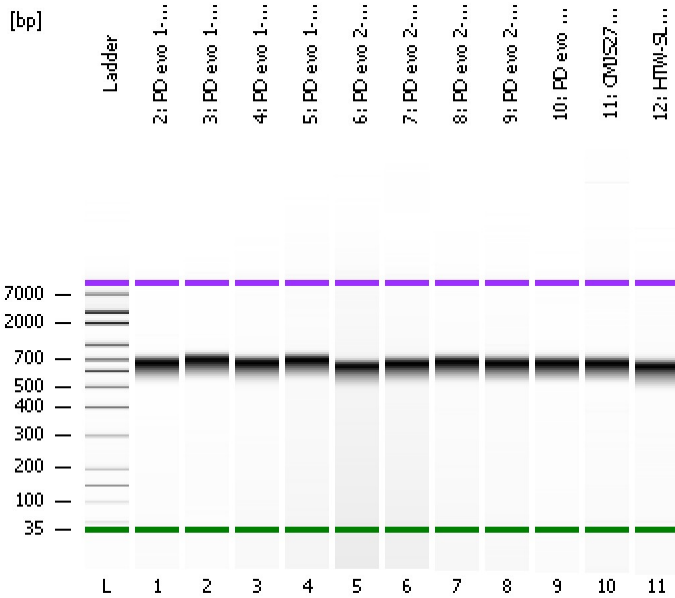


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
Modified: 1/19/2017 10:53:11 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

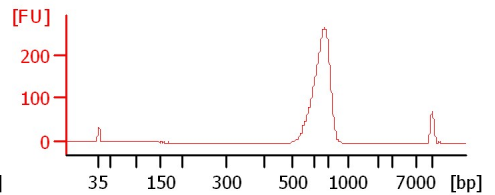
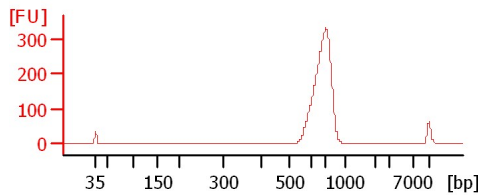
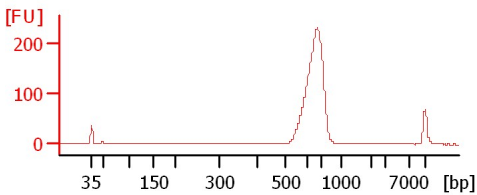
Chip Information:

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

2: PD evo 1-5-1 Size Sel. Library

3: PD evo 1-8-1 Size Sel. Library

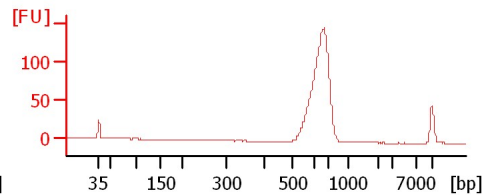
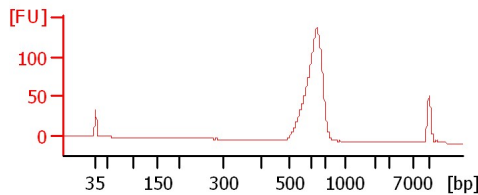
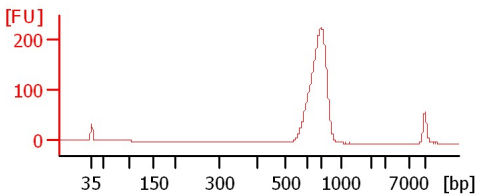
4: PD evo 1-11-1 Size Sel. Library



5: PD evo 1-12-1 Size Sel. Library

6: PD evo 2-3-1 Size Sel. Library

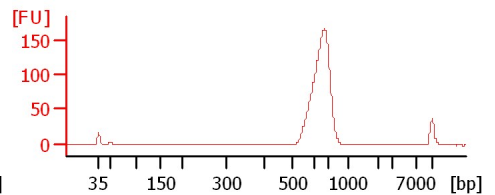
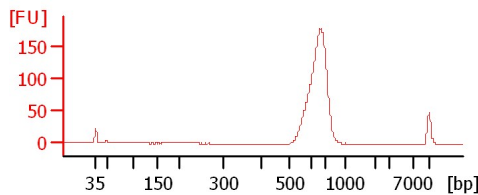
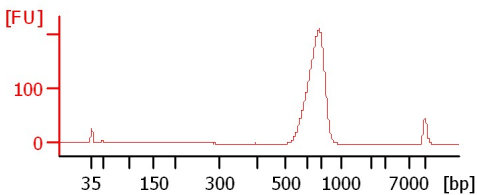
7: PD evo 2-5-1 Size Sel. Library



8: PD evo 2-8-1 Size Sel. Library

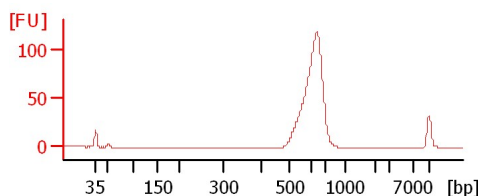
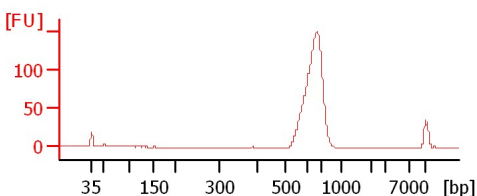
9: PD evo 2-11-1 Size Sel. Library

10: PD evo 2-14-1 Size Sel. Library



11: CMJS271 Size Sel. Library

12: HTW-SL1344 Size Sel. Library



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
2: PD evo 1-5-1 Size Sel. Library		<input type="checkbox"/>	✓			
3: PD evo 1-8-1 Size Sel. Library		<input type="checkbox"/>	✓			
4: PD evo 1-11-1 Size Sel. Library		<input type="checkbox"/>	✓			
5: PD evo 1-12-1 Size Sel. Library		<input type="checkbox"/>	✓			
6: PD evo 2-3-1 Size Sel. Library		<input type="checkbox"/>	✓			
7: PD evo 2-5-1 Size Sel. Library		<input type="checkbox"/>	✓			
8: PD evo 2-8-1 Size Sel. Library		<input type="checkbox"/>	✓			
9: PD evo 2-11-1 Size Sel. Library		<input type="checkbox"/>	✓			
10: PD evo 2-14-1 Size Sel.		<input type="checkbox"/>	✓			
11: CMJS271 Size Sel. Library		<input type="checkbox"/>	✓			
12: HTW-SL1344 Size Sel. Library		<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\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
Modified: 1/19/2017 10:53:11 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

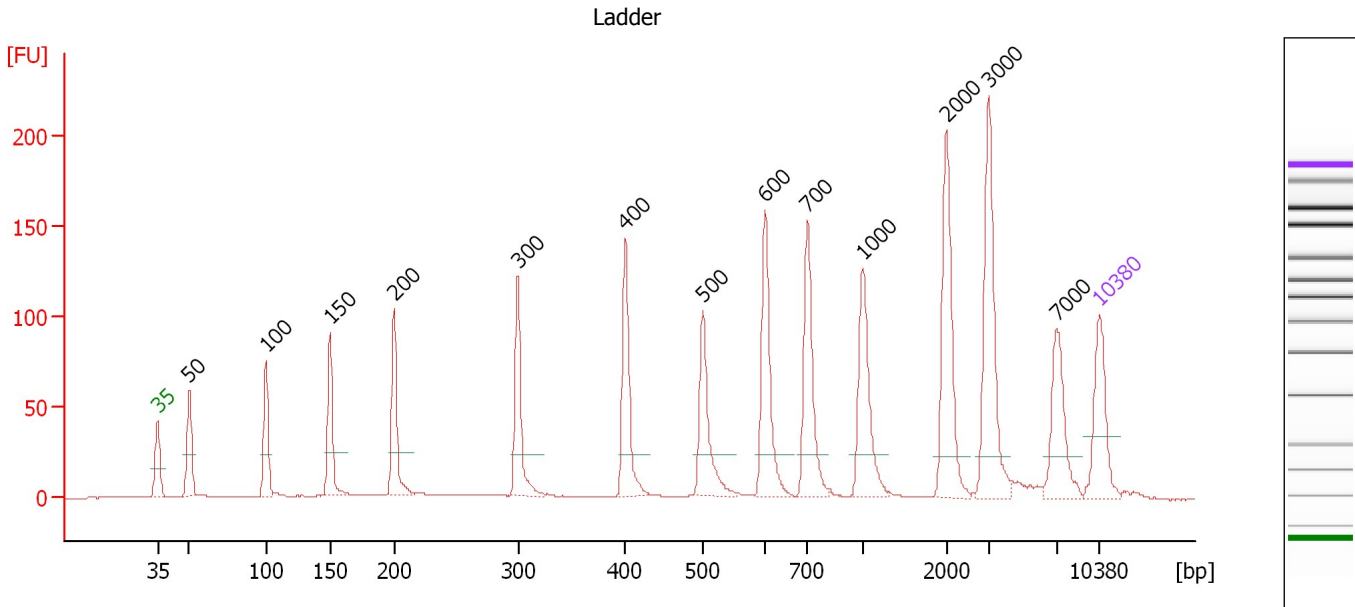
Ladder

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\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

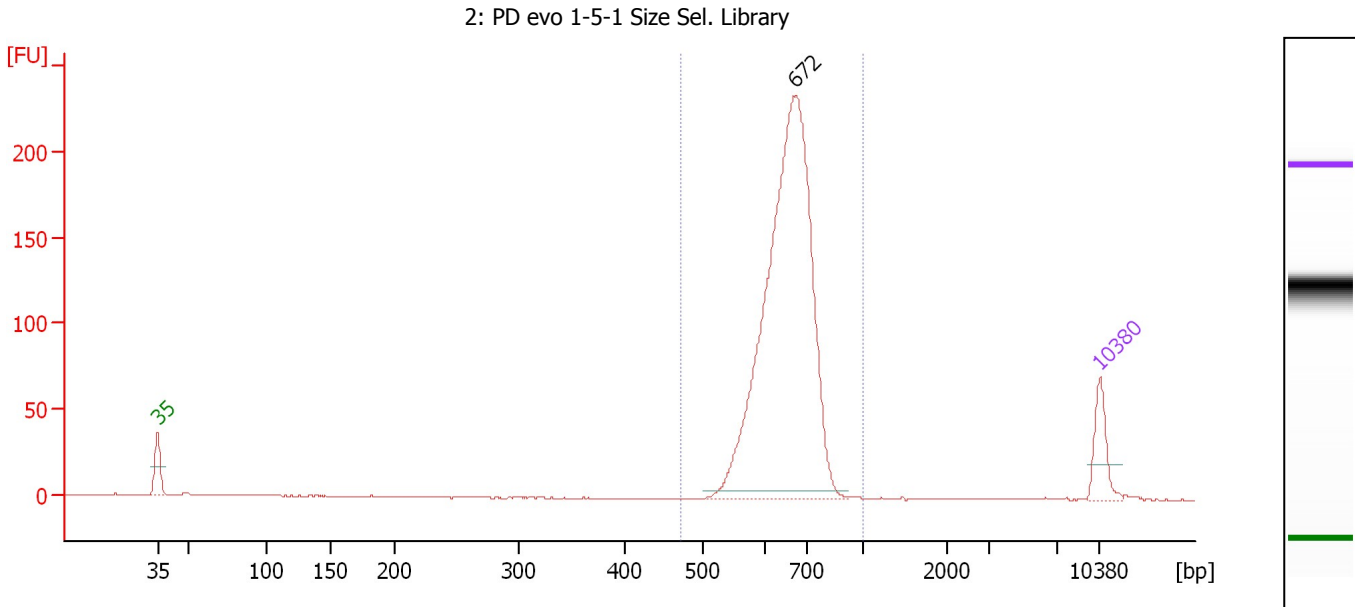
Peak table for Ladder

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	51.02
4	150	150.00	1,515.2	Ladder Peak	55.80
5	200	150.00	1,136.4	Ladder Peak	60.54
6	300	150.00	757.6	Ladder Peak	69.75
7	400	150.00	568.2	Ladder Peak	77.78
8	500	150.00	454.5	Ladder Peak	83.51
9	600	150.00	378.8	Ladder Peak	88.09
10	700	150.00	324.7	Ladder Peak	91.28
11	1,000	150.00	227.3	Ladder Peak	95.41
12	2,000	150.00	113.6	Ladder Peak	101.59
13	3,000	150.00	75.8	Ladder Peak	104.78
14	7,000	150.00	32.5	Ladder Peak	109.76
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : 2: PD evo 1-5-1 Size Sel. Library

Number of peaks found: 1 Corr. Area 1: 1,032.0
 Noise: 0.2

Peak table for sample 1 : 2: PD evo 1-5-1 Size Sel. Library

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	1,601.99	3,609.4		90.40
3	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 1 : 2: PD evo 1-5-1 Size Sel. Library

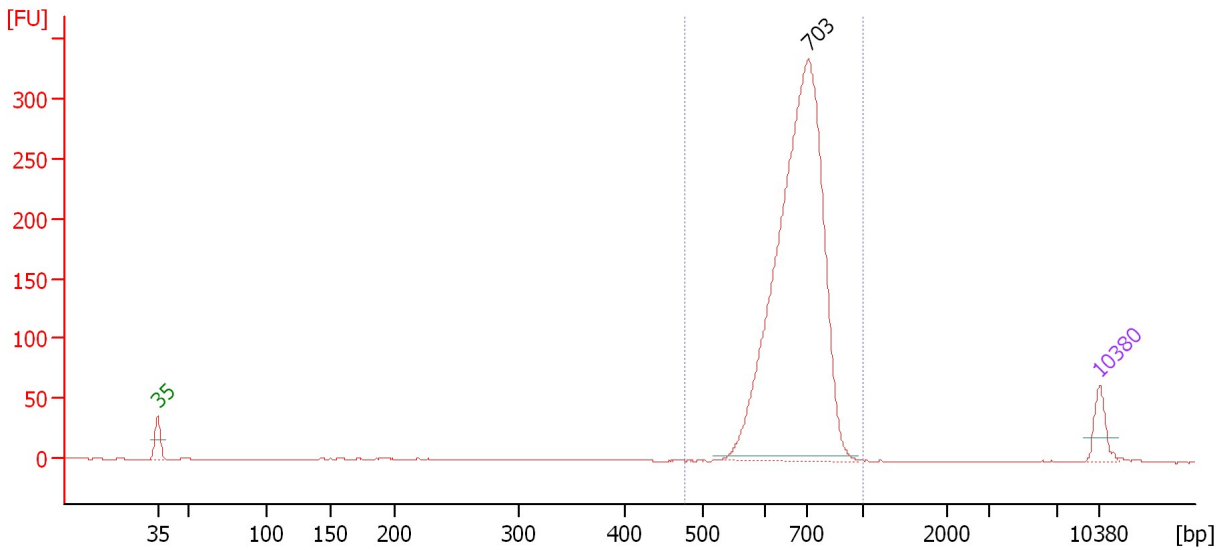
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
472	1,000	655	1,032.0	3,706.8	1,593.03	98	8.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 AM

Electropherogram Summary Continued ...

3: PD evo 1-8-1 Size Sel. Library



Overall Results for sample 2 : 3: PD evo 1-8-1 Size Sel. Library

Number of peaks found: 1 Corr. Area 1: 1,479.4
 Noise: 0.2

Peak table for sample 2 : 3: PD evo 1-8-1 Size Sel. Library

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	703	2,553.60	5,501.6		91.32
3	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 2 : 3: PD evo 1-8-1 Size Sel. Library

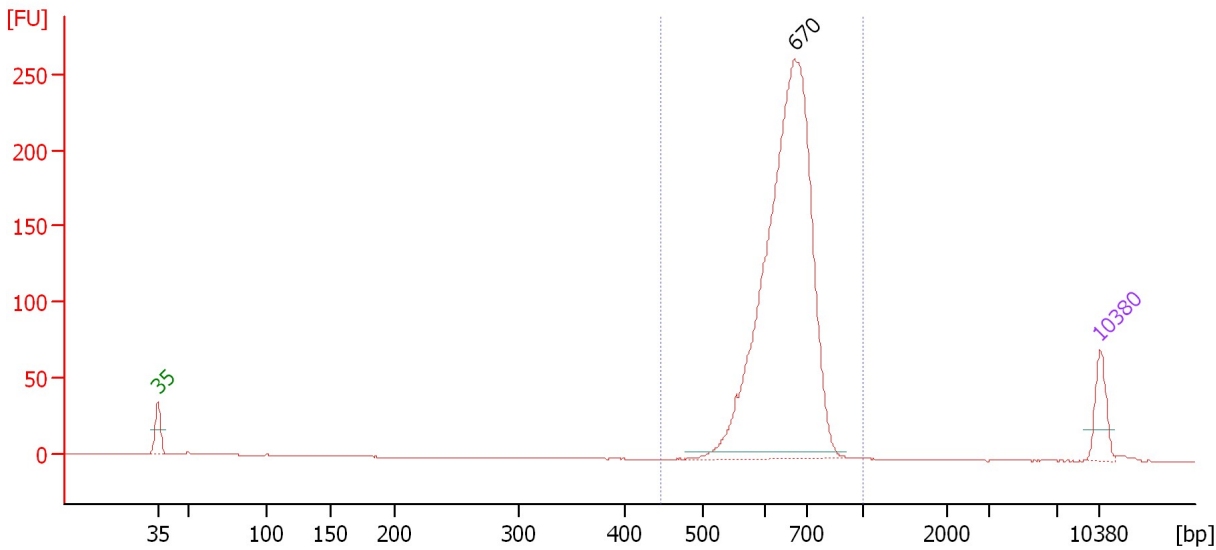
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
478	1,000	689	1,479.4	5,610.1	2,527.12	98	10.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 AM

Electropherogram Summary Continued ...

4: PD evo 1-11-1 Size Sel. Library



Overall Results for sample 3 : 4: PD evo 1-11-1 Size Sel. Library

Number of peaks found: 1 Corr. Area 1: 1,164.7
 Noise: 0.2

Peak table for sample 3 : 4: PD evo 1-11-1 Size Sel. Library

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	670	1,872.64	4,236.5		90.31
3	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 3 : 4: PD evo 1-11-1 Size Sel. Library

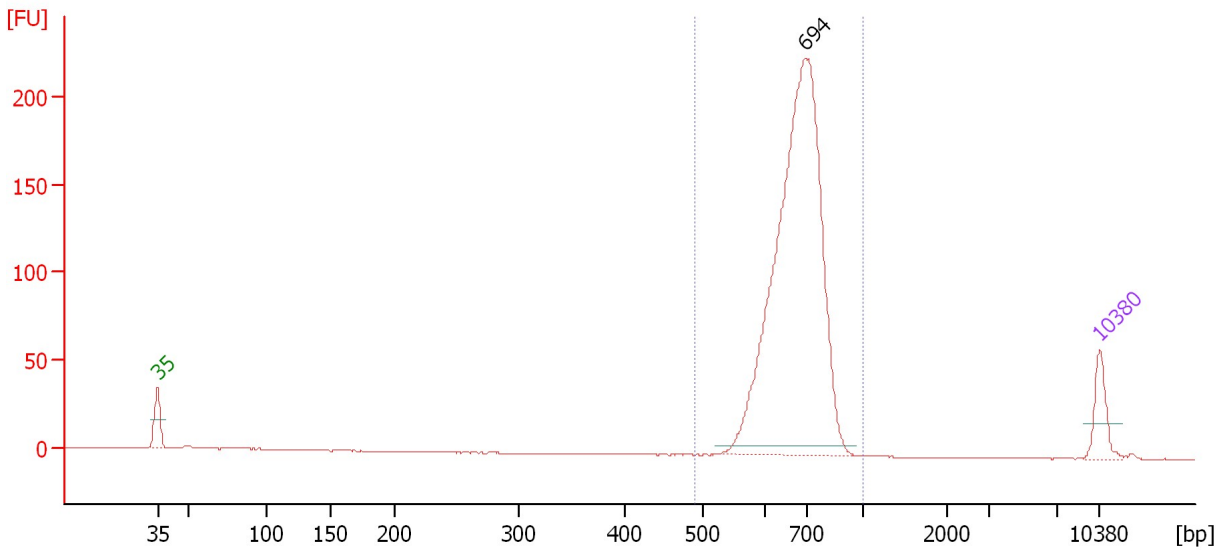
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
446	1,000	654	1,164.7	4,379.1	1,877.08	99	8.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 AM

Electropherogram Summary Continued ...

5: PD evo 1-12-1 Size Sel. Library



Overall Results for sample 4 : 5: PD evo 1-12-1 Size Sel. Library

Number of peaks found: 1 Corr. Area 1: 975.0
 Noise: 0.2

Peak table for sample 4 : 5: PD evo 1-12-1 Size Sel. Library

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	694	1,768.23	3,860.5		91.09
3	10,380	75.00	10.9	Upper Marker	113.00

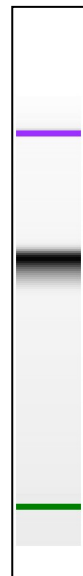
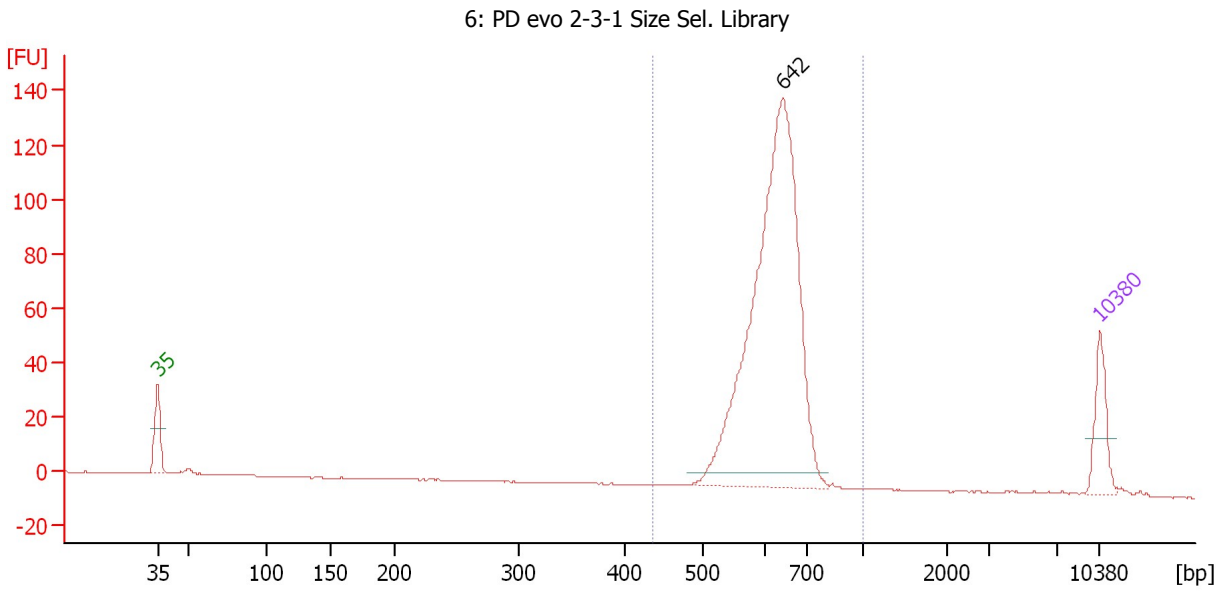
Region table for sample 4 : 5: PD evo 1-12-1 Size Sel. Library

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
491	1,000	685	975.0	3,921.1	1,756.55	96	9.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : 6: PD evo 2-3-1 Size Sel. Library

Number of peaks found: 1 Corr. Area 1: 602.0
 Noise: 0.2

Peak table for sample 5 : 6: PD evo 2-3-1 Size Sel. Library

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	642	1,162.42	2,742.2		89.44
3	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 5 : 6: PD evo 2-3-1 Size Sel. Library

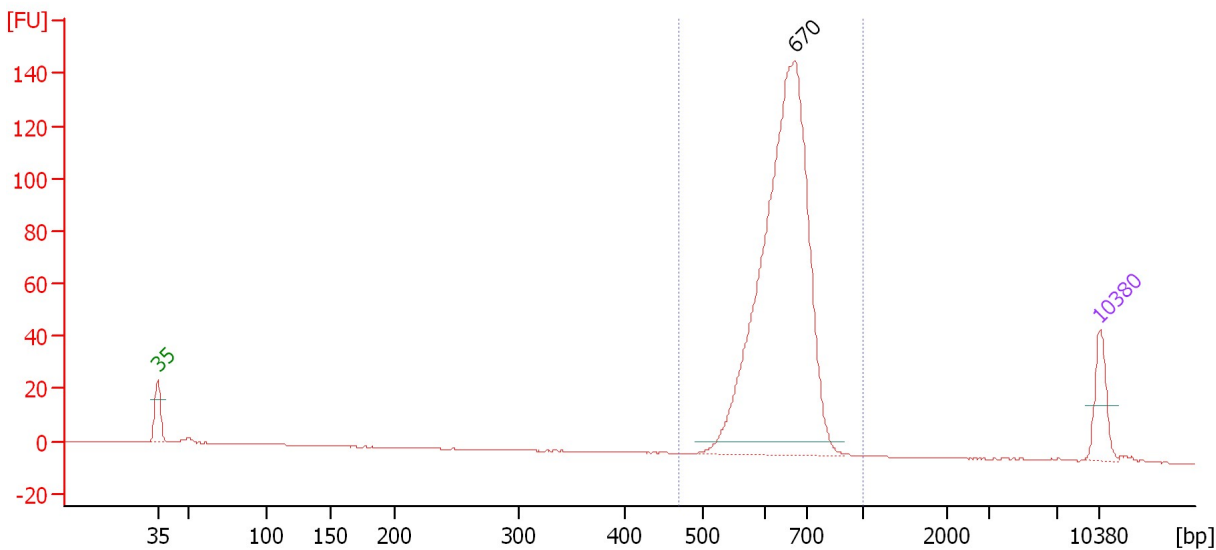
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
435	1,000	622	602.0	2,876.7	1,175.16	99	7.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 AM

Electropherogram Summary Continued ...

7: PD evo 2-5-1 Size Sel. Library



Overall Results for sample 6 : 7: PD evo 2-5-1 Size Sel. Library

Number of peaks found: 1 Corr. Area 1: 640.1
 Noise: 0.2

Peak table for sample 6 : 7: PD evo 2-5-1 Size Sel. Library

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	670	1,477.30	3,341.5		90.32
3	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 6 : 7: PD evo 2-5-1 Size Sel. Library

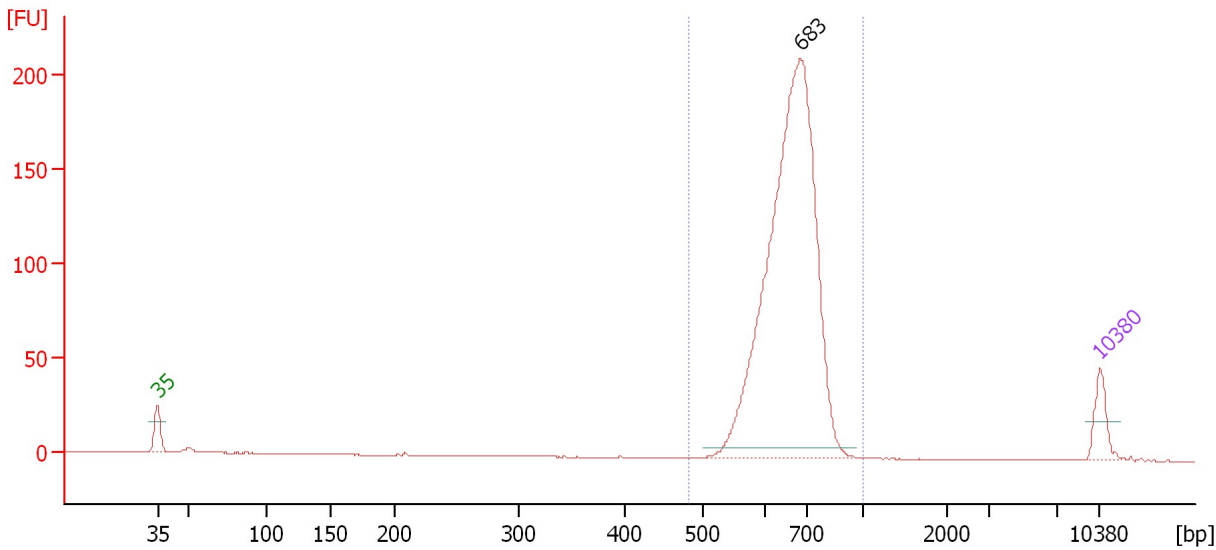
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
470	1,000	645	640.1	3,463.7	1,465.94	98	8.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 AM

Electropherogram Summary Continued ...

8: PD evo 2-8-1 Size Sel. Library



Overall Results for sample 7 : 8: PD evo 2-8-1 Size Sel. Library

Number of peaks found: 1 Corr. Area 1: 916.9
 Noise: 0.1

Peak table for sample 7 : 8: PD evo 2-8-1 Size Sel. Library

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	683	2,152.41	4,775.0		90.73
3	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 7 : 8: PD evo 2-8-1 Size Sel. Library

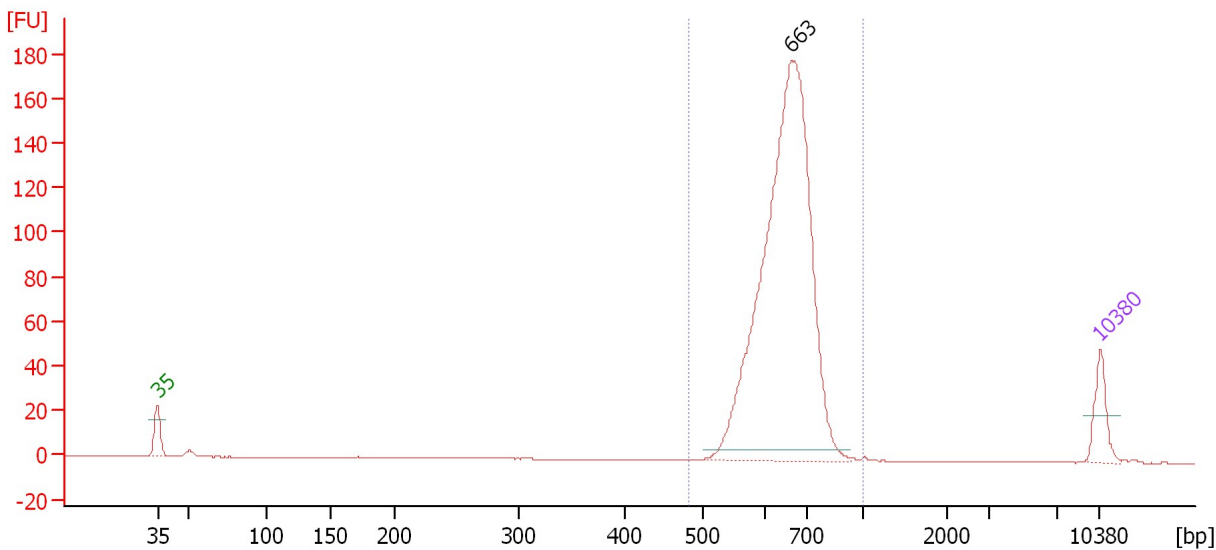
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
483	1,000	665	916.9	4,887.1	2,129.59	99	9.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 AM

Electropherogram Summary Continued ...

9: PD evo 2-11-1 Size Sel. Library



Overall Results for sample 8 : 9: PD evo 2-11-1 Size Sel. Library

Number of peaks found: 1 Corr. Area 1: 779.9
 Noise: 0.1

Peak table for sample 8 : 9: PD evo 2-11-1 Size Sel. Library

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	663	1,772.75	4,048.3		90.11
3	10,380	75.00	10.9	Upper Marker	113.00

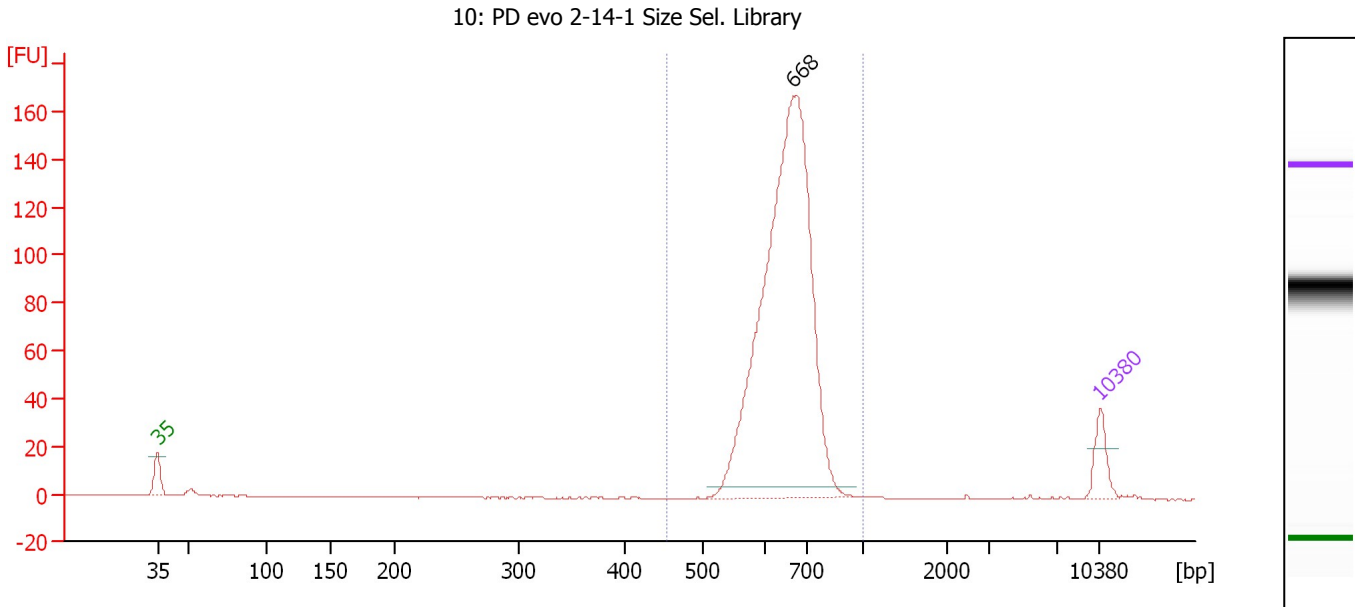
Region table for sample 8 : 9: PD evo 2-11-1 Size Sel. Library

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
482	1,000	652	779.9	4,153.2	1,775.05	99	9.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 AM

Electropherogram Summary Continued ...



Overall Results for sample 9 : 10: PD evo 2-14-1 Size Sel. Library

Number of peaks found: 1 Corr. Area 1: 729.3
 Noise: 0.1

Peak table for sample 9 : 10: PD evo 2-14-1 Size Sel. Library

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	2,180.76	4,949.0		90.25
3	10,380	75.00	10.9	Upper Marker	113.00

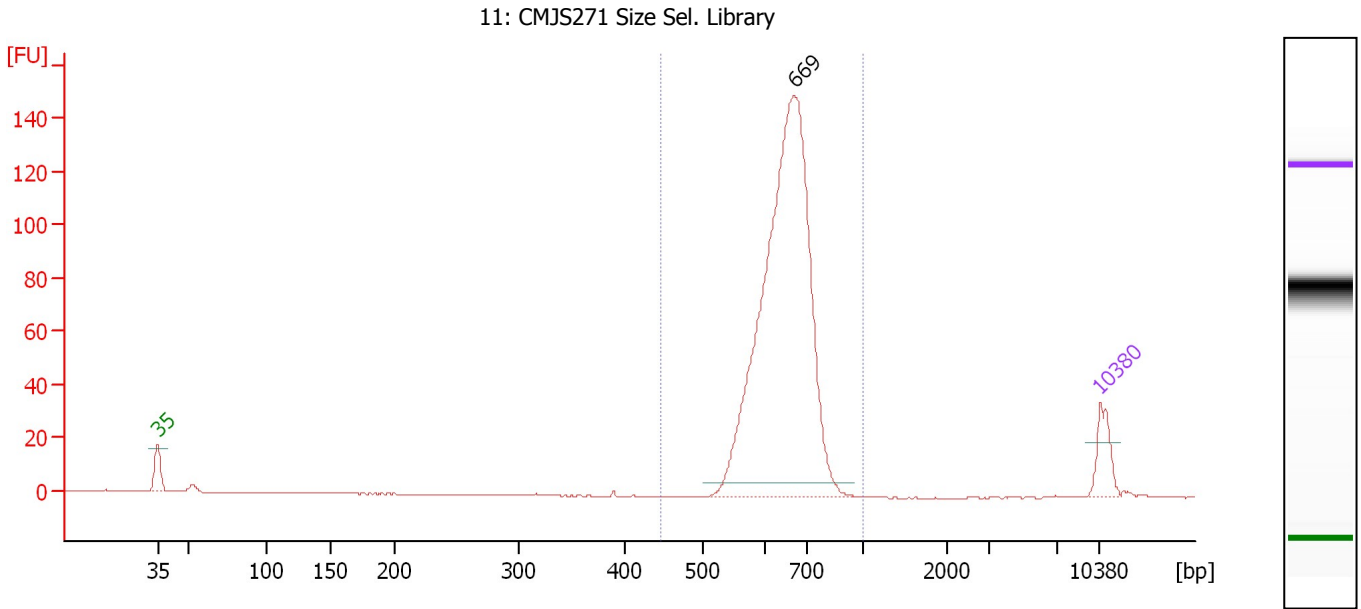
Region table for sample 9 : 10: PD evo 2-14-1 Size Sel. Library

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
454	1,000	653	729.3	5,094.3	2,179.01	99	8.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 AM

Electropherogram Summary Continued ...



Overall Results for sample 10 : 11: CMJS271 Size Sel. Library

Number of peaks found: 1 Corr. Area 1: 632.4
 Noise: 0.1

Peak table for sample 10 : 11: CMJS271 Size Sel. Library

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	669	1,862.10	4,217.5		90.29
3	10,380	75.00	10.9	Upper Marker	113.00

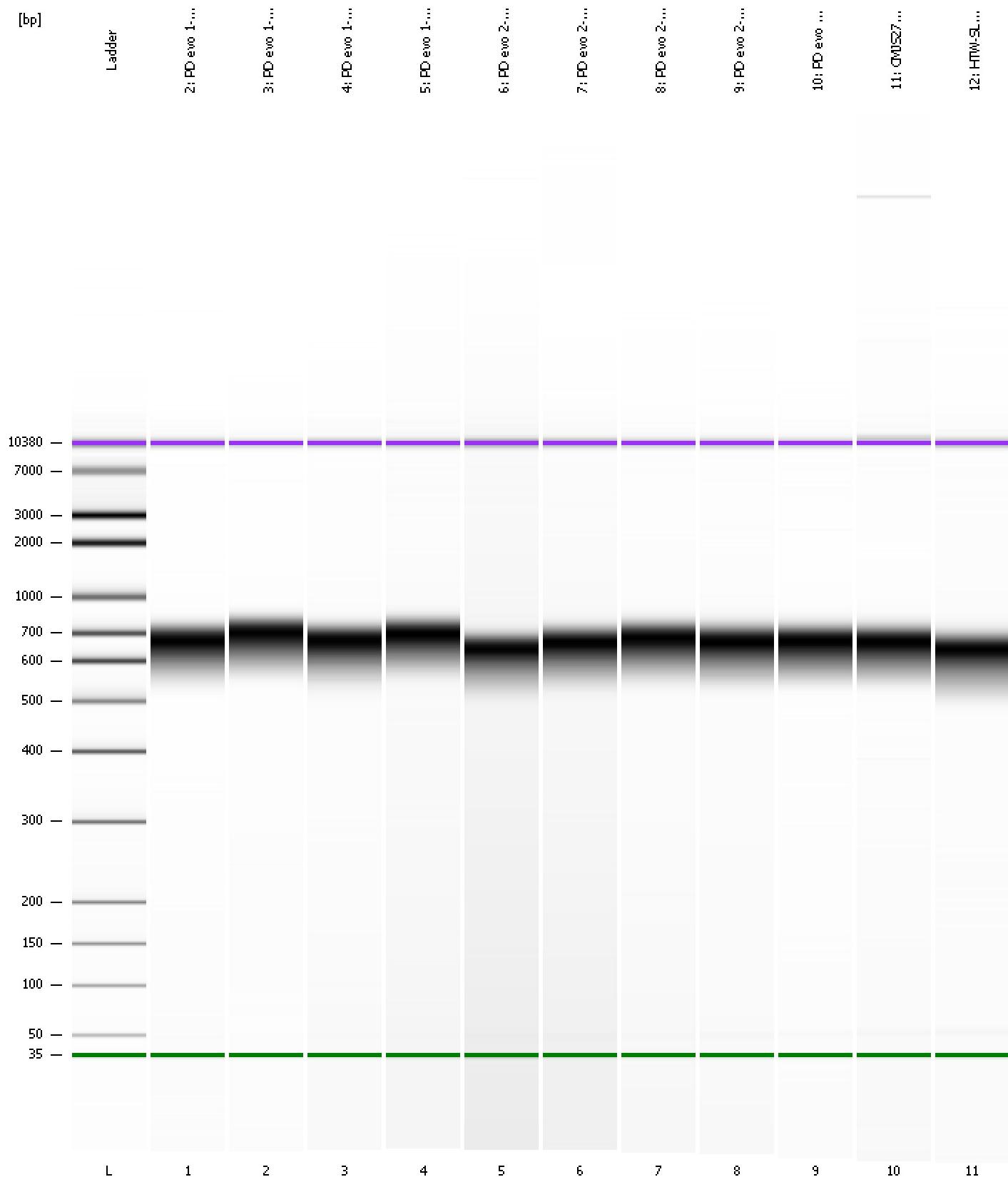
Region table for sample 10 : 11: CMJS271 Size Sel. Library

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
445	1,000	650	632.4	4,308.0	1,836.39	99	8.4

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
Modified: 1/19/2017 10:53:11 AM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-01-19\2017-01-19_001.xad

Created: 1/19/2017 10:09:20 AM
 Modified: 1/19/2017 10:53:11 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		1/19/2017 10:50:38 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\2017-01-19\2017-01-19_001.xad)		Instrument	Run		1/19/2017 10:09:25 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		1/19/2017 10:09:25 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		1/19/2017 10:09:25 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		1/19/2017 10:09:25 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		1/19/2017 10:09:25 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		1/19/2017 10:09:25 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		1/19/2017 10:09:25 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1