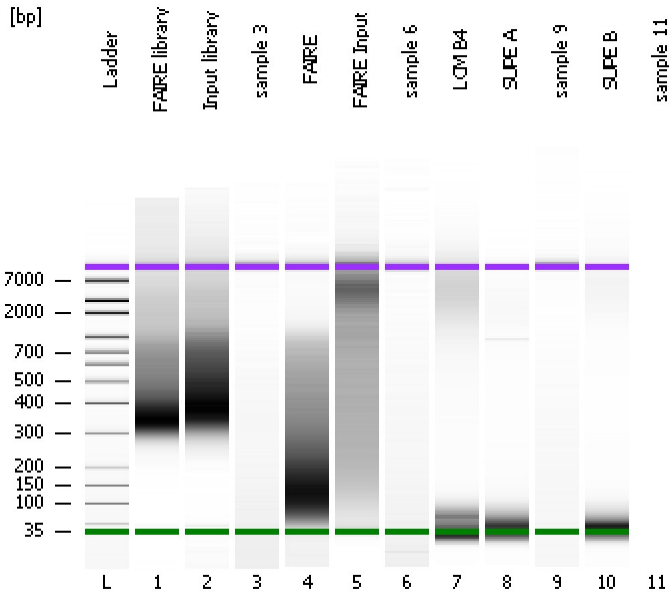


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
Modified: 4/17/2015 11:04:33 AM

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

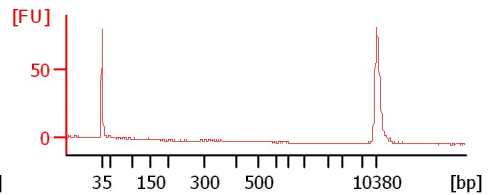
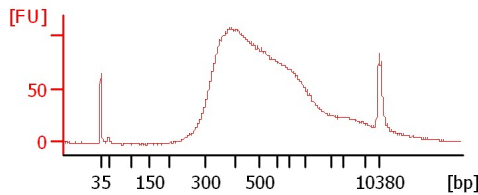
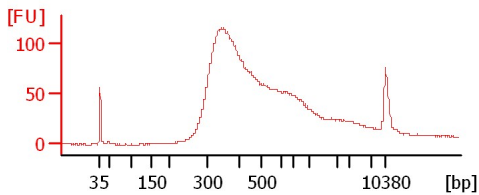
Chip Information:

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

FAIRE library

Input library

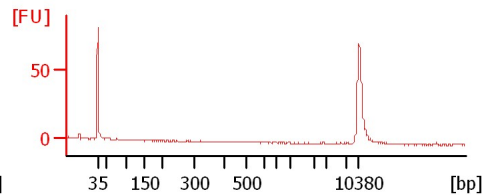
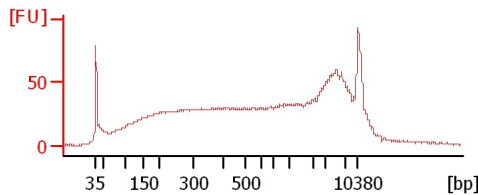
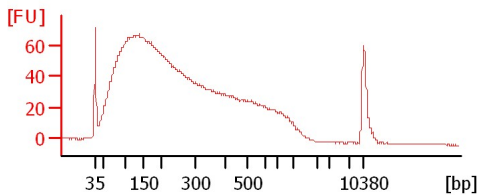
sample 3



FAIRE

FAIRE Input

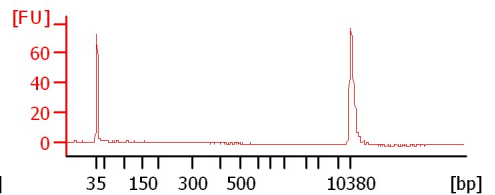
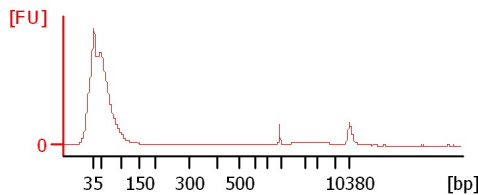
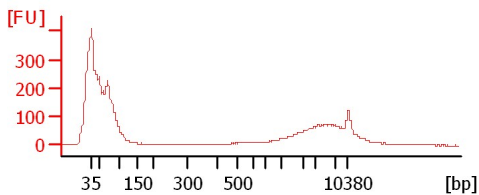
sample 6



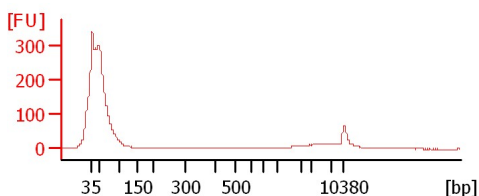
LCM B4

SUPE A

sample 9



SUPE B



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
FAIRE library		<input type="checkbox"/>	✓			
Input library		<input type="checkbox"/>	✓			
sample 3		<input type="checkbox"/>	✓			
FAIRE		<input type="checkbox"/>	✓			
FAIRE Input		<input type="checkbox"/>	✓			
sample 6		<input type="checkbox"/>	✓			
LCM B4		<input type="checkbox"/>	✓			
SUPE A		<input type="checkbox"/>	✓			
sample 9		<input type="checkbox"/>	✓			
SUPE B		<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\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
Modified: 4/17/2015 11:04:33 AM

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

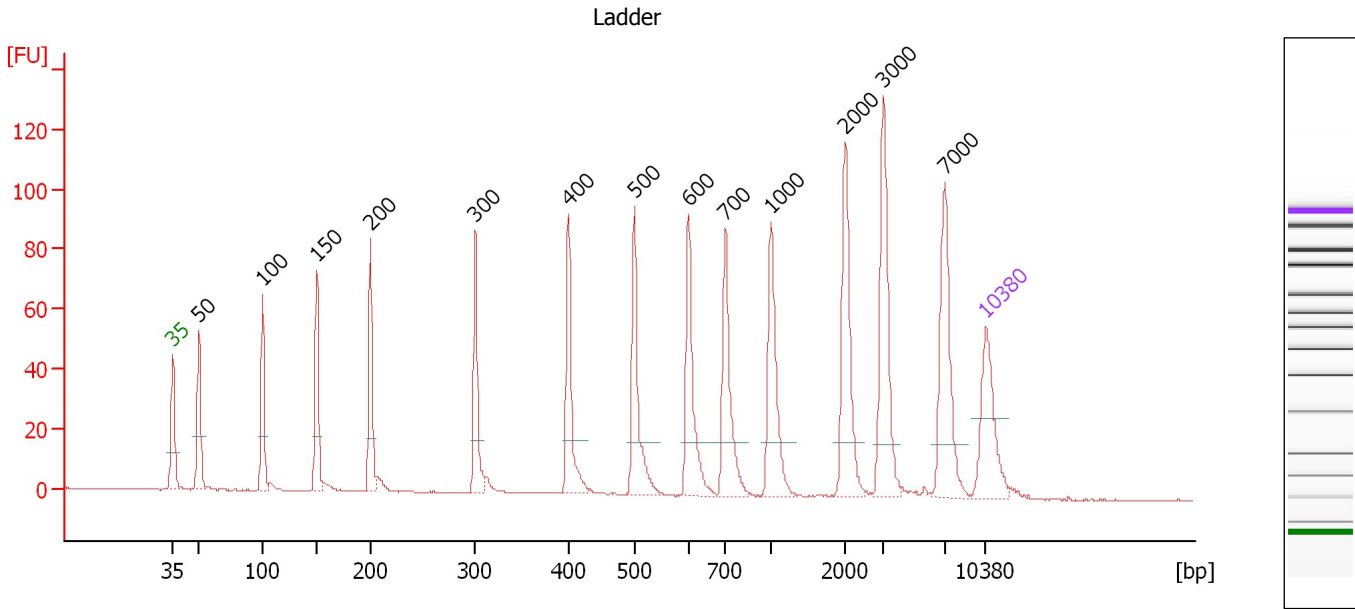
Ladder

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\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

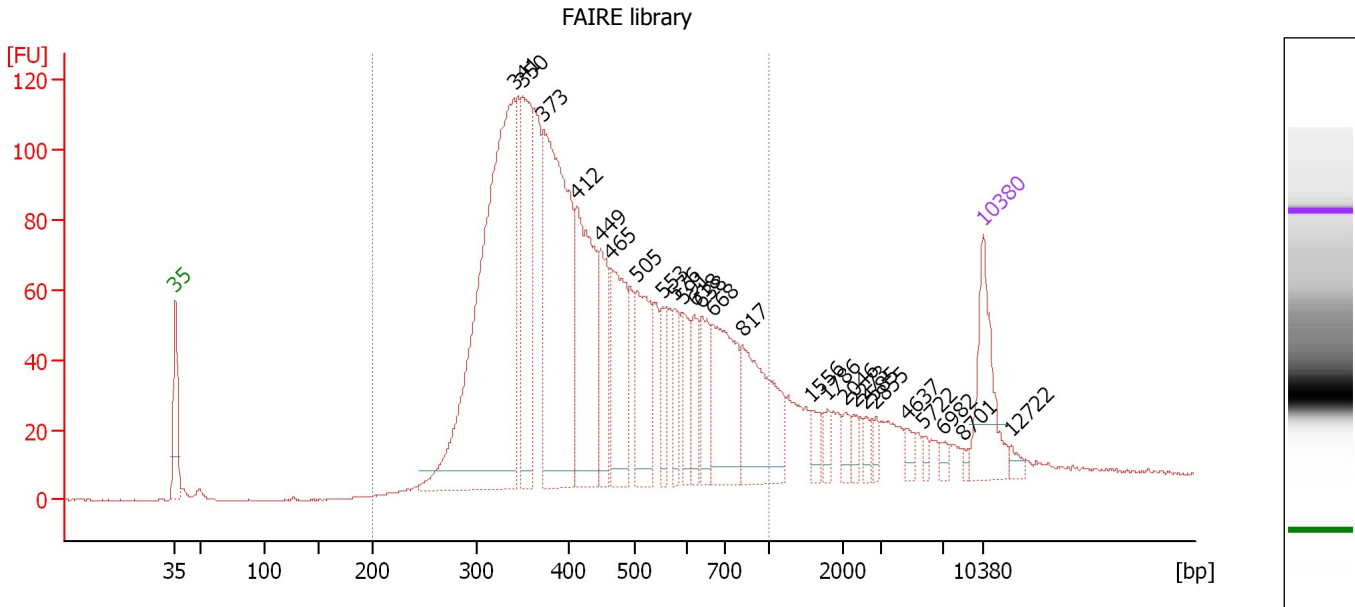
Peak table for Ladder

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	45.25
3	100	150.00	2,272.7	Ladder Peak	50.76
4	150	150.00	1,515.2	Ladder Peak	55.44
5	200	150.00	1,136.4	Ladder Peak	60.03
6	300	150.00	757.6	Ladder Peak	69.07
7	400	150.00	568.2	Ladder Peak	77.06
8	500	150.00	454.5	Ladder Peak	82.75
9	600	150.00	378.8	Ladder Peak	87.39
10	700	150.00	324.7	Ladder Peak	90.60
11	1,000	150.00	227.3	Ladder Peak	94.50
12	2,000	150.00	113.6	Ladder Peak	100.93
13	3,000	150.00	75.8	Ladder Peak	104.19
14	7,000	150.00	32.5	Ladder Peak	109.47
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : FAIRE library

Number of peaks found: 25 Corr. Area 1: 2,392.7
 Noise: 0.2

Peak table for sample 1 : FAIRE library

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	341	725.86	3,226.0		72.34
3	350	195.98	848.0		73.08
4	373	422.67	1,718.1		74.88
5	412	228.21	839.1		77.75
6	449	95.03	321.0		79.83
7	465	138.67	452.0		80.75
8	505	121.10	363.6		82.97
9	553	40.31	110.3		85.23
10	576	38.45	101.1		86.29
11	591	44.68	114.5		86.99
12	618	43.73	107.3		87.96
13	638	51.10	121.4		88.60
14	668	135.91	308.2		89.57
15	817	143.33	266.0		92.12
16	1,556	16.75	16.3		98.08
17	1,786	14.36	12.2		99.55
18	2,046	13.13	9.7		101.08
19	2,273	9.74	6.5		101.82
20	2,585	9.34	5.5		102.83
21	2,855	9.00	4.8		103.71
22	4,637	9.96	3.3		106.35
23	5,722	6.53	1.7		107.78
24	6,982	6.93	1.5		109.44
25	8,701	4.14	0.7		111.24


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...**... Peak table for sample 1 : FAIRE library**

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
27	12,722	0.00	0.0		115.45

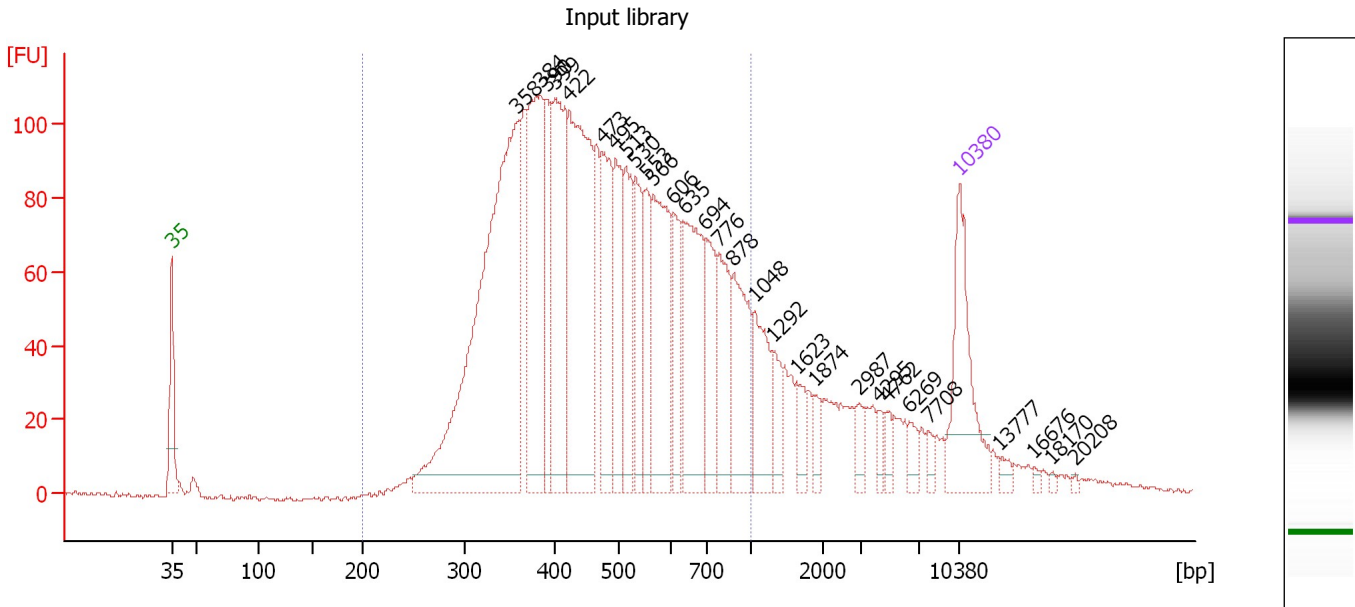
Region table for sample 1 : FAIRE library

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/ μ l]	Color
200	460	1,000	2,392.7	10,313.6	88	33.5	2,767.84	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 : Input library

Number of peaks found: 29 Corr. Area 1: 2,775.3
 Noise: 0.6

Peak table for sample 2 : Input library

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	358	494.80	2,091.7		73.74
3	384	185.47	732.0		75.77
4	390	71.95	279.6		76.25
5	399	166.21	630.6		77.01
6	422	240.83	864.0		78.33
7	473	104.77	335.5		81.22
8	495	79.50	243.5		82.45
9	513	67.81	200.3		83.35
10	530	63.84	182.4		84.16
11	553	49.49	135.6		85.20
12	566	123.24	329.8		85.81
13	606	42.45	106.2		87.57
14	635	113.10	269.8		88.51
15	694	65.80	143.6		90.41
16	776	65.45	127.7		91.59
17	878	78.02	134.6		92.92
18	1,048	53.78	77.7		94.81
19	1,292	23.73	27.8		96.38
20	1,623	16.70	15.6		98.51
21	1,874	11.36	9.2		100.12
22	2,987	11.72	5.9		104.14
23	4,295	8.48	3.0		105.90
24	4,762	9.39	3.0		106.51
25	6,269	9.89	2.4		108.50

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...


... Peak table for sample 2 :

Input library

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	7,708	6.39	1.3		110.21
27	10,380	75.00	10.9	Upper Marker	113.00
28	13,777	0.00	0.0		116.55
29	16,676	0.00	0.0		119.58
30	18,170	0.00	0.0		121.15
31	20,208	0.00	0.0		123.28

Region table for sample 2 :

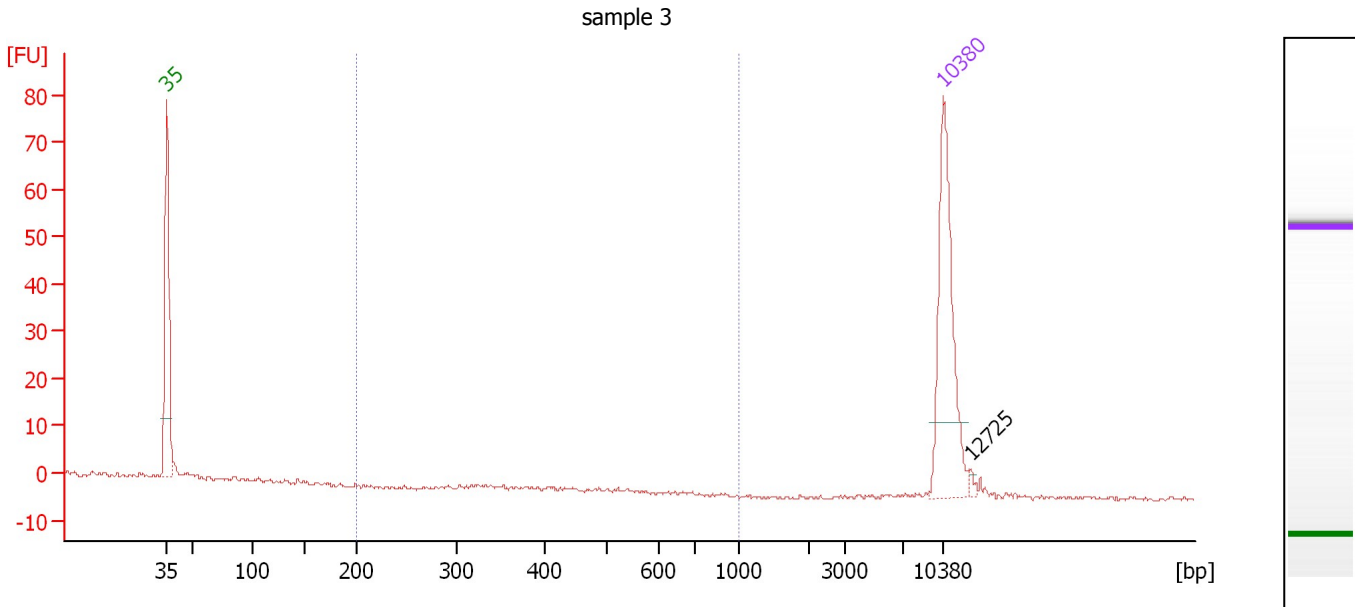
Input library

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/ μ l]	Color
200	496	1,000	2,775.3	7,627.4	84	32.2	2,202.39	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...



Overall Results for sample 3 : sample 3

Number of peaks found: 1 Corr. Area 1: 0.0
 Noise: 0.4

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	10,380	75.00	10.9	Upper Marker	113.00
3	12,725	0.00	0.0		115.45

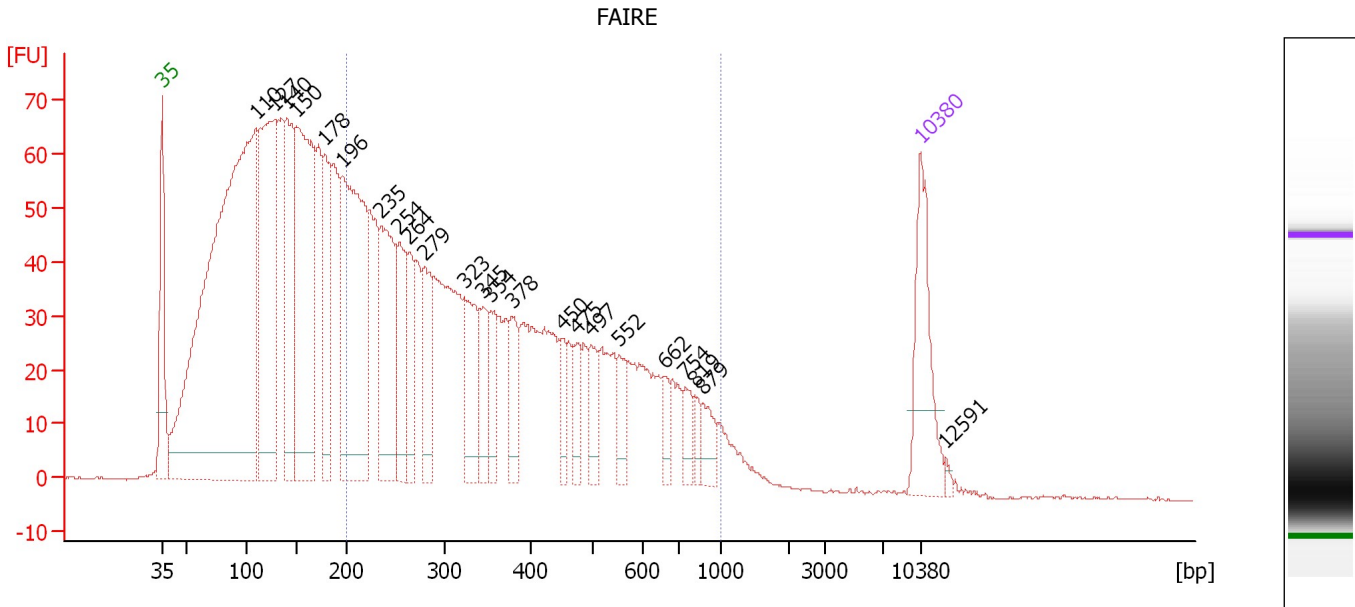
Region table for sample 3 : sample 3

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
200	0	1,000	0.0	0.0	0	0.0	0.00	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...



Overall Results for sample 4 : FAIRE

Number of peaks found: 23 Corr. Area 1: 1,535.5
 Noise: 0.2

Peak table for sample 4 : FAIRE

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	110	1,154.25	15,878.4		51.71
3	127	354.85	4,217.7		53.33
4	140	186.87	2,021.2		54.51
5	150	378.37	3,820.2		55.45
6	178	139.64	1,189.1		58.00
7	196	355.45	2,753.2		59.63
8	235	171.29	1,103.2		63.22
9	254	96.60	576.8		64.89
10	264	68.23	392.3		65.78
11	279	77.54	420.6		67.20
12	323	84.40	396.2		70.89
13	345	48.90	214.8		72.66
14	354	44.07	188.8		73.35
15	378	44.98	180.5		75.27
16	450	26.16	88.1		79.89
17	475	34.40	109.8		81.32
18	497	34.81	106.1		82.60
19	552	29.83	81.9		85.16
20	662	18.76	42.9		89.39
21	754	19.29	38.8		91.31
22	819	12.21	22.6		92.14
23	879	23.56	40.6		92.93
24	10,380	75.00	10.9	Upper Marker	113.00
25	12,591	0.00	0.0		115.31

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...

... Region table for sample 4 :

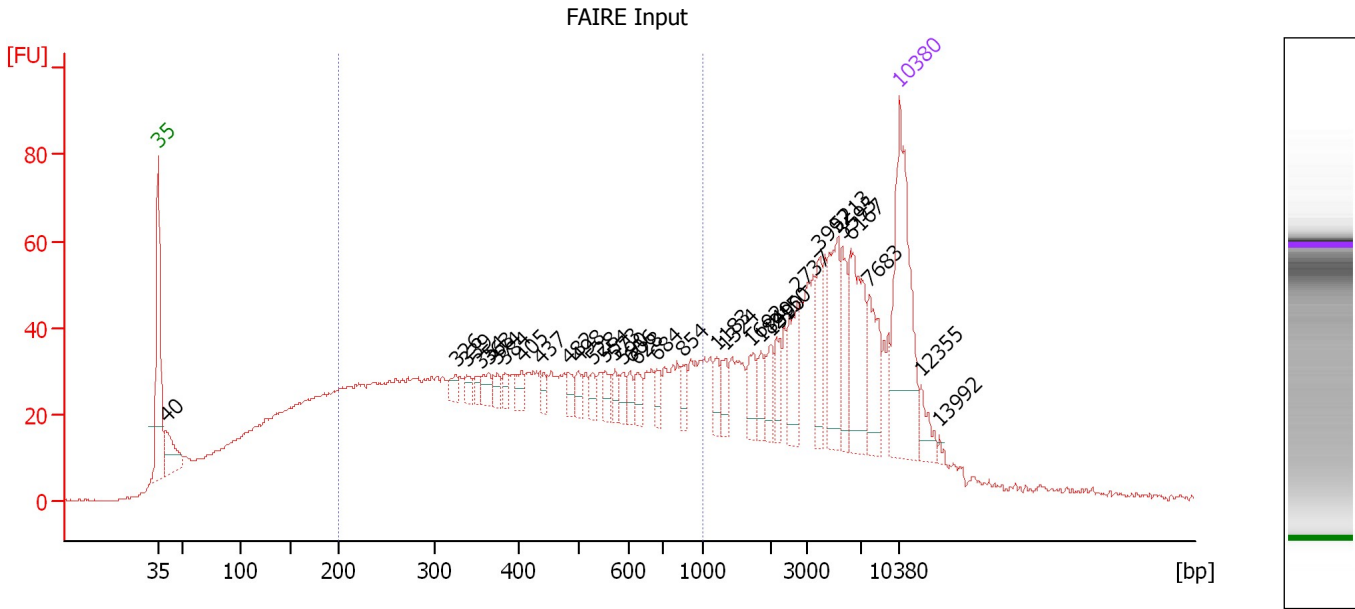
FAIRE

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
200	392	1,000	1,535.5	9,528.5	47	43.3	1,986.75	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : FAIRE Input

Number of peaks found: 33 Corr. Area 1: 1,322.1
 Noise: 0.2

Peak table for sample 5 : FAIRE Input

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	40	37.36	1,425.6		43.71
3	326	6.80	31.6		71.15
4	339	7.29	32.6		72.16
5	354	5.28	22.6		73.37
6	363	9.25	38.6		74.12
7	374	6.31	25.5		74.98
8	384	6.08	24.0		75.78
9	405	8.65	32.4		77.35
10	437	6.53	22.6		79.16
11	482	7.30	22.9		81.73
12	498	8.00	24.3		82.63
13	528	6.73	19.3		84.04
14	554	7.63	20.9		85.25
15	573	7.02	18.5		86.16
16	590	7.33	18.8		86.91
17	606	6.77	16.9		87.57
18	628	9.06	21.9		88.27
19	684	7.85	17.4		90.09
20	854	8.22	14.6		92.60
21	1,183	8.39	10.7		95.68
22	1,324	10.08	11.5		96.58
23	1,692	12.76	11.4		98.95
24	1,849	9.10	7.5		99.96
25	1,990	10.27	7.8		100.86

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad


Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...

... Peak table for sample 5 : FAIRE Input

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,150	9.74	6.9		101.42
27	2,737	22.51	12.5		103.33
28	3,992	19.63	7.5		105.50
29	5,213	35.46	10.3		107.11
30	5,595	17.80	4.8		107.61
31	6,167	44.41	10.9		108.37
32	7,683	27.06	5.3		110.18
33	10,380	75.00	10.9	Upper Marker	113.00
34	12,355	0.00	0.0		115.06
35	13,992	0.00	0.0		116.78

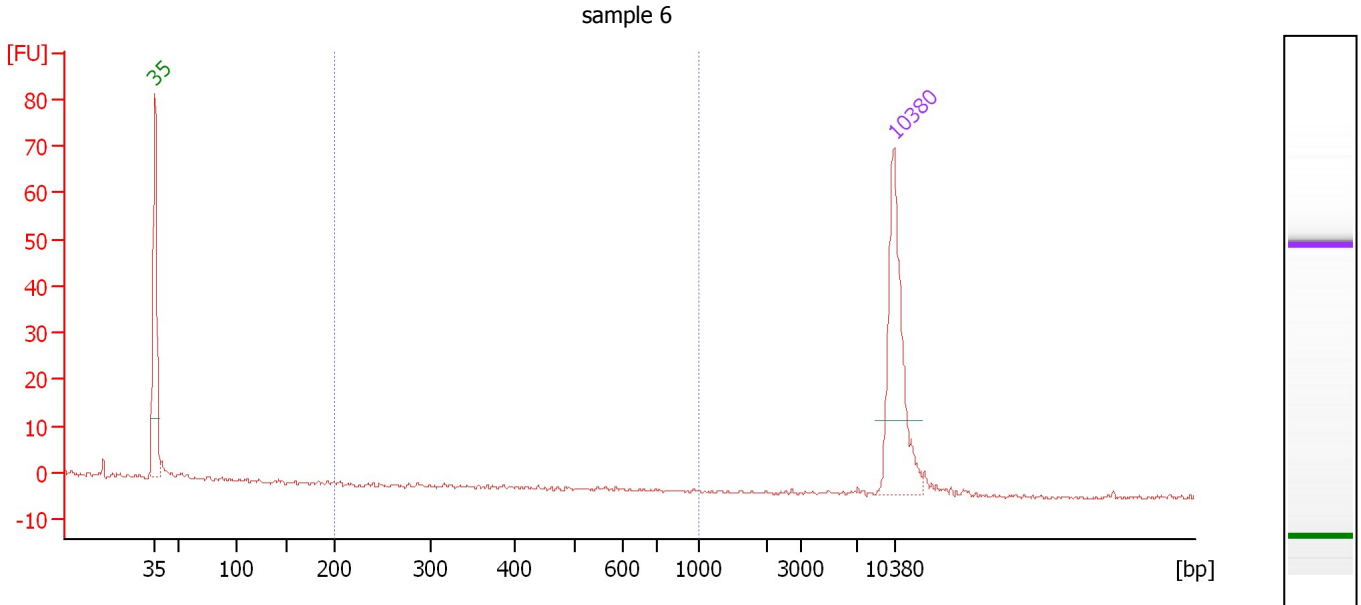
Region table for sample 5 : FAIRE Input

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
200	459	1,000	1,322.1	5,009.3	49	42.9	1,184.24	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...



Overall Results for sample 6 : sample 6

Number of peaks found: 0 Corr. Area 1: 0.0
 Noise: 0.3

Peak table for sample 6 : sample 6

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

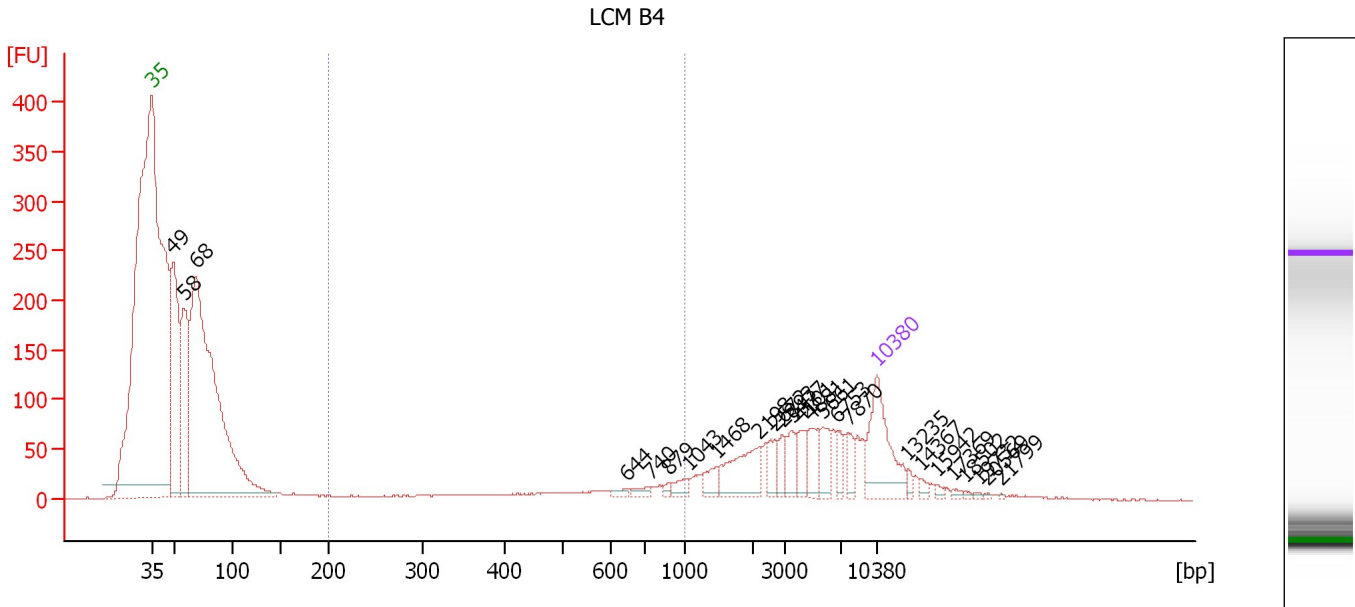
Region table for sample 6 : sample 6

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
200	0	1,000	0.0	0.0	0	0.0	0.00	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...



Overall Results for sample 7 : LCM B4

Number of peaks found: 25 Corr. Area 1: 282.2
 Noise: 0.3

Peak table for sample 7 : LCM B4

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	49	312.25	9,708.6		45.06
3	58	219.11	5,761.9		46.09
4	68	899.23	19,930.0		47.27
5	644	5.22	12.3		88.81
6	740	7.69	15.7		91.13
7	879	3.57	6.2		92.93
8	1,043	10.92	15.9		94.78
9	1,468	15.19	15.7		97.51
10	2,198	56.42	38.9		101.57
11	2,672	18.07	10.2		103.12
12	2,893	15.05	7.9		103.84
13	3,477	21.43	9.3		104.82
14	4,101	20.91	7.7		105.64
15	4,881	27.00	8.4		106.67
16	5,661	22.70	6.1		107.70
17	6,753	15.02	3.4		109.14
18	7,870	13.01	2.5		110.37
19	10,380	75.00	10.9	Upper Marker	113.00
20	13,235	0.00	0.0		115.99
21	14,367	0.00	0.0		117.17
22	15,942	0.00	0.0		118.82
23	17,369	0.00	0.0		120.31
24	18,502	0.00	0.0		121.49
25	19,732	0.00	0.0		122.78

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...**... Peak table for sample 7 : LCM B4**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	20,569	0.00	0.0		123.65
27	21,799	0.00	0.0		124.94

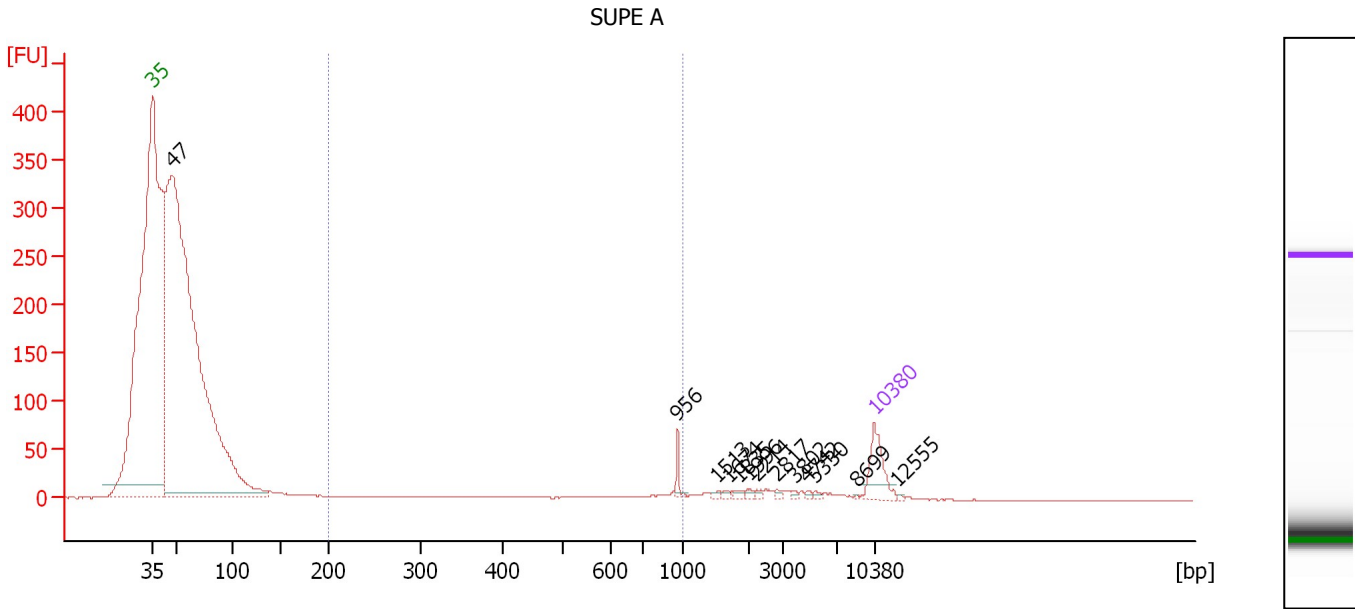
Region table for sample 7 : LCM B4

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/ μ l]	Color
200	570	1,000	282.2	431.1	8	38.8	124.17	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...



Overall Results for sample 8 : SUPE A

Number of peaks found: 13 Corr. Area 1: 97.4
 Noise: 0.3

Peak table for sample 8 : SUPE A

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	47	4,164.82	133,094.9		44.86
3	956	22.00	34.9		93.92
4	1,513	4.90	4.9		97.80
5	1,674	5.78	5.2		98.83
6	1,835	7.46	6.2		99.87
7	1,996	6.09	4.6		100.90
8	2,214	5.08	3.5		101.63
9	2,817	5.70	3.1		103.59
10	3,802	3.95	1.6		105.25
11	4,742	4.09	1.3		106.49
12	5,330	4.39	1.2		107.26
13	8,699	1.91	0.3		111.24
14	10,380	75.00	10.9	Upper Marker	113.00
15	12,555	0.00	0.0		115.27

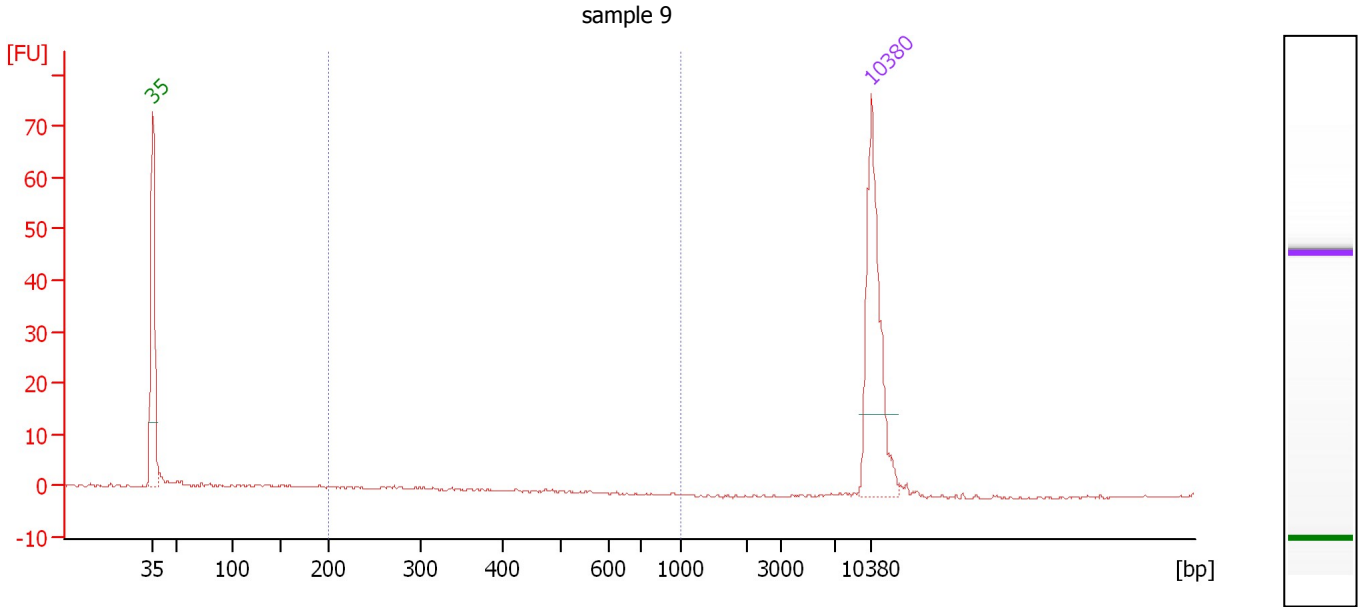
Region table for sample 8 : SUPE A

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
200	636	1,000	97.4	357.9	3	43.4	101.26	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...



Overall Results for sample 9 : sample 9

Number of peaks found: 0 Corr. Area 1: 1.5
 Noise: 0.2

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

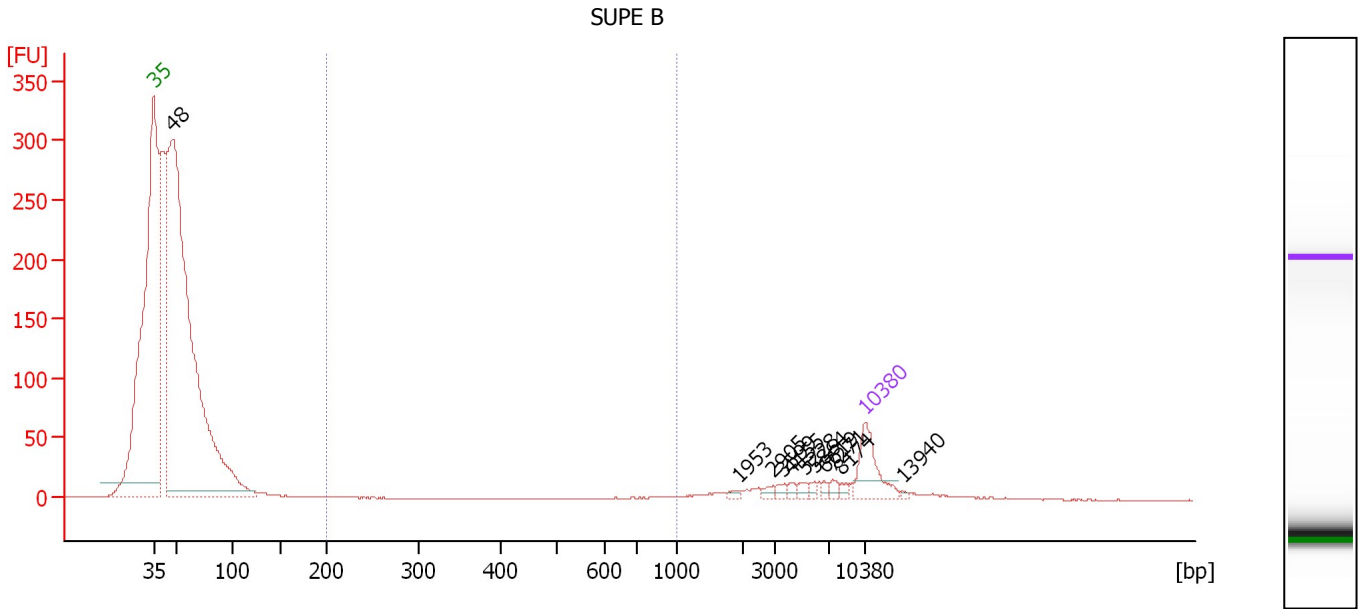
Region table for sample 9 : sample 9

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
200	276	1,000	1.5	10.3	10	20.9	1.80	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Electropherogram Summary Continued ...



Overall Results for sample 10 : SUPE B

Number of peaks found: 11 Corr. Area 1: 14.8
 Noise: 0.2

Peak table for sample 10 : SUPE B

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	48	2,611.61	83,149.3		44.89
3	1,953	5.57	4.3		100.63
4	2,905	8.27	4.3		103.88
5	3,599	8.57	3.6		104.98
6	4,155	8.14	3.0		105.71
7	5,228	10.14	2.9		107.13
8	5,864	5.94	1.5		107.97
9	6,619	6.65	1.5		108.96
10	7,271	8.97	1.9		109.75
11	8,174	6.51	1.2		110.69
12	10,380	75.00	10.9	Upper Marker	113.00
13	13,940	0.00	0.0		116.72

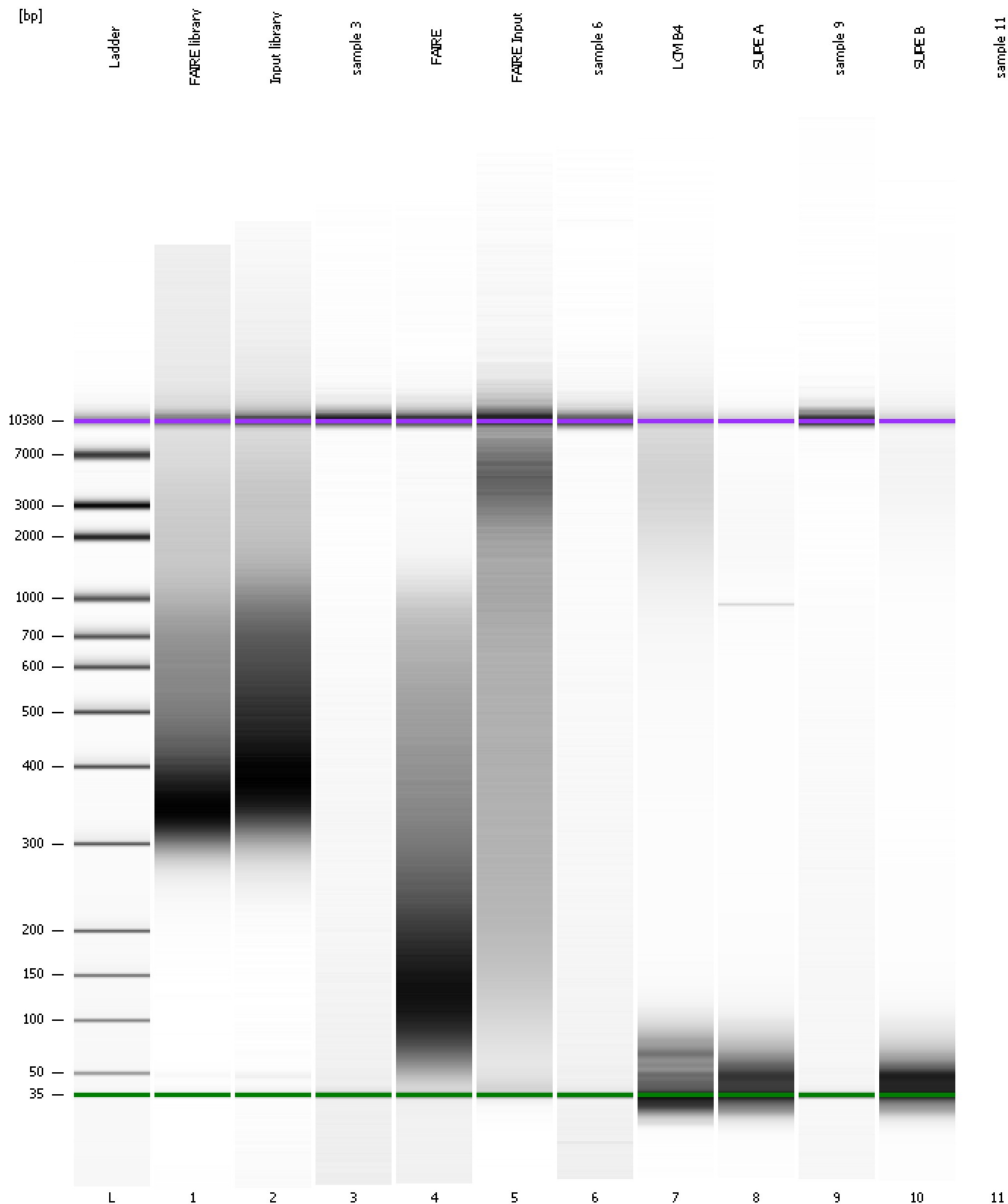
Region table for sample 10 : SUPE B

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
200	671	1,000	14.8	39.0	1	32.4	12.91	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
Modified: 4/17/2015 11:04:33 AM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
Modified: 4/17/2015 11:04:33 AM

Invalid Samples

Sample 11 has not been run, no results available.

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad

Created: 4/17/2015 10:26:01 AM
 Modified: 4/17/2015 11:04:33 AM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 11)		Instrument	Run		4/17/2015 11:04:29 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-04-17\2015-04-17_001.xad)		Instrument	Run		4/17/2015 10:26:07 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		4/17/2015 10:26:07 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		4/17/2015 10:26:07 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		4/17/2015 10:26:07 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		4/17/2015 10:26:07 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		4/17/2015 10:26:07 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		4/17/2015 10:26:07 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1