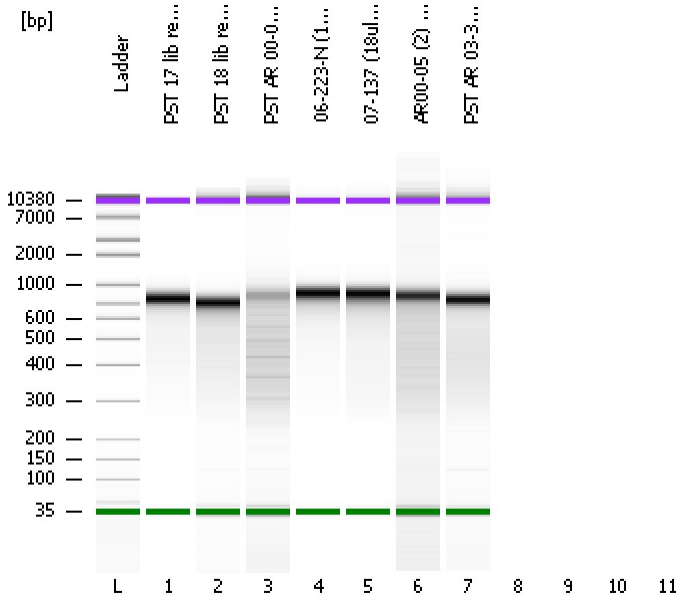


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
Modified: 3/13/2012 11:01:29 AM

Electrophoresis File Run Summary



Instrument Information:

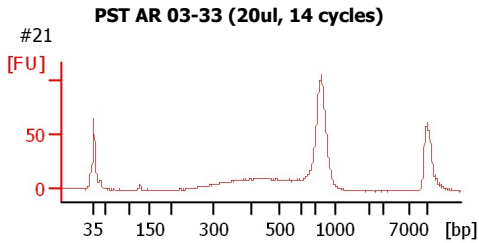
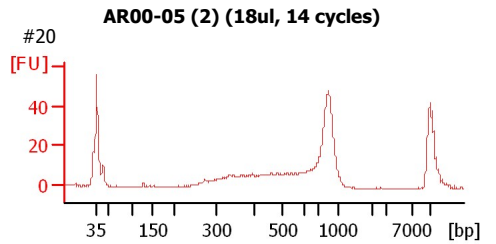
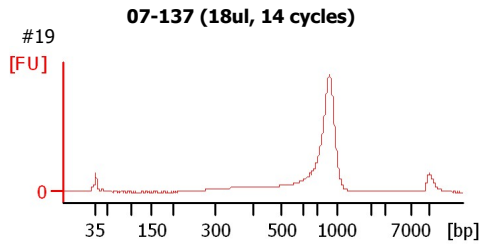
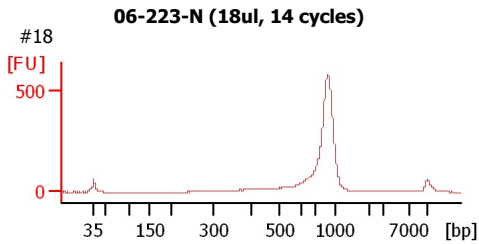
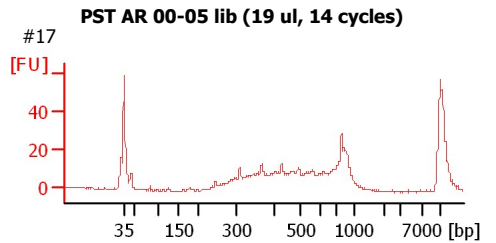
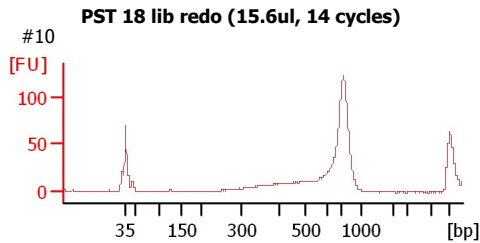
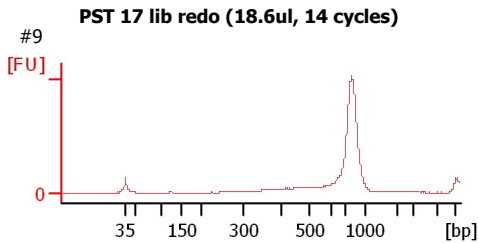
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:\...ents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
 Modified: 3/13/2012 11:01:29 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
PST 17 lib redo (18.6ul, 14 cycles) #9		<input type="checkbox"/>	✓			
PST 18 lib redo (15.6ul, 14 cycles) #10		<input type="checkbox"/>	✓			
PST AR 00-05 lib (19 ul, 14 cycles) #17		<input type="checkbox"/>	✓			
06-223-N (18ul, 14 cycles) #18		<input type="checkbox"/>	✓			
07-137 (18ul, 14 cycles) #19		<input type="checkbox"/>	✓			
AR00-05 (2) (18ul, 14 cycles) #20		<input type="checkbox"/>	✓			
PST AR 03-33 (20ul, 14 cycles) #21		<input type="checkbox"/>	✓			
		<input type="checkbox"/>				
		<input type="checkbox"/>				
		<input type="checkbox"/>				
		<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\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
Modified: 3/13/2012 11:01:29 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

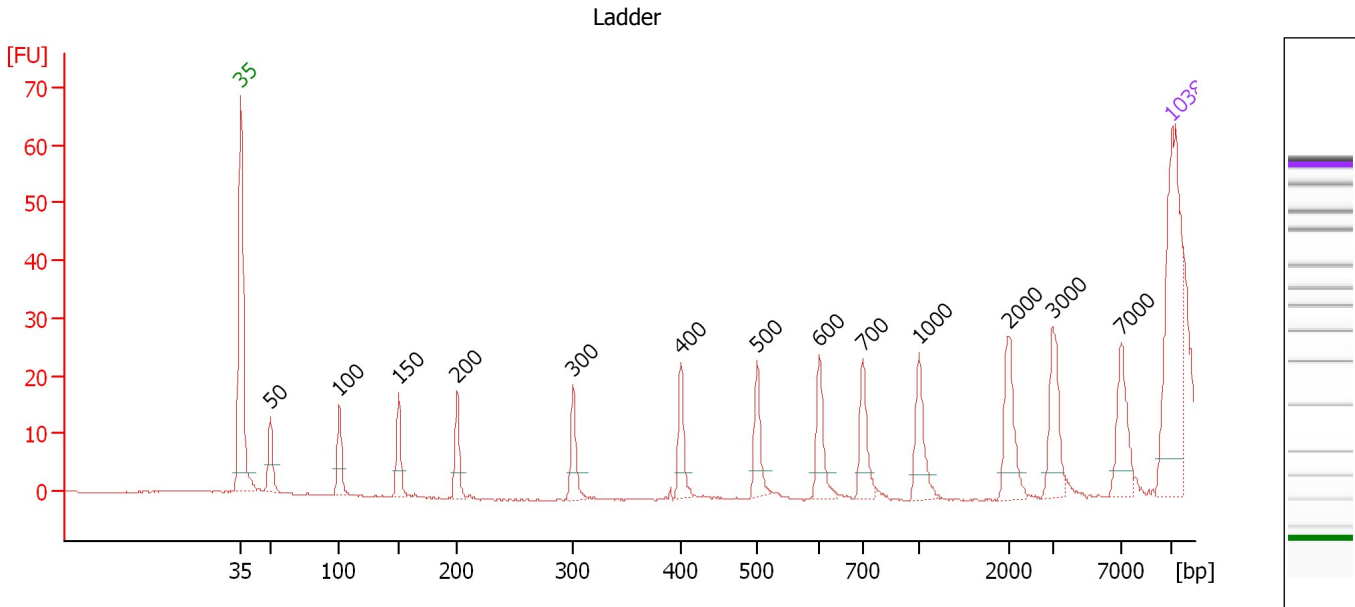
Ladder

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\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
 Modified: 3/13/2012 11:01:29 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

Peak table for Ladder

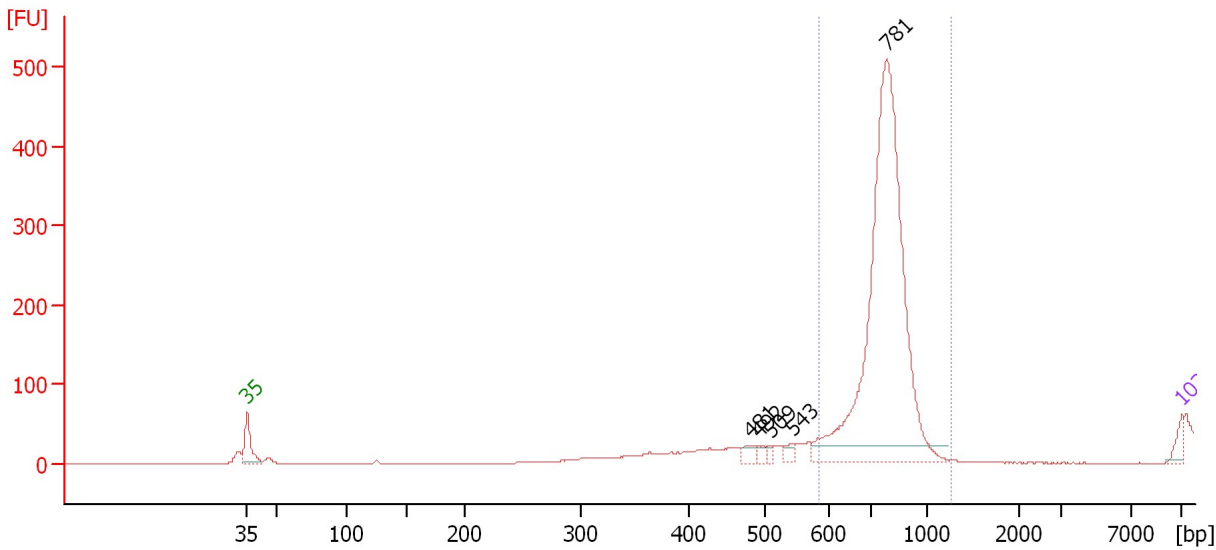
Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	50	150.00	4,545.5	Ladder Peak
3	100	150.00	2,272.7	Ladder Peak
4	150	150.00	1,515.2	Ladder Peak
5	200	150.00	1,136.4	Ladder Peak
6	300	150.00	757.6	Ladder Peak
7	400	150.00	568.2	Ladder Peak
8	500	150.00	454.5	Ladder Peak
9	600	150.00	378.8	Ladder Peak
10	700	150.00	324.7	Ladder Peak
11	1,000	150.00	227.3	Ladder Peak
12	2,000	150.00	113.6	Ladder Peak
13	3,000	150.00	75.8	Ladder Peak
14	7,000	150.00	32.5	Ladder Peak
15	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
 Modified: 3/13/2012 11:01:29 AM

Electropherogram Summary Continued ...

PST 17 lib redo (18.6ul, 14 cycles) [#9]



Setpoint Deviations for sample 1 : PST 17 lib redo (18.6ul, 14 cycles)

Height Threshold [FU] : 20

Overall Results for sample 1 : PST 17 lib redo (18.6ul, 14 cycles)

Number of peaks found: 5 Corr. Area 1: 1,299.1
 Noise: 0.2

Peak table for sample 1 : PST 17 lib redo (18.6ul, 14 cycles)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	481	114.81	361.4	
3	492	64.53	198.8	
4	509	49.50	147.2	
5	543	86.20	240.7	
6	781	5,477.52	10,631.8	
7	10,380	75.00	10.9	Upper Marker

Region table for sample 1 : PST 17 lib redo (18.6ul, 14 cycles)

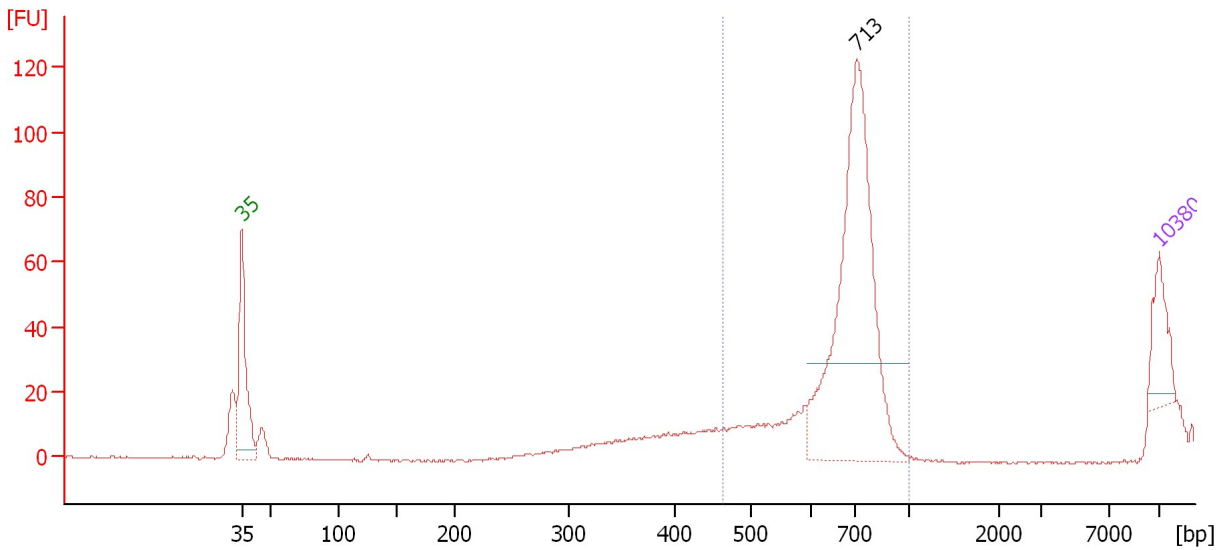
From [bp]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Color
584	8,261.6	782	4,244.12	1,249	1,299.1	99	8.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
 Modified: 3/13/2012 11:01:29 AM

Electropherogram Summary Continued ...

PST 18 lib redo (15.6ul, 14 cycles) [#10]



Setpoint Deviations for sample 2 : PST 18 lib redo (15.6ul, 14 cycles)

Height Threshold [FU] : 30

Overall Results for sample 2 : PST 18 lib redo (15.6ul, 14 cycles)

Number of peaks found: 1 Corr. Area 1: 423.1
 Noise: 0.3

Peak table for sample 2 : PST 18 lib redo (15.6ul, 14 cycles)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	713	772.19	1,641.7	
3	10,380	75.00	10.9	Upper Marker

Region table for sample 2 : PST 18 lib redo (15.6ul, 14 cycles)

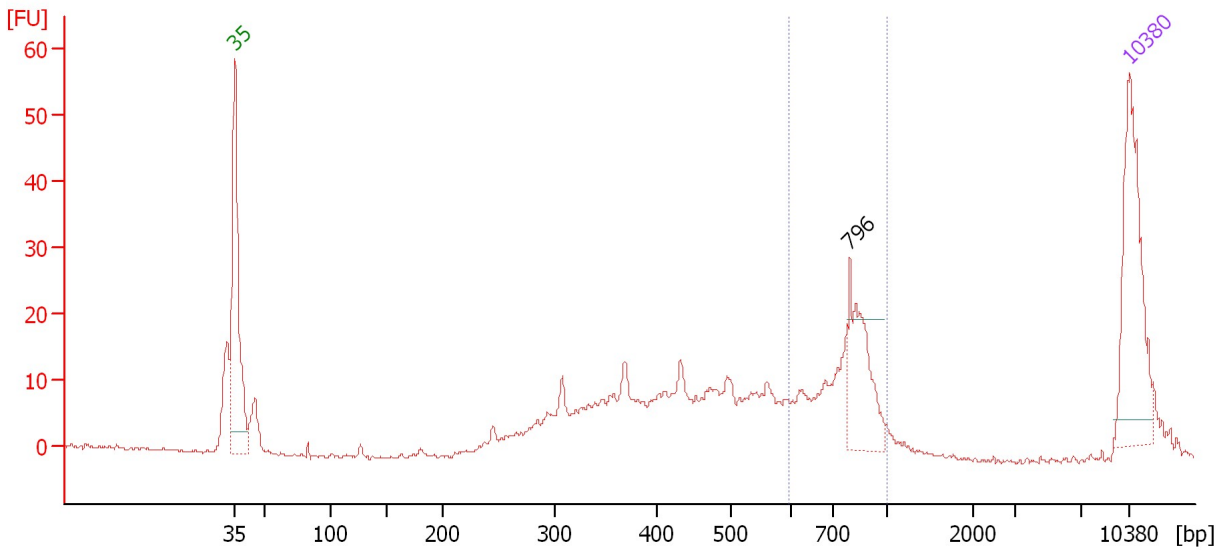
From [bp]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Color
463	1,735.0	695	780.87	1,000	423.1	82	12.1	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
 Modified: 3/13/2012 11:01:29 AM

Electropherogram Summary Continued ...

PST AR 00-05 lib (19 ul, 14 cycles) [#17]



Setpoint Deviations for sample 3 : PST AR 00-05 lib (19 ul, 14 cycles)

Height Threshold [FU] : 20

Overall Results for sample 3 : PST AR 00-05 lib (19 ul, 14 cycles)

Number of peaks found: 1 Corr. Area 1: 104.1
 Noise: 0.3

Peak table for sample 3 : PST AR 00-05 lib (19 ul, 14 cycles)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	796	64.28	122.3	
3	10,380	75.00	10.9	Upper Marker

Region table for sample 3 : PST AR 00-05 lib (19 ul, 14 cycles)

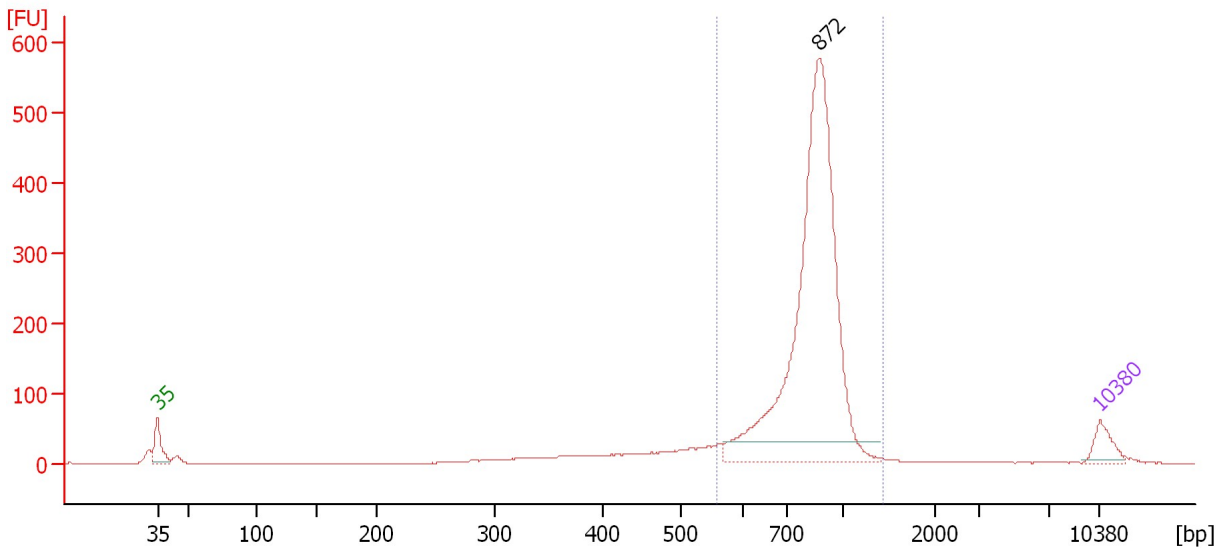
From [bp]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Color
596	249.3	772	124.83	1,000	104.1	27	13.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
 Modified: 3/13/2012 11:01:29 AM

Electropherogram Summary Continued ...

06-223-N (18ul, 14 cycles) [#18]



Setpoint Deviations for sample 4 : 06-223-N (18ul, 14 cycles)

Height Threshold [FU] : 30

Overall Results for sample 4 : 06-223-N (18ul, 14 cycles)

Number of peaks found: 1 Corr. Area 1: 2,120.5
 Noise: 0.4

Peak table for sample 4 : 06-223-N (18ul, 14 cycles)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	872	2,270.50	3,945.6	
3	10,380	75.00	10.9	Upper Marker

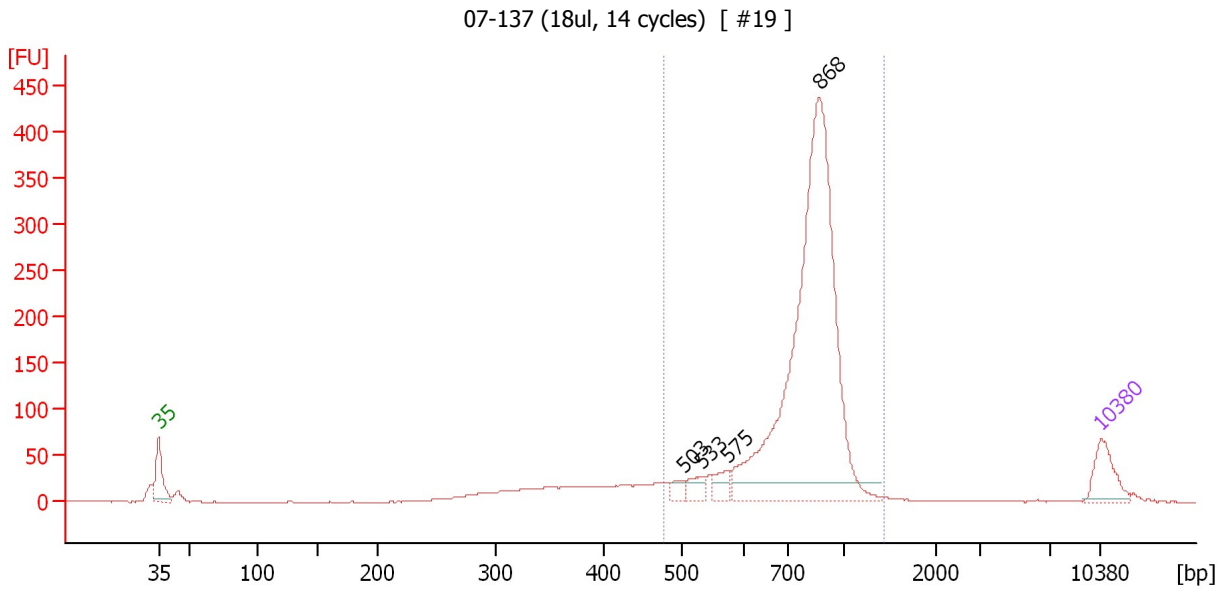
Region table for sample 4 : 06-223-N (18ul, 14 cycles)

From [bp]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Color
557	4,364.5	839	2,361.16	1,437	2,120.5	84	14.7	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
 Modified: 3/13/2012 11:01:29 AM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 5 : 07-137 (18ul, 14 cycles)

Height Threshold [FU] : 20

Overall Results for sample 5 : 07-137 (18ul, 14 cycles)

Number of peaks found: 4 Corr. Area 1: 1,865.1
 Noise: 0.4

Peak table for sample 5 : 07-137 (18ul, 14 cycles)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	503	29.98	90.3	
3	533	44.86	127.6	
4	575	46.01	121.3	
5	868	1,470.40	2,566.5	
6	10,380	75.00	10.9	Upper Marker

Region table for sample 5 : 07-137 (18ul, 14 cycles)

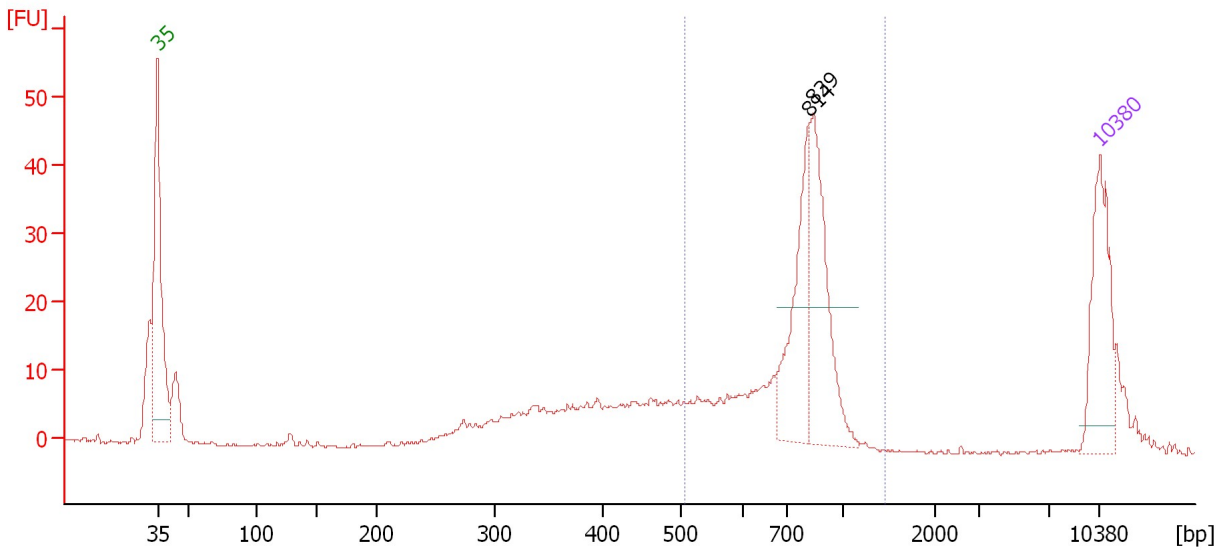
From [bp]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Color
478	3,310.1	802	1,684.32	1,429	1,865.1	82	17.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
 Modified: 3/13/2012 11:01:29 AM

Electropherogram Summary Continued ...

AR00-05 (2) (18ul, 14 cycles) [#20]



Setpoint Deviations for sample 6 : AR00-05 (2) (18ul, 14 cycles)

Height Threshold [FU] : 20

Overall Results for sample 6 : AR00-05 (2) (18ul, 14 cycles)

Number of peaks found: 2 Corr. Area 1: 214.0
 Noise: 0.2

Peak table for sample 6 : AR00-05 (2) (18ul, 14 cycles)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	814	107.43	199.9	
3	839	124.22	224.2	
4	10,380	75.00	10.9	Upper Marker

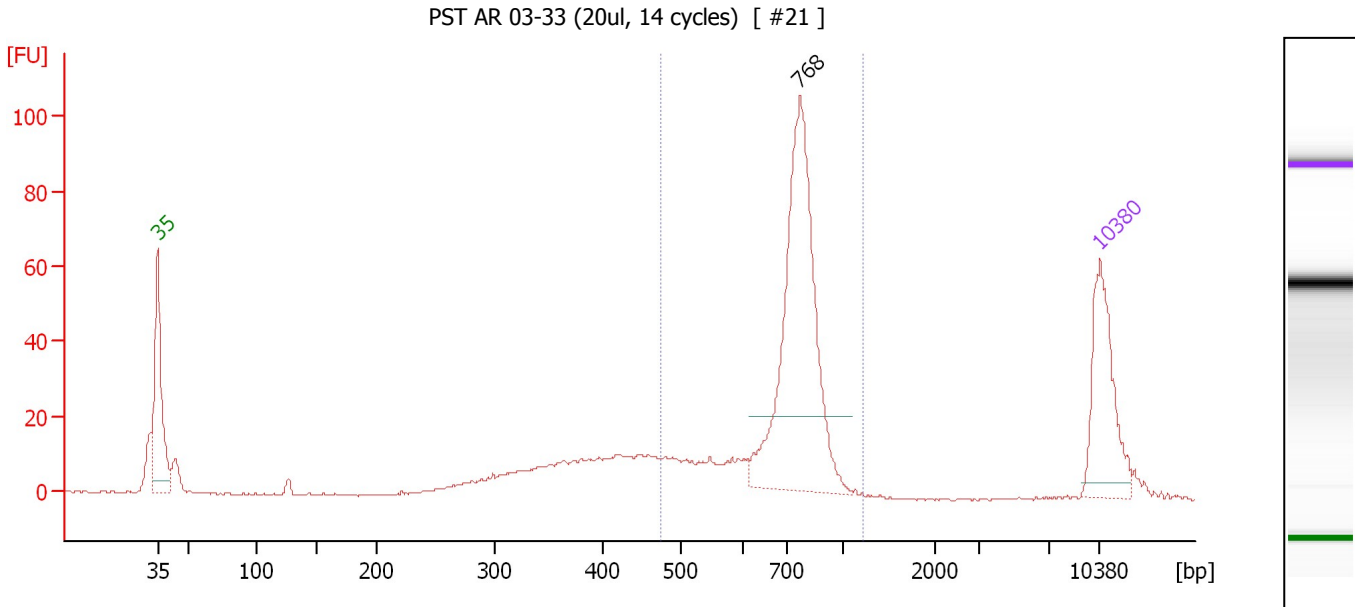
Region table for sample 6 : AR00-05 (2) (18ul, 14 cycles)

From [bp]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Color
508	716.8	764	347.68	1,447	214.0	54	17.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
 Modified: 3/13/2012 11:01:29 AM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 7 : PST AR 03-33 (20ul, 14 cycles)

Height Threshold [FU] : 20

Overall Results for sample 7 : PST AR 03-33 (20ul, 14 cycles)

Number of peaks found: 1 Corr. Area 1: 396.9
 Noise: 0.3

Peak table for sample 7 : PST AR 03-33 (20ul, 14 cycles)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	768	292.89	577.5	
3	10,380	75.00	10.9	Upper Marker

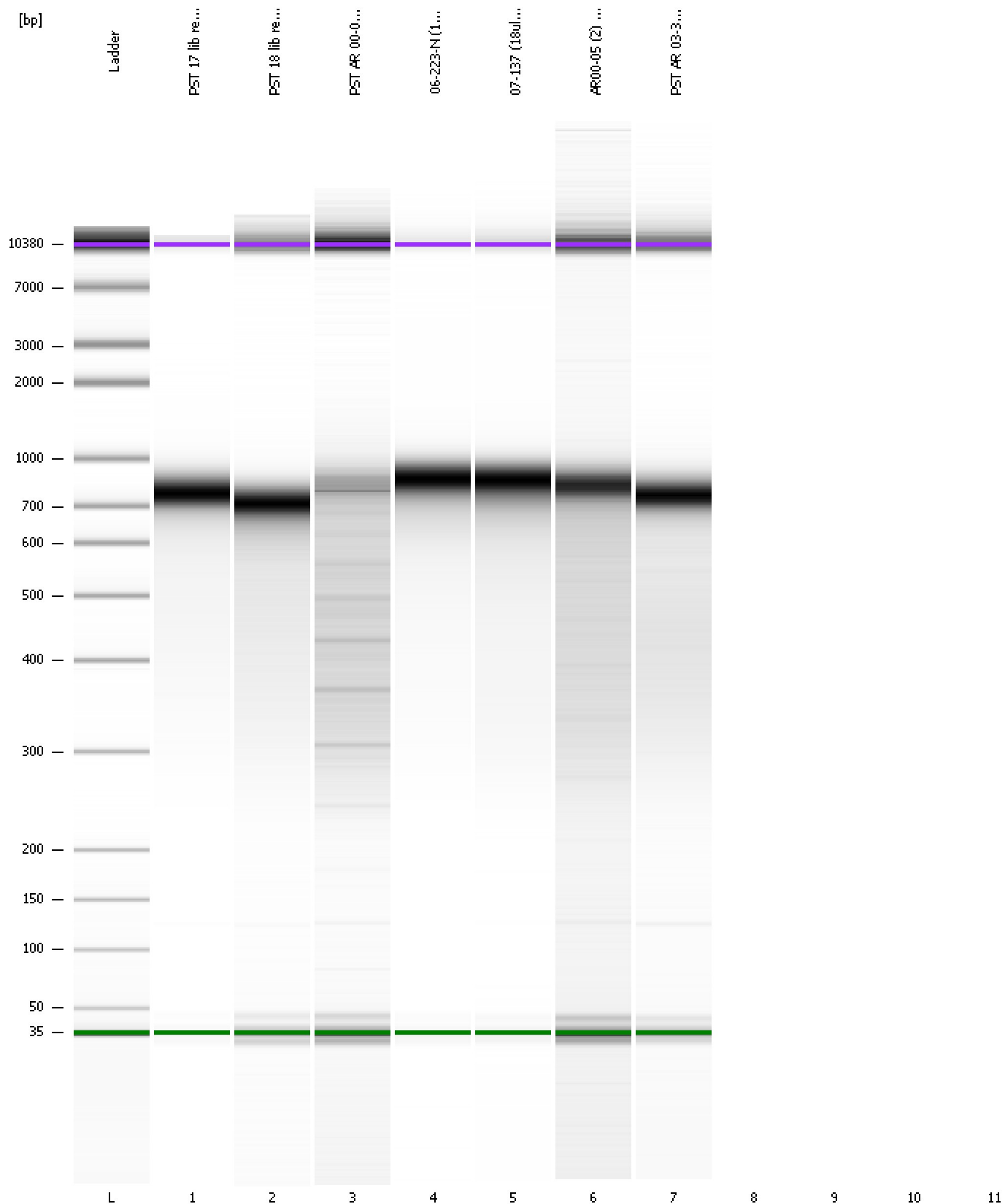
Region table for sample 7 : PST AR 03-33 (20ul, 14 cycles)

From [bp]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Color
474	823.8	728	382.51	1,211	396.9	65	16.0	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
Modified: 3/13/2012 11:01:29 AM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
Modified: 3/13/2012 11:01:29 AM

Invalid Samples

Sample 8 has not been run, no results available.

Sample 9 has not been run, no results available.

Sample 10 has not been run, no results available.

Sample 11 has not been run, no results available.

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad

Created: 3/13/2012 10:29:56 AM
 Modified: 3/13/2012 11:01:29 AM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 8)		Instrument	Run		3/13/2012 10:59:48 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2012-03-13\2012-03-13_002.xad)		Instrument	Run		3/13/2012 10:30:01 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		3/13/2012 10:30:01 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		3/13/2012 10:30:01 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		3/13/2012 10:30:01 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		3/13/2012 10:30:01 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		3/13/2012 10:30:01 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		3/13/2012 10:30:01 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1