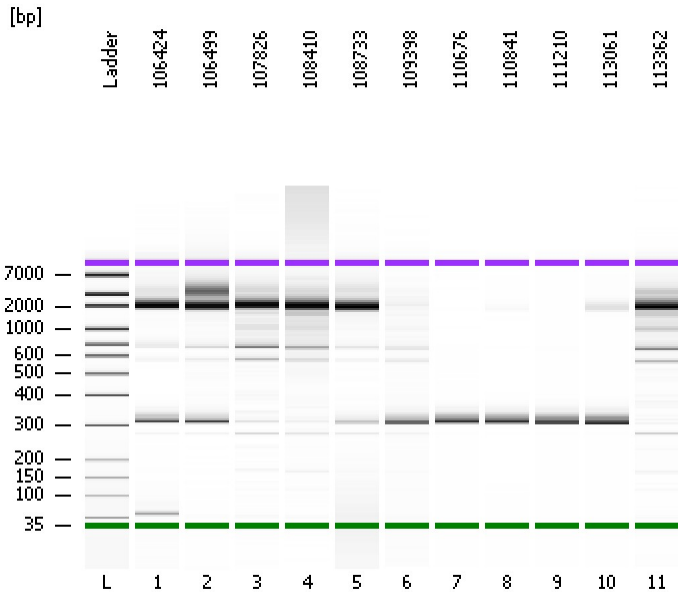


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
Modified: 11/4/2019 1:17:08 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

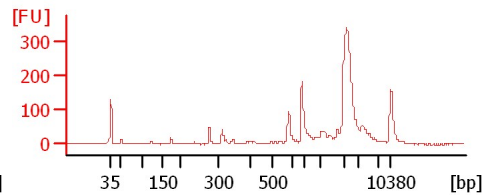
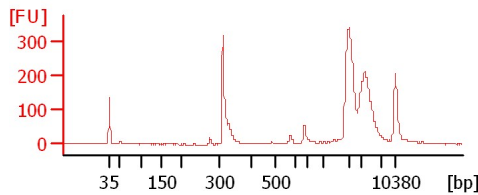
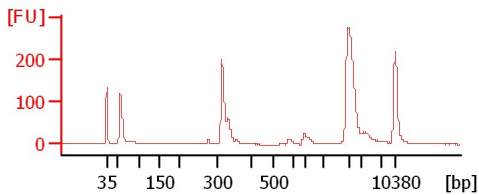
Chip Information:

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

106424

106499

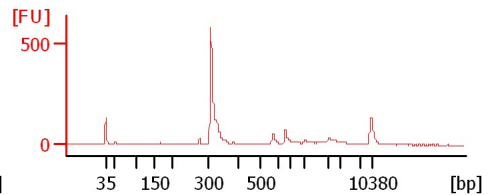
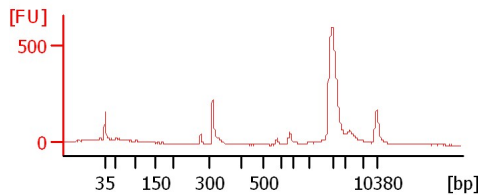
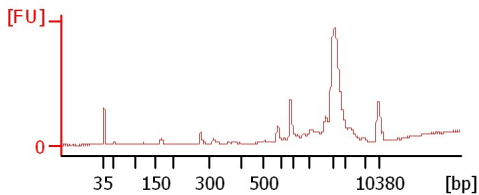
107826



108410

108733

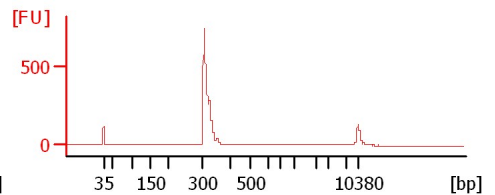
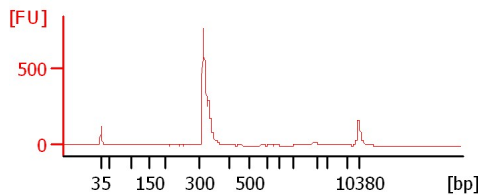
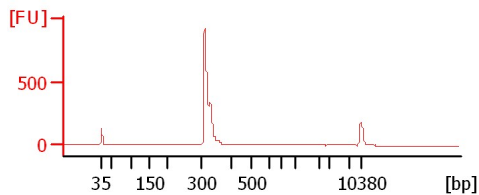
109398



110676

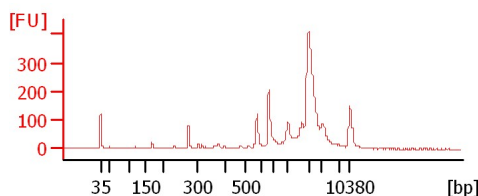
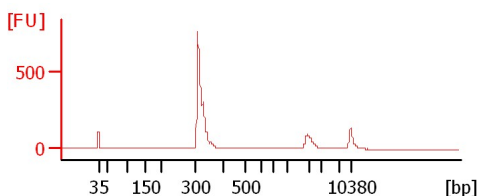
110841

111210



113061

113362



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
 Modified: 11/4/2019 1:17:08 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
106424		<input type="checkbox"/>	✓			
106499		<input type="checkbox"/>	✓			
107826		<input type="checkbox"/>	✓			
108410		<input type="checkbox"/>	✓			
108733		<input type="checkbox"/>	✓			
109398		<input type="checkbox"/>	✓			
110676		<input type="checkbox"/>	✓			
110841		<input type="checkbox"/>	✓			
111210		<input type="checkbox"/>	✓			
113061		<input type="checkbox"/>	✓			
113362		<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\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
Modified: 11/4/2019 1:17:08 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

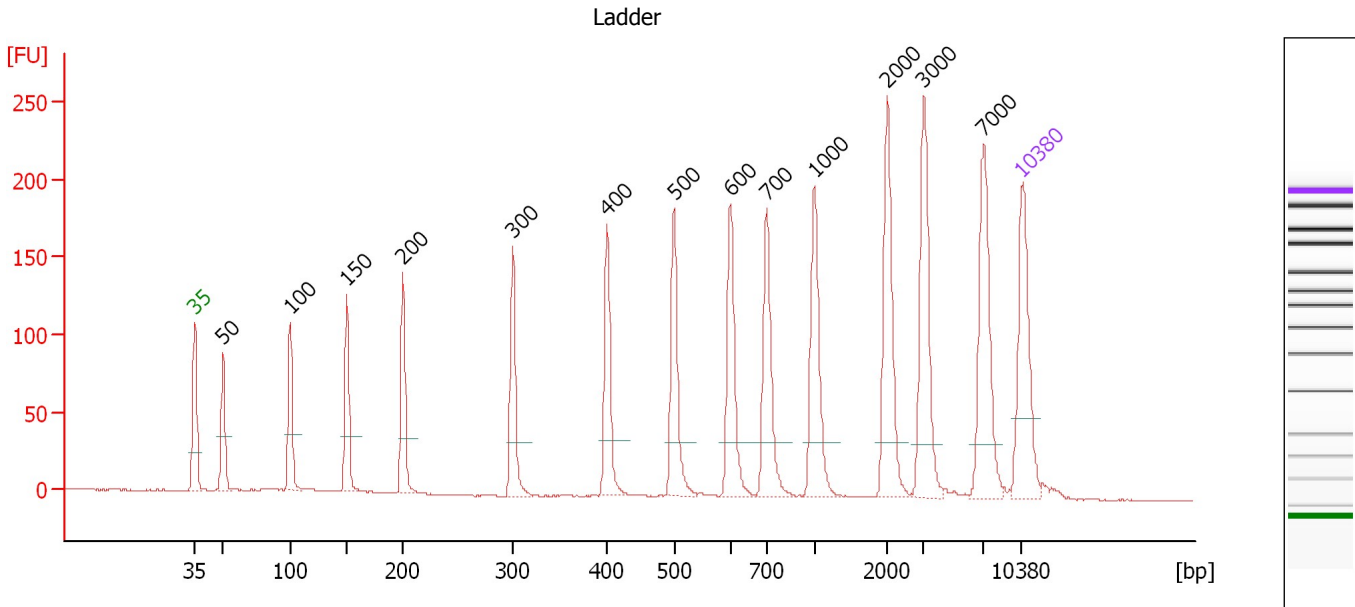
Ladder

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\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

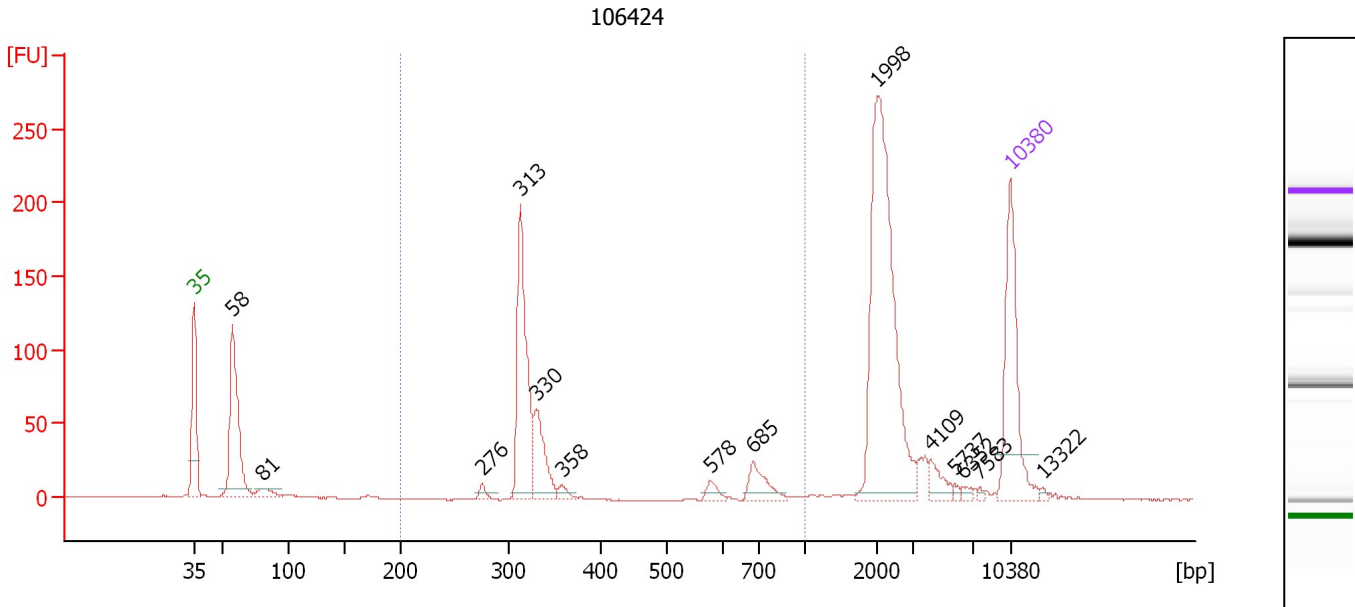
Peak table for Ladder

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	45.39
3	100	150.00	2,272.7	Ladder Peak	51.04
4	150	150.00	1,515.2	Ladder Peak	55.87
5	200	150.00	1,136.4	Ladder Peak	60.61
6	300	150.00	757.6	Ladder Peak	69.92
7	400	150.00	568.2	Ladder Peak	77.91
8	500	150.00	454.5	Ladder Peak	83.55
9	600	150.00	378.8	Ladder Peak	88.34
10	700	150.00	324.7	Ladder Peak	91.41
11	1,000	150.00	227.3	Ladder Peak	95.43
12	2,000	150.00	113.6	Ladder Peak	101.62
13	3,000	150.00	75.8	Ladder Peak	104.74
14	7,000	150.00	32.5	Ladder Peak	109.79
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
 Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : 106424

Number of peaks found: 14 Corr. Area 1: 440.0
 Noise: 0.4

Peak table for sample 1 : 106424

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	58	146.76	3,862.3		46.25
3	81	13.02	243.3		48.90
4	276	5.83	32.1		67.66
5	313	137.12	663.8		70.95
6	330	50.46	231.6		72.33
7	358	6.43	27.3		74.52
8	578	8.14	21.3		87.29
9	685	22.77	50.4		90.95
10	1,998	213.69	162.0		101.61
11	4,109	13.01	4.8		106.14
12	5,737	2.25	0.6		108.20
13	6,352	2.89	0.7		108.97
14	7,583	1.51	0.3		110.35
15	10,380	75.00	10.9	Upper Marker	113.00
16	13,322	0.00	0.0		115.79

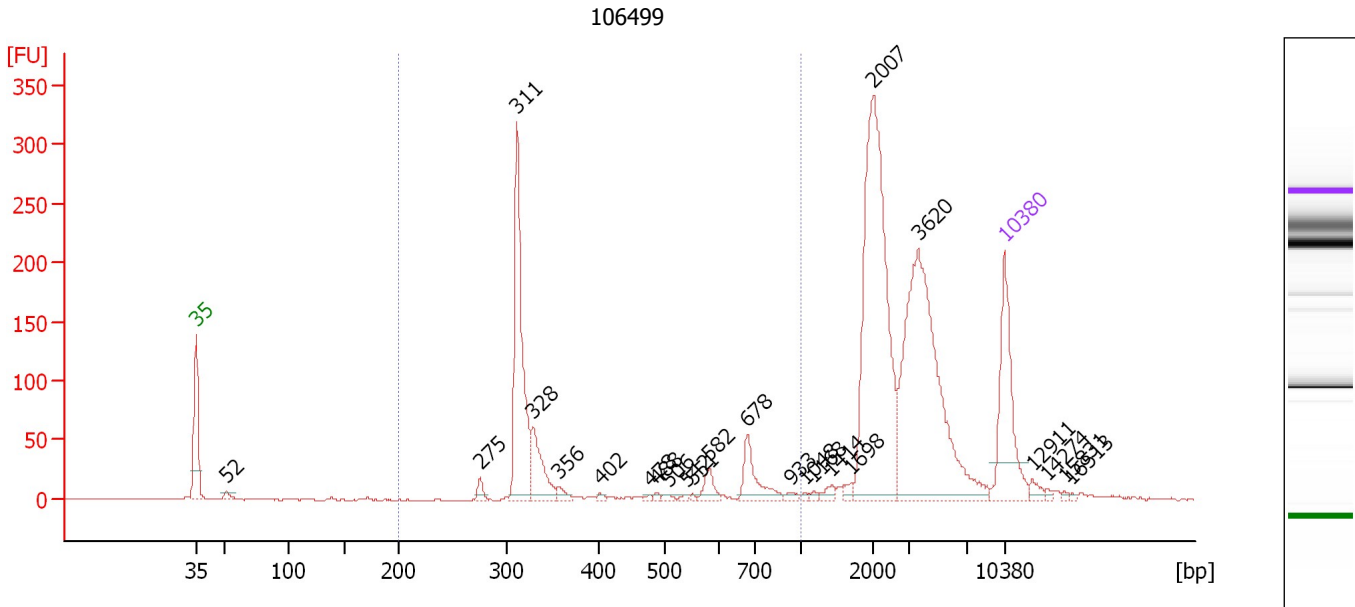
Region table for sample 1 : 106424

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	387	220.49	440.0	972.2	31	38.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

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Electropherogram Summary Continued ...



Overall Results for sample 2 : 106499

Number of peaks found: 24 Corr. Area 1: 625.9
 Noise: 0.5

Peak table for sample 2 : 106499

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	52	6.59	191.5		45.63
3	275	9.59	52.8		67.61
4	311	191.81	934.4		70.80
5	328	53.05	244.8		72.18
6	356	7.79	33.2		74.40
7	402	2.76	10.4		78.00
8	478	1.91	6.1		82.29
9	488	2.34	7.3		82.89
10	506	3.20	9.6		83.86
11	542	2.22	6.2		85.57
12	551	1.57	4.3		85.99
13	582	14.56	37.9		87.47
14	678	35.47	79.2		90.74
15	933	3.84	6.2		94.53
16	1,048	2.13	3.1		95.73
17	1,168	2.31	3.0		96.47
18	1,414	5.59	6.0		97.99
19	1,698	4.51	4.0		99.75
20	2,007	272.58	205.8		101.64
21	3,620	254.20	106.4		105.52
22	10,380	75.00	10.9	Upper Marker	113.00
23	12,911	0.00	0.0		115.40
24	14,274	0.00	0.0		116.69
25	15,831	0.00	0.0		118.17


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...**... Peak table for sample 2 : 106499**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	16,513	0.00	0.0		118.82

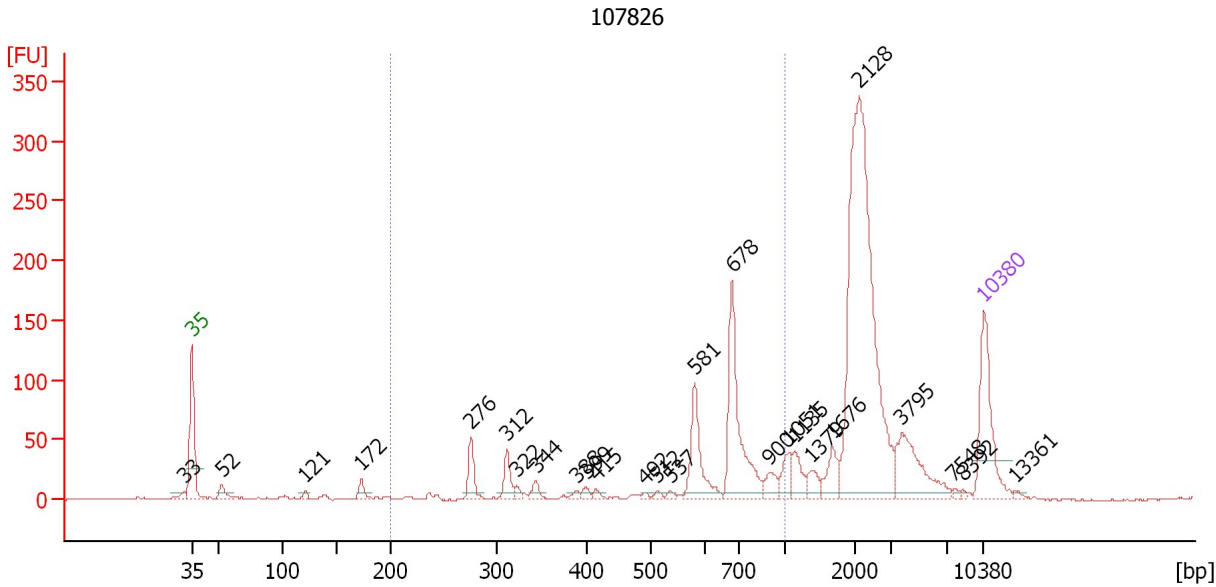
Region table for sample 2 : 106499

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	410	320.81	625.9	1,373.8	 29	39.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
 Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : 107826

Number of peaks found: 26 Corr. Area 1: 550.4
 Noise: 1.0

Peak table for sample 3 : 107826

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	33	0.00	0.0		42.34
2	35	125.00	5,411.3	Lower Marker	43.00
3	52	15.00	435.5		45.64
4	121	4.92	61.8		53.04
5	172	12.88	113.3		57.99
6	276	32.45	178.0		67.70
7	312	25.76	125.2		70.86
8	322	6.14	28.9		71.71
9	344	9.97	44.0		73.40
10	388	3.90	15.2		76.99
11	399	6.04	22.9		77.84
12	415	4.30	15.7		78.78
13	492	2.60	8.0		83.11
14	512	3.34	9.9		84.15
15	537	3.40	9.6		85.33
16	581	66.75	174.0		87.45
17	678	131.61	293.9		90.75
18	900	15.37	25.9		94.10
19	1,051	19.68	28.4		95.75
20	1,135	20.95	28.0		96.27
21	1,379	13.85	15.2		97.77
22	1,676	22.99	20.8		99.61
23	2,128	388.73	276.8		102.02
24	3,795	65.06	26.0		105.74
25	7,548	2.09	0.4		110.31

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
 Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...

... Peak table for sample 3 : 107826

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	8,392	1.57	0.3		111.11
27	10,380	75.00	10.9	Upper Marker	113.00
28	13,361	0.00	0.0		115.83

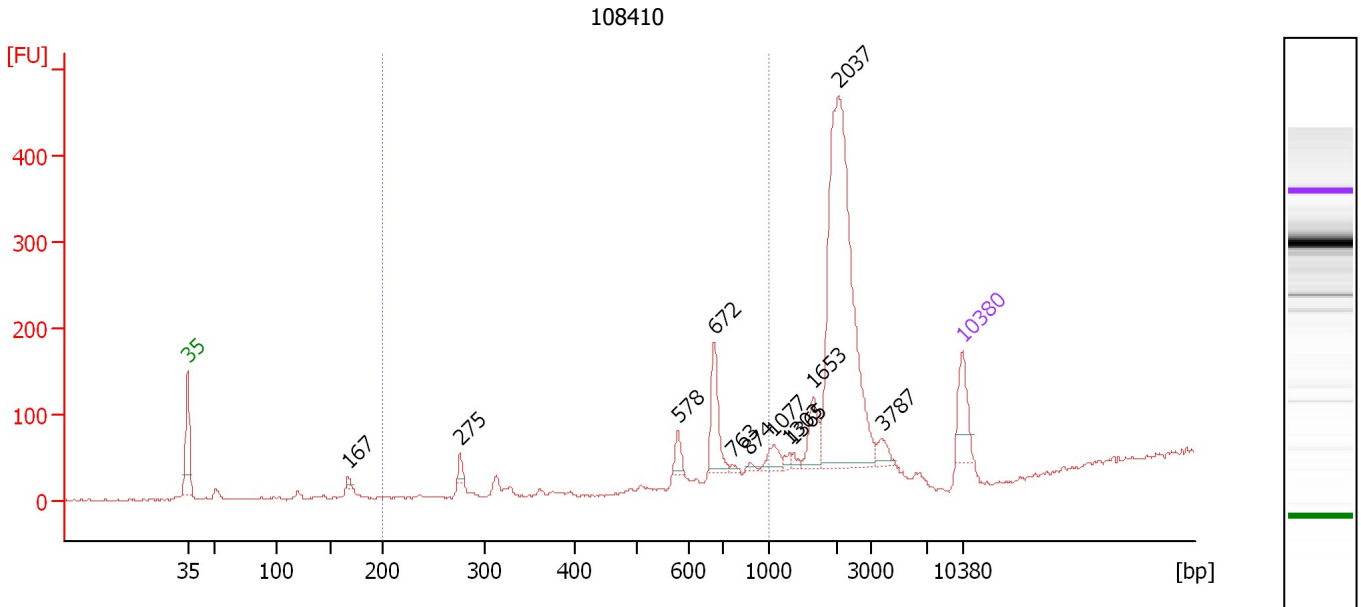
Region table for sample 3 : 107826

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	603	345.62	550.4	1,063.0	31	30.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
 Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : 108410

Number of peaks found: 12 Corr. Area 1: 199.0
 Noise: 1.2

Peak table for sample 4 : 108410

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	167	11.49	104.2		57.49
3	275	24.43	134.5		67.60
4	578	31.36	82.2		87.29
5	672	118.11	266.2		90.57
6	763	5.56	11.0		92.25
7	874	6.93	12.0		93.74
8	1,077	31.98	45.0		95.91
9	1,303	10.53	12.2		97.31
10	1,365	9.21	10.2		97.69
11	1,653	64.99	59.6		99.47
12	2,037	758.84	564.4		101.73
13	3,787	25.62	10.2		105.73
14	10,380	75.00	10.9	Upper Marker	113.00

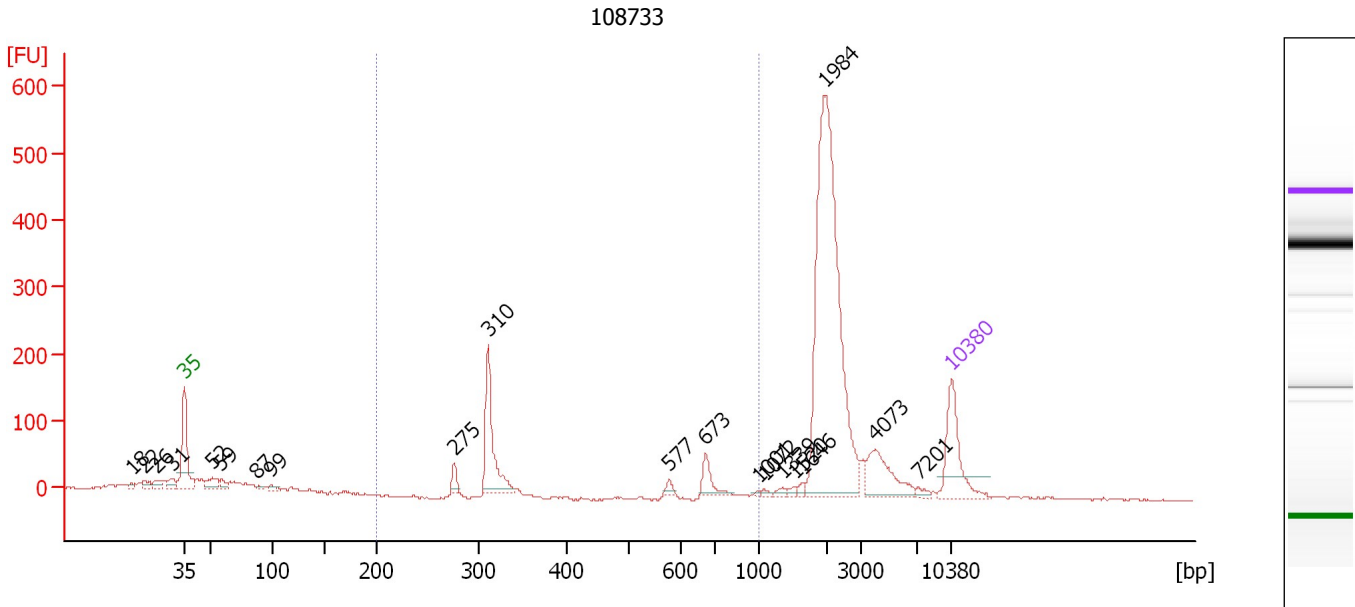
Region table for sample 4 : 108410

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	642	183.94	199.0	498.7	14	21.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

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 Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : 108733

Number of peaks found: 20 Corr. Area 1: 362.4
 Noise: 1.3

Peak table for sample 5 : 108733

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	18	0.00	0.0		38.18
2	22	0.00	0.0		39.40
3	26	0.00	0.0		40.37
4	31	0.00	0.0		41.88
5	35	125.00	5,411.3	Lower Marker	43.00
6	52	29.19	856.4		45.58
7	59	14.24	368.7		46.36
8	87	5.56	96.4		49.62
9	99	8.15	125.3		50.88
10	275	16.75	92.2		67.61
11	310	132.07	645.2		70.73
12	577	7.70	20.2		87.22
13	673	29.36	66.1		90.57
14	1,001	2.01	3.0		95.44
15	1,072	3.41	4.8		95.88
16	1,339	4.60	5.2		97.53
17	1,520	4.33	4.3		98.65
18	1,646	5.19	4.8		99.43
19	1,984	511.10	390.3		101.52
20	4,073	65.32	24.3		106.09
21	7,201	5.91	1.2		109.98
22	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...

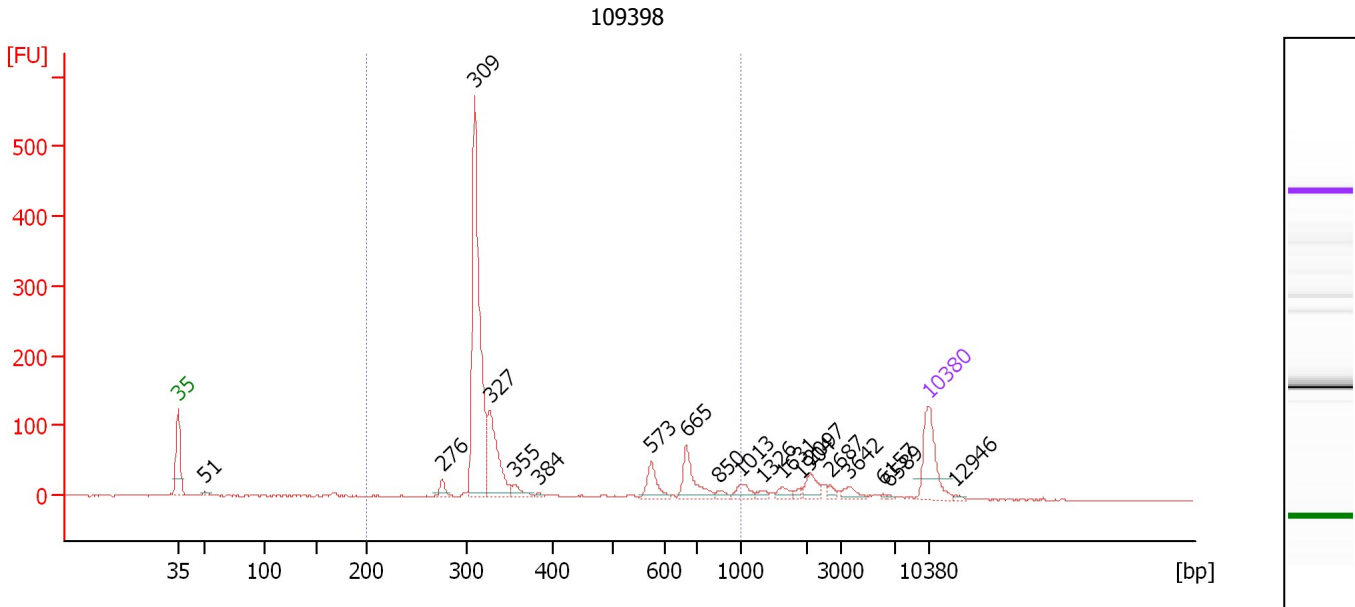
... Region table for sample 5 : 108733

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	418	190.07	362.4	825.6	 15	42.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

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 Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : 109398

Number of peaks found: 19 Corr. Area 1: 1,186.0
 Noise: 1.0

Peak table for sample 6 : 109398

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	51	6.87	204.9		45.48
3	276	18.80	103.4		67.64
4	309	499.50	2,445.6		70.67
5	327	145.50	674.5		72.06
6	355	21.24	90.7		74.30
7	384	4.24	16.7		76.63
8	573	40.43	106.8		87.07
9	665	62.31	141.9		90.35
10	850	8.33	14.9		93.43
11	1,013	19.36	29.0		95.51
12	1,326	7.75	8.9		97.45
13	1,631	11.59	10.8		99.34
14	1,904	5.32	4.2		101.03
15	2,097	23.25	16.8		101.92
16	2,687	7.23	4.1		103.76
17	3,642	13.92	5.8		105.55
18	6,157	2.09	0.5		108.73
19	6,589	1.52	0.3		109.27
20	10,380	75.00	10.9	Upper Marker	113.00
21	12,946	0.00	0.0		115.43

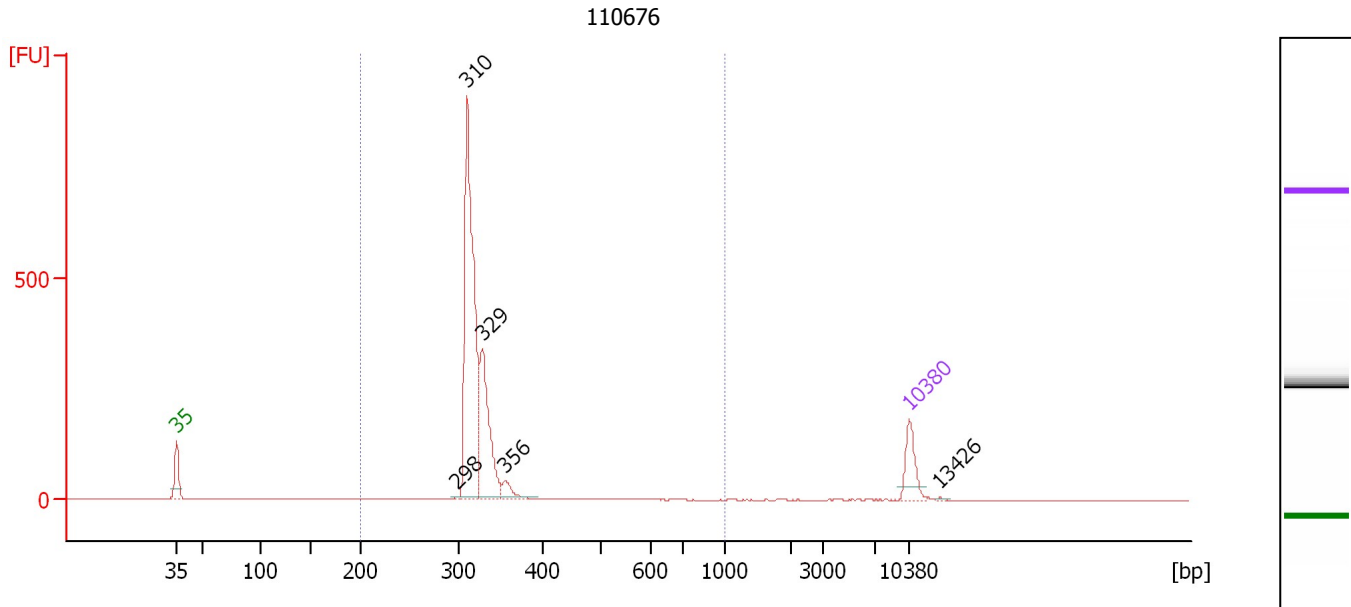
Region table for sample 6 : 109398

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	395	855.41	1,186.0	3,764.6	79	39.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

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 Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : 110676

Number of peaks found: 5 Corr. Area 1: 1,846.4
 Noise: 0.3

Peak table for sample 7 : 110676

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	298	4.34	22.1		69.71
3	310	881.12	4,304.7		70.73
4	329	336.91	1,550.8		72.25
5	356	47.98	204.0		74.43
6	10,380	75.00	10.9	Upper Marker	113.00
7	13,426	0.00	0.0		115.89

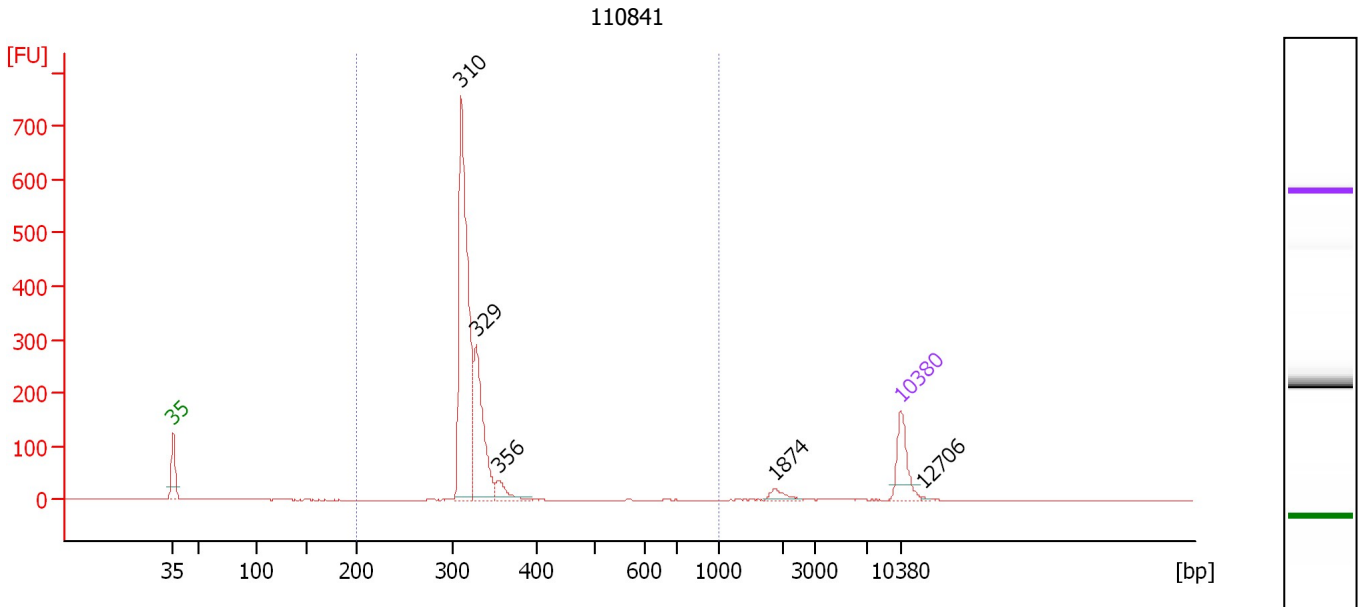
Region table for sample 7 : 110676

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	324	1,271.09	1,846.4	5,970.2	98	10.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
 Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : 110841

Number of peaks found: 5 Corr. Area 1: 1,541.5
 Noise: 0.3

Peak table for sample 8 : 110841

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	310	736.07	3,598.0		70.71
3	329	291.81	1,345.6		72.20
4	356	41.95	178.8		74.36
5	1,874	17.66	14.3		100.84
6	10,380	75.00	10.9	Upper Marker	113.00
7	12,706	0.00	0.0		115.21

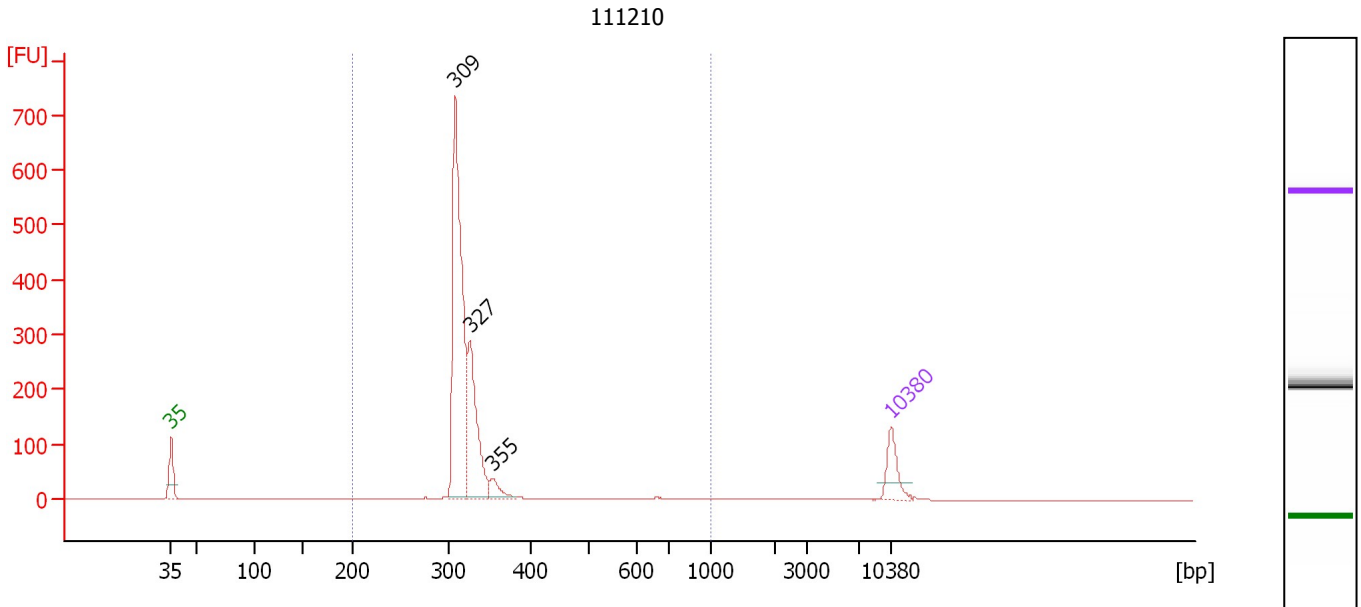
Region table for sample 8 : 110841

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	322	1,062.60	1,541.5	5,004.5	96	6.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
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Electropherogram Summary Continued ...



Overall Results for sample 9 : 111210

Number of peaks found: 3 Corr. Area 1: 1,570.4
 Noise: 0.3

Peak table for sample 9 : 111210

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	309	921.23	4,520.6		70.62
3	327	357.18	1,655.4		72.07
4	355	50.00	213.5		74.30
5	10,380	75.00	10.9	Upper Marker	113.00

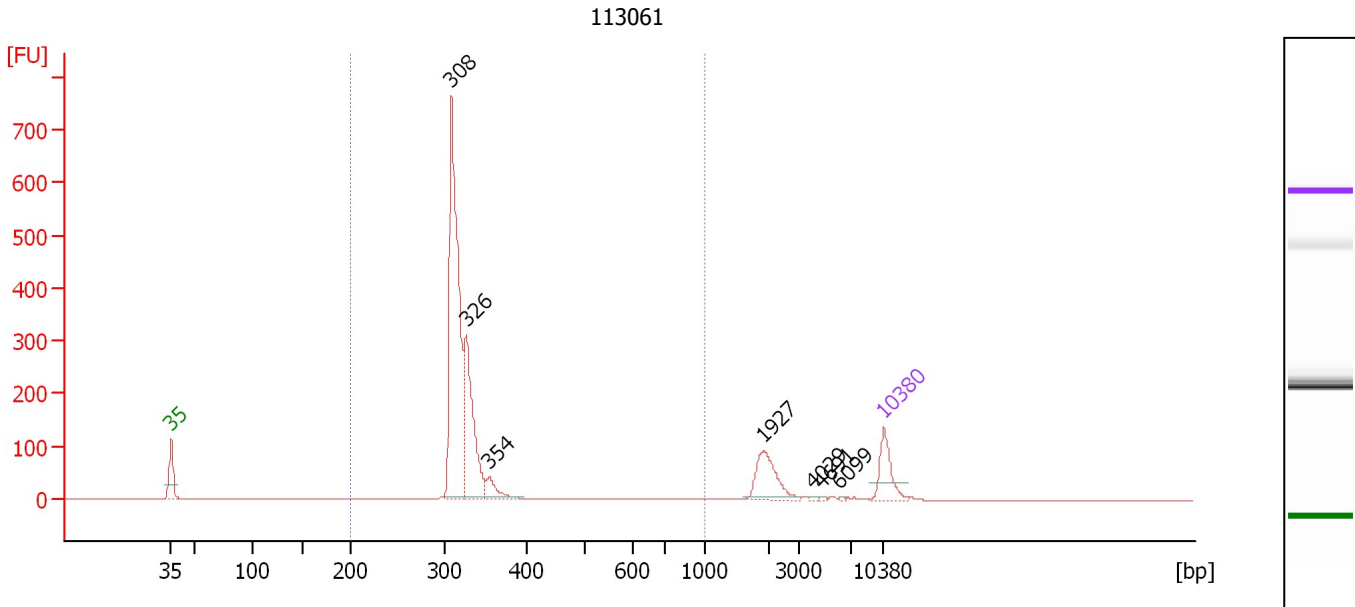
Region table for sample 9 : 111210

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	326	1,353.52	1,570.4	6,366.6	97	15.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
 Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : 113061

Number of peaks found: 7 Corr. Area 1: 1,639.1
 Noise: 0.3

Peak table for sample 10 : 113061

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	308	858.37	4,225.4		70.54
3	326	353.41	1,641.9		72.01
4	354	54.52	233.2		74.26
5	1,927	111.83	87.9		101.17
6	4,029	2.93	1.1		106.04
7	4,691	2.04	0.7		106.87
8	6,099	1.89	0.5		108.65
9	10,380	75.00	10.9	Upper Marker	113.00

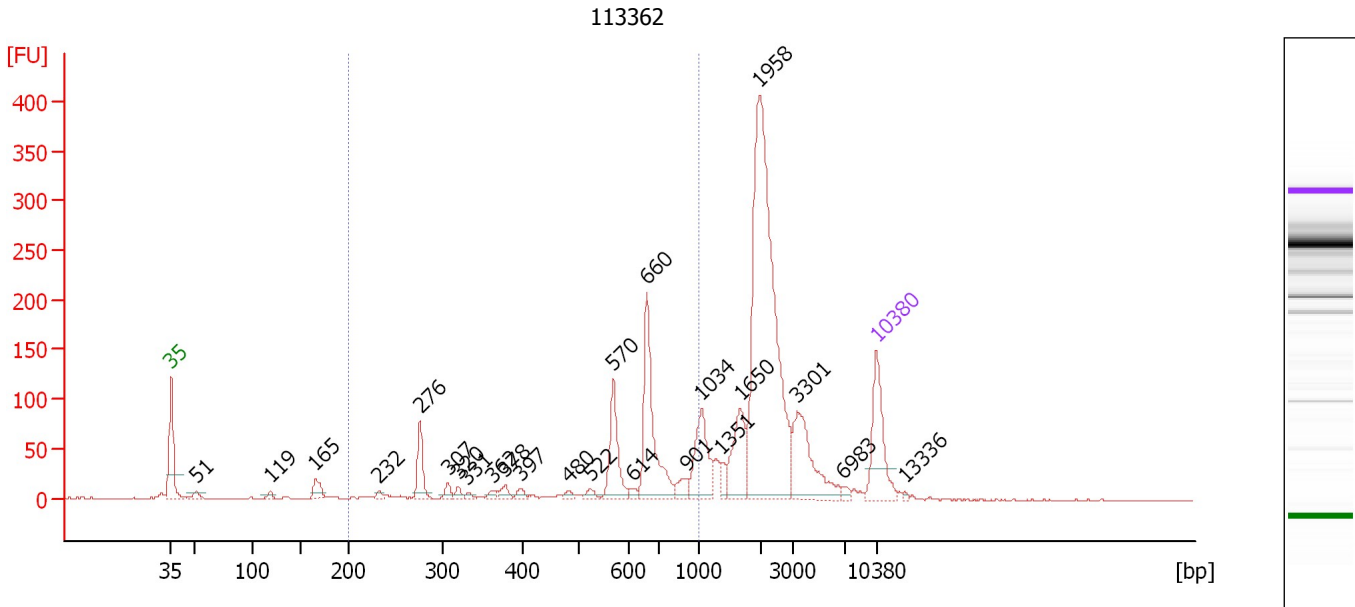
Region table for sample 10 : 113061

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	325	1,283.53	1,639.1	6,043.1	86	14.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
 Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : 113362

Number of peaks found: 24 Corr. Area 1: 661.9
 Noise: 0.4

Peak table for sample 11 : 113362

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	51	11.73	346.5		45.54
3	119	8.26	105.4		52.84
4	165	23.67	216.8		57.34
5	232	6.23	40.6		63.63
6	276	50.83	278.8		67.71
7	307	10.39	51.3		70.46
8	320	7.98	37.8		71.52
9	331	4.90	22.4		72.42
10	362	7.25	30.3		74.90
11	378	11.62	46.5		76.17
12	397	7.82	29.8		77.71
13	480	6.28	19.8		82.42
14	522	8.04	23.3		84.59
15	570	76.50	203.3		86.92
16	614	6.18	15.2		88.77
17	660	149.67	343.4		90.20
18	901	16.76	28.2		94.11
19	1,034	78.06	114.3		95.65
20	1,351	13.49	15.1		97.60
21	1,650	64.82	59.5		99.46
22	1,958	466.73	361.2		101.36
23	3,301	95.28	43.7		105.12
24	6,983	5.53	1.2		109.77
25	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
Modified: 11/4/2019 1:17:08 PM

Electropherogram Summary Continued ...**... Peak table for sample 11 : 113362**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	13,336	0.00	0.0		115.80

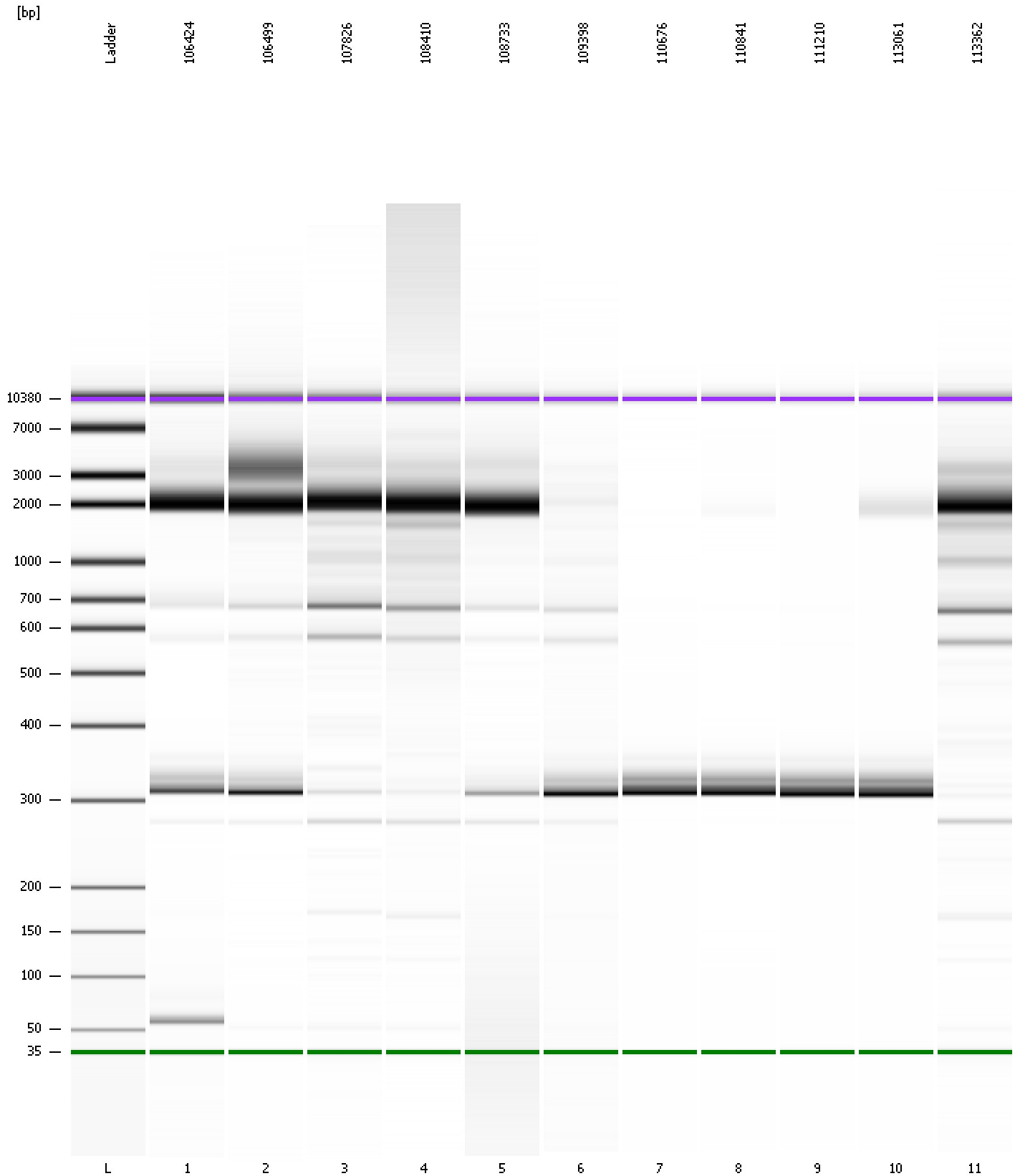
Region table for sample 11 : 113362

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	603	424.44	661.9	1,326.0	 30	32.1

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
Modified: 11/4/2019 1:17:08 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-04\2019-11-04_002.xad

Created: 11/4/2019 12:35:47 PM
 Modified: 11/4/2019 1:17:08 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		11/4/2019 1:17:04 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\2019-11-04\2019-11-04_002.xad)		Instrument	Run		11/4/2019 12:35:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		11/4/2019 12:35:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		11/4/2019 12:35:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		11/4/2019 12:35:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		11/4/2019 12:35:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		11/4/2019 12:35:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		11/4/2019 12:35:52 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1