

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-28\2012-03-28_002.xad

Created: 3/28/2012 1:23:18 PM
Modified: 3/28/2012 2:33:45 PM

Electrophoresis File Run Summary

Instrument Information:

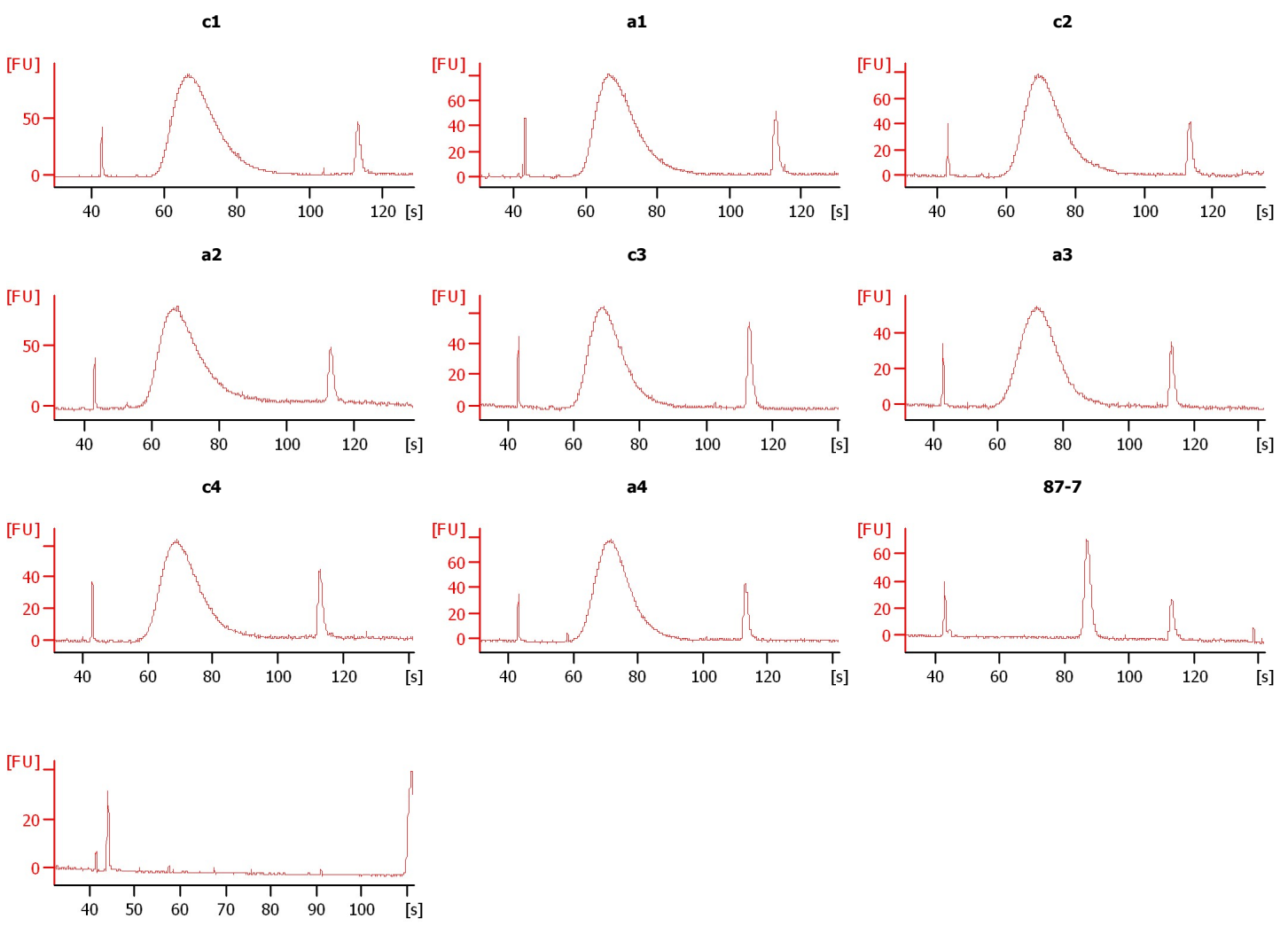
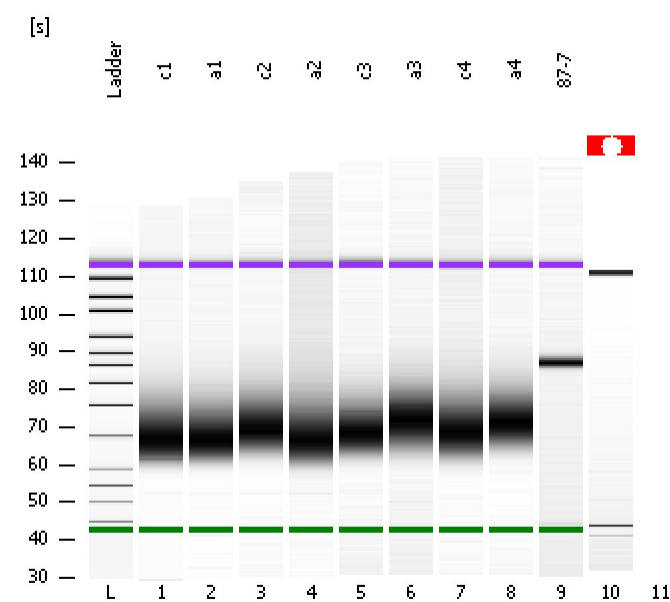
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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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 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-28\2012-03-28_002.xad

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Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
c1		<input type="checkbox"/>	✓			
a1		<input type="checkbox"/>	✓			
c2		<input type="checkbox"/>	✓			
a2		<input type="checkbox"/>	✓			
c3		<input type="checkbox"/>	✓			
a3		<input type="checkbox"/>	✓			
c4		<input type="checkbox"/>	✓			
a4		<input type="checkbox"/>	✓			
87-7		<input type="checkbox"/>	✓			
		<input type="checkbox"/>	✗			
		<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

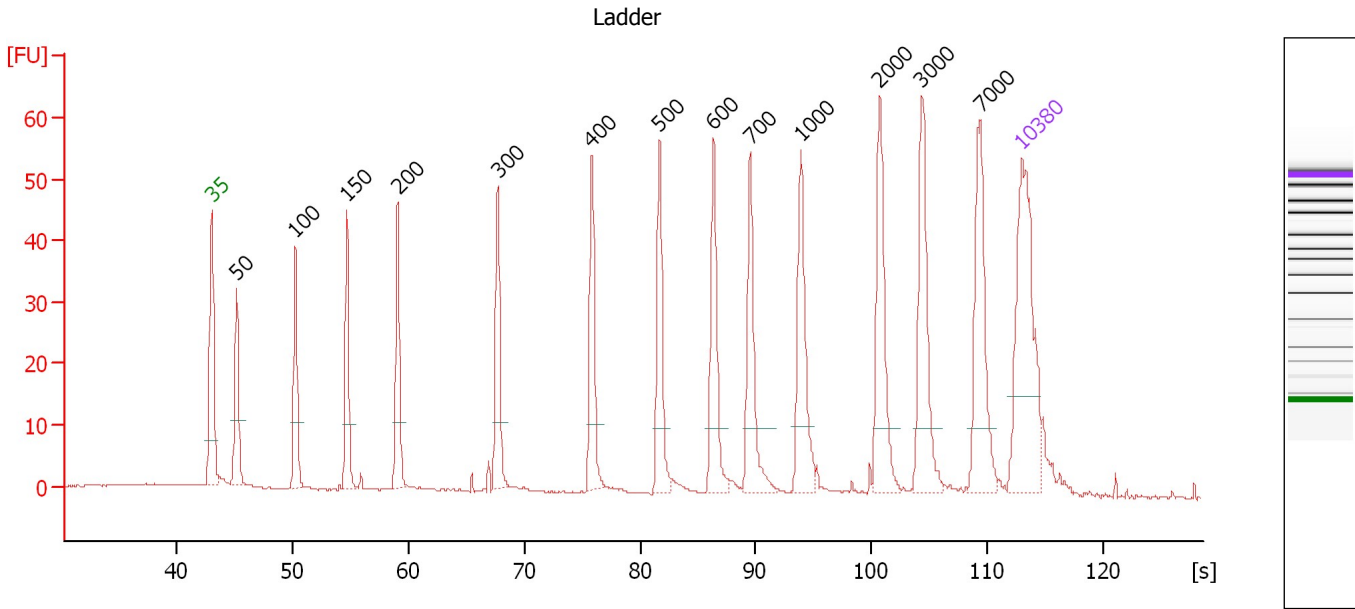
Ladder

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: 0.1

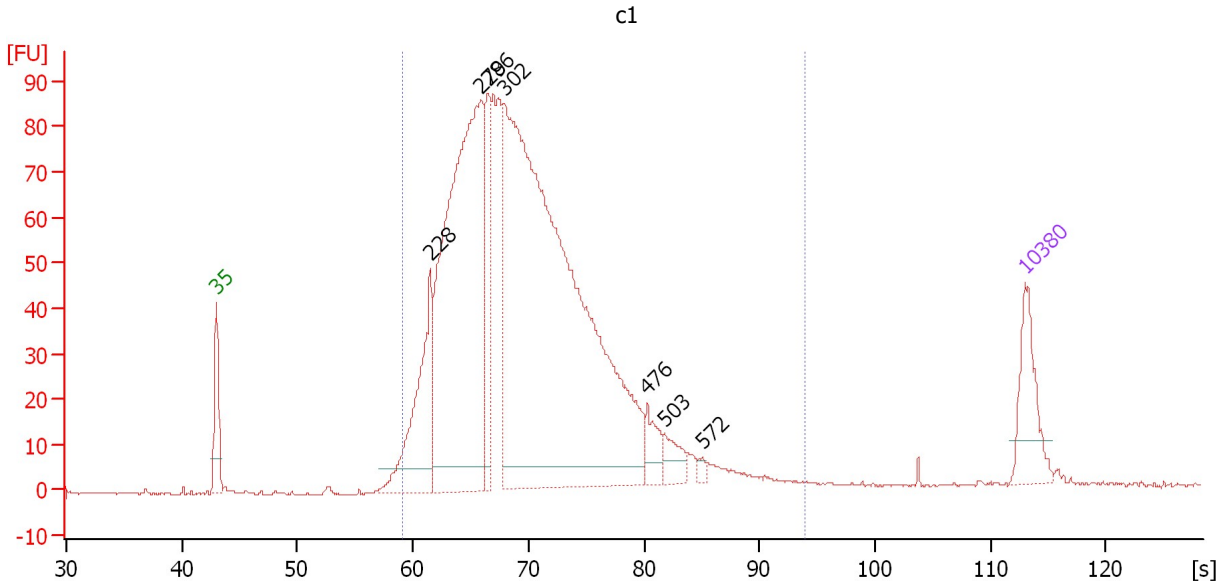
Peak table for Ladder

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

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Electropherogram Summary Continued ...



Overall Results for sample 1 : c1

Number of peaks found: 7 Corr. Area 1: 1,705.6
 Noise: 0.2

Peak table for sample 1 : c1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	228	255.05	1,696.1	
3	279	934.40	5,074.0	
4	286	154.38	817.8	
5	302	1,676.01	8,399.6	
6	476	45.91	146.2	
7	503	36.87	111.1	
8	572	7.16	19.0	
9	10,380	75.00	10.9	Upper Marker

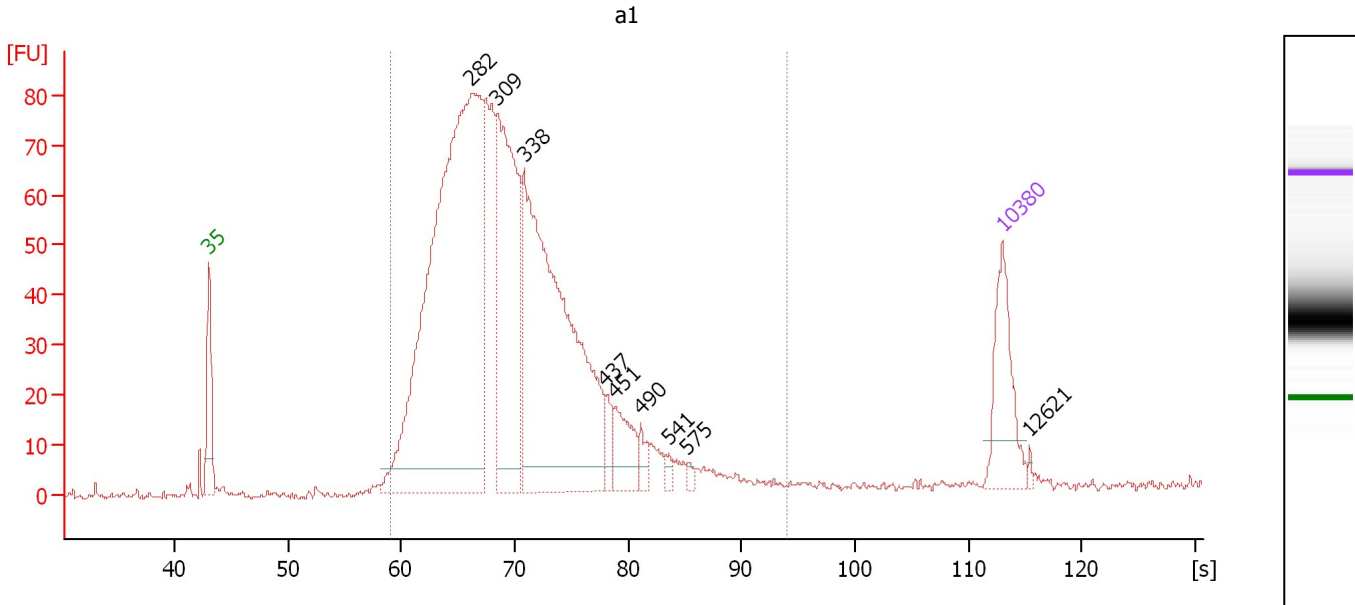
Region table for sample 1 : c1

From [s]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [s]	Corr. Area	% of Total	Size distribution in CV [%]	Color
59.10	16,051.1	330	3,248.09	93.93	1,705.6	99	26.7	Blue

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Electropherogram Summary Continued ...



Overall Results for sample 2 : a1

Number of peaks found: 9 Corr. Area 1: 1,464.4
 Noise: 0.7

Peak table for sample 2 : a1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	282	1,037.79	5,578.7	
3	309	363.01	1,781.6	
4	338	624.13	2,798.7	
5	437	21.03	72.9	
6	451	58.11	195.0	
7	490	19.56	60.4	
8	541	7.48	21.0	
9	575	5.40	14.2	
10	10,380	75.00	10.9	Upper Marker
11	12,621	0.00	0.0	

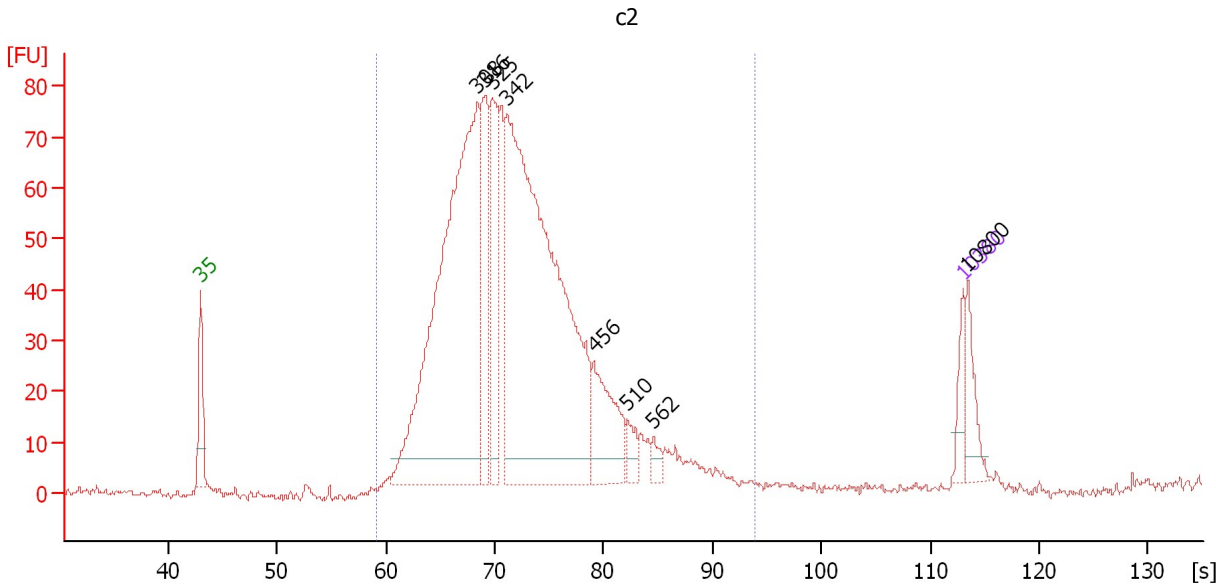
Region table for sample 2 : a1

From [s]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [s]	Corr. Area	% of Total	Size distribution in CV [%]	Color
59.10	11,599.6	327	2,342.13	93.93	1,464.4	99	24.9	Blue

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 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-28\2012-03-28_002.xad

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Electropherogram Summary Continued ...



Overall Results for sample 3 : c2

Number of peaks found: 8 Corr. Area 1: 1,377.8
 Noise: 0.7

Peak table for sample 3 : c2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	308	2,400.63	11,794.9	
3	316	454.50	2,177.3	
4	325	469.93	2,192.2	
5	342	2,714.93	12,013.5	
6	456	333.76	1,109.2	
7	510	71.57	212.5	
8	562	39.19	105.7	
9	10,380	75.00	10.9	Upper Marker
10	10,800	0.00	0.0	

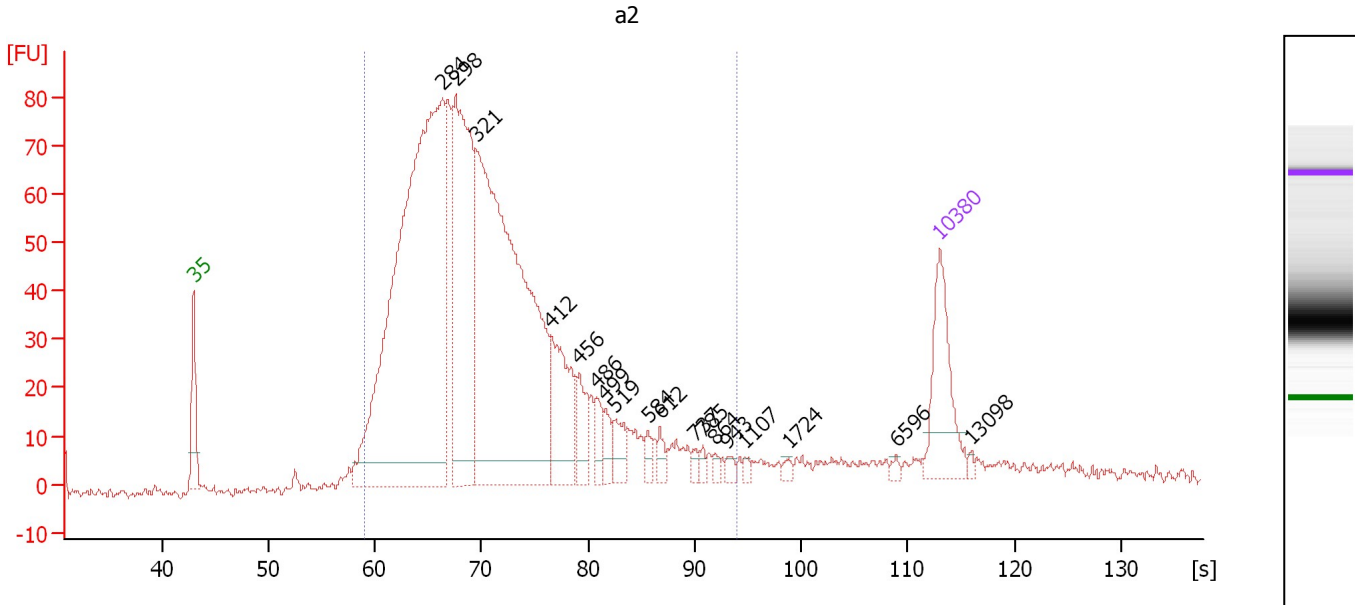
Region table for sample 3 : c2

From [s]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [s]	Corr. Area	% of Total	Size distribution in CV [%]	Color
59.10	32,418.3	359	7,180.23	93.93	1,377.8	97	24.8	Blue

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Electropherogram Summary Continued ...



Overall Results for sample 4 : a2

Number of peaks found: 18 Corr. Area 1: 1,656.7
 Noise: 1.1

Peak table for sample 4 : a2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	284	941.85	5,027.1	
3	298	348.16	1,767.4	
4	321	737.24	3,476.1	
5	412	106.51	391.4	
6	456	41.61	138.2	
7	486	23.88	74.5	
8	499	19.78	60.0	
9	519	26.68	77.9	
10	584	10.17	26.4	
11	612	11.96	29.6	
12	737	6.41	13.2	
13	785	5.58	10.8	
14	864	5.12	9.0	
15	943	6.12	9.8	
16	1,107	4.48	6.1	
17	1,724	4.58	4.0	
18	6,596	3.89	0.9	
19	10,380	75.00	10.9	Upper Marker
20	13,098	0.00	0.0	

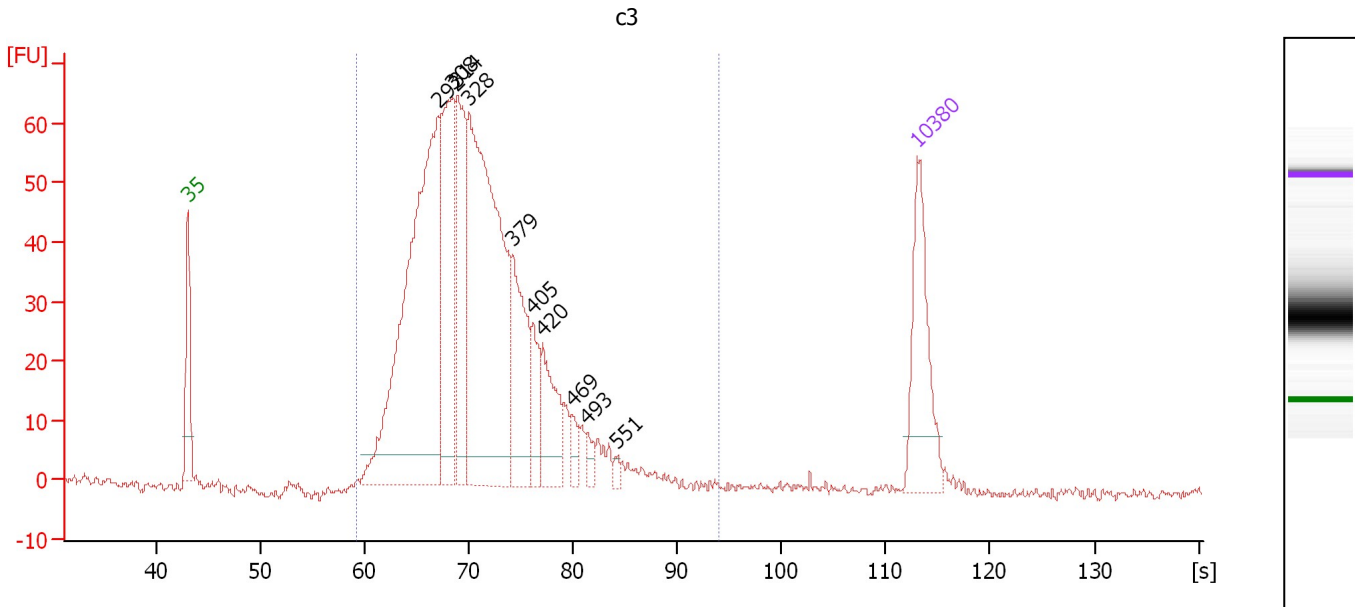
Region table for sample 4 : a2

From [s]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [s]	Corr. Area	% of Total	Size distribution in CV [%]	Color
59.10	12,120.3	347	2,490.27	93.93	1,656.7	93	34.0	Blue

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Electropherogram Summary Continued ...



Overall Results for sample 5 : c3

Number of peaks found: 10 Corr. Area 1: 1,053.1
 Noise: 0.6

Peak table for sample 5 : c3

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	292	473.36	2,452.5	
3	308	170.40	837.0	
4	314	128.61	620.2	
5	328	403.95	1,868.7	
6	379	104.88	419.3	
7	405	35.38	132.3	
8	420	60.39	217.8	
9	469	14.24	46.0	
10	493	10.12	31.1	
11	551	5.18	14.2	
12	10,380	75.00	10.9	Upper Marker

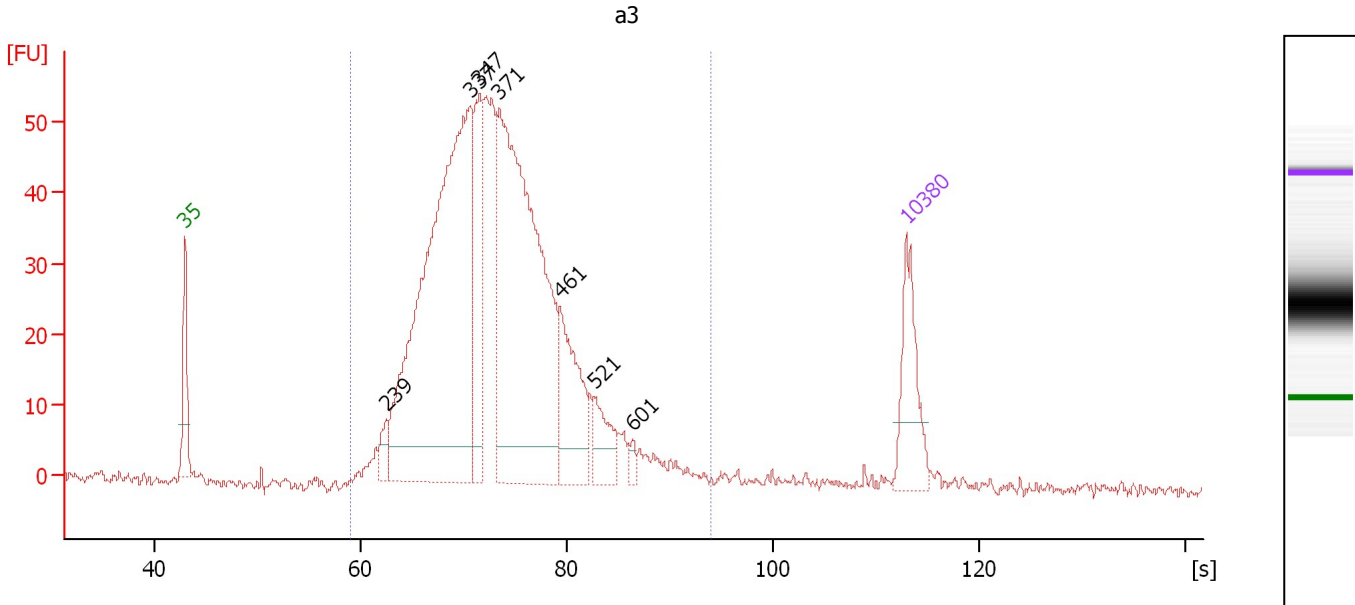
Region table for sample 5 : c3

From [s]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/µl]	To [s]	Corr. Area	% of Total	Size distribution in CV [%]	Color
59.10	6,768.1	338	1,435.23	93.93	1,053.1	100	20.8	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-28\2012-03-28_002.xad

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Electropherogram Summary Continued ...



Overall Results for sample 6 : a3

Number of peaks found: 7 Corr. Area 1: 1,010.0
 Noise: 0.9

Peak table for sample 6 : a3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	239	26.80	169.6	
3	337	822.91	3,699.8	
4	347	141.45	617.1	
5	371	667.35	2,725.2	
6	461	128.85	423.9	
7	521	51.62	150.1	
8	601	8.74	22.0	
9	10,380	75.00	10.9	Upper Marker

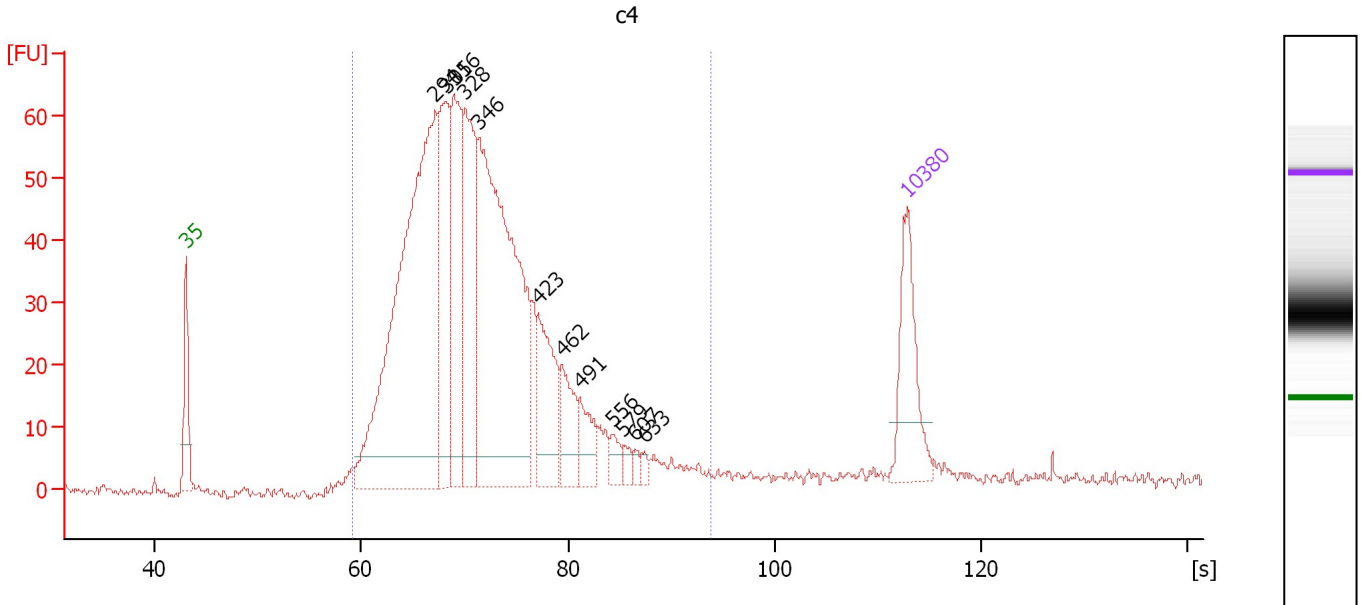
Region table for sample 6 : a3

From [s]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [s]	Corr. Area	% of Total	Size distribution in CV [%]	Color
59.10	9,438.1	378	2,194.87	93.93	1,010.0	97	25.1	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-28\2012-03-28_002.xad

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Electropherogram Summary Continued ...



Overall Results for sample 7 : c4

Number of peaks found: 12 Corr. Area 1: 1,204.3
 Noise: 0.8

Peak table for sample 7 : c4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	294	656.72	3,389.9	
3	305	175.13	869.1	
4	316	179.35	861.1	
5	328	193.46	892.6	
6	346	508.43	2,228.1	
7	423	97.65	350.1	
8	462	54.53	178.9	
9	491	40.81	125.9	
10	556	15.67	42.7	
11	579	9.25	24.2	
12	607	6.76	16.9	
13	633	5.75	13.8	
14	10,380	75.00	10.9	Upper Marker

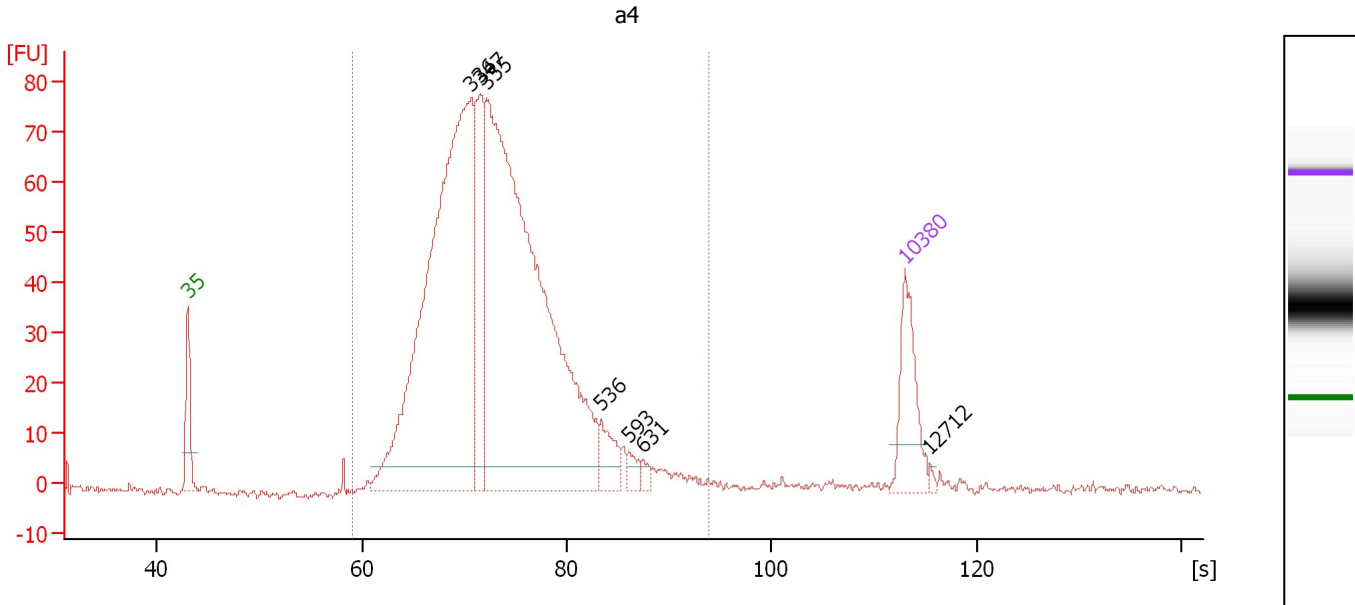
Region table for sample 7 : c4

From [s]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [s]	Corr. Area	% of Total	Size distribution in CV [%]	Color
59.10	9,469.4	354	2,035.91	93.93	1,204.3	97	28.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-28\2012-03-28_002.xad

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Electropherogram Summary Continued ...



Overall Results for sample 8 : a4

Number of peaks found: 7 Corr. Area 1: 1,320.1
 Noise: 0.6

Peak table for sample 8 : a4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	336	932.30	4,206.3	
3	347	170.97	746.9	
4	355	1,077.02	4,602.5	
5	536	41.05	116.0	
6	593	14.20	36.3	
7	631	8.62	20.7	
8	10,380	75.00	10.9	Upper Marker
9	12,712	0.00	0.0	

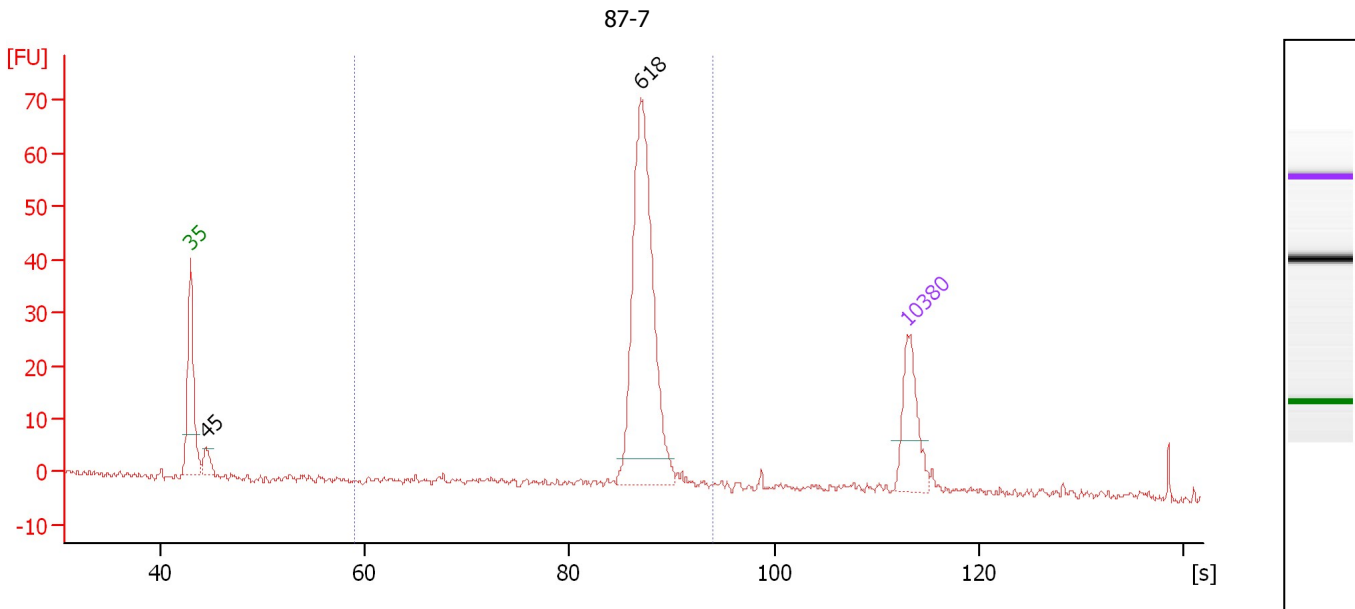
Region table for sample 8 : a4

From [s]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [s]	Corr. Area	% of Total	Size distribution in CV [%]	Color
59.10	9,697.6	374	2,252.36	93.93	1,320.1	99	23.6	Blue

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Electropherogram Summary Continued ...



Overall Results for sample 9 : 87-7

Number of peaks found: 2 Corr. Area 1: 199.2
 Noise: 0.7

Peak table for sample 9 : 87-7

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	45	29.63	995.5	
3	618	418.74	1,026.7	
4	10,380	75.00	10.9	Upper Marker

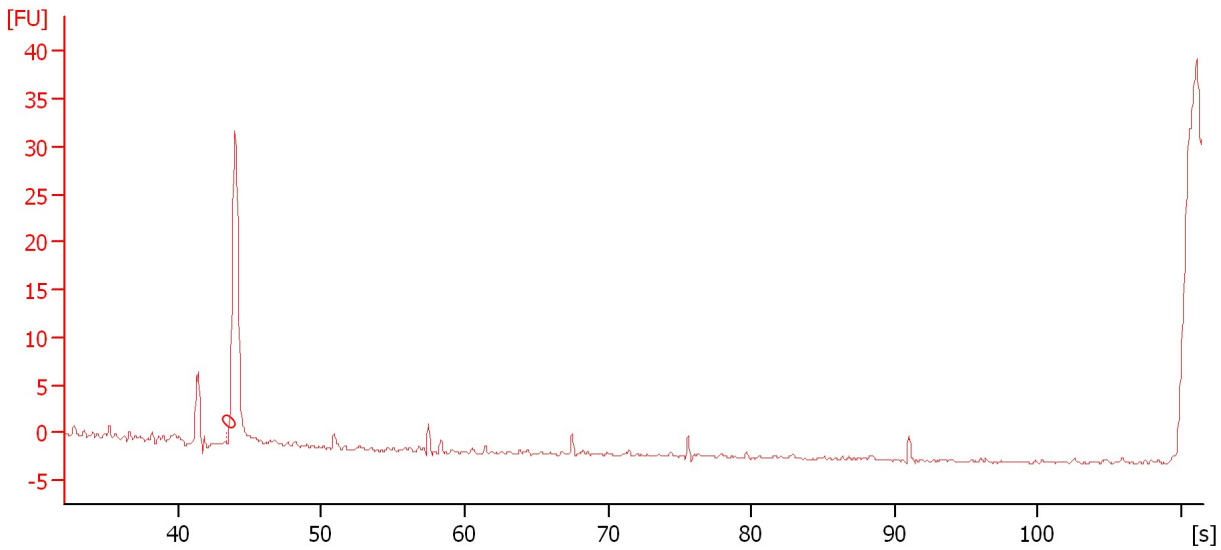
Region table for sample 9 : 87-7

From [s]	Molarity [pmol/l]	Average Size [bp]	Conc. [pg/μl]	To [s]	Corr. Area	% of Total	Size distribution in CV [%]	Color
59.10	1,114.0	626	450.41	93.93	199.2	87	9.9	Blue

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Electropherogram Summary Continued ...



Setpoint Deviations for sample 10 : -

End Analysis Time Range [s]: 111.45

Overall Results for sample 10 : -

Number of peaks found: 0 Noise: 0.2

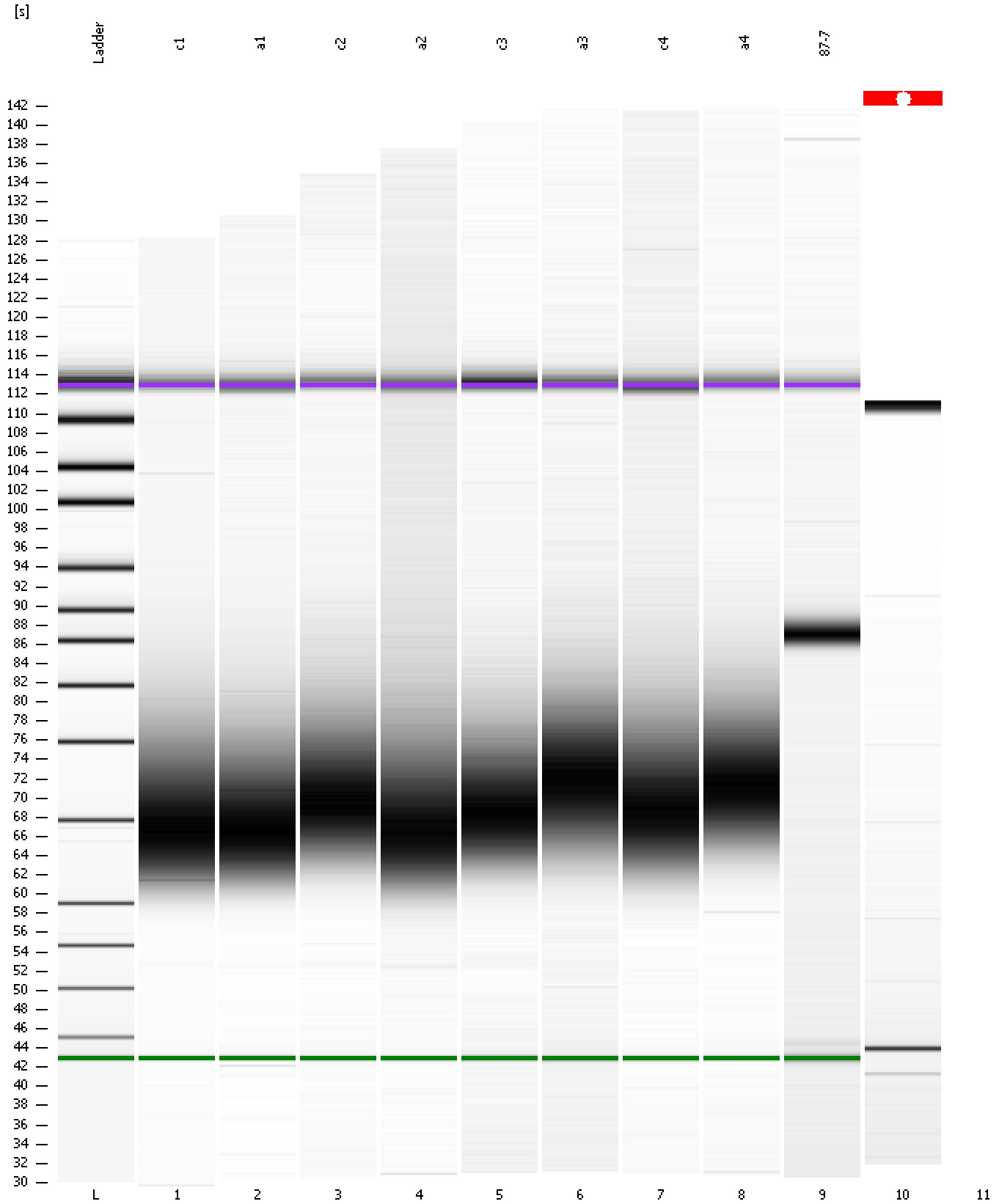
Peak table for sample 10 : -

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	0	0.00	0.0	
2	0	0.00	0.0	
3	0	0.00	0.0	excluded peak

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Gel Image



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Invalid Samples

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 aborted on port 1		Instrument	Run		3/28/2012 2:01:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2012-03-28\2012-03-28_002.xad)		Instrument	Run		3/28/2012 1:23:24 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		3/28/2012 1:23:24 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		3/28/2012 1:23:24 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		3/28/2012 1:23:24 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		3/28/2012 1:23:24 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		3/28/2012 1:23:24 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		3/28/2012 1:23:24 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1