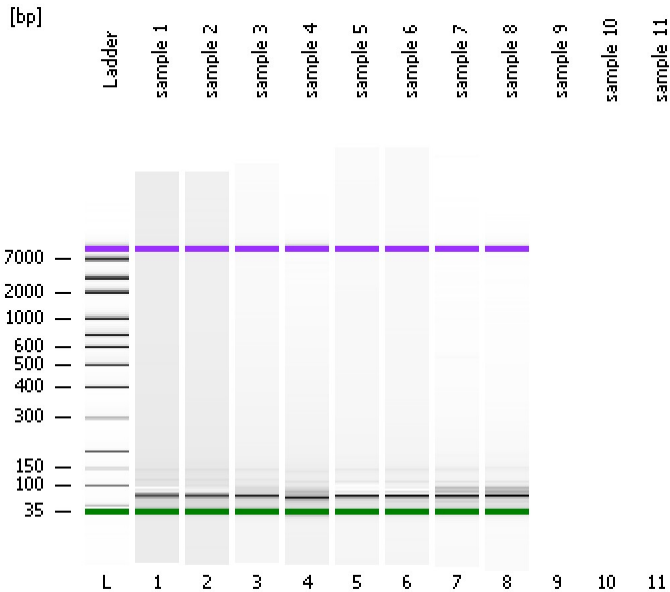


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
Modified: 1/7/2014 12:34:35 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

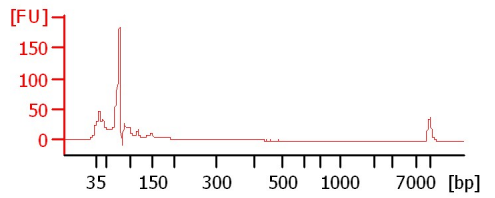
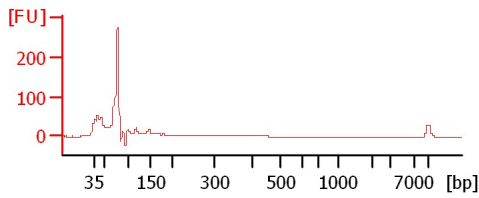
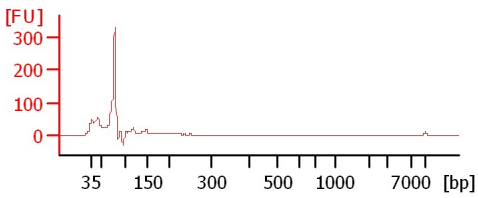
Chip Information:

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

sample 1

sample 2

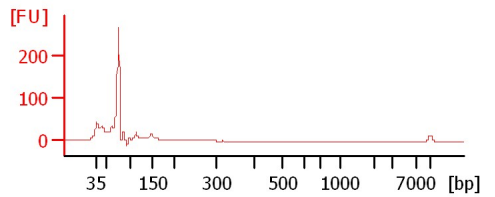
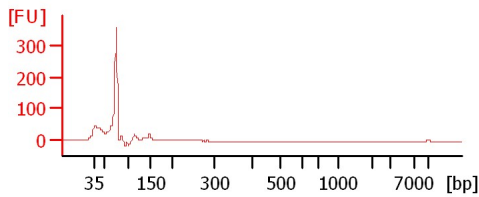
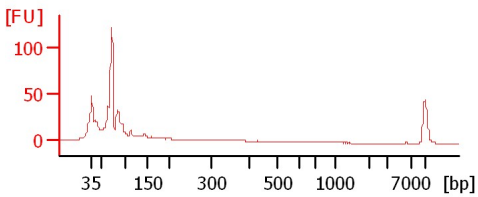
sample 3



sample 4

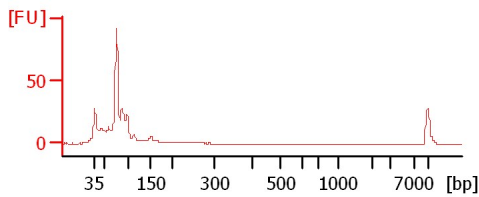
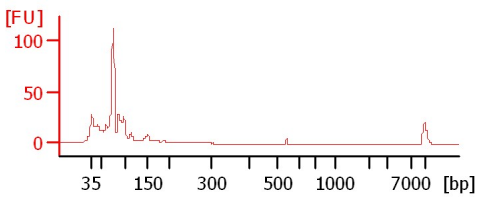
sample 5

sample 6



sample 7

sample 8



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
Modified: 1/7/2014 12:34:35 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:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
Modified: 1/7/2014 12:34:35 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

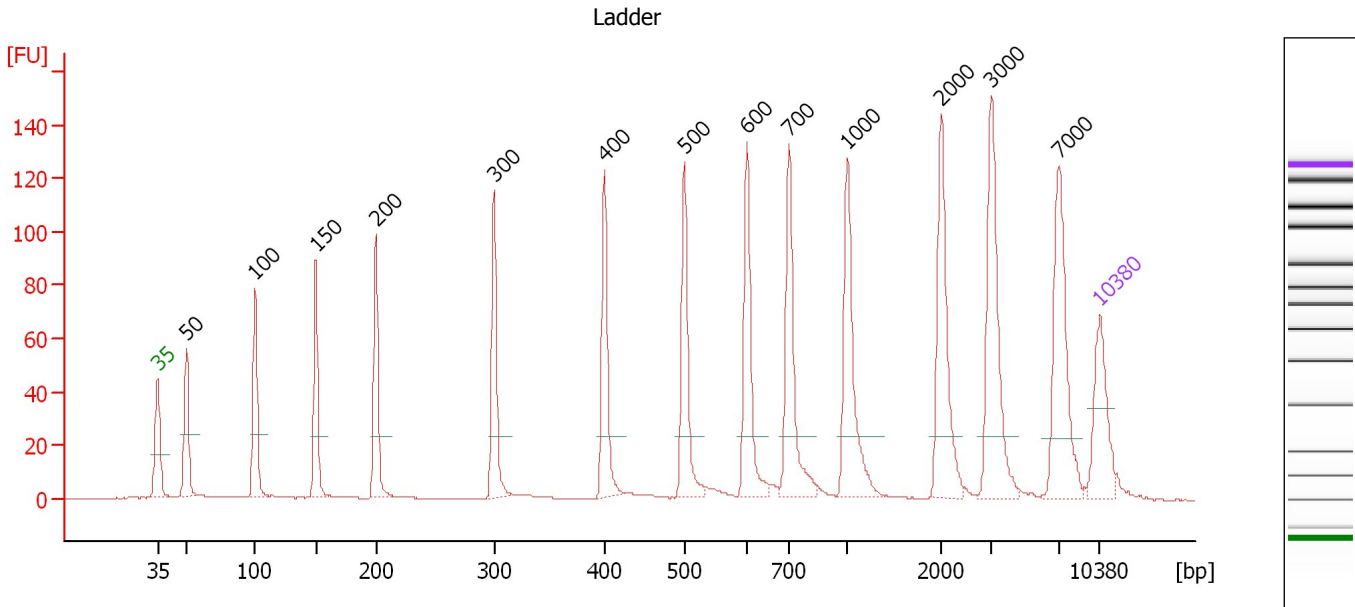
Ladder

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:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
 Modified: 1/7/2014 12:34:35 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

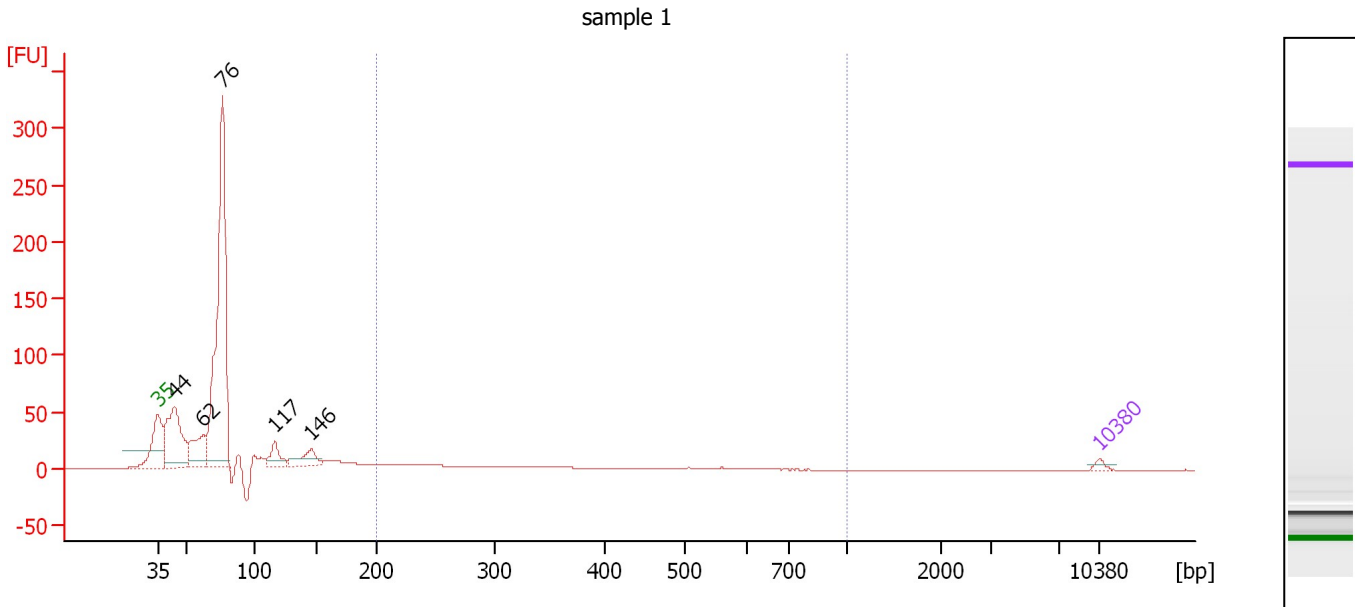
Peak table for Ladder

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	45.12	50	150.00	4,545.5	Ladder Peak
3	50.25	100	150.00	2,272.7	Ladder Peak
4	54.78	150	150.00	1,515.2	Ladder Peak
5	59.22	200	150.00	1,136.4	Ladder Peak
6	67.99	300	150.00	757.6	Ladder Peak
7	76.23	400	150.00	568.2	Ladder Peak
8	82.09	500	150.00	454.5	Ladder Peak
9	86.77	600	150.00	378.8	Ladder Peak
10	89.88	700	150.00	324.7	Ladder Peak
11	94.22	1,000	150.00	227.3	Ladder Peak
12	101.17	2,000	150.00	113.6	Ladder Peak
13	104.92	3,000	150.00	75.8	Ladder Peak
14	109.94	7,000	150.00	32.5	Ladder Peak
15	113.00	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
 Modified: 1/7/2014 12:34:35 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : sample 1

Number of peaks found: 5 Corr. Area 1: 91.2
 Noise: 0.1

Peak table for sample 1 : sample 1

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	44.27	44	2,737.20	94,282.8	
3	46.37	62	1,134.61	27,638.7	
4	47.84	76	8,310.61	164,617.8	
5	51.74	117	505.51	6,574.0	
6	54.43	146	563.26	5,840.2	
7	113.00	10,380	75.00	10.9	Upper Marker

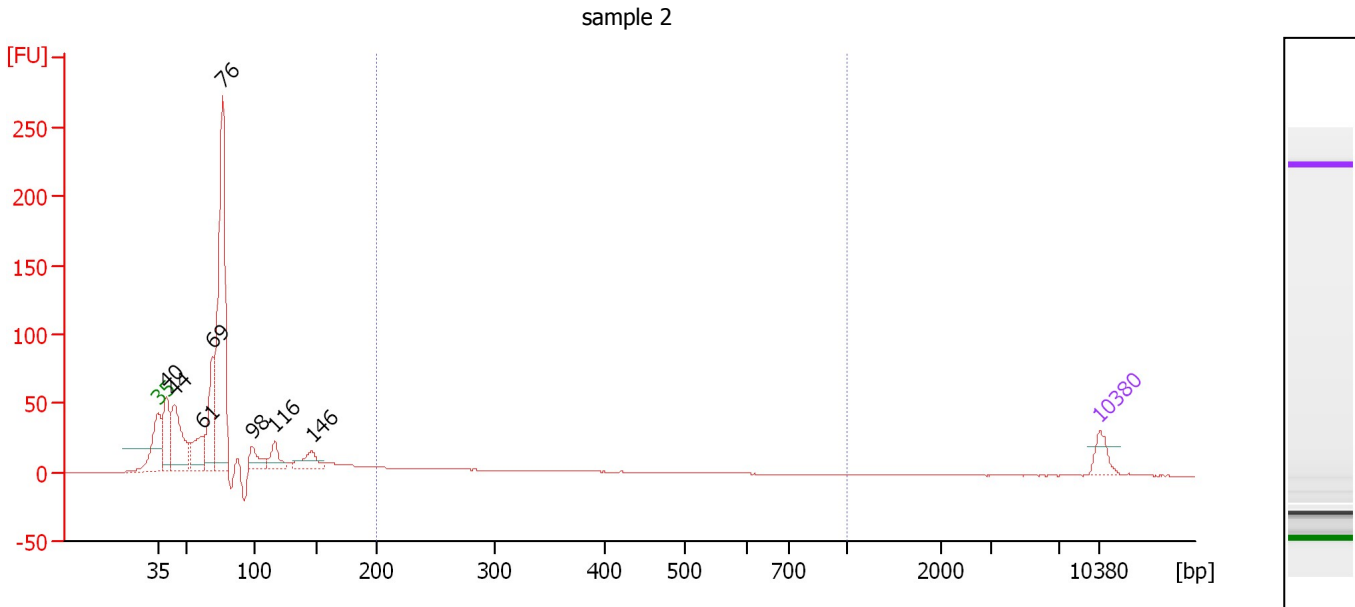
Region table for sample 1 : sample 1

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	340	6,205.4	1,199.08	91.2	9	38.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
 Modified: 1/7/2014 12:34:35 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : sample 2

Number of peaks found: 8 Corr. Area 1: 90.1
 Noise: 0.1

Peak table for sample 2 : sample 2

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	43.64	40	385.45	14,784.0	
3	44.27	44	539.15	18,562.9	
4	46.28	61	242.55	5,993.8	
5	47.06	69	393.34	8,643.2	
6	47.80	76	1,792.17	35,676.1	
7	50.05	98	137.65	2,126.6	
8	51.66	116	143.73	1,883.3	
9	54.45	146	145.98	1,510.9	
10	113.00	10,380	75.00	10.9	Upper Marker

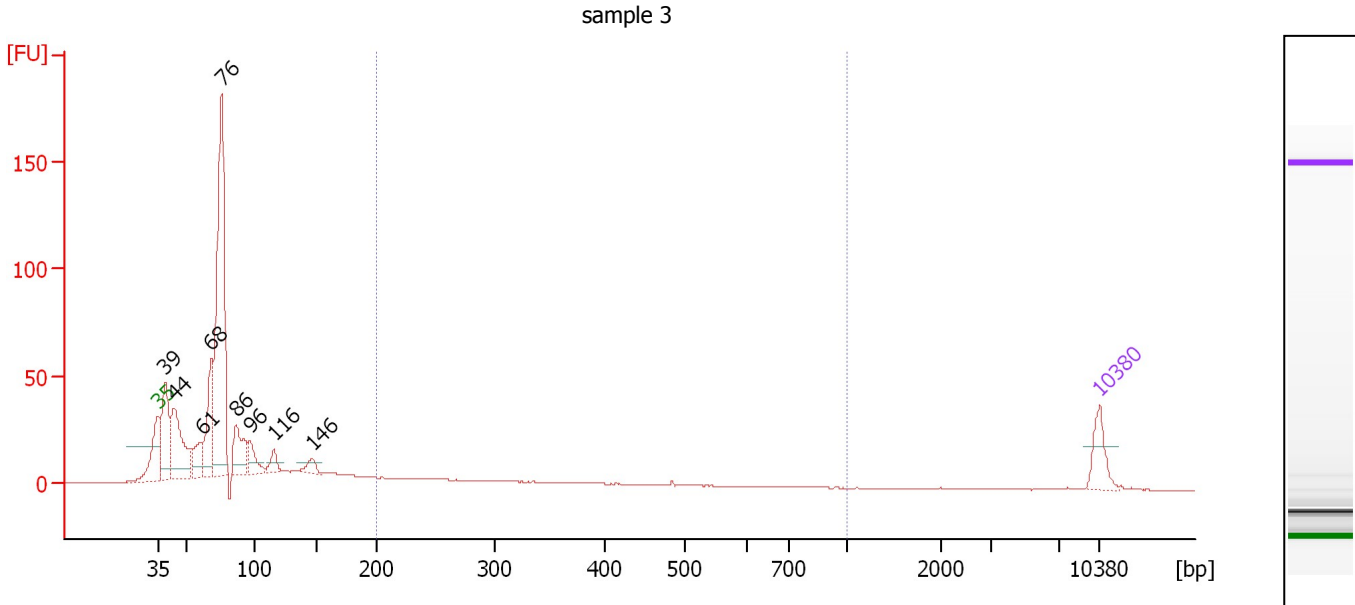
Region table for sample 2 : sample 2

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	337	1,848.1	358.15	90.1	10	36.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
 Modified: 1/7/2014 12:34:35 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : sample 3

Number of peaks found: 9 Corr. Area 1: 74.8
 Noise: 0.1

Peak table for sample 3 : sample 3

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	43.55	39	286.75	11,176.8	
3	44.24	44	315.94	10,928.7	
4	46.28	61	120.13	2,966.9	
5	46.98	68	226.13	5,028.0	
6	47.78	76	1,043.49	20,828.7	
7	48.82	86	175.83	3,094.4	
8	49.87	96	89.07	1,401.6	
9	51.66	116	49.79	652.9	
10	54.44	146	42.77	443.0	
11	113.00	10,380	75.00	10.9	Upper Marker

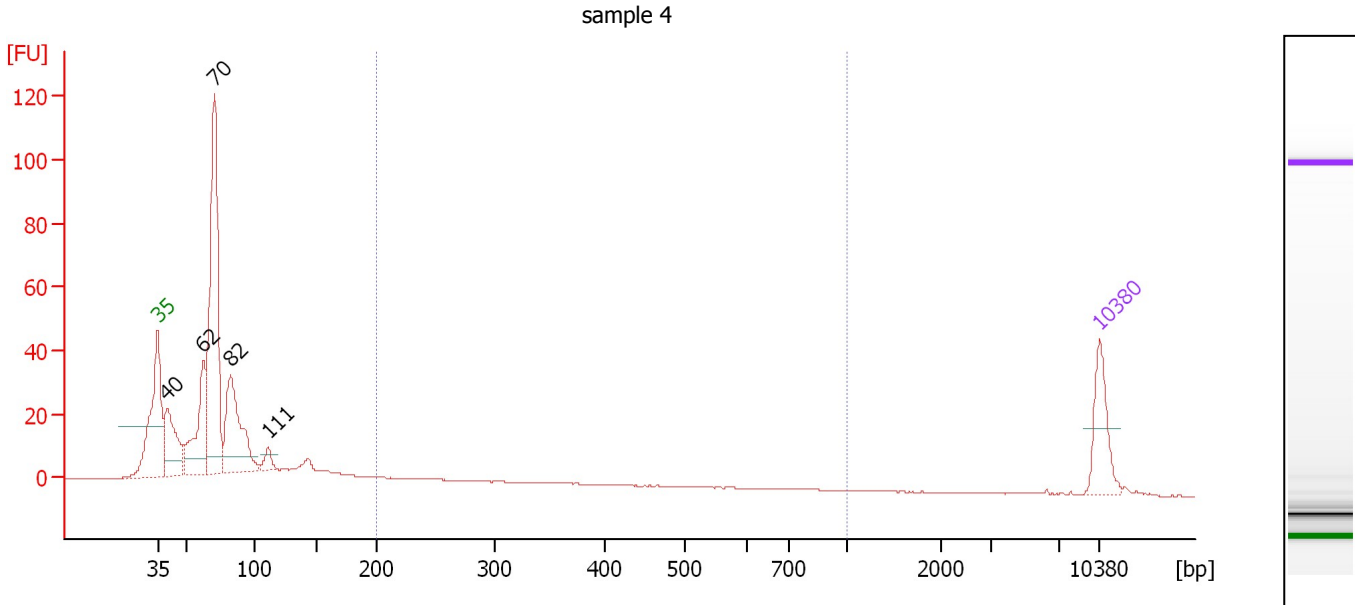
Region table for sample 3 : sample 3

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	346	1,311.2	257.10	74.8	10	38.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
 Modified: 1/7/2014 12:34:35 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : sample 4

Number of peaks found: 5 Corr. Area 1: 35.1
 Noise: 0.1

Peak table for sample 4 : sample 4

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	43.72	40	165.06	6,237.8	
3	46.39	62	190.31	4,620.4	
4	47.21	70	550.50	11,843.1	
5	48.40	82	248.68	4,597.3	
6	51.22	111	26.29	359.5	
7	113.00	10,380	75.00	10.9	Upper Marker

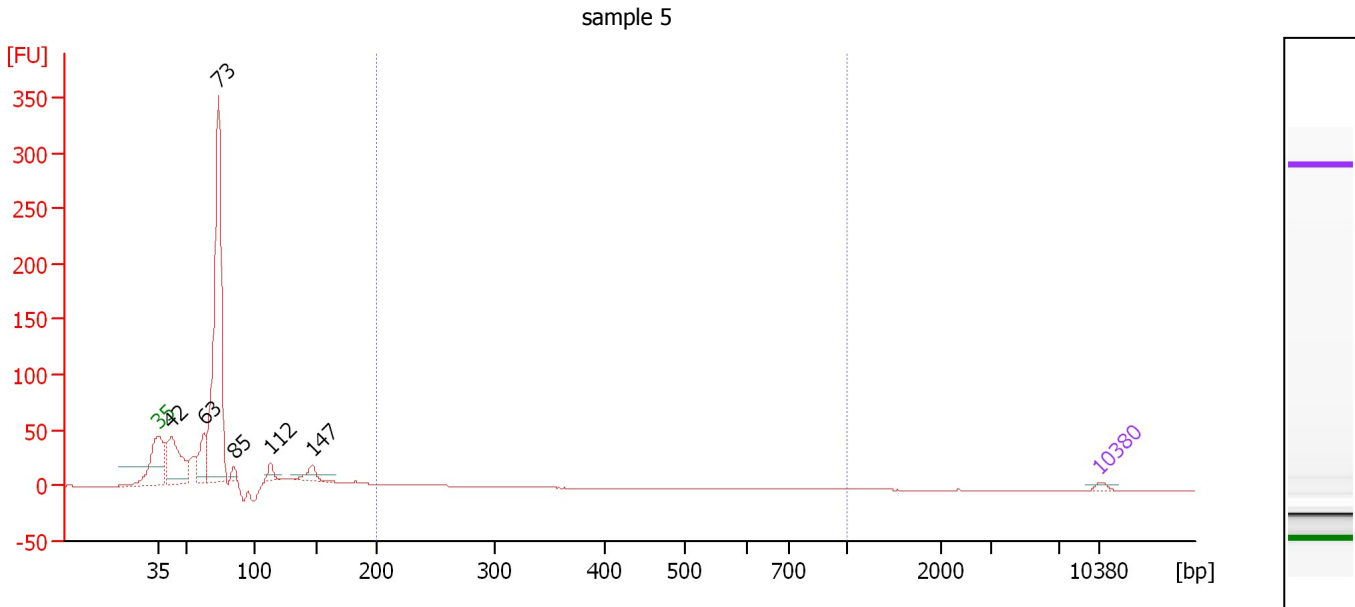
Region table for sample 4 : sample 4

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	323	479.2	90.79	35.1	7	32.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
 Modified: 1/7/2014 12:34:35 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : sample 5

Number of peaks found: 6 Corr. Area 1: 28.0
 Noise: 0.1

Peak table for sample 5 : sample 5

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	43.99	42	2,534.30	91,441.9	
3	46.43	63	1,202.99	29,023.0	
4	47.52	73	9,991.13	206,090.7	
5	48.67	85	239.36	4,286.3	
6	51.37	112	297.30	4,007.0	
7	54.49	147	451.01	4,654.3	
8	113.00	10,380	75.00	10.9	Upper Marker

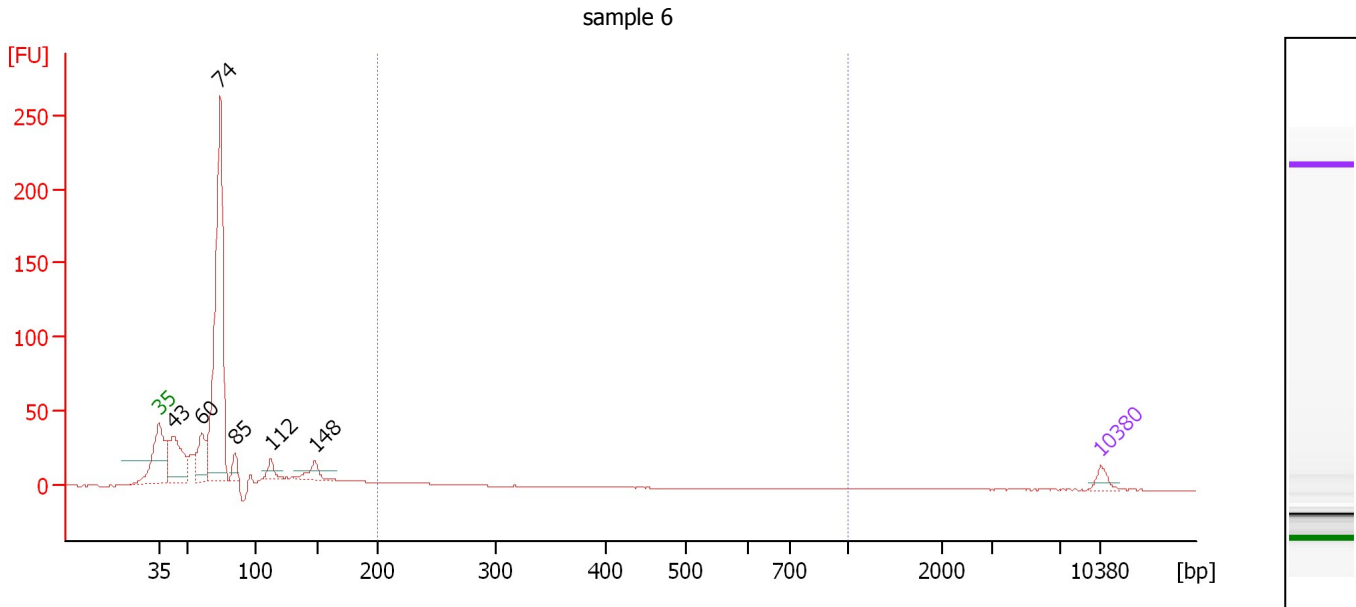
Region table for sample 5 : sample 5

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	251	3,229.7	525.26	28.0	3	16.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
 Modified: 1/7/2014 12:34:35 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : sample 6

Number of peaks found: 6 Corr. Area 1: 29.4
 Noise: 0.1

Peak table for sample 6 : sample 6

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	44.10	43	950.18	33,652.5	
3	46.19	60	535.88	13,426.2	
4	47.55	74	3,963.83	81,434.1	
5	48.71	85	178.57	3,183.6	
6	51.32	112	147.29	1,994.6	
7	54.62	148	197.07	2,014.1	
8	113.00	10,380	75.00	10.9	Upper Marker

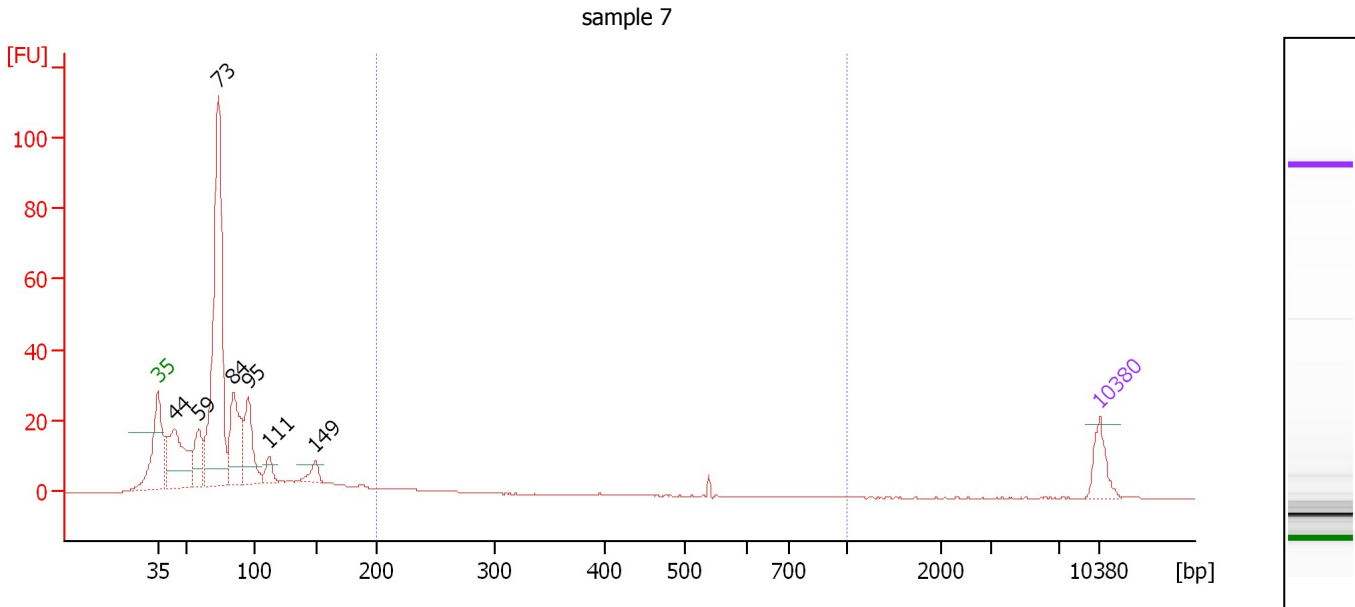
Region table for sample 6 : sample 6

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	260	1,702.5	282.63	29.4	4	21.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
 Modified: 1/7/2014 12:34:35 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : sample 7

Number of peaks found: 7 Corr. Area 1: 15.6
 Noise: 0.1

Peak table for sample 7 : sample 7

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	44.25	44	393.73	13,612.2	
3	46.01	59	173.15	4,467.6	
4	47.52	73	1,276.50	26,342.6	
5	48.61	84	336.68	6,068.2	
6	49.76	95	259.21	4,124.9	
7	51.26	111	57.83	788.0	
8	54.69	149	56.65	576.0	
9	113.00	10,380	75.00	10.9	Upper Marker

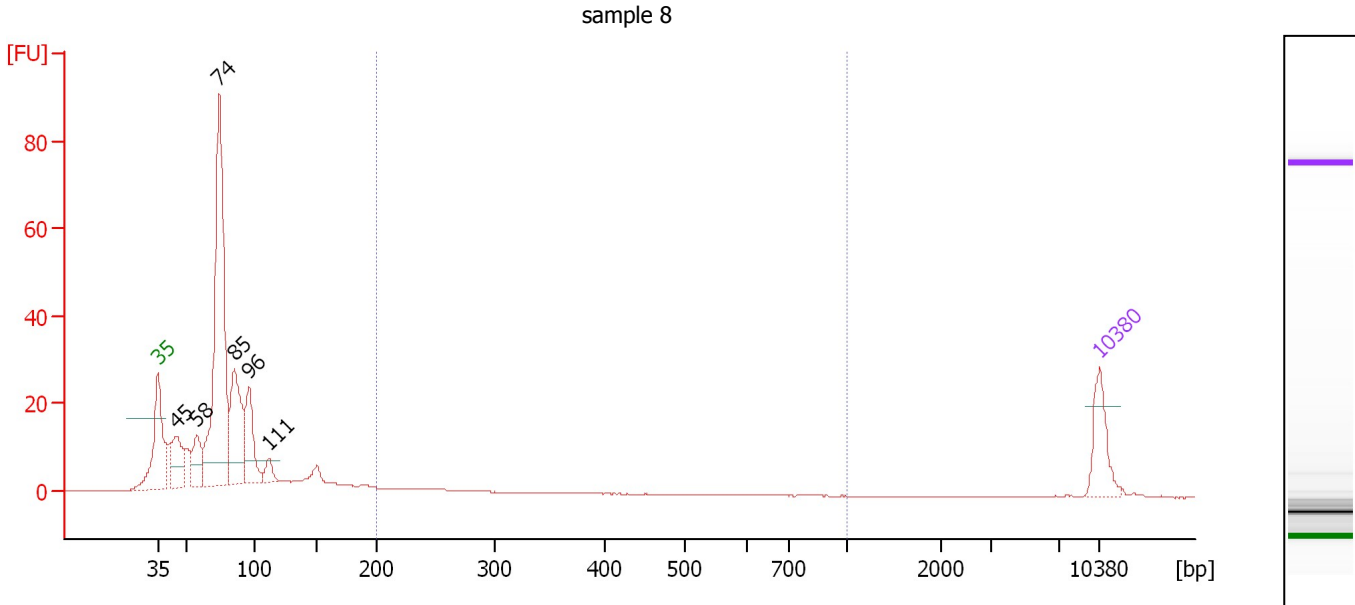
Region table for sample 7 : sample 7

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	290	563.9	96.34	15.6	4	37.7	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
 Modified: 1/7/2014 12:34:35 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : sample 8

Number of peaks found: 6 Corr. Area 1: 12.5
 Noise: 0.1

Peak table for sample 8 : sample 8

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	44.38	45	125.89	4,262.5	
3	45.91	58	88.92	2,335.5	
4	47.59	74	774.31	15,830.5	
5	48.71	85	253.50	4,515.6	
6	49.79	96	159.26	2,526.5	
7	51.27	111	30.46	414.9	
8	113.00	10,380	75.00	10.9	Upper Marker

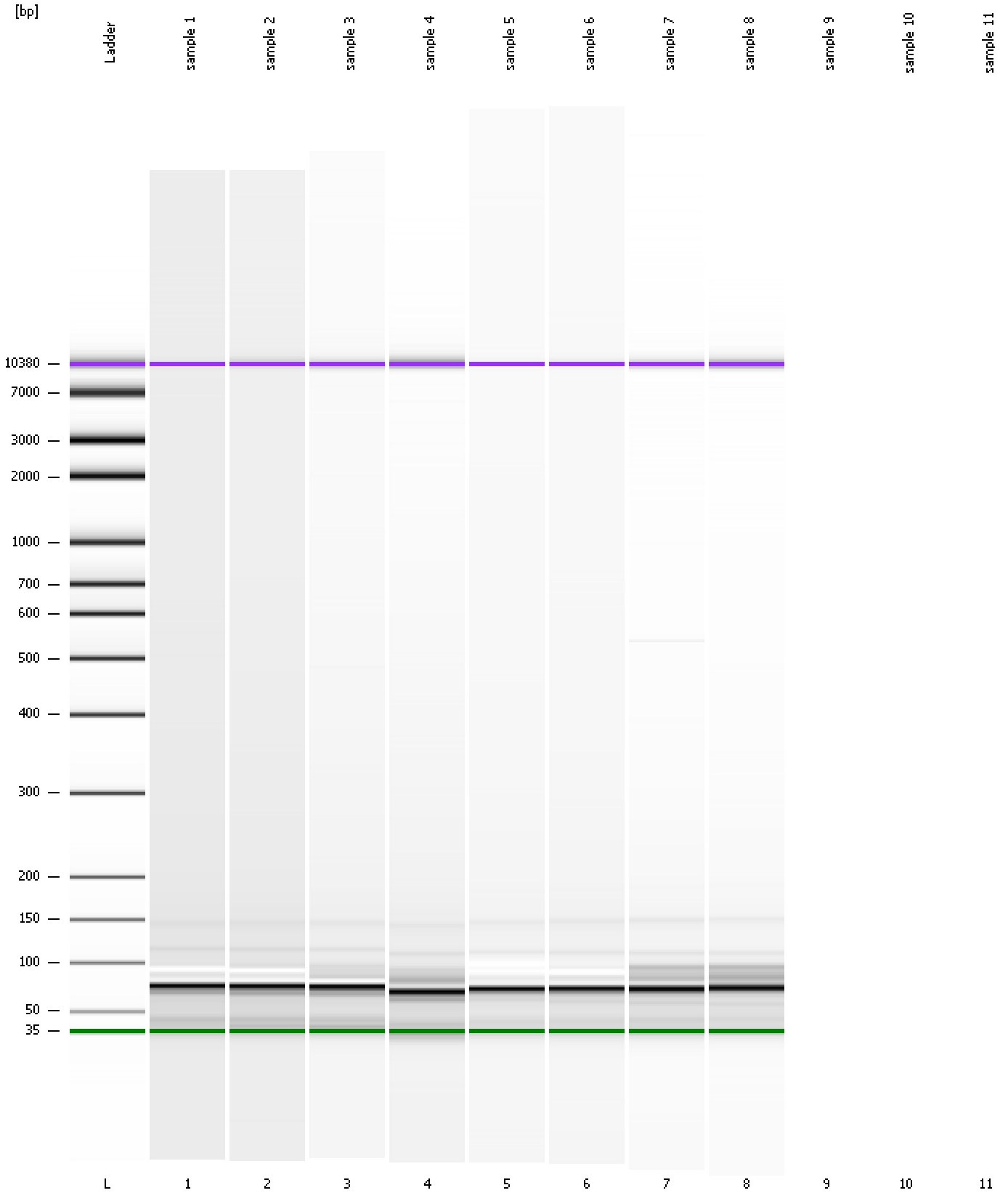
Region table for sample 8 : sample 8

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	268	332.8	56.11	12.5	4	23.6	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
Modified: 1/7/2014 12:34:35 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ttings\Bioanalyzer\2014-01-07\2014-01-07\2014-01-07_001.xad

Created: 1/7/2014 12:01:49 PM
Modified: 1/7/2014 12:34:35 PM

Invalid Samples

Sample 9 has not been run, no results available.

Sample 10 has not been run, no results available.

Sample 11 has not been run, no results available.