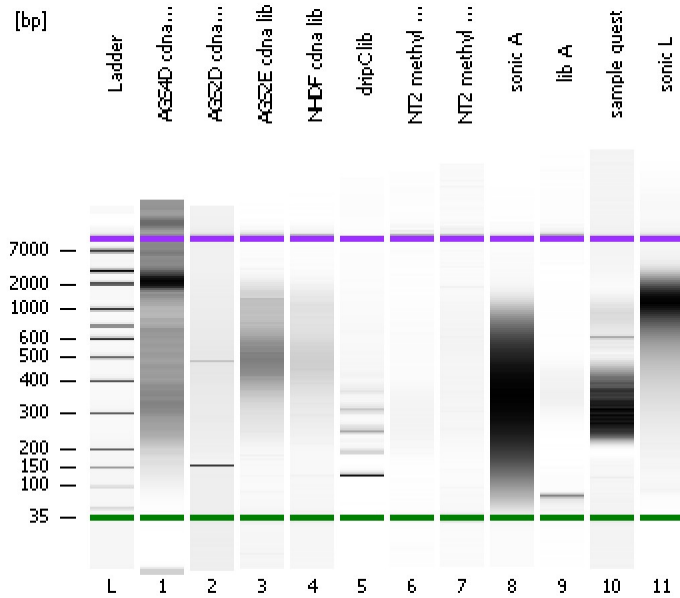


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
Modified: 5/3/2013 2:59:03 PM

Electrophoresis File Run Summary



Instrument Information:

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

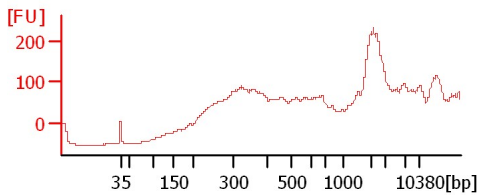
Assay Information:

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

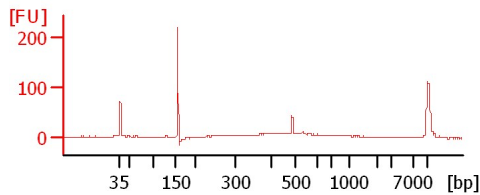
Chip Information:

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

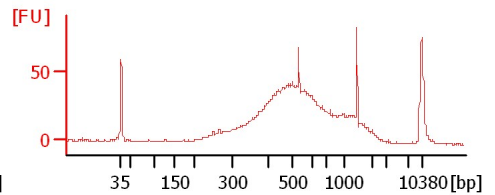
AGS4D cdna lib



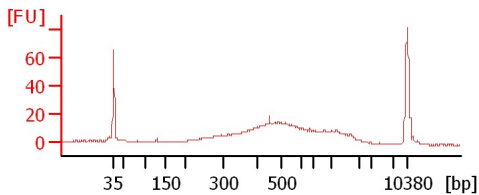
AGS2D cdna lib



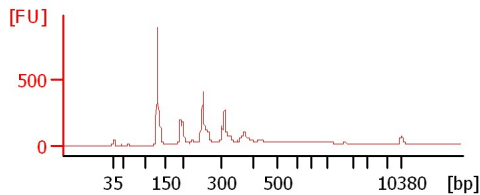
AGS2E cdna lib



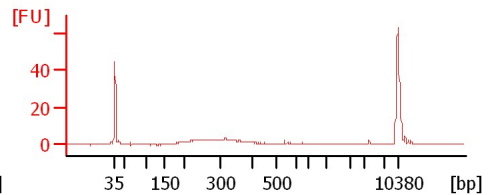
NHDF cdna lib



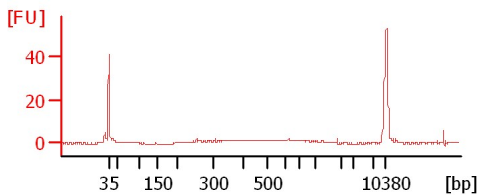
dripC lib



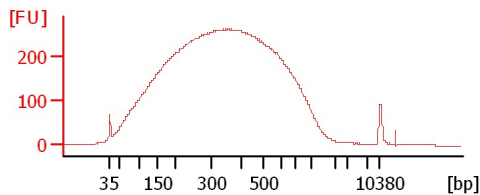
NT2 methyl lib 7



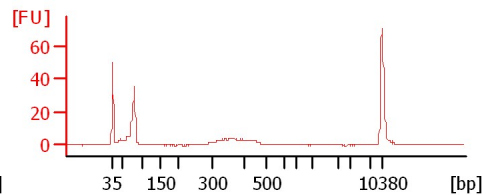
NT2 methyl lib 5



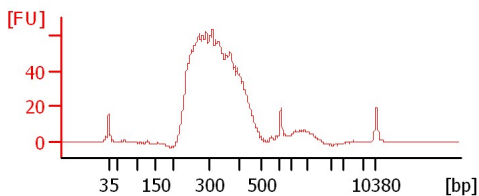
sonic A



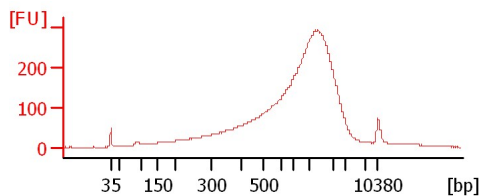
lib A



sample quest



sonic L



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
AGS4D cdna lib		<input type="checkbox"/>	✓			
AGS2D cdna lib		<input type="checkbox"/>	✓			
AGS2E cdna lib		<input type="checkbox"/>	✓			
NHDF cdna lib		<input type="checkbox"/>	✓			
dripC lib		<input type="checkbox"/>	✓			
NT2 methyl lib 7		<input type="checkbox"/>	✓			
NT2 methyl lib 5		<input type="checkbox"/>	✓			
sonic A		<input type="checkbox"/>	✓			
lib A		<input type="checkbox"/>	✓			
sample guest		<input type="checkbox"/>	✓			
sonic L		<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\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 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

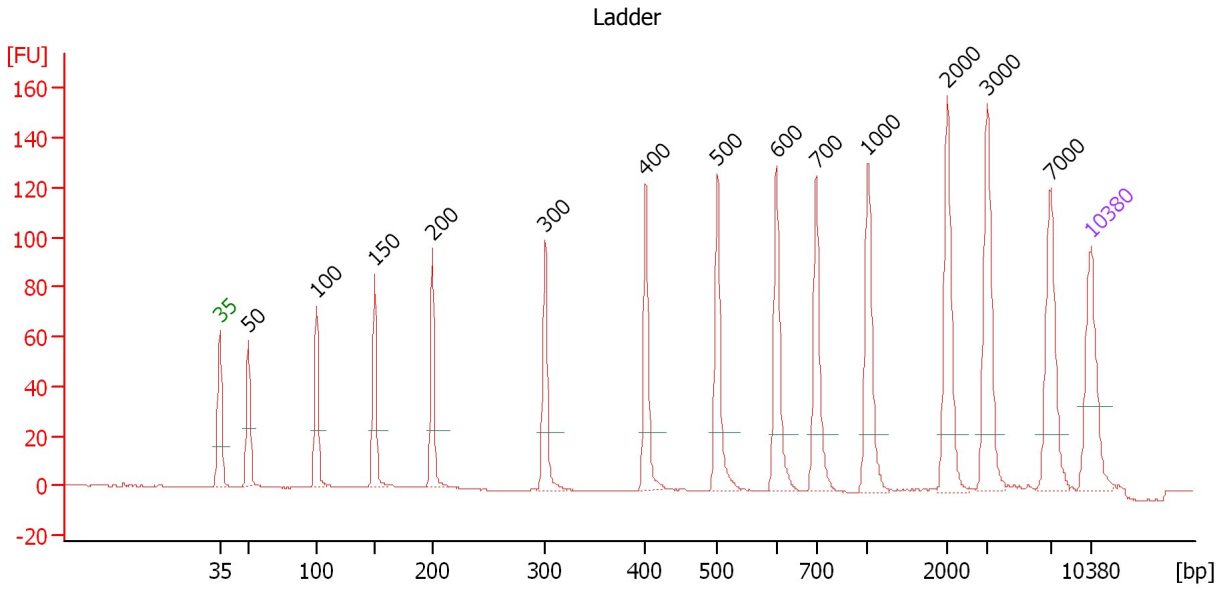
Ladder

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\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

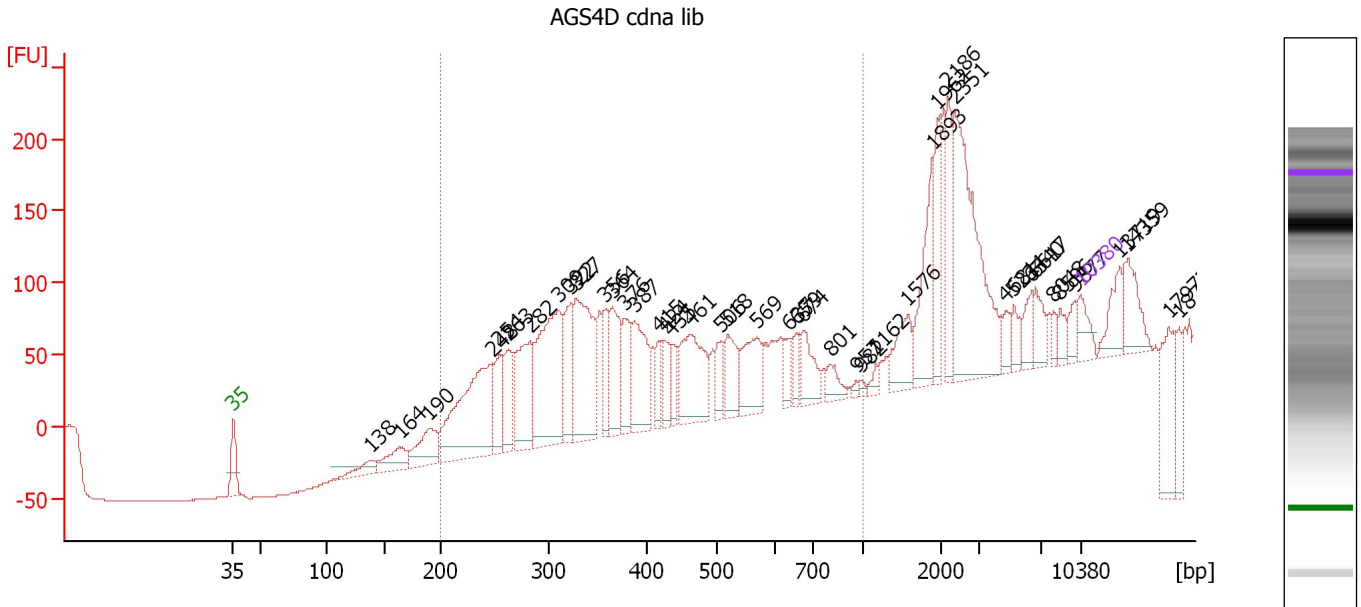
Peak table for Ladder

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\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : AGS4D cdna lib

Number of peaks found: 44 Corr. Area 1: 1,146.3
 Noise: 0.2

Peak table for sample 1 : AGS4D cdna lib

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	138	103.62	1,139.9	
3	164	173.87	1,603.6	
4	190	244.73	1,946.8	
5	248	934.77	5,722.1	
6	254	233.37	1,390.4	
7	263	239.20	1,377.8	
8	282	456.59	2,453.6	
9	309	793.32	3,887.6	
10	322	280.17	1,319.3	
11	327	646.48	2,993.3	
12	356	174.84	744.8	
13	364	291.42	1,211.6	
14	376	206.82	832.4	
15	387	344.41	1,347.0	
16	415	105.57	385.2	
17	424	119.82	427.8	
18	435	99.01	344.8	
19	461	409.42	1,345.9	
20	506	112.18	335.7	
21	518	182.18	532.5	
22	569	250.37	666.8	
23	637	80.14	190.7	
24	659	77.93	179.3	
25	674	164.71	370.5	
26	801	76.86	145.4	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad


Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...

... Peak table for sample 1 : AGS4D cdna lib

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	957	13.85	21.9	
28	982	12.84	19.8	
29	1,162	27.22	35.5	
30	1,576	154.14	148.2	
31	1,893	276.49	221.2	
32	1,962	179.39	138.5	
33	2,186	219.68	152.3	
34	2,351	745.32	480.4	
35	4,681	59.44	19.2	
36	5,234	62.67	18.1	
37	6,340	70.62	16.9	
38	6,617	90.38	20.7	
39	8,048	34.00	6.4	
40	8,596	44.54	7.9	
41	9,877	49.95	7.7	
42	10,380	75.00	10.9	Upper Marker
43	13,719	0.00	0.0	
44	14,359	0.00	0.0	
45	17,972	0.00	0.0	
46	18,795	0.00	0.0	

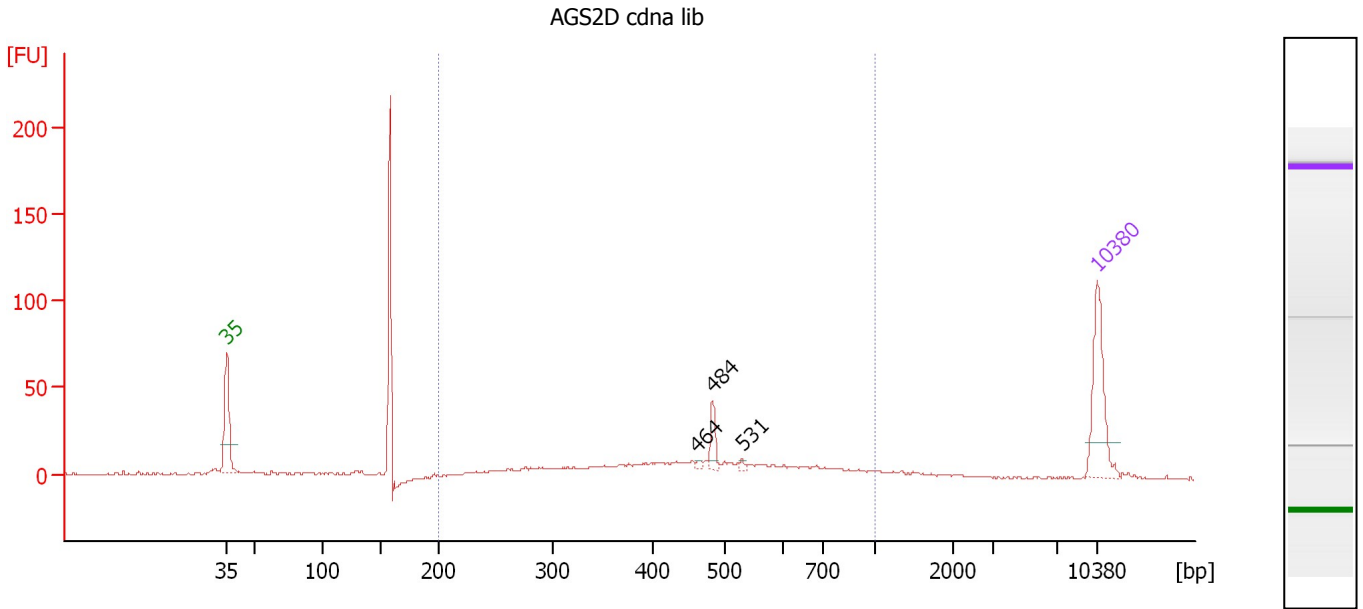
Region table for sample 1 : AGS4D cdna lib

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	392	12,488.4	2,907.03	1,146.3	52	28.9	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : AGS2D cdna lib

Number of peaks found: 3 Corr. Area 1: 269.8
 Noise: 0.5

Peak table for sample 2 : AGS2D cdna lib

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	464	3.58	11.7	
3	484	23.55	73.8	
4	531	3.40	9.7	
5	10,380	75.00	10.9	Upper Marker

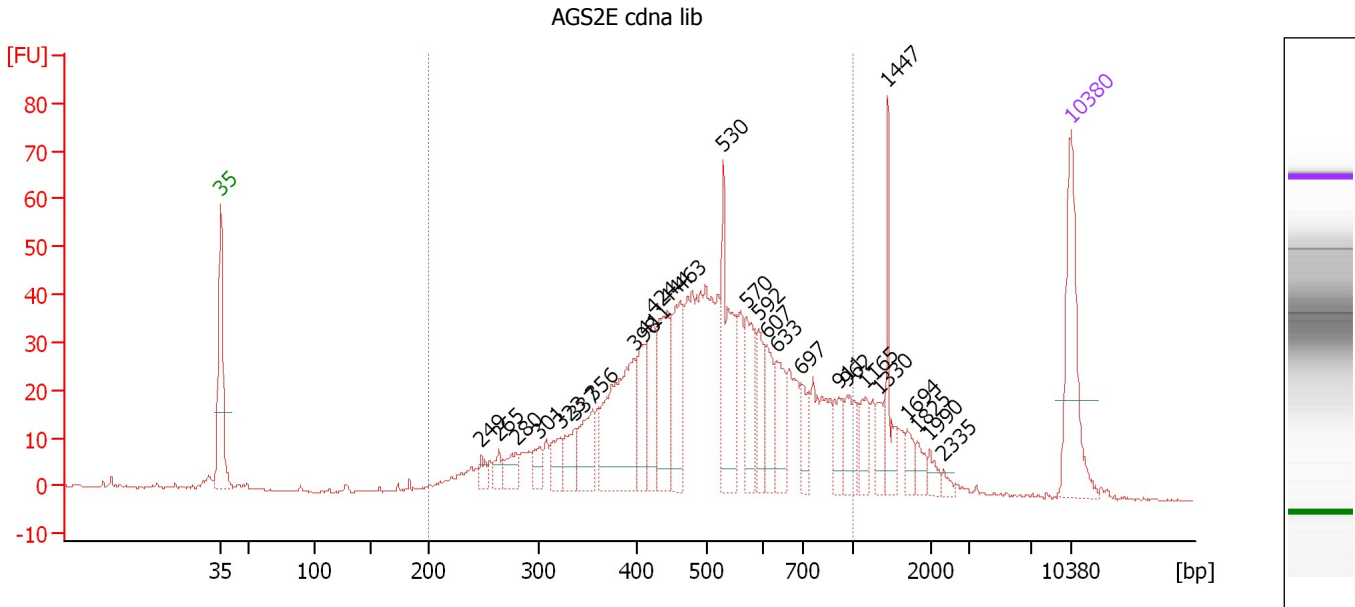
Region table for sample 2 : AGS2D cdna lib

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	481	1,053.3	286.02	269.8	63	33.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : AGS2E cdna lib

Number of peaks found: 27 Corr. Area 1: 941.6
 Noise: 0.2

Peak table for sample 3 : AGS2E cdna lib

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	249	12.07	73.6	
3	265	13.64	78.1	
4	280	22.92	124.0	
5	301	18.08	90.9	
6	323	24.27	113.8	
7	337	28.90	130.0	
8	356	45.79	194.9	
9	398	133.75	509.4	
10	411	42.82	157.7	
11	424	52.72	188.6	
12	444	76.94	262.5	
13	463	74.47	243.7	
14	530	95.18	272.1	
15	570	52.26	138.9	
16	592	31.60	80.9	
17	607	33.77	84.3	
18	633	37.74	90.4	
19	697	26.75	58.2	
20	911	23.64	39.3	
21	962	26.63	41.9	
22	1,165	20.67	26.9	
23	1,330	16.49	18.8	
24	1,447	34.79	36.4	
25	1,694	12.43	11.1	
26	1,825	9.44	7.8	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...

... Peak table for sample 3 : AGS2E cdna lib

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	1,990	8.49	6.5	
28	2,335	4.20	2.7	
29	10,380	75.00	10.9	Upper Marker

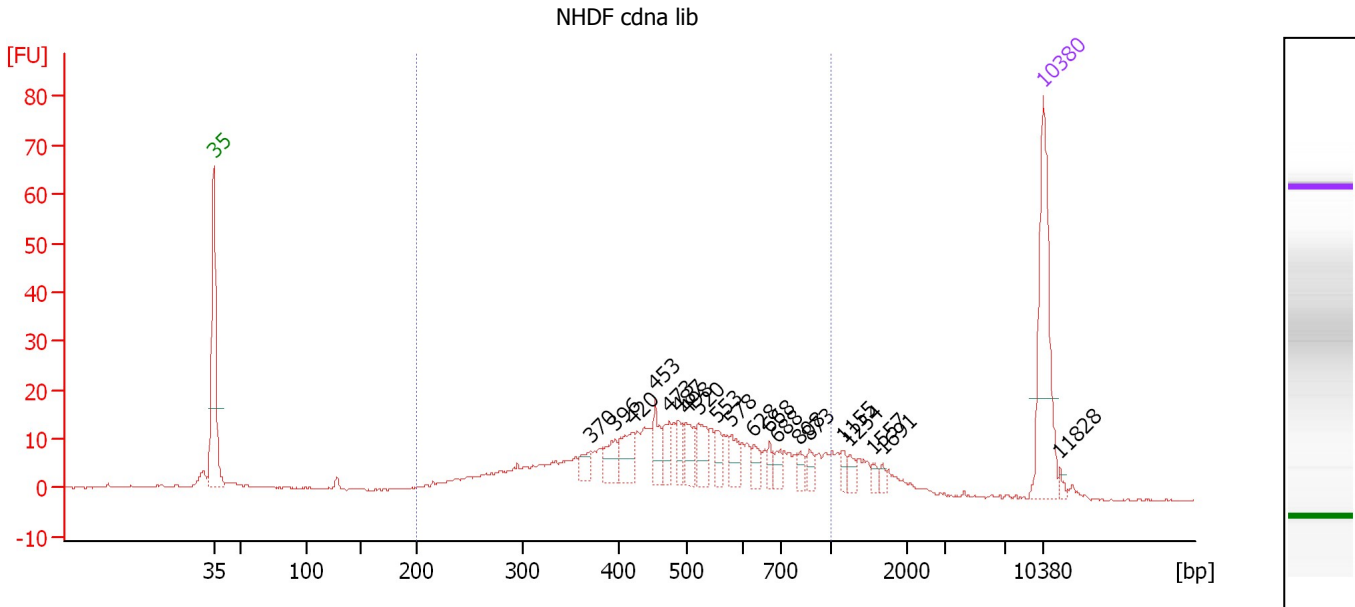
Region table for sample 3 : AGS2E cdna lib

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	515	4,575.1	1,356.57	941.6	86	31.0	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : NHDF cdna lib

Number of peaks found: 20 Corr. Area 1: 376.5
 Noise: 0.2

Peak table for sample 4 : NHDF cdna lib

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	370	12.42	50.8	
3	396	20.50	78.4	
4	420	24.62	88.9	
5	453	20.29	67.9	
6	473	16.08	51.5	
7	487	13.57	42.2	
8	498	19.83	60.3	
9	520	21.75	63.4	
10	553	13.17	36.1	
11	578	14.07	36.9	
12	628	11.37	27.4	
13	668	8.05	18.2	
14	688	8.95	19.7	
15	808	7.13	13.4	
16	873	8.06	14.0	
17	1,155	6.29	8.3	
18	1,254	6.66	8.1	
19	1,557	4.35	4.2	
20	1,691	3.44	3.1	
21	10,380	75.00	10.9	Upper Marker
22	11,828	0.00	0.0	

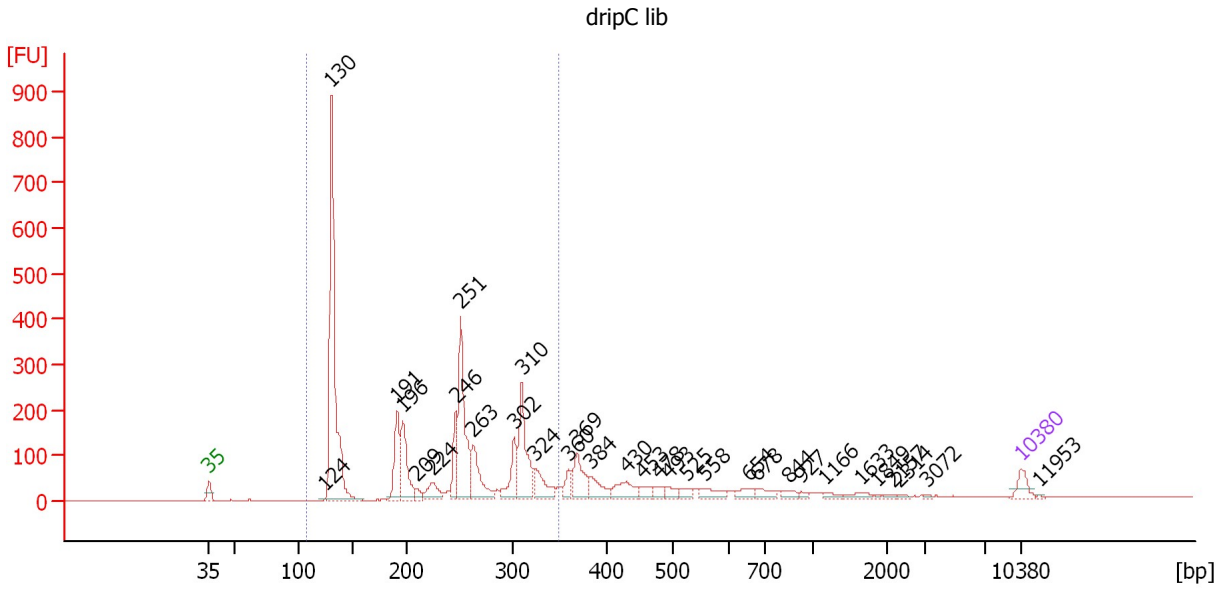
Region table for sample 4 : NHDF cdna lib

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	503	1,958.9	550.36	376.5	78	34.1	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : dripC lib

Number of peaks found: 32 Corr. Area 1: 2,547.9
 Noise: 0.2

Peak table for sample 5 : dripC lib

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	124	12.54	153.1	
3	130	2,328.14	27,130.7	
4	191	437.27	3,471.5	
5	196	435.99	3,365.1	
6	209	50.63	366.4	
7	224	167.14	1,130.4	
8	246	232.44	1,430.0	
9	251	897.20	5,408.9	
10	263	381.76	2,199.7	
11	302	288.54	1,447.1	
12	310	572.02	2,795.0	
13	324	228.70	1,068.4	
14	360	106.02	445.9	
15	369	269.04	1,103.6	
16	384	158.77	627.1	
17	430	183.61	646.4	
18	453	72.78	243.2	
19	478	67.83	215.0	
20	493	61.40	188.7	
21	525	57.51	166.1	
22	558	100.16	272.1	
23	654	72.41	167.8	
24	678	73.62	164.5	
25	844	47.80	85.8	
26	927	20.68	33.8	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad


Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...

... Peak table for sample 5 : dripC lib

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	1,166	31.79	41.3	
28	1,633	40.40	37.5	
29	1,849	6.22	5.1	
30	2,157	12.48	8.8	
31	2,314	9.92	6.5	
32	3,072	6.29	3.1	
33	10,380	75.00	10.9	Upper Marker
34	11,953	0.00	0.0	

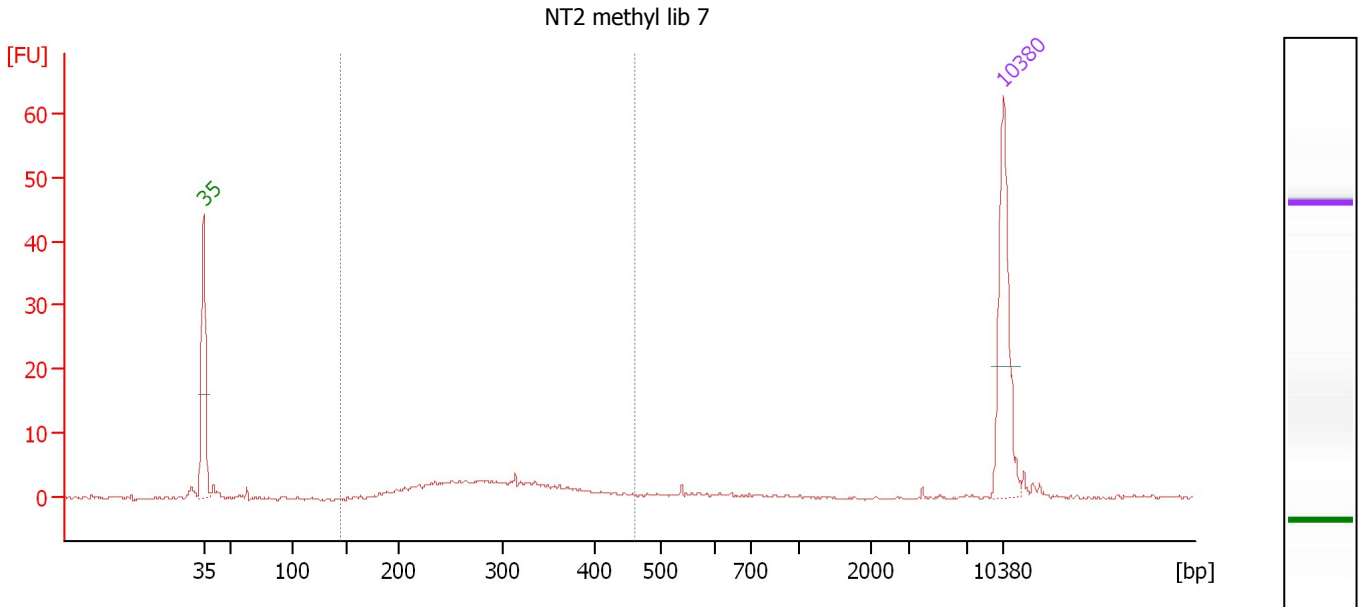
Region table for sample 5 : dripC lib

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
108	350	223	49,264.5	6,077.94	2,547.9	74	31.4	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : NT2 methyl lib 7

Number of peaks found: 0 Corr. Area 1: 56.4
 Noise: 0.2

Peak table for sample 6 : NT2 methyl lib 7

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	10,380	75.00	10.9	Upper Marker

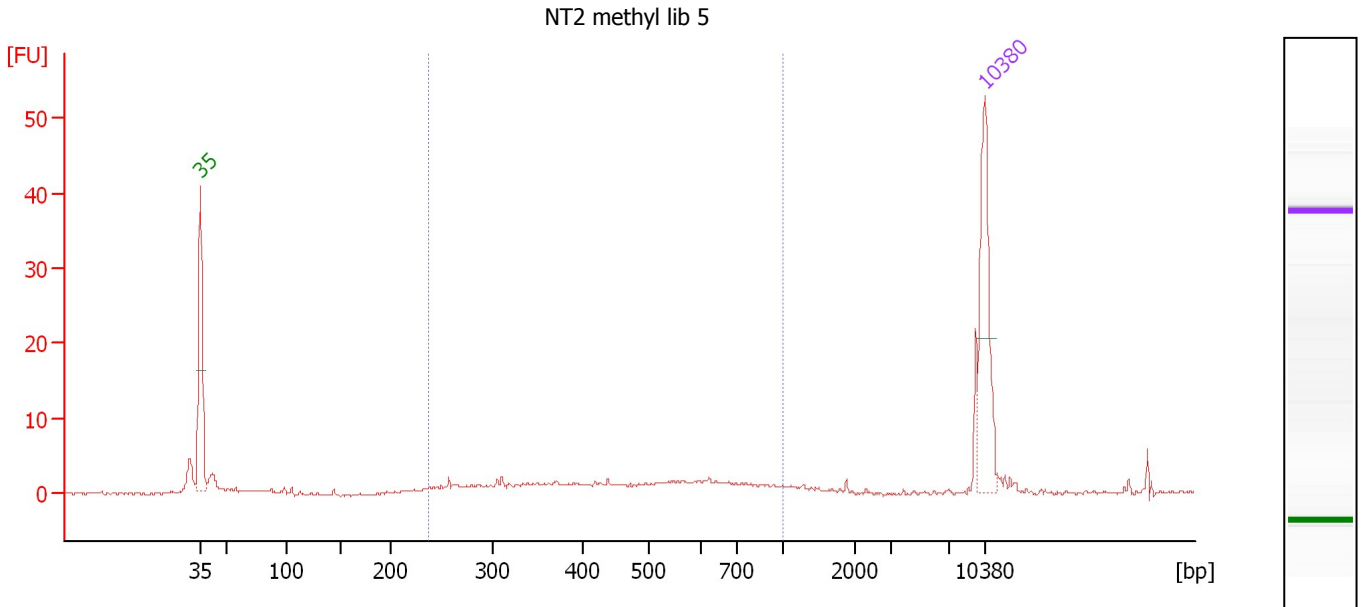
Region table for sample 6 : NT2 methyl lib 7

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
146	462	294	669.9	122.51	56.4	75	21.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : NT2 methyl lib 5

Number of peaks found: 0 Corr. Area 1: 45.2
 Noise: 0.2

Peak table for sample 7 : NT2 methyl lib 5

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	10,380	75.00	10.9	Upper Marker

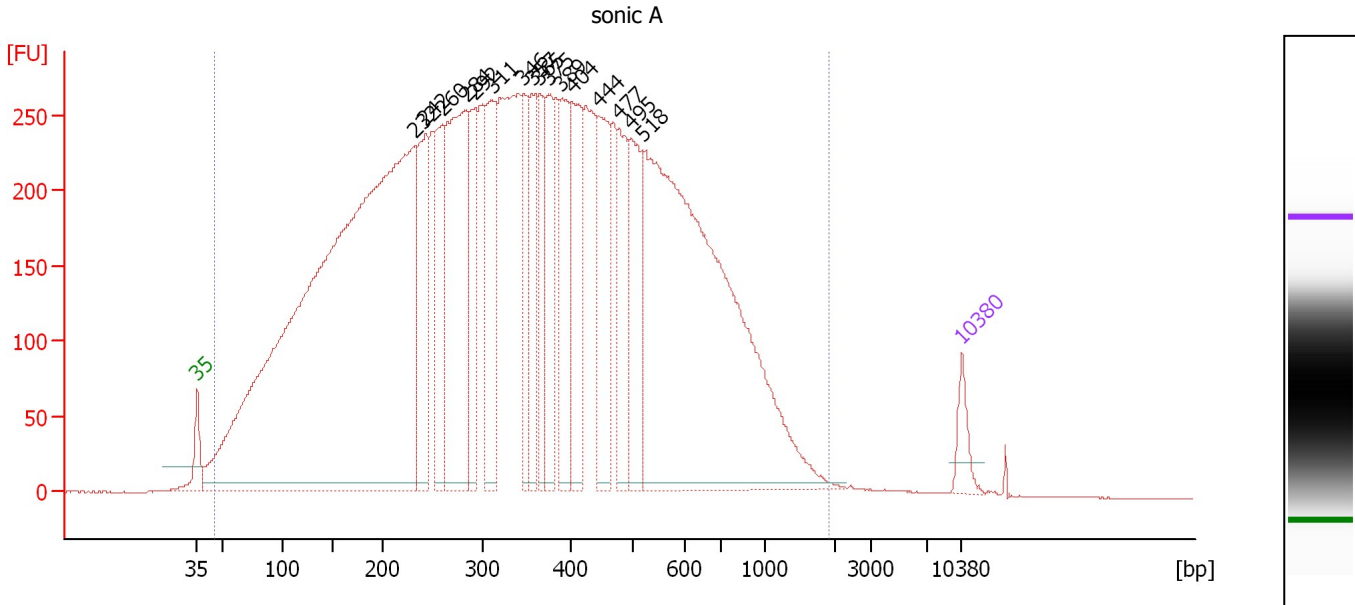
Region table for sample 7 : NT2 methyl lib 5

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
237	1,000	483	415.3	109.91	45.2	58	36.8	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : sonic A

Number of peaks found: 16 Corr. Area 1: 14,022.7
 Noise: 0.2

Peak table for sample 8 : sonic A

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	232	6,249.21	40,890.3	
3	242	565.55	3,536.4	
4	260	534.02	3,117.9	
5	284	1,166.55	6,230.3	
6	292	414.06	2,146.3	
7	311	579.99	2,828.3	
8	346	306.80	1,341.7	
9	357	369.12	1,567.6	
10	365	294.07	1,221.9	
11	375	374.77	1,514.4	
12	389	423.69	1,650.7	
13	404	489.06	1,834.4	
14	444	455.92	1,554.2	
15	477	372.59	1,184.7	
16	495	426.04	1,303.8	
17	518	3,006.48	8,797.1	
18	10,380	75.00	10.9	Upper Marker

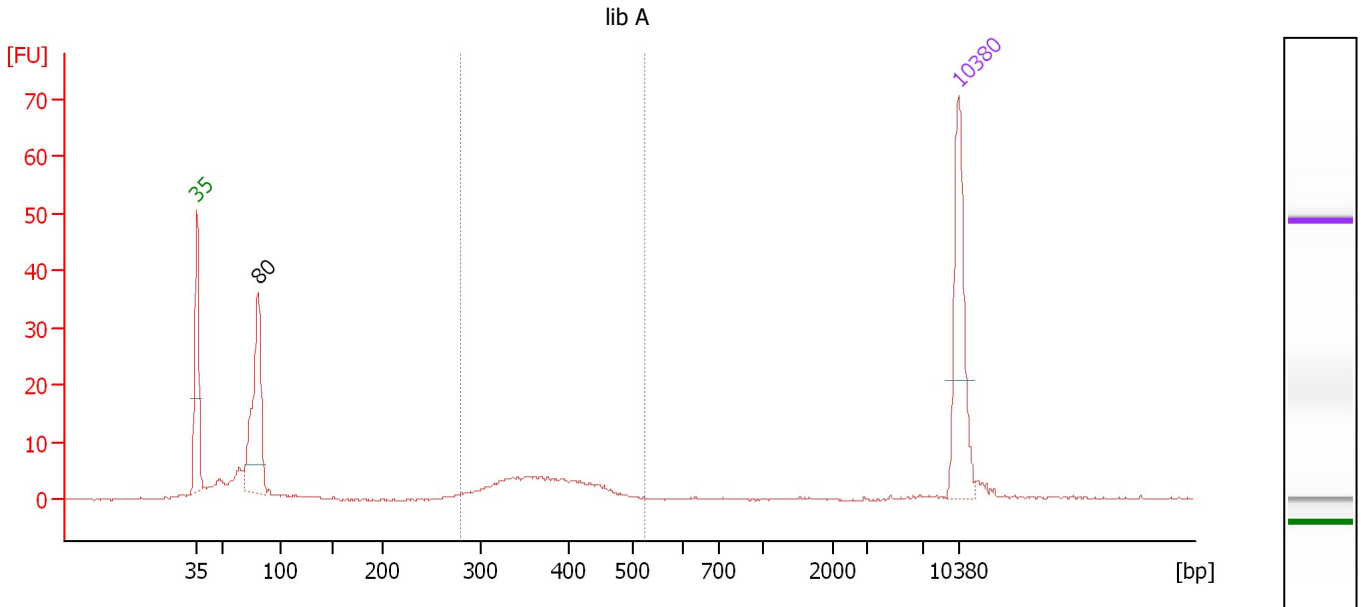
Region table for sample 8 : sonic A

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
45	1,902	384	138,524.2	20,255.45	14,022.7	99	60.1	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : lib A

Number of peaks found: 1 Corr. Area 1: 50.1
 Noise: 0.1

Peak table for sample 9 : lib A

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	80	150.89	2,848.9	
3	10,380	75.00	10.9	Upper Marker

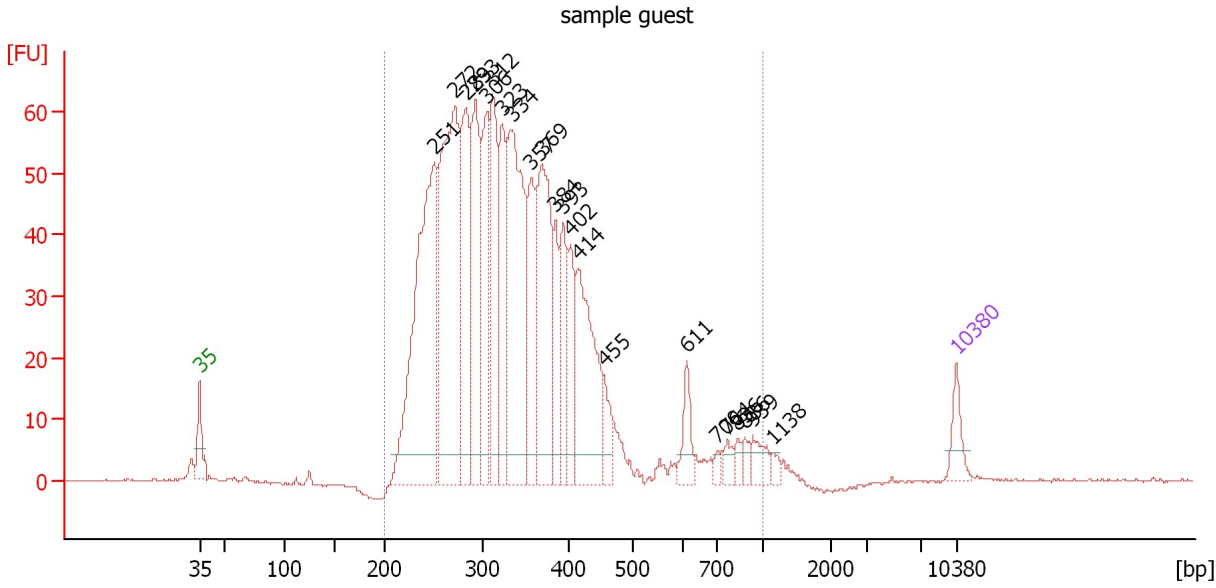
Region table for sample 9 : lib A

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
278	524	373	345.3	83.35	50.1	32	13.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : sample guest

Number of peaks found: 22 Corr. Area 1: 1,268.5
 Noise: 0.1

Peak table for sample 10 : sample guest


Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	251	1,513.73	9,128.2	
3	272	1,413.64	7,877.5	
4	283	616.70	3,298.2	
5	293	603.72	3,121.1	
6	306	512.48	2,540.2	
7	312	545.84	2,647.2	
8	323	417.02	1,957.2	
9	334	1,022.74	4,633.1	
10	357	377.57	1,604.6	
11	369	711.79	2,924.6	
12	384	241.75	953.7	
13	393	269.40	1,037.9	
14	402	228.31	861.1	
15	414	583.58	2,137.6	
16	455	116.97	389.1	
17	611	115.67	286.8	
18	700	27.20	58.8	
19	764	48.28	95.7	
20	836	29.49	53.5	
21	886	31.48	53.9	
22	939	70.47	113.7	
23	1,138	24.22	32.2	
24	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...

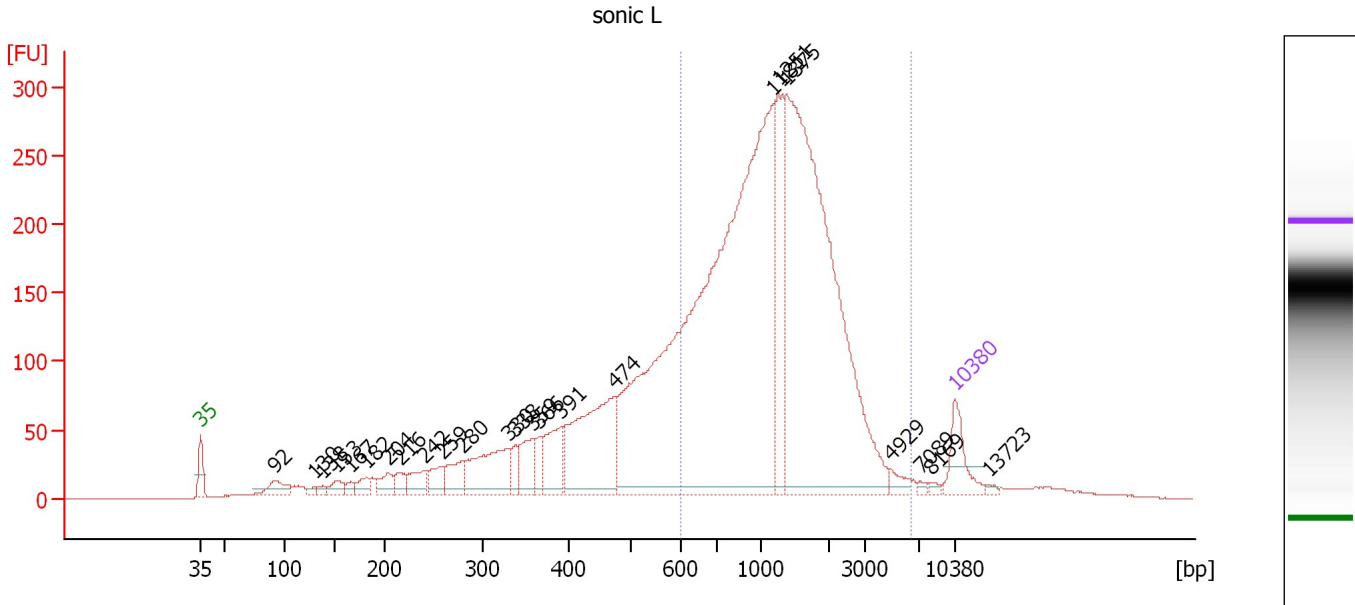
... Region table for sample 10 : sample guest

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/ μ l]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	346	46,661.3	9,702.20	1,268.5	99	33.1	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
 Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : sonic L

Number of peaks found: 24 Corr. Area 1: 3,664.6
 Noise: 0.3

Peak table for sample 11 : sonic L

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	92	75.21	1,237.5	
3	130	14.83	172.8	
4	138	17.28	189.8	
5	153	42.30	419.5	
6	167	20.64	187.6	
7	182	40.51	337.8	
8	204	52.91	393.5	
9	216	37.94	266.5	
10	242	67.21	421.0	
11	259	58.35	340.9	
12	280	81.86	442.8	
13	330	217.94	999.3	
14	338	38.89	174.1	
15	359	96.83	408.3	
16	366	42.97	178.1	
17	391	130.77	506.3	
18	474	409.58	1,310.3	
19	1,181	2,192.36	2,812.5	
20	1,251	260.14	315.1	
21	1,375	1,412.50	1,556.4	
22	4,929	21.87	6.7	
23	7,089	6.60	1.4	
24	8,169	7.42	1.4	
25	10,380	75.00	10.9	Upper Marker
26	13,723	0.00	0.0	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
Modified: 5/3/2013 2:59:03 PM

Electropherogram Summary Continued ...

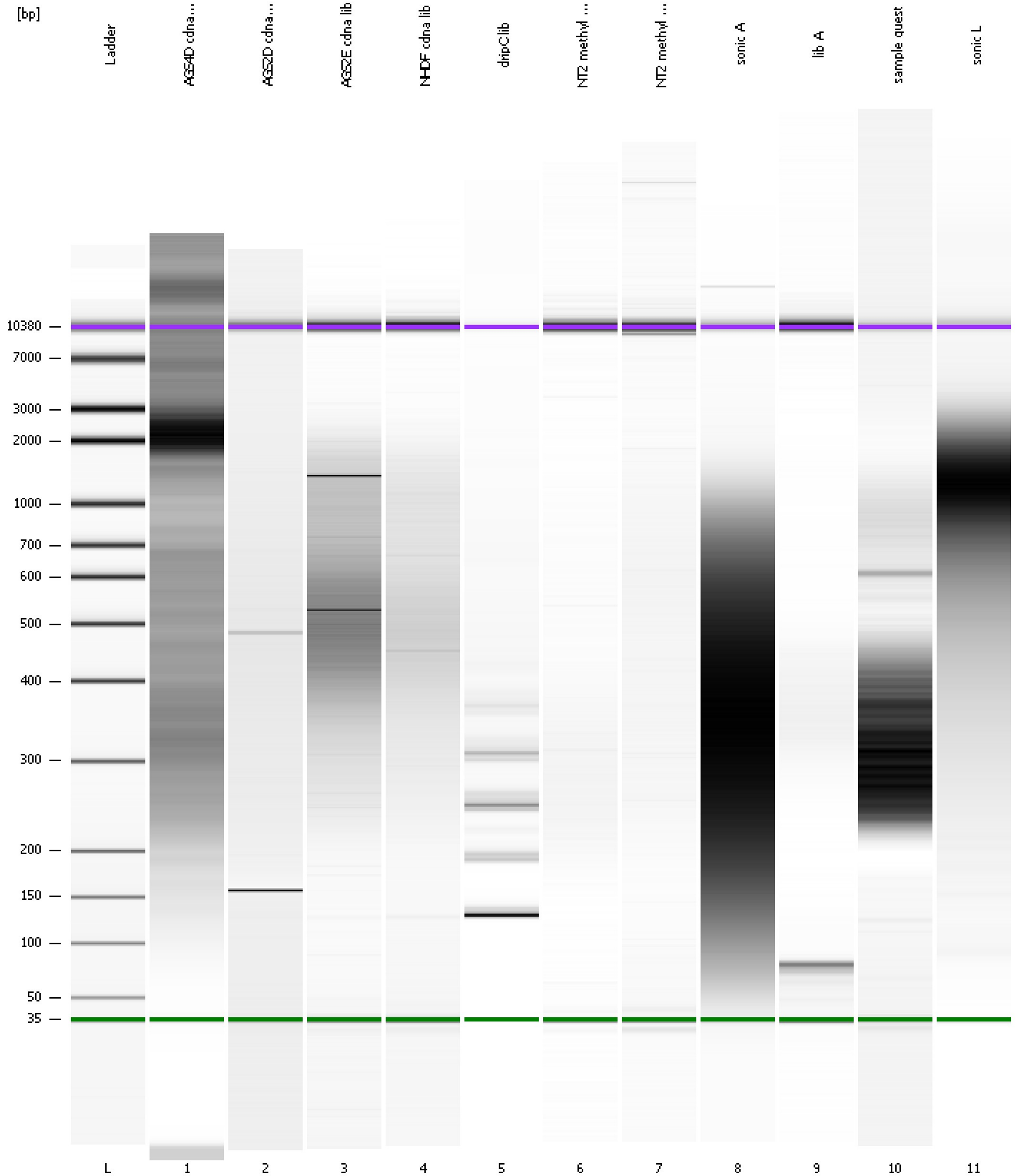
... Region table for sample 11 : sonic L

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
600	6,373	1,400	4,874.7	3,510.31	3,664.6	63	55.6	■

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad

Created: 5/3/2013 2:13:15 PM
Modified: 5/3/2013 2:59:03 PM

Gel Image



Assay Class: High Sensitivity DNA Assay Created: 5/3/2013 2:13:15 PM
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad Modified: 5/3/2013 2:59:03 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		5/3/2013 2:54:36 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2013-05-03\2013-05-03_002.xad)		Instrument	Run		5/3/2013 2:13:21 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		5/3/2013 2:13:21 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		5/3/2013 2:13:21 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		5/3/2013 2:13:21 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		5/3/2013 2:13:21 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		5/3/2013 2:13:21 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		5/3/2013 2:13:21 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1