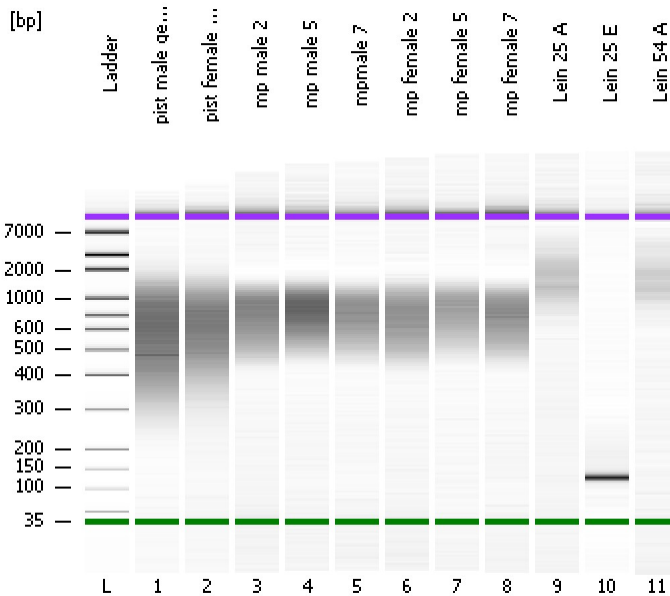


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
Modified: 12/16/2014 2:08:18 PM

**Electrophoresis File Run Summary**



Instrument Information:

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

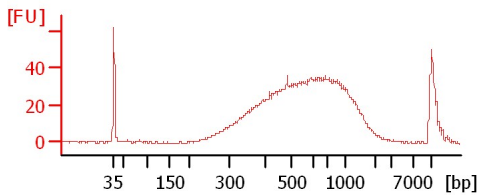
Assay Information:

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

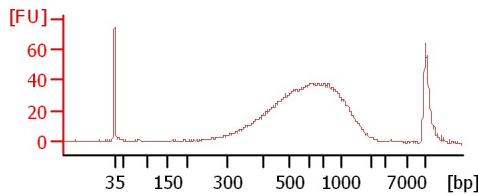
Chip Information:

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

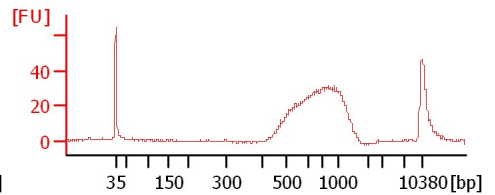
**pist male gen lib**



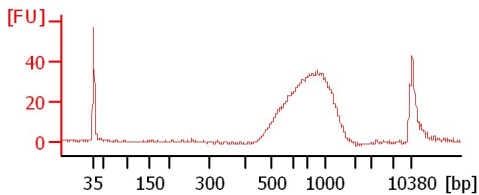
**pist female gen lib**



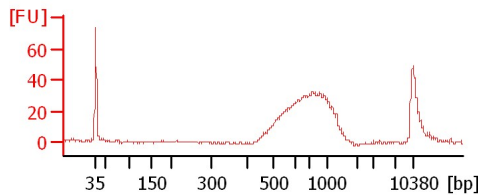
**mp male 2**



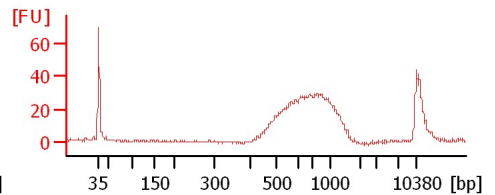
**mp male 5**



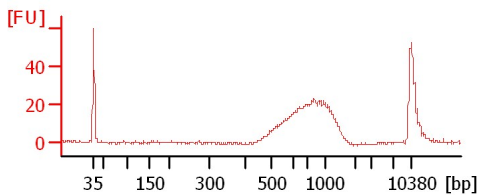
**mpmale 7**



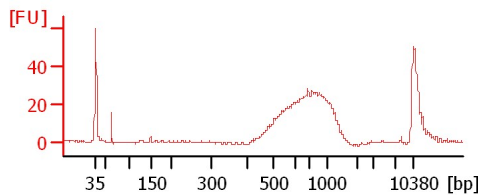
**mp female 2**



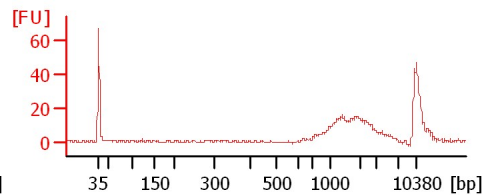
**mp female 5**



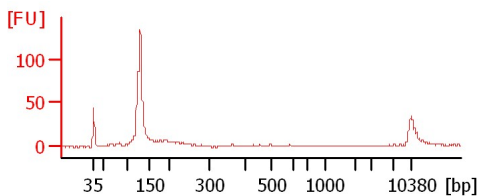
**mp female 7**



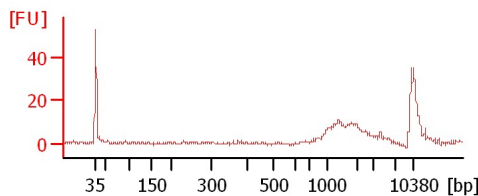
**Lein 25 A**



**Lein 25 E**



**Lein 54 A**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electrophoresis File Run Summary (Chip Summary)**

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
pist male gen lib		<input type="checkbox"/>	✓			
pist female gen lib		<input type="checkbox"/>	✓			
mp male 2		<input type="checkbox"/>	✓			
mp male 5		<input type="checkbox"/>	✓			
mpmale 7		<input type="checkbox"/>	✓			
mp female 2		<input type="checkbox"/>	✓			
mp female 5		<input type="checkbox"/>	✓			
mp female 7		<input type="checkbox"/>	✓			
Lein 25 A		<input type="checkbox"/>	✓			
Lein 25 E		<input type="checkbox"/>	✓			
Lein 54 A		<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\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
Modified: 12/16/2014 2:08:18 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

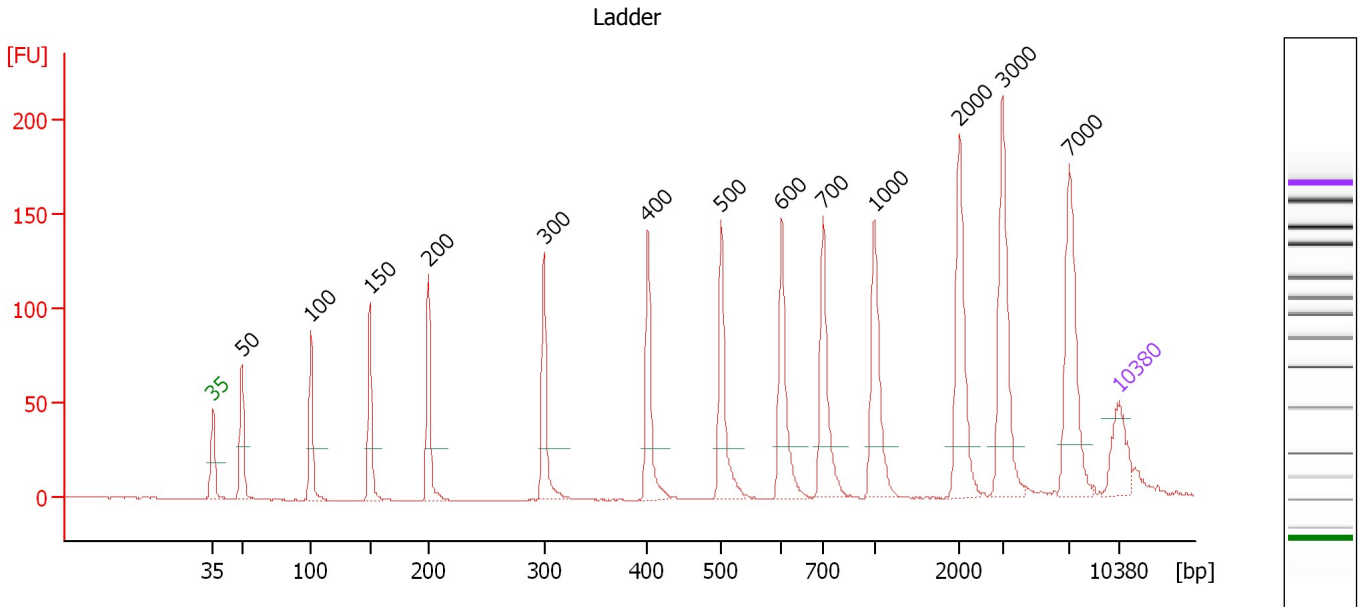
**Ladder**

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\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.2

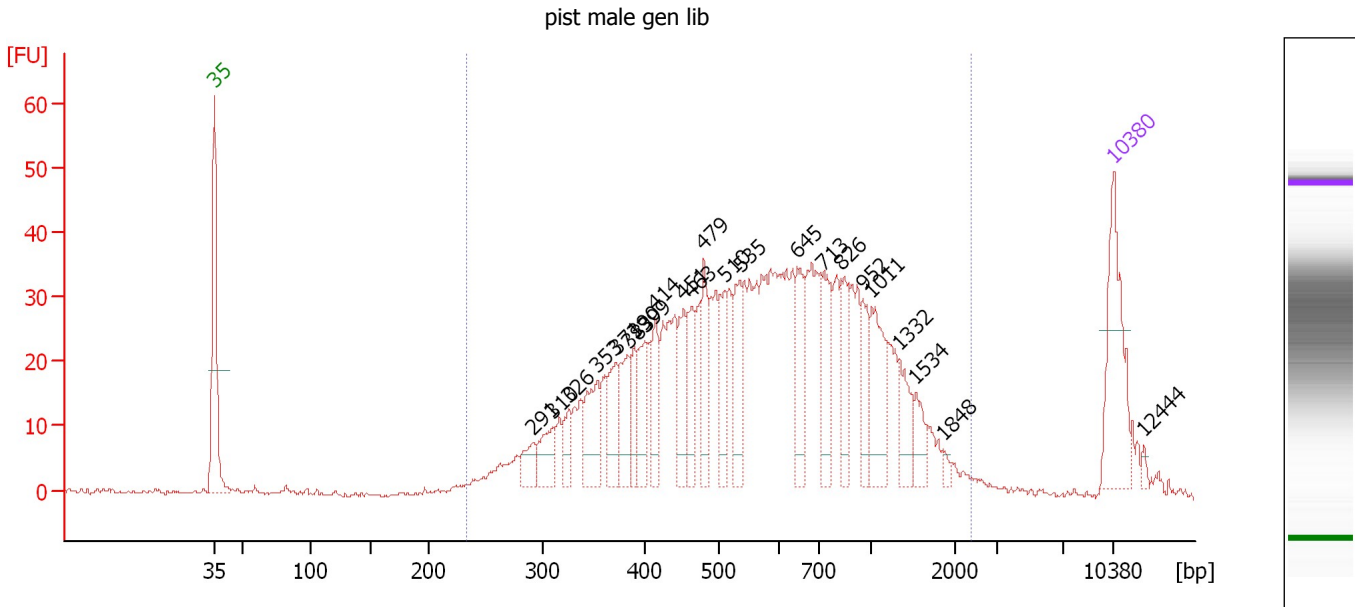
**Peak table for Ladder**

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	45.22
3	100	150.00	2,272.7	Ladder Peak	50.54
4	150	150.00	1,515.2	Ladder Peak	55.11
5	200	150.00	1,136.4	Ladder Peak	59.65
6	300	150.00	757.6	Ladder Peak	68.54
7	400	150.00	568.2	Ladder Peak	76.58
8	500	150.00	454.5	Ladder Peak	82.26
9	600	150.00	378.8	Ladder Peak	86.92
10	700	150.00	324.7	Ladder Peak	90.13
11	1,000	150.00	227.3	Ladder Peak	94.09
12	2,000	150.00	113.6	Ladder Peak	100.68
13	3,000	150.00	75.8	Ladder Peak	103.98
14	7,000	150.00	32.5	Ladder Peak	109.13
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : pist male gen lib**

Number of peaks found: 23                      Corr. Area 1: 1,038.3  
 Noise: 0.3

**Peak table for sample 1 : pist male gen lib**

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	291	24.27	126.2		67.77
3	310	37.26	182.1		69.35
4	326	22.26	103.6		70.59
5	353	55.26	237.3		72.79
6	371	44.08	179.8		74.29
7	383	45.84	181.4		75.20
8	390	29.86	116.1		75.74
9	399	42.32	160.8		76.49
10	414	40.50	148.3		77.36
11	451	46.27	155.5		79.47
12	463	38.93	127.5		80.14
13	479	47.02	148.6		81.09
14	510	43.54	129.2		82.75
15	535	49.33	139.6		83.91
16	645	45.34	106.6		88.35
17	713	49.37	104.9		90.30
18	826	37.94	69.6		91.80
19	952	33.10	52.7		93.46
20	1,011	60.01	89.9		94.16
21	1,332	32.52	37.0		96.28
22	1,534	19.22	19.0		97.61
23	1,848	5.08	4.2		99.68
24	10,380	75.00	10.9	Upper Marker	113.00
25	12,444	0.00	0.0		115.37

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**

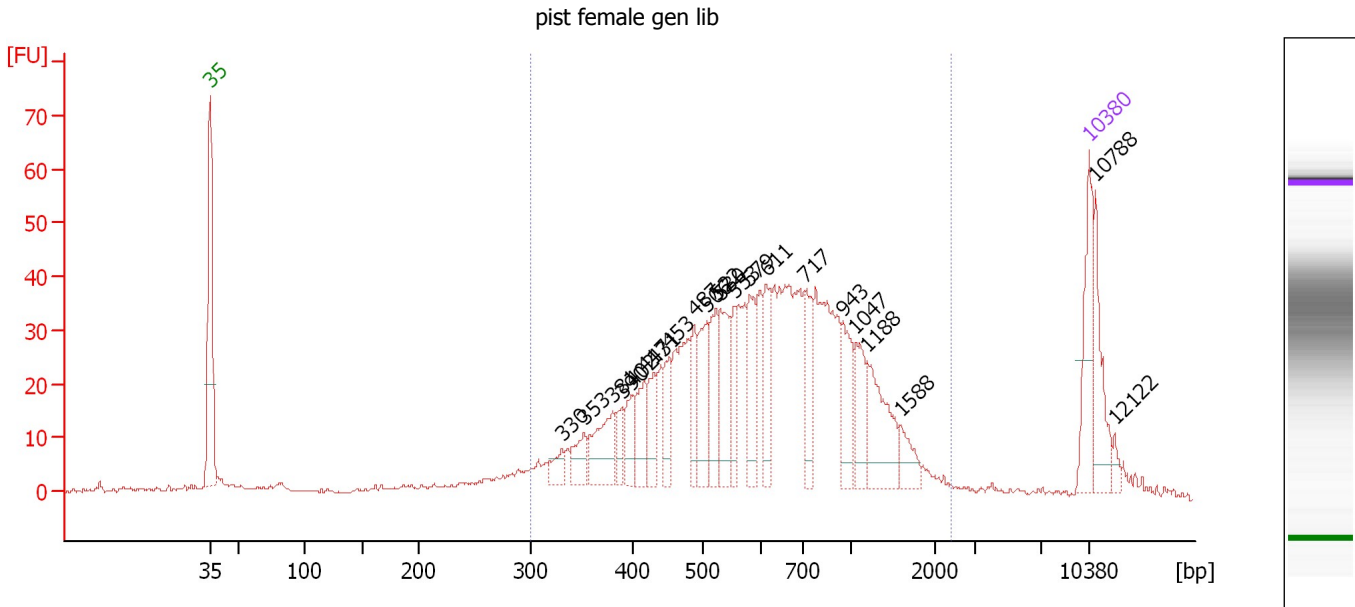
... Region table for sample 1 : pist male gen lib

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
233	665	2,385	1,038.3	97	52.7	1,841.35	5,502.0	■

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : pist female gen lib**

Number of peaks found: 22                      Corr. Area 1: 896.9  
 Noise: 0.3

**Peak table for sample 2 : pist female gen lib**

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	330	25.23	115.7		70.99
3	353	41.13	176.7		72.78
4	381	86.24	342.8		75.07
5	390	27.38	106.3		75.79
6	402	41.80	157.6		76.68
7	417	62.52	227.3		77.53
8	431	51.84	182.2		78.34
9	453	48.87	163.6		79.57
10	487	58.57	182.2		81.53
11	506	75.43	225.9		82.54
12	522	70.97	206.2		83.27
13	530	87.58	250.5		83.65
14	553	53.57	146.9		84.71
15	579	75.50	197.6		85.94
16	611	60.24	149.5		87.26
17	717	66.81	141.1		90.36
18	943	67.51	108.5		93.33
19	1,047	60.17	87.1		94.40
20	1,188	97.13	123.8		95.33
21	1,588	31.07	29.6		97.96
22	10,380	75.00	10.9	Upper Marker	113.00
23	10,788	0.00	0.0		113.47
24	12,122	0.00	0.0		115.00

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**

... Region table for sample 2 : pist female gen lib

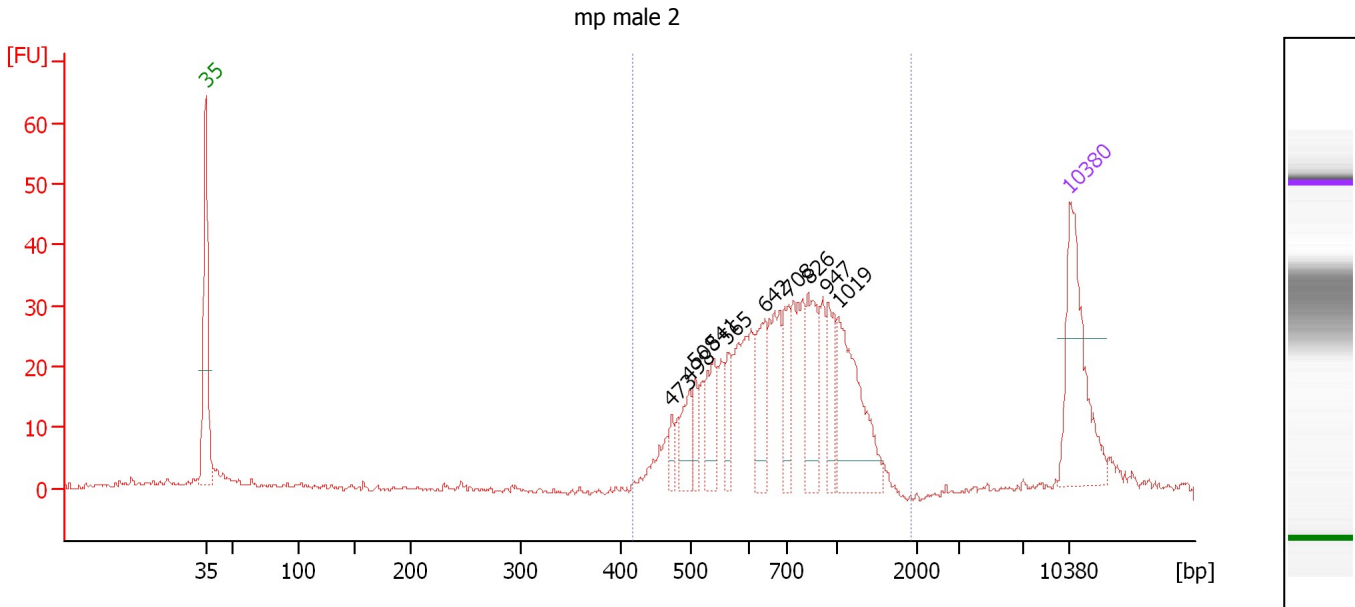
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/ $\mu$ l]	Molarity [pmol/l]	Color
299	695	2,382	896.9	90	47.0	2,101.60	5,659.5	■



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : mp male 2**

Number of peaks found: 10                      Corr. Area 1: 456.4  
 Noise: 0.3

**Peak table for sample 3 : mp male 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	473	11.65	37.3		80.74
3	498	26.70	81.2		82.17
4	508	17.72	52.8		82.64
5	541	35.96	100.8		84.15
6	565	20.80	55.8		85.28
7	642	44.52	105.1		88.26
8	708	32.85	70.3		90.24
9	826	53.38	97.9		91.80
10	947	31.58	50.5		93.39
11	1,019	82.66	122.9		94.22
12	10,380	75.00	10.9	Upper Marker	113.00

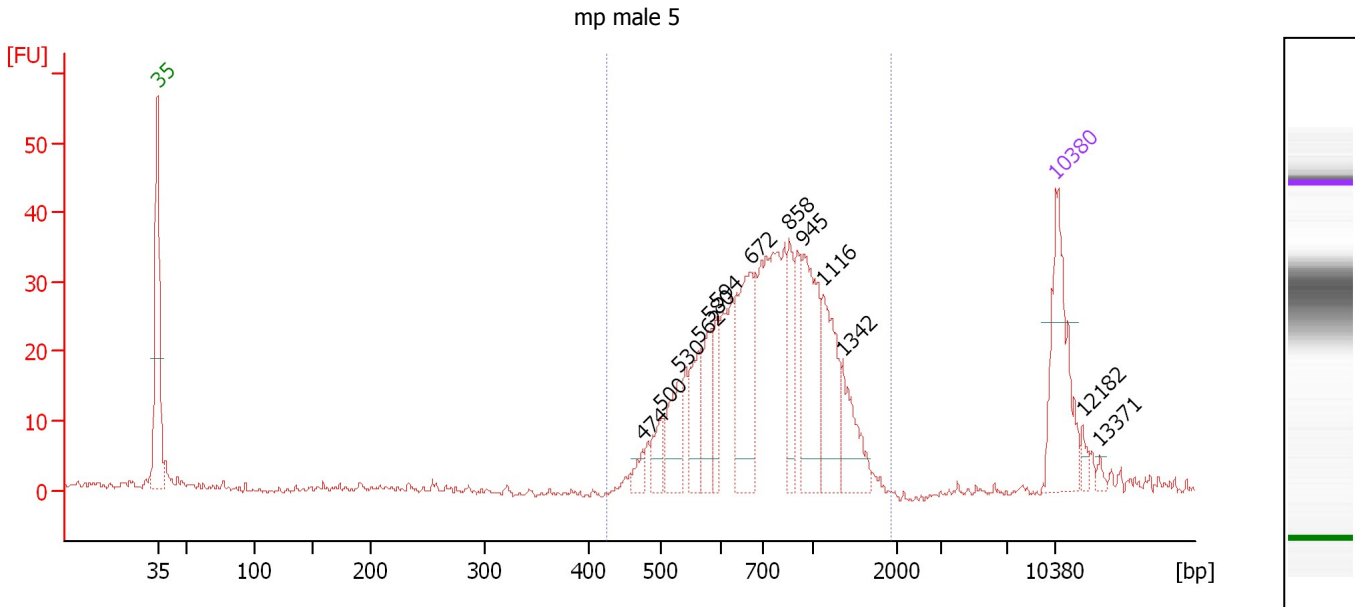
**Region table for sample 3 : mp male 2**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
419	762	1,912	456.4	91	32.5	602.58	1,340.9	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : mp male 5**

Number of peaks found: 13                      Corr. Area 1: 466.4  
 Noise: 0.5

**Peak table for sample 4 : mp male 5**

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	474	12.52	40.0		80.79
3	500	21.03	63.8		82.24
4	530	42.13	120.5		83.65
5	562	40.53	109.3		85.14
6	580	37.26	97.4		85.98
7	594	28.91	73.8		86.63
8	672	101.09	228.0		89.23
9	858	40.19	71.0		92.22
10	945	86.58	138.9		93.36
11	1,116	60.17	81.7		94.85
12	1,342	40.42	45.6		96.35
13	10,380	75.00	10.9	Upper Marker	113.00
14	12,182	0.00	0.0		115.07
15	13,371	0.00	0.0		116.43

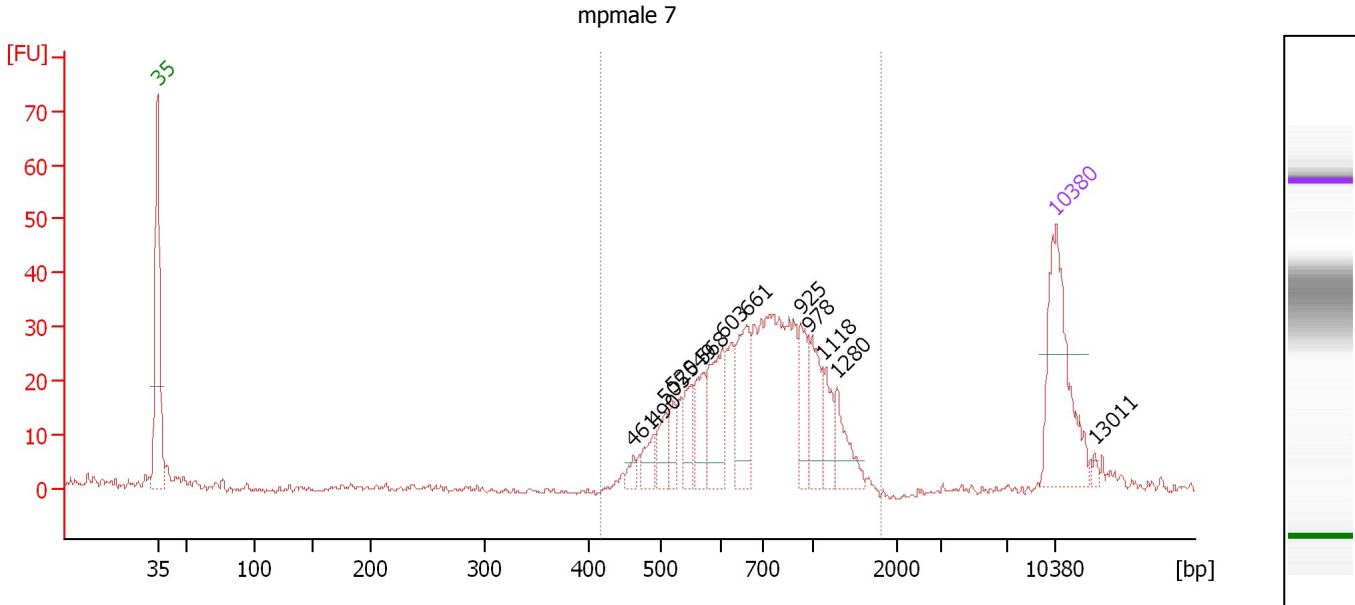
**Region table for sample 4 : mp male 5**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
426	810	1,921	466.4	94	33.0	786.15	1,647.8	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : mpmale 7**

Number of peaks found: 13                      Corr. Area 1: 431.2  
 Noise: 0.5

**Peak table for sample 5 : mpmale 7**

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	461	6.85	22.5		80.06
3	490	16.17	50.0		81.70
4	505	19.35	58.0		82.50
5	520	15.45	45.0		83.21
6	549	21.45	59.2		84.54
7	568	29.14	77.7		85.43
8	603	51.74	130.0		87.02
9	661	51.72	118.5		88.88
10	925	31.16	51.1		93.09
11	978	33.13	51.3		93.80
12	1,118	20.57	27.9		94.87
13	1,280	23.73	28.1		95.93
14	10,380	75.00	10.9	Upper Marker	113.00
15	13,011	0.00	0.0		116.01

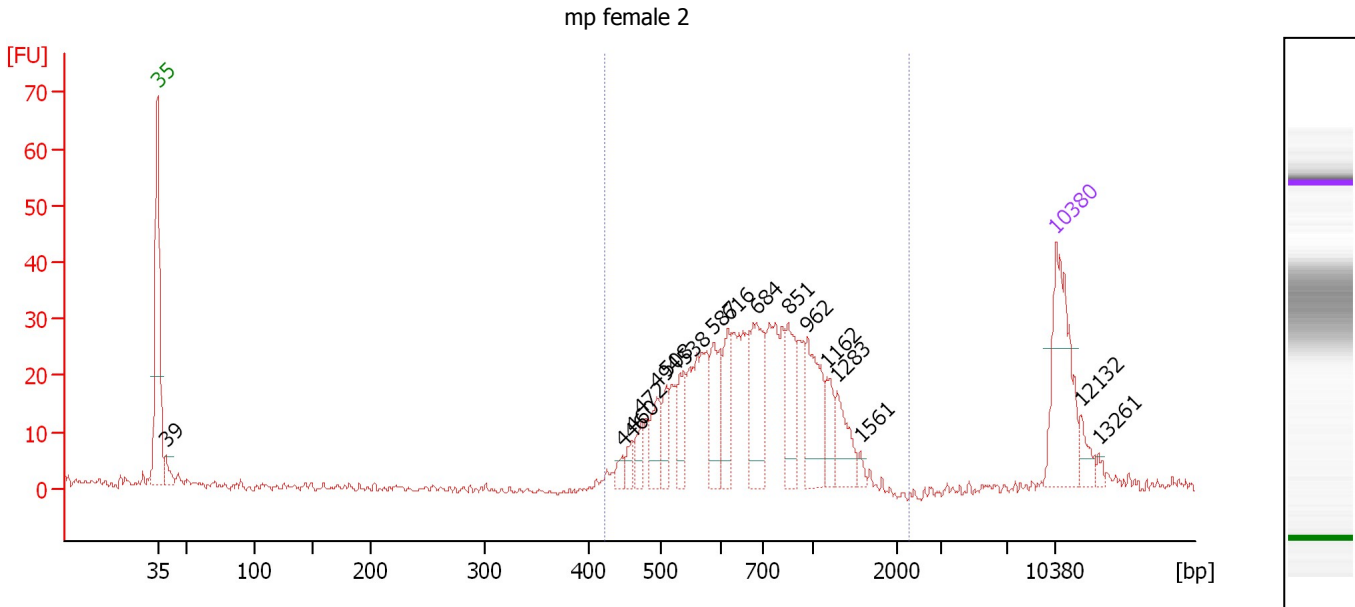
**Region table for sample 5 : mpmale 7**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
417	767	1,800	431.2	93	31.3	521.29	1,140.2	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : mp female 2**

Number of peaks found: 17      Corr. Area 1: 424.3  
 Noise: 0.7

**Peak table for sample 6 : mp female 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	39	14.73	569.3		43.62
3	446	9.02	30.7		79.18
4	460	10.09	33.2		79.99
5	472	15.99	51.3		80.70
6	494	27.70	85.0		81.90
7	508	24.07	71.9		82.62
8	538	24.94	70.2		84.05
9	587	43.23	111.5		86.32
10	616	37.11	91.3		87.44
11	684	67.63	149.8		89.62
12	851	43.66	77.8		92.12
13	962	56.59	89.1		93.59
14	1,162	21.13	27.6		95.15
15	1,283	29.73	35.1		95.96
16	1,561	4.32	4.2		97.79
17	10,380	75.00	10.9	Upper Marker	113.00
18	12,132	0.00	0.0		115.01
19	13,261	0.00	0.0		116.30

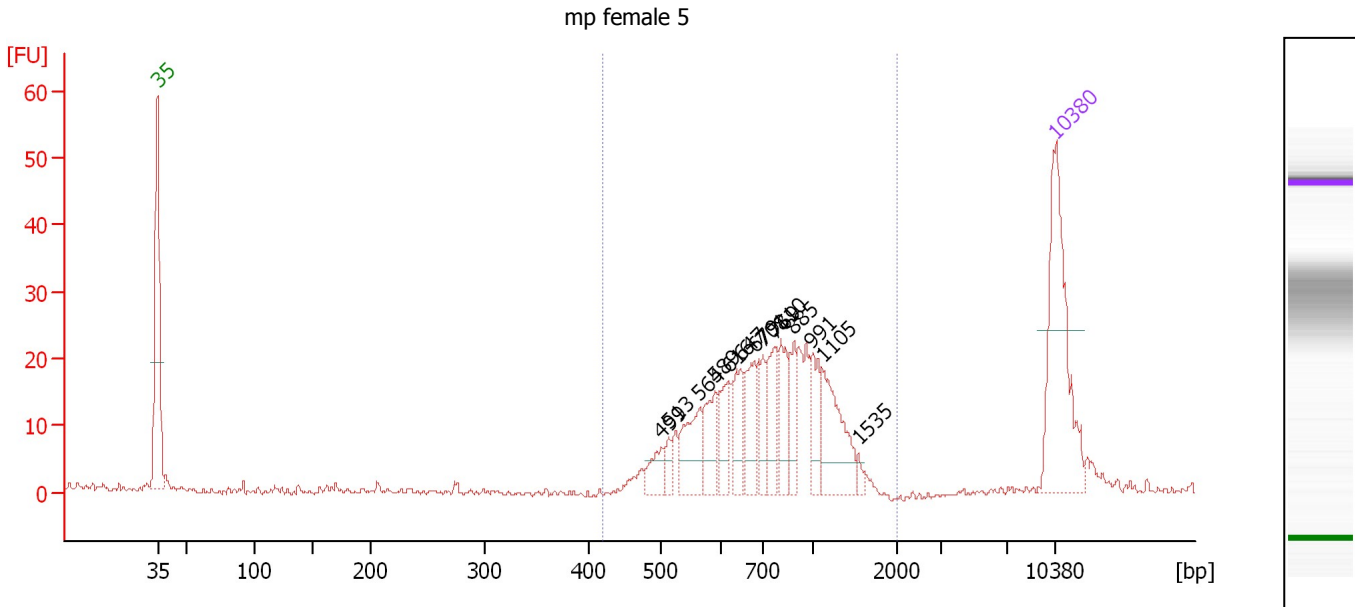
**Region table for sample 6 : mp female 2**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
422	750	2,265	424.3	95	33.0	660.95	1,493.9	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : mp female 5**

Number of peaks found: 14      Corr. Area 1: 263.4  
 Noise: 0.4

**Peak table for sample 7 : mp female 5**

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	499	14.75	44.8		82.23
3	513	7.99	23.6		82.86
4	564	31.71	85.2		85.24
5	589	25.36	65.2		86.41
6	616	19.89	48.9		87.45
7	647	23.59	55.2		88.44
8	679	30.23	67.4		89.47
9	701	19.42	42.0		90.15
10	769	28.10	55.4		91.05
11	810	25.15	47.0		91.59
12	885	17.45	29.9		92.58
13	991	21.89	33.5		93.97
14	1,105	43.27	59.3		94.78
15	1,535	3.03	3.0		97.61
16	10,380	75.00	10.9	Upper Marker	113.00

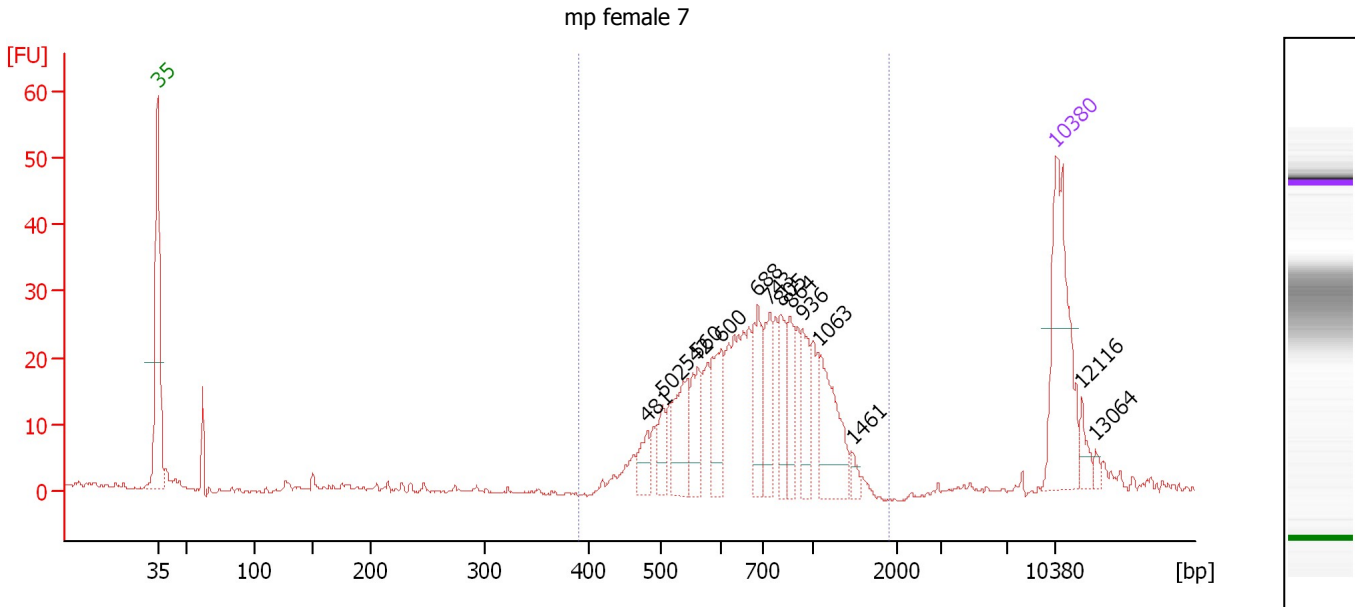
**Region table for sample 7 : mp female 5**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
420	807	1,986	263.4	98	31.2	332.80	693.5	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : mp female 7**

Number of peaks found: 14      Corr. Area 1: 354.5  
 Noise: 0.4

**Peak table for sample 8 : mp female 7**

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	481	16.47	51.9		81.17
3	502	15.69	47.4		82.34
4	542	39.50	110.3		84.24
5	560	28.45	77.0		85.05
6	600	35.80	90.4		86.91
7	688	35.25	77.6		89.76
8	743	32.47	66.2		90.71
9	805	28.03	52.7		91.52
10	864	23.10	40.5		92.29
11	936	25.13	40.7		93.24
12	1,063	50.33	71.7		94.51
13	1,461	5.52	5.7		97.13
14	10,380	75.00	10.9	Upper Marker	113.00
15	12,116	0.00	0.0		114.99
16	13,064	0.00	0.0		116.07

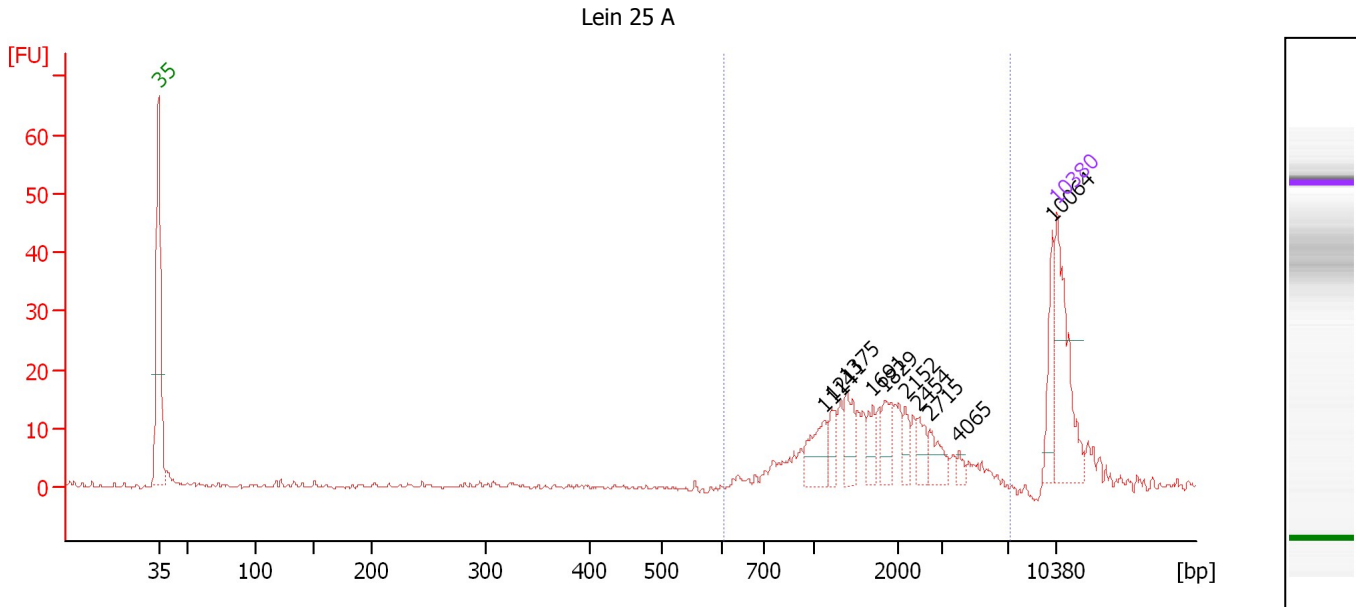
**Region table for sample 8 : mp female 7**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
391	756	1,900	354.5	93	30.7	486.01	1,077.3	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : Lein 25 A**

Number of peaks found: 10                      Corr. Area 1: 158.9  
 Noise: 0.3

**Peak table for sample 9 : Lein 25 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	1,114	34.52	46.9		94.84
3	1,211	15.37	19.2		95.48
4	1,375	27.09	29.8		96.56
5	1,691	16.00	14.3		98.65
6	1,829	20.99	17.4		99.55
7	2,152	11.91	8.4		101.18
8	2,454	15.46	9.5		102.18
9	2,715	17.75	9.9		103.04
10	4,065	5.48	2.0		105.35
11	10,064	31.95	4.8		112.64
12	10,380	75.00	10.9	Upper Marker	113.00

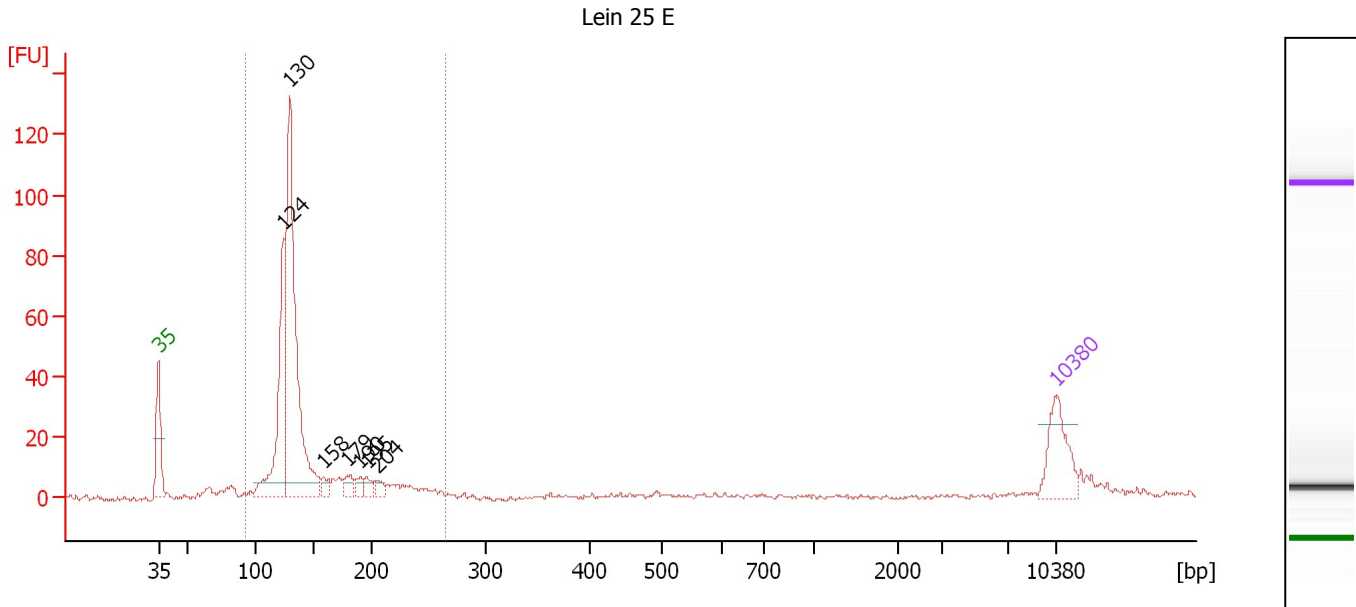
**Region table for sample 9 : Lein 25 A**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
603	1,939	7,132	158.9	81	54.7	280.59	277.9	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : Lein 25 E**

Number of peaks found: 7                      Corr. Area 1: 382.1  
 Noise: 0.8

**Peak table for sample 10 : Lein 25 E**

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	124	283.70	3,463.1		52.75
3	130	648.87	7,585.8		53.25
4	158	17.89	172.0		55.80
5	179	28.00	237.4		57.71
6	190	19.69	157.3		58.71
7	195	18.66	144.8		59.21
8	204	17.54	130.4		59.99
9	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 10 : Lein 25 E**

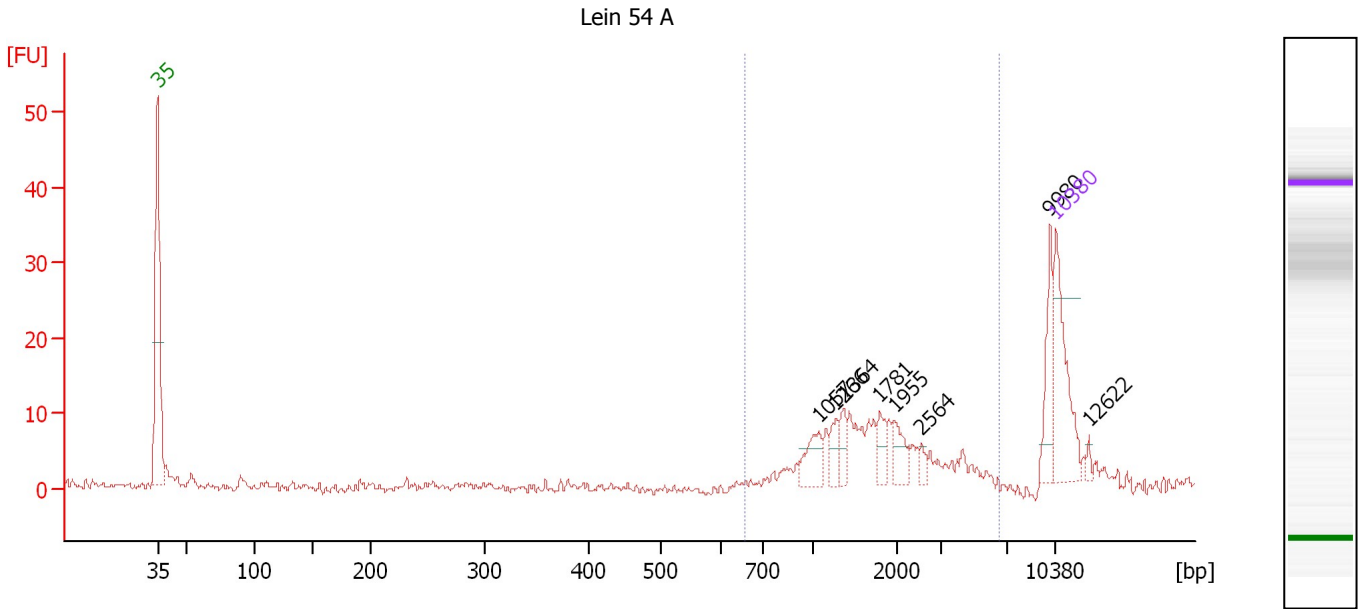
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
93	141	264	382.1	92	20.7	1,067.75	11,942.0	Blue



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : Lein 54 A**

Number of peaks found: 8                      Corr. Area 1: 99.2  
 Noise: 0.5

**Peak table for sample 11 : Lein 54 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	1,057	29.63	42.5		94.47
3	1,266	16.42	19.7		95.84
4	1,364	15.05	16.7		96.49
5	1,781	17.12	14.6		99.24
6	1,955	20.17	15.6		100.39
7	2,564	6.27	3.7		102.54
8	9,980	40.44	6.1		112.54
9	10,380	75.00	10.9	Upper Marker	113.00
10	12,622	0.00	0.0		115.57

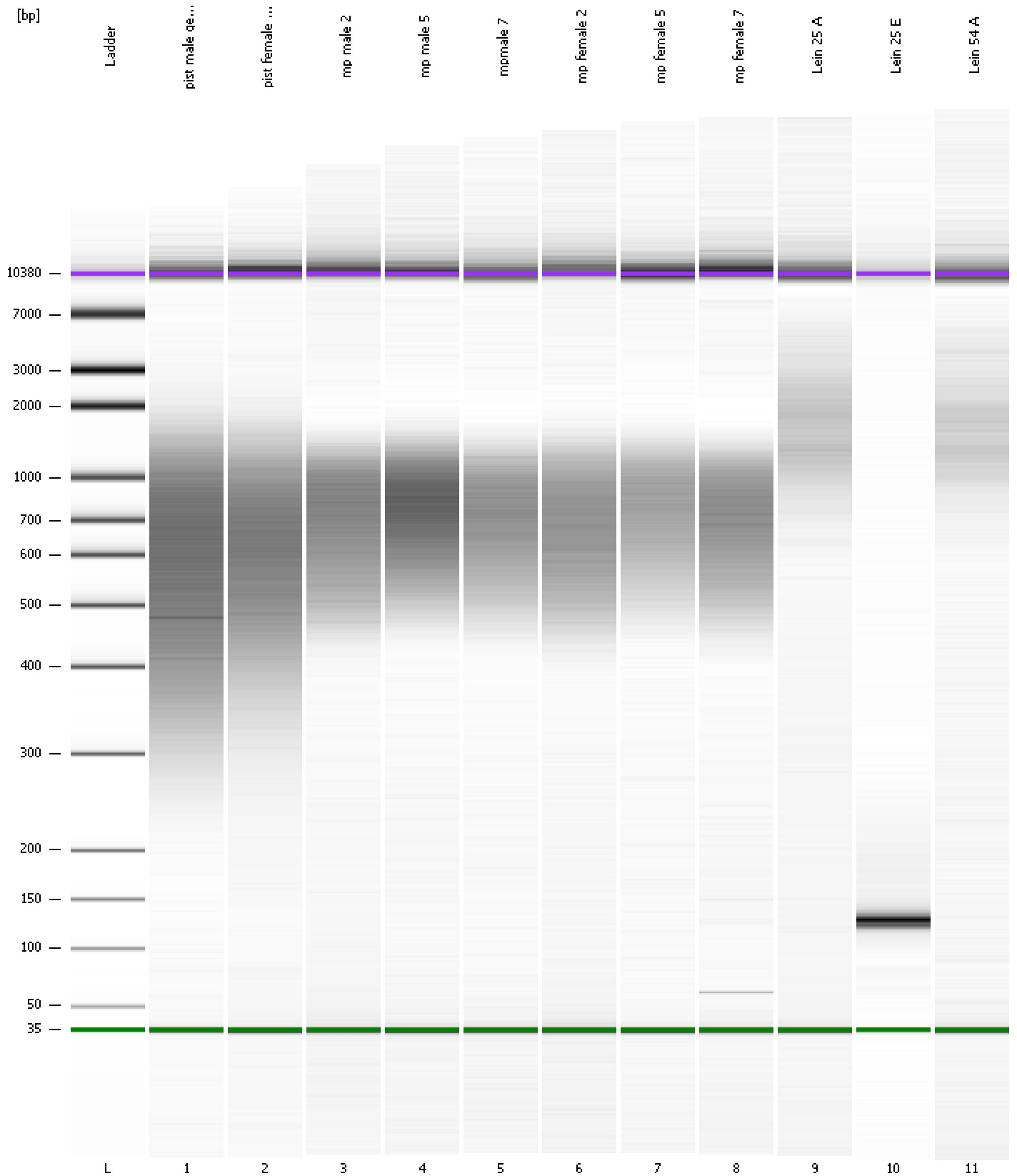
**Region table for sample 11 : Lein 54 A**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
656	1,983	6,543	99.2	73	57.1	246.81	238.7	Blue

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
Modified: 12/16/2014 2:08:18 PM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-16\2014-12-16\_003.xad

Created: 12/16/2014 1:25:01 PM  
 Modified: 12/16/2014 2:08:18 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		12/16/2014 2:06:19 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2014-12-16\2014-12-16_003.xad)		Instrument	Run		12/16/2014 1:25:06 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		12/16/2014 1:25:06 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		12/16/2014 1:25:06 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		12/16/2014 1:25:06 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		12/16/2014 1:25:06 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		12/16/2014 1:25:06 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		12/16/2014 1:25:06 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1