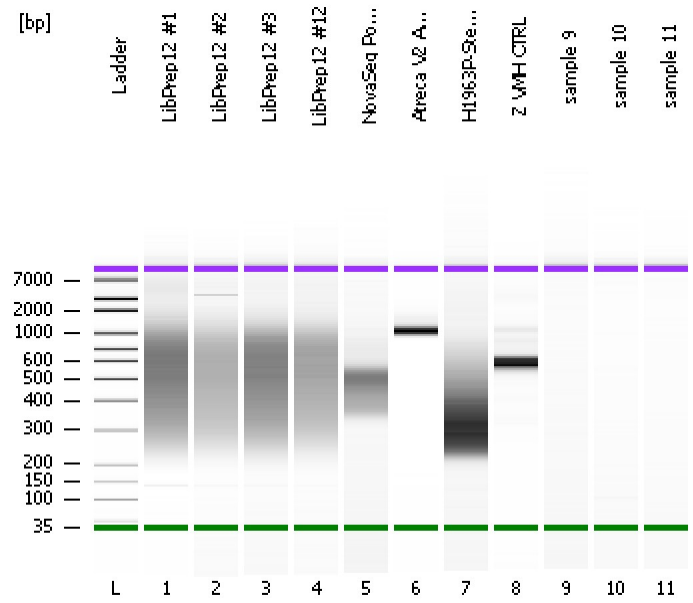
Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 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

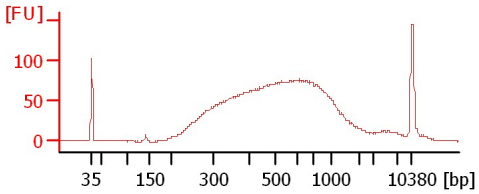
Chip Information:

Chip Lot #:

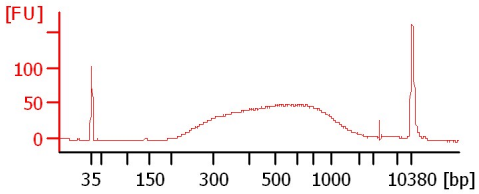
Reagent Kit Lot #:

Chip Comments:

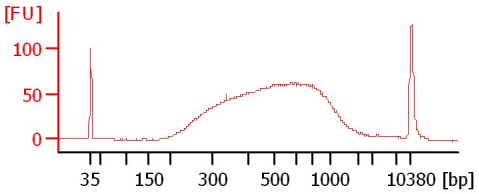
LibPrep12 #1



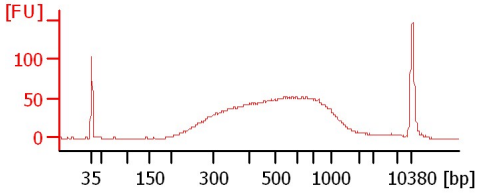
LibPrep12 #2



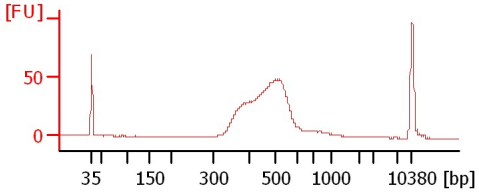
LibPrep12 #3



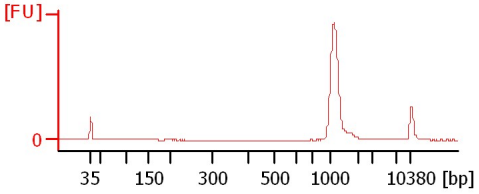
LibPrep12 #12



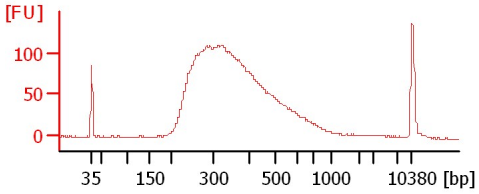
NovaSeq Pool 10-8-19



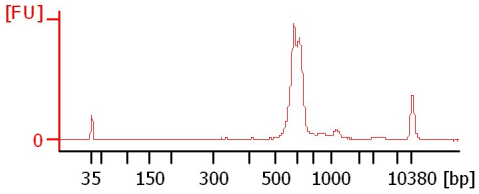
Atreca V2 Amplicon_0.8X clean-up



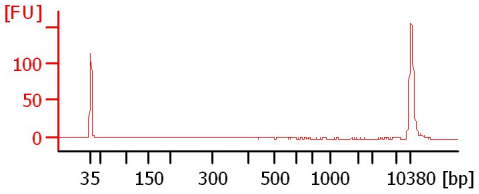
H1963P-Steven 1:10



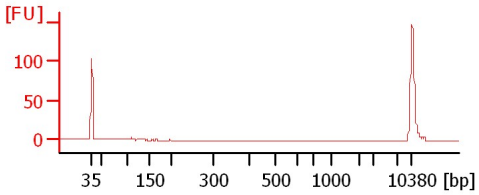
Z VMH CTRL



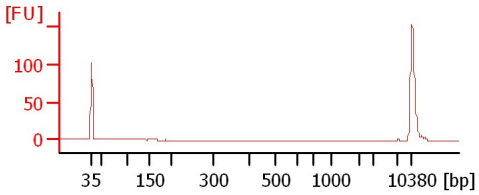
sample 9



sample 10



sample 11



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
LibPrep12 #1		☐	✔			
LibPrep12 #2		☐	✔			
LibPrep12 #3		☐	✔			
LibPrep12 #12		☐	✔			
NovaSeq Pool 10-8-19		☐	✔			
Atreca V2 Amplicon_0.8X clean-up		☐	✔			
H1963P-Steven 1:10		☐	✔			
Z VMH CTRL		☐	✔			
sample 9		☐	✔			
sample 10		☐	✔			
sample 11		☐	✔			
Ladder		☐	✔			

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_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 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

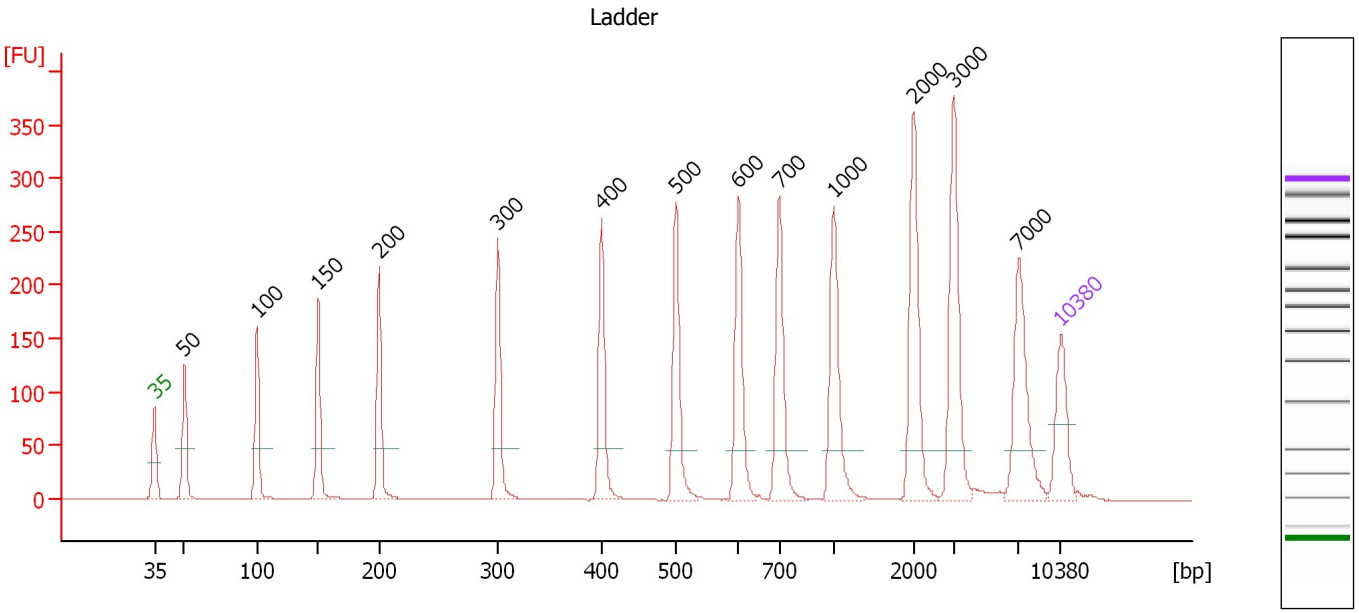
Ladder

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_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

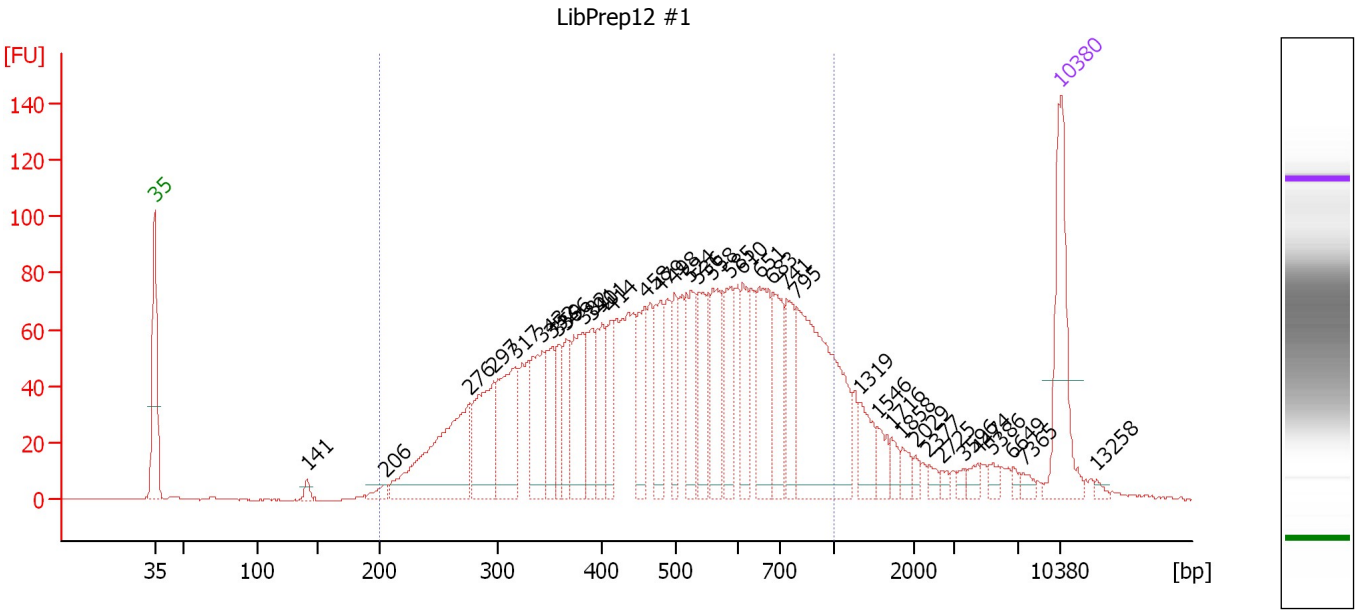
Peak table for Ladder

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	45.30
3		100	150.00	2,272.7	Ladder Peak	50.90
4		150	150.00	1,515.2	Ladder Peak	55.62
5		200	150.00	1,136.4	Ladder Peak	60.35
6		300	150.00	757.6	Ladder Peak	69.53
7		400	150.00	568.2	Ladder Peak	77.51
8		500	150.00	454.5	Ladder Peak	83.25
9		600	150.00	378.8	Ladder Peak	88.11
10		700	150.00	324.7	Ladder Peak	91.24
11		1,000	150.00	227.3	Ladder Peak	95.43
12		2,000	150.00	113.6	Ladder Peak	101.66
13		3,000	150.00	75.8	Ladder Peak	104.70
14		7,000	150.00	32.5	Ladder Peak	109.73
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_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : LibPrep12 #1

Number of peaks found:	38	Corr. Area 1:	2,396.5
Noise:	0.2		

Peak table for sample 1 : LibPrep12 #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	141	8.10	87.0		54.78
3	206	10.22	75.2		60.88
4	276	163.93	898.5		67.36
5	297	92.33	471.6		69.22
6	317	96.39	461.3		70.85
7	343	69.26	305.8		72.97
8	352	46.24	198.8		73.72
9	359	35.55	150.2		74.20
10	366	38.69	160.1		74.82
11	382	71.75	284.3		76.10
12	391	47.53	184.4		76.76
13	401	50.23	189.9		77.56
14	414	36.83	134.8		78.31
15	458	52.58	173.8		80.87
16	479	53.74	169.9		82.06
17	498	39.36	119.6		83.16
18	524	47.76	138.2		84.40
19	536	55.46	156.7		85.02
20	558	65.69	178.3		86.08
21	585	49.86	129.0		87.40
22	610	46.36	115.2		88.42
23	651	72.91	169.8		89.70
24	683	57.74	128.1		90.71
25	741	44.90	91.8		91.81

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...

... Peak table for sample 1 : LibPrep12 #1

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	795	186.71	356.0		92.56
27	1,319	26.96	31.0		97.42
28	1,546	14.22	13.9		98.83
29	1,716	9.14	8.1		99.89
30	1,858	8.86	7.2		100.77
31	2,029	4.86	3.6		101.75
32	2,377	6.27	4.0		102.80
33	2,725	4.54	2.5		103.86
34	3,596	4.11	1.7		105.45
35	4,474	7.25	2.5		106.56
36	5,386	5.77	1.6		107.70
37	6,649	3.59	0.8		109.29
38	7,365	5.60	1.2		110.09
39	10,380	75.00	10.9	Upper Marker	113.00
40	13,258	0.00	0.0		115.78

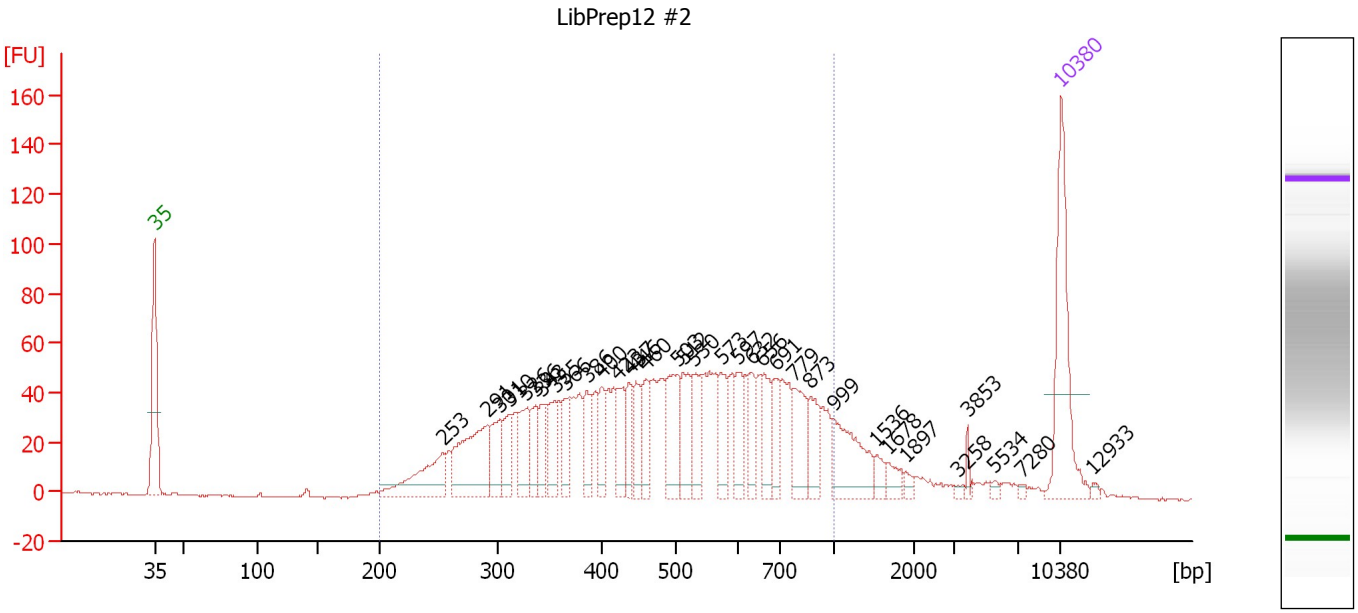
Region table for sample 1 : LibPrep12 #1

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	508	1,863.16	2,396.5	6,683.5	88	35.6

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : LibPrep12 #2

Number of peaks found:	34	Corr. Area 1:	1,619.7
Noise:	0.3		

Peak table for sample 2 : LibPrep12 #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	253	63.45	379.4		65.25
3	291	93.44	486.0		68.73
4	301	32.93	165.7		69.61
5	310	27.65	135.0		70.36
6	326	35.86	166.5		71.63
7	336	25.31	114.0		72.43
8	343	23.67	104.6		72.96
9	355	34.80	148.5		73.93
10	366	25.88	107.1		74.81
11	386	24.06	94.5		76.39
12	400	24.39	92.4		77.49
13	423	27.87	99.9		78.81
14	437	24.53	85.0		79.65
15	446	26.51	90.0		80.18
16	460	24.99	82.2		80.97
17	503	52.42	157.9		83.40
18	512	36.69	108.6		83.84
19	530	33.84	96.7		84.72
20	573	28.22	74.6		86.79
21	597	27.44	69.6		87.98
22	632	25.55	61.2		89.12
23	656	30.18	69.7		89.87
24	691	23.03	50.5		90.97
25	779	37.75	73.5		92.34

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...

... Peak table for sample 2 : LibPrep12 #2

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	873	26.67	46.3		93.66
27	999	50.12	76.0		95.42
28	1,536	8.41	8.3		98.77
29	1,678	9.21	8.3		99.65
30	1,897	4.06	3.2		101.02
31	3,258	1.99	0.9		105.03
32	3,853	3.63	1.4		105.78
33	5,534	2.70	0.7		107.89
34	7,280	1.74	0.4		110.00
35	10,380	75.00	10.9	Upper Marker	113.00
36	12,933	0.00	0.0		115.47

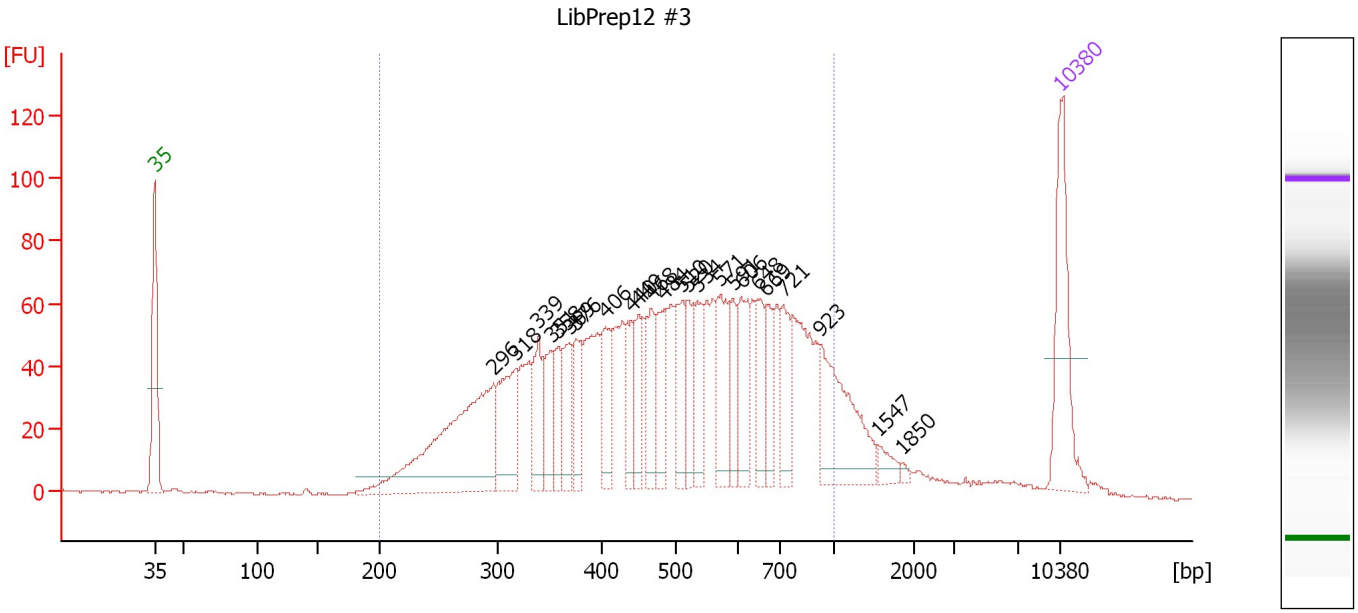
Region table for sample 2 : LibPrep12 #2

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Color % of Total	Size distribution in CV [%]
200	1,000	503	1,193.83	1,619.7	4,306.0	89	35.5

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : LibPrep12 #3

Number of peaks found:	24	Corr. Area 1:	2,001.3
Noise:	0.3		

Peak table for sample 3 : LibPrep12 #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	296	235.59	1,205.4		69.17
3	318	87.18	415.4		70.96
4	339	48.54	217.1		72.62
5	351	44.46	191.9		73.61
6	358	32.64	138.2		74.15
7	369	45.31	186.0		75.04
8	376	40.61	163.7		75.58
9	406	48.87	182.3		77.87
10	440	39.20	134.9		79.84
11	449	37.35	126.0		80.33
12	468	45.67	147.9		81.41
13	484	45.59	142.6		82.35
14	510	48.87	145.2		83.74
15	520	37.68	109.8		84.23
16	534	46.99	133.3		84.90
17	571	60.60	160.8		86.69
18	591	31.44	80.6		87.68
19	606	50.24	125.5		88.31
20	648	42.99	100.5		89.61
21	669	30.31	68.6		90.28
22	721	42.71	89.8		91.53
23	923	99.50	163.3		94.36
24	1,547	11.57	11.3		98.84
25	1,850	2.42	2.0		100.72

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...

... Peak table for sample 3 : LibPrep12 #3

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

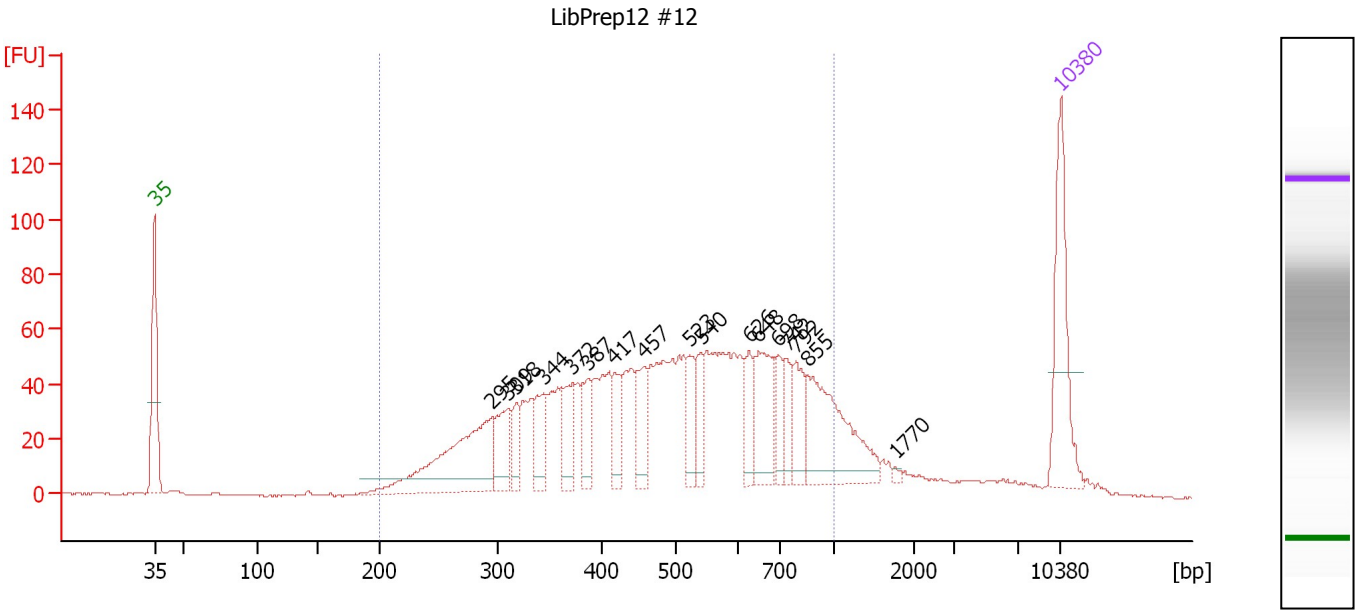
Region table for sample 3 : LibPrep12 #3

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Color % of Total	Size distribution in CV [%]
200	1,000	506	1,715.28	2,001.3	6,173.3	91	35.6

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : LibPrep12 #12

Number of peaks found:	17	Corr. Area 1:	1,652.9
Noise:	0.3		

Peak table for sample 4 : LibPrep12 #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	295	160.84	825.1		69.10
3	309	48.89	240.1		70.21
4	318	25.86	123.3		70.94
5	344	40.63	178.8		73.07
6	372	41.88	170.5		75.28
7	387	32.22	126.2		76.48
8	417	30.52	110.8		78.51
9	457	39.08	129.6		80.77
10	523	29.32	85.0		84.36
11	540	26.97	75.7		85.19
12	626	28.16	68.1		88.93
13	648	61.96	144.8		89.62
14	698	23.95	52.0		91.19
15	742	26.18	53.4		91.83
16	792	35.02	67.0		92.53
17	855	111.35	197.4		93.40
18	1,770	2.31	2.0		100.23
19	10,380	75.00	10.9	Upper Marker	113.00

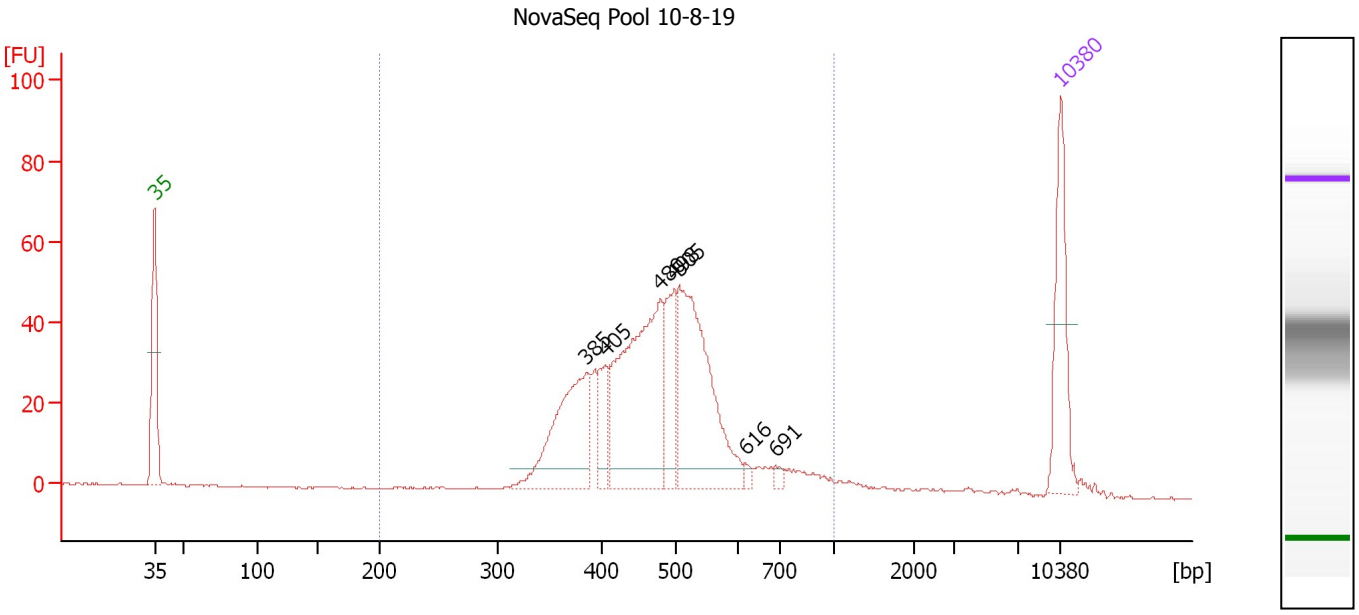
Region table for sample 4 : LibPrep12 #12

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	509	1,345.13	1,652.9	4,799.9	89	35.4

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : NovaSeq Pool 10-8-19

Number of peaks found:	7	Corr. Area 1:	677.0
Noise:	0.2		

Peak table for sample 5 : NovaSeq Pool 10-8-19

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	385	150.27	591.8		76.29
3	405	42.10	157.5		77.81
4	480	246.09	777.5		82.08
5	498	66.26	201.7		83.12
6	505	221.81	665.3		83.50
7	616	5.21	12.8		88.62
8	691	5.80	12.7		90.95
9	10,380	75.00	10.9	Upper Marker	113.00

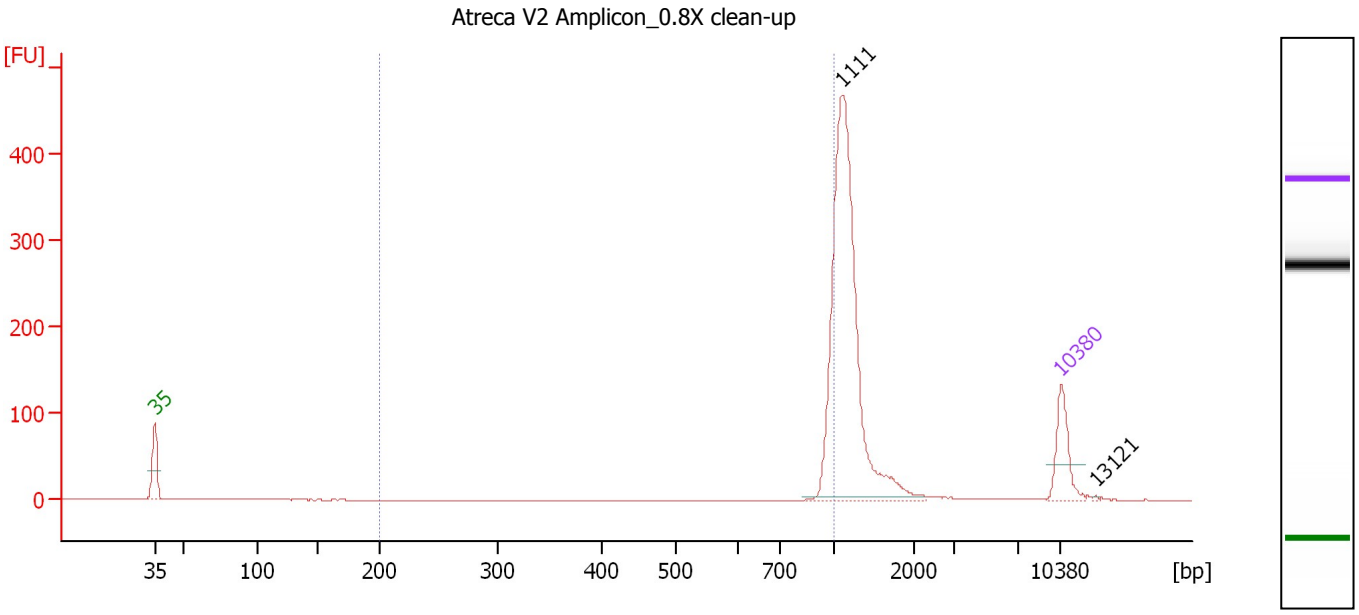
Region table for sample 5 : NovaSeq Pool 10-8-19

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	481	841.27	677.0	2,789.8	95	21.7

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : Atreca V2 Amplicon_0.8X clean-up

Number of peaks found:	2	Corr. Area 1:	131.1
Noise:	0.2		

Peak table for sample 6 : Atreca V2 Amplicon_0.8X clean-up

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	1,111	731.58	997.6		96.13
3	10,380	75.00	10.9	Upper Marker	113.00
4	13,121	0.00	0.0		115.65

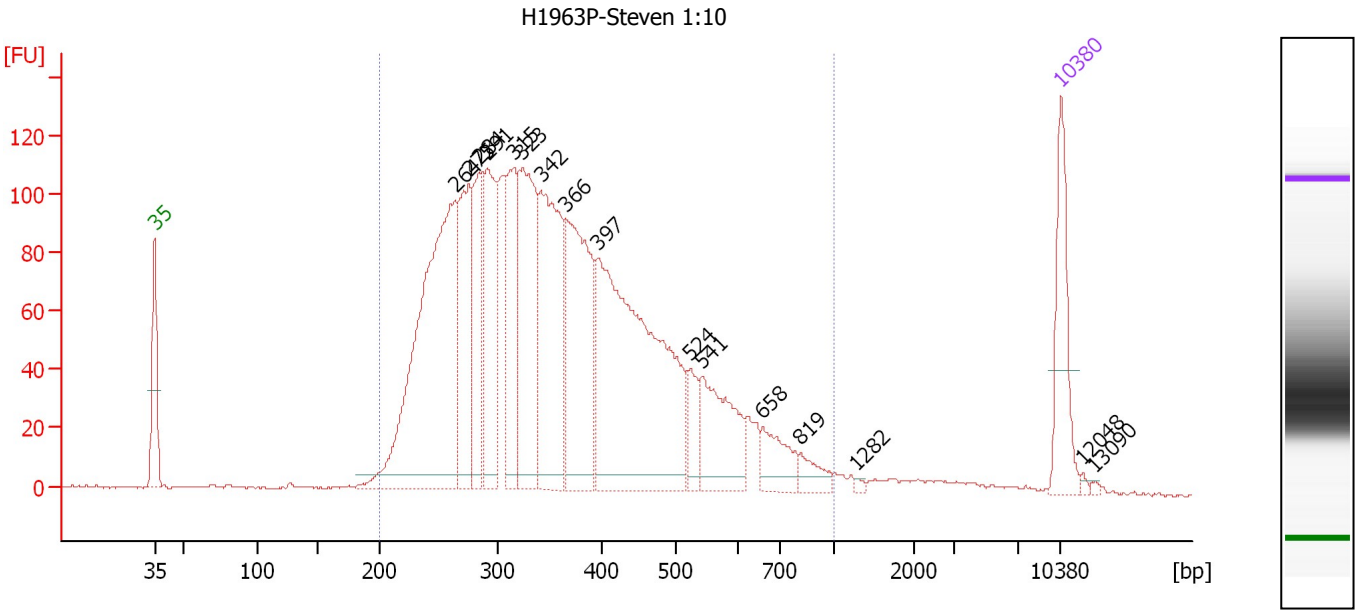
Region table for sample 6 : Atreca V2 Amplicon_0.8X clean-up

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	974	93.19	131.1	145.0	12	2.1

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : H1963P-Steven 1:10

Number of peaks found:	16	Corr. Area 1:	2,846.9
Noise:	0.4		

Peak table for sample 7 : H1963P-Steven 1:10

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	264	528.77	3,035.3		66.22
3	275	166.65	916.6		67.27
4	284	130.17	693.9		68.08
5	291	169.84	884.8		68.68
6	315	146.90	707.3		70.70
7	323	222.49	1,042.1		71.40
8	342	247.68	1,098.0		72.86
9	366	245.51	1,017.1		74.78
10	397	487.43	1,862.1		77.24
11	524	37.72	109.2		84.40
12	541	115.51	323.4		85.25
13	658	47.48	109.3		89.94
14	819	21.43	39.6		92.91
15	1,282	2.96	3.5		97.19
16	10,380	75.00	10.9	Upper Marker	113.00
17	12,048	0.00	0.0		114.61
18	13,090	0.00	0.0		115.62

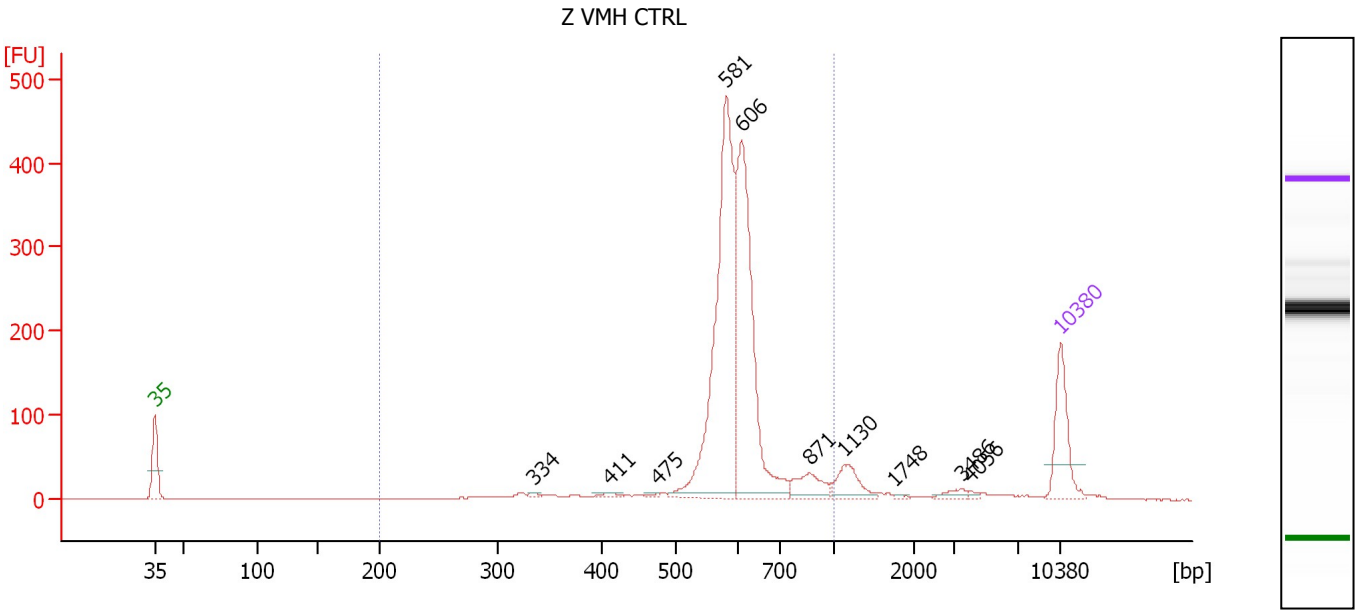
Region table for sample 7 : H1963P-Steven 1:10

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	376	2,762.85	2,846.9	12,627.3	96	33.5

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : Z VMH CTRL

Number of peaks found:	10	Corr. Area 1:	1,820.2
Noise:	0.3		

Peak table for sample 8 : Z VMH CTRL

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	334	3.87	17.6		72.21
3	411	6.89	25.4		78.15
4	475	3.32	10.6		81.79
5	581	482.66	1,259.2		87.17
6	606	391.86	979.5		88.30
7	871	42.18	73.4		93.63
8	1,130	41.04	55.0		96.24
9	1,748	2.38	2.1		100.09
10	3,486	9.33	4.1		105.31
11	4,056	2.96	1.1		106.03
12	10,380	75.00	10.9	Upper Marker	113.00

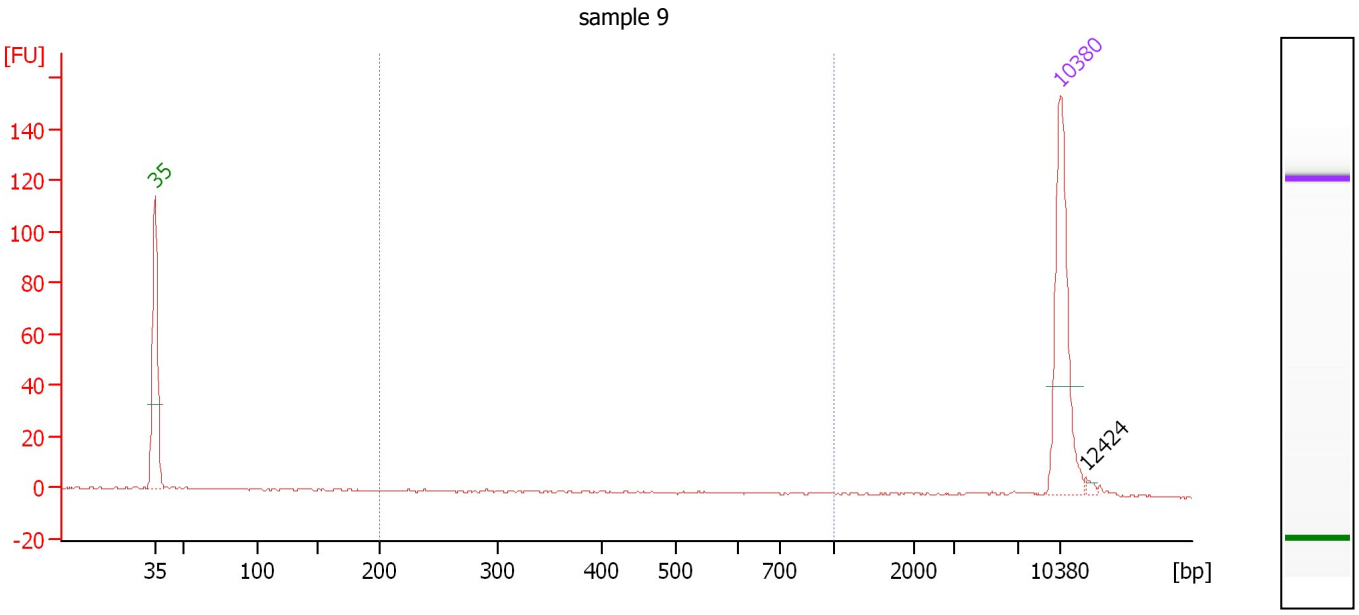
Region table for sample 8 : Z VMH CTRL

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Color % of Total	Size distribution in CV [%]
200	1,000	599	1,011.53	1,820.2	2,653.4	90	14.9

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : sample 9

Number of peaks found:	1	Corr. Area 1:	14.7
Noise:	0.3		

Peak table for sample 9 : sample 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	▶	10,380	75.00	10.9	Upper Marker	113.00
3		12,424	0.00	0.0		114.97

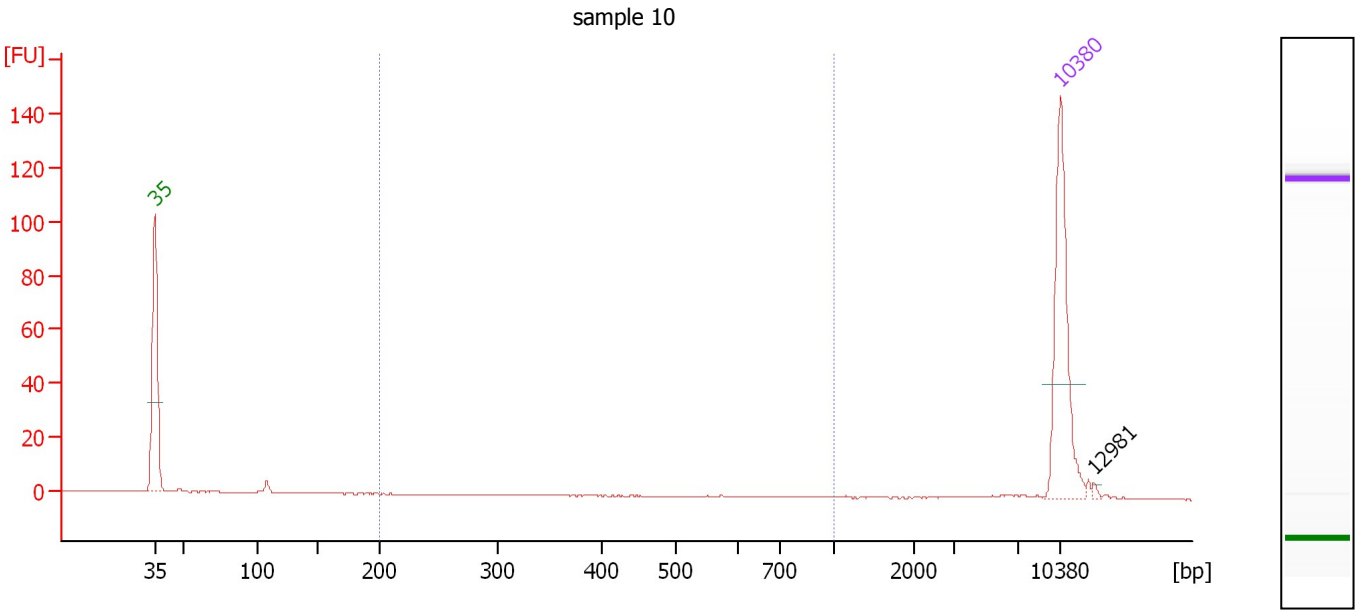
Region table for sample 9 : sample 9

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	511	10.37	14.7	39.1	27	40.0

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : sample 10

Number of peaks found:	1	Corr. Area 1:	0.0
Noise:	0.1		

Peak table for sample 10 : sample 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	10,380	75.00	10.9	Upper Marker	113.00
3	12,981	0.00	0.0		115.51

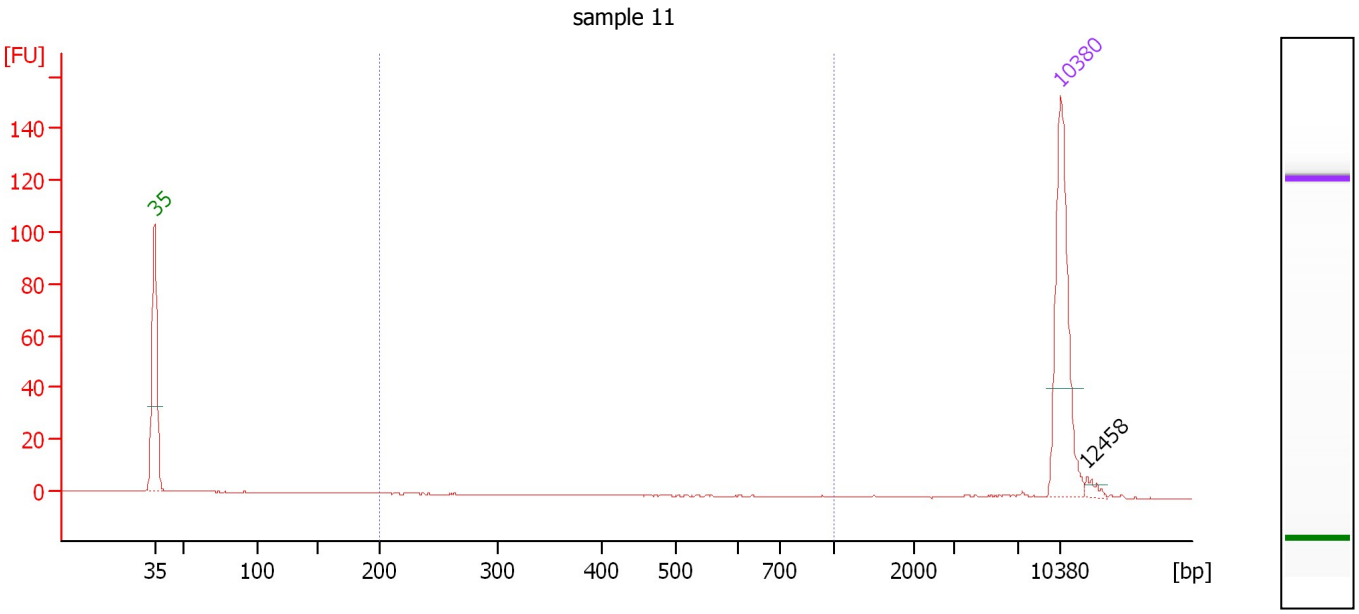
Region table for sample 10 : sample 10

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	572	0.00	0.0	0.0	0	0.1

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : sample 11

Number of peaks found:	1	Corr. Area 1:	0.0
Noise:	0.1		

Peak table for sample 11 : sample 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	10,380	75.00	10.9	Upper Marker	113.00
3	12,458	0.00	0.0		115.01

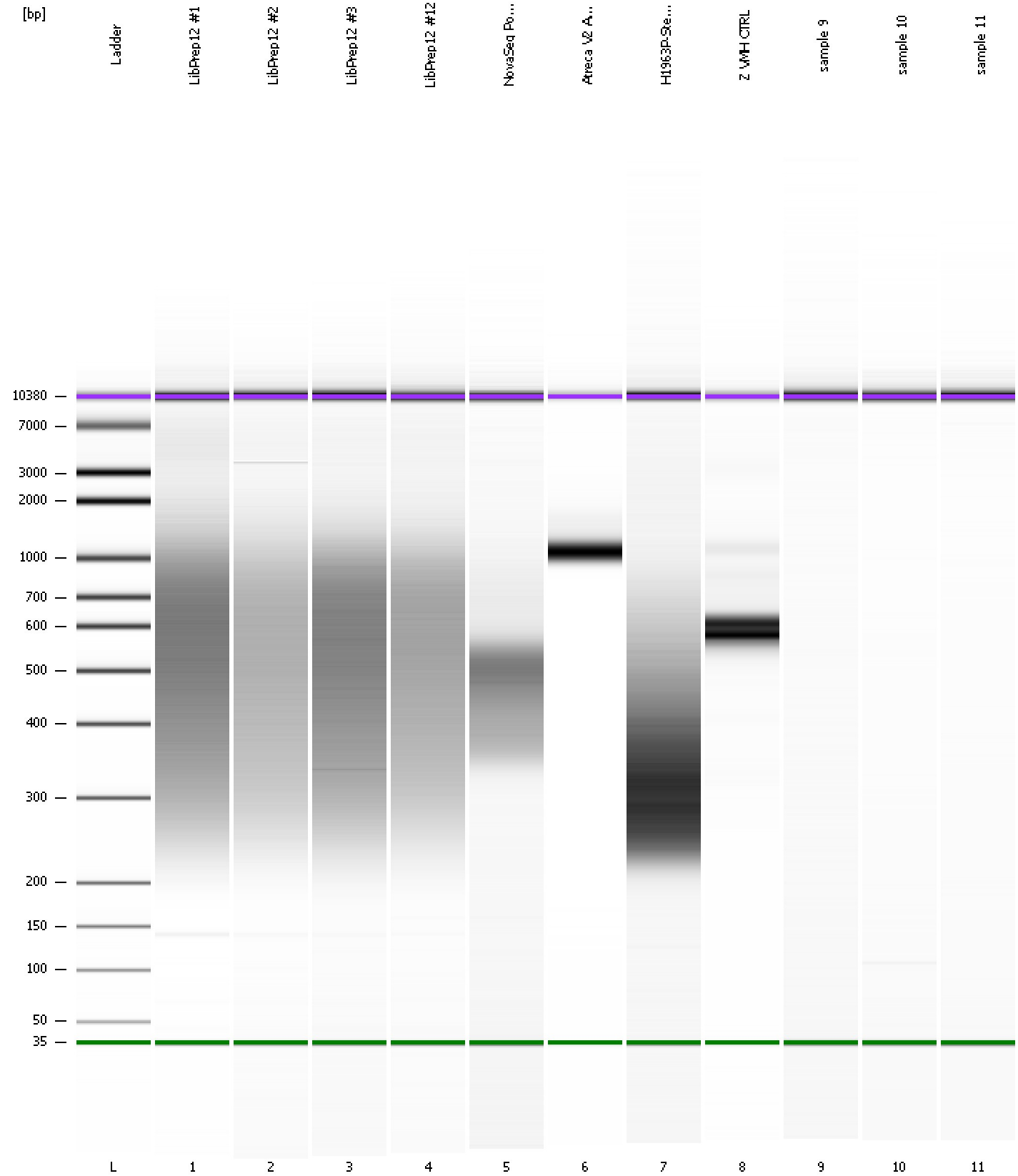
Region table for sample 11 : sample 11

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	922	0.03	0.0	0.0	0	6.6

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2019-10-15\2019-10-15_001.xad

Created: 10/15/2019 12:07:20 PM
Modified: 10/15/2019 4:03:52 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 12:48:36 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_001.xad)		Instrument	Run		10/15/2019 12:07:25 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938C		Instrument	Run		10/15/2019 12:07:25 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		10/15/2019 12:07:25 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		10/15/2019 12:07:25 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE34903152		Instrument	Run		10/15/2019 12:07:25 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		10/15/2019 12:07:25 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		10/15/2019 12:07:25 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1