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Created:

Modified:

Printed:

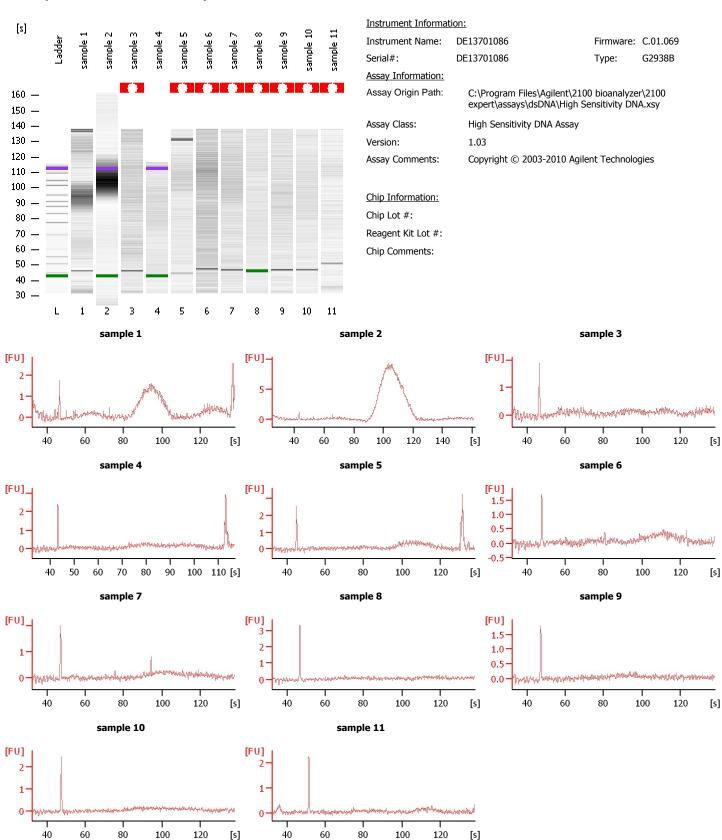
8/20/2013 12:00:33 PM

8/20/2013 11:17:52 AM

8/20/2013 11:59:48 AM

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad

Electrophoresis File Run Summary



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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:17:52 AM Modified: 8/20/2013 11:59:48 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest		Observation	Result Label	Result Color
sample 1		\sqcup	~			
sample 2		<u></u>	~			
sample 3			~			
sample 4			~			
sample 5			~			
sample 6			~			
sample 7			~			
sample 8			~			
sample 9			~			
sample 10			~			
sample 11			~			
Ladder			~			
Chip Lot #			Reage	ent Kit Lot #		

Chip Comments:

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Assay Class: High Sensitivity DNA Assay Created: 8/20/2013 11:17:52 AM Data Path: C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:59:48 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

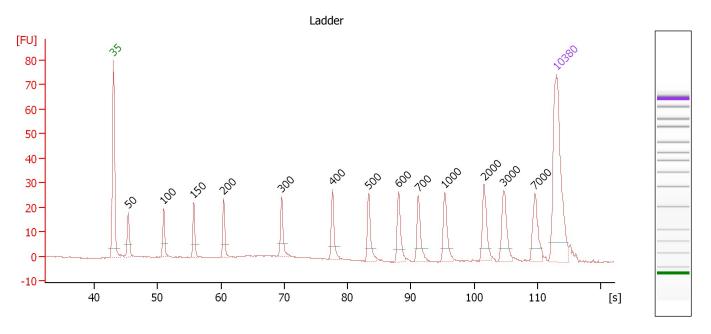
Ladder

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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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:17:52 AM Modified: 8/20/2013 11:59:48 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

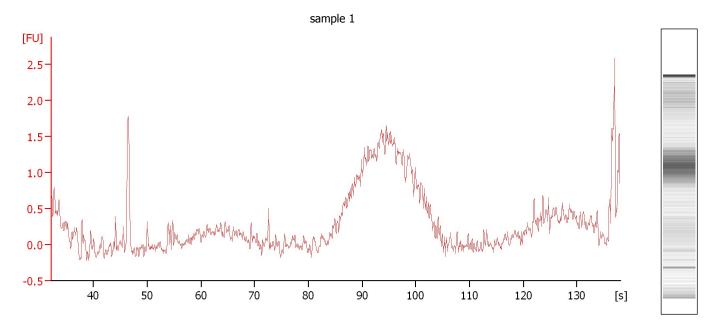
Peak	table	for	Ladder
rcan	cavic	101	Lauuci

Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	18	35	125.00	5,411.3	Lower Marker
2		50	150.00	4,545.5	Ladder Peak
3	L	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	L	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	L	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	L	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	A	10,380	75.00	10.9	Upper Marker

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:17:52 AM Modified: 8/20/2013 11:59:48 AM

Electropherogram Summary Continued ...



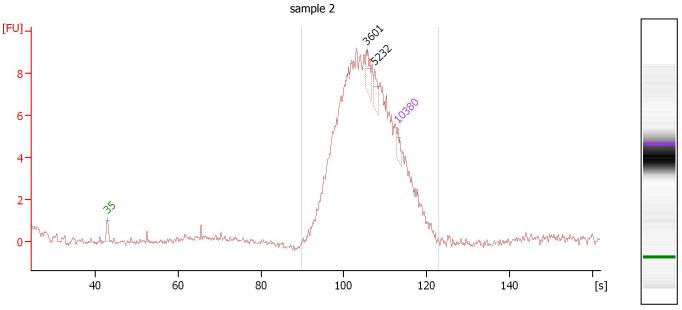
Overall Results for sample 1 : sample 1

Noise: 0.1

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad 8/20/2013 11:17:52 AM 8/20/2013 11:59:48 AM Created: Data Path: Modified:

Electropherogram Summary Continued ...



Setpoint Deviations for sample 2: sample 2

Height Threshold [FU]:

Overall Results for sample 2: sample 2

Number of peaks found: 2 Corr. Area 1: 113.0

sample 2

Conc. [pg/µl]

5,461.8

Noise: 0.1

Size [bp]

20,075 5,181

Peak table for sample 2:

Peak

653

1 2 3	•	35 3,601 5,232	125.00 155.16 109.09	5,411. 65.3 31.6	.3		Lower Marker
4		10,380	75.00	10.9		Upper Marker	
Region From [bp]		ole for sample 2 : o [bp] Average Siz [bp]		Conc. [pg/µl]	Corr. Area	% of Total	Size distribution in CV [%]

9,604.26

Molarity [pmol/l]

113.0

98

Observations

79.1

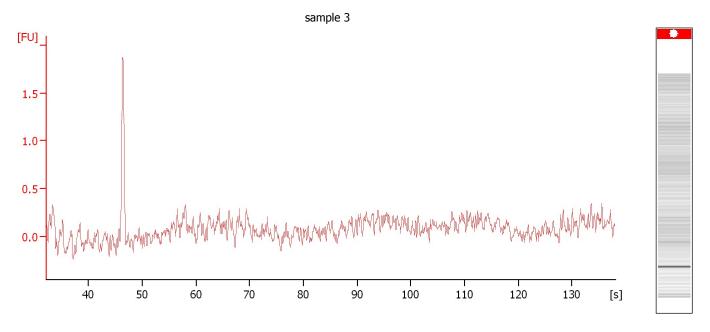
Co

lor

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:17:52 AM Modified: 8/20/2013 11:59:48 AM

Electropherogram Summary Continued ...



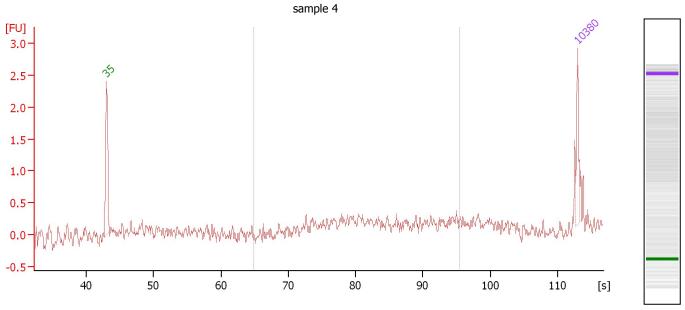
Overall Results for sample 3: sample 3

Noise: 0.1

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Assay Class: High Sensitivity DNA Assay Created: 8/20/2013 11:17:52 AM Data Path: C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:17:52 AM Modified: 8/20/2013 11:59:48 AM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 4: sample 4

Height Threshold [FU]:

Overall Results for sample 4: sample 4

Number of peaks found: 0 Corr. Area 1: 3.3

Noise: 0.1

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

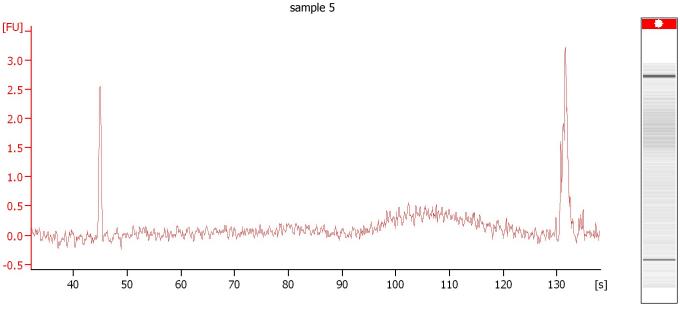
Region table for sample 4: sample 4

From To [bp] Average Size Molarity Conc. Corr. % of Size distribution in Co [pmol/l] **Total CV** [%] [bp] [bp] [pg/µl] Area lor 249 1,000 542 1,351.2 424.40 3.3 56 33.2

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:17:52 AM Modified: 8/20/2013 11:59:48 AM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 5 : <u>sample 5</u>

4

Height Threshold [FU]:

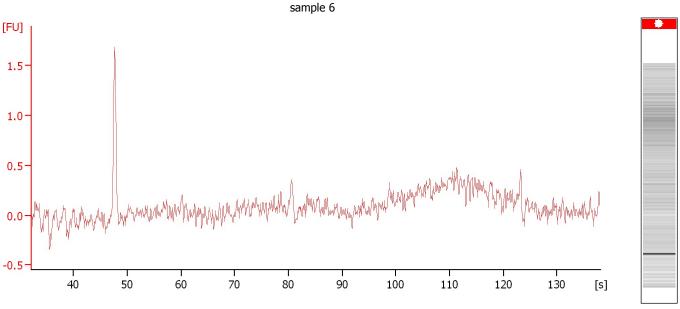
Overall Results for sample 5 : sample 5

Noise: 0.1

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:17:52 AM Modified: 8/20/2013 11:59:48 AM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 6 : <u>sample 6</u>

2

Height Threshold [FU]:

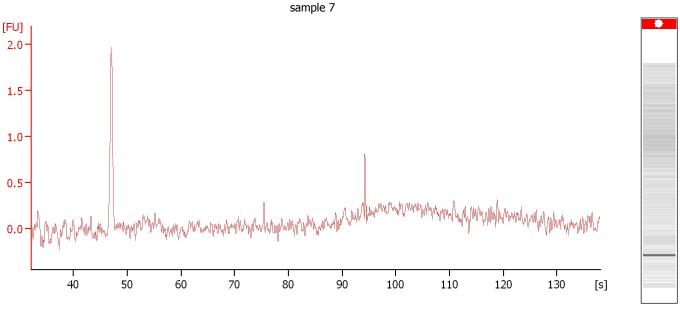
Overall Results for sample 6: sample 6

Noise: 0.1

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:17:52 AM Modified: 8/20/2013 11:59:48 AM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 7 : <u>sample 7</u>

3

Height Threshold [FU]:

Overall Results for sample 7: sample 7

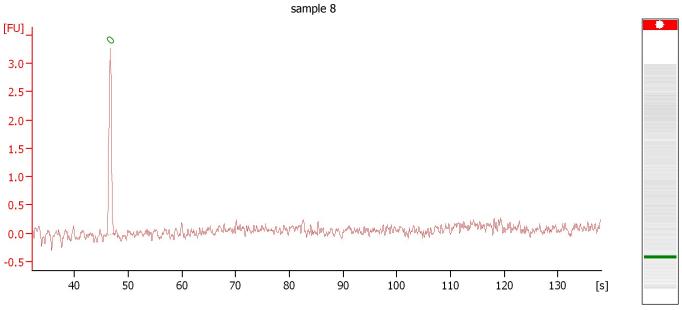
Noise: 0.1

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Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad

Created: 8/20/2013 11:17:52 AM
Modified: 8/20/2013 11:59:48 AM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 8 : sample 8

Height Threshold [FU]: 2

Overall Results for sample 8: sample 8

Number of peaks found: 0 Noise: 0.1

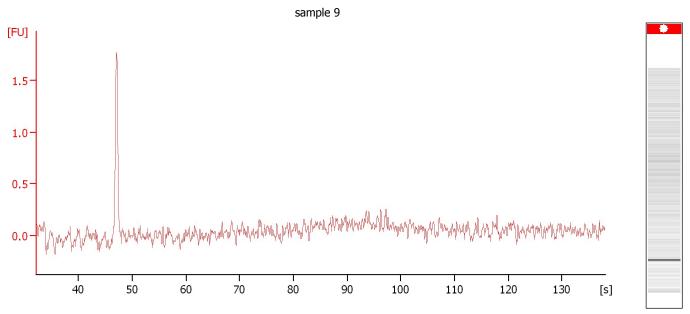
Peak table for sample 8 : <u>sample 8</u>

PeakSize [bp]Conc. [pg/μl]Molarity [pmol/l]Observations1 \bullet 0.000.0Lower Marker

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:17:52 AM Modified: 8/20/2013 11:59:48 AM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 9 : <u>sample 9</u>

Height Threshold [FU]: 3

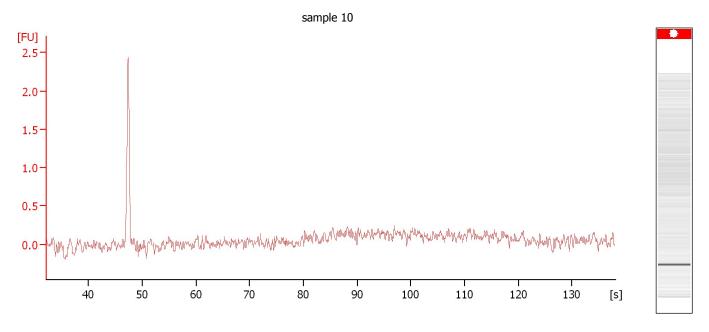
Overall Results for sample 9: sample 9

Noise: 0.1

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:17:52 AM Modified: 8/20/2013 11:59:48 AM

Electropherogram Summary Continued ...



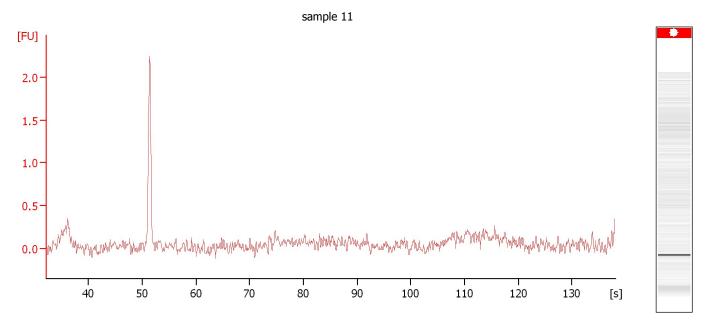
Overall Results for sample 10 : sample 10

Noise: 0.1

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:17:52 AM Modified: 8/20/2013 11:59:48 AM

Electropherogram Summary Continued ...



Overall Results for sample 11 : sample 11

Noise: 0.1

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High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad 8/20/2013 11:17:52 AM 8/20/2013 11:59:48 AM Assay Class: Created: Modified: Data Path: **Gel Image** [5] sample 10 sample 11 sample 2 sample 4 sample 8 sample 1 Ladder 160 — 155 — 150 — 145 — 140 — 135 — 130 — 125 — 120 — 115 -110 105 — 100 -95 -90 85 80 -75 — 70 — 65 — 60 — 55 — 50 **—** 45 — 40 — 35 — 30 — 25 — L 2 3 5 10 11

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2013-08-20\2013-08-20_001.xad Created: 8/20/2013 11:17:52 AM Modified: 8/20/2013 11:59:48 AM

Run Logbook

Description Run ended on port 1 (Number of wells acquired: 12)	Number	Source Instrument	Category Run	Sub Category	Time 8/20/2013 11:59:09 AM	Time Zone (GMT07:00) Pacific Standard Time		Host D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanal yzer\2013-08-20\2013-08-20_00 1.xad))	Instrument	Run		8/20/2013 11:17:57 AM	(GMT07:00) Pacific Standard Time		D8XSMGH1
Product Number : G2938B		Instrument	Run		8/20/2013 11:17:57 AM	(GMT07:00) Pacific Standard Time		D8XSMGH1
Name :		Instrument	Run		8/20/2013 11:17:57 AM	(GMT07:00) Pacific Standard Time		D8XSMGH1
Vendor : Agilent Technologies	:	Instrument	Run		8/20/2013 11:17:57 AM	(GMT07:00) Pacific Standard Time		D8XSMGH1
Serial# : DE13701086		Instrument	Run		8/20/2013 11:17:57 AM	(GMT07:00) Pacific Standard Time		D8XSMGH1
Firmware : C.01.069		Instrument	Run		8/20/2013 11:17:57 AM	(GMT07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		8/20/2013 11:17:57 AM	(GMT07:00) Pacific Standard Time	UC Davis	D8XSMGH1