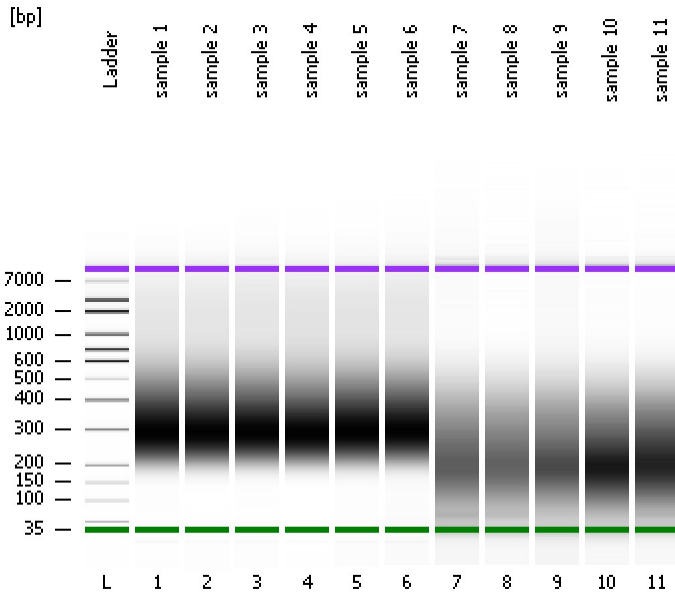


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
Modified: 8/3/2018 12:01:32 PM

Electrophoresis File Run Summary



Instrument Information:

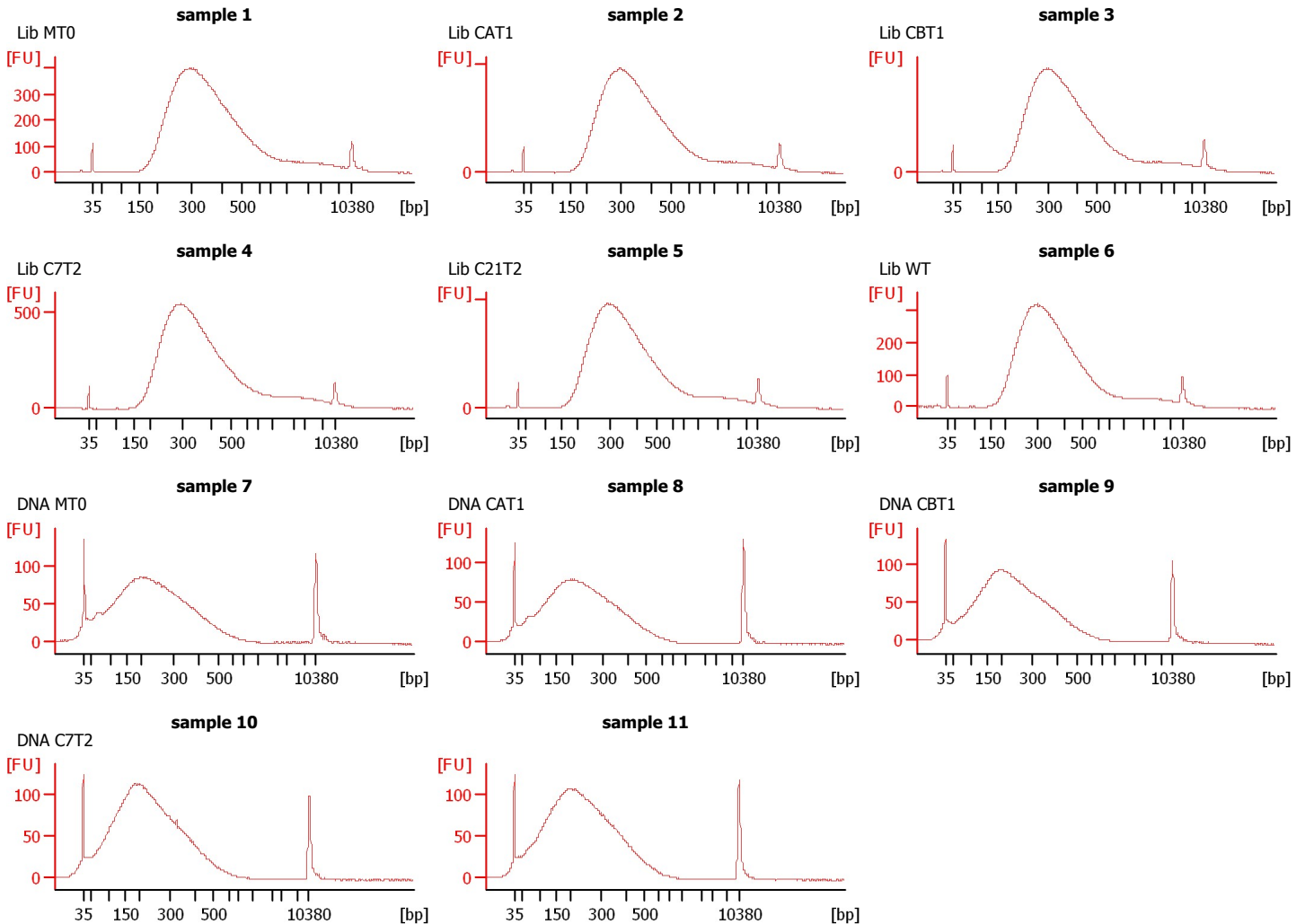
Instrument Name: DE34903152 Firmware: C.01.069
Serial#: DE34903152 Type: G2938C

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:



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
 Modified: 8/3/2018 12:01:32 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
sample 1	Lib MT0	<input type="checkbox"/>	✓			
sample 2	Lib CAT1	<input type="checkbox"/>	✓			
sample 3	Lib CBT1	<input type="checkbox"/>	✓			
sample 4	Lib C7T2	<input type="checkbox"/>	✓			
sample 5	Lib C21T2	<input type="checkbox"/>	✓			
sample 6	Lib WT	<input type="checkbox"/>	✓			
sample 7	DNA MT0	<input type="checkbox"/>	✓			
sample 8	DNA CAT1	<input type="checkbox"/>	✓			
sample 9	DNA CBT1	<input type="checkbox"/>	✓			
sample 10	DNA C7T2	<input type="checkbox"/>	✓			
sample 11		<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\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
Modified: 8/3/2018 12:01:32 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

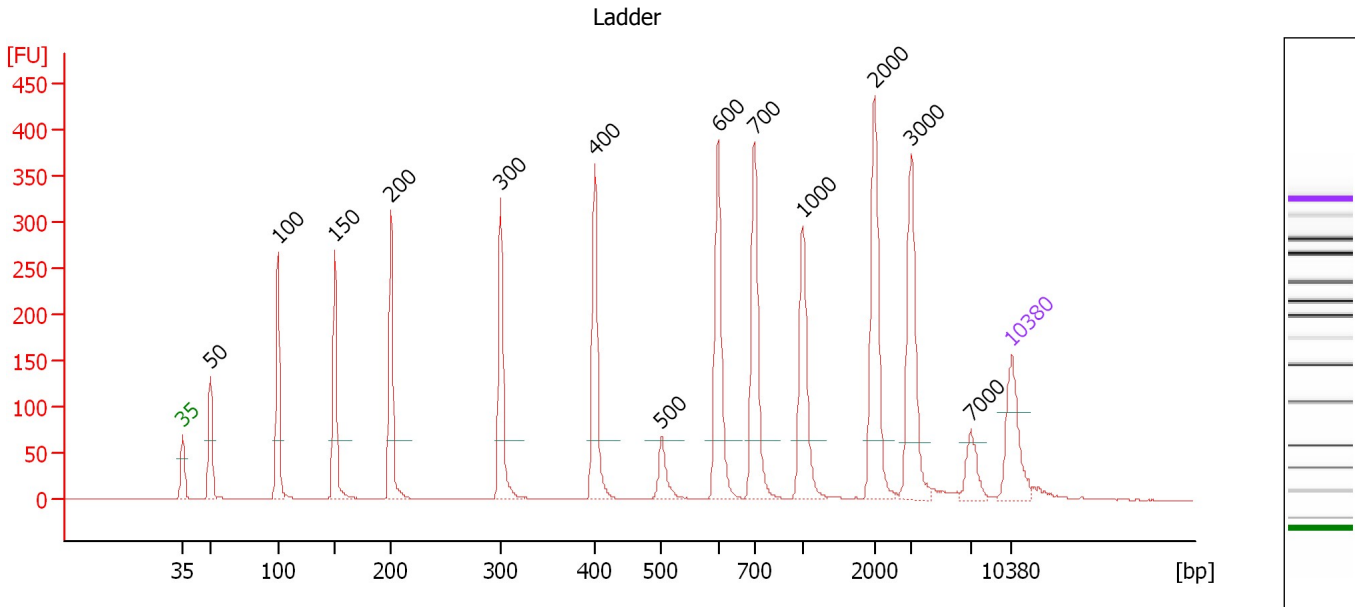
Ladder

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\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
 Modified: 8/3/2018 12:01:32 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.3

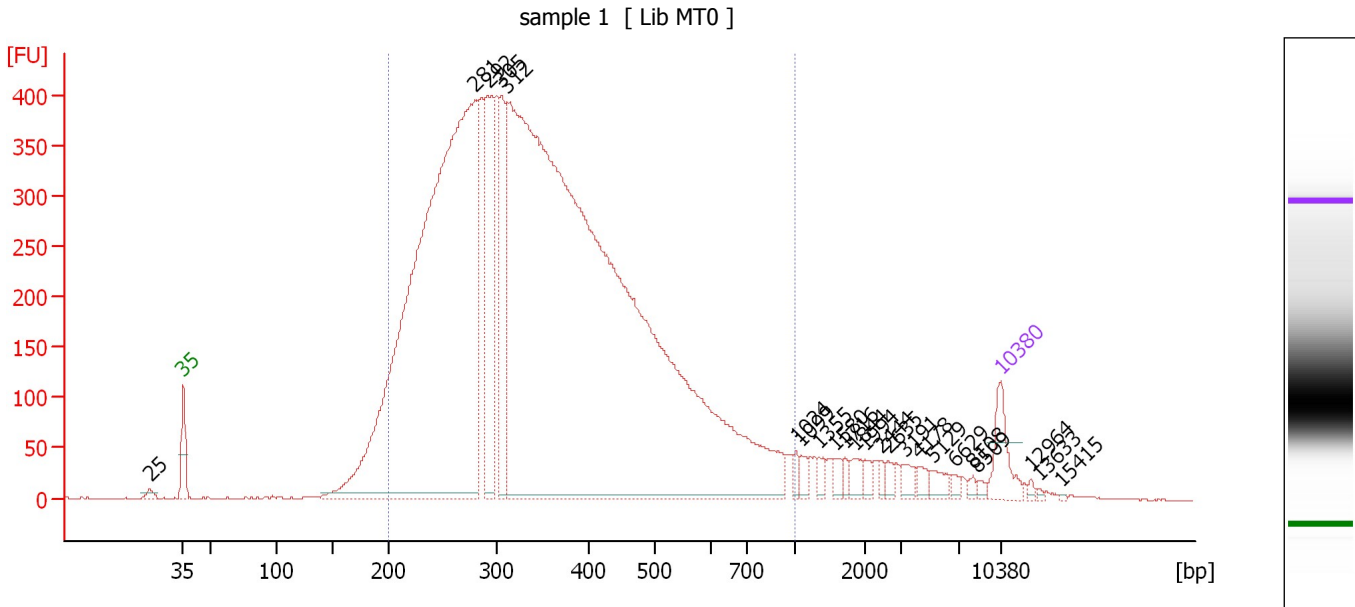
Peak table for Ladder

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	45.34
3	100	150.00	2,272.7	Ladder Peak	51.01
4	150	150.00	1,515.2	Ladder Peak	55.83
5	200	150.00	1,136.4	Ladder Peak	60.56
6	300	150.00	757.6	Ladder Peak	69.83
7	400	150.00	568.2	Ladder Peak	77.80
8	500	150.00	454.5	Ladder Peak	83.42
9	600	150.00	378.8	Ladder Peak	88.20
10	700	150.00	324.7	Ladder Peak	91.26
11	1,000	150.00	227.3	Ladder Peak	95.35
12	2,000	150.00	113.6	Ladder Peak	101.43
13	3,000	150.00	75.8	Ladder Peak	104.54
14	7,000	150.00	32.5	Ladder Peak	109.53
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
 Modified: 8/3/2018 12:01:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : sample 1

Number of peaks found: 23 Corr. Area 1: 11,288.2
 Noise: 0.7

Peak table for sample 1 : sample 1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	25	0.00	0.0		40.12
2	35	125.00	5,411.3	Lower Marker	43.00
3	281	3,295.58	17,789.2		68.04
4	292	456.75	2,369.8		69.09
5	305	346.76	1,722.3		70.23
6	312	5,740.67	27,883.8		70.78
7	1,024	18.39	27.2		95.50
8	1,099	21.03	29.0		95.96
9	1,355	19.34	21.6		97.51
10	1,580	20.68	19.8		98.88
11	1,716	14.20	12.5		99.70
12	1,844	25.61	21.0		100.48
13	1,994	19.09	14.5		101.39
14	2,444	12.76	7.9		102.81
15	2,635	15.01	8.6		103.40
16	3,191	25.09	11.9		104.78
17	4,178	18.20	6.6		106.01
18	5,129	26.87	7.9		107.20
19	6,629	11.28	2.6		109.07
20	8,108	9.02	1.7		110.67
21	8,509	8.10	1.4		111.08
22	10,380	75.00	10.9	Upper Marker	113.00
23	12,964	0.00	0.0		115.65
24	13,633	0.00	0.0		116.34
25	15,415	0.00	0.0		118.16

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
Modified: 8/3/2018 12:01:32 PM

Electropherogram Summary Continued ...

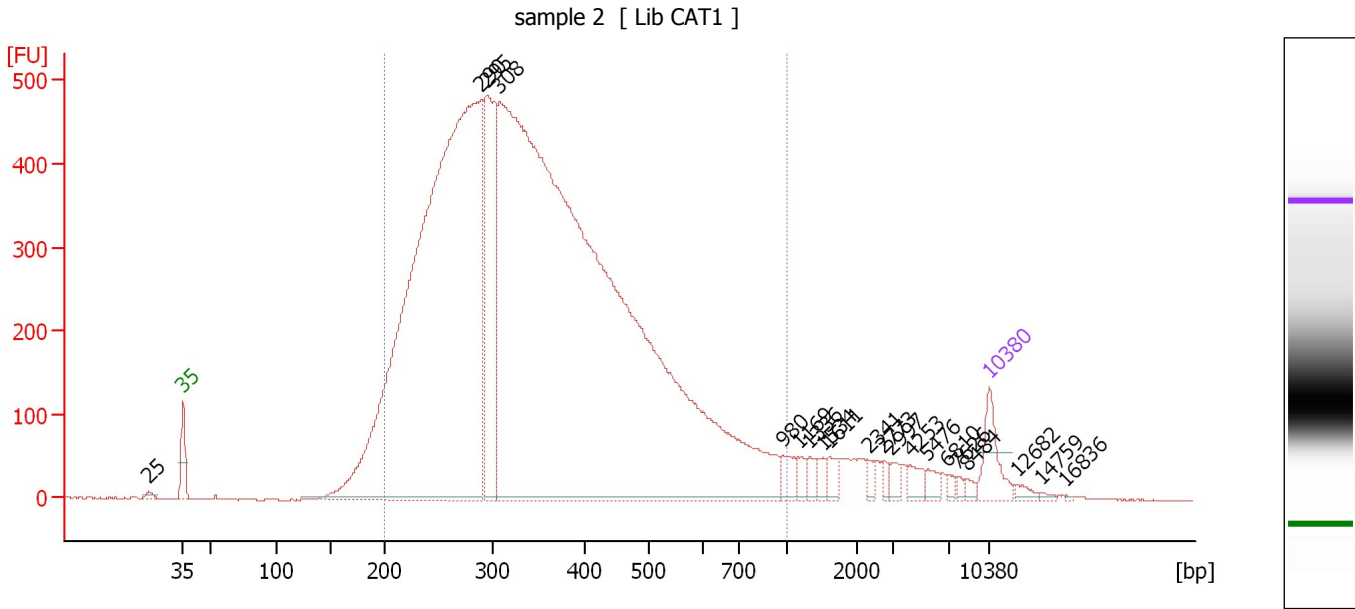
... Region table for sample 1 : sample 1

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	366	9,324.17	11,288.1	44,158.1	 91	35.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
 Modified: 8/3/2018 12:01:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : sample 2

Number of peaks found: 20 Corr. Area 1: 13,209.0
 Noise: 0.4

Peak table for sample 2 : sample 2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	25	0.00	0.0		40.04
2	35	125.00	5,411.3	Lower Marker	43.00
3	290	3,593.44	18,790.3		68.88
4	295	601.42	3,086.4		69.39
5	308	5,930.47	29,189.7		70.45
6	980	42.01	64.9		95.08
7	1,169	23.63	30.6		96.38
8	1,336	22.57	25.6		97.40
9	1,534	21.77	21.5		98.60
10	1,611	28.68	27.0		99.06
11	2,341	13.55	8.8		102.49
12	2,773	13.89	7.6		103.83
13	2,997	20.94	10.6		104.53
14	4,253	28.43	10.1		106.10
15	5,476	20.84	5.8		107.63
16	6,810	10.84	2.4		109.30
17	7,626	8.51	1.7		110.18
18	8,484	11.06	2.0		111.06
19	10,380	75.00	10.9	Upper Marker	113.00
20	12,682	0.00	0.0		115.36
21	14,759	0.00	0.0		117.49
22	16,836	0.00	0.0		119.62

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
Modified: 8/3/2018 12:01:32 PM

Electropherogram Summary Continued ...

... Region table for sample 2 :

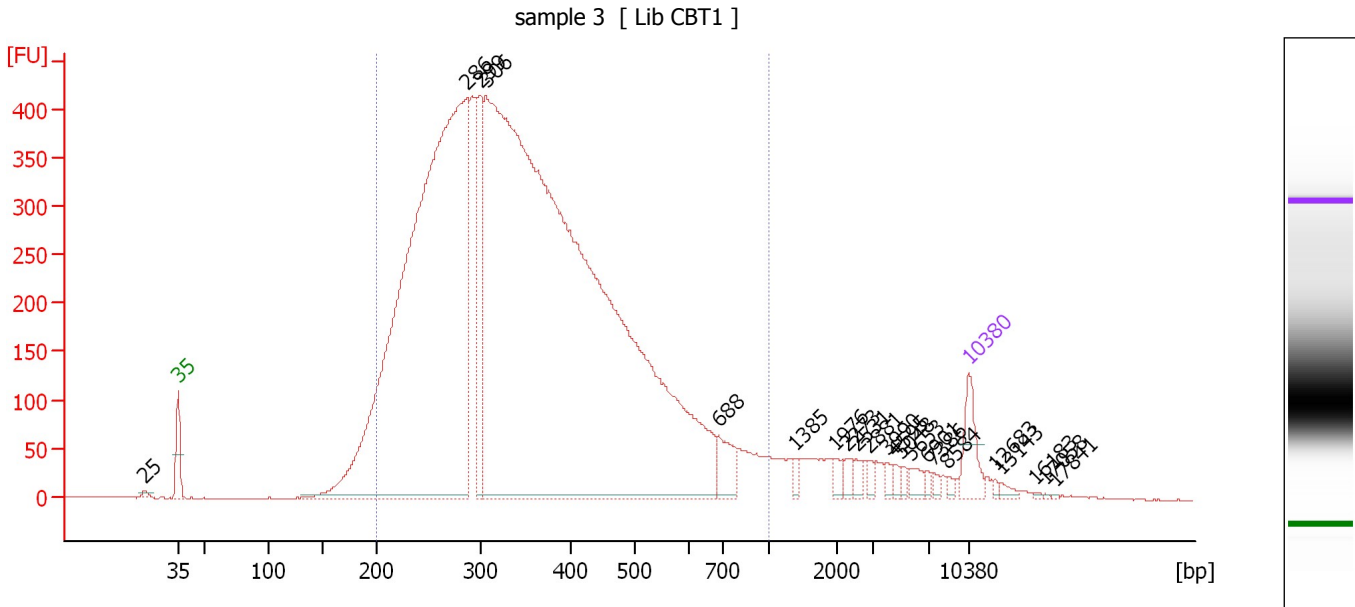
sample 2

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	365	9,129.66	13,209.0	43,272.2	 92	35.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
 Modified: 8/3/2018 12:01:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : sample 3

Number of peaks found: 22 Corr. Area 1: 11,251.8
 Noise: 0.6

Peak table for sample 3 : sample 3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	25	0.00	0.0		40.07
2	35	125.00	5,411.3	Lower Marker	43.00
3	286	3,408.63	18,043.0		68.55
4	299	341.50	1,730.7		69.73
5	306	5,791.91	28,686.3		70.30
6	688	82.60	181.9		90.89
7	1,385	15.96	17.5		97.70
8	1,976	19.66	15.1		101.29
9	2,273	18.24	12.2		102.28
10	2,531	18.83	11.3		103.08
11	2,881	13.12	6.9		104.17
12	3,990	13.91	5.3		105.77
13	4,595	11.49	3.8		106.53
14	5,048	11.68	3.5		107.10
15	5,653	23.07	6.2		107.85
16	6,901	9.17	2.0		109.41
17	7,386	9.57	2.0		109.93
18	8,584	8.07	1.4		111.16
19	10,380	75.00	10.9	Upper Marker	113.00
20	12,683	0.00	0.0		115.36
21	13,143	0.00	0.0		115.83
22	16,183	0.00	0.0		118.95
23	17,058	0.00	0.0		119.85
24	17,841	0.00	0.0		120.65

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
Modified: 8/3/2018 12:01:32 PM

Electropherogram Summary Continued ...

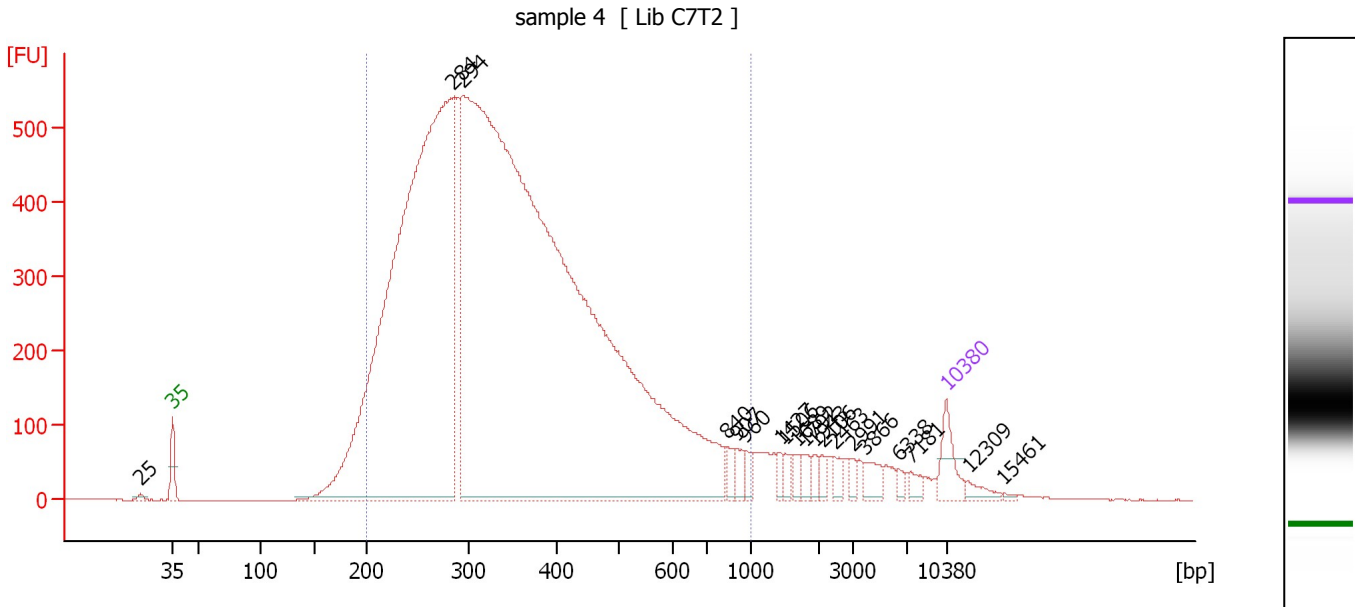
... Region table for sample 3 : sample 3

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	366	9,060.21	11,251.8	42,803.4	 92	35.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
 Modified: 8/3/2018 12:01:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : sample 4

Number of peaks found: 19 Corr. Area 1: 14,477.9
 Noise: 0.5

Peak table for sample 4 : sample 4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	25	0.00	0.0		40.11
2	35	125.00	5,411.3	Lower Marker	43.00
3	284	3,970.28	21,157.7		68.38
4	294	7,621.77	39,321.5		69.24
5	840	38.39	69.2		93.17
6	907	35.76	59.7		94.09
7	960	33.16	52.3		94.81
8	1,427	22.12	23.5		97.95
9	1,506	24.62	24.8		98.43
10	1,689	25.16	22.6		99.54
11	1,792	29.13	24.6		100.17
12	1,943	19.32	15.1		101.08
13	2,106	21.46	15.4		101.76
14	2,463	26.12	16.1		102.87
15	2,991	18.17	9.2		104.51
16	3,866	41.86	16.4		105.62
17	6,338	14.12	3.4		108.71
18	7,181	22.98	4.8		109.72
19	10,380	75.00	10.9	Upper Marker	113.00
20	12,309	0.00	0.0		114.98
21	15,461	0.00	0.0		118.21

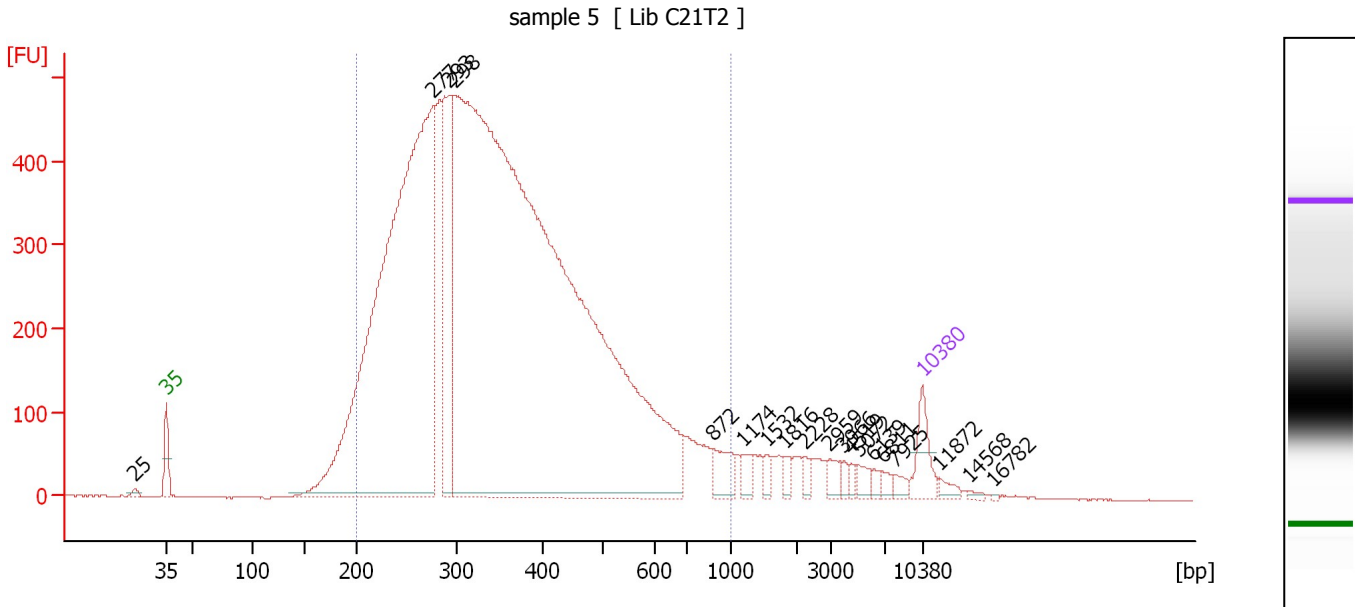
Region table for sample 4 : sample 4

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	364	10,650.00	14,477.9	50,761.5	91	36.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
 Modified: 8/3/2018 12:01:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : sample 5

Number of peaks found: 19 Corr. Area 1: 13,084.2
 Noise: 0.7

Peak table for sample 5 : sample 5

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	25	0.00	0.0		40.09
2	35	125.00	5,411.3	Lower Marker	43.00
3	277	3,132.86	17,145.7		67.68
4	293	512.97	2,654.3		69.16
5	298	6,590.25	33,491.5		69.66
6	872	78.04	135.7		93.60
7	1,174	34.58	44.6		96.41
8	1,532	19.52	19.3		98.59
9	1,816	21.96	18.3		100.31
10	2,228	19.48	13.2		102.14
11	2,959	30.85	15.8		104.41
12	3,966	13.65	5.2		105.74
13	4,519	13.00	4.4		106.43
14	5,072	25.71	7.7		107.13
15	6,139	13.28	3.3		108.46
16	6,811	16.42	3.7		109.30
17	7,925	18.59	3.6		110.48
18	10,380	75.00	10.9	Upper Marker	113.00
19	11,872	0.00	0.0		114.53
20	14,568	0.00	0.0		117.29
21	16,782	0.00	0.0		119.57

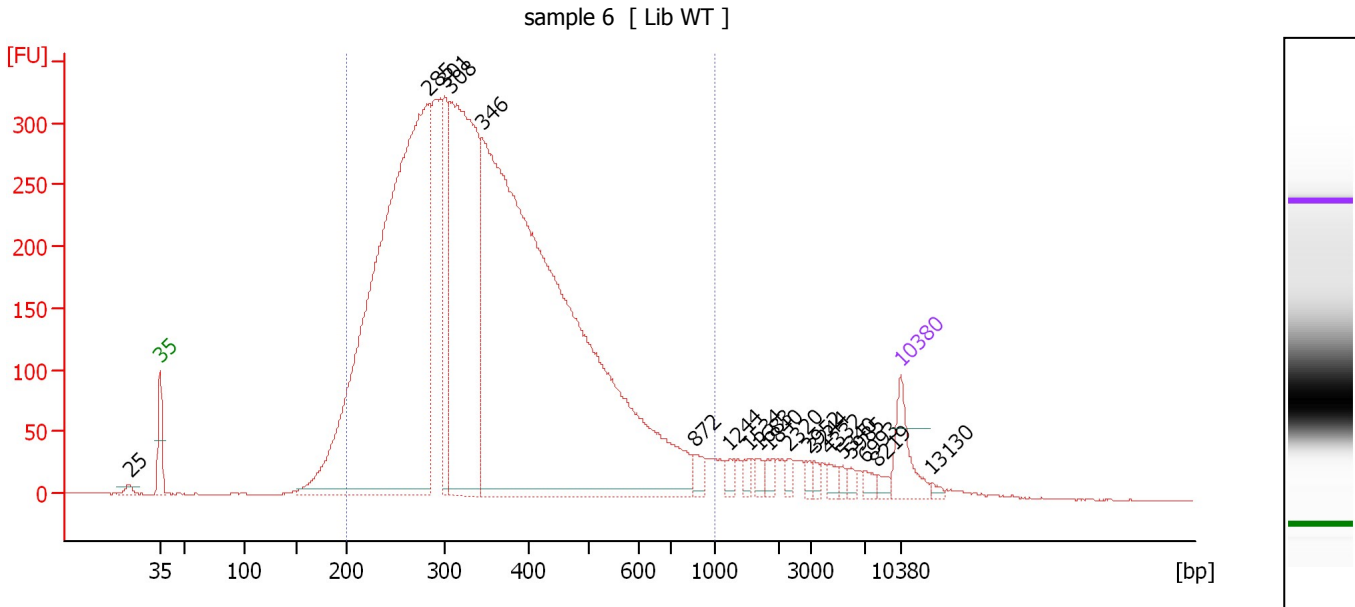
Region table for sample 5 : sample 5

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	367	9,591.31	13,084.2	45,216.4	91	35.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
 Modified: 8/3/2018 12:01:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : sample 6

Number of peaks found: 19 Corr. Area 1: 8,565.0
 Noise: 0.4

Peak table for sample 6 : sample 6

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	25	0.00	0.0		40.13
2	35	125.00	5,411.3	Lower Marker	43.00
3	285	2,560.06	13,613.6		68.43
4	301	272.90	1,371.8		69.94
5	308	1,149.68	5,660.5		70.45
6	346	3,288.09	14,412.9		73.47
7	872	26.66	46.3		93.61
8	1,244	19.54	23.8		96.83
9	1,534	17.65	17.4		98.60
10	1,683	17.18	15.5		99.50
11	1,840	16.67	13.7		100.46
12	2,320	11.71	7.7		102.42
13	2,952	11.02	5.7		104.39
14	3,244	13.15	6.1		104.84
15	4,332	16.46	5.8		106.20
16	5,340	9.18	2.6		107.46
17	5,985	12.60	3.2		108.27
18	6,993	14.22	3.1		109.53
19	8,219	11.78	2.2		110.78
20	10,380	75.00	10.9	Upper Marker	113.00
21	13,130	0.00	0.0		115.82

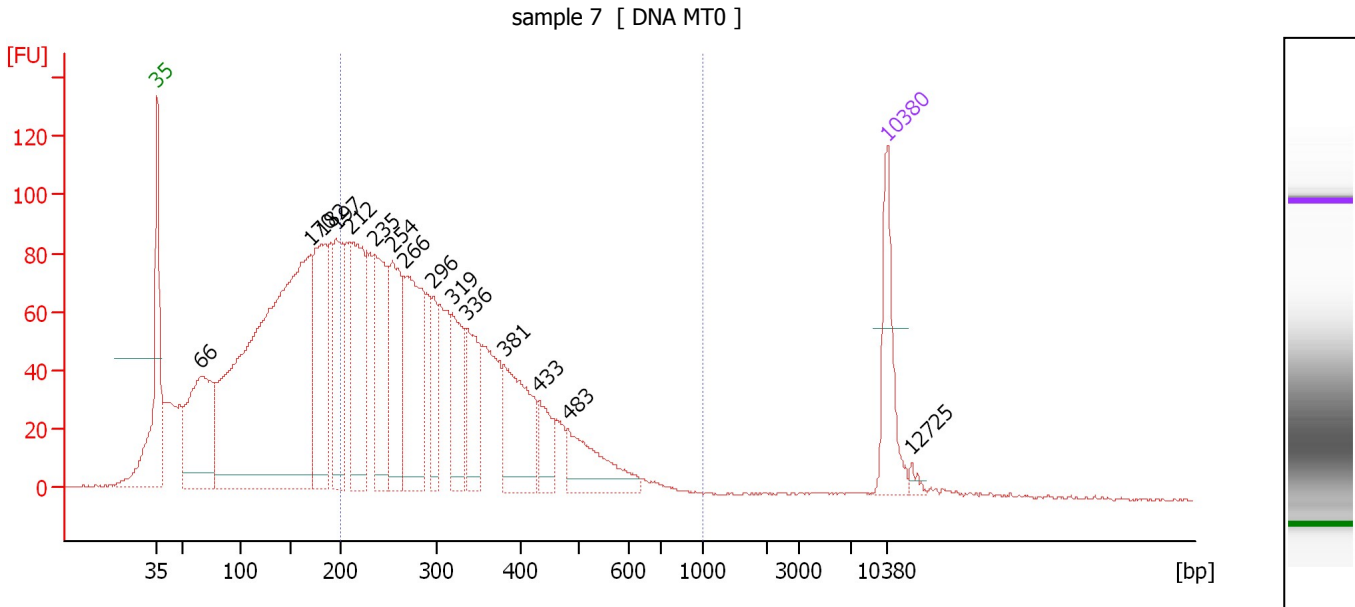
Region table for sample 6 : sample 6

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	365	7,323.11	8,565.0	34,480.8	92	34.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
 Modified: 8/3/2018 12:01:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : sample 7

Number of peaks found: 15 Corr. Area 1: 1,988.7
 Noise: 0.4

Peak table for sample 7 : sample 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	66	286.35	6,583.4		47.14
3	170	1,107.80	9,863.3		57.74
4	182	243.79	2,028.6		58.86
5	197	163.80	1,261.8		60.24
6	212	230.81	1,653.2		61.63
7	235	184.50	1,188.3		63.83
8	254	182.23	1,086.9		65.57
9	266	233.49	1,329.2		66.69
10	296	63.45	324.9		69.45
11	319	110.79	526.1		71.35
12	336	97.66	439.9		72.73
13	381	148.39	589.6		76.31
14	433	51.28	179.6		79.64
15	483	95.26	299.0		82.45
16	10,380	75.00	10.9	Upper Marker	113.00
17	12,725	0.00	0.0		115.40

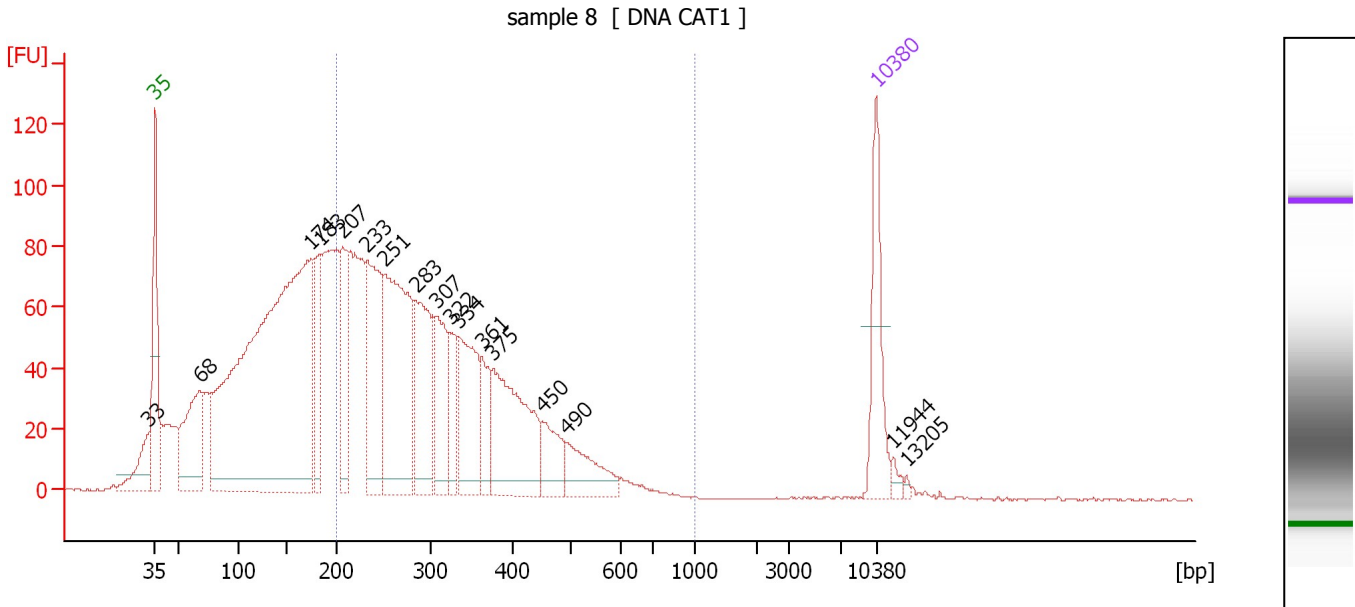
Region table for sample 7 : sample 7

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	319	1,966.55	1,988.7	10,373.2	52	31.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

Created: 8/3/2018 11:20:11 AM
 Modified: 8/3/2018 12:01:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : sample 8

Number of peaks found: 17 Corr. Area 1: 1,787.2
 Noise: 0.3

Peak table for sample 8 : sample 8

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	33	0.00	0.0		42.33
2	35	125.00	5,411.3	Lower Marker	43.00
3	68	179.29	4,014.0		47.35
4	174	1,095.30	9,532.6		58.11
5	183	101.60	839.4		58.99
6	207	112.19	823.2		61.16
7	233	207.28	1,349.5		63.59
8	251	318.42	1,921.0		65.30
9	283	165.53	884.7		68.30
10	307	98.44	486.2		70.37
11	322	59.11	277.9		71.61
12	334	132.24	599.8		72.54
13	361	55.07	230.9		74.71
14	375	189.26	764.8		75.80
15	450	54.43	183.2		80.61
16	490	65.98	203.8		82.89
17	10,380	75.00	10.9	Upper Marker	113.00
18	11,944	0.00	0.0		114.60
19	13,205	0.00	0.0		115.90

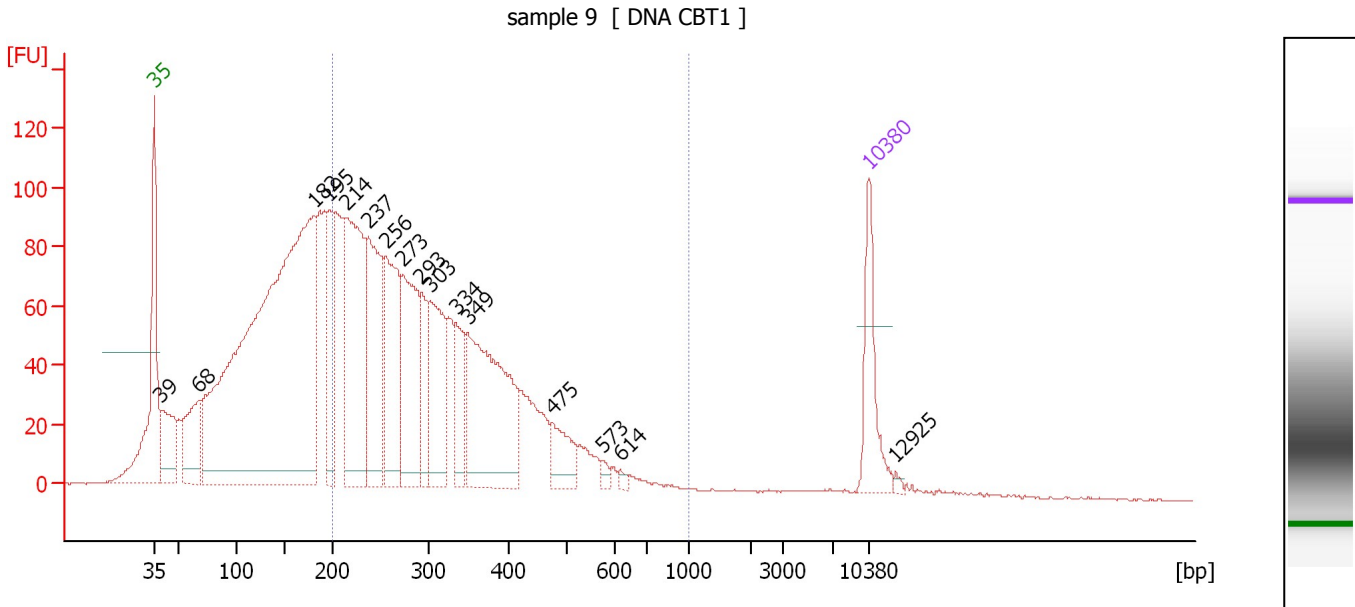
Region table for sample 8 : sample 8

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	313	1,755.05	1,787.2	9,345.6	52	29.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 9 : sample 9

Number of peaks found: 16 Corr. Area 1: 2,014.2
 Noise: 0.4

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	39	131.67	5,113.7		43.63
3	68	146.86	3,249.4		47.44
4	182	1,445.06	12,056.9		58.82
5	195	159.88	1,239.7		60.12
6	214	340.59	2,412.7		61.84
7	237	248.57	1,589.4		63.98
8	256	213.22	1,264.2		65.71
9	273	223.33	1,239.5		67.33
10	293	86.03	445.3		69.15
11	303	162.72	814.7		70.04
12	334	77.65	352.2		72.55
13	349	314.62	1,365.3		73.75
14	475	56.41	179.7		82.05
15	573	8.70	23.0		86.90
16	614	5.75	14.2		88.62
17	10,380	75.00	10.9	Upper Marker	113.00
18	12,925	0.00	0.0		115.61

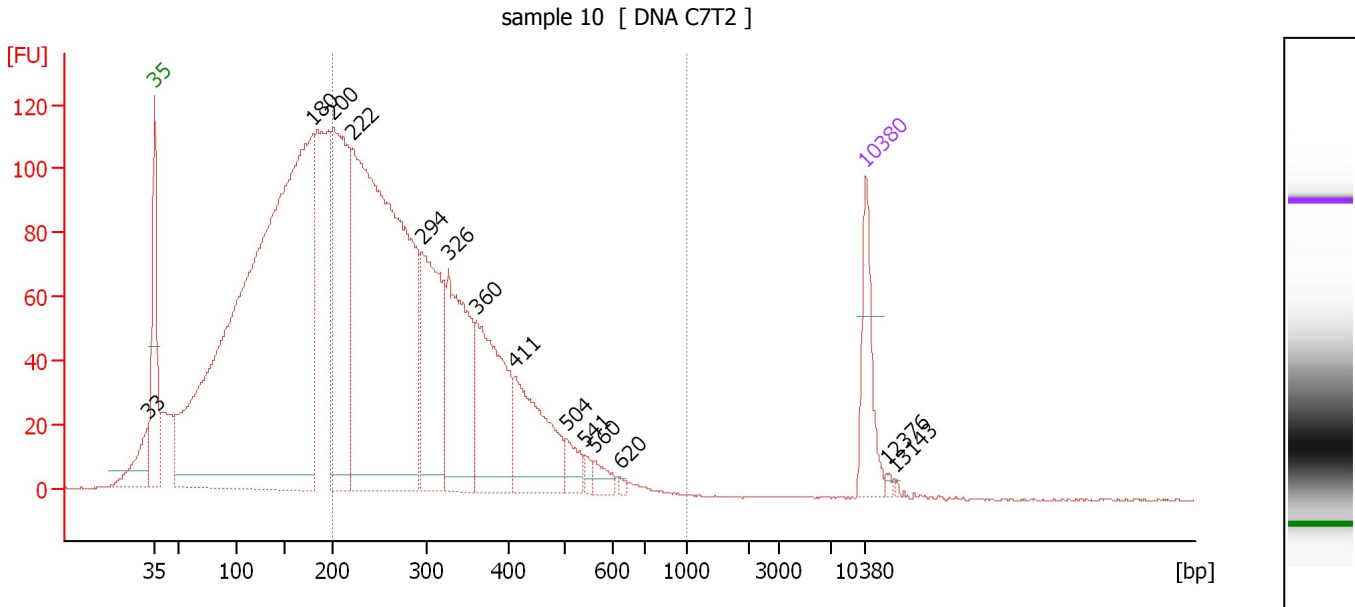
Region table for sample 9 : sample 9

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	319	2,195.58	2,014.2	11,647.2	53	32.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 10 : sample 10

Number of peaks found: 14 Corr. Area 1: 2,243.1
 Noise: 0.3

Peak table for sample 10 : sample 10

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	33	0.00	0.0		42.37
2	35	125.00	5,411.3	Lower Marker	43.00
3	180	2,097.61	17,686.2		58.64
4	200	470.82	3,562.5		60.58
5	222	1,278.76	8,737.6		62.57
6	294	306.24	1,577.3		69.29
7	326	270.96	1,258.8		71.91
8	360	252.15	1,060.1		74.64
9	411	177.14	652.9		78.42
10	504	33.31	100.2		83.61
11	541	9.63	27.0		85.40
12	560	20.49	55.4		86.29
13	620	4.09	10.0		88.81
14	10,380	75.00	10.9	Upper Marker	113.00
15	12,376	0.00	0.0		115.05
16	13,143	0.00	0.0		115.83

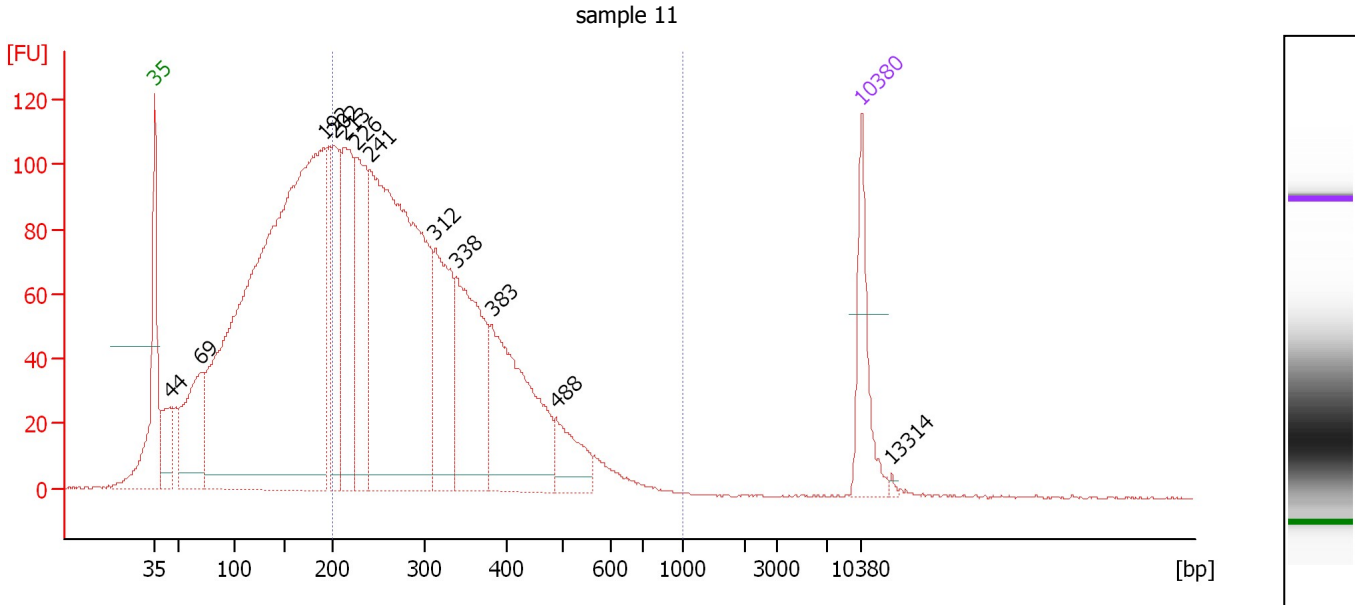
Region table for sample 10 : sample 10

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	309	2,676.06	2,243.1	14,434.5	50	29.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 11 : sample 11

Number of peaks found: 12 Corr. Area 1: 2,342.5
 Noise: 0.3

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	44	88.98	3,031.7		44.48
3	69	205.76	4,493.2		47.54
4	192	1,676.62	13,240.2		59.79
5	202	204.07	1,531.0		60.74
6	213	235.31	1,671.2		61.79
7	226	237.58	1,589.7		63.01
8	241	916.15	5,754.1		64.38
9	312	205.77	997.8		70.82
10	338	263.43	1,179.9		72.88
11	383	276.51	1,094.8		76.42
12	488	67.31	209.0		82.75
13	10,380	75.00	10.9	Upper Marker	113.00
14	13,314	0.00	0.0		116.01

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	314	2,327.20	2,342.5	12,385.9	54	30.1

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-03\2018-08-03_001.xad

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Gel Image

