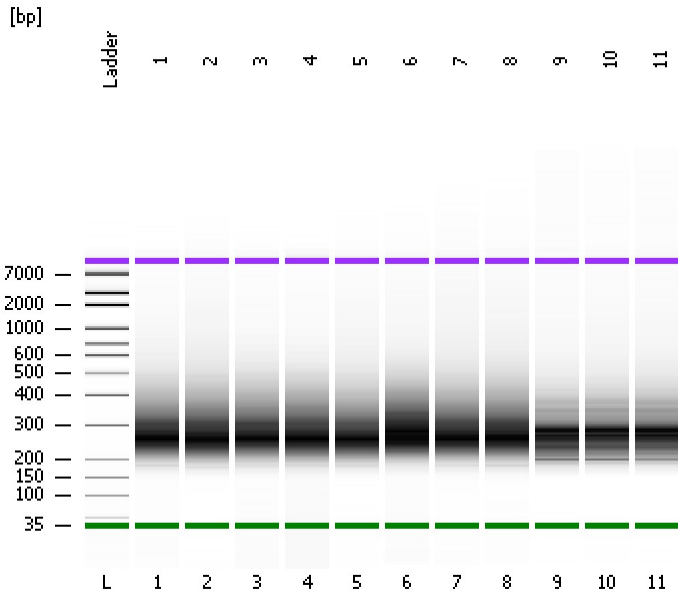


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
Modified: 3/7/2019 5:02:36 PM

**Electrophoresis File Run Summary**



Instrument Information:

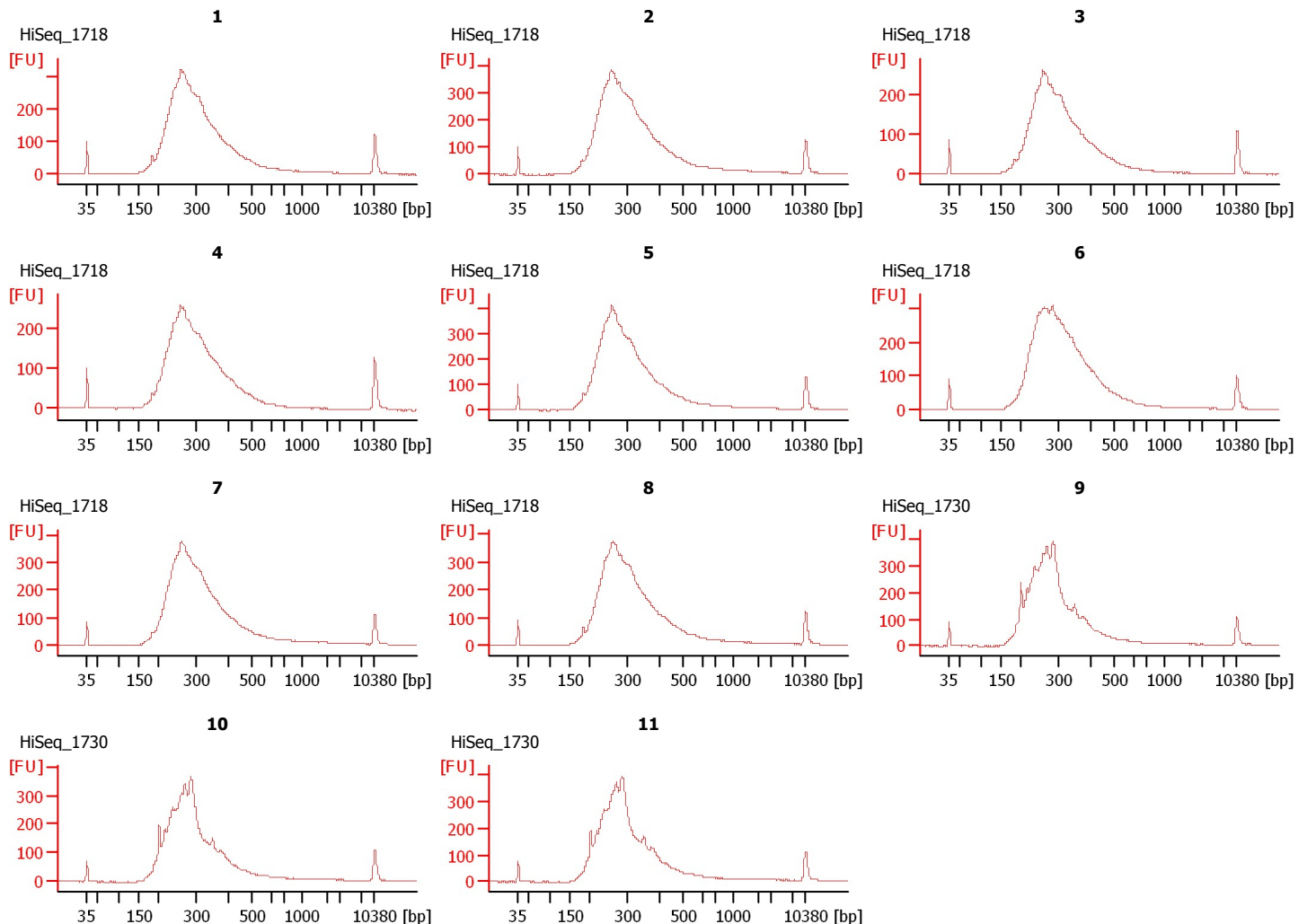
Instrument Name: DE34903152      Firmware: C.01.069  
Serial#: DE34903152      Type: G2938C

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\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
 Modified: 3/7/2019 5:02:36 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
1	HiSeq_1718	<input type="checkbox"/>	✓			
2	HiSeq_1718	<input type="checkbox"/>	✓			
3	HiSeq_1718	<input type="checkbox"/>	✓			
4	HiSeq_1718	<input type="checkbox"/>	✓			
5	HiSeq_1718	<input type="checkbox"/>	✓			
6	HiSeq_1718	<input type="checkbox"/>	✓			
7	HiSeq_1718	<input type="checkbox"/>	✓			
8	HiSeq_1718	<input type="checkbox"/>	✓			
9	HiSeq_1730	<input type="checkbox"/>	✓			
10	HiSeq_1730	<input type="checkbox"/>	✓			
11	HiSeq_1730	<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\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
Modified: 3/7/2019 5:02:36 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

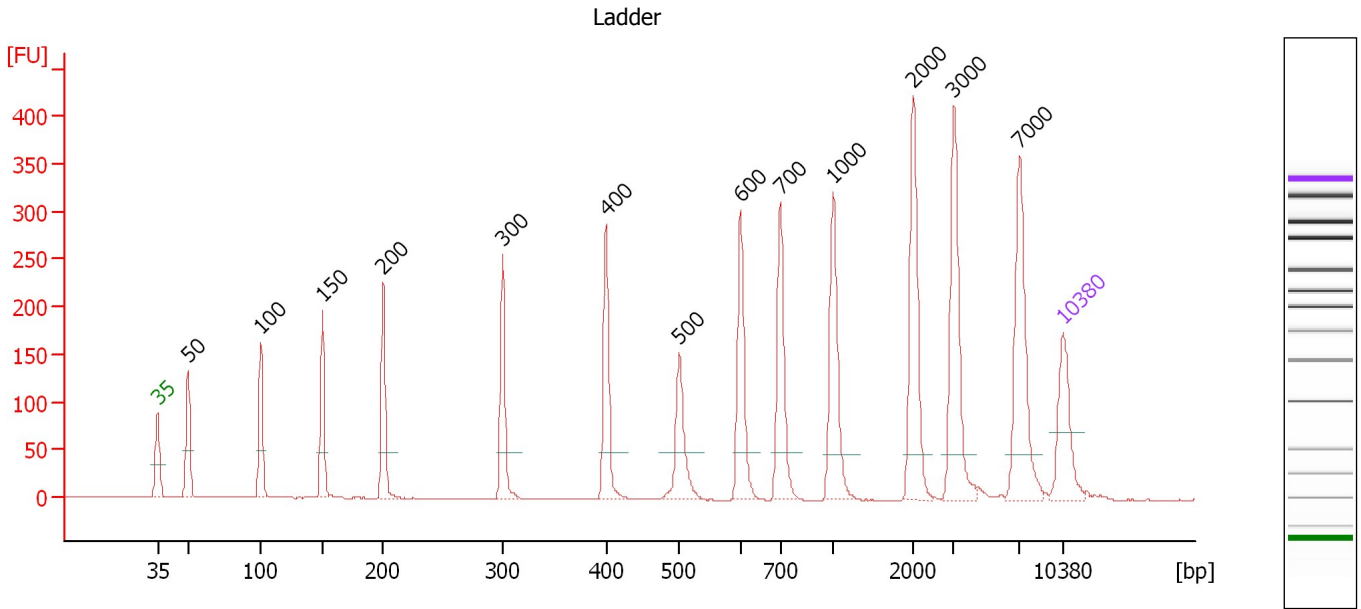
**Ladder**

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\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
 Modified: 3/7/2019 5:02:36 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.6

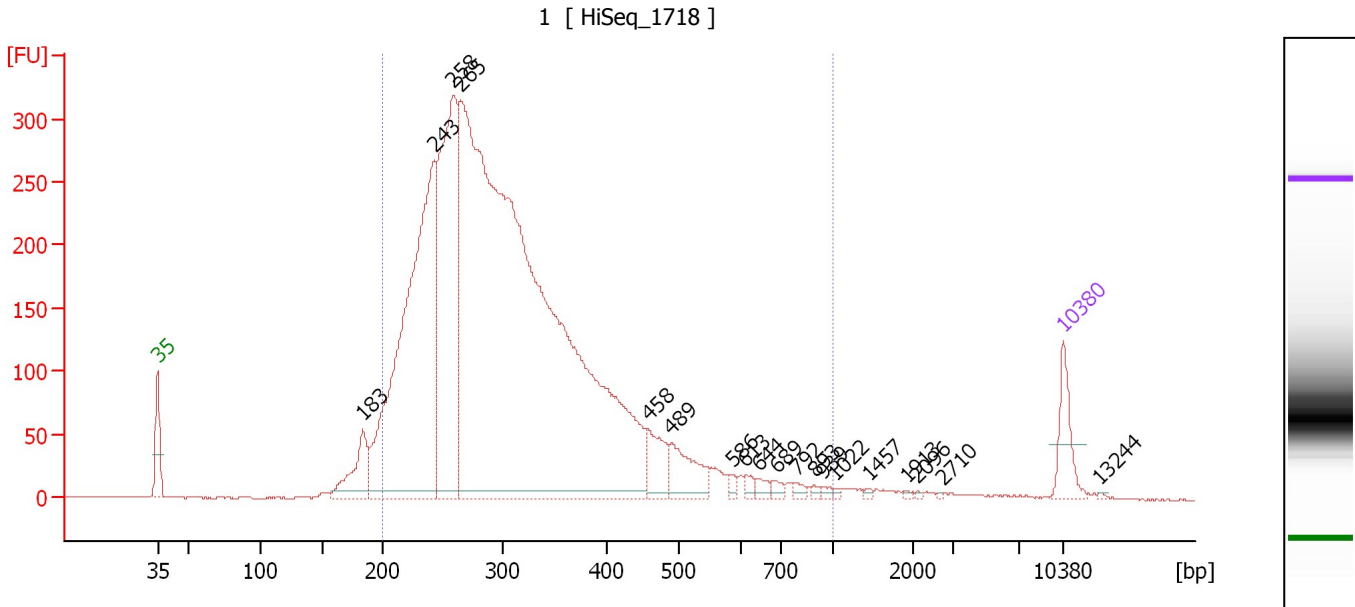
**Peak table for Ladder**

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	45.34
3	100	150.00	2,272.7	Ladder Peak	51.00
4	150	150.00	1,515.2	Ladder Peak	55.76
5	200	150.00	1,136.4	Ladder Peak	60.44
6	300	150.00	757.6	Ladder Peak	69.69
7	400	150.00	568.2	Ladder Peak	77.65
8	500	150.00	454.5	Ladder Peak	83.32
9	600	150.00	378.8	Ladder Peak	88.08
10	700	150.00	324.7	Ladder Peak	91.15
11	1,000	150.00	227.3	Ladder Peak	95.22
12	2,000	150.00	113.6	Ladder Peak	101.41
13	3,000	150.00	75.8	Ladder Peak	104.56
14	7,000	150.00	32.5	Ladder Peak	109.67
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
 Modified: 3/7/2019 5:02:36 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : 1**

Number of peaks found: 19                      Corr. Area 1: 5,880.6  
 Noise: 0.5

**Peak table for sample 1 : 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	183	158.90	1,313.1		58.88
3	243	1,424.48	8,891.2		64.39
4	258	923.53	5,423.8		65.80
5	265	4,214.36	24,126.7		66.42
6	458	108.30	358.4		80.93
7	489	117.94	365.4		82.70
8	586	11.94	30.9		87.42
9	613	15.22	37.6		88.48
10	644	17.57	41.3		89.45
11	689	14.52	31.9		90.81
12	792	12.87	24.6		92.40
13	893	6.88	11.7		93.77
14	939	7.57	12.2		94.39
15	1,022	4.45	6.6		95.36
16	1,457	4.47	4.7		98.05
17	1,913	3.53	2.8		100.87
18	2,096	2.30	1.7		101.71
19	2,710	1.88	1.1		103.65
20	10,380	75.00	10.9	Upper Marker	113.00
21	13,244	0.00	0.0		115.82

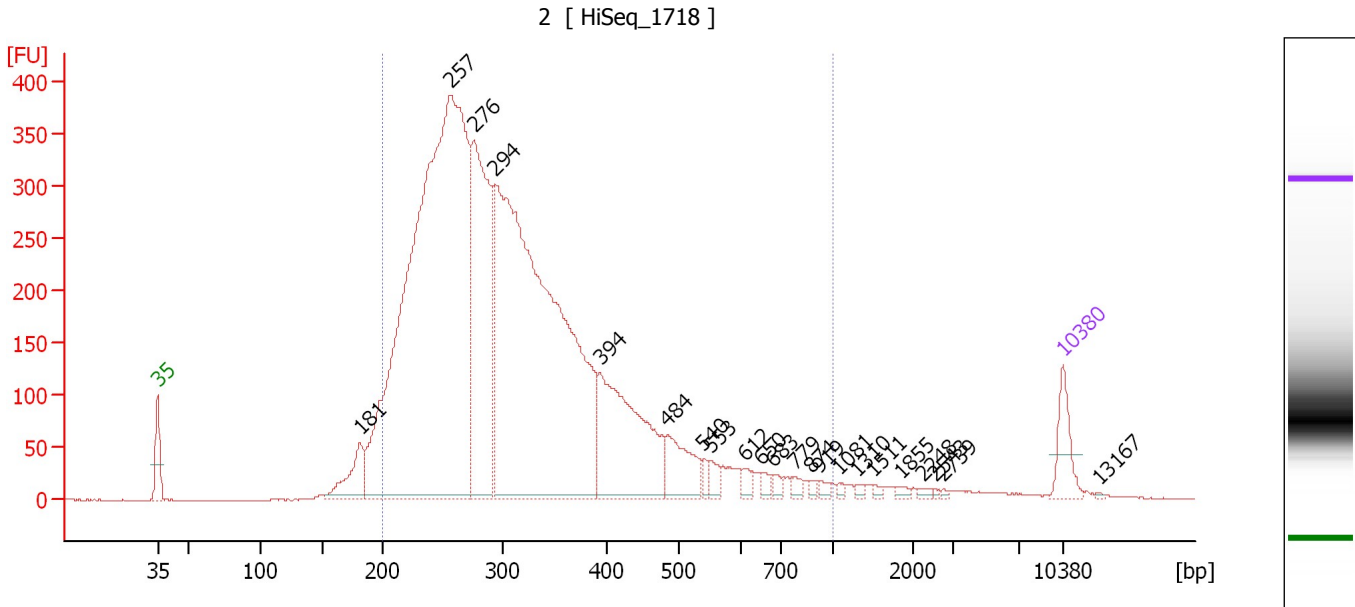
**Region table for sample 1 : 1**

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	319	6,349.51	5,880.6	33,025.4	95	31.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
 Modified: 3/7/2019 5:02:36 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : 2**

Number of peaks found: 22                      Corr. Area 1: 7,450.5  
 Noise: 0.5

**Peak table for sample 2 : 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	181	150.41	1,260.6		58.64
3	257	3,266.16	19,279.1		65.68
4	276	817.27	4,481.5		67.50
5	294	2,461.03	12,703.1		69.09
6	394	539.09	2,071.3		77.20
7	484	152.24	476.3		82.43
8	540	21.97	61.6		85.22
9	553	30.45	83.4		85.84
10	612	24.42	60.4		88.46
11	650	19.44	45.3		89.61
12	683	18.92	42.0		90.63
13	779	16.69	32.5		92.22
14	874	9.21	16.0		93.51
15	919	13.19	21.7		94.13
16	1,081	7.70	10.8		95.72
17	1,310	8.13	9.4		97.14
18	1,511	6.15	6.2		98.38
19	1,855	9.75	8.0		100.51
20	2,248	7.98	5.4		102.19
21	2,543	3.52	2.1		103.12
22	2,739	3.88	2.1		103.74
23	10,380	75.00	10.9	Upper Marker	113.00
24	13,167	0.00	0.0		115.75

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
Modified: 3/7/2019 5:02:36 PM

**Electropherogram Summary Continued ...**

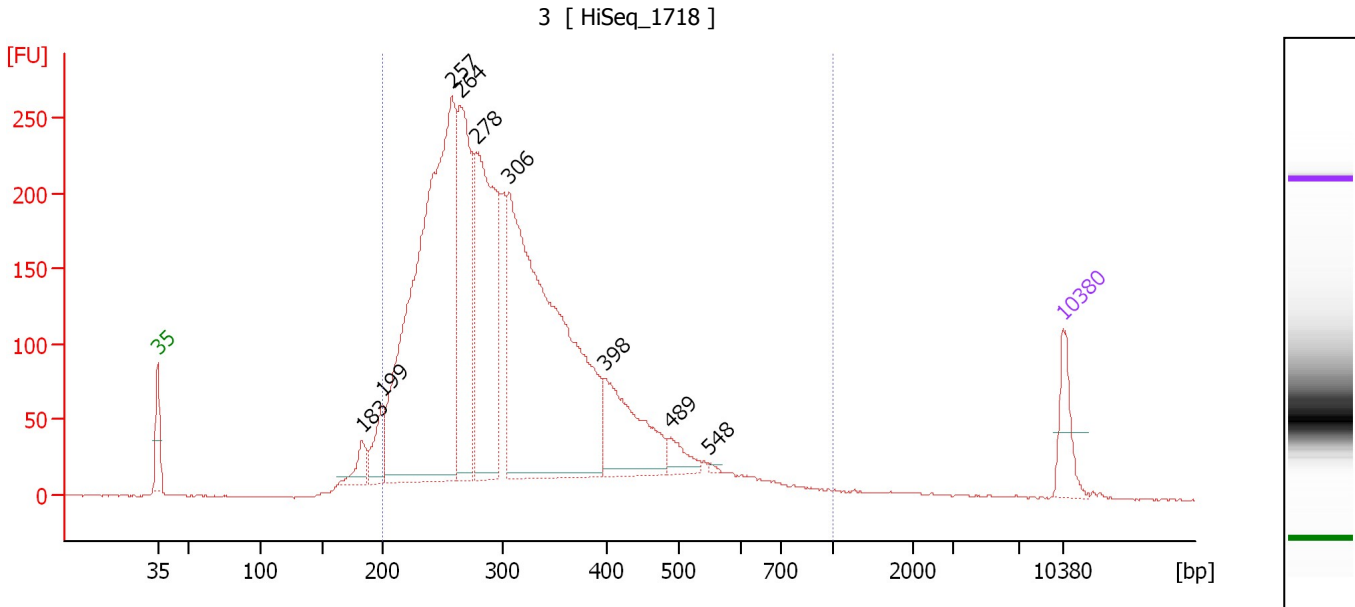
... Region table for sample 2 : 2

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	324	7,231.12	7,450.5	37,478.0	 94	34.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
 Modified: 3/7/2019 5:02:36 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : 3**

Number of peaks found: 9                      Corr. Area 1: 4,870.3  
 Noise: 0.5

**Peak table for sample 3 : 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	183	65.52	543.6		58.81
3	199	114.96	873.1		60.39
4	257	1,718.68	10,117.5		65.75
5	264	572.62	3,284.0		66.38
6	278	710.35	3,874.1		67.64
7	306	1,500.00	7,429.9		70.16
8	398	294.80	1,121.9		77.50
9	489	52.82	163.8		82.68
10	548	5.19	14.4		85.61
11	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 3 : 3**

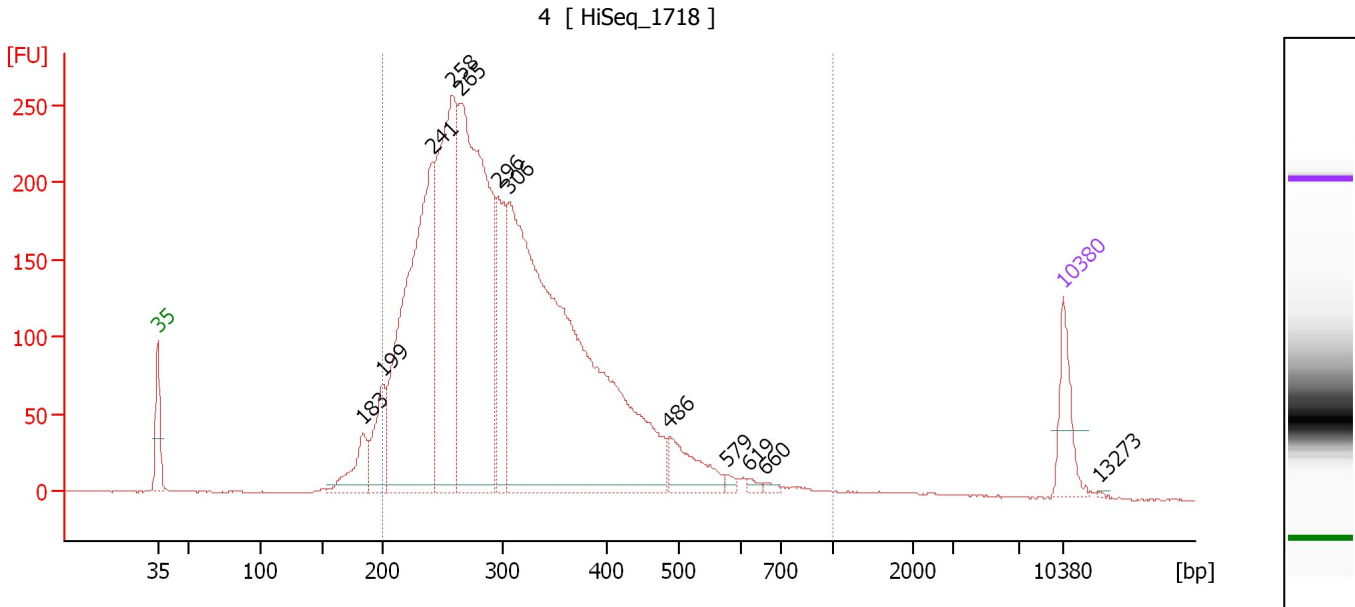
From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	321	5,699.34	4,870.3	29,459.6	95	31.0



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
 Modified: 3/7/2019 5:02:36 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : 4**

Number of peaks found: 12                      Corr. Area 1: 4,807.2  
 Noise: 0.4

**Peak table for sample 4 : 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	183	105.23	870.2		58.87
3	199	143.50	1,090.0		60.39
4	241	922.45	5,806.8		64.20
5	258	716.54	4,206.7		65.81
6	265	1,081.10	6,180.5		66.46
7	296	226.75	1,161.3		69.31
8	306	1,831.07	9,062.7		70.18
9	486	110.78	345.0		82.55
10	579	9.94	26.0		87.06
11	619	9.47	23.2		88.67
12	660	7.10	16.3		89.91
13	10,380	75.00	10.9	Upper Marker	113.00
14	13,273	0.00	0.0		115.85

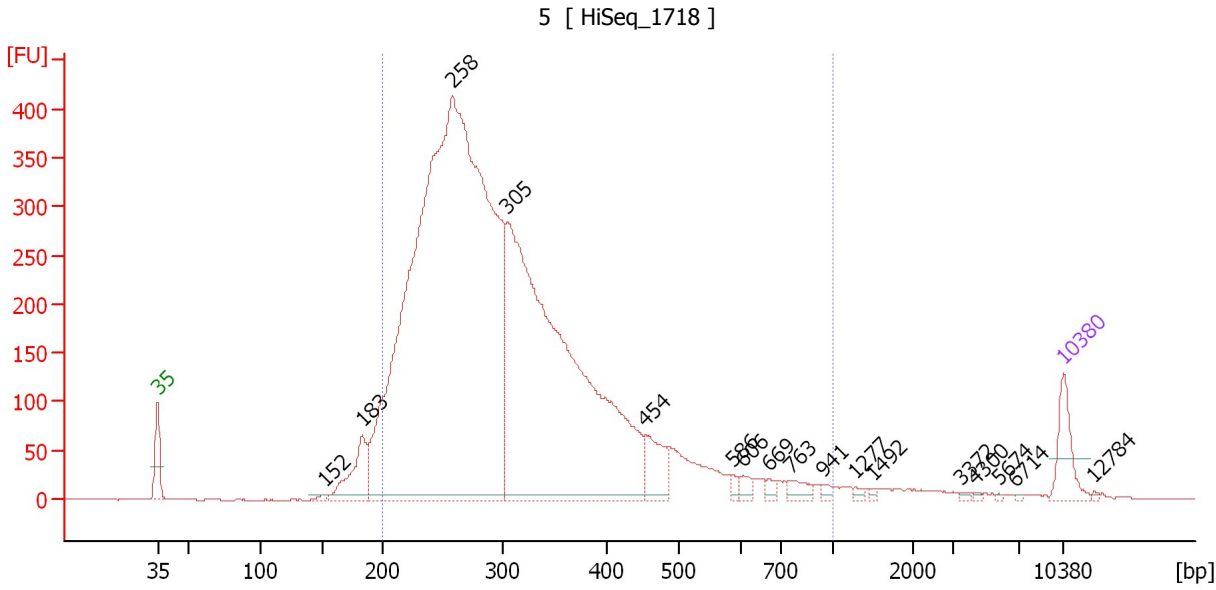
**Region table for sample 4 : 4**

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	316	4,886.87	4,807.2	25,464.7	95	29.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
 Modified: 3/7/2019 5:02:36 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : 5**

Number of peaks found: 17                      Corr. Area 1: 7,256.3  
 Noise: 0.4

**Peak table for sample 5 : 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	152	8.42	84.0		55.94
3	183	168.06	1,392.8		58.83
4	258	4,276.28	25,122.4		65.80
5	305	2,190.26	10,894.0		70.06
6	454	107.94	360.1		80.73
7	586	13.09	33.9		87.41
8	606	21.96	54.9		88.26
9	669	14.91	33.8		90.20
10	763	29.21	58.0		92.00
11	941	11.01	17.7		94.42
12	1,277	7.50	8.9		96.93
13	1,492	4.77	4.8		98.26
14	3,372	4.35	2.0		105.04
15	4,300	2.97	1.0		106.22
16	5,674	2.10	0.6		107.98
17	6,714	1.90	0.4		109.30
18	10,380	75.00	10.9	Upper Marker	113.00
19	12,784	0.00	0.0		115.37

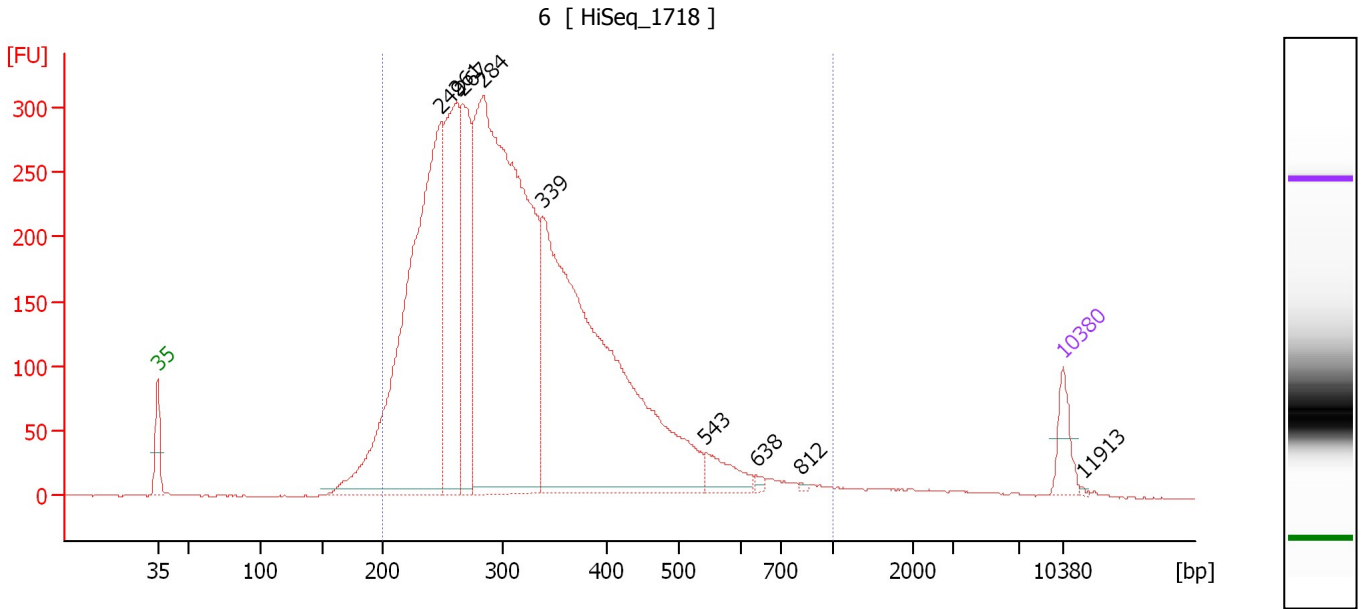
**Region table for sample 5 : 5**

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	319	6,531.78	7,256.3	34,212.9	94	33.6

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
 Modified: 3/7/2019 5:02:36 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : 6**

Number of peaks found: 9                      Corr. Area 1: 6,428.5  
 Noise: 0.4

**Peak table for sample 6 : 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	249	2,049.00	12,487.4		64.94
3	261	819.09	4,750.4		66.11
4	267	569.62	3,238.3		66.59
5	284	2,708.03	14,453.0		68.20
6	339	2,277.30	10,183.6		72.78
7	543	103.56	289.0		85.36
8	638	11.00	26.1		89.26
9	812	4.87	9.1		92.67
10	10,380	75.00	10.9	Upper Marker	113.00
11	11,913	0.00	0.0		114.51

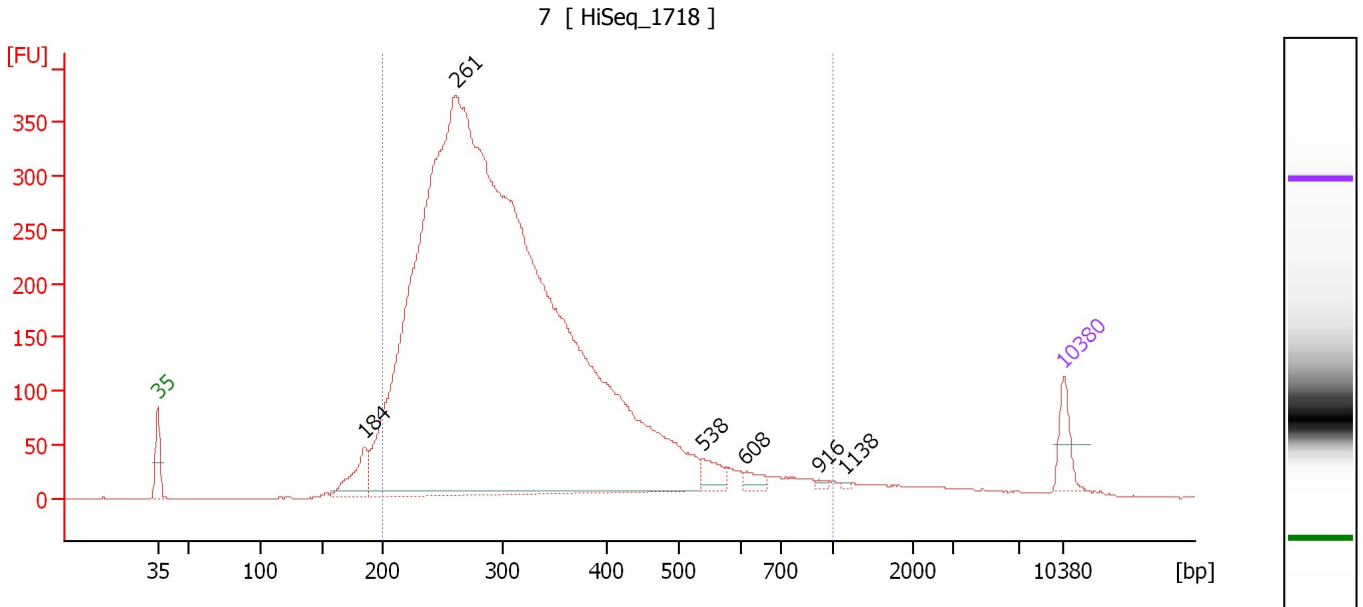
**Region table for sample 6 : 6**

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	326	8,286.41	6,428.5	42,229.8	96	30.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
 Modified: 3/7/2019 5:02:36 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : 7**

Number of peaks found: 6                      Corr. Area 1: 6,854.1  
 Noise: 0.4

**Peak table for sample 7 : 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	184	170.80	1,403.4		58.98
3	261	9,411.07	54,668.3		66.07
4	538	68.19	192.0		85.14
5	608	32.72	81.5		88.34
6	916	8.66	14.3		94.08
7	1,138	4.02	5.4		96.07
8	10,380	75.00	10.9	Upper Marker	113.00

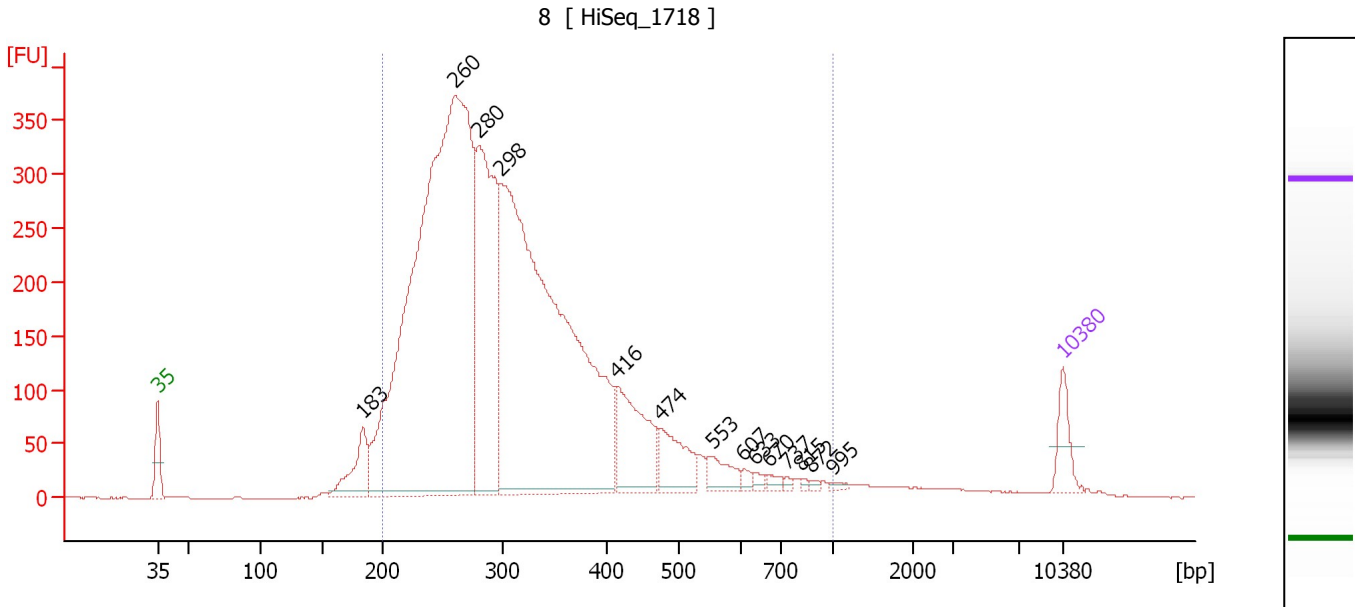
**Region table for sample 7 : 7**

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	326	9,045.55	6,854.1	46,586.6	93	34.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
 Modified: 3/7/2019 5:02:36 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : 8**

Number of peaks found: 14                      Corr. Area 1: 6,919.7  
 Noise: 0.4

**Peak table for sample 8 : 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	183	201.12	1,663.8		58.86
3	260	3,633.88	21,187.6		65.98
4	280	1,041.33	5,642.3		67.81
5	298	2,802.31	14,260.0		69.49
6	416	337.94	1,232.3		78.53
7	474	179.26	573.2		81.84
8	553	82.51	226.0		85.85
9	607	17.94	44.8		88.29
10	633	16.53	39.5		89.11
11	670	18.76	42.4		90.23
12	737	8.58	17.7		91.65
13	815	7.39	13.7		92.72
14	872	7.47	13.0		93.48
15	995	8.54	13.0		95.16
16	10,380	75.00	10.9	Upper Marker	113.00

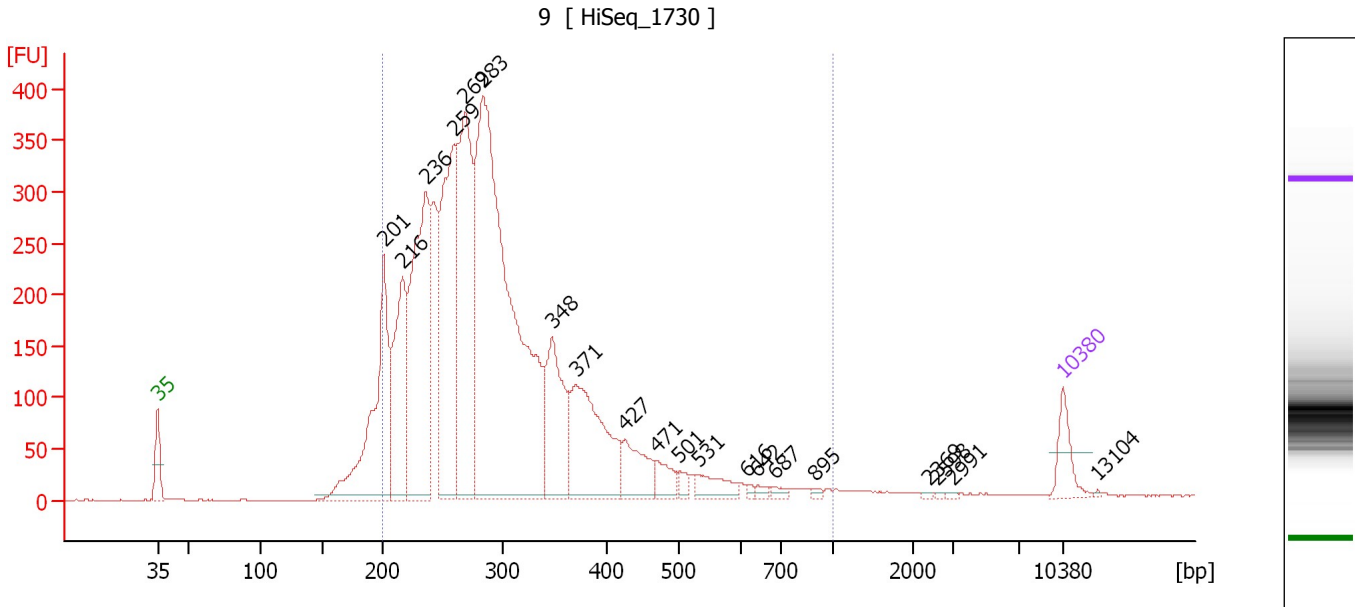
**Region table for sample 8 : 8**

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	8,162.80	6,919.7	41,951.8	93	33.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
 Modified: 3/7/2019 5:02:36 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : 9**

Number of peaks found: 20                      Corr. Area 1: 6,007.9  
 Noise: 0.4

**Peak table for sample 9 : 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	201	704.36	5,317.5		60.50
3	216	497.00	3,481.2		61.95
4	236	921.68	5,921.6		63.75
5	259	861.34	5,034.1		65.92
6	269	832.67	4,694.9		66.80
7	283	2,228.17	11,920.0		68.14
8	348	380.17	1,655.6		73.51
9	371	488.20	1,992.3		75.37
10	427	154.68	548.8		79.19
11	471	69.23	222.8		81.67
12	501	26.32	79.6		83.37
13	531	75.26	214.6		84.81
14	616	9.63	23.7		88.58
15	642	13.17	31.1		89.36
16	687	15.98	35.2		90.75
17	895	8.75	14.8		93.80
18	2,369	3.85	2.5		102.57
19	2,598	2.76	1.6		103.29
20	2,991	3.81	1.9		104.53
21	10,380	75.00	10.9	Upper Marker	113.00
22	13,104	0.00	0.0		115.68

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

Created: 3/7/2019 4:21:18 PM  
Modified: 3/7/2019 5:02:36 PM

**Electropherogram Summary Continued ...**

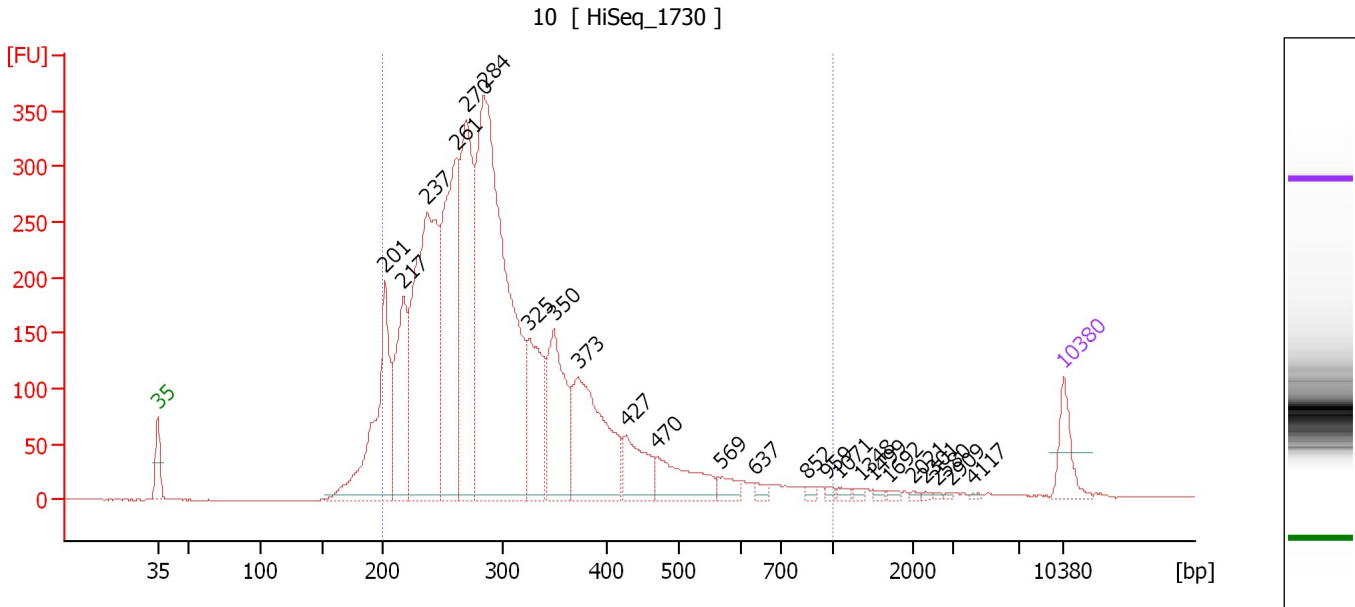
... Region table for sample 9 : 9

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	304	7,119.87	6,007.9	38,394.3	 93	31.2

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : 10**

Number of peaks found: 24                      Corr. Area 1: 5,511.6  
 Noise: 0.5

**Peak table for sample 10 : 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	201	555.37	4,180.9		60.55
3	217	390.20	2,724.9		62.01
4	237	1,064.18	6,798.8		63.88
5	261	763.88	4,439.3		66.06
6	270	674.42	3,789.0		66.89
7	284	1,699.06	9,055.8		68.24
8	325	295.04	1,373.7		71.72
9	350	352.93	1,529.8		73.64
10	373	457.30	1,857.4		75.51
11	427	150.62	534.2		79.19
12	470	161.95	521.8		81.64
13	569	35.03	93.2		86.62
14	637	17.14	40.8		89.22
15	852	11.26	20.0		93.22
16	959	8.91	14.1		94.67
17	1,071	9.54	13.5		95.66
18	1,348	7.24	8.1		97.37
19	1,499	6.48	6.5		98.30
20	1,692	6.93	6.2		99.50
21	2,021	4.23	3.2		101.47
22	2,301	4.09	2.7		102.35
23	2,580	3.56	2.1		103.24
24	2,909	3.37	1.8		104.28
25	4,117	3.38	1.2		105.99



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

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**Electropherogram Summary Continued ...****... Peak table for sample 10 : 10**

Peak	Size [bp]	Conc. [pg/ $\mu$ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	10,380	75.00	10.9	Upper Marker	113.00

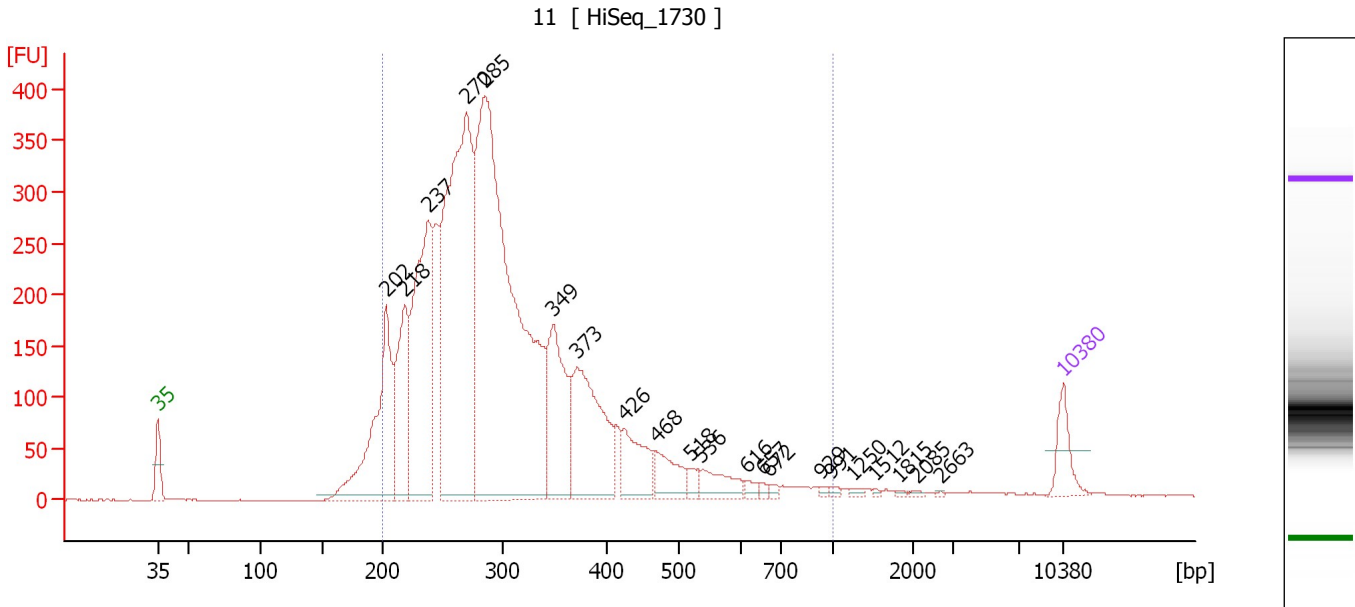
**Region table for sample 10 : 10**

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	311	6,305.71	5,511.6	33,528.2	93	32.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : 11**

Number of peaks found: 21                      Corr. Area 1: 6,042.2  
 Noise: 0.5

**Peak table for sample 11 : 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	202	611.24	4,579.1		60.64
3	218	393.53	2,734.6		62.11
4	237	783.98	5,007.6		63.88
5	270	1,553.14	8,718.7		66.91
6	285	2,236.33	11,907.1		68.26
7	349	389.79	1,692.5		73.59
8	373	476.17	1,936.6		75.47
9	426	170.89	607.9		79.12
10	468	112.89	365.3		81.52
11	518	29.65	86.7		84.19
12	536	87.38	247.1		85.02
13	616	16.15	39.7		88.57
14	657	9.87	22.8		89.82
15	672	10.81	24.4		90.29
16	929	7.21	11.8		94.26
17	991	7.89	12.1		95.10
18	1,250	8.64	10.5		96.77
19	1,512	4.19	4.2		98.38
20	1,815	4.06	3.4		100.26
21	2,085	4.64	3.4		101.67
22	2,663	2.54	1.4		103.50
23	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

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**Electropherogram Summary Continued ...**

... Region table for sample **11** : 11

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	313	6,832.00	6,042.2	36,081.4	 94	31.9

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-03-07\2019-03-07\_006.xad

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**Gel Image**

