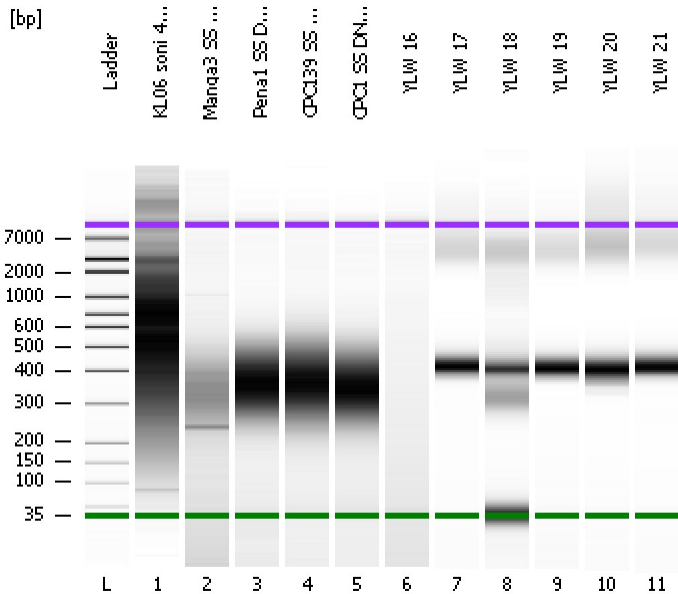


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
Modified: 12/10/2015 5:14:14 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

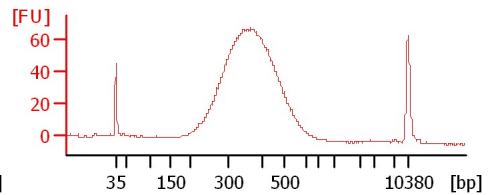
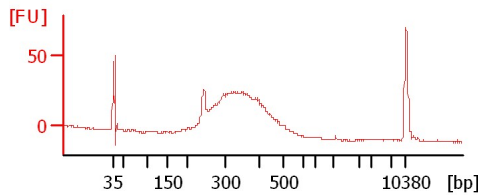
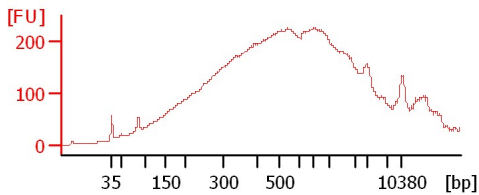
Chip Information:

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

KL06 soni 40 sec

Manga3 SS DNA (0.65X R/0.8X L)

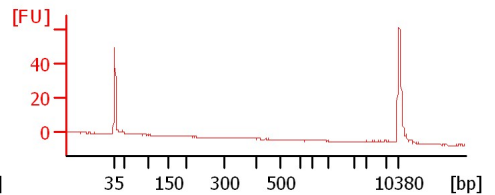
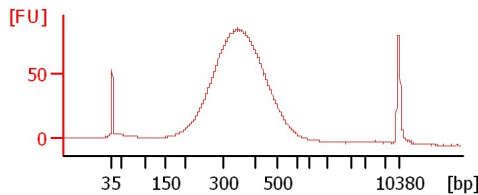
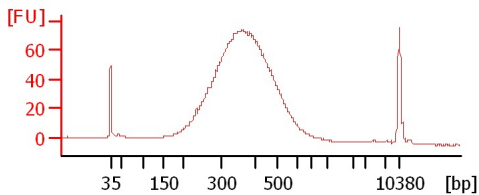
Pena1 SS DNA (0.65X R/0.8X L)



CPC139 SS DNA (0.65X R/0.8X L)

CPC1 SS DNA (0.65X R/0.8X L)

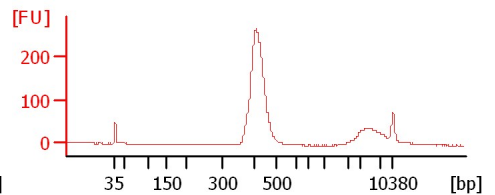
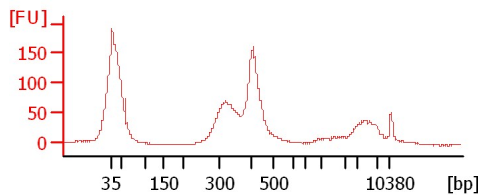
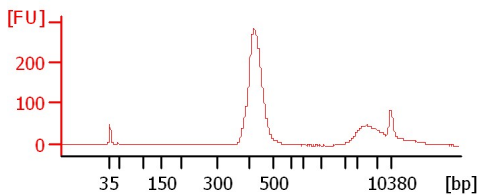
YLW 16



YLW 17

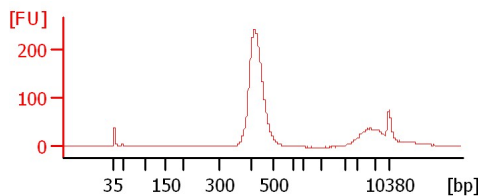
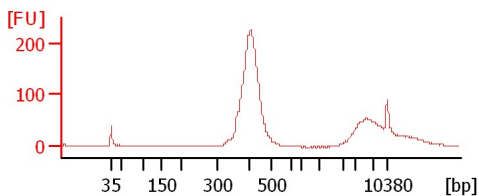
YLW 18

YLW 19



YLW 20

YLW 21



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
Modified: 12/10/2015 5:14:14 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
KL06 soni 40 sec		<input type="checkbox"/>	✓			
Manga3 SS DNA (0.65X R/0.8X L)		<input type="checkbox"/>	✓			
Pena1 SS DNA (0.65X R/0.8X L)		<input type="checkbox"/>	✓			
CPC139 SS DNA (0.65X R/0.8X L)		<input type="checkbox"/>	✓			
CPC1 SS DNA (0.65X R/0.8X L)		<input type="checkbox"/>	✓			
YLW 16		<input type="checkbox"/>	✓			
YLW 17		<input type="checkbox"/>	✓			
YLW 18		<input type="checkbox"/>	✓			
YLW 19		<input type="checkbox"/>	✓			
YLW 20		<input type="checkbox"/>	✓			
YLW 21		<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\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
Modified: 12/10/2015 5:14:14 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

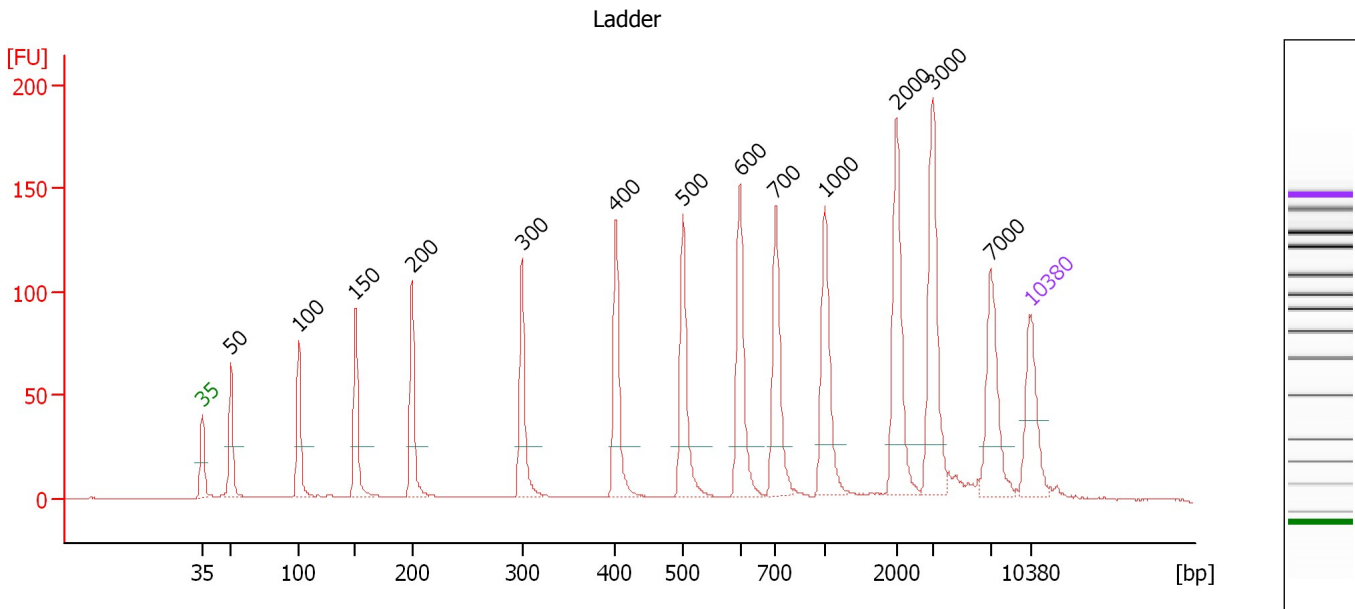
Ladder

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\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

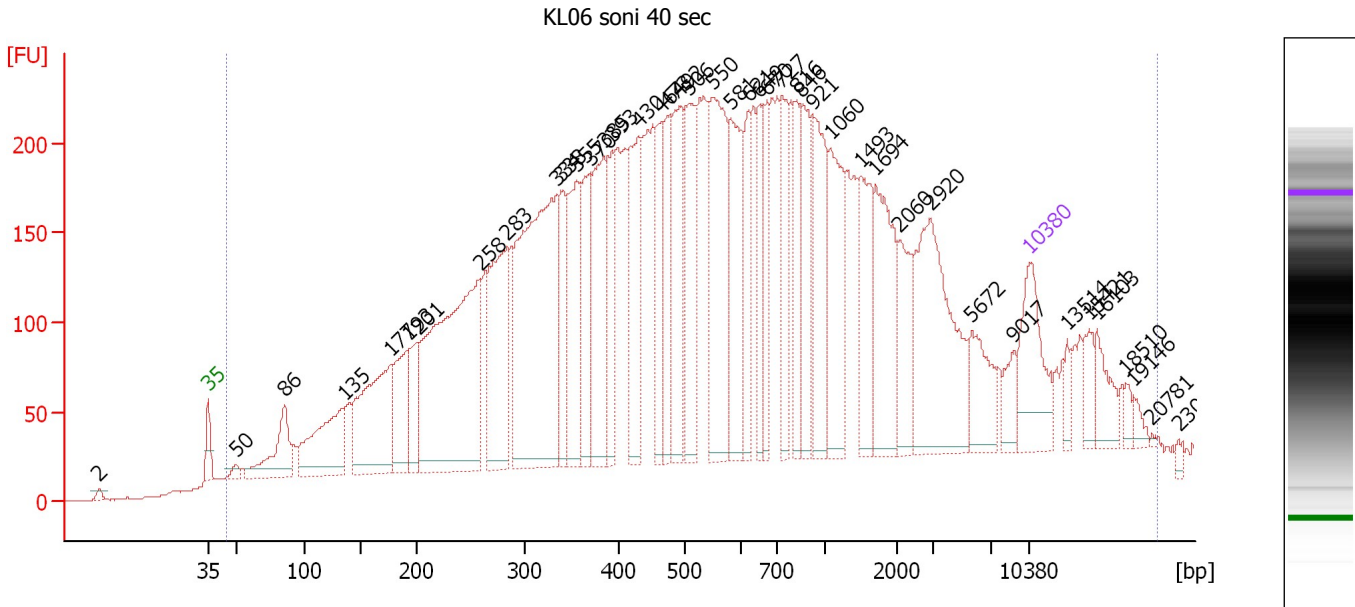
Peak table for Ladder

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	45.39
3	100	150.00	2,272.7	Ladder Peak	51.16
4	150	150.00	1,515.2	Ladder Peak	55.94
5	200	150.00	1,136.4	Ladder Peak	60.71
6	300	150.00	757.6	Ladder Peak	70.00
7	400	150.00	568.2	Ladder Peak	77.93
8	500	150.00	454.5	Ladder Peak	83.61
9	600	150.00	378.8	Ladder Peak	88.43
10	700	150.00	324.7	Ladder Peak	91.45
11	1,000	150.00	227.3	Ladder Peak	95.60
12	2,000	150.00	113.6	Ladder Peak	101.69
13	3,000	150.00	75.8	Ladder Peak	104.75
14	7,000	150.00	32.5	Ladder Peak	109.62
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : KL06 soni 40 sec

Number of peaks found: 43 Corr. Area 1: 10,841.7
 Noise: 0.3

Peak table for sample 1 : KL06 soni 40 sec


Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	2	0.00	0.0		33.69
2	35	125.00	5,411.3	Lower Marker	43.00
3	50	11.34	346.9		45.32
4	86	94.85	1,679.0		49.50
5	135	151.72	1,698.2		54.54
6	177	212.05	1,818.0		58.49
7	193	105.41	826.0		60.08
8	201	68.95	518.5		60.85
9	258	446.60	2,620.4		66.12
10	283	193.85	1,037.2		68.44
11	334	426.75	1,935.1		72.71
12	338	94.13	421.7		73.03
13	355	140.97	601.1		74.39
14	370	97.64	400.2		75.52
15	385	152.98	602.8		76.71
16	393	83.23	320.8		77.39
17	430	126.79	447.2		79.61
18	464	79.66	260.1		81.57
19	473	80.47	257.9		82.07
20	492	138.61	426.9		83.16
21	506	132.67	397.6		83.88
22	550	198.77	547.7		86.02
23	581	130.39	340.1		87.52
24	621	62.56	152.7		89.06
25	649	64.68	150.9		89.92

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad


Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary Continued ...

... Peak table for sample 1 : KL06 soni 40 sec

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	670	65.31	147.6		90.56
27	727	80.25	167.2		91.83
28	816	61.20	113.6		93.06
29	846	95.16	170.5		93.47
30	921	117.92	193.9		94.51
31	1,060	111.88	159.9		95.97
32	1,493	86.34	87.6		98.60
33	1,694	113.48	101.5		99.83
34	2,060	59.37	43.7		101.87
35	2,920	174.80	90.7		104.51
36	5,672	52.15	13.9		108.00
37	9,017	21.77	3.7		111.64
38	 10,380	75.00	10.9	Upper Marker	113.00
39	13,514	0.00	0.0		116.13
40	15,421	0.00	0.0		118.04
41	16,103	0.00	0.0		118.72
42	18,510	0.00	0.0		121.13
43	19,146	0.00	0.0		121.77
44	20,781	0.00	0.0		123.40
45	23,006	0.00	0.0		125.63

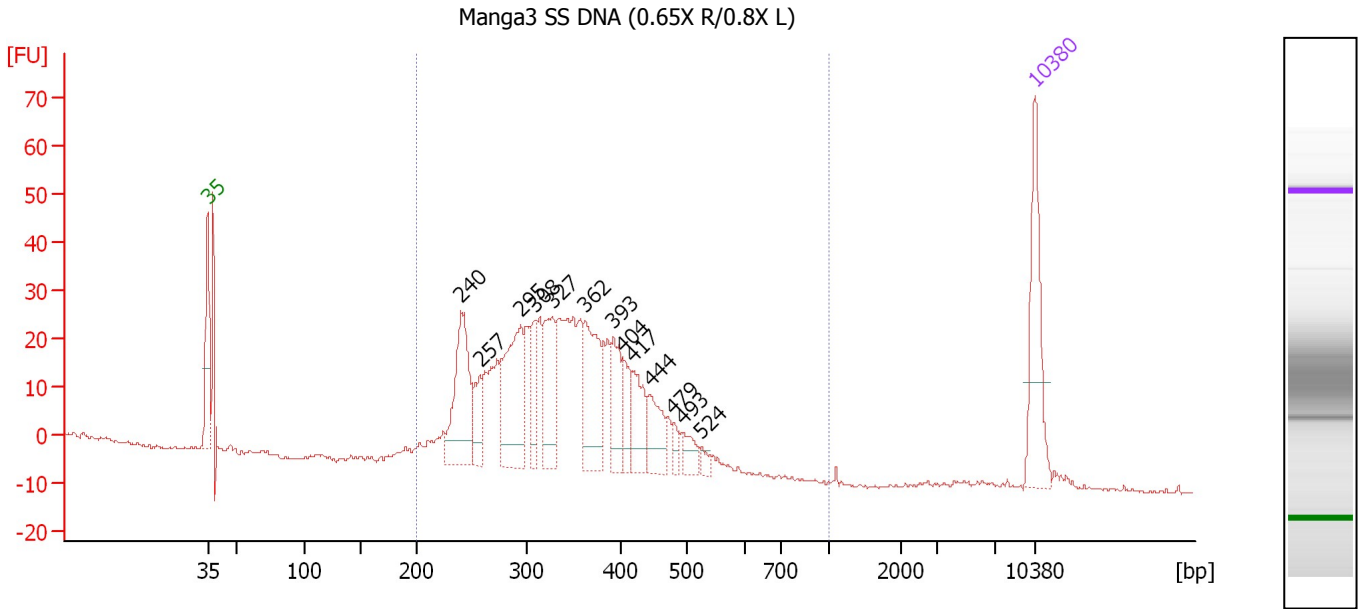
Region table for sample 1 : KL06 soni 40 sec

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
45	21,170	1,864	10,841.7	27,369.6	 5,772.14	99	100.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : Manga3 SS DNA (0.65X R/0.8X L)

Number of peaks found: 13 Corr. Area 1: 625.5
 Noise: 0.4

Peak table for sample 2 : Manga3 SS DNA (0.65X R/0.8X L)

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	240	135.91	857.7		64.44
3	257	36.02	212.3		66.02
4	295	136.06	699.9		69.49
5	308	44.60	219.1		70.67
6	327	80.75	374.0		72.16
7	362	100.46	419.9		74.95
8	393	50.86	196.3		77.35
9	404	34.53	129.5		78.16
10	417	48.22	175.0		78.93
11	444	44.40	151.6		80.41
12	479	12.00	38.0		82.40
13	493	21.46	66.0		83.21
14	524	7.82	22.6		84.79
15	10,380	75.00	10.9	Upper Marker	113.00

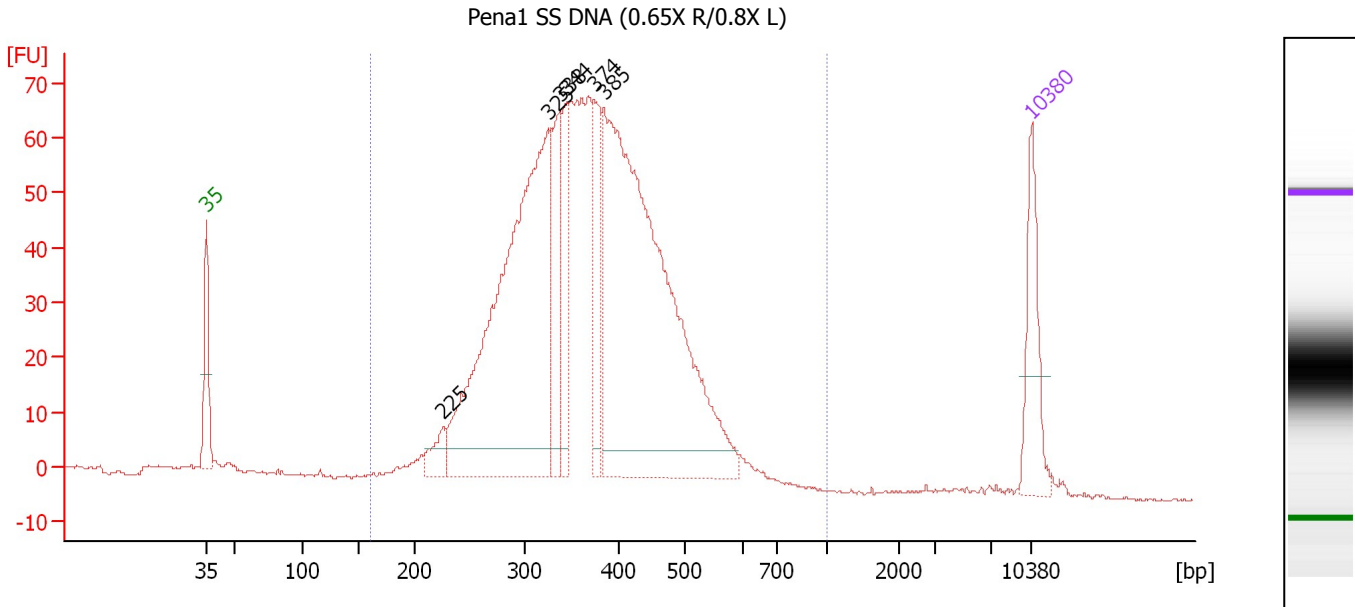
Region table for sample 2 : Manga3 SS DNA (0.65X R/0.8X L)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. Ior [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	342	625.5	4,948.1	1,053.01	95	20.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : Pena1 SS DNA (0.65X R/0.8X L)

Number of peaks found: 6 Corr. Area 1: 1,471.2
 Noise: 0.3

Peak table for sample 3 : Pena1 SS DNA (0.65X R/0.8X L)

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	225	37.02	248.8		63.08
3	325	765.46	3,568.1		71.99
4	338	126.87	569.4		72.98
5	344	93.57	411.6		73.52
6	374	101.70	411.9		75.87
7	385	868.80	3,420.1		76.73
8	10,380	75.00	10.9	Upper Marker	113.00

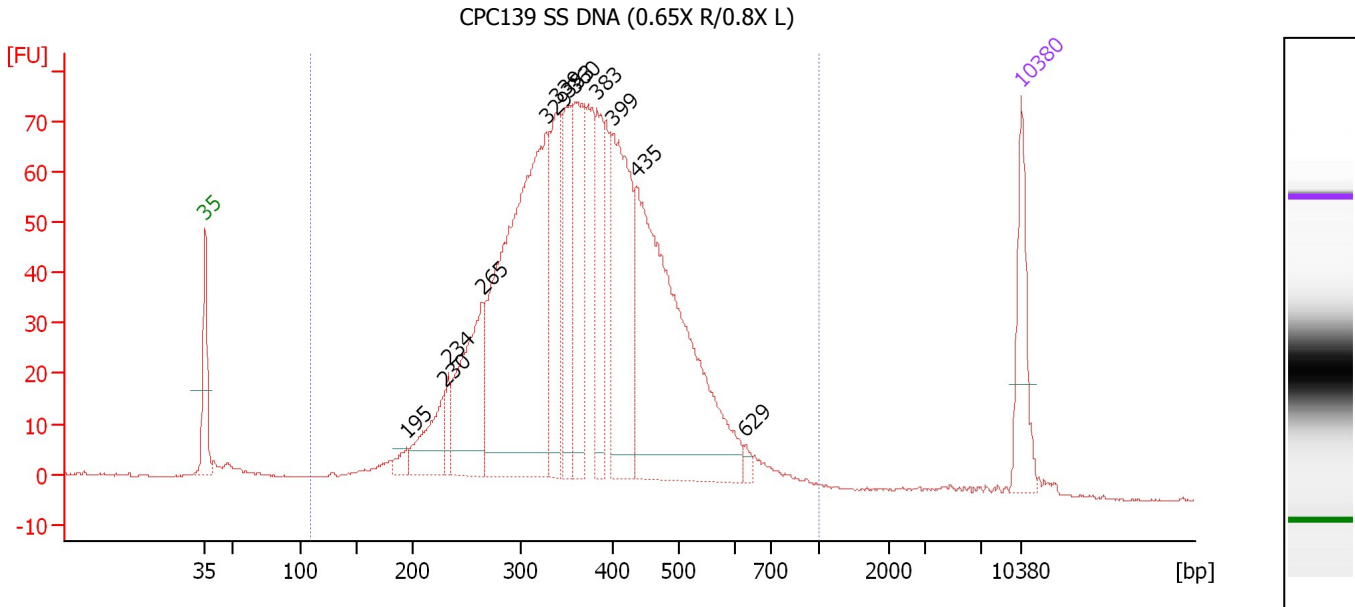
Region table for sample 3 : Pena1 SS DNA (0.65X R/0.8X L)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
160	1,000	374	1,471.2	10,764.1	2,471.53	98	22.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : CPC139 SS DNA (0.65X R/0.8X L)

Number of peaks found: 12 Corr. Area 1: 1,764.9
 Noise: 0.2

Peak table for sample 4 : CPC139 SS DNA (0.65X R/0.8X L)

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	195	19.57	151.7		60.28
3	230	93.29	615.0		63.48
4	234	33.45	216.8		63.85
5	265	211.75	1,211.7		66.73
6	329	732.25	3,371.2		72.31
7	339	173.98	776.5		73.13
8	353	149.84	642.6		74.23
9	360	196.74	828.8		74.73
10	383	122.99	486.2		76.61
11	399	279.96	1,062.0		77.89
12	435	578.97	2,014.8		79.94
13	629	8.42	20.3		89.32
14	10,380	75.00	10.9	Upper Marker	113.00

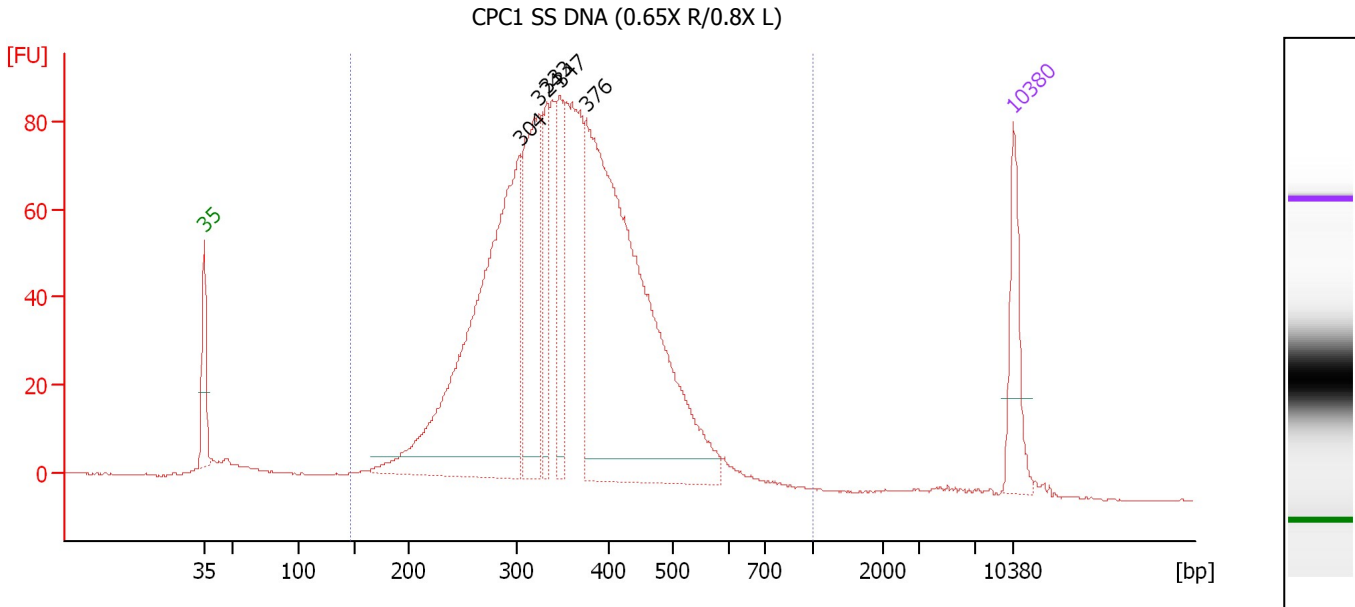
Region table for sample 4 : CPC139 SS DNA (0.65X R/0.8X L)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
109	1,000	377	1,764.9	13,553.2	3,036.83	98	27.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : CPC1 SS DNA (0.65X R/0.8X L)

Number of peaks found: 5 Corr. Area 1: 1,858.1
 Noise: 0.2

Peak table for sample 5 : CPC1 SS DNA (0.65X R/0.8X L)

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	802.76	3,996.3		70.35
3	324	292.86	1,369.0		71.91
4	333	110.27	501.1		72.65
5	347	125.83	548.9		73.76
6	376	954.96	3,849.6		76.02
7	10,380	75.00	10.9	Upper Marker	113.00

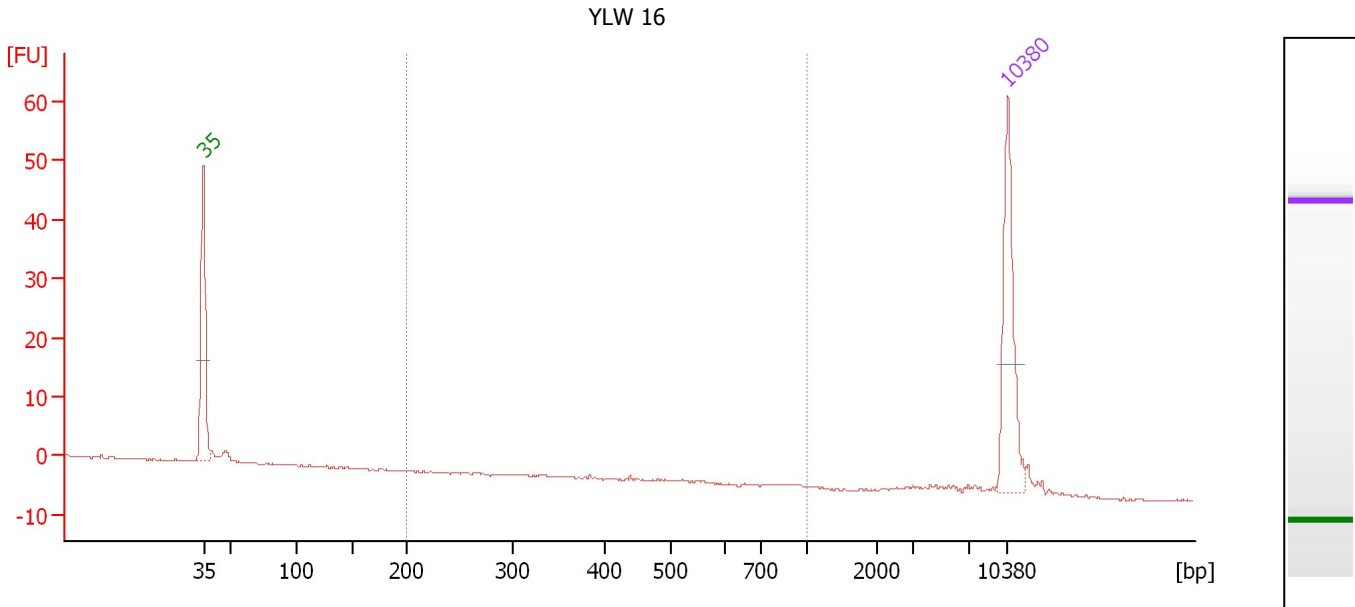
Region table for sample 5 : CPC1 SS DNA (0.65X R/0.8X L)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
147	1,000	360	1,858.1	13,099.8	2,868.62	97	24.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : YLW 16

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

Peak table for sample 6 : YLW 16

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

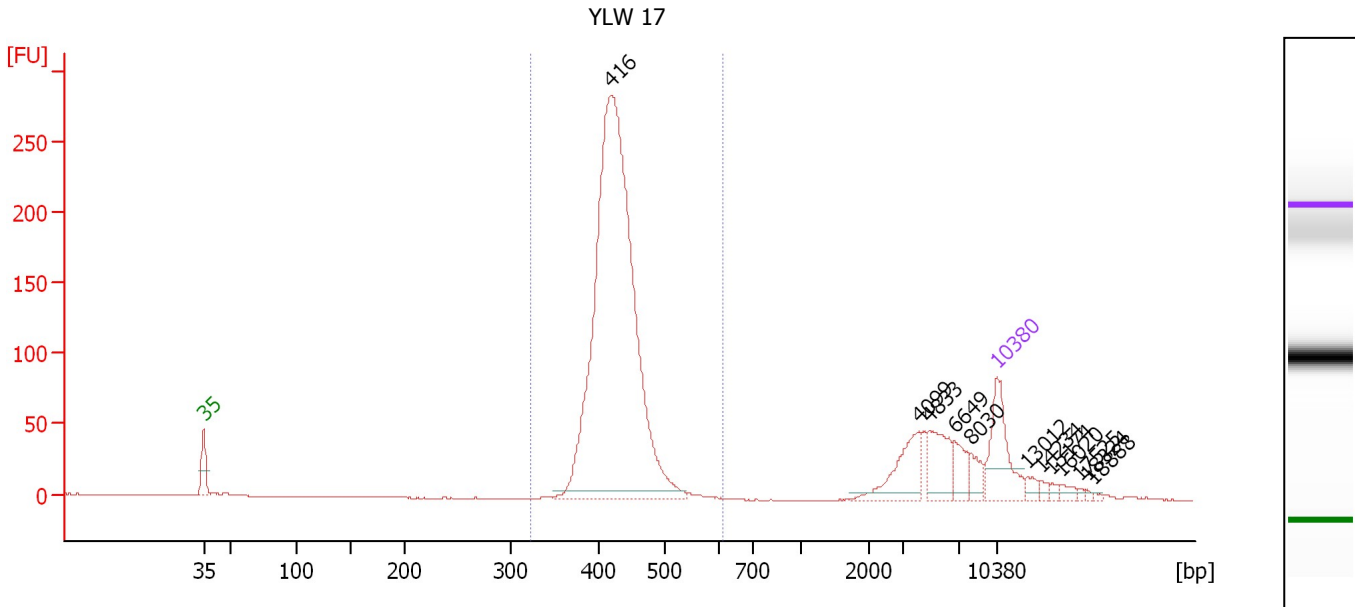
Region table for sample 6 : YLW 16

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	420	0.2	1.5	0.40	1	14.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : YLW 17

Number of peaks found: 12 Corr. Area 1: 1,440.5
 Noise: 0.2

Peak table for sample 7 : YLW 17

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	416	1,153.97	4,206.3		78.82
3	4,099	68.63	25.4		106.09
4	4,833	59.41	18.6		106.98
5	6,649	30.42	6.9		109.19
6	8,030	20.14	3.8		110.65
7	10,380	75.00	10.9	Upper Marker	113.00
8	13,012	0.00	0.0		115.63
9	14,234	0.00	0.0		116.85
10	15,174	0.00	0.0		117.80
11	16,020	0.00	0.0		118.64
12	17,525	0.00	0.0		120.15
13	18,324	0.00	0.0		120.94
14	18,888	0.00	0.0		121.51

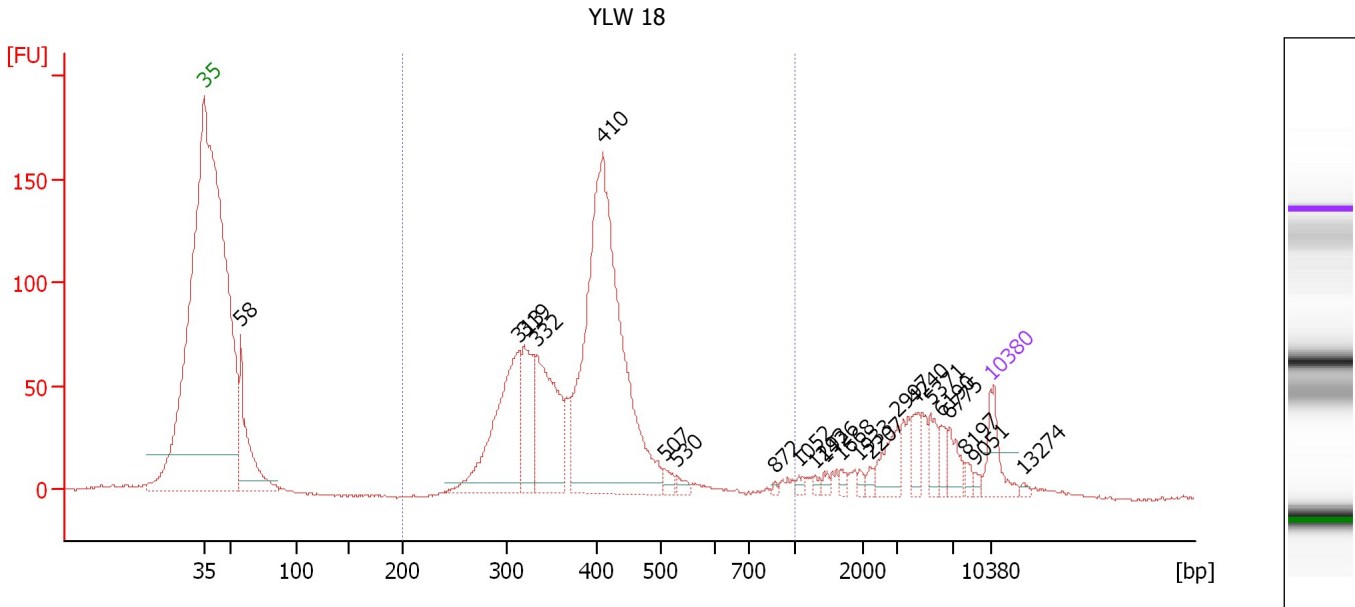
Region table for sample 7 : YLW 17

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
324	614	426	1,440.5	4,082.2	1,143.20	79	7.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : YLW 18

Number of peaks found: 22 Corr. Area 1: 1,430.6
 Noise: 0.6


Peak table for sample 8 : YLW 18

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	58	359.25	9,437.7		46.27
3	313	511.77	2,480.4		71.00
4	319	215.34	1,022.2		71.52
5	332	396.99	1,809.8		72.57
6	410	1,369.58	5,061.7		78.50
7	507	25.51	76.2		83.96
8	530	18.99	54.3		85.05
9	872	5.82	10.1		93.83
10	1,052	11.83	17.0		95.92
11	1,293	9.78	11.5		97.39
12	1,426	11.37	12.1		98.19
13	1,668	10.92	9.9		99.66
14	1,933	10.06	7.9		101.28
15	2,207	12.30	8.4		102.32
16	2,997	72.20	36.5		104.74
17	4,240	44.96	16.1		106.26
18	5,371	43.37	12.2		107.64
19	6,190	28.08	6.9		108.63
20	6,775	47.91	10.7		109.35
21	8,197	14.32	2.6		110.82
22	9,051	9.35	1.6		111.67
23	10,380	75.00	10.9	Upper Marker	113.00
24	13,274	0.00	0.0		115.89

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
Modified: 12/10/2015 5:14:14 PM

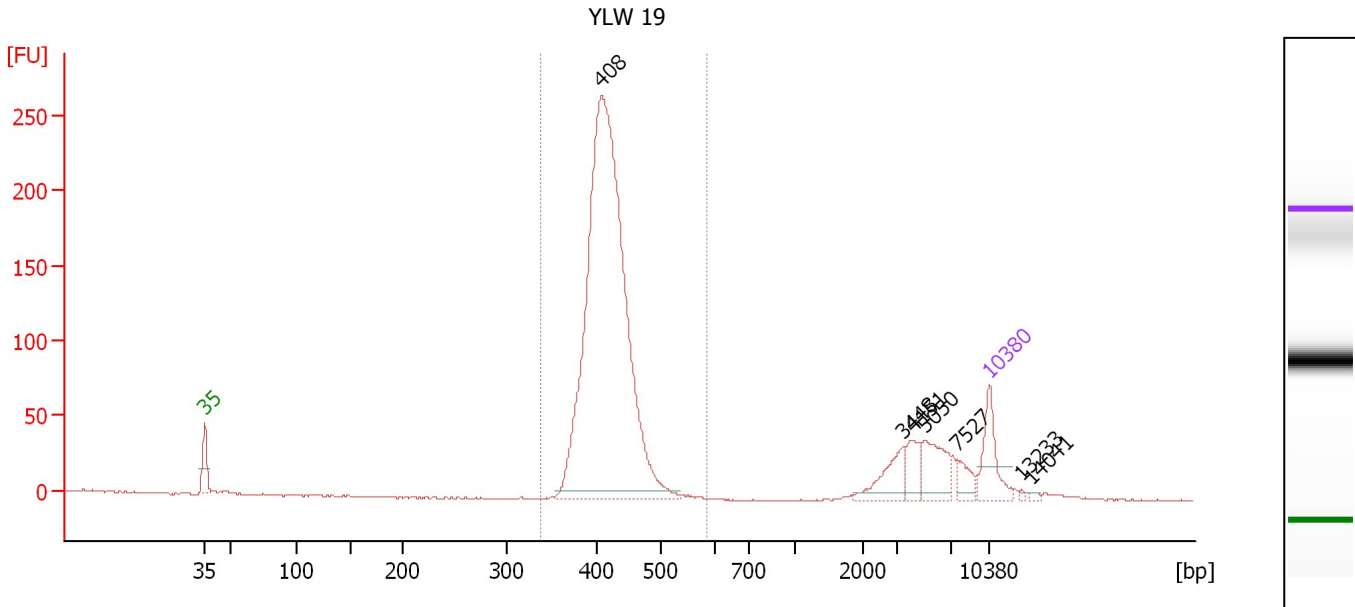
Electropherogram Summary Continued ...**... Region table for sample 8 : YLW 18**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	391	1,430.6	10,841.5	 2,659.63	74	22.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : YLW 19

Number of peaks found: 7 Corr. Area 1: 1,315.7
 Noise: 0.5

Peak table for sample 9 : YLW 19

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	408	1,679.41	6,238.3		78.38
3	3,448	68.49	30.1		105.30
4	4,151	44.55	16.3		106.15
5	5,050	82.01	24.6		107.25
6	7,527	30.76	6.2		110.15
7	10,380	75.00	10.9	Upper Marker	113.00
8	13,233	0.00	0.0		115.85
9	14,041	0.00	0.0		116.66

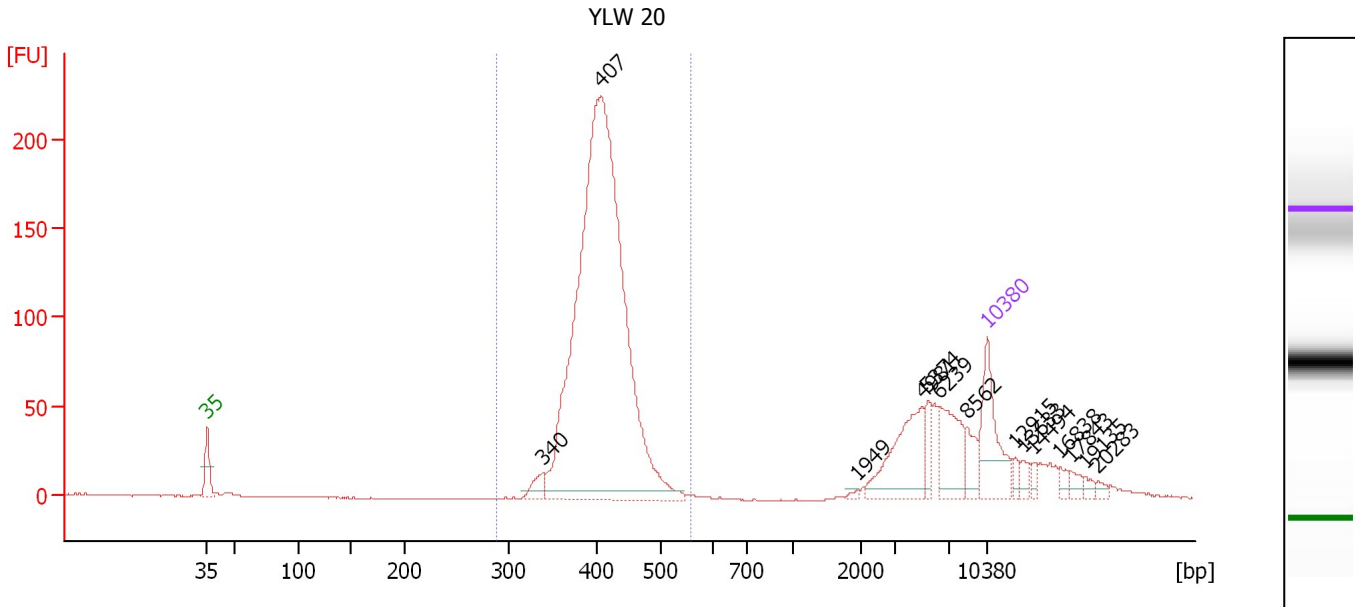
Region table for sample 9 : YLW 19

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
337	585	418	1,315.7	5,917.7	1,628.22	83	6.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : YLW 20

Number of peaks found: 14 Corr. Area 1: 1,375.0
 Noise: 0.3

Peak table for sample 10 : YLW 20

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	340	24.62	109.8		73.14
3	407	1,220.00	4,545.4		78.31
4	1,949	2.47	1.9		101.37
5	4,981	99.70	30.3		107.16
6	5,374	22.02	6.2		107.64
7	6,239	67.30	16.3		108.69
8	8,562	25.18	4.5		111.18
9	10,380	75.00	10.9	Upper Marker	113.00
10	12,915	0.00	0.0		115.54
11	13,633	0.00	0.0		116.25
12	14,494	0.00	0.0		117.11
13	16,838	0.00	0.0		119.46
14	17,843	0.00	0.0		120.46
15	19,135	0.00	0.0		121.76
16	20,283	0.00	0.0		122.90

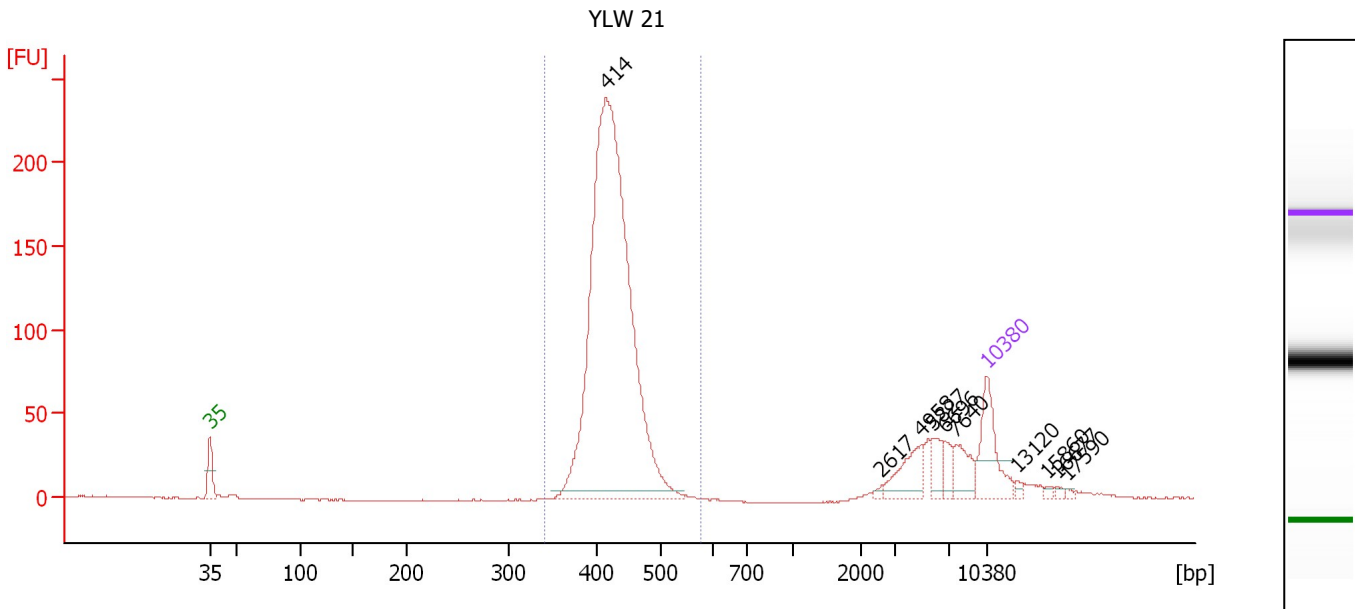
Region table for sample 10 : YLW 20

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
289	558	410	1,375.0	4,527.3	1,216.13	74	8.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : YLW 21

Number of peaks found: 10 Corr. Area 1: 1,208.9
 Noise: 0.3

Peak table for sample 11 : YLW 21

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	414	1,286.37	4,709.1		78.72
3	2,617	4.44	2.6		103.58
4	4,958	54.89	16.8		107.13
5	5,827	26.79	7.0		108.19
6	6,696	23.79	5.4		109.25
7	7,640	38.39	7.6		110.26
8	10,380	75.00	10.9	Upper Marker	113.00
9	13,120	0.00	0.0		115.74
10	15,860	0.00	0.0		118.48
11	16,677	0.00	0.0		119.30
12	17,590	0.00	0.0		120.21

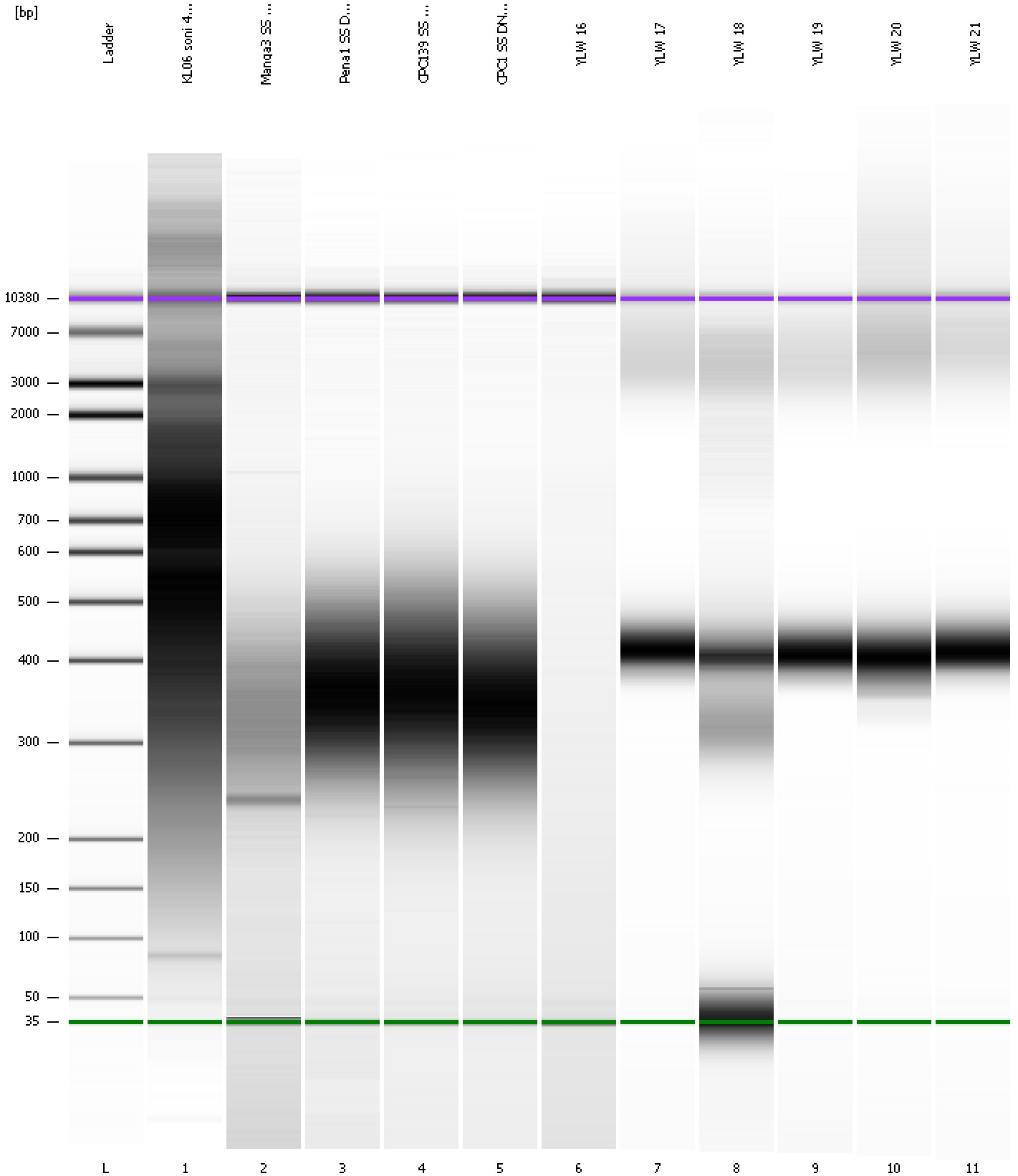
Region table for sample 11 : YLW 21

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
340	574	424	1,208.9	4,519.8	1,261.50	83	6.8

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
Modified: 12/10/2015 5:14:14 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad

Created: 12/10/2015 4:31:58 PM
 Modified: 12/10/2015 5:14:14 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		12/10/2015 5:13:48 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-12-10\2015-12-10_004.xad)		Instrument	Run		12/10/2015 4:32:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		12/10/2015 4:32:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		12/10/2015 4:32:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		12/10/2015 4:32:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		12/10/2015 4:32:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		12/10/2015 4:32:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		12/10/2015 4:32:03 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1