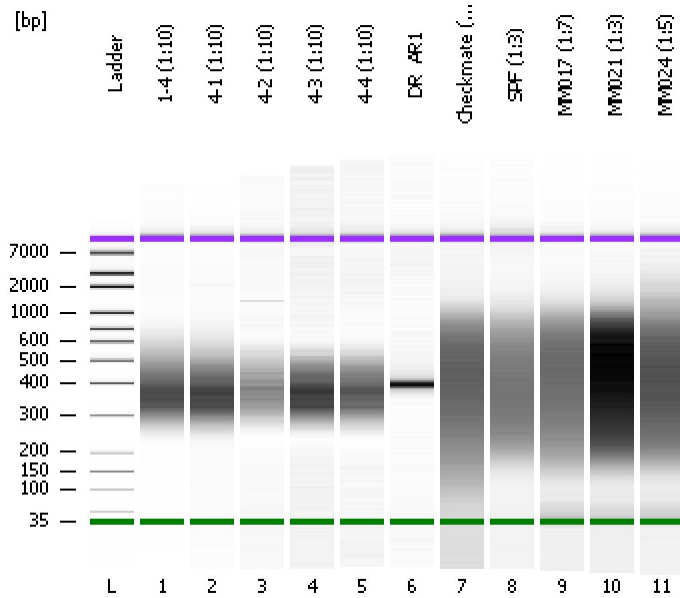


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
Modified: 10/6/2014 4:37:36 PM

Electrophoresis File Run Summary



Instrument Information:

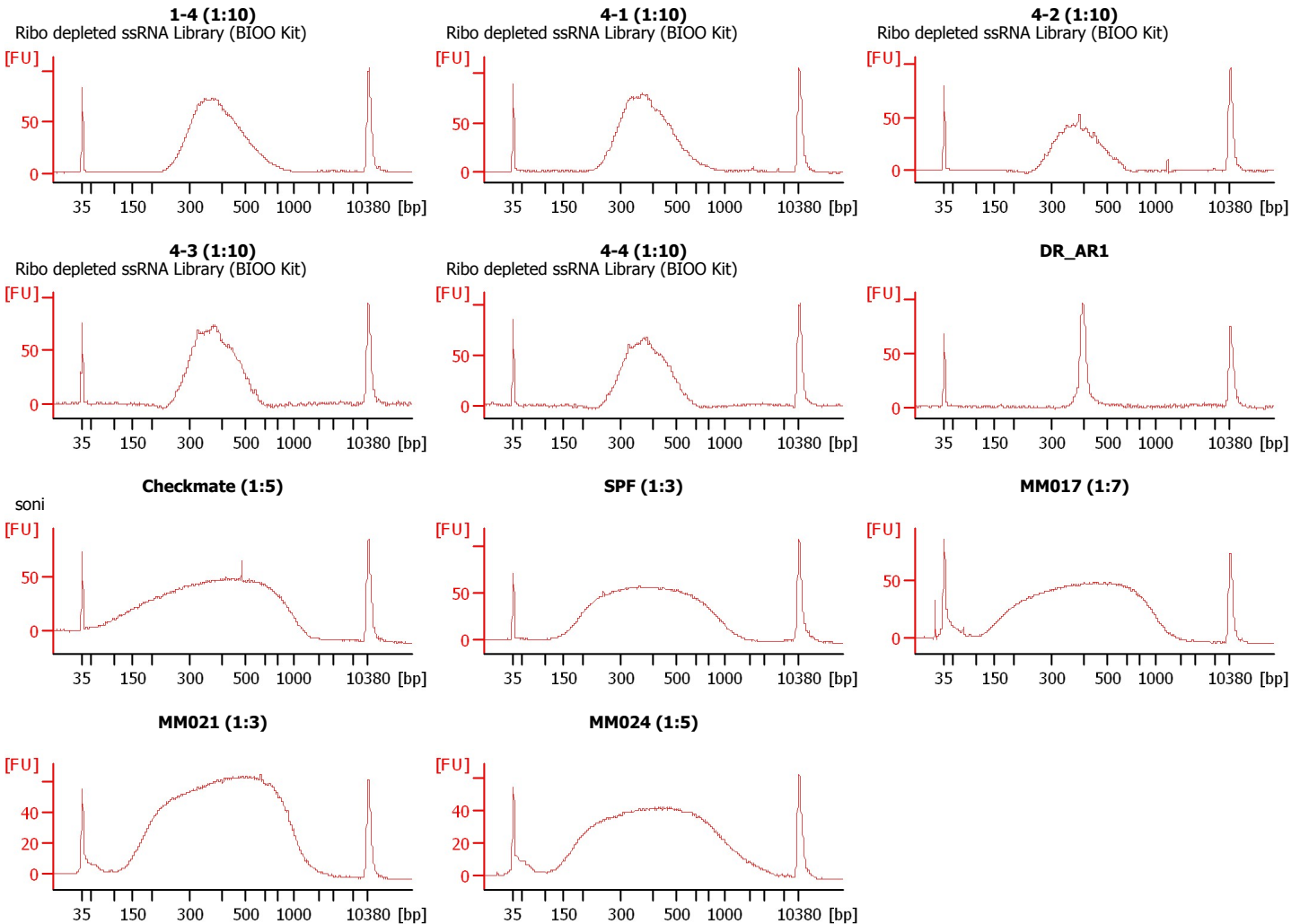
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
Modified: 10/6/2014 4:37:36 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
1-4 (1:10)	Ribo depleted ssRNA Library (BIOO Kit)	<input type="checkbox"/>	✓			
4-1 (1:10)	Ribo depleted ssRNA Library (BIOO Kit)	<input type="checkbox"/>	✓			
4-2 (1:10)	Ribo depleted ssRNA Library (BIOO Kit)	<input type="checkbox"/>	✓			
4-3 (1:10)	Ribo depleted ssRNA Library (BIOO Kit)	<input type="checkbox"/>	✓			
4-4 (1:10)	Ribo depleted ssRNA Library (BIOO Kit)	<input type="checkbox"/>	✓			
DR_AR1		<input type="checkbox"/>	✓			
Checkmate (1:5)	soni	<input type="checkbox"/>	✓			
SPF (1:3)		<input type="checkbox"/>	✓			
MM017 (1:7)		<input type="checkbox"/>	✓			
MM021 (1:3)		<input type="checkbox"/>	✓			
MM024 (1:5)		<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\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
Modified: 10/6/2014 4:37:36 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

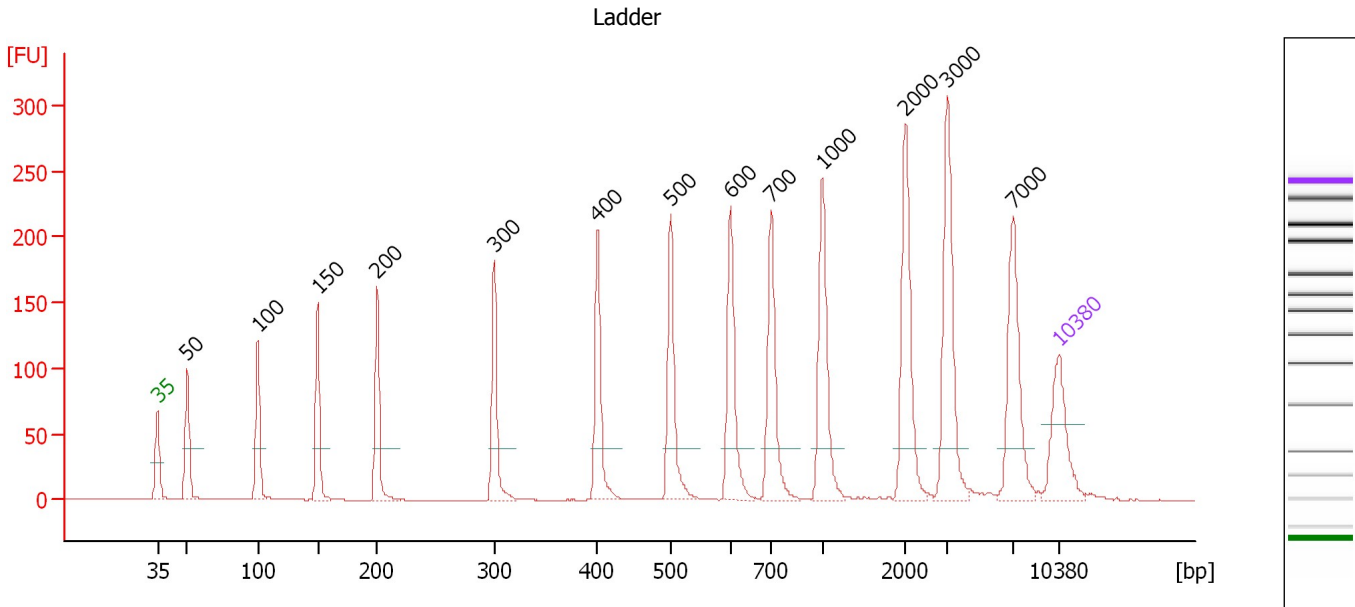
Ladder

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\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

Peak table for Ladder

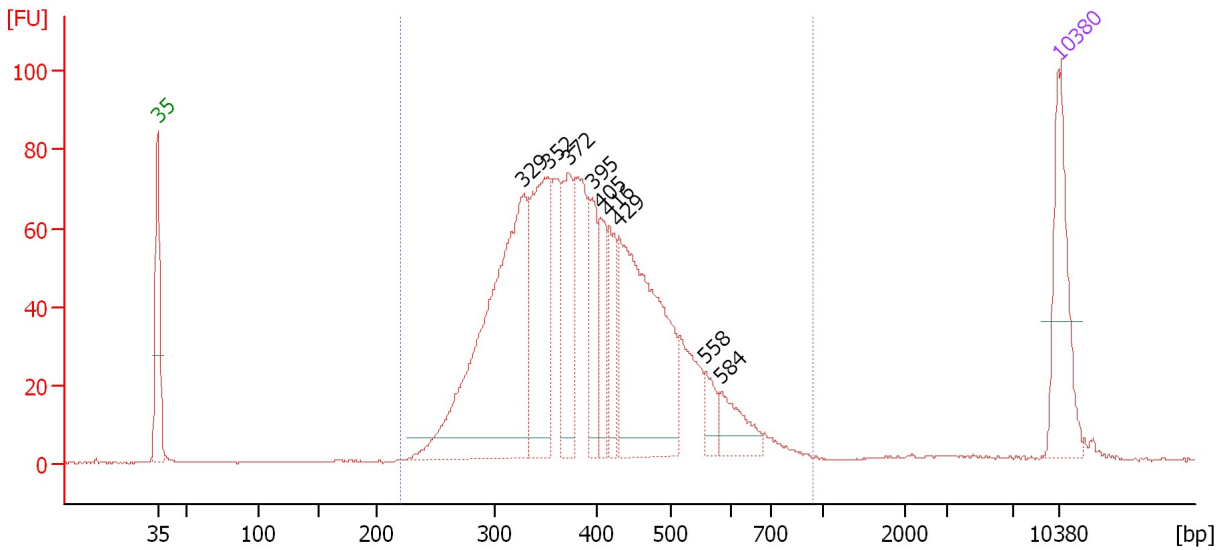
Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	50	150.00	4,545.5	Ladder Peak
3	100	150.00	2,272.7	Ladder Peak
4	150	150.00	1,515.2	Ladder Peak
5	200	150.00	1,136.4	Ladder Peak
6	300	150.00	757.6	Ladder Peak
7	400	150.00	568.2	Ladder Peak
8	500	150.00	454.5	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:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...

1-4 (1:10) [Ribo depleted ssRNA Library (BIOO Kit)]



Overall Results for sample 1 : 1-4 (1:10)

Number of peaks found: 9 Corr. Area 1: 1,482.0
 Noise: 0.1

Peak table for sample 1 : 1-4 (1:10)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	329	381.82	1,760.0	
3	352	169.53	729.4	
4	372	93.73	381.5	
5	395	58.13	222.9	
6	405	50.54	189.1	
7	416	47.38	172.6	
8	429	250.54	884.1	
9	558	22.51	61.1	
10	584	36.85	95.6	
11	10,380	75.00	10.9	Upper Marker

Region table for sample 1 : 1-4 (1:10)

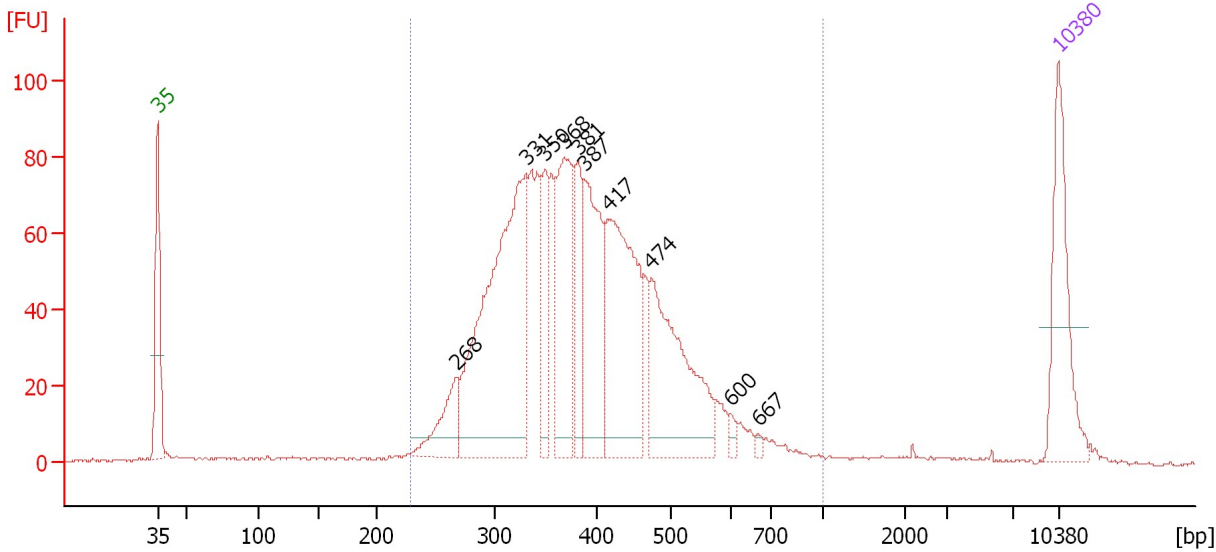
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
220	401	942	1,482.0	97	23.8	1,442.44	5,838.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...

4-1 (1:10) [Ribo depleted ssRNA Library (BIOO Kit)]



Overall Results for sample 2 : 4-1 (1:10)

Number of peaks found: 10 Corr. Area 1: 1,587.5
 Noise: 0.3

Peak table for sample 2 : 4-1 (1:10)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	268	52.60	297.2	
3	331	342.47	1,567.7	
4	350	60.78	263.4	
5	368	133.19	549.1	
6	381	61.65	245.1	
7	387	125.96	492.8	
8	417	192.58	700.0	
9	474	161.27	515.7	
10	600	6.44	16.3	
11	667	3.33	7.6	
12	10,380	75.00	10.9	Upper Marker

Region table for sample 2 : 4-1 (1:10)

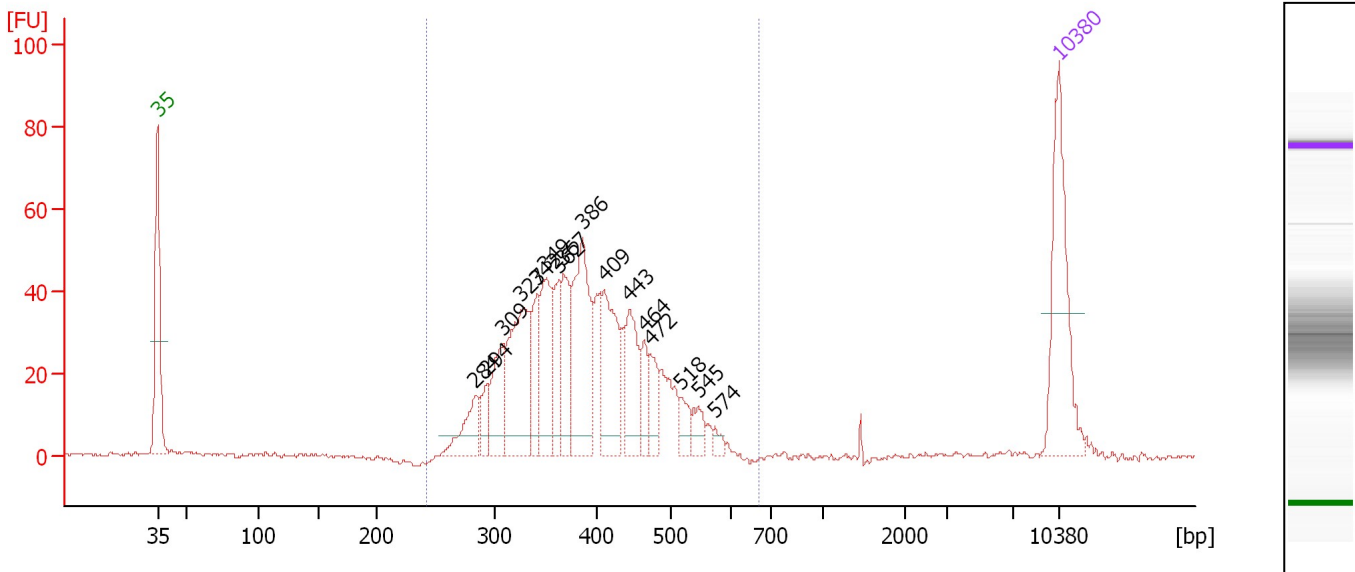
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
228	396	1,000	1,587.5	95	24.4	1,436.75	5,890.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...

4-2 (1:10) [Ribo depleted ssRNA Library (BIOO Kit)]



Overall Results for sample 3 : 4-2 (1:10)

Number of peaks found: 16 Corr. Area 1: 753.2
 Noise: 0.5

Peak table for sample 3 : 4-2 (1:10)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	284	36.13	193.1	
3	294	18.51	95.3	
4	309	43.44	213.1	
5	327	98.07	454.9	
6	342	38.19	169.1	
7	349	61.03	264.9	
8	362	35.05	146.6	
9	367	51.96	214.3	
10	386	100.92	396.5	
11	409	72.65	269.3	
12	443	48.73	166.5	
13	464	20.86	68.2	
14	472	20.94	67.3	
15	518	14.02	41.0	
16	545	13.12	36.4	
17	574	5.72	15.1	
18	10,380	75.00	10.9	Upper Marker

Region table for sample 3 : 4-2 (1:10)

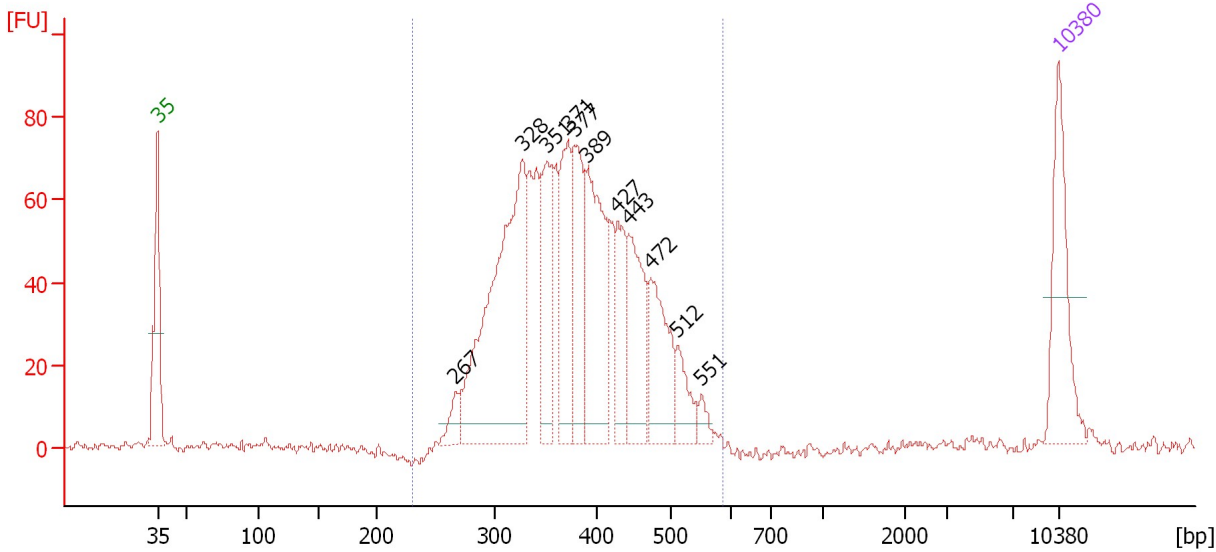
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
243	391	672	753.2	96	17.7	759.23	3,064.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...

4-3 (1:10) [Ribo depleted ssRNA Library (BIOO Kit)]



Overall Results for sample 4 : 4-3 (1:10)

Number of peaks found: 11 Corr. Area 1: 1,215.6
 Noise: 1.0

Peak table for sample 4 : 4-3 (1:10)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	267	22.71	128.9	
3	328	335.99	1,554.2	
4	351	90.80	392.0	
5	371	101.17	412.7	
6	377	99.03	397.7	
7	389	155.80	606.9	
8	427	64.90	230.2	
9	443	92.26	315.7	
10	472	88.54	284.0	
11	512	33.72	99.9	
12	551	10.86	29.9	
13	10,380	75.00	10.9	Upper Marker

Region table for sample 4 : 4-3 (1:10)

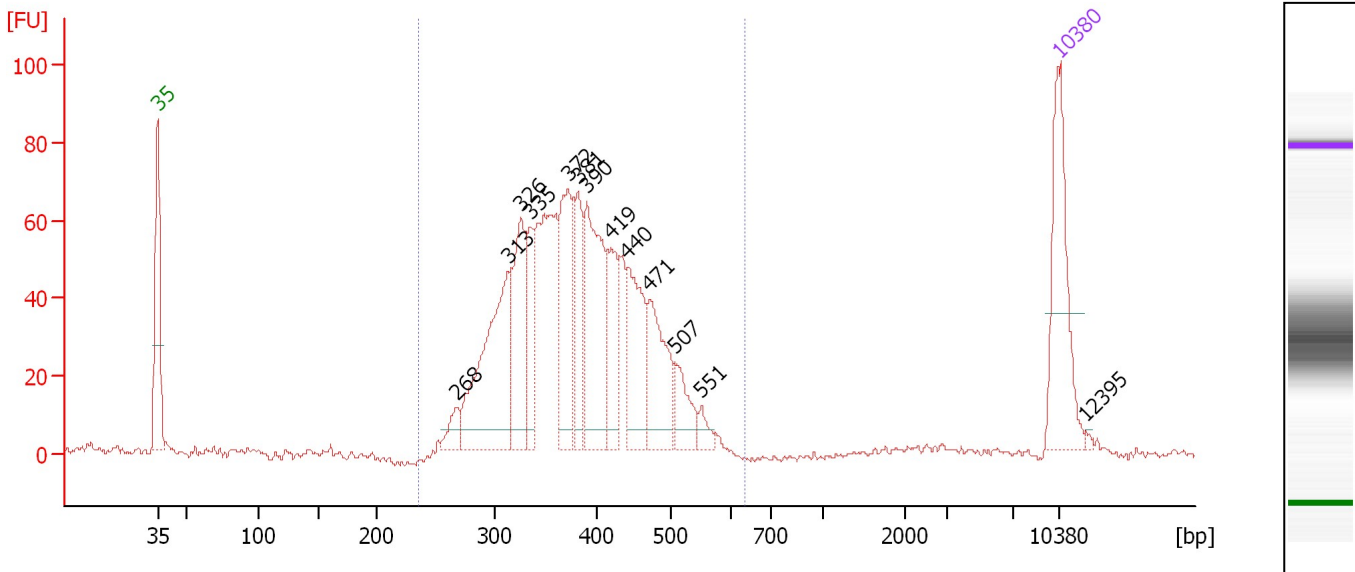
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
231	382	587	1,215.6	98	17.1	1,310.24	5,393.3	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...

4-4 (1:10) [Ribo depleted ssRNA Library (BIOO Kit)]



Overall Results for sample 5 : 4-4 (1:10)

Number of peaks found: 13 Corr. Area 1: 1,102.0
 Noise: 1.0

Peak table for sample 5 : 4-4 (1:10)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	268	16.51	93.5	
3	313	157.14	759.8	
4	326	86.53	402.4	
5	335	48.04	217.0	
6	372	93.34	380.6	
7	381	56.48	224.5	
8	390	123.44	479.5	
9	419	53.47	193.5	
10	440	76.86	264.4	
11	471	72.26	232.3	
12	507	31.01	92.7	
13	551	9.99	27.5	
14	10,380	75.00	10.9	Upper Marker
15	12,395	0.00	0.0	

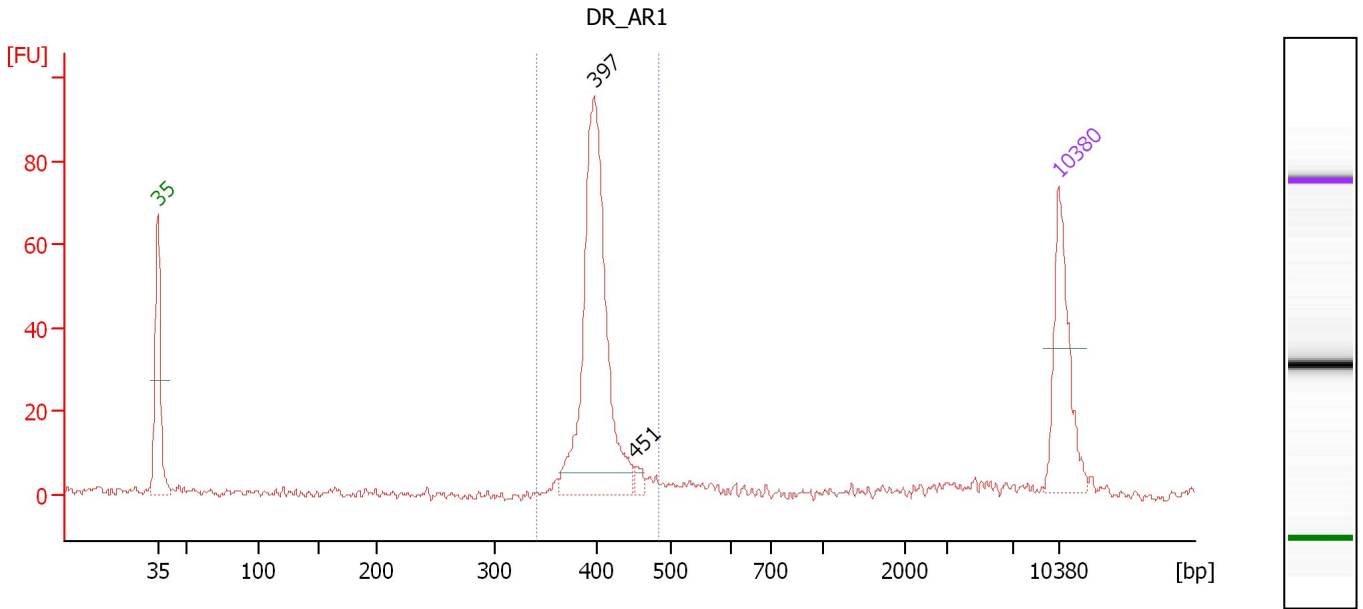
Region table for sample 5 : 4-4 (1:10)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
236	385	633	1,102.0	97	17.2	1,032.42	4,224.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : DR_AR1

Number of peaks found: 2 Corr. Area 1: 260.4
 Noise: 1.0

Peak table for sample 6 : DR_AR1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	397	306.79	1,170.5	
3	451	7.77	26.1	
4	10,380	75.00	10.9	Upper Marker

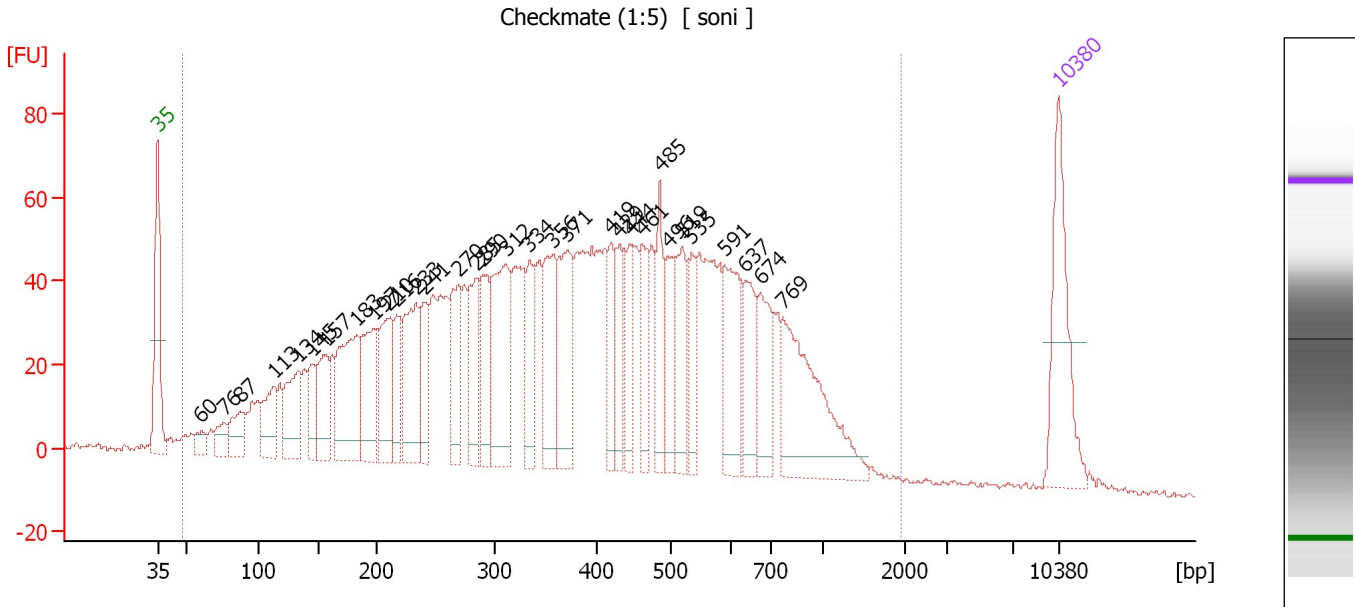
Region table for sample 6 : DR_AR1

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
341	402	484	260.4	87	4.8	315.88	1,192.7	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : Checkmate (1:5)

Number of peaks found: 32 Corr. Area 1: 2,606.8
 Noise: 0.8

Peak table for sample 7 : Checkmate (1:5)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	60	15.95	403.7	
3	76	23.08	461.4	
4	87	34.18	593.1	
5	113	49.65	664.0	
6	134	66.55	754.0	
7	145	34.88	363.8	
8	157	58.81	568.4	
9	183	118.90	984.6	
10	197	77.80	598.0	
11	210	72.05	520.0	
12	216	43.21	302.6	
13	233	93.32	606.8	
14	241	44.60	281.0	
15	270	66.47	373.7	
16	285	59.93	318.5	
17	290	61.19	319.2	
18	312	116.52	566.5	
19	334	48.65	220.7	
20	356	78.75	334.8	
21	371	84.35	344.6	
22	419	42.34	153.2	
23	429	41.84	147.8	
24	444	41.61	141.9	
25	461	41.10	135.0	
26	485	50.60	158.0	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...

... Peak table for sample 7 : Checkmate (1:5)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	496	47.26	144.3	
28	519	58.02	169.3	
29	535	36.64	103.7	
30	591	75.35	193.3	
31	637	49.48	117.7	
32	674	53.96	121.4	
33	769	133.82	263.5	
34	▶ 10,380	75.00	10.9	Upper Marker

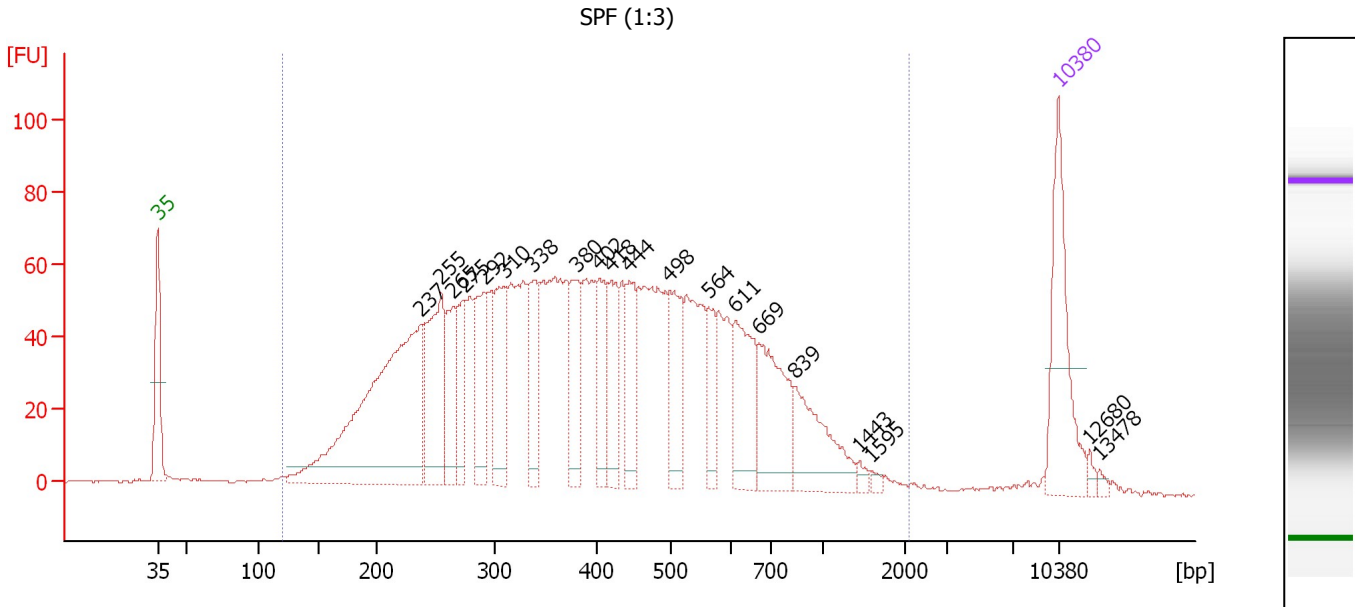
Region table for sample 7 : Checkmate (1:5)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
48	422	1,954	2,606.8	98	55.0	2,795.57	16,671.7	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : SPF (1:3)

Number of peaks found: 20 Corr. Area 1: 2,437.5
 Noise: 0.4

Peak table for sample 8 : SPF (1:3)

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	237	346.09	2,215.2	
3	255	119.03	708.5	
4	265	62.88	359.7	
5	275	49.49	273.1	
6	292	74.29	385.0	
7	310	80.86	394.8	
8	338	61.37	274.8	
9	380	56.20	223.8	
10	402	53.72	202.3	
11	418	52.61	190.8	
12	444	51.43	175.7	
13	498	65.22	198.2	
14	564	32.90	88.5	
15	611	73.39	181.8	
16	669	84.61	191.7	
17	839	68.01	122.8	
18	1,443	4.88	5.1	
19	1,595	2.94	2.8	
20	10,380	75.00	10.9	Upper Marker
21	12,680	0.00	0.0	
22	13,478	0.00	0.0	

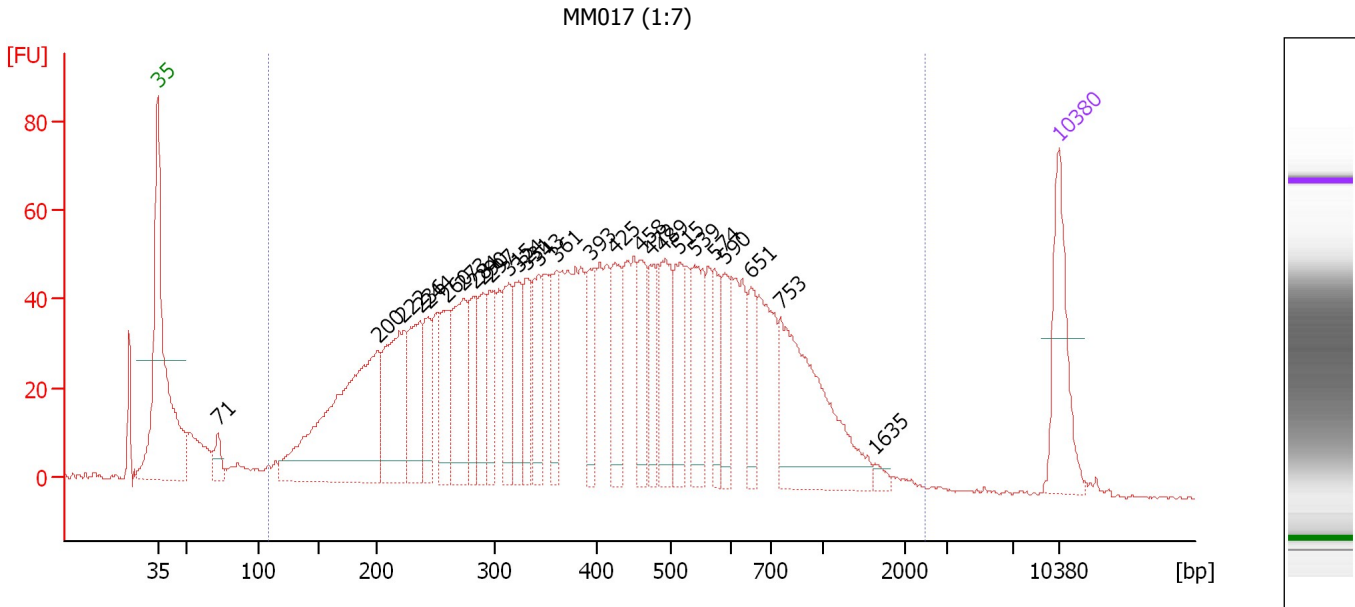
Region table for sample 8 : SPF (1:3)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Color
121	445	2,068	2,437.5	97	53.3	2,148.20	9,848.8	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : MM017 (1:7)

Number of peaks found: 27 Corr. Area 1: 2,266.8
 Noise: 0.2

Peak table for sample 9 : MM017 (1:7)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	71	22.55	478.0	
3	200	343.80	2,598.3	
4	222	158.48	1,080.6	
5	236	98.46	631.2	
6	244	58.81	365.3	
7	260	72.09	420.6	
8	273	111.33	617.2	
9	284	60.37	322.5	
10	290	60.52	316.0	
11	297	60.46	308.8	
12	315	68.23	328.6	
13	324	63.64	297.7	
14	331	55.11	252.1	
15	343	62.68	276.5	
16	361	57.47	241.4	
17	393	58.27	224.6	
18	425	69.16	246.7	
19	458	56.99	188.4	
20	472	55.53	178.2	
21	489	84.88	263.2	
22	515	60.72	178.8	
23	539	71.22	200.2	
24	574	41.16	108.7	
25	590	45.77	117.6	
26	651	44.87	104.5	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...**... Peak table for sample 9 : MM017 (1:7)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	753	195.83	394.2	
28	1,635	6.37	5.9	
29	▶ 10,380	75.00	10.9	Upper Marker

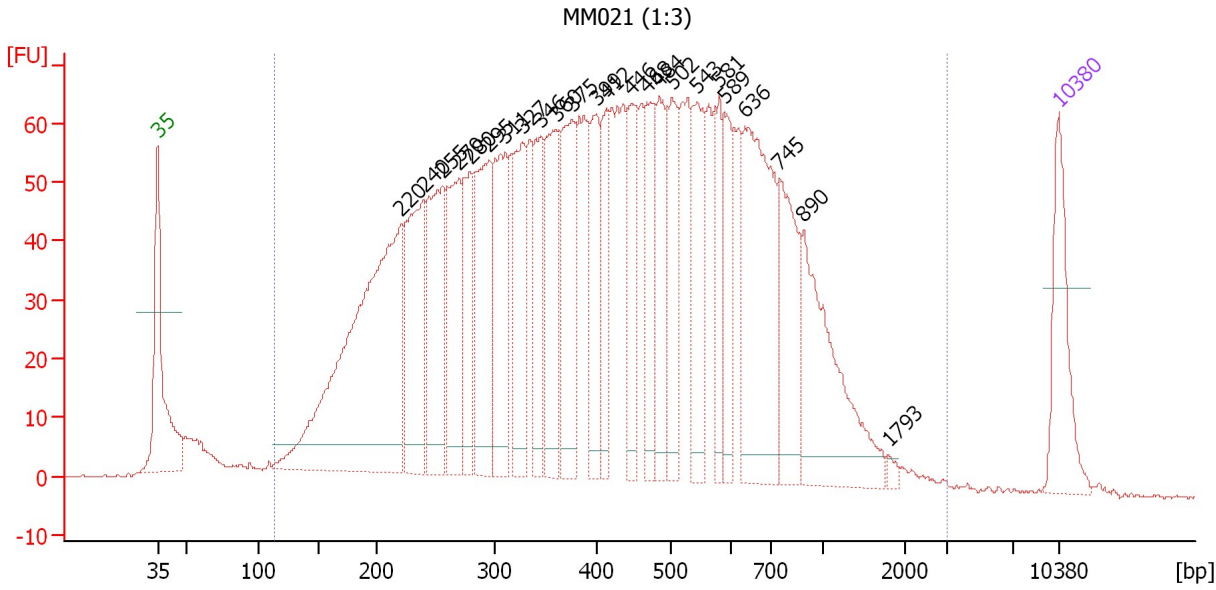
Region table for sample 9 : MM017 (1:7)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
109	463	2,455	2,266.8	95	57.6	2,987.40	14,135.7	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : MM021 (1:3)

Number of peaks found: 24 Corr. Area 1: 2,827.0
 Noise: 0.2


Peak table for sample 10 : MM021 (1:3)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	220	564.19	3,885.8	
3	240	200.94	1,270.4	
4	255	182.88	1,084.6	
5	270	140.18	785.9	
6	280	98.01	530.2	
7	295	164.68	844.8	
8	311	157.73	768.9	
9	327	126.30	585.5	
10	346	88.74	388.7	
11	360	128.51	540.7	
12	375	144.77	584.2	
13	399	100.98	383.6	
14	412	73.11	268.7	
15	446	95.45	324.2	
16	468	94.44	305.9	
17	484	99.05	310.0	
18	502	92.28	278.6	
19	543	106.33	296.4	
20	581	62.03	161.8	
21	589	64.22	165.1	
22	636	241.50	575.5	
23	745	119.62	243.4	
24	890	176.10	299.7	
25	1,793	5.04	4.3	
26	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
Modified: 10/6/2014 4:37:36 PM

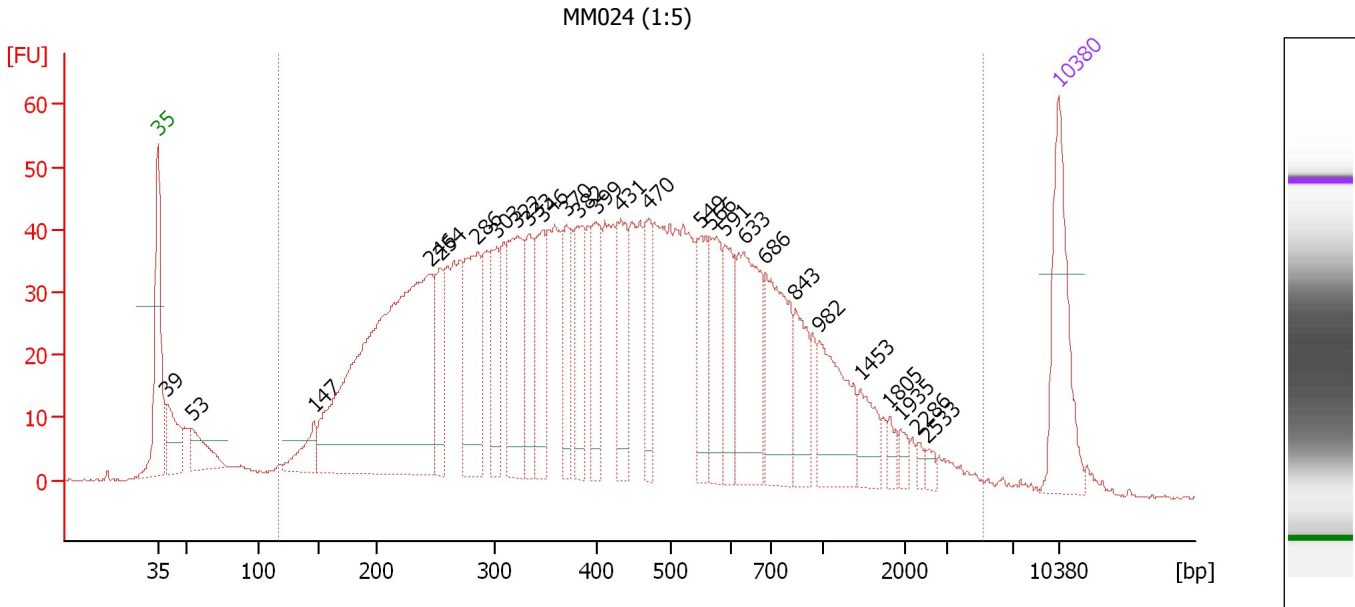
Electropherogram Summary Continued ...**... Region table for sample 10 : MM021 (1:3)**

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Co lor
114	476	3,034	2,827.0	98	59.8	4,318.77	19,715.0	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : MM024 (1:5)

Number of peaks found: 27 Corr. Area 1: 1,954.8
 Noise: 0.1

Peak table for sample 11 : MM024 (1:5)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	39	55.20	2,118.3	
3	53	43.15	1,228.2	
4	147	34.67	358.1	
5	246	508.24	3,130.5	
6	254	52.79	315.3	
7	286	116.92	619.6	
8	303	64.21	321.5	
9	322	108.52	510.1	
10	333	51.37	233.4	
11	346	66.06	288.9	
12	370	49.15	201.0	
13	382	66.77	265.1	
14	399	57.90	219.9	
15	431	63.94	224.6	
16	470	52.11	167.8	
17	549	56.55	156.1	
18	566	57.74	154.5	
19	591	50.91	130.6	
20	633	110.72	264.9	
21	686	99.55	219.8	
22	843	44.72	80.3	
23	982	72.34	111.6	
24	1,453	27.06	28.2	
25	1,805	8.81	7.4	
26	1,935	7.14	5.6	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
Modified: 10/6/2014 4:37:36 PM

Electropherogram Summary Continued ...**... Peak table for sample 11 : MM024 (1:5)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	2,286	4.54	3.0	
28	2,533	5.01	3.0	
29	▶ 10,380	75.00	10.9	Upper Marker

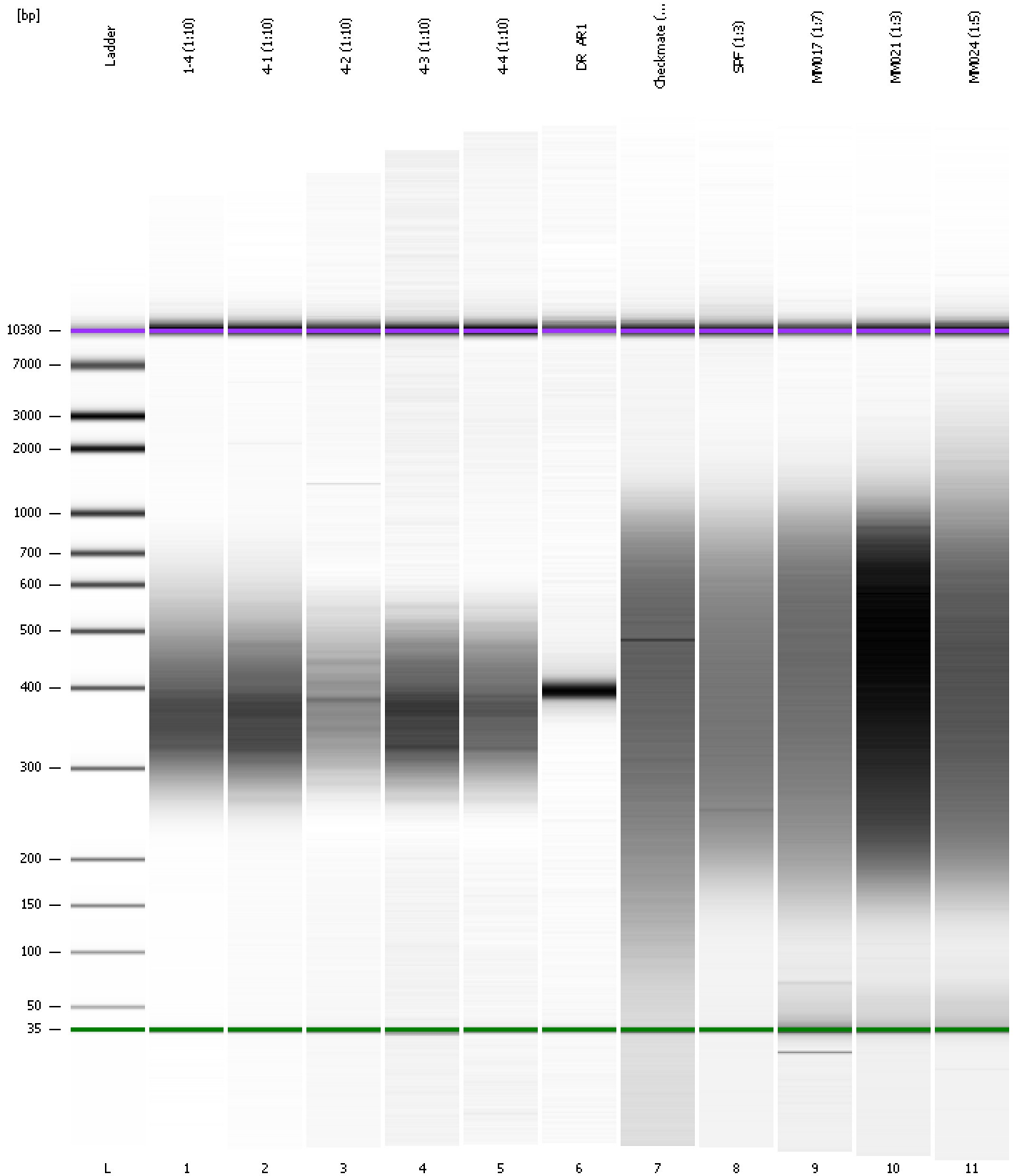
Region table for sample 11 : MM024 (1:5)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
118	547	5,134	1,954.8	94	89.8	2,890.71	13,181.9	■

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
Modified: 10/6/2014 4:37:36 PM

Gel Image

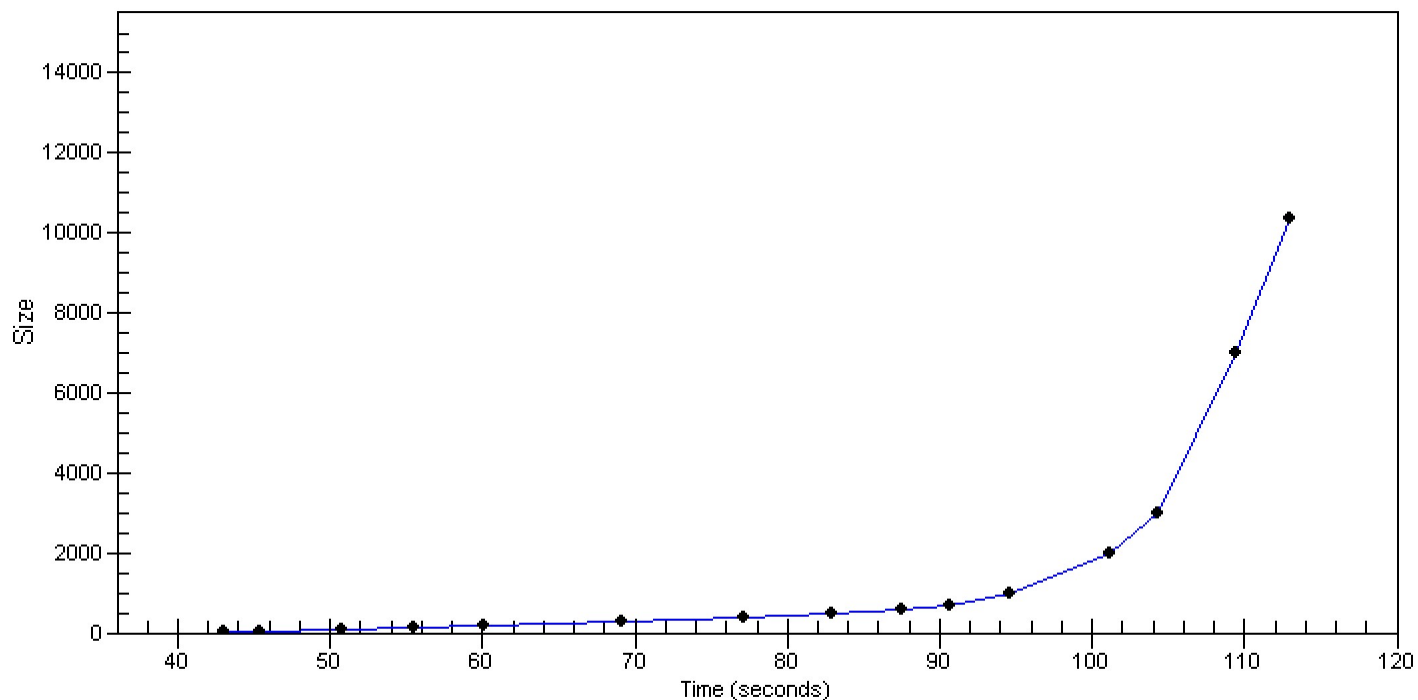


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
Modified: 10/6/2014 4:37:36 PM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad

Created: 10/6/2014 3:51:32 PM
 Modified: 10/6/2014 4:37:36 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		10/6/2014 4:31:59 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2014-10-06\2014-10-06_003.xad)		Instrument	Run		10/6/2014 3:51:38 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		10/6/2014 3:51:38 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		10/6/2014 3:51:38 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		10/6/2014 3:51:38 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		10/6/2014 3:51:38 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		10/6/2014 3:51:38 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		10/6/2014 3:51:38 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1