

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-05\2017-01-05_003.xad

Created: 1/5/2017 4:07:07 PM
Modified: 1/5/2017 4:25:35 PM

Electrophoresis File Run Summary

Instrument Information:

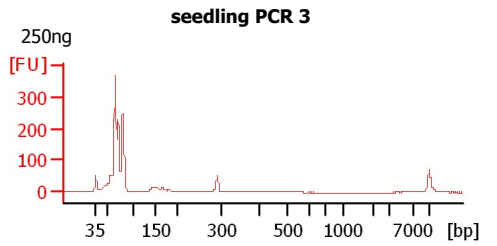
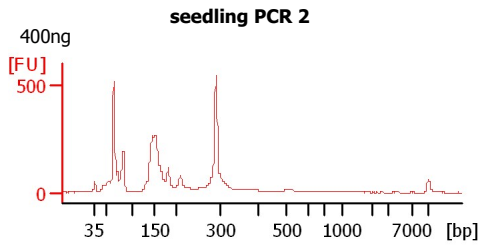
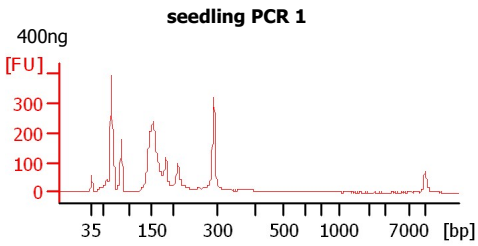
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:



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Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
seedling PCR 1	400ng	<input type="checkbox"/>	✓			
seedling PCR 2	400ng	<input type="checkbox"/>	✓			
seedling PCR 3	250ng	<input type="checkbox"/>	✓			
sample 4		<input type="checkbox"/>				
sample 5		<input type="checkbox"/>				
sample 6		<input type="checkbox"/>				
sample 7		<input type="checkbox"/>				
sample 8		<input type="checkbox"/>				
sample 9		<input type="checkbox"/>				
sample 10		<input type="checkbox"/>				
sample 11		<input type="checkbox"/>				
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

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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

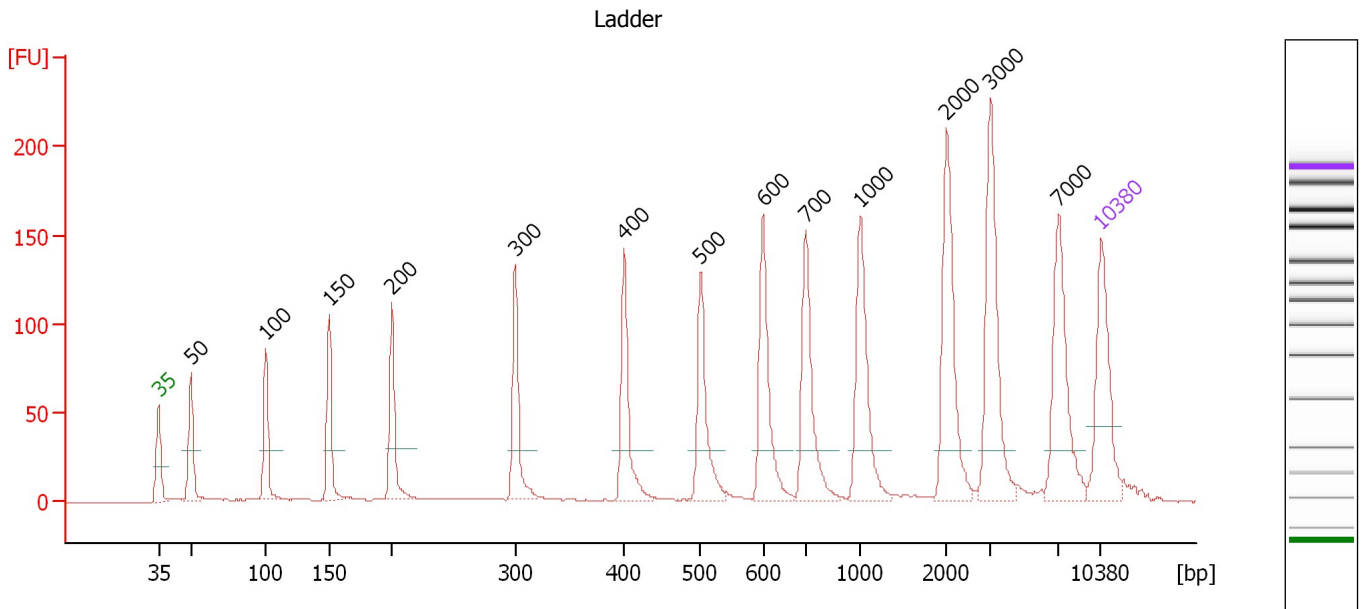
Ladder

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

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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

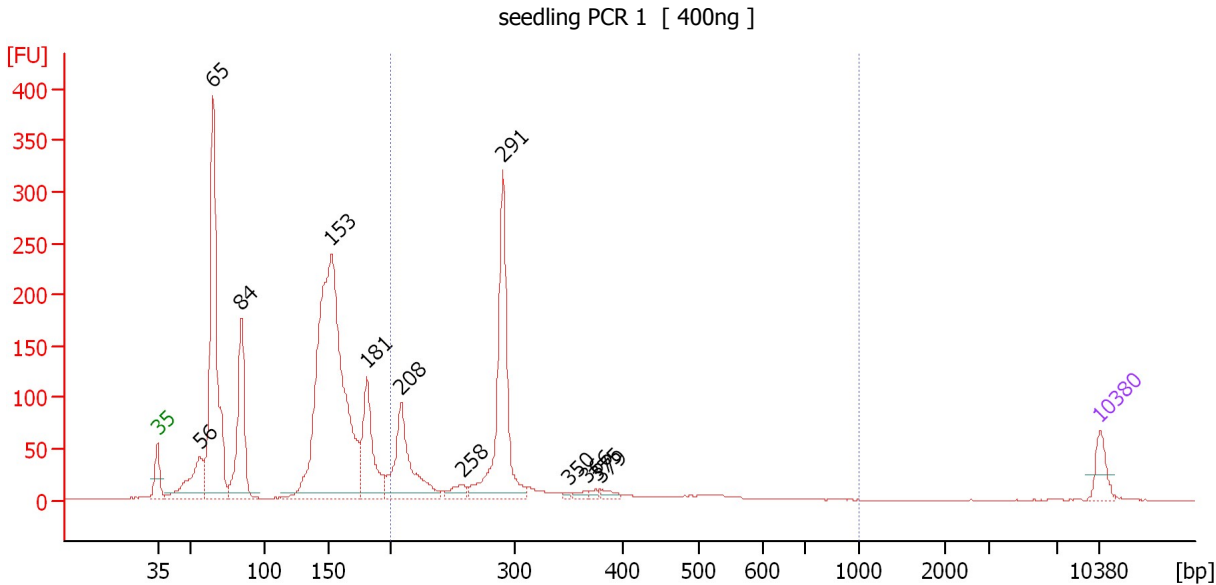
Peak table for Ladder

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	50.95
4	150	150.00	1,515.2	Ladder Peak	55.69
5	200	150.00	1,136.4	Ladder Peak	60.33
6	300	150.00	757.6	Ladder Peak	69.49
7	400	150.00	568.2	Ladder Peak	77.58
8	500	150.00	454.5	Ladder Peak	83.25
9	600	150.00	378.8	Ladder Peak	87.89
10	700	150.00	324.7	Ladder Peak	91.07
11	1,000	150.00	227.3	Ladder Peak	95.13
12	2,000	150.00	113.6	Ladder Peak	101.52
13	3,000	150.00	75.8	Ladder Peak	104.78
14	7,000	150.00	32.5	Ladder Peak	109.87
15	10,380	75.00	10.9	Upper Marker	113.00

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Electropherogram Summary Continued ...



Overall Results for sample 1 : seedling PCR 1

Number of peaks found: 12 Corr. Area 1: 808.5
 Noise: 0.2

Peak table for sample 1 : seedling PCR 1

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	56	319.42	8,571.5		46.13
3	65	1,469.68	34,014.4		47.13
4	84	636.65	11,444.1		49.21
5	153	2,560.30	25,434.0		55.92
6	181	454.72	3,810.2		58.55
7	208	521.23	3,792.0		61.09
8	258	64.60	379.7		65.62
9	291	872.69	4,544.7		68.66
10	350	9.55	41.3		73.56
11	366	20.87	86.4		74.83
12	375	15.49	62.6		75.55
13	379	28.47	113.7		75.91
14	10,380	75.00	10.9	Upper Marker	113.00

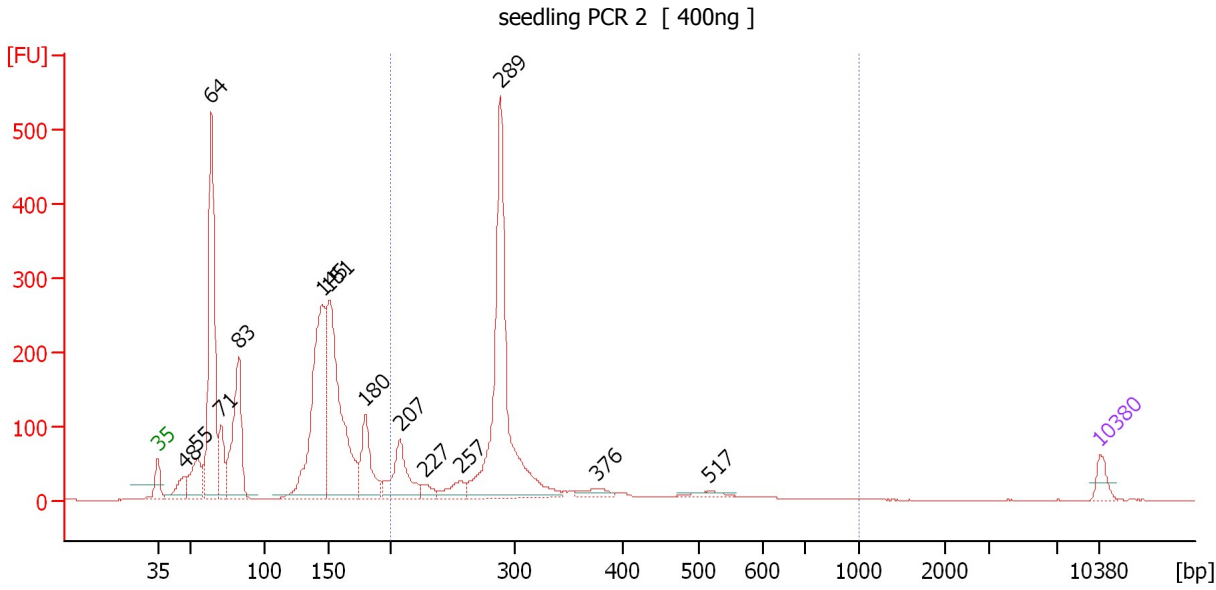
Region table for sample 1 : seedling PCR 1

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	300	808.5	9,569.0	1,769.50	27	31.2

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Electropherogram Summary Continued ...



Overall Results for sample 2 : seedling PCR 2

Number of peaks found: 14 Corr. Area 1: 1,372.8
 Noise: 0.3

Peak table for sample 2 : seedling PCR 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	48	208.51	6,647.4		45.01
3	55	365.94	10,141.2		45.93
4	64	1,866.84	44,073.2		46.98
5	71	309.11	6,616.0		47.72
6	83	962.41	17,616.4		49.04
7	145	1,686.58	17,610.1		55.22
8	151	1,809.56	18,165.8		55.77
9	180	462.89	3,907.1		58.43
10	207	457.52	3,345.5		60.99
11	227	73.39	489.4		62.82
12	257	132.68	783.1		65.52
13	289	2,139.52	11,209.5		68.50
14	376	79.47	320.2		75.64
15	517	51.69	151.6		84.02
16	10,380	75.00	10.9	Upper Marker	113.00

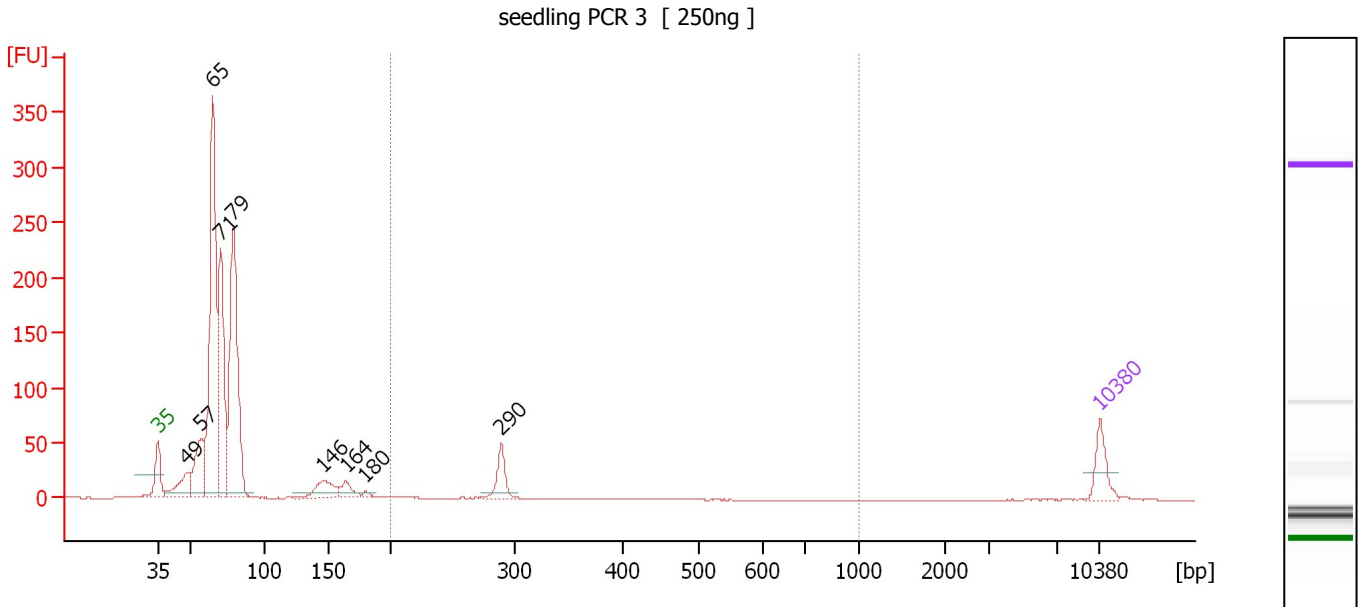
Region table for sample 2 : seedling PCR 2

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	312	1,372.8	17,353.7	3,335.46	34	31.0

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Electropherogram Summary Continued ...



Overall Results for sample 3 : seedling PCR 3

Number of peaks found: 9 Corr. Area 1: 68.9
 Noise: 0.2

Peak table for sample 3 : seedling PCR 3

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	49	135.84	4,224.8		45.21
3	57	256.96	6,798.5		46.22
4	65	1,160.82	26,996.5		47.09
5	71	584.21	12,548.0		47.69
6	79	938.88	18,043.3		48.61
7	146	119.25	1,236.7		55.32
8	164	60.27	555.6		57.02
9	180	15.77	133.0		58.44
10	290	106.34	556.0		68.55
11	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 3 : seedling PCR 3

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	285	68.9	765.6	143.25	5	10.6

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Gel Image

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Invalid Samples

Sample 4 has not been run, no results available.

Sample 5 has not been run, no results available.

Sample 6 has not been run, no results available.

Sample 7 has not been run, no results available.

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.

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Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 4)		Instrument	Run		1/5/2017 4:25:33 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2017-01-05\2017-01-05_003.xad)		Instrument	Run		1/5/2017 4:07:12 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		1/5/2017 4:07:12 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		1/5/2017 4:07:12 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		1/5/2017 4:07:12 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		1/5/2017 4:07:12 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		1/5/2017 4:07:12 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		1/5/2017 4:07:12 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1