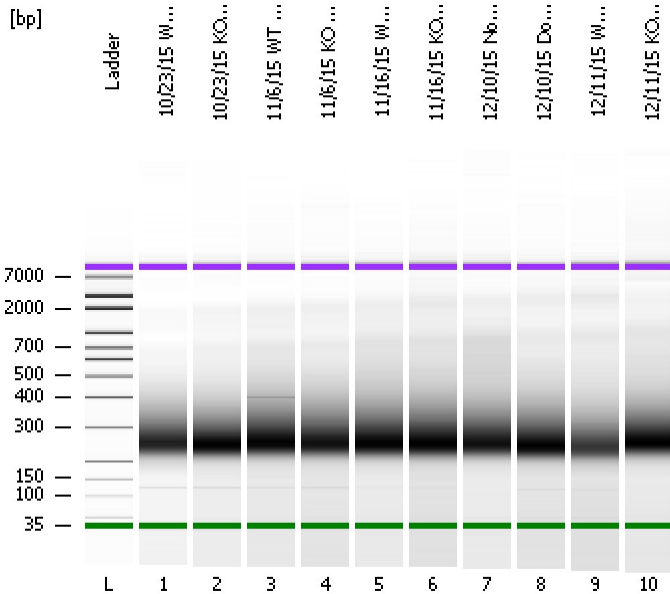


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
Modified: 5/13/2016 8:48:35 AM

Electrophoresis File Run Summary



Instrument Information:

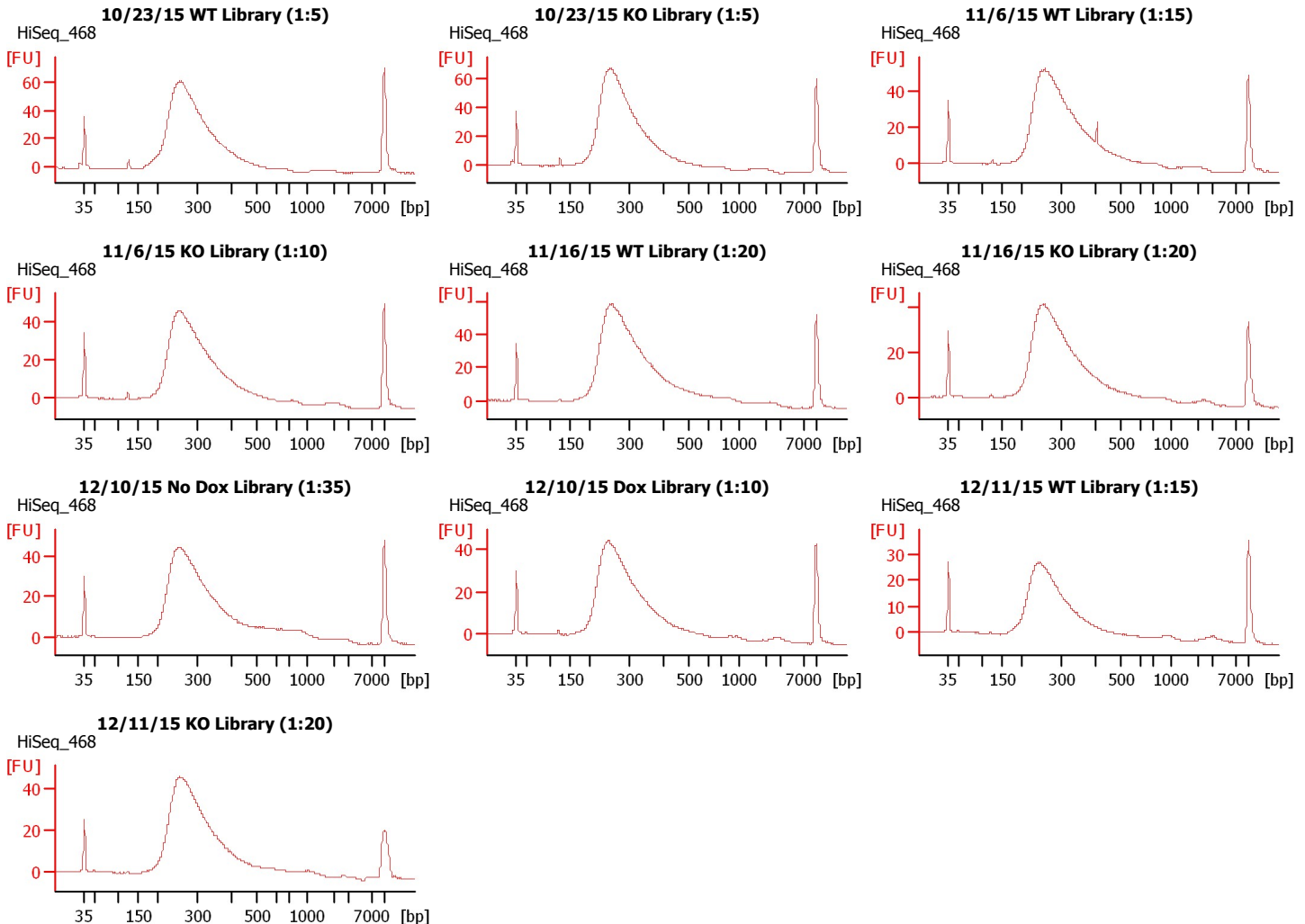
Instrument Name: DE13701086      Firmware: C.01.069  
Serial#: DE13701086      Type: G2938B

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:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
Modified: 5/13/2016 8:48:35 AM

**Electrophoresis File Run Summary (Chip Summary)**

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
10/23/15 WT Library (1:5)	HiSeq_468	<input type="checkbox"/>	✓			
10/23/15 KO Library (1:5)	HiSeq_468	<input type="checkbox"/>	✓			
11/6/15 WT Library (1:15)	HiSeq_468	<input type="checkbox"/>	✓			
11/6/15 KO Library (1:10)	HiSeq_468	<input type="checkbox"/>	✓			
11/16/15 WT Library (1:20)	HiSeq_468	<input type="checkbox"/>	✓			
11/16/15 KO Library (1:20)	HiSeq_468	<input type="checkbox"/>	✓			
12/10/15 No Dox Library (1:35)	HiSeq_468	<input type="checkbox"/>	✓			
12/10/15 Dox Library (1:10)	HiSeq_468	<input type="checkbox"/>	✓			
12/11/15 WT Library (1:15)	HiSeq_468	<input type="checkbox"/>	✓			
12/11/15 KO Library (1:20)	HiSeq_468	<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

**Chip Lot #**

**Reagent Kit Lot #**

**Chip Comments :**

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
Modified: 5/13/2016 8:48:35 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

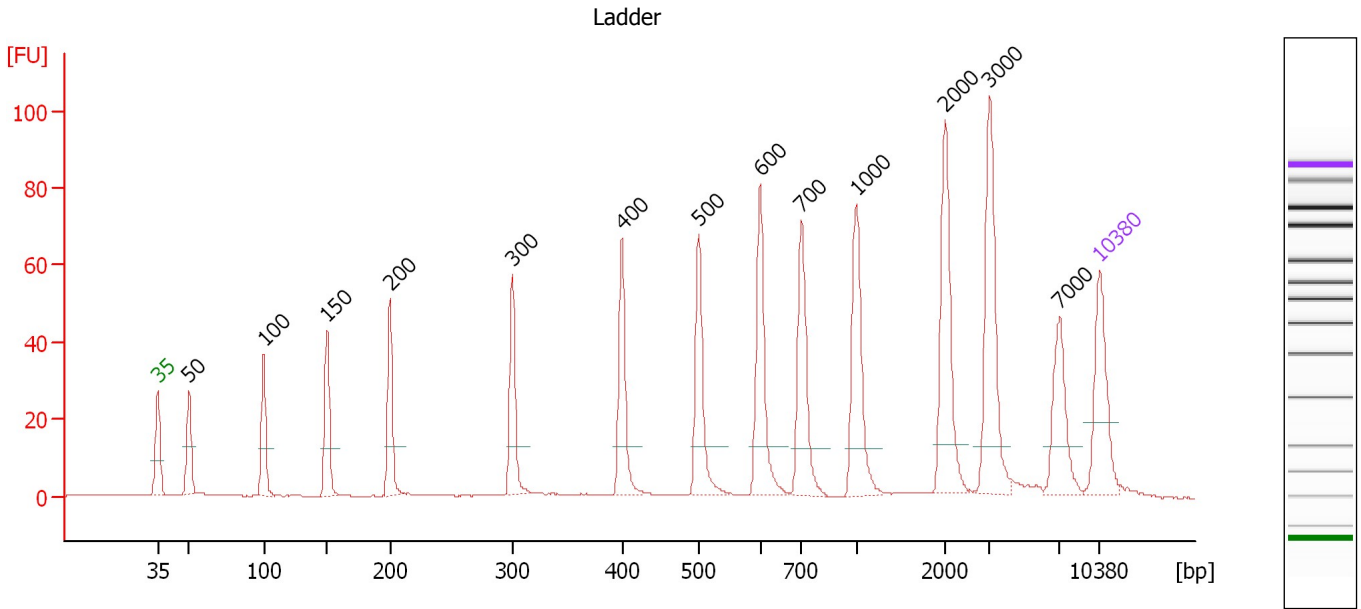
**Ladder**

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:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
 Modified: 5/13/2016 8:48:35 AM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.2

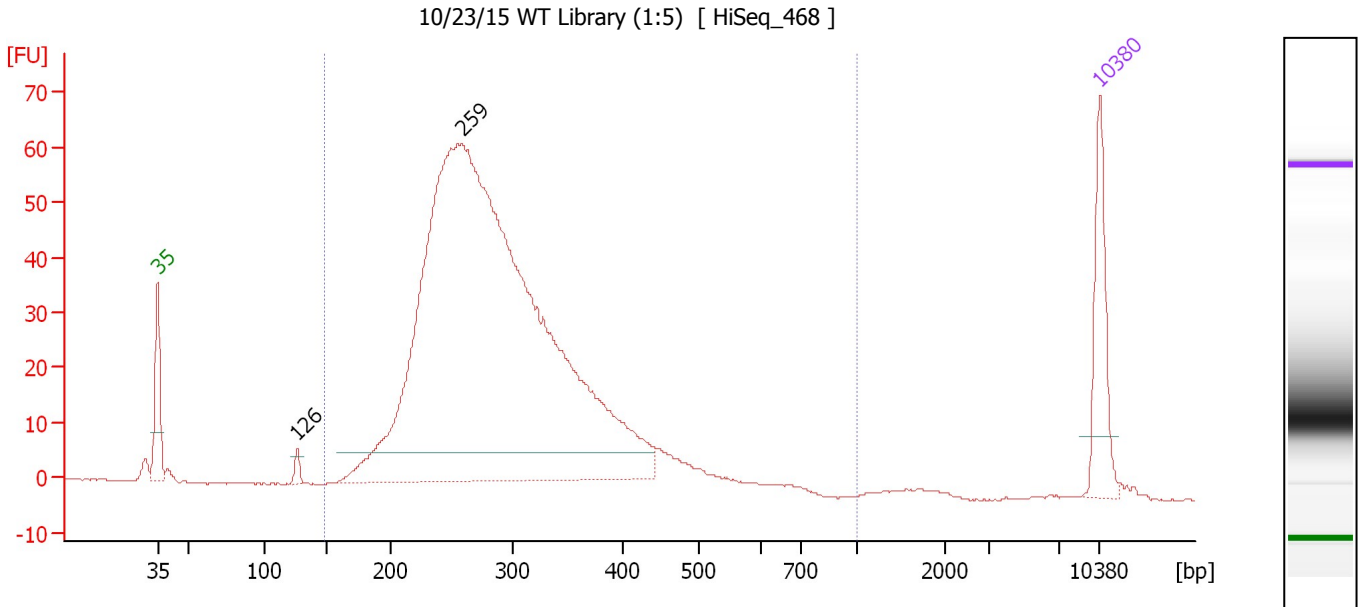
**Peak table for Ladder**

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	50.89
4	150	150.00	1,515.2	Ladder Peak	55.57
5	200	150.00	1,136.4	Ladder Peak	60.26
6	300	150.00	757.6	Ladder Peak	69.37
7	400	150.00	568.2	Ladder Peak	77.52
8	500	150.00	454.5	Ladder Peak	83.22
9	600	150.00	378.8	Ladder Peak	87.75
10	700	150.00	324.7	Ladder Peak	90.85
11	1,000	150.00	227.3	Ladder Peak	94.88
12	2,000	150.00	113.6	Ladder Peak	101.49
13	3,000	150.00	75.8	Ladder Peak	104.80
14	7,000	150.00	32.5	Ladder Peak	110.00
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
 Modified: 5/13/2016 8:48:35 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : 10/23/15 WT Library (1:5)**

Number of peaks found: 2                      Corr. Area 1: 964.0  
 Noise: 0.1

**Peak table for sample 1 : 10/23/15 WT Library (1:5)**

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	126	12.76	152.9		53.37
3	259	1,919.09	11,228.0		65.63
4	10,380	75.00	10.9	Upper Marker	113.00

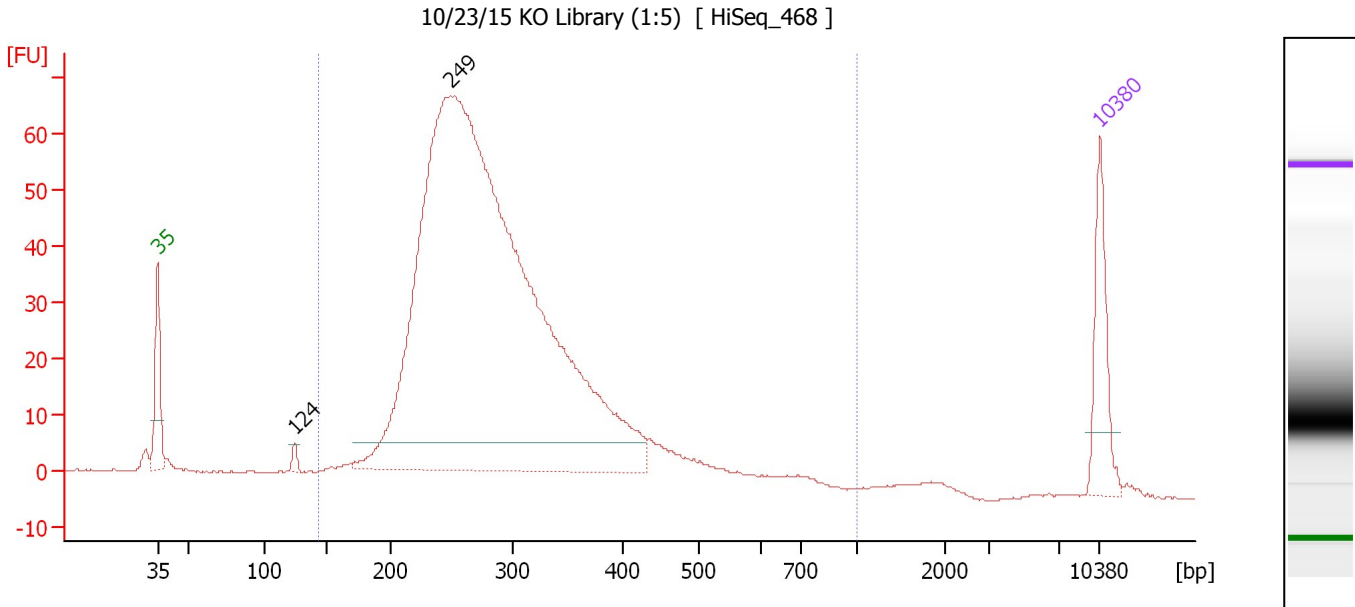
**Region table for sample 1 : 10/23/15 WT Library (1:5)**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
148	1,000	294	964.0	10,407.2	1,905.73	99	22.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
 Modified: 5/13/2016 8:48:35 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : 10/23/15 KO Library (1:5)**

Number of peaks found: 2                      Corr. Area 1: 1,054.7  
 Noise: 0.2

**Peak table for sample 2 : 10/23/15 KO Library (1:5)**

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	124	11.43	139.4		53.16
3	249	2,226.40	13,526.0		64.76
4	10,380	75.00	10.9	Upper Marker	113.00

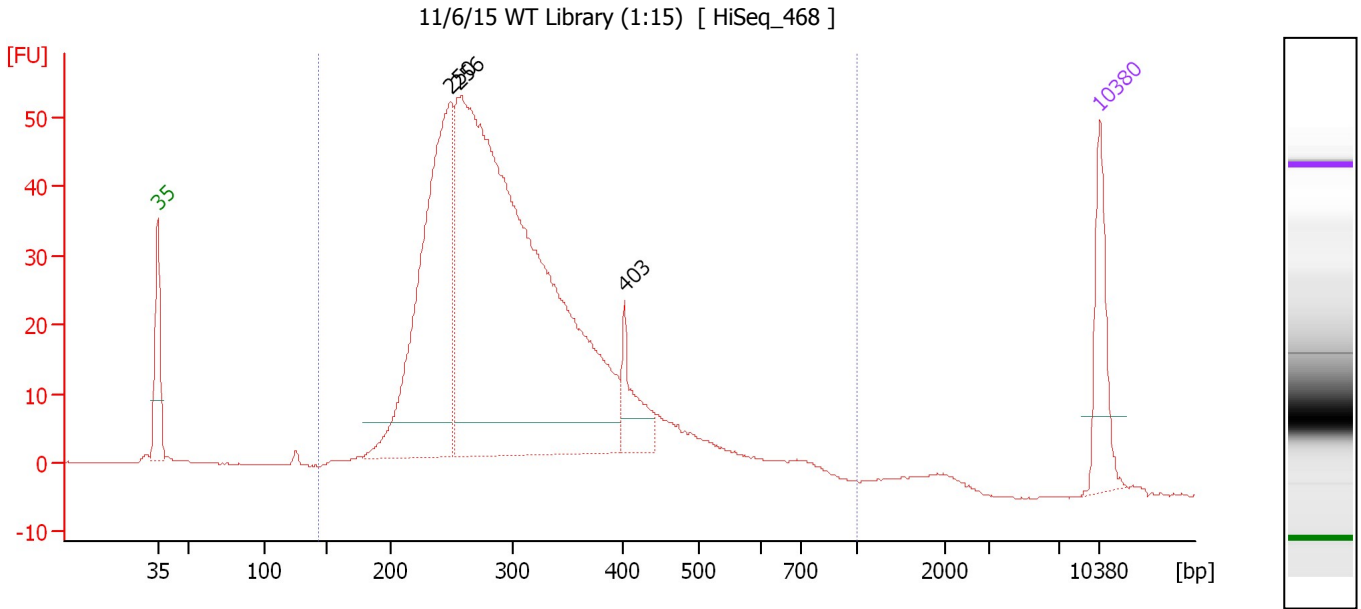
**Region table for sample 2 : 10/23/15 KO Library (1:5)**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
143	1,000	295	1,054.7	12,759.6	2,314.17	96	26.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
 Modified: 5/13/2016 8:48:35 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : 11/6/15 WT Library (1:15)**

Number of peaks found: 3                      Corr. Area 1: 957.4  
 Noise: 0.1

**Peak table for sample 3 : 11/6/15 WT Library (1:15)**

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	250	603.75	3,664.4		64.78
3	256	1,565.80	9,271.7		65.35
4	403	72.65	272.8		77.72
5	10,380	75.00	10.9	Upper Marker	113.00

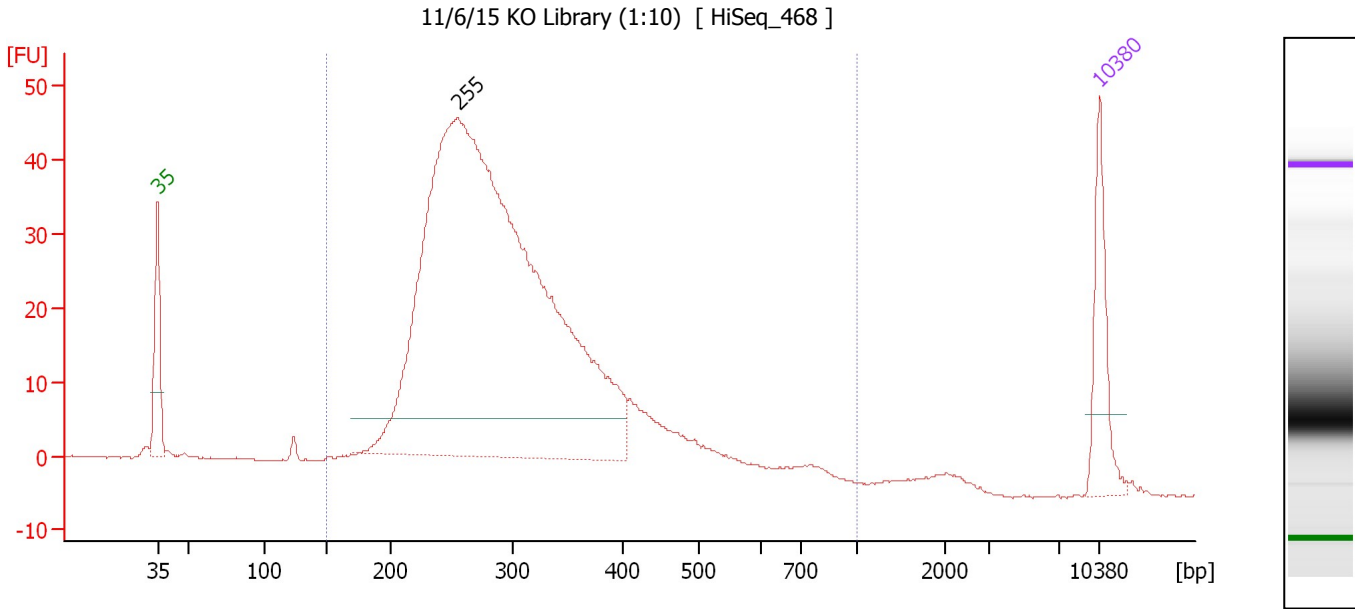
**Region table for sample 3 : 11/6/15 WT Library (1:15)**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
144	1,000	317	957.4	13,290.0	2,524.84	96	31.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
 Modified: 5/13/2016 8:48:35 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : 11/6/15 KO Library (1:10)**

Number of peaks found: 1                      Corr. Area 1: 788.0  
 Noise: 0.1

**Peak table for sample 4 : 11/6/15 KO Library (1:10)**

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	255	1,857.37	11,026.9		65.29
3	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 4 : 11/6/15 KO Library (1:10)**

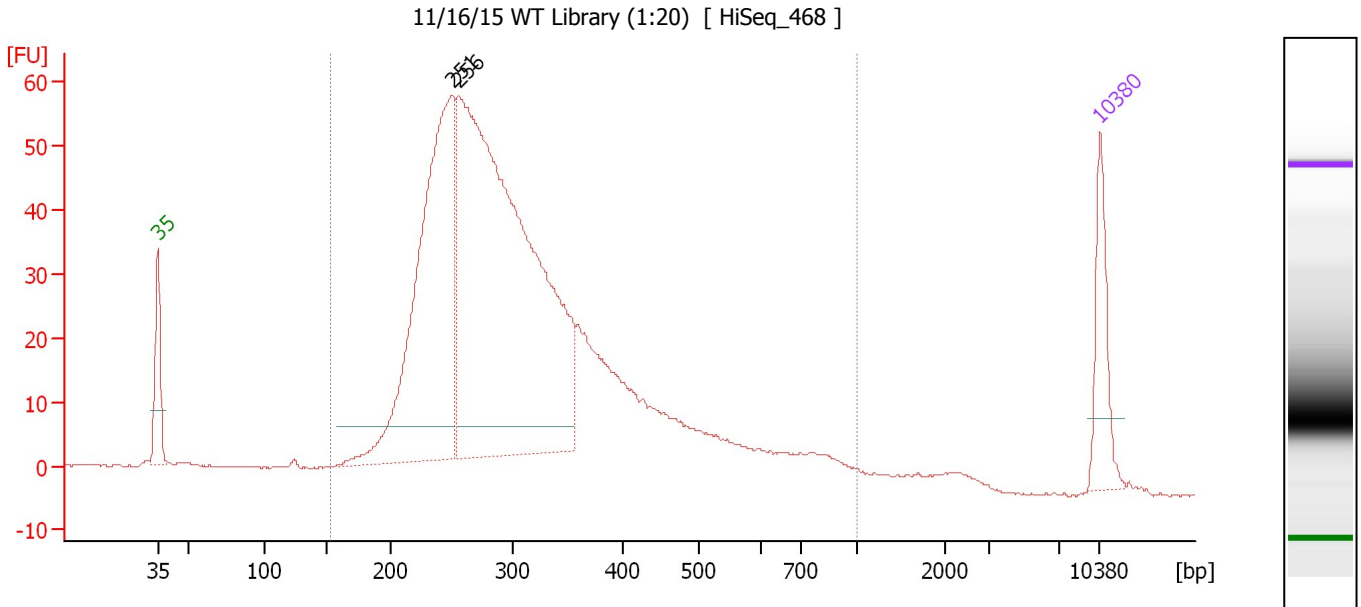
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
150	1,000	308	788.0	10,887.0	2,037.40	97	28.9



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
 Modified: 5/13/2016 8:48:35 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : 11/16/15 WT Library (1:20)**

Number of peaks found: 2                      Corr. Area 1: 1,056.8  
 Noise: 0.1

**Peak table for sample 5 : 11/16/15 WT Library (1:20)**

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	251	703.77	4,253.4		64.88
3	256	1,337.41	7,918.8		65.35
4	10,380	75.00	10.9	Upper Marker	113.00

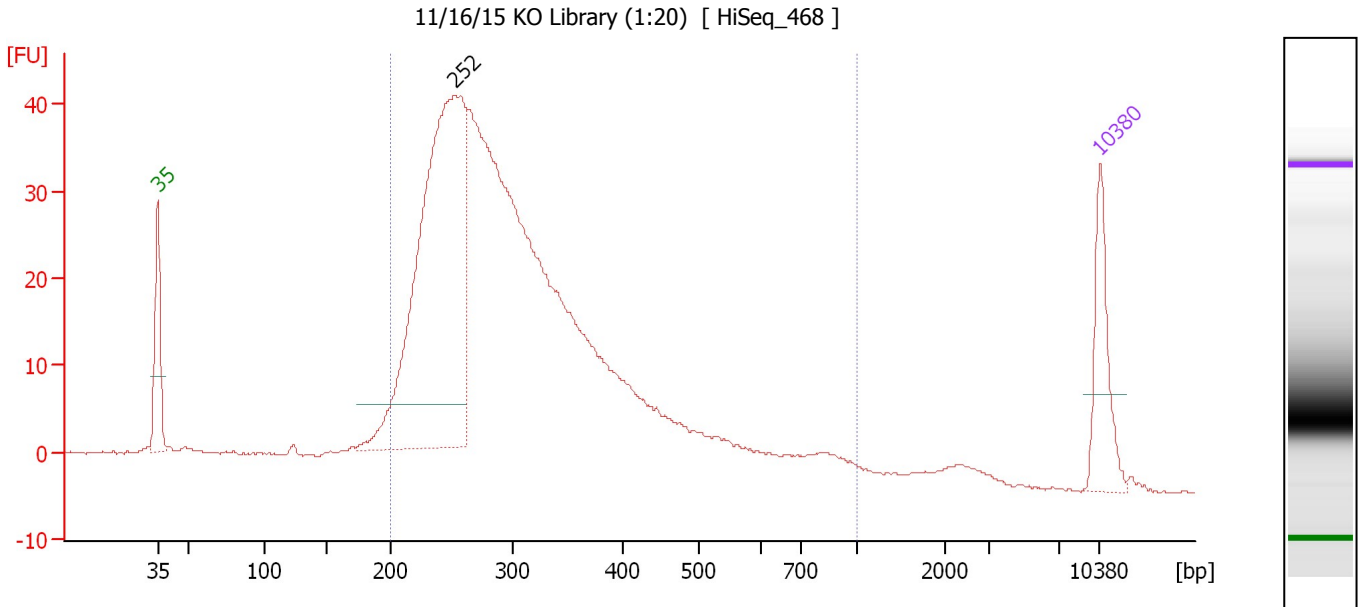
**Region table for sample 5 : 11/16/15 WT Library (1:20)**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
153	1,000	326	1,056.8	14,017.8	2,678.80	96	36.6

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
 Modified: 5/13/2016 8:48:35 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : 11/16/15 KO Library (1:20)**

Number of peaks found: 1                      Corr. Area 1: 726.3  
 Noise: 0.2

**Peak table for sample 6 : 11/16/15 KO Library (1:20)**

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	252	824.95	4,955.7		65.02
3	10,380	75.00	10.9	Upper Marker	113.00

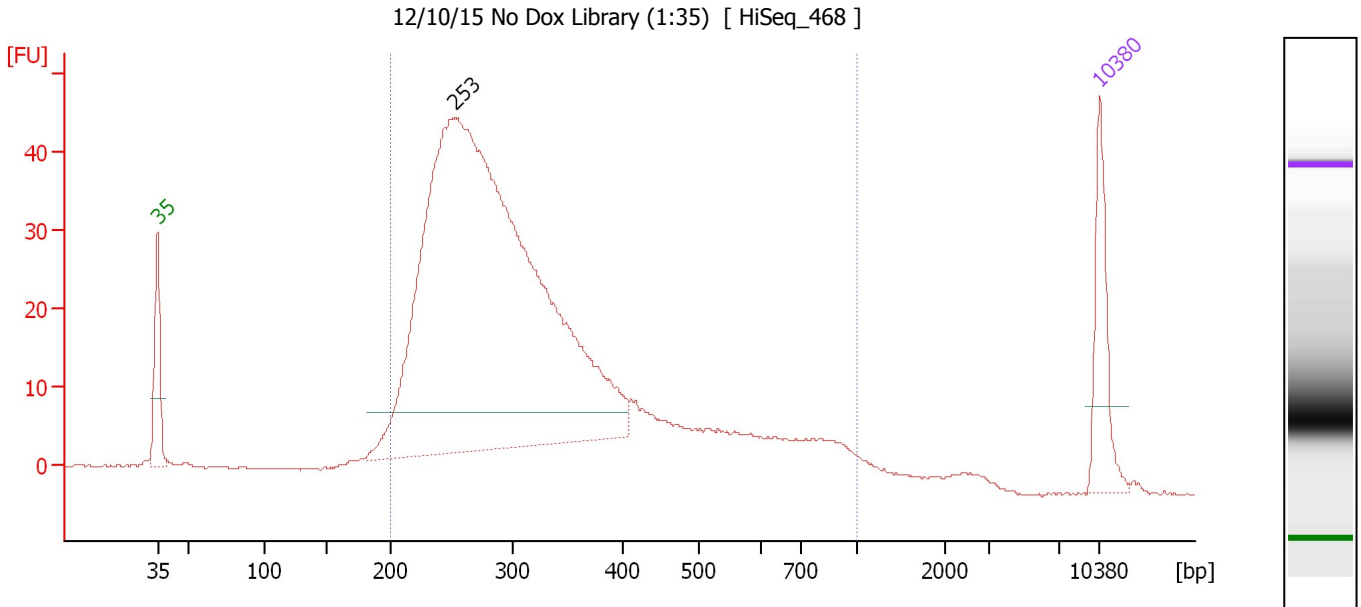
**Region table for sample 6 : 11/16/15 KO Library (1:20)**

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	323	726.3	12,299.5	2,361.95	93	35.3

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
 Modified: 5/13/2016 8:48:35 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : 12/10/15 No Dox Library (1:35)**

Number of peaks found: 1                      Corr. Area 1: 794.8  
 Noise: 0.2

**Peak table for sample 7 : 12/10/15 No Dox Library (1:35)**

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	253	1,760.69	10,541.5		65.09
3	10,380	75.00	10.9	Upper Marker	113.00

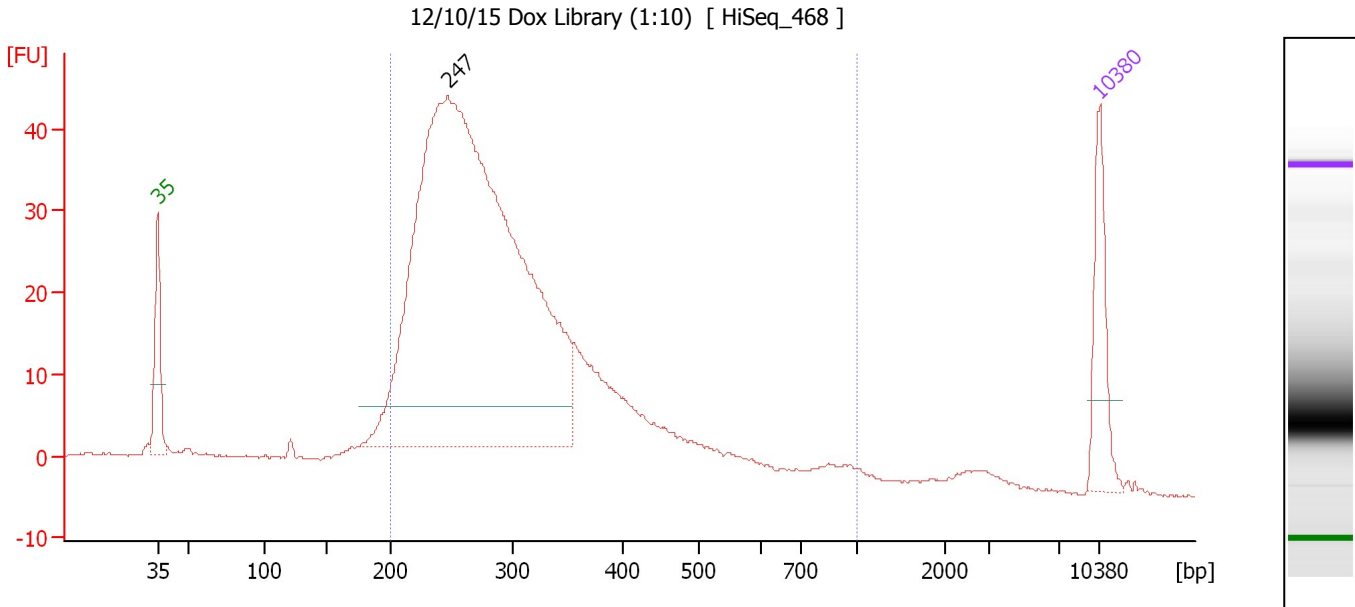
**Region table for sample 7 : 12/10/15 No Dox Library (1:35)**

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	341	794.8	11,028.2	2,160.23	95	40.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
 Modified: 5/13/2016 8:48:35 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : 12/10/15 Dox Library (1:10)**

Number of peaks found: 1                      Corr. Area 1: 717.7  
 Noise: 0.2

**Peak table for sample 8 : 12/10/15 Dox Library (1:10)**

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	247	1,745.71	10,690.4		64.58
3	10,380	75.00	10.9	Upper Marker	113.00

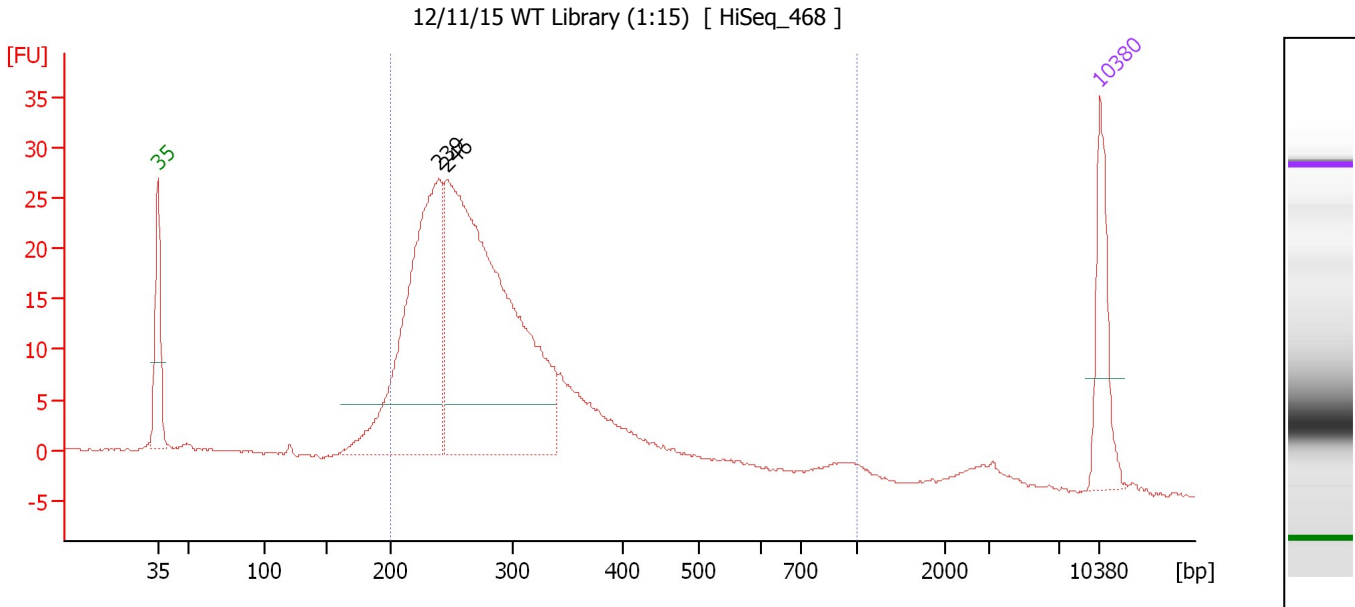
**Region table for sample 8 : 12/10/15 Dox Library (1:10)**

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	307	717.7	11,388.5	2,117.79	92	32.2

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
 Modified: 5/13/2016 8:48:35 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : 12/11/15 WT Library (1:15)**

Number of peaks found: 2                      Corr. Area 1: 416.3  
 Noise: 0.1

**Peak table for sample 9 : 12/11/15 WT Library (1:15)**

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	239	497.92	3,150.1		63.86
3	246	829.00	5,096.1		64.49
4	10,380	75.00	10.9	Upper Marker	113.00

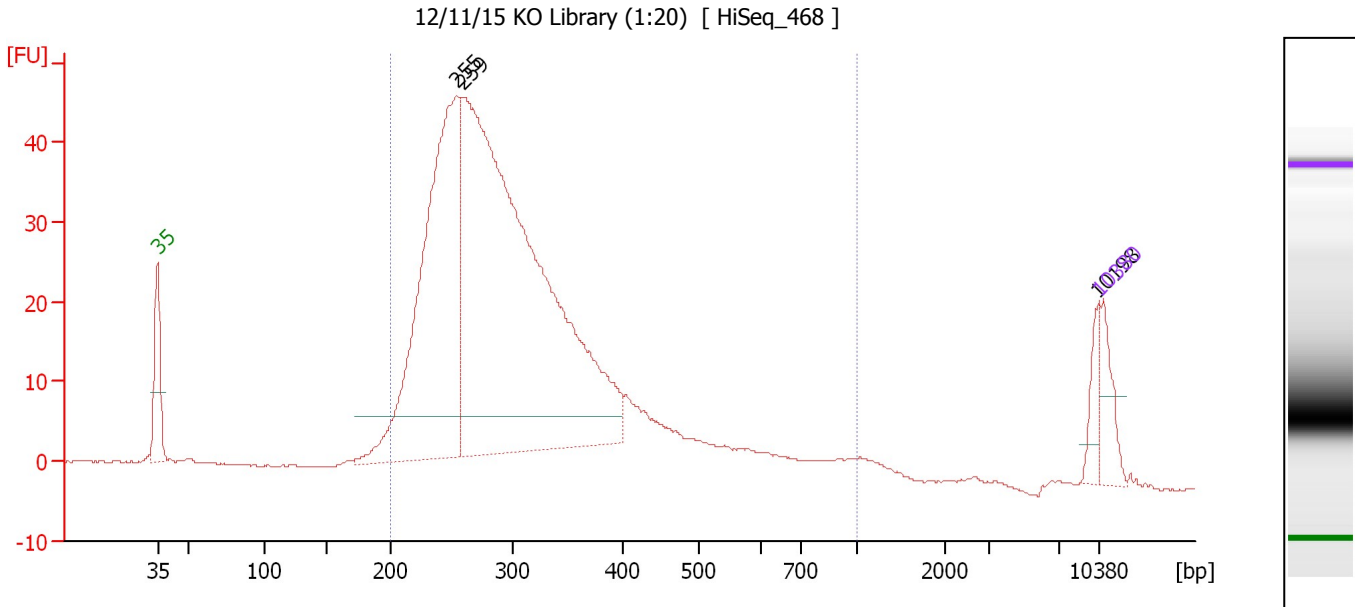
**Region table for sample 9 : 12/11/15 WT Library (1:15)**

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	298	416.3	8,390.6	1,521.20	90	33.4

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
 Modified: 5/13/2016 8:48:35 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : 12/11/15 KO Library (1:20)**

Number of peaks found: 3                      Corr. Area 1: 754.0  
 Noise: 0.1

**Peak table for sample 10 : 12/11/15 KO Library (1:20)**

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	259	1,225.40	7,291.8		65.24
3	259	2,401.28	14,027.9		65.67
4	10,198	45.80	6.8		112.84
5	10,380	75.00	10.9	Upper Marker	113.00

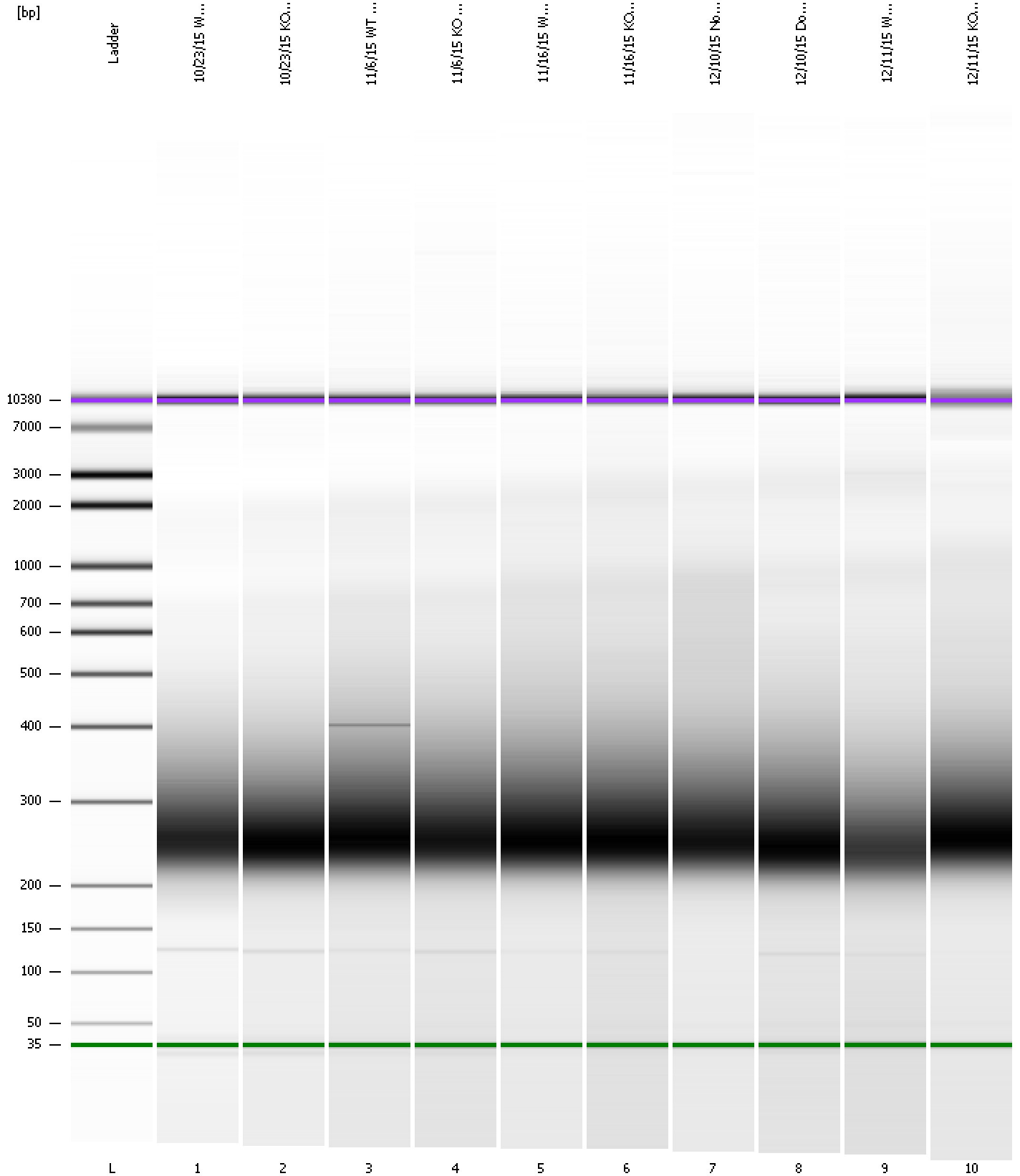
**Region table for sample 10 : 12/11/15 KO Library (1:20)**

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	324	754.0	21,668.1	4,181.20	92	35.5

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad

Created: 5/12/2016 3:39:57 PM  
Modified: 5/13/2016 8:48:35 AM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...data\2016-05-12\2016-05-12\_001\_HiSeq\_468\_Libraries\_1-10.xad  
 Created: 5/12/2016 3:39:57 PM  
 Modified: 5/13/2016 8:48:35 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		5/12/2016 4:21:15 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-05-12\2016-05-12_001.xad)		Instrument	Run		5/12/2016 3:40:02 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		5/12/2016 3:40:02 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		5/12/2016 3:40:02 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		5/12/2016 3:40:02 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		5/12/2016 3:40:02 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		5/12/2016 3:40:02 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		5/12/2016 3:40:02 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1