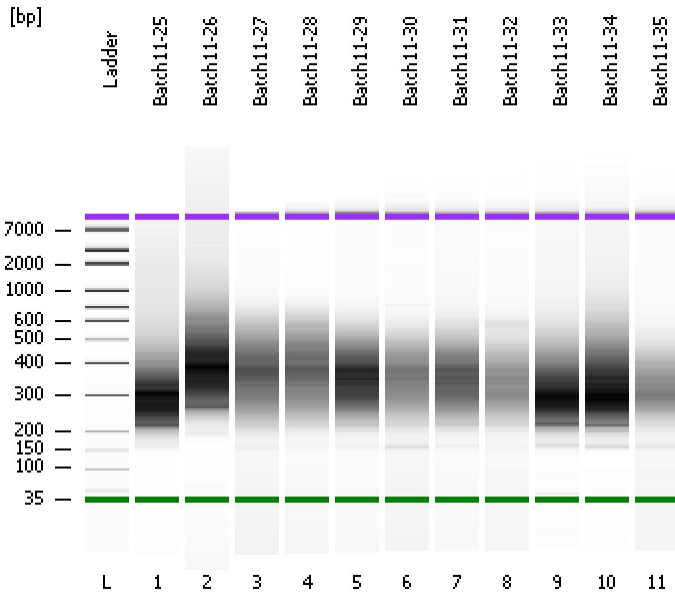


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
Modified: 10/22/2018 4:30:48 PM

Electrophoresis File Run Summary



Instrument Information:

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

Assay Information:

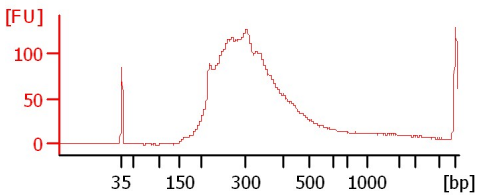
Assay Origin Path: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\assays\dsDNA\High Sensitivity DNA.xsy

Assay Class: High Sensitivity DNA Assay
Version: 1.03
Assay Comments: Copyright © 2003-2010 Agilent Technologies

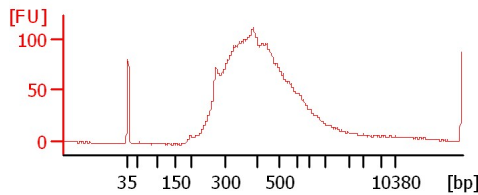
Chip Information:

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

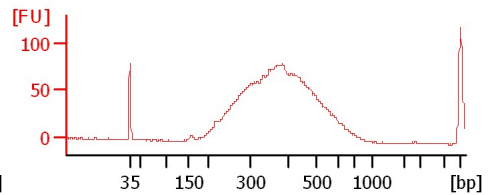
Batch11-25



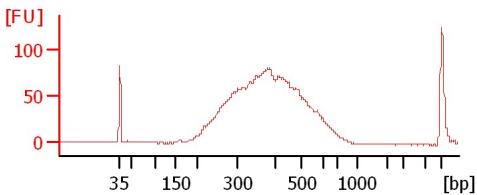
Batch11-26



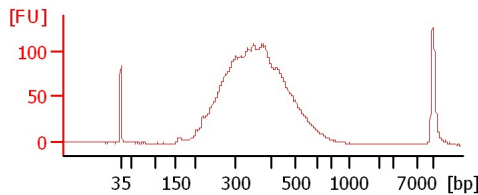
Batch11-27



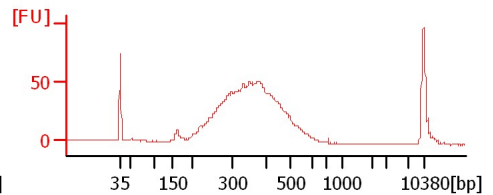
Batch11-28



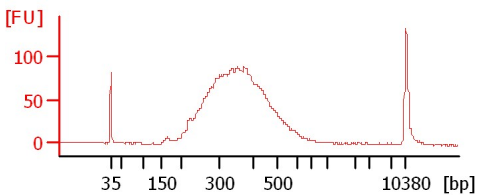
Batch11-29



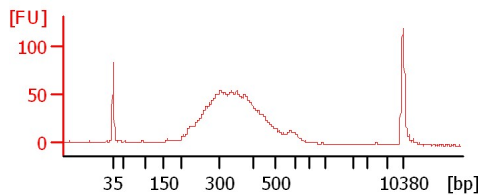
Batch11-30



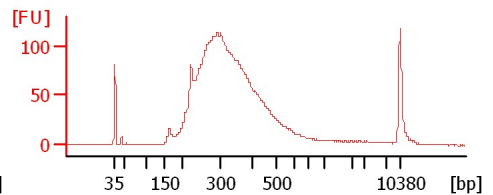
Batch11-31



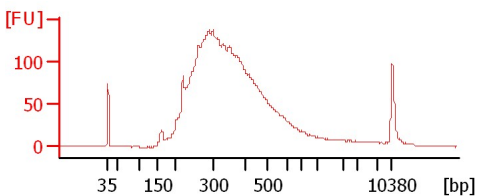
Batch11-32



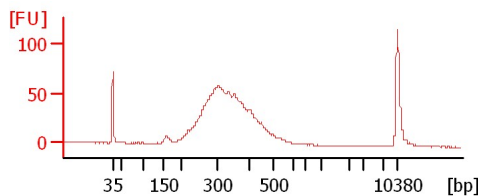
Batch11-33



Batch11-34



Batch11-35



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
 Modified: 10/22/2018 4:30:48 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Batch11-25		<input type="checkbox"/>	✓			
Batch11-26		<input type="checkbox"/>	✓			
Batch11-27		<input type="checkbox"/>	✓			
Batch11-28		<input type="checkbox"/>	✓			
Batch11-29		<input type="checkbox"/>	✓			
Batch11-30		<input type="checkbox"/>	✓			
Batch11-31		<input type="checkbox"/>	✓			
Batch11-32		<input type="checkbox"/>	✓			
Batch11-33		<input type="checkbox"/>	✓			
Batch11-34		<input type="checkbox"/>	✓			
Batch11-35		<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\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
Modified: 10/22/2018 4:30:48 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/ μ l] : 1950
Uses Standard Area for Ladder Fragments
Lower Marker Concentration [pg/ μ l] : 125
Upper Marker Concentration [pg/ μ l] : 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

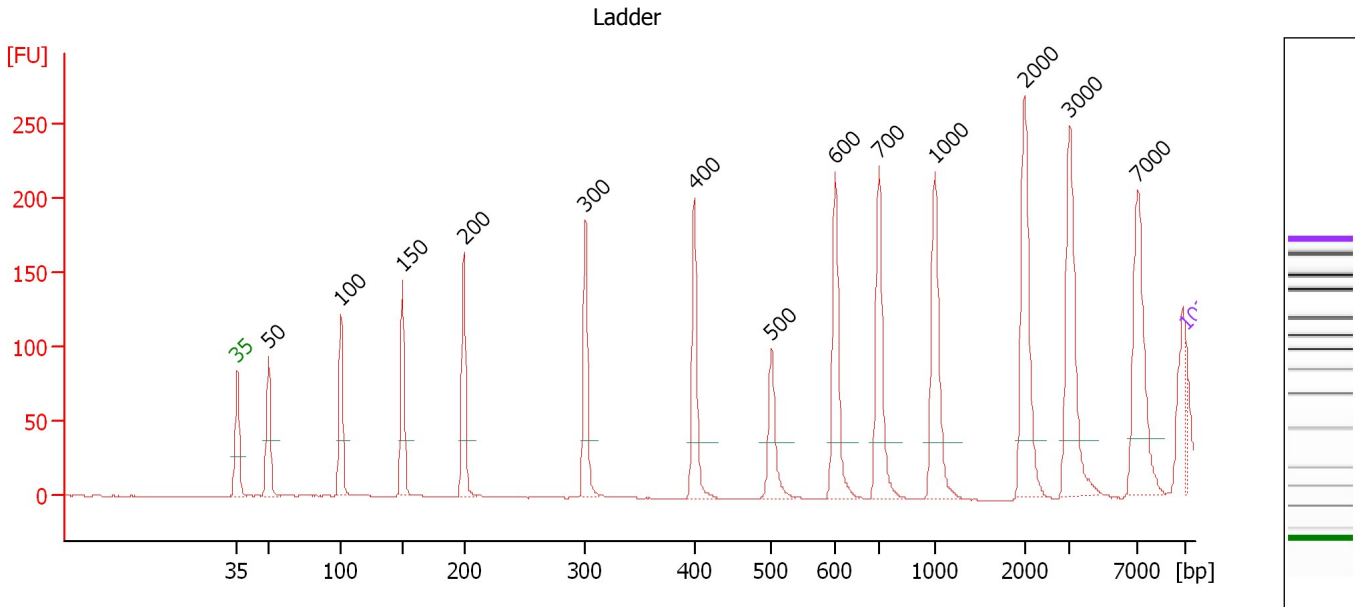
Ladder

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\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
 Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.3

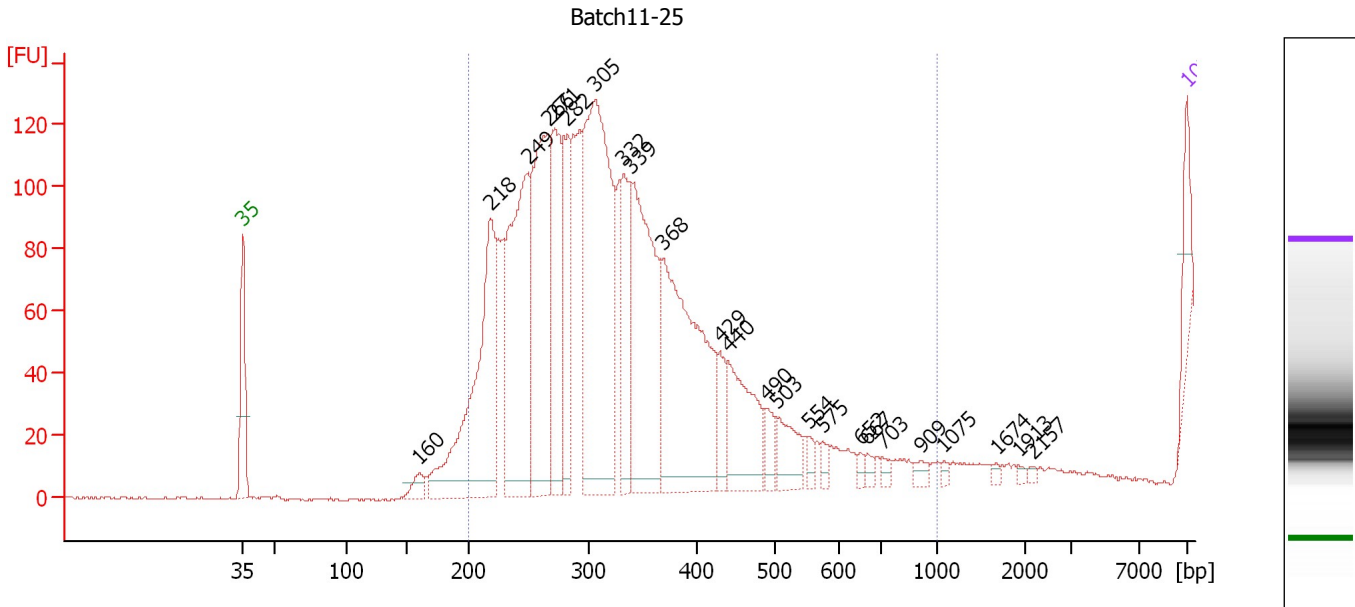
Peak table for Ladder

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	45.32
3	100	150.00	2,272.7	Ladder Peak	50.67
4	150	150.00	1,515.2	Ladder Peak	55.20
5	200	150.00	1,136.4	Ladder Peak	59.72
6	300	150.00	757.6	Ladder Peak	68.69
7	400	150.00	568.2	Ladder Peak	76.72
8	500	150.00	454.5	Ladder Peak	82.43
9	600	150.00	378.8	Ladder Peak	87.15
10	700	150.00	324.7	Ladder Peak	90.37
11	1,000	150.00	227.3	Ladder Peak	94.47
12	2,000	150.00	113.6	Ladder Peak	101.08
13	3,000	150.00	75.8	Ladder Peak	104.46
14	7,000	150.00	32.5	Ladder Peak	109.46
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
 Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : Batch11-25

Number of peaks found: 24 Corr. Area 1: 985.9
 Noise: 0.3


Peak table for sample 1 : Batch11-25

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	160	44.76	424.2		56.09
3	218	797.75	5,541.0		61.35
4	249	749.73	4,554.0		64.16
5	266	694.27	3,958.3		65.62
6	271	412.31	2,301.2		66.14
7	282	288.83	1,551.5		67.08
8	305	1,054.55	5,245.9		69.06
9	332	281.05	1,283.9		71.24
10	339	652.18	2,918.5		71.79
11	368	790.59	3,253.8		74.16
12	429	89.82	317.0		78.40
13	440	258.50	890.7		78.99
14	490	49.48	152.9		81.88
15	503	109.01	328.6		82.55
16	554	26.32	72.0		84.96
17	575	22.67	59.8		85.95
18	652	13.24	30.8		88.84
19	667	16.34	37.1		89.31
20	703	18.84	40.6		90.42
21	909	18.16	30.3		93.23
22	1,075	9.88	13.9		94.97
23	1,674	7.46	6.8		98.92
24	1,913	6.53	5.2		100.50
25	2,157	6.77	4.8		101.61

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...**... Peak table for sample 1 : Batch11-25**

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

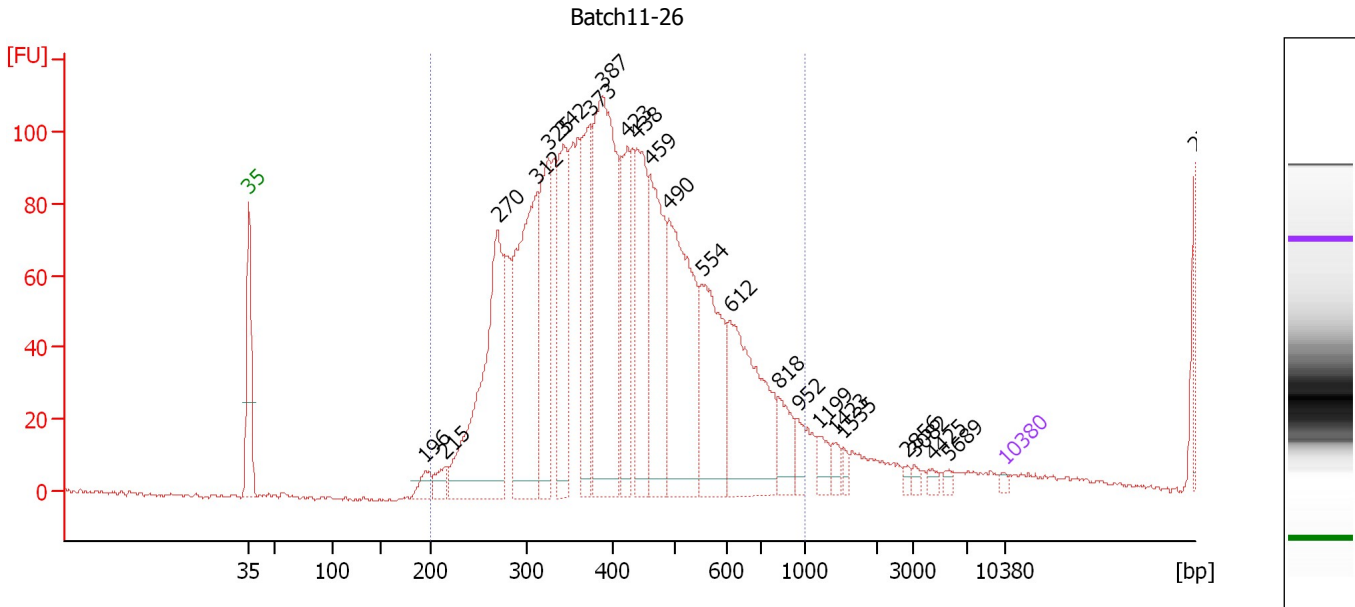
Region table for sample 1 : Batch11-25

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	284	2,443.92	985.9	13,298.8	 100	13.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
 Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : Batch11-26

Number of peaks found: 24 Corr. Area 1: 2,375.3
 Noise: 0.4

Peak table for sample 2 : Batch11-26

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	541.76	4,195.7		59.33
3	215	538.71	3,789.4		61.10
4	270	7,771.97	43,551.6		66.04
5	312	7,422.56	36,002.1		69.69
6	325	4,211.29	19,615.9		70.72
7	342	3,960.10	17,519.1		72.11
8	373	3,367.40	13,670.5		74.57
9	387	8,856.80	34,698.7		75.66
10	423	2,787.35	9,987.7		78.02
11	438	3,629.43	12,568.2		78.86
12	459	4,416.06	14,572.2		80.10
13	490	6,274.15	19,389.2		81.87
14	554	3,903.96	10,675.5		84.98
15	612	4,697.34	11,623.6		87.55
16	818	972.38	1,801.0		91.99
17	952	423.04	673.4		93.81
18	1,199	453.76	573.3		95.78
19	1,423	277.42	295.4		97.26
20	1,535	175.99	173.7		98.00
21	2,856	97.91	52.0		103.97
22	3,082	106.49	52.3		104.56
23	4,425	146.15	50.0		106.24
24	5,689	103.51	27.6		107.82
25	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...**... Peak table for sample 2 : Batch11-26**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	27,048	0.00	0.0		130.46

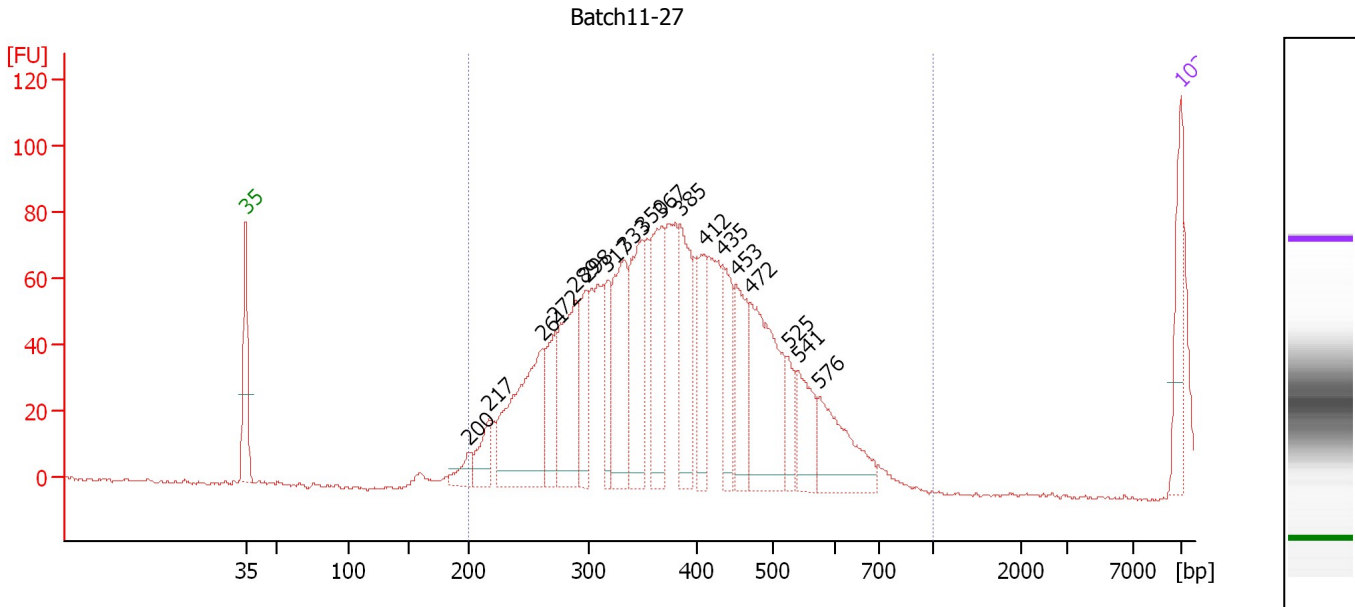
Region table for sample 2 : Batch11-26

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	425	67,509.56	2,375.3	270,473.1	 98	31.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
 Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : Batch11-27

Number of peaks found: 18 Corr. Area 1: 1,006.4
 Noise: 0.5

Peak table for sample 3 : Batch11-27

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	200	41.06	311.6		59.69
3	217	71.12	497.6		61.21
4	261	347.35	2,015.9		65.20
5	272	114.22	635.8		66.20
6	289	256.11	1,342.2		67.72
7	298	121.46	617.5		68.52
8	317	89.37	427.6		70.03
9	333	217.23	988.1		71.35
10	350	234.27	1,015.6		72.67
11	367	213.42	880.1		74.11
12	385	197.22	775.5		75.54
13	412	101.68	373.7		77.42
14	435	114.56	398.7		78.74
15	453	144.44	483.2		79.74
16	472	285.28	916.3		80.82
17	525	55.97	161.5		83.61
18	541	102.26	286.3		84.37
19	576	150.93	397.1		86.01
20	10,380	75.00	10.9	Upper Marker	113.00

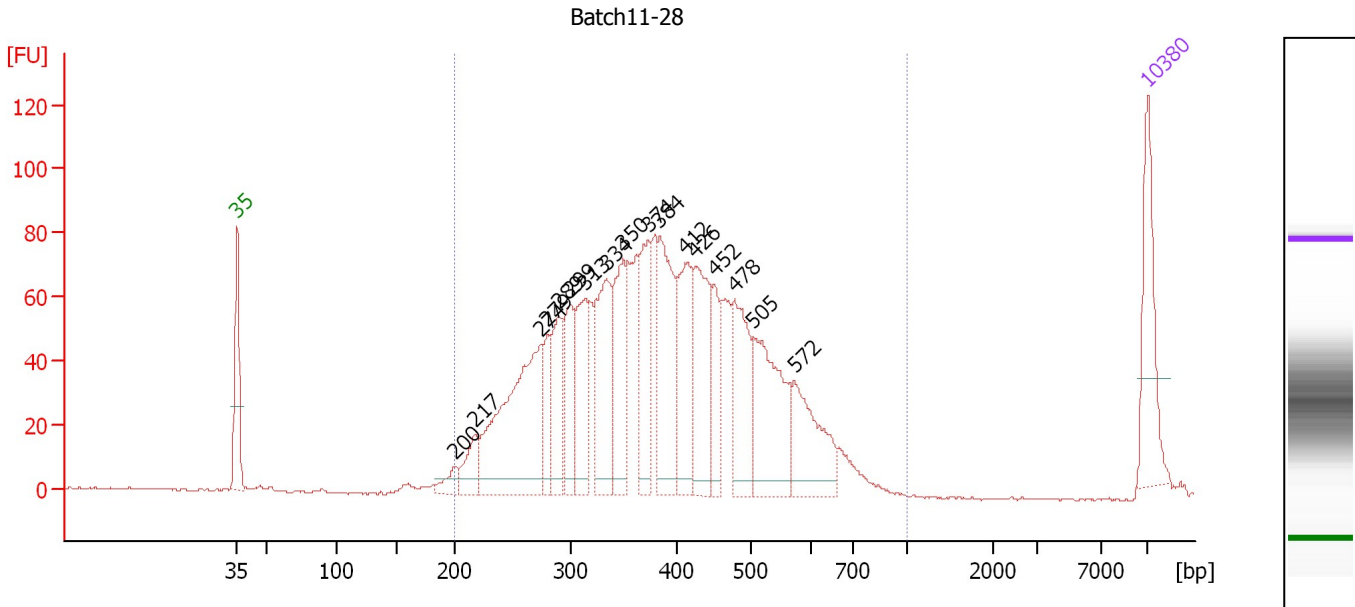
Region table for sample 3 : Batch11-27

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	1,780.42	1,006.4	7,763.1	99	18.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
 Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : Batch11-28

Number of peaks found: 17 Corr. Area 1: 2,014.0
 Noise: 0.4

Peak table for sample 4 : Batch11-28

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	200	21.62	163.4		59.76
3	217	43.16	300.9		61.28
4	274	280.57	1,549.0		66.40
5	279	49.75	269.7		66.85
6	289	75.50	395.8		67.71
7	299	83.77	424.4		68.61
8	313	95.87	463.6		69.76
9	334	142.82	648.4		71.40
10	350	111.36	482.7		72.67
11	374	99.03	401.7		74.60
12	384	151.19	596.1		75.46
13	412	110.55	406.2		77.43
14	426	124.86	444.1		78.20
15	452	53.41	179.1		79.68
16	478	109.02	345.2		81.20
17	505	150.86	452.4		82.67
18	572	96.59	255.8		85.83
19	10,380	75.00	10.9	Upper Marker	113.00

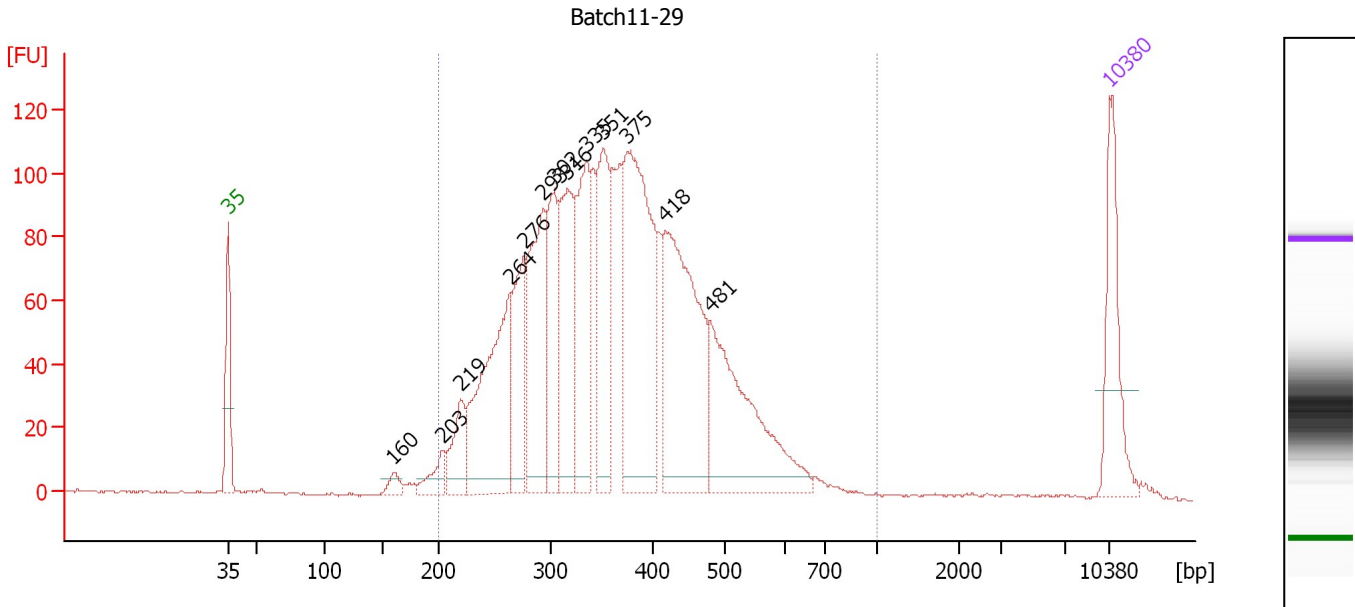
Region table for sample 4 : Batch11-28

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	395	2,061.34	2,014.0	8,768.4	99	27.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
 Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : Batch11-29

Number of peaks found: 13 Corr. Area 1: 2,587.2
 Noise: 0.3

Peak table for sample 5 : Batch11-29

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	160	14.58	137.8		56.13
3	203	29.31	218.9		59.98
4	219	60.78	419.8		61.46
5	264	237.42	1,363.8		65.44
6	276	114.22	628.0		66.50
7	293	185.45	960.4		68.03
8	302	116.13	582.1		68.87
9	316	162.49	779.2		69.98
10	335	163.95	741.6		71.50
11	351	144.77	624.4		72.81
12	375	324.85	1,314.2		74.68
13	418	280.76	1,018.7		77.72
14	481	216.43	682.2		81.32
15	10,380	75.00	10.9	Upper Marker	113.00

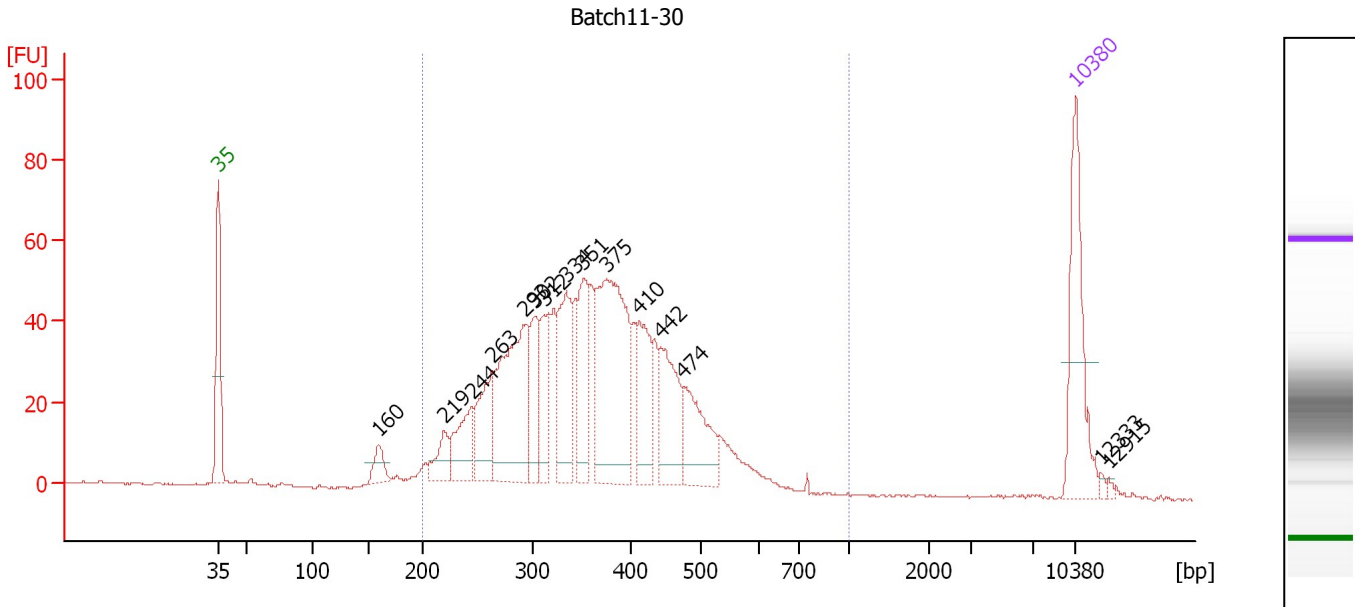
Region table for sample 5 : Batch11-29

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	370	2,291.94	2,587.2	10,243.3	97	26.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
 Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : Batch11-30

Number of peaks found: 15 Corr. Area 1: 1,191.0
 Noise: 0.3

Peak table for sample 6 : Batch11-30

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	160	22.65	214.5		56.10
3	219	30.15	208.7		61.41
4	244	54.74	339.8		63.68
5	263	68.94	397.9		65.33
6	293	167.66	868.2		68.03
7	302	54.23	272.6		68.81
8	312	57.21	278.0		69.64
9	334	93.07	422.2		71.43
10	351	74.80	323.0		72.78
11	375	202.65	818.2		74.74
12	410	65.56	242.1		77.30
13	442	72.44	248.2		79.13
14	474	67.97	217.1		80.96
15	10,380	75.00	10.9	Upper Marker	113.00
16	12,333	0.00	0.0		115.05
17	12,915	0.00	0.0		115.66

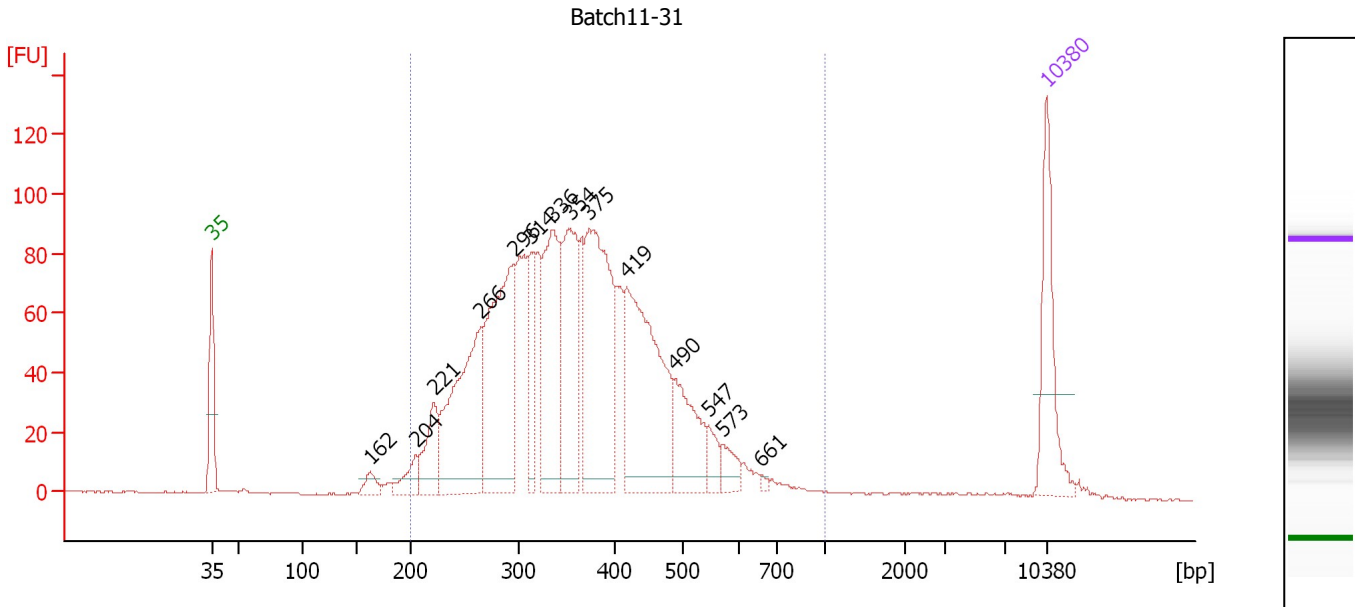
Region table for sample 6 : Batch11-30

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	1,288.30	1,191.0	5,781.3	96	24.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
 Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : Batch11-31

Number of peaks found: 14 Corr. Area 1: 2,165.8
 Noise: 0.4

Peak table for sample 7 : Batch11-31

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	162	18.01	168.0		56.32
3	204	27.75	205.7		60.12
4	221	63.65	435.6		61.64
5	266	232.11	1,323.5		65.62
6	296	259.80	1,329.4		68.34
7	314	64.22	309.8		69.82
8	336	177.17	799.5		71.56
9	354	159.55	682.7		73.04
10	375	257.48	1,041.1		74.69
11	419	248.20	896.9		77.82
12	490	86.37	267.2		81.84
13	547	19.92	55.1		84.66
14	573	21.55	57.0		85.87
15	661	2.86	6.5		89.13
16	10,380	75.00	10.9	Upper Marker	113.00

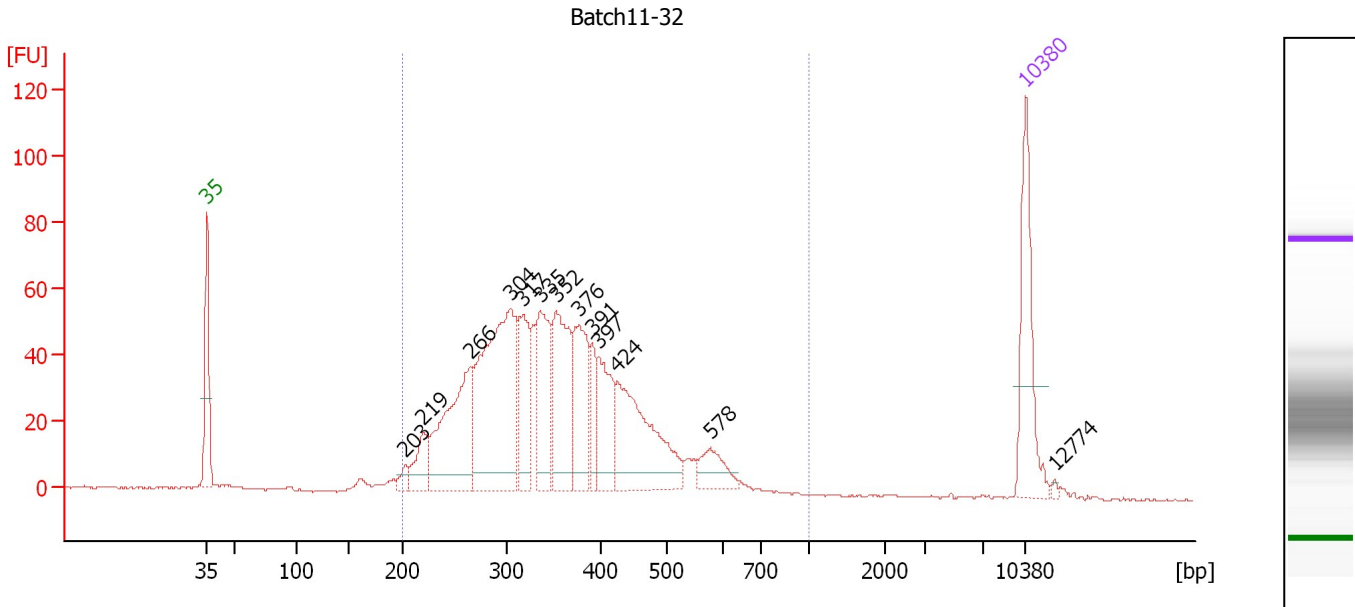
Region table for sample 7 : Batch11-31

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	369	1,951.00	2,165.8	8,775.4	97	27.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

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 Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : Batch11-32

Number of peaks found: 13 Corr. Area 1: 1,260.4
 Noise: 0.3

Peak table for sample 8 : Batch11-32

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	203	12.49	93.3		59.96
3	219	38.88	268.4		61.47
4	266	162.92	929.0		65.62
5	304	277.75	1,385.6		68.99
6	317	78.01	372.4		70.09
7	335	99.68	450.9		71.50
8	352	121.90	524.7		72.87
9	376	91.67	369.0		74.83
10	391	31.94	123.9		75.97
11	397	72.52	276.5		76.52
12	424	143.73	513.1		78.11
13	578	30.24	79.3		86.09
14	10,380	75.00	10.9	Upper Marker	113.00
15	12,774	0.00	0.0		115.51

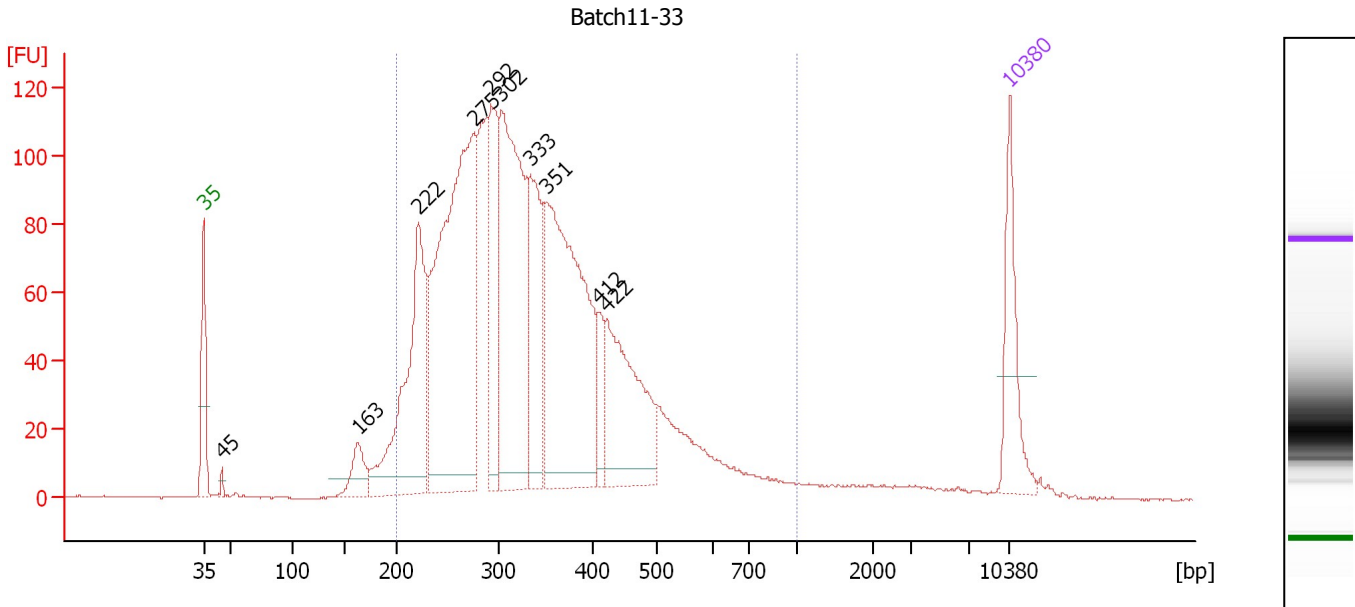
Region table for sample 8 : Batch11-32

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	359	1,278.58	1,260.4	5,879.3	96	26.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

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 Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : Batch11-33

Number of peaks found: 10 Corr. Area 1: 2,584.2
 Noise: 0.3

Peak table for sample 9 : Batch11-33

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	45	7.36	245.3		44.62
3	163	50.48	469.0		56.38
4	222	324.16	2,216.8		61.66
5	275	639.63	3,521.1		66.47
6	292	172.14	894.0		67.95
7	302	420.81	2,113.1		68.83
8	333	179.66	816.4		71.38
9	351	455.49	1,967.6		72.77
10	412	41.94	154.3		77.40
11	422	204.31	732.8		78.00
12	10,380	75.00	10.9	Upper Marker	113.00

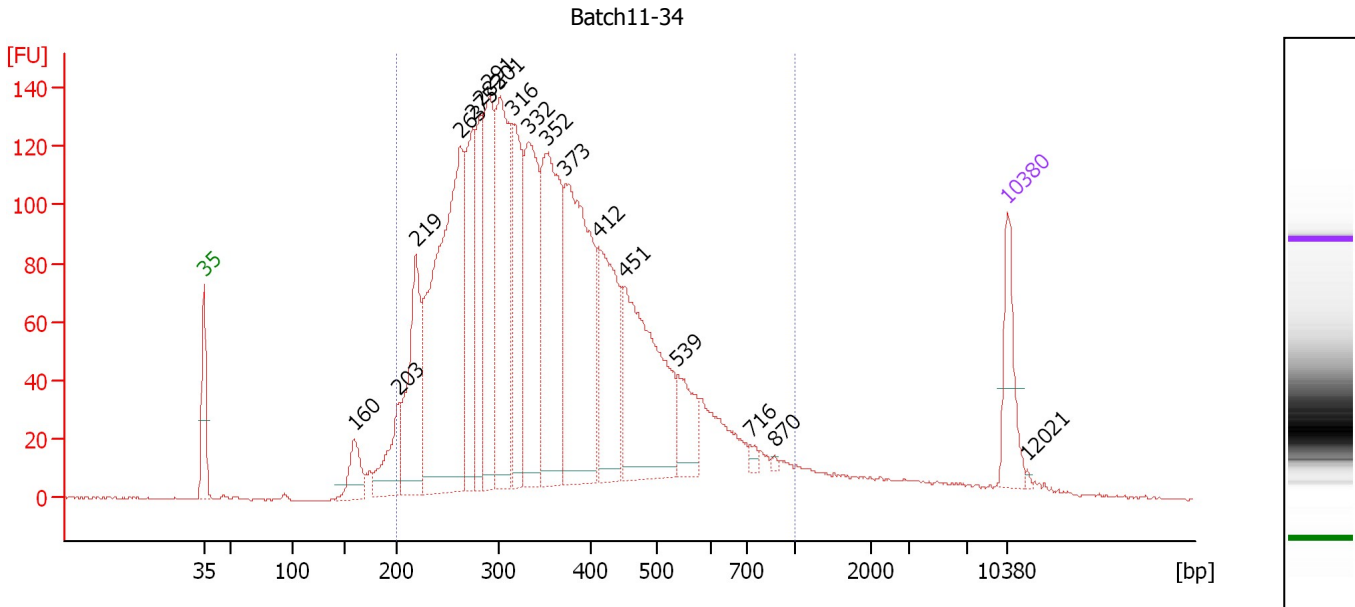
Region table for sample 9 : Batch11-33

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	344	2,795.58	2,584.2	13,647.7	94	31.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

Created: 10/22/2018 3:34:00 PM
 Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : Batch11-34

Number of peaks found: 18 Corr. Area 1: 3,504.8
 Noise: 0.3

Peak table for sample 10 : Batch11-34

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	160	72.37	685.1		56.11
3	203	103.67	772.9		60.01
4	219	253.20	1,749.5		61.45
5	263	761.06	4,387.8		65.36
6	275	231.03	1,274.2		66.43
7	282	188.50	1,011.0		67.12
8	291	281.43	1,466.4		67.87
9	301	343.82	1,732.4		68.75
10	316	219.71	1,054.3		69.96
11	332	311.97	1,423.9		71.26
12	352	392.74	1,689.4		72.89
13	373	456.44	1,856.6		74.51
14	412	232.85	856.7		77.40
15	451	365.77	1,229.0		79.63
16	539	84.54	237.6		84.27
17	716	9.48	20.1		90.60
18	870	3.51	6.1		92.69
19	10,380	75.00	10.9	Upper Marker	113.00
20	12,021	0.00	0.0		114.72

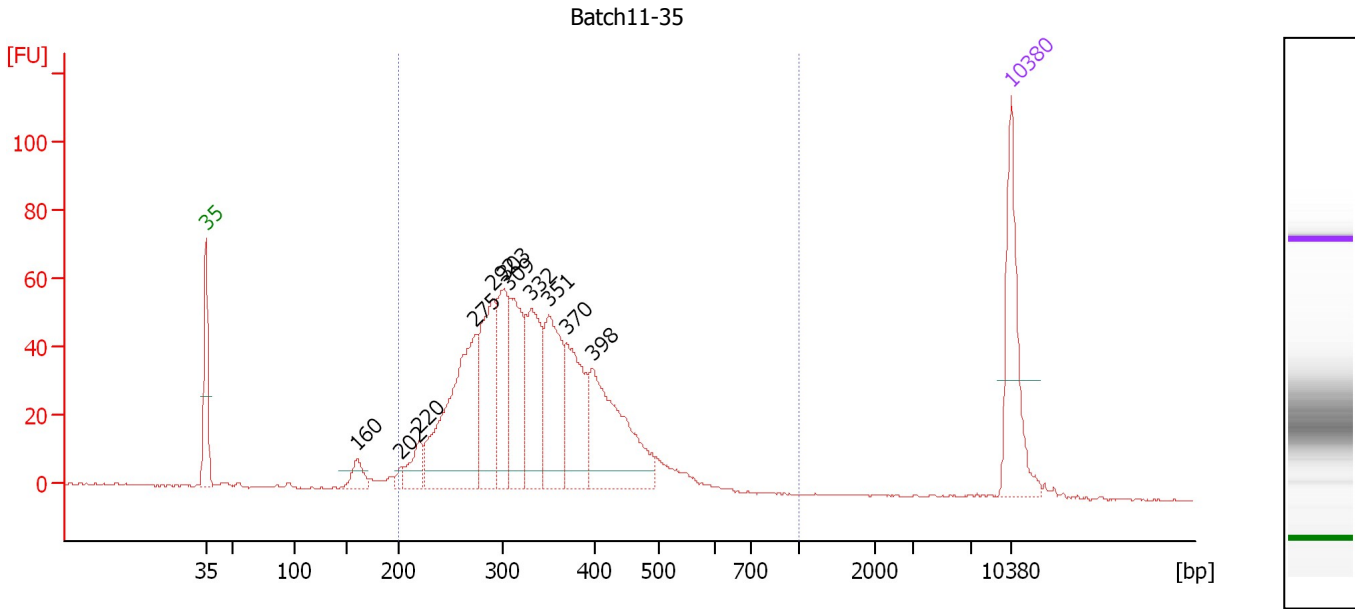
Region table for sample 10 : Batch11-34

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	370	4,660.01	3,504.8	21,721.3	93	34.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

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 Modified: 10/22/2018 4:30:48 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : Batch11-35

Number of peaks found: 11 Corr. Area 1: 1,125.3
 Noise: 0.3

Peak table for sample 11 : Batch11-35

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	160	22.39	211.8		56.12
3	202	6.89	51.8		59.87
4	220	30.66	210.9		61.54
5	275	214.00	1,181.0		66.41
6	292	118.87	617.6		67.94
7	303	93.86	469.8		68.91
8	309	108.94	535.0		69.38
9	332	106.47	485.6		71.28
10	351	114.52	494.0		72.81
11	370	98.31	402.2		74.34
12	398	145.36	553.3		76.56
13	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 11 : Batch11-35

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	1,103.48	1,125.3	5,220.3	96	22.9

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-22\2018-10-22_002.xad

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Gel Image

