

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-11-18\2016-11-18_005.xad

Created: 11/18/2016 2:15:26 PM
Modified: 11/18/2016 2:49:14 PM

Electrophoresis File Run Summary

Instrument Information:

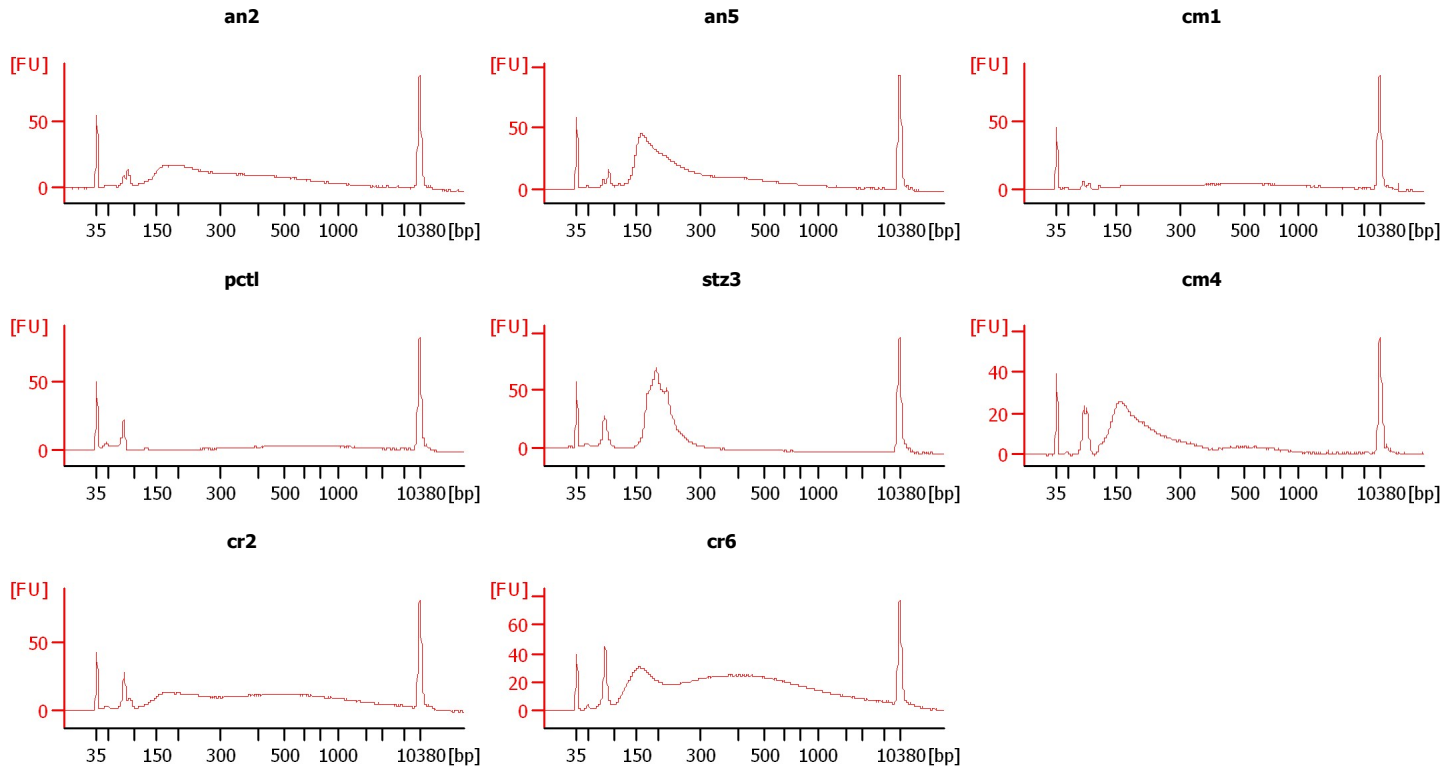
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:



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Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
an2		<input type="checkbox"/>	✓			
an5		<input type="checkbox"/>	✓			
cm1		<input type="checkbox"/>	✓			
pctl		<input type="checkbox"/>	✓			
stz3		<input type="checkbox"/>	✓			
cm4		<input type="checkbox"/>	✓			
cr2		<input type="checkbox"/>	✓			
cr6		<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

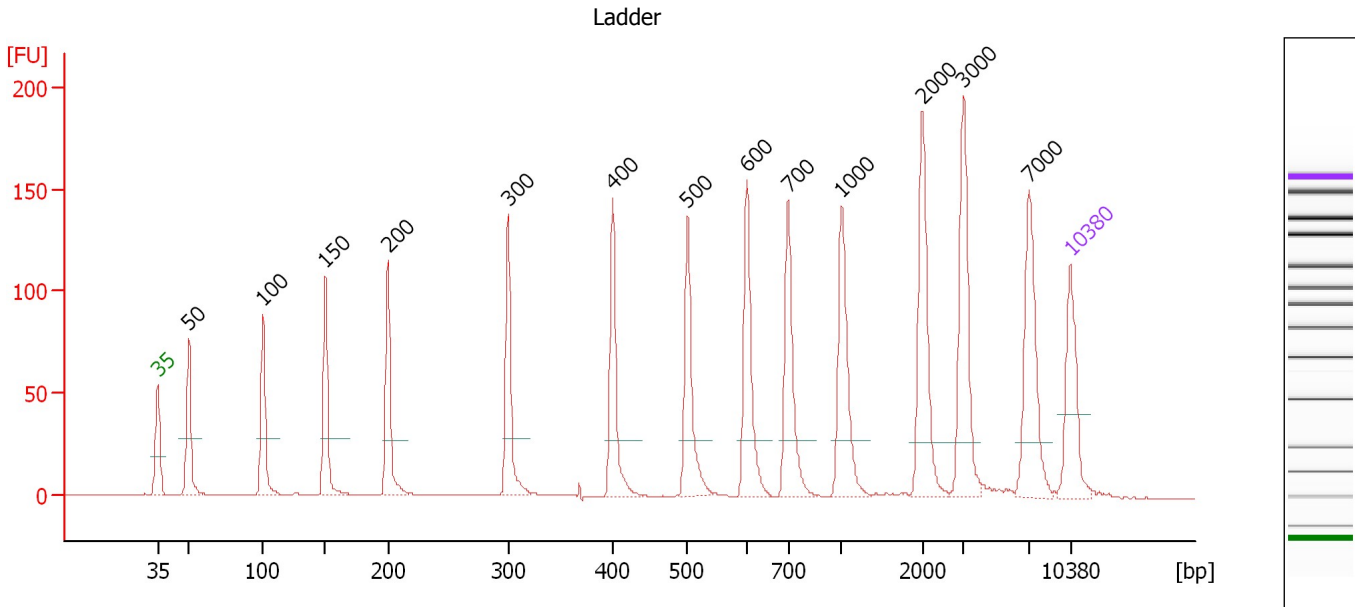
Ladder

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

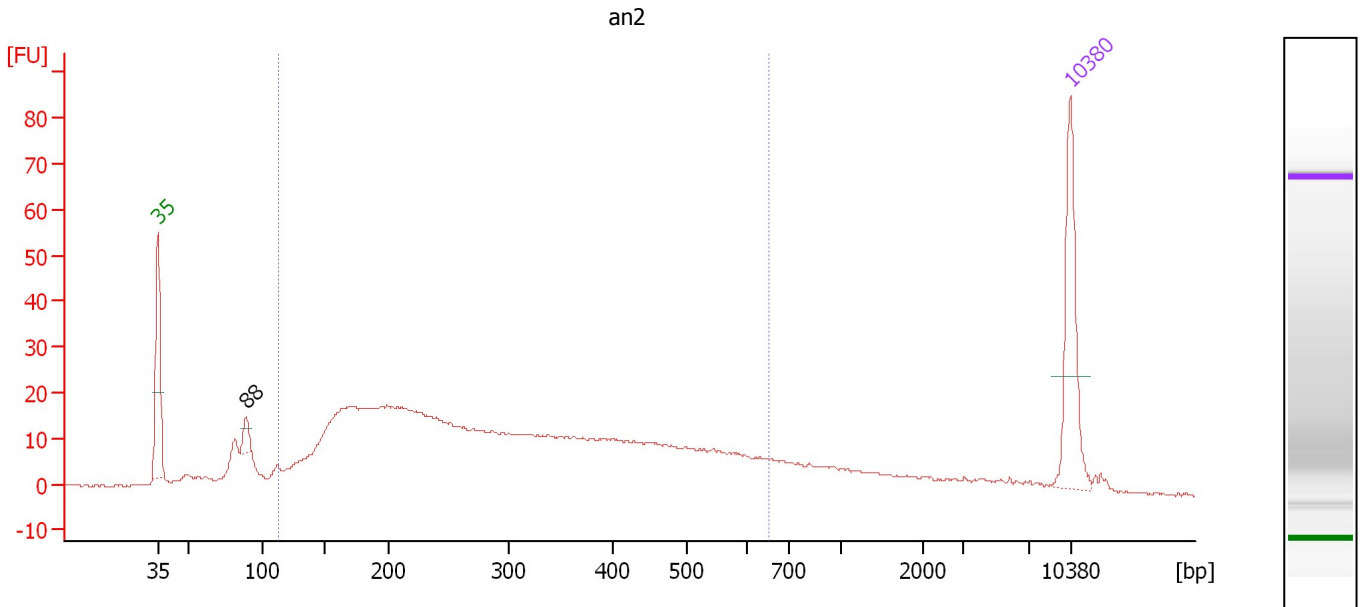
Peak table for Ladder

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	45.39
3	100	150.00	2,272.7	Ladder Peak	51.09
4	150	150.00	1,515.2	Ladder Peak	55.88
5	200	150.00	1,136.4	Ladder Peak	60.67
6	300	150.00	757.6	Ladder Peak	69.88
7	400	150.00	568.2	Ladder Peak	77.92
8	500	150.00	454.5	Ladder Peak	83.63
9	600	150.00	378.8	Ladder Peak	88.21
10	700	150.00	324.7	Ladder Peak	91.36
11	1,000	150.00	227.3	Ladder Peak	95.49
12	2,000	150.00	113.6	Ladder Peak	101.65
13	3,000	150.00	75.8	Ladder Peak	104.80
14	7,000	150.00	32.5	Ladder Peak	109.84
15	10,380	75.00	10.9	Upper Marker	113.00

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Electropherogram Summary Continued ...



Overall Results for sample 1 : an2

Number of peaks found: 1 Corr. Area 1: 654.1
 Noise: 0.2

Peak table for sample 1 : an2

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	88	16.65	285.4		49.77
3	10,380	75.00	10.9	Upper Marker	113.00

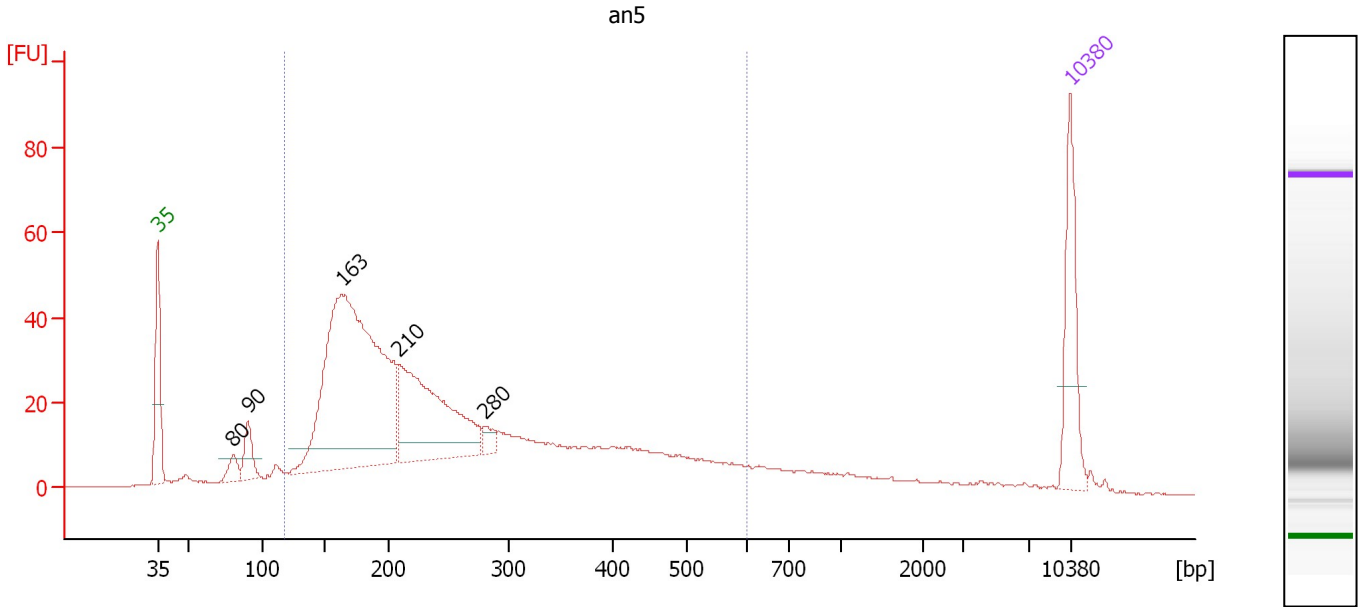
Region table for sample 1 : an2

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
112	653	314	654.1	7,316.8	1,160.58	79	42.3

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Electropherogram Summary Continued ...



Overall Results for sample 2 : an5

Number of peaks found: 5 Corr. Area 1: 899.3
 Noise: 0.1

Peak table for sample 2 : an5

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	80	20.67	391.3		48.82
3	90	40.65	687.2		49.91
4	163	665.87	6,185.2		57.14
5	210	258.47	1,868.2		61.55
6	280	14.34	77.6		68.05
7	10,380	75.00	10.9	Upper Marker	113.00

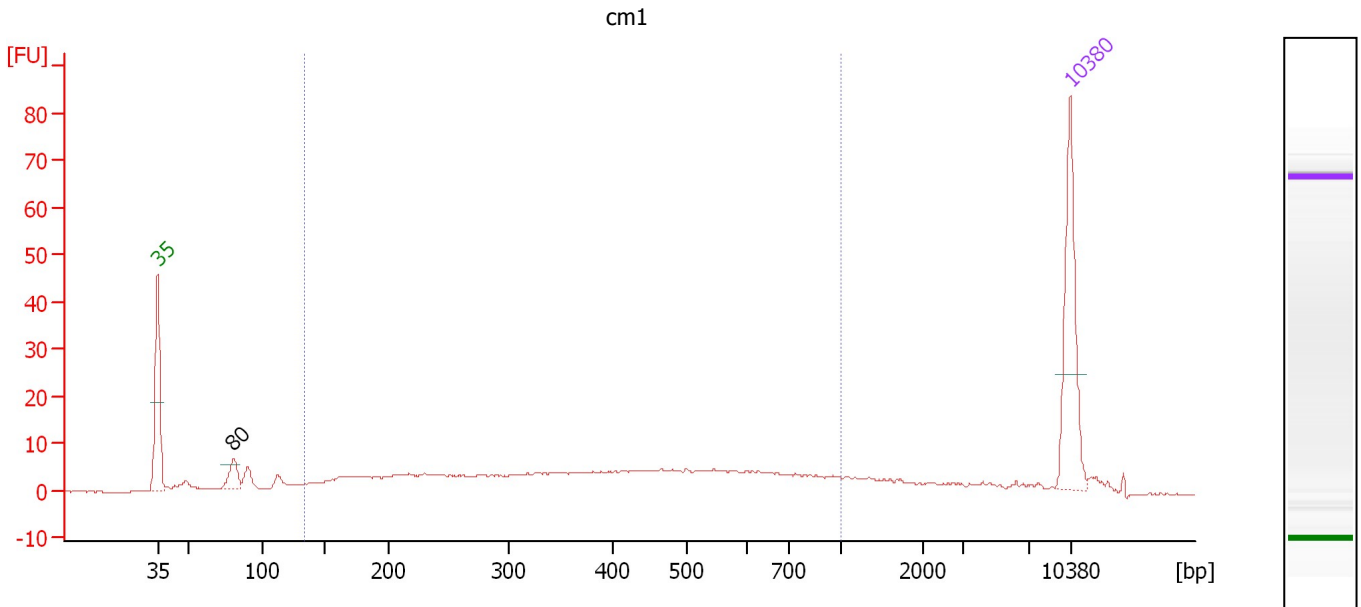
Region table for sample 2 : an5

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
117	600	267	899.3	10,945.6	1,553.33	83	42.6

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Electropherogram Summary Continued ...



Overall Results for sample 3 : cm1

Number of peaks found: 1 Corr. Area 1: 223.8
 Noise: 0.2

Peak table for sample 3 : cm1

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	80	22.58	427.4		48.82
3	10,380	75.00	10.9	Upper Marker	113.00

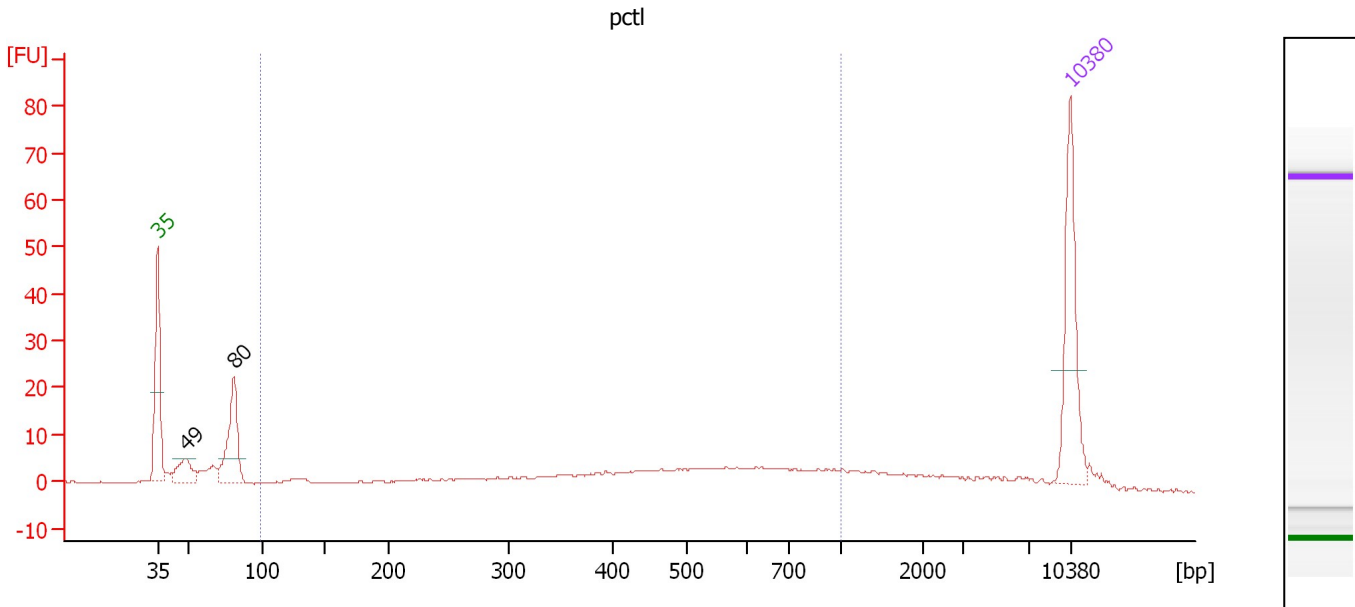
Region table for sample 3 : cm1

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
133	1,000	430	223.8	1,944.3	391.44	70	46.4

Assay Class: High Sensitivity DNA Assay
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Electropherogram Summary Continued ...



Overall Results for sample 4 : pctI

Number of peaks found: 2 Corr. Area 1: 143.5
 Noise: 0.2

Peak table for sample 4 : pctI

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	33.61	1,048.7		45.16
3	80	87.88	1,659.1		48.84
4	10,380	75.00	10.9	Upper Marker	113.00

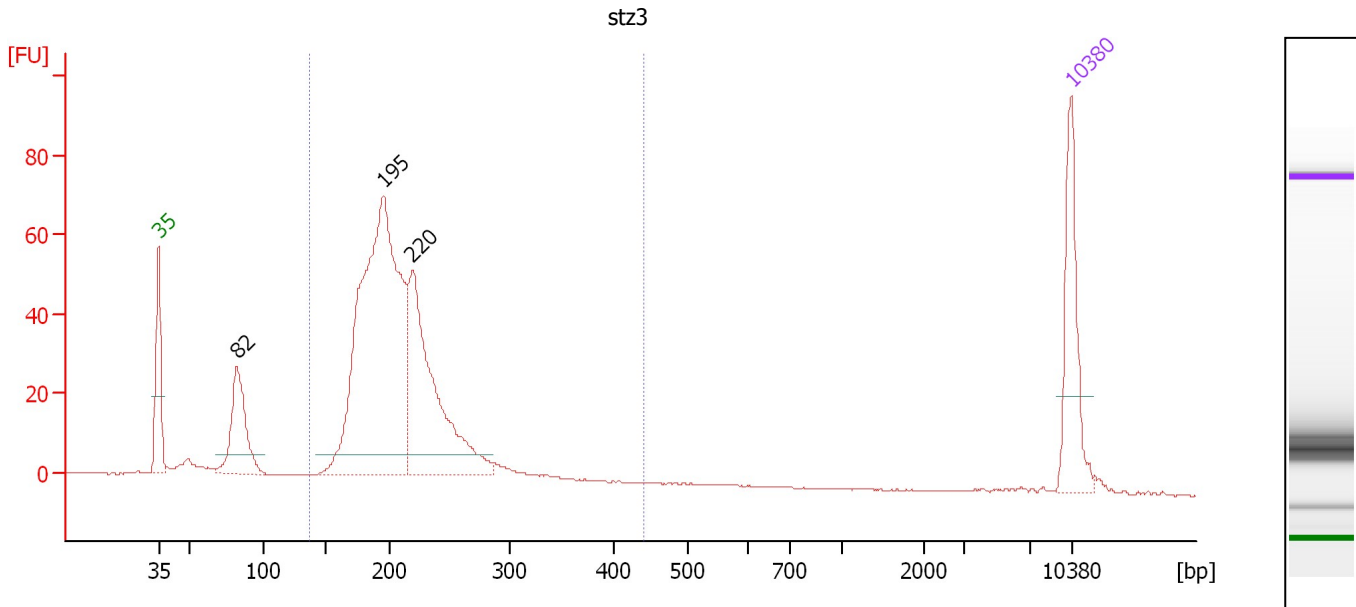
Region table for sample 4 : pctI

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
99	1,000	503	143.5	1,005.6	236.86	51	40.7

Assay Class: High Sensitivity DNA Assay
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Electropherogram Summary Continued ...



Overall Results for sample 5 : stz3

Number of peaks found: 3 Corr. Area 1: 637.3
 Noise: 0.1

Peak table for sample 5 : stz3

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	82	116.97	2,171.4		49.00
3	195	593.70	4,603.7		60.22
4	220	281.07	1,938.3		62.48
5	10,380	75.00	10.9	Upper Marker	113.00

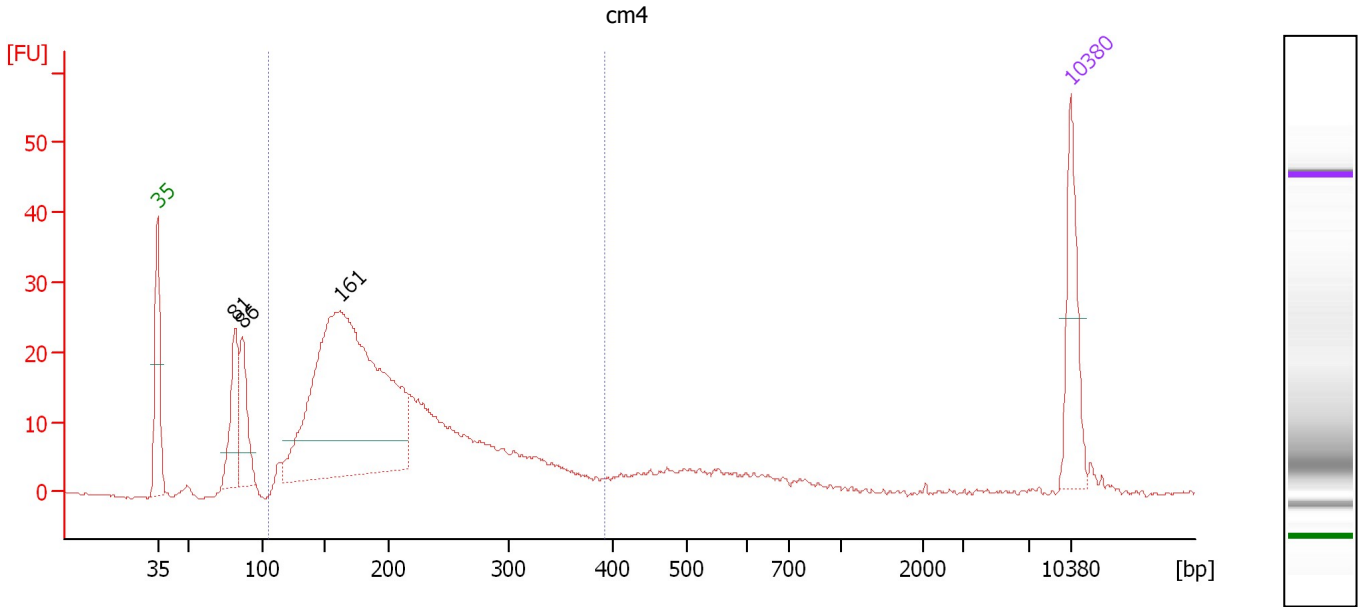
Region table for sample 5 : stz3

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
136	440	211	637.3	6,939.2	939.87	85	17.1

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Electropherogram Summary Continued ...



Overall Results for sample 6 : cm4

Number of peaks found: 3 Corr. Area 1: 420.7
 Noise: 0.2

Peak table for sample 6 : cm4

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	81	109.28	2,046.4		48.92
3	86	96.05	1,695.0		49.48
4	161	781.45	7,332.9		56.98
5	10,380	75.00	10.9	Upper Marker	113.00

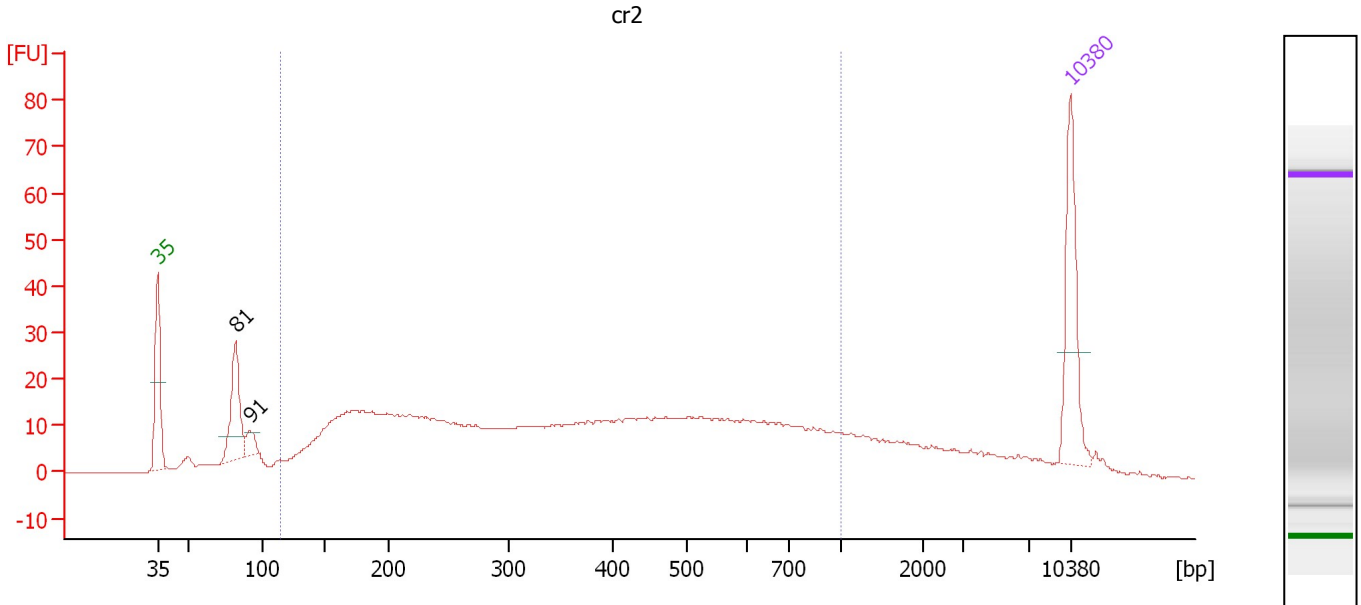
Region table for sample 6 : cm4

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
105	392	207	420.7	10,315.1	1,267.28	79	29.7

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Electropherogram Summary Continued ...



Overall Results for sample 7 : cr2

Number of peaks found: 2 Corr. Area 1: 680.3
 Noise: 0.2

Peak table for sample 7 : cr2

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	81	91.40	1,699.7		48.98
3	91	18.58	309.1		50.07
4	10,380	75.00	10.9	Upper Marker	113.00

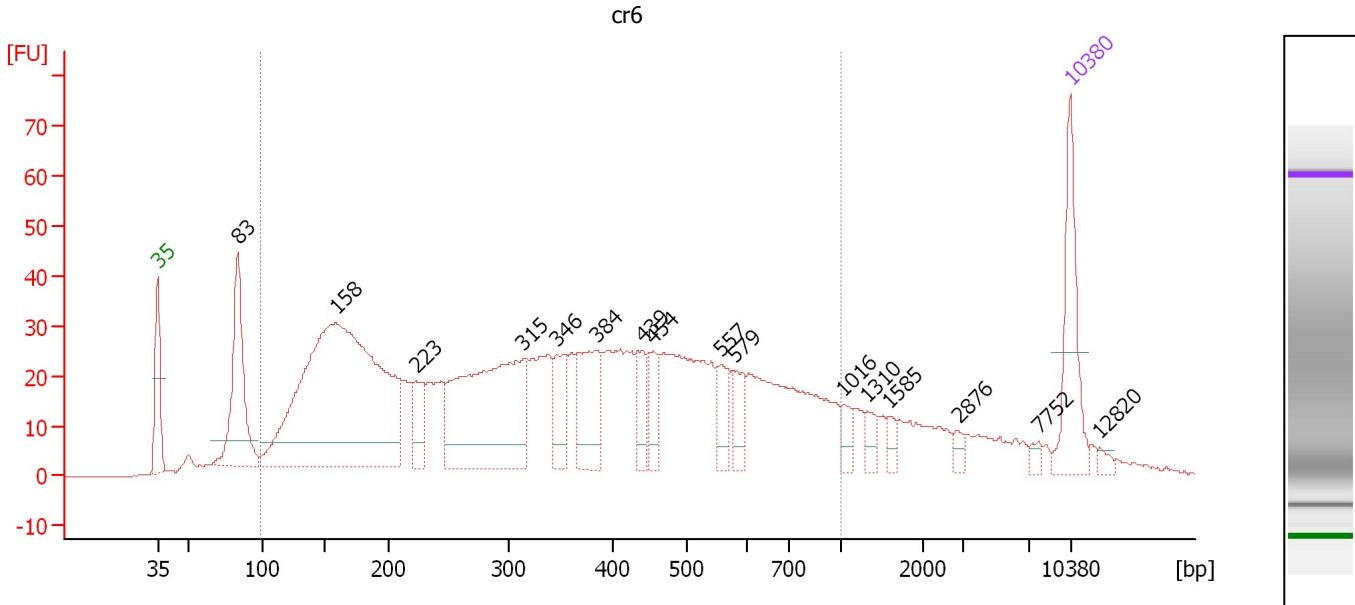
Region table for sample 7 : cr2

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
114	1,000	407	680.3	6,890.5	1,251.69	72	50.5

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Electropherogram Summary Continued ...



Overall Results for sample 8 : cr6

Number of peaks found: 16 Corr. Area 1: 1,326.3
 Noise: 0.1

Peak table for sample 8 : cr6

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	83	181.30	3,316.8		49.14
3	158	719.69	6,889.2		56.67
4	223	43.01	291.9		62.81
5	315	282.85	1,362.6		71.05
6	346	48.33	211.3		73.62
7	384	89.44	352.7		76.66
8	439	35.87	123.7		80.16
9	454	32.88	109.8		80.98
10	557	33.81	92.0		86.24
11	579	25.69	67.3		87.23
12	1,016	15.07	22.5		95.59
13	1,310	14.59	16.9		97.40
14	1,585	9.86	9.4		99.09
15	2,876	7.24	3.8		104.41
16	7,752	6.15	1.2		110.55
17	10,380	75.00	10.9	Upper Marker	113.00
18	12,820	0.00	0.0		115.28

Region table for sample 8 : cr6

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
99	1,000	383	1,326.3	14,174.7	2,353.79	78	51.8

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Gel Image

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Invalid Samples

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: 9)		Instrument	Run		11/18/2016 2:48:07 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\2016-11-18\2016-11-18_005.xad)		Instrument	Run		11/18/2016 2:15:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		11/18/2016 2:15:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		11/18/2016 2:15:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		11/18/2016 2:15:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		11/18/2016 2:15:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		11/18/2016 2:15:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		11/18/2016 2:15:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1