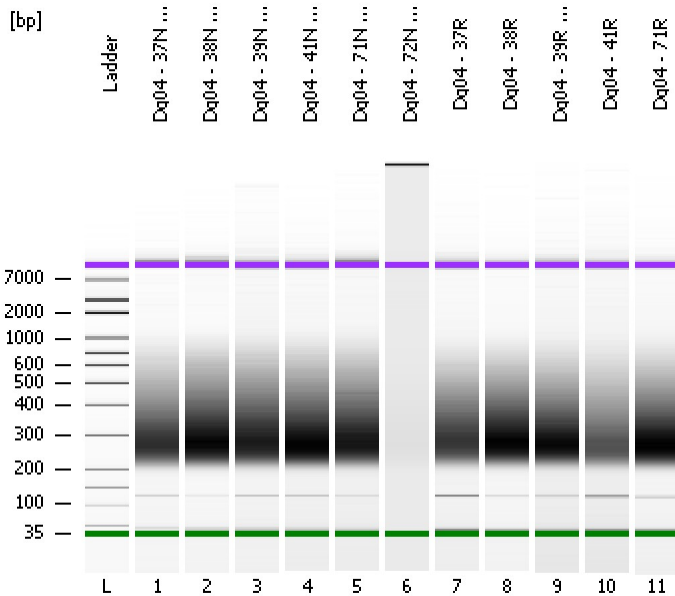


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
Modified: 4/2/2015 3:16:58 PM

Electrophoresis File Run Summary



Instrument Information:

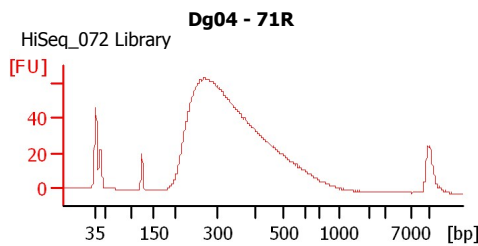
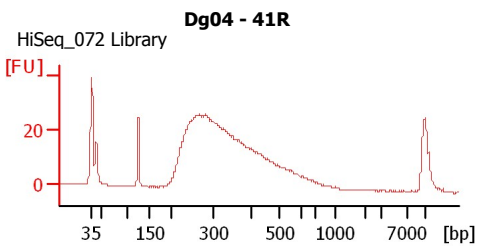
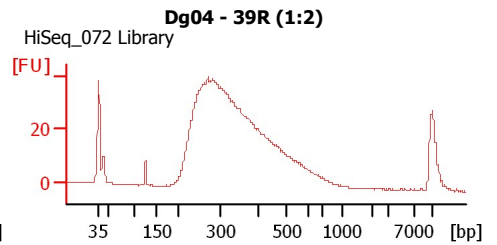
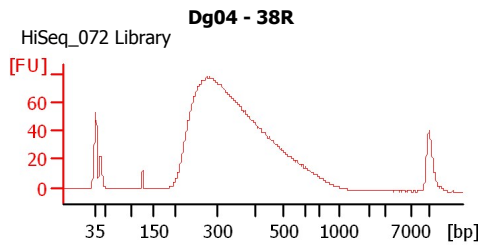
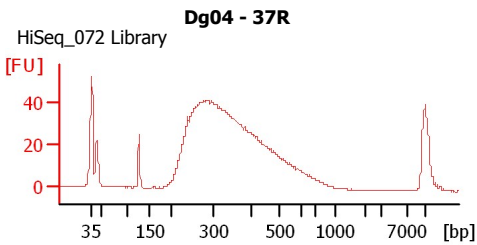
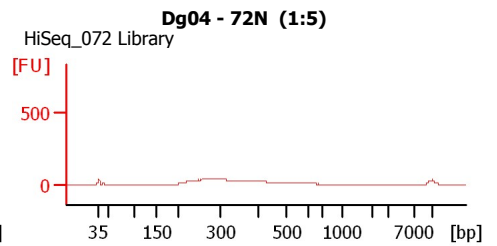
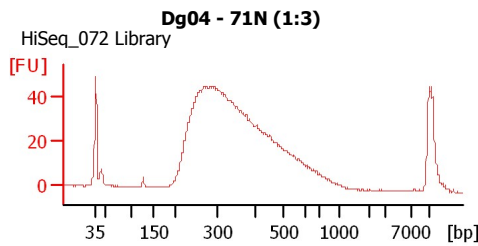
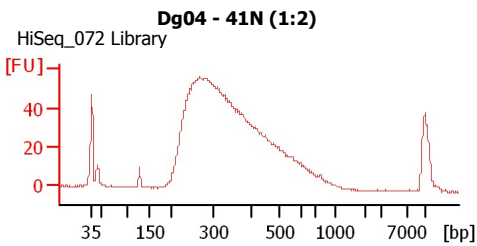
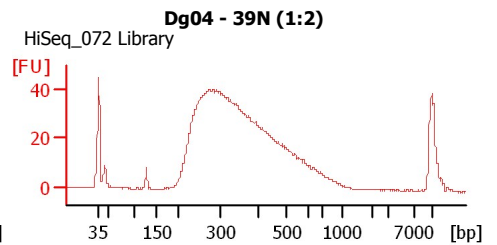
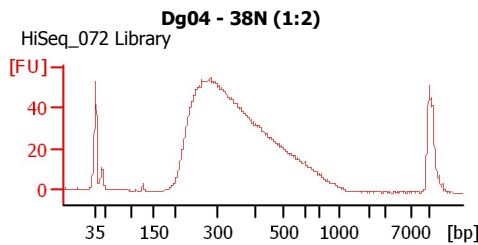
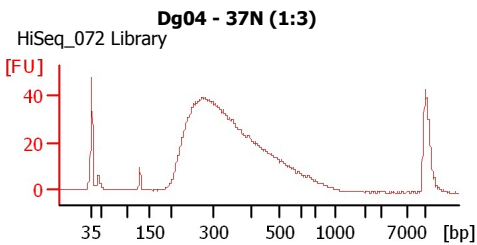
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

Assay Information:

Assay Origin Path: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\assays\dsDNA\High Sensitivity DNA.xsy
Assay Class: High Sensitivity DNA Assay
Version: 1.03
Assay Comments: Copyright © 2003-2010 Agilent Technologies

Chip Information:

Chip Lot #:
Reagent Kit Lot #:
Chip Comments:



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
Modified: 4/2/2015 3:16:58 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Dg04 - 37N (1:3)	HiSeq_072 Library	<input type="checkbox"/>	✓			
Dg04 - 38N (1:2)	HiSeq_072 Library	<input type="checkbox"/>	✓			
Dg04 - 39N (1:2)	HiSeq_072 Library	<input type="checkbox"/>	✓			
Dg04 - 41N (1:2)	HiSeq_072 Library	<input type="checkbox"/>	✓			
Dg04 - 71N (1:3)	HiSeq_072 Library	<input type="checkbox"/>	✓			
Dg04 - 72N (1:5)	HiSeq_072 Library	<input type="checkbox"/>	✓			
Dg04 - 37R	HiSeq_072 Library	<input type="checkbox"/>	✓			
Dg04 - 38R	HiSeq_072 Library	<input type="checkbox"/>	✓			
Dg04 - 39R (1:2)	HiSeq_072 Library	<input type="checkbox"/>	✓			
Dg04 - 41R	HiSeq_072 Library	<input type="checkbox"/>	✓			
Dg04 - 71R	HiSeq_072 Library	<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
Modified: 4/2/2015 3:16:58 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/ μ l] : 1950
Uses Standard Area for Ladder Fragments
Lower Marker Concentration [pg/ μ l] : 125
Upper Marker Concentration [pg/ μ l] : 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

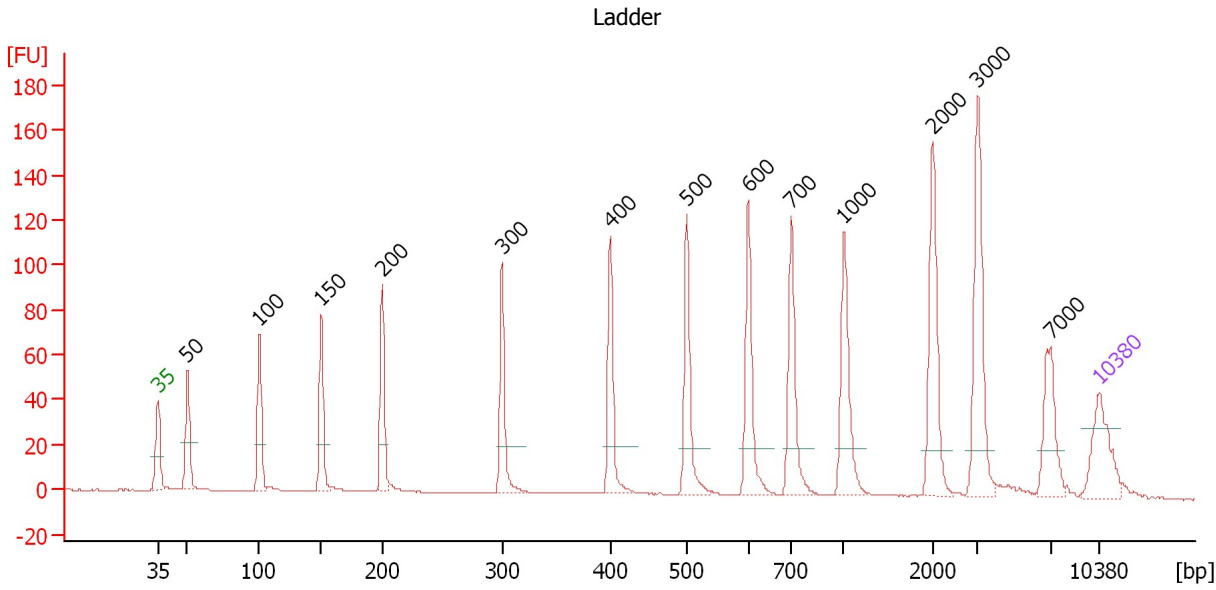
Ladder

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

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
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Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

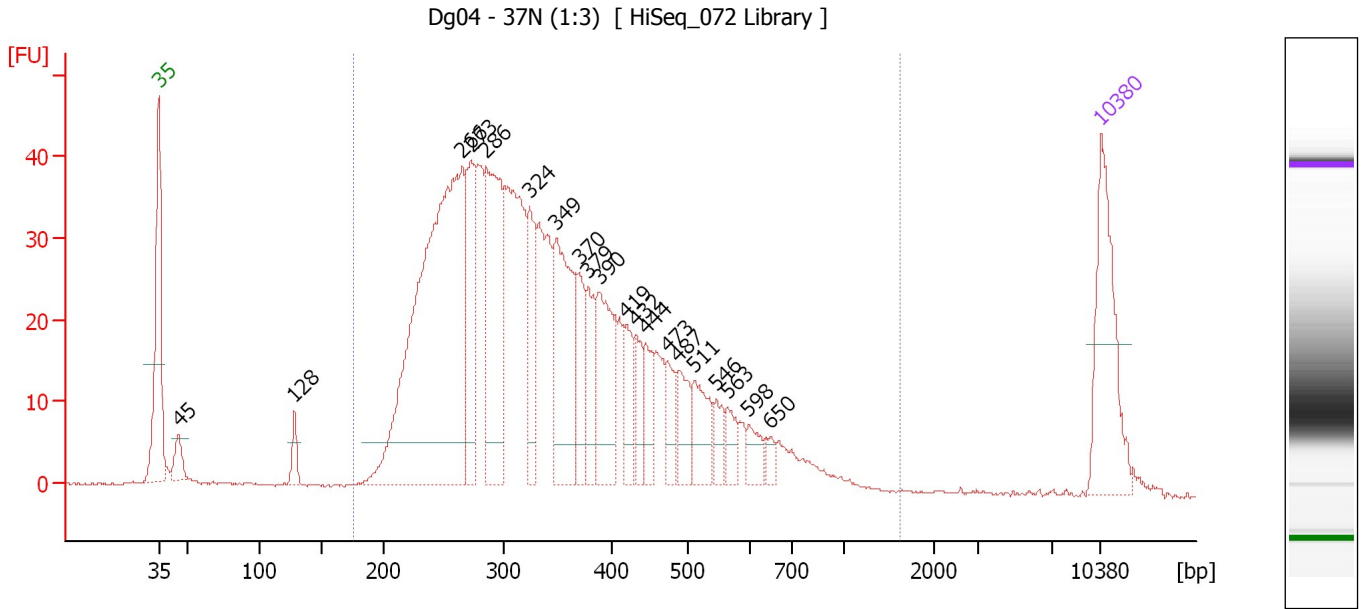
Peak table for Ladder

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	50	150.00	4,545.5	Ladder Peak	45.21
3	100	150.00	2,272.7	Ladder Peak	50.55
4	150	150.00	1,515.2	Ladder Peak	55.16
5	200	150.00	1,136.4	Ladder Peak	59.72
6	300	150.00	757.6	Ladder Peak	68.61
7	400	150.00	568.2	Ladder Peak	76.62
8	500	150.00	454.5	Ladder Peak	82.33
9	600	150.00	378.8	Ladder Peak	86.89
10	700	150.00	324.7	Ladder Peak	90.07
11	1,000	150.00	227.3	Ladder Peak	93.98
12	2,000	150.00	113.6	Ladder Peak	100.57
13	3,000	150.00	75.8	Ladder Peak	103.93
14	7,000	150.00	32.5	Ladder Peak	109.32
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 1 : Dg04 - 37N (1:3)

Number of peaks found: 20 Corr. Area 1: 1,003.4
 Noise: 0.2

Peak table for sample 1 : Dg04 - 37N (1:3)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	45	22.75	765.2		44.48
3	128	17.33	205.9		53.09
4	266	493.57	2,815.8		65.55
5	273	80.64	447.7		66.19
6	286	161.13	853.9		67.35
7	324	59.78	279.4		70.55
8	349	122.00	529.6		72.54
9	370	41.20	168.5		74.25
10	379	45.47	182.0		74.90
11	390	76.09	296.0		75.78
12	419	30.48	110.3		77.68
13	432	24.25	85.0		78.46
14	444	28.02	95.5		79.16
15	473	20.73	66.4		80.78
16	487	30.65	95.3		81.61
17	511	38.46	114.1		82.81
18	546	16.52	45.8		84.44
19	563	14.92	40.1		85.22
20	598	15.75	39.9		86.80
21	650	8.34	19.5		88.46
22	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
Modified: 4/2/2015 3:16:58 PM

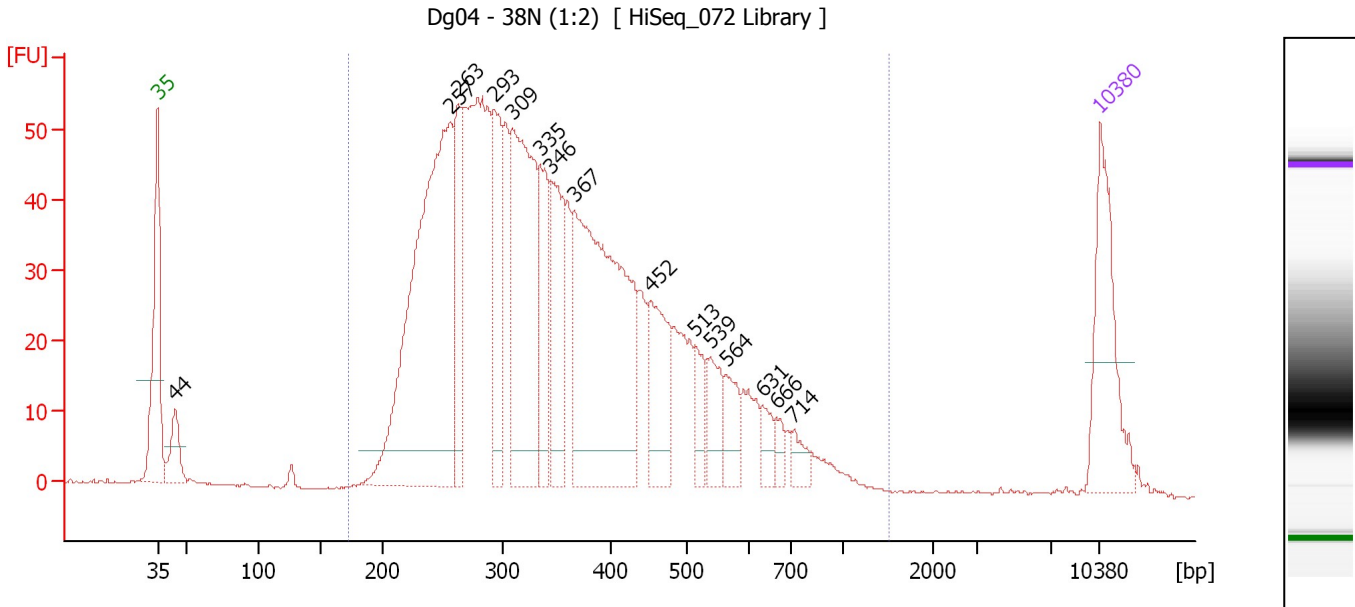
Electropherogram Summary Continued ...**... Region table for sample 1 : Dg04 - 37N (1:3)**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Co lor
176	362	1,622	1,003.4	9,080.1	97	37.0	1,896.53	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
 Modified: 4/2/2015 3:16:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : Dg04 - 38N (1:2)

Number of peaks found: 15 Corr. Area 1: 1,445.6
 Noise: 0.1

Peak table for sample 2 : Dg04 - 38N (1:2)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	44	41.20	1,419.6		44.32
3	257	456.39	2,693.4		64.76
4	263	84.87	488.8		65.33
5	293	108.00	558.8		67.97
6	309	232.45	1,139.2		69.34
7	335	68.91	311.6		71.42
8	346	96.90	424.7		72.27
9	367	329.74	1,361.8		73.96
10	452	77.65	260.3		79.58
11	513	25.90	76.5		82.93
12	539	34.68	97.5		84.11
13	564	28.98	77.9		85.25
14	631	15.80	37.9		87.89
15	666	9.13	20.8		88.97
16	714	14.37	30.5		90.25
17	10,380	75.00	10.9	Upper Marker	113.00

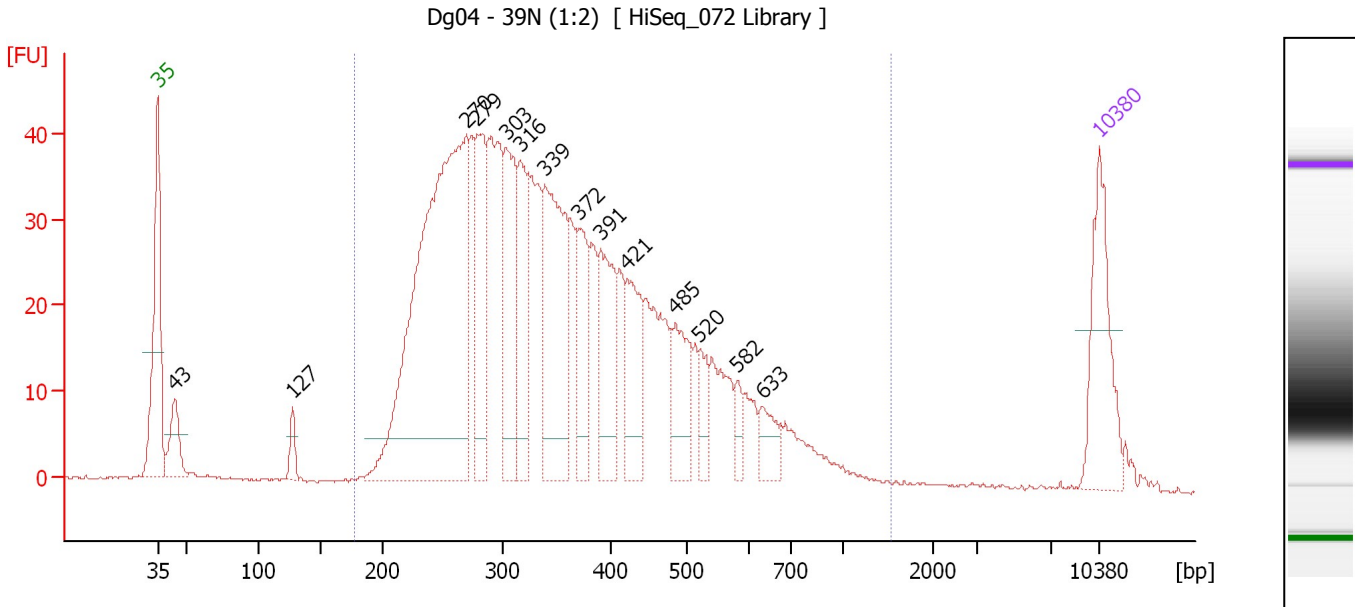
Region table for sample 2 : Dg04 - 38N (1:2)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
173	370	1,498	1,445.6	10,333.2	97	37.0	2,191.28	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
 Modified: 4/2/2015 3:16:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : Dg04 - 39N (1:2)

Number of peaks found: 14 Corr. Area 1: 1,092.5
 Noise: 0.2

Peak table for sample 3 : Dg04 - 39N (1:2)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	43	50.19	1,750.0		44.25
3	127	18.86	225.6		53.01
4	270	613.70	3,449.1		65.90
5	279	114.71	622.3		66.76
6	303	132.19	660.3		68.87
7	316	95.41	456.8		69.93
8	339	184.27	823.1		71.75
9	372	65.70	268.0		74.33
10	391	92.62	358.7		75.92
11	421	78.01	280.6		77.83
12	485	62.39	194.9		81.47
13	520	26.24	76.4		83.25
14	582	14.74	38.4		86.07
15	633	25.84	61.8		87.94
16	10,380	75.00	10.9	Upper Marker	113.00

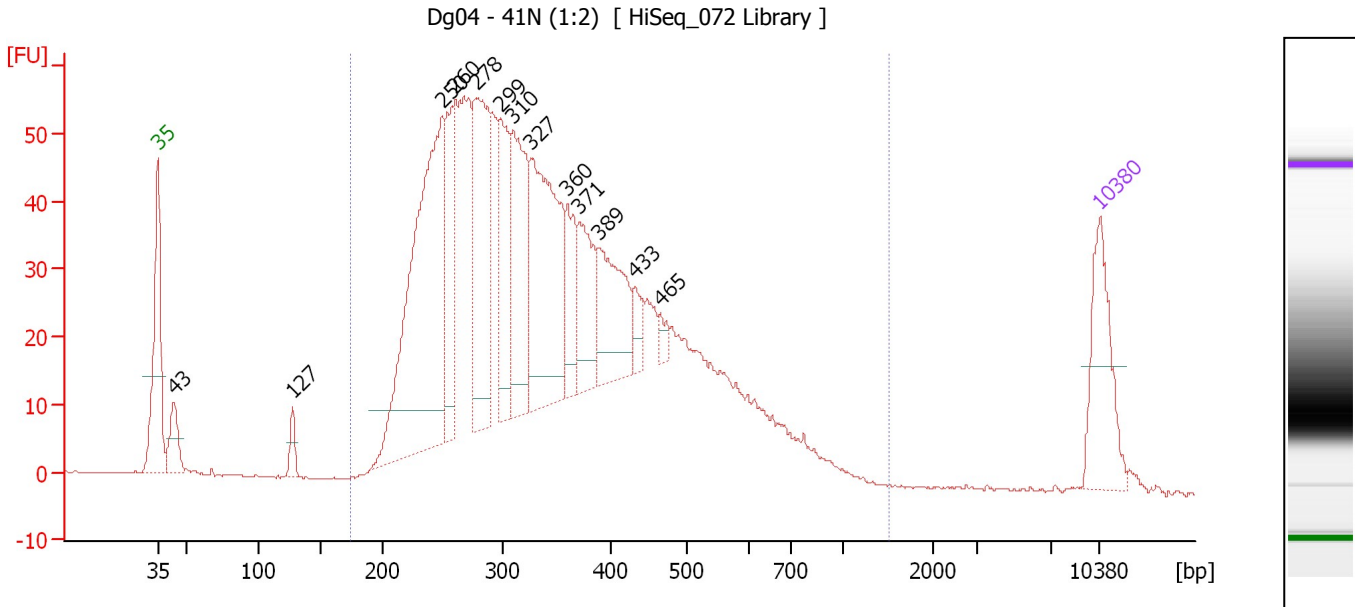
Region table for sample 3 : Dg04 - 39N (1:2)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
178	373	1,539	1,092.5	10,911.1	96	37.6	2,327.40	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
 Modified: 4/2/2015 3:16:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : Dg04 - 41N (1:2)

Number of peaks found: 13 Corr. Area 1: 1,460.6
 Noise: 0.2

Peak table for sample 4 : Dg04 - 41N (1:2)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	43	50.91	1,781.3		44.22
3	127	20.21	241.3		53.03
4	250	444.62	2,696.4		64.15
5	260	118.22	688.1		65.08
6	278	215.48	1,174.7		66.64
7	299	102.39	519.1		68.50
8	310	159.71	781.3		69.38
9	327	242.48	1,124.1		70.76
10	360	61.17	257.1		73.45
11	371	82.41	336.7		74.28
12	389	105.70	411.6		75.75
13	433	22.34	78.2		78.49
14	465	11.40	37.1		80.35
15	10,380	75.00	10.9	Upper Marker	113.00

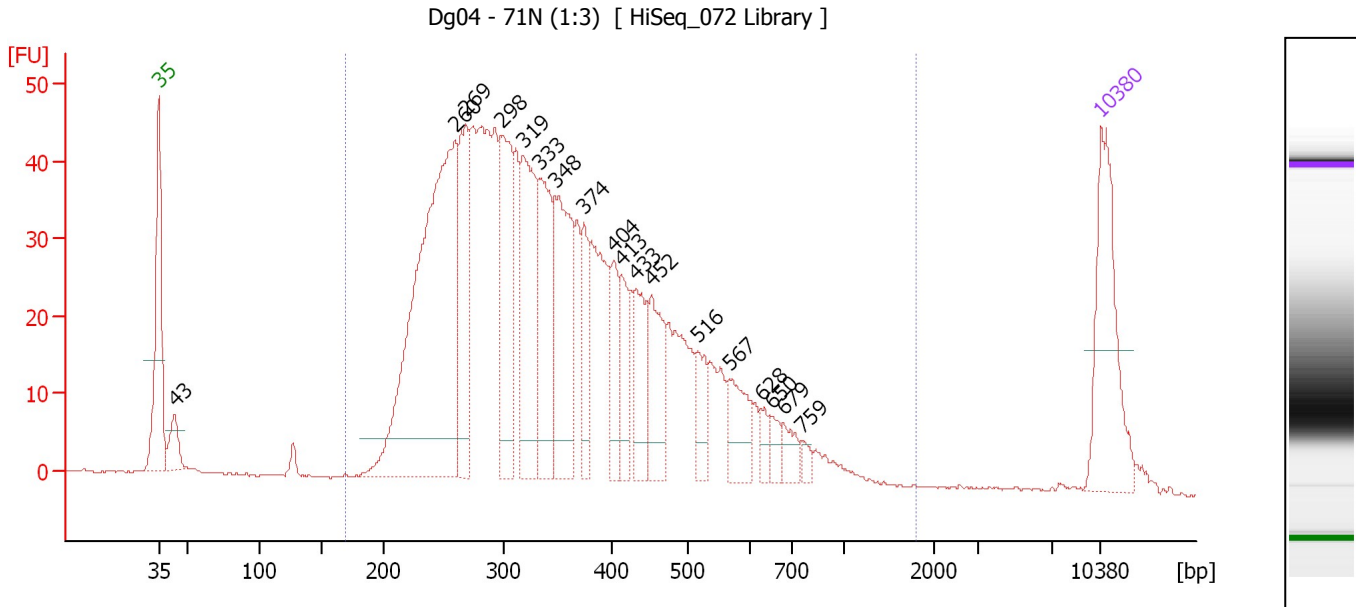
Region table for sample 4 : Dg04 - 41N (1:2)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
173	366	1,512	1,460.5	13,394.1	97	37.3	2,813.44	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
 Modified: 4/2/2015 3:16:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : Dg04 - 71N (1:3)

Number of peaks found: 18 Corr. Area 1: 1,197.2
 Noise: 0.2

Peak table for sample 5 : Dg04 - 71N (1:3)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	43	28.45	999.0		44.20
3	260	406.26	2,371.0		65.02
4	269	107.16	603.1		65.87
5	298	108.68	552.8		68.42
6	319	119.46	567.6		70.12
7	333	103.59	471.0		71.27
8	348	111.77	486.3		72.47
9	374	43.14	174.5		74.57
10	404	40.37	151.6		76.82
11	413	34.86	127.8		77.37
12	433	48.09	168.4		78.48
13	452	58.21	195.2		79.58
14	516	25.67	75.3		83.08
15	567	34.37	91.9		85.38
16	628	9.31	22.5		87.78
17	650	10.77	25.1		88.48
18	679	13.44	30.0		89.38
19	759	5.92	11.8		90.83
20	10,380	75.00	10.9	Upper Marker	113.00

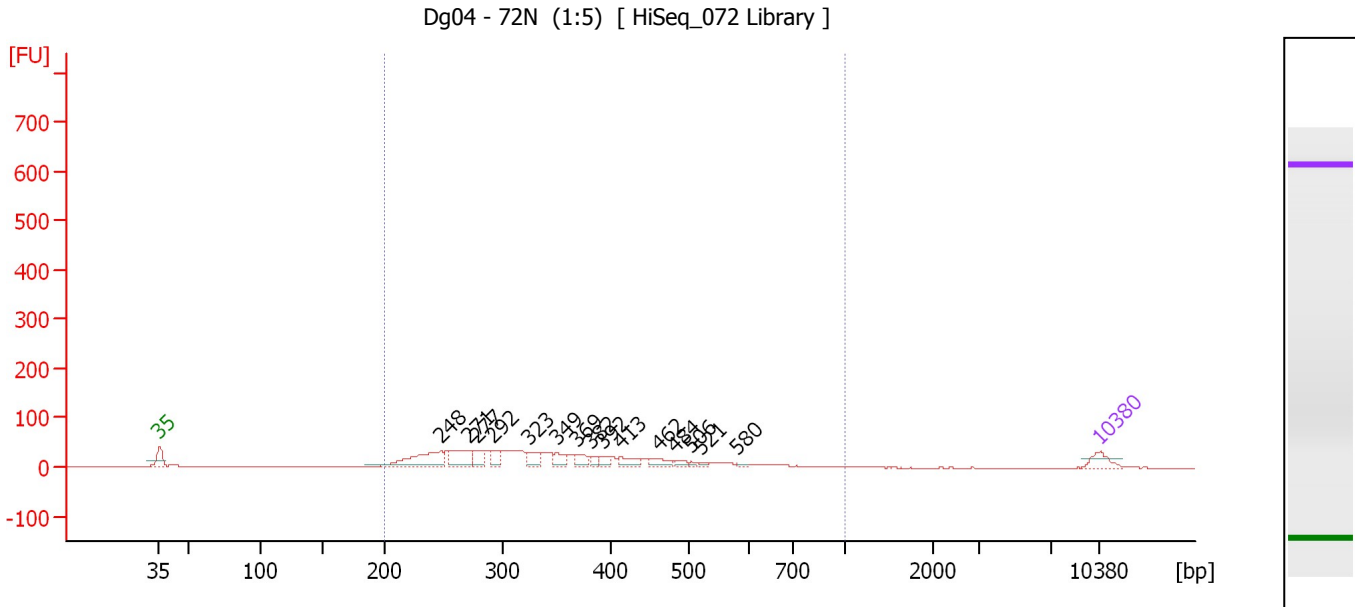
Region table for sample 5 : Dg04 - 71N (1:3)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
168	373	1,804	1,197.2	9,005.9	97	39.2	1,919.28	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
 Modified: 4/2/2015 3:16:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : Dg04 - 72N (1:5)

Number of peaks found: 15 Corr. Area 1: 489.0
 Noise: 0.2

Peak table for sample 6 : Dg04 - 72N (1:5)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	248	416.73	2,545.8		63.98
3	271	237.77	1,328.2		66.05
4	277	124.15	679.3		66.55
5	292	96.63	502.1		67.86
6	323	109.50	514.1		70.43
7	349	87.02	377.7		72.54
8	369	86.23	354.5		74.10
9	382	52.06	206.6		75.16
10	392	55.47	214.5		75.96
11	413	96.12	352.5		77.37
12	462	81.25	266.7		80.14
13	484	40.16	125.8		81.40
14	506	20.94	62.7		82.60
15	521	21.80	63.3		83.31
16	580	16.74	43.7		85.98
17	10,380	75.00	10.9	Upper Marker	113.00

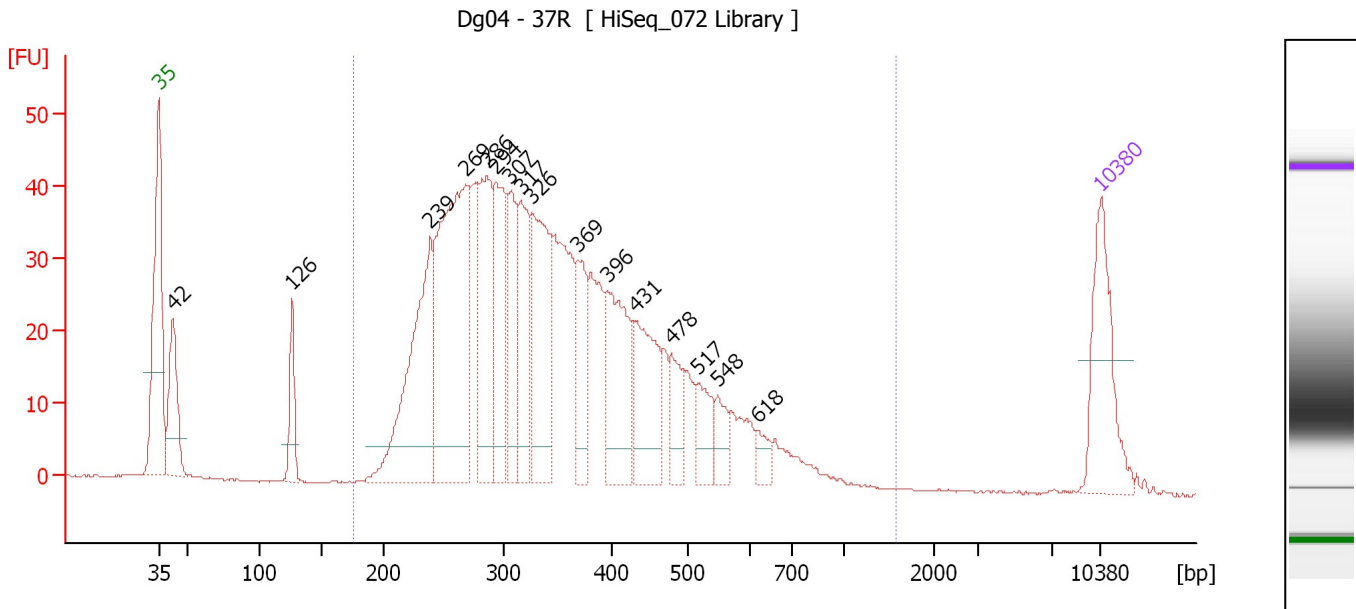
Region table for sample 6 : Dg04 - 72N (1:5)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
200	308	1,000	489.0	6,619.1	81	18.8	1,286.99	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

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 Modified: 4/2/2015 3:16:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : Dg04 - 37R

Number of peaks found: 16 Corr. Area 1: 1,060.7
 Noise: 0.2

Peak table for sample 7 : Dg04 - 37R

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	42	93.55	3,354.1		44.07
3	126	48.93	589.1		52.93
4	239	219.22	1,390.3		63.17
5	269	312.88	1,764.2		65.82
6	286	139.13	735.9		67.40
7	294	115.36	594.6		68.07
8	307	74.46	367.9		69.14
9	317	96.56	461.8		69.95
10	326	143.71	667.3		70.71
11	369	63.93	262.6		74.13
12	396	110.82	423.7		76.32
13	431	91.44	321.2		78.41
14	478	37.11	117.7		81.06
15	517	37.02	108.5		83.09
16	548	23.47	64.9		84.52
17	618	14.44	35.4		87.48
18	10,380	75.00	10.9	Upper Marker	113.00

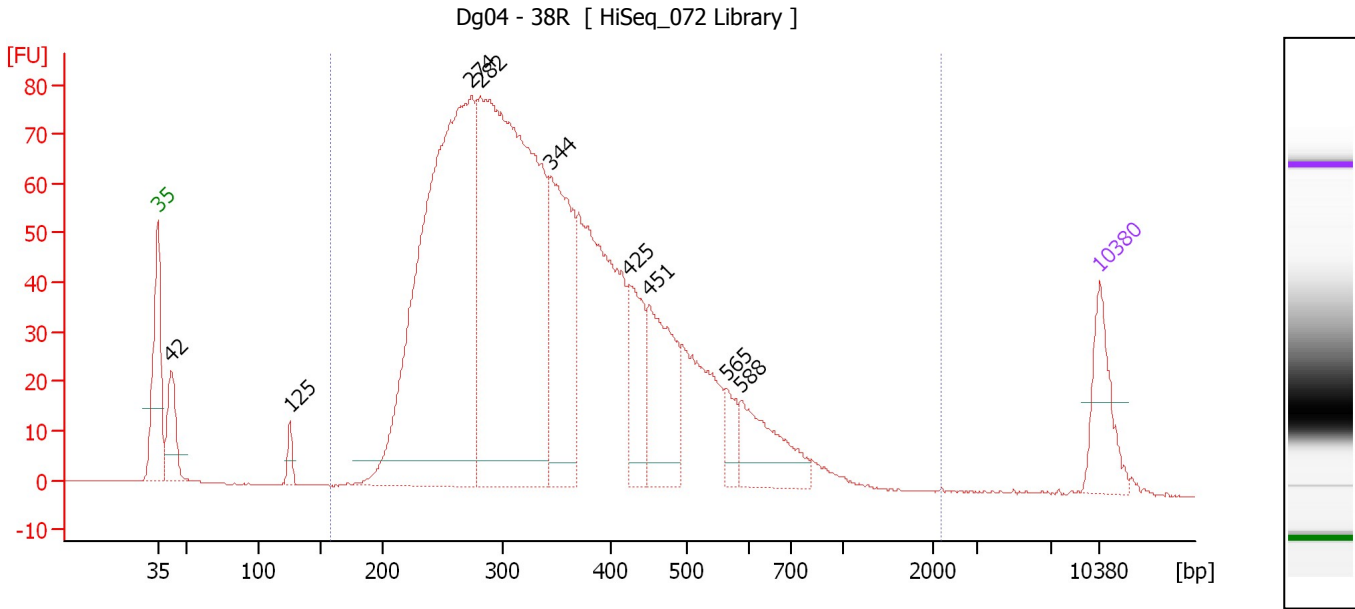
Region table for sample 7 : Dg04 - 37R

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
176	364	1,575	1,060.7	9,382.9	94	34.7	1,990.91	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

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 Modified: 4/2/2015 3:16:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : Dg04 - 38R

Number of peaks found: 9 Corr. Area 1: 1,933.0
 Noise: 0.1

Peak table for sample 8 : Dg04 - 38R

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	42	98.08	3,542.6		44.02
3	125	25.69	312.0		52.83
4	274	1,165.24	6,442.2		66.30
5	282	1,199.68	6,443.0		67.02
6	344	329.74	1,452.0		72.14
7	425	121.73	434.3		78.03
8	451	188.39	633.4		79.51
9	565	36.31	97.4		85.30
10	588	117.19	302.2		86.32
11	10,380	75.00	10.9	Upper Marker	113.00

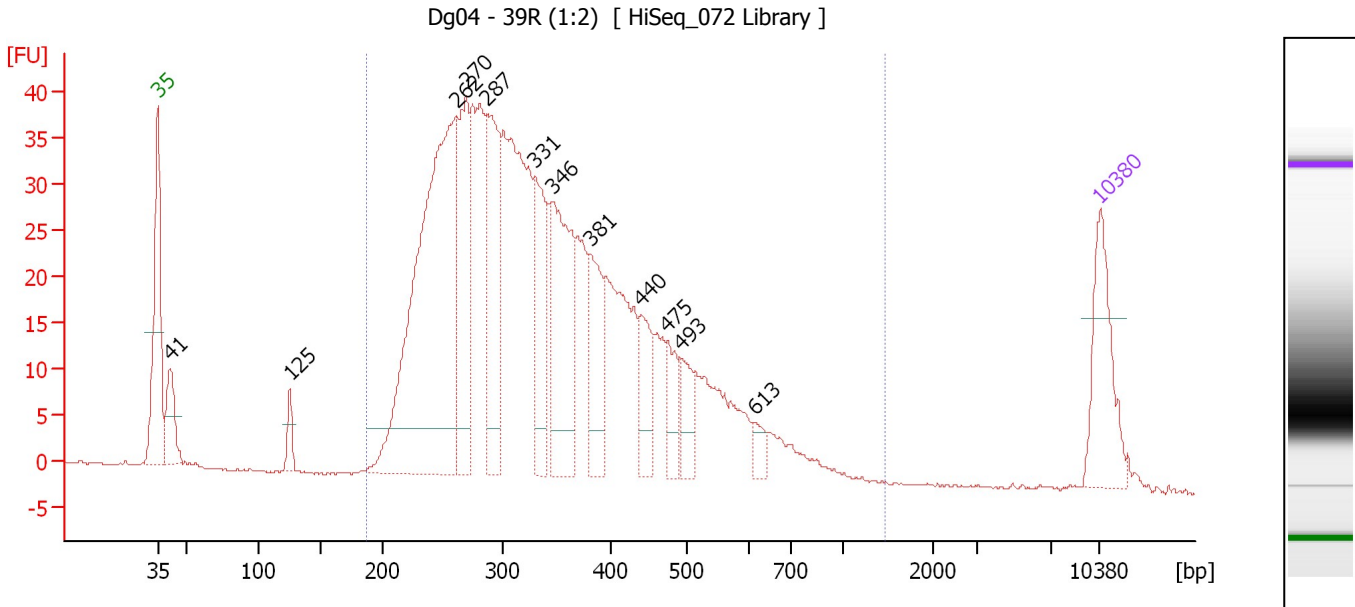
Region table for sample 8 : Dg04 - 38R

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
158	364	2,194	1,933.0	18,205.1	97	37.6	3,845.00	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
 Modified: 4/2/2015 3:16:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : Dg04 - 39R (1:2)

Number of peaks found: 12 Corr. Area 1: 916.0
 Noise: 0.1

Peak table for sample 9 : Dg04 - 39R (1:2)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	41	62.62	2,297.9		43.93
3	125	24.33	295.6		52.83
4	262	582.05	3,372.4		65.18
5	270	161.82	909.4		65.90
6	287	166.92	881.3		67.45
7	331	121.95	559.0		71.05
8	346	184.20	806.8		72.29
9	381	88.06	350.0		75.12
10	440	54.60	188.2		78.88
11	475	41.59	132.8		80.88
12	493	38.51	118.4		81.91
13	613	16.16	39.9		87.32
14	10,380	75.00	10.9	Upper Marker	113.00

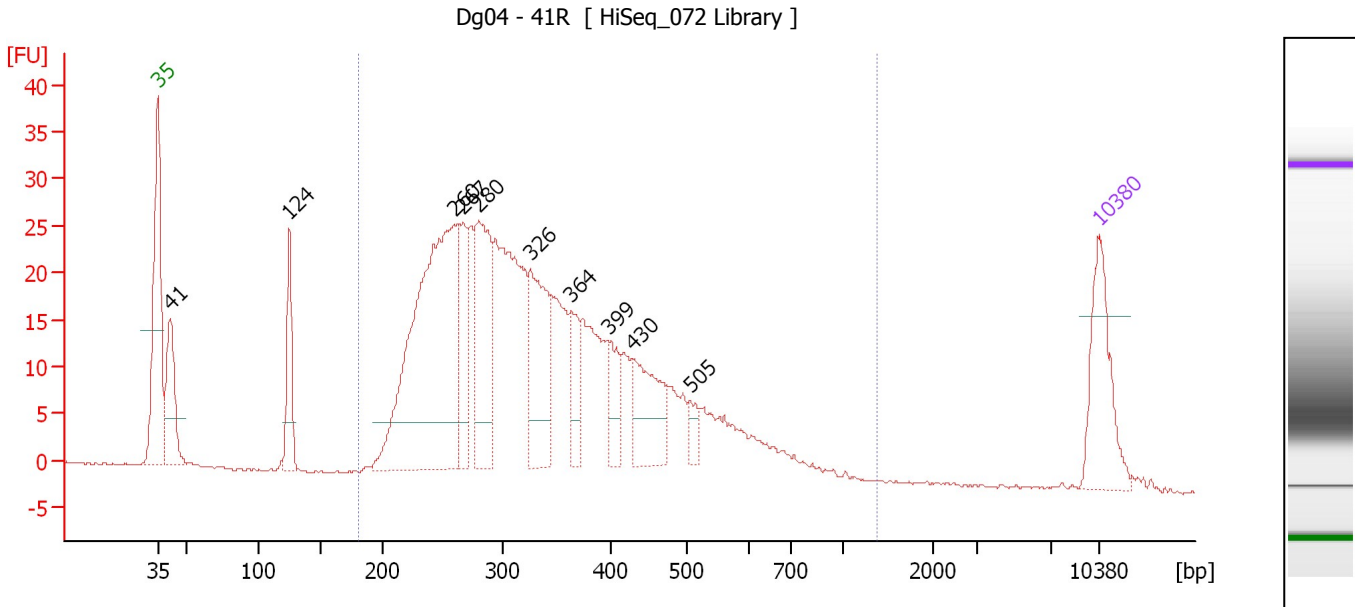
Region table for sample 9 : Dg04 - 39R (1:2)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
186	355	1,452	916.0	12,008.7	97	33.4	2,502.67	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
 Modified: 4/2/2015 3:16:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : Dg04 - 41R

Number of peaks found: 10 Corr. Area 1: 625.1
 Noise: 0.1

Peak table for sample 10 : Dg04 - 41R

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	41	101.39	3,718.2		43.93
3	124	81.01	989.1		52.77
4	260	475.17	2,772.2		65.02
5	267	83.49	473.3		65.70
6	280	153.02	827.8		66.83
7	326	143.01	664.1		70.71
8	364	44.39	184.6		73.76
9	399	41.29	156.7		76.55
10	430	88.26	311.3		78.31
11	505	18.09	54.3		82.55
12	10,380	75.00	10.9	Upper Marker	113.00

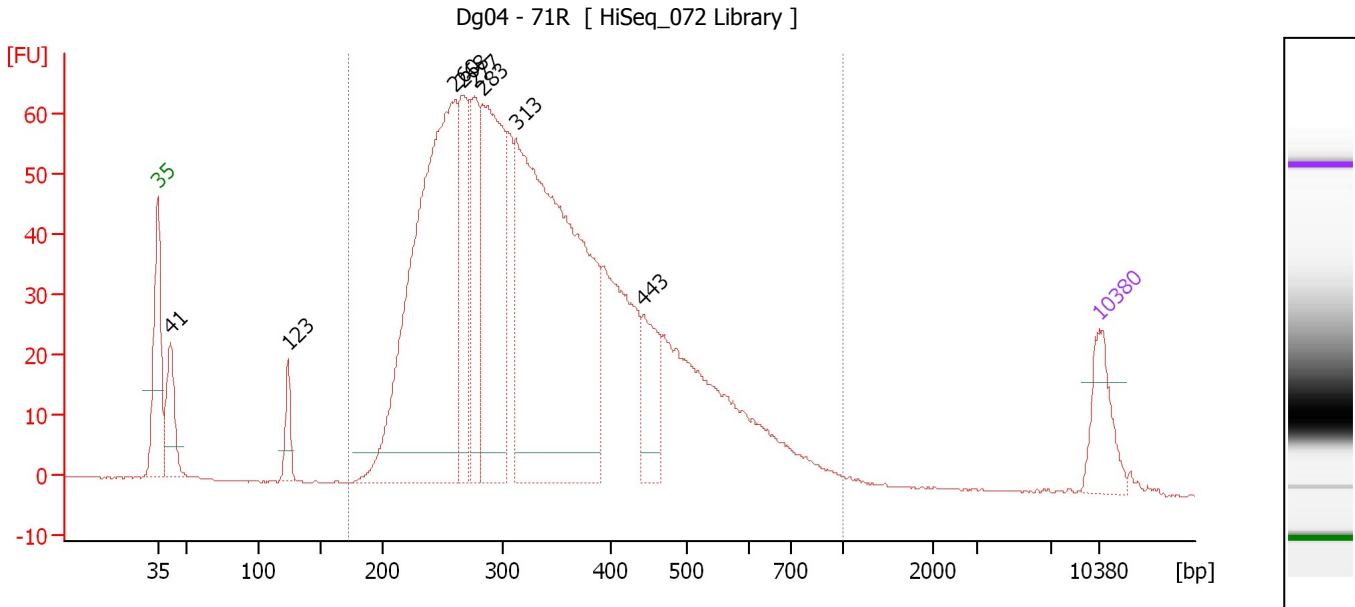
Region table for sample 10 : Dg04 - 41R

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
180	354	1,377	625.1	8,932.2	92	34.7	1,844.95	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
 Modified: 4/2/2015 3:16:58 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : Dg04 - 71R

Number of peaks found: 8 Corr. Area 1: 1,514.8
 Noise: 0.1

Peak table for sample 11 : Dg04 - 71R

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	41	137.79	5,045.3		43.94
3	123	60.89	750.5		52.66
4	260	1,104.48	6,437.8		65.04
5	268	228.25	1,292.4		65.72
6	277	207.13	1,133.0		66.56
7	283	502.64	2,686.8		67.13
8	313	1,185.59	5,740.3		69.64
9	443	120.18	410.7		79.10
10	10,380	75.00	10.9	Upper Marker	113.00

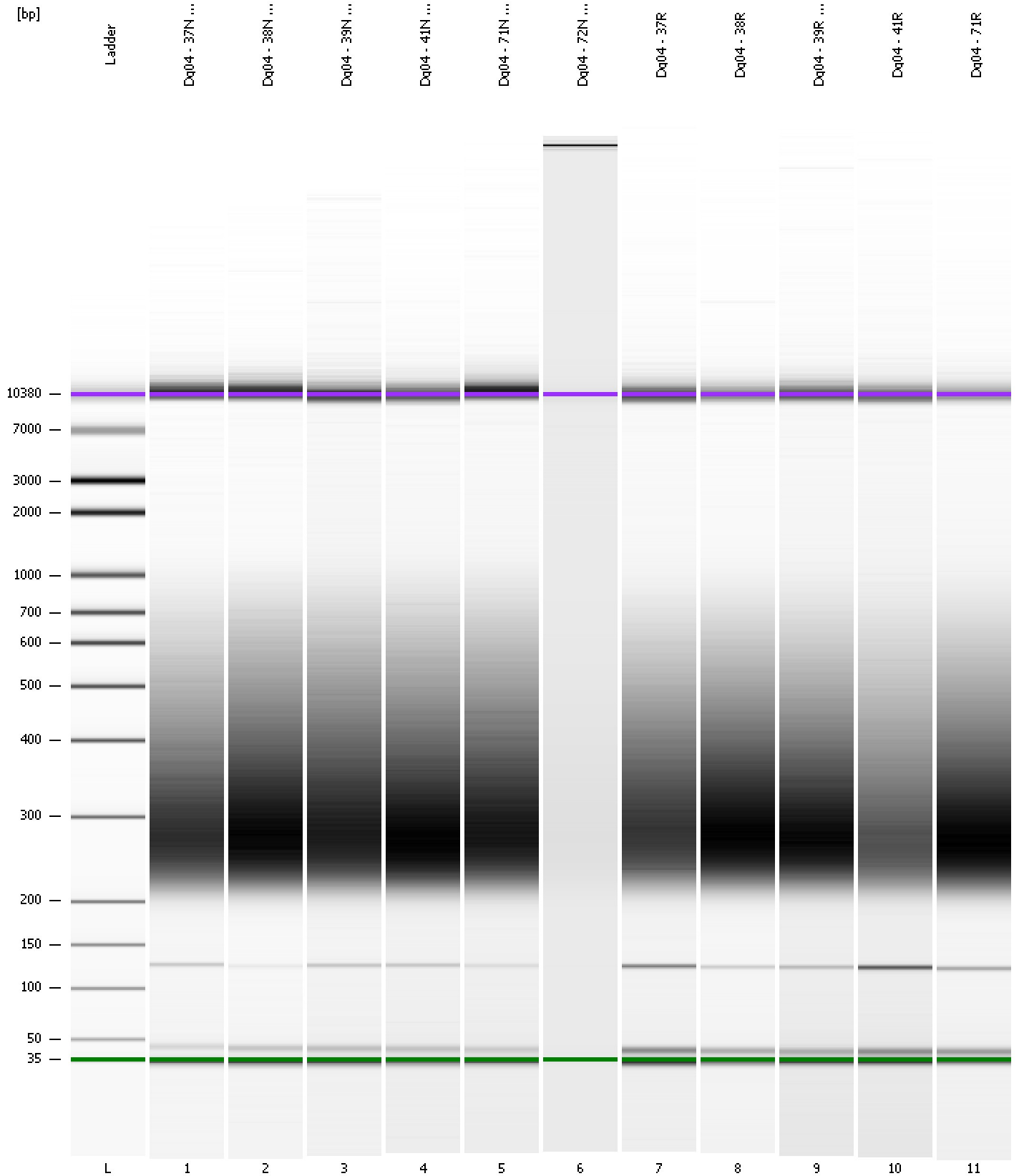
Region table for sample 11 : Dg04 - 71R

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Color
171	353	1,000	1,514.8	20,870.8	96	33.1	4,309.28	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
Modified: 4/2/2015 3:16:58 PM

Gel Image

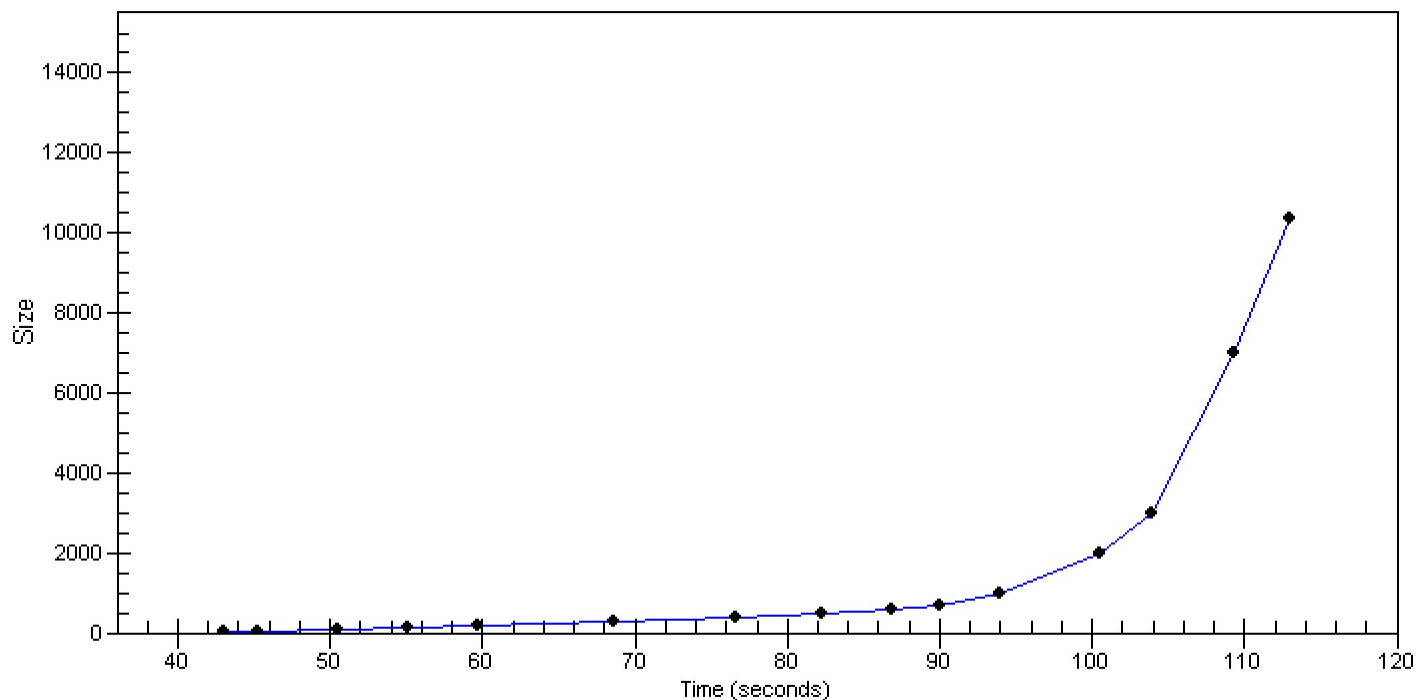


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad

Created: 4/2/2015 2:28:25 PM
Modified: 4/2/2015 3:16:58 PM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay Created: 4/2/2015 2:28:25 PM
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad Modified: 4/2/2015 3:16:58 PM

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		4/2/2015 3:09:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-04-02\2015-04-02_003.xad)		Instrument	Run		4/2/2015 2:28:30 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		4/2/2015 2:28:30 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		4/2/2015 2:28:30 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		4/2/2015 2:28:30 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		4/2/2015 2:28:30 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		4/2/2015 2:28:30 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		4/2/2015 2:28:30 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1