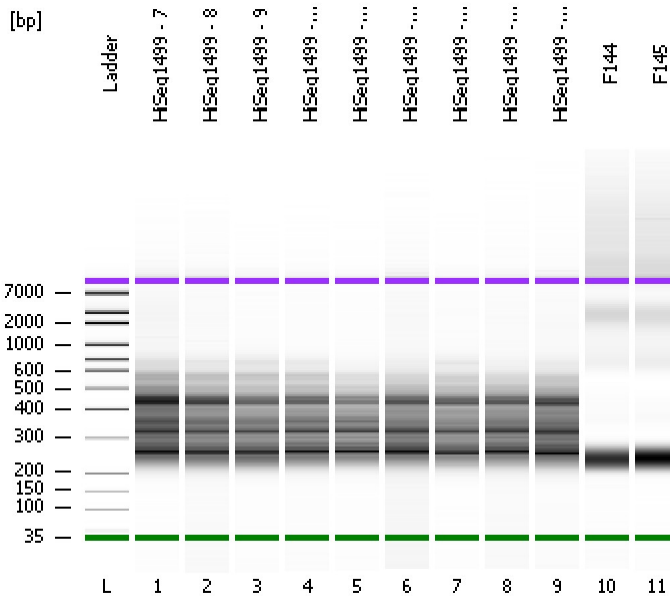


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

Created: 8/27/2018 2:41:08 PM
Modified: 8/27/2018 3:28:42 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

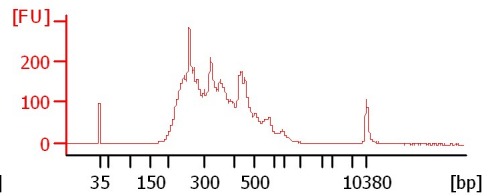
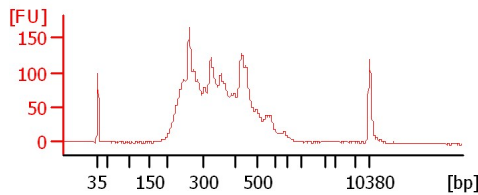
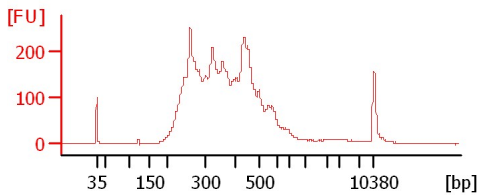
Chip Information:

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

HiSeq1499 - 7

HiSeq1499 - 8

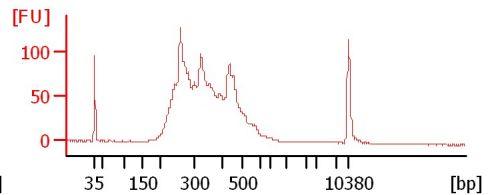
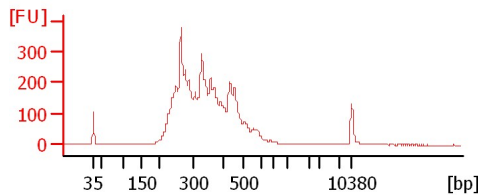
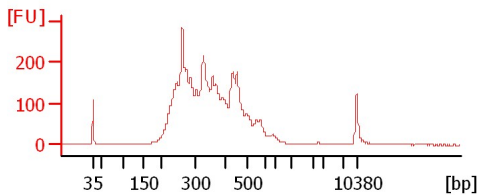
HiSeq1499 - 9



HiSeq1499 - 10

HiSeq1499 - 11

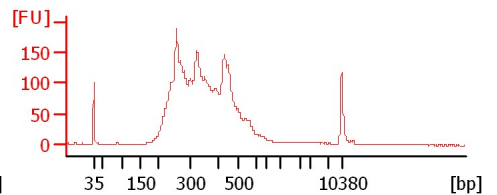
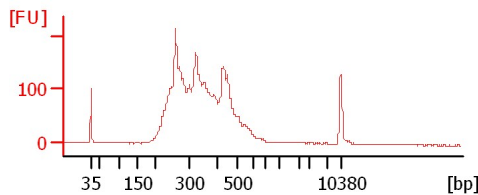
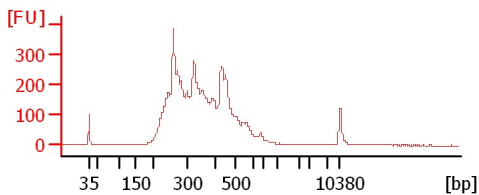
HiSeq1499 - 12



HiSeq1499 - 13

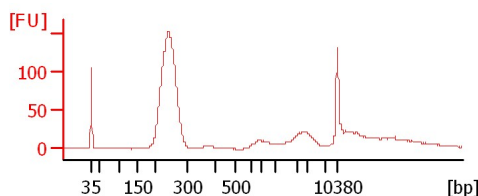
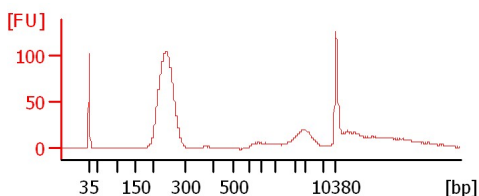
HiSeq1499 - 14

HiSeq1499 - 15



F144

F145



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

Created: 8/27/2018 2:41:08 PM
 Modified: 8/27/2018 3:28:42 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
HiSeq1499 - 7		<input type="checkbox"/>	✓			
HiSeq1499 - 8		<input type="checkbox"/>	✓			
HiSeq1499 - 9		<input type="checkbox"/>	✓			
HiSeq1499 - 10		<input type="checkbox"/>	✓			
HiSeq1499 - 11		<input type="checkbox"/>	✓			
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HiSeq1499 - 13		<input type="checkbox"/>	✓			
HiSeq1499 - 14		<input type="checkbox"/>	✓			
HiSeq1499 - 15		<input type="checkbox"/>	✓			
F144		<input type="checkbox"/>	✓			
F145		<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\2018-08-27\2018-08-27_003.xad

Created: 8/27/2018 2:41:08 PM
Modified: 8/27/2018 3:28:42 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

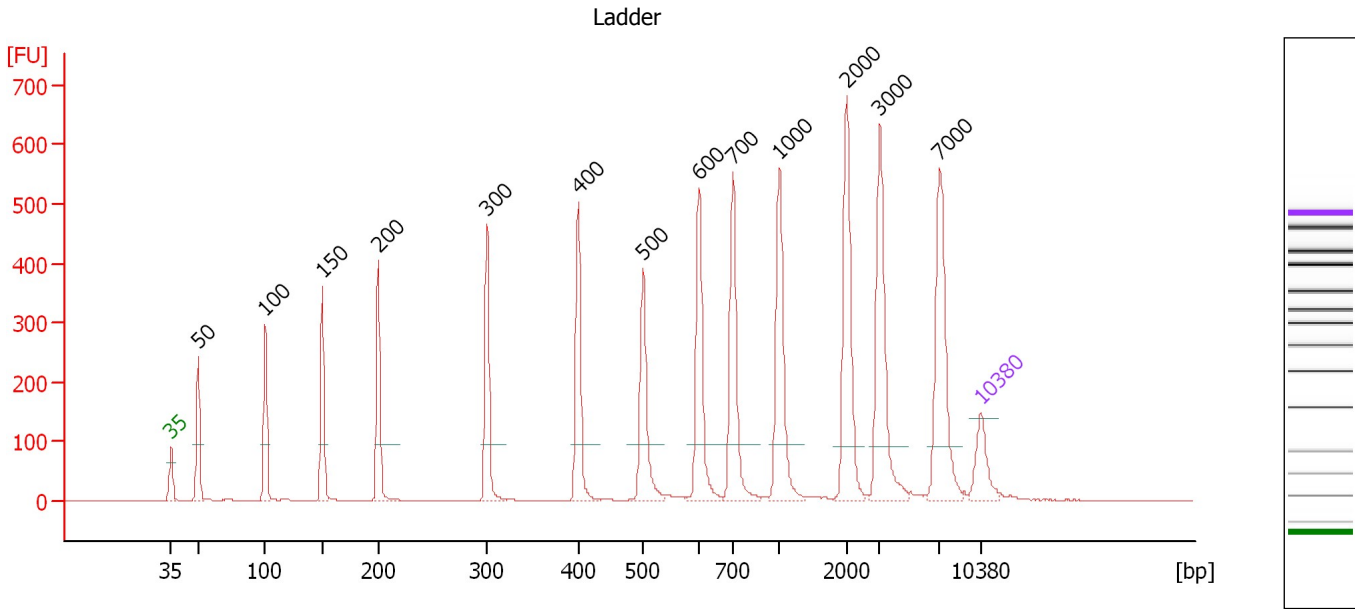
Ladder

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\2018-08-27\2018-08-27_003.xad

Created: 8/27/2018 2:41:08 PM
 Modified: 8/27/2018 3:28:42 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.3

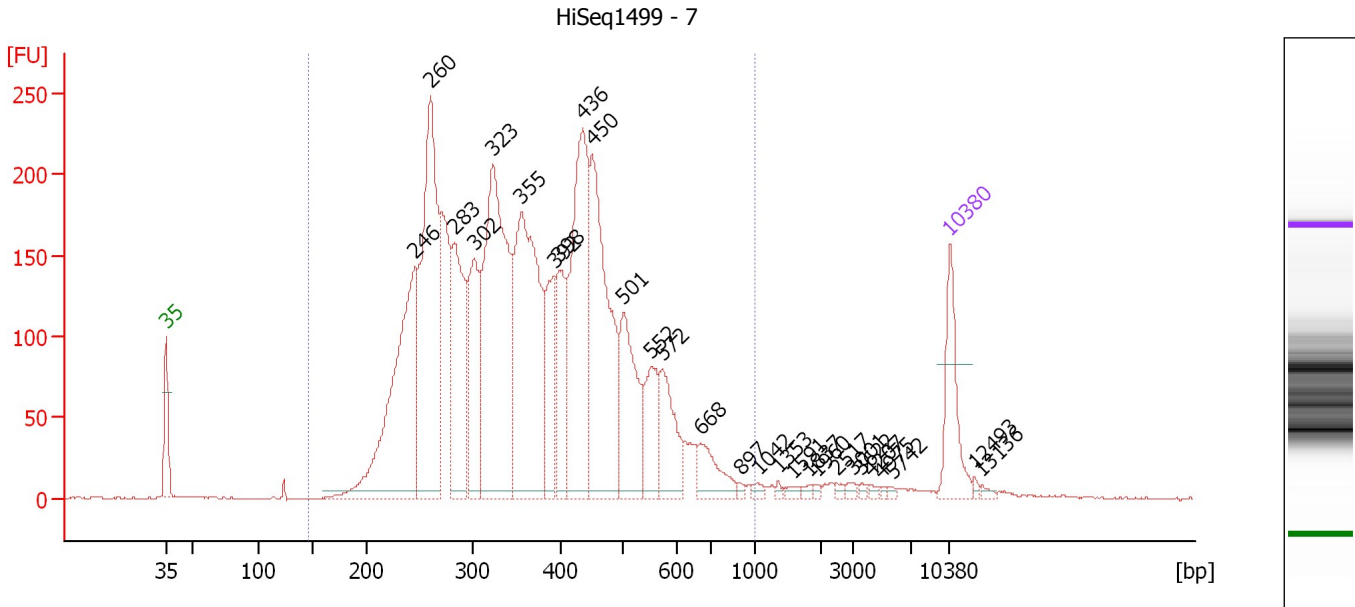
Peak table for Ladder

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	50	150.00	4,545.5	Ladder Peak	45.35
3	100	150.00	2,272.7	Ladder Peak	51.16
4	150	150.00	1,515.2	Ladder Peak	56.10
5	200	150.00	1,136.4	Ladder Peak	60.94
6	300	150.00	757.6	Ladder Peak	70.35
7	400	150.00	568.2	Ladder Peak	78.23
8	500	150.00	454.5	Ladder Peak	83.81
9	600	150.00	378.8	Ladder Peak	88.65
10	700	150.00	324.7	Ladder Peak	91.60
11	1,000	150.00	227.3	Ladder Peak	95.62
12	2,000	150.00	113.6	Ladder Peak	101.43
13	3,000	150.00	75.8	Ladder Peak	104.28
14	7,000	150.00	32.5	Ladder Peak	109.50
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

Created: 8/27/2018 2:41:08 PM
 Modified: 8/27/2018 3:28:42 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : HiSeq1499 - 7

Number of peaks found: 28 Corr. Area 1: 5,411.2
 Noise: 0.3

Peak table for sample 1 : HiSeq1499 - 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	246	442.66	2,729.6		65.24
3	260	476.31	2,776.9		66.57
4	283	239.25	1,280.1		68.76
5	302	163.49	819.5		70.52
6	323	511.80	2,397.8		72.19
7	355	434.65	1,853.0		74.71
8	392	107.88	416.7		77.62
9	398	110.21	419.2		78.10
10	436	338.33	1,175.8		80.24
11	450	332.15	1,117.1		81.05
12	501	159.10	481.2		83.86
13	552	82.32	225.9		86.33
14	572	92.87	246.1		87.29
15	668	54.16	122.8		90.67
16	897	4.30	7.3		94.24
17	1,042	6.42	9.3		95.86
18	1,353	3.00	3.4		97.67
19	1,591	5.33	5.1		99.05
20	1,837	4.28	3.5		100.48
21	1,960	3.23	2.5		101.19
22	2,517	4.24	2.6		102.90
23	3,001	4.52	2.3		104.29
24	3,622	3.78	1.6		105.10
25	4,207	3.60	1.3		105.86

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

Created: 8/27/2018 2:41:08 PM
Modified: 8/27/2018 3:28:42 PM

Electropherogram Summary Continued ...

... Peak table for sample 1 : HiSeq1499 - 7

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	4,975	2.28	0.7		106.86
27	5,742	2.77	0.7		107.86
28	10,380	75.00	10.9	Upper Marker	113.00
29	12,493	0.00	0.0		115.19
30	13,136	0.00	0.0		115.86

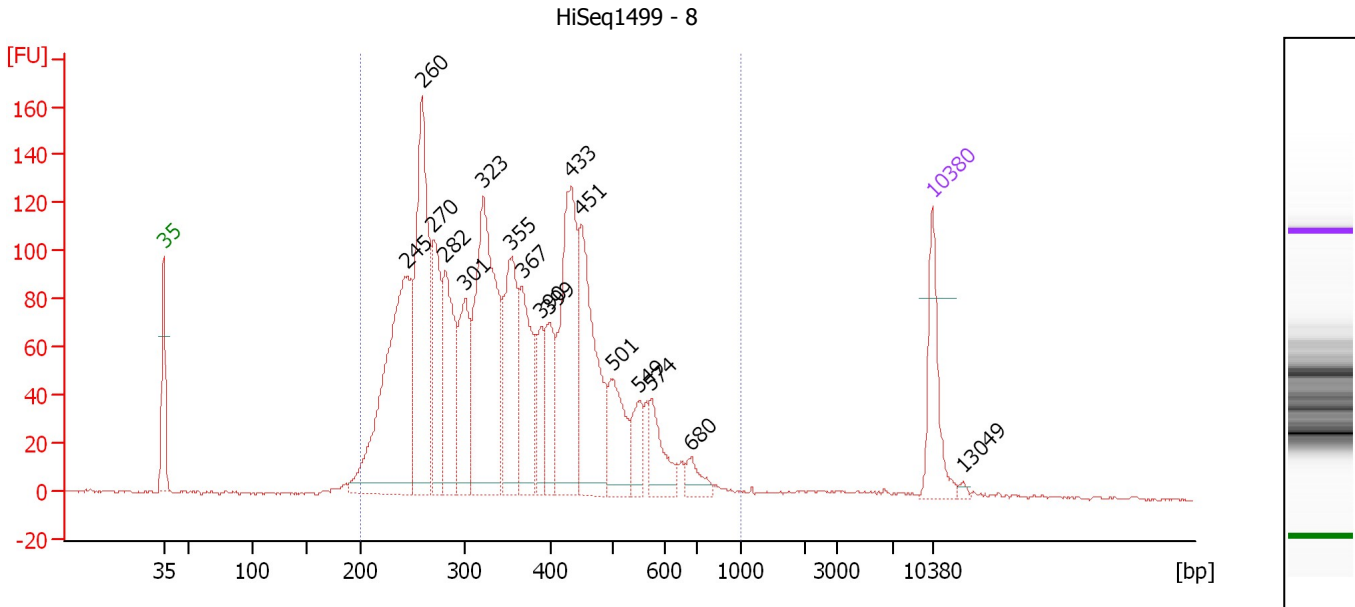
Region table for sample 1 : HiSeq1499 - 7

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
146	1,000	378	3,802.72	5,411.2	17,329.4	97	31.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

Created: 8/27/2018 2:41:08 PM
 Modified: 8/27/2018 3:28:42 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : HiSeq1499 - 8

Number of peaks found: 17 Corr. Area 1: 2,963.8
 Noise: 0.6

Peak table for sample 2 : HiSeq1499 - 8

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	245	460.56	2,852.6		65.14
3	260	366.35	2,138.3		66.54
4	270	158.11	885.8		67.56
5	282	179.57	963.8		68.68
6	301	141.07	710.0		70.43
7	323	391.34	1,838.1		72.13
8	355	183.87	785.7		74.65
9	367	141.93	586.1		75.62
10	390	66.28	257.7		77.42
11	399	86.48	328.5		78.15
12	433	257.22	899.5		80.09
13	451	232.56	781.9		81.06
14	501	94.37	285.2		83.87
15	549	46.46	128.1		86.20
16	574	63.45	167.3		87.42
17	680	26.21	58.4		91.01
18	10,380	75.00	10.9	Upper Marker	113.00
19	13,049	0.00	0.0		115.77

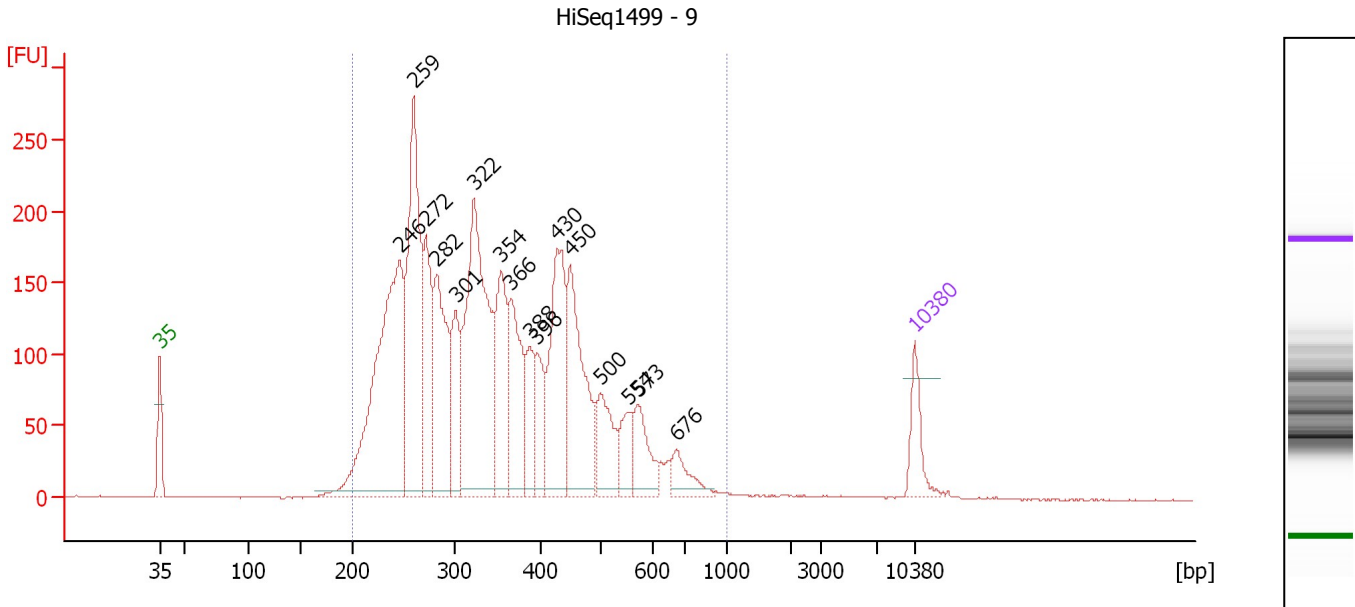
Region table for sample 2 : HiSeq1499 - 8

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	365	2,944.06	2,963.8	13,770.9	97	30.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

Created: 8/27/2018 2:41:08 PM
 Modified: 8/27/2018 3:28:42 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : HiSeq1499 - 9

Number of peaks found: 16 Corr. Area 1: 4,744.7
 Noise: 0.4

Peak table for sample 3 : HiSeq1499 - 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	246	968.44	5,969.8		65.25
3	259	700.03	4,088.0		66.53
4	272	273.19	1,524.3		67.67
5	282	386.11	2,074.1		68.66
6	301	206.90	1,040.9		70.44
7	322	766.37	3,600.8		72.12
8	354	290.51	1,241.8		74.64
9	366	269.85	1,116.0		75.58
10	388	134.22	523.7		77.31
11	396	123.15	470.7		77.95
12	430	375.76	1,325.5		79.88
13	450	371.81	1,252.1		81.02
14	500	157.68	477.4		83.83
15	554	86.48	236.7		86.40
16	573	116.40	307.8		87.34
17	676	70.46	157.9		90.90
18	10,380	75.00	10.9	Upper Marker	113.00

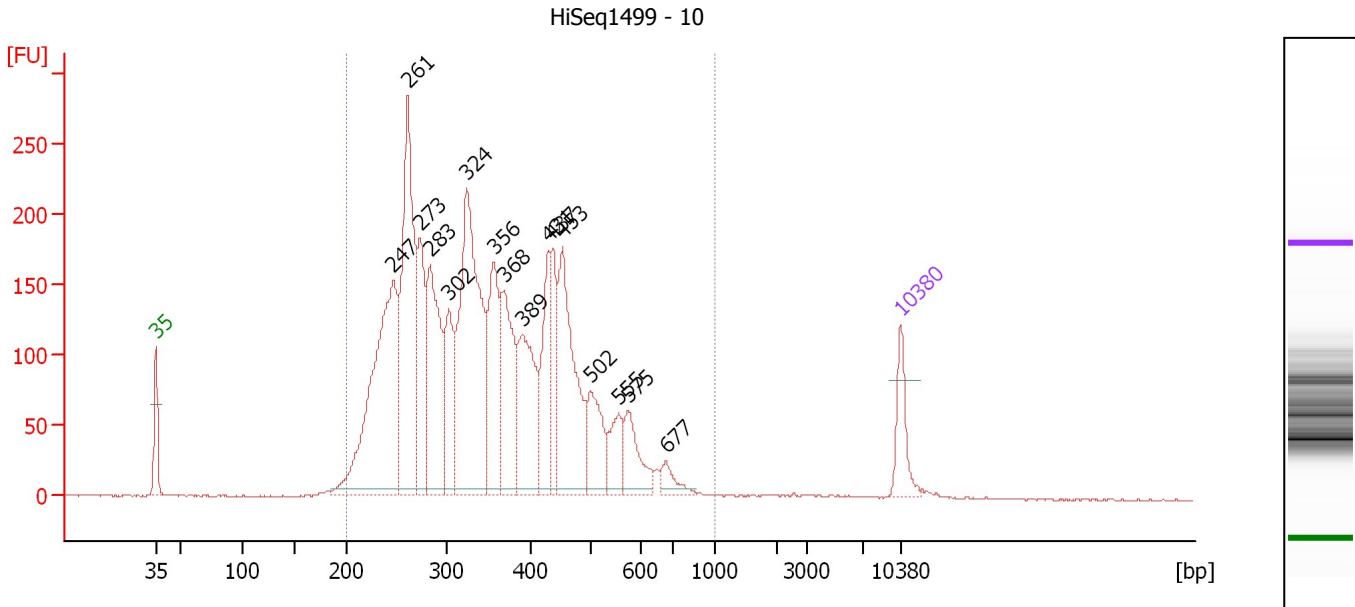
Region table for sample 3 : HiSeq1499 - 9

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	361	5,349.19	4,744.7	25,428.0	98	31.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

Created: 8/27/2018 2:41:08 PM
 Modified: 8/27/2018 3:28:42 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : HiSeq1499 - 10

Number of peaks found: 16 Corr. Area 1: 4,695.9
 Noise: 0.5

Peak table for sample 4 : HiSeq1499 - 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	247	809.57	4,975.9		65.31
3	261	658.12	3,821.9		66.67
4	273	253.44	1,405.7		67.82
5	283	397.47	2,125.8		68.77
6	302	202.52	1,014.9		70.53
7	324	735.59	3,440.5		72.23
8	356	280.83	1,194.0		74.79
9	368	279.85	1,152.8		75.69
10	389	280.66	1,091.9		77.40
11	431	199.72	702.3		79.96
12	437	137.19	475.4		80.31
13	453	382.08	1,276.8		81.21
14	502	144.62	436.3		83.92
15	555	89.52	244.4		86.47
16	575	107.03	282.2		87.43
17	677	40.01	89.5		90.94
18	10,380	75.00	10.9	Upper Marker	113.00

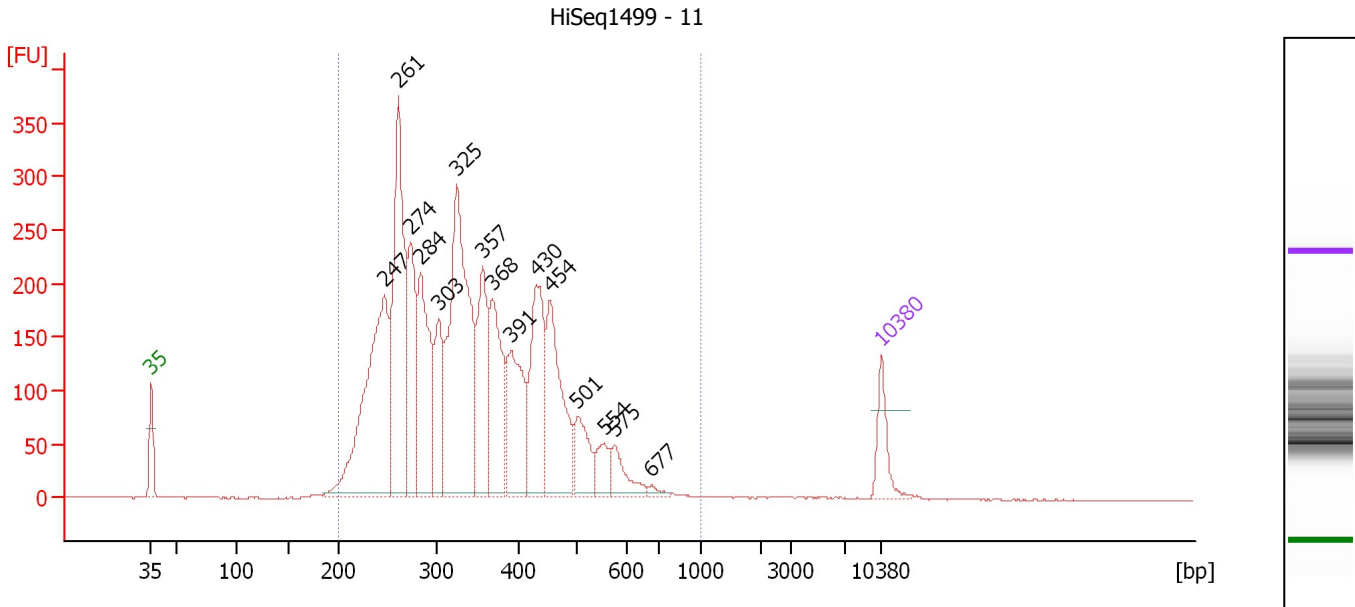
Region table for sample 4 : HiSeq1499 - 10

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	360	5,067.63	4,695.9	23,840.0	98	29.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

Created: 8/27/2018 2:41:08 PM
 Modified: 8/27/2018 3:28:42 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : HiSeq1499 - 11

Number of peaks found: 15 Corr. Area 1: 5,487.2
 Noise: 0.4

Peak table for sample 5 : HiSeq1499 - 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	247	806.22	4,948.7		65.34
3	261	689.38	4,002.3		66.67
4	274	303.84	1,680.0		67.90
5	284	408.46	2,180.6		68.82
6	303	211.17	1,056.9		70.56
7	325	832.80	3,885.2		72.30
8	357	299.25	1,271.7		74.80
9	368	306.73	1,262.1		75.72
10	391	285.06	1,104.9		77.51
11	430	320.05	1,127.1		79.92
12	454	351.05	1,171.4		81.25
13	501	126.10	381.4		83.85
14	554	70.82	193.8		86.41
15	575	73.40	193.5		87.43
16	677	14.01	31.4		90.91
17	10,380	75.00	10.9	Upper Marker	113.00

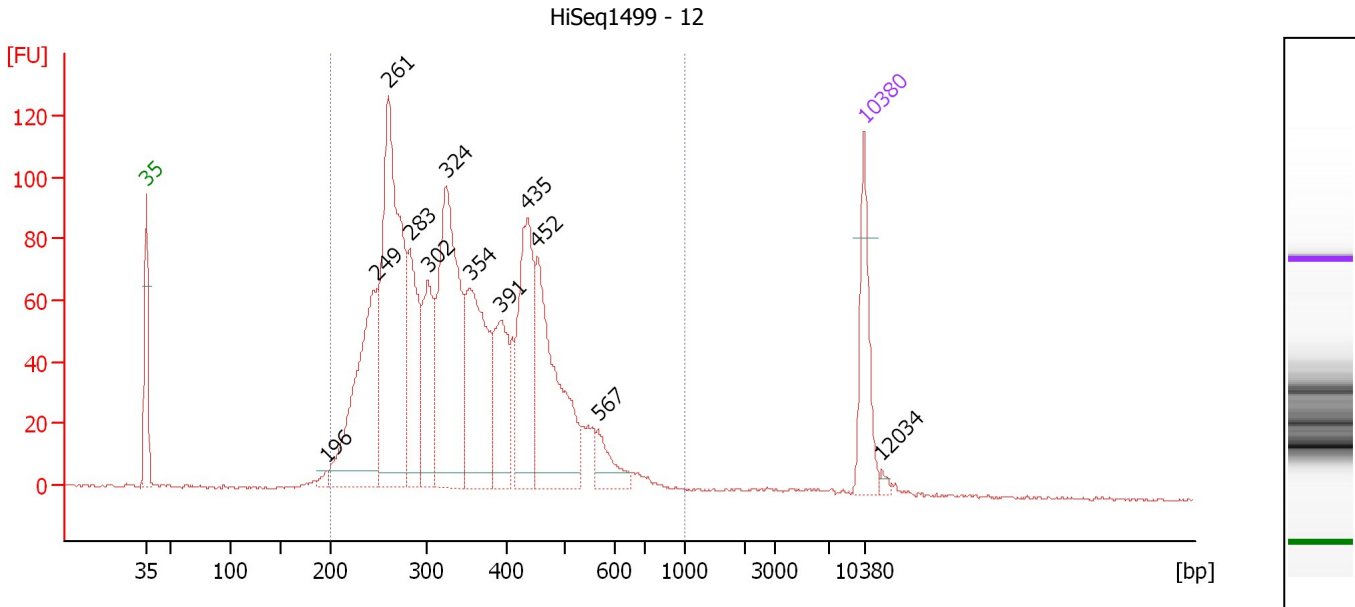
Region table for sample 5 : HiSeq1499 - 11

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	351	5,142.65	5,487.2	24,496.9	99	27.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

Created: 8/27/2018 2:41:08 PM
 Modified: 8/27/2018 3:28:42 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : HiSeq1499 - 12

Number of peaks found: 12 Corr. Area 1: 2,144.1
 Noise: 0.5

Peak table for sample 6 : HiSeq1499 - 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	196	9.38	72.5		60.56
3	249	295.75	1,803.1		65.50
4	261	474.71	2,759.3		66.65
5	283	157.84	845.8		68.72
6	302	143.04	718.0		70.49
7	324	336.13	1,570.7		72.26
8	354	230.79	988.1		74.60
9	391	113.95	441.8		77.51
10	435	179.97	627.5		80.16
11	452	231.36	775.2		81.14
12	567	38.70	103.4		87.07
13	10,380	75.00	10.9	Upper Marker	113.00
14	12,034	0.00	0.0		114.71

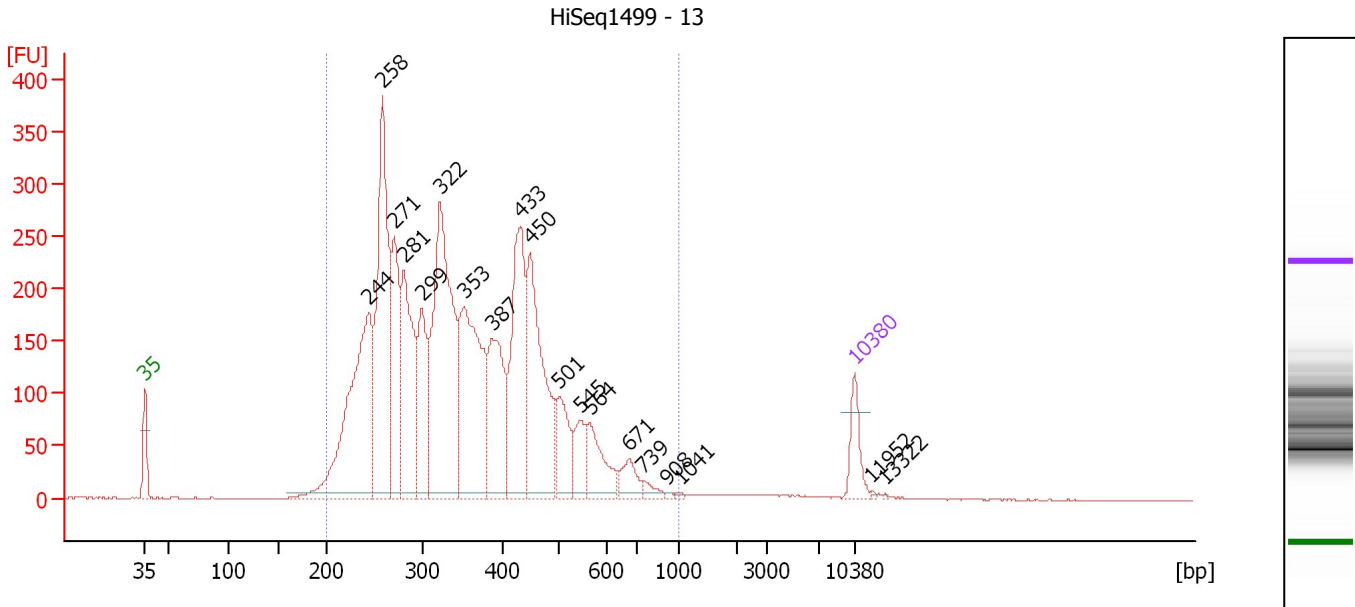
Region table for sample 6 : HiSeq1499 - 12

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	357	2,316.31	2,144.1	10,912.4	97	28.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

Created: 8/27/2018 2:41:08 PM
 Modified: 8/27/2018 3:28:42 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : HiSeq1499 - 13

Number of peaks found: 19 Corr. Area 1: 5,981.0
 Noise: 0.3

Peak table for sample 7 : HiSeq1499 - 13

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	244	865.60	5,372.4		65.09
3	258	824.27	4,838.8		66.40
4	271	345.25	1,930.6		67.61
5	281	473.18	2,551.2		68.56
6	299	286.13	1,447.7		70.30
7	322	913.47	4,297.7		72.08
8	353	636.15	2,732.6		74.50
9	387	351.93	1,376.4		77.24
10	433	476.71	1,667.7		80.08
11	450	512.64	1,725.8		81.02
12	501	166.46	503.3		83.86
13	545	92.83	258.3		85.97
14	564	150.89	405.3		86.91
15	671	66.68	150.5		90.75
16	739	25.47	52.2		92.12
17	908	4.44	7.4		94.38
18	1,041	4.55	6.6		95.85
19	10,380	75.00	10.9	Upper Marker	113.00
20	11,952	0.00	0.0		114.63
21	13,322	0.00	0.0		116.05

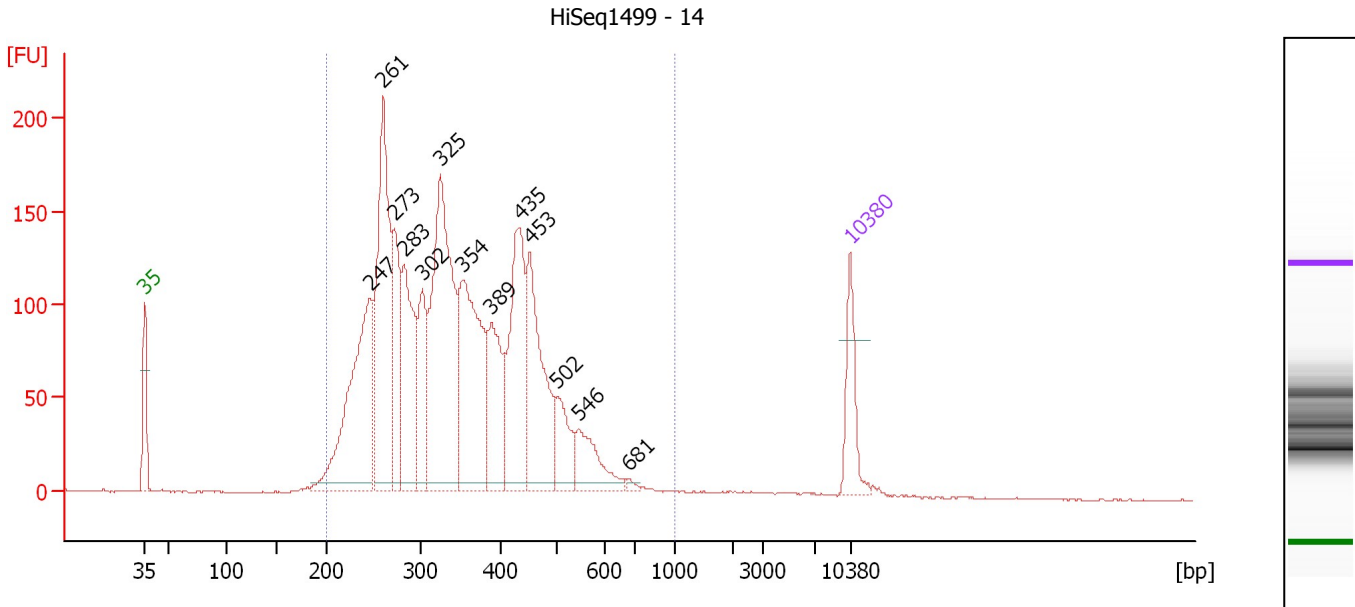
Region table for sample 7 : HiSeq1499 - 13

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	363	6,191.96	5,981.0	29,005.4	98	30.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 8 : HiSeq1499 - 14

Number of peaks found: 13 Corr. Area 1: 3,461.4
 Noise: 0.5

Peak table for sample 8 : HiSeq1499 - 14

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	247	462.56	2,834.8		65.38
3	261	462.03	2,684.9		66.65
4	273	199.86	1,111.1		67.76
5	283	253.84	1,360.6		68.71
6	302	157.75	791.1		70.51
7	325	545.59	2,544.0		72.31
8	354	375.16	1,606.6		74.59
9	389	165.83	645.3		77.39
10	435	289.29	1,007.1		80.20
11	453	264.26	883.4		81.20
12	502	92.53	279.4		83.90
13	546	94.78	263.2		86.02
14	681	6.31	14.0		91.04
15	10,380	75.00	10.9	Upper Marker	113.00

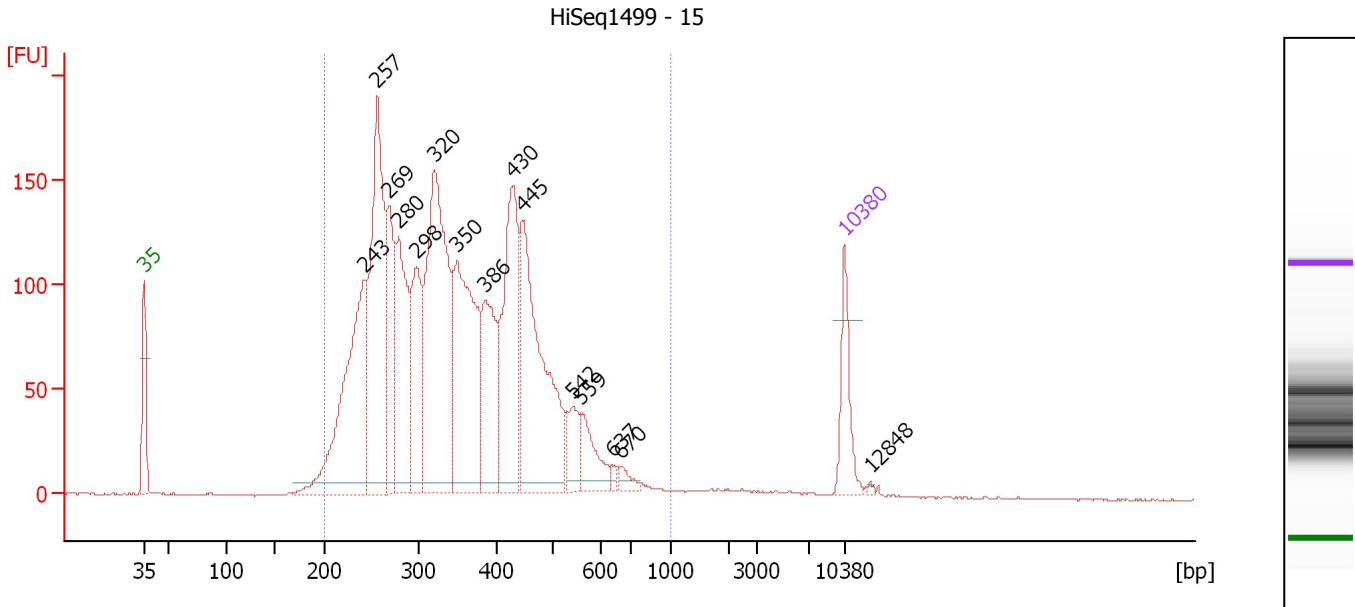
Region table for sample 8 : HiSeq1499 - 14

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	356	3,449.22	3,461.4	16,235.5	97	28.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 9 : HiSeq1499 - 15

Number of peaks found: 15 Corr. Area 1: 3,508.6
 Noise: 0.3

Peak table for sample 9 : HiSeq1499 - 15

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	243	515.43	3,216.4		64.96
3	257	528.51	3,116.2		66.30
4	269	216.27	1,216.1		67.47
5	280	292.62	1,585.4		68.43
6	298	217.56	1,104.9		70.19
7	320	598.86	2,832.5		71.95
8	350	430.80	1,864.5		74.29
9	386	209.35	821.9		77.12
10	430	328.86	1,159.3		79.89
11	445	441.82	1,504.1		80.75
12	542	61.25	171.1		85.86
13	559	78.18	211.9		86.66
14	637	9.10	21.6		89.76
15	670	18.48	41.8		90.72
16	10,380	75.00	10.9	Upper Marker	113.00
17	12,848	0.00	0.0		115.56

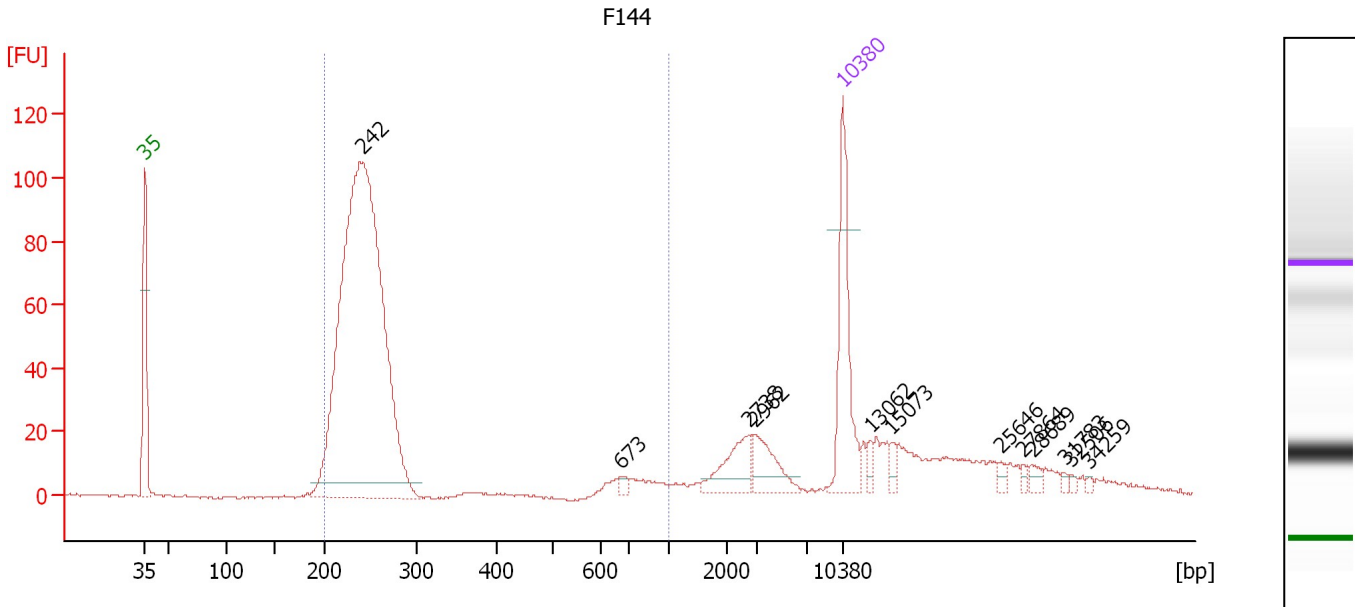
Region table for sample 9 : HiSeq1499 - 15

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	360	4,009.70	3,508.6	18,817.9	97	29.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 10 : F144

Number of peaks found: 12 Corr. Area 1: 829.8
 Noise: 0.2

Peak table for sample 10 : F144

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	242	836.30	5,240.0		64.87
3	673	4.69	10.6		90.81
4	2,738	34.47	19.1		103.53
5	2,962	29.97	15.3		104.18
6	10,380	75.00	10.9	Upper Marker	113.00
7	13,062	0.00	0.0		115.78
8	15,073	0.00	0.0		117.87
9	25,646	0.00	0.0		128.83
10	27,864	0.00	0.0		131.13
11	28,689	0.00	0.0		131.98
12	31,783	0.00	0.0		135.19
13	32,506	0.00	0.0		135.94
14	34,259	0.00	0.0		137.76

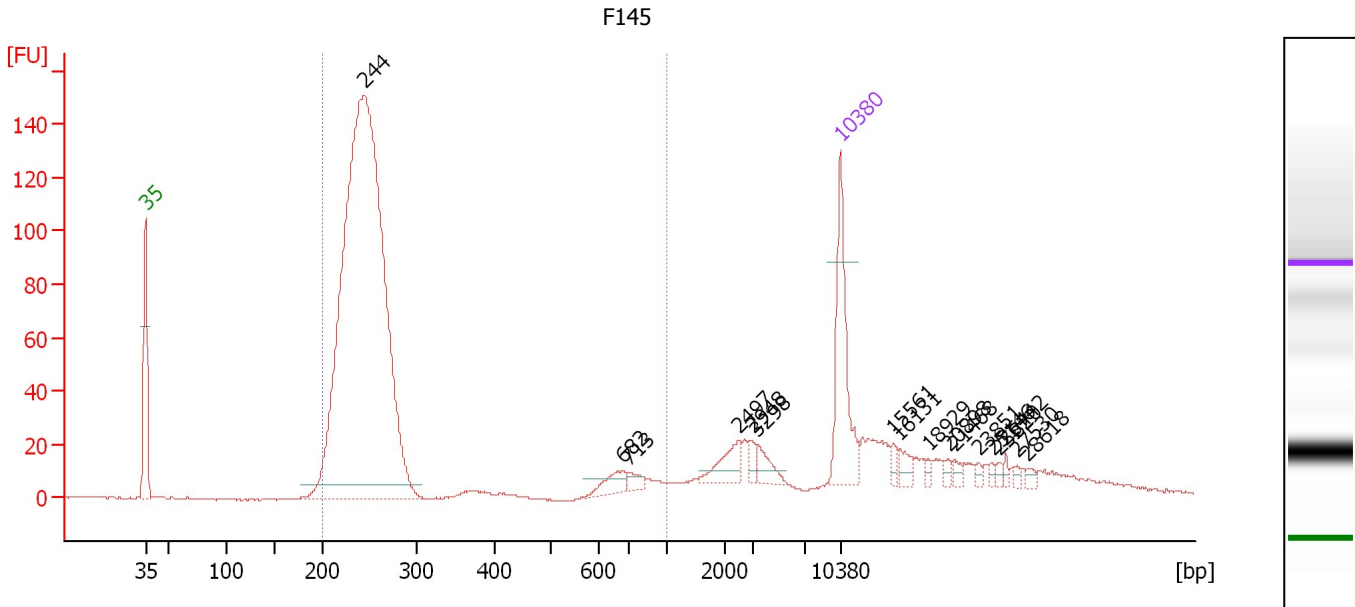
Region table for sample 10 : F144

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	269	838.32	829.8	5,156.7	71	44.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 11 : F145

Number of peaks found: 17 Corr. Area 1: 1,179.5
 Noise: 0.3

Peak table for sample 11 : F145

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	244	1,146.24	7,116.9		65.08
3	682	19.27	42.8		91.08
4	713	8.48	18.0		91.78
5	2,497	22.34	13.6		102.85
6	2,948	7.47	3.8		104.14
7	3,298	13.43	6.2		104.67
8	10,380	75.00	10.9	Upper Marker	113.00
9	15,561	0.00	0.0		118.37
10	16,131	0.00	0.0		118.96
11	18,929	0.00	0.0		121.86
12	20,898	0.00	0.0		123.91
13	21,468	0.00	0.0		124.50
14	23,851	0.00	0.0		126.97
15	25,146	0.00	0.0		128.31
16	25,820	0.00	0.0		129.01
17	26,442	0.00	0.0		129.65
18	27,530	0.00	0.0		130.78
19	28,618	0.00	0.0		131.91

Region table for sample 11 : F145

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	275	1,173.34	1,179.5	7,129.8	75	45.8

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-27\2018-08-27_003.xad

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Gel Image

