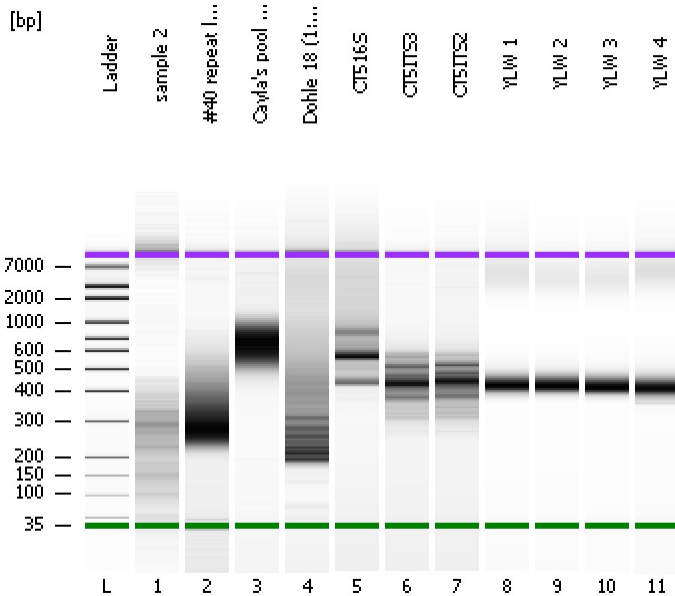


Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**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

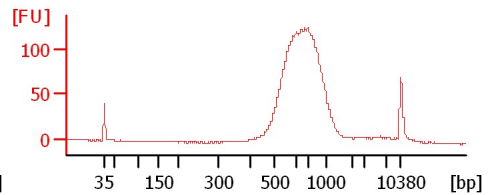
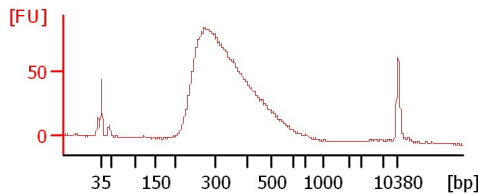
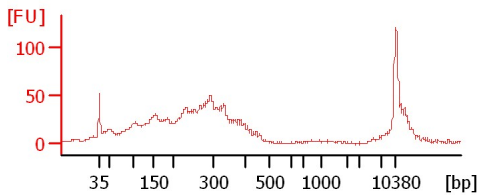
Chip Information:

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

sample 2

#40 repeat library (1:3)

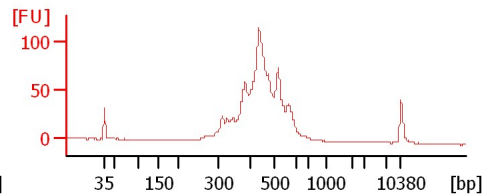
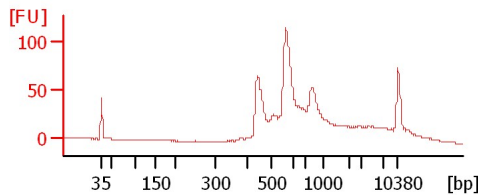
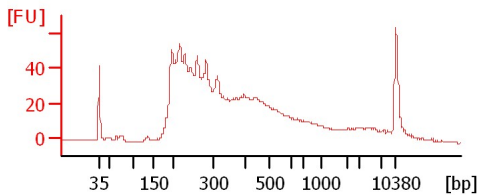
Cayla's pool (1:6)



Dohle 18 (1:10)

CT5165

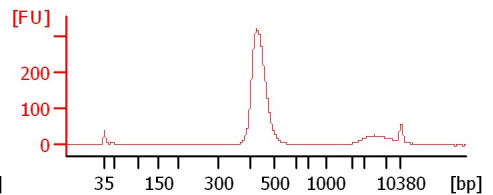
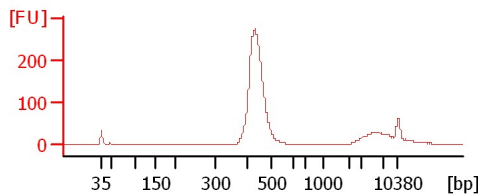
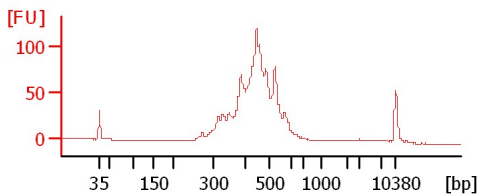
CT51TS3



CT51TS2

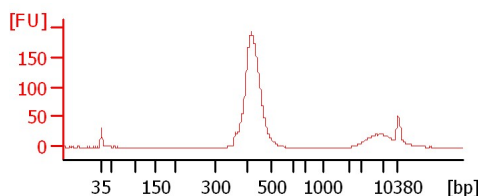
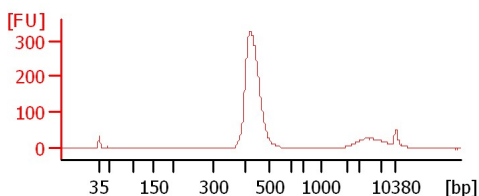
YLW 1

YLW 2



YLW 3

YLW 4



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
Modified: 12/9/2015 11:40:51 AM

**Electrophoresis File Run Summary (Chip Summary)**

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
sample 2		<input type="checkbox"/>	✓			
#40 repeat library (1:3)		<input type="checkbox"/>	✓			
Cayla's pool (1:6)		<input type="checkbox"/>	✓			
Dohle 18 (1:10)		<input type="checkbox"/>	✓			
CT516S		<input type="checkbox"/>	✓			
CT5ITS3		<input type="checkbox"/>	✓			
CT5ITS2		<input type="checkbox"/>	✓			
YLW 1		<input type="checkbox"/>	✓			
YLW 2		<input type="checkbox"/>	✓			
YLW 3		<input type="checkbox"/>	✓			
YLW 4		<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\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
Modified: 12/9/2015 11:40:51 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

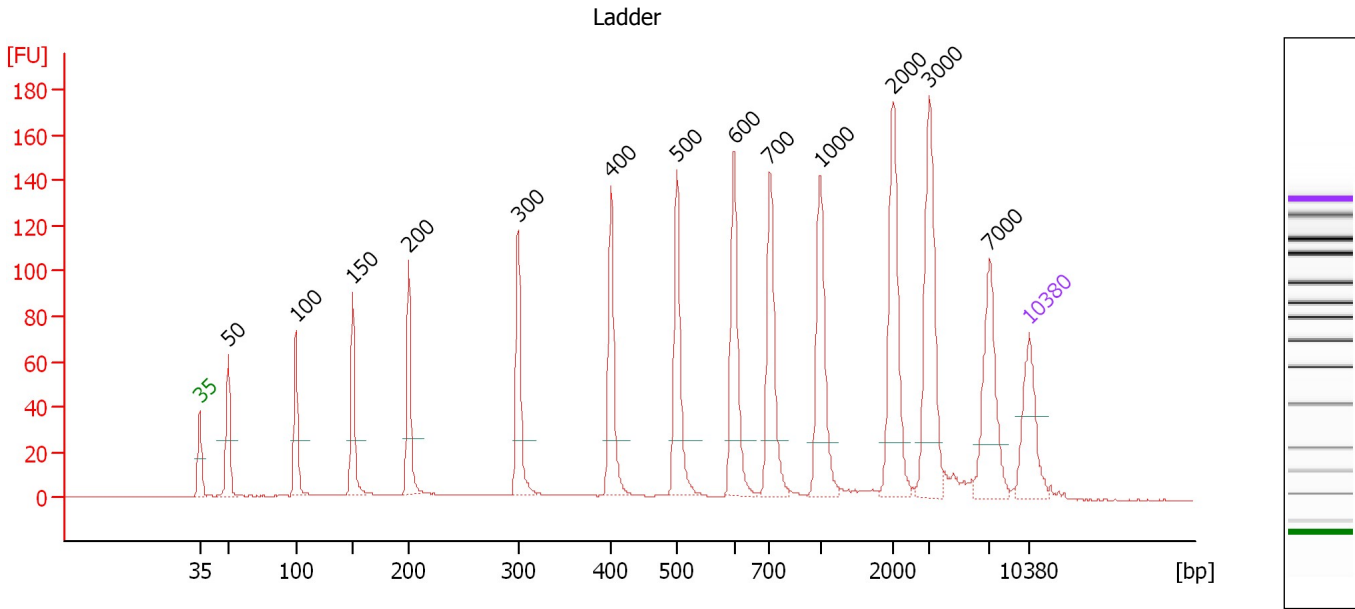
**Ladder**

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: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.2

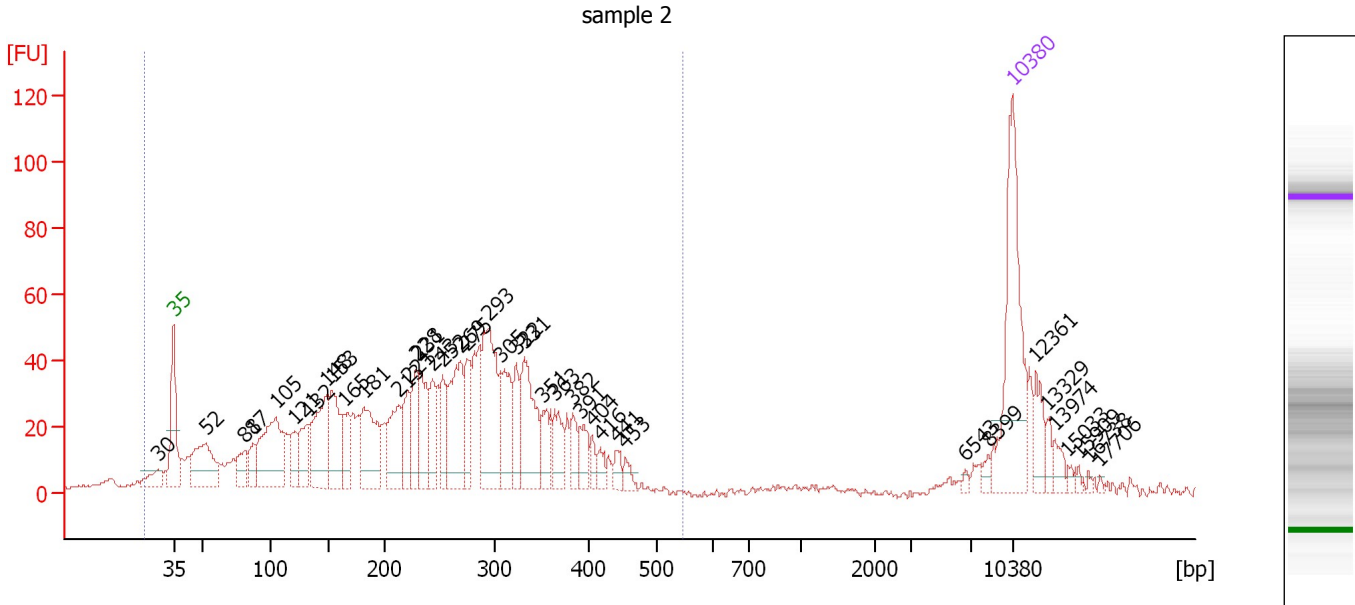
**Peak table for Ladder**

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	45.38
3	100	150.00	2,272.7	Ladder Peak	51.10
4	150	150.00	1,515.2	Ladder Peak	55.91
5	200	150.00	1,136.4	Ladder Peak	60.63
6	300	150.00	757.6	Ladder Peak	69.86
7	400	150.00	568.2	Ladder Peak	77.69
8	500	150.00	454.5	Ladder Peak	83.26
9	600	150.00	378.8	Ladder Peak	88.08
10	700	150.00	324.7	Ladder Peak	91.09
11	1,000	150.00	227.3	Ladder Peak	95.37
12	2,000	150.00	113.6	Ladder Peak	101.48
13	3,000	150.00	75.8	Ladder Peak	104.54
14	7,000	150.00	32.5	Ladder Peak	109.63
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : sample 2**

Number of peaks found: 40                      Corr. Area 1: 1,454.3  
 Noise: 0.6

**Peak table for sample 1 : sample 2**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	30	0.00	0.0		41.67
2	35	125.00	5,411.3	Lower Marker	43.00
3	52	59.47	1,717.1		45.67
4	81	21.78	405.1		48.98
5	87	17.21	298.0		49.67
6	105	89.29	1,291.6		51.55
7	121	19.76	248.4		53.07
8	132	32.23	368.7		54.22
9	148	66.05	675.2		55.74
10	153	52.21	516.8		56.20
11	165	26.50	243.7		57.30
12	181	60.12	502.4		58.87
13	213	43.74	311.5		61.81
14	221	25.78	176.6		62.59
15	228	34.76	230.8		63.24
16	231	40.38	264.6		63.51
17	243	29.73	185.2		64.62
18	252	25.89	155.6		65.44
19	269	63.78	359.8		66.96
20	275	28.85	159.2		67.51
21	293	82.04	424.2		69.22
22	305	42.78	212.3		70.27
23	322	23.57	110.8		71.61
24	331	57.97	265.2		72.30
25	351	19.86	85.7		73.86

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**

... Peak table for sample 1 : sample 2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	363	19.37	80.9		74.78
27	382	13.18	52.2		76.30
28	391	16.43	63.7		76.99
29	404	7.67	28.8		77.91
30	416	7.40	26.9		78.60
31	441	7.97	27.4		79.98
32	453	6.50	21.7		80.67
33	6,543	1.71	0.4		109.04
34	8,399	3.47	0.6		111.02
35	10,380	75.00	10.9	Upper Marker	113.00
36	12,361	0.00	0.0		114.98
37	13,329	0.00	0.0		115.94
38	13,974	0.00	0.0		116.59
39	15,033	0.00	0.0		117.65
40	15,909	0.00	0.0		118.52
41	16,738	0.00	0.0		119.35
42	17,706	0.00	0.0		120.31

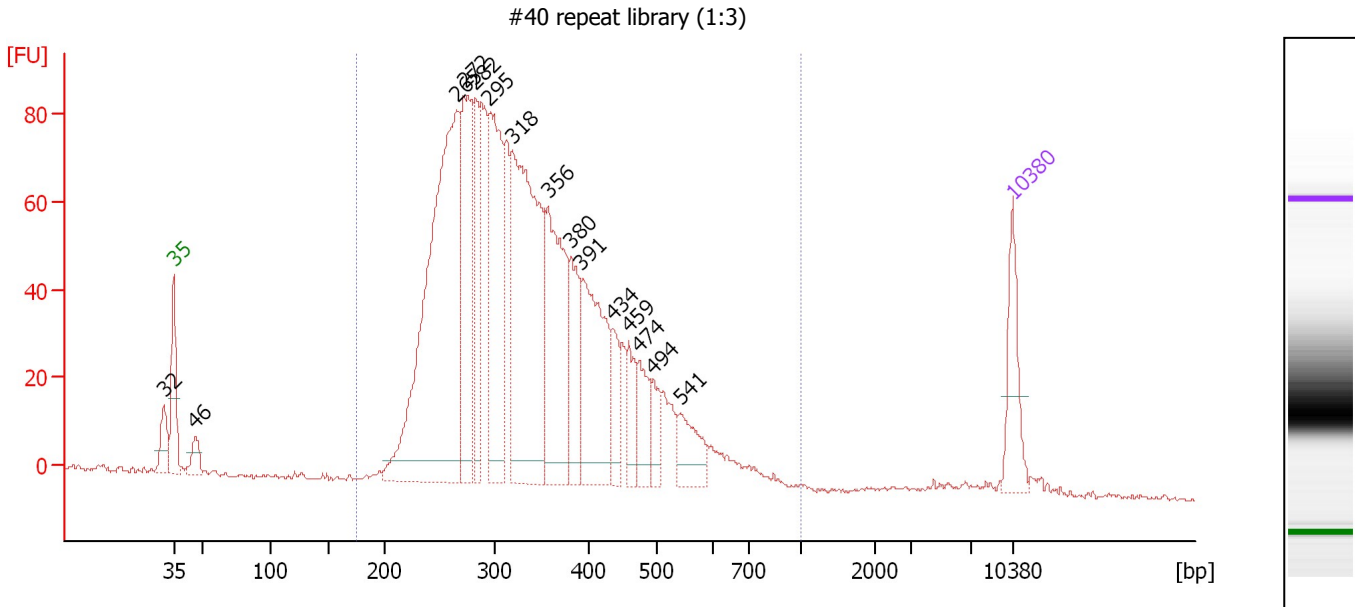
Region table for sample 1 : sample 2

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
20	547	231	1,454.3	14,319.8	1,308.65	91	45.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : #40 repeat library (1:3)**

Number of peaks found: 15      Corr. Area 1: 1,791.2  
 Noise: 0.5

**Peak table for sample 2 : #40 repeat library (1:3)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.22
2	35	125.00	5,411.3	Lower Marker	43.00
3	46	41.99	1,373.3		44.80
4	265	883.07	5,056.1		66.59
5	272	293.21	1,632.5		67.29
6	282	168.78	906.4		68.21
7	295	330.90	1,701.5		69.37
8	318	593.76	2,829.9		71.26
9	356	291.89	1,241.5		74.26
10	380	123.94	493.6		76.15
11	391	266.17	1,031.3		76.98
12	434	61.17	213.7		79.57
13	459	61.80	204.2		80.96
14	474	67.87	216.8		81.83
15	494	41.03	125.8		82.94
16	541	71.92	201.3		85.25
17	10,380	75.00	10.9	Upper Marker	113.00

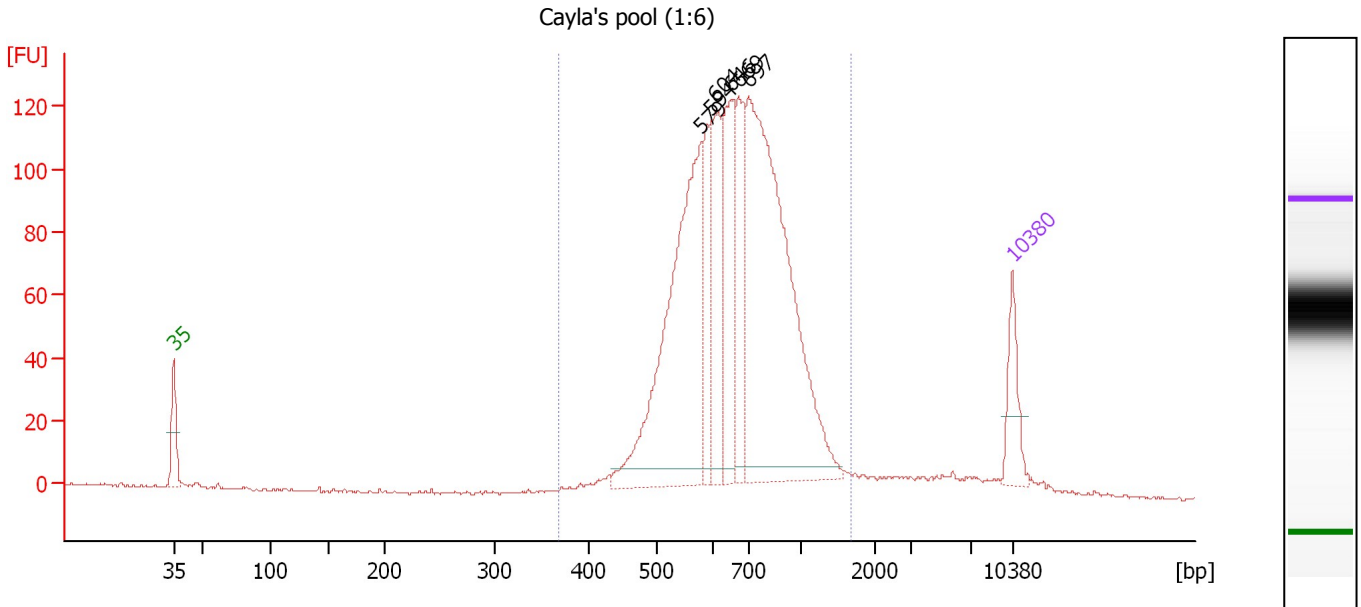
**Region table for sample 2 : #40 repeat library (1:3)**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
175	1,000	346	1,791.2	17,711.3	3,699.60	97	28.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : Cayla's pool (1:6)**

Number of peaks found: 6                      Corr. Area 1: 1,533.1  
 Noise: 0.3

**Peak table for sample 3 : Cayla's pool (1:6)**

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	579	706.24	1,848.2		87.07
3	594	144.60	369.1		87.77
4	604	211.72	531.4		88.19
5	646	240.31	564.1		89.45
6	669	188.01	426.0		90.15
7	697	980.88	2,133.3		90.99
8	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 3 : Cayla's pool (1:6)**

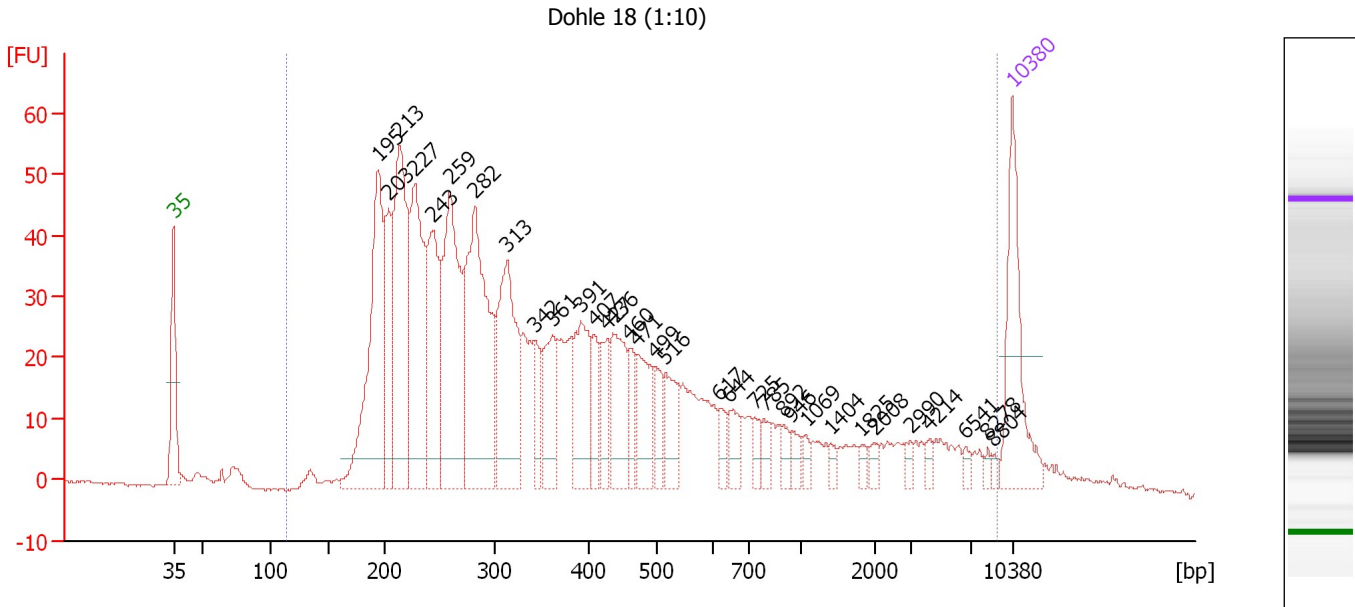
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
367	1,678	704	1,533.1	6,035.3	2,609.40	94	27.3



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : Dohle 18 (1:10)**

Number of peaks found: 33                      Corr. Area 1: 1,487.6  
 Noise: 0.3

**Peak table for sample 4 : Dohle 18 (1:10)**

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	195	235.04	1,830.7		60.12
3	203	87.21	650.3		60.93
4	213	217.48	1,546.9		61.84
5	227	187.41	1,251.1		63.12
6	243	119.31	743.9		64.60
7	259	204.99	1,199.1		66.08
8	282	241.41	1,298.1		68.18
9	313	155.05	751.4		70.85
10	342	32.29	142.8		73.18
11	361	63.42	266.3		74.61
12	391	78.46	303.9		77.00
13	407	31.94	118.8		78.10
14	427	35.18	124.8		79.19
15	436	68.41	238.0		79.67
16	460	26.27	86.5		81.05
17	471	55.37	177.9		81.67
18	499	23.54	71.5		83.20
19	516	39.17	114.9		84.06
20	617	12.50	30.7		88.59
21	644	19.66	46.3		89.40
22	725	11.07	23.1		91.45
23	785	12.65	24.4		92.31
24	892	11.39	19.3		93.83
25	946	10.58	16.9		94.59

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**

**... Peak table for sample 4 : Dohle 18 (1:10)**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	1,069	6.86	9.7		95.79
27	1,404	5.41	5.8		97.84
28	1,825	5.37	4.5		100.41
29	2,008	6.14	4.6		101.51
30	2,990	5.46	2.8		104.51
31	4,214	5.61	2.0		106.09
32	6,541	4.81	1.1		109.04
33	8,278	3.66	0.7		110.90
34	8,804	3.29	0.6		111.43
35	10,380	75.00	10.9	Upper Marker	113.00

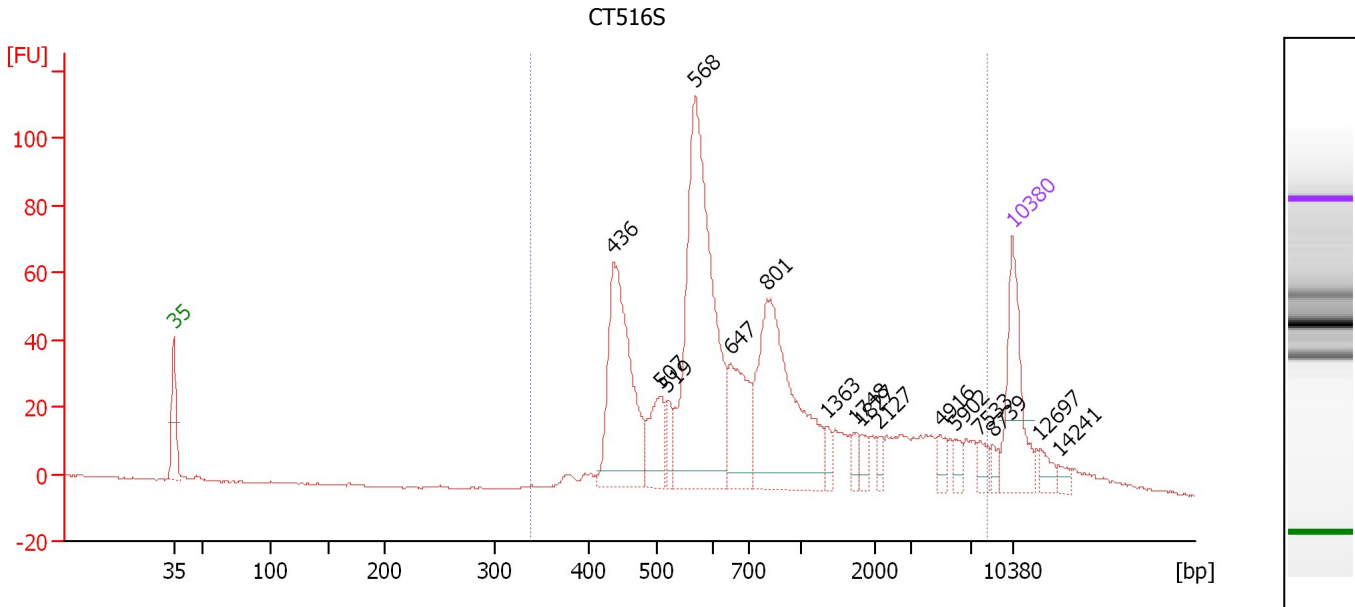
**Region table for sample 4 : Dohle 18 (1:10)**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
114	9,175	738	1,487.6	12,005.7	2,403.32	97	100.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : CT516S**

Number of peaks found: 16                      Corr. Area 1: 1,123.3  
 Noise: 0.2

**Peak table for sample 5 : CT516S**

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	436	219.34	761.7		79.71
3	507	55.71	166.4		83.61
4	519	21.65	63.2		84.19
5	568	375.11	1,000.1		86.55
6	647	85.77	200.9		89.49
7	801	237.49	449.4		92.53
8	1,363	11.10	12.3		97.58
9	1,748	11.67	10.1		99.94
10	1,827	11.23	9.3		100.43
11	2,127	8.69	6.2		101.87
12	4,916	12.53	3.9		106.98
13	5,902	10.39	2.7		108.23
14	7,533	11.80	2.4		110.16
15	8,739	8.12	1.4		111.36
16	10,380	75.00	10.9	Upper Marker	113.00
17	12,697	0.00	0.0		115.31
18	14,241	0.00	0.0		116.85

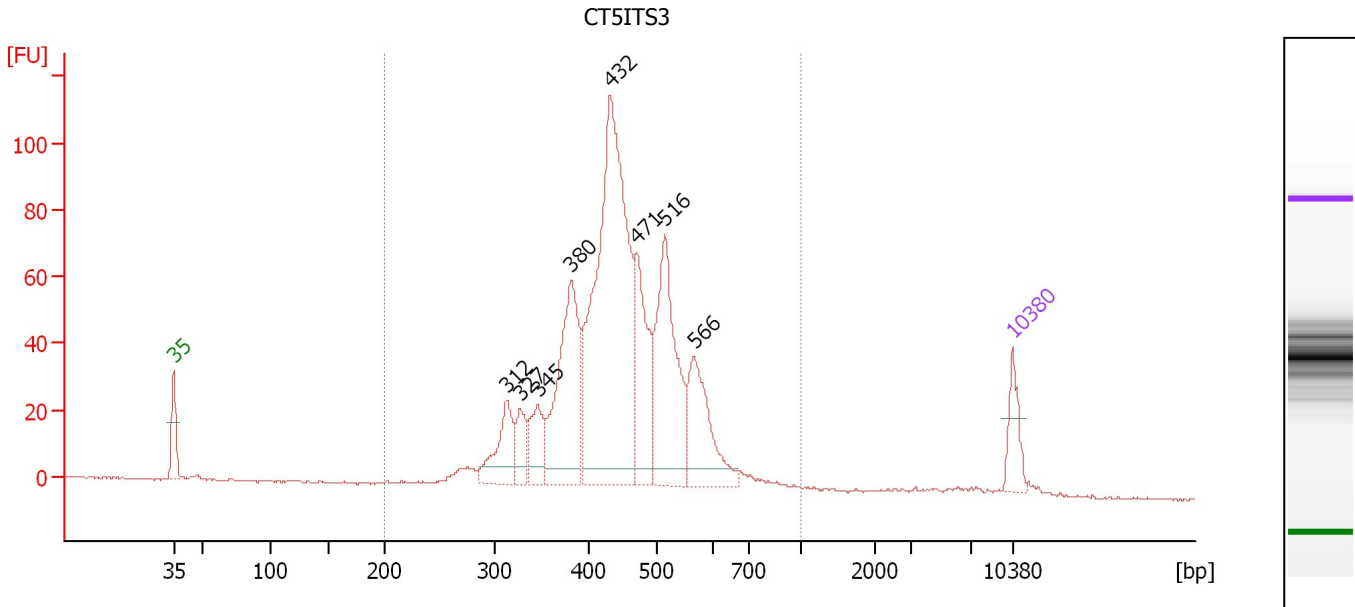
**Region table for sample 5 : CT516S**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
338	8,189	1,292	1,123.3	2,685.3	1,190.59	93	100.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : CT5ITS3**

Number of peaks found: 8                      Corr. Area 1: 1,152.8  
 Noise: 0.3

**Peak table for sample 6 : CT5ITS3**

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	312	190.82	926.4		70.80
3	327	96.47	446.9		71.98
4	345	132.03	579.4		73.40
5	380	500.15	1,992.8		76.14
6	432	1,312.14	4,602.4		79.47
7	471	318.10	1,022.3		81.67
8	516	504.61	1,482.5		84.02
9	566	272.05	728.8		86.42
10	10,380	75.00	10.9	Upper Marker	113.00

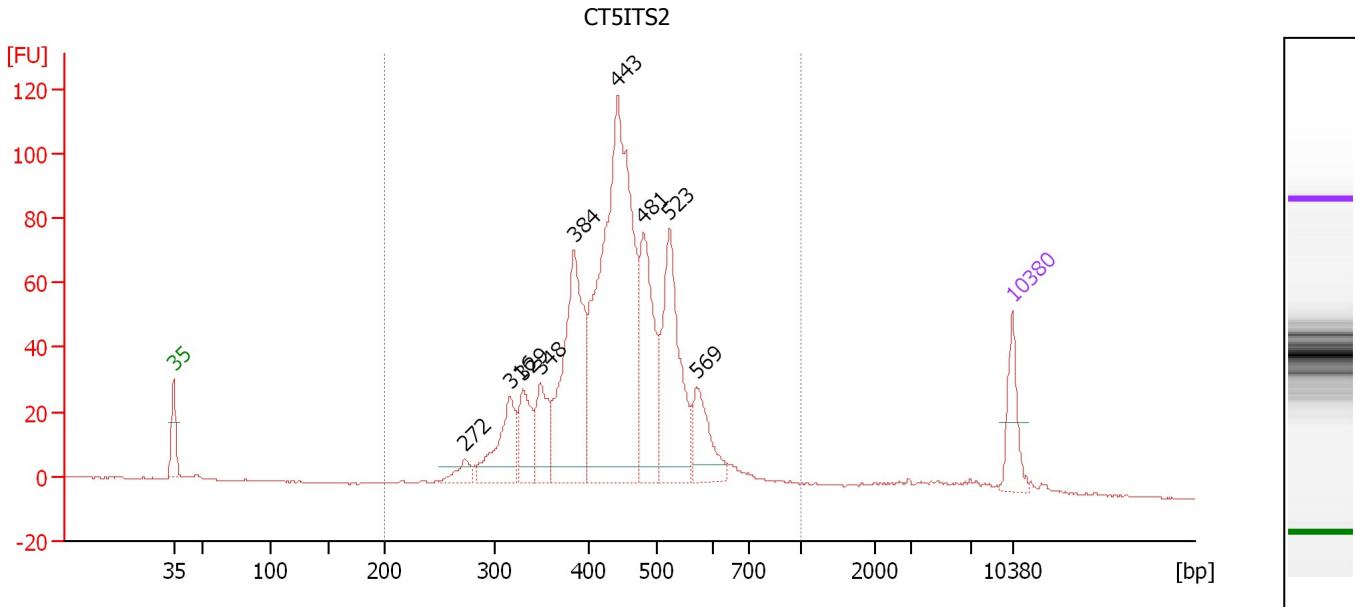
**Region table for sample 6 : CT5ITS3**

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	449	1,152.8	12,395.9	3,495.11	98	19.4

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : CT5ITS2**

Number of peaks found: 9                      Corr. Area 1: 1,236.4  
 Noise: 0.2

**Peak table for sample 7 : CT5ITS2**

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	272	40.99	228.5		67.26
3	316	173.99	835.3		71.08
4	329	119.08	548.5		72.12
5	348	120.66	525.4		73.61
6	384	452.23	1,784.1		76.44
7	443	1,035.87	3,546.5		80.06
8	481	290.50	915.5		82.19
9	523	345.53	1,000.8		84.37
10	569	122.39	325.6		86.61
11	10,380	75.00	10.9	Upper Marker	113.00

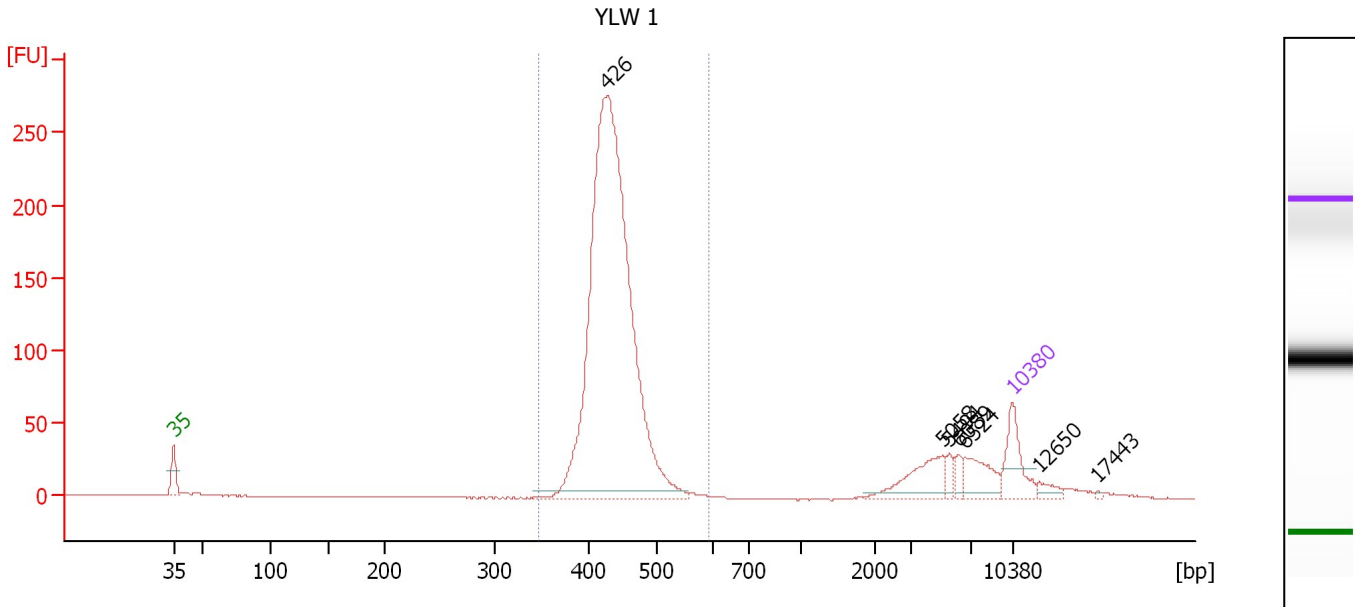
**Region table for sample 7 : CT5ITS2**

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	448	1,236.4	10,363.7	2,901.28	95	20.4

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : YLW 1**

Number of peaks found: 7                      Corr. Area 1: 1,334.9  
 Noise: 0.2

**Peak table for sample 8 : YLW 1**

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	426	1,701.13	6,057.4		79.11
3	5,058	92.96	27.8		107.16
4	5,494	17.30	4.8		107.71
5	6,089	18.29	4.6		108.47
6	6,524	65.09	15.1		109.02
7	10,380	75.00	10.9	Upper Marker	113.00
8	12,650	0.00	0.0		115.27
9	17,443	0.00	0.0		120.05

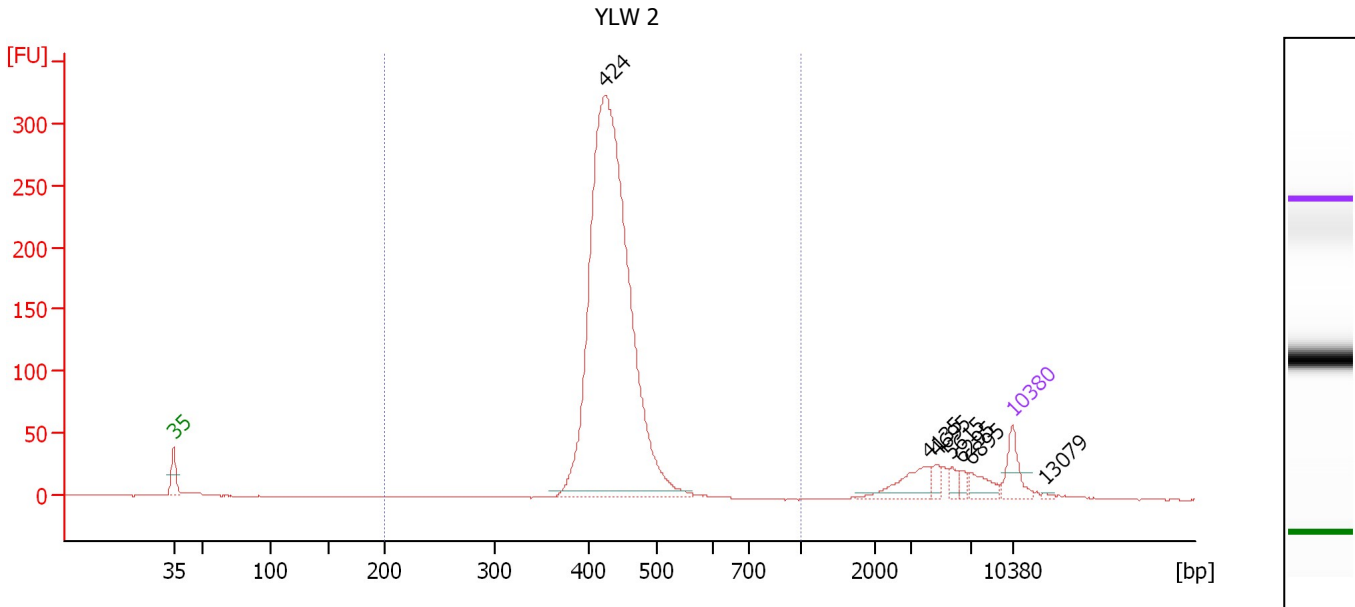
**Region table for sample 8 : YLW 1**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
347	592	435	1,334.9	5,858.0	1,677.69	84	7.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : YLW 2**

Number of peaks found: 7                      Corr. Area 1: 1,560.6  
 Noise: 0.3

**Peak table for sample 9 : YLW 2**

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	424	2,715.85	9,697.5		79.04
3	4,135	100.91	37.0		105.98
4	4,695	24.67	8.0		106.70
5	5,615	28.18	7.6		107.87
6	6,295	17.82	4.3		108.73
7	6,895	51.03	11.2		109.49
8	10,380	75.00	10.9	Upper Marker	113.00
9	13,079	0.00	0.0		115.69

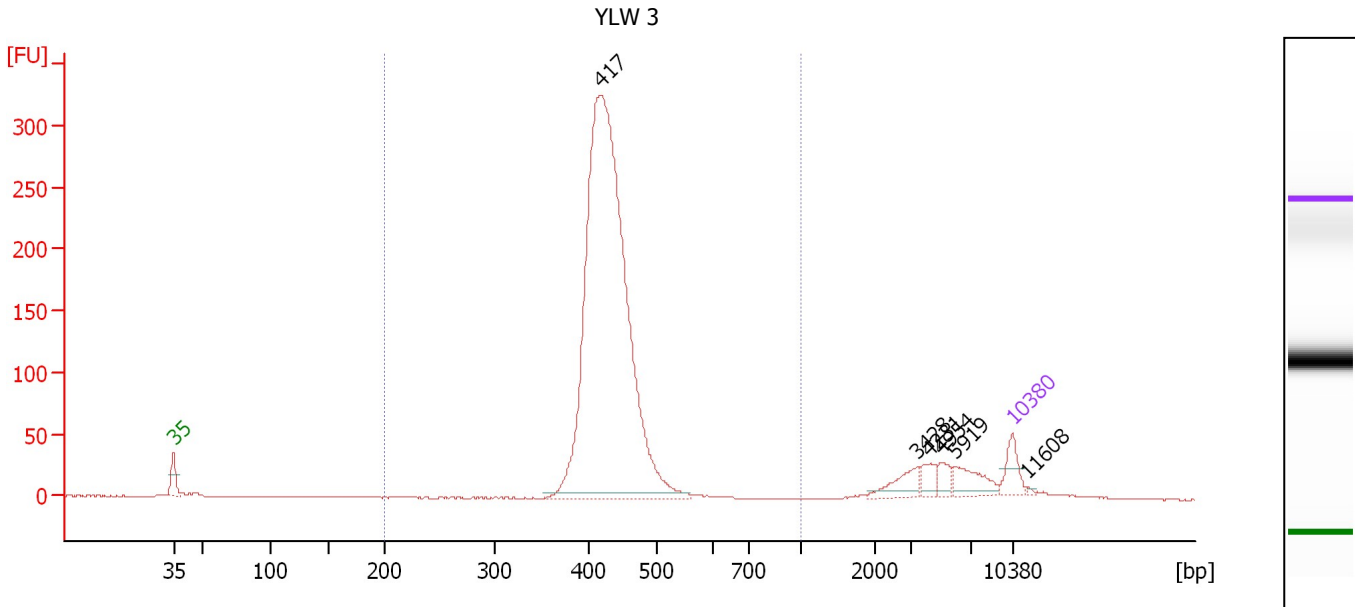
**Region table for sample 9 : YLW 2**

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	435	1,560.6	9,462.3	2,708.24	88	7.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : YLW 3**

Number of peaks found: 6                      Corr. Area 1: 1,595.4  
 Noise: 0.2

**Peak table for sample 10 : YLW 3**

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	417	3,383.42	12,291.0		78.64
3	3,428	86.71	38.3		105.09
4	4,231	52.74	18.9		106.11
5	4,954	46.49	14.2		107.03
6	5,919	100.52	25.7		108.25
7	10,380	75.00	10.9	Upper Marker	113.00
8	11,608	0.00	0.0		114.23

**Region table for sample 10 : YLW 3**

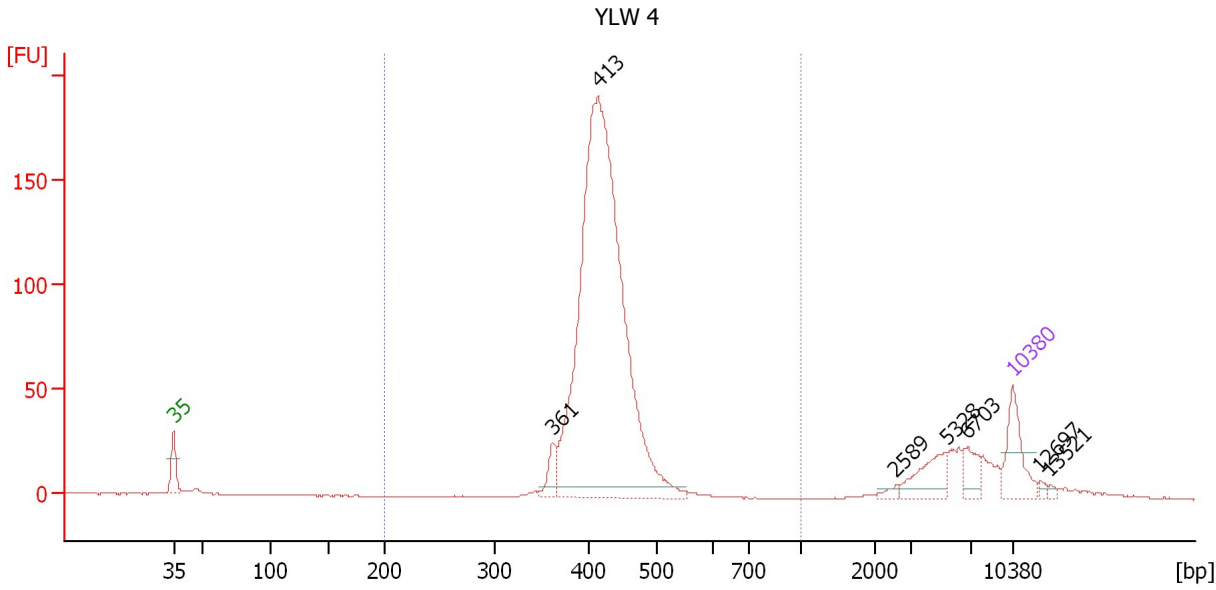
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	430	1,595.4	11,889.9	3,361.51	87	7.3



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 AM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : YLW 4**

Number of peaks found: 7                      Corr. Area 1: 1,002.0  
 Noise: 0.2

**Peak table for sample 11 : YLW 4**

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	361	46.69	195.8		74.66
3	413	1,607.10	5,895.6		78.41
4	2,589	9.15	5.4		103.29
5	5,328	66.87	19.0		107.50
6	6,703	35.98	8.1		109.25
7	10,380	75.00	10.9	Upper Marker	113.00
8	12,697	0.00	0.0		115.31
9	13,521	0.00	0.0		116.14

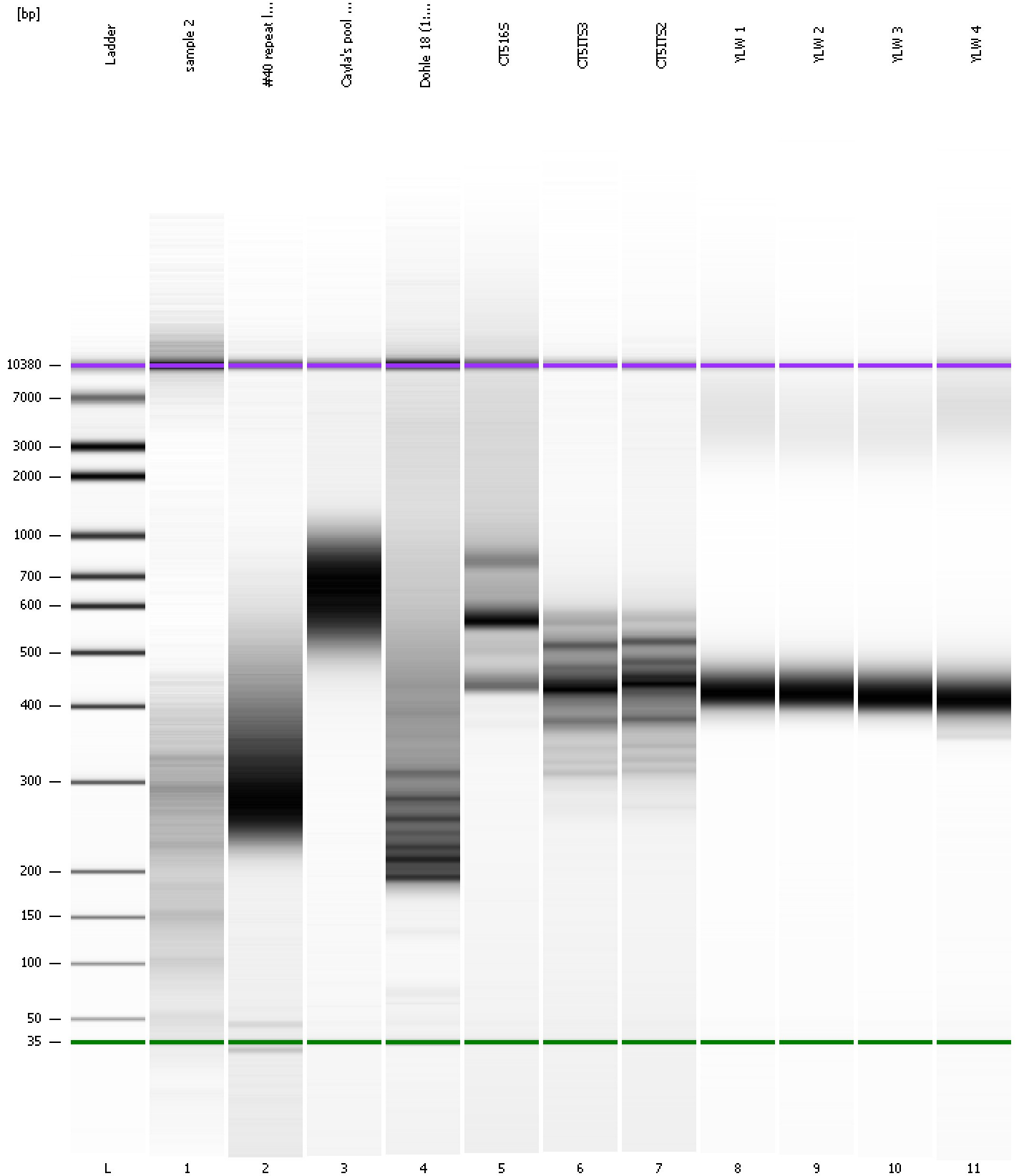
**Region table for sample 11 : YLW 4**

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	421	1,002.0	5,928.6	1,640.11	85	7.9

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
Modified: 12/9/2015 11:40:51 AM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-12-09\2015-12-09\_002.xad

Created: 12/9/2015 11:00:29 AM  
 Modified: 12/9/2015 11:40:51 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		12/9/2015 11:40:51 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-12-09\2015-12-09_002.xad)		Instrument	Run		12/9/2015 11:00:29 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		12/9/2015 11:00:29 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		12/9/2015 11:00:29 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		12/9/2015 11:00:29 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		12/9/2015 11:00:29 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		12/9/2015 11:00:29 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		12/9/2015 11:00:29 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1