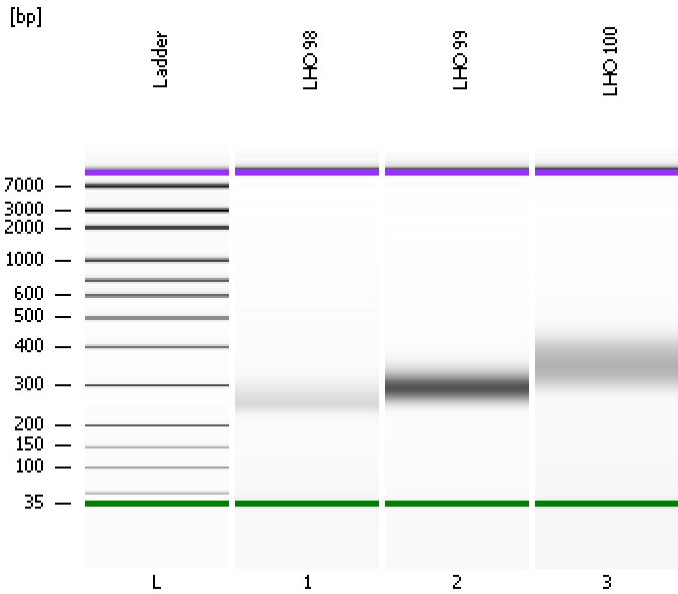


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... and Settings\Bioanalyzer\2013-11-15\2013-11-15\_001\_LHO.xad

Created: 11/15/2013 10:26:18 AM  
Modified: 11/15/2013 11:34:31 AM

**Electrophoresis File Run Summary**



Instrument Information:

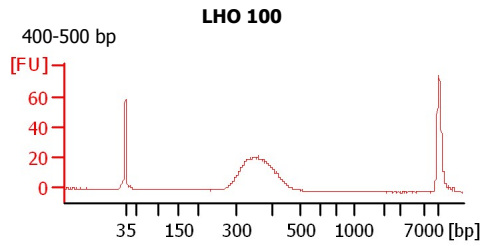
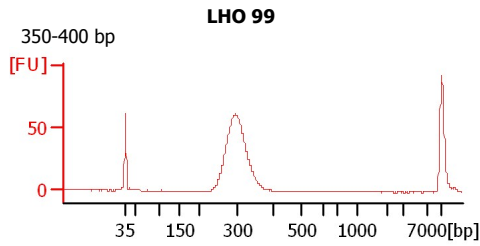
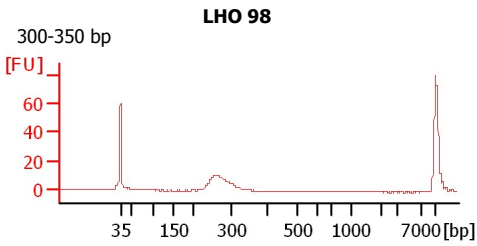
Instrument Name: DE13701086      Firmware: C.01.069  
Serial#: DE13701086      Type: G2938B

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:



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... and Settings\Bioanalyzer\2013-11-15\2013-11-15\_001\_LHO.xad

Created: 11/15/2013 10:26:18 AM  
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**Electrophoresis File Run Summary (Chip Summary)**

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
LHO 98	300-350 bp	<input type="checkbox"/>	✓			
LHO 99	350-400 bp	<input type="checkbox"/>	✓			
LHO 100	400-500 bp	<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

**Chip Lot #**

**Reagent Kit Lot #**

**Chip Comments :**

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... and Settings\Bioanalyzer\2013-11-15\2013-11-15\_001\_LHO.xad

Created: 11/15/2013 10:26:18 AM  
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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

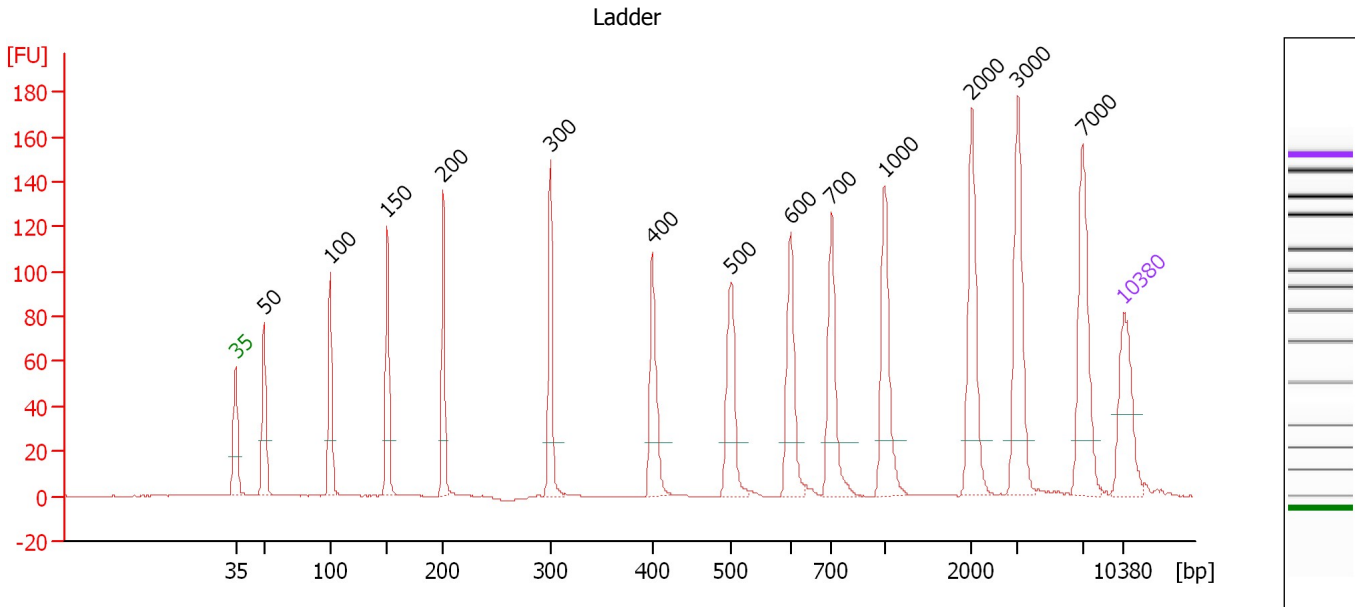
**Ladder**

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

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... and Settings\Bioanalyzer\2013-11-15\2013-11-15\_001\_LHO.xad

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**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.1

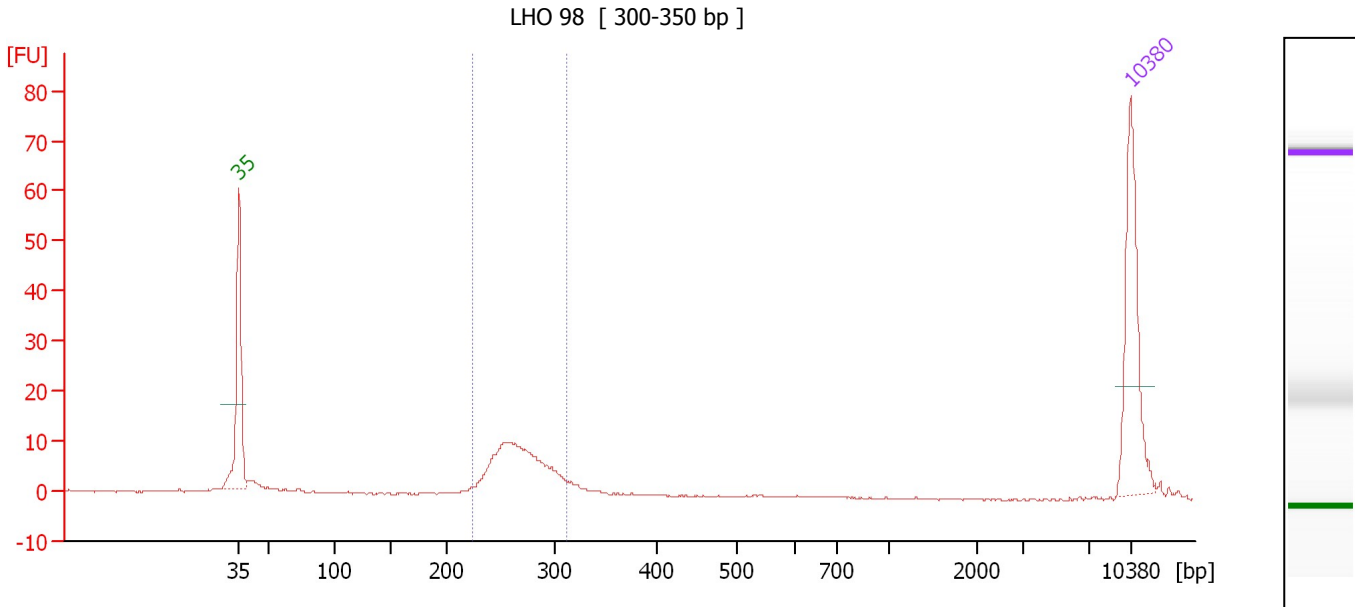
**Peak table for Ladder**

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	45.27	50	150.00	4,545.5	Ladder Peak
3	50.48	100	150.00	2,272.7	Ladder Peak
4	54.97	150	150.00	1,515.2	Ladder Peak
5	59.39	200	150.00	1,136.4	Ladder Peak
6	67.83	300	150.00	757.6	Ladder Peak
7	75.82	400	150.00	568.2	Ladder Peak
8	82.08	500	150.00	454.5	Ladder Peak
9	86.74	600	150.00	378.8	Ladder Peak
10	89.97	700	150.00	324.7	Ladder Peak
11	94.13	1,000	150.00	227.3	Ladder Peak
12	101.03	2,000	150.00	113.6	Ladder Peak
13	104.64	3,000	150.00	75.8	Ladder Peak
14	109.72	7,000	150.00	32.5	Ladder Peak
15	113.00	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... and Settings\Bioanalyzer\2013-11-15\2013-11-15\_001\_LHO.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : LHO 98**

Number of peaks found: 0                      Corr. Area 1: 75.6  
 Noise: 0.2

**Peak table for sample 1 : LHO 98**

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	113.00	10,380	75.00	10.9	Upper Marker

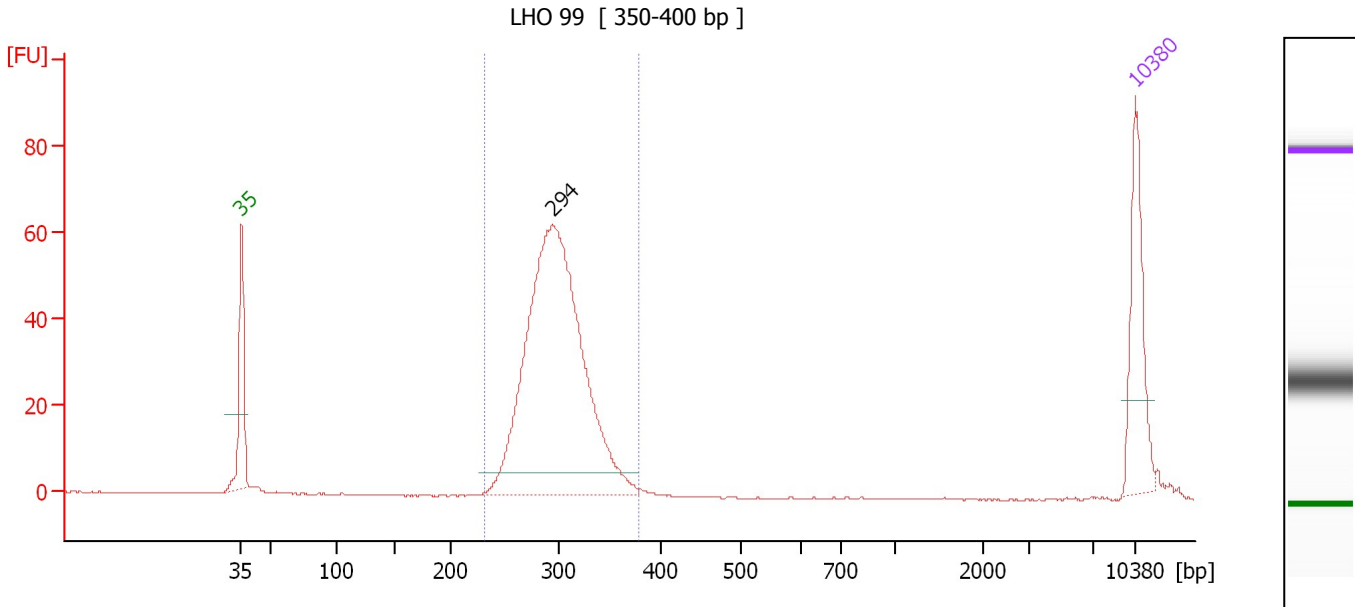
**Region table for sample 1 : LHO 98**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
223	311	267	685.1	120.26	75.6	81	7.6	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... and Settings\Bioanalyzer\2013-11-15\2013-11-15\_001\_LHO.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : LHO 99**

Number of peaks found: 1                      Corr. Area 1: 488.2  
 Noise: 0.2

**Peak table for sample 2 : LHO 99**

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	67.29	294	666.56	3,439.6	
3	113.00	10,380	75.00	10.9	Upper Marker

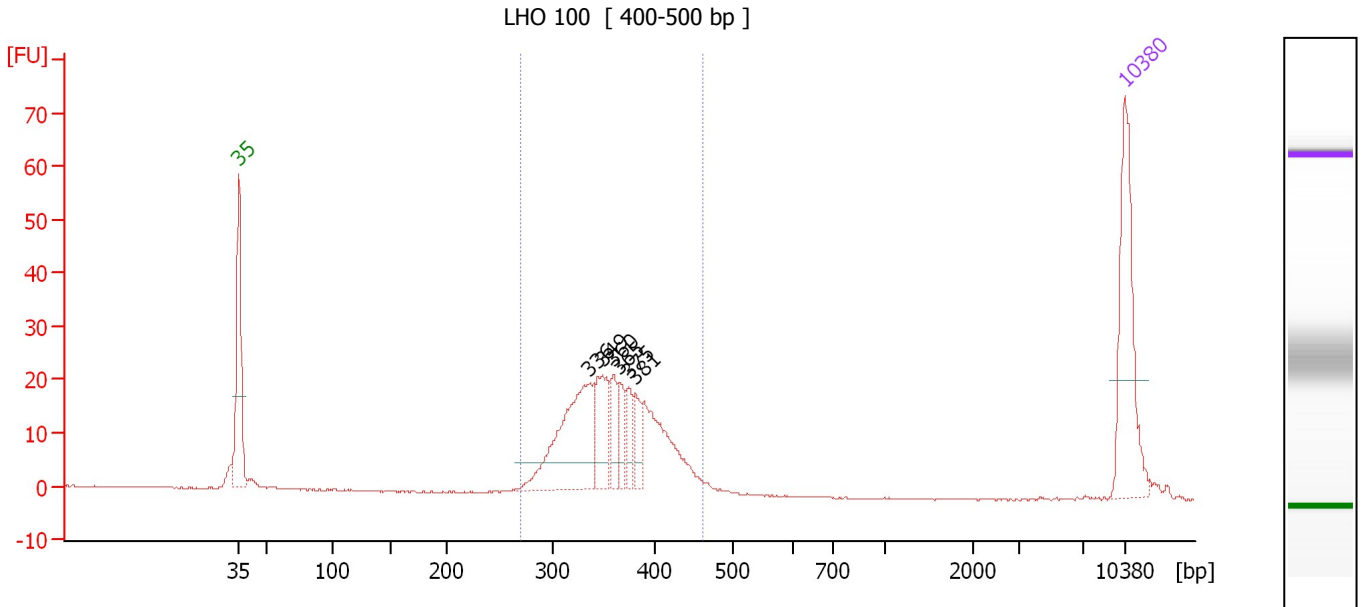
**Region table for sample 2 : LHO 99**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
231	379	298	3,329.0	650.45	488.2	98	8.5	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... and Settings\Bioanalyzer\2013-11-15\2013-11-15\_001\_LHO.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : LHO 100**

Number of peaks found: 6                      Corr. Area 1: 261.9  
 Noise: 0.2

**Peak table for sample 3 : LHO 100**

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	70.73	336	126.79	571.2	
3	71.74	349	49.29	214.0	
4	72.63	360	23.87	100.5	
5	73.05	365	20.67	85.7	
6	73.81	375	19.28	77.9	
7	74.31	381	20.22	80.4	
8	113.00	10,380	75.00	10.9	Upper Marker

**Region table for sample 3 : LHO 100**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
269	461	360	1,548.0	362.97	261.9	94	11.0	Blue

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... and Settings\Bioanalyzer\2013-11-15\2013-11-15\_001\_LHO.xad

Created: 11/15/2013 10:26:18 AM  
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**Gel Image**

