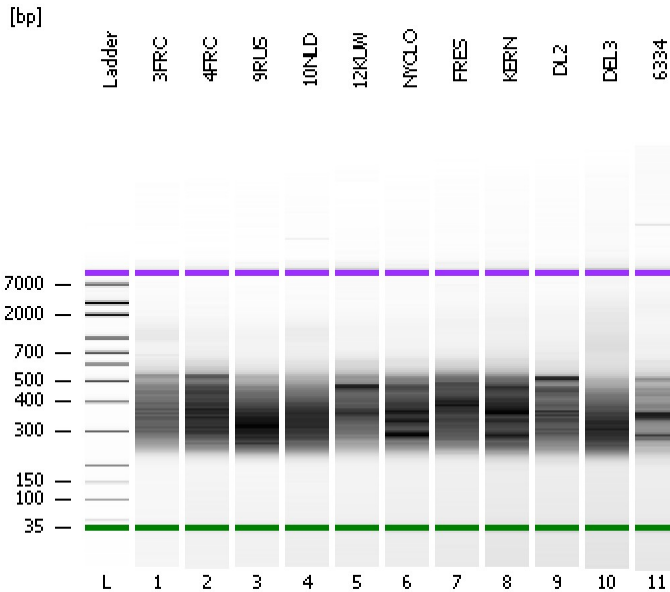


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
Modified: 7/3/2016 3:52:25 PM

Electrophoresis File Run Summary



Instrument Information:

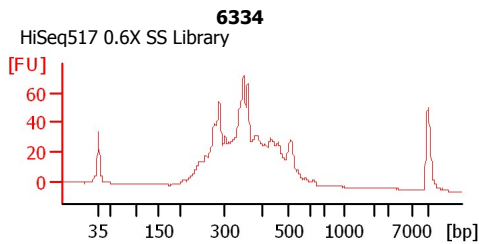
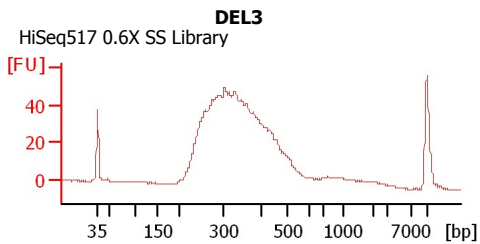
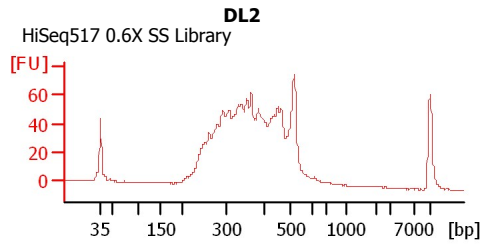
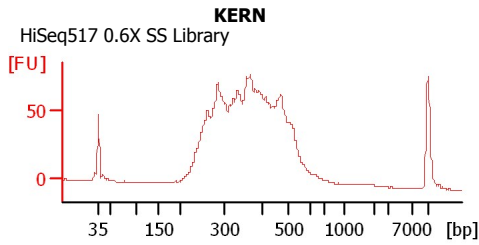
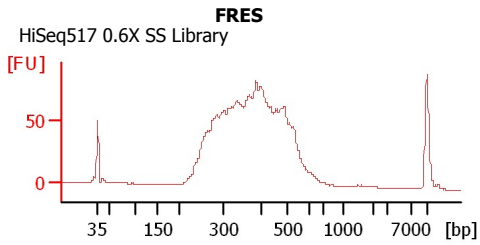
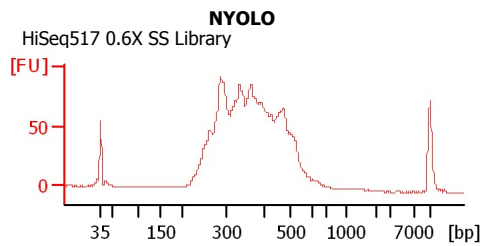
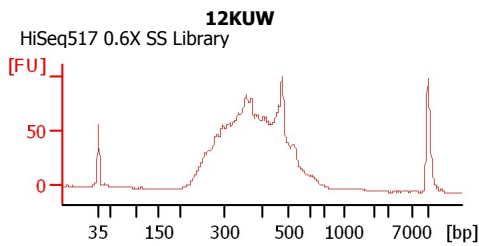
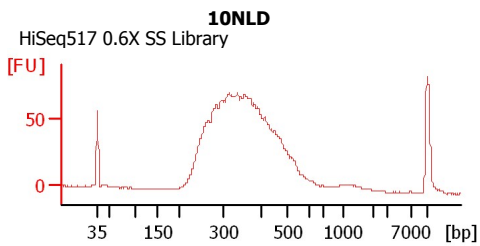
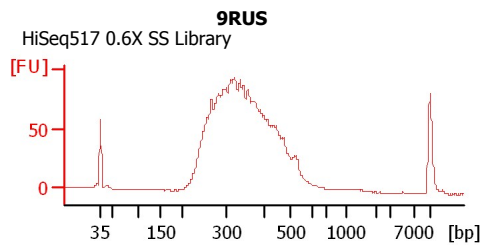
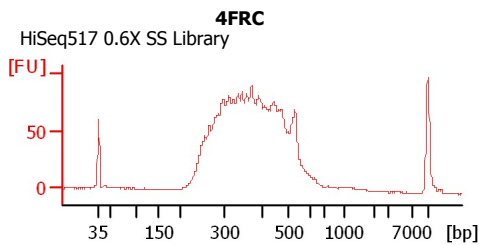
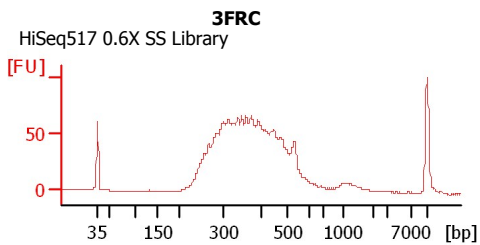
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
Modified: 7/3/2016 3:52:25 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
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4FRC	HiSeq517 0.6X SS Library	<input type="checkbox"/>	✓			
9RUS	HiSeq517 0.6X SS Library	<input type="checkbox"/>	✓			
10NLD	HiSeq517 0.6X SS Library	<input type="checkbox"/>	✓			
12KUW	HiSeq517 0.6X SS Library	<input type="checkbox"/>	✓			
NYOLO	HiSeq517 0.6X SS Library	<input type="checkbox"/>	✓			
FRES	HiSeq517 0.6X SS Library	<input type="checkbox"/>	✓			
KERN	HiSeq517 0.6X SS Library	<input type="checkbox"/>	✓			
DL2	HiSeq517 0.6X SS Library	<input type="checkbox"/>	✓			
DEL3	HiSeq517 0.6X SS Library	<input type="checkbox"/>	✓			
6334	HiSeq517 0.6X SS Library	<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
Modified: 7/3/2016 3:52:25 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

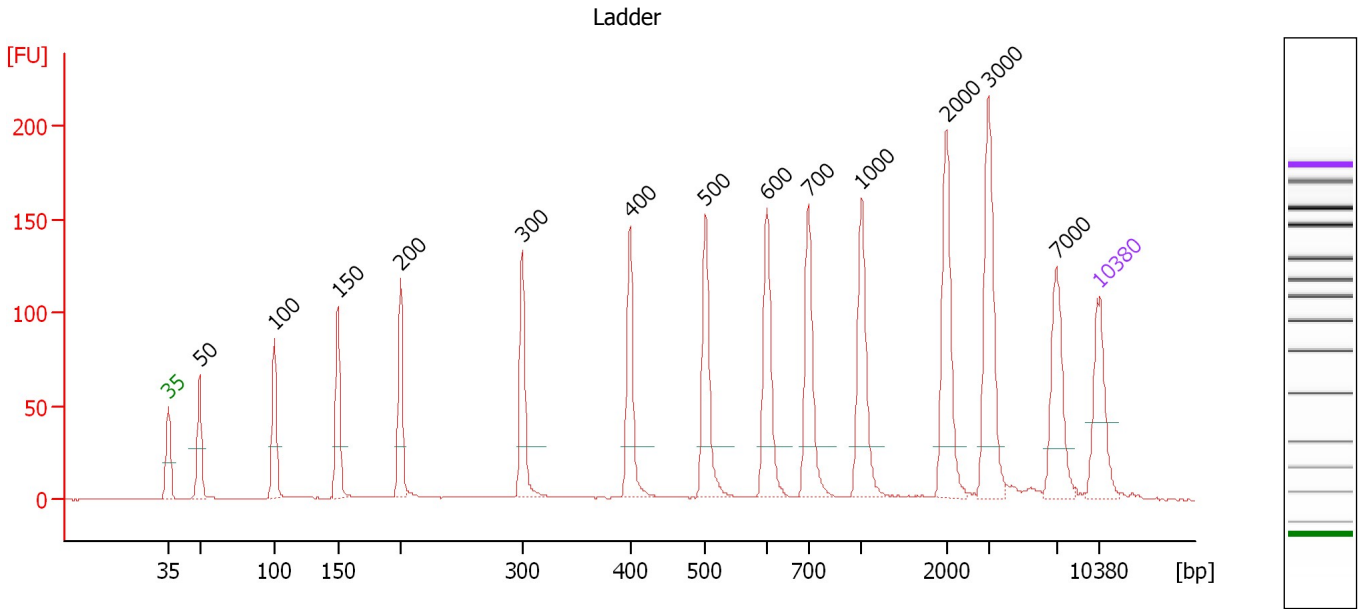
Ladder

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:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
 Modified: 7/3/2016 3:52:25 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

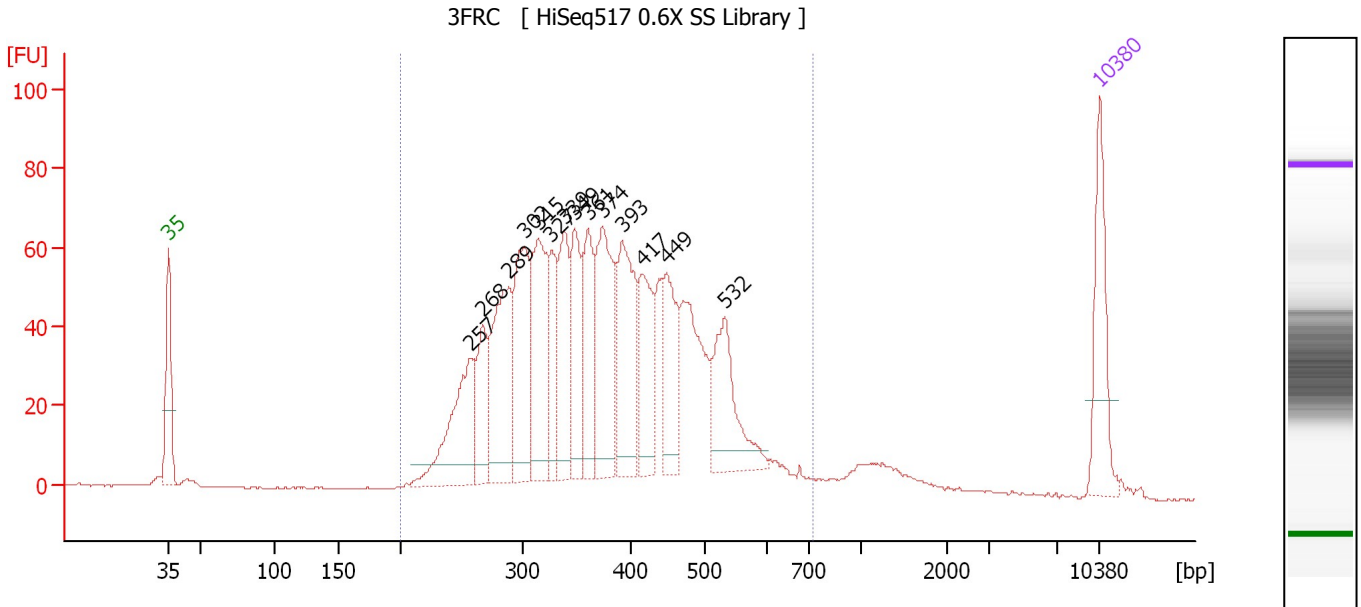
Peak table for Ladder

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	45.35
3	100	150.00	2,272.7	Ladder Peak	50.98
4	150	150.00	1,515.2	Ladder Peak	55.74
5	200	150.00	1,136.4	Ladder Peak	60.44
6	300	150.00	757.6	Ladder Peak	69.60
7	400	150.00	568.2	Ladder Peak	77.68
8	500	150.00	454.5	Ladder Peak	83.36
9	600	150.00	378.8	Ladder Peak	87.97
10	700	150.00	324.7	Ladder Peak	91.10
11	1,000	150.00	227.3	Ladder Peak	95.07
12	2,000	150.00	113.6	Ladder Peak	101.44
13	3,000	150.00	75.8	Ladder Peak	104.62
14	7,000	150.00	32.5	Ladder Peak	109.72
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
 Modified: 7/3/2016 3:52:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : 3FRC

Number of peaks found: 14 Corr. Area 1: 1,535.1
 Noise: 0.2

Peak table for sample 1 : 3FRC

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	257	151.05	888.9		65.70
3	268	82.00	463.9		66.65
4	289	171.51	899.4		68.59
5	302	150.96	756.8		69.78
6	315	158.32	762.7		70.77
7	327	77.58	359.7		71.77
8	339	106.63	476.4		72.76
9	349	98.98	429.8		73.55
10	361	101.23	424.6		74.55
11	374	157.79	639.0		75.59
12	393	146.03	563.6		77.08
13	417	95.21	346.3		78.62
14	449	89.32	301.5		80.46
15	532	108.10	307.9		84.83
16	10,380	75.00	10.9	Upper Marker	113.00

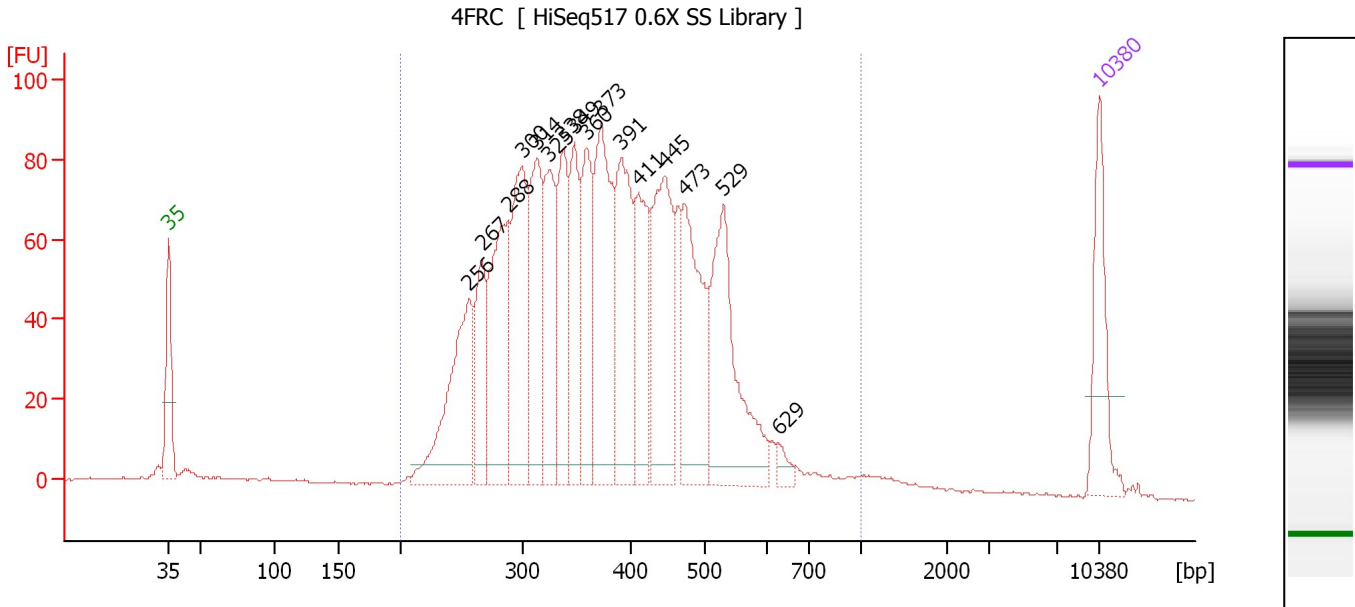
Region table for sample 1 : 3FRC

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	728	381	1,535.1	8,894.3	2,068.57	96	23.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
 Modified: 7/3/2016 3:52:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : 4FRC

Number of peaks found: 16 Corr. Area 1: 2,099.3
 Noise: 0.4

Peak table for sample 2 : 4FRC

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	256	214.46	1,269.2		65.57
3	267	105.90	601.2		66.57
4	288	209.66	1,102.6		68.51
5	300	215.03	1,085.7		69.60
6	314	172.64	832.4		70.75
7	325	144.14	671.3		71.65
8	338	130.59	584.9		72.69
9	349	121.50	527.8		73.54
10	360	125.58	528.7		74.44
11	373	212.14	862.1		75.48
12	391	186.09	720.5		76.98
13	411	127.94	471.3		78.32
14	445	194.71	662.2		80.27
15	473	178.29	571.5		81.81
16	529	217.49	622.9		84.70
17	629	12.21	29.4		88.89
18	10,380	75.00	10.9	Upper Marker	113.00

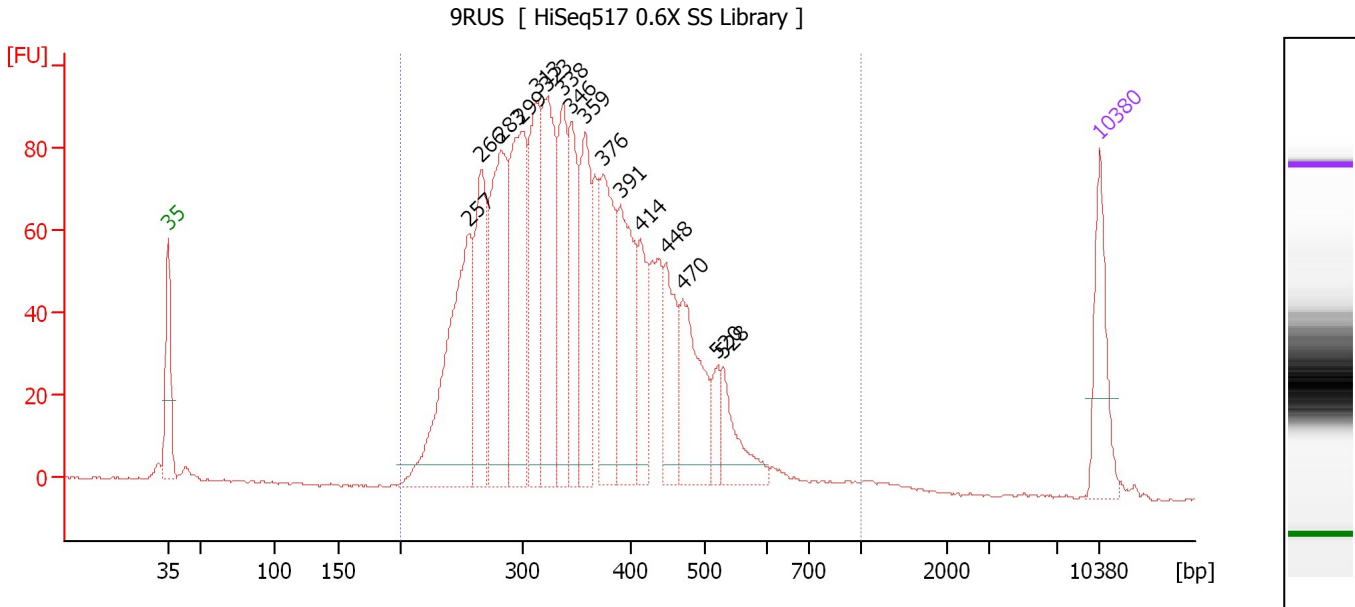
Region table for sample 2 : 4FRC

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	387	2,099.3	11,616.6	2,715.13	98	26.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
 Modified: 7/3/2016 3:52:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : 9RUS

Number of peaks found: 16 Corr. Area 1: 1,994.5
 Noise: 0.3

Peak table for sample 3 : 9RUS

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	257	348.07	2,053.3		65.65
3	266	175.28	999.8		66.45
4	283	284.38	1,524.4		68.01
5	299	268.64	1,360.6		69.52
6	313	188.88	915.1		70.63
7	323	235.96	1,107.9		71.43
8	338	148.37	665.8		72.64
9	346	137.75	603.7		73.29
10	359	159.45	673.3		74.35
11	376	178.09	718.4		75.71
12	391	175.39	679.3		76.97
13	414	76.60	280.3		78.48
14	448	97.23	329.1		80.39
15	470	143.87	464.0		81.65
16	520	31.62	92.2		84.27
17	528	69.55	199.4		84.67
18	10,380	75.00	10.9	Upper Marker	113.00

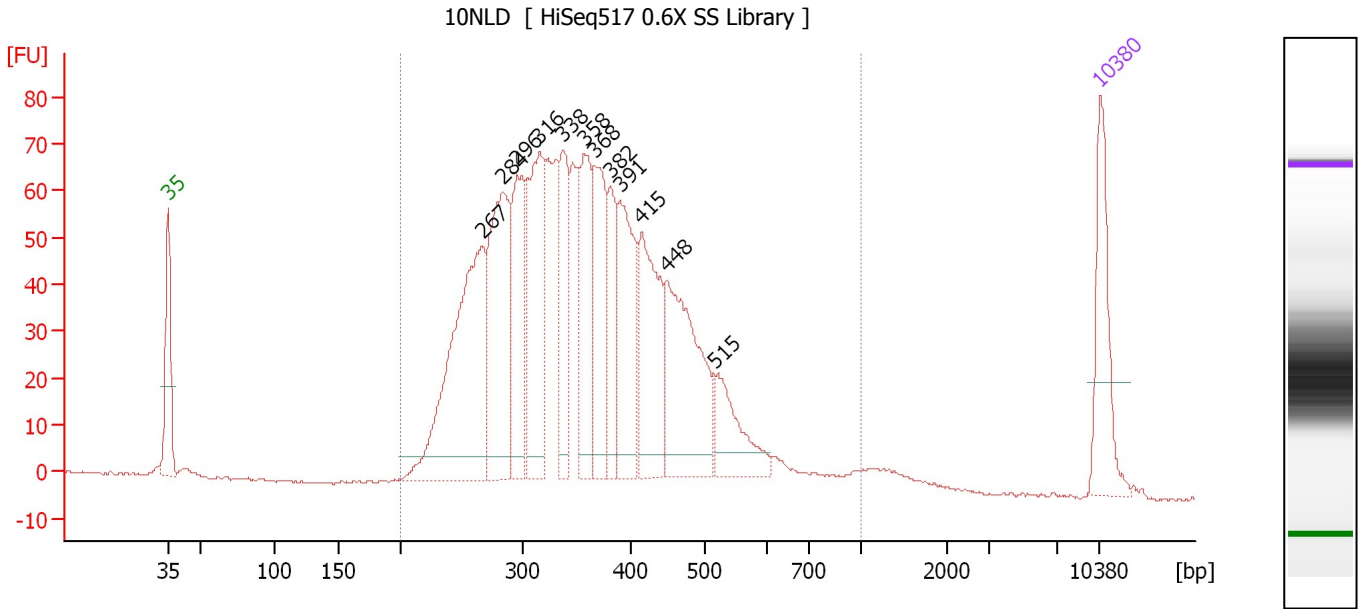
Region table for sample 3 : 9RUS

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	359	1,994.5	13,662.3	2,993.51	98	25.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
 Modified: 7/3/2016 3:52:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : 10NLD

Number of peaks found: 12 Corr. Area 1: 1,595.3
 Noise: 0.3

Peak table for sample 4 : 10NLD

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	267	341.36	1,936.2		66.59
3	284	225.39	1,203.0		68.12
4	296	151.34	774.3		69.24
5	316	188.25	903.2		70.88
6	338	101.24	453.9		72.66
7	358	121.57	515.2		74.25
8	368	125.74	518.2		75.06
9	382	75.90	300.9		76.24
10	391	139.97	542.4		76.95
11	415	153.24	559.4		78.54
12	448	191.52	647.3		80.43
13	515	75.40	221.9		84.05
14	10,380	75.00	10.9	Upper Marker	113.00

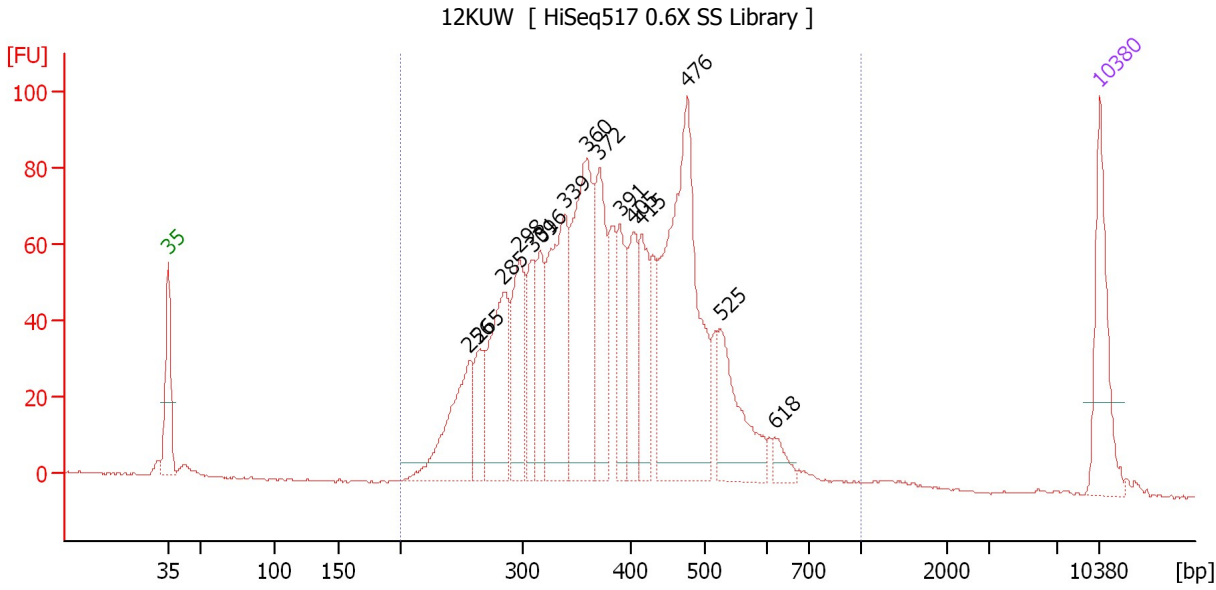
Region table for sample 4 : 10NLD

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	367	1,595.3	9,989.4	2,230.41	98	26.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
 Modified: 7/3/2016 3:52:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : 12KUW

Number of peaks found: 15 Corr. Area 1: 1,703.6
 Noise: 0.3

Peak table for sample 5 : 12KUW

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	256	133.45	788.4		65.61
3	265	55.26	316.0		66.38
4	285	149.56	796.2		68.19
5	298	113.23	576.6		69.37
6	309	73.61	360.6		70.35
7	316	74.67	357.7		70.92
8	339	191.62	855.8		72.77
9	360	225.47	949.9		74.42
10	372	126.75	516.6		75.40
11	391	83.62	324.1		76.94
12	405	71.50	267.4		77.97
13	415	82.35	300.6		78.54
14	476	350.72	1,116.7		81.99
15	525	105.96	305.8		84.52
16	618	14.95	36.6		88.53
17	10,380	75.00	10.9	Upper Marker	113.00

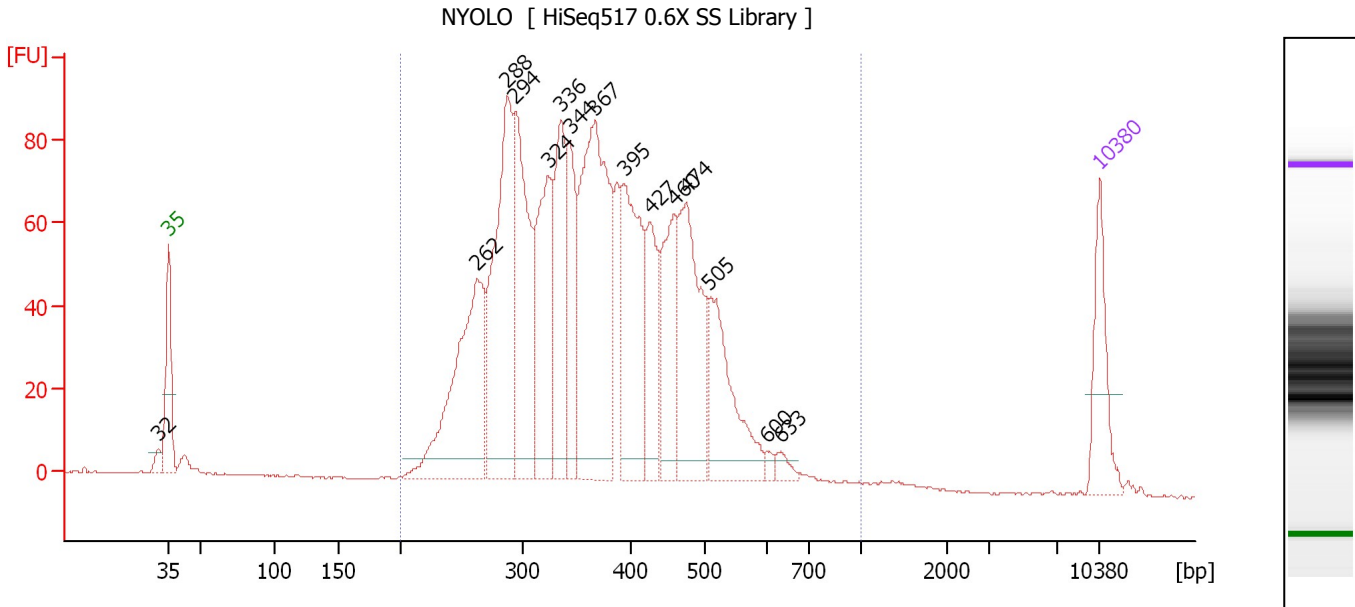
Region table for sample 5 : 12KUW

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	391	1,703.6	8,519.2	2,031.23	99	24.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
 Modified: 7/3/2016 3:52:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : NYOLO

Number of peaks found: 15 Corr. Area 1: 1,880.6
 Noise: 0.2

Peak table for sample 6 : NYOLO

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.27
2	35	125.00	5,411.3	Lower Marker	43.00
3	262	341.53	1,971.6		66.16
4	288	368.91	1,941.0		68.50
5	294	277.41	1,431.4		69.02
6	324	221.56	1,035.2		71.56
7	336	190.52	857.9		72.55
8	344	131.73	579.9		73.17
9	367	439.37	1,812.4		75.04
10	395	229.66	881.1		77.27
11	427	124.11	440.8		79.19
12	460	146.87	484.3		81.06
13	474	226.39	723.4		81.89
14	505	172.98	518.7		83.61
15	600	7.41	18.7		87.97
16	633	13.33	31.9		89.01
17	10,380	75.00	10.9	Upper Marker	113.00

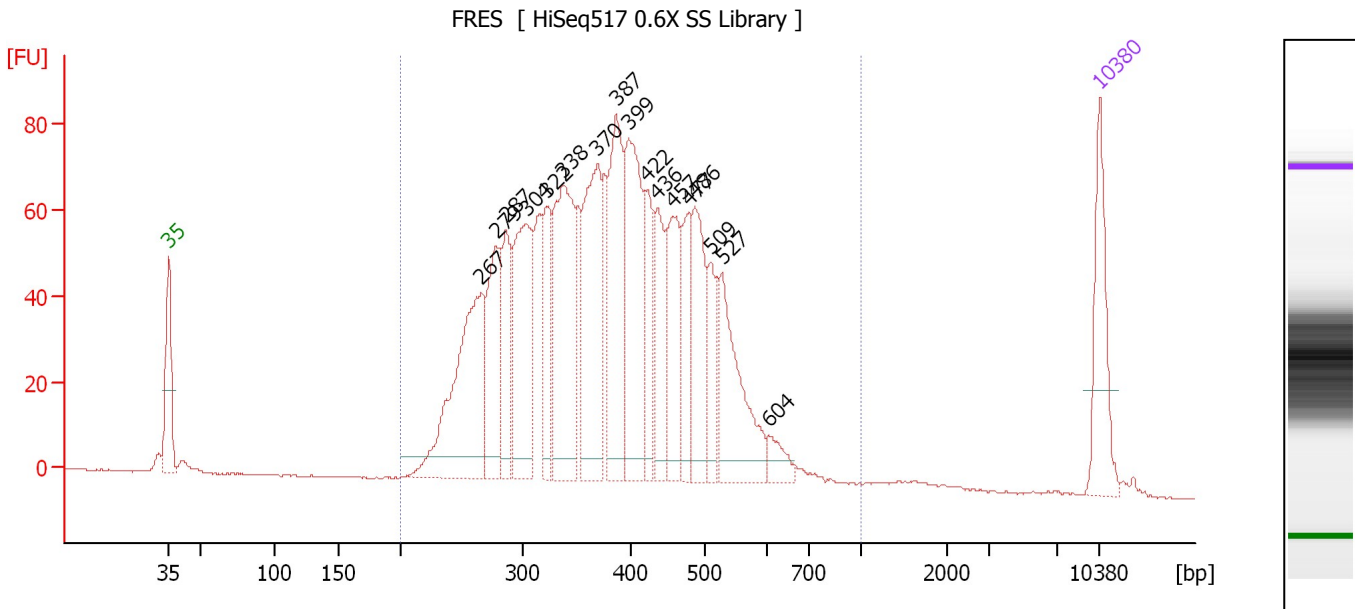
Region table for sample 6 : NYOLO

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	376	1,880.6	13,315.0	3,058.91	98	24.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
 Modified: 7/3/2016 3:52:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : FRES

Number of peaks found: 17 Corr. Area 1: 1,766.6
 Noise: 0.2

Peak table for sample 7 : FRES

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	267	266.31	1,513.2		66.54
3	279	136.55	740.8		67.70
4	287	97.02	512.7		68.38
5	304	187.26	933.9		69.91
6	322	94.02	442.4		71.38
7	338	230.34	1,033.7		72.64
8	370	198.93	814.3		75.27
9	387	173.99	681.1		76.63
10	399	189.57	720.4		77.58
11	422	75.52	271.0		78.95
12	436	90.47	314.3		79.73
13	457	114.13	378.1		80.94
14	477	67.48	214.4		82.05
15	486	118.43	369.2		82.57
16	509	56.94	169.5		83.78
17	527	142.55	409.6		84.62
18	604	22.04	55.3		88.09
19	10,380	75.00	10.9	Upper Marker	113.00

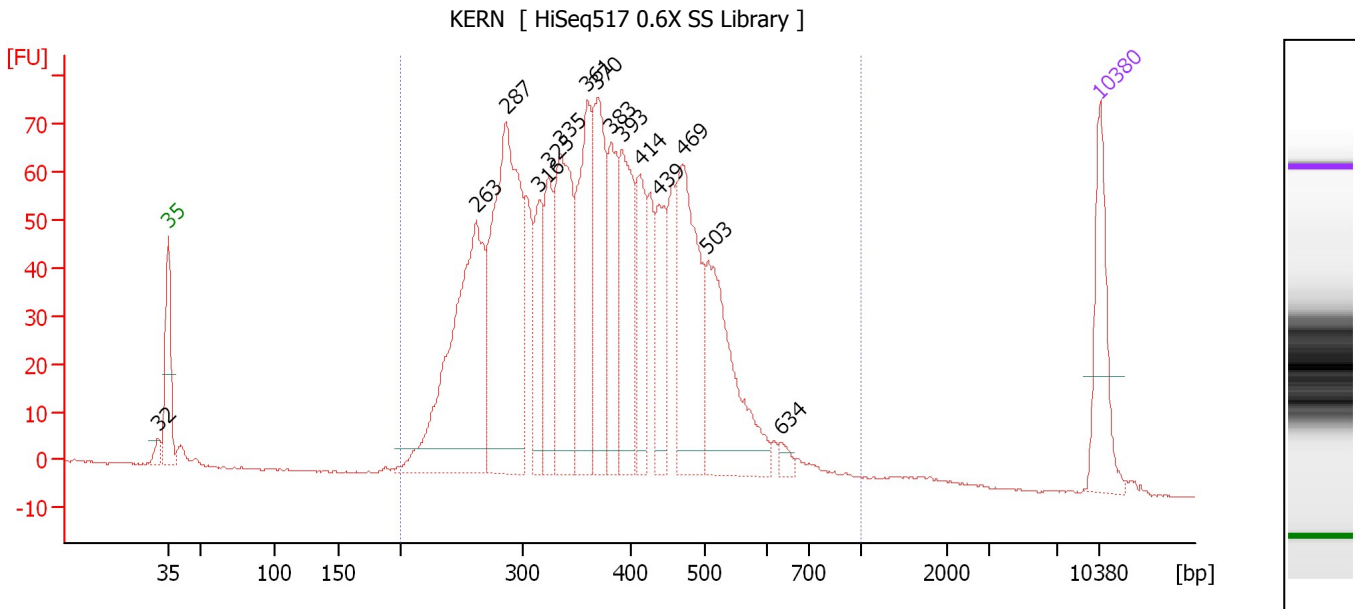
Region table for sample 7 : FRES

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	389	1,766.6	10,557.2	2,493.46	99	24.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
 Modified: 7/3/2016 3:52:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : KERN

Number of peaks found: 15 Corr. Area 1: 1,730.3
 Noise: 0.2

Peak table for sample 8 : KERN

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.20
2	35	125.00	5,411.3	Lower Marker	43.00
3	263	380.37	2,194.6		66.17
4	287	400.93	2,116.8		68.41
5	316	89.84	430.6		70.90
6	325	101.47	473.5		71.60
7	335	209.62	947.5		72.45
8	361	171.98	722.1		74.52
9	370	159.04	651.1		75.26
10	383	100.83	399.3		76.27
11	393	144.55	557.2		77.12
12	414	92.36	338.4		78.45
13	439	77.79	268.6		79.89
14	469	200.82	649.1		81.59
15	503	192.50	579.9		83.50
16	634	9.88	23.6		89.03
17	10,380	75.00	10.9	Upper Marker	113.00

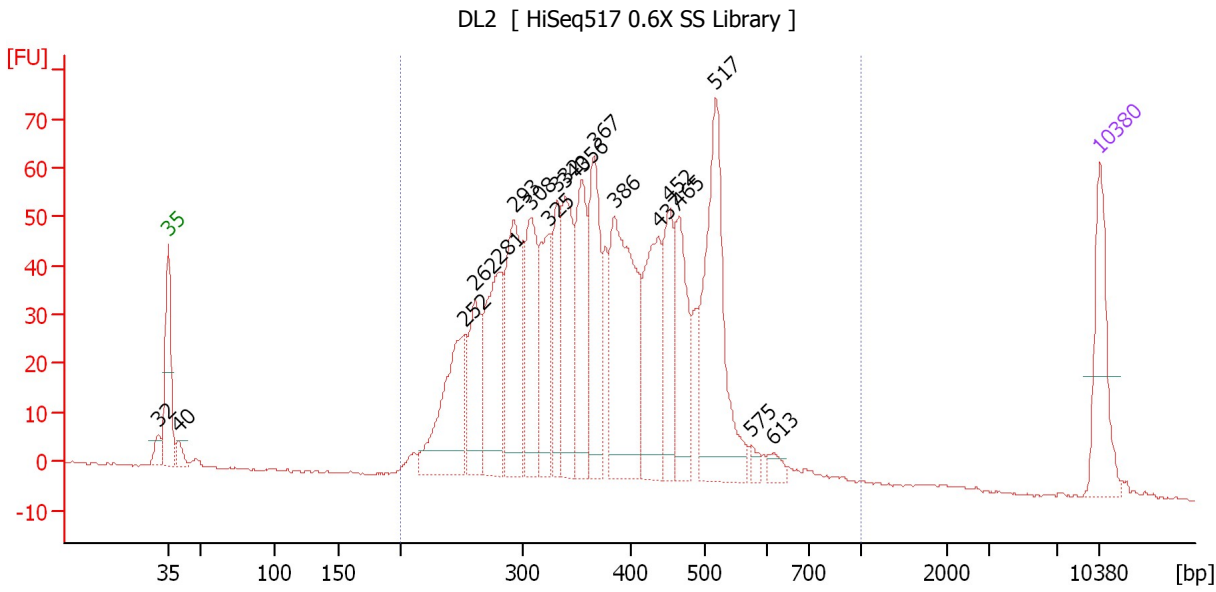
Region table for sample 8 : KERN

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	380	1,730.3	11,428.1	2,620.24	98	26.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
 Modified: 7/3/2016 3:52:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : DL2

Number of peaks found: 19 Corr. Area 1: 1,354.5
 Noise: 0.2

Peak table for sample 9 : DL2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.25
2	35	125.00	5,411.3	Lower Marker	43.00
3	40	16.80	639.9		43.75
4	252	169.33	1,017.0		65.23
5	262	102.11	591.4		66.08
6	281	169.93	916.5		67.85
7	293	187.03	966.5		68.98
8	308	152.82	751.3		70.26
9	325	104.06	485.5		71.60
10	332	92.26	421.0		72.19
11	340	135.08	602.0		72.83
12	356	133.92	570.1		74.12
13	367	146.20	604.4		74.97
14	386	252.41	991.5		76.53
15	437	148.00	512.9		79.79
16	452	102.60	343.7		80.65
17	465	112.05	364.8		81.40
18	517	248.44	728.5		84.13
19	575	9.19	24.2		86.81
20	613	12.56	31.1		88.36
21	10,380	75.00	10.9	Upper Marker	113.00

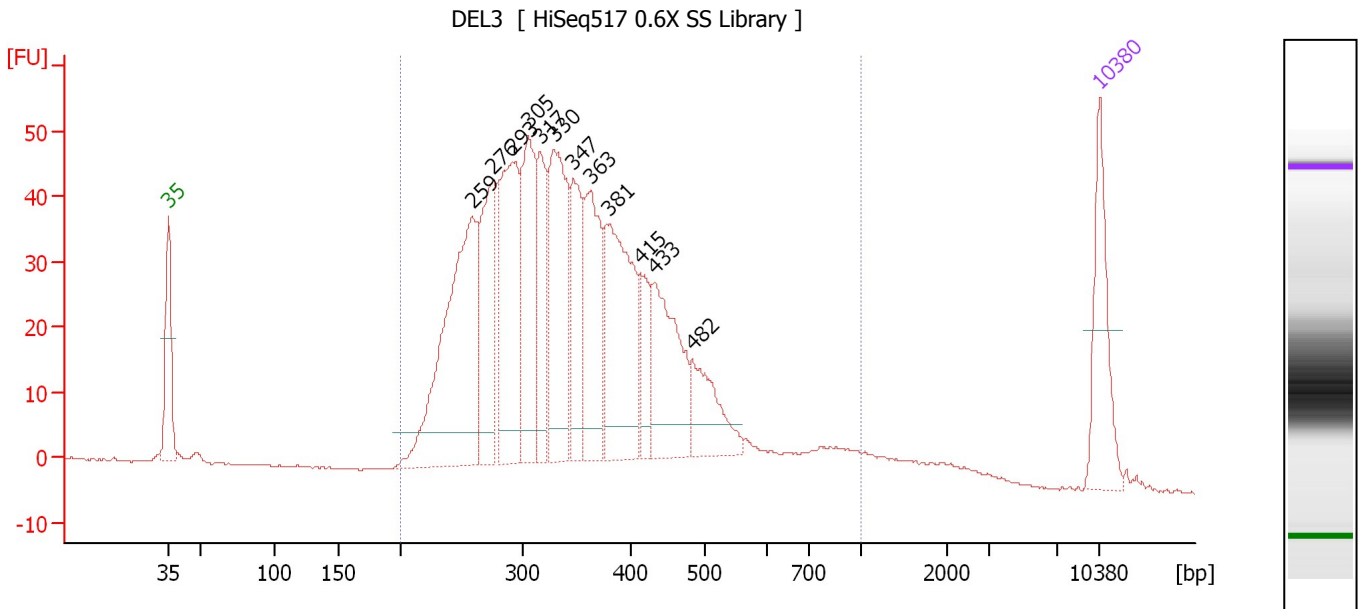
Region table for sample 9 : DL2

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	382	1,354.5	10,383.6	2,402.58	99	24.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
 Modified: 7/3/2016 3:52:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : DEL3

Number of peaks found: 12 Corr. Area 1: 1,086.6
 Noise: 0.2

Peak table for sample 10 : DEL3

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	259	344.93	2,016.0		65.86
3	276	152.45	837.7		67.38
4	293	218.94	1,132.7		68.95
5	305	159.69	792.5		70.03
6	317	102.78	491.8		70.95
7	330	195.71	898.5		72.03
8	347	98.36	428.9		73.43
9	363	144.69	603.1		74.73
10	381	196.72	782.6		76.14
11	415	46.84	171.1		78.51
12	433	137.97	483.1		79.54
13	482	73.07	229.6		82.35
14	10,380	75.00	10.9	Upper Marker	113.00

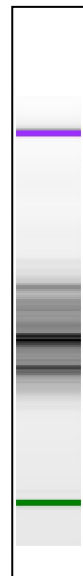
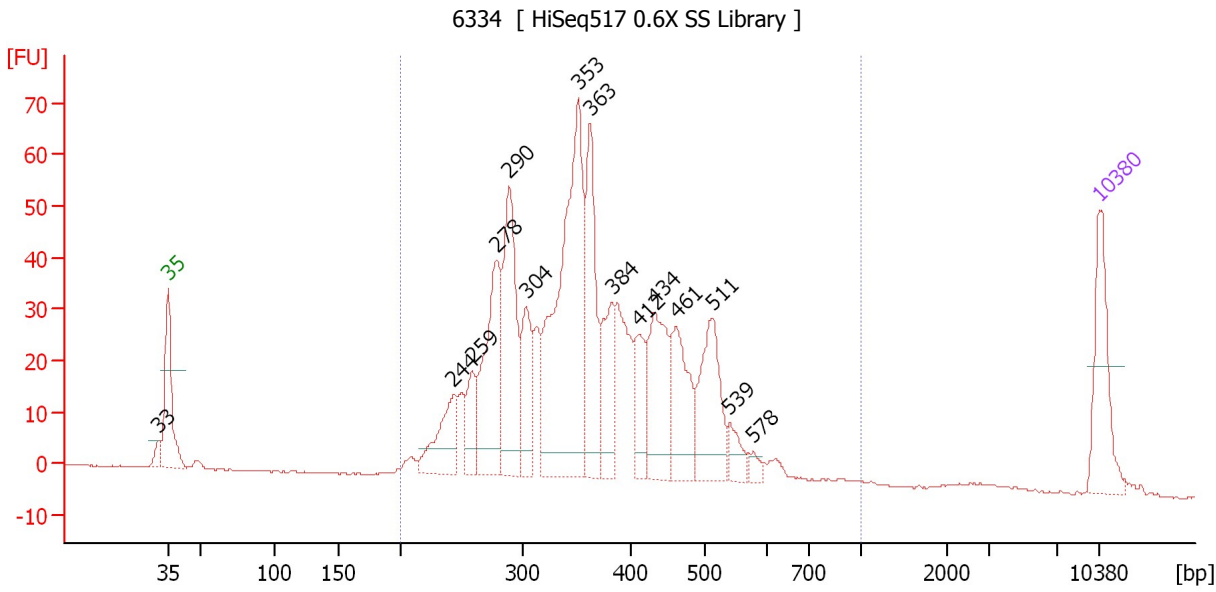
Region table for sample 10 : DEL3

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	363	1,086.6	9,826.7	2,130.76	96	31.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
 Modified: 7/3/2016 3:52:25 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : 6334

Number of peaks found: 15 Corr. Area 1: 905.0
 Noise: 0.1

Peak table for sample 11 : 6334

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	33	0.00	0.0		42.29
2	35	125.00	5,411.3	Lower Marker	43.00
3	244	86.11	535.5		64.44
4	259	52.60	308.2		65.80
5	278	174.80	951.8		67.61
6	290	187.74	982.2		68.65
7	304	82.56	411.7		69.91
8	353	390.46	1,678.2		73.84
9	363	144.79	603.8		74.72
10	384	81.92	323.6		76.36
11	412	59.68	219.3		78.38
12	434	120.43	420.9		79.59
13	461	102.29	335.9		81.17
14	511	118.04	350.3		83.85
15	539	24.39	68.5		85.16
16	578	10.02	26.2		86.97
17	10,380	75.00	10.9	Upper Marker	113.00

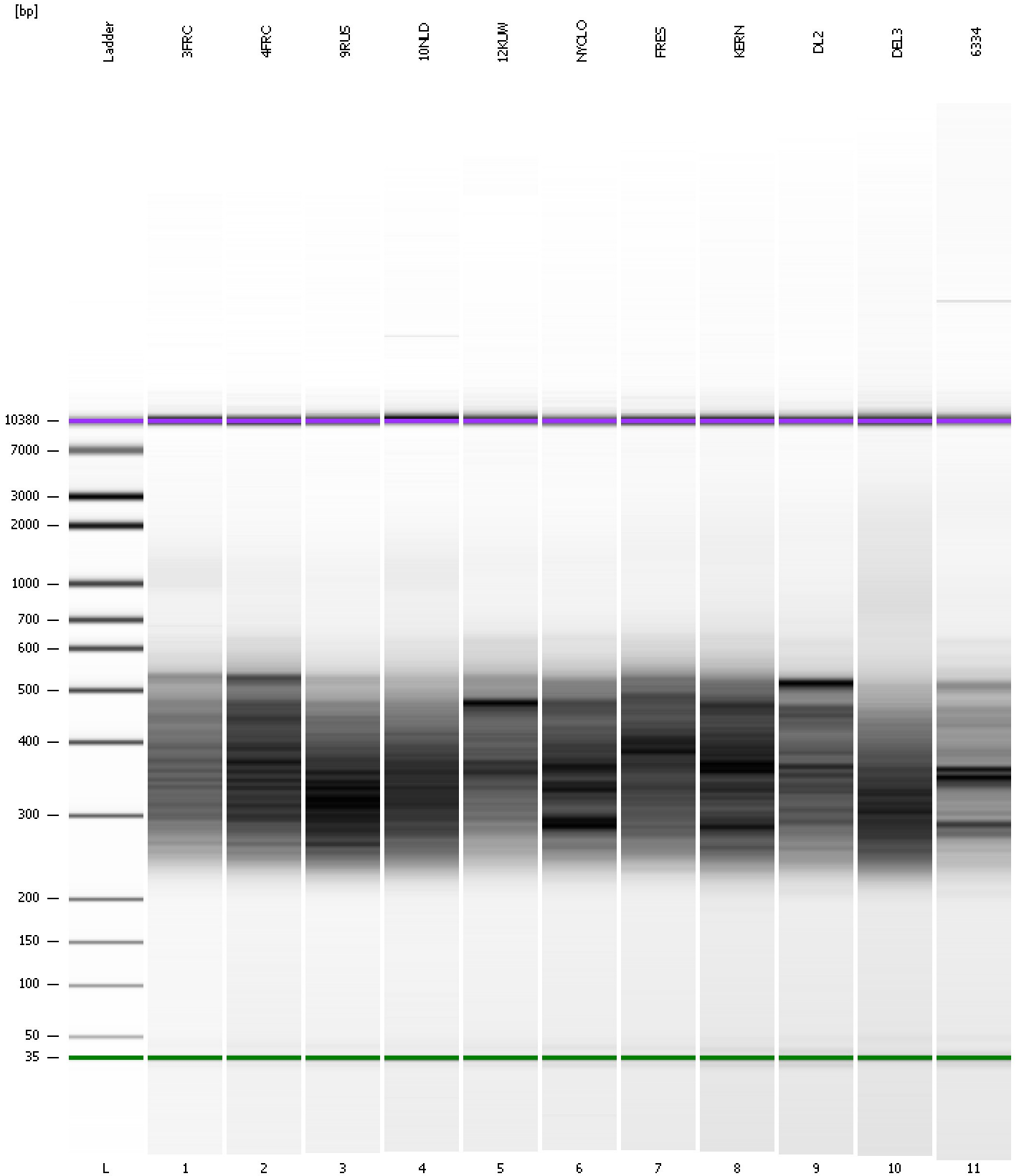
Region table for sample 11 : 6334

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	367	905.0	8,164.3	1,845.25	99	22.7

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
Modified: 7/3/2016 3:52:25 PM

Gel Image

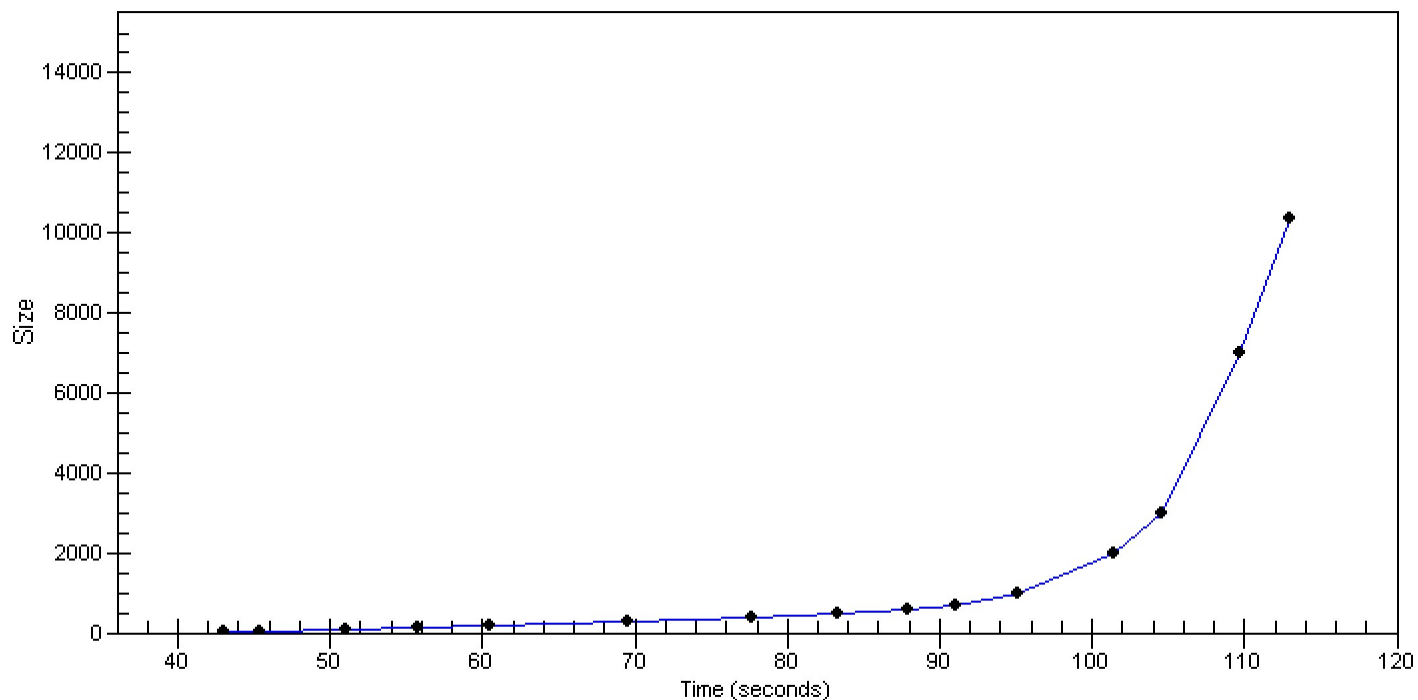


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad

Created: 7/3/2016 2:49:37 PM
Modified: 7/3/2016 3:52:25 PM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay Created: 7/3/2016 2:49:37 PM
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad Modified: 7/3/2016 3:52:25 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		7/3/2016 3:30:54 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-07-03\2016-07-03_001.xad)		Instrument	Run		7/3/2016 2:49:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		7/3/2016 2:49:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		7/3/2016 2:49:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		7/3/2016 2:49:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		7/3/2016 2:49:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		7/3/2016 2:49:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		7/3/2016 2:49:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1