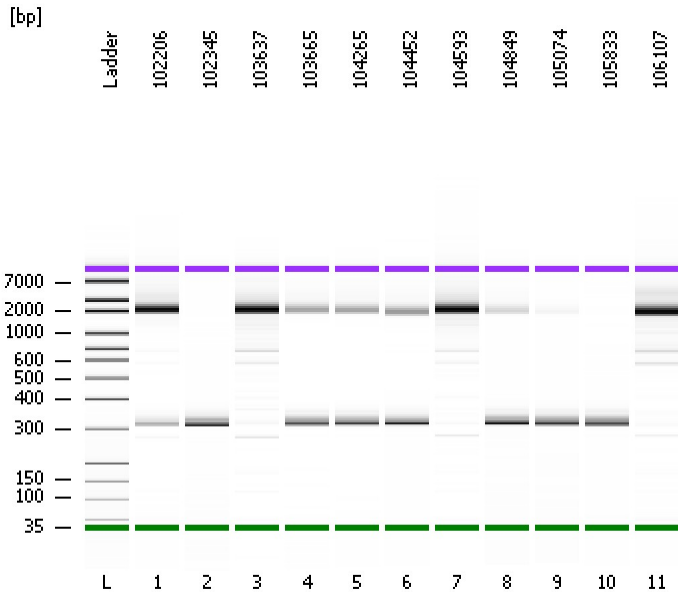


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
Modified: 11/1/2019 12:19:32 PM

**Electrophoresis File Run Summary**



Instrument Information:

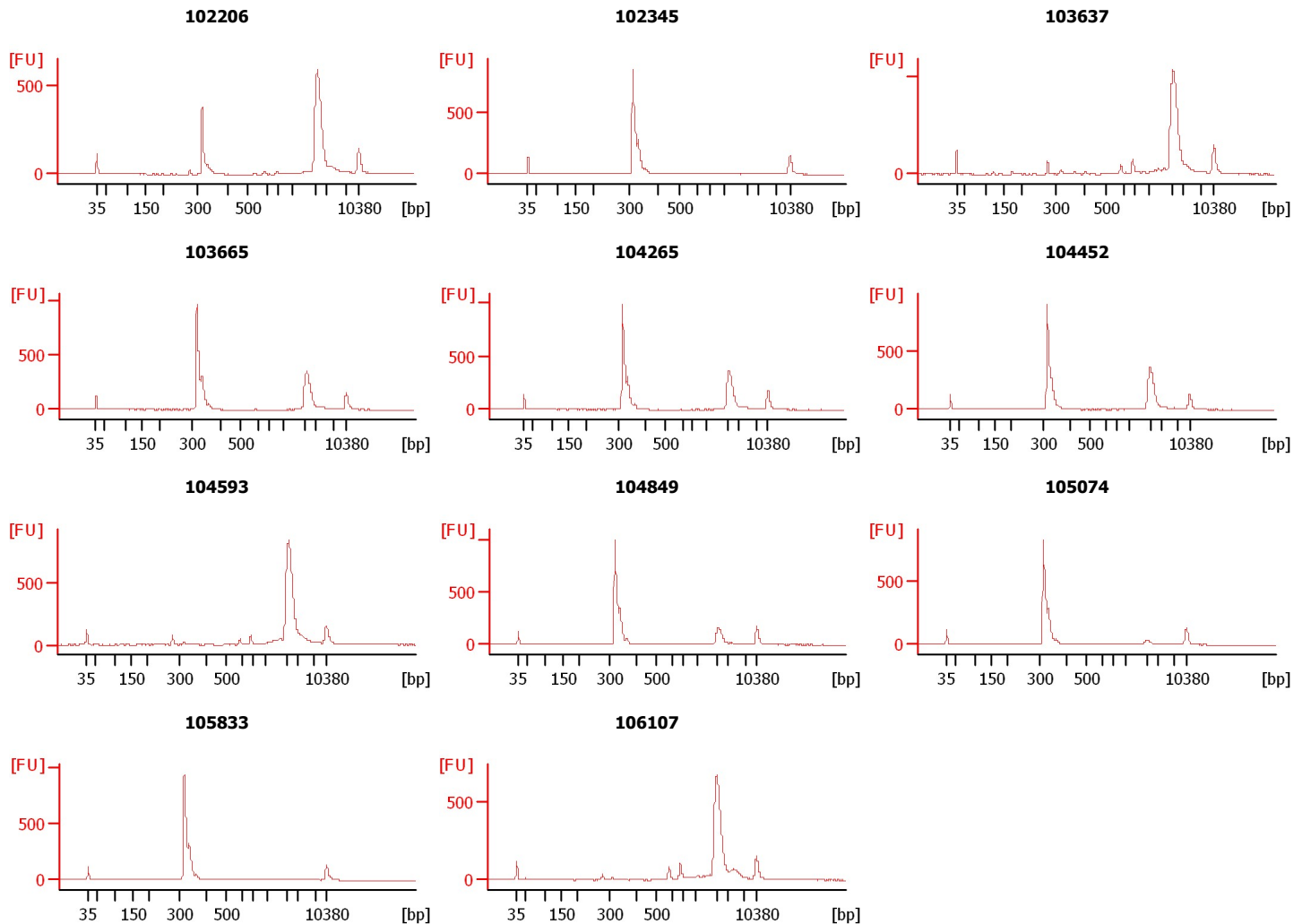
Instrument Name: DE34903152      Firmware: C.01.069  
Serial#: DE34903152      Type: G2938C

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:



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electrophoresis File Run Summary (Chip Summary)**

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
102206		<input type="checkbox"/>	✓			
102345		<input type="checkbox"/>	✓			
103637		<input type="checkbox"/>	✓			
103665		<input type="checkbox"/>	✓			
104265		<input type="checkbox"/>	✓			
104452		<input type="checkbox"/>	✓			
104593		<input type="checkbox"/>	✓			
104849		<input type="checkbox"/>	✓			
105074		<input type="checkbox"/>	✓			
105833		<input type="checkbox"/>	✓			
106107		<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\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
Modified: 11/1/2019 12:19:32 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

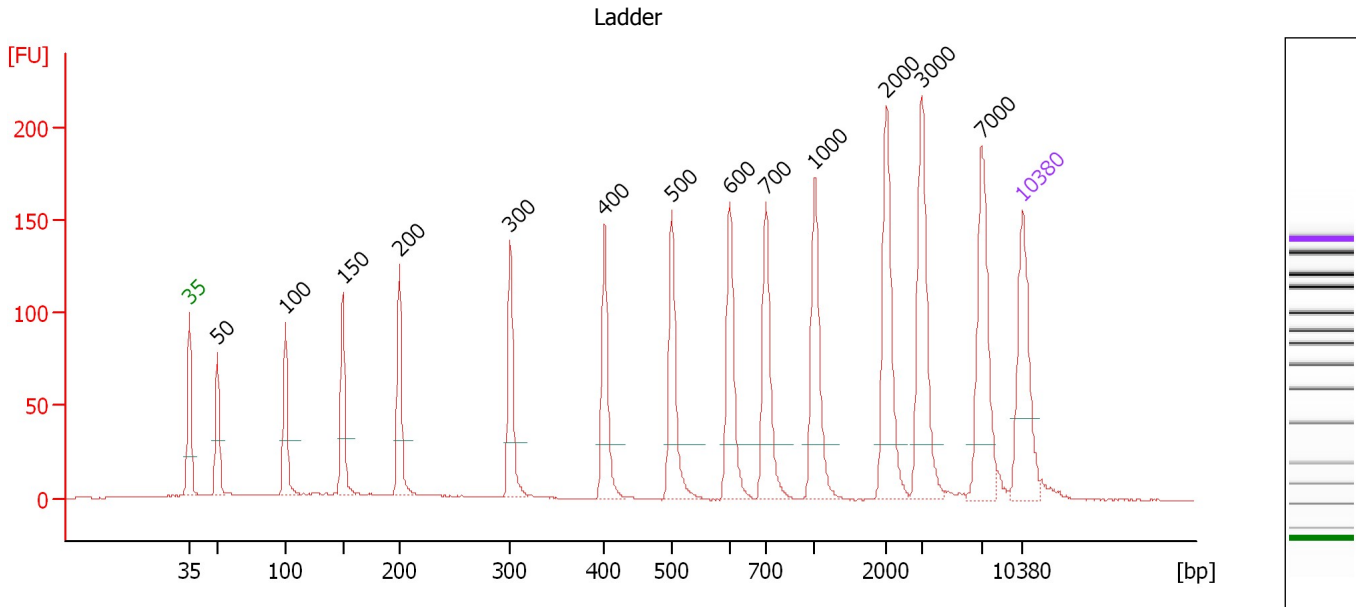
**Ladder**

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\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.3

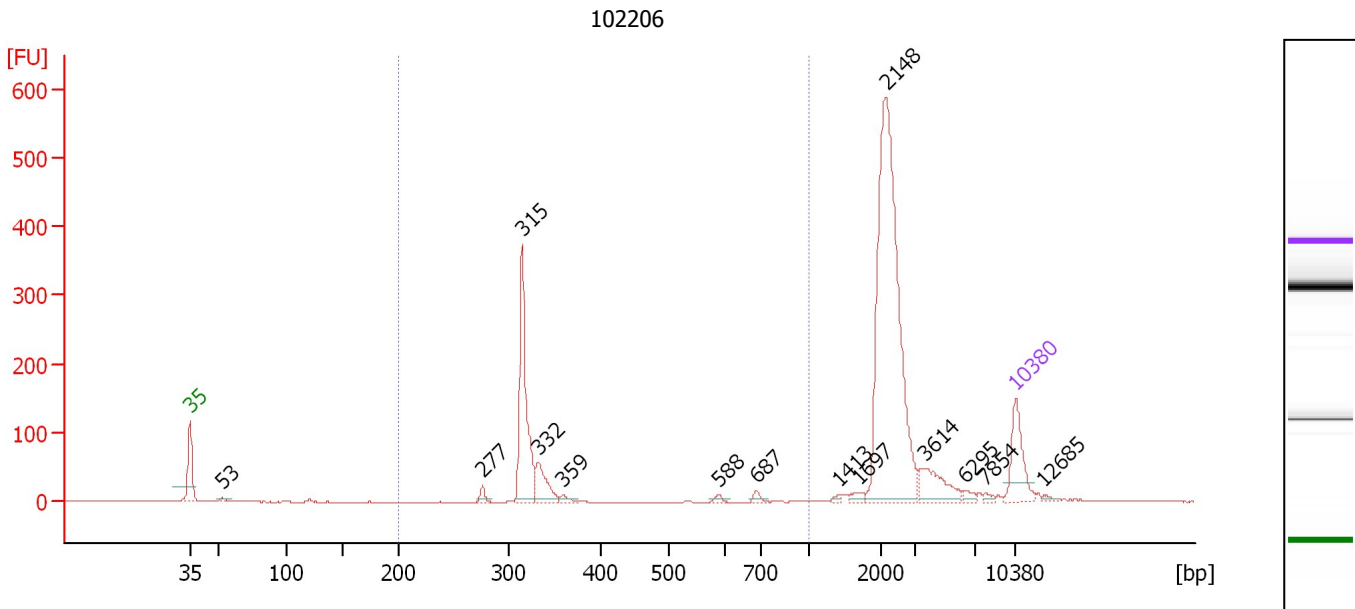
**Peak table for Ladder**

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	45.37
3	100	150.00	2,272.7	Ladder Peak	51.11
4	150	150.00	1,515.2	Ladder Peak	55.90
5	200	150.00	1,136.4	Ladder Peak	60.65
6	300	150.00	757.6	Ladder Peak	69.96
7	400	150.00	568.2	Ladder Peak	77.89
8	500	150.00	454.5	Ladder Peak	83.53
9	600	150.00	378.8	Ladder Peak	88.41
10	700	150.00	324.7	Ladder Peak	91.41
11	1,000	150.00	227.3	Ladder Peak	95.53
12	2,000	150.00	113.6	Ladder Peak	101.58
13	3,000	150.00	75.8	Ladder Peak	104.49
14	7,000	150.00	32.5	Ladder Peak	109.55
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : 102206**

Number of peaks found: 14                      Corr. Area 1: 504.4  
 Noise: 0.3

**Peak table for sample 1 : 102206**

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	53	8.87	251.8		45.76
3	277	15.13	82.9		67.78
4	315	283.65	1,363.2		71.17
5	332	72.04	328.4		72.53
6	359	10.85	45.8		74.61
7	588	7.29	18.8		87.81
8	687	9.49	20.9		91.02
9	1,413	4.36	4.7		98.03
10	1,697	10.26	9.2		99.75
11	2,148	626.96	442.2		102.01
12	3,614	64.11	26.9		105.27
13	6,295	8.72	2.1		108.66
14	7,854	5.66	1.1		110.42
15	10,380	75.00	10.9	Upper Marker	113.00
16	12,685	0.00	0.0		115.35

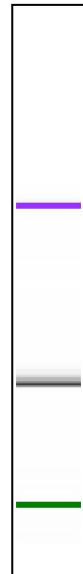
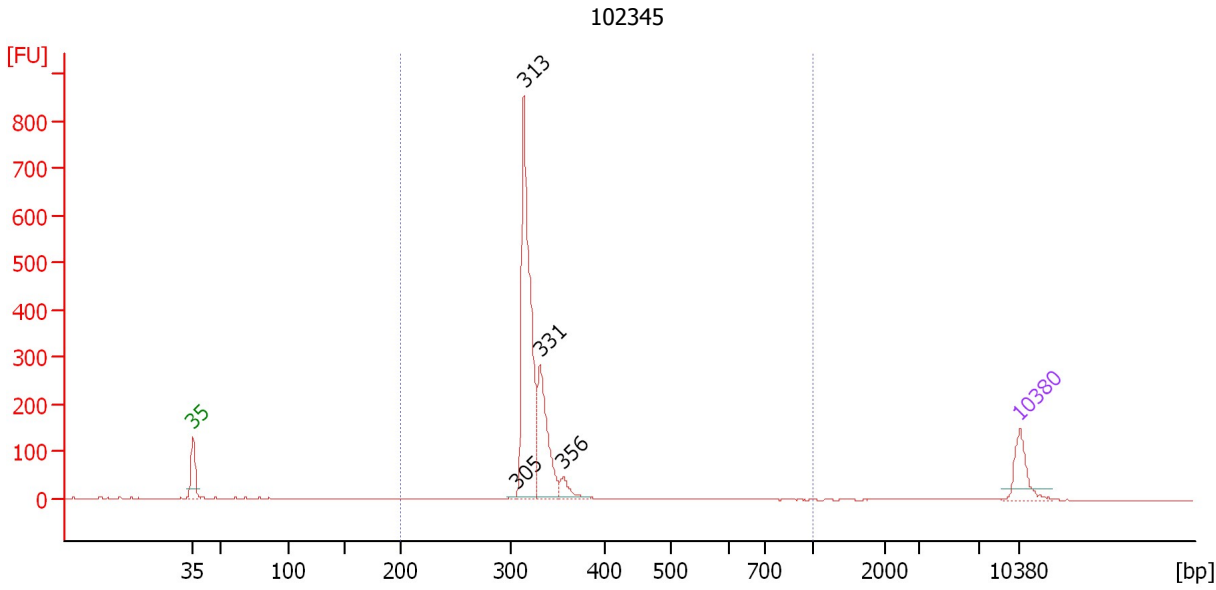
**Region table for sample 1 : 102206**

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	343	385.56	504.4	1,783.1	26	26.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : 102345**

Number of peaks found: 4                      Corr. Area 1: 1,560.8  
 Noise: 0.9

**Peak table for sample 2 : 102345**

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	305	4.00	19.9		70.35
3	313	760.10	3,674.2		71.03
4	331	288.15	1,320.9		72.38
5	356	47.79	203.6		74.37
6	10,380	75.00	10.9	Upper Marker	113.00

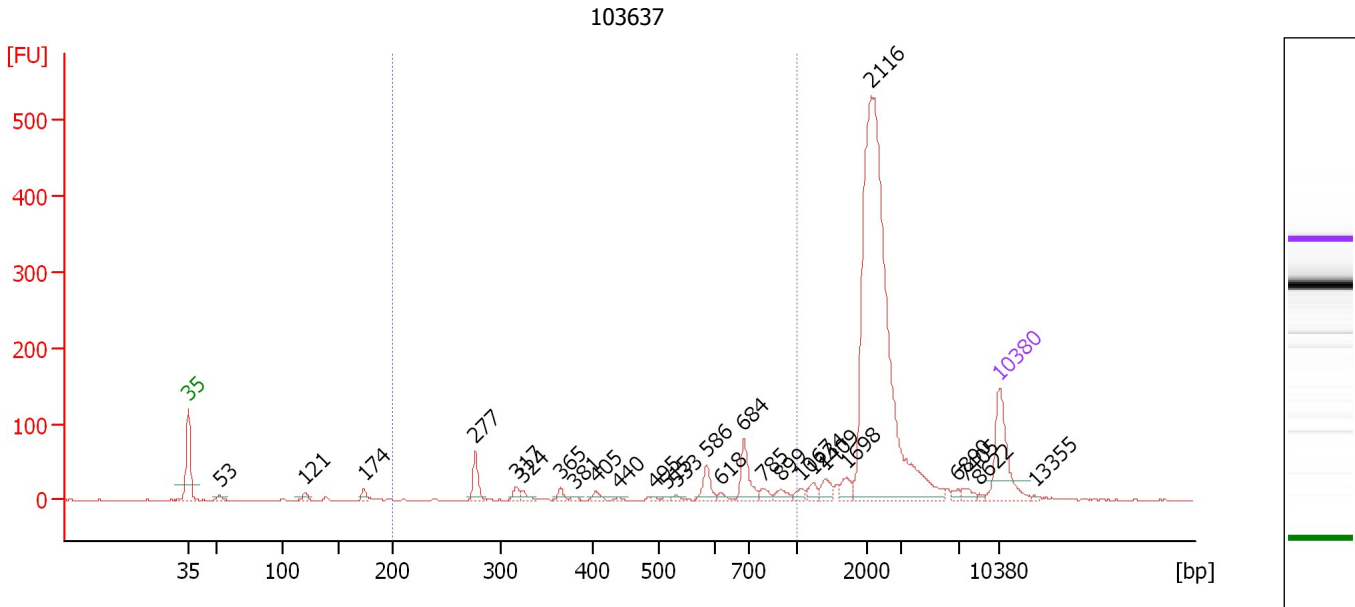
**Region table for sample 2 : 102345**

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	1,106.41	1,560.8	5,165.2	97	11.4

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : 103637**

Number of peaks found: 27                      Corr. Area 1: 297.1  
 Noise: 0.5

**Peak table for sample 3 : 103637**

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	53	7.85	224.5		45.71
3	121	10.38	130.0		53.12
4	174	11.25	97.9		58.18
5	277	39.16	214.4		67.79
6	317	12.29	58.7		71.33
7	324	9.37	43.8		71.88
8	365	9.46	39.2		75.15
9	381	3.82	15.2		76.34
10	405	8.74	32.7		78.18
11	440	3.29	11.3		80.16
12	495	2.75	8.4		83.24
13	515	2.66	7.8		84.25
14	533	4.82	13.7		85.13
15	586	24.99	64.6		87.75
16	618	6.22	15.2		88.95
17	684	44.89	99.5		90.92
18	785	9.98	19.3		92.58
19	899	11.03	18.6		94.14
20	1,067	8.17	11.6		95.94
21	1,234	11.50	14.1		96.95
22	1,409	15.82	17.0		98.01
23	1,698	14.55	13.0		99.75
24	2,116	627.81	449.6		101.92
25	6,890	4.70	1.0		109.41

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

### Electropherogram Summary Continued ...

#### ... Peak table for sample 3 : 103637

Peak	Size [bp]	Conc. [pg/ $\mu$ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	7,405	8.68	1.8		109.96
27	8,622	2.51	0.4		111.21
28	10,380	75.00	10.9	Upper Marker	113.00
29	13,355	0.00	0.0		116.04

#### Region table for sample 3 : 103637

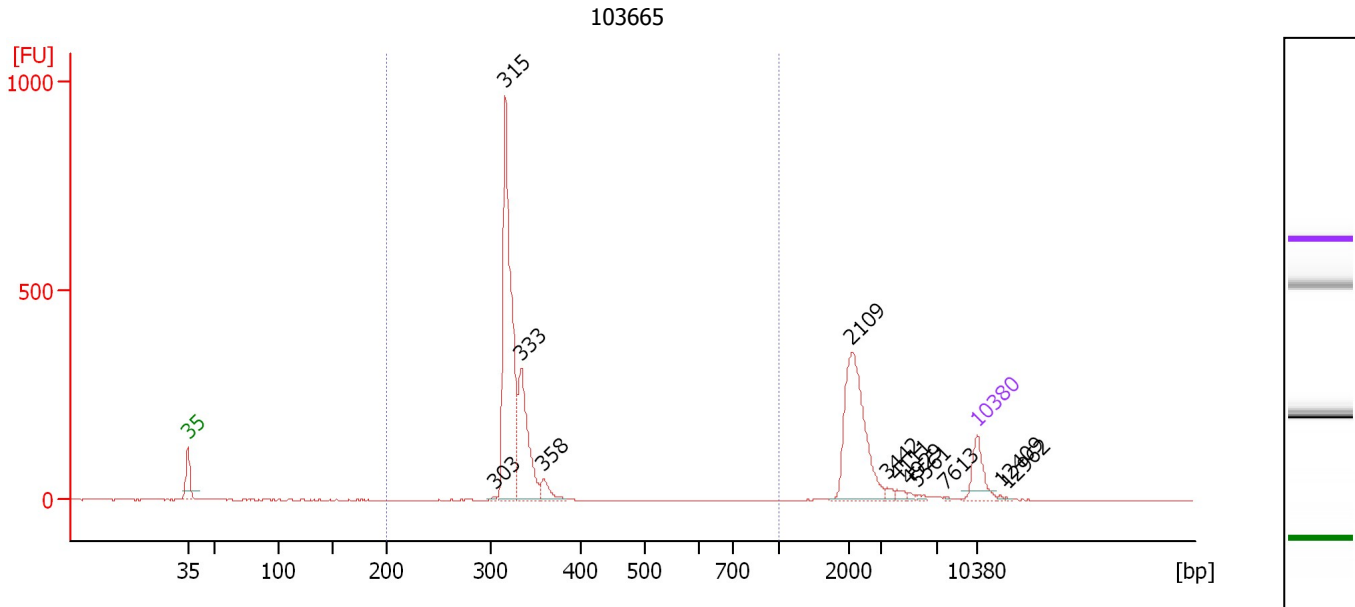
From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ $\mu$ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	565	188.12	297.1	639.7	16	35.5



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : 103665**

Number of peaks found: 12                      Corr. Area 1: 1,714.0  
 Noise: 0.7

**Peak table for sample 4 : 103665**

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	3.60	18.0		70.19
3	315	838.46	4,028.4		71.18
4	333	322.16	1,465.1		72.59
5	358	47.47	200.9		74.57
6	2,109	355.49	255.4		101.90
7	3,442	11.97	5.3		105.05
8	4,111	11.14	4.1		105.90
9	4,929	5.19	1.6		106.93
10	5,561	5.26	1.4		107.73
11	7,613	2.17	0.4		110.18
12	10,380	75.00	10.9	Upper Marker	113.00
13	12,409	0.00	0.0		115.07
14	12,962	0.00	0.0		115.63

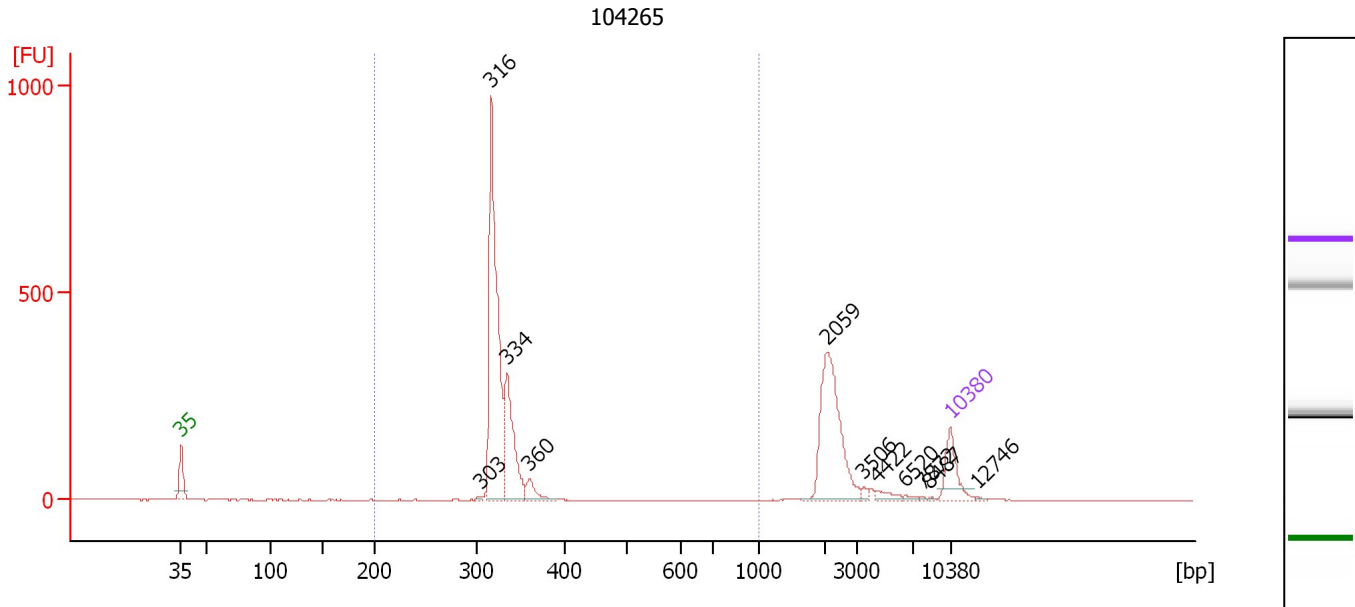
**Region table for sample 4 : 103665**

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	1,209.52	1,714.0	5,629.0	66	5.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : 104265**

Number of peaks found: 11                      Corr. Area 1: 1,695.4  
 Noise: 0.7

**Peak table for sample 5 : 104265**

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	4.13	20.7		70.18
3	316	744.68	3,568.9		71.24
4	334	274.46	1,245.8		72.64
5	360	40.74	171.4		74.72
6	2,059	318.42	234.3		101.75
7	3,506	8.60	3.7		105.13
8	4,422	17.74	6.1		106.29
9	6,520	5.96	1.4		108.94
10	7,872	2.17	0.4		110.44
11	8,487	1.90	0.3		111.07
12	10,380	75.00	10.9	Upper Marker	113.00
13	12,746	0.00	0.0		115.41

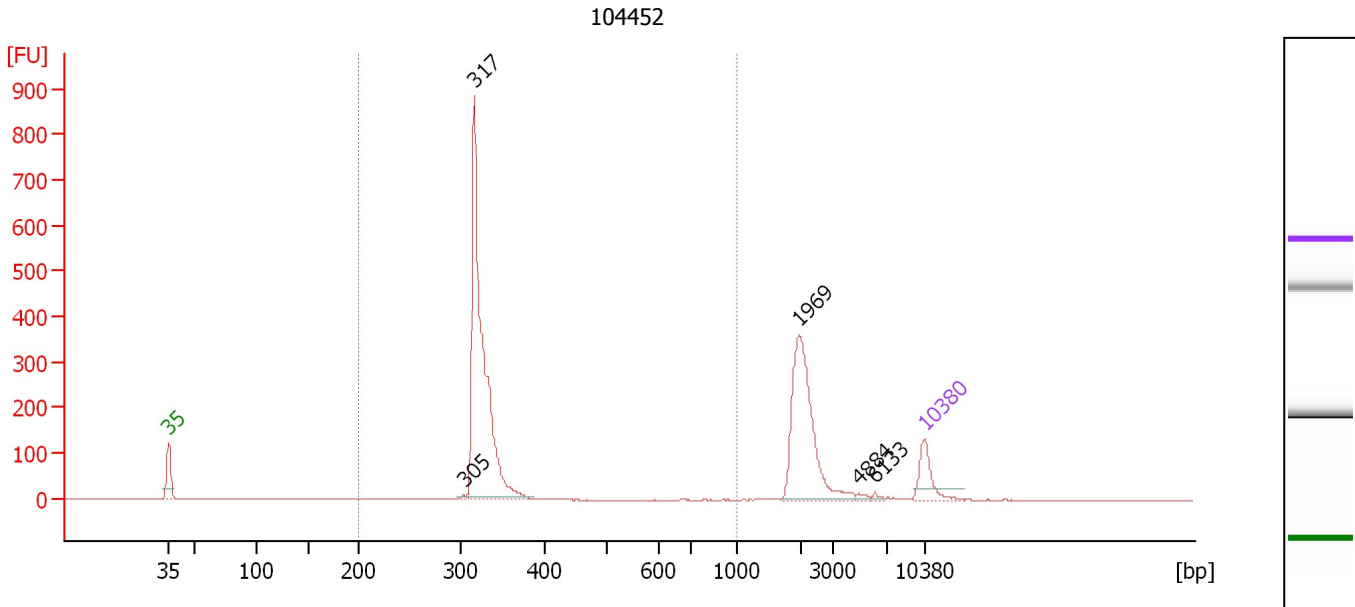
**Region table for sample 5 : 104265**

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	1,072.67	1,695.4	4,978.3	65	9.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : 104452**

Number of peaks found: 5                      Corr. Area 1: 1,454.3  
 Noise: 0.5

**Peak table for sample 6 : 104452**

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	305	6.29	31.3		70.32
3	317	1,081.84	5,171.3		71.31
4	1,969	400.84	308.5		101.39
5	4,884	8.39	2.6		106.87
6	6,133	5.57	1.4		108.46
7	10,380	75.00	10.9	Upper Marker	113.00

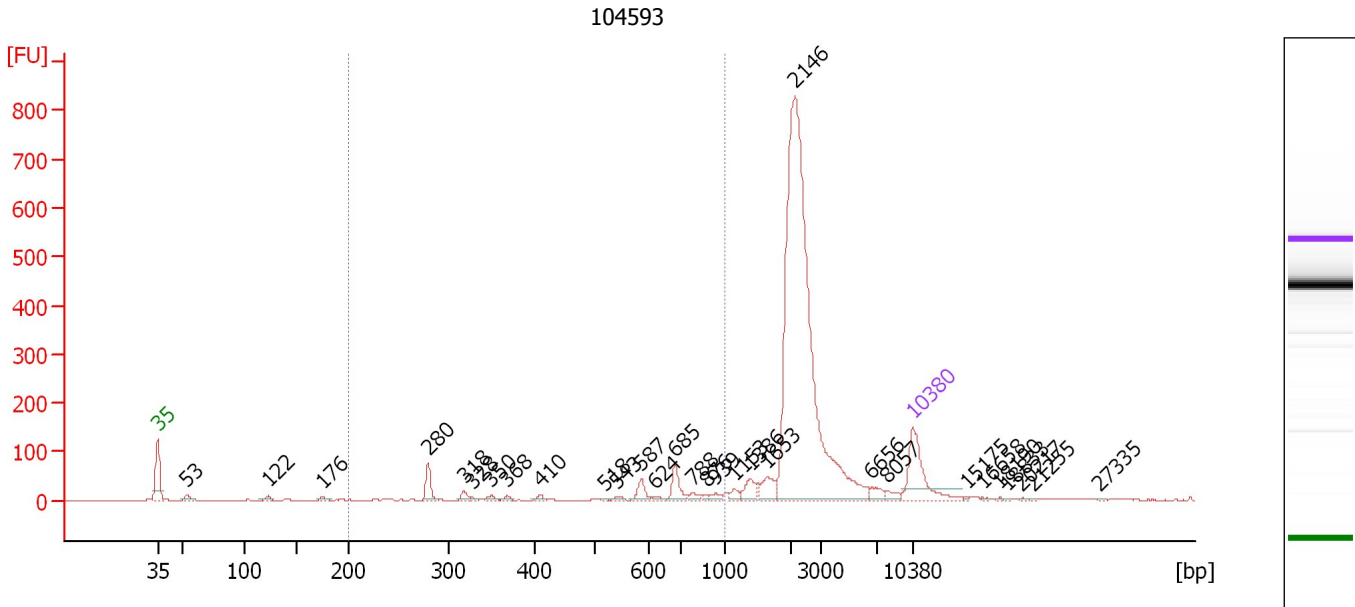
**Region table for sample 6 : 104452**

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	1,071.42	1,454.3	4,981.4	64	4.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : 104593**

Number of peaks found: 30                      Corr. Area 1: 287.1  
 Noise: 0.9

**Peak table for sample 7 : 104593**

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	53	9.91	283.1		45.72
3	122	6.27	77.7		53.24
4	176	5.18	44.7		58.33
5	280	34.72	188.1		68.06
6	318	9.28	44.2		71.39
7	328	2.50	11.6		72.15
8	350	5.53	23.9		73.91
9	368	4.37	18.0		75.38
10	410	4.70	17.4		78.45
11	518	1.79	5.2		84.40
12	543	4.25	11.9		85.62
13	587	18.05	46.6		87.78
14	624	3.68	8.9		89.15
15	685	30.07	66.5		90.96
16	788	7.50	14.4		92.63
17	876	3.75	6.5		93.84
18	939	6.80	11.0		94.69
19	1,153	8.82	11.6		96.46
20	1,386	20.76	22.7		97.87
21	1,653	26.60	24.4		99.48
22	2,146	777.44	548.8		102.01
23	6,656	13.64	3.1		109.12
24	8,057	8.39	1.6		110.63
25	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
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### Electropherogram Summary Continued ...

#### ... Peak table for sample 7 : 104593

Peak	Size [bp]	Conc. [pg/ $\mu$ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	15,175	0.00	0.0		117.89
27	16,658	0.00	0.0		119.40
28	18,190	0.00	0.0		120.97
29	18,833	0.00	0.0		121.62
30	20,217	0.00	0.0		123.04
31	21,255	0.00	0.0		124.10
32	27,335	0.00	0.0		130.30

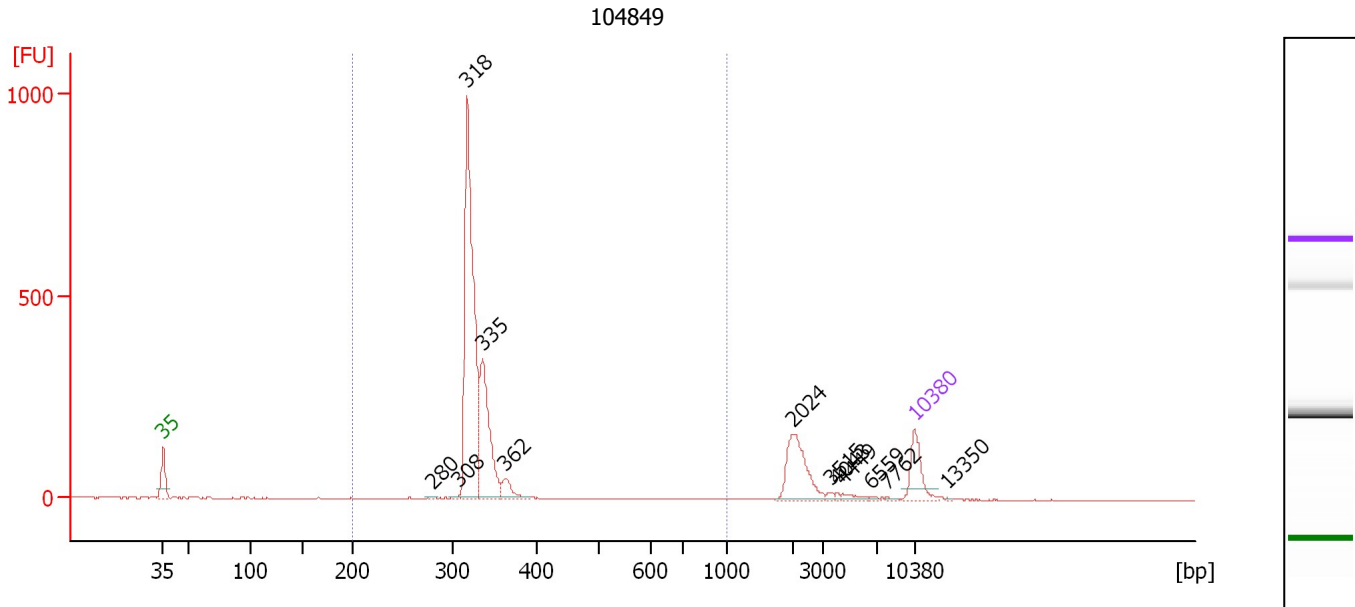
#### Region table for sample 7 : 104593

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ $\mu$ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	561	142.31	287.1	503.4	 11	38.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : 104849**

Number of peaks found: 12                      Corr. Area 1: 1,913.8  
 Noise: 0.9

**Peak table for sample 8 : 104849**

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	280	4.12	22.3		68.11
3	308	5.61	27.6		70.58
4	318	852.18	4,066.5		71.35
5	335	318.63	1,441.1		72.74
6	362	49.34	206.4		74.89
7	2,024	150.86	112.9		101.65
8	3,515	5.69	2.5		105.14
9	4,043	4.53	1.7		105.81
10	4,449	11.94	4.1		106.32
11	6,559	2.69	0.6		108.99
12	7,762	2.71	0.5		110.33
13	10,380	75.00	10.9	Upper Marker	113.00
14	13,350	0.00	0.0		116.03

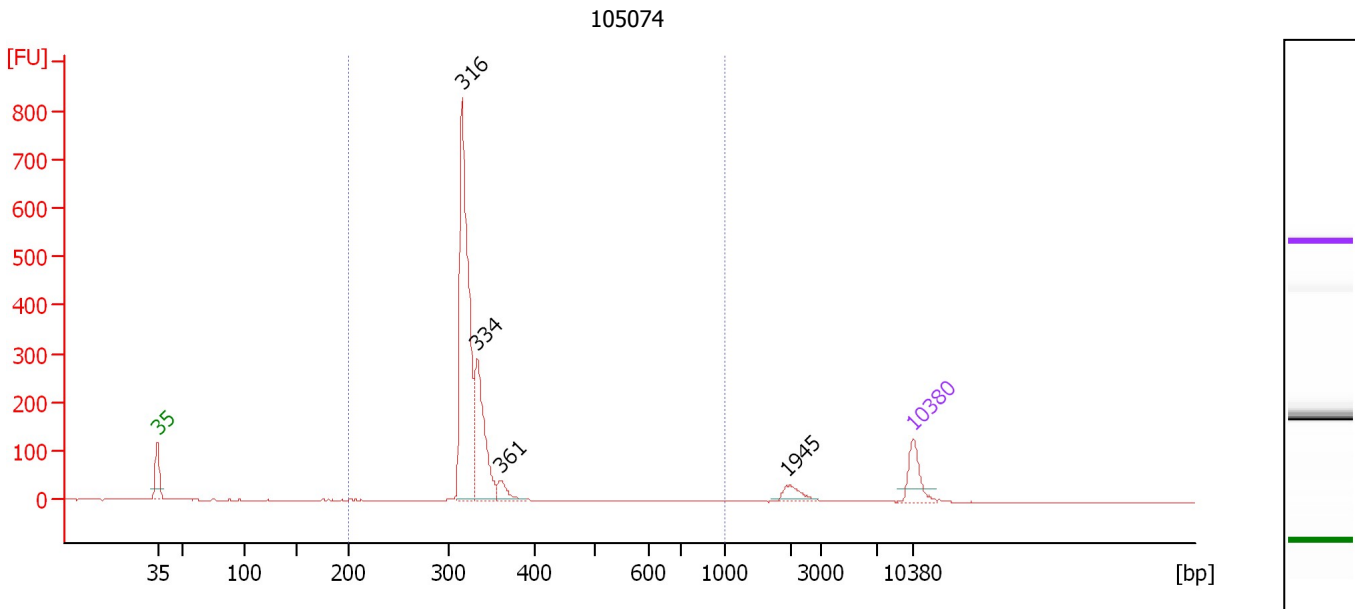
**Region table for sample 8 : 104849**

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	332	1,246.40	1,913.8	5,741.0	80	13.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : 105074**

Number of peaks found: 4                      Corr. Area 1: 1,582.9  
 Noise: 0.4

**Peak table for sample 9 : 105074**

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	903.43	4,330.0		71.24
3	334	352.09	1,598.0		72.64
4	361	50.88	213.7		74.78
5	1,945	37.60	29.3		101.25
6	10,380	75.00	10.9	Upper Marker	113.00

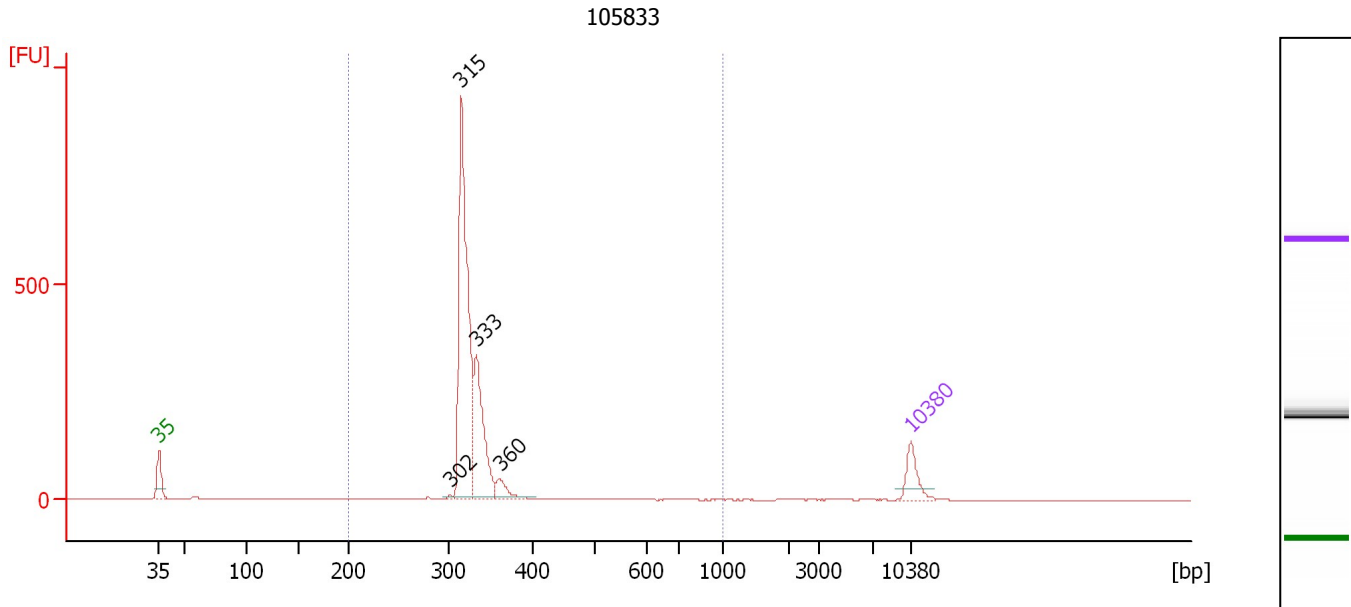
**Region table for sample 9 : 105074**

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	327	1,314.27	1,582.9	6,104.8	94	6.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : 105833**

Number of peaks found: 4                      Corr. Area 1: 1,802.8  
 Noise: 0.3

**Peak table for sample 10 : 105833**

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	302	7.83	39.3		70.12
3	315	1,014.86	4,877.9		71.17
4	333	401.22	1,824.9		72.59
5	360	64.98	273.2		74.74
6	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 10 : 105833**

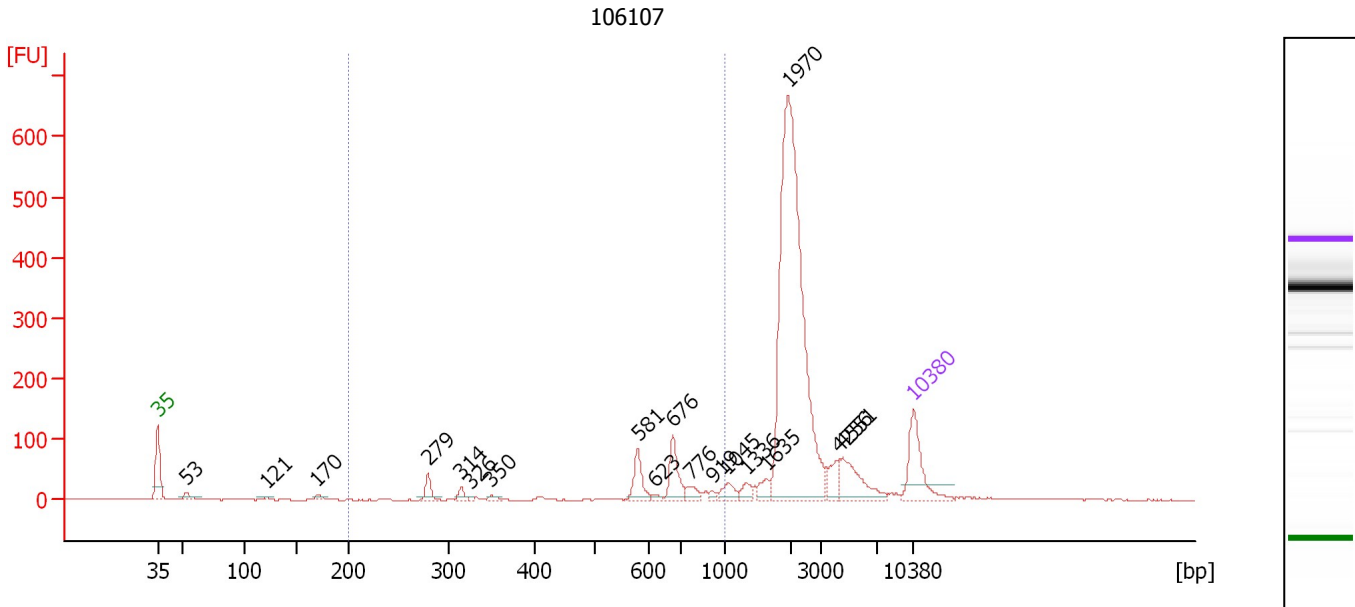
From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	328	1,491.43	1,802.8	6,926.4	97	9.1



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : 106107**

Number of peaks found: 18                      Corr. Area 1: 341.7  
 Noise: 0.3

**Peak table for sample 11 : 106107**

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	53	17.50	505.0		45.66
3	121	4.57	57.3		53.11
4	170	8.06	71.8		57.80
5	279	24.68	134.0		68.02
6	314	12.10	58.3		71.11
7	326	2.95	13.7		72.01
8	350	5.99	25.9		73.93
9	581	44.04	114.8		87.50
10	623	4.51	11.0		89.10
11	676	55.81	125.1		90.70
12	776	16.05	31.4		92.45
13	919	5.27	8.7		94.42
14	1,045	19.37	28.1		95.81
15	1,336	13.31	15.1		97.56
16	1,635	18.04	16.7		99.37
17	1,970	660.68	508.3		101.40
18	4,256	28.14	10.0		106.08
19	4,551	65.18	21.7		106.45
20	10,380	75.00	10.9	Upper Marker	113.00

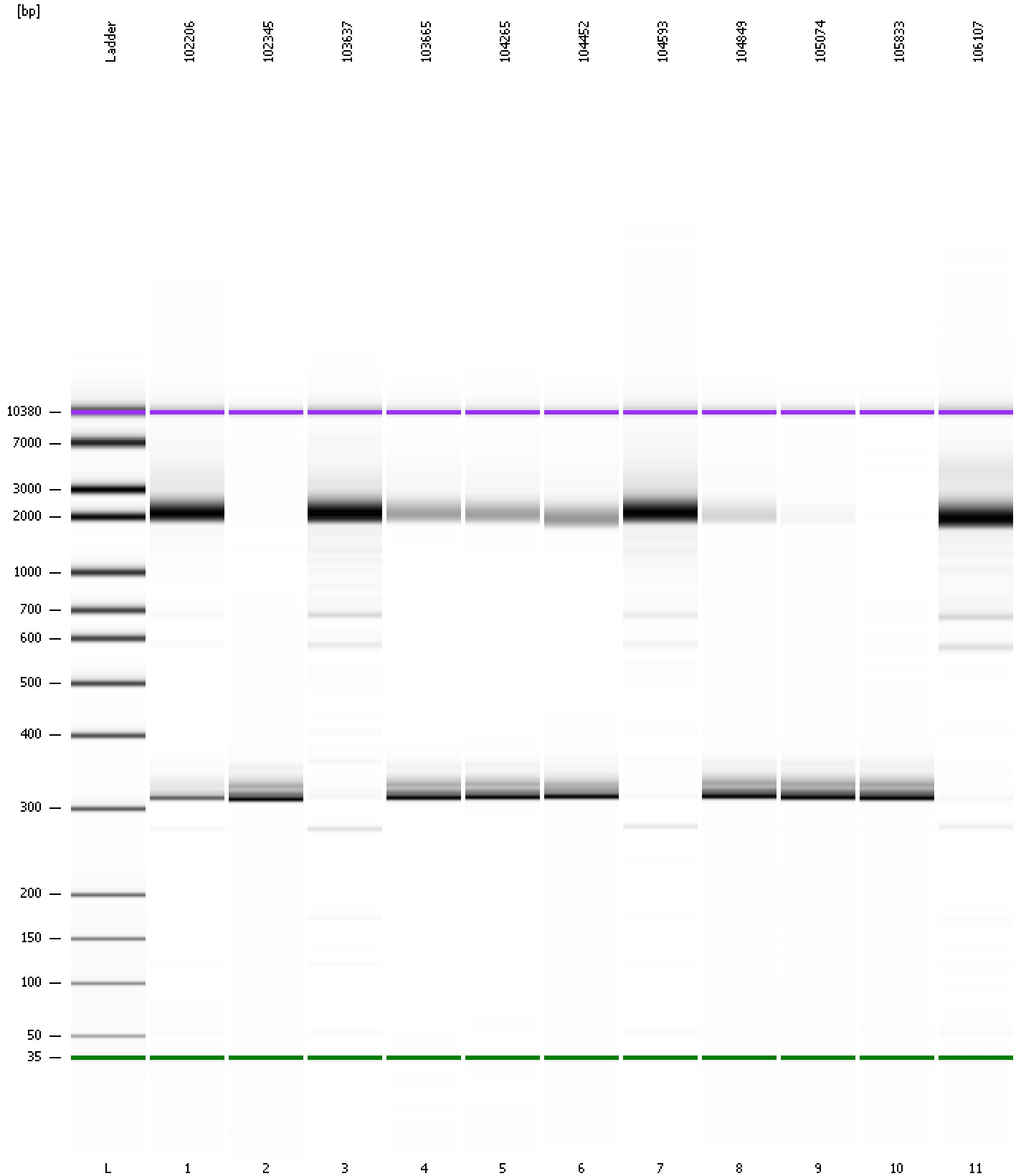
**Region table for sample 11 : 106107**

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	617	187.98	341.7	560.0	15	29.2

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
Modified: 11/1/2019 12:19:32 PM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-11-01\2019-11-01\_002.xad

Created: 11/1/2019 11:39:05 AM  
 Modified: 11/1/2019 12:19:32 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		11/1/2019 12:19:31 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\2019-11-01\2019-11-01_002.xad)		Instrument	Run		11/1/2019 11:39:10 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		11/1/2019 11:39:10 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		11/1/2019 11:39:10 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		11/1/2019 11:39:10 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		11/1/2019 11:39:10 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		11/1/2019 11:39:10 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		11/1/2019 11:39:10 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1