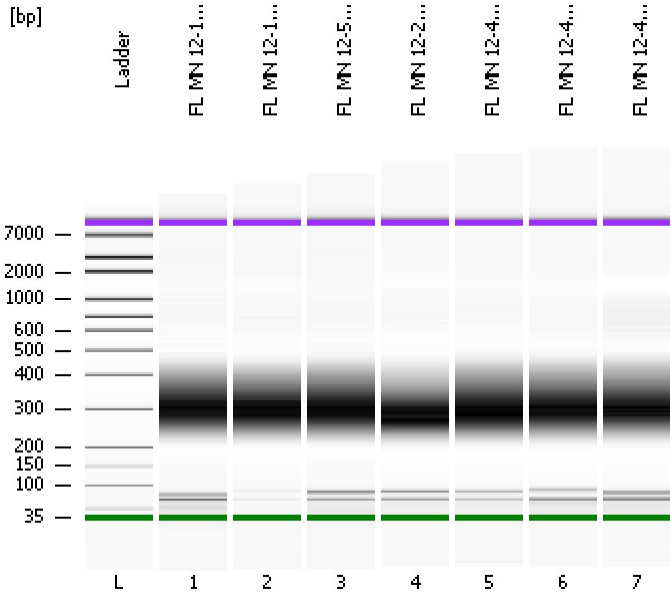


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad

Created: 3/7/2013 2:56:49 PM  
Modified: 3/7/2013 3:44:56 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

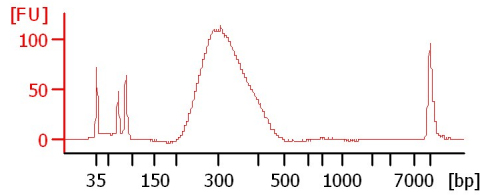
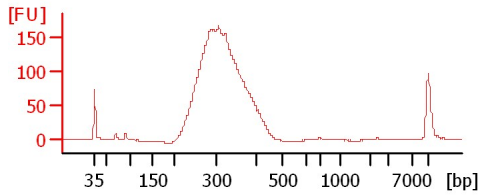
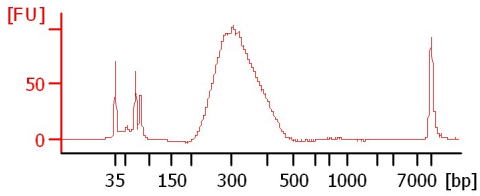
Chip Information:

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

**FL MN 12-13 mRNA Lib 1:2**

**FL MN 12-16 mRNA Lib 1:2**

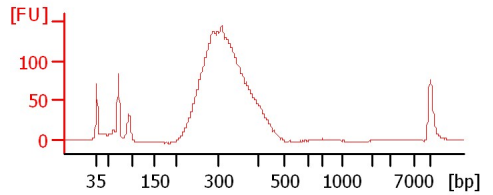
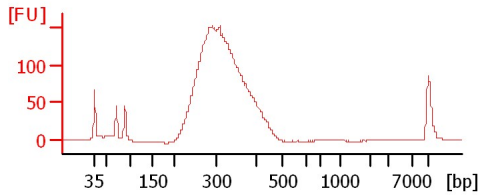
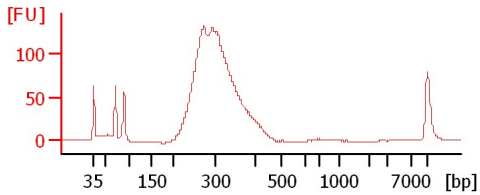
**FL MN 12-58 mRNA Lib 1:2**



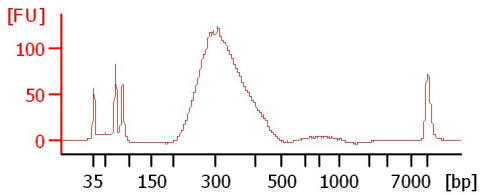
**FL MN 12-21 mRNA Lib 1:2**

**FL MN 12-43 mRNA Lib 1:2**

**FL MN 12-42 mRNA Lib 1:2**



**FL MN 12-45 mRNA Lib 1:2**



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad

Created: 3/7/2013 2:56:49 PM  
Modified: 3/7/2013 3:44:56 PM

**Electrophoresis File Run Summary (Chip Summary)**

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
FL MN 12-13 mRNA Lib 1:2		<input type="checkbox"/>	✓			
FL MN 12-16 mRNA Lib 1:2		<input type="checkbox"/>	✓			
FL MN 12-58 mRNA Lib 1:2		<input type="checkbox"/>	✓			
FL MN 12-21 mRNA Lib 1:2		<input type="checkbox"/>	✓			
FL MN 12-43 mRNA Lib 1:2		<input type="checkbox"/>	✓			
FL MN 12-42 mRNA Lib 1:2		<input type="checkbox"/>	✓			
FL MN 12-45 mRNA Lib 1:2		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

**Chip Lot #** **Reagent Kit Lot #**

**Chip Comments :**

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad

Created: 3/7/2013 2:56:49 PM  
Modified: 3/7/2013 3:44:56 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

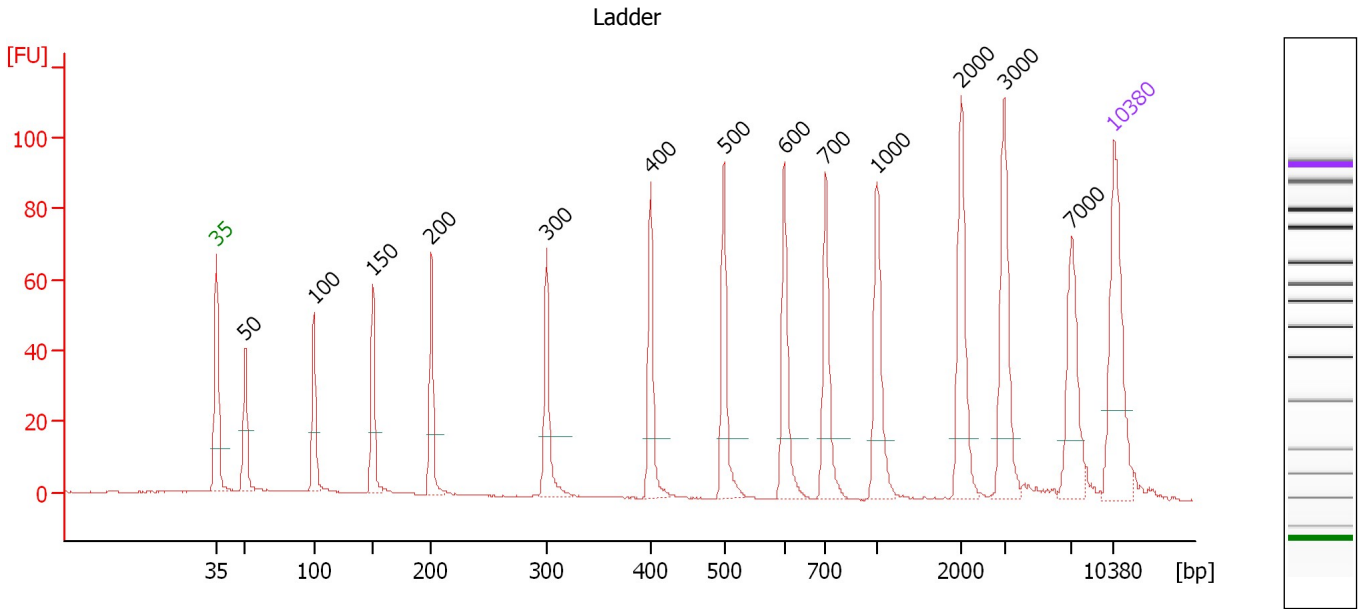
**Ladder**

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

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad

Created: 3/7/2013 2:56:49 PM  
 Modified: 3/7/2013 3:44:56 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.1

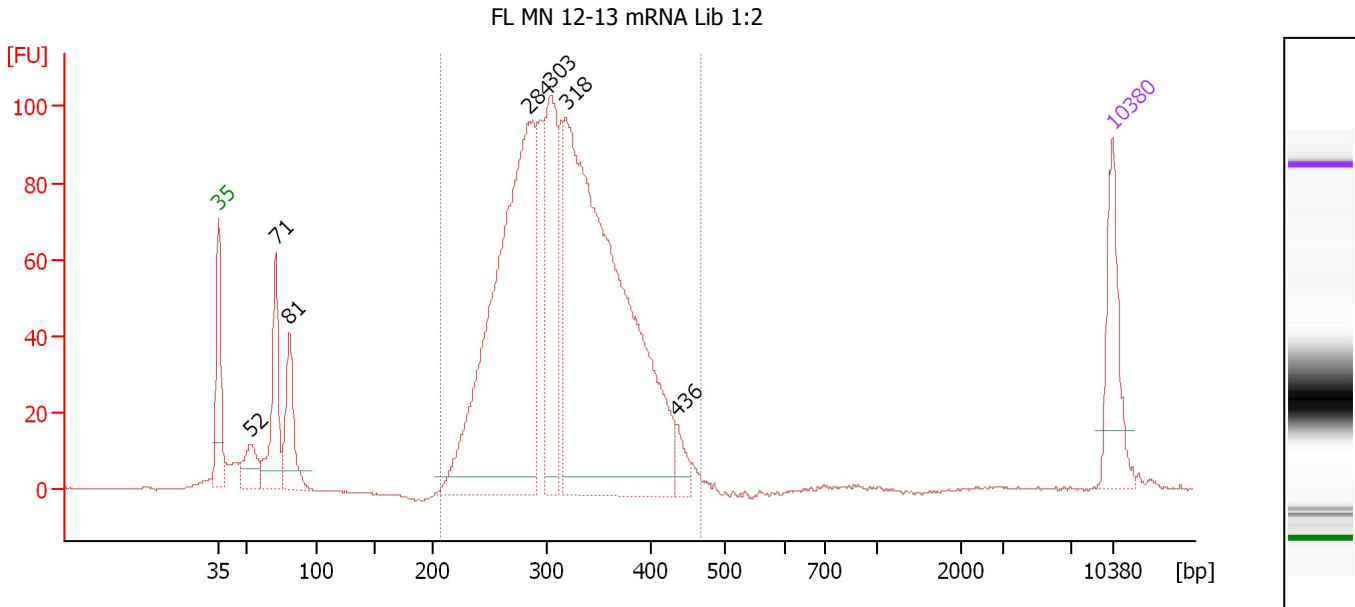
**Peak table for Ladder**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	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

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad

Created: 3/7/2013 2:56:49 PM  
 Modified: 3/7/2013 3:44:56 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : FL MN 12-13 mRNA Lib 1:2**

Number of peaks found: 7                      Corr. Area 1: 1,631.2  
 Noise: 0.2

**Peak table for sample 1 : FL MN 12-13 mRNA Lib 1:2**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	52	57.87	1,673.0	
3	71	146.05	3,130.3	
4	81	115.92	2,173.6	
5	284	761.58	4,058.8	
6	303	230.25	1,152.6	
7	318	986.83	4,700.1	
8	436	25.05	87.1	
9	10,380	75.00	10.9	Upper Marker

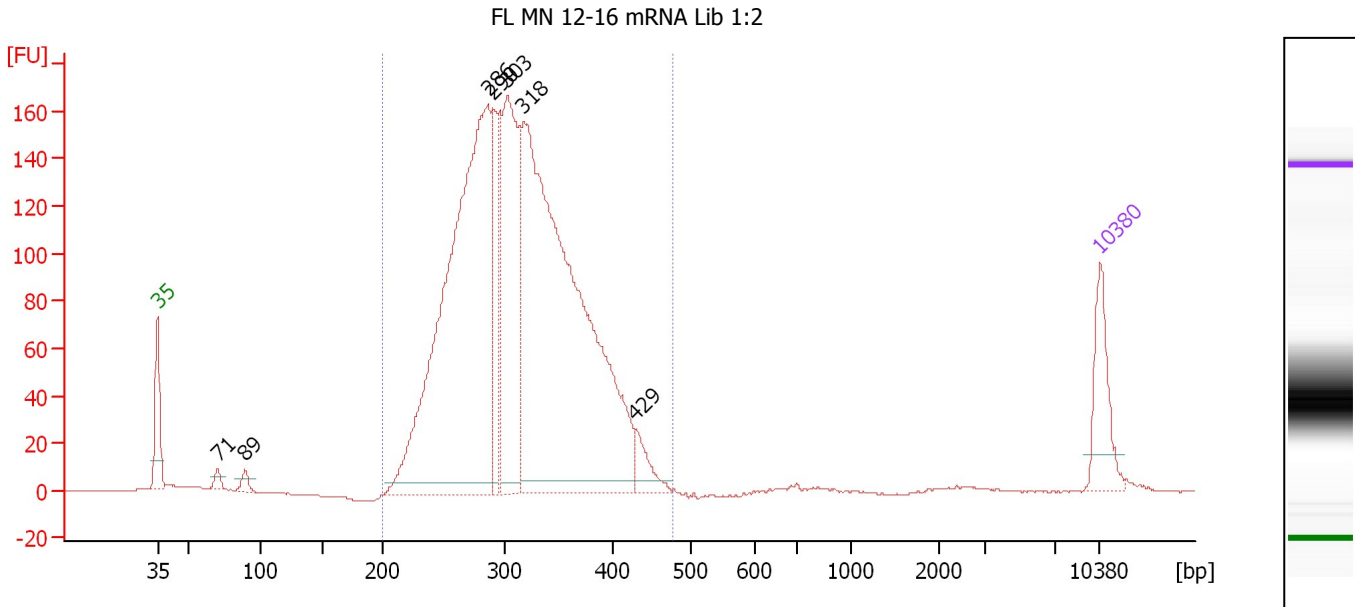
**Region table for sample 1 : FL MN 12-13 mRNA Lib 1:2**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
207	469	318	10,338.9	2,101.50	1,631.2	88	15.6	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad

Created: 3/7/2013 2:56:49 PM  
 Modified: 3/7/2013 3:44:56 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : FL MN 12-16 mRNA Lib 1:2**

Number of peaks found: 7                      Corr. Area 1: 2,607.6  
 Noise: 0.2

**Peak table for sample 2 : FL MN 12-16 mRNA Lib 1:2**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	71	14.29	306.6	
3	89	17.02	288.5	
4	286	1,133.95	6,000.1	
5	290	162.88	850.5	
6	303	400.67	2,004.6	
7	318	1,283.83	6,123.7	
8	429	42.73	151.1	
9	10,380	75.00	10.9	Upper Marker

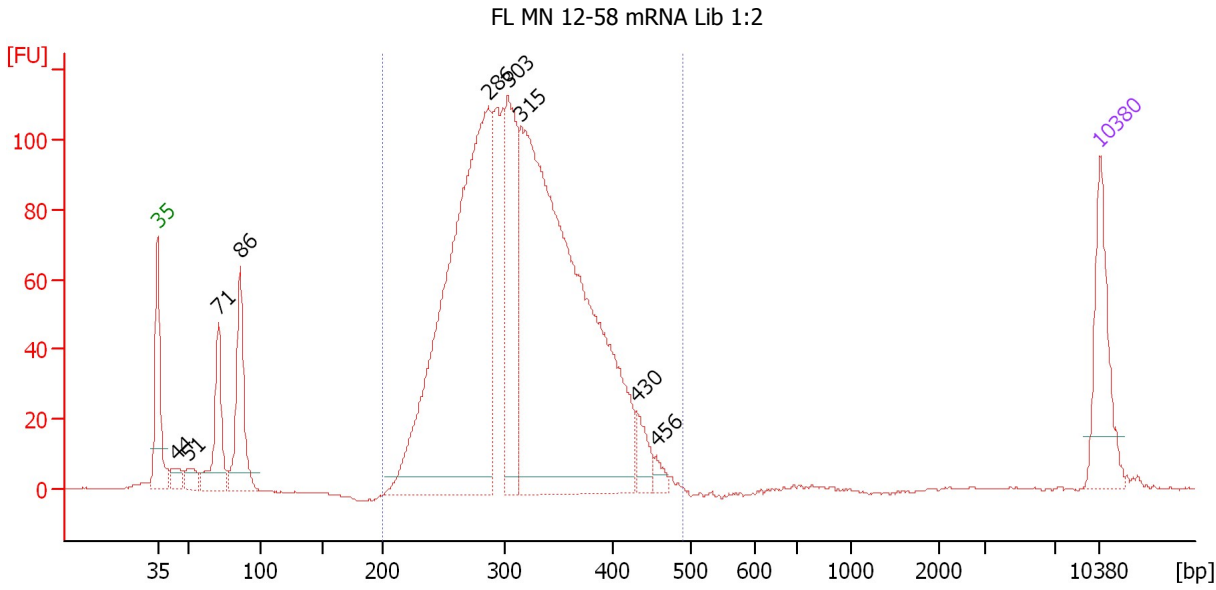
**Region table for sample 2 : FL MN 12-16 mRNA Lib 1:2**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	476	314	14,765.9	2,964.52	2,607.6	98	15.6	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad

Created: 3/7/2013 2:56:49 PM  
 Modified: 3/7/2013 3:44:56 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : FL MN 12-58 mRNA Lib 1:2**

Number of peaks found: 9                      Corr. Area 1: 1,820.7  
 Noise: 0.2

**Peak table for sample 3 : FL MN 12-58 mRNA Lib 1:2**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	44	19.09	656.2	
3	51	21.85	653.0	
4	71	106.48	2,268.6	
5	86	140.92	2,476.5	
6	286	807.65	4,271.6	
7	303	234.79	1,172.6	
8	315	977.76	4,707.1	
9	430	32.39	114.2	
10	456	12.28	40.8	
11	10,380	75.00	10.9	Upper Marker

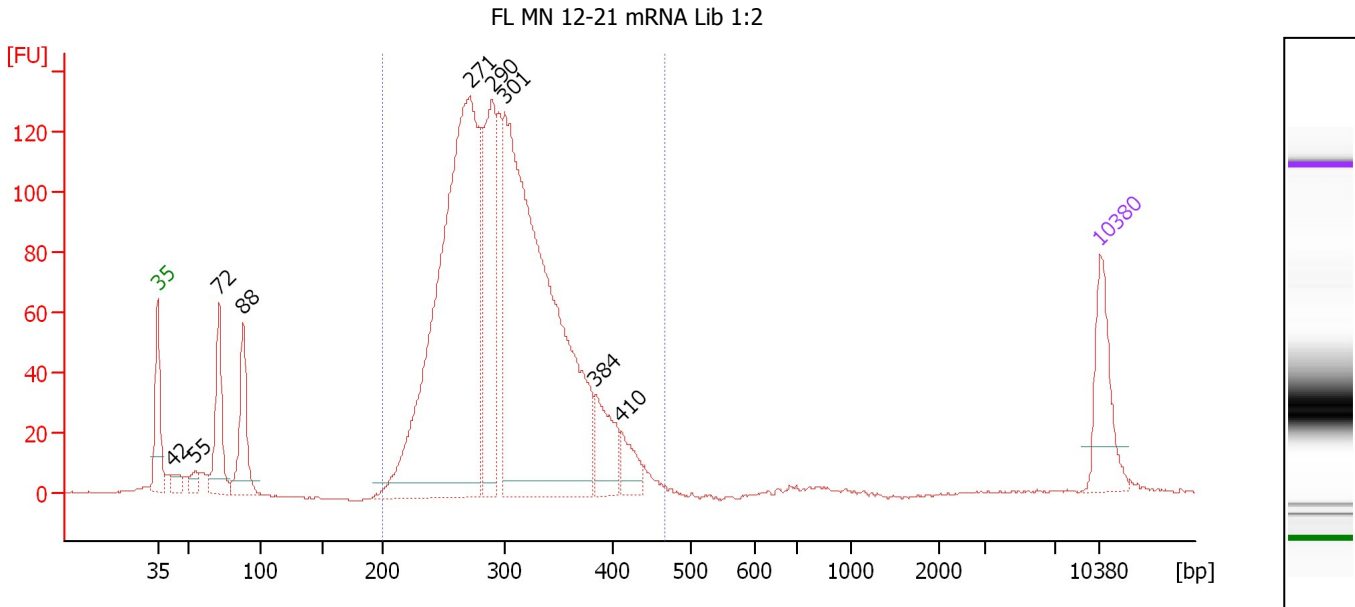
**Region table for sample 3 : FL MN 12-58 mRNA Lib 1:2**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	490	317	10,576.9	2,140.64	1,820.7	90	16.2	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad

Created: 3/7/2013 2:56:49 PM  
 Modified: 3/7/2013 3:44:56 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : FL MN 12-21 mRNA Lib 1:2**

Number of peaks found: 9                      Corr. Area 1: 1,892.6  
 Noise: 0.2

**Peak table for sample 4 : FL MN 12-21 mRNA Lib 1:2**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	42	22.93	822.7	
3	55	20.65	572.0	
4	72	133.24	2,822.7	
5	88	128.82	2,209.6	
6	271	997.39	5,566.6	
7	290	266.04	1,389.7	
8	301	998.75	5,033.2	
9	384	81.24	321.0	
10	410	39.67	146.7	
11	10,380	75.00	10.9	Upper Marker

**Region table for sample 4 : FL MN 12-21 mRNA Lib 1:2**

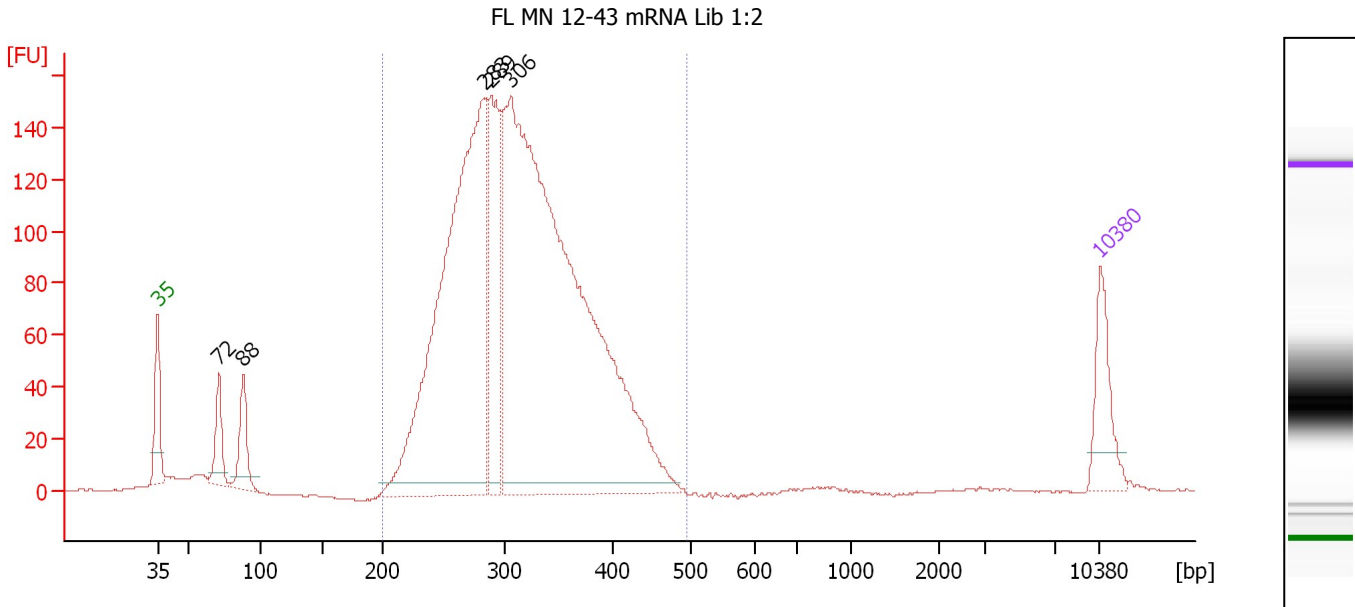
From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	467	302	12,656.6	2,453.44	1,892.6	90	15.5	Blue



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad

Created: 3/7/2013 2:56:49 PM  
 Modified: 3/7/2013 3:44:56 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : FL MN 12-43 mRNA Lib 1:2**

Number of peaks found: 5                      Corr. Area 1: 2,483.3  
 Noise: 0.3

**Peak table for sample 5 : FL MN 12-43 mRNA Lib 1:2**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	72	82.24	1,742.8	
3	88	94.15	1,617.0	
4	283	1,117.63	5,979.1	
5	289	268.20	1,406.8	
6	306	1,793.21	8,868.6	
7	10,380	75.00	10.9	Upper Marker

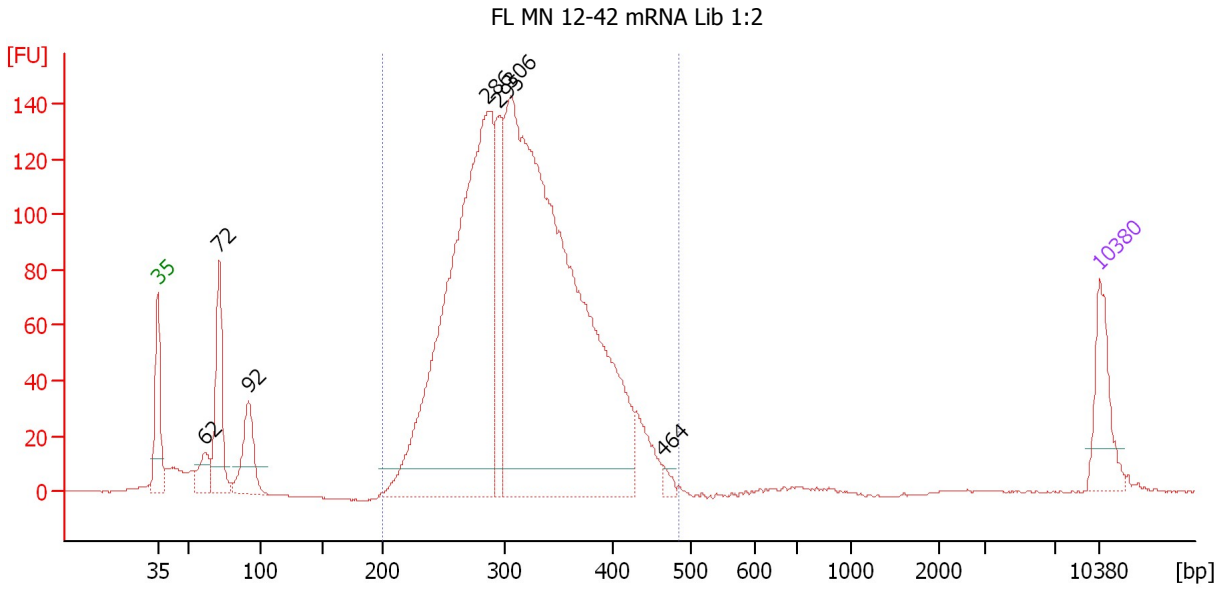
**Region table for sample 5 : FL MN 12-43 mRNA Lib 1:2**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	496	315	15,234.7	3,058.19	2,483.3	93	16.6	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad

Created: 3/7/2013 2:56:49 PM  
 Modified: 3/7/2013 3:44:56 PM

**Electropherogram Summary Continued ...**



**Setpoint Deviations for sample 6 : FL MN 12-42 mRNA Lib 1:2**

Height Threshold [FU] : 10

**Overall Results for sample 6 : FL MN 12-42 mRNA Lib 1:2**

Number of peaks found: 7                      Corr. Area 1: 2,197.2  
 Noise: 0.3

**Peak table for sample 6 : FL MN 12-42 mRNA Lib 1:2**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	62	55.40	1,352.7	
3	72	190.76	4,036.6	
4	92	121.85	2,006.7	
5	286	1,182.01	6,259.6	
6	295	174.86	898.3	
7	306	1,740.21	8,604.4	
8	464	12.80	41.8	
9	10,380	75.00	10.9	Upper Marker

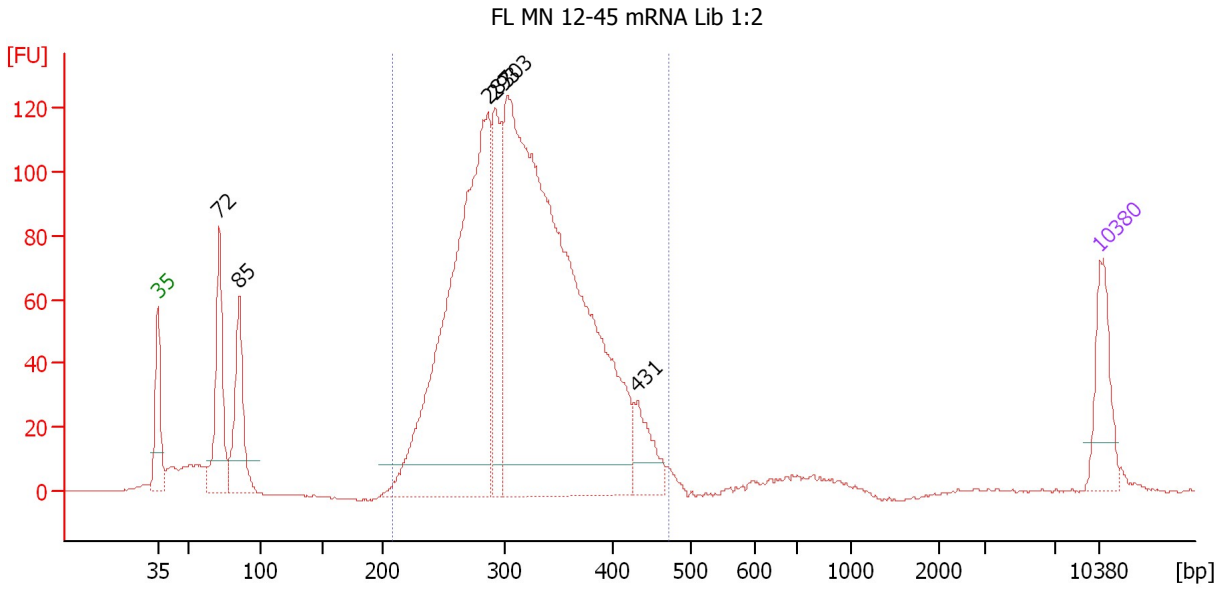
**Region table for sample 6 : FL MN 12-42 mRNA Lib 1:2**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	484	317	14,823.3	3,003.09	2,197.2	90	16.3	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad

Created: 3/7/2013 2:56:49 PM  
 Modified: 3/7/2013 3:44:56 PM

**Electropherogram Summary Continued ...**



**Setpoint Deviations for sample 7 : FL MN 12-45 mRNA Lib 1:2**

Height Threshold [FU] : 10

**Overall Results for sample 7 : FL MN 12-45 mRNA Lib 1:2**

Number of peaks found: 6                      Corr. Area 1: 1,863.7  
 Noise: 0.2

**Peak table for sample 7 : FL MN 12-45 mRNA Lib 1:2**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	72	212.02	4,472.1	
3	85	182.41	3,237.9	
4	287	942.79	4,985.5	
5	293	208.82	1,080.7	
6	303	1,582.00	7,911.5	
7	431	76.99	270.7	
8	10,380	75.00	10.9	Upper Marker

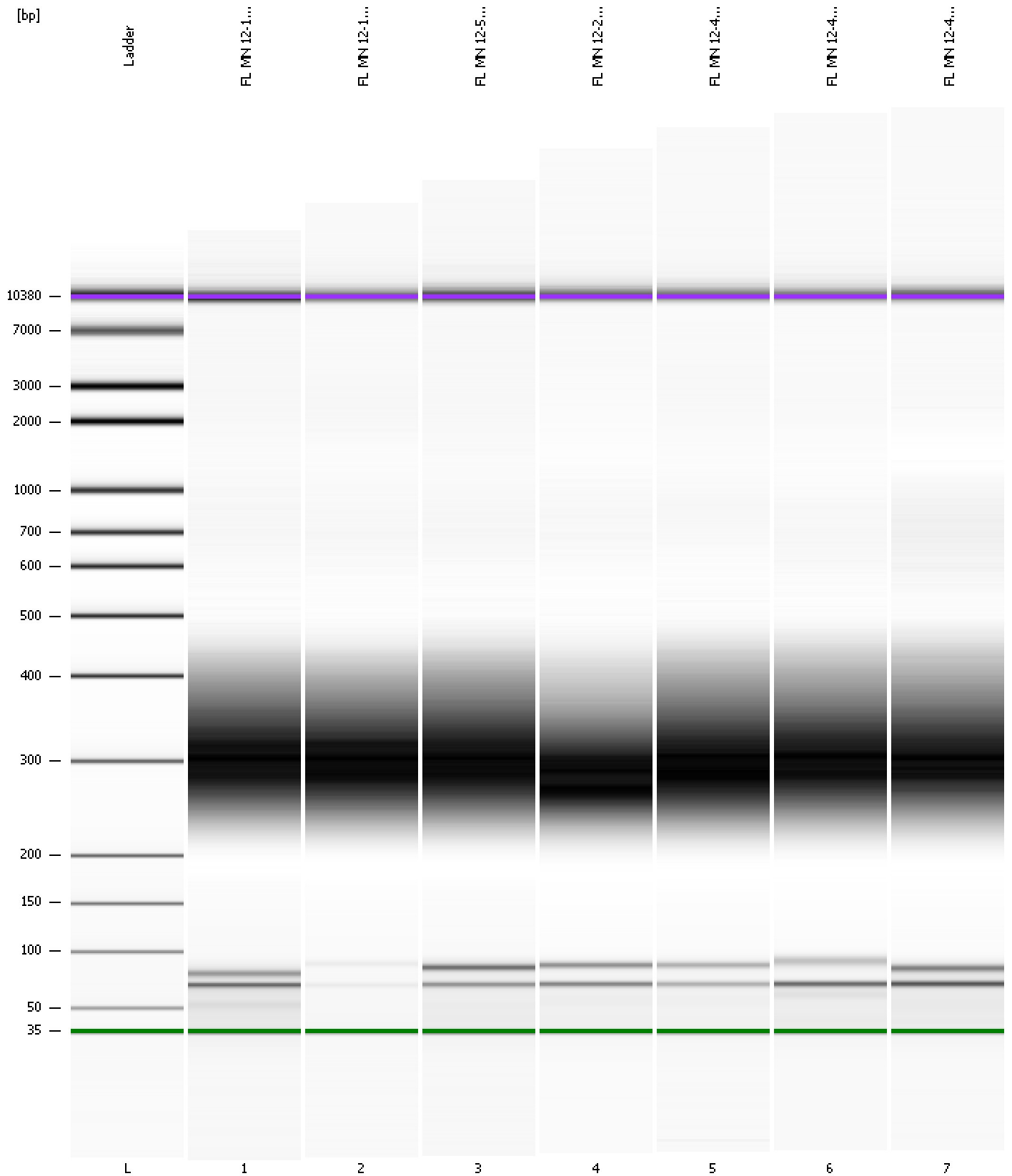
**Region table for sample 7 : FL MN 12-45 mRNA Lib 1:2**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
208	472	319	13,066.4	2,658.99	1,863.7	86	16.2	Blue

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad

Created: 3/7/2013 2:56:49 PM  
Modified: 3/7/2013 3:44:56 PM

**Gel Image**



Assay Class: High Sensitivity DNA Assay Created: 3/7/2013 2:56:49 PM  
 Data Path: C:\...ttings\Bioanalyzer\2013-03-07\2013-03-07\_002\_SSmRNA\_Lib.xad Modified: 3/7/2013 3:44:56 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		3/7/2013 3:38:08 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2013-03-07\2013-03-07_002.xad)		Instrument	Run		3/7/2013 2:56:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		3/7/2013 2:56:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		3/7/2013 2:56:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		3/7/2013 2:56:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		3/7/2013 2:56:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		3/7/2013 2:56:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		3/7/2013 2:56:55 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1