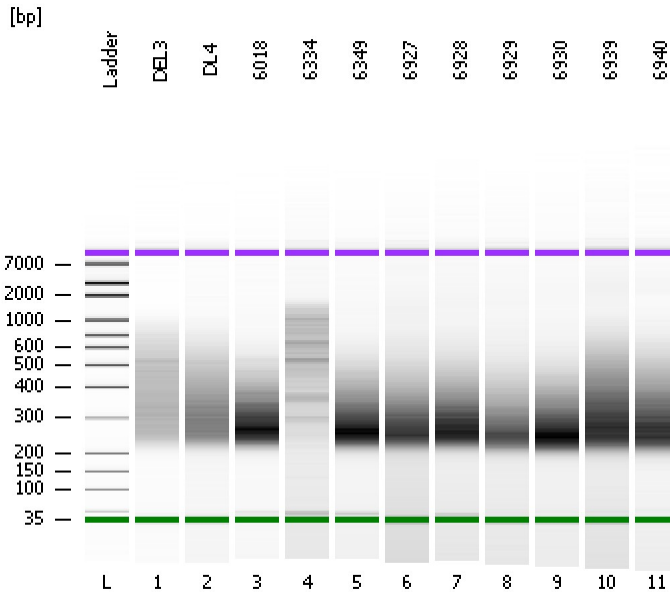


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
Modified: 6/21/2016 4:21:51 PM

Electrophoresis File Run Summary



Instrument Information:

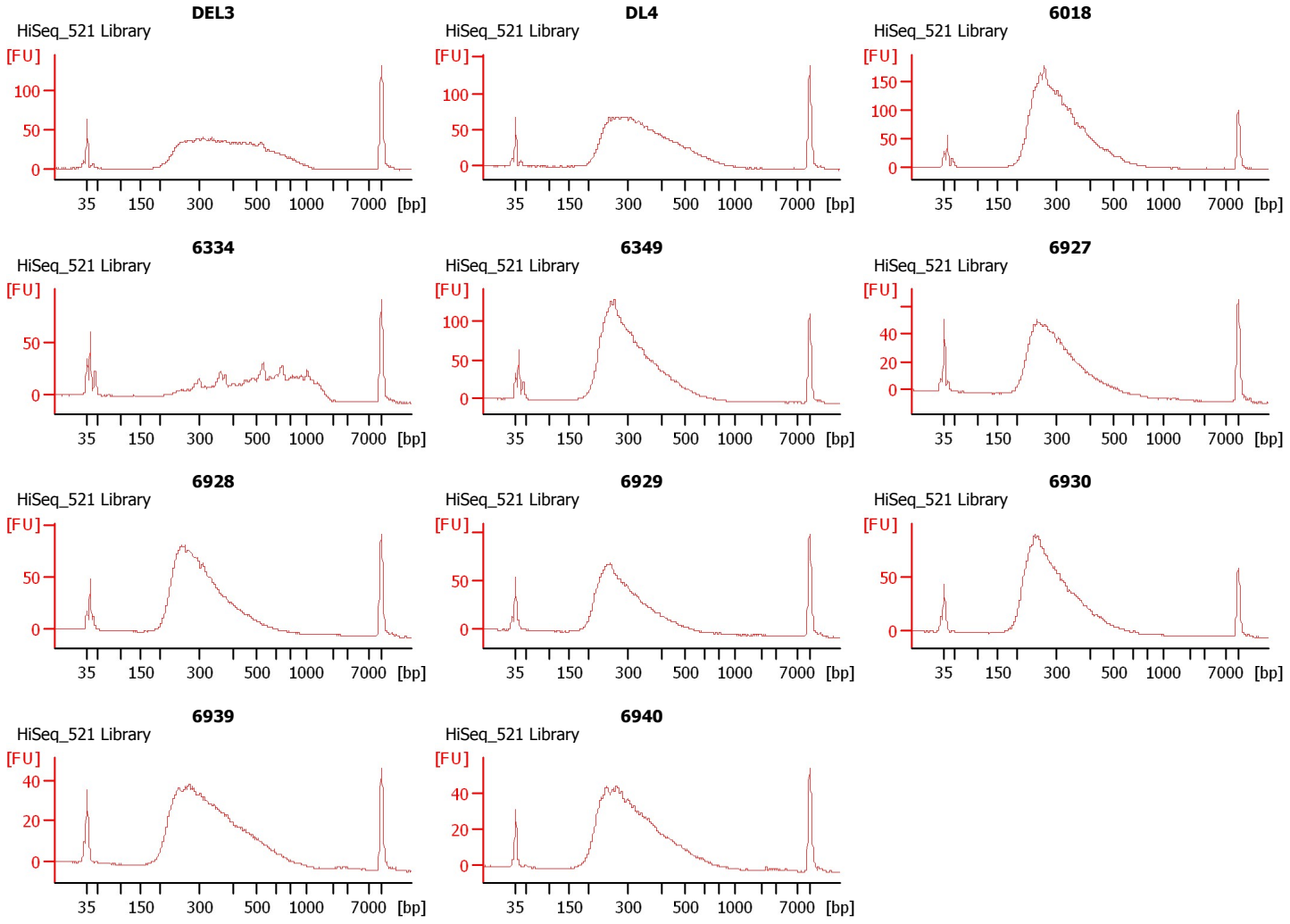
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
DEL3	HiSeq_521 Library	<input type="checkbox"/>	✓			
DL4	HiSeq_521 Library	<input type="checkbox"/>	✓			
6018	HiSeq_521 Library	<input type="checkbox"/>	✓			
6334	HiSeq_521 Library	<input type="checkbox"/>	✓			
6349	HiSeq_521 Library	<input type="checkbox"/>	✓			
6927	HiSeq_521 Library	<input type="checkbox"/>	✓			
6928	HiSeq_521 Library	<input type="checkbox"/>	✓			
6929	HiSeq_521 Library	<input type="checkbox"/>	✓			
6930	HiSeq_521 Library	<input type="checkbox"/>	✓			
6939	HiSeq_521 Library	<input type="checkbox"/>	✓			
6940	HiSeq_521 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\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
Modified: 6/21/2016 4:21:51 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

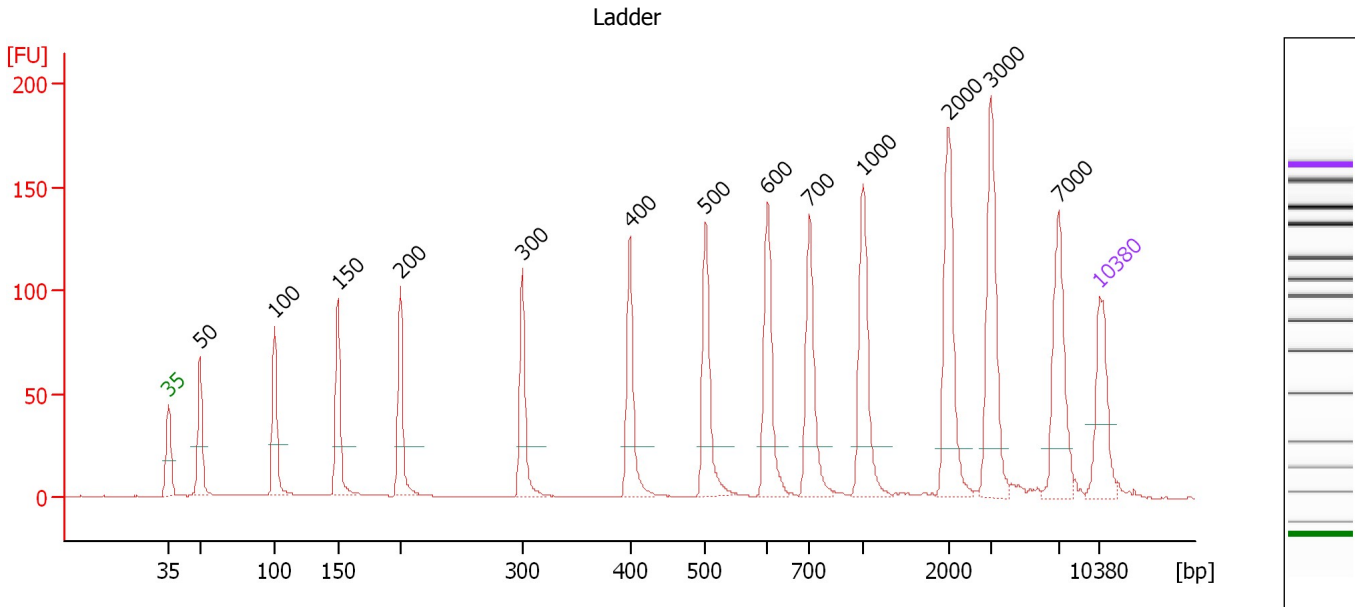
Ladder

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\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

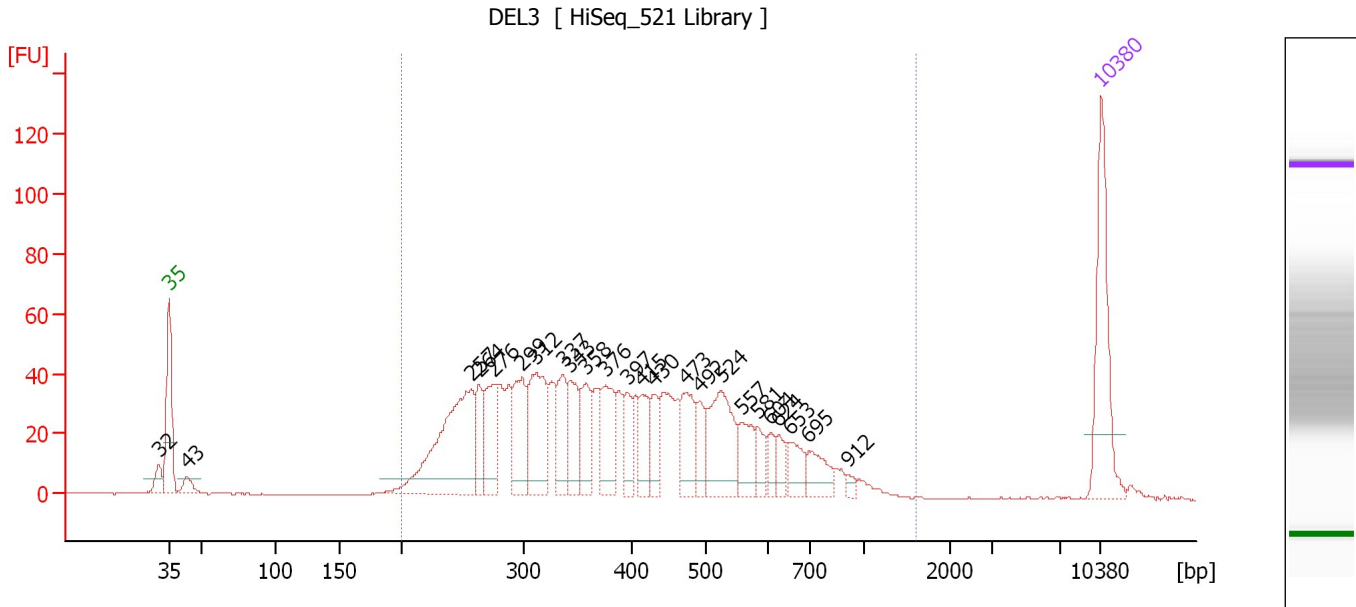
Peak table for Ladder

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	45.37
3	100	150.00	2,272.7	Ladder Peak	50.98
4	150	150.00	1,515.2	Ladder Peak	55.73
5	200	150.00	1,136.4	Ladder Peak	60.43
6	300	150.00	757.6	Ladder Peak	69.59
7	400	150.00	568.2	Ladder Peak	77.67
8	500	150.00	454.5	Ladder Peak	83.37
9	600	150.00	378.8	Ladder Peak	88.02
10	700	150.00	324.7	Ladder Peak	91.20
11	1,000	150.00	227.3	Ladder Peak	95.19
12	2,000	150.00	113.6	Ladder Peak	101.60
13	3,000	150.00	75.8	Ladder Peak	104.83
14	7,000	150.00	32.5	Ladder Peak	109.91
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : DEL3

Number of peaks found: 24 Corr. Area 1: 1,280.9
 Noise: 0.2

Peak table for sample 1 : DEL3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.18
2	35	125.00	5,411.3	Lower Marker	43.00
3	43	14.64	509.9		44.35
4	257	182.23	1,073.3		65.68
5	264	39.32	226.1		66.25
6	276	58.63	321.7		67.41
7	299	63.29	321.1		69.47
8	312	88.37	428.9		70.58
9	337	46.59	209.4		72.60
10	343	48.82	215.6		73.08
11	358	39.14	165.7		74.28
12	376	54.47	219.2		75.77
13	397	25.52	97.5		77.40
14	415	30.57	111.7		78.50
15	430	28.38	100.0		79.37
16	473	49.15	157.5		81.82
17	492	25.08	77.2		82.92
18	524	77.62	224.6		84.46
19	557	30.70	83.6		86.00
20	581	17.66	46.0		87.15
21	604	12.48	31.3		88.16
22	624	15.40	37.4		88.79
23	653	22.95	53.3		89.70
24	695	26.19	57.1		91.04
25	912	5.01	8.3		94.02

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...

... Peak table for sample 1 : DEL3

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

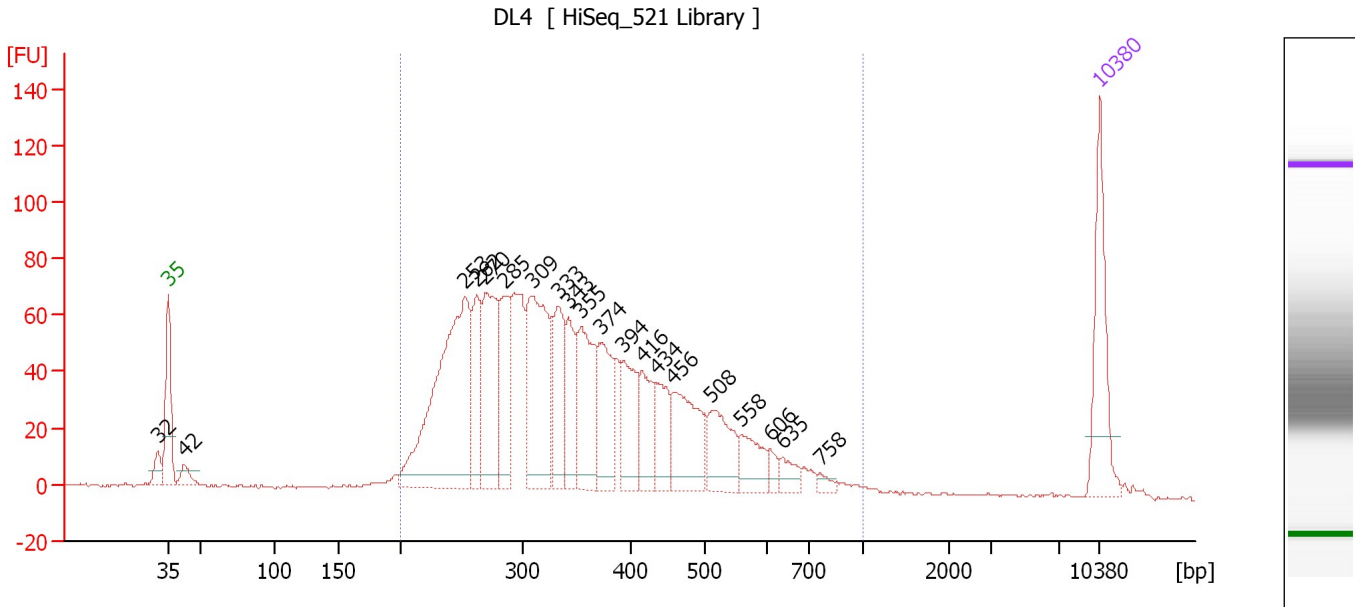
Region table for sample 1 : DEL3

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
200	1,601	414	1,280.9	5,366.3	1,246.60	97	37.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : DL4

Number of peaks found: 20 Corr. Area 1: 1,793.2
 Noise: 0.4

Peak table for sample 2 : DL4

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.18
2	35	125.00	5,411.3	Lower Marker	43.00
3	42	19.02	681.5		44.15
4	253	341.48	2,045.8		65.28
5	262	85.25	492.4		66.14
6	270	149.71	841.2		66.81
7	285	87.78	466.9		68.21
8	309	182.23	893.7		70.32
9	333	88.63	402.9		72.29
10	343	65.74	290.5		73.05
11	355	108.44	463.2		74.02
12	374	83.10	336.9		75.55
13	394	69.18	266.0		77.18
14	416	57.48	209.4		78.58
15	434	50.14	174.8		79.63
16	456	96.50	320.4		80.88
17	508	66.40	197.9		83.76
18	558	40.05	108.8		86.07
19	606	10.05	25.1		88.23
20	635	17.24	41.1		89.14
21	758	7.84	15.7		91.97
22	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...

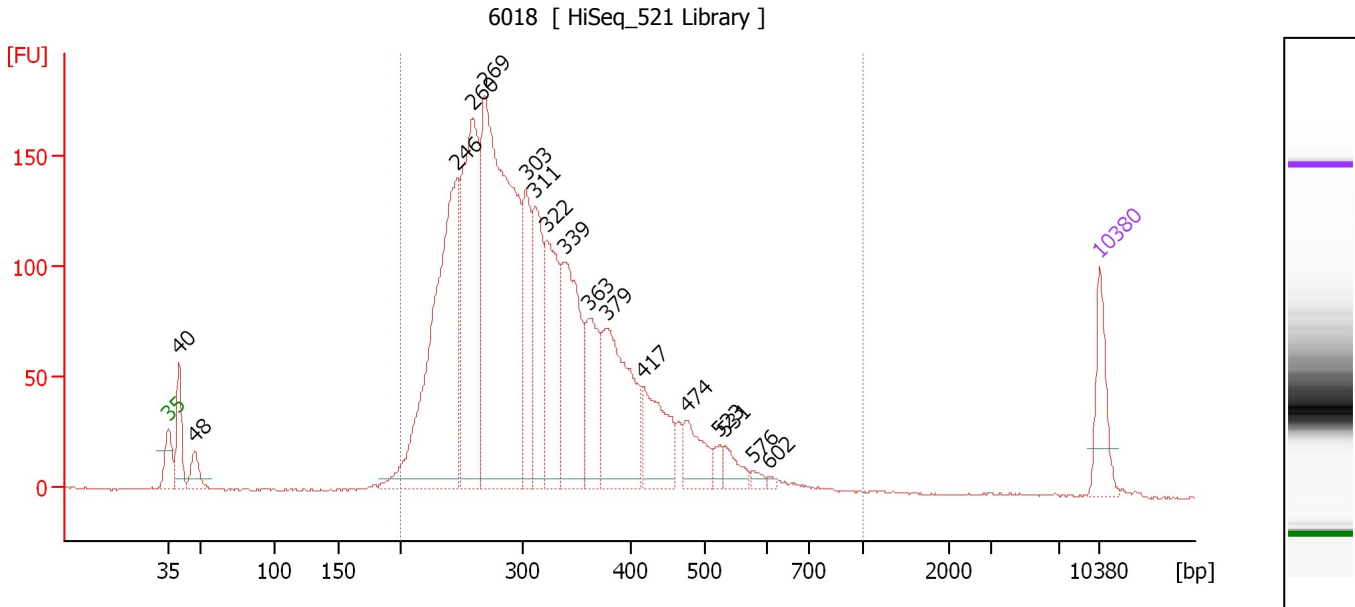
... Region table for sample 2 : DL4

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	1,793.2	8,497.2	■ 1,808.94	96	32.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : 6018

Number of peaks found: 17 Corr. Area 1: 3,074.0
 Noise: 0.4

Peak table for sample 3 : 6018

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	40	118.95	4,489.1		43.82
3	48	54.65	1,735.1		45.01
4	246	799.70	4,919.7		64.67
5	260	544.87	3,176.6		65.92
6	269	1,080.76	6,080.5		66.78
7	303	212.67	1,062.9		69.85
8	311	252.98	1,230.6		70.52
9	322	249.87	1,175.1		71.38
10	339	318.55	1,422.1		72.77
11	363	164.44	687.2		74.64
12	379	321.32	1,286.0		75.94
13	417	160.38	583.0		78.62
14	474	88.11	281.7		81.88
15	523	20.85	60.4		84.42
16	531	40.28	114.9		84.81
17	576	12.28	32.3		86.92
18	602	4.73	11.9		88.07
19	10,380	75.00	10.9	Upper Marker	113.00

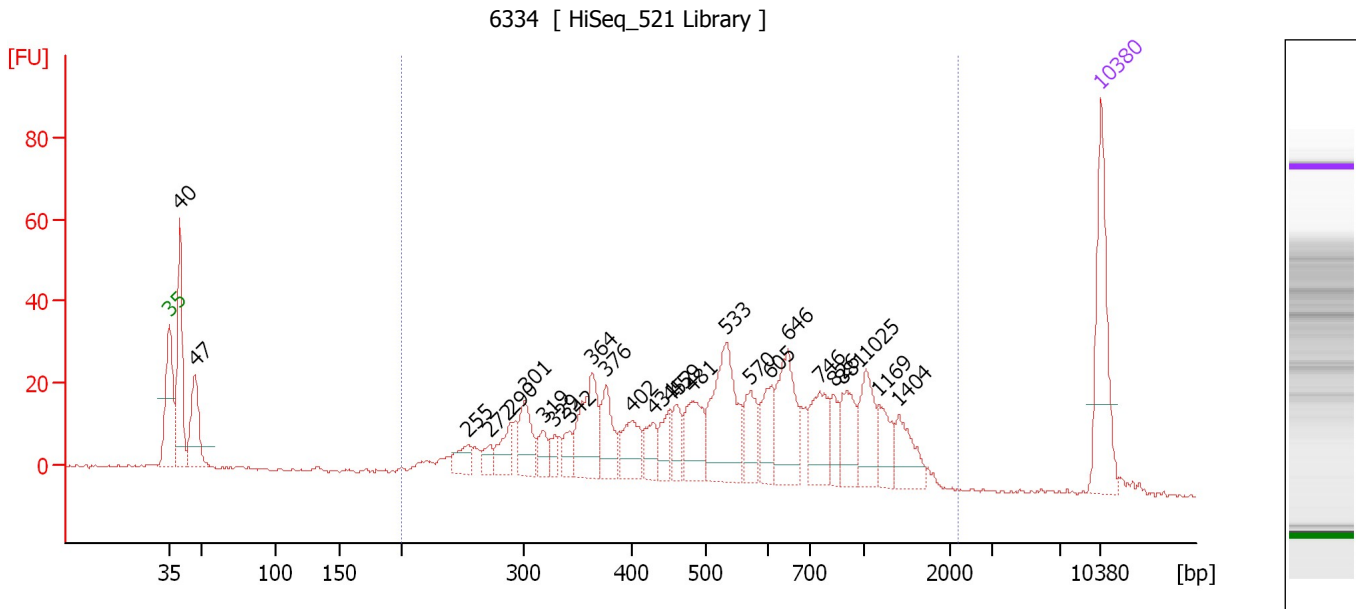
Region table for sample 3 : 6018

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	324	3,074.0	22,137.8	4,373.42	96	27.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : 6334

Number of peaks found: 26 Corr. Area 1: 785.0
 Noise: 0.4

Peak table for sample 4 : 6334

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	40	139.91	5,313.0		43.78
3	47	76.83	2,479.6		44.89
4	255	23.92	142.3		65.44
5	272	13.96	77.7		67.04
6	290	29.32	153.4		68.64
7	301	43.53	218.8		69.71
8	319	19.13	90.9		71.12
9	329	12.78	58.8		71.94
10	342	18.13	80.3		73.01
11	364	75.05	312.0		74.80
12	376	41.04	165.1		75.77
13	402	39.51	148.8		77.81
14	431	24.68	86.9		79.41
15	452	25.26	84.7		80.62
16	459	23.30	76.8		81.05
17	481	52.47	165.4		82.27
18	533	108.25	307.9		84.88
19	570	36.15	96.1		86.63
20	605	37.04	92.7		88.18
21	646	72.84	170.8		89.49
22	746	46.85	95.1		91.82
23	826	19.48	35.7		92.88
24	881	38.36	66.0		93.61
25	1,025	43.25	63.9		95.35

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...

... Peak table for sample 4 : 6334

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	1,169	26.56	34.4		96.28
27	1,404	32.41	35.0		97.78
28	10,380	75.00	10.9	Upper Marker	113.00

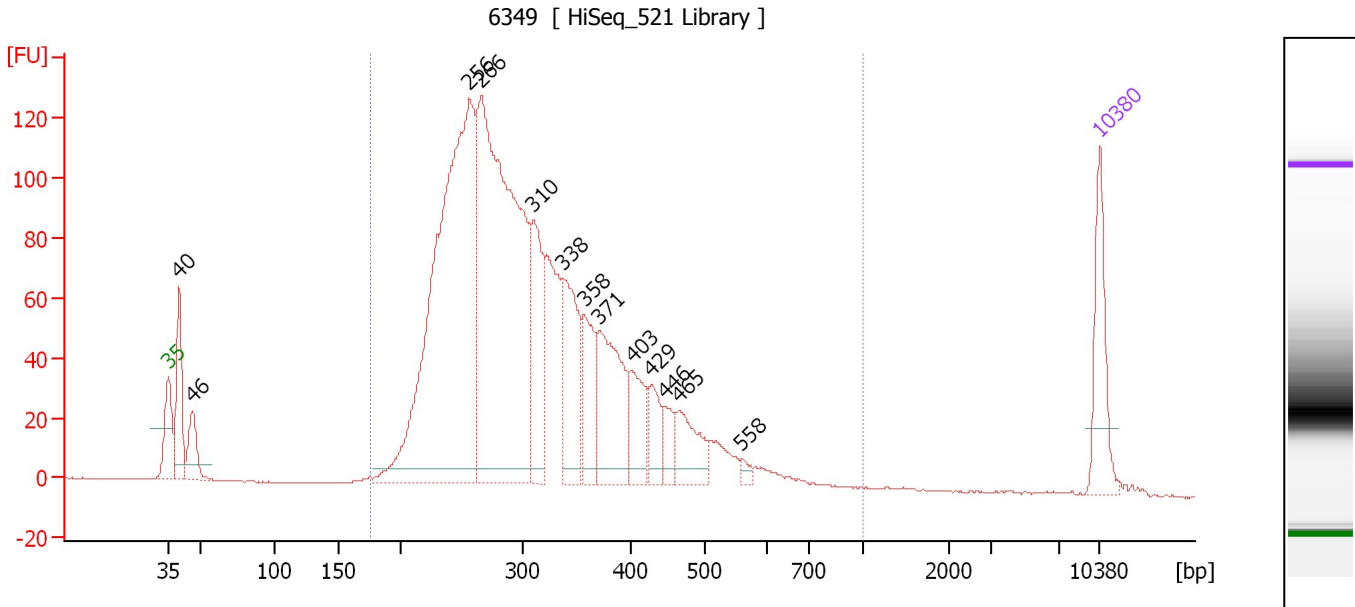
Region table for sample 4 : 6334

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
200	2,190	623	785.0	3,500.3	1,048.87	83	53.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : 6349

Number of peaks found: 13 Corr. Area 1: 2,267.0
 Noise: 0.3

Peak table for sample 5 : 6349

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	40	120.30	4,562.3		43.78
3	46	64.65	2,108.9		44.81
4	256	925.49	5,472.7		65.58
5	266	859.04	4,886.2		66.51
6	310	149.88	731.7		70.43
7	338	153.69	688.4		72.69
8	358	90.40	382.3		74.30
9	371	169.49	692.1		75.33
10	403	74.88	281.6		77.83
11	429	47.58	168.2		79.30
12	446	33.80	114.9		80.28
13	465	67.72	220.8		81.36
14	558	9.03	24.5		86.06
15	10,380	75.00	10.9	Upper Marker	113.00

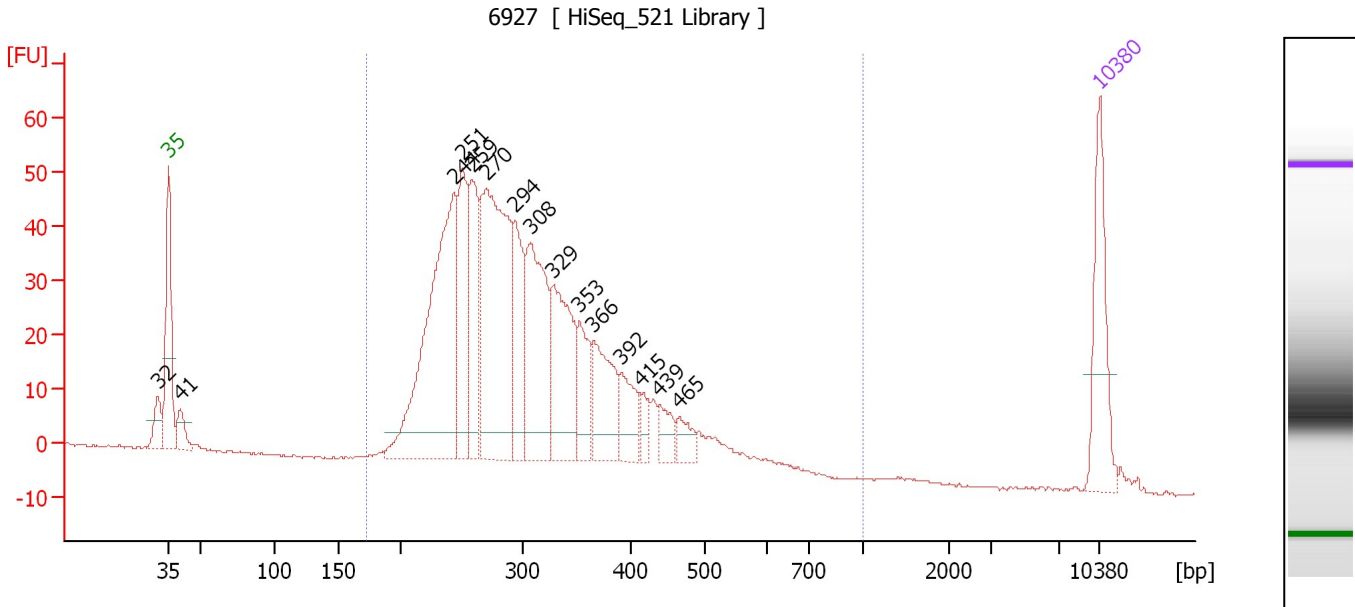
Region table for sample 5 : 6349

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
176	1,000	319	2,267.0	14,942.3	2,898.23	95	28.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : 6927

Number of peaks found: 15 Corr. Area 1: 945.6
 Noise: 0.2

Peak table for sample 6 : 6927

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	41	29.74	1,107.5		43.90
4	244	369.15	2,291.9		64.47
5	251	123.15	744.6		65.07
6	259	118.10	691.4		65.82
7	270	314.55	1,763.6		66.87
8	294	105.40	542.7		69.07
9	308	175.65	864.8		70.22
10	329	144.48	665.7		71.92
11	353	57.99	248.9		73.87
12	366	84.93	351.6		74.92
13	392	48.44	187.2		77.02
14	415	16.35	59.7		78.53
15	439	24.64	85.1		79.88
16	465	21.99	71.6		81.38
17	10,380	75.00	10.9	Upper Marker	113.00

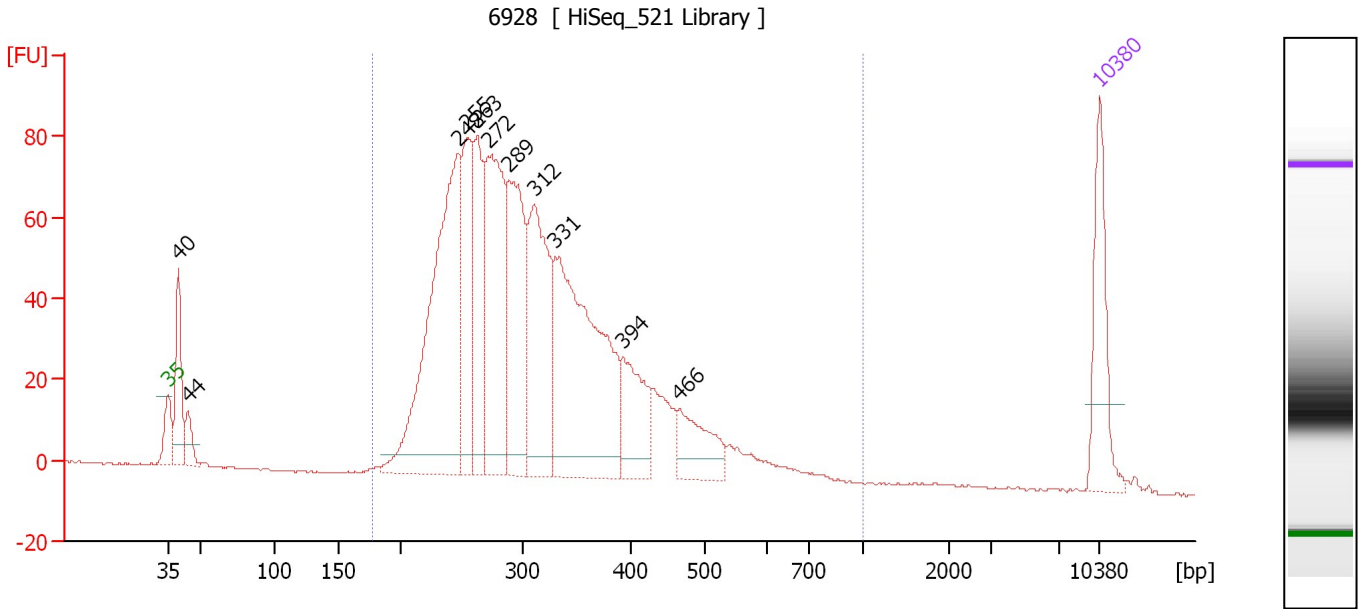
Region table for sample 6 : 6927

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
173	1,000	313	945.6	9,022.1	1,735.73	97	25.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : 6928

Number of peaks found: 11 Corr. Area 1: 1,553.2
 Noise: 0.3

Peak table for sample 7 : 6928

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	40	98.29	3,746.2		43.75
3	44	30.49	1,045.5		44.45
4	248	440.91	2,695.6		64.81
5	255	159.33	946.9		65.46
6	263	137.08	789.3		66.22
7	272	270.09	1,505.1		67.02
8	289	208.16	1,089.8		68.62
9	312	217.35	1,057.0		70.53
10	331	375.28	1,718.7		72.08
11	394	89.55	344.8		77.15
12	466	68.36	222.4		81.41
13	10,380	75.00	10.9	Upper Marker	113.00

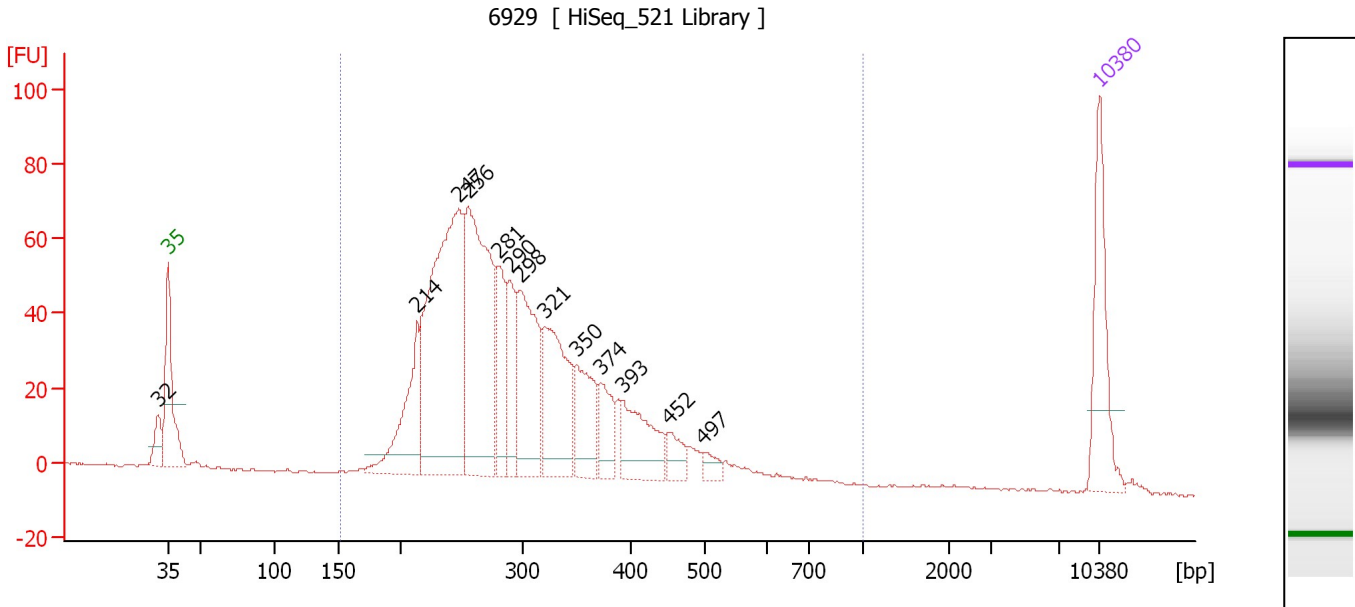
Region table for sample 7 : 6928

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
178	1,000	321	1,553.2	10,510.9	2,055.74	95	27.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : 6929

Number of peaks found: 13 Corr. Area 1: 1,203.1
 Noise: 0.2

Peak table for sample 8 : 6929

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.23
2	35	125.00	5,411.3	Lower Marker	43.00
3	214	115.44	818.1		61.69
4	247	396.72	2,429.4		64.78
5	256	305.83	1,811.3		65.55
6	281	79.55	428.9		67.86
7	290	68.20	356.3		68.68
8	298	152.12	773.8		69.40
9	321	145.88	688.4		71.30
10	350	78.18	338.7		73.61
11	374	41.80	169.4		75.56
12	393	76.45	294.8		77.10
13	452	24.66	82.6		80.64
14	497	12.76	38.9		83.21
15	10,380	75.00	10.9	Upper Marker	113.00

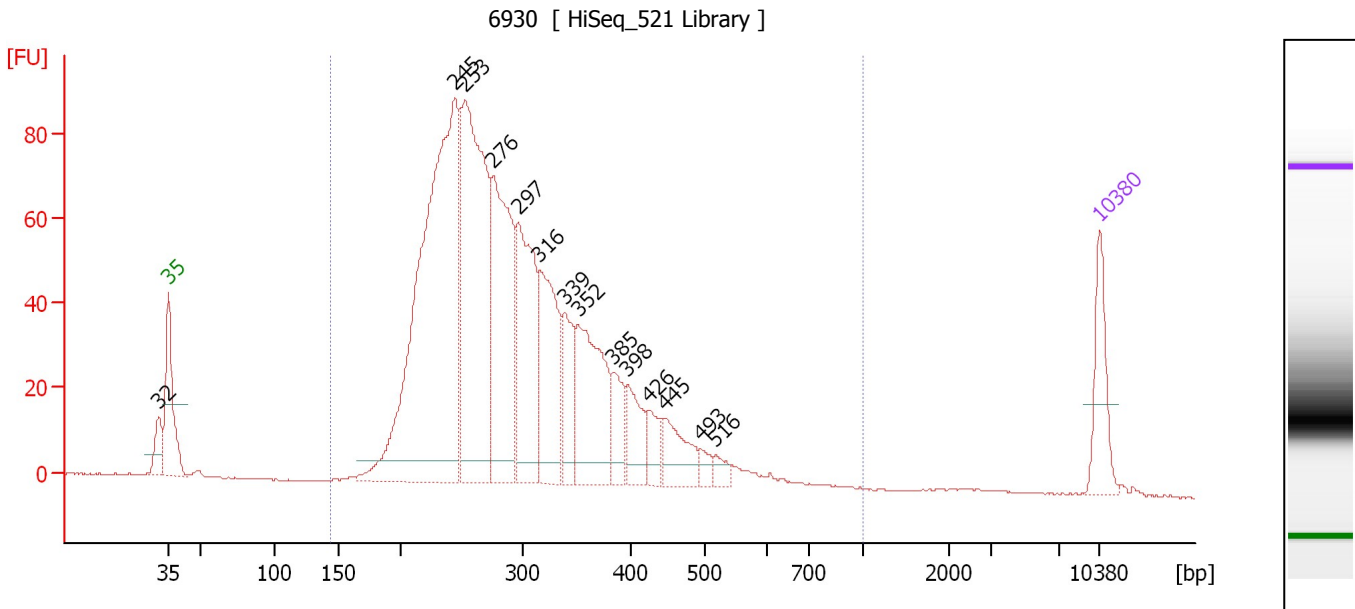
Region table for sample 8 : 6929

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
152	1,000	304	1,203.1	8,386.9	1,561.48	98	27.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : 6930

Number of peaks found: 14 Corr. Area 1: 1,494.0
 Noise: 0.1

Peak table for sample 9 : 6930

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	245	1,067.12	6,609.6		64.52
4	253	656.62	3,930.6		65.30
5	276	401.76	2,203.1		67.42
6	297	305.15	1,558.5		69.29
7	316	232.21	1,112.9		70.90
8	339	110.10	491.7		72.76
9	352	243.44	1,047.6		73.80
10	385	64.92	255.6		76.44
11	398	77.67	295.9		77.48
12	426	40.71	144.8		79.14
13	445	80.60	274.5		80.23
14	493	17.53	53.9		82.98
15	516	20.47	60.1		84.12
16	10,380	75.00	10.9	Upper Marker	113.00

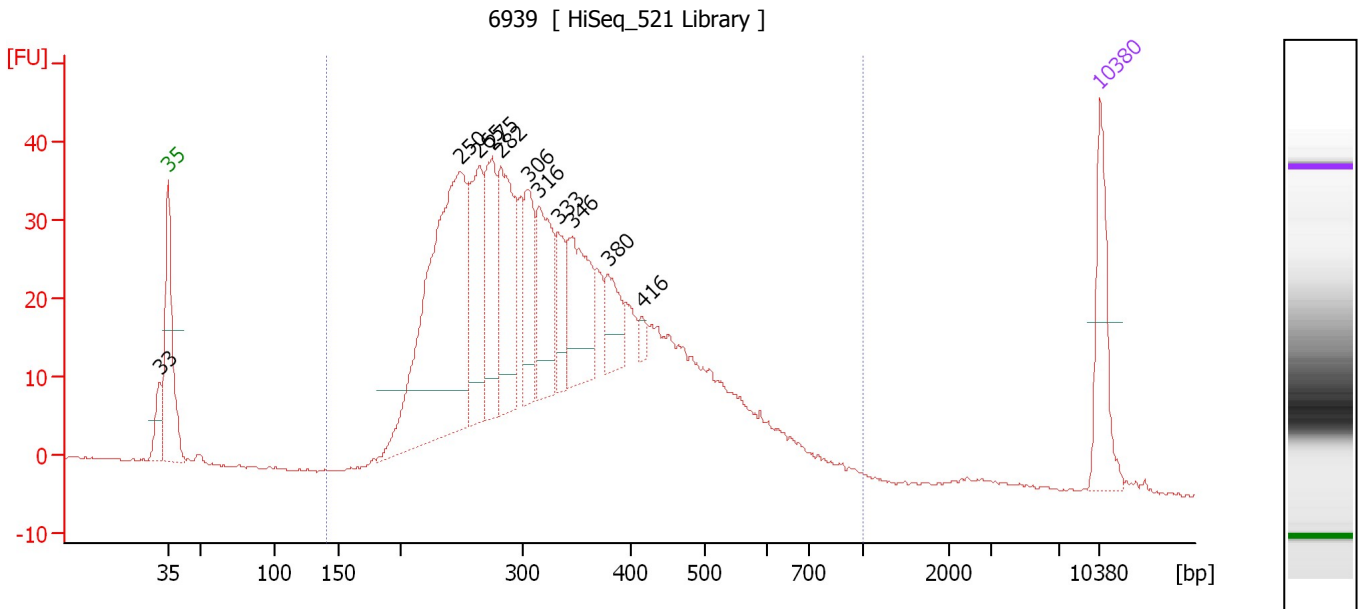
Region table for sample 9 : 6930

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
145	1,000	302	1,494.0	18,508.8	3,417.12	98	27.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : 6939

Number of peaks found: 11 Corr. Area 1: 935.7
 Noise: 0.2

Peak table for sample 10 : 6939

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	33	0.00	0.0		42.32
2	35	125.00	5,411.3	Lower Marker	43.00
3	250	525.70	3,191.0		64.98
4	265	164.58	941.0		66.39
5	275	134.79	742.0		67.33
6	282	178.41	958.3		67.95
7	306	91.64	454.5		70.04
8	316	127.68	612.5		70.87
9	333	53.49	243.1		72.28
10	346	123.12	538.8		73.33
11	380	54.47	217.3		76.04
12	416	10.87	39.6		78.60
13	10,380	75.00	10.9	Upper Marker	113.00

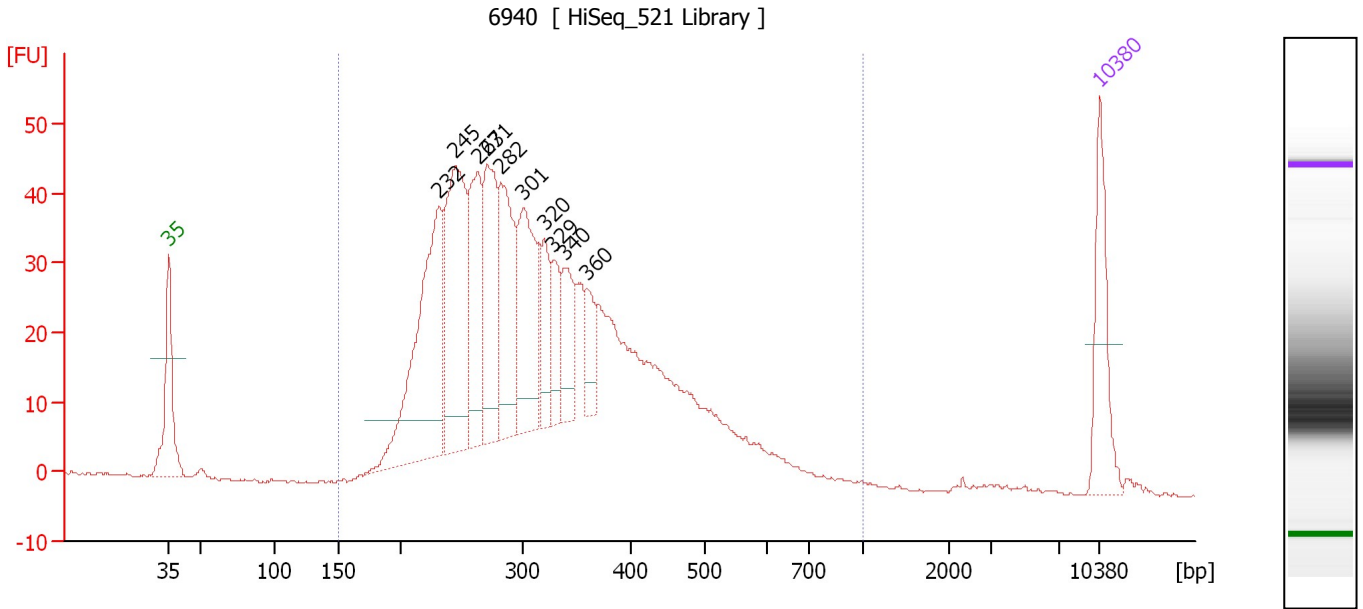
Region table for sample 10 : 6939

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
141	1,000	351	935.7	12,937.6	2,643.39	97	33.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : 6940

Number of peaks found: 10 Corr. Area 1: 961.0
 Noise: 0.2

Peak table for sample 11 : 6940

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	232	319.59	2,091.2		63.32
3	245	272.97	1,686.5		64.57
4	263	155.16	892.3		66.25
5	271	161.35	900.6		66.98
6	282	157.76	846.8		67.97
7	301	173.32	871.8		69.69
8	320	69.61	329.6		71.21
9	329	53.61	246.8		71.94
10	340	69.28	308.7		72.83
11	360	40.33	169.7		74.45
12	10,380	75.00	10.9	Upper Marker	113.00

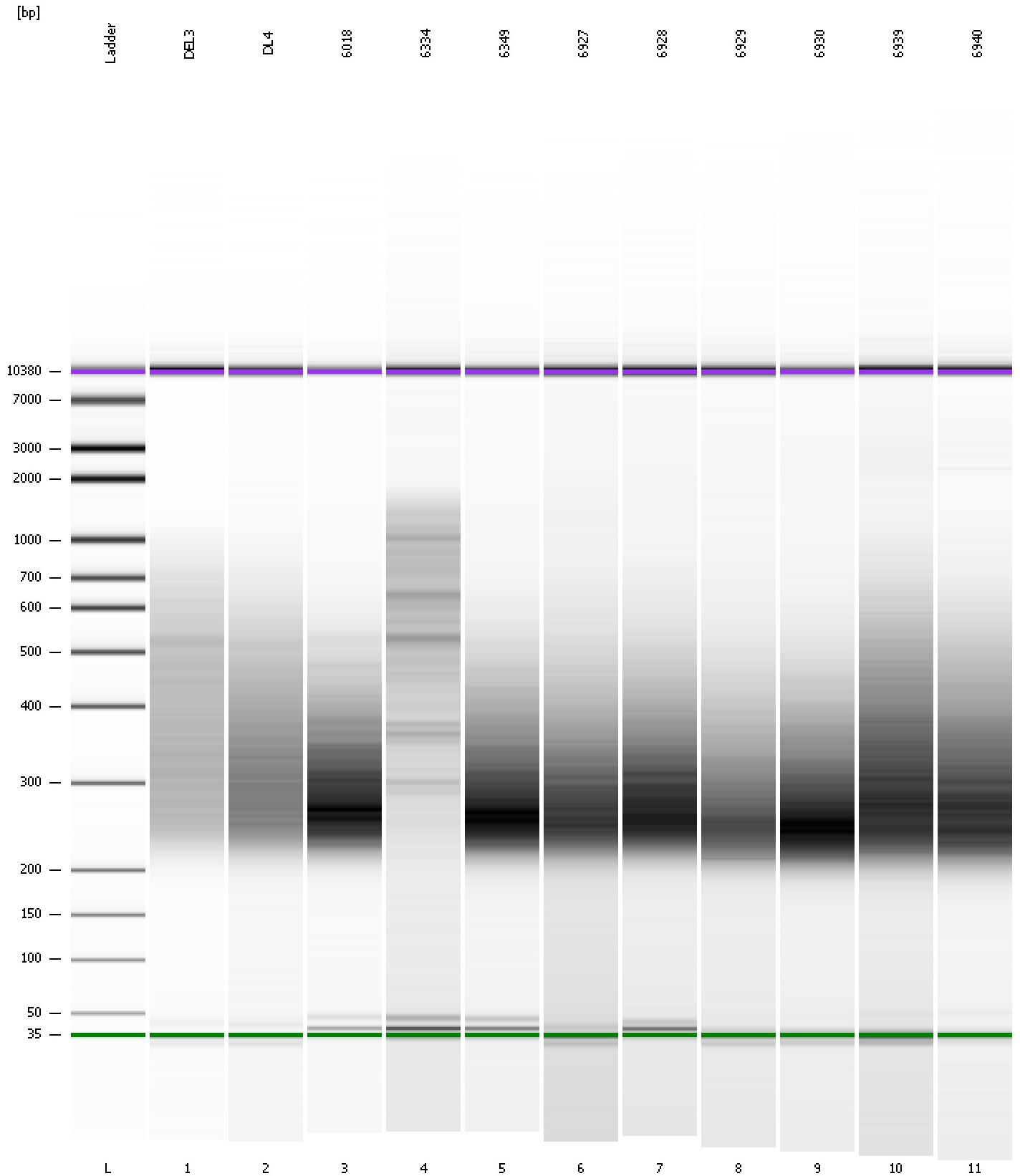
Region table for sample 11 : 6940

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
150	1,000	331	961.0	12,022.8	2,359.76	99	31.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
Modified: 6/21/2016 4:21:51 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-06-21\2016-06-21_005.xad

Created: 6/21/2016 3:32:54 PM
 Modified: 6/21/2016 4:21:51 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		6/21/2016 4:13:23 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\2016-06-21\2016-06-21_005.xad)		Instrument	Run		6/21/2016 3:33:00 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		6/21/2016 3:33:00 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		6/21/2016 3:33:00 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		6/21/2016 3:33:00 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		6/21/2016 3:33:00 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		6/21/2016 3:33:00 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		6/21/2016 3:33:00 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1