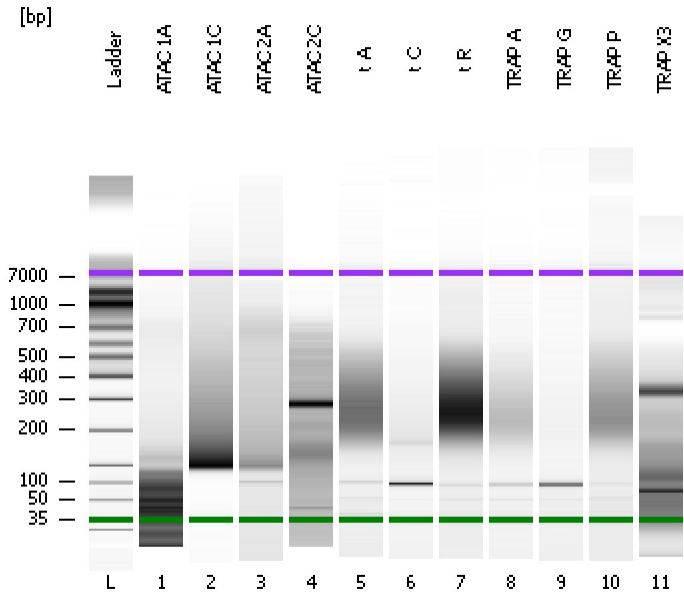


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

Created: 2/1/2017 11:36:13 AM  
Modified: 2/1/2017 12:17:31 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

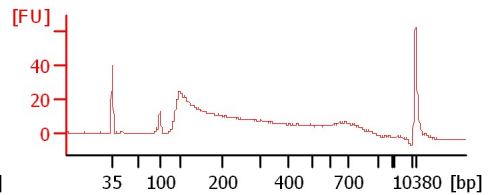
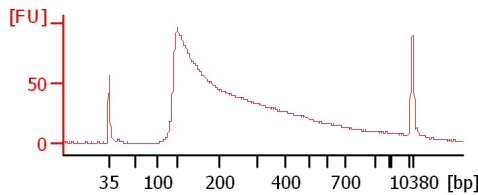
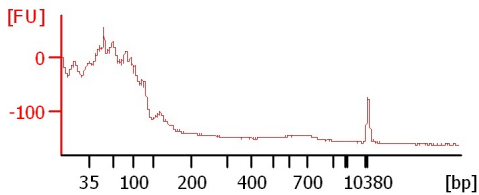
Chip Information:

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

**ATAC 1A**

**ATAC 1C**

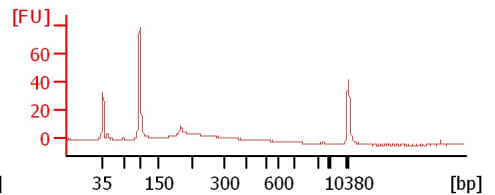
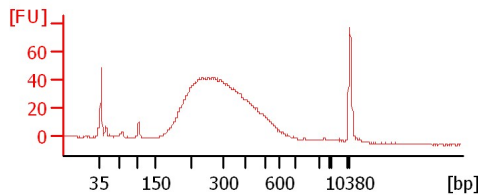
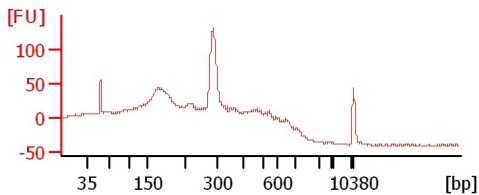
**ATAC 2A**



**ATAC 2C**

**t A**

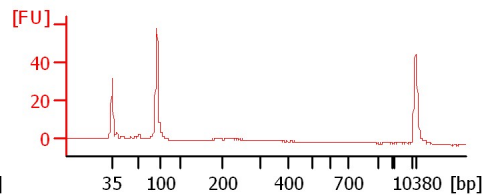
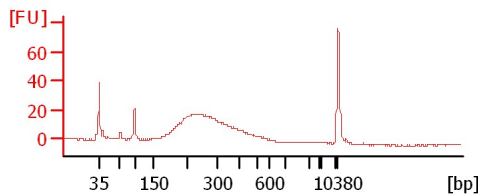
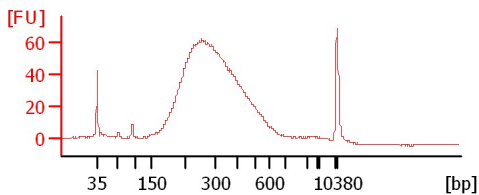
**t C**



**t R**

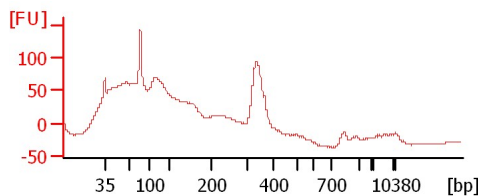
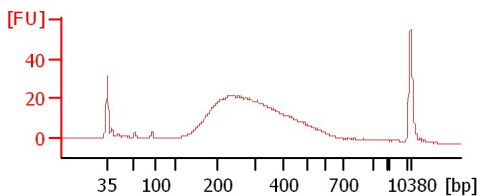
**TRAP A**

**TRAP G**



**TRAP P**

**TRAP X3**



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

Created: 2/1/2017 11:36:13 AM  
Modified: 2/1/2017 12:17:31 PM

**Electrophoresis File Run Summary (Chip Summary)**

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
ATAC 1A		<input type="checkbox"/>	✓			
ATAC 1C		<input type="checkbox"/>	✓			
ATAC 2A		<input type="checkbox"/>	✓			
ATAC 2C		<input type="checkbox"/>	✓			
t A		<input type="checkbox"/>	✓			
t C		<input type="checkbox"/>	✓			
t R		<input type="checkbox"/>	✓			
TRAP A		<input type="checkbox"/>	✓			
TRAP G		<input type="checkbox"/>	✓			
TRAP P		<input type="checkbox"/>	✓			
TRAP X3		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

**Chip Lot #**

**Reagent Kit Lot #**

**Chip Comments :**

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

Created: 2/1/2017 11:36:13 AM  
Modified: 2/1/2017 12:17:31 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

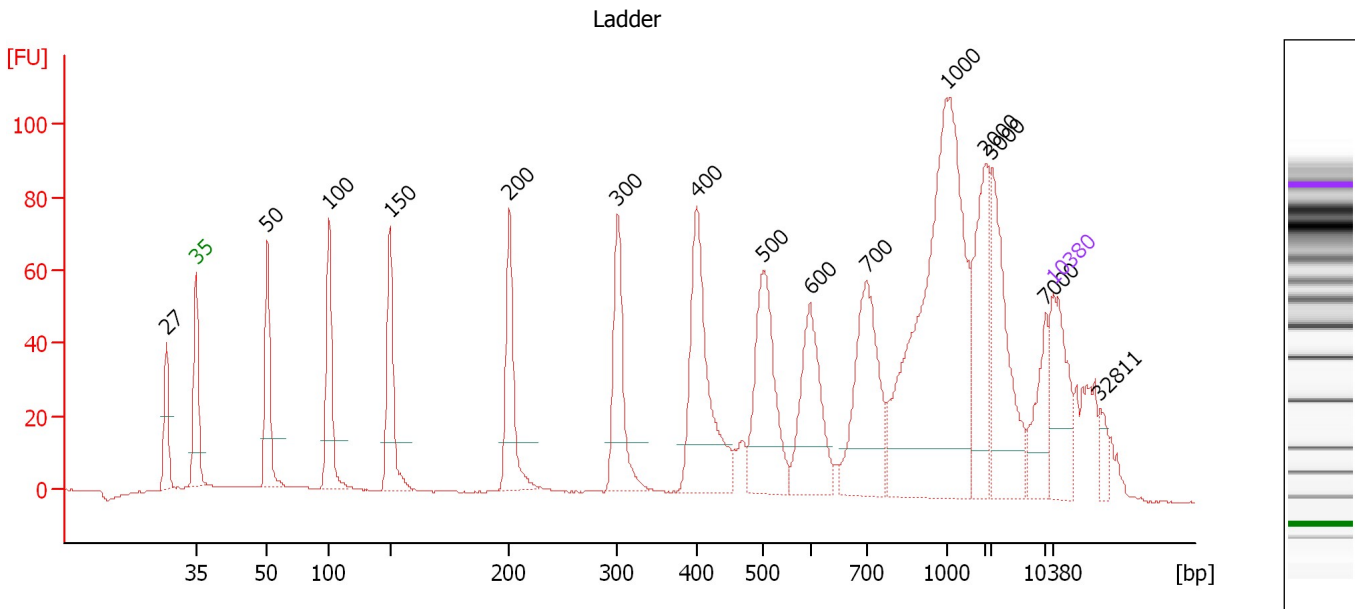
**Ladder**

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:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

Created: 2/1/2017 11:36:13 AM  
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**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.2

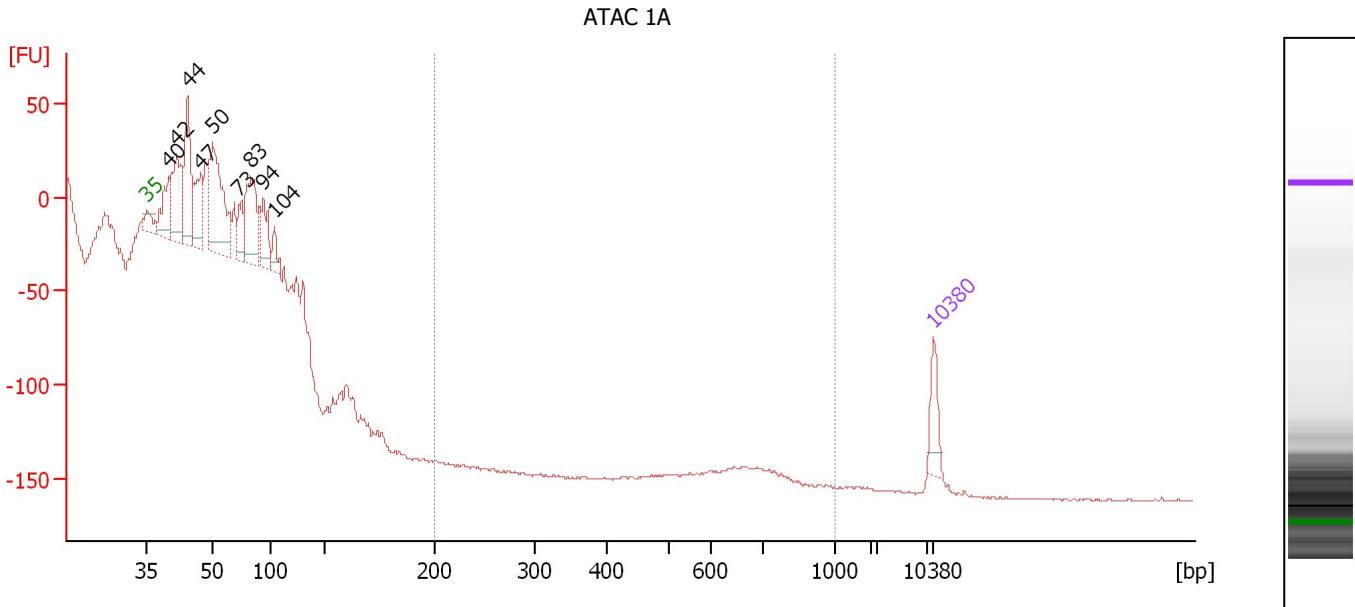
**Peak table for Ladder**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	27	0.00	0.0		40.58
2	35	125.00	5,411.3	Lower Marker	43.00
3	50	150.00	4,545.5	Ladder Peak	48.84
4	100	150.00	2,272.7	Ladder Peak	53.89
5	150	150.00	1,515.2	Ladder Peak	58.84
6	200	150.00	1,136.4	Ladder Peak	68.63
7	300	150.00	757.6	Ladder Peak	77.42
8	400	150.00	568.2	Ladder Peak	83.89
9	500	150.00	454.5	Ladder Peak	89.37
10	600	150.00	378.8	Ladder Peak	93.16
11	700	150.00	324.7	Ladder Peak	97.74
12	1,000	150.00	227.3	Ladder Peak	104.32
13	2,000	150.00	113.6	Ladder Peak	107.47
14	3,000	150.00	75.8	Ladder Peak	108.00
15	7,000	150.00	32.5	Ladder Peak	112.42
16	10,380	75.00	10.9	Upper Marker	113.00
17	32,811	0.00	0.0		116.84

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

Created: 2/1/2017 11:36:13 AM  
 Modified: 2/1/2017 12:17:31 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : ATAC 1A**

Number of peaks found: 9                      Corr. Area 1: 0.0  
 Noise: 0.5

**Peak table for sample 1 : ATAC 1A**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	40	225.52	8,570.9		44.90
3	42	329.11	11,995.2		45.56
4	44	371.63	12,725.3		46.60
5	47	220.83	7,130.2		47.64
6	50	569.88	17,280.2		48.83
7	73	110.88	2,291.3		51.20
8	83	336.00	6,120.8		52.19
9	94	164.58	2,640.8		53.33
10	104	61.51	893.0		54.33
11	10,380	75.00	10.9	Upper Marker	113.00

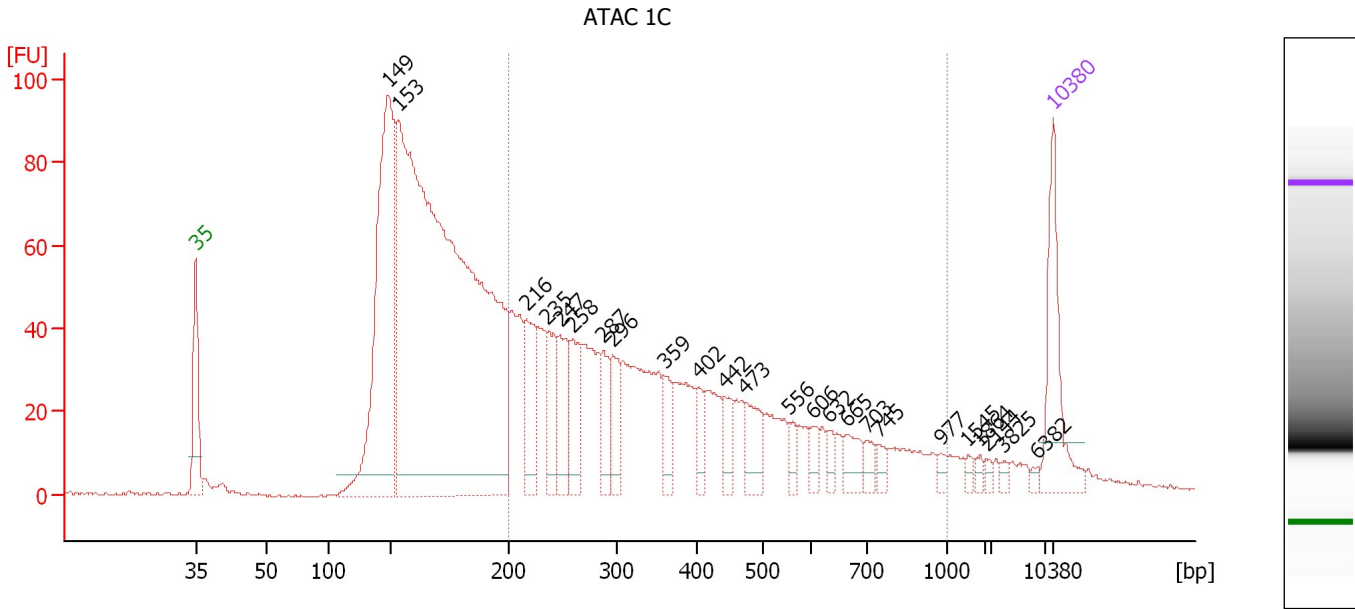
**Region table for sample 1 : ATAC 1A**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	0	0.0	0.0	0.00	0	0.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : ATAC 1C**

Number of peaks found: 24                      Corr. Area 1: 1,060.5  
 Noise: 0.3

**Peak table for sample 2 : ATAC 1C**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	149	486.33	4,961.4		58.70
3	153	1,615.70	15,960.8		59.50
4	216	100.02	703.0		70.00
5	235	69.17	446.9		71.67
6	247	73.22	449.9		72.73
7	258	69.50	408.0		73.74
8	287	47.22	249.5		76.26
9	296	47.99	245.7		77.07
10	359	35.81	151.0		81.26
11	402	24.38	91.8		84.03
12	442	28.25	96.8		86.20
13	473	43.92	140.5		87.92
14	556	14.11	38.4		91.50
15	606	15.51	38.8		93.42
16	632	11.73	28.1		94.63
17	665	25.98	59.2		96.14
18	703	13.45	29.0		97.81
19	745	12.02	24.4		98.72
20	977	7.50	11.6		103.81
21	1,545	5.53	5.4		106.04
22	1,864	4.75	3.9		107.04
23	2,144	4.40	3.1		107.55
24	3,825	4.45	1.8		108.91
25	6,382	3.99	0.9		111.74

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

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 Modified: 2/1/2017 12:17:31 PM

### Electropherogram Summary Continued ...

#### ... Peak table for sample 2 : ATAC 1C

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	10,380	75.00	10.9	Upper Marker	113.00

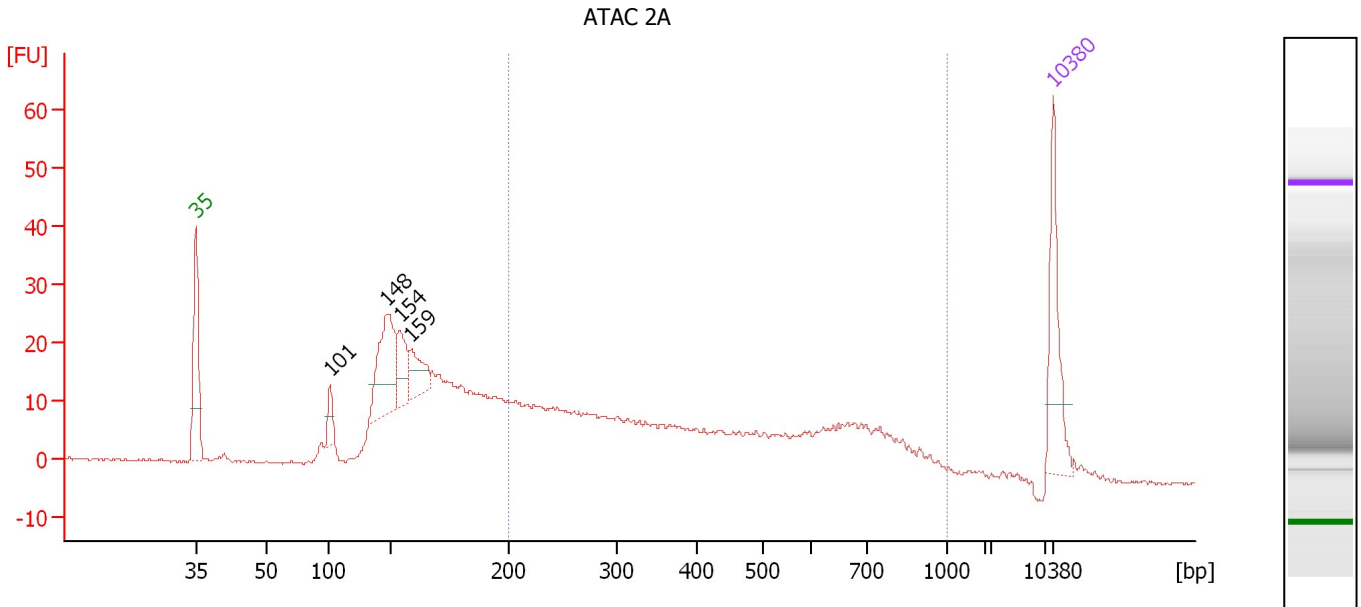
#### Region table for sample 2 : ATAC 1C

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	407	1,060.5	6,725.1	1,427.17	45	46.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

Created: 2/1/2017 11:36:13 AM  
 Modified: 2/1/2017 12:17:31 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : ATAC 2A**

Number of peaks found: 4                      Corr. Area 1: 330.8  
 Noise: 0.3

**Peak table for sample 3 : ATAC 2A**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	101	26.82	403.5		53.96
3	148	141.63	1,454.8		58.60
4	154	57.87	568.4		59.68
5	159	53.70	511.8		60.60
6	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 3 : ATAC 2A**

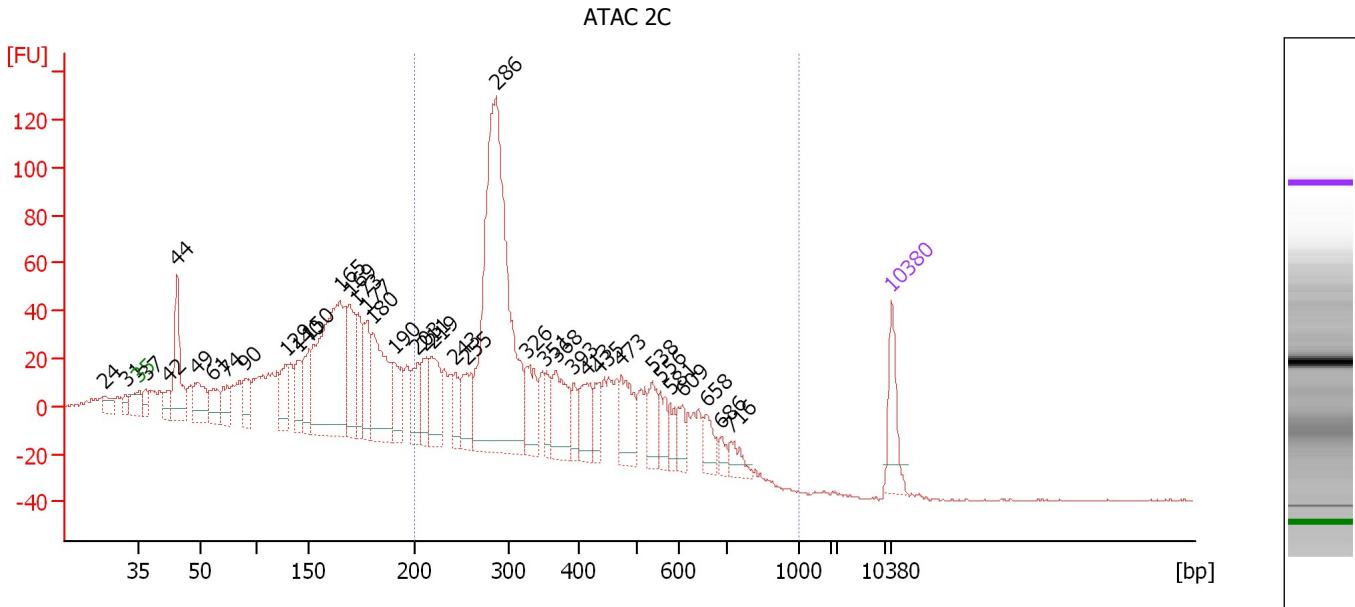
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	451	330.8	3,566.6	812.62	50	44.7



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

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 Modified: 2/1/2017 12:17:31 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : ATAC 2C**

Number of peaks found: 38                      Corr. Area 1: 1,310.3  
 Noise: 0.3

**Peak table for sample 4 : ATAC 2C**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	24	0.00	0.0		39.88
2	31	0.00	0.0		41.81
3	35	125.00	5,411.3	Lower Marker	43.00
4	37	48.12	1,989.0		43.64
5	42	53.15	1,905.6		45.83
6	44	238.51	8,182.5		46.57
7	49	122.23	3,779.2		48.45
8	61	80.28	1,998.2		49.94
9	74	68.42	1,407.9		51.23
10	90	85.43	1,433.3		52.92
11	129	95.98	1,130.1		56.73
12	140	96.84	1,050.3		57.82
13	150	110.45	1,114.8		58.86
14	165	621.72	5,716.0		61.74
15	169	195.68	1,750.7		62.63
16	173	123.43	1,081.7		63.33
17	177	122.75	1,048.1		64.22
18	180	275.53	2,312.9		64.81
19	190	104.44	831.2		66.75
20	203	89.87	669.5		68.93
21	211	86.60	621.1		69.62
22	219	139.54	967.2		70.27
23	243	62.82	391.0		72.45
24	255	102.88	610.7		73.49
25	286	1,040.35	5,506.1		76.22

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

Created: 2/1/2017 11:36:13 AM  
 Modified: 2/1/2017 12:17:31 PM

### Electropherogram Summary Continued ...

#### ... Peak table for sample 4 : ATAC 2C

Peak	Size [bp]	Conc. [pg/ $\mu$ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	326	108.75	505.8		79.09
27	351	52.96	228.5		80.73
28	368	140.93	580.4		81.82
29	393	46.02	177.4		83.45
30	413	90.18	331.0		84.59
31	435	50.84	177.3		85.78
32	473	113.98	364.8		87.92
33	538	60.04	169.2		90.79
34	556	48.60	132.5		91.48
35	581	31.33	81.8		92.43
36	609	39.96	99.4		93.57
37	658	44.32	102.1		95.80
38	686	20.07	44.3		97.09
39	716	36.50	77.3		98.08
40	▶ 10,380	75.00	10.9	Upper Marker	113.00

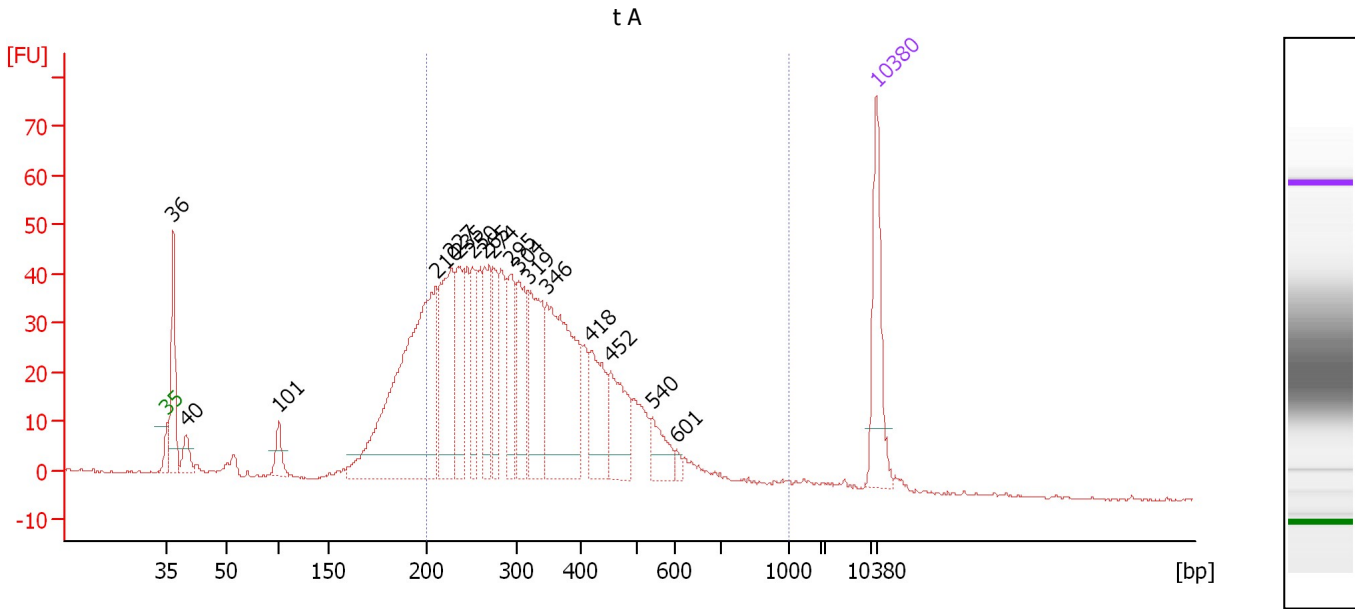
#### Region table for sample 4 : ATAC 2C

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ $\mu$ l]	% of Total	Size distribution in CV [%]
200	1,000	368	1,310.3	11,473.6	■ 2,421.69	51	34.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : t A**

Number of peaks found: 17                      Corr. Area 1: 993.4  
 Noise: 0.3

**Peak table for sample 5 : t A**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	36	132.10	5,485.5		43.58
3	40	31.73	1,210.1		44.84
4	101	32.50	487.3		54.00
5	210	472.56	3,406.6		69.53
6	227	175.89	1,174.3		71.00
7	235	104.85	676.8		71.68
8	250	72.93	441.5		73.05
9	265	76.62	438.6		74.32
10	274	68.98	381.1		75.16
11	295	73.03	375.6		76.95
12	304	91.54	456.1		77.68
13	319	129.41	615.3		78.63
14	346	230.51	1,008.4		80.42
15	418	81.94	296.8		84.89
16	452	78.72	263.9		86.74
17	540	35.04	98.3		90.89
18	601	6.01	15.2		93.21
19	10,380	75.00	10.9	Upper Marker	113.00

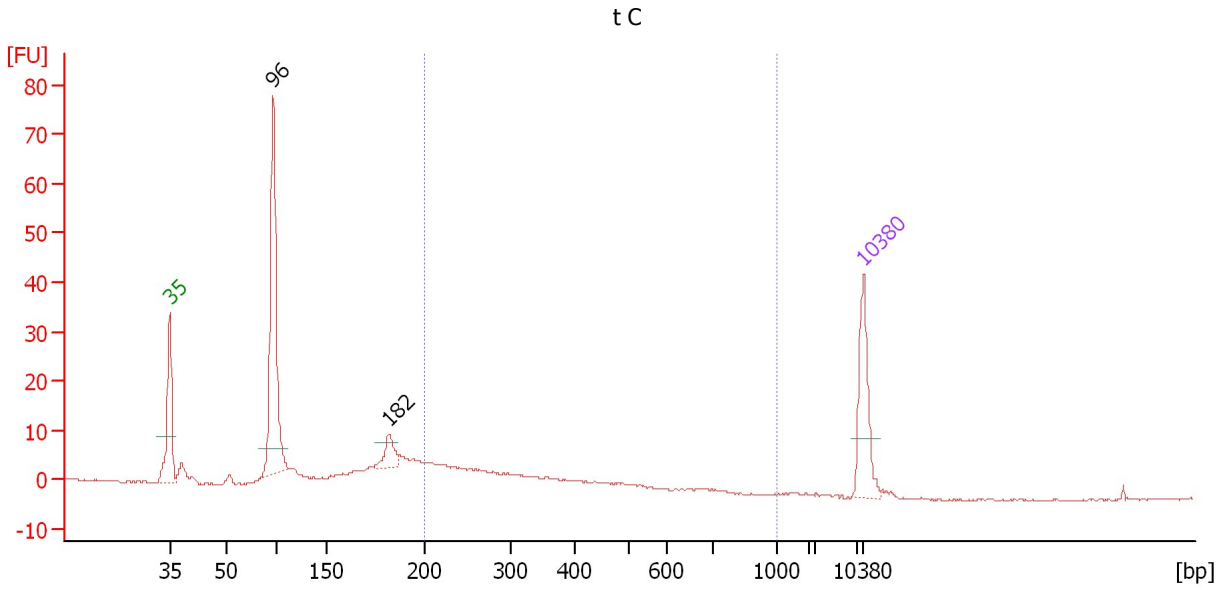
**Region table for sample 5 : t A**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	341	993.4	8,626.2	1,713.67	76	34.2

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : t C**

Number of peaks found: 2                      Corr. Area 1: 67.4  
 Noise: 0.2

**Peak table for sample 6 : t C**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	96	360.04	5,686.3		53.48
3	182	40.77	339.0		65.15
4	10,380	75.00	10.9	Upper Marker	113.00

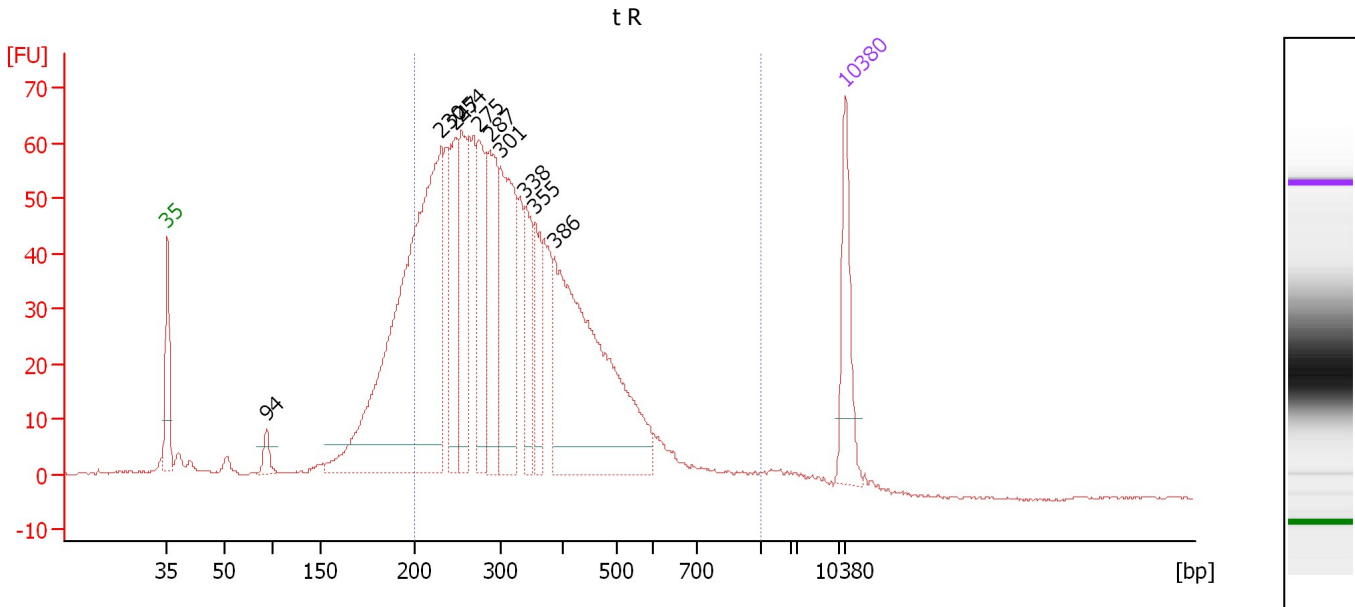
**Region table for sample 6 : t C**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	308	67.4	1,066.8	196.46	26	31.2

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

Created: 2/1/2017 11:36:13 AM  
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**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : t R**

Number of peaks found: 10                      Corr. Area 1: 1,337.7  
 Noise: 0.2

**Peak table for sample 7 : t R**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	94	25.14	407.2		53.24
3	230	851.29	5,596.9		71.31
4	247	167.15	1,026.4		72.74
5	254	185.49	1,108.1		73.35
6	275	147.15	810.9		75.22
7	287	177.15	933.7		76.32
8	301	242.38	1,220.6		77.48
9	338	96.83	433.7		79.90
10	355	81.70	348.4		81.00
11	386	473.11	1,857.4		82.98
12	10,380	75.00	10.9	Upper Marker	113.00

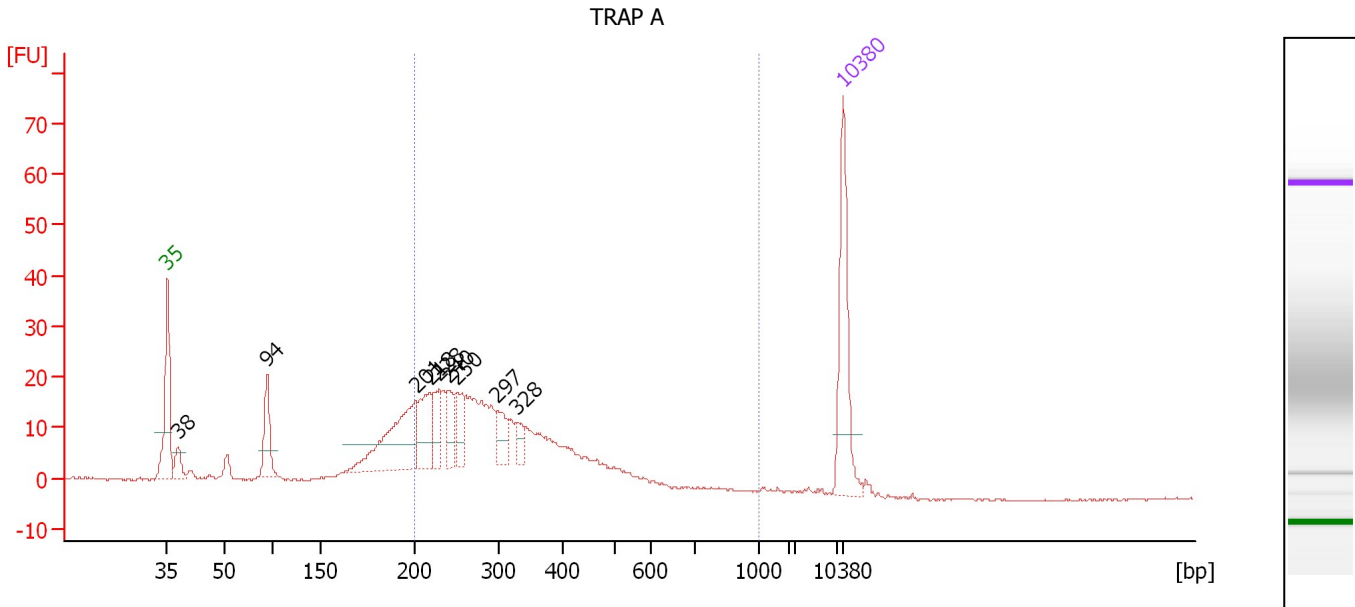
**Region table for sample 7 : t R**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	342	1,337.7	12,641.8	2,506.78	80	36.3

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

Created: 2/1/2017 11:36:13 AM  
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**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : TRAP A**

Number of peaks found: 9                      Corr. Area 1: 332.5  
 Noise: 0.2

**Peak table for sample 8 : TRAP A**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	38	24.70	988.9		44.10
3	94	54.55	875.3		53.33
4	201	132.94	1,000.6		68.75
5	218	59.02	409.7		70.24
6	228	35.13	233.1		71.12
7	240	29.53	186.7		72.12
8	250	31.82	192.6		73.06
9	297	28.62	145.8		77.20
10	328	14.65	67.6		79.24
11	10,380	75.00	10.9	Upper Marker	113.00

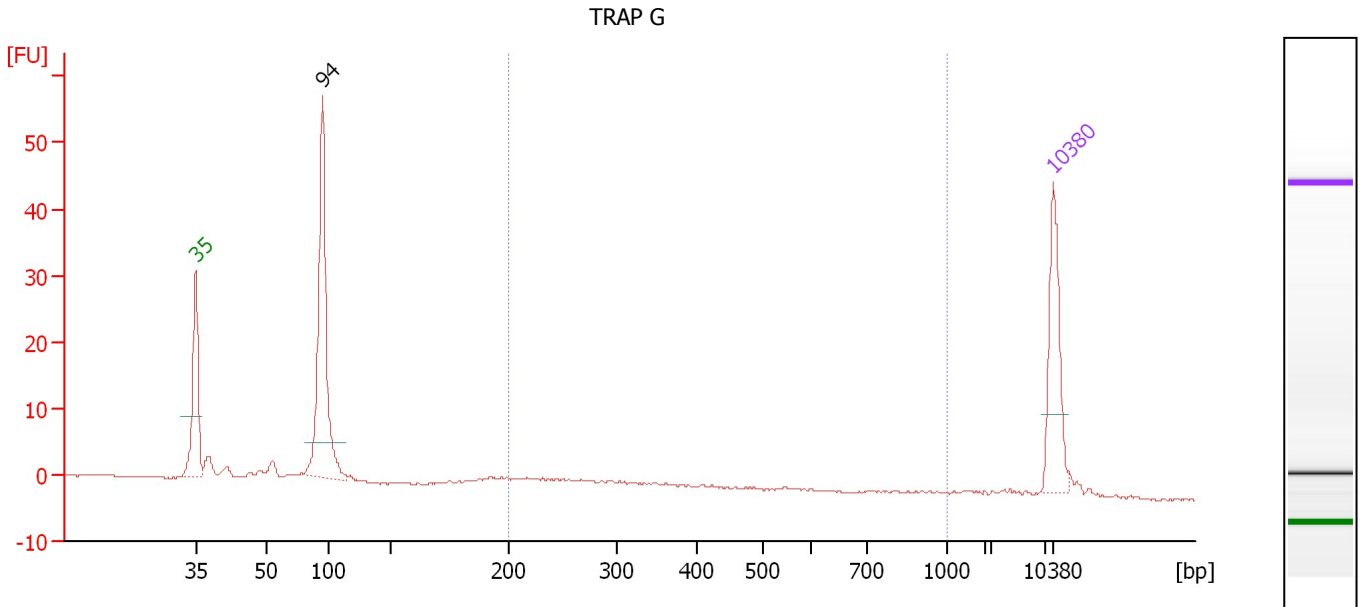
**Region table for sample 8 : TRAP A**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	312	332.5	3,186.5	598.81	69	29.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : TRAP G**

Number of peaks found: 1                      Corr. Area 1: 4.9  
 Noise: 0.2

**Peak table for sample 9 : TRAP G**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	94	286.91	4,603.7		53.33
3	10,380	75.00	10.9	Upper Marker	113.00

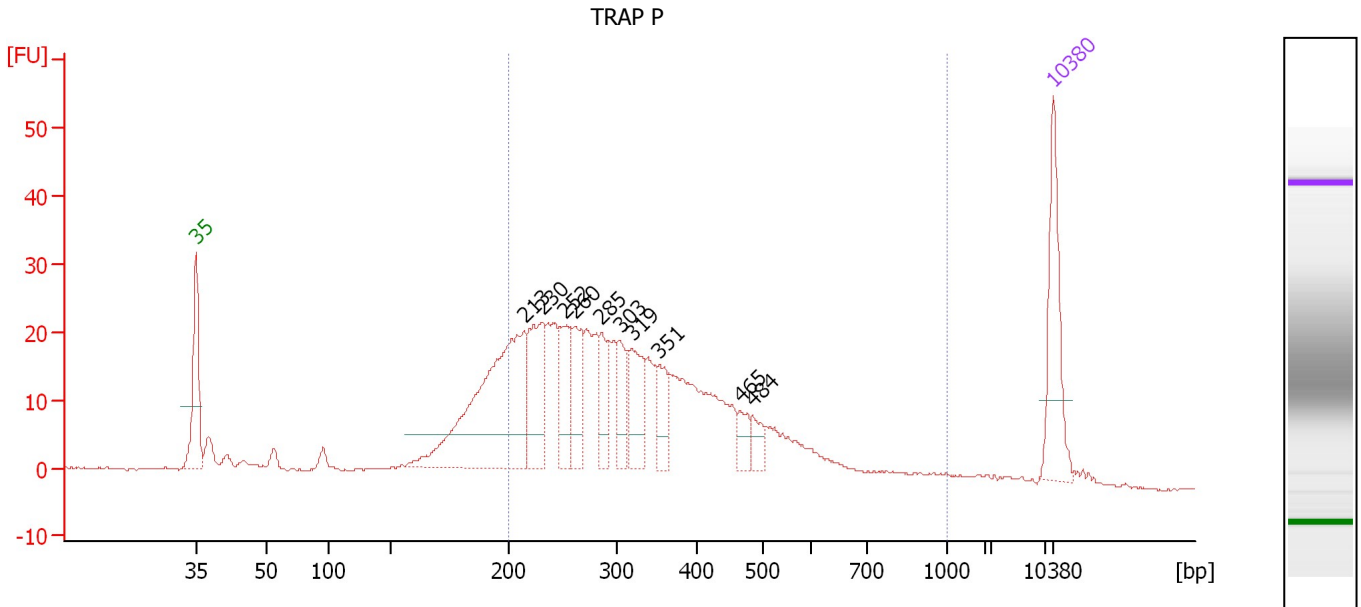
**Region table for sample 9 : TRAP G**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	249	4.9	95.0	15.44	5	13.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : TRAP P**

Number of peaks found: 10                      Corr. Area 1: 449.7  
 Noise: 0.2

**Peak table for sample 10 : TRAP P**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	213	339.09	2,407.2		69.81
3	230	107.33	707.5		71.26
4	252	63.14	379.7		73.20
5	260	61.02	355.3		73.92
6	285	47.60	252.7		76.14
7	303	46.30	231.2		77.64
8	319	67.16	319.2		78.64
9	351	43.29	186.7		80.75
10	465	21.21	69.1		87.46
11	484	18.87	59.0		88.52
12	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 10 : TRAP P**

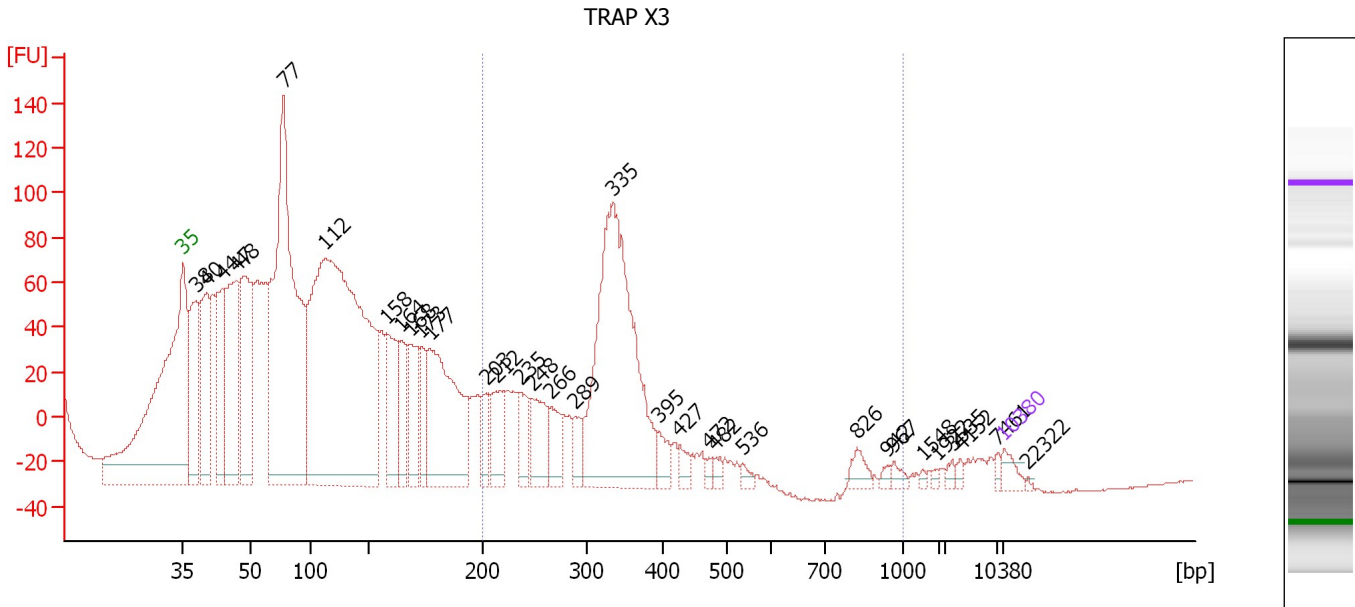
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	331	449.7	5,638.0	1,100.80	79	32.3



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : TRAP X3**

Number of peaks found: 33                      Corr. Area 1: 826.9  
 Noise: 0.4

**Peak table for sample 11 : TRAP X3**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	38	1,324.40	52,921.6		44.14
3	40	1,262.59	47,664.3		45.00
4	44	1,170.55	40,644.1		46.36
5	47	1,678.88	54,504.2		47.55
6	48	1,550.67	48,757.1		48.14
7	77	4,895.98	96,087.9		51.59
8	112	7,173.59	96,968.2		55.09
9	158	844.02	8,093.6		60.41
10	164	475.48	4,398.0		61.55
11	168	575.55	5,191.2		62.36
12	173	377.54	3,313.6		63.27
13	177	1,791.18	15,369.6		64.05
14	203	270.12	2,014.6		68.91
15	212	440.64	3,142.3		69.73
16	235	294.78	1,898.8		71.73
17	248	475.30	2,908.2		72.82
18	266	288.52	1,645.1		74.41
19	289	187.43	982.6		76.45
20	335	3,056.92	13,829.2		79.68
21	395	154.45	592.0		83.59
22	427	94.81	336.5		85.36
23	472	46.74	150.1		87.82
24	482	58.49	184.0		88.36
25	536	53.71	151.9		90.73

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

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 Modified: 2/1/2017 12:17:31 PM

**Electropherogram Summary Continued ...**

**... Peak table for sample 11 : TRAP X3**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	826	114.36	209.8		100.50
27	942	33.33	53.6		103.05
28	967	54.39	85.2		103.59
29	1,548	18.02	17.6		106.05
30	1,922	17.61	13.9		107.23
31	3,535	30.00	12.9		108.59
32	4,152	29.57	10.8		109.27
33	7,461	29.50	6.0		112.50
34	10,380	75.00	10.9	Upper Marker	113.00
35	22,322	0.00	0.0		115.05

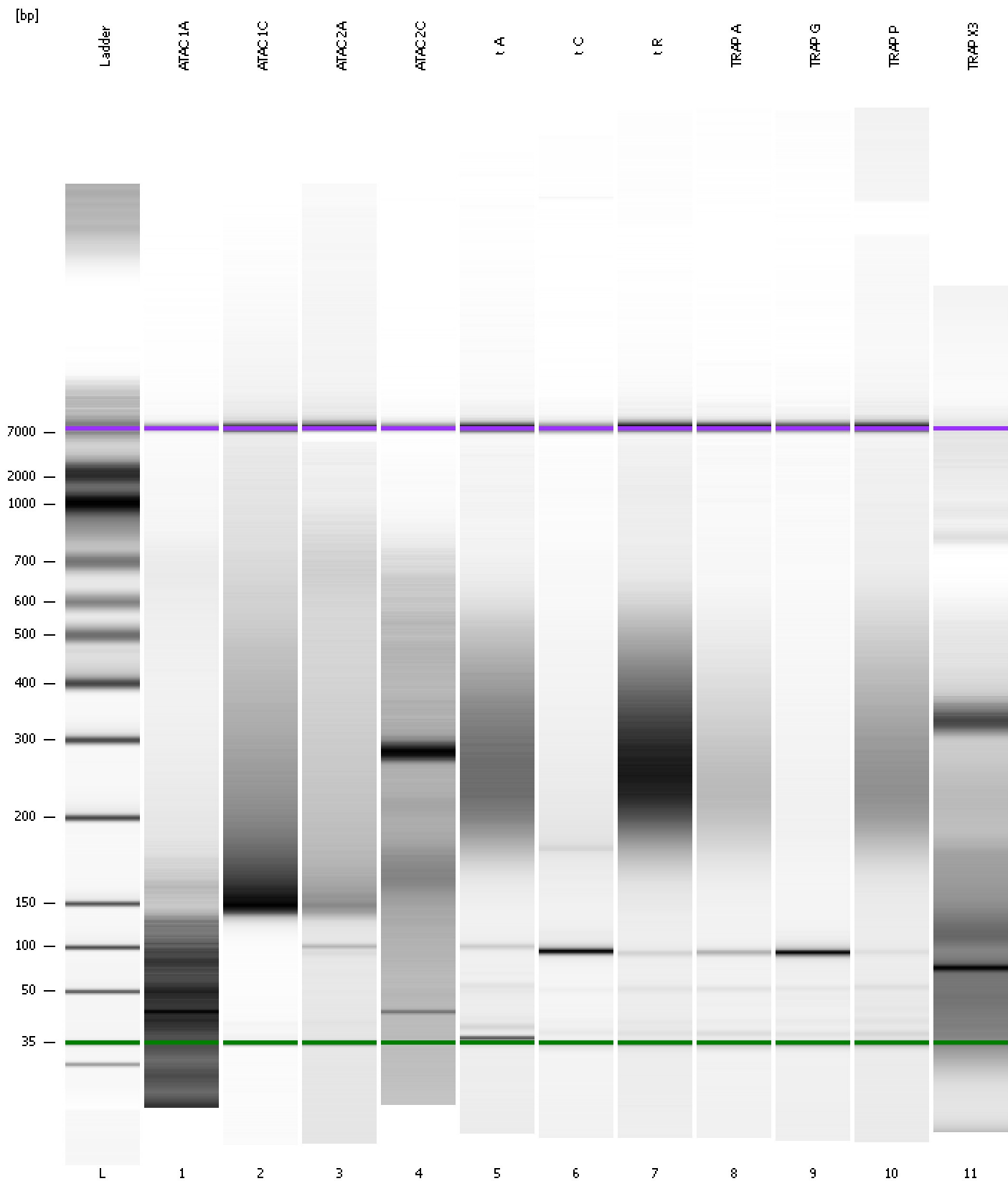
**Region table for sample 11 : TRAP X3**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	320	826.9	21,800.4	4,302.31	21	27.2

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

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**Gel Image**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-01\2017-02-01\_001.xad

Created: 2/1/2017 11:36:13 AM  
 Modified: 2/1/2017 12:17:31 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		2/1/2017 12:17:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2017-02-01\2017-02-01_001.xad)		Instrument	Run		2/1/2017 11:36:19 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		2/1/2017 11:36:19 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		2/1/2017 11:36:19 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		2/1/2017 11:36:19 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		2/1/2017 11:36:19 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		2/1/2017 11:36:19 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		2/1/2017 11:36:19 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1