Assay Class: High Sensitivity DNA Assay Created: 9/15/2017 12:39:32 PM Data Path: C:\...gh Sensitivity DNA Assay_DE13805510_2017-09-15_12-39-33.xad Created: 9/15/2017 12:058 PM

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]: 3
Area Threshold: 0.1
Width Threshold [s]: 0.6
Baseline Plateau [s]: 0.5

Filter Settings

Filter Width [s]: 0.5 Polynomial Order: 4

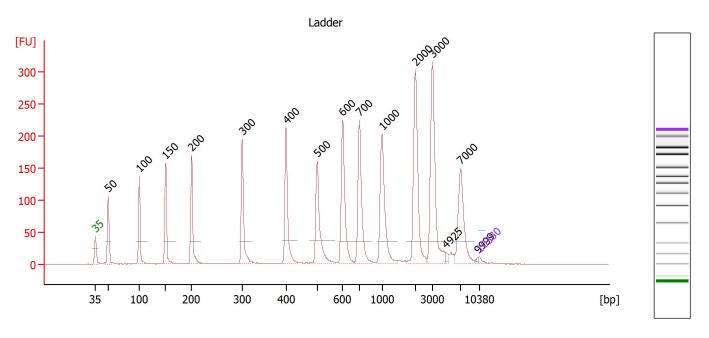
Ladder

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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Electropherogram Summary



Overall Results for Ladder

Noise: 0.3

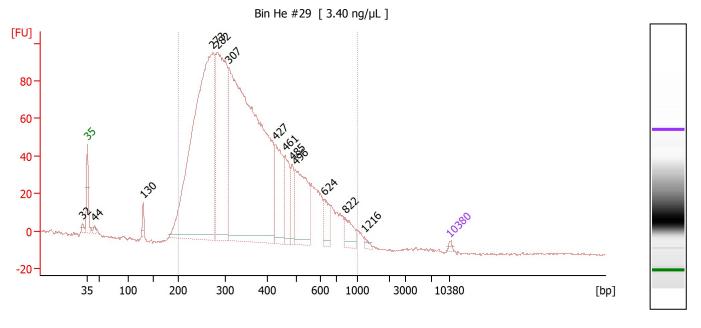
Doak	table	for	Ladder
reak	table	тог	Lauuer

Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2	L	50	150.00	4,545.5	Ladder Peak
3		100	150.00	2,272.7	Ladder Peak
4	L	150	150.00	1,515.2	Ladder Peak
5		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		500	150.00	454.5	Ladder Peak
9		600	150.00	378.8	Ladder Peak
10		700	150.00	324.7	Ladder Peak
11		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		4,925	0.00	0.0	
15		7,000	150.00	32.5	Ladder Peak
16		9,999	0.00	0.0	
17	8	10,380	75.00	10.9	Upper Marker

Page 3 of 11

Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DE13805510_2017-09-15_12-39-33.xad Created: 9/15/2017 12:39:32 PM Modified: 9/15/2017 1:20:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 4: Bin He #29

Number of peaks found: 13 Corr. Area 1: 2,497.5

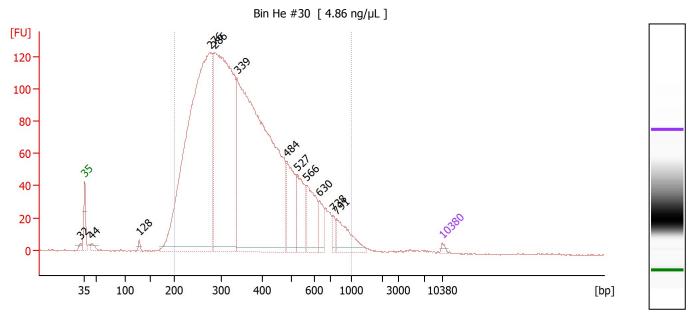
Noise: 0.4

Peak ta	ble	for sample 4:	Bin He #29		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		32	0.00	0.0	
2	4	35	125.00	5,411.3	Lower Marker
3		44	202.52	7,036.3	
4		130	469.07	5,486.0	
5		273	22,884.44	127,167.4	
6		282	11,296.88	60,786.9	
7		307	28,149.27	138,886.0	
8		427	3,133.02	11,108.2	
9		461	1,944.53	6,393.1	
10		485	993.65	3,102.3	
11		496	3,757.41	11,474.0	
12		624	795.85	1,933.3	
13		822	819.53	1,510.4	
14		1,216	129.18	161.0	
15		10,380	75.00	10.9	Upper Marker

Region table for sample 4: Bin He #29

From To [bp] Corr.		p] Corr. % of Average Size		Size distribution in	Conc.	Molarity	Co	
[bp]		Area	Total	[bp]	CV [%]	[pg/µl]	[pmol/l]	lor
200	1,000	2,497.5	97	371	36.3	74,732.77	351,685.1	

Electropherogram Summary Continued ...



Overall Results for sample 5 : Bin He #30

Number of peaks found: 12 Corr. Area 1: 3,331.6

Noise: 0.4

Peak t	able	for sample 5:	<u>Bin He #30</u>				
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations		
1		32	0.00	0.0			
2	4	35	125.00	5,411.3	Lower Marker		
3		44	270.62	9,377.7			
4		128	151.49	1,788.4			
5		276	23,456.69	128,684.7			
6		286	17,433.80	92,427.6			
7		339	23,105.23	103,410.1			
8		484	2,508.69	7,859.4			
9		527	1,790.77	5,145.5			
10		566	1,939.37	5,194.0			
11		630	697.99	1,678.3			
12		738	301.56	619.2			
13		791	1,342.89	2,572.3			
14		10,380	75.00	10.9	Upper Marker		

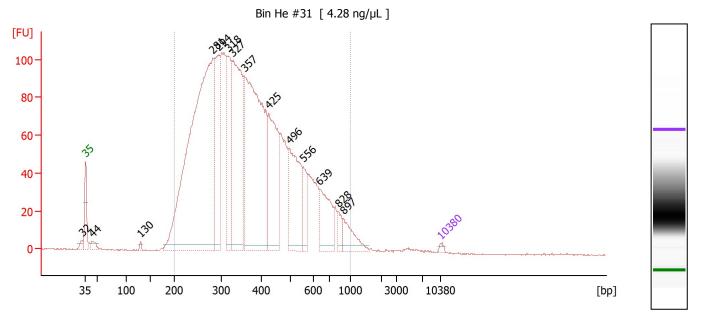
Region table for sample 5: Bin He #30

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	1.000	3.331.6	96	373	36.6	71.060.59	333.633.9	

Page 5 of 11

Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DE13805510_2017-09-15_12-39-33.xad Created: 9/15/2017 12:39:32 PM Modified: 9/15/2017 1:20:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : Bin He #31

Number of peaks found: 14 Corr. Area 1: 2,967.7

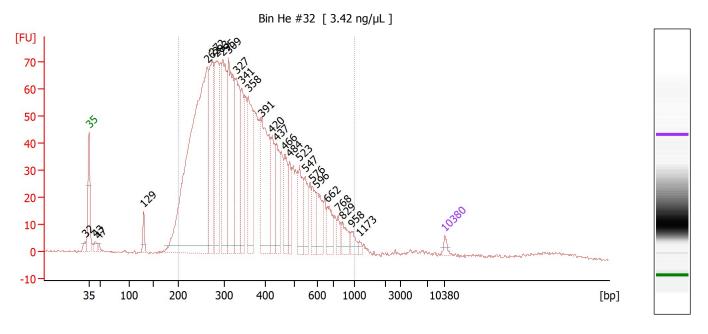
Noise: 0.3

Peak t	able	for sample 6:	Bin He #31		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		32	0.00	0.0	
2	4	35	125.00	5,411.3	Lower Marker
3		44	438.93	15,184.9	
4		130	130.57	1,524.1	
5		281	23,752.42	127,967.7	
6		294	4,971.74	25,592.9	
7		318	3,824.41	18,229.7	
8		327	8,294.45	38,419.7	
9		357	13,815.16	58,555.5	
10		425	5,035.54	17,969.1	
11		496	4,113.50	12,569.6	
12		556	1,198.36	3,263.2	
13		639	2,114.83	5,013.6	
14		828	503.91	921.8	
15		897	1,021.53	1,725.4	
16		10,380	75.00	10.9	Upper Marker

Region table for sample 6: Bin He #31

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	1,000	2,967.7	97	388	36.5	81,092.97	367,771.1	

Electropherogram Summary Continued ...



Overall Results for sample 7: Bin He #32

Number of peaks found: 26 Corr. Area 1: 1,957.9

Noise: 0.3

Peak table	for sample 7:	Bin He #32		
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	32	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	43	170.22	5,984.4	
4	47	94.07	3,023.4	
5	129	326.55	3,821.4	
6	262	10,764.42	62,194.1	
7	272	2,442.44	13,582.2	
8	283	2,411.22	12,923.5	
9	295	2,543.58	13,055.7	
10	309	3,266.96	16,024.1	
11	327	2,235.54	10,347.7	
12	341	1,305.77	5,799.2	
13	358	1,833.54	7,768.5	
14	391	2,366.88	9,183.1	
15	420	956.95	3,452.8	
16	437	881.53	3,059.7	
17	466	821.66	2,671.3	
18	484	546.42	1,712.0	
19	523	908.34	2,631.7	
20	547	570.16	1,579.8	
21	576	459.17	1,207.1	
22	596	585.20	1,488.5	
23	662	450.94	1,032.1	
24	768	177.36	349.9	
25	829	309.39	565.6	
26	958	173.86	274.9	

7 of 11 Page

Assay Class: High Sensitivity DNA Assay

9/15/2017 12:39:32 PM 9/15/2017 1:20:58 PM Created: C:\...gh Sensitivity DNA Assay_DE13805510_2017-09-15_12-39-33.xad Data Path: Modified:

Electropherogram Summary Continued ...

... Peak table for sample 7: Bin He #32

Peak Size [bp] Molarity [pmol/l] **Observations** Conc. [pg/µl] 27 1,173 58.65 75.7

10,380 Upper Marker 28 75.00 10.9

Region table for sample 7: Bin He #32

To [bp] Corr. Size distribution in % of Average Size Conc. Molarity Co [bp] Area Total [bp] CV [%] [pg/µl] [pmol/I] lor 200 1,000 36.9 42,749.18 199,838.9 1,957.9 94 376

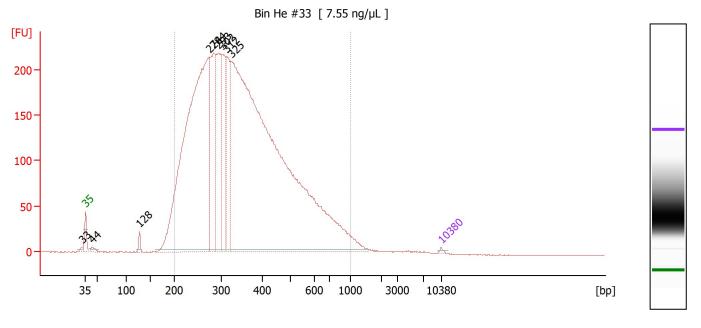
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Page 8 of 11

Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DE13805510_2017-09-15_12-39-33.xad Created: 9/15/2017 12:39:32 PM Modified: 9/15/2017 1:20:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : <u>Bin He #33</u>

Number of peaks found: 9 Corr. Area 1: 6,043.4

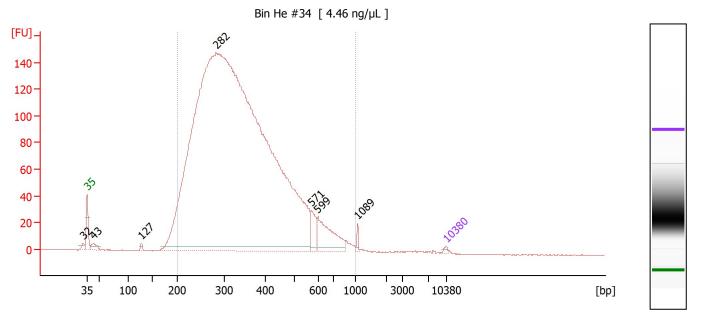
Noise: 0.3

Peak to	able	for sample 8:	Bin He #33		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		33	0.00	0.0	
2	4	35	125.00	5,411.3	Lower Marker
3		44	399.39	13,719.4	
4		128	520.89	6,173.6	
5		274	44,433.22	246,062.7	
6		284	9,196.06	49,091.2	
7		293	9,039.68	46,668.4	
8		302	6,395.69	32,053.1	
9		312	6,516.24	31,624.5	
10		325	79,131.16	368,500.9	
11		10,380	75.00	10.9	Upper Marker

Region table for sample 8: Bin He #33

From	To [bp] C	Corr.	% of	Average Size	Size distribution in CV [%]	Conc.	Molarity	Co
[bp]	A	Area	Total	[bp]		[pg/µl]	[pmol/l]	lor
200	1 000 6	043.4	96	368	36.2	143 522 23	679 469 9	

Electropherogram Summary Continued ...



Overall Results for sample 9: Bin He #34

Number of peaks found: 7 Corr. Area 1: 3,595.5

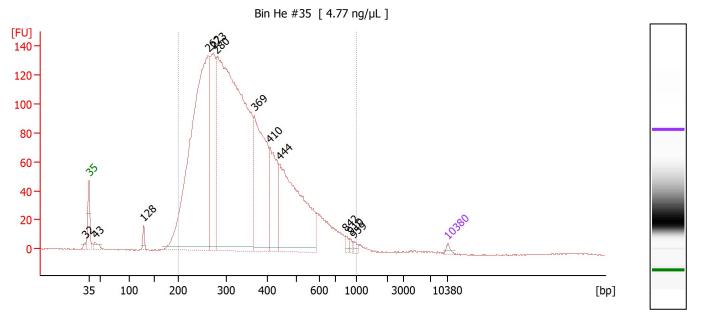
Noise: 0.3

Peak table for sample 9:			Bin He #34		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		32	0.00	0.0	
2	4	35	125.00	5,411.3	Lower Marker
3		43	587.87	20,546.1	
4		127	149.27	1,776.5	
5		282	104,802.54	562,469.6	
6		571	907.22	2,409.2	
7		599	2,276.55	5,761.5	
8		1,089	140.38	195.4	
9		10,380	75.00	10.9	Upper Marker

Region table for sample 9: Bin He #34

From	To [bp]	Corr.	% of	Average Size	Size distribution in	Conc.	Molarity	Co	
[bp]		Area	Total	[bp]	CV [%]	[pg/µl]	[pmol/l]	lor	
200	1 000	3 595 5	97	352	32.6	98 549 20	475 022 9		

Electropherogram Summary Continued ...



Overall Results for sample 10 : Bin He #35

Number of peaks found: 12 Corr. Area 1: 3,268.6

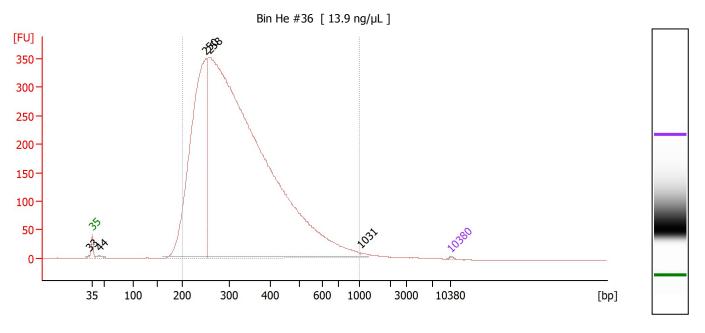
Noise: 0.3

Peak table for sample 10 : <u>Bin He #35</u>									
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations				
1		32	0.00	0.0					
2	4	35	125.00	5,411.3	Lower Marker				
3		43	270.11	9,512.5					
4		128	281.89	3,324.4					
5		262	16,080.52	92,907.4					
6		273	5,041.60	27,986.8					
7		280	22,329.18	120,716.0					
8		369	5,559.53	22,821.3					
9		410	2,543.77	9,406.7					
10		444	6,253.14	21,360.0					
11		842	120.96	217.7					
12		910	95.81	159.5					
13		959	93.14	147.2					
14		10,380	75.00	10.9	Upper Marker				

Region table for sample 10 : Bin He #35

From [bp]	To [bp	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	1,000	3,268.6	97	354	34.2	57,396.20	278,054.5	

Electropherogram Summary Continued ...



Overall Results for sample 11 : Bin He #36

Number of peaks found: 5 Corr. Area 1: 7,908.0

Noise: 0.3

Peak t	:able	for sample 11	: <u>Bin He #36</u>	Bin He #36		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	
1		33	0.00	0.0		
2	4	35	125.00	5,411.3	Lower Marker	
3		44	682.48	23,659.9		
4		250	67,065.69	406,111.3		
5		258	219,523.53	1,288,651.0		
6		1,031	261.49	384.3		
7		10,380	75.00	10.9	Upper Marker	

Region table for sample 11: Bin He #36

From	To [bp]	[bp] Corr. % of Average Size Size distribution in		Conc.	Molarity	Co		
[bp]		Area	Total	[bp]	CV [%]	[pg/µl]	[pmol/l]	lor
200	1.000	7.908.0	97	336	33.2	248.933.28	1.254.674.9	