D1000 ScreenTape® Page 1 of 6

# $Filename: 2020-09-24-01.D1000\_i5\_L1LotterhosLab\_shortAdapter.D1000$

# Gel Image



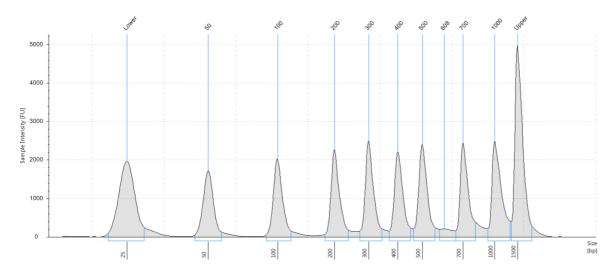
Default image (Contrast 100%), Image is Scaled to view larger Molecular Weight range

#### Sample Info

Well	Conc. [ng/µl]	Sample Description	Alert	Observations
Al	26.9	Ladder	<u> </u>	Caution! Expired ScreenTape device; Issue with ladder peak detection (too many peaks detected); Ladder
B1	2.54	Zymo i5	<u> </u>	Caution! Expired ScreenTape device
Cl	8.32	short custom i5	<u> </u>	Caution! Expired ScreenTape device
D1		blank	<u> </u>	Marker(s) not detected; Caution! Expired ScreenTape device

D1000 ScreenTape® Page 2 of 6

# A1: Ladder



# Sample Table

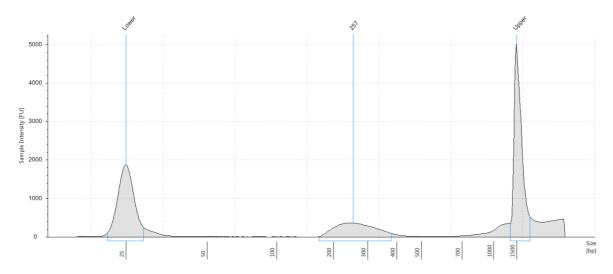
Well	Conc. [ng/µl]	Sample Description	Alert	Observations
Al	26.9	Ladder	<u>^</u>	Caution! Expired ScreenTape device; Issue with ladder peak detection (too many peaks detected); Ladder

#### Peak Table

Size [bp]	Calibrated Conc. [ng/µl]	Assigned Conc. [ng/µl]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	5.32	-	327	-		Lower Marker
50	2.97	-	91.5	11.06		
100	3.12	-	48.0	11.60		
200	3.17	-	24.4	11.81		
300	3.39	-	17.4	12.63		
400	3.18	-	12.2	11.82		
500	3.45	-	10.6	12.83		
608	0.444	-	1.12	1.65		
700	3.29	-	7.23	12.24		
1000	3.86	-	5.94	14.36		
1500	6.50	6.50	6.67	-		Upper Marker

D1000 ScreenTape® Page 3 of 6

# B1: Zymo i5



# Sample Table

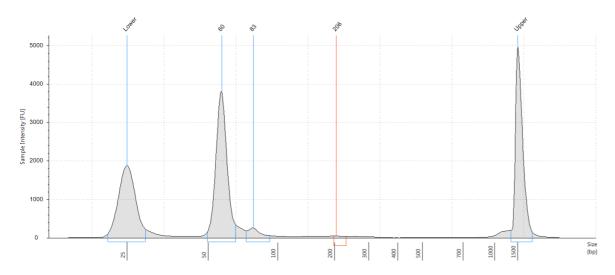
Well	Conc. [ng/µl]	Sample Description	Alert	Observations
Bl	2.54	Zymo i5		Caution! Expired ScreenTape device

#### Peak Table

Size [bp]	Calibrated Conc. [ng/µl]	Assigned Conc. [ng/µl]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
25	5.18	-	319	-		Lower Marker
257	2.54	-	15.2	100.00		
1500	6.50	6.50	6.67	-		Upper Marker

D1000 ScreenTape® Page 4 of 6

# C1: short custom i5



# Sample Table

Well	Conc. [ng/µl]	Sample Description	Alert	Observations
Cl	8.32	short custom i5		Caution! Expired ScreenTape device

#### Peak Table

Siz	e [bp]	Calibrated Conc. [ng/µl]	Assigned Conc. [ng/µl]	Peak Molarity [nmol/l]	% Integrated Area	Peak Comment	Observations
	25	5.80	-	357	-		Lower Marker
	60	7.65	-	197	92.01		
	83	0.582	-	10.9	7.00		
2	206	0.0825	-	0.616	0.99		
1	500	6.50	6.50	6.67	-		Upper Marker

D1000 ScreenTape® Page 5 of 6

# D1: blank



# Sample Table

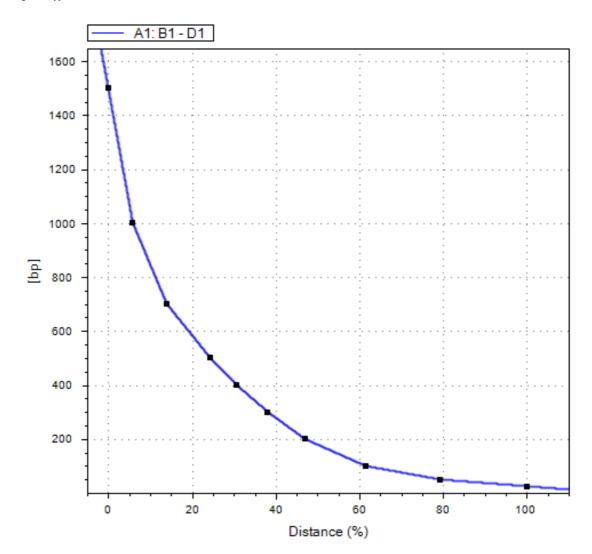
Well	Conc. [ng/µl]	Sample Description	Alert	Observations
D1		blank	<u> </u>	Marker(s) not detected; Caution! Expired ScreenTape device

D1000 ScreenTape® Page 6 of 6

#### 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