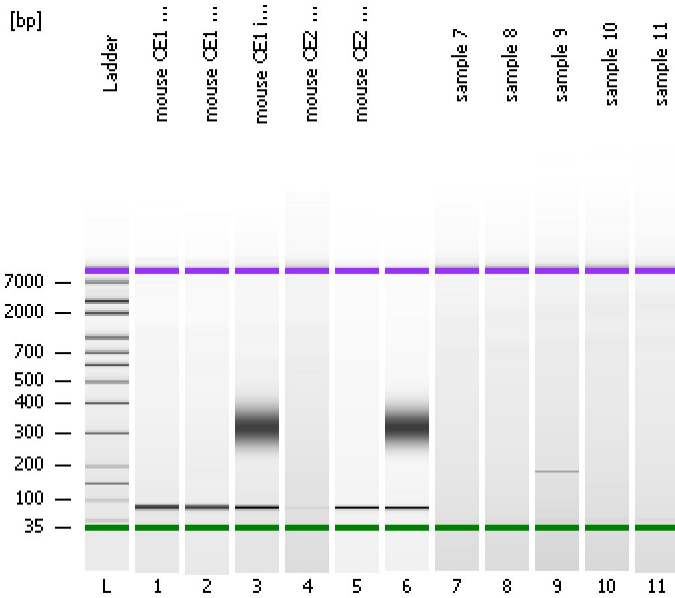


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
Modified: 2/18/2016 4:59:31 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

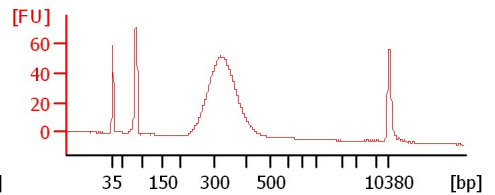
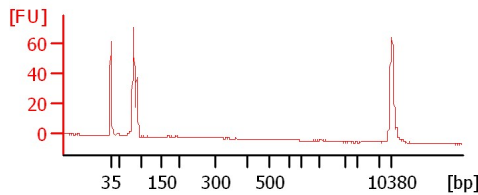
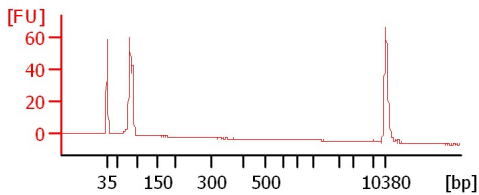
Chip Information:

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

mouse OE1 Mecp2 ChIP

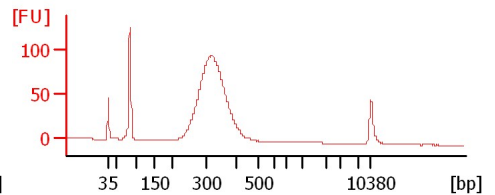
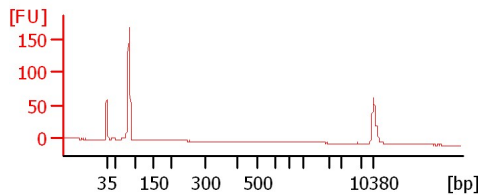
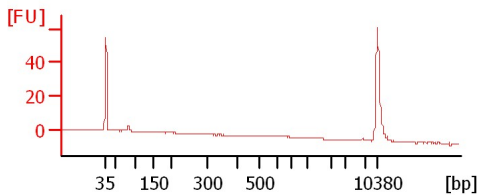
mouse OE1 IgG ChIP

mouse OE1 input



mouse OE2 Mecp2 ChIP

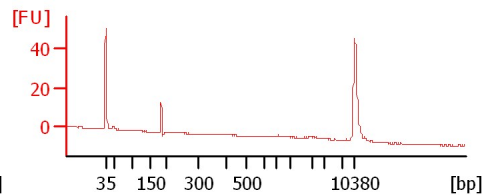
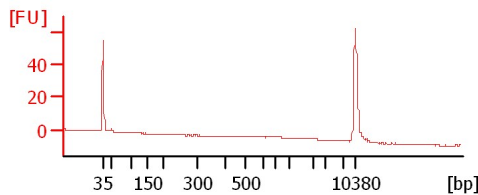
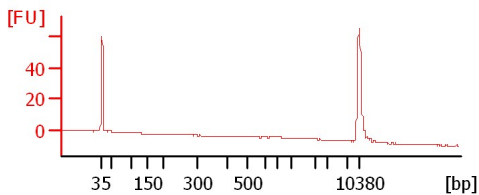
mouse OE2 IgG ChIP



sample 7

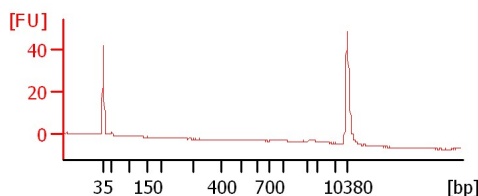
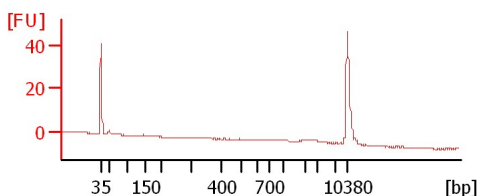
sample 8

sample 9



sample 10

sample 11



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
mouse OE1 Mecp2 ChIP		<input type="checkbox"/>	✓			
mouse OE1 IgG ChIP		<input type="checkbox"/>	✓			
mouse OE1 input		<input type="checkbox"/>	✓			
mouse OE2 Mecp2 ChIP		<input type="checkbox"/>	✓			
mouse OE2 IgG ChIP		<input type="checkbox"/>	✓			
sample 7		<input type="checkbox"/>	✓			
sample 8		<input type="checkbox"/>	✓			
sample 9		<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\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 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

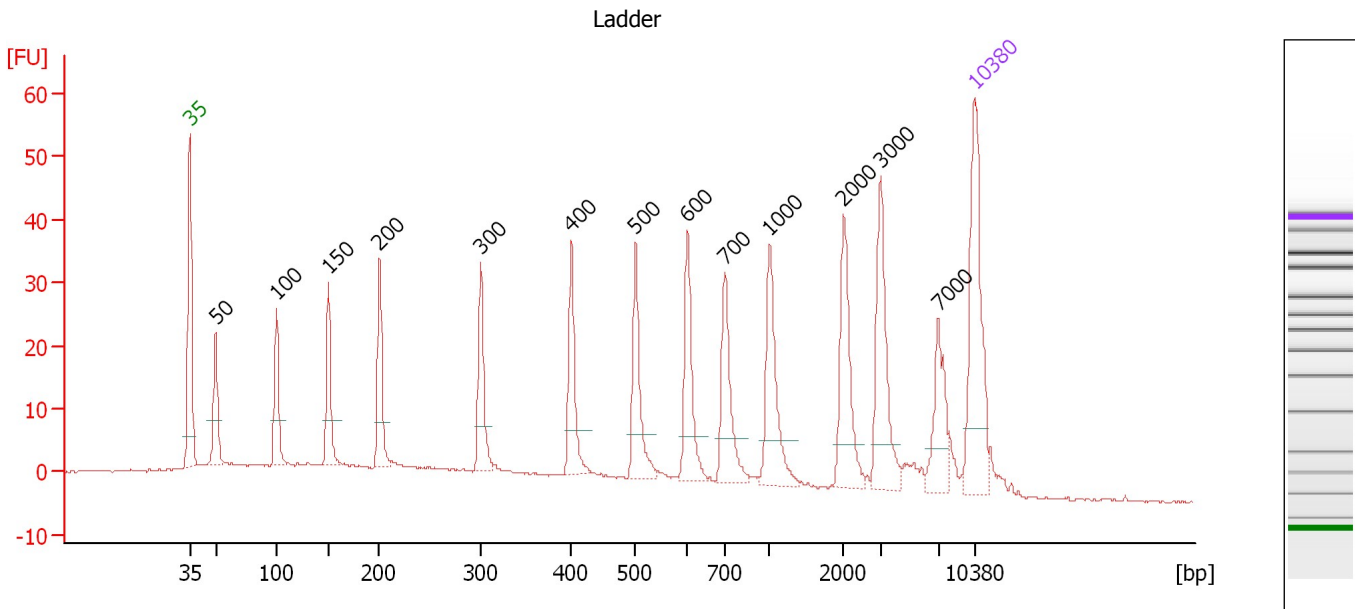
Ladder

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\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

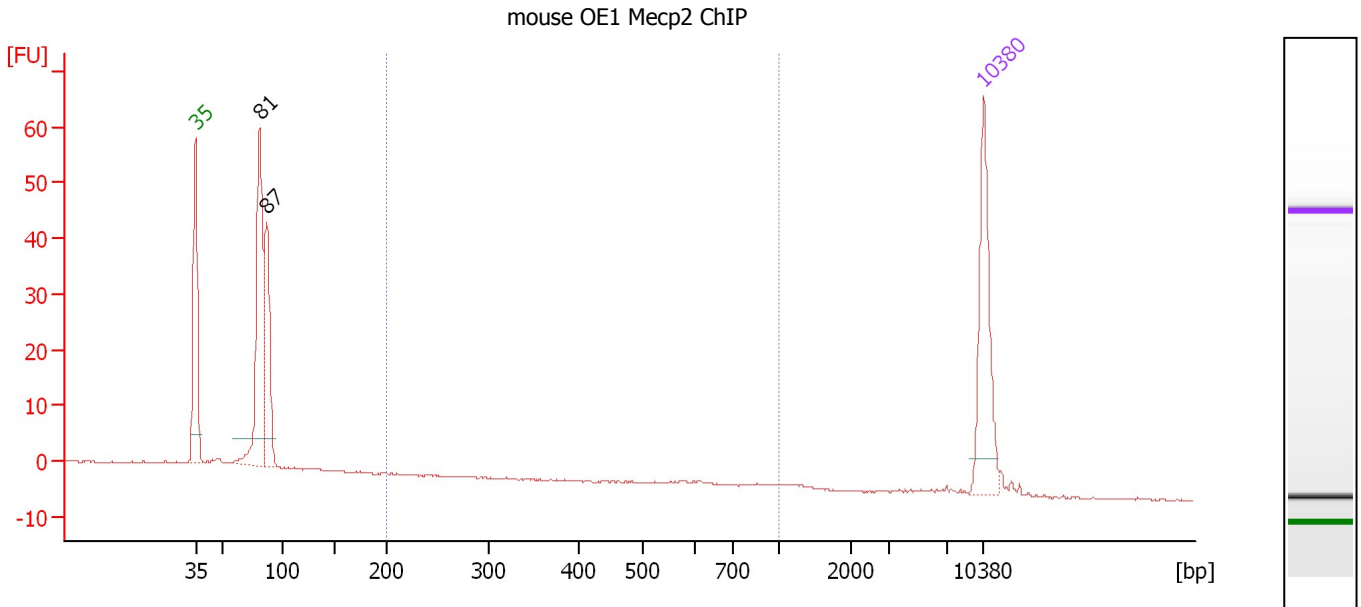
Peak table for Ladder

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	45.33
3	100	150.00	2,272.7	Ladder Peak	50.75
4	150	150.00	1,515.2	Ladder Peak	55.36
5	200	150.00	1,136.4	Ladder Peak	59.93
6	300	150.00	757.6	Ladder Peak	68.96
7	400	150.00	568.2	Ladder Peak	77.05
8	500	150.00	454.5	Ladder Peak	82.76
9	600	150.00	378.8	Ladder Peak	87.42
10	700	150.00	324.7	Ladder Peak	90.74
11	1,000	150.00	227.3	Ladder Peak	94.74
12	2,000	150.00	113.6	Ladder Peak	101.30
13	3,000	150.00	75.8	Ladder Peak	104.63
14	7,000	150.00	32.5	Ladder Peak	109.77
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : mouse OE1 Mecp2 ChIP

Number of peaks found: 2 Corr. Area 1: 1.2
 Noise: 0.2

Peak table for sample 1 : mouse OE1 Mecp2 ChIP

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	204.42	3,827.3		48.68
3	87	114.46	2,002.5		49.30
4	10,380	75.00	10.9	Upper Marker	113.00

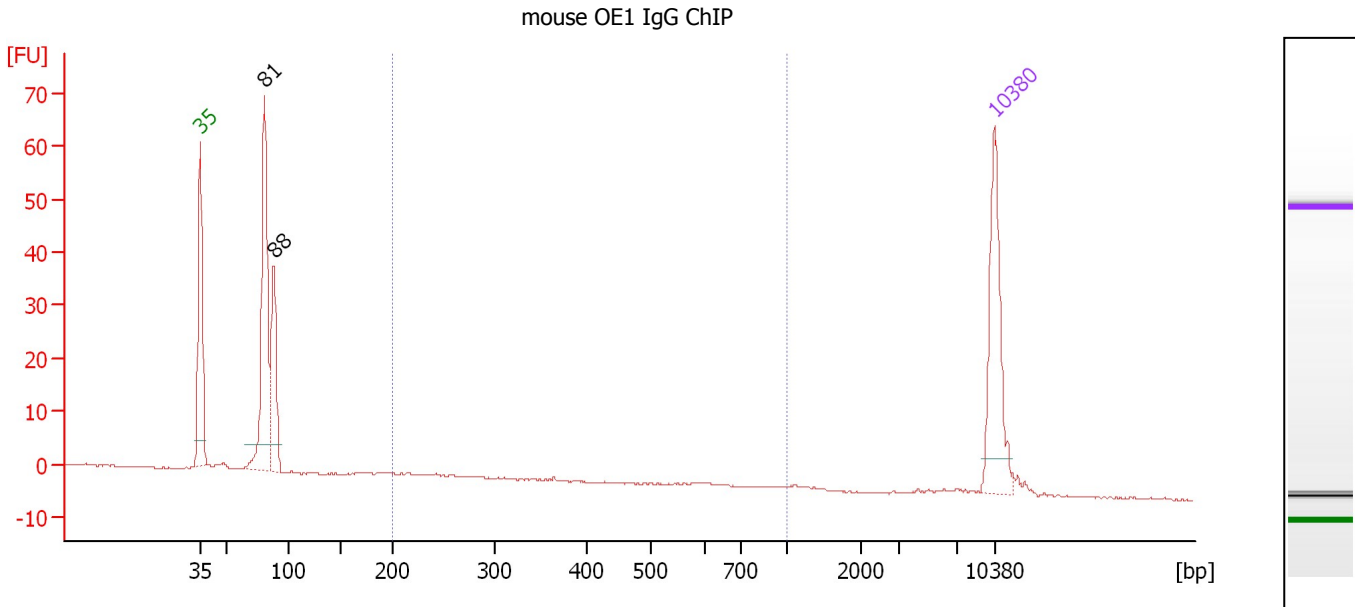
Region table for sample 1 : mouse OE1 Mecp2 ChIP

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	705	1.2	3.8	1.69	1	21.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : mouse OE1 IgG ChIP

Number of peaks found: 2 Corr. Area 1: 2.9
 Noise: 0.2

Peak table for sample 2 : mouse OE1 IgG ChIP

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	206.58	3,886.3		48.64
3	88	93.34	1,600.9		49.49
4	10,380	75.00	10.9	Upper Marker	113.00

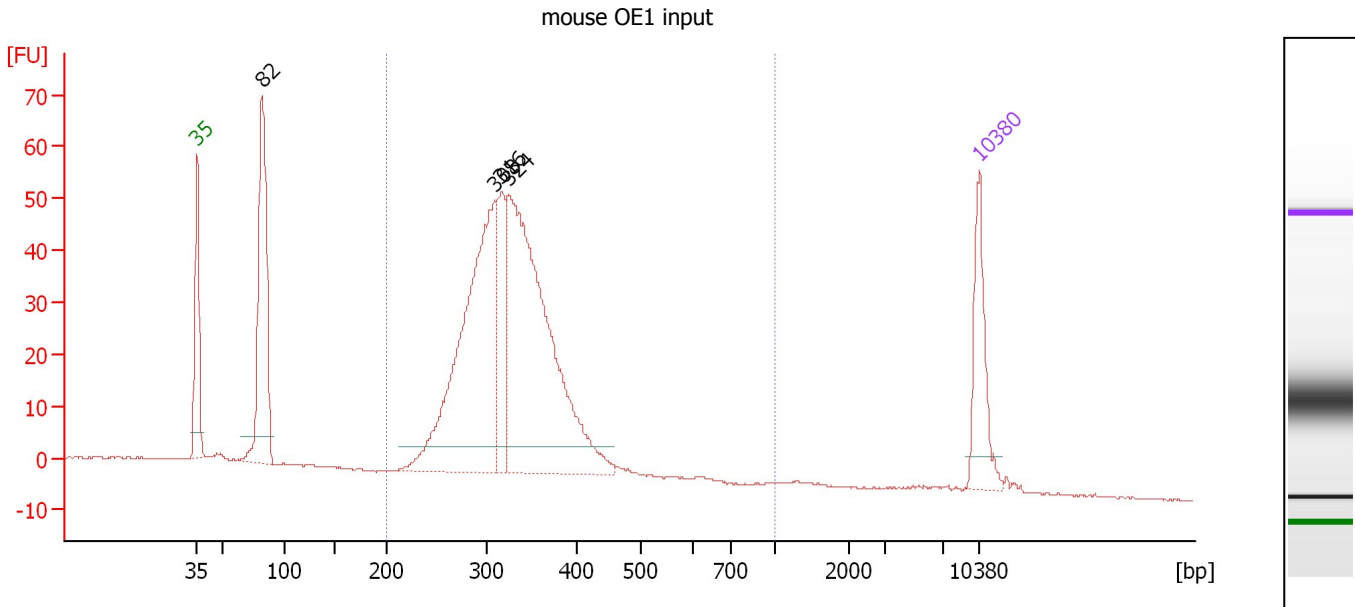
Region table for sample 2 : mouse OE1 IgG ChIP

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	388	2.9	27.3	5.04	2	53.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : mouse OE1 input

Number of peaks found: 4 Corr. Area 1: 673.0
 Noise: 0.1

Peak table for sample 3 : mouse OE1 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	82	318.78	5,907.5		48.77
3	308	507.43	2,494.8		69.63
4	316	110.45	529.8		70.25
5	324	609.52	2,854.5		70.87
6	10,380	75.00	10.9	Upper Marker	113.00

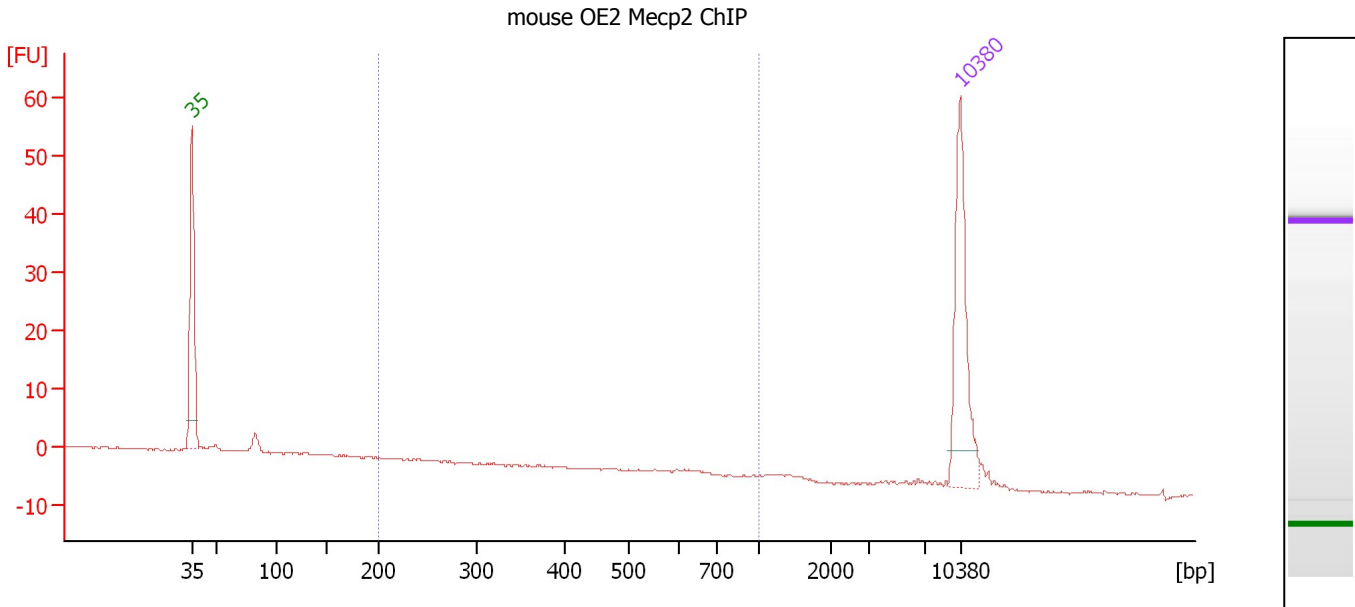
Region table for sample 3 : mouse OE1 input

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	329	673.0	5,964.0	1,259.30	80	16.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : mouse OE2 Mecp2 ChIP

Number of peaks found: 0 Corr. Area 1: 12.3
 Noise: 0.2

Peak table for sample 4 : mouse OE2 Mecp2 ChIP

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	10,380	75.00	10.9	Upper Marker	113.00

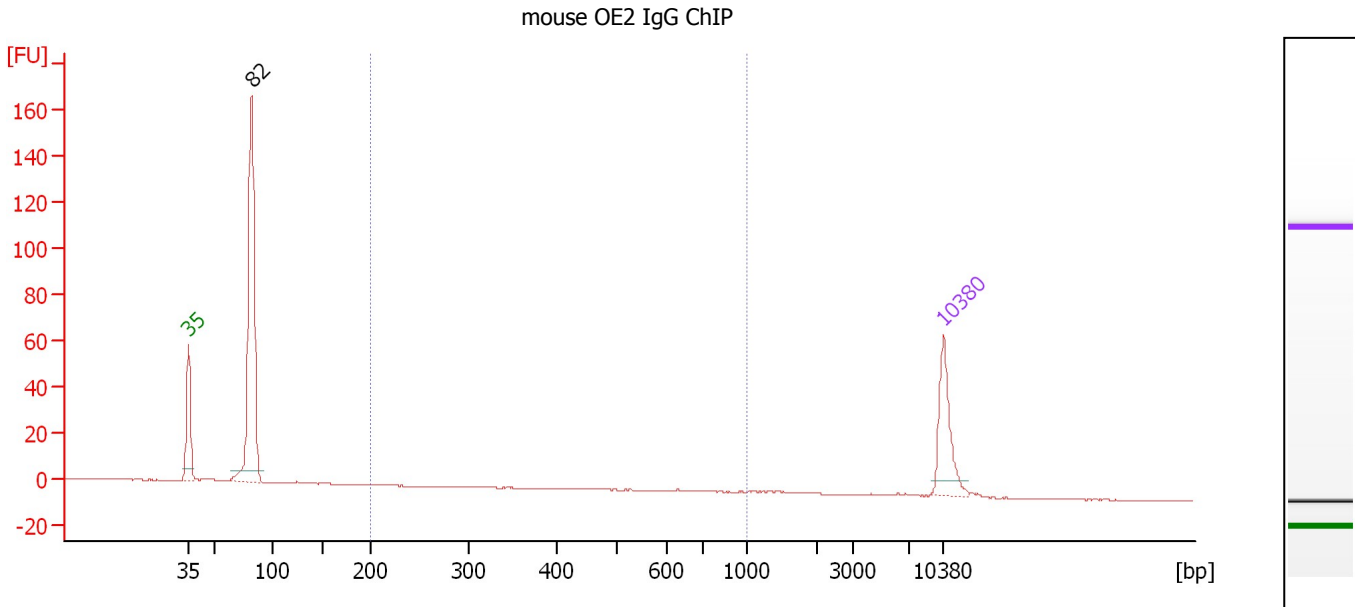
Region table for sample 4 : mouse OE2 Mecp2 ChIP

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	438	12.3	87.7	19.75	26	40.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : mouse OE2 IgG ChIP

Number of peaks found: 1 Corr. Area 1: 0.9
 Noise: 0.2

Peak table for sample 5 : mouse OE2 IgG ChIP

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	82	494.05	9,094.4		48.83
3	10,380	75.00	10.9	Upper Marker	113.00

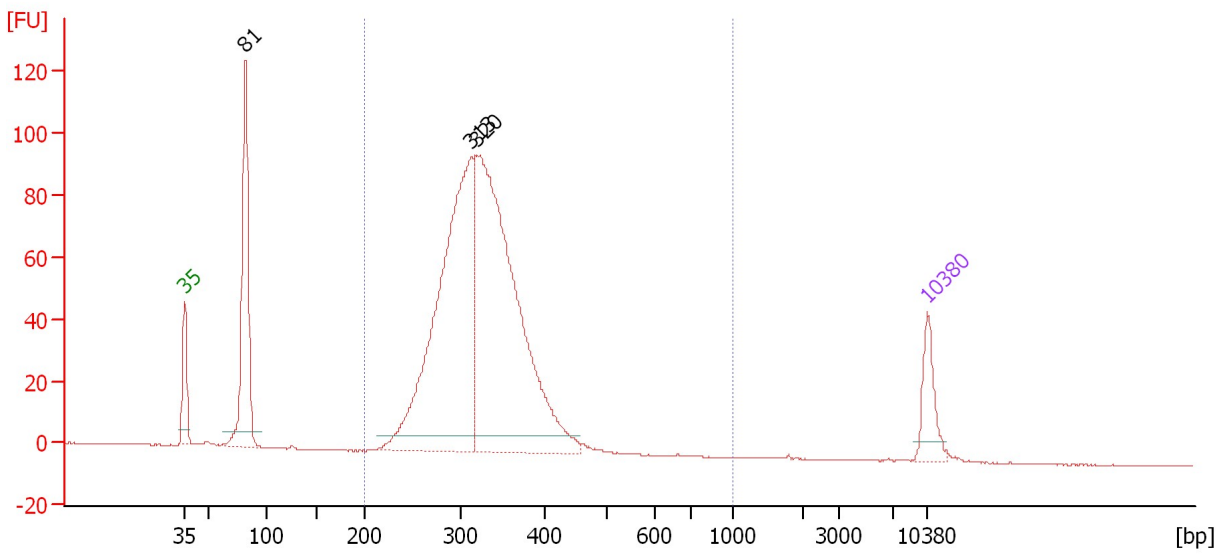
Region table for sample 5 : mouse OE2 IgG ChIP

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	657	0.9	2.8	1.21	0	10.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 :

Number of peaks found: 3 Corr. Area 1: 1,111.8
 Noise: 0.1

Peak table for sample 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	81	524.11	9,762.3		48.73
3	313	1,136.92	5,499.3		70.04
4	320	1,348.15	6,381.7		70.59
5	10,380	75.00	10.9	Upper Marker	113.00

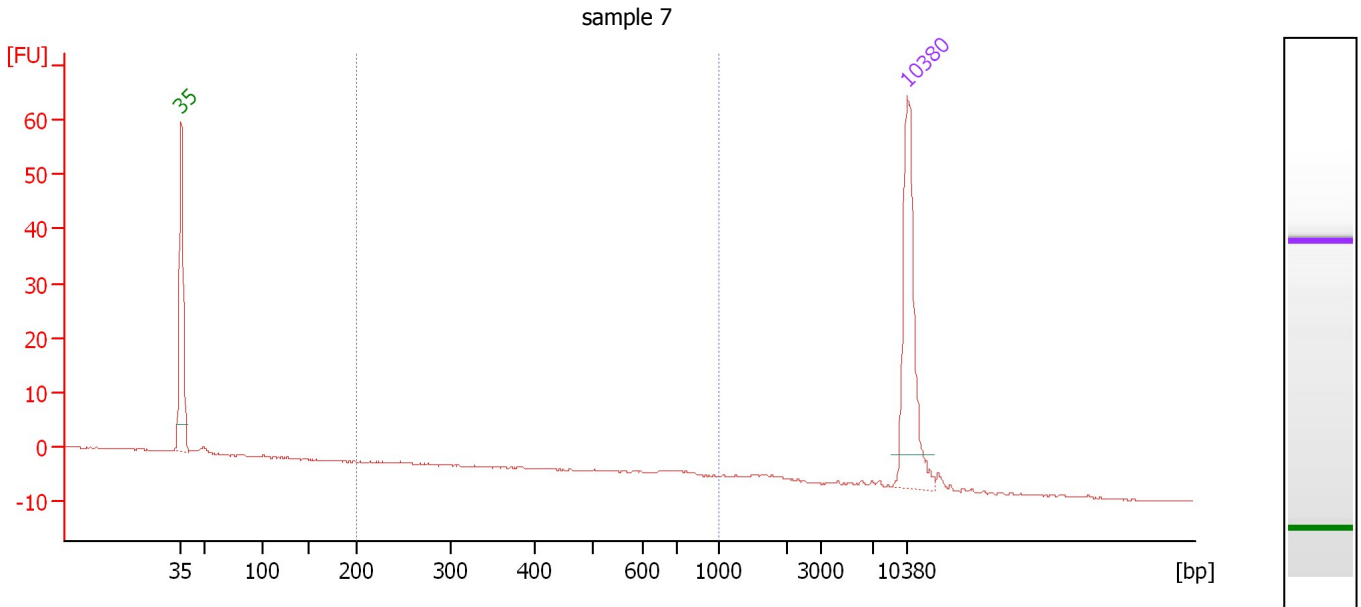
Region table for sample 6 :

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	325	1,111.8	11,744.2	2,467.63	86	13.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : sample 7

Number of peaks found: 0 Corr. Area 1: 7.6
 Noise: 0.1

Peak table for sample 7 : sample 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	10,380	75.00	10.9	Upper Marker	113.00

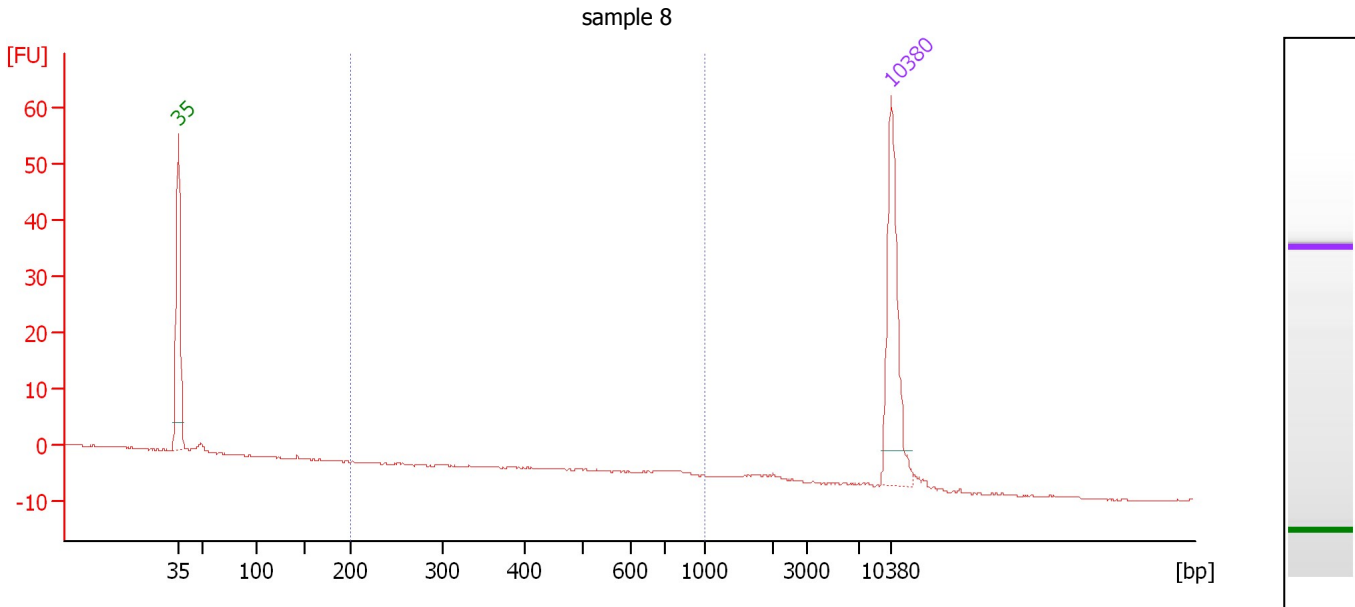
Region table for sample 7 : sample 7

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	649	7.6	23.7	9.45	40	22.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : sample 8

Number of peaks found: 0 Corr. Area 1: 2.9
 Noise: 0.1

Peak table for sample 8 : sample 8

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	10,380	75.00	10.9	Upper Marker	113.00

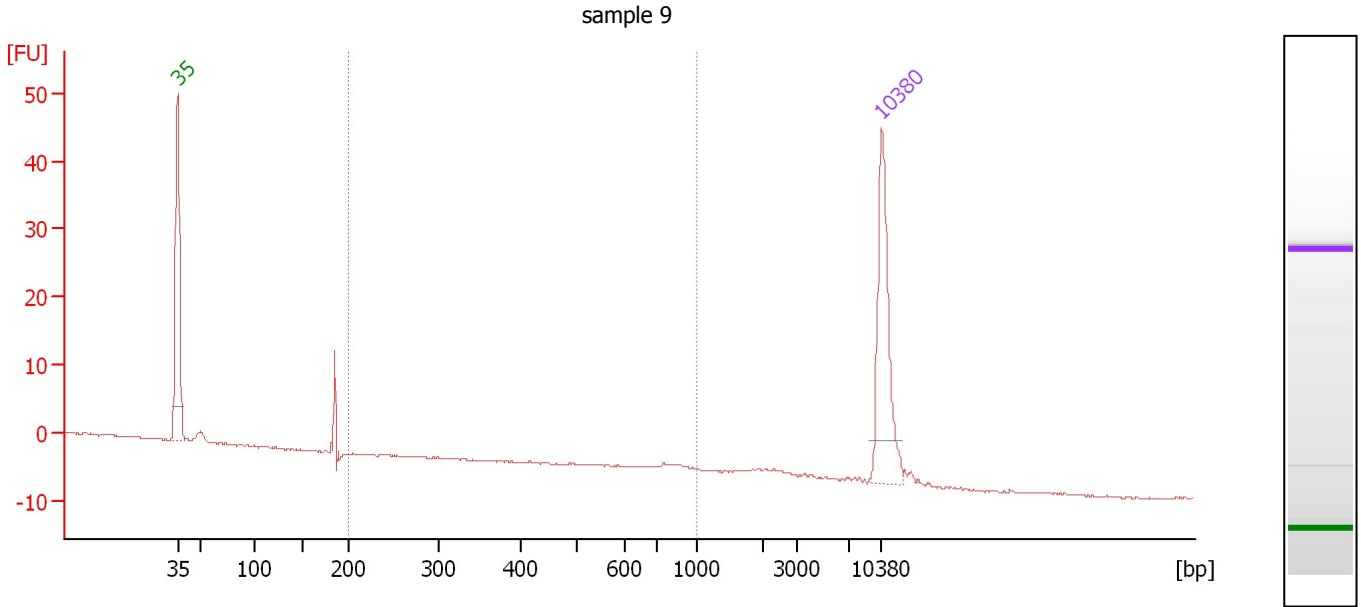
Region table for sample 8 : sample 8

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	725	2.9	8.7	4.06	30	14.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : sample 9

Number of peaks found: 0 Corr. Area 1: 1.4
 Noise: 0.1

Peak table for sample 9 : sample 9

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	10,380	75.00	10.9	Upper Marker	113.00

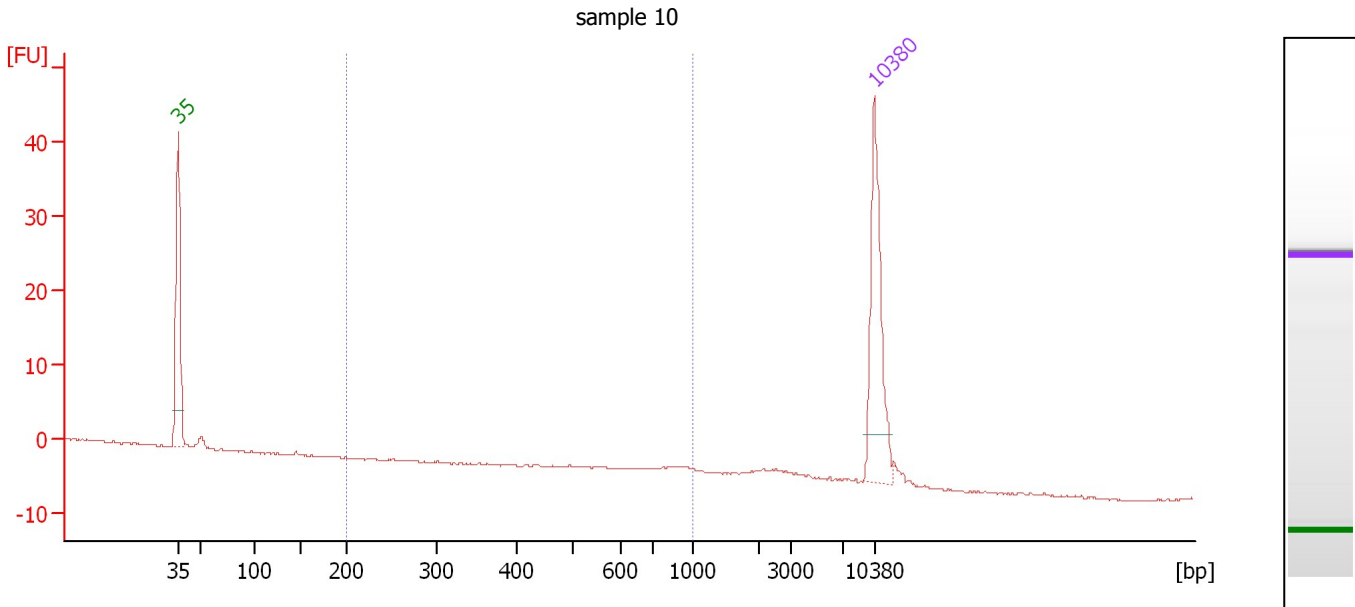
Region table for sample 9 : sample 9

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	827	1.4	4.4	2.42	12	9.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : sample 10

Number of peaks found: 0 Corr. Area 1: 3.2
 Noise: 0.1

Peak table for sample 10 : sample 10

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	10,380	75.00	10.9	Upper Marker	113.00

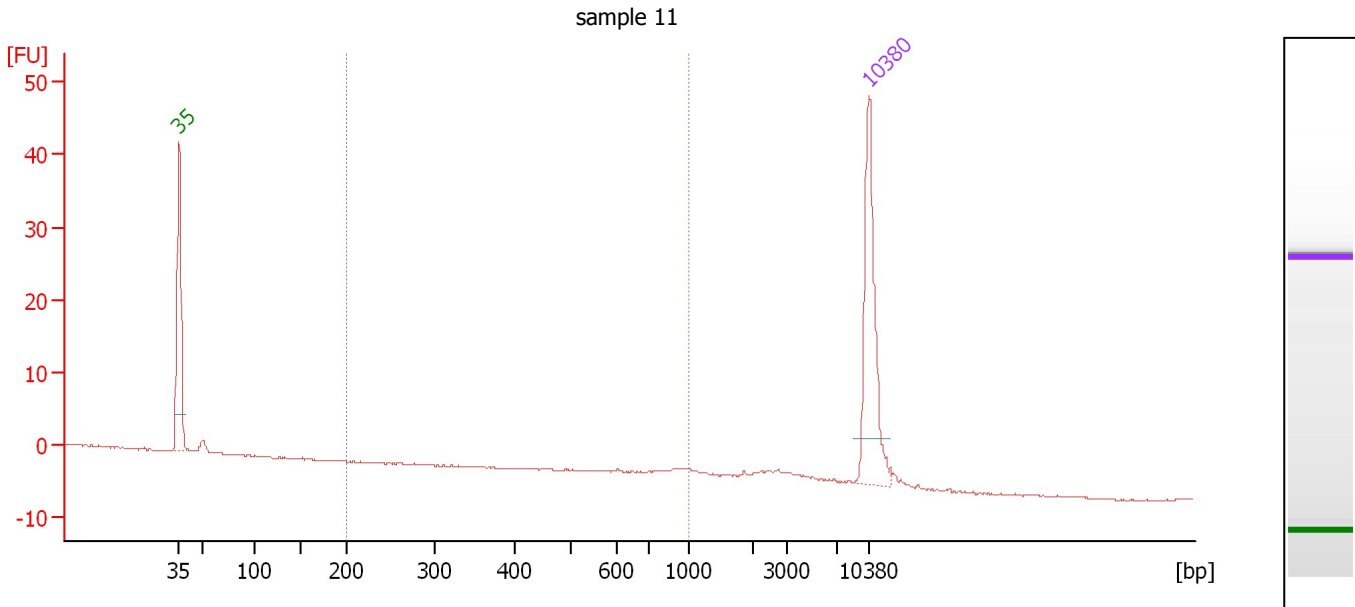
Region table for sample 10 : sample 10

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	811	3.2	10.9	5.67	23	15.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : sample 11

Number of peaks found: 0 Corr. Area 1: 2.5
 Noise: 0.1

Peak table for sample 11 : sample 11

Pea k	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	10,380	75.00	10.9	Upper Marker	113.00

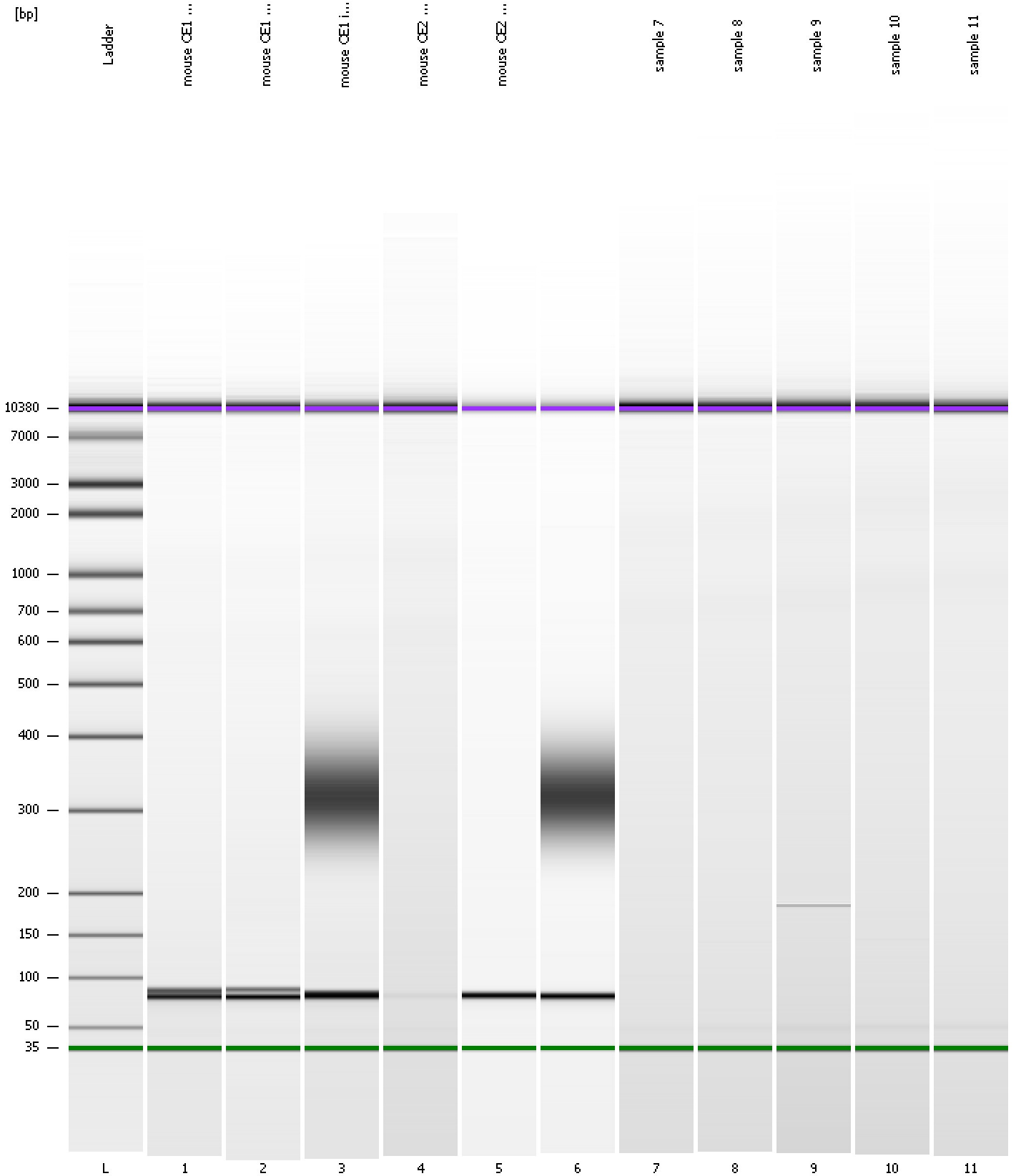
Region table for sample 11 : sample 11

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	851	2.5	7.6	4.22	18	12.8

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
Modified: 2/18/2016 4:59:31 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad

Created: 2/18/2016 4:13:38 PM
 Modified: 2/18/2016 4:59:31 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		2/18/2016 4:54:54 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2016-02-18\2016-02-18_002.xad)		Instrument	Run		2/18/2016 4:13:43 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		2/18/2016 4:13:43 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		2/18/2016 4:13:43 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		2/18/2016 4:13:43 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		2/18/2016 4:13:43 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		2/18/2016 4:13:43 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		2/18/2016 4:13:43 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1