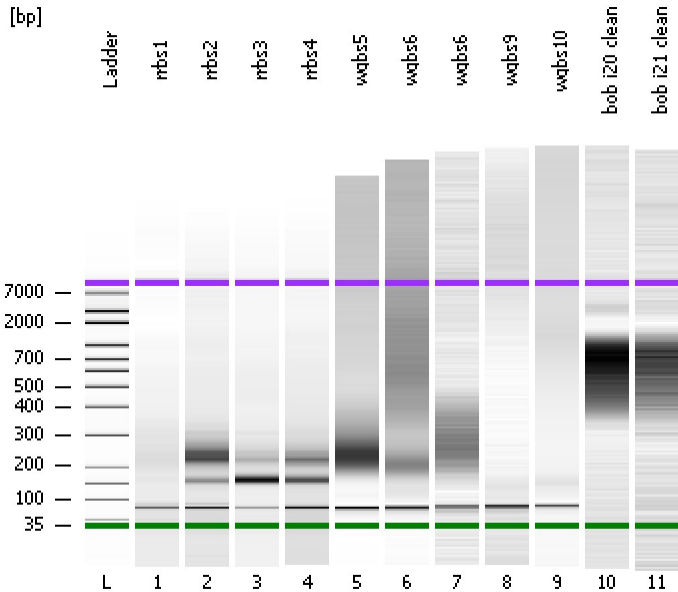


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

Created: 2/19/2016 3:56:58 PM
Modified: 2/19/2016 4:39:07 PM

Electrophoresis File Run Summary



Instrument Information:

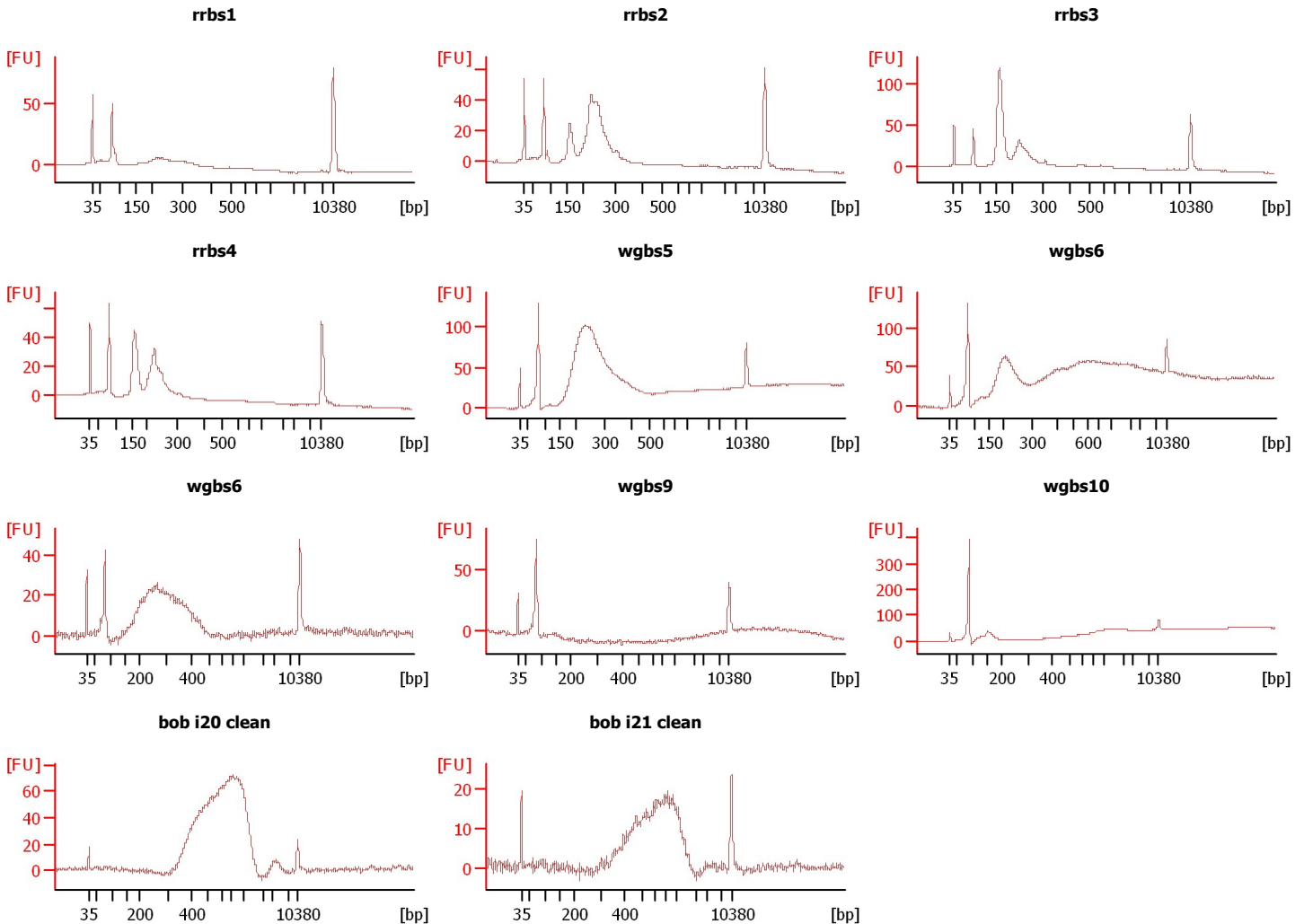
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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\2016-02-19\2016-02-19_001.xad

Created: 2/19/2016 3:56:58 PM
Modified: 2/19/2016 4:39:07 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
rrbs1		<input type="checkbox"/>	✓			
rrbs2		<input type="checkbox"/>	✓			
rrbs3		<input type="checkbox"/>	✓			
rrbs4		<input type="checkbox"/>	✓			
wgbs5		<input type="checkbox"/>	✓			
wgbs6		<input type="checkbox"/>	✓			
wgbs6		<input type="checkbox"/>	✓			
wgbs9		<input type="checkbox"/>	✓			
wgbs10		<input type="checkbox"/>	✓			
bob i20 clean		<input type="checkbox"/>	✓			
bob i21 clean		<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\2016-02-19\2016-02-19_001.xad

Created: 2/19/2016 3:56:58 PM
Modified: 2/19/2016 4:39:07 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

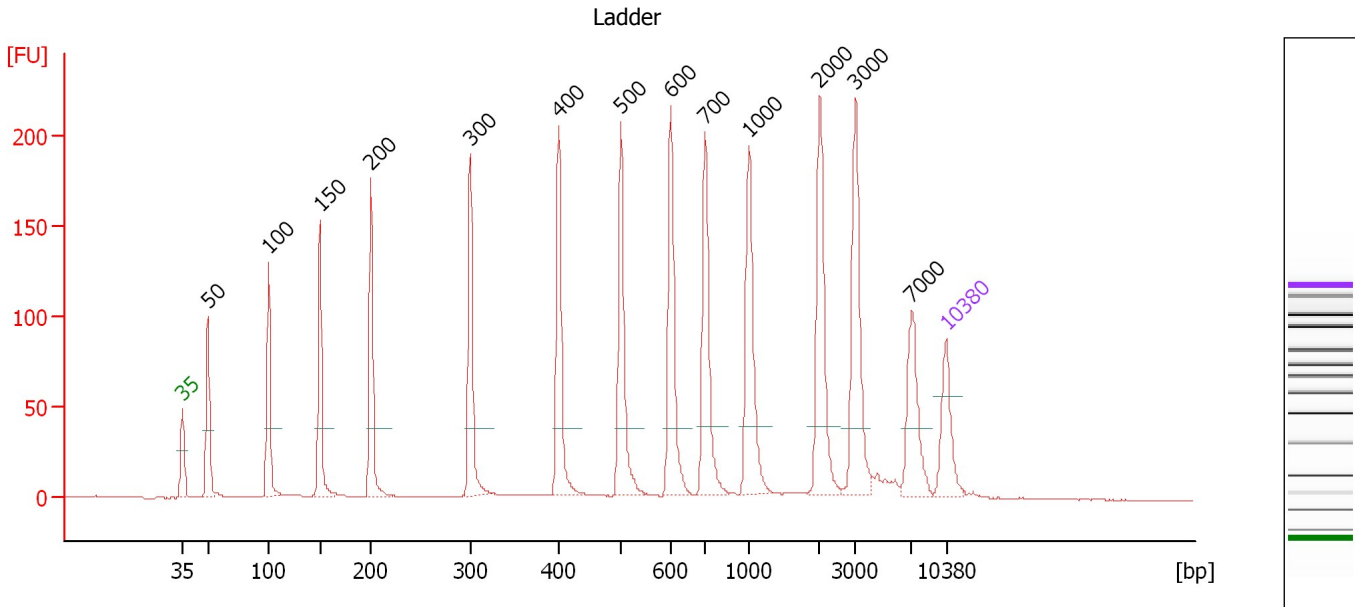
Ladder

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\2016-02-19\2016-02-19_001.xad

Created: 2/19/2016 3:56:58 PM
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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

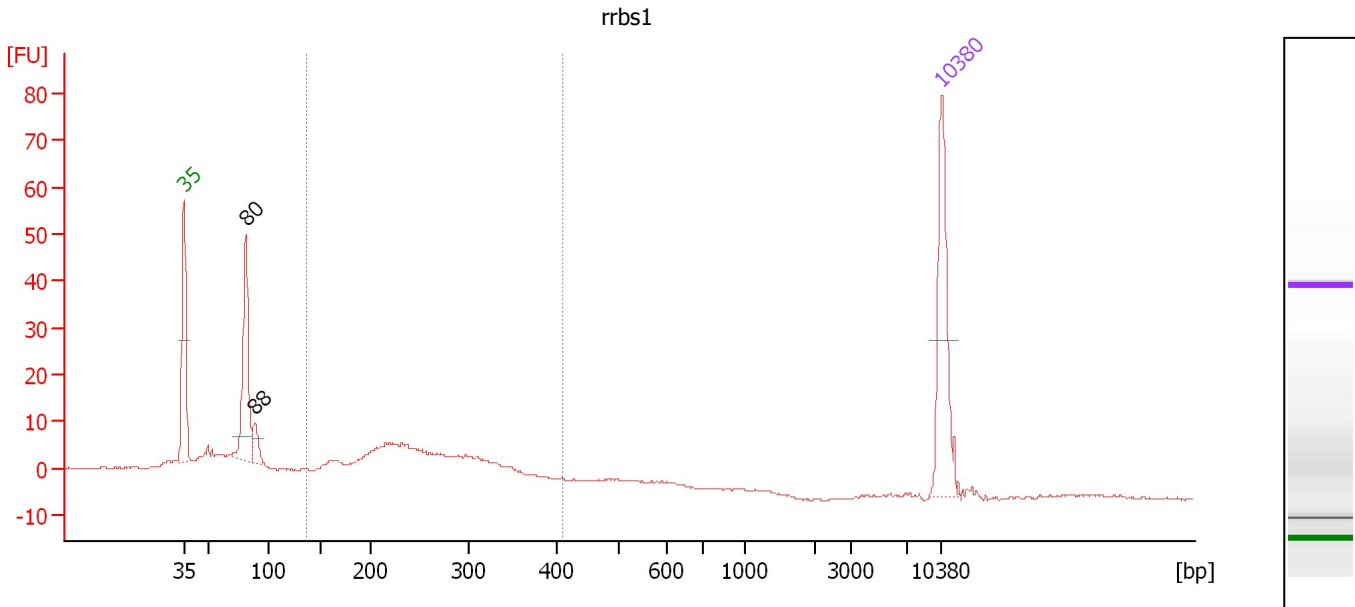
Peak table for Ladder

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	45.34
3	100	150.00	2,272.7	Ladder Peak	50.91
4	150	150.00	1,515.2	Ladder Peak	55.59
5	200	150.00	1,136.4	Ladder Peak	60.28
6	300	150.00	757.6	Ladder Peak	69.36
7	400	150.00	568.2	Ladder Peak	77.46
8	500	150.00	454.5	Ladder Peak	83.17
9	600	150.00	378.8	Ladder Peak	87.71
10	700	150.00	324.7	Ladder Peak	90.89
11	1,000	150.00	227.3	Ladder Peak	94.89
12	2,000	150.00	113.6	Ladder Peak	101.38
13	3,000	150.00	75.8	Ladder Peak	104.65
14	7,000	150.00	32.5	Ladder Peak	109.78
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 1 : rrbs1

Number of peaks found: 2 Corr. Area 1: 139.9
 Noise: 0.2

Peak table for sample 1 : rrbs1

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	80	121.30	2,289.0		48.71
3	88	21.77	375.7		49.55
4	10,380	75.00	10.9	Upper Marker	113.00

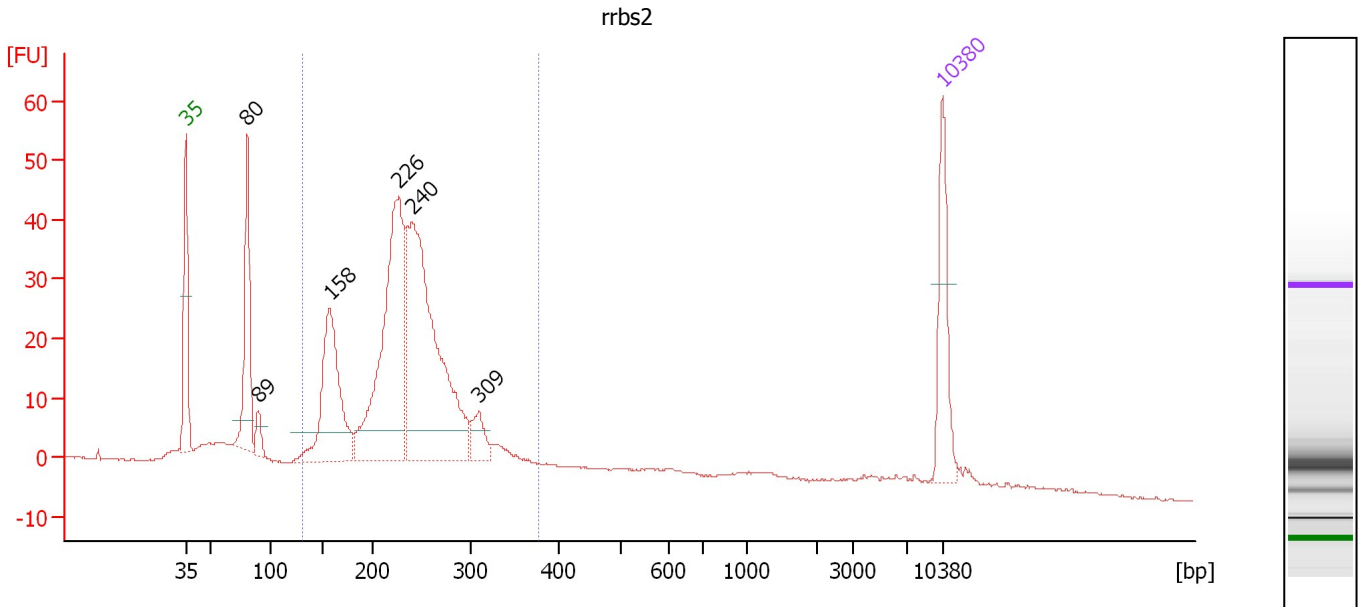
Region table for sample 1 : rrbs1

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
136	411	254	139.9	1,492.6	232.38	48	23.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 2 : rrbs2

Number of peaks found: 6 Corr. Area 1: 537.9
 Noise: 0.2

Peak table for sample 2 : rrbs2

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	80	174.82	3,314.3		48.67
3	89	24.38	413.9		49.71
4	158	233.72	2,245.0		56.32
5	226	389.58	2,606.5		62.68
6	240	459.23	2,898.7		63.92
7	309	32.25	158.2		70.08
8	10,380	75.00	10.9	Upper Marker	113.00

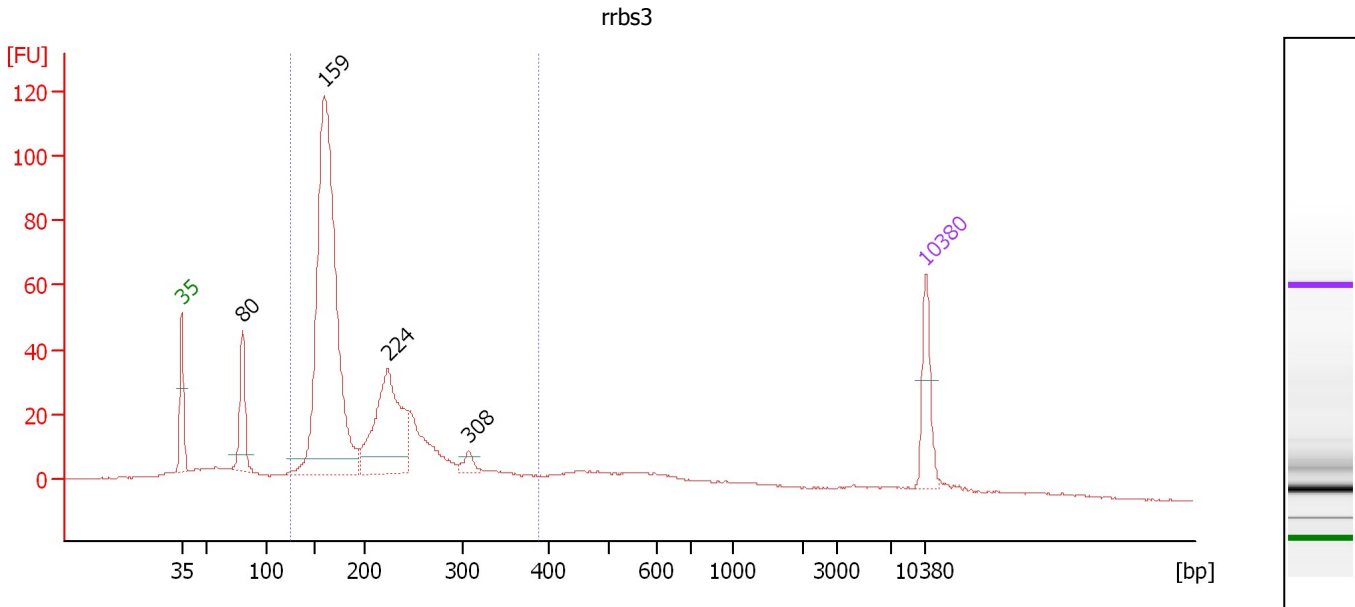
Region table for sample 2 : rrbs2

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
131	378	234	537.9	8,782.2	1,280.75	73	20.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

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 Modified: 2/19/2016 4:39:07 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : rrbs3

Number of peaks found: 4 Corr. Area 1: 786.0
 Noise: 0.2

Peak table for sample 3 : rrbs3

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	80	152.22	2,870.5		48.72
3	159	1,157.03	11,018.0		56.45
4	224	350.97	2,378.7		62.42
5	308	23.94	117.8		70.00
6	10,380	75.00	10.9	Upper Marker	113.00

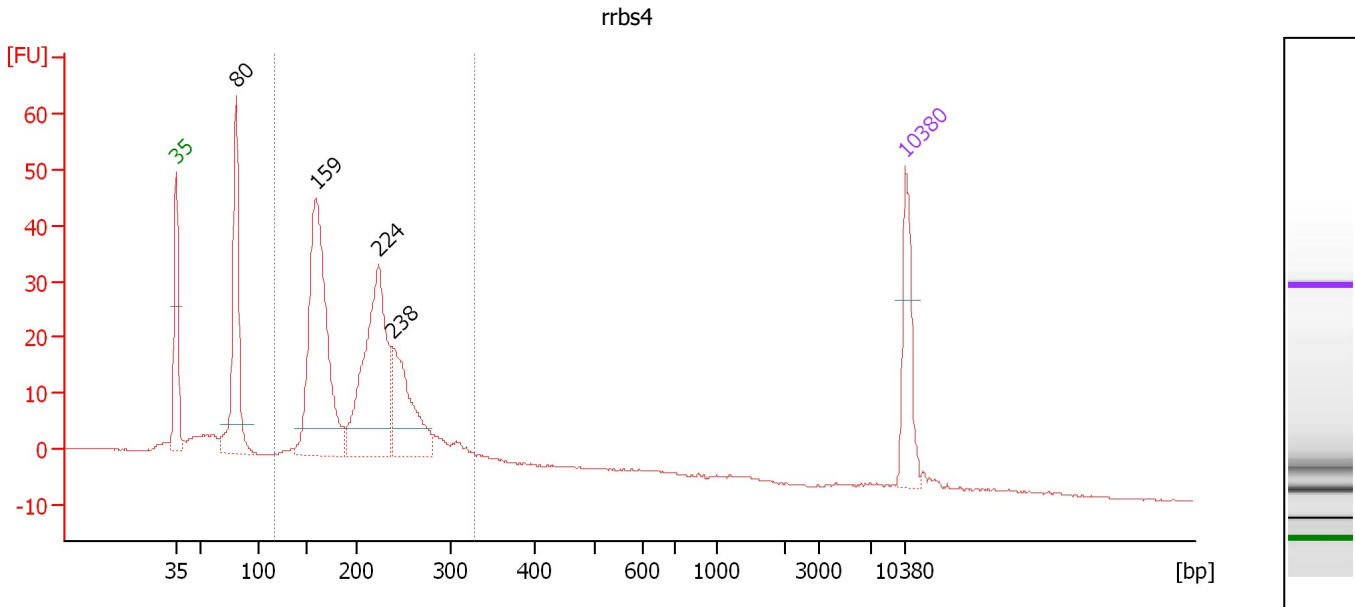
Region table for sample 3 : rrbs3

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
124	389	204	786.0	16,221.1	1,995.32	72	27.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 4 : rrbs4

Number of peaks found: 4 Corr. Area 1: 411.9
 Noise: 0.2

Peak table for sample 4 : rrbs4

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	80	276.05	5,197.2		48.74
3	159	466.94	4,456.0		56.42
4	224	347.28	2,349.1		62.46
5	238	170.37	1,084.2		63.74
6	10,380	75.00	10.9	Upper Marker	113.00

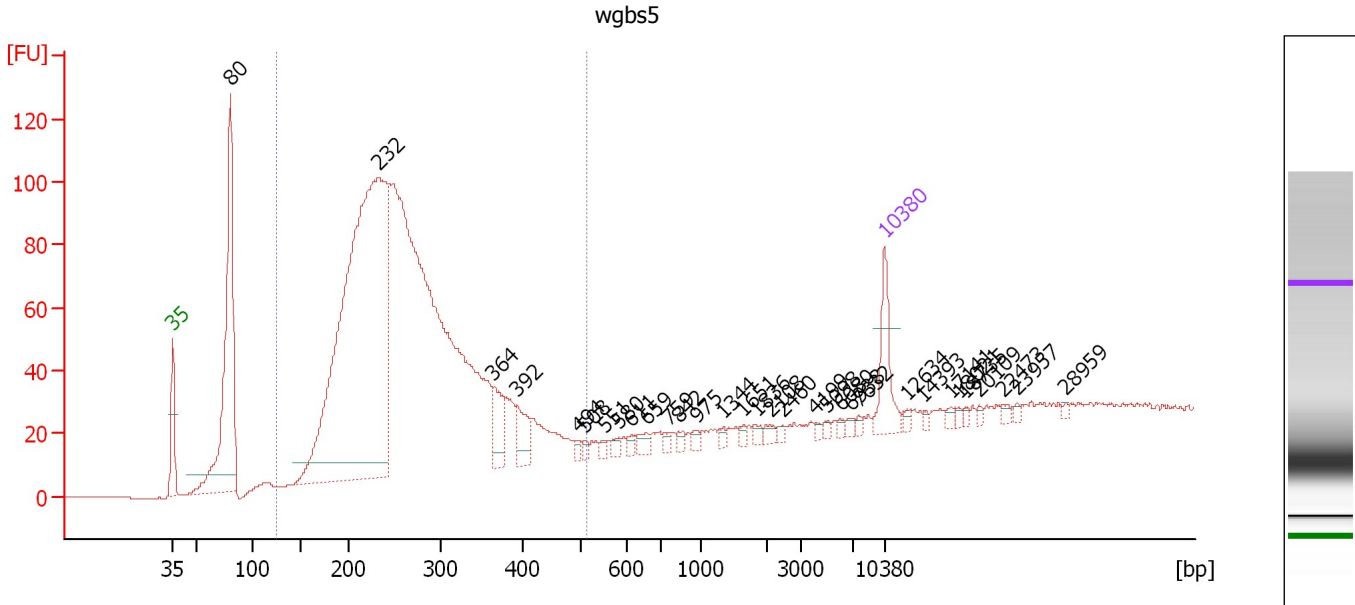
Region table for sample 4 : rrbs4

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
116	327	208	411.9	8,542.9	1,102.46	73	21.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 5 : wgbs5

Number of peaks found: 32 Corr. Area 1: 1,639.8
 Noise: 0.2

Peak table for sample 5 : wgbs5

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	80	662.39	12,578.3		48.66
3	232	1,816.59	11,871.5		63.17
4	364	68.64	285.7		74.54
5	392	55.71	215.1		76.85
6	494	9.81	30.1		82.82
7	508	9.35	27.9		83.55
8	551	10.09	27.7		85.49
9	580	12.51	32.7		86.80
10	611	11.43	28.3		88.06
11	659	17.51	40.3		89.58
12	759	8.49	16.9		91.68
13	842	10.86	19.6		92.78
14	975	9.31	14.5		94.56
15	1,344	7.59	8.6		97.12
16	1,651	7.04	6.5		99.12
17	1,836	9.15	7.6		100.32
18	2,108	10.16	7.3		101.74
19	2,460	5.36	3.3		102.89
20	4,199	5.52	2.0		106.19
21	5,098	5.64	1.7		107.34
22	6,080	6.90	1.7		108.60
23	6,938	7.14	1.6		109.70
24	7,632	6.62	1.3		110.38
25	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad


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Electropherogram Summary Continued ...

... Peak table for sample 5 : wgbs5

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	12,634	0.00	0.0		115.15
27	14,393	0.00	0.0		116.82
28	17,141	0.00	0.0		119.44
29	18,021	0.00	0.0		120.28
30	18,735	0.00	0.0		120.96
31	20,109	0.00	0.0		122.27
32	22,473	0.00	0.0		124.53
33	23,957	0.00	0.0		125.94
34	28,959	0.00	0.0		130.71

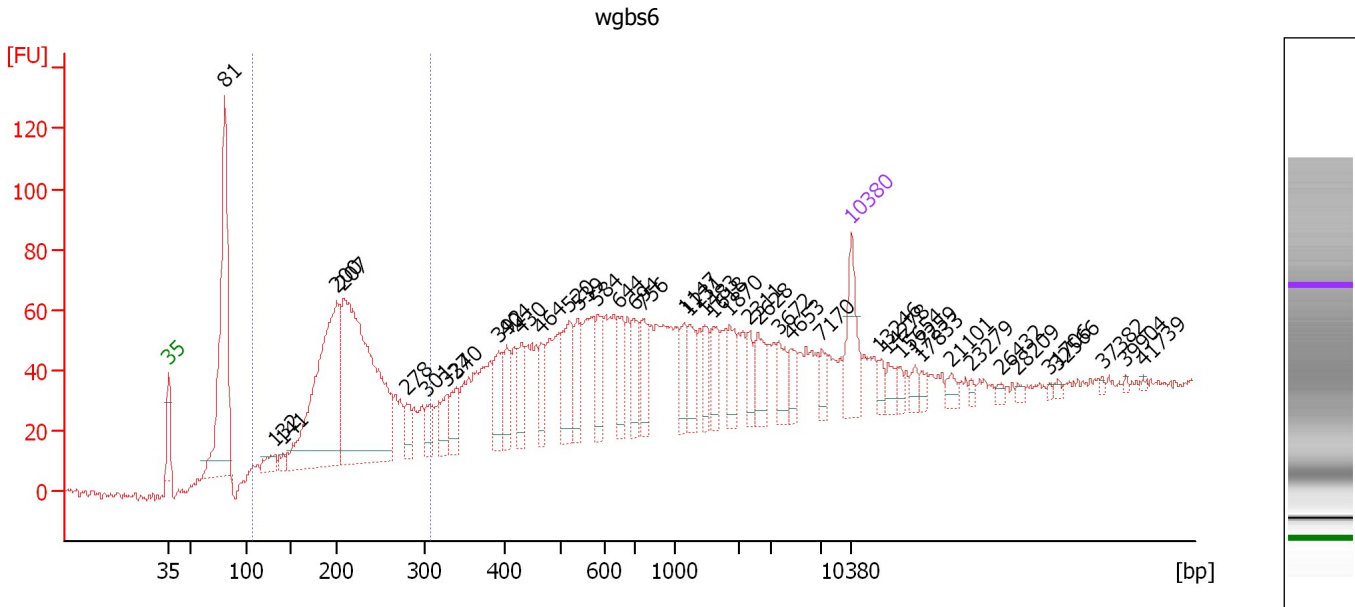
Region table for sample 5 : wgbs5

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
125	510	268	1,639.8	23,159.5	 3,829.38	80	24.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 6 : wgbs6

Number of peaks found: 43 Corr. Area 1: 697.3
 Noise: 1.2

Peak table for sample 6 : wgbs6

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	81	550.61	10,300.5		48.79
3	132	34.92	401.9		53.87
4	141	16.70	179.5		54.75
5	200	545.99	4,139.0		60.27
6	207	722.58	5,286.8		60.92
7	278	42.07	229.2		67.37
8	301	35.57	179.0		69.45
9	327	50.85	235.3		71.58
10	340	55.15	246.1		72.56
11	392	69.58	268.9		76.83
12	404	67.59	253.4		77.70
13	430	55.96	197.2		79.17
14	464	50.31	164.1		81.14
15	520	105.01	305.8		84.09
16	539	63.30	177.8		84.97
17	584	61.32	159.1		86.99
18	644	62.50	147.0		89.12
19	694	57.08	124.6		90.70
20	756	50.75	101.7		91.63
21	1,147	40.01	52.9		95.84
22	1,231	49.38	60.8		96.39
23	1,483	36.10	36.9		98.03
24	1,618	44.51	41.7		98.90
25	1,870	43.71	35.4		100.54

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

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Electropherogram Summary Continued ...

... Peak table for sample 6 : wgbs6

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,311	28.51	18.7		102.40
27	2,628	44.11	25.4		103.44
28	3,672	37.34	15.4		105.51
29	4,653	28.66	9.3		106.77
30	7,170	21.87	4.6		109.94
31	10,380	75.00	10.9	Upper Marker	113.00
32	13,246	0.00	0.0		115.73
33	14,278	0.00	0.0		116.72
34	15,654	0.00	0.0		118.03
35	17,259	0.00	0.0		119.56
36	17,833	0.00	0.0		120.10
37	21,101	0.00	0.0		123.22
38	23,279	0.00	0.0		125.30
39	26,432	0.00	0.0		128.30
40	28,209	0.00	0.0		129.99
41	31,706	0.00	0.0		133.33
42	32,566	0.00	0.0		134.15
43	37,382	0.00	0.0		138.74
44	39,904	0.00	0.0		141.14
45	41,739	0.00	0.0		142.89

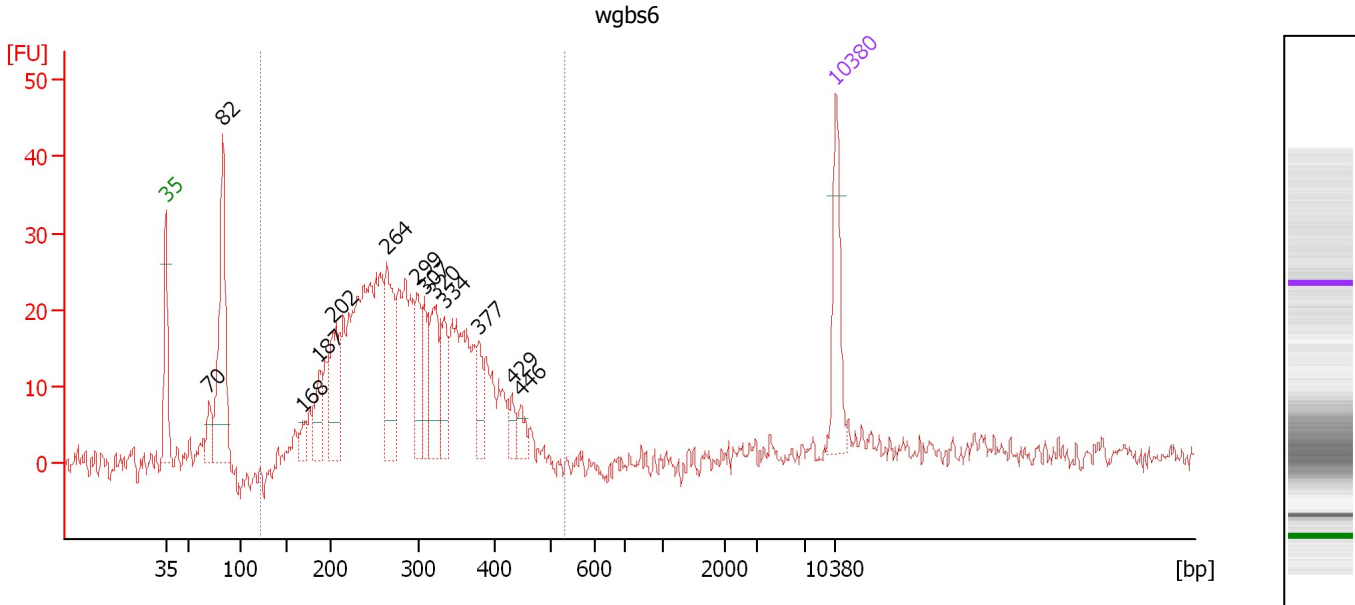
Region table for sample 6 : wgbs6

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
107	307	218	697.3	11,098.4	1,530.73	28	18.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 7 : wgs6

Number of peaks found: 13 Corr. Area 1: 508.2
 Noise: 1.6

Peak table for sample 7 : wgs6

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	70	39.77	861.1		47.57
3	82	270.46	4,968.0		48.96
4	168	21.22	191.0		57.31
5	187	58.79	477.0		59.04
6	202	113.23	848.2		60.49
7	264	146.23	840.5		66.05
8	299	73.95	374.5		69.28
9	307	63.10	311.1		69.95
10	320	89.76	424.5		71.01
11	334	73.41	332.9		72.12
12	377	44.77	179.7		75.63
13	429	17.94	63.3		79.14
14	446	20.51	69.7		80.09
15	10,380	75.00	10.9	Upper Marker	113.00

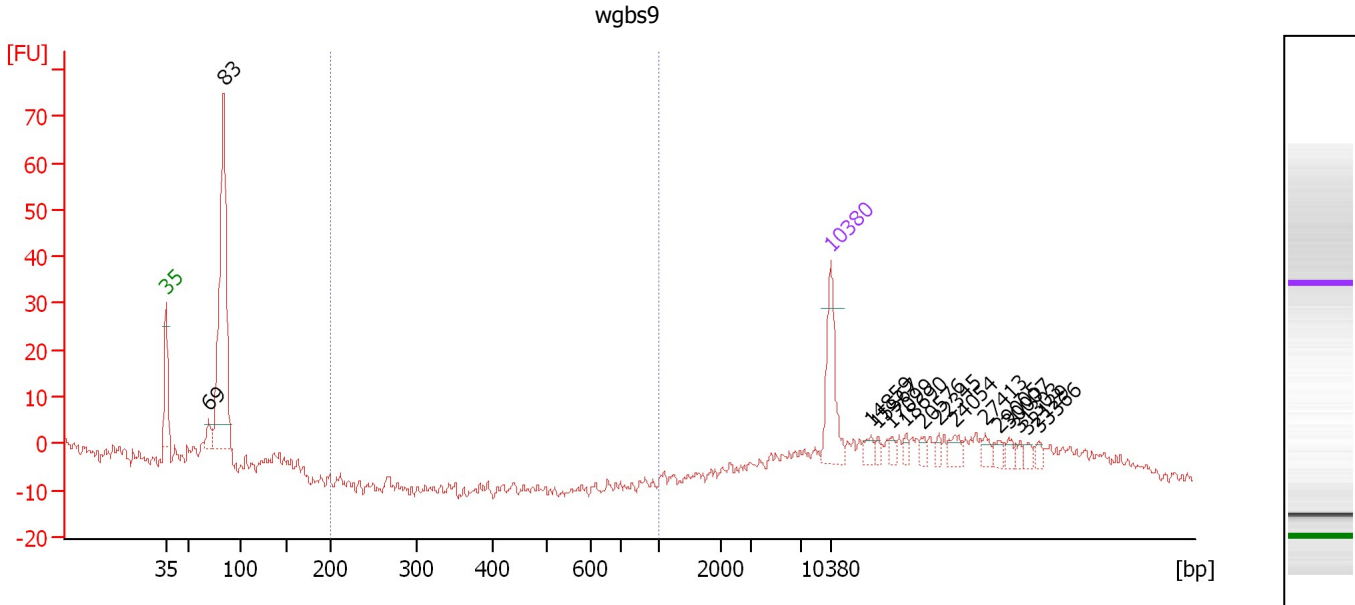
Region table for sample 7 : wgs6

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
120	532	291	508.2	9,666.6	1,720.21	84	23.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 8 : wgbs9

Number of peaks found: 15 Corr. Area 1: 0.0
 Noise: 1.2

Peak table for sample 8 : wgbs9

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	69	26.79	585.5		47.49
3	83	475.61	8,634.0		49.07
4	10,380	75.00	10.9	Upper Marker	113.00
5	14,859	0.00	0.0		117.27
6	15,567	0.00	0.0		117.94
7	17,099	0.00	0.0		119.40
8	18,690	0.00	0.0		120.92
9	20,576	0.00	0.0		122.72
10	22,345	0.00	0.0		124.40
11	24,054	0.00	0.0		126.03
12	27,413	0.00	0.0		129.24
13	29,005	0.00	0.0		130.75
14	30,007	0.00	0.0		131.71
15	31,303	0.00	0.0		132.94
16	32,129	0.00	0.0		133.73
17	33,366	0.00	0.0		134.91

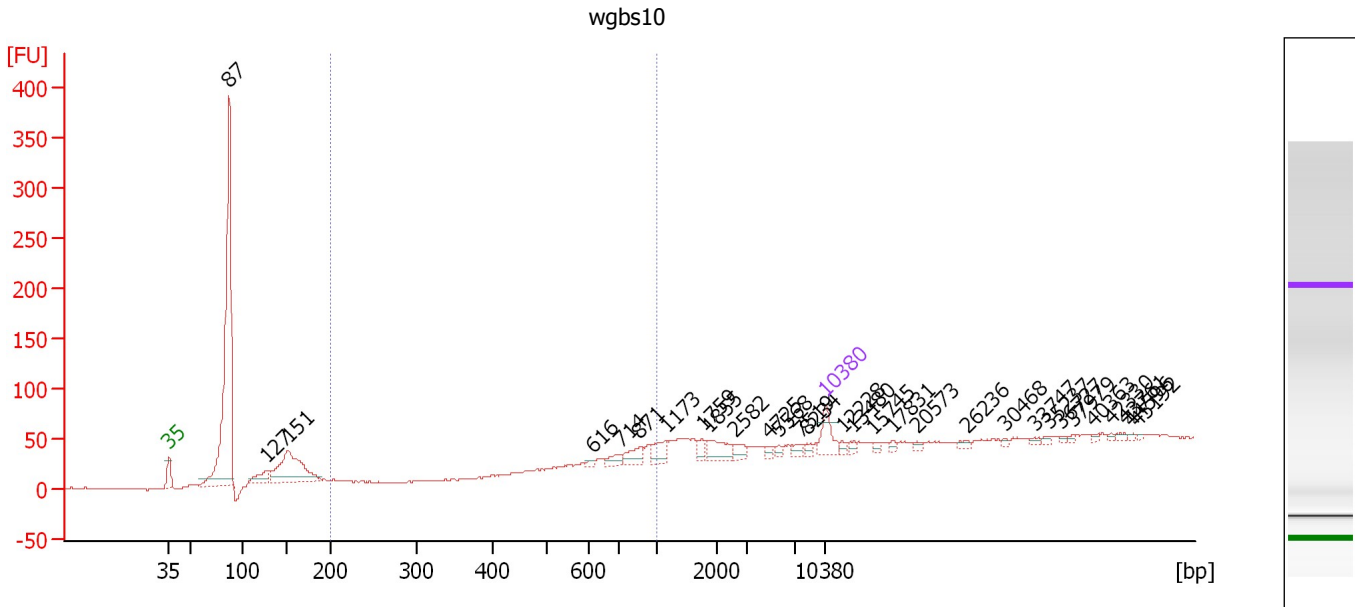
Region table for sample 8 : wgbs9

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:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 9 : wgbs10

Number of peaks found: 30 Corr. Area 1: 71.8
 Noise: 0.8

Peak table for sample 9 : wgbs10

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	87	1,532.14	26,795.5		49.42
3	127	87.28	1,044.9		53.40
4	151	405.51	4,074.0		55.67
5	616	11.91	29.3		88.23
6	714	33.53	71.2		91.07
7	871	59.07	102.7		93.17
8	1,173	59.43	76.8		96.01
9	1,759	28.40	24.5		99.82
10	1,855	69.28	56.6		100.44
11	2,582	30.54	17.9		103.28
12	4,725	13.51	4.3		106.86
13	5,568	12.22	3.3		107.94
14	7,519	19.11	3.9		110.27
15	8,234	14.85	2.7		110.95
16	10,380	75.00	10.9	Upper Marker	113.00
17	12,228	0.00	0.0		114.76
18	13,480	0.00	0.0		115.95
19	15,745	0.00	0.0		118.11
20	17,831	0.00	0.0		120.10
21	20,573	0.00	0.0		122.72
22	26,236	0.00	0.0		128.11
23	30,468	0.00	0.0		132.15
24	33,747	0.00	0.0		135.27
25	35,237	0.00	0.0		136.69

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad


Created: 2/19/2016 3:56:58 PM
 Modified: 2/19/2016 4:39:07 PM

Electropherogram Summary Continued ...

... Peak table for sample 9 : wgbs10

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	36,727	0.00	0.0		138.11
27	37,979	0.00	0.0		139.31
28	40,363	0.00	0.0		141.58
29	42,330	0.00	0.0		143.45
30	43,761	0.00	0.0		144.82
31	44,596	0.00	0.0		145.61
32	45,192	0.00	0.0		146.18

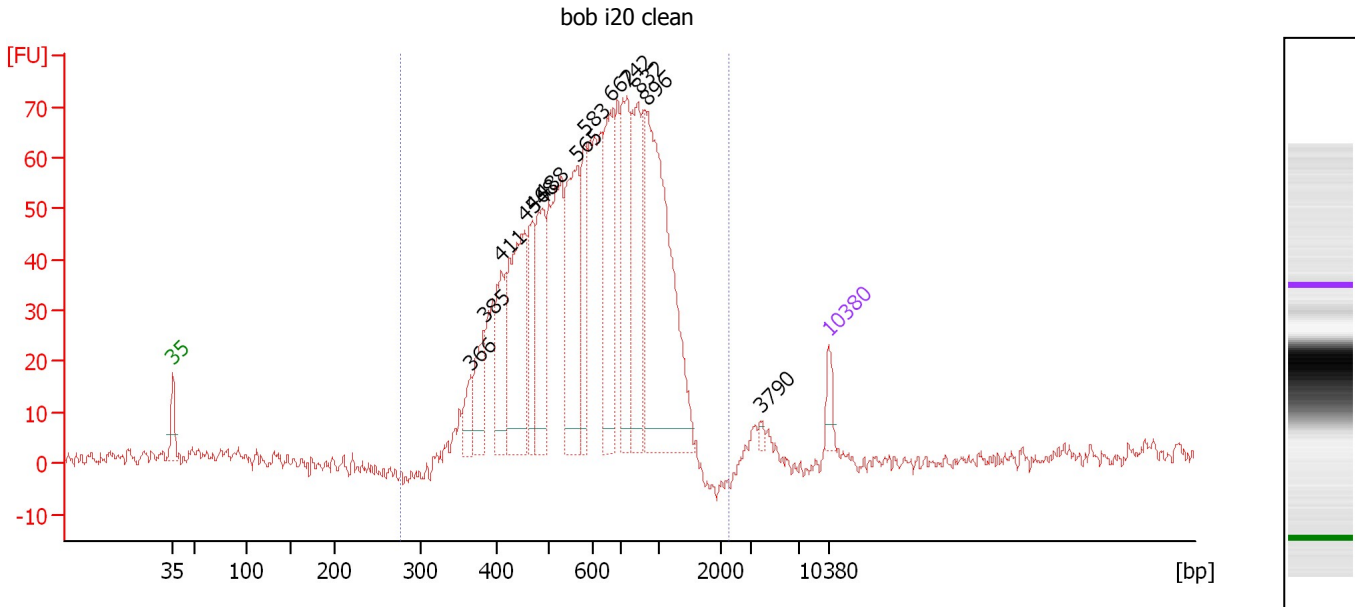
Region table for sample 9 : wgbs10

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	810	71.8	249.7	 130.46	7	14.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

Created: 2/19/2016 3:56:58 PM
 Modified: 2/19/2016 4:39:07 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : bob i20 clean

Number of peaks found: 13 Corr. Area 1: 1,293.6
 Noise: 1.7

Peak table for sample 10 : bob i20 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	366	152.51	630.9		74.73
3	385	271.83	1,071.1		76.21
4	411	410.56	1,513.9		78.09
5	454	900.37	3,006.8		80.53
6	468	350.04	1,134.1		81.33
7	488	516.03	1,603.6		82.46
8	565	779.49	2,092.1		86.10
9	583	396.68	1,030.4		86.96
10	662	651.01	1,489.7		89.69
11	742	667.27	1,362.4		91.45
12	832	563.38	1,026.5		92.64
13	896	1,514.75	2,562.9		93.50
14	3,790	22.70	9.1		105.66
15	10,380	75.00	10.9	Upper Marker	113.00

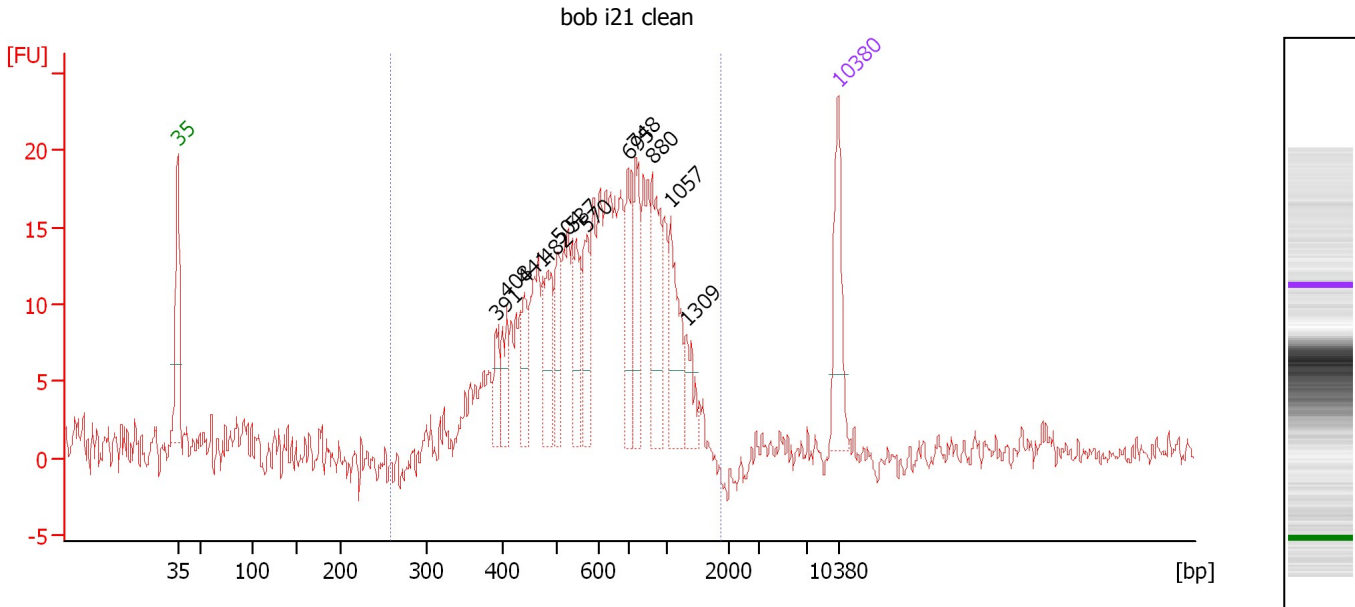
Region table for sample 10 : bob i20 clean

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
278	2,301	668	1,293.6	26,826.0	10,214.59	93	36.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

Created: 2/19/2016 3:56:58 PM
 Modified: 2/19/2016 4:39:07 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : bob i21 clean

Number of peaks found: 12 Corr. Area 1: 267.1
 Noise: 0.8

Peak table for sample 11 : bob i21 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	391	36.33	140.8		76.73
3	408	43.30	160.8		77.92
4	441	53.09	182.6		79.78
5	482	85.31	268.1		82.15
6	504	61.73	185.7		83.34
7	537	79.43	224.0		84.86
8	570	64.84	172.5		86.33
9	695	76.43	166.5		90.74
10	748	87.21	176.6		91.53
11	880	110.92	191.1		93.28
12	1,057	88.70	127.1		95.26
13	1,309	36.01	41.7		96.90
14	10,380	75.00	10.9	Upper Marker	113.00

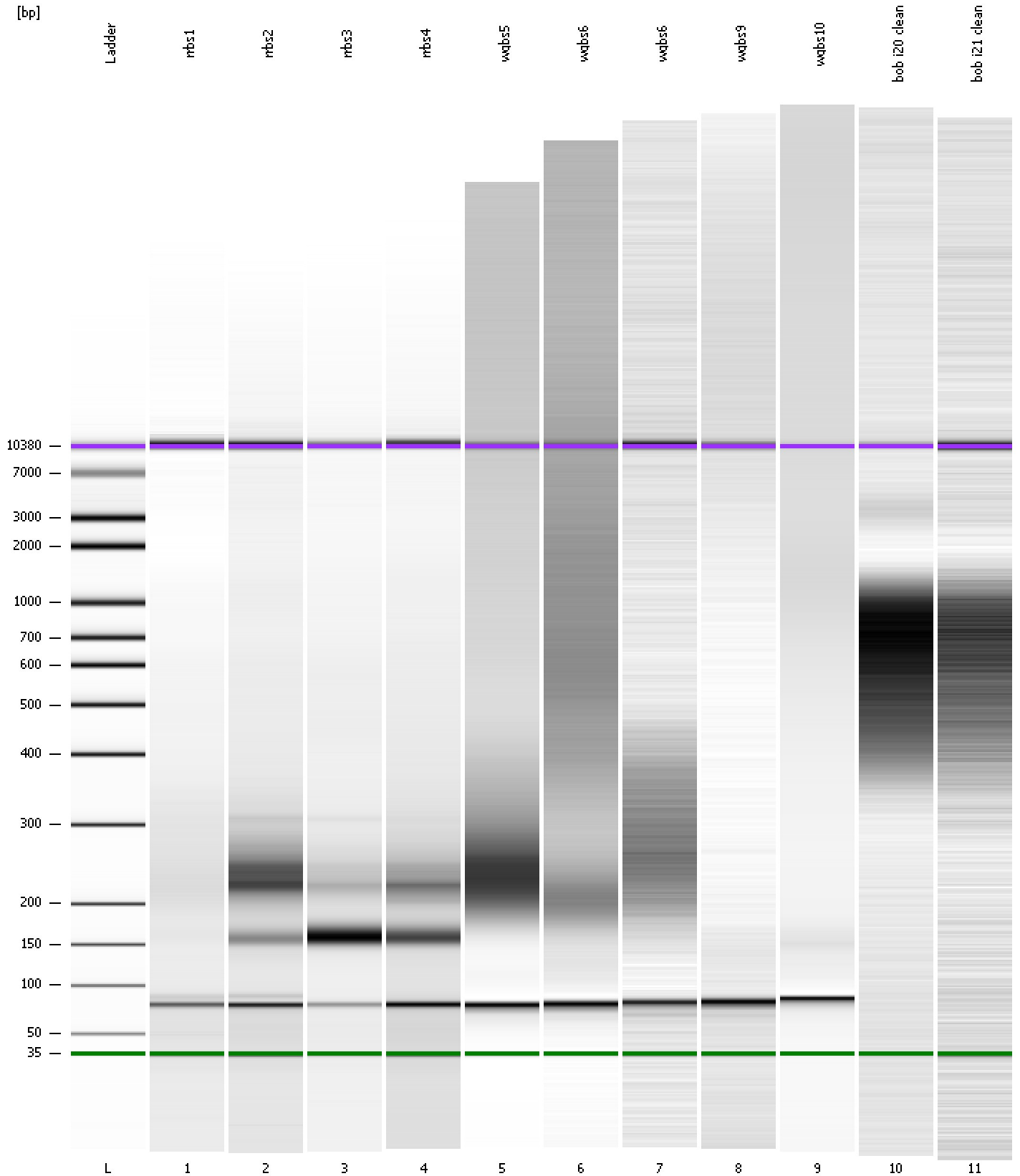
Region table for sample 11 : bob i21 clean

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
259	1,883	679	267.1	3,699.0	1,446.30	100	35.0

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

Created: 2/19/2016 3:56:58 PM
Modified: 2/19/2016 4:39:07 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad

Created: 2/19/2016 3:56:58 PM
 Modified: 2/19/2016 4:39:07 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		2/19/2016 4:38:16 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2016-02-19\2016-02-19_001.xad)		Instrument	Run		2/19/2016 3:57:04 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		2/19/2016 3:57:04 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		2/19/2016 3:57:04 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		2/19/2016 3:57:04 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		2/19/2016 3:57:04 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		2/19/2016 3:57:04 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		2/19/2016 3:57:04 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1