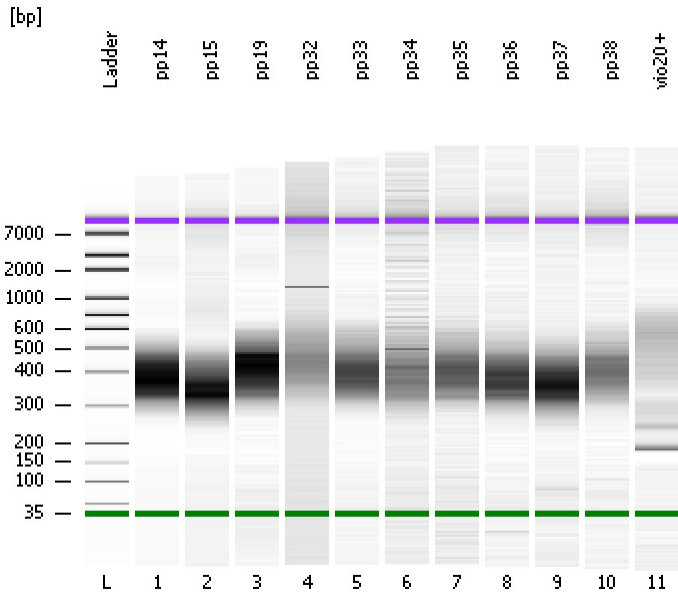


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

Created: 2/7/2014 11:41:00 AM
Modified: 2/7/2014 12:23:14 PM

Electrophoresis File Run Summary



Instrument Information:

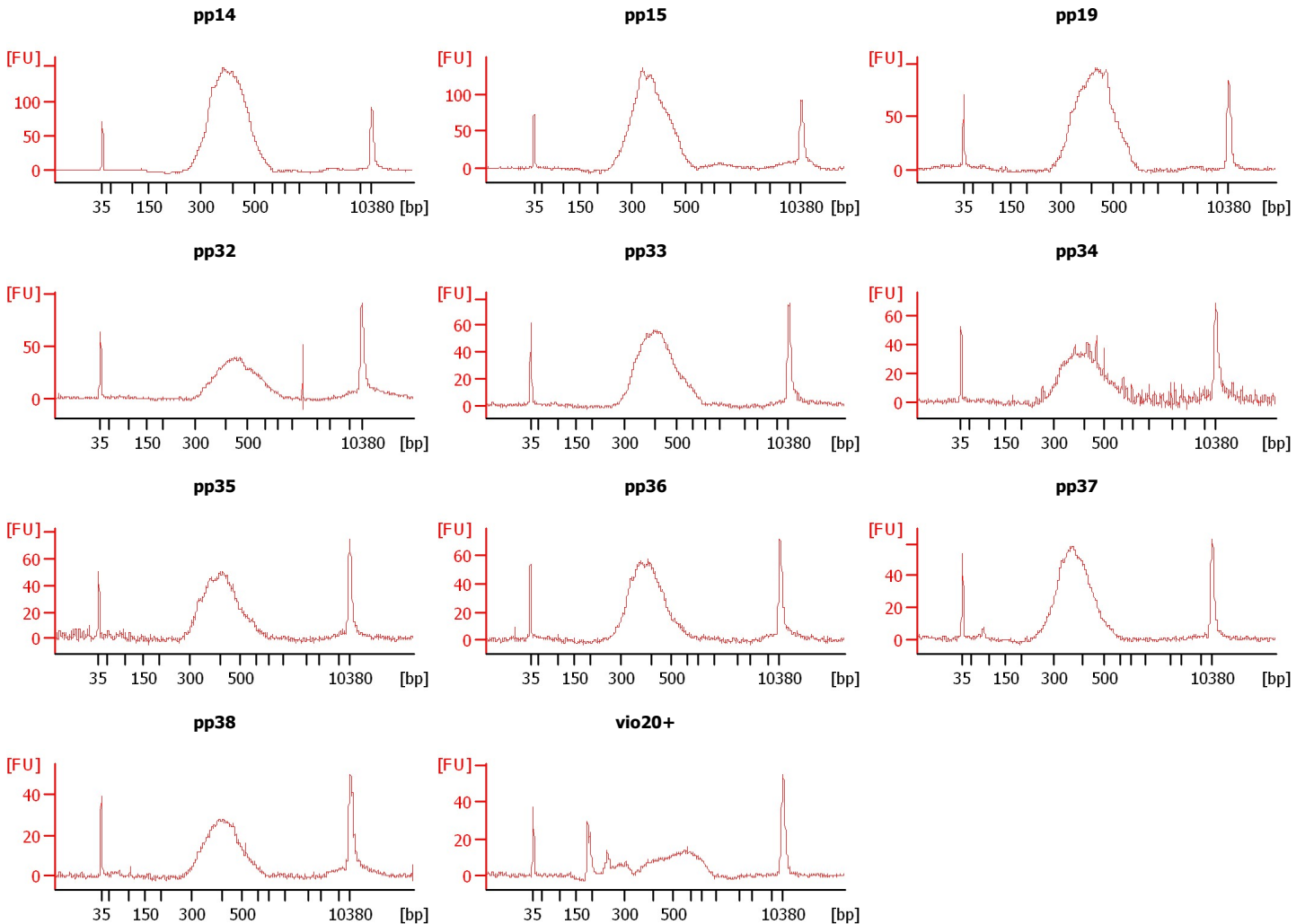
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

Created: 2/7/2014 11:41:00 AM
Modified: 2/7/2014 12:23:14 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
pp14		<input type="checkbox"/>	✓			
pp15		<input type="checkbox"/>	✓			
pp19		<input type="checkbox"/>	✓			
pp32		<input type="checkbox"/>	✓			
pp33		<input type="checkbox"/>	✓			
pp34		<input type="checkbox"/>	✓			
pp35		<input type="checkbox"/>	✓			
pp36		<input type="checkbox"/>	✓			
pp37		<input type="checkbox"/>	✓			
pp38		<input type="checkbox"/>	✓			
vio20+		<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\2014-02-07\2014-02-07_001.xad

Created: 2/7/2014 11:41:00 AM
Modified: 2/7/2014 12:23:14 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

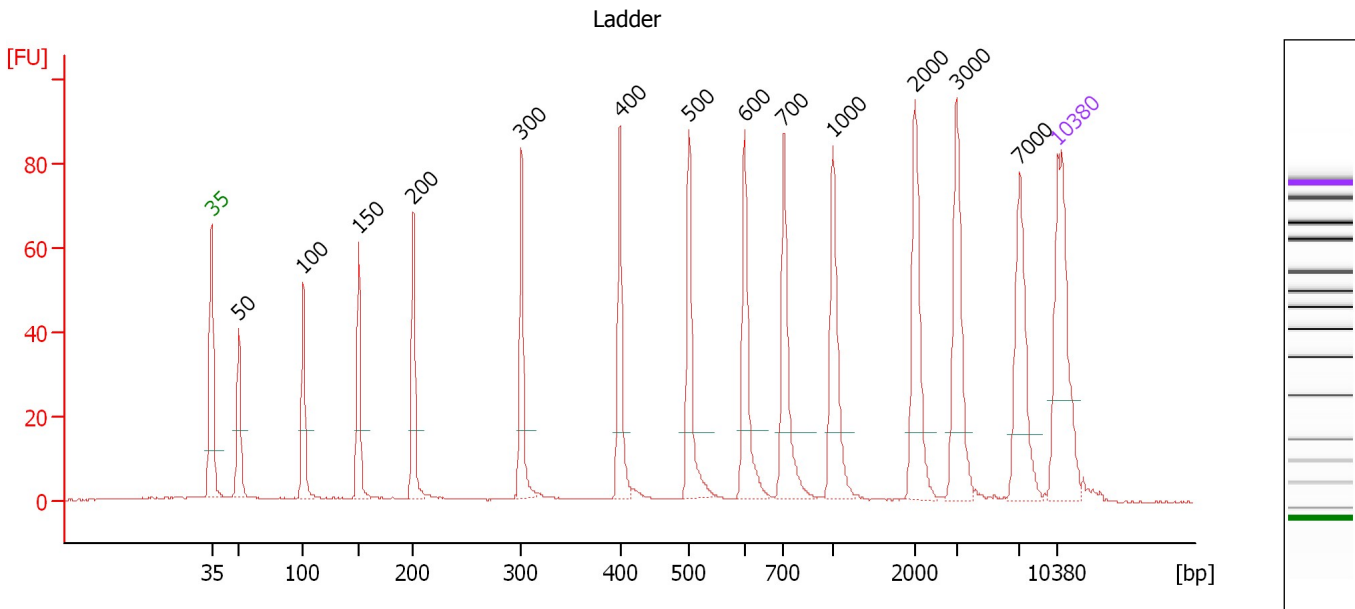
Ladder

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\2014-02-07\2014-02-07_001.xad

Created: 2/7/2014 11:41:00 AM
 Modified: 2/7/2014 12:23:14 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

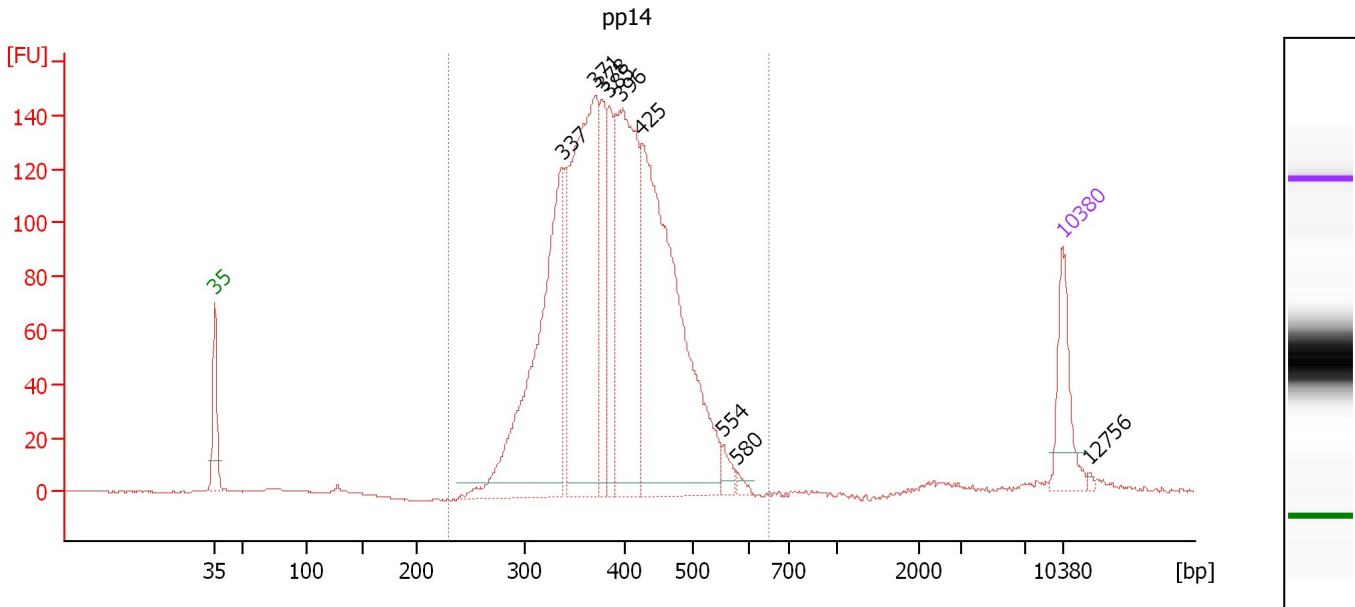
Peak table for Ladder

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	45.25	50	150.00	4,545.5	Ladder Peak
3	50.59	100	150.00	2,272.7	Ladder Peak
4	55.17	150	150.00	1,515.2	Ladder Peak
5	59.67	200	150.00	1,136.4	Ladder Peak
6	68.63	300	150.00	757.6	Ladder Peak
7	76.79	400	150.00	568.2	Ladder Peak
8	82.48	500	150.00	454.5	Ladder Peak
9	87.11	600	150.00	378.8	Ladder Peak
10	90.33	700	150.00	324.7	Ladder Peak
11	94.39	1,000	150.00	227.3	Ladder Peak
12	101.18	2,000	150.00	113.6	Ladder Peak
13	104.62	3,000	150.00	75.8	Ladder Peak
14	109.87	7,000	150.00	32.5	Ladder Peak
15	113.00	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

Created: 2/7/2014 11:41:00 AM
 Modified: 2/7/2014 12:23:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : pp14

Number of peaks found: 9 Corr. Area 1: 2,268.5
 Noise: 0.1

Peak table for sample 1 : pp14

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	71.66	337	542.29	2,436.9	
3	74.39	371	603.10	2,465.5	
4	75.01	378	141.18	565.6	
5	75.58	385	126.72	498.4	
6	76.46	396	450.93	1,725.4	
7	78.22	425	704.66	2,511.1	
8	85.00	554	19.94	54.5	
9	86.19	580	7.83	20.5	
10	113.00	10,380	75.00	10.9	Upper Marker
11	115.20	12,756	0.00	0.0	

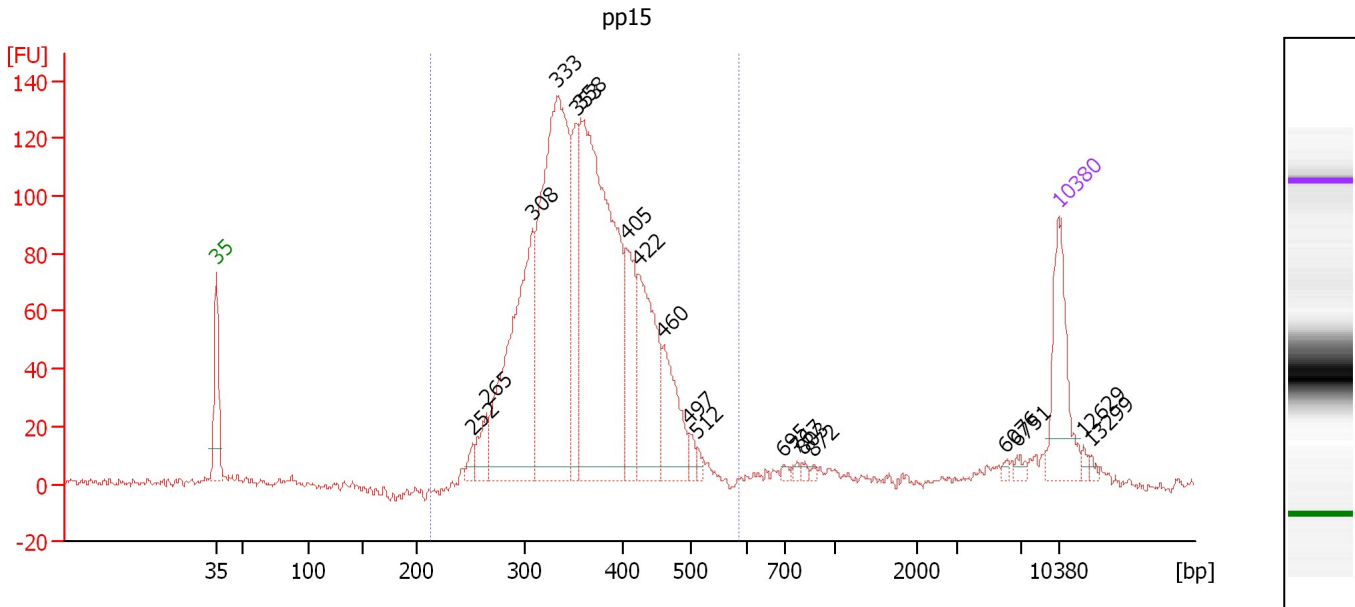
Region table for sample 1 : pp14

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
230	651	395	10,284.6	2,601.04	2,268.5	98	15.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

Created: 2/7/2014 11:41:00 AM
 Modified: 2/7/2014 12:23:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : pp15

Number of peaks found: 19 Corr. Area 1: 1,909.3
 Noise: 1.4

Peak table for sample 2 : pp15

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	64.32	252	13.18	79.2	
3	65.48	265	38.75	221.7	
4	69.29	308	319.37	1,570.5	
5	71.28	333	521.26	2,374.9	
6	72.92	353	99.81	428.8	
7	73.32	358	561.38	2,378.9	
8	77.09	405	103.83	388.1	
9	78.07	422	155.19	556.6	
10	80.19	460	87.19	287.2	
11	82.32	497	10.64	32.4	
12	83.03	512	6.85	20.3	
13	90.17	695	3.80	8.3	
14	91.23	767	3.19	6.3	
15	91.72	803	3.34	6.3	
16	92.65	872	2.74	4.8	
17	108.66	6,076	2.69	0.7	
18	109.54	6,751	5.11	1.1	
19	113.00	10,380	75.00	10.9	Upper Marker
20	115.08	12,629	0.00	0.0	
21	115.70	13,299	0.00	0.0	

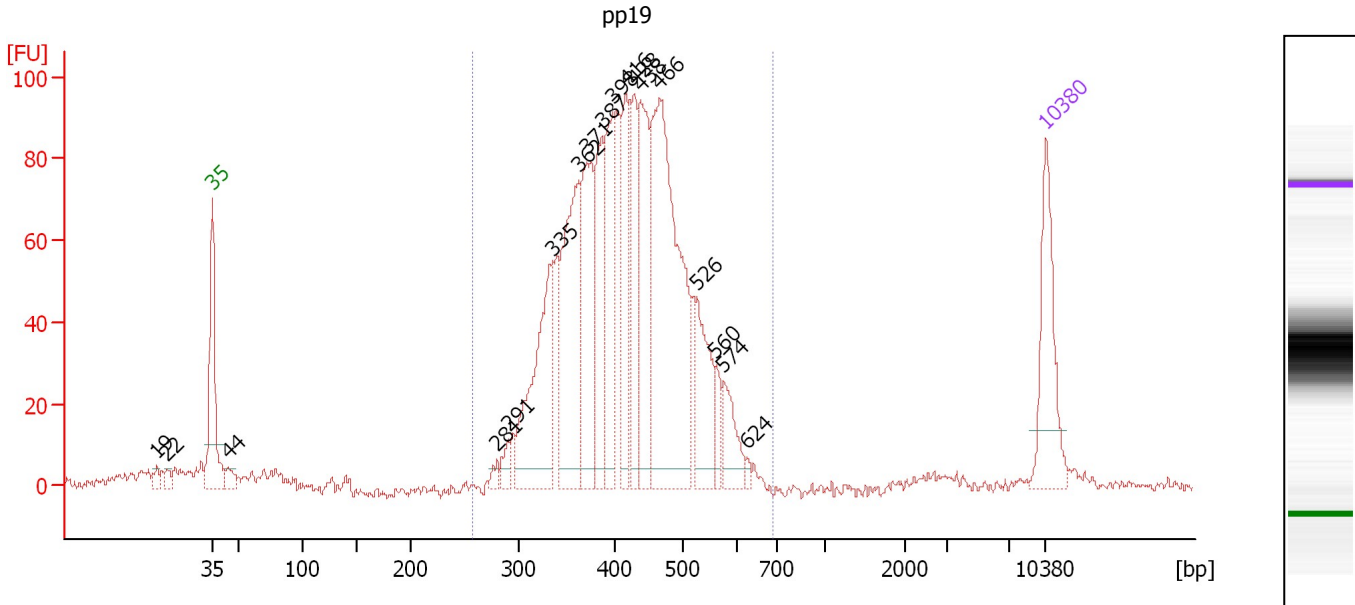
Region table for sample 2 : pp15

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
213	586	363	8,273.0	1,921.90	1,909.3	95	15.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

Created: 2/7/2014 11:41:00 AM
 Modified: 2/7/2014 12:23:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : pp19

Number of peaks found: 18 Corr. Area 1: 1,523.3
 Noise: 1.4

Peak table for sample 3 : pp19

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	38.30	19	0.00	0.0	
2	39.15	22	0.00	0.0	
3	43.00	35	125.00	5,411.3	Lower Marker
4	44.34	44	17.21	593.1	
5	66.92	281	8.54	46.1	
6	67.86	291	15.25	79.3	
7	71.48	335	175.32	792.9	
8	73.72	362	189.03	790.2	
9	74.39	371	139.44	570.0	
10	75.69	387	102.43	401.4	
11	76.63	398	131.44	500.2	
12	77.71	416	93.32	339.7	
13	78.38	428	80.51	285.0	
14	78.96	438	131.68	455.3	
15	80.53	466	360.35	1,172.2	
16	83.67	526	82.46	237.7	
17	85.23	560	20.55	55.7	
18	85.90	574	40.66	107.3	
19	87.88	624	4.55	11.1	
20	113.00	10,380	75.00	10.9	Upper Marker

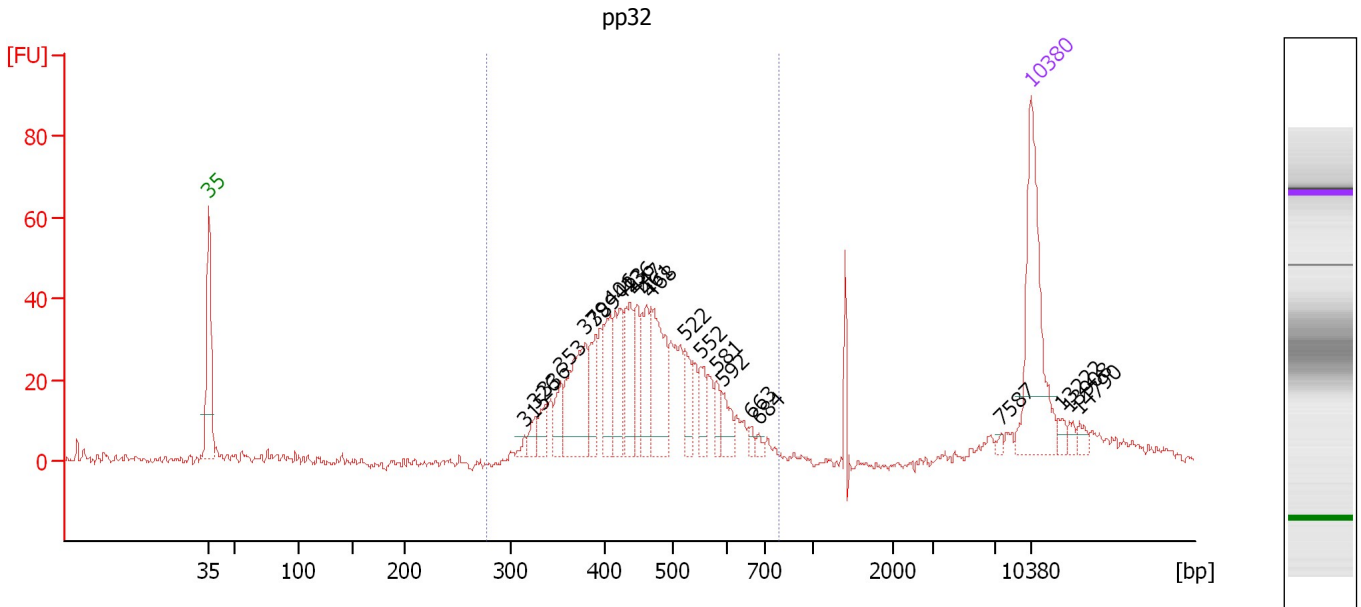
Region table for sample 3 : pp19

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
258	689	423	6,413.2	1,726.01	1,523.3	94	16.8	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

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 Modified: 2/7/2014 12:23:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : pp32

Number of peaks found: 22 Corr. Area 1: 600.2
 Noise: 1.2

Peak table for sample 4 : pp32


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	69.87	315	4.85	23.3	
3	70.78	326	10.94	50.8	
4	71.55	336	14.38	64.9	
5	72.96	353	20.27	87.0	
6	75.05	379	68.91	275.7	
7	75.91	389	27.56	107.3	
8	77.14	406	35.59	132.8	
9	78.09	423	39.04	139.9	
10	78.82	436	37.99	132.1	
11	79.45	447	28.79	97.6	
12	80.27	461	32.20	105.8	
13	80.63	468	61.11	198.0	
14	83.49	522	19.49	56.6	
15	84.90	552	18.28	50.1	
16	86.22	581	10.92	28.5	
17	86.72	592	13.86	35.5	
18	89.12	663	4.28	9.8	
19	89.80	684	3.46	7.7	
20	110.41	7,587	1.96	0.4	
21	113.00	10,380	75.00	10.9	Upper Marker
22	115.63	13,222	0.00	0.0	
23	116.27	13,908	0.00	0.0	
24	117.09	14,790	0.00	0.0	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

Created: 2/7/2014 11:41:00 AM
Modified: 2/7/2014 12:23:14 PM

Electropherogram Summary Continued ...

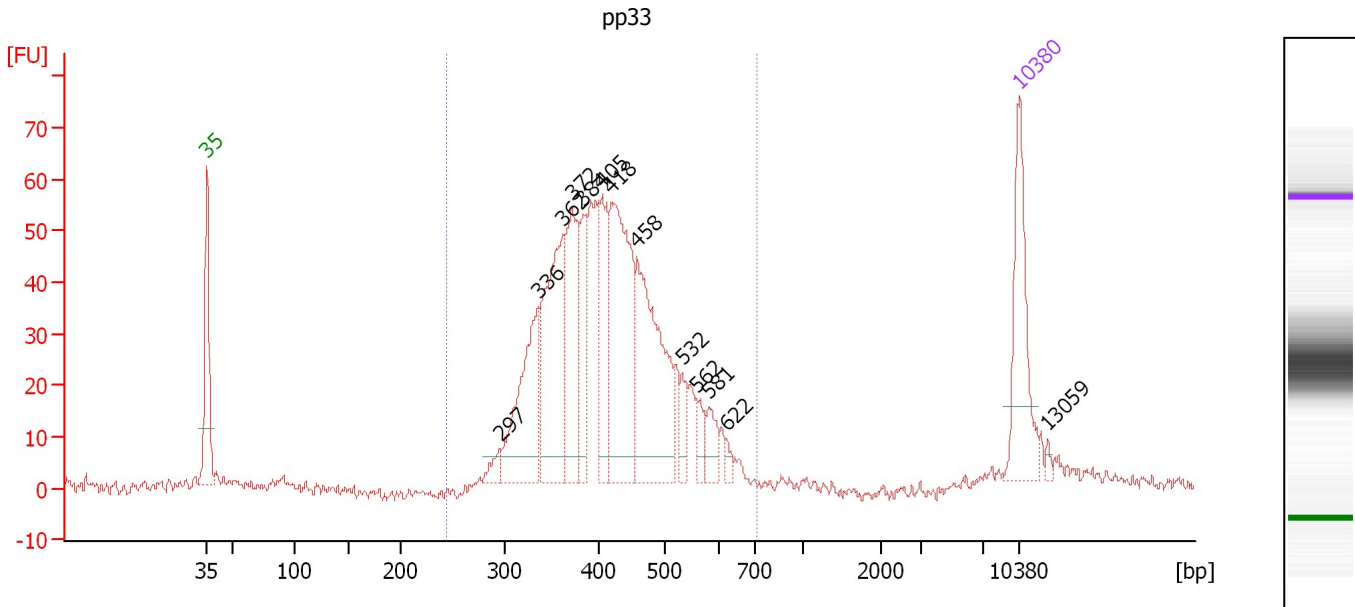
... Region table for sample 4 : pp32

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/ μ l]	Corr. Area	% of Total	Size distribution in CV [%]	Color
277	784	458	2,020.0	584.21	600.2	87	18.6	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

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 Modified: 2/7/2014 12:23:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : pp33

Number of peaks found: 13 Corr. Area 1: 876.3
 Noise: 1.0

Peak table for sample 5 : pp33

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	68.33	297	14.72	75.2	
3	71.59	336	133.24	600.3	
4	73.66	362	172.98	724.8	
5	74.48	372	113.47	462.4	
6	75.45	384	56.06	221.4	
7	77.06	405	78.10	292.4	
8	77.84	418	194.01	702.4	
9	80.09	458	181.36	599.9	
10	83.95	532	19.82	56.5	
11	85.33	562	13.26	35.8	
12	86.25	581	22.94	59.8	
13	87.81	622	5.95	14.5	
14	113.00	10,380	75.00	10.9	Upper Marker
15	115.48	13,059	0.00	0.0	

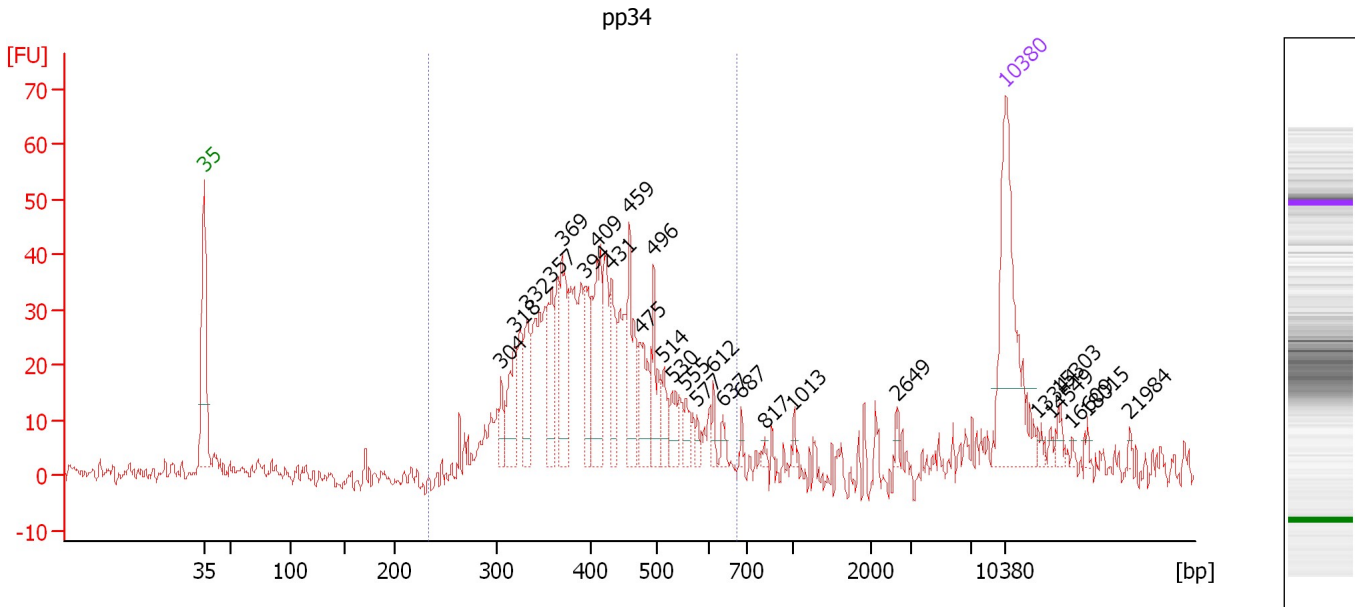
Region table for sample 5 : pp33

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
244	704	420	4,428.4	1,181.01	876.3	96	17.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

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 Modified: 2/7/2014 12:23:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : pp34

Number of peaks found: 27 Corr. Area 1: 603.5
 Noise: 1.7

Peak table for sample 6 : pp34

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	68.92	304	13.97	69.7	
3	70.09	318	33.70	160.6	
4	71.20	332	28.81	131.7	
5	73.30	357	30.43	129.1	
6	74.23	369	46.67	191.8	
7	76.28	394	29.29	112.7	
8	77.30	409	52.51	194.5	
9	78.56	431	27.00	94.9	
10	80.14	459	40.75	134.5	
11	81.07	475	31.04	99.0	
12	82.24	496	27.57	84.3	
13	83.12	514	13.02	38.4	
14	83.86	530	13.73	39.2	
15	85.03	555	9.12	24.9	
16	86.05	577	5.38	14.1	
17	87.49	612	5.59	13.8	
18	88.29	637	5.43	12.9	
19	89.91	687	3.48	7.7	
20	91.92	817	2.09	3.9	
21	94.48	1,013	3.10	4.6	
22	103.41	2,649	3.24	1.9	
23	113.00	10,380	75.00	10.9	Upper Marker
24	115.75	13,344	0.00	0.0	
25	116.86	14,549	0.00	0.0	


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

Created: 2/7/2014 11:41:00 AM
Modified: 2/7/2014 12:23:14 PM

Electropherogram Summary Continued ...**... Peak table for sample 6 : pp34**

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations
26	117.56	15,303	0.00	0.0	
27	118.77	16,609	0.00	0.0	
28	120.07	18,015	0.00	0.0	
29	123.75	21,984	0.00	0.0	

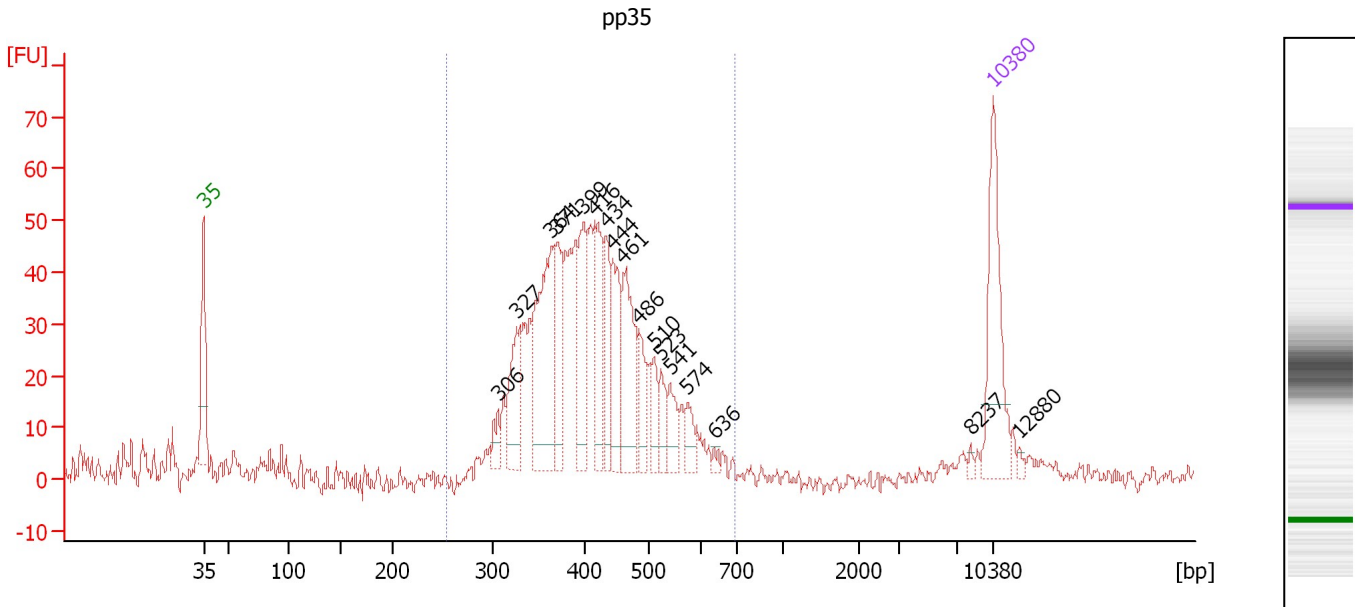
Region table for sample 6 : pp34

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/ μ l]	Corr. Area	% of Total	Size distribution in CV [%]	Color
233	676	412	2,726.5	707.80	603.5	89	18.9	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

Created: 2/7/2014 11:41:00 AM
 Modified: 2/7/2014 12:23:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : pp35

Number of peaks found: 17 Corr. Area 1: 707.7
 Noise: 1.8

Peak table for sample 7 : pp35

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	69.09	306	16.27	80.6	
3	70.79	327	54.22	251.6	
4	73.82	364	139.54	581.5	
5	74.38	371	61.18	250.2	
6	76.70	399	73.08	277.6	
7	77.69	416	64.92	236.5	
8	78.73	434	45.79	159.8	
9	79.30	444	59.12	201.7	
10	80.25	461	82.00	269.6	
11	81.66	486	26.18	81.7	
12	82.94	510	20.03	59.5	
13	83.55	523	20.66	59.8	
14	84.36	541	22.38	62.7	
15	85.92	574	16.89	44.6	
16	88.28	636	3.94	9.4	
17	111.01	8,237	2.50	0.5	
18	113.00	10,380	75.00	10.9	Upper Marker
19	115.32	12,880	0.00	0.0	

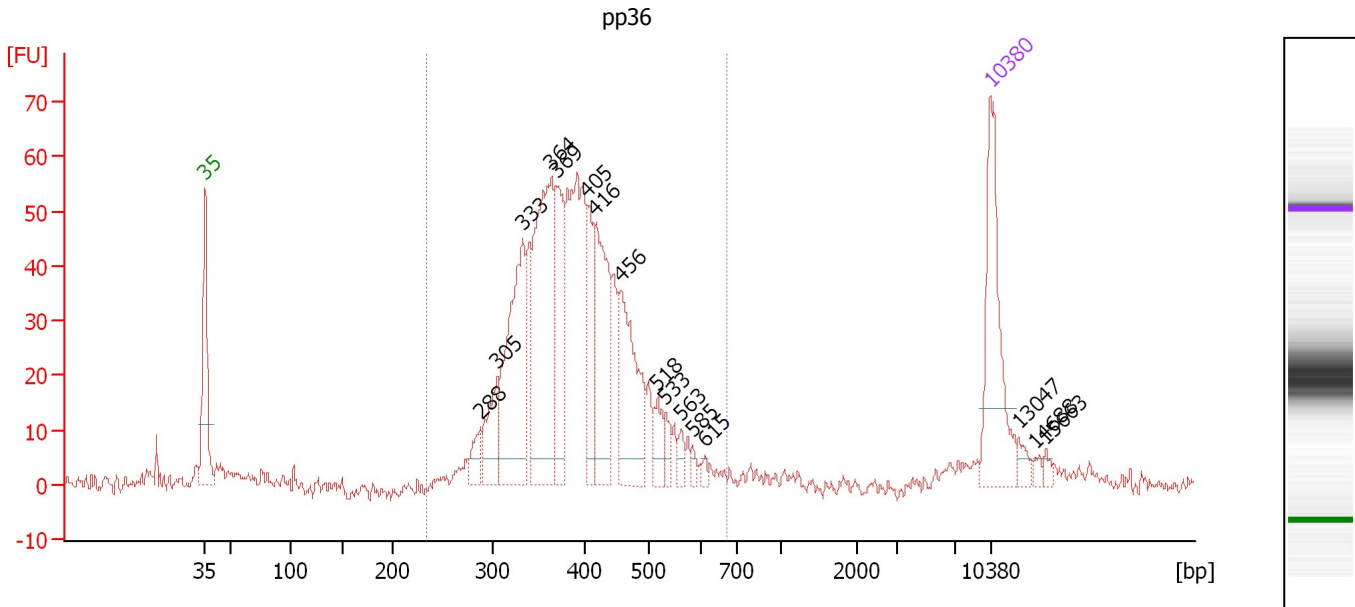
Region table for sample 7 : pp35

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
254	694	417	3,649.5	969.89	707.7	92	17.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

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 Modified: 2/7/2014 12:23:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : pp36

Number of peaks found: 16 Corr. Area 1: 833.7
 Noise: 1.4

Peak table for sample 8 : pp36

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	67.52	288	17.99	94.7	
3	69.04	305	45.58	226.3	
4	71.32	333	143.09	651.0	
5	73.84	364	181.63	756.2	
6	74.27	369	76.36	313.4	
7	77.07	405	47.68	178.4	
8	77.69	416	104.36	380.2	
9	79.97	456	96.54	320.8	
10	83.30	518	19.68	57.6	
11	84.01	533	10.04	28.5	
12	85.39	563	7.93	21.3	
13	86.44	585	4.59	11.9	
14	87.58	615	4.02	9.9	
15	113.00	10,380	75.00	10.9	Upper Marker
16	115.47	13,047	0.00	0.0	
17	116.99	14,688	0.00	0.0	
18	117.89	15,663	0.00	0.0	

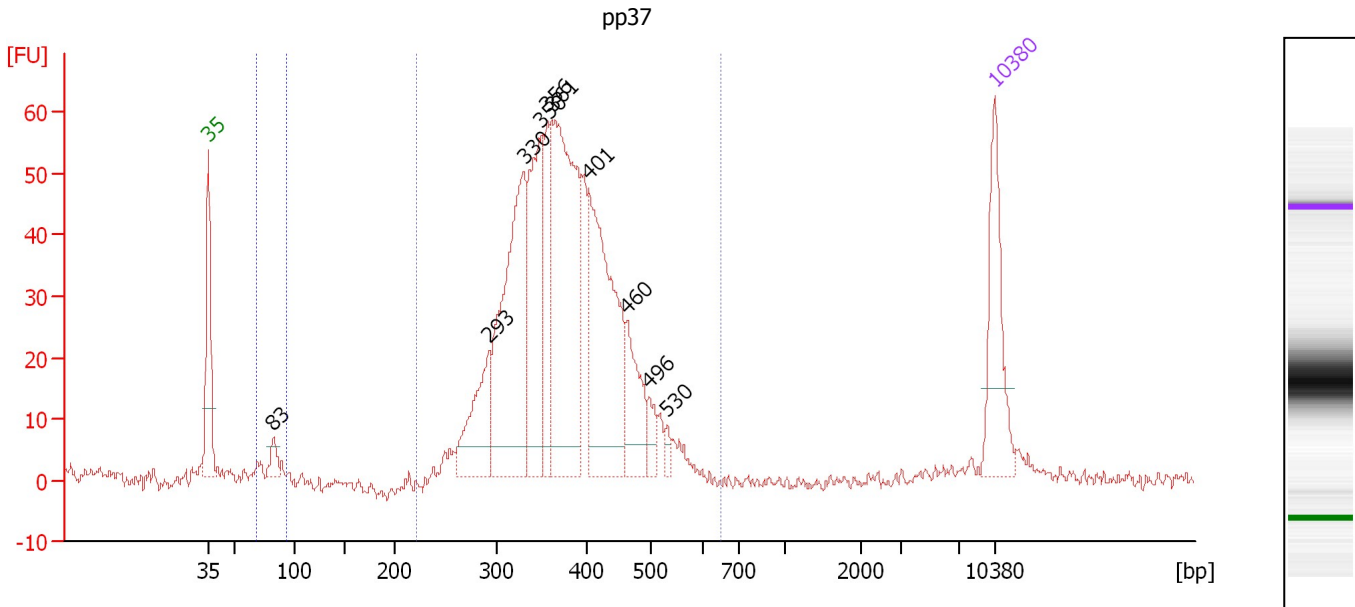
Region table for sample 8 : pp36

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
234	673	396	4,090.1	1,031.18	833.7	90	17.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

Created: 2/7/2014 11:41:00 AM
 Modified: 2/7/2014 12:23:14 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : pp37

Number of peaks found: 10 Corr. Area 1: 842.0
 Noise: 0.8 Corr. Area 2: 10.5

Peak table for sample 9 : pp37

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	48.79	83	20.04	364.9	
3	68.03	293	98.51	508.9	
4	71.11	330	270.01	1,237.9	
5	72.68	350	173.66	752.5	
6	73.20	356	78.38	333.5	
7	73.58	361	327.25	1,374.5	
8	76.86	401	235.68	889.8	
9	80.18	460	72.27	238.2	
10	82.23	496	19.55	59.8	
11	83.89	530	8.71	24.9	
12	113.00	10,380	75.00	10.9	Upper Marker

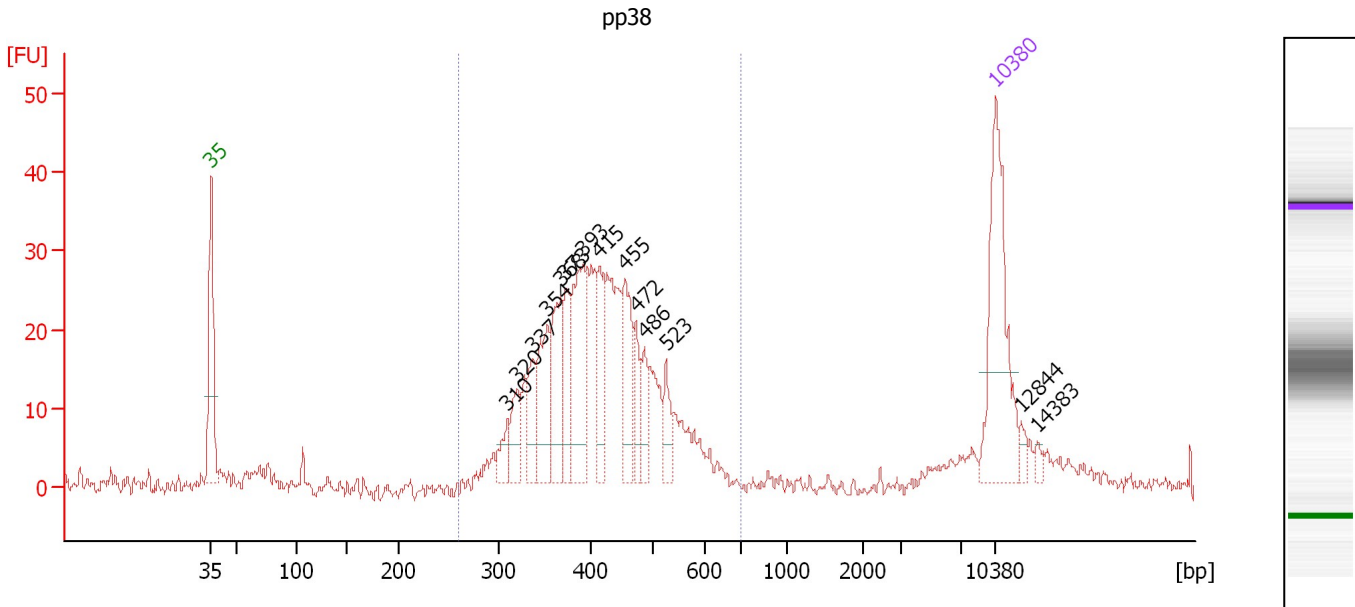
Region table for sample 9 : pp37

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
69	94	82	443.1	23.88	10.5	1	7.3	Blue
220	645	377	5,737.1	1,380.18	842.0	96	16.2	Dark Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 10 : pp38

Number of peaks found: 14 Corr. Area 1: 414.6
 Noise: 1.1

Peak table for sample 10 : pp38

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	69.44	310	15.40	75.3	
3	70.30	320	24.98	118.1	
4	71.68	337	24.92	111.9	
5	73.01	354	47.43	203.2	
6	74.15	368	51.69	213.0	
7	74.62	373	30.15	122.3	
8	76.24	393	72.36	278.8	
9	77.67	415	31.06	113.3	
10	79.90	455	42.91	143.0	
11	80.90	472	20.35	65.3	
12	81.66	486	17.62	55.0	
13	83.56	523	18.43	53.3	
14	113.00	10,380	75.00	10.9	Upper Marker
15	115.28	12,844	0.00	0.0	
16	116.71	14,383	0.00	0.0	

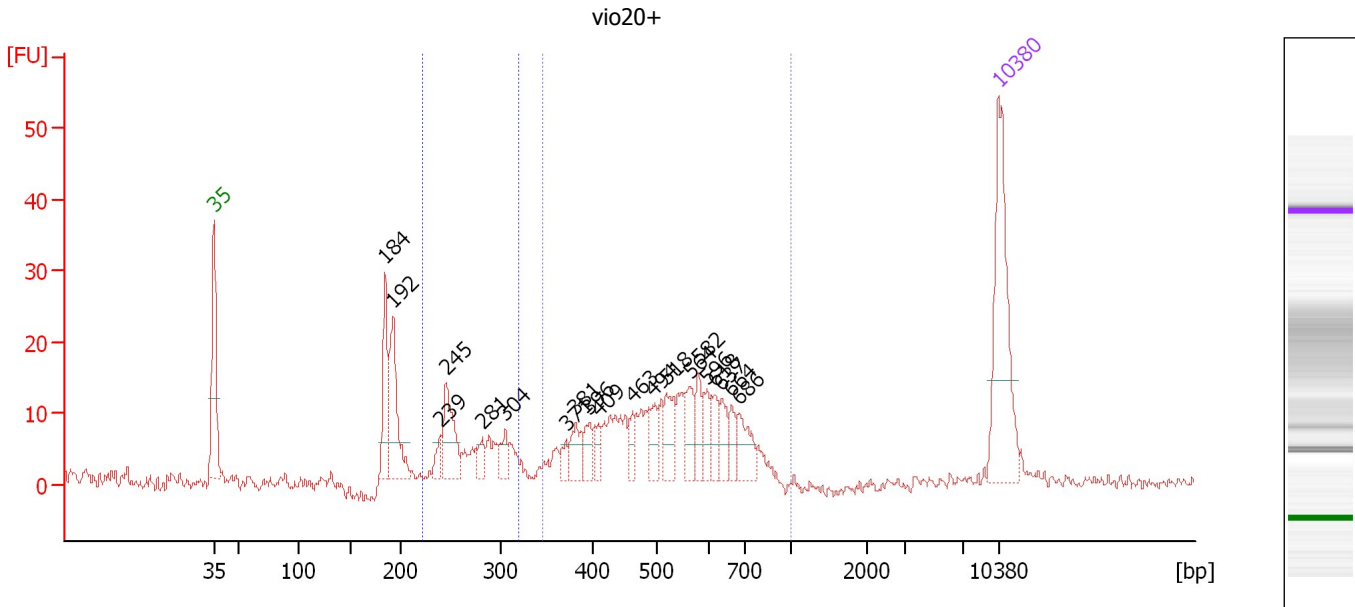
Region table for sample 10 : pp38

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
260	699	419	2,319.4	618.23	414.6	89	17.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

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Electropherogram Summary Continued ...



Overall Results for sample 11 : vio20+

Number of peaks found: 20 Corr. Area 1: 202.7
 Noise: 1.0 Corr. Area 2: 59.0

Peak table for sample 11 : vio20+



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	58.28	184	52.50	431.2	
3	58.99	192	64.17	505.3	
4	63.18	239	10.51	66.6	
5	63.70	245	44.20	273.4	
6	66.94	281	11.95	64.4	
7	68.98	304	12.76	63.5	
8	74.41	371	8.43	34.4	
9	75.26	381	18.08	71.9	
10	76.50	396	14.05	53.7	
11	77.31	409	10.83	40.1	
12	80.36	463	11.40	37.3	
13	82.16	494	17.72	54.3	
14	83.31	518	23.58	69.0	
15	85.45	564	20.92	56.2	
16	86.26	582	16.87	44.0	
17	86.92	596	14.14	36.0	
18	87.68	618	14.42	35.4	
19	88.30	637	14.64	34.8	
20	89.16	664	11.28	25.7	
21	89.87	686	23.08	51.0	
22	113.00	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

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Electropherogram Summary Continued ...

... Region table for sample 11 : vio20+

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
221	320	271	649.8	115.57	59.0	18	9.3	
346	1,000	534	993.5	330.30	202.7	62	20.9	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-07\2014-02-07_001.xad

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Gel Image

