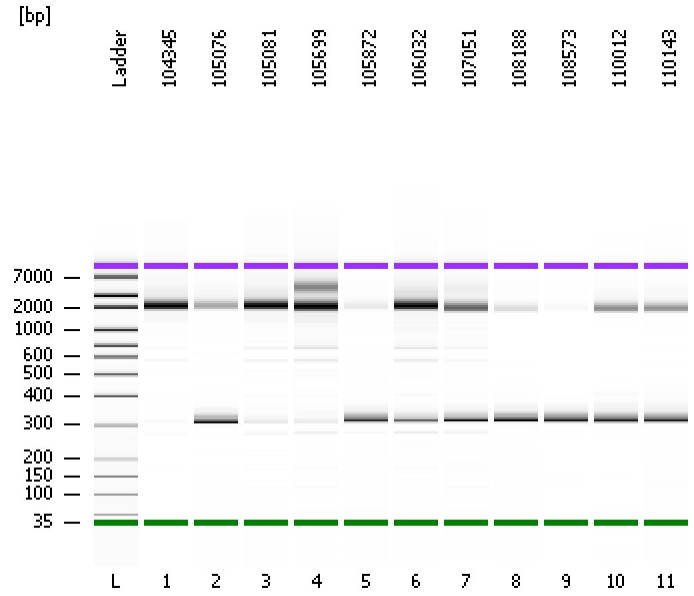
Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electrophoresis File Run Summary



Instrument Information:

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

Assay Information:

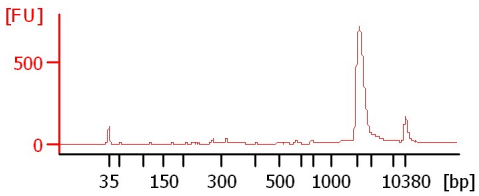
Assay Origin Path: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\assays\dsDNA\High Sensitivity DNA.xsy

Assay Class: High Sensitivity DNA Assay
Version: 1.03
Assay Comments: Copyright © 2003-2010 Agilent Technologies

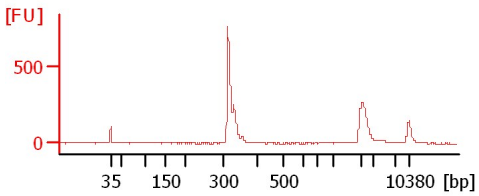
Chip Information:

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

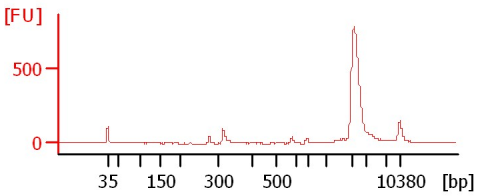
104345



105076



105081



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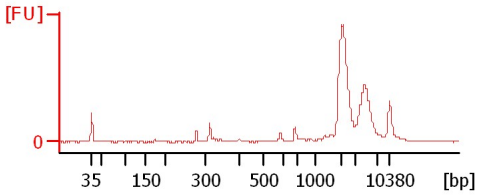

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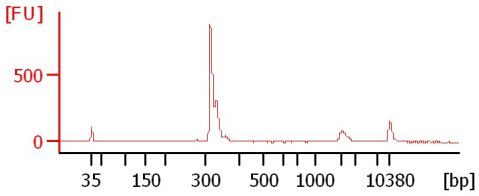

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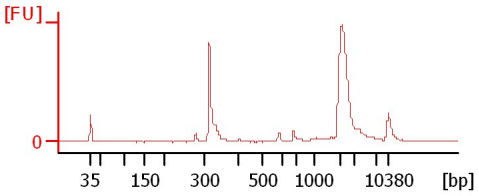
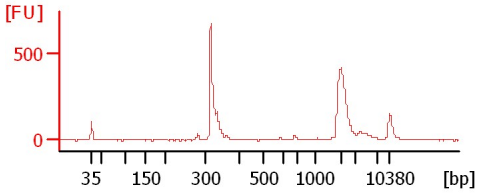
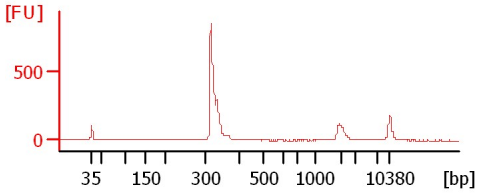


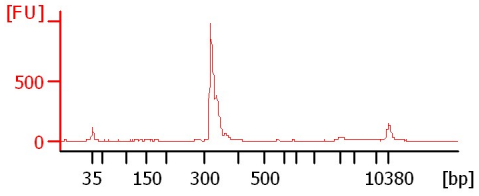
107051



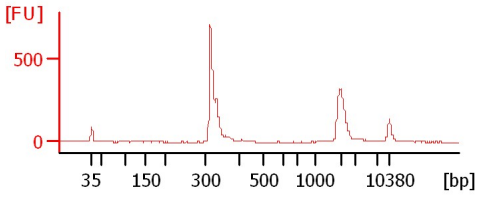
108188



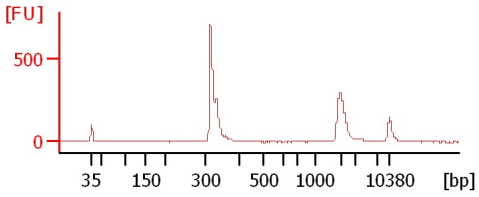
108573



110012



110143



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
104345		☐	✔			
105076		☐	✔			
105081		☐	✔			
105699		☐	✔			
105872		☐	✔			
106032		☐	✔			
107051		☐	✔			
108188		☐	✔			
108573		☐	✔			
110012		☐	✔			
110143		☐	✔			
Ladder		☐	✔			

Chip Lot # **Reagent Kit Lot #**

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 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

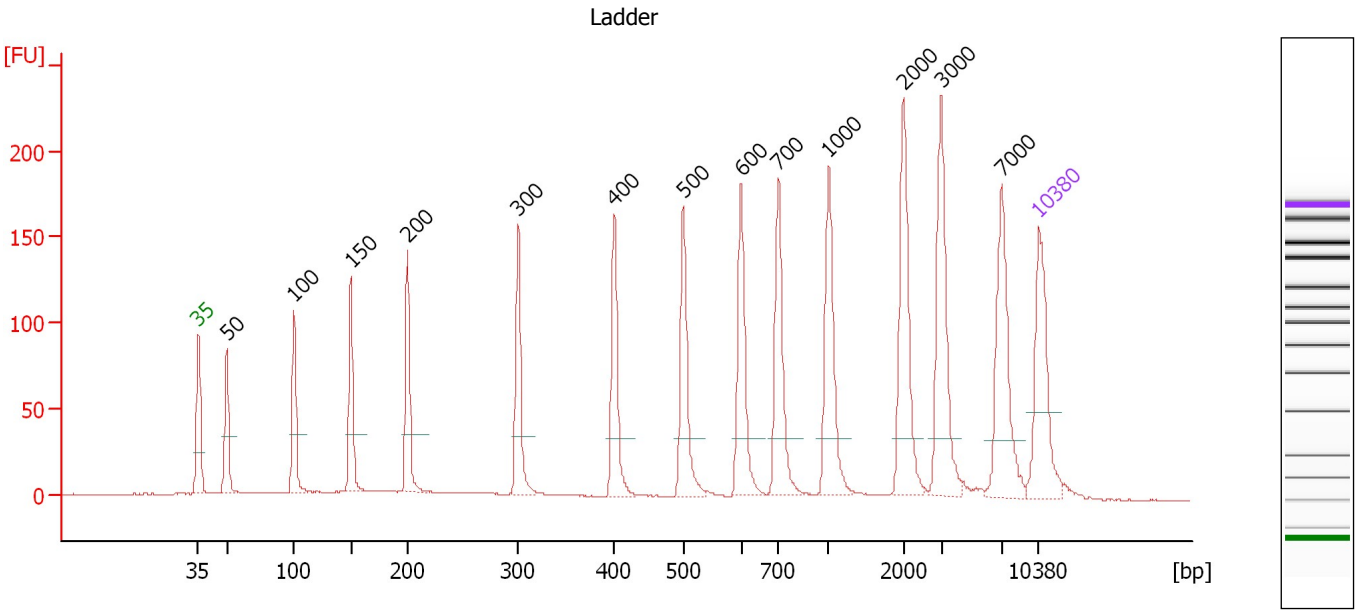
Ladder

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\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

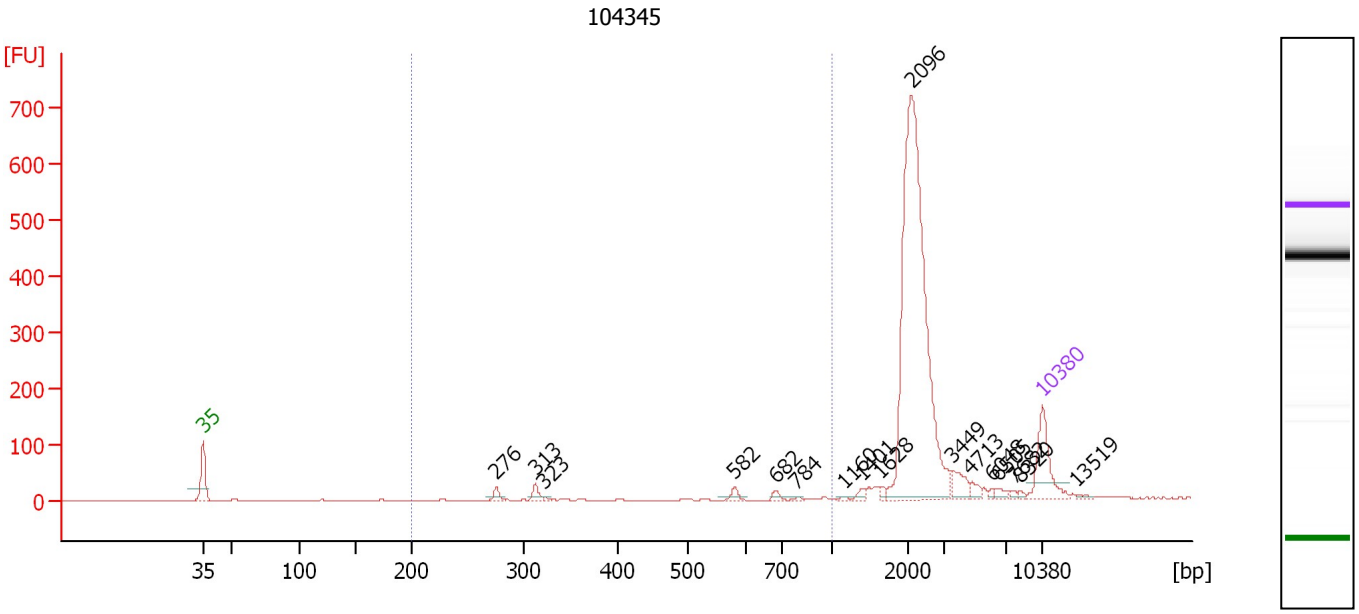
Peak table for Ladder

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	45.35
3		100	150.00	2,272.7	Ladder Peak	50.95
4		150	150.00	1,515.2	Ladder Peak	55.70
5		200	150.00	1,136.4	Ladder Peak	60.41
6		300	150.00	757.6	Ladder Peak	69.65
7		400	150.00	568.2	Ladder Peak	77.64
8		500	150.00	454.5	Ladder Peak	83.37
9		600	150.00	378.8	Ladder Peak	88.22
10		700	150.00	324.7	Ladder Peak	91.32
11		1,000	150.00	227.3	Ladder Peak	95.50
12		2,000	150.00	113.6	Ladder Peak	101.76
13		3,000	150.00	75.8	Ladder Peak	104.87
14		7,000	150.00	32.5	Ladder Peak	109.94
15		10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : 104345

Number of peaks found:	17	Corr. Area 1:	84.9
Noise:	0.2		

Peak table for sample 1 : 104345

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	276	13.87	76.1		67.46
3	313	16.82	81.4		70.71
4	323	2.32	10.9		71.46
5	582	11.55	30.1		87.34
6	682	8.08	18.0		90.76
7	784	2.55	4.9		92.50
8	1,160	2.68	3.5		96.50
9	1,401	8.08	8.7		98.01
10	1,628	6.17	5.7		99.44
11	2,096	710.34	513.6		102.06
12	3,449	29.32	12.9		105.44
13	4,713	11.19	3.6		107.04
14	6,048	4.33	1.1		108.73
15	6,505	7.96	1.9		109.31
16	7,682	3.84	0.8		110.55
17	8,320	4.02	0.7		111.13
18	10,380	75.00	10.9	Upper Marker	113.00
19	13,519	0.00	0.0		115.85

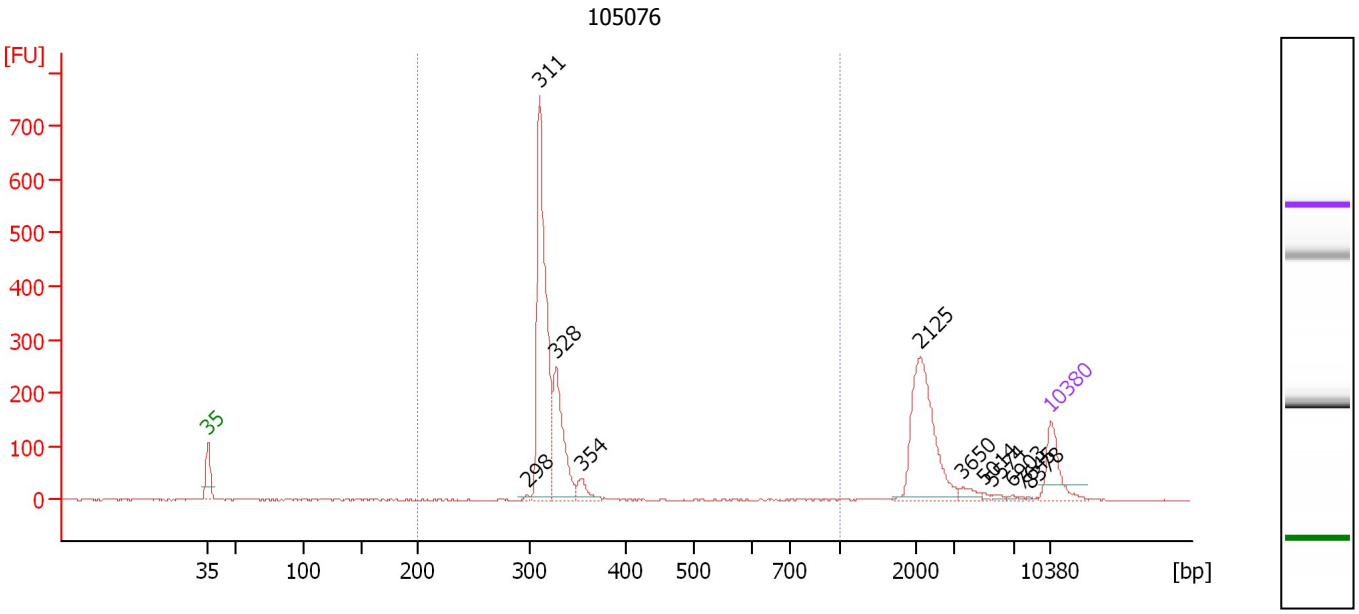
Region table for sample 1 : 104345

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	471	54.59	84.9	223.8	5	42.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : 105076

Number of peaks found:	11	Corr. Area 1:	1,374.5
Noise:	1.0		

Peak table for sample 2 : 105076

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	3.92	19.9		69.48
3	311	655.60	3,192.7		70.54
4	328	245.98	1,135.2		71.91
5	354	37.57	160.7		73.99
6	2,125	273.78	195.2		102.15
7	3,650	17.86	7.4		105.69
8	5,014	4.07	1.2		107.42
9	5,574	5.15	1.4		108.13
10	6,903	3.08	0.7		109.81
11	7,645	1.89	0.4		110.52
12	8,378	1.72	0.3		111.18
13	10,380	75.00	10.9	Upper Marker	113.00

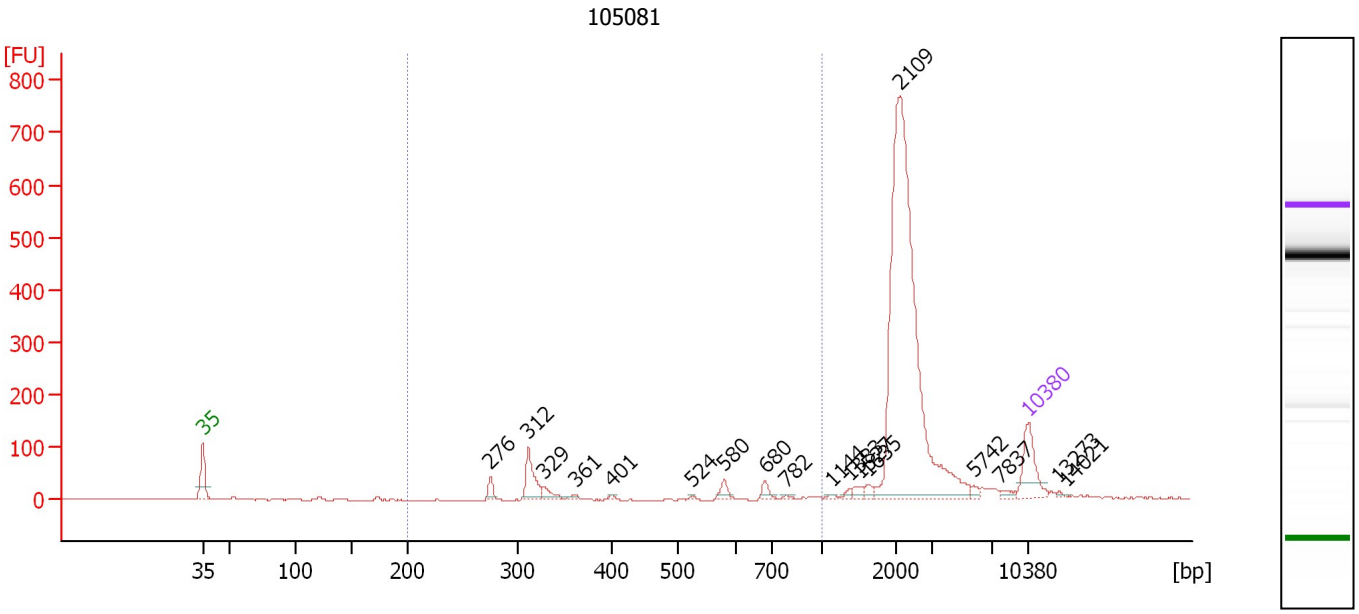
Region table for sample 2 : 105076

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	328	956.20	1,374.5	4,485.3	66	16.6

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : 105081

Number of peaks found:	18	Corr. Area 1:	254.1
Noise:	0.6		

Peak table for sample 3 : 105081

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	276	21.78	119.6		67.42
3	312	83.28	404.1		70.63
4	329	22.14	102.1		71.94
5	361	5.07	21.3		74.56
6	401	2.68	10.1		77.73
7	524	1.65	4.8		84.56
8	580	17.79	46.5		87.22
9	680	15.37	34.2		90.71
10	782	2.30	4.4		92.47
11	1,144	3.23	4.3		96.40
12	1,383	6.17	6.8		97.90
13	1,527	11.09	11.0		98.80
14	1,635	11.10	10.3		99.48
15	2,109	916.57	658.4		102.10
16	5,742	7.46	2.0		108.34
17	7,837	8.11	1.6		110.69
18	10,380	75.00	10.9	Upper Marker	113.00
19	13,273	0.00	0.0		115.62
20	14,021	0.00	0.0		116.30

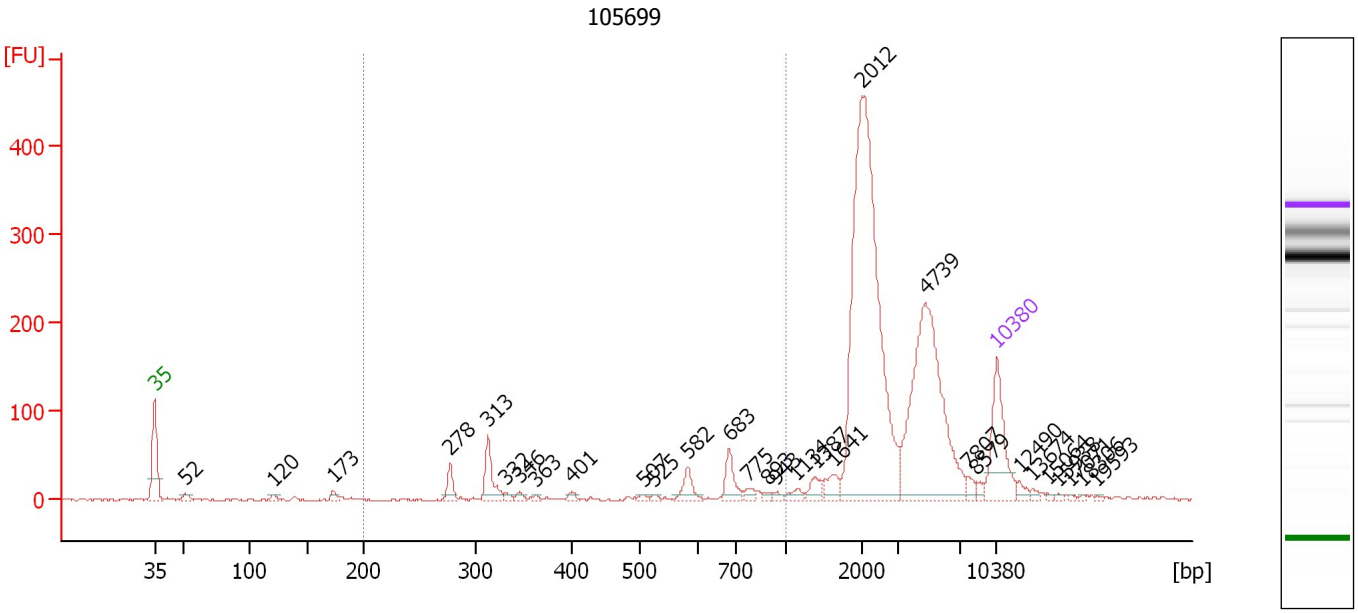
Region table for sample 3 : 105081

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	427	182.50	254.1	771.6	11	40.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : 105699

Number of peaks found:	30	Corr. Area 1:	248.0
Noise:	1.2		

Peak table for sample 4 : 105699

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	8.81	257.2		45.57
3		120	6.09	76.6		52.89
4		173	7.62	66.8		57.84
5		278	22.76	124.2		67.59
6		313	47.61	230.2		70.72
7		332	5.27	24.0		72.21
8		346	5.56	24.3		73.33
9		363	2.53	10.5		74.69
10		401	5.17	19.6		77.67
11		507	3.33	10.0		83.69
12		525	3.03	8.7		84.58
13		582	21.57	56.2		87.33
14		683	29.60	65.7		90.79
15		775	7.55	14.8		92.37
16		893	3.53	6.0		94.01
17		943	4.15	6.7		94.71
18		1,134	7.86	10.5		96.34
19		1,387	14.11	15.4		97.93
20		1,641	17.73	16.4		99.51
21		2,012	507.68	382.3		101.80
22		4,739	284.98	91.1		107.07
23		7,807	8.39	1.6		110.67
24		8,579	4.89	0.9		111.37
25	▶	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...

... Peak table for sample 4 : 105699

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	12,490	0.00	0.0		114.91
27	13,674	0.00	0.0		115.99
28	15,064	0.00	0.0		117.25
29	15,938	0.00	0.0		118.04
30	17,071	0.00	0.0		119.07
31	18,306	0.00	0.0		120.19
32	19,593	0.00	0.0		121.35

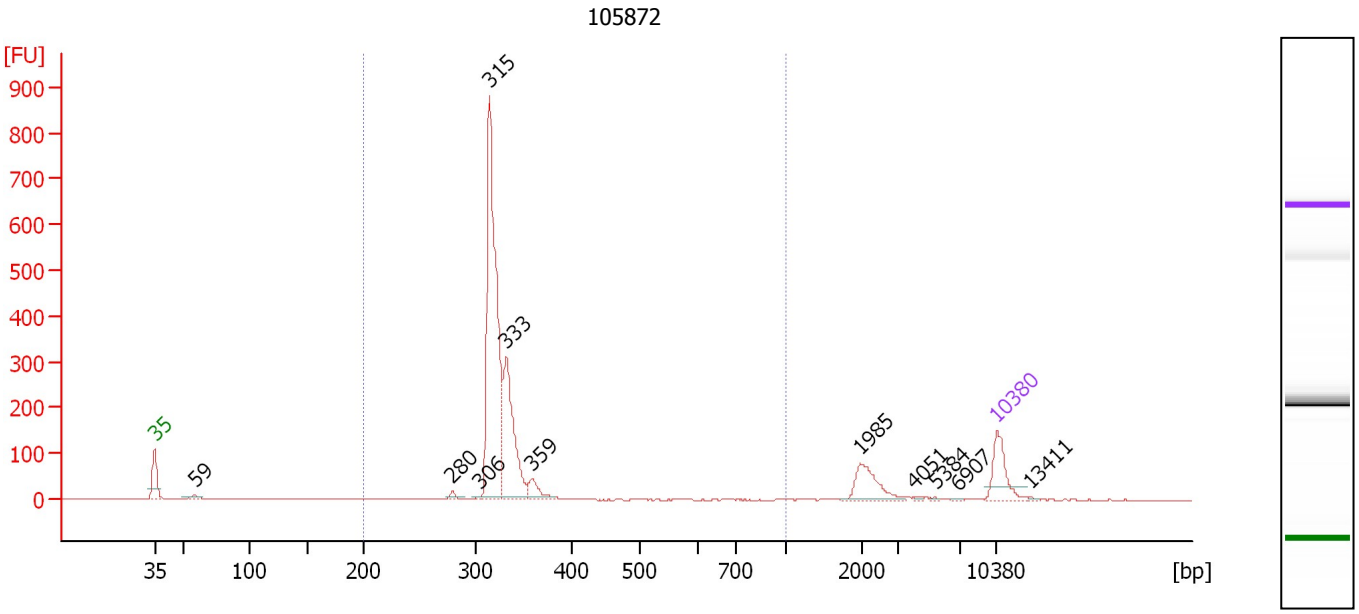
Region table for sample 4 : 105699

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	522	154.00	248.0	568.2	12	39.1

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : 105872

Number of peaks found:	11	Corr. Area 1:	1,633.7
Noise:	1.0		

Peak table for sample 5 : 105872

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	59	15.47	399.4		46.33
3	280	12.60	68.3		67.78
4	306	4.17	20.7		70.09
5	315	793.47	3,814.2		70.87
6	333	317.16	1,444.4		72.26
7	359	47.28	199.4		74.38
8	1,985	81.05	61.9		101.67
9	4,051	2.63	1.0		106.20
10	5,384	1.69	0.5		107.89
11	6,907	1.94	0.4		109.82
12	10,380	75.00	10.9	Upper Marker	113.00
13	13,411	0.00	0.0		115.75

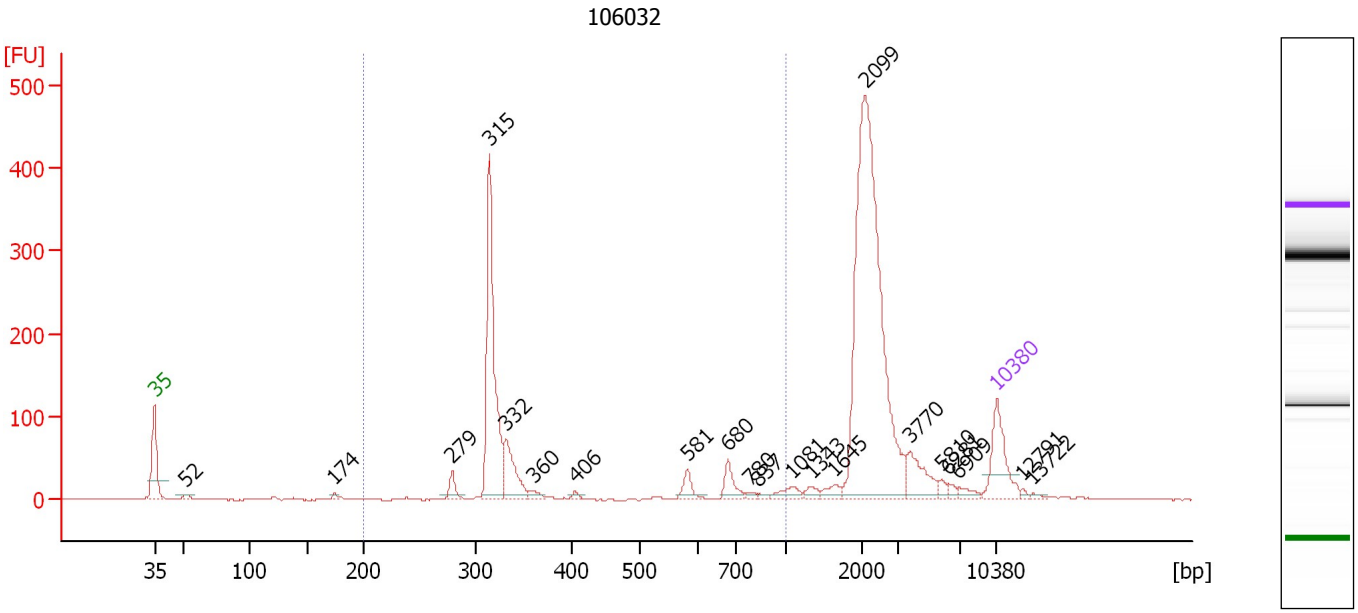
Region table for sample 5 : 105872

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	325	1,155.10	1,633.7	5,384.3	89	5.1

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : 106032

Number of peaks found:	21	Corr. Area 1:	726.2
Noise:	0.6		

Peak table for sample 6 : 106032

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	11.42	329.7		45.63
3	174	6.58	57.4		57.95
4	279	27.36	148.4		67.74
5	315	357.93	1,720.1		70.87
6	332	92.06	420.1		72.21
7	360	12.32	51.8		74.45
8	406	6.21	23.2		77.98
9	581	25.29	65.9		87.32
10	680	33.50	74.7		90.69
11	780	5.85	11.4		92.43
12	837	4.62	8.4		93.23
13	1,081	19.90	27.9		96.01
14	1,343	11.45	12.9		97.65
15	1,645	15.74	14.5		99.54
16	2,099	648.66	468.3		102.07
17	3,770	54.48	21.9		105.85
18	5,810	8.56	2.2		108.43
19	6,281	7.29	1.8		109.03
20	6,909	12.40	2.7		109.82
21	10,380	75.00	10.9	Upper Marker	113.00
22	12,791	0.00	0.0		115.19
23	13,722	0.00	0.0		116.03

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...

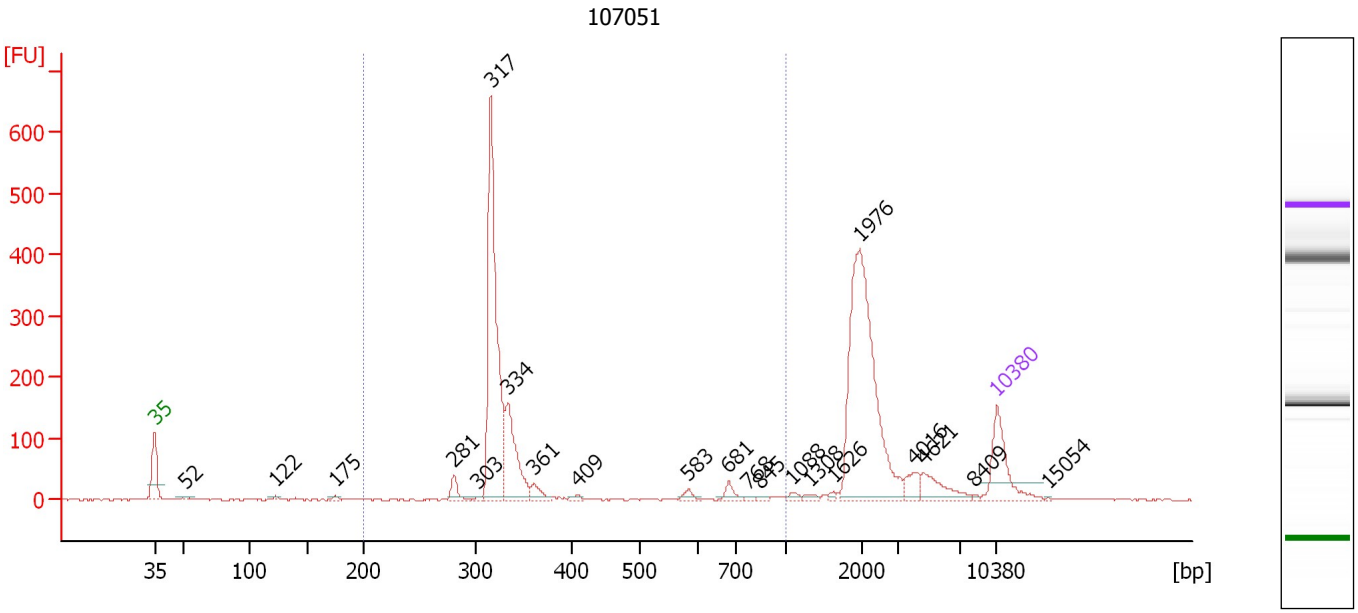
... Region table for sample 6 : 106032

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	399	590.49	726.2	2,576.1	34	40.8

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : 107051

Number of peaks found:	21	Corr. Area 1:	1,152.6
Noise:	1.1		

Peak table for sample 7 : 107051

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	7.79	226.0		45.60
3	122	4.40	54.5		53.06
4	175	4.53	39.2		58.06
5	281	23.52	126.7		67.92
6	303	4.23	21.2		69.91
7	317	460.69	2,204.5		70.98
8	334	144.46	655.6		72.36
9	361	21.33	89.4		74.55
10	409	4.90	18.1		78.18
11	583	10.00	26.0		87.37
12	681	14.76	32.8		90.74
13	768	3.53	7.0		92.27
14	845	3.47	6.2		93.34
15	1,088	5.76	8.0		96.05
16	1,308	5.79	6.7		97.43
17	1,626	3.44	3.2		99.42
18	1,976	402.07	308.3		101.61
19	4,016	24.43	9.2		106.16
20	4,621	39.88	13.1		106.92
21	8,409	2.46	0.4		111.21
22	10,380	75.00	10.9	Upper Marker	113.00
23	15,054	0.00	0.0		117.24

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...

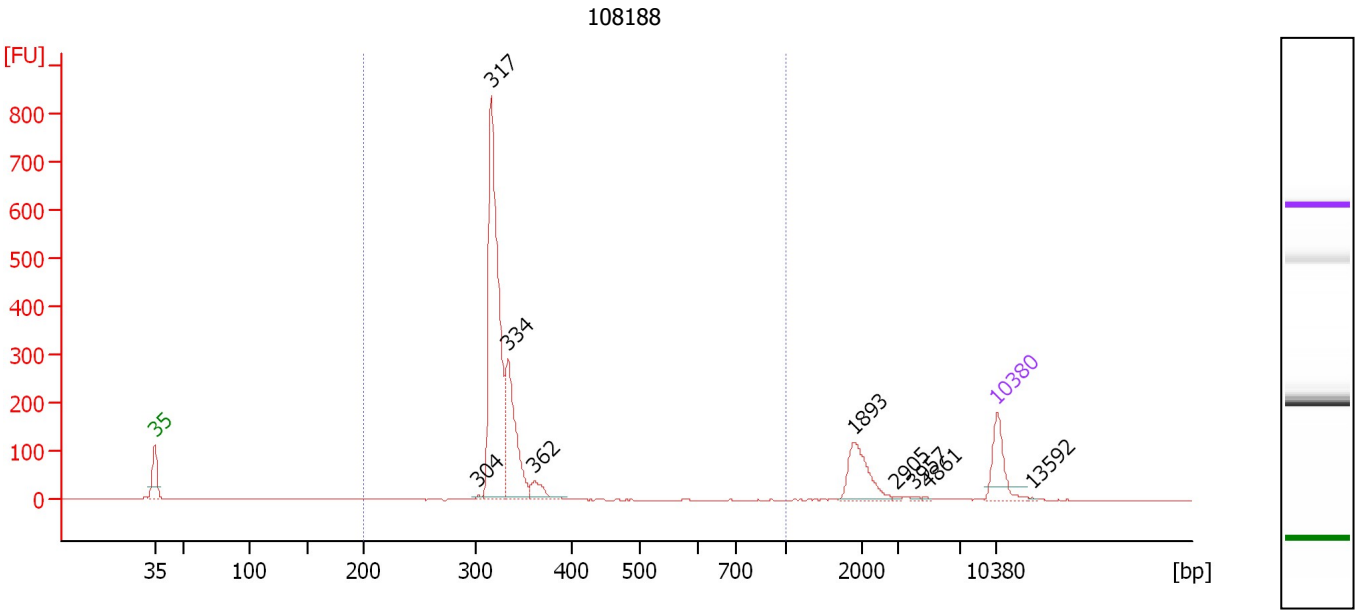
... Region table for sample 7 : 107051

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	355	697.84	1,152.6	3,163.0	48	30.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : 108188

Number of peaks found:	9	Corr. Area 1:	1,573.0
Noise:	0.6		

Peak table for sample 8 : 108188

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	304	5.05	25.2		69.94
3	317	730.38	3,495.1		70.98
4	334	264.78	1,200.5		72.38
5	362	44.84	187.9		74.57
6	1,893	104.79	83.9		101.09
7	2,905	2.64	1.4		104.58
8	3,957	2.49	1.0		106.08
9	4,861	1.84	0.6		107.23
10	10,380	75.00	10.9	Upper Marker	113.00
11	13,592	0.00	0.0		115.91

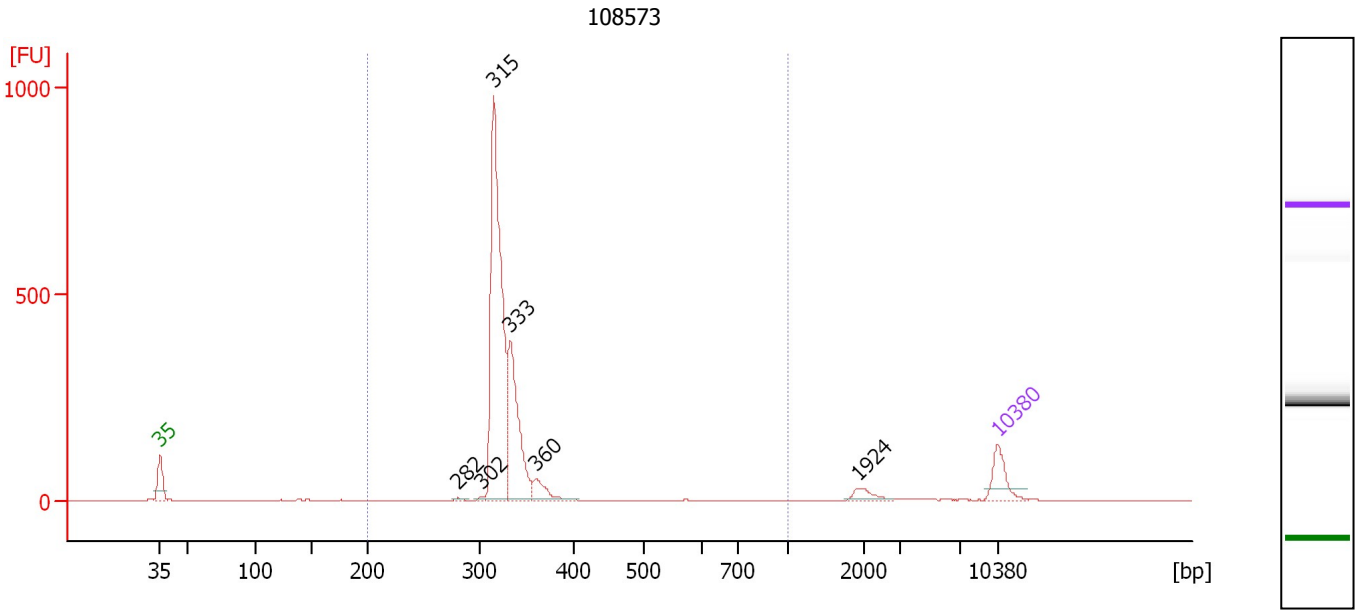
Region table for sample 8 : 108188

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	327	1,034.96	1,573.0	4,791.6	85	4.8

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : 108573

Number of peaks found:	6	Corr. Area 1:	2,033.1
Noise:	0.4		

Peak table for sample 9 : 108573

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	282	5.75	30.9		67.94
3	302	6.85	34.3		69.83
4	315	1,056.21	5,073.2		70.88
5	333	423.55	1,926.1		72.30
6	360	72.80	306.3		74.46
7	1,924	30.35	23.9		101.29
8	10,380	75.00	10.9	Upper Marker	113.00

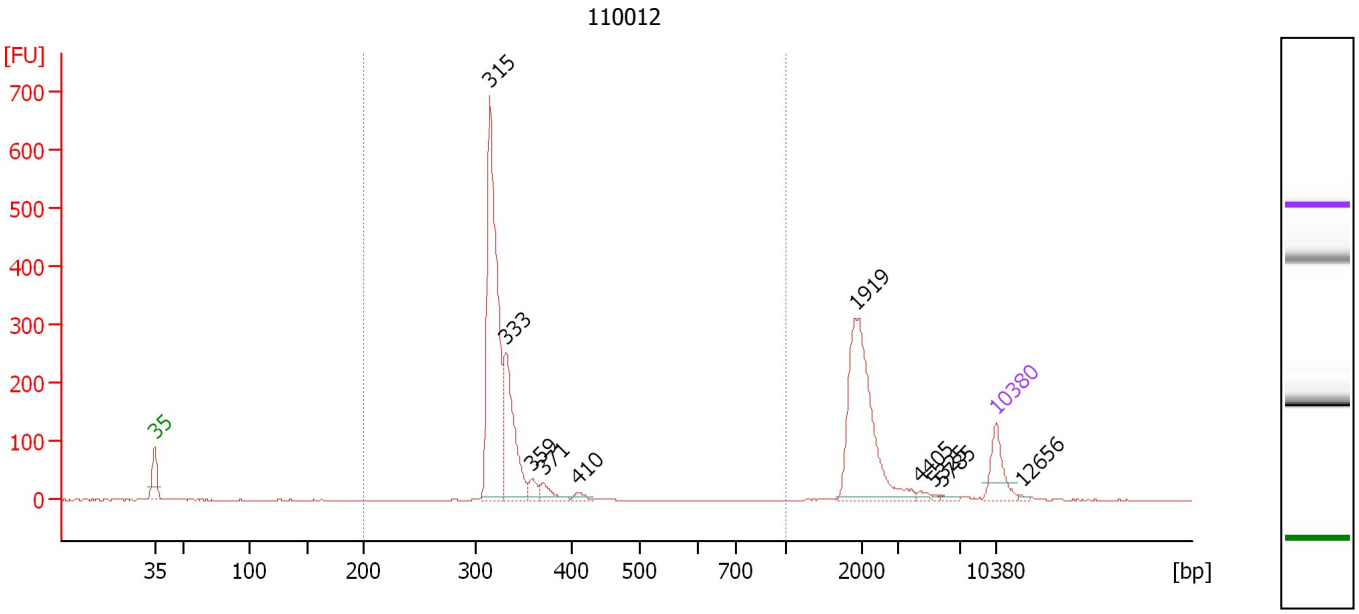
Region table for sample 9 : 108573

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	329	1,570.23	2,033.1	7,271.4	93	10.0

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : 110012

Number of peaks found:	10	Corr. Area 1:	1,399.0
Noise:	0.3		

Peak table for sample 10 : 110012

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	315	827.35	3,974.1		70.88
3	333	304.80	1,388.3		72.26
4	359	30.98	130.7		74.38
5	371	37.41	152.8		75.33
6	410	14.24	52.6		78.24
7	1,919	411.28	324.8		101.25
8	4,405	8.32	2.9		106.65
9	5,325	3.37	1.0		107.81
10	5,785	5.70	1.5		108.40
11	10,380	75.00	10.9	Upper Marker	113.00
12	12,656	0.00	0.0		115.06

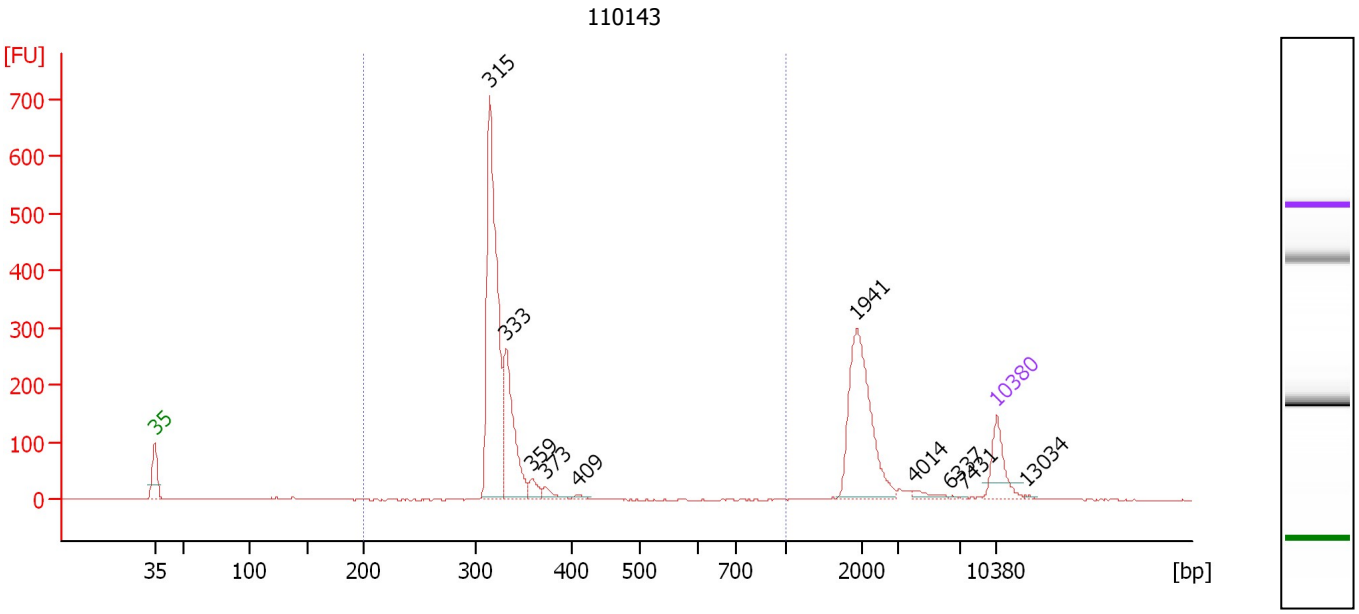
Region table for sample 10 : 110012

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	328	1,212.27	1,399.0	5,604.9	65	6.0

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : 110143

Number of peaks found:	10	Corr. Area 1:	1,380.5
Noise:	0.6		

Peak table for sample 11 : 110143

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		315	709.19	3,409.6		70.86
3		333	268.82	1,224.8		72.25
4		359	28.52	120.3		74.39
5		373	19.80	80.5		75.46
6		409	6.09	22.6		78.13
7		1,941	314.73	245.6		101.40
8		4,014	14.35	5.4		106.16
9		6,337	2.11	0.5		109.10
10		7,431	1.45	0.3		110.33
11	▶	10,380	75.00	10.9	Upper Marker	113.00
12		13,034	0.00	0.0		115.41

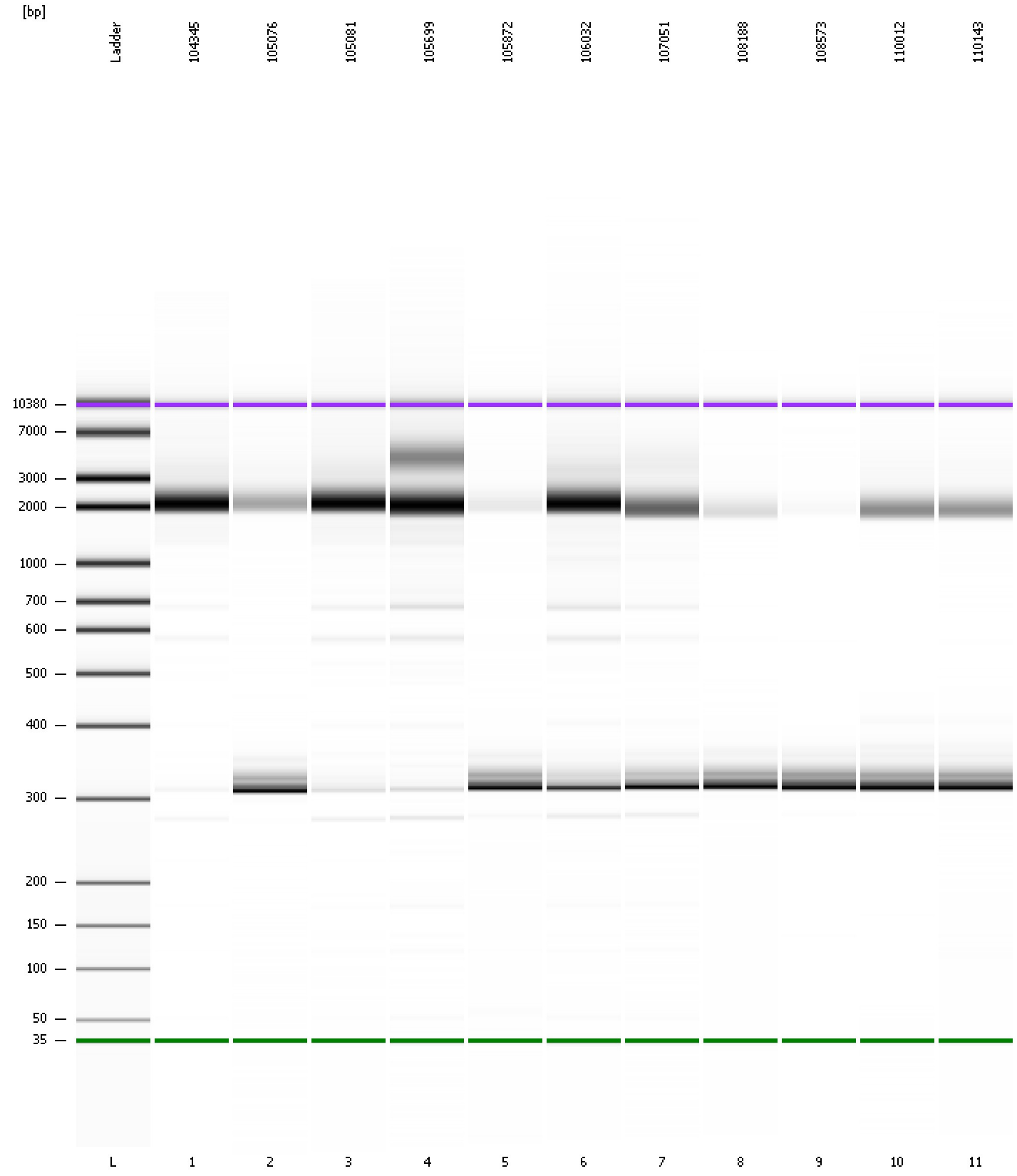
Region table for sample 11 : 110143

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	329	1,043.14	1,380.5	4,826.3	65	8.9

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-31\2019-10-31_002.xad

Created: 10/31/2019 3:31:44 PM
Modified: 10/31/2019 4:12:06 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		10/31/2019 4:12:05 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\2019-10-31\2019-10-31_002.xad)		Instrument	Run		10/31/2019 3:31:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		10/31/2019 3:31:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		10/31/2019 3:31:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		10/31/2019 3:31:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		10/31/2019 3:31:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		10/31/2019 3:31:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		10/31/2019 3:31:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1