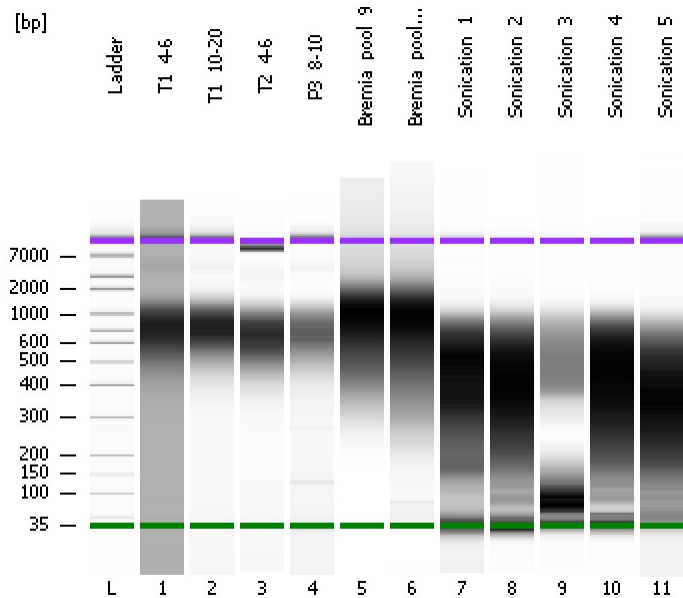


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
Modified: 7/8/2013 12:16:55 PM

Electrophoresis File Run Summary



Instrument Information:

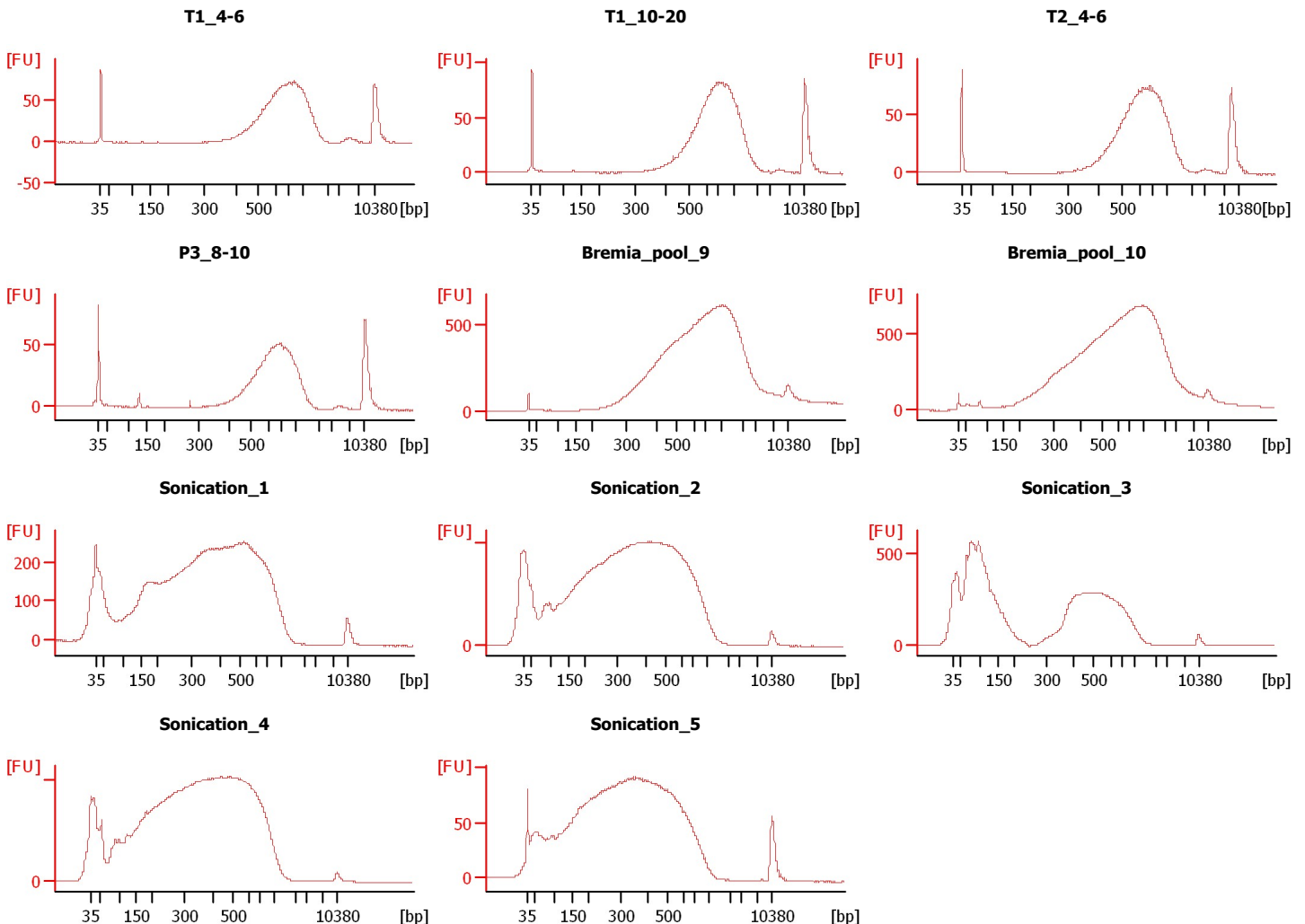
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
T1_4-6		<input type="checkbox"/>	✓			
T1_10-20		<input type="checkbox"/>	✓			
T2_4-6		<input type="checkbox"/>	✓			
P3_8-10		<input type="checkbox"/>	✓			
Bremia_pool_9		<input type="checkbox"/>	✓			
Bremia_pool_10		<input type="checkbox"/>	✓			
Sonication_1		<input type="checkbox"/>	✓			
Sonication_2		<input type="checkbox"/>	✓			
Sonication_3		<input type="checkbox"/>	✓			
Sonication_4		<input type="checkbox"/>	✓			
Sonication_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\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
Modified: 7/8/2013 12:16:55 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

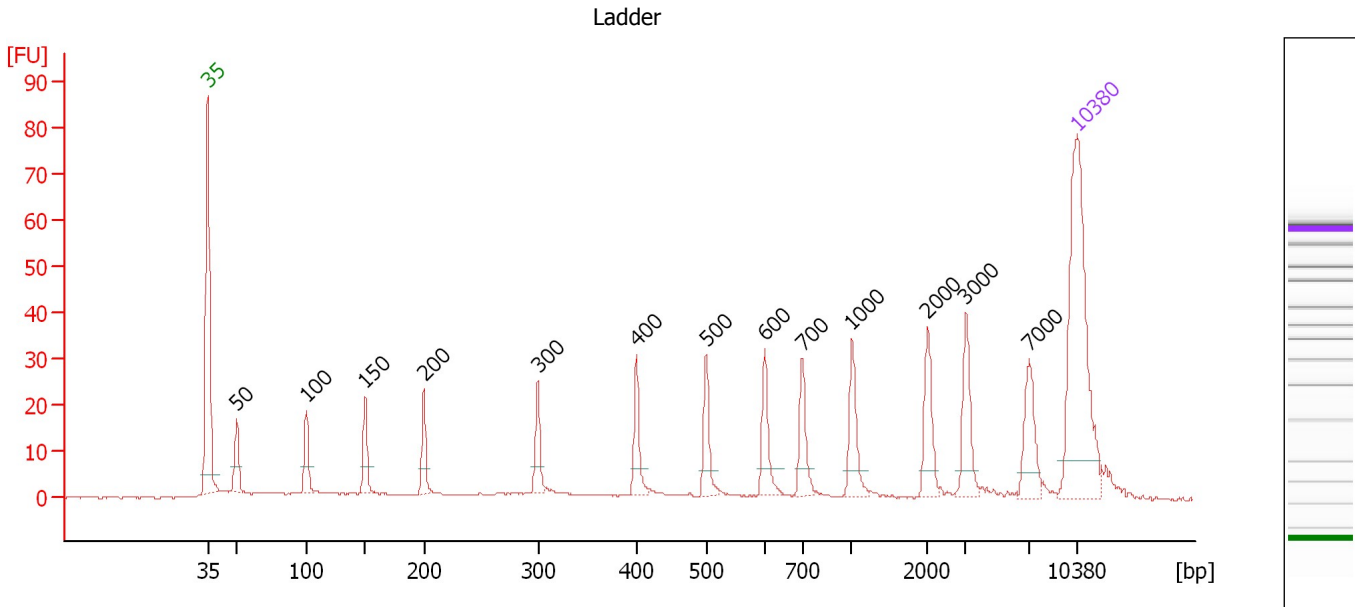
Ladder

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\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

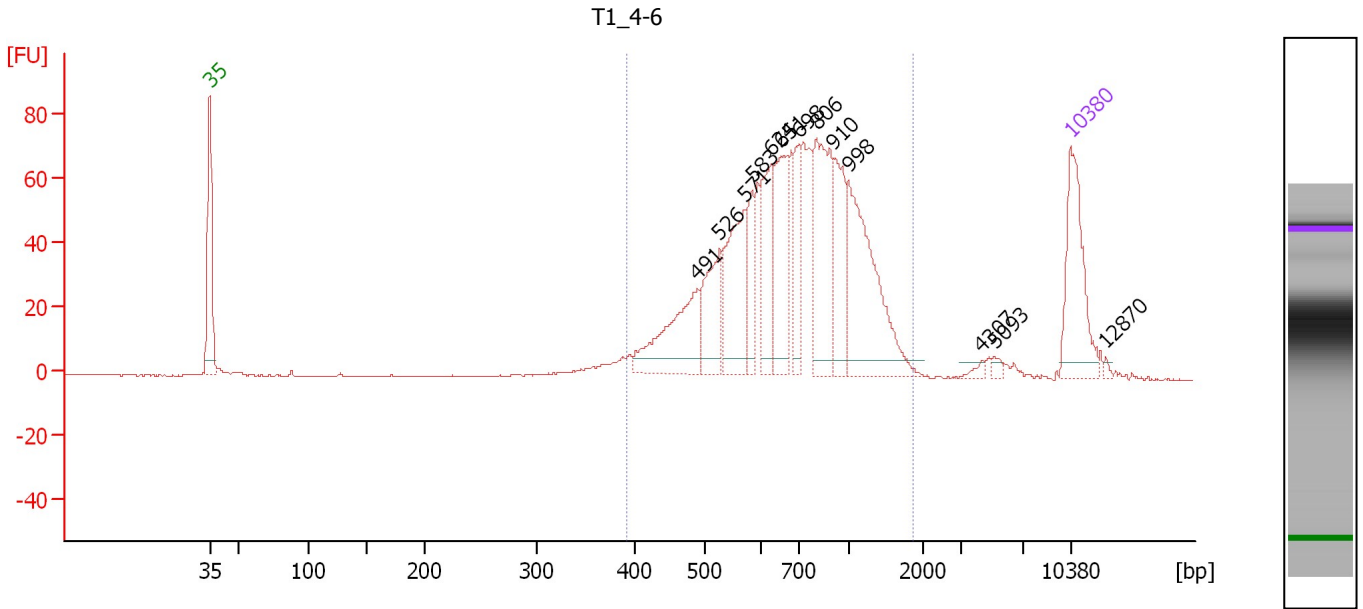
Peak table for Ladder

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\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : T1_4-6

Number of peaks found: 13 Corr. Area 1: 1,109.2
 Noise: 0.2

Peak table for sample 1 : T1_4-6

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	491	105.63	326.1	
3	526	69.36	199.8	
4	571	100.72	267.1	
5	583	39.75	103.2	
6	624	74.59	181.1	
7	651	86.70	201.8	
8	698	44.16	95.9	
9	806	123.87	232.9	
10	910	68.55	114.1	
11	998	149.44	226.9	
12	4,307	3.62	1.3	
13	5,093	4.24	1.3	
14	10,380	75.00	10.9	Upper Marker
15	12,870	0.00	0.0	

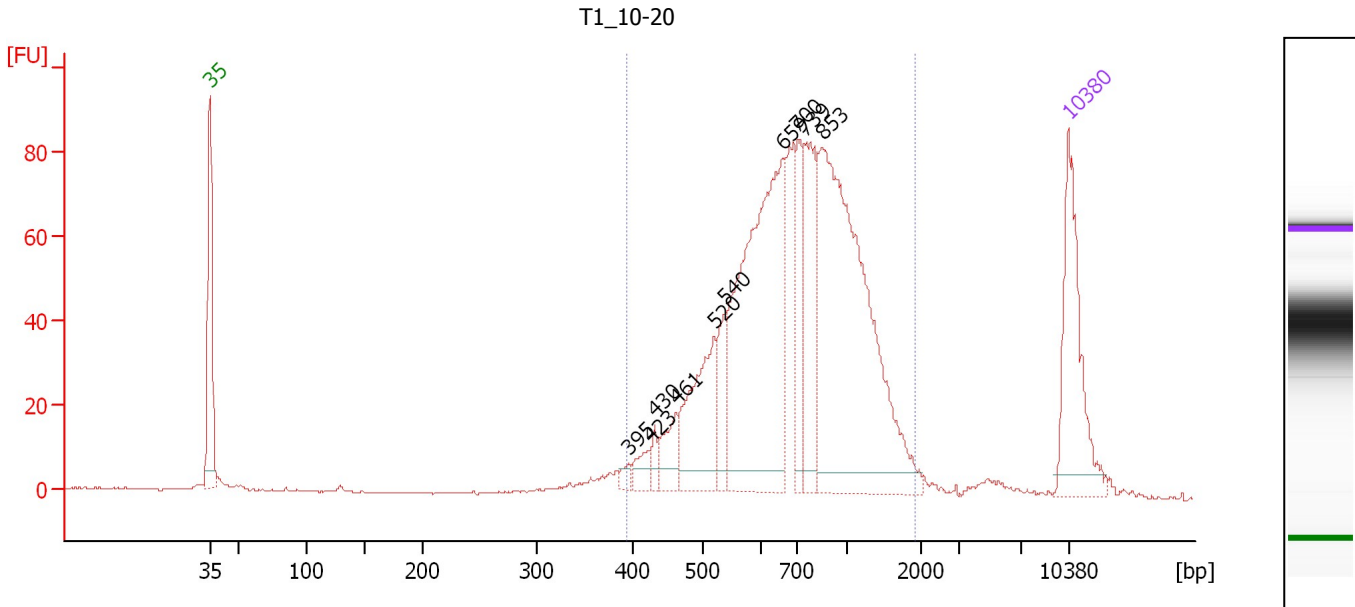
Region table for sample 1 : T1_4-6

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
393	1,859	766	2,328.0	1,032.31	1,109.2	93	35.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : T1_10-20

Number of peaks found: 10 Corr. Area 1: 1,236.6
 Noise: 0.2

Peak table for sample 2 : T1_10-20

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	395	6.25	23.9	
3	423	14.17	50.7	
4	430	8.51	30.0	
5	461	26.31	86.4	
6	520	86.80	252.7	
7	540	33.11	92.9	
8	659	277.99	639.5	
9	700	50.24	108.7	
10	739	84.05	172.3	
11	853	319.69	567.6	
12	10,380	75.00	10.9	Upper Marker

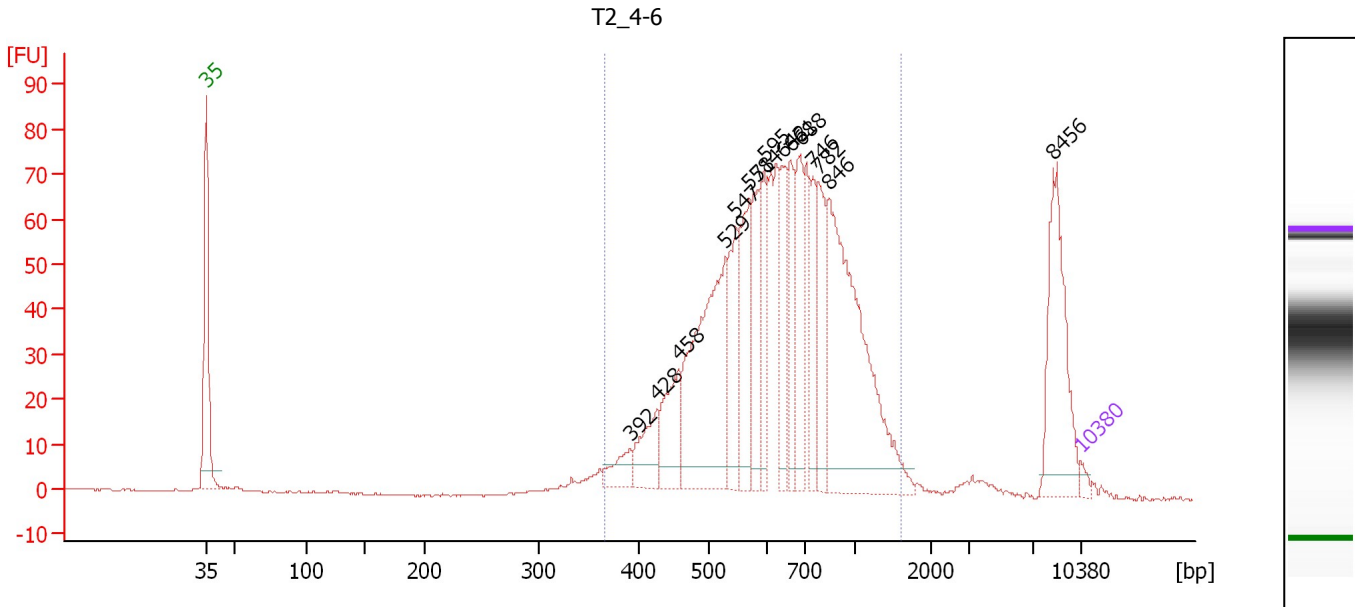
Region table for sample 2 : T1_10-20

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
394	1,915	792	2,116.9	961.05	1,236.6	94	37.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : T2_4-6

Number of peaks found: 15 Corr. Area 1: 1,183.3
 Noise: 0.2

Peak table for sample 3 : T2_4-6

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	392	454.55	1,755.8	
3	428	831.28	2,944.7	
4	458	1,173.09	3,880.3	
5	529	3,972.60	11,380.9	
6	547	1,266.87	3,508.5	
7	571	1,629.48	4,325.4	
8	584	1,219.23	3,160.7	
9	595	1,036.24	2,637.0	
10	642	1,076.98	2,542.1	
11	661	1,007.72	2,308.2	
12	688	1,333.47	2,936.3	
13	746	937.86	1,903.9	
14	782	1,296.19	2,512.7	
15	846	4,920.80	8,816.1	
16	8,456	1,872.12	335.4	
17	10,380	75.00	10.9	Upper Marker

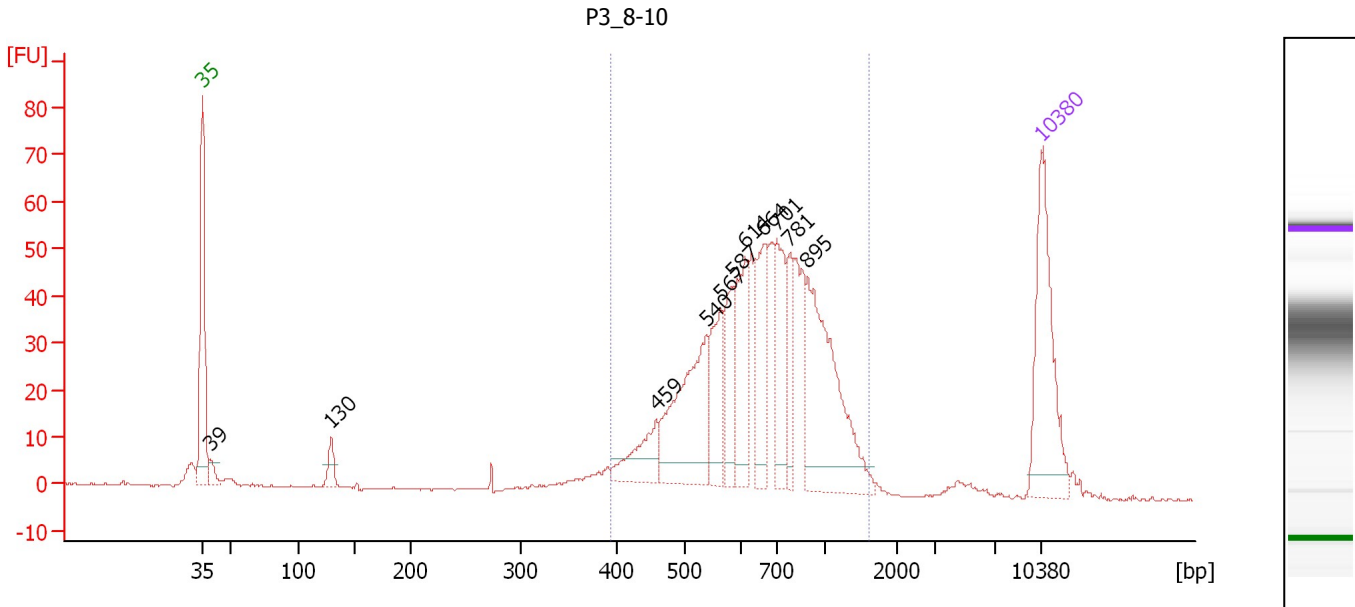
Region table for sample 3 : T2_4-6

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
367	1,606	689	62,009.1	25,240.89	1,183.3	88	32.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : P3 8-10

Number of peaks found: 11 Corr. Area 1: 753.7
 Noise: 0.2

Peak table for sample 4 : P3 8-10

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	39	11.66	449.3	
3	130	15.87	185.2	
4	459	36.81	121.5	
5	540	107.94	302.8	
6	567	43.67	116.8	
7	587	37.86	97.8	
8	614	57.10	140.8	
9	664	56.02	127.8	
10	701	56.80	122.7	
11	781	29.68	57.6	
12	895	128.26	217.2	
13	10,380	75.00	10.9	Upper Marker

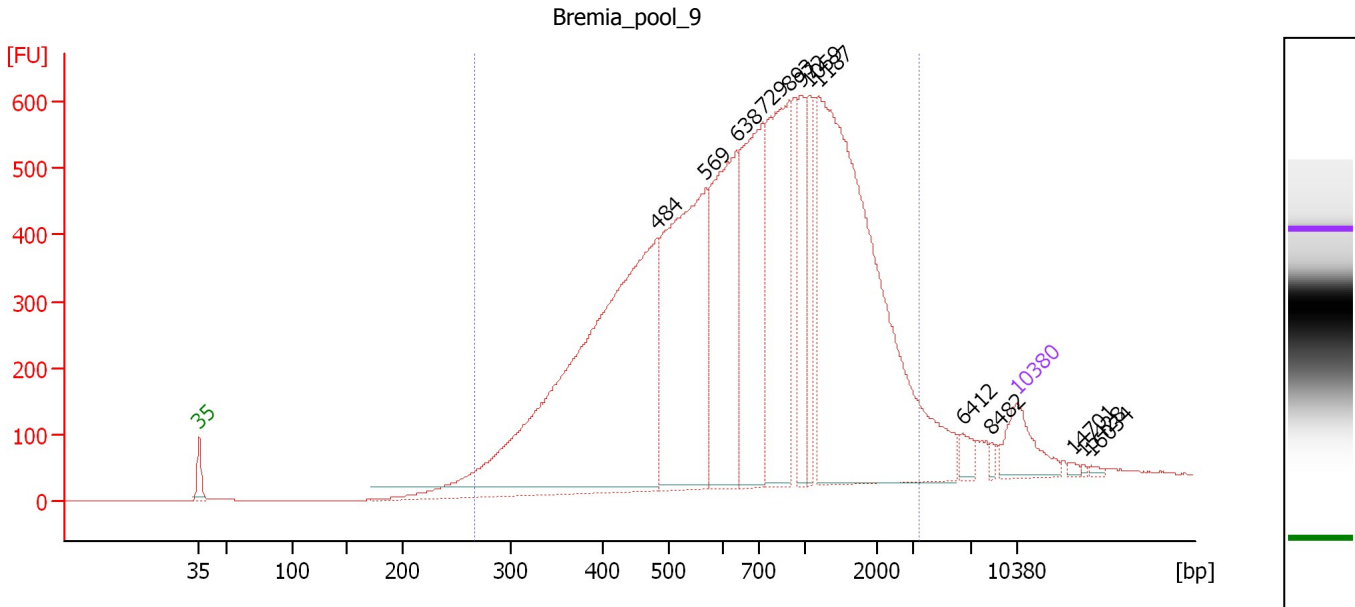
Region table for sample 4 : P3 8-10

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
395	1,603	718	1,636.9	699.48	753.7	90	31.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : Bremia_pool_9

Number of peaks found: 13 Corr. Area 1: 14,986.1
 Noise: 0.2

Peak table for sample 5 : Bremia_pool_9

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	484	1,413.92	4,430.8	
3	569	682.74	1,818.2	
4	638	469.25	1,114.9	
5	729	428.66	891.2	
6	893	433.69	735.5	
7	972	137.77	214.7	
8	1,059	108.88	155.7	
9	1,187	1,097.85	1,401.7	
10	6,412	22.86	5.4	
11	8,482	7.70	1.4	
12	10,380	75.00	10.9	Upper Marker
13	14,701	0.00	0.0	
14	15,428	0.00	0.0	
15	16,034	0.00	0.0	

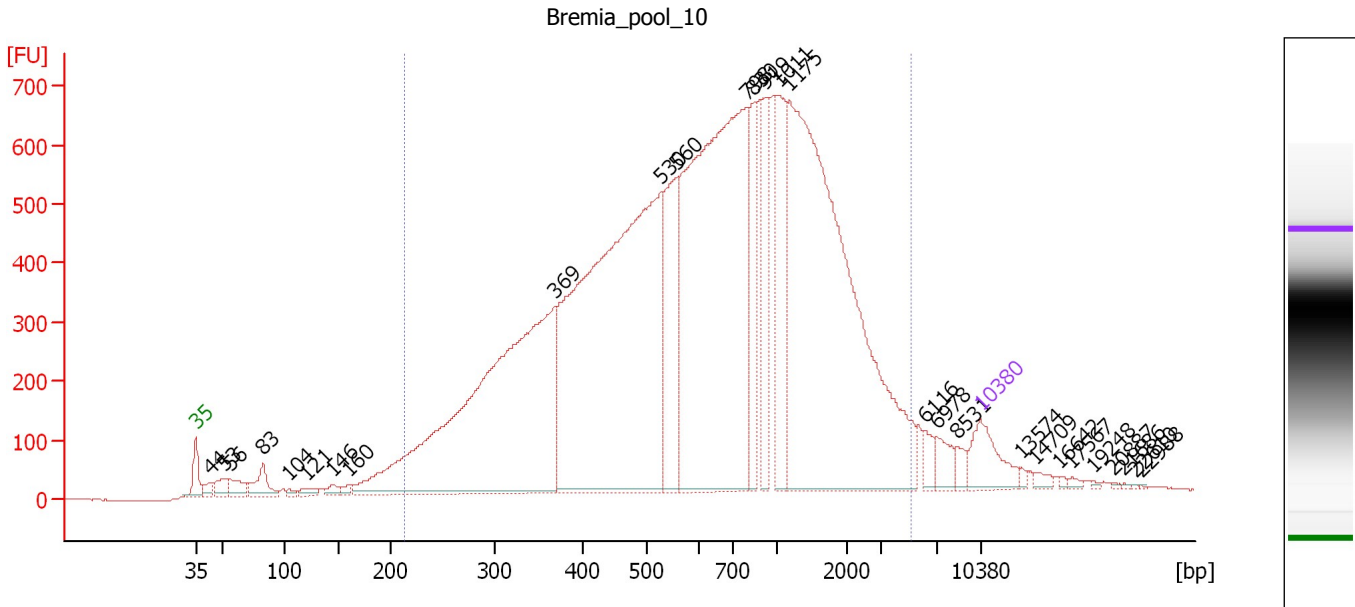
Region table for sample 5 : Bremia_pool_9

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/µl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
267	3,395	907	12,061.4	4,837.49	14,986.1	96	63.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : Bremia_pool_10

Number of peaks found: 28 Corr. Area 1: 20,544.2
 Noise: 1.5

Peak table for sample 6 : Bremia_pool_10

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	44	27.08	941.0	
3	53	39.69	1,142.8	
4	56	46.84	1,257.7	
5	83	69.10	1,260.8	
6	104	9.27	135.2	
7	121	18.43	231.7	
8	146	19.28	200.5	
9	160	12.25	116.2	
10	369	1,328.64	5,452.8	
11	530	1,595.45	4,564.8	
12	560	270.45	731.7	
13	798	1,309.11	2,485.5	
14	830	149.81	273.4	
15	919	144.51	238.1	
16	1,011	216.67	324.7	
17	1,175	1,279.96	1,650.8	
18	6,116	24.78	6.1	
19	6,978	36.21	7.9	
20	8,531	18.01	3.2	
21	10,380	75.00	10.9	Upper Marker
22	13,574	0.00	0.0	
23	14,709	0.00	0.0	
24	16,642	0.00	0.0	
25	17,567	0.00	0.0	
26	19,248	0.00	0.0	


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...**... Peak table for sample 6 : Bremia_pool_10**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	20,887	0.00	0.0	
28	21,686	0.00	0.0	
29	22,610	0.00	0.0	
30	22,988	0.00	0.0	

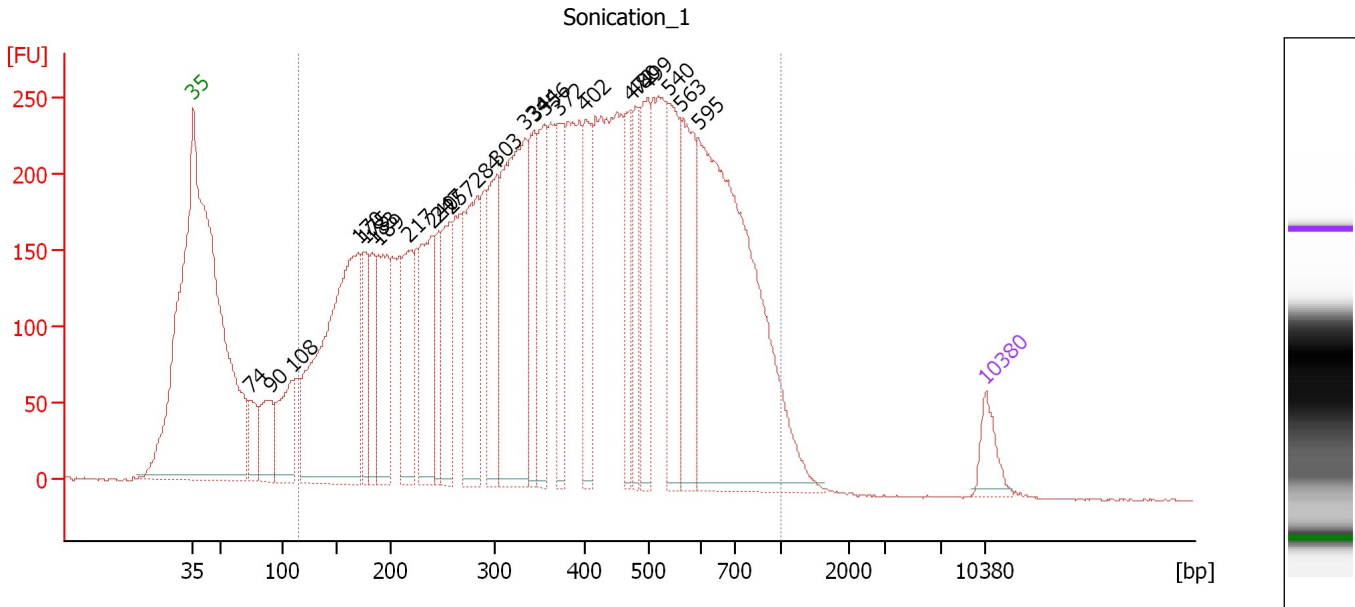
Region table for sample 6 : Bremia_pool_10

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
214	5,131	908	19,399.3	6,762.02	20,544.2	93	81.7	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : Sonication_1

Number of peaks found: 24 Corr. Area 1: 11,254.6
 Noise: 0.5

Peak table for sample 7 : Sonication_1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	74	167.00	3,430.0	
3	90	235.42	3,977.4	
4	108	312.75	4,401.6	
5	170	1,545.30	13,774.1	
6	175	234.77	2,027.2	
7	183	227.74	1,886.1	
8	189	461.01	3,697.1	
9	217	367.99	2,571.6	
10	240	458.55	2,896.3	
11	247	209.50	1,287.5	
12	257	322.64	1,899.9	
13	284	538.94	2,875.6	
14	303	390.57	1,954.3	
15	334	904.77	4,101.6	
16	344	283.99	1,249.7	
17	356	335.05	1,427.7	
18	372	275.42	1,123.1	
19	402	292.27	1,102.3	
20	472	222.03	712.6	
21	480	207.28	653.7	
22	499	303.38	921.4	
23	540	384.22	1,078.3	
24	563	393.47	1,058.8	
25	595	1,572.14	4,001.7	
26	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...

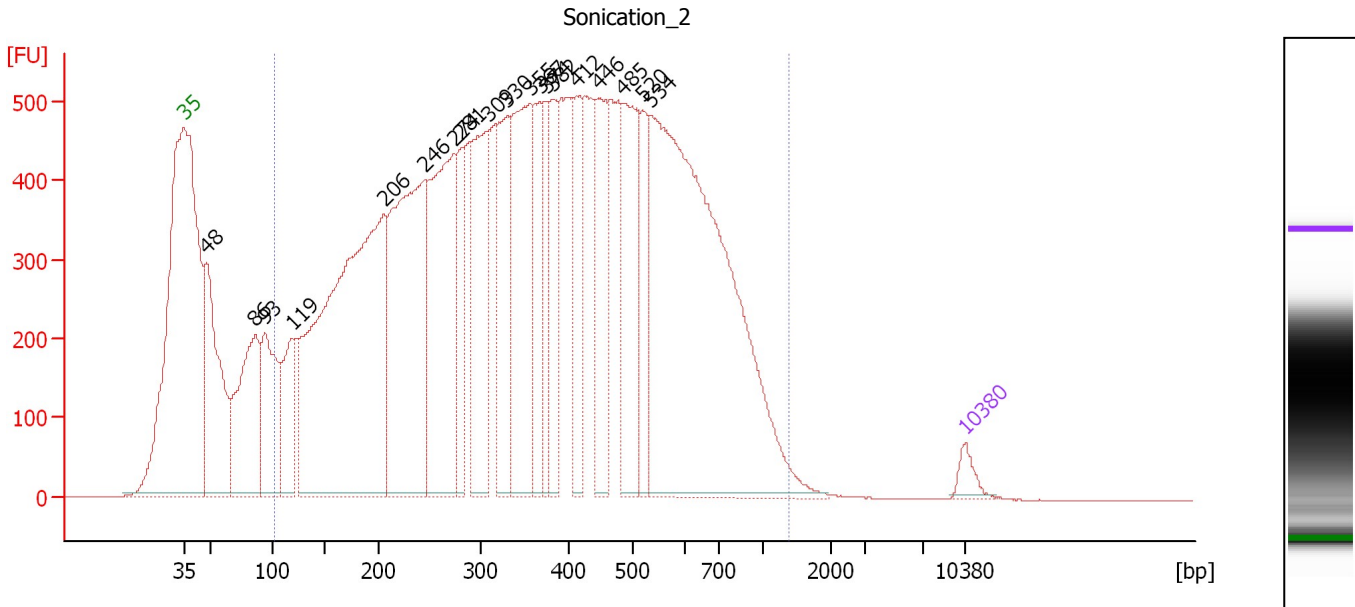
... Region table for sample 7 : Sonication 1

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
114	1,000	409	70,258.0	13,439.96	11,254.6	95	44.6	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : Sonication_2

Number of peaks found: 19 Corr. Area 1: 24,449.7
 Noise: 0.2

Peak table for sample 8 : Sonication_2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	48	1,619.49	50,822.0	
3	86	1,410.35	24,772.8	
4	93	998.03	16,177.2	
5	119	688.89	8,807.3	
6	206	4,789.79	35,298.3	
7	246	2,592.39	15,992.4	
8	274	1,990.66	11,000.4	
9	281	525.35	2,828.0	
10	309	1,241.18	6,081.9	
11	330	907.35	4,162.9	
12	355	1,476.69	6,293.9	
13	367	625.80	2,584.2	
14	374	446.52	1,808.4	
15	382	542.30	2,151.3	
16	412	585.51	2,154.1	
17	446	818.49	2,781.9	
18	485	990.44	3,095.2	
19	520	482.47	1,407.1	
20	534	4,615.70	13,101.3	
21	10,380	75.00	10.9	Upper Marker

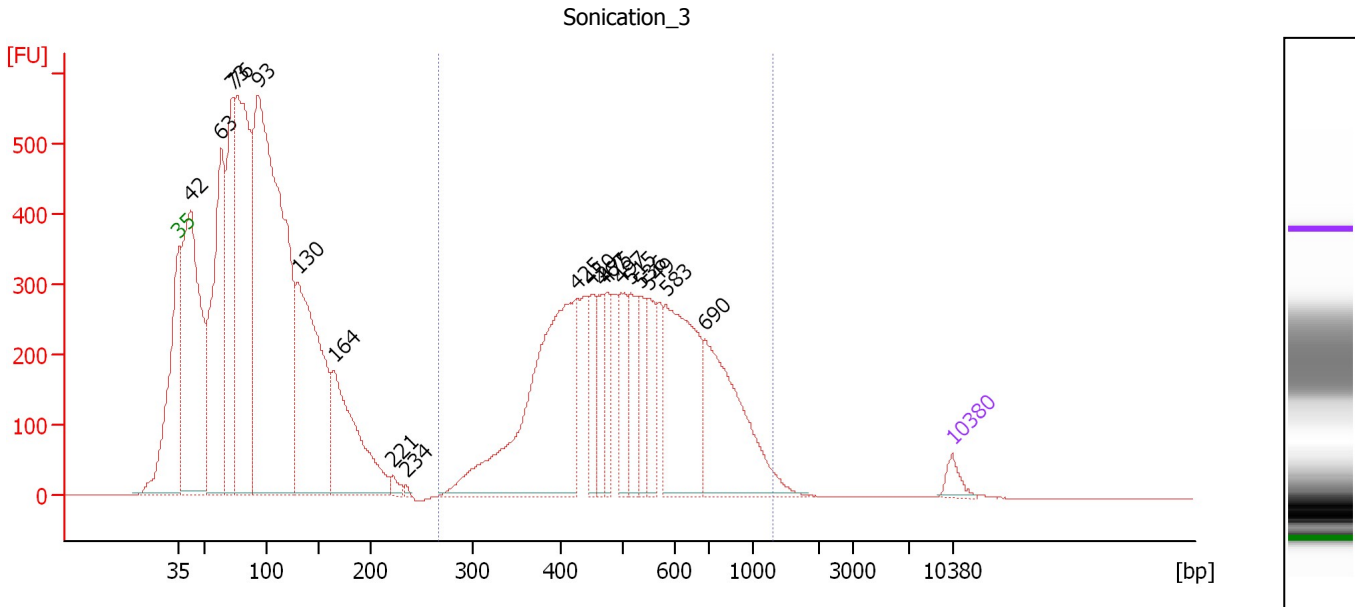
Region table for sample 8 : Sonication_2

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
101	1,367	400	153,740.1	27,813.72	24,449.7	90	48.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : Sonication_3

Number of peaks found: 19 Corr. Area 1: 6,741.4
 Noise: 0.2

Peak table for sample 9 : Sonication_3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	42	3,464.78	125,393.5	
3	63	2,664.15	63,805.1	
4	73	1,768.86	36,866.1	
5	76	3,425.24	68,171.3	
6	93	6,930.44	113,091.1	
7	130	2,719.64	31,749.6	
8	164	1,517.12	14,016.2	
9	221	70.32	481.5	
10	234	17.24	111.7	
11	425	2,440.77	8,698.1	
12	450	339.78	1,143.9	
13	467	336.25	1,090.5	
14	475	291.56	930.2	
15	497	418.03	1,273.9	
16	515	426.03	1,252.9	
17	536	266.24	752.9	
18	549	355.08	979.6	
19	583	1,249.51	3,246.1	
20	690	1,246.28	2,737.7	
21	10,380	75.00	10.9	Upper Marker

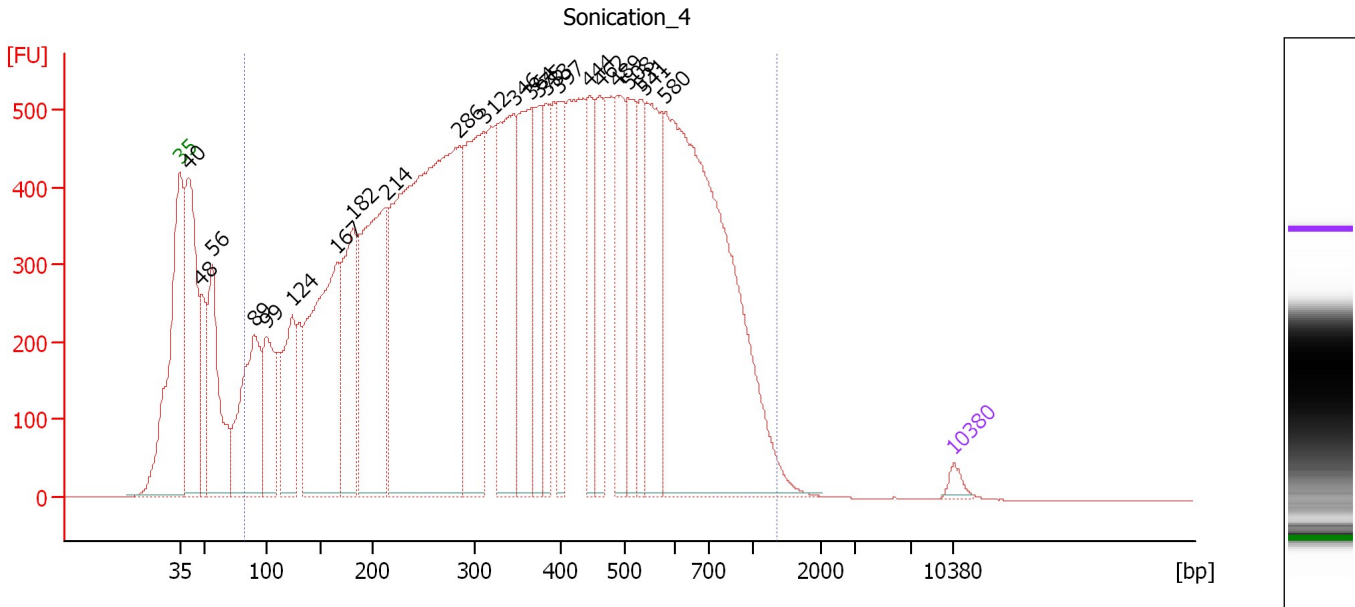
Region table for sample 9 : Sonication_3

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
266	1,314	544	26,453.4	8,492.18	6,741.4	36	31.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : Sonication_4

Number of peaks found: 23 Corr. Area 1: 26,596.0
 Noise: 0.1

Peak table for sample 10 : Sonication_4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	40	3,360.60	126,354.6	
3	48	1,076.51	34,139.2	
4	56	2,191.01	59,803.8	
5	89	2,451.05	41,569.1	
6	99	1,394.14	21,296.4	
7	124	1,447.56	17,757.9	
8	167	3,934.19	35,724.8	
9	182	2,099.60	17,513.4	
10	214	3,654.07	25,915.7	
11	286	8,837.49	46,821.8	
12	312	2,843.91	13,825.0	
13	346	2,322.44	10,180.7	
14	364	1,765.63	7,340.6	
15	375	1,308.94	5,292.2	
16	383	800.48	3,169.8	
17	397	1,020.64	3,899.4	
18	444	931.48	3,178.8	
19	462	977.17	3,204.9	
20	489	1,240.41	3,840.4	
21	508	1,152.58	3,438.9	
22	531	685.89	1,955.3	
23	541	1,741.43	4,879.4	
24	580	6,786.37	17,732.4	
25	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...

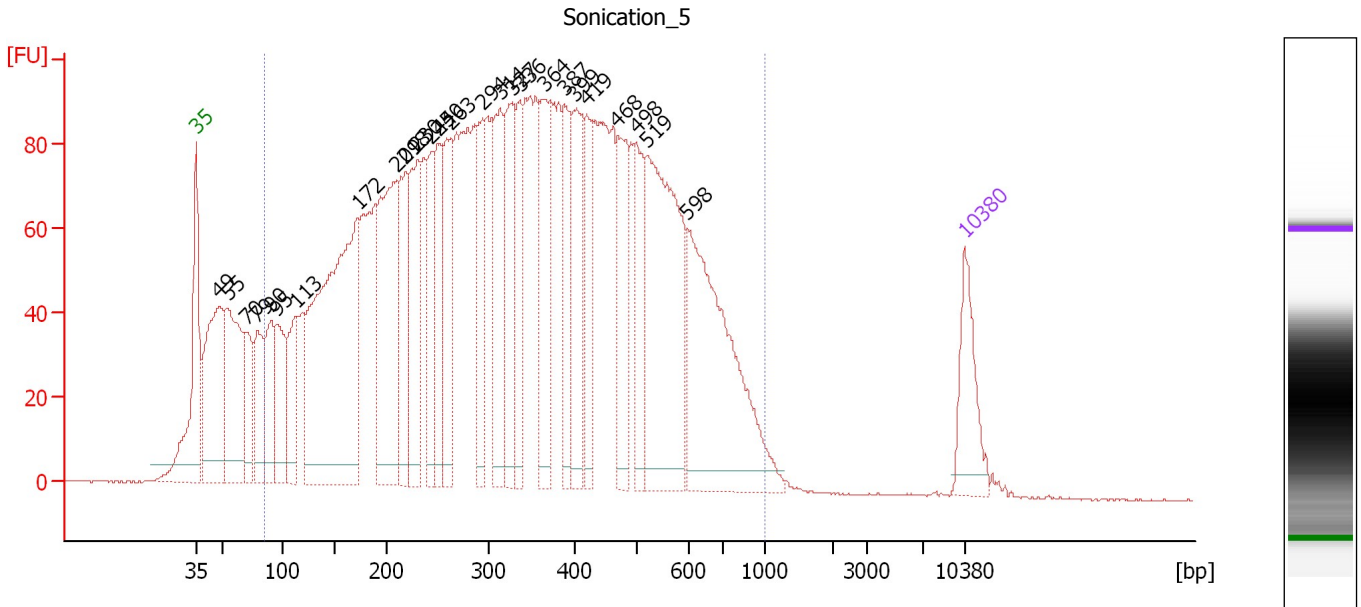
... Region table for sample 10 : Sonication 4

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
82	1,356	402	326,692.0	54,579.45	26,596.0	90	51.5	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : Sonication_5

Number of peaks found: 26 Corr. Area 1: 4,406.3
 Noise: 0.2

Peak table for sample 11 : Sonication_5

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	49	312.75	9,769.8	
3	55	265.26	7,327.9	
4	70	105.83	2,278.3	
5	79	129.84	2,489.9	
6	90	121.37	2,038.2	
7	95	137.95	2,200.8	
8	113	128.81	1,734.6	
9	172	730.18	6,446.9	
10	209	334.72	2,422.2	
11	218	171.94	1,196.0	
12	230	208.58	1,374.2	
13	244	118.42	736.3	
14	250	155.79	944.0	
15	263	162.24	933.8	
16	294	150.07	772.3	
17	314	181.06	874.5	
18	327	179.00	829.0	
19	336	117.84	530.9	
20	364	182.99	760.8	
21	387	126.27	494.3	
22	399	172.06	653.9	
23	419	104.22	377.0	
24	468	151.24	489.6	
25	498	113.97	346.6	
26	519	397.22	1,160.6	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
Modified: 7/8/2013 12:16:55 PM

Electropherogram Summary Continued ...**... Peak table for sample 11 : Sonication_5**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations
27	598	374.62	948.5	
28	▶ 10,380	75.00	10.9	Upper Marker

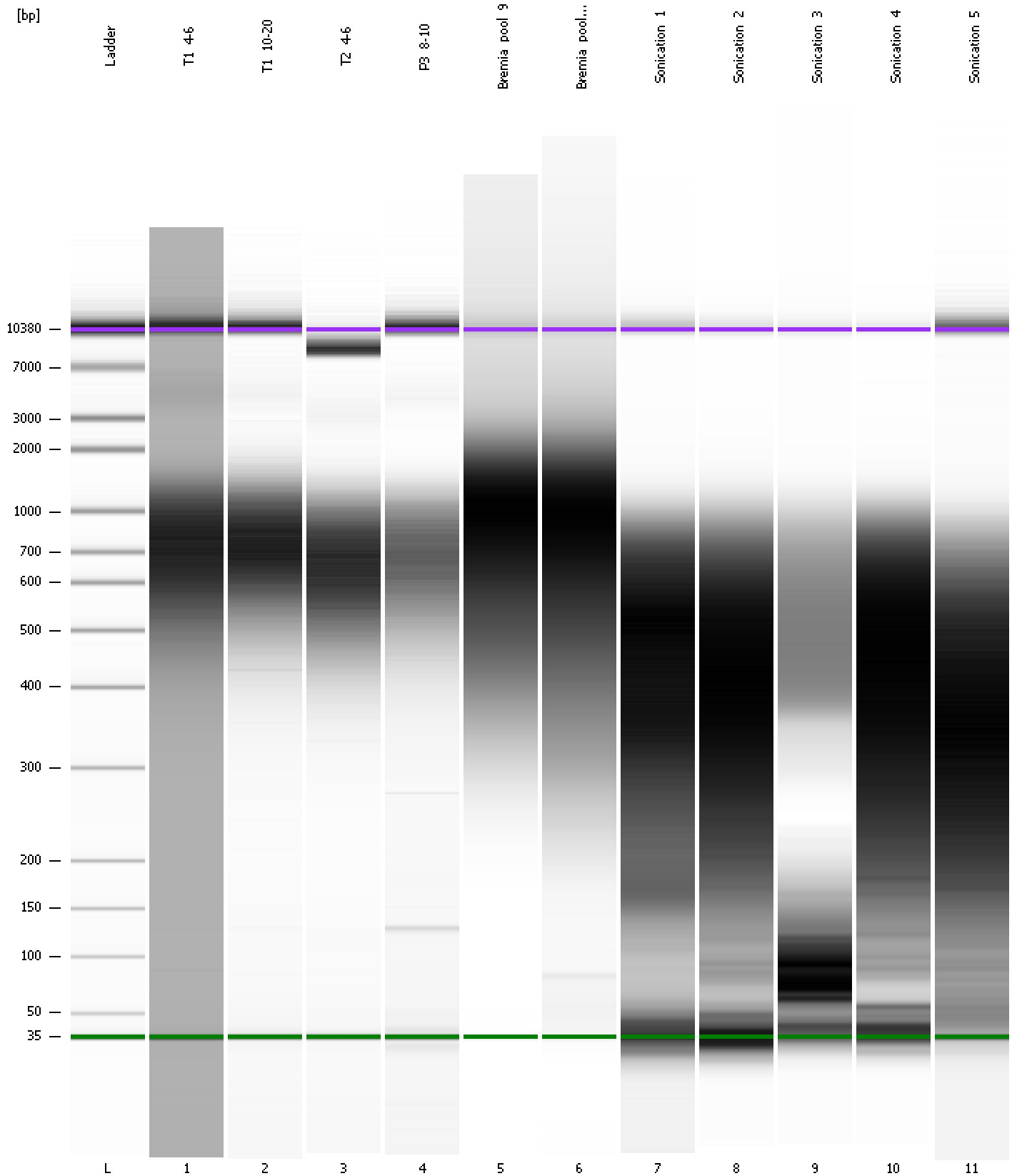
Region table for sample 11 : Sonication_5

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/ μ l]	Corr. Area	% of Total	Size distribution in CV [%]	Color
85	1,000	359	40,128.6	6,408.79	4,406.3	90	47.0	■

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
Modified: 7/8/2013 12:16:55 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad

Created: 7/8/2013 11:21:03 AM
 Modified: 7/8/2013 12:16:55 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		7/8/2013 12:02:23 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2013-07-08\2013-07-08_003.xad)		Instrument	Run		7/8/2013 11:21:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		7/8/2013 11:21:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		7/8/2013 11:21:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		7/8/2013 11:21:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		7/8/2013 11:21:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		7/8/2013 11:21:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		7/8/2013 11:21:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1