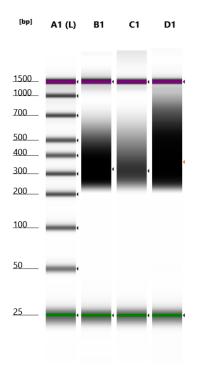
Filename: 2021-07-23 - 13.52.17.D1000



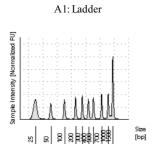
Default image (Contrast 100%)

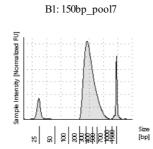
Sample Info

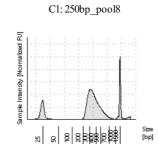
Well	Conc. [ng/µl]	Sample Description	Alert	Observations
A1	21.4	Ladder		Ladder
B1	59.0	150bp_pool7		
Cl	25.4	250bp_pool8		
D1	63.6	350bp pool9		

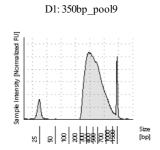
Compact Peak Table

Well	Sample Description	Size [bp]	Calibrated Conc. [ng/µl]	Assigned Conc. [ng/µl]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
A1	Ladder	25	6.47	•	398	•		Lower Marker
A1	Ladder	50	2.53	-	77.9	11.85		
A1	Ladder	100	2.64	-	40.7	12.37		
A1	Ladder	200	2.69	-	20.7	12.61		
A1	Ladder	300	2.74	-	14.0	12.82		
A1	Ladder	400	2.58	-	9.90	12.06		
A1	Ladder	500	2.70	-	8.32	12.66		
A1	Ladder	700	2.69	-	5.91	12.60		
A1	Ladder	1000	2.79	-	4.29	13.04		
A1	Ladder	1500	6.50	6.50	6.67	-		Upper Marker
B1	150bp_pool7	25	6.04	-	372	-		Lower Marker
B1	150bp_pool7	326	59.0	-	279	100.00		
B1	150bp_pool7	1500	6.50	6.50	6.67	-		Upper Marker
C1	250bp_pool8	25	6.37	-	392	-		Lower Marker
C1	250bp_pool8	317	25.4	-	123	100.00		
C1	250bp_pool8	1500	6.50	6.50	6.67	-		Upper Marker
D1	350bp_pool9	25	6.12	-	377	-		Lower Marker
D1	350bp_pool9	365	63.6	-	268	100.00		
D1	350bp_pool9	1500	6.50	6.50	6.67	-		Upper Marker

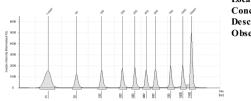








Peak: A1: Ladder



A1 Location Concentration Description Observations

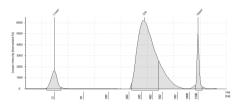
21.4 Ladder Ladder

Size [bp]	Calibrated Conc. [ng/µl]	Assigned Conc. [ng/µl]	PeakMolarity [nmol/l]	% Integrated Area	PeakComment	Observations
25	6.47	-	398	-		Lower Marker
50	2.53	-	77.9	11.85		
100	2.64	-	40.7	12.37		
200	2.69	-	20.7	12.61		
300	2.74	-	14.0	12.82		
400	2.58	-	9.90	12.06		
500	2.70	-	8.32	12.66		
700	2.69	-	5.91	12.60		
1000	2.79	-	4.29	13.04		
1500	6.50	6.50	6.67	-		Upper Marker

Peak: B1: 150bp_pool7

1500

Peak: D1: 350bp_pool9



6.50

B1 Location 59.0 Concentration 150bp_pool7 Description Observations

Calibrated Conc. Assigned Conc. PeakMolarity % Integrated Area Size [bp] PeakComment Observations [nmol/I] 372 [**ng/µl**] Lower Marker 326 59.0 279 100.00 6.50 Upper Marker

6.67

Peak: C1: 250bp_pool8



Location Concentration Description Observations

C1 25.4 250bp_pool8

Location

Concentration Description Observations

D1 63.6 350bp pool9

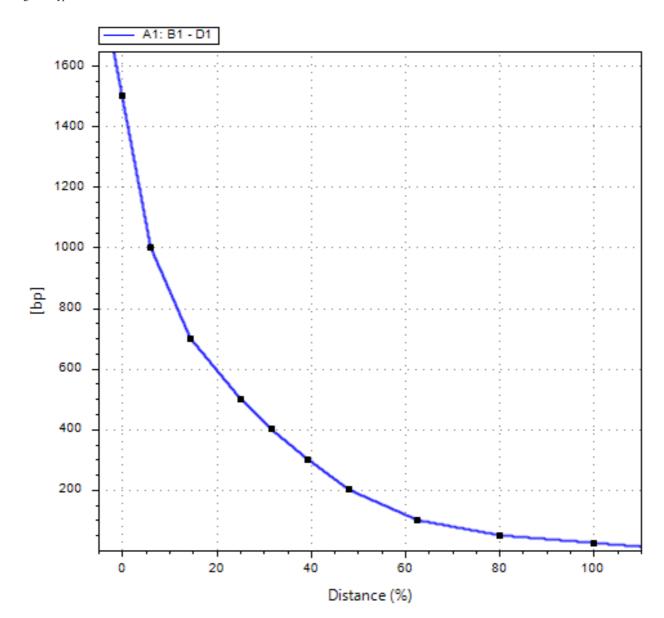
Calibrated Conc. Assigned Conc. PeakMolarity Size [bp] % Integrated Area PeakComment Observations [nmol/l] 25 317 1500 392 123 6.67 Lower Marker 6.37 25.4 100.00 Upper Marker

Size [bp]	Calibrated Conc. [ng/µl]	Assigned Conc. [ng/µl]	PeakMolarity [nmol/l]	% Integrated Area	PeakComment	Observations
25	6.12	-	377	-		Lower Marker
365	63.6	-	268	100.00		
1500	6.50	6.50	6.67	-		Unner Marker

Calibration

Molecular Weight Settings

Fitting type: Piecewise Interpolation Alignment type: Between markers



Concentration Settings

Calibration mode: Upper Marker Normalise peaks from: Upper Marker Fitting type: Linear Regression

Experiment Properties

Run Properties

Analysis Software Version 3.2.0.37131

 $C. Users admin Documents Agilent \ Tape Station\ Data \ 2021-07-23 \ 2021-07-23 \ -\ 13.52.17. D1000$ **Filename**

D1000 Assay

23-Jul-2021 1:59 PM 3.2.0.37131 Run End Date

Last Saved Under Version

Study Comments

ScreenTape Device 1

admin Username

Screen Tape Device ID 01-8029-210531-01-003595

Expiry Date ScreenTape Device History 04-Aug-2021

First run 21-Jul-2021, 2 run(s) performed

Temperature [°C] 24.1 Electrophoresis Time [s] 151

Instrument Type 4200 TapeStation Instrument Serial Number DEDAA01963

Notes

ScreenTape Device Run Date

Reagent Lot # ScreenTape Lot# 23-Jul-2021 1:52 PM

Controller Environment

DESKTOP-NBEV0I6 Computer Instrument Controller Software Version 3.2.0.37131 First Run Analysis Version 3.2.0.37131

Microsoft Windows 10 Pro Operating System

Generated: 23-Jul-2021