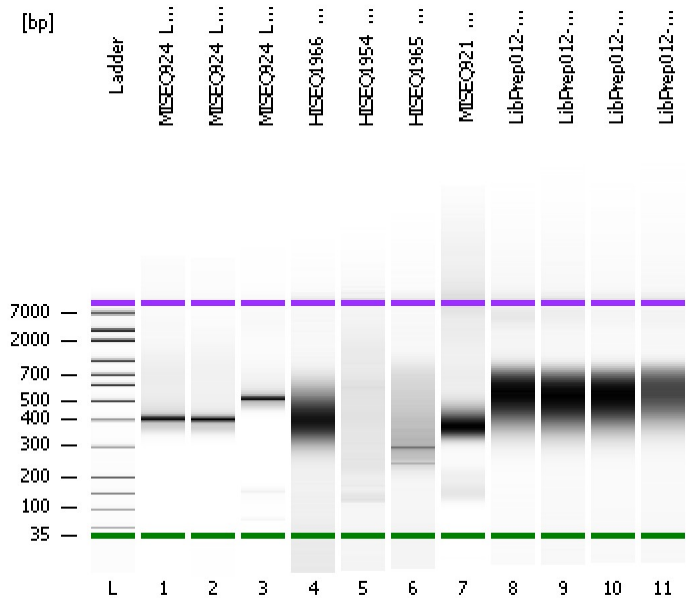


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
Modified: 10/15/2019 4:43:55 PM

Electrophoresis File Run Summary



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

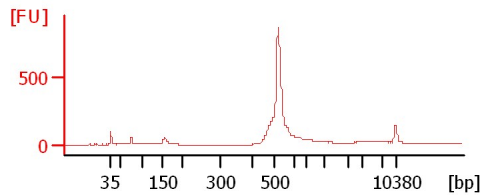
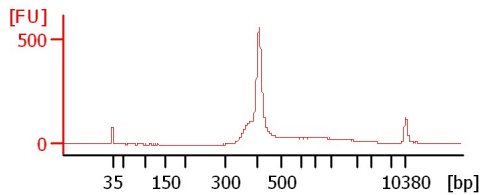
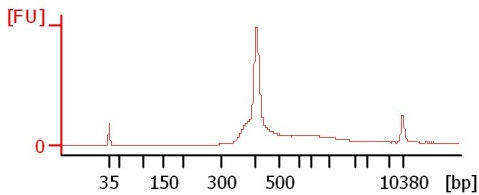
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 Assay Origin Path: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\assays\dsDNA\High Sensitivity DNA.xsy  
 Assay Class: High Sensitivity DNA Assay  
 Version: 1.03  
 Assay Comments: Copyright © 2003-2010 Agilent Technologies

**Chip Information:**  
 Chip Lot #:   
 Reagent Kit Lot #:   
 Chip Comments:

MISEQ924\_Library1

MISEQ924\_Library2

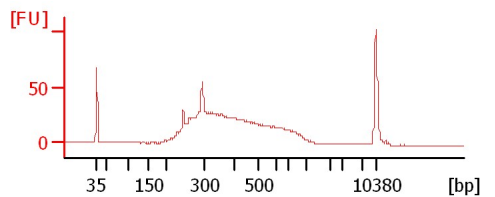
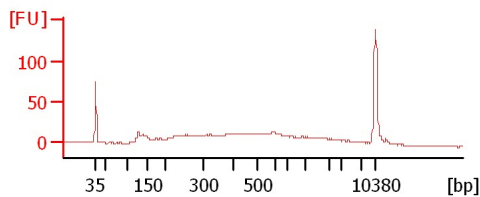
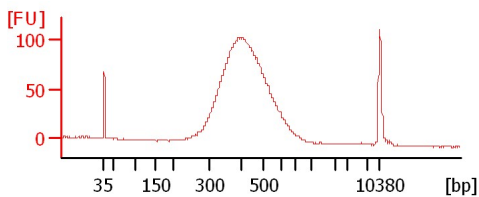
MISEQ924\_Library3



HISEQ1966\_Bzisolates.10.9.2019\_BC

HISEQ1954\_RFFJ5M01\_BC

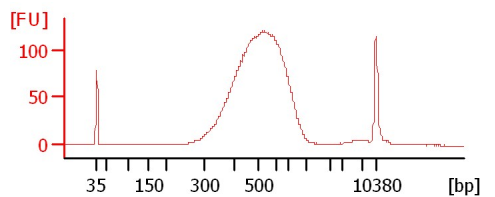
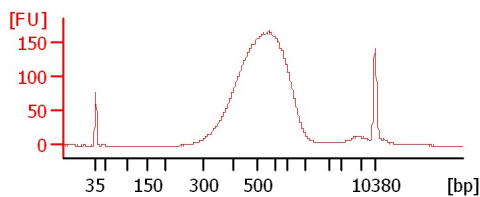
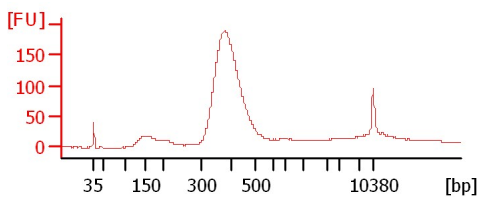
HISEQ1965\_PS2VB\_BC



MISEQ921\_Mac2\_BC

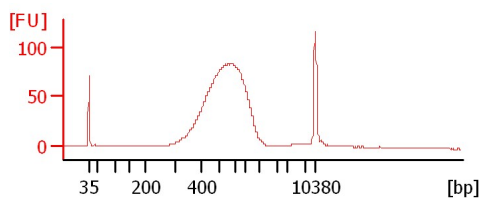
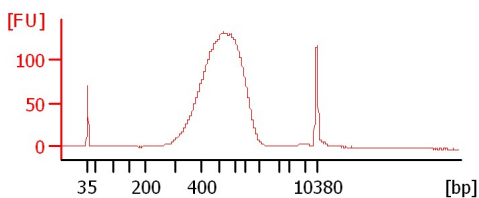
LibPrep012-1(1:4)

LibPrep012-2(1:6)



LibPrep012-3(1:5)

LibPrep012-12(1:6)



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
MISEQ924_Library1		<input type="checkbox"/>	✓			
MISEQ924_Library2		<input type="checkbox"/>	✓			
MISEQ924_Library3		<input type="checkbox"/>	✓			
HISEQ1966_Bzisolates.10.9.2019_ BC		<input type="checkbox"/>	✓			
HISEQ1954_RFFJ5M01_BC		<input type="checkbox"/>	✓			
HISEQ1965_PS2VB_BC		<input type="checkbox"/>	✓			
MISEQ921_Mac2_BC		<input type="checkbox"/>	✓			
LibPrep012-1(1:4)		<input type="checkbox"/>	✓			
LibPrep012-2(1:6)		<input type="checkbox"/>	✓			
LibPrep012-3(1:5)		<input type="checkbox"/>	✓			
LibPrep012-12(1:6)		<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\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
Modified: 10/15/2019 4:43:55 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

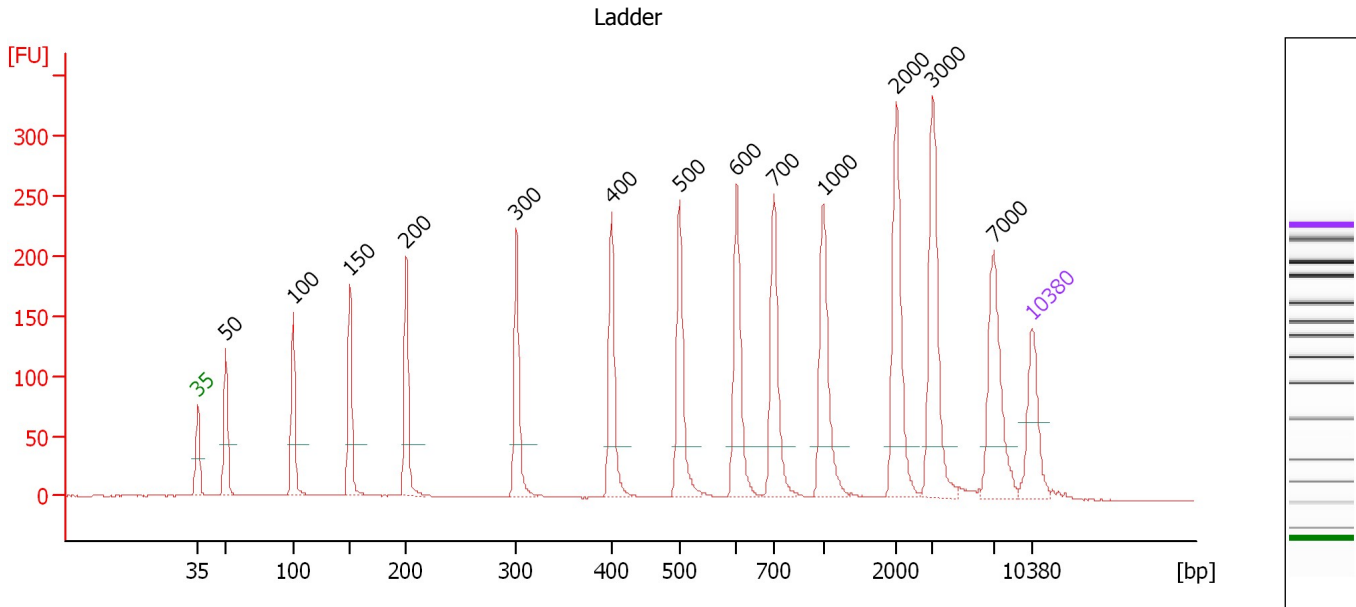
**Ladder**

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\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.2

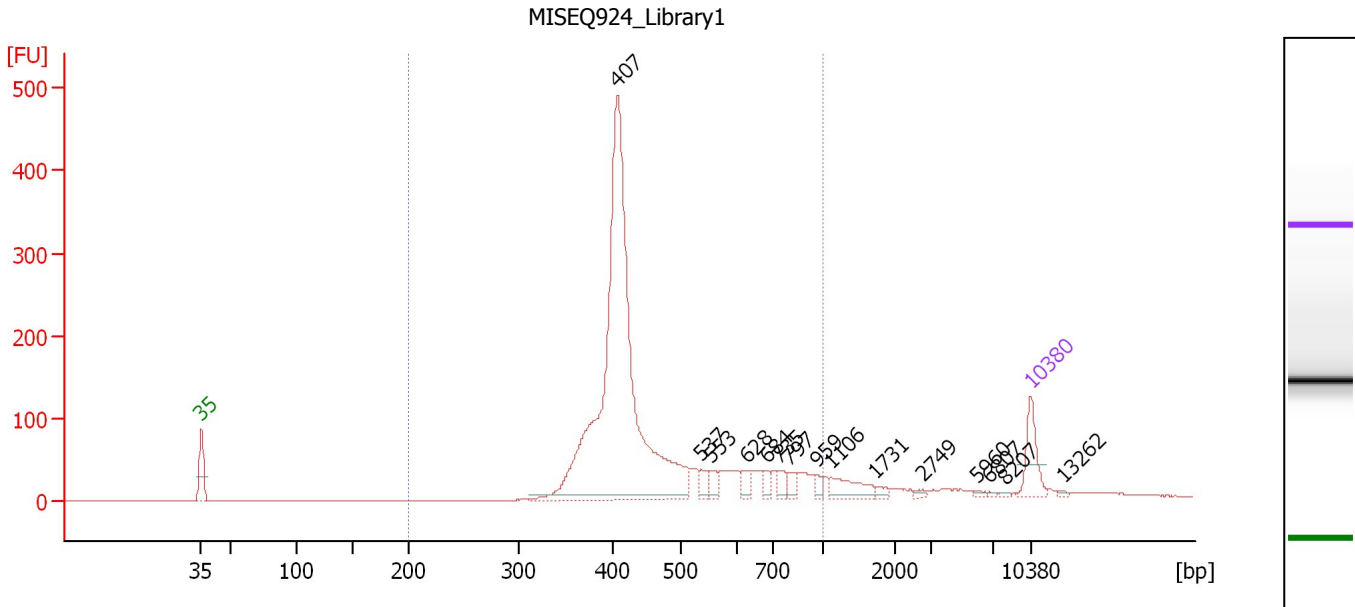
**Peak table for Ladder**

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	45.37
3	100	150.00	2,272.7	Ladder Peak	51.01
4	150	150.00	1,515.2	Ladder Peak	55.79
5	200	150.00	1,136.4	Ladder Peak	60.49
6	300	150.00	757.6	Ladder Peak	69.75
7	400	150.00	568.2	Ladder Peak	77.71
8	500	150.00	454.5	Ladder Peak	83.43
9	600	150.00	378.8	Ladder Peak	88.22
10	700	150.00	324.7	Ladder Peak	91.31
11	1,000	150.00	227.3	Ladder Peak	95.47
12	2,000	150.00	113.6	Ladder Peak	101.59
13	3,000	150.00	75.8	Ladder Peak	104.68
14	7,000	150.00	32.5	Ladder Peak	109.78
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : MISEQ924 Library1**

Number of peaks found: 15                      Corr. Area 1: 2,136.4  
 Noise: 0.3

**Peak table for sample 1 : MISEQ924 Library1**

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	407	1,664.14	6,194.5		78.11
3	537	29.72	83.8		85.23
4	553	29.71	81.4		85.99
5	628	27.85	67.1		89.10
6	684	26.64	59.0		90.81
7	735	25.47	52.5		91.80
8	797	27.88	53.0		92.65
9	959	16.27	25.7		94.90
10	1,106	63.79	87.3		96.12
11	1,731	11.63	10.2		99.95
12	2,749	6.86	3.8		103.91
13	5,960	5.45	1.4		108.45
14	6,807	3.18	0.7		109.53
15	8,207	3.53	0.7		110.93
16	10,380	75.00	10.9	Upper Marker	113.00
17	13,262	0.00	0.0		115.75

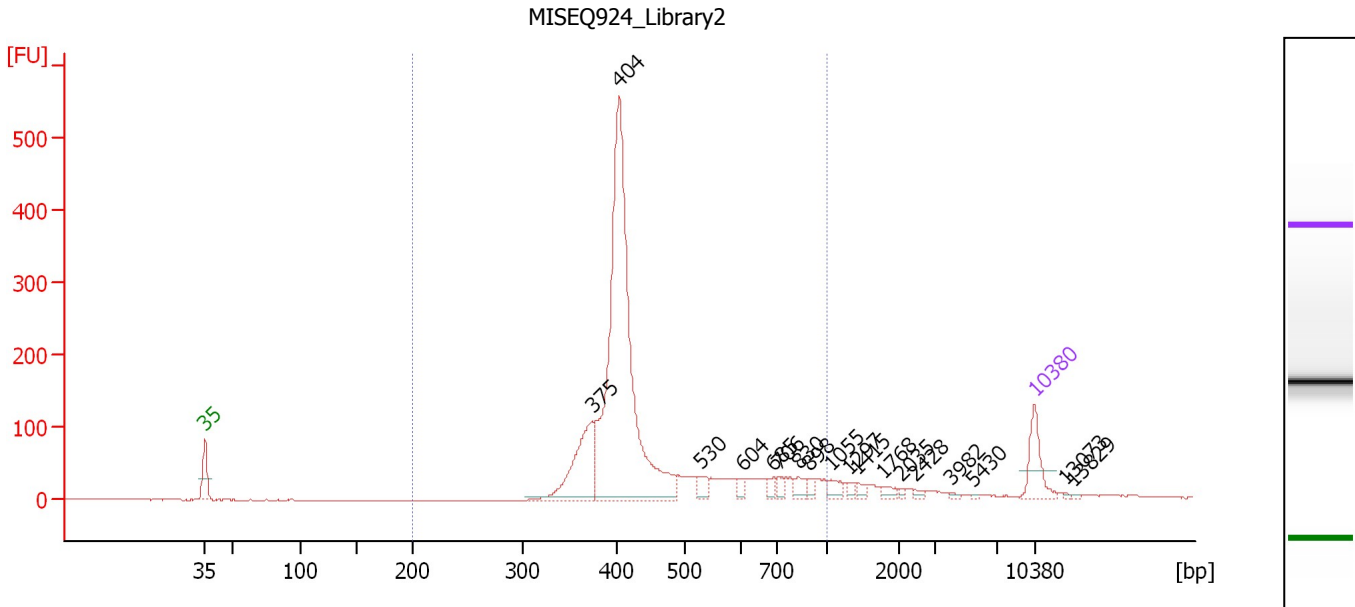
**Region table for sample 1 : MISEQ924 Library1**

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	473	2,025.90	2,136.4	6,958.9	90	27.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : MISEQ924\_Library2**

Number of peaks found: 18                      Corr. Area 1: 2,152.0  
 Noise: 0.4

**Peak table for sample 2 : MISEQ924\_Library2**

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	375	262.58	1,062.2		75.68
3	404	1,286.36	4,825.6		77.93
4	530	33.25	95.1		84.86
5	604	18.75	47.1		88.33
6	685	18.37	40.6		90.85
7	706	19.89	42.7		91.39
8	830	26.94	49.2		93.10
9	898	14.60	24.6		94.05
10	1,055	24.05	34.5		95.80
11	1,297	9.76	11.4		97.29
12	1,415	13.36	14.3		98.01
13	1,768	15.99	13.7		100.17
14	2,035	5.17	3.9		101.70
15	2,428	7.54	4.7		102.92
16	3,982	4.47	1.7		105.93
17	5,430	2.48	0.7		107.78
18	10,380	75.00	10.9	Upper Marker	113.00
19	13,073	0.00	0.0		115.57
20	13,829	0.00	0.0		116.29

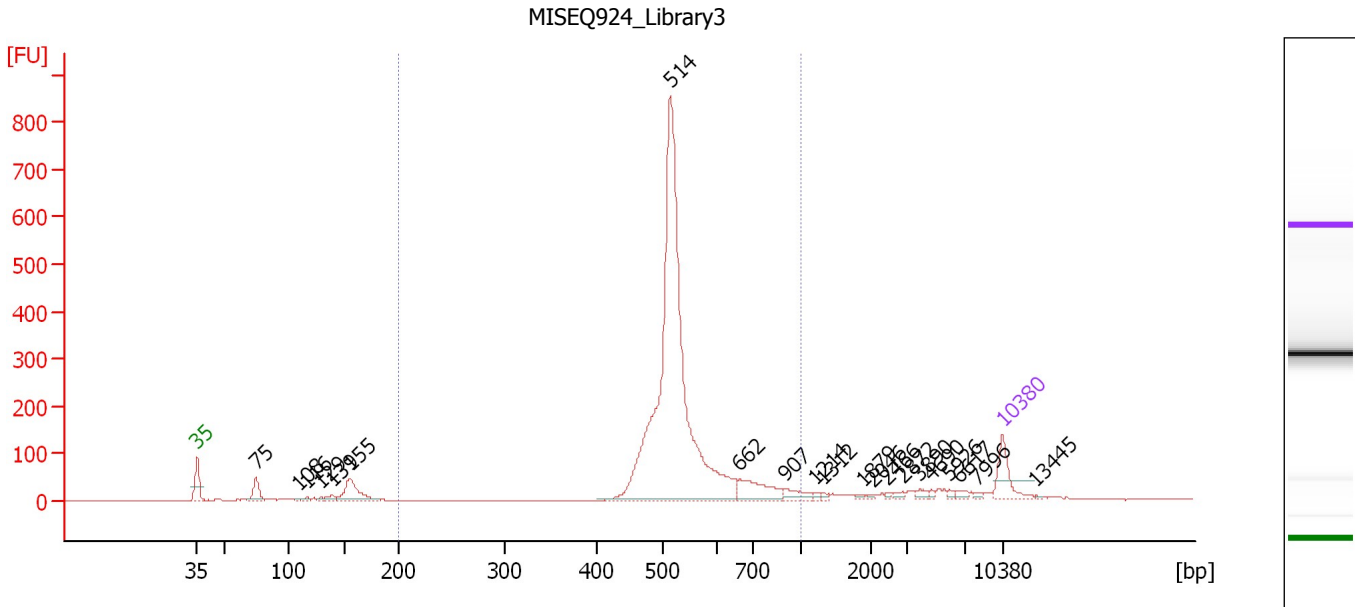
**Region table for sample 2 : MISEQ924\_Library2**

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	459	1,841.75	2,152.0	6,490.4	91	27.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : MISEQ924 Library3**

Number of peaks found: 21                      Corr. Area 1: 2,461.7  
 Noise: 0.6

**Peak table for sample 3 : MISEQ924 Library3**

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	75	71.81	1,458.1		48.15
3	108	6.25	87.3		51.81
4	116	10.40	135.7		52.55
5	129	8.26	97.2		53.75
6	139	17.97	196.0		54.73
7	155	129.37	1,265.0		56.26
8	514	1,767.27	5,206.4		84.12
9	662	106.69	244.0		90.15
10	907	33.69	56.3		94.18
11	1,214	8.02	10.0		96.77
12	1,312	5.91	6.8		97.38
13	1,879	6.71	5.4		100.85
14	2,046	5.36	4.0		101.74
15	2,466	4.67	2.9		103.03
16	2,872	9.06	4.8		104.28
17	3,890	13.13	5.1		105.81
18	4,690	6.61	2.1		106.83
19	5,926	8.20	2.1		108.41
20	6,617	11.19	2.6		109.29
21	7,996	6.46	1.2		110.73
22	10,380	75.00	10.9	Upper Marker	113.00
23	13,445	0.00	0.0		115.92

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary Continued ...****... Region table for sample 3 : MISEQ924 Library3**

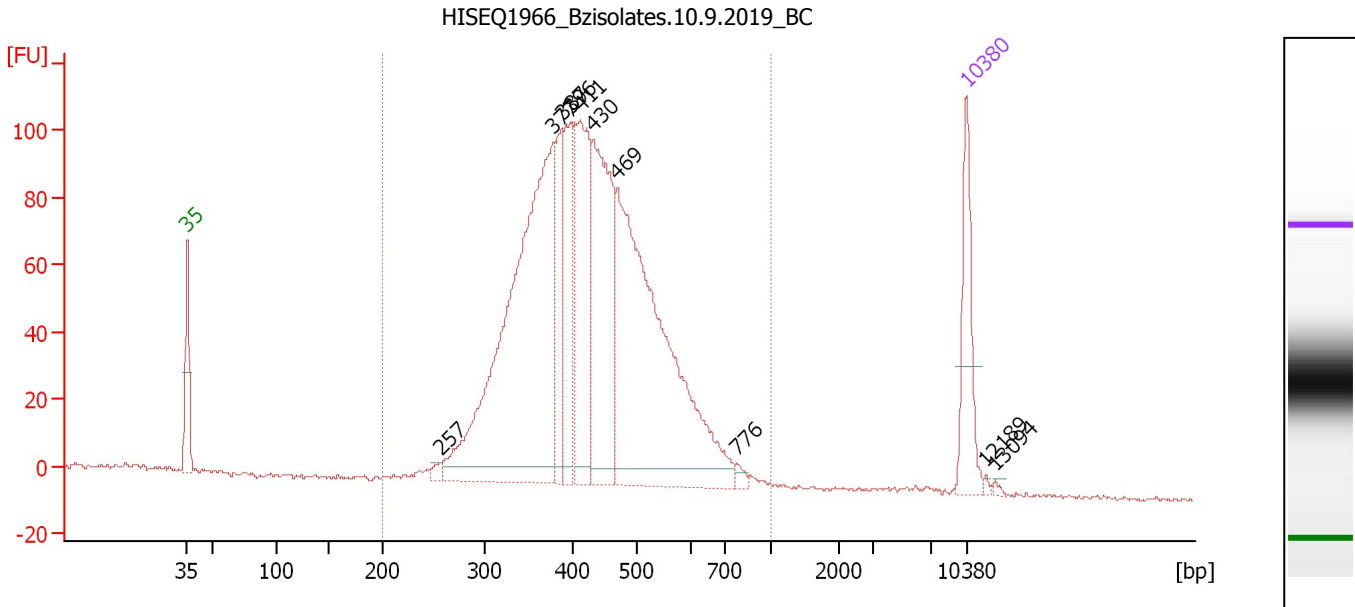
From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ $\mu$ l]	Corr. Area	Molarity [pmol/l]	Co lor % of Total	Size distribution in CV [%]
200	1,000	542	1,855.58	2,461.7	5,296.5	 83	15.2



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : HISEQ1966 Bzisolates.10.9.2019 BC**

Number of peaks found: 10                      Corr. Area 1: 1,889.5  
 Noise: 0.5

**Peak table for sample 4 : HISEQ1966 Bzisolates.10.9.2019 BC**

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	257	9.16	54.1		65.74
3	377	685.82	2,759.3		75.85
4	387	105.24	412.2		76.66
5	396	126.67	484.1		77.43
6	411	206.52	761.5		78.34
7	430	276.58	974.2		79.44
8	469	572.40	1,850.6		81.64
9	776	7.58	14.8		92.36
10	10,380	75.00	10.9	Upper Marker	113.00
11	12,189	0.00	0.0		114.72
12	13,094	0.00	0.0		115.59

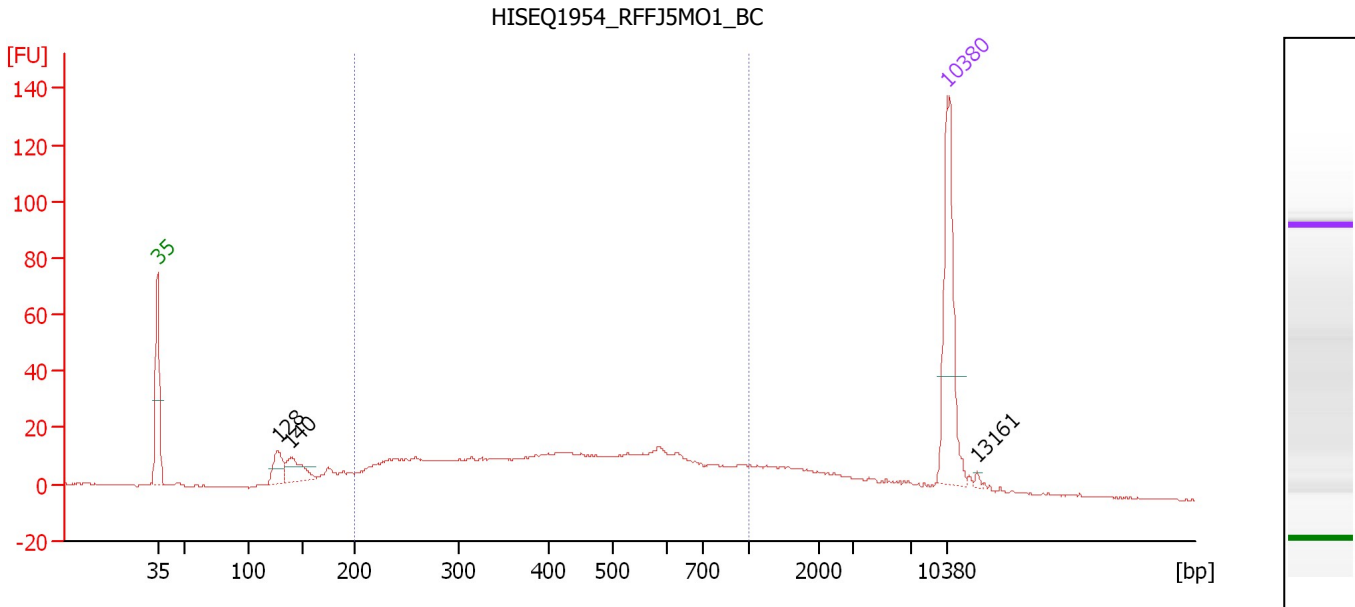
**Region table for sample 4 : HISEQ1966 Bzisolates.10.9.2019 BC**

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	431	2,006.35	1,889.5	7,471.3	99	21.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : HISEQ1954 RFFJ5M01 BC**

Number of peaks found: 3                      Corr. Area 1: 518.4  
 Noise: 0.3

**Peak table for sample 5 : HISEQ1954 RFFJ5M01 BC**

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	128	24.84	294.9		53.65
3	140	30.95	334.6		54.85
4	10,380	75.00	10.9	Upper Marker	113.00
5	13,161	0.00	0.0		115.65

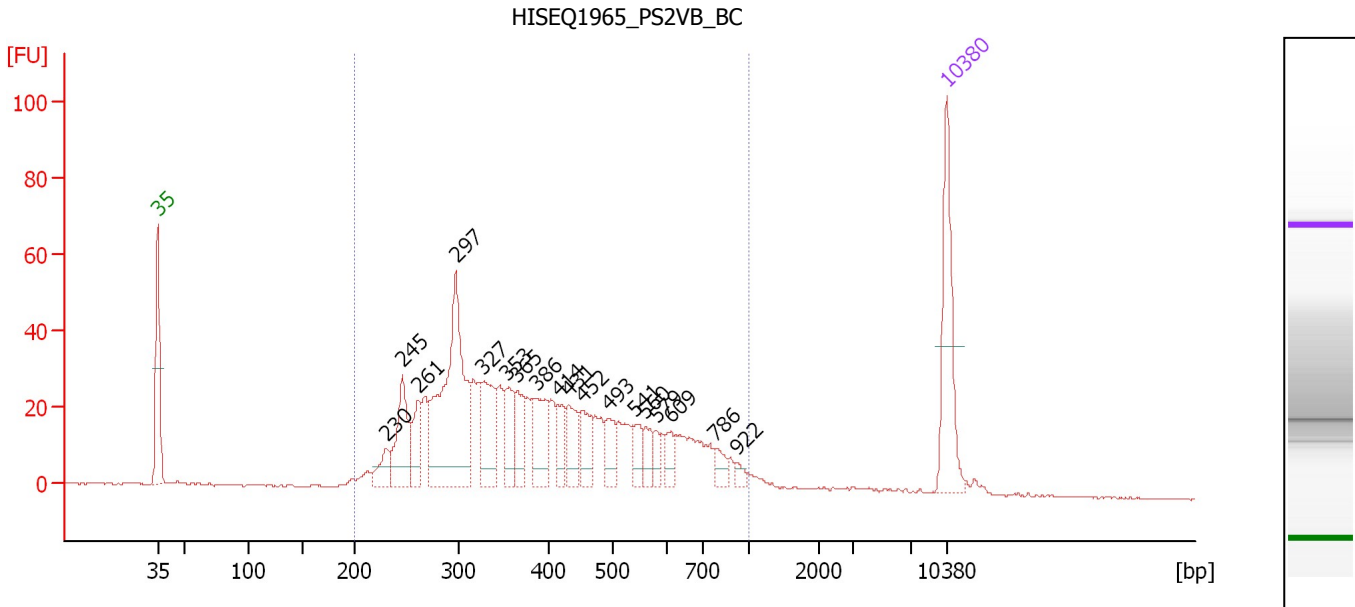
**Region table for sample 5 : HISEQ1954 RFFJ5M01 BC**

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	461	459.28	518.4	1,887.2	68	40.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : HISEQ1965 PS2VB BC**

Number of peaks found: 18                      Corr. Area 1: 877.1  
 Noise: 0.2

**Peak table for sample 6 : HISEQ1965 PS2VB BC**

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	230	25.85	170.1		63.29
3	245	73.89	456.1		64.70
4	261	41.40	240.1		66.16
5	297	233.80	1,193.8		69.45
6	327	68.61	317.9		71.90
7	353	42.82	183.7		73.98
8	365	37.35	155.0		74.92
9	386	53.44	209.8		76.59
10	414	23.04	84.3		78.52
11	431	38.63	135.9		79.46
12	452	32.81	109.9		80.71
13	493	25.92	79.7		83.01
14	541	19.96	55.9		85.41
15	560	17.35	47.0		86.29
16	579	14.04	36.7		87.23
17	609	16.15	40.2		88.48
18	786	12.81	24.7		92.50
19	922	6.39	10.5		94.38
20	10,380	75.00	10.9	Upper Marker	113.00

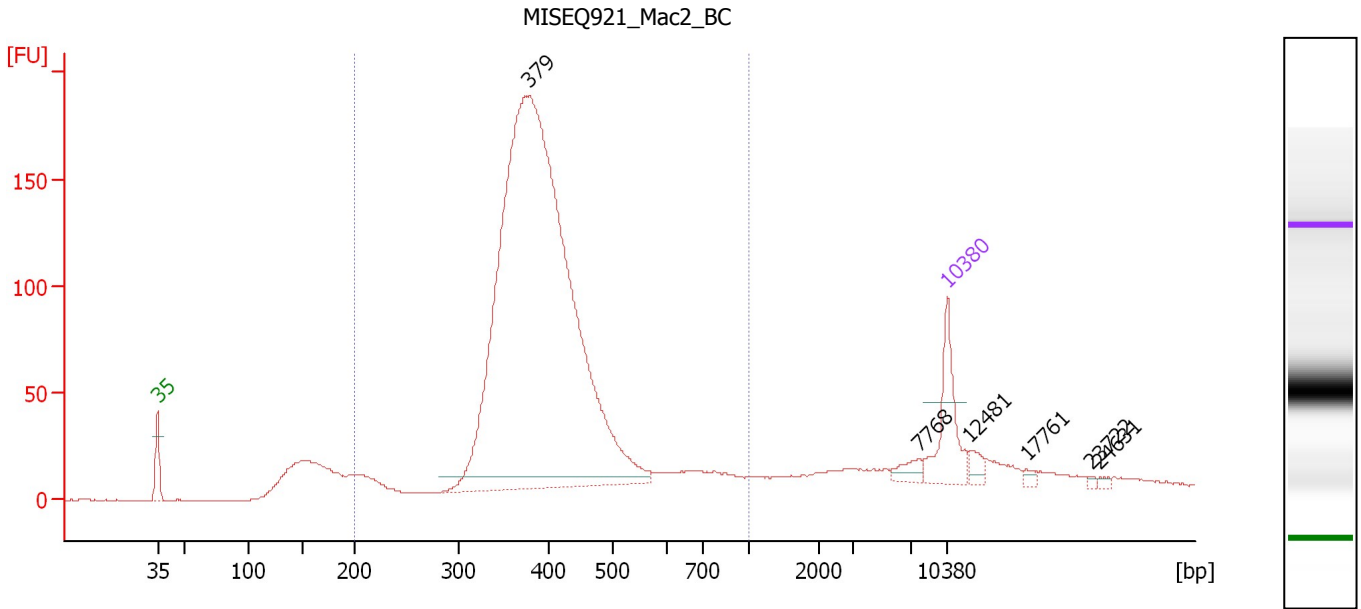
**Region table for sample 6 : HISEQ1965 PS2VB BC**

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	417	1,159.80	877.1	4,951.0	94	37.3

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : MISEQ921 Mac2 BC**

Number of peaks found: 6                      Corr. Area 1: 2,047.0  
 Noise: 0.2

**Peak table for sample 7 : MISEQ921 Mac2 BC**

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	379	1,810.31	7,246.5		76.00
3	7,768	17.22	3.4		110.51
4	10,380	75.00	10.9	Upper Marker	113.00
5	12,481	0.00	0.0		115.00
6	17,761	0.00	0.0		120.03
7	23,722	0.00	0.0		125.71
8	24,631	0.00	0.0		126.58

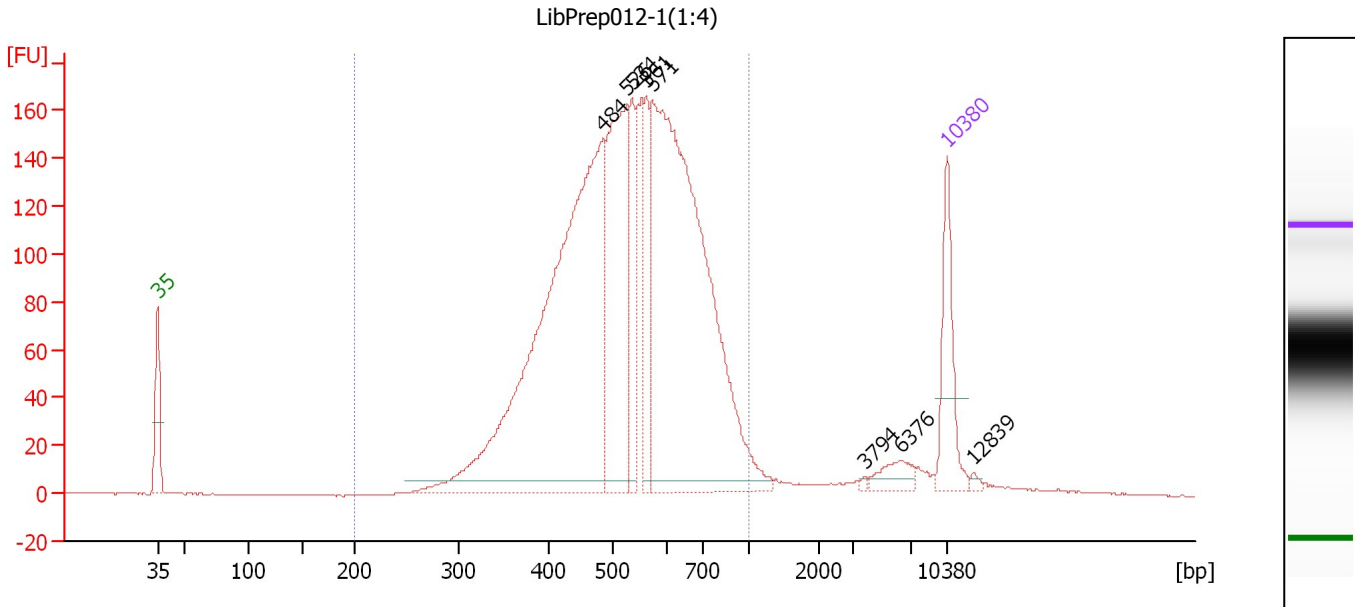
**Region table for sample 7 : MISEQ921 Mac2 BC**

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	412	2,118.98	2,047.0	8,302.9	78	25.3

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : LibPrep012-1(1:4)**

Number of peaks found: 8                      Corr. Area 1: 2,908.7  
 Noise: 0.2

**Peak table for sample 8 : LibPrep012-1(1:4)**

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	484	866.94	2,716.7		82.49
3	526	294.89	849.9		84.67
4	534	106.75	303.0		85.06
5	561	103.24	279.0		86.34
6	571	780.91	2,071.5		86.84
7	3,794	2.22	0.9		105.69
8	6,376	23.71	5.6		108.98
9	10,380	75.00	10.9	Upper Marker	113.00
10	12,839	0.00	0.0		115.34

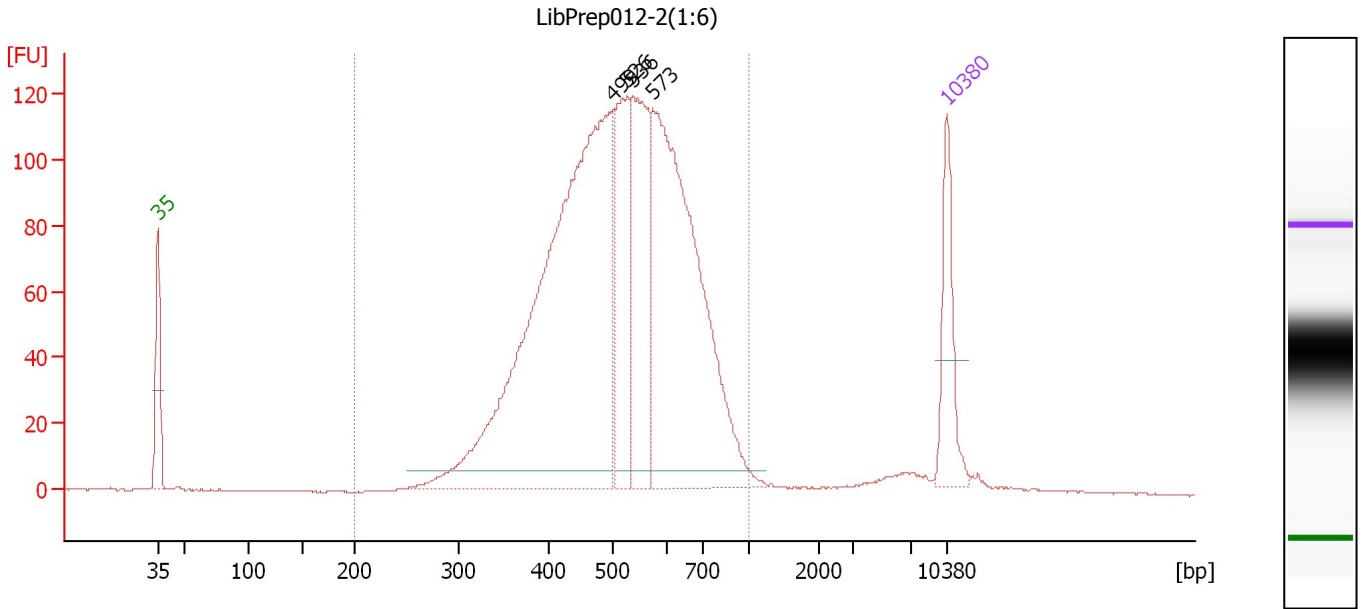
**Region table for sample 8 : LibPrep012-1(1:4)**

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	540	2,302.18	2,908.7	6,975.9	93	24.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : LibPrep012-2(1:6)**

Number of peaks found: 4                      Corr. Area 1: 2,150.2  
 Noise: 0.2

**Peak table for sample 9 : LibPrep012-2(1:6)**

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	499	965.65	2,934.9		83.35
3	526	193.06	556.6		84.66
4	536	218.61	617.6		85.17
5	573	569.92	1,506.6		86.93
6	10,380	75.00	10.9	Upper Marker	113.00

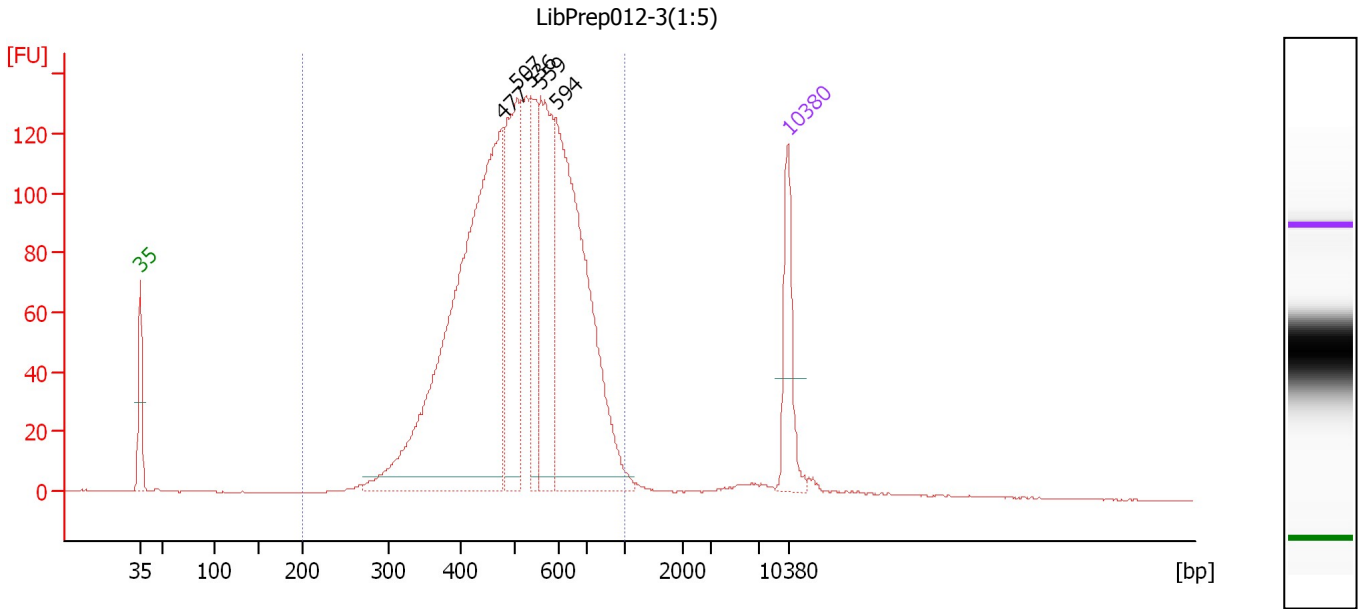
**Region table for sample 9 : LibPrep012-2(1:6)**

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	521	2,085.76	2,150.2	6,530.4	95	23.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : LibPrep012-3(1:5)**

Number of peaks found: 5                      Corr. Area 1: 2,340.4  
 Noise: 0.2

**Peak table for sample 10 : LibPrep012-3(1:5)**

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	477	839.13	2,665.3		82.12
3	507	250.04	746.6		83.79
4	536	127.81	361.1		85.17
5	559	219.20	593.9		86.27
6	594	504.28	1,286.1		87.94
7	10,380	75.00	10.9	Upper Marker	113.00

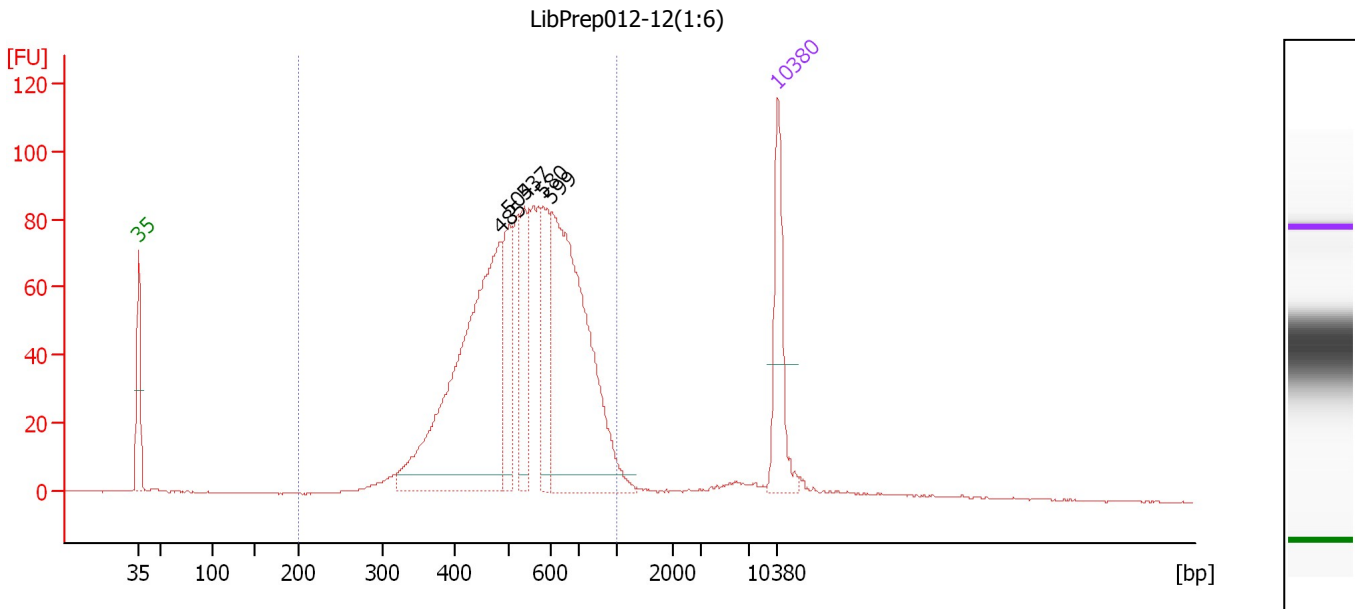
**Region table for sample 10 : LibPrep012-3(1:5)**

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	526	2,183.95	2,340.4	6,758.9	97	23.2

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : LibPrep012-12(1:6)**

Number of peaks found: 5                      Corr. Area 1: 1,441.5  
 Noise: 0.2

**Peak table for sample 11 : LibPrep012-12(1:6)**

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	485	443.36	1,384.7		82.58
3	504	81.84	246.0		83.63
4	537	83.77	236.3		85.21
5	580	79.39	207.5		87.25
6	599	357.73	904.4		88.19
7	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 11 : LibPrep012-12(1:6)**

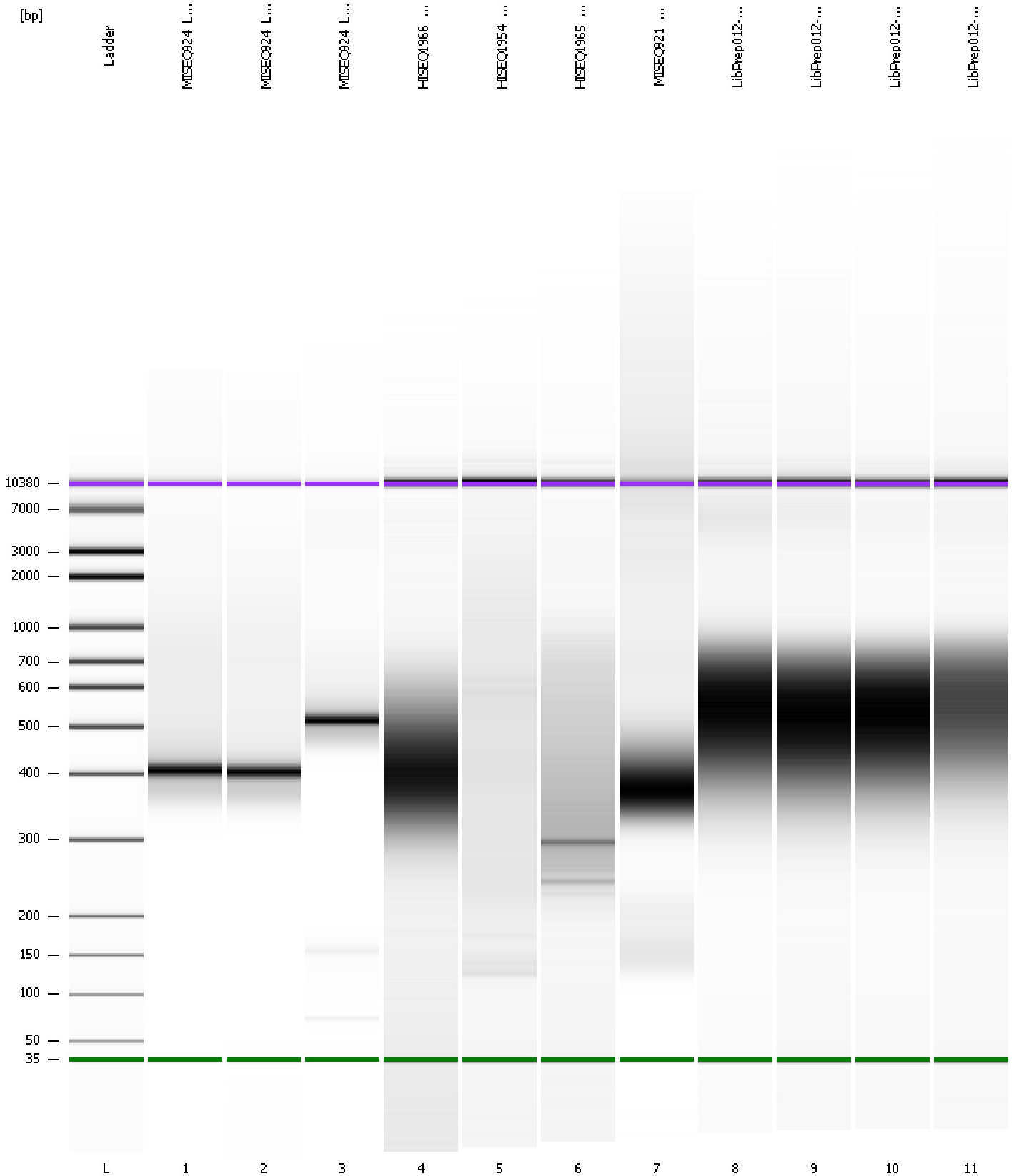
From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	552	1,302.17	1,441.5	3,856.1	94	23.8



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
Modified: 10/15/2019 4:43:55 PM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15\_003.xad

Created: 10/15/2019 4:01:12 PM  
 Modified: 10/15/2019 4:43:55 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		10/15/2019 4:42:30 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\2019-10-15\2019-10-15_003.xad)		Instrument	Run		10/15/2019 4:01:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		10/15/2019 4:01:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		10/15/2019 4:01:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		10/15/2019 4:01:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		10/15/2019 4:01:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		10/15/2019 4:01:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		10/15/2019 4:01:17 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1