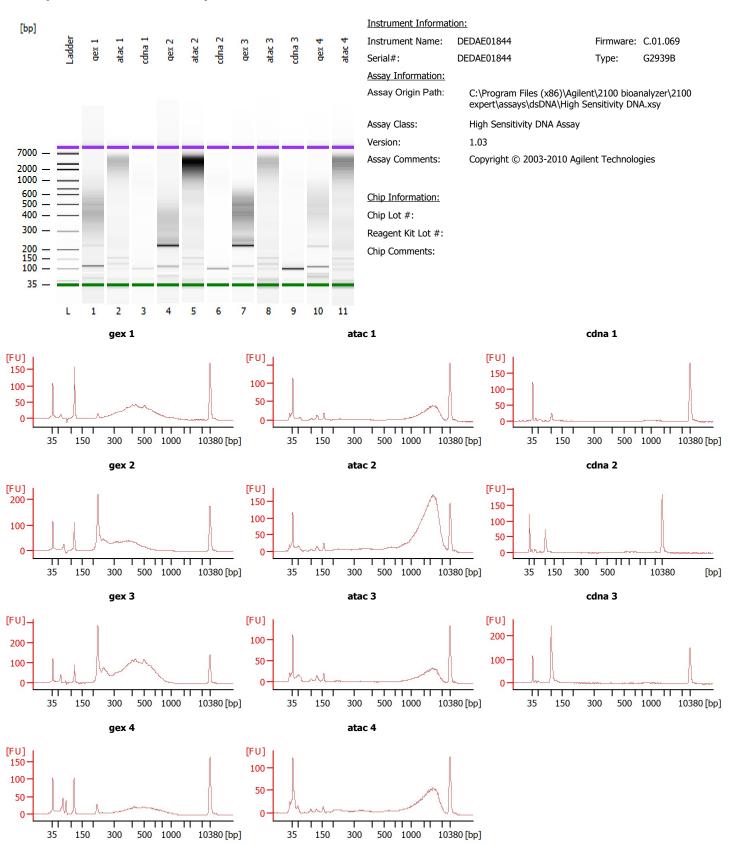
Assay Class: High Sensitivity DNA Assay Created: 28/10/2021 10:55:44
Data Path: C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-10-28_10-55-45.xad Created: 28/10/2021 11:46:18

Electrophoresis File Run Summary



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High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-10-28_10-55-45.xad Assay Class: Created: 28/10/2021 10:55:44 Modified: Data Path: 28/10/2021 11:46:18 **Electrophoresis File Run Summary (Chip Summary) Sample Name Sample Comment** Rest. Digest Status Observation **Result Label** Res ult Col or gex 1 atac 1 cdna 1 gex 2 atac 2 cdna 2 gex 3 atac 3 cdna 3 gex 4 atac 4 Ladder Chip Lot # Reagent Kit Lot #

Chip Comments:

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Assay Class: High Sensitivity DNA Assay Created: 28/10/2021 10:55:44
Data Path: C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-10-28_10-55-45.xad Created: 28/10/2021 11:46:18

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

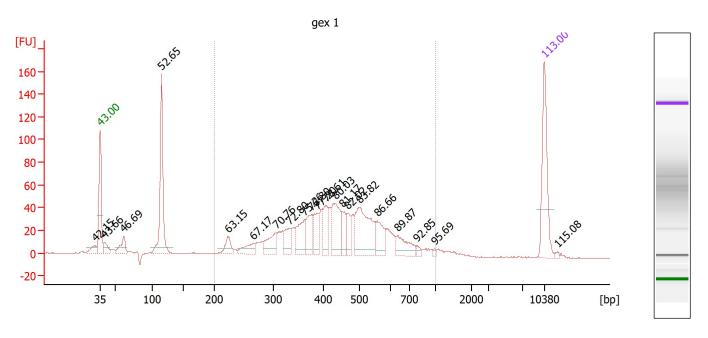
Filter Settings

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

Electropherogram Summary



Overall Results for sample 1: gex 1

Number of peaks found: 21 Corr. Area 1: 1,027.6

Noise: 0.3

Peak to	able for sample	e 1 : <u>gex 1</u>		
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	32	0.00	0.0	
2	◀ 35	125.00	5,411.3	Lower Marker
3	39	21.32	823.7	
4	61	32.41	798.9	
5	115	183.41	2,416.0	
6	223	24.86	168.6	
7	267	25.54	145.1	
8	306	38.17	188.8	
9	332	31.09	142.0	
10	363	47.89	200.1	
11	376	40.90	165.0	
12	387	35.55	139.2	
13	407	36.26	135.1	
14	432	67.82	238.0	
15	452	27.68	92.8	
16	467	26.17	84.9	
17	499	111.39	338.5	
18	556	37.90	103.2	
19	637	32.67	77.8	
20	783	5.18	10.0	
21	993	2.87	4.4	
22	1 0,380	75.00	10.9	Upper Marker
23	12,423	0.00	0.0	

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Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-10-28_10-55-45.xad Created: 28/10/2021 10:55:44 Modified: 28/10/2021 11:46:18

Electropherogram Summary Continued ...

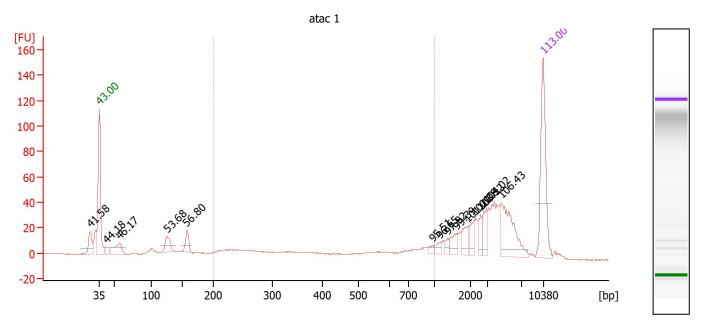
... Region table for sample 1: gex 1

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	1,000	1,027.6	76	450	29.8	792.77	3,020.8	

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Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-10-28_10-55-45.xad Created: 28/10/2021 10:55:44 Modified: 28/10/2021 11:46:18

Electropherogram Summary Continued ...



Overall Results for sample 2: <u>atac 1</u>

Number of peaks found: 14 Corr. Area 1: 164.2

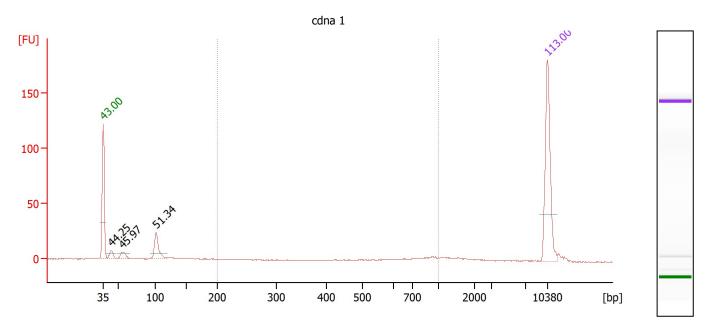
Noise: 0.3

Peak t	able	for sample 2:	atac 1		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		30	0.00	0.0	
2	4	35	125.00	5,411.3	Lower Marker
3		43	12.48	444.7	
4		57	33.49	890.7	
5		125	27.89	336.8	
6		157	21.02	202.8	
7		980	3.94	6.1	
8		1,151	5.86	7.7	
9		1,375	5.24	5.8	
10		1,632	12.62	11.7	
11		1,930	16.05	12.6	
12		2,212	12.29	8.4	
13		2,586	11.08	6.5	
14		2,909	17.50	9.1	
15		4,633	59.63	19.5	
16		10,380	75.00	10.9	Upper Marker

Region table for sample 2: <u>atac 1</u>

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	1.000 164.2	23	504	47.4	139.88	586.2	

Electropherogram Summary Continued ...



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

Number of peaks found: 3 Corr. Area 1: 27.3

Noise: 0.2

Peak	Peak table for sample 3:		<u>cdna 1</u>			
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations	
1	4	35	125.00	5,411.3	Lower Marker	
2		43	13.28	468.8		
3		55	14.45	396.1		
4		102	39.66	590.1		
5		10,380	75.00	10.9	Upper Marker	

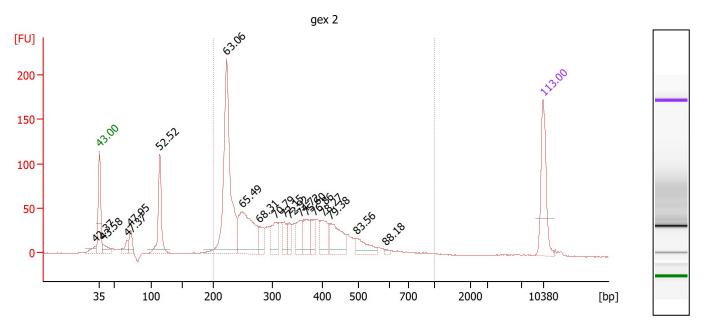
Region table for sample 3: <u>cdna 1</u>

From [bp]	To [bp] C A	orr. rea	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	1,000 2	7.3	18	684	30.8	18.09	51.7	

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Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-10-28_10-55-45.xad Created: 28/10/2021 10:55:44 Modified: 28/10/2021 11:46:18

Electropherogram Summary Continued ...



Overall Results for sample 4: gex 2

Number of peaks found: 18 Corr. Area 1: 1,465.6

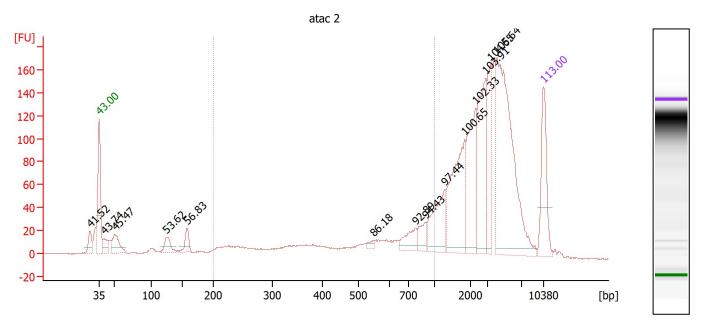
Noise: 0.4

Peak ta	able	for sample 4:	gex 2		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		33	0.00	0.0	
2	◀	35	125.00	5,411.3	Lower Marker
3		39	27.09	1,060.5	
4		67	22.42	504.3	
5		72	34.62	724.8	
6		114	133.32	1,776.0	
7		222	385.00	2,621.7	
8		249	178.72	1,089.1	
9		279	30.87	167.7	
10		307	47.56	235.1	
11		324	28.47	133.3	
12		333	27.13	123.3	
13		356	38.94	165.8	
14		367	47.11	194.5	
15		383	34.86	138.0	
16		401	46.95	177.6	
17		420	69.44	250.3	
18		494	37.33	114.5	
19		587	3.45	8.9	
20		10,380	75.00	10.9	Upper Marker

Region table for sample 4: <u>gex 2</u>

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	1,000	1,465.6	83	335	34.7	1,155.00	5,980.1	

Electropherogram Summary Continued ...



Overall Results for sample 5: <u>atac 2</u>

Number of peaks found: 14 Corr. Area 1: 499.8

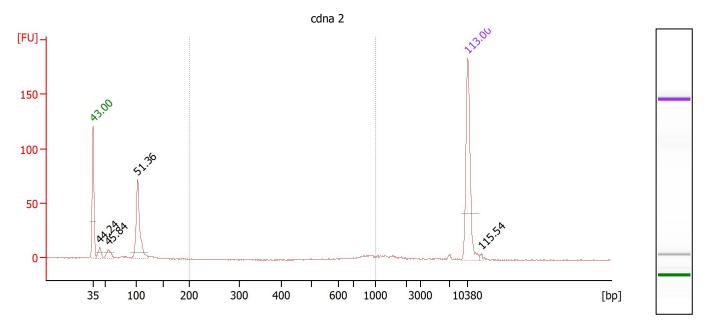
Noise: 0.5

Peak ta	ble	for sample 5:	atac 2		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		30	0.00	0.0	
2	4	35	125.00	5,411.3	Lower Marker
3		40	29.37	1,120.6	
4		51	57.11	1,697.8	
5		125	32.51	394.6	
6		157	28.92	278.4	
7		546	6.35	17.6	
8		786	32.12	61.9	
9		900	24.74	41.7	
10		1,290	78.67	92.4	
11		1,853	141.30	115.6	
12		2,301	113.30	74.6	
13		2,869	123.32	65.1	
14		3,286	80.97	37.3	
15		4,034	297.09	111.6	
16		10,380	75.00	10.9	Upper Marker

Region table for sample 5: <u>atac 2</u>

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	1,000	499.8	22	560	41.4	397.33	1,451.6	

Electropherogram Summary Continued ...



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

Number of peaks found: 4 Corr. Area 1: 12.4

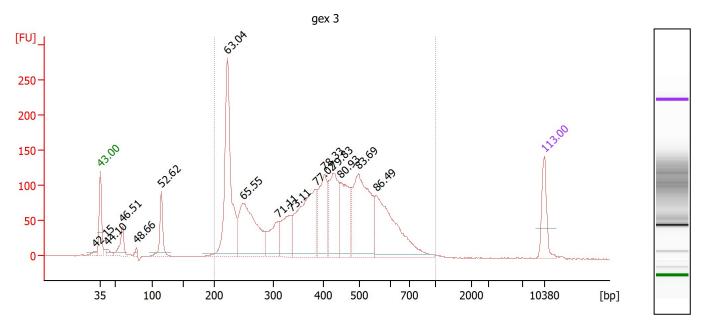
Noise: 0.4

Peak table for sample 6:		<u>cdna 2</u>				
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations		
1	35	125.00	5,411.3	Lower Marker		
2	43	16.60	586.0			
3	54	17.98	503.2			
4	102	118.88	1,764.8			
5	10,380	75.00	10.9	Upper Marker		
6	12,872	0.00	0.0			

Region table for sample 6: cdna 2

From	To [bp] Corr.	% of Total	Average Size	Size distribution in CV [%]		Molarity	Co lor
[bp]	Area	iotai	[nh]	CV [90]	[P9/ pi]	[pmol/l]	101
200	1 000 12 4	6	835	13.0	7 54	13.0	

Electropherogram Summary Continued ...



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

Number of peaks found: 15 Corr. Area 1: 3,256.0

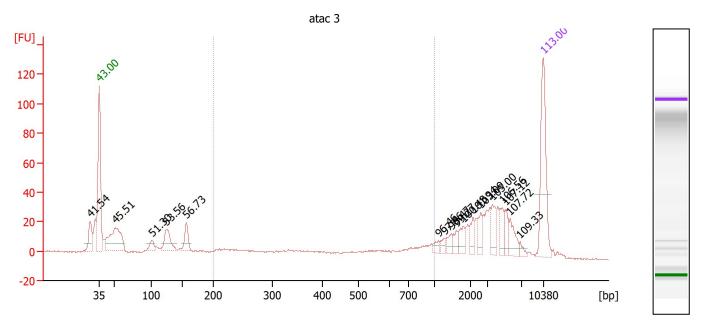
Noise: 0.4

Peak ta	able	for sample 7:	<u>gex 3</u>		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		32	0.00	0.0	
2	4	35	125.00	5,411.3	Lower Marker
3		42	23.57	850.1	
4		60	92.15	2,330.8	
5		78	11.98	231.3	
6		115	122.58	1,618.7	
7		222	556.72	3,794.7	
8		249	398.23	2,420.7	
9		311	130.72	637.6	
10		336	147.87	667.3	
11		385	351.41	1,383.7	
12		402	206.87	780.5	
13		428	233.16	825.1	
14		448	186.48	631.2	
15		496	388.85	1,187.1	
16		553	344.89	945.1	
17		10,380	75.00	10.9	Upper Marker

Region table for sample 7: <u>gex 3</u>

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	1,000	3,256.0	92	408	32.9	2,992.96	13,192.1	

Electropherogram Summary Continued ...



Overall Results for sample 8: <u>atac 3</u>

Number of peaks found: 17 Corr. Area 1: 128.9

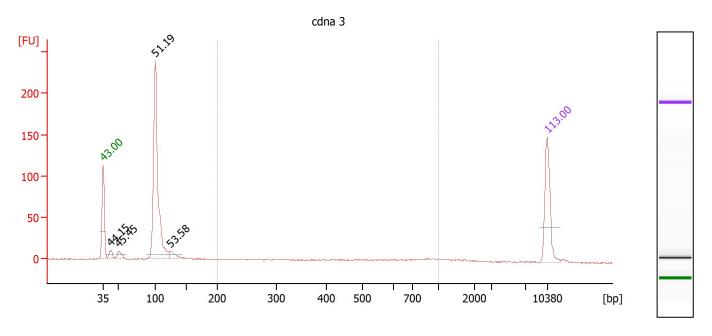
Noise: 0.4

Peak tal	ble	for sample 8:	<u>atac 3</u>		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1		30	0.00	0.0	
2	4	35	125.00	5,411.3	Lower Marker
3		51	104.69	3,089.3	
4		101	16.06	240.0	
5		124	43.03	524.8	
6		156	27.03	262.0	
7		1,118	6.44	8.7	
8		1,294	5.66	6.6	
9		1,470	8.22	8.5	
10		1,611	9.16	8.6	
11		1,814	13.86	11.6	
12		2,160	12.69	8.9	
13		2,576	14.16	8.3	
14		3,553	18.42	7.9	
15		4,734	14.42	4.6	
16		5,152	13.14	3.9	
17		5,609	17.96	4.9	
18		6,828	4.66	1.0	
19		10,380	75.00	10.9	Upper Marker

Region table for sample 8: <u>atac 3</u>

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	1,000	128.9	19	535	44.3	120.69	472.7	

Electropherogram Summary Continued ...



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

Number of peaks found: 4 Corr. Area 1: 60.0

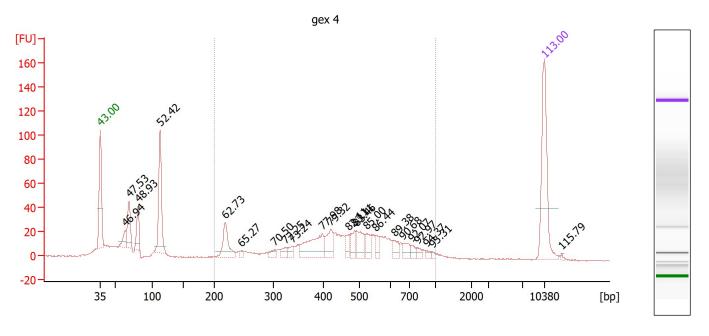
Noise: 0.4

Peak ta	able	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	19.40	694.8	
3		51	21.17	631.9	
4		100	462.93	6,994.4	
5		124	21.30	259.3	
6		10,380	75.00	10.9	Upper Marker

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

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
L-61			r-61	0. [· · ·]	FB 21 E-7	[[, .]	
200	1 000 60 0	11	527	42.2	49 32	189 N	

Electropherogram Summary Continued ...



Overall Results for sample 10: gex 4

Number of peaks found: 23 Corr. Area 1: 546.9

Noise: 0.5

Peak tab	Peak table for sample 10: <u>gex 4</u>									
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations					
1	4 3	35	125.00	5,411.3	Lower Marker					
2	6	64	18.61	443.4						
3	6	69	49.25	1,085.6						
4	8	81	33.37	626.0						
5		113	105.47	1,418.2						
6	2	219	43.54	301.4						
7	2	246	4.55	28.0						
8	3	303	7.87	39.4						
9	3	325	9.22	43.0						
10	3	337	8.53	38.3						
11	3	397	54.83	209.4						
12	4	419	27.70	100.1						
13	4	468	10.98	35.5						
14	4	486	16.63	51.8						
15	4	492	24.85	76.5						
16	į	522	15.61	45.3						
17	į	552	9.64	26.5						
18	6	620	11.41	27.9						
19	(664	11.89	27.1						
20	7	726	6.70	14.0						
21	7	792	5.00	9.6						
22	8	895	3.70	6.3						
23	9	965	2.32	3.6						
24)	10,380	75.00	10.9	Upper Marker					
25		13,119	0.00	0.0						

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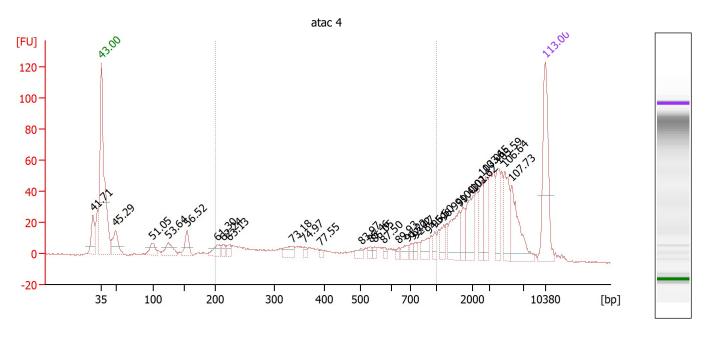
Assay Class: Data Path: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-10-28_10-55-45.xad Created: Modified: 28/10/2021 10:55:44 28/10/2021 11:46:18

Electropherogram Summary Continued ...

... Region table for sample 10: <u>gex 4</u>

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	1,000	546.9	61	472	33.4	405.91	1,558.3	

Electropherogram Summary Continued ...



Overall Results for sample 11: <u>atac 4</u>

Number of peaks found: 30 Corr. Area 1: 302.6

Noise: 0.5

Peak table for sample 11: <u>atac 4</u>									
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations					
1	30	0.00	0.0						
2	35	125.00	5,411.3	Lower Marker					
3	50	46.73	1,428.9						
4	99	21.41	327.3						
5	125	27.91	338.2						
6	154	26.42	259.6						
7	203	18.62	138.6						
8	214	9.07	64.4						
9	223	9.89	67.1						
10	337	17.10	77.0						
11	359	5.39	22.8						
12	391	3.82	14.8						
13	501	8.28	25.0						
14	532	6.29	17.9						
15	544	4.76	13.3						
16	573	4.36	11.5						
17	639	4.28	10.2						
18	683	10.04	22.3						
19	722	5.86	12.3						
20	755	6.29	12.6						
21	913	15.14	25.1						
22	986	8.63	13.3						
23	1,212	14.17	17.7						
24	1,639	39.06	36.1						
25	1,743	15.96	13.9						
26	2,045	35.46	26.3						

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Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DEDAE01844_2021-10-28_10-55-45.xad Created: 28/10/2021 10:55:44 Modified: 28/10/2021 11:46:18

Electropherogram Summary Continued ...

Peak t	Peak table for sample 11: <u>atac 4</u>							
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations				
27	2,545	28.01	16.7					
28	2,777	23.51	12.8					
29	3,998	29.69	11.3					
30	4,789	38.47	12.2					
31	5,617	48.50	13.1					
32	10,380	75.00	10.9	Upper Marker				

Region table for sample 11: <u>atac 4</u>

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	1,000	302.6	29	505	46.0	282.03	1,160.9	

