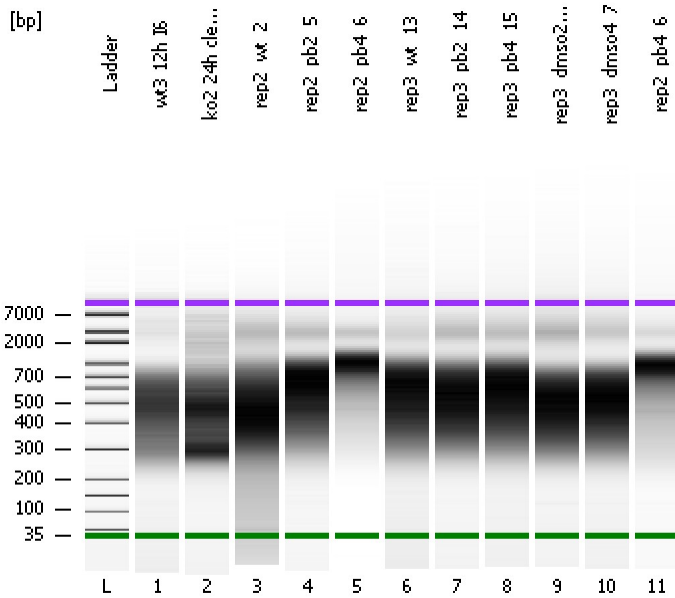


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
Modified: 3/20/2018 11:47:53 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

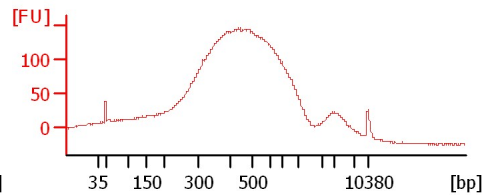
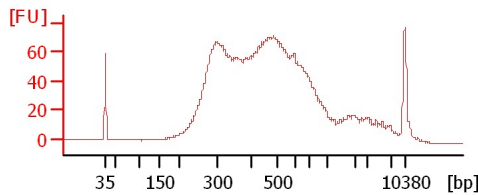
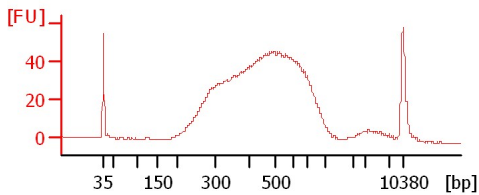
Chip Information:

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

wt3 12h I6

ko2 24h clean I19

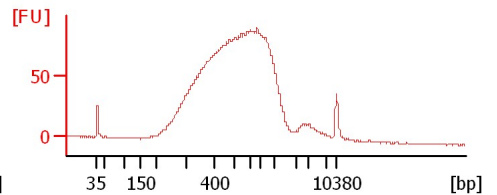
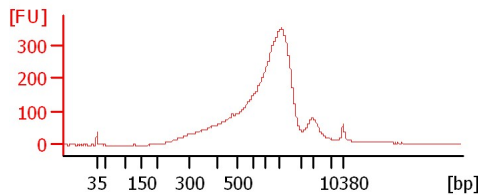
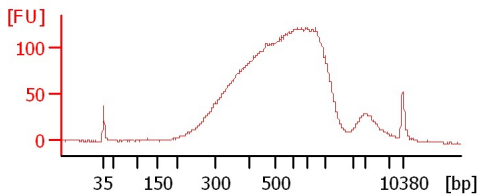
rep2_wt_2



rep2_pb2_5

rep2_pb4_6

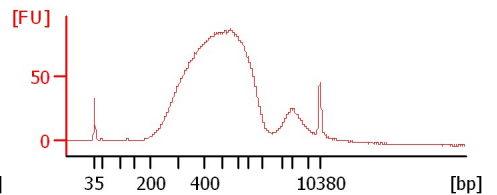
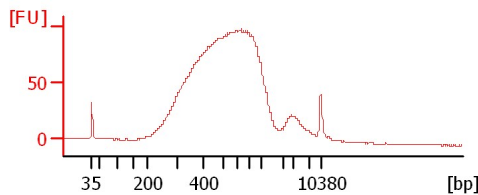
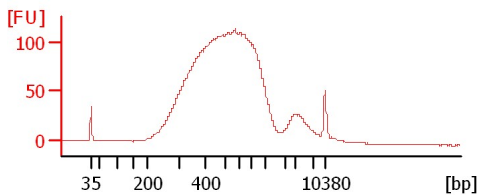
rep3_wt_13



rep3_pb2_14

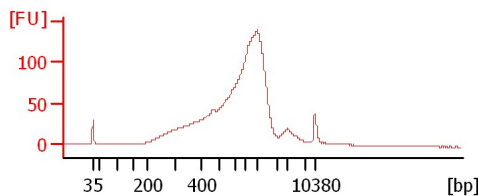
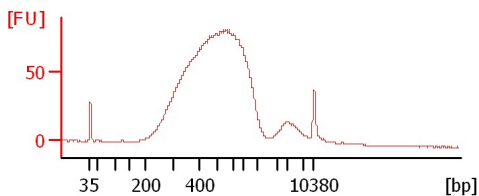
rep3_pb4_15

rep3_dms02_12



rep3_dms04_7

rep2_pb4_6



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
 Modified: 3/20/2018 11:47:53 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
wt3 12h I6		<input type="checkbox"/>	✓			
ko2 24h clean I19		<input type="checkbox"/>	✓			
rep2_wt_2		<input type="checkbox"/>	✓			
rep2_pb2_5		<input type="checkbox"/>	✓			
rep2_pb4_6		<input type="checkbox"/>	✓			
rep3_wt_13		<input type="checkbox"/>	✓			
rep3_pb2_14		<input type="checkbox"/>	✓			
rep3_pb4_15		<input type="checkbox"/>	✓			
rep3_dmso2_12		<input type="checkbox"/>	✓			
rep3_dmso4_7		<input type="checkbox"/>	✓			
rep2_pb4_6		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

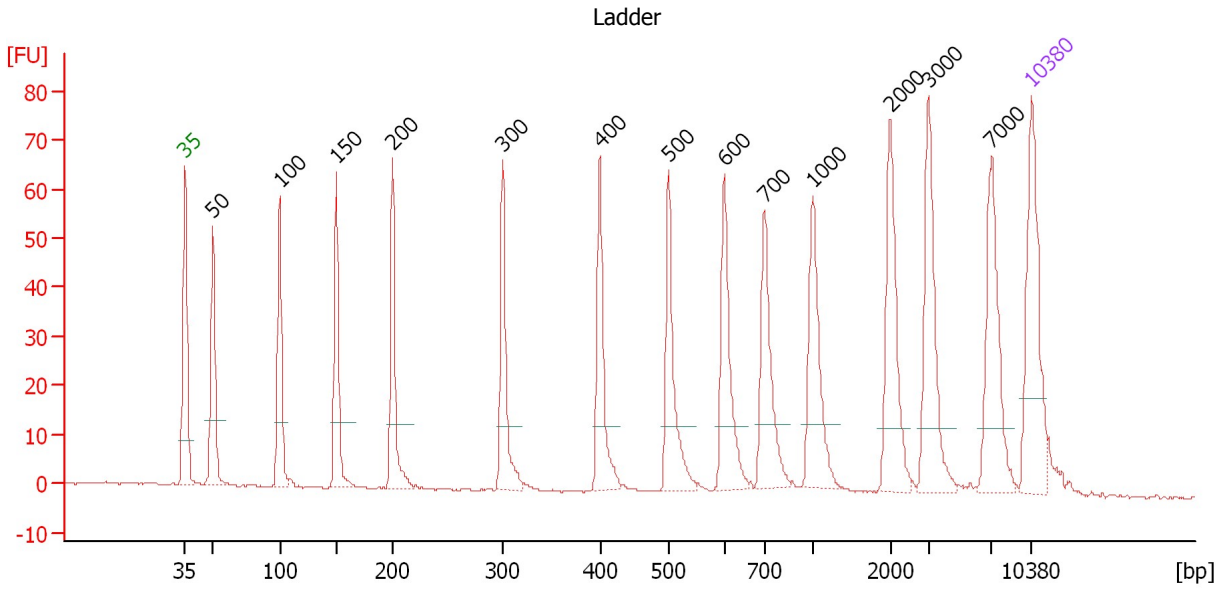
Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

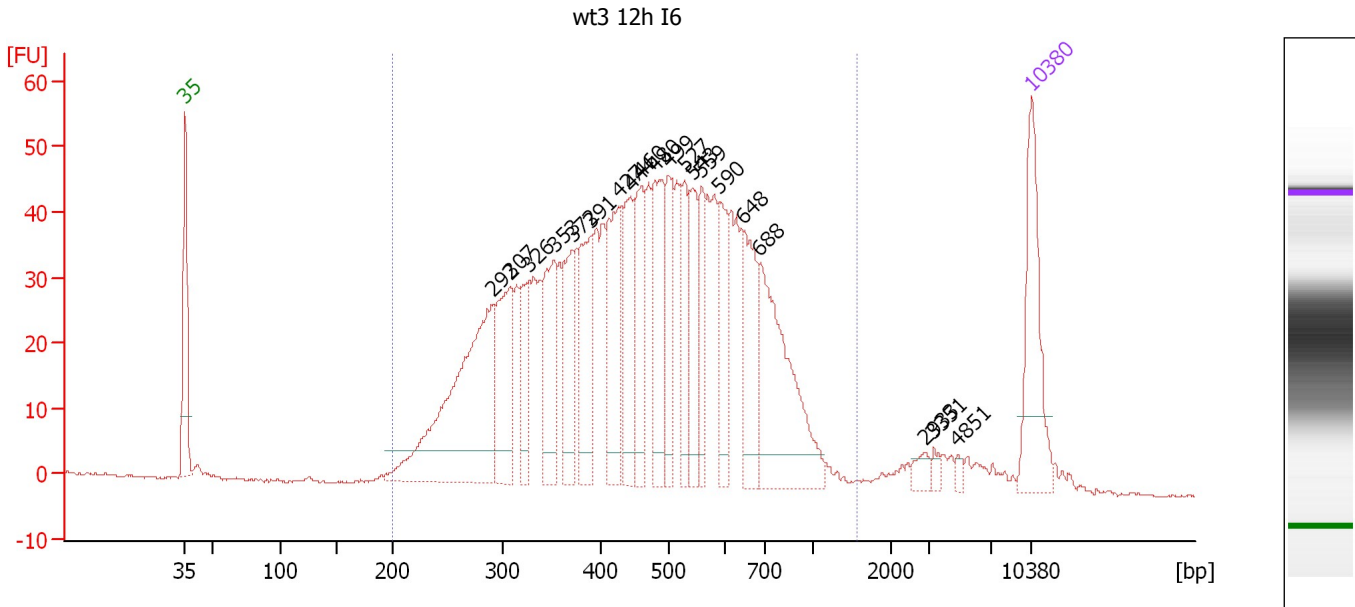
Peak table for Ladder

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	50	150.00	4,545.5	Ladder Peak	45.28
3	100	150.00	2,272.7	Ladder Peak	50.80
4	150	150.00	1,515.2	Ladder Peak	55.50
5	200	150.00	1,136.4	Ladder Peak	60.15
6	300	150.00	757.6	Ladder Peak	69.26
7	400	150.00	568.2	Ladder Peak	77.30
8	500	150.00	454.5	Ladder Peak	83.01
9	600	150.00	378.8	Ladder Peak	87.62
10	700	150.00	324.7	Ladder Peak	90.91
11	1,000	150.00	227.3	Ladder Peak	94.93
12	2,000	150.00	113.6	Ladder Peak	101.33
13	3,000	150.00	75.8	Ladder Peak	104.47
14	7,000	150.00	32.5	Ladder Peak	109.75
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
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Electropherogram Summary Continued ...



Overall Results for sample 1 : wt3 12h I6

Number of peaks found: 20 Corr. Area 1: 1,380.8
 Noise: 0.2

Peak table for sample 1 : wt3 12h I6


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	292	284.43	1,478.0		68.49
3	307	107.24	528.8		69.84
4	326	56.00	259.9		71.39
5	353	88.90	381.7		73.51
6	372	68.36	278.3		75.06
7	391	91.38	353.8		76.60
8	427	105.93	375.5		78.87
9	444	73.40	250.2		79.83
10	460	79.03	260.5		80.70
11	480	93.04	293.8		81.86
12	499	58.87	178.6		82.97
13	527	51.15	146.9		84.28
14	543	60.62	169.1		85.00
15	559	47.97	130.1		85.72
16	590	53.80	138.1		87.17
17	648	80.79	188.9		89.20
18	688	163.15	359.5		90.50
19	2,933	8.94	4.6		104.26
20	3,351	5.58	2.5		104.94
21	4,851	3.98	1.2		106.92
22	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
Modified: 3/20/2018 11:47:53 AM

Electropherogram Summary Continued ...

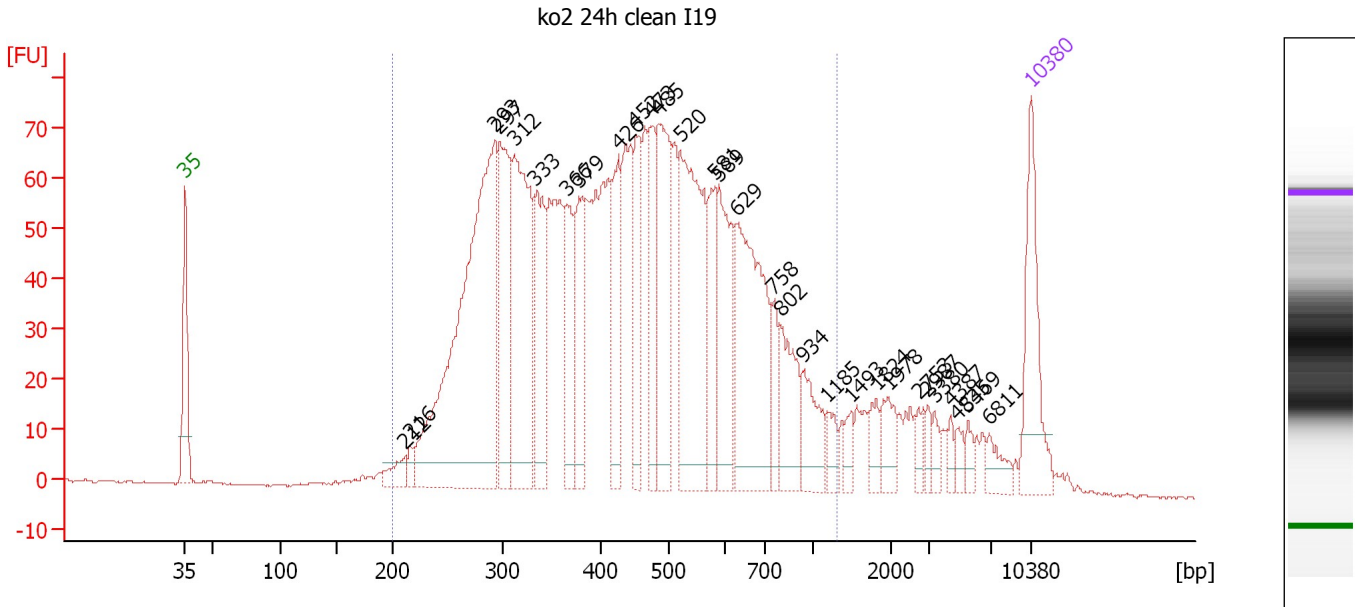
... Region table for sample 1 : wt3 12h I6

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Conc. [pg/ μ l]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,570	483	1,380.8	2,122.07	7,842.1	 96	35.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
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Electropherogram Summary Continued ...



Overall Results for sample 2 : ko2 24h clean I19

Number of peaks found: 30 Corr. Area 1: 2,238.3
 Noise: 0.2

Peak table for sample 2 : ko2 24h clean I19

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	212	24.12	172.6		61.21
3	216	12.57	88.1		61.61
4	293	536.13	2,774.6		68.60
5	297	151.62	773.3		68.99
6	312	239.52	1,163.1		70.22
7	333	118.81	540.9		71.90
8	366	81.32	336.2		74.60
9	379	85.99	344.0		75.59
10	426	89.45	318.1		78.79
11	452	84.74	284.1		80.26
12	473	74.98	240.0		81.50
13	485	147.06	459.0		82.18
14	520	227.87	663.3		83.96
15	581	57.88	150.8		86.76
16	589	115.04	296.0		87.11
17	629	192.30	462.9		88.58
18	758	30.81	61.6		91.68
19	802	65.27	123.3		92.28
20	934	45.83	74.3		94.05
21	1,185	17.29	22.1		96.12
22	1,493	13.25	13.4		98.08
23	1,824	16.90	14.0		100.20
24	1,978	24.31	18.6		101.19
25	2,753	11.25	6.2		103.70

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
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Electropherogram Summary Continued ...

... Peak table for sample 2 : ko2 24h clean I19

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,987	9.70	4.9		104.43
27	3,380	9.75	4.4		104.98
28	4,387	7.85	2.7		106.31
29	4,835	9.58	3.0		106.90
30	5,469	8.90	2.5		107.73
31	6,811	16.43	3.7		109.50
32	10,380	75.00	10.9	Upper Marker	113.00

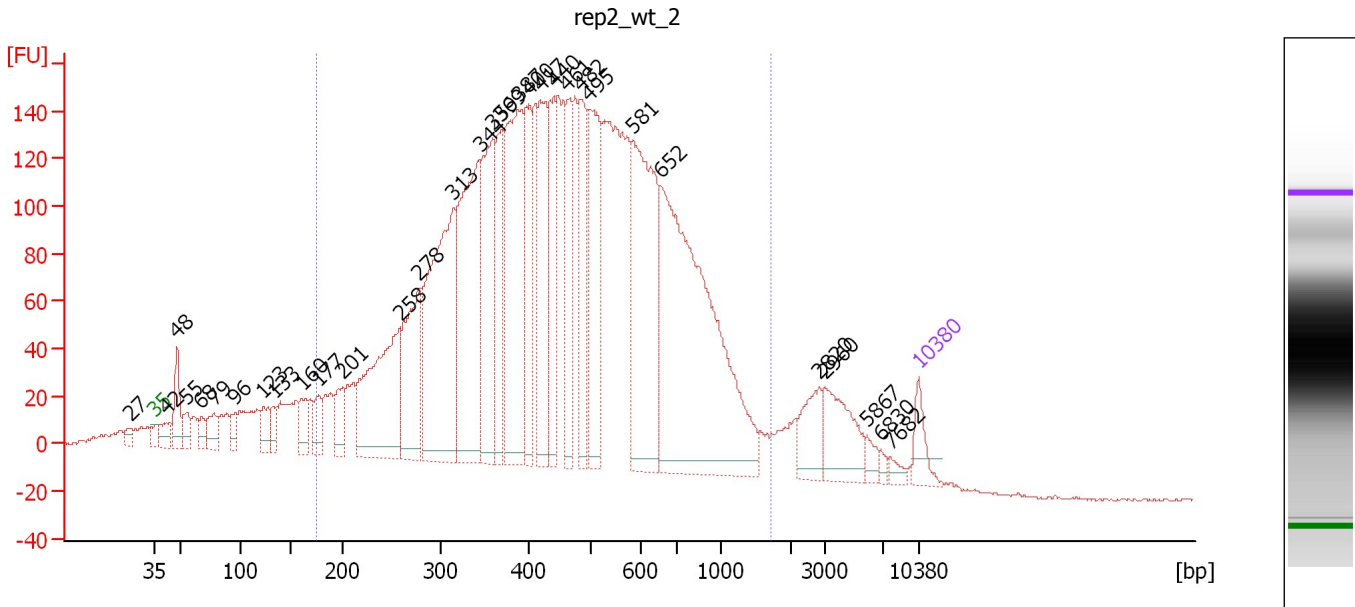
Region table for sample 2 : ko2 24h clean I19

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,320	468	2,238.3	3,031.65	11,730.6	91	39.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
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Electropherogram Summary Continued ...



Overall Results for sample 3 : rep2_wt_2

Number of peaks found: 32 Corr. Area 1: 5,299.9
 Noise: 0.7

Peak table for sample 3 : rep2_wt_2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	27	0.00	0.0		40.70
2	35	125.00	5,411.3	Lower Marker	43.00
3	42	92.61	3,334.0		44.08
4	48	189.80	5,965.9		45.00
5	55	93.36	2,548.9		45.88
6	68	62.65	1,398.3		47.25
7	79	113.56	2,191.6		48.43
8	96	71.89	1,137.3		50.33
9	123	107.73	1,331.4		52.92
10	133	79.72	908.2		53.90
11	160	119.39	1,129.5		56.44
12	177	111.36	953.4		58.01
13	201	140.55	1,061.4		60.21
14	258	776.25	4,557.5		65.44
15	278	483.99	2,638.5		67.25
16	313	1,084.22	5,243.4		70.33
17	344	945.72	4,169.2		72.77
18	359	603.93	2,549.6		73.99
19	369	344.41	1,415.7		74.77
20	387	877.82	3,432.7		76.29
21	400	370.85	1,406.1		77.27
22	417	463.68	1,683.1		78.29
23	440	344.56	1,187.4		79.56
24	461	294.25	967.0		80.79
25	482	332.86	1,047.3		81.96

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
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Electropherogram Summary Continued ...

... Peak table for sample 3 : rep2_wt_2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	495	533.15	1,631.1		82.74
27	581	871.42	2,271.8		86.75
28	652	1,636.03	3,799.8		89.34
29	2,820	134.49	72.3		103.91
30	2,960	207.18	106.1		104.35
31	5,867	34.32	8.9		108.26
32	6,830	15.53	3.4		109.53
33	7,682	24.17	4.8		110.41
34	10,380	75.00	10.9	Upper Marker	113.00

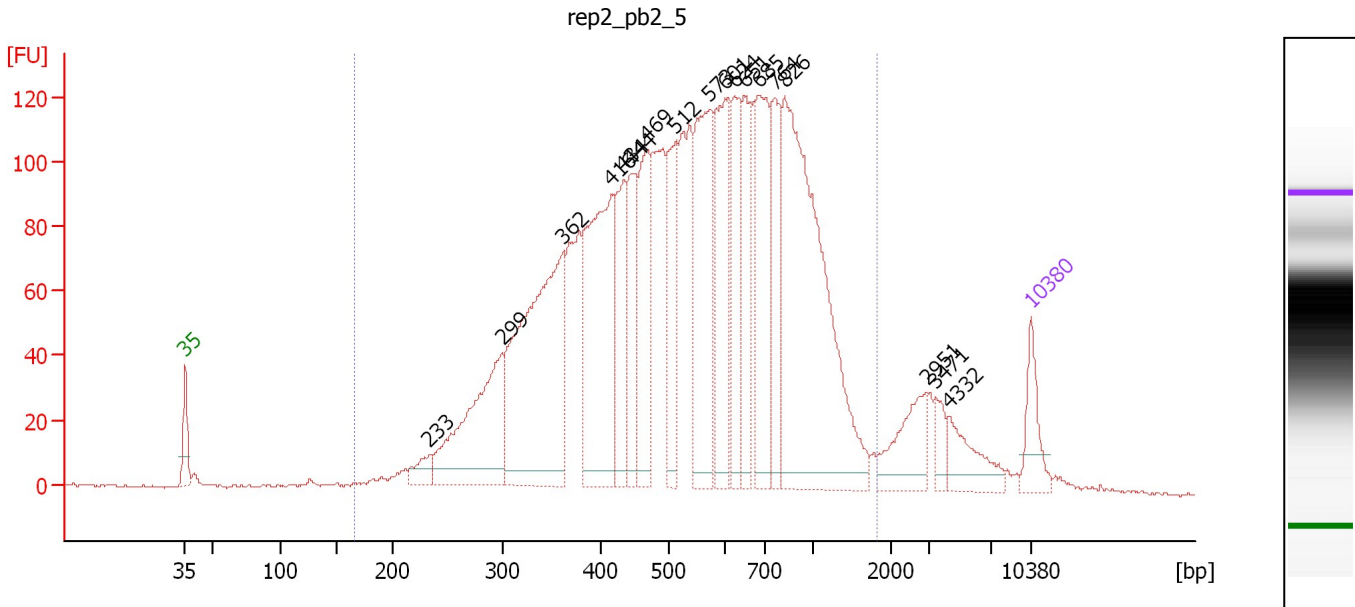
Region table for sample 3 : rep2_wt_2

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
174	1,719	497	5,299.9	12,569.50	48,483.8	85	45.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
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Electropherogram Summary Continued ...



Overall Results for sample 4 : rep2_pb2_5

Number of peaks found: 18 Corr. Area 1: 3,437.4
 Noise: 0.4

Peak table for sample 4 : rep2_pb2_5

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	233	54.46	353.4		63.20
3	299	453.92	2,297.9		69.19
4	362	819.62	3,429.0		74.25
5	416	586.74	2,135.2		78.23
6	434	219.93	767.0		79.27
7	444	214.70	732.1		79.83
8	469	287.18	928.2		81.23
9	512	207.80	615.3		83.55
10	573	414.81	1,096.0		86.39
11	601	321.52	811.1		87.63
12	624	196.64	477.4		88.41
13	651	209.65	488.1		89.29
14	685	342.11	756.4		90.42
15	764	214.28	425.0		91.77
16	826	949.01	1,741.5		92.59
17	2,951	120.45	61.8		104.32
18	3,471	39.63	17.3		105.10
19	4,332	91.94	32.2		106.23
20	10,380	75.00	10.9	Upper Marker	113.00

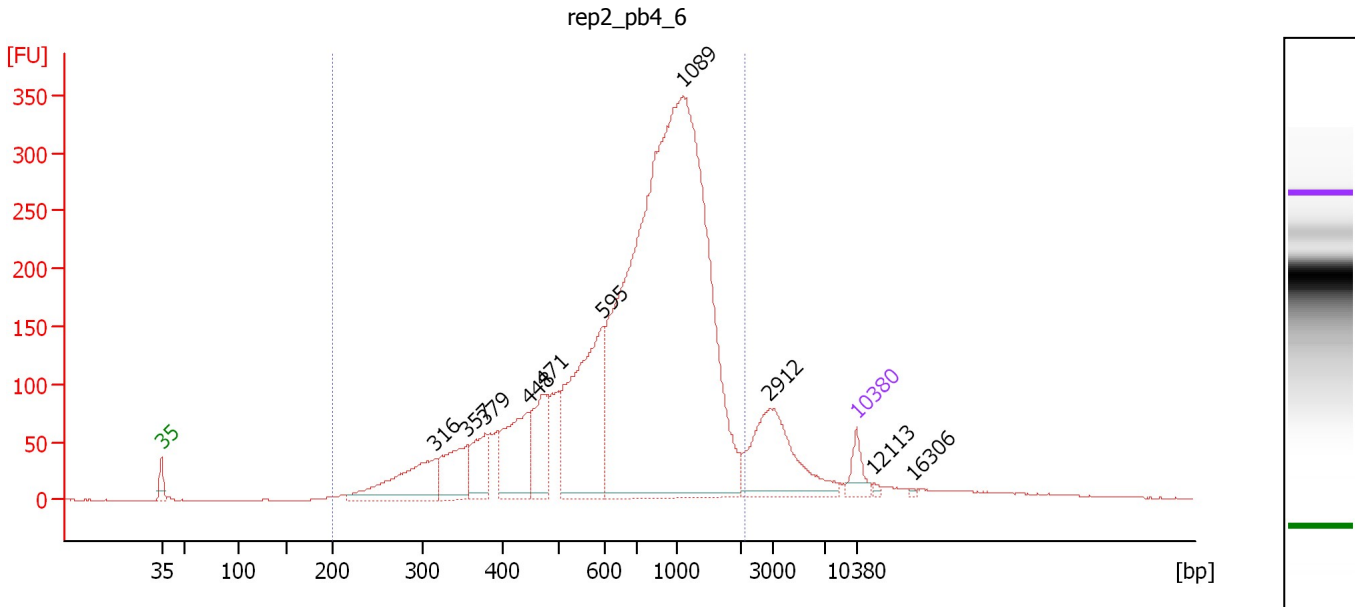
Region table for sample 4 : rep2_pb2_5

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
166	1,824	591	3,437.4	6,824.90	22,048.1	93	44.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
 Modified: 3/20/2018 11:47:53 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : rep2_pb4_6

Number of peaks found: 10 Corr. Area 1: 5,092.8
 Noise: 0.4

Peak table for sample 5 : rep2_pb4_6

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	316	474.50	2,275.5		70.54
3	357	305.31	1,297.0		73.81
4	379	236.42	944.3		75.64
5	448	429.95	1,454.2		80.04
6	471	326.80	1,052.4		81.33
7	595	954.50	2,429.9		87.39
8	1,089	4,447.39	6,187.7		95.50
9	2,912	478.35	248.9		104.20
10	10,380	75.00	10.9	Upper Marker	113.00
11	12,113	0.00	0.0		114.66
12	16,306	0.00	0.0		118.69

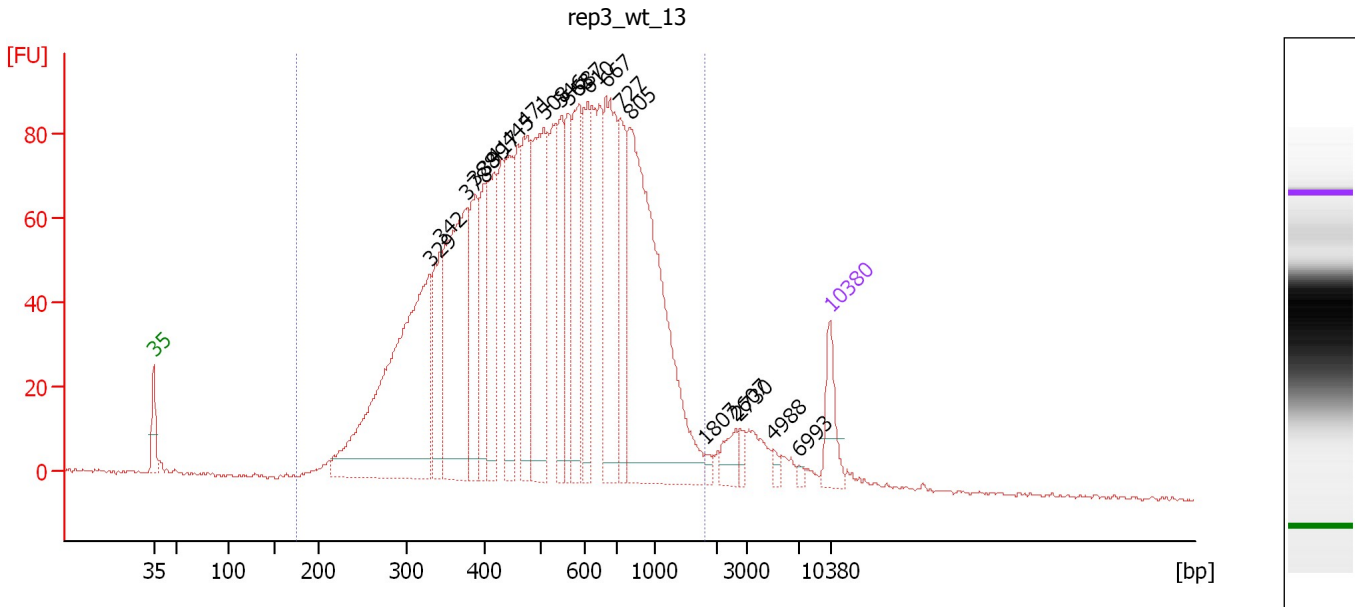
Region table for sample 5 : rep2_pb4_6

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	2,156	842	5,092.8	7,959.18	19,540.4	91	46.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
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Electropherogram Summary Continued ...



Overall Results for sample 6 : rep3 wt 13

Number of peaks found: 21 Corr. Area 1: 2,601.6
 Noise: 0.5

Peak table for sample 6 : rep3 wt 13


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	329	1,014.96	4,676.8		71.57
3	342	211.82	938.8		72.62
4	378	594.30	2,385.1		75.49
5	388	232.33	907.7		76.32
6	399	210.19	798.6		77.20
7	417	255.30	928.4		78.25
8	445	239.16	815.0		79.85
9	471	291.84	939.5		81.34
10	508	414.33	1,236.0		83.38
11	546	211.22	585.8		85.14
12	563	180.66	486.2		85.92
13	587	259.92	670.9		87.02
14	610	230.60	572.4		87.96
15	667	393.52	893.6		89.83
16	727	206.04	429.6		91.27
17	805	906.18	1,706.0		92.31
18	1,807	12.78	10.7		100.09
19	2,607	41.81	24.3		103.24
20	2,730	18.14	10.1		103.62
21	4,988	13.50	4.1		107.10
22	6,993	8.92	1.9		109.75
23	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
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Electropherogram Summary Continued ...

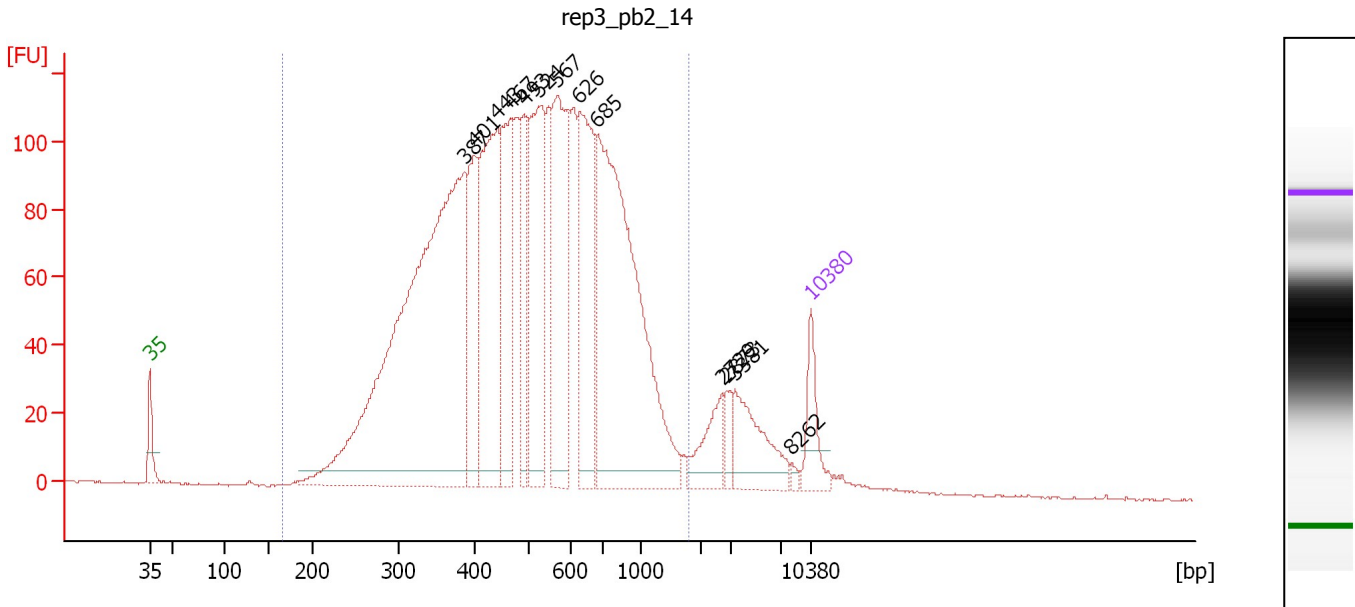
... Region table for sample 6 : rep3 wt 13

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Conc. [pg/ μ l]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
175	1,811	569	2,601.6	6,931.90	23,015.2	 95	44.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

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Electropherogram Summary Continued ...



Overall Results for sample 7 : rep3_pb2_14

Number of peaks found: 13 Corr. Area 1: 3,257.7
 Noise: 0.4

Peak table for sample 7 : rep3_pb2_14

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	387	1,793.62	7,029.4		76.22
3	401	287.33	1,085.9		77.35
4	443	538.68	1,840.6		79.78
5	467	304.96	989.1		81.14
6	493	194.71	598.6		82.60
7	524	406.29	1,174.2		84.13
8	567	440.08	1,175.5		86.11
9	626	391.12	946.3		88.48
10	685	1,021.86	2,261.7		90.40
11	2,729	87.95	48.8		103.62
12	2,873	32.64	17.2		104.07
13	3,381	142.95	64.1		104.98
14	8,262	7.90	1.4		110.97
15	10,380	75.00	10.9	Upper Marker	113.00

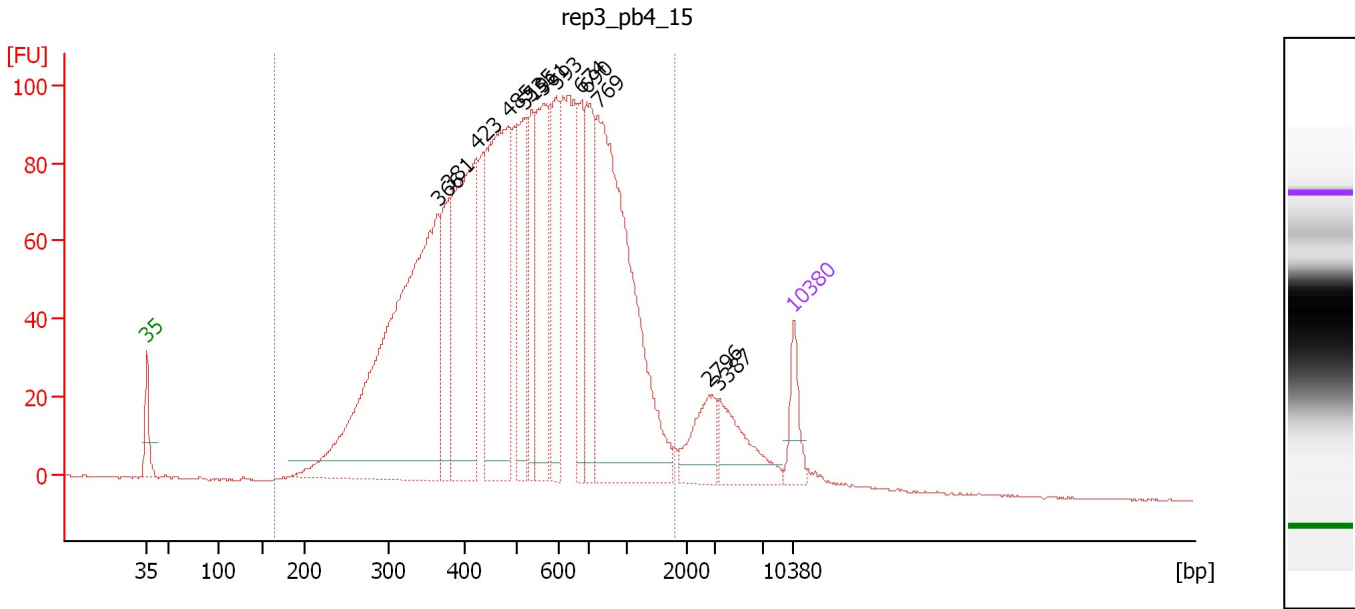
Region table for sample 7 : rep3_pb2_14

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
167	1,806	549	3,257.7	6,194.85	20,912.9	93	43.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

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Electropherogram Summary Continued ...



Overall Results for sample 8 : rep3_pb4_15

Number of peaks found: 13 Corr. Area 1: 2,847.1
 Noise: 0.2

Peak table for sample 8 : rep3_pb4_15

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	366	1,552.93	6,433.8		74.54
3	381	243.99	970.9		75.75
4	423	731.03	2,616.1		78.63
5	485	766.32	2,394.4		82.15
6	519	293.00	855.6		83.88
7	535	199.95	566.1		84.63
8	561	421.76	1,138.2		85.84
9	593	278.26	711.2		87.28
10	674	202.92	456.2		90.05
11	690	272.96	599.7		90.57
12	769	1,066.99	2,101.5		91.84
13	2,796	119.82	64.9		103.83
14	3,387	143.78	64.3		104.99
15	10,380	75.00	10.9	Upper Marker	113.00

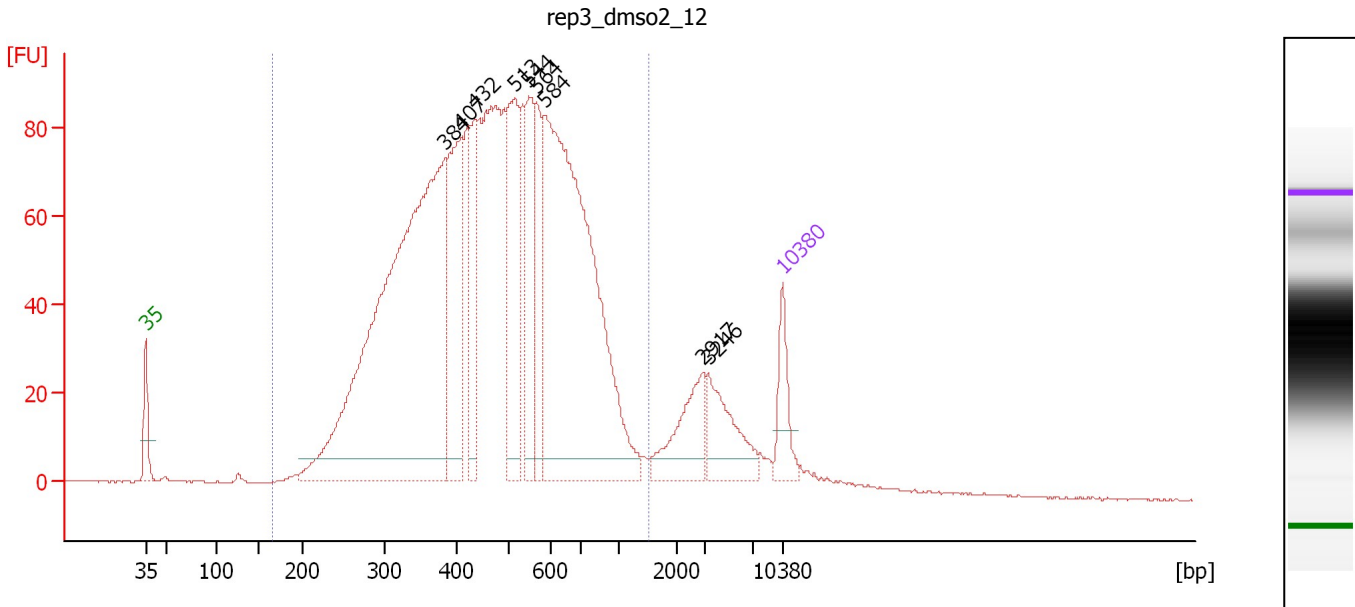
Region table for sample 8 : rep3_pb4_15

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
165	1,820	574	2,847.1	7,213.86	23,872.3	94	45.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

Created: 3/20/2018 11:05:32 AM
 Modified: 3/20/2018 11:47:53 AM

Electropherogram Summary Continued ...



Overall Results for sample 9 : rep3_dms02_12

Number of peaks found: 9 Corr. Area 1: 2,439.1
 Noise: 0.2

Peak table for sample 9 : rep3_dms02_12

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	384	1,708.44	6,748.0		75.98
3	407	374.54	1,395.5		77.68
4	432	197.12	690.9		79.14
5	513	310.60	918.1		83.59
6	544	236.52	658.2		85.06
7	564	153.87	413.7		85.94
8	584	1,208.20	3,135.2		86.87
9	2,917	130.65	67.9		104.21
10	3,246	128.12	59.8		104.80
11	10,380	75.00	10.9	Upper Marker	113.00

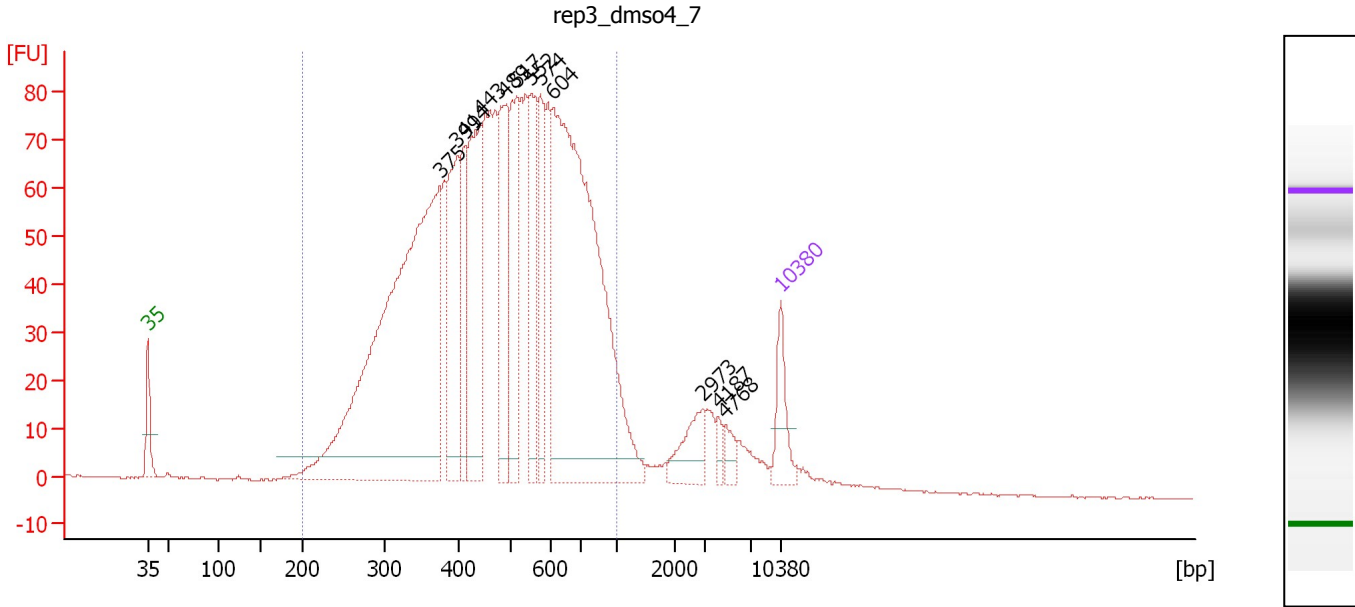
Region table for sample 9 : rep3_dms02_12

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
167	1,531	499	2,439.1	5,584.51	20,002.9	91	36.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

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Electropherogram Summary Continued ...



Overall Results for sample 10 : rep3_dms04_7

Number of peaks found: 12 Corr. Area 1: 2,126.3
 Noise: 0.2

Peak table for sample 10 : rep3_dms04_7

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	375	1,596.16	6,455.4		75.26
3	399	315.37	1,198.1		77.20
4	414	177.21	648.8		78.09
5	443	391.86	1,341.1		79.74
6	489	320.33	992.2		82.39
7	517	258.36	756.8		83.81
8	552	221.23	607.4		85.40
9	574	172.42	455.4		86.40
10	604	1,257.98	3,153.7		87.76
11	2,973	80.65	41.1		104.39
12	4,187	20.29	7.3		106.04
13	4,768	28.43	9.0		106.81
14	10,380	75.00	10.9	Upper Marker	113.00

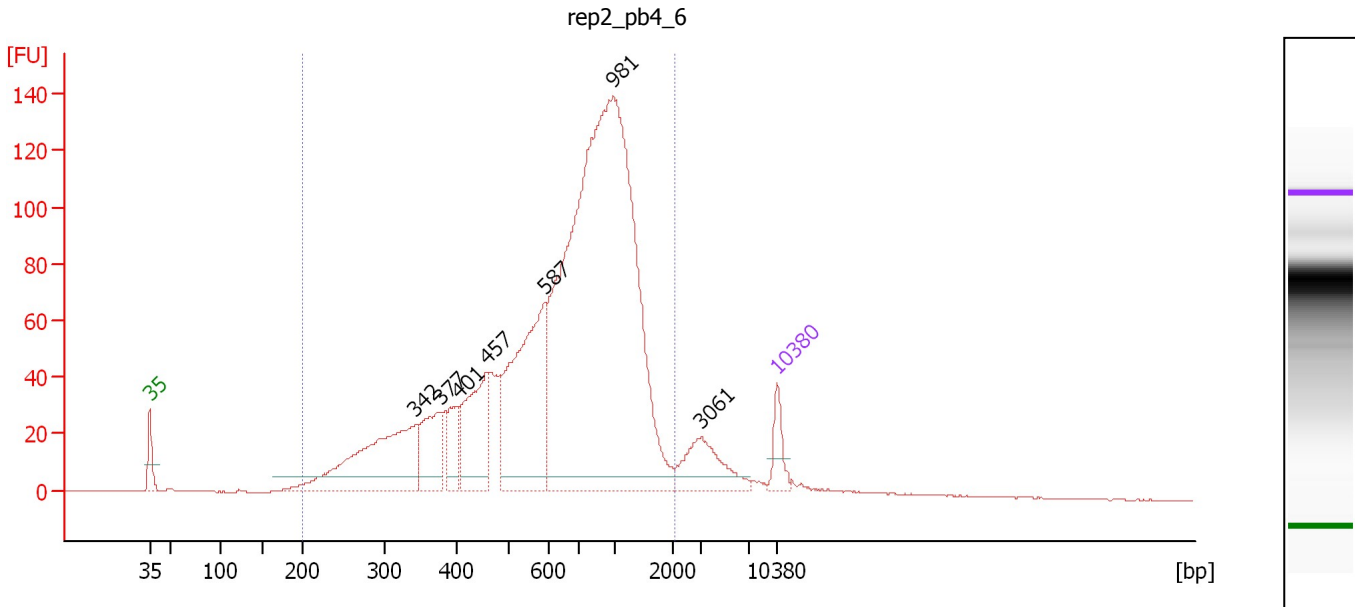
Region table for sample 10 : rep3_dms04_7

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	501	2,126.3	5,871.71	20,561.8	92	32.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

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Electropherogram Summary Continued ...



Overall Results for sample 11 : rep2_pb4_6

Number of peaks found: 7 Corr. Area 1: 2,196.7
 Noise: 0.2

Peak table for sample 11 : rep2_pb4_6

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	342	708.29	3,139.7		72.62
3	377	250.12	1,006.4		75.41
4	401	143.11	540.2		77.38
5	457	388.82	1,290.5		80.53
6	587	796.89	2,057.5		87.01
7	981	2,992.29	4,619.4		94.68
8	3,061	188.47	93.3		104.55
9	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 11 : rep2_pb4_6

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Conc. [pg/μl]	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	2,092	758	2,196.7	5,995.19	16,417.5	93	46.8

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-03-20\2018-03-20_002.xad

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Gel Image

