

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...\Bioanalyzer1\_High Sensitivity DNA Assay\_2023-03-03\_001.xad

Created: 3/3/2023 11:02:31 AM  
Modified: 3/3/2023 11:45:30 AM

**Electrophoresis File Run Summary**

Instrument Information:

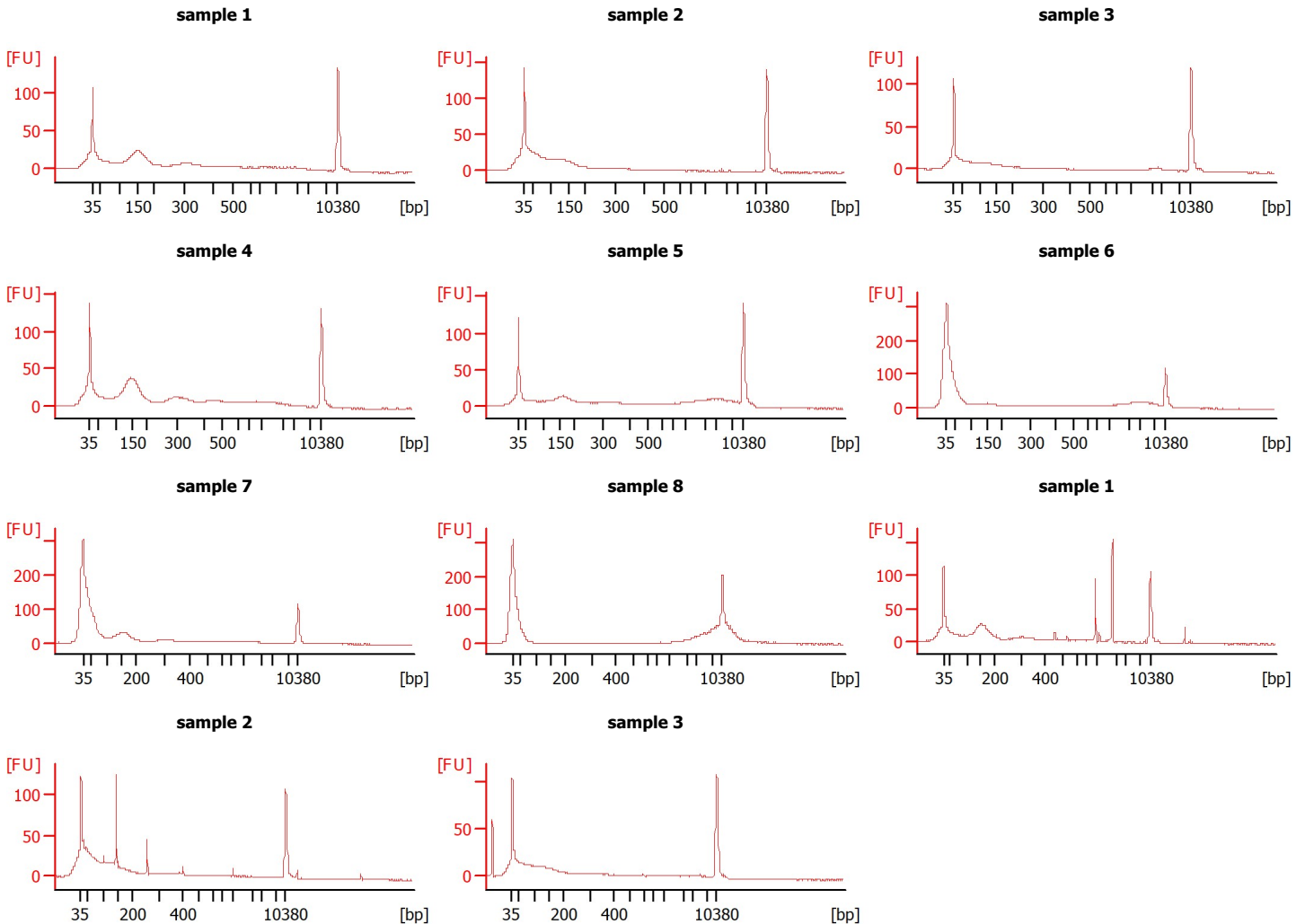
Instrument Name: DE34903152      Firmware: C.01.069  
Serial#: DE34903152      Type: G2938C

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 Observations	Result Label	Result Color
sample 1		<input type="checkbox"/>	✓		
sample 2		<input type="checkbox"/>	✓		
sample 3		<input type="checkbox"/>	✓		
sample 4		<input type="checkbox"/>	✓		
sample 5		<input type="checkbox"/>	✓		
sample 6		<input type="checkbox"/>	✓		
sample 7		<input type="checkbox"/>	✓		
sample 8		<input type="checkbox"/>	✓		
sample 1		<input type="checkbox"/>	✓		
sample 2		<input type="checkbox"/>	✓		
sample 3		<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

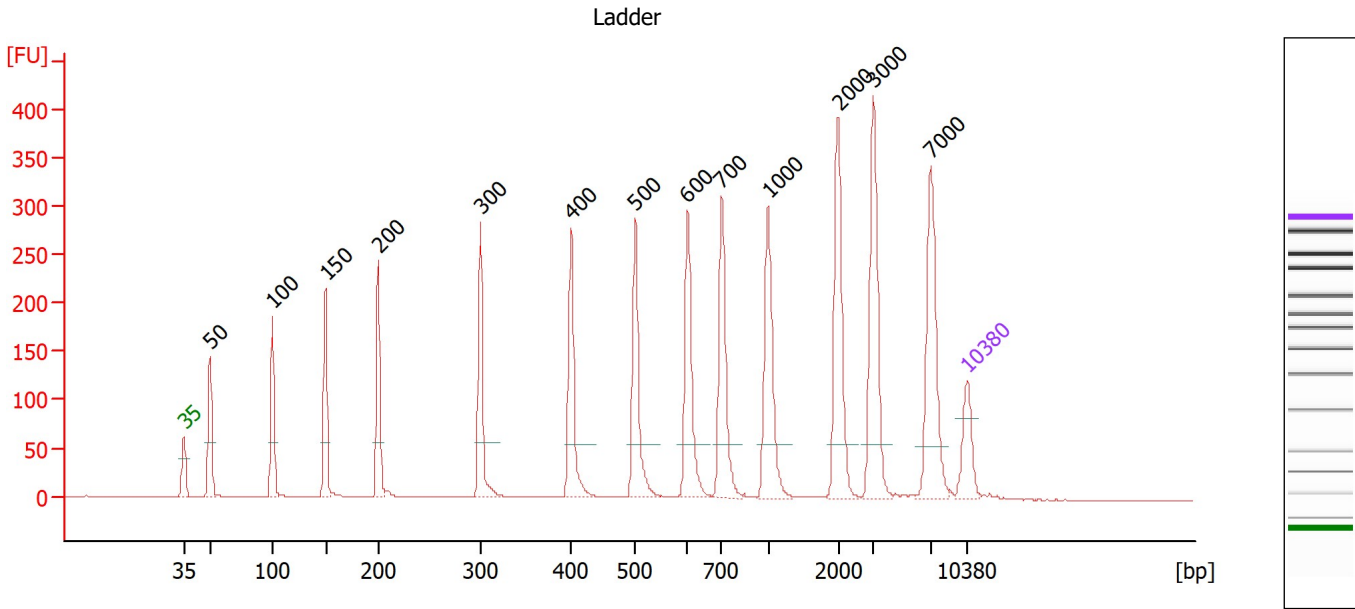
**Ladder**

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

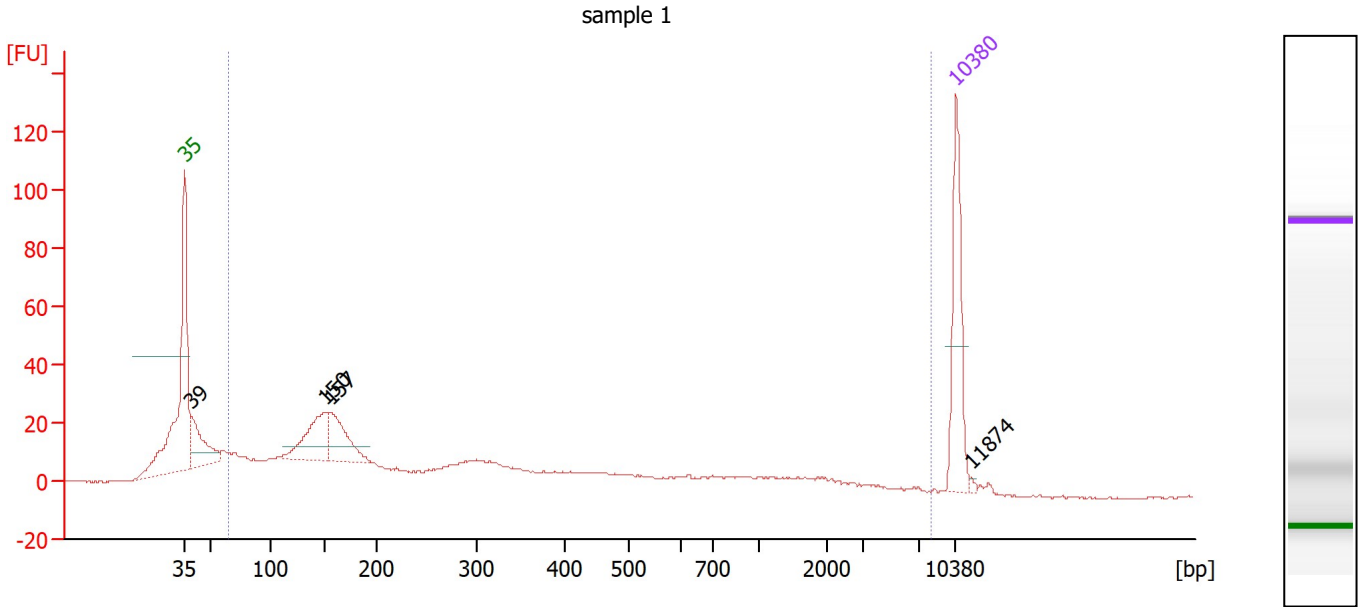
**Peak table for Ladder**

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 1 : sample 1**

Number of peaks found: 4                      Corr. Area 1: 655.3  
 Noise: 0.3

**Peak table for sample 1 : sample 1**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	39	81.28	3,154.0	
3	150	89.42	902.2	
4	157	73.34	706.2	
5	10,380	75.00	10.9	Upper Marker
6	11,874	0.00	0.0	

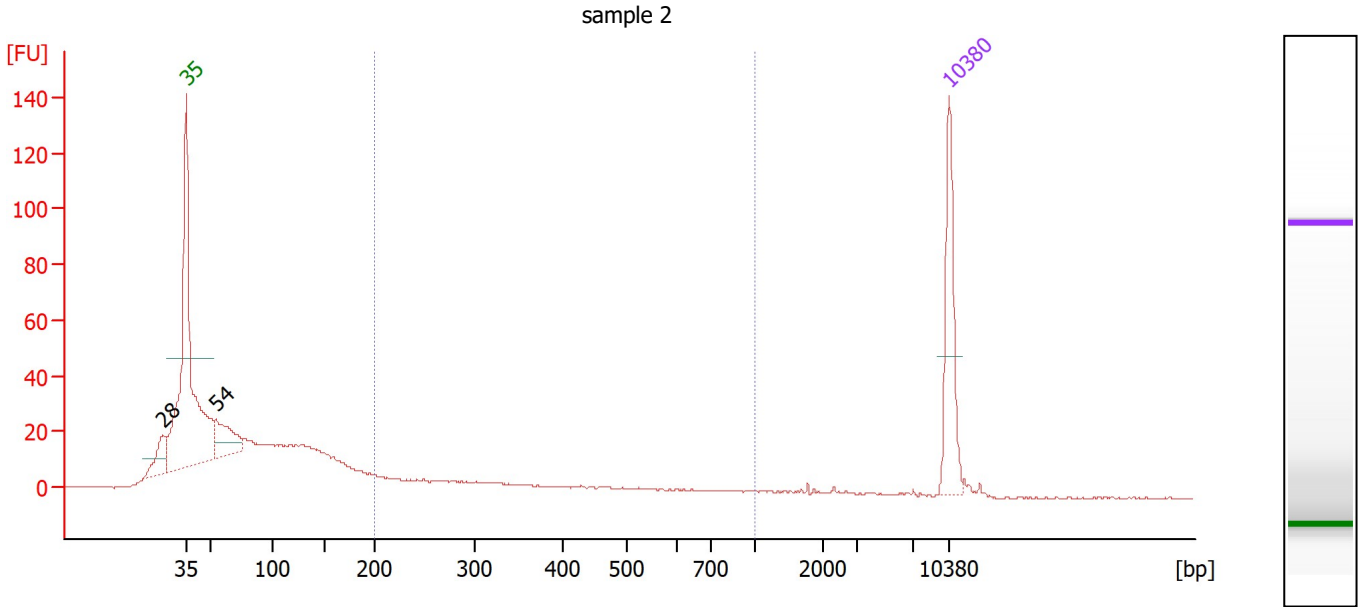
**Region table for sample 1 : sample 1**

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
66	8,077	655.3	84	531	100.0	772.79	6,873.1	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : sample 2**

Number of peaks found: 2                      Corr. Area 1: 110.9  
 Noise: 0.3

**Peak table for sample 2 : sample 2**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	28	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	54	71.42	2,001.6	
4	10,380	75.00	10.9	Upper Marker

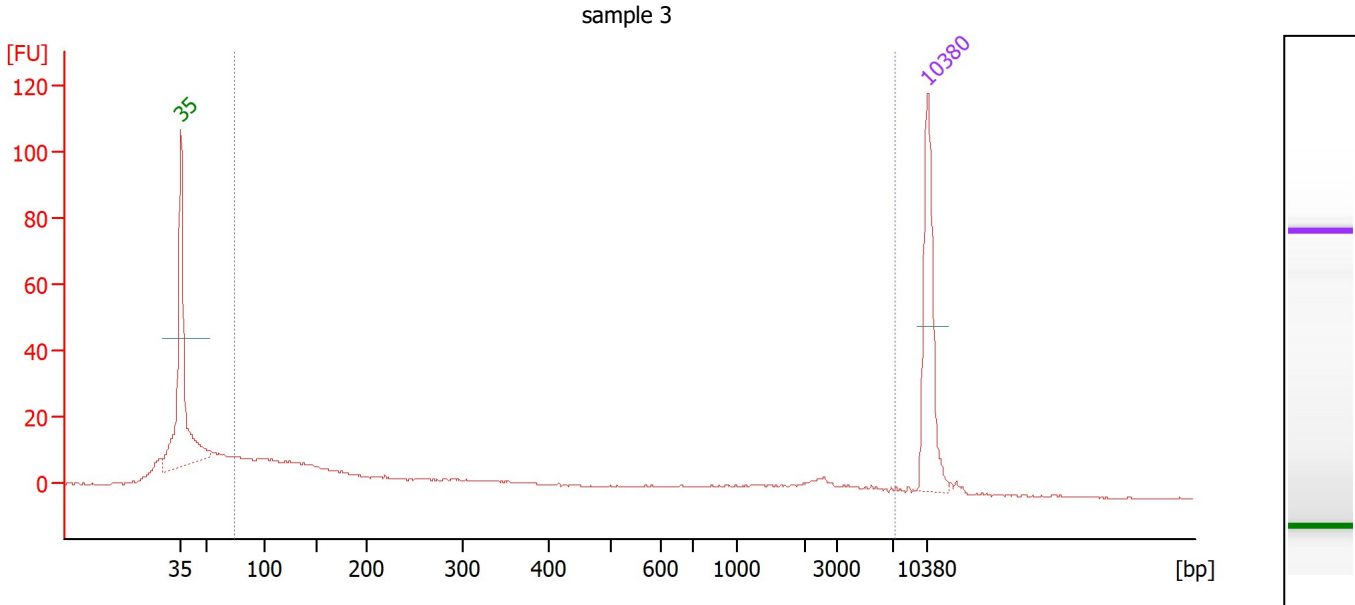
**Region table for sample 2 : sample 2**

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	110.9	18	375	44.4	115.87	572.6	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : sample 3**

Number of peaks found: 0                      Corr. Area 1: 287.1  
 Noise: 0.2

**Peak table for sample 3 : sample 3**

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

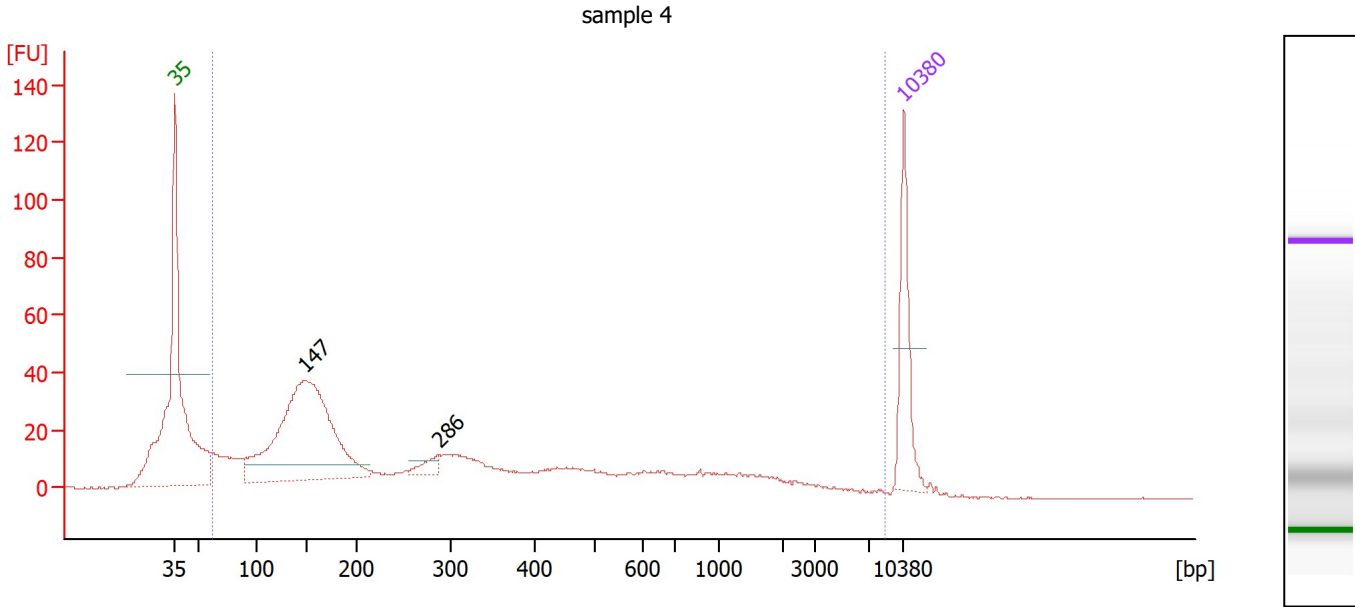
**Region table for sample 3 : sample 3**

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
75	7,264	287.1	78	766	100.0	337.56	3,233.5	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : sample 4**

Number of peaks found: 2                      Corr. Area 1: 934.4  
 Noise: 0.2

**Peak table for sample 4 : sample 4**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	147	502.68	5,177.0	
3	286	19.40	102.6	
4	10,380	75.00	10.9	Upper Marker

**Region table for sample 4 : sample 4**

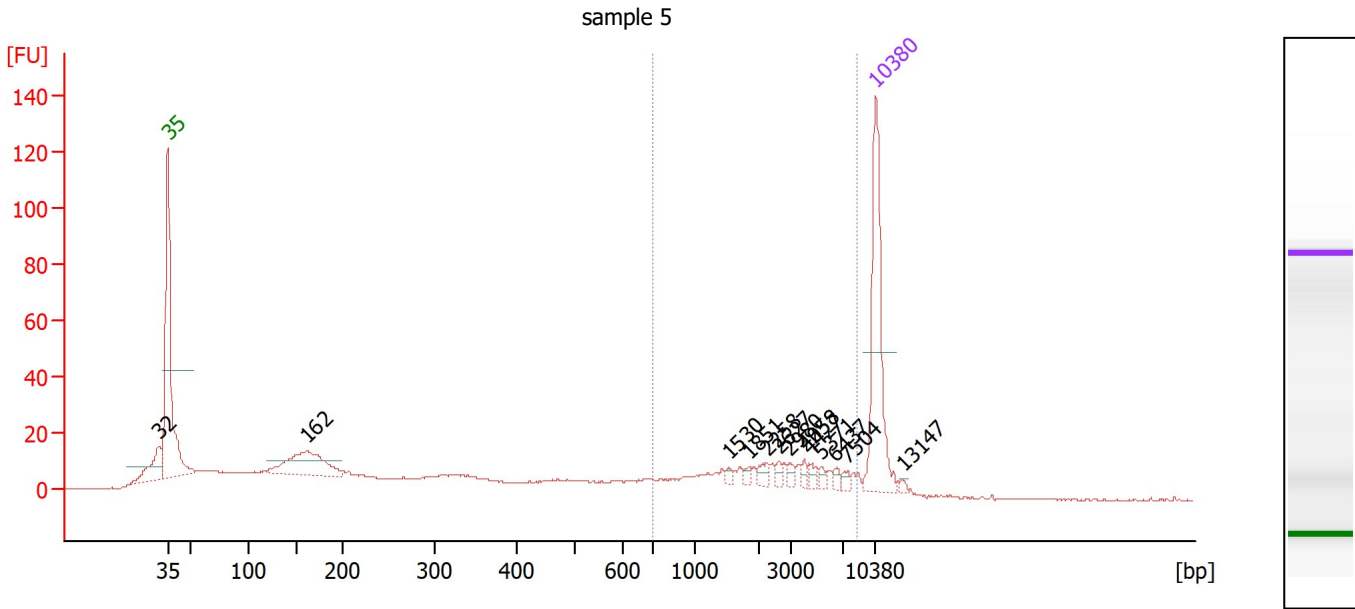
From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
61	8,524	934.4	99	508	100.0	1,090.58	9,847.2	Blue



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**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : sample 5**

Number of peaks found: 13                      Corr. Area 1: 172.5  
 Noise: 0.2

**Peak table for sample 5 : sample 5**

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	162	71.29	668.1	
4	1,530	2.18	2.2	
5	1,851	2.83	2.3	
6	2,258	5.51	3.7	
7	2,627	3.29	1.9	
8	2,980	3.37	1.7	
9	4,058	3.36	1.3	
10	4,427	4.22	1.4	
11	5,371	3.12	0.9	
12	6,437	3.23	0.8	
13	7,504	3.19	0.6	
14	10,380	75.00	10.9	Upper Marker
15	13,147	0.00	0.0	

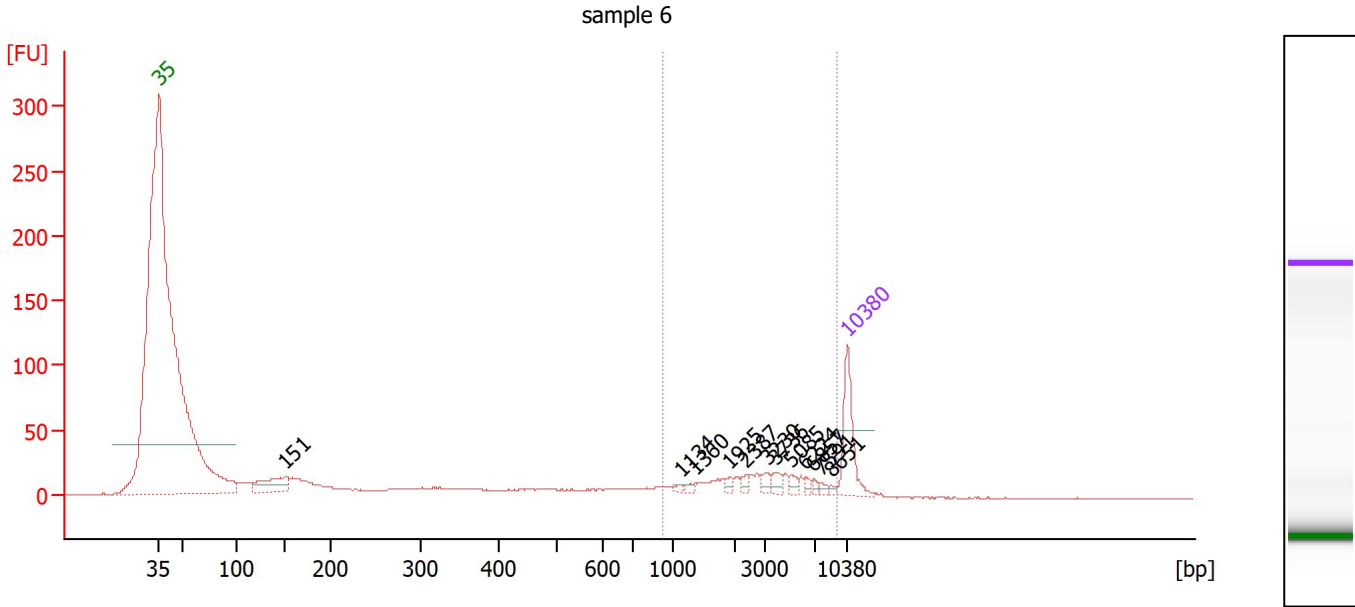
**Region table for sample 5 : sample 5**

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
698	8,578	172.5	25	3,058	68.5	108.07	88.3	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : sample 6**

Number of peaks found: 12                      Corr. Area 1: 229.8  
 Noise: 0.2

**Peak table for sample 6 : sample 6**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	151	80.07	801.5	
3	1,134	4.44	5.9	
4	1,360	5.48	6.1	
5	1,925	6.57	5.2	
6	2,387	6.91	4.4	
7	3,230	11.28	5.3	
8	3,736	12.16	4.9	
9	5,085	8.58	2.6	
10	6,224	6.24	1.5	
11	6,857	4.96	1.1	
12	7,591	5.23	1.0	
13	8,651	3.28	0.6	
14	10,380	75.00	10.9	Upper Marker

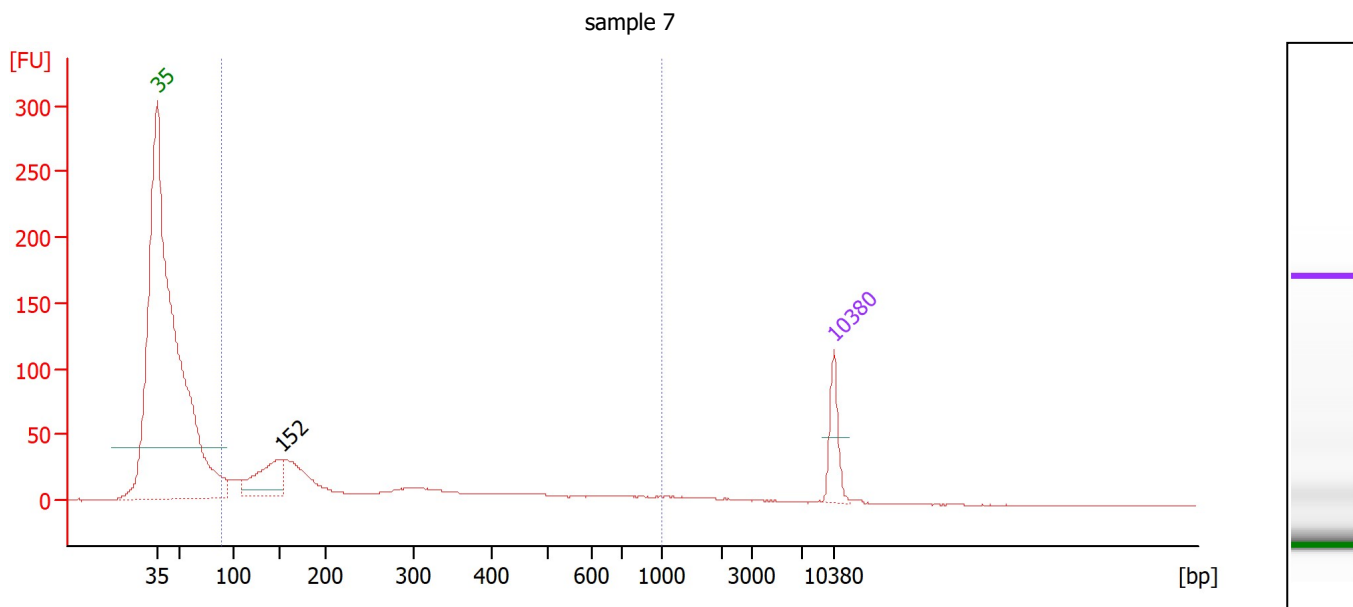
**Region table for sample 6 : sample 6**

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
932	9,242	229.8	34	3,520	61.4	169.25	106.2	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : sample 7**

Number of peaks found: 1                      Corr. Area 1: 710.8  
 Noise: 0.2

**Peak table for sample 7 : sample 7**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	152	219.76	2,184.1	
3	10,380	75.00	10.9	Upper Marker

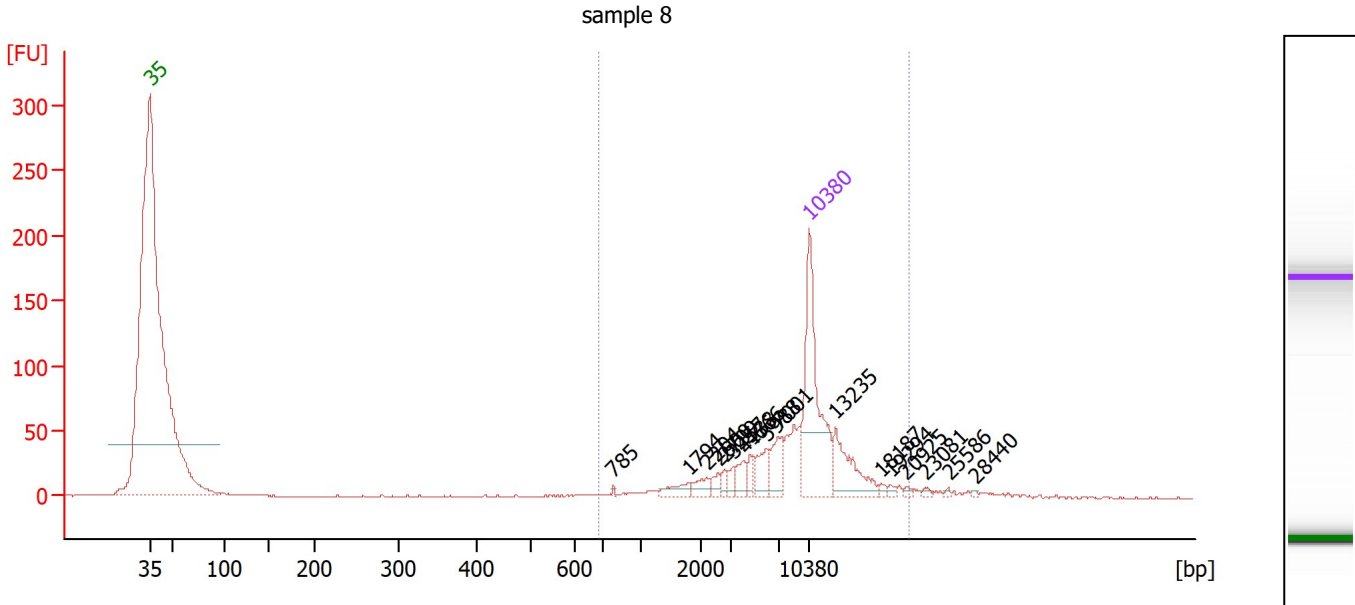
**Region table for sample 7 : sample 7**

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
88	1,000	710.8	91	285	67.1	922.44	7,701.6	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...\Bioanalyzer1\_High Sensitivity DNA Assay\_2023-03-03\_001.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : sample 8**

Number of peaks found: 17                      Corr. Area 1: 500.7  
 Noise: 0.2

**Peak table for sample 8 : sample 8**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	785	0.98	1.9	
3	1,794	6.92	5.8	
4	2,204	6.59	4.5	
5	2,600	4.81	2.8	
6	2,799	3.82	2.1	
7	3,257	4.57	2.1	
8	4,138	8.40	3.1	
9	4,666	5.88	1.9	
10	5,988	14.41	3.6	
11	7,001	16.33	3.5	
12	10,380	75.00	10.9	Upper Marker
13	13,235	0.00	0.0	
14	18,187	0.00	0.0	
15	19,294	0.00	0.0	
16	20,925	0.00	0.0	
17	23,081	0.00	0.0	
18	25,586	0.00	0.0	
19	28,440	0.00	0.0	

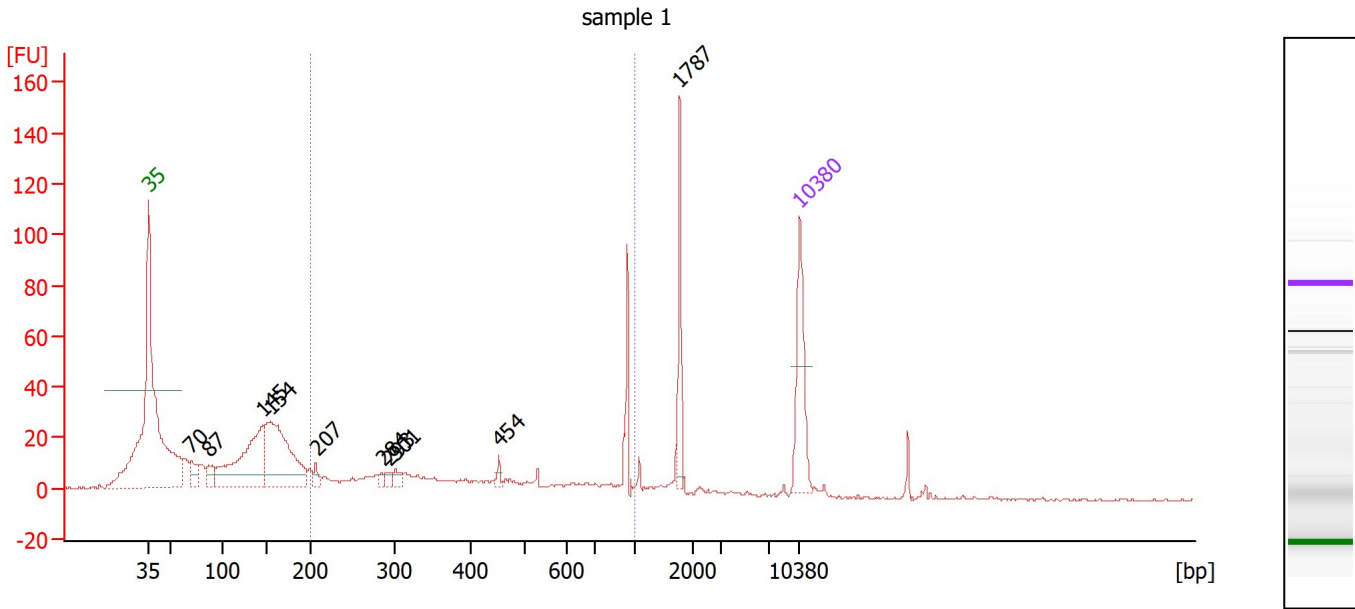
**Region table for sample 8 : sample 8**

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
686	21,380	500.7	84	9,143	49.3	141.69	52.9	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : sample 1**

Number of peaks found: 10                      Corr. Area 1: 250.0  
 Noise: 0.5

**Peak table for sample 9 : sample 1**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	70	32.98	711.6	
3	87	28.14	488.9	
4	145	212.57	2,217.9	
5	154	227.49	2,233.9	
6	207	11.46	83.9	
7	284	7.58	40.4	
8	293	8.14	42.1	
9	301	12.11	61.0	
10	454	6.40	21.4	
11	1,787	47.82	40.5	
12	10,380	75.00	10.9	Upper Marker

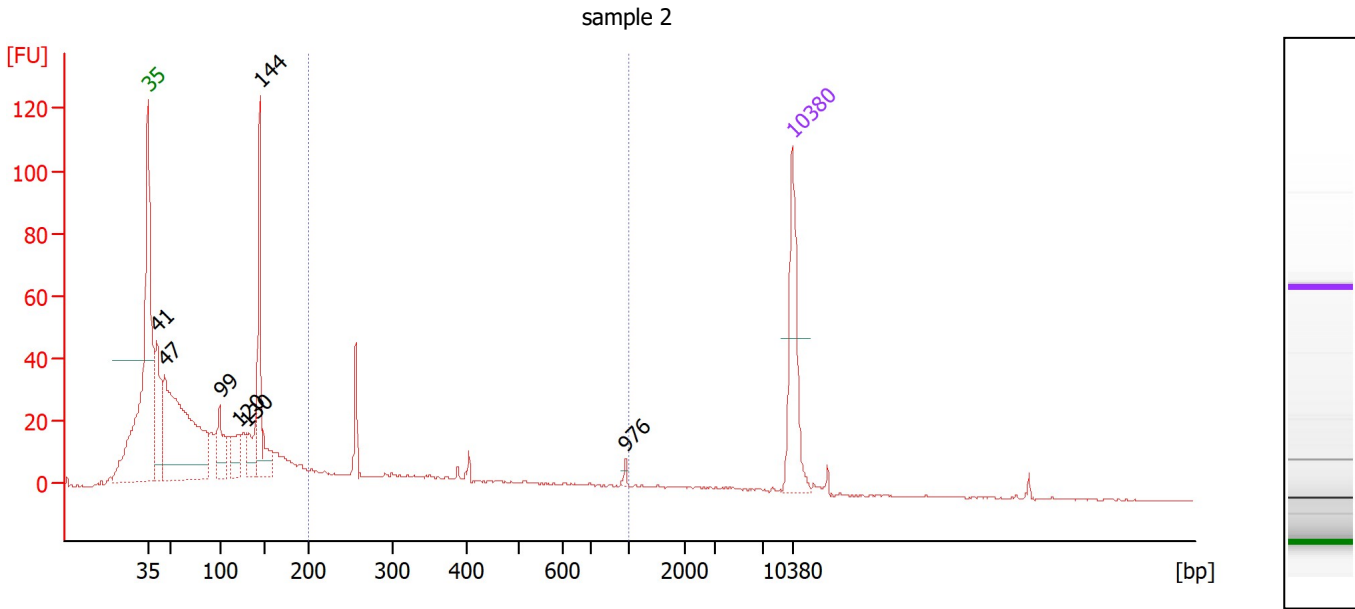
**Region table for sample 9 : sample 1**

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	250.0	35	495	50.5	325.07	1,371.0	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : sample 2**

Number of peaks found: 7                      Corr. Area 1: 191.7  
 Noise: 0.2

**Peak table for sample 10 : sample 2**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	41	115.55	4,274.2	
3	47	394.88	12,853.6	
4	99	56.70	870.0	
5	120	38.14	479.8	
6	130	37.81	440.1	
7	144	138.95	1,460.3	
8	976	2.68	4.2	
9	10,380	75.00	10.9	Upper Marker

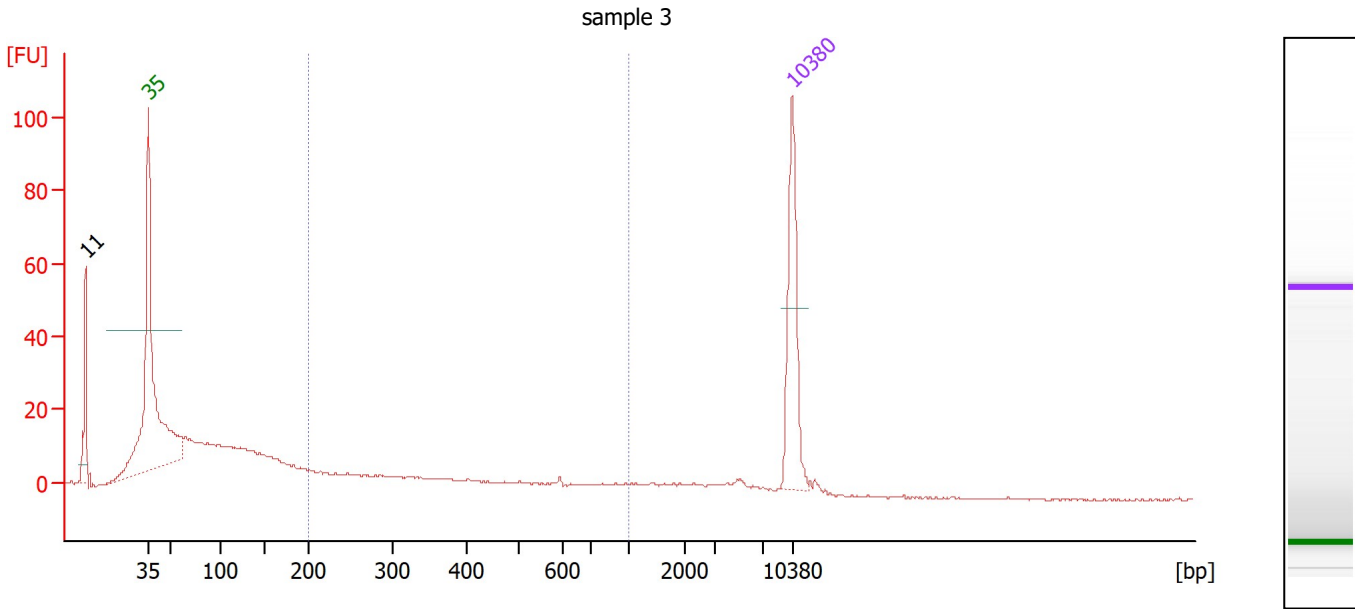
**Region table for sample 10 : sample 2**

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	191.7	22	403	47.0	234.46	1,109.5	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : sample 3**

Number of peaks found: 1                      Corr. Area 1: 121.4  
 Noise: 0.3

**Peak table for sample 11 : sample 3**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	11	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	10,380	75.00	10.9	Upper Marker

**Region table for sample 11 : sample 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	121.4	27	407	45.7	136.89	643.0	Blue

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**Gel Image**



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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		3/3/2023 11:43:50 AM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Run started on port 1 (File: C:\Users\sbsuser\Desktop\2023-03-03\Bioanalyzer1_High Sensitivity DNA Assay_2023-03-03_001.xad)		Instrument	Run		3/3/2023 11:02:37 AM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Product Number : G2938C		Instrument	Run		3/3/2023 11:02:37 AM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Name :		Instrument	Run		3/3/2023 11:02:37 AM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Vendor : Agilent Technologies		Instrument	Run		3/3/2023 11:02:37 AM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Serial# : DE34903152		Instrument	Run		3/3/2023 11:02:37 AM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Firmware : C.01.069		Instrument	Run		3/3/2023 11:02:37 AM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Cartridge : Electrode		Instrument	Run		3/3/2023 11:02:37 AM	(GMT --08:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB