

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-19\2016-05-19_003.xad

Created: 5/19/2016 1:30:45 PM
Modified: 5/19/2016 2:11:12 PM

Electrophoresis File Run Summary

Instrument Information:

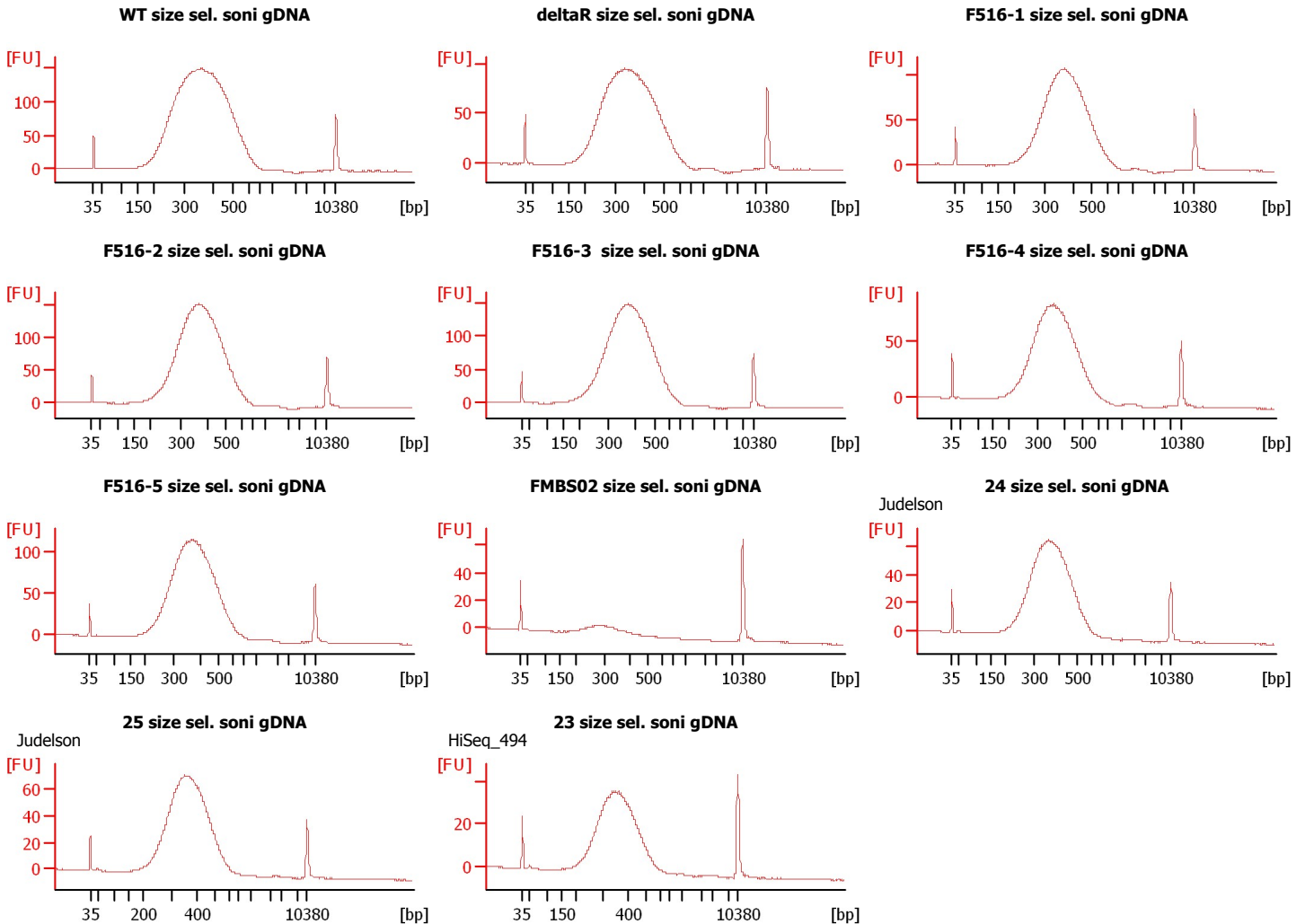
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

Assay Information:

Assay Origin Path: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\assays\dsDNA\High Sensitivity DNA.xsy
Assay Class: High Sensitivity DNA Assay
Version: 1.03
Assay Comments: Copyright © 2003-2010 Agilent Technologies

Chip Information:

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



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Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
WT size sel. soni gDNA		<input type="checkbox"/>	✓			
deltaR size sel. soni gDNA		<input type="checkbox"/>	✓			
F516-1 size sel. soni gDNA		<input type="checkbox"/>	✓			
F516-2 size sel. soni gDNA		<input type="checkbox"/>	✓			
F516-3 size sel. soni gDNA		<input type="checkbox"/>	✓			
F516-4 size sel. soni gDNA		<input type="checkbox"/>	✓			
F516-5 size sel. soni gDNA		<input type="checkbox"/>	✓			
FMBS02 size sel. soni gDNA		<input type="checkbox"/>	✓			
24 size sel. soni gDNA	Judelson	<input type="checkbox"/>	✓			
25 size sel. soni gDNA	Judelson	<input type="checkbox"/>	✓			
23 size sel. soni gDNA	HiSeq_494	<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

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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] : 5
Area Threshold : 0.1
Width Threshold [s] : 0.6
Baseline Plateau [s] : 0.5

Filter Settings

Filter Width [s] : 0.5
Polynomial Order : 4

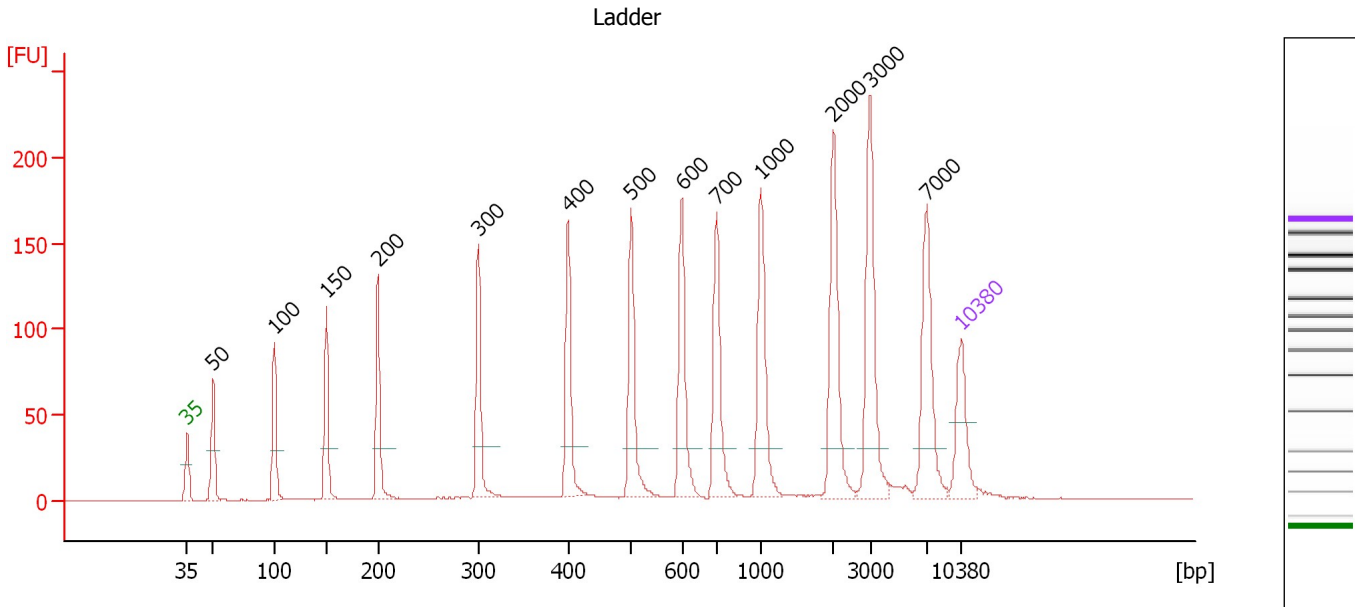
Ladder

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

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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

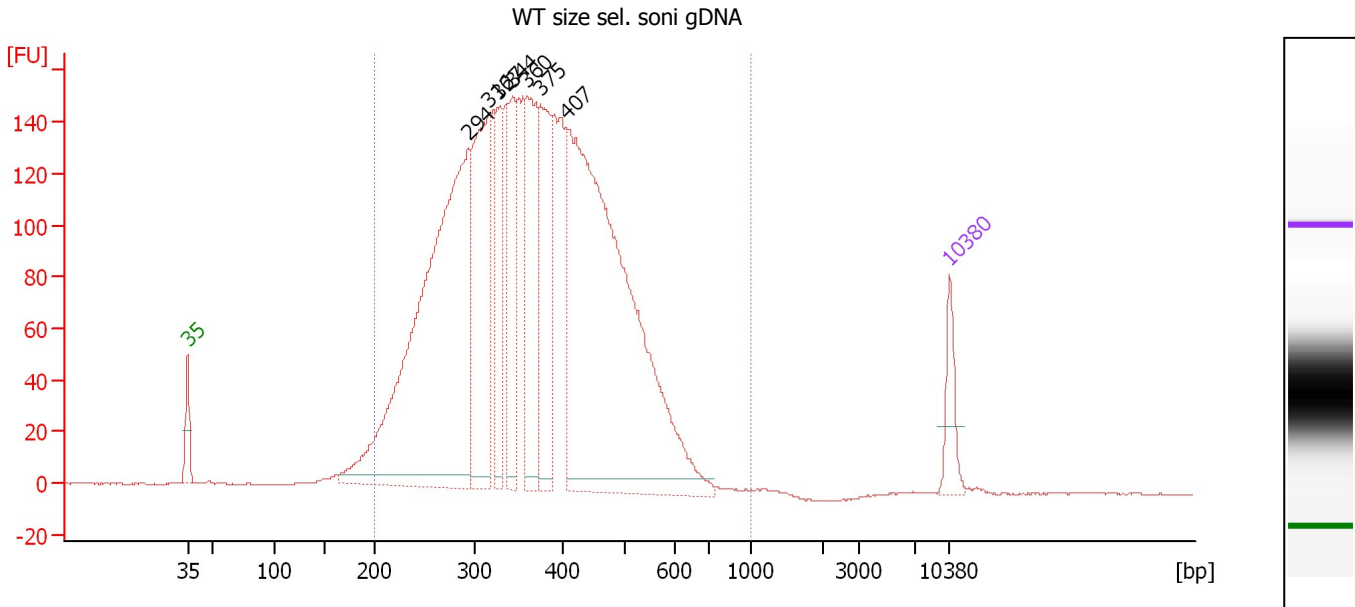
Peak table for Ladder

Peak	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	45.36
3	100	150.00	2,272.7	Ladder Peak	50.90
4	150	150.00	1,515.2	Ladder Peak	55.57
5	200	150.00	1,136.4	Ladder Peak	60.25
6	300	150.00	757.6	Ladder Peak	69.35
7	400	150.00	568.2	Ladder Peak	77.45
8	500	150.00	454.5	Ladder Peak	83.13
9	600	150.00	378.8	Ladder Peak	87.71
10	700	150.00	324.7	Ladder Peak	90.84
11	1,000	150.00	227.3	Ladder Peak	94.84
12	2,000	150.00	113.6	Ladder Peak	101.39
13	3,000	150.00	75.8	Ladder Peak	104.71
14	7,000	150.00	32.5	Ladder Peak	109.82
15	10,380	75.00	10.9	Upper Marker	113.00

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Electropherogram Summary Continued ...



Overall Results for sample 1 : WT size sel. soni gDNA

Number of peaks found: 7 Corr. Area 1: 3,866.3
 Noise: 0.2

Peak table for sample 1 : WT size sel. soni gDNA

Peak	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	294	1,788.74	9,232.1		68.77
3	316	614.70	2,944.0		70.68
4	327	240.77	1,116.8		71.51
5	344	287.31	1,267.0		72.88
6	360	406.18	1,709.8		74.20
7	375	353.63	1,428.5		75.43
8	407	1,867.97	6,958.6		77.83
9	10,380	75.00	10.9	Upper Marker	113.00

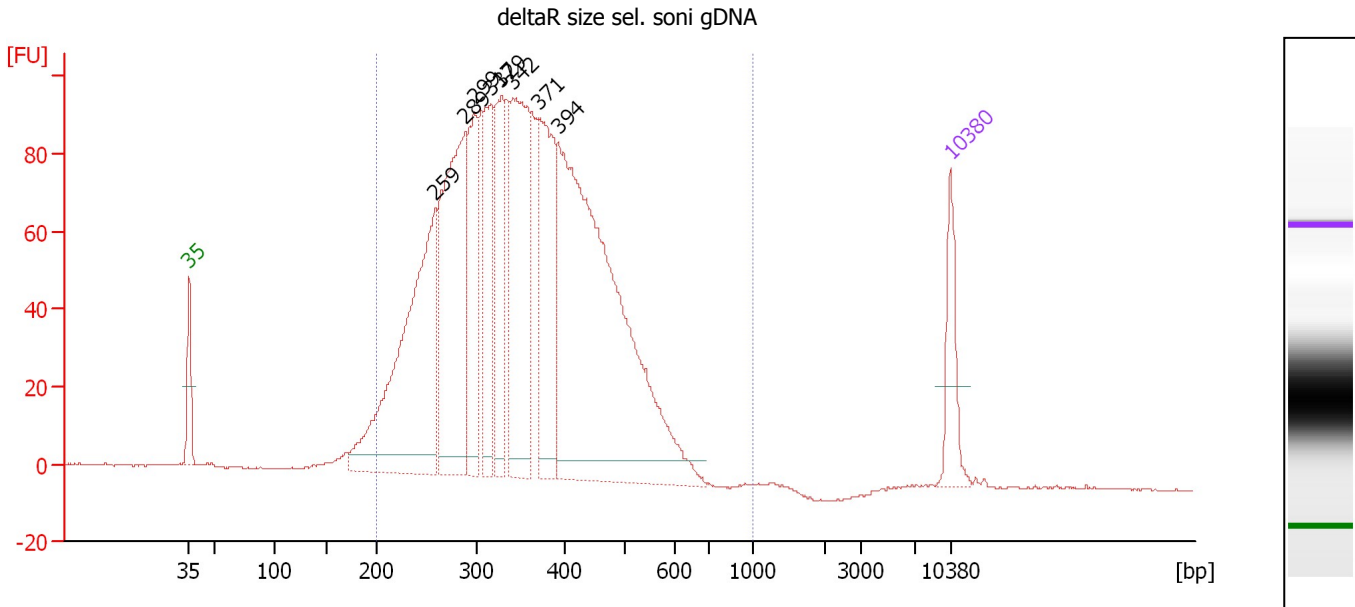
Region table for sample 1 : WT size sel. soni gDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	371	3,866.3	28,600.2	6,421.03	98	25.0

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Electropherogram Summary Continued ...



Overall Results for sample 2 : deltaR size sel. soni gDNA

Number of peaks found: 8 Corr. Area 1: 2,404.5
 Noise: 0.2

Peak table for sample 2 : deltaR size sel. soni gDNA

Peak	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	259	661.53	3,865.4		65.65
3	289	539.53	2,824.2		68.39
4	299	229.49	1,164.5		69.23
5	317	196.65	941.1		70.70
6	329	205.00	943.2		71.73
7	342	406.34	1,800.0		72.75
8	371	310.11	1,266.1		75.11
9	394	1,097.96	4,227.4		76.92
10	10,380	75.00	10.9	Upper Marker	113.00

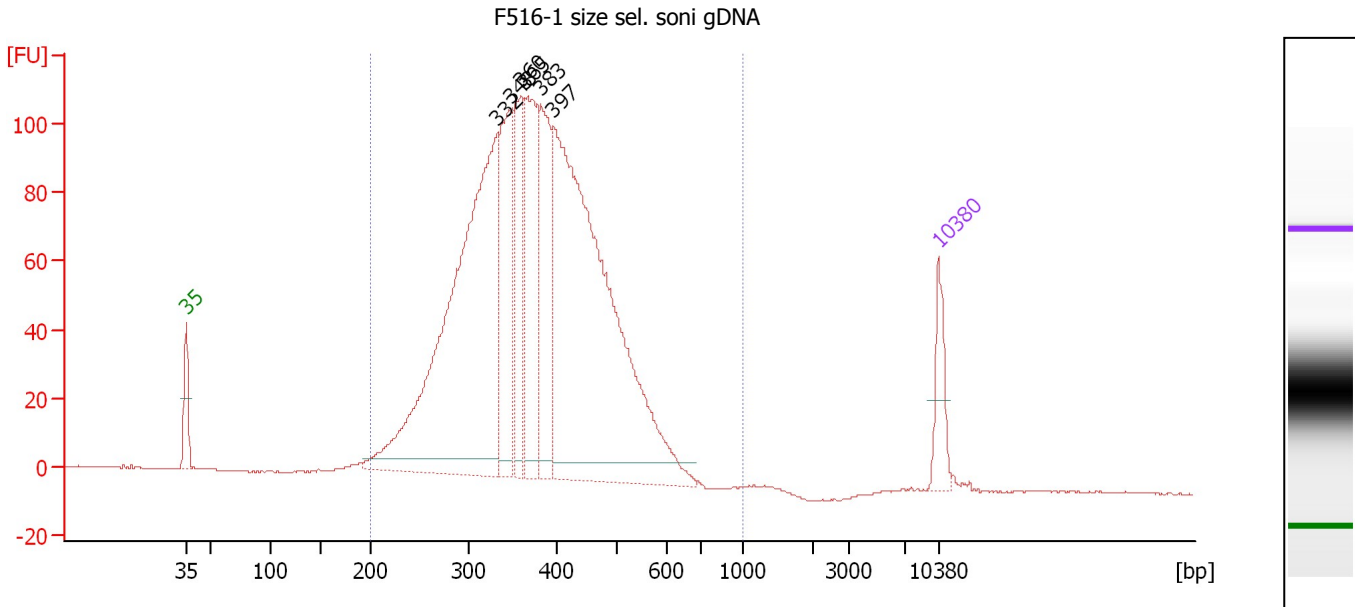
Region table for sample 2 : deltaR size sel. soni gDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	357	2,404.5	17,877.6	3,884.90	97	24.3

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Electropherogram Summary Continued ...



Overall Results for sample 3 : F516-1 size sel. soni gDNA

Number of peaks found: 6 Corr. Area 1: 2,206.8
 Noise: 0.2

Peak table for sample 3 : F516-1 size sel. soni gDNA

Peak	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	332	1,391.79	6,344.6		71.97
3	348	347.22	1,510.4		73.26
4	360	227.32	956.8		74.20
5	365	380.41	1,579.7		74.60
6	383	300.24	1,188.9		76.04
7	397	1,495.58	5,711.7		77.18
8	10,380	75.00	10.9	Upper Marker	113.00

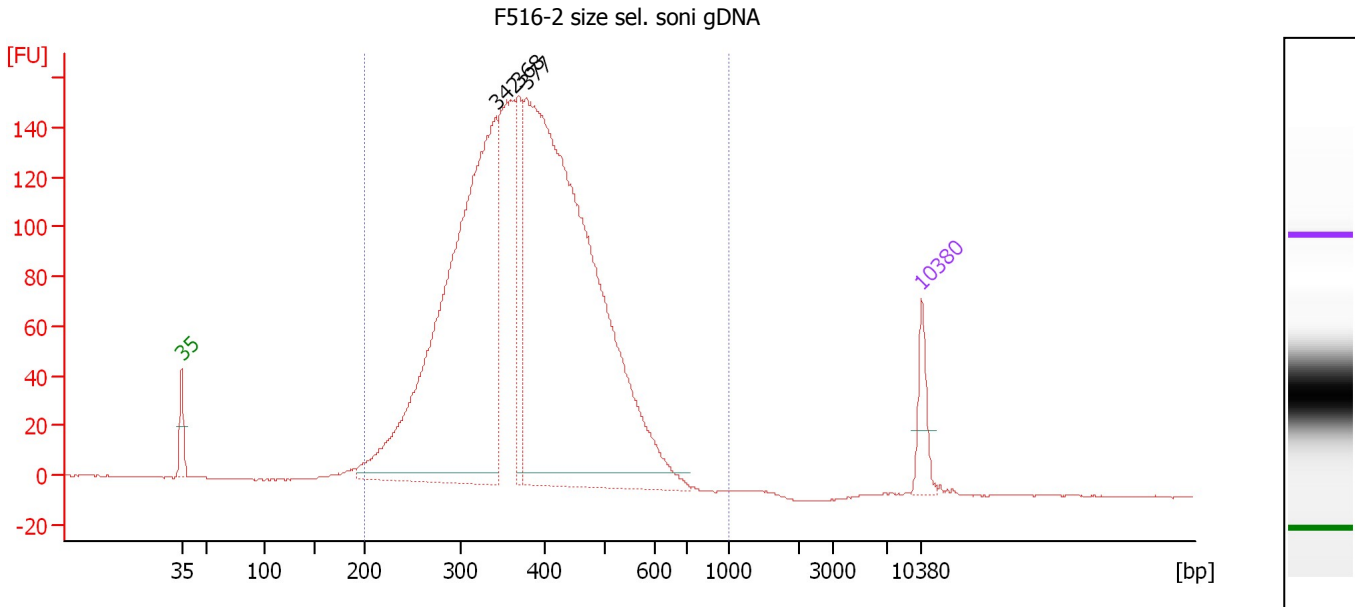
Region table for sample 3 : F516-1 size sel. soni gDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	379	2,206.8	18,387.8	4,320.10	99	21.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-19\2016-05-19_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 4 : F516-2 size sel. soni gDNA

Number of peaks found: 3 Corr. Area 1: 3,174.2
 Noise: 0.2

Peak table for sample 4 : F516-2 size sel. soni gDNA

Peak	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	342	2,048.92	9,085.7		72.73
3	368	251.67	1,034.8		74.90
4	377	2,677.20	10,753.1		75.60
5	10,380	75.00	10.9	Upper Marker	113.00

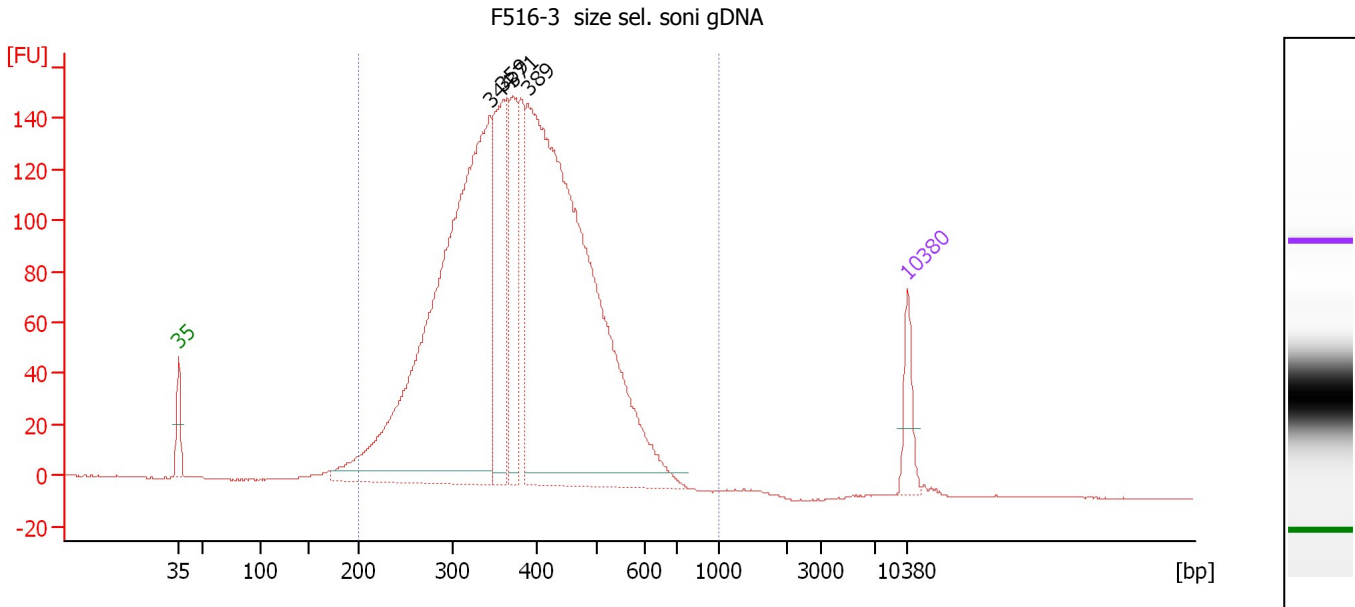
Region table for sample 4 : F516-2 size sel. soni gDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	382	3,174.2	23,633.6	5,576.87	99	21.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-19\2016-05-19_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 5 : F516-3 size sel. soni gDNA

Number of peaks found: 4 Corr. Area 1: 3,205.4
 Noise: 0.2

Peak table for sample 5 : F516-3 size sel. soni gDNA

Peak	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	344	2,095.58	9,240.1		72.88
3	359	448.88	1,892.2		74.16
4	371	293.14	1,197.8		75.08
5	389	2,314.33	9,025.6		76.52
6	10,380	75.00	10.9	Upper Marker	113.00

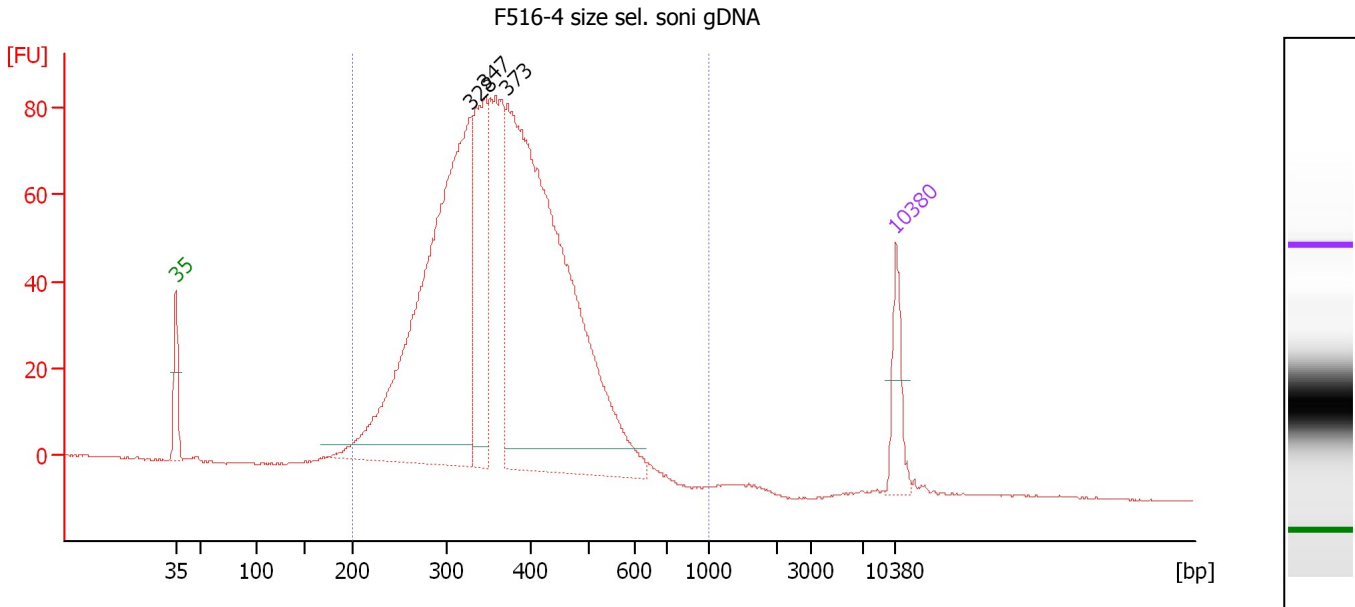
Region table for sample 5 : F516-3 size sel. soni gDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	384	3,205.4	23,353.2	5,507.63	99	22.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-19\2016-05-19_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 6 : F516-4 size sel. soni gDNA

Number of peaks found: 3 Corr. Area 1: 1,738.2
 Noise: 0.2

Peak table for sample 6 : F516-4 size sel. soni gDNA

Peak	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	328	1,348.06	6,226.3		71.62
3	347	387.51	1,693.9		73.13
4	373	1,626.25	6,607.9		75.25
5	10,380	75.00	10.9	Upper Marker	113.00

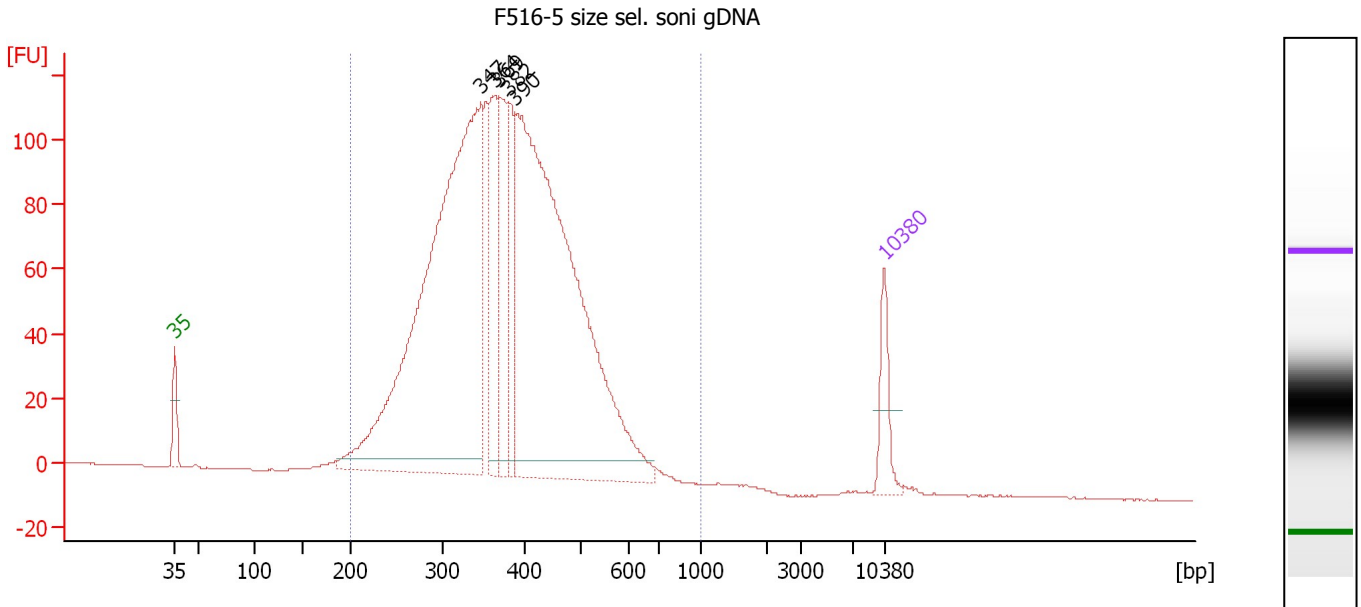
Region table for sample 6 : F516-4 size sel. soni gDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	369	1,738.2	16,964.2	3,869.91	99	21.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-19\2016-05-19_003.xad

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Electropherogram Summary Continued ...



Overall Results for sample 7 : F516-5 size sel. soni gDNA

Number of peaks found: 5 Corr. Area 1: 2,445.5
 Noise: 0.2

Peak table for sample 7 : F516-5 size sel. soni gDNA

Peak	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	347	1,838.66	8,019.4		73.19
3	364	253.31	1,053.6		74.56
4	369	264.43	1,084.4		74.98
5	382	181.71	719.8		76.03
6	390	1,742.26	6,775.2		76.61
7	10,380	75.00	10.9	Upper Marker	113.00

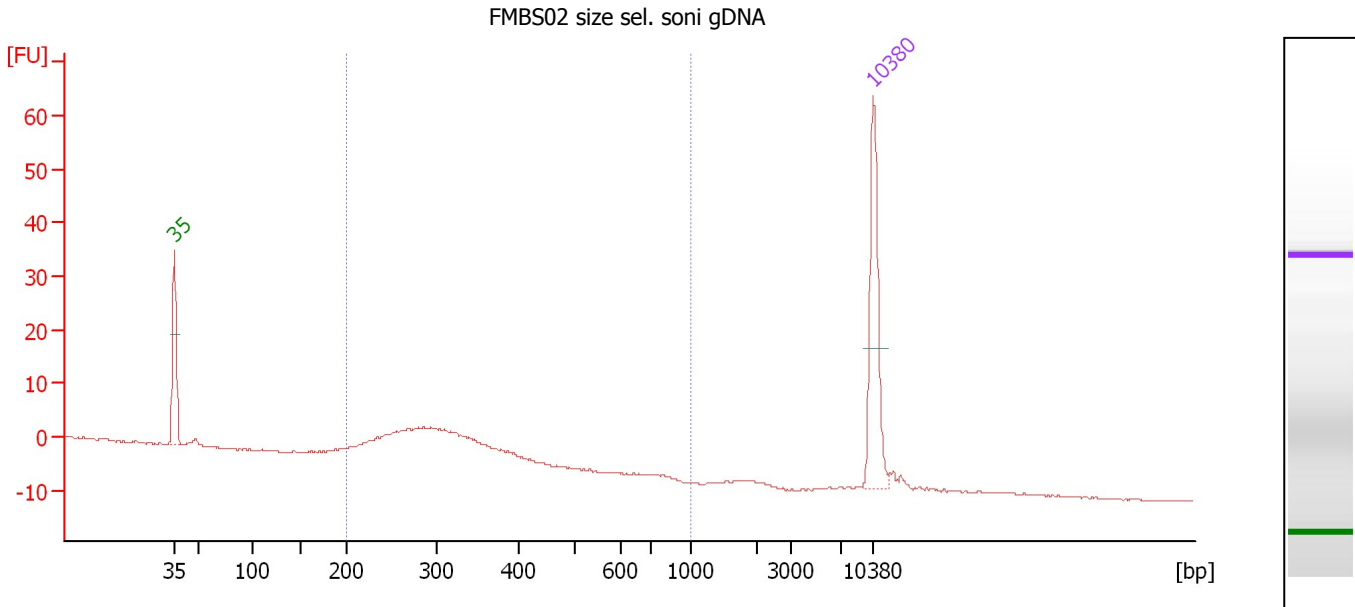
Region table for sample 7 : F516-5 size sel. soni gDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	383	2,445.5	19,410.7	4,565.30	99	23.0

Assay Class: High Sensitivity DNA Assay
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Electropherogram Summary Continued ...



Overall Results for sample 8 : FMBS02 size sel. soni gDNA

Number of peaks found: 0 Corr. Area 1: 92.3
 Noise: 0.1

Peak table for sample 8 : FMBS02 size sel. soni gDNA

Peak	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	10,380	75.00	10.9	Upper Marker	113.00

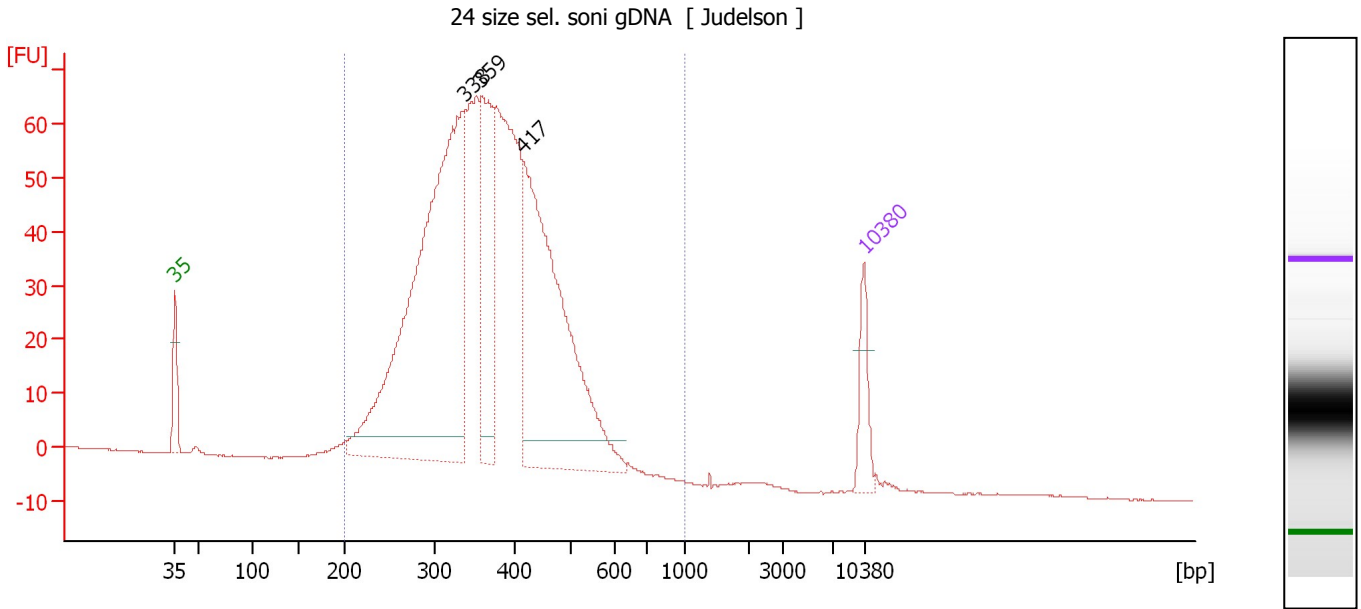
Region table for sample 8 : FMBS02 size sel. soni gDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	300	92.3	937.6	179.48	94	16.8

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Electropherogram Summary Continued ...



Overall Results for sample 9 : 24 size sel. soni gDNA

Number of peaks found: 3 Corr. Area 1: 1,335.6
 Noise: 0.1

Peak table for sample 9 : 24 size sel. soni gDNA

Peak	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	338	1,546.01	6,938.2		72.40
3	359	407.05	1,717.9		74.13
4	417	951.78	3,459.2		78.41
5	10,380	75.00	10.9	Upper Marker	113.00

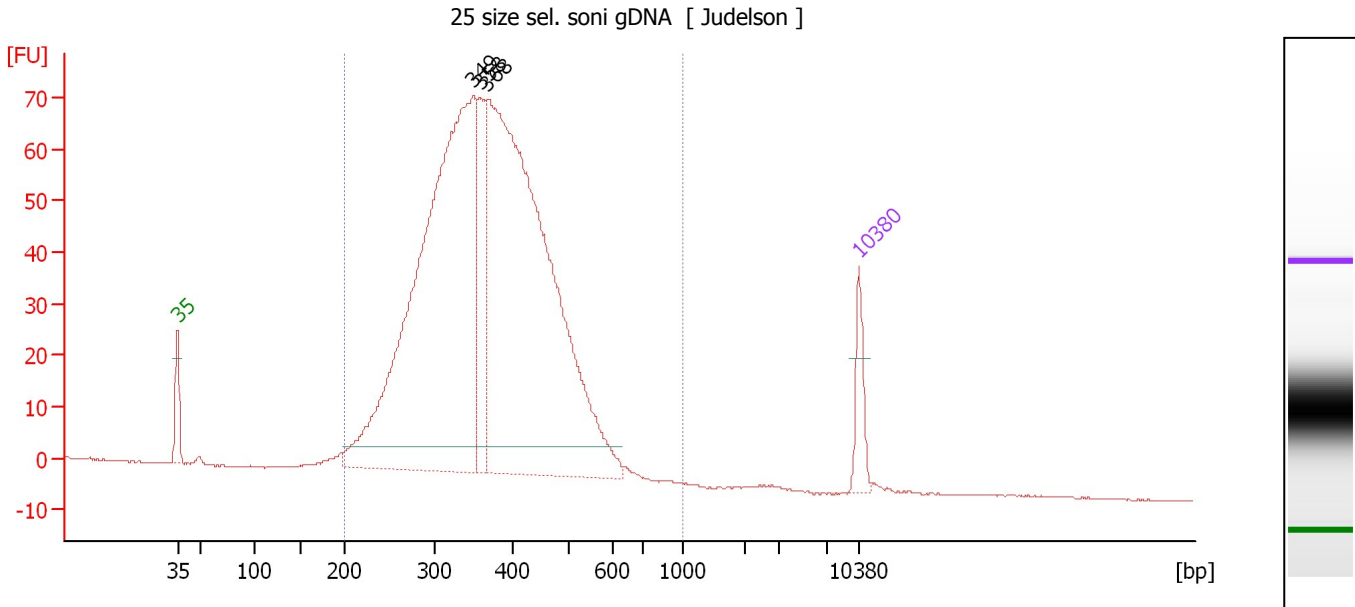
Region table for sample 9 : 24 size sel. soni gDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	371	1,335.6	18,077.0	4,166.19	99	21.1

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Electropherogram Summary Continued ...



Overall Results for sample 10 : 25 size sel. soni gDNA

Number of peaks found: 3 Corr. Area 1: 1,419.6
 Noise: 0.1

Peak table for sample 10 : 25 size sel. soni gDNA

Peak	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	349	2,091.13	9,087.7		73.29
3	358	283.03	1,197.4		74.06
4	368	2,185.56	9,008.5		74.82
5	10,380	75.00	10.9	Upper Marker	113.00

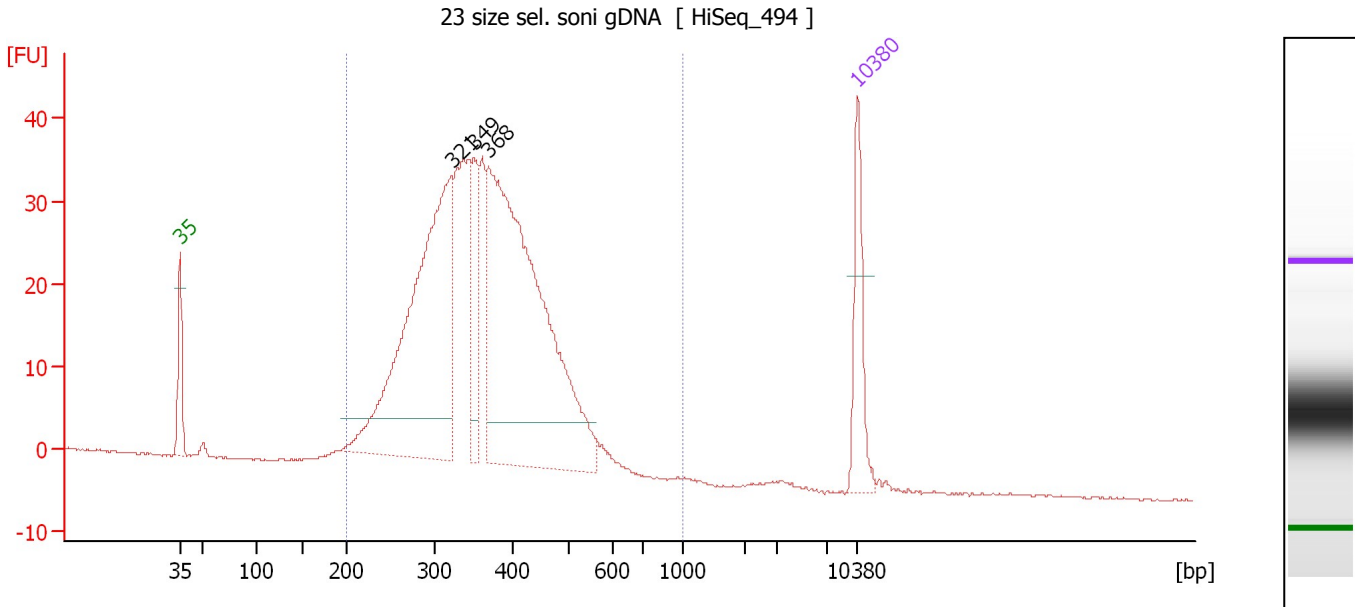
Region table for sample 10 : 25 size sel. soni gDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	372	1,419.6	20,250.9	4,676.14	99	21.4

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Electropherogram Summary Continued ...



Overall Results for sample 11 : 23 size sel. soni gDNA

Number of peaks found: 3 Corr. Area 1: 708.8
 Noise: 0.1

Peak table for sample 11 : 23 size sel. soni gDNA

Peak	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	321	624.94	2,945.7		71.09
3	349	104.05	451.3		73.35
4	368	761.42	3,131.6		74.89
5	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 11 : 23 size sel. soni gDNA

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	362	708.8	8,490.1	1,913.22	99	21.0

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Gel Image

Assay Class: High Sensitivity DNA Assay
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Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 12)		Instrument	Run		5/19/2016 2:11:12 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-05-19\2016-05-19_003.xad)		Instrument	Run		5/19/2016 1:30:51 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		5/19/2016 1:30:51 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		5/19/2016 1:30:51 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		5/19/2016 1:30:51 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		5/19/2016 1:30:51 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		5/19/2016 1:30:51 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		5/19/2016 1:30:51 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1