

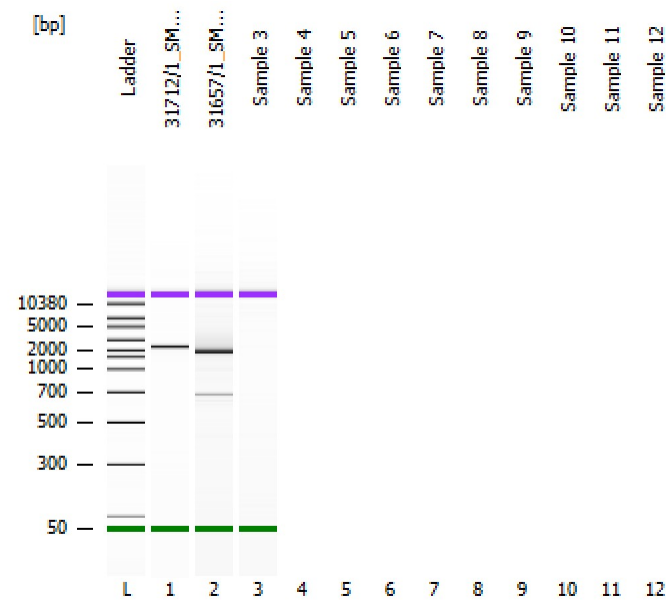
Assay Class: DNA 12000

Data Path: C:\...ta\2023-05-23\Bioanalyzer\_DNA 12000\_2023-05-23\_15-24-15.xad

Created: 5/23/2023 3:24:15 PM

Modified: 5/23/2023 3:58:07 PM

Electrophoresis File Run Summary



Instrument Information:

Instrument Name: DE34903176

Serial#: DE34903176

Firmware: C.01.069

Type: G2938C

Assay Information:

Assay Origin Path: C:\Program Files (x86)\Agilent\2100 bioanalyzer\2100 expert\assays\dsDNA\DNA 12000 Series II.xsy

Assay Class: DNA 12000

Version: 2.4

Assay Comments: DNA Analysis 100 -12000 bp

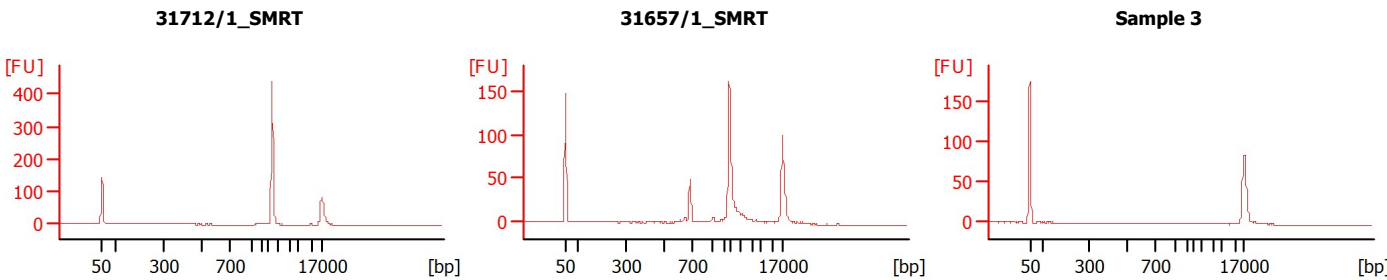
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Chip Information:

Chip Lot #:

Reagent Kit Lot #:

Chip Comments:



Assay Class: DNA 12000  
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Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	St at us	Observation	Result Label	Result Color
31712/1_SMR T		<input type="checkbox"/>		✓		
31657/1_SMR T		<input type="checkbox"/>		✓		
Sample 3		<input type="checkbox"/>		✓		
Sample 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"/>				
Sample 12		<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.) : 13  
Minimum Visible Range [s] : 20  
Maximum Visible Range [s] : 99  
Start Analysis Time Range [s] : 20  
End Analysis Time Range [s] : 98.95  
Ladder Concentration [ng/μl] : 44  
Uses Standard Area for Ladder Fragments  
Lower Marker Concentration [ng/μl] : 8.3  
Upper Marker Concentration [ng/μl] : 4.2  
Used Upper Marker for Quantitation  
This is a Qualitative Assay Only  
Standard Curve Fit is Point to Point  
Show Data Aligned to Lower and Upper Marker

### Integrator Settings

Integration Start Time [s] : 20  
Integration End Time [s] : 98.95  
Slope Threshold : 0.8  
Height Threshold [FU] : 20  
Area Threshold : 0.1  
Width Threshold [s] : 0.5  
Baseline Plateau [s] : 0.5

### Filter Settings

Filter Width [s] : 0.5  
Polynomial Order : 4

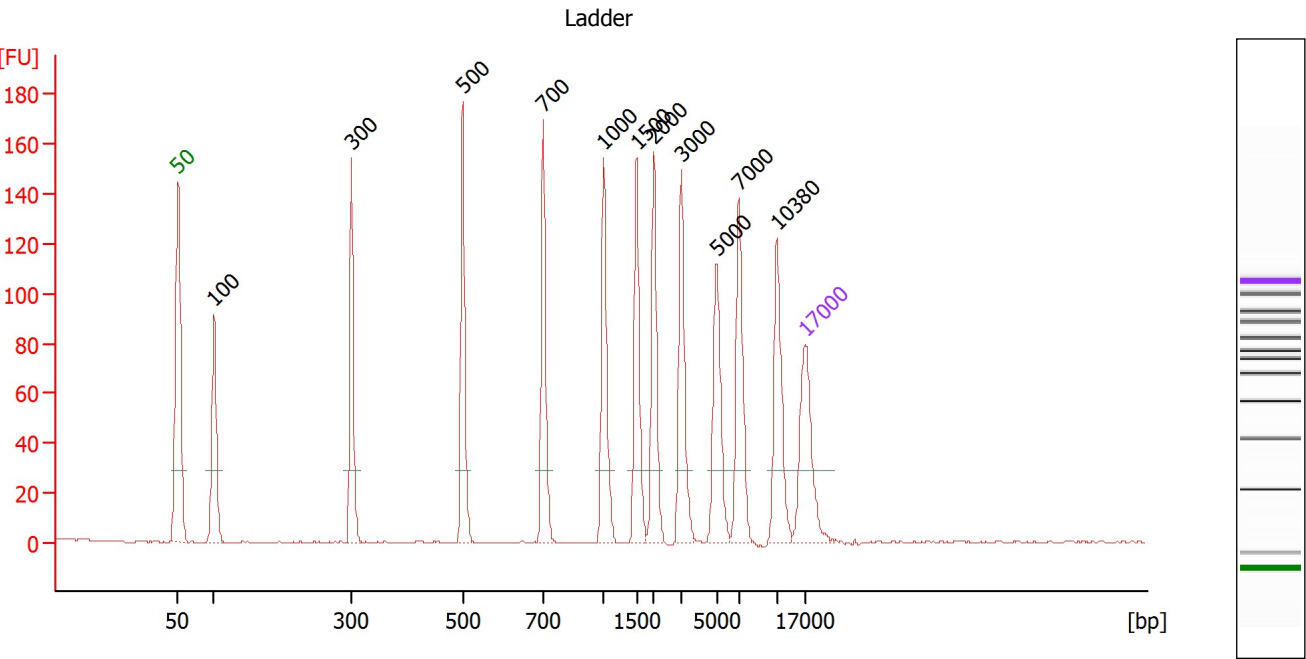
### Ladder

Ladder Peak	Size	Area
1	50	120
2	100	48
3	300	61
4	500	77
5	700	81
6	1000	86
7	1500	92
8	2000	92
9	3000	96
10	5000	100
11	7000	98
12	10380	102
13	17000	110

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Electropherogram Summary

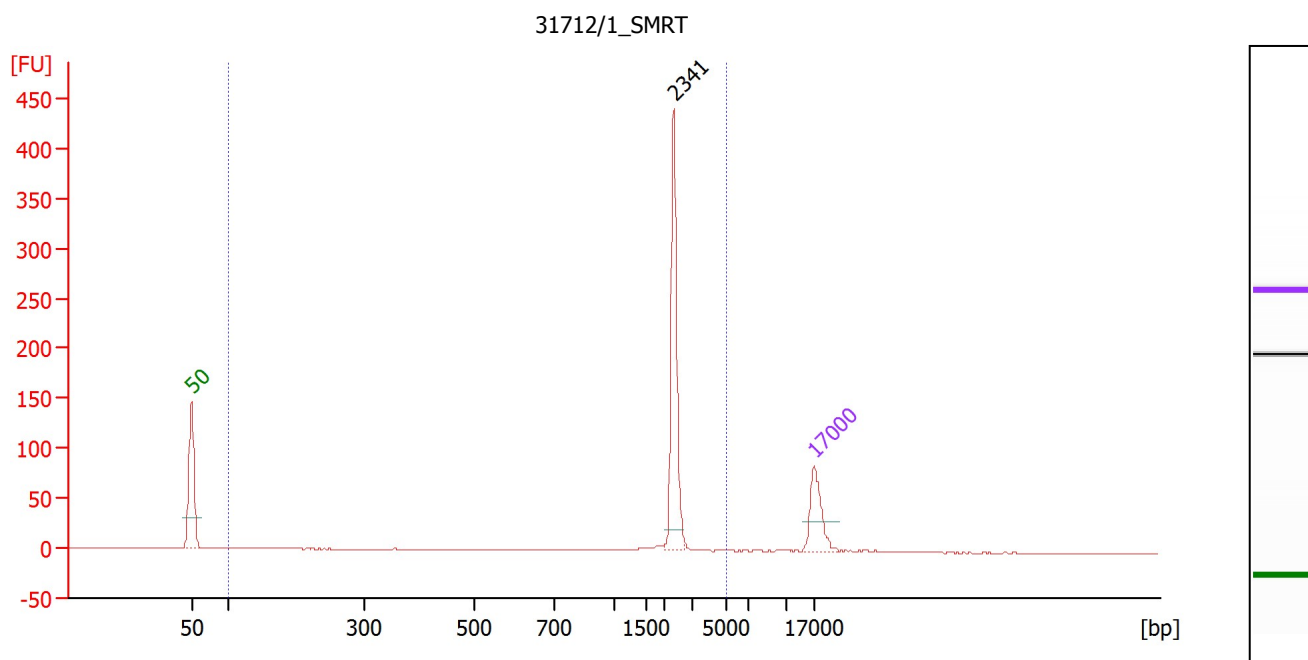


Peak table for Ladder

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
2	100	4.00	60.6	Ladder Peak
3	300	4.00	20.2	Ladder Peak
4	500	4.00	12.1	Ladder Peak
5	700	4.00	8.7	Ladder Peak
6	1,000	4.00	6.1	Ladder Peak
7	1,500	4.00	4.0	Ladder Peak
8	2,000	4.00	3.0	Ladder Peak
9	3,000	4.00	2.0	Ladder Peak
10	5,000	4.00	1.2	Ladder Peak
11	7,000	4.00	0.9	Ladder Peak
12	10,380	4.00	0.6	Ladder Peak

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**Electropherogram Summary Continued ...****Overall Results for sample 1 : 31712/1 SMRT**

Number of peaks found: 1 Area 1: 237.6

**Peak table for sample 1 : 31712/1 SMRT**

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
2	2,341	13.23	8.6	

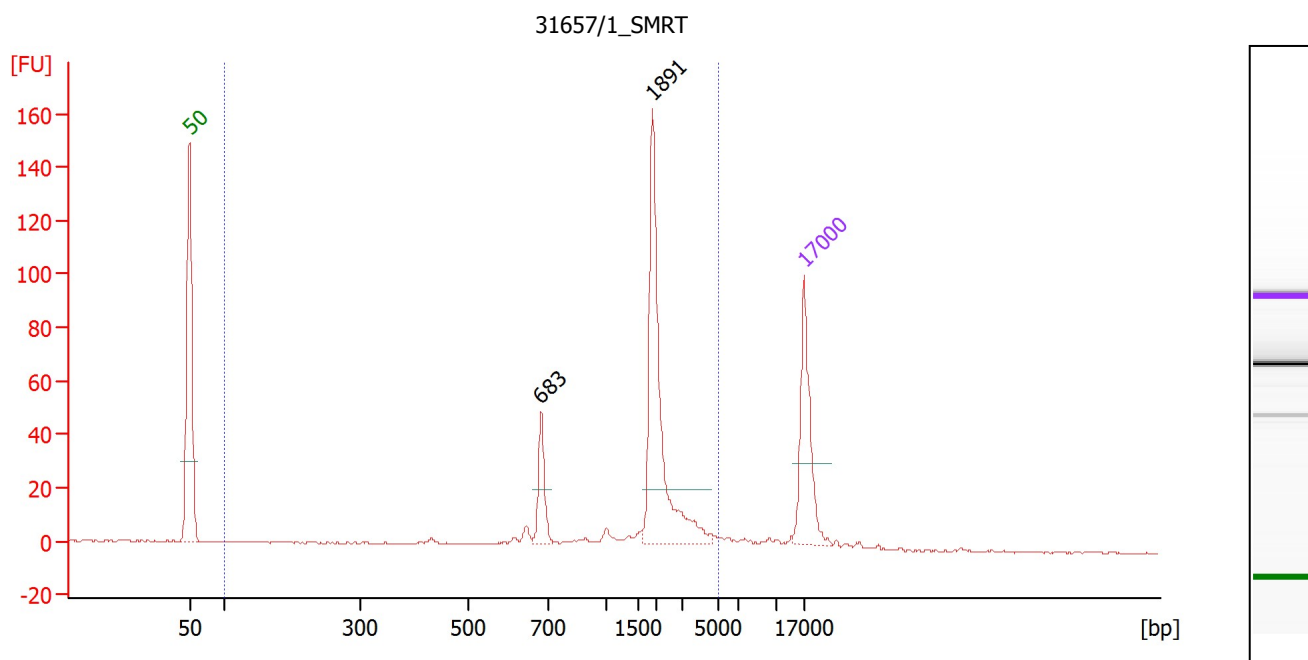
**Region table for sample 1 : 31712/1 SMRT**

From [bp]	To [bp]	Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [ng/μl]	Color
100	5,000	237.6	95	2,307	16.6	14.59	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : 31657/1 SMRT**

Number of peaks found: 2 Area 1: 224.6

**Peak table for sample 2 : 31657/1 SMRT**

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
2	683	1.72	3.8	
3	1,891	9.10	7.3	

**Region table for sample 2 : 31657/1 SMRT**

From [bp]	To [bp]	Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [ng/μl]	Color
100	5,000	224.6	90	1,833	48.8	14.04	Blue

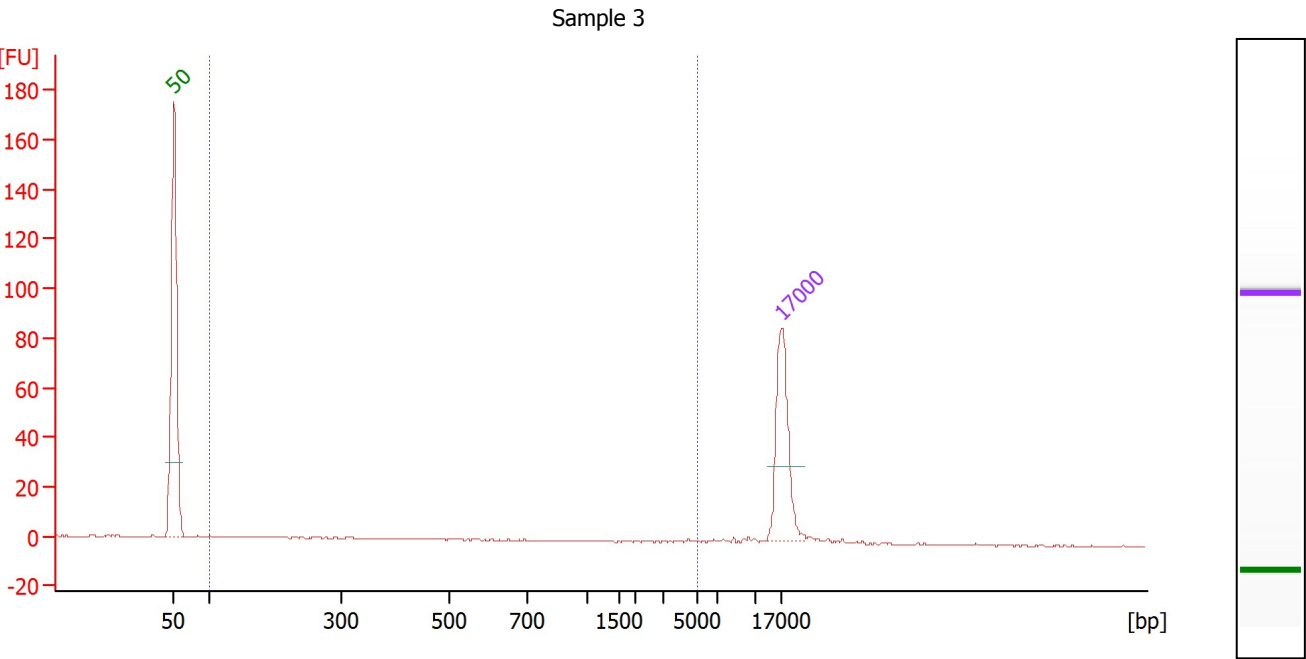
Assay Class: DNA 12000

Data Path: C:\...ta\2023-05-23\Bioanalyzer\_DNA 12000\_2023-05-23\_15-24-15.xad

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Electropherogram Summary Continued ...



Overall Results for sample 3 : Sample 3

Number of peaks found: 0

Area 1: 8.5

Region table for sample 3 : Sample 3

From [bp]	To [bp]	Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [ng/μl]	Col or
100	5,000	8.5	41	870	100.0	0.66	<div></div>

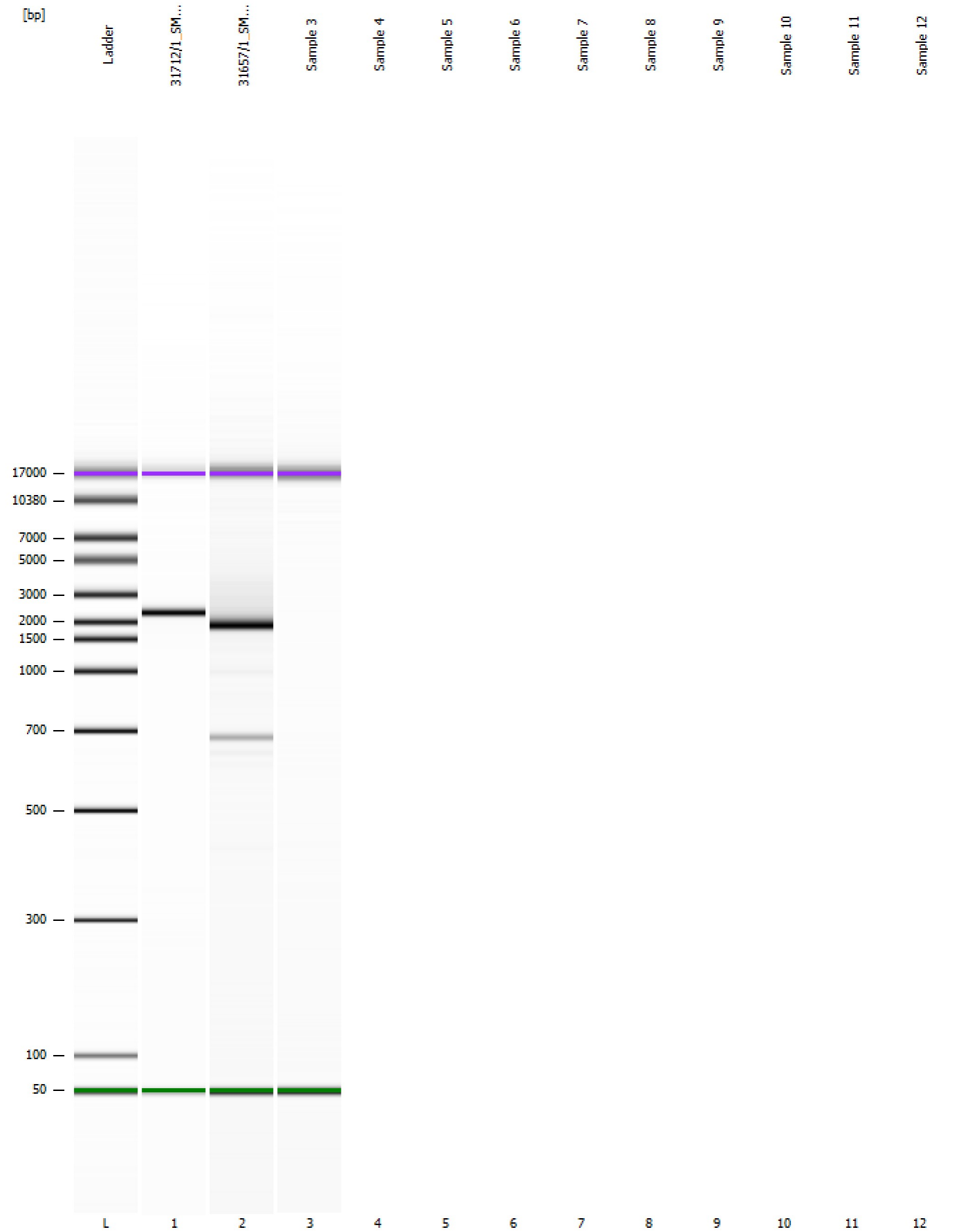
Assay Class: DNA 12000

Data Path: C:\...ta\2023-05-23\Bioanalyzer\_DNA 12000\_2023-05-23\_15-24-15.xad

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Gel Image





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**Invalid Samples**

Sample 4 has not been run, no results available.

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.

Sample 12 has not been run, no results available.

Assay Class: DNA 12000  
Data Path: C:\...ta\2023-05-23\Bioanalzyer\_DNA 12000\_2023-05-23\_15-24-15.xad

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# Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 2 (Number of wells acquired: 4)		Instrument	Run		5/23/2023 3:39:47 PM	(GMT +02:00) W. Europe Standard Time	transcriptomics	FGCZ-NET
Run started on port 2 (File: C:\Program Files (x86)\Agilent\2100 bioanalzyer\2100 expert\data\2023-05-23\Bioanalzyer_DNA 12000_2023-05-23_15-24-15.xad)		Instrument	Run		5/23/2023 3:24:20 PM	(GMT +02:00) W. Europe Standard Time	transcriptomics	FGCZ-NET
Product Number : G2938C		Instrument	Run		5/23/2023 3:24:20 PM	(GMT +02:00) W. Europe Standard Time	transcriptomics	FGCZ-NET
Name :		Instrument	Run		5/23/2023 3:24:20 PM	(GMT +02:00) W. Europe Standard Time	transcriptomics	FGCZ-NET
Vendor : Agilent Technologies		Instrument	Run		5/23/2023 3:24:20 PM	(GMT +02:00) W. Europe Standard Time	transcriptomics	FGCZ-NET
Serial# : DE34903176		Instrument	Run		5/23/2023 3:24:20 PM	(GMT +02:00) W. Europe Standard Time	transcriptomics	FGCZ-NET
Firmware : C.01.069		Instrument	Run		5/23/2023 3:24:20 PM	(GMT +02:00) W. Europe Standard Time	transcriptomics	FGCZ-NET
Cartridge : Electrode		Instrument	Run		5/23/2023 3:24:20 PM	(GMT +02:00) W. Europe Standard Time	transcriptomics	FGCZ-NET