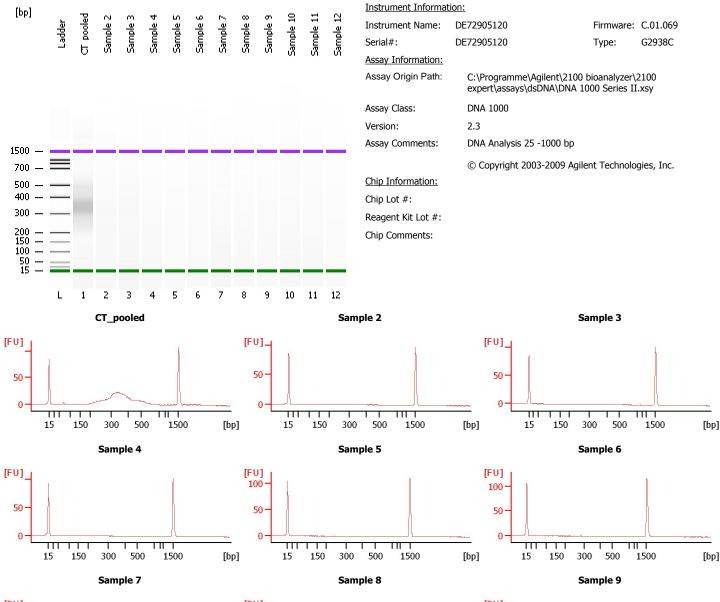
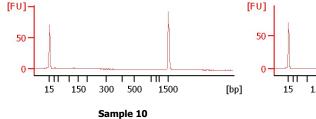
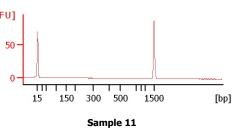
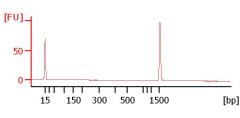
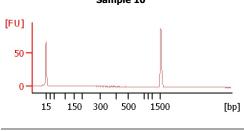
Electrophoresis File Run Summary

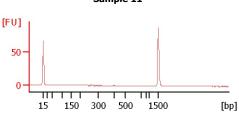


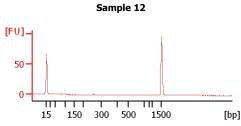












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Assay Class: DNA 1000 Created: 02.08.2024 10:10:45 Data Path: C:\...-02\2100 expert_DNA 1000_DE72905120_2024-08-02_10-10-45.xad Modified: 02.08.2024 11:02:35

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Statu Observation s	Result Label	Result Color
CT_pooled			~		
Sample 2			✓		
Sample 3			✓		
Sample 4			✓		
Sample 5			✓		
Sample 6			✓		
Sample 7			✓		
Sample 8			✓		
Sample 9			✓		
Sample 10			✓		
Sample 11			✓		
Sample 12			~		
Chip Lot #			R	eagent Kit Lot #	

Chip Comments:

Electrophoresis Assay Details

General Analysis Settings

Number of Available Sample and Ladder Wells (Max.): 13

Minimum Visible Range [s]: 30

Maximum Visible Range [s]: 129

Start Analysis Time Range [s]: 30

End Analysis Time Range [s]: 128,95

Ladder Concentration [ng/µl]: 44

Uses Standard Area for Ladder Fragments

Lower Marker Concentration [ng/µl]: 4,2

Upper Marker Concentration [ng/µl]: 2,1

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]: 30 Integration End Time [s]: 128,95

Slope Threshold: 0,5 Height Threshold [FU]: 20 Area Threshold: 0,1 Width Threshold [s]: 0,5 Baseline Plateau [s]: 0,5

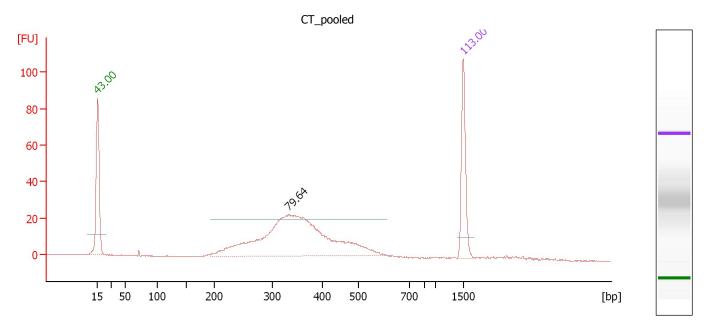
Filter Settings

Filter Width [s]: 0,5 Polynomial Order: 4

Ladder

Ladder Peak	Size	Area
1	15	25
2	25	26
3	50	34
4	100	41
5	150	45
6	200	52
7	300	63
8	400	76
9	500	83
10	700	88
11	850	86
12	1000	90
13	1500	52

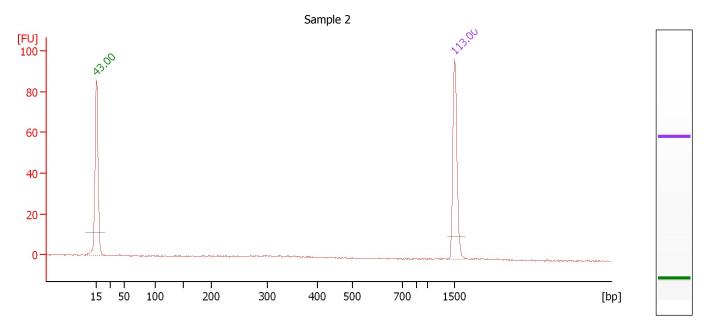
Electropherogram Summary



Overall Results for sample 1: <u>CT pooled</u>

Peak table for sample 1: <u>CT_pooled</u>				CT_pooled		
P	eak		Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1		€ .	15	4,20	424,2	Lower Marker
2			333	10,07	45,8	
3		•	1.500	2.10	2.1	Upper Marker

Electropherogram Summary Continued ...



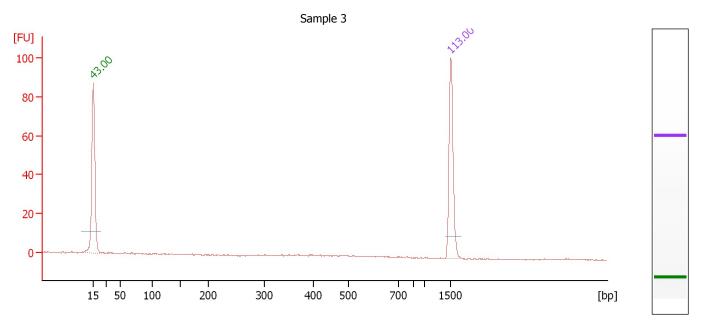
Overall Results for sample 2 : <u>Sample 2</u>

Peak table for sample 2:		for sample 2:	Sample 2		
Peak		Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1	4	15	4,20	424,2	Lower Marker
2		1.500	2,10	2,1	Upper Marker

Assay Class: DNA 1000 C:\...-02\2100 expert_DNA 1000_DE72905120_2024-08-02_10-10-45.xad N

Created: 02.08.2024 10:10:45 Modified: 02.08.2024 11:02:35

Electropherogram Summary Continued ...



Overall Results for sample 3: Sample 3

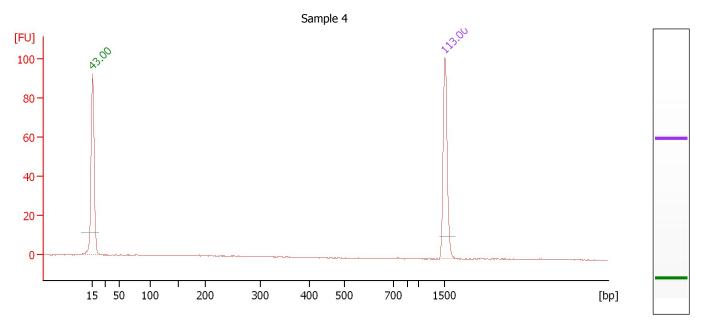
Peak table for sample 3:		for sample 3:	Sample 3		
Peak		Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1	4	15	4,20	424,2	Lower Marker
2		1.500	2,10	2,1	Upper Marker

Assay Class: DNA 1000

Data Path: C:\...-02\2100 expert_DNA 1000_DE72905120_2024-08-02_10-10-45.xad

Created: Modified: 02.08.2024 10:10:45 02.08.2024 11:02:35

Electropherogram Summary Continued ...



Overall Results for sample 4: Sample 4

Number of peaks found:

Peak table for sample 4:

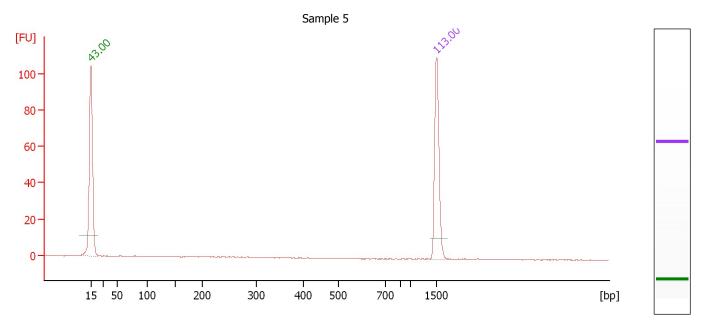
Sample 4

Peak		Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1	4	15	4,20	424,2	Lower Marker
2		1.500	2,10	2,1	Upper Marker

Assay Class: DNA 1000 C:\...-02\2100 expert_DNA 1000_DE72905120_2024-08-02_10-10-45.xad

Created: 02.08.2024 10:10:45 Modified: 02.08.2024 11:02:35

Electropherogram Summary Continued ...



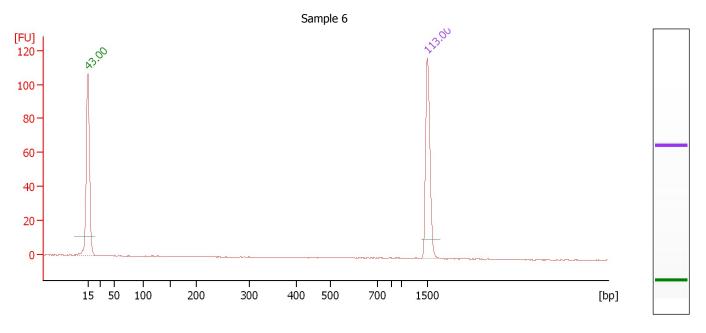
Overall Results for sample 5 : <u>Sample 5</u>

Peak table for sample 5:		for sample 5:	Sample 5		
Peak		Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1	4	15	4,20	424,2	Lower Marker
2		1.500	2,10	2,1	Upper Marker

Assay Class: DNA 1000
Data Path: C:\...-02\2100 expert_DNA 1000_DE72905120_2024-08-02_10-10-45.xad

Created: 02.08.2024 10:10:45 Modified: 02.08.2024 11:02:35

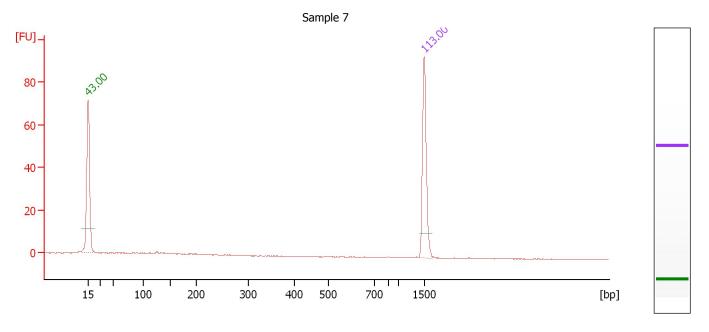
Electropherogram Summary Continued ...



Overall Results for sample 6: Sample 6

Peak table for sample 6:		for sample 6:	Sample 6		
Peak		Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1	4	15	4,20	424,2	Lower Marker
2		1.500	2,10	2,1	Upper Marker

Electropherogram Summary Continued ...



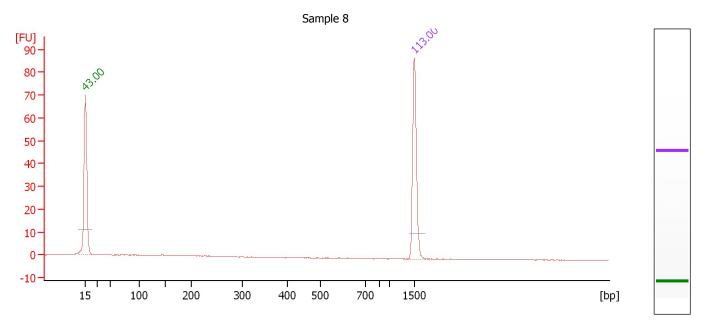
Overall Results for sample 7: Sample 7

Number of peaks found:

Dook table for sample 7

Peak table	e for Sample 7	: <u>Sampie 7</u>			
Peak	Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations	
1	15	4,20	424,2	Lower Marker	
2	1.500	2,10	2,1	Upper Marker	

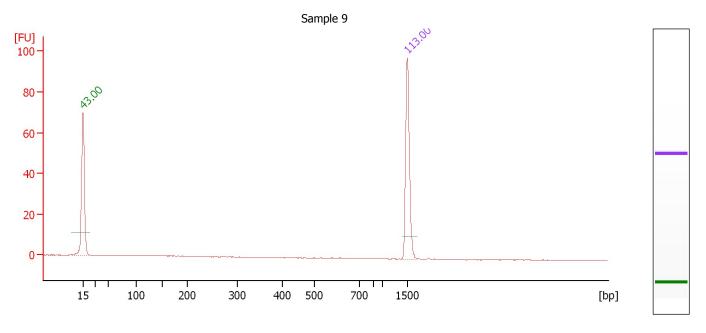
Electropherogram Summary Continued ...



Overall Results for sample 8: Sample 8

Peak table for sample 8:		for sample 8:	Sample 8		
Peak		Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1	•	15	4,20	424,2	Lower Marker
2		1.500	2,10	2,1	Upper Marker

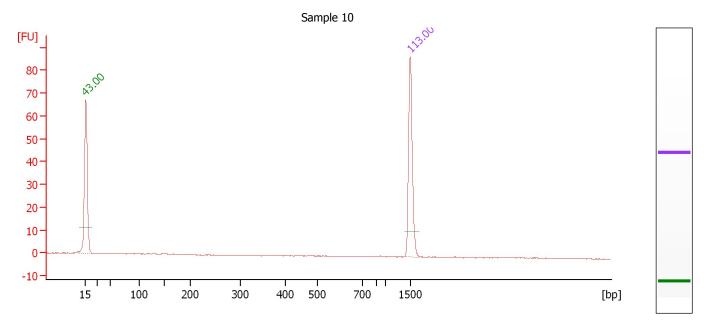
Electropherogram Summary Continued ...



Overall Results for sample 9: Sample 9

Peak table for sample 9:		for sample 9:	Sample 9		
Peak		Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1	4	15	4,20	424,2	Lower Marker
2		1.500	2,10	2,1	Upper Marker

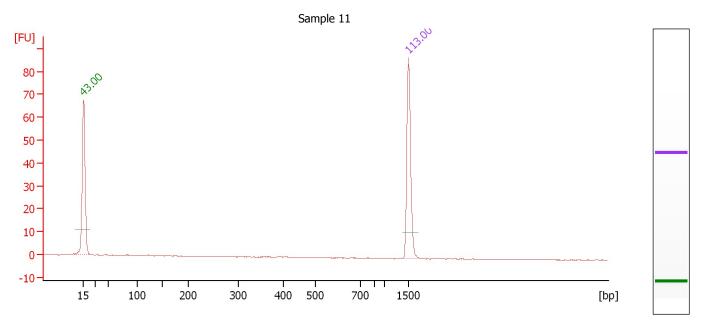
Electropherogram Summary Continued ...



Overall Results for sample 10 : Sample 10

Peak table for sample 10 : <u>Sample 10</u>						
Peak		Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations	
1	4	15	4,20	424,2	Lower Marker	
2		1.500	2,10	2,1	Upper Marker	

Electropherogram Summary Continued ...



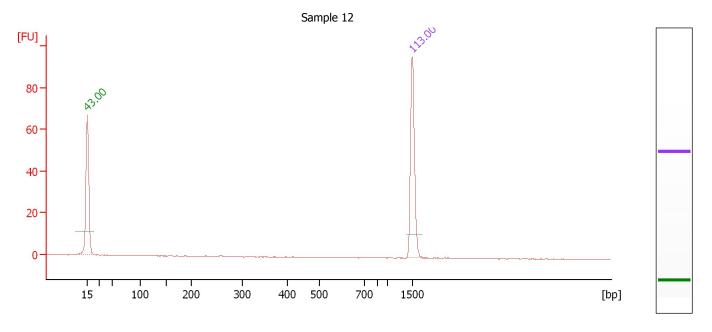
Overall Results for sample 11: Sample 11

Number of peaks found:

Dook table for cample 11

reak table for Sample 11: Sample 11							
Peak		Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations		
1	∢ .	15	4,20	424,2	Lower Marker		
2		1.500	2,10	2,1	Upper Marker		

Electropherogram Summary Continued ...



Overall Results for sample 12 : Sample 12

Number of peaks found:

Dook table for cample 12

reak table for Sample 12: Sample 12							
Peak	Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations			
1	15	4,20	424,2	Lower Marker			
2	1.500	2,10	2,1	Upper Marker			

02.08.2024 10:10:45 02.08.2024 11:02:35 Assay Class: DNA 1000 Created: C:\...-02\2100 expert_DNA 1000_DE72905120_2024-08-02_10-10-45.xad Modified: Data Path: **Gel Image** [bp] Sample 12 Sample 2 CT pooled Ladder 1500 — 1000 850 **—** 700 — 500 — 400 -300 **—** 200 — 150 - • 100 — 50 **—** 25 — L 1 2 3 4 5 6 7 8 9 10 11 12

Run Logbook

Description	Number	Source	Category	Sub Category	Time Stamp	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 13)		Instrument	Run		02.08.2024 10:52:32	(GMT +02:00) Westeuropäisch e Normalzeit	bq_egerstmann	AD
Run started on port 1 (File: C:\Programme\ Agilent\2100 bioanalyzer\210 0 expert\Data\20 4-08-02\2100 expert_DNA 1000_DE72905 20_2024-08-02 10-10-45.xad)	1	Instrument	Run		02.08.2024 10:10:51	(GMT +02:00) Westeuropäisch e Normalzeit	bq_egerstmann	AD
Product Numbe : G2938C	r	Instrument	Run		02.08.2024 10:10:51	(GMT +02:00) Westeuropäisch e Normalzeit	bq_egerstmann	AD
Name :		Instrument	Run		02.08.2024 10:10:51	(GMT +02:00) Westeuropäisch e Normalzeit	bq_egerstmann	AD
Vendor : Agilen Technologies	t	Instrument	Run		02.08.2024 10:10:51	(GMT +02:00) Westeuropäisch e Normalzeit	bq_egerstmann	AD
Serial# : DE72905120		Instrument	Run		02.08.2024 10:10:51	(GMT +02:00) Westeuropäisch e Normalzeit	bq_egerstmann	AD
Firmware : C.01.069		Instrument	Run		02.08.2024 10:10:51	(GMT +02:00) Westeuropäisch e Normalzeit	bq_egerstmann	AD
Cartridge : Electrode		Instrument	Run		02.08.2024 10:10:51	(GMT +02:00) Westeuropäisch e Normalzeit	bq_egerstmann	AD