2018-01-31_	_003.xad			I	Page	1 0	f	9
Assay Class: Data Path:	High Sensitivity DNA Assay C:\ bioanalyzer\2100 expert\data\2018-01-31\2018-	01-31_003.xad	Crea Modi			3:21:0 3:42:1		
Electrophore	esis File Run Summary							
		Instrument Informat	tion:					
		Instrument Name:	DE34903152	Firmware	: C.01	.069		
		Serial#:	DE34903152	Type:	G29	38C		
		Assay Information:						
		Assay Origin Path:	C:\Program Files\Agilen expert\assays\dsDNA\H					

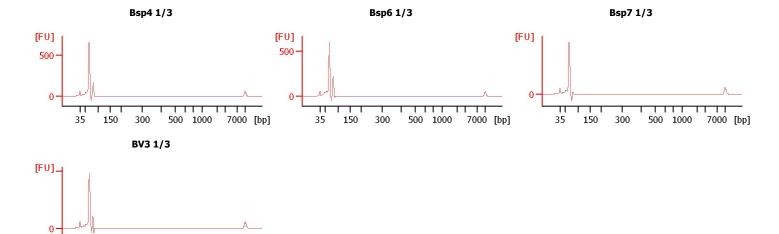
Assay Class:

Chip Information: Chip Lot #: Reagent Kit Lot #: Chip Comments:

Version: Assay Comments: High Sensitivity DNA Assay

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1.03



11 35

Т

150

300

Т

500 1000 7000 [bp]

Assay Class: Data Path:	High Sensitivity DNA Assay C:\ bioanalyzer\2100 expert\data\2018-01-31\2018-0	01-31_003.xad	Created: Modified:		L8 3:21:00 PM L8 3:42:19 PM
Electrophore	esis File Run Summary (Chip Summary)				
Sample Name	Sample Comment Rest. Digest	Status Observation	Result	Label	Result Color
Bsp4 1/3		×			
Bsp6 1/3	$\Box$	×			
Bsp7 1/3	$\Box$	×			
BV3 1/3	$\Box$	×			
sample 5	$\Box$				
sample 6	$\Box$				
sample 7	$\Box$				
sample 8	$\Box$				
sample 9	$\Box$				
sample 10	$\Box$				
sample 11	$\Box$				
Ladder	Ō	*			
Chip Lot #		Reagent Kit Lot #			

**Chip Comments :** 

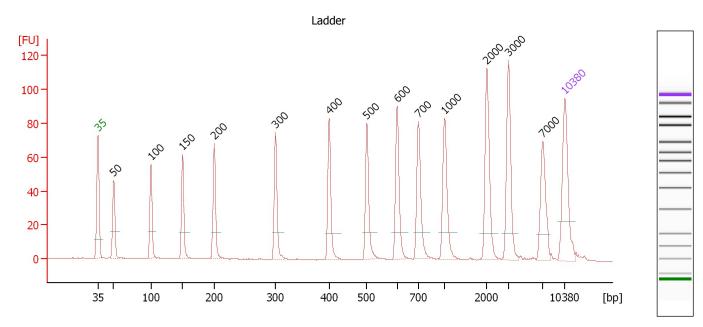
2018-01-31\_003.xad

#### High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-01-31\2018-01-31\_003.xad Assay Class: Data Path:

## 1/31/2018 3:21:00 PM 1/31/2018 3:42:19 PM Created:

# Modified:

#### **Electropherogram Summary**



## **Overall Results for Ladder**

Noise:

0.2

#### Peak table for Ladder

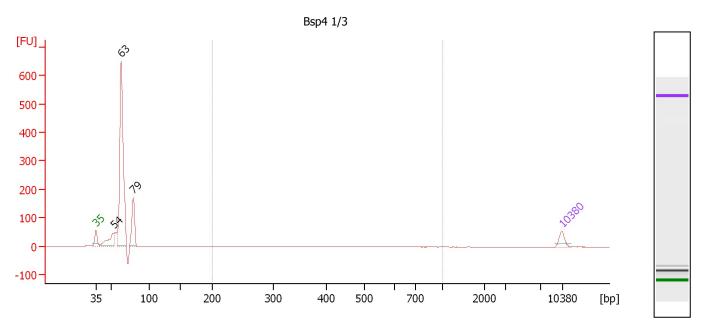
Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1		35	125.00	5,411.3	Lower Marker	43.00
2	L	50	150.00	4,545.5	Ladder Peak	45.33
3		100	150.00	2,272.7	Ladder Peak	50.95
4		150	150.00	1,515.2	Ladder Peak	55.70
5	L	200	150.00	1,136.4	Ladder Peak	60.41
6		300	150.00	757.6	Ladder Peak	69.57
7		400	150.00	568.2	Ladder Peak	77.62
8	L	500	150.00	454.5	Ladder Peak	83.29
9		600	150.00	378.8	Ladder Peak	87.84
10		700	150.00	324.7	Ladder Peak	91.03
11		1,000	150.00	227.3	Ladder Peak	94.98
12		2,000	150.00	113.6	Ladder Peak	101.31
13		3,000	150.00	75.8	Ladder Peak	104.55
14		7,000	150.00	32.5	Ladder Peak	109.71
15		10,380	75.00	10.9	Upper Marker	113.00

# Assay Class: High Sensitivity DNA Assay Data Path: C:\... bioanalyzer\2100 expert\data\2018-01-31\2018-01-31\_003.xad

#### Created: 1/31/2018 3 Modified: 1/31/2018 3

## 1/31/2018 3:21:00 PM 1/31/2018 3:42:19 PM

# Electropherogram Summary Continued ...



# Overall Results for sample 1 : <u>Bsp4 1/3</u>

Number of peaks found:	3	Corr. Area 1:	1.1
Noise:	0.2		

Peak table for sample 1	:	<u>Bsp4 1/3</u>
-------------------------	---	-----------------

Pea k		Size [bp]	Conc. [pg	<b>յ/μ</b> ]]	Molarity [pm	ol/l]	Observa	tions	Aligr	ned Migration Time [s]	]
1		35	125.00		5,411.3		Lower Ma	arker	43.00	)	
2		54	387.86		10,930.8				45.75	5	
3		63	2,726.92		65,758.7				46.77	7	
4		79	581.37		11,128.5				48.60	)	
5		10,380	75.00		10.9		Upper Ma	arker	113.0	00	
Regi	on t	able for sar	nple 1:	Bsp4	1/3						
<b>From</b> [bp] 200	I	<b>To [bp] Ave</b> [bp 1,000 614	]	<b>Corr.</b> Area 1.1	<b>Conc.</b> [pg/μl] 2.33	<b>Mola</b> [pmo 6.2	•	<b>Co % o</b> lor Tota		Size distribution in CV [%] 26.6	

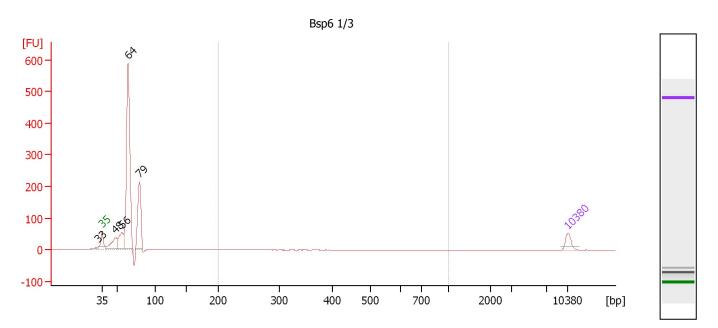
1/31/2018 3:21:00 PM 1/31/2018 3:42:19 PM

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Modified:

# Assay Class: High Sensitivity DNA Assay Data Path: C:\... bioanalyzer\2100 expert\data\2018-01-31\2018-01-31\_003.xad

Electropherogram Summary Continued ...



# Overall Results for sample 2 : <u>Bsp6 1/3</u>

Number of peaks found:	5	Corr. Area 1:	3.1
Noise:	0.2		

# Peak table for sample 2 : <u>Bsp6 1/3</u>

Pea k		Size [bp]	Conc. [pg	<b>յ/μ</b> ]]	Molarity [pmo	ol/l] Obse	rvations	Alig	ned Migration Time [s]
1		33	0.00		0.0			42.4	4
2		35	125.00		5,411.3	Lower	<sup>-</sup> Marker	43.0	0
3		48	306.93		9,666.2			45.0	3
4		56	337.22		9,125.2			46.0	0
5		64	2,376.44		56,137.2			46.9	2
6		79	821.33		15,736.3			48.6	0
7		10,380	75.00		10.9	Upper	<sup>-</sup> Marker	113.	00
Regi	on t	table for sar	nple 2 :	Bsp6	<u>1/3</u>				
<b>From</b> [bp] 200	1	<b>To [bp] Ave</b> [bp 1,000 399	]	Corr. Area 3.1	<b>Conc. [pg/μl]</b> 7.19	Molarity [pmol/l] 30.5	<b>Co %</b> ( lor Tot		Size distribution in CV [%] 27.4

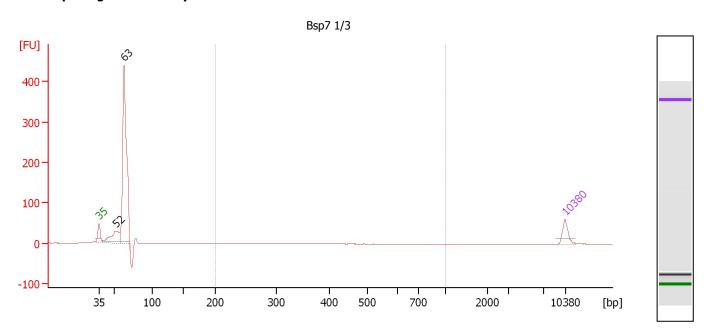
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Modified:

# Assay Class: High Sensitivity DNA Assay Data Path: C:\... bioanalyzer\2100 expert\data\2018-01-31\2018-01-31\_003.xad

# Electropherogram Summary Continued ...



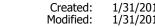
# Overall Results for sample 3 : <u>Bsp7 1/3</u>

Number of peaks found:	2	Corr. Area 1:	0.0
Noise:	0.2		

Peak	tał	ble for samp	le 3 : <u>Bsp7</u>	<u>1/3</u>		
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		52	312.49	9,147.9		45.53
3		63	1,870.02	45,140.4		46.76
4		10,380	75.00	10.9	Upper Marker	113.00
Regi	on t	table for sar	nple 3 : <u>Bs</u> r	<u> 71/3</u>		
Erom			orago Sizo Corr	Conc Mo	larity Co %	of Size distribution in

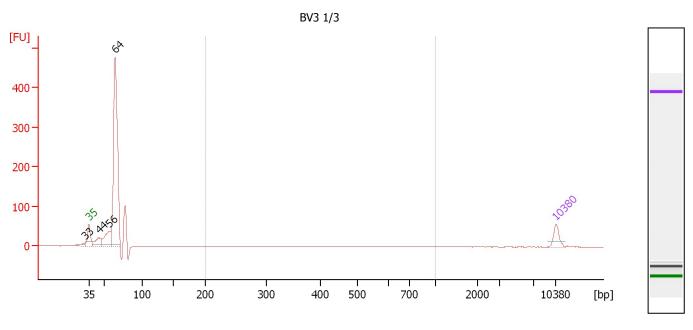
From [bp]	To [bp]	] Average Size [bp]	Corr. Area	Conc. [pg/µl]	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	0	0.0	0.00	0.0	0	0.0

# Assay Class: High Sensitivity DNA Assay Data Path: C:\... bioanalyzer\2100 expert\data\2018-01-31\2018-01-31\_003.xad



## 1/31/2018 3:21:00 PM 1/31/2018 3:42:19 PM

Electropherogram Summary Continued ...



# Overall Results for sample 4 : <u>BV3 1/3</u>

Number of peaks found:	4	Corr. Area 1:	1.0
Noise:	0.1		

Peak table fo	or sample 4	:	BV3 1/3
---------------	-------------	---	---------

Pea k		Size [bp]	Conc. [pg	/µl] N	Iolarity [pmd	ol/I] Obs	ervations	Aligı	ned Migration Time [s]
1		33	0.00	0	0.0			42.37	7
2	•	35	125.00	5	,411.3	Lowe	er Marker	43.00	0
3		44	156.88	5	,344.0			44.47	7
4		56	277.26	7	,447.8			46.05	5
5		64	2,155.82	5	1,126.1			46.89	9
6		10,380	75.00	1	0.9	Uppe	er Marker	113.0	00
Region table for sample 4 : <u>BV3 1/3</u>									
<b>From</b> [bp] 200	1	To [bp] Ave [bp 1,000 550	-	Corr. Area 1.0	<b>Conc. [pg/µl]</b> 1.95	Molarity [pmol/l] 5.6	Co % c lor Tota 0		Size distribution in CV [%] 15.4

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.

Created: 1/31/2018 3:21:00 PM Modified: 1/31/2018 3:42:19 PM