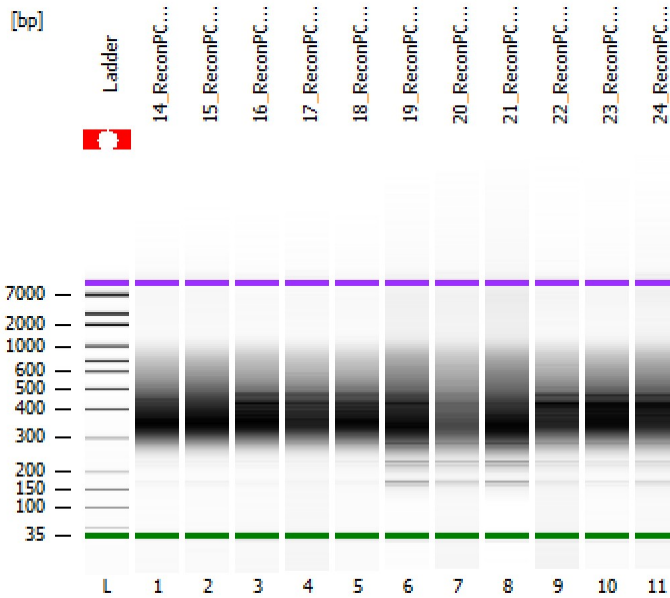


Assay Class: High Sensitivity DNA Assay
Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
Modified: 8/26/2020 11:38:04 AM

Electrophoresis File Run Summary



Instrument Information:

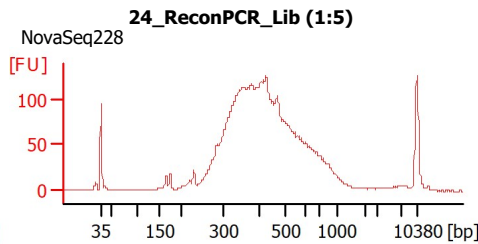
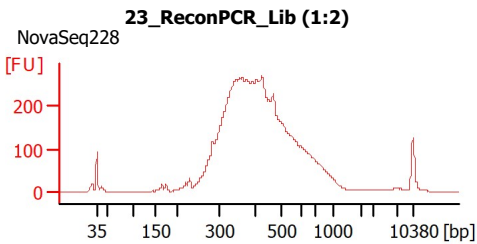
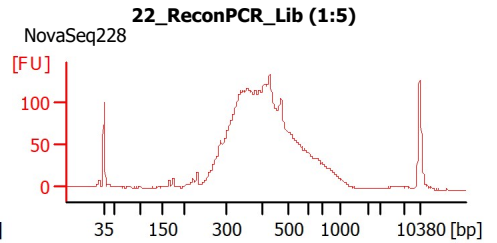
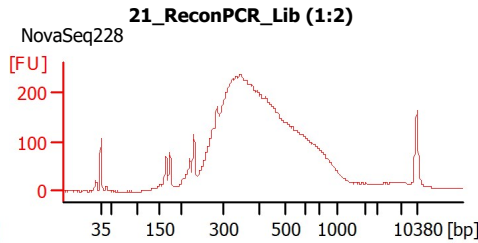
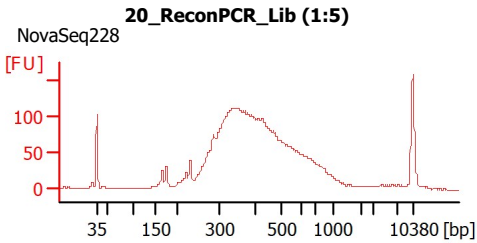
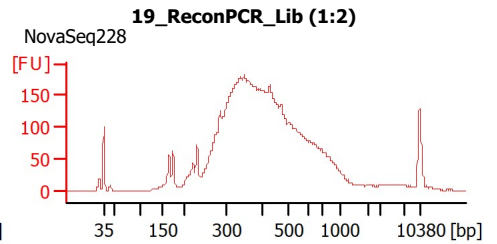
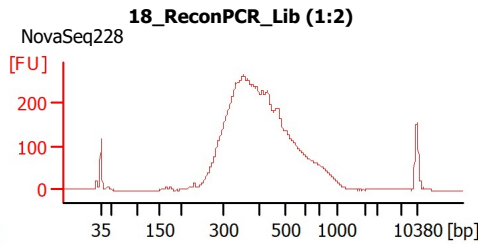
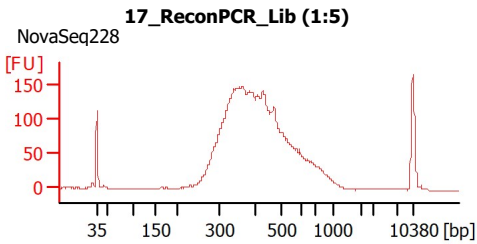
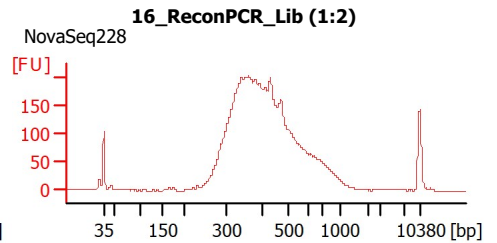
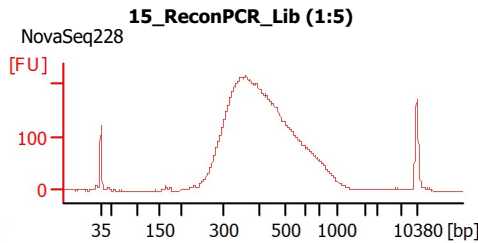
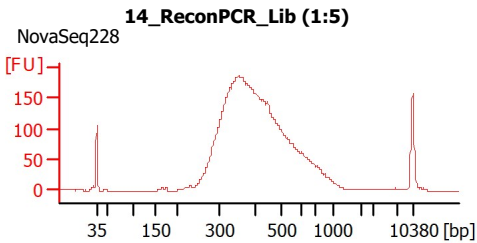
Instrument Name: DE34903152 Firmware: C.01.069
Serial#: DE34903152 Type: G2938C

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:



Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
 Modified: 8/26/2020 11:38:04 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
14_ReconPCR_Lib (1:5)	NovaSeq228	<input type="checkbox"/>	✓			
15_ReconPCR_Lib (1:5)	NovaSeq228	<input type="checkbox"/>	✓			
16_ReconPCR_Lib (1:2)	NovaSeq228	<input type="checkbox"/>	✓			
17_ReconPCR_Lib (1:5)	NovaSeq228	<input type="checkbox"/>	✓			
18_ReconPCR_Lib (1:2)	NovaSeq228	<input type="checkbox"/>	✓			
19_ReconPCR_Lib (1:2)	NovaSeq228	<input type="checkbox"/>	✓			
20_ReconPCR_Lib (1:5)	NovaSeq228	<input type="checkbox"/>	✓			
21_ReconPCR_Lib (1:2)	NovaSeq228	<input type="checkbox"/>	✓			
22_ReconPCR_Lib (1:5)	NovaSeq228	<input type="checkbox"/>	✓			
23_ReconPCR_Lib (1:2)	NovaSeq228	<input type="checkbox"/>	✓			
24_ReconPCR_Lib (1:5)	NovaSeq228	<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
 Modified: 8/26/2020 11:38:04 AM

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

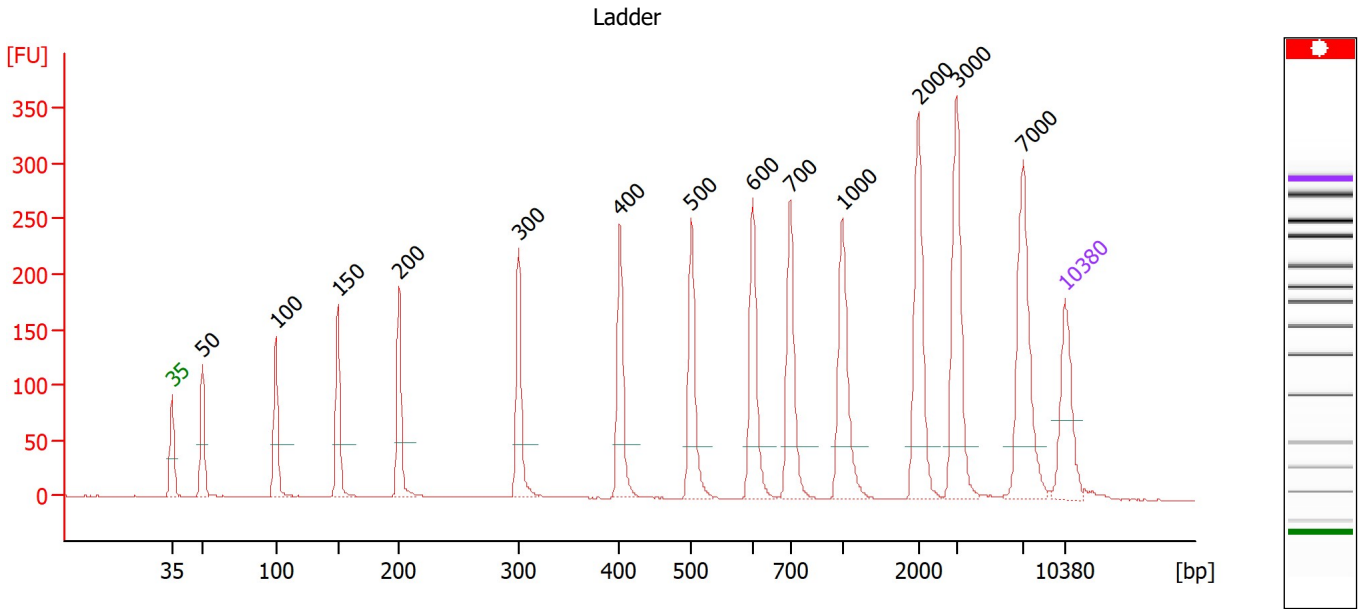
Ladder

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: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
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Electropherogram Summary



Overall Results for Ladder

Noise: 0.4

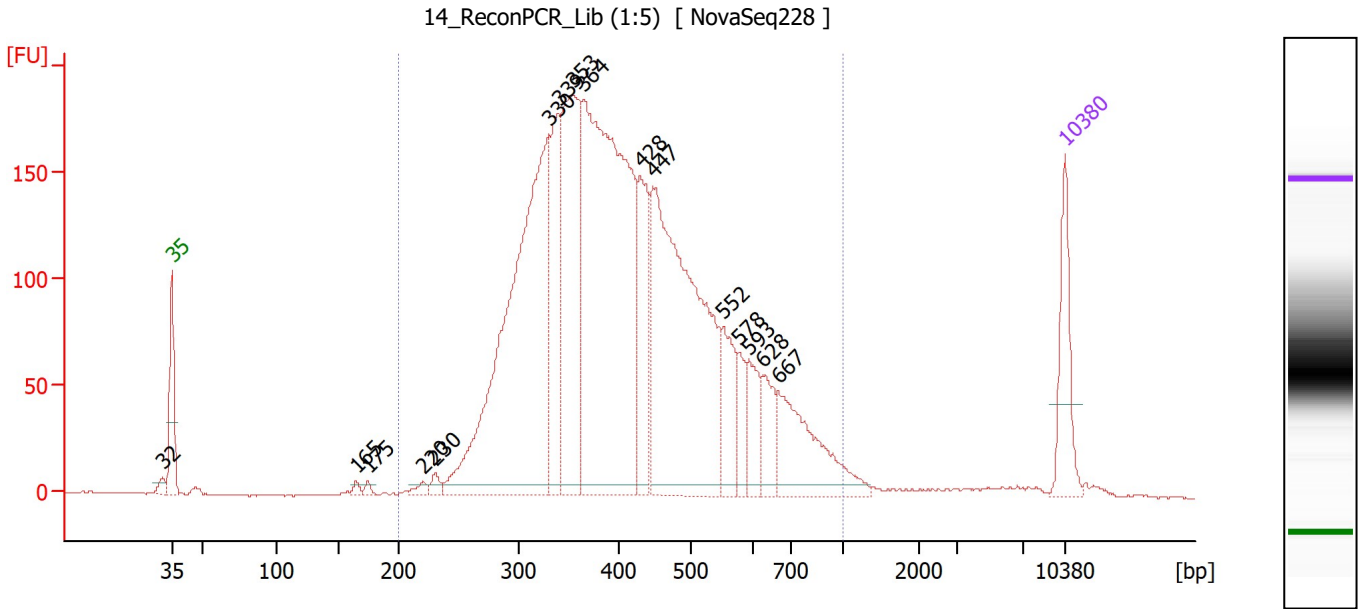
Peak table for Ladder

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: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
 Modified: 8/26/2020 11:38:04 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : 14 ReconPCR Lib (1:5)

Number of peaks found: 16 Corr. Area 1: 4,015.2
 Noise: 0.4

Peak table for sample 1 : 14 ReconPCR Lib (1:5)

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	165	8.09	74.2	
4	175	7.12	61.8	
5	220	11.18	76.9	
6	230	13.56	89.2	
7	330	792.39	3,639.2	
8	339	185.20	828.3	
9	353	347.70	1,490.4	
10	364	881.39	3,667.9	
11	428	154.11	545.1	
12	447	634.63	2,152.5	
13	552	86.19	236.4	
14	578	38.42	100.6	
15	593	61.72	157.7	
16	628	54.18	130.7	
17	667	160.66	364.7	
18	10,380	75.00	10.9	Upper Marker

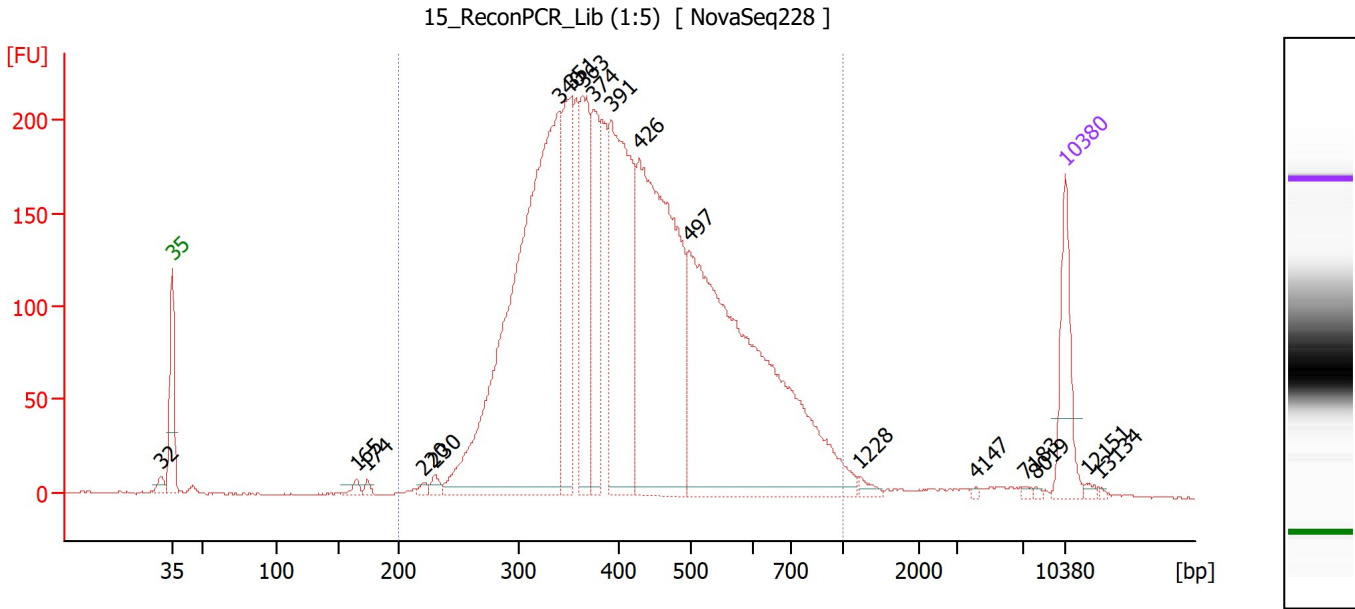
Region table for sample 1 : 14 ReconPCR Lib (1:5)

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	4,015.2	97	426	29.5	3,410.82	13,380.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
 Modified: 8/26/2020 11:38:04 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 : 15 ReconPCR Lib (1:5)

Number of peaks found: 18 Corr. Area 1: 4,785.6
 Noise: 0.5

Peak table for sample 2 : 15 ReconPCR Lib (1:5)

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	165	9.94	91.1	
4	174	6.79	59.1	
5	220	6.90	47.4	
6	230	12.04	79.4	
7	340	982.13	4,378.7	
8	351	230.22	993.0	
9	363	213.84	891.7	
10	374	172.23	698.6	
11	391	404.92	1,569.0	
12	426	607.30	2,157.5	
13	497	837.67	2,552.8	
14	1,228	8.79	10.8	
15	4,147	2.09	0.8	
16	7,183	2.92	0.6	
17	8,019	2.17	0.4	
18	10,380	75.00	10.9	Upper Marker
19	12,151	0.00	0.0	
20	13,134	0.00	0.0	

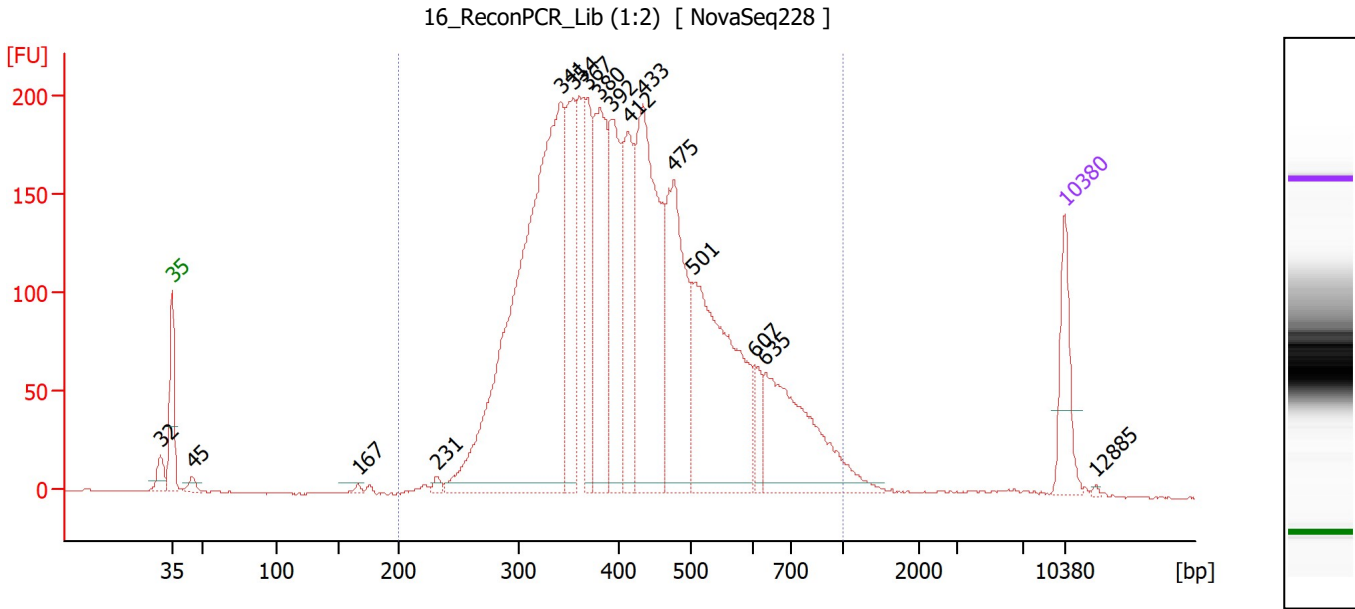
Region table for sample 2 : 15 ReconPCR Lib (1:5)

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	4,785.6	97	433	29.8	3,636.08	14,095.7	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
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Electropherogram Summary Continued ...



Overall Results for sample 3 : 16 ReconPCR Lib (1:2)

Number of peaks found: 16 Corr. Area 1: 4,374.8
 Noise: 0.4

Peak table for sample 3 : 16 ReconPCR Lib (1:2)

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	45	17.17	580.0	
4	167	7.51	68.3	
5	231	11.10	72.7	
6	341	1,152.18	5,112.0	
7	354	217.36	930.9	
8	367	172.10	710.0	
9	380	290.22	1,156.7	
10	392	250.21	967.6	
11	412	196.34	722.4	
12	433	469.49	1,644.2	
13	475	312.22	995.6	
14	501	442.69	1,337.8	
15	607	40.43	101.0	
16	635	263.14	628.3	
17	10,380	75.00	10.9	Upper Marker
18	12,885	0.00	0.0	

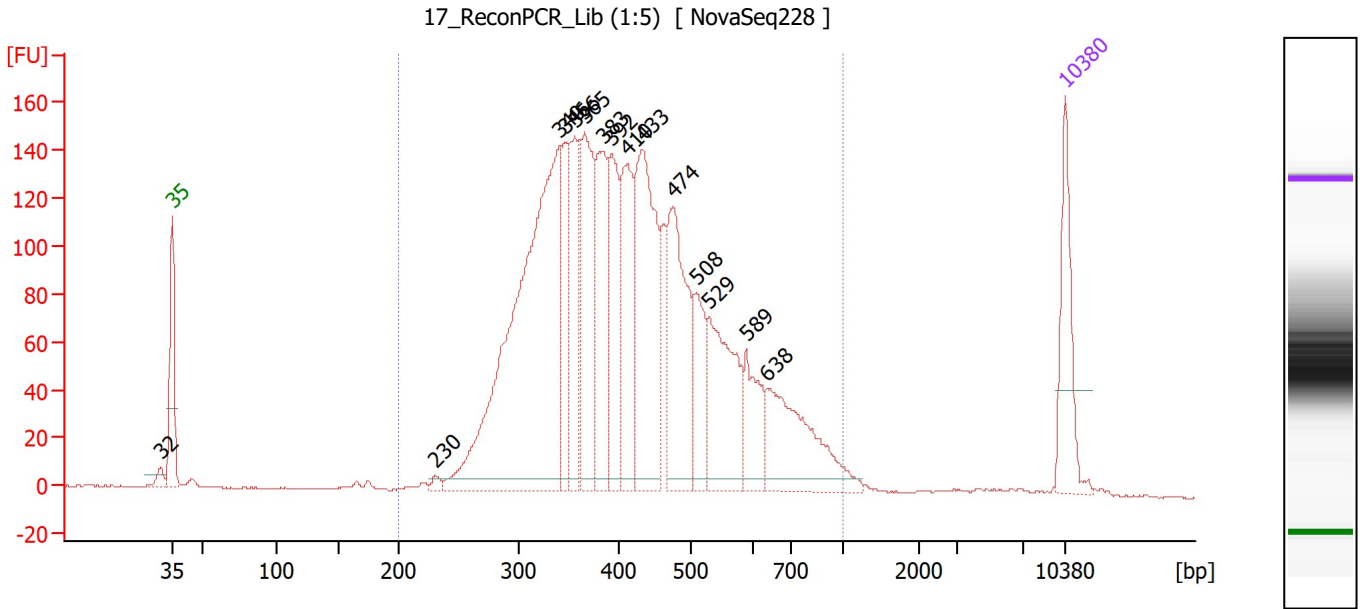
Region table for sample 3 : 16 ReconPCR Lib (1:2)

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	4,374.8	97	432	29.6	4,076.20	15,772.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
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Electropherogram Summary Continued ...



Overall Results for sample 4 : 17 ReconPCR Lib (1:5)

Number of peaks found: 15 Corr. Area 1: 3,222.4
 Noise: 0.3

Peak table for sample 4 : 17 ReconPCR Lib (1:5)

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	230	7.25	47.7	
4	340	668.83	2,979.7	
5	346	106.71	466.8	
6	356	130.09	554.1	
7	365	188.87	783.8	
8	383	154.01	608.9	
9	392	126.17	487.7	
10	410	146.97	543.3	
11	433	258.53	904.9	
12	474	201.52	643.6	
13	508	83.47	248.8	
14	529	155.61	445.9	
15	589	72.01	185.1	
16	638	148.26	352.2	
17	10,380	75.00	10.9	Upper Marker

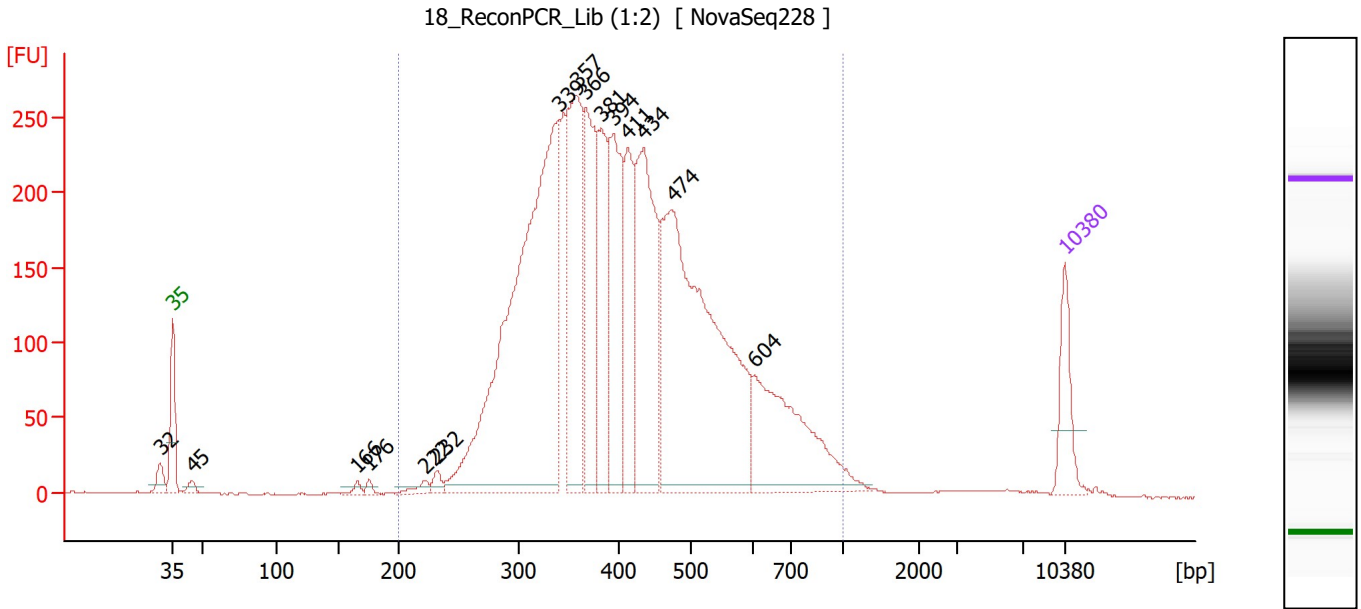
Region table for sample 4 : 17 ReconPCR Lib (1:5)

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	3,222.4	97	431	29.2	2,568.92	9,951.3	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
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Electropherogram Summary Continued ...



Overall Results for sample 5 : 18 ReconPCR Lib (1:2)

Number of peaks found: 15 Corr. Area 1: 5,521.3
 Noise: 0.3

Peak table for sample 5 : 18 ReconPCR Lib (1:2)

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	45	17.13	581.7	
4	166	11.74	107.3	
5	176	9.60	82.8	
6	222	17.82	121.6	
7	232	18.58	121.5	
8	339	1,214.67	5,423.0	
9	357	412.54	1,752.7	
10	366	286.13	1,183.8	
11	381	240.43	956.3	
12	394	288.11	1,108.7	
13	411	202.16	745.2	
14	434	414.68	1,448.8	
15	474	919.53	2,937.0	
16	604	311.94	782.3	
17	10,380	75.00	10.9	Upper Marker

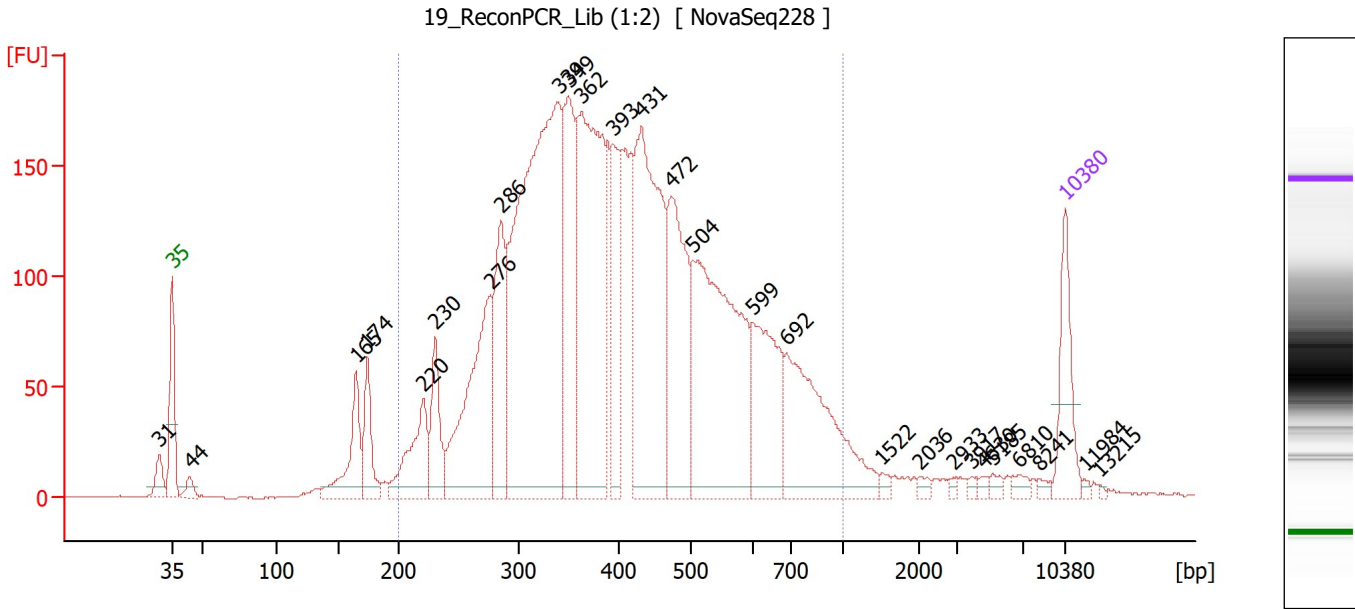
Region table for sample 5 : 18 ReconPCR Lib (1:2)

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	5,521.3	97	427	29.4	4,608.05	18,038.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
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Electropherogram Summary Continued ...



Overall Results for sample 6 : 19 ReconPCR Lib (1:2)

Number of peaks found: 27 Corr. Area 1: 4,603.9
 Noise: 0.4

Peak table for sample 6 : 19 ReconPCR Lib (1:2)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	31	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	44	22.84	788.9	
4	165	121.00	1,111.4	
5	174	89.18	776.2	
6	220	133.98	921.3	
7	230	99.10	652.0	
8	276	318.73	1,748.4	
9	286	200.01	1,061.1	
10	339	935.56	4,178.1	
11	349	248.92	1,080.4	
12	362	513.25	2,147.1	
13	393	161.42	622.4	
14	431	464.56	1,633.9	
15	472	262.40	841.5	
16	504	480.52	1,445.9	
17	599	187.62	474.4	
18	692	270.69	593.1	
19	1,522	7.38	7.3	
20	2,036	6.85	5.1	
21	2,933	4.03	2.1	
22	3,917	5.04	1.9	
23	4,630	5.19	1.7	
24	5,185	7.41	2.2	
25	6,810	9.57	2.1	
26	8,241	5.64	1.0	

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
 Modified: 8/26/2020 11:38:04 AM

Electropherogram Summary Continued ...

... Peak table for sample 6 : 19 ReconPCR Lib (1:2)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
27	10,380	75.00	10.9	Upper Marker
28	11,984	0.00	0.0	
29	13,215	0.00	0.0	

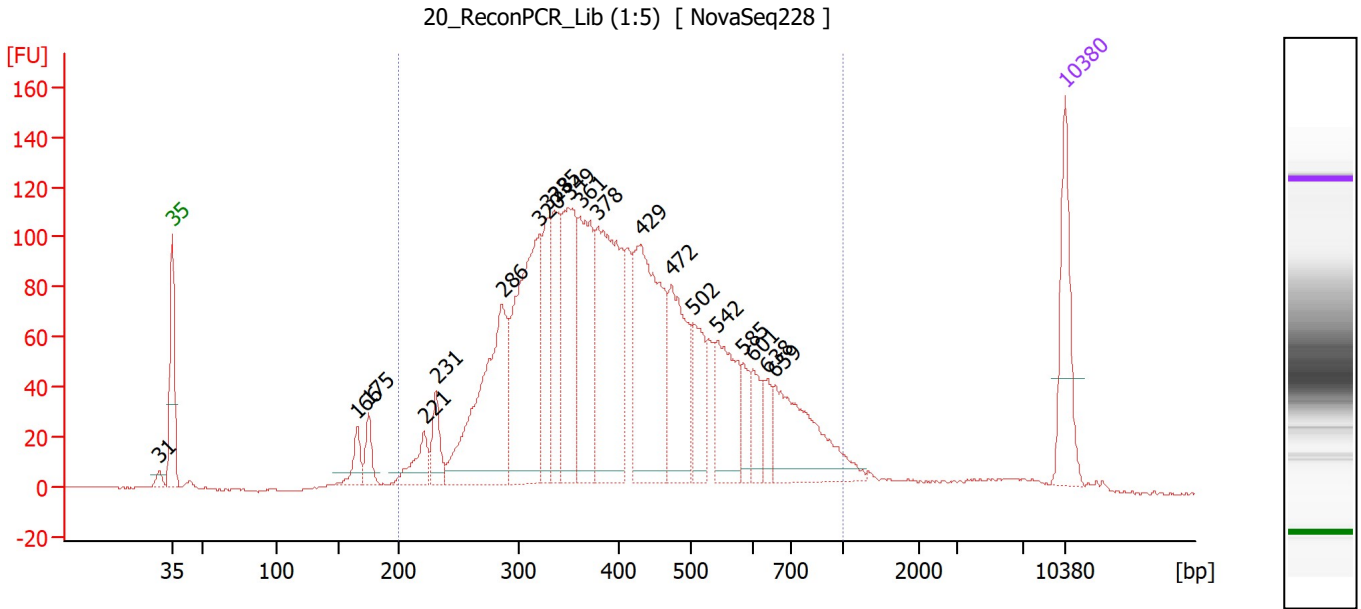
Region table for sample 6 : 19 ReconPCR Lib (1:2)

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	4,603.9	91	428	34.1	4,468.81	18,161.9	■

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
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Electropherogram Summary Continued ...



Overall Results for sample 7 : 20 ReconPCR Lib (1:5)

Number of peaks found: 20 Corr. Area 1: 2,783.1
 Noise: 0.4

Peak table for sample 7 : 20 ReconPCR Lib (1:5)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	31	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	166	34.02	310.9	
4	175	31.82	275.5	
5	221	46.19	317.0	
6	231	43.76	287.2	
7	286	270.98	1,436.8	
8	320	282.06	1,334.3	
9	328	100.67	464.7	
10	335	118.33	535.4	
11	349	180.16	782.6	
12	361	174.78	734.1	
13	378	273.02	1,094.5	
14	429	258.00	911.5	
15	472	137.78	442.3	
16	502	73.21	220.8	
17	542	106.92	299.1	
18	585	31.14	80.6	
19	601	36.42	91.9	
20	638	24.68	58.6	
21	659	139.45	320.6	
22	10,380	75.00	10.9	Upper Marker

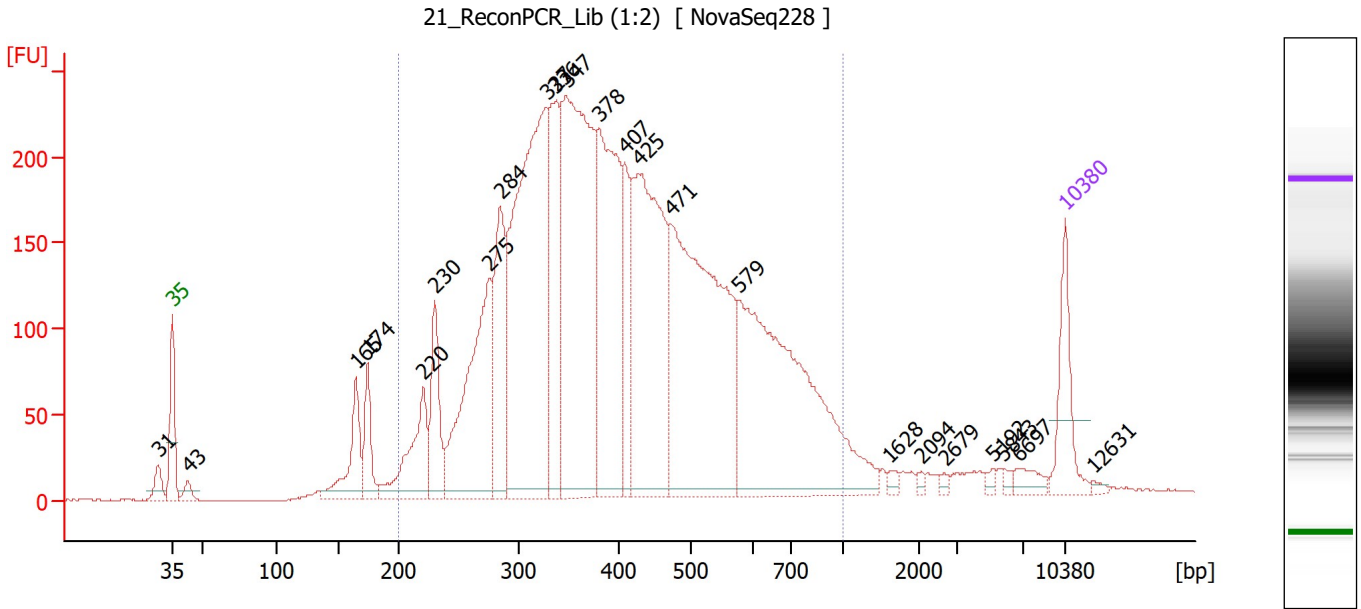
Region table for sample 7 : 20 ReconPCR Lib (1:5)

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	2,783.1	92	428	33.3	2,522.75	10,183.1	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
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Electropherogram Summary Continued ...



Overall Results for sample 8 : 21_ReconPCR Lib (1:2)

Number of peaks found: 23 Corr. Area 1: 5,993.0
 Noise: 0.4

Peak table for sample 8 : 21_ReconPCR Lib (1:2)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	31	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	43	20.97	743.0	
4	165	118.07	1,083.8	
5	174	79.75	692.6	
6	220	152.65	1,050.0	
7	230	120.18	792.0	
8	275	364.15	2,003.9	
9	284	219.76	1,170.7	
10	327	751.26	3,478.1	
11	336	221.21	997.5	
12	347	699.94	3,058.4	
13	378	435.09	1,742.4	
14	407	128.76	478.9	
15	425	505.13	1,798.8	
16	471	663.89	2,135.1	
17	579	584.67	1,529.9	
18	1,628	7.06	6.6	
19	2,094	5.31	3.8	
20	2,679	4.81	2.7	
21	5,192	5.86	1.7	
22	5,843	6.49	1.7	
23	6,697	18.50	4.2	
24	10,380	75.00	10.9	Upper Marker
25	12,631	0.00	0.0	

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
 Modified: 8/26/2020 11:38:04 AM

Electropherogram Summary Continued ...

... Region table for sample 8 :

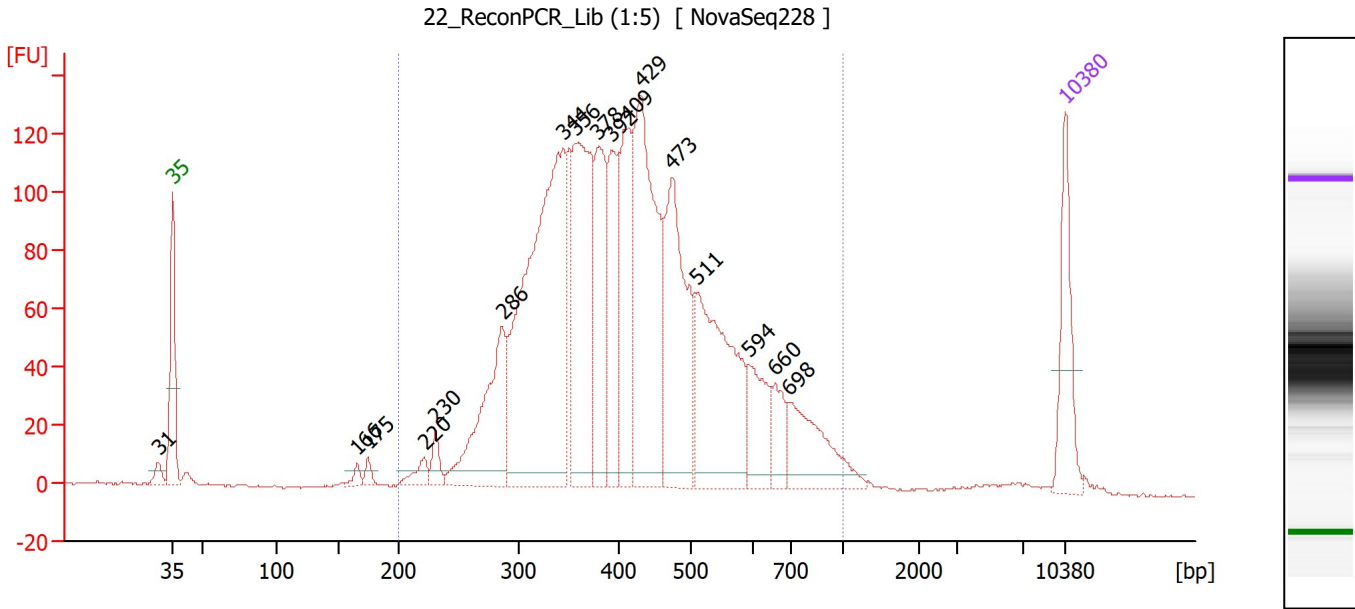
21 ReconPCR Lib (1:2)

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	5,993.0	90	427	34.7	4,767.94	19,534.8	■

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
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Electropherogram Summary Continued ...



Overall Results for sample 9 : 22_ReconPCR Lib (1:5)

Number of peaks found: 17 Corr. Area 1: 2,677.2
 Noise: 0.4

Peak table for sample 9 : 22_ReconPCR Lib (1:5)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	31	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	166	11.41	104.2	
4	175	10.93	94.8	
5	220	21.98	151.1	
6	230	24.07	158.4	
7	286	190.69	1,010.9	
8	344	568.40	2,501.8	
9	356	286.39	1,219.7	
10	378	154.78	620.1	
11	392	134.21	519.1	
12	409	154.28	571.0	
13	429	342.12	1,209.4	
14	473	239.67	768.1	
15	511	256.81	761.9	
16	594	73.32	187.0	
17	660	40.04	92.0	
18	698	104.87	227.7	
19	10,380	75.00	10.9	Upper Marker

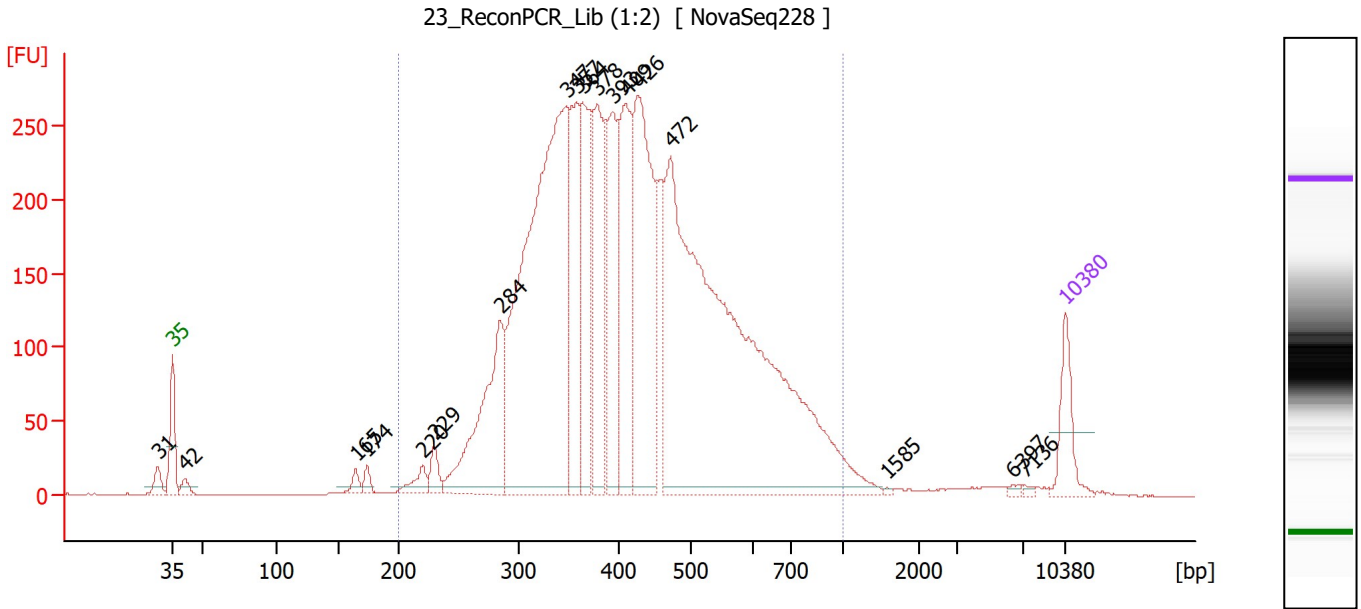
Region table for sample 9 : 22_ReconPCR Lib (1:5)

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	2,677.2	96	431	30.0	2,713.35	10,581.8	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
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Electropherogram Summary Continued ...



Overall Results for sample 10 : 23 ReconPCR Lib (1:2)

Number of peaks found: 18 Corr. Area 1: 6,031.9
 Noise: 0.3

Peak table for sample 10 : 23 ReconPCR Lib (1:2)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	31	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	42	25.55	932.0	
4	165	22.54	206.9	
5	174	22.41	195.1	
6	220	41.43	285.9	
7	229	40.38	266.7	
8	284	402.66	2,145.8	
9	347	1,349.25	5,888.7	
10	357	305.79	1,298.7	
11	364	281.89	1,174.7	
12	378	341.16	1,367.6	
13	393	308.75	1,190.2	
14	409	340.45	1,262.6	
15	426	555.10	1,974.4	
16	472	1,804.33	5,798.1	
17	1,585	2.55	2.4	
18	6,397	5.38	1.3	
19	7,136	3.97	0.8	
20	10,380	75.00	10.9	Upper Marker

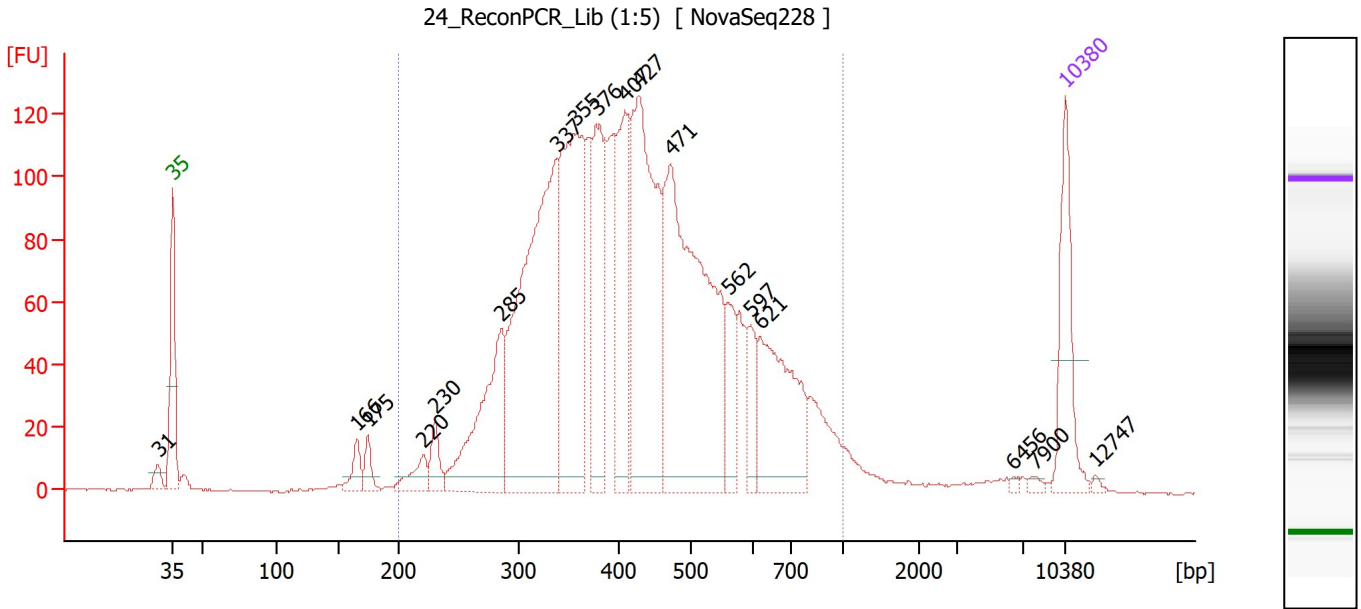
Region table for sample 10 : 23 ReconPCR Lib (1:2)

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	6,031.9	96	434	30.3	5,909.79	22,918.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
 Modified: 8/26/2020 11:38:04 AM

Electropherogram Summary Continued ...



Overall Results for sample 11 : 24 ReconPCR Lib (1:5)

Number of peaks found: 18 Corr. Area 1: 2,742.1
 Noise: 0.3

Peak table for sample 11 : 24 ReconPCR Lib (1:5)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	31	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	166	25.92	237.1	
4	175	22.86	198.2	
5	220	31.66	218.0	
6	230	29.72	195.8	
7	285	181.42	963.5	
8	337	451.38	2,028.8	
9	355	312.60	1,334.0	
10	376	157.41	633.6	
11	407	160.00	596.1	
12	427	323.11	1,146.0	
13	471	439.17	1,412.5	
14	562	49.09	132.4	
15	597	37.71	95.7	
16	621	154.22	376.6	
17	6,456	2.31	0.5	
18	7,900	3.91	0.7	
19	10,380	75.00	10.9	Upper Marker
20	12,747	0.00	0.0	

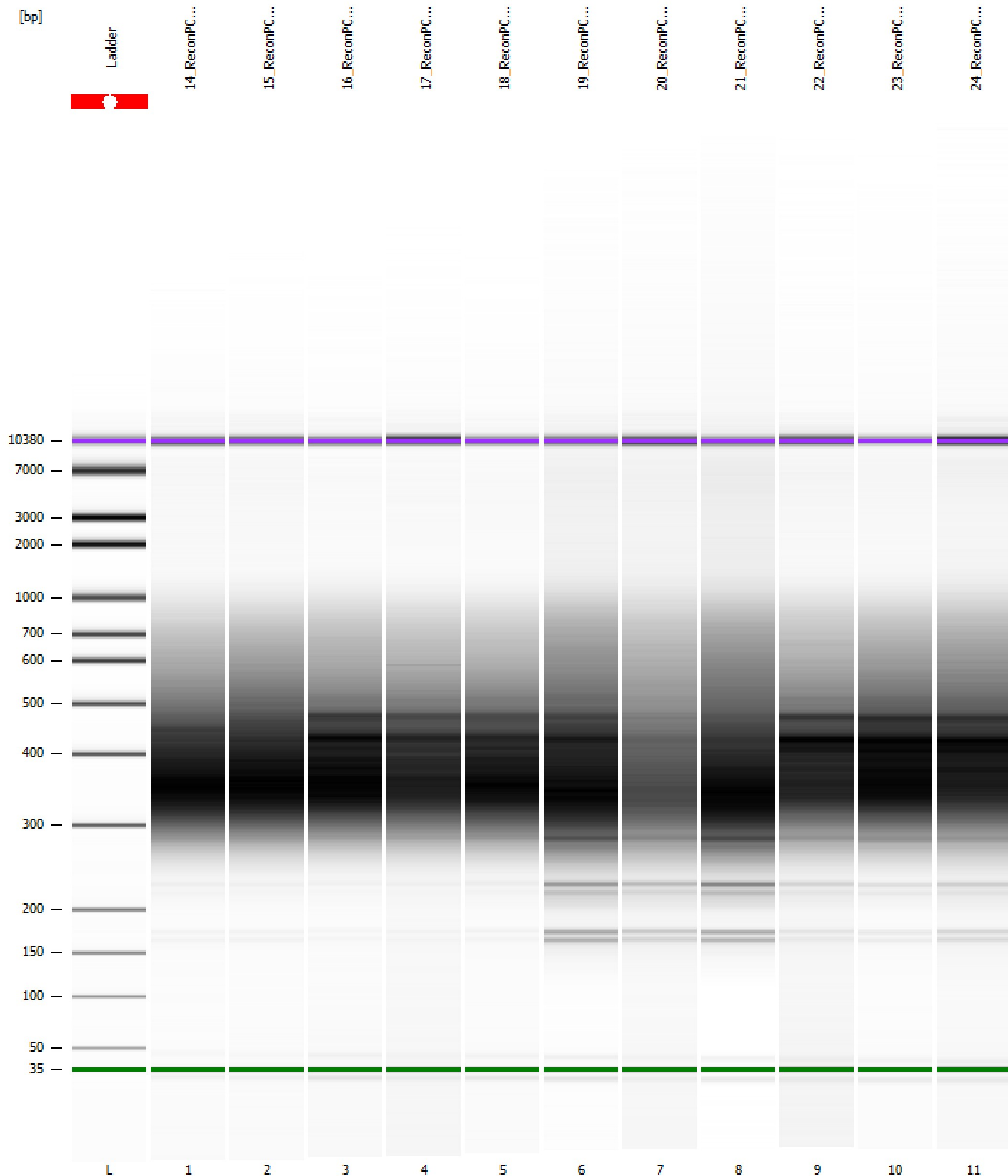
Region table for sample 11 : 24 ReconPCR Lib (1:5)

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	2,742.1	94	441	31.2	2,634.28	10,160.8	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
Modified: 8/26/2020 11:38:04 AM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: Z:\...\Bioanalyzer1_High Sensitivity DNA Assay_2020-08-26_001.xad

Created: 8/26/2020 10:56:36 AM
 Modified: 8/26/2020 11:38:04 AM

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		8/26/2020 11:37:57 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Run started on port 1 (File: Z:\XADs\2020-08-26\Bioanalyze r1_High Sensitivity DNA Assay_2020-08-26_001.xad)		Instrument	Run		8/26/2020 10:56:41 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Product Number : G2938C		Instrument	Run		8/26/2020 10:56:41 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Name :		Instrument	Run		8/26/2020 10:56:41 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Vendor : Agilent Technologies		Instrument	Run		8/26/2020 10:56:41 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Serial# : DE34903152		Instrument	Run		8/26/2020 10:56:41 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Firmware : C.01.069		Instrument	Run		8/26/2020 10:56:41 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB
Cartridge : Electrode		Instrument	Run		8/26/2020 10:56:41 AM	(GMT --07:00) Pacific Standard Time	sbsuser	DESKTOP-4UNV VOB