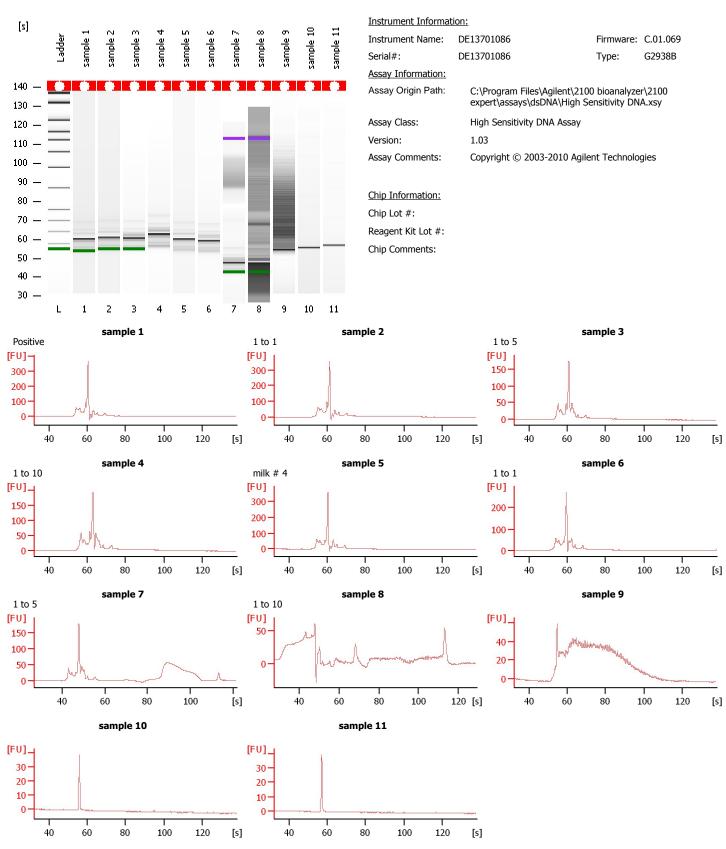
2014-01-07_001.xad Page 1 of 17

Assay Class: High Sensitivity DNA Assay Created: 1/7/2014 11:06:30 AM Data Path: C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad I:d7/2014 11:47:52 AM

Electrophoresis File Run Summary



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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
sample 1	Positive		~			
sample 2	1 to 1		~			
sample 3	1 to 5		~			
sample 4	1 to 10		~			
sample 5	milk # 4		~			
sample 6	1 to 1		~			
sample 7	1 to 5		~			
sample 8	1 to 10		~			
sample 9			~			
sample 10			~			
sample 11			~			
Ladder			~			
Chip Lot #			Reagen	t Kit Lot #		

Chip Comments:

2014-01-07_001.xad 3 of 17 Page

Assay Class: High Sensitivity DNA Assay

1/7/2014 11:06:30 AM Created: C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad 1/7/2014 11:47:52 AM 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

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

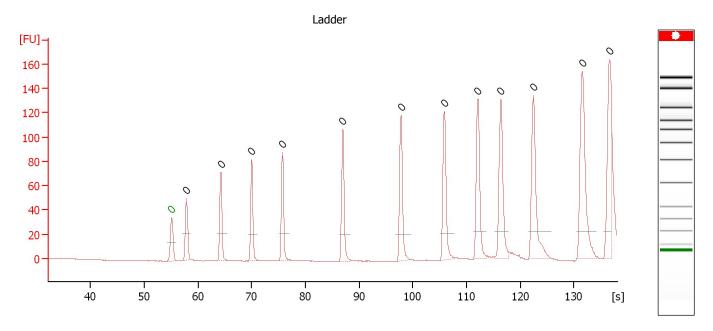
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1/7/2014 12:36:13 PM

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

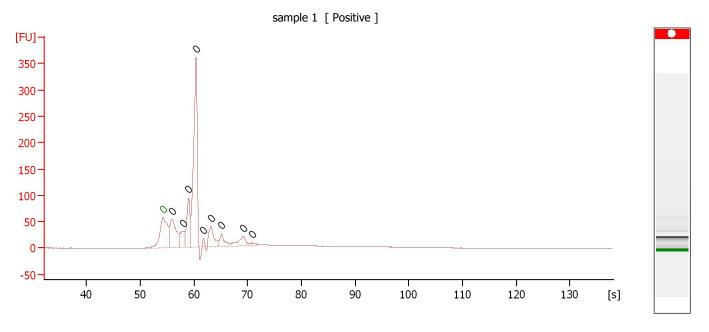
Peak table for Ladder

Pea k		Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	18	55.05	0	0.00	0.0	Lower Marker
2		57.85	0	0.00	0.0	Ladder Peak
3		64.25	0	0.00	0.0	Ladder Peak
4		70.00	0	0.00	0.0	Ladder Peak
5	L	75.70	0	0.00	0.0	Ladder Peak
6		86.95	0	0.00	0.0	Ladder Peak
7		97.80	0	0.00	0.0	Ladder Peak
8		105.85	0	0.00	0.0	Ladder Peak
9		112.10	0	0.00	0.0	Ladder Peak
10		116.40	0	0.00	0.0	Ladder Peak
11		122.45	0	0.00	0.0	Ladder Peak
12		131.65	0	0.00	0.0	Ladder Peak
13	L	136.65	0	0.00	0.0	Ladder Peak

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electropherogram Summary Continued ...



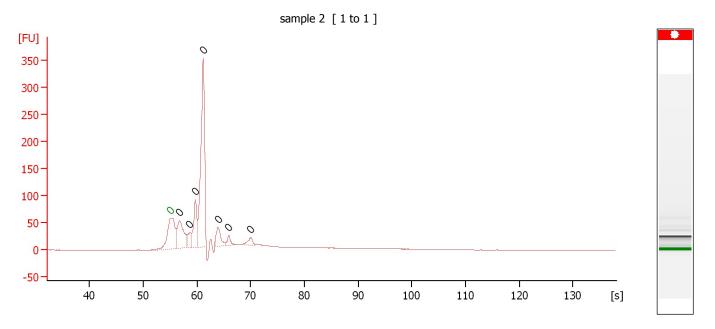
Overall Results for sample 1 : sample 1

Peak tal	Peak table for sample 1 : sample 1								
Pea	Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations				
k									
1 🔥	54.20	0	0.00	0.0	Lower Marker				
2	55.90	0	0.00	0.0					
3	57.95	0	0.00	0.0					
4	58.95	0	0.00	0.0					
5	60.35	0	0.00	0.0					
6	61.75	0	0.00	0.0					
7	63.15	0	0.00	0.0					
8	65.15	0	0.00	0.0					
9	69.15	0	0.00	0.0					
10	70.80	0	0.00	0.0					

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electropherogram Summary Continued ...



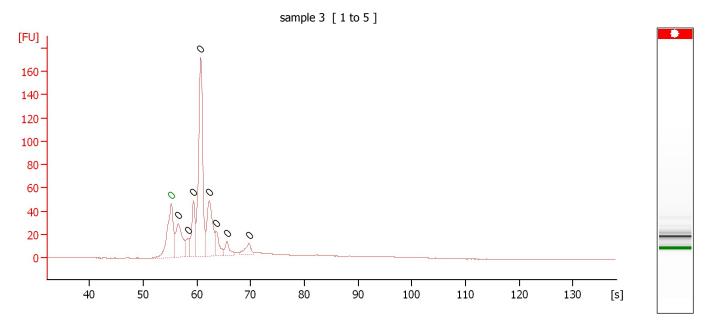
Overall Results for sample 2 : sample 2

Peak	Peak table for sample 2 : sample 2								
Pea		Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations			
k									
1	18	55.05	0	0.00	0.0	Lower Marker			
2		56.75	0	0.00	0.0				
3		58.65	0	0.00	0.0				
4		59.65	0	0.00	0.0				
5		61.15	0	0.00	0.0				
6		63.90	0	0.00	0.0				
7		65.90	0	0.00	0.0				
8		70.00	0	0.00	0.0				

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electropherogram Summary Continued ...



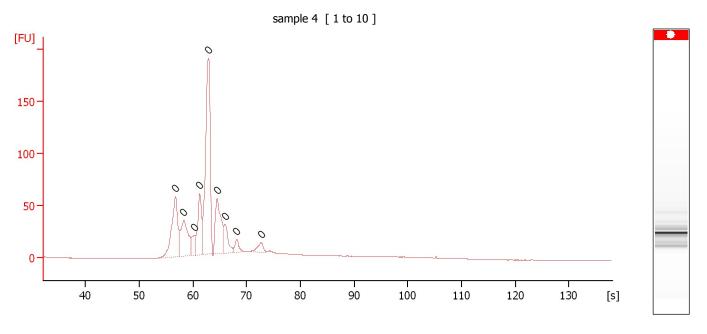
Overall Results for sample 3: sample 3

Peak t	Peak table for sample 3: <u>sample 3</u>								
Pea	Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations				
k		_							
1	§ 55.20	0	0.00	0.0	Lower Marker				
2	56.45	0	0.00	0.0					
3	58.35	0	0.00	0.0					
4	59.30	0	0.00	0.0					
5	60.70	0	0.00	0.0					
6	62.25	0	0.00	0.0					
7	63.55	0	0.00	0.0					
8	65.55	0	0.00	0.0					
9	69.65	0	0.00	0.0					

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electropherogram Summary Continued ...



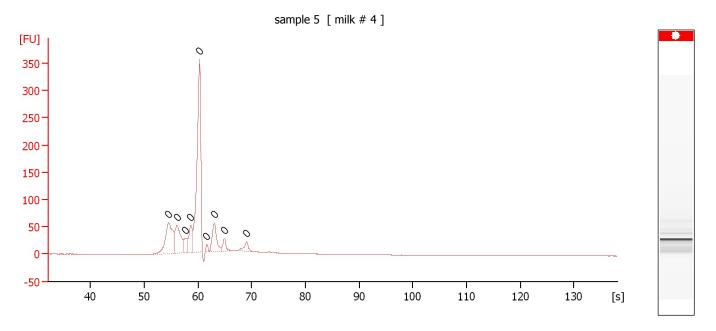
Overall Results for sample 4: sample 4

Peak t	Peak table for sample 4: sample 4									
Pea k	Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations					
1	56.70	0	0.00	0.0						
2	58.25	0	0.00	0.0						
3	60.15	0	0.00	0.0						
4	61.25	0	0.00	0.0						
5	62.85	0	0.00	0.0						
6	64.45	0	0.00	0.0						
7	65.95	0	0.00	0.0						
8	68.15	0	0.00	0.0						
9	72.65	0	0.00	0.0						

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electropherogram Summary Continued ...



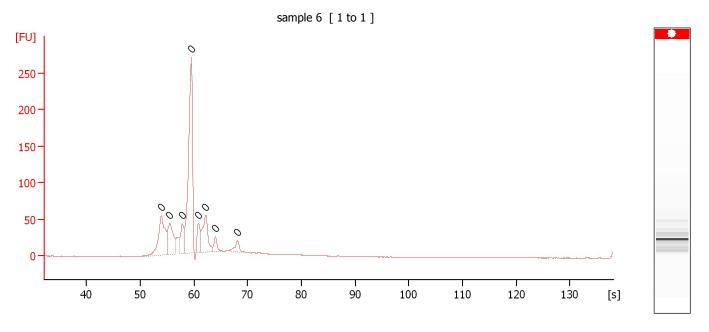
Overall Results for sample 5 : sample 5

Peak t	Peak table for sample 5 : sample 5								
Pea k	Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations				
1	54.50	0	0.00	0.0					
2	56.05	0	0.00	0.0					
3	57.70	0	0.00	0.0					
4	58.65	0	0.00	0.0					
5	60.25	0	0.00	0.0					
6	61.60	0	0.00	0.0					
7	63.00	0	0.00	0.0					
8	64.90	0	0.00	0.0					
9	69.05	0	0.00	0.0					

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electropherogram Summary Continued ...



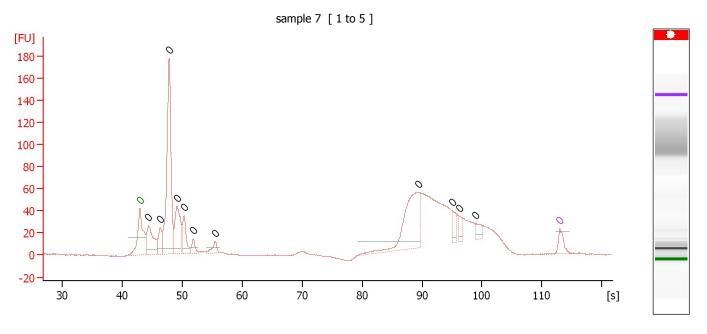
Overall Results for sample 6: sample 6

Peak ta	Peak table for sample 6: sample 6								
Pea	Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations				
k									
1	53.90	0	0.00	0.0					
2	55.45	0	0.00	0.0					
3	57.80	0	0.00	0.0					
4	59.45	0	0.00	0.0					
5	60.80	0	0.00	0.0					
6	62.15	0	0.00	0.0					
7	64.00	0	0.00	0.0					
8	68.10	0	0.00	0.0					

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electropherogram Summary Continued ...



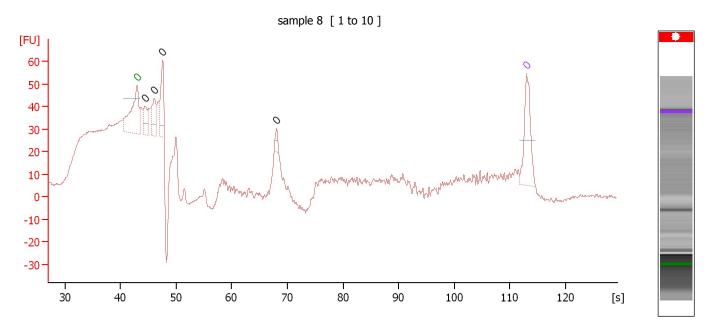
Overall Results for sample 7: sample 7

Peak	Peak table for sample 7: sample 7							
Pea k		Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations		
1	18	43.00	0	0.00	0.0	Lower Marker		
2		44.48	0	0.00	0.0			
3		46.35	0	0.00	0.0			
4		47.83	0	0.00	0.0			
5		49.17	0	0.00	0.0			
6		50.34	0	0.00	0.0			
7		51.90	0	0.00	0.0			
8		55.52	0	0.00	0.0			
9		89.47	0	0.00	0.0			
10		95.06	0	0.00	0.0			
11		96.18	0	0.00	0.0			
12		99.00	0	0.00	0.0			
13		113.00	0	0.00	0.0	Upper Marker		

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electropherogram Summary Continued ...



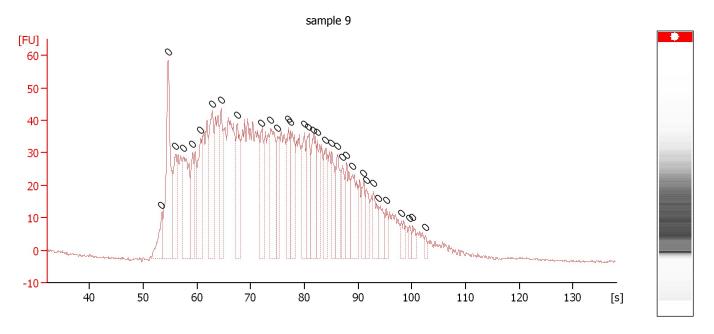
Overall Results for sample 8 : sample 8

Peal	Peak table for sample 8 : sample 8								
Pea		Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations			
k									
1	10	43.00	0	0.00	0.0	Lower Marker			
2		44.40	0	0.00	0.0				
3		46.04	0	0.00	0.0				
4		47.58	0	0.00	0.0				
5		68.04	0	0.00	0.0				
6	8	113.00	0	0.00	0.0	Upper Marker			

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electropherogram Summary Continued ...



Overall Results for sample 9: sample 9

Peak tal	Peak table for sample 9: sample 9							
Pea k	Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations			
1	53.40	0	0.00	0.0				
2	54.60	0	0.00	0.0				
3	56.00	0	0.00	0.0				
4	57.40	0	0.00	0.0				
5	59.15	0	0.00	0.0				
6	60.70	0	0.00	0.0				
7	62.80	0	0.00	0.0				
8	64.50	0	0.00	0.0				
9	67.45	0	0.00	0.0				
10	72.05	0	0.00	0.0				
11	73.70	0	0.00	0.0				
12	75.05	0	0.00	0.0				
13	77.05	0	0.00	0.0				
14	77.40	0	0.00	0.0				
15	79.95	0	0.00	0.0				
16	80.80	0	0.00	0.0				
17	81.60	0	0.00	0.0				
18	82.40	0	0.00	0.0				
19	83.90	0	0.00	0.0				
20	85.10	0	0.00	0.0				
21	86.15	0	0.00	0.0				
22	87.05	0	0.00	0.0				
23	87.85	0	0.00	0.0				
24	88.90	0	0.00	0.0				
25	91.05	0	0.00	0.0				
26	91.55	0	0.00	0.0				
27	92.85	0	0.00	0.0				

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

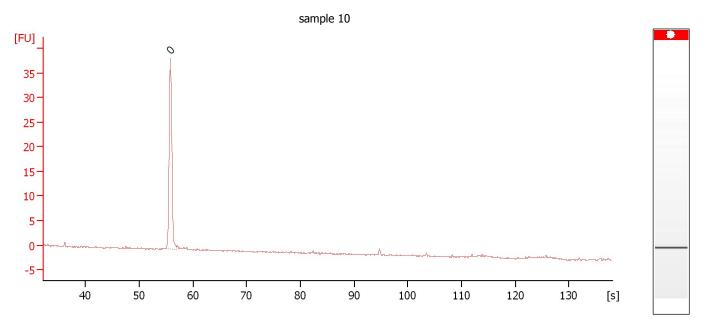
Electropherogram Summary Continued ...

Peak table for sample 9 : <u>sample 9</u>								
	Pea	Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations		
	k							
	28	93.85	0	0.00	0.0			
:	29	95.20	0	0.00	0.0			
:	30	98.05	0	0.00	0.0			
	31	99.60	0	0.00	0.0			
	32	100.20	0	0.00	0.0			
:	33	102.55	0	0.00	0.0			

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electropherogram Summary Continued ...



Overall Results for sample 10 : sample 10

Number of peaks found: 0 Noise: 0.1

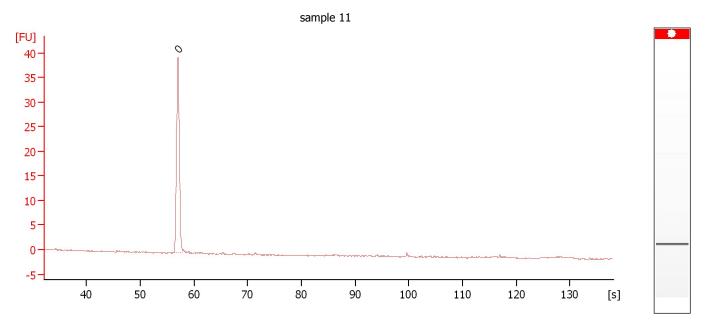
Peak table for sample 10 : <u>sample 10</u>

Pea	Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	55.75	0	0.00	0.0	

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Assay Class: High Sensitivity DNA Assay C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad C:\...ents and Settings\Bioanalyzer\2014-01-07\2014-01-07_001.xad

Electropherogram Summary Continued ...



Overall Results for sample 11 : sample 11

Number of peaks found: 0 Noise: 0.1

Peak table for sample 11 : <u>sample 11</u>

Pea k	Aligned Migration Time [s]	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	57.00	0	0.00	0.0	

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1/7/2014 11:06:30 AM 1/7/2014 11:47:52 AM Assay Class: High Sensitivity DNA Assay Created: C:\\'...ents and \(\'Settings\\Bio\'analyzer\\2014-01-07\\2014-01-07_001.xad\) Data Path: Modified:

Gel Image

