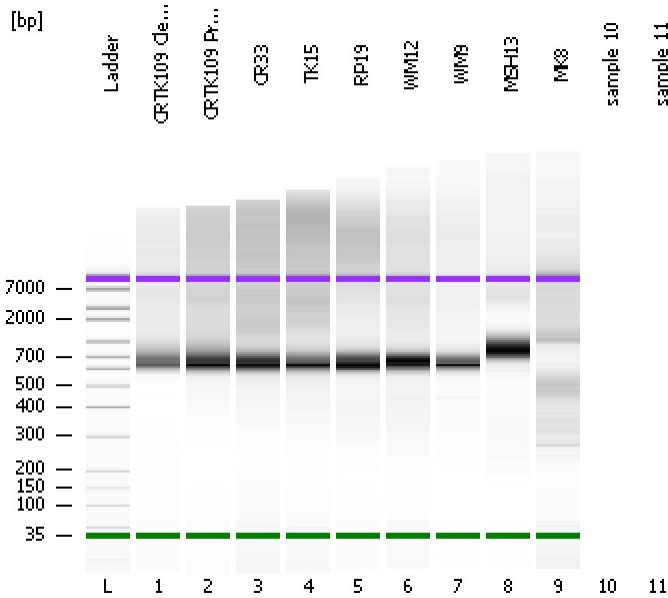


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 AM

Electrophoresis File Run Summary



Instrument Information:

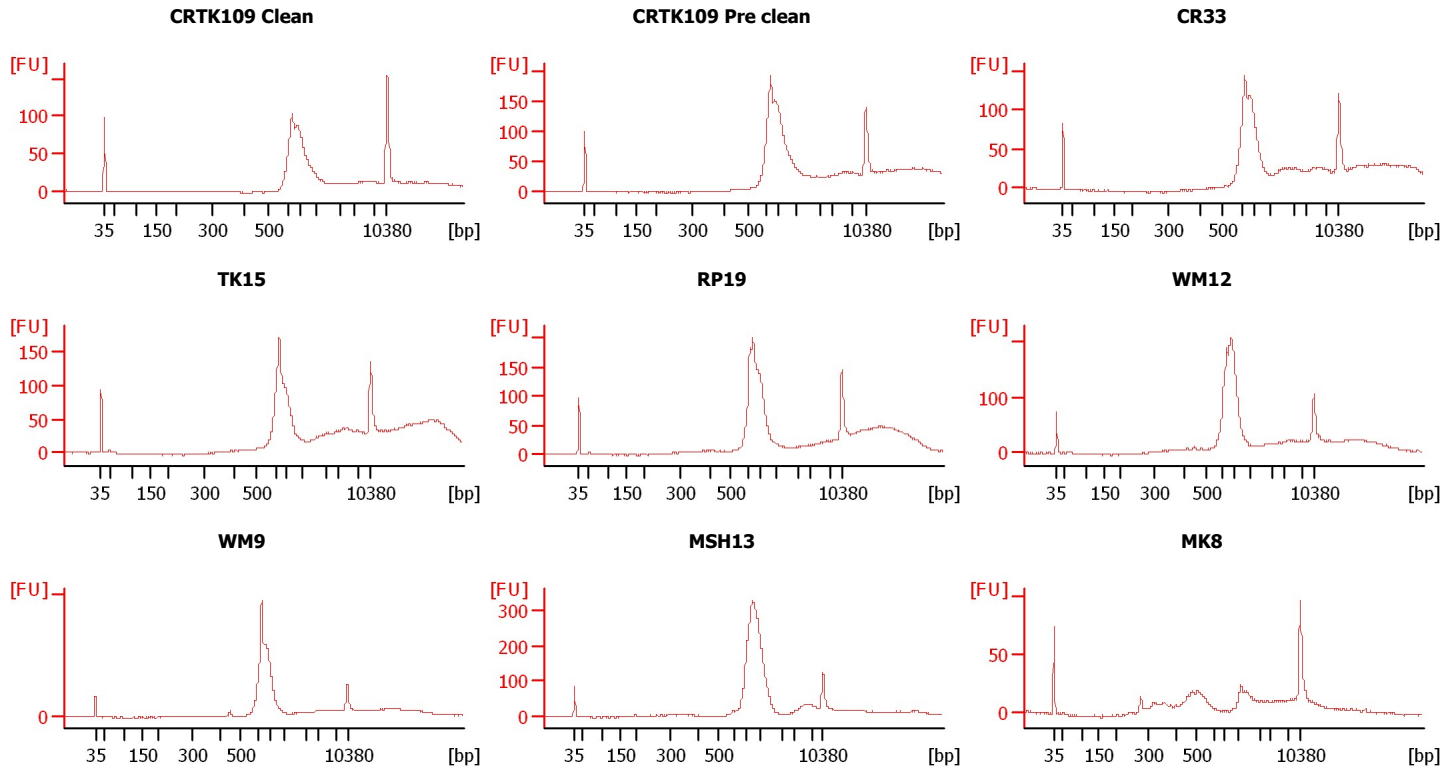
Instrument Name: DE34903152 Firmware: C.01.069
Serial#: DE34903152 Type: G2938C

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:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
CRTK109 Clean		<input type="checkbox"/>	✓			
CRTK109 Pre clean		<input type="checkbox"/>	✓			
CR33		<input type="checkbox"/>	✓			
TK15		<input type="checkbox"/>	✓			
RP19		<input type="checkbox"/>	✓			
WM12		<input type="checkbox"/>	✓			
WM9		<input type="checkbox"/>	✓			
MSH13		<input type="checkbox"/>	✓			
MK8		<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\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 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

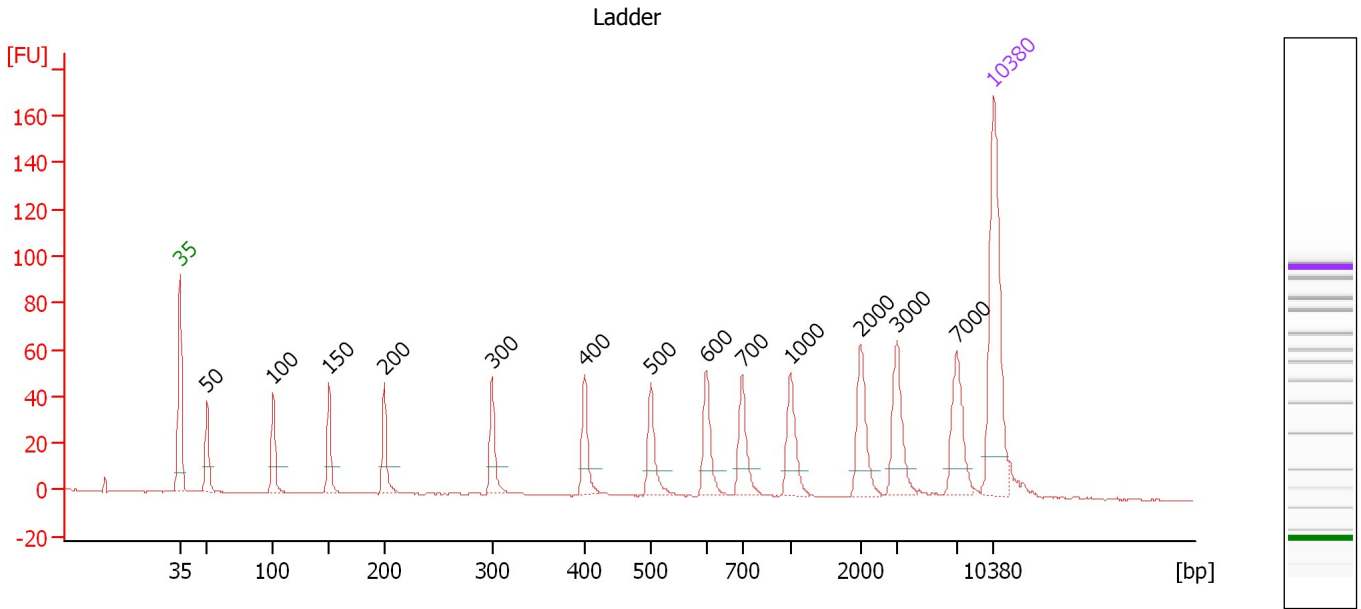
Ladder

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\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
 Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

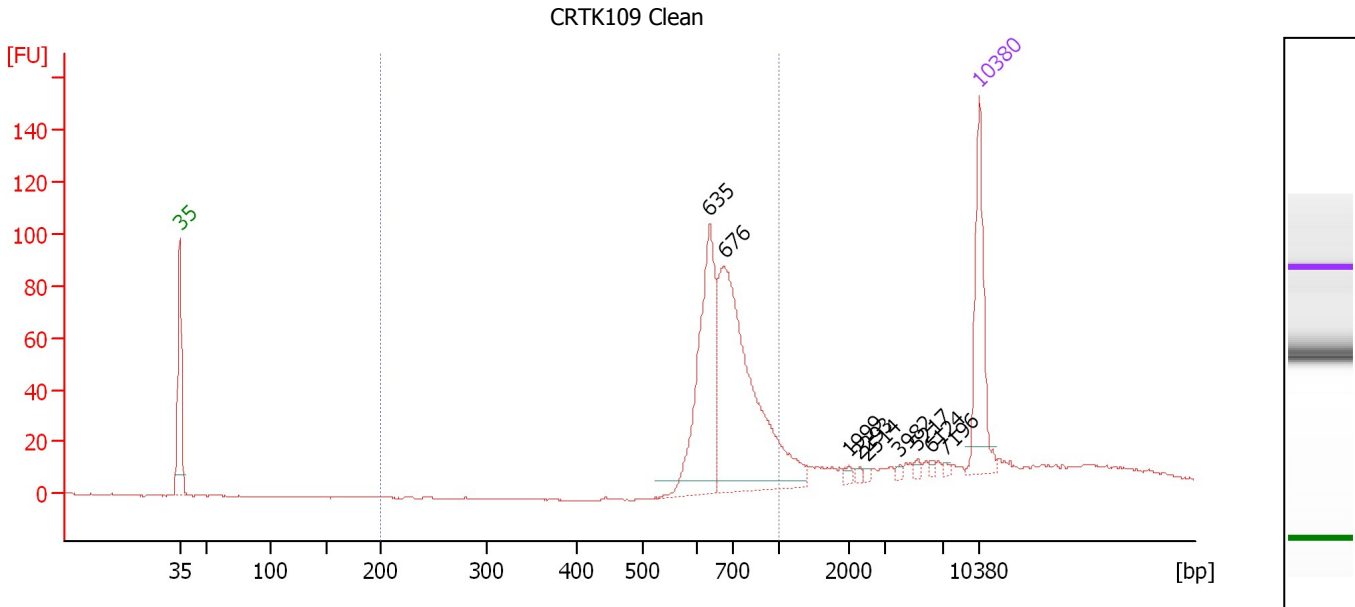
Peak table for Ladder

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	50	150.00	4,545.5	Ladder Peak	45.34
3	100	150.00	2,272.7	Ladder Peak	51.03
4	150	150.00	1,515.2	Ladder Peak	55.81
5	200	150.00	1,136.4	Ladder Peak	60.58
6	300	150.00	757.6	Ladder Peak	69.85
7	400	150.00	568.2	Ladder Peak	77.84
8	500	150.00	454.5	Ladder Peak	83.53
9	600	150.00	378.8	Ladder Peak	88.30
10	700	150.00	324.7	Ladder Peak	91.38
11	1,000	150.00	227.3	Ladder Peak	95.51
12	2,000	150.00	113.6	Ladder Peak	101.57
13	3,000	150.00	75.8	Ladder Peak	104.74
14	7,000	150.00	32.5	Ladder Peak	109.88
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
 Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : CRTK109 Clean

Number of peaks found: 9 Corr. Area 1: 491.9
 Noise: 0.2

Peak table for sample 1 : CRTK109 Clean

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	635	178.19	425.1		89.39
3	676	299.24	670.5		90.65
4	1,999	3.70	2.8		101.57
5	2,293	2.83	1.9		102.50
6	2,514	2.47	1.5		103.20
7	3,982	2.26	0.9		106.00
8	5,217	3.08	0.9		107.59
9	6,124	2.63	0.7		108.75
10	7,196	2.18	0.5		110.06
11	10,380	75.00	10.9	Upper Marker	113.00

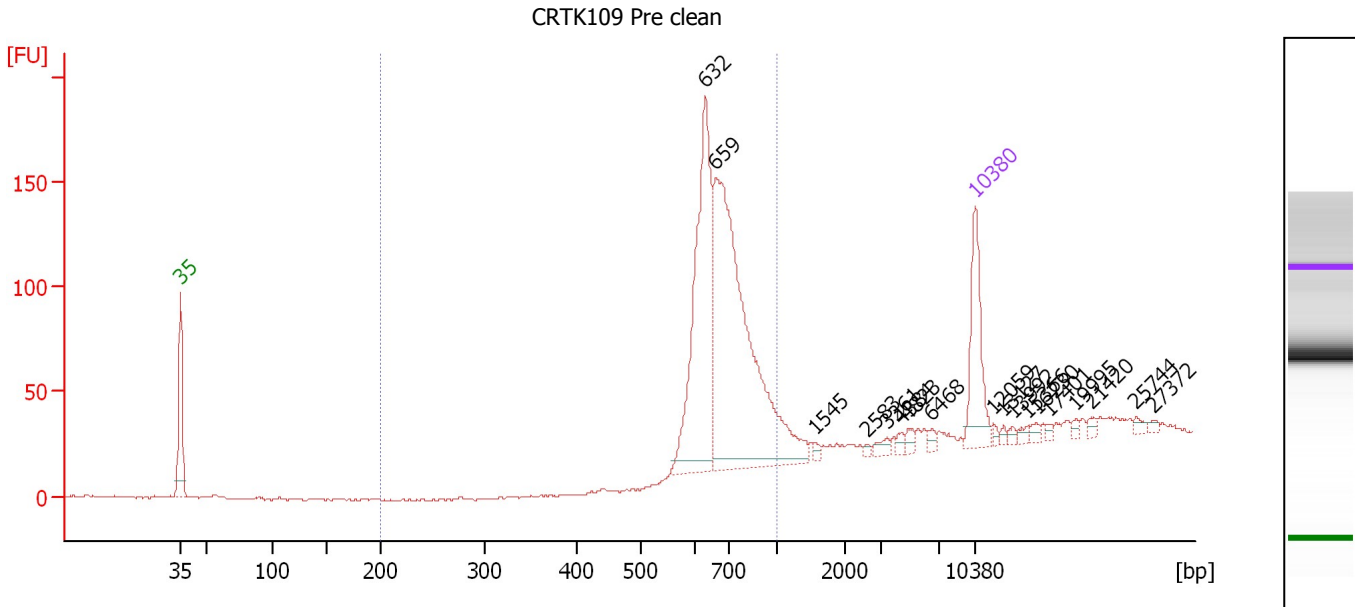
Region table for sample 1 : CRTK109 Clean

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	706	414.65	491.9	904.9	73	13.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
 Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 : CRTK109 Pre clean

Number of peaks found: 18 Corr. Area 1: 790.8
 Noise: 0.5

Peak table for sample 2 : CRTK109 Pre clean

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	632	309.32	742.1		89.28
3	659	499.88	1,149.2		90.12
4	1,545	4.50	4.4		98.81
5	2,583	2.65	1.6		103.42
6	3,361	7.54	3.4		105.20
7	4,384	5.25	1.8		106.52
8	4,823	7.03	2.2		107.08
9	6,468	5.22	1.2		109.19
10	10,380	75.00	10.9	Upper Marker	113.00
11	12,059	0.00	0.0		114.55
12	13,127	0.00	0.0		115.54
13	13,992	0.00	0.0		116.34
14	15,366	0.00	0.0		117.60
15	16,230	0.00	0.0		118.40
16	17,401	0.00	0.0		119.48
17	19,995	0.00	0.0		121.88
18	21,420	0.00	0.0		123.19
19	25,744	0.00	0.0		127.19
20	27,372	0.00	0.0		128.69

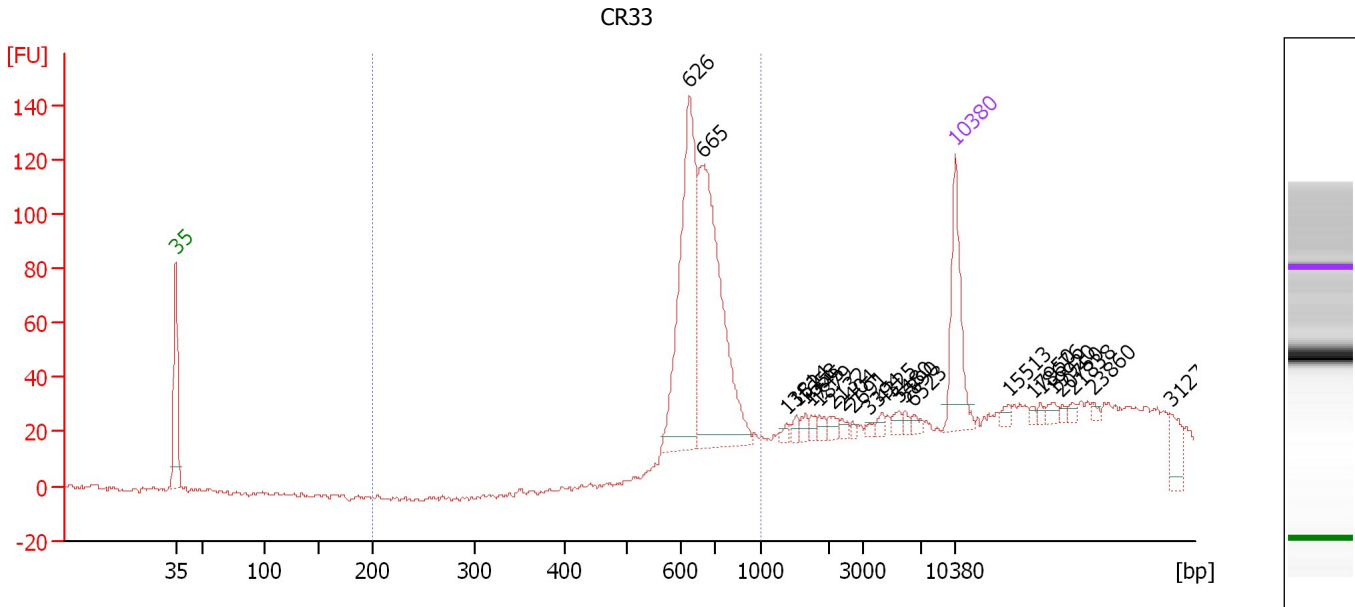
Region table for sample 2 : CRTK109 Pre clean

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	698	703.59	790.8	1,552.8	80	13.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
 Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...



Overall Results for sample 3 : CR33

Number of peaks found: 23 Corr. Area 1: 562.4
 Noise: 0.8

Peak table for sample 3 : CR33

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	626	278.46	673.8		89.11
3	665	322.80	735.4		90.31
4	1,381	5.10	5.6		97.82
5	1,524	6.61	6.6		98.68
6	1,658	6.99	6.4		99.50
7	1,745	6.46	5.6		100.02
8	1,879	6.38	5.1		100.84
9	2,132	7.86	5.6		101.99
10	2,404	5.17	3.3		102.85
11	2,691	3.00	1.7		103.76
12	3,394	3.86	1.7		105.24
13	4,325	5.58	2.0		106.44
14	5,480	7.05	1.9		107.92
15	5,890	4.27	1.1		108.45
16	6,523	5.35	1.2		109.27
17	10,380	75.00	10.9	Upper Marker	113.00
18	15,513	0.00	0.0		117.74
19	17,950	0.00	0.0		119.99
20	18,676	0.00	0.0		120.66
21	19,920	0.00	0.0		121.81
22	20,750	0.00	0.0		122.58
23	21,838	0.00	0.0		123.58
24	23,860	0.00	0.0		125.45
25	31,275	0.00	0.0		132.30

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...

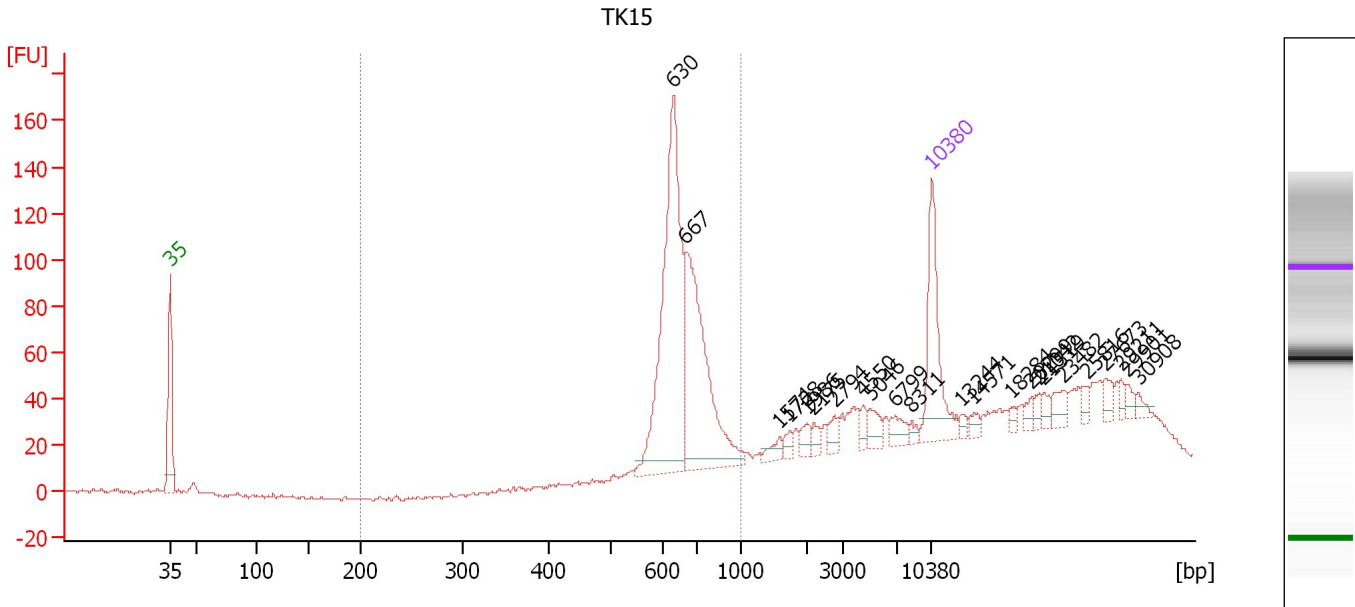
... Region table for sample 3 : CR33

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	674	620.39	562.4	1,407.8	 66	11.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
 Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...



Overall Results for sample 4 : TK15

Number of peaks found: 23 Corr. Area 1: 588.5
 Noise: 0.7

Peak table for sample 4 : TK15


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	630	290.30	697.6		89.24
3	667	204.08	463.5		90.37
4	1,574	8.83	8.5		98.99
5	1,728	6.87	6.0		99.92
6	1,986	9.09	6.9		101.49
7	2,175	7.65	5.3		102.13
8	2,794	9.29	5.0		104.08
9	4,550	8.52	2.8		106.73
10	5,046	13.12	3.9		107.37
11	6,799	12.59	2.8		109.62
12	8,311	4.59	0.8		111.09
13	10,380	75.00	10.9	Upper Marker	113.00
14	13,244	0.00	0.0		115.65
15	14,571	0.00	0.0		116.87
16	18,284	0.00	0.0		120.30
17	20,299	0.00	0.0		122.16
18	21,042	0.00	0.0		122.85
19	21,519	0.00	0.0		123.29
20	23,482	0.00	0.0		125.10
21	25,816	0.00	0.0		127.25
22	27,673	0.00	0.0		128.97
23	29,211	0.00	0.0		130.39
24	29,901	0.00	0.0		131.03
25	30,908	0.00	0.0		131.96

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...

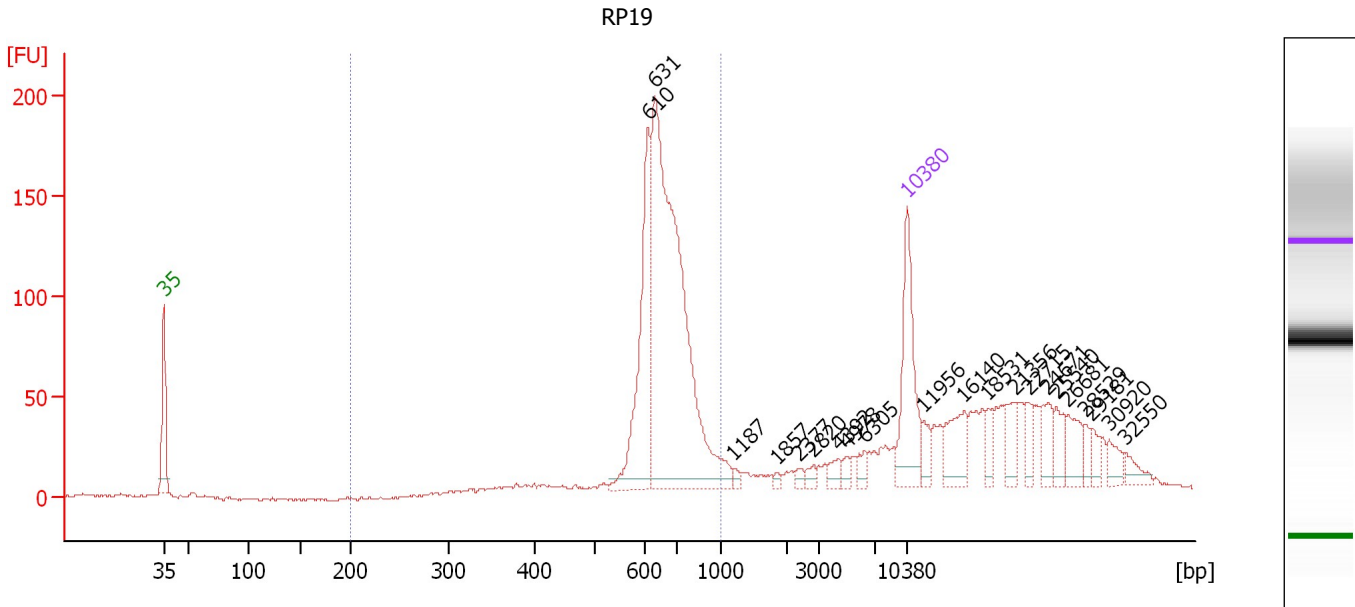
... Region table for sample 4 : TK15

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	672	485.39	588.5	1,106.9	 47	11.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
 Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : RP19

Number of peaks found: 21 Corr. Area 1: 970.9
 Noise: 0.8

Peak table for sample 5 : RP19


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	610	153.15	380.4		88.61
3	631	445.55	1,069.5		89.27
4	1,187	3.95	5.0		96.64
5	1,857	3.05	2.5		100.71
6	2,377	4.14	2.6		102.76
7	2,820	5.61	3.0		104.17
8	4,392	8.24	2.8		106.53
9	4,978	5.90	1.8		107.28
10	6,305	8.65	2.1		108.99
11	10,380	75.00	10.9	Upper Marker	113.00
12	11,956	0.00	0.0		114.46
13	16,140	0.00	0.0		118.32
14	18,531	0.00	0.0		120.53
15	21,356	0.00	0.0		123.14
16	22,715	0.00	0.0		124.39
17	24,671	0.00	0.0		126.20
18	25,540	0.00	0.0		127.00
19	26,681	0.00	0.0		128.05
20	28,529	0.00	0.0		129.76
21	29,181	0.00	0.0		130.36
22	30,920	0.00	0.0		131.97
23	32,550	0.00	0.0		133.47

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...

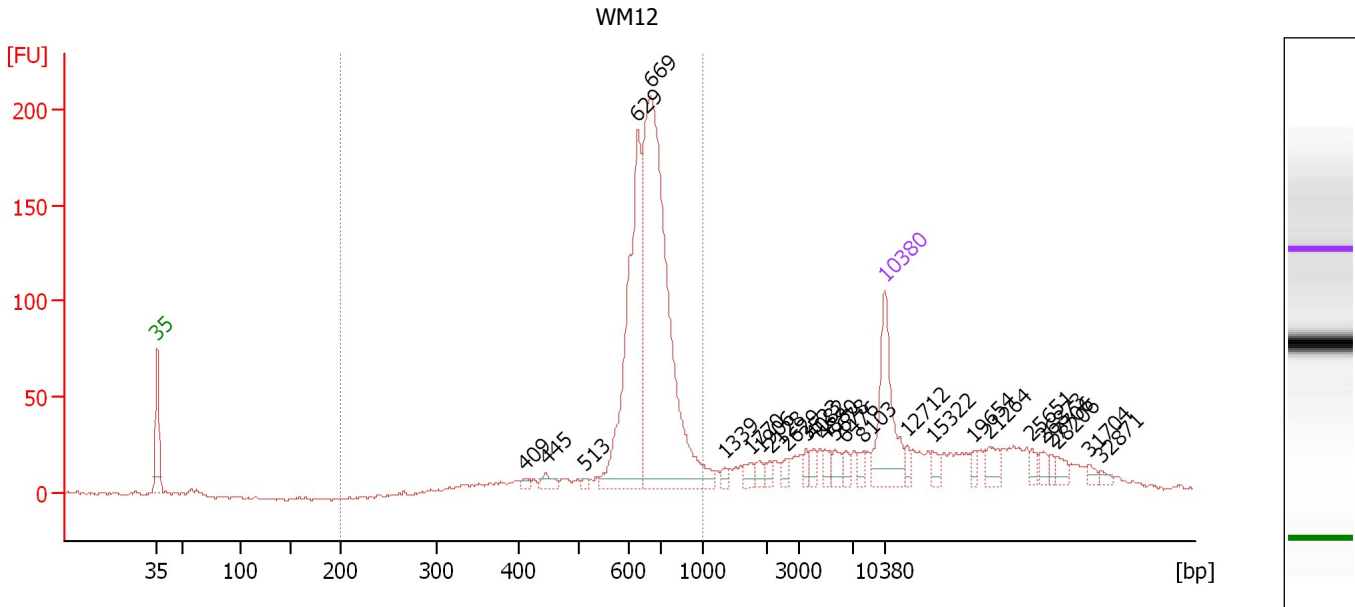
... Region table for sample 5 : RP19

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	665	617.75	970.9	1,447.1	 56	14.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
 Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...



Overall Results for sample 6 : WM12

Number of peaks found: 26 Corr. Area 1: 1,058.1
 Noise: 0.7

Peak table for sample 6 : WM12

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	409	5.64	20.9		78.33
3	445	11.72	39.9		80.38
4	513	3.13	9.3		84.13
5	629	249.94	601.8		89.21
6	669	486.38	1,101.1		90.44
7	1,339	4.80	5.4		97.56
8	1,770	9.08	7.8		100.18
9	1,906	6.68	5.3		101.00
10	2,128	5.61	4.0		101.97
11	2,629	6.95	4.0		103.56
12	3,523	7.11	3.1		105.41
13	4,082	8.54	3.2		106.13
14	4,880	8.43	2.6		107.15
15	5,678	10.79	2.9		108.18
16	6,476	7.88	1.8		109.21
17	8,103	8.22	1.5		110.90
18	10,380	75.00	10.9	Upper Marker	113.00
19	12,712	0.00	0.0		115.15
20	15,322	0.00	0.0		117.56
21	19,654	0.00	0.0		121.56
22	21,264	0.00	0.0		123.05
23	25,651	0.00	0.0		127.10
24	26,873	0.00	0.0		128.23
25	27,706	0.00	0.0		129.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...**... Peak table for sample 6 : WM12**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	28,206	0.00	0.0		129.46
27	31,704	0.00	0.0		132.69
28	32,871	0.00	0.0		133.77

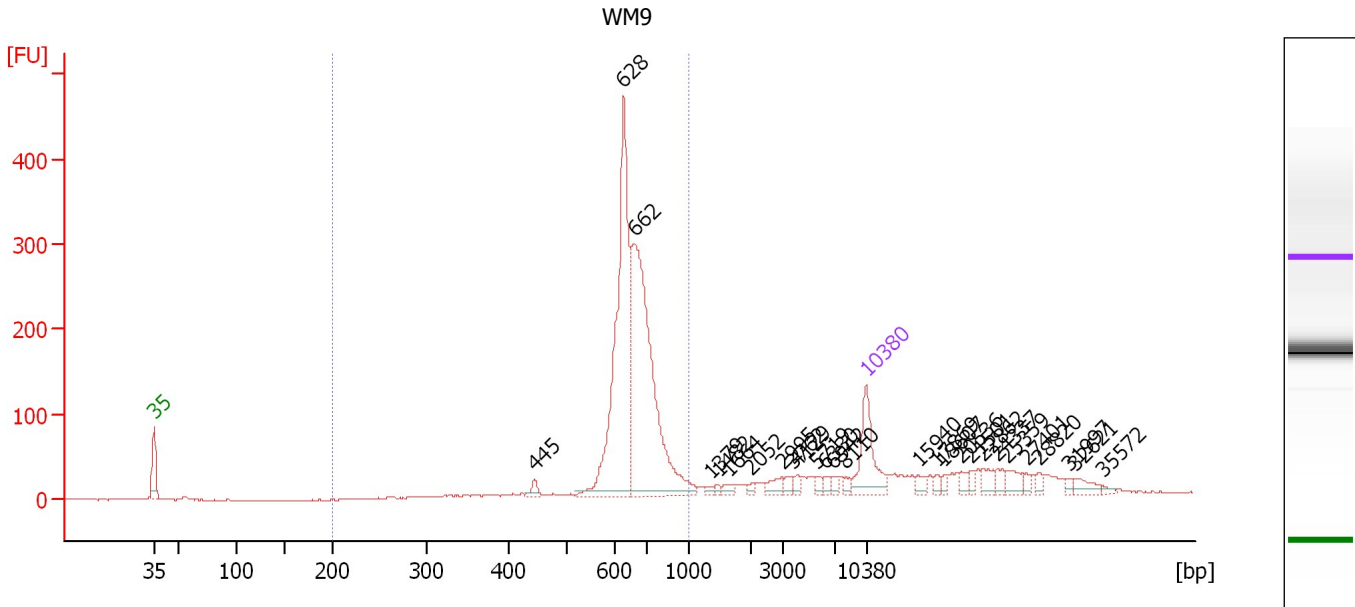
Region table for sample 6 : WM12

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	662	801.93	1,058.1	1,907.0	 66	15.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
 Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...



Overall Results for sample 7 : WM9

Number of peaks found: 27 Corr. Area 1: 1,593.0
 Noise: 1.0

Peak table for sample 7 : WM9

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	445	12.11	41.2		80.41
3	628	465.51	1,123.5		89.16
4	662	498.38	1,140.9		90.21
5	1,379	5.07	5.6		97.81
6	1,482	4.29	4.4		98.43
7	1,664	8.70	7.9		99.53
8	2,052	4.85	3.6		101.74
9	2,995	16.66	8.4		104.72
10	3,762	10.35	4.2		105.72
11	4,129	7.37	2.7		106.19
12	5,719	7.55	2.0		108.23
13	6,330	7.35	1.8		109.02
14	6,942	7.34	1.6		109.80
15	8,110	7.18	1.3		110.90
16	10,380	75.00	10.9	Upper Marker	113.00
17	15,940	0.00	0.0		118.13
18	17,869	0.00	0.0		119.92
19	18,607	0.00	0.0		120.60
20	20,536	0.00	0.0		122.38
21	21,501	0.00	0.0		123.27
22	22,862	0.00	0.0		124.53
23	24,337	0.00	0.0		125.89
24	25,359	0.00	0.0		126.83
25	27,401	0.00	0.0		128.72

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...**... Peak table for sample 7 : WM9**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	28,820	0.00	0.0		130.03
27	31,997	0.00	0.0		132.96
28	32,621	0.00	0.0		133.54
29	35,572	0.00	0.0		136.26

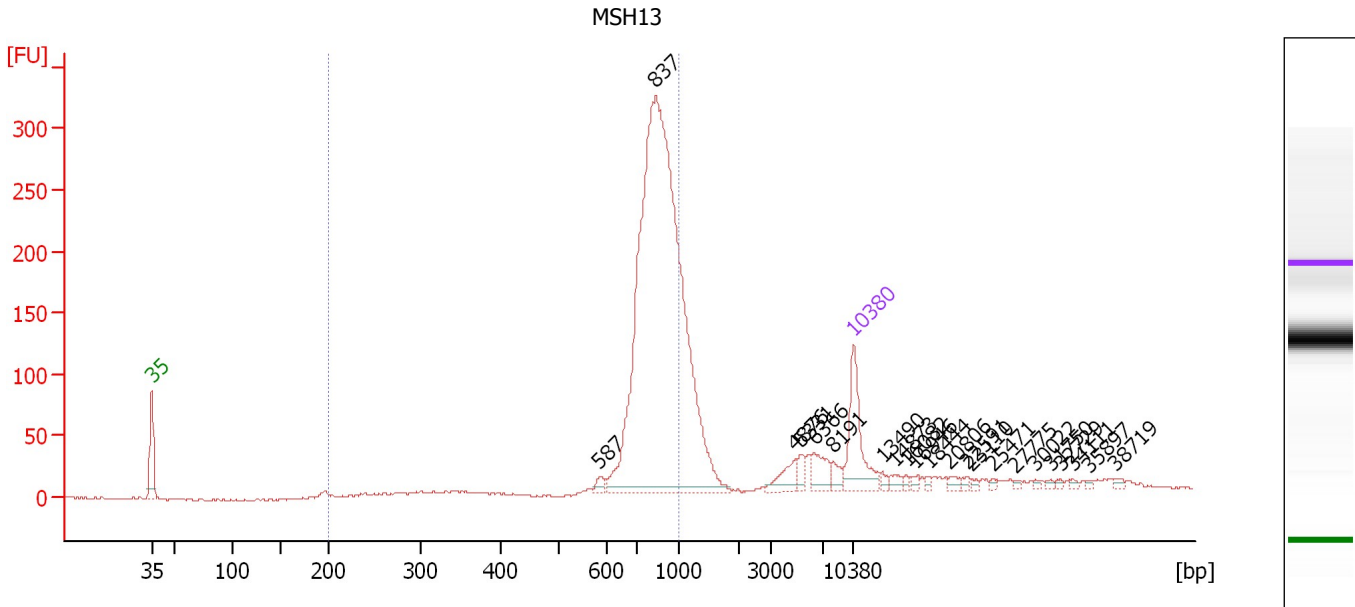
Region table for sample 7 : WM9

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	663	990.07	1,593.0	2,316.8	 71	13.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
 Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...



Overall Results for sample 8 : MSH13

Number of peaks found: 22 Corr. Area 1: 1,397.1
 Noise: 0.8

Peak table for sample 8 : MSH13

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	587	10.79	27.9		87.68
3	837	1,156.81	2,094.2		93.27
4	4,876	24.40	7.6		107.15
5	5,331	12.75	3.6		107.73
6	6,366	28.57	6.8		109.06
7	8,191	11.99	2.2		110.98
8	10,380	75.00	10.9	Upper Marker	113.00
9	13,490	0.00	0.0		115.87
10	14,873	0.00	0.0		117.15
11	16,082	0.00	0.0		118.27
12	16,946	0.00	0.0		119.06
13	18,444	0.00	0.0		120.45
14	20,806	0.00	0.0		122.63
15	22,591	0.00	0.0		124.28
16	23,110	0.00	0.0		124.76
17	25,471	0.00	0.0		126.94
18	27,775	0.00	0.0		129.06
19	30,022	0.00	0.0		131.14
20	31,750	0.00	0.0		132.73
21	32,729	0.00	0.0		133.64
22	34,111	0.00	0.0		134.91
23	35,897	0.00	0.0		136.56
24	38,719	0.00	0.0		139.17

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 AM

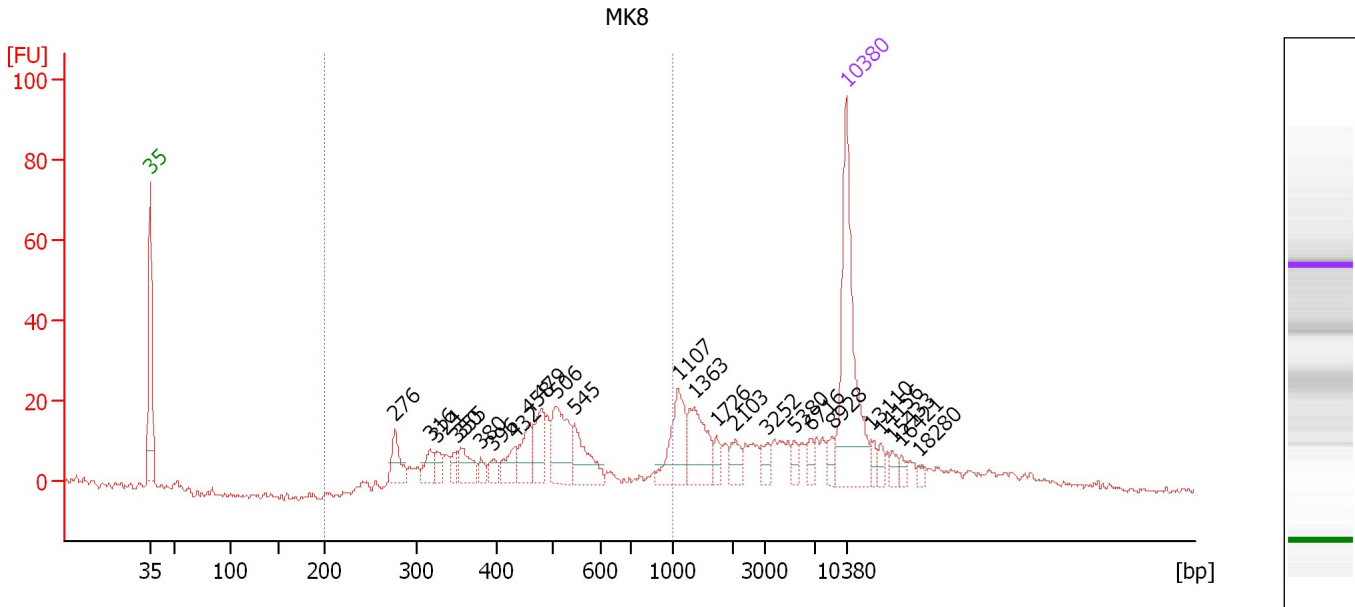
Electropherogram Summary Continued ...**... Region table for sample 8 : MSH13**

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	817	939.61	1,397.1	1,803.0	 67	13.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
 Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...



Overall Results for sample 9 : MK8

Number of peaks found: 25 Corr. Area 1: 268.9
 Noise: 0.7

Peak table for sample 9 : MK8

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	276	19.58	107.5		67.62
3	316	11.89	57.1		71.11
4	324	8.79	41.0		71.80
5	350	6.68	28.9		73.84
6	355	17.50	74.8		74.22
7	380	5.89	23.5		76.26
8	396	7.62	29.2		77.49
9	432	13.03	45.7		79.64
10	458	22.06	73.0		81.14
11	479	23.02	72.9		82.32
12	506	39.45	118.1		83.82
13	545	26.53	73.7		85.70
14	1,107	32.74	44.8		96.16
15	1,363	31.00	34.5		97.71
16	1,726	5.75	5.0		99.91
17	2,103	9.01	6.5		101.90
18	3,252	6.56	3.1		105.06
19	5,380	4.70	1.3		107.80
20	6,716	5.29	1.2		109.51
21	8,928	6.03	1.0		111.66
22	10,380	75.00	10.9	Upper Marker	113.00
23	13,110	0.00	0.0		115.52
24	14,156	0.00	0.0		116.49
25	15,433	0.00	0.0		117.67

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 AM

Electropherogram Summary Continued ...**... Peak table for sample 9 : MK8**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	16,421	0.00	0.0		118.58
27	18,280	0.00	0.0		120.30

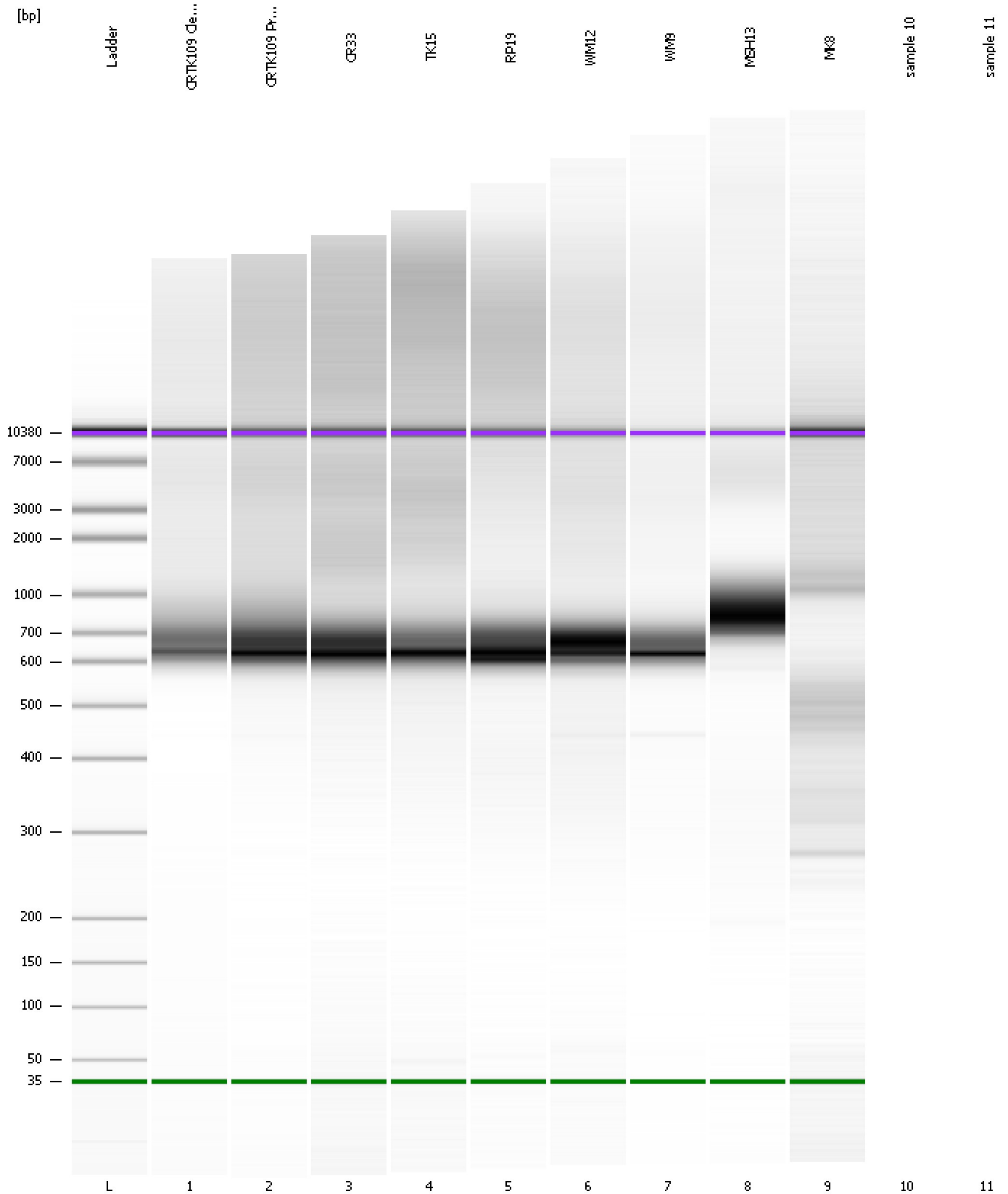
Region table for sample 9 : MK8

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	479	248.87	268.9	888.9	 50	32.0

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 AM

Gel Image

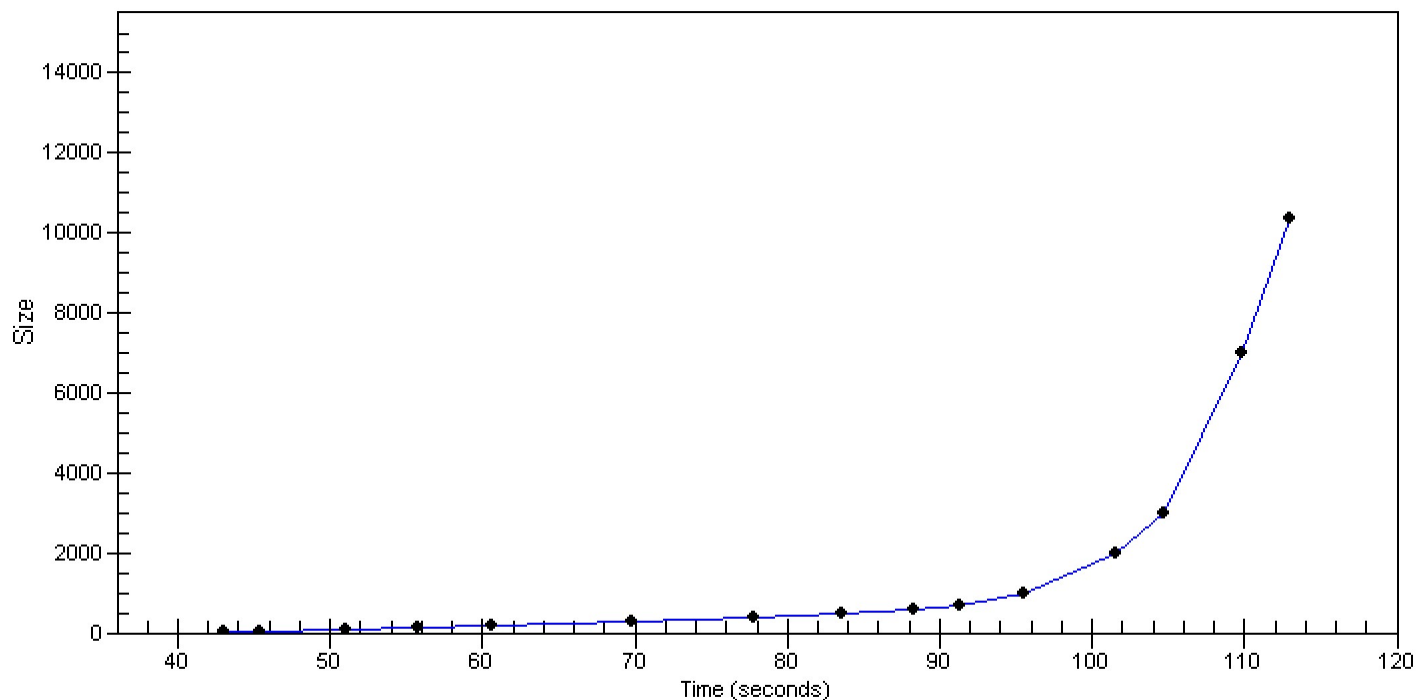


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 AM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-04-26\2018-04-26_001.xad

Created: 4/26/2018 9:03:13 AM
Modified: 4/26/2018 9:44:40 AM

Invalid Samples

Sample 10 has not been run, no results available.

Sample 11 has not been run, no results available.