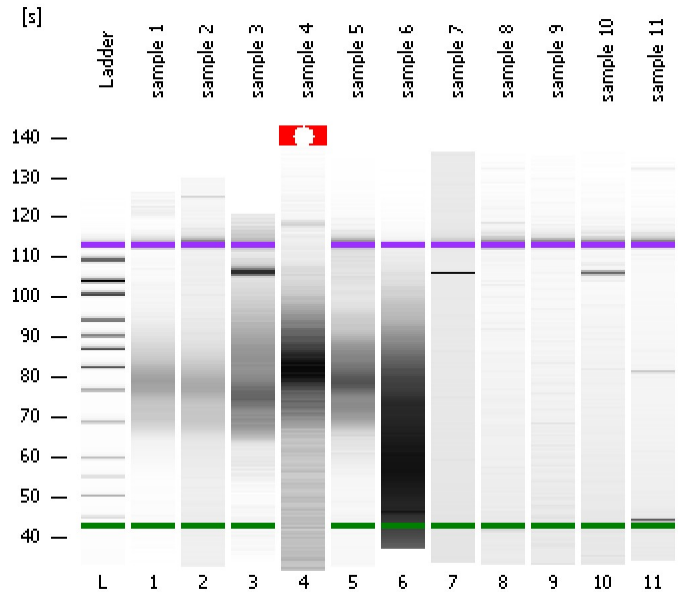


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

Created: 1/7/2015 11:48:58 AM
Modified: 1/7/2015 12:30:19 PM

Electrophoresis File Run Summary



Instrument Information:

Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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

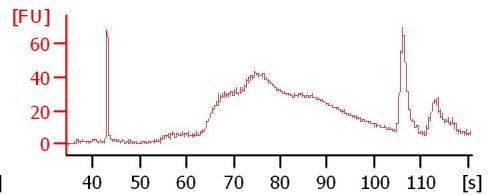
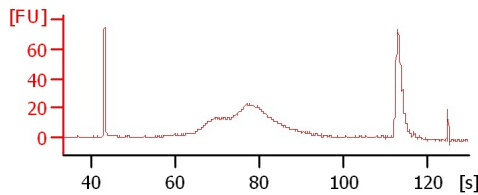
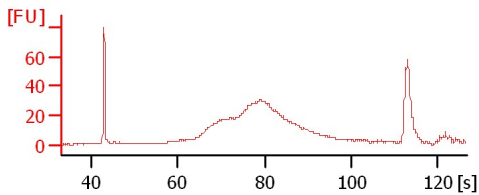
Chip Information:

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

sample 1

sample 2

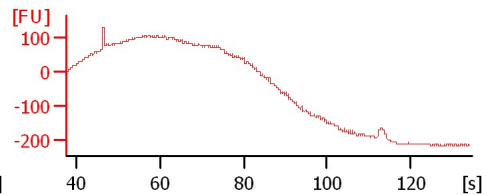
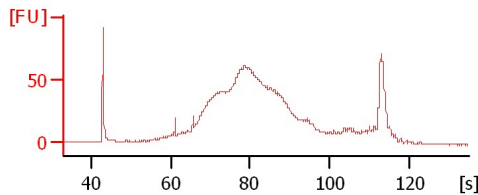
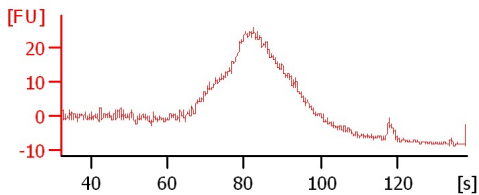
sample 3



sample 4

sample 5

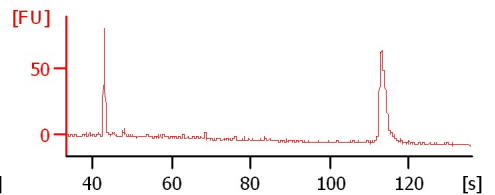
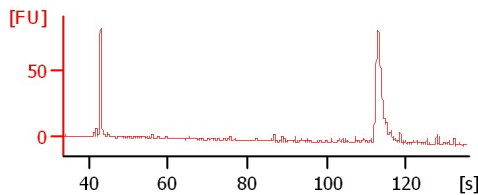
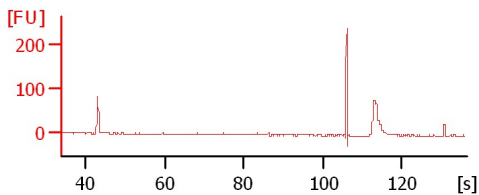
sample 6



sample 7

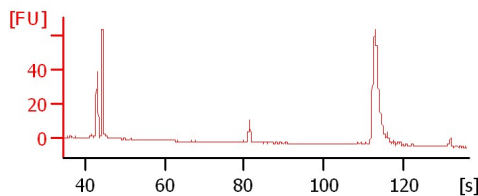
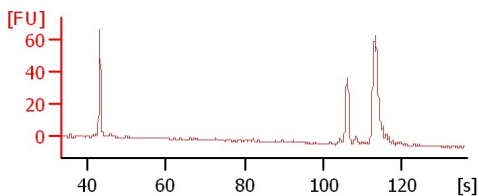
sample 8

sample 9



sample 10

sample 11



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

Created: 1/7/2015 11:48:58 AM
Modified: 1/7/2015 12:30:19 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
sample 1		<input type="checkbox"/>	✓			
sample 2		<input type="checkbox"/>	✓			
sample 3		<input type="checkbox"/>	✓			
sample 4		<input type="checkbox"/>	✓			
sample 5		<input type="checkbox"/>	✓			
sample 6		<input type="checkbox"/>	✓			
sample 7		<input type="checkbox"/>	✓			
sample 8		<input type="checkbox"/>	✓			
sample 9		<input type="checkbox"/>	✓			
sample 10		<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:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

Created: 1/7/2015 11:48:58 AM
Modified: 1/7/2015 12:30:19 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

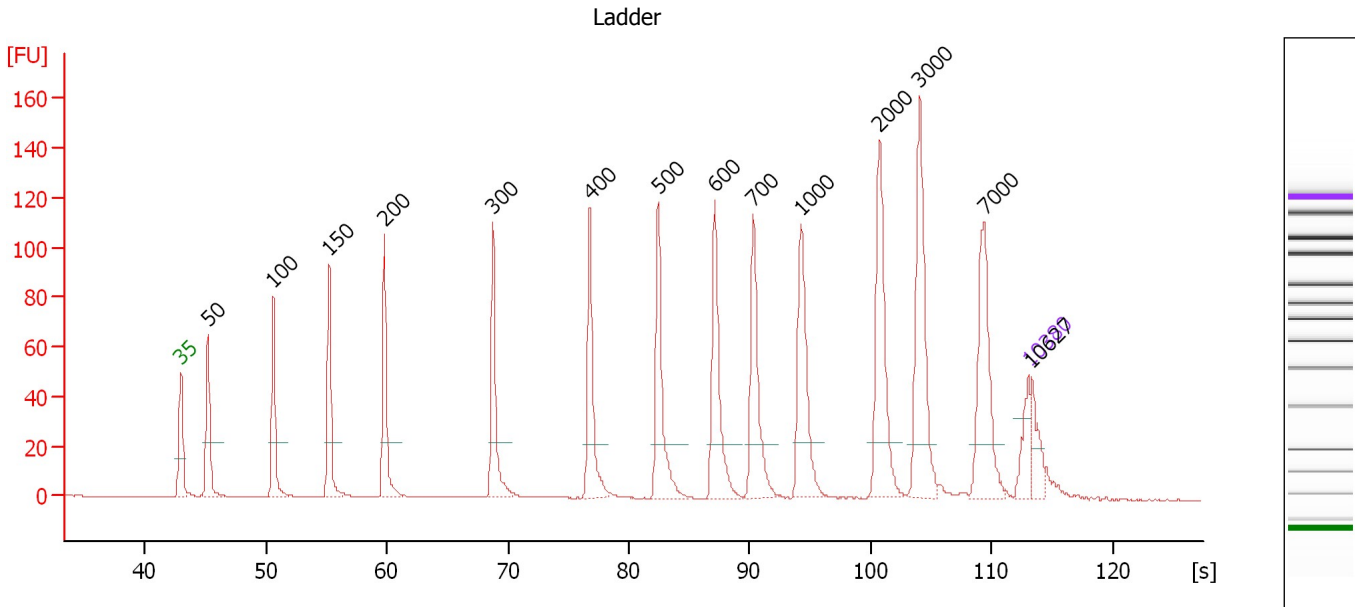
Ladder

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:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

Created: 1/7/2015 11:48:58 AM
 Modified: 1/7/2015 12:30:19 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

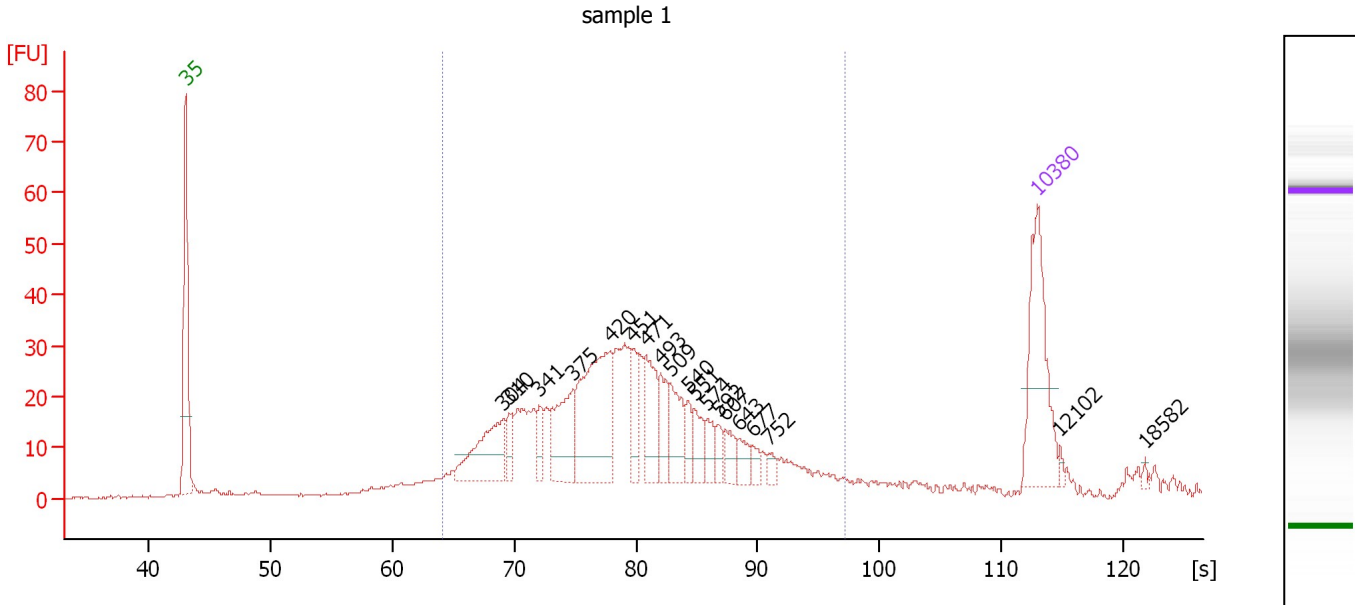
Peak table for Ladder

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	45.22
3	100	150.00	2,272.7	Ladder Peak	50.63
4	150	150.00	1,515.2	Ladder Peak	55.24
5	200	150.00	1,136.4	Ladder Peak	59.81
6	300	150.00	757.6	Ladder Peak	68.77
7	400	150.00	568.2	Ladder Peak	76.80
8	500	150.00	454.5	Ladder Peak	82.44
9	600	150.00	378.8	Ladder Peak	87.09
10	700	150.00	324.7	Ladder Peak	90.29
11	1,000	150.00	227.3	Ladder Peak	94.24
12	2,000	150.00	113.6	Ladder Peak	100.71
13	3,000	150.00	75.8	Ladder Peak	104.04
14	7,000	150.00	32.5	Ladder Peak	109.36
15	10,380	75.00	10.9	Upper Marker	113.00
16	10,627	0.00	0.0		113.27

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

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 Modified: 1/7/2015 12:30:19 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : sample 1

Number of peaks found: 19 Corr. Area 1: 655.7
 Noise: 0.2

Peak table for sample 1 : sample 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	304	72.58	361.2		69.13
3	310	16.78	82.0		69.57
4	341	17.34	77.1		72.04
5	375	63.49	256.7		74.78
6	420	129.07	466.0		77.91
7	451	25.96	87.2		79.68
8	471	47.37	152.2		80.82
9	493	25.81	79.4		82.02
10	509	36.95	110.0		82.85
11	540	13.94	39.1		84.31
12	551	21.17	58.3		84.80
13	574	14.38	37.9		85.90
14	592	9.39	24.0		86.74
15	607	14.47	36.1		87.31
16	643	12.17	28.7		88.46
17	677	7.78	17.4		89.56
18	752	7.41	14.9		90.98
19	10,380	75.00	10.9	Upper Marker	113.00
20	12,102	0.00	0.0		114.85
21	18,582	0.00	0.0		121.83

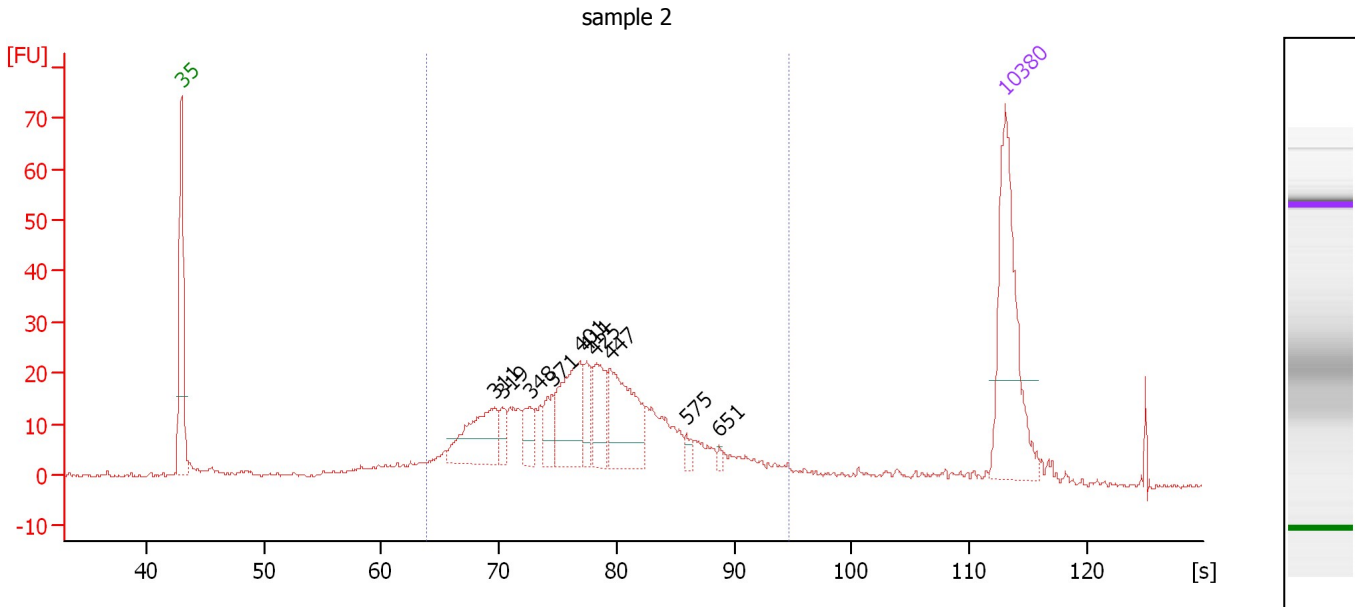
Region table for sample 1 : sample 1

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
247	469	1,463	655.7	90	36.6	875.05	3,227.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 2 : sample 2

Number of peaks found: 10 Corr. Area 1: 490.2
 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	35	125.00	5,411.3	Lower Marker	43.00
2	311	53.27	259.4		69.67
3	319	10.72	51.0		70.26
4	348	16.71	72.7		72.63
5	371	17.09	69.7		74.51
6	401	55.96	211.2		76.88
7	411	15.13	55.8		77.43
8	425	28.12	100.3		78.21
9	447	61.43	208.3		79.44
10	575	5.13	13.5		85.92
11	651	2.53	5.9		88.71
12	10,380	75.00	10.9	Upper Marker	113.00

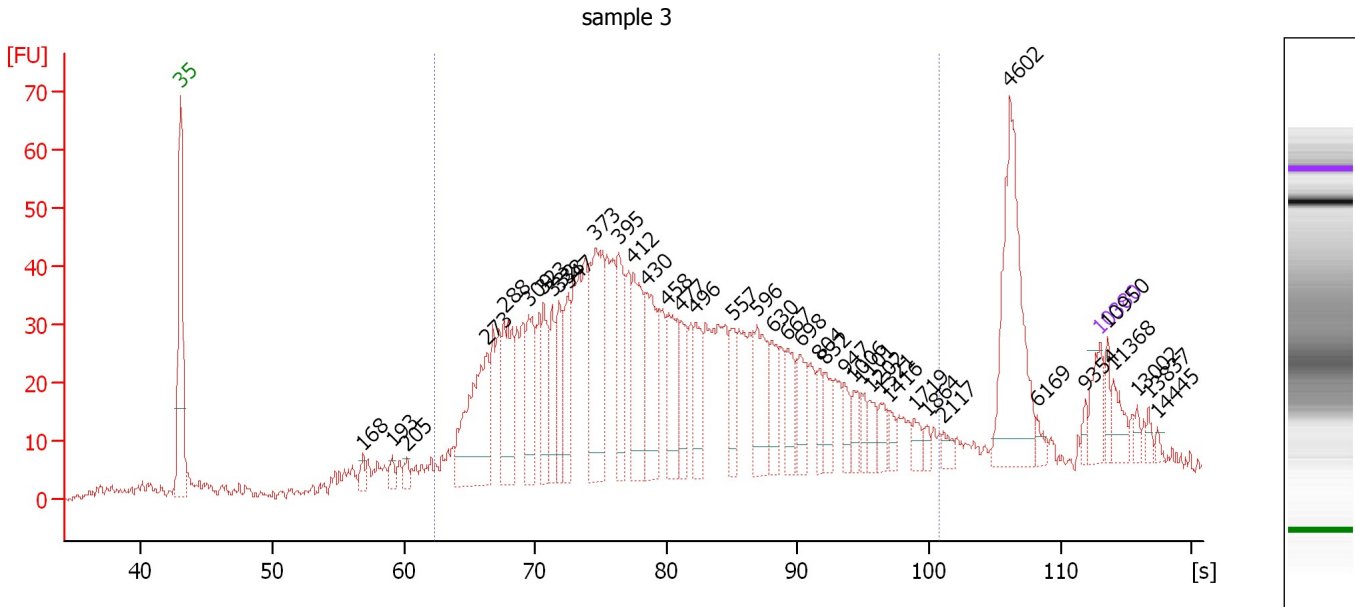
Region table for sample 2 : sample 2

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
244	438	1,059	490.2	85	30.2	465.32	1,788.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 3 : sample 3

Number of peaks found: 41 Corr. Area 1: 1,251.1
 Noise: 1.0

Peak table for sample 3 : sample 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	168	39.64	358.1		56.86
3	193	37.67	296.1		59.15
4	205	32.56	240.8		60.25
5	273	458.77	2,542.6		66.39
6	288	282.40	1,488.1		67.66
7	309	201.55	988.3		69.50
8	323	176.39	826.8		70.64
9	332	139.40	636.4		71.34
10	338	144.60	648.1		71.83
11	347	159.11	695.4		72.52
12	373	425.70	1,730.8		74.61
13	395	219.32	842.2		76.36
14	412	259.19	953.6		77.47
15	430	250.97	884.4		78.49
16	458	196.01	649.1		80.04
17	477	128.63	408.5		81.15
18	496	123.81	378.2		82.21
19	557	103.18	280.9		85.07
20	596	173.02	439.7		86.91
21	630	103.54	248.9		88.06
22	667	96.14	218.3		89.24
23	698	106.26	230.6		90.23
24	804	55.20	104.0		91.66
25	832	66.93	121.9		92.02

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad


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 Modified: 1/7/2015 12:30:19 PM

Electropherogram Summary Continued ...

... Peak table for sample 3 : sample 3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	947	50.75	81.2		93.54
27	1,006	46.49	70.0		94.27
28	1,101	35.07	48.3		94.89
29	1,202	43.83	55.3		95.54
30	1,321	43.03	49.3		96.32
31	1,416	29.29	31.3		96.93
32	1,719	29.46	26.0		98.89
33	1,864	17.96	14.6		99.83
34	2,117	22.50	16.1		101.10
35	4,602	415.50	136.8		106.17
36	6,169	21.58	5.3		108.26
37	9,354	18.06	2.9		111.90
38	10,380	75.00	10.9	Upper Marker	113.00
39	10,950	0.00	0.0		113.61
40	11,368	0.00	0.0		114.06
41	13,002	0.00	0.0		115.82
42	13,837	0.00	0.0		116.72
43	14,445	0.00	0.0		117.38

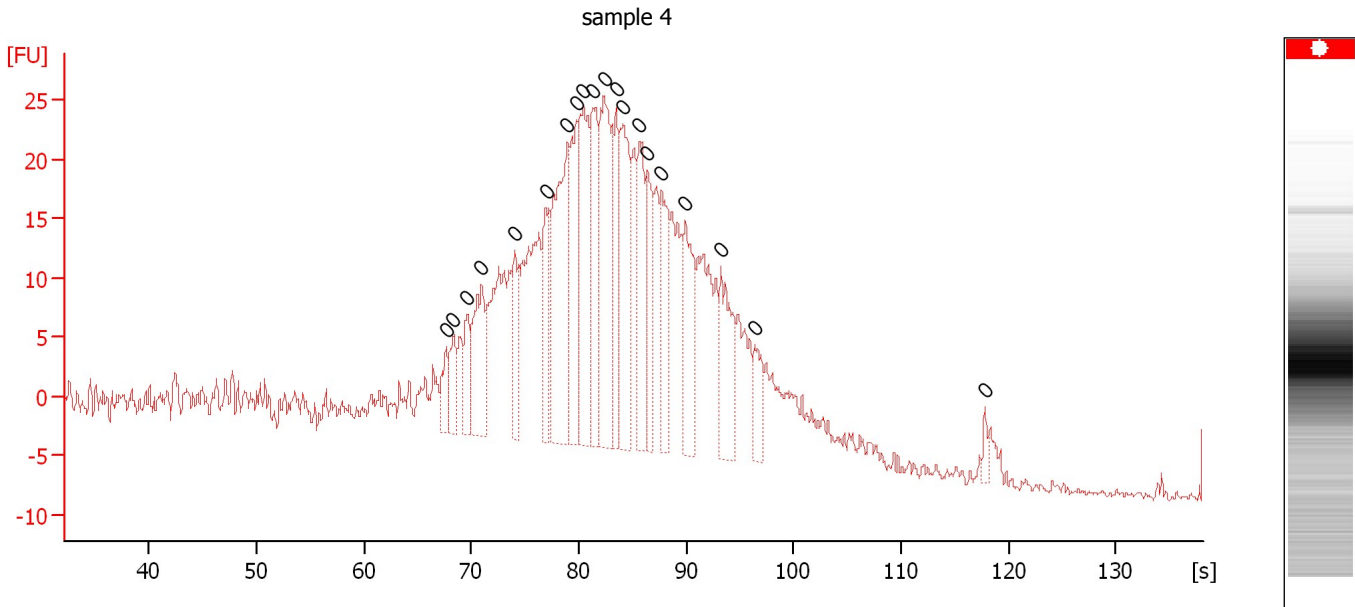
Region table for sample 3 : sample 3

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
228	539	1,999	1,251.1	81	59.2	6,888.52	25,553.8	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 4 : sample 4

Number of peaks found: 0 Noise: 0.4

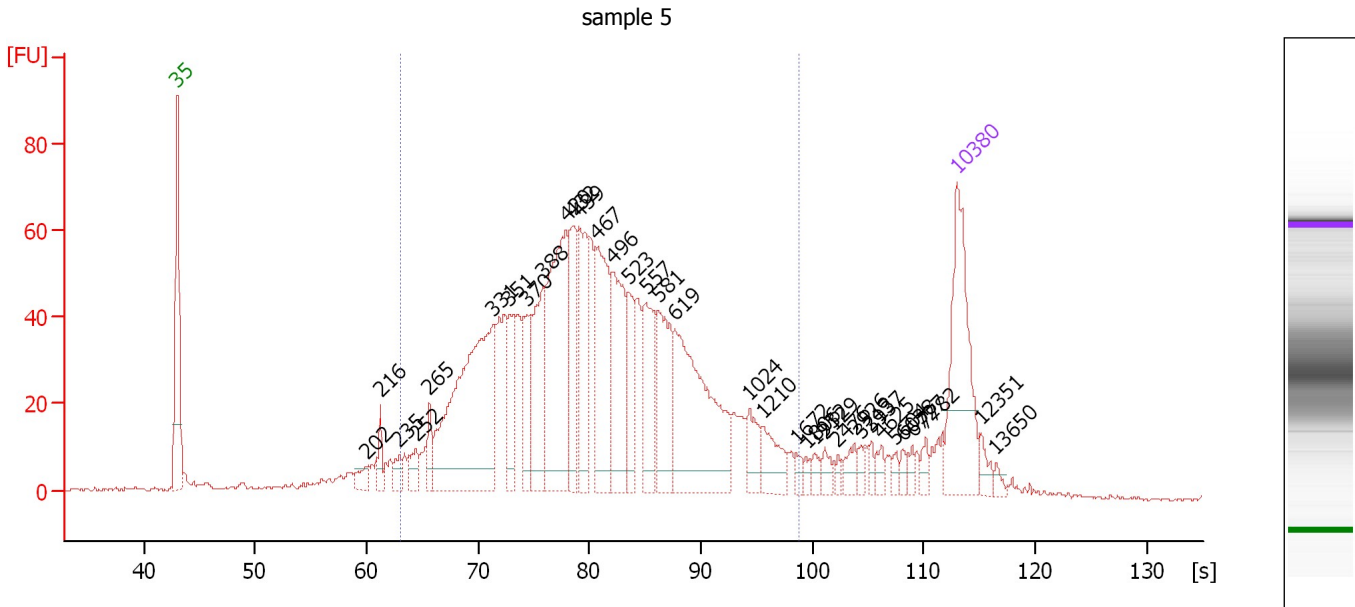
Peak table for sample 4 : sample 4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	0	0.00	0.0		67.65
2	0	0.00	0.0		68.30
3	0	0.00	0.0		69.60
4	0	0.00	0.0		70.95
5	0	0.00	0.0		74.05
6	0	0.00	0.0		76.95
7	0	0.00	0.0		78.95
8	0	0.00	0.0		79.80
9	0	0.00	0.0		80.40
10	0	0.00	0.0		81.30
11	0	0.00	0.0		82.40
12	0	0.00	0.0		83.45
13	0	0.00	0.0		84.05
14	0	0.00	0.0		85.65
15	0	0.00	0.0		86.40
16	0	0.00	0.0		87.70
17	0	0.00	0.0		89.95
18	0	0.00	0.0		93.25
19	0	0.00	0.0		96.40
20	0	0.00	0.0		117.80

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 5 : sample 5

Number of peaks found: 35 Corr. Area 1: 1,521.6
 Noise: 0.5

Peak table for sample 5 : sample 5

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	202	11.79	88.6		59.97
3	216	13.45	94.5		61.22
4	235	10.66	68.7		62.96
5	252	14.73	88.7		64.45
6	265	15.96	91.2		65.66
7	331	215.06	983.1		71.30
8	351	40.27	174.0		72.84
9	370	35.64	145.8		74.43
10	388	71.43	279.0		75.83
11	420	136.82	494.1		77.90
12	432	55.98	196.5		78.58
13	439	65.08	224.5		79.01
14	467	87.33	283.1		80.60
15	496	74.61	228.0		82.19
16	523	38.40	111.3		83.50
17	557	53.49	145.5		85.09
18	581	55.80	145.6		86.20
19	619	134.61	329.7		87.69
20	1,024	17.72	26.2		94.39
21	1,210	22.36	28.0		95.60
22	1,672	4.89	4.4		98.59
23	1,806	4.07	3.4		99.45
24	1,932	4.90	3.8		100.27
25	2,129	6.40	4.6		101.14

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad


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Electropherogram Summary Continued ...

... Peak table for sample 5 : sample 5

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,476	3.70	2.3		102.30
27	2,926	8.11	4.2		103.79
28	3,213	4.55	2.1		104.32
29	3,937	4.47	1.7		105.29
30	4,625	5.83	1.9		106.20
31	5,604	3.96	1.1		107.50
32	6,038	4.78	1.2		108.08
33	6,727	4.05	0.9		109.00
34	7,782	5.93	1.2		110.20
35	10,380	75.00	10.9	Upper Marker	113.00
36	12,351	0.00	0.0		115.12
37	13,650	0.00	0.0		116.52

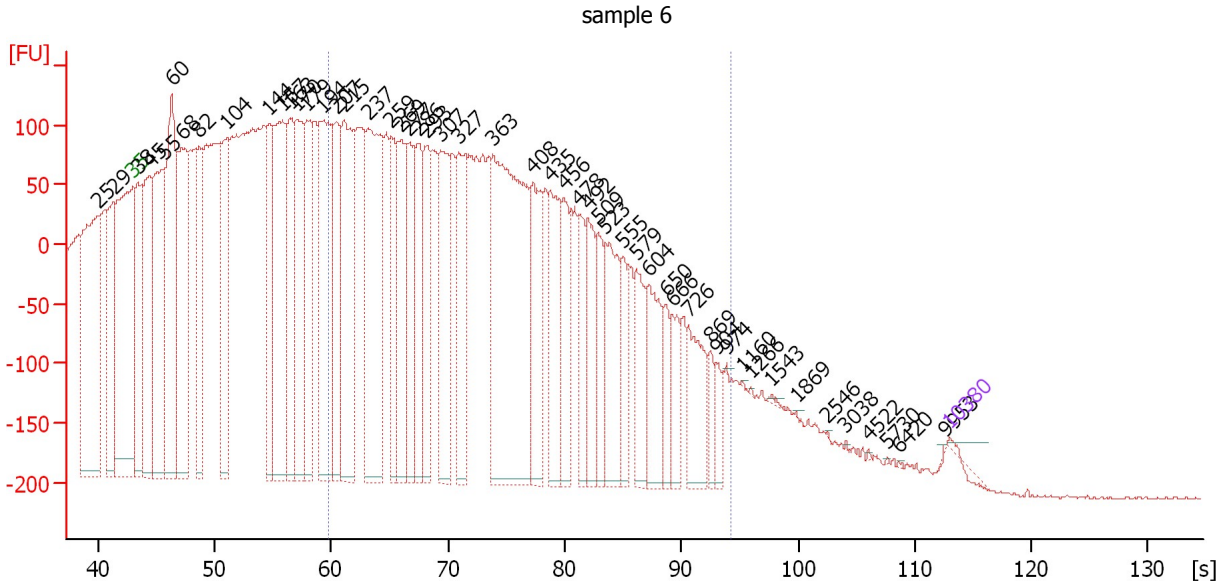
Region table for sample 5 : sample 5

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
236	514	1,699	1,521.6	86	45.9	1,397.85	4,966.9	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 6 : sample 6

Number of peaks found: 52 Corr. Area 1: 5,933.5
 Noise: 1.1

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		40.11
2	29	0.00	0.0		41.35
3	35	125.00	5,411.3	Lower Marker	43.00
4	38	8,412.99	334,510.3		43.46
5	45	11,742.88	398,665.0		44.42
6	55	14,269.53	396,376.3		45.71
7	60	15,125.60	381,565.1		46.31
8	68	12,330.77	274,246.3		47.18
9	82	7,796.96	144,589.3		48.65
10	104	8,111.12	117,689.7		51.04
11	144	6,944.65	73,197.8		54.67
12	157	12,319.48	119,074.5		55.86
13	163	7,067.34	65,576.2		56.46
14	170	8,865.16	79,092.8		57.06
15	179	7,131.36	60,236.8		57.93
16	194	11,694.24	91,118.6		59.31
17	207	6,897.89	50,450.4		60.45
18	215	10,041.62	70,815.0		61.14
19	237	12,795.32	81,838.6		63.12
20	259	5,026.98	29,415.5		65.09
21	269	5,557.83	31,283.2		66.01
22	277	5,411.12	29,611.6		66.70
23	286	4,590.06	24,352.1		67.48
24	293	5,120.15	26,452.2		68.17
25	307	7,012.43	34,567.2		69.36

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad


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Electropherogram Summary Continued ...

... Peak table for sample 6 : sample 6

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	327	6,054.68	28,020.7		70.97
27	363	18,513.79	77,184.2		73.87
28	408	4,994.67	18,538.5		77.27
29	435	5,168.45	17,997.3		78.78
30	456	3,568.06	11,868.5		79.93
31	478	2,592.51	8,212.0		81.22
32	492	3,373.29	10,384.3		82.00
33	509	2,818.15	8,384.1		82.87
34	523	4,320.56	12,514.5		83.51
35	555	1,968.27	5,376.7		84.98
36	579	2,761.60	7,222.8		86.13
37	604	3,463.63	8,683.8		87.23
38	650	1,357.56	3,162.7		88.70
39	666	1,798.13	4,089.7		89.21
40	726	3,317.29	6,922.1		90.63
41	869	913.57	1,592.5		92.51
42	904	888.09	1,488.3		92.97
43	974	18.42	28.7		93.89
44	1,160	18.61	24.3		95.27
45	1,266	22.03	26.4		95.96
46	1,543	71.85	70.6		97.75
47	1,869	11.88	9.6		99.86
48	2,546	10.55	6.3		102.53
49	3,038	16.78	8.4		104.09
50	4,522	26.72	9.0		106.06
51	5,730	16.37	4.3		107.67
52	6,420	18.98	4.5		108.59
53	9,953	11.50	1.8		112.54
54	10,380	75.00	10.9	Upper Marker	113.00

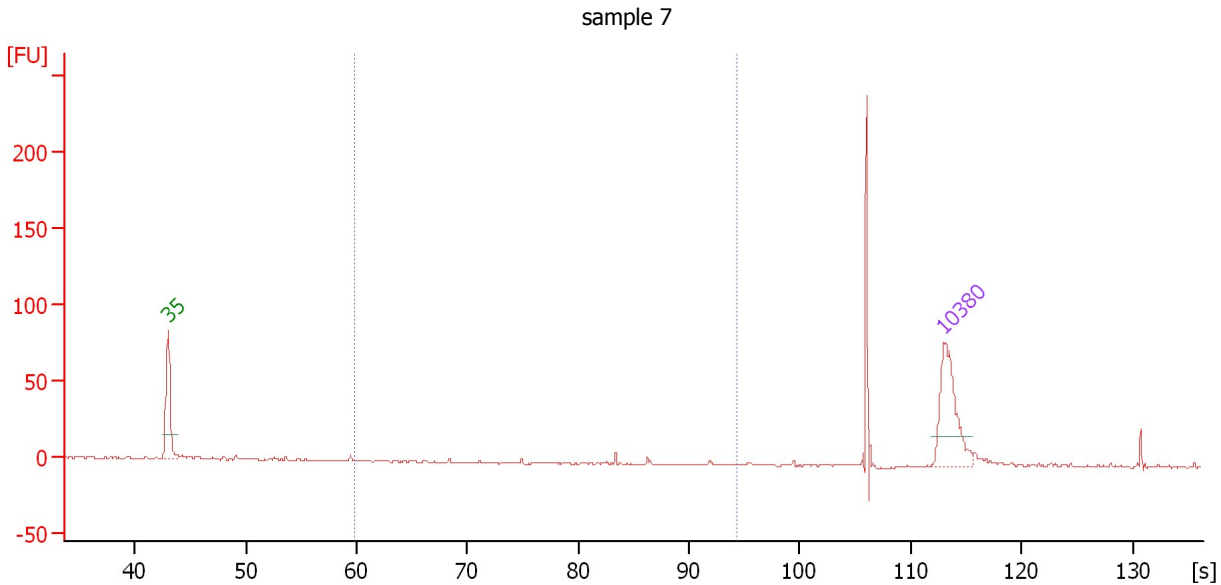
Region table for sample 6 : sample 6

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	381	1,000	5,933.5	57	36.6	84,684.02	398,452.0	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 7 : sample 7

Number of peaks found: 0 Corr. Area 1: 2.2
 Noise: 0.7

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	10,380	75.00	10.9	Upper Marker	113.00

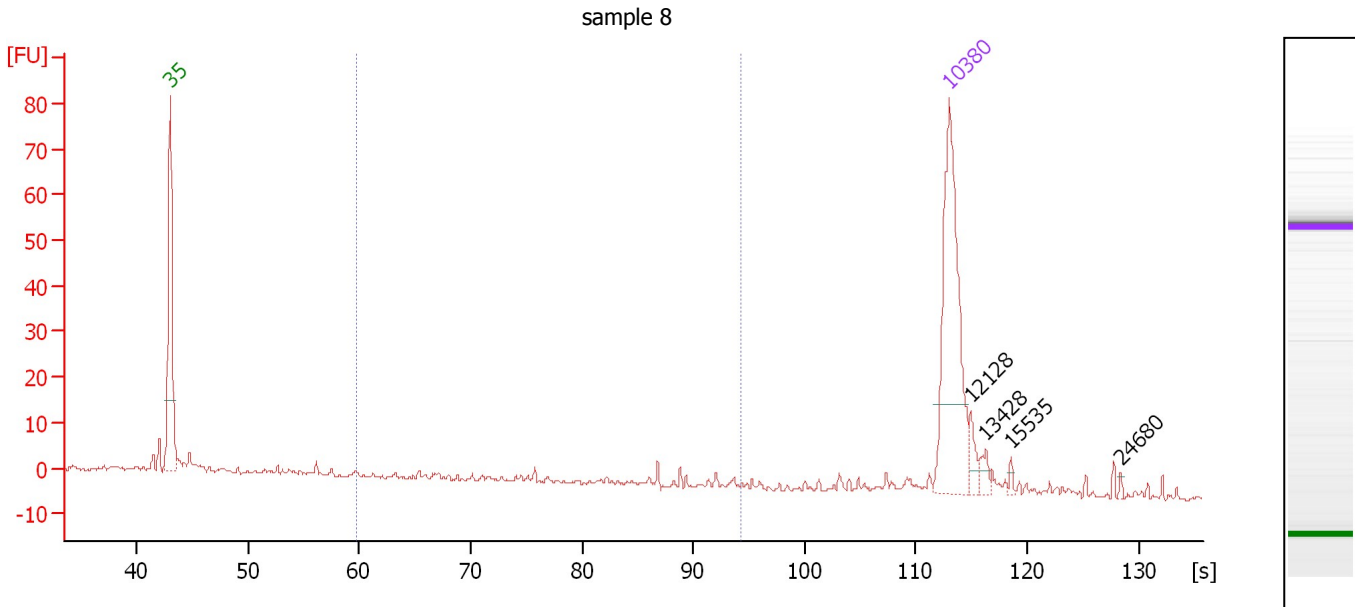
Region table for sample 7 : sample 7

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	517	1,000	2.2	3	21.0	1.86	5.8	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 8 : sample 8

Number of peaks found: 4 Corr. Area 1: 10.4
 Noise: 0.4

Peak table for sample 8 : sample 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	10,380	75.00	10.9	Upper Marker	113.00
3	12,128	0.00	0.0		114.88
4	13,428	0.00	0.0		116.28
5	15,535	0.00	0.0		118.55
6	24,680	0.00	0.0		128.39

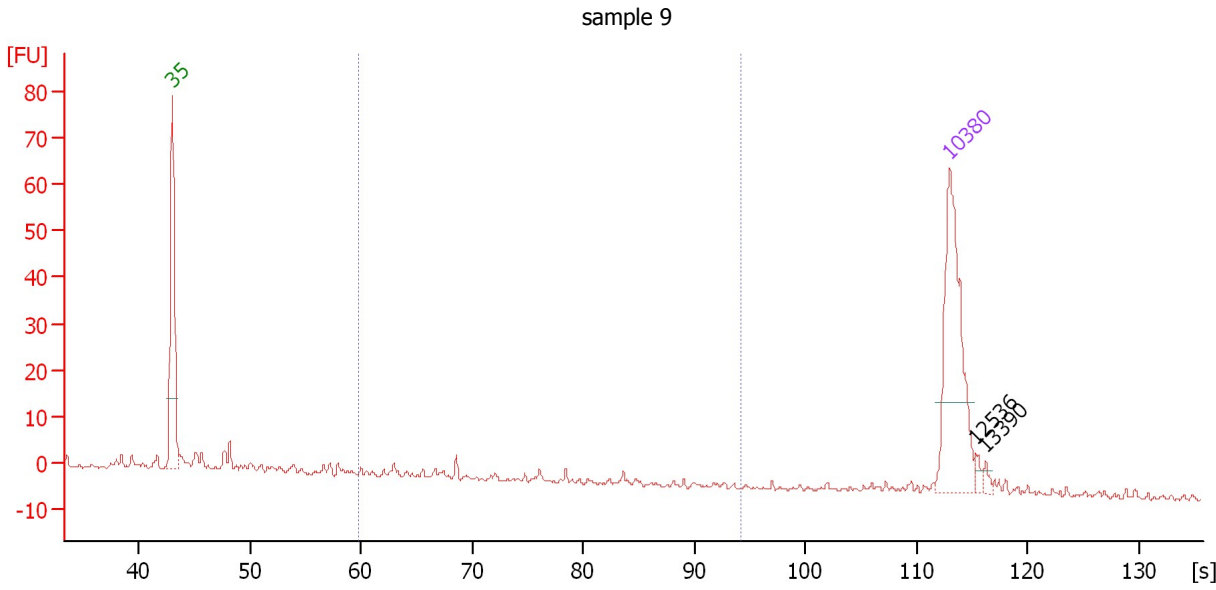
Region table for sample 8 : sample 8

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	531	1,000	10.4	13	40.4	9.04	33.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 9 : sample 9

Number of peaks found: 2 Corr. Area 1: 14.2
 Noise: 0.6

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	12,536	0.00	0.0		115.32
4	13,390	0.00	0.0		116.24

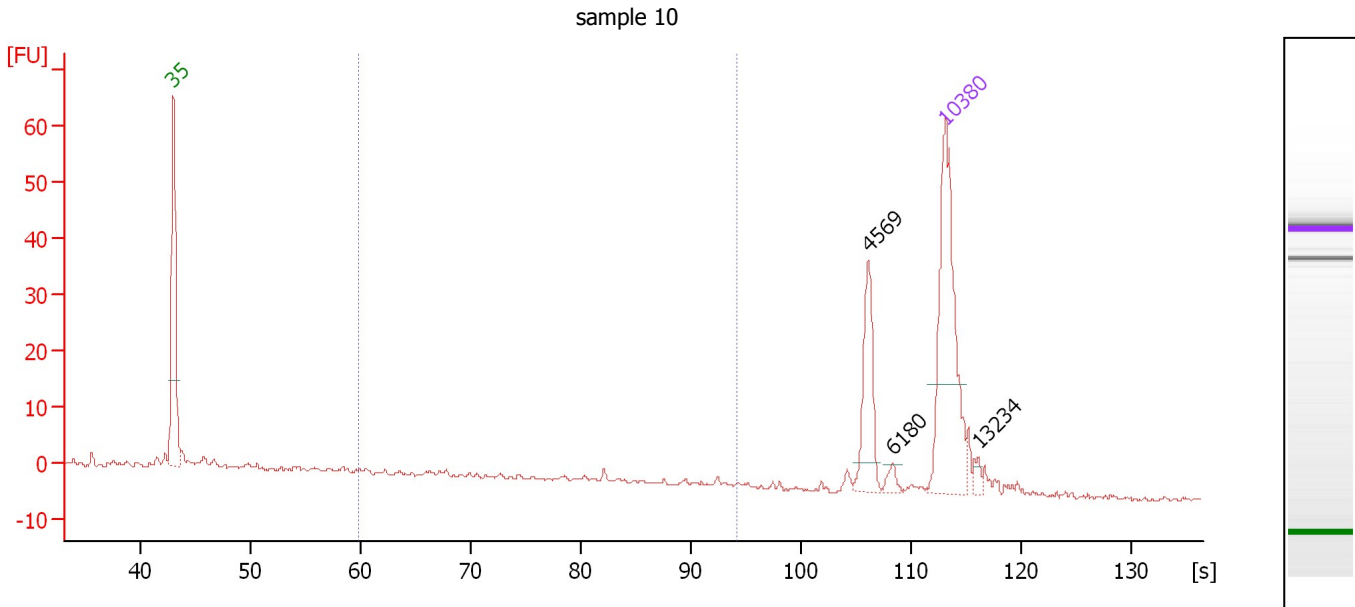
Region table for sample 9 : sample 9

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	342	1,000	14.2	17	38.9	15.62	80.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

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 Modified: 1/7/2015 12:30:19 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : sample 10

Number of peaks found: 3 Corr. Area 1: 2.6
 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	35	125.00	5,411.3	Lower Marker	43.00
2	4,569	32.38	10.7		106.13
3	6,180	3.68	0.9		108.27
4	10,380	75.00	10.9	Upper Marker	113.00
5	13,234	0.00	0.0		116.07

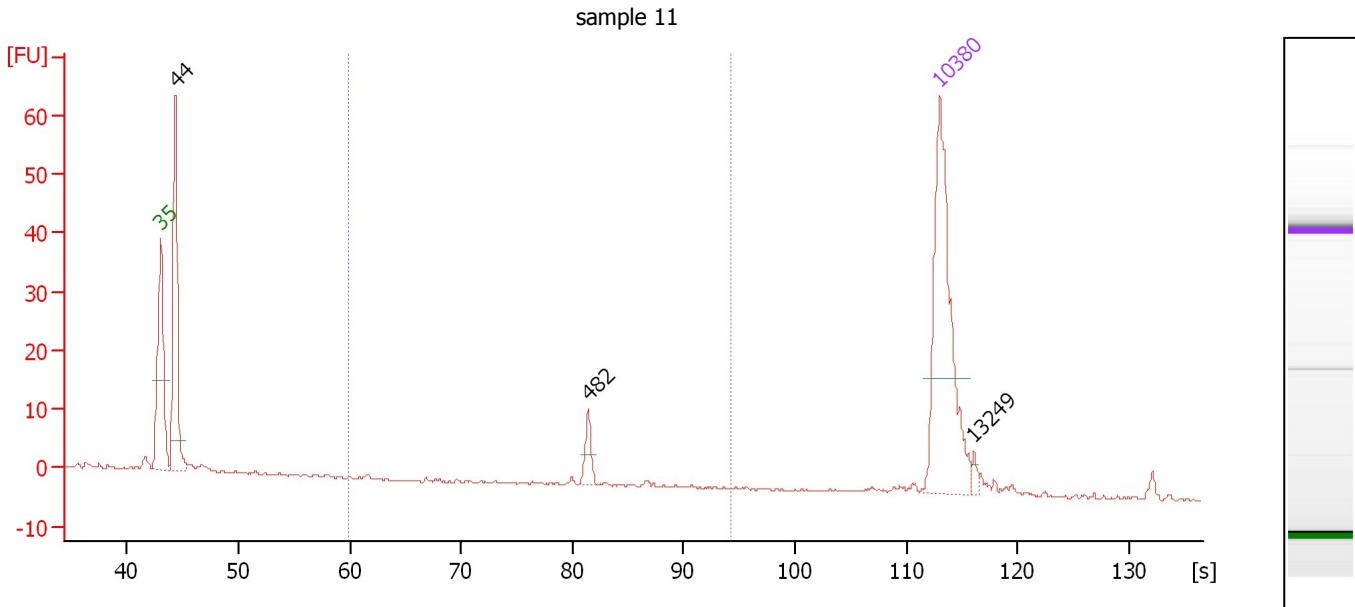
Region table for sample 10 : sample 10

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	480	1,000	2.6	4	43.2	2.99	12.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 11 : sample 11

Number of peaks found: 3 Corr. Area 1: 9.9
 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	44	108.30	3,718.0		44.35
3	482	9.82	30.8		81.45
4	10,380	75.00	10.9	Upper Marker	113.00
5	13,249	0.00	0.0		116.09

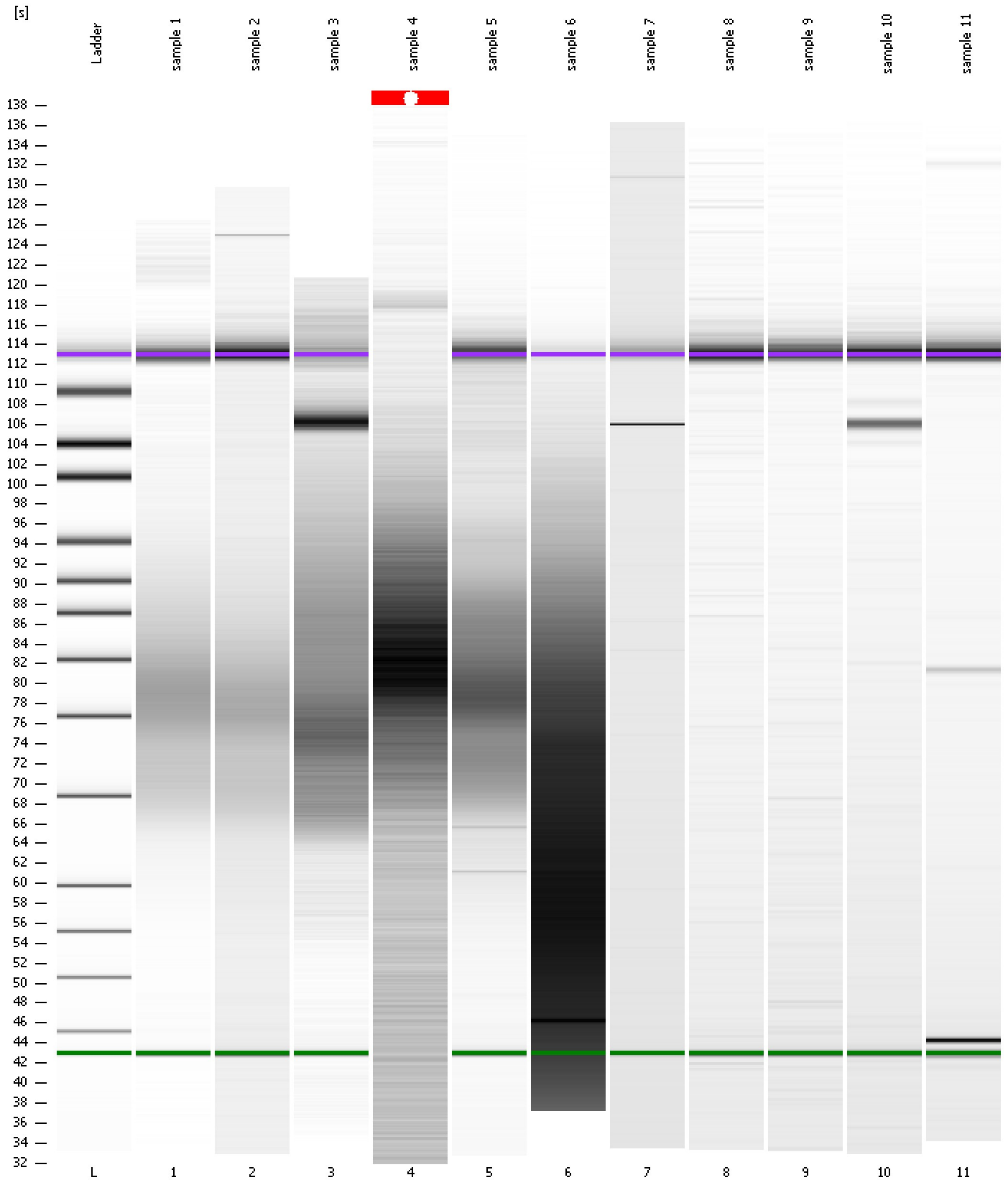
Region table for sample 11 : sample 11

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	483	1,000	9.9	9	6.4	9.83	31.2	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

Created: 1/7/2015 11:48:58 AM
Modified: 1/7/2015 12:30:19 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad

Created: 1/7/2015 11:48:58 AM
 Modified: 1/7/2015 12:30:19 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		1/7/2015 12:30:15 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-01-07\2015-01-07_001.xad)		Instrument	Run		1/7/2015 11:49:03 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		1/7/2015 11:49:03 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		1/7/2015 11:49:03 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		1/7/2015 11:49:03 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		1/7/2015 11:49:03 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		1/7/2015 11:49:03 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		1/7/2015 11:49:03 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1