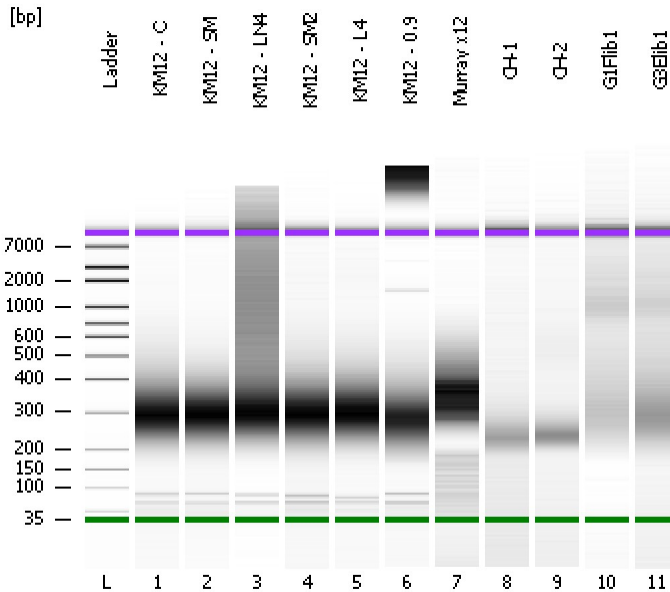


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
Modified: 10/17/2014 11:20:39 AM

Electrophoresis File Run Summary



Instrument Information:

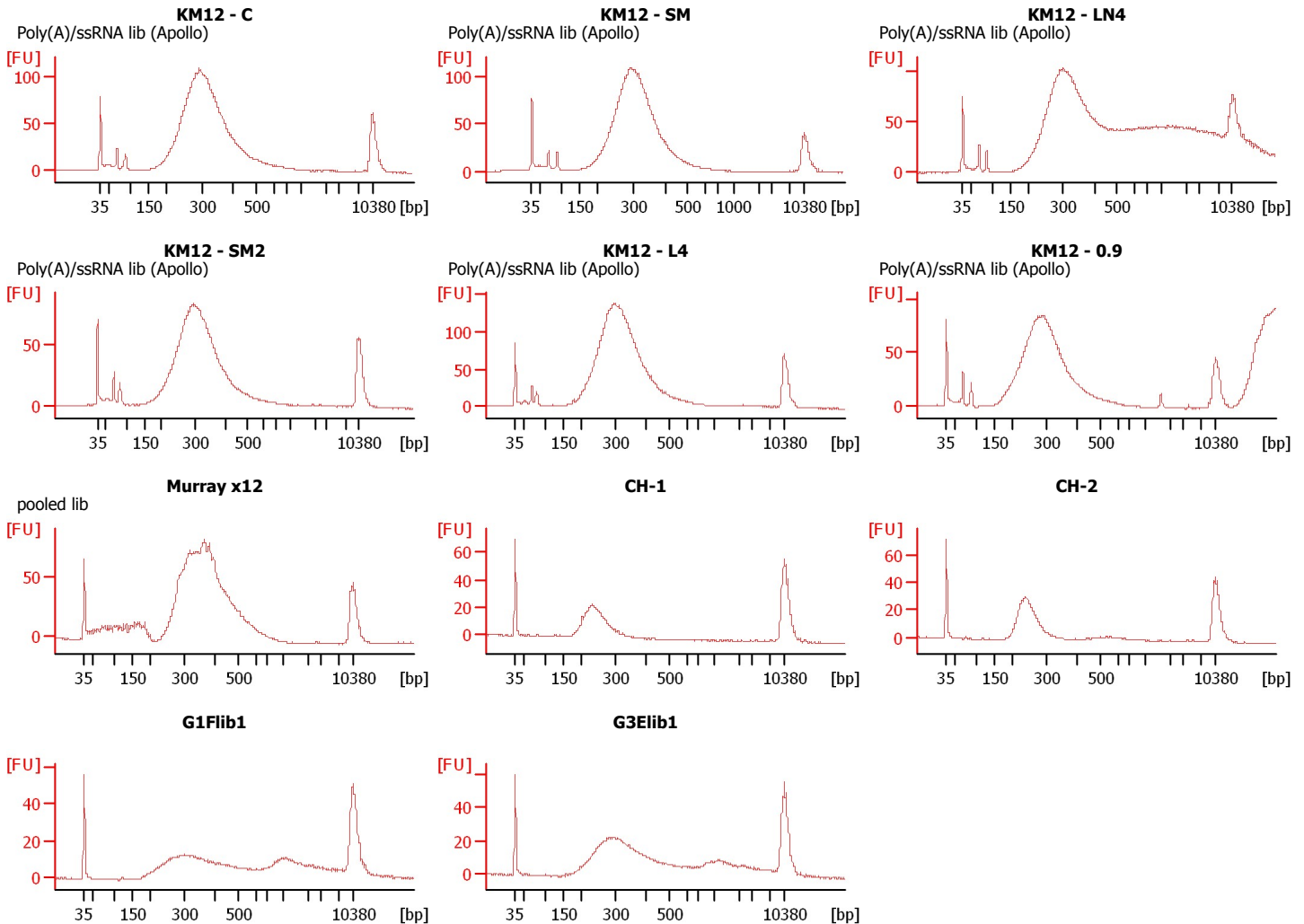
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
Modified: 10/17/2014 11:20:39 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
KM12 - C	Poly(A)/ssRNA lib (Apollo)	<input type="checkbox"/>	✓			
KM12 - SM	Poly(A)/ssRNA lib (Apollo)	<input type="checkbox"/>	✓			
KM12 - LN4	Poly(A)/ssRNA lib (Apollo)	<input type="checkbox"/>	✓			
KM12 - SM2	Poly(A)/ssRNA lib (Apollo)	<input type="checkbox"/>	✓			
KM12 - L4	Poly(A)/ssRNA lib (Apollo)	<input type="checkbox"/>	✓			
KM12 - 0.9	Poly(A)/ssRNA lib (Apollo)	<input type="checkbox"/>	✓			
Murray x12	pooled lib	<input type="checkbox"/>	✓			
CH-1		<input type="checkbox"/>	✓			
CH-2		<input type="checkbox"/>	✓			
G1Flib1		<input type="checkbox"/>	✓			
G3Elib1		<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\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
Modified: 10/17/2014 11:20:39 AM

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

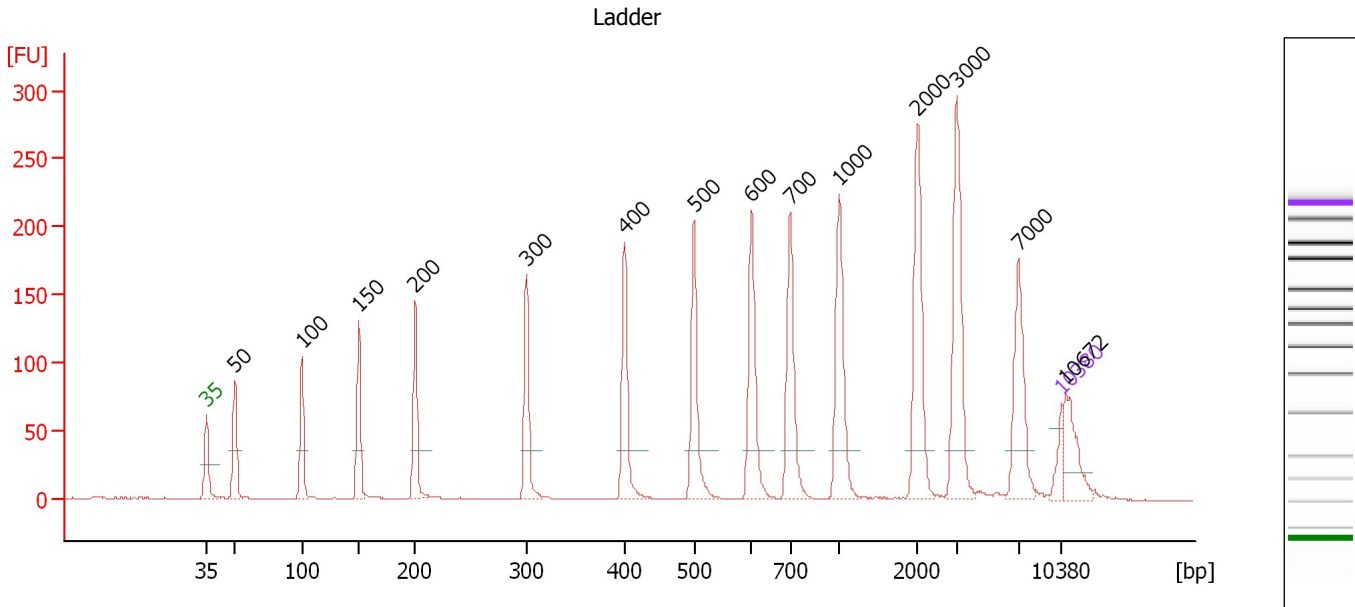
Ladder

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\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

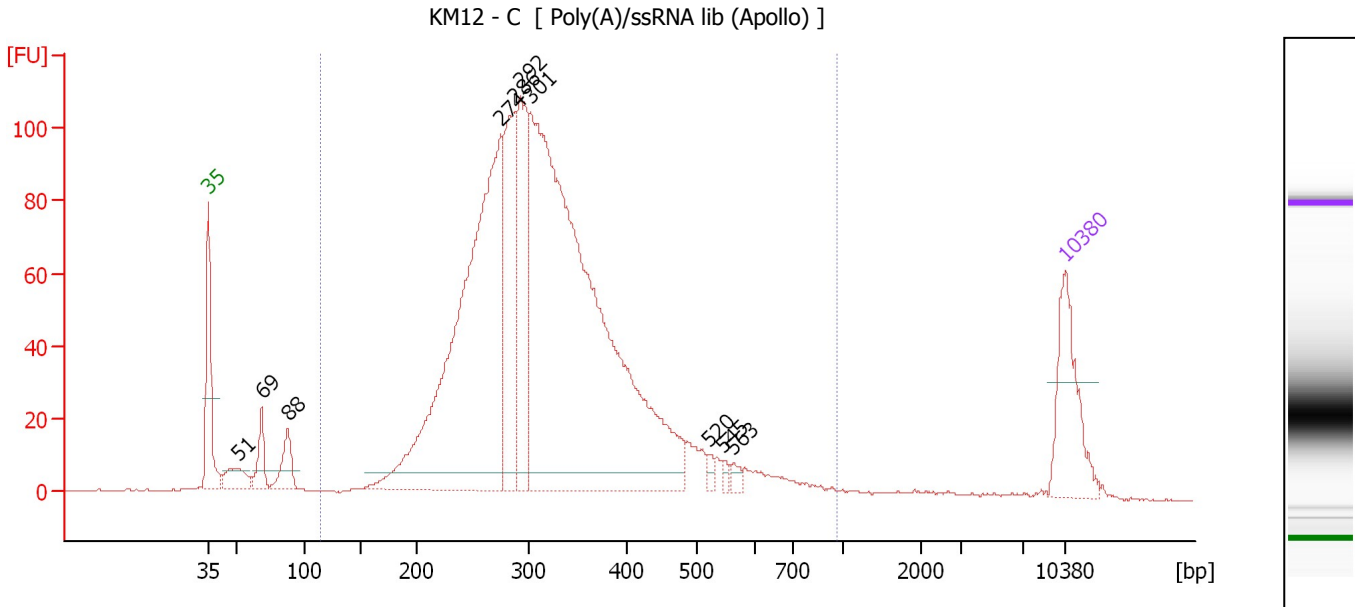
Peak table for Ladder

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
16	10,672	0.00	0.0	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : KM12 - C

Number of peaks found: 10 Corr. Area 1: 2,076.9
 Noise: 0.2

Peak table for sample 1 : KM12 - C

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	51	36.47	1,089.2	
3	69	39.25	867.9	
4	88	45.58	785.6	
5	274	720.80	3,981.4	
6	286	197.15	1,043.6	
7	292	185.20	961.1	
8	301	1,113.56	5,602.5	
9	520	7.91	23.1	
10	545	5.38	14.9	
11	563	7.74	20.8	
12	10,380	75.00	10.9	Upper Marker

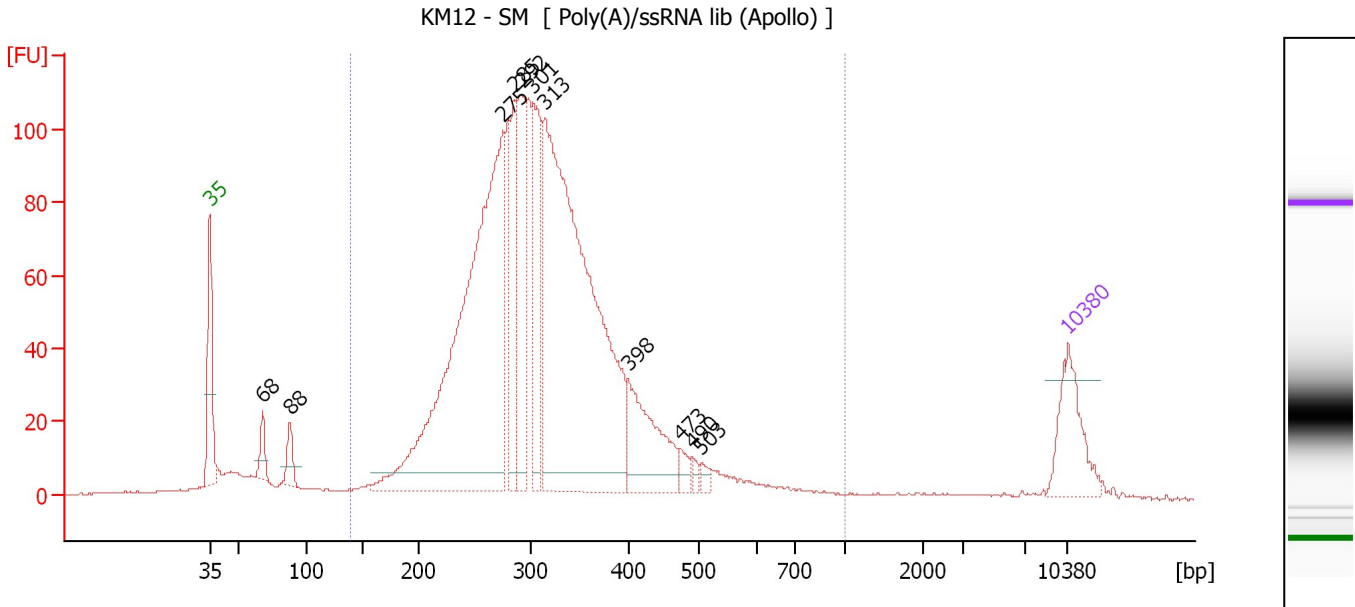
Region table for sample 1 : KM12 - C

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
114	327	969	2,076.9	93	29.2	2,318.55	11,765.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 : KM12 - SM

Number of peaks found: 11 Corr. Area 1: 2,025.3
 Noise: 0.1

Peak table for sample 2 : KM12 - SM

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	68	35.67	793.5	
3	88	38.93	668.3	
4	275	951.24	5,248.3	
5	285	141.78	753.5	
6	292	187.18	972.0	
7	301	187.48	942.2	
8	313	981.33	4,744.3	
9	398	154.81	588.7	
10	473	16.79	53.8	
11	490	8.39	25.9	
12	503	10.09	30.4	
13	10,380	75.00	10.9	Upper Marker

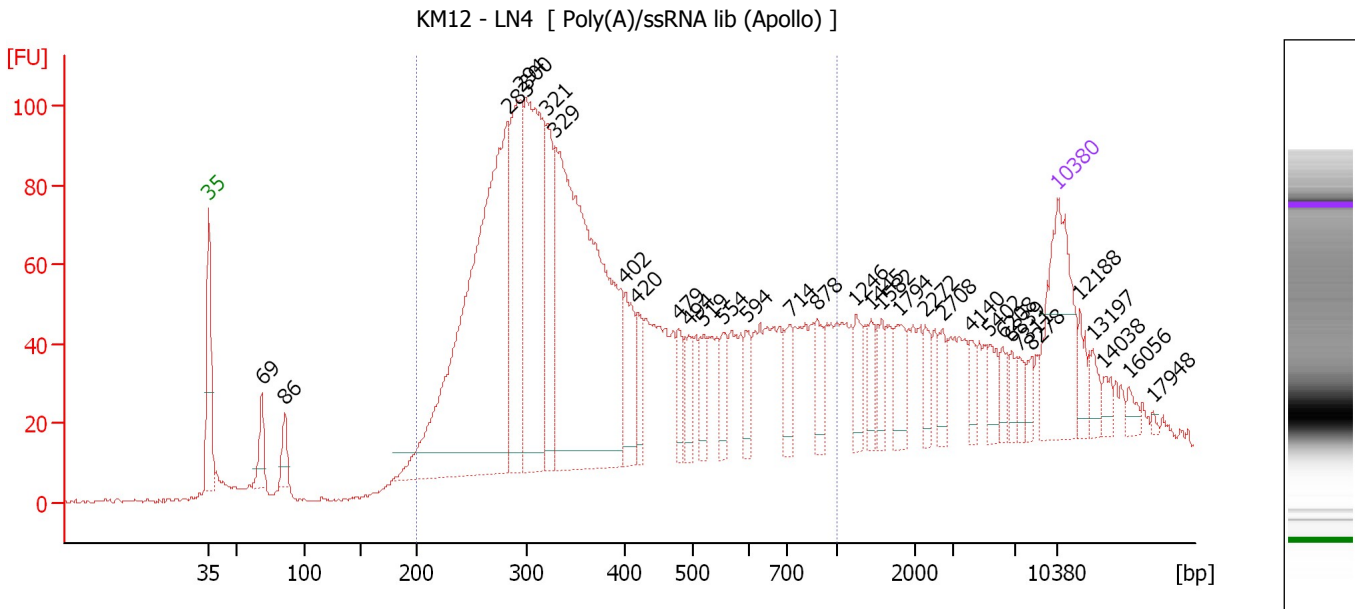
Region table for sample 2 : KM12 - SM

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
140	318	1,000	2,025.3	92	26.3	3,049.66	15,671.8	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...



Overall Results for sample 3 : KM12 - LN4

Number of peaks found: 33 Corr. Area 1: 2,234.2
 Noise: 0.5

Peak table for sample 3 : KM12 - LN4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	69	31.16	688.7	
3	86	23.38	413.2	
4	283	516.73	2,763.0	
5	294	153.74	791.1	
6	300	236.34	1,192.6	
7	321	82.02	387.7	
8	329	468.64	2,159.9	
9	402	46.94	176.8	
10	420	26.00	93.8	
11	479	21.44	67.8	
12	494	20.51	62.9	
13	519	21.19	61.8	
14	554	20.14	55.1	
15	594	20.09	51.2	
16	714	24.52	52.1	
17	878	22.65	39.1	
18	1,246	22.11	26.9	
19	1,445	14.93	15.7	
20	1,582	15.34	14.7	
21	1,794	21.65	18.3	
22	2,272	13.10	8.7	
23	2,708	12.49	7.0	
24	4,140	9.77	3.6	
25	5,402	13.46	3.8	
26	6,208	9.91	2.4	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad


Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...

... Peak table for sample 3 : KM12 - LN4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	6,839	7.69	1.7	
28	7,311	8.78	1.8	
29	8,278	7.83	1.4	
30	10,380	75.00	10.9	Upper Marker
31	12,188	0.00	0.0	
32	13,197	0.00	0.0	
33	14,038	0.00	0.0	
34	16,056	0.00	0.0	
35	17,948	0.00	0.0	

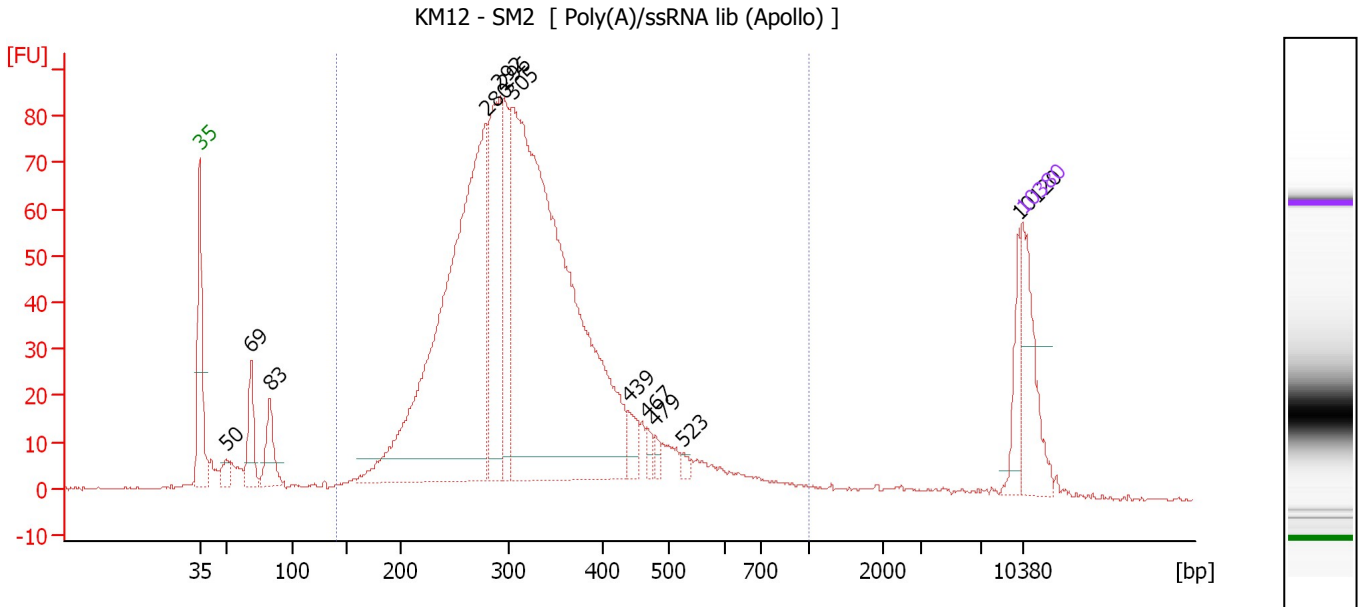
Region table for sample 3 : KM12 - LN4

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	412	1,000	2,234.2	79	42.0	2,013.82	8,902.1	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...



Overall Results for sample 4 : KM12 - SM2

Number of peaks found: 12 Corr. Area 1: 1,639.7
 Noise: 0.3

Peak table for sample 4 : KM12 - SM2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	50	21.14	644.3	
3	69	70.41	1,556.4	
4	83	67.34	1,229.7	
5	280	859.68	4,654.5	
6	292	261.54	1,355.8	
7	296	130.57	667.8	
8	305	1,139.06	5,652.2	
9	439	27.00	93.1	
10	467	11.27	36.6	
11	479	10.09	31.9	
12	523	7.57	21.9	
13	10,120	36.46	5.5	
14	10,380	75.00	10.9	Upper Marker

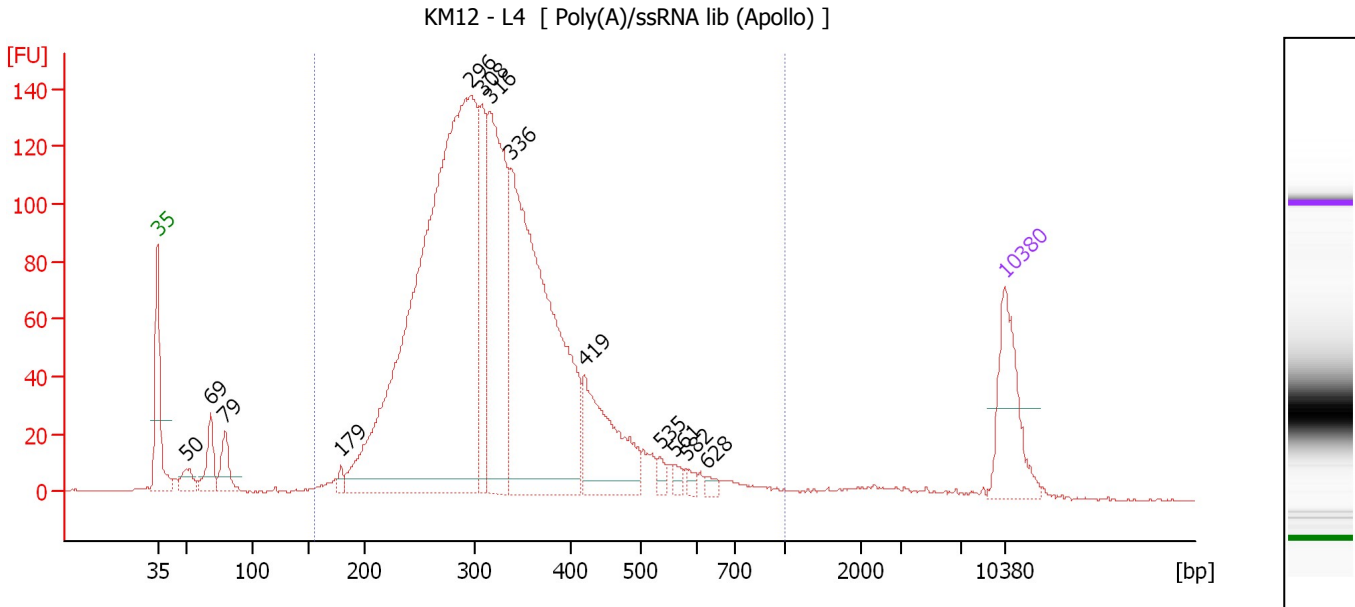
Region table for sample 4 : KM12 - SM2

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
142	328	1,000	1,639.7	90	29.9	2,723.82	13,811.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : KM12 - L4

Number of peaks found: 13 Corr. Area 1: 2,604.8
 Noise: 0.3

Peak table for sample 5 : KM12 - L4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	50	25.72	773.1	
3	69	47.27	1,039.8	
4	79	43.51	830.4	
5	179	8.69	73.4	
6	296	1,145.78	5,862.9	
7	308	121.62	599.0	
8	316	300.80	1,443.5	
9	336	580.74	2,615.4	
10	419	140.99	509.6	
11	535	9.06	25.6	
12	561	7.53	20.4	
13	582	6.83	17.8	
14	628	6.55	15.8	
15	10,380	75.00	10.9	Upper Marker

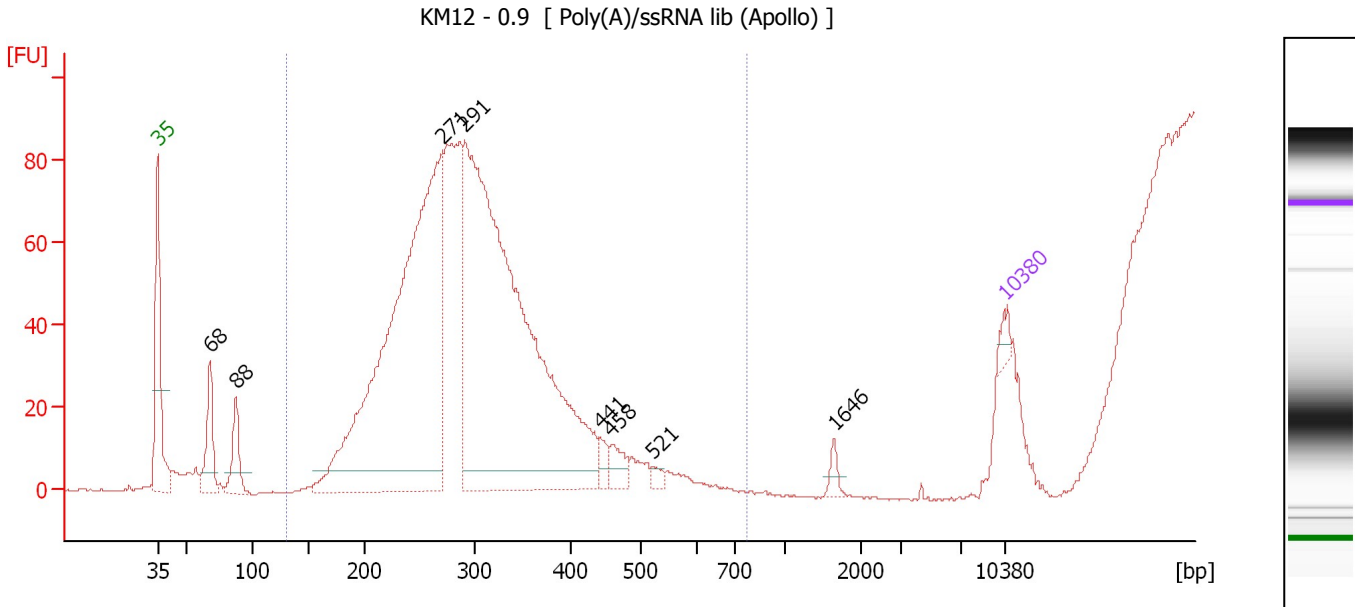
Region table for sample 5 : KM12 - L4

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
156	330	1,000	2,604.8	94	28.0	2,463.19	12,261.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...



Overall Results for sample 6 : KM12 - 0.9

Number of peaks found: 8 Corr. Area 1: 576.4
 Noise: 0.2

Peak table for sample 6 : KM12 - 0.9

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	68	629.30	13,974.8	
3	88	481.94	8,311.1	
4	271	7,110.04	39,786.8	
5	291	8,402.29	43,686.6	
6	441	125.96	432.4	
7	458	182.93	604.6	
8	521	61.25	178.1	
9	1,646	79.83	73.5	
10	10,380	75.00	10.9	Upper Marker

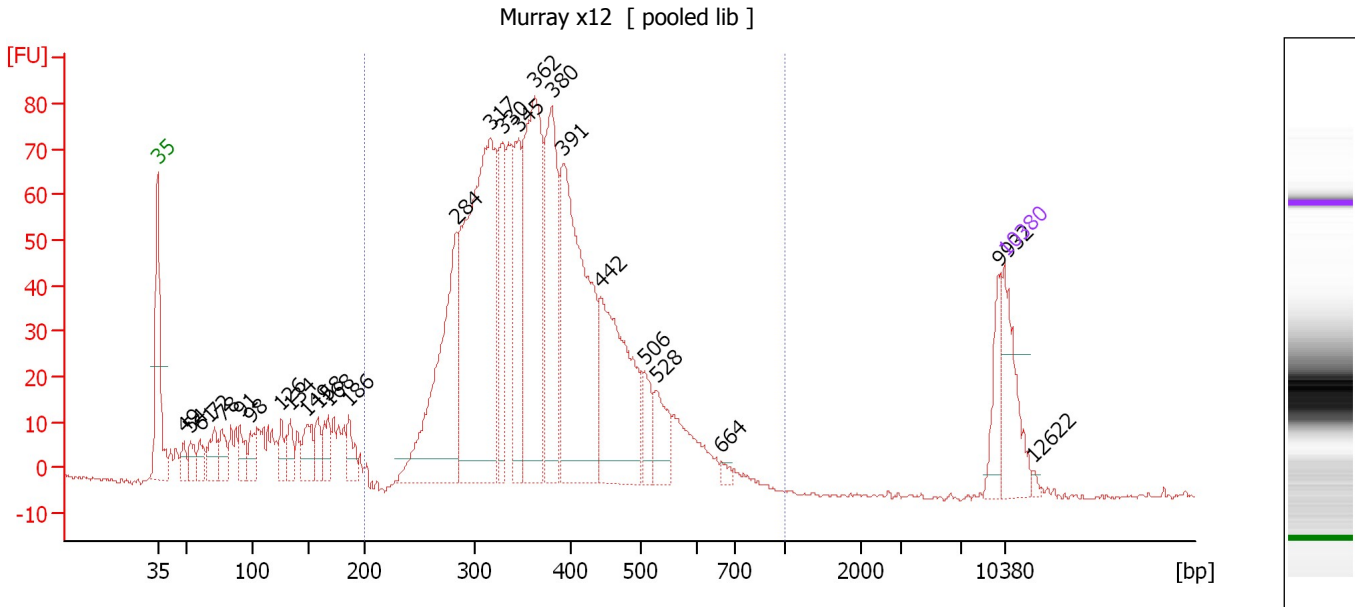
Region table for sample 6 : KM12 - 0.9

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
131	281	773	576.4	98	10.6	6,683.75	36,460.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...



Overall Results for sample 7 : Murray x12

Number of peaks found: 26 Corr. Area 1: 1,505.7
 Noise: 0.5

Peak table for sample 7 : Murray x12

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	49	24.49	752.6	
3	54	29.09	820.9	
4	61	27.54	684.6	
5	72	48.49	1,026.1	
6	78	38.48	751.5	
7	91	29.27	488.5	
8	98	37.74	580.7	
9	126	31.64	380.3	
10	134	36.37	410.7	
11	149	51.72	524.7	
12	158	31.05	296.9	
13	168	40.47	364.7	
14	186	36.89	300.8	
15	284	315.96	1,686.7	
16	317	568.00	2,717.7	
17	330	120.60	554.5	
18	345	144.57	635.5	
19	362	343.62	1,437.4	
20	380	231.10	922.1	
21	391	397.44	1,538.5	
22	442	256.00	876.8	
23	506	42.42	127.0	
24	528	54.45	156.2	
25	664	7.57	17.3	
26	9,932	42.36	6.5	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...**... Peak table for sample 7 : Murray x12**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	10,380	75.00	10.9	Upper Marker
28	12,622	0.00	0.0	

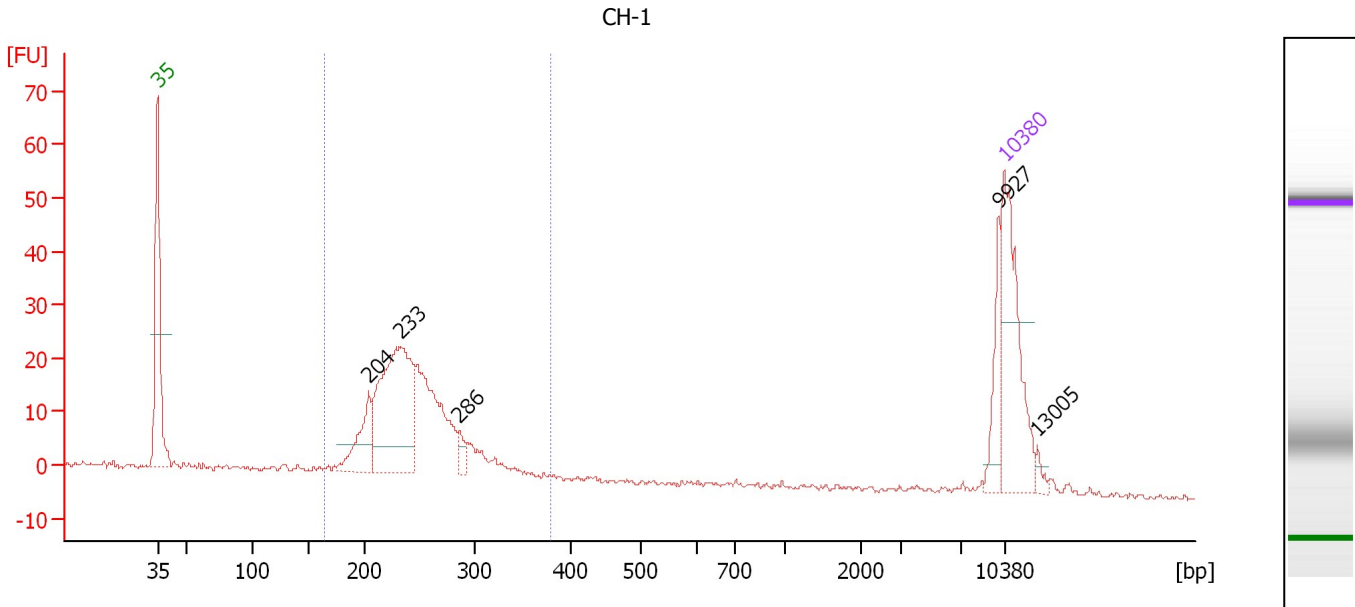
Region table for sample 7 : Murray x12

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	378	1,000	1,505.7	84	22.6	2,734.06	11,657.2	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...



Overall Results for sample 8 : CH-1

Number of peaks found: 5 Corr. Area 1: 254.9
 Noise: 0.5

Peak table for sample 8 : CH-1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	204	46.47	344.4	
3	233	168.46	1,093.3	
4	286	11.48	60.7	
5	9,927	23.83	3.6	
6	10,380	75.00	10.9	Upper Marker
7	13,005	0.00	0.0	

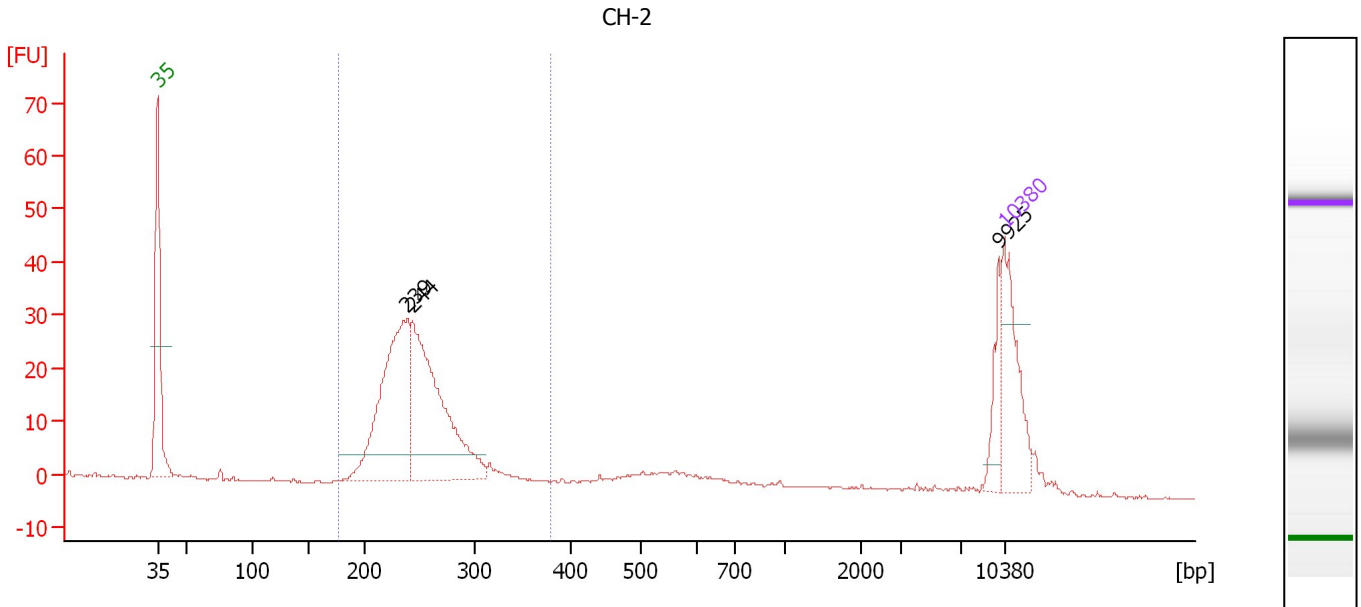
Region table for sample 8 : CH-1

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
164	245	378	254.9	85	13.9	371.68	2,332.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...



Overall Results for sample 9 : CH-2

Number of peaks found: 3 Corr. Area 1: 288.7
 Noise: 0.2

Peak table for sample 9 : CH-2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	239	261.02	1,656.9	
3	244	269.13	1,671.9	
4	9,925	32.57	5.0	
5	10,380	75.00	10.9	Upper Marker

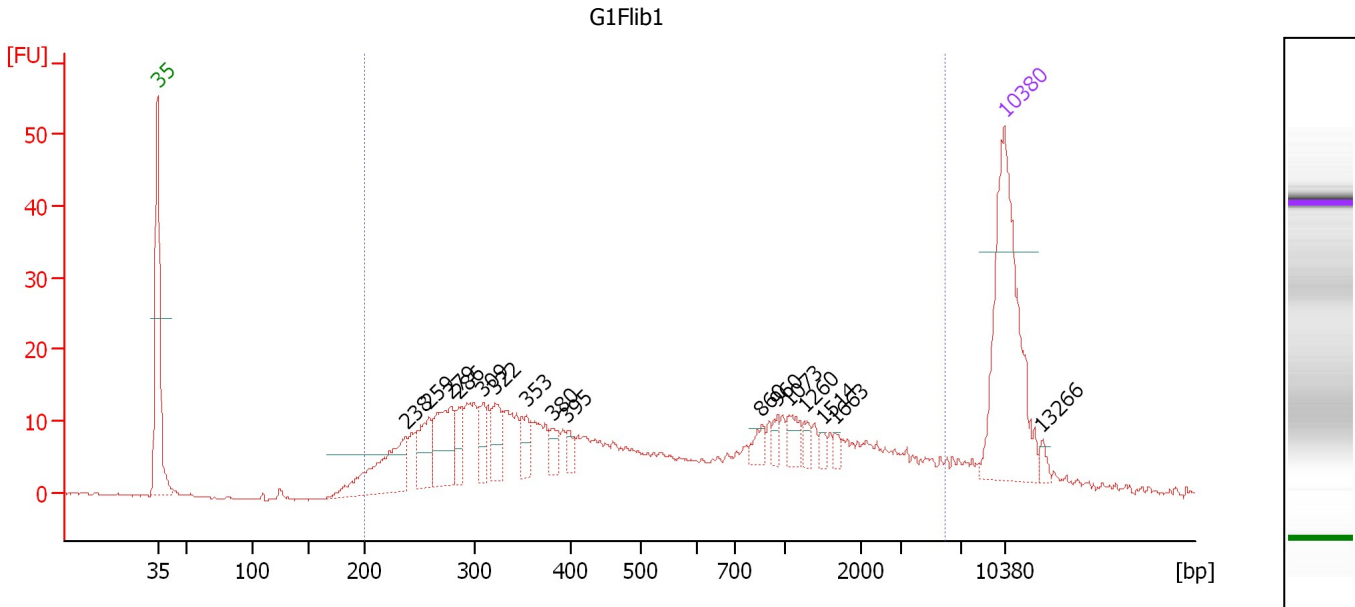
Region table for sample 9 : CH-2

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
178	250	379	288.7	75	12.6	572.58	3,505.8	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...



Overall Results for sample 10 : G1Flib1

Number of peaks found: 16 Corr. Area 1: 450.9
 Noise: 0.1

Peak table for sample 10 : G1Flib1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	238	47.21	300.2	
3	259	23.78	139.2	
4	279	35.71	194.1	
5	286	12.30	65.1	
6	309	11.66	57.2	
7	322	15.93	75.0	
8	353	11.22	48.1	
9	380	7.58	30.2	
10	395	4.89	18.7	
11	860	5.94	10.5	
12	960	4.21	6.7	
13	1,073	7.94	11.2	
14	1,260	3.77	4.5	
15	1,514	3.01	3.0	
16	1,663	2.87	2.6	
17	10,380	75.00	10.9	Upper Marker
18	13,266	0.00	0.0	

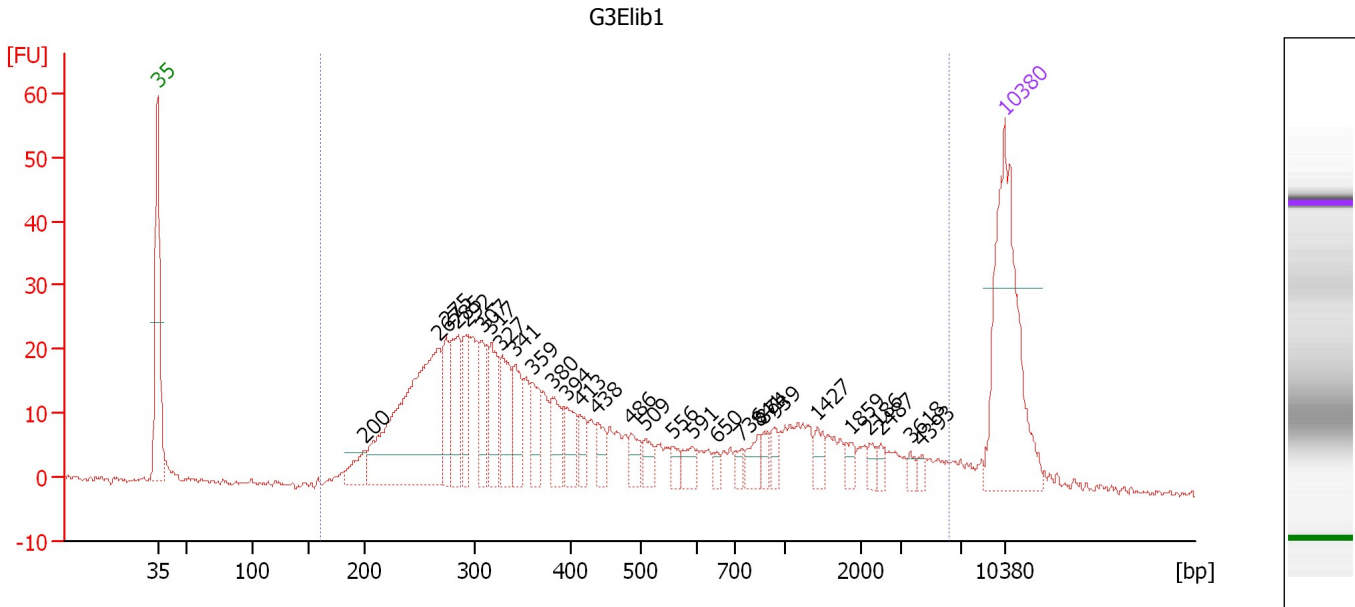
Region table for sample 10 : G1Flib1

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	885	5,893	450.9	94	100.0	490.30	1,873.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...



Overall Results for sample 11 : G3Elib1

Number of peaks found: 29 Corr. Area 1: 665.5
 Noise: 0.5

Peak table for sample 11 : G3Elib1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	200	13.35	101.1	
3	267	144.00	816.7	
4	275	23.73	130.9	
5	285	32.65	173.7	
6	292	23.54	122.1	
7	307	23.36	115.1	
8	317	26.37	126.2	
9	327	27.71	128.2	
10	341	22.43	99.5	
11	359	17.52	74.0	
12	380	16.73	66.7	
13	394	16.00	61.5	
14	413	10.31	37.8	
15	438	10.00	34.6	
16	486	10.09	31.5	
17	509	7.86	23.4	
18	556	6.00	16.3	
19	591	8.03	20.6	
20	650	3.48	8.1	
21	736	3.95	8.1	
22	841	8.35	15.0	
23	874	4.98	8.6	
24	939	5.12	8.3	
25	1,427	7.00	7.4	
26	1,859	4.67	3.8	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad


Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 AM

Electropherogram Summary Continued ...

... Peak table for sample 11 : G3Elib1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	2,186	3.50	2.4	
28	2,487	3.40	2.1	
29	3,618	2.92	1.2	
30	4,393	2.37	0.8	
31	10,380	75.00	10.9	Upper Marker

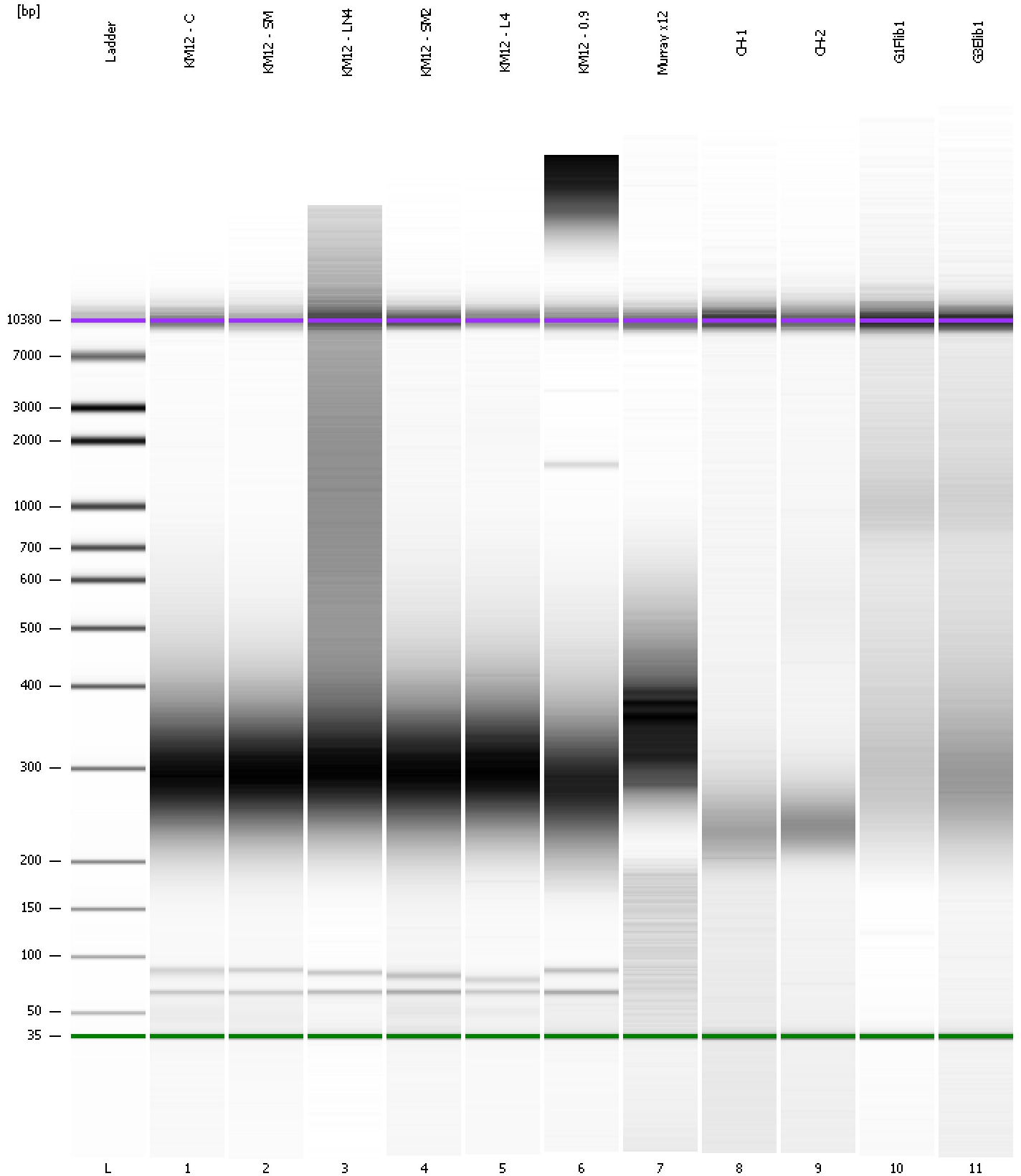
Region table for sample 11 : G3Elib1

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
162	752	6,243	665.5	96	100.0	661.20	2,831.7	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
Modified: 10/17/2014 11:20:39 AM

Gel Image

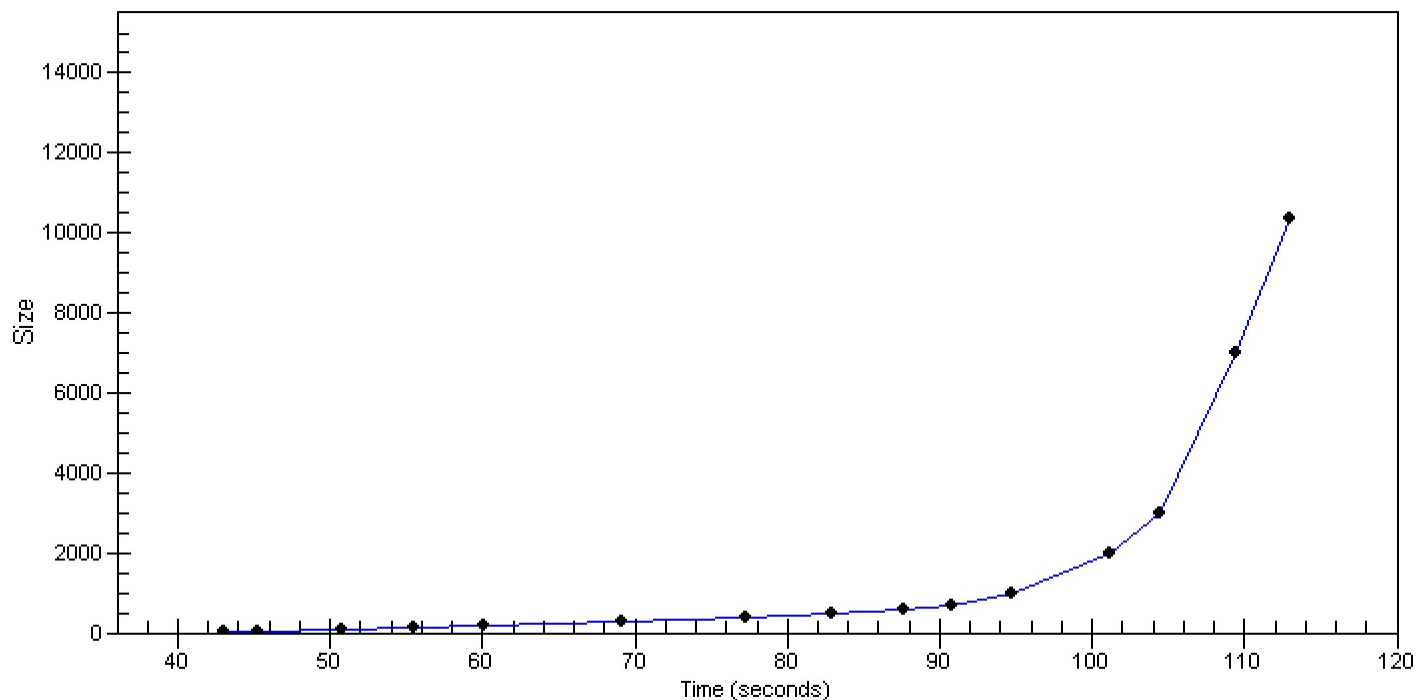


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
Modified: 10/17/2014 11:20:39 AM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad

Created: 10/17/2014 10:37:43 AM
 Modified: 10/17/2014 11:20:39 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		10/17/2014 11:19:03 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2014-10-17\2014-10-17_001.xad)		Instrument	Run		10/17/2014 10:37:48 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		10/17/2014 10:37:48 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		10/17/2014 10:37:48 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		10/17/2014 10:37:48 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		10/17/2014 10:37:48 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		10/17/2014 10:37:48 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		10/17/2014 10:37:48 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1