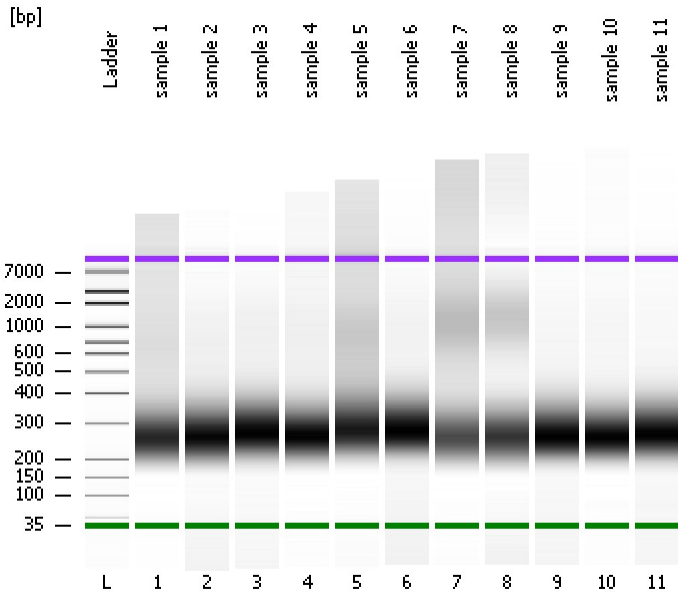


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
Modified: 8/2/2019 3:44:06 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

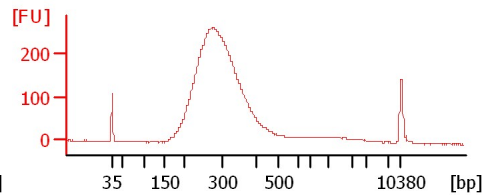
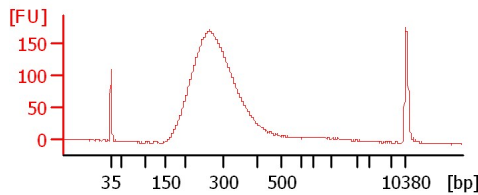
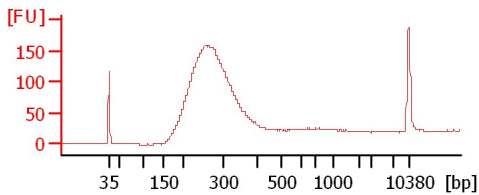
Chip Information:

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

sample 1

sample 2

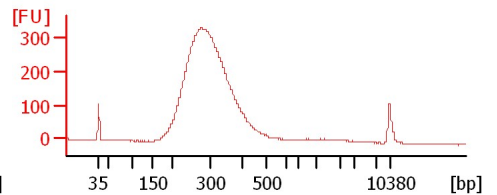
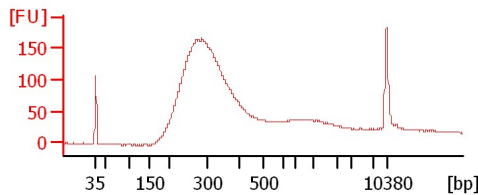
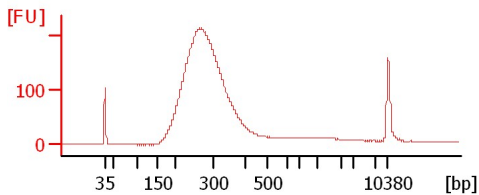
sample 3



sample 4

sample 5

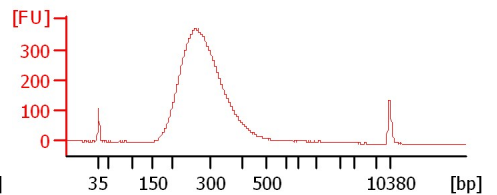
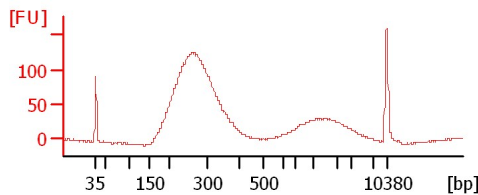
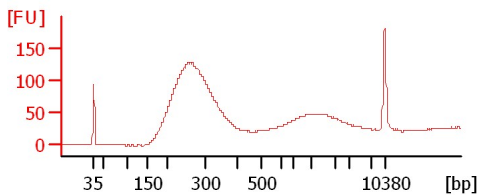
sample 6



sample 7

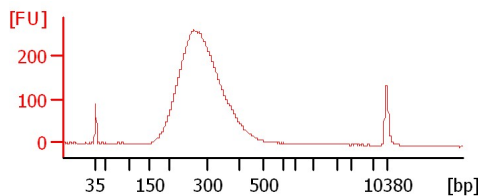
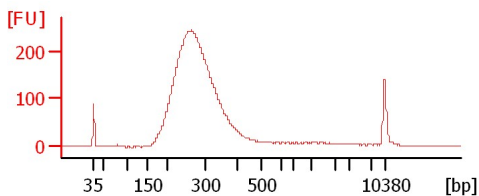
sample 8

sample 9



sample 10

sample 11



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
Modified: 8/2/2019 3:44:06 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\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
Modified: 8/2/2019 3:44:06 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

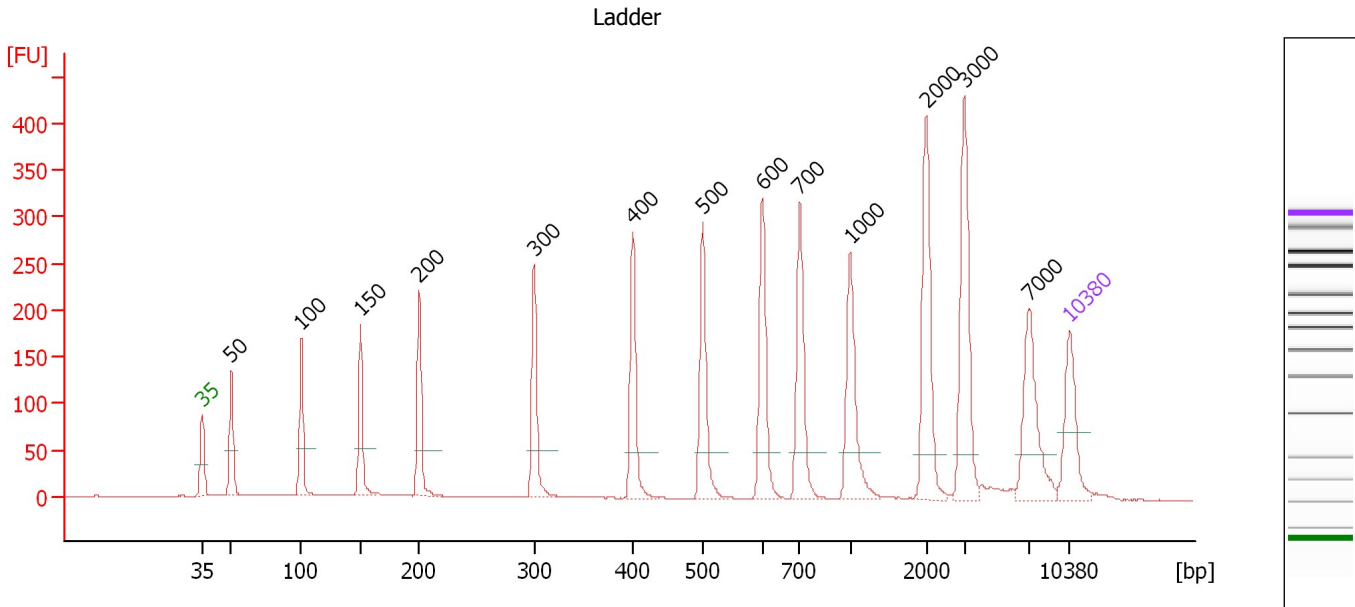
Ladder

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\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
 Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.3

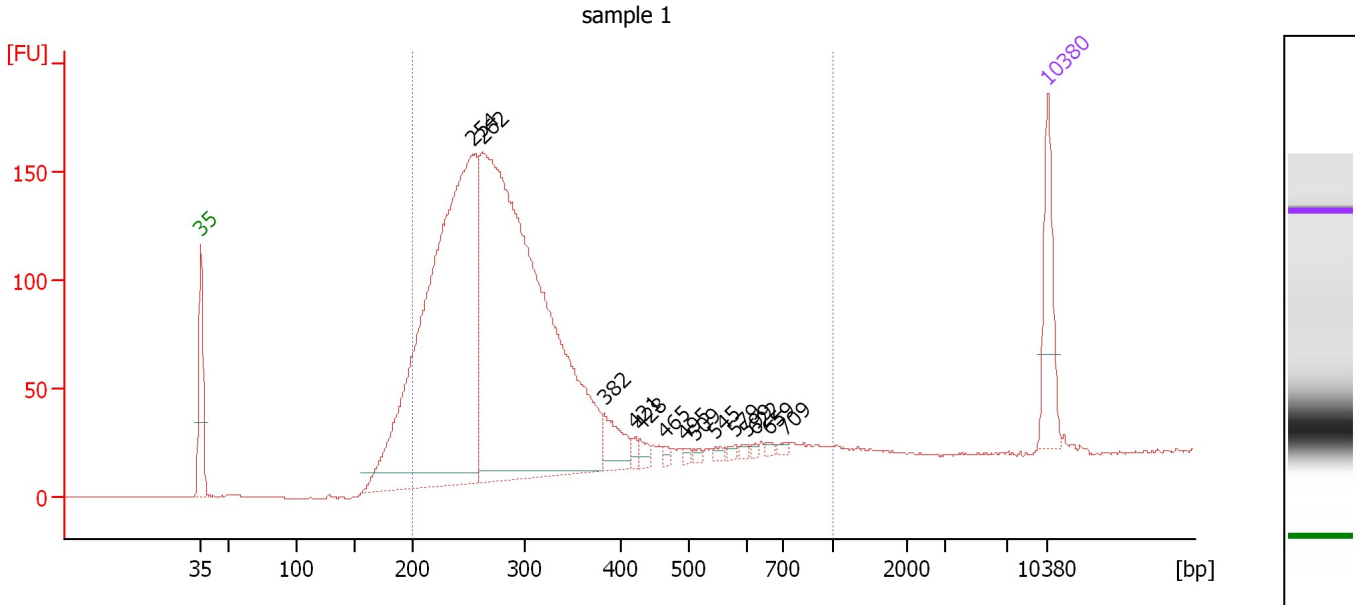
Peak table for Ladder

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	45.32
3	100	150.00	2,272.7	Ladder Peak	50.96
4	150	150.00	1,515.2	Ladder Peak	55.78
5	200	150.00	1,136.4	Ladder Peak	60.51
6	300	150.00	757.6	Ladder Peak	69.76
7	400	150.00	568.2	Ladder Peak	77.72
8	500	150.00	454.5	Ladder Peak	83.36
9	600	150.00	378.8	Ladder Peak	88.18
10	700	150.00	324.7	Ladder Peak	91.19
11	1,000	150.00	227.3	Ladder Peak	95.27
12	2,000	150.00	113.6	Ladder Peak	101.38
13	3,000	150.00	75.8	Ladder Peak	104.48
14	7,000	150.00	32.5	Ladder Peak	109.69
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
 Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : sample 1

Number of peaks found: 14 Corr. Area 1: 2,741.3
 Noise: 0.3

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	254	1,164.80	6,942.5		65.53
3	262	1,392.05	8,040.7		66.27
4	382	51.44	203.8		76.32
5	421	8.65	31.2		78.88
6	428	12.00	42.4		79.32
7	465	5.95	19.4		81.39
8	495	4.66	14.3		83.07
9	509	5.20	15.5		83.77
10	545	5.29	14.7		85.54
11	579	5.15	13.5		87.17
12	599	4.77	12.1		88.14
13	622	3.23	7.9		88.84
14	659	4.49	10.3		89.95
15	709	3.88	8.3		91.31
16	10,380	75.00	10.9	Upper Marker	113.00

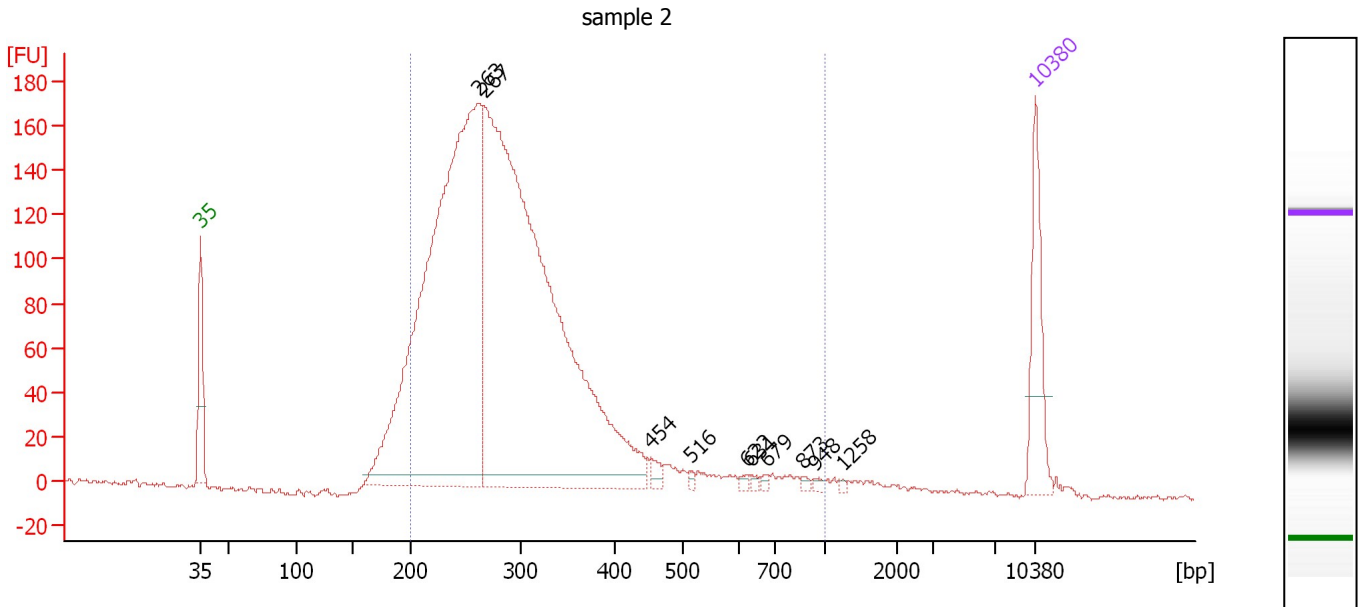
Region table for sample 1 : sample 1

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	312	2,557.03	2,741.3	13,835.0	92	38.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
 Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : sample 2

Number of peaks found: 10 Corr. Area 1: 3,126.2
 Noise: 1.2

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	263	1,173.52	6,770.3		66.30
3	267	1,411.30	7,995.0		66.75
4	454	11.11	37.1		80.75
5	516	4.16	12.2		84.15
6	622	4.60	11.2		88.85
7	631	3.93	9.4		89.12
8	679	3.91	8.7		90.55
9	873	3.56	6.2		93.54
10	948	3.77	6.0		94.57
11	1,258	2.12	2.6		96.85
12	10,380	75.00	10.9	Upper Marker	113.00

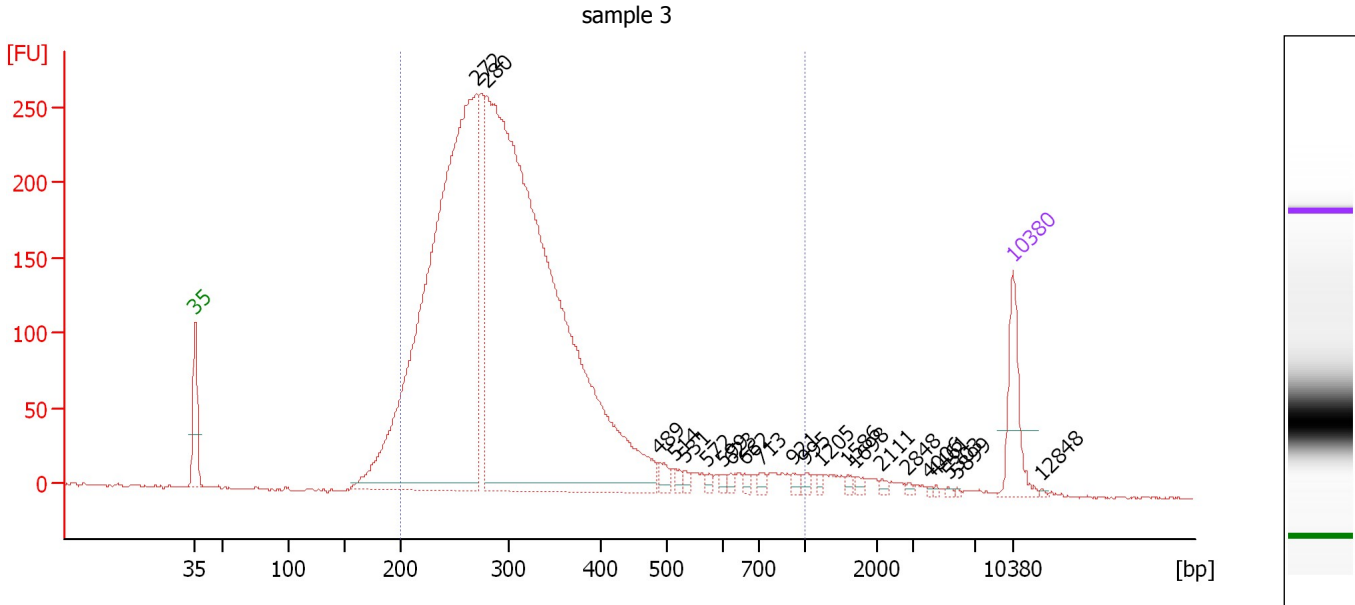
Region table for sample 2 : sample 2

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	304	2,505.79	3,126.2	13,541.5	92	32.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
 Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : sample 3

Number of peaks found: 22 Corr. Area 1: 4,865.6
 Noise: 0.8

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	272	1,751.32	9,767.8		67.14
3	280	2,338.47	12,651.9		67.92
4	489	18.16	56.2		82.75
5	514	10.67	31.5		84.02
6	531	7.28	20.8		84.84
7	572	7.58	20.1		86.81
8	599	5.82	14.7		88.13
9	623	6.53	15.9		88.86
10	662	8.01	18.3		90.05
11	713	9.29	19.7		91.37
12	921	9.38	15.4		94.20
13	995	6.71	10.2		95.20
14	1,205	5.46	6.9		96.53
15	1,586	5.15	4.9		98.85
16	1,698	6.33	5.6		99.54
17	2,111	4.49	3.2		101.73
18	2,848	3.24	1.7		104.01
19	4,006	1.91	0.7		105.79
20	4,461	1.89	0.6		106.38
21	5,303	1.99	0.6		107.48
22	5,899	1.55	0.4		108.25
23	10,380	75.00	10.9	Upper Marker	113.00
24	12,848	0.00	0.0		115.42

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...

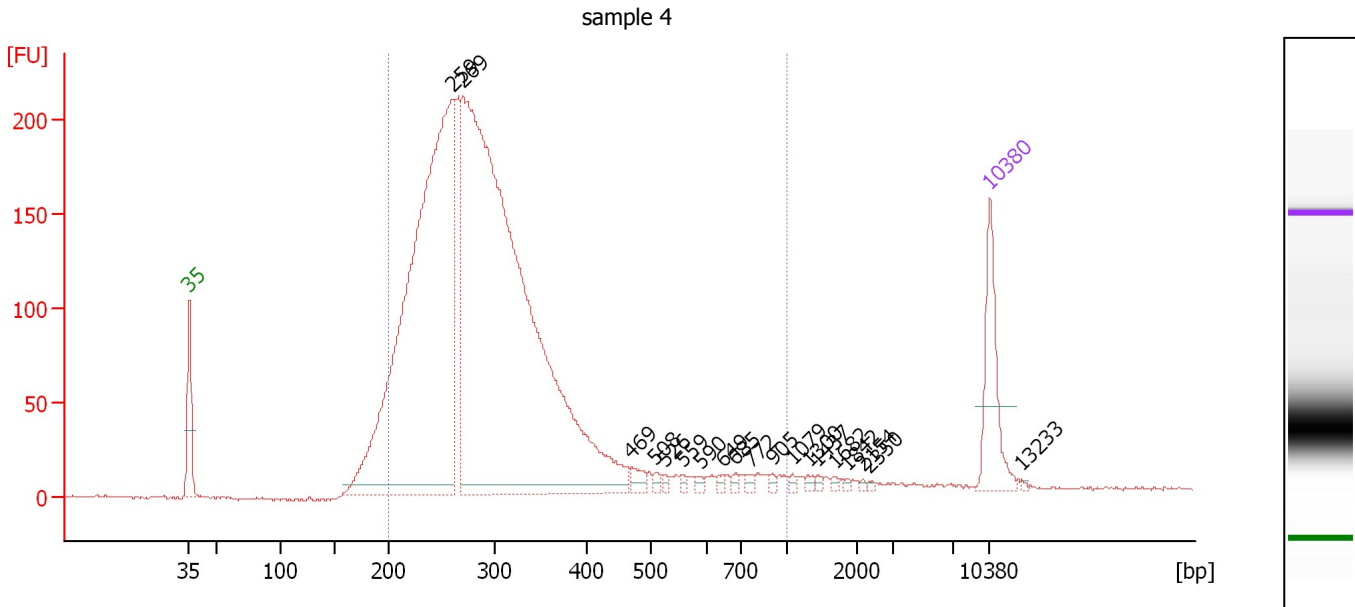
... Region table for sample 3 : sample 3

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	315	4,100.02	4,865.6	21,394.0	 94	32.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
 Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : sample 4

Number of peaks found: 19 Corr. Area 1: 3,644.3
 Noise: 0.8

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	259	1,190.96	6,964.9		65.98
3	269	1,661.32	9,351.9		66.91
4	469	14.90	48.1		81.64
5	508	6.55	19.5		83.73
6	526	4.46	12.8		84.62
7	559	4.48	12.1		86.20
8	590	5.39	13.8		87.69
9	649	4.75	11.1		89.65
10	685	4.72	10.4		90.72
11	772	5.06	9.9		92.17
12	905	3.53	5.9		93.99
13	1,079	3.60	5.1		95.76
14	1,300	3.71	4.3		97.11
15	1,437	2.73	2.9		97.95
16	1,682	2.73	2.5		99.44
17	1,842	2.20	1.8		100.42
18	2,154	1.81	1.3		101.86
19	2,350	1.76	1.1		102.47
20	10,380	75.00	10.9	Upper Marker	113.00
21	13,233	0.00	0.0		115.80

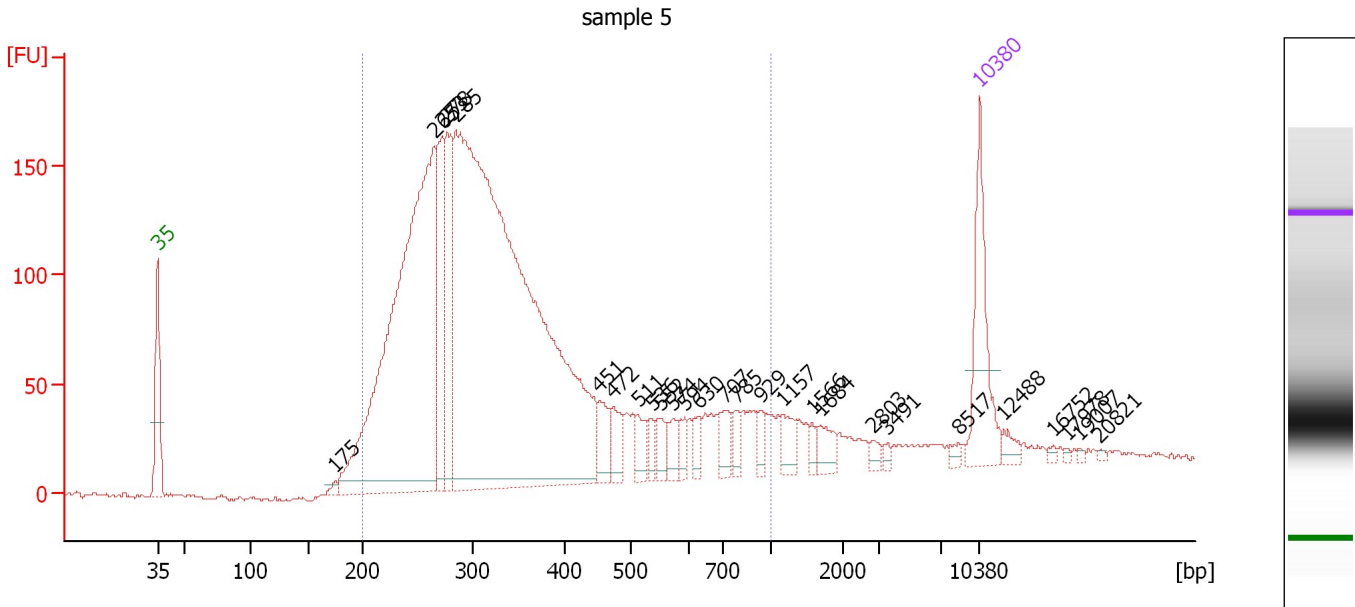
Region table for sample 4 : sample 4

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	304	2,818.18	3,644.3	15,187.4	93	32.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
 Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : sample 5

Number of peaks found: 27 Corr. Area 1: 3,369.9
 Noise: 1.0

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	175	6.14	53.2		58.13
3	265	730.99	4,176.3		66.54
4	273	120.09	666.5		67.26
5	278	128.97	703.8		67.69
6	285	1,311.97	6,963.8		68.42
7	451	30.45	102.2		80.62
8	472	28.58	91.8		81.77
9	511	22.14	65.6		83.89
10	536	12.88	36.4		85.09
11	552	17.18	47.2		85.86
12	574	17.26	45.6		86.91
13	594	14.15	36.1		87.87
14	630	14.35	34.5		89.07
15	707	21.24	45.5		91.28
16	785	13.03	25.2		92.34
17	929	11.61	18.9		94.31
18	1,157	18.68	24.5		96.23
19	1,566	7.00	6.8		98.73
20	1,684	16.93	15.2		99.45
21	2,803	6.43	3.5		103.87
22	3,491	3.85	1.7		105.12
23	8,517	4.70	0.8		111.17
24	10,380	75.00	10.9	Upper Marker	113.00
25	12,488	0.00	0.0		115.07

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...**... Peak table for sample 5 : sample 5**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	16,752	0.00	0.0		119.25
27	17,978	0.00	0.0		120.45
28	19,007	0.00	0.0		121.46
29	20,821	0.00	0.0		123.23

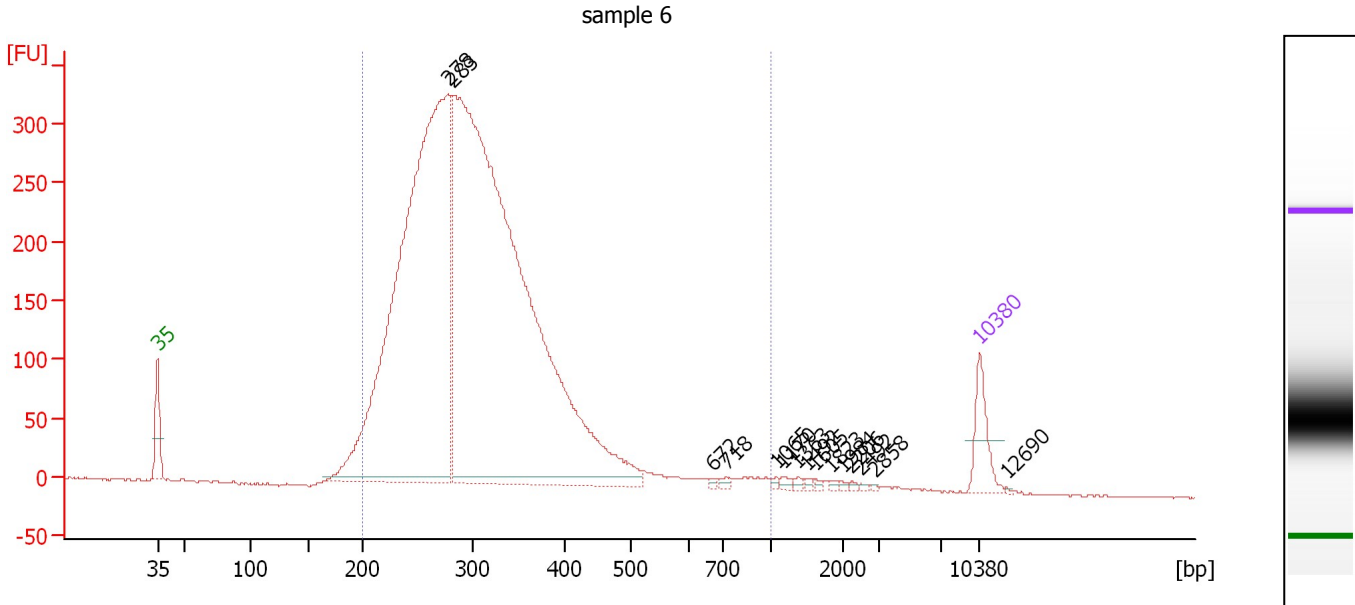
Region table for sample 5 : sample 5

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	360	2,344.97	3,369.9	11,453.9	 90	41.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
 Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : sample 6

Number of peaks found: 15 Corr. Area 1: 5,926.2
 Noise: 0.7

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	278	2,497.80	13,594.9		67.76
3	283	3,487.72	18,661.8		68.20
4	672	5.65	12.7		90.36
5	718	7.77	16.4		91.44
6	1,065	5.19	7.4		95.67
7	1,170	8.14	10.5		96.31
8	1,363	5.60	6.2		97.49
9	1,492	4.92	5.0		98.28
10	1,605	3.86	3.6		98.97
11	1,823	4.80	4.0		100.30
12	1,984	3.28	2.5		101.28
13	2,206	3.93	2.7		102.02
14	2,492	2.83	1.7		102.91
15	2,858	2.33	1.2		104.04
16	10,380	75.00	10.9	Upper Marker	113.00
17	12,690	0.00	0.0		115.26

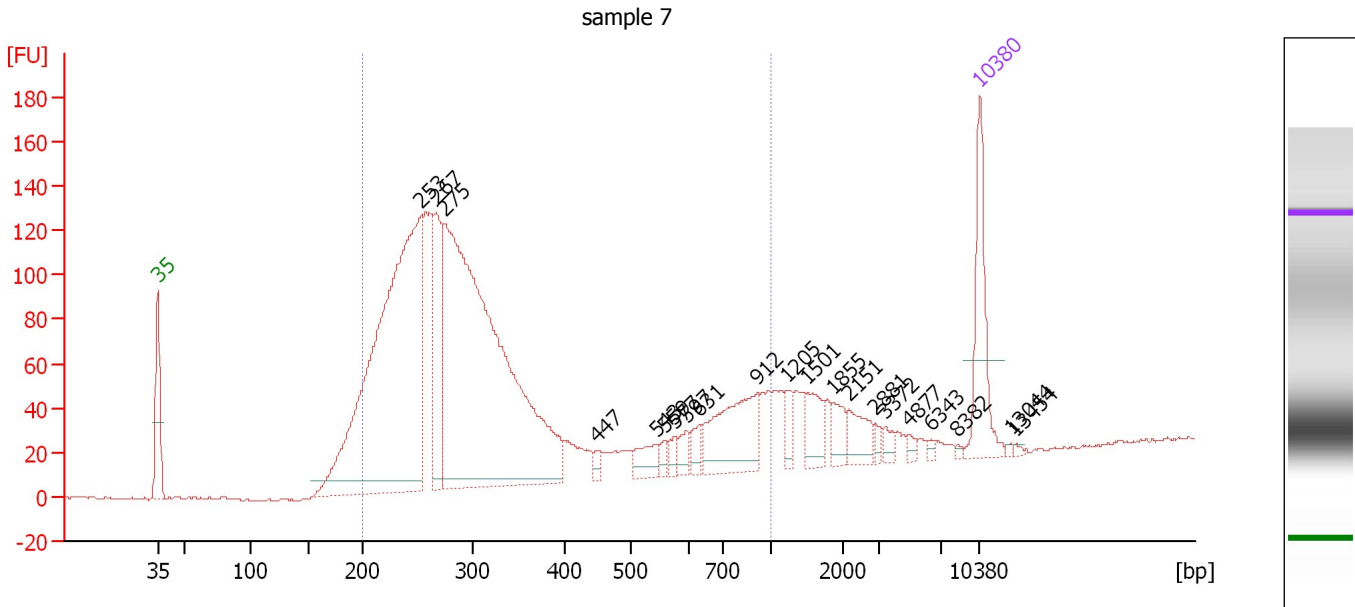
Region table for sample 6 : sample 6

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	313	5,826.61	5,926.2	30,098.6	97	28.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
 Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : sample 7

Number of peaks found: 21 Corr. Area 1: 2,224.8
 Noise: 0.5

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	253	768.82	4,597.8		65.45
3	267	133.18	754.4		66.75
4	275	875.14	4,820.5		67.45
5	447	8.92	30.2		80.36
6	543	25.67	71.6		85.43
7	559	8.57	23.2		86.19
8	577	9.69	25.4		87.09
9	597	14.99	38.0		88.04
10	631	16.07	38.6		89.10
11	912	100.42	166.9		94.07
12	1,205	17.16	21.6		96.53
13	1,501	33.06	33.4		98.34
14	1,855	19.56	16.0		100.50
15	2,151	27.33	19.2		101.85
16	2,881	6.14	3.2		104.11
17	3,372	8.93	4.0		104.97
18	4,877	4.85	1.5		106.92
19	6,343	3.66	0.9		108.83
20	8,382	2.06	0.4		111.04
21	10,380	75.00	10.9	Upper Marker	113.00
22	13,044	0.00	0.0		115.61
23	13,454	0.00	0.0		116.01


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...

... Region table for sample 7 :

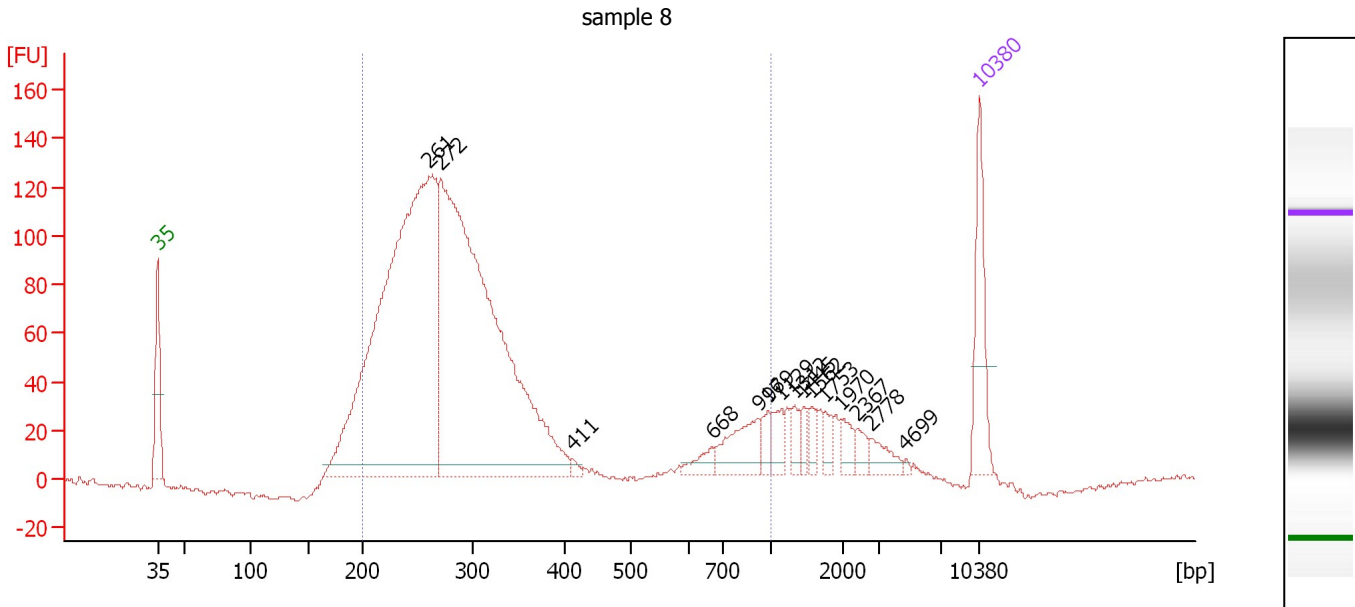
sample 7

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	352	1,915.24	2,224.8	9,997.9	 84	50.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
 Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : sample 8

Number of peaks found: 15 Corr. Area 1: 2,036.8
 Noise: 1.0

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	261	1,108.40	6,436.2		66.15
3	272	1,073.49	5,981.4		67.16
4	411	5.33	19.7		78.36
5	668	19.66	44.6		90.21
6	916	57.82	95.7		94.13
7	979	19.54	30.2		94.99
8	1,129	21.81	29.3		96.06
9	1,312	17.06	19.7		97.18
10	1,445	13.29	13.9		97.99
11	1,562	15.15	14.7		98.70
12	1,753	14.24	12.3		99.88
13	1,970	17.61	13.6		101.20
14	2,367	13.34	8.5		102.52
15	2,778	19.80	10.8		103.79
16	4,699	2.12	0.7		106.69
17	10,380	75.00	10.9	Upper Marker	113.00

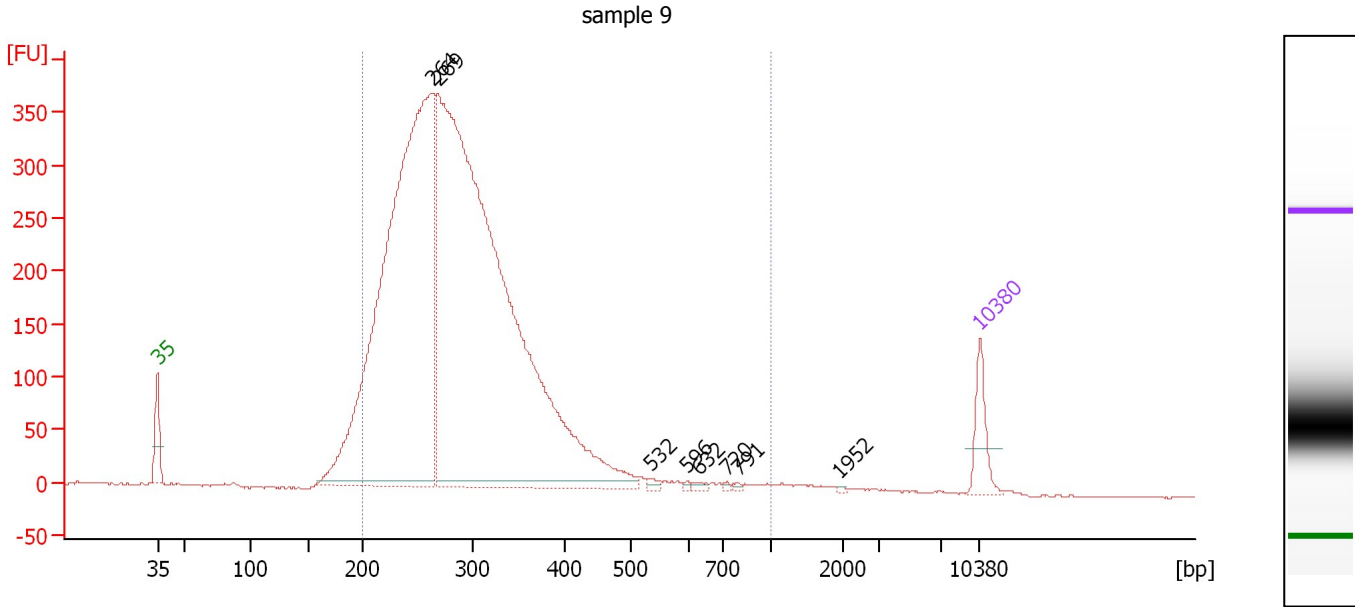
Region table for sample 8 : sample 8

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Total	Size distribution in CV [%]
200	1,000	322	2,145.11	2,036.8	11,623.1	85	48.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
 Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : sample 9

Number of peaks found: 8 Corr. Area 1: 6,231.5
 Noise: 0.9

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	264	2,645.07	15,198.5		66.40
3	269	3,613.90	20,377.8		66.87
4	532	10.92	31.1		84.91
5	596	4.68	11.9		88.00
6	632	8.74	21.0		89.13
7	720	5.01	10.5		91.45
8	791	5.08	9.7		92.43
9	1,952	2.60	2.0		101.09
10	10,380	75.00	10.9	Upper Marker	113.00

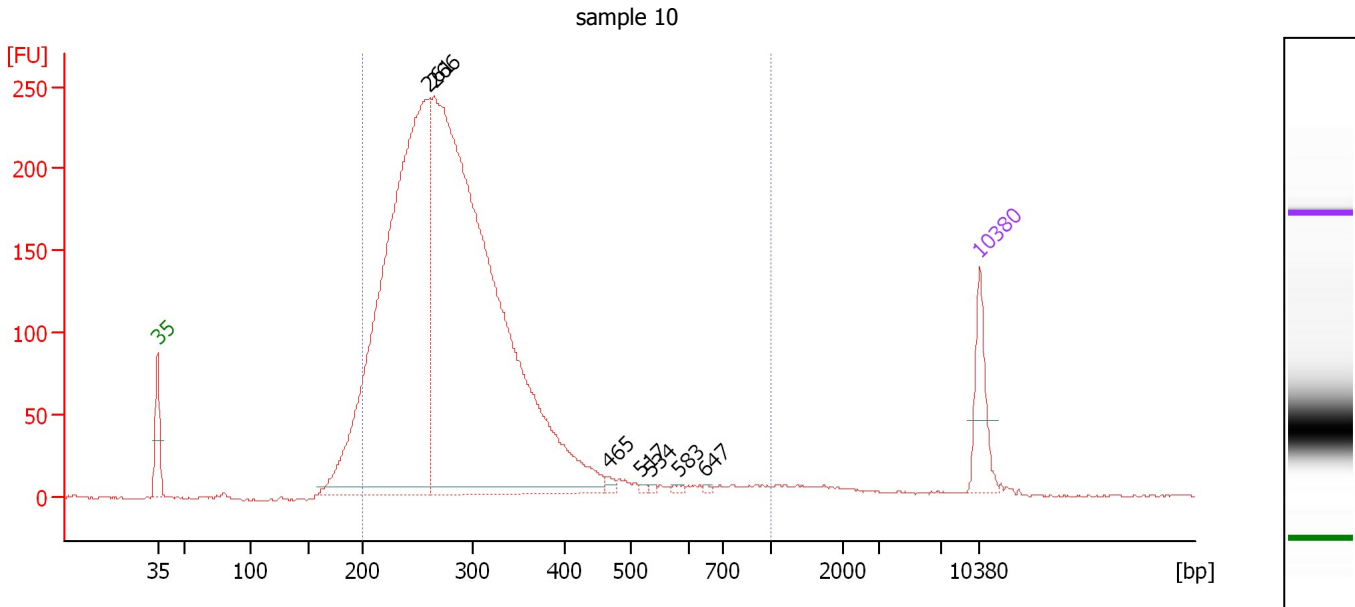
Region table for sample 9 : sample 9

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	296	5,870.17	6,231.5	31,789.9	95	26.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
 Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : sample 10

Number of peaks found: 7 Corr. Area 1: 3,883.1
 Noise: 0.7

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	261	1,683.94	9,793.4		66.11
3	266	2,327.21	13,278.2		66.57
4	465	8.92	29.1		81.36
5	517	4.33	12.7		84.15
6	534	3.76	10.7		84.98
7	583	5.50	14.3		87.36
8	647	3.21	7.5		89.58
9	10,380	75.00	10.9	Upper Marker	113.00

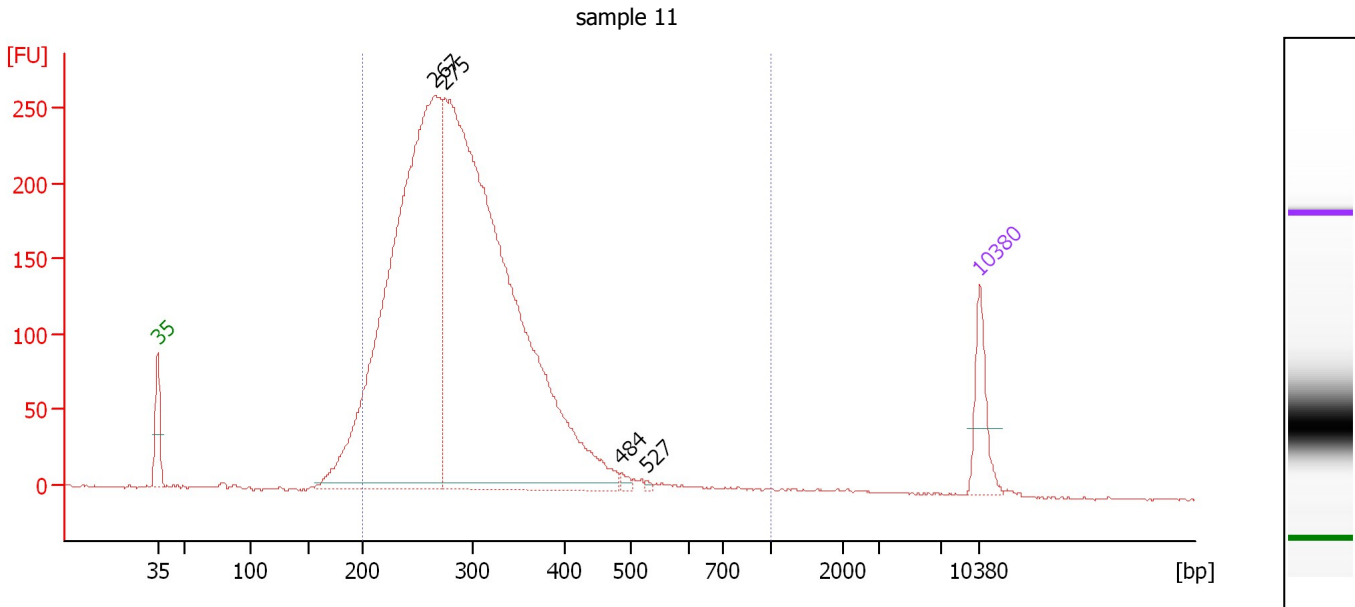
Region table for sample 10 : sample 10

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	295	3,861.85	3,883.1	21,120.0	94	29.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
 Modified: 8/2/2019 3:44:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : sample 11

Number of peaks found: 4 Corr. Area 1: 4,370.4
 Noise: 0.7

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	267	1,860.80	10,578.8		66.66
3	275	2,402.52	13,238.6		67.45
4	484	10.00	31.3		82.46
5	527	3.71	10.7		84.65
6	10,380	75.00	10.9	Upper Marker	113.00

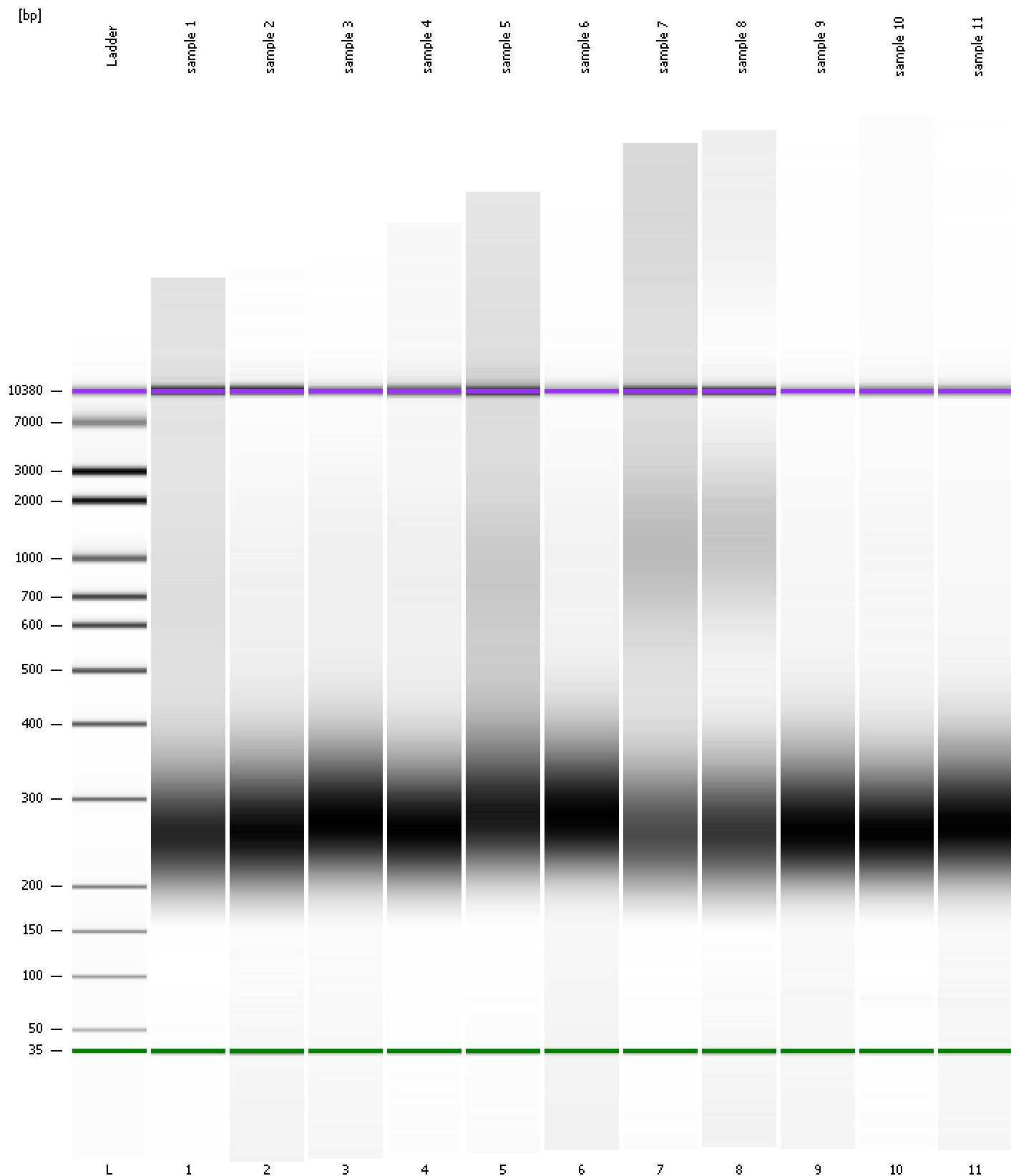
Region table for sample 11 : sample 11

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	299	4,038.88	4,370.4	21,594.7	95	24.4

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad

Created: 8/2/2019 3:02:47 PM
Modified: 8/2/2019 3:44:06 PM

Gel Image



Assay Class: High Sensitivity DNA Assay Created: 8/2/2019 3:02:47 PM
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-08-02\2019-08-02_004.xad Modified: 8/2/2019 3:44:06 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		8/2/2019 3:44:05 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\2019-08-02\2019-08-02_004.xad)		Instrument	Run		8/2/2019 3:02:53 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		8/2/2019 3:02:53 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		8/2/2019 3:02:53 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		8/2/2019 3:02:53 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		8/2/2019 3:02:53 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		8/2/2019 3:02:53 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		8/2/2019 3:02:53 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1