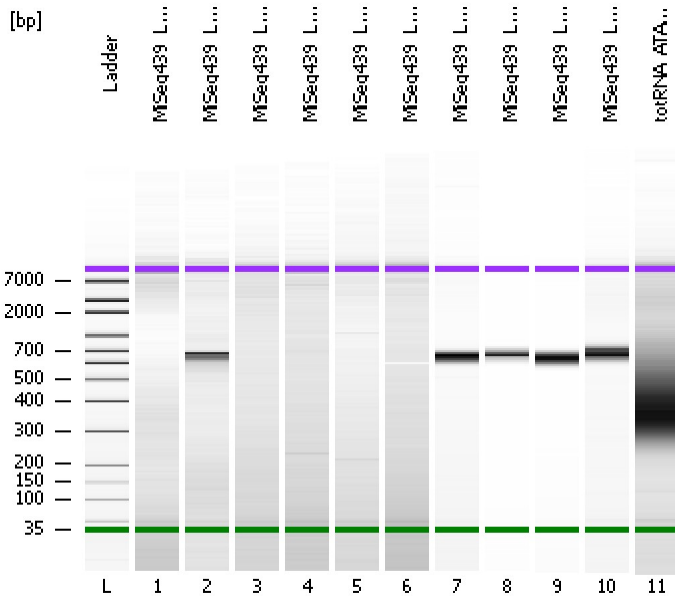


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
Modified: 2/14/2017 10:16:52 AM

**Electrophoresis File Run Summary**



Instrument Information:

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

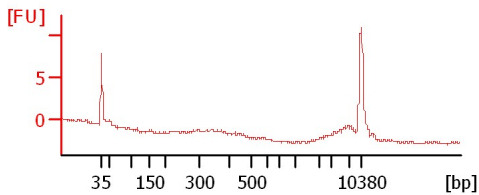
Assay Information:

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

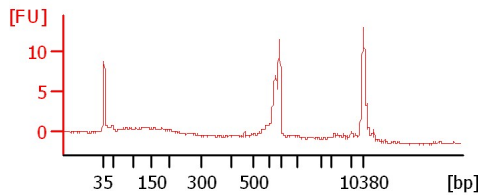
Chip Information:

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

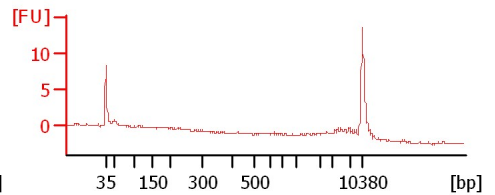
MiSeq439\_LIB1\_SS



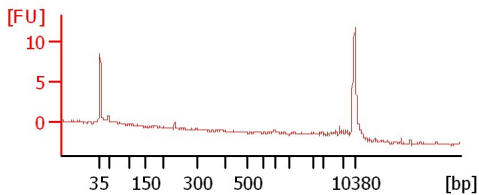
MiSeq439\_LIB8\_SS



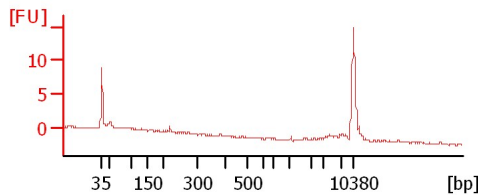
MiSeq439\_LIB9\_SS



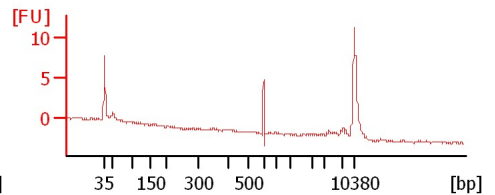
MiSeq439\_LIB11\_SS



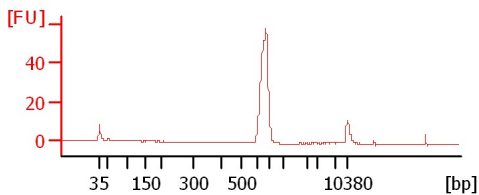
MiSeq439\_LIB10\_SS



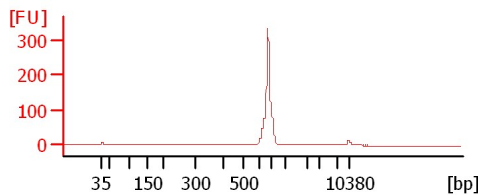
MiSeq439\_LIB12\_SS



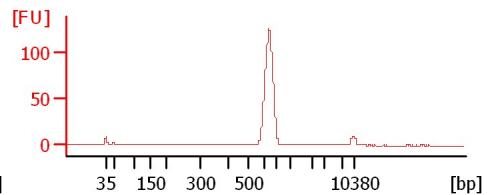
MiSeq439\_LIB13\_SS



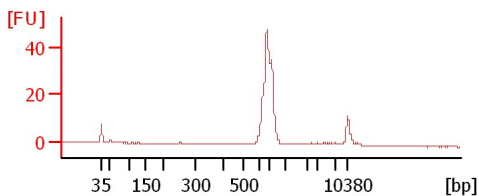
MiSeq439\_LIB14\_SS



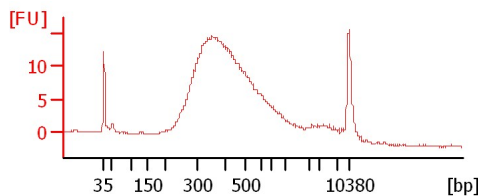
MiSeq439\_LIB15\_SS



MiSeq439\_LIB16\_SS



totRNA\_ATAC\_AY\_BC



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
Modified: 2/14/2017 10:16:52 AM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
MiSeq439_LIB1_SS		<input type="checkbox"/>	✓			
MiSeq439_LIB8_SS		<input type="checkbox"/>	✓			
MiSeq439_LIB9_SS		<input type="checkbox"/>	✓			
MiSeq439_LIB11_SS		<input type="checkbox"/>	✓			
MiSeq439_LIB10_SS		<input type="checkbox"/>	✓			
MiSeq439_LIB12_SS		<input type="checkbox"/>	✓			
MiSeq439_LIB13_SS		<input type="checkbox"/>	✓			
MiSeq439_LIB14_SS		<input type="checkbox"/>	✓			
MiSeq439_LIB15_SS		<input type="checkbox"/>	✓			
MiSeq439_LIB16_SS		<input type="checkbox"/>	✓			
totRNA_ATAC_AY_BC		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

**Chip Lot #**

**Reagent Kit Lot #**

**Chip Comments :**

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
Modified: 2/14/2017 10:16:52 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

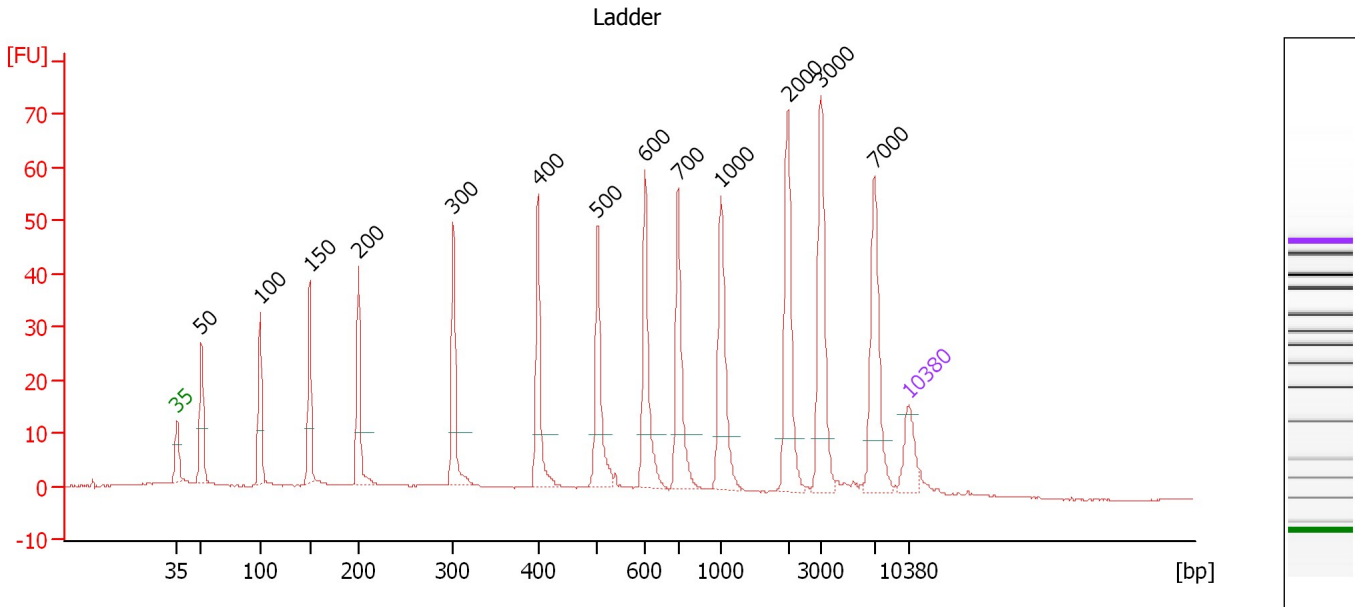
**Ladder**

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:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
 Modified: 2/14/2017 10:16:52 AM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.1

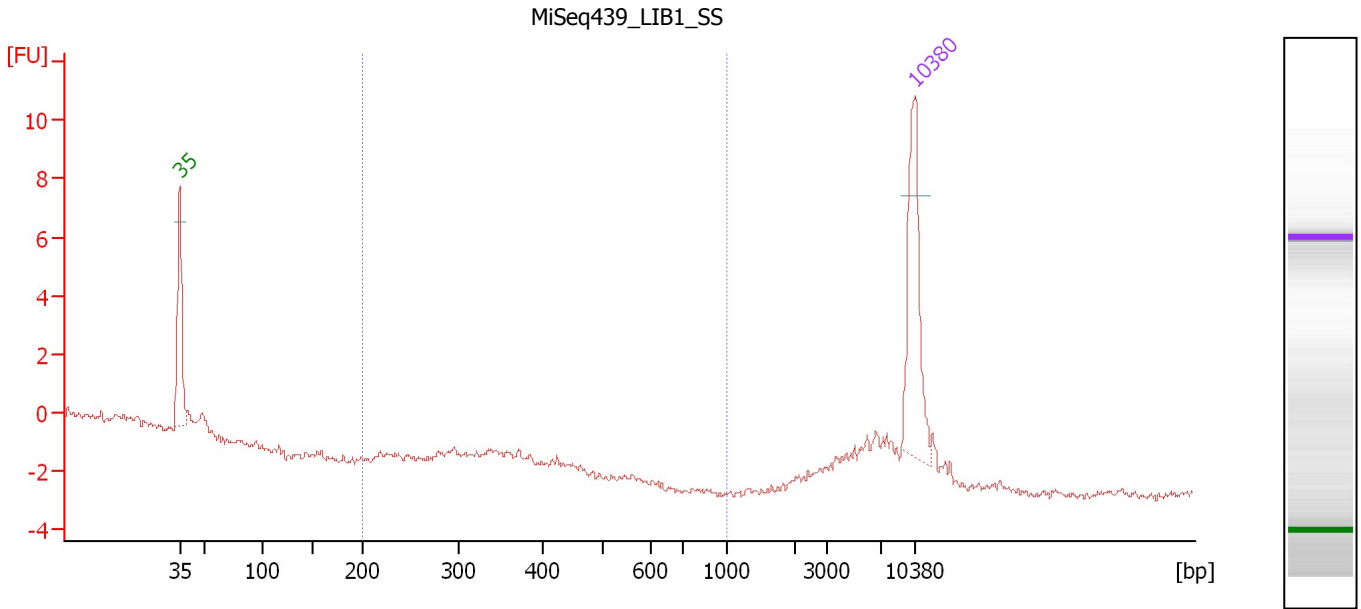
**Peak table for Ladder**

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	45.35
3	100	150.00	2,272.7	Ladder Peak	50.96
4	150	150.00	1,515.2	Ladder Peak	55.70
5	200	150.00	1,136.4	Ladder Peak	60.40
6	300	150.00	757.6	Ladder Peak	69.48
7	400	150.00	568.2	Ladder Peak	77.54
8	500	150.00	454.5	Ladder Peak	83.26
9	600	150.00	378.8	Ladder Peak	87.80
10	700	150.00	324.7	Ladder Peak	90.96
11	1,000	150.00	227.3	Ladder Peak	95.09
12	2,000	150.00	113.6	Ladder Peak	101.47
13	3,000	150.00	75.8	Ladder Peak	104.63
14	7,000	150.00	32.5	Ladder Peak	109.73
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
 Modified: 2/14/2017 10:16:52 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : MiSeq439\_LIB1\_SS**

Number of peaks found: 0    Corr. Area 1: 0.0  
 Noise: 0.1

**Peak table for sample 1 : MiSeq439\_LIB1\_SS**

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

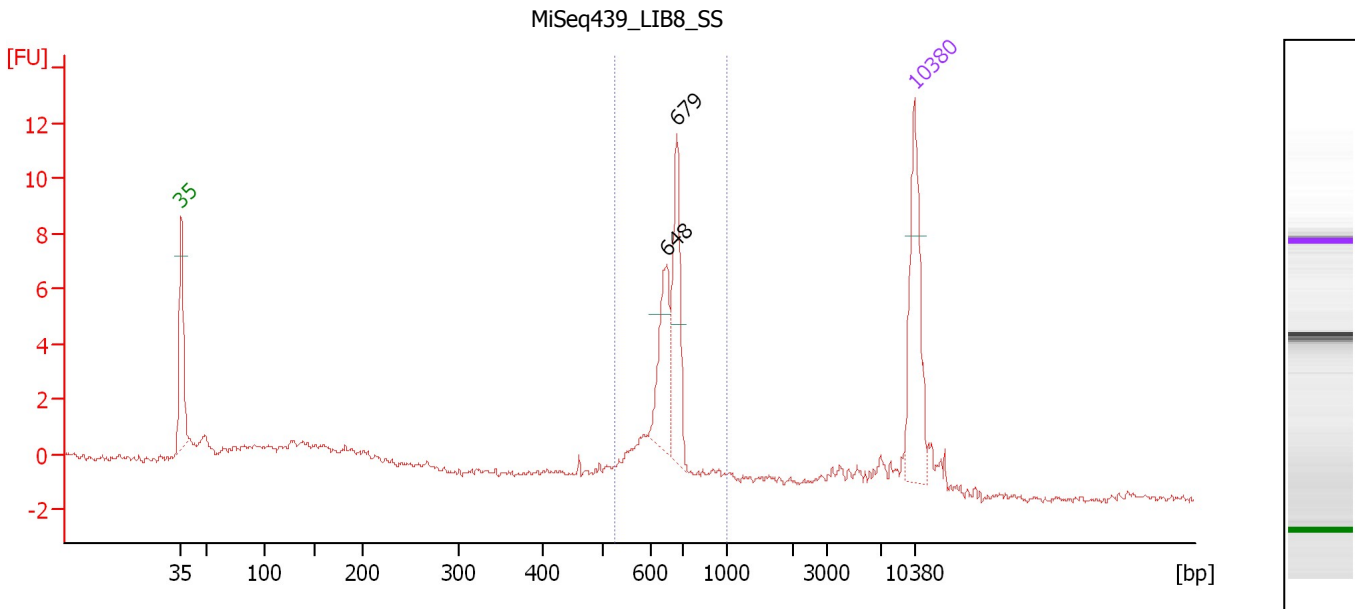
**Region table for sample 1 : MiSeq439\_LIB1\_SS**

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:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
 Modified: 2/14/2017 10:16:52 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : MiSeq439 LIB8 SS**

Number of peaks found: 2                      Corr. Area 1: 27.7  
 Noise: 0.1

**Peak table for sample 2 : MiSeq439 LIB8 SS**

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	648	61.83	144.5		89.33
3	679	75.30	168.0		90.29
4	10,380	75.00	10.9	Upper Marker	113.00

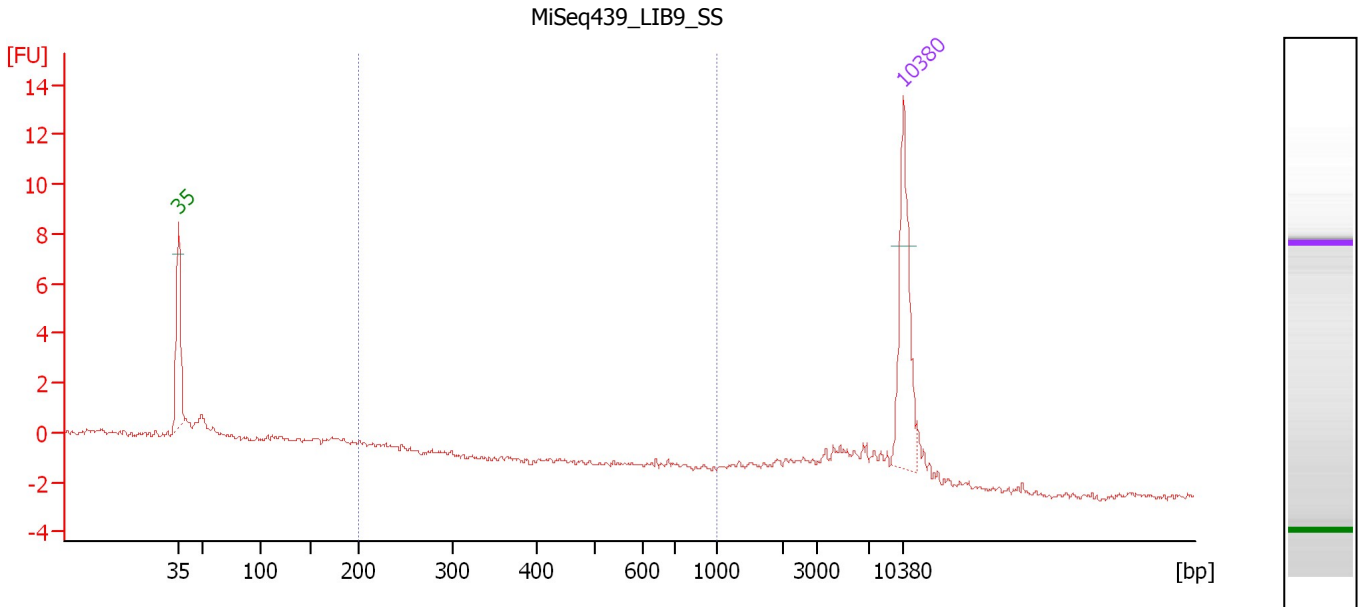
**Region table for sample 2 : MiSeq439 LIB8 SS**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
523	1,000	654	27.7	452.7	194.08	50	9.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
 Modified: 2/14/2017 10:16:52 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : MiSeq439\_LIB9\_SS**

Number of peaks found: 0                                  Corr. Area 1: 1.7  
 Noise: 0.1

**Peak table for sample 3 : MiSeq439\_LIB9\_SS**

Peak	Size [bp]	Conc. [pg/ $\mu$ l]	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

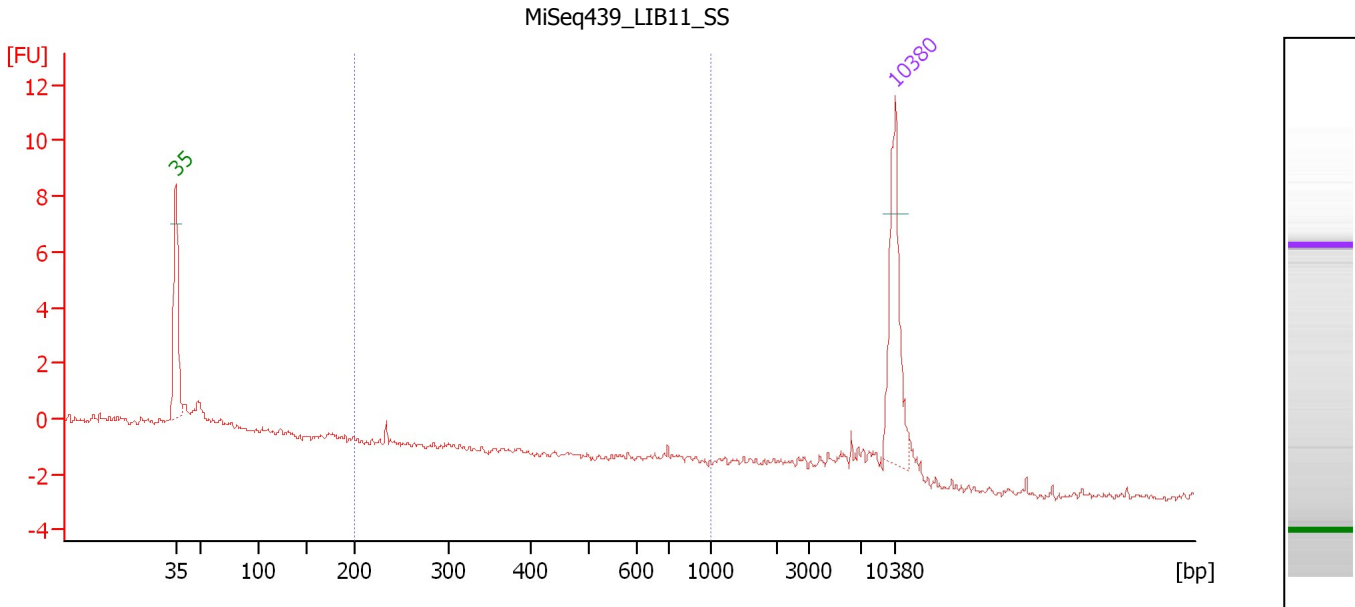
**Region table for sample 3 : MiSeq439\_LIB9\_SS**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ $\mu$ l]	% of Total	Size distribution in CV [%]
200	1,000	332	1.7	91.0	15.12	7	59.6

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
 Modified: 2/14/2017 10:16:52 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : MiSeq439 LIB11 SS**

Number of peaks found: 0                      Corr. Area 1: 1.5  
 Noise: 0.1

**Peak table for sample 4 : MiSeq439 LIB11 SS**

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 : MiSeq439 LIB11 SS**

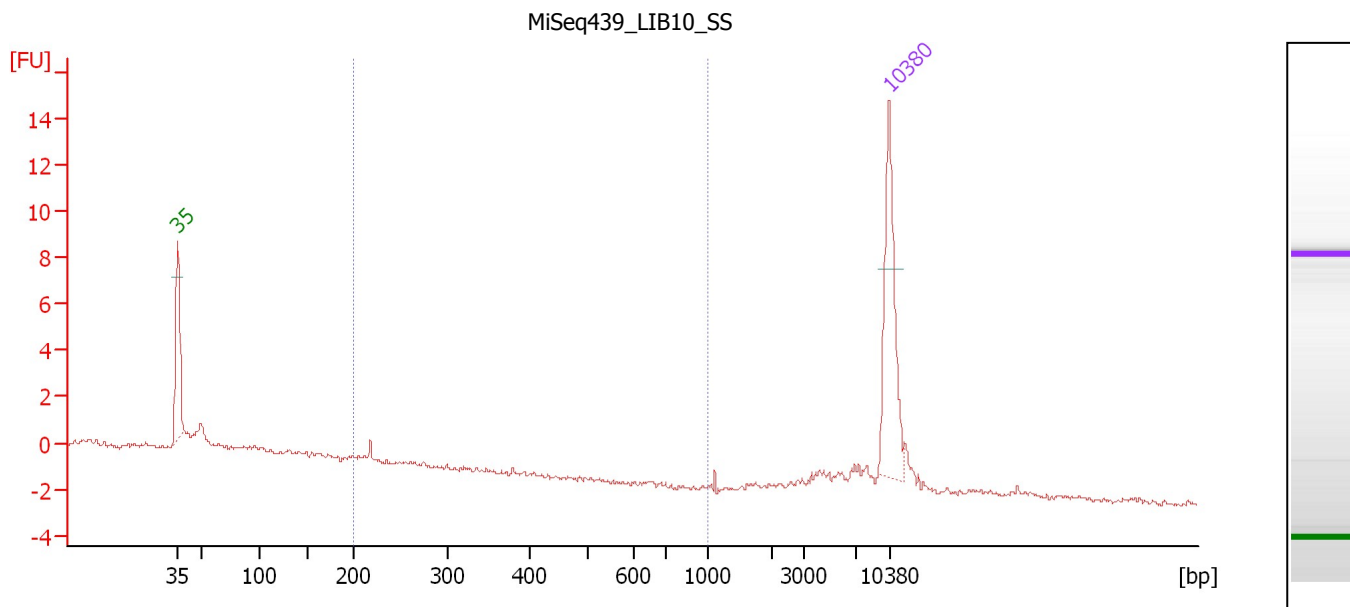
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	665	1.5	36.6	12.23	10	29.7



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
 Modified: 2/14/2017 10:16:52 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : MiSeq439\_LIB10\_SS**

Number of peaks found: 0                              Corr. Area 1: 0.6  
 Noise: 0.1

**Peak table for sample 5 : MiSeq439\_LIB10\_SS**

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

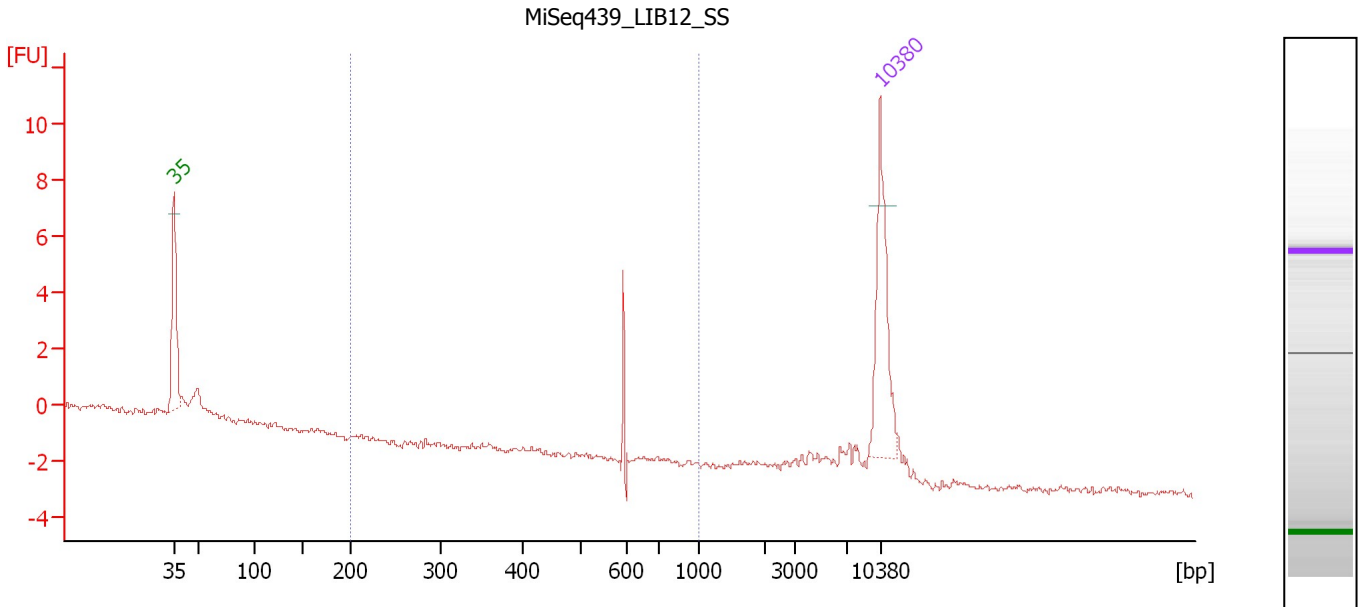
**Region table for sample 5 : MiSeq439\_LIB10\_SS**

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	216	0.6	35.9	5.11	4	4.6

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
 Modified: 2/14/2017 10:16:52 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : MiSeq439 LIB12 SS**

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

**Peak table for sample 6 : MiSeq439 LIB12 SS**

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

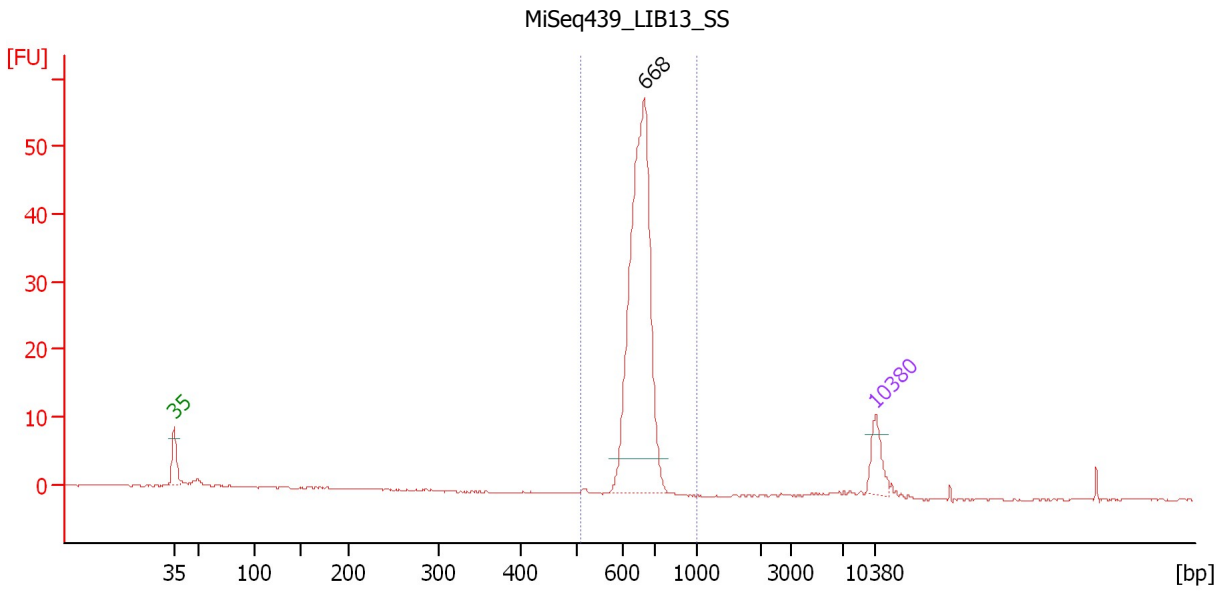
**Region table for sample 6 : MiSeq439 LIB12 SS**

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	594	1.4	27.1	10.61	14	0.3

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
 Modified: 2/14/2017 10:16:52 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : MiSeq439 LIB13 SS**

Number of peaks found: 1                      Corr. Area 1: 148.7  
 Noise: 0.1

**Peak table for sample 7 : MiSeq439 LIB13 SS**

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	668	1,181.77	2,680.0		89.95
3	10,380	75.00	10.9	Upper Marker	113.00

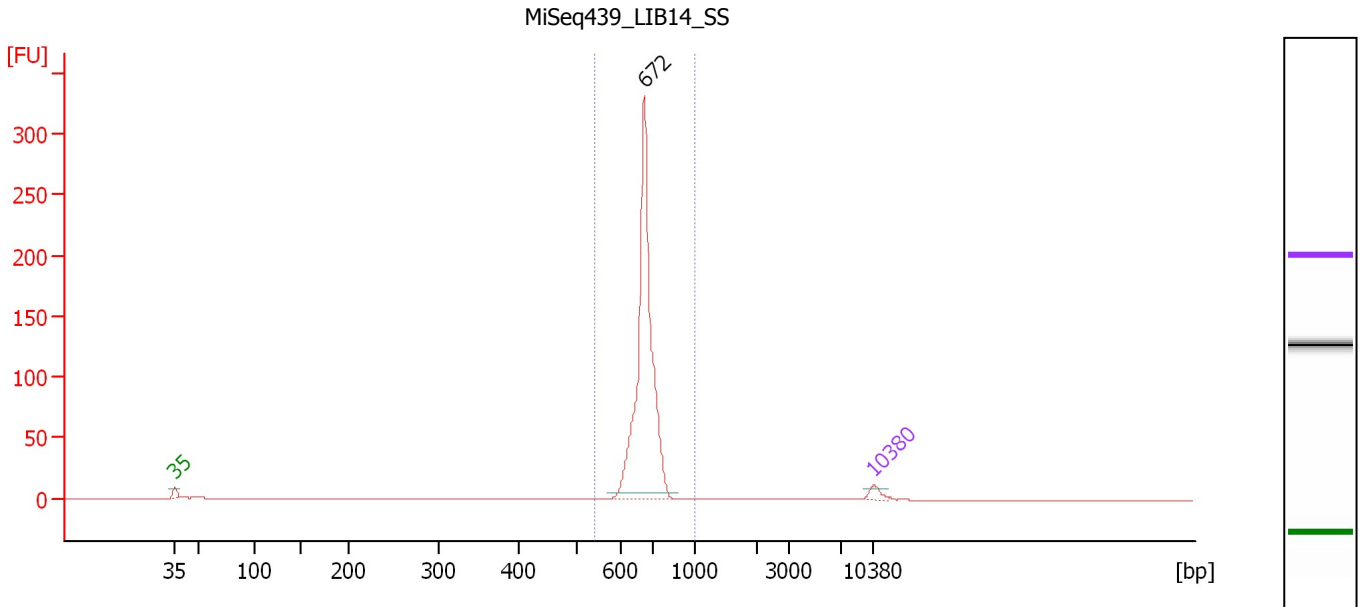
**Region table for sample 7 : MiSeq439 LIB13 SS**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
509	1,000	653	148.7	2,732.2	1,175.79	94	4.6

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
 Modified: 2/14/2017 10:16:52 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : MiSeq439 LIB14 SS**

Number of peaks found: 1                      Corr. Area 1: 532.3  
 Noise: 0.1

**Peak table for sample 8 : MiSeq439 LIB14 SS**

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	672	3,792.99	8,557.9		90.06
3	10,380	75.00	10.9	Upper Marker	113.00

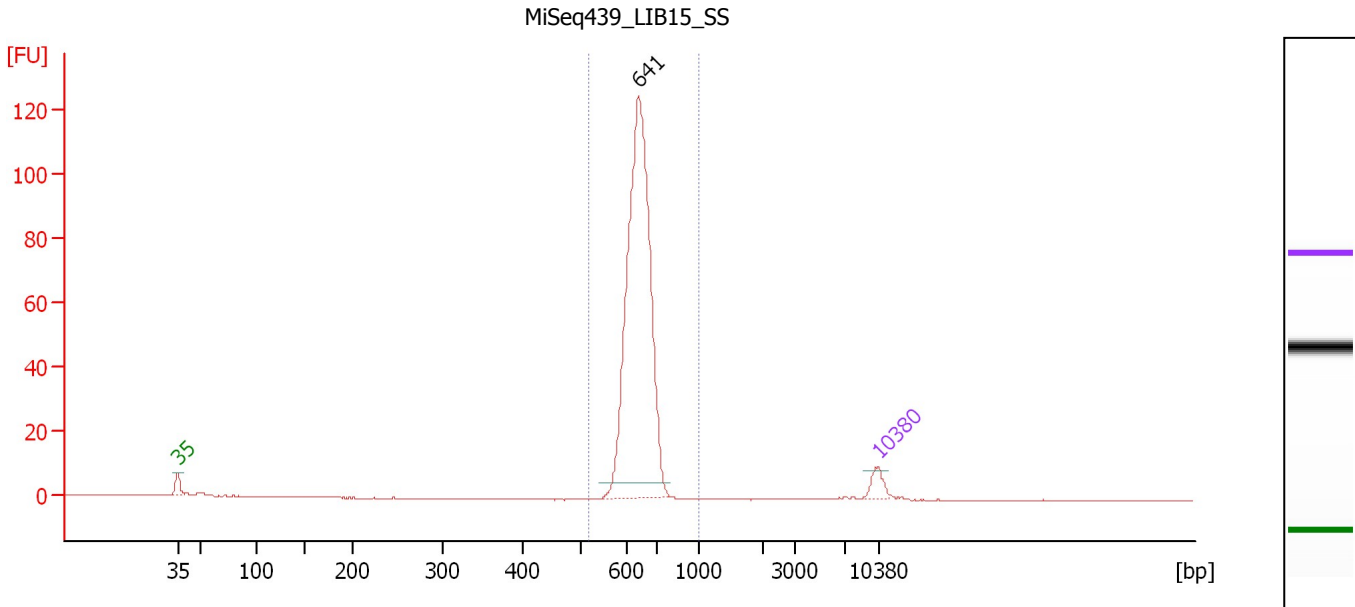
**Region table for sample 8 : MiSeq439 LIB14 SS**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
541	1,000	677	532.3	8,514.8	3,797.12	96	5.4

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
 Modified: 2/14/2017 10:16:52 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : MiSeq439\_LIB15\_SS**

Number of peaks found: 1                      Corr. Area 1: 365.3  
 Noise: 0.1

**Peak table for sample 9 : MiSeq439\_LIB15\_SS**

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	641	3,033.80	7,175.4		89.08
3	10,380	75.00	10.9	Upper Marker	113.00

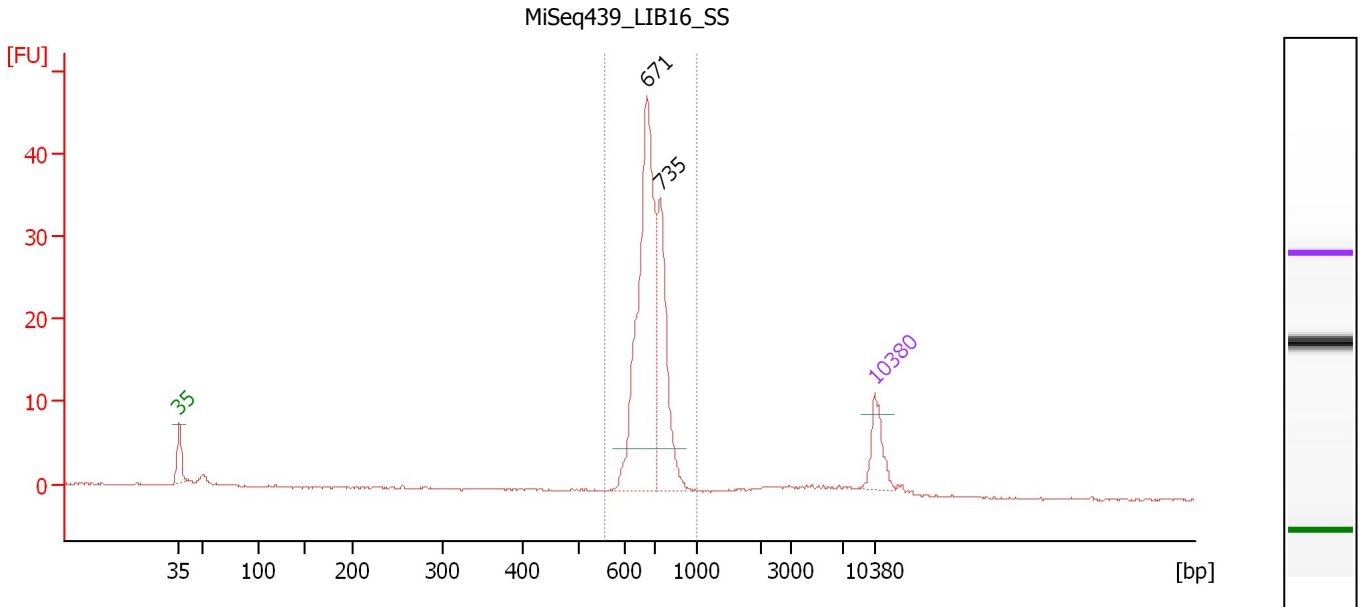
**Region table for sample 9 : MiSeq439\_LIB15\_SS**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
517	1,000	641	365.3	7,174.5	3,030.91	98	5.2

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
 Modified: 2/14/2017 10:16:52 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : MiSeq439 LIB16 SS**

Number of peaks found: 2                      Corr. Area 1: 142.2  
 Noise: 0.1

**Peak table for sample 10 : MiSeq439 LIB16 SS**

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	671	752.50	1,699.1		90.04
3	735	315.58	650.8		91.44
4	10,380	75.00	10.9	Upper Marker	113.00

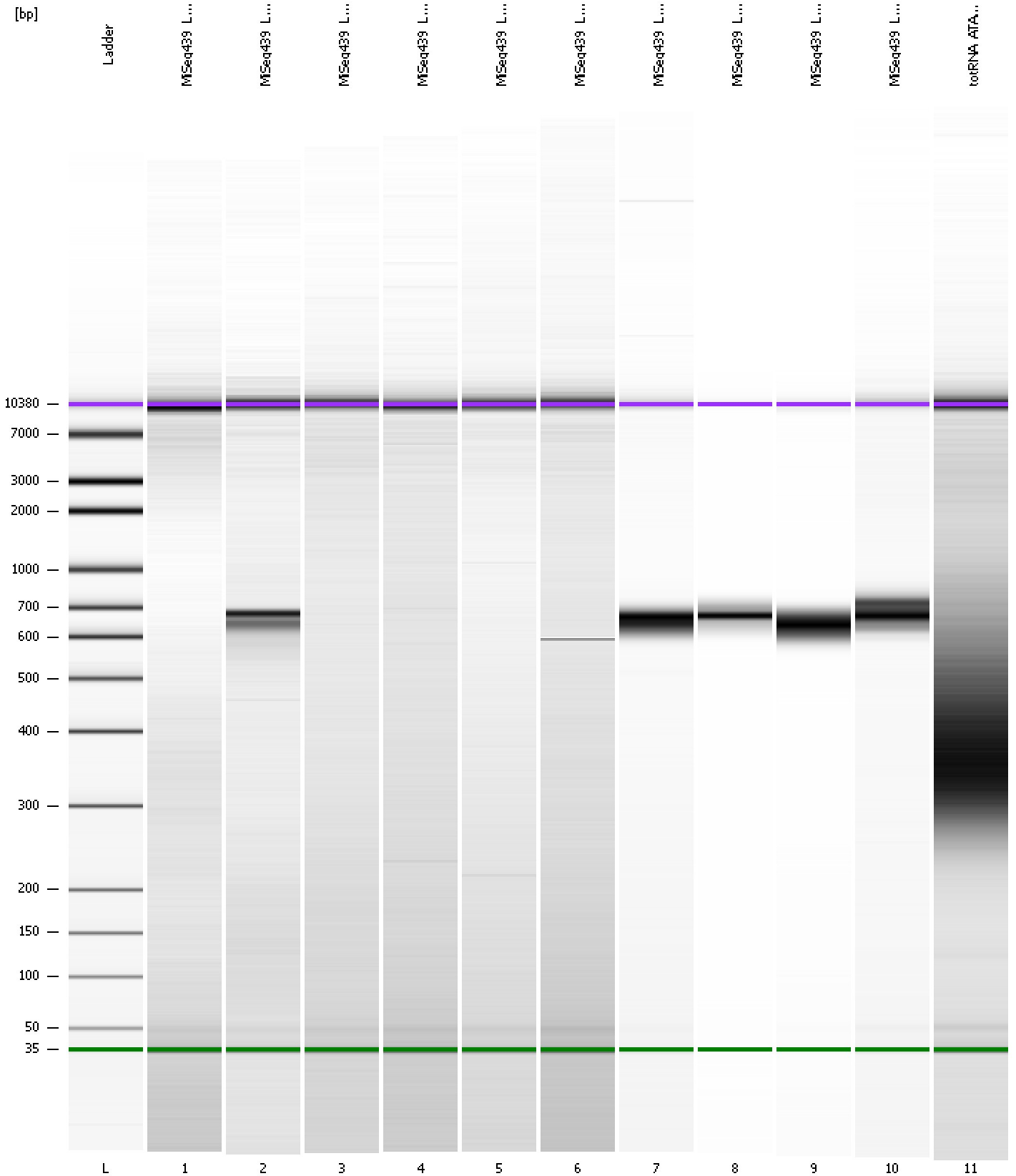
**Region table for sample 10 : MiSeq439 LIB16 SS**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
558	1,000	694	142.2	2,415.2	1,101.04	86	8.2

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
Modified: 2/14/2017 10:16:52 AM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-02-14\2017-02-14\_002.xad

Created: 2/14/2017 9:28:54 AM  
 Modified: 2/14/2017 10:16:52 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		2/14/2017 10:10:12 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2017-02-14\2017-02-14_002.xad)		Instrument	Run		2/14/2017 9:28:59 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		2/14/2017 9:28:59 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		2/14/2017 9:28:59 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		2/14/2017 9:28:59 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		2/14/2017 9:28:59 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		2/14/2017 9:28:59 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		2/14/2017 9:28:59 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1