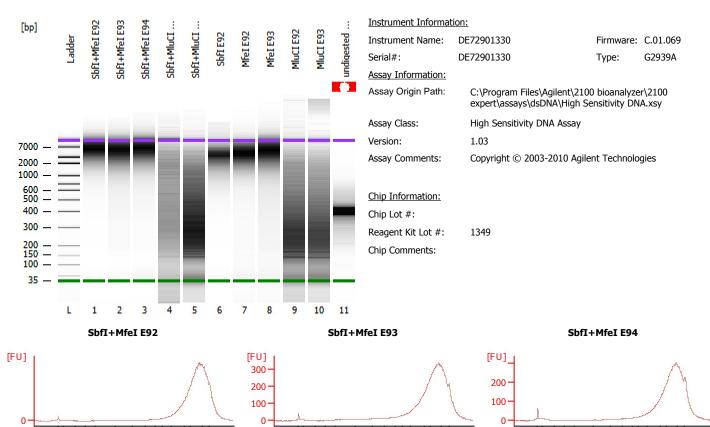
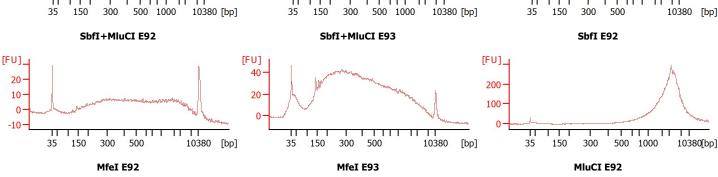
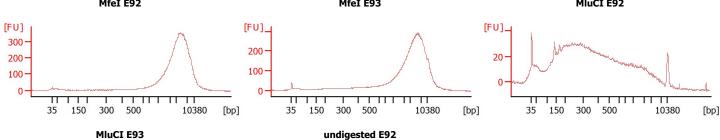
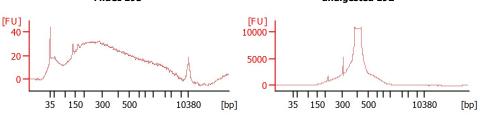
Assay Class: High Sensitivity DNA Assay Created: 4/7/2014 2:22:30 PM Data Path: C:\...ENR\Desktop\2014-04-07_surfclam HS DNA Assay_DE72901330.xad Modified: 4/8/2014 2:53:53 PM

Electrophoresis File Run Summary









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Assay Class: High Sensitivity DNA Assay C:\...ENR\Desktop\2014-04-07_surfclam HS DNA Assay_DE72901330.xad Created: 4/7/2014 2:22:30 PM Modified: 4/8/2014 2:53:53 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name SbfI+MfeI E92 SbfI+MfeI E93 SbfI+MfeI E94 SbfI+MluCI E92 SbfI+MluCI E93 SbfI E92 MfeI E92 MfeI E93 MluCI E93 undigested E92 Ladder	Sample Comment	Rest. Digest	Statu Observation s * * * * * * * * * * * *	Result Label Result Color
Chip Lot #		_		Reagent Kit Lot # 1349

Chip Comments:

Assay Class: High Sensitivity DNA Assay Created: 4/7/2014 2:22:30 PM Data Path: C:\...ENR\Desktop\2014-04-07_surfclam HS DNA Assay_DE72901330.xad Modified: 4/8/2014 2:53:53 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]: 5
Area Threshold: 0.1
Width Threshold [s]: 0.6
Baseline Plateau [s]: 0.5

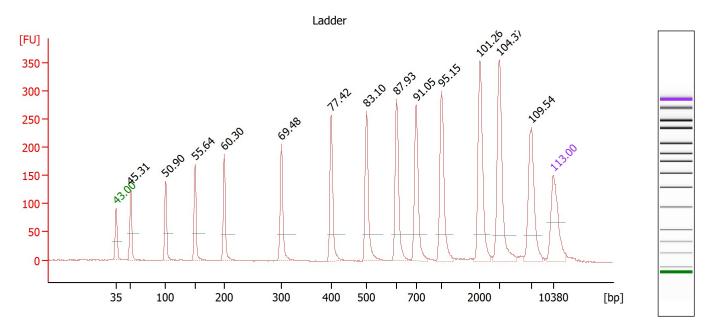
Filter Settings

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

Electropherogram Summary



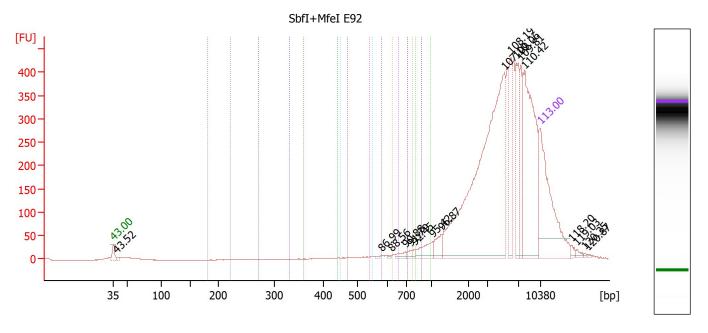
Overall Results for Ladder

Noise: 0.2

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		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		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		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 ...



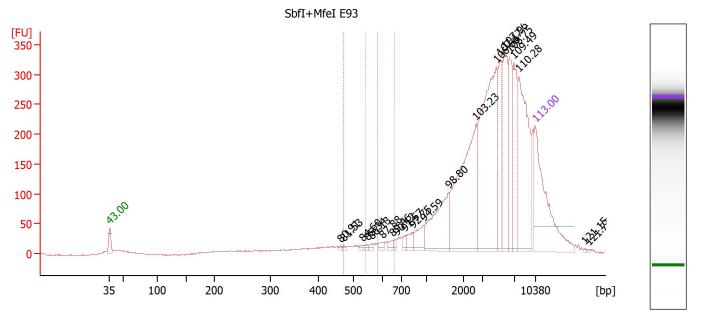
Overall Results for sample 1 : SbfI+MfeI E92

Number of peaks found:	17	Corr. Area 5:	34.8
Noise:	0.3	Corr. Area 6:	43.1
Corr. Area 1:	0.0	Corr. Area 7:	45.5
Corr. Area 2:	0.0	Corr. Area 8:	69.6
Corr. Area 3:	1.6	Corr. Area 9:	21.7
Corr. Area 4:	15.2		

Peak table	for sample 1:	SbfI+MfeI E92		
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	38	2.42	95.3	
3	580	0.95	2.5	
4	620	0.88	2.2	
5	695	4.02	8.8	
6	755	2.52	5.1	
7	803	1.83	3.5	
8	1,045	12.40	18.0	
9	1,281	10.05	11.9	
10	5,109	263.86	78.3	
11	5,955	38.06	9.7	
12	6,632	37.32	8.5	
13	7,262	29.32	6.1	
14	7,860	108.11	20.8	
15	10,380	75.00	10.9	Upper Marker
16	15,464	0.00	0.0	
17	16,275	0.00	0.0	
18	17,557	0.00	0.0	
19	18,069	0.00	0.0	

Regio	on table	e for sam _l	ole 1 :	<u>SbfI+MfeI E</u>	<u>92</u>			
From [bp]	To [b	p] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
180	220	0.0	0	0	0.0	0.00	0.0	
270	330	0.0	0	0	0.0	0.00	0.0	
360	440	1.6	0	425	2.3	0.28	1.0	
450	550	15.2	0	507	5.5	2.68	8.0	
471	581	21.7	1	535	6.0	3.73	10.6	
540	660	34.8	1	602	5.6	5.76	14.6	
630	770	43.1	1	694	5.6	7.21	15.8	
720	880	45.5	1	802	5.9	7.39	14.0	
810	990	69.6	2	906	5.8	10.80	18.1	

Electropherogram Summary Continued ...



Overall Results for sample 2 : SbfI+MfeI E93

 Number of peaks found:
 21
 Corr. Area 1:
 103.0

 Noise:
 0.7
 Corr. Area 2:
 96.3

Peak ta	able	for sample 2:	SbfI+MfeI E93		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2		461	1.18	3.9	
3		472	1.52	4.9	
4		531	1.80	5.1	
5		546	1.77	4.9	
6		563	2.16	5.8	
7		599	2.96	7.5	
8		649	3.76	8.8	
9		670	2.69	6.1	
10		738	3.74	7.7	
11		825	8.24	15.1	
12		959	14.27	22.5	
13		1,598	55.28	52.4	
14		2,632	122.28	70.4	
15		4,724	159.16	51.0	
16		5,165	33.73	9.9	
17		5,776	54.56	14.3	
18		6,387	41.09	9.7	
19		6,964	41.46	9.0	
20		7,725	97.07	19.0	
21		10,380	75.00	10.9	Upper Marker
22		18,346	0.00	0.0	
23		18,945	0.00	0.0	

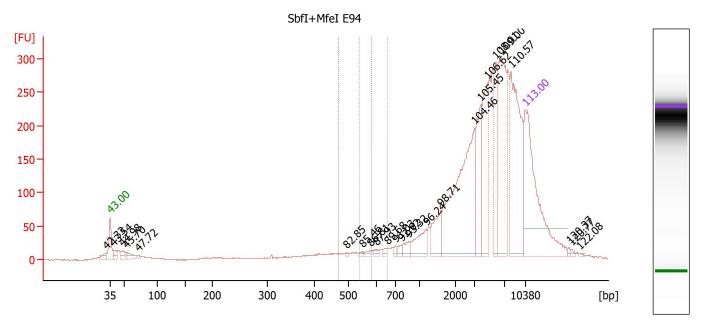
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Assay Class: High Sensitivity DNA Assay C:\...ENR\Desktop\2014-04-07_surfclam HS DNA Assay_DE72901330.xad Created: 4/7/2014 2:22:30 PM Modified: 4/8/2014 2:53:53 PM

Regio	on table for samp	ole 2 :	SbfI+MfeI E	<u> 93</u>		
From	To [bp] Corr.	% of	Average Size	Size distribution in	Conc.	

From [bp]	To [b _l	o] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co <u>lor</u>
471	581	96.3	2	526	6.2	22.43	64.9	
540	660	103.0	2	597	5.8	22.96	58.5	

Electropherogram Summary Continued ...



Overall Results for sample 3: SbfI+MfeI E94

 Number of peaks found:
 24
 Corr. Area 1:
 64.7

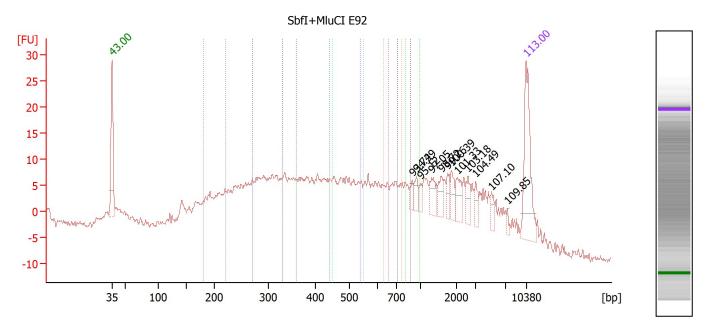
 Noise:
 0.5
 Corr. Area 2:
 57.8

Peak ta	ble 1	for sample 3:	SbfI+MfeI E94		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		33	0.00	0.0	
2	◀ -	35	125.00	5,411.3	Lower Marker
3		39	6.14	241.5	
4		48	6.37	201.5	
5		53	8.94	253.2	
6		72	1.86	39.4	
7		496	0.77	2.4	
8		549	1.12	3.1	
9		577	2.20	5.8	
10		600	2.04	5.2	
11		656	2.22	5.1	
12		699	1.78	3.9	
13		771	2.97	5.8	
14		866	4.87	8.5	
15		1,179	14.96	19.2	
16		1,584	17.50	16.7	
17		3,070	99.23	49.0	
18		3,835	32.63	12.9	
19		4,739	41.06	13.1	
20		5,817	23.59	6.1	
21		6,583	66.21	15.2	
22		8,010	74.76	14.1	
23		10,380	75.00	10.9	Upper Marker
24		17,579	0.00	0.0	
25		17,974	0.00	0.0	
26		19,247	0.00	0.0	

Assay Class: Data Path: High Sensitivity DNA Assay C:\...ENR\Desktop\2014-04-07_surfclam HS DNA Assay_DE72901330.xad 4/7/2014 2:22:30 PM 4/8/2014 2:53:53 PM Created: Modified:

Region table for sample 3:			ple 3:	SbfI+MfeI I	<u> 94</u>			
From [bp]	To [b	p] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
471	581	57.8	2	528	6.2	11.08	32.0	
540	660	64 7	2	598	5.8	11 87	30.2	

Electropherogram Summary Continued ...



Overall Results for sample 4: SbfI+MluCI E92

Number of peaks found:	12	Corr. Area 4:	77.8
Noise:	0.6	Corr. Area 5:	65.9
Corr. Area 1:	45.1	Corr. Area 6:	42.3
Corr. Area 2:	86.6	Corr. Area 7:	28.6
Corr. Area 3:	85.7	Corr. Area 8:	33.3

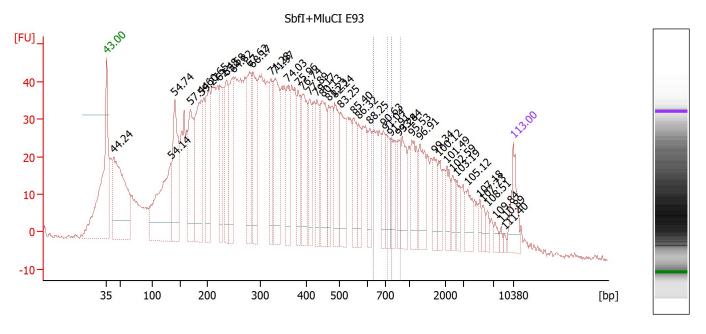
Peak ta	able	for sample 4:	SbfI+MluCI E92		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2		896	9.37	15.9	
3		952	11.97	19.1	
4		998	7.73	11.7	
5		1,312	20.48	23.6	
6		1,585	16.00	15.3	
7		1,755	11.42	9.9	
8		1,858	15.51	12.6	
9		2,025	19.75	14.8	
10		2,617	14.58	8.4	
11		3,088	11.49	5.6	
12		5,110	10.81	3.2	
13		7,299	4.16	0.9	
14		10,380	75.00	10.9	Upper Marker

Region table for sample 4: SbfI+MluCI E92

			_					
From [bp]	To [b	p] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
180	220	45.1	5	200	5.8	163.11	1,238.4	
270	330	86.6	10	299	5.8	280.11	1,426.0	
360	440	85.7	10	397	5.8	249.03	955.3	
450	550	77.8	9	497	5.8	219.91	673.5	
540	660	65.9	8	592	5.8	175.28	450.3	

Regi	on table	e for sam _l	ple 4 :	SbfI+MluCI	E92			
From [bp]	To [b	p] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
630	770	42.3	5	685	5.6	113.30	251.5	
720	880	28.6	3	797	5.8	74.50	142.3	
810	990	33.3	4	898	5.9	82.96	140.5	

Electropherogram Summary Continued ...



Overall Results for sample 5 : SbfI+MluCI E93

 Number of peaks found:
 44
 Corr. Area 1:
 72.2

 Noise:
 0.6
 Corr. Area 2:
 109.8

for sample 5:	SbfI+MluCI E93		
Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
35	125.00	5,411.3	Lower Marker
43	680.06	23,934.8	
134	513.10	5,794.9	
140	369.28	3,984.0	
169	306.55	2,743.2	
189	389.60	3,123.1	
204	273.33	2,031.8	
224	190.41	1,289.1	
236	215.46	1,384.6	
249	252.49	1,534.8	
280	243.35	1,318.1	
286	249.18	1,321.5	
323	144.57	678.8	
331	269.59	1,232.7	
357	167.30	709.4	
382	125.56	498.6	
391	196.94	762.4	
408	122.67	455.4	
431	103.53	364.2	
448	210.51	712.5	
467	97.39	315.9	
485	154.65	483.3	
503	178.12	536.5	
548	129.45	358.1	
567	74.99	200.5	
610	74.50	185.0	
	35 43 134 140 169 189 204 224 236 249 280 286 323 331 357 382 391 408 431 448 467 485 503 548 567	Size [bp] Conc. [pg/μl] 35 125.00 43 680.06 134 513.10 140 369.28 169 306.55 189 389.60 204 273.33 224 190.41 236 215.46 249 252.49 280 243.35 286 249.18 323 144.57 331 269.59 357 167.30 382 125.56 391 196.94 408 122.67 431 103.53 448 210.51 467 97.39 485 154.65 503 178.12 548 129.45 567 74.99	Size [bp] Conc. [pg/μl] Molarity [pmol/l] 35 125.00 5,411.3 43 680.06 23,934.8 134 513.10 5,794.9 140 369.28 3,984.0 169 306.55 2,743.2 189 389.60 3,123.1 204 273.33 2,031.8 224 190.41 1,289.1 236 215.46 1,384.6 249 252.49 1,534.8 280 243.35 1,318.1 286 249.18 1,321.5 323 144.57 678.8 331 269.59 1,232.7 357 167.30 709.4 382 125.56 498.6 391 196.94 762.4 408 122.67 455.4 431 103.53 364.2 448 210.51 712.5 467 97.39 315.9 485 154.65 483.3

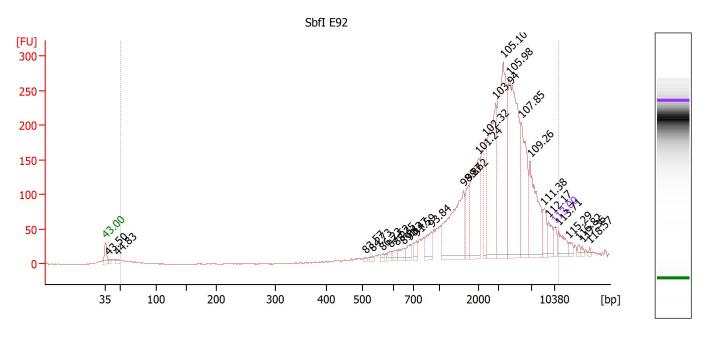
Electropherogram Summary Continued ...

Peak table for sample 5: <u>SbfI+MluCI E93</u>						
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	
27		687	68.52	151.2		
28		700	92.04	199.3		
29		763	85.68	170.1		
30		861	58.13	102.3		
31		904	105.74	177.2		
32		1,063	109.93	156.6		
33		1,288	107.91	126.9		
34		1,686	54.76	49.2		
35		1,814	45.18	37.7		
36		2,077	49.49	36.1		
37		2,429	36.36	22.7		
38		2,621	37.29	21.6		
39		3,575	53.88	22.8		
40		5,171	18.97	5.6		
41		5,597	20.91	5.7		
42		6,200	16.83	4.1		
43		7,290	14.84	3.1		
44		8,320	8.56	1.6		
45		8,812	6.25	1.1		
46		10,380	75.00	10.9	Upper Marker	

Region table for sample 5 : <u>SbfI+MluCI E93</u>

From [bp]	To [b	p] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
630	770	109.8	4	684	5.5	356.92	792.5	
720	880	72.2	3	796	6.0	228.62	436.4	

Electropherogram Summary Continued ...



Overall Results for sample 6 : SbfI E92

Number of peaks found: 27 Corr. Area 1: 2,621.5

Noise: 0.3

Peak table	e for sample 6:	SbfI E92		
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	38	68.10	2,697.9	
3	47	86.76	2,803.2	
4	510	17.32	51.5	
5	534	31.41	89.2	
6	565	30.78	82.6	
7	591	53.68	137.5	
8	613	65.47	161.7	
9	648	114.16	266.9	
10	681	89.29	198.5	
11	740	79.88	163.6	
12	904	215.82	361.7	
13	1,609	1,037.23	976.7	
14	1,732	248.05	217.1	
15	1,997	809.38	614.1	
16	2,341	355.36	230.0	
17	2,861	1,091.46	578.1	
18	3,565	1,511.38	642.4	
19	4,240	1,636.54	584.8	
20	5,688	786.57	209.5	
21	6,782	712.45	159.2	
22	8,796	175.31	30.2	
23	9,568	137.50	21.8	
24	10,380	75.00	10.9	Upper Marker
25	11,070	0.00	0.0	
26	12,614	0.00	0.0	

Observations

Electropherogram Summary Continued ...

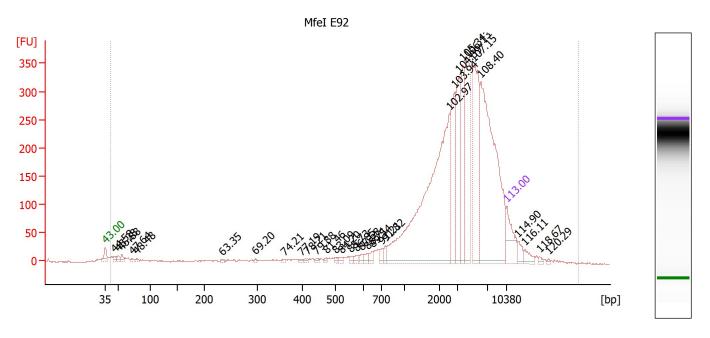
Peak ta	ble for sample 6	:	SbfI E92
Dook	Ciza [hn]	Conc	[na/ul]

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]
27	14,116	0.00	0.0
28	14,644	0.00	0.0
29	15,822	0.00	0.0

Region table for sample 6: SbfI E92

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
50	11.000 2.621.5	97	3,739	60.7	9,590.07	8,392,4	

Electropherogram Summary Continued ...



Overall Results for sample 7: MfeI E92

Number of peaks found: 32 Corr. Area 1: 4,435.0

Noise: 1.0

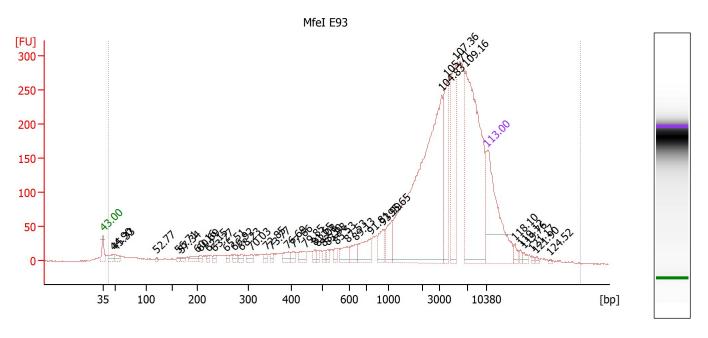
Peak table	for sample 7:	MfeI E92		
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	45	16.71	559.3	
3	50	17.84	542.7	
4	55	17.68	485.9	
5	71	10.11	216.1	
6	78	9.07	175.3	
7	233	5.28	34.3	
8	297	4.10	20.9	
9	360	4.82	20.3	
10	397	4.94	18.9	
11	414	6.73	24.6	
12	443	6.09	20.8	
13	471	6.91	22.2	
14	500	6.12	18.5	
15	523	9.06	26.3	
16	554	7.70	21.0	
17	578	10.59	27.8	
18	595	10.01	25.5	
19	625	13.59	32.9	
20	652	14.26	33.2	
21	712	16.03	34.1	
22	756	15.38	30.8	
23	2,549	788.99	469.0	
24	2,862	157.85	83.6	
25	3,170	173.43	82.9	
26	3,745	157.05	63.5	

Peak table for sample 7: <u>MfeI E92</u>							
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations			
27	4,357	212.18	73.8				
28	5,148	251.63	74.1				
29	6,118	613.72	152.0				
30	10,380	75.00	10.9	Upper Marker			
31	12,241	0.00	0.0				
32	13,421	0.00	0.0				
33	15,917	0.00	0.0				
34	17,505	0.00	0.0				

Region	table	for	sample :	7	:	MfeI E92
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From	To [bp] Corr.	% of	Average Size	Size distribution in	Conc.	Molarity	Co
[bp]	Area	Total	[bp]	CV [%]	[pg/µl]	[pmol/l]	lor
41	22,660 4,435.0	99	4,483	71.8	2,982.51	5,778.7	

Electropherogram Summary Continued ...



Overall Results for sample 8: MfeI E93

Number of peaks found: 38 Corr. Area 1: 4,224.9

Noise: 0.7

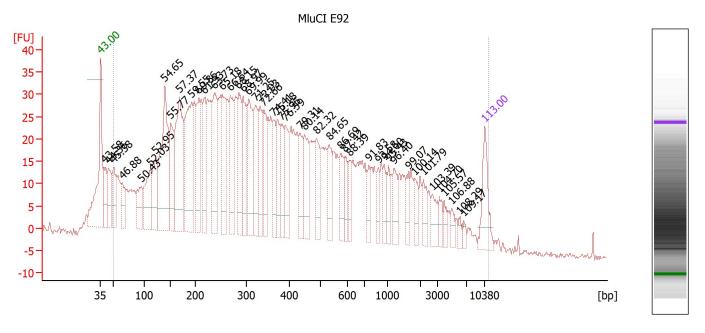
Peak table for sample 8: MfeI E93							
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations			
1	35	125.00	5,411.3	Lower Marker			
2	47	10.50	336.1				
3	50	8.30	250.3				
4	120	2.23	28.2				
5	161	2.37	22.2				
6	168	3.52	31.7				
7	199	8.02	61.2				
8	204	3.50	26.0				
9	220	3.29	22.7				
10	232	3.48	22.7				
11	257	3.49	20.6				
12	272	3.96	22.1				
13	286	4.09	21.7				
14	307	5.71	28.2				
15	342	3.73	16.5				
16	354	3.15	13.5				
17	391	7.41	28.7				
18	404	4.53	17.0				
19	443	8.40	28.8				
20	474	4.24	13.5				
21	482	3.76	11.8				
22	504	3.64	10.9				
23	518	3.74	10.9				
24	550	5.08	14.0				
25	592	10.48	26.8				
26	639	10.12	24.0				

Peal	k tab	ole for sample 8	: <u>MfeI E93</u>		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
27		755	22.56	45.3	
28		916	13.88	23.0	
29		1,082	17.15	24.0	
30		3,356	254.31	114.8	
31		4,034	43.88	16.5	
32		5,313	61.17	17.4	
33		6,705	168.69	38.1	
34		10,380	75.00	10.9	Upper Marker
35		15,367	0.00	0.0	
36		16,364	0.00	0.0	
37		16,982	0.00	0.0	
38		18,359	0.00	0.0	
39		19,072	0.00	0.0	
40		21,636	0.00	0.0	

Region table for	r sample 8:	<u>MfeI E93</u>
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	To [bp] Corr.	% of		Size distribution in		Molarity	Co
[bp]	Area	Total	[bp]	CV [%]	[pg/µl]	[pmol/l]	lor
41	27,131 4,224.9	99	5,041	81.7	957.04	2,436.3	

Electropherogram Summary Continued ...



Overall Results for sample 9: MluCI E92

Number of peaks found: 47 Corr. Area 1: 1,895.8

Noise: 0.7

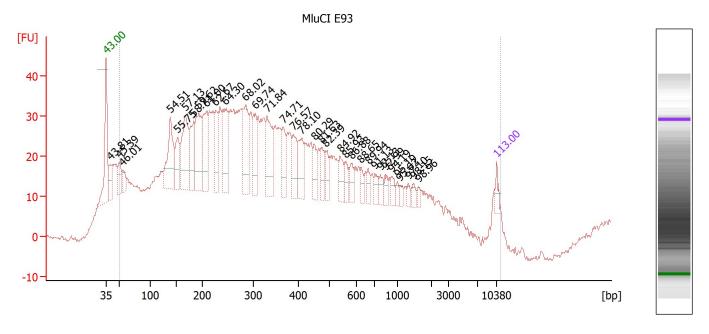
for sample 9:	MluCI E92		
Size [bp]	Conc. [pg/µl]	Molarity [pmol/I]	Observations
35	125.00	5,411.3	Lower Marker
39	117.30	4,582.0	
44	117.55	4,062.6	
51	96.63	2,890.7	
64	79.12	1,870.4	
96	106.10	1,678.7	
112	166.67	2,256.9	
122	141.23	1,759.3	
140	427.58	4,642.6	
151	164.03	1,642.1	
169	368.97	3,316.8	
192	330.16	2,605.3	
206	163.29	1,200.0	
218	140.39	976.6	
237	216.05	1,379.1	
253	179.64	1,074.9	
269	171.38	965.0	
285	149.76	794.8	
294	149.72	770.4	
306	181.45	897.2	
322	134.86	633.9	
332	94.19	429.7	
340	103.35	460.5	
372	83.76	341.3	
382	74.23	294.8	
	Size [bp] 35 39 44 51 64 96 112 122 140 151 169 192 206 218 237 253 269 285 294 306 322 332 340 362	Size [bp] Conc. [pg/μl] 35 125.00 39 117.30 44 117.55 51 96.63 64 79.12 96 106.10 112 166.67 122 141.23 140 427.58 151 164.03 169 368.97 192 330.16 206 163.29 218 140.39 237 216.05 253 179.64 269 171.38 285 149.76 294 149.72 306 181.45 322 134.86 332 94.19 340 103.35 362 80.11 372 83.76	Size [bp] Conc. [pg/μl] Molarity [pmol/l] 35 125.00 5,411.3 39 117.30 4,582.0 44 117.55 4,062.6 51 96.63 2,890.7 64 79.12 1,870.4 96 106.10 1,678.7 112 166.67 2,256.9 122 141.23 1,759.3 140 427.58 4,642.6 151 164.03 1,642.1 169 368.97 3,316.8 192 330.16 2,605.3 206 163.29 1,200.0 218 140.39 976.6 237 216.05 1,379.1 253 179.64 1,074.9 269 171.38 965.0 285 149.76 794.8 294 149.72 770.4 306 181.45 897.2 322 134.86 633.9 332 94.19 429.7 </td

Pea	ak tal	ole for sample	9: MluCI E	<u>92</u>	
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
27		390	103.64	403.1	
28		433	83.67	292.6	
29		448	99.70	337.4	
30		486	84.44	263.1	
31		532	79.61	226.7	
32		574	68.89	181.7	
33		589	42.33	108.8	
34		615	43.82	108.0	
35		758	36.61	73.2	
36		885	33.55	57.4	
37		960	37.58	59.3	
38		1,046	30.57	44.3	
39		1,205	49.97	62.8	
40		1,642	26.50	24.5	
41		1,817	30.68	25.6	
42		2,170	34.47	24.1	
43		2,684	34.70	19.6	
44		3,253	20.69	9.6	
45		3,929	16.98	6.5	
46		4,943	16.29	5.0	
47		6,033	11.13	2.8	
48		6,709	10.49	2.4	
49		10,380	75.00	10.9	Upper Marker

Region table for sample 9: <u>MluCI</u>

From	To [bp] Corr.	% of		Size distribution in		Molarity	Co
[bp]	Area	Total	[bp]	CV [%]	[pg/µl]	[pmol/l]	lor
50	11,000 1,895.8	96	886	100.0	7,328.20	52,001.9	

Electropherogram Summary Continued ...



Overall Results for sample 10 : MluCI E93

Number of peaks found: 34 Corr. Area 1: 1,730.0

Noise: 0.5

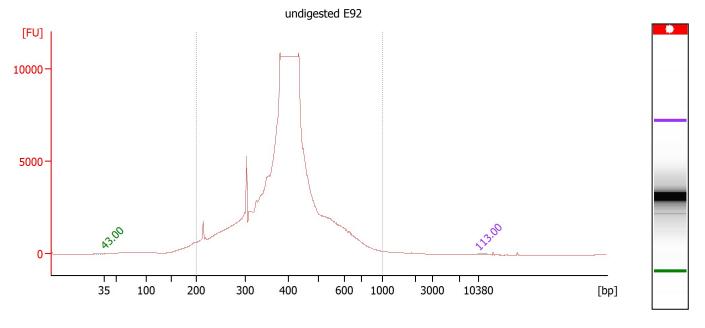
Peak table for sample 10 : MluCI E93							
Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations				
35	125.00	5,411.3	Lower Marker				
40	355.77	13,382.7					
51	268.49	8,019.7					
56	181.55	4,888.2					
138	957.44	10,509.3					
151	459.42	4,605.3					
166	1,051.09	9,592.3					
183	561.88	4,653.4					
193	877.44	6,899.4					
208	717.36	5,233.9					
226	419.59	2,814.6					
244	757.24	4,710.9					
284	793.84	4,233.5					
303	444.33	2,219.9					
	635.00	2,918.0					
366	323.56	1,340.3					
389	344.01	1,339.1					
412	351.11	1,291.8					
450	224.00	753.4					
474	178.53	570.6					
	151.24	470.1					
538	201.13	566.7					
		393.6					
623	102.16	248.4					
664	105.67	241.0					
	Size [bp] 35 40 51 56 138 151 166 183 193 208 226 244 284 303 330 366 389 412 450 474 487 538 558 578 623	Size [bp] Conc. [pg/μl] 35 125.00 40 355.77 51 268.49 56 181.55 138 957.44 151 459.42 166 1,051.09 183 561.88 193 877.44 208 717.36 226 419.59 244 757.24 284 793.84 303 444.33 330 635.00 366 323.56 389 344.01 412 351.11 450 224.00 474 178.53 487 151.24 538 201.13 558 141.83 578 150.22 623 102.16	Size [bp] Conc. [pg/μl] Molarity [pmol/l] 35 125.00 5,411.3 40 355.77 13,382.7 51 268.49 8,019.7 56 181.55 4,888.2 138 957.44 10,509.3 151 459.42 4,605.3 166 1,051.09 9,592.3 183 561.88 4,653.4 193 877.44 6,899.4 208 717.36 5,233.9 226 419.59 2,814.6 244 757.24 4,710.9 284 793.84 4,233.5 303 444.33 2,219.9 330 635.00 2,918.0 366 323.56 1,340.3 389 344.01 1,339.1 412 351.11 1,291.8 450 224.00 753.4 474 178.53 570.6 487 151.24 470.1 538 141.83 38				

Peak	table for sa	mple 10 : <u>Mlu</u>	<u>CI E93</u>	
Peak	Size [bp	o] Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	706	113.95	244.5	
28	779	98.38	191.2	
29	842	88.28	158.8	
30	930	95.39	155.5	
31	1,007	50.11	75.4	
32	1,077	49.41	69.5	
33	1,320	50.24	57.7	
34	1,492	65.99	67.0	
35	1,625	41.34	38.6	
36	10,380	75.00	10.9	Upper Marker

Region tab	le f	or samp	le 10 :	M	<u>luCI E93</u>
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From	To [bp] Corr.	% of	Average Size	Size distribution in	Conc.	Molarity	Co
[bp]	Area	Total	[bp]	CV [%]	[pg/µl]	[pmol/l]	lor
50	11,000 1,730.0	95	560	100.0	33,880.71	265,990.5	

Electropherogram Summary Continued ...



Overall Results for sample 11: undigested E92

Number of peaks found: 0 Corr. Area 1: 140,160.1

Noise: 1.5

Peak table for sample 11: undigested E92

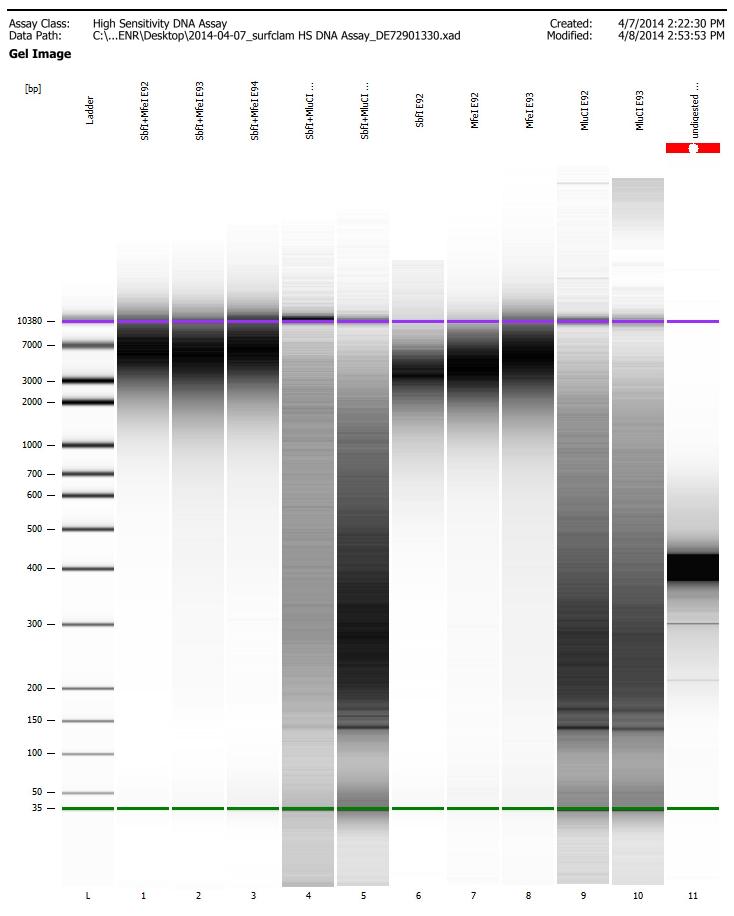
 Peak
 Size [bp]
 Conc. [pg/μl]
 Molarity [pmol/l]
 Observations

 1
 %
 35
 125.00
 5,411.3
 Lower Marker

 2
 %
 10,380
 75.00
 10.9
 Upper Marker

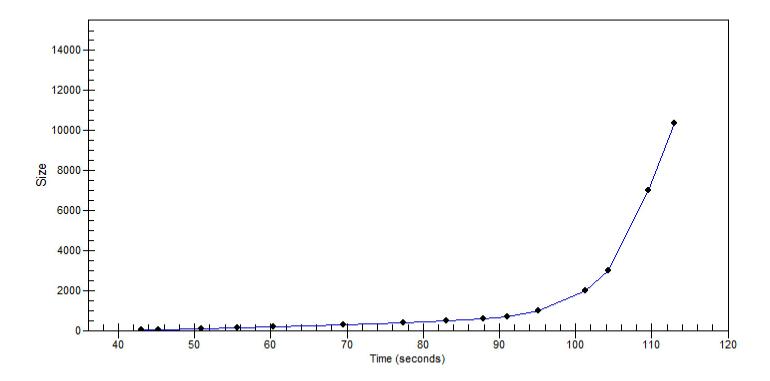
Region table for sample 11: <u>undigested E92</u>

From To [bp] Corr. % of **Average Size** Size distribution in Conc. [pg/µl] Molarity Co **Total** [bp] Area [bp] CV [%] [pmol/l] lor 200 1,000 140,160.1 96 120,186,832.0 484,540,768.0 408 24.5



Curves

Standard Curve



Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 12)		Instrument	Run		4/7/2014 3:01:52 PM	(GMT04:00) Eastern Standard Time	foran103	RUTGERS-4341 E48
Instrument error occurred on port 1, Optical signal too high (1605h)	559	Instrument	Run	Sample 11	4/7/2014 3:01:52 PM	(GMT04:00) Eastern Standard Time	foran103	RUTGERS-4341 E48
Run started on port 1 (File: C:\Program Files\Agilent\21 00 bioanalyzer\210 0 expert\Data\20: 4-04-07\Pinsky_High Sensitivity DNA Assay_DE72901 330_2014-04-0: _14-19-53.xad)) 1 -	Instrument	Run		4/7/2014 2:22:30 PM	(GMT04:00) Eastern Standard Time	foran103	RUTGERS-4341 E48
Product Number: G2939A	r	Instrument	Run		4/7/2014 2:22:30 PM	(GMT04:00) Eastern Standard Time	foran103	RUTGERS-4341 E48
Name :		Instrument	Run		4/7/2014 2:22:30 PM	(GMT04:00) Eastern Standard Time	foran103	RUTGERS-4341 E48
Vendor : Agilent Technologies	t	Instrument	Run		4/7/2014 2:22:30 PM	(GMT04:00) Eastern Standard Time	foran103	RUTGERS-4341 E48
Serial#: DE72901330		Instrument	Run		4/7/2014 2:22:30 PM	(GMT04:00) Eastern Standard Time	foran103	RUTGERS-4341 E48
Firmware : C.01.069		Instrument	Run		4/7/2014 2:22:30 PM	(GMT04:00) Eastern Standard Time	foran103	RUTGERS-4341 E48
Cartridge : Electrode		Instrument	Run		4/7/2014 2:22:30 PM	(GMT04:00) Eastern Standard Time	foran103	RUTGERS-4341 E48