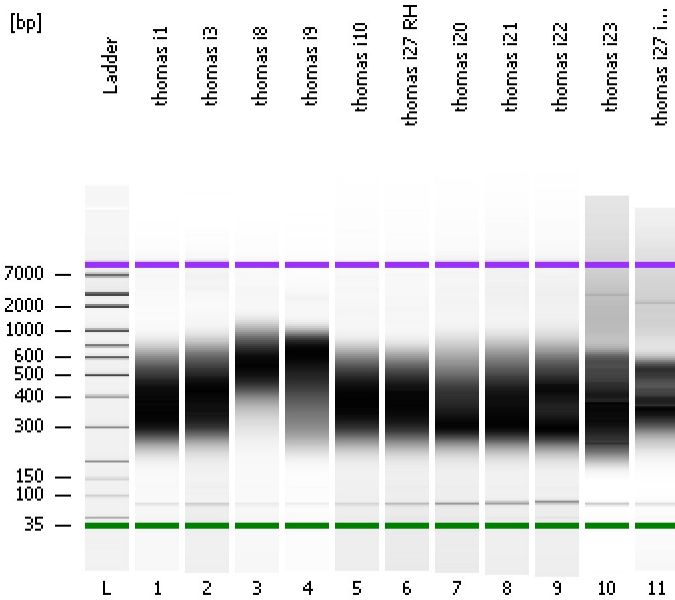


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
Modified: 3/7/2016 10:41:16 AM

Electrophoresis File Run Summary



Instrument Information:

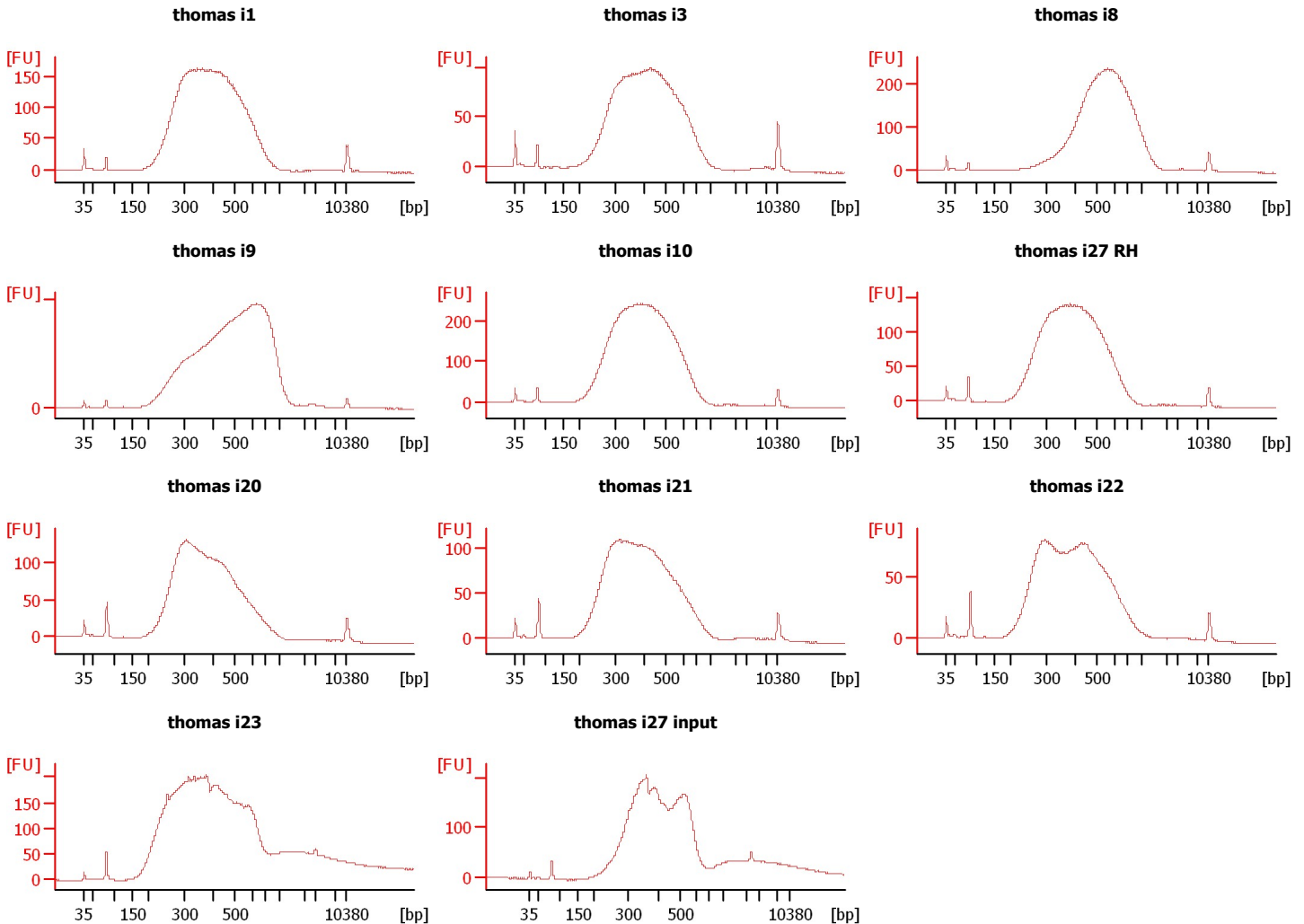
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
thomas i1		<input type="checkbox"/>	✓			
thomas i3		<input type="checkbox"/>	✓			
thomas i8		<input type="checkbox"/>	✓			
thomas i9		<input type="checkbox"/>	✓			
thomas i10		<input type="checkbox"/>	✓			
thomas i27 RH		<input type="checkbox"/>	✓			
thomas i20		<input type="checkbox"/>	✓			
thomas i21		<input type="checkbox"/>	✓			
thomas i22		<input type="checkbox"/>	✓			
thomas i23		<input type="checkbox"/>	✓			
thomas i27 input		<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\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
Modified: 3/7/2016 10:41:16 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

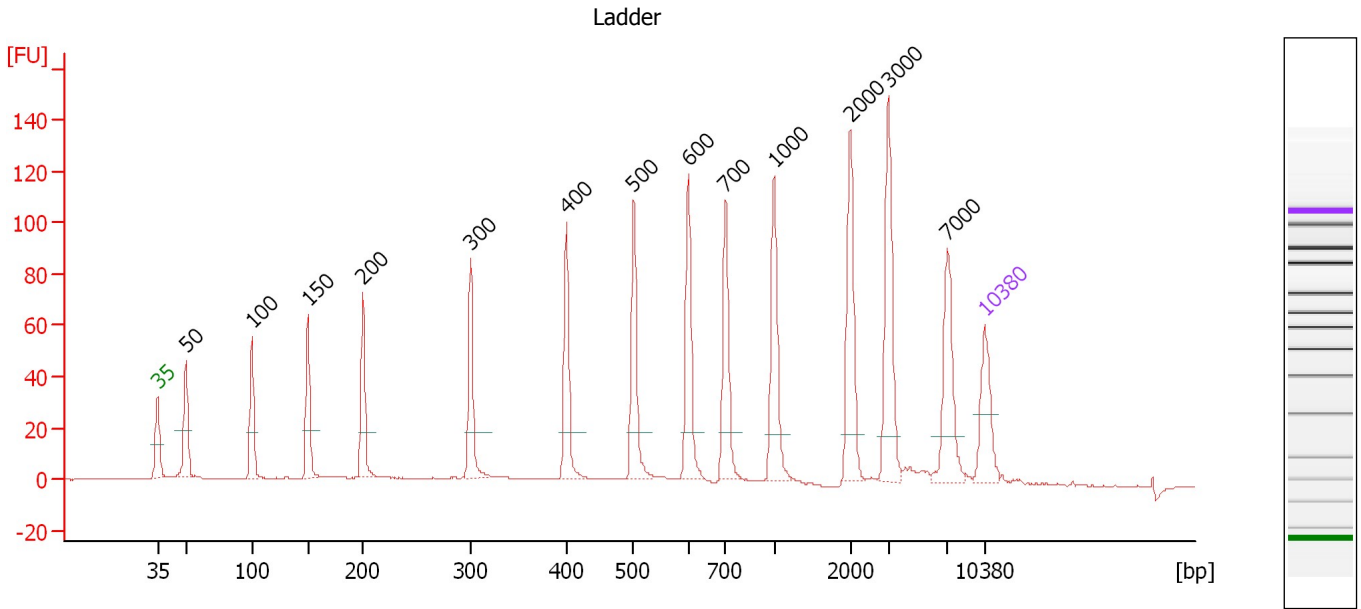
Ladder

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:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

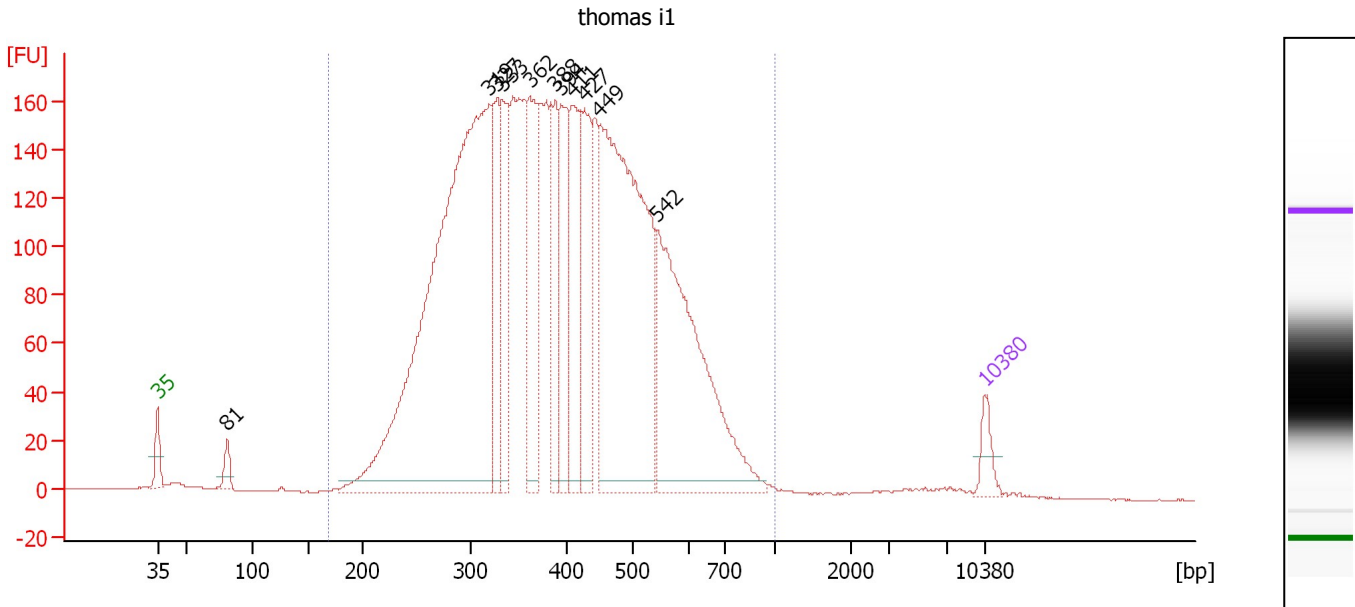
Peak table for Ladder

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	45.41
3	100	150.00	2,272.7	Ladder Peak	51.01
4	150	150.00	1,515.2	Ladder Peak	55.70
5	200	150.00	1,136.4	Ladder Peak	60.38
6	300	150.00	757.6	Ladder Peak	69.50
7	400	150.00	568.2	Ladder Peak	77.61
8	500	150.00	454.5	Ladder Peak	83.31
9	600	150.00	378.8	Ladder Peak	87.94
10	700	150.00	324.7	Ladder Peak	91.08
11	1,000	150.00	227.3	Ladder Peak	95.19
12	2,000	150.00	113.6	Ladder Peak	101.66
13	3,000	150.00	75.8	Ladder Peak	104.89
14	7,000	150.00	32.5	Ladder Peak	109.86
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : thomas i1

Number of peaks found: 11 Corr. Area 1: 4,594.8
 Noise: 0.2

Peak table for sample 1 : thomas i1

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	81	89.83	1,677.3		48.90
3	319	3,953.64	18,775.4		71.05
4	327	493.19	2,281.9		71.73
5	333	519.37	2,363.9		72.17
6	362	580.00	2,429.2		74.51
7	388	370.15	1,444.6		76.66
8	394	502.04	1,929.5		77.15
9	411	599.50	2,212.0		78.22
10	427	503.11	1,785.6		79.15
11	449	2,171.05	7,323.5		80.41
12	542	1,445.76	4,043.7		85.24
13	10,380	75.00	10.9	Upper Marker	113.00

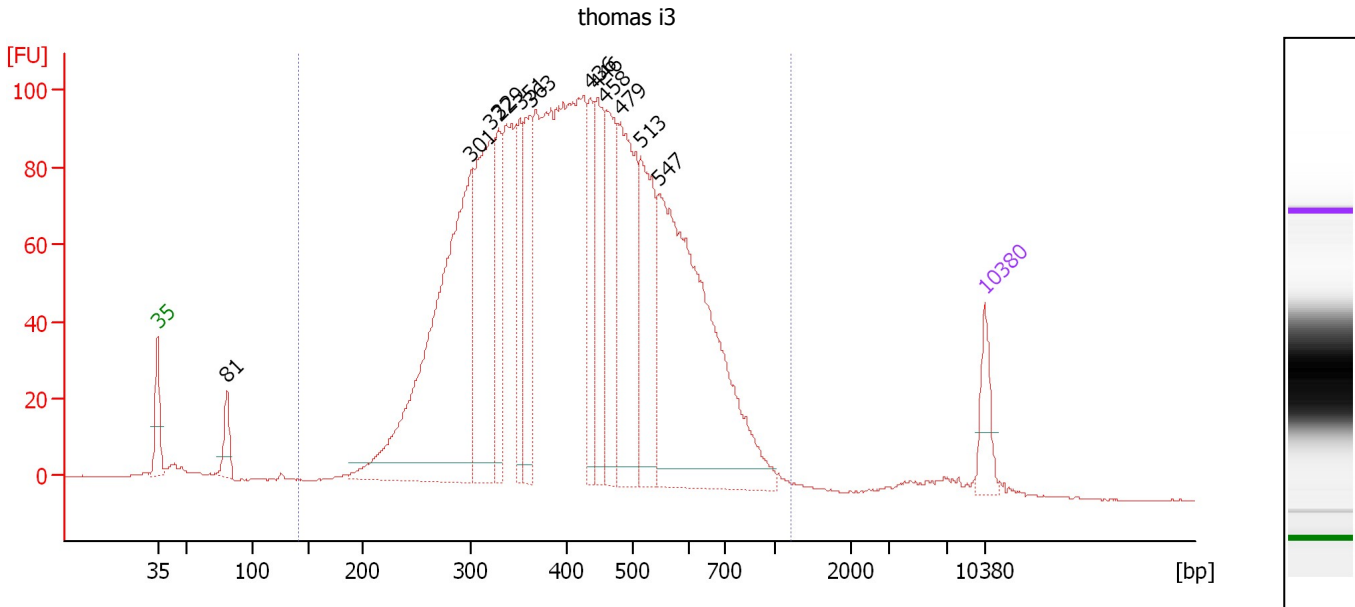
Region table for sample 1 : thomas i1

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
169	1,000	407	4,594.8	55,505.9	13,325.50	98	28.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 : thomas i3

Number of peaks found: 12 Corr. Area 1: 2,804.6
 Noise: 0.3

Peak table for sample 2 : thomas i3

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	81	90.76	1,699.2		48.88
3	301	1,314.91	6,628.3		69.55
4	322	574.22	2,704.3		71.26
5	329	238.75	1,099.6		71.85
6	351	217.51	938.1		73.66
7	363	263.87	1,100.2		74.64
8	436	202.03	702.7		79.64
9	446	275.03	934.5		80.23
10	458	309.06	1,022.6		80.91
11	479	499.76	1,579.4		82.14
12	513	342.35	1,011.6		83.90
13	547	1,142.39	3,166.6		85.47
14	10,380	75.00	10.9	Upper Marker	113.00

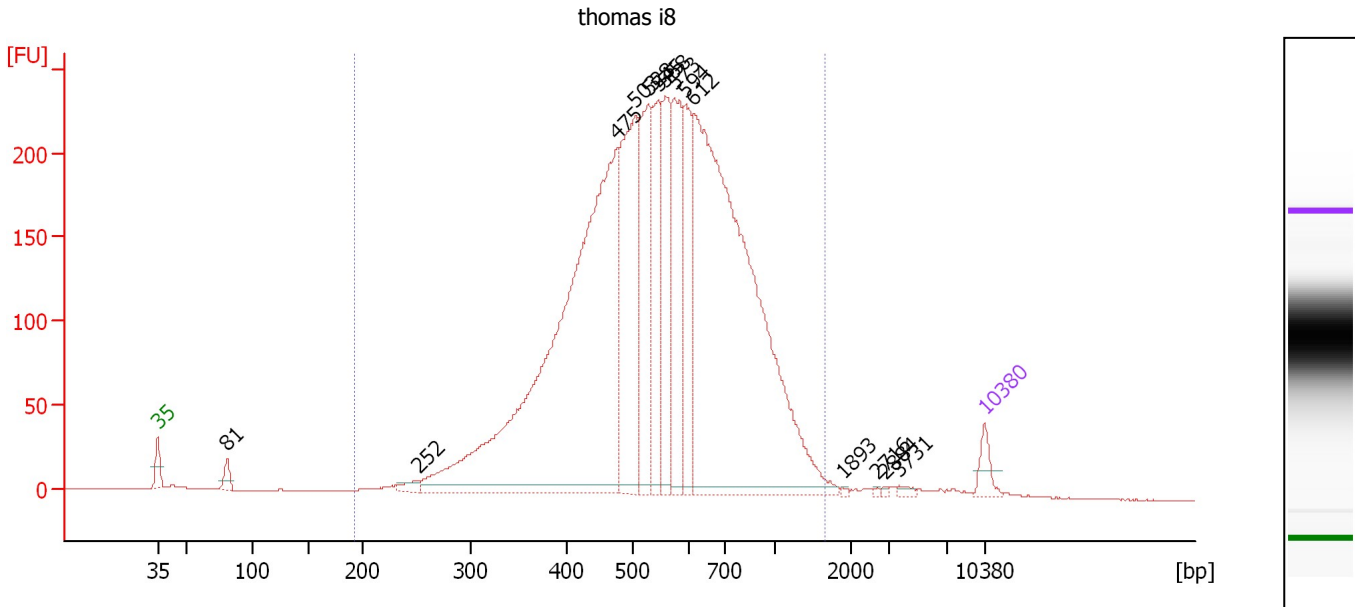
Region table for sample 2 : thomas i3

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
143	1,212	431	2,804.6	29,570.7	7,434.80	97	30.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...



Overall Results for sample 3 : thomas i8

Number of peaks found: 14 Corr. Area 1: 4,723.3
 Noise: 0.2

Peak table for sample 3 : thomas i8

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	81	79.97	1,498.4		48.87
3	252	52.08	313.5		65.10
4	475	4,049.54	12,923.7		81.87
5	503	1,236.76	3,725.1		83.45
6	528	734.60	2,109.9		84.59
7	545	624.73	1,738.2		85.37
8	558	584.75	1,586.7		86.02
9	573	669.04	1,768.2		86.71
10	594	511.05	1,304.6		87.64
11	612	3,698.87	9,151.1		88.33
12	1,893	6.31	5.0		100.96
13	2,716	5.94	3.3		103.97
14	2,884	6.72	3.5		104.52
15	3,731	17.71	7.2		105.80
16	10,380	75.00	10.9	Upper Marker	113.00

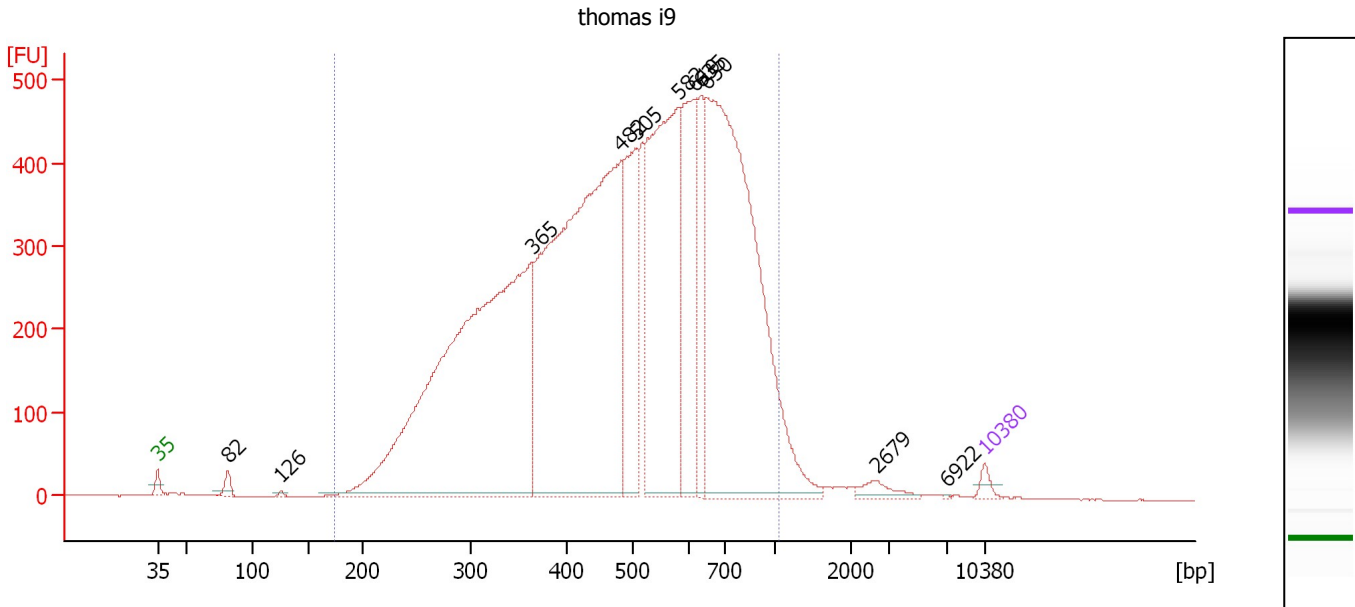
Region table for sample 3 : thomas i8

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
193	1,666	585	4,723.3	37,009.6	12,434.31	98	35.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...



Overall Results for sample 4 : thomas i9

Number of peaks found: 11 Corr. Area 1: 12,665.2
 Noise: 0.2

Peak table for sample 4 : thomas i9

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	82	145.96	2,707.0		48.96
3	126	21.60	259.5		53.46
4	365	9,330.80	38,712.9		74.79
5	482	8,971.61	28,216.5		82.27
6	505	1,751.27	5,256.1		83.53
7	582	4,140.27	10,773.7		87.12
8	619	1,899.01	4,649.0		88.54
9	635	997.16	2,379.3		89.04
10	650	7,077.34	16,509.9		89.50
11	2,679	141.24	79.9		103.85
12	6,922	6.33	1.4		109.77
13	10,380	75.00	10.9	Upper Marker	113.00

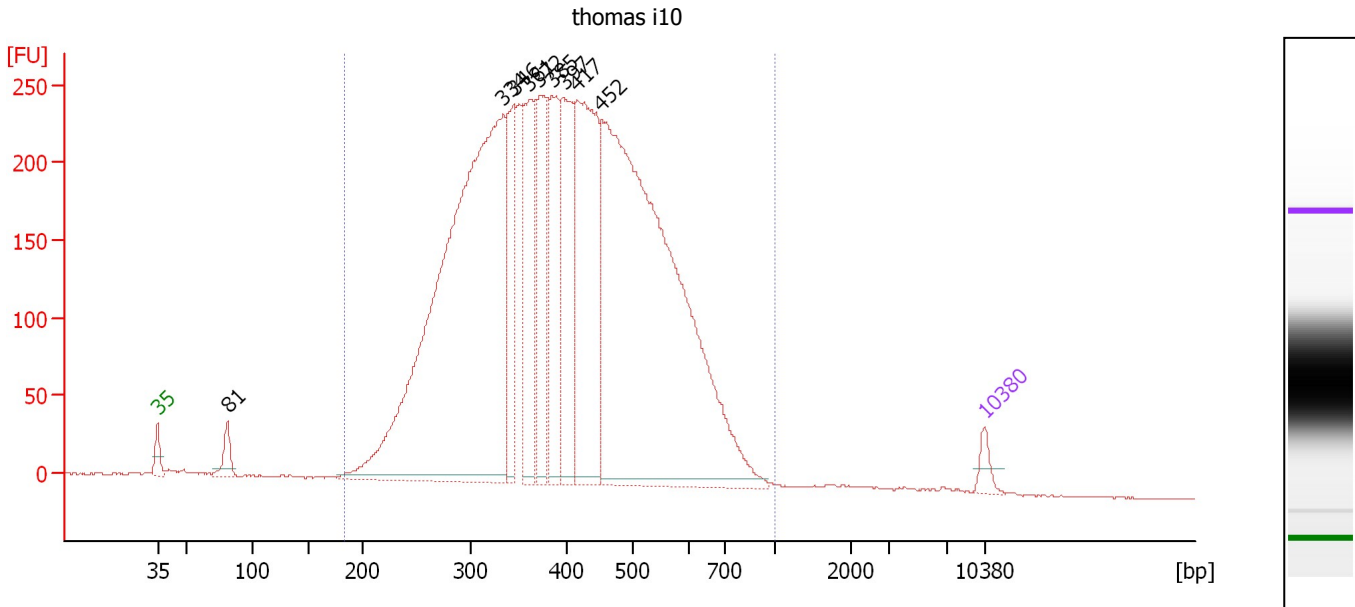
Region table for sample 4 : thomas i9

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
174	1,066	514	12,665.2	128,242.1	36,368.70	97	34.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : thomas i10

Number of peaks found: 9 Corr. Area 1: 6,599.5
 Noise: 0.5

Peak table for sample 5 : thomas i10

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	81	199.28	3,710.8		48.93
3	334	6,211.21	28,159.3		72.28
4	346	693.00	3,038.7		73.20
5	361	1,072.41	4,497.4		74.47
6	372	867.76	3,534.3		75.34
7	385	842.27	3,312.7		76.42
8	397	1,042.14	3,975.3		77.39
9	417	1,917.68	6,973.7		78.56
10	452	5,851.22	19,630.1		80.55
11	10,380	75.00	10.9	Upper Marker	113.00

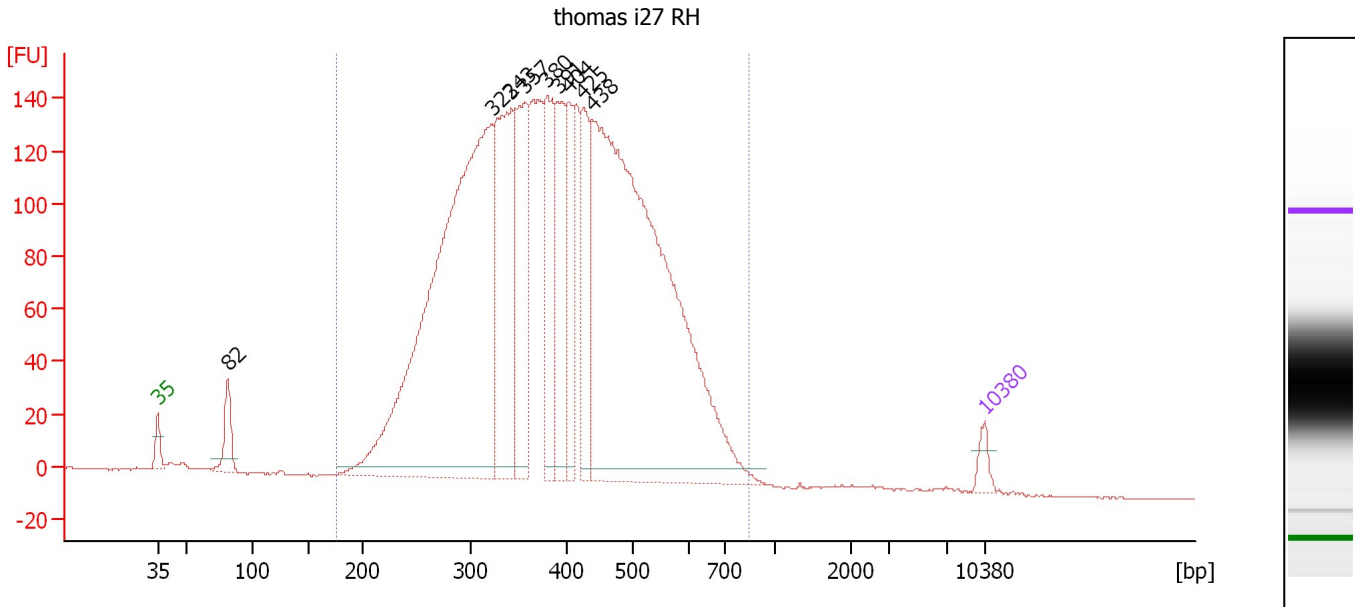
Region table for sample 5 : thomas i10

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
184	1,000	412	6,599.5	79,217.8	19,390.82	98	27.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...



Overall Results for sample 6 : thomas i27 RH

Number of peaks found: 9 Corr. Area 1: 3,798.5
 Noise: 0.2

Peak table for sample 6 : thomas i27 RH

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	82	314.61	5,843.7		48.95
3	322	5,647.29	26,560.4		71.30
4	343	1,722.59	7,618.9		72.96
5	357	1,078.50	4,582.4		74.09
6	380	896.54	3,572.8		76.01
7	391	779.55	3,020.4		76.89
8	404	581.76	2,184.0		77.82
9	425	643.80	2,293.0		79.06
10	438	5,873.26	20,311.8		79.78
11	10,380	75.00	10.9	Upper Marker	113.00

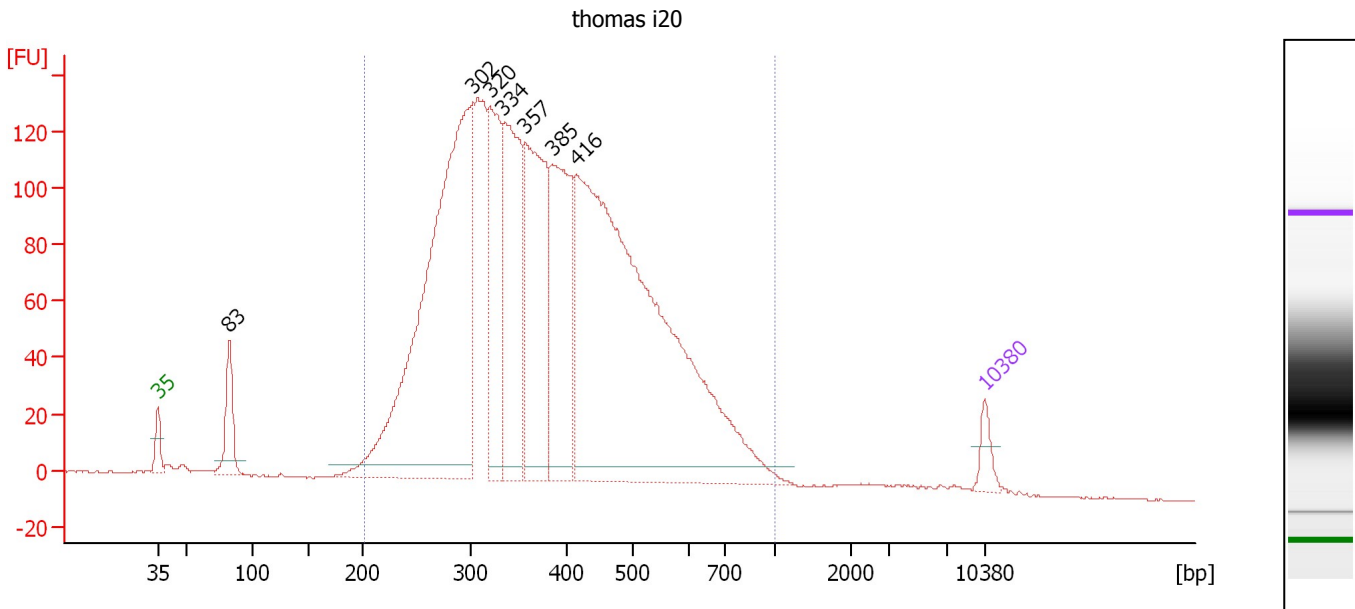
Region table for sample 6 : thomas i27 RH

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
177	845	402	3,798.5	80,118.9	19,224.99	98	26.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...



Overall Results for sample 7 : thomas i20

Number of peaks found: 7 Corr. Area 1: 3,298.0
 Noise: 0.2

Peak table for sample 7 : thomas i20

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	83	327.86	6,019.1		49.06
3	302	3,389.93	16,980.7		69.70
4	320	802.35	3,798.8		71.13
5	334	1,140.52	5,169.2		72.29
6	357	1,193.23	5,063.8		74.13
7	385	1,044.00	4,109.1		76.39
8	416	4,066.89	14,827.9		78.50
9	10,380	75.00	10.9	Upper Marker	113.00

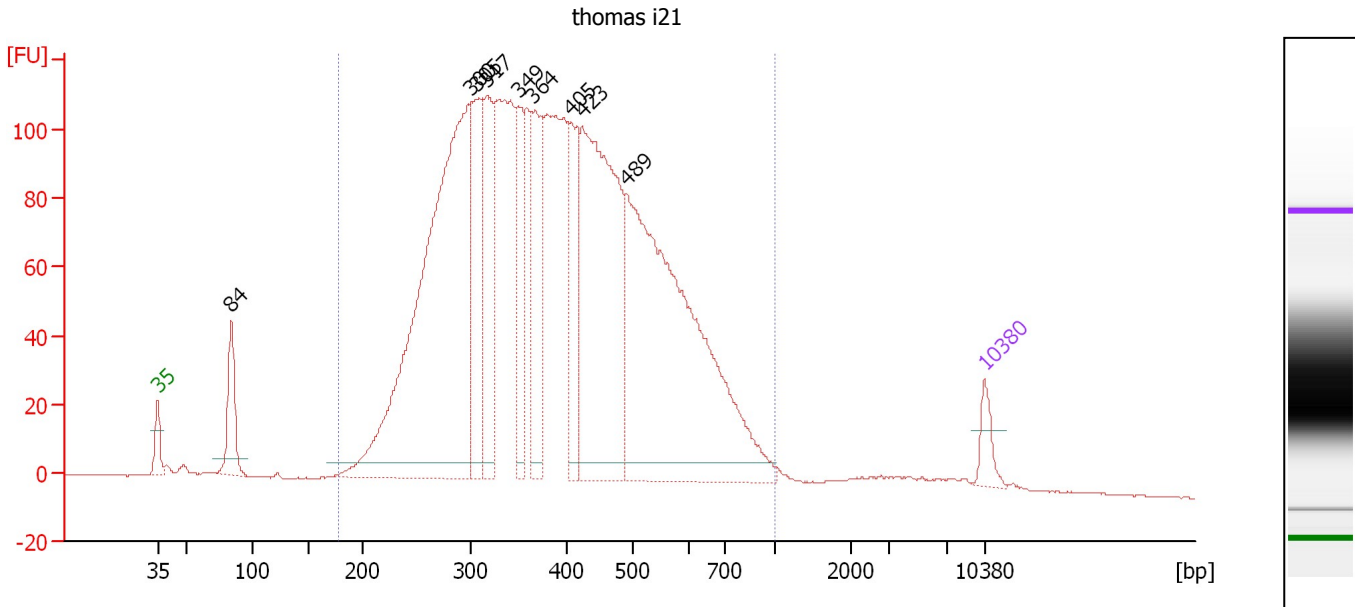
Region table for sample 7 : thomas i20

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
201	1,000	397	3,298.0	53,615.8	12,545.50	96	30.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...



Overall Results for sample 8 : thomas i21

Number of peaks found: 9 Corr. Area 1: 3,094.3
 Noise: 0.2

Peak table for sample 8 : thomas i21

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	84	323.83	5,837.1		49.23
3	300	3,040.91	15,354.3		69.51
4	305	614.97	3,051.7		69.94
5	317	566.82	2,708.0		70.89
6	349	393.79	1,708.1		73.50
7	364	546.98	2,274.3		74.73
8	405	436.89	1,632.9		77.92
9	423	1,648.12	5,901.6		78.93
10	489	2,423.89	7,503.0		82.71
11	10,380	75.00	10.9	Upper Marker	113.00

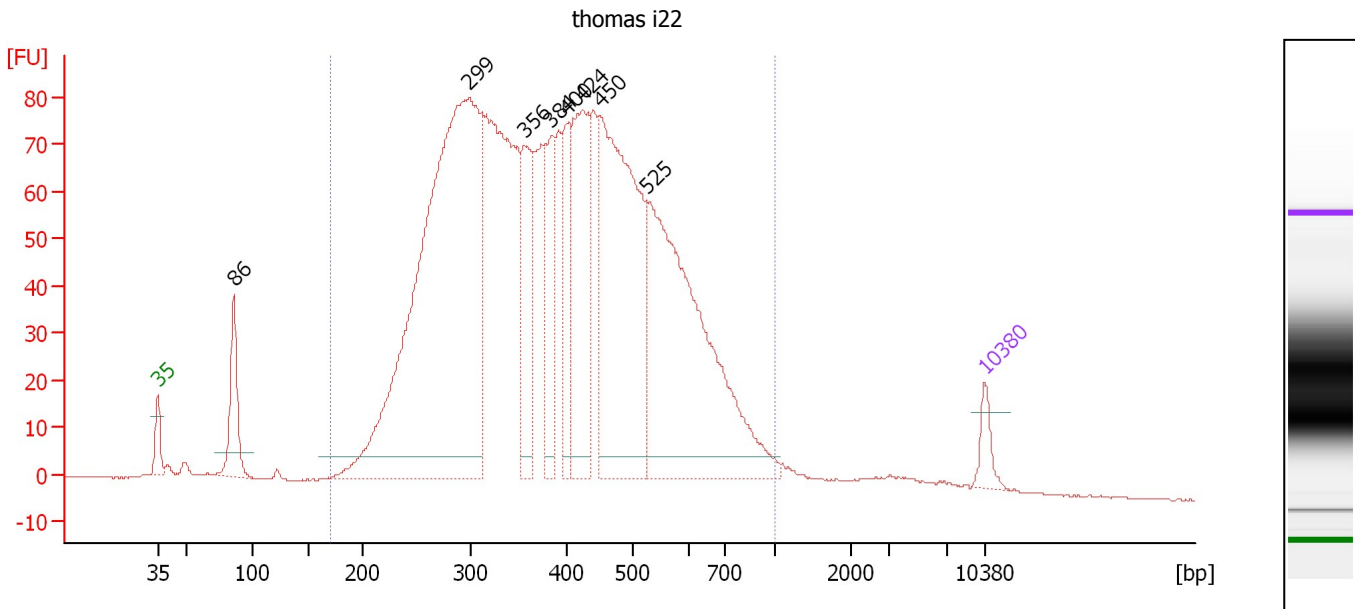
Region table for sample 8 : thomas i21

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
179	1,000	408	3,094.3	52,691.8	12,489.32	95	31.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...



Overall Results for sample 9 : thomas i22

Number of peaks found: 8 Corr. Area 1: 2,341.4
 Noise: 0.1

Peak table for sample 9 : thomas i22

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	86	396.98	7,008.2		49.43
3	299	4,123.92	20,909.9		69.40
4	356	544.25	2,314.4		74.07
5	384	367.31	1,447.5		76.35
6	400	343.53	1,300.4		77.63
7	424	802.13	2,869.4		78.96
8	450	1,727.60	5,821.1		80.44
9	525	1,935.31	5,582.9		84.48
10	10,380	75.00	10.9	Upper Marker	113.00

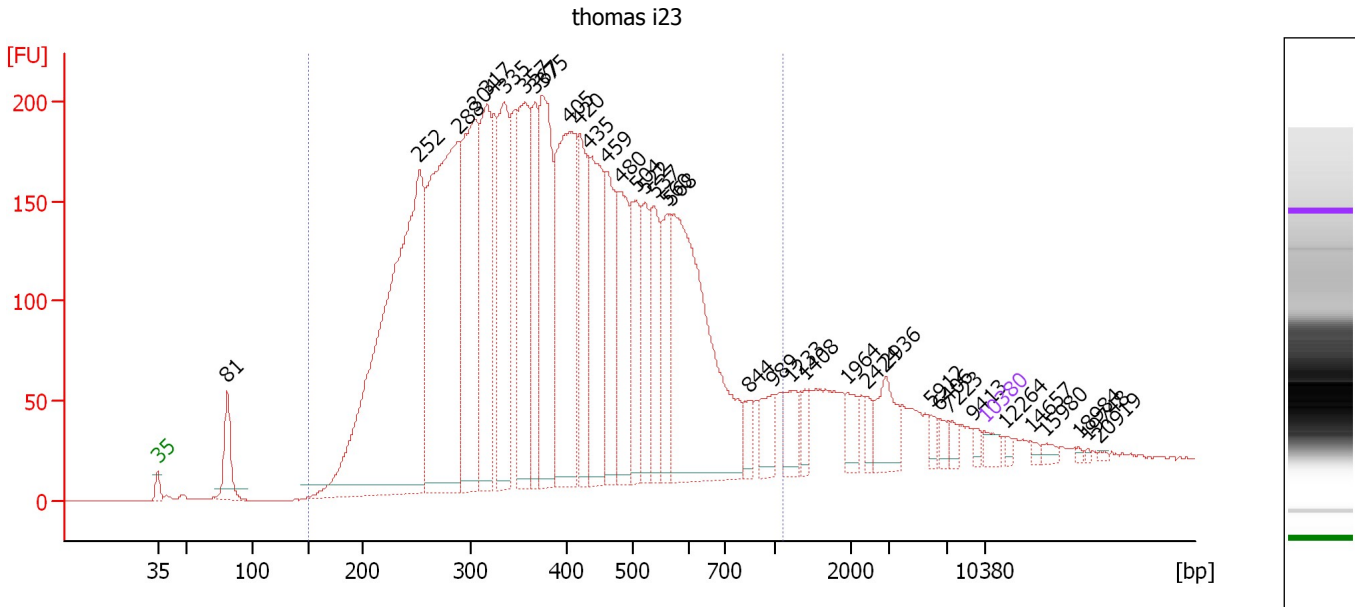
Region table for sample 9 : thomas i22

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
172	1,000	411	2,341.4	56,475.4	13,359.27	95	32.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...



Overall Results for sample 10 : thomas i23

Number of peaks found: 36 Corr. Area 1: 6,298.6
 Noise: 0.1

Peak table for sample 10 : thomas i23

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	81	447.84	8,356.5		48.91
3	252	5,158.15	30,954.7		65.17
4	288	3,623.49	19,048.3		68.43
5	304	1,914.14	9,532.9		69.85
6	317	1,387.97	6,632.9		70.89
7	335	1,338.27	6,050.6		72.35
8	357	1,281.35	5,443.0		74.10
9	367	707.71	2,920.4		74.95
10	375	1,400.31	5,652.8		75.61
11	405	1,875.77	7,022.6		77.88
12	420	883.12	3,188.6		78.73
13	435	1,157.98	4,037.3		79.58
14	459	801.16	2,646.7		80.95
15	480	786.27	2,480.8		82.18
16	504	615.16	1,848.5		83.51
17	522	558.72	1,623.1		84.31
18	537	564.91	1,594.3		85.02
19	560	503.16	1,360.6		86.11
20	568	2,383.73	6,353.2		86.48
21	844	127.96	229.7		93.05
22	989	209.76	321.3		95.04
23	1,233	195.61	240.4		96.69
24	1,408	98.95	106.4		97.83
25	1,964	147.72	114.0		101.42

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...

... Peak table for sample 10 : thomas i23

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,424	82.24	51.4		103.03
27	2,936	268.77	138.7		104.68
28	5,912	56.69	14.5		108.51
29	6,406	66.39	15.7		109.12
30	7,223	60.90	12.8		110.07
31	9,413	37.88	6.1		112.10
32	10,380	75.00	10.9	Upper Marker	113.00
33	12,264	0.00	0.0		114.75
34	14,657	0.00	0.0		116.97
35	15,980	0.00	0.0		118.20
36	18,984	0.00	0.0		120.99
37	19,748	0.00	0.0		121.70
38	20,919	0.00	0.0		122.78

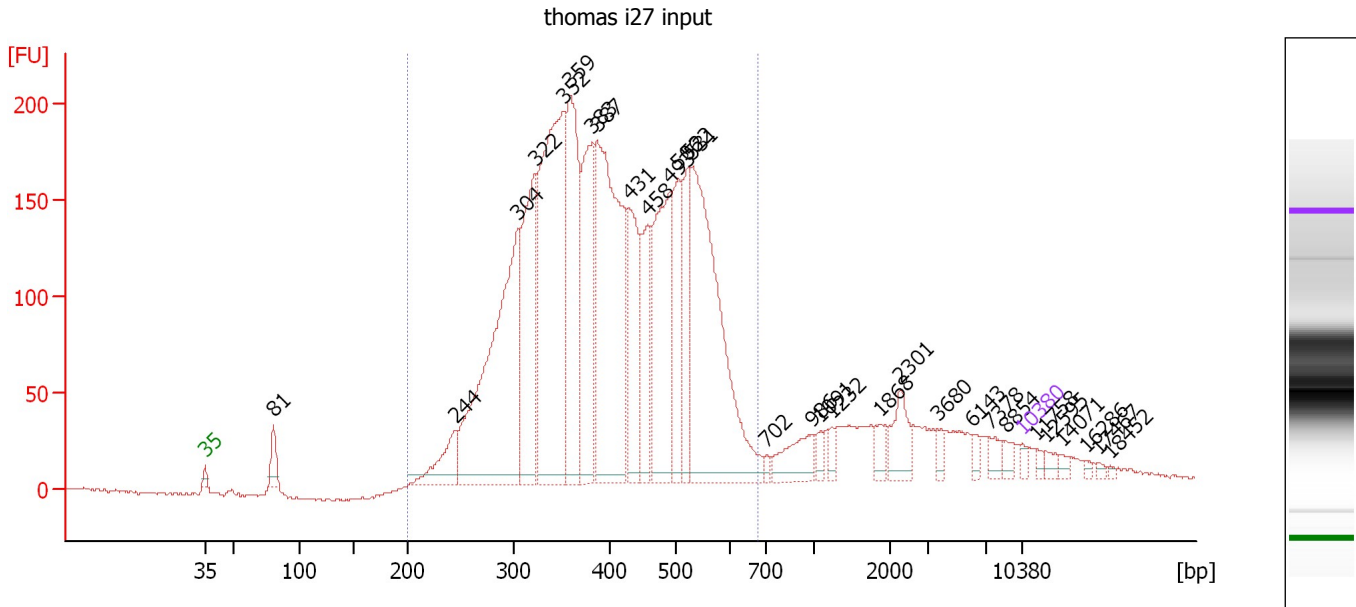
Region table for sample 10 : thomas i23

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
151	1,125	406	6,298.6	127,285.7	28,704.08	90	37.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...



Overall Results for sample 11 : thomas i27 input

Number of peaks found: 30 Corr. Area 1: 4,346.0
 Noise: 0.7

Peak table for sample 11 : thomas i27 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	81	543.52	10,208.7		48.85
3	244	1,017.80	6,310.1		64.43
4	304	7,230.99	36,049.4		69.82
5	322	3,576.93	16,833.9		71.28
6	352	7,345.87	31,586.6		73.75
7	359	3,771.11	15,910.2		74.30
8	383	3,131.97	12,396.9		76.22
9	387	6,486.87	25,377.4		76.58
10	431	1,981.76	6,969.3		79.37
11	458	1,527.41	5,051.7		80.92
12	493	3,481.92	10,709.7		82.89
13	506	1,729.52	5,181.9		83.57
14	522	1,454.37	4,217.7		84.35
15	531	6,325.61	18,038.2		84.76
16	702	95.17	205.3		91.11
17	986	708.14	1,088.0		95.00
18	1,091	159.09	221.0		95.77
19	1,232	167.28	205.7		96.69
20	1,868	256.97	208.4		100.80
21	2,301	569.29	374.9		102.63
22	3,680	152.92	63.0		105.73
23	6,143	125.03	30.8		108.80
24	7,378	212.21	43.6		110.21
25	8,854	152.57	26.1		111.58

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

Electropherogram Summary Continued ...

... Peak table for sample 11 : thomas i27 input

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	10,380	75.00	10.9	Upper Marker	113.00
27	11,758	0.00	0.0		114.28
28	12,595	0.00	0.0		115.06
29	14,071	0.00	0.0		116.43
30	16,286	0.00	0.0		118.48
31	17,467	0.00	0.0		119.58
32	18,452	0.00	0.0		120.49

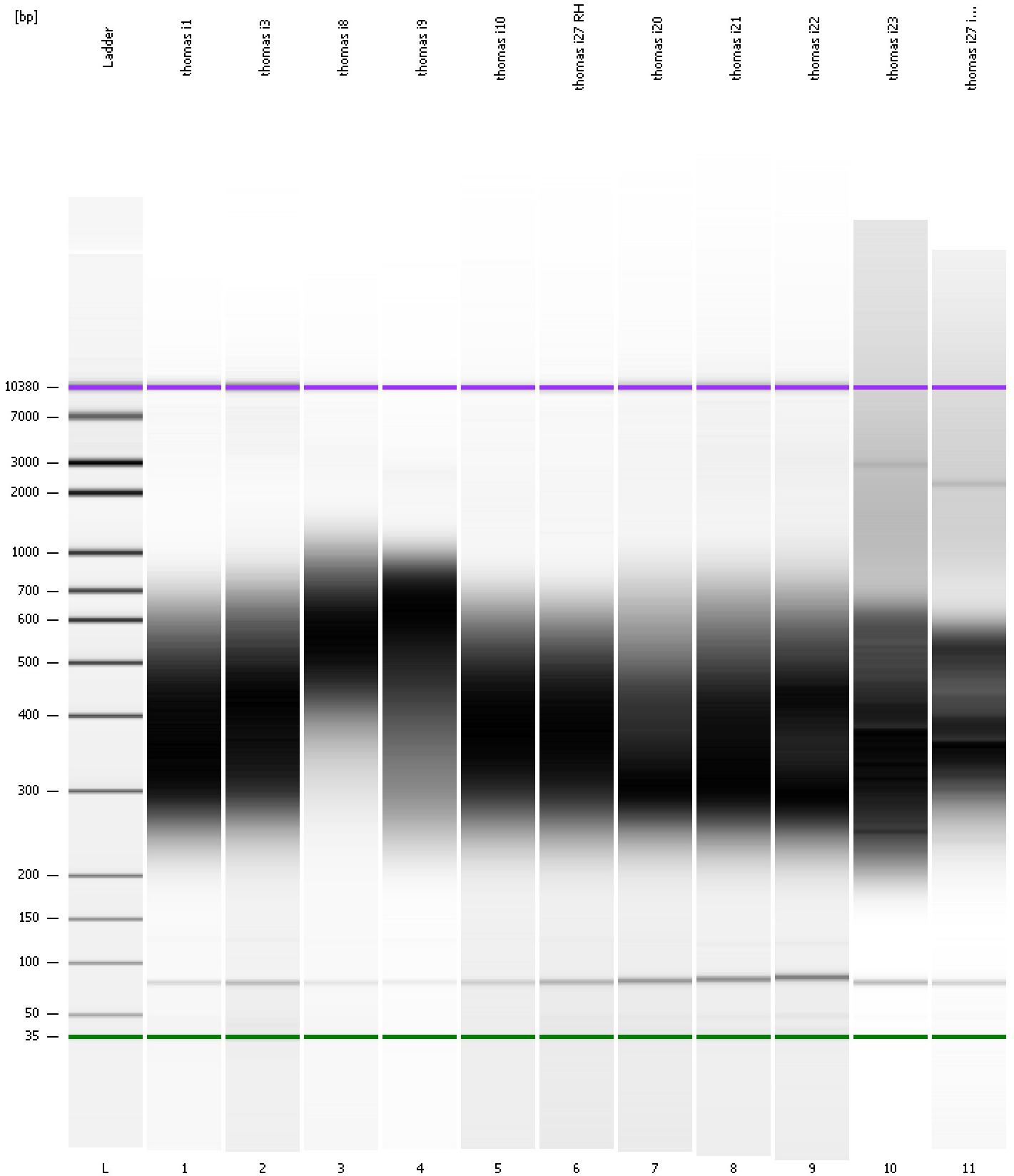
Region table for sample 11 : thomas i27 input

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	678	409	4,346.0	202,026.5	50,384.71	87	23.4

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
Modified: 3/7/2016 10:41:16 AM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad

Created: 3/7/2016 9:58:58 AM
 Modified: 3/7/2016 10:41:16 AM

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		3/7/2016 10:40:16 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-03-07\2016-03-07_001.xad)		Instrument	Run		3/7/2016 9:59:04 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		3/7/2016 9:59:04 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		3/7/2016 9:59:04 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		3/7/2016 9:59:04 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		3/7/2016 9:59:04 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		3/7/2016 9:59:04 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		3/7/2016 9:59:04 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1