

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2017-10-11\2017-10-11\_001.xad

Created: 10/11/2017 11:16:18 AM  
Modified: 10/11/2017 11:40:32 AM

**Electrophoresis File Run Summary**

Instrument Information:

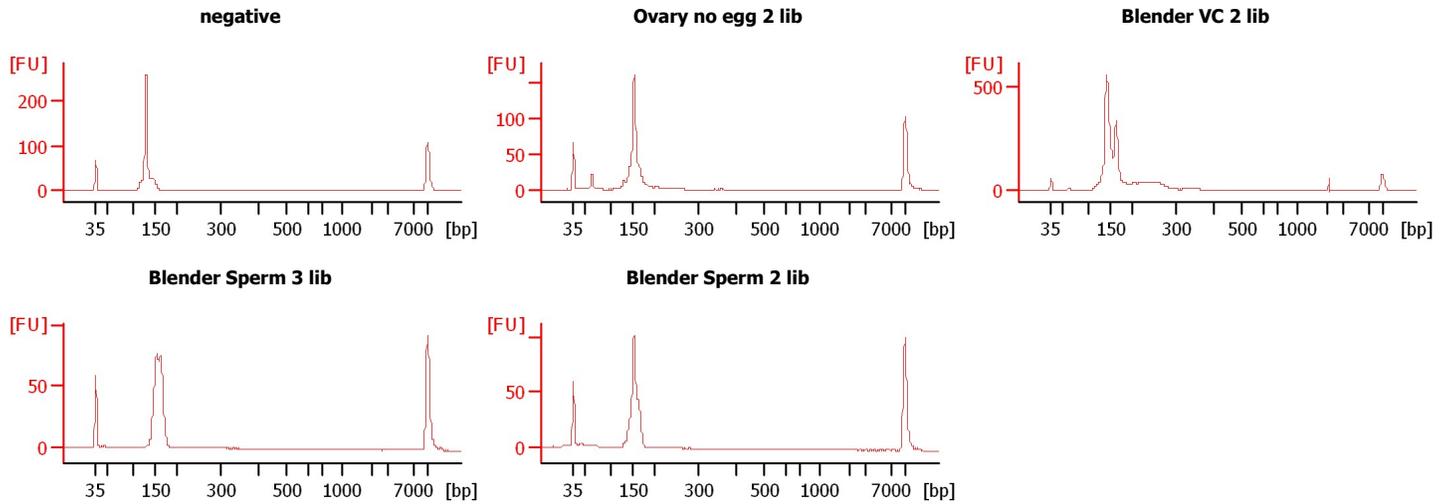
Instrument Name: DE13701086                      Firmware: C.01.069  
Serial#: DE13701086                              Type: G2938B

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:



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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
negative		<input type="checkbox"/>	✓			
Ovary no egg 2 lib		<input type="checkbox"/>	✓			
Blender VC 2 lib		<input type="checkbox"/>	✓			
Blender Sperm 3 lib		<input type="checkbox"/>	✓			
Blender Sperm 2 lib		<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 :**

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**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

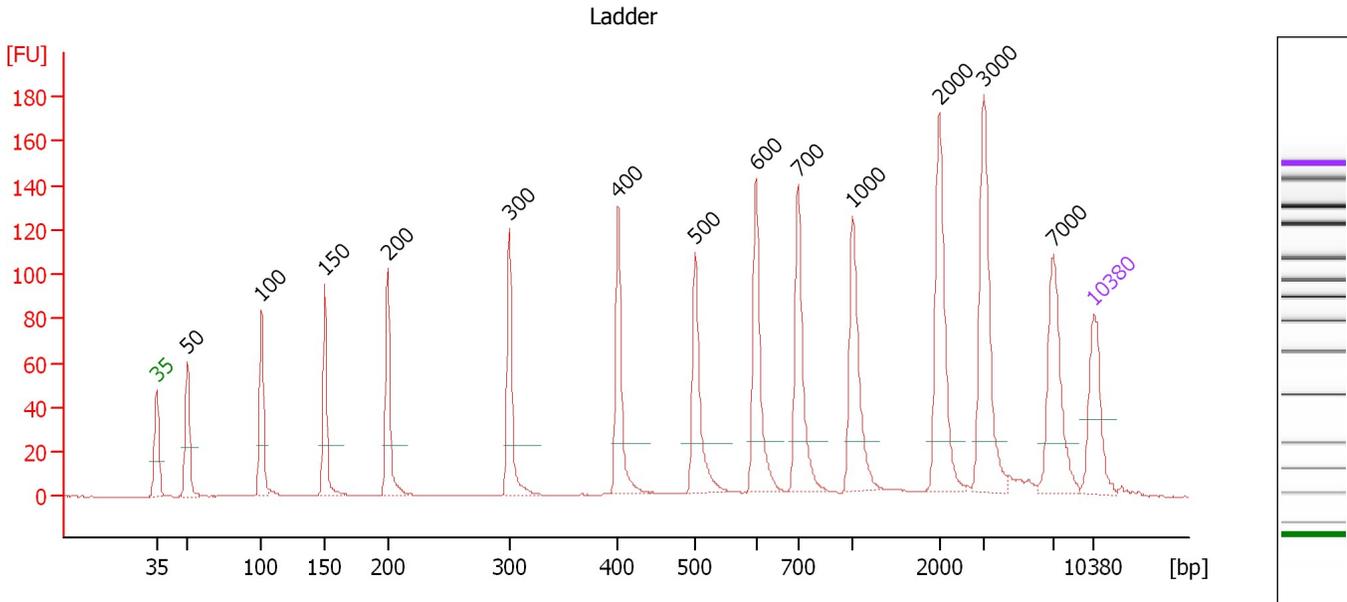
**Ladder**

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

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**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.1

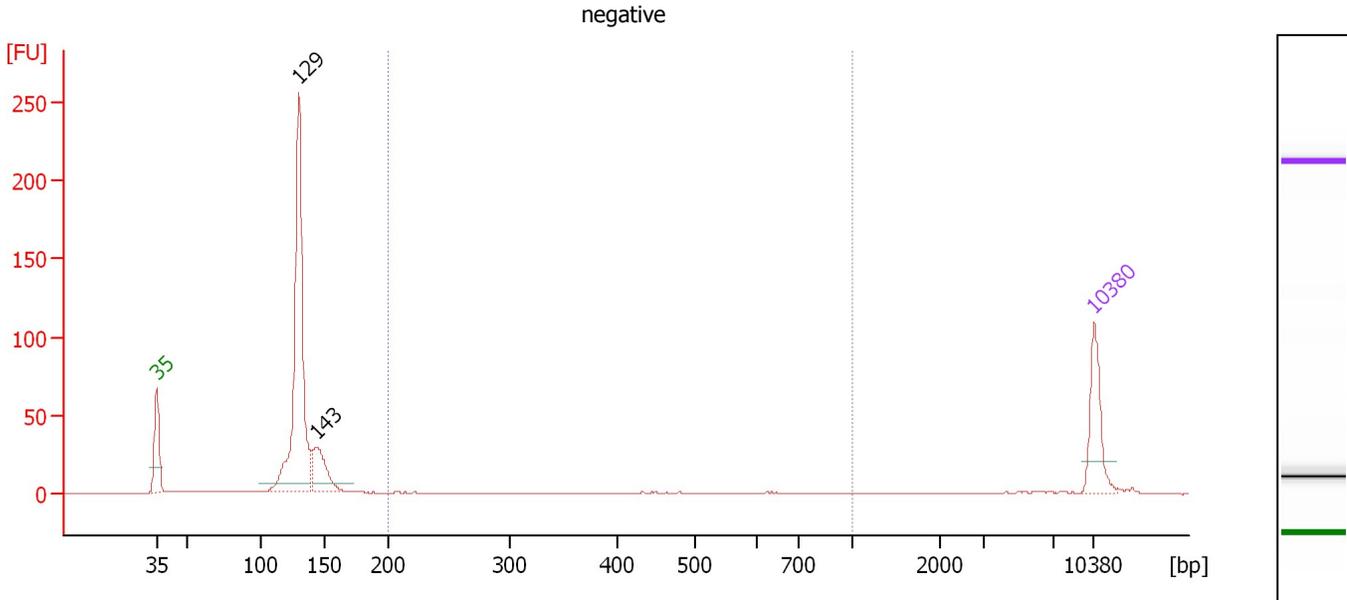
**Peak table for Ladder**

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	45.36
3	100	150.00	2,272.7	Ladder Peak	50.86
4	150	150.00	1,515.2	Ladder Peak	55.58
5	200	150.00	1,136.4	Ladder Peak	60.24
6	300	150.00	757.6	Ladder Peak	69.38
7	400	150.00	568.2	Ladder Peak	77.44
8	500	150.00	454.5	Ladder Peak	83.18
9	600	150.00	378.8	Ladder Peak	87.75
10	700	150.00	324.7	Ladder Peak	90.89
11	1,000	150.00	227.3	Ladder Peak	94.97
12	2,000	150.00	113.6	Ladder Peak	101.46
13	3,000	150.00	75.8	Ladder Peak	104.75
14	7,000	150.00	32.5	Ladder Peak	109.95
15	10,380	75.00	10.9	Upper Marker	113.00

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**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : negative**

Number of peaks found: 2                      Corr. Area 1: 33.1  
 Noise: 0.2

**Peak table for sample 1 : negative**

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	129	588.34	6,903.9		53.61
3	143	107.48	1,141.1		54.89
4	10,380	75.00	10.9	Upper Marker	113.00

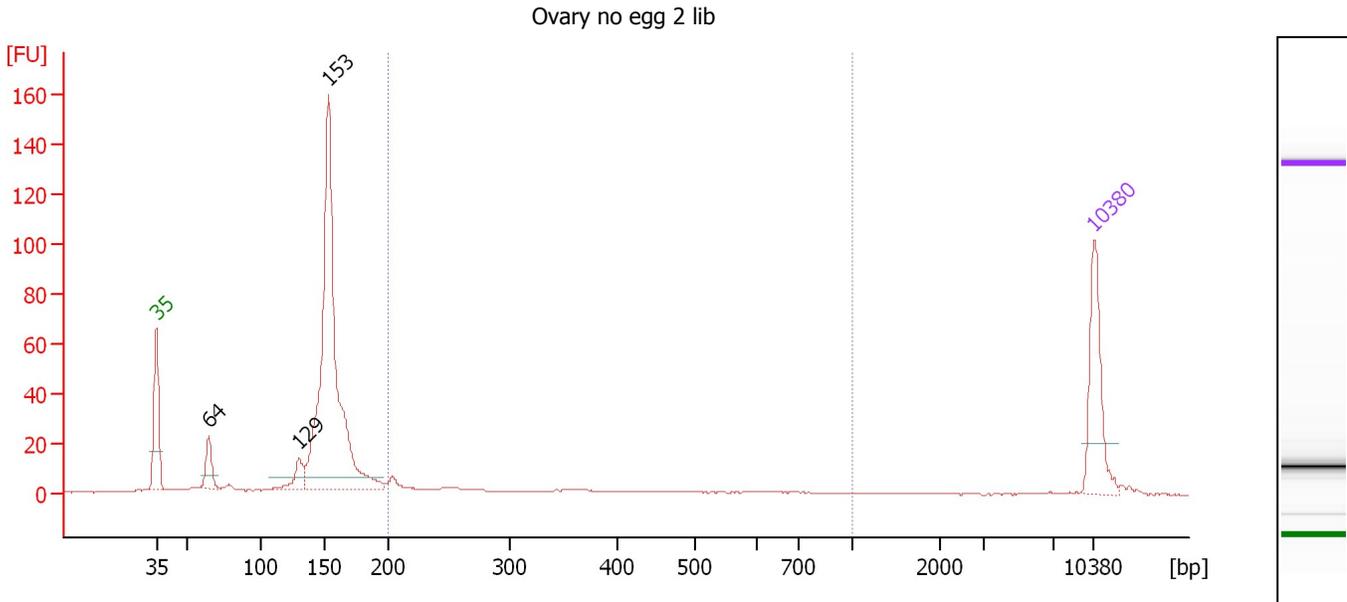
**Region table for sample 1 : negative**

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	481	33.1	168.9	41.80	7	39.6

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**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : Ovary no egg 2 lib**

Number of peaks found: 3                      Corr. Area 1: 78.3  
 Noise: 0.2

**Peak table for sample 2 : Ovary no egg 2 lib**

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	64	38.12	900.6		46.91
3	129	33.30	391.0		53.60
4	153	559.08	5,546.1		55.83
5	10,380	75.00	10.9	Upper Marker	113.00

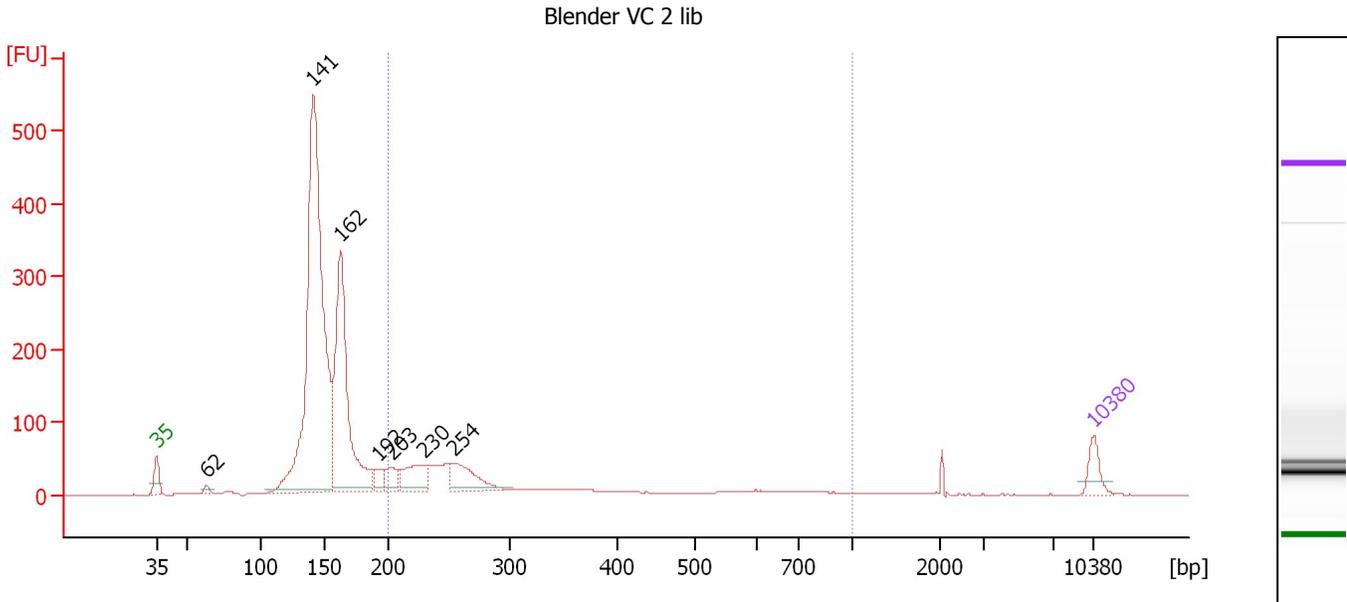
**Region table for sample 2 : Ovary no egg 2 lib**

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	398	78.3	490.9	101.00	14	45.5

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**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : Blender VC 2 lib**

Number of peaks found: 7                      Corr. Area 1: 672.7  
 Noise: 0.1

**Peak table for sample 3 : Blender VC 2 lib**

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	62	27.04	656.6		46.72
3	141	2,940.16	31,617.2		54.72
4	162	1,326.24	12,377.0		56.73
5	192	68.46	540.3		59.49
6	203	108.24	806.4		60.55
7	230	208.87	1,377.2		62.96
8	254	217.81	1,299.3		65.18
9	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 3 : Blender VC 2 lib**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	333	672.7	6,444.7	1,164.06	23	49.2

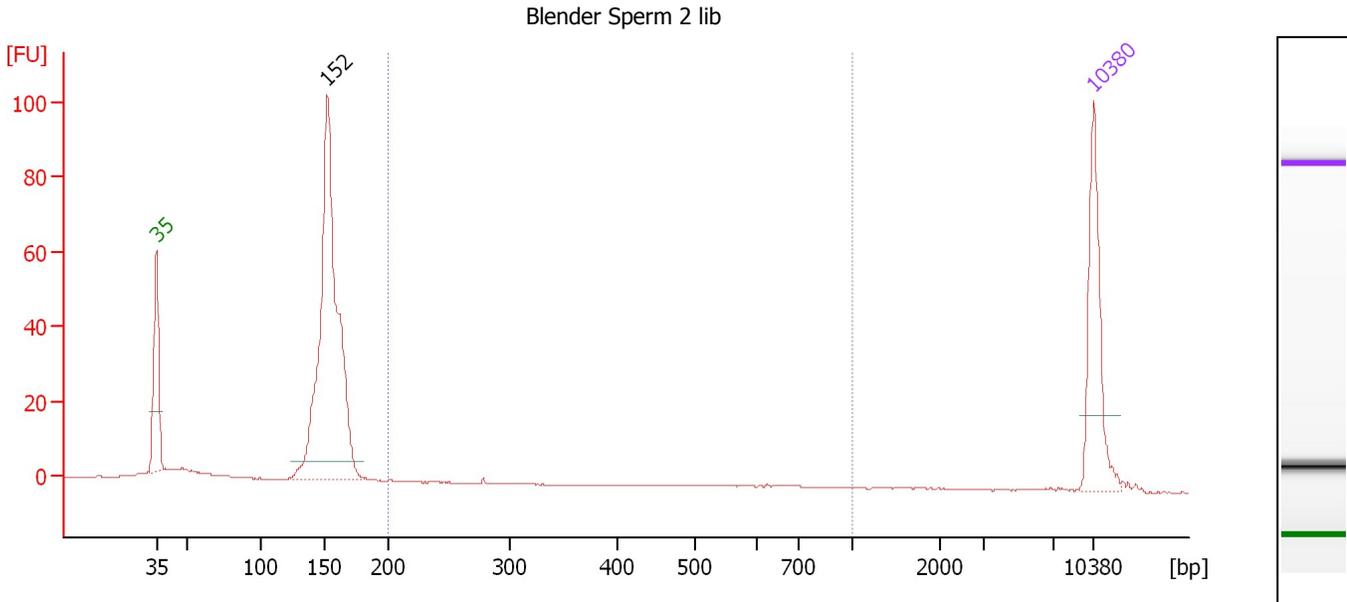




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**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : Blender Sperm 2 lib**

Number of peaks found: 1                      Corr. Area 1: 4.2  
 Noise: 0.1

**Peak table for sample 5 : Blender Sperm 2 lib**

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	152	433.92	4,323.1		55.77
3	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 5 : Blender Sperm 2 lib**

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	558	4.2	18.6	5.03	1	35.3

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**Gel Image**

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**Invalid Samples**

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.