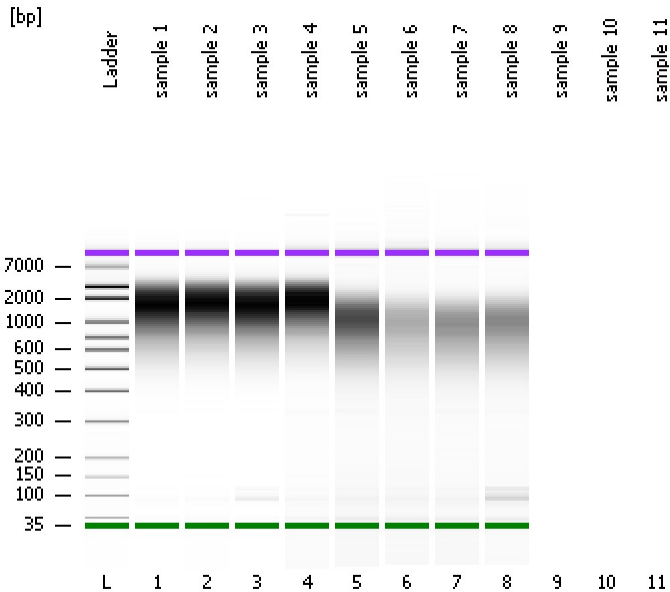


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
Modified: 9/27/2019 2:48:33 PM

Electrophoresis File Run Summary



Instrument Information:

Instrument Name: DE34903152 Firmware: C.01.069
Serial#: DE34903152 Type: G2938C

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

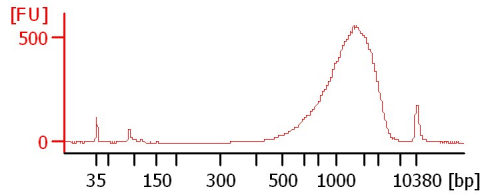
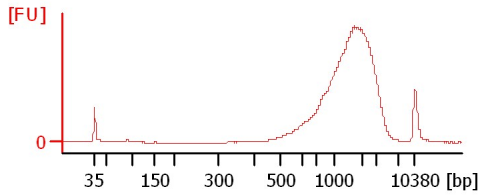
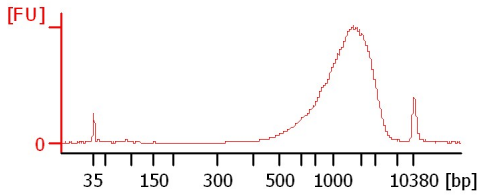
Chip Information:

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

sample 1

sample 2

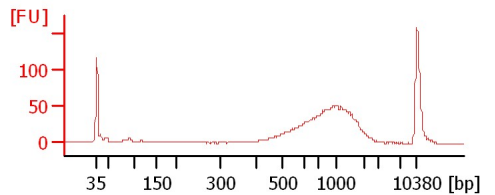
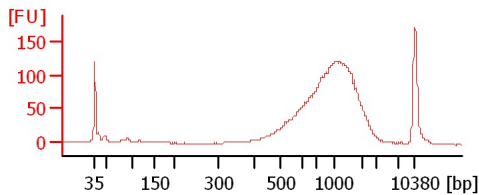
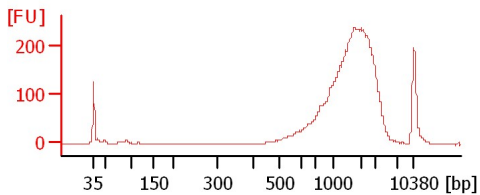
sample 3



sample 4

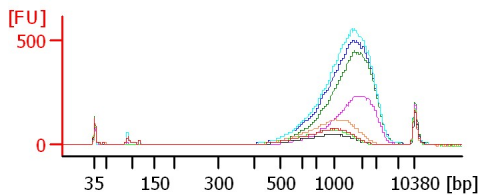
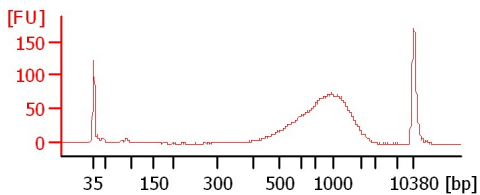
sample 5

sample 6



sample 7

sample 8



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
 Modified: 9/27/2019 2:48:33 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
sample 1		<input type="checkbox"/>	✓			
sample 2		<input type="checkbox"/>	✓			
sample 3		<input type="checkbox"/>	✓			
sample 4		<input type="checkbox"/>	✓			
sample 5		<input type="checkbox"/>	✓			
sample 6		<input type="checkbox"/>	✓			
sample 7		<input type="checkbox"/>	✓			
sample 8		<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\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
Modified: 9/27/2019 2:48:33 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

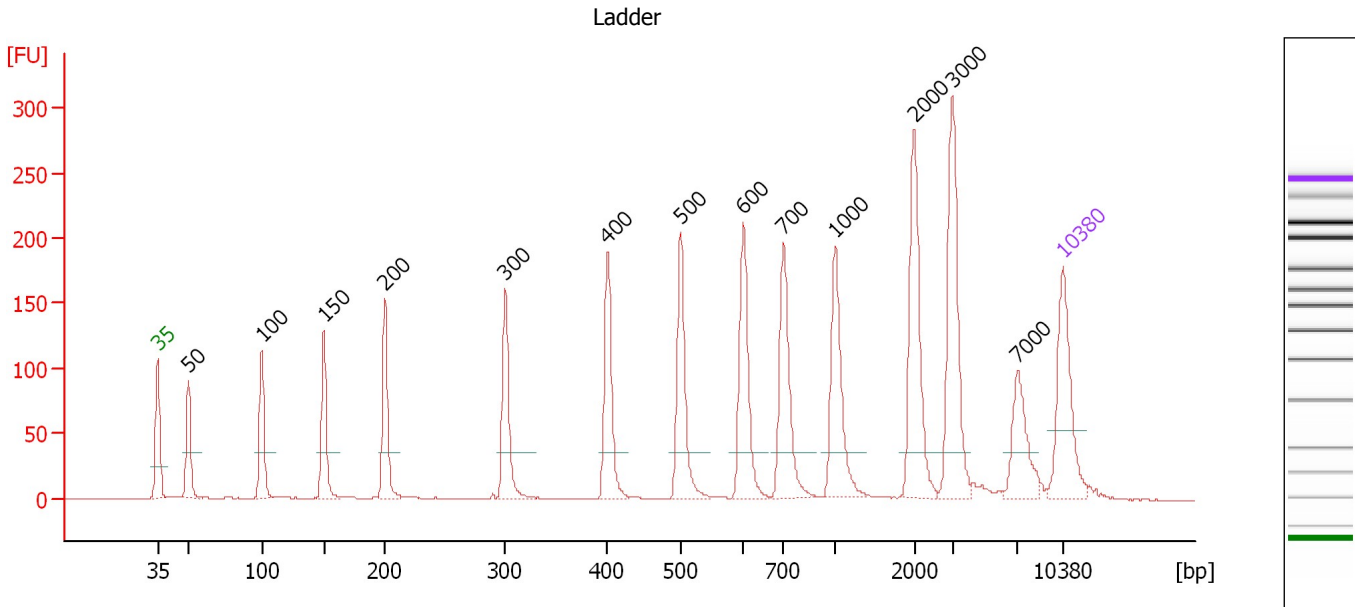
Ladder

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\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
 Modified: 9/27/2019 2:48:33 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

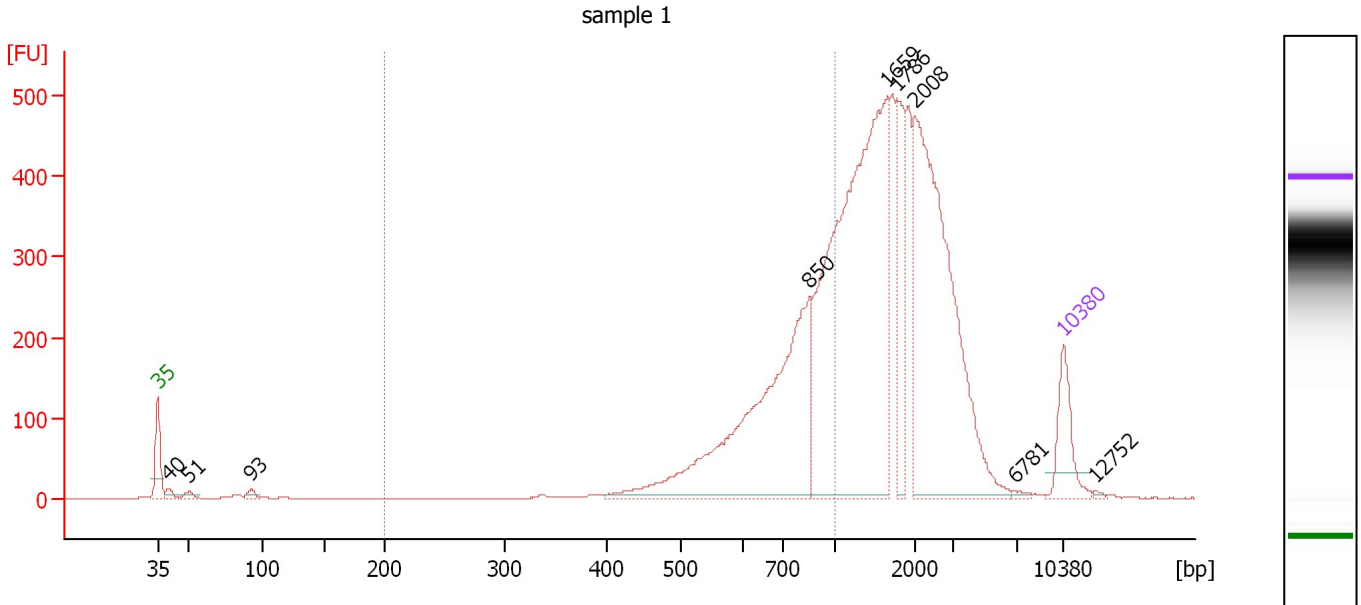
Peak table for Ladder

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	45.38
3	100	150.00	2,272.7	Ladder Peak	51.05
4	150	150.00	1,515.2	Ladder Peak	55.87
5	200	150.00	1,136.4	Ladder Peak	60.59
6	300	150.00	757.6	Ladder Peak	69.90
7	400	150.00	568.2	Ladder Peak	77.78
8	500	150.00	454.5	Ladder Peak	83.44
9	600	150.00	378.8	Ladder Peak	88.26
10	700	150.00	324.7	Ladder Peak	91.36
11	1,000	150.00	227.3	Ladder Peak	95.41
12	2,000	150.00	113.6	Ladder Peak	101.48
13	3,000	150.00	75.8	Ladder Peak	104.45
14	7,000	150.00	32.5	Ladder Peak	109.49
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
 Modified: 9/27/2019 2:48:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : sample 1

Number of peaks found: 9 Corr. Area 1: 2,032.9
 Noise: 0.2

Peak table for sample 1 : sample 1

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	40	14.21	532.2		43.87
3	51	13.22	395.1		45.46
4	93	13.61	222.9		50.21
5	850	636.51	1,134.1		93.39
6	1,659	944.60	862.9		99.41
7	1,786	126.33	107.2		100.19
8	2,008	625.92	472.2		101.51
9	6,781	4.63	1.0		109.21
10	10,380	75.00	10.9	Upper Marker	113.00
11	12,752	0.00	0.0		115.46

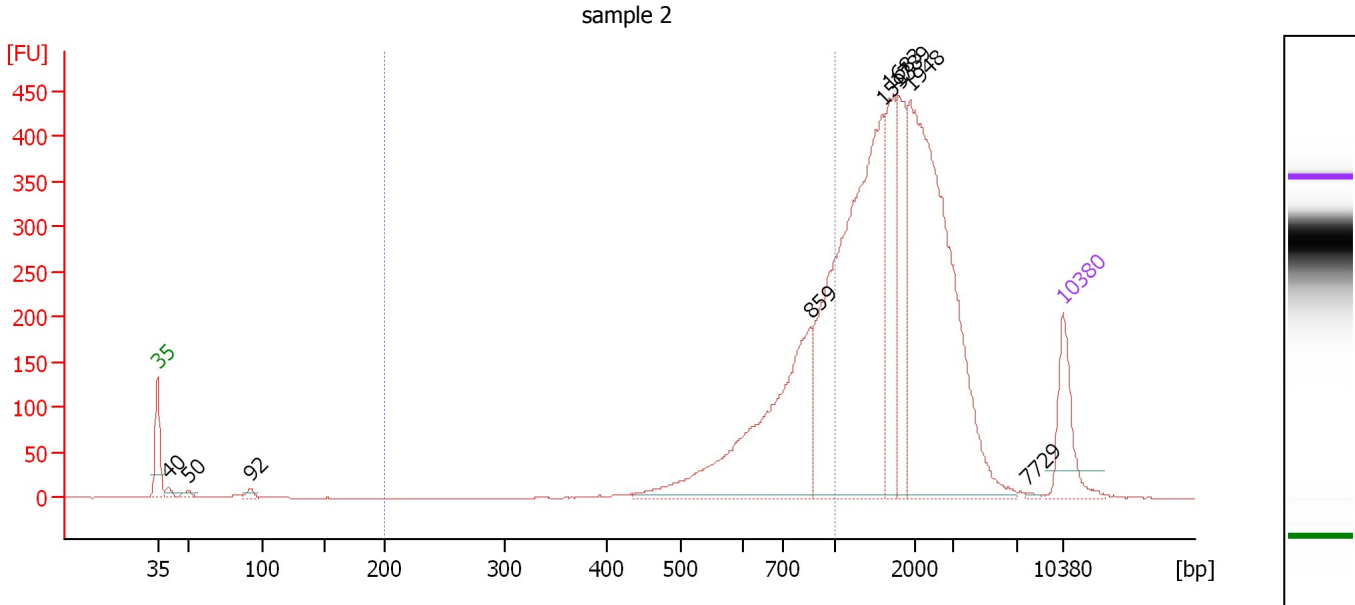
Region table for sample 1 : sample 1

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	743	959.86	2,032.9	2,088.3	31	21.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
 Modified: 9/27/2019 2:48:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : sample 2

Number of peaks found: 9 Corr. Area 1: 1,449.8
 Noise: 0.3

Peak table for sample 2 : sample 2

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	40	11.75	443.0		43.83
3	50	9.83	295.5		45.43
4	92	10.27	169.0		50.16
5	859	396.81	700.1		93.50
6	1,593	647.67	616.2		99.01
7	1,683	140.57	126.5		99.56
8	1,789	138.45	117.3		100.20
9	1,948	589.69	458.8		101.17
10	7,729	2.20	0.4		110.25
11	10,380	75.00	10.9	Upper Marker	113.00

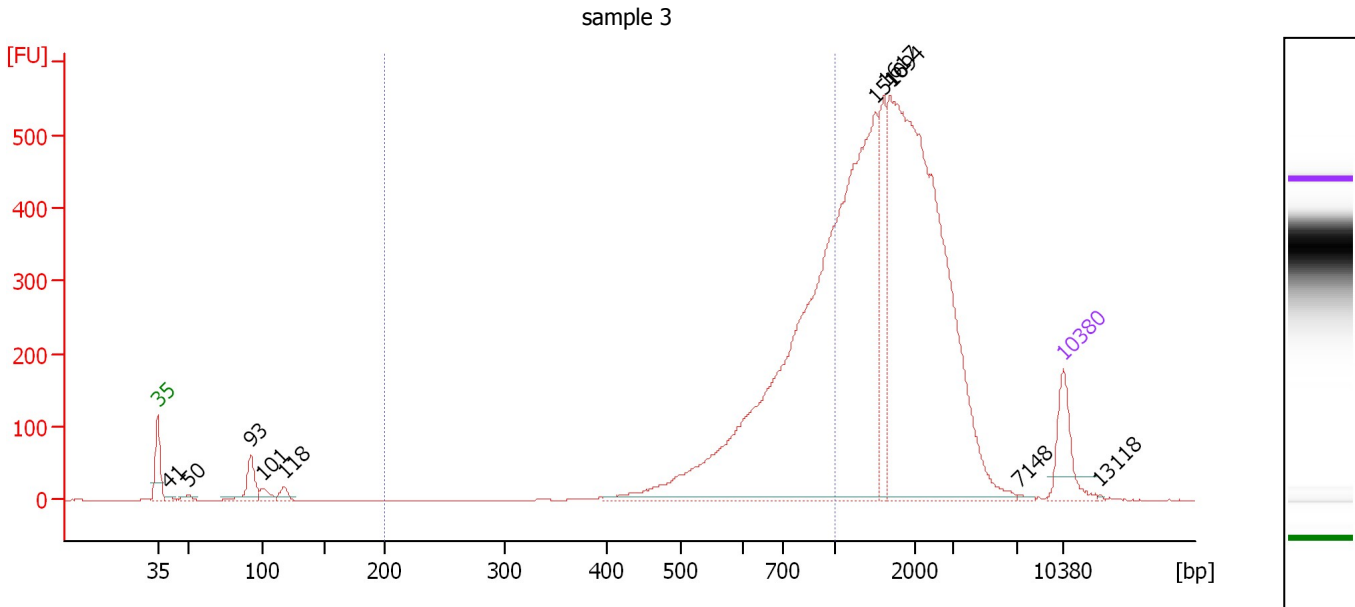
Region table for sample 2 : sample 2

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	760	609.75	1,449.8	1,285.2	27	20.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
 Modified: 9/27/2019 2:48:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : sample 3

Number of peaks found: 10 Corr. Area 1: 2,219.1
 Noise: 0.2

Peak table for sample 3 : sample 3

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	41	6.69	249.9		43.89
3	50	10.72	325.1		45.38
4	93	77.22	1,262.2		50.22
5	101	23.47	353.5		51.11
6	118	21.48	275.8		52.79
7	1,510	1,539.77	1,545.3		98.51
8	1,617	163.24	152.9		99.16
9	1,694	1,174.95	1,050.9		99.62
10	7,148	3.03	0.6		109.64
11	10,380	75.00	10.9	Upper Marker	113.00
12	13,118	0.00	0.0		115.84

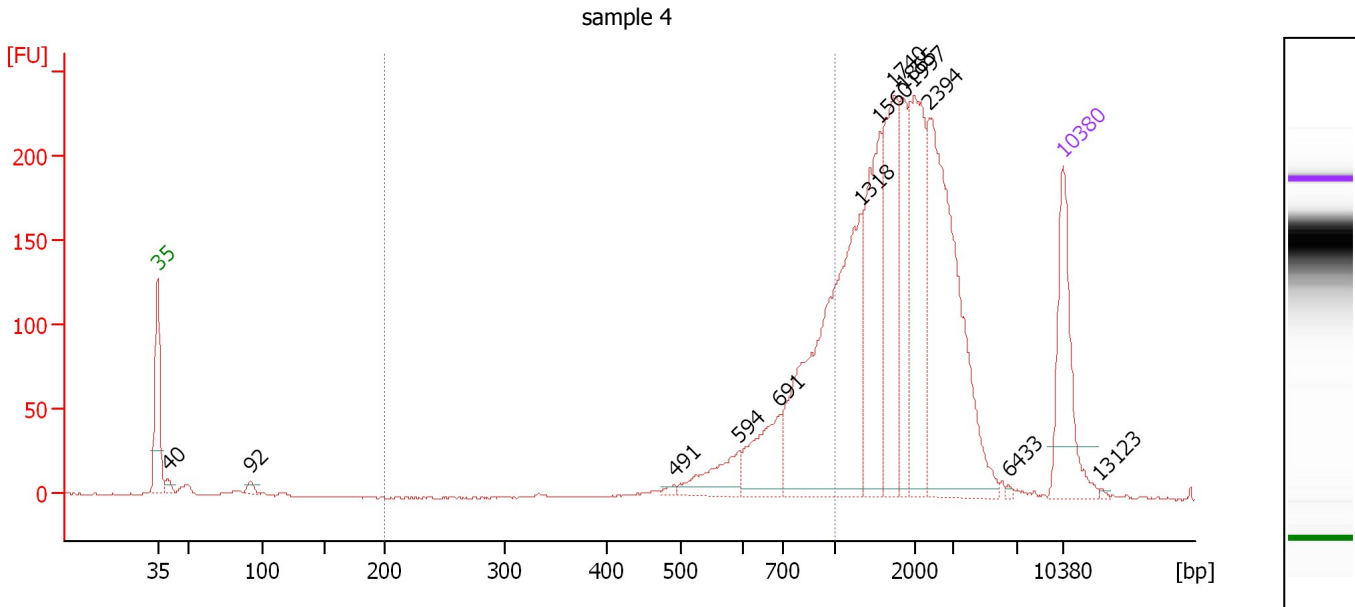
Region table for sample 3 : sample 3

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	752	1,089.30	2,219.1	2,328.7	31	20.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
 Modified: 9/27/2019 2:48:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : sample 4

Number of peaks found: 13 Corr. Area 1: 627.9
 Noise: 0.3

Peak table for sample 4 : sample 4

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	40	7.81	295.5		43.81
3	92	6.72	110.5		50.17
4	491	3.52	10.9		82.91
5	594	39.00	99.4		87.99
6	691	62.81	137.8		91.07
7	1,318	276.81	318.3		97.34
8	1,560	118.80	115.4		98.81
9	1,740	114.31	99.6		99.90
10	1,865	59.23	48.1		100.66
11	1,997	118.72	90.1		101.47
12	2,394	232.08	146.9		102.65
13	6,433	1.40	0.3		108.78
14	10,380	75.00	10.9	Upper Marker	113.00
15	13,123	0.00	0.0		115.85

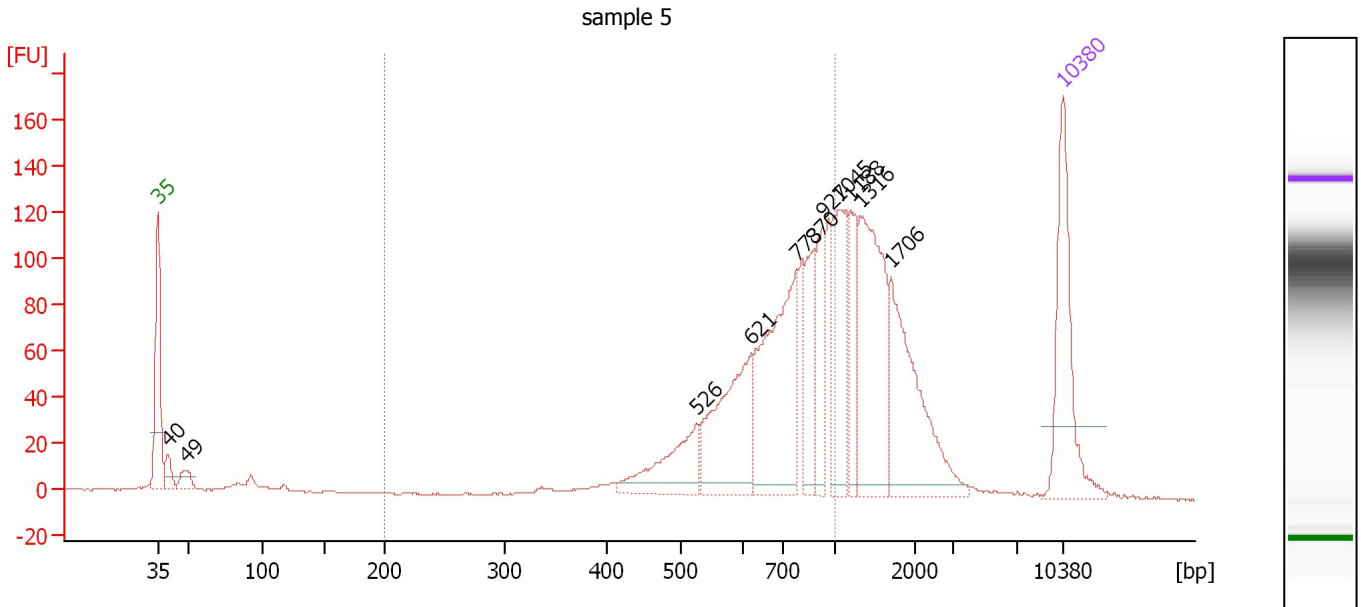
Region table for sample 4 : sample 4

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	767	283.05	627.9	588.0	23	19.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
 Modified: 9/27/2019 2:48:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : sample 5

Number of peaks found: 11 Corr. Area 1: 1,002.4
 Noise: 0.2

Peak table for sample 5 : sample 5

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	40	17.16	652.1		43.77
3	49	14.12	439.3		45.18
4	526	62.86	181.0		84.70
5	621	104.94	256.0		88.91
6	773	150.25	294.6		92.34
7	870	48.19	84.0		93.65
8	927	44.78	73.2		94.42
9	1,045	72.81	105.6		95.68
10	1,188	39.34	50.2		96.55
11	1,316	123.04	141.7		97.33
12	1,706	106.59	94.7		99.70
13	10,380	75.00	10.9	Upper Marker	113.00

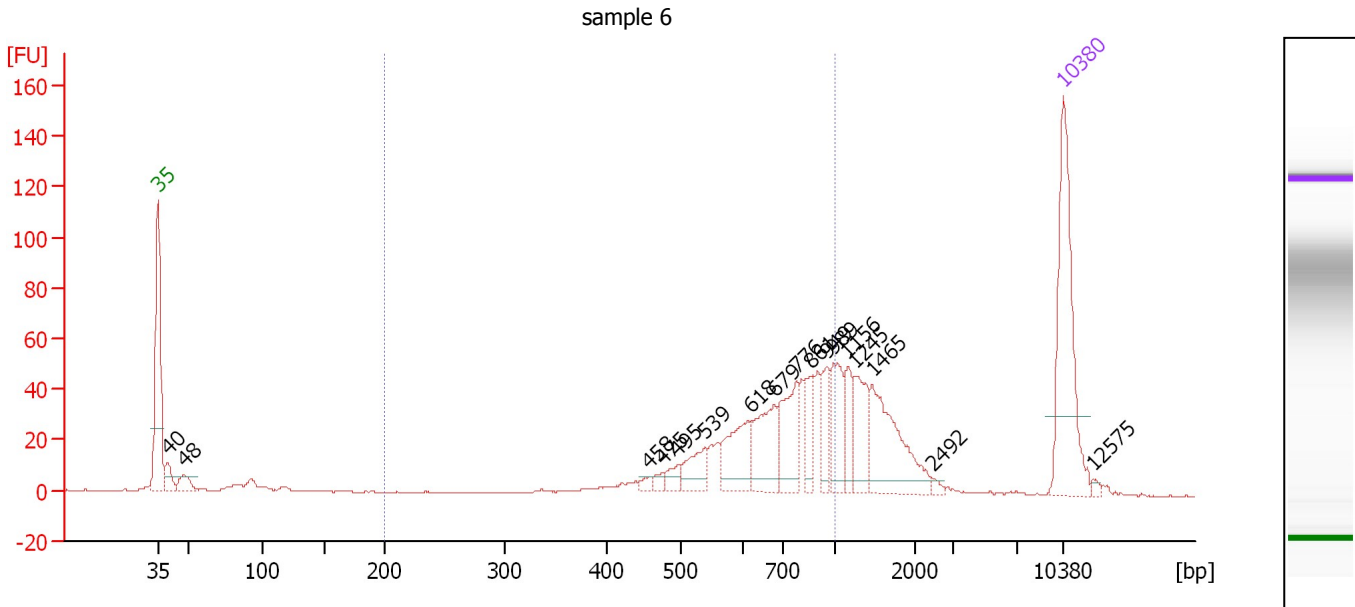
Region table for sample 5 : sample 5

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	712	512.90	1,002.4	1,173.2	54	22.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
 Modified: 9/27/2019 2:48:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : sample 6

Number of peaks found: 17 Corr. Area 1: 491.9
 Noise: 0.2

Peak table for sample 6 : sample 6

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	40	13.86	525.1		43.79
3	48	12.45	395.0		45.03
4	458	3.52	11.7		81.05
5	475	4.34	13.8		82.04
6	495	7.41	22.7		83.18
7	539	20.12	56.6		85.31
8	618	36.64	89.8		88.82
9	679	40.85	91.2		90.71
10	776	38.94	76.0		92.39
11	861	17.31	30.5		93.53
12	949	17.67	28.2		94.72
13	989	28.27	43.3		95.26
14	1,156	18.88	24.8		96.35
15	1,245	27.44	33.4		96.90
16	1,465	55.96	57.9		98.24
17	2,492	2.11	1.3		102.94
18	10,380	75.00	10.9	Upper Marker	113.00
19	12,575	0.00	0.0		115.28

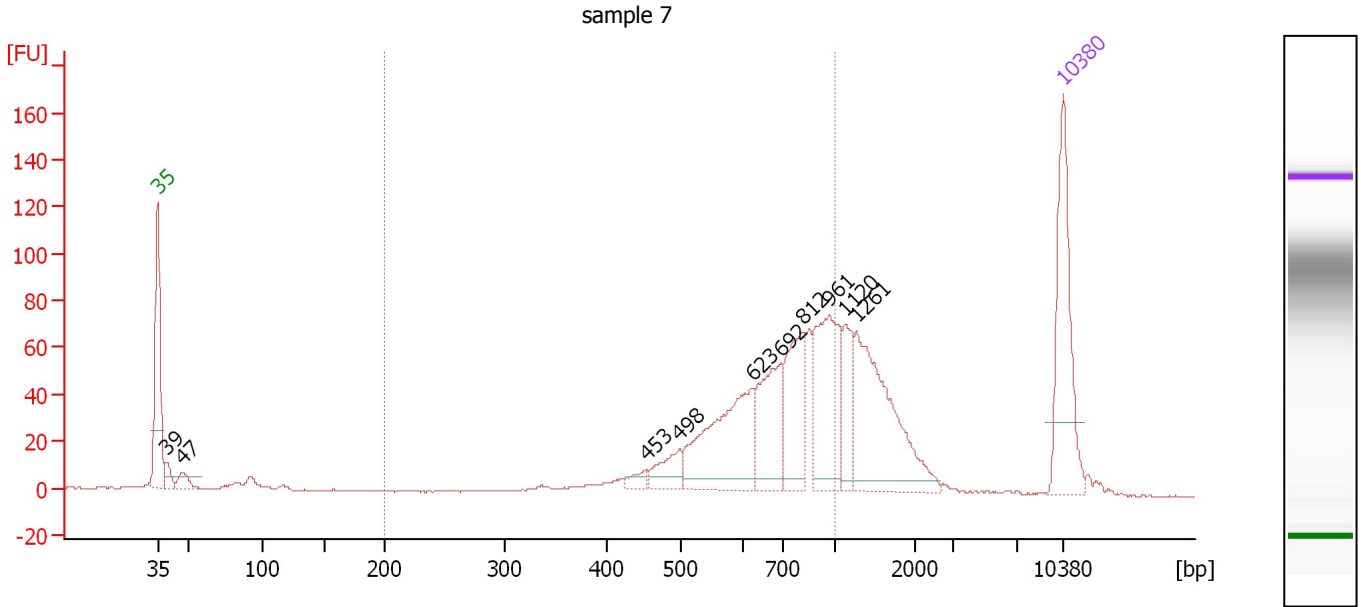
Region table for sample 6 : sample 6

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	693	280.27	491.9	671.3	58	24.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
 Modified: 9/27/2019 2:48:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : sample 7

Number of peaks found: 10 Corr. Area 1: 718.7
 Noise: 0.2

Peak table for sample 7 : sample 7

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	39	12.88	494.9		43.70
3	47	13.68	441.1		44.91
4	453	8.02	26.8		80.78
5	498	23.79	72.3		83.34
6	623	106.58	259.3		88.96
7	692	68.14	149.1		91.12
8	812	66.34	123.7		92.88
9	961	89.42	141.0		94.89
10	1,120	33.15	44.8		96.14
11	1,261	118.62	142.6		96.99
12	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 7 : sample 7

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	696	406.67	718.7	956.6	63	23.3

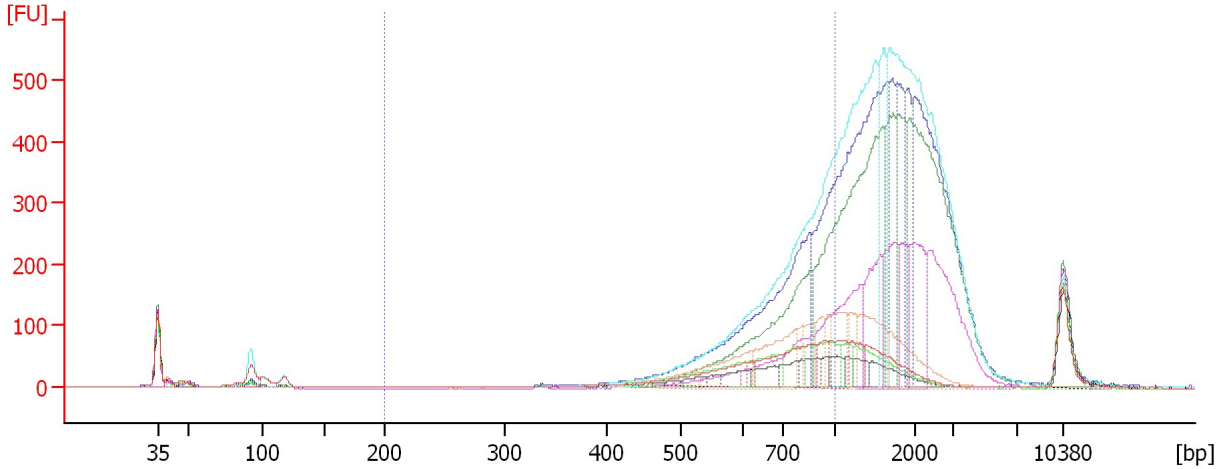
Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
 Modified: 9/27/2019 2:48:33 PM

Electropherogram Summary Continued ...

sample 8

● 8: sample 8	● 2: sample 2	● 4: sample 4	● 6: sample 6
● 1: sample 1	● 3: sample 3	● 5: sample 5	● 7: sample 7



Overall Results for sample 8 : sample 8

Number of peaks found: 12 Corr. Area 1: 683.0
 Noise: 0.2

Peak table for sample 8 : sample 8

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	40	13.44	511.9		43.76
3	46	17.98	586.1		44.82
4	84	9.43	171.1		49.18
5	93	51.68	842.7		50.25
6	101	24.91	373.2		51.16
7	118	20.77	266.8		52.78
8	452	4.33	14.5		80.71
9	605	101.06	253.0		88.42
10	861	155.06	272.8		93.54
11	929	40.22	65.6		94.45
12	1,209	44.30	55.5		96.68
13	1,401	106.94	115.7		97.84
14	10,380	75.00	10.9	Upper Marker	113.00

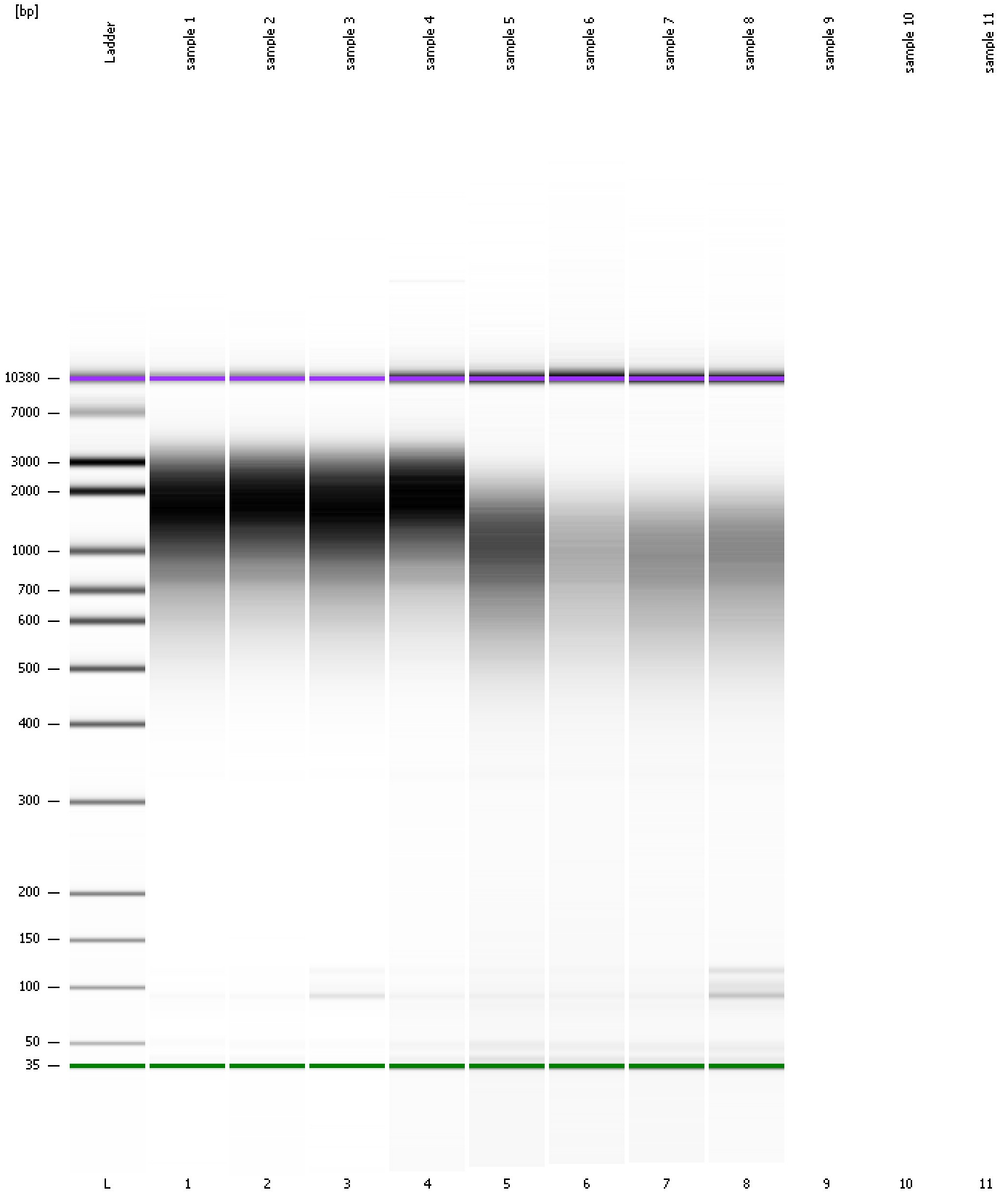
Region table for sample 8 : sample 8

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	703	400.66	683.0	933.0	54	23.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
Modified: 9/27/2019 2:48:33 PM

Gel Image

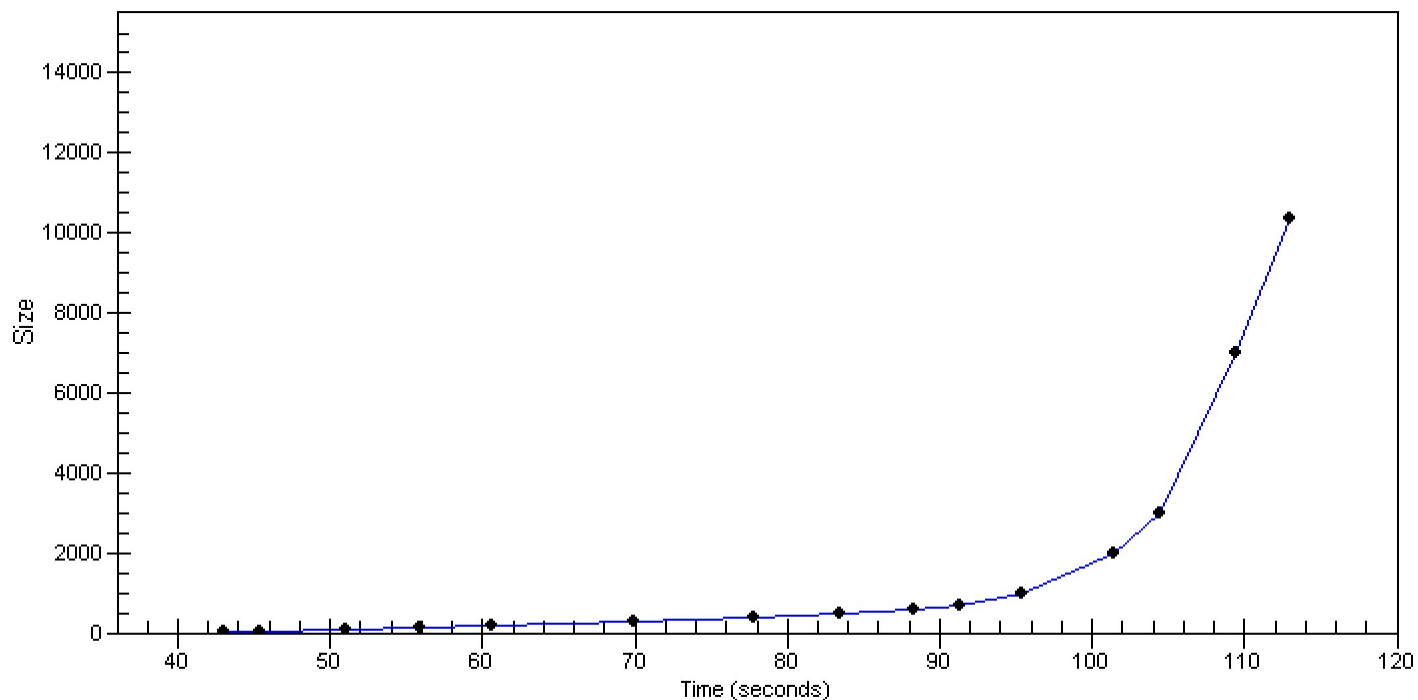


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
Modified: 9/27/2019 2:48:33 PM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
Modified: 9/27/2019 2:48:33 PM

Invalid Samples

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.

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-09-27\2019-09-27_001.xad

Created: 9/27/2019 2:15:50 PM
 Modified: 9/27/2019 2:48:33 PM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 9)		Instrument	Run		9/27/2019 2:48:32 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\2019-09-27\2019-09-27_001.xad)		Instrument	Run		9/27/2019 2:15:56 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		9/27/2019 2:15:56 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		9/27/2019 2:15:56 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		9/27/2019 2:15:56 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		9/27/2019 2:15:56 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		9/27/2019 2:15:56 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		9/27/2019 2:15:56 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1