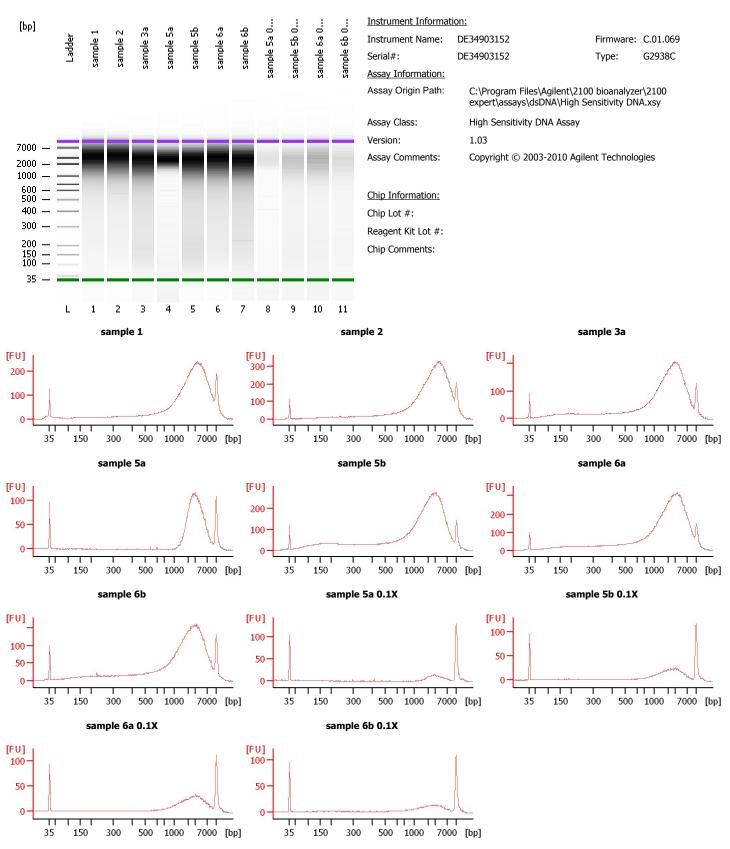
2018-08-31_001.xad Page 1 of 16

Assay Class: High Sensitivity DNA Assay Created: 8/31/2018 11:38:57 AM Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad Created: 8/31/2018 12:33:16 PM

Electrophoresis File Run Summary



2018-08-31_001.xad Page 2 of 16

Assay Class: High Sensitivity DNA Assay Created: 8/31/2018 11:38:57 AM C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad Data Path: Modified: 8/31/2018 12:33:16 PM **Electrophoresis File Run Summary (Chip Summary) Sample Name** Sample Comment Rest. Digest Status Observation **Result Label Result Color** sample 1 sample 2 sample 3a sample 5a sample 5b sample 6a sample 6b

Chip Lot # Reagent Kit Lot #

Chip Comments:

sample 5a 0.1X sample 5b 0.1X sample 6a 0.1X sample 6b 0.1X Ladder

2018-08-31_001.xad Page 3 of 16

Assay Class: High Sensitivity DNA Assay Created: 8/31/2018 11:38:57 AM Data Path: C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad Modified: 8/31/2018 12:33:16 PM

Electrophoresis Assay Details

General Analysis Settings

Number of Available Sample and Ladder Wells (Max.): 12

Minimum Visible Range [s]: 32
Maximum Visible Range [s]: 138
Start Analysis Time Range [s]: 33
End Analysis Time Range [s]: 137.5
Ladder Concentration [pg/µl]: 1950
Uses Standard Area for Ladder Fragments
Lower Marker Concentration [pg/µl]: 125
Upper Marker Concentration [pg/µl]: 75
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]: 33.05
Integration End Time [s]: 137
Slope Threshold: 0.8
Height Threshold [FU]: 90
Area Threshold: 0.1
Width Threshold [s]: 0.6
Baseline Plateau [s]: 0.5

Filter Settings

Filter Width [s]: 0.5 Polynomial Order: 4

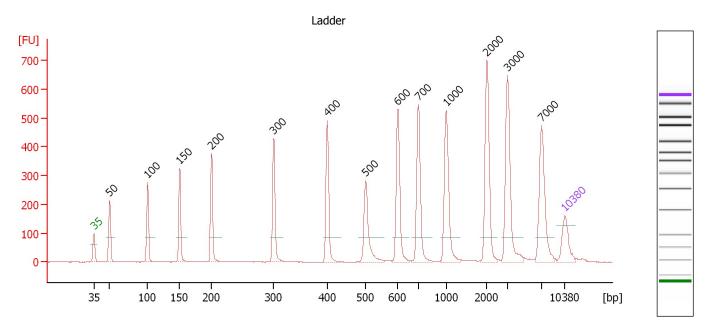
Ladder

Ladder Peak	Size	Area
1	35	160
2	50	210
3	100	208
4	150	221
5	200	242
6	300	270
7	400	305
8	500	306
9	600	336
10	700	321
11	1000	366
12	2000	413
13	3000	411
14	7000	400
15	10380	214

2018-08-31_001.xad Page 4 of 16

Assay Class: High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad S/31/2018 12:33:16 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.3

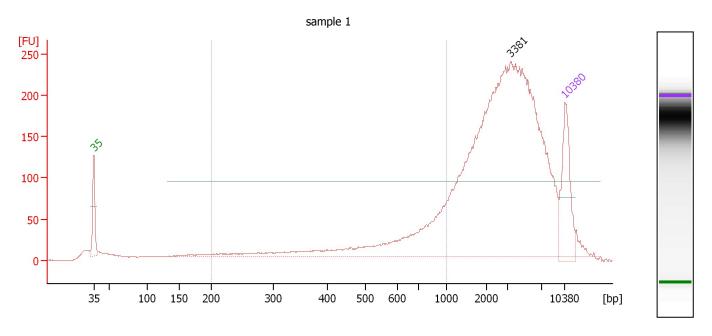
Peak table for Ladder

Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	4	35	125.00	5,411.3	Lower Marker	43.00
2	L	50	150.00	4,545.5	Ladder Peak	45.31
3	L	100	150.00	2,272.7	Ladder Peak	50.95
4		150	150.00	1,515.2	Ladder Peak	55.74
5		200	150.00	1,136.4	Ladder Peak	60.49
6		300	150.00	757.6	Ladder Peak	69.73
7		400	150.00	568.2	Ladder Peak	77.68
8		500	150.00	454.5	Ladder Peak	83.37
9	L	600	150.00	378.8	Ladder Peak	88.16
10		700	150.00	324.7	Ladder Peak	91.23
11	L	1,000	150.00	227.3	Ladder Peak	95.38
12		2,000	150.00	113.6	Ladder Peak	101.41
13		3,000	150.00	75.8	Ladder Peak	104.45
14	L	7,000	150.00	32.5	Ladder Peak	109.49
15		10,380	75.00	10.9	Upper Marker	113.00

2018-08-31_001.xad Page 5 of 16

Assay Class: High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad S/31/2018 12:33:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : sample 1

Number of peaks found: 1 Corr. Area 1: 834.8

Noise: 0.5

Peak table for sample 1 : <u>sample 1</u>

Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	4	35	125.00	5,411.3	Lower Marker	43.00
2		3,381	1,029.18	461.2		104.93
3		10,380	75.00	10.9	Upper Marker	113.00

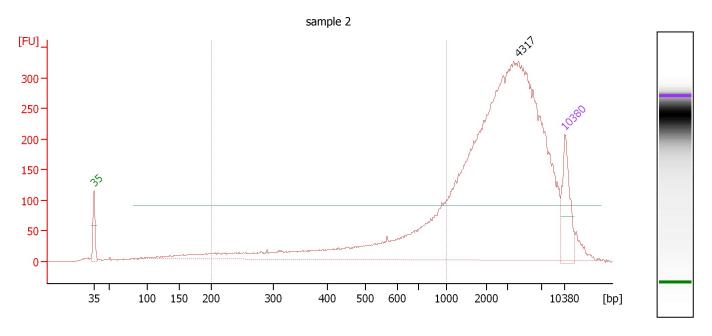
Region table for sample 1 : <u>sample 1</u>

From	To [bp] Average Siz	e Conc.	Corr.	Molarity	Co % of	Size distribution in
[bp]	[bp]	[pg/µl]	Area	[pmol/l]	lor Total	CV [%]
200	1.000 588	339.03	834.8	1.160.3	21	38.9

2018-08-31_001.xad Page 6 of 16

Assay Class: High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad S/31/2018 12:33:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : sample 2

Number of peaks found: 1 Corr. Area 1: 1,291.9

Noise: 0.8

Peak table for sample 2 : <u>sample 2</u>

Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	4	35	125.00	5,411.3	Lower Marker	43.00
2		4,317	1,534.11	538.4		106.11
3		10,380	75.00	10.9	Upper Marker	113.00

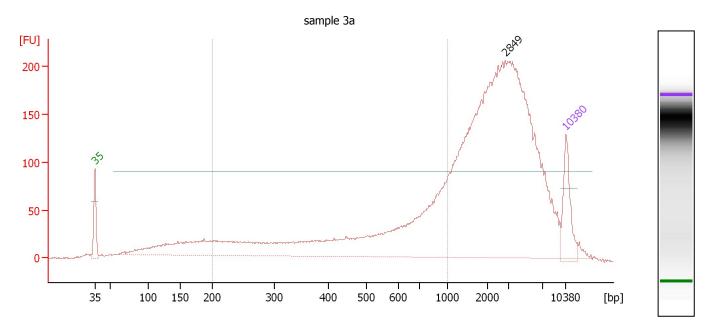
Region table for sample 2: <u>sample 2</u>

From [bp]	To [bp] Avera	age Size Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distributio	n in
[nb]	[pb]	[ba/bi]	Aica	[hilloi/i]		CV [/0]	
200	1.000 580	539.00	1.291.9	1.876.2	24	39.5	

2018-08-31_001.xad Page 7 of 16

Assay Class: High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad S/31/2018 12:33:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 3: sample 3a

Number of peaks found: 1 Corr. Area 1: 1,233.6

Noise: 0.9

Peak table for sample 3: sample 3a

Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	4	35	125.00	5,411.3	Lower Marker	43.00
2		2,849	1,832.49	974.7		103.99
3		10,380	75.00	10.9	Upper Marker	113.00

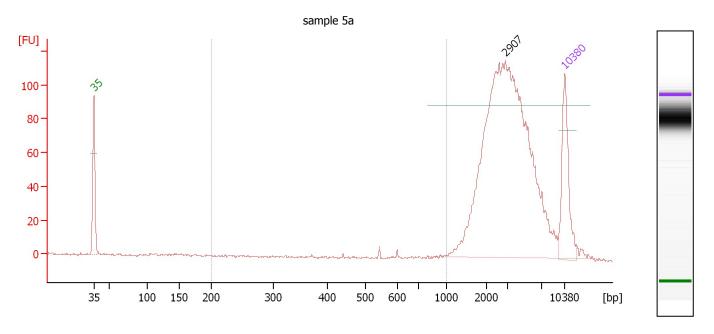
Region table for sample 3: <u>sample 3a</u>

From	To [bp] Average Siz	e Conc.	Corr.	Molarity	Co % of	Size distribution in
[bp]	[bp]	[pg/µl]	Area	[pmol/l]	<u>lor</u> Total	CV [%]
200	1.000 552	889 11	1 233 6	3 314 4	30	42 1

2018-08-31_001.xad Page 8 of 16

Assay Class: High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad S/31/2018 12:33:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 4: <u>sample 5a</u>

Number of peaks found: 1 Corr. Area 1: 16.8

Noise: 0.5

Peak tab	ole for sam	le 4:	<u>sample 5a</u>
----------	-------------	-------	------------------

Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	4	35	125.00	5,411.3	Lower Marker	43.00
2		2,907	709.55	369.8		104.17
3		10,380	75.00	10.9	Upper Marker	113.00

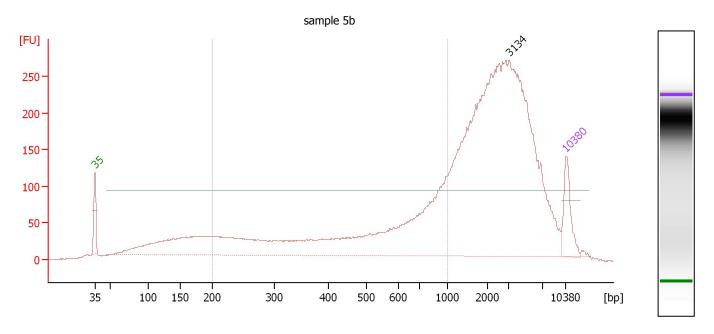
Region table for sample 4: <u>sample 5</u>	<u>а</u>
--	----------

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/µl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	639	14.61	16.8	42.7	2	32.7

2018-08-31_001.xad Page 9 of 16

Assay Class: High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad S/31/2018 12:33:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 5: sample 5b

Number of peaks found: 1 Corr. Area 1: 1,776.9

Noise: 0.4

Peak table for sample 5: sample 5b

Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	4	35	125.00	5,411.3	Lower Marker	43.00
2		3,134	2,324.24	1,123.7		104.62
3		10,380	75.00	10.9	Upper Marker	113.00

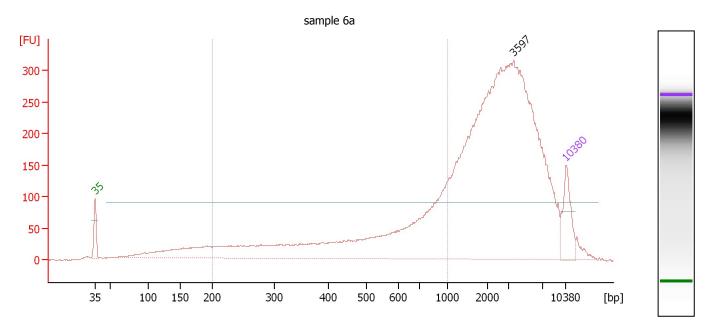
Region table for sample 5: <u>sample 5b</u>

From	To [bp] Average Size	Conc.	Corr.	Molarity	Co % of	Size distribution in
[bp]	[bp]	[pg/µl]	Area	[pmol/l]	<u>lor</u> Total	CV [%]
200	1.000 541	1.258.90	1.776.9	4.827.8	31	43.3

2018-08-31_001.xad Page 10 of 16

Assay Class: High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad S/31/2018 12:33:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 6: <u>sample 6a</u>

Number of peaks found: 1 Corr. Area 1: 1,683.0

Noise: 0.5

Peak table for sample 6 : <u>sample 6a</u>

Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	4	35	125.00	5,411.3	Lower Marker	43.00
2		3,597	2,191.37	923.0		105.20
3		10,380	75.00	10.9	Upper Marker	113.00

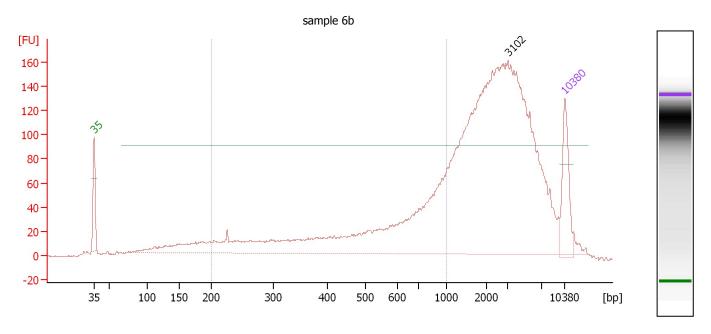
Region table for sample 6 : <u>sample 6a</u>

From	To [bp] Average	Size Conc.	Corr.	Molarity	Co % of	Size distribution in
[bp]	[bp]	[pg/µl]	Area	[pmol/l]	<u>lor</u> Total	CV [%]
200	1.000 561	969.12	1.683.0	3.520.3	29	41.2

2018-08-31_001.xad Page 11 of 16

Assay Class: High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad S/31/2018 12:33:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 7: <u>sample 6b</u>

Number of peaks found: 1 Corr. Area 1: 1,028.4

Noise: 0.7

Peak table for sample 7: <u>sample 6b</u>

Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	4	35	125.00	5,411.3	Lower Marker	43.00
2		3,102	1,555.14	759.5		104.58
3		10,380	75.00	10.9	Upper Marker	113.00

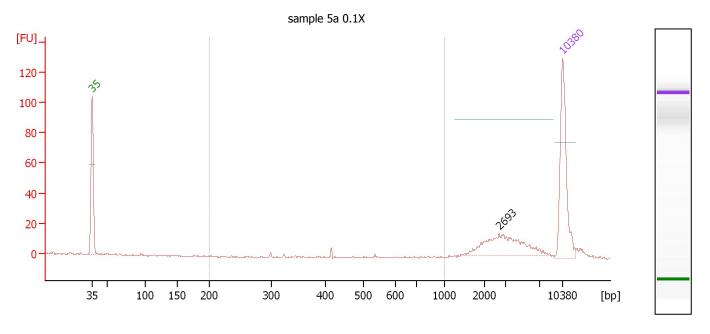
Region table for sample 7 : <u>sample 6b</u>

From	To [bp] Average Size		Corr.	Molarity	Co % of	Size distribution in
[bp]	[bp]	[pg/µl]	Area	[pmol/l]	<u>lor</u> Total	CV [%]
200	1.000 557	821.76	1.028.4	3.000.2	32	41.2

2018-08-31_001.xad Page 12 of 16

Assay Class: High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad S/31/2018 12:33:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : <u>sample 5a 0.1X</u>

Number of peaks found: 1 Corr. Area 1: 5.0

Noise: 0.4

Peak table for sample 8 : <u>sample 5a 0.1X</u>

Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	4	35	125.00	5,411.3	Lower Marker	43.00
2		2,693	57.77	32.5		103.52
3		10,380	75.00	10.9	Upper Marker	113.00

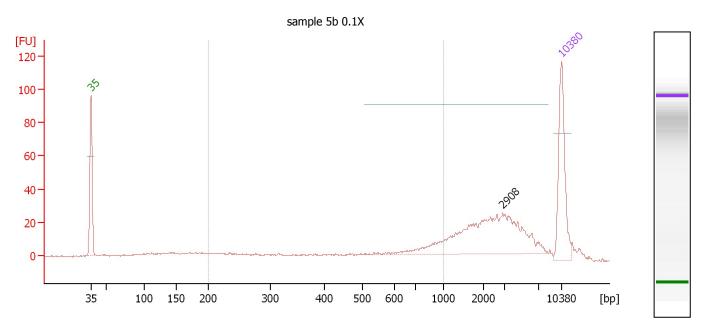
Region table for sample 8 : <u>sample 5a 0.1X</u>

From	To [bp] Average Size	Conc.	Corr.	Molarity	Co % of	Size distribution in
[bp]	[bp]	[pg/µl]	Area	[pmol/l]	<u>lor</u> Total	CV [%]
200	1.000 586	4.16	5.0	13.4	3	38.5

2018-08-31_001.xad Page 13 of 16

Assay Class: High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad S/31/2018 12:33:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : <u>sample 5b 0.1X</u>

Number of peaks found: 1 Corr. Area 1: 134.4

Noise: 0.4

		_		
Peak table	for sample 9	9 :	sample	5b 0.1X

Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	4	35	125.00	5,411.3	Lower Marker	43.00
2		2,908	166.97	87.0		104.17
3		10,380	75.00	10.9	Upper Marker	113.00

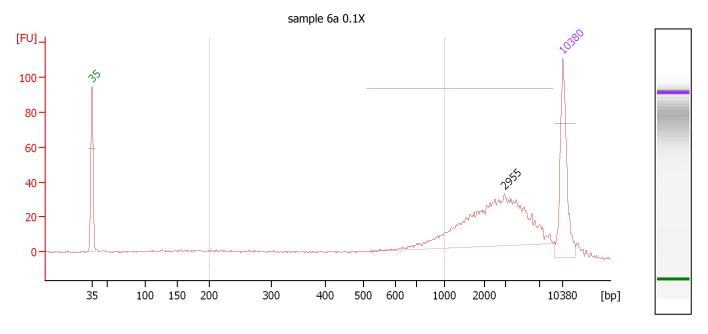
Region table for sample 9:	sample 5b 0.1X
----------------------------	----------------

From	To [bp] Average Size		Corr.	Molarity	Co % of	Size distribution in
[bp]	[bp]	[pg/µl]	Area	[pmol/l]	<u>lor</u> Total	CV [%]
200	1.000 578	123.24	134.4	442.4	28	41.0

2018-08-31_001.xad Page 14 of 16

Assay Class: High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad S/31/2018 12:33:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 10: sample 6a 0.1X

Number of peaks found: 1 Corr. Area 1: 129.0

Noise: 0.3

Peak table for sample 10 : sample 6a 0.1X

Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	4	35	125.00	5,411.3	Lower Marker	43.00
2		2,955	195.40	100.2		104.31
3		10,380	75.00	10.9	Upper Marker	113.00

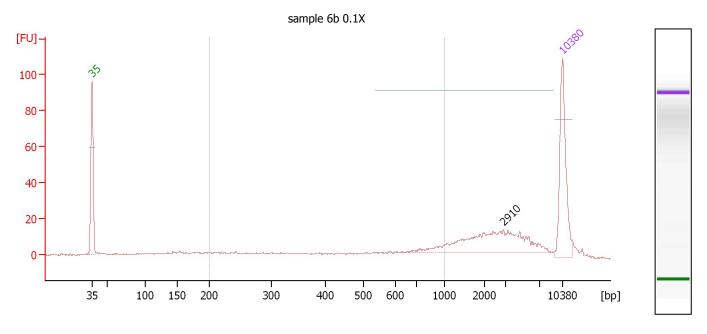
Region table for sample 10: sample 6a 0.1X

From	To [bp] Average Size	Conc.	Corr.	Molarity	Co % of	Size distribution in
[bp]	[bp]	[pg/µl]	Area	[pmol/l]	lor Total	CV [%]
200	1.000 602	116.14	129.0	398.8	23	39.6

2018-08-31_001.xad Page 15 of 16

Assay Class: High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad S/31/2018 12:33:16 PM

Electropherogram Summary Continued ...



Overall Results for sample 11: sample 6b 0.1X

Number of peaks found: 1 Corr. Area 1: 95.7

Noise: 0.3

Peak table for sample 11: sample 6b 0.1X

Pea k		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	4	35	125.00	5,411.3	Lower Marker	43.00
2		2,910	96.35	50.2		104.18
3		10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 11: sample 6b 0.1X

From	To [bp] Average Siz	e Conc.	Corr.	Molarity	Co % of	Size distribution in
[bp]	[bp]	[pg/µl]	Area	[pmol/l]	<u>lor</u> Total	CV [%]
200	1.000 548	98.71	95.7	372.6	31	42.8

2018-08-31_001.xad Page 16 of 16

High Sensitivity DNA Assay C:\... bioanalyzer\2100 expert\data\2018-08-31\2018-08-31_001.xad Created: Modified: 8/31/2018 11:38:57 AM 8/31/2018 12:33:16 PM Assay Class: Data Path: **Gel Image** sample 6b 0... sample 5b 0... sample 6a 0... [bp] sample 5a 0... sample 5b sample 5a sample 6a sample 6b sample 2 10380 — 7000 -3000 -2000 -1000 -700 — 600 — 500 — 400 — 300 — = 200 — 150 — 100 — 35 — L 1 2 3 5 6 7 8 9 10 11