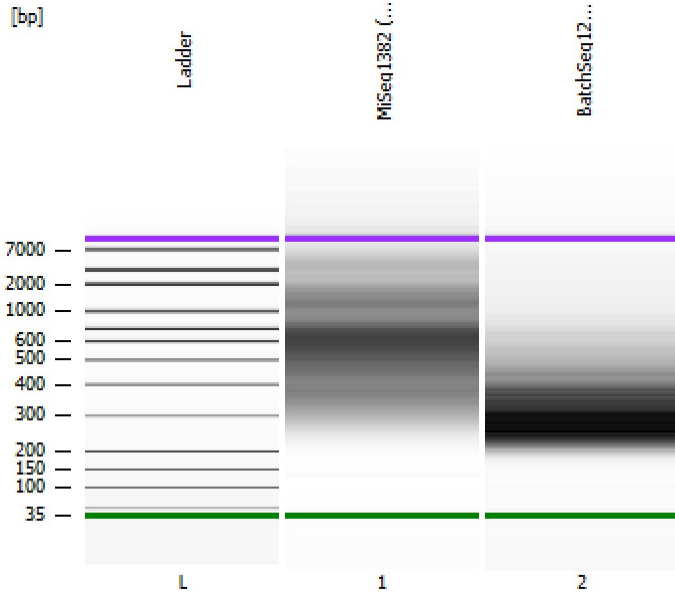


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...zer\2100 expert\data\2023-10-12\MiSeq1382P_BatchSeq121P.xad

Created: 10/12/2023 4:45:15 PM
Modified: 10/12/2023 5:33:25 PM

Electrophoresis File Run Summary



Instrument Information:

Instrument Name: DE34903152 Firmware: C.01.069
Serial#: DE34903152 Type: G2938C

Assay Information:

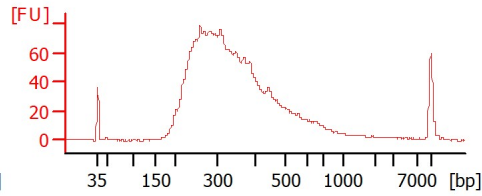
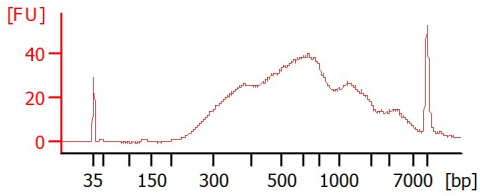
Assay Origin Path: C:\Program Files (x86)\Agilent\2100 bioanalyzer\2100 expert\assays\dsDNA\High Sensitivity DNA.xsy
Assay Class: High Sensitivity DNA Assay
Version: 1.03
Assay Comments: Copyright © 2003-2010 Agilent Technologies

Chip Information:

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

MiSeq1382 (1x BC)

BatchSeq121_1xBC (1:3)



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...zer\2100 expert\data\2023-10-12\MiSeq1382P_BatchSeq121P.xad

Created: 10/12/2023 4:45:15 PM
 Modified: 10/12/2023 5:33:25 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
MiSeq1382 (1x BC)		<input type="checkbox"/>		✓		
BatchSeq121_1xBC (1:3)		<input type="checkbox"/>		✓		
Ladder		<input type="checkbox"/>		✓		

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...zer\2100 expert\data\2023-10-12\MiSeq1382P_BatchSeq121P.xad

Created: 10/12/2023 4:45:15 PM
Modified: 10/12/2023 5:33:25 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/ μ l] : 1950
Uses Standard Area for Ladder Fragments
Lower Marker Concentration [pg/ μ l] : 125
Upper Marker Concentration [pg/ μ l] : 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

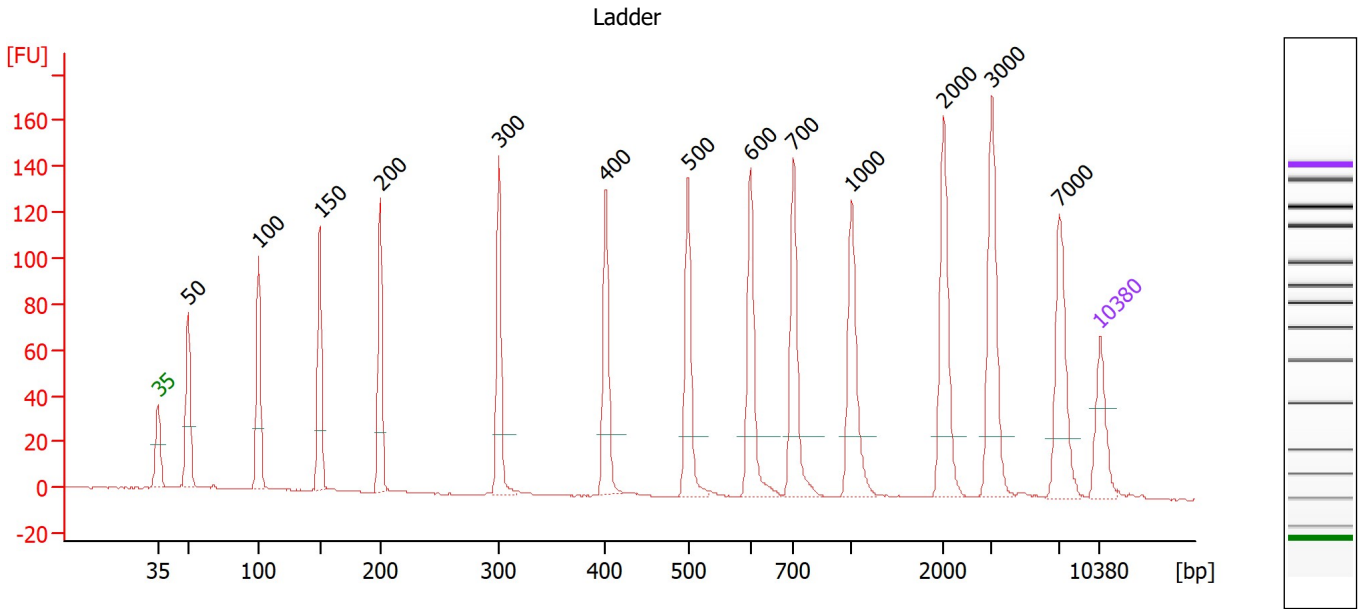
Ladder

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

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...zer\2100 expert\data\2023-10-12\MiSeq1382P_BatchSeq121P.xad

Created: 10/12/2023 4:45:15 PM
 Modified: 10/12/2023 5:33:25 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

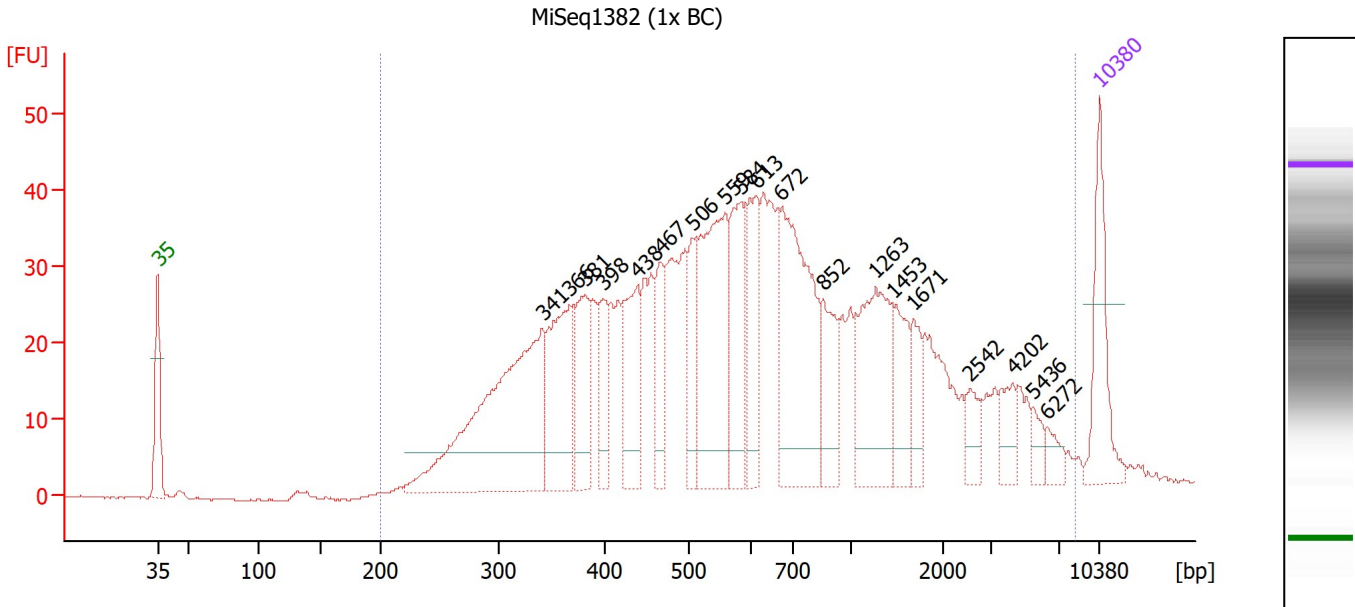
Peak table for Ladder

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	50	150.00	4,545.5	Ladder Peak
3	100	150.00	2,272.7	Ladder Peak
4	150	150.00	1,515.2	Ladder Peak
5	200	150.00	1,136.4	Ladder Peak
6	300	150.00	757.6	Ladder Peak
7	400	150.00	568.2	Ladder Peak
8	500	150.00	454.5	Ladder Peak
9	600	150.00	378.8	Ladder Peak
10	700	150.00	324.7	Ladder Peak
11	1,000	150.00	227.3	Ladder Peak
12	2,000	150.00	113.6	Ladder Peak
13	3,000	150.00	75.8	Ladder Peak
14	7,000	150.00	32.5	Ladder Peak
15	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...zer\2100 expert\data\2023-10-12\MiSeq1382P_BatchSeq121P.xad

Created: 10/12/2023 4:45:15 PM
 Modified: 10/12/2023 5:33:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : MiSeq1382 (1x BC)

Number of peaks found: 19 Corr. Area 1: 1,244.3
 Noise: 0.1

Peak table for sample 1 : MiSeq1382 (1x BC)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	341	380.11	1,691.3	
3	366	158.13	654.1	
4	381	88.43	351.3	
5	398	53.67	204.5	
6	438	99.17	343.4	
7	467	63.73	206.9	
8	506	59.27	177.3	
9	559	214.46	580.9	
10	584	111.15	288.3	
11	613	88.17	217.9	
12	672	244.00	550.4	
13	852	65.12	115.9	
14	1,263	131.79	158.1	
15	1,453	55.53	57.9	
16	1,671	34.24	31.0	
17	2,542	25.11	15.0	
18	4,202	27.34	9.9	
19	5,436	13.85	3.9	
20	6,272	15.27	3.7	
21	10,380	75.00	10.9	Upper Marker

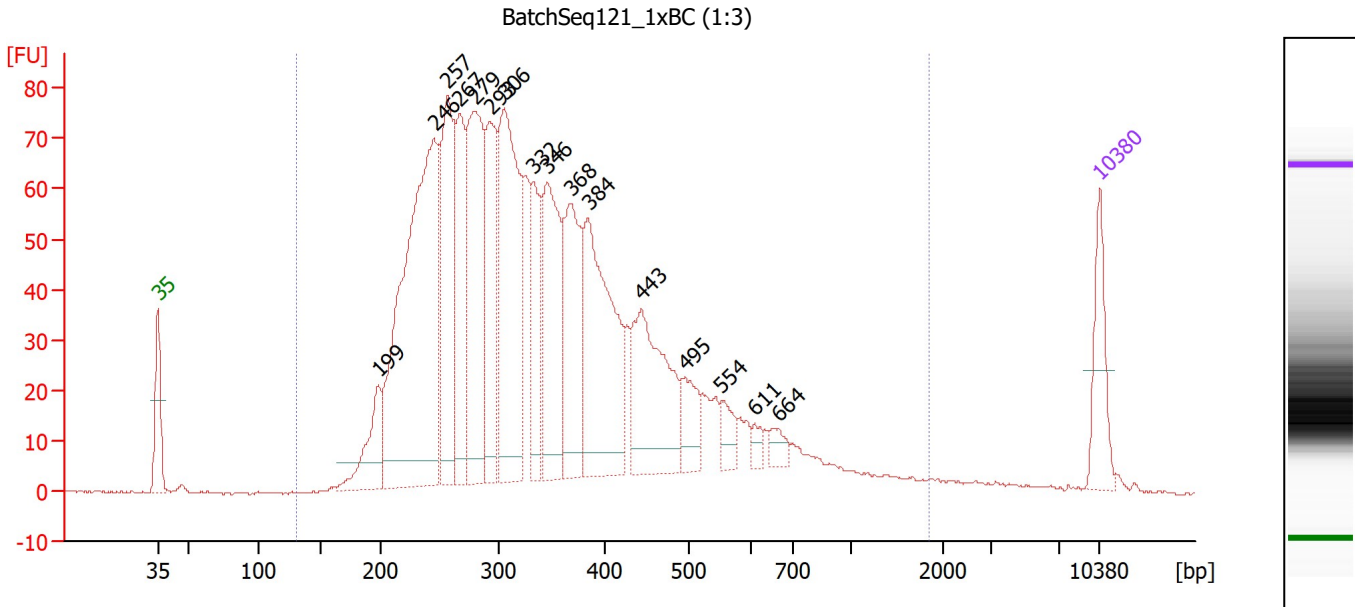
Region table for sample 1 : MiSeq1382 (1x BC)

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	8,258	1,244.3	98	1,033	100.0	2,829.58	8,255.3	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...zer\2100 expert\data\2023-10-12\MiSeq1382P_BatchSeq121P.xad

Created: 10/12/2023 4:45:15 PM
 Modified: 10/12/2023 5:33:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : BatchSeq121 1xBC (1:3)

Number of peaks found: 16 Corr. Area 1: 1,907.3
 Noise: 0.3

Peak table for sample 2 : BatchSeq121 1xBC (1:3)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	199	119.98	912.9	
3	246	786.61	4,848.2	
4	257	309.14	1,825.0	
5	267	217.14	1,232.7	
6	279	369.62	2,004.2	
7	293	225.42	1,167.6	
8	306	430.25	2,133.0	
9	332	139.49	636.1	
10	346	268.62	1,177.9	
11	368	209.80	864.5	
12	384	354.37	1,399.9	
13	443	249.93	854.1	
14	495	69.06	211.5	
15	554	34.34	93.9	
16	611	16.08	39.9	
17	664	21.40	48.8	
18	10,380	75.00	10.9	Upper Marker

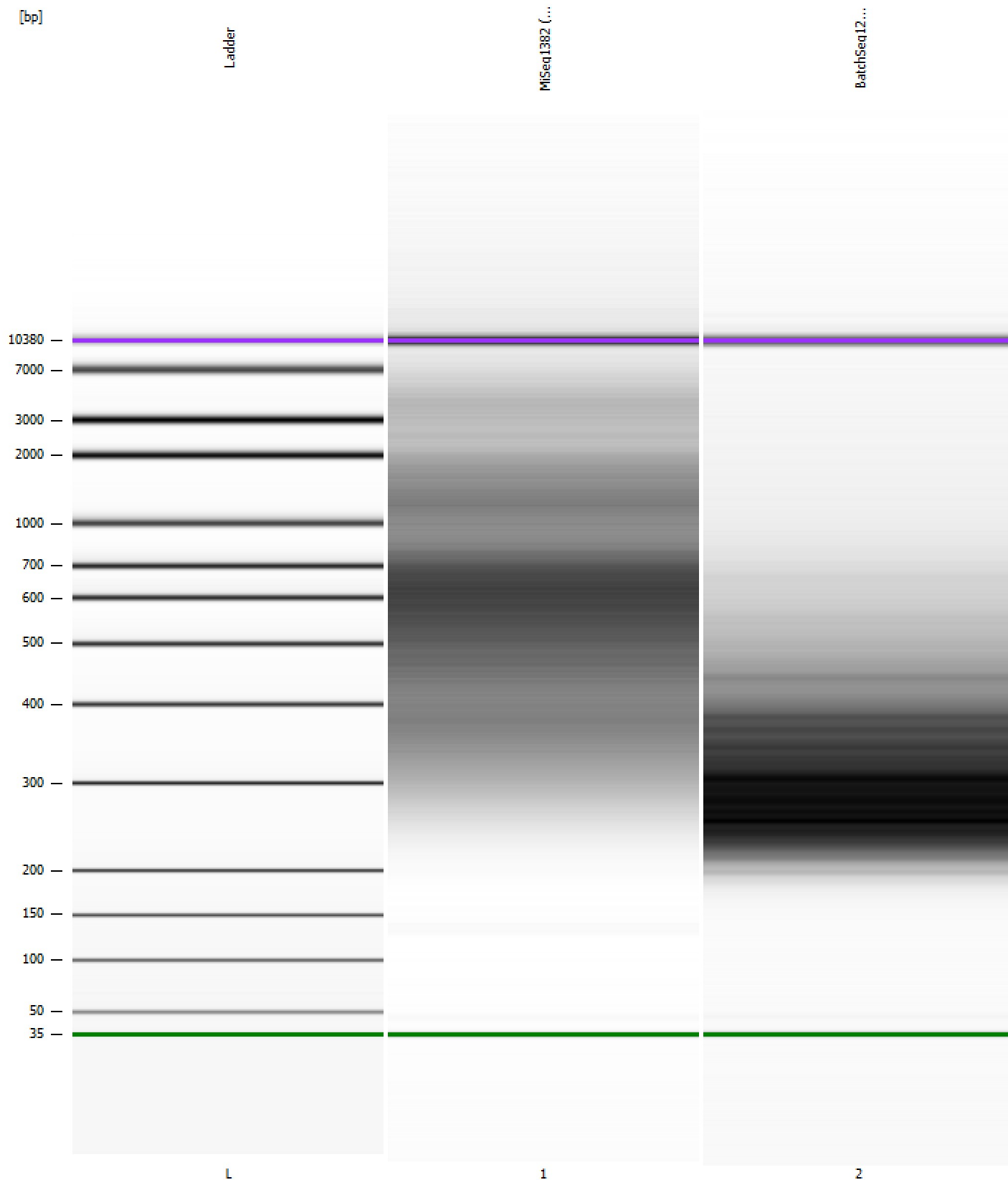
Region table for sample 2 : BatchSeq121 1xBC (1:3)

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
131	1,844	1,907.3	98	370	49.4	4,530.58	22,117.4	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...zer\2100 expert\data\2023-10-12\MiSeq1382P_BatchSeq121P.xad

Created: 10/12/2023 4:45:15 PM
Modified: 10/12/2023 5:33:25 PM

Gel Image

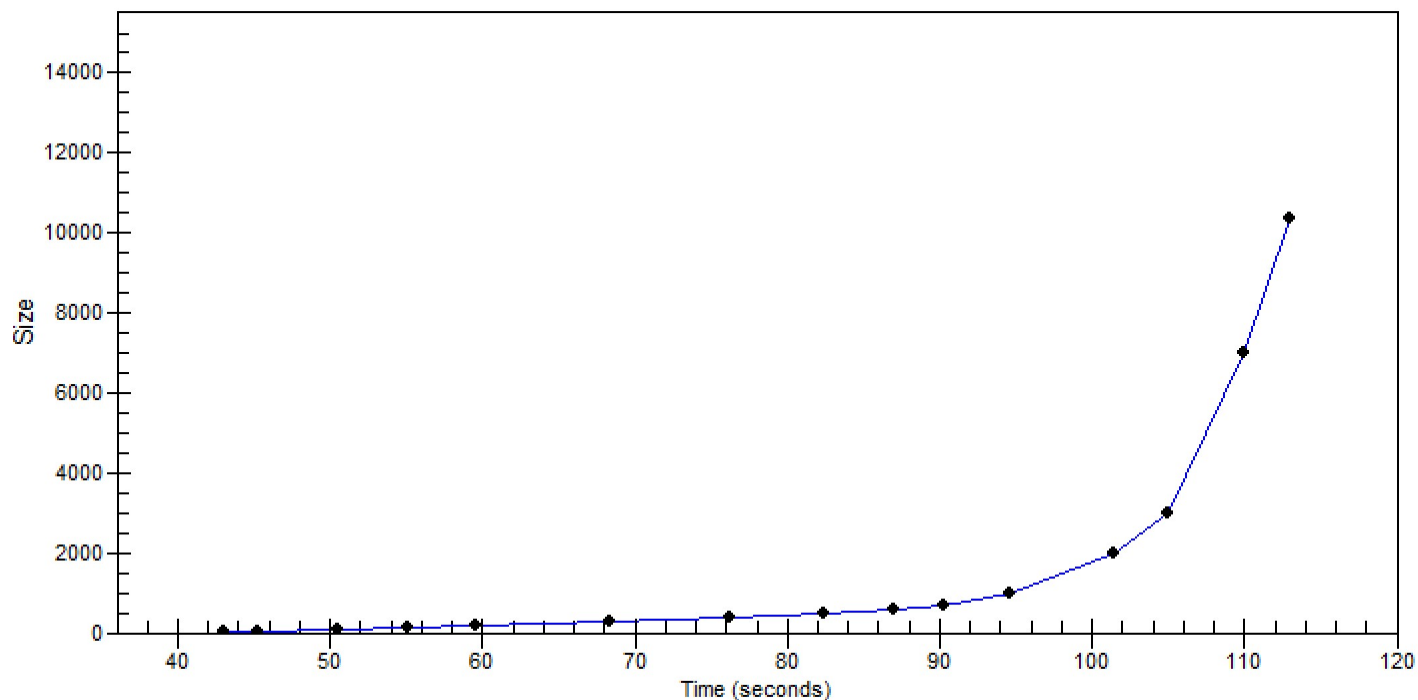


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...zer\2100 expert\data\2023-10-12\MiSeq1382P_BatchSeq121P.xad

Created: 10/12/2023 4:45:15 PM
Modified: 10/12/2023 5:33:25 PM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...zer\2100 expert\data\2023-10-12\MiSeq1382P_BatchSeq121P.xad

Created: 10/12/2023 4:45:15 PM
 Modified: 10/12/2023 5:33:25 PM

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		10/12/2023 5:26:39 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Run started on port 1 (File: C:\Program Files (x86)\Agilent\2100 bioanalyzer\2100 expert\data\2023-10-12\Bioanalyzer1_High Sensitivity DNA Assay_DE34903152_2023-10-12_001.xad)		Instrument	Run		10/12/2023 4:45:23 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Product Number : G2938C		Instrument	Run		10/12/2023 4:45:23 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Name :		Instrument	Run		10/12/2023 4:45:23 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Vendor : Agilent Technologies		Instrument	Run		10/12/2023 4:45:23 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Serial# : DE34903152		Instrument	Run		10/12/2023 4:45:23 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Firmware : C.01.069		Instrument	Run		10/12/2023 4:45:23 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Cartridge : Electrode		Instrument	Run		10/12/2023 4:45:23 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB