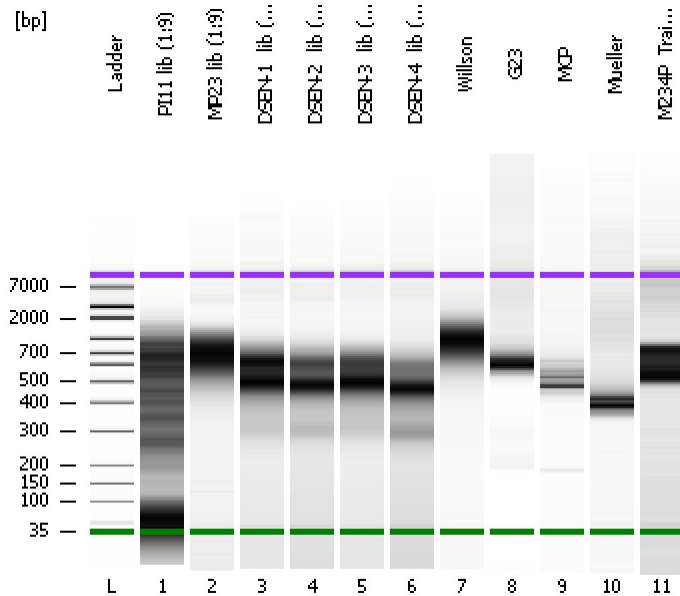


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
Modified: 12/7/2015 11:13:14 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

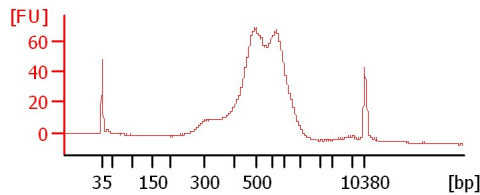
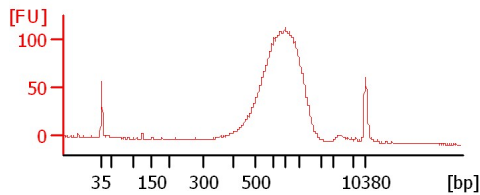
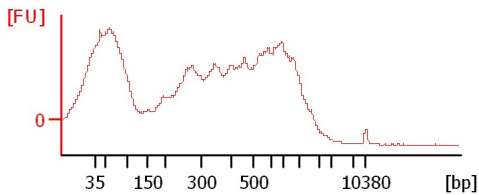
Chip Information:

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

PI11 lib (1:9)

MP23 lib (1:9)

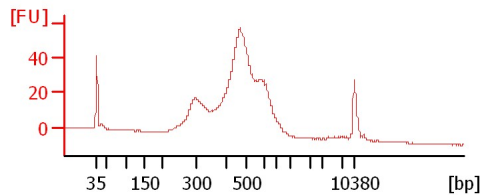
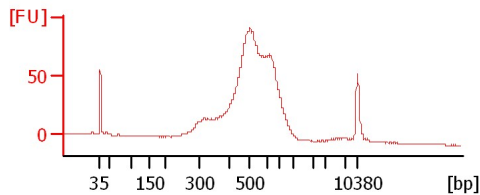
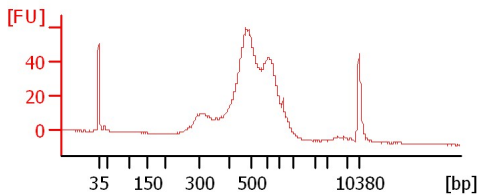
DSEN-1 lib (1:2)



DSEN-2 lib (1:2)

DSEN-3 lib (1:4)

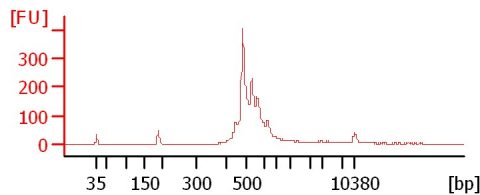
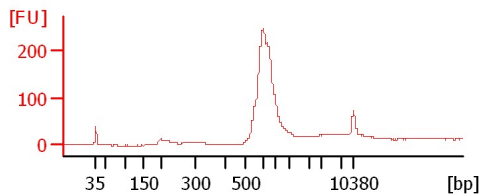
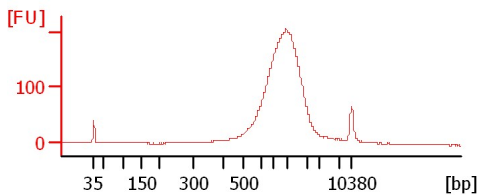
DSEN-4 lib (1:2)



Willson

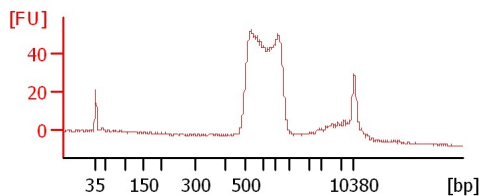
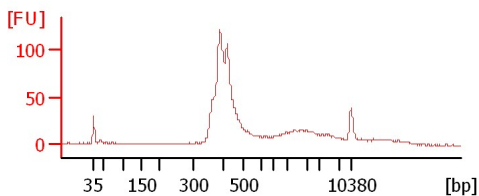
G23

MCP



Mueller

M234P_Trainor BP



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
Modified: 12/7/2015 11:13:14 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
PI11 lib (1:9)		<input type="checkbox"/>	✓			
MP23 lib (1:9)		<input type="checkbox"/>	✓			
DSEN-1 lib (1:2)		<input type="checkbox"/>	✓			
DSEN-2 lib (1:2)		<input type="checkbox"/>	✓			
DSEN-3 lib (1:4)		<input type="checkbox"/>	✓			
DSEN-4 lib (1:2)		<input type="checkbox"/>	✓			
Willson		<input type="checkbox"/>	✓			
G23		<input type="checkbox"/>	✓			
MCP		<input type="checkbox"/>	✓			
Mueller		<input type="checkbox"/>	✓			
M234P_Trainor BP		<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\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
Modified: 12/7/2015 11:13:14 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

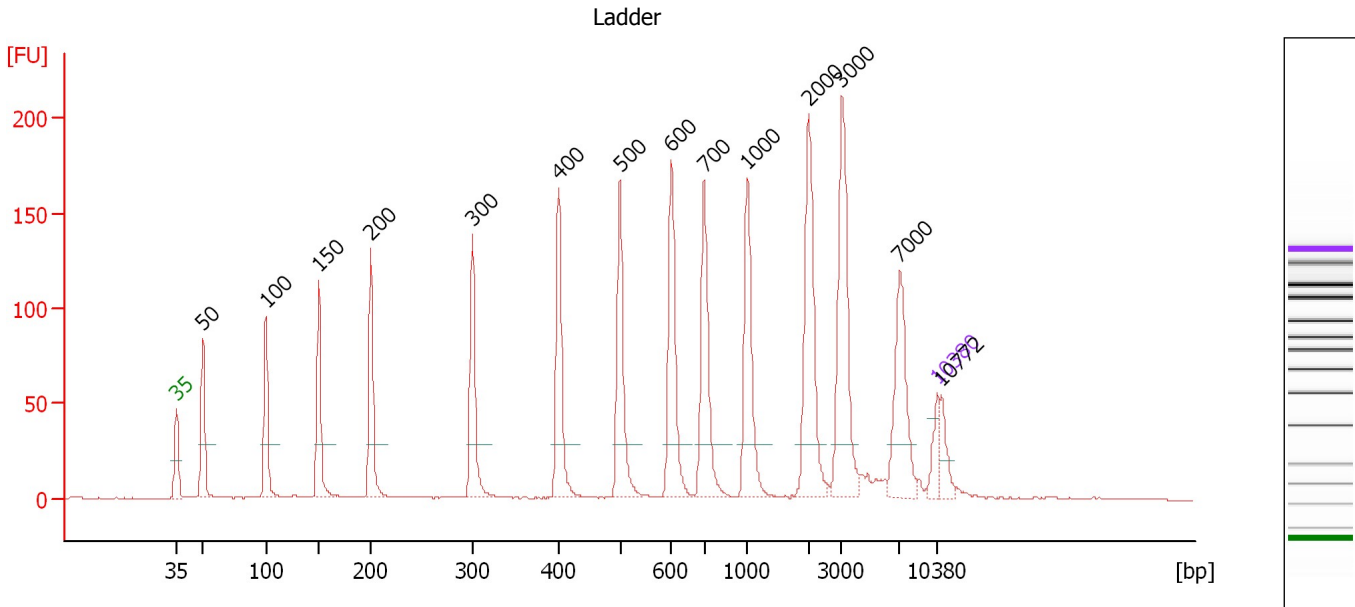
Ladder

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\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
 Modified: 12/7/2015 11:13:14 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

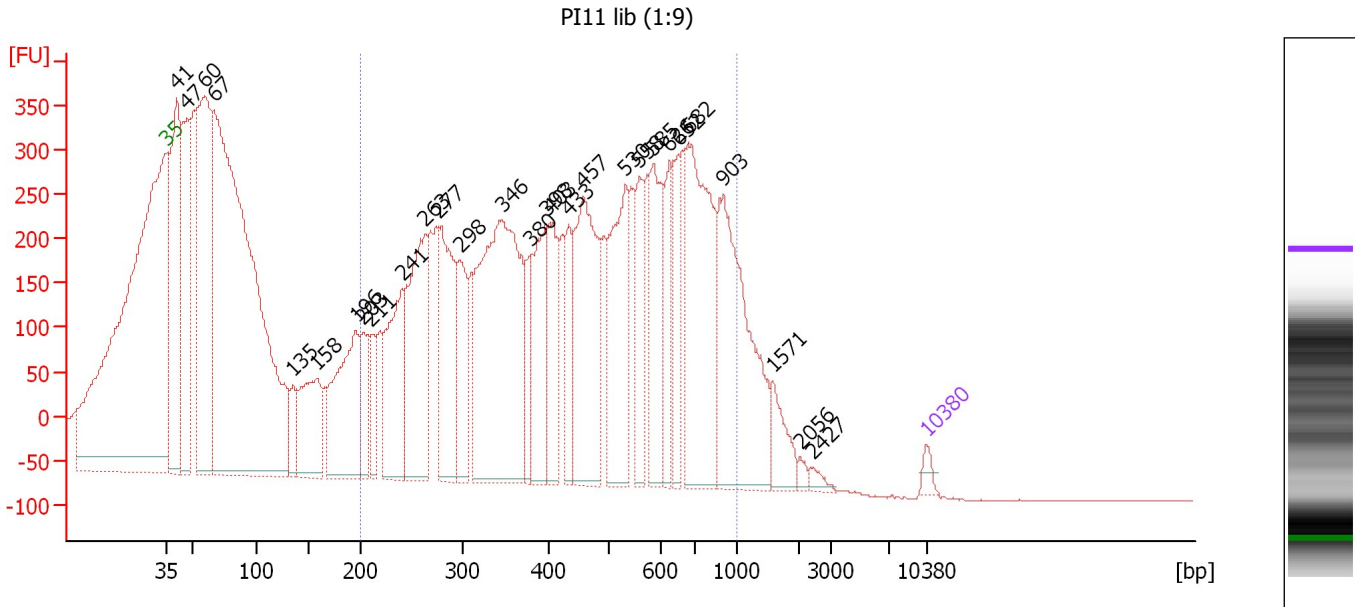
Peak table for Ladder

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	45.45
3	100	150.00	2,272.7	Ladder Peak	51.25
4	150	150.00	1,515.2	Ladder Peak	56.11
5	200	150.00	1,136.4	Ladder Peak	60.92
6	300	150.00	757.6	Ladder Peak	70.24
7	400	150.00	568.2	Ladder Peak	78.20
8	500	150.00	454.5	Ladder Peak	83.84
9	600	150.00	378.8	Ladder Peak	88.55
10	700	150.00	324.7	Ladder Peak	91.60
11	1,000	150.00	227.3	Ladder Peak	95.57
12	2,000	150.00	113.6	Ladder Peak	101.22
13	3,000	150.00	75.8	Ladder Peak	104.26
14	7,000	150.00	32.5	Ladder Peak	109.61
15	10,380	75.00	10.9	Upper Marker	113.00
16	10,772	0.00	0.0		113.39

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
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Electropherogram Summary Continued ...



Overall Results for sample 1 : PI11 lib (1:9)

Number of peaks found: 29 Corr. Area 1: 10,926.5
 Noise: 0.4

Peak table for sample 1 : PI11 lib (1:9)

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	41	3,613.35	133,571.9		43.98
3	47	2,315.72	74,692.4		44.96
4	60	4,335.83	110,076.0		46.58
5	67	12,302.64	278,758.7		47.41
6	135	366.36	4,108.7		54.66
7	158	1,348.70	12,944.9		56.86
8	196	1,912.53	14,819.6		60.49
9	203	547.99	4,095.2		61.17
10	211	464.11	3,338.6		61.91
11	241	1,561.23	9,833.2		64.70
12	263	2,144.80	12,349.4		66.81
13	277	1,707.86	9,330.7		68.13
14	298	931.80	4,732.3		70.09
15	346	4,108.32	17,985.5		73.91
16	380	478.70	1,908.8		76.60
17	398	1,107.55	4,218.0		78.02
18	403	873.91	3,285.4		78.37
19	433	577.78	2,019.9		80.08
20	457	2,051.39	6,803.8		81.40
21	530	1,528.26	4,365.6		85.27
22	558	685.71	1,860.4		86.60
23	585	1,106.17	2,862.6		87.87
24	626	548.69	1,328.4		89.34
25	652	617.76	1,436.5		90.12

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
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Electropherogram Summary Continued ...

... Peak table for sample 1 : PI11 lib (1:9)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	682	2,306.12	5,122.1		91.05
27	903	2,236.77	3,753.2		94.29
28	1,571	350.76	338.4		98.79
29	2,056	52.96	39.0		101.39
30	2,427	66.17	41.3		102.52
31	10,380	75.00	10.9	Upper Marker	113.00

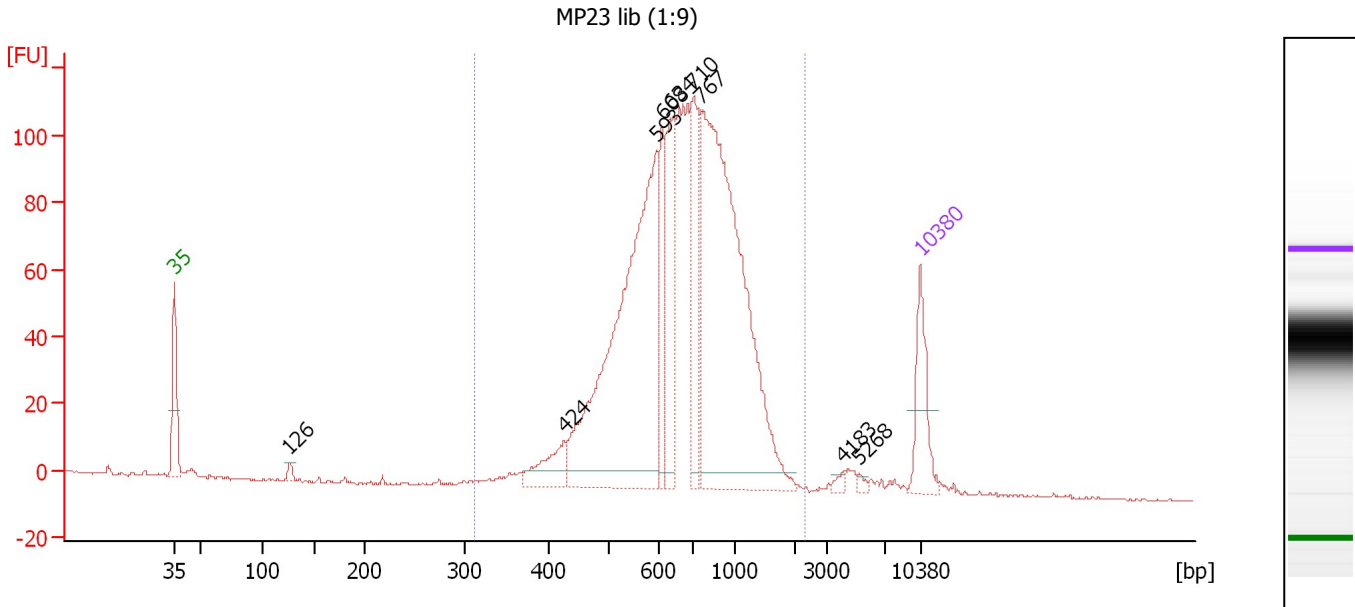
Region table for sample 1 : PI11 lib (1:9)

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	487	10,926.5	94,441.8	24,052.95	59	39.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
 Modified: 12/7/2015 11:13:14 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 : MP23 lib (1:9)

Number of peaks found: 9 Corr. Area 1: 1,575.8
 Noise: 0.3

Peak table for sample 2 : MP23 lib (1:9)

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	126	10.74	128.8		53.81
3	424	68.59	244.9		79.58
4	593	711.39	1,817.1		88.23
5	608	111.29	277.5		88.78
6	634	170.49	407.5		89.58
7	710	161.41	344.3		91.73
8	767	799.14	1,578.7		92.49
9	4,183	6.34	2.3		105.84
10	5,268	4.82	1.4		107.30
11	10,380	75.00	10.9	Upper Marker	113.00

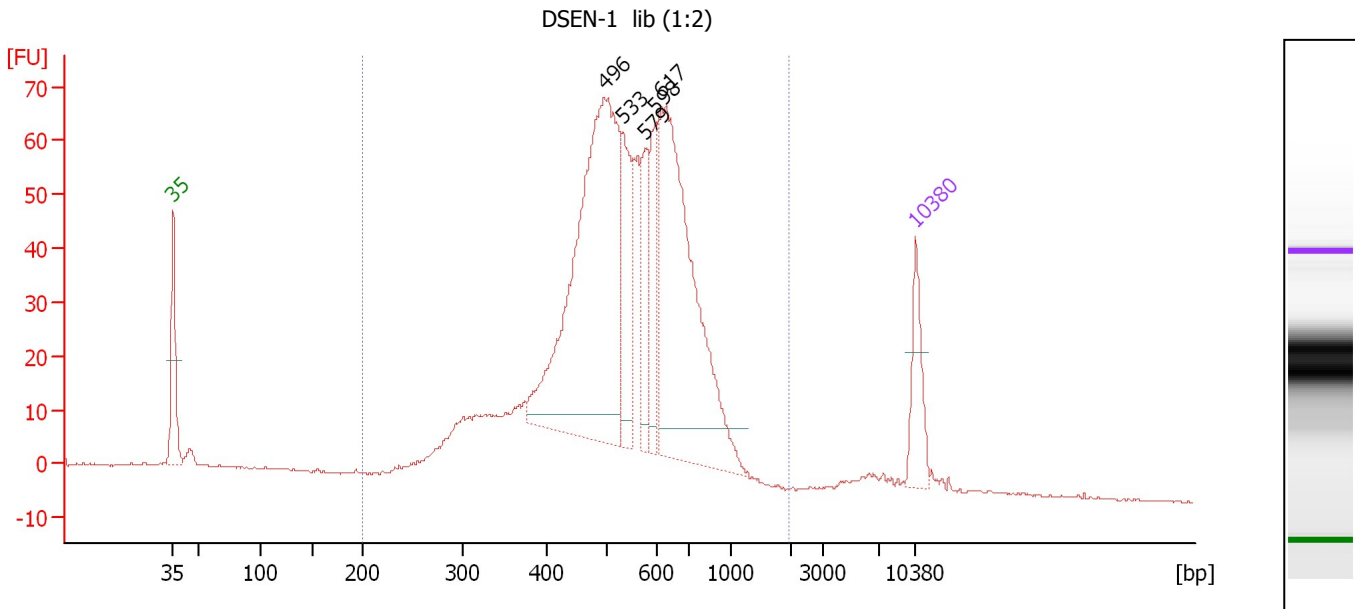
Region table for sample 2 : MP23 lib (1:9)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
311	2,309	747	1,575.8	5,264.2	2,295.58	97	35.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
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Electropherogram Summary Continued ...



Overall Results for sample 3 : DSEN-1 lib (1:2)

Number of peaks found: 5 Corr. Area 1: 1,189.2
 Noise: 0.2

Peak table for sample 3 : DSEN-1 lib (1:2)

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	496	909.37	2,776.3		83.63
3	533	176.40	501.5		85.39
4	579	104.24	272.9		87.55
5	598	131.76	333.8		88.46
6	617	661.17	1,624.3		89.06
7	10,380	75.00	10.9	Upper Marker	113.00

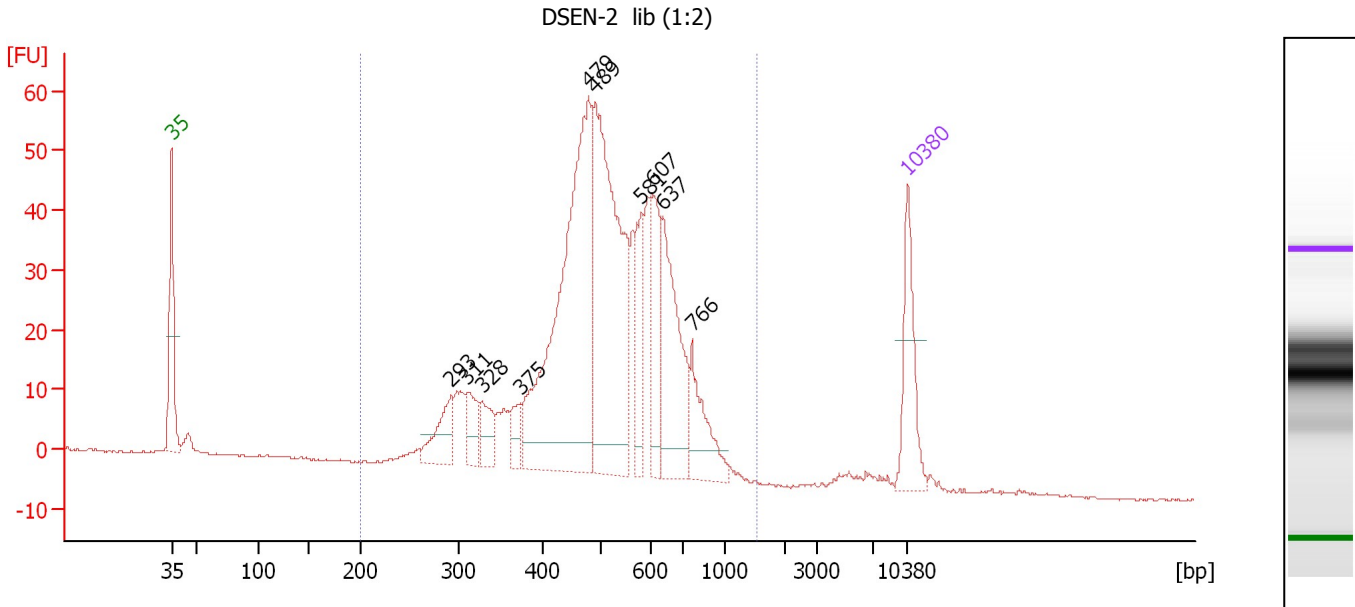
Region table for sample 3 : DSEN-1 lib (1:2)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,977	550	1,189.2	8,903.9	2,902.47	96	28.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
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Electropherogram Summary Continued ...



Overall Results for sample 4 : DSEN-2 lib (1:2)

Number of peaks found: 10 Corr. Area 1: 893.1
 Noise: 0.2

Peak table for sample 4 : DSEN-2 lib (1:2)

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	293	67.45	349.2		69.56
3	311	43.70	212.8		71.13
4	328	44.28	204.3		72.50
5	375	29.84	120.6		76.20
6	479	539.07	1,706.3		82.64
7	489	430.86	1,333.8		83.25
8	581	80.13	209.0		87.66
9	607	89.22	222.7		88.77
10	637	174.59	415.2		89.68
11	766	77.70	153.7		92.47
12	10,380	75.00	10.9	Upper Marker	113.00

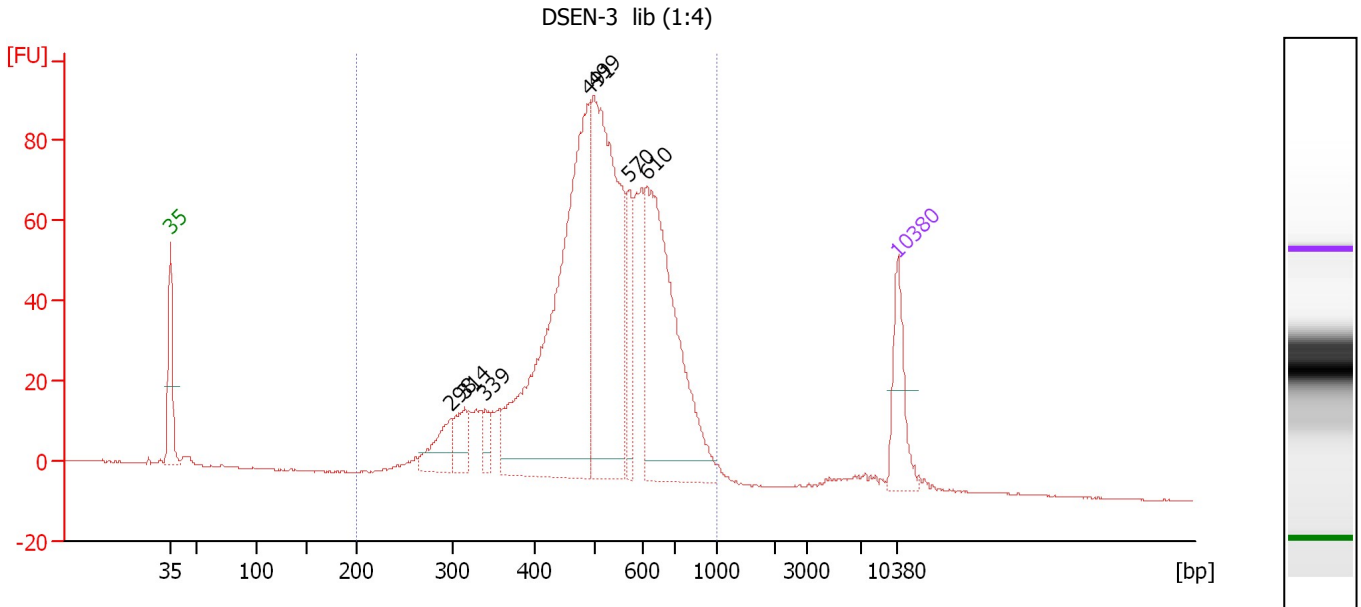
Region table for sample 4 : DSEN-2 lib (1:2)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,544	517	893.1	5,877.6	1,819.30	97	25.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 5 : DSEN-3 lib (1:4)

Number of peaks found: 7 Corr. Area 1: 1,400.2
 Noise: 0.2

Peak table for sample 5 : DSEN-3 lib (1:4)

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	298	71.52	363.9		70.03
3	314	52.45	253.0		71.37
4	339	25.10	112.3		73.32
5	491	737.95	2,276.2		83.35
6	499	548.21	1,663.2		83.81
7	570	90.87	241.5		87.15
8	610	441.07	1,096.4		88.84
9	10,380	75.00	10.9	Upper Marker	113.00

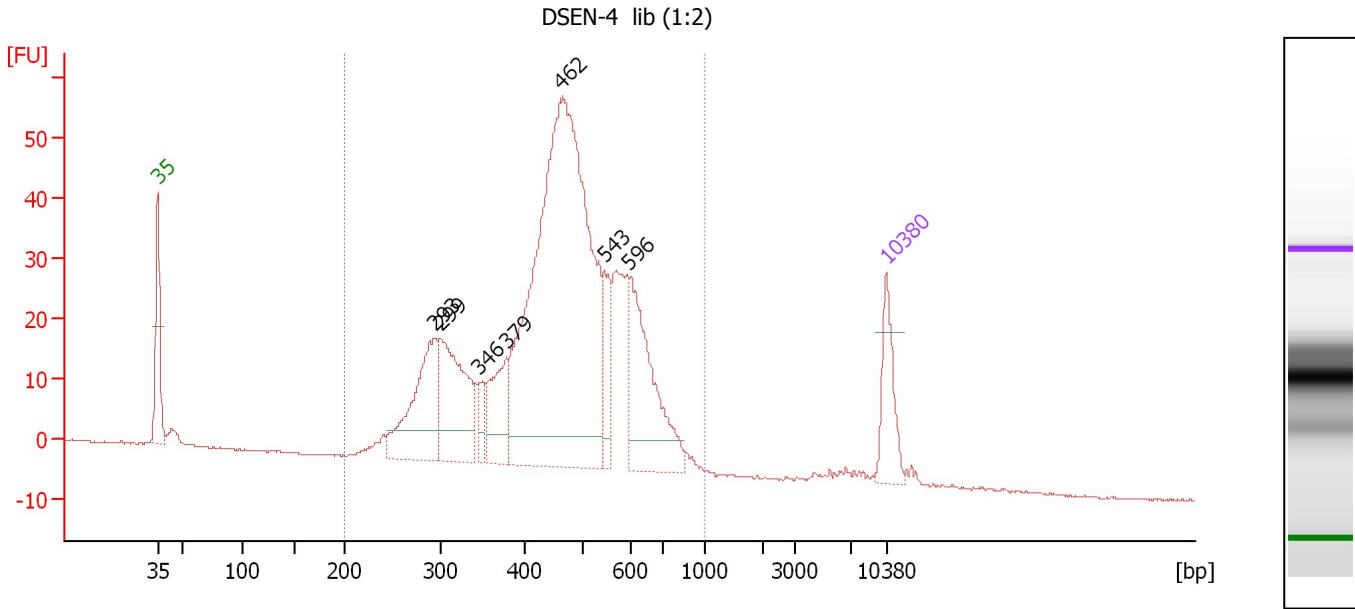
Region table for sample 5 : DSEN-3 lib (1:4)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	523	1,400.2	7,155.5	2,265.70	98	23.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
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Electropherogram Summary Continued ...



Overall Results for sample 6 : DSEN-4 lib (1:2)

Number of peaks found: 7 Corr. Area 1: 870.7
 Noise: 0.2

Peak table for sample 6 : DSEN-4 lib (1:2)

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	293	244.17	1,264.3		69.56
3	299	259.26	1,312.5		70.18
4	346	35.53	155.5		73.92
5	379	118.45	473.7		76.52
6	462	1,239.06	4,060.5		81.72
7	543	86.65	241.7		85.87
8	596	241.53	614.0		88.37
9	10,380	75.00	10.9	Upper Marker	113.00

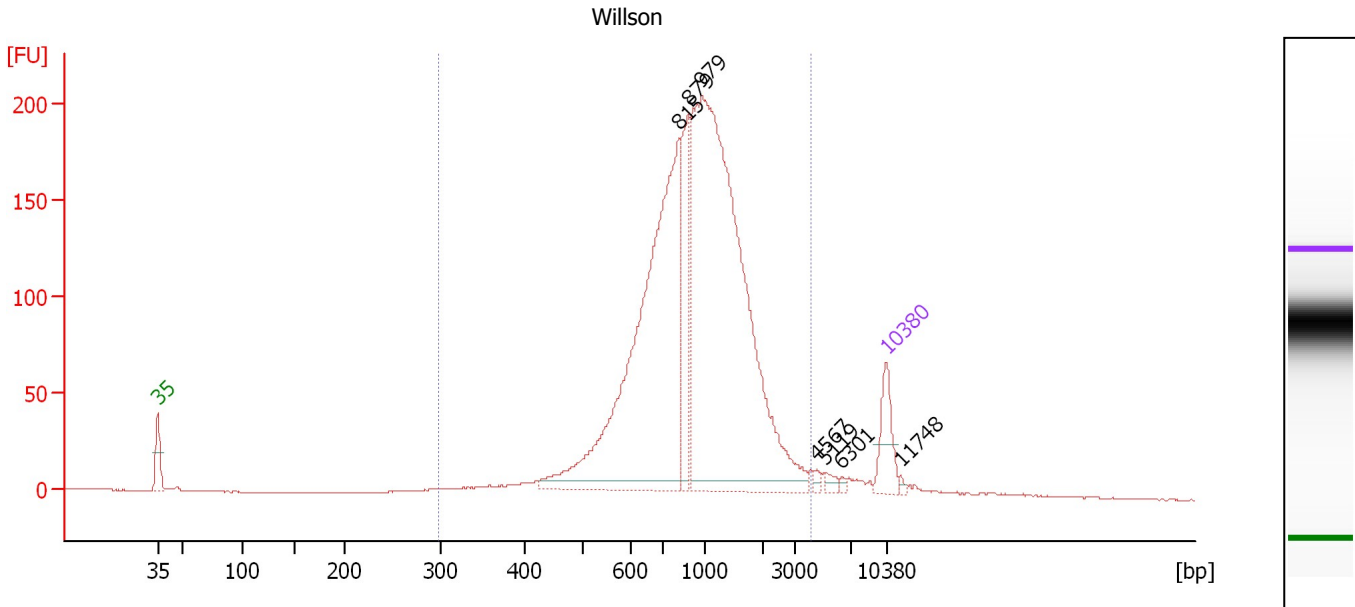
Region table for sample 6 : DSEN-4 lib (1:2)

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	466	870.7	8,721.2	2,427.57	98	25.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
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Electropherogram Summary Continued ...



Overall Results for sample 7 : Willson

Number of peaks found: 7 Corr. Area 1: 2,382.5
 Noise: 0.1

Peak table for sample 7 : Willson

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	815	1,148.58	2,134.0		93.13
3	879	200.40	345.4		93.97
4	979	1,481.01	2,293.2		95.29
5	4,567	8.41	2.8		106.36
6	5,119	12.83	3.8		107.10
7	6,301	5.37	1.3		108.68
8	10,380	75.00	10.9	Upper Marker	113.00
9	11,748	0.00	0.0		114.37

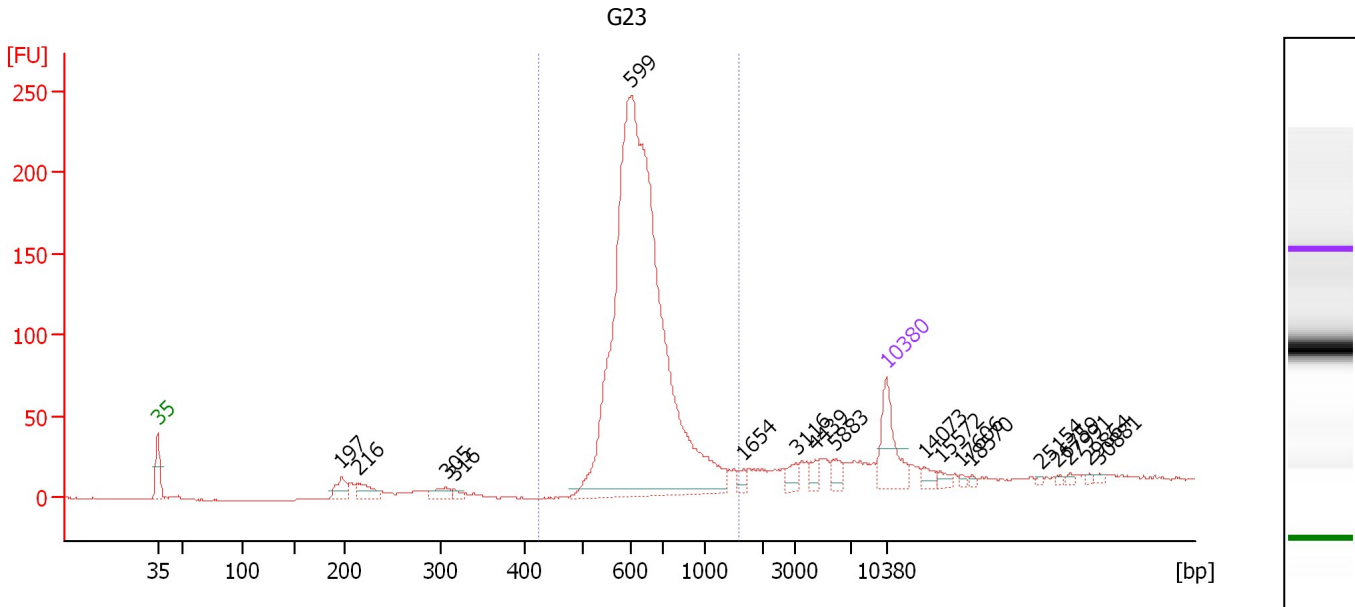
Region table for sample 7 : Willson

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
298	4,061	1,068	2,382.5	5,337.8	2,996.95	96	49.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 8 : G23

Number of peaks found: 18 Corr. Area 1: 1,348.3
 Noise: 0.3

Peak table for sample 8 : G23

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	197	34.98	268.8		60.65
3	216	34.35	241.5		62.36
4	305	22.98	114.0		70.68
5	316	9.03	43.3		71.54
6	599	1,478.81	3,743.3		88.49
7	1,654	9.87	9.0		99.27
8	3,116	16.02	7.8		104.42
9	4,439	12.68	4.3		106.19
10	5,883	18.01	4.6		108.12
11	10,380	75.00	10.9	Upper Marker	113.00
12	14,073	0.00	0.0		116.70
13	15,572	0.00	0.0		118.20
14	17,606	0.00	0.0		120.24
15	18,570	0.00	0.0		121.21
16	25,154	0.00	0.0		127.80
17	26,759	0.00	0.0		129.41
18	27,991	0.00	0.0		130.65
19	29,864	0.00	0.0		132.52
20	30,881	0.00	0.0		133.54

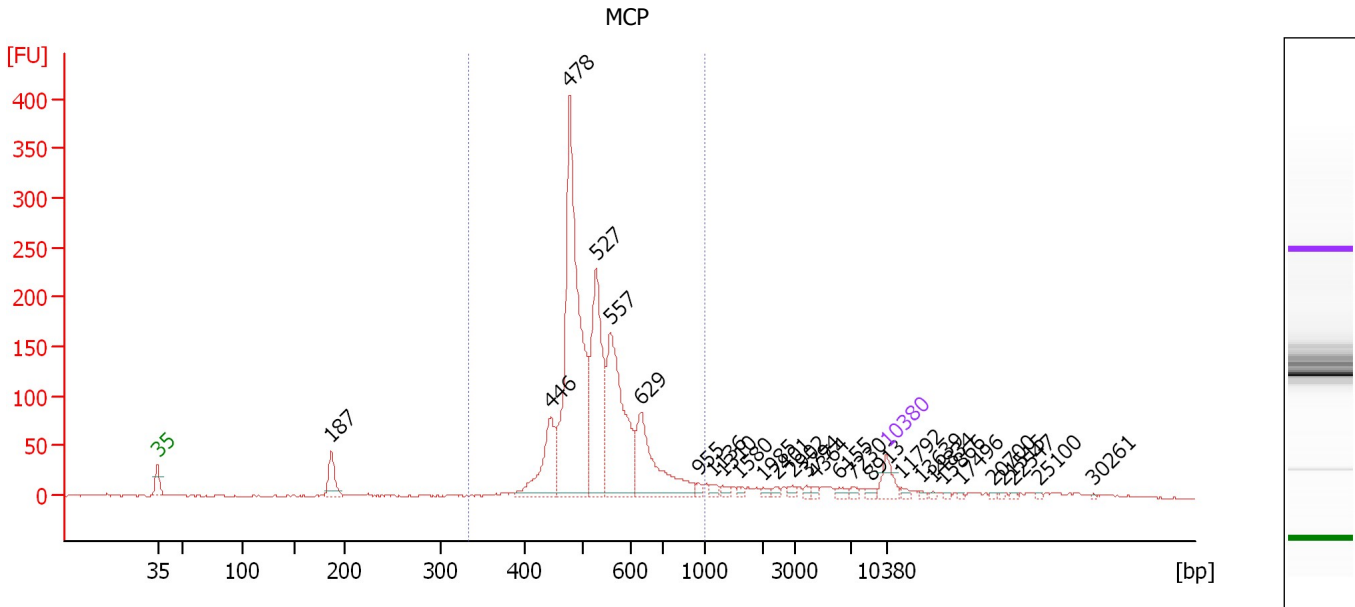
Region table for sample 8 : G23

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
427	1,575	663	1,348.3	3,380.7	1,426.96	81	22.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
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Electropherogram Summary Continued ...



Overall Results for sample 9 : MCP

Number of peaks found: 28 Corr. Area 1: 1,779.1
 Noise: 1.1

Peak table for sample 9 : MCP

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	187	152.35	1,235.1		59.66
3	446	344.24	1,170.2		80.78
4	478	1,697.52	5,386.2		82.57
5	527	727.95	2,091.5		85.13
6	557	775.91	2,109.2		86.55
7	629	475.47	1,145.7		89.43
8	955	19.67	31.2		94.98
9	1,136	23.63	31.5		96.34
10	1,310	21.02	24.3		97.32
11	1,580	16.57	15.9		98.85
12	1,985	15.57	11.9		101.13
13	2,401	15.19	9.6		102.44
14	2,902	16.23	8.5		103.96
15	3,794	12.61	5.0		105.33
16	4,364	16.54	5.7		106.09
17	6,155	19.65	4.8		108.48
18	7,230	15.87	3.3		109.84
19	8,913	16.49	2.8		111.53
20	10,380	75.00	10.9	Upper Marker	113.00
21	11,792	0.00	0.0		114.42
22	13,639	0.00	0.0		116.27
23	14,834	0.00	0.0		117.46
24	15,866	0.00	0.0		118.50
25	17,496	0.00	0.0		120.13

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

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Electropherogram Summary Continued ...

... Peak table for sample 9 : MCP

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	20,700	0.00	0.0		123.34
27	21,515	0.00	0.0		124.16
28	22,547	0.00	0.0		125.19
29	25,100	0.00	0.0		127.75
30	30,261	0.00	0.0		132.92

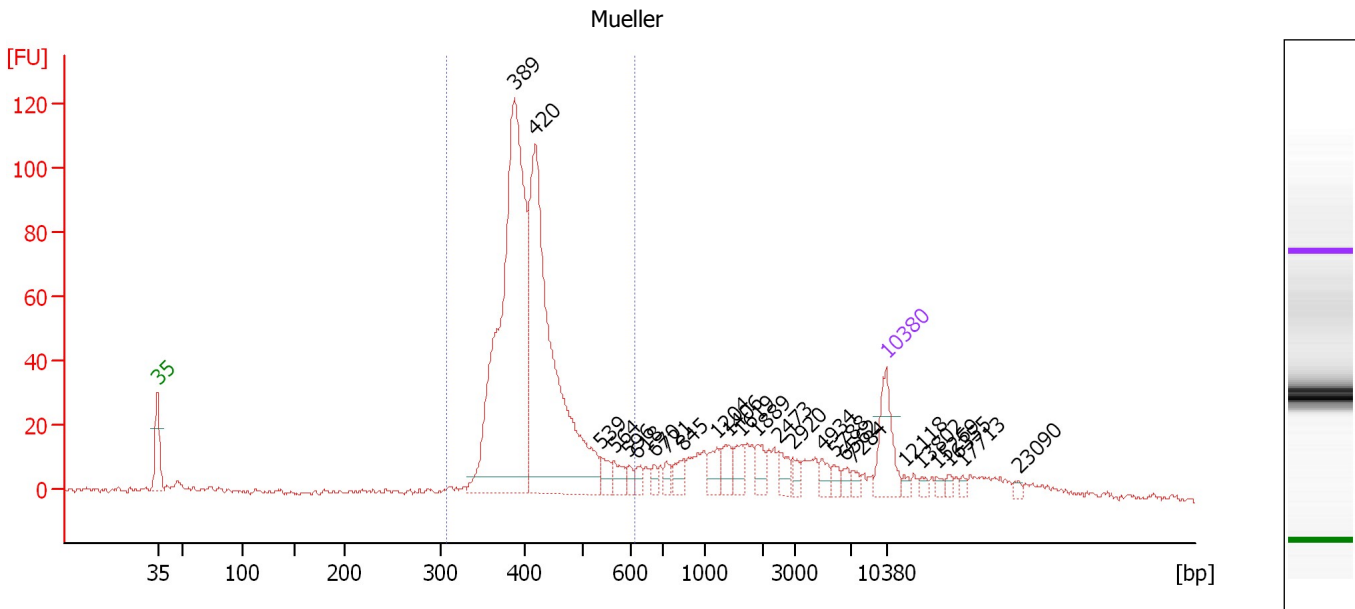
Region table for sample 9 : MCP

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
332	1,000	536	1,779.1	11,816.5	4,063.08	84	16.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 10 : Mueller

Number of peaks found: 25 Corr. Area 1: 798.2
 Noise: 0.8

Peak table for sample 10 : Mueller

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	389	932.86	3,636.9		77.29
3	420	774.55	2,796.4		79.31
4	539	29.19	82.1		85.67
5	564	27.15	72.9		86.87
6	596	14.16	36.0		88.34
7	618	14.09	34.5		89.10
8	670	13.53	30.6		90.68
9	721	16.82	35.3		91.88
10	845	25.75	46.2		93.51
11	1,204	29.72	37.4		96.72
12	1,406	32.99	35.5		97.87
13	1,619	25.21	23.6		99.07
14	1,889	24.96	20.0		100.59
15	2,473	23.39	14.3		102.66
16	2,920	13.00	6.7		104.02
17	4,934	17.08	5.2		106.85
18	5,788	13.18	3.4		107.99
19	6,562	11.61	2.7		109.03
20	7,284	9.57	2.0		109.90
21	10,380	75.00	10.9	Upper Marker	113.00
22	12,118	0.00	0.0		114.74
23	13,802	0.00	0.0		116.43
24	15,269	0.00	0.0		117.90
25	16,355	0.00	0.0		118.99


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

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Electropherogram Summary Continued ...**... Peak table for sample 10 : Mueller**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	17,713	0.00	0.0		120.35
27	23,090	0.00	0.0		125.74

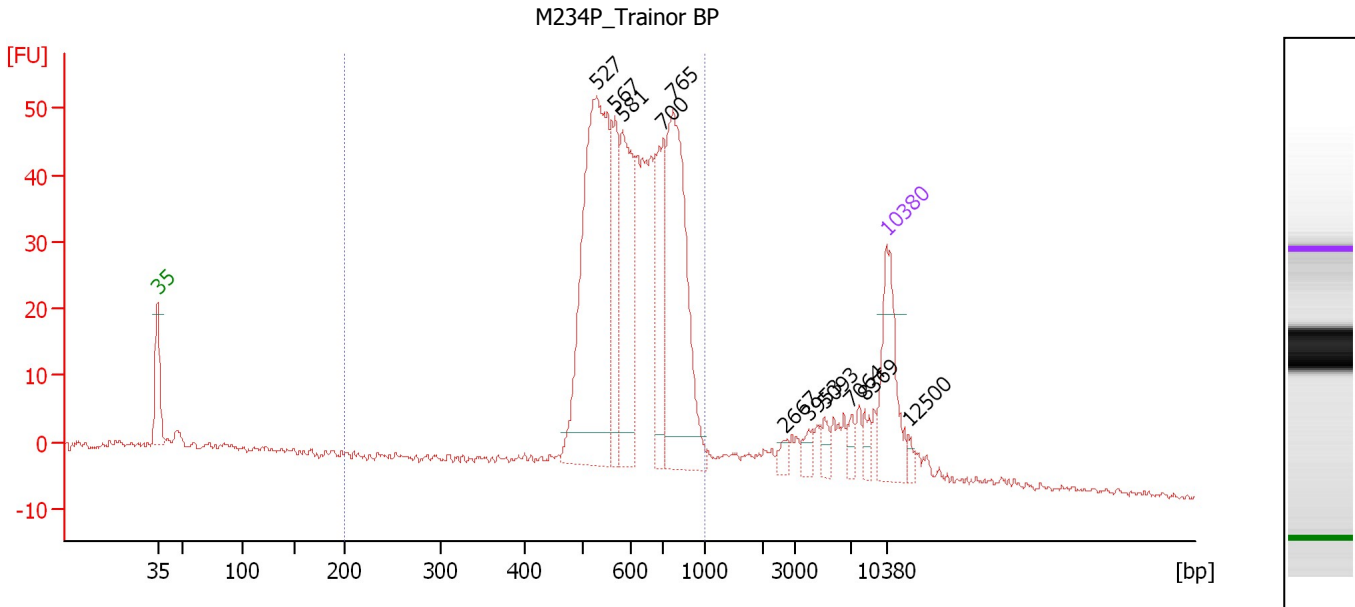
Region table for sample 10 : Mueller

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
306	607	418	798.2	6,558.0	 1,781.85	70	12.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 11 : M234P Trainor BP

Number of peaks found: 11 Corr. Area 1: 612.4
 Noise: 0.5

Peak table for sample 11 : M234P Trainor BP

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	527	402.75	1,158.2		85.11
3	567	87.83	234.6		87.02
4	581	157.82	411.4		87.67
5	700	109.15	236.3		91.59
6	765	259.04	512.8		92.46
7	2,667	8.38	4.8		103.25
8	3,953	11.45	4.4		105.54
9	5,093	10.47	3.1		107.06
10	7,064	9.92	2.1		109.68
11	8,369	10.18	1.8		110.98
12	10,380	75.00	10.9	Upper Marker	113.00
13	12,500	0.00	0.0		115.12

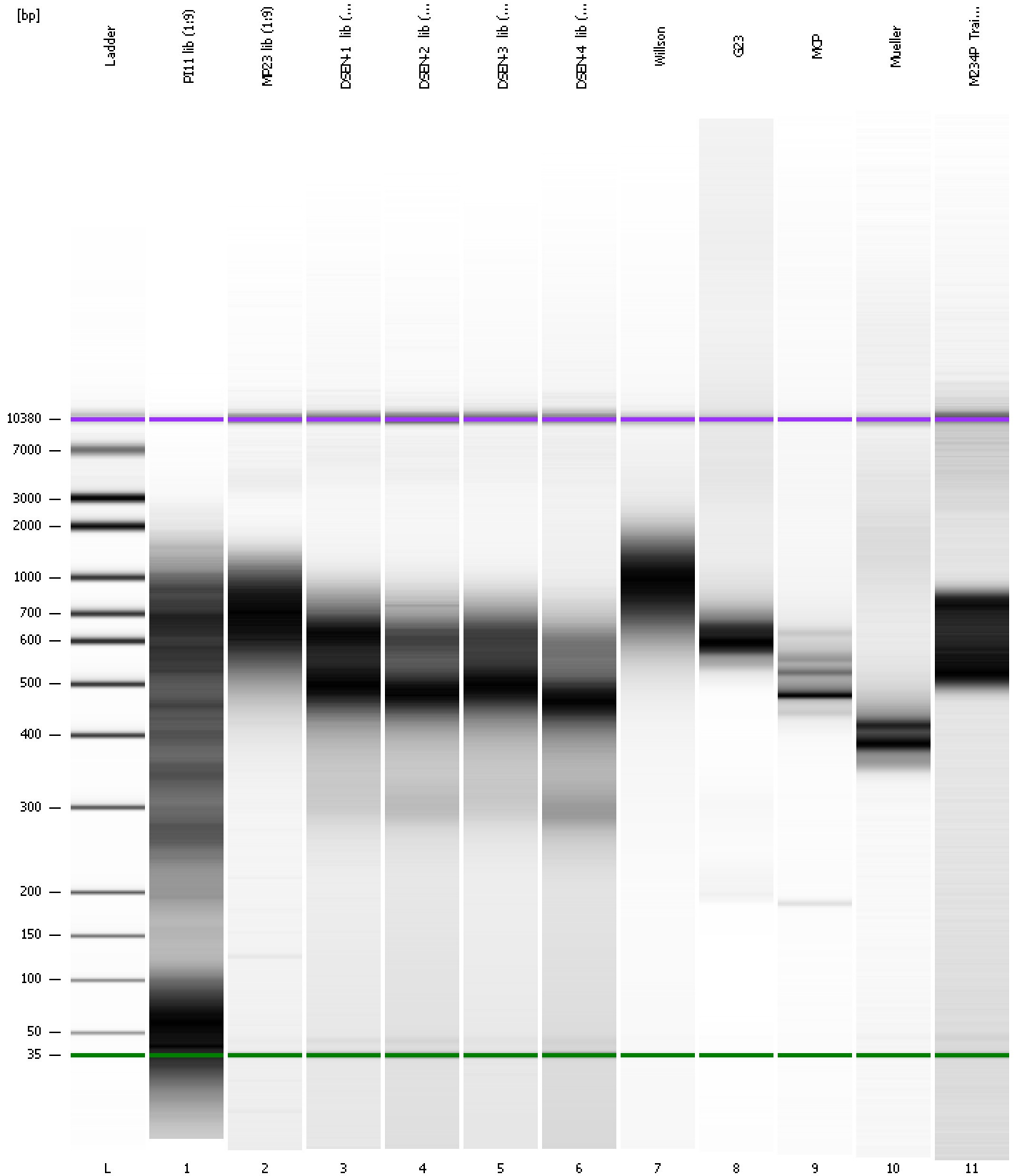
Region table for sample 11 : M234P Trainor BP

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	628	612.4	3,224.8	1,265.88	82	19.5

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
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Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad

Created: 12/7/2015 10:31:26 AM
 Modified: 12/7/2015 11:13:14 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		12/7/2015 11:13:13 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-12-07\2015-12-07_001.xad)		Instrument	Run		12/7/2015 10:31:31 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		12/7/2015 10:31:31 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		12/7/2015 10:31:31 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		12/7/2015 10:31:31 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		12/7/2015 10:31:31 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		12/7/2015 10:31:31 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		12/7/2015 10:31:31 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1