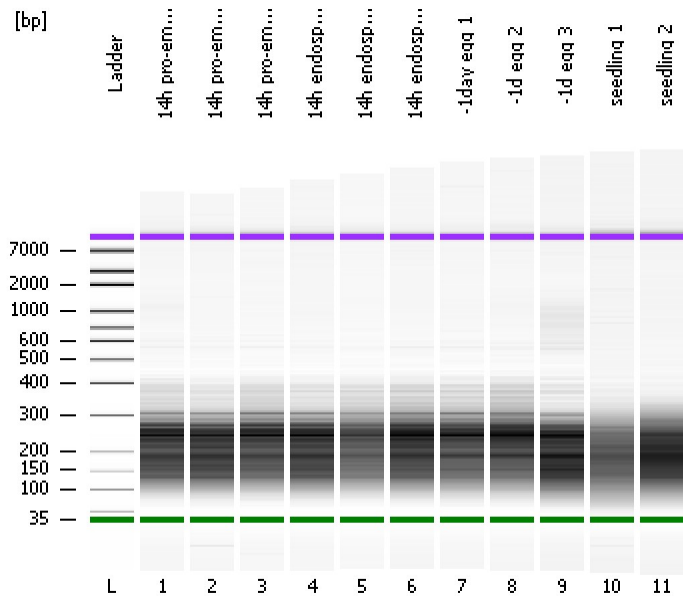


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
Modified: 4/17/2014 5:09:53 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

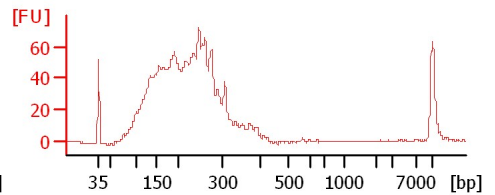
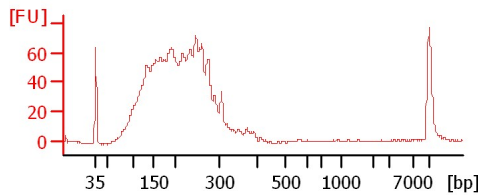
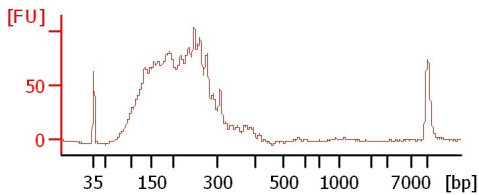
Chip Information:

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

14h pro-embryo 1

14h pro-embryo 2

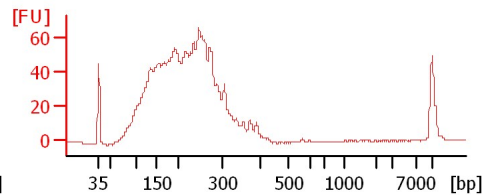
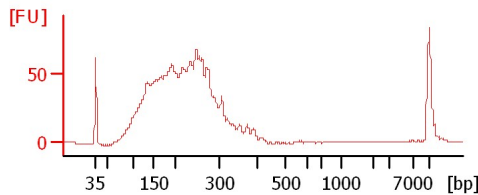
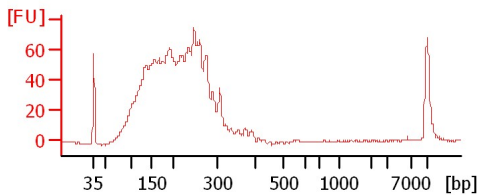
14h pro-embryo 3



14h endosperm 2

14h endosperm 3

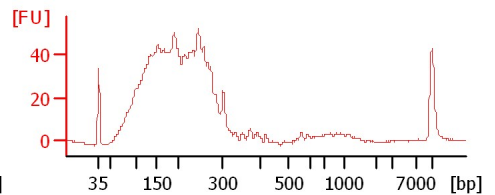
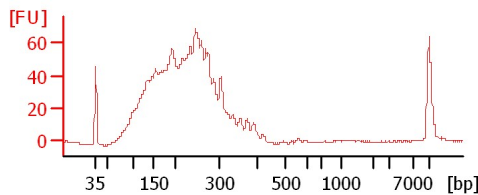
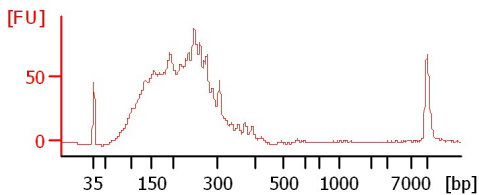
14h endosperm 4



-1day egg 1

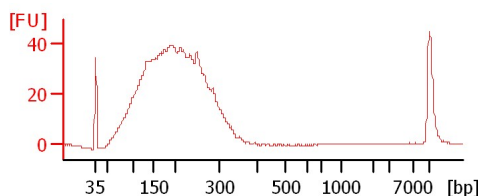
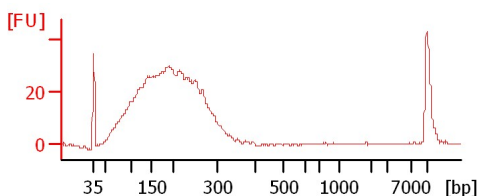
-1d egg 2

-1d egg 3



seedling 1

seedling 2



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
14h pro-embryo 1		<input type="checkbox"/>	✓			
14h pro-embryo 2		<input type="checkbox"/>	✓			
14h pro-embryo 3		<input type="checkbox"/>	✓			
14h endosperm 2		<input type="checkbox"/>	✓			
14h endosperm 3		<input type="checkbox"/>	✓			
14h endosperm 4		<input type="checkbox"/>	✓			
-1day egg 1		<input type="checkbox"/>	✓			
-1d egg 2		<input type="checkbox"/>	✓			
-1d egg 3		<input type="checkbox"/>	✓			
seedling 1		<input type="checkbox"/>	✓			
seedling 2		<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\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
Modified: 4/17/2014 5:09:53 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

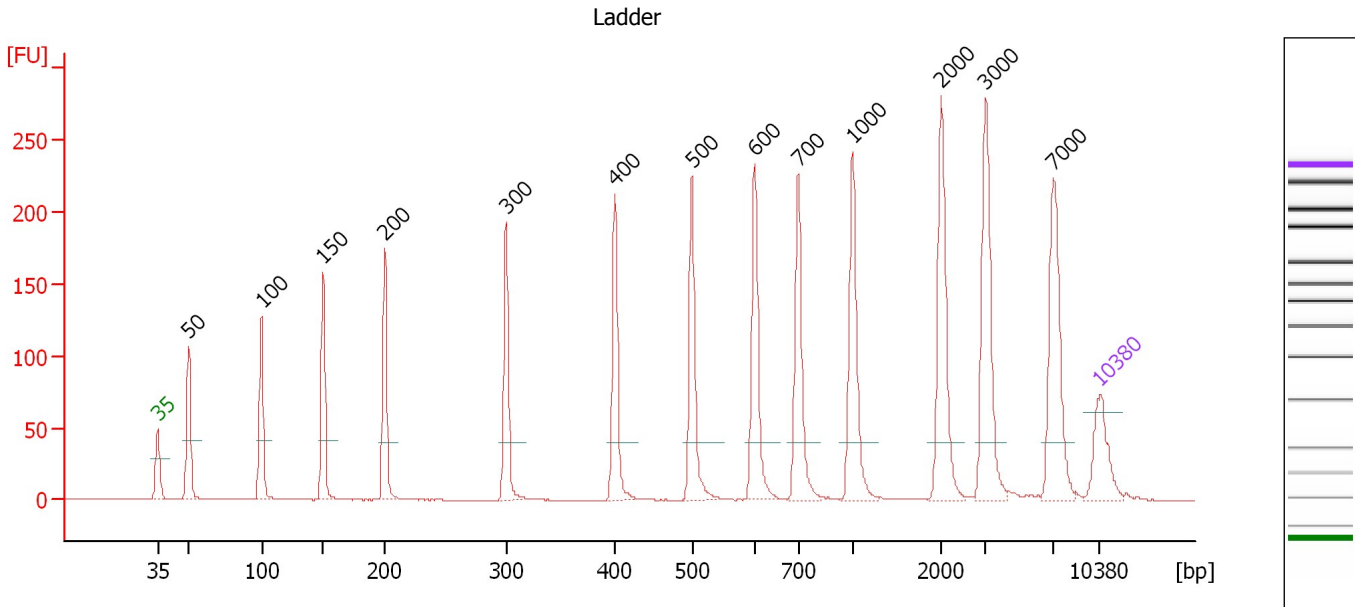
Ladder

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\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

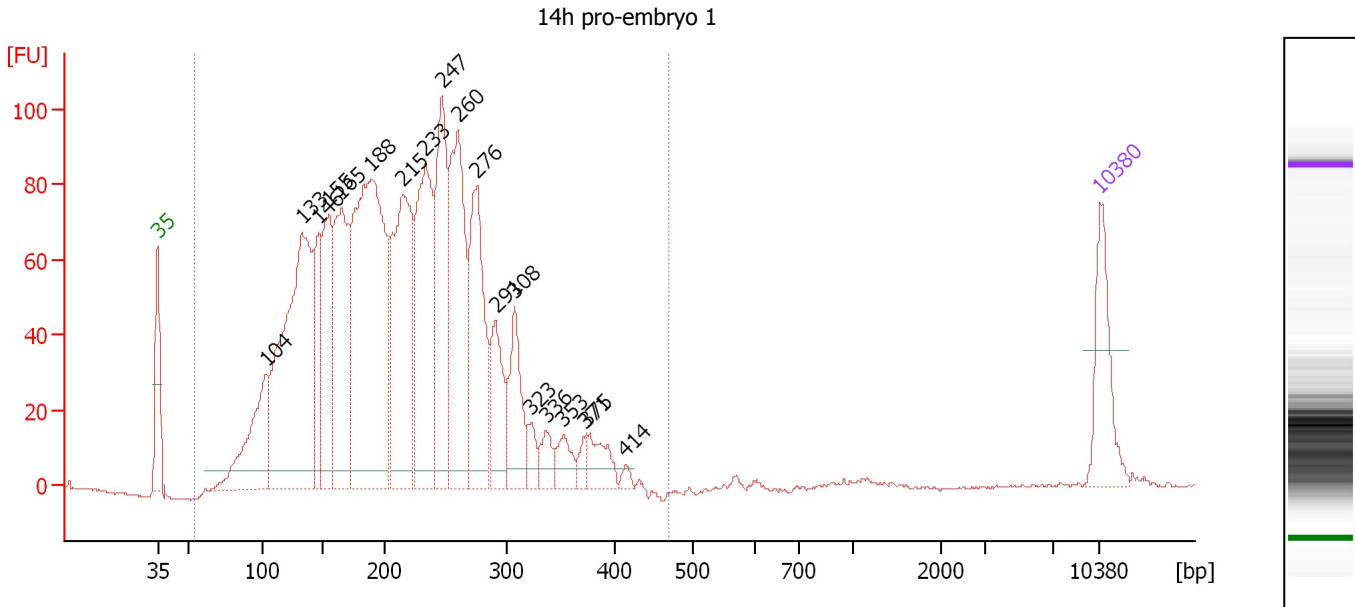
Peak table for Ladder

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	45.29	50	150.00	4,545.5	Ladder Peak
3	50.72	100	150.00	2,272.7	Ladder Peak
4	55.31	150	150.00	1,515.2	Ladder Peak
5	59.89	200	150.00	1,136.4	Ladder Peak
6	68.89	300	150.00	757.6	Ladder Peak
7	76.96	400	150.00	568.2	Ladder Peak
8	82.70	500	150.00	454.5	Ladder Peak
9	87.33	600	150.00	378.8	Ladder Peak
10	90.59	700	150.00	324.7	Ladder Peak
11	94.61	1,000	150.00	227.3	Ladder Peak
12	101.22	2,000	150.00	113.6	Ladder Peak
13	104.49	3,000	150.00	75.8	Ladder Peak
14	109.52	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\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : 14h pro-embryo 1

Number of peaks found: 19 Corr. Area 1: 2,340.0
 Noise: 0.3

Peak table for sample 1 : 14h pro-embryo 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	51.05	104	187.06	2,736.1	
3	53.70	133	588.85	6,733.3	
4	54.98	146	127.83	1,322.2	
5	55.74	155	203.65	1,994.9	
6	56.66	165	280.83	2,581.9	
7	58.83	188	643.56	5,175.2	
8	61.22	215	345.78	2,439.9	
9	62.90	233	307.72	1,997.6	
10	64.14	247	241.30	1,479.2	
11	65.33	260	290.11	1,687.8	
12	66.75	276	226.07	1,240.4	
13	68.12	291	102.67	533.9	
14	69.53	308	103.59	509.7	
15	70.77	323	24.77	116.1	
16	71.83	336	29.31	132.0	
17	73.16	353	34.75	149.2	
18	74.66	371	14.91	60.8	
19	74.93	375	43.33	175.2	
20	77.76	414	7.12	26.1	
21	113.00	10,380	75.00	10.9	Upper Marker

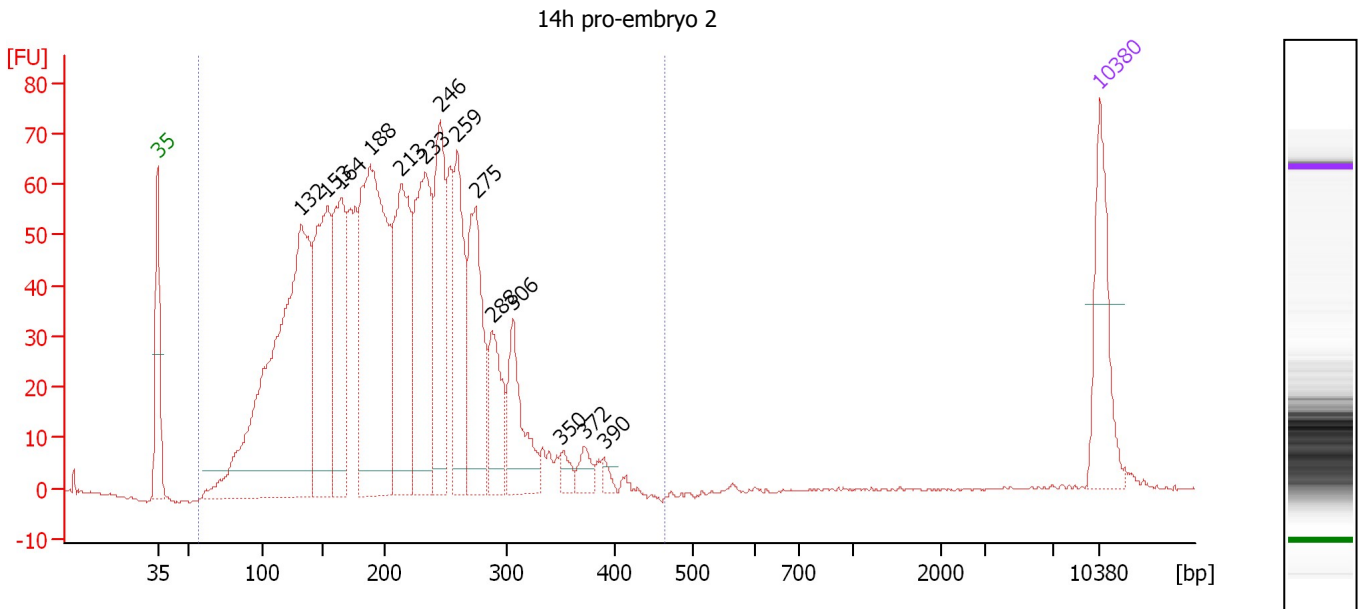
Region table for sample 1 : 14h pro-embryo 1

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
54	471	214	31,097.9	3,769.82	2,340.0	100	31.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : 14h pro-embryo 2

Number of peaks found: 14 Corr. Area 1: 1,748.7
 Noise: 0.2

Peak table for sample 2 : 14h pro-embryo 2

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	53.65	132	664.59	7,628.9	
3	55.59	153	260.74	2,580.3	
4	56.60	164	197.65	1,824.5	
5	58.81	188	449.30	3,618.4	
6	61.05	213	235.06	1,673.2	
7	62.90	233	246.09	1,597.5	
8	64.00	246	174.93	1,079.0	
9	65.23	259	147.00	858.8	
10	66.64	275	165.32	910.9	
11	67.83	288	83.16	437.2	
12	69.42	306	95.36	471.4	
13	72.94	350	14.67	63.5	
14	74.70	372	18.90	77.0	
15	76.15	390	8.06	31.3	
16	113.00	10,380	75.00	10.9	Upper Marker

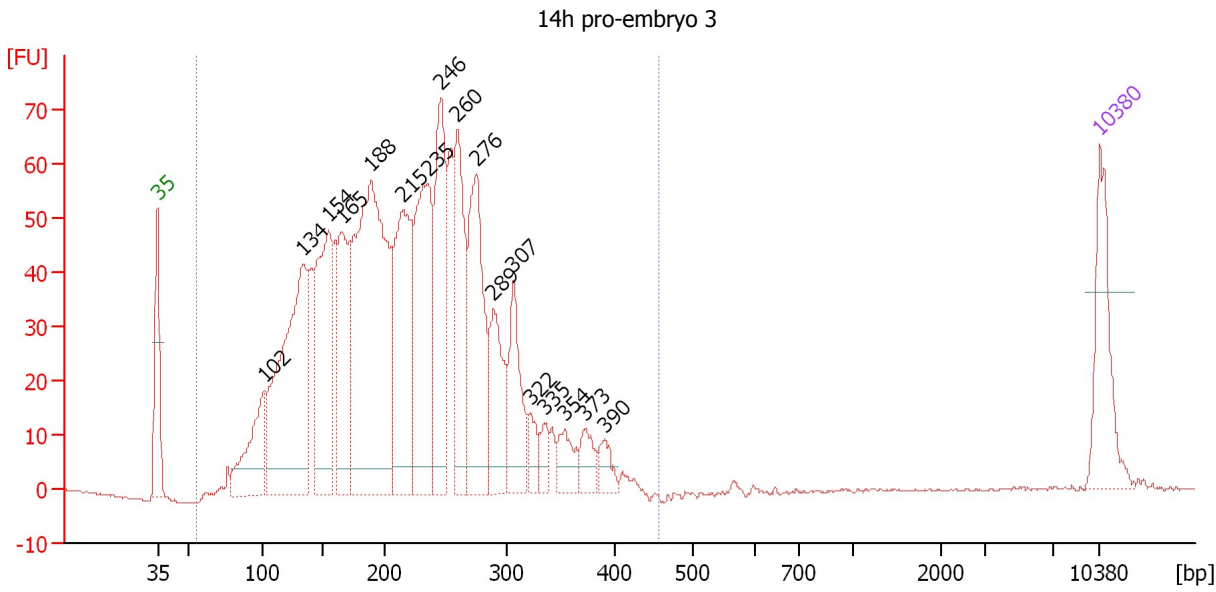
Region table for sample 2 : 14h pro-embryo 2

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
56	465	208	25,071.4	2,957.85	1,748.7	100	30.7

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : 14h pro-embryo 3

Number of peaks found: 17 Corr. Area 1: 1,590.4
 Noise: 0.2

Peak table for sample 3 : 14h pro-embryo 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	50.92	102	121.76	1,805.9	
3	53.82	134	387.08	4,381.8	
4	55.70	154	235.26	2,309.7	
5	56.69	165	164.92	1,514.0	
6	58.83	188	542.54	4,362.0	
7	61.20	215	241.57	1,705.8	
8	63.08	235	243.16	1,564.8	
9	64.07	246	195.32	1,201.1	
10	65.27	260	162.20	946.0	
11	66.71	276	201.60	1,107.9	
12	67.91	289	99.89	523.5	
13	69.43	307	95.13	469.9	
14	70.64	322	27.18	128.0	
15	71.72	335	21.41	96.8	
16	73.24	354	34.77	148.9	
17	74.76	373	28.71	116.7	
18	76.14	390	22.07	85.8	
19	113.00	10,380	75.00	10.9	Upper Marker

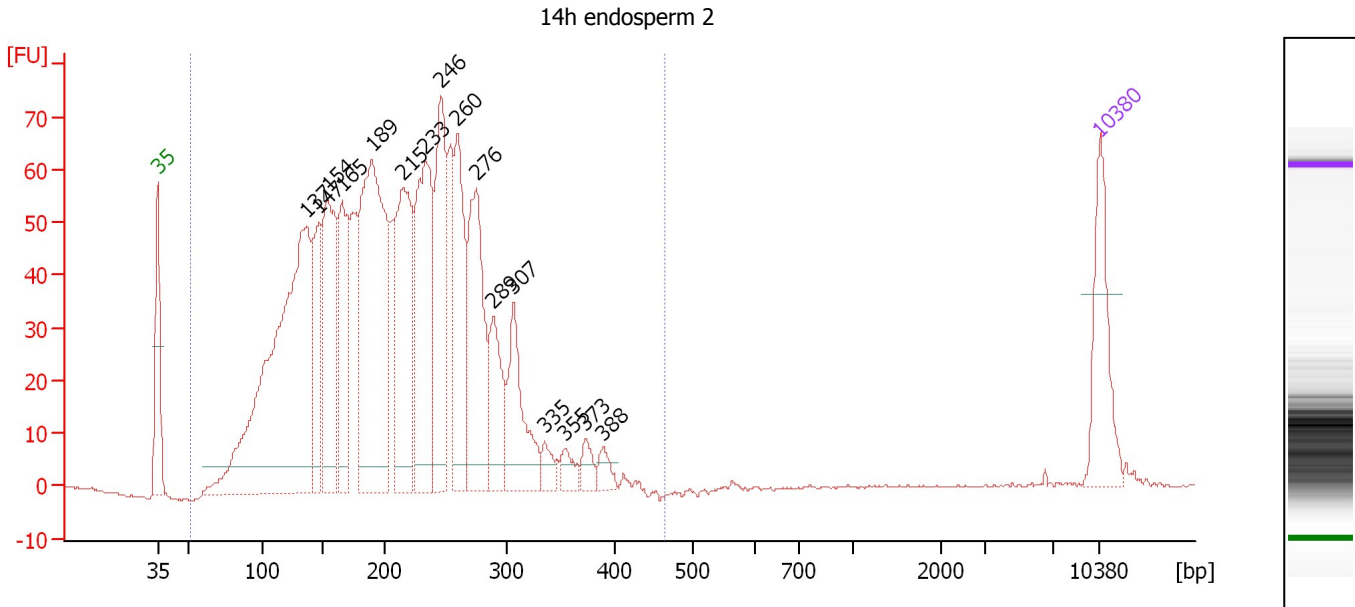
Region table for sample 3 : 14h pro-embryo 3

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
55	457	219	24,556.3	3,036.97	1,590.4	100	30.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : 14h endosperm 2

Number of peaks found: 16 Corr. Area 1: 1,694.6
 Noise: 0.1

Peak table for sample 4 : 14h endosperm 2

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	54.08	137	679.83	7,536.5	
3	54.99	147	130.04	1,344.2	
4	55.63	154	227.10	2,241.5	
5	56.72	165	139.51	1,278.0	
6	58.90	189	437.93	3,507.8	
7	61.26	215	245.57	1,729.0	
8	62.85	233	251.14	1,634.0	
9	64.03	246	198.02	1,219.7	
10	65.26	260	161.61	943.1	
11	66.71	276	194.38	1,067.9	
12	67.94	289	88.98	465.9	
13	69.44	307	112.75	556.9	
14	71.75	335	20.15	91.0	
15	73.30	355	19.33	82.6	
16	74.80	373	18.43	74.8	
17	76.02	388	16.26	63.4	
18	113.00	10,380	75.00	10.9	Upper Marker

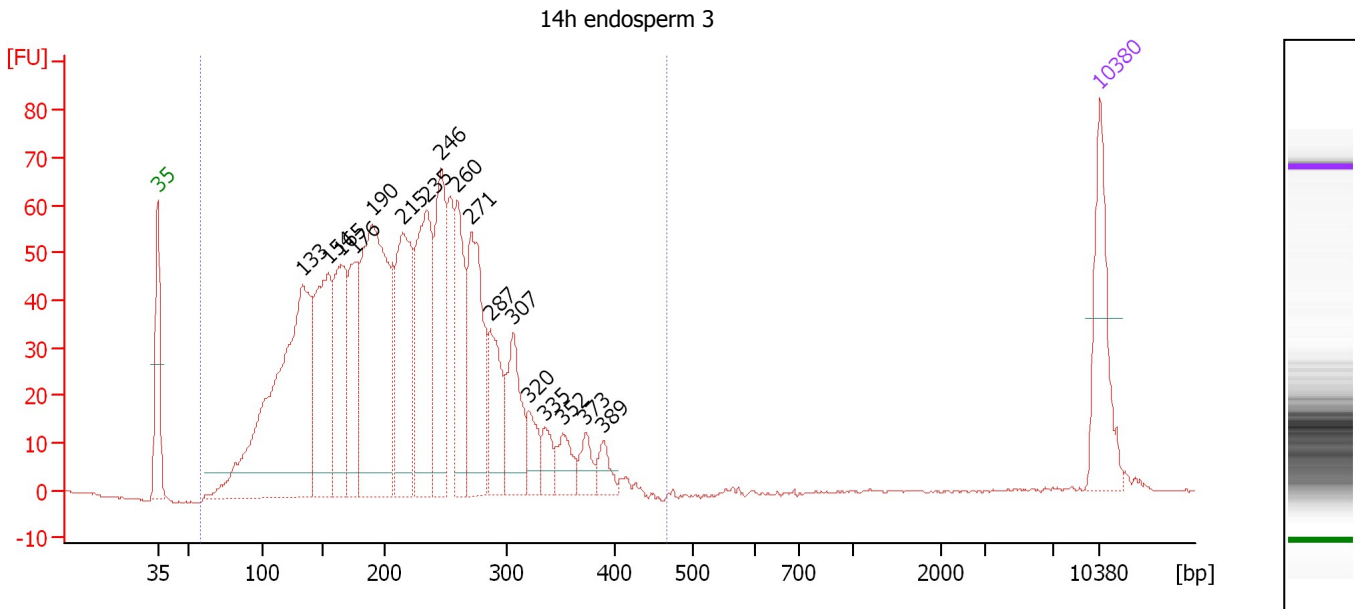
Region table for sample 4 : 14h endosperm 2

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
51	466	210	26,730.4	3,173.28	1,694.6	100	30.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : 14h endosperm 3

Number of peaks found: 17 Corr. Area 1: 1,571.4
 Noise: 0.2

Peak table for sample 5 : 14h endosperm 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	53.79	133	511.70	5,808.5	
3	55.68	154	209.00	2,055.2	
4	56.65	165	158.84	1,461.8	
5	57.66	176	129.03	1,112.7	
6	58.96	190	384.00	3,065.9	
7	61.21	215	202.85	1,431.6	
8	63.01	235	205.58	1,327.3	
9	64.07	246	157.27	966.9	
10	65.27	260	123.05	717.7	
11	66.29	271	167.27	935.0	
12	67.76	287	85.99	453.2	
13	69.42	307	86.23	426.2	
14	70.53	320	34.15	161.6	
15	71.73	335	27.58	124.7	
16	73.11	352	31.24	134.4	
17	74.77	373	24.12	98.0	
18	76.11	389	19.96	77.6	
19	113.00	10,380	75.00	10.9	Upper Marker

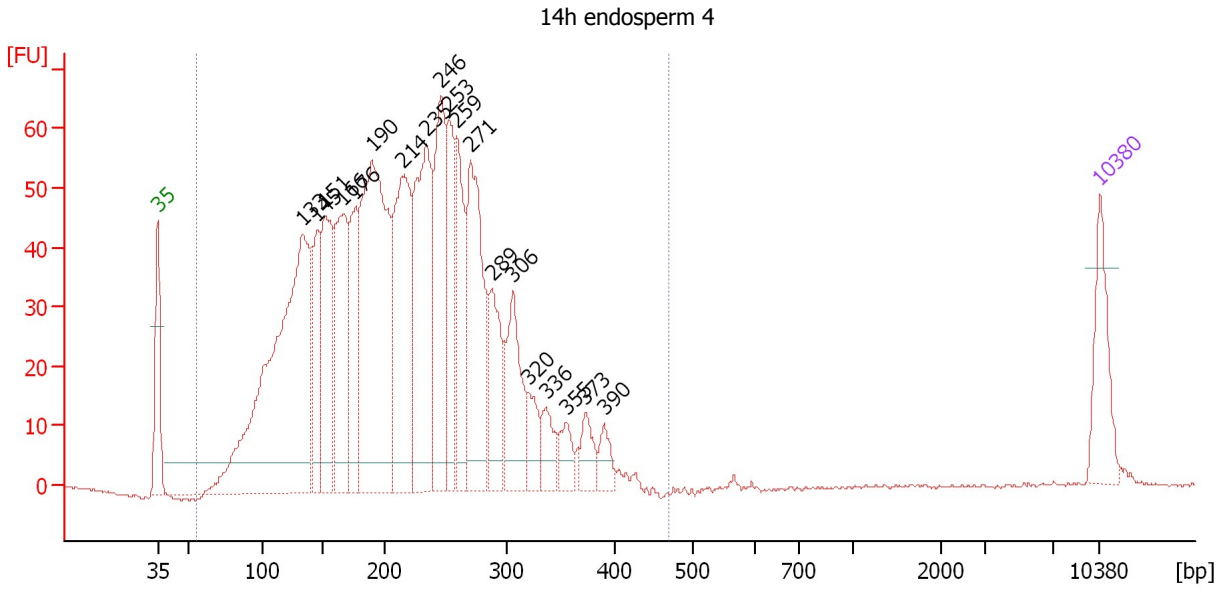
Region table for sample 5 : 14h endosperm 3

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
58	466	218	20,865.0	2,568.16	1,571.4	100	30.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : 14h endosperm 4

Number of peaks found: 19 Corr. Area 1: 1,550.8
 Noise: 0.2

Peak table for sample 6 : 14h endosperm 4

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	53.70	133	858.39	9,815.6	
3	54.87	145	147.11	1,534.7	
4	55.43	151	235.23	2,354.9	
5	56.79	166	259.84	2,370.0	
6	57.67	176	167.44	1,443.2	
7	58.93	190	610.93	4,883.4	
8	61.18	214	342.67	2,423.1	
9	63.00	235	347.96	2,248.0	
10	64.07	246	262.63	1,614.5	
11	64.68	253	163.77	979.9	
12	65.20	259	161.98	947.8	
13	66.27	271	289.58	1,619.7	
14	67.86	289	127.89	671.6	
15	69.40	306	147.64	730.3	
16	70.52	320	51.50	243.7	
17	71.83	336	50.37	226.8	
18	73.33	355	38.60	164.8	
19	74.78	373	39.38	160.0	
20	76.18	390	30.53	118.5	
21	113.00	10,380	75.00	10.9	Upper Marker

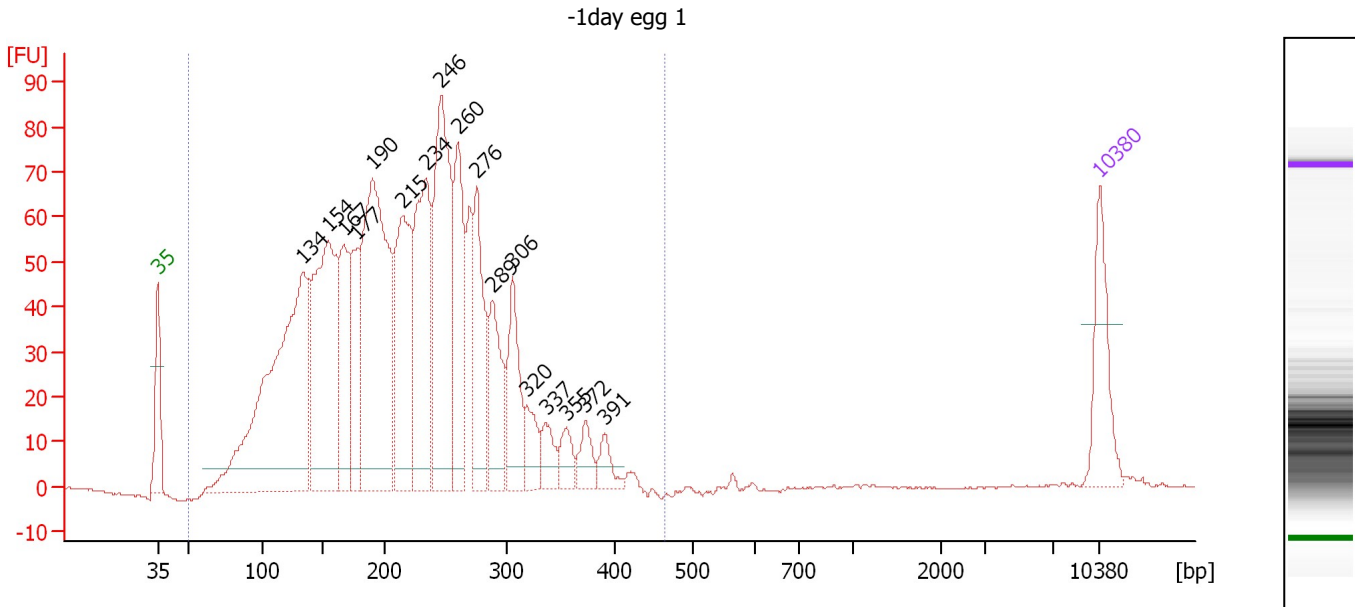
Region table for sample 6 : 14h endosperm 4

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
56	470	217	35,264.4	4,298.47	1,550.8	100	31.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : -1day egg 1

Number of peaks found: 17 Corr. Area 1: 1,835.4
 Noise: 0.2

Peak table for sample 7 : -1day egg 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	53.79	134	681.81	7,736.6	
3	55.65	154	428.77	4,225.7	
4	56.84	167	161.42	1,467.1	
5	57.79	177	128.46	1,099.2	
6	58.98	190	514.00	4,098.3	
7	61.26	215	248.64	1,750.6	
8	62.93	234	280.76	1,820.3	
9	64.07	246	344.11	2,116.2	
10	65.30	260	187.49	1,092.1	
11	66.73	276	161.63	887.4	
12	67.87	289	121.36	637.0	
13	69.39	306	115.26	570.3	
14	70.49	320	44.43	210.5	
15	71.87	337	37.52	168.8	
16	73.34	355	32.06	136.8	
17	74.72	372	34.06	138.6	
18	76.24	391	27.43	106.3	
19	113.00	10,380	75.00	10.9	Upper Marker

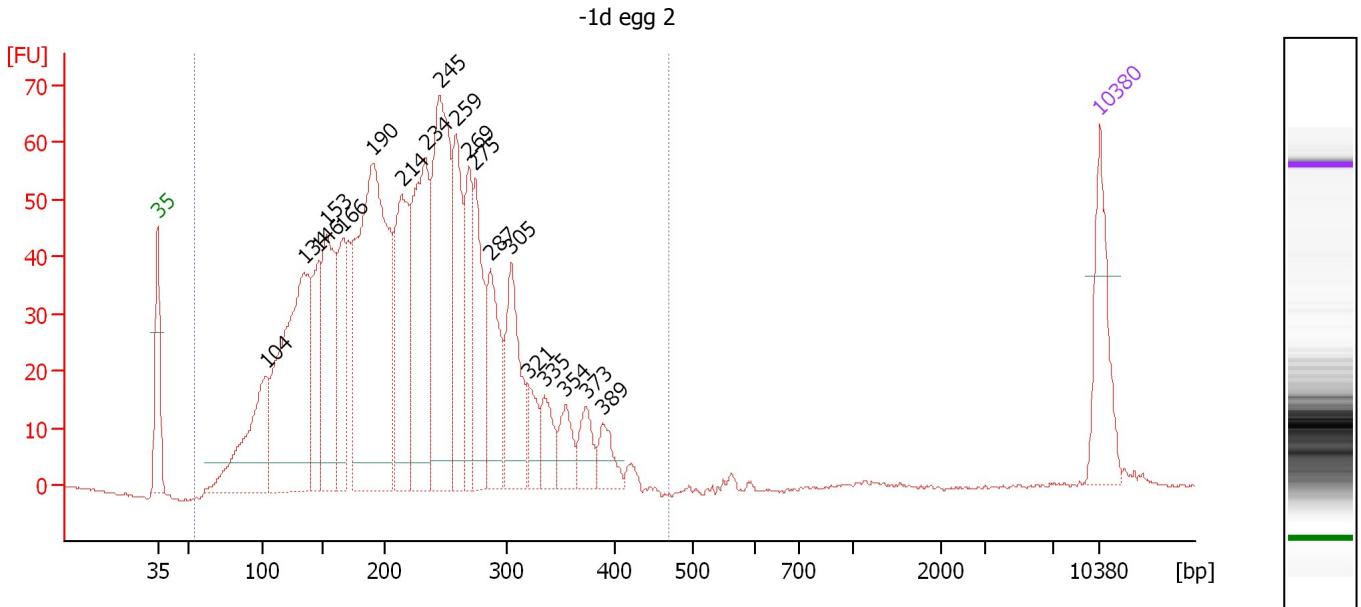
Region table for sample 7 : -1day egg 1

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
51	466	217	29,759.9	3,631.10	1,835.4	100	31.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : -1d egg 2

Number of peaks found: 19 Corr. Area 1: 1,519.3
 Noise: 0.1

Peak table for sample 8 : -1d egg 2

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	51.10	104	196.33	2,856.3	
3	53.85	134	413.91	4,676.8	
4	54.96	146	104.96	1,087.9	
5	55.58	153	216.38	2,142.6	
6	56.74	166	126.15	1,154.0	
7	58.96	190	546.71	4,364.4	
8	61.13	214	207.80	1,473.3	
9	62.91	234	275.43	1,787.0	
10	63.97	245	320.71	1,980.8	
11	65.18	259	153.65	899.9	
12	66.14	269	109.51	615.8	
13	66.62	275	129.17	712.2	
14	67.73	287	123.16	649.9	
15	69.27	305	130.24	647.6	
16	70.58	321	37.76	178.3	
17	71.73	335	44.74	202.3	
18	73.28	354	39.80	170.2	
19	74.82	373	38.11	154.6	
20	76.07	389	31.53	122.8	
21	113.00	10,380	75.00	10.9	Upper Marker

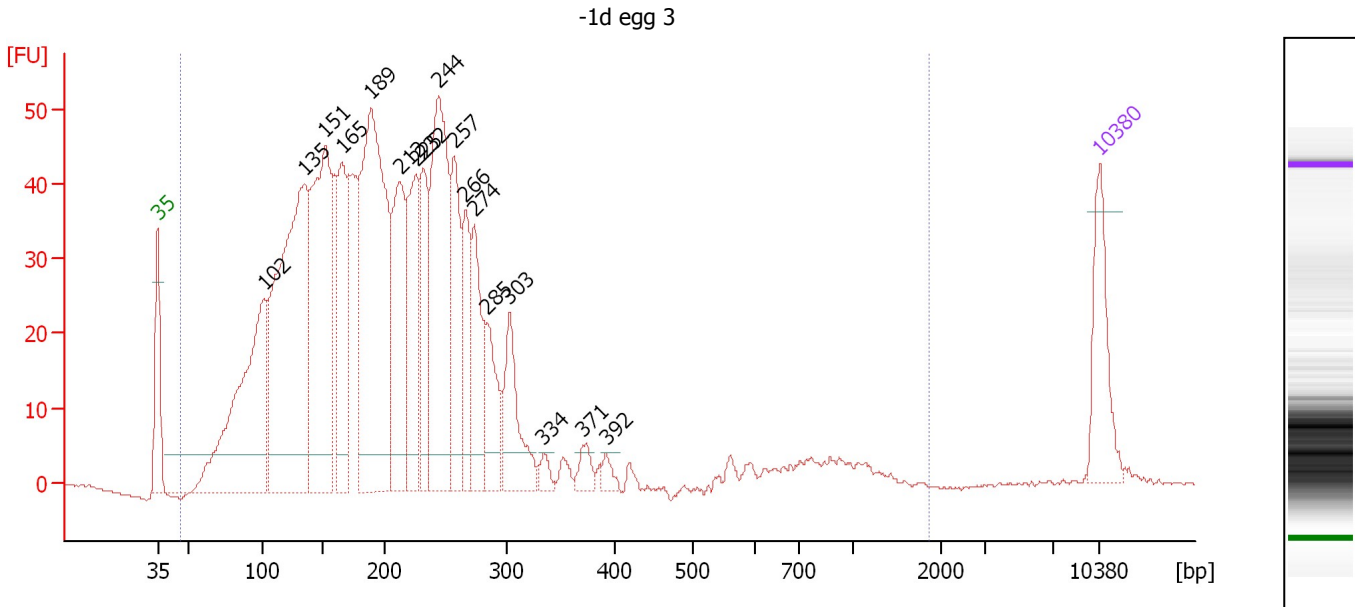
Region table for sample 8 : -1d egg 2

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
54	471	222	26,247.3	3,263.04	1,519.3	100	31.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : -1d egg 3

Number of peaks found: 17 Corr. Area 1: 1,287.8
 Noise: 0.2

Peak table for sample 9 : -1d egg 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	50.91	102	426.89	6,334.4	
3	53.92	135	627.92	7,051.7	
4	55.43	151	450.20	4,507.8	
5	56.69	165	185.81	1,705.4	
6	58.87	189	543.86	4,362.6	
7	61.01	212	212.82	1,518.1	
8	62.17	225	166.27	1,117.9	
9	62.76	232	130.56	853.3	
10	63.87	244	335.61	2,082.0	
11	65.04	257	153.71	905.6	
12	65.86	266	95.27	541.9	
13	66.54	274	119.22	659.5	
14	67.51	285	88.64	471.7	
15	69.12	303	110.24	551.6	
16	71.64	334	14.78	67.0	
17	74.60	371	21.12	86.3	
18	76.30	392	12.59	48.7	
19	113.00	10,380	75.00	10.9	Upper Marker

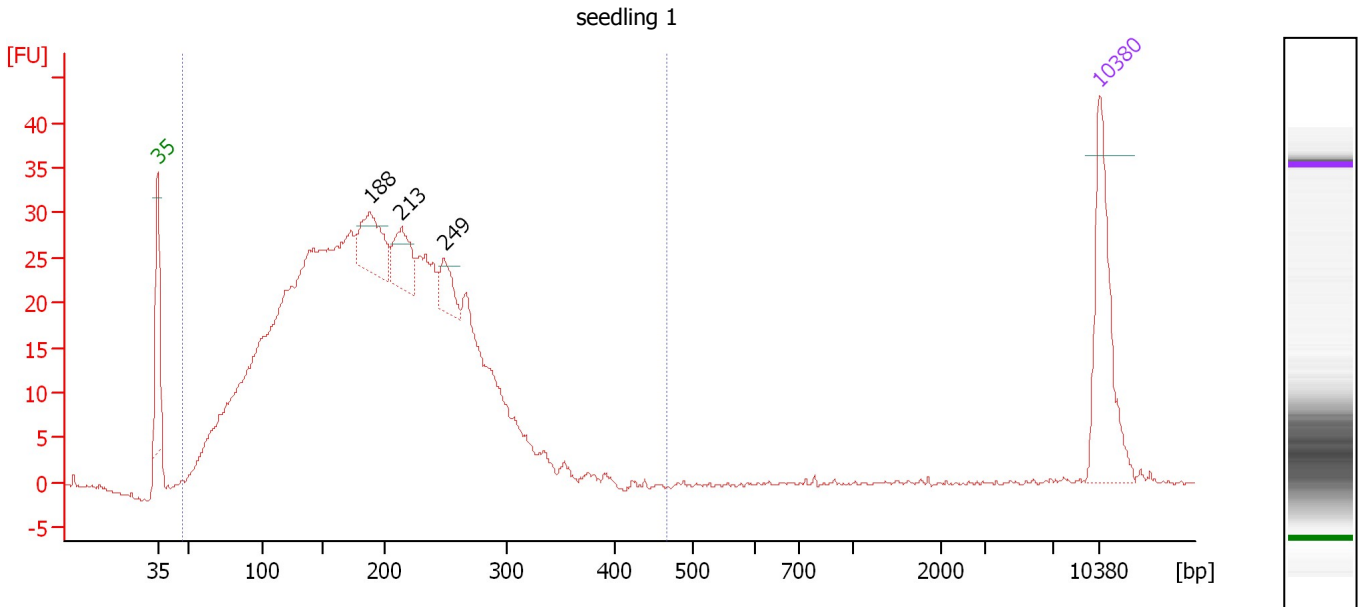
Region table for sample 9 : -1d egg 3

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
47	1,858	217	35,536.8	3,920.62	1,287.8	100	62.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : seedling 1

Number of peaks found: 3 Corr. Area 1: 804.3
 Noise: 0.2

Peak table for sample 10 : seedling 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	58.80	188	60.39	486.6	
3	61.10	213	42.91	304.7	
4	64.29	249	25.79	157.0	
5	113.00	10,380	75.00	10.9	Upper Marker

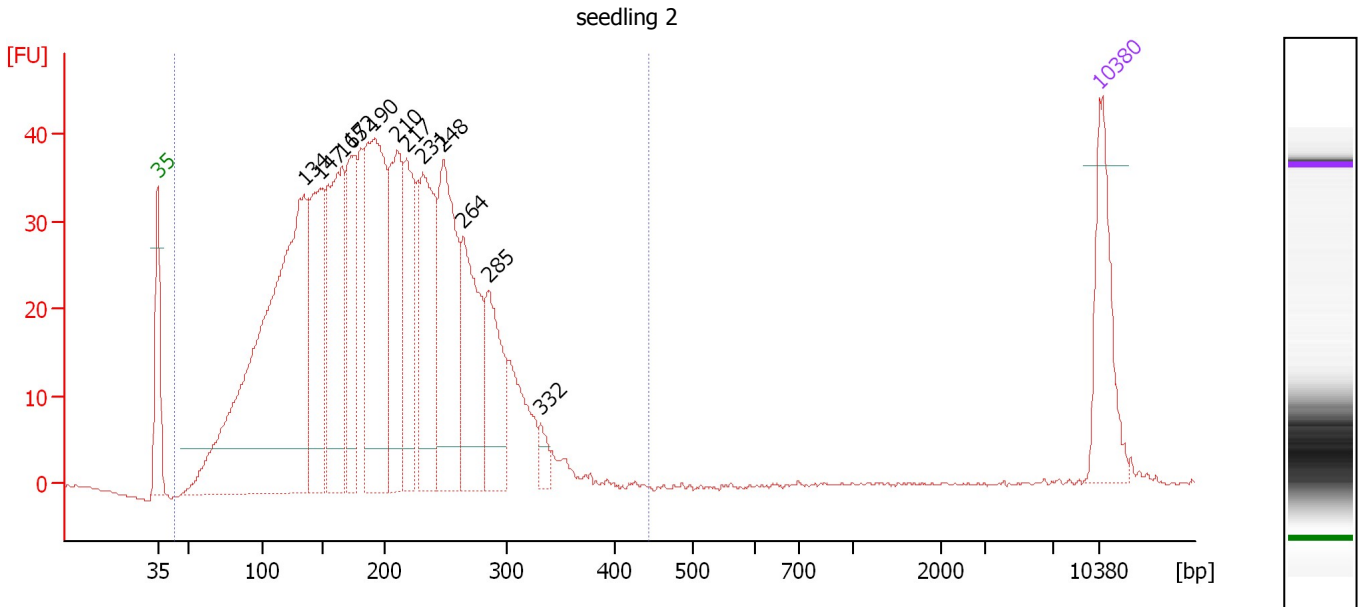
Region table for sample 10 : seedling 1

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
47	467	190	22,124.6	2,276.22	804.3	99	34.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : seedling 2

Number of peaks found: 12 Corr. Area 1: 1,051.3
 Noise: 0.1

Peak table for sample 11 : seedling 2

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	53.85	134	729.44	8,240.5	
3	55.04	147	185.52	1,911.1	
4	56.67	165	241.39	2,218.0	
5	57.37	172	129.02	1,133.5	
6	59.00	190	292.77	2,331.3	
7	60.78	210	168.04	1,212.9	
8	61.43	217	130.97	914.2	
9	62.67	231	180.17	1,182.7	
10	64.20	248	214.19	1,309.2	
11	65.69	264	158.76	909.7	
12	67.52	285	109.92	584.8	
13	71.44	332	17.20	78.6	
14	113.00	10,380	75.00	10.9	Upper Marker

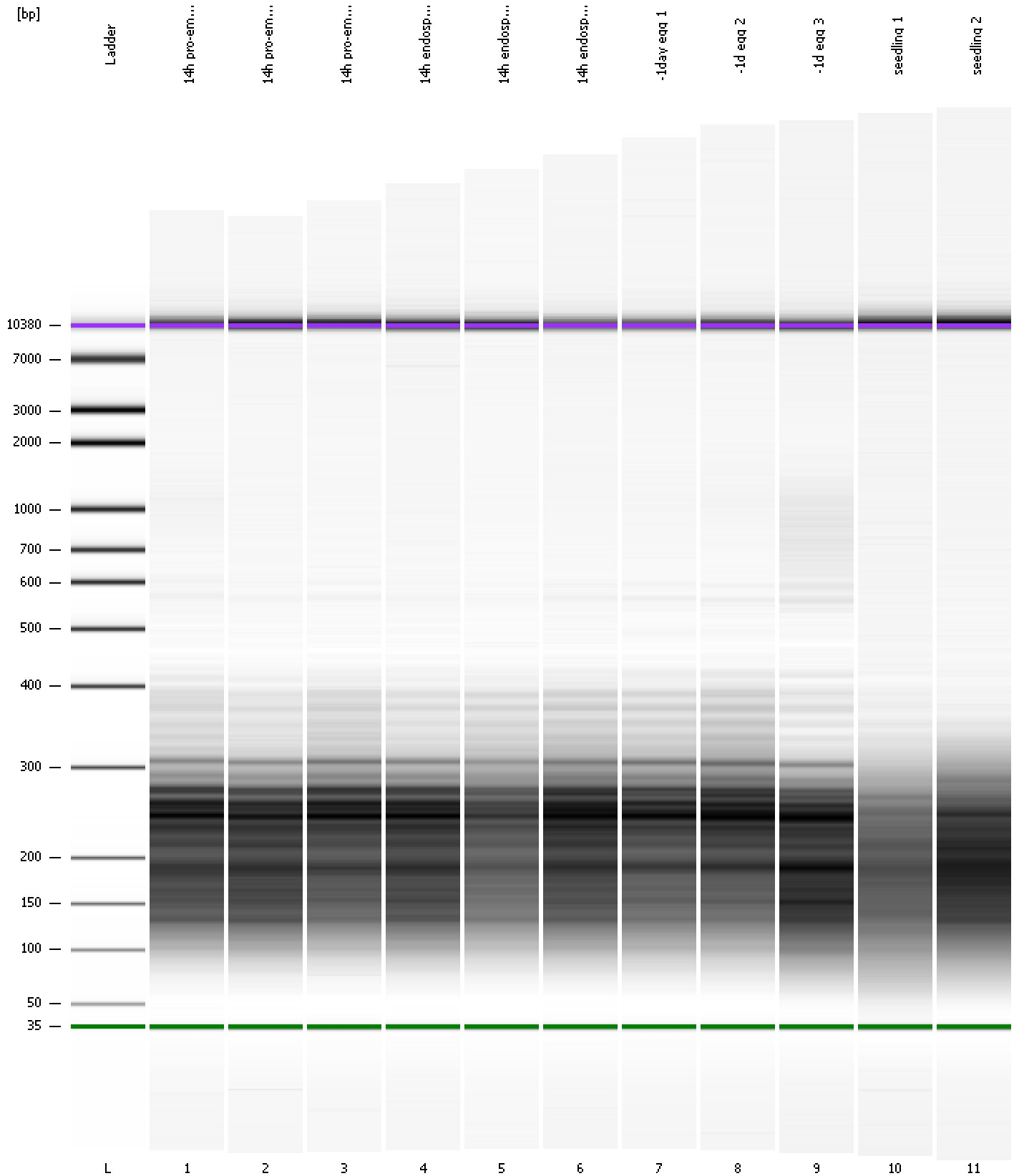
Region table for sample 11 : seedling 2

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Co Corr. lor Area	% of Total	Size distribution in CV [%]
43	443	197	25,375.7	2,764.31	1,051.3	100	32.4

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
Modified: 4/17/2014 5:09:53 PM

Gel Image

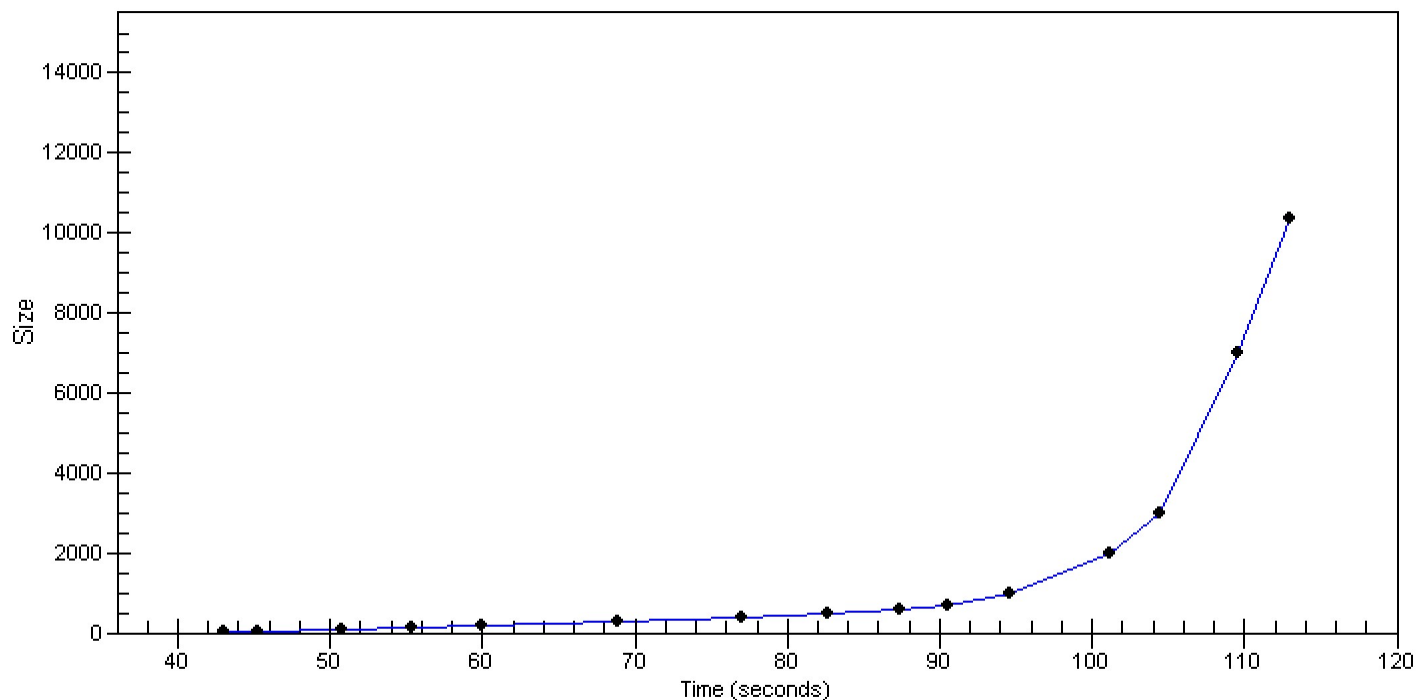


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
Modified: 4/17/2014 5:09:53 PM

Curves

Standard Curve



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad

Created: 4/17/2014 4:27:43 PM
 Modified: 4/17/2014 5:09:53 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		4/17/2014 5:09:03 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2014-04-17\2014-04-17_004.xad)		Instrument	Run		4/17/2014 4:27:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		4/17/2014 4:27:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		4/17/2014 4:27:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		4/17/2014 4:27:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		4/17/2014 4:27:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		4/17/2014 4:27:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		4/17/2014 4:27:49 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1