

Assay Class: High Sensitivity DNA Assay  
Data Path: Z:\...\Bioanalyzer1\_High Sensitivity DNA Assay\_2020-07-02\_001.xad

Created: 7/2/2020 2:43:10 PM  
Modified: 7/2/2020 3:24:32 PM

**Electrophoresis File Run Summary**

Instrument Information:

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

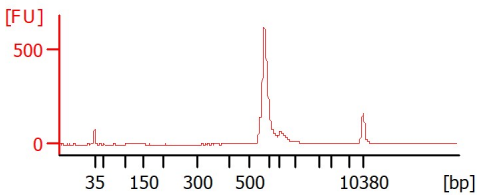
Assay Information:

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

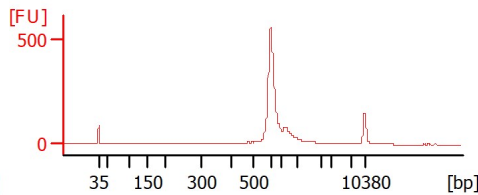
Chip Information:

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

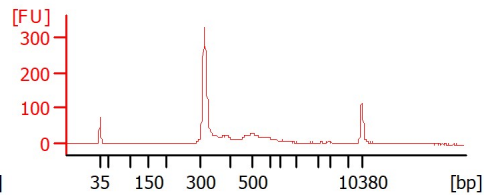
**LongDNA (1:100)**



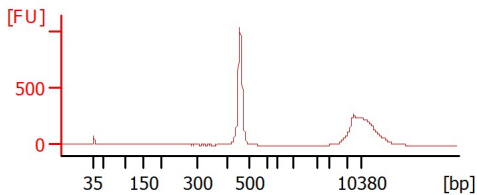
**GolayDNA(1:100)**



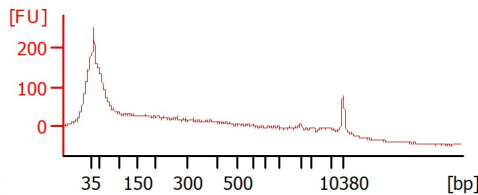
**HYJ423(1:60)**



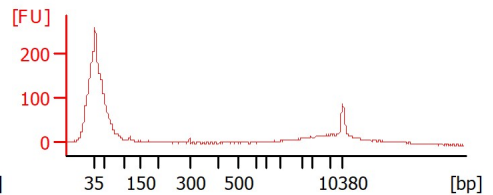
**F165\_redo**



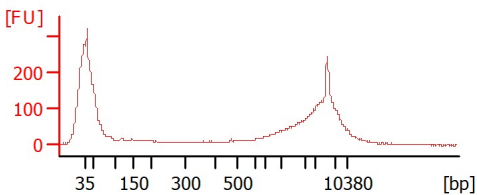
**NovaSeq223-12 (1:15)**



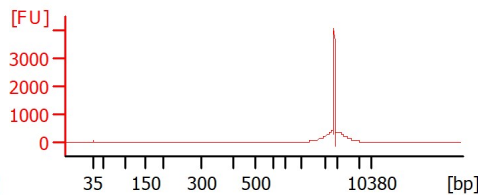
**NovaSeq223-53 (1:20)**



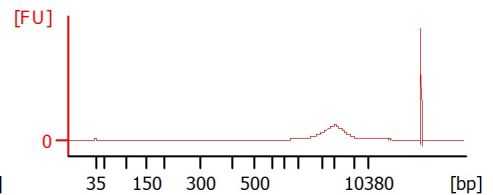
**NovaSeq223-83 (1:30)**



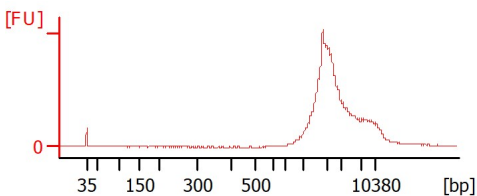
**NovaSeq208-10 (1:2)**



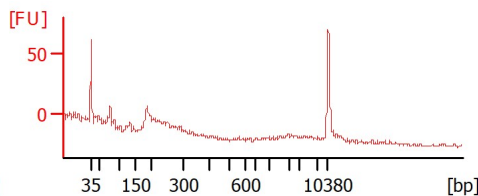
**NovaSeq208-11 (1:3)**



**NovaSeq208-12 (1:20)**



**H2109P\_Wellendorf**



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**Electrophoresis File Run Summary (Chip Summary)**

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
LongDNA (1:100)		<input type="checkbox"/>	✓			
GolayDNA(1:100)		<input type="checkbox"/>	✓			
HYJ423(1:60)		<input type="checkbox"/>	✓			
FI65_redo		<input type="checkbox"/>	✓			
NovaSeq223-12 (1:15)		<input type="checkbox"/>	✓			
NovaSeq223-53 (1:20)		<input type="checkbox"/>	✓			
NovaSeq223-83 (1:30)		<input type="checkbox"/>	✓			
NovaSeq208-10 (1:2)		<input type="checkbox"/>	✓			
NovaSeq208-11 (1:3)		<input type="checkbox"/>	✓			
NovaSeq208-12 (1:20)		<input type="checkbox"/>	✓			
H2109P_Wellendorf		<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

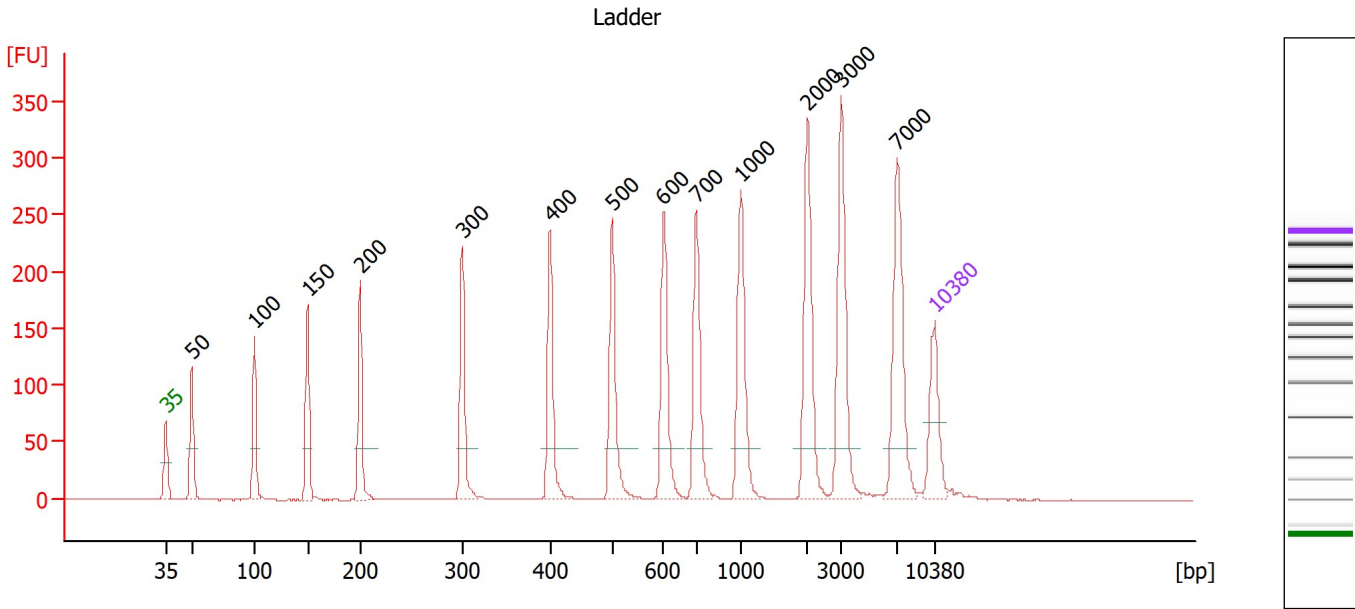
**Ladder**

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: 0.3

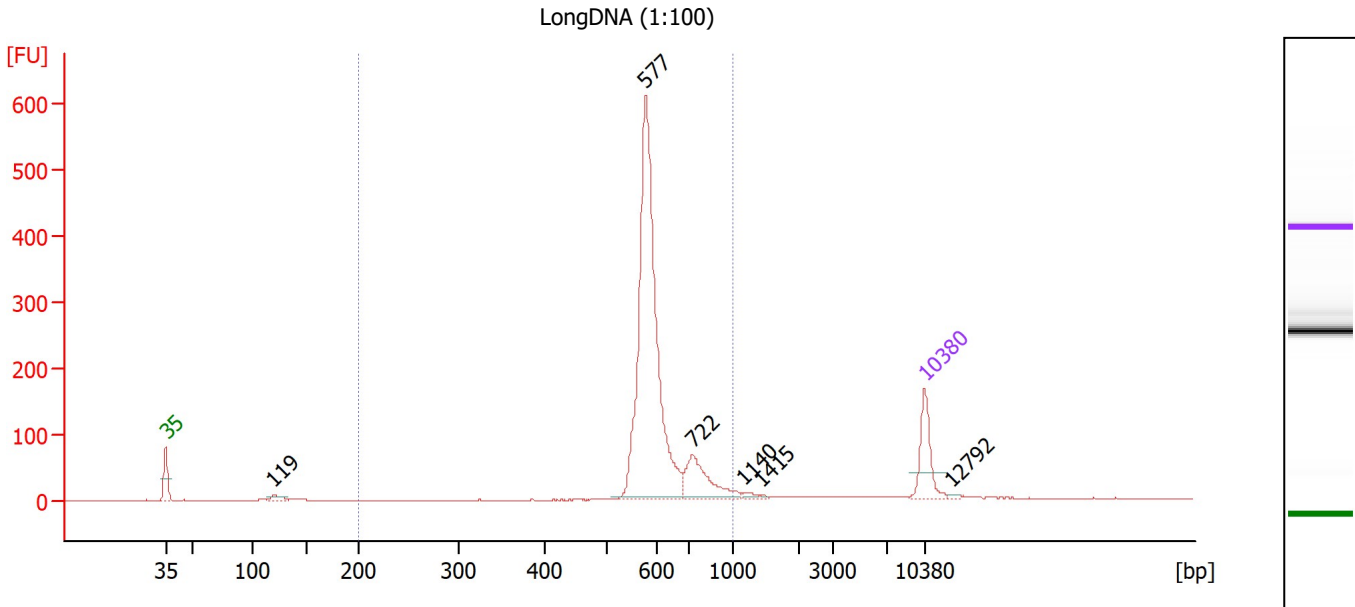
**Peak table for Ladder**

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

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**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : LongDNA (1:100)**

Number of peaks found: 6                      Corr. Area 1: 1,488.0  
 Noise: 0.2

**Peak table for sample 1 : LongDNA (1:100)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	119	19.30	246.0	
3	577	815.33	2,139.7	
4	722	122.65	257.3	
5	1,140	7.95	10.6	
6	1,415	2.41	2.6	
7	10,380	75.00	10.9	Upper Marker
8	12,792	0.00	0.0	

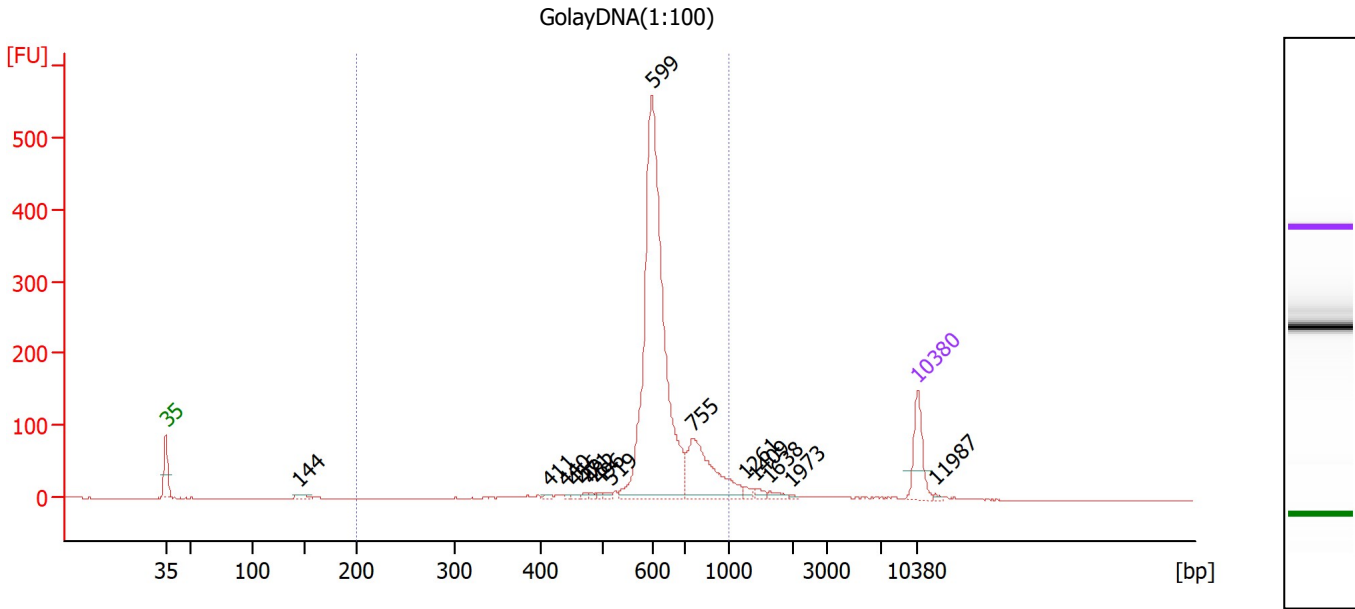
**Region table for sample 1 : LongDNA (1:100)**

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	1,488.0	92	612	12.5	922.53	2,313.9	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : GolayDNA(1:100)**

Number of peaks found: 15                      Corr. Area 1: 1,655.2  
 Noise: 0.5

**Peak table for sample 2 : GolayDNA(1:100)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	144	11.91	125.0	
3	411	5.06	18.6	
4	440	3.75	12.9	
5	456	5.28	17.6	
6	471	4.72	15.2	
7	482	5.49	17.3	
8	496	5.53	16.9	
9	519	7.27	21.3	
10	599	989.29	2,503.8	
11	755	210.42	422.2	
12	1,261	10.28	12.3	
13	1,409	10.33	11.1	
14	1,638	10.77	10.0	
15	1,973	2.59	2.0	
16	10,380	75.00	10.9	Upper Marker
17	11,987	0.00	0.0	

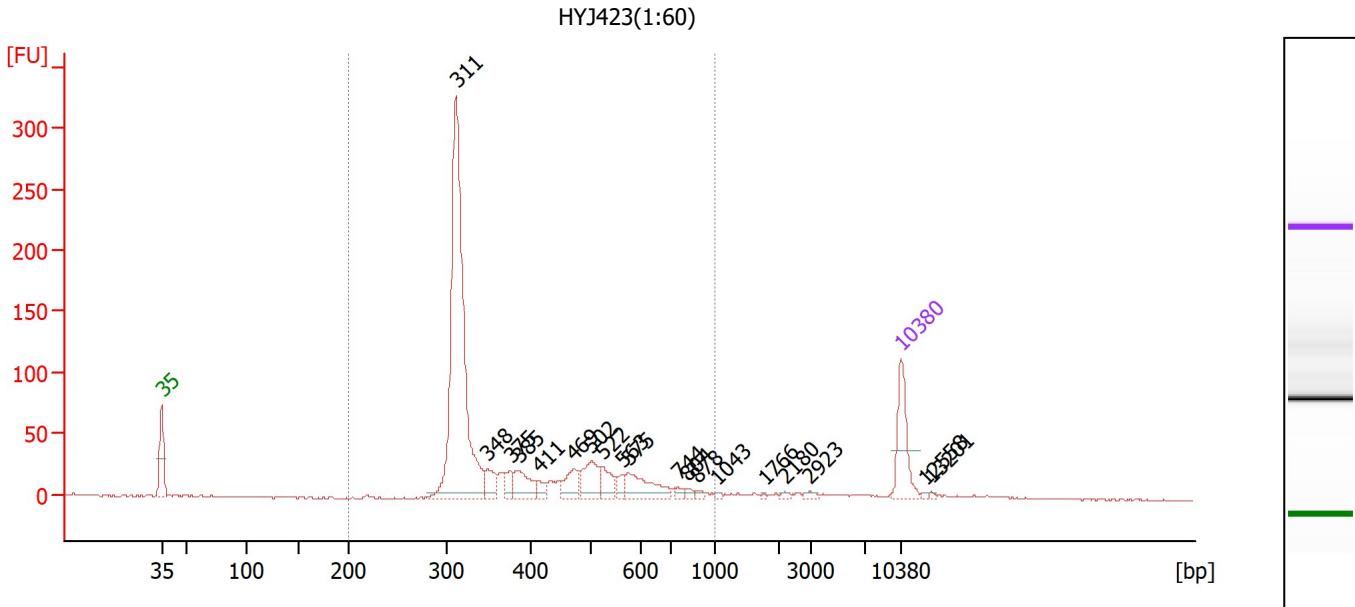
**Region table for sample 2 : GolayDNA(1:100)**

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	1,655.2	92	637	15.3	1,243.55	3,039.4	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : HYJ423(1:60)**

Number of peaks found: 19                      Corr. Area 1: 1,048.6  
 Noise: 0.5

**Peak table for sample 3 : HYJ423(1:60)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	311	718.82	3,503.8	
3	348	39.20	170.7	
4	375	26.75	108.0	
5	385	54.61	215.2	
6	411	16.84	62.0	
7	469	41.80	134.9	
8	502	61.29	185.0	
9	522	38.06	110.4	
10	563	15.87	42.7	
11	575	66.94	176.4	
12	744	8.33	17.0	
13	804	6.96	13.1	
14	878	5.70	9.8	
15	1,043	3.07	4.5	
16	1,766	2.06	1.8	
17	2,180	3.96	2.8	
18	2,923	5.69	2.9	
19	10,380	75.00	10.9	Upper Marker
20	12,558	0.00	0.0	
21	13,201	0.00	0.0	

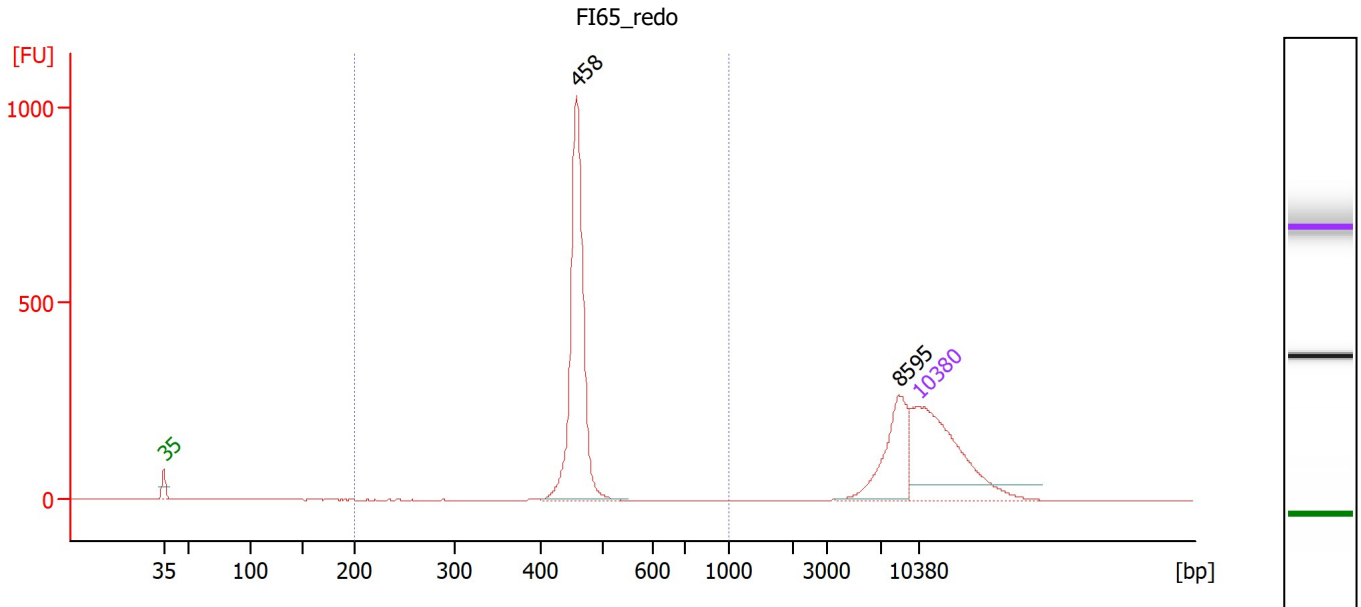
**Region table for sample 3 : HYJ423(1:60)**

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	1,048.6	92	398	32.0	1,110.54	4,674.8	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : FI65\_redo**

Number of peaks found: 2                      Corr. Area 1: 1,768.5  
 Noise: 0.3

**Peak table for sample 4 : FI65\_redo**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	458	163.47	540.7	
3	8,595	42.80	7.5	
4	10,380	75.00	10.9	Upper Marker

**Region table for sample 4 : FI65\_redo**

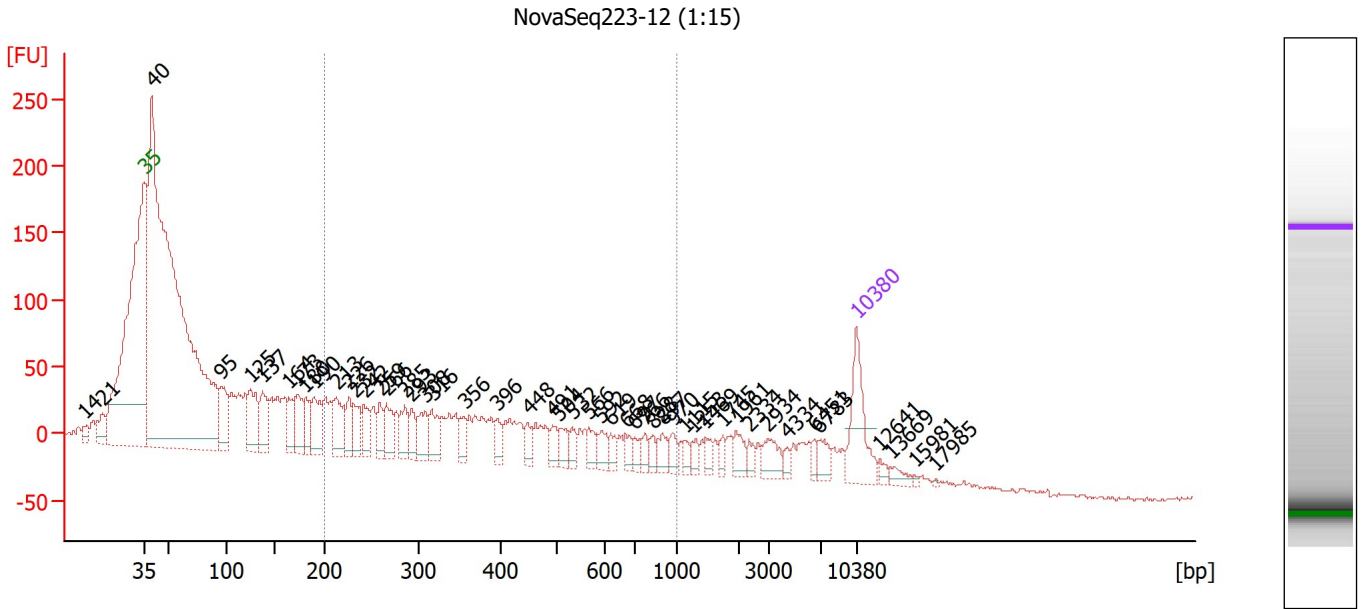
From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	1,768.5	74	459	2.8	161.83	534.1	Blue



Assay Class: High Sensitivity DNA Assay  
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**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : NovaSeq223-12 (1:15)**

Number of peaks found: 49                      Corr. Area 1: 1,213.6  
 Noise: 1.5

**Peak table for sample 5 : NovaSeq223-12 (1:15)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	14	0.00	0.0	
2	21	0.00	0.0	
3	35	125.00	5,411.3	Lower Marker
4	40	2,103.95	79,782.4	
5	95	95.33	1,520.1	
6	125	96.04	1,164.7	
7	137	73.87	814.9	
8	164	54.80	506.5	
9	173	69.67	611.4	
10	180	44.93	377.6	
11	190	69.36	552.8	
12	213	67.71	481.9	
13	226	38.42	257.0	
14	232	41.14	269.2	
15	242	41.62	260.3	
16	259	33.36	195.0	
17	266	47.07	268.1	
18	285	43.26	229.8	
19	293	34.29	177.6	
20	308	43.93	216.3	
21	316	44.77	214.5	
22	356	26.93	114.6	
23	396	25.53	97.7	
24	448	19.80	66.9	
25	491	22.89	70.6	
26	504	24.51	73.7	

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**Electropherogram Summary Continued ...**

**... Peak table for sample 5 : NovaSeq223-12 (1:15)**

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
27	532	18.43	52.5	
28	566	20.86	55.8	
29	592	21.77	55.7	
30	619	15.12	37.0	
31	668	14.33	32.5	
32	697	14.59	31.7	
33	766	13.04	25.8	
34	828	13.16	24.1	
35	897	22.31	37.7	
36	970	17.14	26.8	
37	1,115	10.34	14.1	
38	1,253	13.91	16.8	
39	1,469	12.93	13.3	
40	1,745	11.15	9.7	
41	1,961	22.23	17.2	
42	2,334	10.79	7.0	
43	2,934	27.81	14.4	
44	4,334	8.88	3.1	
45	6,451	10.30	2.4	
46	6,783	19.71	4.4	
47	10,380	75.00	10.9	Upper Marker
48	12,641	0.00	0.0	
49	13,669	0.00	0.0	
50	15,981	0.00	0.0	
51	17,985	0.00	0.0	

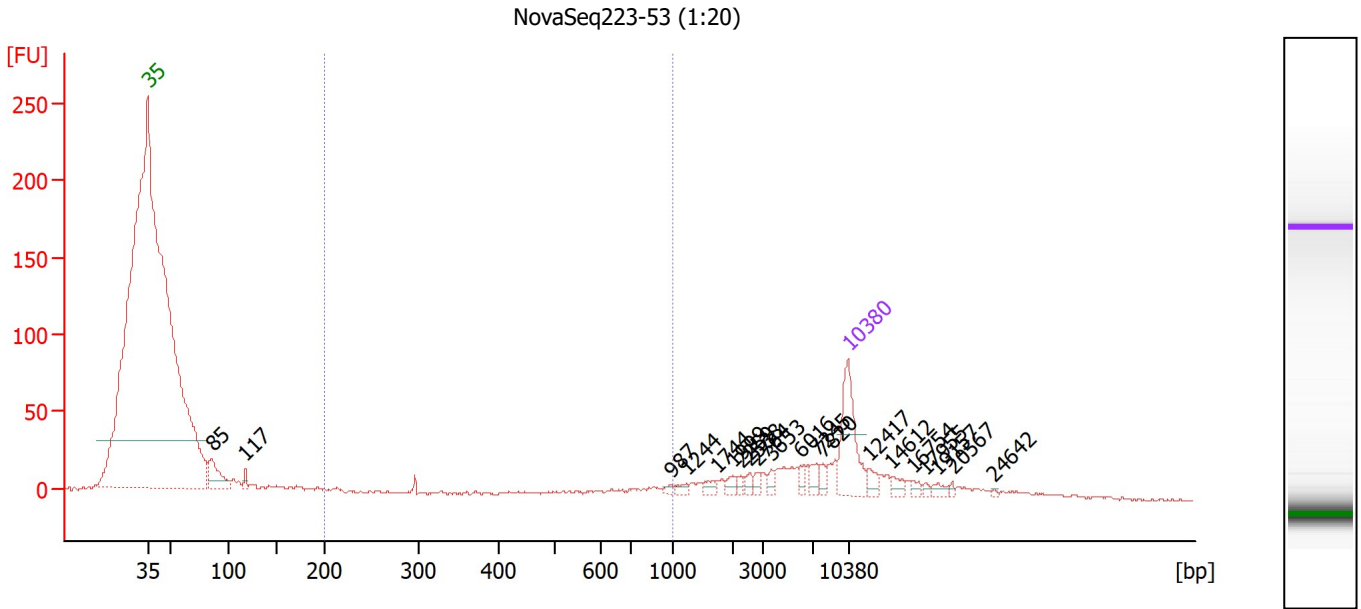
**Region table for sample 5 : NovaSeq223-12 (1:15)**

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	1,213.6	30	425	44.5	927.99	4,205.4	<span style="color: blue;">■</span>

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**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : NovaSeq223-53 (1:20)**

Number of peaks found: 20                      Corr. Area 1: 37.9  
 Noise: 1.0

**Peak table for sample 6 : NovaSeq223-53 (1:20)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	85	67.77	1,203.2	
3	117	11.75	151.6	
4	987	3.94	6.0	
5	1,244	5.33	6.5	
6	1,744	8.41	7.3	
7	1,999	7.43	5.6	
8	2,259	5.10	3.4	
9	2,538	6.81	4.1	
10	2,764	6.91	3.8	
11	3,653	6.41	2.7	
12	6,016	7.91	2.0	
13	7,245	11.58	2.4	
14	7,820	8.81	1.7	
15	10,380	75.00	10.9	Upper Marker
16	12,417	0.00	0.0	
17	14,612	0.00	0.0	
18	16,754	0.00	0.0	
19	17,955	0.00	0.0	
20	19,157	0.00	0.0	
21	20,567	0.00	0.0	
22	24,642	0.00	0.0	

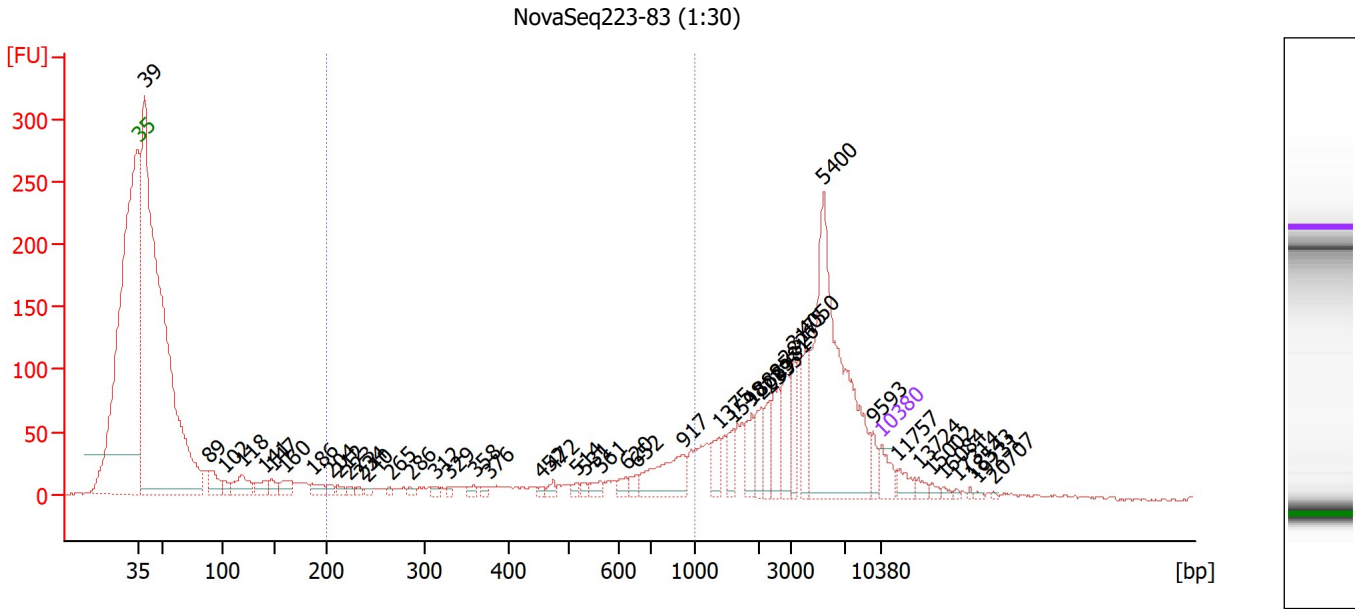
**Region table for sample 6 : NovaSeq223-53 (1:20)**

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	37.9	8	685	31.2	31.42	89.1	Blue

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**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : NovaSeq223-83 (1:30)**

Number of peaks found: 46                      Corr. Area 1: 490.0  
 Noise: 0.9

**Peak table for sample 7 : NovaSeq223-83 (1:30)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	39	6,158.00	238,560.9	
3	89	138.31	2,363.8	
4	102	50.11	746.5	
5	118	161.24	2,066.3	
6	141	80.86	869.2	
7	147	68.86	709.0	
8	160	83.66	791.4	
9	186	66.85	543.8	
10	204	30.71	227.9	
11	212	33.35	238.0	
12	223	21.97	149.2	
13	234	20.96	135.5	
14	240	16.14	101.7	
15	265	17.55	100.4	
16	286	22.70	120.4	
17	312	22.57	109.7	
18	329	16.36	75.3	
19	358	26.00	110.0	
20	376	19.66	79.3	
21	452	14.14	47.4	
22	472	30.27	97.1	
23	514	21.12	62.3	
24	531	24.42	69.7	
25	561	36.75	99.2	
26	620	38.77	94.7	

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**Electropherogram Summary Continued ...**

**... Peak table for sample 7 : NovaSeq223-83 (1:30)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	652	39.27	91.2	
28	917	258.72	427.6	
29	1,375	93.35	102.8	
30	1,598	73.00	69.2	
31	1,888	100.04	80.3	
32	2,089	102.66	74.5	
33	2,335	83.75	54.4	
34	2,581	136.51	80.2	
35	2,826	144.72	77.6	
36	3,175	115.44	55.1	
37	4,050	148.33	55.5	
38	5,400	1,144.74	321.2	
39	9,593	53.94	8.5	
40	10,380	75.00	10.9	Upper Marker
41	11,757	0.00	0.0	
42	13,724	0.00	0.0	
43	15,002	0.00	0.0	
44	16,084	0.00	0.0	
45	17,314	0.00	0.0	
46	18,543	0.00	0.0	
47	19,231	0.00	0.0	
48	20,707	0.00	0.0	

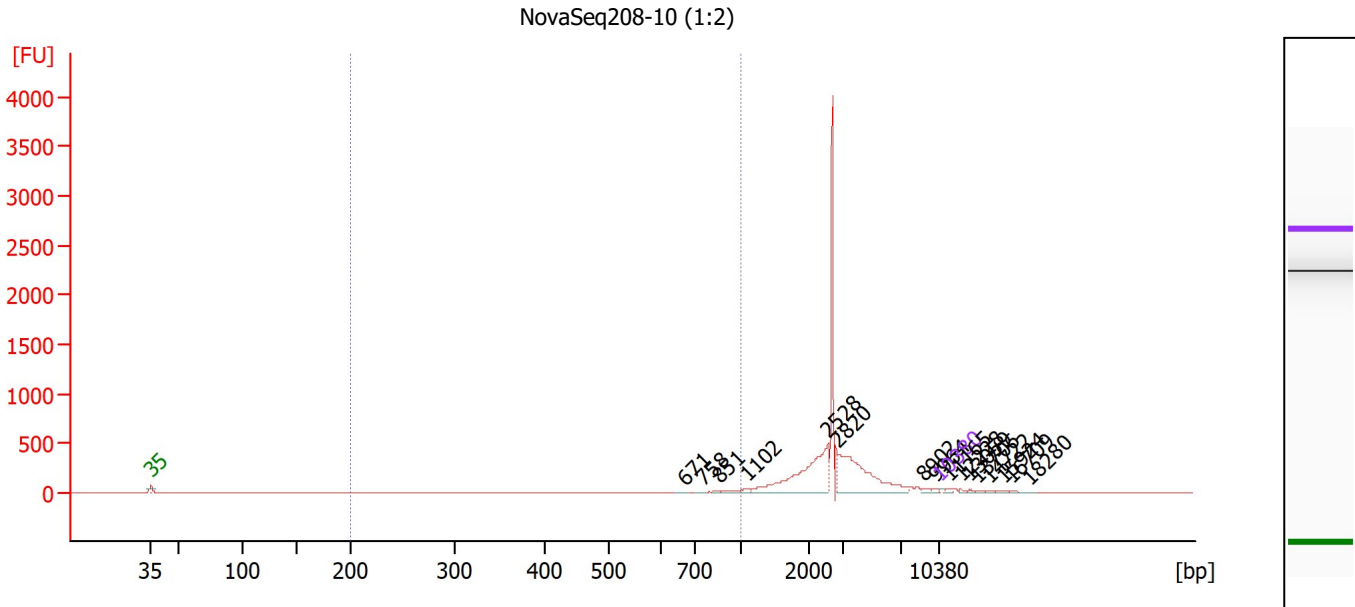
**Region table for sample 7 : NovaSeq223-83 (1:30)**

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	490.0	12	576	40.8	1,110.15	3,979.1	

Assay Class: High Sensitivity DNA Assay  
 Data Path: Z:\...\Bioanalyzer1\_High Sensitivity DNA Assay\_2020-07-02\_001.xad

Created: 7/2/2020 2:43:10 PM  
 Modified: 7/2/2020 3:24:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : NovaSeq208-10 (1:2)**

Number of peaks found: 16                      Corr. Area 1: 75.4  
 Noise: 0.7

**Peak table for sample 8 : NovaSeq208-10 (1:2)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	671	31.43	71.0	
3	758	43.33	86.7	
4	851	55.79	99.3	
5	1,102	305.22	419.8	
6	2,528	4,910.01	2,943.1	
7	2,820	4,804.16	2,580.8	
8	8,902	134.61	22.9	
9	9,964	102.79	15.6	
10	10,380	75.00	10.9	Upper Marker
11	11,165	0.00	0.0	
12	12,228	0.00	0.0	
13	13,059	0.00	0.0	
14	13,706	0.00	0.0	
15	14,722	0.00	0.0	
16	15,924	0.00	0.0	
17	16,709	0.00	0.0	
18	18,280	0.00	0.0	

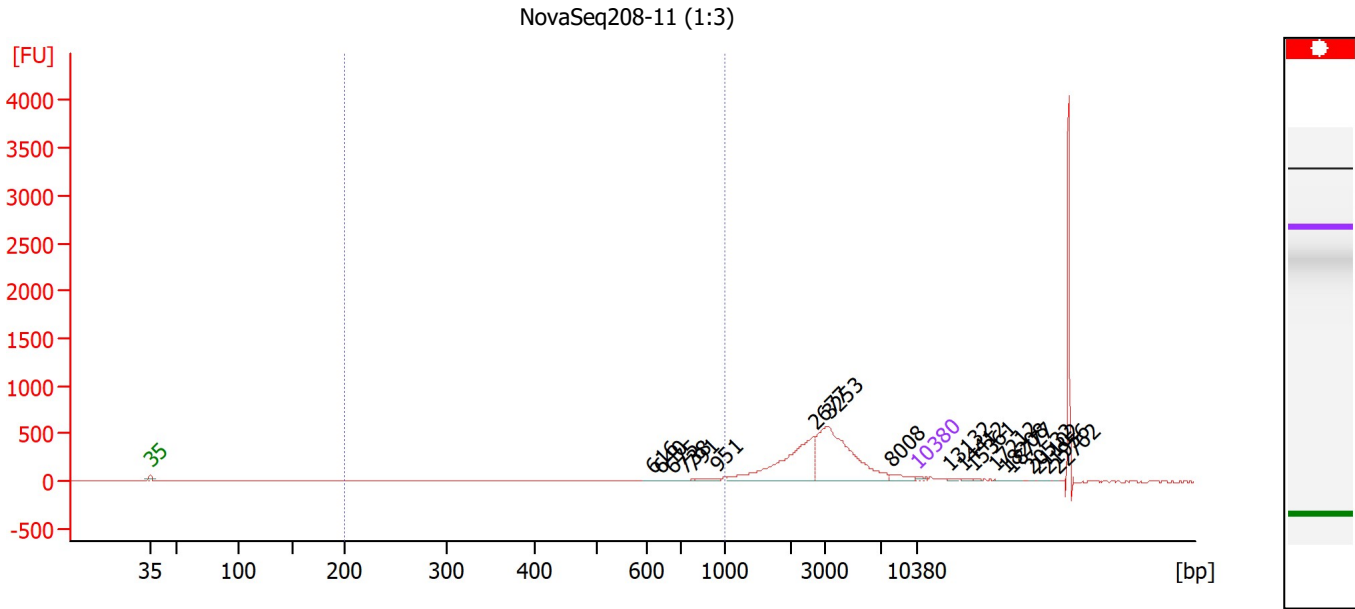
**Region table for sample 8 : NovaSeq208-10 (1:2)**

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	75.4	2	853	13.1	331.14	598.1	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: Z:\...\Bioanalyzer1\_High Sensitivity DNA Assay\_2020-07-02\_001.xad

Created: 7/2/2020 2:43:10 PM  
 Modified: 7/2/2020 3:24:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : NovaSeq208-11 (1:3)**

Number of peaks found: 19                      Corr. Area 1: 192.0  
 Noise: 0.9

**Peak table for sample 9 : NovaSeq208-11 (1:3)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	616	17.18	42.3	
3	640	19.66	46.5	
4	675	24.53	55.1	
5	738	44.54	91.5	
6	791	27.94	53.5	
7	951	135.72	216.2	
8	2,677	2,744.13	1,552.9	
9	3,253	3,619.18	1,685.6	
10	8,008	229.06	43.3	
11	10,380	75.00	10.9	Upper Marker
12	13,132	0.00	0.0	
13	14,412	0.00	0.0	
14	15,361	0.00	0.0	
15	17,212	0.00	0.0	
16	18,208	0.00	0.0	
17	18,777	0.00	0.0	
18	20,532	0.00	0.0	
19	21,102	0.00	0.0	
20	21,956	0.00	0.0	
21	22,762	0.00	0.0	

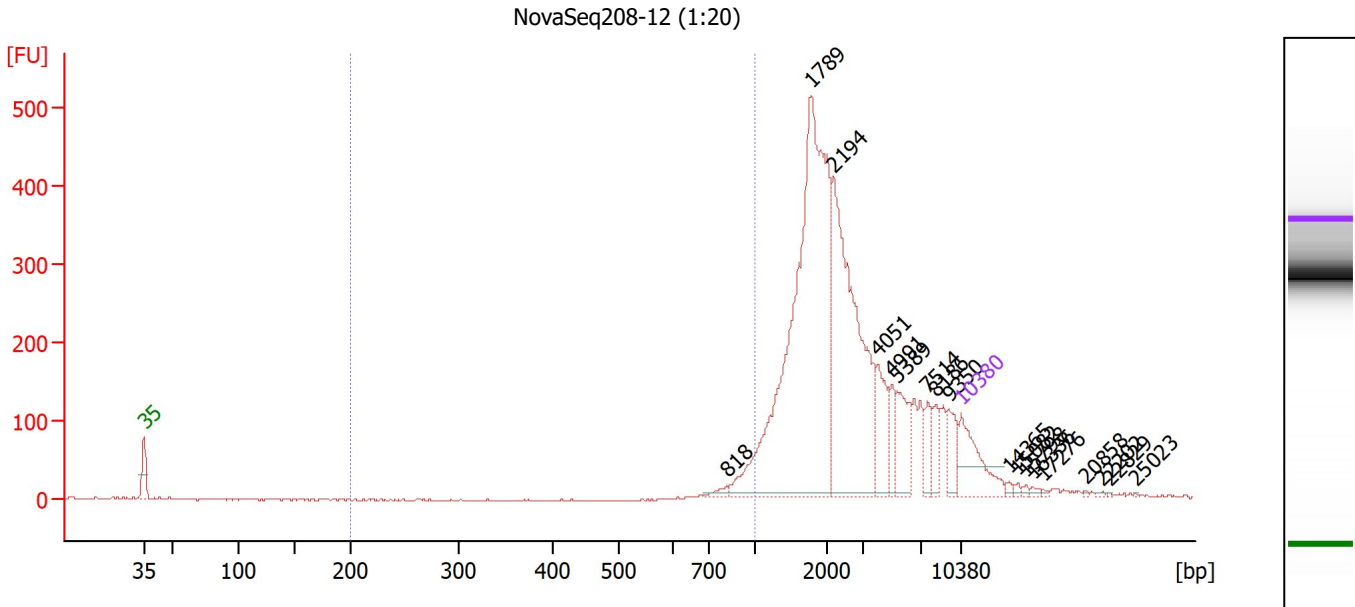
**Region table for sample 9 : NovaSeq208-11 (1:3)**

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	192.0	4	763	22.9	400.40	893.2	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: Z:\...\Bioanalyzer1\_High Sensitivity DNA Assay\_2020-07-02\_001.xad

Created: 7/2/2020 2:43:10 PM  
 Modified: 7/2/2020 3:24:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : NovaSeq208-12 (1:20)**

Number of peaks found: 18                      Corr. Area 1: 100.7  
 Noise: 0.7

**Peak table for sample 10 : NovaSeq208-12 (1:20)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	818	10.06	18.6	
3	1,789	754.56	638.9	
4	2,194	405.30	279.8	
5	4,051	68.76	25.7	
6	4,991	32.12	9.7	
7	5,389	67.40	18.9	
8	7,514	30.35	6.1	
9	8,186	29.32	5.4	
10	9,350	38.26	6.2	
11	10,380	75.00	10.9	Upper Marker
12	14,365	0.00	0.0	
13	15,082	0.00	0.0	
14	15,798	0.00	0.0	
15	16,336	0.00	0.0	
16	17,276	0.00	0.0	
17	20,858	0.00	0.0	
18	22,202	0.00	0.0	
19	22,829	0.00	0.0	
20	25,023	0.00	0.0	

**Region table for sample 10 : NovaSeq208-12 (1:20)**

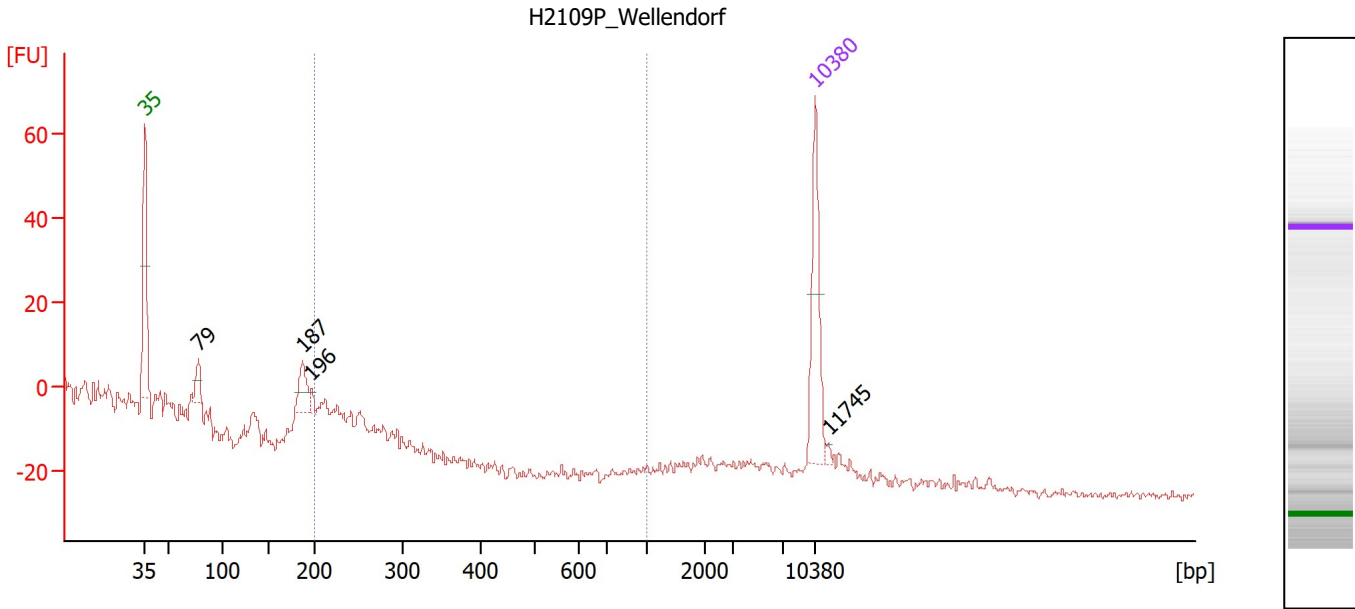
From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	100.7	3	900	8.9	45.02	76.0	Blue



Assay Class: High Sensitivity DNA Assay  
 Data Path: Z:\...\Bioanalyzer1\_High Sensitivity DNA Assay\_2020-07-02\_001.xad

Created: 7/2/2020 2:43:10 PM  
 Modified: 7/2/2020 3:24:32 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : H2109P Wellendorf**

Number of peaks found: 4                      Corr. Area 1: 21.9  
 Noise: 1.2

**Peak table for sample 11 : H2109P Wellendorf**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	79	26.41	509.2	
3	187	37.72	305.4	
4	196	8.94	69.0	
5	10,380	75.00	10.9	Upper Marker
6	11,745	0.00	0.0	

**Region table for sample 11 : H2109P Wellendorf**

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	21.9	18	223	7.6	41.91	284.4	Blue

Assay Class: High Sensitivity DNA Assay  
Data Path: Z:\...\Bioanalyzer1\_High Sensitivity DNA Assay\_2020-07-02\_001.xad

Created: 7/2/2020 2:43:10 PM  
Modified: 7/2/2020 3:24:32 PM

**Gel Image**

Assay Class: High Sensitivity DNA Assay Created: 7/2/2020 2:43:10 PM  
 Data Path: Z:\...\Bioanalyzer1\_High Sensitivity DNA Assay\_2020-07-02\_001.xad Modified: 7/2/2020 3:24:32 PM

**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		7/2/2020 3:24:29 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Instrument error occurred on port 1, Optical signal too high (1605h)	559	Instrument	Run	Sample 9	7/2/2020 3:18:28 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Run started on port 1 (File: Z:\XADs\2020-07-02\Bioanalyze r1_High Sensitivity DNA Assay_2020-07-02_001.xad)		Instrument	Run		7/2/2020 2:43:15 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Product Number : G2938C		Instrument	Run		7/2/2020 2:43:15 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Name :		Instrument	Run		7/2/2020 2:43:15 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Vendor : Agilent Technologies		Instrument	Run		7/2/2020 2:43:15 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Serial# : DE34903152		Instrument	Run		7/2/2020 2:43:15 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Firmware : C.01.069		Instrument	Run		7/2/2020 2:43:15 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Cartridge : Electrode		Instrument	Run		7/2/2020 2:43:15 PM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB