

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-02\2015-12-02_005.xad

Created: 12/2/2015 1:57:53 PM
Modified: 12/2/2015 2:31:28 PM

Electrophoresis File Run Summary

Instrument Information:

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

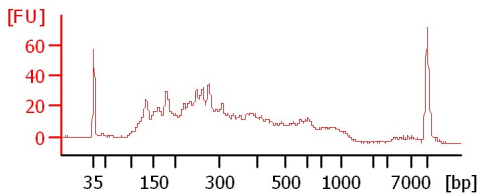
Assay Information:

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

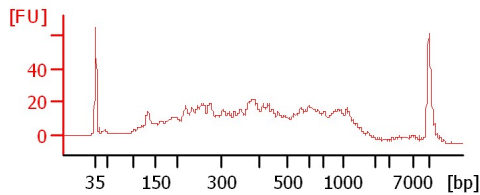
Chip Information:

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

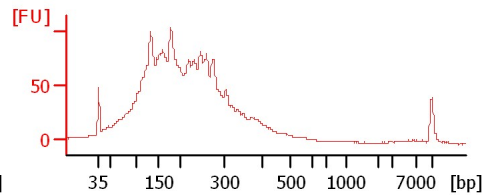
9h JI-1 cDNA



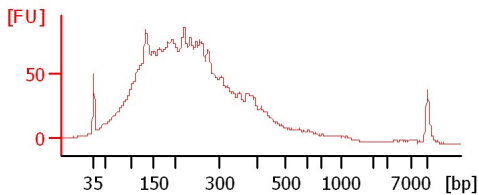
9h JI-2 cDNA



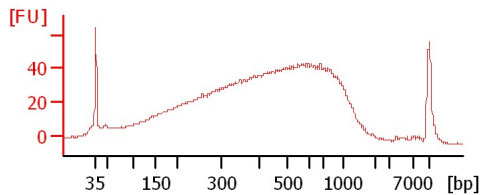
9h JI-1 sheared



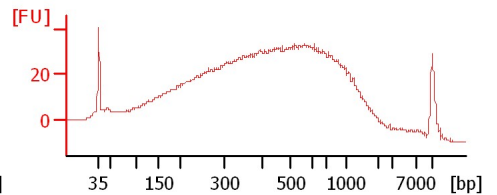
9h JI-2 sheared



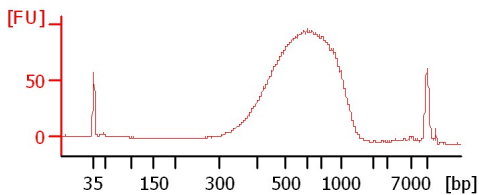
P111 soni



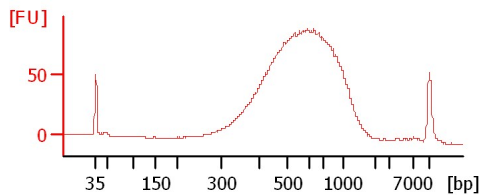
MP23 soni



P111 soni SS



MP23 soni SS



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Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
9h JI-1 cDNA		<input type="checkbox"/>	✓			
9h JI-2 cDNA		<input type="checkbox"/>	✓			
9h JI-1 sheared		<input type="checkbox"/>	✓			
9h JI-2 sheared		<input type="checkbox"/>	✓			
P111 soni		<input type="checkbox"/>	✓			
MP23 soni		<input type="checkbox"/>	✓			
P111 soni SS		<input type="checkbox"/>	✓			
MP23 soni SS		<input type="checkbox"/>	✓			
sample 9		<input type="checkbox"/>				
sample 10		<input type="checkbox"/>				
sample 11		<input type="checkbox"/>				
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

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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

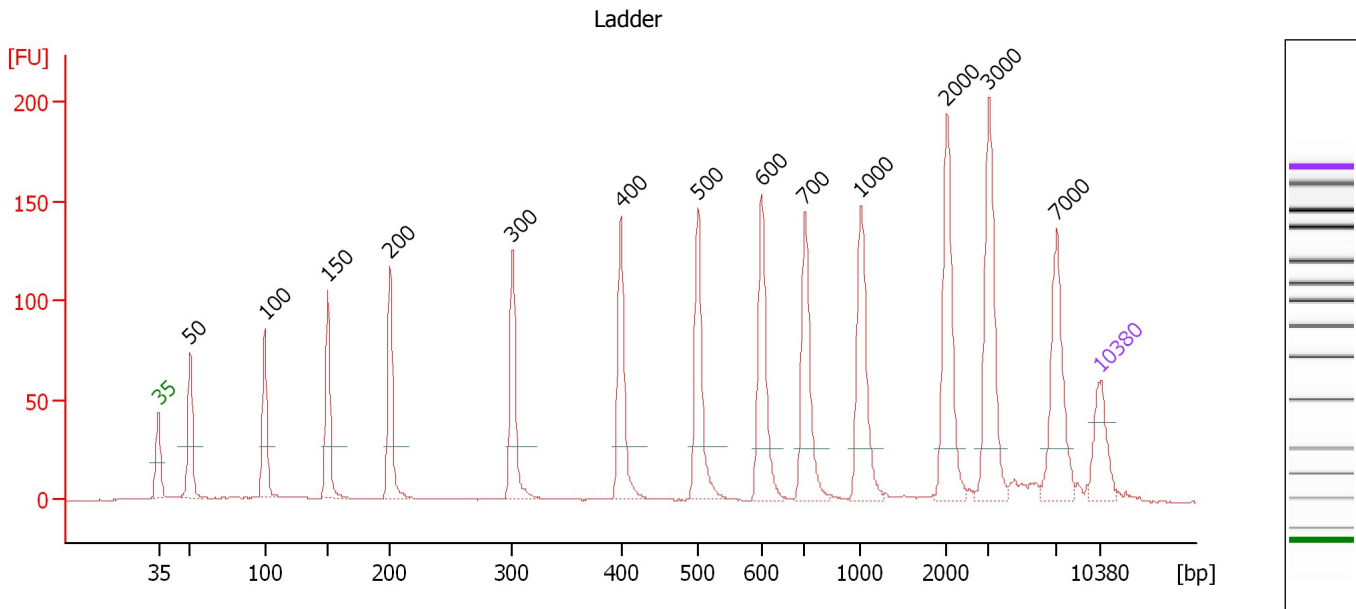
Ladder

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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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

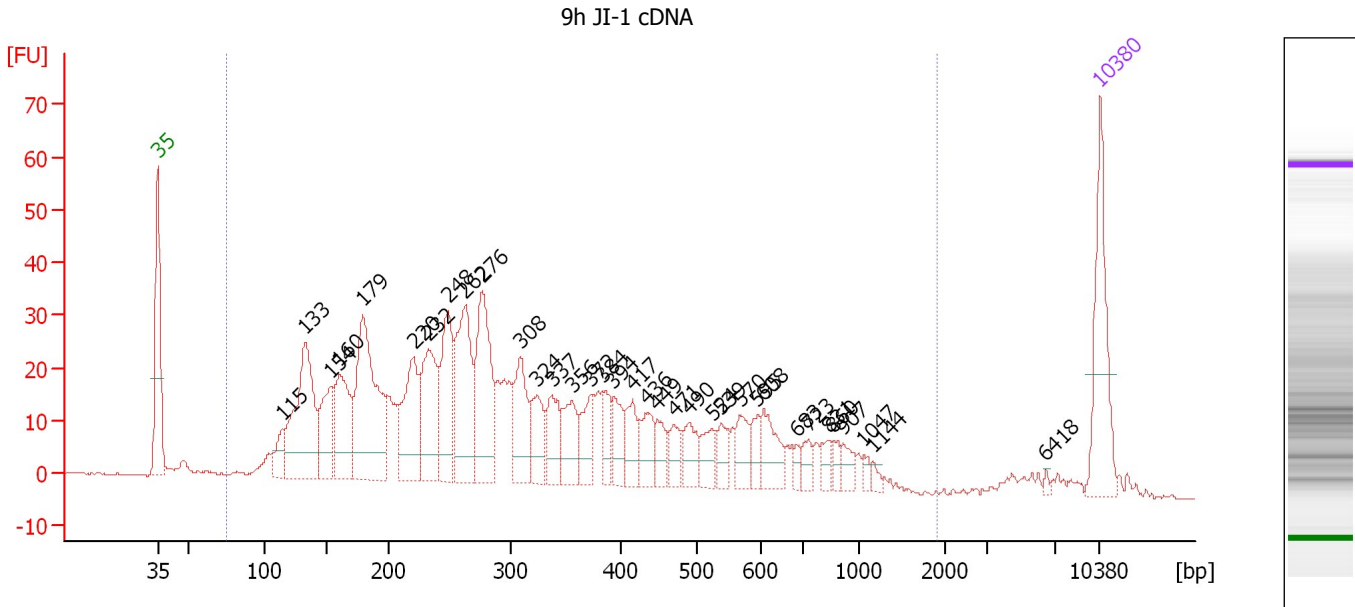
Peak table for Ladder

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	45.35
3	100	150.00	2,272.7	Ladder Peak	50.88
4	150	150.00	1,515.2	Ladder Peak	55.57
5	200	150.00	1,136.4	Ladder Peak	60.21
6	300	150.00	757.6	Ladder Peak	69.28
7	400	150.00	568.2	Ladder Peak	77.34
8	500	150.00	454.5	Ladder Peak	83.04
9	600	150.00	378.8	Ladder Peak	87.78
10	700	150.00	324.7	Ladder Peak	91.01
11	1,000	150.00	227.3	Ladder Peak	95.17
12	2,000	150.00	113.6	Ladder Peak	101.54
13	3,000	150.00	75.8	Ladder Peak	104.68
14	7,000	150.00	32.5	Ladder Peak	109.73
15	10,380	75.00	10.9	Upper Marker	113.00

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Electropherogram Summary Continued ...



Overall Results for sample 1 : 9h JI-1 cDNA

Number of peaks found: 35 Corr. Area 1: 1,066.1
 Noise: 0.2

Peak table for sample 1 : 9h JI-1 cDNA

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	115	29.69	392.4		52.25
3	133	186.21	2,121.8		53.97
4	154	70.78	697.9		55.91
5	160	99.50	943.2		56.48
6	179	210.54	1,784.2		58.24
7	220	104.02	716.2		62.03
8	232	108.67	710.6		63.09
9	248	98.60	601.9		64.59
10	262	133.42	770.7		65.86
11	276	133.53	732.0		67.14
12	308	70.86	348.7		69.92
13	324	44.07	206.3		71.19
14	337	45.64	204.9		72.30
15	356	46.59	198.3		73.79
16	372	38.20	155.4		75.11
17	384	26.60	104.8		76.08
18	394	31.23	120.2		76.83
19	417	35.07	127.5		78.29
20	436	32.65	113.5		79.39
21	449	23.27	78.5		80.14
22	471	20.73	66.7		81.37
23	490	28.35	87.7		82.47
24	524	23.92	69.1		84.19
25	539	19.76	55.5		84.89

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Electropherogram Summary Continued ...

... Peak table for sample 1 : 9h JI-1 cDNA

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	570	26.37	70.1		86.35
27	595	17.80	45.3		87.54
28	608	38.11	95.0		88.02
29	683	9.20	20.4		90.44
30	723	13.98	29.3		91.33
31	831	11.43	20.8		92.82
32	860	9.77	17.2		93.22
33	907	12.82	21.4		93.88
34	1,047	5.77	8.3		95.47
35	1,144	4.92	6.5		96.08
36	6,418	2.49	0.6		108.99
37	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 1 : 9h JI-1 cDNA

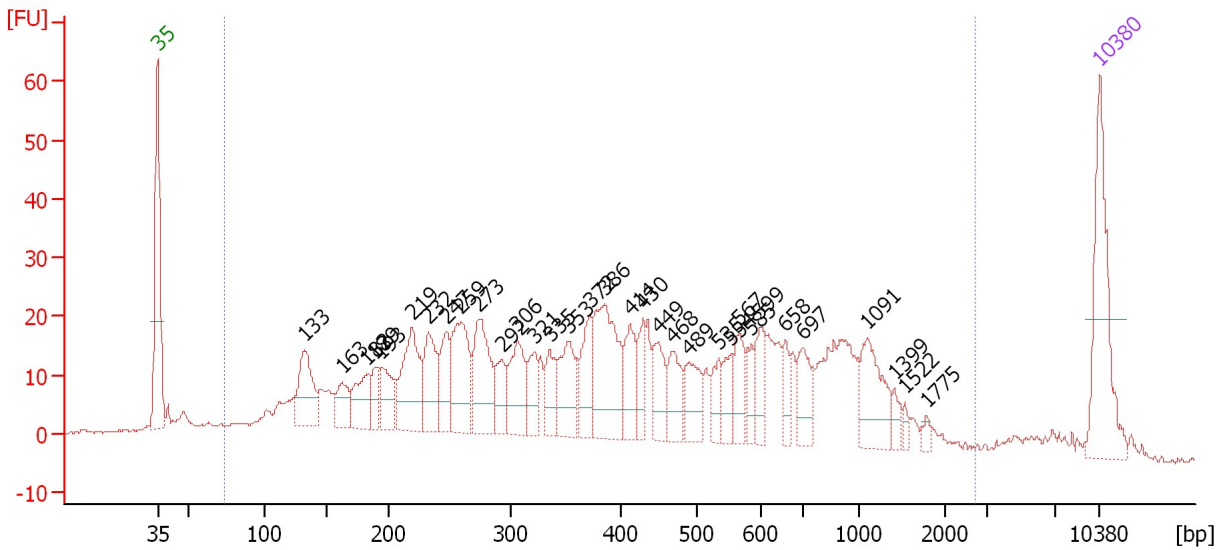
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
75	1,894	355	1,066.1	13,328.6	2,077.21	96	59.3

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Electropherogram Summary Continued ...

9h JI-2 cDNA



Overall Results for sample 2 : 9h JI-2 cDNA

Number of peaks found: 33 Corr. Area 1: 1,038.9
 Noise: 0.2

Peak table for sample 2 : 9h JI-2 cDNA

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	133	67.57	772.1		53.93
3	163	30.41	282.9		56.76
4	182	43.47	361.7		58.55
5	189	20.50	164.6		59.16
6	193	35.71	279.8		59.60
7	219	83.73	580.2		61.90
8	232	52.68	343.9		63.12
9	247	39.35	241.4		64.47
10	259	77.28	452.1		65.56
11	273	75.18	417.4		66.83
12	292	27.35	141.8		68.57
13	306	50.89	252.2		69.75
14	321	28.17	133.0		70.97
15	335	30.05	135.7		72.14
16	353	47.74	205.0		73.54
17	372	41.88	170.5		75.10
18	386	95.38	374.1		76.24
19	414	38.90	142.5		78.11
20	430	22.21	78.2		79.07
21	449	34.09	114.9		80.16
22	468	33.81	109.4		81.25
23	489	34.05	105.5		82.42
24	535	19.94	56.5		84.69
25	554	21.61	59.1		85.60

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Electropherogram Summary Continued ...

... Peak table for sample 2 : 9h JI-2 cDNA

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	567	29.12	77.8		86.21
27	585	15.92	41.2		87.08
28	599	26.35	66.6		87.74
29	658	16.06	37.0		89.65
30	697	30.66	66.6		90.92
31	1,091	47.87	66.5		95.75
32	1,399	8.70	9.4		97.71
33	1,522	4.98	5.0		98.49
34	1,775	4.86	4.1		100.11
35	10,380	75.00	10.9	Upper Marker	113.00

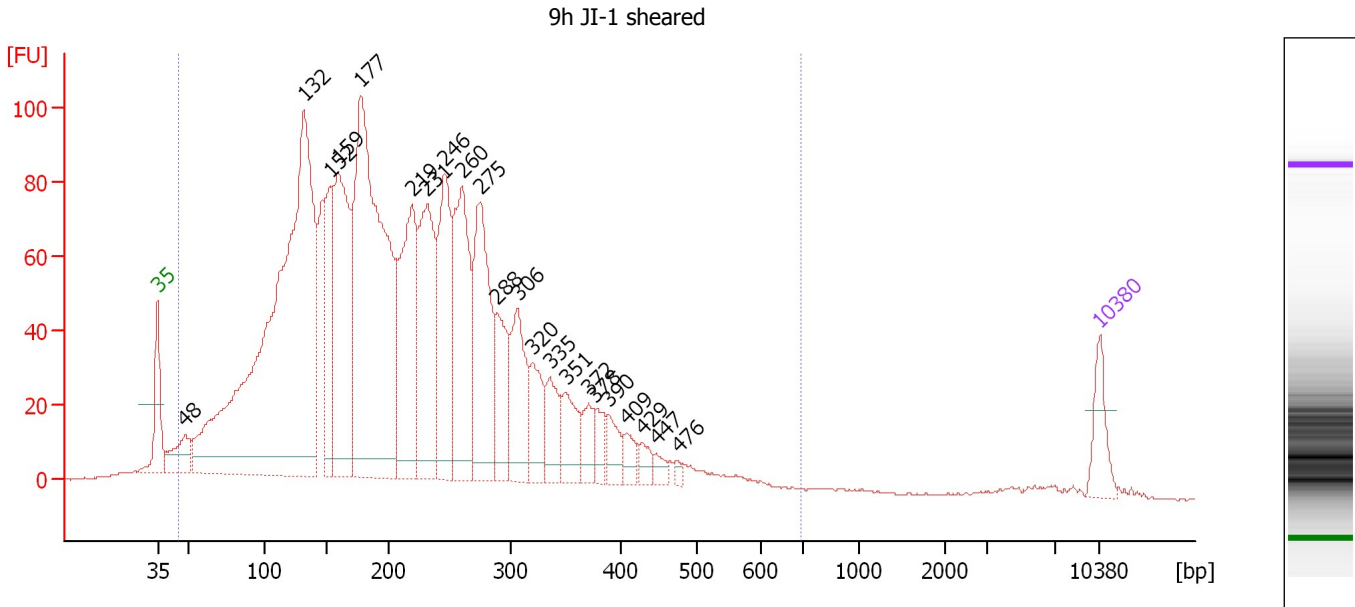
Region table for sample 2 : 9h JI-2 cDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
74	2,704	501	1,038.9	9,962.3	1,809.58	94	72.0

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Electropherogram Summary Continued ...



Overall Results for sample 3 : 9h JI-1 sheared

Number of peaks found: 22 Corr. Area 1: 2,909.3
 Noise: 0.3

Peak table for sample 3 : 9h JI-1 sheared

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	48	124.06	3,918.4		45.03
3	132	2,594.44	29,862.1		53.84
4	152	295.78	2,941.8		55.78
5	159	691.59	6,571.7		56.44
6	177	1,559.78	13,351.7		58.08
7	219	537.11	3,720.7		61.91
8	231	502.66	3,291.9		63.06
9	246	432.78	2,666.2		64.38
10	260	516.66	3,010.5		65.66
11	275	487.37	2,689.1		66.98
12	288	213.76	1,123.7		68.21
13	306	250.77	1,242.2		69.76
14	320	143.12	678.5		70.86
15	335	121.01	546.6		72.14
16	351	123.57	533.7		73.37
17	372	69.70	284.2		75.05
18	378	60.00	240.8		75.53
19	390	63.11	245.4		76.50
20	409	48.71	180.7		77.82
21	429	38.82	137.2		78.97
22	447	26.40	89.4		80.03
23	476	14.17	45.1		81.66
24	10,380	75.00	10.9	Upper Marker	113.00

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Electropherogram Summary Continued ...

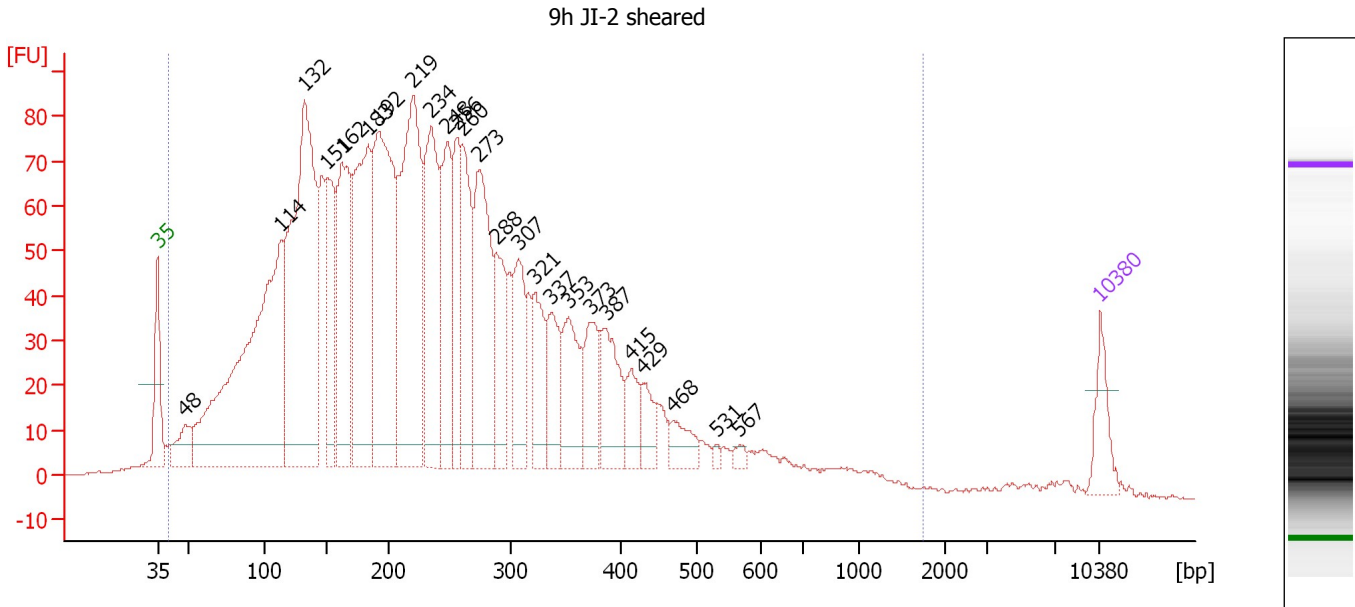
... Region table for sample 3 : 9h JI-1 sheared

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
45	694	220	2,909.3	92,730.2	9,910.09	97	44.8

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Electropherogram Summary Continued ...



Overall Results for sample 4 : 9h JI-2 sheared

Number of peaks found: 25 Corr. Area 1: 3,072.5
 Noise: 0.2

Peak table for sample 4 : 9h JI-2 sheared

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	48	110.87	3,498.0		45.04
3	114	1,403.60	18,661.7		52.19
4	132	1,250.37	14,320.9		53.90
5	151	312.88	3,147.3		55.62
6	162	517.70	4,832.6		56.71
7	183	643.20	5,318.1		58.66
8	192	828.20	6,551.7		59.43
9	219	827.35	5,717.3		61.96
10	234	488.96	3,169.7		63.27
11	248	342.38	2,094.4		64.54
12	256	241.61	1,431.8		65.26
13	260	297.36	1,735.1		65.62
14	273	463.22	2,569.6		66.85
15	288	214.96	1,132.5		68.16
16	307	213.52	1,054.4		69.83
17	321	171.25	807.2		71.01
18	337	155.13	697.1		72.28
19	353	196.59	844.0		73.54
20	373	156.48	635.4		75.17
21	387	183.29	717.3		76.30
22	415	86.16	314.4		78.20
23	429	70.54	248.9		79.02
24	468	69.35	224.4		81.24
25	531	10.26	29.3		84.49

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Electropherogram Summary Continued ...

... Peak table for sample 4 : 9h JI-2 sheared

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	567	16.68	44.6		86.21
27	10,380	75.00	10.9	Upper Marker	113.00

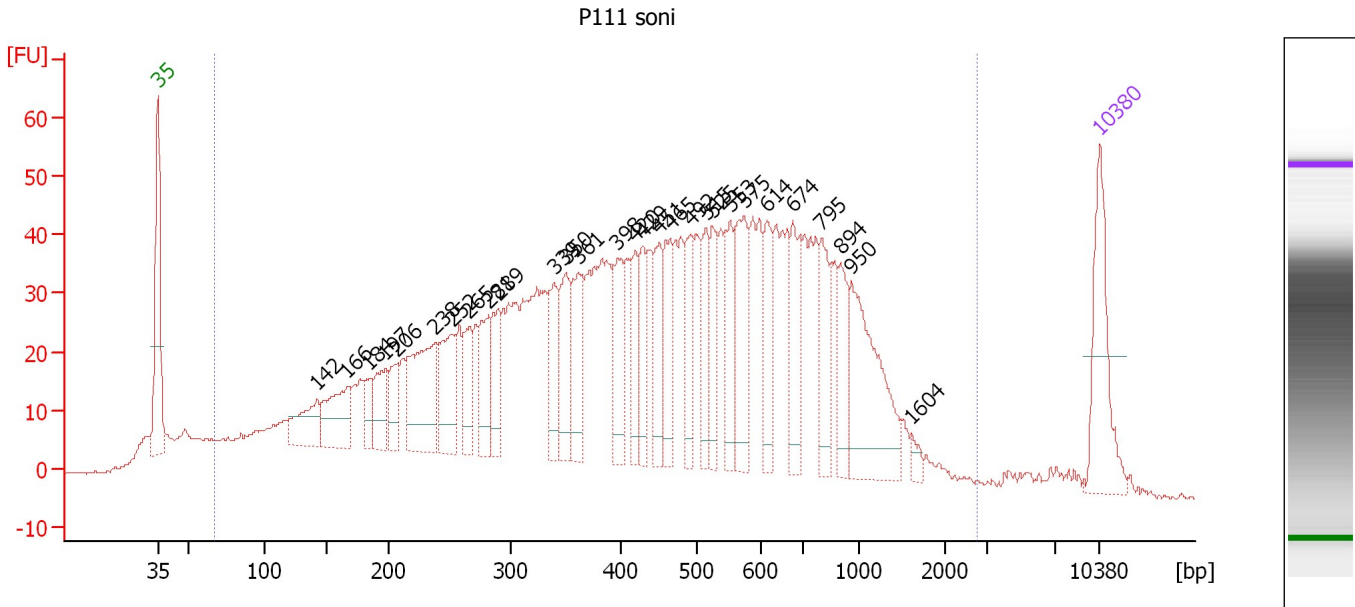
Region table for sample 4 : 9h JI-2 sheared

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
40	1,738	257	3,072.5	99,576.6	11,057.19	98	62.3

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Electropherogram Summary Continued ...



Overall Results for sample 5 : P111 soni

Number of peaks found: 29 Corr. Area 1: 1,991.9
 Noise: 0.2

Peak table for sample 5 : P111 soni

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	142	61.18	652.3		54.83
3	166	78.44	715.0		57.07
4	184	28.67	236.1		58.72
5	197	45.37	349.3		59.92
6	206	37.17	273.7		60.74
7	238	127.13	810.5		63.63
8	252	75.91	456.8		64.91
9	265	44.16	252.1		66.15
10	281	55.36	298.3		67.57
11	289	50.80	266.1		68.30
12	339	54.34	242.8		72.43
13	350	65.76	284.3		73.35
14	361	65.82	276.0		74.22
15	398	59.43	226.4		77.15
16	420	45.96	165.8		78.48
17	429	46.92	165.7		78.99
18	451	50.53	169.6		80.27
19	465	58.59	190.9		81.05
20	492	51.69	159.3		82.56
21	515	51.10	150.4		83.75
22	525	47.38	136.8		84.21
23	553	56.84	155.8		85.54
24	575	81.43	214.6		86.60
25	614	47.32	116.7		88.25

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Electropherogram Summary Continued ...

... Peak table for sample 5 : P111 soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	674	60.12	135.1		90.17
27	795	53.16	101.3		92.33
28	894	43.03	72.9		93.70
29	950	131.41	209.5		94.48
30	1,604	7.42	7.0		99.02
31	10,380	75.00	10.9	Upper Marker	113.00

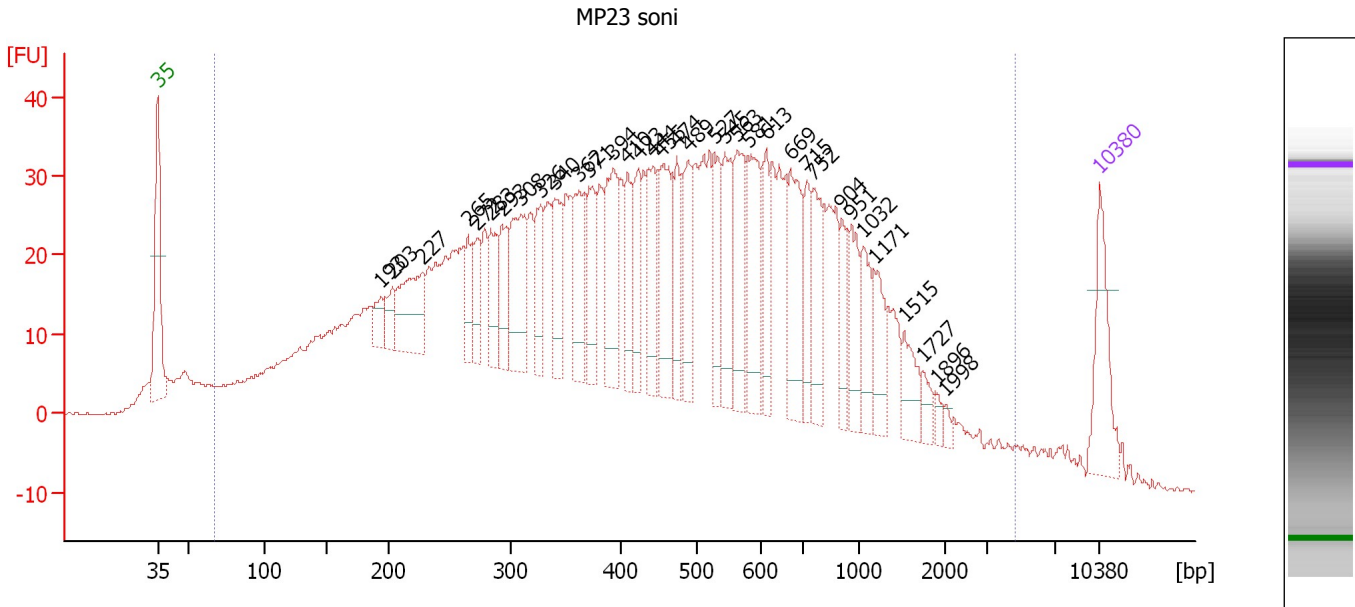
Region table for sample 5 : P111 soni

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
67	2,718	507	1,991.9	18,636.8	3,559.66	95	62.9

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Electropherogram Summary Continued ...



Overall Results for sample 6 : MP23 soni

Number of peaks found: 35 Corr. Area 1: 1,838.4
 Noise: 0.2

Peak table for sample 6 : MP23 soni

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	193	29.13	228.7		59.56
3	203	32.38	242.2		60.44
4	227	116.49	776.8		62.68
5	265	48.83	279.5		66.08
6	272	49.29	274.6		66.74
7	283	59.74	320.1		67.72
8	293	61.25	317.3		68.60
9	308	120.18	590.5		69.96
10	326	66.65	309.5		71.40
11	340	69.55	309.8		72.52
12	362	88.79	371.4		74.29
13	371	66.19	270.0		75.04
14	394	111.76	429.7		76.86
15	410	62.11	229.7		77.88
16	423	68.10	243.7		78.68
17	444	89.41	305.2		79.84
18	455	119.60	398.7		80.45
19	474	64.10	204.9		81.57
20	489	79.77	247.3		82.41
21	527	70.44	202.6		84.32
22	545	83.69	232.8		85.16
23	563	93.39	251.2		86.04
24	581	108.99	284.2		86.88
25	613	60.54	149.7		88.19

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-02\2015-12-02_005.xad

Created: 12/2/2015 1:57:53 PM
 Modified: 12/2/2015 2:31:28 PM

Electropherogram Summary Continued ...

... Peak table for sample 6 : MP23 soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	669	109.32	247.6		90.01
27	715	55.48	117.5		91.22
28	752	77.04	155.1		91.73
29	904	43.30	72.6		93.83
30	951	58.26	92.8		94.49
31	1,032	53.59	78.7		95.37
32	1,171	48.71	63.0		96.26
33	1,515	37.47	37.5		98.45
34	1,727	15.87	13.9		99.80
35	1,896	8.16	6.5		100.87
36	1,998	6.99	5.3		101.53
37	10,380	75.00	10.9	Upper Marker	113.00

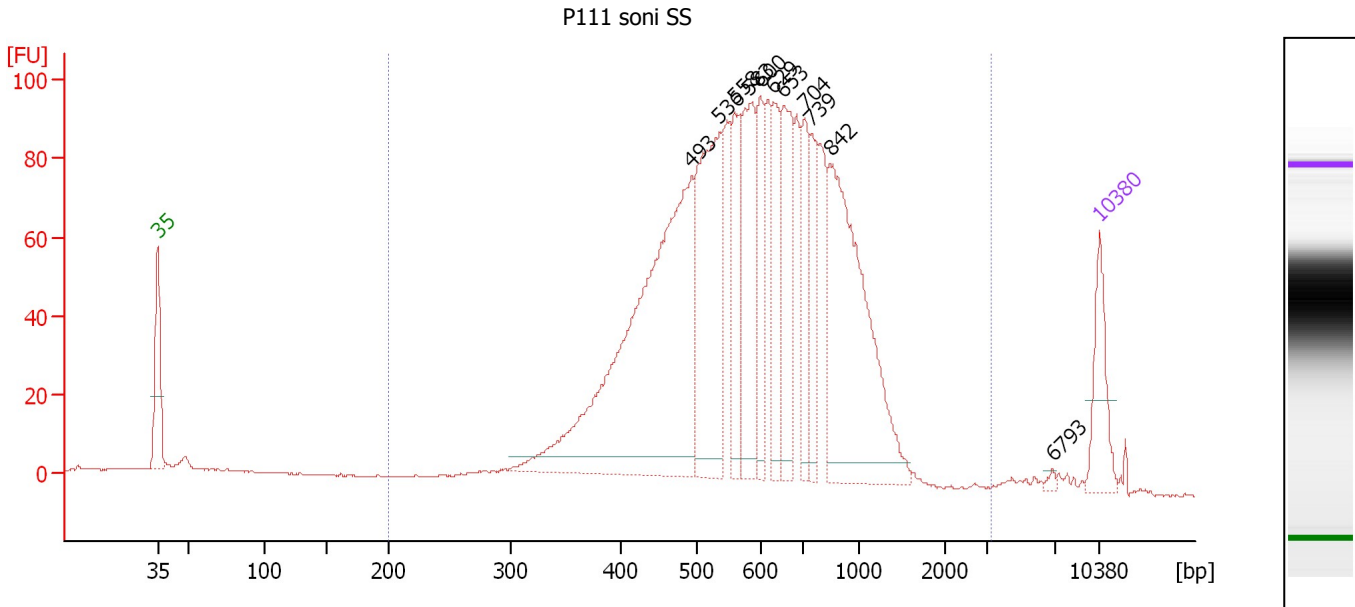
Region table for sample 6 : MP23 soni

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
67	4,599	566	1,838.4	30,164.6	5,809.30	95	85.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-02\2015-12-02_005.xad

Created: 12/2/2015 1:57:53 PM
 Modified: 12/2/2015 2:31:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : P111 soni SS

Number of peaks found: 11 Corr. Area 1: 1,927.4
 Noise: 0.2

Peak table for sample 7 : P111 soni SS

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	493	869.95	2,672.3		82.66
3	536	366.74	1,036.2		84.76
4	558	130.75	354.7		85.81
5	583	215.14	559.4		86.96
6	600	111.17	280.8		87.77
7	629	127.60	307.2		88.73
8	653	168.05	389.9		89.49
9	704	105.61	227.2		91.07
10	739	100.07	205.2		91.55
11	842	461.17	829.7		92.98
12	6,793	4.33	1.0		109.46
13	10,380	75.00	10.9	Upper Marker	113.00

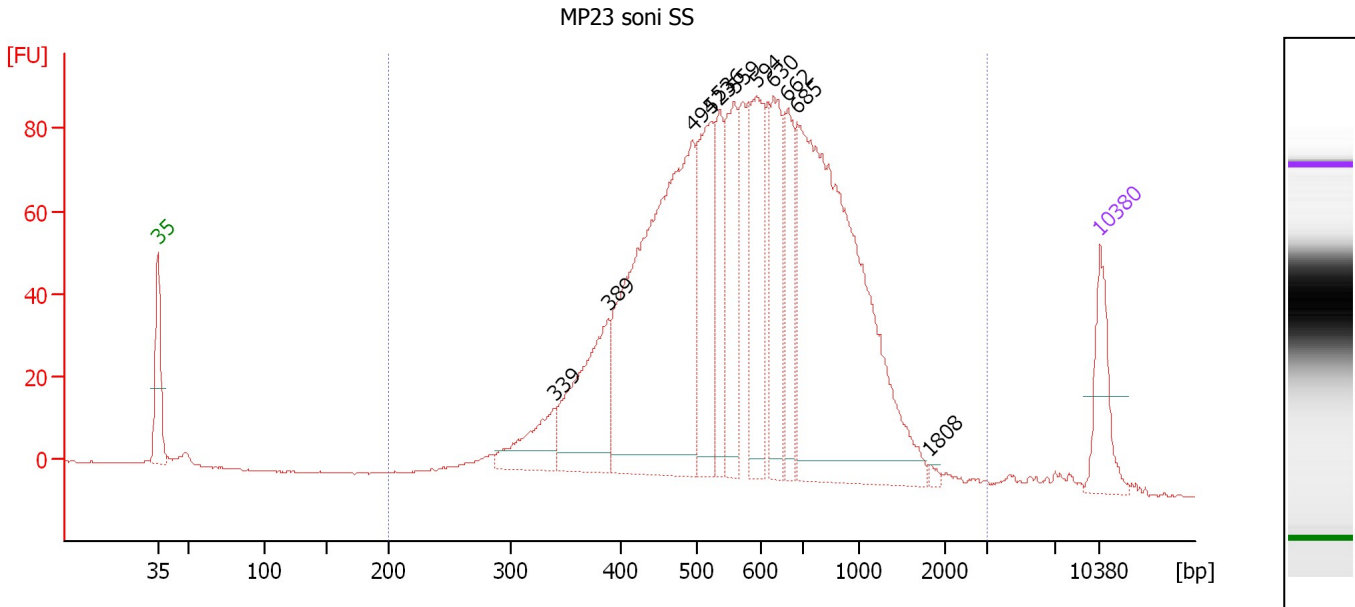
Region table for sample 7 : P111 soni SS

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	3,124	641	1,927.4	9,057.5	3,302.58	94	38.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-02\2015-12-02_005.xad

Created: 12/2/2015 1:57:53 PM
 Modified: 12/2/2015 2:31:28 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : MP23 soni SS

Number of peaks found: 11 Corr. Area 1: 1,982.6
 Noise: 0.2

Peak table for sample 8 : MP23 soni SS

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	339	101.47	452.9		72.46
3	389	233.07	907.4		76.46
4	494	807.40	2,475.9		82.71
5	523	230.02	666.7		84.12
6	536	117.00	330.6		84.76
7	559	172.73	468.3		85.83
8	594	209.28	534.0		87.49
9	630	151.28	363.7		88.76
10	662	116.16	265.9		89.78
11	685	857.63	1,898.0		90.51
12	1,808	5.35	4.5		100.32
13	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 8 : MP23 soni SS

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	2,998	640	1,982.6	9,294.6	3,305.71	99	43.5

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-02\2015-12-02_005.xad

Created: 12/2/2015 1:57:53 PM
Modified: 12/2/2015 2:31:28 PM

Gel Image

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-02\2015-12-02_005.xad

Created: 12/2/2015 1:57:53 PM
Modified: 12/2/2015 2:31:28 PM

Invalid Samples

Sample 9 has not been run, no results available.

Sample 10 has not been run, no results available.

Sample 11 has not been run, no results available.

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-02\2015-12-02_005.xad

Created: 12/2/2015 1:57:53 PM
 Modified: 12/2/2015 2:31:28 PM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 9)		Instrument	Run		12/2/2015 2:30:37 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-12-02\2015-12-02_005.xad)		Instrument	Run		12/2/2015 1:57:58 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		12/2/2015 1:57:58 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		12/2/2015 1:57:58 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		12/2/2015 1:57:58 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		12/2/2015 1:57:58 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		12/2/2015 1:57:58 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		12/2/2015 1:57:58 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1