

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...alyzer\2100 expert\data\2016-04-01\2016-04-01_003_Sarah.xad

Created: 4/1/2016 4:29:01 PM
Modified: 4/1/2016 5:11:26 PM

Electrophoresis File Run Summary

Instrument Information:

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

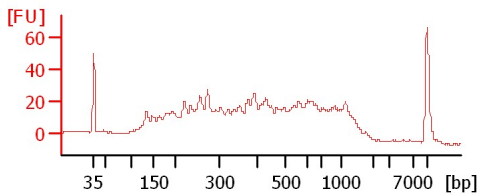
Assay Information:

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

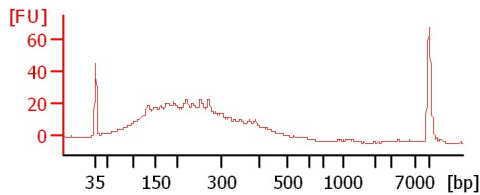
Chip Information:

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

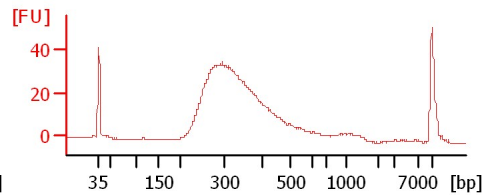
9h JI 3 cDNA



9h JI 3 sheared



9h JI 3 library



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Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
9h JI 3 cDNA		<input type="checkbox"/>				
9h JI 3 sheared		<input type="checkbox"/>				
9h JI 3 library		<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/ 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

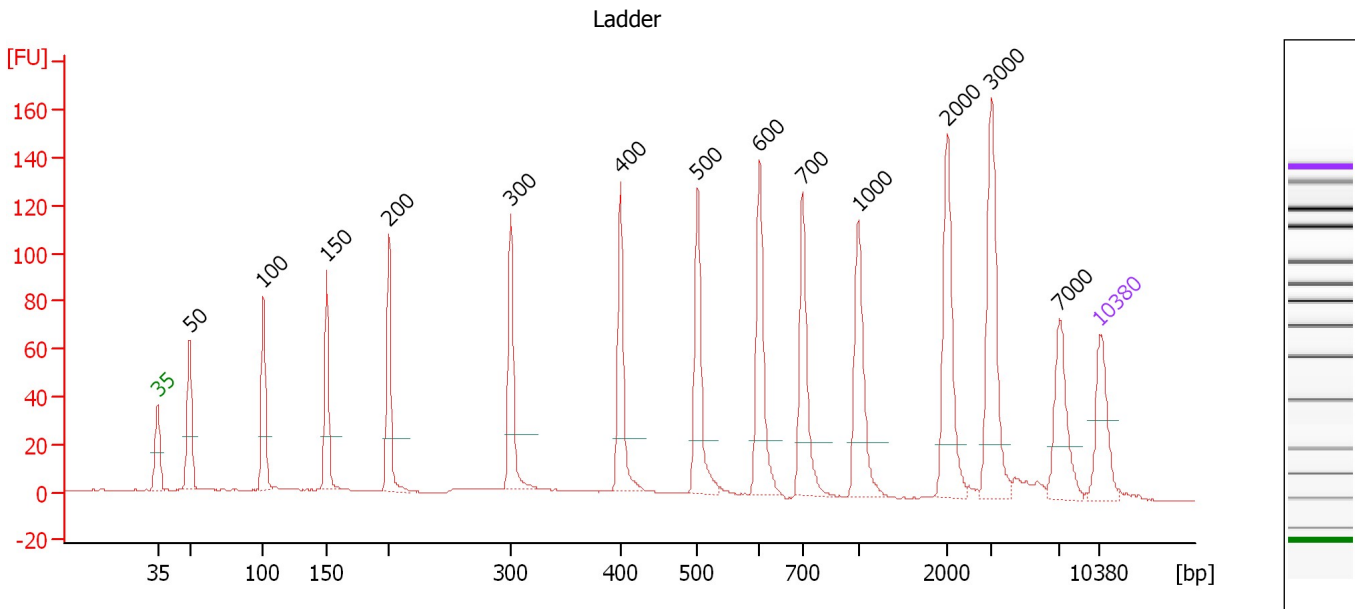
Ladder

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

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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

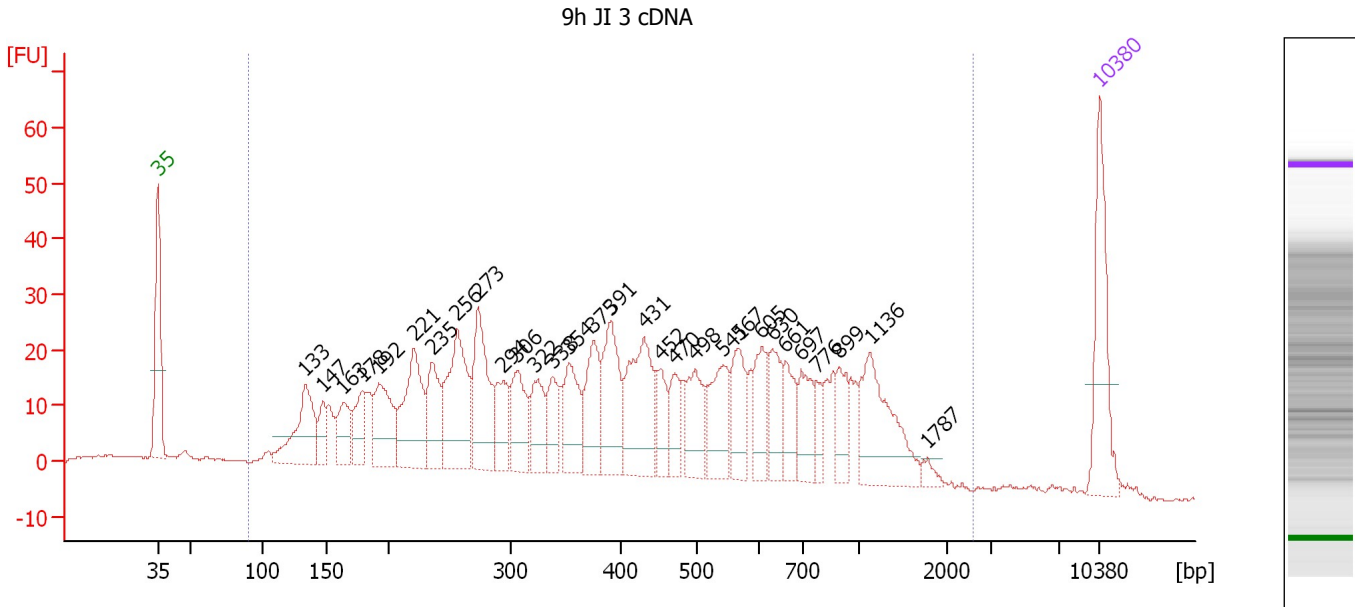
Peak table for Ladder

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	45.38
3	100	150.00	2,272.7	Ladder Peak	50.86
4	150	150.00	1,515.2	Ladder Peak	55.57
5	200	150.00	1,136.4	Ladder Peak	60.19
6	300	150.00	757.6	Ladder Peak	69.24
7	400	150.00	568.2	Ladder Peak	77.38
8	500	150.00	454.5	Ladder Peak	83.10
9	600	150.00	378.8	Ladder Peak	87.71
10	700	150.00	324.7	Ladder Peak	90.90
11	1,000	150.00	227.3	Ladder Peak	95.05
12	2,000	150.00	113.6	Ladder Peak	101.62
13	3,000	150.00	75.8	Ladder Peak	104.95
14	7,000	150.00	32.5	Ladder Peak	109.95
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 3 cDNA

Number of peaks found: 30 Corr. Area 1: 1,069.8
 Noise: 0.2

Peak table for sample 1 : 9h JI 3 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	101.57	1,156.3		53.98
3	147	28.84	296.8		55.31
4	163	48.43	449.9		56.78
5	178	46.80	398.4		58.16
6	192	85.79	677.3		59.44
7	221	111.98	769.0		62.06
8	235	63.59	409.4		63.39
9	256	134.02	792.1		65.29
10	273	106.41	590.3		66.81
11	294	47.56	245.4		68.66
12	306	64.25	318.4		69.71
13	322	49.51	233.3		70.99
14	338	39.91	178.7		72.37
15	354	65.92	282.5		73.60
16	375	65.67	265.2		75.36
17	391	86.51	335.3		76.65
18	431	109.12	383.4		79.16
19	452	31.84	106.7		80.35
20	470	34.56	111.3		81.40
21	498	51.94	158.1		82.97
22	541	63.42	177.5		85.01
23	567	51.12	136.6		86.20
24	605	42.83	107.3		87.86
25	630	41.26	99.2		88.67

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Electropherogram Summary Continued ...

... Peak table for sample 1 : 9h JI 3 cDNA

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	661	37.68	86.4		89.67
27	697	42.32	92.0		90.81
28	776	20.87	40.8		91.95
29	899	33.71	56.8		93.66
30	1,136	94.82	126.5		95.94
31	1,787	7.22	6.1		100.22
32	10,380	75.00	10.9	Upper Marker	113.00

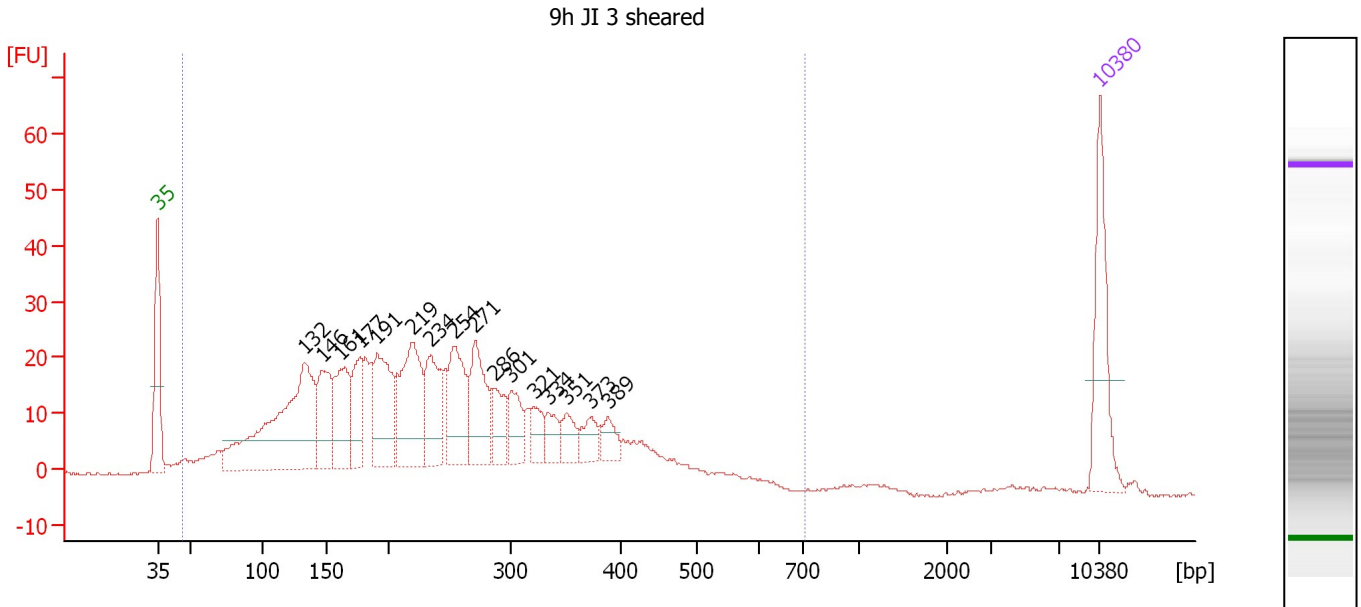
Region table for sample 1 : 9h JI 3 cDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
89	2,566	491	1,069.8	10,376.6	2,022.07	97	63.7

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Electropherogram Summary Continued ...



Overall Results for sample 2 : 9h JI 3 sheared

Number of peaks found: 16 Corr. Area 1: 806.8
 Noise: 0.2

Peak table for sample 2 : 9h JI 3 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	132	313.48	3,593.2		53.89
3	146	89.23	925.2		55.21
4	161	93.22	875.7		56.62
5	177	63.70	546.7		58.02
6	191	111.93	889.0		59.34
7	219	140.31	972.5		61.87
8	234	82.47	533.6		63.28
9	254	95.51	570.0		65.07
10	271	90.42	505.5		66.62
11	286	43.66	231.2		67.98
12	301	41.85	210.9		69.29
13	321	27.14	127.9		70.98
14	334	25.27	114.6		72.01
15	351	24.94	107.7		73.38
16	373	24.81	100.7		75.21
17	389	22.50	87.7		76.47
18	10,380	75.00	10.9	Upper Marker	113.00

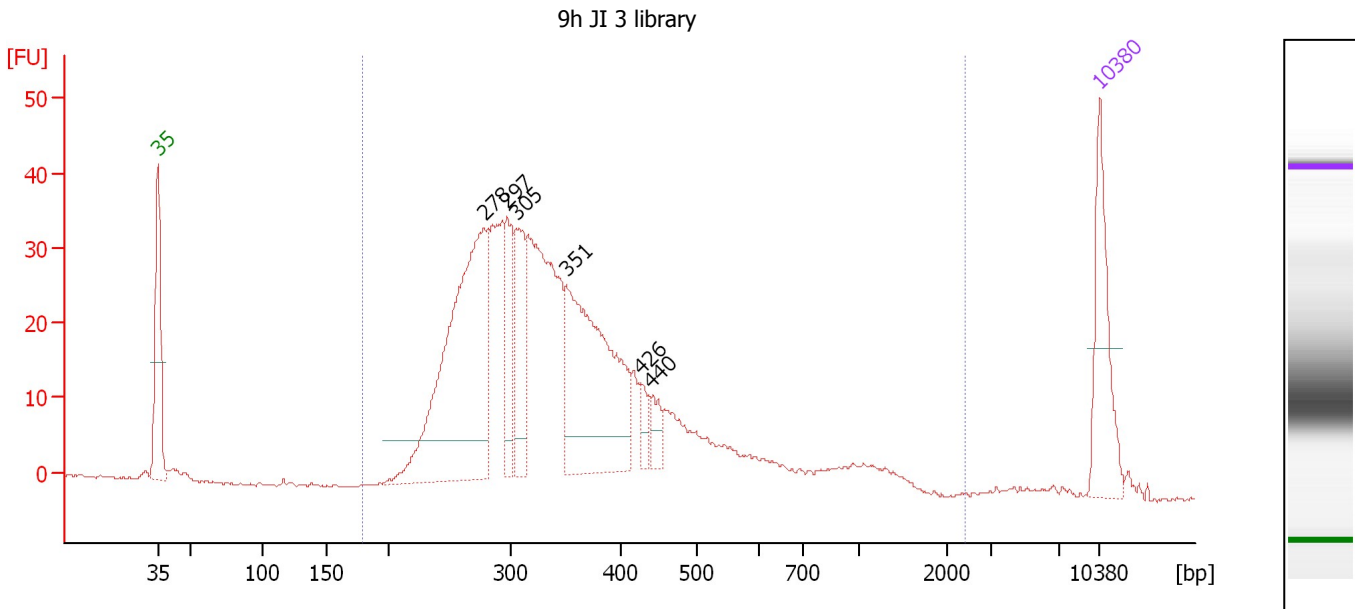
Region table for sample 2 : 9h JI 3 sheared

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
47	710	246	806.8	15,075.3	1,792.85	98	42.9

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Electropherogram Summary Continued ...



Overall Results for sample 3 : 9h JI 3 library

Number of peaks found: 6 Corr. Area 1: 703.1
 Noise: 0.2

Peak table for sample 3 : 9h JI 3 library

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	278	403.48	2,195.9		67.28
3	297	76.33	389.7		68.95
4	305	93.21	463.6		69.61
5	351	275.51	1,190.3		73.37
6	426	20.01	71.1		78.88
7	440	21.70	74.8		79.64
8	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 3 : 9h JI 3 library

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
178	2,425	382	703.1	7,289.2	1,579.33	98	48.9

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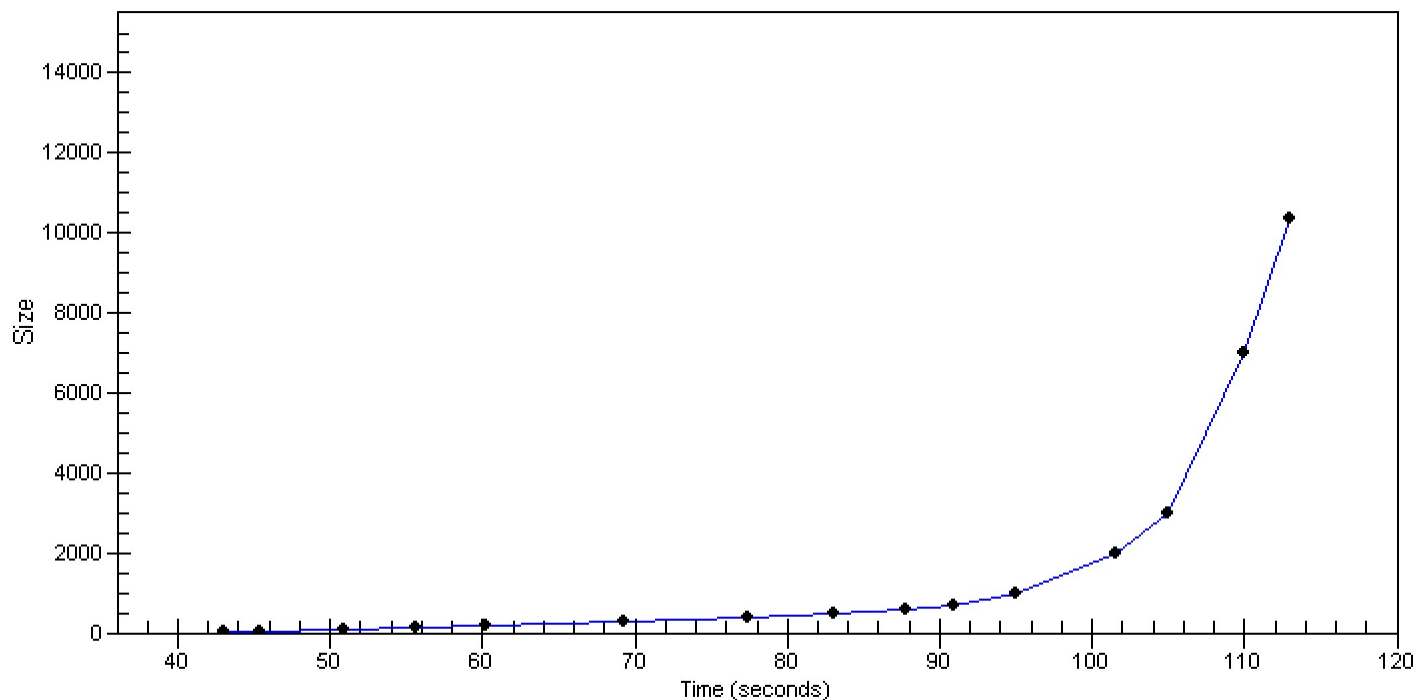
Gel Image

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Curves

Standard Curve



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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		4/1/2016 5:10:21 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-04-01\2016-04-01_003.xad)		Instrument	Run		4/1/2016 4:29:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		4/1/2016 4:29:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		4/1/2016 4:29:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		4/1/2016 4:29:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		4/1/2016 4:29:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		4/1/2016 4:29:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		4/1/2016 4:29:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1