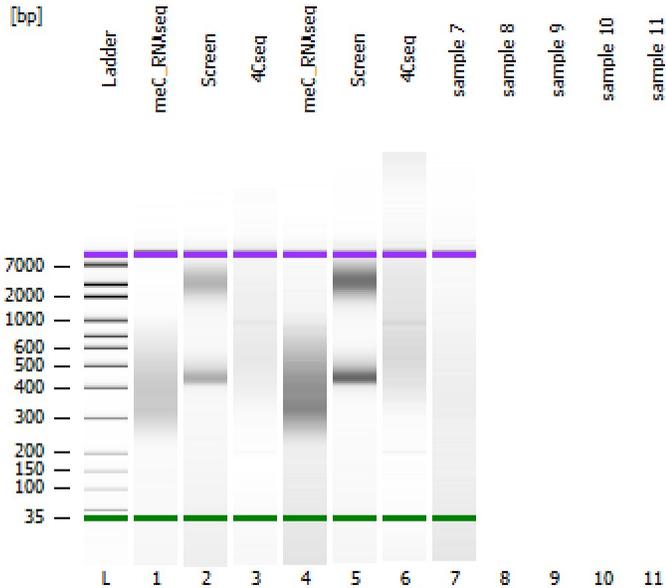


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...analyzer1_High Sensitivity DNA Assay_2023-04-20_001_Lib.xad

Created: 4/20/2023 2:28:05 PM
Modified: 4/20/2023 3:07:10 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

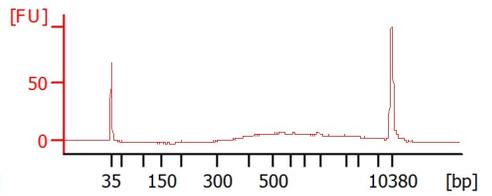
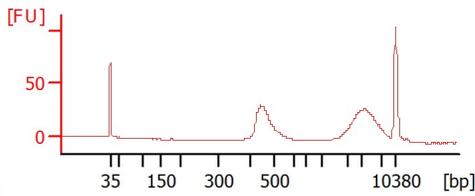
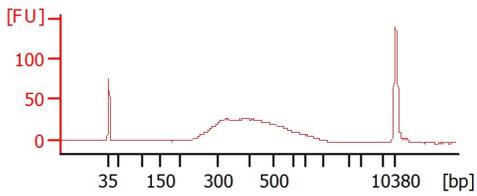
Chip Information:

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

meC_RNAseq

Screen

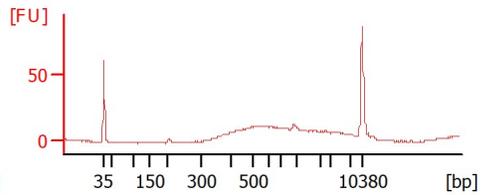
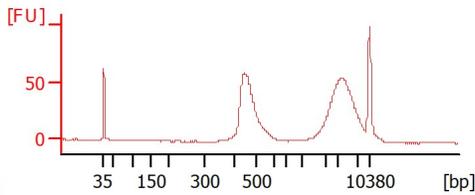
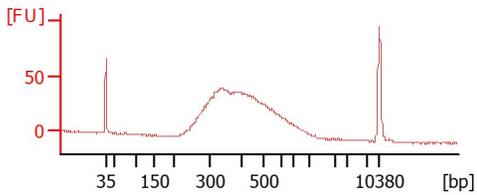
4Cseq



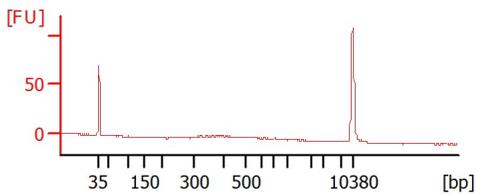
meC_RNAseq

Screen

4Cseq



sample 7



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...analyzer1_High Sensitivity DNA Assay_2023-04-20_001_Lib.xad

Created: 4/20/2023 2:28:05 PM
 Modified: 4/20/2023 3:07:10 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
meC_RNAseq		<input type="checkbox"/>	✓			
Screen		<input type="checkbox"/>	✓			
4Cseq		<input type="checkbox"/>	✓			
meC_RNAseq		<input type="checkbox"/>	✓			
Screen		<input type="checkbox"/>	✓			
4Cseq		<input type="checkbox"/>	✓			
sample 7		<input type="checkbox"/>	✓			
sample 8		<input type="checkbox"/>				
sample 9		<input type="checkbox"/>				
sample 10		<input type="checkbox"/>				
sample 11		<input type="checkbox"/>				
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...analyzer1_High Sensitivity DNA Assay_2023-04-20_001_Lib.xad

Created: 4/20/2023 2:28:05 PM
 Modified: 4/20/2023 3:07:10 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] : 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

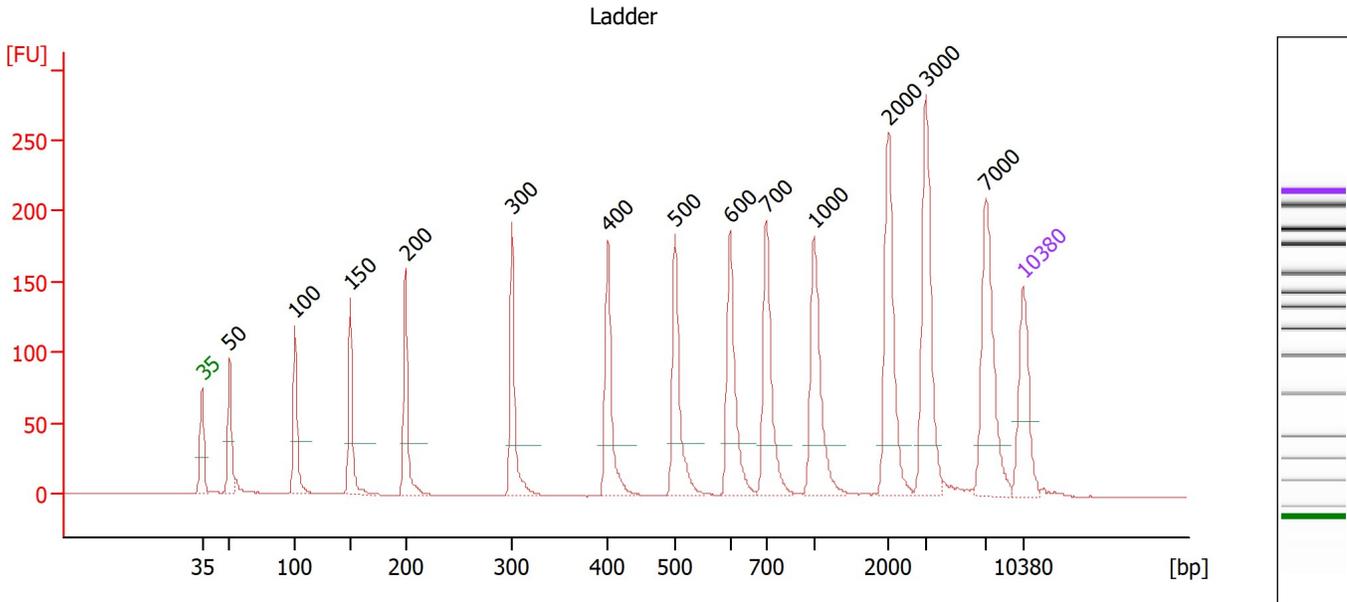
Ladder

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:\...analyzer1_High Sensitivity DNA Assay_2023-04-20_001_Lib.xad

Created: 4/20/2023 2:28:05 PM
 Modified: 4/20/2023 3:07:10 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.3

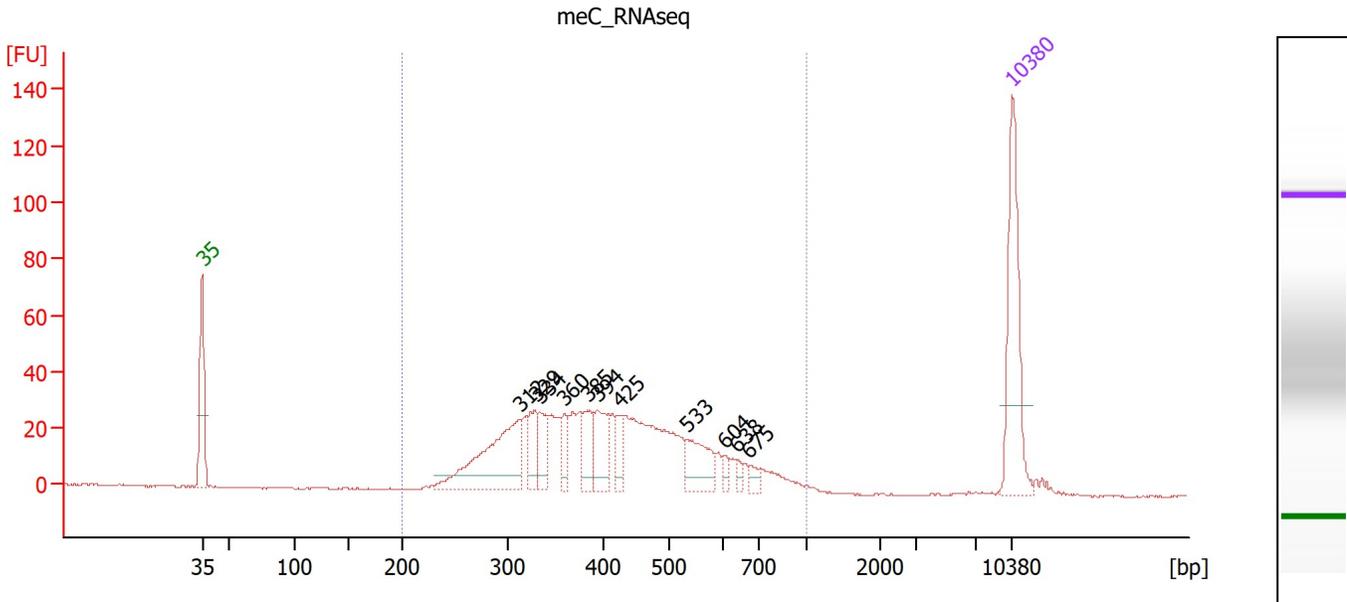
Peak table for Ladder

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:\...analyzer1_High Sensitivity DNA Assay_2023-04-20_001_Lib.xad

Created: 4/20/2023 2:28:05 PM
 Modified: 4/20/2023 3:07:10 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : meC_RNAseq

Number of peaks found: 11 Corr. Area 1: 726.9
 Noise: 0.2

Peak table for sample 1 : meC_RNAseq

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	312	115.71	561.4	
3	329	30.21	139.1	
4	334	25.10	114.0	
5	360	17.68	74.4	
6	385	28.66	112.9	
7	394	36.93	142.0	
8	425	15.96	56.9	
9	533	38.21	108.6	
10	604	6.13	15.4	
11	638	5.76	13.7	
12	675	7.39	16.6	
13	10,380	75.00	10.9	Upper Marker

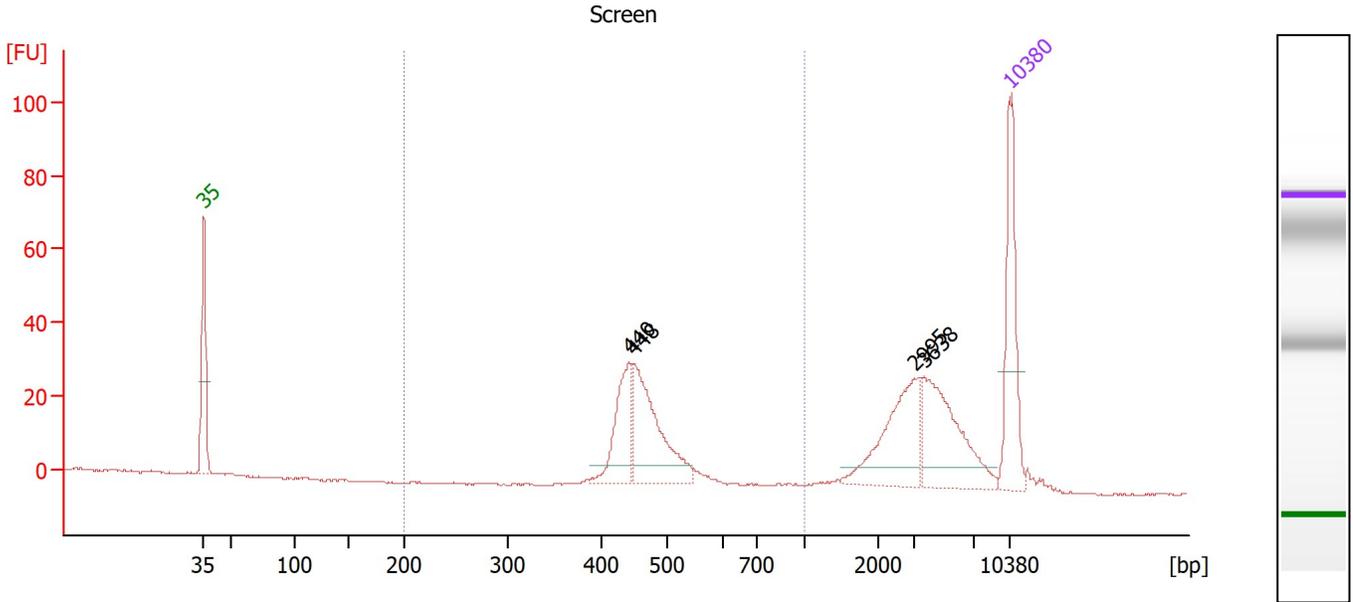
Region table for sample 1 : meC_RNAseq

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	726.9	99	430	30.0	583.37	2,291.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...analyzer1_High Sensitivity DNA Assay_2023-04-20_001_Lib.xad

Created: 4/20/2023 2:28:05 PM
 Modified: 4/20/2023 3:07:10 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : Screen

Number of peaks found: 4 Corr. Area 1: 173.1
 Noise: 0.2

Peak table for sample 2 : Screen

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	440	75.82	260.9	
3	448	118.99	402.8	
4	2,995	82.05	41.5	
5	3,638	89.68	37.4	
6	10,380	75.00	10.9	Upper Marker

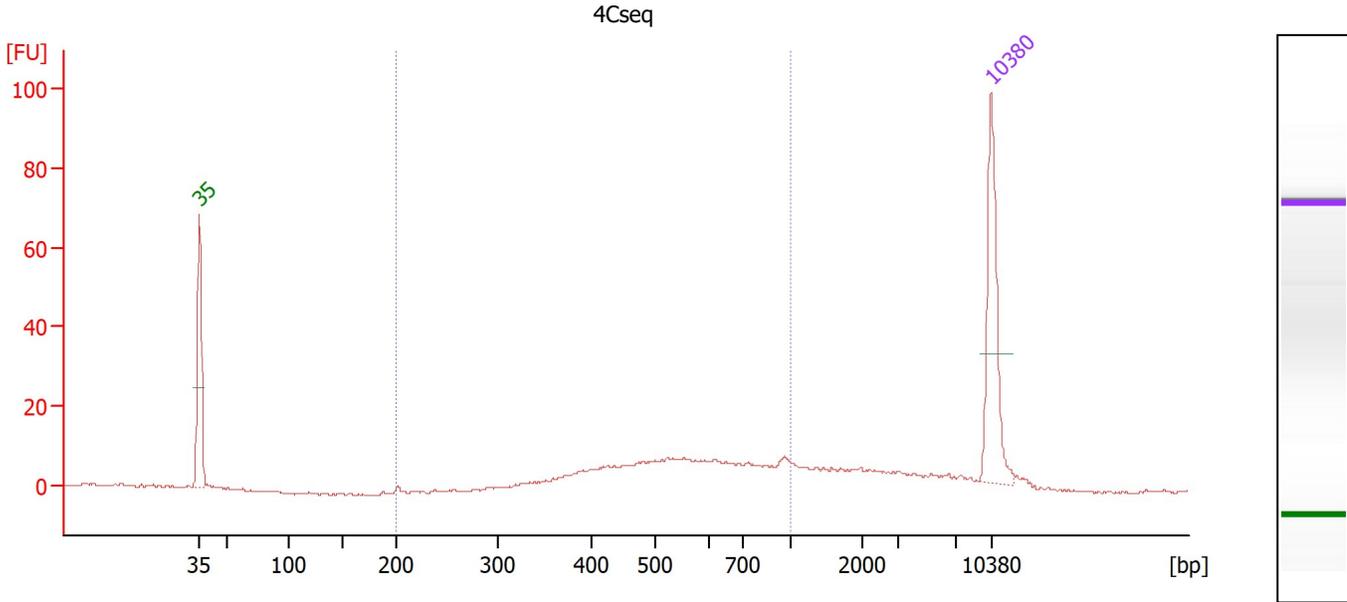
Region table for sample 2 : Screen

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	173.1	44	464	9.6	196.60	645.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...analyzer1_High Sensitivity DNA Assay_2023-04-20_001_Lib.xad

Created: 4/20/2023 2:28:05 PM
 Modified: 4/20/2023 3:07:10 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : 4Cseq

Number of peaks found: 0 Corr. Area 1: 148.9
 Noise: 0.2

Peak table for sample 3 : 4Cseq

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

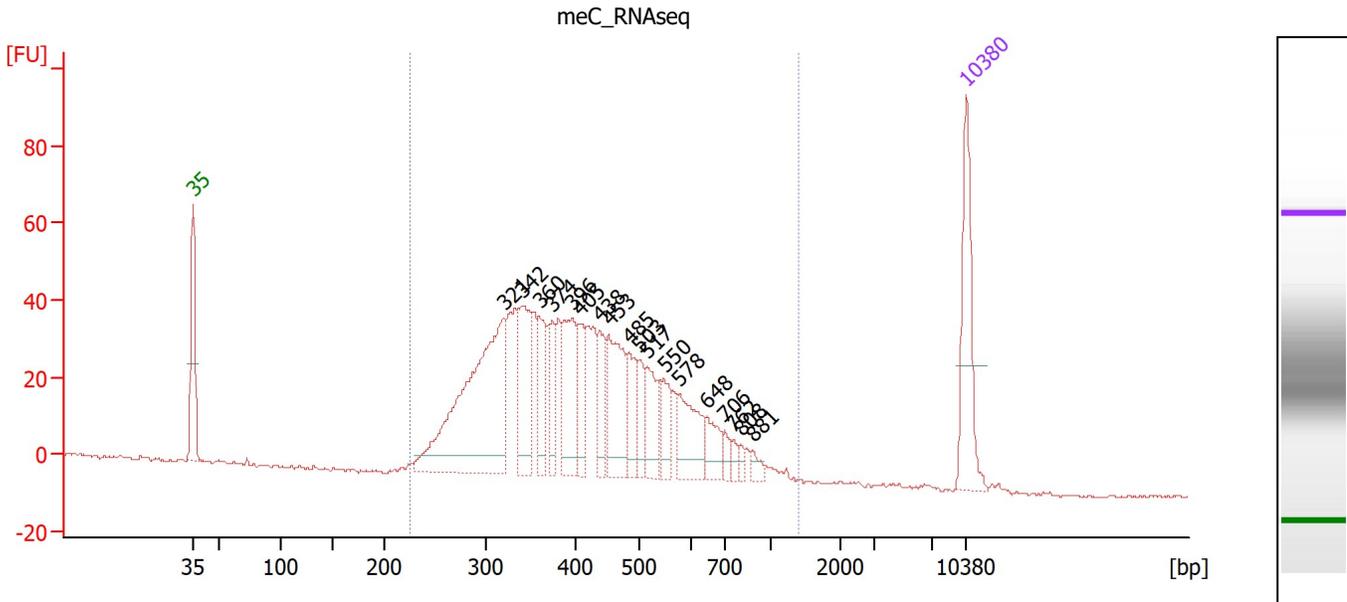
Region table for sample 3 : 4Cseq

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	148.9	66	581	28.7	176.44	509.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...analyzer1_High Sensitivity DNA Assay_2023-04-20_001_Lib.xad

Created: 4/20/2023 2:28:05 PM
 Modified: 4/20/2023 3:07:10 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : meC_RNAseq

Number of peaks found: 18 Corr. Area 1: 1,050.9
 Noise: 0.3

Peak table for sample 4 : meC_RNAseq

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	321	318.01	1,503.3	
3	342	92.78	411.5	
4	360	49.80	209.7	
5	374	41.10	166.3	
6	396	91.60	350.3	
7	405	38.31	143.1	
8	438	43.35	149.8	
9	453	94.59	316.6	
10	485	44.38	138.7	
11	503	29.03	87.5	
12	517	46.04	134.9	
13	550	31.91	87.9	
14	578	59.80	156.6	
15	648	28.77	67.2	
16	706	10.41	22.3	
17	762	7.59	15.1	
18	808	6.60	12.4	
19	881	7.94	13.7	
20	10,380	75.00	10.9	Upper Marker

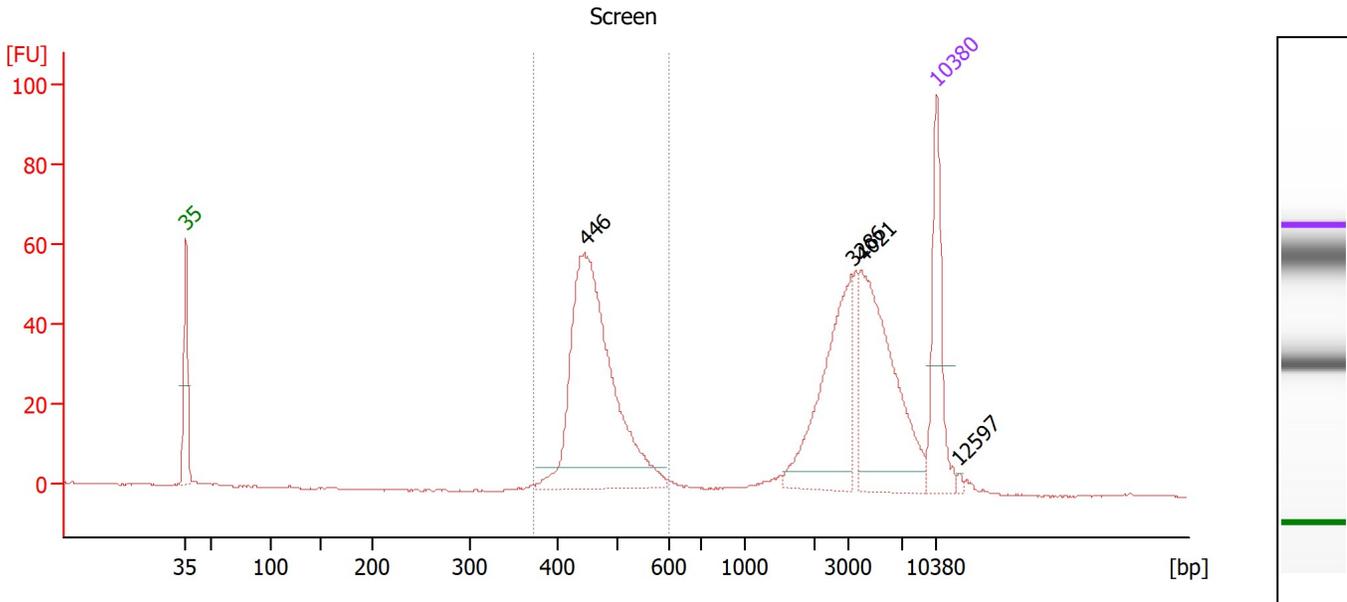
Region table for sample 4 : meC_RNAseq

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
225	1,403	1,050.9	100	434	32.6	1,310.90	5,144.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...analyzer1_High Sensitivity DNA Assay_2023-04-20_001_Lib.xad

Created: 4/20/2023 2:28:05 PM
 Modified: 4/20/2023 3:07:10 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : Screen

Number of peaks found: 4 Corr. Area 1: 332.3
 Noise: 0.2

Peak table for sample 5 : Screen

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	446	375.60	1,275.4	
3	3,286	137.12	63.2	
4	4,021	162.45	61.2	
5	10,380	75.00	10.9	Upper Marker
6	12,597	0.00	0.0	

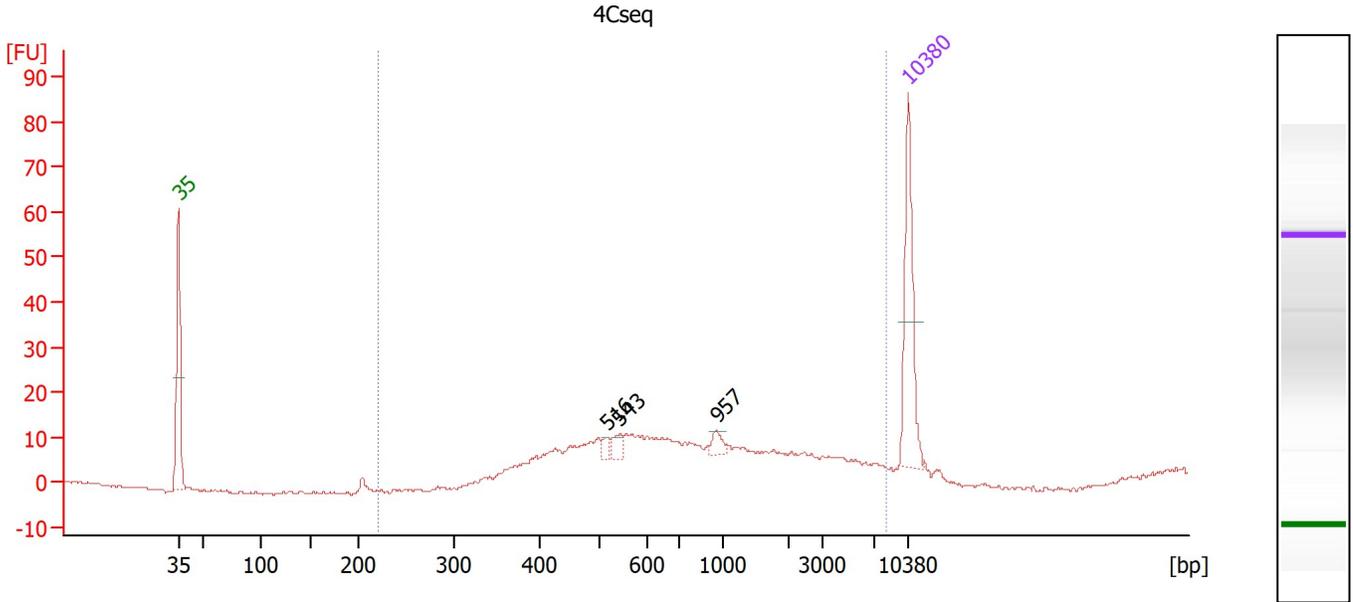
Region table for sample 5 : Screen

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
374	604	332.3	45	466	8.5	374.22	1,225.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...analyzer1_High Sensitivity DNA Assay_2023-04-20_001_Lib.xad

Created: 4/20/2023 2:28:05 PM
 Modified: 4/20/2023 3:07:10 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : 4Cseq

Number of peaks found: 3 Corr. Area 1: 224.7
 Noise: 0.3

Peak table for sample 6 : 4Cseq

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	516	7.56	22.2	
3	543	9.80	27.3	
4	957	8.81	13.9	
5	10,380	75.00	10.9	Upper Marker

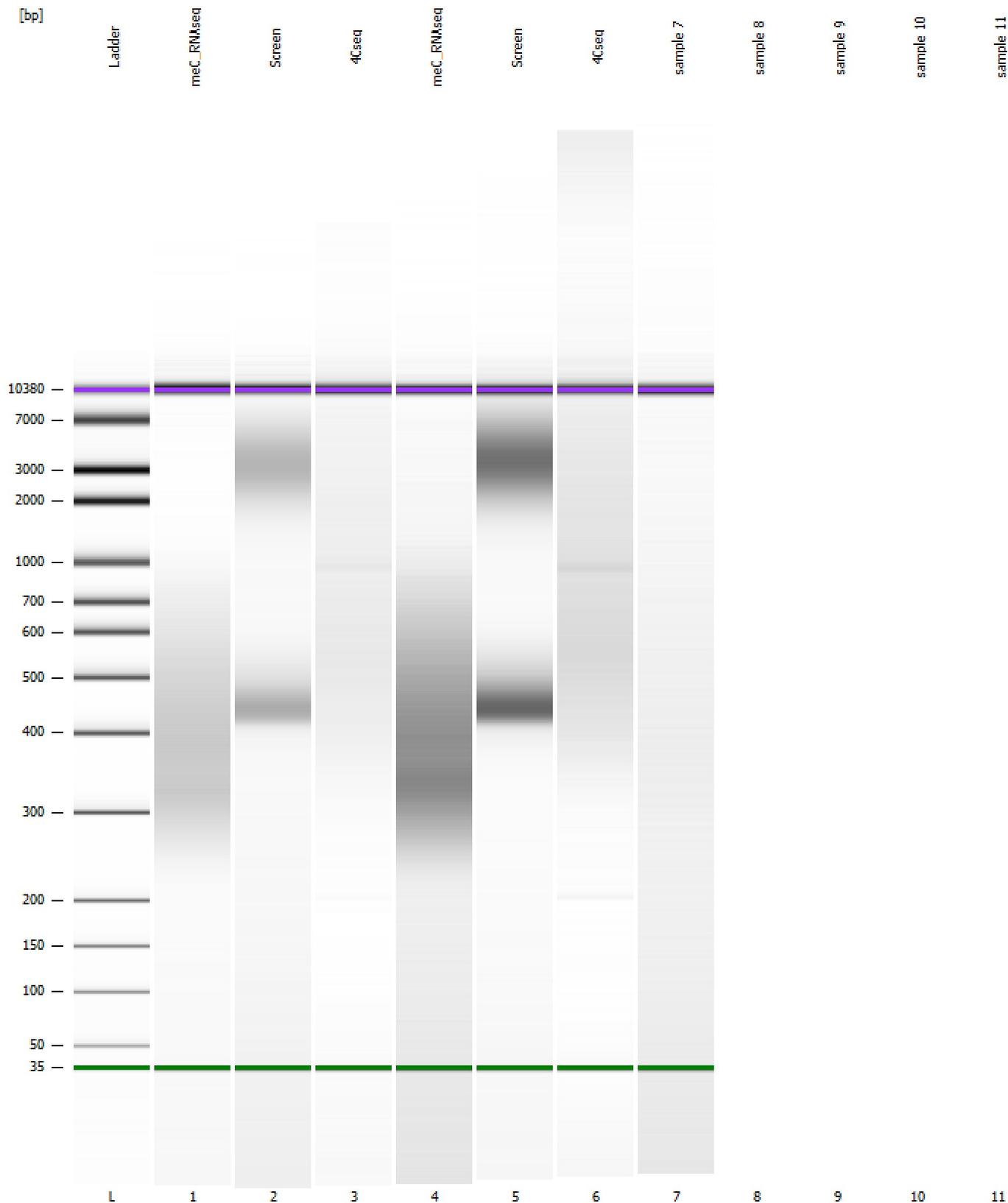
Region table for sample 6 : 4Cseq

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
220	8,070	224.7	99	1,211	100.0	334.23	772.7	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...analyzer1_High Sensitivity DNA Assay_2023-04-20_001_Lib.xad

Created: 4/20/2023 2:28:05 PM
Modified: 4/20/2023 3:07:10 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...analyzer1_High Sensitivity DNA Assay_2023-04-20_001_Lib.xad

Created: 4/20/2023 2:28:05 PM
 Modified: 4/20/2023 3:07:10 PM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
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Run started on port 1 (File: C:\Users\sbsuser\Desktop\2023-04-20\Bioanalyzer1_High Sensitivity DNA Assay_2023-04-20_001.xad)		Instrument	Run		4/20/2023 2:28:11 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Product Number : G2938C		Instrument	Run		4/20/2023 2:28:11 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Name :		Instrument	Run		4/20/2023 2:28:11 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Vendor : Agilent Technologies		Instrument	Run		4/20/2023 2:28:11 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Serial# : DE34903152		Instrument	Run		4/20/2023 2:28:11 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Firmware : C.01.069		Instrument	Run		4/20/2023 2:28:11 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Cartridge : Electrode		Instrument	Run		4/20/2023 2:28:11 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB