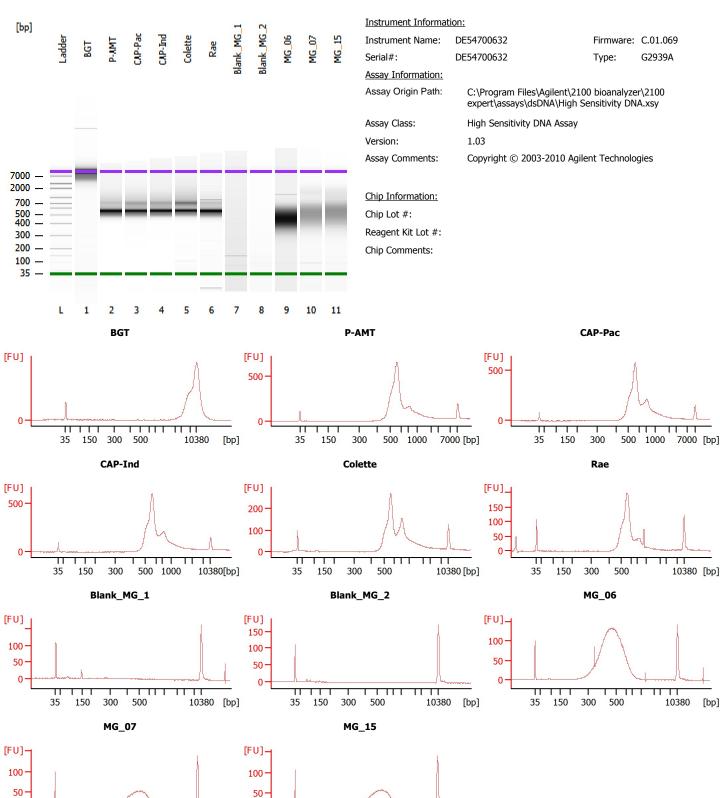
Assay Class: High Sensitivity DNA Assay Created: 1/18/2022 11:37:01 AM Data Path: C:\...gh Sensitivity DNA Assay_DE54700632_2022-01-18_11-37-01.xad C:\...gh Sensitivity DNA Assay_DE54700632_2022-01-18_11-37-01.xad Created: 1/18/2022 11:49:12 PM

Electrophoresis File Run Summary



300

500

10380

[bp]

35 150

35 150

500

10380

[bp]

300

Page 2 of 17

Assay Class: Data Path:	High Sensitivity DNA Assay C:\gh Sensitivity DNA Assay_DE547	00632_2022-01-18_11-37-01.xad	Created: Modified:	1/18/2022 11:37:01 AM 1/18/2022 1:49:12 PM
Electrophore	sis File Run Summary (Chip Sum	mary)		
Sample Name	Sample Rest. Sta Observation Comment Digest tus	Result Color Label		
BGT P-AMT CAP-Pac CAP-Ind Colette Rae Blank_MG_1 Blank_MG_2 MG_06 MG_07 MG_15 Ladder				
Chip Lot #		Reagent Kit Lot #		

Chip Comments:

Printed:

Page 3 of 17

Assay Class: High Sensitivity DNA Assay Created: 1/18/2022 11:37:01 AM Data Path: C:\...gh Sensitivity DNA Assay_DE54700632_2022-01-18_11-37-01.xad C:\...gh Sensitivity DNA Assay_DE54700632_2022-01-18_11-37-01.xad Created: 1/18/2022 11:49:12 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

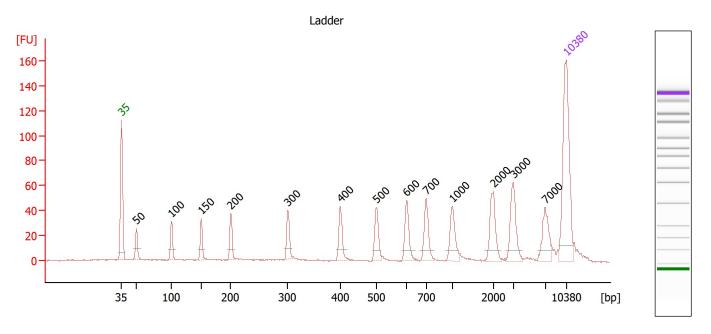
Ladder

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

Printed:

1/18/2022 2:03:21 PM

Electropherogram Summary



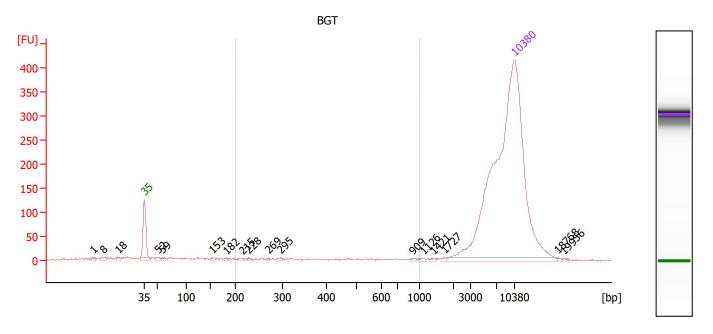
Overall Results for Ladder

Noise: 0.4

Peak	table	for	Ladder
rcan	cavic	101	Lauuci

Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	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

Electropherogram Summary Continued ...



Overall Results for sample 1: BGT

Number of peaks found: 17 Corr. Area 1: 124.7

Noise: 0.5

Peak table for sample 1: <u>BGT</u>							
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations			
1	1	0.00	0.0				
2	8	0.00	0.0				
3	18	0.00	0.0				
4	35	125.00	5,411.3	Lower Marker			
5	52	0.64	18.5				
6	59	0.53	13.7				
7	153	0.37	3.7				
8	182	0.40	3.3				
9	215	0.46	3.2				
10	228	0.32	2.2				
11	269	0.24	1.4				
12	295	0.31	1.6				
13	909	0.21	0.3				
14	1,126	0.24	0.3				
15	1,421	0.20	0.2				
16	1,727	0.20	0.2				
17	10,380	75.00	10.9	Upper Marker			
18	18,768	0.00	0.0				
19	19,956	0.00	0.0				

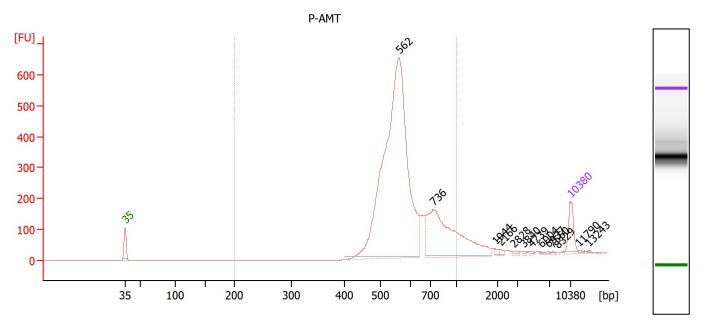
Region table for sample 1: BGT

From [bp]	To [bp] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	1,000	124.7	30	453	45.2	7.09	30.6	

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Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DE54700632_2022-01-18_11-37-01.xad C:\...gh Sensitivity DNA A

Electropherogram Summary Continued ...



Overall Results for sample 2: P-AMT

Number of peaks found: 13 Corr. Area 1: 3,961.3

Noise: 0.4

Peak ta	ble	for sample 2:	P-AMT		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	◀	35	125.00	5,411.3	Lower Marker
2		562	2,174.97	5,862.9	
3		736	625.08	1,286.9	
4		1,944	9.20	7.2	
5		2,166	7.47	5.2	
6		2,828	7.16	3.8	
7		3,840	3.74	1.5	
8		4,739	4.57	1.5	
9		6,004	3.60	0.9	
10		6,937	4.10	0.9	
11		7,560	2.53	0.5	
12		8,329	2.00	0.4	
13	8	10,380	75.00	10.9	Upper Marker
14		11,790	0.00	0.0	
15		13,243	0.00	0.0	

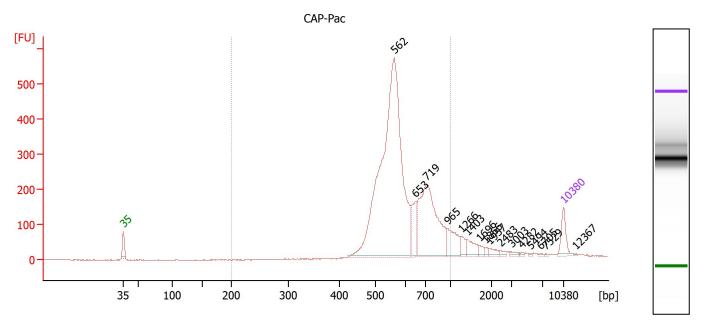
Region table for sample 2: P-AMT

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	3,961.3	91	594	18.1	2,623.45	6,910.1	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...gh Sensitivity DNA Assay_DE54700632_2022-01-18_11-37-01.xad

Created: 1/18/2022 11:37:01 AM Modified: 1/18/2022 1:49:12 PM

Electropherogram Summary Continued ...



Overall Results for sample 3: <u>CAP-Pac</u>

Number of peaks found: 16 Corr. Area 1: 3,579.8

Noise: 1.0

Peak table	for sample 3:	<u>CAP-Pac</u>		
Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	562	2,286.27	6,162.1	
3	653	134.02	311.0	
4	719	703.17	1,482.6	
5	965	134.29	210.8	
6	1,266	39.87	47.7	
7	1,403	63.26	68.3	
8	1,696	21.53	19.2	
9	1,866	11.55	9.4	
10	1,957	23.92	18.5	
11	2,483	11.27	6.9	
12	3,003	12.96	6.5	
13	4,282	5.38	1.9	
14	5,494	3.46	1.0	
15	6,705	3.88	0.9	
16	7,529	2.59	0.5	
17 🍓	10,380	75.00	10.9	Upper Marker
18	12,367	0.00	0.0	

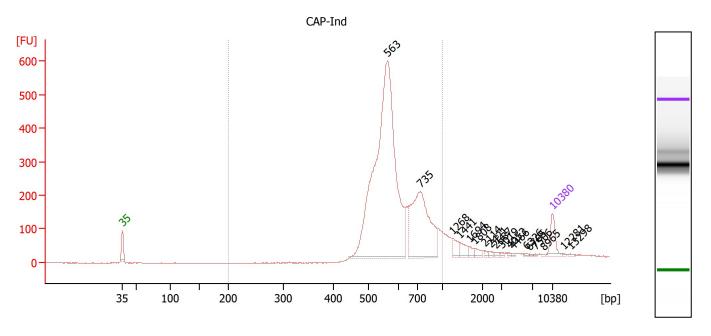
Region table for sample 3: <u>CAP-Pac</u>

From	To [bp] Co	rr. % o	f Average Siz	e Size distribution in	Conc.	Molarity	Co
[bp]	Are	ea Tota	ıl [bp]	CV [%]	[pg/µl]	[pmol/l]	lor
200	1 000 3 5	79 8 89	608	18.8	3 237 80	8 359 5	

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Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DE54700632_2022-01-18_11-37-01.xad C:\...gh Sensitivity DNA A

Electropherogram Summary Continued ...



Overall Results for sample 4: <u>CAP-Ind</u>

Number of peaks found: 18 Corr. Area 1: 3,648.2

Noise: 0.7

for sample 4:	CAP-Ind		
Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
35	125.00	5,411.3	Lower Marker
563	2,356.43	6,337.0	
735	698.02	1,438.6	
1,268	44.84	53.6	
1,441	44.52	46.8	
1,694	20.04	17.9	
1,808	22.66	19.0	
2,114	8.99	6.4	
2,421	8.98	5.6	
2,687	9.29	5.2	
3,019	6.38	3.2	
4,053	4.40	1.6	
4,466	5.55	1.9	
6,326	5.22	1.2	
6,705	3.35	0.8	
7,506	2.91	0.6	
8,965	2.38	0.4	
10,380	75.00	10.9	Upper Marker
12,281	0.00	0.0	
13,298	0.00	0.0	
	35 563 735 1,268 1,441 1,694 1,808 2,114 2,421 2,687 3,019 4,053 4,466 6,326 6,705 7,506 8,965 10,380 12,281	Size [bp] Conc. [pg/μl] 35 125.00 563 2,356.43 735 698.02 1,268 44.84 1,441 44.52 1,694 20.04 1,808 22.66 2,114 8.99 2,421 8.98 2,687 9.29 3,019 6.38 4,053 4.40 4,466 5.55 6,326 5.22 6,705 3.35 7,506 2.91 8,965 2.38 10,380 75.00 12,281 0.00	Size [bp] Conc. [pg/μl] Molarity [pmol/l] 35 125.00 5,411.3 563 2,356.43 6,337.0 735 698.02 1,438.6 1,268 44.84 53.6 1,441 44.52 46.8 1,694 20.04 17.9 1,808 22.66 19.0 2,114 8.99 6.4 2,421 8.98 5.6 2,687 9.29 5.2 3,019 6.38 3.2 4,053 4.40 1.6 4,466 5.55 1.9 6,326 5.22 1.2 6,705 3.35 0.8 7,506 2.91 0.6 8,965 2.38 0.4 10,380 75.00 10.9 12,281 0.00 0.0

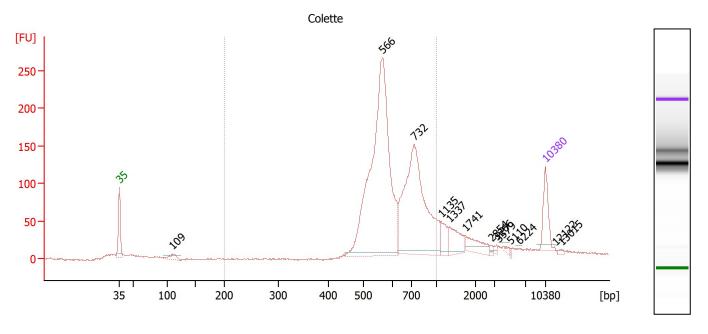
Region table for sample 4: <u>CAP-Ind</u>

From [bp]	To [bp	o] Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]		Molarity [pmol/l]	Co lor
200	1,000	3,648.2	88	611	18.6	3,243.70		

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Assay Class: High Sensitivity DNA Assay C:\...gh Sensitivity DNA Assay_DE54700632_2022-01-18_11-37-01.xad C:\...gh Sensitivity DNA A

Electropherogram Summary Continued ...



Overall Results for sample 5 : <u>Colette</u>

Number of peaks found: 13 Corr. Area 1: 1,734.4

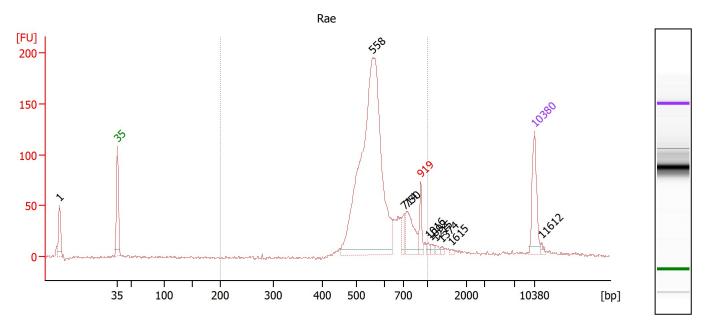
Noise: 0.9

Peak ta	able	for sample 5:	<u>Colette</u>		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2		109	29.23	405.4	
3		566	1,122.63	3,005.0	
4		732	693.64	1,435.4	
5		1,135	51.09	68.2	
6		1,337	77.34	87.7	
7		1,741	54.07	47.1	
8		3,266	11.17	5.2	
9		2,854	6.20	3.3	
10		3,579	6.21	2.6	
11		5,110	1.43	0.4	
12		6,224	0.60	0.1	
13	8	10,380	75.00	10.9	Upper Marker
14		12,122	0.00	0.0	
15		13,015	0.00	0.0	

Region table for sample 5: Colette

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	1,734.4	81	633	19.2	1,797.56	4,471.7	

Electropherogram Summary Continued ...



Overall Results for sample 6: Rae

Number of peaks found: 10 Corr. Area 1: 1,180.4

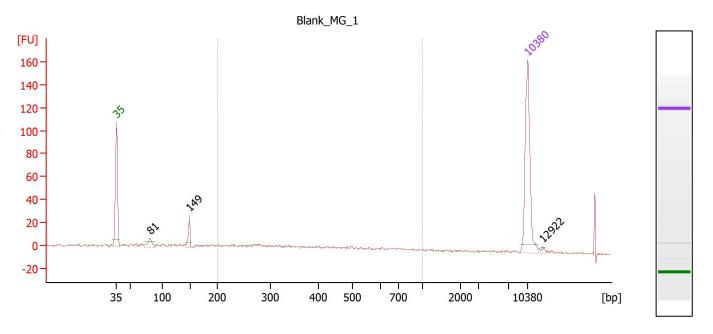
Noise: 1.1

Peak t	Peak table for sample 6: Rae										
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations						
1		1	0.00	0.0							
2	4	35	125.00	5,411.3	Lower Marker						
3		558	954.28	2,592.9							
4		714	26.36	55.9							
5		750	78.78	159.1							
6		919	33.10	54.6	excluded peak						
7		1,016	6.65	9.9							
8		1,092	7.56	10.5							
9		1,215	6.17	7.7							
10		1,374	5.21	5.7							
11		1,615	3.08	2.9							
12	Pa	10,380	75.00	10.9	Upper Marker						
13		11.612	0.00	0.0							

Region table for sample 6: Rae

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	1,180.4	84	592	18.4	1,231.21	3,260.2	

Electropherogram Summary Continued ...



Overall Results for sample 7: Blank MG 1

Number of peaks found: 3 Corr. Area 1: 94.2

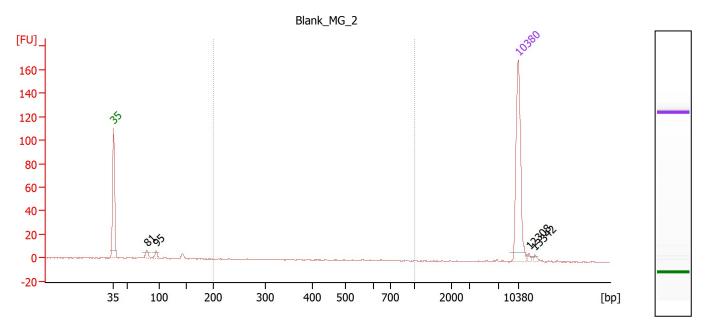
Noise: 0.6

Peak t	table	for sample 7:	Blank MG 1				
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations		
1	4	35	125.00	5,411.3	Lower Marker		
2		81	16.20	304.6			
3		149	19.35	197.2			
4		10,380	75.00	10.9	Upper Marker		
5		12,922	0.00	0.0			

Region table for sample 7: Blank MG 1

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 94.2	46	421	42.0	74.45	332.2	

Electropherogram Summary Continued ...



Overall Results for sample 8: Blank MG 2

Number of peaks found: 4 Corr. Area 1: 9.6

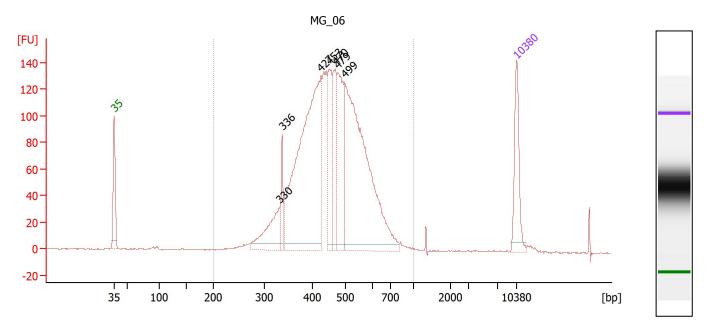
Noise: 0.4

Peak ta	able	for sample 8:	Blank MG 2				
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/I]	Observations		
1	4	35	125.00	5,411.3	Lower Marker		
2		81	9.07	169.6			
3		95	7.16	113.8			
4		10,380	75.00	10.9	Upper Marker		
5		12,308	0.00	0.0			
6		13,342	0.00	0.0			

Region table for sample 8: Blank MG 2

From [bp]	To [bp] Corr. Area		Average Size [bp]	Size distribution in CV [%]	Conc. [pg/µl]	Molarity [pmol/l]	Co lor
200	1 000 9 6	14	456	38.0	7 33	30.3	

Electropherogram Summary Continued ...



Overall Results for sample 9: MG 06

Number of peaks found: 7 Corr. Area 1: 2,303.7

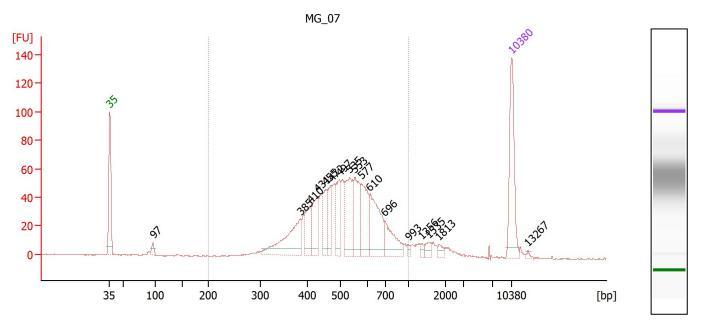
Noise: 0.4

Peak '	table	for sample 9	: <u>MG_06</u>		
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	4	35	125.00	5,411.3	Lower Marker
2		330	114.15	523.3	
3		336	42.41	191.2	
4		427	641.09	2,277.3	
5		453	129.75	434.4	
6		470	100.91	325.6	
7		479	181.08	573.4	
8		499	610.49	1,854.3	
9		10,380	75.00	10.9	Upper Marker

Region table for sample 9: MG 06

From	To [bp] Corr.		% of Average Size	Average Size	Size distribution in	Conc.	Molarity	Co
[bp]		Area	Total	[bp]	CV [%]	[pg/µl]	[pmol/l]	lor
200	1,000	2,303.7	98	463	20.4	1,991.25	6,889.4	

Electropherogram Summary Continued ...



Overall Results for sample 10 : MG 07

Number of peaks found: 17 Corr. Area 1: 974.8

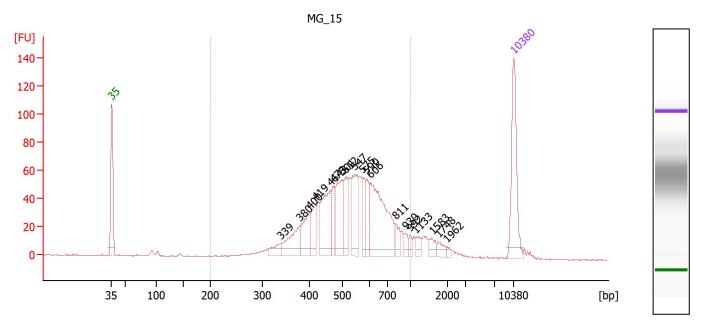
Noise: 0.4

Peak t	Peak table for sample 10 : MG 07									
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations					
1	4	35	125.00	5,411.3	Lower Marker					
2		97	11.64	181.8						
3		385	96.08	378.5						
4		410	43.65	161.2						
5		431	55.80	196.0						
6		455	46.04	153.3						
7		472	35.96	115.4						
8		497	53.40	163.0						
9		535	79.52	225.4						
10		553	64.37	176.3						
11		577	69.03	181.2						
12		610	83.45	207.3						
13		696	47.88	104.2						
14		993	4.11	6.3						
15		1,356	3.90	4.4						
16		1,535	8.77	8.7						
17		1,813	5.96	5.0						
18		10,380	75.00	10.9	Upper Marker					
19		13,267	0.00	0.0						

Region table for sample 10 : MG 07

From [bp]	То [bp] Corr. Area	% of Total	Average Size [bp]	CV [%]		Molarity [pmol/l]	Co lor
200	1,000	974.8	91	522	23.8	826.97	2,579.1	

Electropherogram Summary Continued ...



Overall Results for sample 11: MG 15

Number of peaks found: 19 Corr. Area 1: 1,066.5

Noise: 0.3

Peak table for sample 11: MG 15							
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations		
1	4	35	125.00	5,411.3	Lower Marker		
2		339	17.21	76.9			
3		380	48.90	195.1			
4		400	49.68	188.4			
5		419	32.31	116.9			
6		464	85.46	278.8			
7		473	31.53	101.1			
8		501	75.76	229.0			
9		512	41.80	123.7			
10		547	64.91	179.8			
11		575	29.20	76.9			
12		590	31.89	81.9			
13		606	142.81	356.8			
14		811	15.33	28.7			
15		930	9.14	14.9			
16		992	7.12	10.9			
17		1,133	11.01	14.7			
18		1,583	11.57	11.1			
19		1,748	8.72	7.6			
20		1,962	3.40	2.6			
21		10,380	75.00	10.9	Upper Marker		

Region table for sample 11: MG 15

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	1,066.5	89	539	25.5	839.50	2.576.2	

1/18/2022 11:37:01 AM 1/18/2022 1:49:12 PM Assay Class: High Sensitivity DNA Assay Created: C:\...gh Sensitivity DNA Assay_DE54700632_2022-01-18_11-37-01.xad Data Path: Modified: **Gel Image** [bp] Blank MG 2 Blank MG_1 MG_07 MG_15 Ladder Colette 8 BGT 10380 7000 3000 -2000 -1000 — 700 --600 — 500 — 400 ---300 — 200 ---150 ---100 --2 3 5 6 8 10 11

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		1/18/2022 12:18:27 PM	(GMT08:00) Pacific Standard Time	Rohan	HEIDELBERG-G EL
Run started on port 1 (File: C:\Program Files\Agilent\21 00 bioanalyzer\210 0 expert\Data\202 2-01-18\2100 expert_High Sensitivity DNA Assay_DE54700 632_2022-01-18_11-37-01.xad)	2	Instrument	Run		1/18/2022 11:37:06 AM	(GMT08:00) Pacific Standard Time	Rohan	HEIDELBERG-G EL
Product Number : G2939A		Instrument	Run		1/18/2022 11:37:06 AM	(GMT08:00) Pacific Standard Time	Rohan	HEIDELBERG-G EL
Name :		Instrument	Run		1/18/2022 11:37:06 AM	(GMT08:00) Pacific Standard Time	Rohan	HEIDELBERG-G EL
Vendor : Agilent Technologies	:	Instrument	Run		1/18/2022 11:37:06 AM	(GMT08:00) Pacific Standard Time	Rohan	HEIDELBERG-G EL
Serial# : DE54700632		Instrument	Run		1/18/2022 11:37:06 AM	(GMT08:00) Pacific Standard Time		HEIDELBERG-G EL
Firmware : C.01.069		Instrument	Run		1/18/2022 11:37:06 AM	(GMT08:00) Pacific Standard Time		HEIDELBERG-G EL
Cartridge : Electrode		Instrument	Run		1/18/2022 11:37:06 AM	(GMT08:00) Pacific Standard Time	Rohan	HEIDELBERG-G EL