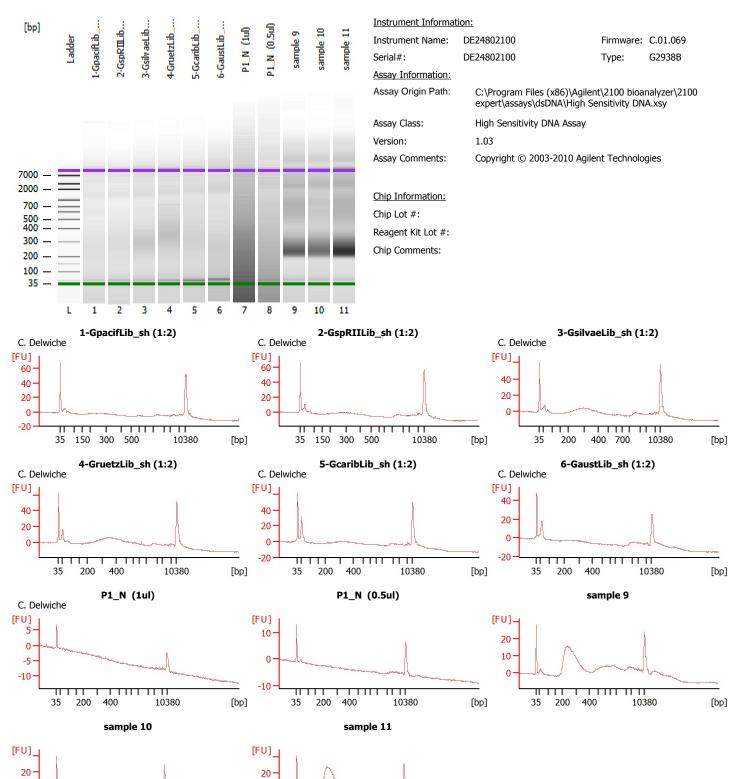
Assay Class: High Sensitivity DNA Assay Created: 6/14/2017 4:35:24 PM Data Path: C:\...gh Sensitivity DNA Assay_DE24802100_2017-06-14_16-35-25.xad Created: 6/15/2017 1:56:38 PM

Electrophoresis File Run Summary



10380

35 200 400

10

10380

[bp]

10

35 200 400

[bp]

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Assay Class: High Sensitivity DNA Assay
Data Path: C:\...gh Sensitivity DNA Assay_DE24802100_2017-06-14_16-35-25.xad

Created: 6/14/2017 4:35:24 PM Modified: 6/15/2017 1:56:38 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Statu Observation s	Result Label Result Color
1-GpacifLib_sh (1:2)	C. Delwiche		✓	
2-GspRIILib_sh (1:2)	C. Delwiche		✓	
3-GsilvaeLib_sh (1:2)	C. Delwiche		✓	
4-GruetzLib_sh (1:2)	C. Delwiche		✓	
5-GcaribLib_sh (1:2)	C. Delwiche		✓	
6-GaustLib_sh (1:2)	C. Delwiche		✓	
P1_N (1ul)	C. Delwiche		✓	
P1_N (0.5ul)			✓	
sample 9			✓	
sample 10			✓	
sample 11			✓	
Ladder			~	
Chip Lot #				Reagent Kit Lot #

Chip Comments:

Printed:

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Assay Class: High Sensitivity DNA Assay

6/14/2017 4:35:24 PM Created: C:\...gh Sensitivity DNA Assay_DE24802100_2017-06-14_16-35-25.xad 6/15/2017 1:56:38 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

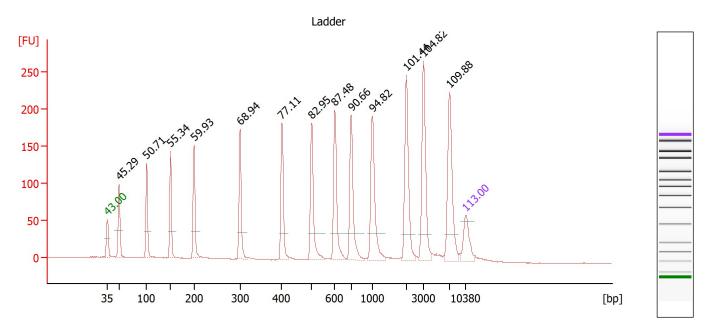
Filter Width [s]: 0.5 Polynomial Order: 4

Ladder

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

Printed:

Electropherogram Summary



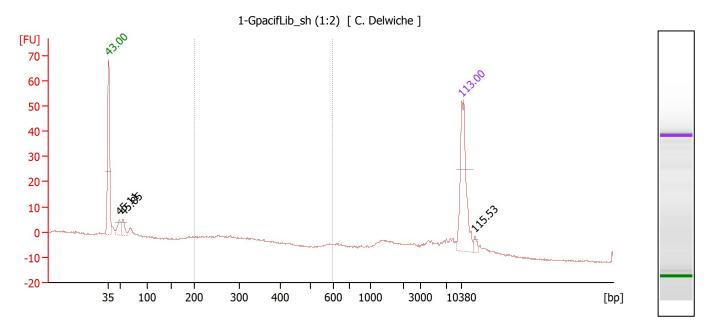
Overall Results for Ladder

Noise: 0.3

Doak	table	for	Ladder
reak	table	TOF	Lauuer

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		7,000	150.00	32.5	Ladder Peak
15		10,380	75.00	10.9	Upper Marker

Electropherogram Summary Continued ...



Overall Results for sample 1 : 1-GpacifLib_sh (1:2)

Number of peaks found: 3 Corr. Area 1: 31.4

Noise: 0.3

Peak table for sample 1: 1-GpacifLib sh (1:2)

Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2		49	21.47	666.3	
3		55	19.53	536.4	
4		10,380	75.00	10.9	Upper Marker
5		13,121	0.00	0.0	

Region table for sample 1: 1-GpacifLib sh (1:2)

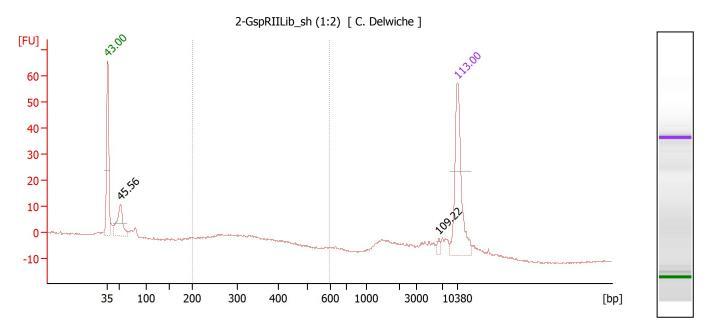
From [bp]	To [b _l	o] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	597	31.4	23	309	27.6	52.89	279.7	

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Electropherogram Summary Continued ...



Overall Results for sample 2: 2-GspRIILib sh (1:2)

Number of peaks found: 2 Corr. Area 1: 47.2

Noise: 0.3

Peak table for sample 2: 2-GspRIILib sh (1:2)

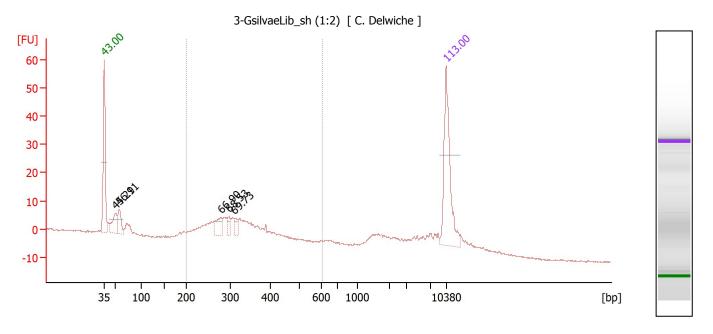
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2		52	59.38	1,715.3	
3		6,480	3.22	0.8	
4		10,380	75.00	10.9	Upper Marker

Region table for sample 2: 2-GspRIILib_sh (1:2)

From [bp]	To [b _l	p] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	597	47 2	32	317	18.8	64 10	321 1	

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Electropherogram Summary Continued ...



Overall Results for sample 3: 3-GsilvaeLib_sh (1:2)

Number of peaks found: 5 Corr. Area 1: 153.3

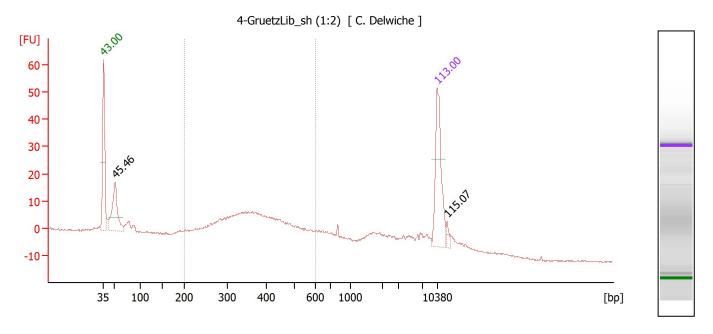
Noise: 0.3

Peak table for sample 3:			3-GsilvaeLib s	<u>sh (1:2)</u>		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	
1	4	35	125.00	5,411.3	Lower Marker	
2		50	32.51	985.1		
3		58	27.21	716.3		
4		277	22.73	124.2		
5		296	9.62	49.3		
6		310	9.50	46.4		
7		10,380	75.00	10.9	Upper Marker	

Region table for sample 3: 3-GsilvaeLib_sh (1:2)

From [bp]	To [b	p] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	606	153.3	53	332	24.7	235.27	1,155.1	

Electropherogram Summary Continued ...



Overall Results for sample 4: 4-GruetzLib sh (1:2)

Number of peaks found: 2 Corr. Area 1: 238.7

Noise: 0.2

Peak table for sample 4: 4-GruetzLib sh (1:2)

Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2		52	98.00	2,880.1	
3		10,380	75.00	10.9	Upper Marker
4		12,617	0.00	0.0	

Region table for sample 4: 4-GruetzLib sh (1:2)

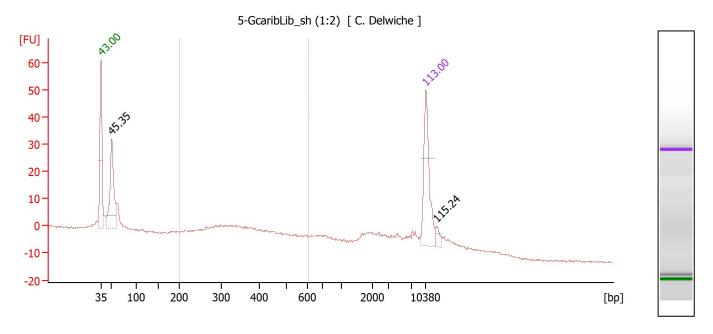
From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	599	238.7	53	380	24.6	374 44	1 630 9	

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Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DE24802100_2017-06-14_16-35-25.xad Created: 6/14/2017 4:35:24 PM Modified: 6/15/2017 1:56:38 PM

Electropherogram Summary Continued ...



Overall Results for sample 5: <u>5-GcaribLib_sh (1:2)</u>

Number of peaks found: 2 Corr. Area 1: 107.3

Noise: 0.3

Peak table for sample 5: <u>5-GcaribLib sh (1:2)</u>

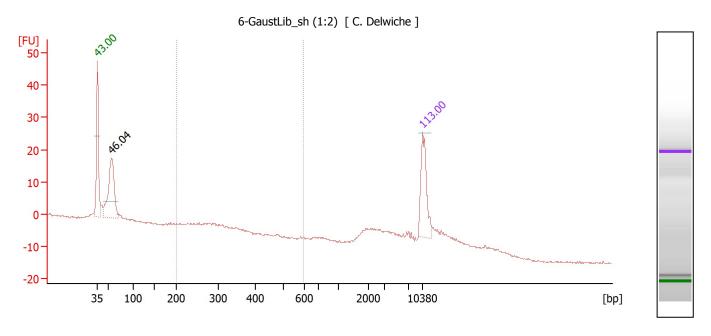
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2		51	146.87	4,401.1	
3		10,380	75.00	10.9	Upper Marker
4		12,800	0.00	0.0	

Region table for sample 5 : <u>5-GcaribLib_sh (1:2)</u>

From [bp]	To [b _l	p] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	606	107 3	33	374	26.4	182 22	814 6	

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Electropherogram Summary Continued ...



Overall Results for sample 6: 6-GaustLib sh (1:2)

Number of peaks found: 1 Corr. Area 1: 23.6

Noise: 0.3

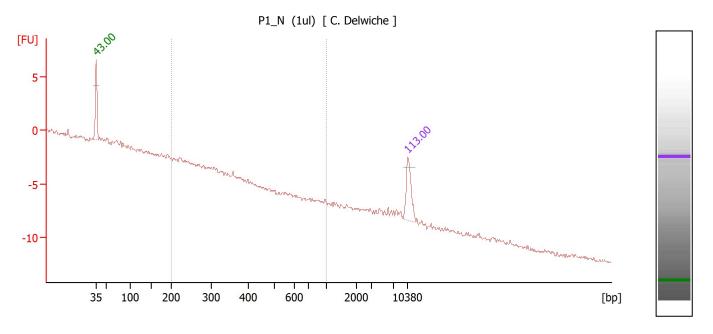
Peak table for sample 6: 6-GaustLib sh (1:2)

Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2		57	211.32	5,625.5	
3		10,380	75.00	10.9	Upper Marker

Region table for sample 6: 6-GaustLib sh (1:2)

From [bp]	To [b _l	p] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	596	23.6	13	284	15.8	71.34	389.7	

Electropherogram Summary Continued ...



Overall Results for sample 7: P1 N (1ul)

Number of peaks found: 0 Corr. Area 1: 0.8

Noise: 0.2

Peak table for sample 7 : P1 N (1ul)

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

 1
 ◀
 35
 125.00
 5,411.3
 Lower Marker

 2
 ▶
 10,380
 75.00
 10.9
 Upper Marker

Region table for sample 7: P1 N (1ul)

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	1 000 0 8	12	251	16.8	16 53	102 1	

High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DE24802100_2017-06-14_16-35-25.xad 6/14/2017 4:35:24 PM 6/15/2017 1:56:38 PM Assay Class: Created: Modified: Data Path: **Gel Image** 4-GruetzLib_... 1-GpacifLib_... 2-GspRIILib... 3-GsilvaeLib... 5-GcaribLib_... 6-GaustLib_... [bp] P1_N (0.5ul) P1_N (1ul) sample 10 sample 11 sample 9 Ladder 10380 7000 3000 2000 1000 700 -600 500 -400 -300 200 150 100 -50 -35 -1 2 3 4 5 6 7 8 9 10 11

Assay Class: Data Path:

High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DE24802100_2017-06-14_16-35-25.xad

Created: Modified:

6/14/2017 4:35:24 PM 6/15/2017 1:56:38 PM

Curves

Standard Curve

