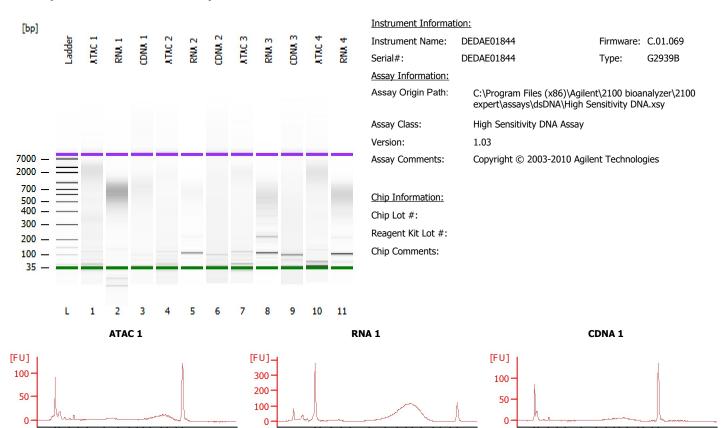
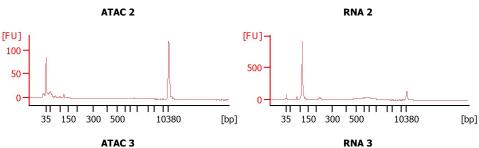
Assay Class: High Sensitivity DNA Assay Created: 23/09/2021 15:46:53
Data Path: C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-09-23_15-46-53.xad Created: 23/09/2021 16:28:15

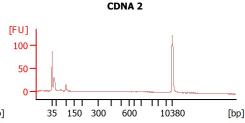
Electrophoresis File Run Summary





[bp]

35 150

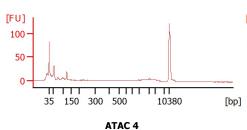


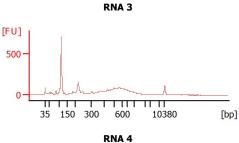
500

10380

[bp]

35 150 300

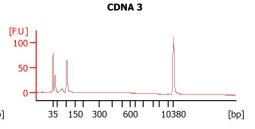


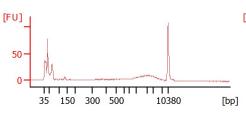


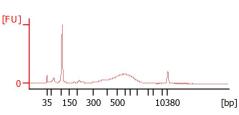
300

500

10380[bp]







35 150 300

500

10380

Page 2 of 17

High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-09-23_15-46-53.xad Assay Class: Created: 23/09/2021 15:46:53 Modified: Data Path: 23/09/2021 16:28:15 **Electrophoresis File Run Summary (Chip Summary) Sample Name Sample Comment** Rest. Digest Status Observation **Result Label** Res ult Col or ATAC 1 RNA 1 CDNA 1 ATAC 2 RNA 2 CDNA 2 ATAC 3 RNA 3 CDNA 3 ATAC 4 RNA 4 Ladder

Reagent Kit Lot #

Chip Comments:

Chip Lot #

Page 3 of 17

Assay Class: High Sensitivity DNA Assay Created: 23/09/2021 15:46:53
Data Path: C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-09-23_15-46-53.xad C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-09-23_15-46-53.xad Modified: 23/09/2021 16:28:15

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

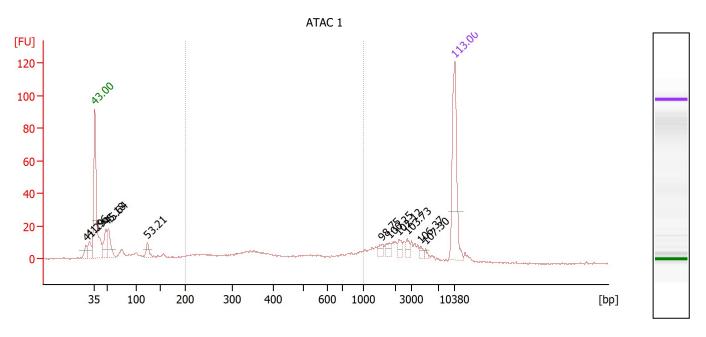
Filter Settings

Filter Width [s]: 0.5 Polynomial Order: 4

Ladder

Size	Area
35	160
50	210
100	208
150	221
200	242
300	270
400	305
500	306
600	336
700	321
1000	366
2000	413
3000	411
7000	400
10380	214
	35 50 100 150 200 300 400 500 600 700 1000 2000 3000 7000

Electropherogram Summary



Overall Results for sample 1 : ATAC 1

Number of peaks found: 11 Corr. Area 1: 168.7

Noise: 0.4

Peak ta	ble	for sample 1:	ATAC 1		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		29	0.00	0.0	
2		31	0.00	0.0	
3	◀	35	125.00	5,411.3	Lower Marker
4		49	38.41	1,194.8	
5		52	41.47	1,201.3	
6		123	16.10	198.4	
7		1,560	6.75	6.6	
8		1,800	6.99	5.9	
9		2,194	6.90	4.8	
10		2,703	6.61	3.7	
11		4,330	3.85	1.3	
12		5,057	2.53	0.8	
13		10,380	75.00	10.9	Upper Marker

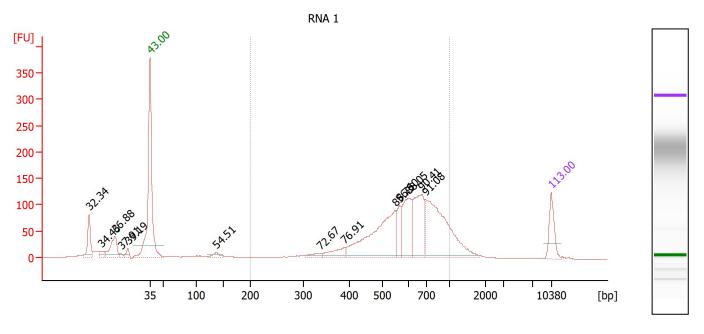
Region table for sample 1: ATAC 1

From	To [bp] Corr.		Average Size	Size distribution in		Molarity	Co
[bp]		Area	Total	[bp]	CV [%]	[pg/µl]	[pmol/l]	lor
200	1,000	168.7	34	465	44.8	174.22	731.6	

Page 5 of 17

Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-09-23_15-46-53.xad Created: 23/09/2021 15:46:53 Modified: 23/09/2021 16:28:15

Electropherogram Summary Continued ...



Overall Results for sample 2: RNA 1

Number of peaks found: 13 Corr. Area 1: 1,449.3

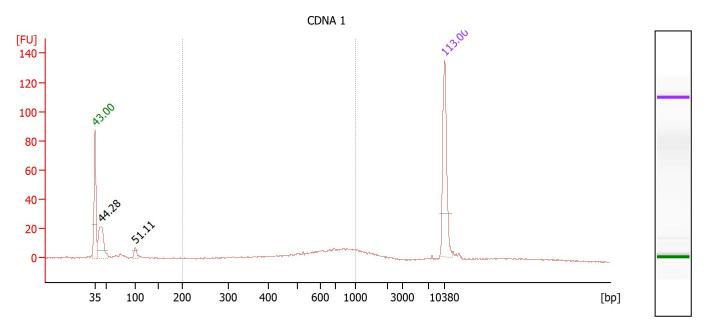
Noise: 0.4

Peak ta	able	for sample 2:	<u>RNA 1</u>		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		0	0.00	0.0	
2		5	0.00	0.0	
3		14	0.00	0.0	
4		17	0.00	0.0	
5		22	0.00	0.0	
6	18	35	125.00	5,411.3	Lower Marker
7		137	28.88	320.5	
8		337	25.78	116.0	
9		390	64.58	251.1	
10		548	434.62	1,201.7	
11		566	73.90	197.7	
12		598	197.80	501.2	
13		675	247.16	554.4	
14		698	485.52	1,054.5	
15	B	10,380	75.00	10.9	Upper Marker

Region table for sample 2: RNA 1

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	1 000 1 449 3	73	612	26.3	1 524 97	4 185 3	

Electropherogram Summary Continued ...



Overall Results for sample 3: <u>CDNA 1</u>

Number of peaks found: 2 Corr. Area 1: 126.8

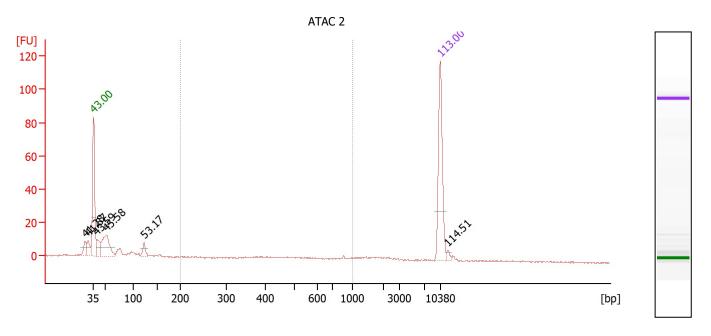
Noise: 0.4

Peak t	able	for sample 3:	CDNA 1					
Peak	Peak Size [bp]		eak Size [bp] Conc. [pg/µ		Conc. [pg/µl]	Molarity [pmol/l]] Observations	
1	4	35	125.00	5,411.3	Lower Marker			
2		43	86.68	3,049.8				
3		101	13.07	196.2				
4		10.380	75.00	10.9	Upper Marker			

Region table for sample 3: CDNA 1

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	1 000 126 8	41	599	32.4	118 09	358 2	

Electropherogram Summary Continued ...



Overall Results for sample 4: <u>ATAC 2</u>

Number of peaks found: 6 Corr. Area 1: 18.0

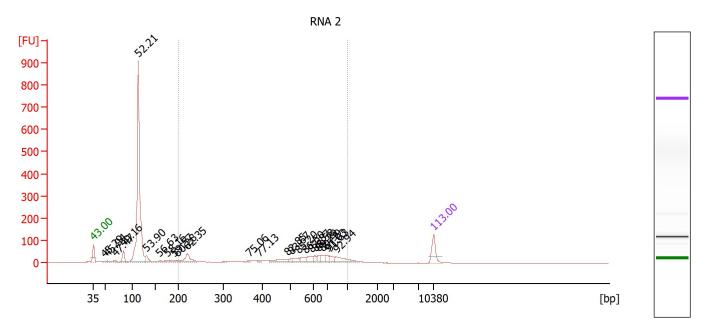
Noise: 0.4

Peak ta	able	for sample 4:	ATAC 2		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		29	0.00	0.0	
2		31	0.00	0.0	
3	4	35	125.00	5,411.3	Lower Marker
4		39	29.44	1,152.0	
5		52	74.19	2,170.8	
6		123	16.09	199.0	
7		10,380	75.00	10.9	Upper Marker
8		11,966	0.00	0.0	

Region table for sample 4: ATAC 2

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	1,000	18.0	10	524	37.6	19.43	68.5	

Electropherogram Summary Continued ...



Overall Results for sample 5: RNA 2

Number of peaks found: 24 Corr. Area 1: 685.3

Noise: 0.4

able	for sample 5:	<u>RNA 2</u>		
	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
4	35	125.00	5,411.3	Lower Marker
	49	12.49	383.0	
	55	15.30	421.6	
	69	27.04	598.0	
	83	79.58	1,444.4	
	112	1,550.45	20,886.4	
	130	70.67	822.4	
	159	19.84	189.2	
	175	18.19	157.6	
	190	24.57	196.2	
	197	10.22	78.6	
	220	101.67	701.3	
	367	8.48	35.1	
	392	6.87	26.5	
	488	49.40	153.3	
	503	12.78	38.5	
	536	30.23	85.5	
	564	29.78	80.0	
	594	37.88	96.6	
	614	21.37	52.7	
	638	22.03	52.3	
	661	31.62	72.4	
	696	22.40	48.8	
	735	32.27	66.5	
	831	74.44	135.7	
	10,380	75.00	10.9	Upper Marker
	•	 ■ 35 49 55 69 83 112 130 159 175 190 197 220 367 392 488 503 536 564 594 614 638 661 696 735 831 	Size [bp] Conc. [pg/µl] 35 125.00 49 12.49 55 15.30 69 27.04 83 79.58 112 1,550.45 130 70.67 159 19.84 175 18.19 190 24.57 197 10.22 220 101.67 367 8.48 392 6.87 488 49.40 503 12.78 536 30.23 564 29.78 594 37.88 614 21.37 638 22.03 661 31.62 696 22.40 735 32.27 831 74.44	Size [bp] Conc. [pg/μl] Molarity [pmol/l] 49 12.49 383.0 55 15.30 421.6 69 27.04 598.0 83 79.58 1,444.4 112 1,550.45 20,886.4 130 70.67 822.4 159 19.84 189.2 175 18.19 157.6 190 24.57 196.2 197 10.22 78.6 220 101.67 701.3 367 8.48 35.1 392 6.87 26.5 488 49.40 153.3 503 12.78 38.5 536 30.23 85.5 564 29.78 80.0 594 37.88 96.6 614 21.37 52.7 638 22.03 52.3 661 31.62 72.4 696 22.40 48.8 735 32.27 66.5 831 74.44 135.7

2100 expert_High Sensitivity DNA Assay_DEDAE01844_2021-09-23_15-46-53.xad

Page 9 of 17

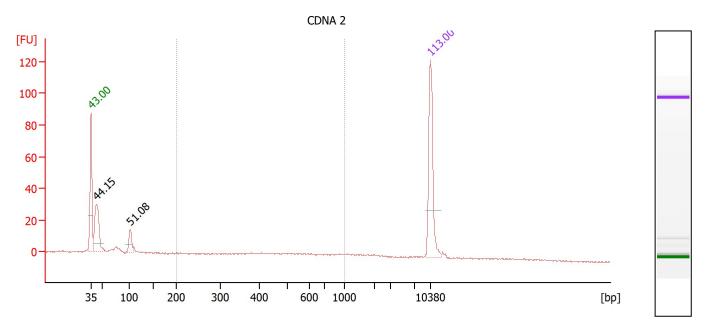
Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-09-23_15-46-53.xad Created: 23/09/2021 15:46:53 Modified: 23/09/2021 16:28:15

Electropherogram Summary Continued ...

... Region table for sample 5 : RNA 2

From To [bp] Corr. % of **Average Size** Size distribution in Conc. Molarity Co Area Total [bp] **CV** [%] [pg/µl] [pmol/l] lor [bp] 200 1,000 685.3 33 532 38.0 708.91 2,680.3

Electropherogram Summary Continued ...



Overall Results for sample 6: <u>CDNA 2</u>

Number of peaks found: 2 Corr. Area 1: 50.5

Noise: 0.5

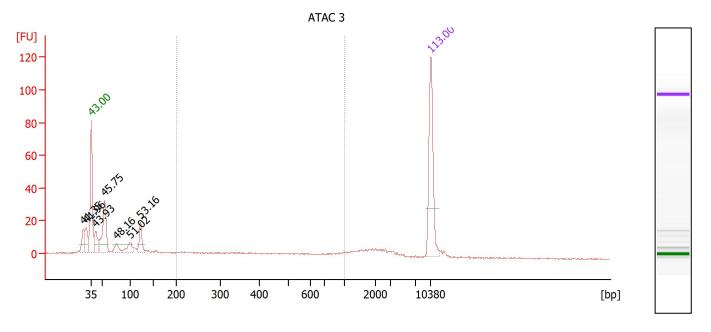
Peak tab	le for sample 6:	CDNA 2
Peak	Size [bp]	Conc. [pg/µl]

Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2		42	102.48	3,673.6	
3		101	25.34	381.9	
4		10,380	75.00	10.9	Upper Marker

Region table for sample 6: CDNA 2

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	1.000 50.5	23	512	37.9	49.16	180.1	

Electropherogram Summary Continued ...



Overall Results for sample 7: <u>ATAC 3</u>

Number of peaks found: 7 Corr. Area 1: 59.0

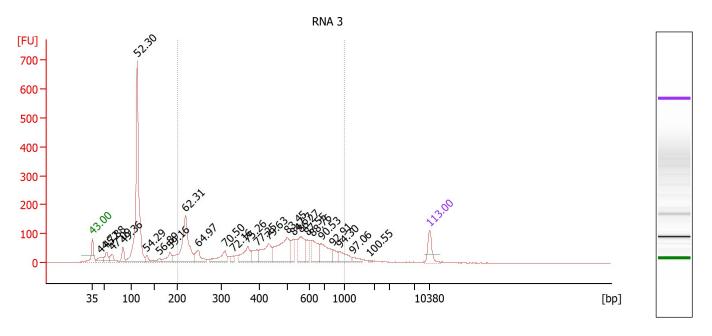
Noise: 0.3

•	Peak t	able	for sample 7	: <u>ATAC 3</u>		
ı	Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	1		29	0.00	0.0	
2	2		31	0.00	0.0	
3	3	- ◀	35	125.00	5,411.3	Lower Marker
4	4		41	34.94	1,295.1	
į	5		53	88.95	2,532.3	
6	5		75	18.65	378.6	
7	7		100	21.35	323.8	
8	3		122	32.06	396.9	
9	9		10,380	75.00	10.9	Upper Marker

Region table for sample 7: ATAC 3

From	To [bp] Corr.	% of Average Size	Size distribution in	Conc.	Molarity	Co	
[bp]	Area	Total	[bp]	CV [%]	[pg/µl]	[pmol/l]	lor
200	1 000 59 0	18	465	45 4	61 59	262 4	

Electropherogram Summary Continued ...



Overall Results for sample 8: RNA 3

Number of peaks found: 25 Corr. Area 1: 2,188.3

Noise: 0.4

Peak table	for sample 8:	<u>RNA 3</u>		
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	46	85.84	2,839.4	
3	54	88.00	2,451.3	
4	65	94.30	2,192.1	
5	85	95.62	1,698.9	
6	113	1,453.27	19,426.8	
7	134	53.75	606.7	
8	162	36.32	340.7	
9	185	101.19	827.1	
10	219	552.70	3,820.2	
11	248	120.18	733.7	
12	310	129.34	632.6	
13	330	28.91	132.5	
14	369	147.16	604.1	
15	394	86.38	332.2	
16	433	197.67	692.3	
17	498	315.03	957.6	
18	524	73.28	211.7	
19	559	164.45	445.5	
20	587	61.55	158.9	
21	620	126.83	309.7	
22	679	153.09	341.4	
23	829	55.80	101.9	
24	931	79.08	128.7	
25	1,291	29.71	34.9	
26	1,847	4.65	3.8	

Page 13 of 17

Assay Class: High Sensitivity DNA Assay

Created: 23/09/2021 15:46:53 C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-09-23_15-46-53.xad Data Path: Modified: 23/09/2021 16:28:15

Electropherogram Summary Continued ...

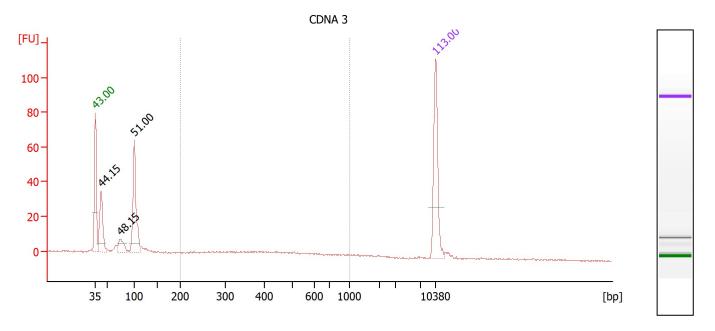
... Peak table for sample 8: <u>RNA 3</u>

Peak Size [bp] Molarity [pmol/l] **Observations** Conc. [pg/µl] 27 10,380 75.00 10.9 Upper Marker

Region table for sample 8: RNA 3

From To [bp] Corr. % of **Average Size** Size distribution in Conc. Molarity Co **CV** [%] [bp] Area Total [bp] [pg/µl] [pmol/l] lor 200 1,000 2,188.3 59 476 38.7 2,415.93 9,946.3

Electropherogram Summary Continued ...



Overall Results for sample 9: <u>CDNA 3</u>

Number of peaks found: 3 Corr. Area 1: 50.2

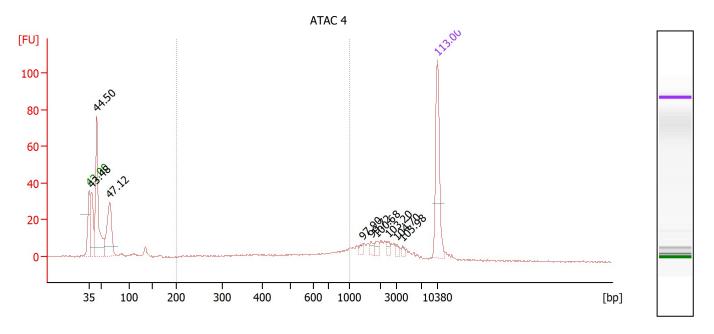
Noise: 0.3

Peak table for sample 9:			: <u>CDNA 3</u>		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2		42	95.24	3,415.5	
3		75	30.73	624.5	
4		100	145.06	2,202.2	
5		10,380	75.00	10.9	Upper Marker

Region table for sample 9: <u>CDNA 3</u>

From [bp]	To [bp] Corr Area		Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	1,000 50.2	19	439	37.4	55.83	229.5	

Electropherogram Summary Continued ...



Overall Results for sample 10 : ATAC 4

Number of peaks found: 9 Corr. Area 1: 75.2

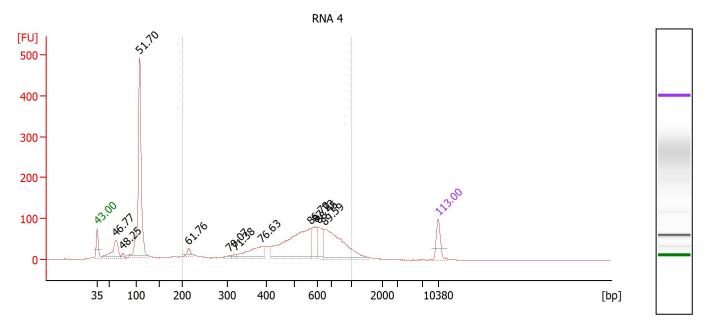
Noise: 0.4

Peak	Peak table for sample 10: <u>ATAC 4</u>									
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations					
1	•	35	125.00	5,411.3	Lower Marker					
2		38	87.77	3,496.2						
3		44	206.28	7,032.4						
4		65	123.43	2,858.1						
5		1,424	4.76	5.1						
6		1,715	5.83	5.1						
7		1,868	6.16	5.0						
8		2,536	5.62	3.4						
9		3,027	3.26	1.6						
10		4,030	3.03	1.1						
11		10,380	75.00	10.9	Upper Marker					

Region table for sample 10: <u>ATAC 4</u>

From	To [bp] Corr.	% of	Average Size	Size distribution in CV [%]	Conc.	Molarity	Co
[bp]	Area	Total	[bp]		[pg/µl]	[pmol/l]	lor
200	1 000 75 2	17	560	38 7	88 85	306.7	

Electropherogram Summary Continued ...



Overall Results for sample 11: RNA 4

Number of peaks found: 11 Corr. Area 1: 1,422.9

Noise: 0.4

Peak table for sample 11: RNA 4									
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations					
1	35	125.00	5,411.3	Lower Marker					
2	62	236.65	5,752.5						
3	75	24.30	488.3						
4	107	1,156.58	16,372.2						
5	213	36.73	261.1						
6	304	13.85	68.9						
7	321	15.76	74.4						
8	386	168.57	661.3						
9	568	540.04	1,439.4						
10	591	107.96	276.8						
11	610	104.82	260.4						
12	648	415.57	971.5						
13	10,380	75.00	10.9	Upper Marker					

Region table for sample 11: RNA 4

From	To [bp	[bp] Corr. % of Ave	Average Size	Average Size Size distribution in	Conc.	Molarity	Co	
[bp]		Area	Total	[bp]	CV [%]	[pg/µl]	[pmol/l]	lor
200	1,000	1,422.9	58	547	30.5	1,684.35	5,514.5	

High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-09-23_15-46-53.xad 23/09/2021 15:46:53 23/09/2021 16:28:15 Assay Class: Created: Modified: Data Path: **Gel Image** [bp] RNA 4 10380 7000 3000 2000 -1000 -700 -600 -500 — 400 - -300 - -200 - -150 - -100 -50 -35 — 1 3 5 6 10 11