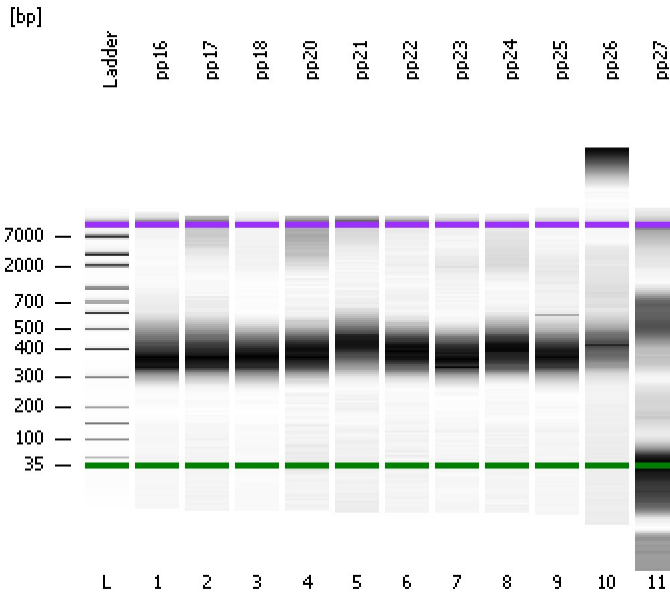


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
Modified: 2/6/2014 1:38:57 PM

Electrophoresis File Run Summary



Instrument Information:

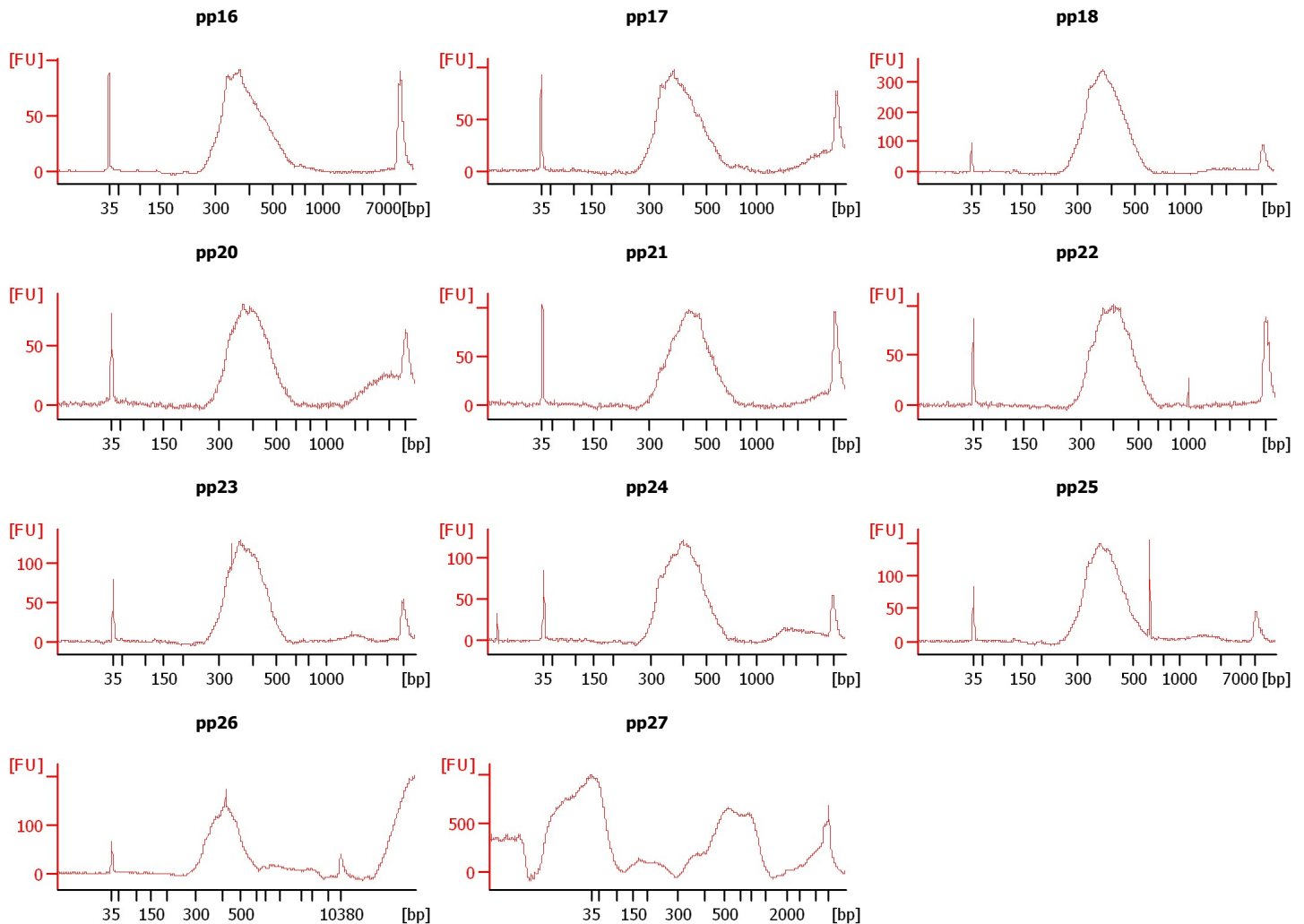
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
pp16		<input type="checkbox"/>	✓			
pp17		<input type="checkbox"/>	✓			
pp18		<input type="checkbox"/>	✓			
pp20		<input type="checkbox"/>	✓			
pp21		<input type="checkbox"/>	✓			
pp22		<input type="checkbox"/>	✓			
pp23		<input type="checkbox"/>	✓			
pp24		<input type="checkbox"/>	✓			
pp25		<input type="checkbox"/>	✓			
pp26		<input type="checkbox"/>	✓			
pp27		<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\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 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

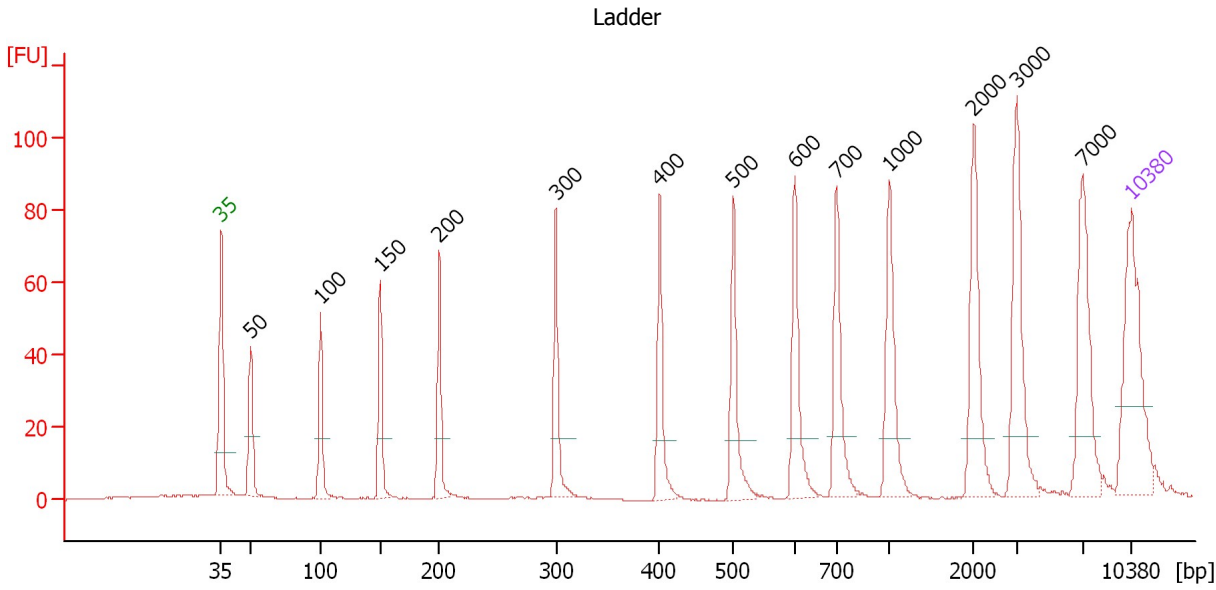
Ladder

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\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

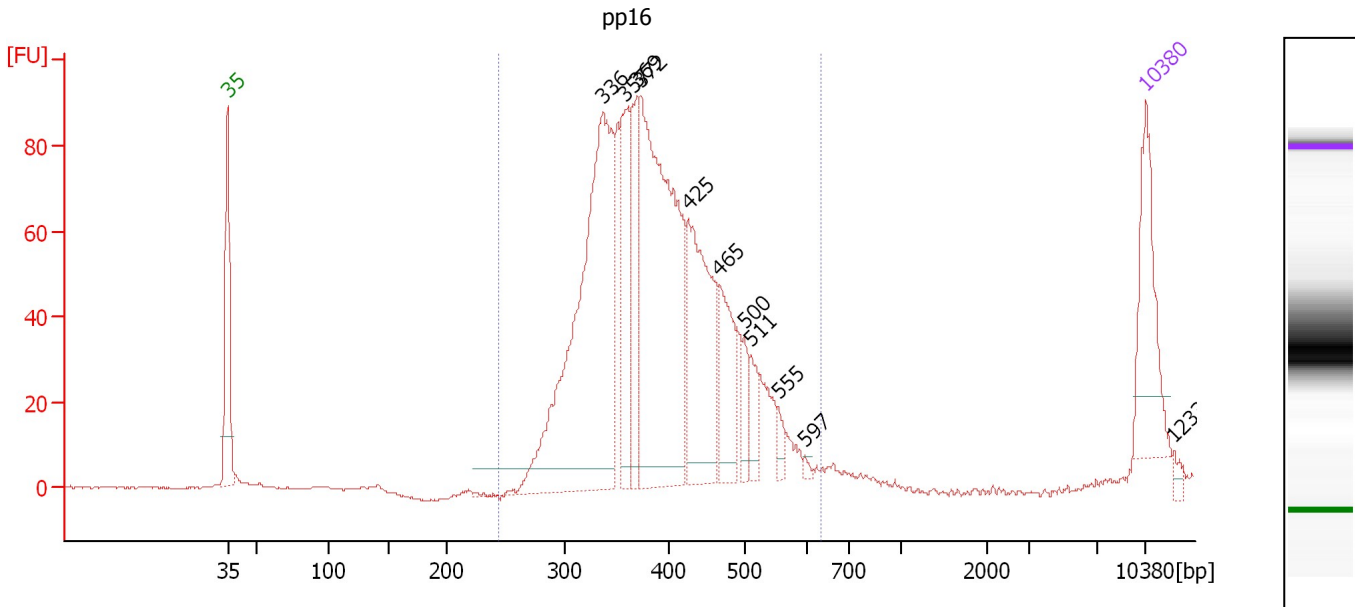
Peak table for Ladder

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	45.26	50	150.00	4,545.5	Ladder Peak
3	50.67	100	150.00	2,272.7	Ladder Peak
4	55.22	150	150.00	1,515.2	Ladder Peak
5	59.77	200	150.00	1,136.4	Ladder Peak
6	68.75	300	150.00	757.6	Ladder Peak
7	76.71	400	150.00	568.2	Ladder Peak
8	82.41	500	150.00	454.5	Ladder Peak
9	87.12	600	150.00	378.8	Ladder Peak
10	90.32	700	150.00	324.7	Ladder Peak
11	94.38	1,000	150.00	227.3	Ladder Peak
12	100.90	2,000	150.00	113.6	Ladder Peak
13	104.18	3,000	150.00	75.8	Ladder Peak
14	109.27	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\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : pp16

Number of peaks found: 11 Corr. Area 1: 1,388.0
 Noise: 0.2

Peak table for sample 1 : pp16

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	71.62	336	484.31	2,183.9	
3	73.33	357	108.47	459.7	
4	74.26	369	85.35	350.3	
5	74.50	372	411.46	1,674.5	
6	78.12	425	180.44	643.6	
7	80.44	465	78.86	256.7	
8	82.39	500	22.76	69.0	
9	82.92	511	26.61	78.9	
10	84.99	555	12.95	35.4	
11	86.98	597	3.23	8.2	
12	113.00	10,380	75.00	10.9	Upper Marker
13	115.15	12,331	0.00	0.0	

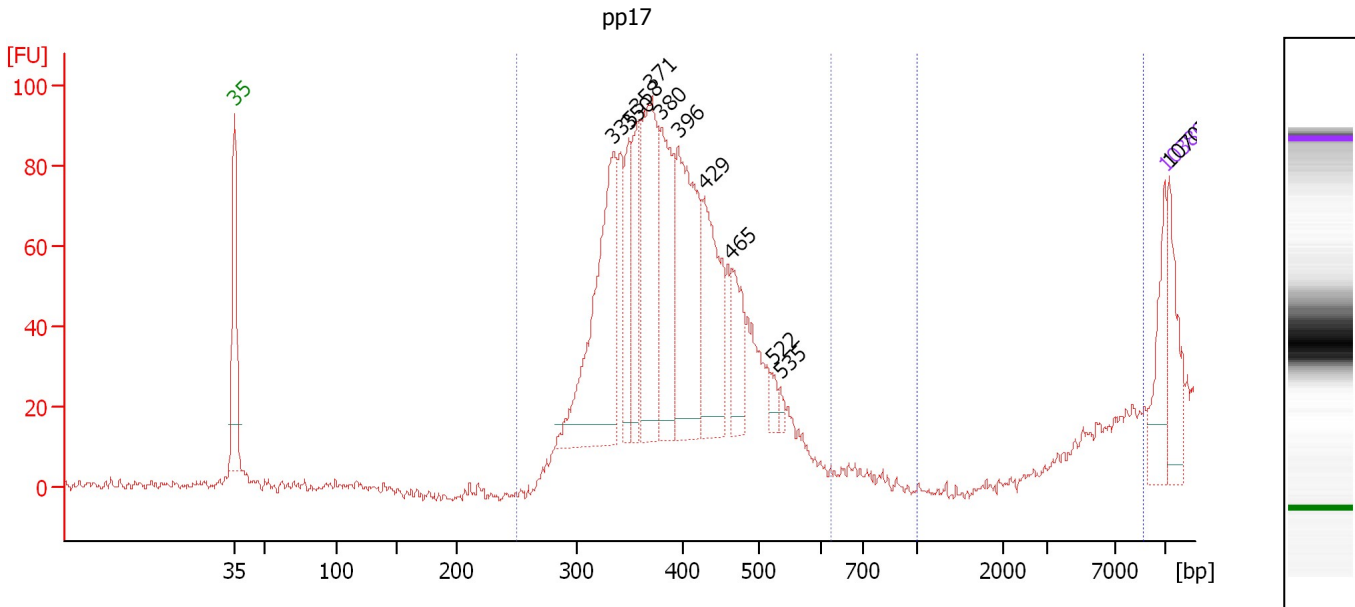
Region table for sample 1 : pp16

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
243	634	394	6,005.3	1,505.37	1,388.0	99	17.2	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : pp17

Number of peaks found: 11 Corr. Area 1: 1,056.7
 Noise: 1.1 Corr. Area 2: 0.0

Peak table for sample 2 : pp17

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	71.54	335	435.23	1,968.4	
3	72.70	350	120.28	521.2	
4	73.34	358	134.18	568.3	
5	74.39	371	319.00	1,303.4	
6	75.15	380	241.70	962.7	
7	76.39	396	338.10	1,293.4	
8	78.36	429	226.51	800.0	
9	80.41	465	99.19	323.3	
10	83.46	522	20.10	58.3	
11	84.06	535	10.88	30.8	
12	113.00	10,380	75.00	10.9	Upper Marker
13	113.36	10,707	0.00	0.0	

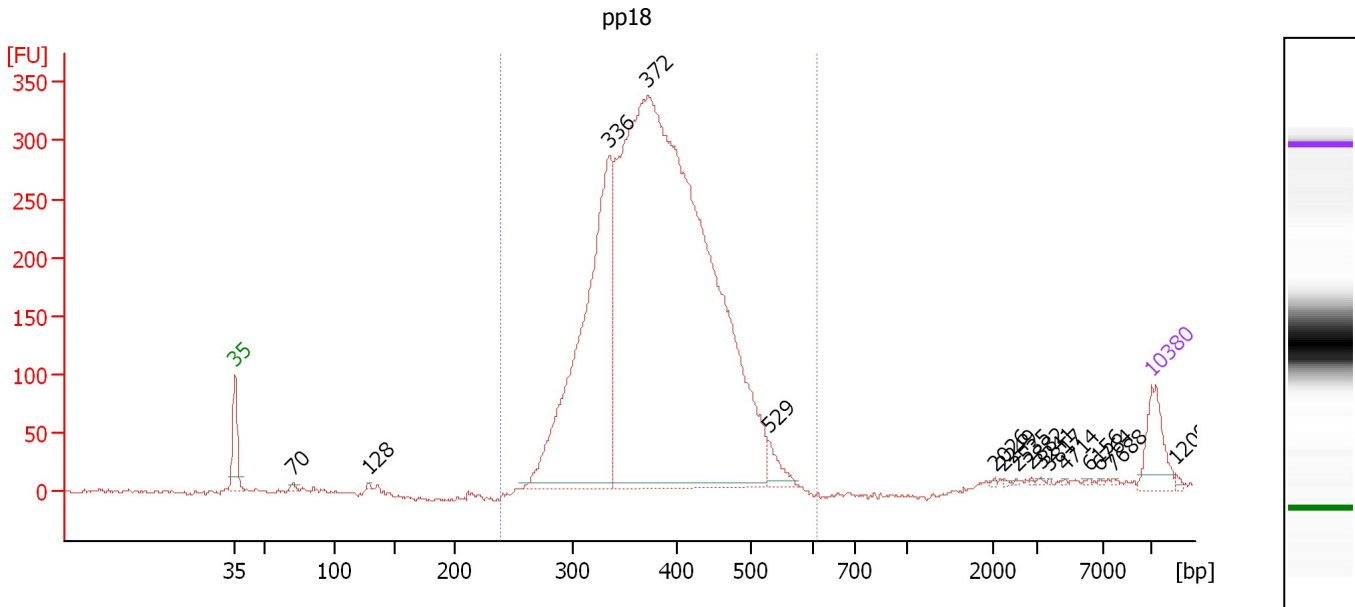
Region table for sample 2 : pp17

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
250	622	388	8,184.1	2,057.15	1,056.7	97	13.3	Blue
996	8,931	0	0.0	0.00	0.0	0	0.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : pp18

Number of peaks found: 16 Corr. Area 1: 4,891.2
 Noise: 1.6

Peak table for sample 3 : pp18

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	47.44	70	7.28	157.2	
3	53.18	128	4.42	52.5	
4	71.59	336	1,056.59	4,770.0	
5	74.52	372	3,293.85	13,398.8	
6	83.76	529	43.71	125.3	
7	100.99	2,026	1.85	1.4	
8	101.72	2,249	2.24	1.5	
9	102.66	2,535	1.47	0.9	
10	103.80	2,882	1.79	0.9	
11	104.49	3,241	2.56	1.2	
12	105.22	3,817	1.44	0.6	
13	106.36	4,714	1.35	0.4	
14	108.19	6,156	2.21	0.5	
15	108.97	6,764	1.58	0.4	
16	110.03	7,688	1.76	0.3	
17	113.00	10,380	75.00	10.9	Upper Marker
18	114.79	12,003	0.00	0.0	

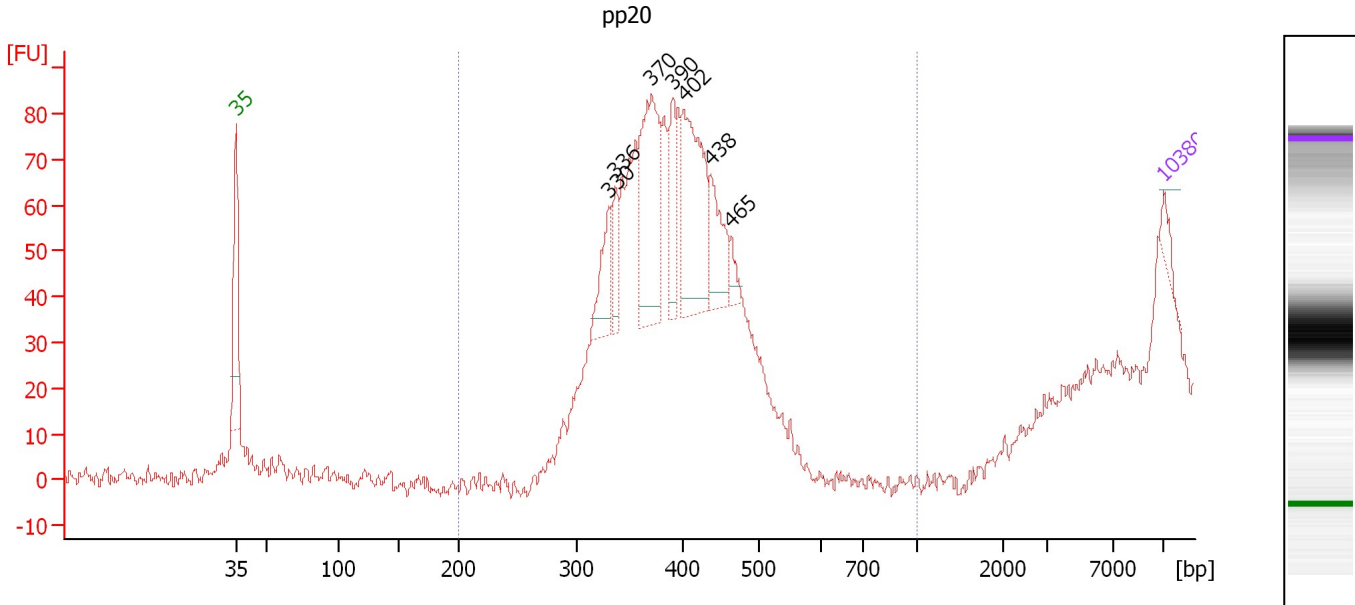
Region table for sample 3 : pp18

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
239	611	385	17,689.1	4,371.57	4,891.2	97	14.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 4 : pp20

Height Threshold [FU] : 4

Overall Results for sample 4 : pp20

Number of peaks found: 7 Corr. Area 1: 927.6
 Noise: 2.2

Peak table for sample 4 : pp20

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	71.14	330	451.32	2,072.4	
3	71.62	336	279.35	1,259.6	
4	74.36	370	1,299.42	5,314.7	
5	75.93	390	423.59	1,644.9	
6	76.81	402	1,239.41	4,673.4	
7	78.87	438	496.08	1,716.7	
8	80.40	465	115.62	377.0	
9	113.00	10,380	75.00	10.9	Upper Marker

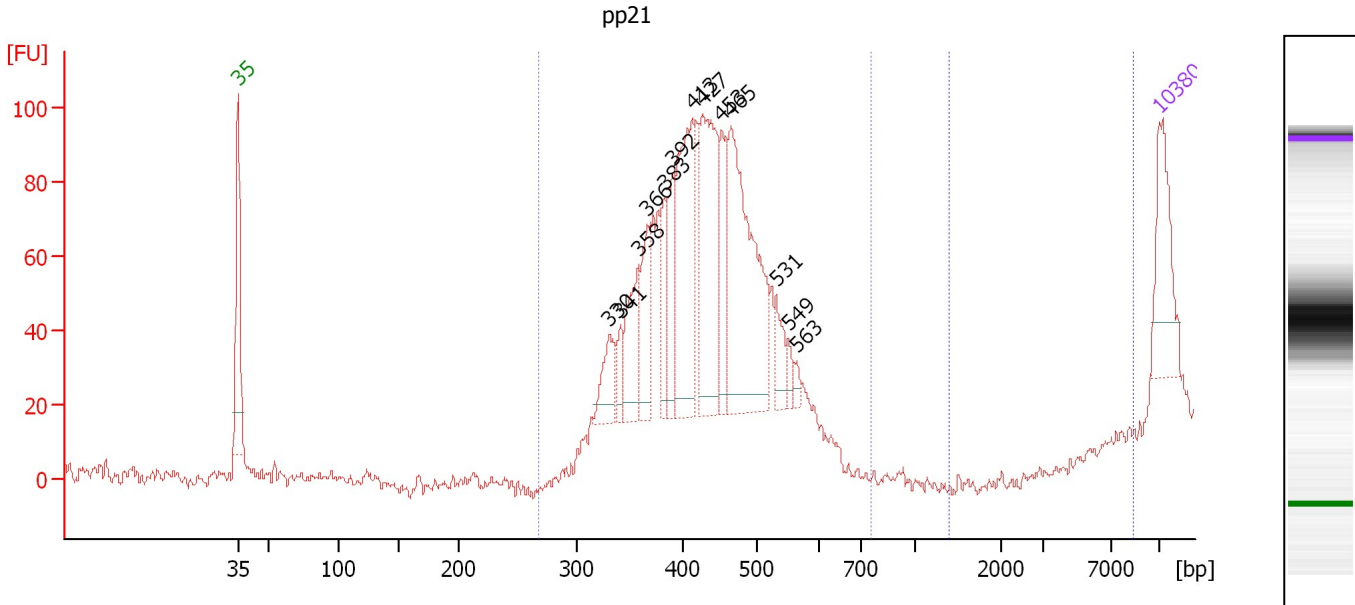
Region table for sample 4 : pp20

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
200	1,000	390	41,717.8	10,570.26	927.6	96	12.3	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : pp21

Number of peaks found: 13 Corr. Area 1: 1,105.0
 Noise: 2.0 Corr. Area 2: 0.0

Peak table for sample 5 : pp21

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	71.18	330	54.53	250.0	
3	72.03	341	26.03	115.6	
4	73.40	358	82.53	348.8	
5	74.01	366	70.89	293.4	
6	75.35	383	57.18	226.2	
7	76.04	392	65.40	253.1	
8	77.45	413	197.23	723.4	
9	78.22	427	211.69	751.9	
10	79.72	453	67.75	226.7	
11	80.41	465	302.17	984.7	
12	83.85	531	33.54	95.8	
13	84.74	549	12.88	35.5	
14	85.39	563	9.31	25.0	
15	113.00	10,380	75.00	10.9	Upper Marker

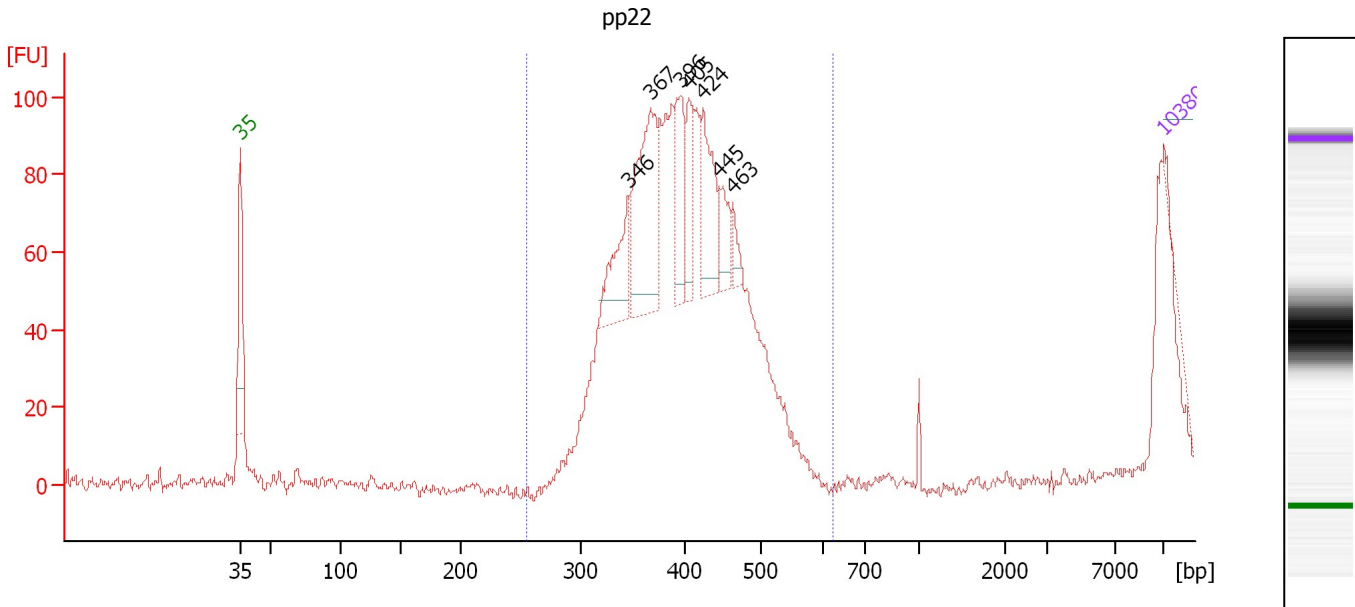
Region table for sample 5 : pp21

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
268	757	429	5,293.6	1,464.62	1,105.0	97	13.7	Blue
1,388	8,552	0	0.0	0.00	0.0	0	0.0	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : pp22

Number of peaks found: 7 Corr. Area 1: 1,237.0
 Noise: 1.5

Peak table for sample 6 : pp22

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	72.38	346	1,971.13	8,641.1	
3	74.08	367	4,290.47	17,715.1	
4	76.42	396	1,970.57	7,531.6	
5	76.99	405	1,288.92	4,822.8	
6	78.08	424	2,167.98	7,745.7	
7	79.29	445	851.55	2,897.1	
8	80.30	463	425.99	1,393.8	
9	113.00	10,380	75.00	10.9	Upper Marker

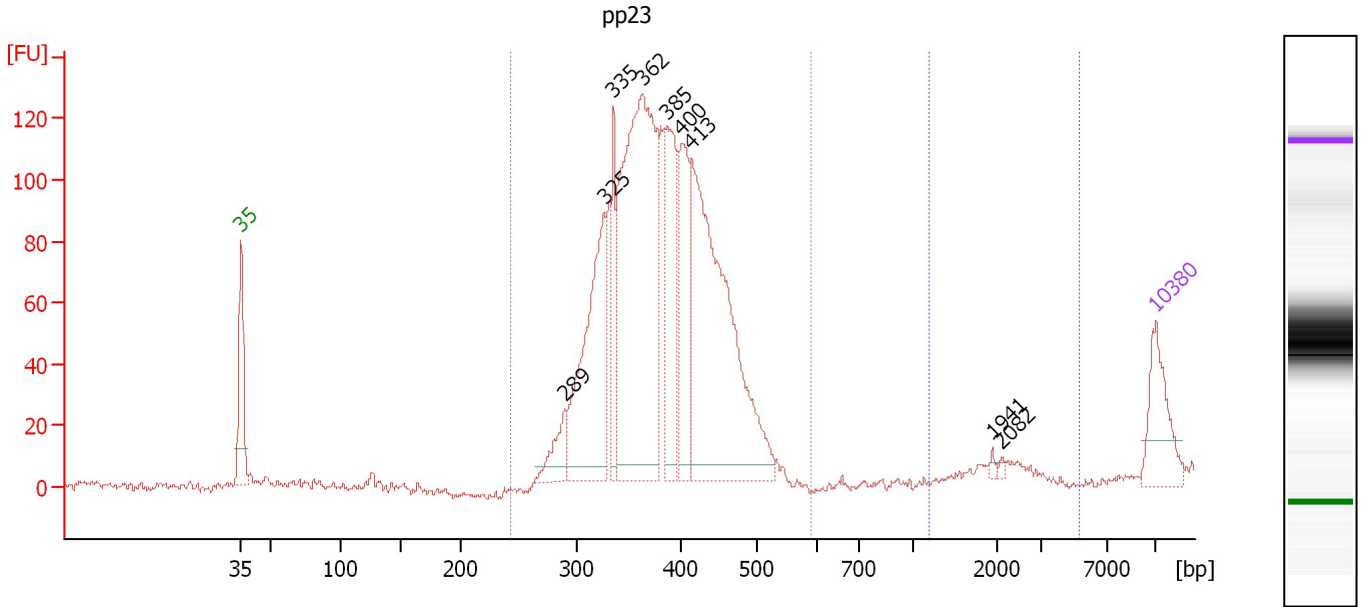
Region table for sample 6 : pp22

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
255	626	400	152,956.2	39,564.13	1,237.0	97	13.1	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : pp23

Number of peaks found: 9 Corr. Area 1: 1,704.7
 Noise: 1.3 Corr. Area 2: 9.0

Peak table for sample 7 : pp23

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	67.76	289	61.00	319.9	
3	70.78	325	355.72	1,655.9	
4	71.51	335	114.98	520.5	
5	73.72	362	720.47	3,012.2	
6	75.55	385	220.95	868.5	
7	76.69	400	179.90	681.7	
8	77.47	413	629.49	2,307.4	
9	100.52	1,941	3.98	3.1	
10	101.17	2,082	3.89	2.8	
11	113.00	10,380	75.00	10.9	Upper Marker

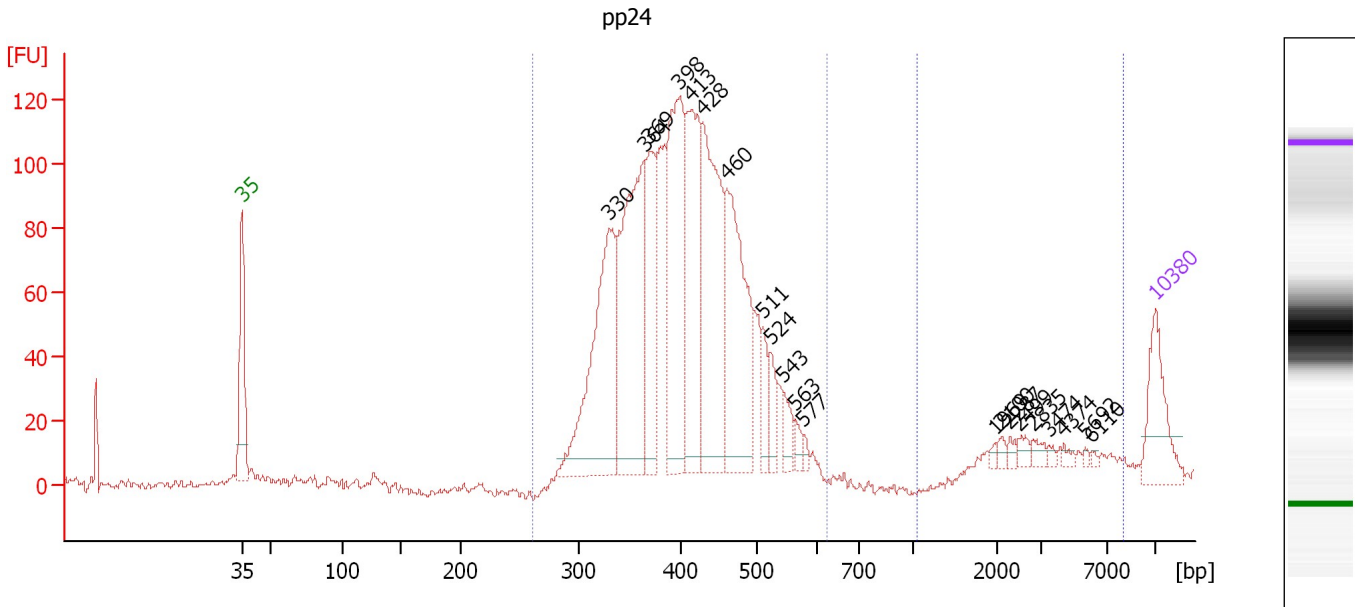
Region table for sample 7 : pp23

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
243	589	382	9,793.4	2,415.71	1,704.7	97	13.6	Blue
1,195	5,324	2,113	6.6	9.10	9.0	1	11.9	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : pp24

Number of peaks found: 21 Corr. Area 1: 1,759.1
 Noise: 1.4 Corr. Area 2: 92.2



Peak table for sample 8 : pp24

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	71.16	330	366.06	1,679.7	
3	73.81	364	403.44	1,681.2	
4	74.22	369	178.82	734.8	
5	76.55	398	324.16	1,234.0	
6	77.45	413	264.54	970.5	
7	78.31	428	334.84	1,185.2	
8	80.15	460	289.83	954.0	
9	82.92	511	39.23	116.3	
10	83.54	524	42.51	122.9	
11	84.44	543	32.60	91.0	
12	85.38	563	14.37	38.7	
13	86.03	577	8.36	22.0	
14	100.70	1,969	4.41	3.4	
15	101.23	2,100	6.38	4.6	
16	101.84	2,287	5.90	3.9	
17	102.54	2,499	9.45	5.7	
18	103.64	2,835	9.96	5.3	
19	104.79	3,474	4.63	2.0	
20	105.93	4,374	5.82	2.0	
21	107.61	5,692	2.71	0.7	
22	108.14	6,110	2.39	0.6	
23	113.00	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
Modified: 2/6/2014 1:38:57 PM

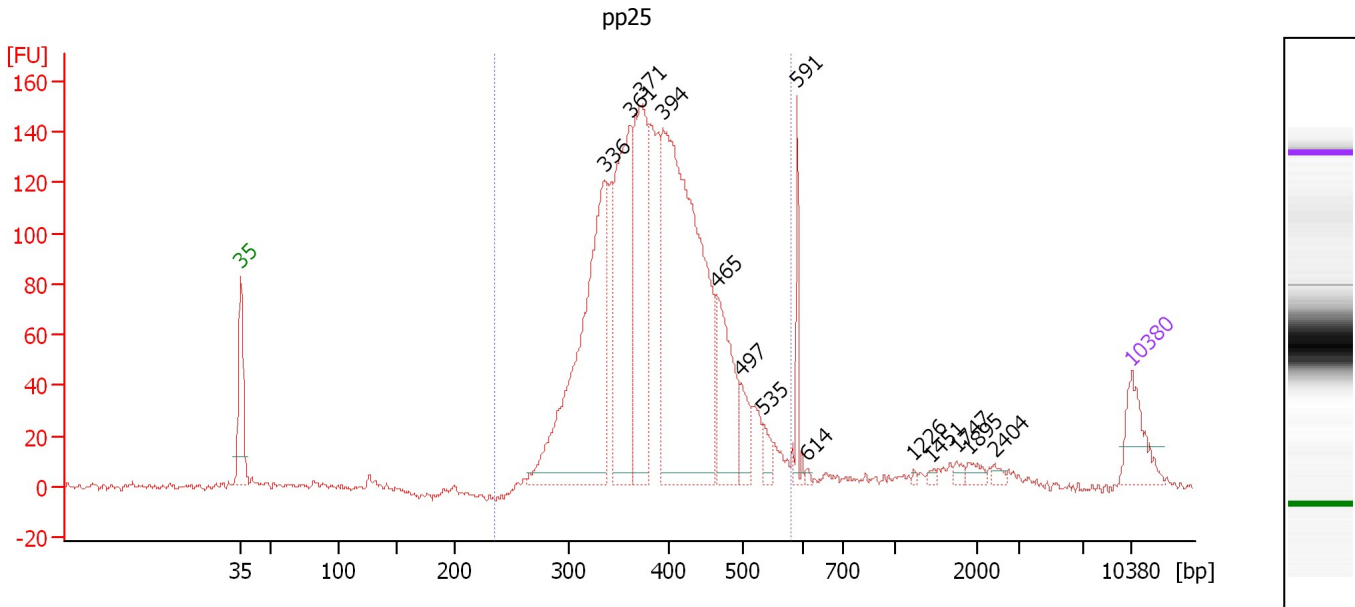
Electropherogram Summary Continued ...**... Region table for sample 8 : pp24**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
262	622	409	9,851.6	2,580.48	1,759.1	93	15.5	
1,046	8,203	3,713	48.9	98.78	92.2	5	48.5	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : pp25

Number of peaks found: 14 Corr. Area 1: 2,163.7
 Noise: 1.2

Peak table for sample 9 : pp25

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	71.59	336	791.59	3,573.6	
3	73.64	361	473.45	1,984.7	
4	74.40	371	466.29	1,904.7	
5	76.20	394	1,094.82	4,214.7	
6	80.39	465	225.33	734.9	
7	82.23	497	65.41	199.4	
8	84.04	535	30.81	87.3	
9	86.72	591	62.96	161.3	
10	87.56	614	5.05	12.5	
11	95.86	1,226	3.44	4.3	
12	97.32	1,451	6.16	6.4	
13	99.25	1,747	9.52	8.3	
14	100.22	1,895	15.20	12.2	
15	102.23	2,404	8.63	5.4	
16	113.00	10,380	75.00	10.9	Upper Marker

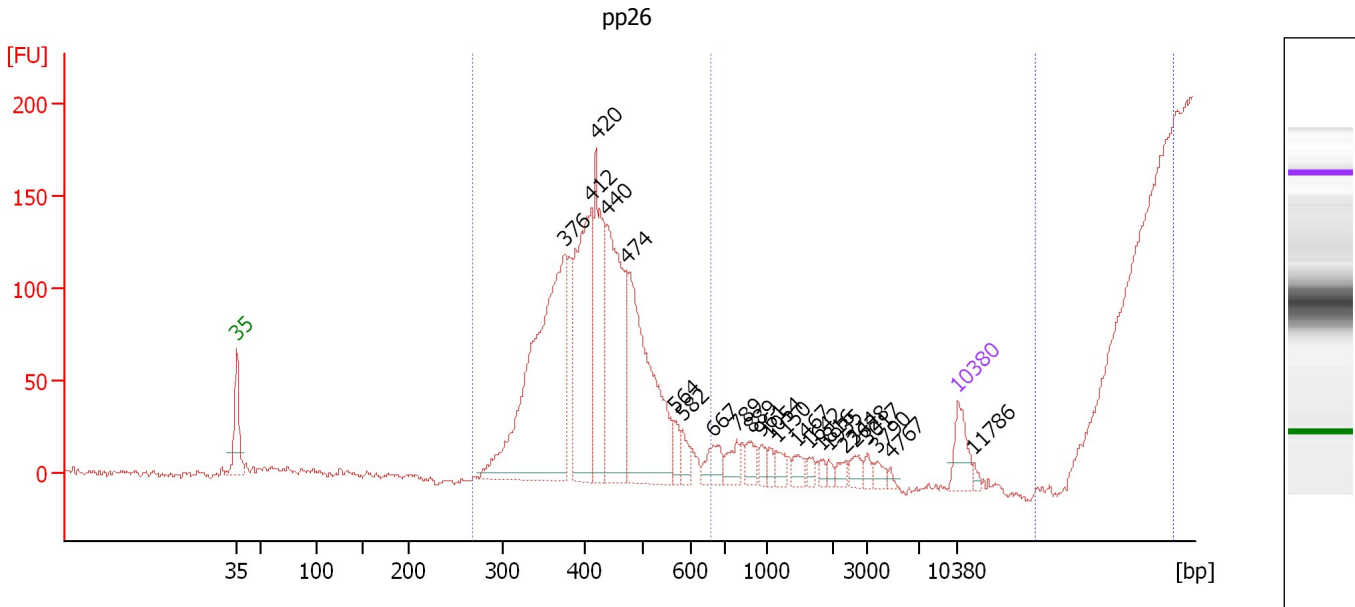
Region table for sample 9 : pp25

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
235	583	391	14,854.7	3,720.93	2,163.7	92	15.4	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

Created: 2/6/2014 12:49:32 PM
 Modified: 2/6/2014 1:38:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : pp26

Number of peaks found: 23 Corr. Area 1: 265.5
 Noise: 1.9 Corr. Area 2: 0.0



Peak table for sample 10 : pp26

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	74.77	376	1,163.05	4,691.8	
3	77.41	412	613.53	2,255.1	
4	77.87	420	428.11	1,543.0	
5	79.01	440	590.14	2,030.5	
6	80.92	474	687.16	2,196.8	
7	85.42	564	43.20	116.1	
8	86.25	582	45.24	117.9	
9	89.25	667	72.13	164.0	
10	91.53	789	55.34	106.3	
11	92.87	889	45.91	78.3	
12	93.86	961	28.65	45.2	
13	94.74	1,054	21.17	30.4	
14	95.36	1,150	33.27	43.8	
15	97.43	1,467	30.62	31.6	
16	98.57	1,642	16.76	15.5	
17	99.70	1,816	14.12	11.8	
18	100.48	1,935	14.31	11.2	
19	102.03	2,344	20.38	13.2	
20	102.96	2,628	29.34	16.9	
21	104.20	3,017	18.06	9.1	
22	105.19	3,790	23.77	9.5	
23	106.43	4,767	8.89	2.8	
24	113.00	10,380	75.00	10.9	Upper Marker
25	114.55	11,786	0.00	0.0	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

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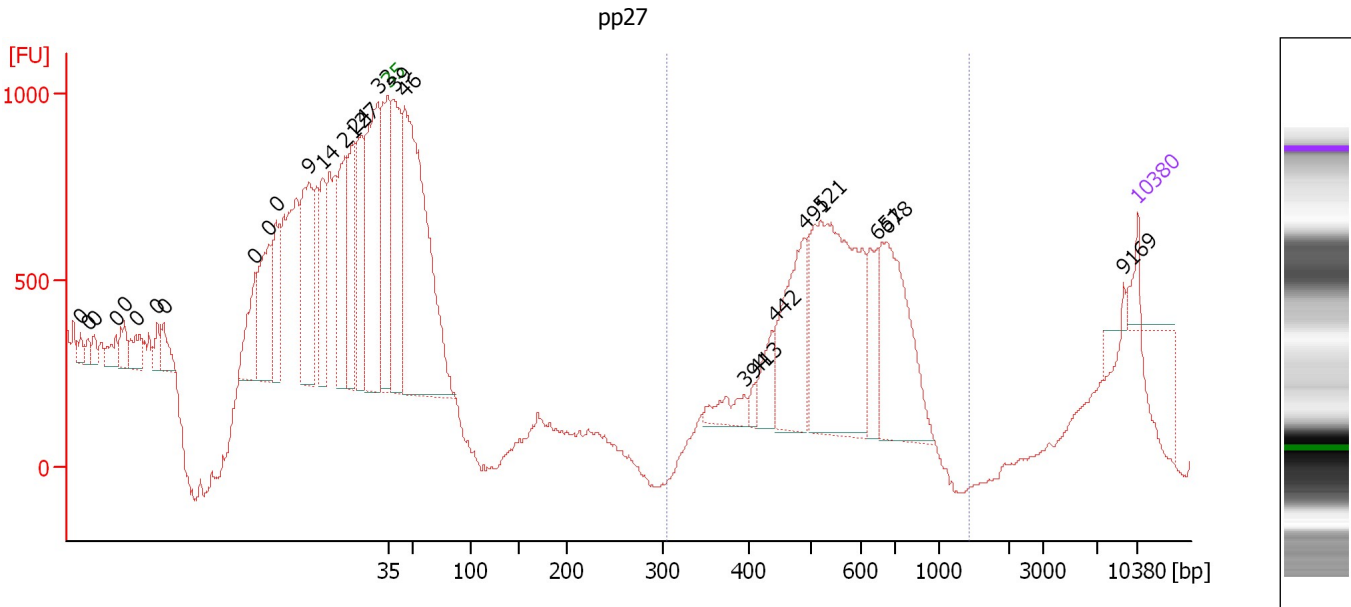
Electropherogram Summary Continued ...**... Region table for sample 10 : pp26**

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
269	659	412	1,936.0	524.11	265.5	100	6.8	
17,228	29,314	0	0.0	0.00	0.0	0	0.0	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

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 Modified: 2/6/2014 1:38:57 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 11 : pp27

Height Threshold [FU] : 1

Overall Results for sample 11 : pp27

Number of peaks found: 27 Corr. Area 1: 6,988.1
 Noise: 8.5

Peak table for sample 11 : pp27


Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	14.34	0	0.00	0.0	
2	15.03	0	0.00	0.0	
3	15.57	0	0.00	0.0	
4	17.60	0	0.00	0.0	
5	18.45	0	0.00	0.0	
6	19.49	0	0.00	0.0	
7	21.47	0	0.00	0.0	
8	22.06	0	0.00	0.0	
9	30.62	0	0.00	0.0	
10	31.91	0	0.00	0.0	
11	32.60	0	0.00	0.0	
12	35.62	9	0.00	0.0	
13	36.91	14	0.00	0.0	
14	38.99	21	0.00	0.0	
15	39.93	24	0.00	0.0	
16	40.57	27	0.00	0.0	
17	42.06	32	0.00	0.0	
18	43.00	35	125.00	5,411.3	Lower Marker
19	43.59	39	1,283.65	49,932.7	
20	44.58	46	3,145.49	104,662.6	
21	76.27	394	179.42	689.2	
22	77.46	413	50.25	184.3	

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

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Electropherogram Summary Continued ...

... Peak table for sample 11 : pp27

Peak	Aligned Migration Time [s]	Size [bp]	Conc. [pg/ μ l]	Molarity [pmol/l]	Observations
23	79.09	442	217.11	744.6	
24	81.91	491	821.05	2,532.2	
25	83.40	521	1,798.62	5,231.2	
26	88.74	651	319.88	745.0	
27	89.63	678	951.50	2,124.9	
28	111.66	9,169	17.42	2.9	
29	 113.00	10,380	75.00	10.9	Upper Marker

Region table for sample 11 : pp27

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/ μ l]	Corr. Area	% of Total	Size distribution in CV [%]	Color
303	1,432	578	11,806.7	4,284.49	6,988.1	29	20.2	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-02-06\2014-02-06_003.xad

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Gel Image

