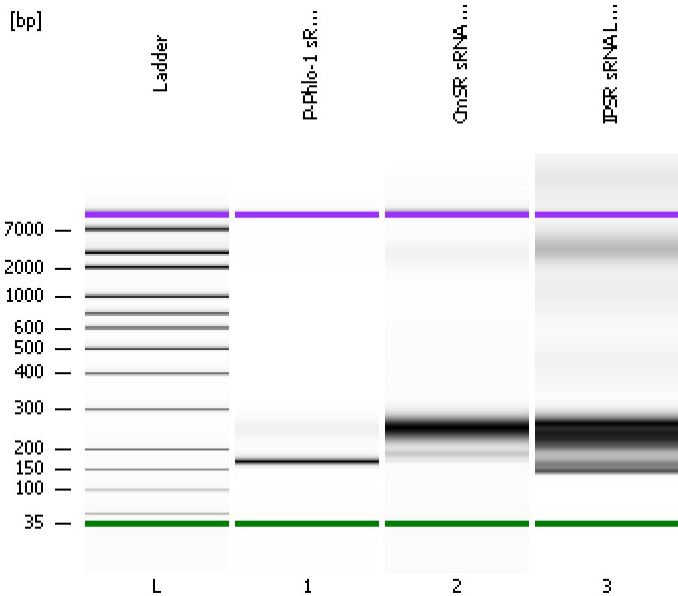


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ngs\Bioanalyzer\2013-02-04\2013-02-04\_004\_P-M\_sRNA\_Libs.xad

Created: 2/4/2013 3:37:22 PM  
Modified: 2/6/2013 3:58:46 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

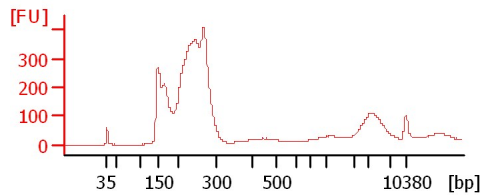
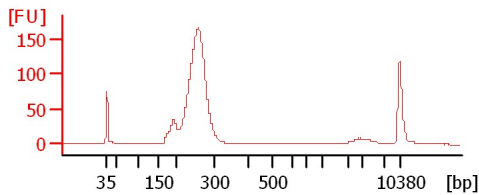
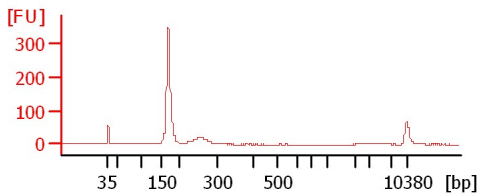
Chip Information:

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

**P-Phlo-1 sRNA Lib**

**CmSR sRNA Lib 1:2**

**IPSR sRNA Lib 1:2**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ngs\Bioanalyzer\2013-02-04\2013-02-04\_004\_P-M\_sRNA\_Libs.xad

Created: 2/4/2013 3:37:22 PM  
 Modified: 2/6/2013 3:58:46 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
P-Phlo-1 sRNA Lib		<input type="checkbox"/>				
CmSR sRNA Lib 1:2		<input type="checkbox"/>				
IPSR sRNA 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:\...ngs\Bioanalyzer\2013-02-04\2013-02-04\_004\_P-M\_sRNA\_Libs.xad

Created: 2/4/2013 3:37:22 PM  
 Modified: 2/6/2013 3:58:46 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

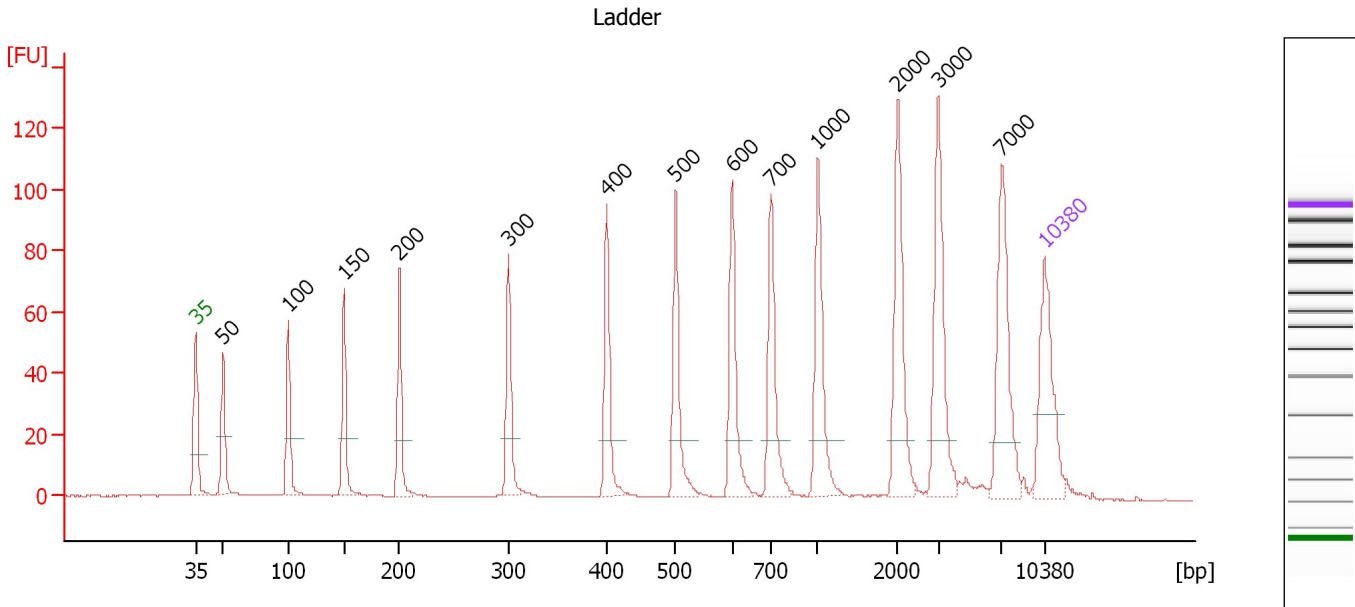
**Ladder**

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:\...ngs\Bioanalyzer\2013-02-04\2013-02-04\_004\_P-M\_sRNA\_Libs.xad

Created: 2/4/2013 3:37:22 PM  
 Modified: 2/6/2013 3:58:46 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.1

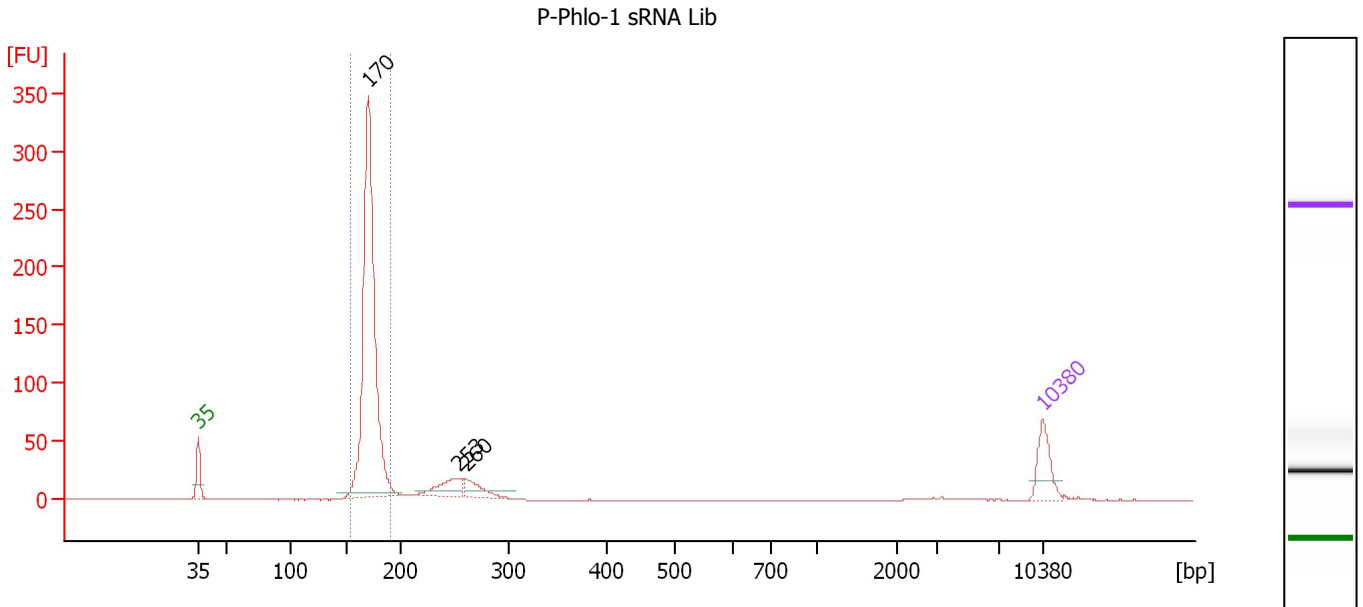
**Peak table for Ladder**

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:\...ngs\Bioanalyzer\2013-02-04\2013-02-04\_004\_P-M\_sRNA\_Libs.xad

Created: 2/4/2013 3:37:22 PM  
 Modified: 2/6/2013 3:58:46 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : P-Phlo-1 sRNA Lib**

Number of peaks found: 3                      Corr. Area 1: 699.7  
 Noise: 0.2

**Peak table for sample 1 : P-Phlo-1 sRNA Lib**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	170	1,308.31	11,640.2	
3	253	98.00	587.9	
4	260	75.74	441.5	
5	10,380	75.00	10.9	Upper Marker

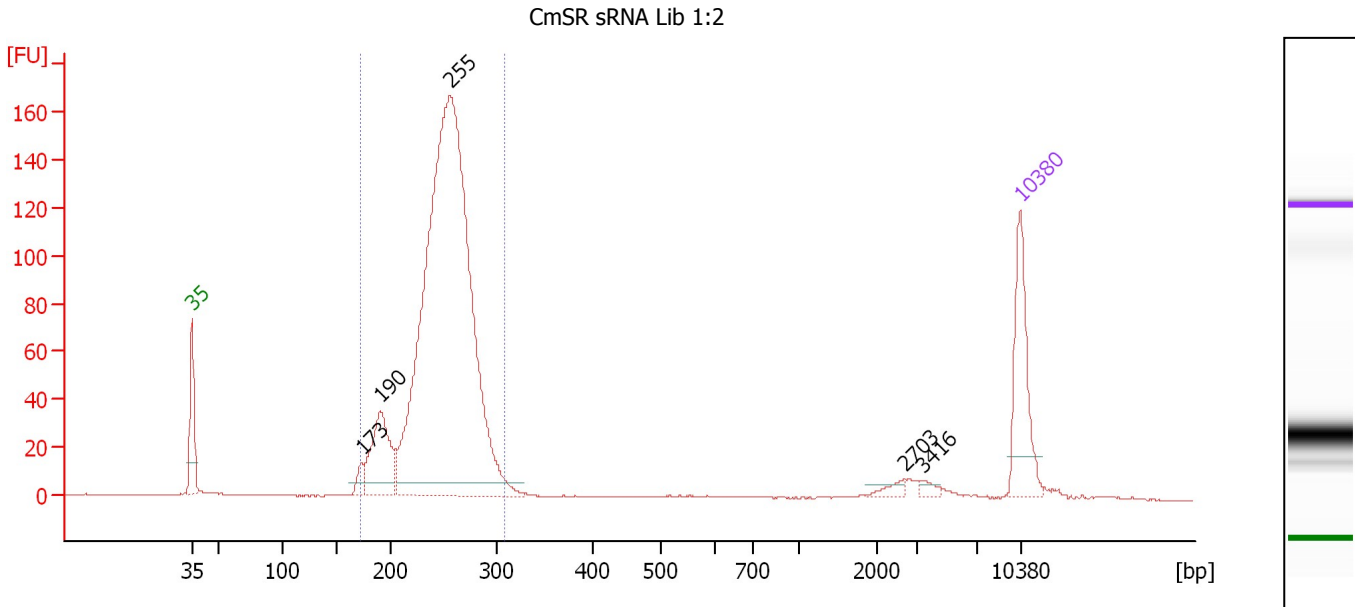
**Region table for sample 1 : P-Phlo-1 sRNA Lib**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
154	190	171	11,460.1	1,296.83	699.7	80	3.4	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ngs\Bioanalyzer\2013-02-04\2013-02-04\_004\_P-M\_sRNA\_Libs.xad

Created: 2/4/2013 3:37:22 PM  
 Modified: 2/6/2013 3:58:46 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : CmSR sRNA Lib 1:2**

Number of peaks found: 5                      Corr. Area 1: 1,349.1  
 Noise: 0.1

**Peak table for sample 2 : CmSR sRNA Lib 1:2**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	173	17.01	149.1	
3	190	115.85	924.6	
4	255	1,156.97	6,865.3	
5	2,703	7.22	4.0	
6	3,416	6.03	2.7	
7	10,380	75.00	10.9	Upper Marker

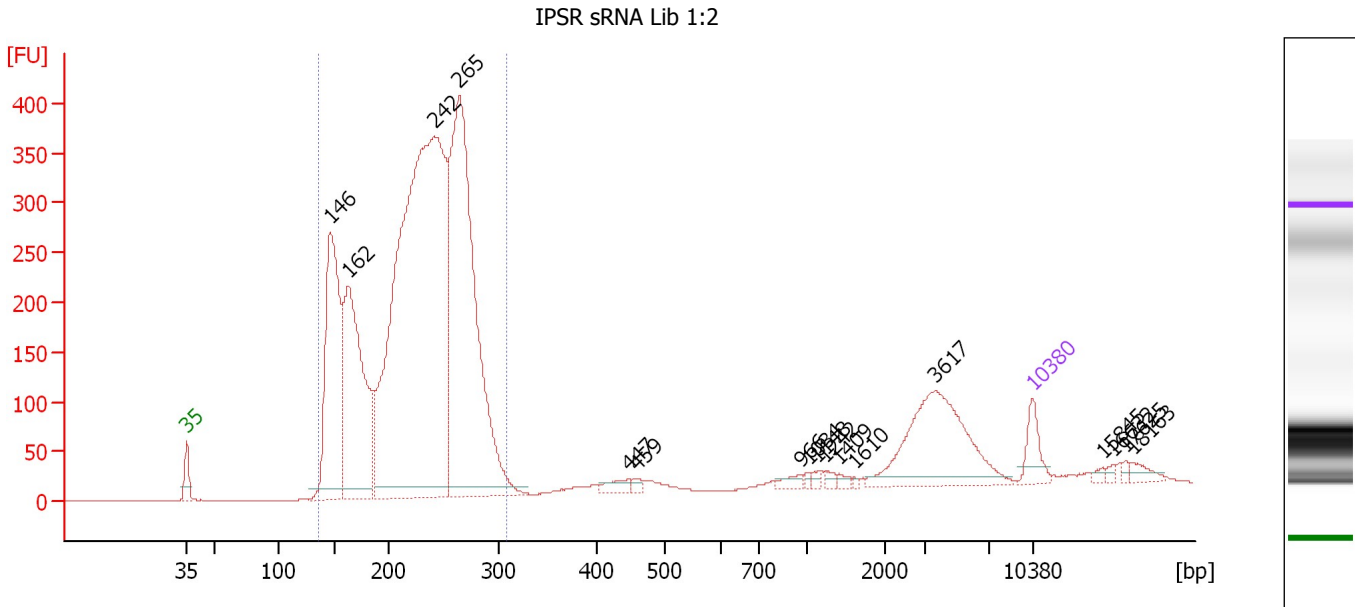
**Region table for sample 2 : CmSR sRNA Lib 1:2**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
172	307	247	8,035.2	1,297.93	1,349.1	91	10.6	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ngs\Bioanalyzer\2013-02-04\2013-02-04\_004\_P-M\_sRNA\_Libs.xad

Created: 2/4/2013 3:37:22 PM  
 Modified: 2/6/2013 3:58:46 PM

**Electropherogram Summary Continued ...**



**Setpoint Deviations for sample 3 : IPSP sRNA Lib 1:2**

Height Threshold [FU] : 10

**Overall Results for sample 3 : IPSP sRNA Lib 1:2**

Number of peaks found: 17                      Corr. Area 1: 5,973.2  
 Noise: 0.1

**Peak table for sample 3 : IPSP sRNA Lib 1:2**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	146	1,117.77	11,577.5	
3	162	1,204.35	11,243.2	
4	242	3,861.47	24,178.6	
5	265	2,123.49	12,163.3	
6	447	45.76	155.1	
7	459	18.78	61.9	
8	966	29.86	46.8	
9	1,034	9.61	14.1	
10	1,148	14.61	19.3	
11	1,242	15.38	18.8	
12	1,409	14.16	15.2	
13	1,610	5.63	5.3	
14	3,617	462.50	193.7	
15	10,380	75.00	10.9	Upper Marker
16	15,845	0.00	0.0	
17	16,632	0.00	0.0	
18	17,625	0.00	0.0	
19	18,163	0.00	0.0	

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ngs\Bioanalyzer\2013-02-04\2013-02-04\_004\_P-M\_sRNA\_Libs.xad

Created: 2/4/2013 3:37:22 PM  
Modified: 2/6/2013 3:58:46 PM

**Electropherogram Summary Continued ...**

... Region table for sample 3 : IPSR sRNA Lib 1:2

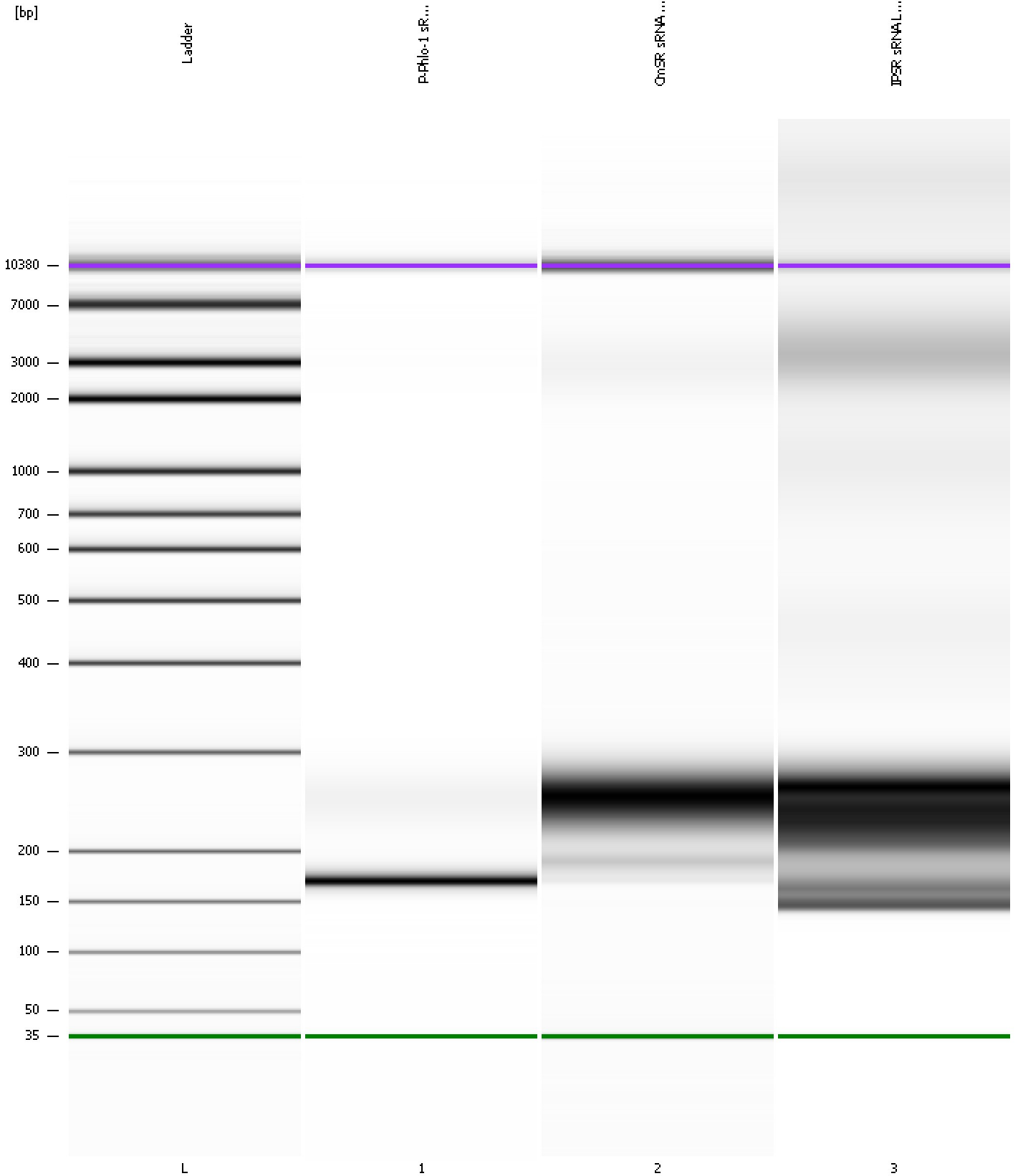
From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
135	309	224	59,423.2	8,296.07	5,973.2	87	18.9	■



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ngs\Bioanalyzer\2013-02-04\2013-02-04\_004\_P-M\_sRNA\_Libs.xad

Created: 2/4/2013 3:37:22 PM  
Modified: 2/6/2013 3:58:46 PM

**Gel Image**



Assay Class: High Sensitivity DNA Assay Created: 2/4/2013 3:37:22 PM  
 Data Path: C:\...ngs\Bioanalyzer\2013-02-04\2013-02-04\_004\_P-M\_sRNA\_Libs.xad Modified: 2/6/2013 3:58:46 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		2/4/2013 4:18:41 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2013-02-04\2013-02-04_004.xad)		Instrument	Run		2/4/2013 3:37:28 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		2/4/2013 3:37:28 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		2/4/2013 3:37:28 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		2/4/2013 3:37:28 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		2/4/2013 3:37:28 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		2/4/2013 3:37:28 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		2/4/2013 3:37:28 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1