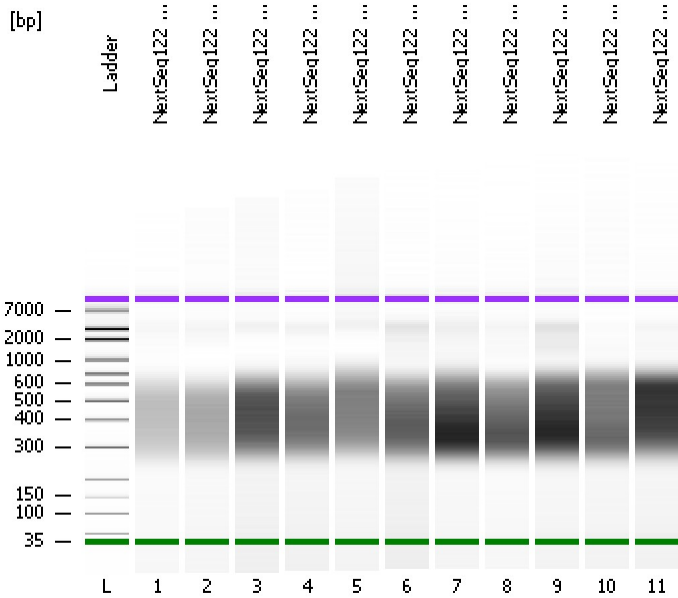


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
Modified: 5/15/2019 6:02:33 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

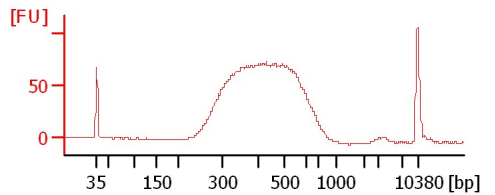
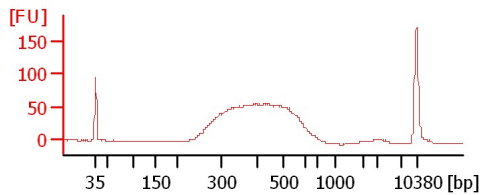
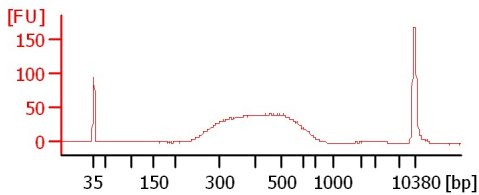
Chip Information:

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

NextSeq122 - 1 size sel lib

NextSeq122 - 2 size sel lib

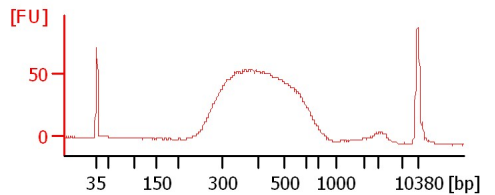
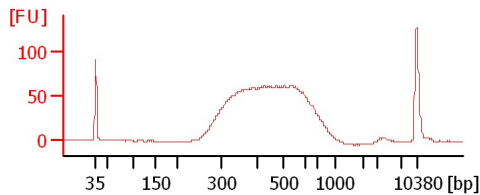
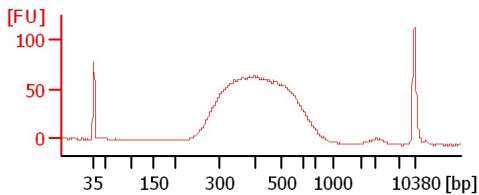
NextSeq122 - 3 size sel lib (1:2)



NextSeq122 - 4 size sel lib (1:2)

NextSeq122 - 5 size sel lib (1:7)

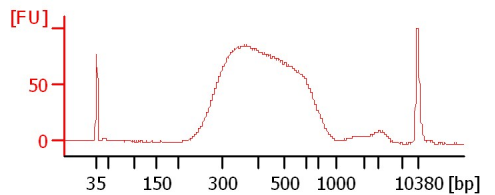
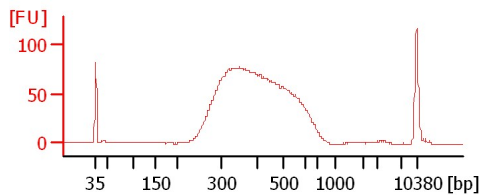
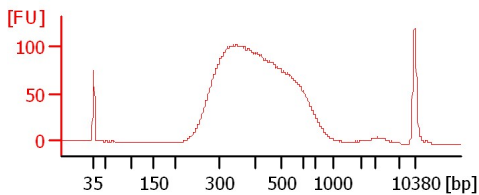
NextSeq122 - 6 size sel lib



NextSeq122 - 7 size sel lib

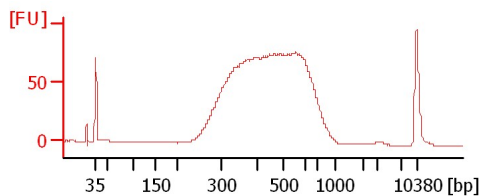
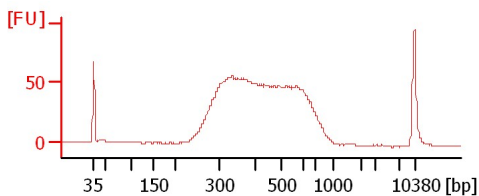
NextSeq122 - 8 size sel lib (1:3)

NextSeq122 - 9 size sel lib



NextSeq122 - 10 size sel lib (1:6)

NextSeq122 - 11 size sel lib



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
NextSeq122 - 1 size sel lib		<input type="checkbox"/>	✓			
NextSeq122 - 2 size sel lib		<input type="checkbox"/>	✓			
NextSeq122 - 3 size sel lib (1:2)		<input type="checkbox"/>	✓			
NextSeq122 - 4 size sel lib (1:2)		<input type="checkbox"/>	✓			
NextSeq122 - 5 size sel lib (1:7)		<input type="checkbox"/>	✓			
NextSeq122 - 6 size sel lib		<input type="checkbox"/>	✓			
NextSeq122 - 7 size sel lib		<input type="checkbox"/>	✓			
NextSeq122 - 8 size sel lib (1:3)		<input type="checkbox"/>	✓			
NextSeq122 - 9 size sel lib		<input type="checkbox"/>	✓			
NextSeq122 - 10 size sel lib (1:6)		<input type="checkbox"/>	✓			
NextSeq122 - 11 size sel lib		<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\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
Modified: 5/15/2019 6:02:33 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

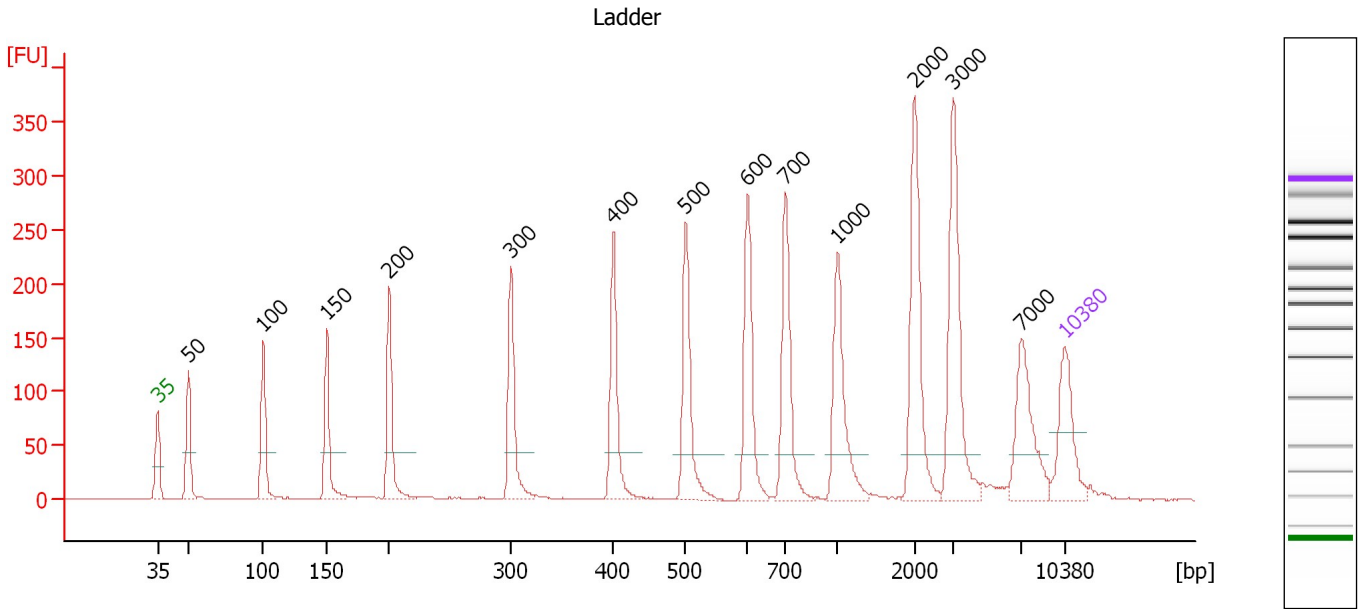
**Ladder**

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\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.5

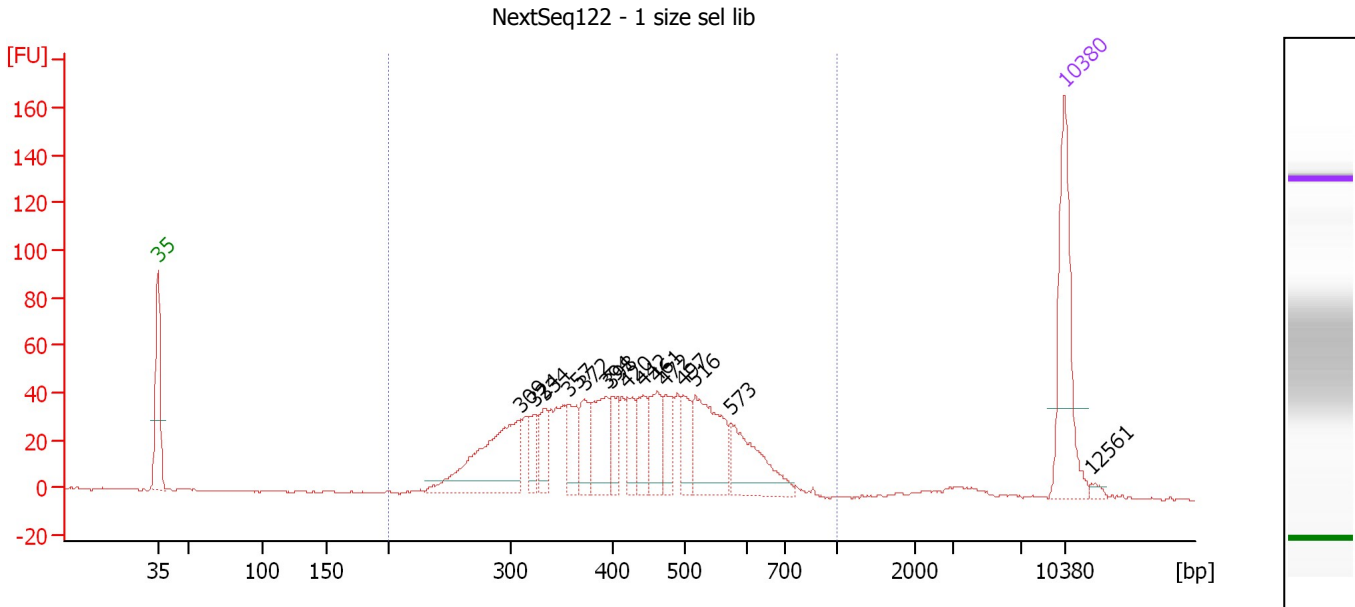
**Peak table for Ladder**

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	51.15
4	150	150.00	1,515.2	Ladder Peak	56.07
5	200	150.00	1,136.4	Ladder Peak	60.90
6	300	150.00	757.6	Ladder Peak	70.26
7	400	150.00	568.2	Ladder Peak	78.17
8	500	150.00	454.5	Ladder Peak	83.76
9	600	150.00	378.8	Ladder Peak	88.54
10	700	150.00	324.7	Ladder Peak	91.48
11	1,000	150.00	227.3	Ladder Peak	95.49
12	2,000	150.00	113.6	Ladder Peak	101.42
13	3,000	150.00	75.8	Ladder Peak	104.41
14	7,000	150.00	32.5	Ladder Peak	109.72
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : NextSeq122 - 1 size sel lib**

Number of peaks found: 15                      Corr. Area 1: 1,060.1  
 Noise: 0.4

**Peak table for sample 1 : NextSeq122 - 1 size sel 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	309	121.70	596.9		70.96
3	324	23.23	108.7		72.14
4	334	24.06	109.2		72.93
5	357	32.94	139.9		74.75
6	372	31.39	128.0		75.93
7	394	54.74	210.8		77.66
8	398	24.30	92.4		78.05
9	420	25.37	91.6		79.28
10	442	32.02	109.8		80.51
11	461	37.59	123.5		81.59
12	472	28.33	91.0		82.18
13	497	27.52	83.8		83.61
14	516	80.61	236.5		84.55
15	573	58.94	155.8		87.25
16	10,380	75.00	10.9	Upper Marker	113.00
17	12,561	0.00	0.0		115.12

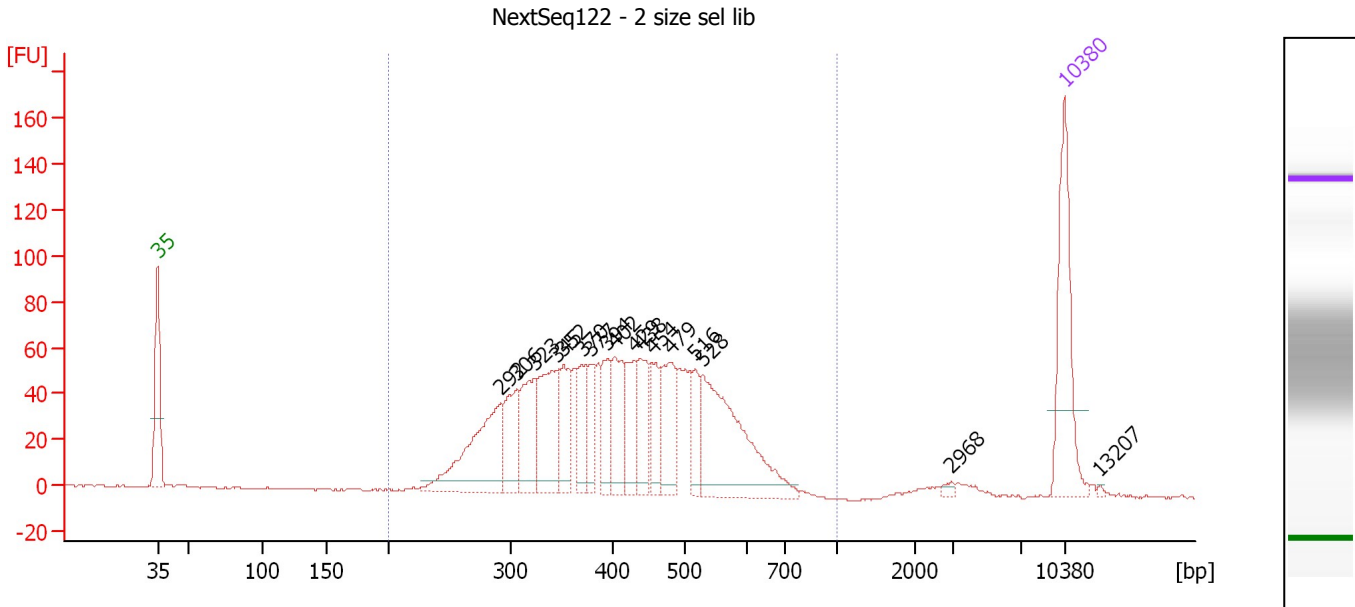
**Region table for sample 1 : NextSeq122 - 1 size sel lib**

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	428	712.68	1,060.1	2,760.5	98	24.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : NextSeq122 - 2 size sel lib**

Number of peaks found: 17                      Corr. Area 1: 1,435.4  
 Noise: 0.4

**Peak table for sample 2 : NextSeq122 - 2 size sel 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	292	121.58	631.2		69.49
3	306	50.86	251.7		70.74
4	323	64.77	304.3		72.04
5	345	87.05	382.1		73.83
6	352	46.00	198.3		74.33
7	370	32.68	133.9		75.78
8	377	32.48	130.4		76.38
9	394	37.11	142.6		77.73
10	402	54.15	204.1		78.27
11	429	40.67	143.8		79.77
12	438	45.52	157.6		80.27
13	454	34.62	115.6		81.17
14	479	56.39	178.2		82.62
15	516	26.81	78.8		84.51
16	528	149.41	428.6		85.11
17	2,968	3.19	1.6		104.32
18	10,380	75.00	10.9	Upper Marker	113.00
19	13,207	0.00	0.0		115.74

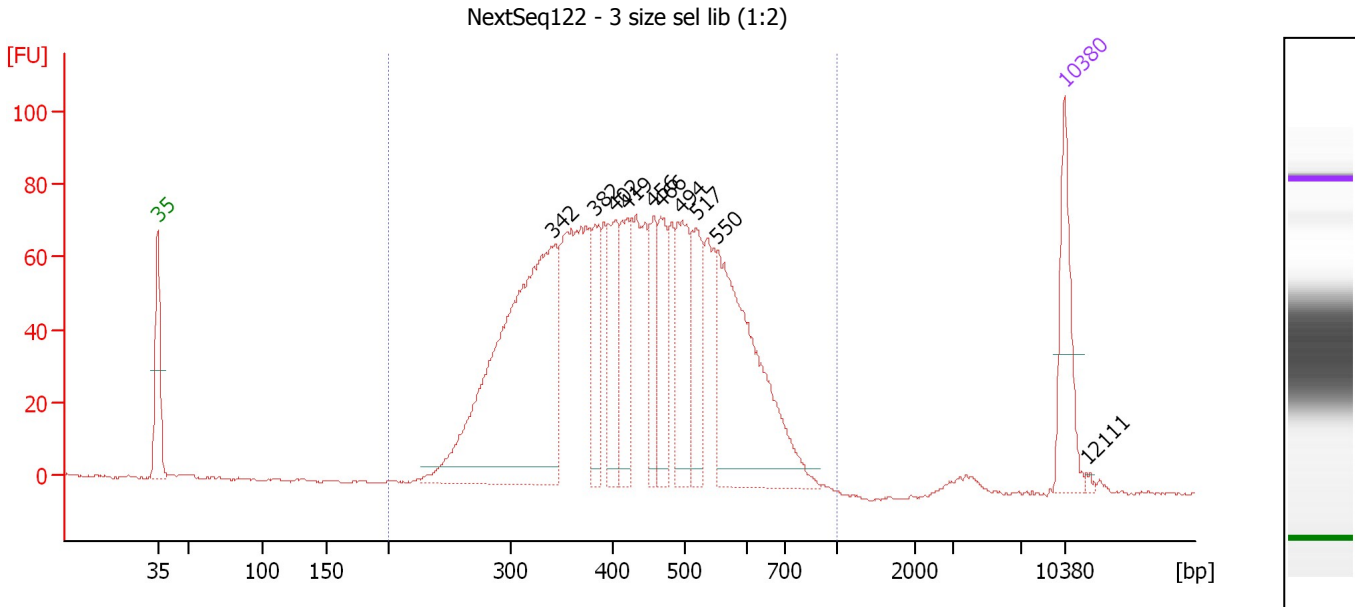
**Region table for sample 2 : NextSeq122 - 2 size sel lib**

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	417	956.38	1,435.4	3,774.7	98	23.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : NextSeq122 - 3 size sel lib (1:2)**

Number of peaks found: 10                      Corr. Area 1: 1,857.4  
 Noise: 0.5

**Peak table for sample 3 : NextSeq122 - 3 size sel lib (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	342	605.84	2,687.5		73.55
3	382	79.39	314.8		76.75
4	402	99.47	374.9		78.28
5	419	104.55	377.8		79.25
6	456	69.48	231.0		81.28
7	466	84.78	275.9		81.84
8	494	122.71	376.5		83.42
9	517	86.62	253.7		84.59
10	550	312.31	859.8		86.17
11	10,380	75.00	10.9	Upper Marker	113.00
12	12,111	0.00	0.0		114.68

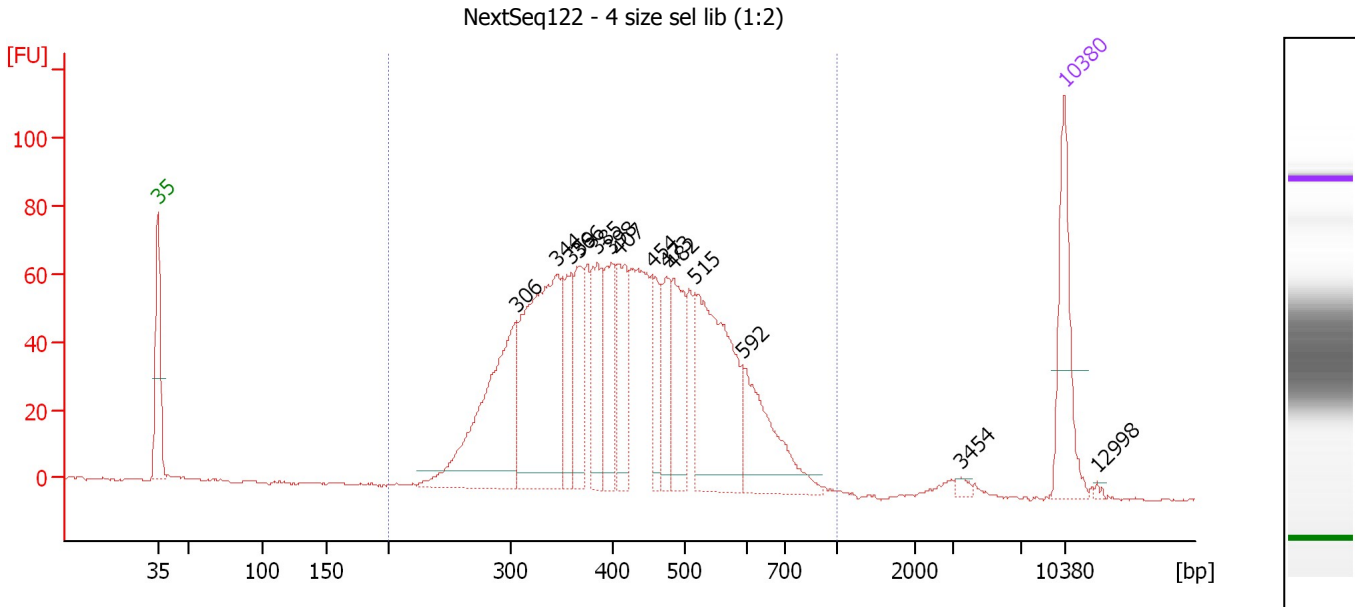
**Region table for sample 3 : NextSeq122 - 3 size sel lib (1:2)**

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	435	2,185.95	1,857.4	8,322.7	99	24.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : NextSeq122 - 4 size sel lib (1:2)**

Number of peaks found: 14                      Corr. Area 1: 1,609.2  
 Noise: 0.3

**Peak table for sample 4 : NextSeq122 - 4 size sel lib (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	306	240.95	1,192.6		70.74
3	344	291.88	1,285.1		73.75
4	359	64.61	272.5		74.94
5	366	84.51	349.5		75.51
6	385	77.50	305.2		76.96
7	398	84.76	322.8		78.00
8	407	79.55	296.0		78.57
9	454	52.13	173.8		81.21
10	473	51.40	164.7		82.25
11	482	87.45	274.8		82.77
12	515	213.37	627.7		84.48
13	592	107.33	274.7		88.16
14	3,454	4.67	2.0		105.01
15	10,380	75.00	10.9	Upper Marker	113.00
16	12,998	0.00	0.0		115.54

**Region table for sample 4 : NextSeq122 - 4 size sel lib (1:2)**

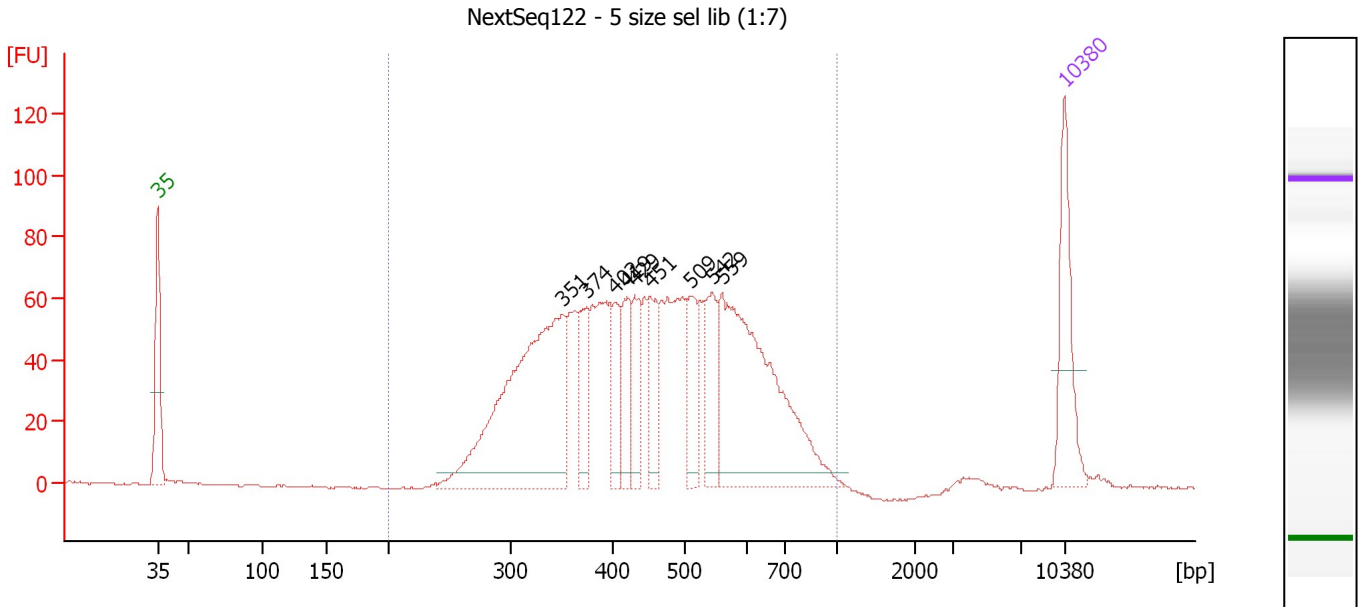
From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	428	1,646.61	1,609.2	6,342.2	99	24.9



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : NextSeq122 - 5 size sel lib (1:7)**

Number of peaks found: 9                      Corr. Area 1: 1,638.6  
 Noise: 0.4

**Peak table for sample 5 : NextSeq122 - 5 size sel lib (1:7)**

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	351	390.75	1,685.1		74.32
3	374	59.17	239.5		76.13
4	402	53.67	202.4		78.27
5	419	57.71	208.7		79.23
6	429	53.72	189.5		79.81
7	451	56.61	190.0		81.04
8	509	61.39	182.8		84.19
9	542	69.08	193.0		85.79
10	559	317.01	859.0		86.59
11	10,380	75.00	10.9	Upper Marker	113.00

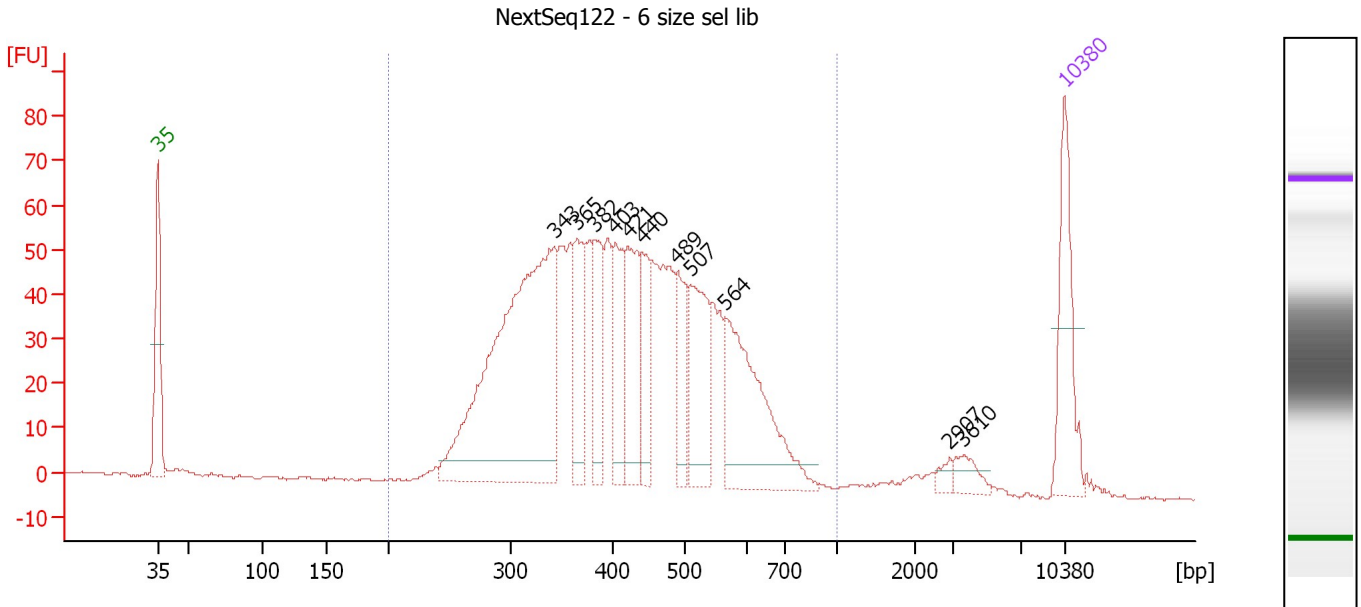
**Region table for sample 5 : NextSeq122 - 5 size sel lib (1:7)**

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	465	1,541.06	1,638.6	5,574.2	98	27.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : NextSeq122 - 6 size sel lib**

Number of peaks found: 11                      Corr. Area 1: 1,342.3  
 Noise: 0.3

**Peak table for sample 6 : NextSeq122 - 6 size sel 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	343	494.09	2,185.6		73.62
3	365	84.79	352.4		75.36
4	382	67.29	267.1		76.72
5	403	72.28	271.5		78.35
6	421	105.52	379.9		79.33
7	440	55.24	190.1		80.42
8	489	59.12	183.3		83.14
9	507	98.98	295.6		84.12
10	564	170.46	457.6		86.84
11	2,907	7.67	4.0		104.13
12	3,610	15.18	6.4		105.22
13	10,380	75.00	10.9	Upper Marker	113.00

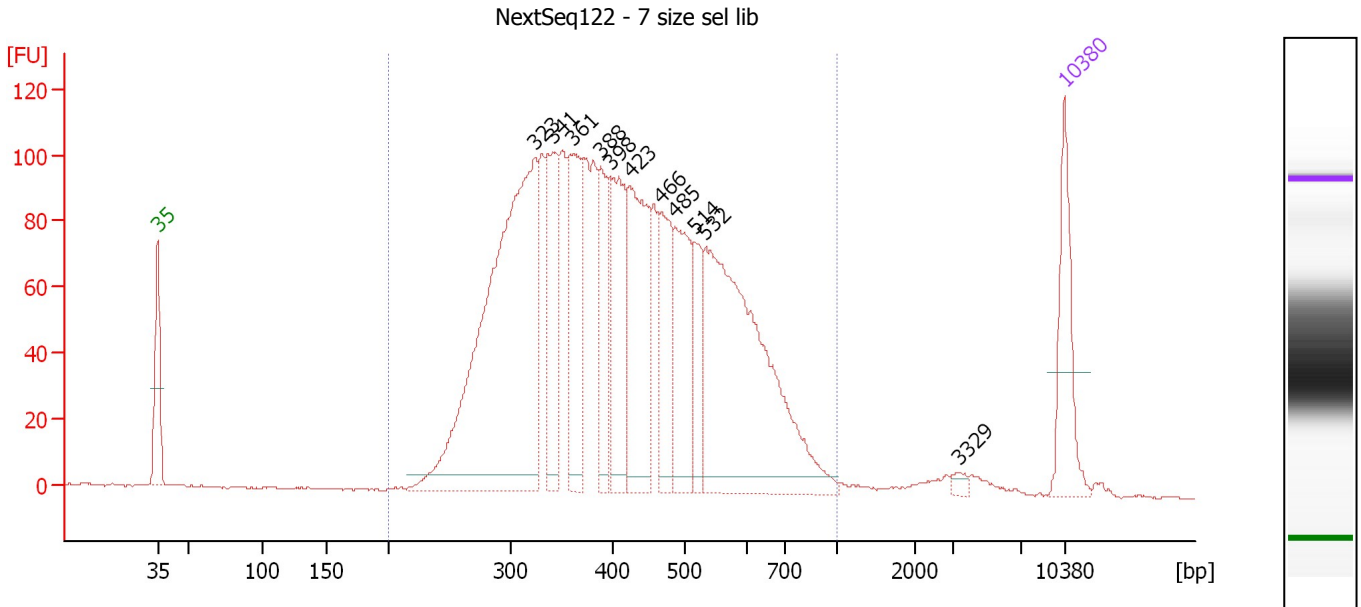
**Region table for sample 6 : NextSeq122 - 6 size sel lib**

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	424	1,672.76	1,342.3	6,523.2	97	25.2

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : NextSeq122 - 7 size sel lib**

Number of peaks found: 11                      Corr. Area 1: 2,595.5  
 Noise: 0.3

**Peak table for sample 7 : NextSeq122 - 7 size sel 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	323	666.11	3,124.3		72.08
3	341	135.56	602.1		73.51
4	361	143.96	603.7		75.11
5	388	95.23	372.1		77.20
6	398	148.86	567.4		77.97
7	423	191.46	685.7		79.46
8	466	107.97	350.8		81.88
9	485	135.84	424.3		82.93
10	514	65.32	192.7		84.42
11	532	444.13	1,264.7		85.30
12	3,329	6.03	2.7		104.85
13	10,380	75.00	10.9	Upper Marker	113.00

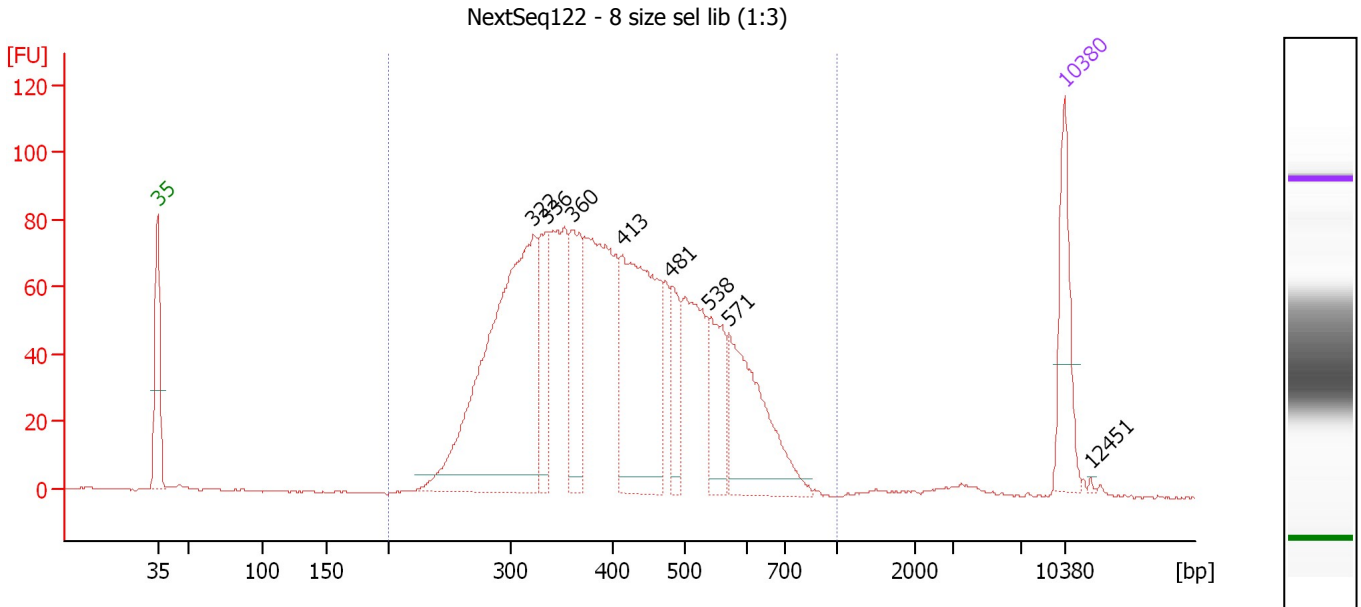
**Region table for sample 7 : NextSeq122 - 7 size sel lib**

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	426	2,590.08	2,595.5	10,205.4	98	28.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : NextSeq122 - 8 size sel lib (1:3)**

Number of peaks found: 8                      Corr. Area 1: 1,886.6  
 Noise: 0.3

**Peak table for sample 8 : NextSeq122 - 8 size sel lib (1:3)**

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	322	563.22	2,647.1		72.03
3	336	103.98	469.2		73.09
4	360	131.83	555.2		74.99
5	413	305.53	1,121.0		78.89
6	481	61.71	194.5		82.69
7	538	90.16	253.8		85.59
8	571	176.76	469.1		87.15
9	10,380	75.00	10.9	Upper Marker	113.00
10	12,451	0.00	0.0		115.01

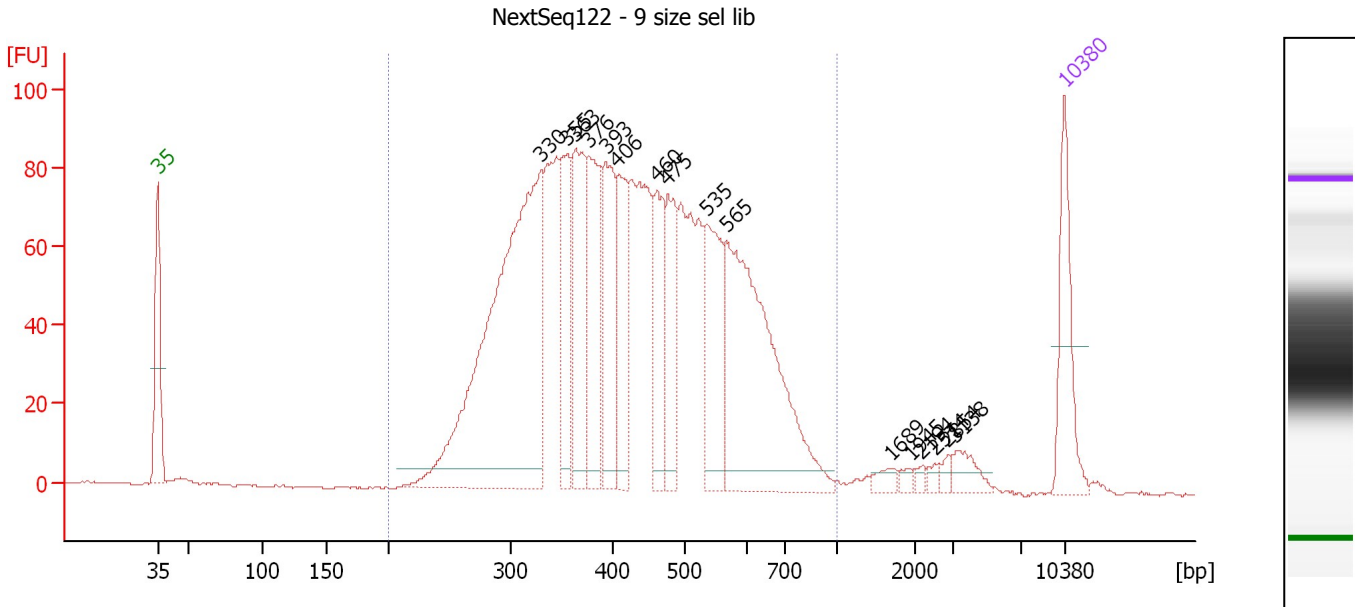
**Region table for sample 8 : NextSeq122 - 8 size sel lib (1:3)**

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	416	2,150.08	1,886.6	8,576.0	98	25.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : NextSeq122 - 9 size sel lib**

Number of peaks found: 16                      Corr. Area 1: 2,172.5  
 Noise: 0.3

**Peak table for sample 9 : NextSeq122 - 9 size sel 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	330	659.87	3,031.7		72.61
3	355	120.30	512.8		74.64
4	363	159.08	663.5		75.26
5	376	136.02	547.9		76.28
6	393	142.78	550.1		77.63
7	406	108.65	405.9		78.48
8	460	92.32	304.1		81.53
9	475	95.65	305.0		82.37
10	535	138.91	393.7		85.42
11	565	351.57	942.3		86.88
12	1,689	10.22	9.2		99.58
13	1,945	5.89	4.6		101.10
14	2,194	4.41	3.0		102.00
15	2,514	5.37	3.2		102.96
16	2,854	6.17	3.3		103.98
17	3,138	19.81	9.6		104.60
18	10,380	75.00	10.9	Upper Marker	113.00

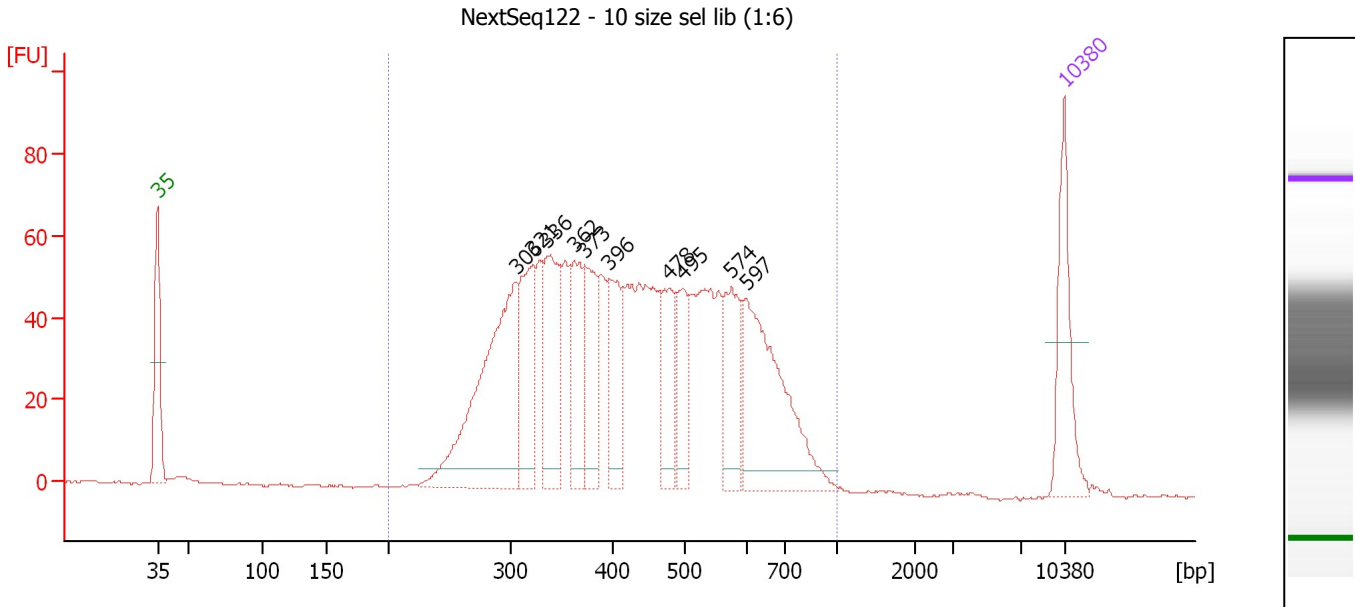
**Region table for sample 9 : NextSeq122 - 9 size sel lib**

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	435	2,673.28	2,172.5	10,315.3	96	28.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : NextSeq122 - 10 size sel lib (1:6)**

Number of peaks found: 10                      Corr. Area 1: 1,520.9  
 Noise: 0.4

**Peak table for sample 10 : NextSeq122 - 10 size sel lib (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	306	310.61	1,540.4		70.69
3	321	119.41	563.2		71.94
4	336	133.11	601.0		73.07
5	362	97.52	408.1		75.17
6	373	87.88	357.1		76.02
7	396	83.32	318.5		77.89
8	478	75.24	238.5		82.53
9	495	66.23	202.7		83.49
10	574	89.13	235.4		87.29
11	597	204.08	517.5		88.42
12	10,380	75.00	10.9	Upper Marker	113.00

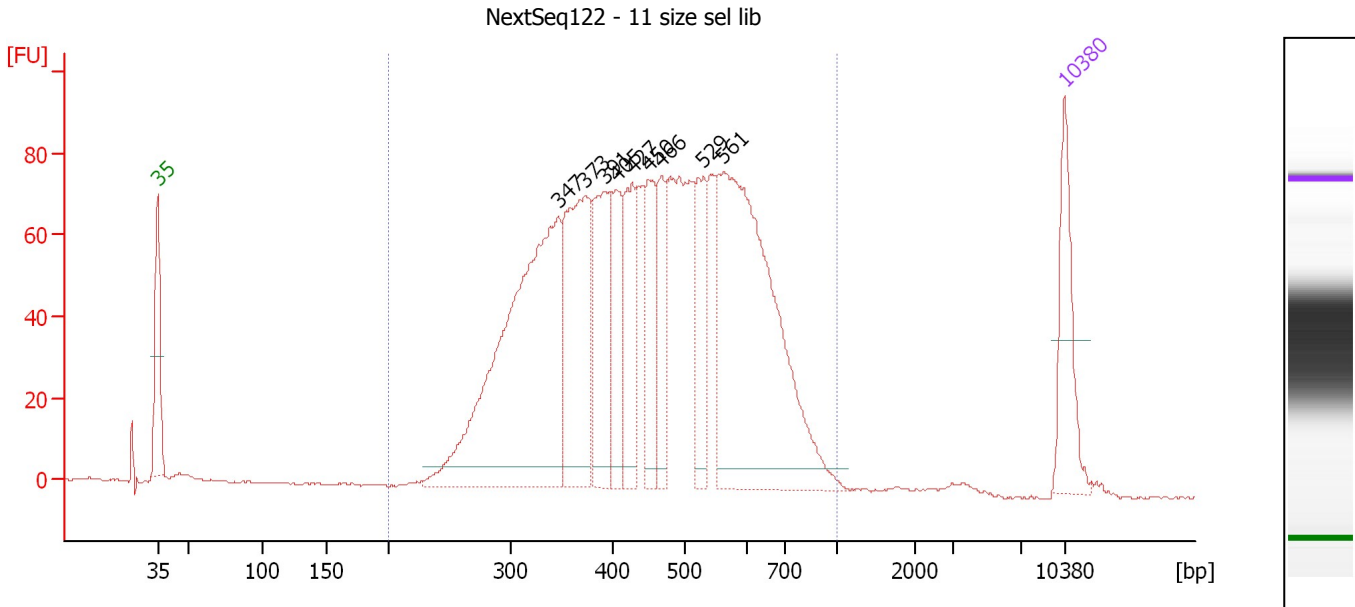
**Region table for sample 10 : NextSeq122 - 10 size sel lib (1:6)**

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	443	1,872.70	1,520.9	7,173.8	99	29.3

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : NextSeq122 - 11 size sel lib**

Number of peaks found: 9                      Corr. Area 1: 1,984.3  
 Noise: 0.3

**Peak table for sample 11 : NextSeq122 - 11 size sel 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	347	551.58	2,411.6		73.94
3	373	228.98	931.2		76.00
4	391	138.42	535.9		77.49
5	405	89.14	333.4		78.46
6	427	112.14	398.3		79.66
7	450	94.91	319.5		80.97
8	466	76.82	249.5		81.89
9	529	88.60	253.8		85.15
10	561	490.07	1,322.9		86.69
11	10,380	75.00	10.9	Upper Marker	113.00

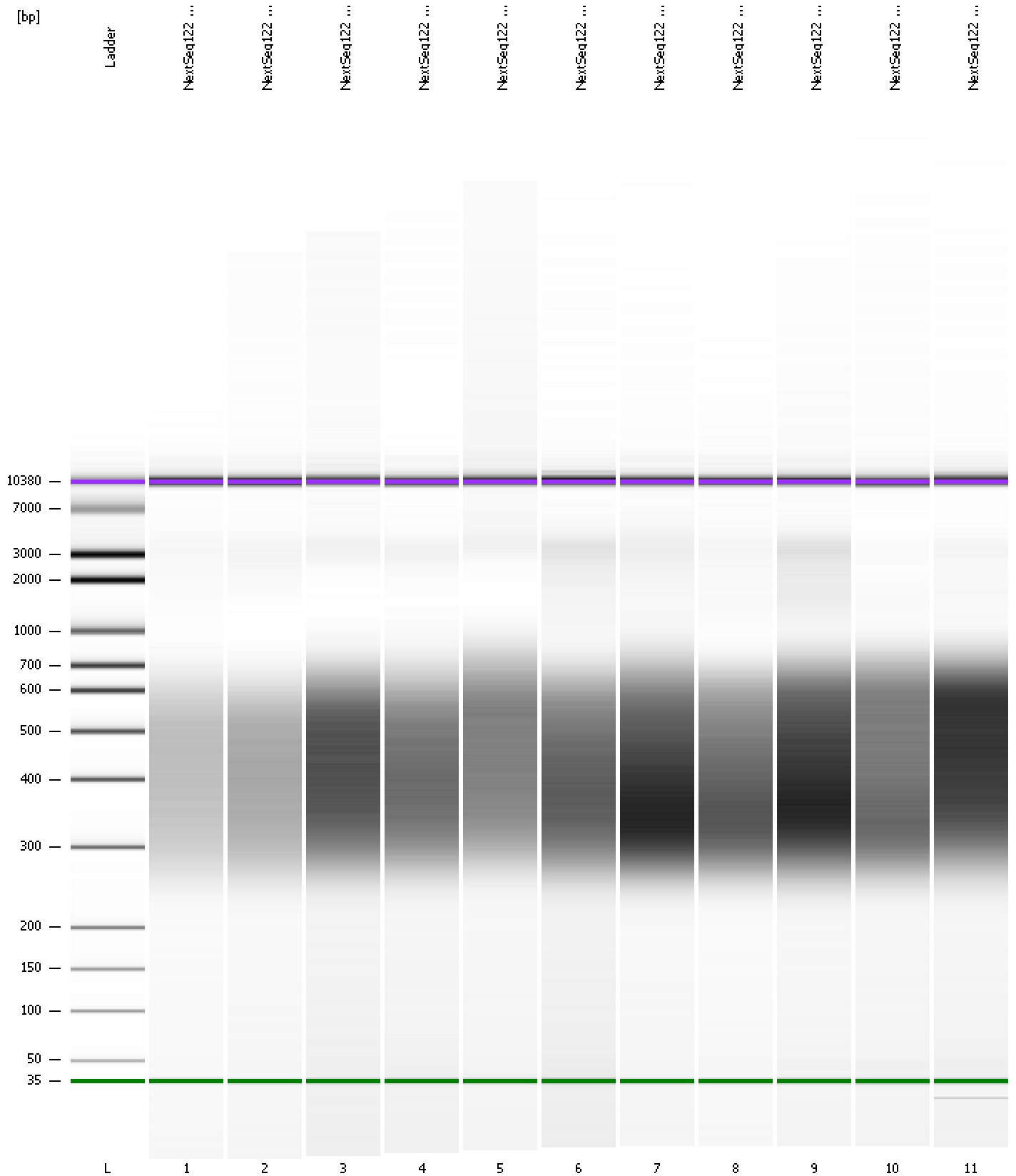
**Region table for sample 11 : NextSeq122 - 11 size sel lib**

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	462	2,289.39	1,984.3	8,329.0	98	27.4

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
Modified: 5/15/2019 6:02:33 PM

**Gel Image**





Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2019-05-15\2019-05-15\_008.xad

Created: 5/15/2019 5:22:09 PM  
 Modified: 5/15/2019 6:02:33 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		5/15/2019 6:02:32 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\2019-05-15\2019-05-15_008.xad)		Instrument	Run		5/15/2019 5:22:10 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		5/15/2019 5:22:10 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		5/15/2019 5:22:10 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		5/15/2019 5:22:10 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		5/15/2019 5:22:10 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		5/15/2019 5:22:10 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		5/15/2019 5:22:10 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1