

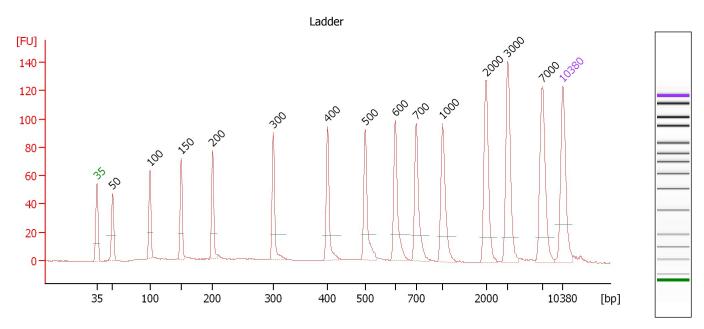
2017-09-22_	001.xad					Pag	ge 2	of	8
Assay Class: Data Path: Electrophore	High Sensitivity DNA Assay C:\ bioanalyzer\2100 expert\data\20 esis File Run Summary (Chip Sum	-	'-09-22_	001.xad	Created: Modified:	9/22/20 9/22/20			
Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result L	abel	Res Colo		
sample 1 sample 2 sample 3 sample 4 sample 5 sample 6 sample 7 sample 8 sample 9 sample 10 sample 11 Ladder			* * *						
Chip Lot #			Reage	ent Kit Lot #					

Chip Comments :

Created: Modified:

# 9/22/2017 1:01:56 PM 9/22/2017 1:20:24 PM

## **Electropherogram Summary**



# **Overall Results for Ladder**

Noise:

0.1

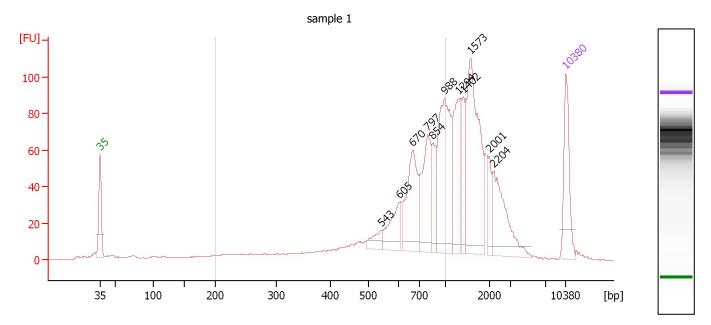
# Peak table for Ladder

Pea k	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	50	150.00	4,545.5	Ladder Peak	45.35
3	100	150.00	2,272.7	Ladder Peak	50.94
4	150	150.00	1,515.2	Ladder Peak	55.63
5	200	150.00	1,136.4	Ladder Peak	60.38
6	300	150.00	757.6	Ladder Peak	69.51
7	400	150.00	568.2	Ladder Peak	77.60
8	500	150.00	454.5	Ladder Peak	83.29
9	600	150.00	378.8	Ladder Peak	87.79
10	700	150.00	324.7	Ladder Peak	90.93
11	1,000	150.00	227.3	Ladder Peak	94.93
12	2,000	150.00	113.6	Ladder Peak	101.47
13	3,000	150.00	75.8	Ladder Peak	104.71
14	7,000	150.00	32.5	Ladder Peak	109.95
15	10,380	75.00	10.9	Upper Marker	113.00

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# **Electropherogram Summary Continued ...**



Overall Results for sample	± .	Sample I		
Number of peaks found:	11		Corr. Area 1:	662.0
Noise:	0.2			

comple 1

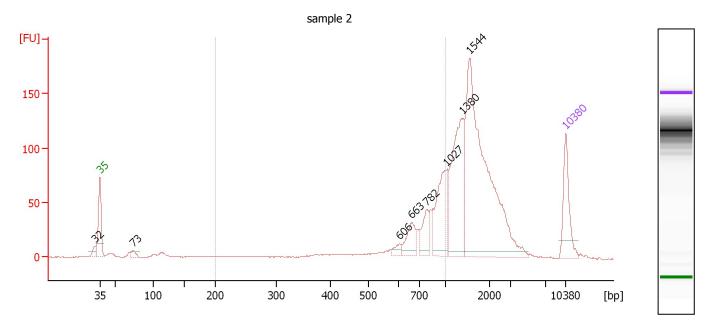
#### Peak table for sample 1 : sample 1

Overall Recults for cample 1

Pea k		Size [bp]	Conc. [pg	/µI]	Molarity [pmol/l]	Observatio	ons	Align	ed Migration Time [s]
1		35	125.00		5,411.3	Lower Mark	er	43.00	
2		543	24.69		68.9			85.21	
3		605	62.73		157.0			87.96	
4		670	137.70		311.4			89.99	
5		797	118.70		225.6			92.22	
6		854	49.71		88.2			92.99	
7		988	197.04		302.2			94.76	
8		1,294	110.76		129.7			96.85	
9		1,402	56.58		61.1			97.56	
10		1,573	222.91		214.7			98.67	
11		2,001	30.07		22.8			101.47	7
12		2,204	100.10		68.8			102.13	3
13		10,380	75.00		10.9	Upper Mark	er	113.00	)
Regi	on t	able for san	nple 1 :	sam	ple 1				
<b>From</b> [ <b>bp</b> ] 200	1	To [bp] Ave [bp 1,000 686	]	<b>Corr.</b> <b>Area</b> 662.0	<b>Molarity</b> [pmol/l] 2,061.8	Co Conc. lor [pg/µl] 773.91	<b>% of</b> Tota 46	1 (	Size distribution in CV [%] 28.7
200		1,000 000		002.0	2,001.0	// 5.91	10	4	20.7

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# **Electropherogram Summary Continued ...**

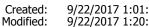


#### Overall Results for sample 2 : sample 2

Number of peaks found:	8	Corr. Area 1:	385.3
Noise:	0.2		

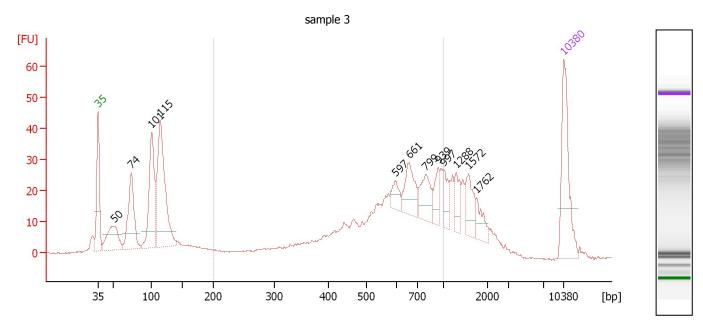
#### Peak table for sample 2 : sample 2

1 32 0.00 0.0 42.23   2 35 125.00 5,411.3 Lower Marker 43.00   3 73 15.84 326.6 47.97   4 606 12.99 32.5 87.97   5 663 50.10 114.5 89.77	on Time [s]
3 73 15.84 326.6 47.97   4 606 12.99 32.5 87.97   5 663 50.10 114.5 89.77	
4   606   12.99   32.5   87.97     5   663   50.10   114.5   89.77	
5 663 50.10 114.5 89.77	
C 702 40.01 0C F 02.02	
6 782 49.81 96.5 92.03	
7 1,027 135.76 200.3 95.10	
8 1,380 222.63 244.5 97.41	
9 1,544 518.80 508.9 98.49	
10 🕨 10,380 75.00 10.9 Upper Marker 113.00	
Region table for sample 2 : <u>sample 2</u>	
From   To [bp]   Average Size   Corr.   Molarity   Co Conc.   % of   Size distribution     [bp]   [bp]   Area   [pmol/I]   lor [pg/μ]   Total   CV [%]     200   1,000   745   385.3   873.0   365.35   27   25.7	ution in



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# **Electropherogram Summary Continued ...**



<b>Overall Results for sam</b>	ple 3 :	sample 3		
Number of peaks found:	12		Corr. Area 1:	411.6
Noise:	0.2			

#### Peak table for sample 3 : sample 3

Pea k		Size [bp]	Conc. [pg	/µI]	Molarity [pmol/l	] Observatio	ns A	ligned Migration Time [s]
1		35	125.00		5,411.3	Lower Marke	er 4	13.00
2		50	75.30		2,268.4		4	15.38
3		74	115.73		2,373.0		4	18.02
4		101	149.94		2,240.3		5	51.07
5		115	227.62		2,994.0		5	52.36
6		597	18.57		47.1		8	37.65
7		661	45.80		104.9		8	39.72
8		799	39.21		74.3		ç	92.25
9		939	21.67		35.0		ç	94.12
10		997	21.63		32.9		ç	94.89
11		1,288	21.51		25.3		ç	96.81
12		1,572	31.63		30.5		ç	98.67
13		1,762	19.11		16.4		ç	99.91
14		10,380	75.00		10.9	Upper Marke	er 1	.13.00
Regi	on t	able for san	nple 3 :	<u>sam</u>	<u>ple 3</u>			
<b>From</b> [ <b>bp]</b> 200	1	<b>To [bp] Ave</b> [bp 1,000 618	]	<b>Corr.</b> <b>Area</b> 411.6	<b>Molarity</b> [pmol/l] 1,714.5	<b>Co Conc.</b> lor [pg/μl] 598.05	<b>% of</b> Total 45	Size distribution in CV [%] 30.0

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Assay Class: Data Path: <b>Gel Image</b>	підп : C:\	bioanalyze	DNA Assay er\2100 exp	ert\data\20	17-09-22\2	2017-09-22	_001.xad		M	reated: odified:	9/22/2017 9/22/2017	1:20:24 PM
[bp]	Ladder	sample 1	sample 2	sample 3	sample 4	sample 5	sample 6	sample 7	sample 8	sample 9	sample 10	sample 11
10380 —	_											
3000 — — 2000 — —	=	-										
1000 —	_		-									
600 — <b>—</b>	_											
400 — —	_											
300 — —	_											
200 —												
150 — — 100 — —				_								
50 — 35 — —		_										
	L	1	2	3	4	5	6	7	8	9	10	11

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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.