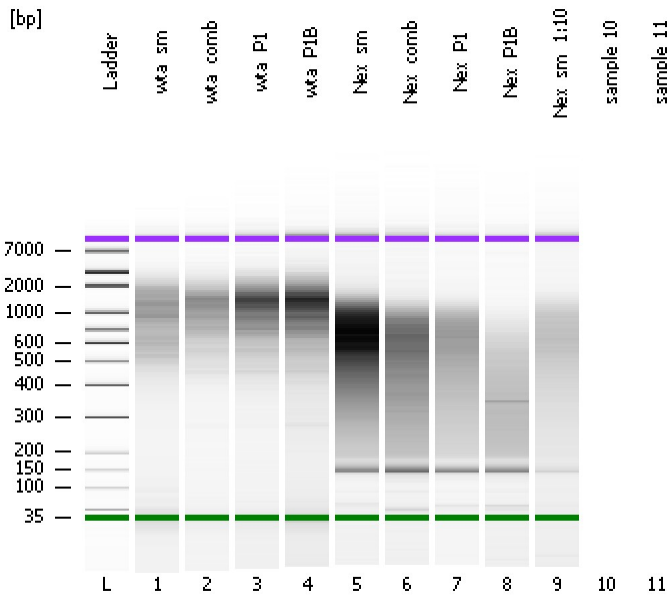


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
Modified: 4/10/2017 11:45:00 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

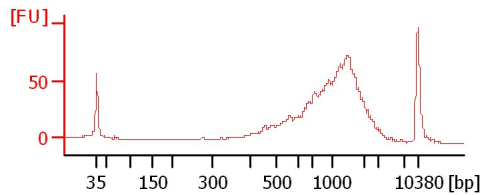
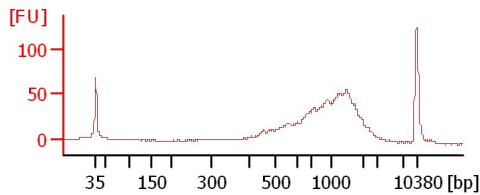
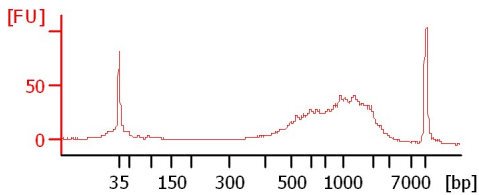
Chip Information:

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

wta_sm

wta_comb

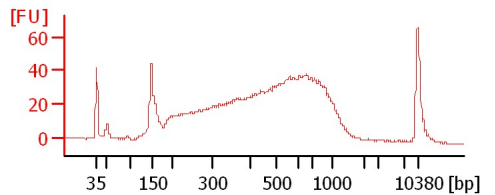
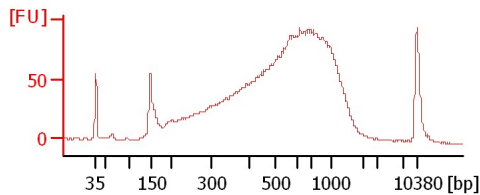
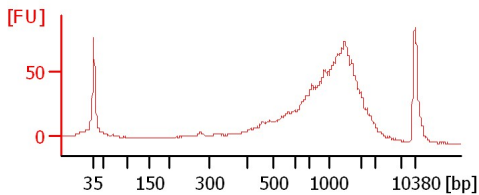
wta_P1



wta_P1B

Nex_sm

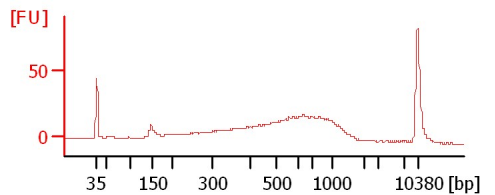
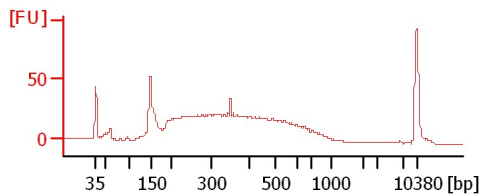
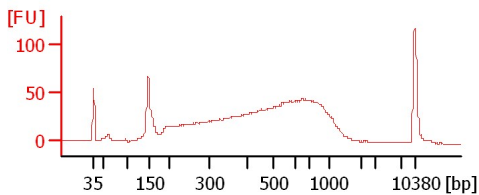
Nex_comb



Nex_P1

Nex_P1B

Nex_sm_1:10



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
Modified: 4/10/2017 11:45:00 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
wta_sm		<input type="checkbox"/>	✓			
wta_comb		<input type="checkbox"/>	✓			
wta_P1		<input type="checkbox"/>	✓			
wta_P1B		<input type="checkbox"/>	✓			
Nex_sm		<input type="checkbox"/>	✓			
Nex_comb		<input type="checkbox"/>	✓			
Nex_P1		<input type="checkbox"/>	✓			
Nex_P1B		<input type="checkbox"/>	✓			
Nex_sm_1:10		<input type="checkbox"/>	✓			
sample 10		<input type="checkbox"/>				
sample 11		<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\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
Modified: 4/10/2017 11:45:00 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

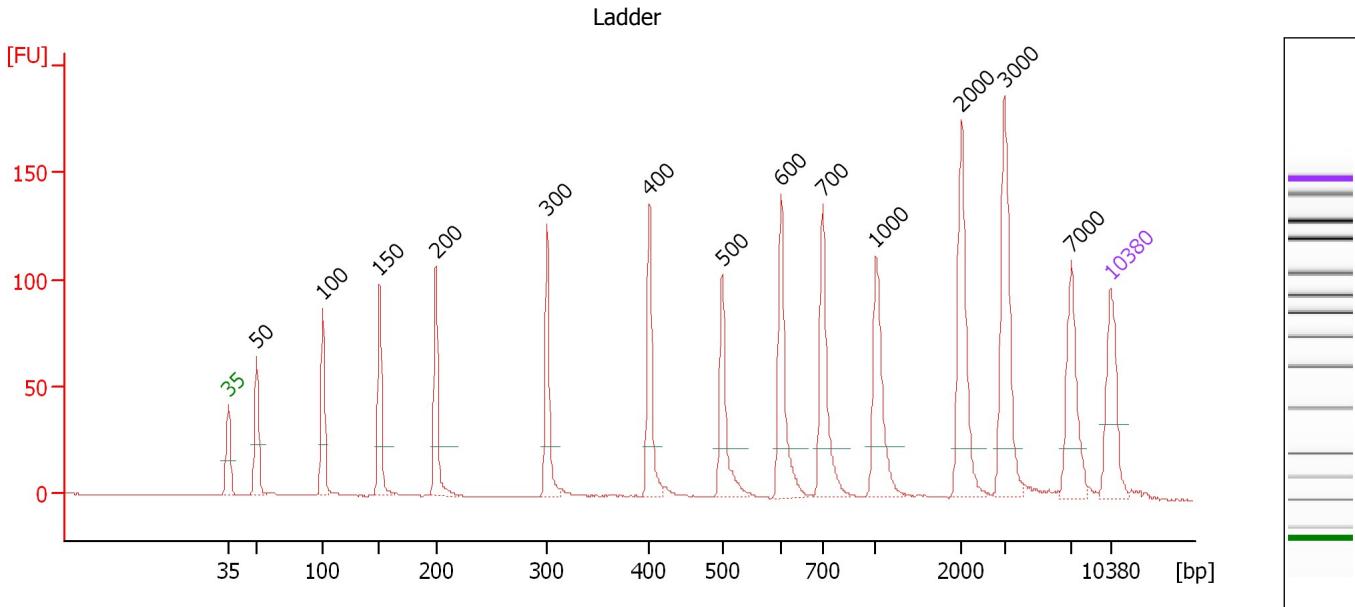
Ladder

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:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
 Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

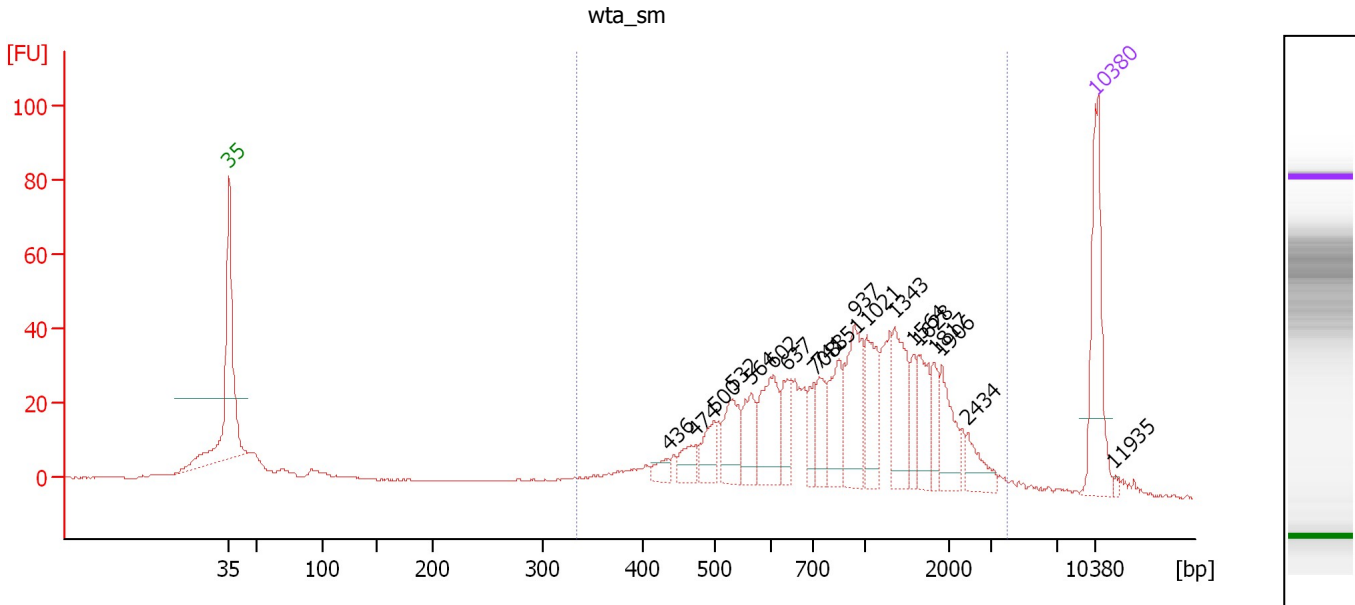
Peak table for Ladder

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	50	150.00	4,545.5	Ladder Peak	45.24
3	100	150.00	2,272.7	Ladder Peak	50.49
4	150	150.00	1,515.2	Ladder Peak	54.98
5	200	150.00	1,136.4	Ladder Peak	59.46
6	300	150.00	757.6	Ladder Peak	68.27
7	400	150.00	568.2	Ladder Peak	76.39
8	500	150.00	454.5	Ladder Peak	82.19
9	600	150.00	378.8	Ladder Peak	86.85
10	700	150.00	324.7	Ladder Peak	90.15
11	1,000	150.00	227.3	Ladder Peak	94.38
12	2,000	150.00	113.6	Ladder Peak	101.19
13	3,000	150.00	75.8	Ladder Peak	104.58
14	7,000	150.00	32.5	Ladder Peak	109.91
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
 Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : wta_sm

Number of peaks found: 19 Corr. Area 1: 806.2
 Noise: 0.1

Peak table for sample 1 : wta_sm

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	436	11.96	41.6		78.45
3	474	21.56	68.9		80.69
4	500	28.78	87.2		82.20
5	532	45.43	129.5		83.66
6	564	35.22	94.6		85.17
7	602	61.31	154.4		86.90
8	637	30.87	73.4		88.06
9	708	21.18	45.3		90.26
10	741	29.74	60.8		90.73
11	851	42.67	76.0		92.28
12	937	68.00	110.0		93.49
13	1,021	49.85	74.0		94.52
14	1,343	54.22	61.2		96.72
15	1,564	18.41	17.8		98.22
16	1,628	35.25	32.8		98.66
17	1,817	18.85	15.7		99.95
18	1,906	36.09	28.7		100.55
19	2,434	20.55	12.8		102.66
20	10,380	75.00	10.9	Upper Marker	113.00
21	11,935	0.00	0.0		114.42

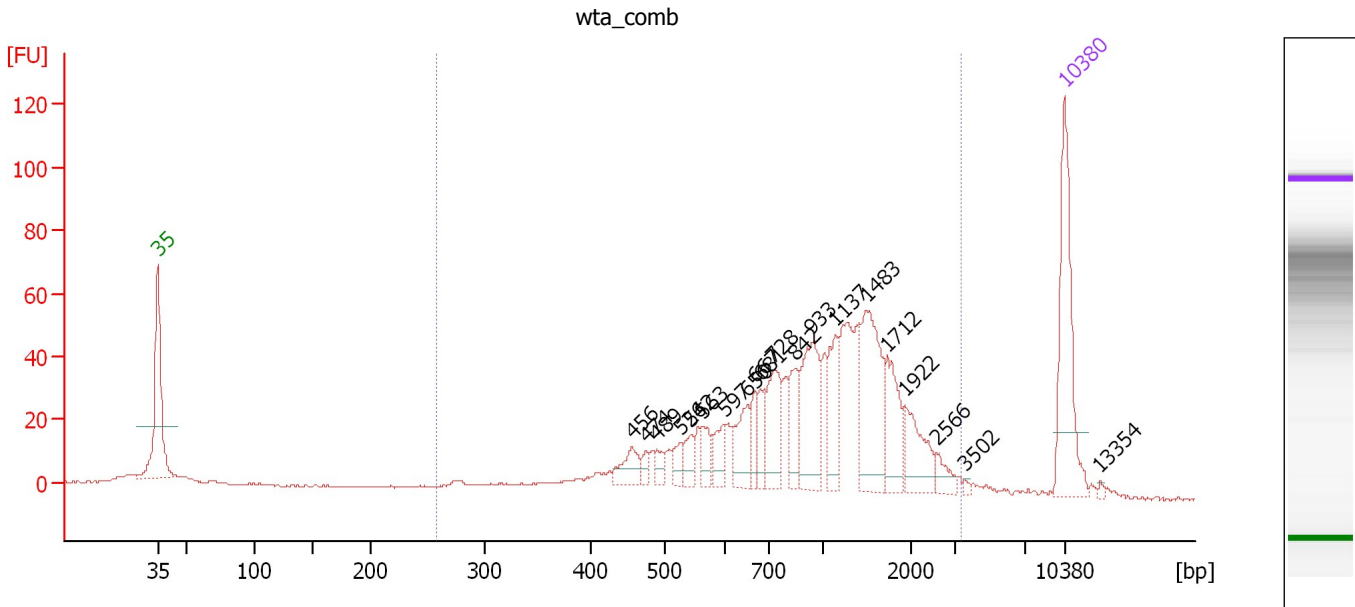
Region table for sample 1 : wta_sm

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
335	3,939	1,076	806.2	1,592.9	806.73	88	55.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
 Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 : wta_comb

Number of peaks found: 20 Corr. Area 1: 878.6
 Noise: 0.3

Peak table for sample 2 : wta_comb

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	456	20.69	68.8		79.63
3	474	6.70	21.4		80.70
4	489	8.92	27.7		81.54
5	526	12.00	34.5		83.41
6	542	13.29	37.2		84.13
7	563	15.29	41.2		85.11
8	597	18.55	47.1		86.71
9	650	31.21	72.8		88.49
10	667	16.75	38.0		89.07
11	681	17.93	39.9		89.52
12	728	44.66	92.9		90.54
13	842	27.57	49.6		92.15
14	933	62.35	101.2		93.44
15	1,137	35.39	47.2		95.31
16	1,483	77.74	79.4		97.67
17	1,712	37.65	33.3		99.23
18	1,922	32.93	26.0		100.66
19	2,566	9.20	5.4		103.11
20	3,502	1.63	0.7		105.25
21	10,380	75.00	10.9	Upper Marker	113.00
22	13,354	0.00	0.0		115.72

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...

... Region table for sample 2 :

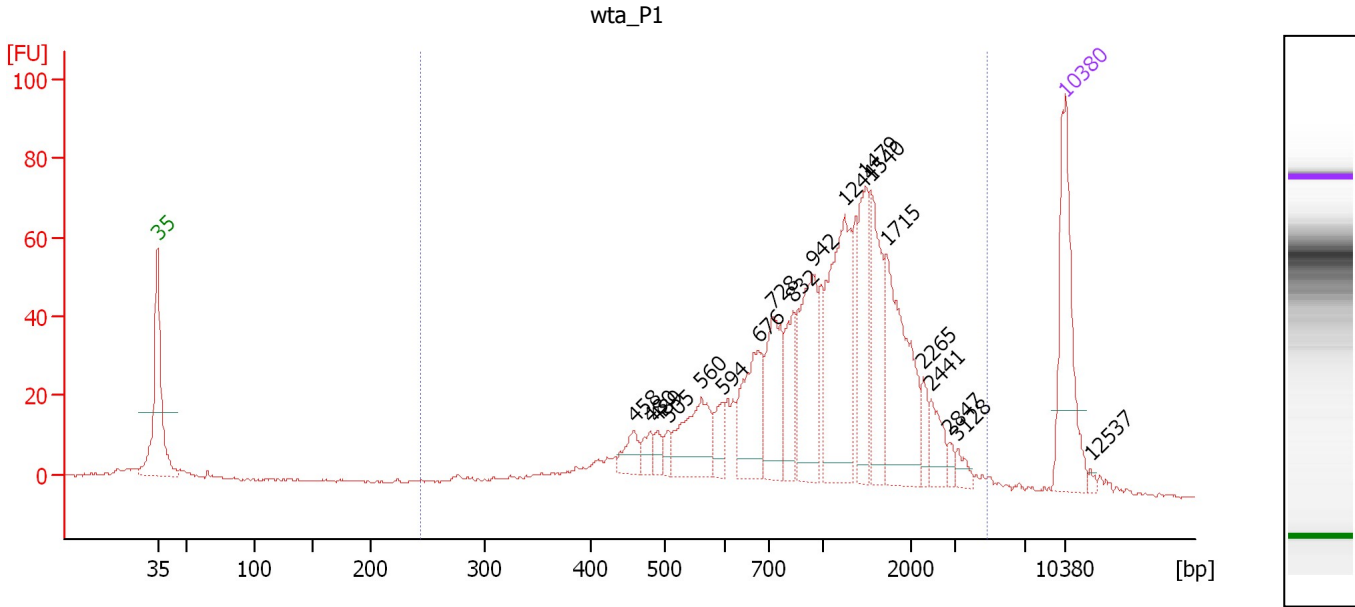
wta_comb

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
258	3,325	1,099	878.6	1,377.4	711.94	93	51.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
 Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...



Overall Results for sample 3 : wta_P1

Number of peaks found: 19 Corr. Area 1: 1,042.9
 Noise: 0.3

Peak table for sample 3 : wta_P1

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	458	18.94	62.6		79.76
3	480	12.97	41.0		81.00
4	489	9.48	29.4		81.55
5	505	8.96	26.9		82.42
6	560	58.10	157.1		84.99
7	594	16.91	43.2		86.55
8	676	66.08	148.2		89.35
9	728	65.36	136.1		90.54
10	832	38.89	70.8		92.00
11	942	79.84	128.4		93.56
12	1,244	119.53	145.6		96.04
13	1,479	54.53	55.9		97.64
14	1,540	58.09	57.2		98.06
15	1,715	95.76	84.6		99.25
16	2,265	13.12	8.8		102.09
17	2,441	18.55	11.5		102.69
18	2,847	4.79	2.5		104.06
19	3,128	7.10	3.4		104.75
20	10,380	75.00	10.9	Upper Marker	113.00
21	12,537	0.00	0.0		114.97

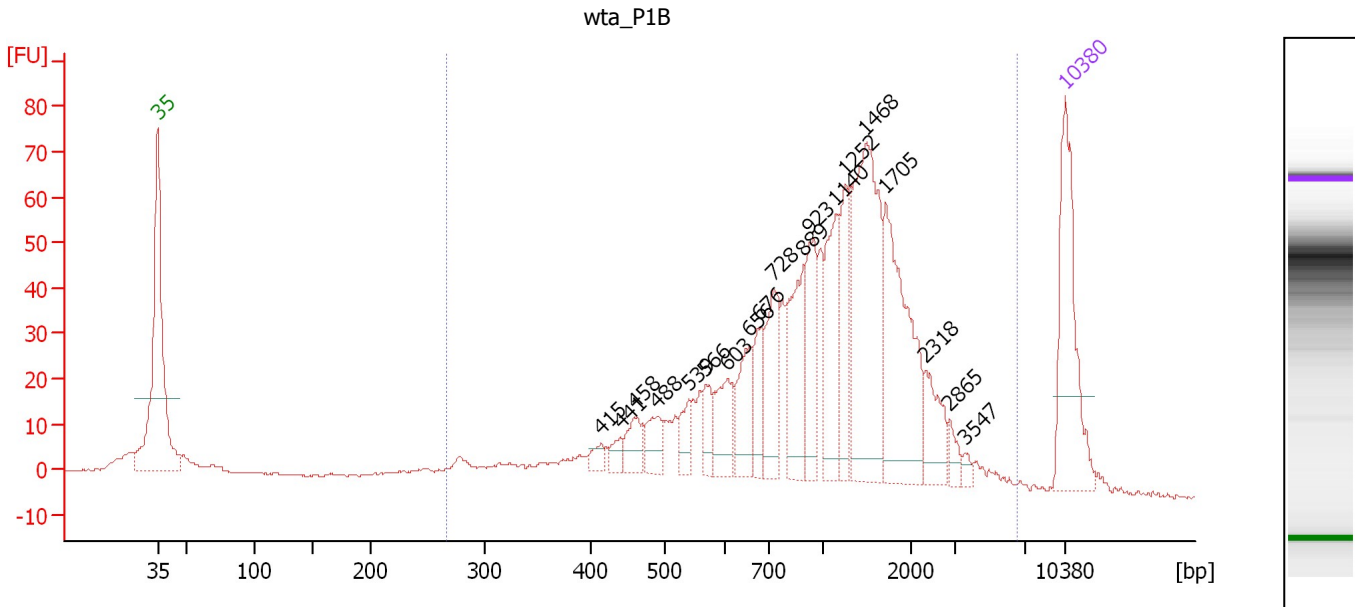
Region table for sample 3 : wta_P1

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
244	4,829	1,196	1,042.9	1,735.7	952.05	95	54.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
 Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...



Overall Results for sample 4 : wta_P1B

Number of peaks found: 19 Corr. Area 1: 1,088.7
 Noise: 0.2

Peak table for sample 4 : wta_P1B

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	415	8.68	31.7		77.24
3	441	9.39	32.3		78.76
4	458	19.04	63.0		79.76
5	488	21.11	65.5		81.51
6	539	16.93	47.6		83.98
7	566	18.29	49.0		85.27
8	603	32.22	81.0		86.93
9	656	37.62	86.9		88.69
10	676	27.26	61.1		89.35
11	728	49.21	102.5		90.54
12	889	61.44	104.7		92.82
13	923	43.12	70.8		93.29
14	1,140	56.97	75.7		95.33
15	1,252	40.58	49.1		96.09
16	1,468	137.26	141.7		97.57
17	1,705	103.51	92.0		99.18
18	2,318	27.41	17.9		102.27
19	2,865	7.19	3.8		104.12
20	3,547	3.99	1.7		105.31
21	10,380	75.00	10.9	Upper Marker	113.00

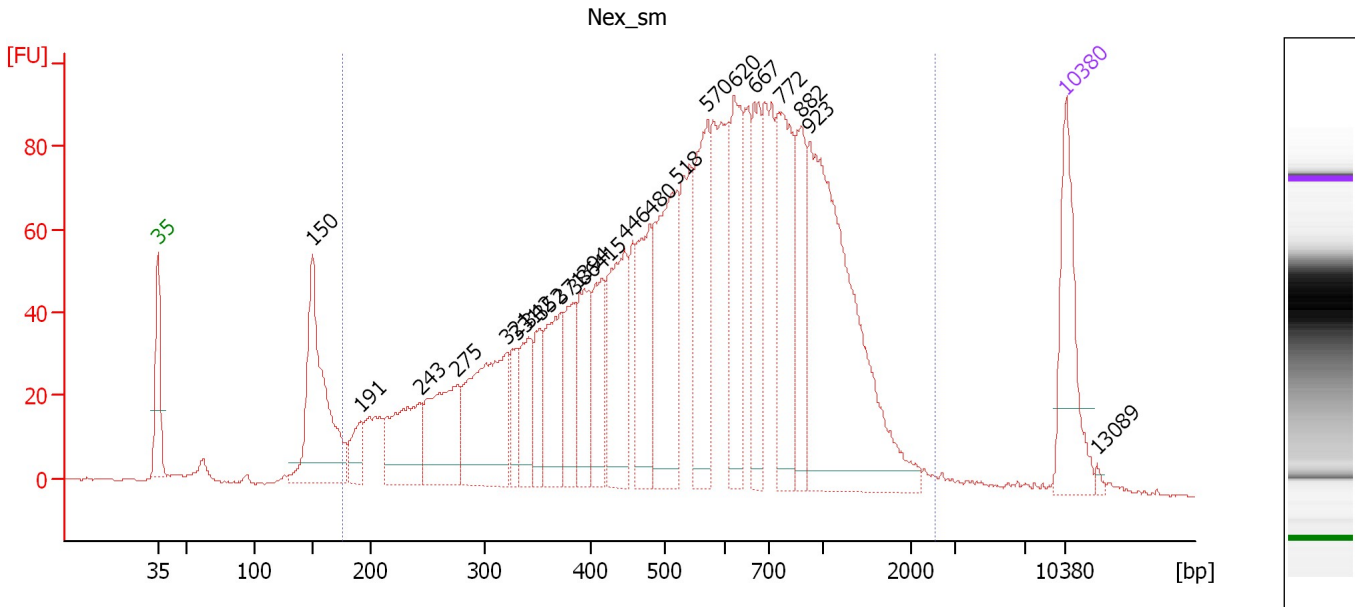
Region table for sample 4 : wta_P1B

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
266	6,597	1,223	1,088.7	1,800.9	946.09	94	63.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
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Electropherogram Summary Continued ...



Overall Results for sample 5 : Nex_sm

Number of peaks found: 22 Corr. Area 1: 2,616.4
 Noise: 0.3

Peak table for sample 5 : Nex_sm

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	150	203.50	2,061.0		54.94
3	191	32.65	258.6		58.68
4	243	93.77	584.8		63.24
5	275	108.10	594.5		66.11
6	321	152.95	722.8		69.94
7	331	32.95	150.9		70.77
8	342	50.77	225.2		71.64
9	352	42.63	183.3		72.51
10	371	83.12	339.1		74.07
11	386	54.62	214.5		75.23
12	394	66.22	254.6		75.91
13	415	62.12	226.7		77.27
14	446	113.88	386.7		79.07
15	480	90.19	284.9		81.01
16	518	155.79	455.3		83.05
17	570	118.78	316.0		85.43
18	620	98.20	239.9		87.51
19	667	80.45	182.7		89.07
20	772	120.71	237.0		91.16
21	882	75.39	129.6		92.71
22	923	307.45	504.7		93.29
23	10,380	75.00	10.9	Upper Marker	113.00
24	13,089	0.00	0.0		115.48


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...

... Region table for sample 5 :

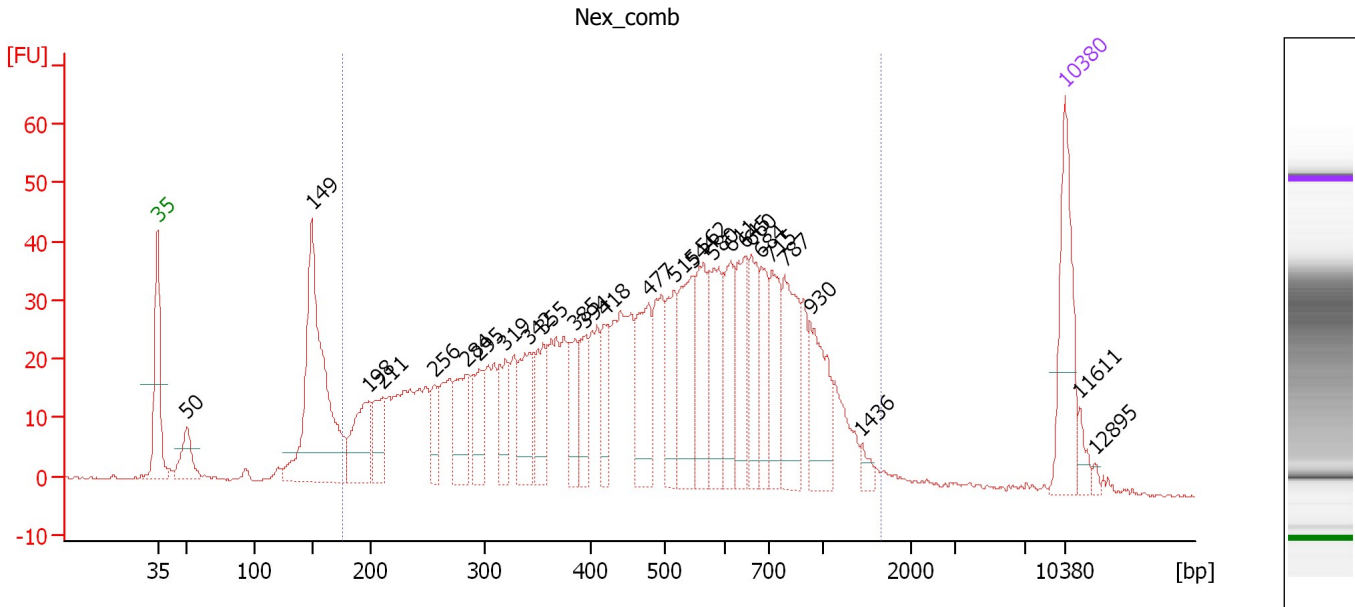
Nex_sm

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
176	2,537	638	2,616.4	8,202.9	 2,449.83	92	53.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
 Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...



Overall Results for sample 6 : Nex_comb

Number of peaks found: 28 Corr. Area 1: 1,272.6
 Noise: 0.3

Peak table for sample 6 : Nex_comb

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	50	43.54	1,317.6		45.25
3	149	295.70	3,007.0		54.89
4	198	74.20	567.7		59.29
5	211	46.45	333.8		60.41
6	256	31.21	184.8		64.38
7	284	61.88	330.6		66.82
8	295	43.37	222.6		67.85
9	319	46.81	222.3		69.81
10	342	65.26	289.2		71.67
11	355	53.37	227.7		72.74
12	385	37.21	146.4		75.19
13	394	37.28	143.3		75.92
14	418	39.39	142.8		77.44
15	477	76.39	242.6		80.86
16	515	58.54	172.4		82.87
17	544	84.87	236.4		84.24
18	562	65.46	176.5		85.07
19	580	72.89	190.5		85.90
20	611	56.55	140.1		87.22
21	645	64.55	151.5		88.35
22	660	58.01	133.1		88.84
23	684	43.62	96.6		89.62
24	715	57.28	121.5		90.35
25	787	83.71	161.1		91.38

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
 Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...

... Peak table for sample 6 : Nex_comb

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	930	61.72	100.6		93.38
27	1,436	9.09	9.6		97.35
28	10,380	75.00	10.9	Upper Marker	113.00
29	11,611	0.00	0.0		114.13
30	12,895	0.00	0.0		115.30

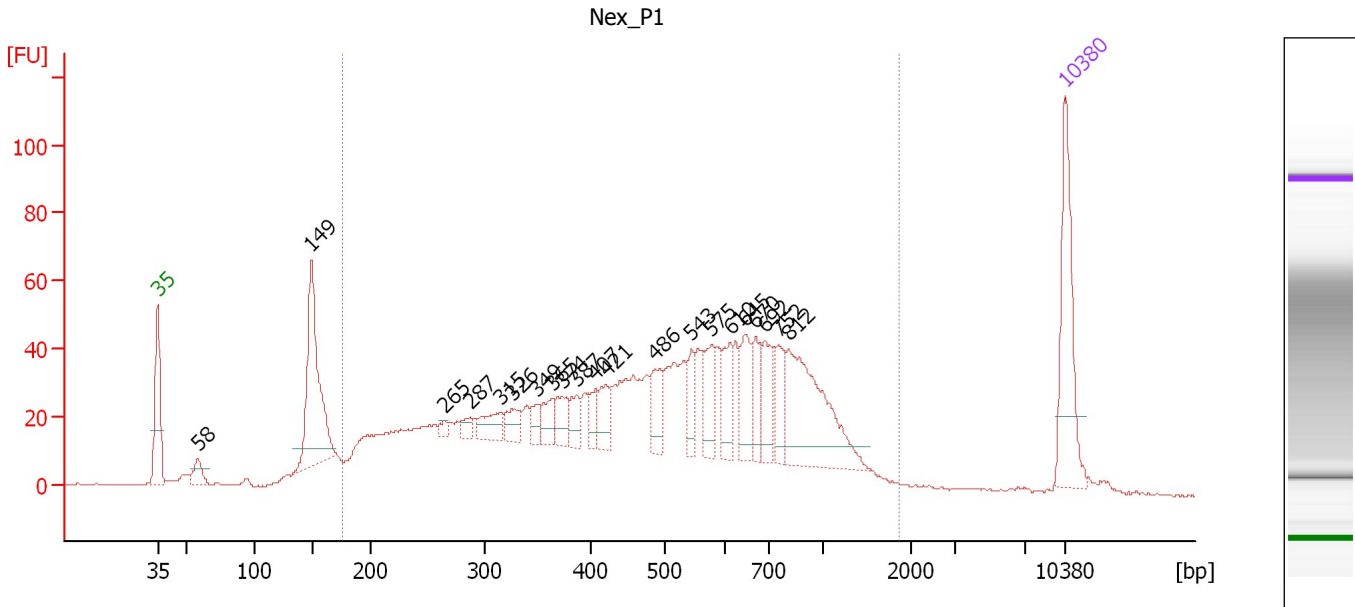
Region table for sample 6 : Nex_comb

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
176	1,643	531	1,272.6	8,129.7	2,102.51	87	47.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
 Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...



Overall Results for sample 7 : Nex_P1

Number of peaks found: 21 Corr. Area 1: 1,456.0
 Noise: 0.2

Peak table for sample 7 : Nex_P1

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	58	20.66	536.8		46.12
3	149	165.18	1,681.5		54.87
4	265	6.02	34.5		65.16
5	287	10.31	54.4		67.14
6	315	25.65	123.2		69.52
7	326	18.97	88.3		70.36
8	349	12.44	54.0		72.24
9	365	19.89	82.6		73.52
10	374	21.80	88.3		74.27
11	387	18.71	73.2		75.35
12	407	15.01	55.9		76.79
13	421	24.88	89.5		77.63
14	486	30.05	93.6		81.39
15	543	23.40	65.2		84.21
16	575	33.30	87.7		85.69
17	610	31.15	77.4		87.18
18	645	37.39	87.9		88.31
19	670	27.57	62.4		89.16
20	692	36.61	80.1		89.90
21	752	31.22	62.9		90.89
22	812	123.87	231.1		91.73
23	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...

... Region table for sample 7 :

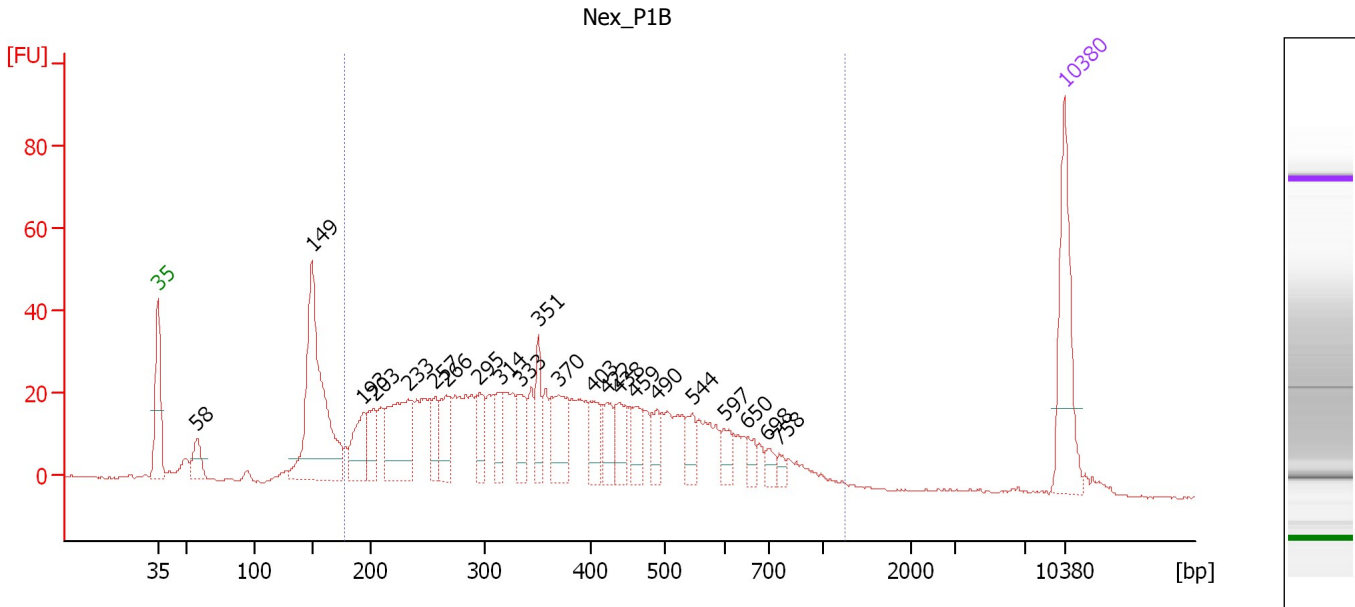
Nex P1

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
175	1,865	548	1,456.0	6,051.5	1,581.78	86	49.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
 Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...



Overall Results for sample 8 : Nex_P1B

Number of peaks found: 22 Corr. Area 1: 881.5
 Noise: 0.2

Peak table for sample 8 : Nex_P1B

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	58	27.26	714.5		46.06
3	149	238.21	2,419.1		54.91
4	193	48.59	380.9		58.86
5	203	33.26	248.6		59.70
6	233	100.17	651.5		62.36
7	257	32.06	189.4		64.44
8	266	43.31	246.6		65.28
9	295	33.01	169.4		67.85
10	314	32.23	155.3		69.43
11	333	30.07	136.7		70.96
12	351	34.57	149.3		72.39
13	370	55.01	225.5		73.92
14	403	28.88	108.5		76.59
15	422	26.30	94.4		77.68
16	438	27.46	95.1		78.57
17	459	27.57	91.0		79.80
18	490	23.98	74.2		81.58
19	544	19.84	55.2		84.25
20	597	17.18	43.6		86.72
21	650	12.23	28.5		88.50
22	698	9.45	20.5		90.08
23	758	7.28	14.5		90.97
24	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...

... Region table for sample 8 :

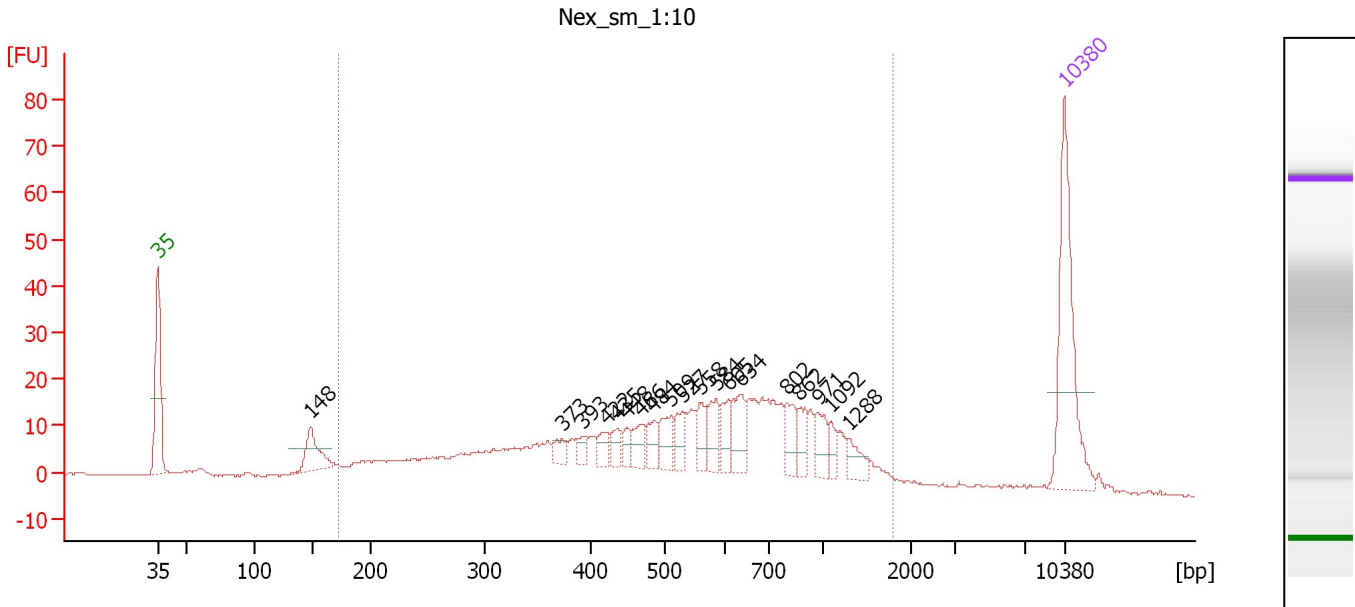
Nex P1B

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
178	1,246	400	881.5	5,662.6	1,206.87	81	41.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
 Modified: 4/10/2017 11:45:00 AM

Electropherogram Summary Continued ...



Overall Results for sample 9 : Nex sm 1:10

Number of peaks found: 19 Corr. Area 1: 551.1
 Noise: 0.2

Peak table for sample 9 : Nex sm 1:10

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	148	33.36	341.5		54.80
3	373	8.71	35.3		74.23
4	393	8.58	33.1		75.82
5	422	10.35	37.2		77.65
6	435	9.60	33.4		78.45
7	448	8.10	27.4		79.19
8	466	14.09	45.8		80.23
9	484	13.86	43.4		81.27
10	509	18.24	54.3		82.61
11	527	15.37	44.2		83.45
12	558	15.12	41.1		84.89
13	584	19.27	50.0		86.08
14	605	14.35	35.9		87.02
15	634	23.65	56.5		87.96
16	802	17.33	32.7		91.58
17	862	13.40	23.6		92.43
18	971	15.74	24.6		93.96
19	1,092	7.63	10.6		95.00
20	1,288	10.57	12.4		96.34
21	10,380	75.00	10.9	Upper Marker	113.00

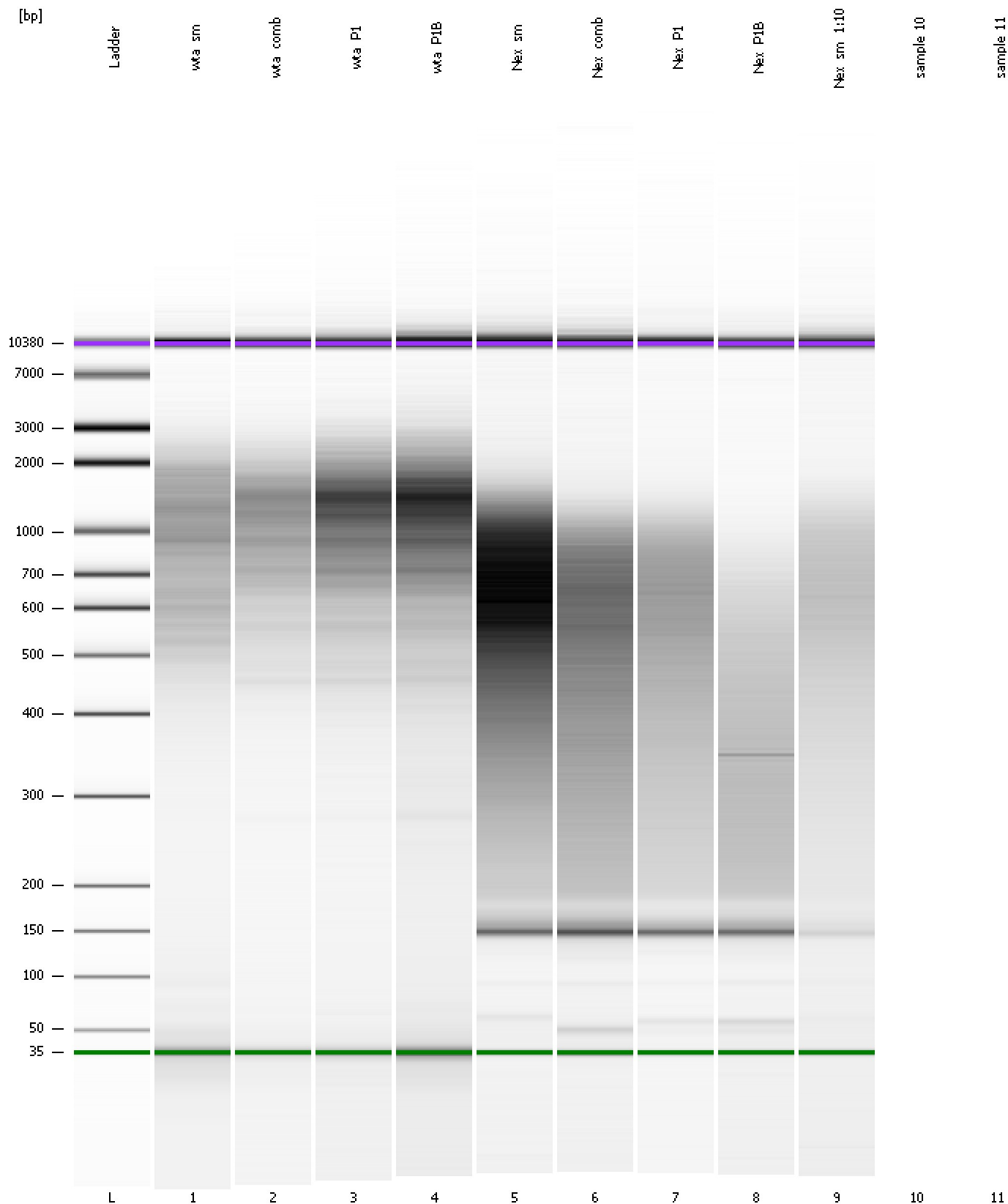
Region table for sample 9 : Nex sm 1:10

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
172	1,787	609	551.1	2,357.3	674.61	88	50.4

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
Modified: 4/10/2017 11:45:00 AM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
Modified: 4/10/2017 11:45:00 AM

Invalid Samples

Sample 10 has not been run, no results available.

Sample 11 has not been run, no results available.

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad

Created: 4/10/2017 11:01:19 AM
 Modified: 4/10/2017 11:45:00 AM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 10)		Instrument	Run		4/10/2017 11:36:55 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2017-04-10\2017-04-10_001.xad)		Instrument	Run		4/10/2017 11:01:24 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		4/10/2017 11:01:24 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		4/10/2017 11:01:24 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		4/10/2017 11:01:24 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		4/10/2017 11:01:24 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		4/10/2017 11:01:24 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		4/10/2017 11:01:24 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1