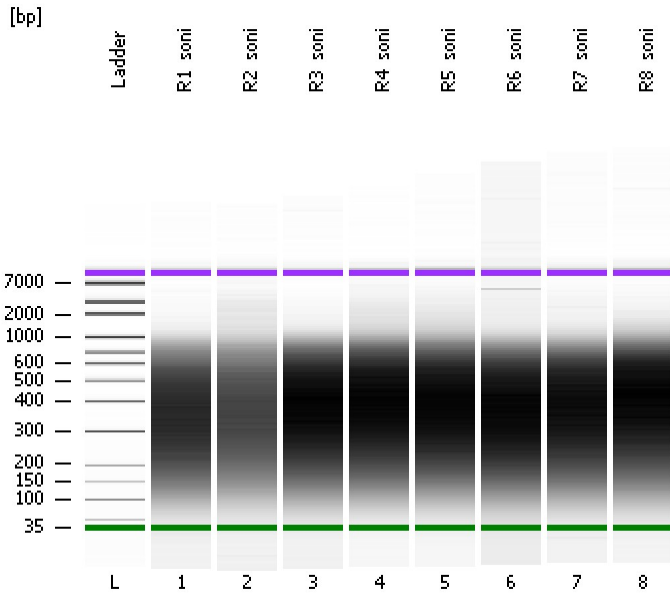


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
Modified: 3/4/2019 3:45:39 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\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

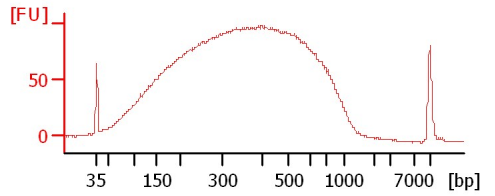
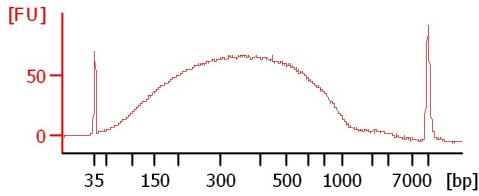
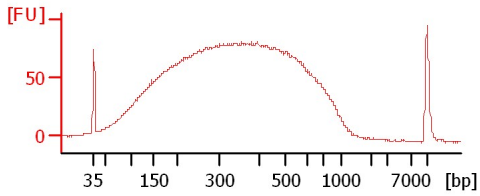
Chip Information:

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

R1_soni

R2_soni

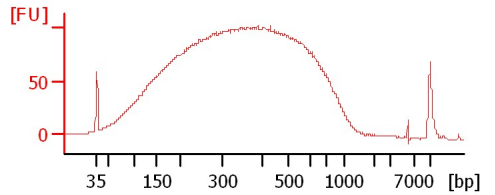
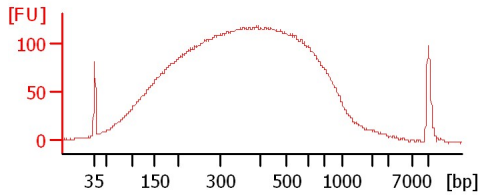
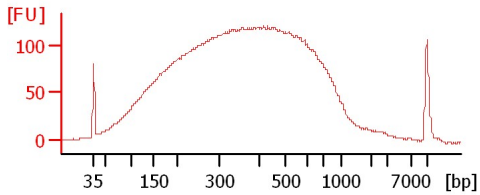
R3_soni



R4_soni

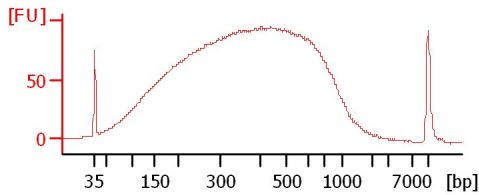
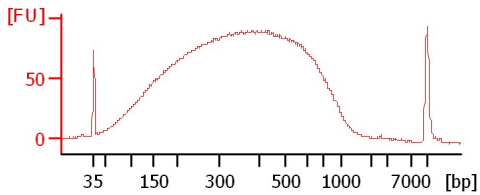
R5_soni

R6_soni



R7_soni

R8_soni



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
R1_soni		<input type="checkbox"/>	✓			
R2_soni		<input type="checkbox"/>	✓			
R3_soni		<input type="checkbox"/>	✓			
R4_soni		<input type="checkbox"/>	✓			
R5_soni		<input type="checkbox"/>	✓			
R6_soni		<input type="checkbox"/>	✓			
R7_soni		<input type="checkbox"/>	✓			
R8_soni		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
Modified: 3/4/2019 3:45:39 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/ μ l] : 1950
Uses Standard Area for Ladder Fragments
Lower Marker Concentration [pg/ μ l] : 125
Upper Marker Concentration [pg/ μ l] : 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

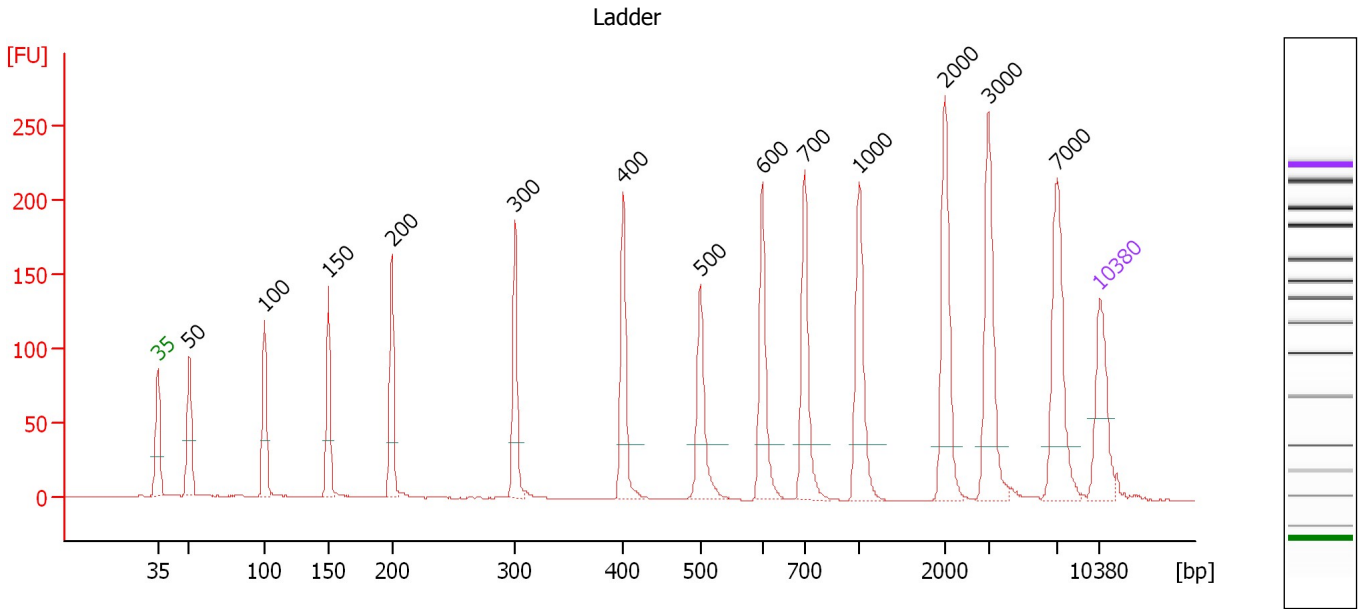
Ladder

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:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.4

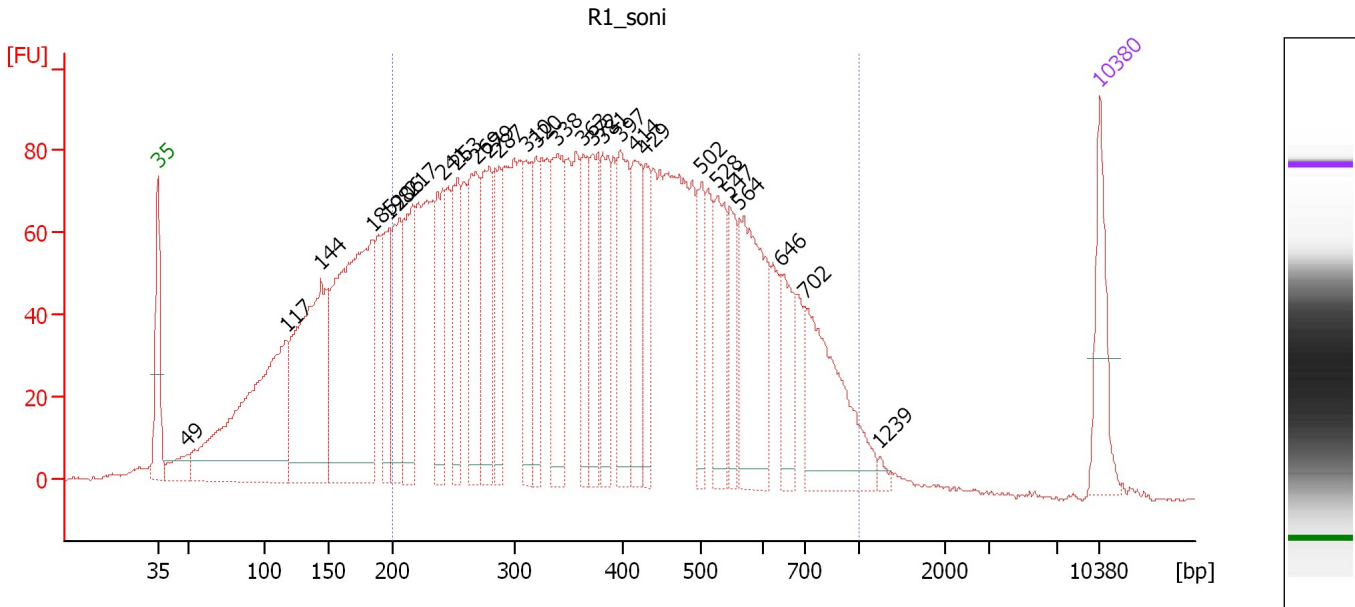
Peak table for Ladder

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	50	150.00	4,545.5	Ladder Peak	45.34
3	100	150.00	2,272.7	Ladder Peak	50.95
4	150	150.00	1,515.2	Ladder Peak	55.67
5	200	150.00	1,136.4	Ladder Peak	60.39
6	300	150.00	757.6	Ladder Peak	69.56
7	400	150.00	568.2	Ladder Peak	77.60
8	500	150.00	454.5	Ladder Peak	83.31
9	600	150.00	378.8	Ladder Peak	87.89
10	700	150.00	324.7	Ladder Peak	91.07
11	1,000	150.00	227.3	Ladder Peak	95.09
12	2,000	150.00	113.6	Ladder Peak	101.50
13	3,000	150.00	75.8	Ladder Peak	104.72
14	7,000	150.00	32.5	Ladder Peak	109.82
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : R1_soni

Number of peaks found: 28 Corr. Area 1: 3,203.2
 Noise: 0.4

Peak table for sample 1 : R1_soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	49	43.81	1,356.5		45.17
3	117	524.84	6,809.1		52.53
4	144	427.66	4,491.4		55.13
5	185	546.56	4,482.4		58.95
6	198	117.83	900.6		60.23
7	206	144.26	1,061.5		60.94
8	217	163.91	1,145.9		61.93
9	241	155.49	977.8		64.15
10	253	130.06	778.0		65.28
11	269	171.53	967.1		66.70
12	279	169.64	921.1		67.64
13	287	117.58	620.1		68.39
14	310	113.89	556.3		70.38
15	320	103.81	492.2		71.13
16	338	185.04	828.7		72.64
17	363	95.04	396.7		74.63
18	372	114.86	468.1		75.33
19	381	119.27	474.8		76.04
20	397	148.38	566.3		77.36
21	414	117.65	430.6		78.40
22	429	89.32	315.6		79.25
23	502	73.37	221.4		83.40
24	528	113.74	326.5		84.58
25	547	72.50	200.7		85.48

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...

... Peak table for sample 1 : R1_soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	564	205.10	551.0		86.24
27	646	74.00	173.6		89.35
28	702	205.85	444.2		91.10
29	1,239	7.05	8.6		96.62
30	10,380	75.00	10.9	Upper Marker	113.00

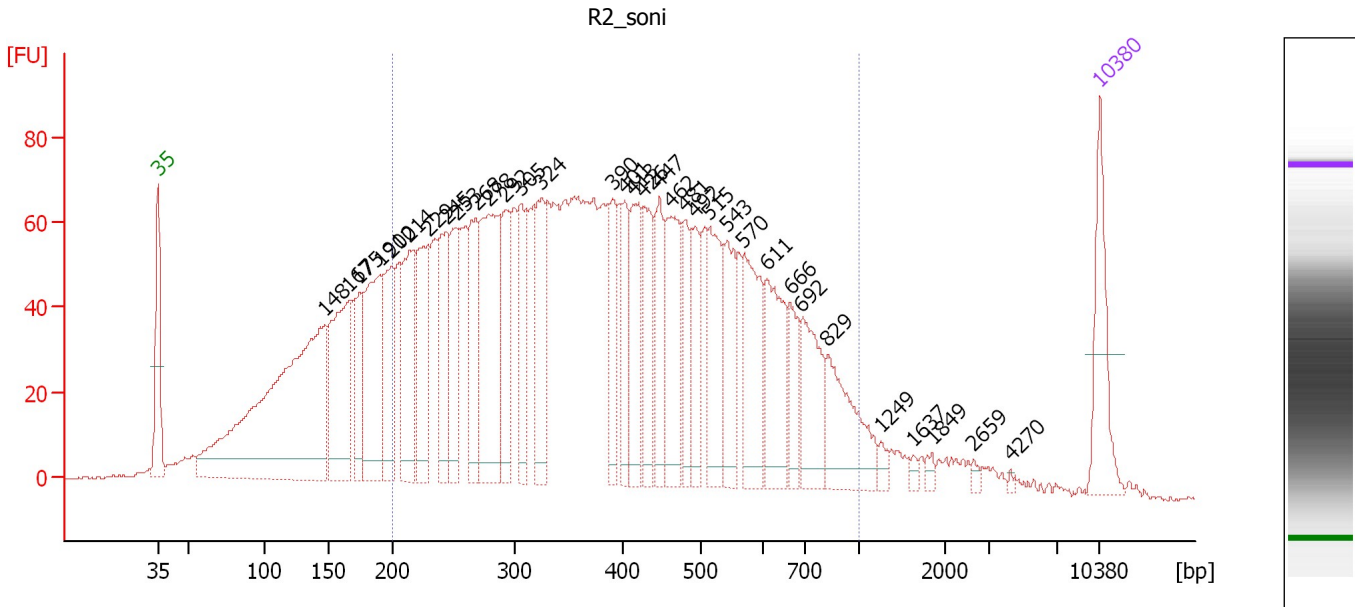
Region table for sample 1 : R1_soni

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	413	4,567.60	3,203.2	20,273.8	75	38.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : R2_soni

Number of peaks found: 34 Corr. Area 1: 2,652.2
 Noise: 0.4

Peak table for sample 2 : R2_soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	148	677.22	6,938.8		55.47
3	167	221.29	2,010.9		57.25
4	175	92.25	800.1		58.00
5	191	200.60	1,594.9		59.50
6	200	124.15	942.9		60.35
7	214	155.57	1,099.8		61.71
8	229	141.12	933.0		63.07
9	245	128.83	798.3		64.47
10	253	100.28	601.3		65.22
11	268	128.14	724.3		66.63
12	278	226.93	1,235.6		67.57
13	292	104.86	544.9		68.79
14	305	89.67	445.5		69.96
15	324	120.65	563.9		71.51
16	390	81.09	315.0		76.80
17	401	72.76	275.1		77.65
18	413	105.20	385.8		78.35
19	426	86.32	306.8		79.10
20	447	84.99	288.2		80.27
21	462	137.38	450.1		81.16
22	481	69.14	218.0		82.20
23	492	71.91	221.4		82.85
24	515	118.03	347.5		83.98
25	543	83.02	231.5		85.29

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...

... Peak table for sample 2 : R2_soni

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	570	124.29	330.4		86.51
27	611	112.57	279.1		88.24
28	666	42.86	97.5		89.98
29	692	91.71	200.7		90.82
30	829	111.32	203.6		92.79
31	1,249	9.70	11.8		96.68
32	1,637	5.21	4.8		99.17
33	1,849	6.92	5.7		100.53
34	2,659	5.11	2.9		103.62
35	4,270	2.34	0.8		106.34
36	10,380	75.00	10.9	Upper Marker	113.00

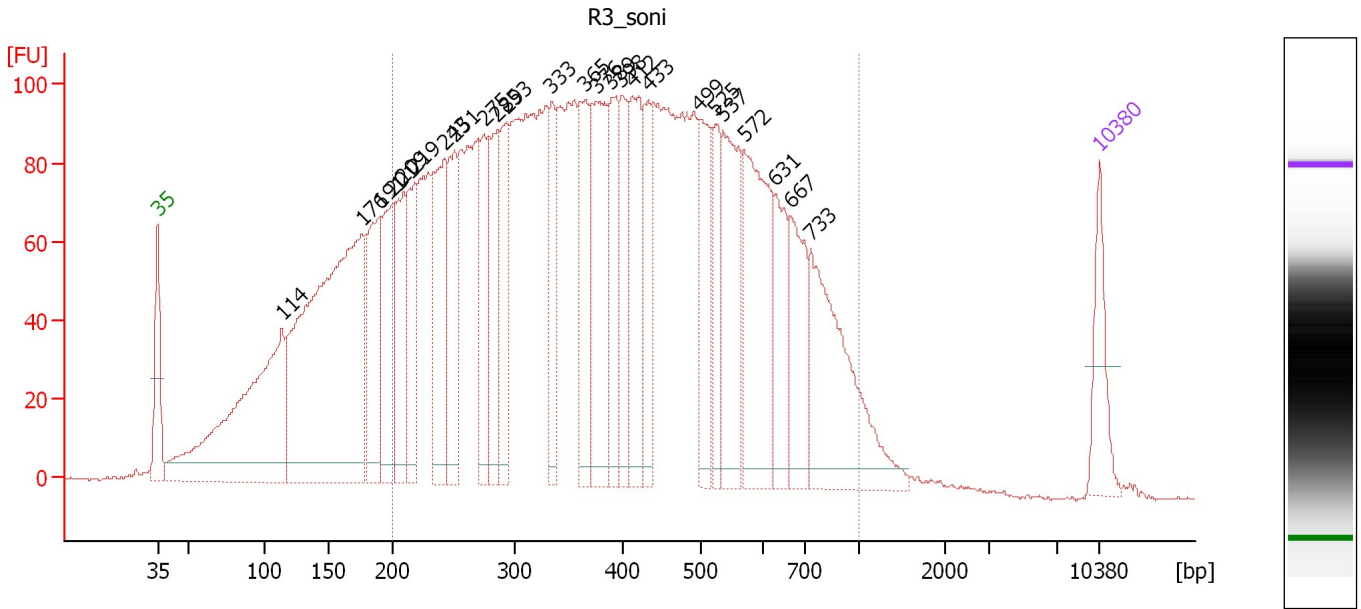
Region table for sample 2 : R2_soni

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	417	3,858.58	2,652.2	17,039.6	75	38.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : R3_soni

Number of peaks found: 25 Corr. Area 1: 3,895.3
 Noise: 0.4

Peak table for sample 3 : R3_soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	114	664.72	8,865.2		52.23
3	176	1,014.00	8,708.6		58.17
4	191	236.37	1,879.0		59.51
5	201	221.52	1,671.8		60.46
6	209	182.99	1,325.9		61.23
7	219	185.02	1,282.9		62.09
8	243	251.47	1,567.7		64.34
9	251	177.53	1,072.2		65.06
10	275	154.84	853.5		67.26
11	285	163.14	866.3		68.22
12	293	152.84	789.9		68.93
13	333	136.59	621.0		72.23
14	365	177.44	737.0		74.77
15	376	268.52	1,081.8		75.68
16	389	160.01	622.9		76.73
17	398	120.43	458.3		77.45
18	412	209.26	770.3		78.26
19	433	142.06	497.6		79.46
20	499	160.54	487.6		83.24
21	525	99.04	286.0		84.44
22	537	229.93	648.6		85.01
23	572	293.33	777.5		86.59
24	631	129.07	309.8		88.89
25	667	146.42	332.4		90.03

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...

... Peak table for sample 3 : R3_soni

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	733	323.91	669.2		91.52
27	10,380	75.00	10.9	Upper Marker	113.00

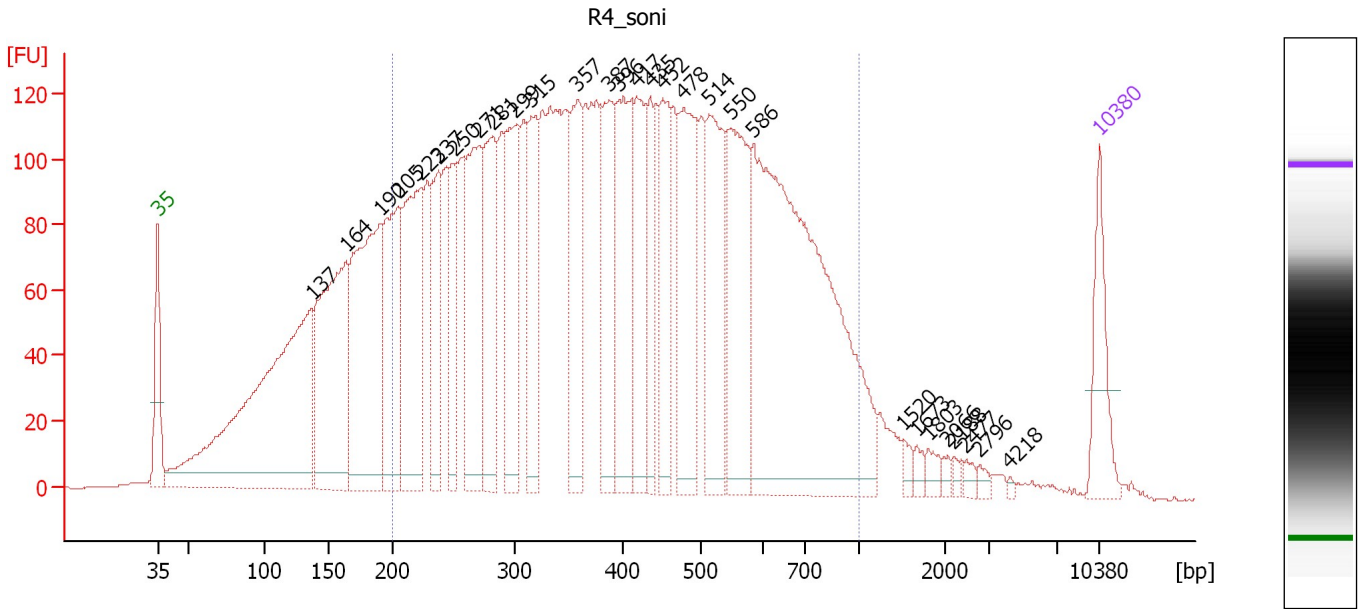
Region table for sample 3 : R3_soni

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	424	6,121.16	3,895.3	26,633.0	 77	38.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : R4_soni

Number of peaks found: 29 Corr. Area 1: 4,750.2
 Noise: 0.4

Peak table for sample 4 : R4_soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	137	826.83	9,148.9		54.44
3	164	436.96	4,025.3		57.04
4	190	454.45	3,615.2		59.49
5	205	269.92	1,993.3		60.87
6	223	323.31	2,193.0		62.54
7	237	151.37	966.5		63.81
8	250	131.71	797.8		64.99
9	271	261.55	1,465.0		66.86
10	281	221.78	1,194.9		67.84
11	299	210.64	1,065.9		69.51
12	315	158.27	760.8		70.78
13	357	185.88	788.1		74.17
14	387	183.80	719.1		76.58
15	396	224.35	857.5		77.31
16	417	160.39	582.4		78.59
17	435	107.76	375.8		79.57
18	452	139.38	467.5		80.55
19	478	234.31	742.1		82.07
20	514	213.34	629.1		83.94
21	550	248.86	685.3		85.61
22	586	807.04	2,088.2		87.23
23	1,520	10.52	10.5		98.42
24	1,673	9.76	8.8		99.40
25	1,803	13.80	11.6		100.24

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...

... Peak table for sample 4 : R4_soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,066	7.80	5.7		101.71
27	2,188	6.05	4.2		102.10
28	2,477	8.29	5.1		103.04
29	2,796	6.76	3.7		104.07
30	4,218	2.82	1.0		106.27
31	10,380	75.00	10.9	Upper Marker	113.00

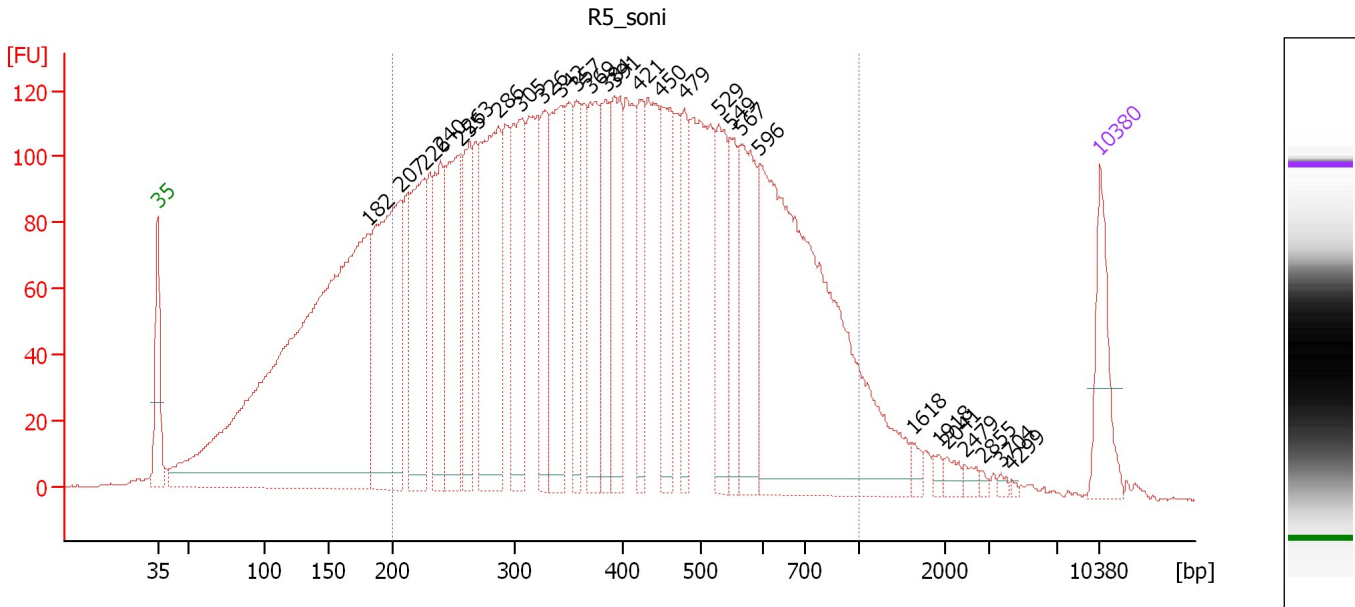
Region table for sample 4 : R4_soni

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	430	5,583.93	4,750.2	24,101.1	76	39.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : R5_soni

Number of peaks found: 28 Corr. Area 1: 4,656.0
 Noise: 0.3

Peak table for sample 5 : R5_soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	182	1,402.15	11,691.1		58.67
3	207	453.73	3,319.9		61.04
4	226	244.68	1,637.7		62.81
5	240	177.01	1,119.3		64.03
6	255	236.89	1,407.2		65.44
7	263	167.17	963.9		66.15
8	286	338.21	1,792.2		68.27
9	305	233.68	1,159.5		69.99
10	326	148.89	691.9		71.66
11	342	202.98	899.8		72.92
12	357	123.60	524.8		74.13
13	369	165.90	680.4		75.14
14	384	134.55	531.0		76.31
15	391	177.17	685.8		76.91
16	421	110.91	399.4		78.78
17	450	147.61	497.1		80.45
18	479	110.63	349.8		82.12
19	529	149.67	428.5		84.65
20	549	87.35	241.0		85.56
21	567	183.76	491.3		86.36
22	596	746.99	1,897.4		87.73
23	1,618	11.88	11.1		99.05
24	1,918	7.05	5.6		100.97
25	2,041	12.15	9.0		101.63

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...

... Peak table for sample 5 : R5_soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,479	8.87	5.4		103.04
27	2,855	3.88	2.1		104.26
28	3,704	4.16	1.7		105.62
29	4,299	2.27	0.8		106.38
30	10,380	75.00	10.9	Upper Marker	113.00

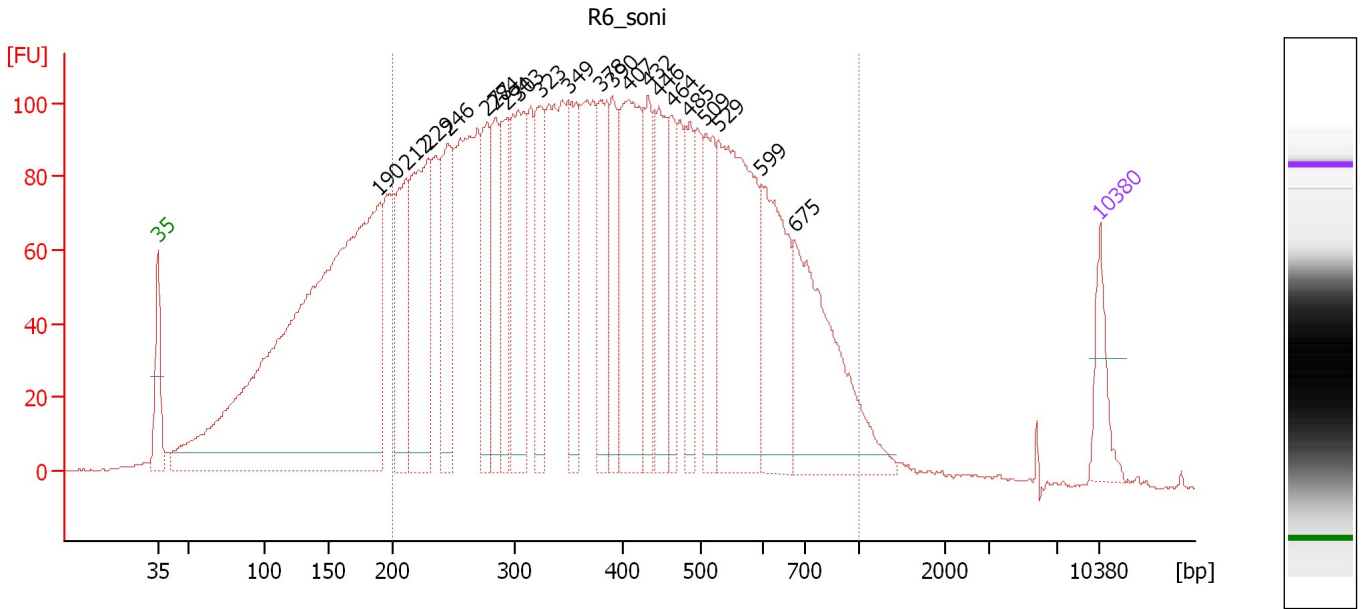
Region table for sample 5 : R5_soni

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	427	5,487.86	4,656.0	23,796.1	76	39.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : R6_soni

Number of peaks found: 21 Corr. Area 1: 3,959.0
 Noise: 0.3

Peak table for sample 6 : R6_soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	190	2,234.31	17,815.4		59.45
3	212	282.43	2,021.1		61.47
4	229	465.78	3,078.7		63.07
5	246	307.36	1,896.2		64.57
6	277	213.22	1,167.7		67.42
7	284	197.95	1,056.1		68.09
8	294	194.29	1,002.7		68.97
9	303	356.53	1,782.9		69.80
10	323	202.01	947.8		71.40
11	349	232.96	1,010.5		73.52
12	378	206.57	827.4		75.85
13	390	187.59	729.1		76.78
14	407	425.24	1,584.9		77.97
15	432	166.69	584.8		79.42
16	446	218.55	741.8		80.25
17	464	145.87	476.7		81.23
18	485	140.00	437.0		82.48
19	509	203.22	605.0		83.72
20	529	572.04	1,637.6		84.65
21	599	319.89	808.8		87.86
22	675	451.89	1,013.7		90.29
23	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...

... Region table for sample 6 :

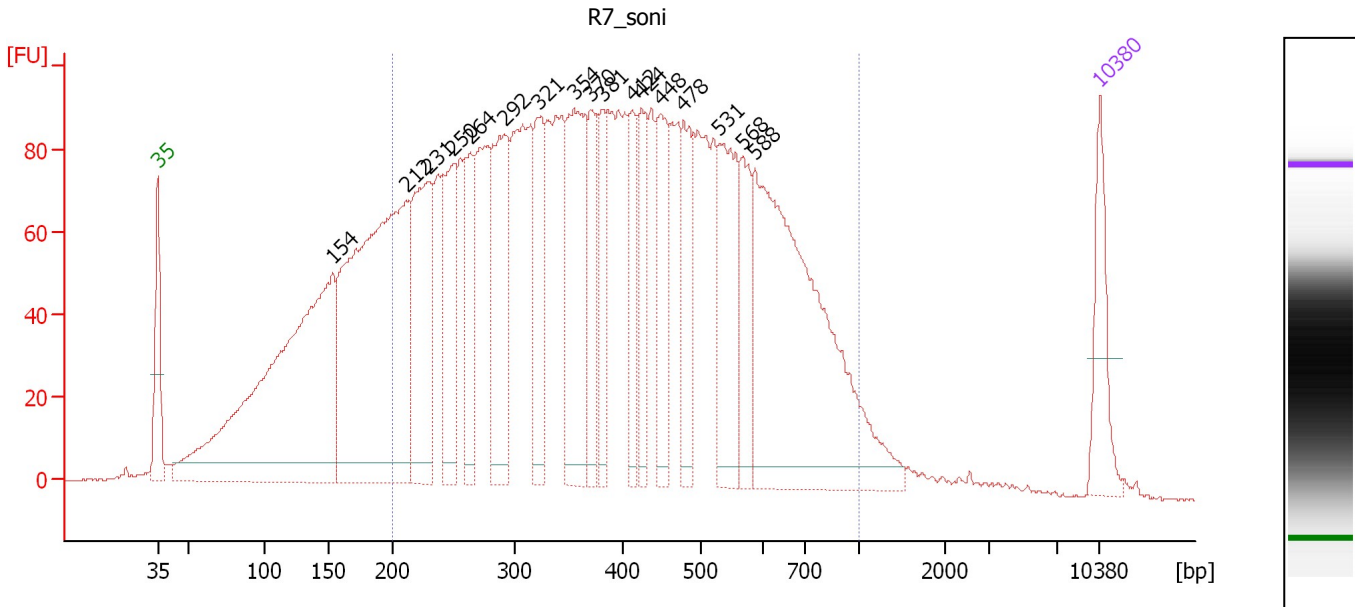
R6_soni

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	416	7,660.41	3,959.0	33,736.5	 76	38.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

Created: 3/4/2019 3:03:00 PM
 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : R7_soni

Number of peaks found: 17 Corr. Area 1: 3,484.8
 Noise: 0.3

Peak table for sample 7 : R7_soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	154	889.09	8,773.8		56.01
3	212	826.02	5,916.6		61.45
4	231	290.09	1,906.3		63.20
5	250	184.15	1,117.8		64.94
6	264	120.65	692.3		66.26
7	292	245.49	1,275.1		68.80
8	321	161.11	759.4		71.29
9	354	253.79	1,085.3		73.93
10	370	114.55	469.0		75.20
11	381	104.46	415.8		76.04
12	412	99.01	364.5		78.26
13	424	98.28	351.5		78.95
14	448	131.41	444.7		80.33
15	478	125.11	396.3		82.07
16	531	184.04	525.4		84.71
17	568	95.03	253.7		86.41
18	588	594.99	1,532.1		87.36
19	10,380	75.00	10.9	Upper Marker	113.00

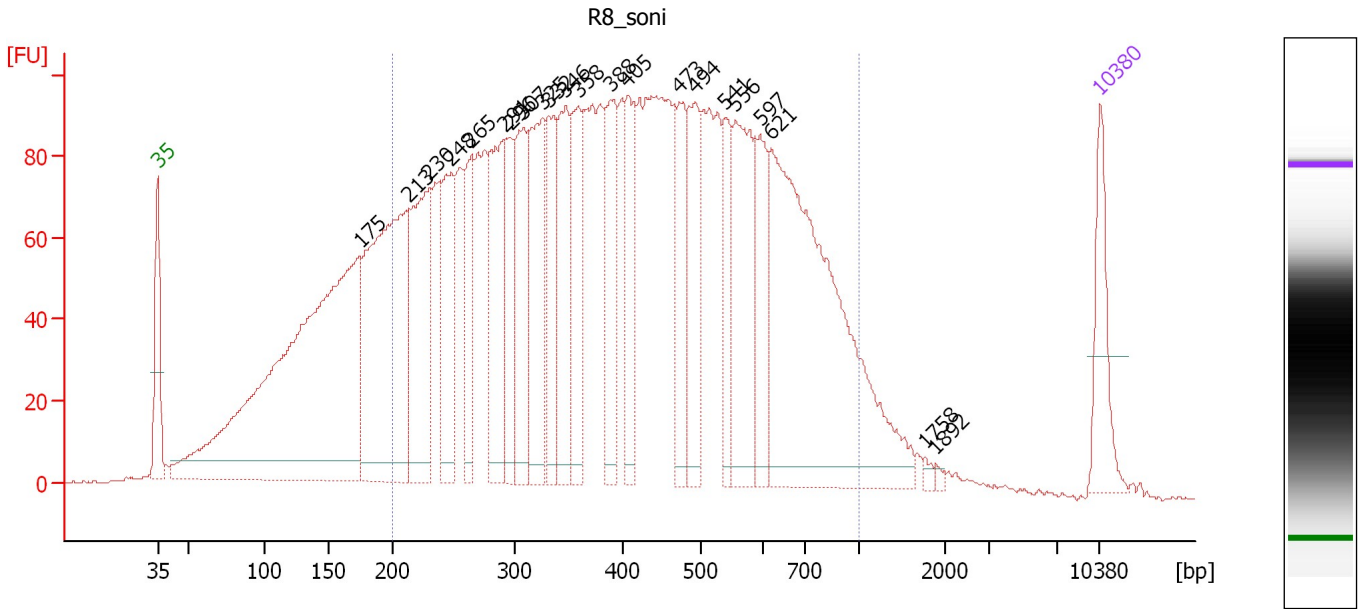
Region table for sample 7 : R7_soni

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	421	4,621.55	3,484.8	20,165.9	77	38.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

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 Modified: 3/4/2019 3:45:39 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : R8_soni

Number of peaks found: 22 Corr. Area 1: 3,686.6
 Noise: 0.2

Peak table for sample 8 : R8_soni

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	35	125.00	5,411.3	Lower Marker	43.00
2	175	1,052.51	9,107.7		58.04
3	213	537.13	3,820.5		61.59
4	230	258.23	1,701.0		63.15
5	248	193.78	1,185.8		64.76
6	265	121.09	693.4		66.32
7	291	191.99	999.7		68.73
8	296	120.33	616.6		69.16
9	307	165.25	815.3		70.13
10	325	197.76	921.6		71.58
11	332	126.50	577.7		72.12
12	346	168.33	736.1		73.30
13	358	147.77	625.7		74.21
14	388	149.11	582.4		76.63
15	405	121.62	455.4		77.87
16	473	121.71	389.6		81.79
17	494	127.17	390.0		82.97
18	541	87.73	245.9		85.17
19	556	200.09	545.3		85.87
20	597	105.69	268.3		87.75
21	621	626.60	1,528.9		88.56
22	1,758	5.51	4.8		99.95
23	1,892	3.46	2.8		100.81
24	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

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Electropherogram Summary Continued ...

... Region table for sample 8 :

R8_soni

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/ μ l]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	435	4,865.76	3,686.6	20,774.5	 78	39.0

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...100 expert\data\2019-03-04\NovaSeq029_Sample1-8_soniDNA.xad

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Gel Image

