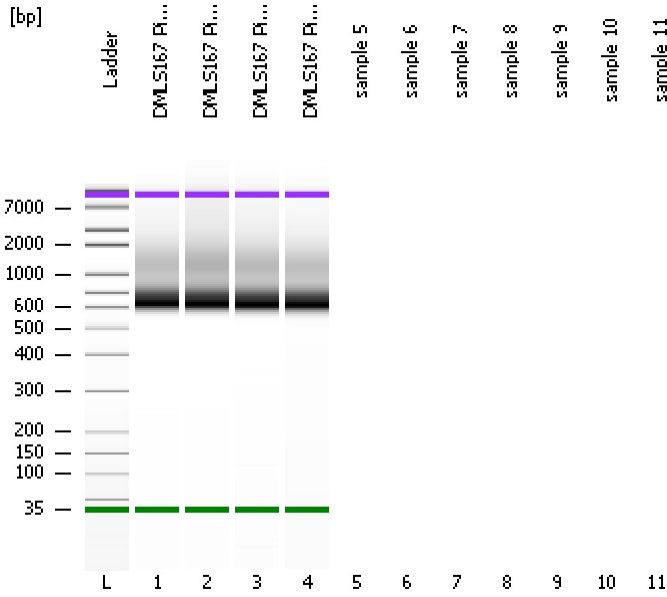


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-17\2018-10-17_002.xad

Created: 10/17/2018 1:06:27 PM
Modified: 10/17/2018 1:27:16 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 #:

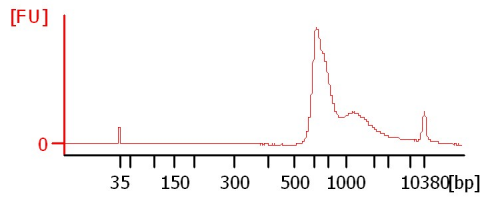
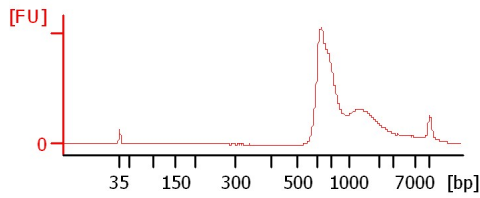
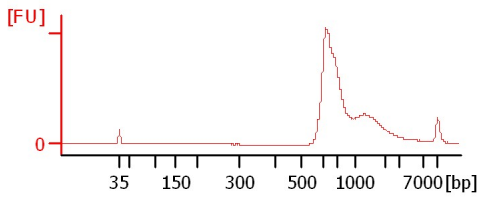
Reagent Kit Lot #:

Chip Comments:

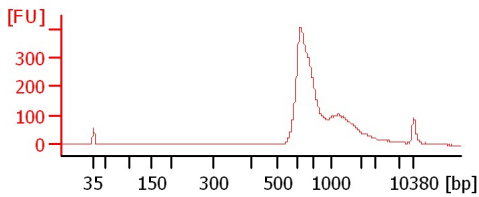
DMLS167 Pippin 1

DMLS167 Pippin 2

DMLS167 Pippin 3



DMLS167 Pippin 4



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-17\2018-10-17_002.xad

Created: 10/17/2018 1:06:27 PM
 Modified: 10/17/2018 1:27:16 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
DMLS167 Pippin 1		<input type="checkbox"/>	✓			
DMLS167 Pippin 2		<input type="checkbox"/>	✓			
DMLS167 Pippin 3		<input type="checkbox"/>	✓			
DMLS167 Pippin 4		<input type="checkbox"/>	✓			
sample 5		<input type="checkbox"/>				
sample 6		<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:\... bioanalyzer\2100 expert\data\2018-10-17\2018-10-17_002.xad

Created: 10/17/2018 1:06:27 PM
Modified: 10/17/2018 1:27:16 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

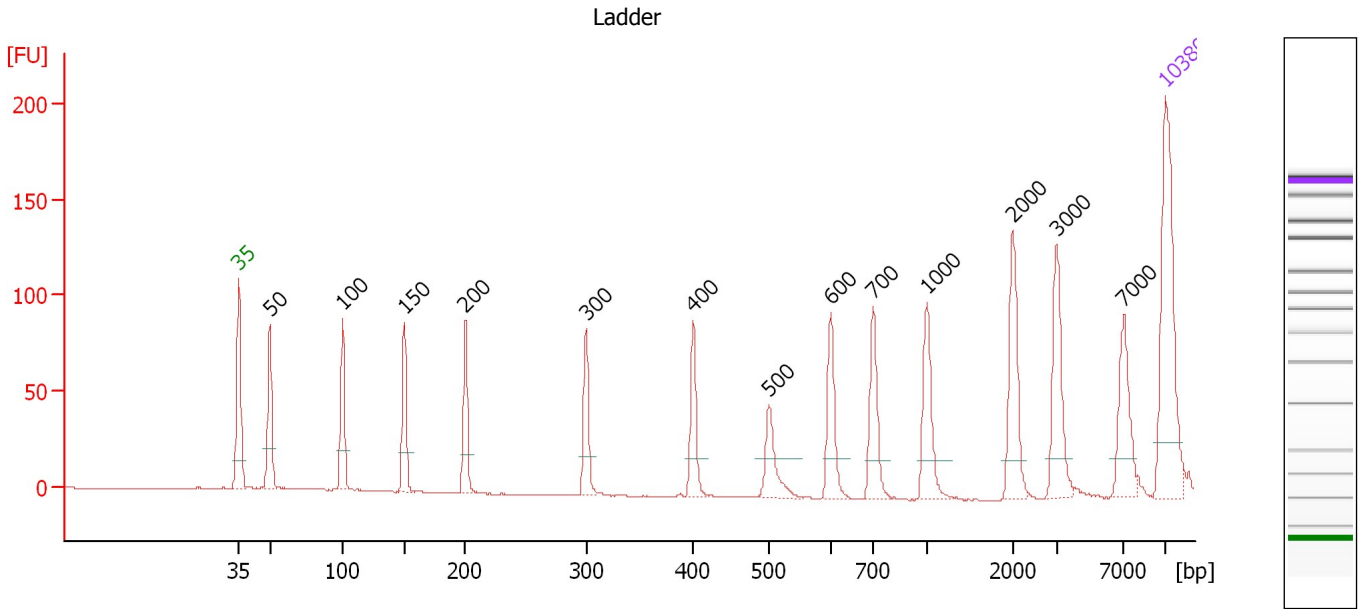
Ladder

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\2018-10-17\2018-10-17_002.xad

Created: 10/17/2018 1:06:27 PM
 Modified: 10/17/2018 1:27:16 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.3

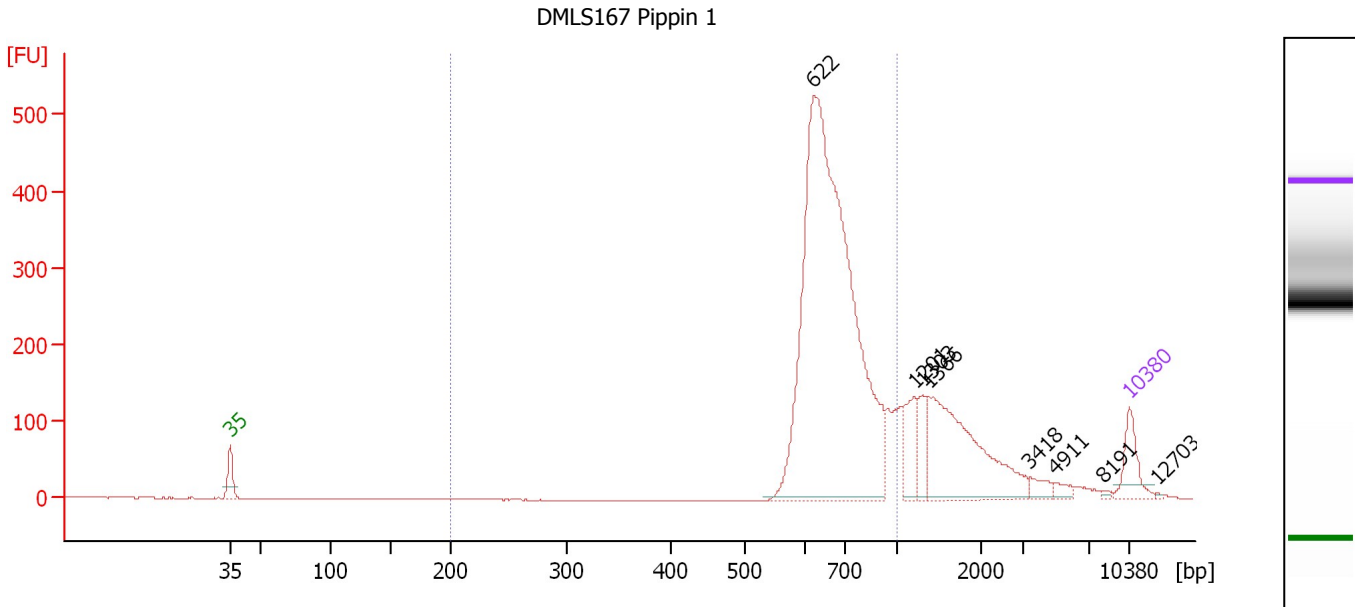
Peak table for Ladder

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	50.85
4	150	150.00	1,515.2	Ladder Peak	55.53
5	200	150.00	1,136.4	Ladder Peak	60.12
6	300	150.00	757.6	Ladder Peak	69.26
7	400	150.00	568.2	Ladder Peak	77.32
8	500	150.00	454.5	Ladder Peak	83.03
9	600	150.00	378.8	Ladder Peak	87.71
10	700	150.00	324.7	Ladder Peak	90.93
11	1,000	150.00	227.3	Ladder Peak	94.96
12	2,000	150.00	113.6	Ladder Peak	101.48
13	3,000	150.00	75.8	Ladder Peak	104.74
14	7,000	150.00	32.5	Ladder Peak	109.86
15	10,388	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-17\2018-10-17_002.xad

Created: 10/17/2018 1:06:27 PM
 Modified: 10/17/2018 1:27:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : DMLS167 Pippin 1

Number of peaks found: 8 Corr. Area 1: 2,725.2
 Noise: 0.3

Peak table for sample 1 : DMLS167 Pippin 1

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	622	2,034.12	4,953.6		88.42
3	1,201	108.69	137.1		96.27
4	1,303	67.86	78.9		96.93
5	1,366	441.93	490.0		97.35
6	3,418	27.97	12.4		105.28
7	4,911	17.97	5.5		107.19
8	8,191	5.76	1.1		110.97
9	10,380	75.00	10.9	Upper Marker	113.00
10	12,703	0.00	0.0		115.16

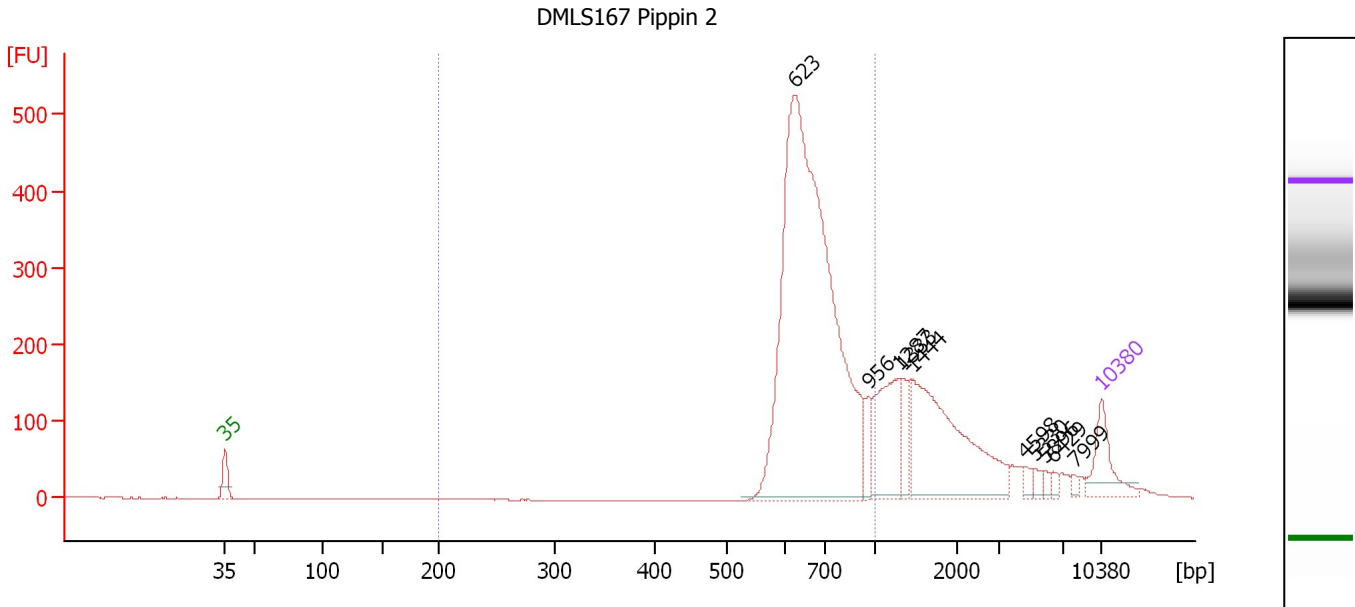
Region table for sample 1 : DMLS167 Pippin 1

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	692	2,073.72	2,725.2	4,619.7	72	14.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-17\2018-10-17_002.xad

Created: 10/17/2018 1:06:27 PM
 Modified: 10/17/2018 1:27:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : DMLS167 Pippin 2

Number of peaks found: 10 Corr. Area 1: 2,797.3
 Noise: 0.2

Peak table for sample 2 : DMLS167 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	623	1,517.00	3,689.5		88.45
3	956	47.53	75.4		94.36
4	1,287	177.68	209.2		96.83
5	1,333	51.21	58.2		97.13
6	1,444	372.79	391.2		97.85
7	4,598	15.39	5.1		106.79
8	5,330	10.89	3.1		107.72
9	5,896	9.62	2.5		108.45
10	6,429	8.55	2.0		109.13
11	7,999	6.58	1.2		110.79
12	10,380	75.00	10.9	Upper Marker	113.00

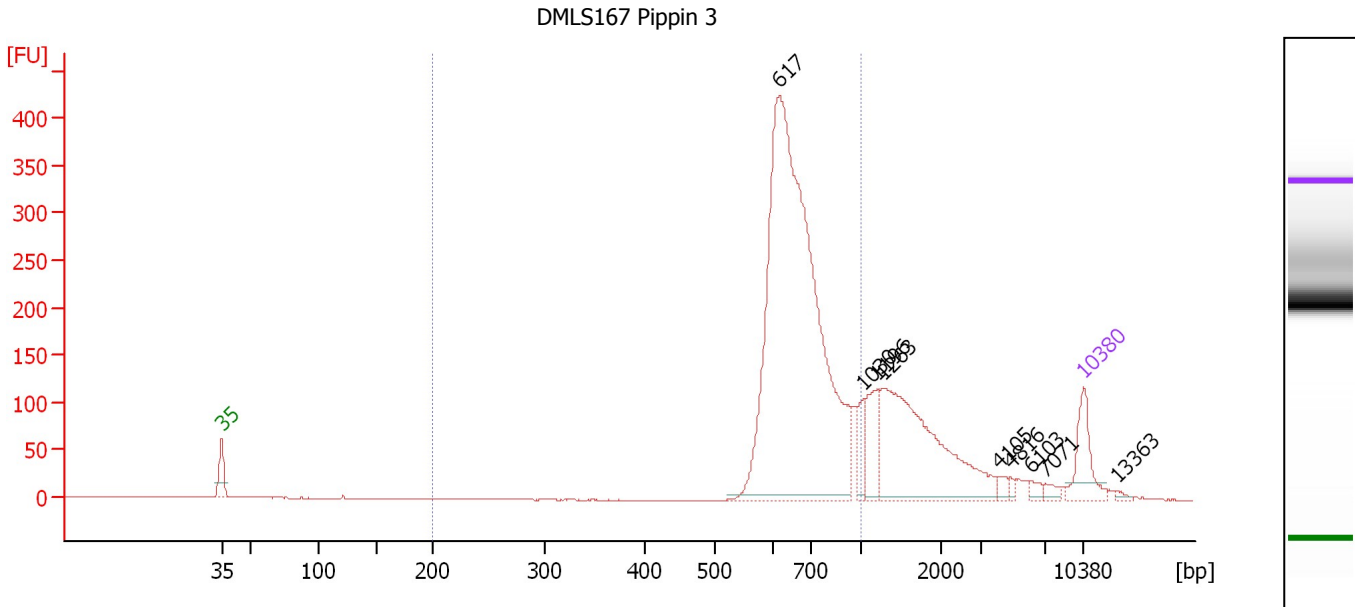
Region table for sample 2 : DMLS167 Pippin 2

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	693	1,546.30	2,797.3	3,444.8	67	14.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-17\2018-10-17_002.xad

Created: 10/17/2018 1:06:27 PM
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Electropherogram Summary Continued ...



Overall Results for sample 3 : DMLS167 Pippin 3

Number of peaks found: 9 Corr. Area 1: 2,238.9
 Noise: 0.2

Peak table for sample 3 : DMLS167 Pippin 3

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	617	1,538.80	3,776.5		88.27
3	1,030	52.14	76.7		95.15
4	1,196	84.43	106.9		96.24
5	1,263	428.60	514.3		96.67
6	4,105	11.28	4.2		106.16
7	4,816	7.20	2.3		107.07
8	6,103	11.14	2.8		108.71
9	7,071	12.45	2.7		109.92
10	10,380	75.00	10.9	Upper Marker	113.00
11	13,363	0.00	0.0		115.77

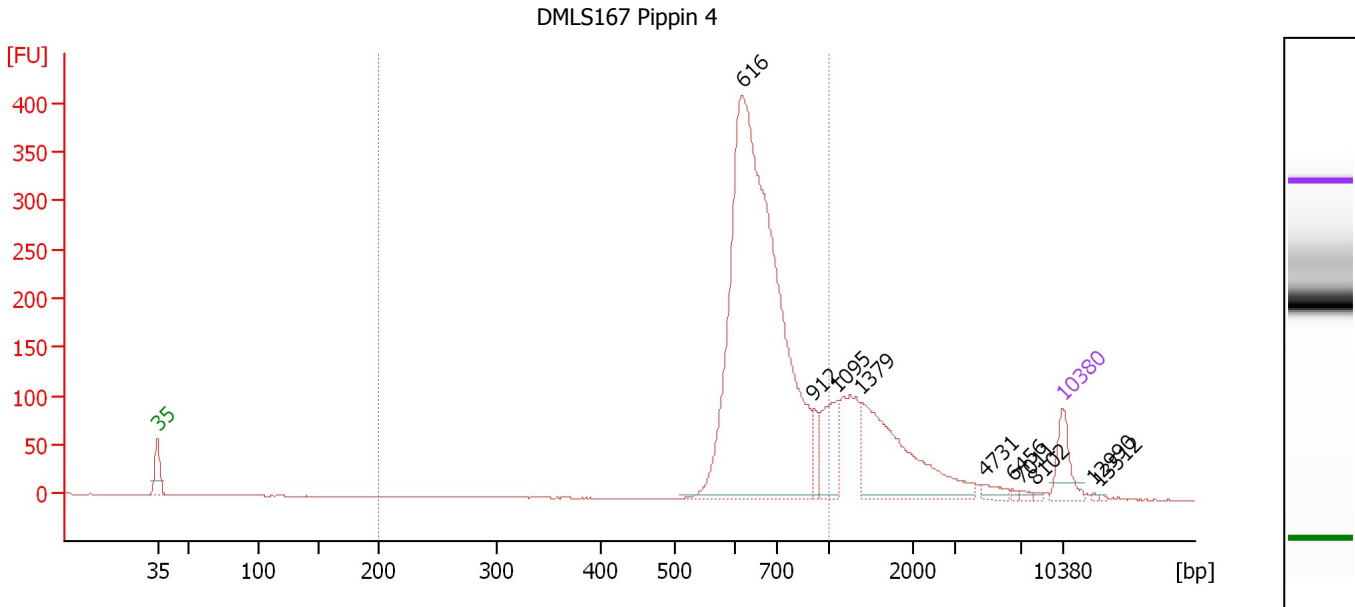
Region table for sample 3 : DMLS167 Pippin 3

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	686	1,564.60	2,238.9	3,519.2	70	14.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-17\2018-10-17_002.xad

Created: 10/17/2018 1:06:27 PM
 Modified: 10/17/2018 1:27:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : DMLS167 Pippin 4

Number of peaks found: 10 Corr. Area 1: 2,126.1
 Noise: 0.2

Peak table for sample 4 : DMLS167 Pippin 4

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	616	2,039.32	5,018.8		88.21
3	912	53.58	89.1		93.77
4	1,095	142.80	197.6		95.58
5	1,379	391.37	430.0		97.43
6	4,731	22.08	7.1		106.96
7	6,456	4.74	1.1		109.16
8	7,011	6.46	1.4		109.87
9	8,102	4.58	0.9		110.88
10	10,380	75.00	10.9	Upper Marker	113.00
11	12,990	0.00	0.0		115.43
12	13,512	0.00	0.0		115.91

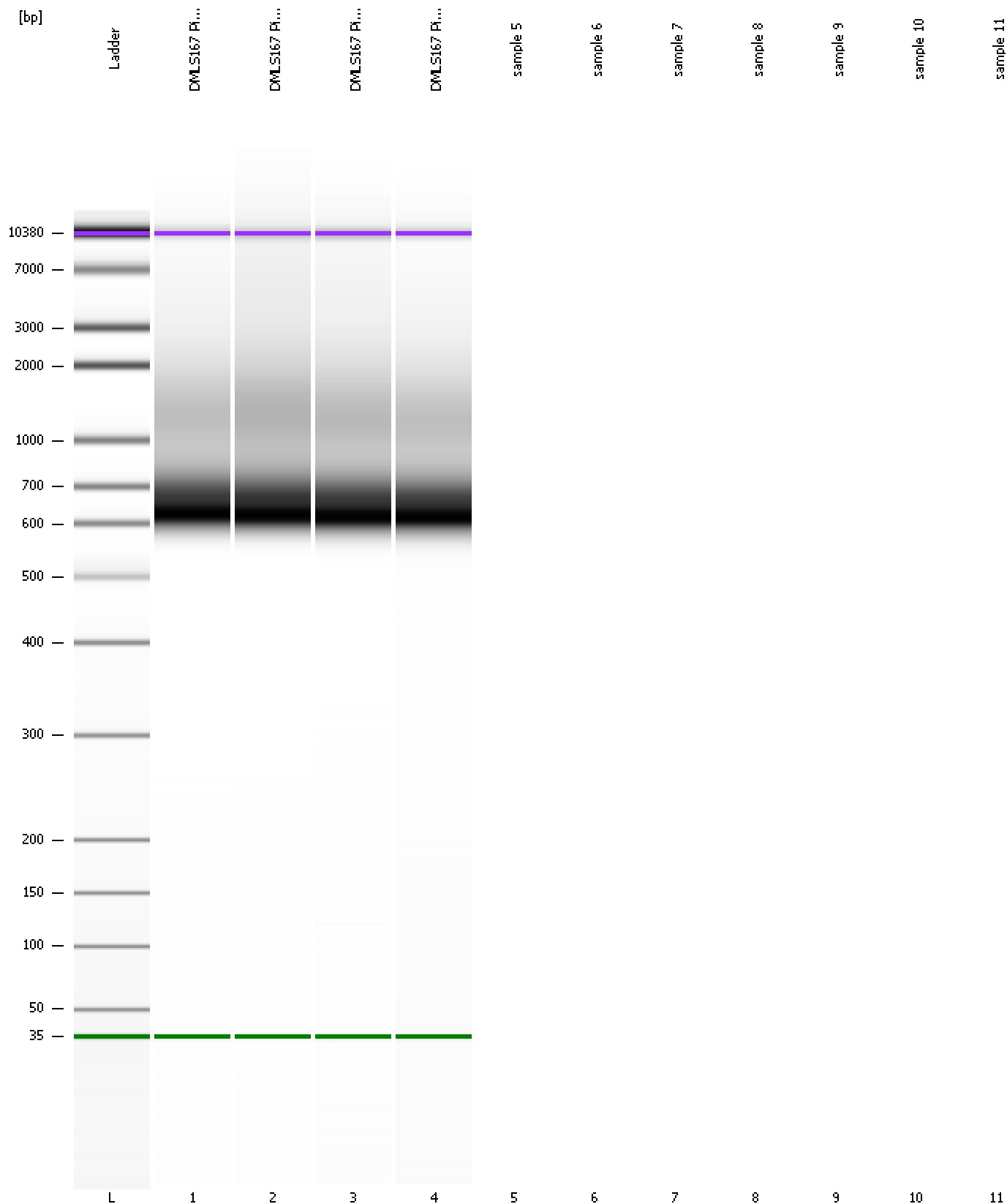
Region table for sample 4 : DMLS167 Pippin 4

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	683	2,143.69	2,126.1	4,845.5	73	14.7

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-17\2018-10-17_002.xad

Created: 10/17/2018 1:06:27 PM
Modified: 10/17/2018 1:27:16 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-17\2018-10-17_002.xad

Created: 10/17/2018 1:06:27 PM
Modified: 10/17/2018 1:27:16 PM

Invalid Samples

Sample 5 has not been run, no results available.

Sample 6 has not been run, no results available.

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.