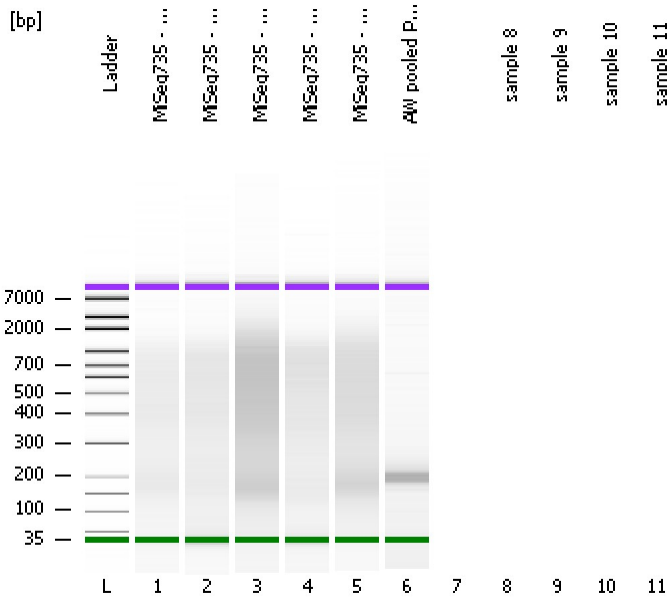


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-11-27\2018-11-27_001.xad

Created: 11/27/2018 9:40:38 AM
Modified: 11/27/2018 12:51:14 PM

Electrophoresis File Run Summary



Instrument Information:

Instrument Name: DE34903152 Firmware: C.01.069
Serial#: DE34903152 Type: G2938C

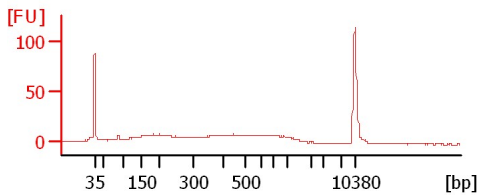
Assay Information:

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

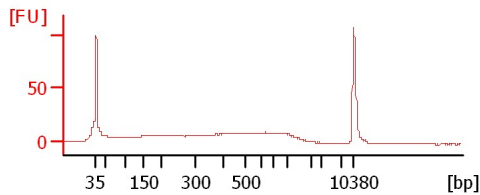
Chip Information:

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

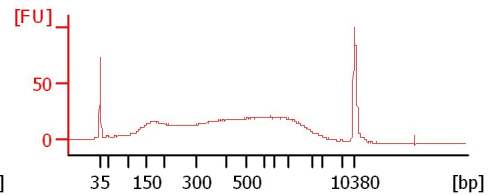
MiSeq735 - 1 soni 20 sec



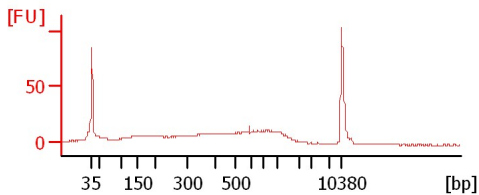
MiSeq735 - 2 soni 20 sec



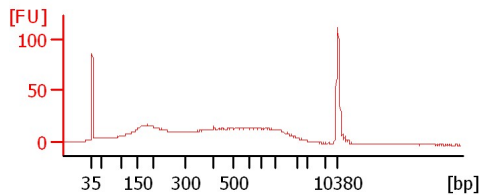
MiSeq735 - 3 soni 20 sec



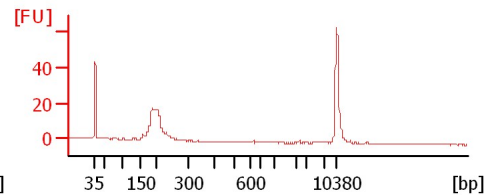
MiSeq735 - 6 soni 20 sec



MiSeq735 - 7 soni 20 sec



AW pooled Pippin 2



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-11-27\2018-11-27_001.xad

Created: 11/27/2018 9:40:38 AM
 Modified: 11/27/2018 12:51:14 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
MiSeq735 - 1 soni 20 sec		<input type="checkbox"/>	✓			
MiSeq735 - 2 soni 20 sec		<input type="checkbox"/>	✓			
MiSeq735 - 3 soni 20 sec		<input type="checkbox"/>	✓			
MiSeq735 - 6 soni 20 sec		<input type="checkbox"/>	✓			
MiSeq735 - 7 soni 20 sec		<input type="checkbox"/>	✓			
AW pooled Pippin 2		<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:\... bioanalyzer\2100 expert\data\2018-11-27\2018-11-27_001.xad

Created: 11/27/2018 9:40:38 AM
Modified: 11/27/2018 12:51:14 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

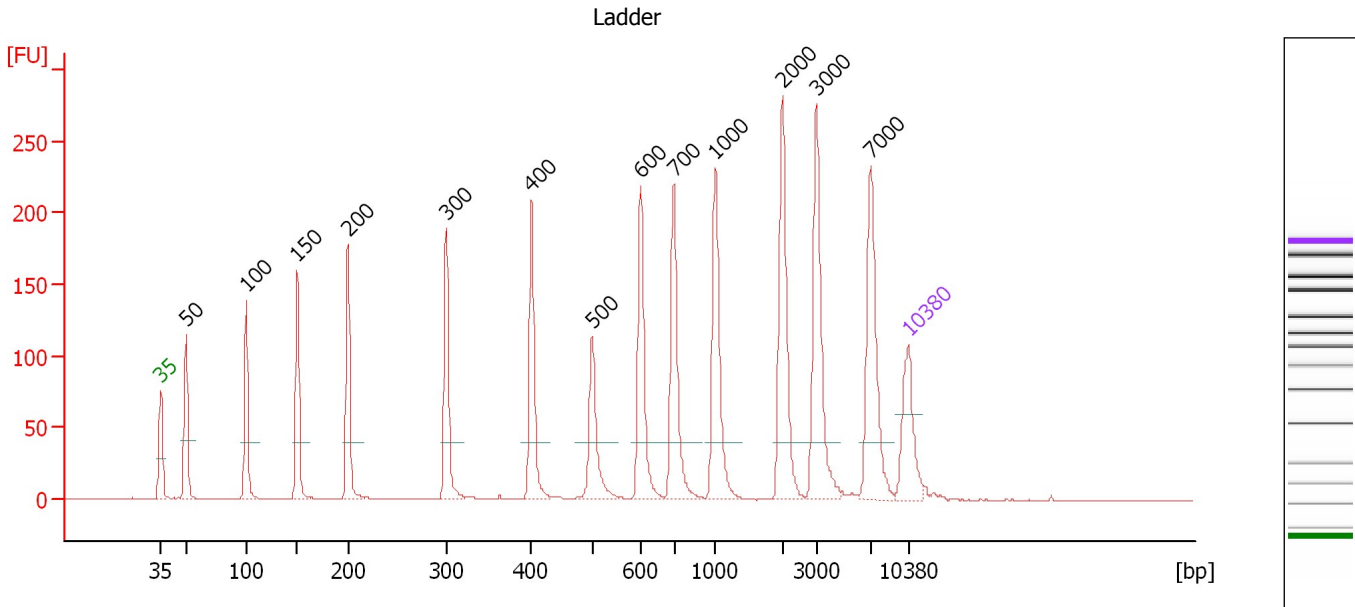
Ladder

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:\... bioanalyzer\2100 expert\data\2018-11-27\2018-11-27_001.xad

Created: 11/27/2018 9:40:38 AM
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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

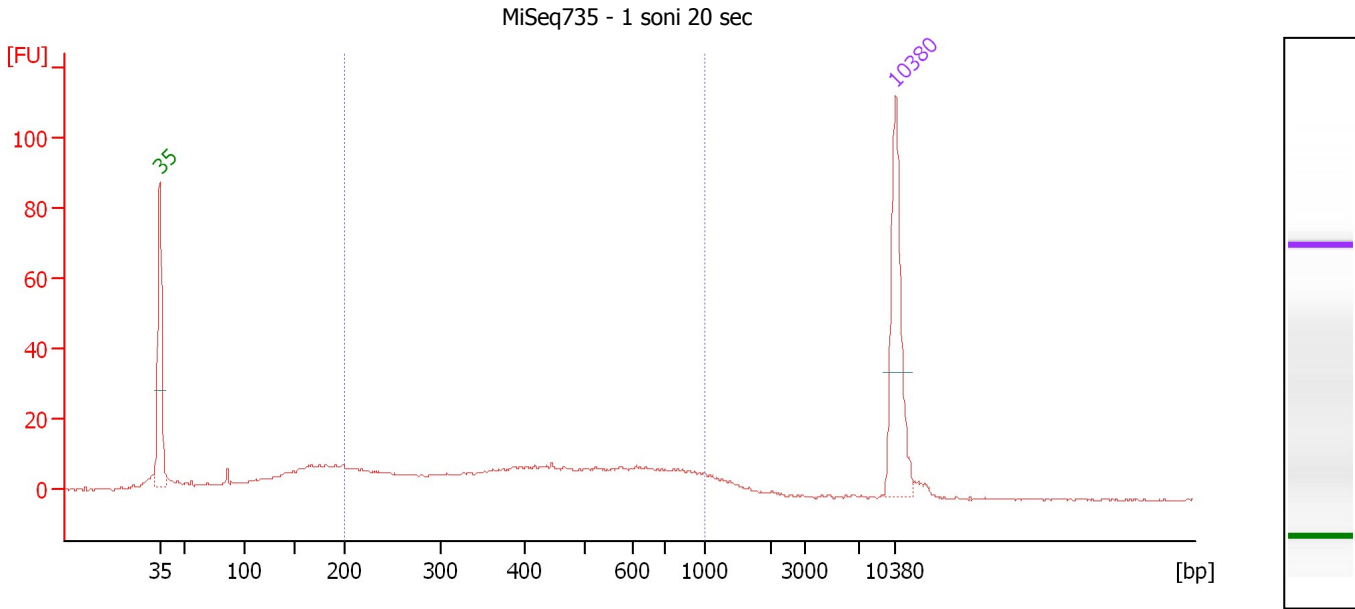
Peak table for Ladder

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	45.35
3	100	150.00	2,272.7	Ladder Peak	50.99
4	150	150.00	1,515.2	Ladder Peak	55.78
5	200	150.00	1,136.4	Ladder Peak	60.52
6	300	150.00	757.6	Ladder Peak	69.71
7	400	150.00	568.2	Ladder Peak	77.70
8	500	150.00	454.5	Ladder Peak	83.39
9	600	150.00	378.8	Ladder Peak	87.94
10	700	150.00	324.7	Ladder Peak	91.03
11	1,000	150.00	227.3	Ladder Peak	94.93
12	2,000	150.00	113.6	Ladder Peak	101.22
13	3,000	150.00	75.8	Ladder Peak	104.36
14	7,000	150.00	32.5	Ladder Peak	109.46
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-11-27\2018-11-27_001.xad

Created: 11/27/2018 9:40:38 AM
 Modified: 11/27/2018 12:51:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : MiSeq735 - 1 soni 20 sec

Number of peaks found: 0 Corr. Area 1: 296.1
 Noise: 0.2

Peak table for sample 1 : MiSeq735 - 1 soni 20 sec

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

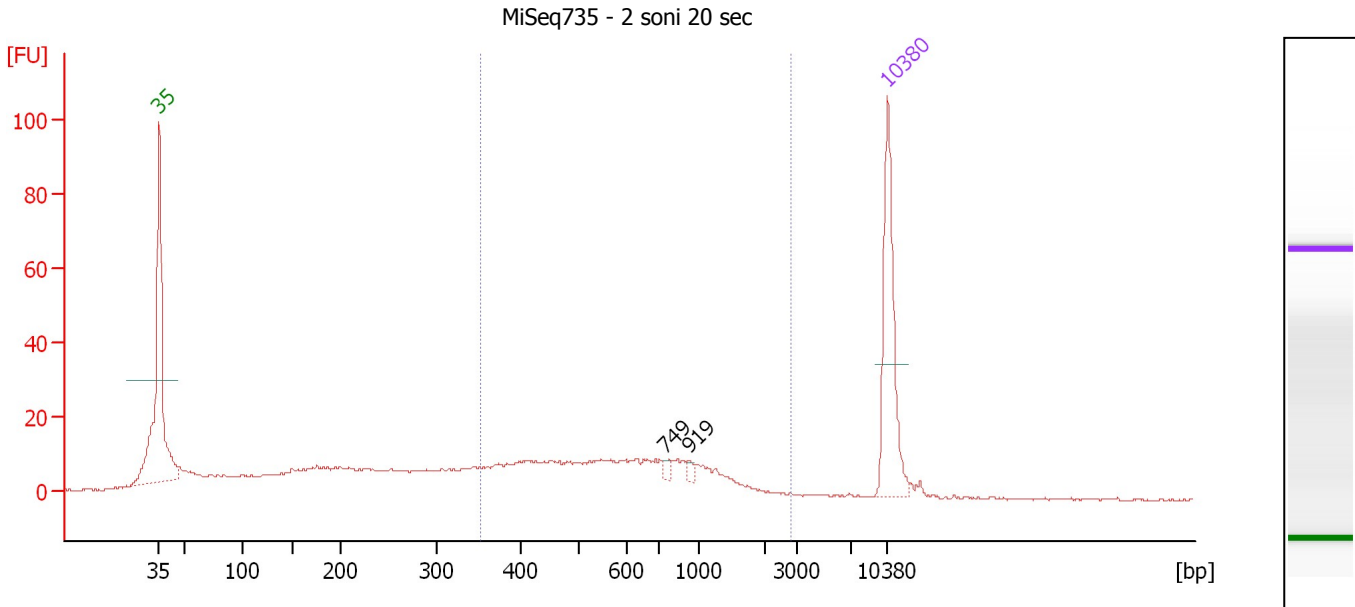
Region table for sample 1 : MiSeq735 - 1 soni 20 sec

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	463	289.66	296.1	1,209.1	63	41.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-11-27\2018-11-27_001.xad

Created: 11/27/2018 9:40:38 AM
 Modified: 11/27/2018 12:51:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : MiSeq735 - 2 soni 20 sec

Number of peaks found: 2 Corr. Area 1: 255.6
 Noise: 0.2

Peak table for sample 2 : MiSeq735 - 2 soni 20 sec

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	749	4.25	8.6		91.67
3	919	3.98	6.6		93.87
4	10,380	75.00	10.9	Upper Marker	113.00

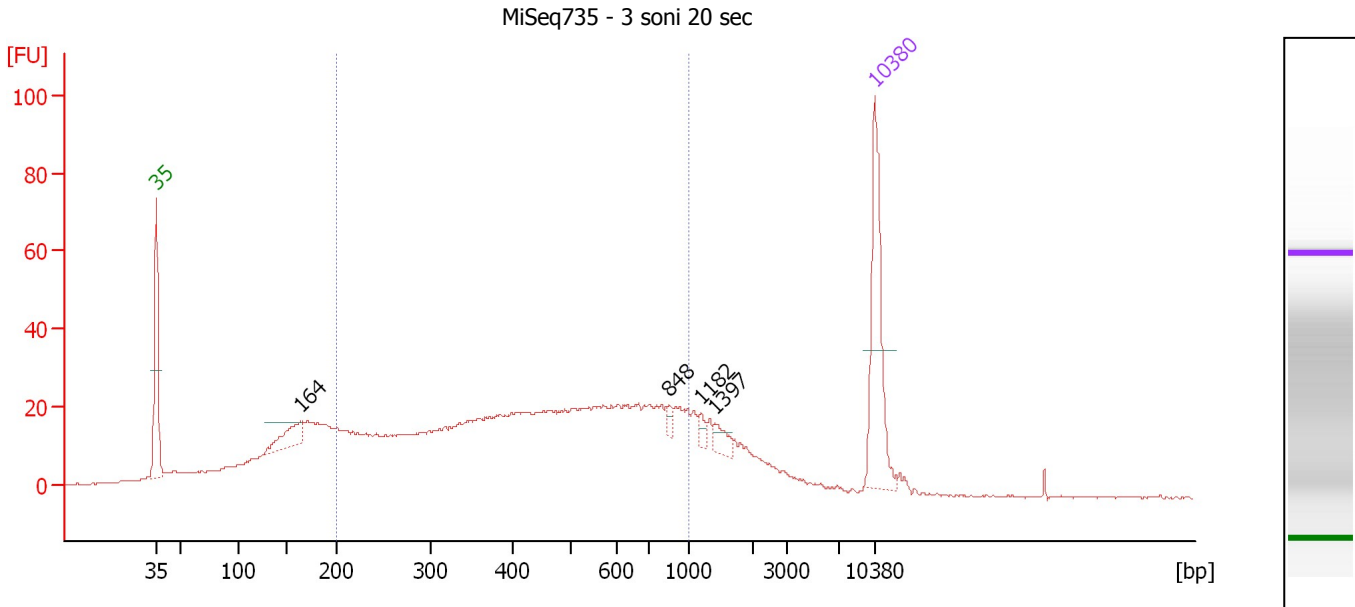
Region table for sample 2 : MiSeq735 - 2 soni 20 sec

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
352	2,782	691	239.11	255.6	661.3	46	53.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-11-27\2018-11-27_001.xad

Created: 11/27/2018 9:40:38 AM
 Modified: 11/27/2018 12:51:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : MiSeq735 - 3 soni 20 sec

Number of peaks found: 4 Corr. Area 1: 807.7
 Noise: 0.2

Peak table for sample 3 : MiSeq735 - 3 soni 20 sec

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	164	33.99	314.8		57.07
3	848	5.50	9.8		92.96
4	1,182	5.69	7.3		96.07
5	1,397	10.12	11.0		97.42
6	10,380	75.00	10.9	Upper Marker	113.00

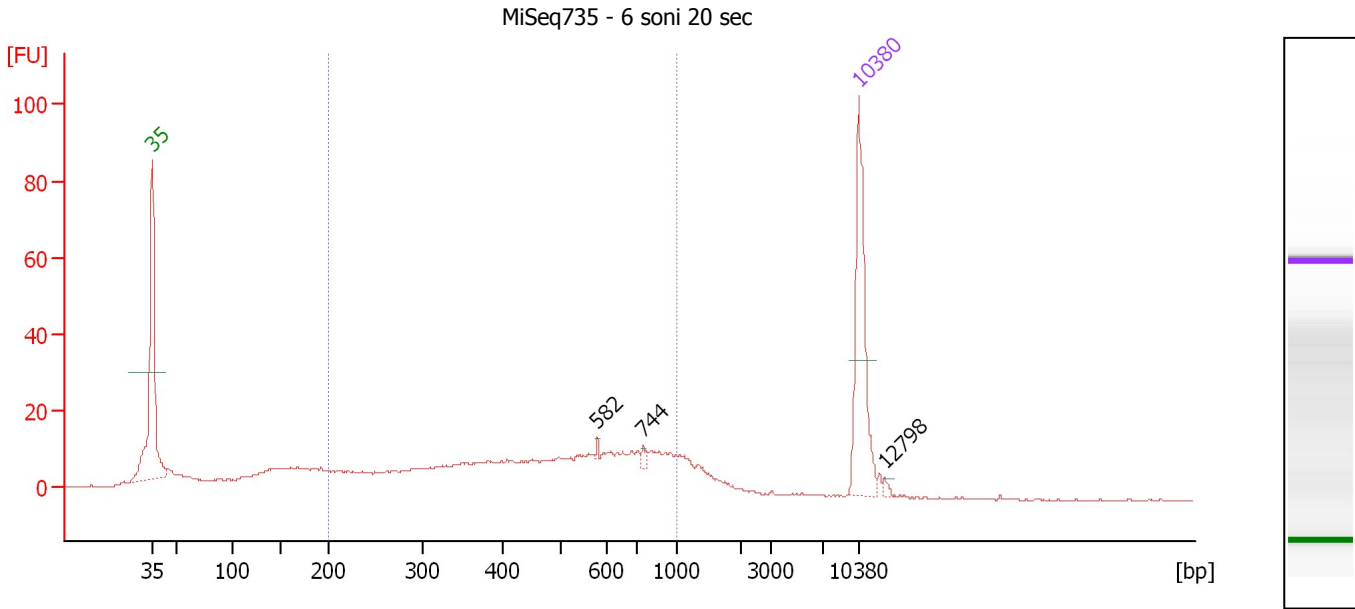
Region table for sample 3 : MiSeq735 - 3 soni 20 sec

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	477	915.10	807.7	3,701.2	65	41.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-11-27\2018-11-27_001.xad

Created: 11/27/2018 9:40:38 AM
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Electropherogram Summary Continued ...



Overall Results for sample 4 : MiSeq735 - 6 soni 20 sec

Number of peaks found: 3 Corr. Area 1: 342.8
 Noise: 0.2

Peak table for sample 4 : MiSeq735 - 6 soni 20 sec

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	582	1.77	4.6		87.11
3	744	3.72	7.6		91.60
4	10,380	75.00	10.9	Upper Marker	113.00
5	12,798	0.00	0.0		115.54

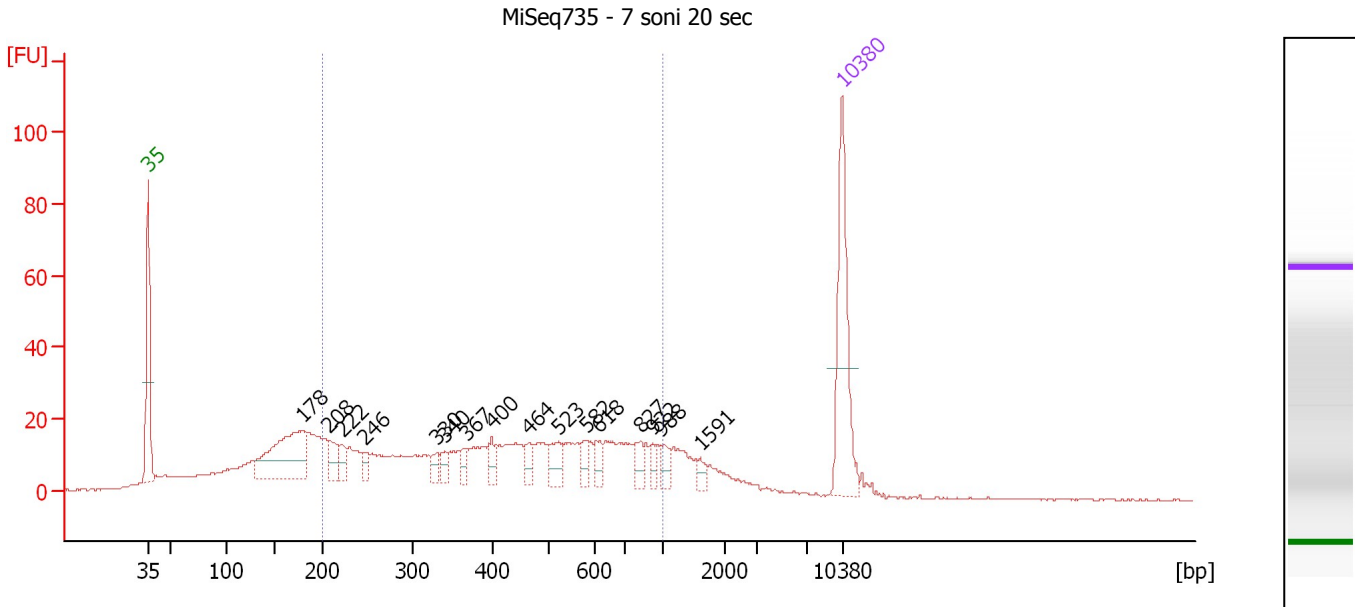
Region table for sample 4 : MiSeq735 - 6 soni 20 sec

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	498	383.86	342.8	1,482.5	66	39.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-11-27\2018-11-27_001.xad

Created: 11/27/2018 9:40:38 AM
 Modified: 11/27/2018 12:51:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : MiSeq735 - 7 soni 20 sec

Number of peaks found: 16 Corr. Area 1: 587.3
 Noise: 0.1

Peak table for sample 5 : MiSeq735 - 7 soni 20 sec

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	178	112.06	951.3		58.48
3	208	18.83	137.0		61.28
4	222	13.89	94.9		62.52
5	246	9.79	60.2		64.77
6	330	10.70	49.2		72.09
7	340	9.31	41.5		72.89
8	367	9.84	40.7		75.04
9	400	11.03	41.8		77.68
10	464	10.40	34.0		81.33
11	523	17.33	50.2		84.45
12	582	10.51	27.3		87.14
13	618	11.59	28.4		88.48
14	827	11.54	21.1		92.68
15	922	7.61	12.5		93.91
16	988	9.83	15.1		94.77
17	1,591	5.85	5.6		98.65
18	10,380	75.00	10.9	Upper Marker	113.00

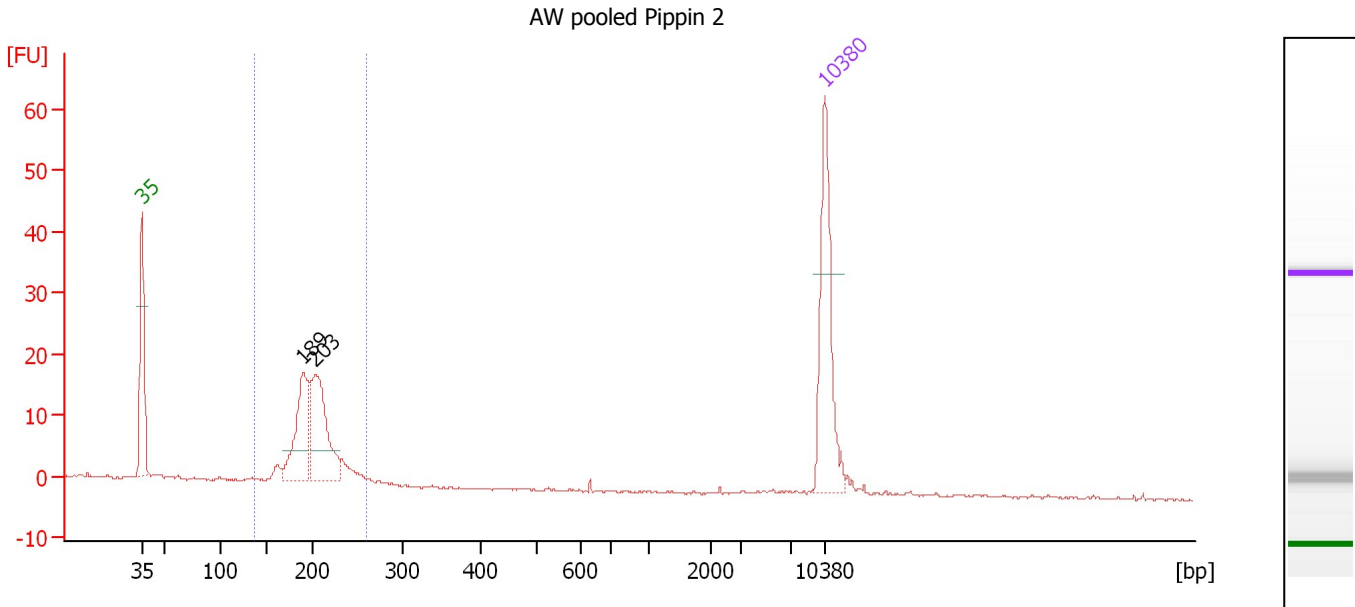
Region table for sample 5 : MiSeq735 - 7 soni 20 sec

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	464	590.96	587.3	2,482.1	60	42.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-11-27\2018-11-27_001.xad

Created: 11/27/2018 9:40:38 AM
 Modified: 11/27/2018 12:51:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : AW pooled Pippin 2

Number of peaks found: 2 Corr. Area 1: 113.5
 Noise: 0.1

Peak table for sample 6 : AW pooled Pippin 2

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	189	85.40	685.9		59.45
3	203	102.01	763.1		60.76
4	10,380	75.00	10.9	Upper Marker	113.00

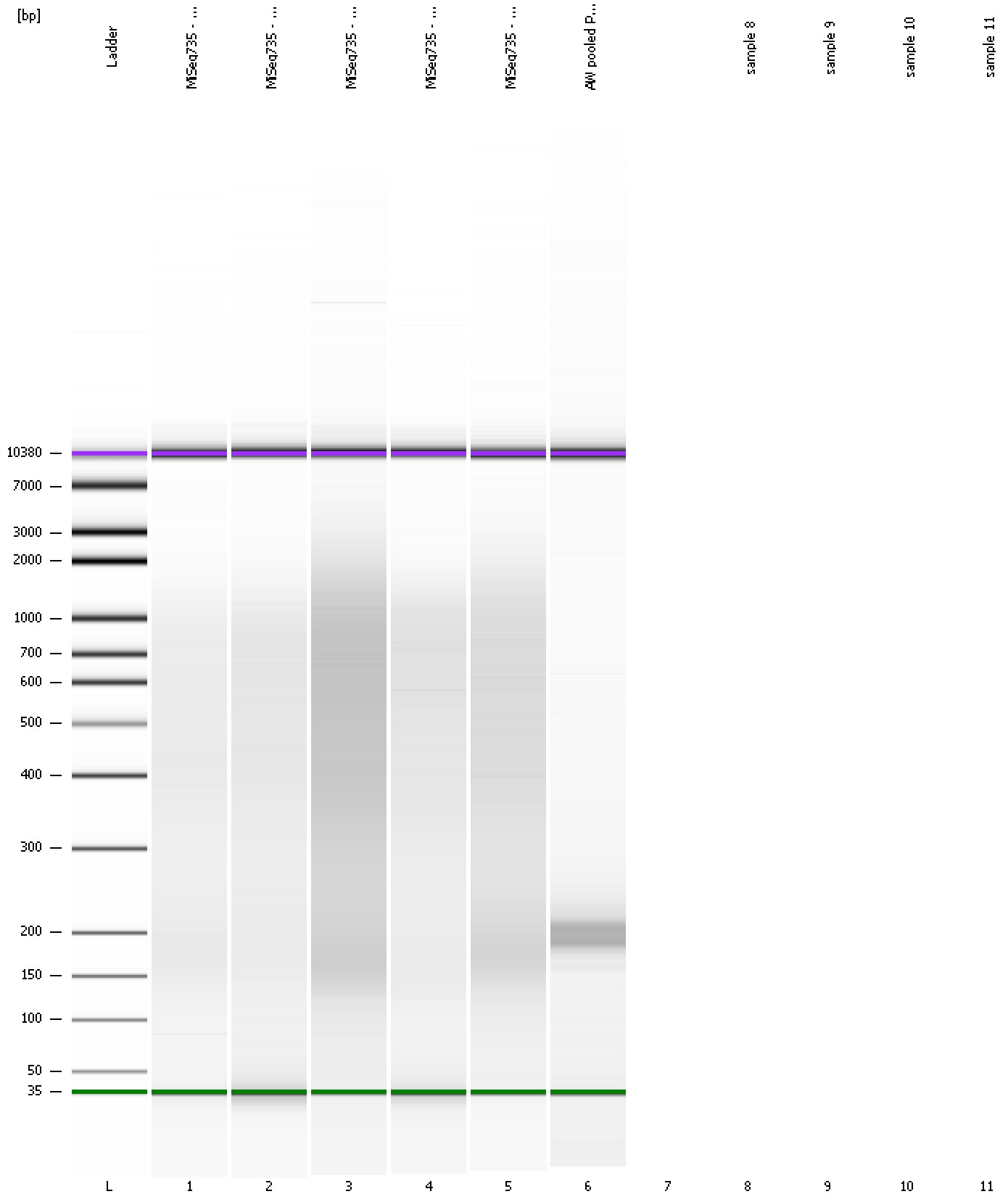
Region table for sample 6 : AW pooled Pippin 2

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
136	261	201	213.29	113.5	1,624.1	94	9.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-11-27\2018-11-27_001.xad

Created: 11/27/2018 9:40:38 AM
Modified: 11/27/2018 12:51:14 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-11-27\2018-11-27_001.xad

Created: 11/27/2018 9:40:38 AM
Modified: 11/27/2018 12:51:14 PM

Invalid Samples

Sample 7 has not been run, no results available.

Sample 8 has not been run, no results available.

Sample 9 has not been run, no results available.

Sample 10 has not been run, no results available.

Sample 11 has not been run, no results available.