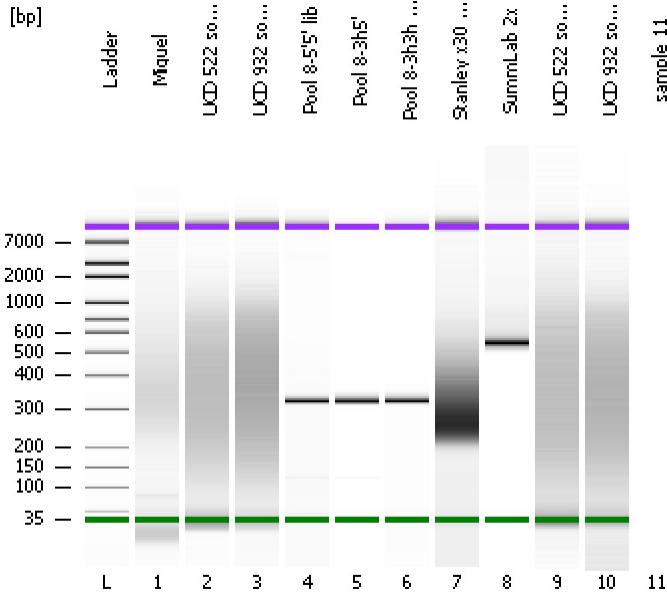


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
Modified: 12/9/2014 11:37:41 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

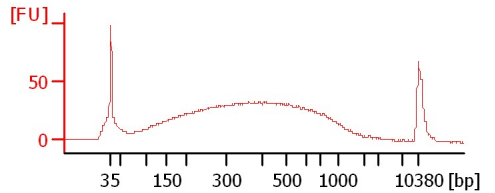
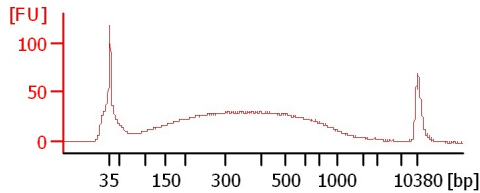
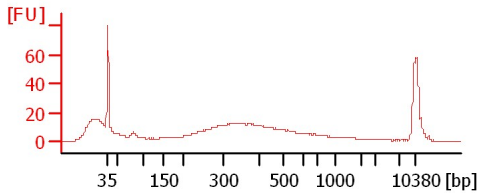
Chip Information:

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

Miguel

UCD 522 soni (1:4)

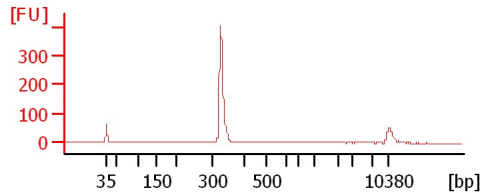
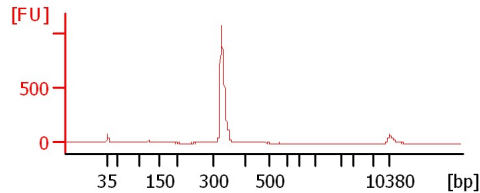
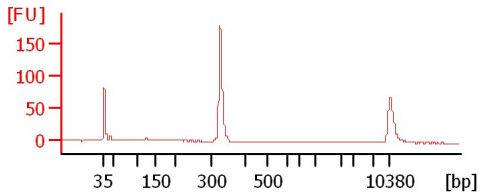
UCD 932 soni (1:4)



Pool 8-5'5' lib

Pool 8-3h5'

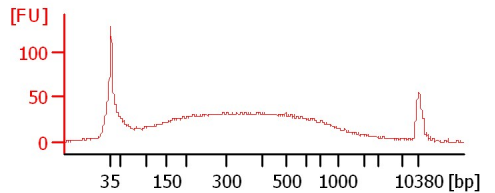
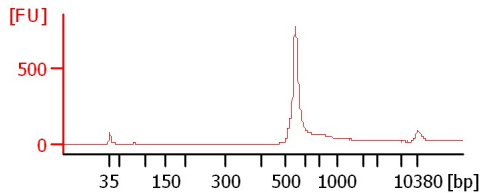
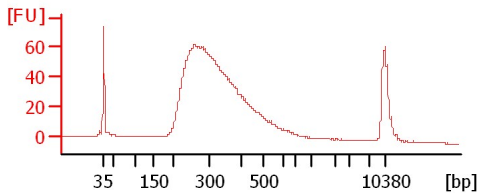
Pool 8-3h3h (1:15)



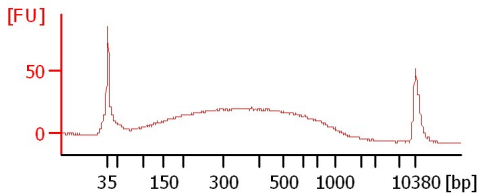
Stanley x30 (1:10) lib. pool

SummLab 2x

UCD 522 soni (1:4)



UCD 932 soni (1:4)



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
Modified: 12/9/2014 11:37:41 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Miguel		<input type="checkbox"/>	✓			
UCD 522 soni (1:4)		<input type="checkbox"/>	✓			
UCD 932 soni (1:4)		<input type="checkbox"/>	✓			
Pool 8-5'5' lib		<input type="checkbox"/>	✓			
Pool 8-3h5'		<input type="checkbox"/>	✓			
Pool 8-3h3h (1:15)		<input type="checkbox"/>	✓			
Stanley x30 (1:10) lib. pool		<input type="checkbox"/>	✓			
SummLab 2x		<input type="checkbox"/>	✓			
UCD 522 soni (1:4)		<input type="checkbox"/>	✓			
UCD 932 soni (1:4)		<input type="checkbox"/>	✓			
sample 11		<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\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
Modified: 12/9/2014 11:37:41 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

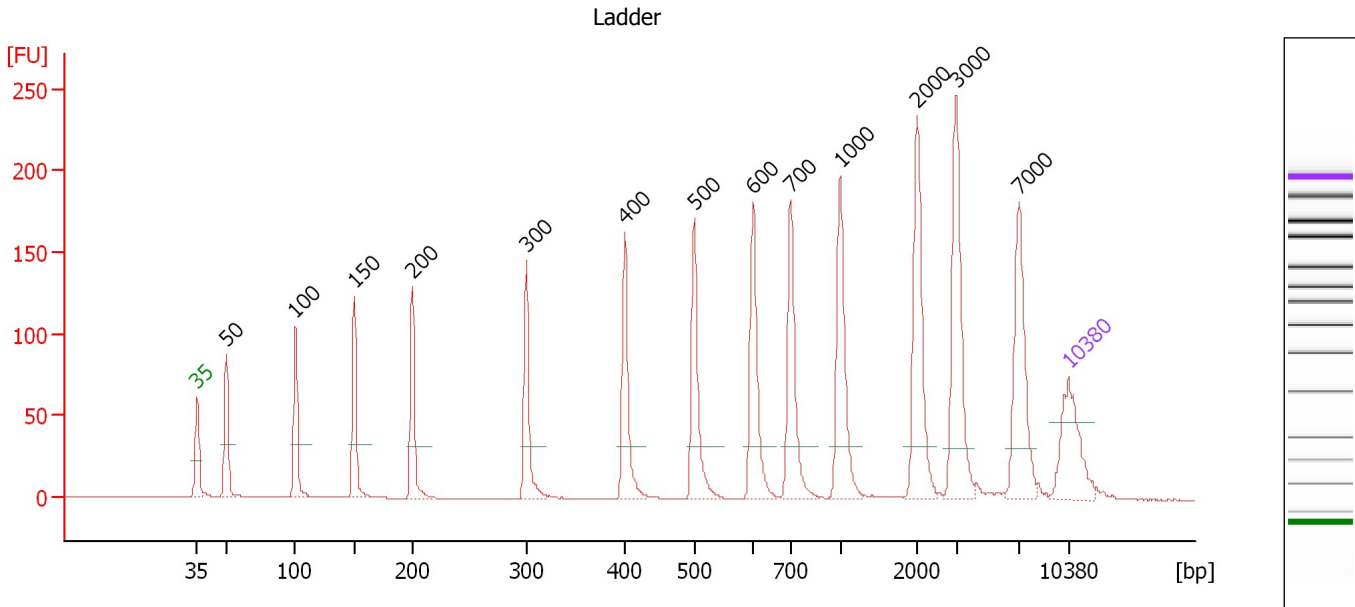
Ladder

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\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

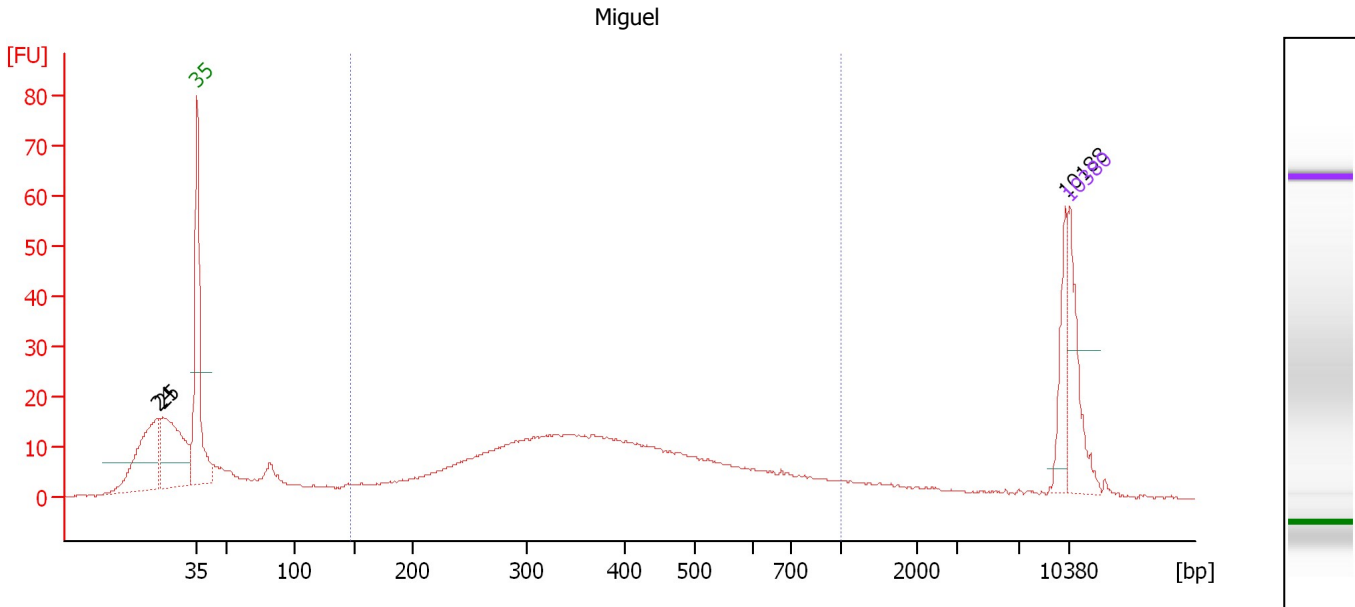
Peak table for Ladder

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	45.33
3	100	150.00	2,272.7	Ladder Peak	50.88
4	150	150.00	1,515.2	Ladder Peak	55.63
5	200	150.00	1,136.4	Ladder Peak	60.29
6	300	150.00	757.6	Ladder Peak	69.44
7	400	150.00	568.2	Ladder Peak	77.36
8	500	150.00	454.5	Ladder Peak	82.95
9	600	150.00	378.8	Ladder Peak	87.66
10	700	150.00	324.7	Ladder Peak	90.65
11	1,000	150.00	227.3	Ladder Peak	94.65
12	2,000	150.00	113.6	Ladder Peak	100.81
13	3,000	150.00	75.8	Ladder Peak	103.98
14	7,000	150.00	32.5	Ladder Peak	109.04
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : Miguel

Number of peaks found: 3 Corr. Area 1: 427.3
 Noise: 0.2

Peak table for sample 1 : Miguel

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	24	0.00	0.0		39.80
2	25	0.00	0.0		40.21
3	35	125.00	5,411.3	Lower Marker	43.00
4	10,188	50.29	7.5		112.77
5	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 1 : Miguel

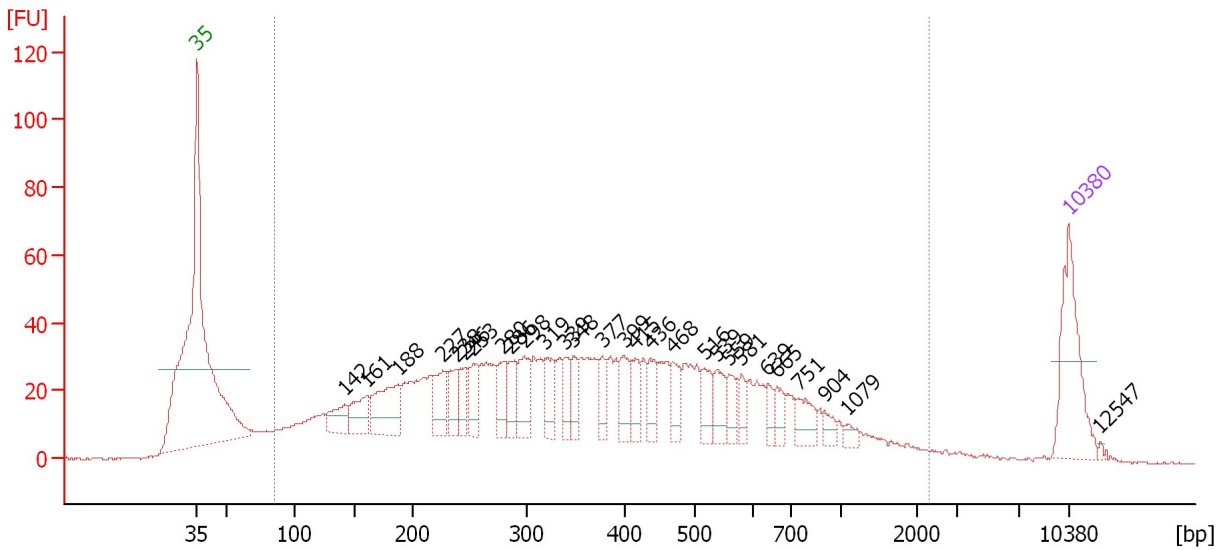
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
147	402	1,000	427.3	55	39.9	953.00	4,437.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...

UCD 522 soni (1:4)



Overall Results for sample 2 : UCD 522 soni (1:4)

Number of peaks found: 28 Corr. Area 1: 1,620.9
 Noise: 0.3

Peak table for sample 2 : UCD 522 soni (1:4)

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	142	35.28	376.1		54.88
3	161	43.23	407.0		56.65
4	188	80.67	651.3		59.14
5	227	46.14	308.6		62.72
6	238	32.03	203.9		63.76
7	246	26.68	164.1		64.53
8	253	33.32	199.3		65.17
9	280	32.72	177.3		67.57
10	286	32.72	173.6		68.12
11	298	44.58	226.7		69.25
12	319	37.29	177.3		70.93
13	339	25.22	112.8		72.51
14	348	23.15	100.8		73.24
15	377	25.14	101.2		75.51
16	399	33.60	127.6		77.27
17	415	23.24	84.7		78.23
18	436	28.81	100.2		79.36
19	468	25.77	83.4		81.17
20	516	27.94	82.0		83.71
21	539	26.23	73.7		84.80
22	559	19.87	53.9		85.71
23	581	13.60	35.5		86.75
24	639	12.67	30.0		88.84
25	665	17.96	40.9		89.61

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad


Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...

... Peak table for sample 2 : UCD 522 soni (1:4)

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	751	24.75	49.9		91.33
27	904	9.70	16.3		93.37
28	1,079	6.82	9.6		95.14
29	10,380	75.00	10.9	Upper Marker	113.00
30	12,547	0.00	0.0		115.54

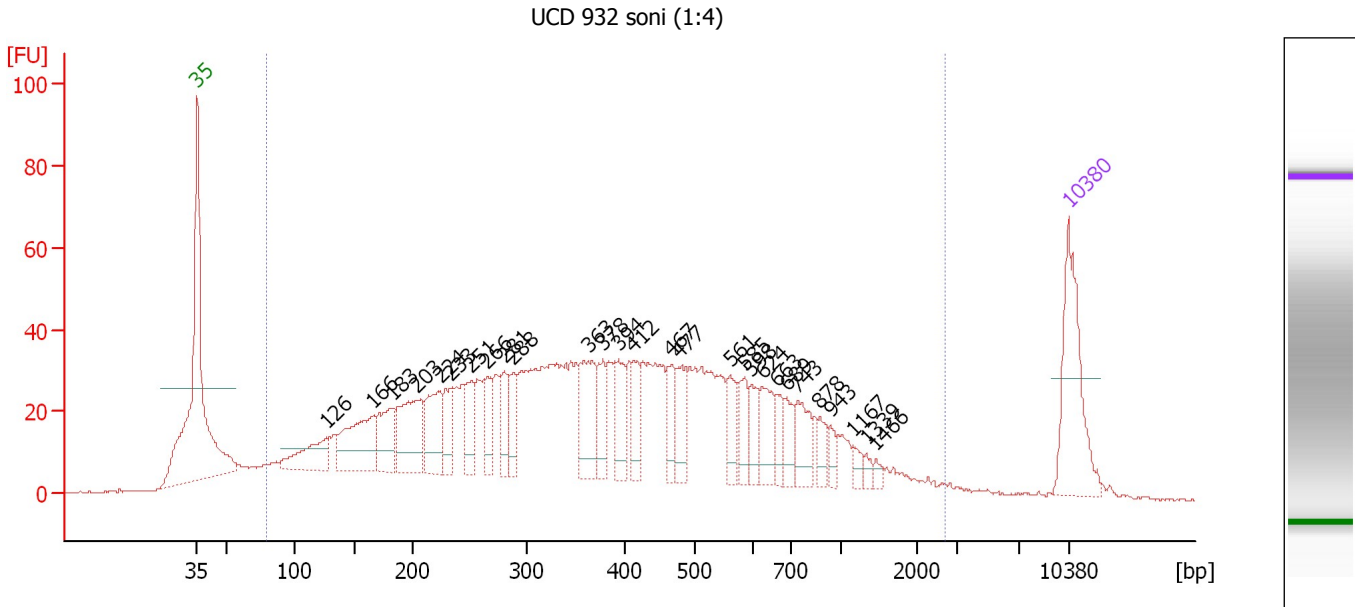
Region table for sample 2 : UCD 522 soni (1:4)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/ μ l]	Molarity [pmol/l]	Color
84	426	2,321	1,620.9	97	69.2	1,945.63	11,515.1	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...



Overall Results for sample 3 : UCD 932 soni (1:4)

Number of peaks found: 28 Corr. Area 1: 1,731.6
 Noise: 0.2

Peak table for sample 3 : UCD 932 soni (1:4)

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	126	58.32	699.3		53.38
3	166	94.11	857.5		57.15
4	183	57.08	473.8		58.66
5	203	89.79	670.8		60.55
6	224	63.83	432.0		62.48
7	233	39.61	257.7		63.30
8	251	34.94	210.5		65.00
9	266	31.07	177.3		66.29
10	281	31.35	169.3		67.67
11	288	30.98	162.9		68.35
12	363	64.93	270.7		74.46
13	378	34.38	137.8		75.61
14	394	35.61	137.1		76.85
15	412	31.09	114.3		78.05
16	467	27.37	88.7		81.12
17	477	32.22	102.3		81.67
18	561	27.95	75.5		85.81
19	585	26.36	68.3		86.96
20	598	23.35	59.2		87.55
21	624	35.33	85.8		88.38
22	663	15.54	35.5		89.53
23	689	20.35	44.8		90.31
24	743	32.82	66.9		91.23
25	878	14.26	24.6		93.02

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad


Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...

... Peak table for sample 3 : UCD 932 soni (1:4)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	943	9.99	16.1		93.89
27	1,167	7.95	10.3		95.68
28	1,339	5.20	5.9		96.74
29	1,466	5.00	5.2		97.52
30	10,380	75.00	10.9	Upper Marker	113.00

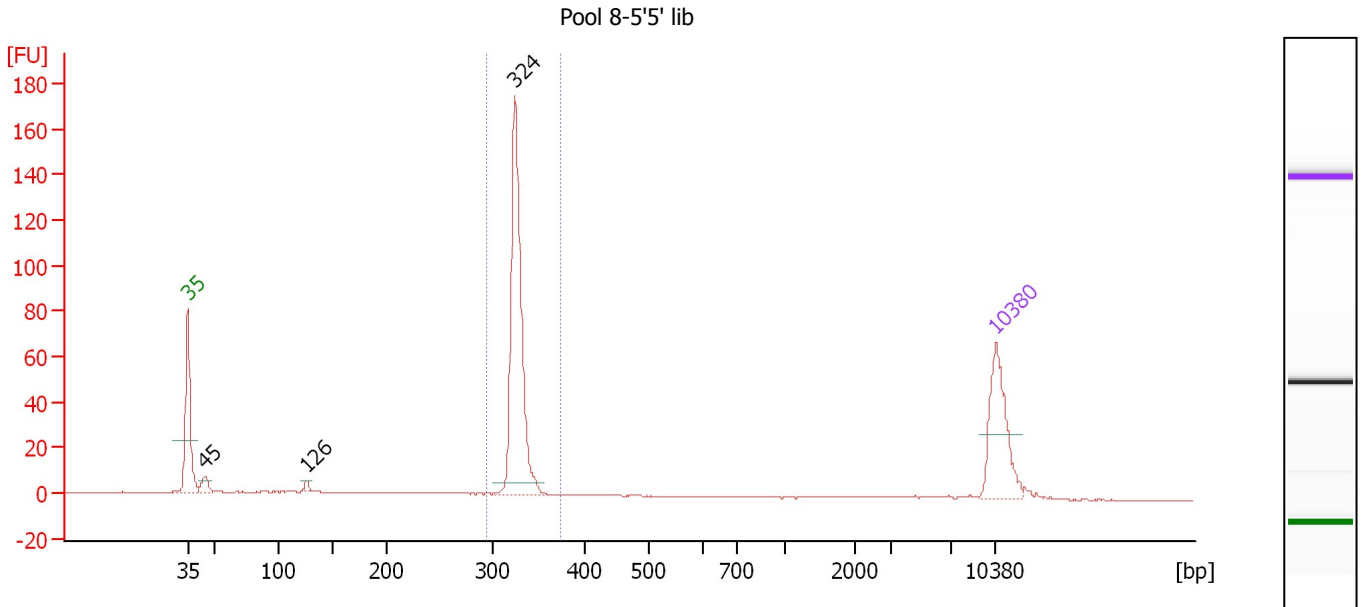
Region table for sample 3 : UCD 932 soni (1:4)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
79	450	2,708	1,731.6	97	72.9	2,103.85	12,211.1	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...



Overall Results for sample 4 : Pool 8-5'5' lib

Number of peaks found: 3 Corr. Area 1: 265.3
 Noise: 0.1

Peak table for sample 4 : Pool 8-5'5' lib

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	45	17.44	589.9		44.52
3	126	6.17	74.2		53.34
4	324	291.02	1,360.5		71.35
5	10,380	75.00	10.9	Upper Marker	113.00

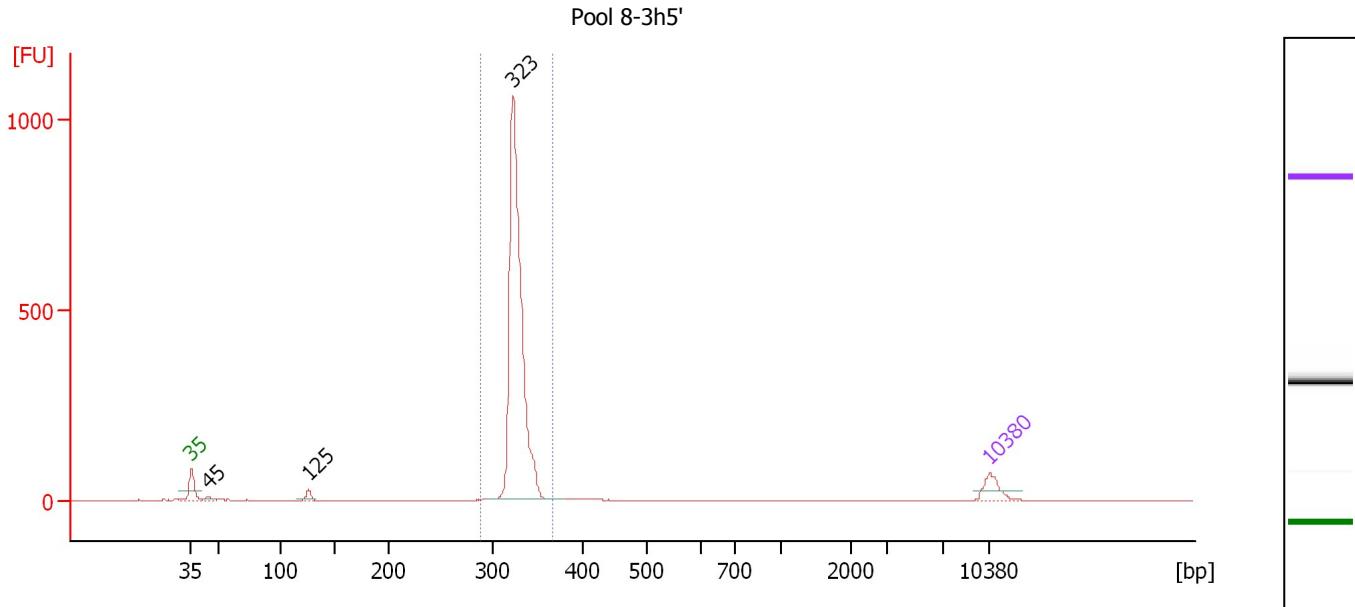
Region table for sample 4 : Pool 8-5'5' lib

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
294	327	375	265.3	72	2.4	299.08	1,386.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : Pool 8-3h5'

Number of peaks found: 3 Corr. Area 1: 1,850.4
 Noise: 0.1

Peak table for sample 5 : Pool 8-3h5'

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	45	18.87	641.3		44.49
3	125	32.99	398.7		53.29
4	323	2,002.51	9,398.2		71.25
5	10,380	75.00	10.9	Upper Marker	113.00

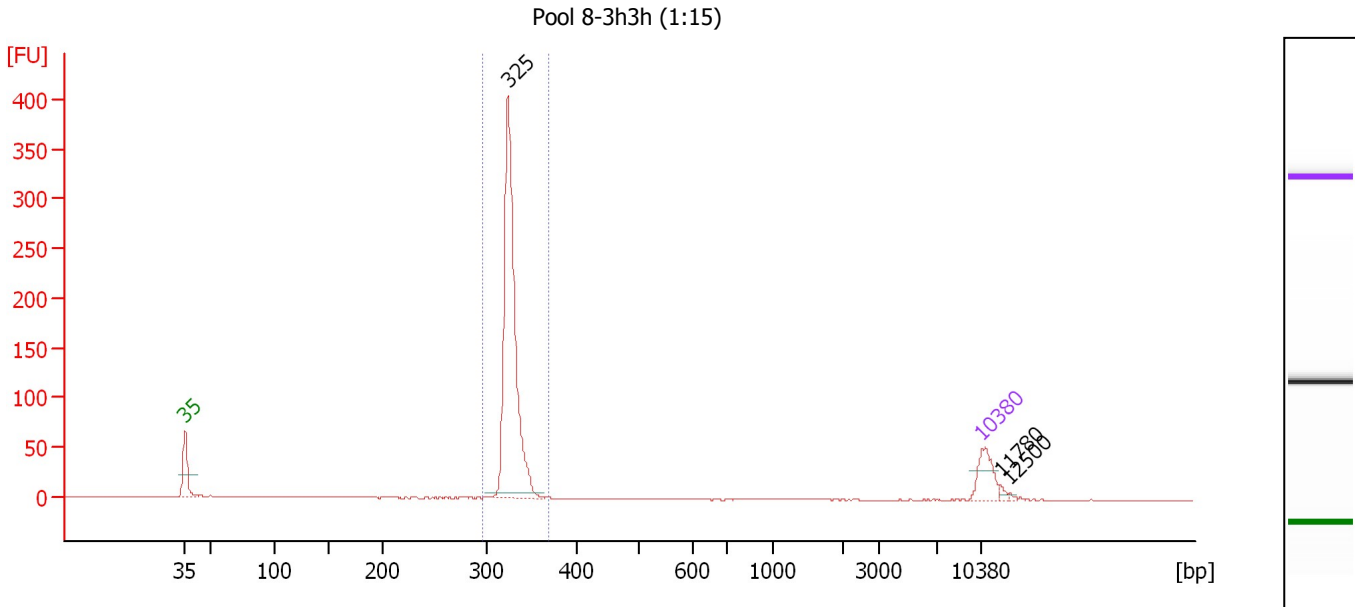
Region table for sample 5 : Pool 8-3h5'

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
289	327	366	1,850.4	93	2.4	2,000.32	9,262.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...



Overall Results for sample 6 : Pool 8-3h3h (1:15)

Number of peaks found: 3 Corr. Area 1: 636.5
 Noise: 0.2

Peak table for sample 6 : Pool 8-3h3h (1:15)

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	325	1,019.98	4,760.5		71.39
3	10,380	75.00	10.9	Upper Marker	113.00
4	11,780	0.00	0.0		114.64
5	12,500	0.00	0.0		115.48

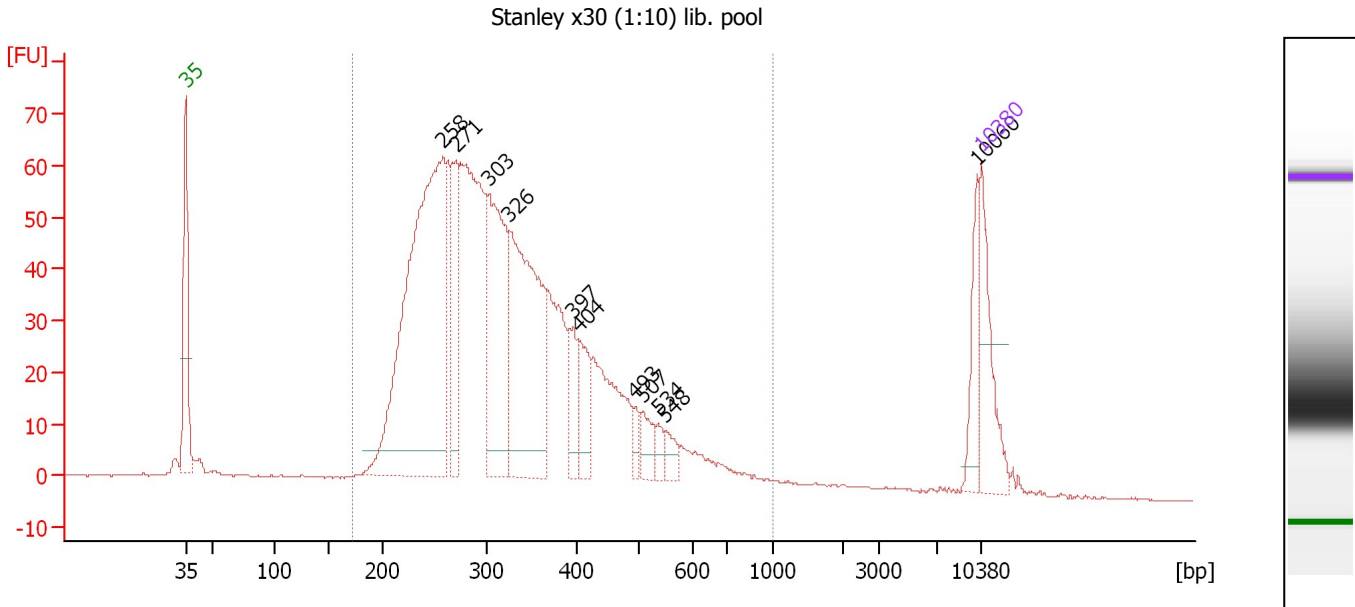
Region table for sample 6 : Pool 8-3h3h (1:15)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
296	328	369	636.5	93	2.3	1,016.58	4,694.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...



Overall Results for sample 7 : Stanley x30 (1:10) lib. pool

Number of peaks found: 11 Corr. Area 1: 1,452.6
 Noise: 0.2

Peak table for sample 7 : Stanley x30 (1:10) lib. pool

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	258	683.95	4,016.6		65.60
3	271	119.54	667.5		66.82
4	303	249.11	1,245.8		69.68
5	326	336.75	1,564.8		71.51
6	397	44.46	169.9		77.09
7	404	58.60	219.6		77.60
8	493	16.00	49.1		82.57
9	507	28.40	84.9		83.27
10	534	13.97	39.6		84.54
11	548	17.35	48.0		85.20
12	10,060	43.83	6.6		112.62
13	10,380	75.00	10.9	Upper Marker	113.00

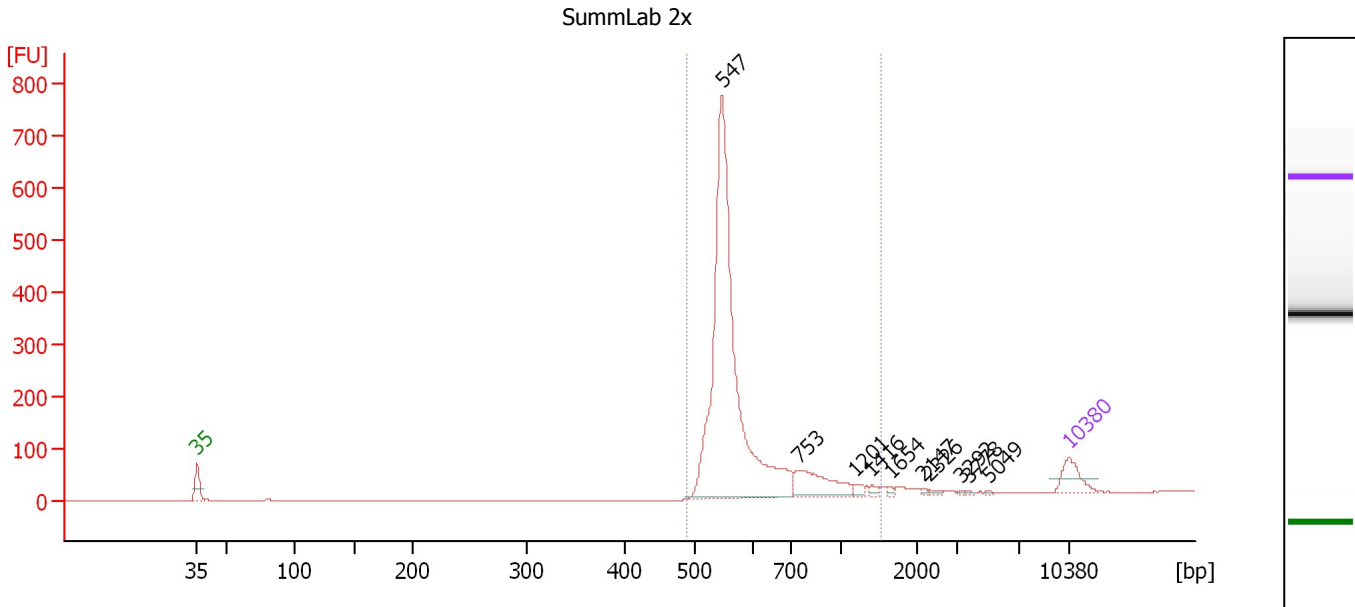
Region table for sample 7 : Stanley x30 (1:10) lib. pool

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
172	337	1,000	1,452.6	93	32.4	2,508.49	12,581.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...



Overall Results for sample 8 : SummLab 2x

Number of peaks found: 10 Corr. Area 1: 1,948.5
 Noise: 0.1

Peak table for sample 8 : SummLab 2x

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	547	1,806.31	5,003.7		85.16
3	753	195.95	394.4		91.35
4	1,201	19.75	24.9		95.89
5	1,416	12.13	13.0		97.21
6	1,654	8.56	7.8		98.68
7	2,147	5.92	4.2		101.28
8	2,326	8.29	5.4		101.85
9	3,292	2.91	1.3		104.35
10	3,778	3.72	1.5		104.96
11	5,049	3.51	1.1		106.57
12	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 8 : SummLab 2x

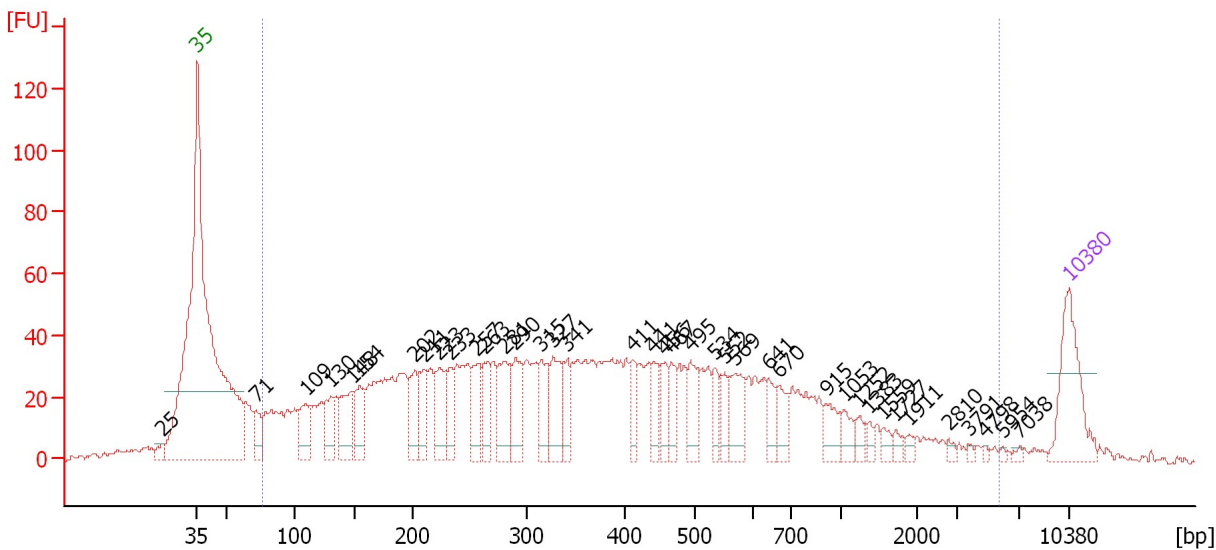
From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
490	608	1,527	1,948.5	96	24.7	1,904.57	4,952.5	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...

UCD 522 soni (1:4)



Overall Results for sample 9 : UCD 522 soni (1:4)

Number of peaks found: 39 Corr. Area 1: 1,903.4
 Noise: 0.7

Peak table for sample 9 : UCD 522 soni (1:4)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	25	0.00	0.0		40.17
2	35	125.00	5,411.3	Lower Marker	43.00
3	71	39.87	849.8		47.67
4	109	53.42	742.9		51.73
5	130	50.59	588.3		53.75
6	148	73.48	753.9		55.41
7	154	49.09	482.3		56.02
8	202	53.10	398.7		60.45
9	211	47.07	338.8		61.25
10	223	62.49	423.8		62.43
11	233	42.30	275.4		63.28
12	257	54.14	319.3		65.50
13	263	44.92	258.7		66.07
14	281	70.57	381.0		67.67
15	290	63.57	331.6		68.57
16	315	50.96	245.0		70.64
17	327	61.70	285.8		71.58
18	341	35.00	155.4		72.72
19	411	30.46	112.4		77.95
20	441	29.44	101.1		79.65
21	456	31.05	103.1		80.50
22	467	28.33	91.9		81.11
23	495	40.16	122.9		82.67
24	534	24.45	69.3		84.56
25	552	22.96	63.0		85.41

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad


Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...

... Peak table for sample 9 : UCD 522 soni (1:4)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	569	49.44	131.6		86.21
27	641	25.56	60.4		88.90
28	670	27.35	61.9		89.75
29	915	25.49	42.2		93.52
30	1,053	17.86	25.7		94.98
31	1,252	11.14	13.5		96.21
32	1,383	7.26	8.0		97.01
33	1,559	8.70	8.5		98.09
34	1,727	7.50	6.6		99.13
35	1,911	5.01	4.0		100.26
36	2,810	4.30	2.3		103.38
37	3,791	2.89	1.2		104.98
38	4,798	2.53	0.8		106.25
39	5,954	2.41	0.6		107.72
40	7,038	3.08	0.7		109.08
41	10,380	75.00	10.9	Upper Marker	113.00

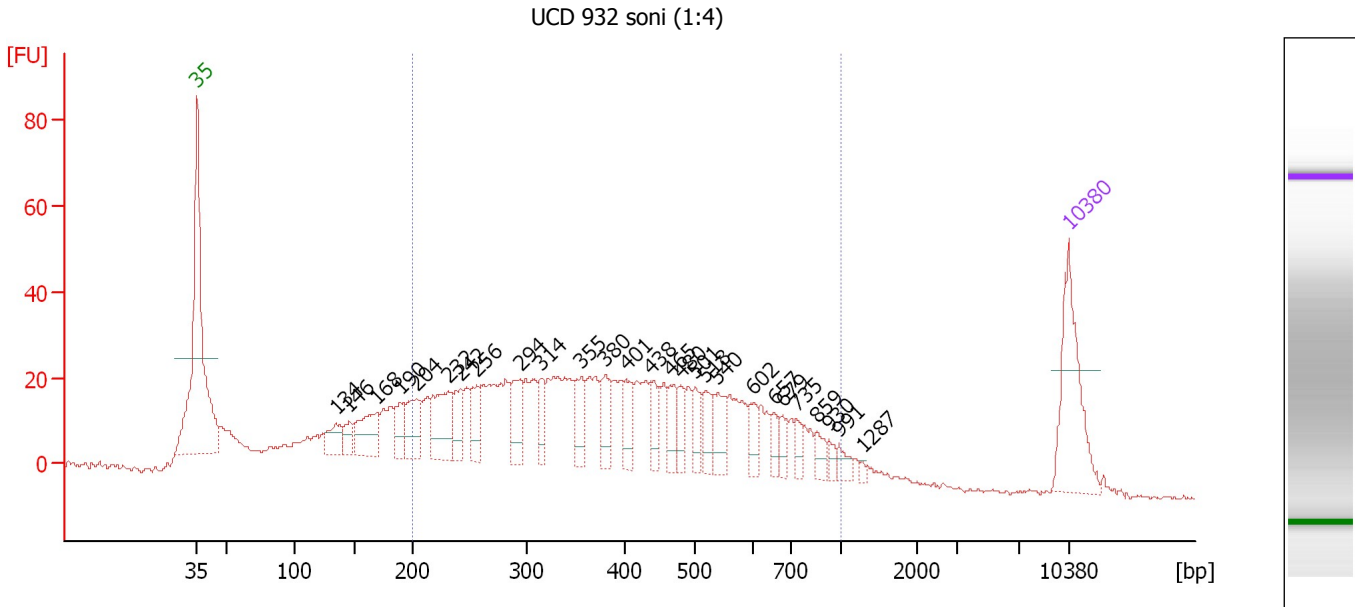
Region table for sample 9 : UCD 522 soni (1:4)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
76	499	5,716	1,903.4	94	100.0	2,518.78	16,164.5	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...



Overall Results for sample 10 : UCD 932 soni (1:4)

Number of peaks found: 27 Corr. Area 1: 939.0
 Noise: 0.2

Peak table for sample 10 : UCD 932 soni (1:4)

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	134	29.10	328.4		54.13
3	146	18.47	192.1		55.22
4	168	57.15	514.0		57.35
5	190	28.58	228.1		59.34
6	204	43.95	327.1		60.62
7	232	68.71	449.6		63.18
8	242	32.65	204.1		64.17
9	256	29.63	175.1		65.45
10	294	42.03	216.5		68.91
11	314	25.52	123.1		70.56
12	355	34.87	148.7		73.83
13	380	32.13	128.1		75.77
14	401	28.40	107.2		77.43
15	438	24.03	83.2		79.47
16	465	24.68	80.4		80.98
17	480	19.75	62.3		81.84
18	501	21.70	65.6		83.02
19	518	22.11	64.7		83.78
20	540	31.33	87.9		84.82
21	602	17.47	44.0		87.71
22	657	12.37	28.5		89.37
23	679	12.86	28.7		90.03
24	735	11.09	22.9		91.12
25	859	12.56	22.1		92.78

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad


Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Electropherogram Summary Continued ...

... Peak table for sample 10 : UCD 932 soni (1:4)

Peak	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	930	6.85	11.2		93.72
27	991	10.16	15.5		94.53
28	1,287	2.98	3.5		96.42
29	10,380	75.00	10.9	Upper Marker	113.00

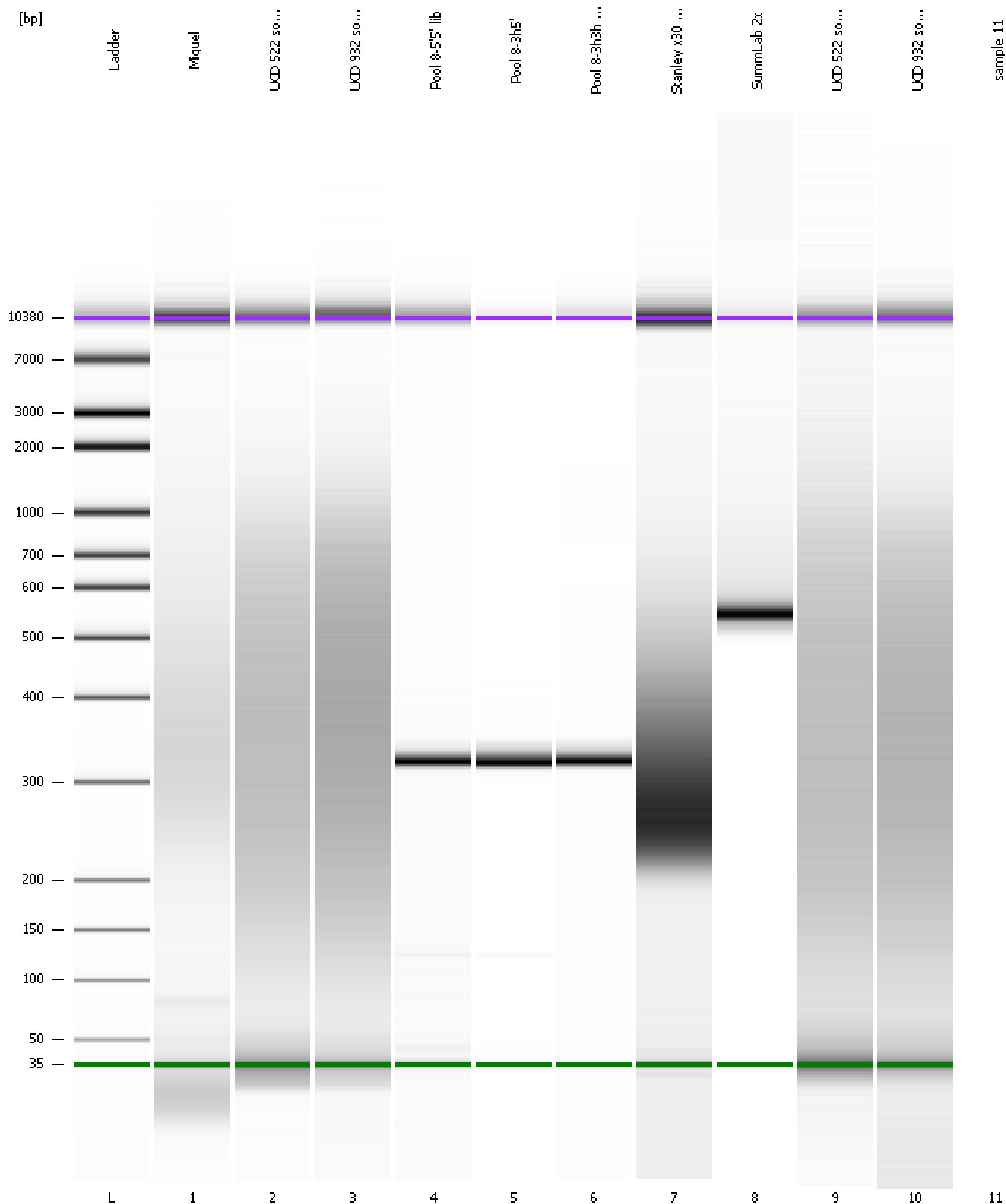
Region table for sample 10 : UCD 932 soni (1:4)

From [bp]	Average Size [bp]	To [bp]	Corr. Area	% of Total	Size distribution in CV [%]	Conc. [pg/ μ l]	Molarity [pmol/l]	Color
200	425	1,000	939.0	73	39.9	1,301.47	5,682.7	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
Modified: 12/9/2014 11:37:41 AM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
Modified: 12/9/2014 11:37:41 AM

Invalid Samples

Sample 11 has not been run, no results available.

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad

Created: 12/9/2014 10:59:11 AM
 Modified: 12/9/2014 11:37:41 AM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 11)		Instrument	Run		12/9/2014 11:37:38 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2014-12-09\2014-12-09_001.xad)		Instrument	Run		12/9/2014 10:59:16 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		12/9/2014 10:59:16 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		12/9/2014 10:59:16 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		12/9/2014 10:59:16 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		12/9/2014 10:59:16 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		12/9/2014 10:59:16 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		12/9/2014 10:59:16 AM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1