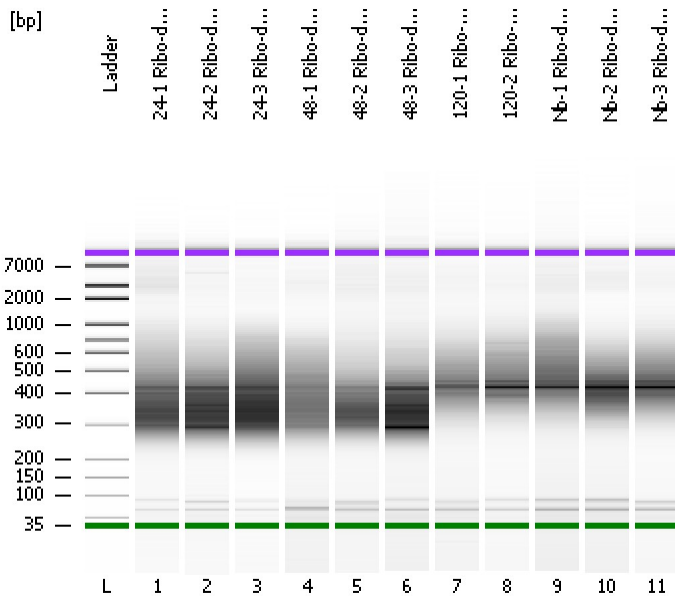


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

Created: 7/14/2014 9:23:23 AM  
Modified: 7/14/2014 10:04:46 AM

### Electrophoresis File Run Summary



#### Instrument Information:

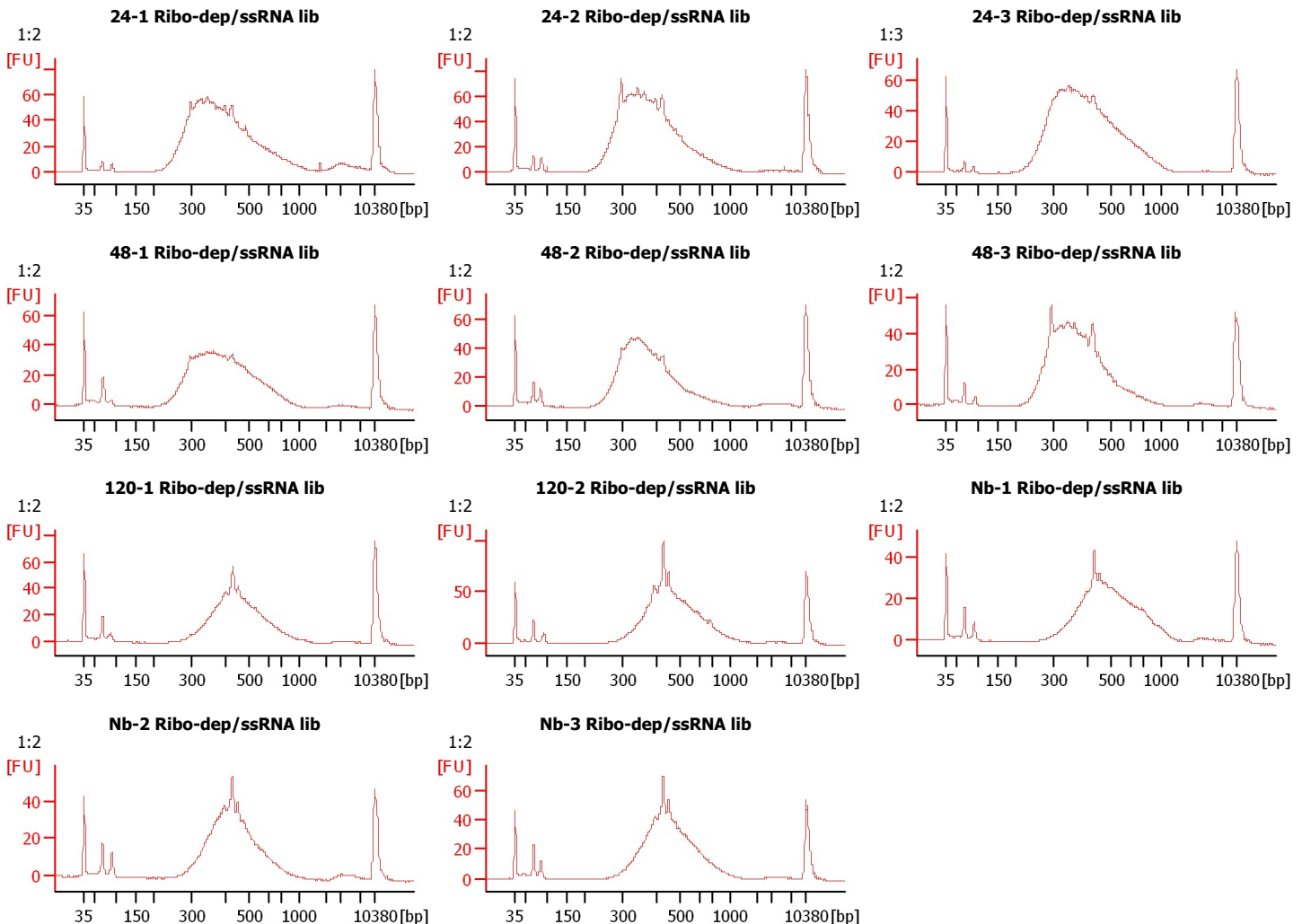
Instrument Name: DE13701086      Firmware: C.01.069  
Serial#: DE13701086      Type: G2938B

#### 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:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

Created: 7/14/2014 9:23:23 AM  
 Modified: 7/14/2014 10:04:46 AM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
24-1 Ribo-dep/ssRNA lib	1:2	<input type="checkbox"/>	✓			
24-2 Ribo-dep/ssRNA lib	1:2	<input type="checkbox"/>	✓			
24-3 Ribo-dep/ssRNA lib	1:3	<input type="checkbox"/>	✓			
48-1 Ribo-dep/ssRNA lib	1:2	<input type="checkbox"/>	✓			
48-2 Ribo-dep/ssRNA lib	1:2	<input type="checkbox"/>	✓			
48-3 Ribo-dep/ssRNA lib	1:2	<input type="checkbox"/>	✓			
120-1 Ribo-dep/ssRNA lib	1:2	<input type="checkbox"/>	✓			
120-2 Ribo-dep/ssRNA lib	1:2	<input type="checkbox"/>	✓			
Nb-1 Ribo-dep/ssRNA lib	1:2	<input type="checkbox"/>	✓			
Nb-2 Ribo-dep/ssRNA lib	1:2	<input type="checkbox"/>	✓			
Nb-3 Ribo-dep/ssRNA 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:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

Created: 7/14/2014 9:23:23 AM  
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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

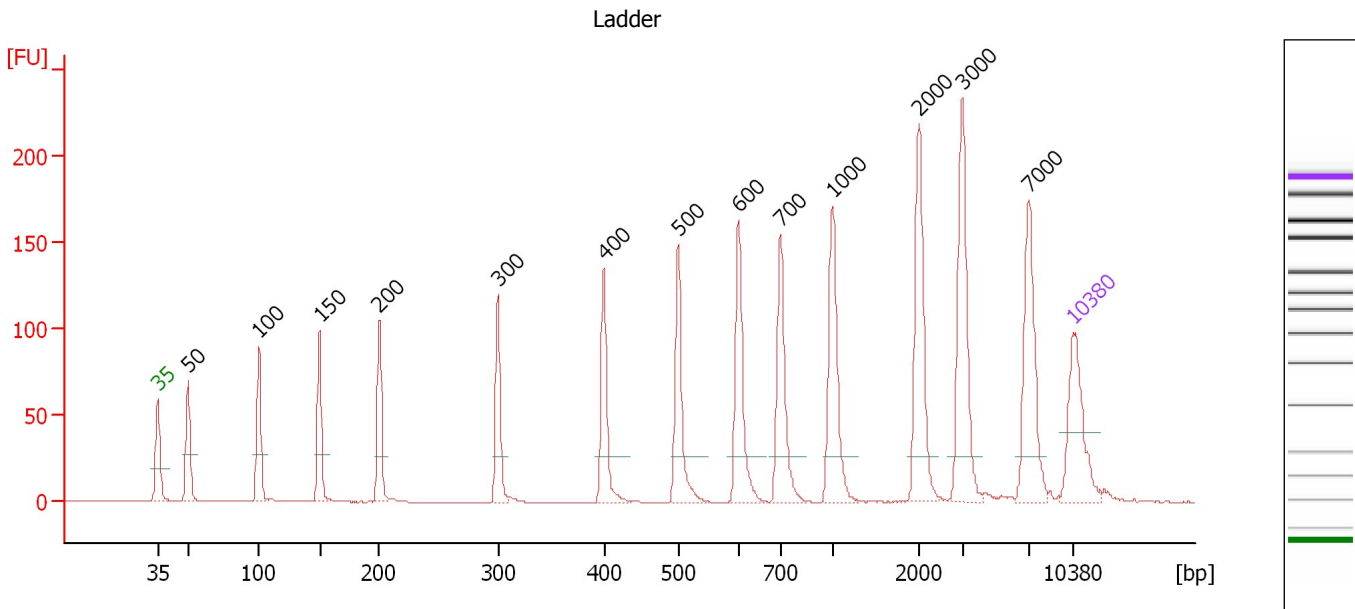
**Ladder**

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

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

Created: 7/14/2014 9:23:23 AM  
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**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.2

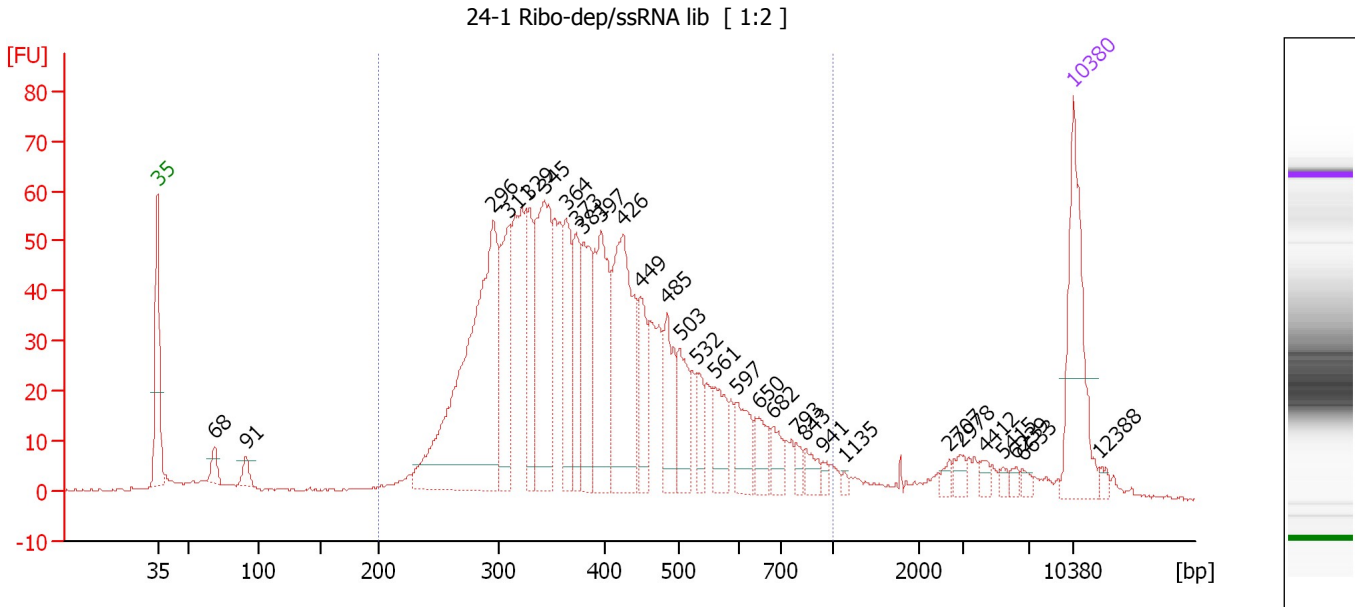
**Peak table for Ladder**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
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:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

Created: 7/14/2014 9:23:23 AM  
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**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : 24-1 Ribo-dep/ssRNA lib**

Number of peaks found: 30                      Corr. Area 1: 1,380.5  
 Noise: 0.2

**Peak table for sample 1 : 24-1 Ribo-dep/ssRNA lib**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	68	14.21	314.5	
3	91	12.81	214.2	
4	296	321.31	1,645.0	
5	311	81.65	397.7	
6	329	61.45	282.6	
7	345	147.06	646.8	
8	364	70.41	293.1	
9	373	54.48	221.0	
10	381	73.90	294.1	
11	397	105.00	400.8	
12	426	141.25	502.6	
13	449	43.58	147.1	
14	485	47.84	149.5	
15	503	40.90	123.3	
16	532	20.65	58.8	
17	561	33.26	89.8	
18	597	29.22	74.1	
19	650	21.71	50.6	
20	682	18.29	40.6	
21	793	7.91	15.1	
22	843	11.90	21.4	
23	941	4.51	7.3	
24	1,135	2.60	3.5	
25	2,707	4.79	2.7	
26	2,978	7.32	3.7	

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad


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### Electropherogram Summary Continued ...

#### ... Peak table for sample 1 : 24-1 Ribo-dep/ssRNA lib

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	4,412	5.16	1.8	
28	5,415	3.44	1.0	
29	6,239	3.86	0.9	
30	6,633	4.21	1.0	
31	10,380	75.00	10.9	Upper Marker
32	12,388	0.00	0.0	

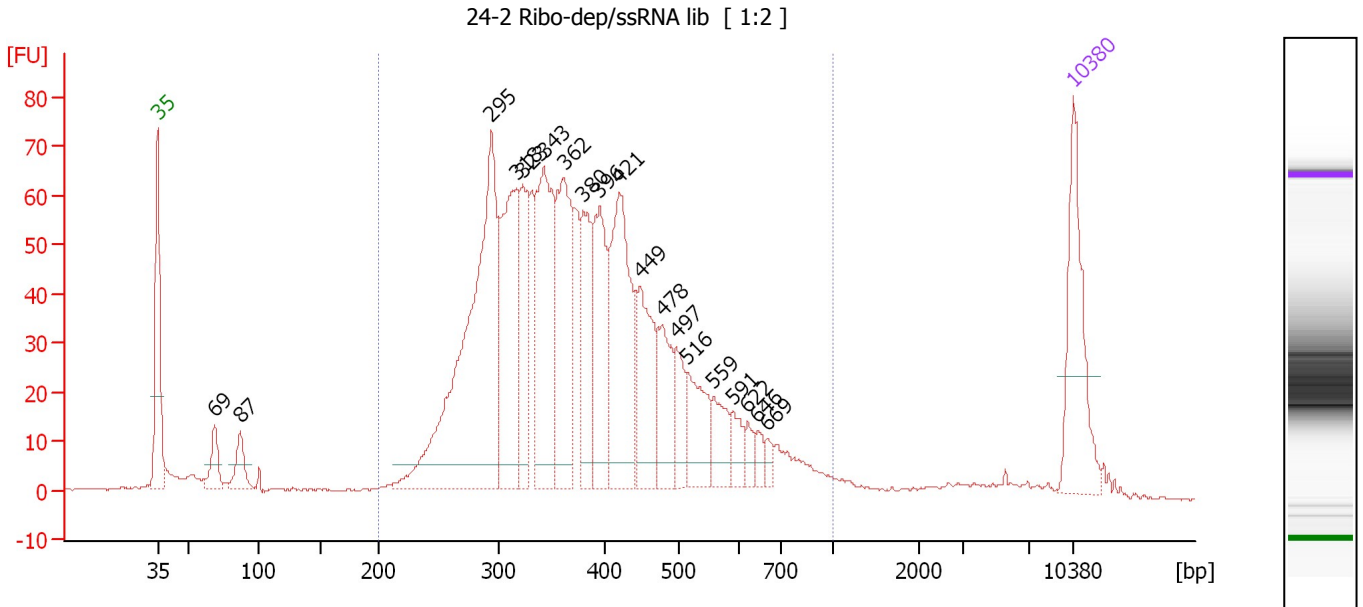
#### Region table for sample 1 : 24-1 Ribo-dep/ssRNA lib

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
200	1,000	410	7,153.9	1,732.90	 1,380.5	89	31.6

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

Created: 7/14/2014 9:23:23 AM  
 Modified: 7/14/2014 10:04:46 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : 24-2 Ribo-dep/ssRNA lib**

Number of peaks found: 19                      Corr. Area 1: 1,484.2  
 Noise: 0.2

**Peak table for sample 2 : 24-2 Ribo-dep/ssRNA lib**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	69	29.28	646.2	
3	87	29.98	523.9	
4	295	368.61	1,895.1	
5	318	159.65	761.6	
6	323	79.52	372.8	
7	343	163.35	720.8	
8	362	138.80	580.3	
9	380	75.19	299.5	
10	396	96.48	369.2	
11	421	150.89	543.5	
12	449	81.11	273.9	
13	478	55.28	175.1	
14	497	31.91	97.2	
15	516	50.52	148.4	
16	559	31.02	84.1	
17	591	17.90	45.9	
18	622	11.89	28.9	
19	646	9.28	21.7	
20	669	7.68	17.4	
21	10,380	75.00	10.9	Upper Marker

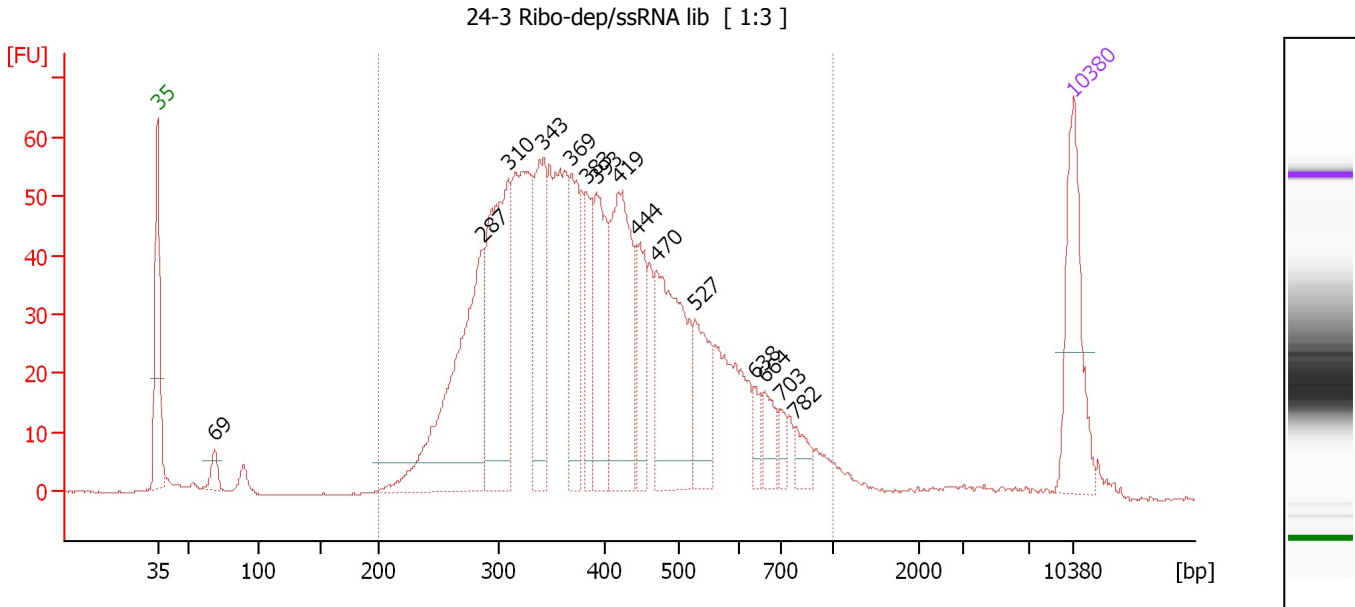
**Region table for sample 2 : 24-2 Ribo-dep/ssRNA lib**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
200	1,000	396	7,447.6	1,769.00	1,484.2	92	29.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

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 Modified: 7/14/2014 10:04:46 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : 24-3 Ribo-dep/ssRNA lib**

Number of peaks found: 15                      Corr. Area 1: 1,420.7  
 Noise: 0.1

**Peak table for sample 3 : 24-3 Ribo-dep/ssRNA lib**

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	69	16.75	369.1	
3	287	256.25	1,354.1	
4	310	211.96	1,035.5	
5	343	116.15	513.0	
6	369	94.95	389.7	
7	383	58.51	231.3	
8	393	98.50	379.8	
9	419	159.50	576.6	
10	444	59.74	203.8	
11	470	157.37	507.1	
12	527	64.60	185.7	
13	638	14.07	33.4	
14	664	23.21	53.0	
15	703	10.95	23.6	
16	782	17.20	33.3	
17	10,380	75.00	10.9	Upper Marker

**Region table for sample 3 : 24-3 Ribo-dep/ssRNA lib**

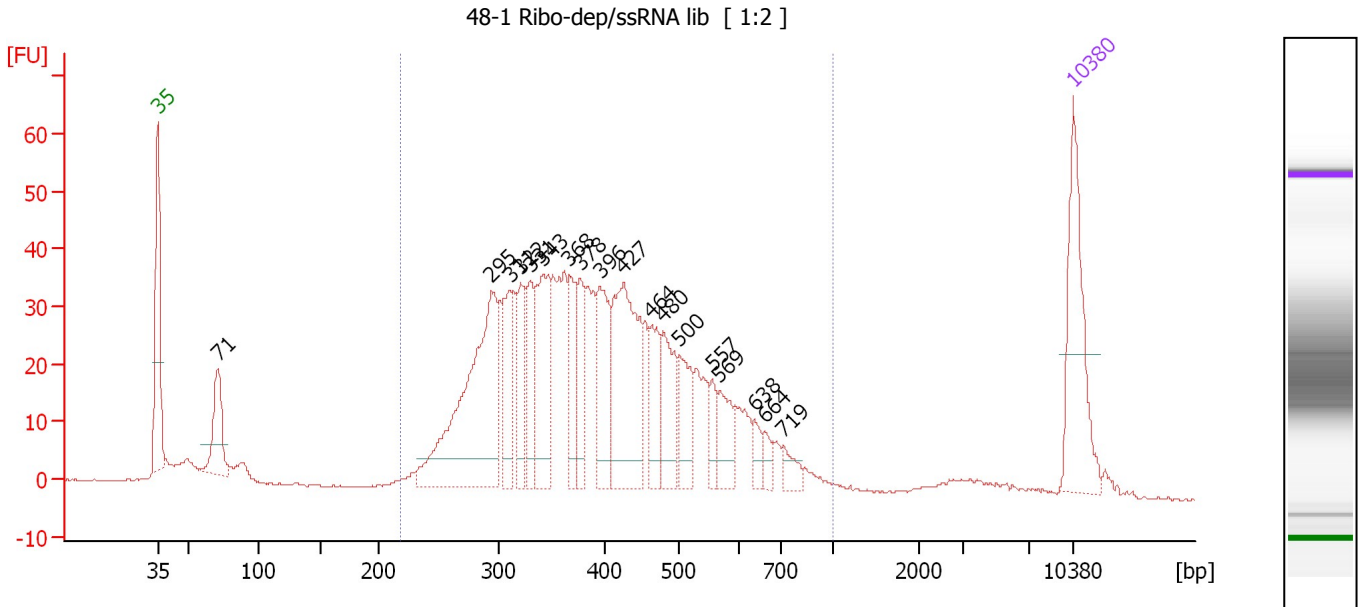
From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/µl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
200	1,000	417	8,021.6	1,964.24	1,420.7	95	31.6



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : 48-1 Ribo-dep/ssRNA lib**

Number of peaks found: 18                      Corr. Area 1: 926.6  
 Noise: 0.2

**Peak table for sample 4 : 48-1 Ribo-dep/ssRNA lib**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	71	60.86	1,300.3	
3	295	225.85	1,161.2	
4	311	55.58	270.4	
5	322	46.11	217.1	
6	331	45.70	209.5	
7	343	91.05	402.5	
8	368	43.02	177.3	
9	378	41.25	165.3	
10	396	63.58	243.2	
11	427	136.98	486.4	
12	464	40.20	131.3	
13	480	53.83	170.1	
14	500	42.91	129.9	
15	557	16.94	46.1	
16	569	30.36	80.9	
17	638	10.56	25.1	
18	664	10.12	23.1	
19	719	12.48	26.3	
20	10,380	75.00	10.9	Upper Marker

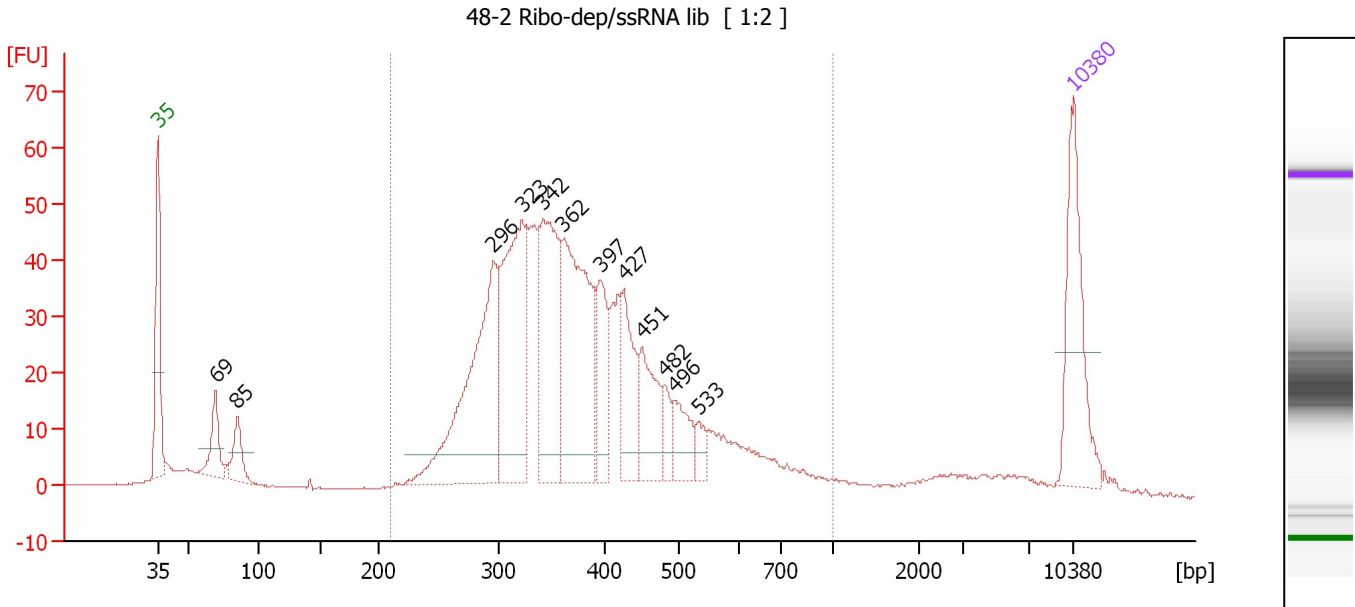
**Region table for sample 4 : 48-1 Ribo-dep/ssRNA lib**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
217	1,000	414	5,332.7	1,317.65	926.6	90	28.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

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 Modified: 7/14/2014 10:04:46 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : 48-2 Ribo-dep/ssRNA lib**

Number of peaks found: 12                      Corr. Area 1: 936.1  
 Noise: 0.1

**Peak table for sample 5 : 48-2 Ribo-dep/ssRNA lib**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	69	41.19	900.5	
3	85	33.92	606.6	
4	296	215.29	1,102.4	
5	323	178.16	835.6	
6	342	139.69	618.1	
7	362	181.09	758.4	
8	397	57.07	217.8	
9	427	61.17	217.1	
10	451	56.11	188.5	
11	482	18.84	59.2	
12	496	32.74	100.1	
13	533	13.13	37.3	
14	10,380	75.00	10.9	Upper Marker

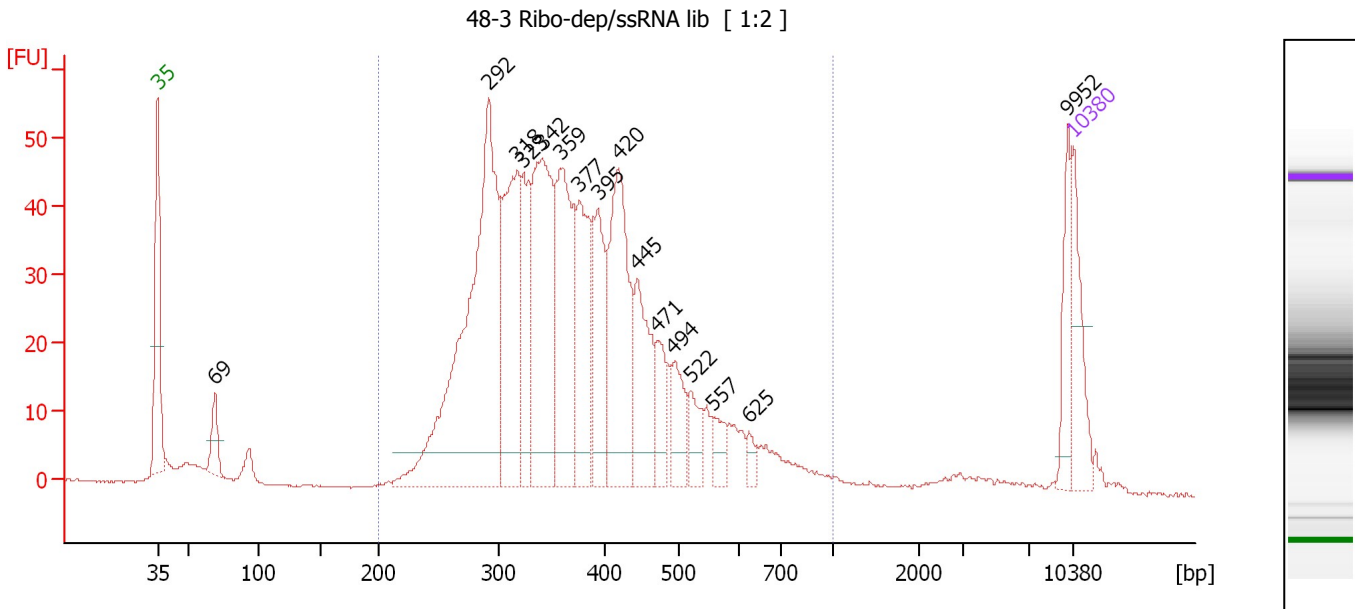
**Region table for sample 5 : 48-2 Ribo-dep/ssRNA lib**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
209	1,000	392	5,265.9	1,251.68	936.1	88	28.4

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

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 Modified: 7/14/2014 10:04:46 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : 48-3 Ribo-dep/ssRNA lib**

Number of peaks found: 16                      Corr. Area 1: 1,028.4  
 Noise: 0.2

**Peak table for sample 6 : 48-3 Ribo-dep/ssRNA lib**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	69	54.27	1,194.2	
3	292	720.53	3,737.4	
4	318	286.19	1,361.5	
5	323	130.07	609.7	
6	342	339.61	1,506.1	
7	359	251.23	1,058.9	
8	377	169.47	681.7	
9	395	146.71	563.4	
10	420	269.18	970.7	
11	445	148.08	504.5	
12	471	63.02	202.7	
13	494	67.66	207.5	
14	522	46.05	133.7	
15	557	32.57	88.5	
16	625	16.32	39.6	
17	9,952	63.99	9.7	
18	10,380	75.00	10.9	Upper Marker

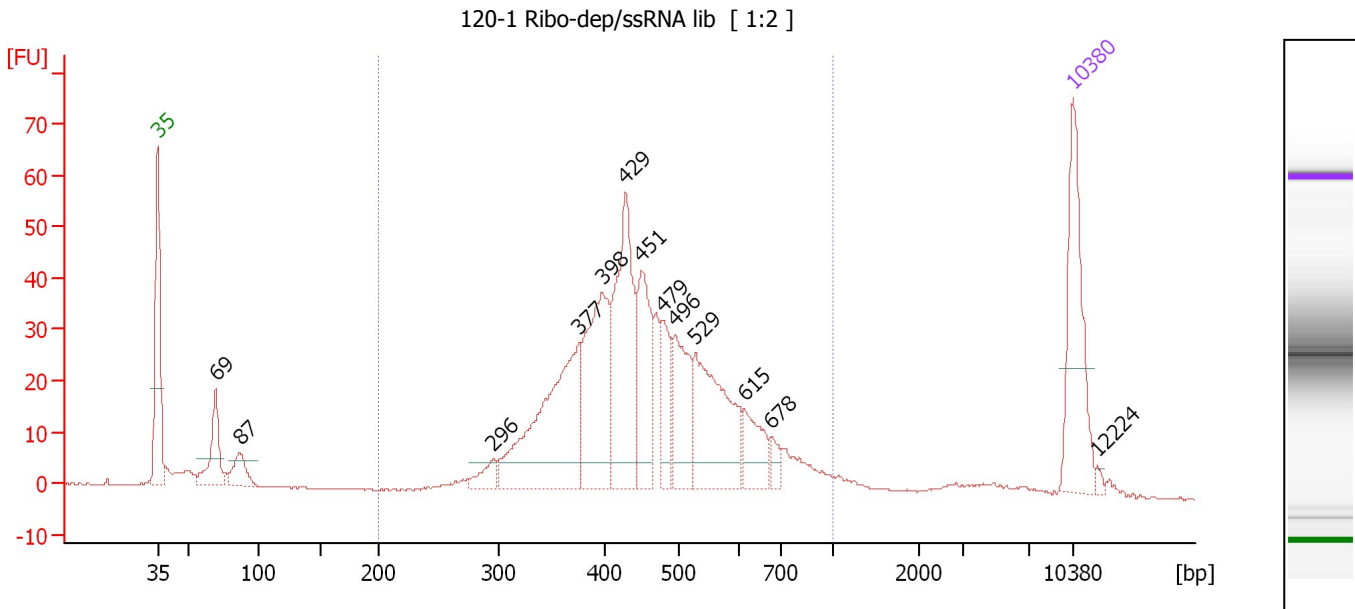
**Region table for sample 6 : 48-3 Ribo-dep/ssRNA lib**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
200	1,000	386	12,772.5	2,988.71	1,028.4	90	28.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : 120-1 Ribo-dep/ssRNA lib**

Number of peaks found: 13                      Corr. Area 1: 709.2  
 Noise: 0.1

**Peak table for sample 7 : 120-1 Ribo-dep/ssRNA lib**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	69	53.27	1,165.4	
3	87	27.52	481.9	
4	296	15.17	77.5	
5	377	168.04	676.1	
6	398	122.35	465.3	
7	429	140.87	497.6	
8	451	73.37	246.7	
9	479	39.72	125.6	
10	496	62.63	191.2	
11	529	109.57	313.9	
12	615	31.88	78.6	
13	678	10.17	22.7	
14	10,380	75.00	10.9	Upper Marker
15	12,224	0.00	0.0	

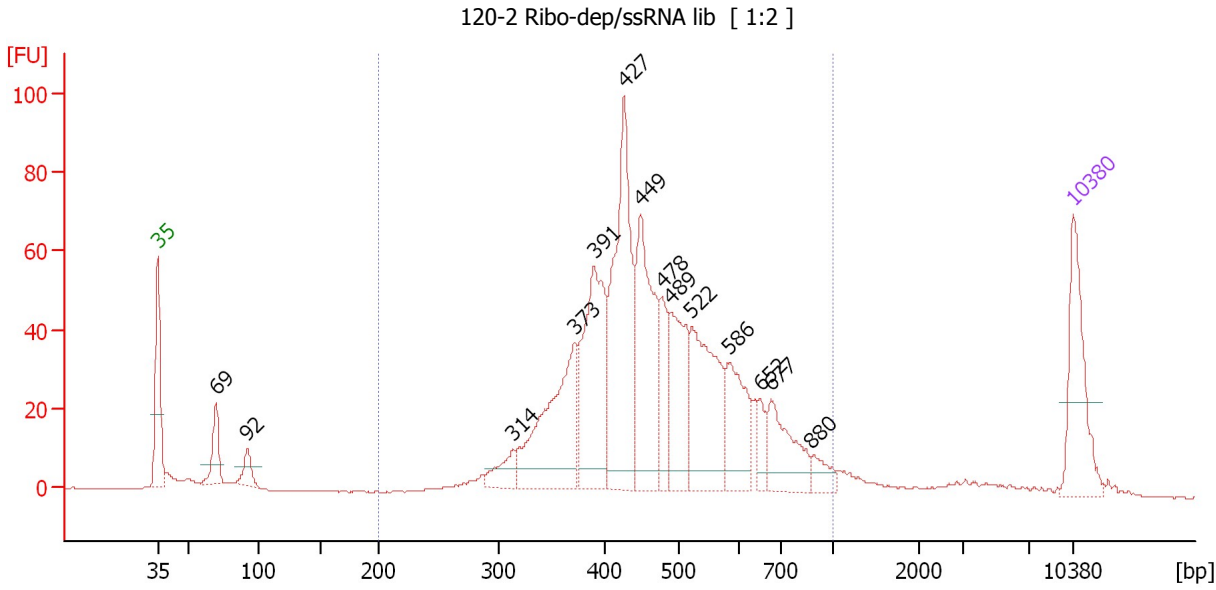
**Region table for sample 7 : 120-1 Ribo-dep/ssRNA lib**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
200	1,000	469	3,068.1	881.90	709.2	87	25.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : 120-2 Ribo-dep/ssRNA lib**

Number of peaks found: 14                      Corr. Area 1: 1,059.8  
 Noise: 0.2

**Peak table for sample 8 : 120-2 Ribo-dep/ssRNA lib**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	69	45.34	991.9	
3	92	24.62	406.3	
4	314	29.97	144.5	
5	373	174.06	708.0	
6	391	182.47	707.3	
7	427	233.17	826.8	
8	449	167.58	565.3	
9	478	51.61	163.6	
10	489	99.89	309.7	
11	522	145.02	421.2	
12	586	78.87	204.1	
13	652	19.05	44.3	
14	677	70.64	158.1	
15	880	17.48	30.1	
16	10,380	75.00	10.9	Upper Marker

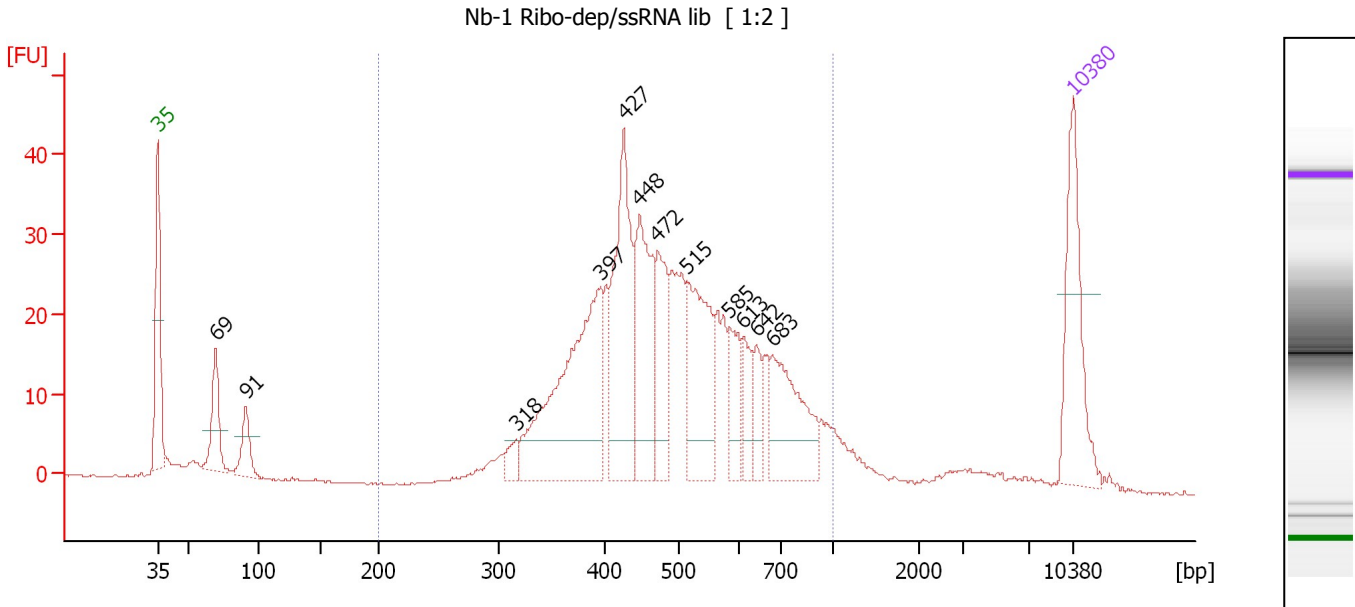
**Region table for sample 8 : 120-2 Ribo-dep/ssRNA lib**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
200	1,000	482	4,592.8	1,353.72	1,059.8	89	25.6

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

Created: 7/14/2014 9:23:23 AM  
 Modified: 7/14/2014 10:04:46 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : Nb-1 Ribo-dep/ssRNA lib**

Number of peaks found: 12                      Corr. Area 1: 584.5  
 Noise: 0.1

**Peak table for sample 9 : Nb-1 Ribo-dep/ssRNA lib**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	69	53.81	1,173.7	
3	91	33.15	553.6	
4	318	13.87	66.1	
5	397	216.48	826.9	
6	427	145.89	517.8	
7	448	113.54	384.0	
8	472	67.32	216.2	
9	515	107.23	315.3	
10	585	35.33	91.5	
11	613	25.34	62.7	
12	642	22.52	53.1	
13	683	86.07	191.0	
14	10,380	75.00	10.9	Upper Marker

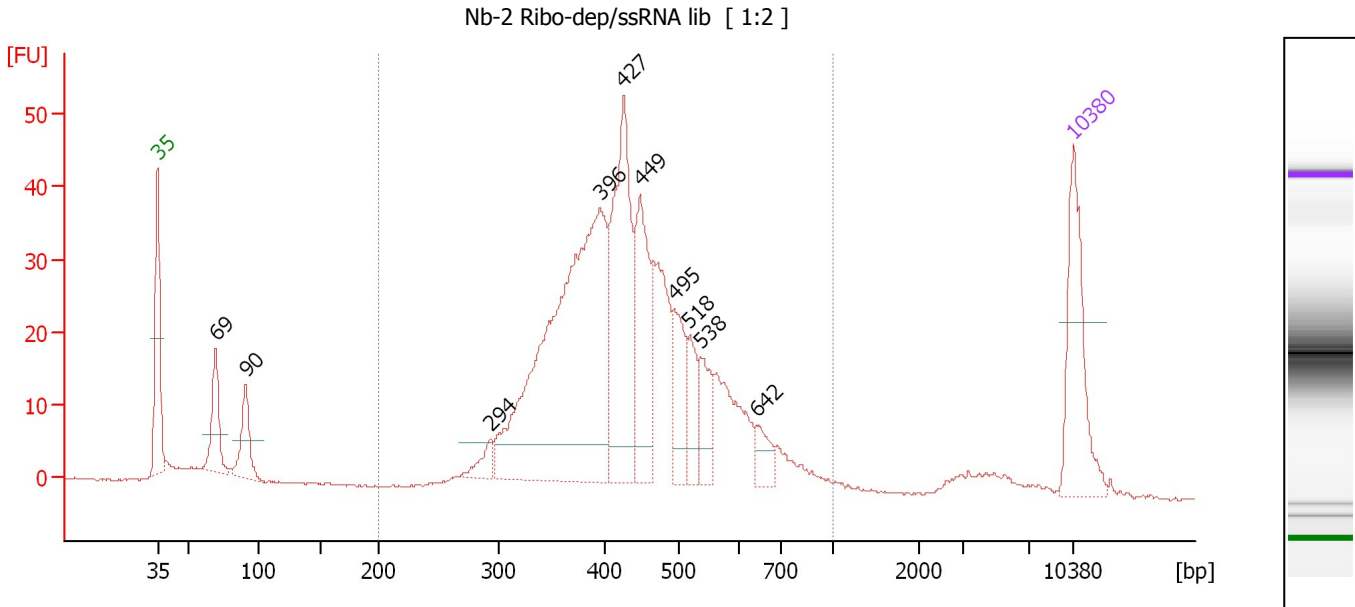
**Region table for sample 9 : Nb-1 Ribo-dep/ssRNA lib**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
200	1,000	508	3,567.5	1,096.23	584.5	86	27.3

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

Created: 7/14/2014 9:23:23 AM  
 Modified: 7/14/2014 10:04:46 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : Nb-2 Ribo-dep/ssRNA lib**

Number of peaks found: 10                      Corr. Area 1: 643.6  
 Noise: 0.1

**Peak table for sample 10 : Nb-2 Ribo-dep/ssRNA lib**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	69	57.64	1,257.5	
3	90	49.74	833.1	
4	294	16.26	83.9	
5	396	447.23	1,710.3	
6	427	189.98	674.8	
7	449	107.36	362.4	
8	495	52.03	159.2	
9	518	34.71	101.5	
10	538	37.70	106.2	
11	642	20.36	48.1	
12	10,380	75.00	10.9	Upper Marker

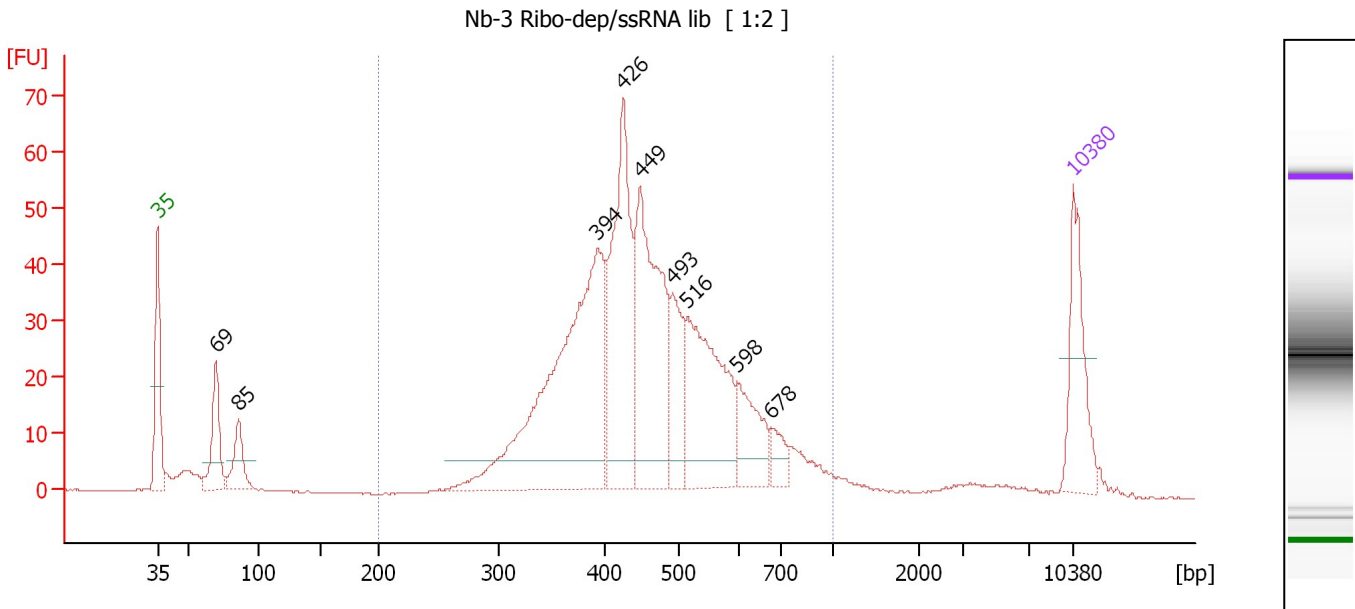
**Region table for sample 10 : Nb-2 Ribo-dep/ssRNA lib**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
200	1,000	446	4,285.4	1,186.86	643.6	89	23.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

Created: 7/14/2014 9:23:23 AM  
 Modified: 7/14/2014 10:04:46 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : Nb-3 Ribo-dep/ssRNA lib**

Number of peaks found: 9                      Corr. Area 1: 805.0  
 Noise: 0.1

**Peak table for sample 11 : Nb-3 Ribo-dep/ssRNA lib**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	69	81.53	1,779.2	
3	85	54.03	957.6	
4	394	450.53	1,733.3	
5	426	244.68	869.7	
6	449	254.62	859.7	
7	493	78.96	242.7	
8	516	203.21	597.2	
9	598	69.05	175.0	
10	678	23.38	52.2	
11	10,380	75.00	10.9	Upper Marker

**Region table for sample 11 : Nb-3 Ribo-dep/ssRNA lib**

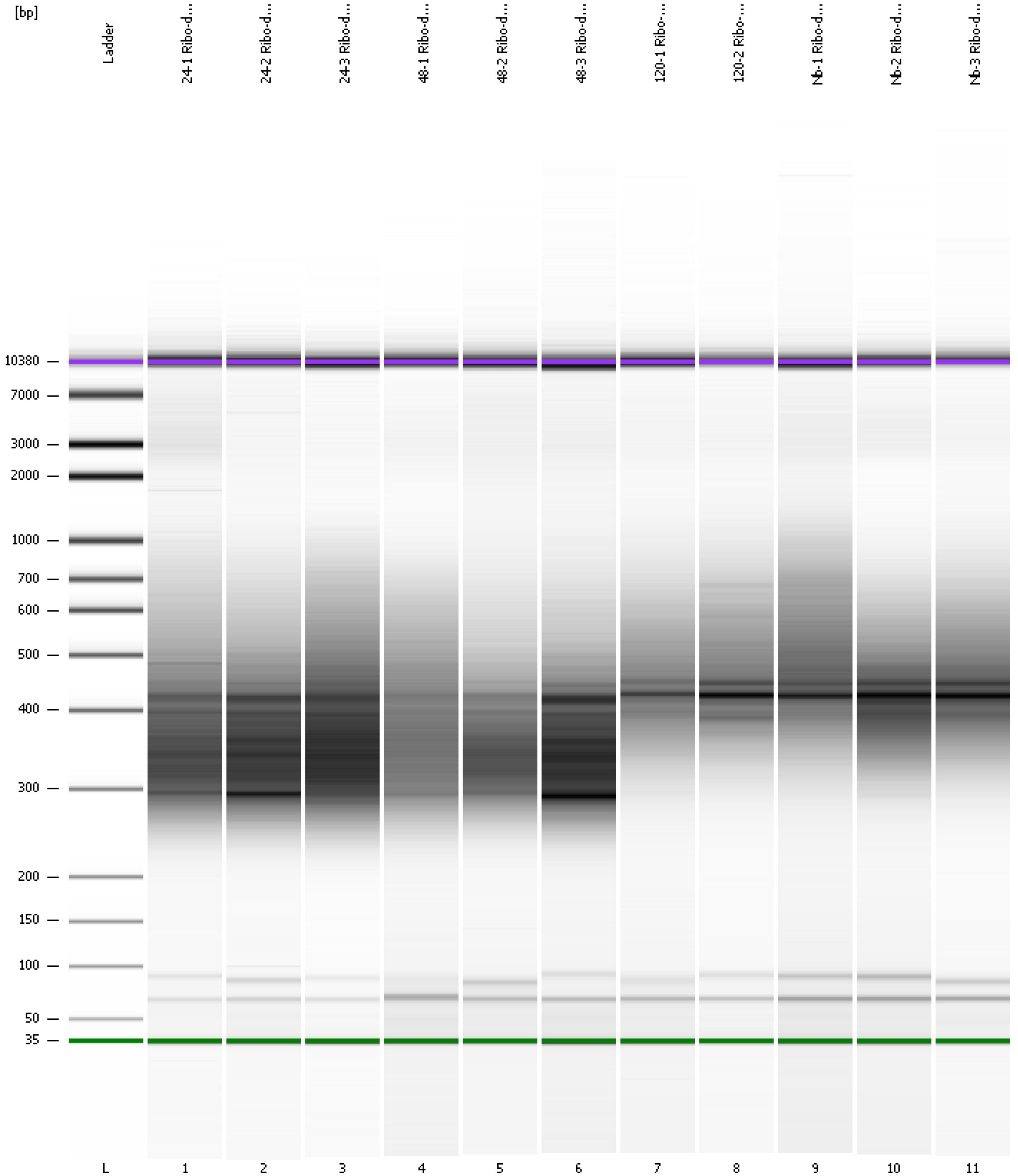
From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
200	1,000	469	5,052.9	1,458.46	805.0	88	24.8



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

Created: 7/14/2014 9:23:23 AM  
Modified: 7/14/2014 10:04:46 AM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14\_001.xad

Created: 7/14/2014 9:23:23 AM  
 Modified: 7/14/2014 10:04:46 AM

**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		7/14/2014 10:04:42 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\Data\2014-07-14\2014-07-14_001.xad)		Instrument	Run		7/14/2014 9:23:28 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		7/14/2014 9:23:28 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		7/14/2014 9:23:28 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		7/14/2014 9:23:28 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		7/14/2014 9:23:28 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		7/14/2014 9:23:28 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		7/14/2014 9:23:28 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1