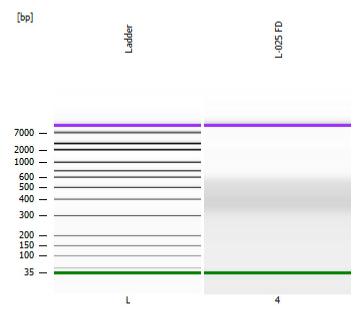
## **Electrophoresis File Run Summary**



### Instrument Information:

 Instrument Name:
 DE13806062
 Firmware:
 C.01.069

 Serial#:
 DE13806062
 Type:
 G2938C

### Assay Information:

Assay Origin Path: C:\Program Files (x86)\Agilent\2100 bioanalyzer\2100

expert\assays\dsDNA\High Sensitivity DNA.xsy

Printed: 10/9/2018 10:20:32 AM

Assay Class: High Sensitivity DNA Assay

Version: 1.03

Assay Comments: Copyright © 2003-2010 Agilent Technologies

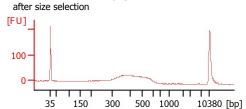
#### Chip Information:

Chip Lot #:

Reagent Kit Lot #:

Chip Comments:





Page 2 of 8

Printed: 10/9/2018 10:20:32 AM

 $\label{light} \begin{tabular}{ll} High Sensitivity DNA Assay \\ C:\...braries\2018-10-09\181009\_L-025\_FD\_after\_size\_selection.xad \\ \end{tabular}$ 10/9/2018 9:26:39 AM 10/9/2018 9:55:16 AM Assay Class: Created: Data Path: Modified: **Electrophoresis File Run Summary (Chip Summary) Sample Name Sample Comment** Rest. Digest Status Observation **Result Label** Res ult Col or L-025 FD after size selection Ladder Chip Lot # Reagent Kit Lot # **Chip Comments:** 

Printed: 10/9/2018 10:20:32 AM

Assay Class: High Sensitivity DNA Assay Created: 10/9/2018 9:26:39 AM Data Path: C:\...braries\2018-10-09\181009\_L-025\_FD\_after\_size\_selection.xad Created: 10/9/2018 9:25:16 AM

## **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**

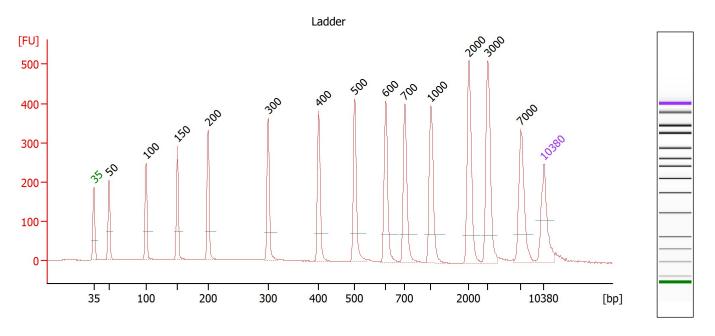
Filter Width [s]: 0.5 Polynomial Order: 4

# 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 C:\...braries\2018-10-09\181009\_L-025\_FD\_after\_size\_selection.xad Created: 10/9/2018 9:26:39 AM Modified: 10/9/2018 9:55:16 AM

### **Electropherogram Summary**



### **Overall Results for Ladder**

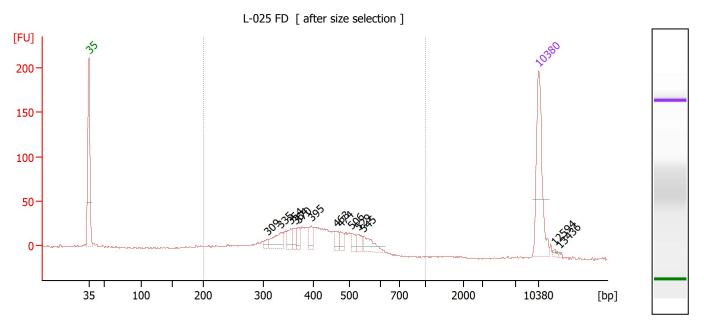
Noise: 0.5

Peak	table	for	Ladder
rcan	cavic	101	Lauuci

Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2	L	50	150.00	4,545.5	Ladder Peak
3	L	100	150.00	2,272.7	Ladder Peak
4	L	150	150.00	1,515.2	Ladder Peak
5	L	200	150.00	1,136.4	Ladder Peak
6	L	300	150.00	757.6	Ladder Peak
7	L	400	150.00	568.2	Ladder Peak
8		500	150.00	454.5	Ladder Peak
9	L	600	150.00	378.8	Ladder Peak
10	L	700	150.00	324.7	Ladder Peak
11		1,000	150.00	227.3	Ladder Peak
12	L	2,000	150.00	113.6	Ladder Peak
13		3,000	150.00	75.8	Ladder Peak
14		7,000	150.00	32.5	Ladder Peak
15		10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay C:\...braries\2018-10-09\181009\_L-025\_FD\_after\_size\_selection.xad Created: 10/9/2018 9:26:39 AM Modified: 10/9/2018 9:55:16 AM

## **Electropherogram Summary Continued ...**



# Overall Results for sample 4: <u>L-025 FD</u>

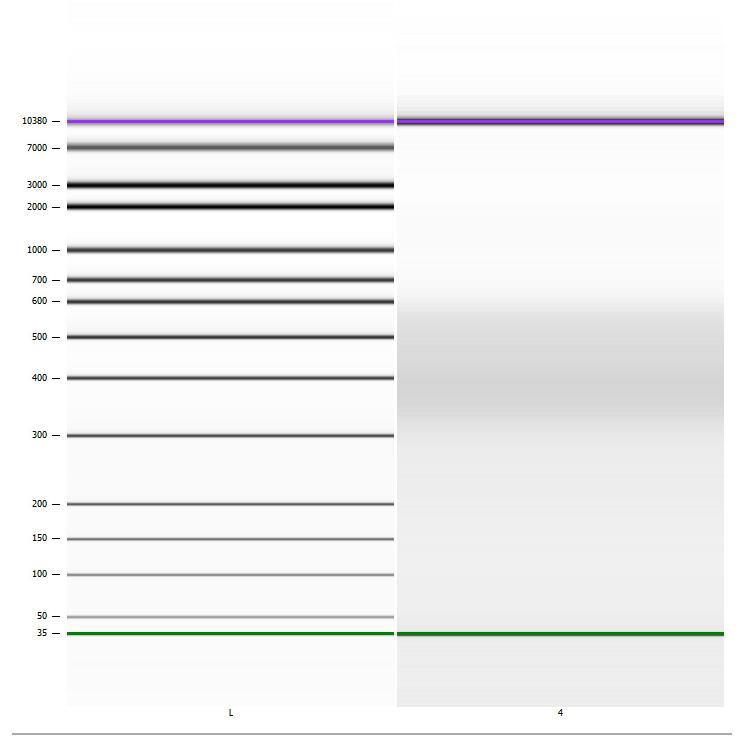
Number of peaks found: 13 Corr. Area 1: 596.9

Noise: 0.6

Peak table for sample 4:		for sample 4:	<u>L-025 FD</u>		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2		309	6.61	32.5	
3		335	26.80	121.1	
4		354	14.13	60.4	
5		364	11.33	47.2	
6		370	11.64	47.7	
7		395	13.43	51.5	
8		463	10.52	34.4	
9		474	10.65	34.0	
10		506	9.49	28.4	
11		529	10.79	30.9	
12		545	19.42	54.0	
13		10,380	75.00	10.9	Upper Marker
14		12,594	0.00	0.0	
15		13,436	0.00	0.0	

Region table for sample 4: <u>L-025 FD</u>

Name	From [bp]	To [bp	] % of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
Region 1	200	1,000	90	408	22.4	337.90	1,360.9	

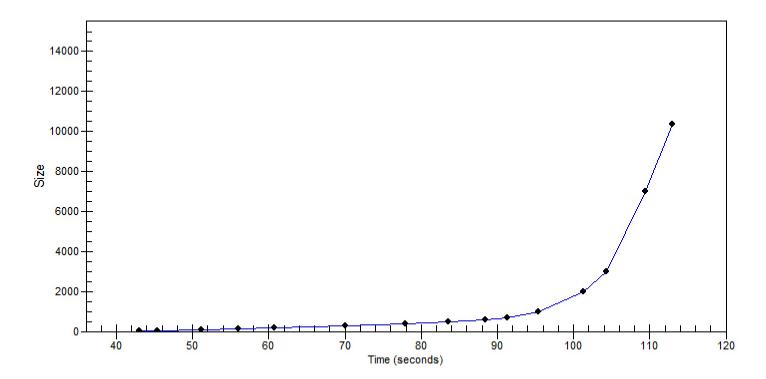


Printed: 10/9/2018 10:20:32 AM

Assay Class: High Sensitivity DNA Assay C:\...braries\2018-10-09\181009\_L-025\_FD\_after\_size\_selection.xad Created: 10/9/2018 9:26:39 AM Modified: 10/9/2018 9:55:16 AM

**Curves** 

# **Standard Curve**



Assay Class: High Sensitivity DNA Assay C:\...braries\2018-10-09\181009\_L-025\_FD\_after\_size\_selection.xad Created: 10/9/2018 9:26:39 AM Modified: 10/9/2018 9:55:16 AM

### **Run Logbook**

<b>Description</b> Run ended on port 5 (Number of wells acquired: 5)	Number	<b>Source</b> Instrument	<b>Category</b> Run	Sub Category	<b>Time</b> 10/9/2018 9:47:59 AM	<b>Time Zone</b> (GMT +02:00) W. Europe Standard Time	<b>User</b> Admin	<b>Host</b> Datasystem01
Run started on port 5 (File: C:\Users\Admin \Desktop\Librari es\2018-10-09\2100 expert_High Sensitivity DNA Assay_DE13806 062_2018-10-09_09-26-40.xad)		Instrument	Run		10/9/2018 9:26:45 AM	(GMT +02:00) W. Europe Standard Time	Admin	Datasystem01
Product Number : G2938C		Instrument	Run		10/9/2018 9:26:45 AM	(GMT +02:00) W. Europe Standard Time	Admin	Datasystem01
Name :		Instrument	Run		10/9/2018 9:26:45 AM	(GMT +02:00) W. Europe Standard Time	Admin	Datasystem01
Vendor : Agilent Technologies		Instrument	Run		10/9/2018 9:26:45 AM	(GMT +02:00) W. Europe Standard Time	Admin	Datasystem01
Serial# : DE13806062		Instrument	Run		10/9/2018 9:26:45 AM	(GMT +02:00) W. Europe Standard Time	Admin	Datasystem01
Firmware : C.01.069		Instrument	Run		10/9/2018 9:26:45 AM	(GMT +02:00) W. Europe Standard Time	Admin	Datasystem01
Cartridge : Electrode		Instrument	Run		10/9/2018 9:26:45 AM	(GMT +02:00) W. Europe Standard Time	Admin	Datasystem01