

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...alyzer\2100 expert\data\2024-12-26\CUSTOM0099 Libraries.xad

Created: 12/26/2024 1:25:12 PM
Modified: 12/26/2024 3:27:29 PM

Electrophoresis File Run Summary

Instrument Information:

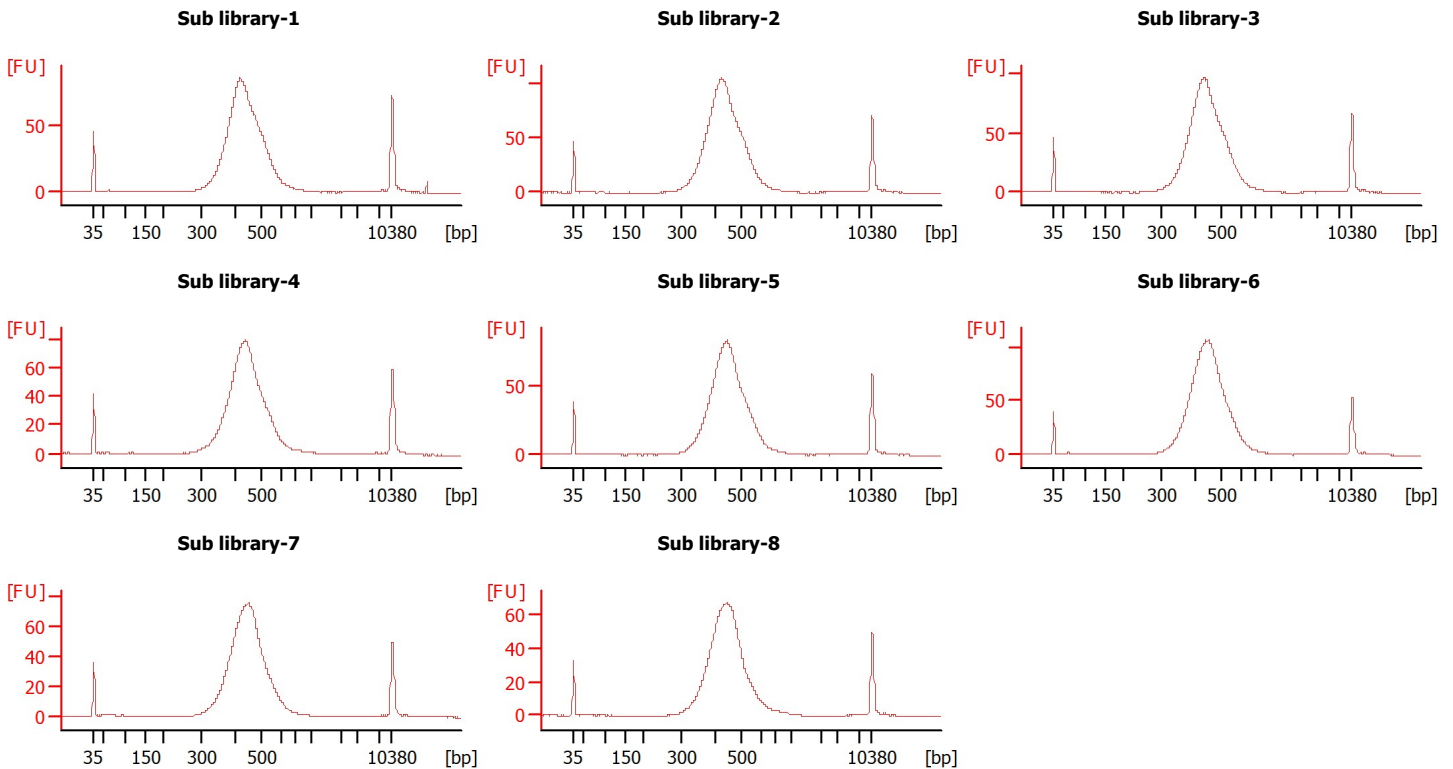
Instrument Name: DE34903152 Firmware: C.01.069
Serial#: DE34903152 Type: G2938C

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:



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Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Sub library-1		<input type="checkbox"/>		✓		
Sub library-2		<input type="checkbox"/>		✓		
Sub library-3		<input type="checkbox"/>		✓		
Sub library-4		<input type="checkbox"/>		✓		
Sub library-5		<input type="checkbox"/>		✓		
Sub library-6		<input type="checkbox"/>		✓		
Sub library-7		<input type="checkbox"/>		✓		
Sub library-8		<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

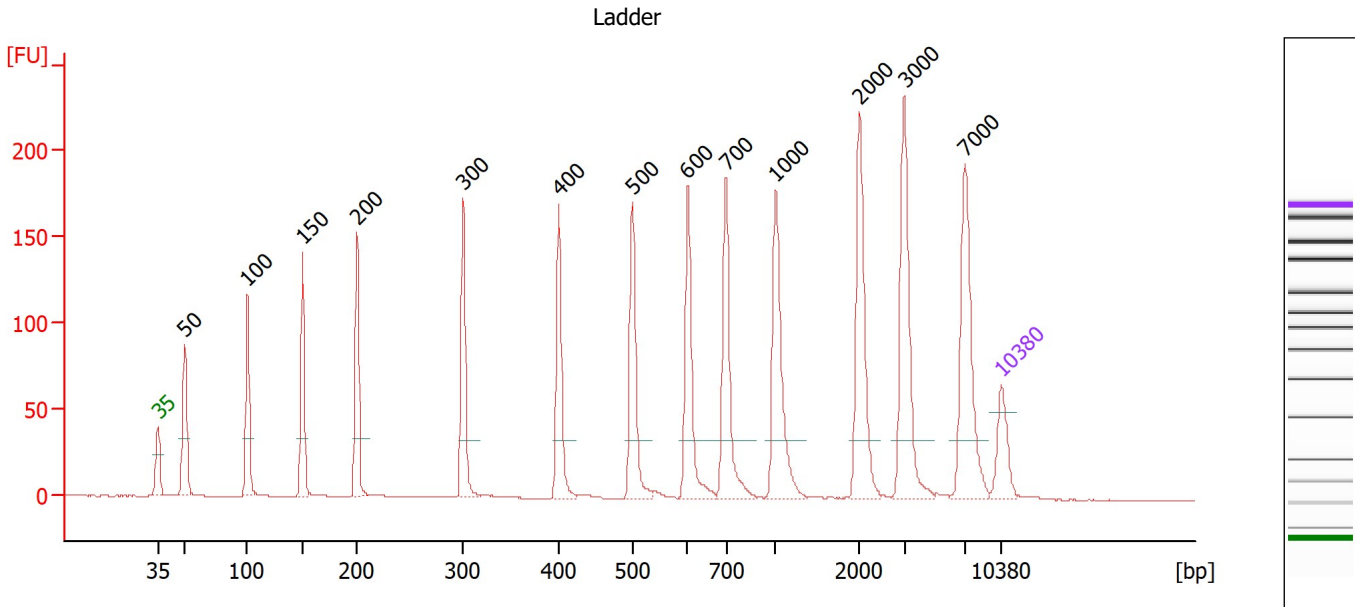
Ladder

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

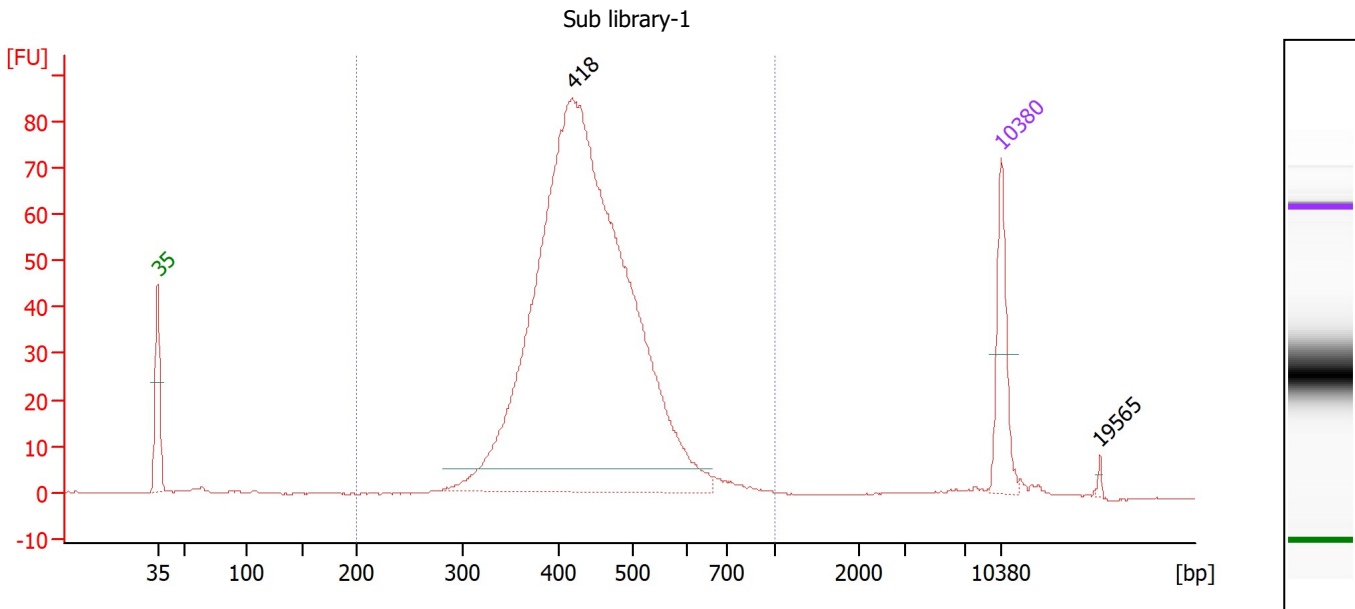
Peak table for Ladder

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
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

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Electropherogram Summary Continued ...



Overall Results for sample 3 : Sub library-1

Number of peaks found: 2 Corr. Area 1: 953.6
 Noise: 0.1

Peak table for sample 3 : Sub library-1

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	418	1,670.24	6,061.1	
3	10,380	75.00	10.9	Upper Marker
4	19,565	0.00	0.0	

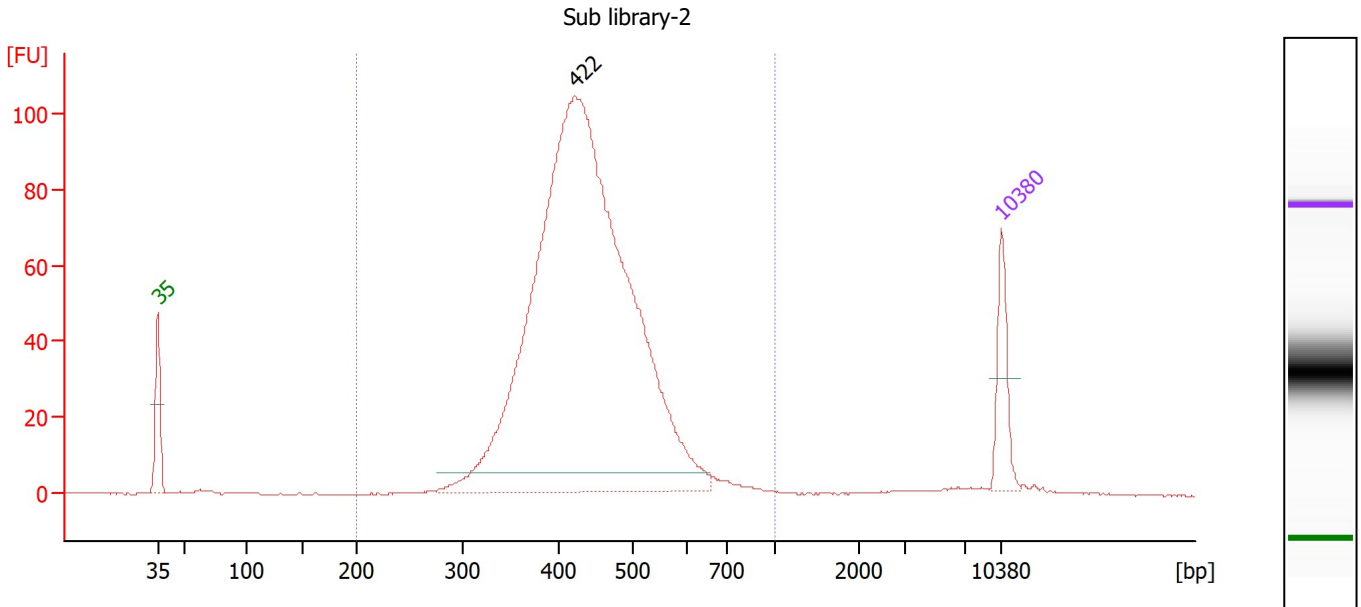
Region table for sample 3 : Sub library-1

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	953.6	95	444	17.7	1,747.56	6,176.1	Blue

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Electropherogram Summary Continued ...



Overall Results for sample 4 : Sub library-2

Number of peaks found: 1 Corr. Area 1: 1,189.1
 Noise: 0.2

Peak table for sample 4 : Sub library-2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	422	2,203.55	7,903.1	
3	10,380	75.00	10.9	Upper Marker

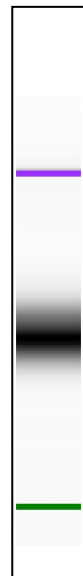
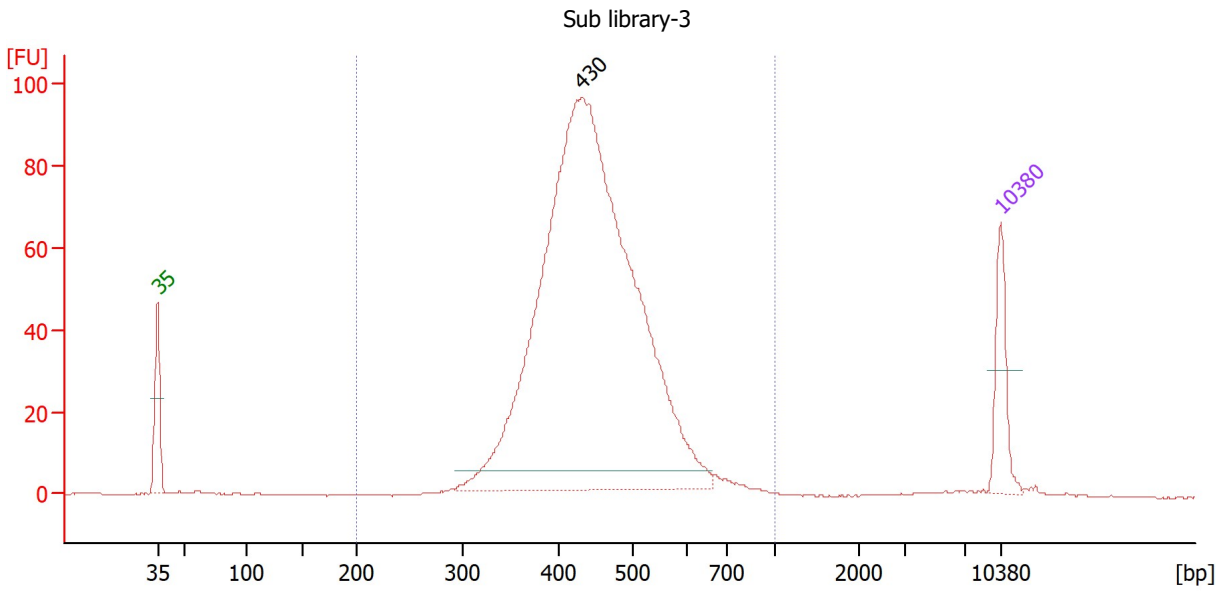
Region table for sample 4 : Sub library-2

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	1,189.1	96	443	17.5	2,291.39	8,112.5	Blue

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 Data Path: C:\...alyzer\2100 expert\data\2024-12-26\CUSTOM0099 Libraries.xad

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Electropherogram Summary Continued ...



Overall Results for sample 5 : Sub library-3

Number of peaks found: 1 Corr. Area 1: 1,084.8
 Noise: 0.2

Peak table for sample 5 : Sub library-3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	430	1,942.09	6,836.1	
3	10,380	75.00	10.9	Upper Marker

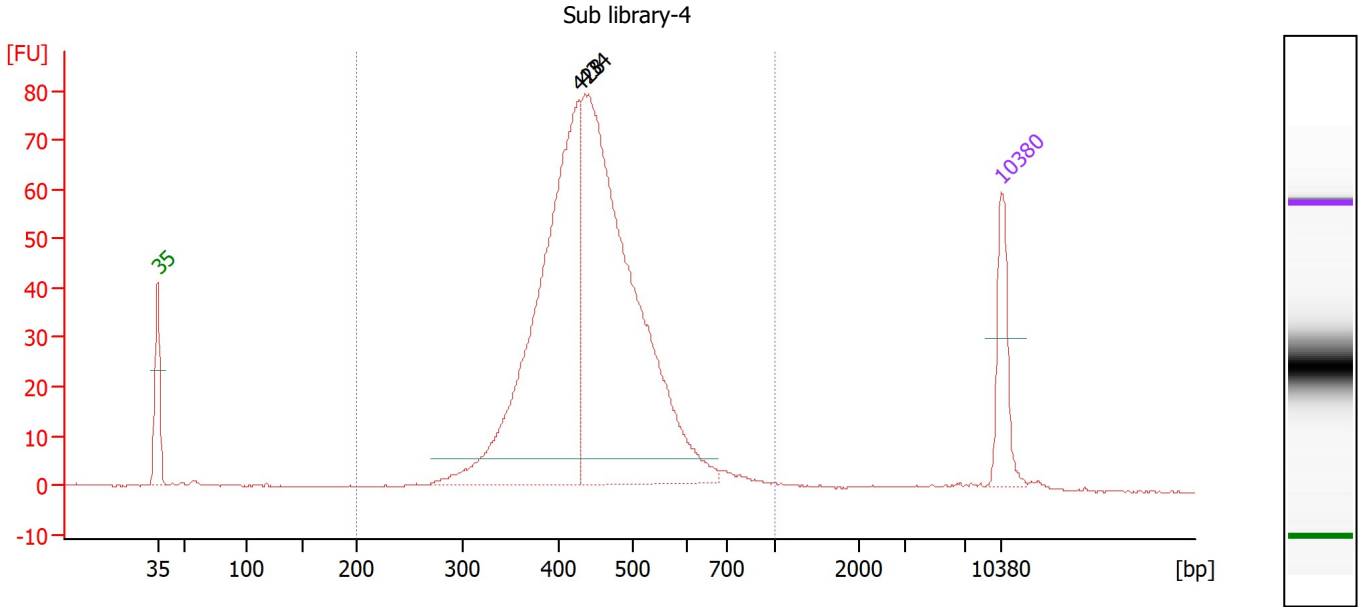
Region table for sample 5 : Sub library-3

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	1,084.8	96	451	17.5	2,052.86	7,141.9	Blue

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 Data Path: C:\...alyzer\2100 expert\data\2024-12-26\CUSTOM0099 Libraries.xad

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Electropherogram Summary Continued ...



Overall Results for sample 6 : Sub library-4

Number of peaks found: 2 Corr. Area 1: 890.3
 Noise: 0.1

Peak table for sample 6 : Sub library-4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	428	735.75	2,607.1	
3	434	930.55	3,245.7	
4	10,380	75.00	10.9	Upper Marker

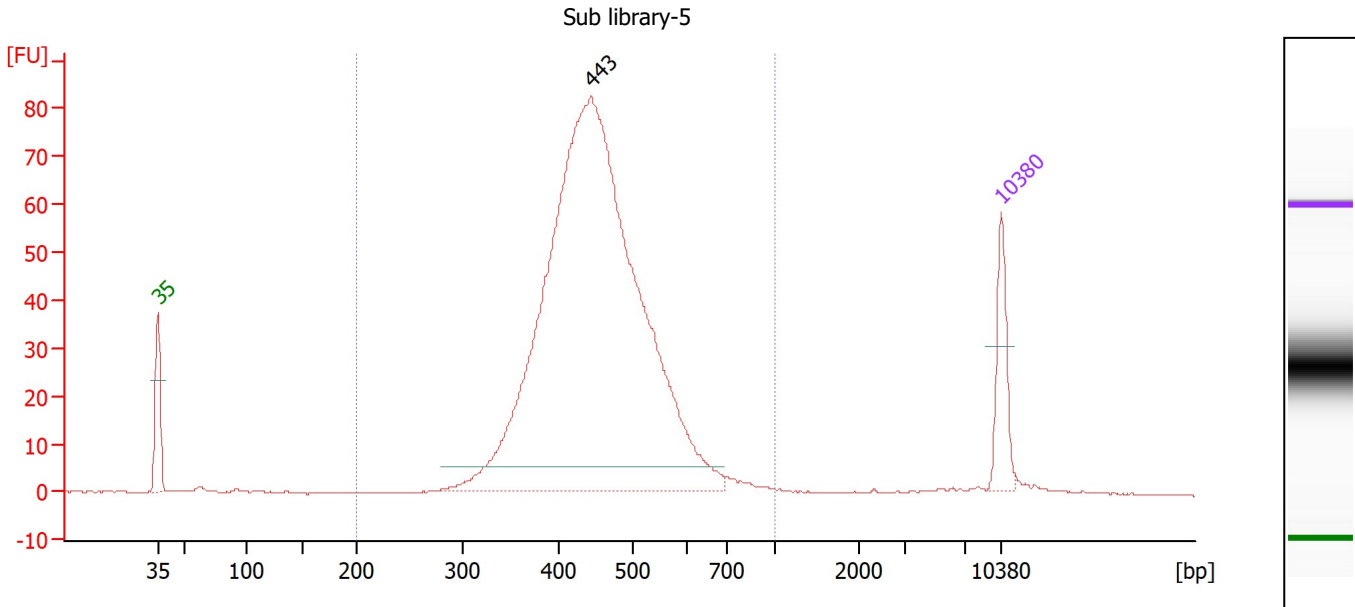
Region table for sample 6 : Sub library-4

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	890.3	95	450	18.5	1,779.21	6,230.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...alyzer\2100 expert\data\2024-12-26\CUSTOM0099 Libraries.xad

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Electropherogram Summary Continued ...



Overall Results for sample 7 : Sub library-5

Number of peaks found: 1 Corr. Area 1: 905.1
 Noise: 0.2

Peak table for sample 7 : Sub library-5

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	443	1,960.02	6,705.9	
3	10,380	75.00	10.9	Upper Marker

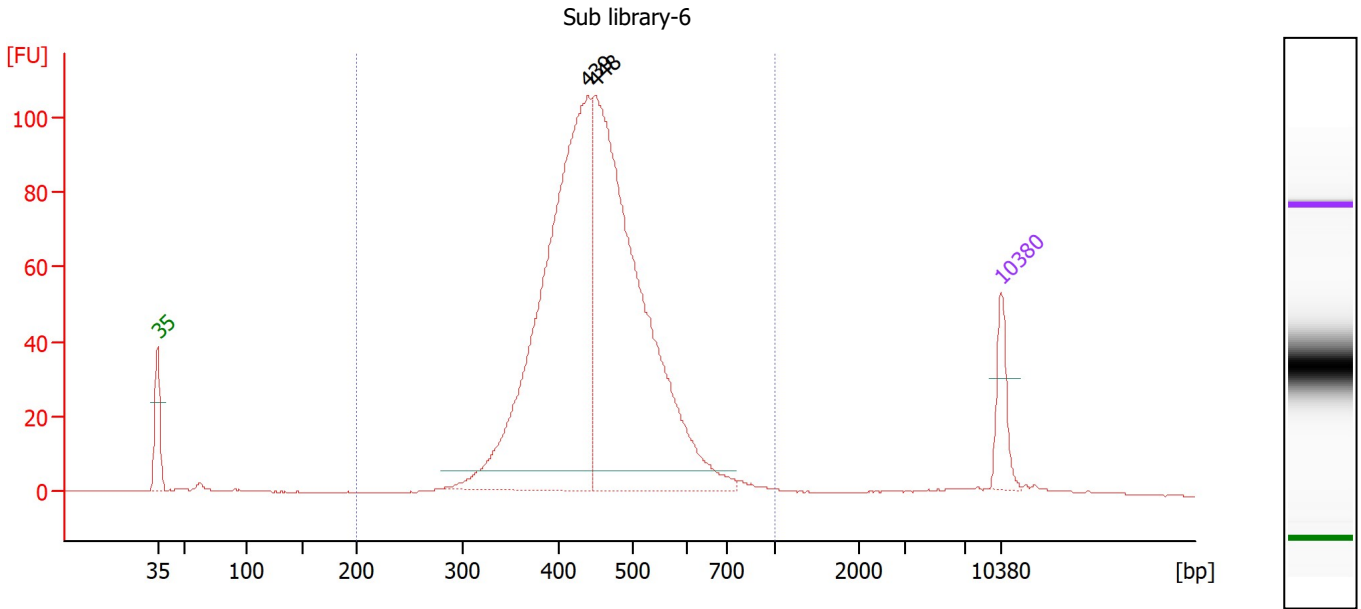
Region table for sample 7 : Sub library-5

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	905.1	95	456	17.7	2,036.58	7,004.3	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...alyzer\2100 expert\data\2024-12-26\CUSTOM0099 Libraries.xad

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Electropherogram Summary Continued ...



Overall Results for sample 8 : Sub library-6

Number of peaks found: 2 Corr. Area 1: 1,198.3
 Noise: 0.1

Peak table for sample 8 : Sub library-6

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	439	1,411.89	4,878.3	
3	448	1,427.33	4,822.3	
4	10,380	75.00	10.9	Upper Marker

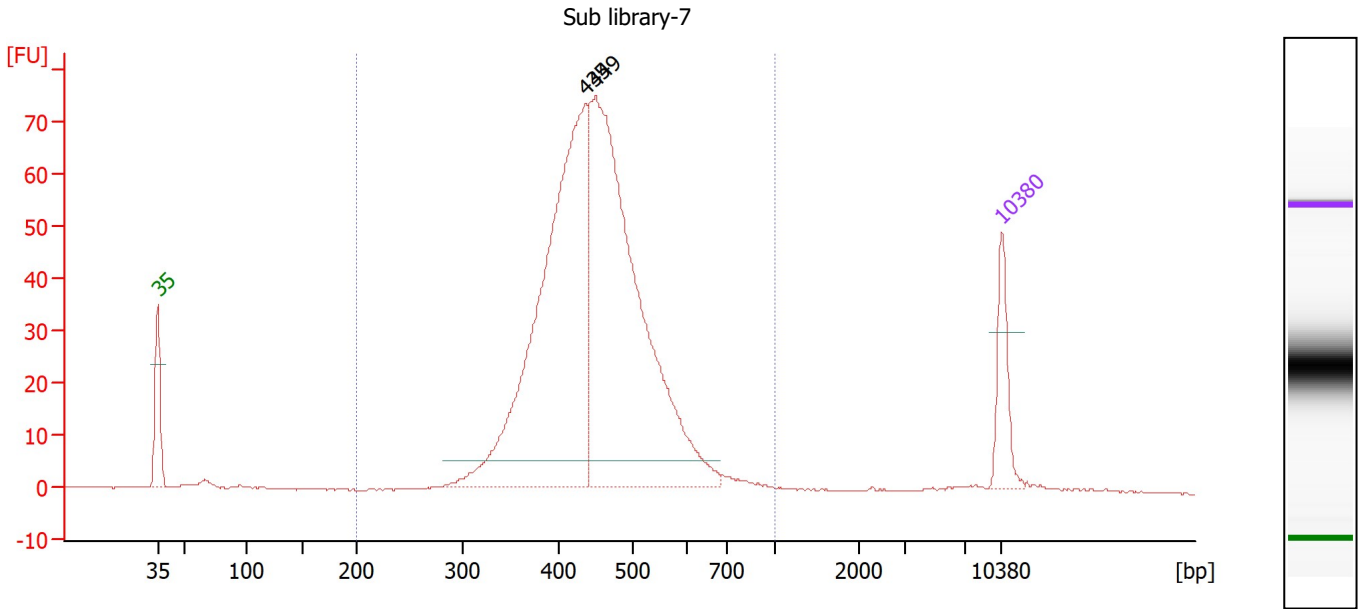
Region table for sample 8 : Sub library-6

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	1,198.3	96	456	17.4	2,944.45	10,129.3	Blue

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 Data Path: C:\...alyzer\2100 expert\data\2024-12-26\CUSTOM0099 Libraries.xad

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Electropherogram Summary Continued ...



Overall Results for sample 9 : Sub library-7

Number of peaks found: 2 Corr. Area 1: 829.3
 Noise: 0.2

Peak table for sample 9 : Sub library-7

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	435	985.70	3,436.5	
3	449	1,061.67	3,585.7	
4	10,380	75.00	10.9	Upper Marker

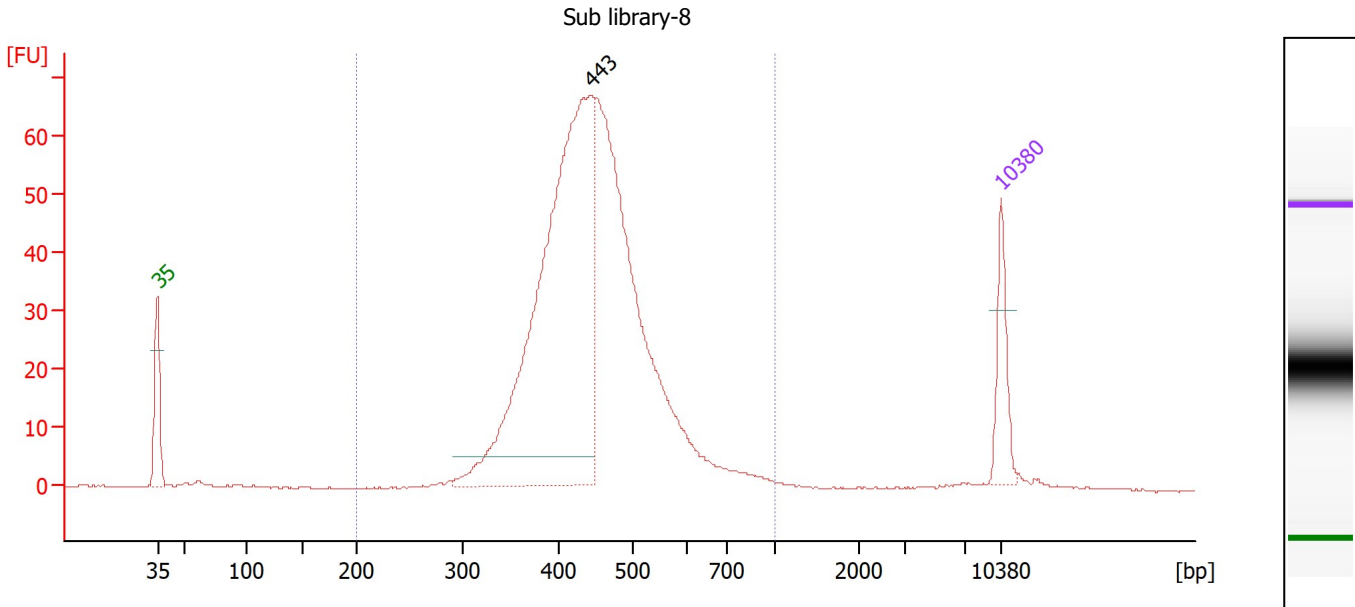
Region table for sample 9 : Sub library-7

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	829.3	96	453	17.2	2,147.71	7,418.7	Blue

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 Data Path: C:\...alyzer\2100 expert\data\2024-12-26\CUSTOM0099 Libraries.xad

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Electropherogram Summary Continued ...



Overall Results for sample 10 : Sub library-8

Number of peaks found: 1 Corr. Area 1: 760.5
 Noise: 0.1

Peak table for sample 10 : Sub library-8

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	443	1,126.38	3,850.9	
3	10,380	75.00	10.9	Upper Marker

Region table for sample 10 : Sub library-8

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	760.5	94	453	19.1	2,182.50	7,590.1	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...alyzer\2100 expert\data\2024-12-26\CUSTOM0099 Libraries.xad

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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		12/26/2024 2:05:40 PM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Run started on port 1 (File: C:\Program Files (x86)\Agilent\2100 bioanalyzer\2100 expert\data\2024-12-26\Bioanalyzer1_High Sensitivity DNA Assay_DE34903152_2024-12-26_002.xad)		Instrument	Run		12/26/2024 1:25:17 PM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Product Number : G2938C		Instrument	Run		12/26/2024 1:25:17 PM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Name :		Instrument	Run		12/26/2024 1:25:17 PM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Vendor : Agilent Technologies		Instrument	Run		12/26/2024 1:25:17 PM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Serial# : DE34903152		Instrument	Run		12/26/2024 1:25:17 PM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Firmware : C.01.069		Instrument	Run		12/26/2024 1:25:17 PM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Cartridge : Electrode		Instrument	Run		12/26/2024 1:25:17 PM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB