195.89	281.3		95.34
18	1,436	89.81	94.8		97.82
19	1,568	166.59	161.0		98.69
20	1,763	82.05	70.5		99.95
21	1,980	157.97	120.9		101.38
22	2,458	82.11	50.6		103.00
23	2,707	190.26	106.5		103.81
24	6,540	5.61	1.3		109.29
25	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
Modified: 7/26/2016 12:35:05 PM

Electropherogram Summary Continued ...**... Peak table for sample 4 : D**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	12,031	0.00	0.0		114.52

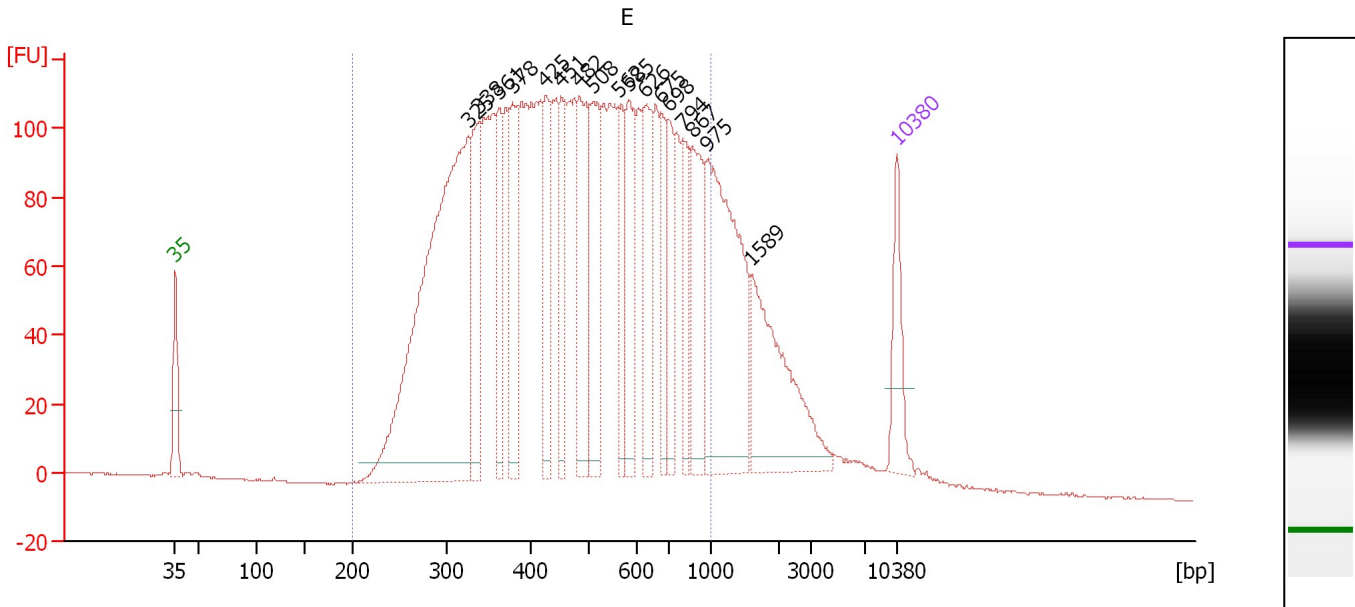
Region table for sample 4 : D

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
200	1,000	525	2,822.7	13,659.3	3,925.55	71	36.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 5 : E

Number of peaks found: 17 Corr. Area 1: 3,751.1
 Noise: 0.3

Peak table for sample 5 : E

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	325	1,068.06	4,976.3		71.40
3	338	202.45	907.7		72.44
4	361	137.37	575.8		74.35
5	378	185.43	743.3		75.70
6	425	127.44	454.5		78.91
7	451	124.27	417.3		80.41
8	482	195.07	613.1		82.16
9	508	197.64	589.3		83.56
10	562	106.96	288.2		86.05
11	585	143.74	372.3		87.08
12	626	163.22	395.2		88.58
13	675	99.07	222.5		90.13
14	698	104.73	227.5		90.86
15	794	93.73	178.8		92.20
16	867	167.51	292.7		93.18
17	975	399.92	621.8		94.63
18	1,589	246.03	234.5		98.82
19	10,380	75.00	10.9	Upper Marker	113.00

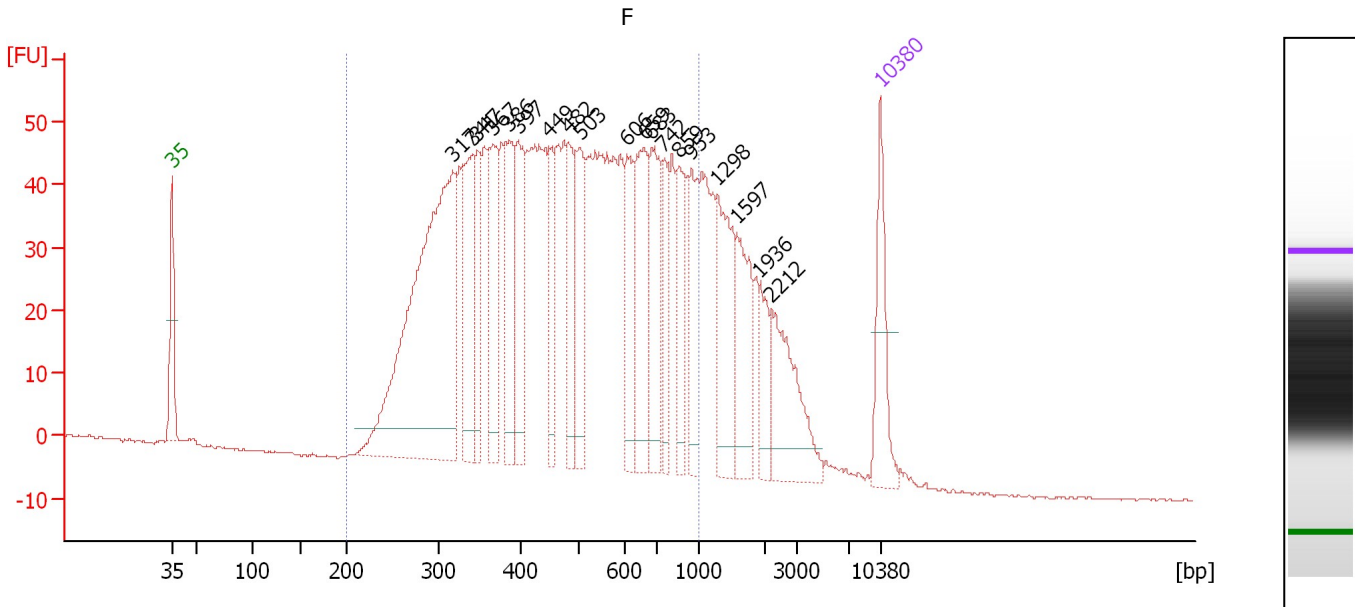
Region table for sample 5 : E

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	497	3,751.1	18,779.1	5,197.62	85	36.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 6 : F

Number of peaks found: 19 Corr. Area 1: 1,719.2
 Noise: 0.2

Peak table for sample 6 : F

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	317	662.98	3,170.1		70.73
3	341	156.01	692.2		72.73
4	347	93.60	409.1		73.15
5	367	126.52	522.7		74.78
6	386	123.17	483.2		76.37
7	397	125.84	480.8		77.21
8	449	82.34	278.0		80.27
9	482	87.07	273.7		82.16
10	503	115.00	346.4		83.32
11	606	93.14	232.8		87.96
12	659	143.48	329.7		89.65
13	683	128.12	284.3		90.39
14	742	66.25	135.3		91.49
15	859	83.54	147.4		93.08
16	953	82.32	130.9		94.34
17	1,298	128.37	149.8		96.92
18	1,597	93.43	88.6		98.87
19	1,936	44.98	35.2		101.09
20	2,212	114.07	78.2		102.19
21	10,380	75.00	10.9	Upper Marker	113.00

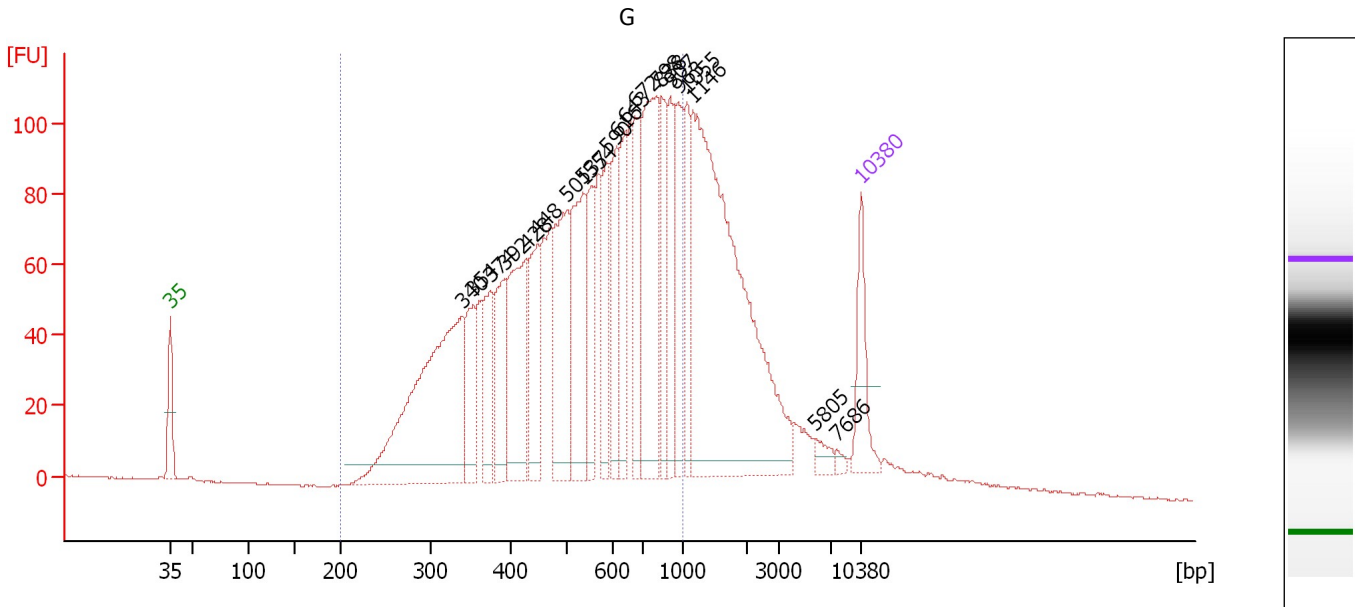
Region table for sample 6 : F

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	498	1,719.2	12,171.8	3,364.06	83	36.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
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Electropherogram Summary Continued ...



Overall Results for sample 7 : G

Number of peaks found: 21 Corr. Area 1: 2,416.9
 Noise: 0.2


Peak table for sample 7 : G

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	586.05	2,612.4		72.60
3	354	118.16	506.0		73.73
4	374	116.18	470.1		75.41
5	392	135.04	521.5		76.87
6	428	232.45	823.0		79.08
7	448	136.89	463.1		80.21
8	501	235.12	710.9		83.24
9	537	208.24	588.1		84.86
10	554	109.70	299.9		85.67
11	590	104.15	267.6		87.29
12	616	114.86	282.7		88.26
13	643	136.72	322.2		89.13
14	672	128.47	289.7		90.04
15	798	287.02	544.7		92.26
16	838	107.62	194.5		92.80
17	907	103.27	172.6		93.72
18	963	129.47	203.8		94.47
19	1,055	95.05	136.5		95.34
20	1,146	757.27	1,000.8		95.93
21	5,805	17.29	4.5		108.35
22	7,686	7.11	1.4		110.52
23	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
Modified: 7/26/2016 12:35:05 PM

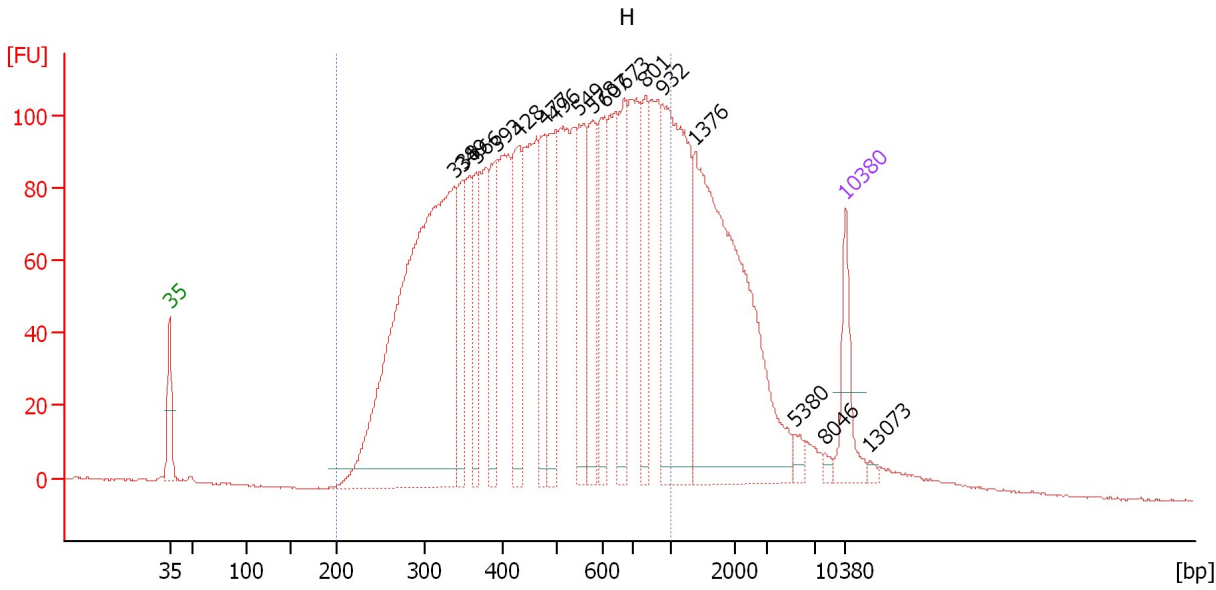
Electropherogram Summary Continued ...**... Region table for sample 7 : G**

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	557	2,416.9	11,701.0	 3,620.23	74	34.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
 Modified: 7/26/2016 12:35:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : H

Number of peaks found: 17 Corr. Area 1: 3,242.7
 Noise: 0.2

Peak table for sample 8 : H

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	338	1,167.45	5,226.7		72.48
3	349	123.10	534.0		73.36
4	366	121.87	504.2		74.74
5	393	154.96	597.9		76.90
6	428	174.12	615.9		79.10
7	477	128.95	409.8		81.86
8	496	164.96	503.8		82.97
9	549	164.44	453.4		85.45
10	578	150.44	394.1		86.78
11	607	139.55	348.3		87.99
12	673	162.30	365.2		90.09
13	801	131.38	248.4		92.30
14	932	426.30	692.8		94.06
15	1,376	649.56	715.1		97.43
16	5,380	13.59	3.8		107.81
17	8,046	8.01	1.5		110.85
18	10,380	75.00	10.9	Upper Marker	113.00
19	13,073	0.00	0.0		115.48

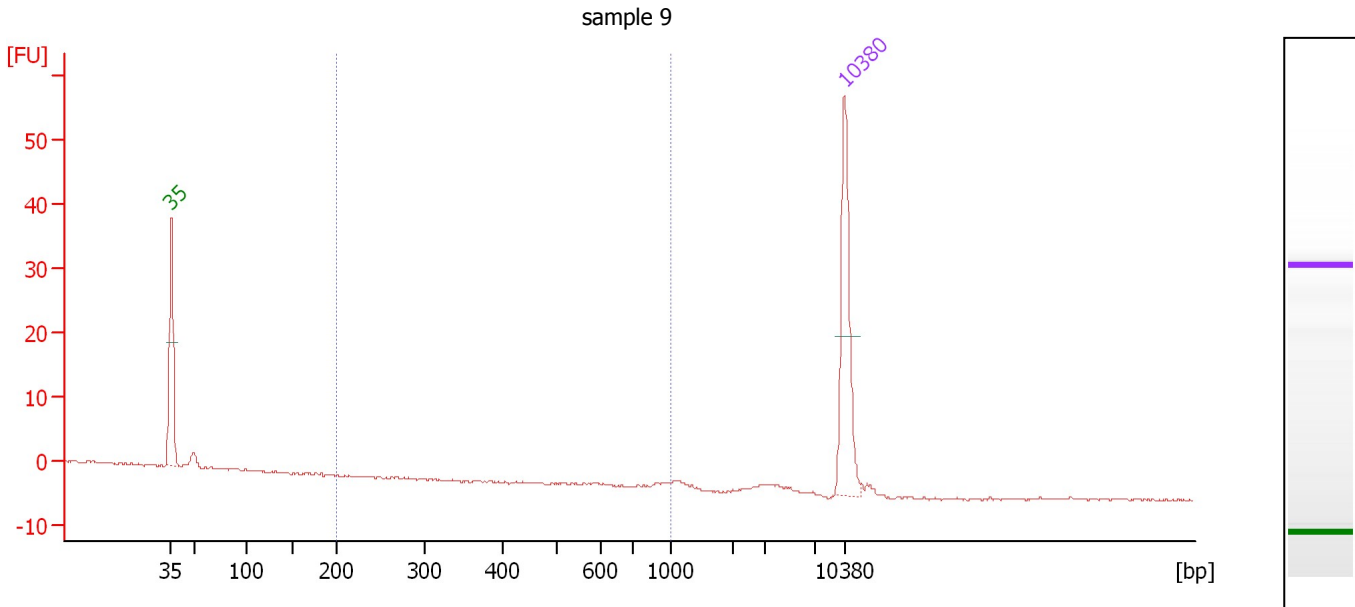
Region table for sample 8 : H

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	512	3,242.7	17,148.1	4,812.78	79	36.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
 Modified: 7/26/2016 12:35:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : sample 9

Number of peaks found: 0 Corr. Area 1: 0.0
 Noise: 0.2

Peak table for sample 9 : sample 9

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	10,380	75.00	10.9	Upper Marker	113.00

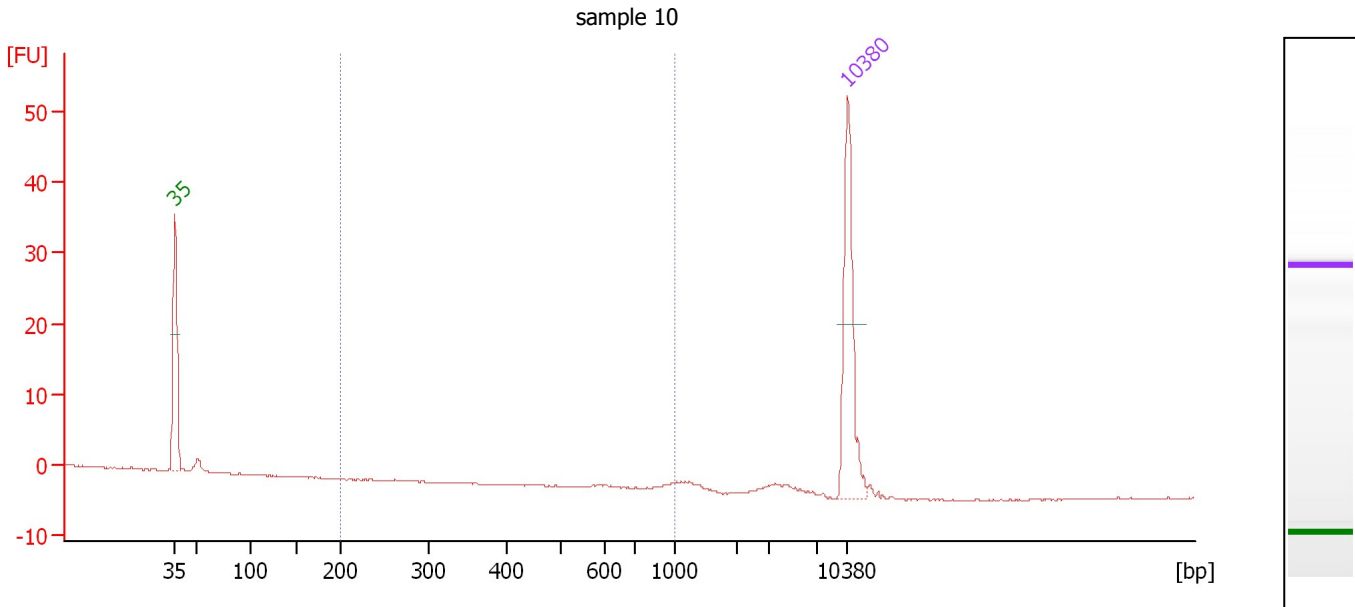
Region table for sample 9 : sample 9

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	0	0.0	0.0	0.00	0	0.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
 Modified: 7/26/2016 12:35:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : sample 10

Number of peaks found: 0 Corr. Area 1: 0.0
 Noise: 0.1

Peak table for sample 10 : sample 10

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	10,380	75.00	10.9	Upper Marker	113.00

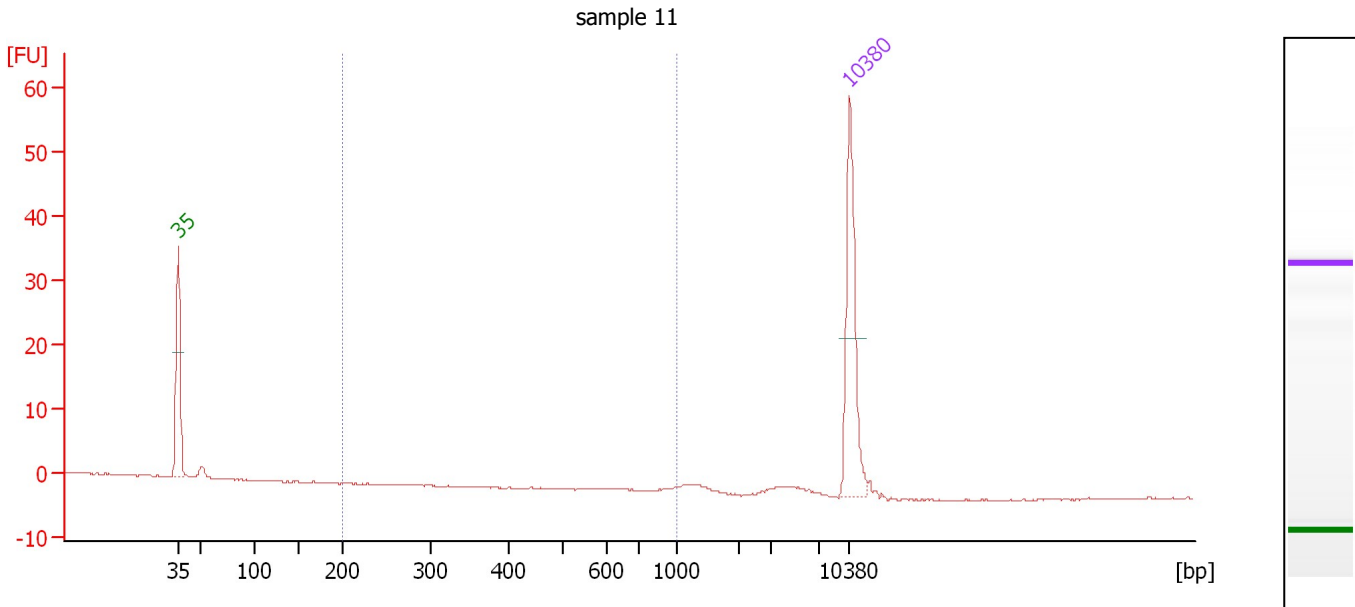
Region table for sample 10 : sample 10

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	986	0.0	0.0	0.00	0	0.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
 Modified: 7/26/2016 12:35:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : sample 11

Number of peaks found: 0 Corr. Area 1: 0.0
 Noise: 0.1

Peak table for sample 11 : sample 11

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	10,380	75.00	10.9	Upper Marker	113.00

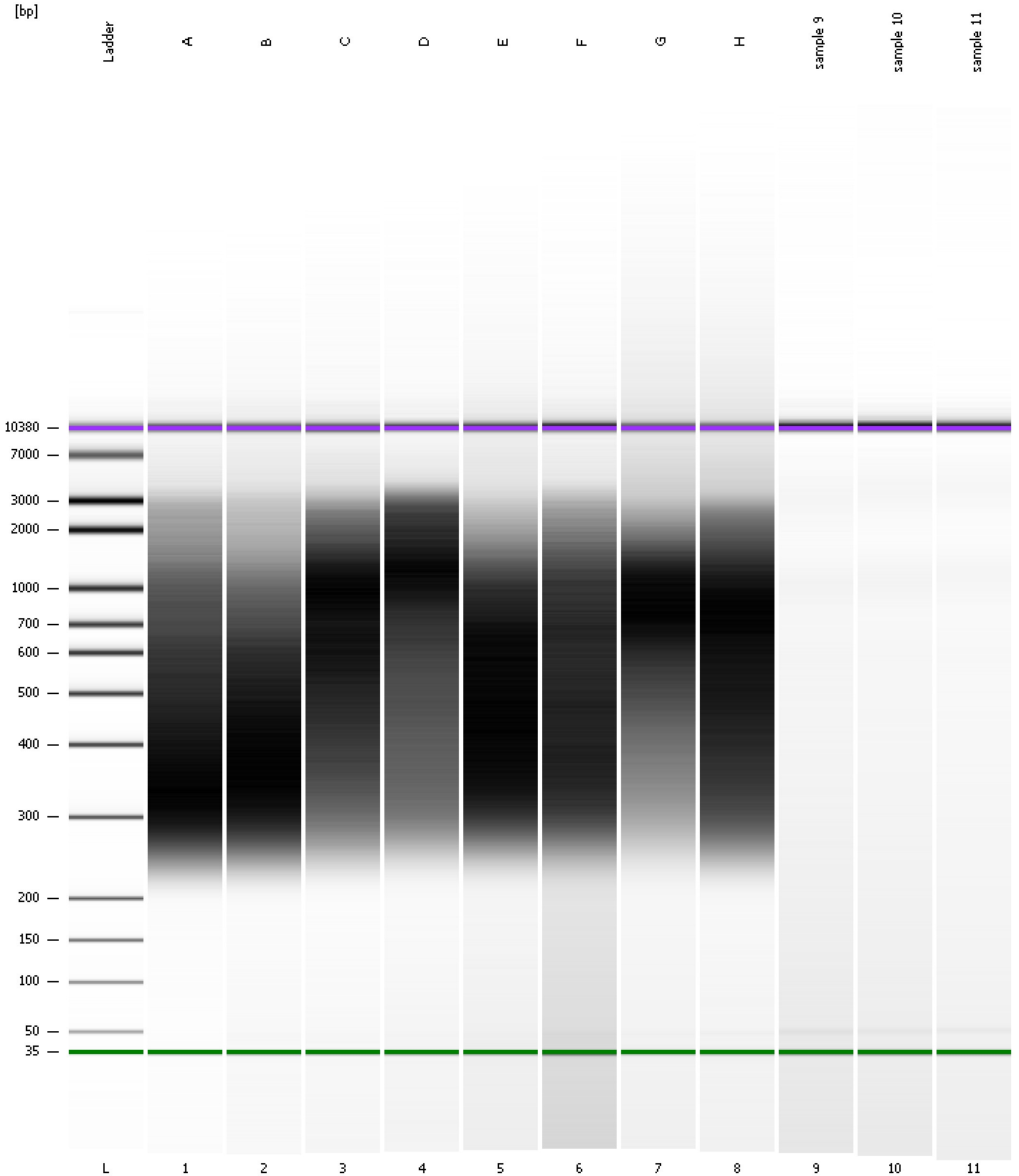
Region table for sample 11 : sample 11

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	0	0.0	0.0	0.00	0	0.0

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
Modified: 7/26/2016 12:35:05 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad

Created: 7/26/2016 11:53:45 AM
 Modified: 7/26/2016 12:35:05 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		7/26/2016 12:35:04 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-07-26\2016-07-26_001.xad)		Instrument	Run		7/26/2016 11:53:51 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		7/26/2016 11:53:51 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		7/26/2016 11:53:51 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		7/26/2016 11:53:51 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		7/26/2016 11:53:51 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		7/26/2016 11:53:51 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		7/26/2016 11:53:51 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1