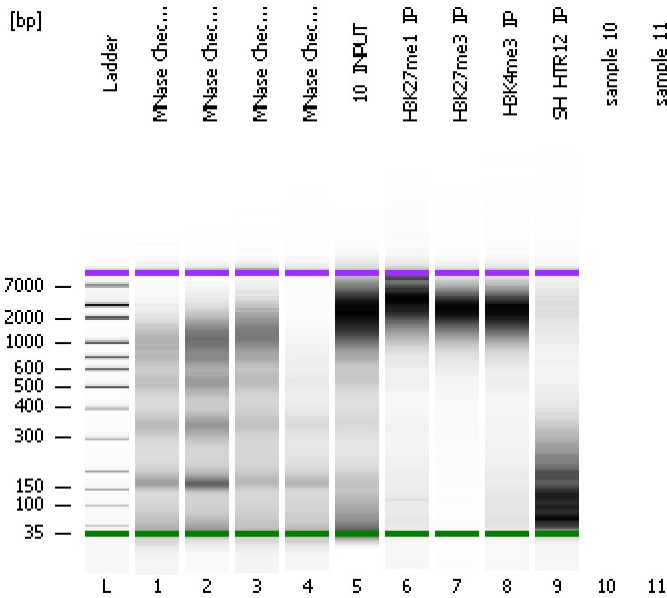


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
Modified: 2/4/2015 2:35:06 PM

Electrophoresis File Run Summary



Instrument Information:

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

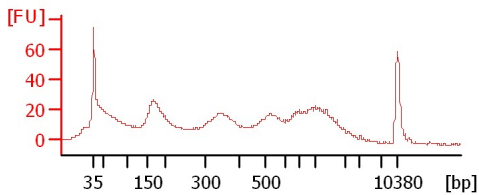
Assay Information:

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

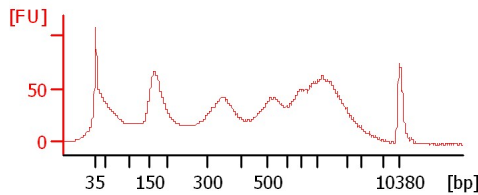
Chip Information:

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

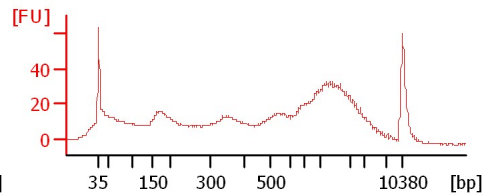
MNase Check A Expt 12/20/14



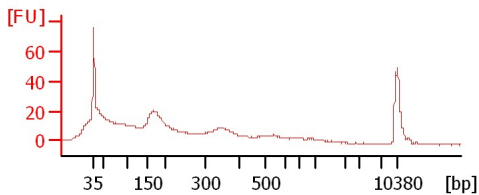
MNase Check B Expt 12/20/14



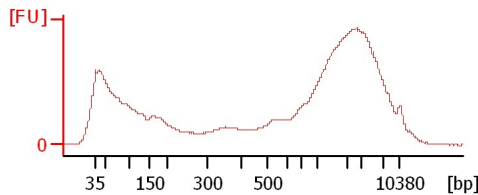
MNase Check Expt 1/20/15



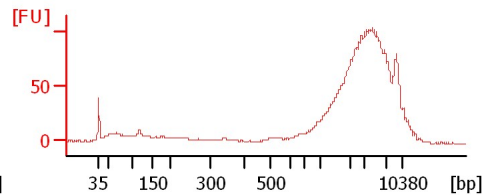
MNase Check Expt 1/31/15



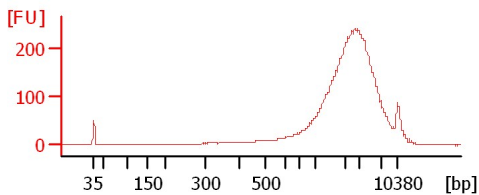
10_INPUT



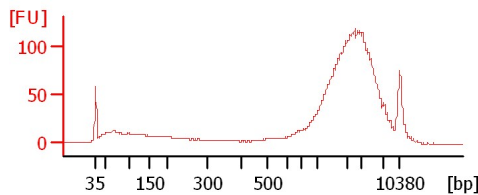
H3K27me1_IP



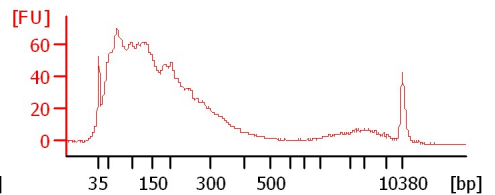
H3K27me3_IP



H3K4me3_IP



SH_HTR12_IP



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
Modified: 2/4/2015 2:35:06 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
MNase Check A Expt 12/20/14		<input type="checkbox"/>	✓			
MNase Check B Expt 12/20/14		<input type="checkbox"/>	✓			
MNase Check Expt 1/20/15		<input type="checkbox"/>	✓			
MNase Check Expt 1/31/15		<input type="checkbox"/>	✓			
10_INPUT		<input type="checkbox"/>	✓			
H3K27me1_IP		<input type="checkbox"/>	✓			
H3K27me3_IP		<input type="checkbox"/>	✓			
H3K4me3_IP		<input type="checkbox"/>	✓			
SH_HTR12_IP		<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:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
Modified: 2/4/2015 2:35:06 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

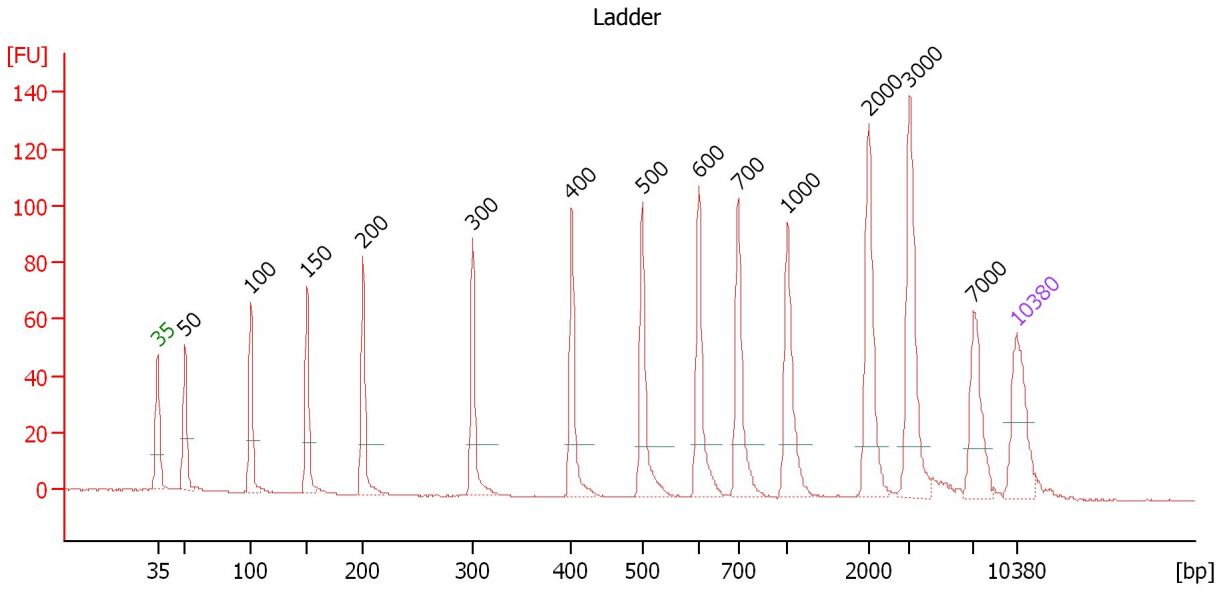
Ladder

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\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

Peak table for Ladder

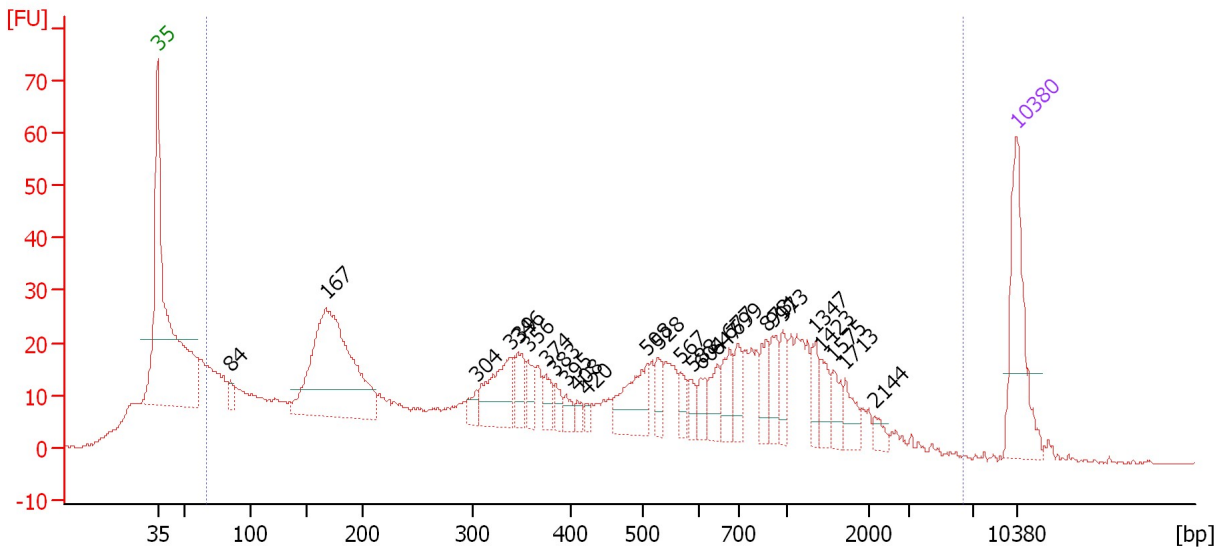
Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	50	150.00	4,545.5	Ladder Peak	45.25
3	100	150.00	2,272.7	Ladder Peak	50.61
4	150	150.00	1,515.2	Ladder Peak	55.19
5	200	150.00	1,136.4	Ladder Peak	59.73
6	300	150.00	757.6	Ladder Peak	68.67
7	400	150.00	568.2	Ladder Peak	76.74
8	500	150.00	454.5	Ladder Peak	82.52
9	600	150.00	378.8	Ladder Peak	87.10
10	700	150.00	324.7	Ladder Peak	90.31
11	1,000	150.00	227.3	Ladder Peak	94.30
12	2,000	150.00	113.6	Ladder Peak	100.94
13	3,000	150.00	75.8	Ladder Peak	104.29
14	7,000	150.00	32.5	Ladder Peak	109.47
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...

MNase Check A Expt 12/20/14



Overall Results for sample 1 : MNase Check A Expt 12/20/14

Number of peaks found: 27 Corr. Area 1: 1,159.2
 Noise: 0.2

Peak table for sample 1 : MNase Check A Expt 12/20/14

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	84	13.65	245.4		48.93
3	167	289.20	2,624.4		56.73
4	304	13.16	65.6		69.00
5	339	66.75	298.5		71.80
6	346	28.73	125.9		72.35
7	356	18.96	80.7		73.18
8	374	15.96	64.7		74.65
9	383	12.62	49.9		75.38
10	395	10.65	40.9		76.30
11	408	8.55	31.7		77.22
12	420	6.38	23.0		77.91
13	508	55.35	165.2		82.87
14	528	17.85	51.2		83.79
15	567	11.96	32.0		85.58
16	588	12.10	31.2		86.54
17	604	15.67	39.3		87.23
18	647	22.03	51.6		88.61
19	677	26.04	58.3		89.57
20	699	20.96	45.5		90.26
21	873	21.41	37.2		92.61
22	911	23.59	39.2		93.11
23	973	19.47	30.3		93.94
24	1,347	18.35	20.6		96.60
25	1,423	17.86	19.0		97.11

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad


Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...

... Peak table for sample 1 : MNase Check A Expt 12/20/14

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	1,575	14.82	14.3		98.12
27	1,713	16.77	14.8		99.04
28	2,144	7.66	5.4		101.43
29	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 1 : MNase Check A Expt 12/20/14

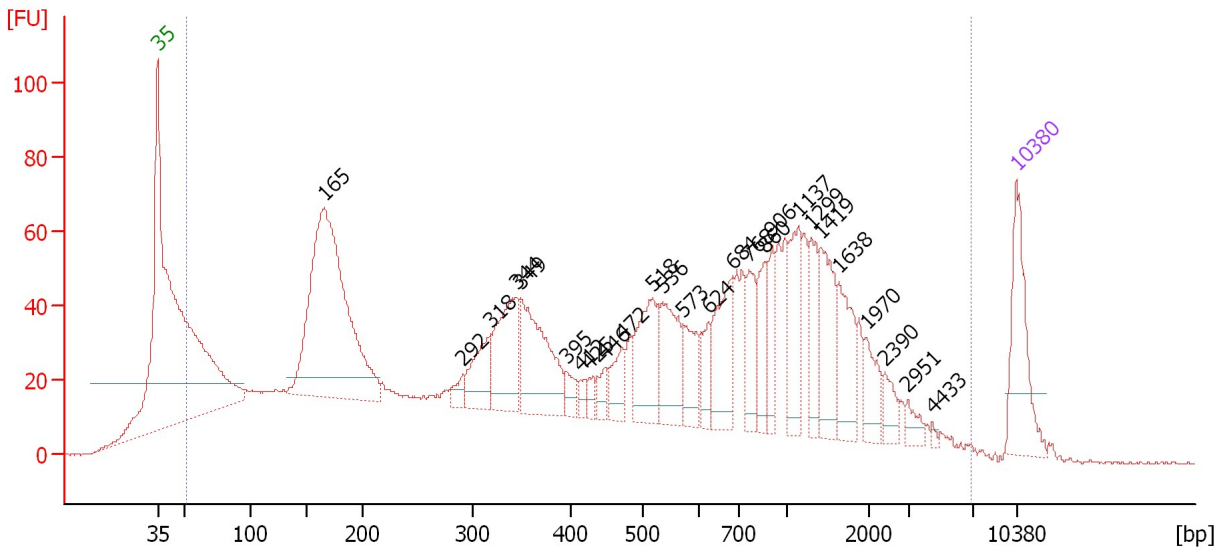
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
67	622	6,379	1,159.2	94	100.0	1,893.95	13,381.1	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...

MNase Check B Expt 12/20/14



Overall Results for sample 2 : MNase Check B Expt 12/20/14

Number of peaks found: 26 Corr. Area 1: 2,543.9
 Noise: 0.2

Peak table for sample 2 : MNase Check B Expt 12/20/14

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	165	511.57	4,693.6		56.57
3	292	16.76	87.1		67.93
4	318	60.05	286.5		70.09
5	344	109.89	484.3		72.20
6	349	135.62	588.8		72.62
7	395	17.10	65.6		76.33
8	412	10.41	38.2		77.46
9	425	11.04	39.3		78.21
10	446	17.13	58.2		79.38
11	472	33.39	107.2		80.89
12	518	90.46	264.7		83.33
13	536	78.84	222.8		84.17
14	573	46.10	121.9		85.86
15	624	28.48	69.1		87.88
16	684	79.39	175.8		89.81
17	768	45.80	90.3		91.22
18	860	43.03	75.8		92.44
19	906	33.39	55.8		93.05
20	1,137	62.81	83.7		95.21
21	1,299	41.71	48.6		96.29
22	1,419	69.91	74.6		97.08
23	1,638	54.32	50.2		98.54
24	1,970	30.05	23.1		100.75
25	2,390	16.73	10.6		102.25

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...

... Peak table for sample 2 : MNase Check B Expt 12/20/14

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,951	12.18	6.3		104.13
27	4,433	2.78	1.0		106.15
28	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 2 : MNase Check B Expt 12/20/14

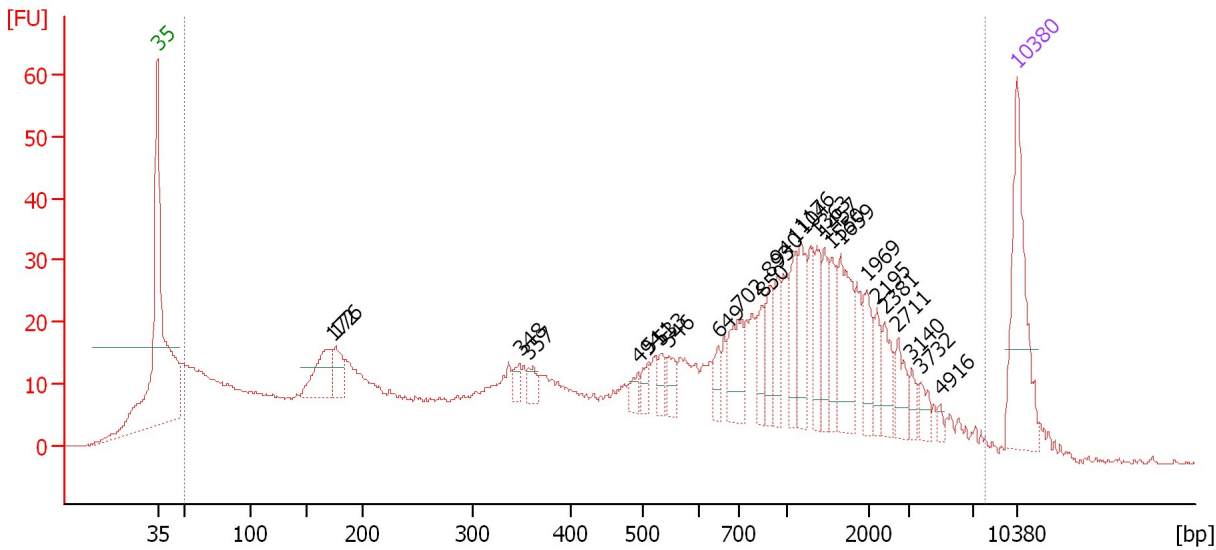
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
51	731	6,875	2,543.9	100	100.0	3,155.23	17,083.1	■

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...

MNase Check Expt 1/20/15



Overall Results for sample 3 : MNase Check Expt 1/20/15

Number of peaks found: 26 Corr. Area 1: 1,244.2
 Noise: 0.1

Peak table for sample 3 : MNase Check Expt 1/20/15

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	172	49.22	433.8		57.18
3	176	26.81	231.3		57.52
4	348	8.97	39.1		72.51
5	357	10.94	46.5		73.23
6	494	9.52	29.2		82.14
7	511	11.24	33.3		83.01
8	533	12.21	34.7		84.01
9	546	14.97	41.5		84.64
10	649	11.83	27.6		88.66
11	702	38.46	83.0		90.34
12	850	19.51	34.8		92.30
13	893	23.66	40.1		92.88
14	940	22.08	35.6		93.50
15	1,104	27.49	37.7		94.98
16	1,176	28.83	37.2		95.46
17	1,363	24.18	26.9		96.71
18	1,457	24.97	26.0		97.33
19	1,550	22.15	21.6		97.96
20	1,659	45.92	42.0		98.67
21	1,969	22.36	17.2		100.73
22	2,195	14.13	9.8		101.60
23	2,381	18.21	11.6		102.22
24	2,711	17.48	9.8		103.32
25	3,140	7.34	3.5		104.47

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad


Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...

... Peak table for sample 3 : MNase Check Expt 1/20/15

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	3,732	8.97	3.6		105.24
27	4,916	3.58	1.1		106.77
28	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 3 : MNase Check Expt 1/20/15

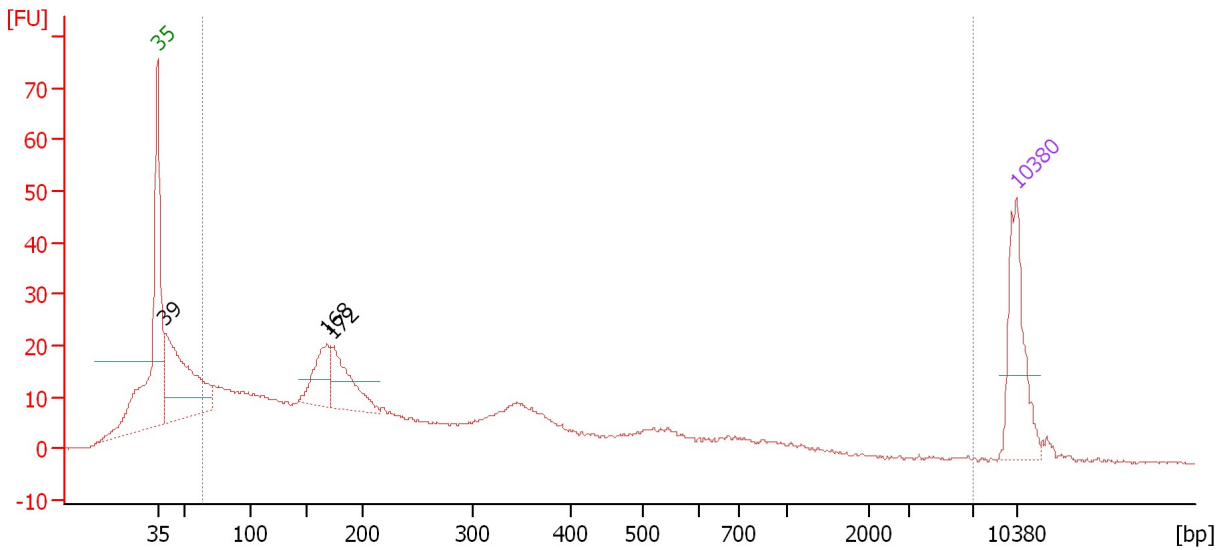
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
49	1,000	7,846	1,244.2	98	100.0	2,088.43	14,778.2	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...

MNase Check Expt 1/31/15



Overall Results for sample 4 : MNase Check Expt 1/31/15

Number of peaks found: 3 Corr. Area 1: 603.7
 Noise: 0.2

Peak table for sample 4 : MNase Check Expt 1/31/15

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	39	231.92	9,027.9		43.59
3	168	68.67	620.3		56.80
4	172	84.92	747.8		57.20
5	10,380	75.00	10.9	Upper Marker	113.00

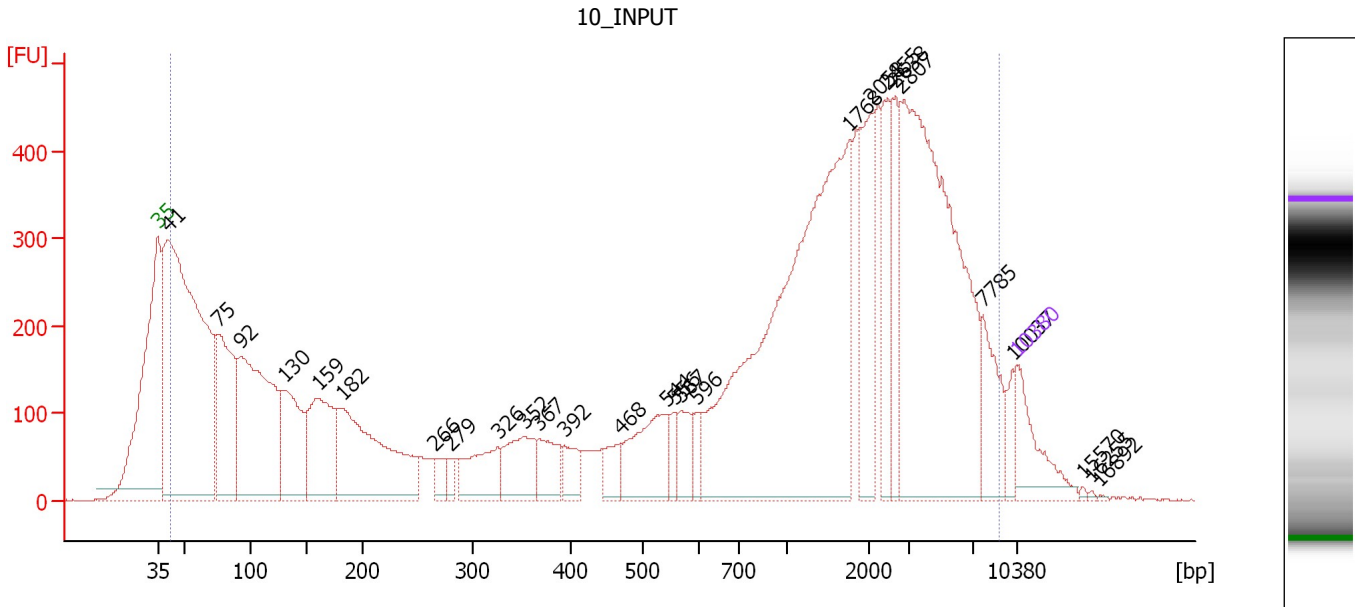
Region table for sample 4 : MNase Check Expt 1/31/15

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
64	316	7,036	603.7	82	100.0	1,199.25	11,556.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : 10_INPUT

Number of peaks found: 27 Corr. Area 1: 14,586.5
 Noise: 0.3

Peak table for sample 5 : 10_INPUT

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	41	1,365.24	50,693.8		43.87
3	75	343.47	6,911.4		47.96
4	92	548.85	8,994.3		49.80
5	130	229.66	2,683.8		53.33
6	159	240.28	2,285.3		56.04
7	182	388.96	3,240.9		58.08
8	266	30.03	170.9		65.65
9	279	22.95	124.5		66.83
10	326	103.74	482.2		70.76
11	352	102.19	439.9		72.86
12	367	71.83	296.4		74.09
13	392	44.52	171.8		76.13
14	468	38.72	125.3		80.68
15	544	136.98	381.7		84.52
16	556	27.17	74.0		85.08
17	567	47.33	126.4		85.59
18	596	25.64	65.2		86.92
19	1,768	821.42	704.1		99.40
20	2,058	143.48	105.6		101.14
21	2,455	111.73	69.0		102.47
22	2,638	80.65	46.3		103.08
23	2,807	617.49	333.4		103.64
24	7,785	81.72	15.9		110.29
25	10,037	26.26	4.0		112.64

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad


Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...

... Peak table for sample 5 : 10 INPUT

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	15,570	0.00	0.0		118.42
28	16,255	0.00	0.0		119.14
29	16,892	0.00	0.0		119.80

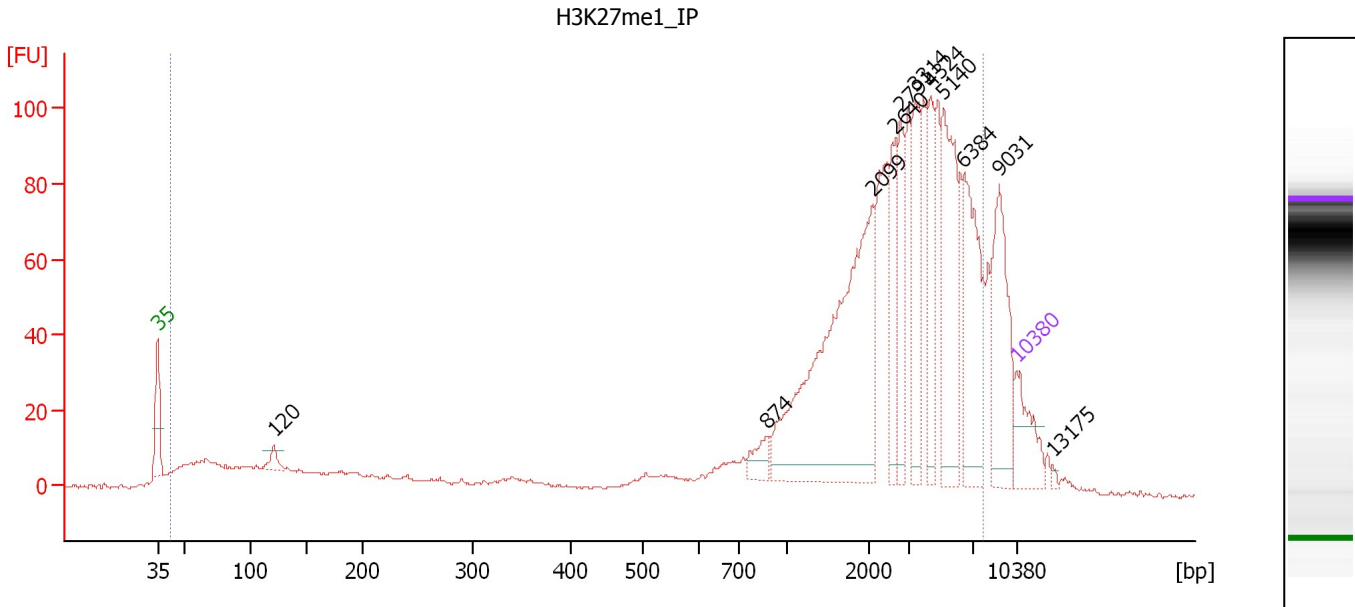
Region table for sample 5 : 10 INPUT

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/ μ l]	Molarity [pmol/l]	Color
42	1,861	9,046	14,586.5	95	100.0	5,584.80	53,777.5	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : H3K27me1_IP

Number of peaks found: 11 Corr. Area 1: 1,467.9
 Noise: 0.3

Peak table for sample 6 : H3K27me1_IP

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	120	28.87	363.9		52.46
3	874	35.58	61.7		92.62
4	2,099	580.23	418.9		101.27
5	2,640	101.61	58.3		103.09
6	2,791	106.17	57.6		103.59
7	3,314	130.45	59.6		104.70
8	4,324	114.09	40.0		106.01
9	5,140	236.47	69.7		107.06
10	6,384	206.95	49.1		108.67
11	9,031	180.61	30.3		111.59
12	10,380	75.00	10.9	Upper Marker	113.00
13	13,175	0.00	0.0		115.92

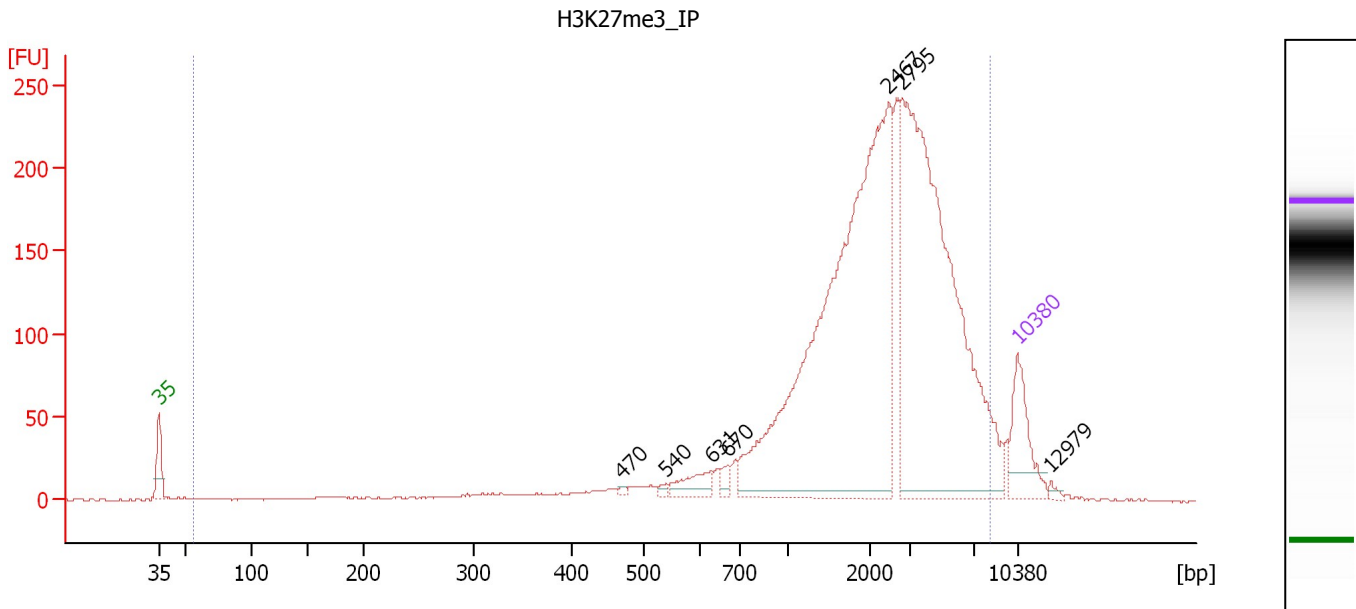
Region table for sample 6 : H3K27me1_IP

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
42	3,022	7,793	1,467.9	89	67.4	3,034.34	12,422.3	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : H3K27me3 IP

Number of peaks found: 7 Corr. Area 1: 2,930.4
 Noise: 0.3

Peak table for sample 7 : H3K27me3 IP

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	470	3.74	12.0		80.79
3	540	5.34	15.0		84.34
4	631	37.62	90.3		88.10
5	670	14.02	31.7		89.35
6	2,467	910.44	559.1		102.51
7	2,795	747.82	405.4		103.60
8	10,380	75.00	10.9	Upper Marker	113.00
9	12,979	0.00	0.0		115.71

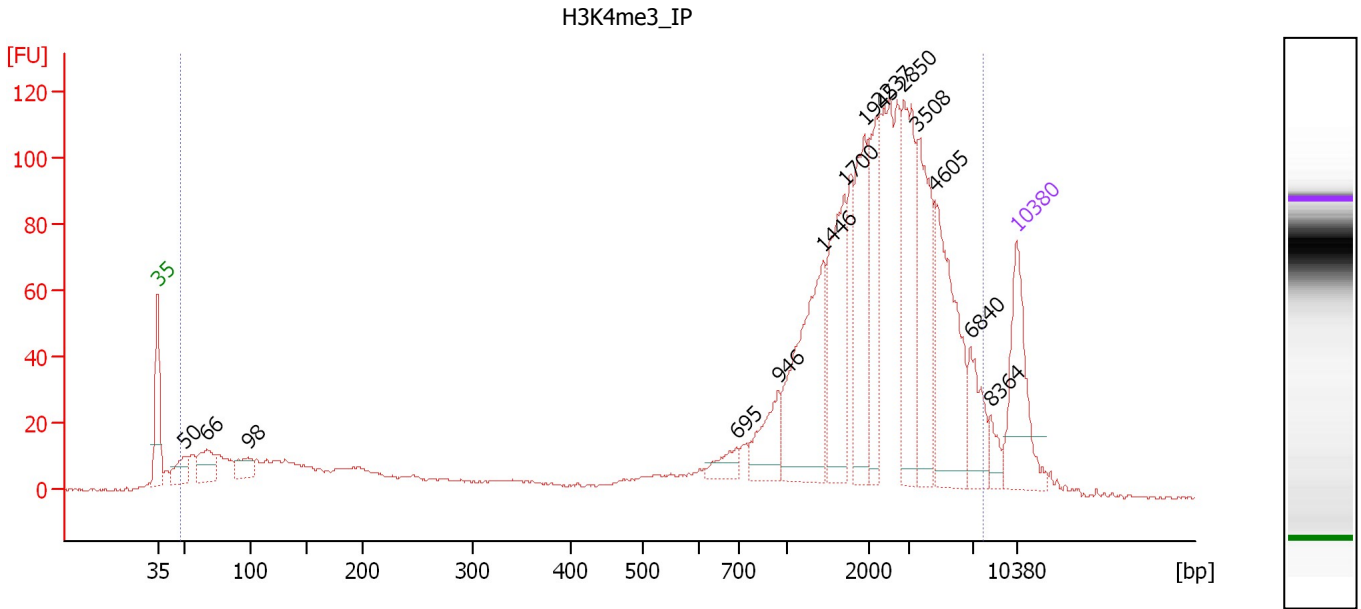
Region table for sample 7 : H3K27me3 IP

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
55	2,778	8,233	2,930.4	97	63.0	2,040.83	2,181.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : H3K4me3 IP

Number of peaks found: 14 Corr. Area 1: 1,810.8
 Noise: 0.4

Peak table for sample 8 : H3K4me3 IP

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	33.73	1,027.2		45.21
3	66	46.30	1,057.3		47.00
4	98	27.24	422.3		50.37
5	695	20.39	44.4		90.16
6	946	47.35	75.8		93.58
7	1,446	148.86	156.0		97.26
8	1,700	105.96	94.5		98.95
9	1,945	92.07	71.7		100.58
10	2,237	79.87	54.1		101.74
11	2,850	108.53	57.7		103.79
12	3,508	103.70	44.8		104.95
13	4,605	125.08	41.2		106.37
14	6,840	42.15	9.3		109.26
15	8,364	14.23	2.6		110.89
16	10,380	75.00	10.9	Upper Marker	113.00

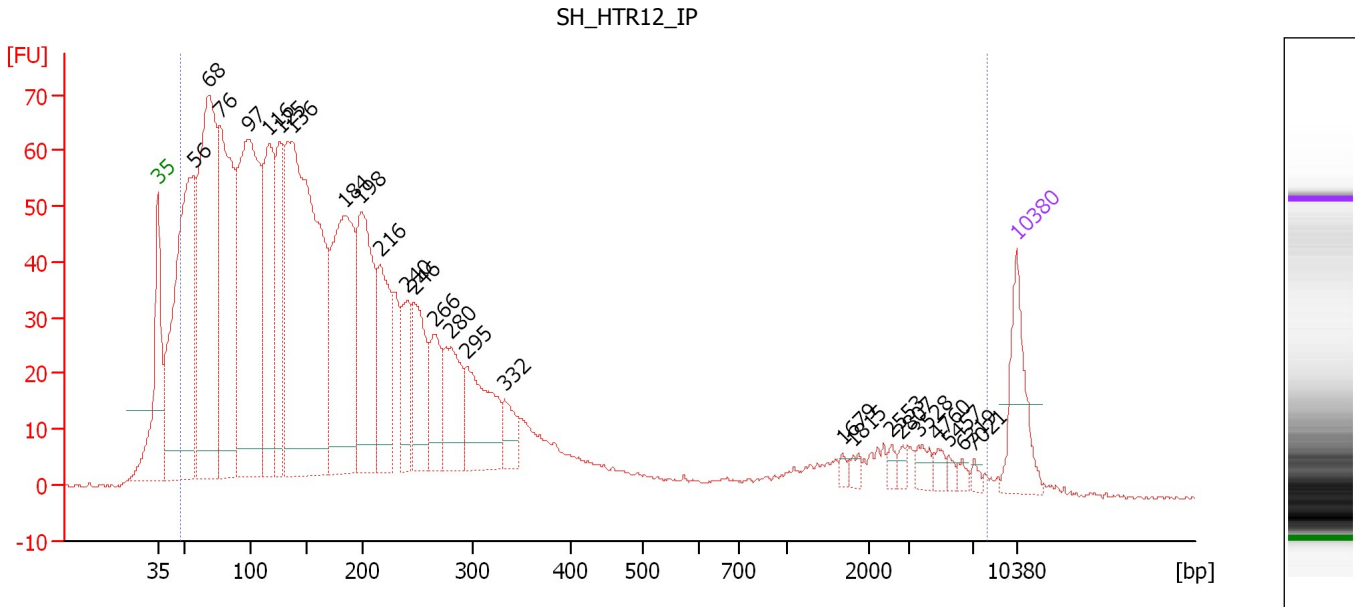
Region table for sample 8 : H3K4me3 IP

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
47	2,376	7,756	1,810.8	96	72.9	1,732.50	8,019.3	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : SH_HTR12_IP

Number of peaks found: 25 Corr. Area 1: 2,375.8
 Noise: 0.2

Peak table for sample 9 : SH_HTR12_IP

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	56	679.82	18,291.5		45.92
3	68	689.65	15,320.7		47.20
4	76	518.30	10,315.1		48.05
5	97	709.11	11,026.6		50.33
6	116	293.73	3,832.2		52.09
7	125	222.40	2,687.0		52.94
8	136	971.99	10,795.1		53.95
9	184	433.70	3,565.4		58.31
10	198	312.23	2,385.0		59.58
11	216	175.13	1,227.5		61.18
12	240	87.15	550.3		63.30
13	246	143.62	882.8		63.89
14	266	94.72	540.5		65.59
15	280	123.39	666.8		66.92
16	295	153.07	787.1		68.19
17	332	42.65	194.5		71.28
18	1,679	6.12	5.5		98.81
19	1,815	8.83	7.4		99.71
20	2,553	8.28	4.9		102.79
21	2,807	7.79	4.2		103.65
22	3,528	15.62	6.7		104.97
23	4,760	9.89	3.1		106.57
24	5,457	7.54	2.1		107.47
25	6,319	6.20	1.5		108.59

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad


Created: 2/4/2015 1:46:44 PM
 Modified: 2/4/2015 2:35:06 PM

Electropherogram Summary Continued ...

... Peak table for sample 9 : SH_HTR12_IP

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	7,021	5.96	1.3		109.49
27	10,380	75.00	10.9	Upper Marker	113.00

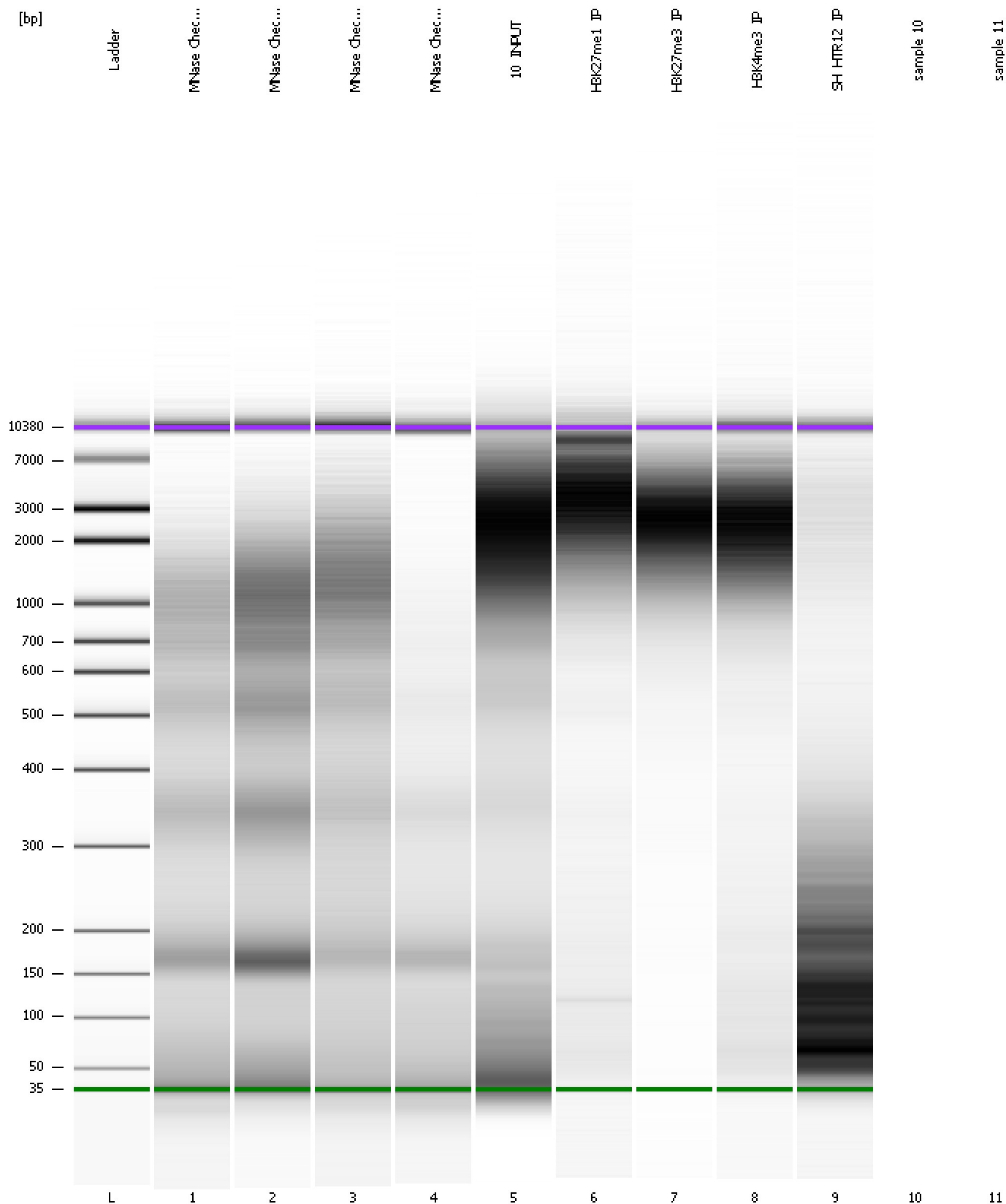
Region table for sample 9 : SH_HTR12_IP

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/ μ l]	Molarity [pmol/l]	Color
47	409	8,033	2,375.8	96	100.0	6,362.91	81,074.2	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
Modified: 2/4/2015 2:35:06 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad

Created: 2/4/2015 1:46:44 PM
Modified: 2/4/2015 2:35:06 PM

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 Created: 2/4/2015 1:46:44 PM
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad Modified: 2/4/2015 2:35:06 PM

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		2/4/2015 2:22:18 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-02-04\2015-02-04_001.xad)		Instrument	Run		2/4/2015 1:46:49 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		2/4/2015 1:46:49 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		2/4/2015 1:46:49 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		2/4/2015 1:46:49 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		2/4/2015 1:46:49 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		2/4/2015 1:46:49 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		2/4/2015 1:46:49 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1