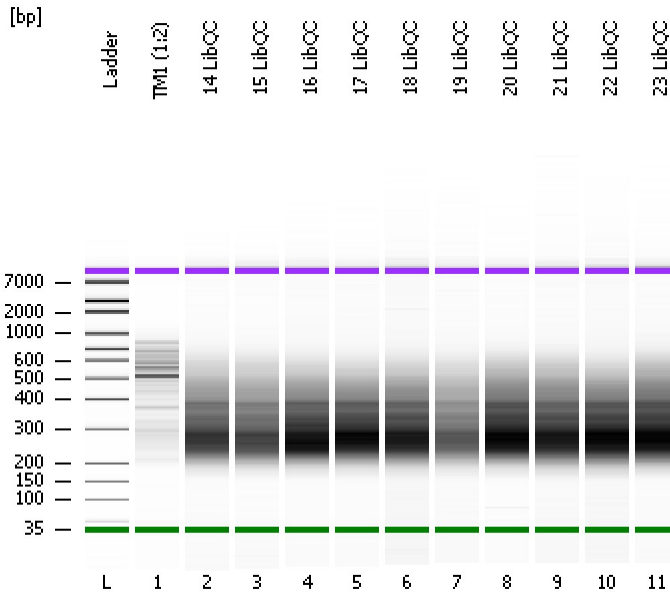


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
Modified: 10/18/2019 12:01:45 PM

**Electrophoresis File Run Summary**



Instrument Information:

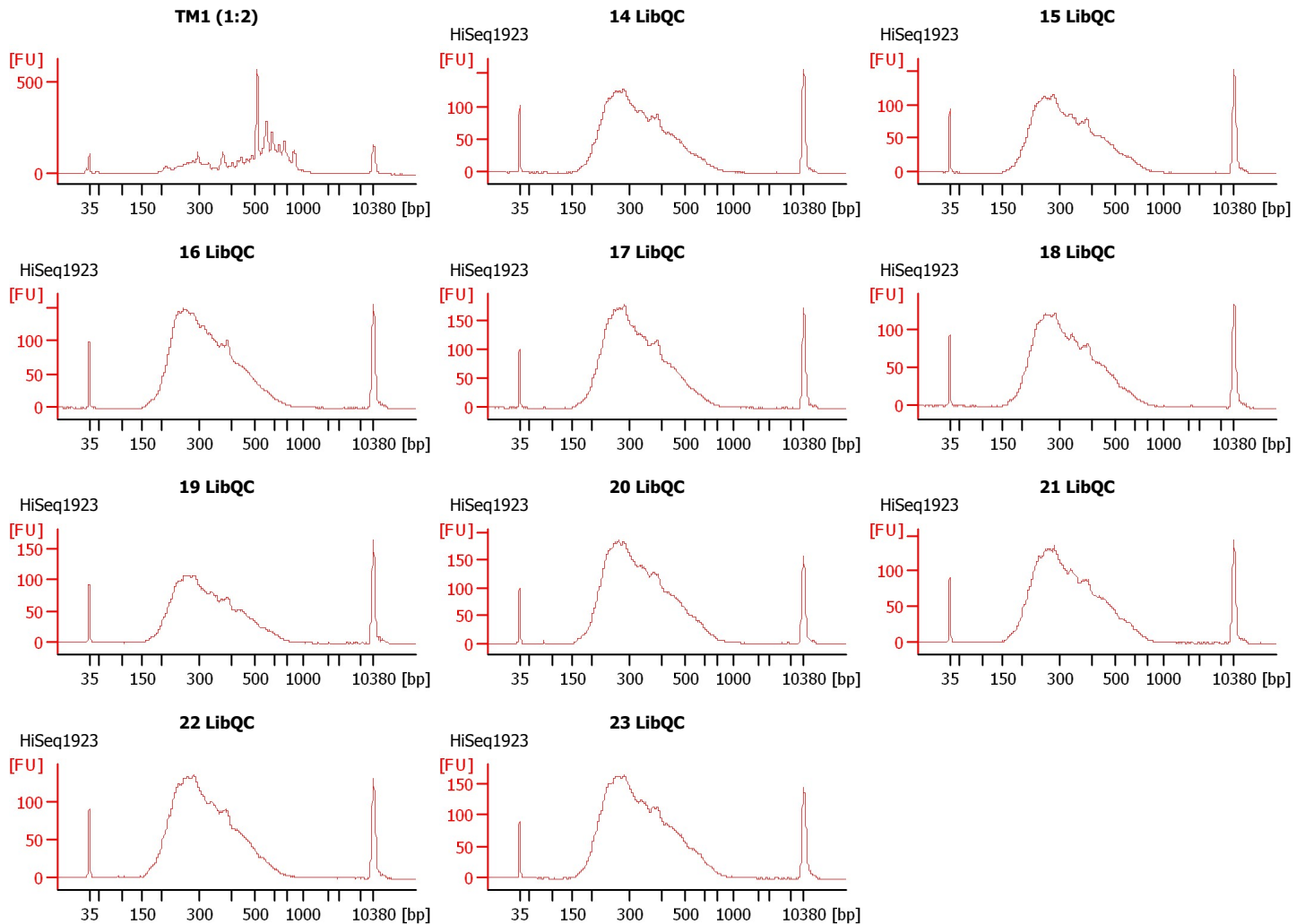
Instrument Name: DE34903152      Firmware: C.01.069  
Serial#: DE34903152      Type: G2938C

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\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
TM1 (1:2)		<input type="checkbox"/>	✓			
14 LibQC	HiSeq1923	<input type="checkbox"/>	✓			
15 LibQC	HiSeq1923	<input type="checkbox"/>	✓			
16 LibQC	HiSeq1923	<input type="checkbox"/>	✓			
17 LibQC	HiSeq1923	<input type="checkbox"/>	✓			
18 LibQC	HiSeq1923	<input type="checkbox"/>	✓			
19 LibQC	HiSeq1923	<input type="checkbox"/>	✓			
20 LibQC	HiSeq1923	<input type="checkbox"/>	✓			
21 LibQC	HiSeq1923	<input type="checkbox"/>	✓			
22 LibQC	HiSeq1923	<input type="checkbox"/>	✓			
23 LibQC	HiSeq1923	<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\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
Modified: 10/18/2019 12:01:45 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

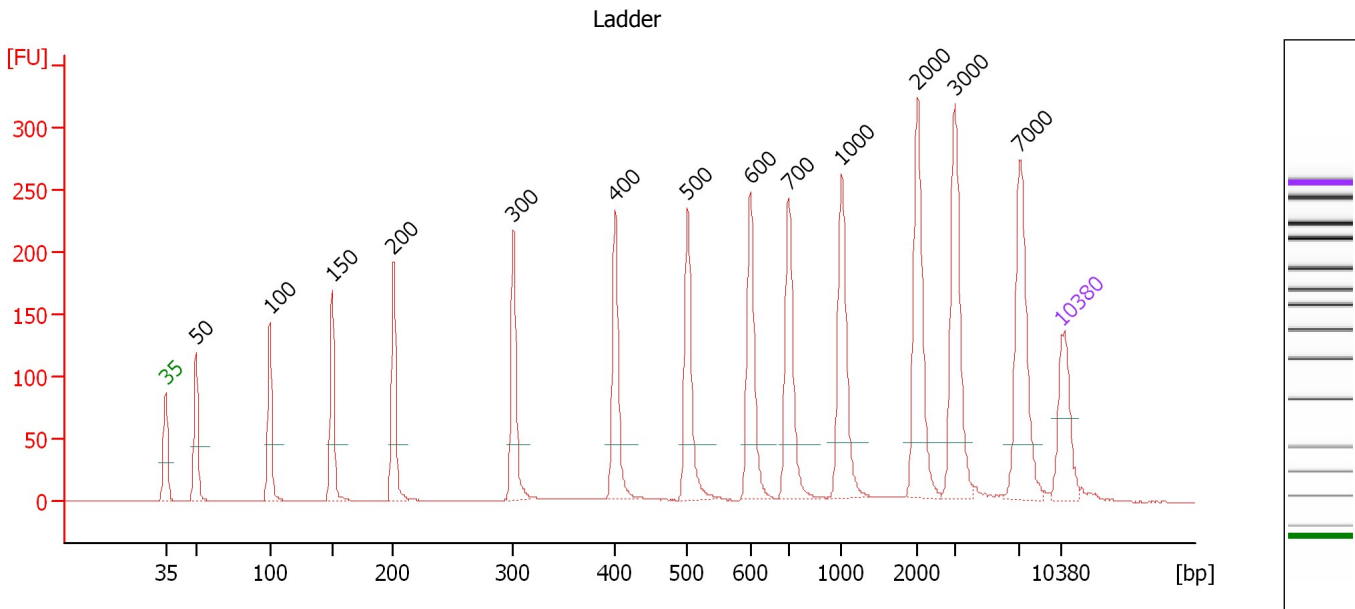
**Ladder**

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\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.3

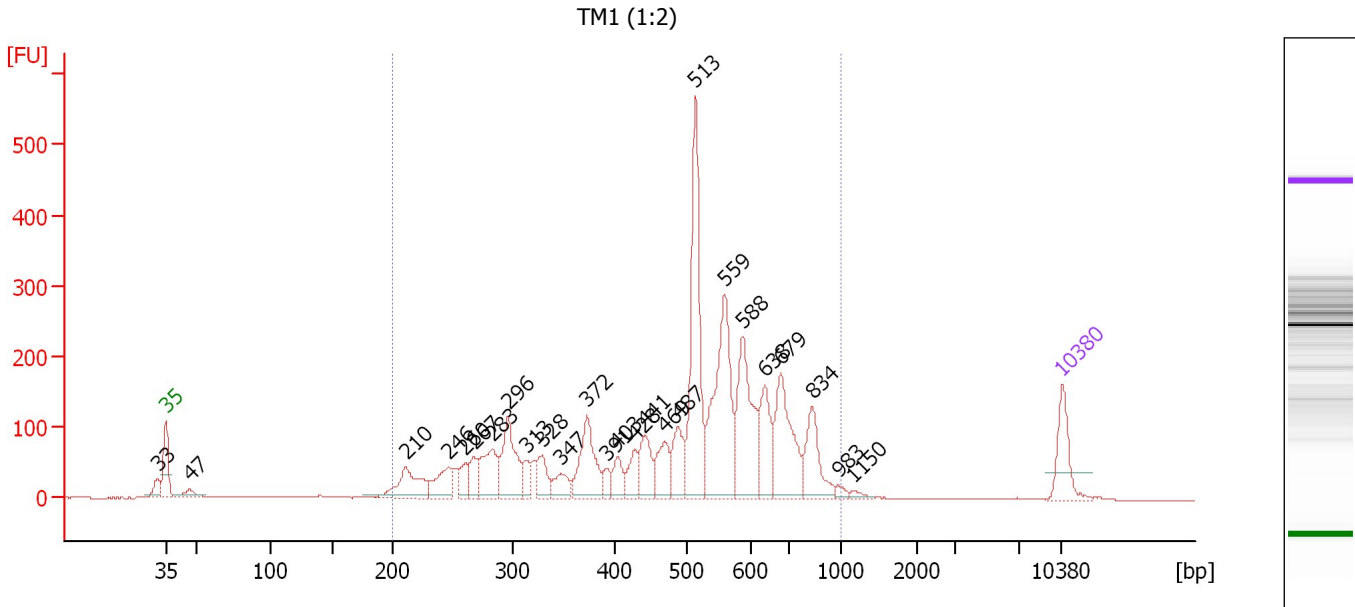
**Peak table for Ladder**

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	45.35
3	100	150.00	2,272.7	Ladder Peak	51.14
4	150	150.00	1,515.2	Ladder Peak	55.99
5	200	150.00	1,136.4	Ladder Peak	60.78
6	300	150.00	757.6	Ladder Peak	70.15
7	400	150.00	568.2	Ladder Peak	78.11
8	500	150.00	454.5	Ladder Peak	83.77
9	600	150.00	378.8	Ladder Peak	88.66
10	700	150.00	324.7	Ladder Peak	91.64
11	1,000	150.00	227.3	Ladder Peak	95.76
12	2,000	150.00	113.6	Ladder Peak	101.73
13	3,000	150.00	75.8	Ladder Peak	104.63
14	7,000	150.00	32.5	Ladder Peak	109.70
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : TM1 (1:2)**

Number of peaks found: 26                      Corr. Area 1: 3,838.9  
 Noise: 0.3

**Peak table for sample 1 : TM1 (1:2)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	33	0.00	0.0		42.36
2	35	125.00	5,411.3	Lower Marker	43.00
3	47	32.89	1,063.4		44.86
4	210	129.22	930.9		61.75
5	246	86.02	530.5		65.06
6	260	49.17	286.3		66.42
7	267	51.62	292.9		67.06
8	283	121.62	652.3		68.51
9	296	163.89	840.0		69.74
10	313	38.17	184.7		71.19
11	328	59.28	273.4		72.42
12	347	51.17	223.6		73.87
13	372	153.16	623.2		75.91
14	391	26.13	101.2		77.41
15	403	46.76	175.9		78.27
16	428	53.14	188.2		79.68
17	441	83.59	287.0		80.45
18	469	79.39	256.7		81.99
19	487	85.35	265.5		83.04
20	513	377.35	1,114.7		84.40
21	559	335.18	907.9		86.67
22	588	227.48	586.0		88.08
23	638	120.14	285.1		89.80
24	679	199.08	443.9		91.03
25	834	115.22	209.4		93.48

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 PM

### Electropherogram Summary Continued ...

#### ... Peak table for sample 1 : TM1 (1:2)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	983	10.58	16.3		95.52
27	1,150	9.64	12.7		96.66
28	10,380	75.00	10.9	Upper Marker	113.00

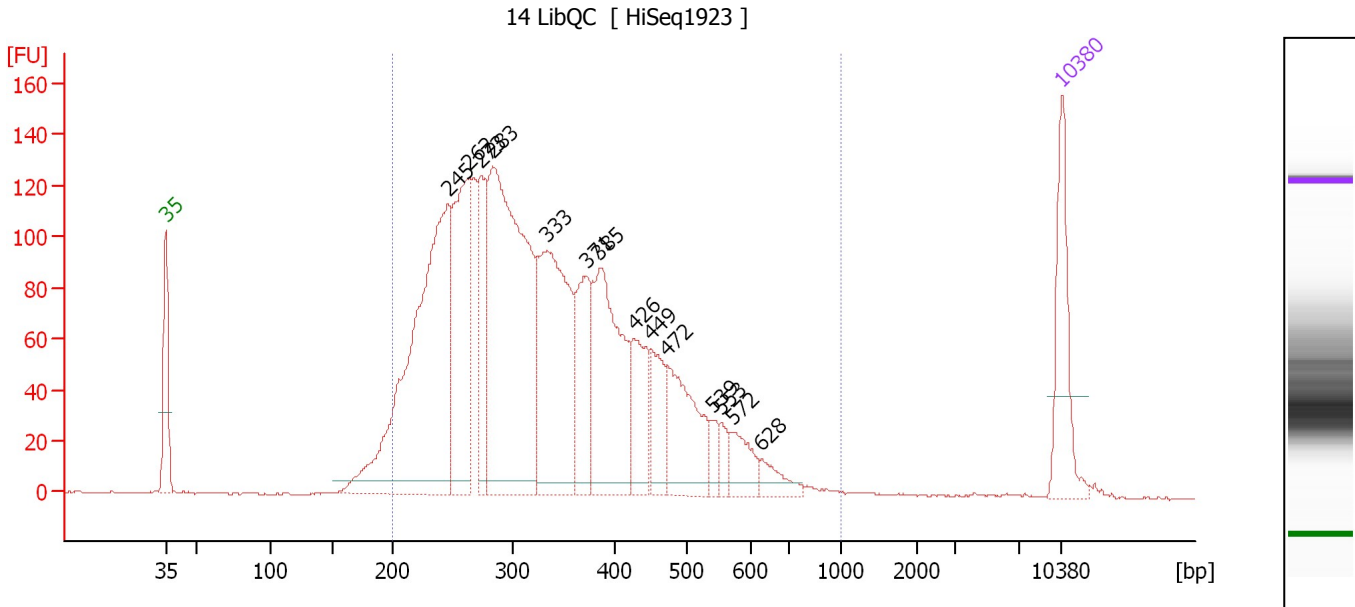
#### Region table for sample 1 : TM1 (1:2)

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	503	2,722.88	3,838.9	9,847.4	95	31.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : 14 LibQC**

Number of peaks found: 14                      Corr. Area 1: 3,114.4  
 Noise: 0.2

**Peak table for sample 2 : 14 LibQC**

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	245	555.37	3,427.8		65.04
3	262	235.11	1,359.5		66.59
4	273	122.65	680.2		67.64
5	283	523.22	2,802.0		68.55
6	333	288.11	1,310.5		72.79
7	371	103.91	423.9		75.84
8	385	225.19	887.2		76.88
9	426	80.32	285.8		79.57
10	449	60.86	205.3		80.89
11	472	121.08	388.9		82.17
12	539	16.56	46.5		85.67
13	553	17.85	48.9		86.36
14	572	38.02	100.8		87.27
15	628	22.47	54.2		89.50
16	10,380	75.00	10.9	Upper Marker	113.00

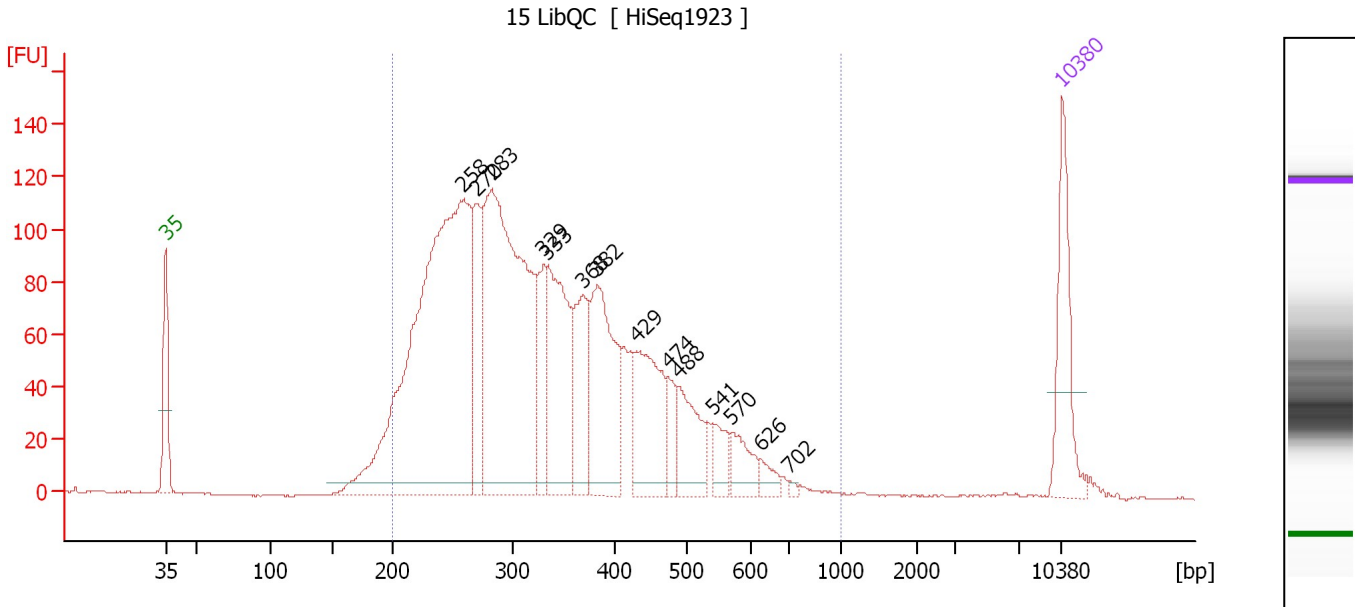
**Region table for sample 2 : 14 LibQC**

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	2,451.92	3,114.4	12,081.9	96	31.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : 15 LibQC**

Number of peaks found: 14                      Corr. Area 1: 2,818.7  
 Noise: 0.3

**Peak table for sample 3 : 15 LibQC**

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	258	782.02	4,587.8		66.24
3	270	108.07	607.2		67.31
4	283	541.59	2,903.9		68.52
5	329	72.71	334.7		72.47
6	333	184.79	841.8		72.75
7	368	96.87	399.2		75.54
8	382	173.57	688.0		76.70
9	429	124.42	439.2		79.77
10	474	27.77	88.9		82.28
11	488	69.05	214.2		83.11
12	541	29.14	81.6		85.76
13	570	34.11	90.6		87.20
14	626	14.31	34.6		89.43
15	702	3.08	6.6		91.67
16	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 3 : 15 LibQC**

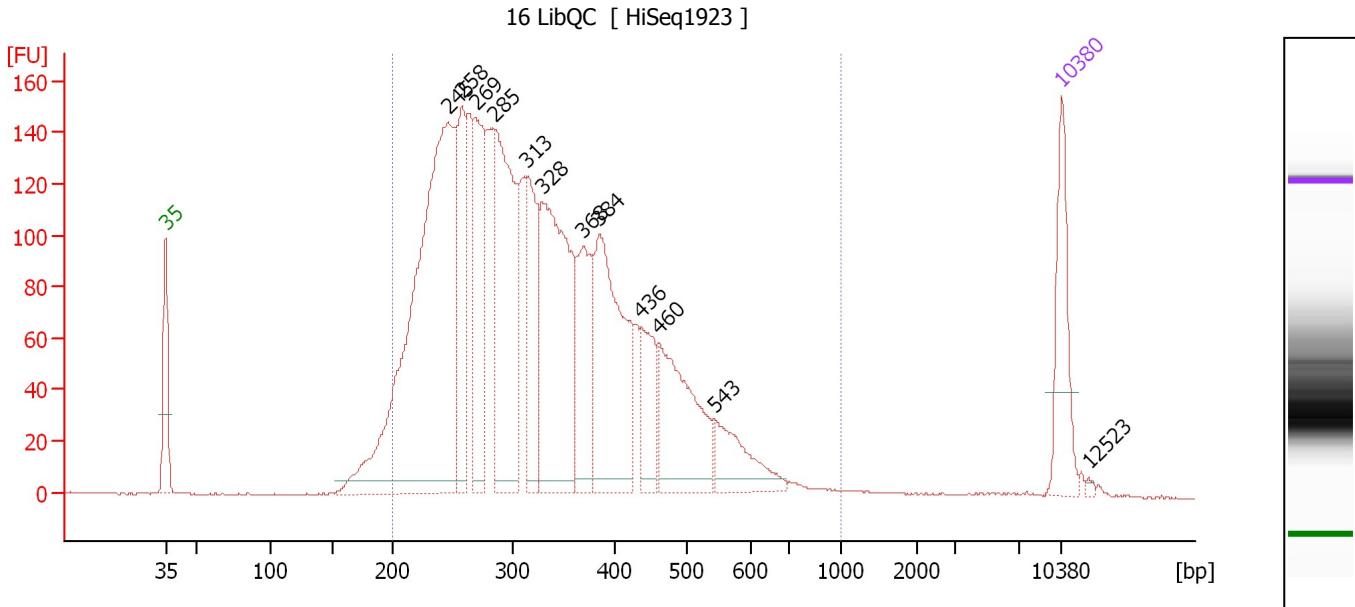
From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	342	2,275.25	2,818.7	11,268.5	96	31.1



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

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 Modified: 10/18/2019 12:01:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : 16 LibQC**

Number of peaks found: 12                      Corr. Area 1: 3,591.9  
 Noise: 0.3

**Peak table for sample 4 : 16 LibQC**

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	245	801.21	4,947.4		65.03
3	258	155.60	915.3		66.17
4	269	184.49	1,040.1		67.22
5	285	312.56	1,661.7		68.74
6	313	142.65	689.6		71.22
7	328	344.50	1,589.6		72.41
8	368	132.18	543.6		75.60
9	384	263.61	1,040.3		76.83
10	436	72.95	253.3		80.17
11	460	172.01	566.8		81.50
12	543	74.18	206.9		85.88
13	10,380	75.00	10.9	Upper Marker	113.00
14	12,523	0.00	0.0		115.09

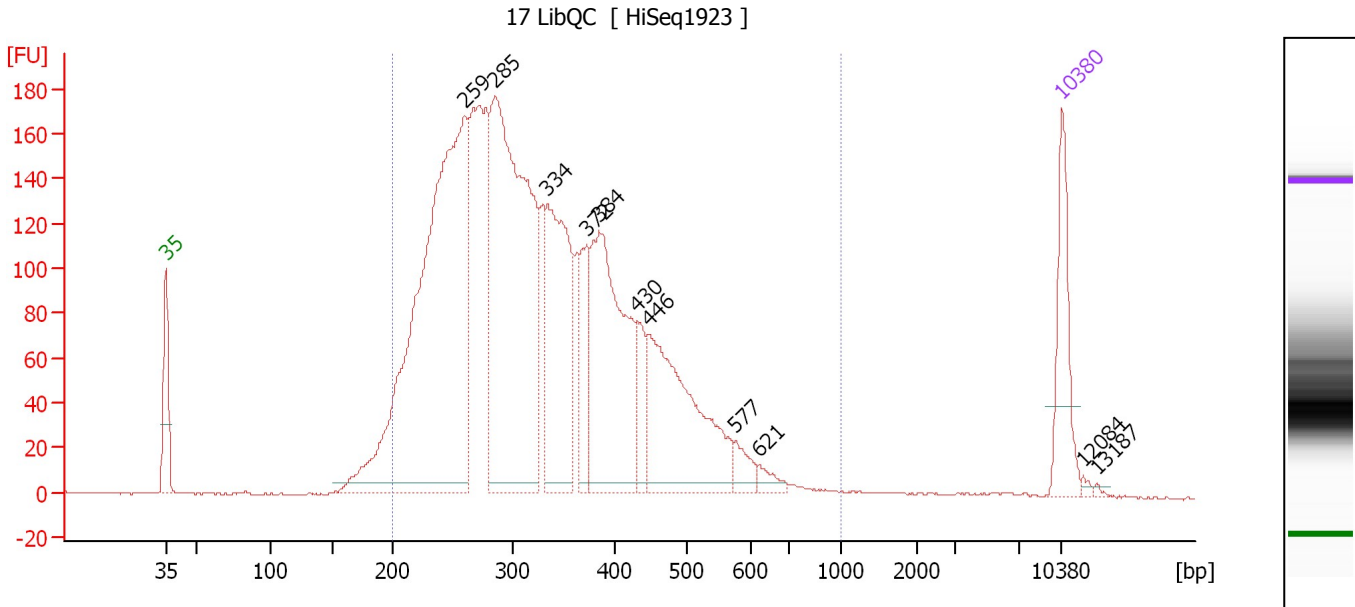
**Region table for sample 4 : 16 LibQC**

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	337	2,985.52	3,591.9	14,901.4	96	30.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : 17 LibQC**

Number of peaks found: 11                      Corr. Area 1: 4,043.6  
 Noise: 0.3

**Peak table for sample 5 : 17 LibQC**

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	926.18	5,416.4		66.32
3	285	702.89	3,740.6		68.72
4	334	276.96	1,257.6		72.83
5	372	86.66	353.2		75.87
6	384	318.96	1,258.1		76.85
7	430	51.18	180.1		79.84
8	446	260.71	885.5		80.72
9	577	24.29	63.8		87.53
10	621	13.98	34.1		89.29
11	10,380	75.00	10.9	Upper Marker	113.00
12	12,084	0.00	0.0		114.67
13	13,187	0.00	0.0		115.74

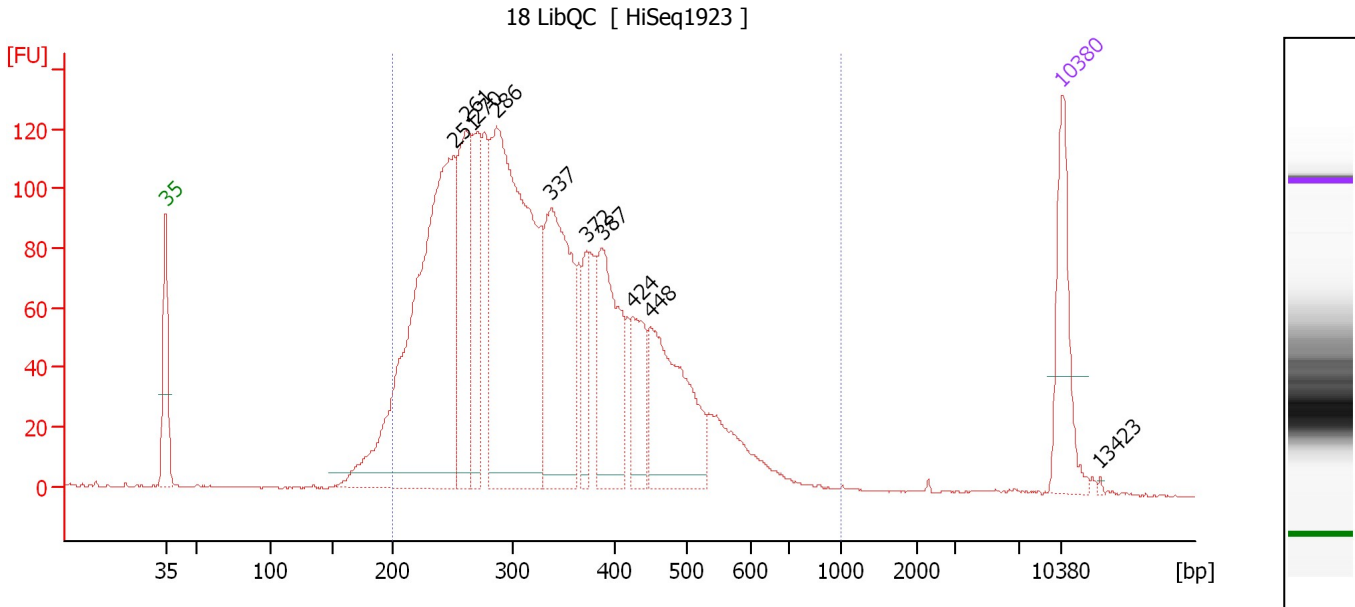
**Region table for sample 5 : 17 LibQC**

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	337	3,059.44	4,043.6	15,190.7	96	29.6

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : 18 LibQC**

Number of peaks found: 10                      Corr. Area 1: 2,900.6  
 Noise: 0.3

**Peak table for sample 6 : 18 LibQC**

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	251	631.53	3,809.3		65.58
3	261	192.12	1,115.8		66.48
4	270	124.78	700.2		67.34
5	286	576.23	3,051.0		68.85
6	337	267.04	1,201.1		73.09
7	372	60.05	244.8		75.86
8	387	164.84	645.6		77.07
9	424	76.94	274.8		79.49
10	448	182.52	616.8		80.85
11	10,380	75.00	10.9	Upper Marker	113.00
12	13,423	0.00	0.0		115.97

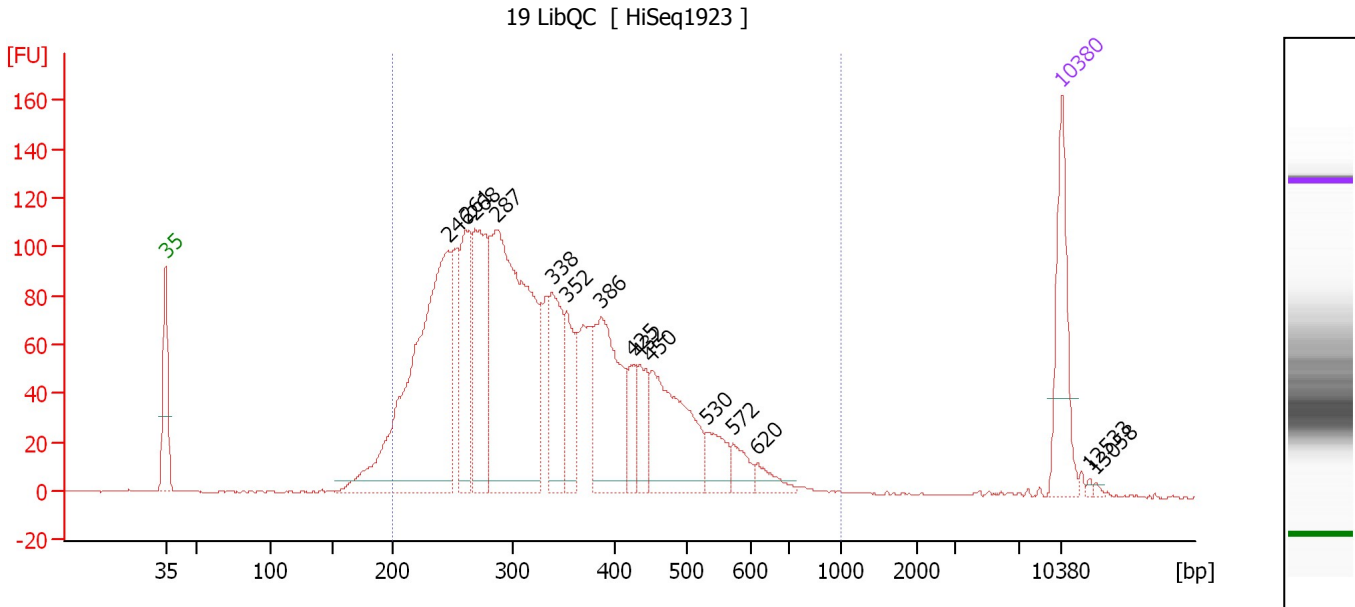
**Region table for sample 6 : 18 LibQC**

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	339	2,516.54	2,900.6	12,478.8	95	30.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : 19 LibQC**

Number of peaks found: 15                      Corr. Area 1: 2,586.7  
 Noise: 0.3

**Peak table for sample 7 : 19 LibQC**

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	246	501.55	3,092.3		65.07
3	261	131.29	763.5		66.45
4	268	187.99	1,064.2		67.12
5	287	472.28	2,494.7		68.92
6	338	103.84	465.5		73.18
7	352	73.43	315.9		74.30
8	386	171.61	673.0		77.02
9	425	35.57	126.8		79.54
10	432	44.66	156.8		79.90
11	450	155.05	522.4		80.93
12	530	37.23	106.4		85.24
13	572	23.94	63.4		87.29
14	620	17.23	42.1		89.24
15	10,380	75.00	10.9	Upper Marker	113.00
16	12,533	0.00	0.0		115.10
17	13,058	0.00	0.0		115.62

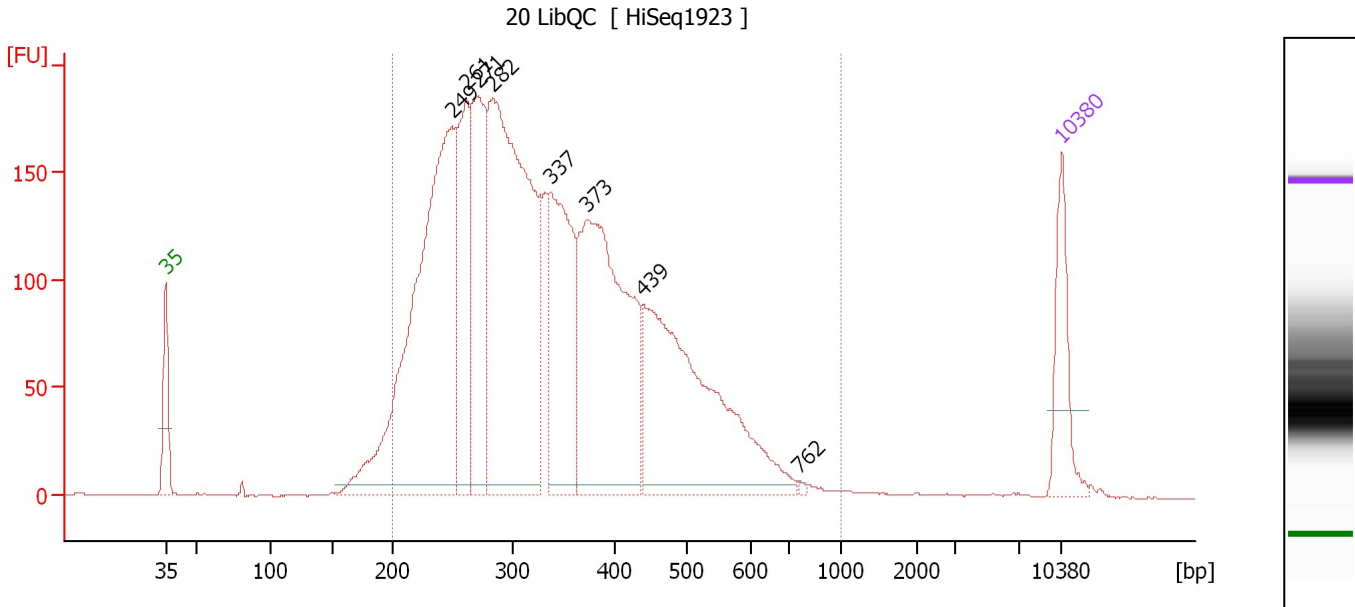
**Region table for sample 7 : 19 LibQC**

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	342	2,145.22	2,586.7	10,596.3	95	30.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
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**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : 20 LibQC**

Number of peaks found: 8                      Corr. Area 1: 4,515.0  
 Noise: 0.2

**Peak table for sample 8 : 20 LibQC**

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	820.81	4,993.1		65.38
3	261	250.96	1,455.2		66.52
4	271	275.63	1,542.5		67.41
5	282	789.67	4,236.6		68.50
6	337	297.30	1,337.4		73.08
7	373	524.73	2,132.8		75.94
8	439	478.52	1,651.7		80.32
9	762	2.51	5.0		92.49
10	10,380	75.00	10.9	Upper Marker	113.00

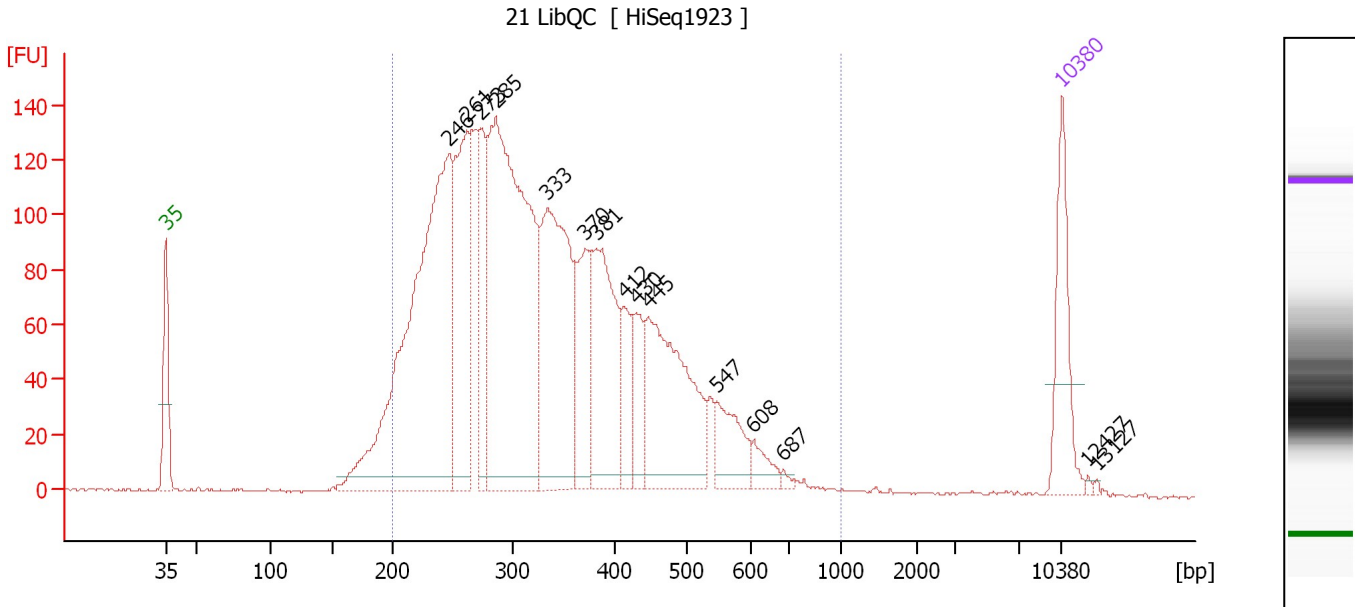
**Region table for sample 8 : 20 LibQC**

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	346	3,426.33	4,515.0	16,771.0	96	31.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : 21 LibQC**

Number of peaks found: 15                      Corr. Area 1: 3,232.7  
 Noise: 0.2

**Peak table for sample 9 : 21 LibQC**

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	246	679.03	4,179.7		65.11
3	261	237.65	1,377.9		66.53
4	273	119.63	665.0		67.58
5	285	610.96	3,242.6		68.79
6	333	303.28	1,379.3		72.79
7	370	114.68	469.4		75.74
8	381	190.95	758.6		76.63
9	412	57.51	211.5		78.79
10	430	58.35	205.8		79.79
11	445	223.83	761.3		80.68
12	547	59.35	164.5		86.05
13	608	20.68	51.5		88.89
14	687	3.85	8.5		91.26
15	10,380	75.00	10.9	Upper Marker	113.00
16	12,427	0.00	0.0		115.00
17	13,127	0.00	0.0		115.68

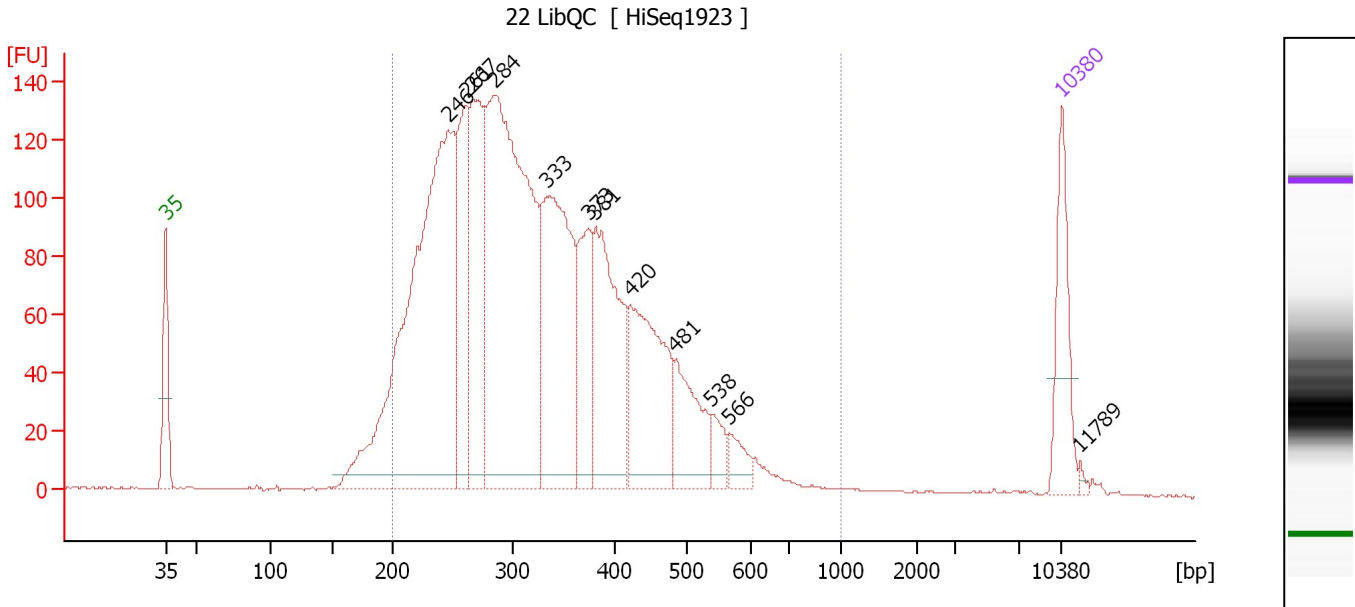
**Region table for sample 9 : 21 LibQC**

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	343	2,762.55	3,232.7	13,623.3	95	31.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : 22 LibQC**

Number of peaks found: 12                      Corr. Area 1: 3,167.3  
 Noise: 0.3

**Peak table for sample 10 : 22 LibQC**

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	246	794.01	4,892.8		65.08
3	261	203.18	1,178.9		66.51
4	267	227.87	1,294.1		67.04
5	284	714.91	3,809.8		68.68
6	333	342.65	1,560.2		72.76
7	373	127.36	516.9		75.99
8	381	230.19	916.3		76.57
9	420	208.37	750.8		79.27
10	481	98.02	308.6		82.71
11	538	28.31	79.7		85.62
12	566	25.20	67.4		87.00
13	10,380	75.00	10.9	Upper Marker	113.00
14	11,789	0.00	0.0		114.38

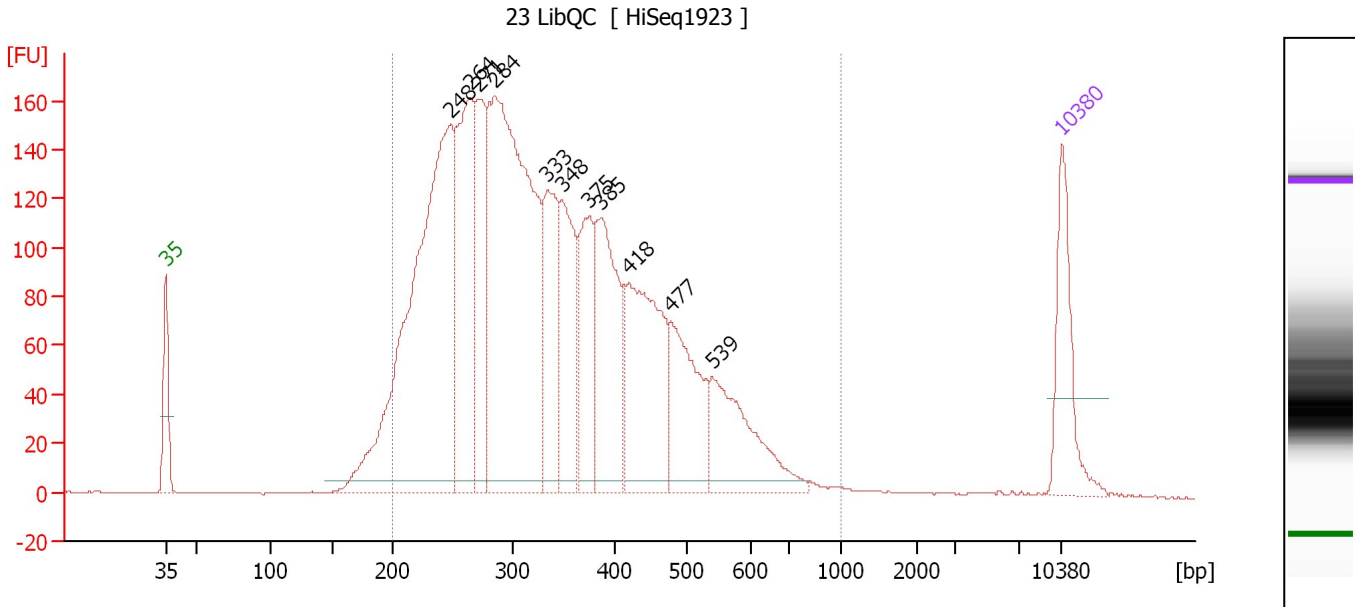
**Region table for sample 10 : 22 LibQC**

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	335	2,963.38	3,167.3	14,808.8	95	29.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : 23 LibQC**

Number of peaks found: 11                      Corr. Area 1: 4,017.3  
 Noise: 0.2

**Peak table for sample 11 : 23 LibQC**

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	248	682.88	4,177.0		65.25
3	264	250.84	1,441.7		66.74
4	271	159.17	889.9		67.43
5	284	654.47	3,490.7		68.66
6	333	159.26	723.8		72.81
7	348	143.00	622.4		73.98
8	375	121.55	491.3		76.11
9	385	193.22	760.7		76.91
10	418	220.35	798.3		79.14
11	477	139.16	442.4		82.44
12	539	131.36	369.0		85.69
13	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 11 : 23 LibQC**

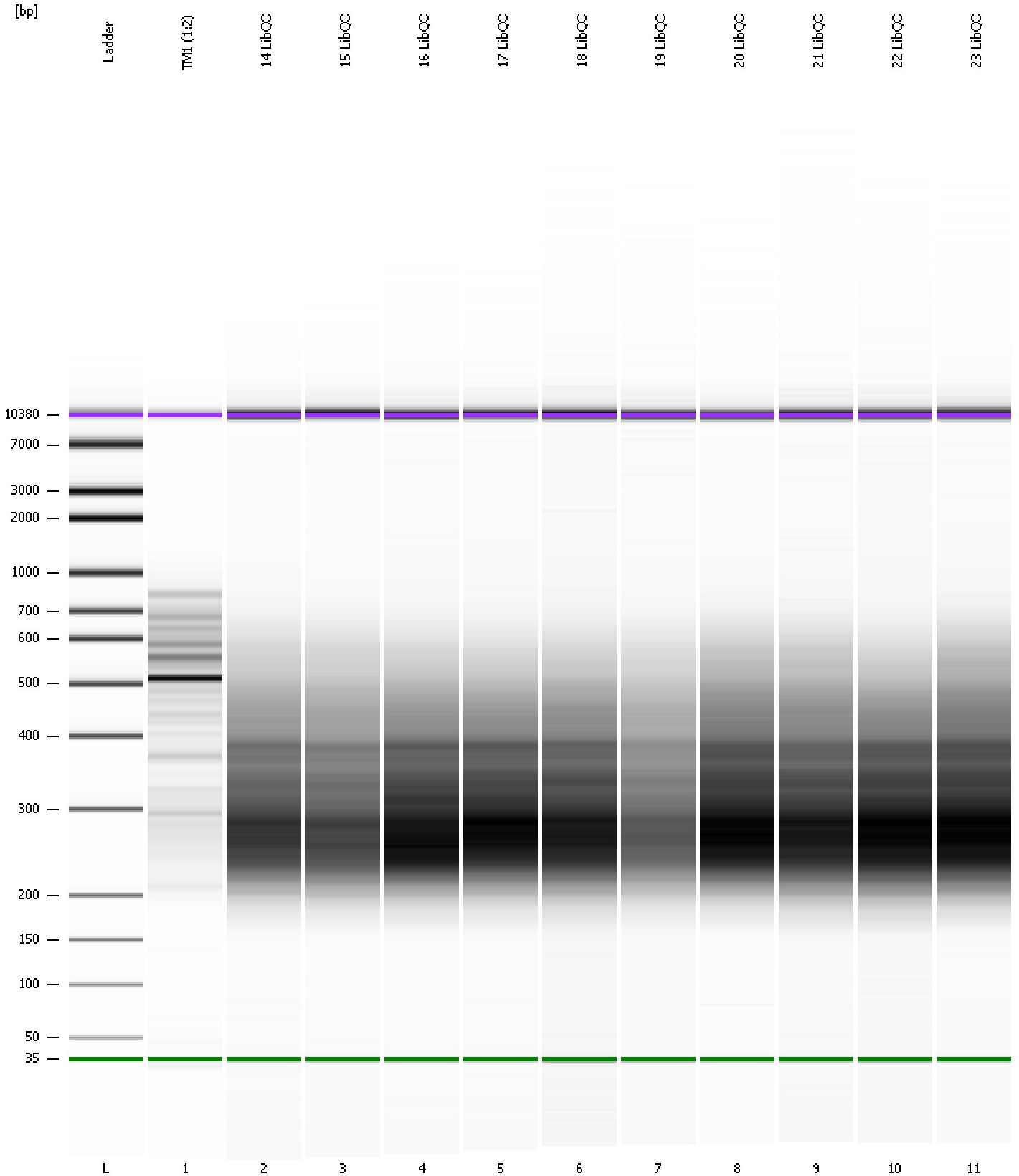
From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	348	2,818.32	4,017.3	13,786.5	96	31.8



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
Modified: 10/18/2019 12:01:45 PM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-18\2019-10-18\_001.xad

Created: 10/18/2019 11:20:24 AM  
 Modified: 10/18/2019 12:01:45 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		10/18/2019 12:01:40 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\2019-10-18\2019-10-18_001.xad)		Instrument	Run		10/18/2019 11:20:29 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		10/18/2019 11:20:29 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		10/18/2019 11:20:29 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		10/18/2019 11:20:29 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		10/18/2019 11:20:29 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		10/18/2019 11:20:29 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		10/18/2019 11:20:29 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1