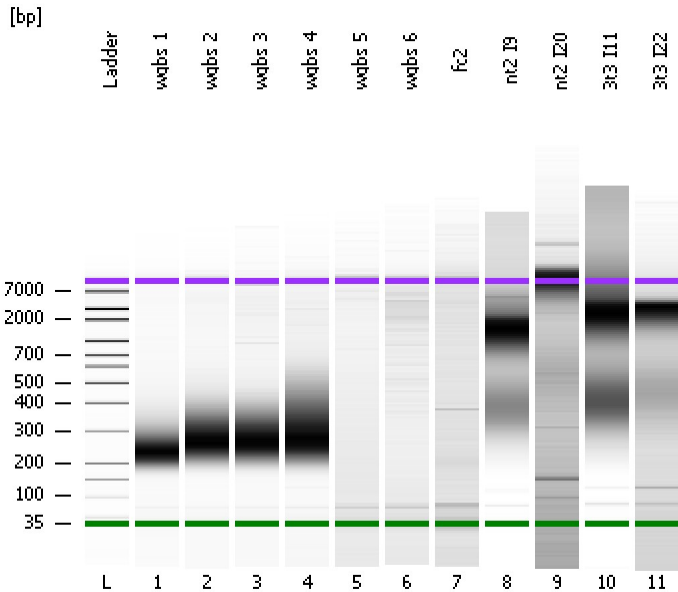


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

Created: 11/5/2015 11:16:21 AM
Modified: 11/5/2015 11:58:21 AM

Electrophoresis File Run Summary



Instrument Information:

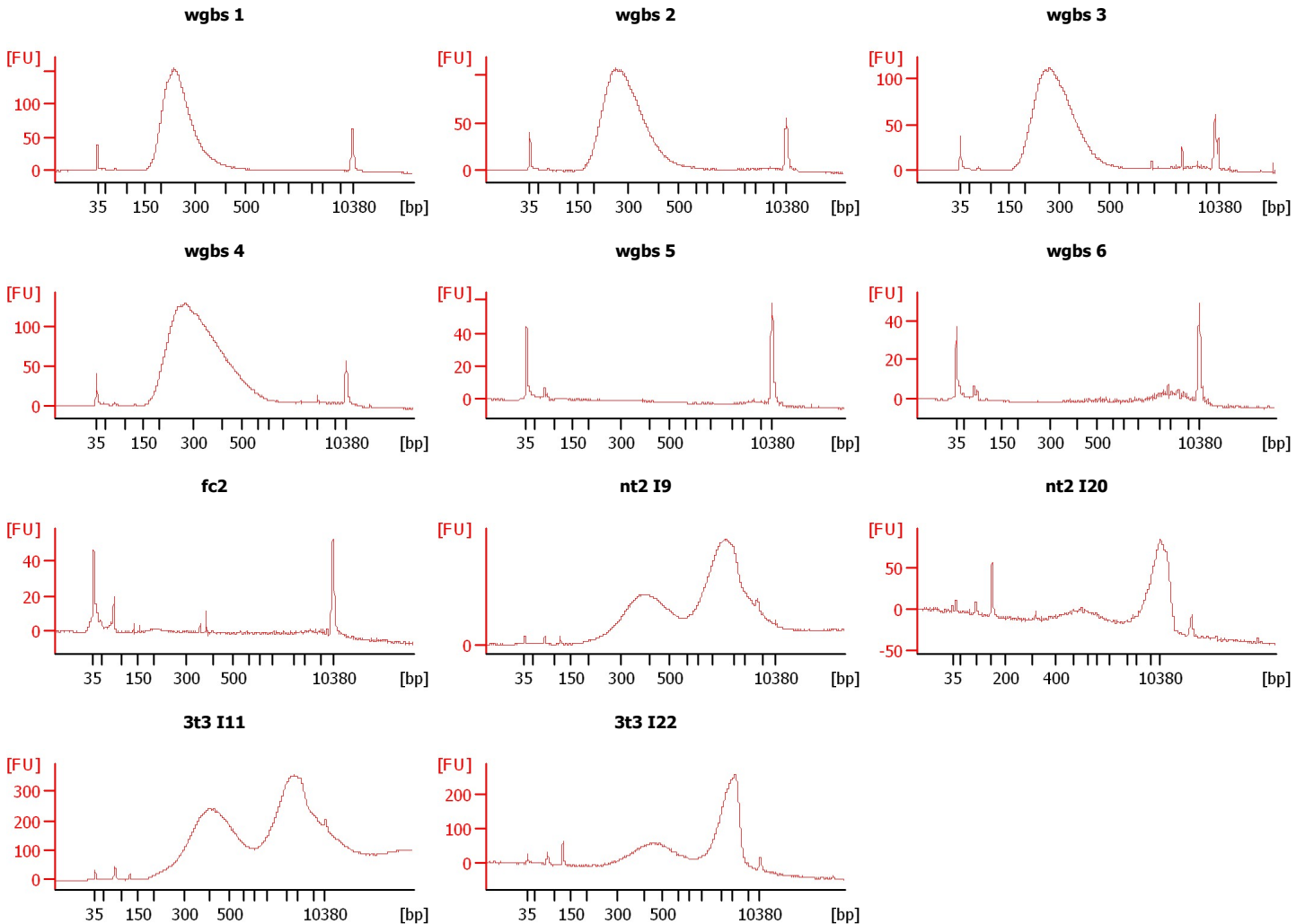
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

Created: 11/5/2015 11:16:21 AM
 Modified: 11/5/2015 11:58:21 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
wgbs 1		<input type="checkbox"/>	✓			
wgbs 2		<input type="checkbox"/>	✓			
wgbs 3		<input type="checkbox"/>	✓			
wgbs 4		<input type="checkbox"/>	✓			
wgbs 5		<input type="checkbox"/>	✓			
wgbs 6		<input type="checkbox"/>	✓			
fc2		<input type="checkbox"/>	✓			
nt2 I9		<input type="checkbox"/>	✓			
nt2 I20		<input type="checkbox"/>	✓			
3t3 I11		<input type="checkbox"/>	✓			
3t3 I22		<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\2015-11-05\2015-11-05_003.xad

Created: 11/5/2015 11:16:21 AM
Modified: 11/5/2015 11:58:21 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

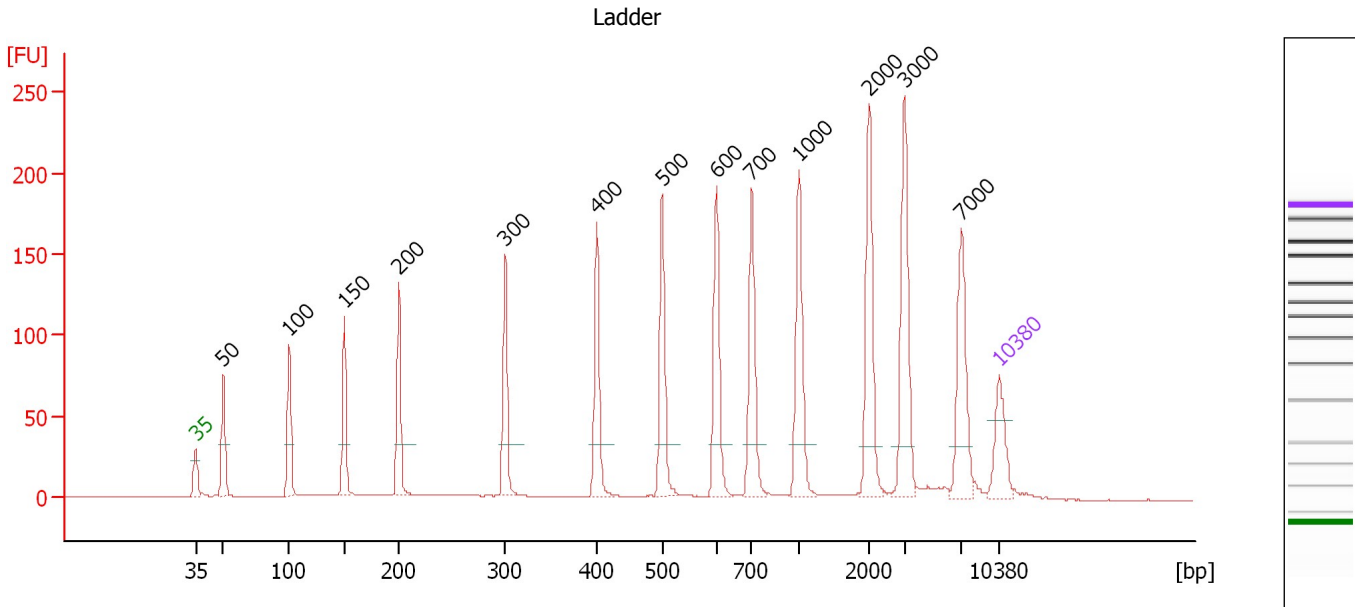
Ladder

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:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

Created: 11/5/2015 11:16:21 AM
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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

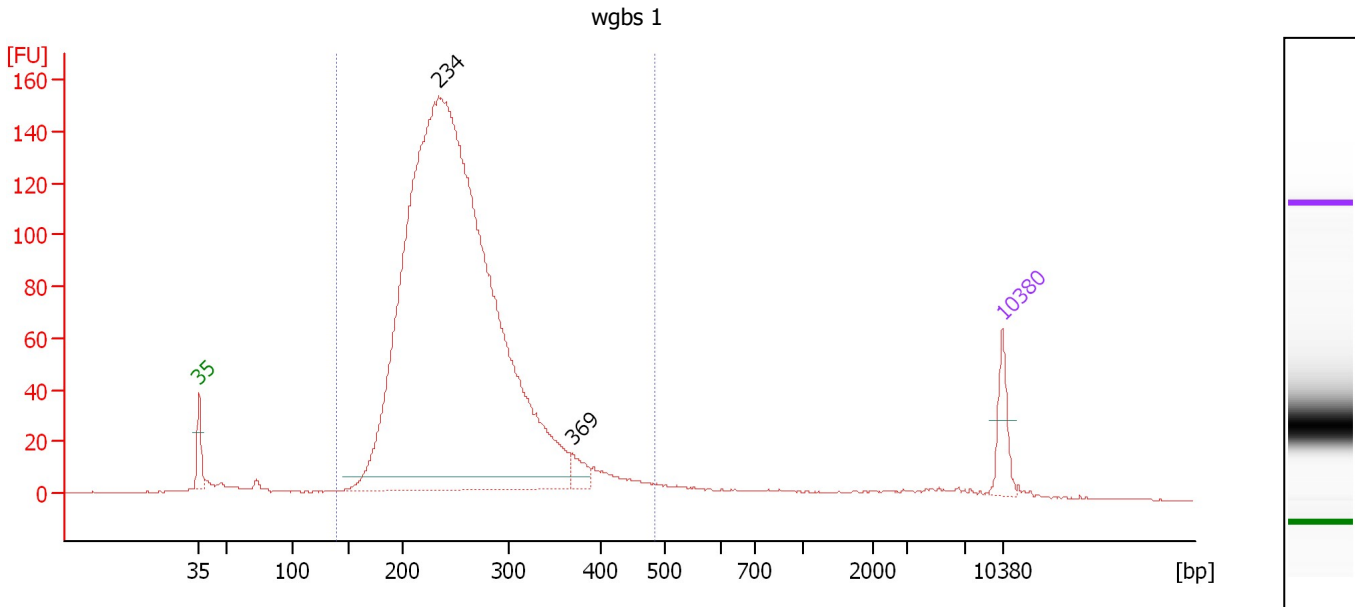
Peak table for Ladder

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	45.42
3	100	150.00	2,272.7	Ladder Peak	51.18
4	150	150.00	1,515.2	Ladder Peak	55.96
5	200	150.00	1,136.4	Ladder Peak	60.70
6	300	150.00	757.6	Ladder Peak	69.99
7	400	150.00	568.2	Ladder Peak	77.98
8	500	150.00	454.5	Ladder Peak	83.64
9	600	150.00	378.8	Ladder Peak	88.38
10	700	150.00	324.7	Ladder Peak	91.45
11	1,000	150.00	227.3	Ladder Peak	95.53
12	2,000	150.00	113.6	Ladder Peak	101.67
13	3,000	150.00	75.8	Ladder Peak	104.73
14	7,000	150.00	32.5	Ladder Peak	109.70
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

Created: 11/5/2015 11:16:21 AM
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Electropherogram Summary Continued ...



Overall Results for sample 1 : wgbs 1

Number of peaks found: 2 Corr. Area 1: 2,283.2
 Noise: 0.2

Peak table for sample 1 : wgbs 1

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	234	5,310.39	34,436.5		63.82
3	369	53.63	220.3		75.49
4	10,380	75.00	10.9	Upper Marker	113.00

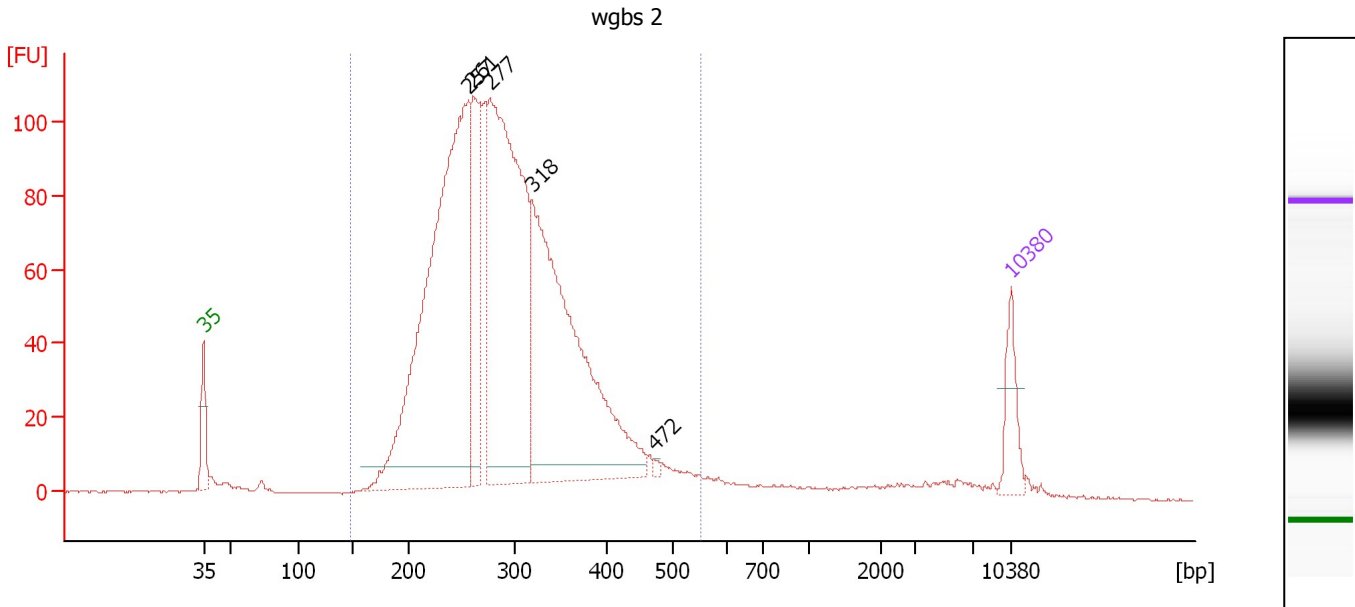
Region table for sample 1 : wgbs 1

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
140	483	256	2,283.2	33,978.0	5,499.84	93	20.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 2 : wgbs 2

Number of peaks found: 5 Corr. Area 1: 2,040.1
 Noise: 0.3

Peak table for sample 2 : wgbs 2

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	257	1,577.32	9,310.4		65.96
3	261	301.24	1,747.7		66.38
4	277	1,222.67	6,697.9		67.81
5	318	1,122.03	5,338.4		71.46
6	472	6.62	21.2		82.04
7	10,380	75.00	10.9	Upper Marker	113.00

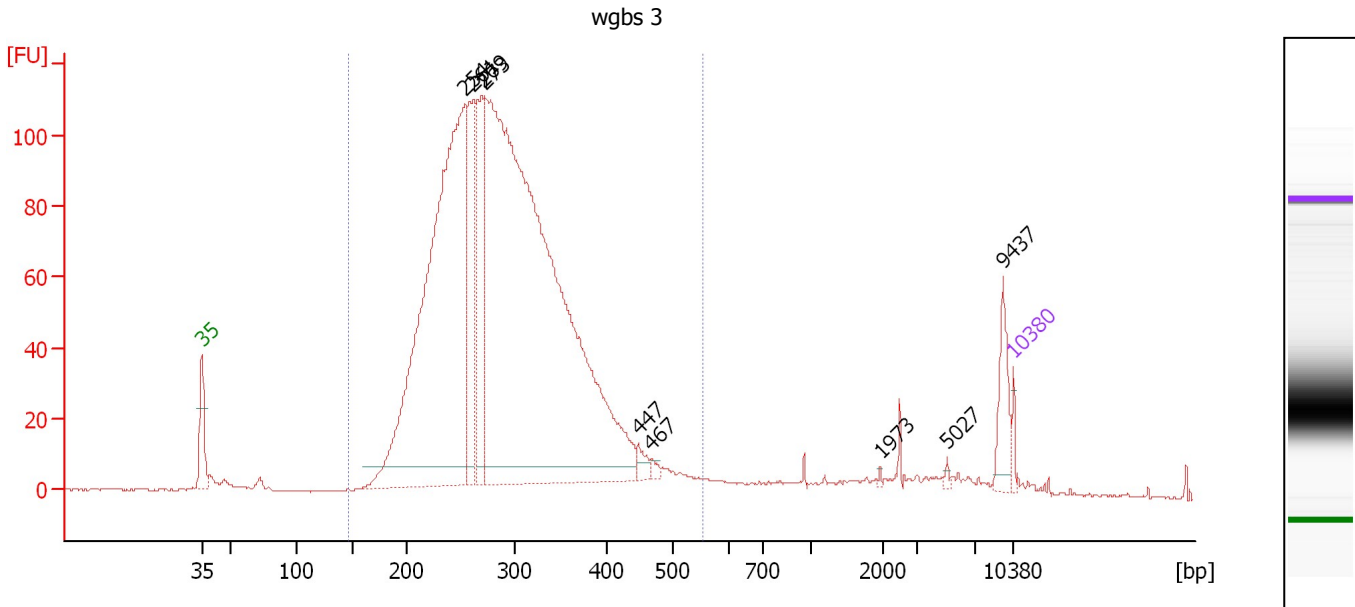
Region table for sample 2 : wgbs 2

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
148	551	295	2,040.1	25,880.3	4,753.40	94	22.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 3 : wgbs 3

Number of peaks found: 9 Corr. Area 1: 2,098.0
 Noise: 0.2

Peak table for sample 3 : wgbs 3

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	254	8,465.04	50,580.8		65.67
3	261	1,568.43	9,105.1		66.36
4	269	1,355.70	7,624.3		67.15
5	273	14,204.62	78,870.8		67.47
6	447	125.02	423.3		80.67
7	467	54.70	177.5		81.77
8	1,973	13.84	10.6		101.50
9	5,027	23.52	7.1		107.25
10	9,437	348.29	55.9		112.08
11	10,380	75.00	10.9	Upper Marker	113.00

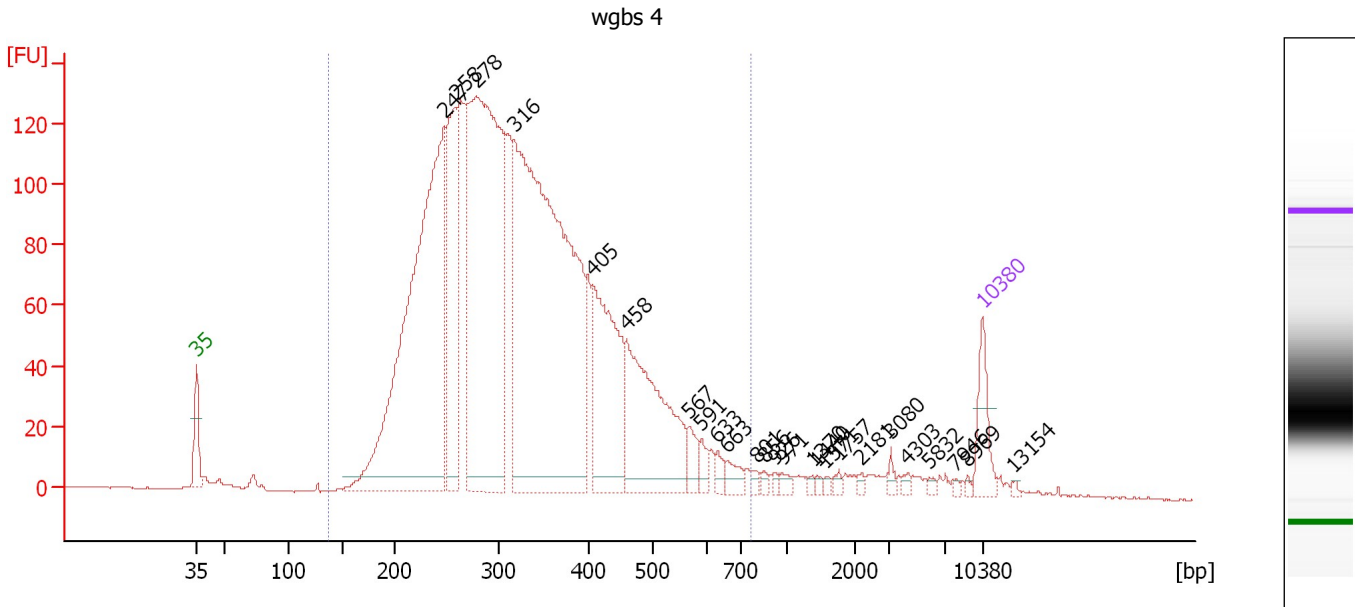
Region table for sample 3 : wgbs 3

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
147	554	292	2,098.0	142,794.7	26,120.53	94	21.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 4 : wgbs 4

Number of peaks found: 25 Corr. Area 1: 3,242.5
 Noise: 0.2

Peak table for sample 4 : wgbs 4

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	247	1,637.25	10,055.7		65.04
3	258	477.72	2,806.1		66.08
4	278	1,458.27	7,936.7		67.98
5	316	1,957.78	9,372.9		71.30
6	405	446.85	1,669.9		78.28
7	458	476.92	1,576.9		81.28
8	567	48.12	128.5		86.83
9	591	29.22	74.9		87.97
10	633	22.47	53.8		89.40
11	663	33.31	76.2		90.30
12	801	11.66	22.1		92.82
13	856	11.25	19.9		93.58
14	926	7.86	12.9		94.53
15	971	14.82	23.1		95.14
16	1,370	6.80	7.5		97.80
17	1,440	6.79	7.1		98.23
18	1,571	6.66	6.4		99.04
19	1,757	9.02	7.8		100.18
20	2,181	7.35	5.1		102.22
21	3,080	10.84	5.3		104.83
22	4,303	8.59	3.0		106.35
23	5,832	6.43	1.7		108.25
24	7,946	5.11	1.0		110.63
25	8,969	4.96	0.8		111.62

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

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Electropherogram Summary Continued ...**... Peak table for sample 4 : wgbs 4**

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	10,380	75.00	10.9	Upper Marker	113.00
27	13,154	0.00	0.0		115.71

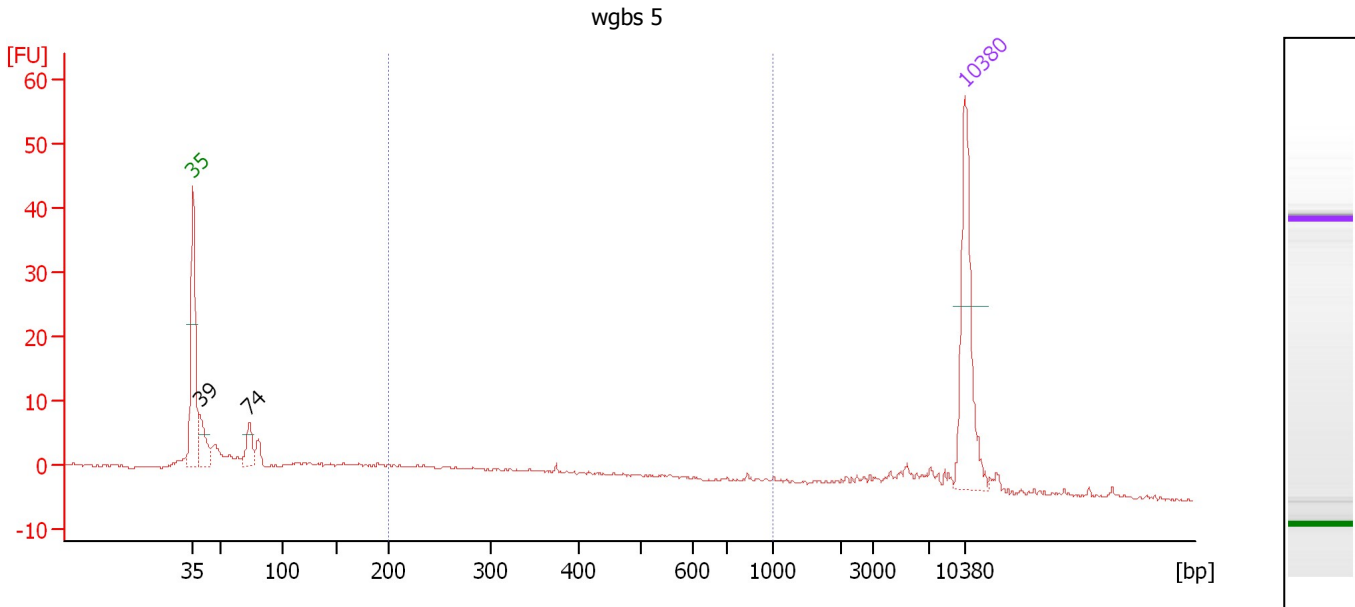
Region table for sample 4 : wgbs 4

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
138	766	332	3,242.5	37,266.3	7,327.05	95	29.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 5 : wgbs 5

Number of peaks found: 2 Corr. Area 1: 54.7
 Noise: 0.2

Peak table for sample 5 : wgbs 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	39	38.08	1,483.2		43.63
3	74	24.88	512.7		48.12
4	10,380	75.00	10.9	Upper Marker	113.00

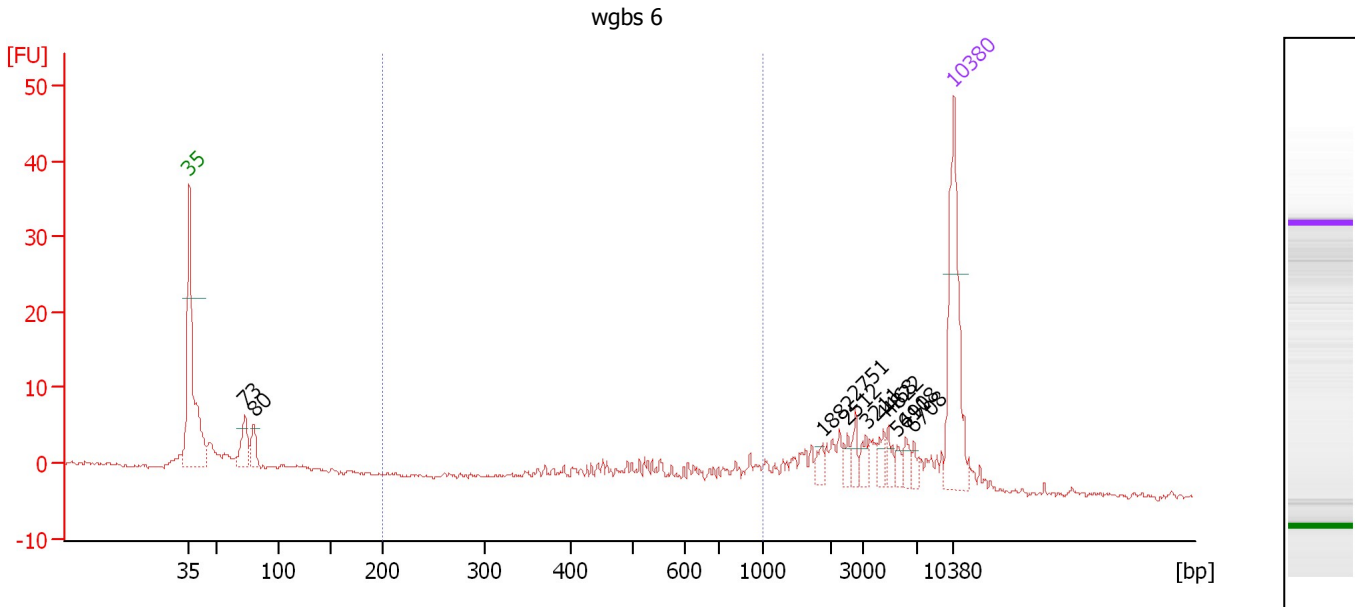
Region table for sample 5 : wgbs 5

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	429	54.7	488.7	109.13	34	44.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 6 : wgbs 6

Number of peaks found: 11 Corr. Area 1: 33.1
 Noise: 0.3

Peak table for sample 6 : wgbs 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	73	31.50	652.8		48.08
3	80	20.05	378.3		48.91
4	1,882	6.91	5.6		100.94
5	2,512	6.34	3.8		103.24
6	2,751	7.80	4.3		103.97
7	3,211	7.75	3.7		104.99
8	4,468	8.42	2.9		106.56
9	4,822	7.64	2.4		107.00
10	5,490	4.97	1.4		107.83
11	6,118	7.25	1.8		108.61
12	6,708	4.80	1.1		109.34
13	10,380	75.00	10.9	Upper Marker	113.00

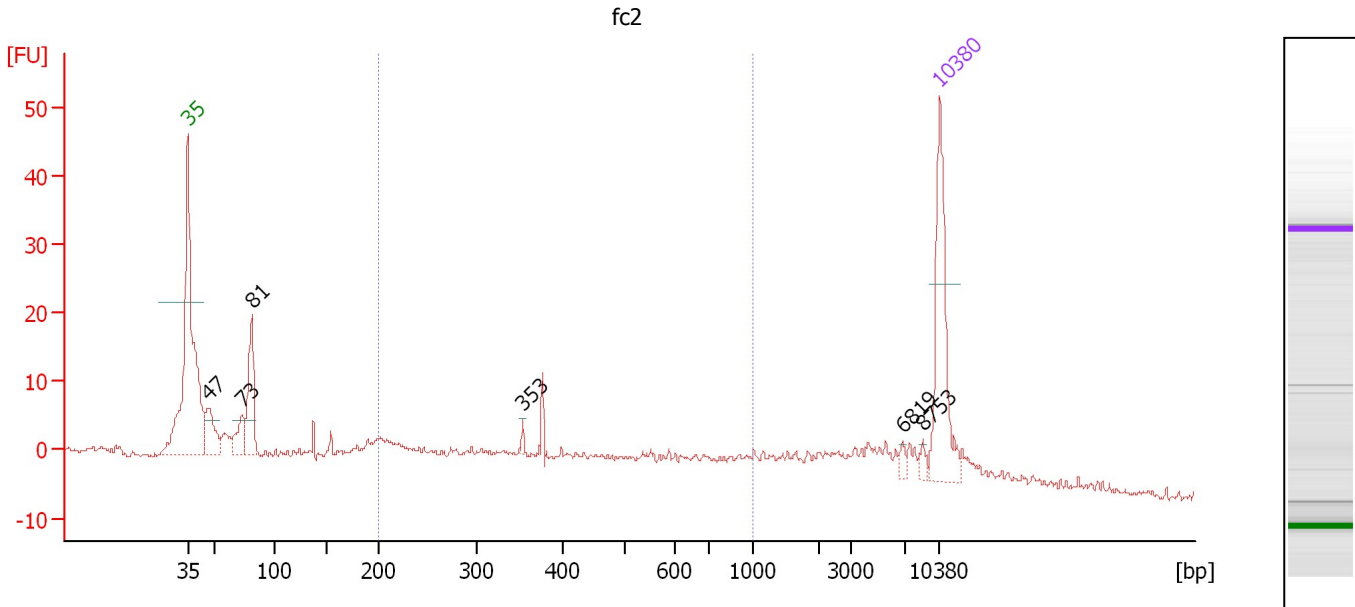
Region table for sample 6 : wgbs 6

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	604	33.1	199.7	70.87	23	30.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 7 : fc2

Number of peaks found: 6 Corr. Area 1: 119.5
 Noise: 0.5

Peak table for sample 7 : fc2

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	47	44.08	1,420.6		44.93
3	73	30.45	632.4		48.06
4	81	68.85	1,292.5		48.95
5	353	3.56	15.3		74.20
6	6,819	4.17	0.9		109.48
7	8,753	4.38	0.8		111.41
8	10,380	75.00	10.9	Upper Marker	113.00

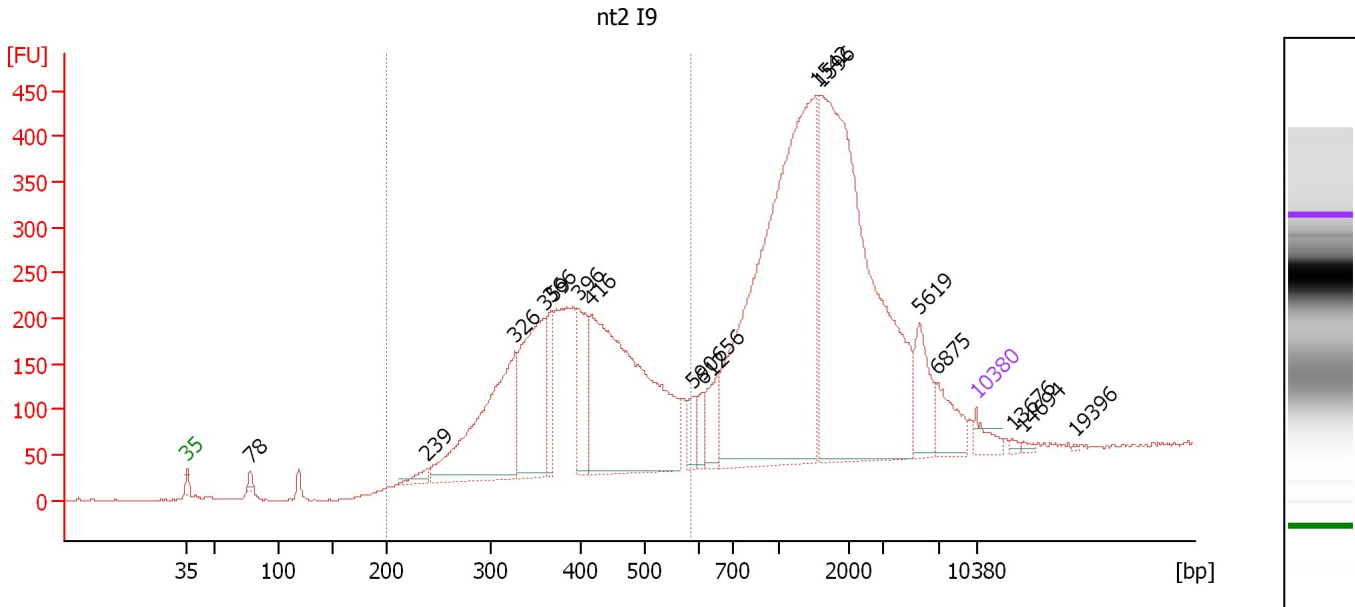
Region table for sample 7 : fc2

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	464	119.5	1,056.5	249.27	39	43.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 8 : nt2 I9

Number of peaks found: 17 Corr. Area 1: 3,571.4
 Noise: 1.0

Peak table for sample 8 : nt2 I9

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	78	45.88	892.5		48.63
3	239	64.77	410.4		64.33
4	326	1,311.07	6,088.0		72.09
5	359	937.36	3,951.2		74.74
6	366	249.41	1,032.6		75.26
7	396	410.56	1,570.3		77.67
8	416	2,065.03	7,516.4		78.90
9	590	113.43	291.5		87.89
10	612	91.10	225.7		88.74
11	656	210.46	485.9		90.11
12	1,542	2,945.00	2,893.7		98.86
13	1,596	2,726.68	2,588.5		99.19
14	5,619	260.76	70.3		107.99
15	6,875	189.89	41.8		109.55
16	10,380	75.00	10.9	Upper Marker	113.00
17	13,676	0.00	0.0		116.22
18	14,694	0.00	0.0		117.21
19	19,396	0.00	0.0		121.80

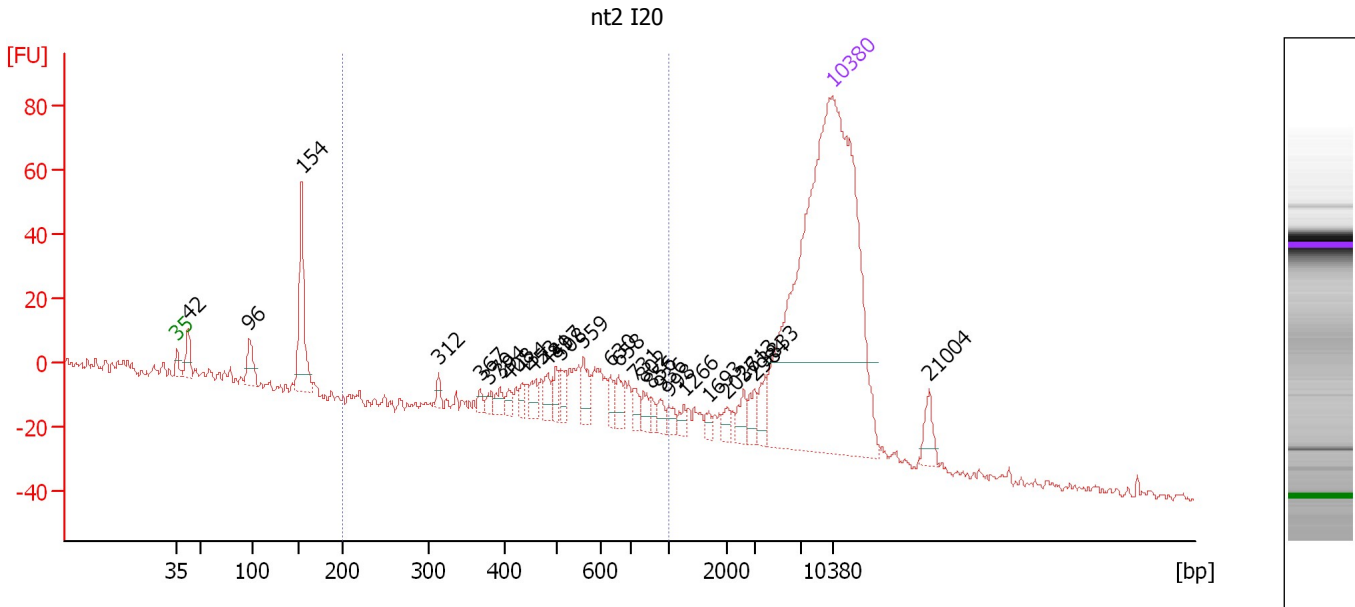
Region table for sample 8 : nt2 I9

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	585	399	3,571.4	23,840.8	5,923.51	40	20.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 9 : nt2 I20

Number of peaks found: 28 Corr. Area 1: 317.3
 Noise: 1.8

Peak table for sample 9 : nt2 I20

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	42	4.34	156.4		44.14
3	96	4.50	70.8		50.75
4	154	14.53	143.1		56.33
5	312	1.16	5.7		70.91
6	367	0.90	3.7		75.35
7	379	1.05	4.2		76.26
8	394	1.43	5.5		77.52
9	408	1.17	4.3		78.43
10	434	1.36	4.8		79.91
11	453	1.98	6.6		80.99
12	481	2.27	7.2		82.59
13	497	2.00	6.1		83.50
14	508	2.22	6.6		84.01
15	559	3.23	8.8		86.46
16	630	1.71	4.1		89.31
17	658	2.45	5.6		90.16
18	731	1.61	3.3		91.87
19	802	1.59	3.0		92.84
20	856	1.06	1.9		93.58
21	936	1.24	2.0		94.66
22	998	1.13	1.7		95.51
23	1,266	1.01	1.2		97.17
24	1,693	0.75	0.7		99.79
25	2,037	0.96	0.7		101.78

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

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Electropherogram Summary Continued ...

... Peak table for sample 9 : nt2 I20

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,613	1.83	1.1		103.55
27	2,984	1.79	0.9		104.68
28	3,833	2.25	0.9		105.77
29	10,380	75.00	10.9	Upper Marker	113.00
30	21,004	0.00	0.0		123.37

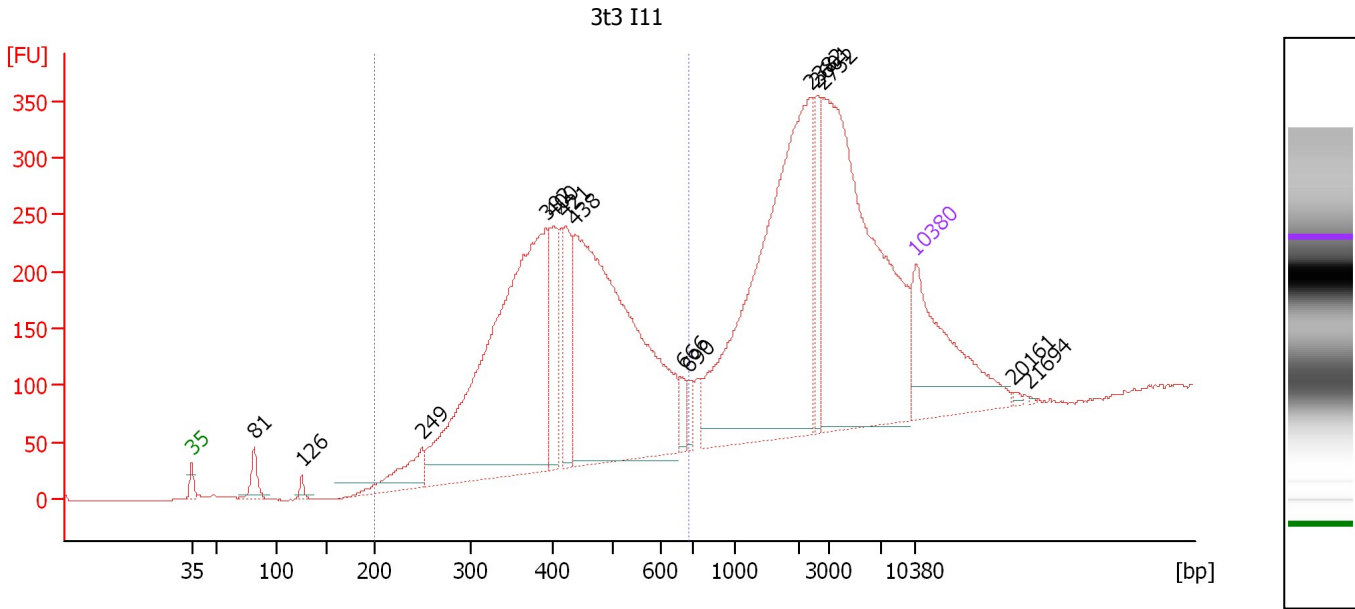
Region table for sample 9 : nt2 I20

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
200	1,000	560	317.3	148.8	48.76	48	28.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 10 : 3t3 I11

Number of peaks found: 14 Corr. Area 1: 3,675.5
 Noise: 0.3

Peak table for sample 10 : 3t3 I11

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	20.05	373.1		49.04
3	126	5.73	69.1		53.63
4	249	38.41	233.4		65.28
5	392	375.33	1,452.3		77.30
6	400	53.13	201.3		77.97
7	421	51.33	184.8		79.16
8	438	341.91	1,182.3		80.14
9	666	8.93	20.3		90.41
10	690	8.59	18.9		91.13
11	2,382	262.10	166.7		102.84
12	2,601	28.40	16.5		103.51
13	2,752	241.29	132.8		103.97
14	10,380	75.00	10.9	Upper Marker	113.00
15	20,161	0.00	0.0		122.54
16	21,694	0.00	0.0		124.04

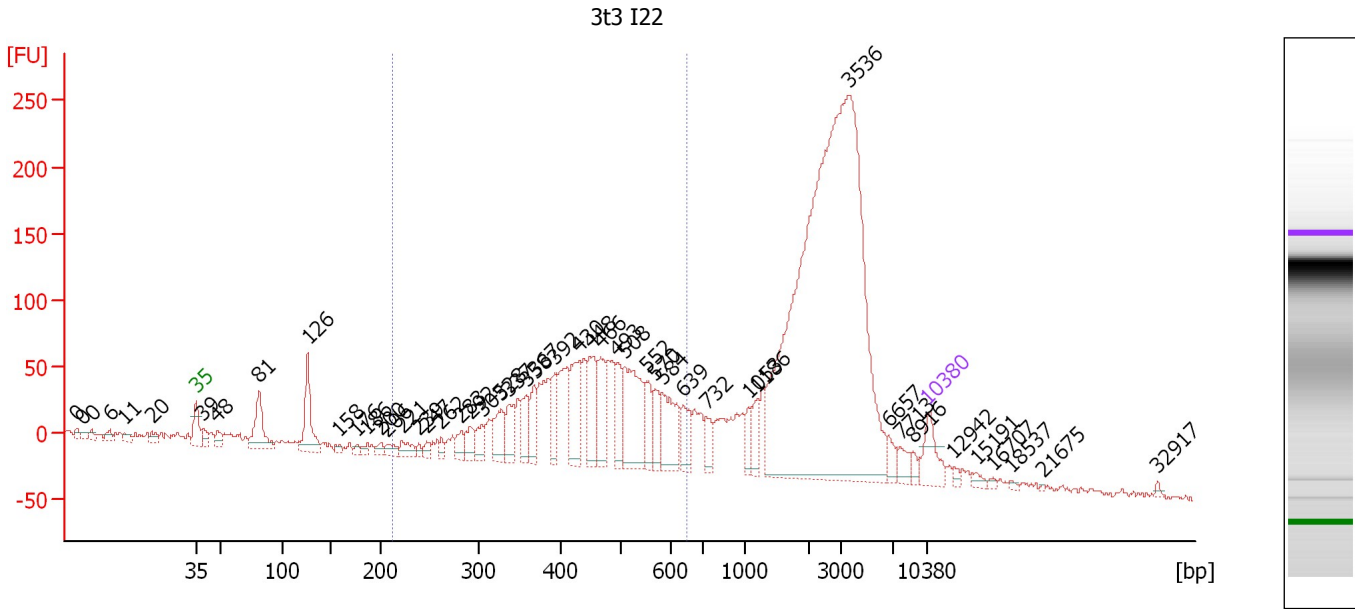
Region table for sample 10 : 3t3 I11

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	686	429	3,675.5	2,893.7	768.14	51	21.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

Created: 11/5/2015 11:16:21 AM
 Modified: 11/5/2015 11:58:21 AM

Electropherogram Summary Continued ...



Overall Results for sample 11 : 3t3 I22

Number of peaks found: 49 Corr. Area 1: 1,526.4
 Noise: 2.6

Peak table for sample 11 : 3t3 I22

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	0	0.00	0.0		31.52
2	0	0.00	0.0		32.03
3	0	0.00	0.0		33.00
4	6	0.00	0.0		34.79
5	11	0.00	0.0		36.11
6	20	0.00	0.0		38.77
7	35	125.00	5,411.3	Lower Marker	43.00
8	39	41.34	1,601.3		43.66
9	48	39.56	1,257.2		45.04
10	81	175.45	3,269.9		49.02
11	126	190.88	2,295.3		53.66
12	158	16.39	157.1		56.72
13	176	13.67	117.8		58.41
14	186	16.88	137.5		59.38
15	200	21.51	163.3		60.65
16	206	29.71	218.4		61.27
17	221	44.70	305.8		62.69
18	239	15.76	99.9		64.33
19	247	20.76	127.5		65.04
20	262	23.39	135.2		66.47
21	283	45.52	243.7		68.41
22	292	55.11	285.6		69.28
23	305	55.34	274.8		70.40
24	328	99.62	460.0		72.23
25	337	85.39	383.8		72.95

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

Created: 11/5/2015 11:16:21 AM
 Modified: 11/5/2015 11:58:21 AM

Electropherogram Summary Continued ...

... Peak table for sample 11 : 3t3 I22

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	358	85.47	362.2		74.58
27	367	81.39	335.9		75.35
28	392	88.77	343.1		77.34
29	430	157.24	553.9		79.68
30	448	148.61	502.5		80.70
31	466	122.20	397.2		81.72
32	493	93.72	288.0		83.26
33	508	267.99	799.4		84.02
34	552	76.64	210.3		86.11
35	570	69.37	184.3		86.98
36	584	155.86	404.1		87.64
37	639	79.75	189.1		89.58
38	732	57.61	119.3		91.88
39	1,053	43.47	62.6		95.86
40	1,186	47.05	60.1		96.67
41	3,536	2,143.47	918.5		105.40
42	6,657	33.76	7.7		109.28
43	7,713	32.11	6.3		110.40
44	8,916	18.19	3.1		111.57
45	10,380	75.00	10.9	Upper Marker	113.00
46	12,942	0.00	0.0		115.50
47	15,191	0.00	0.0		117.69
48	16,707	0.00	0.0		119.17
49	18,537	0.00	0.0		120.96
50	21,675	0.00	0.0		124.02
51	32,917	0.00	0.0		134.99

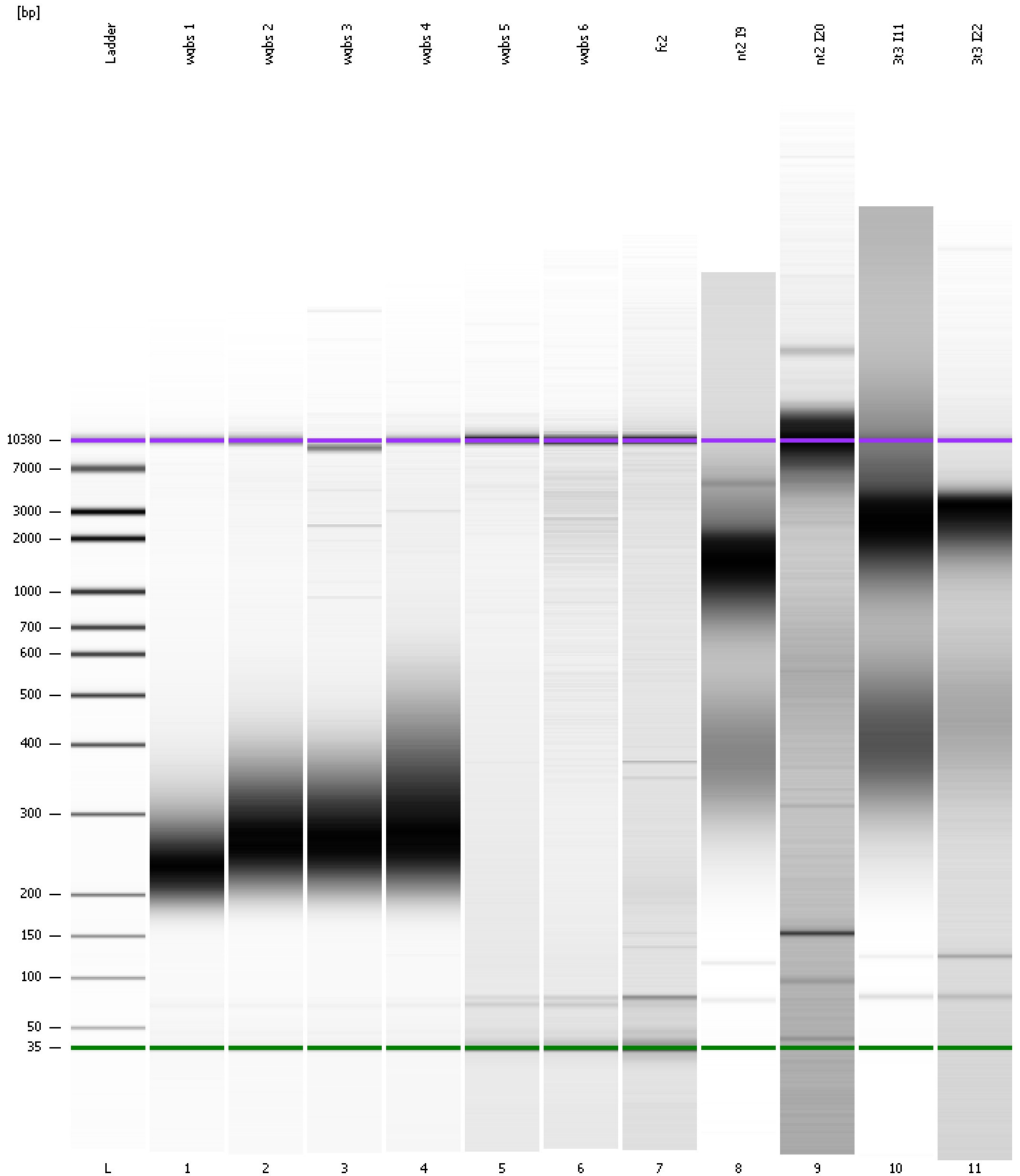
Region table for sample 11 : 3t3 I22

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
212	652	446	1,526.4	8,870.5	2,413.37	37	21.8

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

Created: 11/5/2015 11:16:21 AM
Modified: 11/5/2015 11:58:21 AM

Gel Image

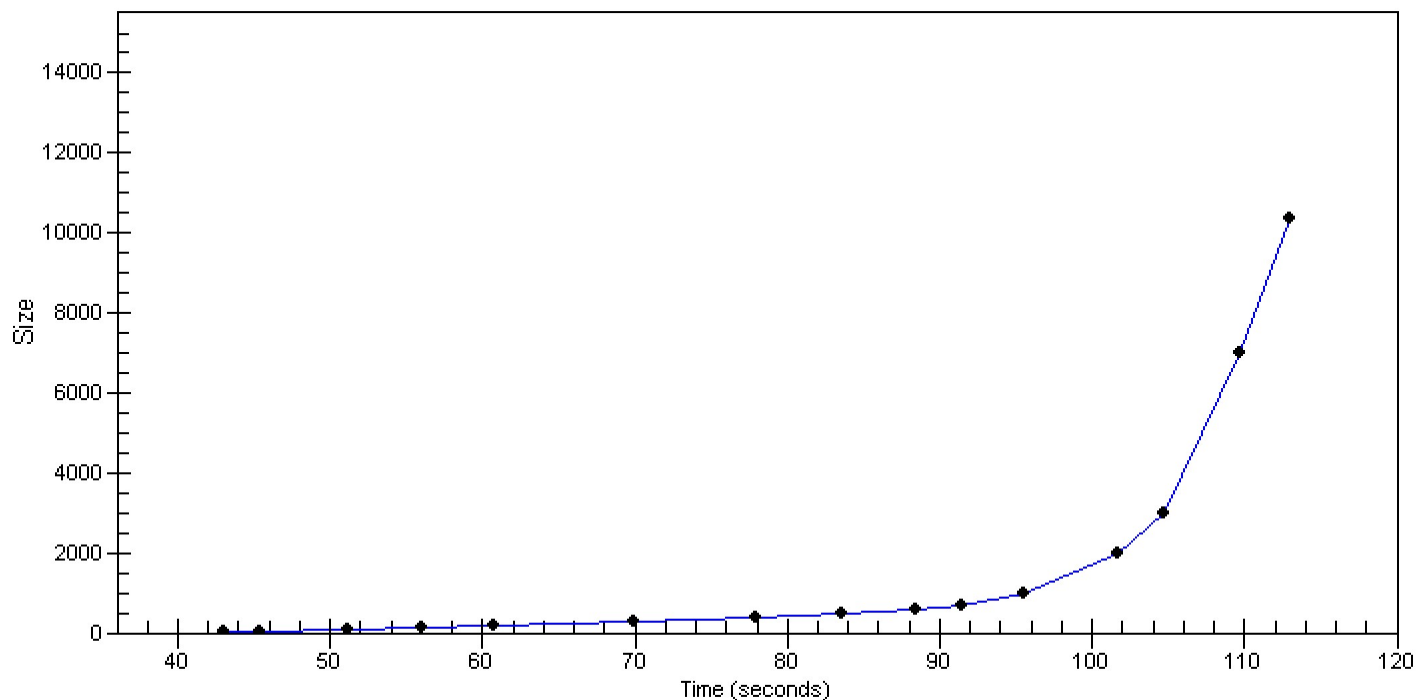


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

Created: 11/5/2015 11:16:21 AM
Modified: 11/5/2015 11:58:21 AM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad

Created: 11/5/2015 11:16:21 AM
 Modified: 11/5/2015 11:58:21 AM

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		11/5/2015 9:57:38 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-11-05\2015-11-05_003.xad)		Instrument	Run		11/5/2015 9:16:27 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		11/5/2015 9:16:27 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		11/5/2015 9:16:27 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		11/5/2015 9:16:27 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		11/5/2015 9:16:27 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		11/5/2015 9:16:27 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		11/5/2015 9:16:27 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1