

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...nd Settings\Bioanalyzer\2016-01-19\2016-01-19_004_BC383.xad

Created: 1/19/2016 3:34:33 PM
Modified: 1/19/2016 4:22:54 PM

Electrophoresis File Run Summary

Instrument Information:

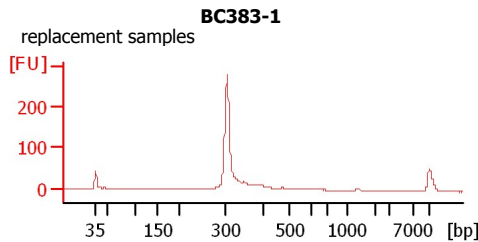
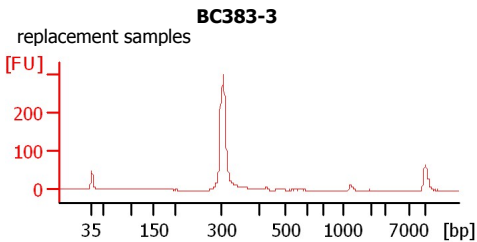
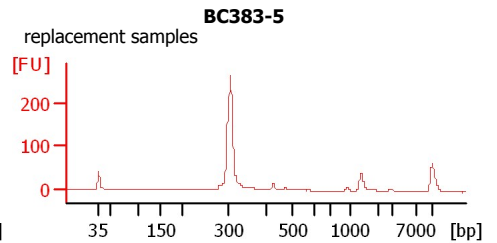
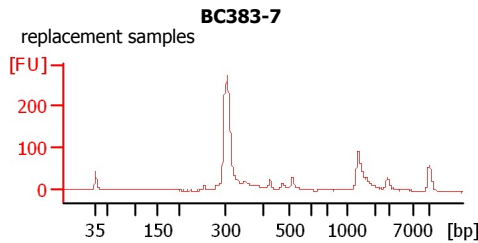
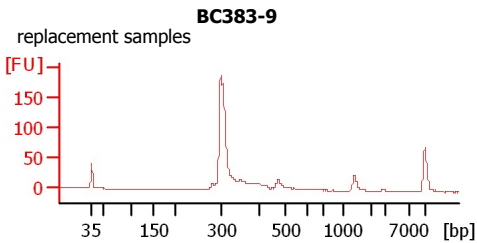
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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
BC383-9	replacement samples	<input type="checkbox"/>	✓			
BC383-7	replacement samples	<input type="checkbox"/>	✓			
BC383-5	replacement samples	<input type="checkbox"/>	✓			
BC383-3	replacement samples	<input type="checkbox"/>	✓			
BC383-1	replacement samples	<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

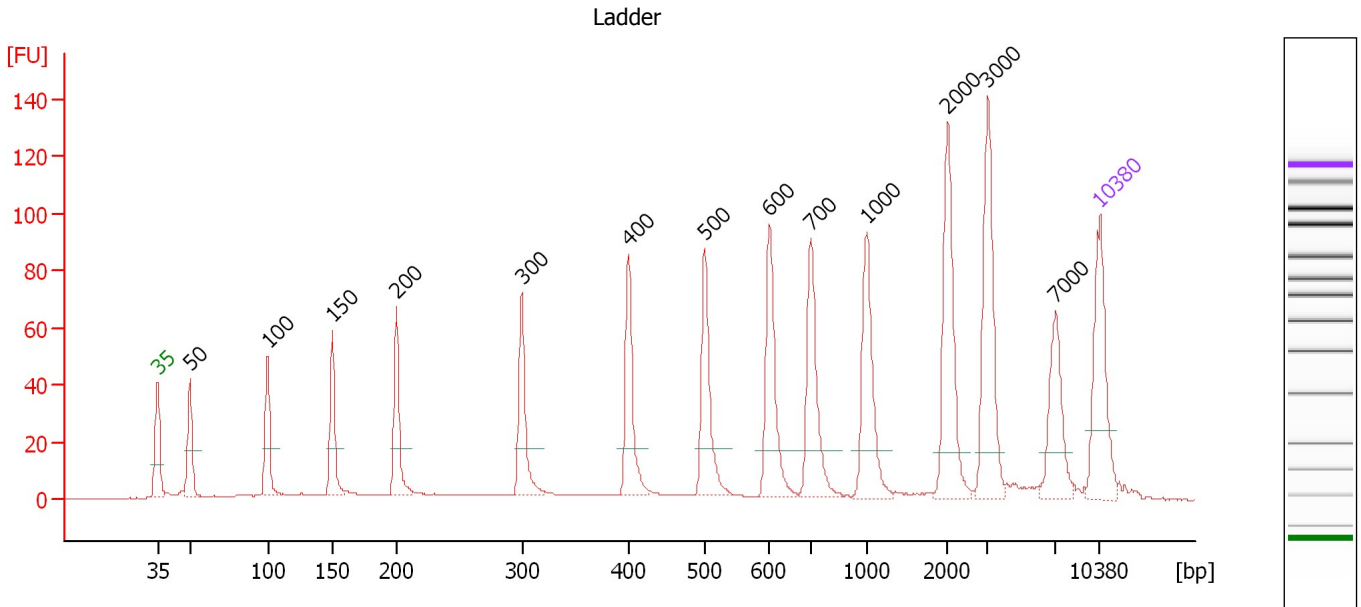
Ladder

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

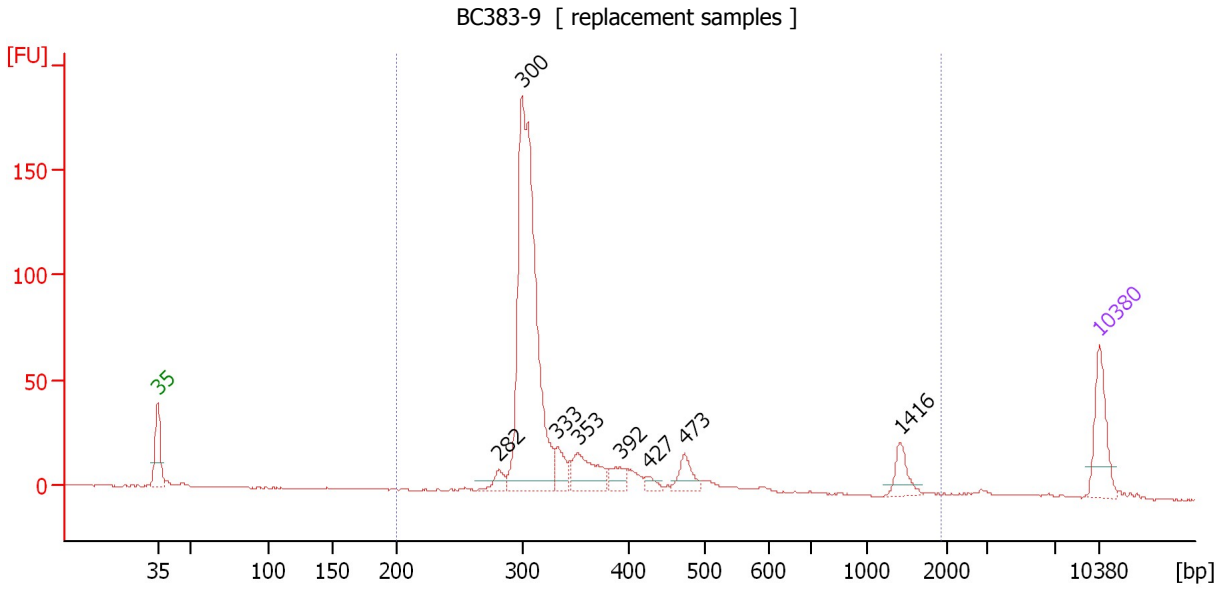
Peak table for Ladder

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	45.42
3	100	150.00	2,272.7	Ladder Peak	51.18
4	150	150.00	1,515.2	Ladder Peak	55.98
5	200	150.00	1,136.4	Ladder Peak	60.73
6	300	150.00	757.6	Ladder Peak	70.05
7	400	150.00	568.2	Ladder Peak	78.00
8	500	150.00	454.5	Ladder Peak	83.67
9	600	150.00	378.8	Ladder Peak	88.46
10	700	150.00	324.7	Ladder Peak	91.52
11	1,000	150.00	227.3	Ladder Peak	95.68
12	2,000	150.00	113.6	Ladder Peak	101.67
13	3,000	150.00	75.8	Ladder Peak	104.68
14	7,000	150.00	32.5	Ladder Peak	109.71
15	10,380	75.00	10.9	Upper Marker	113.00

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Electropherogram Summary Continued ...



Overall Results for sample 2 : BC383-9

Number of peaks found: 8 Corr. Area 1: 657.6
 Noise: 0.2

Peak table for sample 2 : BC383-9

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	282	26.74	143.9		68.33
3	300	812.00	4,096.0		70.08
4	333	45.83	208.3		72.70
5	353	93.46	401.7		74.22
6	392	34.91	135.0		77.36
7	427	15.76	55.9		79.52
8	473	42.95	137.5		82.14
9	1,416	36.90	39.5		98.17
10	10,380	75.00	10.9	Upper Marker	113.00

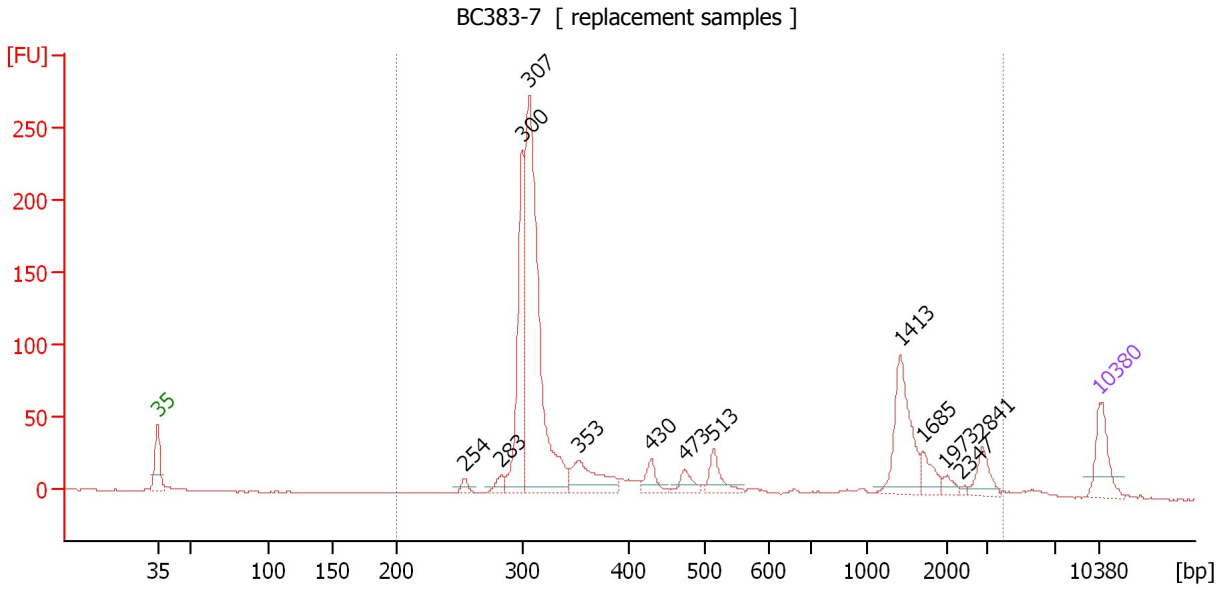
Region table for sample 2 : BC383-9

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,931	410	657.6	5,333.1	1,189.03	97	67.5

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Electropherogram Summary Continued ...



Overall Results for sample 3 : BC383-7

Number of peaks found: 13 Corr. Area 1: 1,175.4
 Noise: 0.2

Peak table for sample 3 : BC383-7

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	254	18.94	112.8		65.81
3	283	26.26	140.6		68.46
4	300	375.04	1,891.1		70.09
5	307	848.18	4,187.1		70.60
6	353	126.35	542.1		74.28
7	430	43.21	152.4		79.68
8	473	34.90	111.8		82.14
9	513	55.56	164.1		84.28
10	1,413	173.30	185.9		98.15
11	1,685	41.65	37.5		99.78
12	1,973	15.90	12.2		101.50
13	2,347	4.33	2.8		102.71
14	2,841	43.87	23.4		104.20
15	10,380	75.00	10.9	Upper Marker	113.00

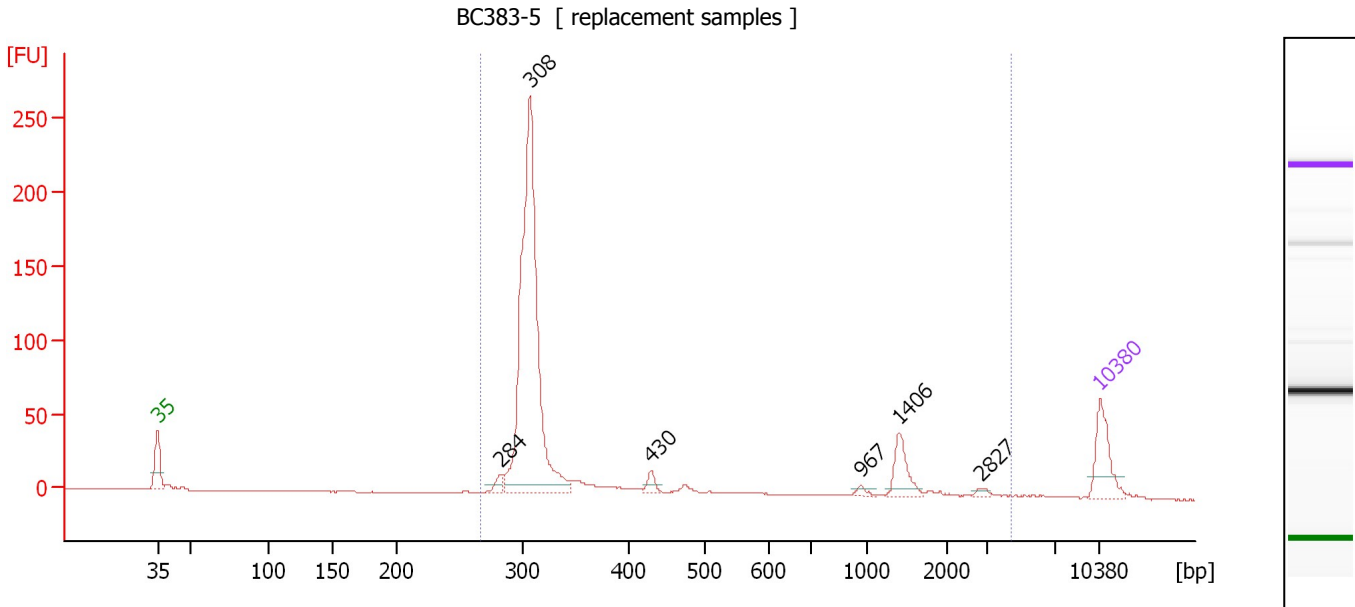
Region table for sample 3 : BC383-7

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	3,921	740	1,175.4	7,752.4	1,941.25	98	96.7

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Electropherogram Summary Continued ...



Overall Results for sample 4 : BC383-5

Number of peaks found: 6 Corr. Area 1: 692.5
 Noise: 0.1

Peak table for sample 4 : BC383-5

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	284	19.94	106.5		68.52
3	308	900.23	4,435.0		70.65
4	430	17.62	62.1		79.68
5	967	7.56	11.8		95.23
6	1,406	58.27	62.8		98.11
7	2,827	7.07	3.8		104.16
8	10,380	75.00	10.9	Upper Marker	113.00

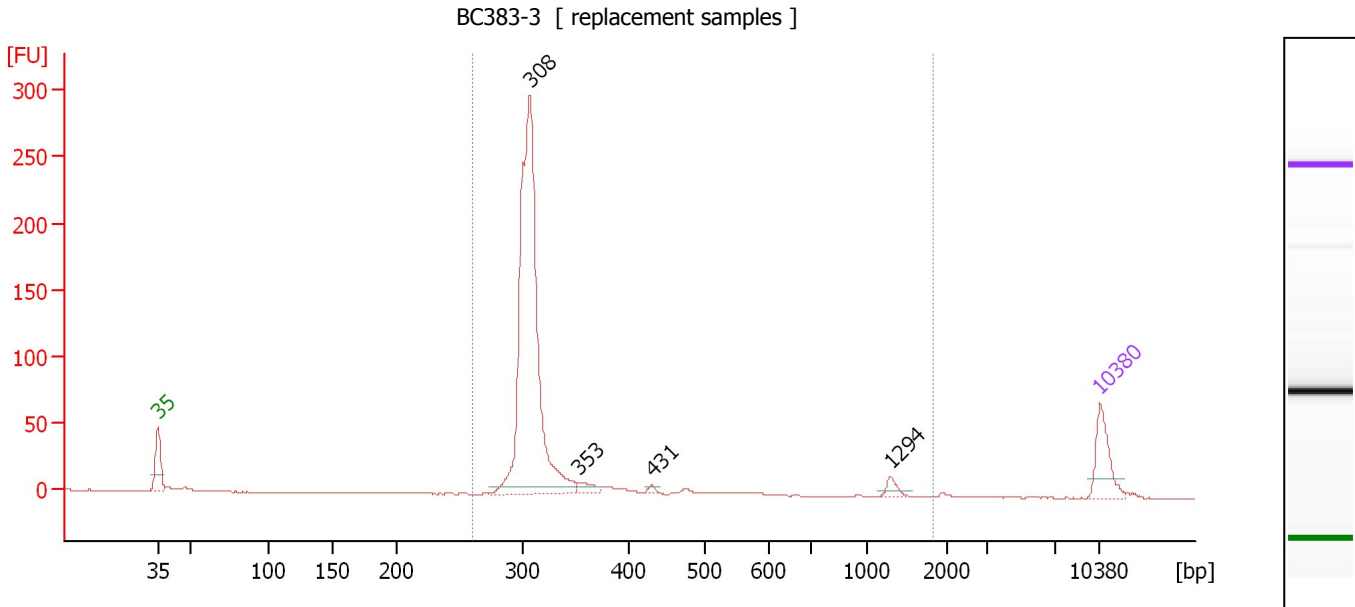
Region table for sample 4 : BC383-5

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
267	4,376	513	692.5	5,044.9	1,124.81	98	100.0

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Electropherogram Summary Continued ...



Overall Results for sample 5 : BC383-3

Number of peaks found: 4 Corr. Area 1: 709.7
 Noise: 0.1

Peak table for sample 5 : BC383-3

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	308	1,073.55	5,287.9		70.65
3	353	25.39	109.0		74.26
4	431	6.89	24.2		79.73
5	1,294	18.45	21.6		97.44
6	10,380	75.00	10.9	Upper Marker	113.00

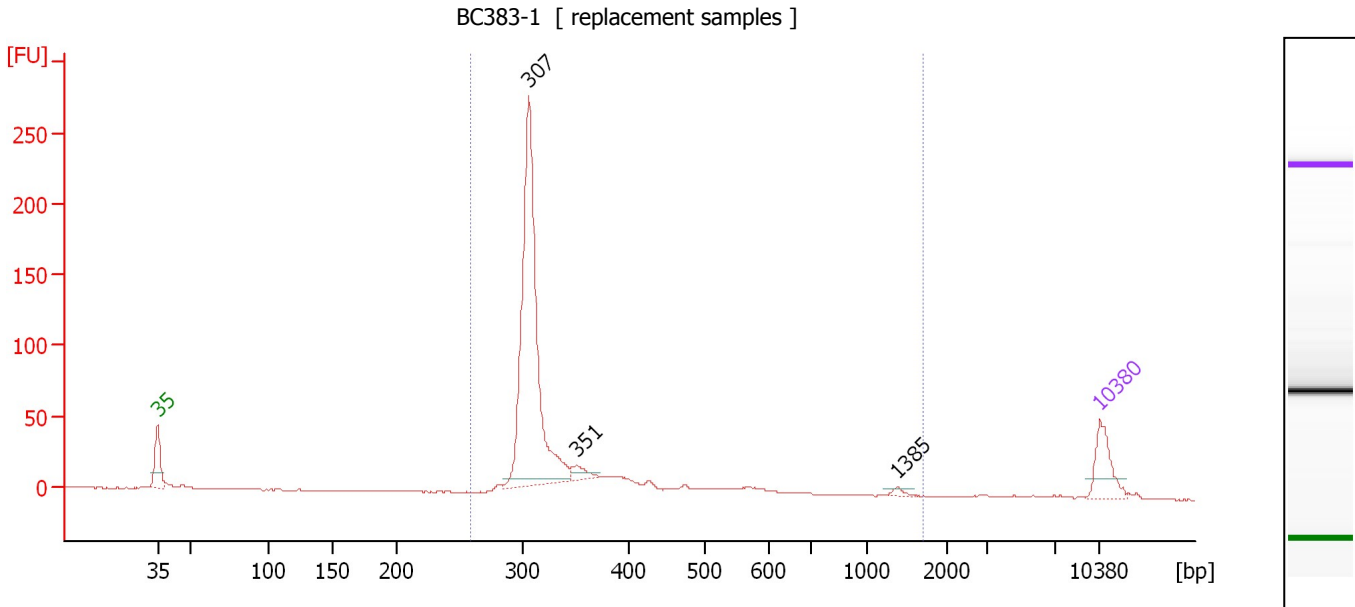
Region table for sample 5 : BC383-3

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
260	1,824	348	709.7	5,595.3	1,177.19	97	49.2

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Electropherogram Summary Continued ...



Overall Results for sample 6 : BC383-1

Number of peaks found: 3 Corr. Area 1: 689.3
 Noise: 0.2

Peak table for sample 6 : BC383-1

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	307	935.64	4,620.4		70.59
3	351	27.87	120.2		74.14
4	1,385	8.68	9.5		97.99
5	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 6 : BC383-1

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
259	1,703	351	689.3	5,785.6	1,253.88	98	38.6

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Gel Image

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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: 12)		Instrument	Run		1/19/2016 4:15:53 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2016-01-19\2016-01-19_004.xad)		Instrument	Run		1/19/2016 3:34:38 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		1/19/2016 3:34:38 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		1/19/2016 3:34:38 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		1/19/2016 3:34:38 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		1/19/2016 3:34:38 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		1/19/2016 3:34:38 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		1/19/2016 3:34:38 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1