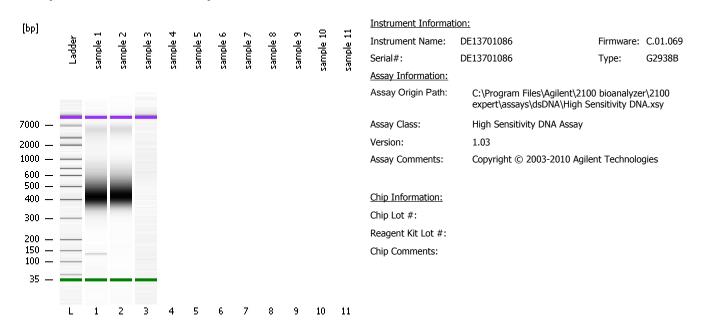
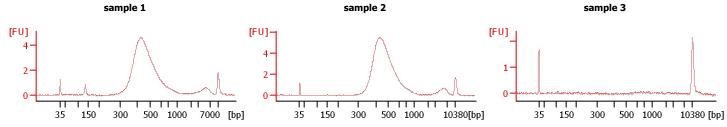
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Assay Class: High Sensitivity DNA Assay Created: 8/20/2012 10:19:17 AM Data Path: C:\...ents and Settings\Bioanalyzer\2012-08-20\2012-08-20_001.xad Modified: 8/20/2012 11:02:38 AM

Electrophoresis File Run Summary





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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2012-08-20\2012-08-20_001.xad Created: 8/20/2012 10:19:17 AM Modified: 8/20/2012 11:02:38 AM Electrophoresis File Run Summary (Chip Summary)

Sample Name Sample Comment Rest. Digest Status Observation Result Label Result Color sample 1

sample 1
sample 2
sample 3
sample 4
sample 5
sample 6
sample 7
sample 8
sample 9
sample 10
sample 10
chip Lot #

Reagent Kit Lot #

Chip Comments :

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Assay Class: High Sensitivity DNA Assay Created: 8/20/2012 10:19:17 AM Data Path: C:\...ents and Settings\Bioanalyzer\2012-08-20\2012-08-20_001.xad Modified: 8/20/2012 11:02:38 AM

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

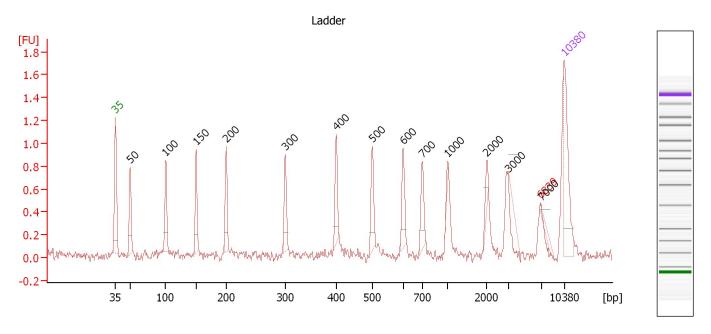
Ladder

1 35 160 2 50 210 3 100 208 4 150 221 5 200 242 6 300 270 7 400 205	Ladder Peak	Size	Area
3 100 208 4 150 221 5 200 242 6 300 270	1	35	160
4 150 221 5 200 242 6 300 270	2	50	210
5 200 242 6 300 270	3	100	208
6 300 270	4	150	221
	5	200	242
7 400	6	300	270
/ 400 305	7	400	305
8 500 306	8	500	306
9 600 336	9	600	336
10 700 321	10	700	321
11 1000 366	11	1000	366
12 2000 413	12	2000	413
13 3000 411	13	3000	411
14 7000 400	14	7000	400
15 10380 214	15	10380	214

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2012-08-20\2012-08-20_001.xad Created: 8/20/2012 10:19:17 AM Modified: 8/20/2012 11:02:38 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.0

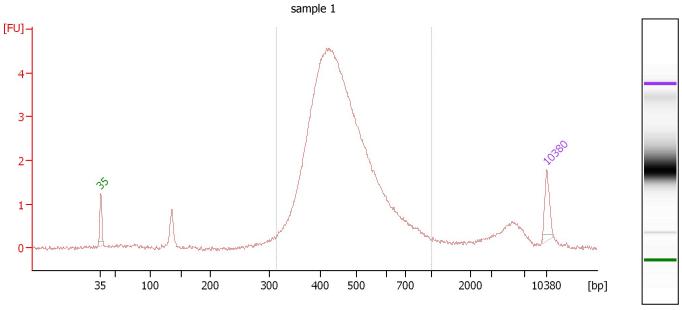
Peak	ta	hle	for	Lad	der
rear	. La	DIE	101	Lau	uei

Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		0	0.00	0.0	
2	10	35	125.00	5,411.3	Lower Marker
3		50	150.00	4,545.5	Ladder Peak
4		100	150.00	2,272.7	Ladder Peak
5		150	150.00	1,515.2	Ladder Peak
6		200	150.00	1,136.4	Ladder Peak
7		300	150.00	757.6	Ladder Peak
8		400	150.00	568.2	Ladder Peak
9		500	150.00	454.5	Ladder Peak
10		600	150.00	378.8	Ladder Peak
11		700	150.00	324.7	Ladder Peak
12		1,000	150.00	227.3	Ladder Peak
13		2,000	150.00	113.6	Ladder Peak
14		3,000	150.00	75.8	Ladder Peak
15	△	6,839	0.00	0.0	excluded peak
16		7,000	150.00	32.5	Ladder Peak
17		10,380	75.00	10.9	Upper Marker

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2012-08-20\2012-08-20_001.xad Created: 8/20/2012 10:19:17 AM Modified: 8/20/2012 11:02:38 AM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 1 : sample 1

Height Threshold [FU]:

Overall Results for sample 1 : sample 1

Number of peaks found: 0 Corr. Area 1: 64.1

Noise: 0.0

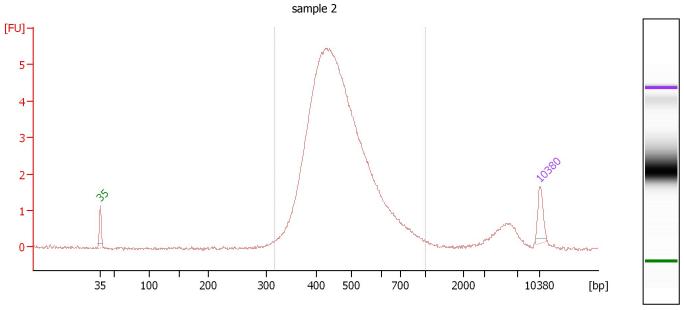
Peak table for sample 1:				<u>sample 1</u>		
	Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
	1	4	35	125.00	5,411.3	Lower Marker
	2		10,380	75.00	10.9	Upper Marker

Region table for sample 1: sample 1 From To [bp] Average Size Molarity Conc. Corr. % of Size distribution in Co [pmol/l] **Total** CV [%] [bp] [bp] [pg/µl] Area lor 314 1,000 469 17,287.3 5,096.59 64.1 90 21.5

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Assay Class: High Sensitivity DNA Assay Created: 8/20/2012 10:19:17 AM Data Path: C:\...ents and Settings\Bioanalyzer\2012-08-20\2012-08-20_001.xad Modified: 8/20/2012 11:02:38 AM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 2: sample 2

Height Threshold [FU]:

Overall Results for sample 2: sample 2

Number of peaks found: 0 Corr. Area 1: 74.5

Noise: 0.0

 Peak table for sample 2 : sample 2

 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

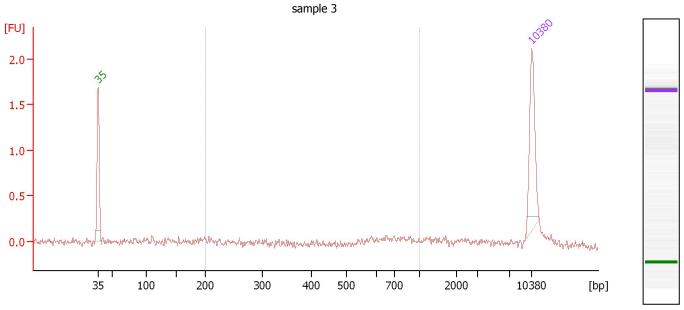
Region table for sample 2 : <u>sample 2</u>

From To [bp] Average Size **Molarity** Conc. Corr. % of Size distribution in Co [pmol/l] Total **CV** [%] [bp] [bp] [pg/µl] Area lor 317 1,000 474 18,029.1 5,396.51 74.5 94 20.4

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Assay Class: High Sensitivity DNA Assay Created: 8/20/2012 10:19:17 AM Data Path: C:\...ents and Settings\Bioanalyzer\2012-08-20\2012-08-20_001.xad Created: 8/20/2012 11:02:38 AM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 3: sample 3

Height Threshold [FU]:

Overall Results for sample 3: sample 3

Number of peaks found: 0 Corr. Area 1: 1.1

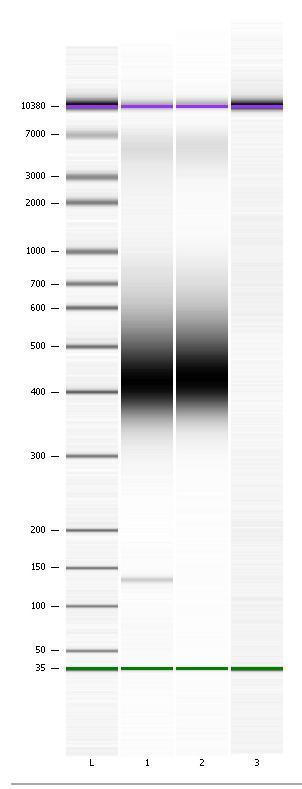
Noise: 0.0

Peak table for sample 3: sample 3 Size [bp] **Observations Peak** Conc. [pg/µl] Molarity [pmol/l] 35 125.00 5,411.3 Lower Marker 1 10,380 75.00 10.9 Upper Marker

Region table for sample 3: sample 3
From To [bp] Average Size Molarity Conc. Corr. % of Size dis

Size distribution in Co [pmol/l] **Total CV** [%] [bp] [bp] [pg/µl] Area lor 200 1,000 579 234.1 64.98 1.1 36 38.1

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Assay Class: High Sensitivity DNA Assay

8/20/2012 10:19:17 AM 8/20/2012 11:02:38 AM Created: C:\\'...ents and \(Settings\\ Bio\(analyzer\\ 2012-08-20\\ 2012-08-20\\ 2012-08-20\\ analyzer\\ 2012-08-20\\ 2012-08-20\\ analyzer\\ 2012-08-20\\ analyzer\\ 2012-08-20\\ analyzer\\ anal 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.