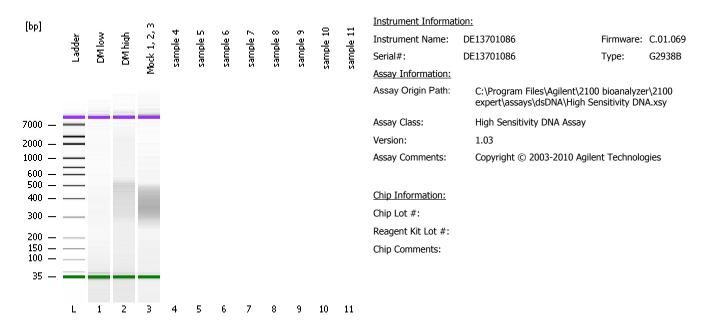
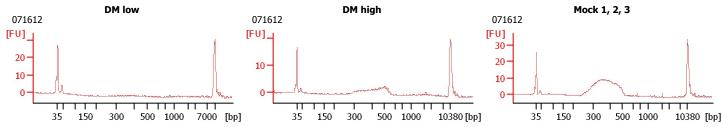
2012-07-16_001.xad Page 1 of 10

Assay Class: High Sensitivity DNA Assay Created: 7/16/2012 3:09:25 PM Data Path: C:\...ents and Settings\Bioanalyzer\2012-07-16\2012-07-16_001.xad Created: 7/16/2012 3:28:30 PM

Electrophoresis File Run Summary





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High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2012-07-16\2012-07-16_001.xad 7/16/2012 3:09:25 PM 7/16/2012 3:28:30 PM Assay Class: Created: Data Path: Modified: **Electrophoresis File Run Summary (Chip Summary) Sample Name** Sample Comment Rest. Digest Status Observation **Result Label Result Color** DM low 071612 DM high 071612 Mock 1, 2, 3 071612 sample 4 sample 5 sample 6

Chip Lot # Reagent Kit Lot #

Chip Comments:

sample 7 sample 8 sample 9 sample 10 sample 11 Ladder

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2012-07-16_001.xad Page 3 of 10

Assay Class: High Sensitivity DNA Assay

7/16/2012 3:09:25 PM Created: C:\...ents and Settings\Bioanalyzer\2012-07-16\2012-07-16_001.xad 7/16/2012 3:28:30 PM Data Path: Modified:

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

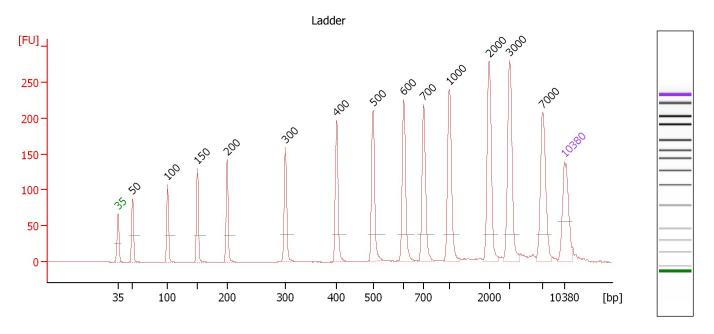
Size	Area
35	160
50	210
100	208
150	221
200	242
300	270
400	305
500	306
600	336
700	321
1000	366
2000	413
3000	411
7000	400
10380	214
	35 50 100 150 200 300 400 500 600 700 1000 2000 3000 7000

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2012-07-16\2012-07-16_001.xad Created: 7/16/2012 3:09:25 PM Modified: 7/16/2012 3:28:30 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

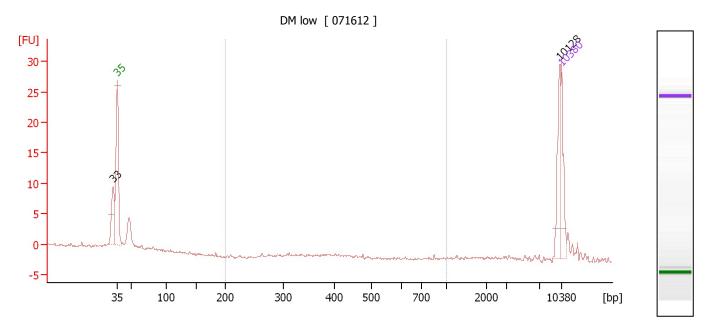
Peak	ta	hle	for	Lad	der
rcar	\ La	DIC	101	Lau	ucı

Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	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	L	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	L	500	150.00	454.5	Ladder Peak
9		600	150.00	378.8	Ladder Peak
10		700	150.00	324.7	Ladder Peak
11	L	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	L	7,000	150.00	32.5	Ladder Peak
15		10,380	75.00	10.9	Upper Marker

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High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2012-07-16\2012-07-16_001.xad 7/16/2012 3:09:25 PM 7/16/2012 3:28:30 PM Assay Class: Created: Data Path: Modified:

Electropherogram Summary Continued ...



Overall Results for sample 1: **DM low**

Number of peaks found: 2 0.0 Corr. Area 1:

DM low

Noise: 0.2

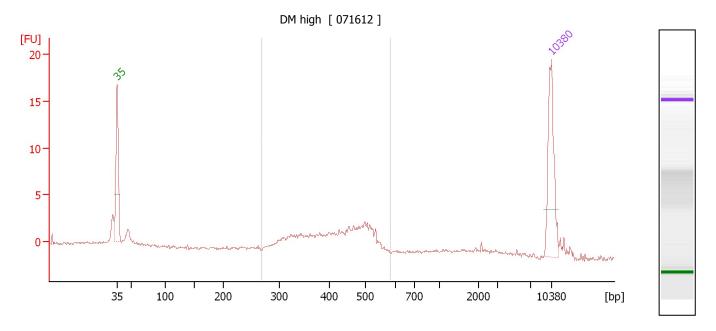
Peak table for sample 1:

Peak		Size [bp]	Conc. [pg/µl]	Mola	rity [pmol	/I]	Observations	
1		33	0.00	0.0				
2	4	35	125.00	5,411	.3		Lower Marker	
3		10,128	70.42	10.5				
4		10,380	75.00	10.9			Upper Marker	
Region	tak	ole for sample 1	: <u>DM low</u>					
From [bp]	Т	o [bp] Average Si [bp]	ize Molarity [pmol/l]	Conc. [pg/µl]	Corr. Area	% of Total	Size distribution in CV [%]	Co lor
200	1	,000 849	0.0	0.00	0.0	0	0.0	

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Assay Class: High Sensitivity DNA Assay Created: 7/16/2012 3:09:25 PM Data Path: C:\...ents and Settings\Bioanalyzer\2012-07-16\2012-07-16_001.xad Created: 7/16/2012 3:28:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 2: <u>DM high</u>

Number of peaks found: 0 Corr. Area 1: 41.4

Noise: 0.1

Peak table for sample 2: DM high Peak Size [bp] Molarity [pmol/l] **Observations** Conc. [pg/µl] 125.00 5,411.3 Lower Marker 1 35 10,380 75.00 Upper Marker 10.9

Region table for sample 2: <u>DM high</u>
From To [bp] Average Size Molarity Conc. Corr. % of

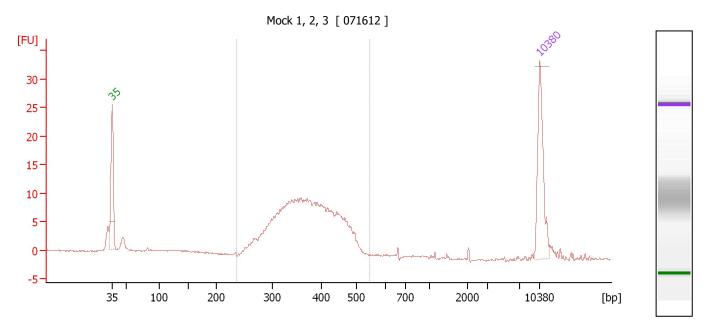
Size distribution in Co [pmol/l] **Total** [bp] [bp] [pg/µl] Area CV [%] lor 585 794.9 41.4 71 17.4 267 422 211.89

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Assay Class: High Sensitivity DNA Assay Created: 7/16/2012 3:09:25 PM Data Path: C:\...ents and Settings\Bioanalyzer\2012-07-16\2012-07-16_001.xad Created: 7/16/2012 3:28:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 3: Mock 1, 2, 3

Number of peaks found: 0 Corr. Area 1: 175.4

Noise: 0.1

 Peak table for sample 3 : Mock 1, 2, 3

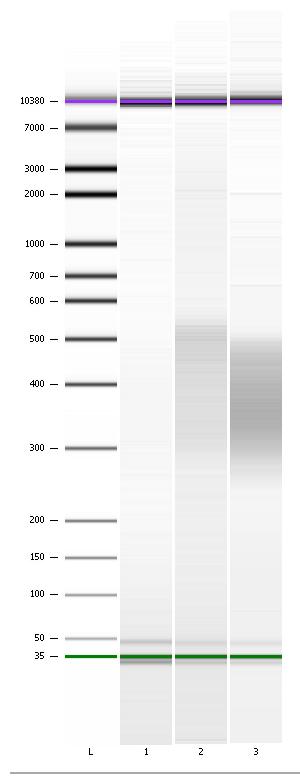
 Peak
 Size [bp]
 Conc. [pg/μl]
 Molarity [pmol/l]
 Observations

 1
 4
 35
 125.00
 5,411.3
 Lower Marker

 2
 10,380
 75.00
 10.9
 Upper Marker

Region table for sample 3: Mock 1, 2, 3 To [bp] Average Size **Molarity** Conc. Corr. % of Size distribution in Co **Total** [bp] [bp] [pmol/l] [pg/µl] Area CV [%] lor 373 2,453.3 175.4 88 16.3 236 543 583.04

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6

10

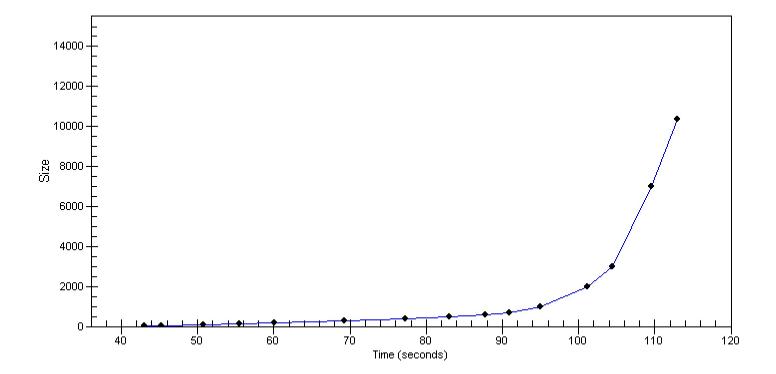
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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2012-07-16\2012-07-16_001.xad Created: 7/16/2012 3:09:25 PM Modified: 7/16/2012 3:28:30 PM

Curves

Standard Curve



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Assay Class:

High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2012-07-16\2012-07-16_001.xad 7/16/2012 3:09:25 PM 7/16/2012 3:28:30 PM Created: Data Path: Modified:

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.

Printed: