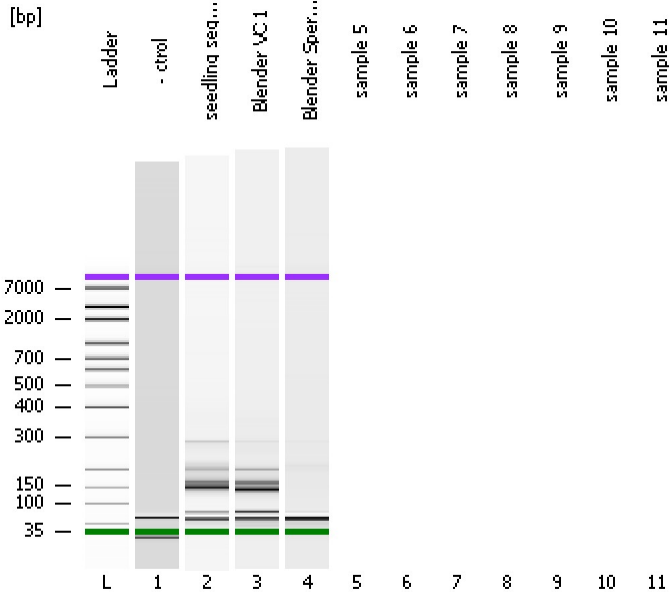


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-10-06\2017-10-06_004.xad

Created: 10/6/2017 3:32:07 PM
Modified: 10/6/2017 3:53:29 PM

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

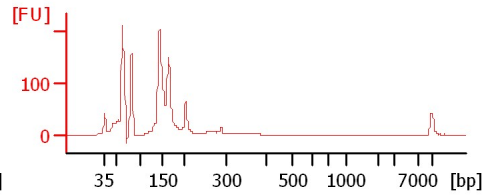
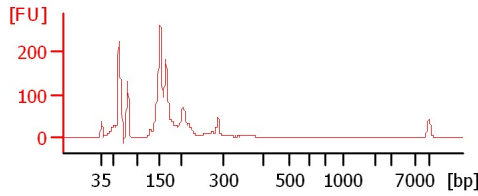
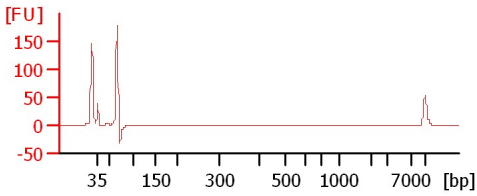
Chip Information:

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

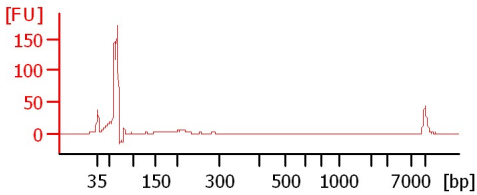
- ctrl

seedling segment 4

Blender VC 1



Blender Sperm 2



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-10-06\2017-10-06_004.xad

Created: 10/6/2017 3:32:07 PM
Modified: 10/6/2017 3:53:29 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
- ctrol		<input type="checkbox"/>	✓			
seedling segment 4		<input type="checkbox"/>	✓			
Blender VC 1		<input type="checkbox"/>	✓			
Blender Sperm 2		<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\2017-10-06\2017-10-06_004.xad

Created: 10/6/2017 3:32:07 PM
Modified: 10/6/2017 3:53:29 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

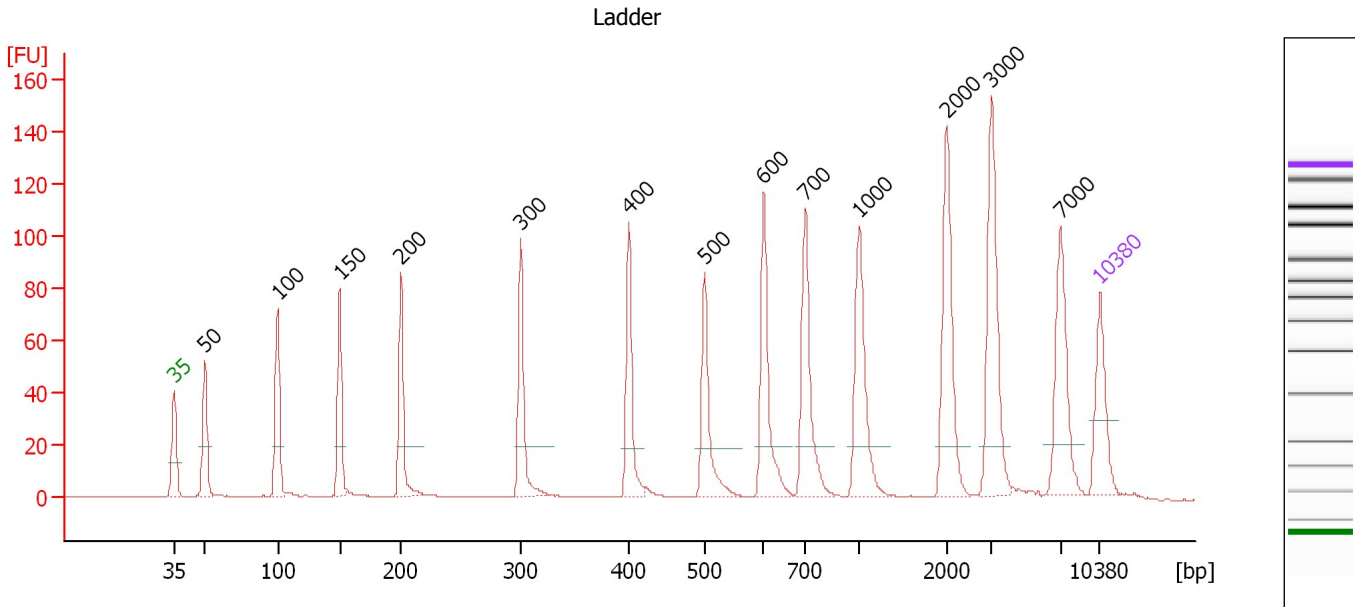
Ladder

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\2017-10-06\2017-10-06_004.xad

Created: 10/6/2017 3:32:07 PM
 Modified: 10/6/2017 3:53:29 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

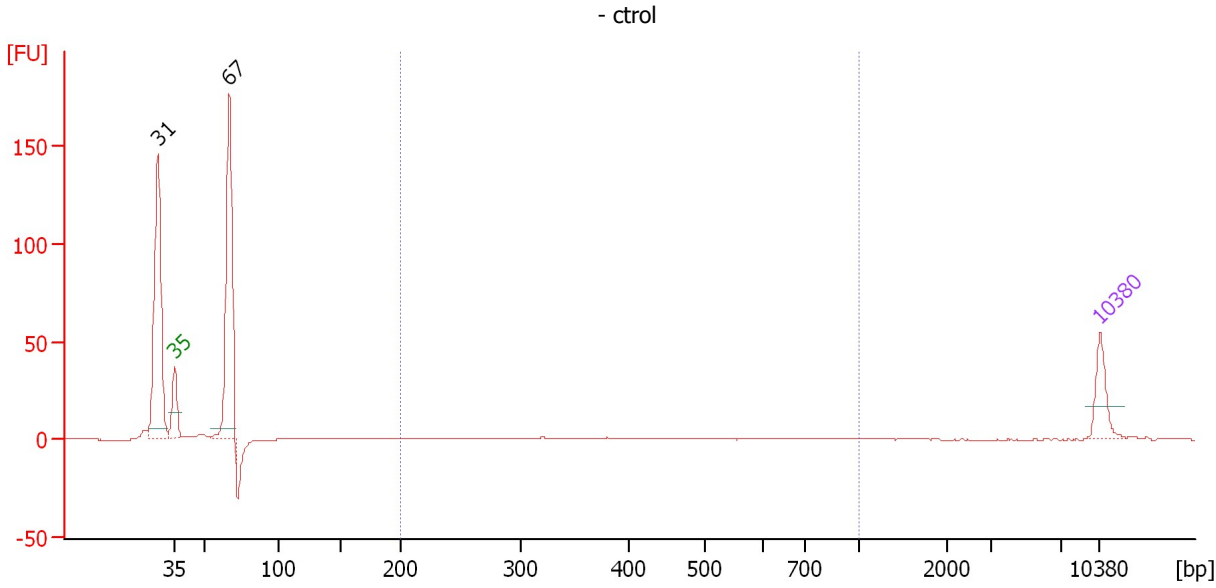
Peak table for Ladder

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	45.33
3	100	150.00	2,272.7	Ladder Peak	50.81
4	150	150.00	1,515.2	Ladder Peak	55.51
5	200	150.00	1,136.4	Ladder Peak	60.16
6	300	150.00	757.6	Ladder Peak	69.21
7	400	150.00	568.2	Ladder Peak	77.38
8	500	150.00	454.5	Ladder Peak	83.12
9	600	150.00	378.8	Ladder Peak	87.56
10	700	150.00	324.7	Ladder Peak	90.72
11	1,000	150.00	227.3	Ladder Peak	94.75
12	2,000	150.00	113.6	Ladder Peak	101.42
13	3,000	150.00	75.8	Ladder Peak	104.83
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:\... bioanalyzer\2100 expert\data\2017-10-06\2017-10-06_004.xad

Created: 10/6/2017 3:32:07 PM
 Modified: 10/6/2017 3:53:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : - ctrol

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

Peak table for sample 1 : - ctrol

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.73
2	35	125.00	5,411.3	Lower Marker	43.00
3	67	701.87	15,927.6		47.16
4	10,380	75.00	10.9	Upper Marker	113.00

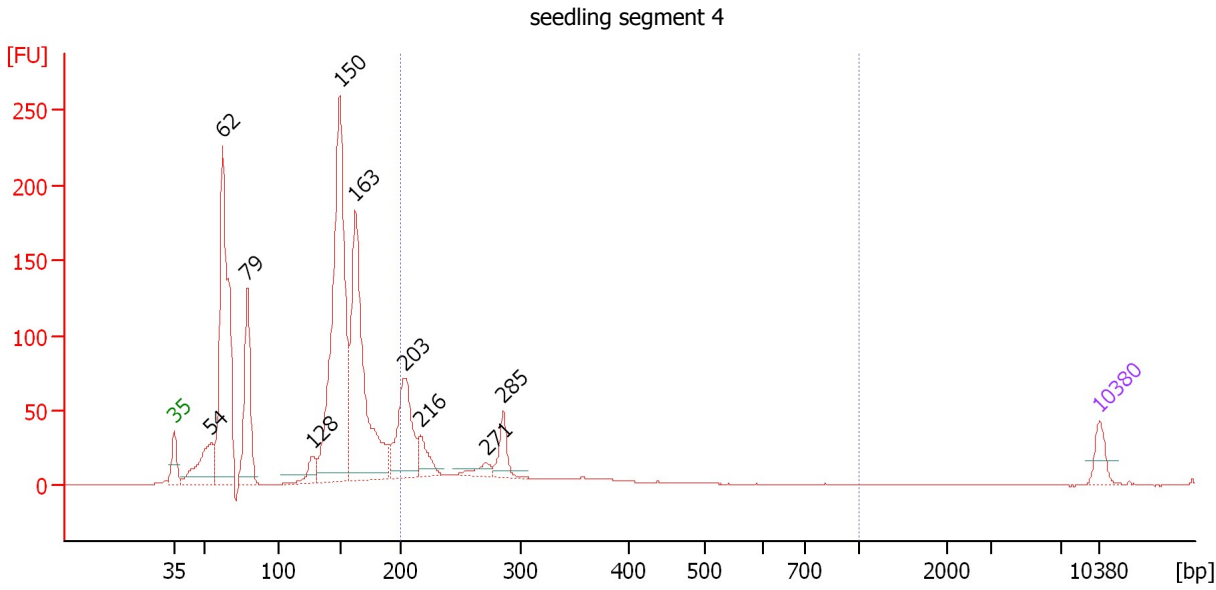
Region table for sample 1 : - ctrol

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	404	27.7	336.2	73.84	6	42.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-10-06\2017-10-06_004.xad

Created: 10/6/2017 3:32:07 PM
 Modified: 10/6/2017 3:53:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : seedling segment 4

Number of peaks found: 10 Corr. Area 1: 361.1
 Noise: 0.1

Peak table for sample 2 : seedling segment 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	54	380.43	10,641.7		45.78
3	62	1,538.15	37,603.1		46.64
4	79	627.69	12,028.4		48.51
5	128	104.76	1,240.6		53.44
6	150	1,988.42	20,068.0		55.52
7	163	1,404.67	13,074.7		56.70
8	203	520.27	3,880.9		60.45
9	216	146.21	1,025.0		61.62
10	271	70.97	397.5		66.55
11	285	151.04	802.0		67.89
12	10,380	75.00	10.9	Upper Marker	113.00

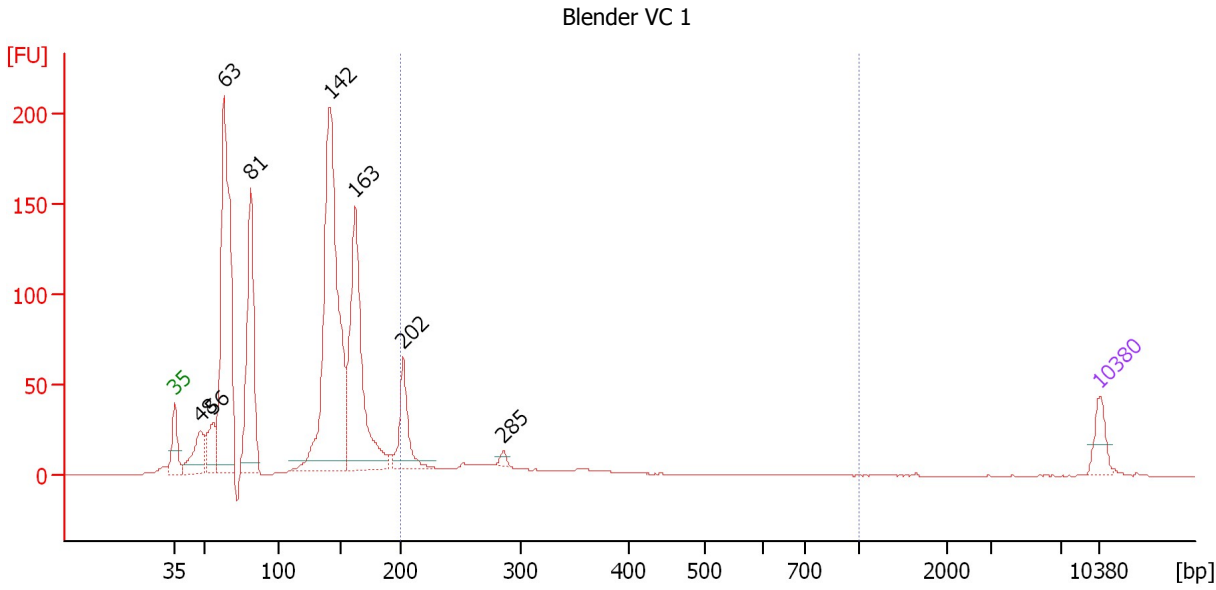
Region table for sample 2 : seedling segment 4

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	280	361.1	7,446.7	1,255.87	19	36.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-10-06\2017-10-06_004.xad

Created: 10/6/2017 3:32:07 PM
 Modified: 10/6/2017 3:53:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : Blender VC 1

Number of peaks found: 8 Corr. Area 1: 180.9
 Noise: 0.1

Peak table for sample 3 : Blender VC 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	48	221.58	7,052.4		44.96
3	56	180.87	4,935.2		45.93
4	63	1,424.19	34,272.5		46.75
5	81	798.66	14,885.1		48.76
6	142	1,780.74	18,963.7		54.78
7	163	1,004.60	9,360.3		56.69
8	202	266.02	1,997.7		60.32
9	285	22.24	118.1		67.87
10	10,380	75.00	10.9	Upper Marker	113.00

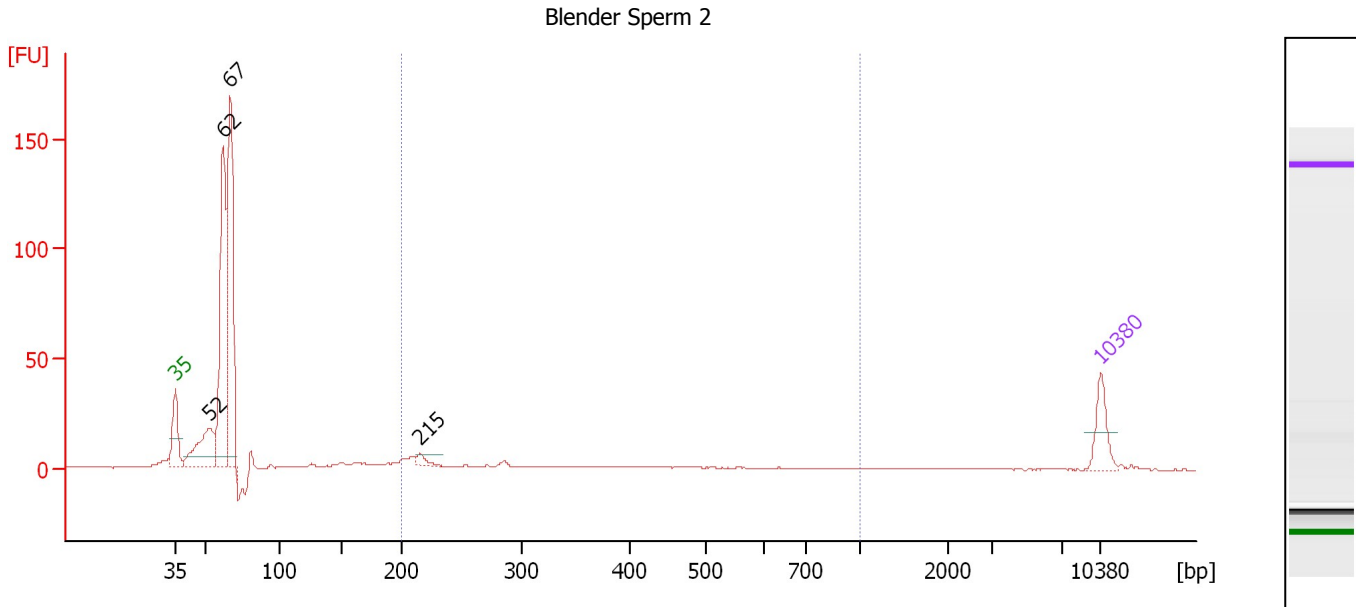
Region table for sample 3 : Blender VC 1

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	291	180.9	3,556.1	604.57	11	40.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2017-10-06\2017-10-06_004.xad

Created: 10/6/2017 3:32:07 PM
 Modified: 10/6/2017 3:53:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : Blender Sperm 2

Number of peaks found: 4 Corr. Area 1: 49.3
 Noise: 0.1

Peak table for sample 4 : Blender Sperm 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	52	246.46	7,140.3		45.58
3	62	686.80	16,835.9		46.62
4	67	721.00	16,350.2		47.17
5	215	23.85	168.4		61.49
6	10,380	75.00	10.9	Upper Marker	113.00

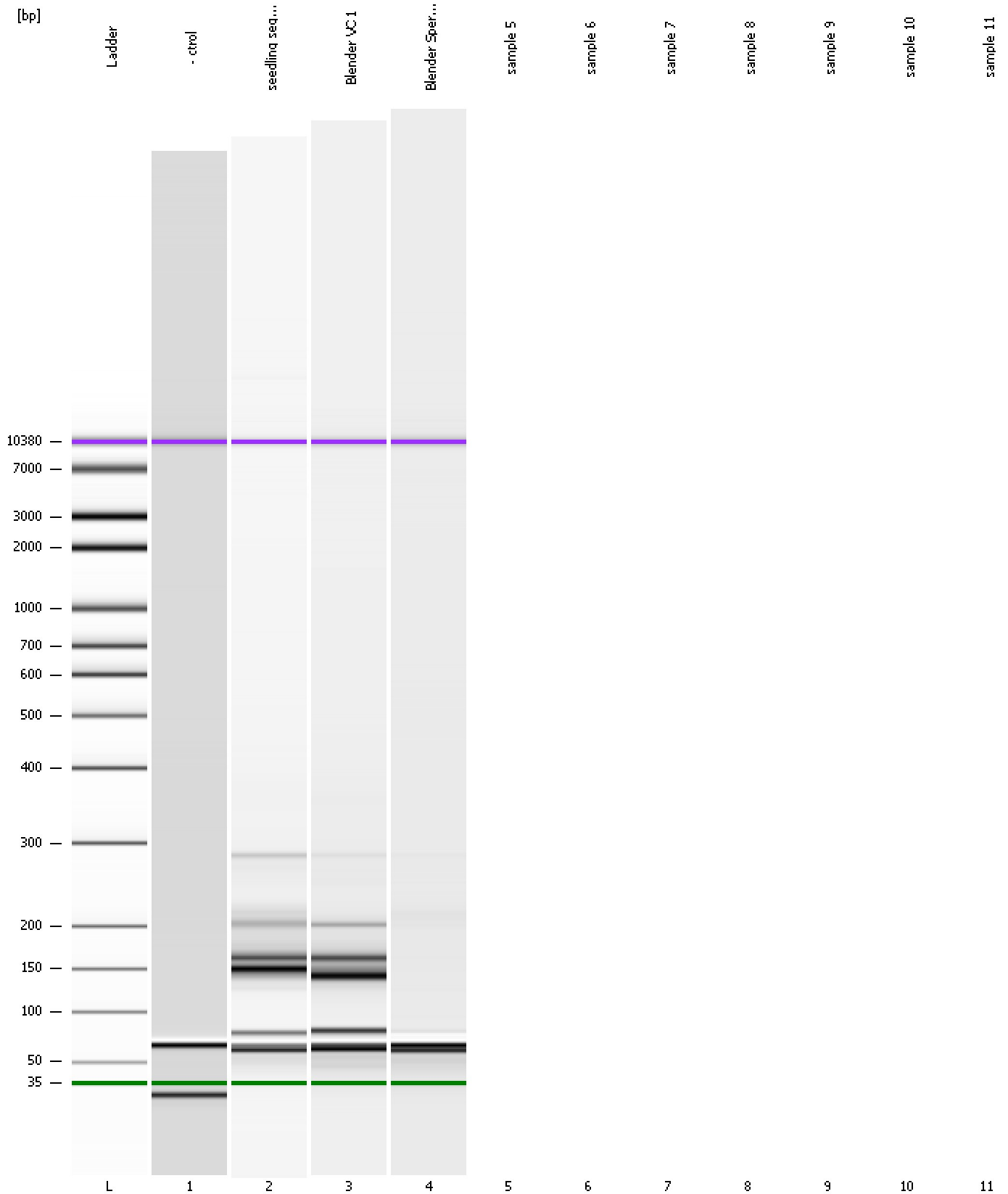
Region table for sample 4 : Blender Sperm 2

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	344	49.3	908.1	166.33	10	47.5

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-10-06\2017-10-06_004.xad

Created: 10/6/2017 3:32:07 PM
Modified: 10/6/2017 3:53:29 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2017-10-06\2017-10-06_004.xad

Created: 10/6/2017 3:32:07 PM
Modified: 10/6/2017 3:53:29 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.