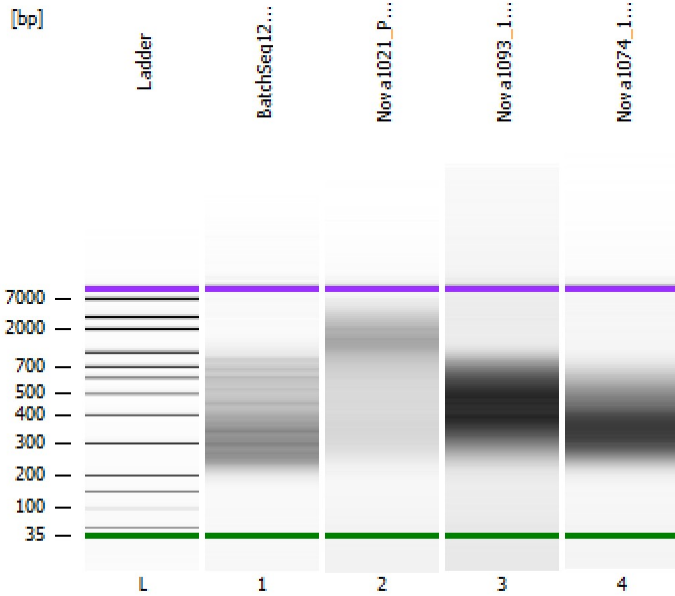


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...pert\data\2024-05-21\BatchSeq128\_N1021P-D\_N1093P\_N1074P.xad

Created: 5/21/2024 11:08:49 AM  
Modified: 5/21/2024 12:24:48 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

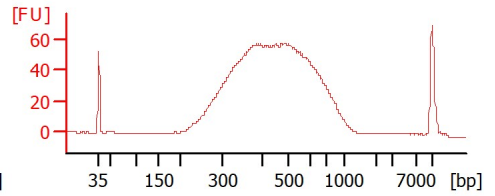
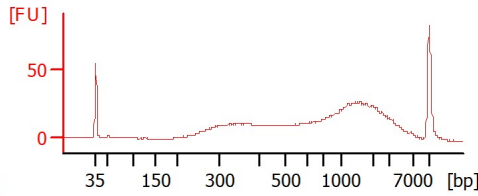
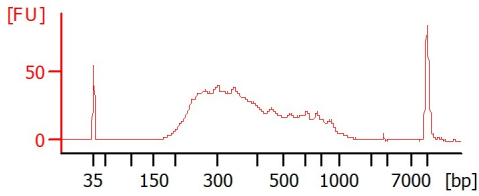
Chip Information:

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

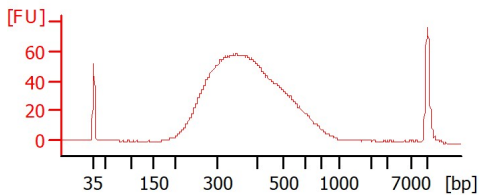
BatchSeq128\_1.2xBC (1:5)

Nova1021\_PoolID\_1.2xBC on 1xBC (1:2)

Nova1093\_1.2xBC (1:2)



Nova1074\_1.2xBC (1:2)



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...pert\data\2024-05-21\BatchSeq128\_N1021P-D\_N1093P\_N1074P.xad

Created: 5/21/2024 11:08:49 AM  
 Modified: 5/21/2024 12:24:48 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
BatchSeq128_1.2xBC (1:5)		<input type="checkbox"/>	✓			
Nova1021_PoolD_1.2xBC on 1xBC (1:2)		<input type="checkbox"/>	✓			
Nova1093_1.2xBC (1:2)		<input type="checkbox"/>	✓			
Nova1074_1.2xBC (1:2)		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...pert\data\2024-05-21\BatchSeq128\_N1021P-D\_N1093P\_N1074P.xad

Created: 5/21/2024 11:08:49 AM  
Modified: 5/21/2024 12:24:48 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

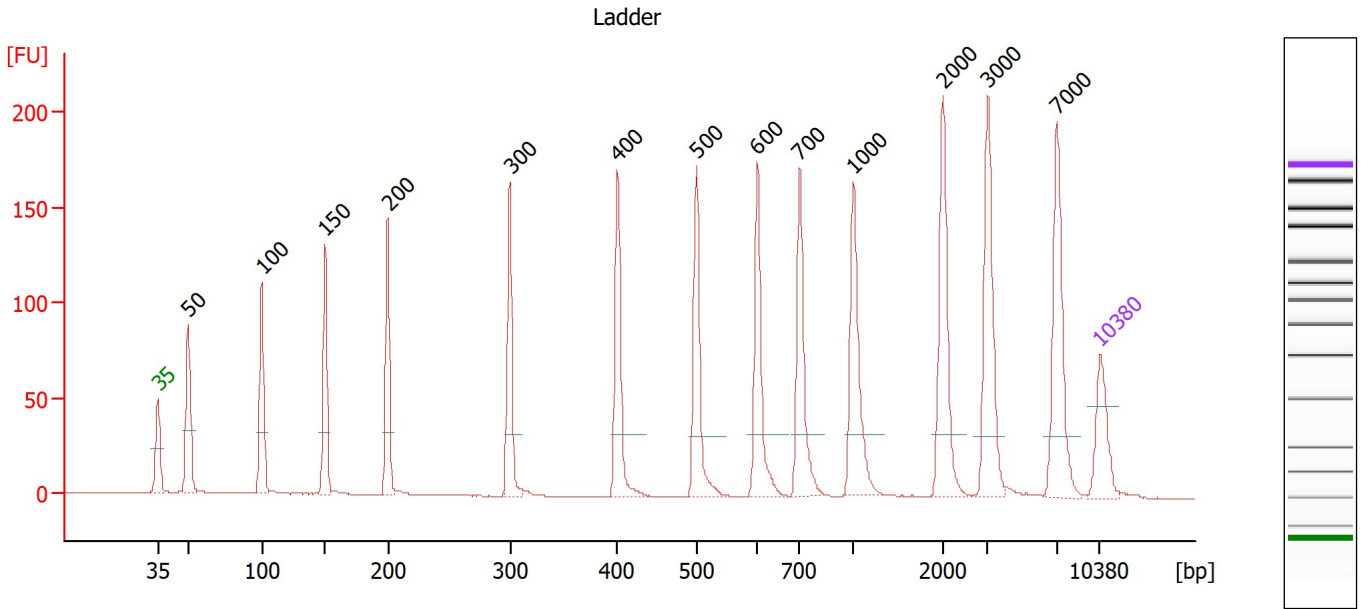
**Ladder**

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

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...pert\data\2024-05-21\BatchSeq128\_N1021P-D\_N1093P\_N1074P.xad

Created: 5/21/2024 11:08:49 AM  
 Modified: 5/21/2024 12:24:48 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.2

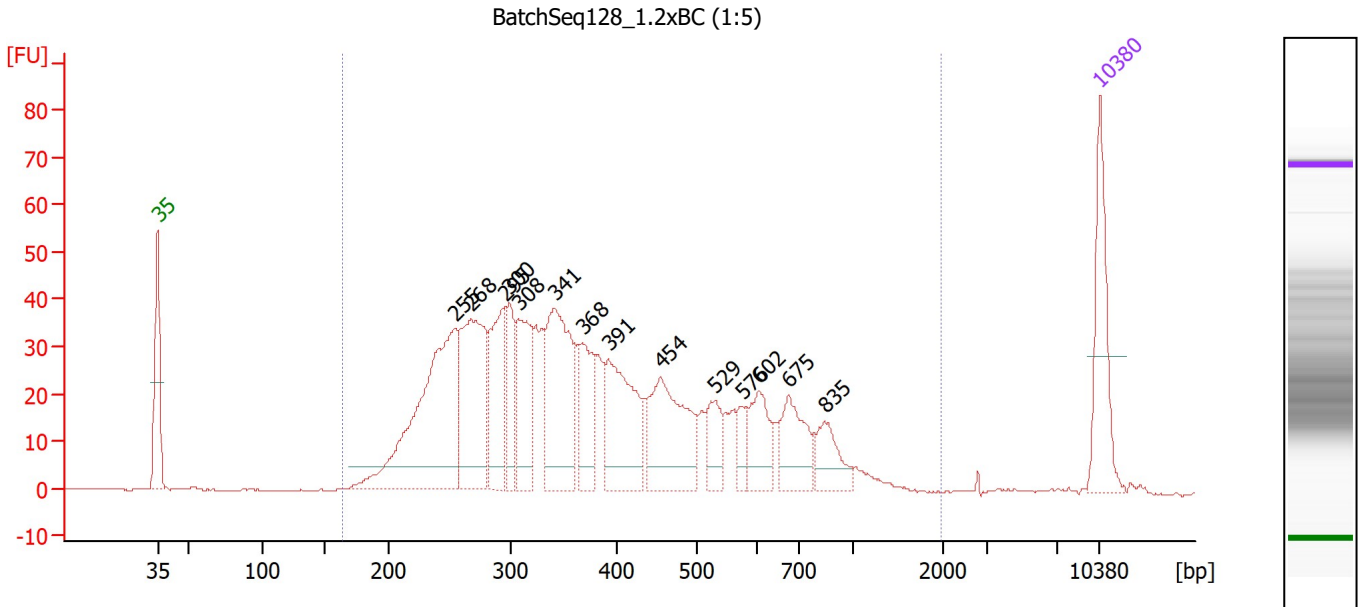
**Peak table for Ladder**

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

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...pert\data\2024-05-21\BatchSeq128\_N1021P-D\_N1093P\_N1074P.xad

Created: 5/21/2024 11:08:49 AM  
 Modified: 5/21/2024 12:24:48 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : BatchSeq128 1.2xBC (1:5)**

Number of peaks found: 14                      Corr. Area 1: 1,131.3  
 Noise: 0.2

**Peak table for sample 1 : BatchSeq128 1.2xBC (1:5)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	255	326.31	1,937.6	
3	268	193.04	1,089.8	
4	295	112.36	576.5	
5	300	63.28	319.7	
6	308	107.16	527.7	
7	341	175.74	780.1	
8	368	75.39	310.5	
9	391	133.15	515.7	
10	454	138.90	463.2	
11	529	38.99	111.6	
12	576	19.39	51.0	
13	602	56.17	141.4	
14	675	64.69	145.2	
15	835	41.37	75.0	
16	10,380	75.00	10.9	Upper Marker

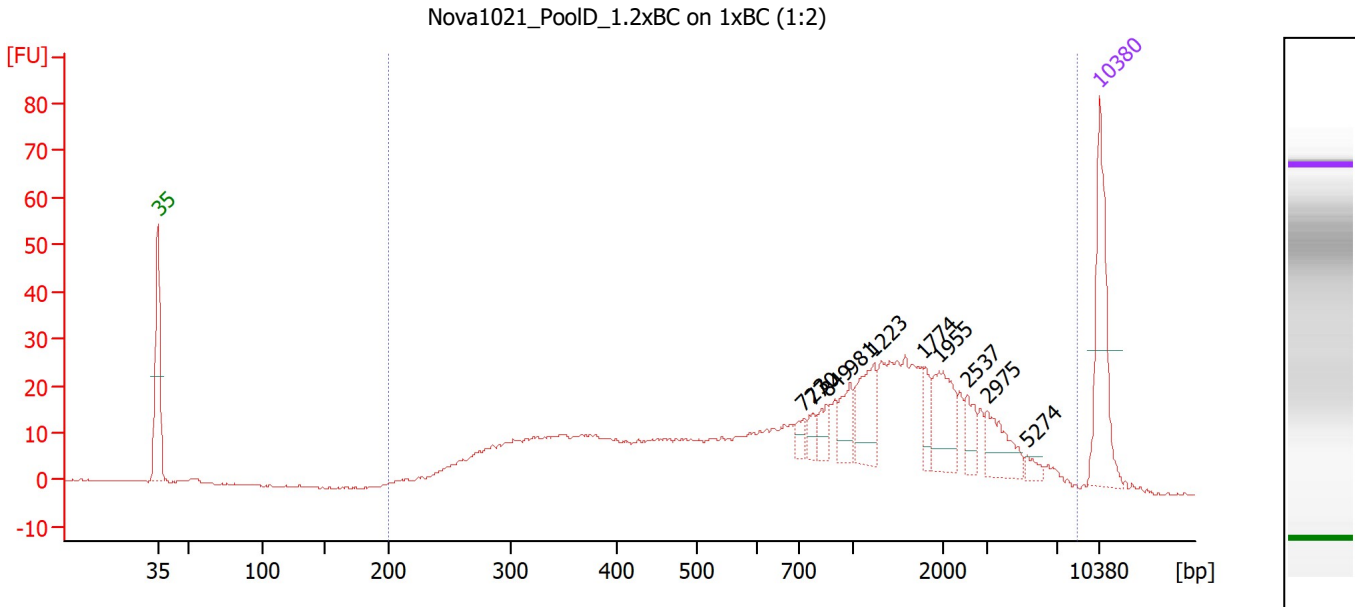
**Region table for sample 1 : BatchSeq128 1.2xBC (1:5)**

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
164	1,977	1,131.2	98	418	47.8	1,827.66	8,196.9	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...pert\data\2024-05-21\BatchSeq128\_N1021P-D\_N1093P\_N1074P.xad

Created: 5/21/2024 11:08:49 AM  
 Modified: 5/21/2024 12:24:48 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : Nova1021 PoolD 1.2xBC on 1xBC (1:2)**

Number of peaks found: 10                      Corr. Area 1: 724.3  
 Noise: 0.2

**Peak table for sample 2 : Nova1021 PoolD 1.2xBC on 1xBC (1:2)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	723	10.58	22.2	
3	770	11.19	22.0	
4	849	16.25	29.0	
5	981	26.12	40.3	
6	1,223	44.50	55.1	
7	1,774	17.87	15.3	
8	1,955	45.01	34.9	
9	2,537	17.71	10.6	
10	2,975	32.60	16.6	
11	5,274	6.66	1.9	
12	10,380	75.00	10.9	Upper Marker

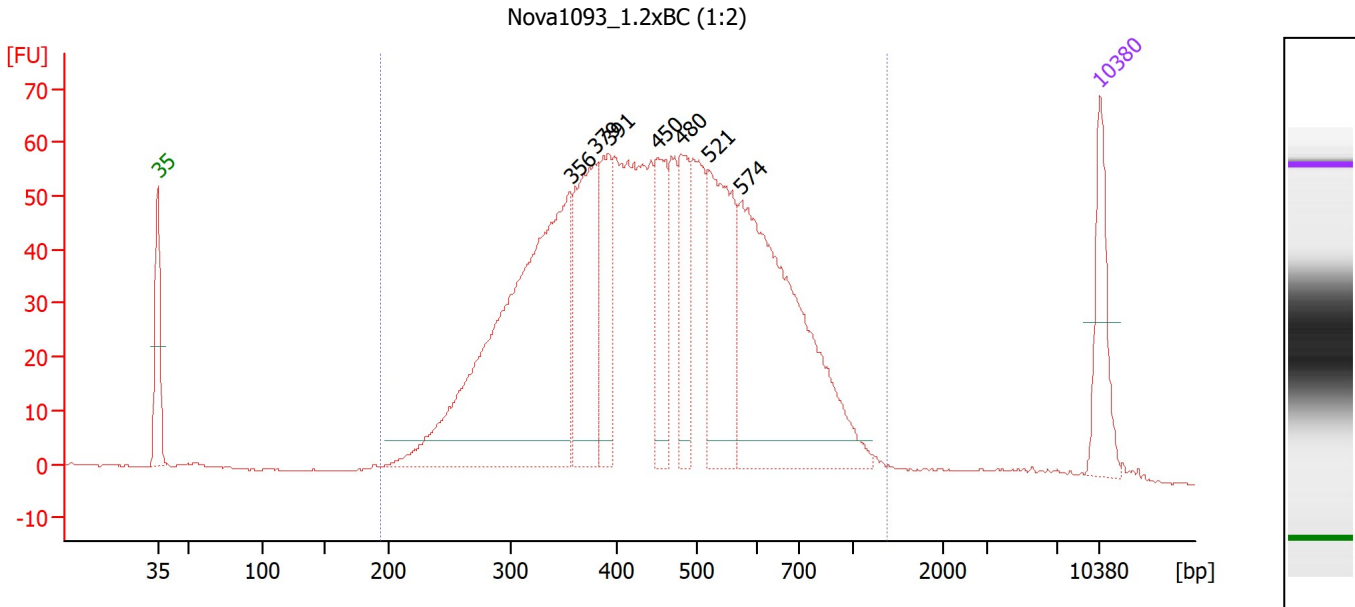
**Region table for sample 2 : Nova1021 PoolD 1.2xBC on 1xBC (1:2)**

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	8,646	724.3	99	1,358	97.3	1,086.61	2,848.2	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...pert\data\2024-05-21\BatchSeq128\_N1021P-D\_N1093P\_N1074P.xad

Created: 5/21/2024 11:08:49 AM  
 Modified: 5/21/2024 12:24:48 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : Nova1093 1.2xBC (1:2)**

Number of peaks found: 7                      Corr. Area 1: 1,660.6  
 Noise: 0.1

**Peak table for sample 3 : Nova1093 1.2xBC (1:2)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	356	719.57	3,065.2	
3	379	236.62	947.0	
4	391	128.97	499.3	
5	450	120.72	406.5	
6	480	100.68	317.5	
7	521	223.84	650.7	
8	574	478.24	1,262.9	
9	10,380	75.00	10.9	Upper Marker

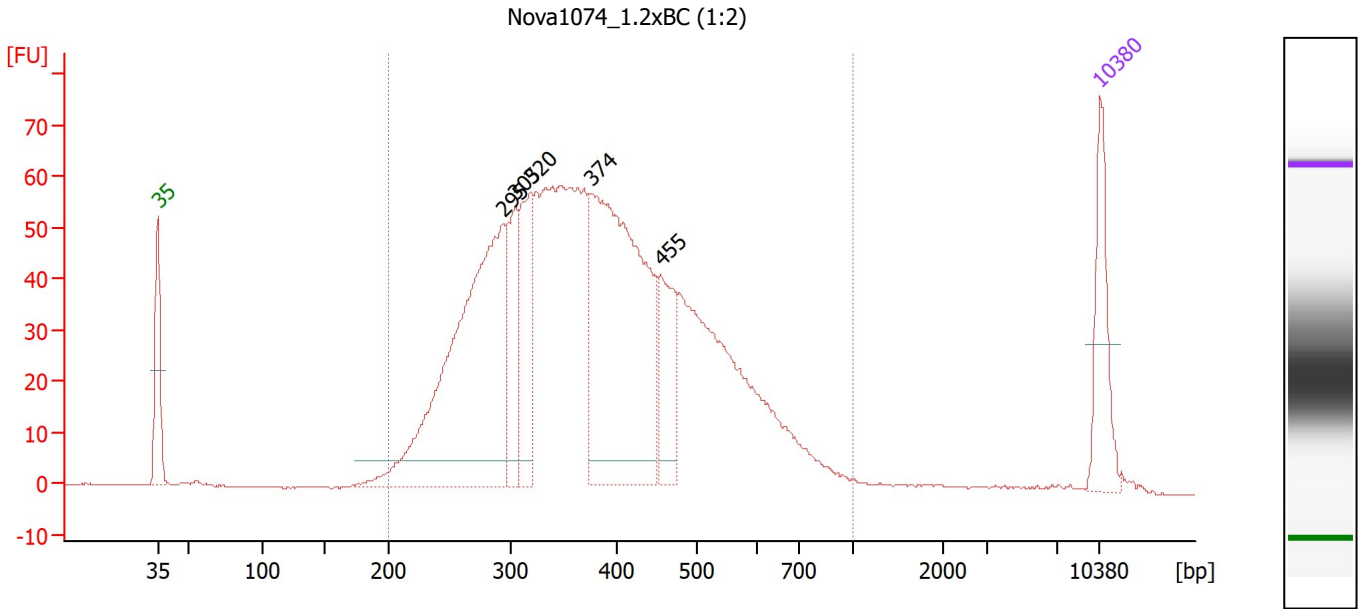
**Region table for sample 3 : Nova1093 1.2xBC (1:2)**

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
195	1,368	1,660.6	96	472	33.9	2,821.02	10,422.1	Blue

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...pert\data\2024-05-21\BatchSeq128\_N1021P-D\_N1093P\_N1074P.xad

Created: 5/21/2024 11:08:49 AM  
 Modified: 5/21/2024 12:24:48 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : Nova1074\_1.2xBC (1:2)**

Number of peaks found: 5                      Corr. Area 1: 1,480.6  
 Noise: 0.2

**Peak table for sample 4 : Nova1074\_1.2xBC (1:2)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	295	584.53	3,006.6	
3	305	101.03	501.8	
4	320	146.60	694.7	
5	374	516.70	2,091.8	
6	455	103.00	343.2	
7	10,380	75.00	10.9	Upper Marker

**Region table for sample 4 : Nova1074\_1.2xBC (1:2)**

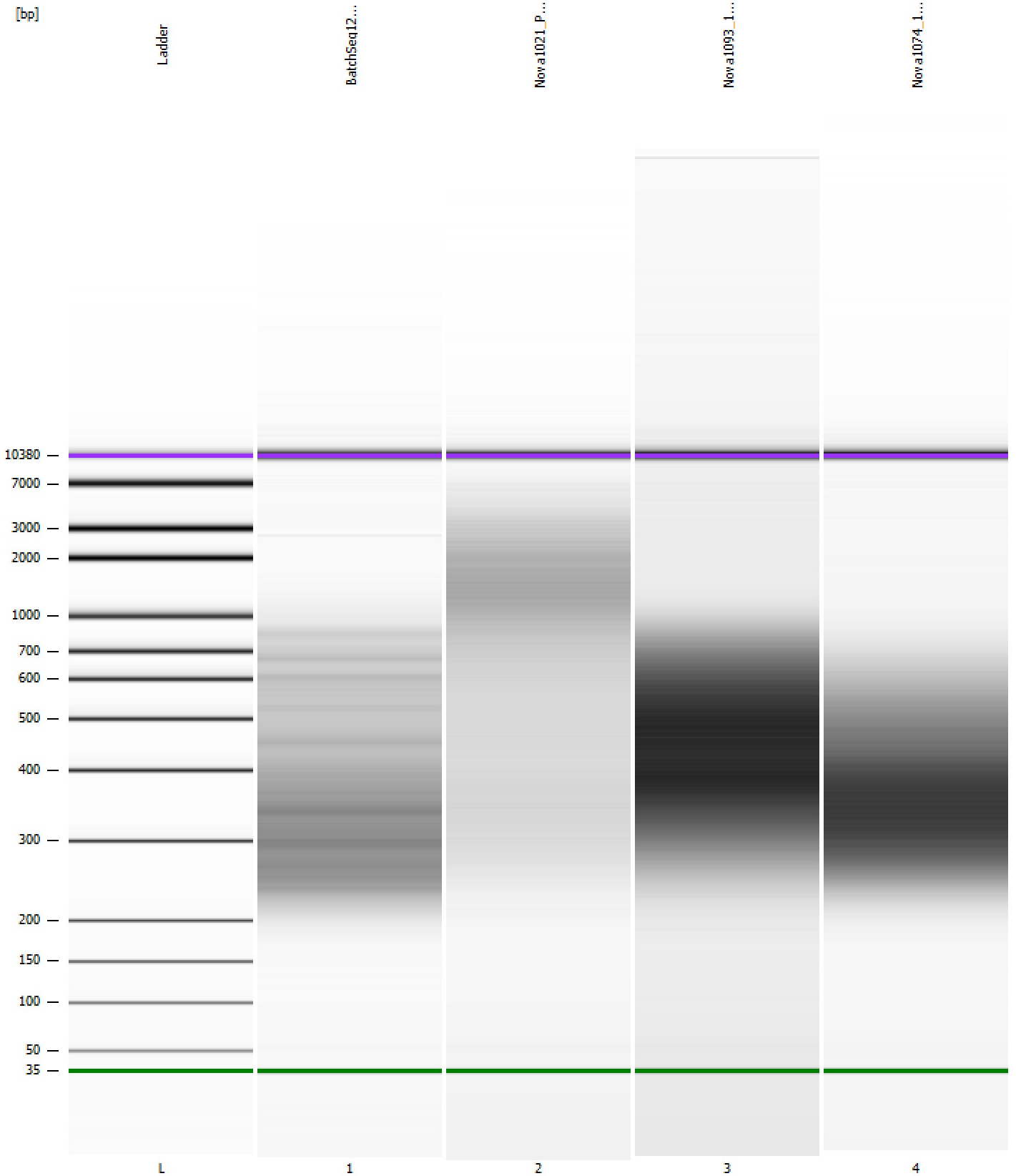
From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,000	1,480.6	98	394	29.7	2,422.26	10,376.2	Blue



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...pert\data\2024-05-21\BatchSeq128\_N1021P-D\_N1093P\_N1074P.xad

Created: 5/21/2024 11:08:49 AM  
Modified: 5/21/2024 12:24:48 PM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...pert\data\2024-05-21\BatchSeq128\_N1021P-D\_N1093P\_N1074P.xad

Created: 5/21/2024 11:08:49 AM  
 Modified: 5/21/2024 12:24:48 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		5/21/2024 11:50:11 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Run started on port 1 (File: C:\Program Files (x86)\Agilent\2100 bioanalyzer\2100 expert\data\2024-05-21\Bioanalyzer1_High Sensitivity DNA Assay_DE34903152_2024-05-21_001.xad)		Instrument	Run		5/21/2024 11:08:56 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Product Number : G2938C		Instrument	Run		5/21/2024 11:08:56 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Name :		Instrument	Run		5/21/2024 11:08:56 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Vendor : Agilent Technologies		Instrument	Run		5/21/2024 11:08:56 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Serial# : DE34903152		Instrument	Run		5/21/2024 11:08:56 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Firmware : C.01.069		Instrument	Run		5/21/2024 11:08:56 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Cartridge : Electrode		Instrument	Run		5/21/2024 11:08:56 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB