

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
Modified: 7/3/2019 1:09:59 PM

**Electrophoresis File Run Summary**

Instrument Information:

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

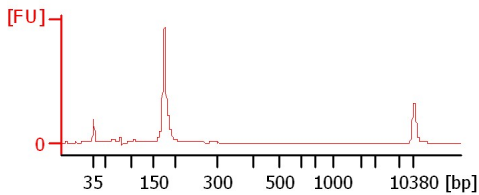
Assay Information:

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

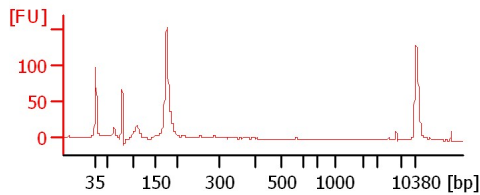
Chip Information:

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

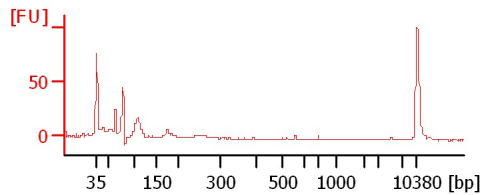
**PoolA\_ST-3\_LIB**



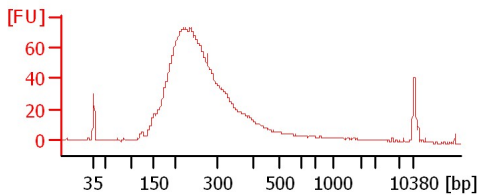
**PoolB\_ST-3\_LIB**



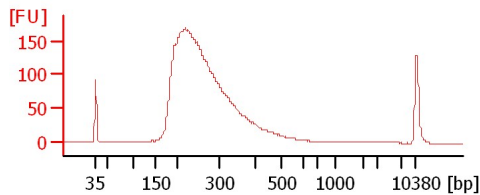
**PoolC\_ST-3\_LIB**



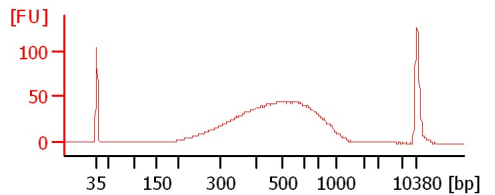
**H1829P\_Ikoh**



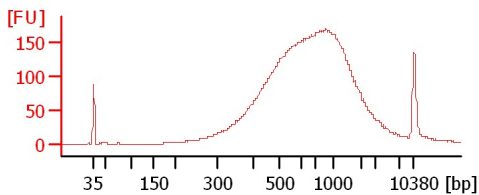
**AW070319 (1:2)**



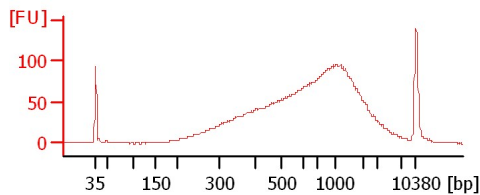
**seq library 6-2MN**



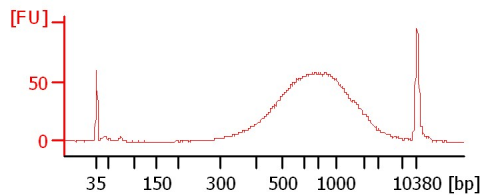
**H1826-10**



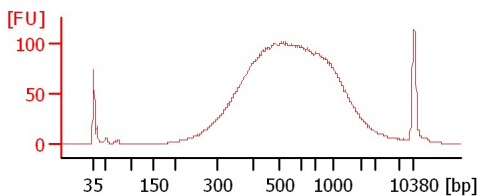
**H1826-12**



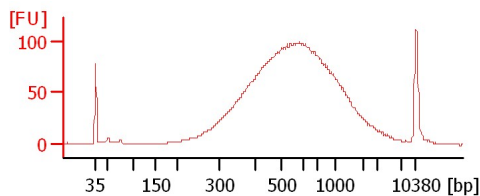
**H1826-16**



**H1828-20**



**H1826-21**



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 Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
PoolA_ST-3_LIB		<input type="checkbox"/>	✓			
PoolB_ST-3_LIB		<input type="checkbox"/>	✓			
PoolC_ST-3_LIB		<input type="checkbox"/>	✓			
H1829P_Ikoh		<input type="checkbox"/>	✓			
AW070319 (1:2)		<input type="checkbox"/>	✓			
seq library 6-2MN		<input type="checkbox"/>	✓			
H1826-10		<input type="checkbox"/>	✓			
H1826-12		<input type="checkbox"/>	✓			
H1826-16		<input type="checkbox"/>	✓			
H1828-20		<input type="checkbox"/>	✓			
H1826-21		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

**Chip Lot #**

**Reagent Kit Lot #**

**Chip Comments :**

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Created: 7/3/2019 12:28:38 PM  
Modified: 7/3/2019 1:09:59 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

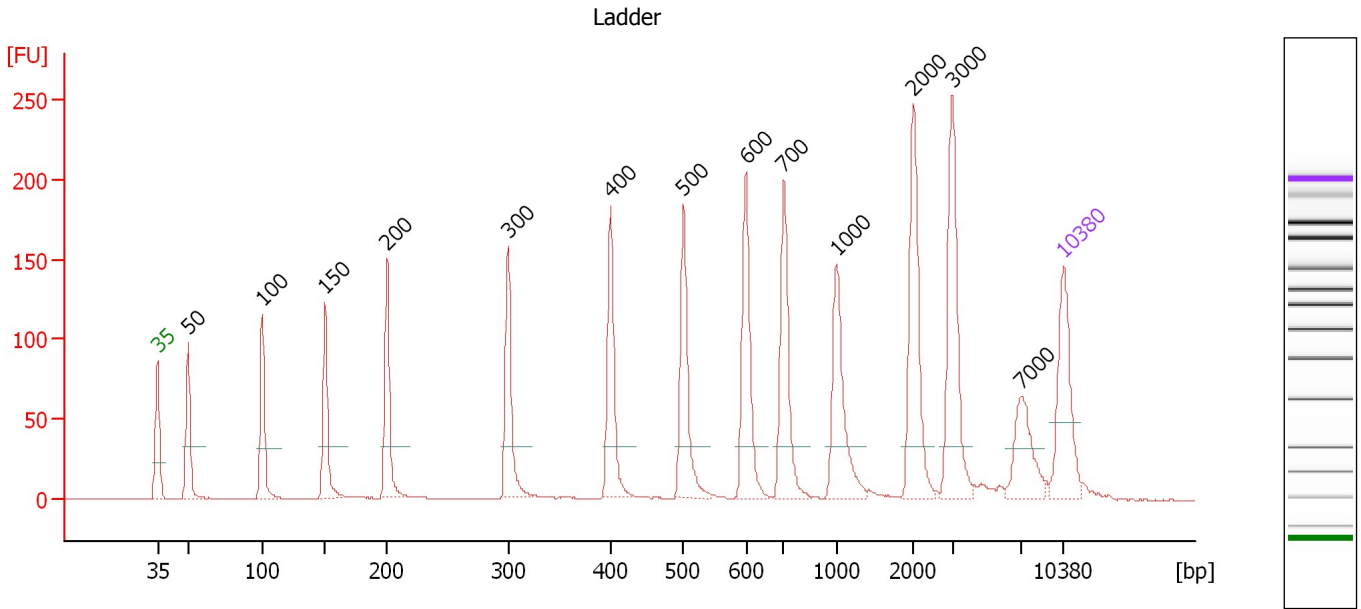
**Ladder**

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

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 Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.3

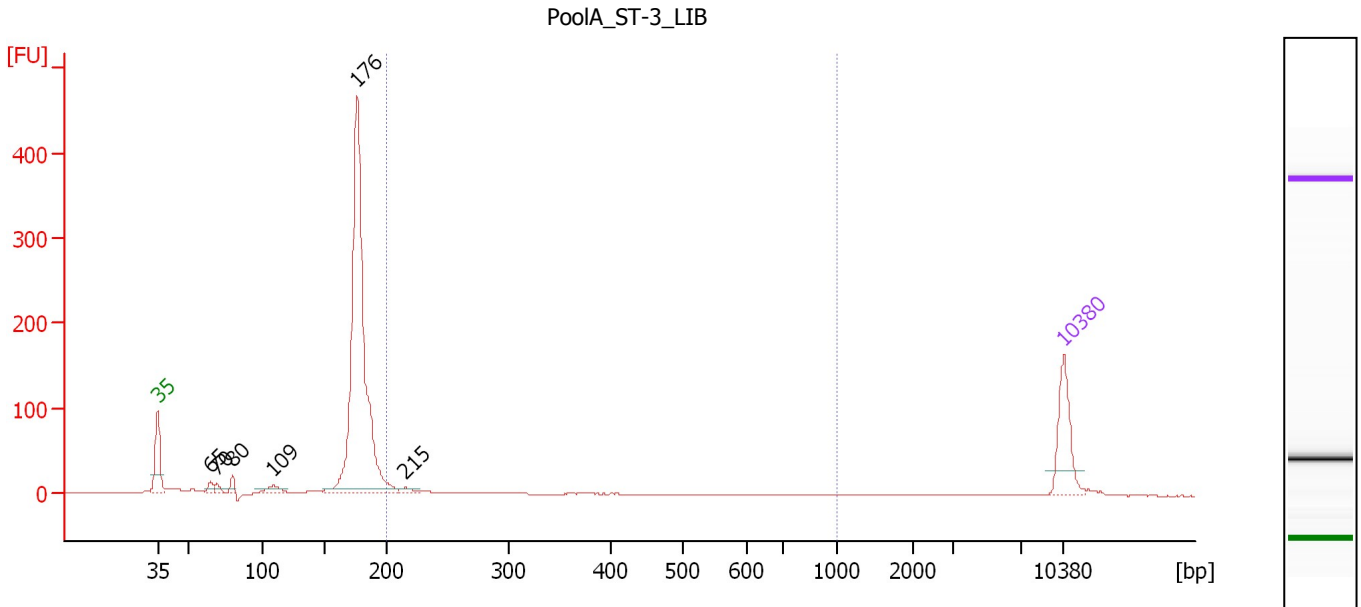
**Peak table for Ladder**

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	45.36
3	100	150.00	2,272.7	Ladder Peak	51.06
4	150	150.00	1,515.2	Ladder Peak	55.96
5	200	150.00	1,136.4	Ladder Peak	60.72
6	300	150.00	757.6	Ladder Peak	70.06
7	400	150.00	568.2	Ladder Peak	77.98
8	500	150.00	454.5	Ladder Peak	83.63
9	600	150.00	378.8	Ladder Peak	88.44
10	700	150.00	324.7	Ladder Peak	91.36
11	1,000	150.00	227.3	Ladder Peak	95.42
12	2,000	150.00	113.6	Ladder Peak	101.36
13	3,000	150.00	75.8	Ladder Peak	104.37
14	7,000	150.00	32.5	Ladder Peak	109.70
15	10,380	75.00	10.9	Upper Marker	113.00

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Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : PoolA ST-3 LIB**

Number of peaks found: 6                      Corr. Area 1: 82.4  
 Noise: 0.3

**Peak table for sample 1 : PoolA ST-3 LIB**

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	65	16.69	386.8		47.11
3	70	11.72	255.4		47.59
4	80	14.90	282.2		48.78
5	109	23.56	327.7		51.94
6	176	822.82	7,085.7		58.43
7	215	11.14	78.5		62.11
8	10,380	75.00	10.9	Upper Marker	113.00

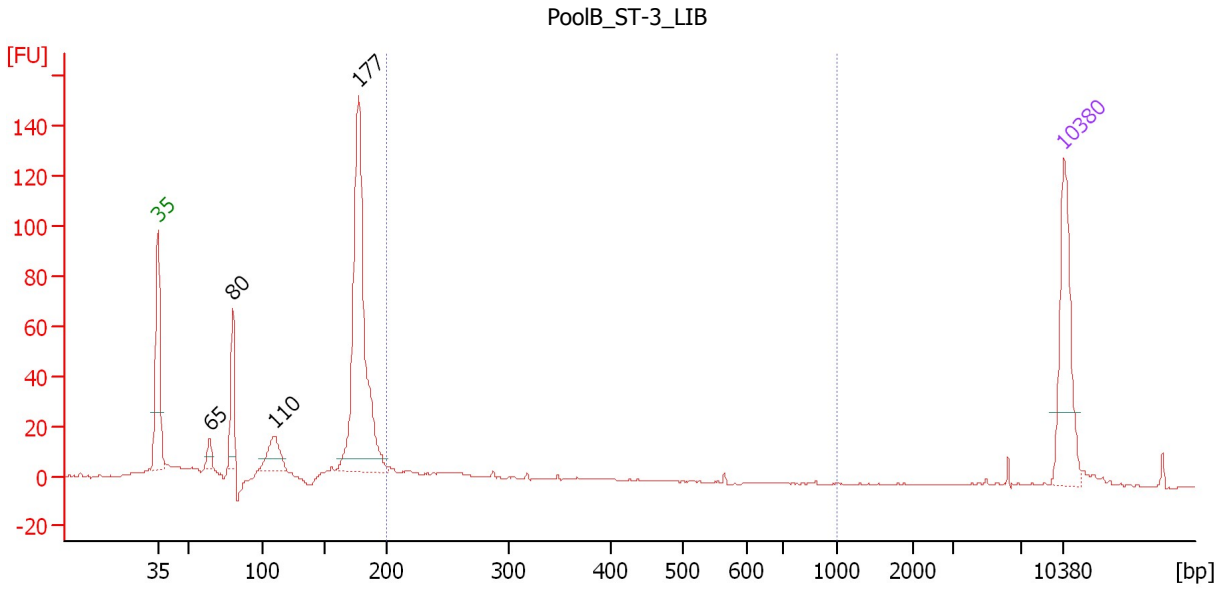
**Region table for sample 1 : PoolA ST-3 LIB**

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	341	64.30	82.4	359.9	7	50.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : PoolB\_ST-3\_LIB**

Number of peaks found: 4                      Corr. Area 1: 43.6  
 Noise: 0.3

**Peak table for sample 2 : PoolB\_ST-3\_LIB**

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	65	13.88	325.4		47.02
3	80	63.92	1,205.6		48.82
4	110	40.10	553.6		52.02
5	177	325.90	2,793.3		58.51
6	10,380	75.00	10.9	Upper Marker	113.00

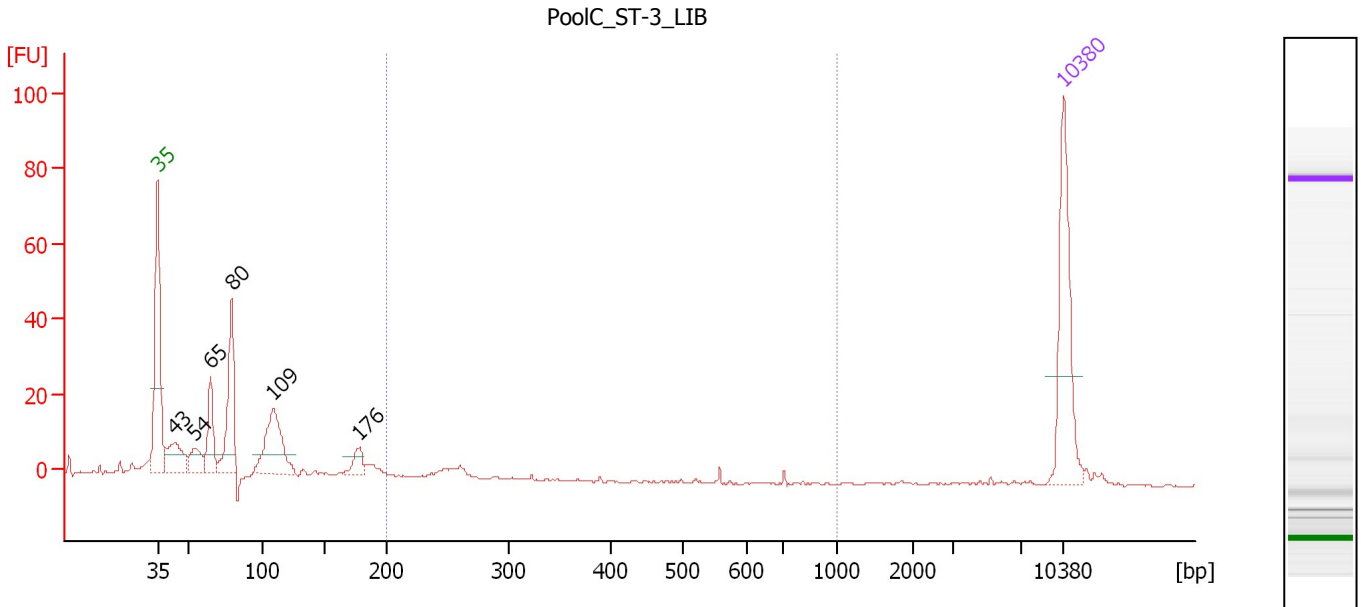
**Region table for sample 2 : PoolB\_ST-3\_LIB**

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	282	46.55	43.6	271.3	9	31.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : PoolC\_ST-3\_LIB**

Number of peaks found: 6                      Corr. Area 1: 11.8  
 Noise: 0.3

**Peak table for sample 3 : PoolC\_ST-3\_LIB**

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	43	49.50	1,724.5		44.33
3	54	26.94	749.6		45.87
4	65	49.29	1,144.0		47.10
5	80	92.29	1,757.3		48.73
6	109	90.93	1,264.2		51.94
7	176	17.12	147.1		58.46
8	10,380	75.00	10.9	Upper Marker	113.00

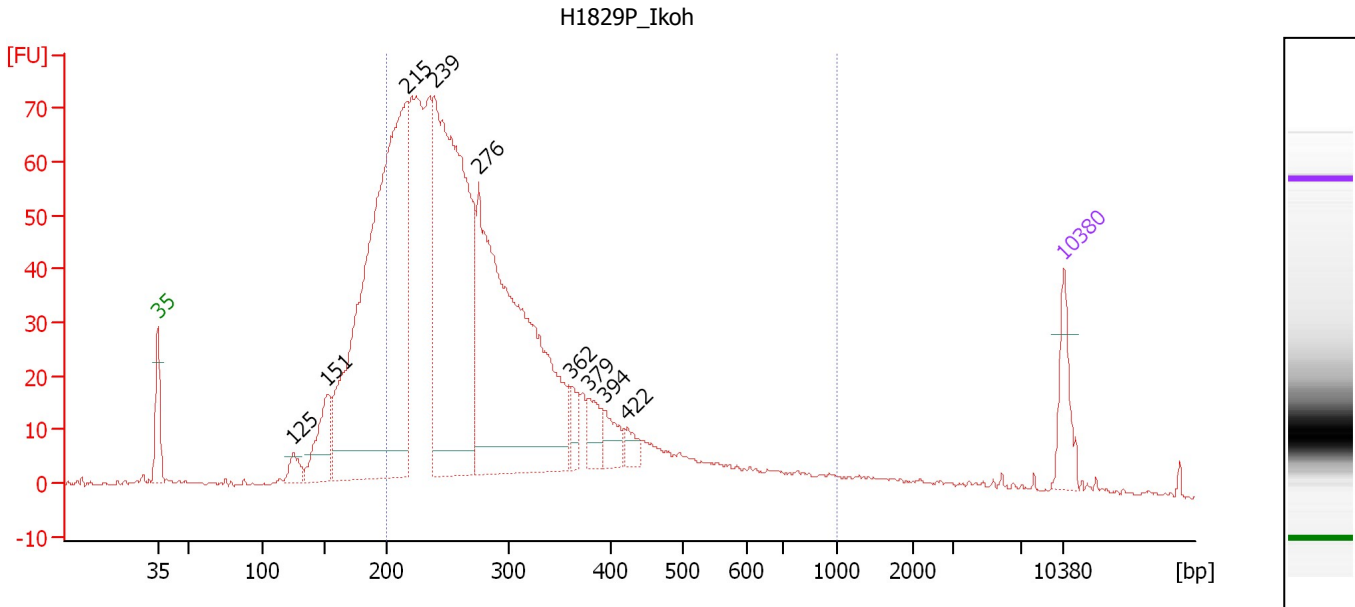
**Region table for sample 3 : PoolC\_ST-3\_LIB**

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	309	16.59	11.8	94.5	5	44.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : H1829P\_Ikoh**

Number of peaks found: 9                      Corr. Area 1: 1,298.1  
 Noise: 0.3

**Peak table for sample 4 : H1829P\_Ikoh**

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	125	45.72	555.3		53.49
3	151	164.73	1,654.3		56.05
4	215	1,698.53	11,943.4		62.17
5	239	1,283.49	8,132.2		64.38
6	276	1,248.78	6,863.0		67.79
7	362	44.91	188.0		74.96
8	379	63.06	252.1		76.32
9	394	62.91	242.1		77.47
10	422	32.52	116.7		79.23
11	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 4 : H1829P\_Ikoh**

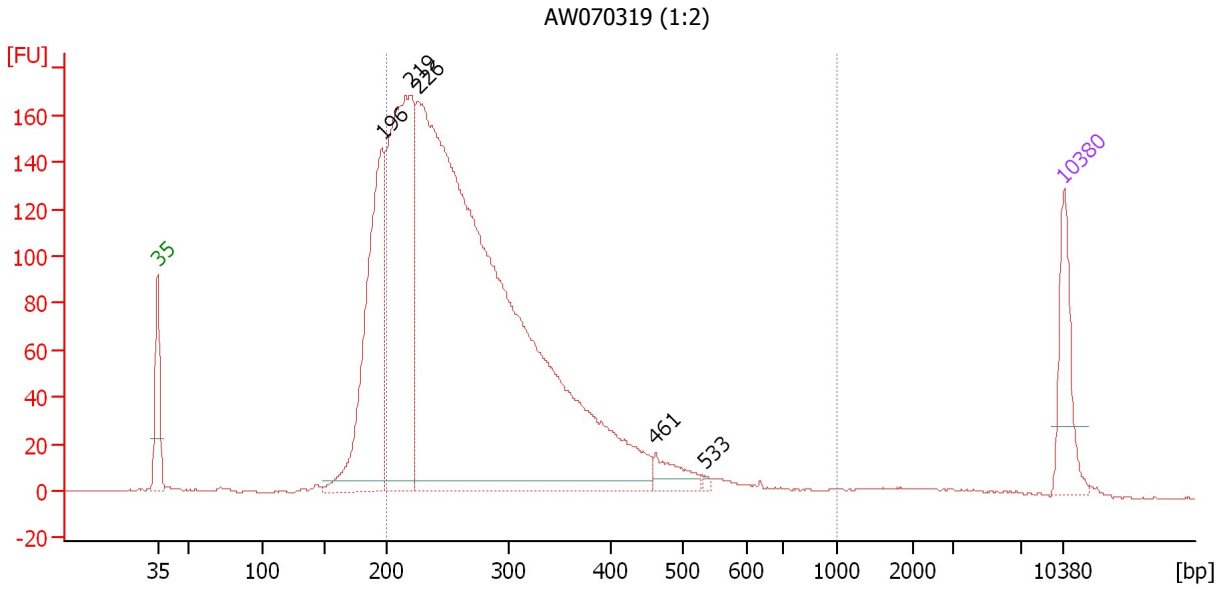
From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	302	4,818.17	1,298.1	26,983.5	77	37.4



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : AW070319 (1:2)**

Number of peaks found: 5                      Corr. Area 1: 2,775.1  
 Noise: 0.2

**Peak table for sample 5 : AW070319 (1:2)**

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	196	466.49	3,610.7		60.32
3	219	655.07	4,523.5		62.54
4	226	2,346.53	15,728.6		63.15
5	461	42.84	140.6		81.45
6	533	4.33	12.3		85.22
7	10,380	75.00	10.9	Upper Marker	113.00

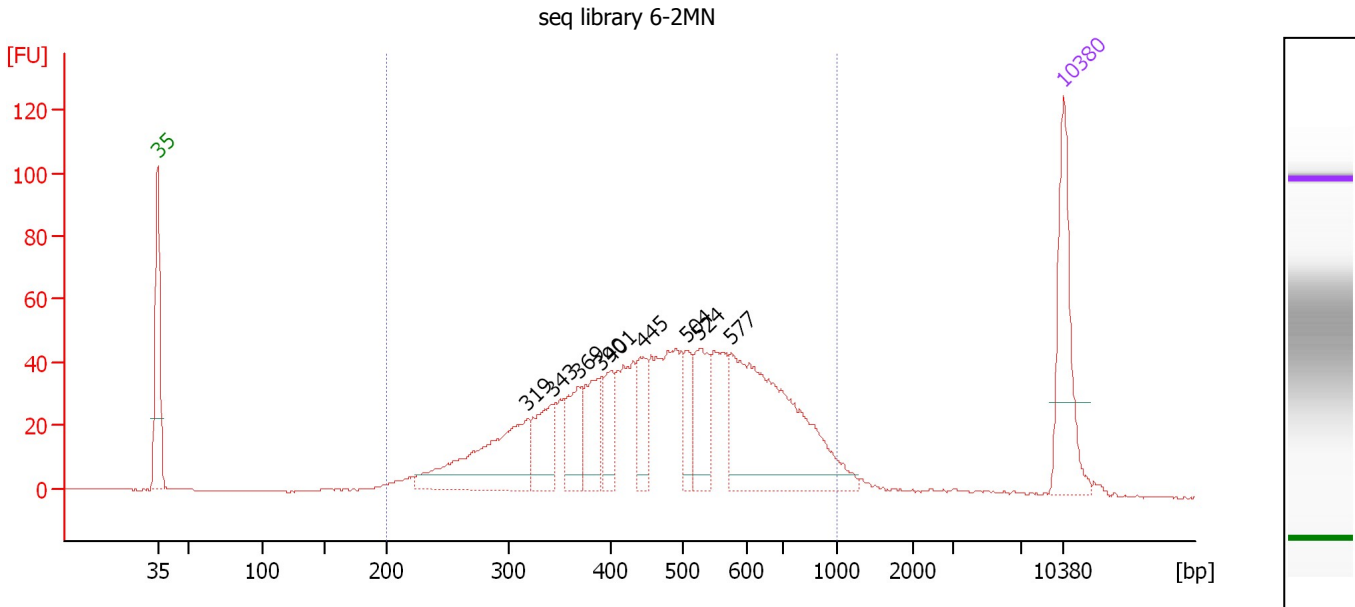
**Region table for sample 5 : AW070319 (1:2)**

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	288	2,788.64	2,775.1	15,864.8	84	31.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : seq library 6-2MN**

Number of peaks found: 9                      Corr. Area 1: 1,212.0  
 Noise: 0.2

**Peak table for sample 6 : seq library 6-2MN**

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	319	143.03	679.5		71.56
3	343	58.43	258.2		73.45
4	369	51.11	210.0		75.50
5	390	54.25	210.7		77.18
6	401	31.95	120.8		78.03
7	445	39.32	134.0		80.50
8	504	37.19	111.9		83.81
9	524	54.27	156.8		84.81
10	577	230.68	605.8		87.34
11	10,380	75.00	10.9	Upper Marker	113.00

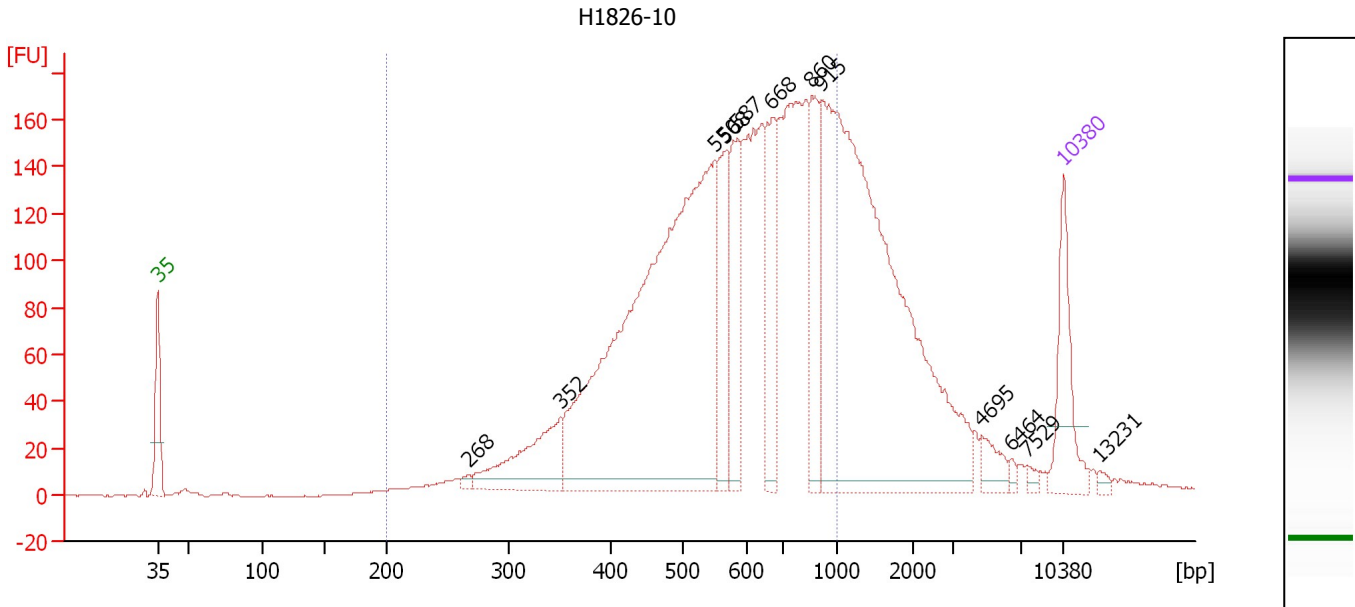
**Region table for sample 6 : seq library 6-2MN**

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	491	1,021.29	1,212.0	3,653.3	96	32.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : H1826-10**

Number of peaks found: 12                      Corr. Area 1: 3,166.6  
 Noise: 0.3

**Peak table for sample 7 : H1826-10**

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	268	7.47	42.3		67.03
3	352	131.75	567.7		74.15
4	550	902.95	2,488.4		86.03
5	568	105.23	280.9		86.88
6	587	125.59	324.4		87.79
7	668	124.98	283.7		90.42
8	860	107.81	190.0		93.52
9	915	816.59	1,352.2		94.27
10	4,695	23.82	7.7		106.63
11	6,464	5.28	1.2		108.99
12	7,529	5.87	1.2		110.22
13	10,380	75.00	10.9	Upper Marker	113.00
14	13,231	0.00	0.0		115.78

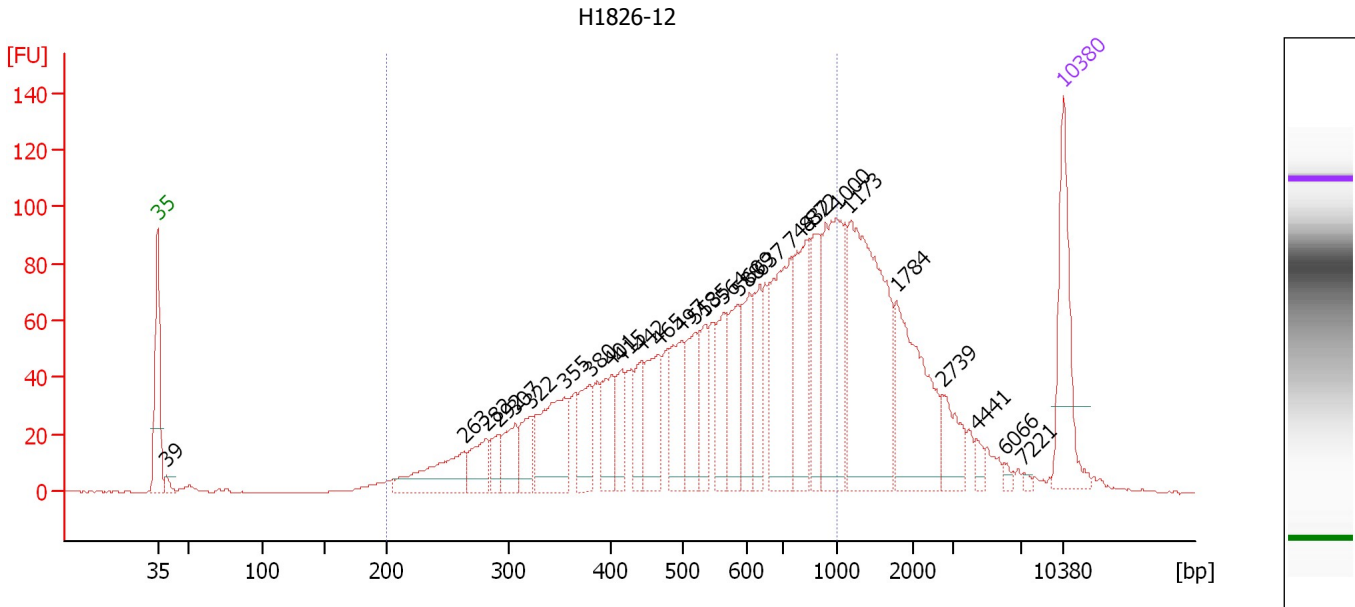
**Region table for sample 7 : H1826-10**

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	597	2,400.28	3,166.6	6,971.5	73	29.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : H1826-12**

Number of peaks found: 29                      Corr. Area 1: 1,820.6  
 Noise: 0.3

**Peak table for sample 8 : H1826-12**

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	39	7.74	297.2		43.70
3	263	77.98	448.6		66.64
4	282	42.00	225.7		68.37
5	292	24.32	126.3		69.29
6	307	39.47	194.5		70.64
7	322	36.72	172.5		71.83
8	355	99.16	423.8		74.38
9	380	50.38	201.0		76.38
10	401	48.19	181.9		78.05
11	415	34.99	127.8		78.81
12	442	39.09	133.9		80.38
13	465	67.26	218.9		81.68
14	497	63.97	195.0		83.46
15	518	53.56	156.7		84.49
16	535	39.74	112.6		85.30
17	564	52.80	141.9		86.71
18	588	59.93	154.5		87.85
19	609	55.67	138.5		88.71
20	637	44.53	105.9		89.52
21	744	125.22	255.0		91.96
22	832	92.60	168.6		93.15
23	872	59.55	103.5		93.69
24	1,000	131.12	198.6		95.42
25	1,173	220.56	284.8		96.45

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Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**

**... Peak table for sample 8 : H1826-12**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	1,784	115.60	98.2		100.07
27	2,739	31.96	17.7		103.59
28	4,441	7.15	2.4		106.29
29	6,066	3.44	0.9		108.46
30	7,221	2.30	0.5		109.92
31	10,380	75.00	10.9	Upper Marker	113.00

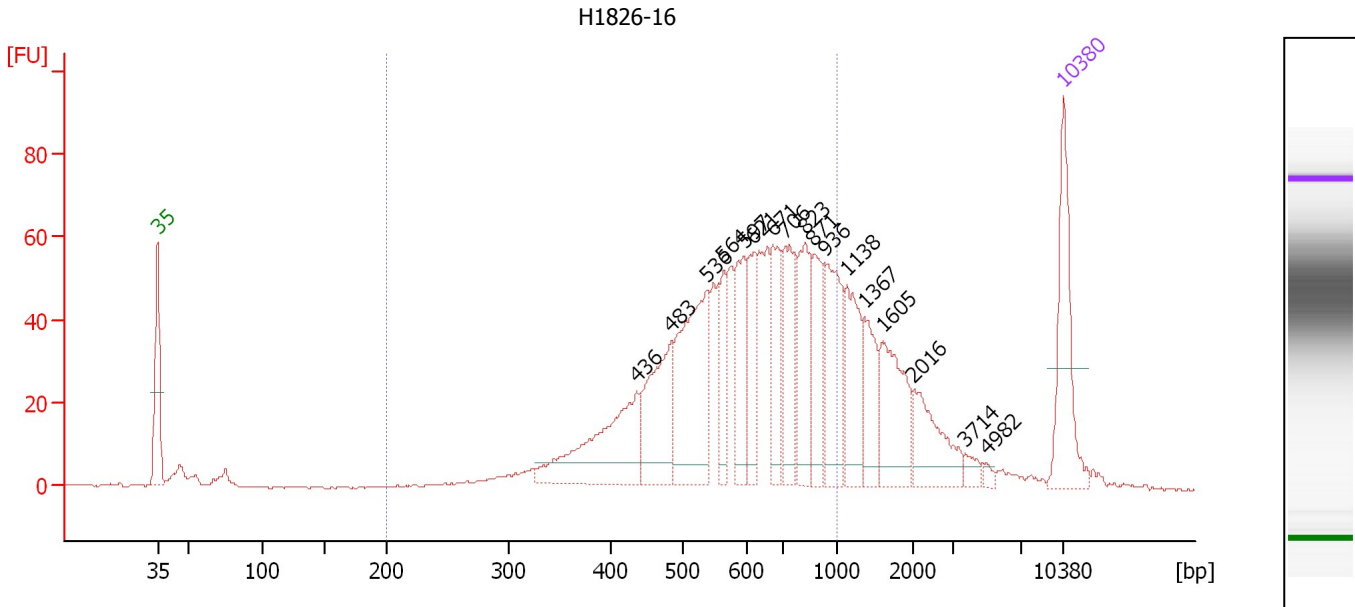
**Region table for sample 8 : H1826-12**

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	565	1,550.06	1,820.6	5,071.4	70	35.4

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Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : H1826-16**

Number of peaks found: 17      Corr. Area 1: 1,008.4  
 Noise: 0.3

**Peak table for sample 9 : H1826-16**

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	436	136.84	475.5		80.01
3	483	102.35	321.0		82.68
4	536	157.02	444.1		85.35
5	564	47.19	126.8		86.71
6	597	62.98	159.9		88.29
7	621	57.20	139.6		89.05
8	671	59.41	134.1		90.52
9	706	74.84	160.6		91.44
10	823	69.54	128.0		93.02
11	871	55.09	95.8		93.68
12	936	83.99	136.0		94.55
13	1,138	70.81	94.3		96.23
14	1,367	42.71	47.4		97.60
15	1,605	72.10	68.1		99.01
16	2,016	52.42	39.4		101.41
17	3,714	8.35	3.4		105.33
18	4,982	4.15	1.3		107.01
19	10,380	75.00	10.9	Upper Marker	113.00

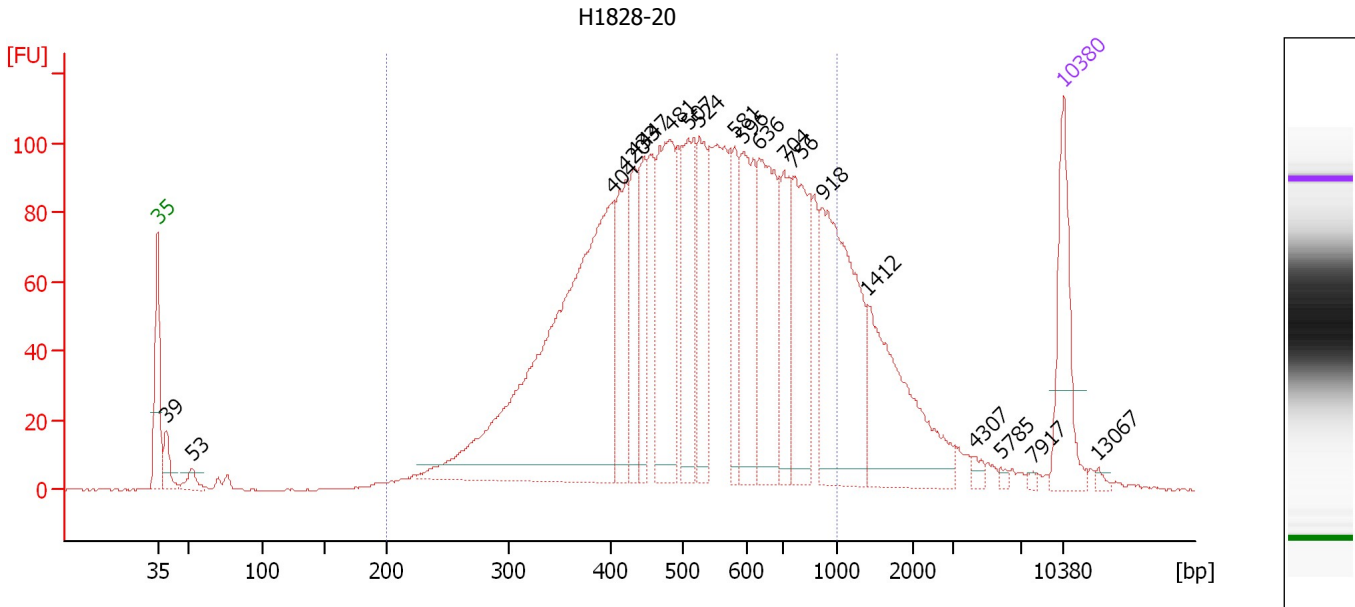
**Region table for sample 9 : H1826-16**

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	614	1,171.69	1,008.4	3,246.1	72	27.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : H1828-20**

Number of peaks found: 20      Corr. Area 1: 2,614.1  
 Noise: 0.3

**Peak table for sample 10 : H1828-20**

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	39	30.77	1,190.1		43.66
3	53	16.61	476.6		45.68
4	404	600.49	2,253.2		78.19
5	420	107.24	386.7		79.12
6	433	77.22	270.3		79.83
7	447	78.87	267.2		80.65
8	481	186.43	587.2		82.56
9	507	122.88	367.0		83.98
10	524	101.59	293.5		84.80
11	581	69.81	182.0		87.54
12	596	128.69	327.2		88.25
13	636	155.18	369.5		89.50
14	704	88.25	190.0		91.42
15	756	119.76	239.9		92.13
16	918	223.10	368.2		94.31
17	1,412	150.76	161.8		97.86
18	4,307	5.96	2.1		106.11
19	5,785	3.15	0.8		108.08
20	7,917	2.57	0.5		110.60
21	10,380	75.00	10.9	Upper Marker	113.00
22	13,067	0.00	0.0		115.62

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**

... Region table for sample 10 : H1828-20

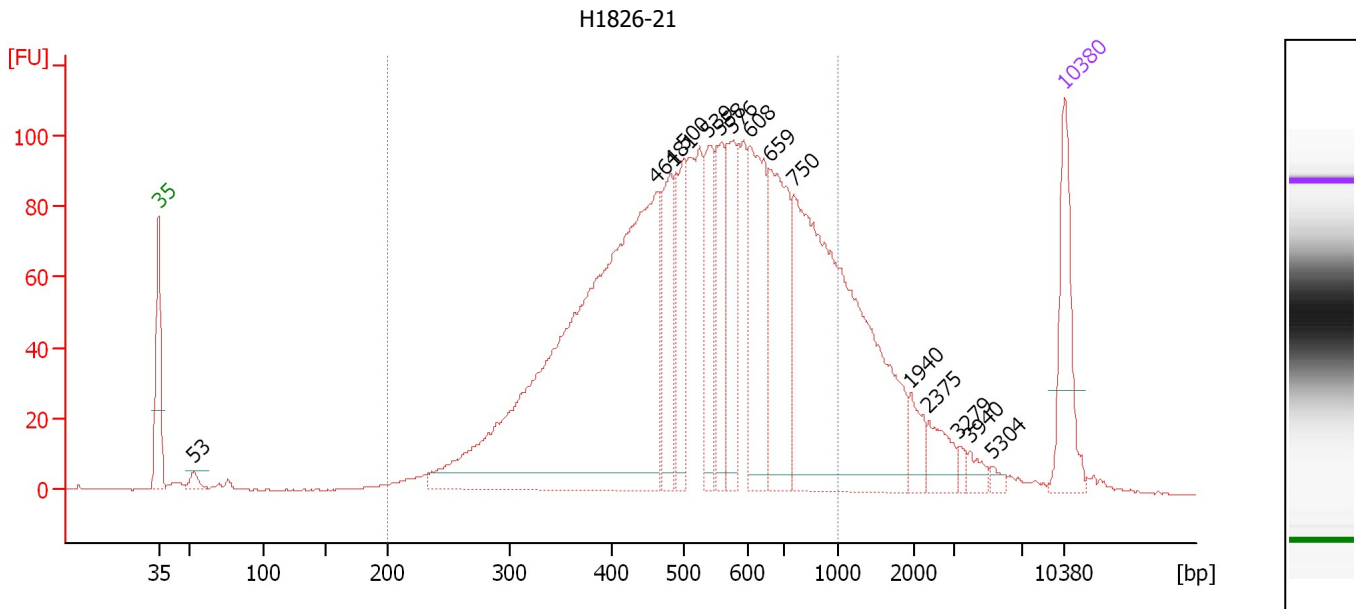
From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ $\mu$ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	535	2,500.80	2,614.1	8,189.4	 83	32.3



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : H1826-21**

Number of peaks found: 15                      Corr. Area 1: 2,337.1  
 Noise: 0.3

**Peak table for sample 11 : H1826-21**

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	12.68	362.5		45.70
3	461	855.30	2,812.7		81.41
4	481	93.39	294.1		82.57
5	500	84.30	255.6		83.61
6	539	90.15	253.6		85.48
7	558	88.35	239.9		86.42
8	576	104.06	273.6		87.30
9	608	150.26	374.4		88.68
10	659	165.56	380.7		90.16
11	750	510.18	1,031.3		92.03
12	1,940	25.01	19.5		101.00
13	2,375	28.71	18.3		102.49
14	3,279	6.40	3.0		104.75
15	3,940	10.96	4.2		105.63
16	5,304	4.69	1.3		107.44
17	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 11 : H1826-21**

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	541	2,314.51	2,337.1	7,478.1	84	31.6

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Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
Modified: 7/3/2019 1:09:59 PM

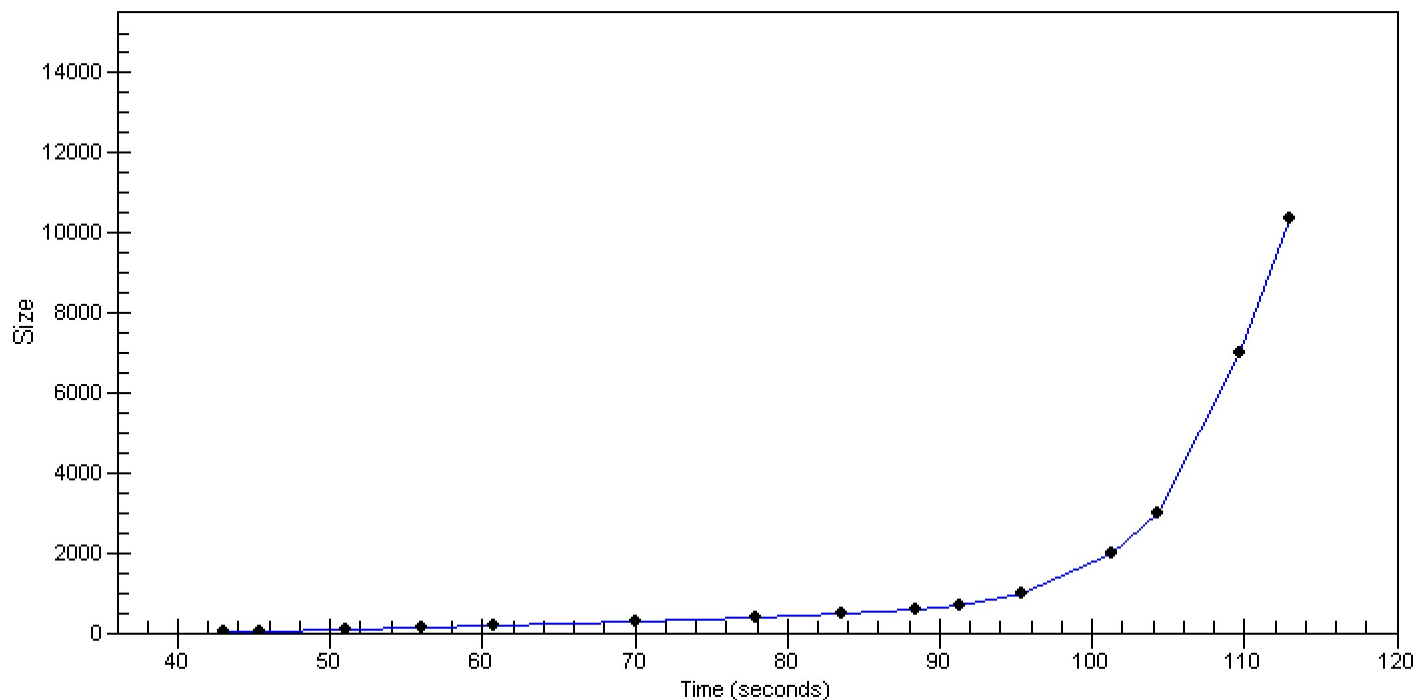
**Gel Image**

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
Modified: 7/3/2019 1:09:59 PM

**Curves**

**Standard Curve**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-07-03\2019-07-03\_001.xad

Created: 7/3/2019 12:28:38 PM  
 Modified: 7/3/2019 1:09:59 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		7/3/2019 1:09:57 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\2019-07-03\2019-07-03_001.xad)		Instrument	Run		7/3/2019 12:28:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		7/3/2019 12:28:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		7/3/2019 12:28:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		7/3/2019 12:28:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		7/3/2019 12:28:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		7/3/2019 12:28:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		7/3/2019 12:28:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1