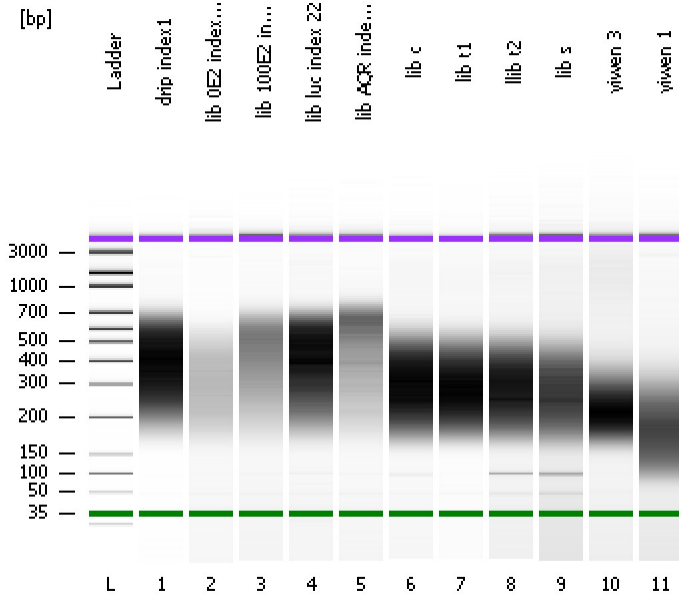


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
Modified: 3/21/2014 12:06:23 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

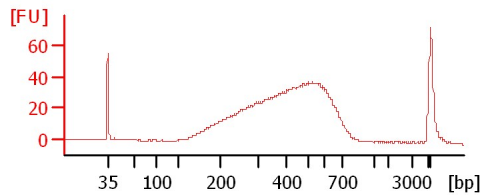
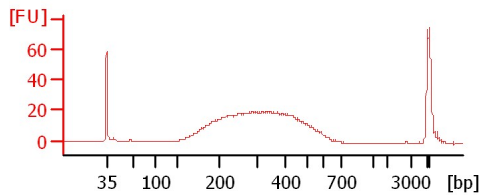
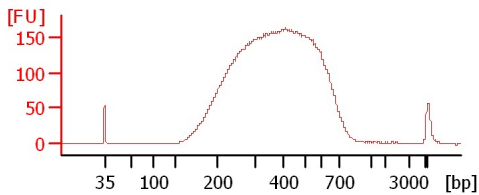
Chip Information:

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

drip index 1

lib 0E2 index 10

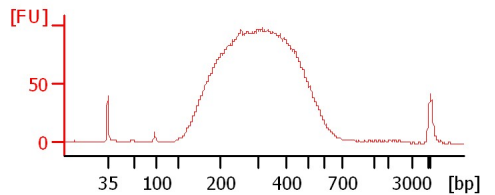
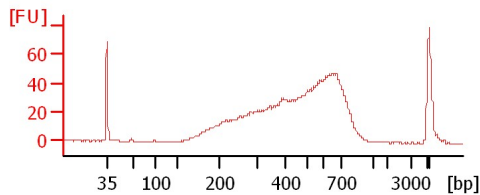
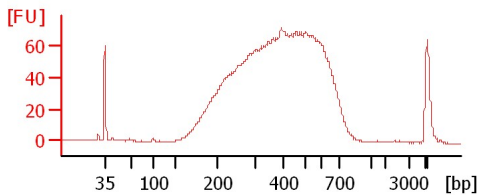
lib 100E2 index 11



lib luc index 22

lib AQR index 23

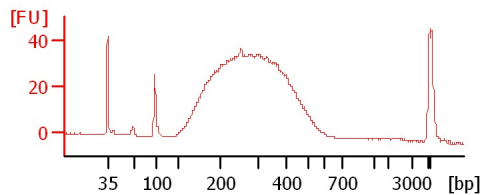
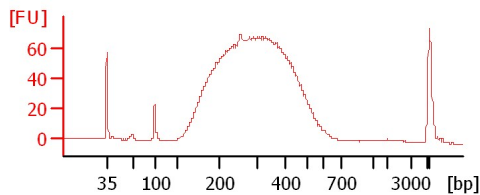
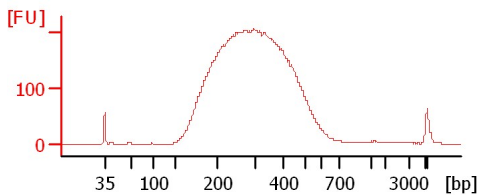
lib c



lib t1

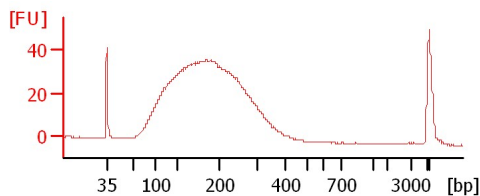
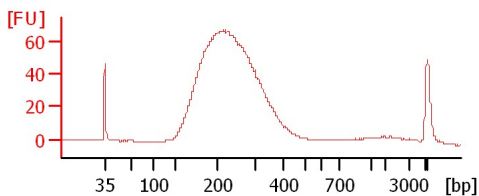
lib t2

lib s



yiwen 3

yiwen 1



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
Modified: 3/21/2014 12:06:23 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
drip index1		<input type="checkbox"/>	✓			
lib 0E2 index 10		<input type="checkbox"/>	✓			
lib 100E2 index 11		<input type="checkbox"/>	✓			
lib luc index 22		<input type="checkbox"/>	✓			
lib AQR index 23		<input type="checkbox"/>	✓			
lib c		<input type="checkbox"/>	✓			
lib t1		<input type="checkbox"/>	✓			
llib t2		<input type="checkbox"/>	✓			
lib s		<input type="checkbox"/>	✓			
yiwen 3		<input type="checkbox"/>	✓			
yiwen 1		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
Modified: 3/21/2014 12:06:23 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

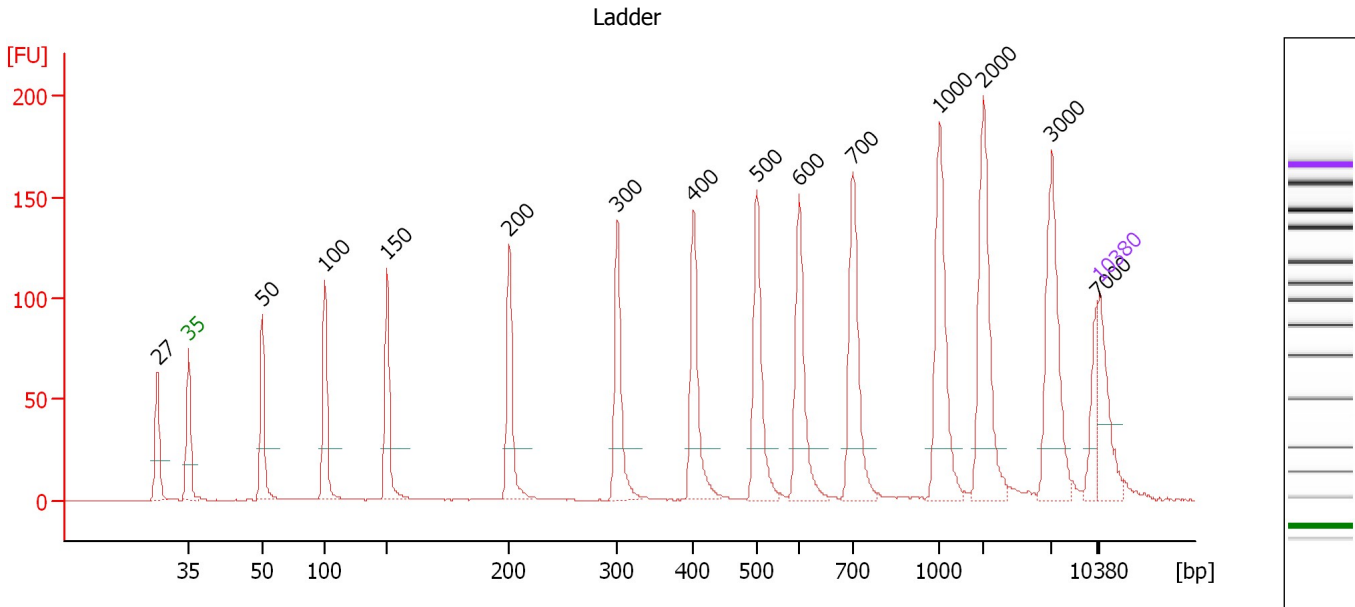
Ladder

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:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

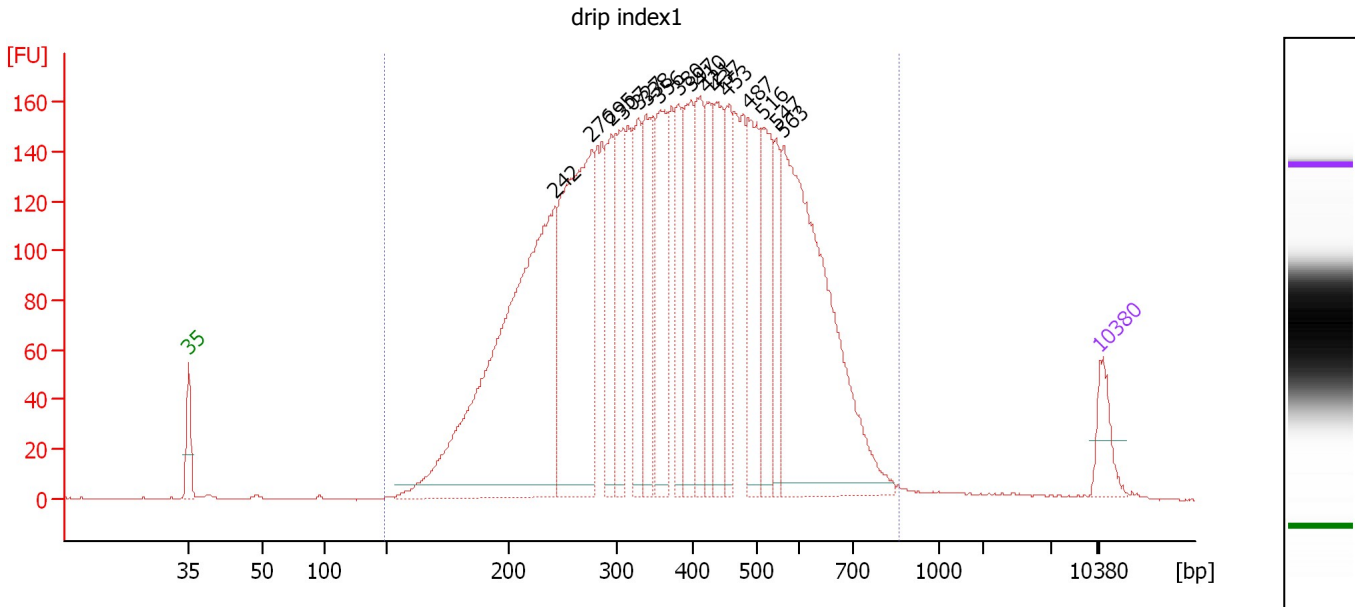
Peak table for Ladder

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	40.61	27	0.00	0.0	
2	43.00	35	125.00	5,411.3	Lower Marker
3	48.68	50	150.00	4,545.5	Ladder Peak
4	53.45	100	150.00	2,272.7	Ladder Peak
5	58.23	150	150.00	1,515.2	Ladder Peak
6	67.64	200	150.00	1,136.4	Ladder Peak
7	75.93	300	150.00	757.6	Ladder Peak
8	81.78	400	150.00	568.2	Ladder Peak
9	86.65	500	150.00	454.5	Ladder Peak
10	89.89	600	150.00	378.8	Ladder Peak
11	94.04	700	150.00	324.7	Ladder Peak
12	100.70	1,000	150.00	227.3	Ladder Peak
13	104.04	2,000	150.00	113.6	Ladder Peak
14	109.31	3,000	150.00	75.8	Ladder Peak
15	112.73	7,000	150.00	32.5	Ladder Peak
16	113.00	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : drip index1

Number of peaks found: 17 Corr. Area 1: 4,995.1
 Noise: 0.2

Peak table for sample 1 : drip index1

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	71.13	242	1,861.37	11,647.3	
3	73.95	276	1,038.78	5,699.9	
4	75.49	295	272.40	1,400.4	
5	76.33	307	254.34	1,256.0	
6	77.52	327	298.80	1,384.0	
7	78.13	338	313.55	1,407.1	
8	79.19	356	339.22	1,445.0	
9	80.60	380	218.73	872.7	
10	81.61	397	289.23	1,103.8	
11	82.27	410	273.06	1,009.1	
12	82.80	421	209.33	753.6	
13	83.59	437	265.13	918.9	
14	84.34	453	231.24	774.2	
15	86.01	487	312.54	972.5	
16	87.16	516	296.55	871.3	
17	88.17	547	155.66	431.2	
18	88.70	563	1,122.73	3,020.5	
19	113.00	10,380	75.00	10.9	Upper Marker

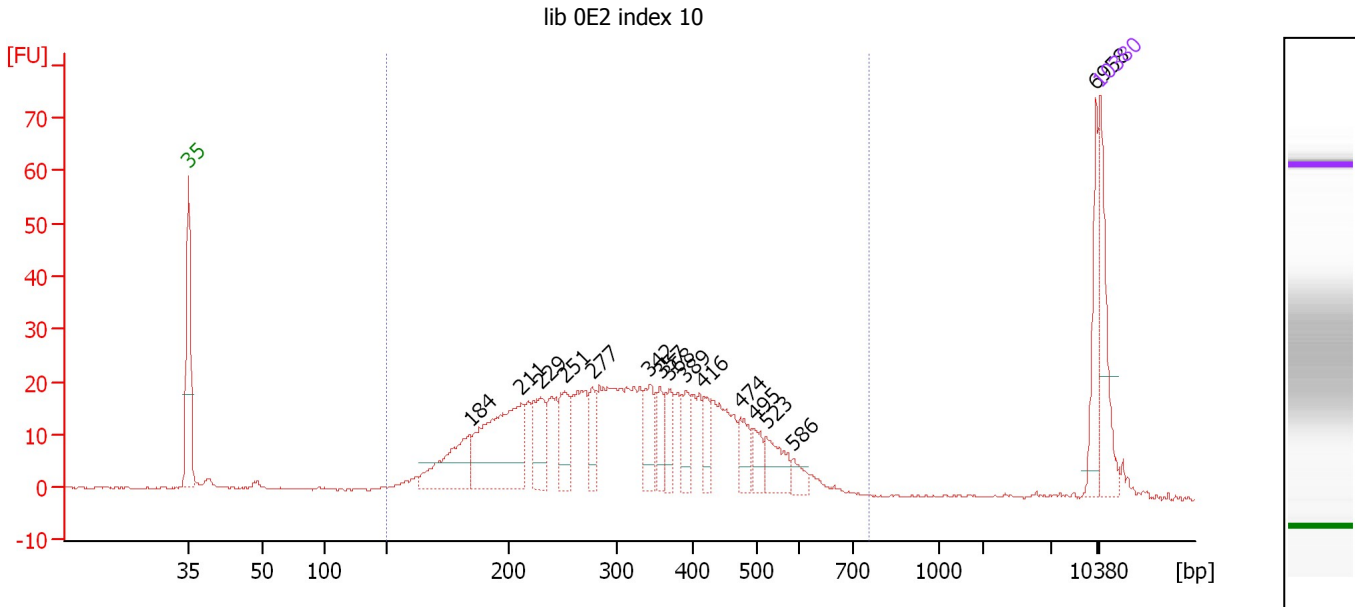
Region table for sample 1 : drip index1

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
148	857	393	43,562.2	9,213.23	4,995.1	99	36.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : lib OE2 index 10

Number of peaks found: 15 Corr. Area 1: 623.6
 Noise: 0.2

Peak table for sample 2 : lib OE2 index 10

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	64.60	184	110.97	914.5	
3	68.58	211	229.34	1,644.4	
4	70.06	229	74.43	491.9	
5	71.90	251	63.14	380.5	
6	74.04	277	38.86	212.4	
7	78.42	342	58.64	259.4	
8	79.25	357	38.13	162.0	
9	79.90	368	34.46	141.9	
10	81.13	389	35.19	137.2	
11	82.57	416	28.13	102.4	
12	85.37	474	35.37	113.1	
13	86.42	495	26.19	80.1	
14	87.38	523	45.47	131.9	
15	89.43	586	16.75	43.3	
16	112.69	6,958	55.00	12.0	
17	113.00	10,380	75.00	10.9	Upper Marker

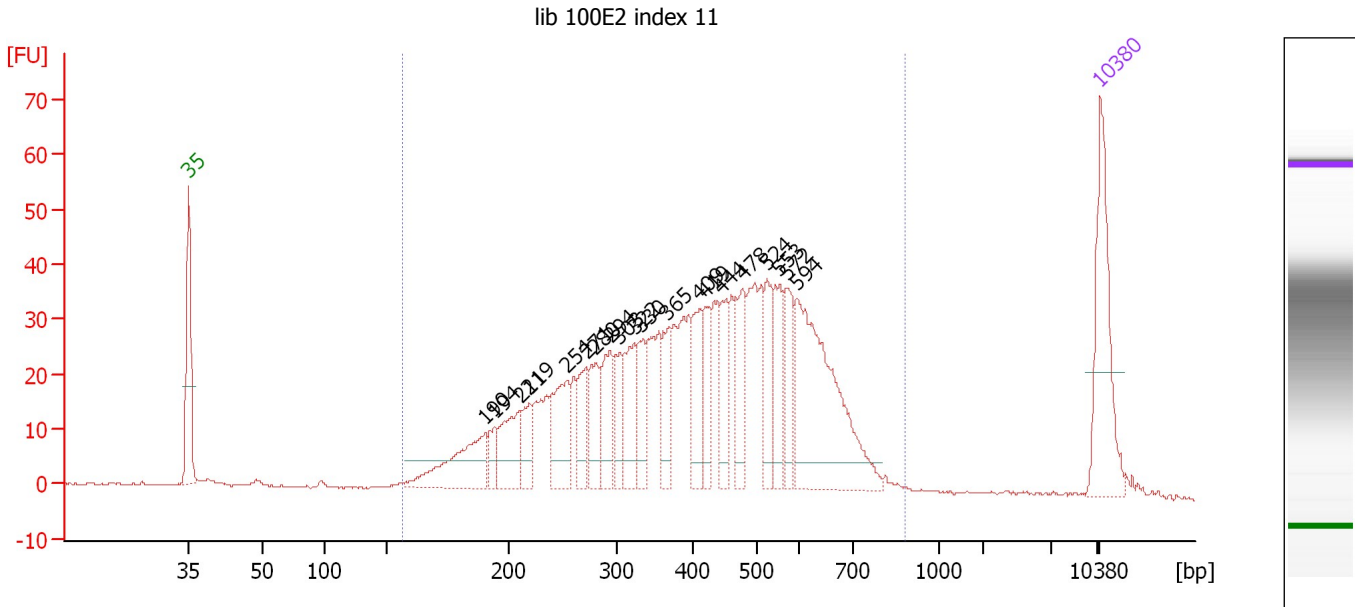
Region table for sample 2 : lib OE2 index 10

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. for Area	% of Total	Size distribution in CV [%]
150	752	327	8,588.3	1,551.88	623.6	92	35.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : lib 100E2 index 11

Number of peaks found: 20 Corr. Area 1: 1,020.6
 Noise: 0.2

Peak table for sample 3 : lib 100E2 index 11

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	65.75	190	88.25	703.8	
3	66.51	194	21.85	170.6	
4	68.52	211	59.99	431.6	
5	69.19	219	31.98	221.6	
6	72.09	254	67.56	403.5	
7	73.52	271	38.35	214.5	
8	74.23	280	42.02	227.8	
9	75.39	294	50.50	260.7	
10	76.10	303	32.30	161.5	
11	77.22	322	50.84	239.2	
12	77.67	330	42.94	197.3	
13	79.76	365	38.28	158.7	
14	82.22	409	45.01	166.8	
15	82.71	419	35.62	128.8	
16	83.91	444	43.82	149.6	
17	85.56	478	44.68	141.7	
18	87.44	524	39.62	114.5	
19	88.37	553	44.92	123.0	
20	89.00	572	40.57	107.4	
21	89.71	594	178.00	453.7	
22	113.00	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary Continued ...

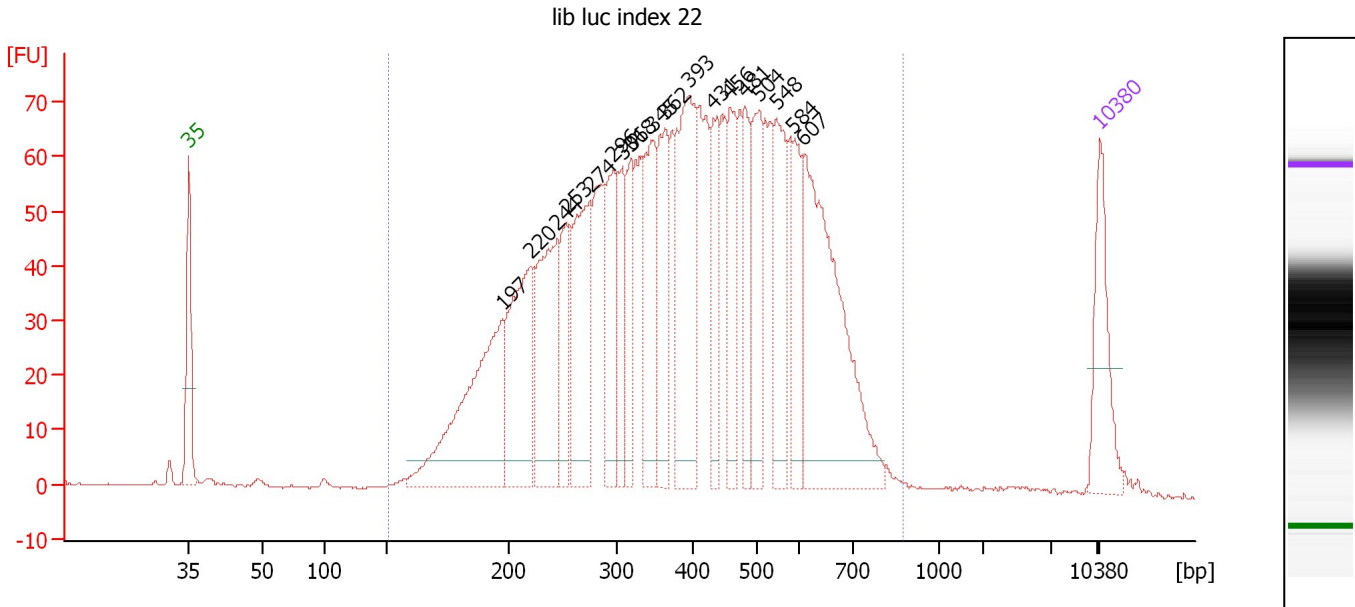
... Region table for sample 3 : lib 100E2 index 11

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
156	880	420	6,730.2	1,493.53	1,020.6	97	36.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : lib luc index 22

Number of peaks found: 18 Corr. Area 1: 2,157.7
 Noise: 0.2

Peak table for sample 4 : lib luc index 22

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	67.12	197	319.00	2,450.3	
3	69.31	220	219.22	1,509.0	
4	71.31	244	202.85	1,258.2	
5	71.99	253	101.44	608.7	
6	73.81	274	199.68	1,102.3	
7	75.63	296	129.31	660.9	
8	76.27	306	79.67	394.7	
9	76.95	318	79.22	378.1	
10	78.59	345	145.69	638.9	
11	79.55	362	128.09	536.4	
12	81.37	393	222.19	856.9	
13	83.28	431	87.03	306.1	
14	84.51	456	99.04	329.1	
15	85.74	481	80.02	251.9	
16	86.78	504	117.44	352.9	
17	88.20	548	117.29	324.5	
18	89.38	584	83.11	215.6	
19	90.20	607	321.62	802.3	
20	113.00	10,380	75.00	10.9	Upper Marker

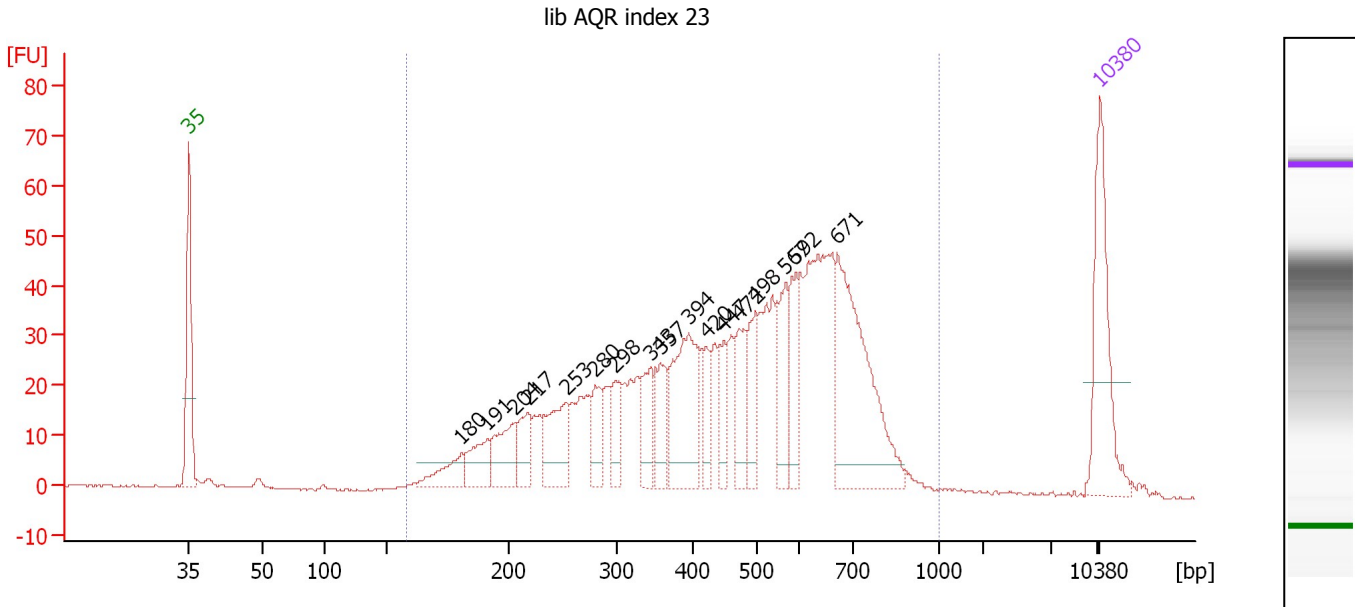
Region table for sample 4 : lib luc index 22

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
151	873	404	16,590.9	3,553.38	2,157.7	98	37.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : lib AQR index 23

Number of peaks found: 17 Corr. Area 1: 1,108.0
 Noise: 0.2

Peak table for sample 5 : lib AQR index 23

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	63.93	180	32.28	271.3	
3	65.96	191	38.83	307.9	
4	68.00	204	50.46	374.2	
5	69.06	217	36.37	253.7	
6	72.03	253	65.43	391.9	
7	74.25	280	35.54	192.5	
8	75.73	298	31.06	158.1	
9	78.46	343	36.51	161.1	
10	79.25	357	37.47	159.1	
11	81.43	394	104.01	400.1	
12	82.77	420	25.46	91.8	
13	84.06	447	25.89	87.8	
14	85.27	472	40.67	130.7	
15	86.56	498	42.61	129.6	
16	88.83	567	47.75	127.5	
17	89.62	592	42.21	108.1	
18	92.81	671	165.00	372.8	
19	113.00	10,380	75.00	10.9	Upper Marker

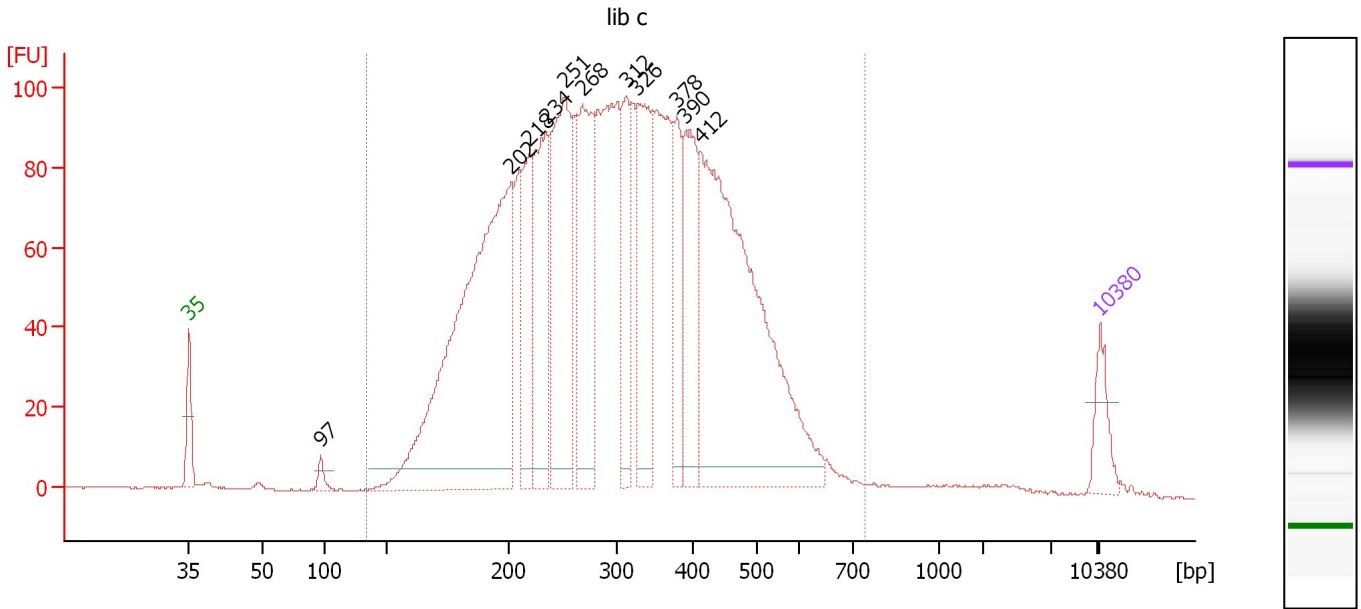
Region table for sample 5 : lib AQR index 23

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
158	1,000	476	5,817.8	1,403.13	1,108.0	98	37.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : lib c

Number of peaks found: 11 Corr. Area 1: 2,950.4
 Noise: 0.1

Peak table for sample 6 : lib c

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	53.15	97	31.43	491.4	
3	67.78	202	1,736.27	13,047.0	
4	69.14	218	320.35	2,225.6	
5	70.45	234	486.77	3,152.3	
6	71.87	251	732.56	4,422.4	
7	73.28	268	510.50	2,886.2	
8	76.61	312	255.70	1,242.9	
9	77.46	326	423.74	1,968.5	
10	80.52	378	209.90	840.6	
11	81.17	390	343.75	1,336.9	
12	82.35	412	1,357.21	4,996.0	
13	113.00	10,380	75.00	10.9	Upper Marker

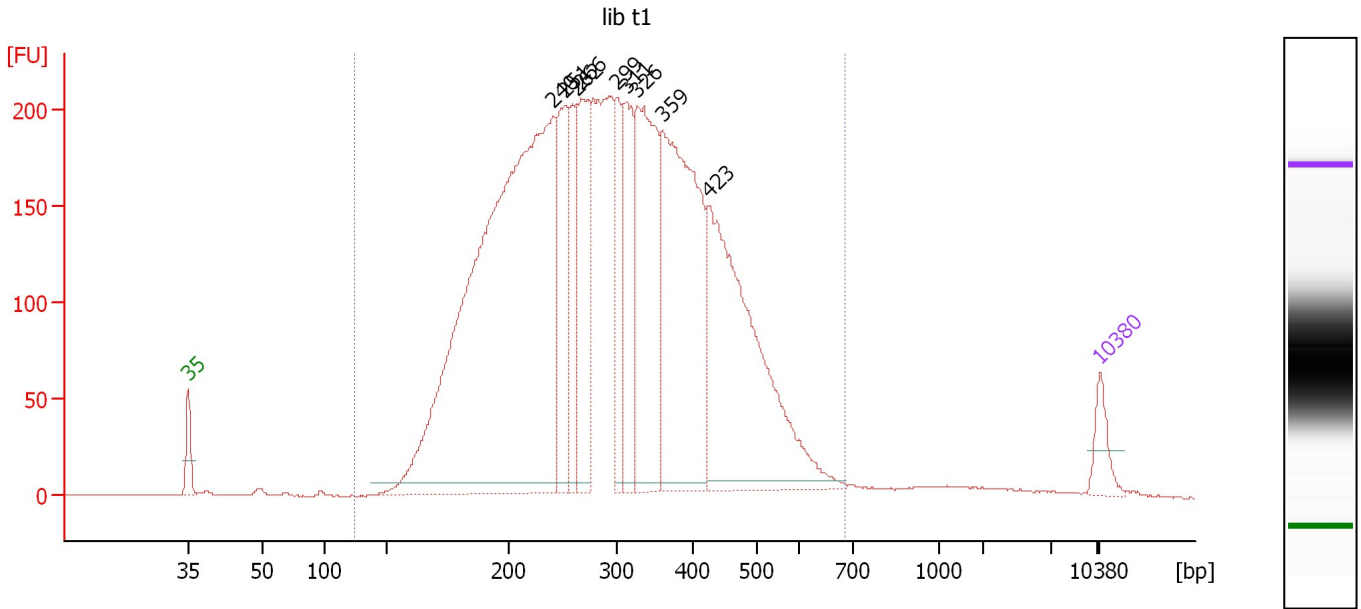
Region table for sample 6 : lib c

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
133	744	320	47,324.0	8,442.64	2,950.4	98	35.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : lib t1

Number of peaks found: 9 Corr. Area 1: 5,890.8
 Noise: 0.2

Peak table for sample 7 : lib t1

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	70.99	240	3,922.56	24,719.5	
3	71.90	251	531.80	3,205.1	
4	72.76	262	341.33	1,975.5	
5	73.10	266	579.29	3,301.8	
6	75.82	299	317.99	1,612.8	
7	76.59	311	423.21	2,060.0	
8	77.45	326	906.14	4,211.6	
9	79.41	359	1,345.44	5,670.7	
10	82.90	423	1,327.76	4,755.5	
11	113.00	10,380	75.00	10.9	Upper Marker

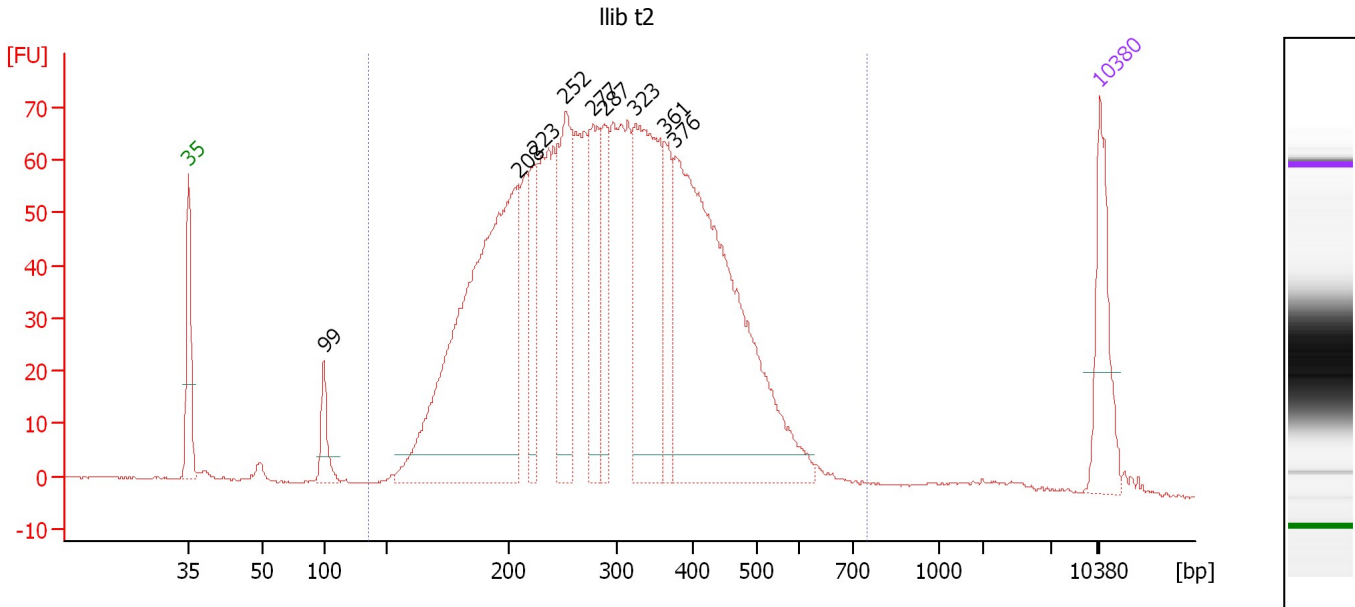
Region table for sample 7 : lib t1

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
125	686	310	63,712.9	11,175.61	5,890.8	98	34.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : lib t2

Number of peaks found: 9 Corr. Area 1: 1,951.5
 Noise: 0.2

Peak table for sample 8 : lib t2

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	53.37	99	49.39	754.6	
3	68.28	208	737.14	5,377.0	
4	69.58	223	101.81	690.4	
5	71.95	252	198.65	1,194.6	
6	74.02	277	144.82	792.2	
7	74.89	287	92.38	486.9	
8	77.30	323	323.10	1,513.6	
9	79.52	361	86.27	361.8	
10	80.39	376	633.00	2,549.7	
11	113.00	10,380	75.00	10.9	Upper Marker

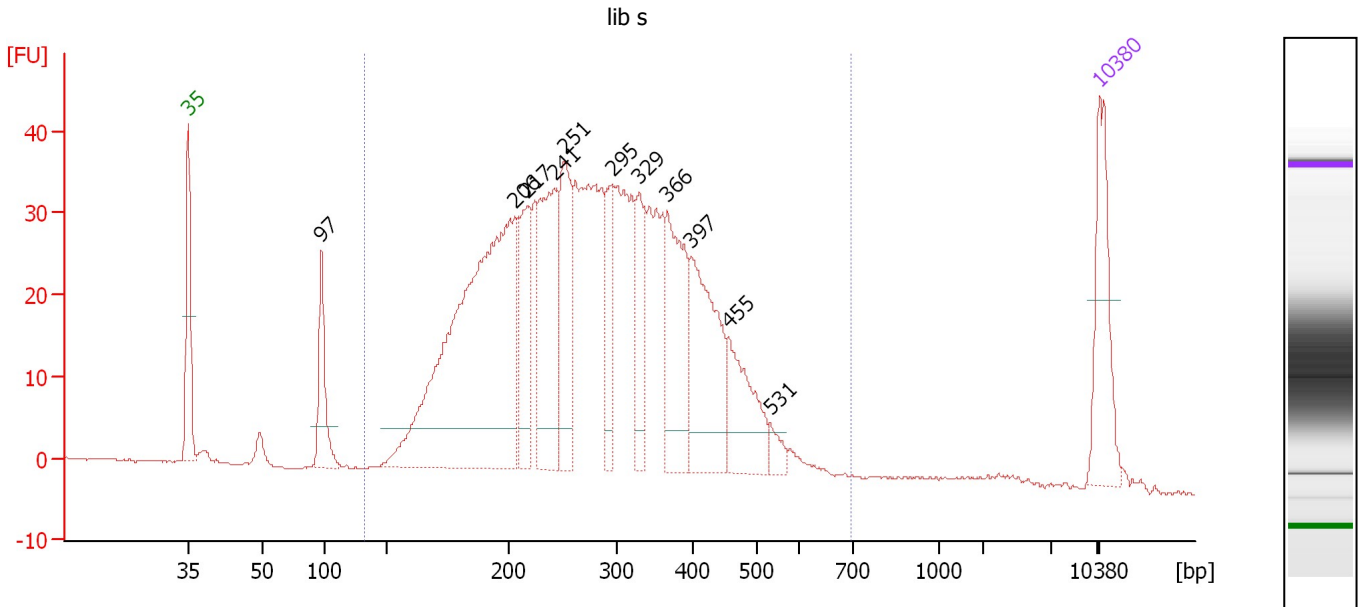
Region table for sample 8 : lib t2

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
135	748	309	18,041.4	3,161.81	1,951.5	97	33.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : lib s

Number of peaks found: 11 Corr. Area 1: 1,011.9
 Noise: 0.1

Peak table for sample 9 : lib s

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	53.18	97	81.06	1,264.0	
3	68.11	206	639.37	4,709.9	
4	69.03	217	103.01	719.9	
5	71.07	241	183.35	1,151.0	
6	71.84	251	138.58	837.5	
7	75.53	295	73.37	376.6	
8	77.61	329	74.95	345.4	
9	79.79	366	143.81	595.3	
10	81.59	397	159.93	610.9	
11	84.45	455	88.83	296.0	
12	87.65	531	14.28	40.8	
13	113.00	10,380	75.00	10.9	Upper Marker

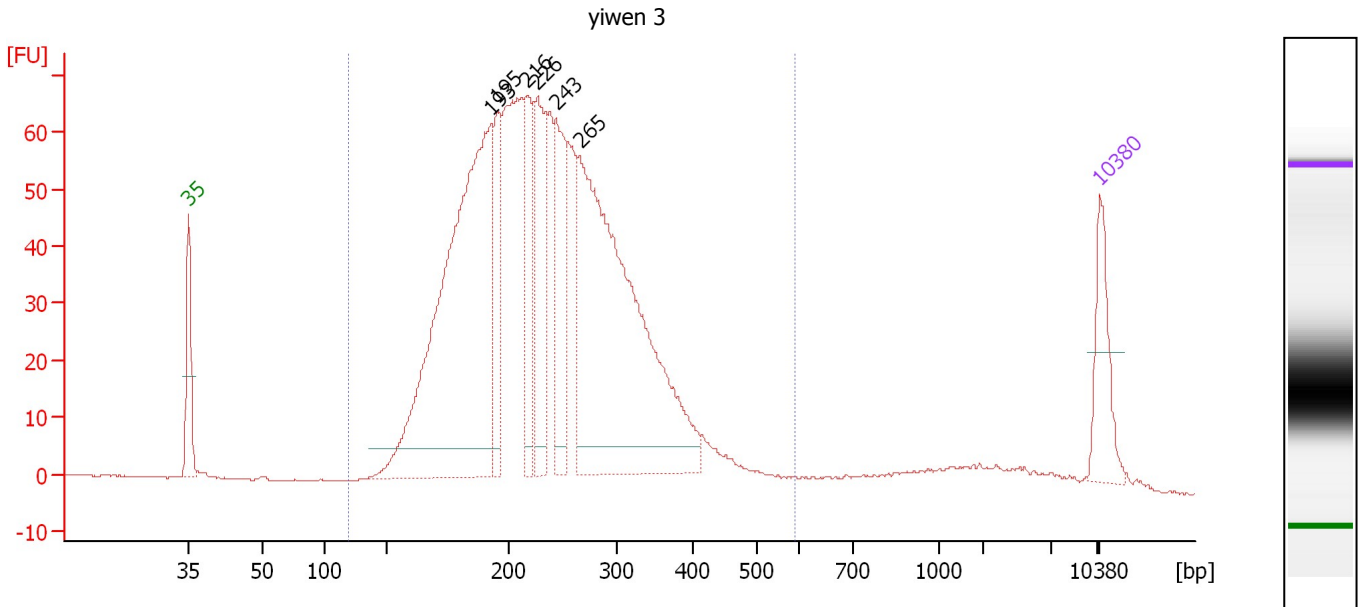
Region table for sample 9 : lib s

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
132	696	295	14,382.0	2,418.63	1,011.9	94	33.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : yiwen 3

Number of peaks found: 6 Corr. Area 1: 1,458.2
 Noise: 0.1

Peak table for sample 10 : yiwen 3

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	66.25	193	1,013.45	7,971.4	
3	66.69	195	153.69	1,194.4	
4	68.94	216	151.67	1,065.5	
5	69.77	226	212.99	1,429.9	
6	71.23	243	199.59	1,242.6	
7	73.04	265	909.62	5,197.3	
8	113.00	10,380	75.00	10.9	Upper Marker

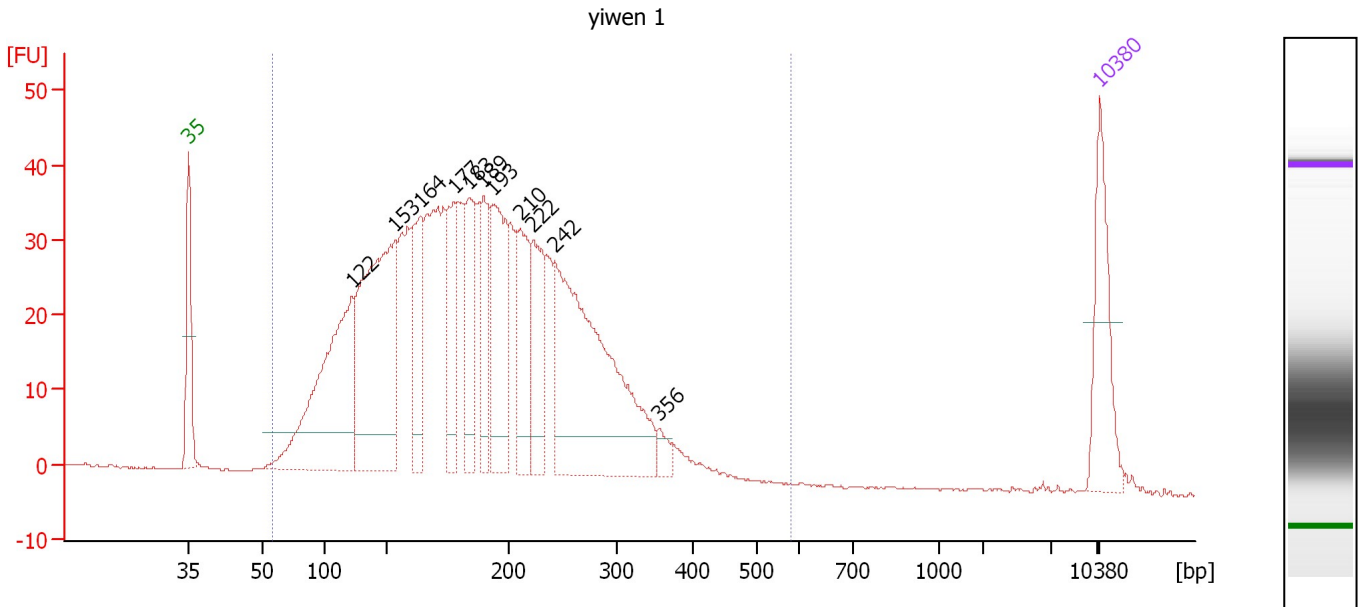
Region table for sample 10 : yiwen 3

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
119	588	245	24,025.9	3,556.15	1,458.2	95	28.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : yiwen 1

Number of peaks found: 11 Corr. Area 1: 1,130.7
 Noise: 0.1

Peak table for sample 11 : yiwen 1

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	43.00	35	125.00	5,411.3	Lower Marker
2	55.59	122	334.55	4,141.8	
3	58.85	153	395.63	3,910.3	
4	60.82	164	124.00	1,147.1	
5	63.39	177	110.14	940.6	
6	64.52	183	109.12	901.3	
7	65.66	189	99.28	793.9	
8	66.35	193	162.43	1,274.2	
9	68.47	210	108.54	782.9	
10	69.46	222	115.12	785.8	
11	71.14	242	423.42	2,648.7	
12	79.18	356	16.28	69.4	
13	113.00	10,380	75.00	10.9	Upper Marker

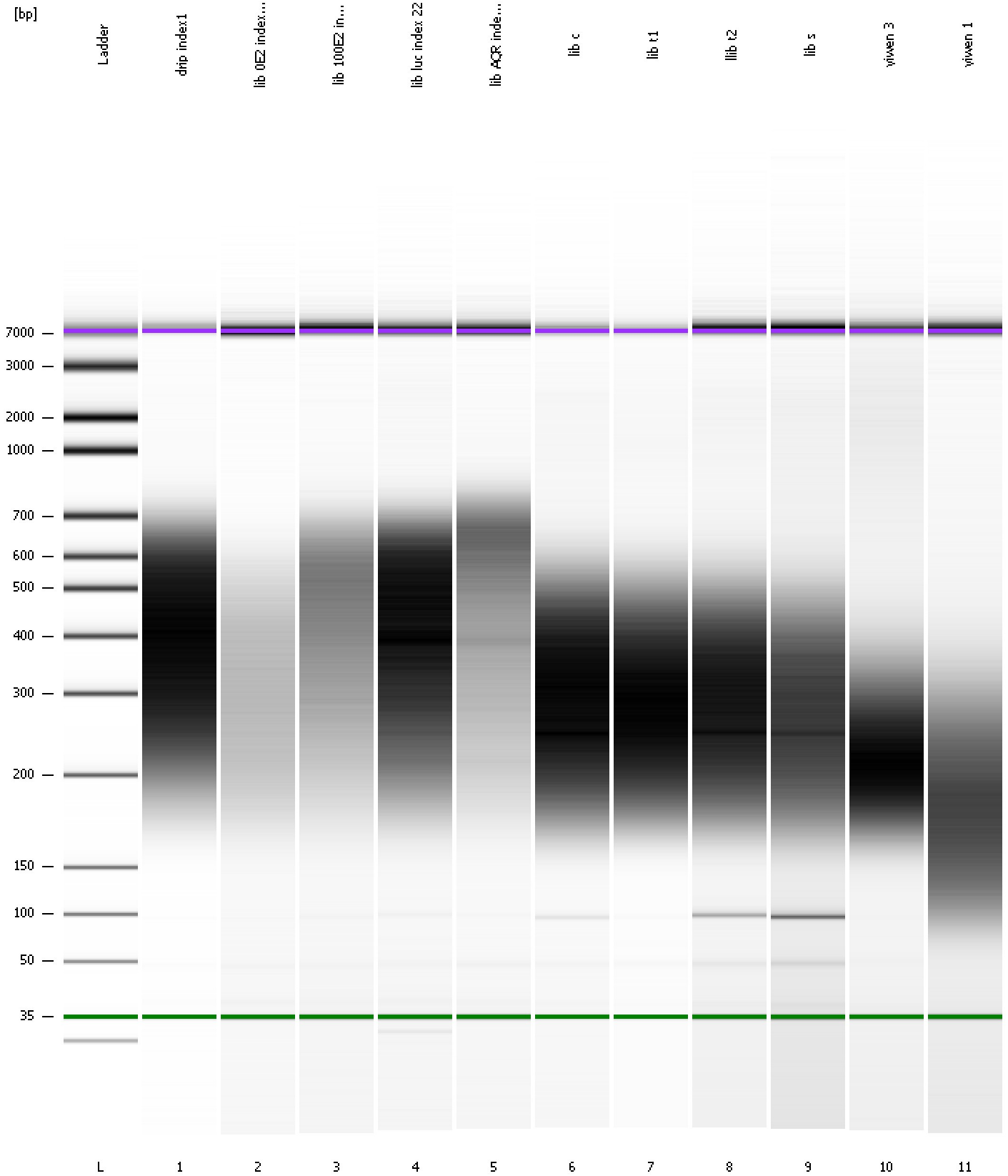
Region table for sample 11 : yiwen 1

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
57	579	197	24,793.3	2,782.58	1,130.7	99	32.8

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
Modified: 3/21/2014 12:06:23 PM

Gel Image

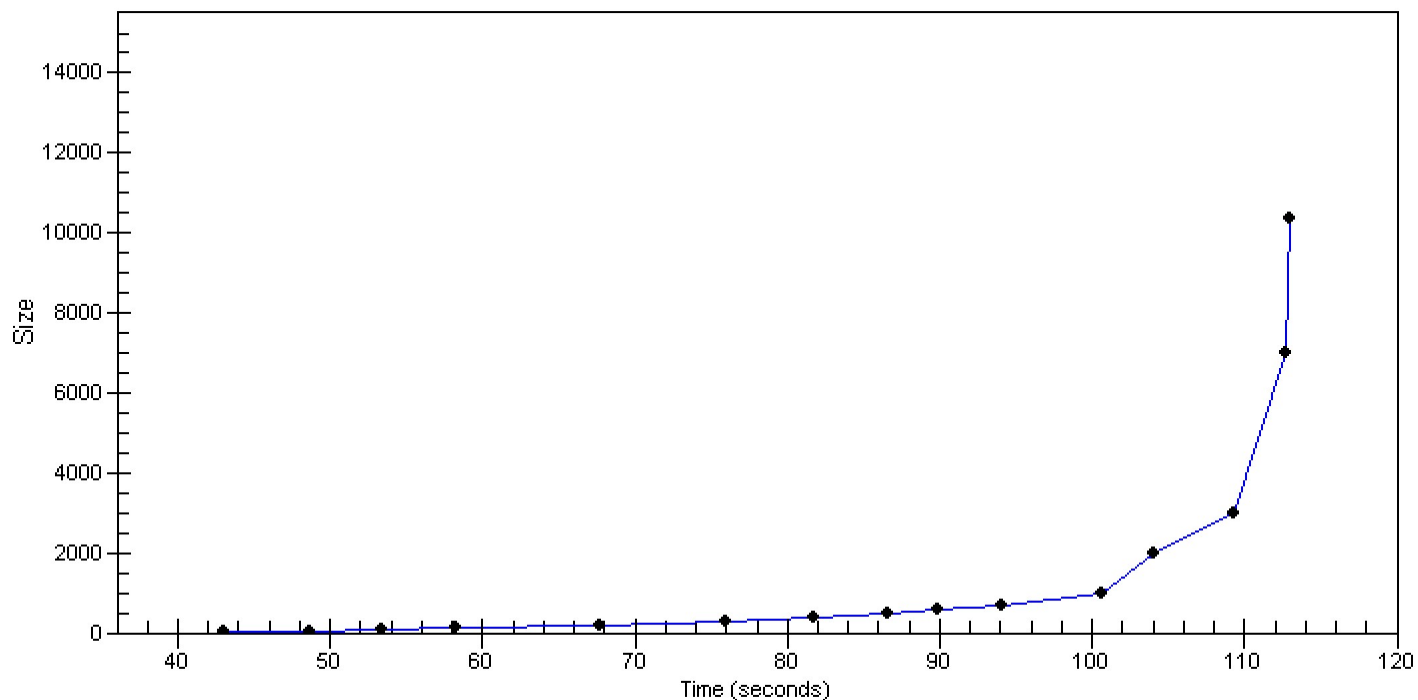


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
Modified: 3/21/2014 12:06:23 PM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad

Created: 3/21/2014 11:25:14 AM
 Modified: 3/21/2014 12:06:23 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		3/21/2014 12:05:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2014-03-21\2014-03-21_002.xad)		Instrument	Run		3/21/2014 11:25:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		3/21/2014 11:25:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		3/21/2014 11:25:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		3/21/2014 11:25:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		3/21/2014 11:25:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		3/21/2014 11:25:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		3/21/2014 11:25:20 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1