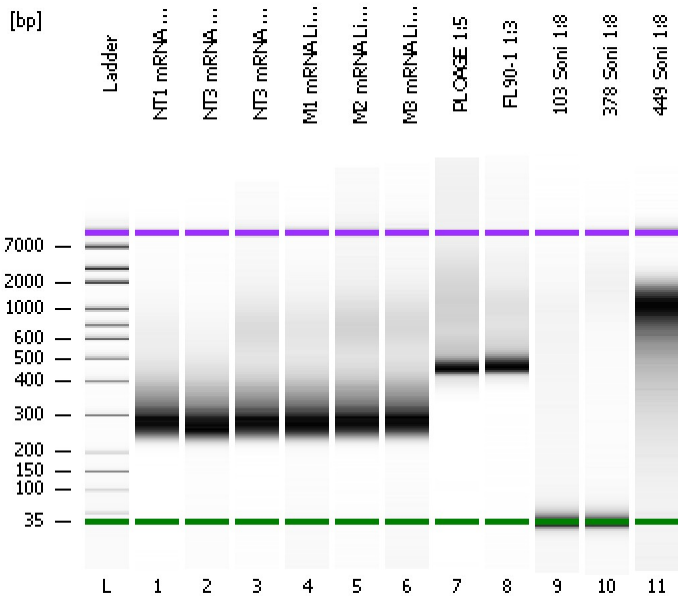


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
Modified: 1/16/2013 5:34:05 PM

Electrophoresis File Run Summary



Instrument Information:

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

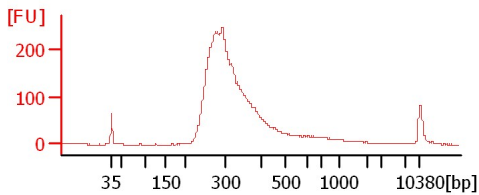
Assay Information:

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

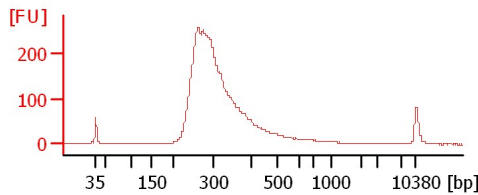
Chip Information:

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

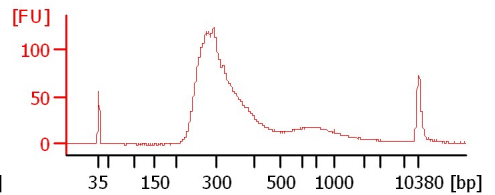
NT1 mRNA Lib 1:4



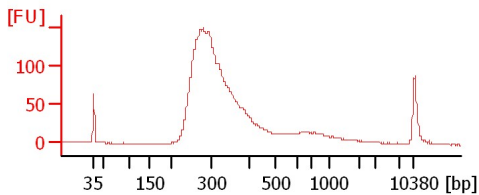
NT3 mRNA Lib 1:2



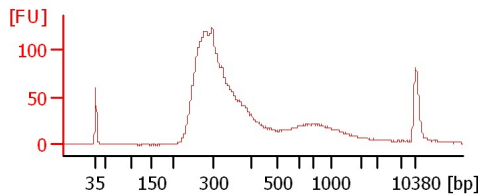
NT3 mRNA Lib 1:6



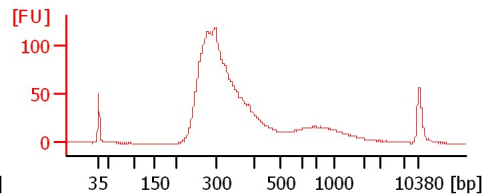
M1 mRNA Lib 1:7



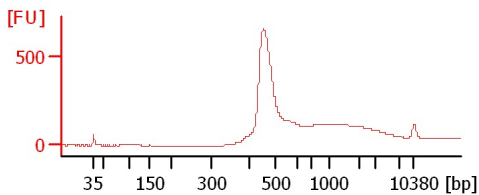
M2 mRNA Lib 1:8



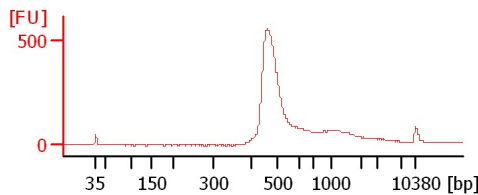
M3 mRNA Lib 1:8



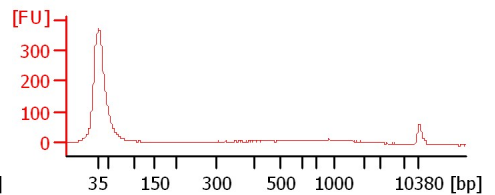
PLOAGE 1:5



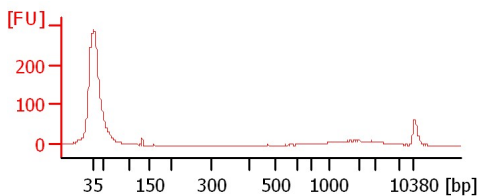
FL90-1 1:3



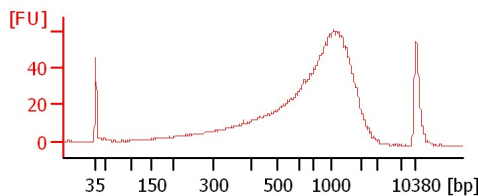
103 Soni 1:8



378 Soni 1:8



449 Soni 1:8



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
NT1 mRNA Lib 1:4		<input type="checkbox"/>	✓			
NT3 mRNA Lib 1:2		<input type="checkbox"/>	✓			
NT3 mRNA Lib 1:6		<input type="checkbox"/>	✓			
M1 mRNA Lib 1:7		<input type="checkbox"/>	✓			
M2 mRNA Lib 1:8		<input type="checkbox"/>	✓			
M3 mRNA Lib 1:8		<input type="checkbox"/>	✓			
PLOAGE 1:5		<input type="checkbox"/>	✓			
FL90-1 1:3		<input type="checkbox"/>	✓			
103 Soni 1:8		<input type="checkbox"/>	✓			
378 Soni 1:8		<input type="checkbox"/>	✓			
449 Soni 1:8		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
Modified: 1/16/2013 5:34:05 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

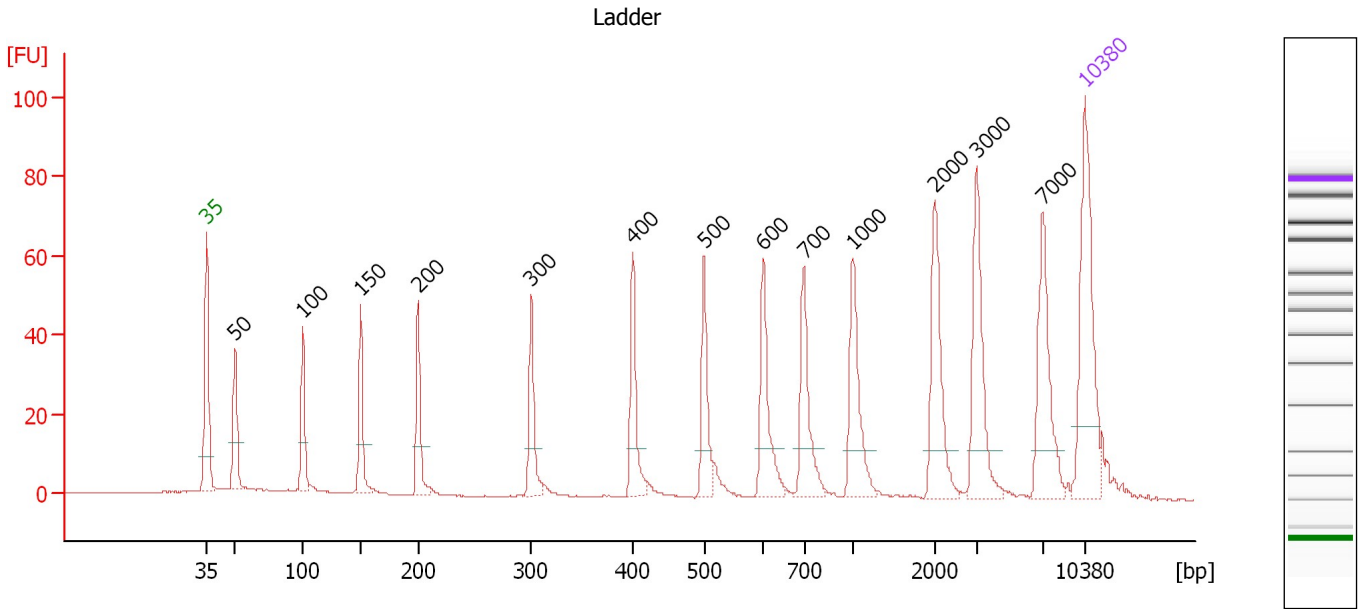
Ladder

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:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

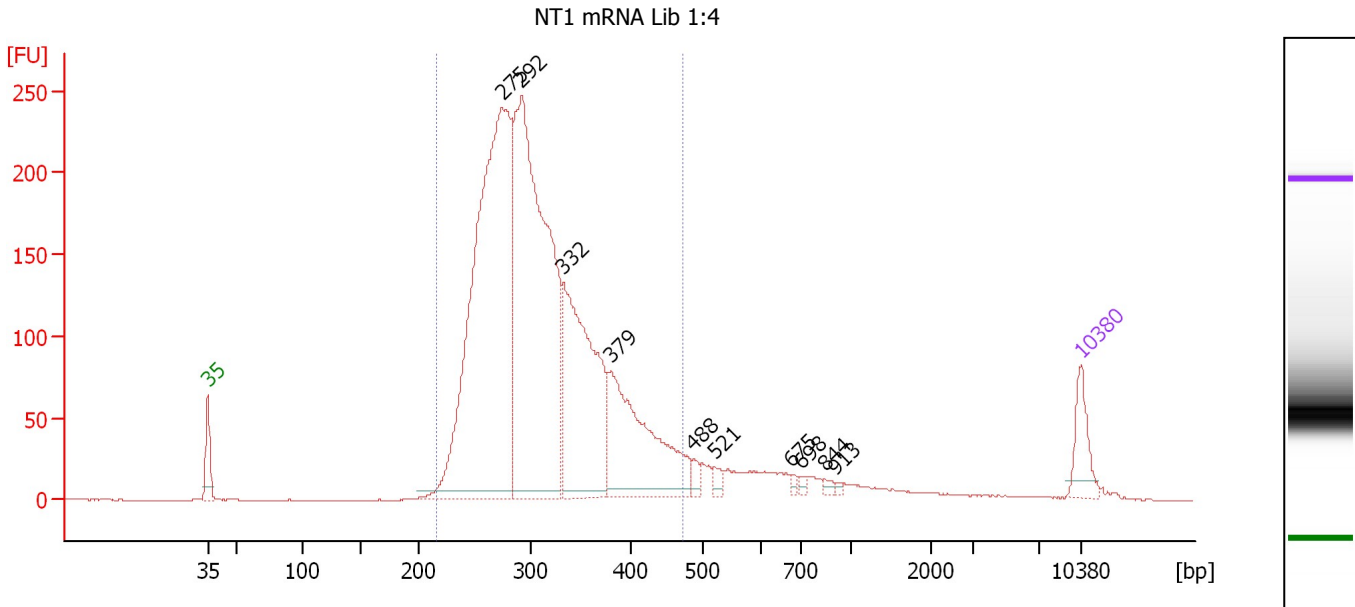
Peak table for Ladder

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:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : NT1 mRNA Lib 1:4

Number of peaks found: 10 Corr. Area 1: 3,458.6
 Noise: 0.2

Peak table for sample 1 : NT1 mRNA Lib 1:4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	275	1,928.42	10,636.3	
3	292	1,646.15	8,547.8	
4	332	760.06	3,467.2	
5	379	533.74	2,132.2	
6	488	25.25	78.5	
7	521	23.46	68.3	
8	675	9.62	21.6	
9	698	10.96	23.8	
10	844	11.55	20.7	
11	913	6.65	11.0	
12	10,380	75.00	10.9	Upper Marker

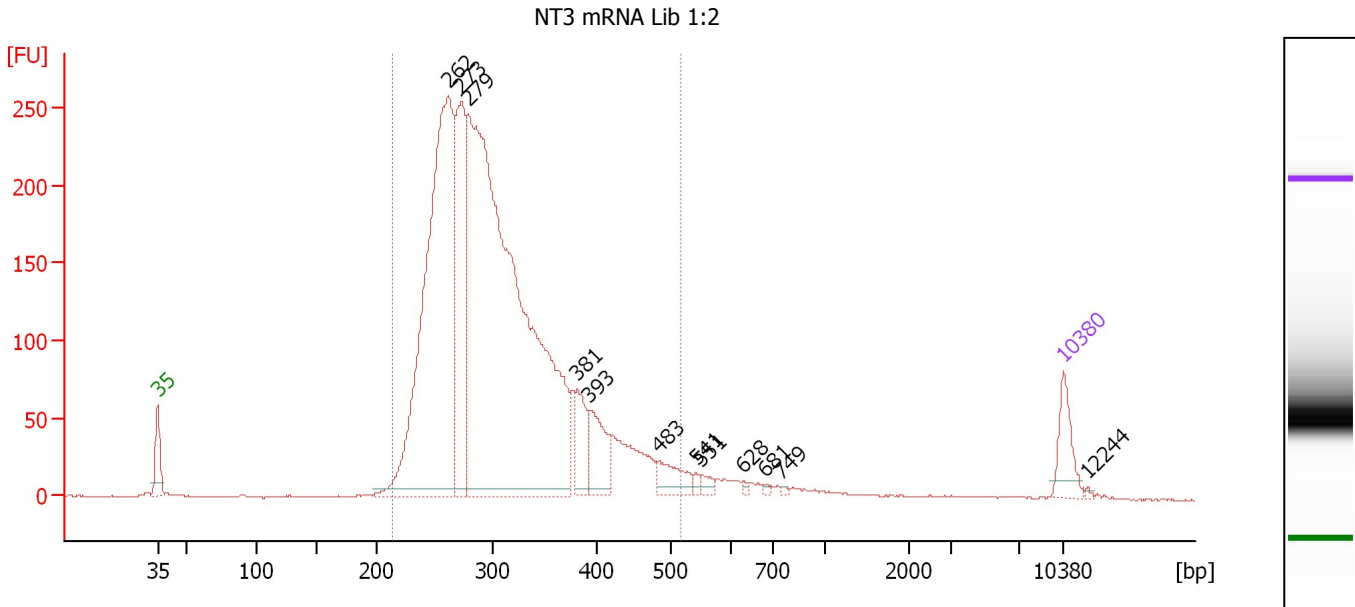
Region table for sample 1 : NT1 mRNA Lib 1:4

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
216	472	311	24,391.9	4,833.81	3,458.6	90	16.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : NT3 mRNA Lib 1:2

Number of peaks found: 12 Corr. Area 1: 3,654.9
 Noise: 0.3

Peak table for sample 2 : NT3 mRNA Lib 1:2

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	262	1,629.06	9,420.3	
3	273	521.17	2,890.9	
4	279	2,771.91	15,055.5	
5	381	120.06	477.1	
6	393	143.35	553.2	
7	483	78.97	247.7	
8	541	13.45	37.7	
9	551	18.73	51.5	
10	628	6.33	15.3	
11	681	5.41	12.0	
12	749	4.18	8.5	
13	10,380	75.00	10.9	Upper Marker
14	12,244	0.00	0.0	

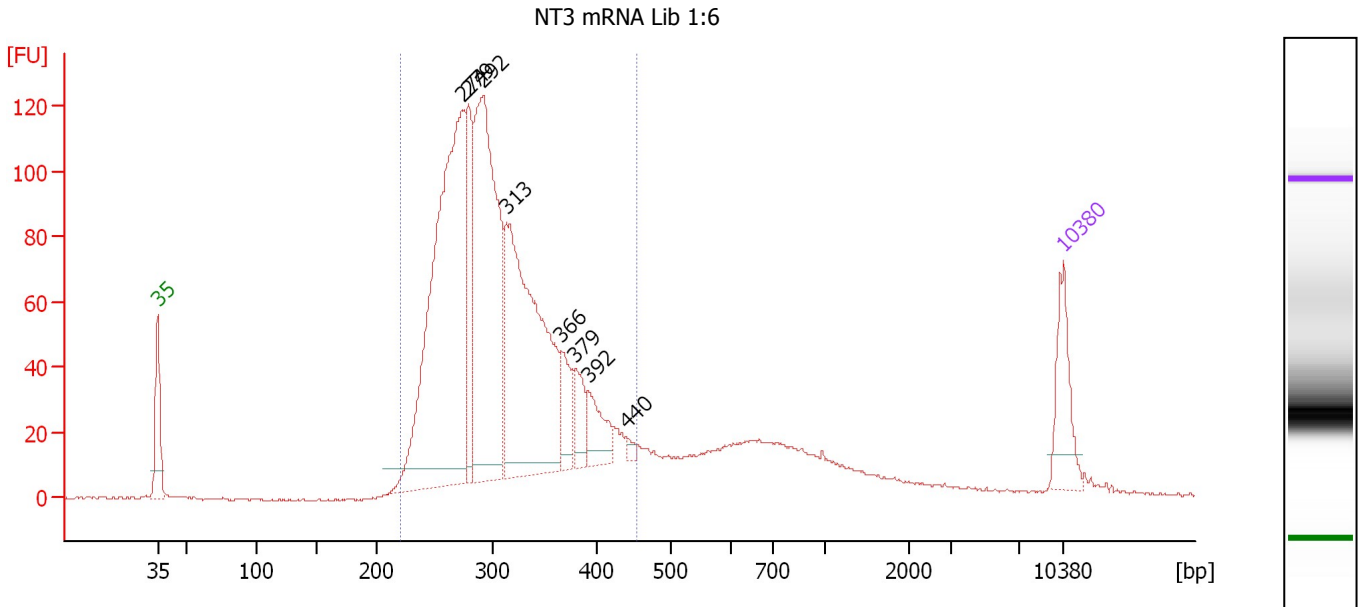
Region table for sample 2 : NT3 mRNA Lib 1:2

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/µl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
214	516	308	27,337.3	5,327.86	3,654.9	94	19.1	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : NT3 mRNA Lib 1:6

Number of peaks found: 8 Corr. Area 1: 1,653.6
 Noise: 0.4

Peak table for sample 3 : NT3 mRNA Lib 1:6

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	274	857.89	4,735.6	
3	279	170.39	924.0	
4	292	602.85	3,126.4	
5	313	578.12	2,797.0	
6	366	69.17	286.5	
7	379	55.35	221.3	
8	392	67.38	260.7	
9	440	9.52	32.8	
10	10,380	75.00	10.9	Upper Marker

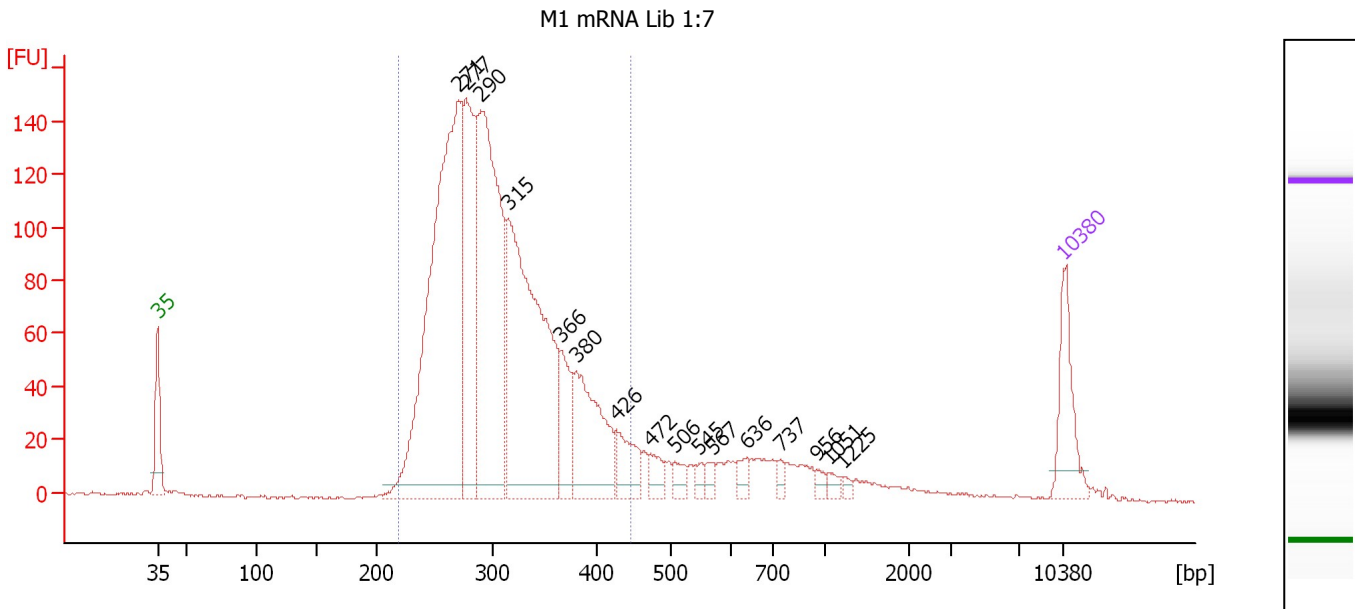
Region table for sample 3 : NT3 mRNA Lib 1:6

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
220	453	308	13,560.0	2,673.81	1,653.6	84	16.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : M1 mRNA Lib 1:7

Number of peaks found: 16 Corr. Area 1: 2,071.2
 Noise: 0.4

Peak table for sample 4 : M1 mRNA Lib 1:7

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	271	867.86	4,851.7	
3	277	306.56	1,676.5	
4	290	605.15	3,160.4	
5	315	612.96	2,950.5	
6	366	93.77	388.4	
7	380	189.09	754.3	
8	426	64.64	229.8	
9	472	28.68	92.1	
10	506	21.37	64.0	
11	545	14.08	39.1	
12	567	12.46	33.3	
13	636	18.73	44.6	
14	737	10.83	22.3	
15	956	8.82	14.0	
16	1,051	10.36	14.9	
17	1,225	5.45	6.7	
18	10,380	75.00	10.9	Upper Marker

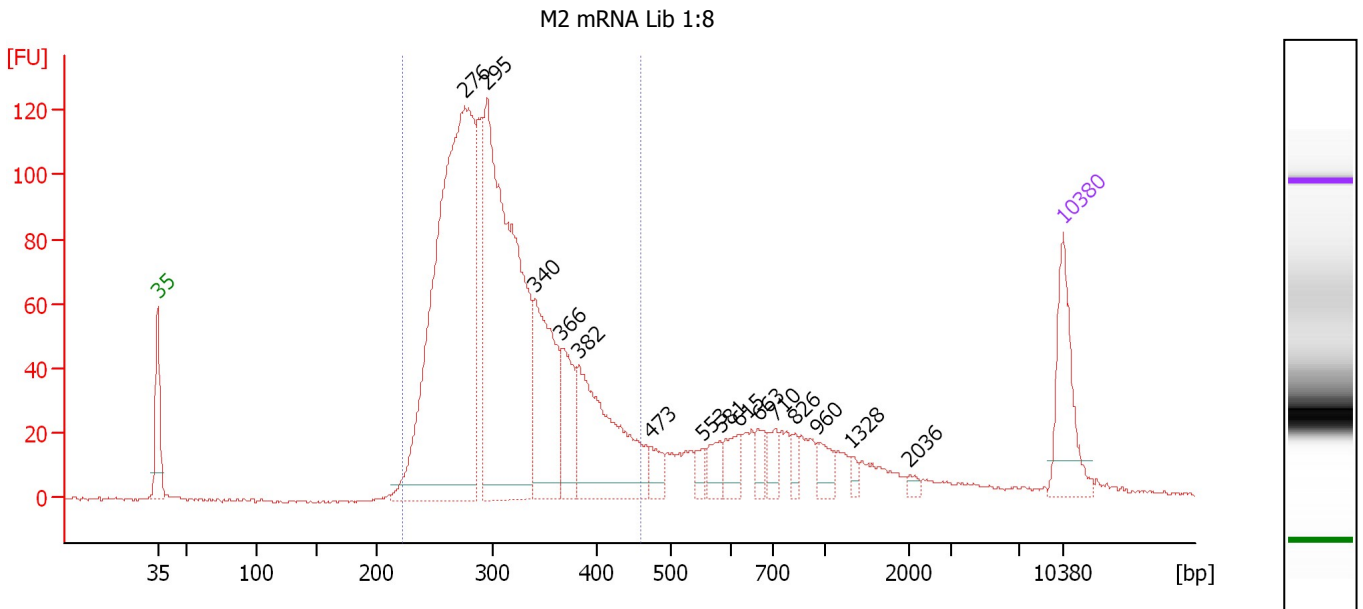
Region table for sample 4 : M1 mRNA Lib 1:7

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
219	446	307	13,698.5	2,689.55	2,071.2	86	15.7	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : M2 mRNA Lib 1:8

Number of peaks found: 15 Corr. Area 1: 1,676.9
 Noise: 0.5

Peak table for sample 5 : M2 mRNA Lib 1:8

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	276	846.24	4,643.8	
3	295	642.12	3,293.2	
4	340	193.41	861.2	
5	366	82.02	339.7	
6	382	218.70	866.9	
7	473	23.55	75.4	
8	553	15.41	42.2	
9	581	24.65	64.3	
10	615	28.77	70.9	
11	663	21.00	48.0	
12	710	20.86	44.5	
13	826	12.64	23.2	
14	960	21.19	33.5	
15	1,328	7.18	8.2	
16	2,036	4.86	3.6	
17	10,380	75.00	10.9	Upper Marker

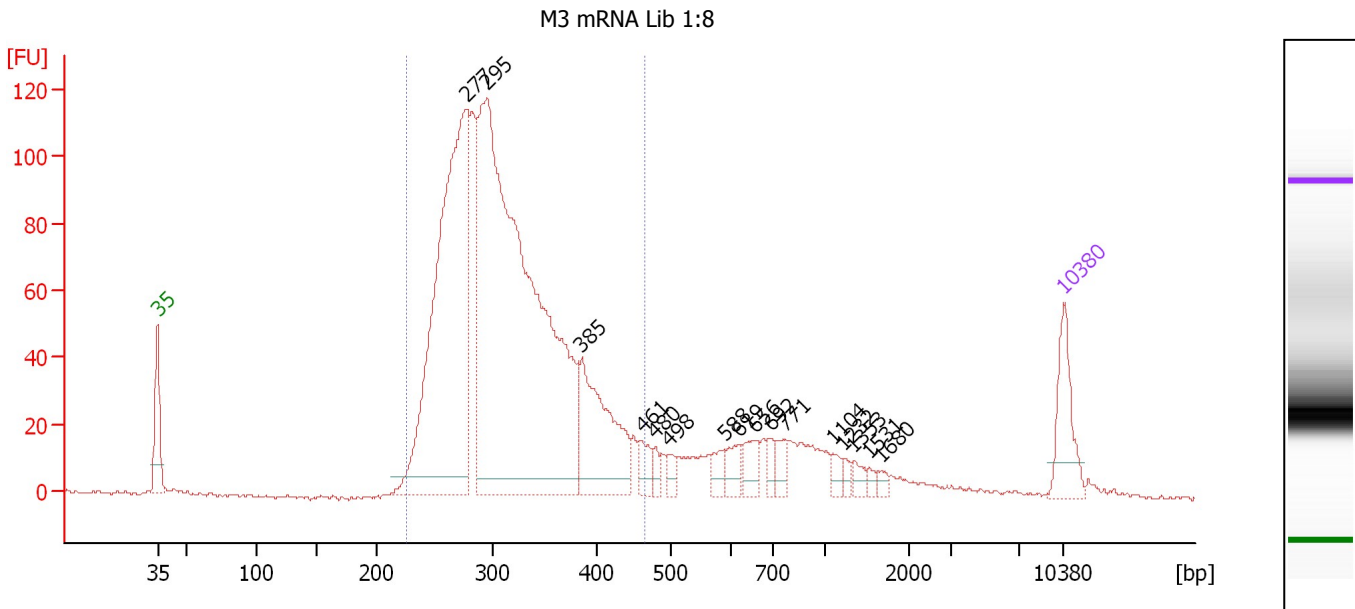
Region table for sample 5 : M2 mRNA Lib 1:8

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
223	458	311	10,160.2	2,017.97	1,676.9	80	16.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : M3 mRNA Lib 1:8

Number of peaks found: 16 Corr. Area 1: 1,611.6
 Noise: 0.5

Peak table for sample 6 : M3 mRNA Lib 1:8

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	277	951.03	5,198.7	
3	295	1,566.58	8,056.1	
4	385	247.86	975.2	
5	461	34.04	111.9	
6	480	16.75	52.9	
7	498	17.99	54.7	
8	588	26.16	67.4	
9	619	30.95	75.8	
10	656	38.12	88.1	
11	692	17.97	39.4	
12	771	26.19	51.4	
13	1,104	14.86	20.4	
14	1,232	10.72	13.2	
15	1,353	14.54	16.3	
16	1,531	8.09	8.0	
17	1,680	8.89	8.0	
18	10,380	75.00	10.9	Upper Marker

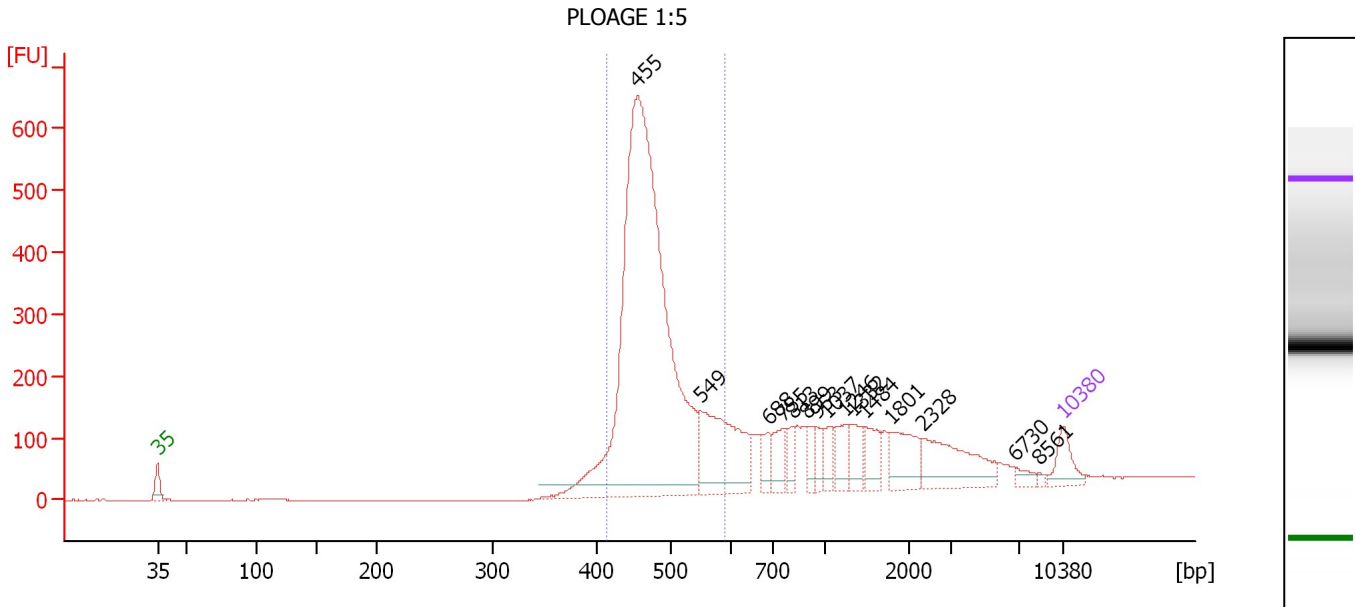
Region table for sample 6 : M3 mRNA Lib 1:8

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
226	466	314	14,535.7	2,914.75	1,611.6	82	16.3	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 7 : PLOAGE 1:5

Height Threshold [FU] : 20

Overall Results for sample 7 : PLOAGE 1:5

Number of peaks found: 15 Corr. Area 1: 3,220.2
 Noise: 0.6

Peak table for sample 7 : PLOAGE 1:5

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	455	2,829.09	9,430.6	
3	549	480.32	1,326.7	
4	688	72.46	159.7	
5	755	98.90	198.4	
6	813	56.95	106.1	
7	899	62.67	105.6	
8	953	55.55	88.3	
9	1,037	61.62	90.1	
10	1,246	94.66	115.1	
11	1,332	81.78	93.0	
12	1,484	91.23	93.2	
13	1,801	149.01	125.3	
14	2,328	241.73	157.3	
15	6,730	28.06	6.3	
16	8,561	7.77	1.4	
17	10,380	75.00	10.9	Upper Marker

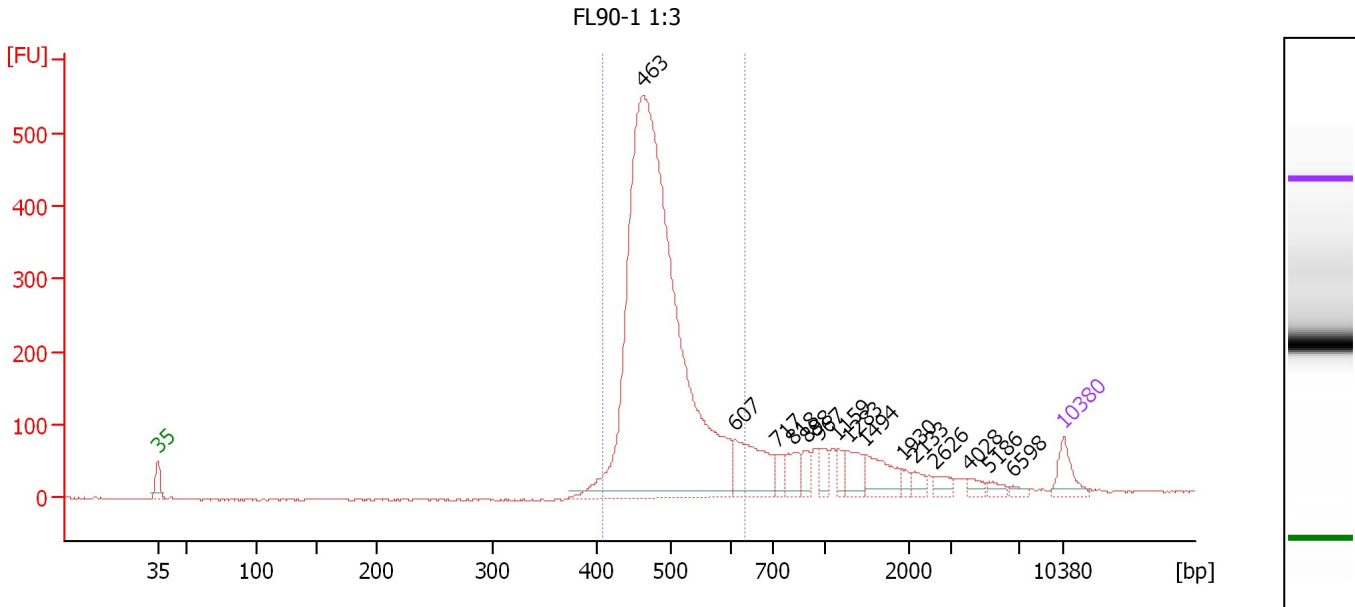
Region table for sample 7 : PLOAGE 1:5

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/µl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
412	590	479	9,020.8	2,832.11	3,220.2	59	8.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 8 : FL90-1 1:3

Height Threshold [FU] : 10

Overall Results for sample 8 : FL90-1 1:3

Number of peaks found: 15 Corr. Area 1: 3,076.2
 Noise: 1.2

Peak table for sample 8 : FL90-1 1:3

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	463	3,574.16	11,693.0	
3	607	271.95	678.7	
4	717	50.55	106.9	
5	818	84.49	156.5	
6	898	51.75	87.3	
7	967	58.71	92.0	
8	1,159	45.86	59.9	
9	1,283	84.56	99.9	
10	1,494	130.64	132.5	
11	1,930	20.54	16.1	
12	2,133	34.93	24.8	
13	2,626	37.74	21.8	
14	4,028	26.60	10.0	
15	5,186	23.17	6.8	
16	6,598	14.88	3.4	
17	10,380	75.00	10.9	Upper Marker

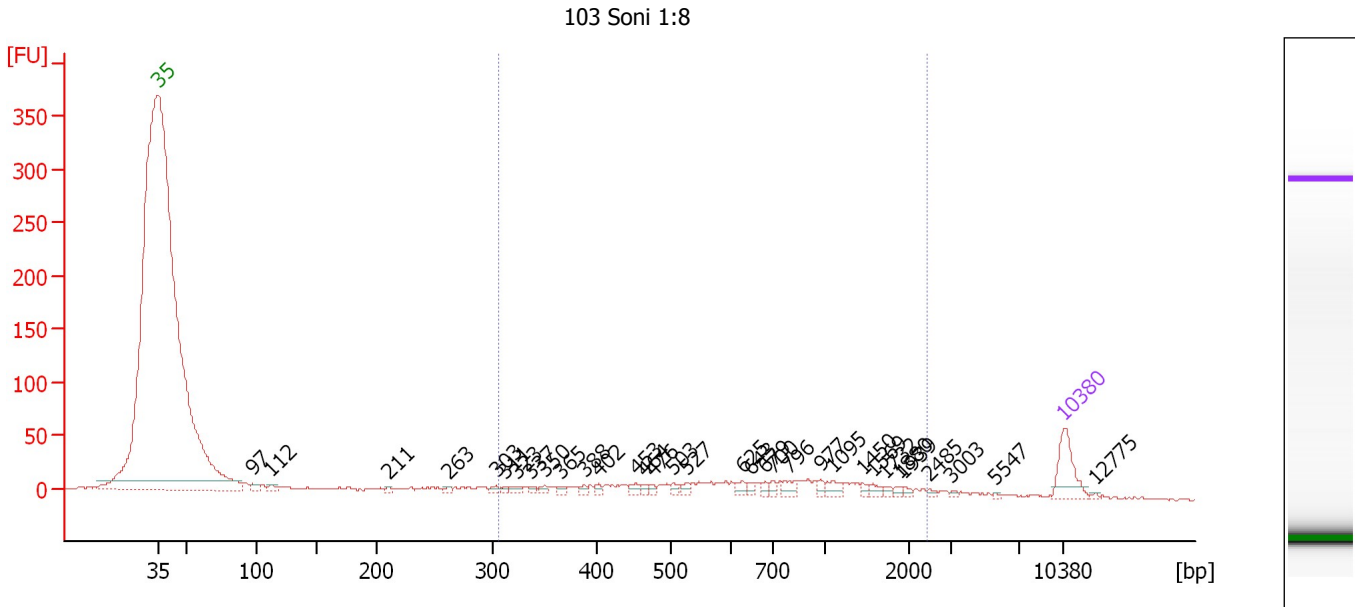
Region table for sample 8 : FL90-1 1:3

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/µl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
407	630	488	10,958.2	3,502.17	3,076.2	74	9.1	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : 103 Soni 1:8

Number of peaks found: 33 Corr. Area 1: 374.7
 Noise: 0.7

Peak table for sample 9 : 103 Soni 1:8

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	97	18.29	285.7	
3	112	15.52	209.8	
4	211	7.57	54.4	
5	263	8.18	47.1	
6	303	11.04	55.1	
7	311	7.46	36.3	
8	323	14.76	69.2	
9	337	7.60	34.1	
10	350	10.38	44.9	
11	365	10.83	45.0	
12	388	8.53	33.3	
13	402	10.65	40.2	
14	453	14.37	48.1	
15	464	9.24	30.1	
16	476	9.20	29.3	
17	503	10.85	32.7	
18	527	12.63	36.3	
19	625	15.39	37.3	
20	643	10.64	25.1	
21	679	11.01	24.6	
22	700	11.57	25.1	
23	796	22.25	42.4	
24	977	10.91	16.9	
25	1,095	22.47	31.1	
26	1,450	8.77	9.2	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad


Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary Continued ...

... Peak table for sample 9 : 103 Soni 1:8

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	1,569	11.46	11.1	
28	1,732	7.41	6.5	
29	1,880	5.92	4.8	
30	1,939	6.70	5.2	
31	2,485	5.07	3.1	
32	3,003	4.36	2.2	
33	5,547	2.47	0.7	
34	10,380	75.00	10.9	Upper Marker
35	12,775	0.00	0.0	

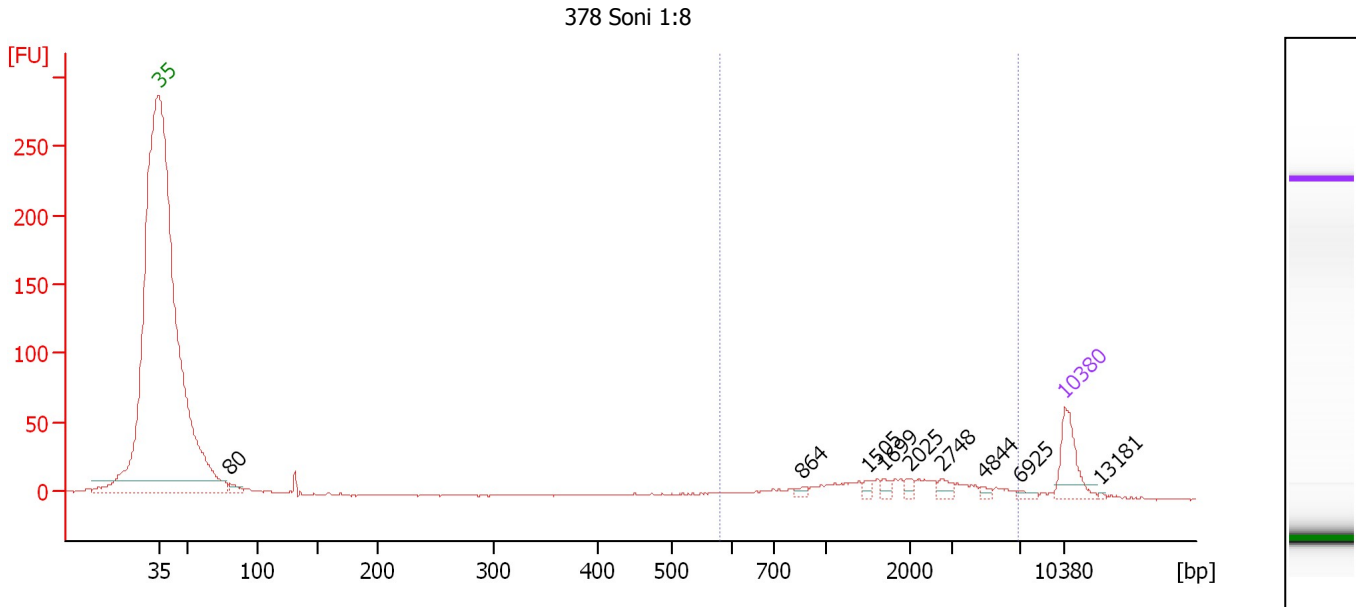
Region table for sample 9 : 103 Soni 1:8

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
307	2,442	814	1,240.3	467.05	374.7	65	59.5	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : 378 Soni 1:8

Number of peaks found: 9 Corr. Area 1: 194.8
 Noise: 0.6

Peak table for sample 10 : 378 Soni 1:8

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	80	21.95	413.4	
3	864	9.29	16.3	
4	1,505	10.17	10.2	
5	1,699	10.43	9.3	
6	2,025	10.40	7.8	
7	2,748	15.36	8.5	
8	4,844	6.78	2.1	
9	6,925	7.73	1.7	
10	10,380	75.00	10.9	Upper Marker
11	13,181	0.00	0.0	

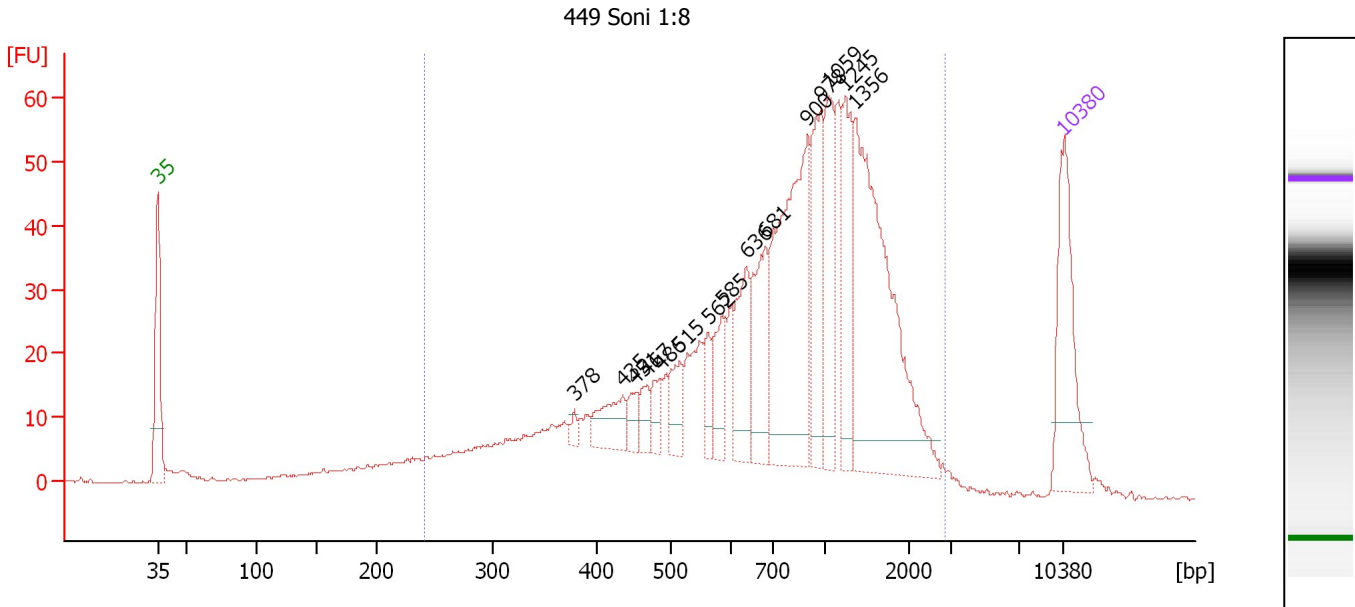
Region table for sample 10 : 378 Soni 1:8

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
579	6,897	2,236	210.2	202.95	194.8	73	65.7	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : 449 Soni 1:8

Number of peaks found: 15 Corr. Area 1: 1,073.7
 Noise: 0.2

Peak table for sample 11 : 449 Soni 1:8

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	378	6.07	24.3	
3	435	36.25	126.2	
4	451	15.50	52.0	
5	467	16.36	53.0	
6	486	18.03	56.2	
7	515	28.73	84.6	
8	562	19.93	53.8	
9	585	34.32	88.9	
10	636	58.66	139.7	
11	681	69.82	155.4	
12	900	184.03	309.7	
13	978	74.73	115.8	
14	1,059	66.01	94.4	
15	1,245	66.33	80.8	
16	1,356	221.85	248.0	
17	10,380	75.00	10.9	Upper Marker

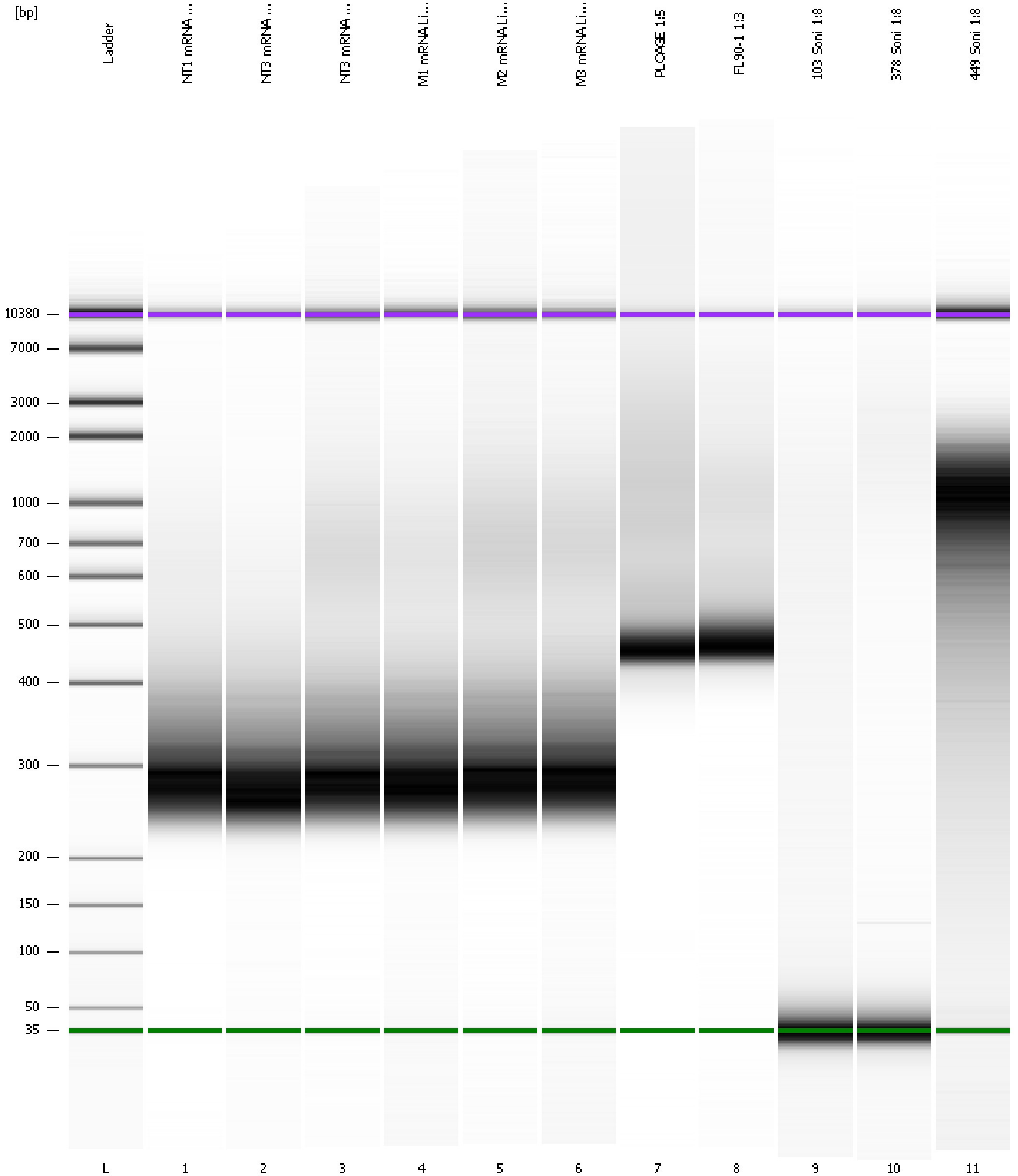
Region table for sample 11 : 449 Soni 1:8

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
242	2,864	929	3,556.2	1,474.71	1,073.7	92	53.7	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
Modified: 1/16/2013 5:34:05 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad

Created: 1/16/2013 4:33:07 PM
 Modified: 1/16/2013 5:34:05 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		1/16/2013 5:14:21 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2013-01-16\2013-01-16_002.xad)		Instrument	Run		1/16/2013 4:33:08 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		1/16/2013 4:33:08 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		1/16/2013 4:33:08 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		1/16/2013 4:33:08 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		1/16/2013 4:33:08 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		1/16/2013 4:33:08 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		1/16/2013 4:33:08 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1