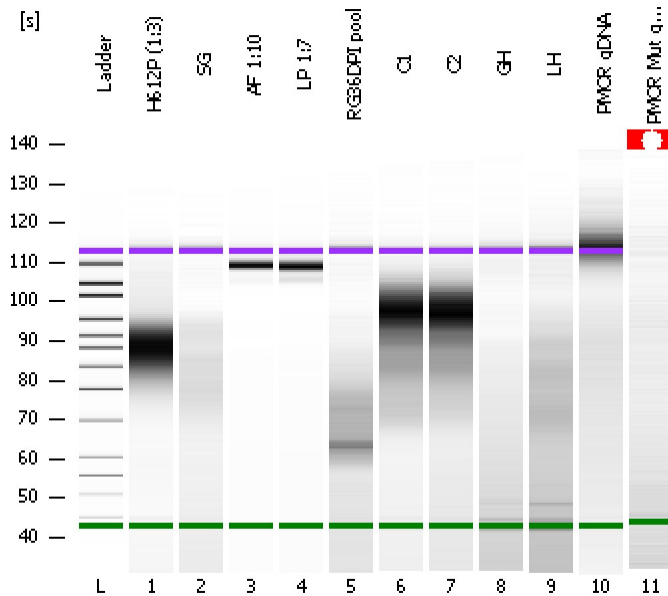


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
Modified: 8/11/2016 5:48:04 PM

Electrophoresis File Run Summary



Instrument Information:

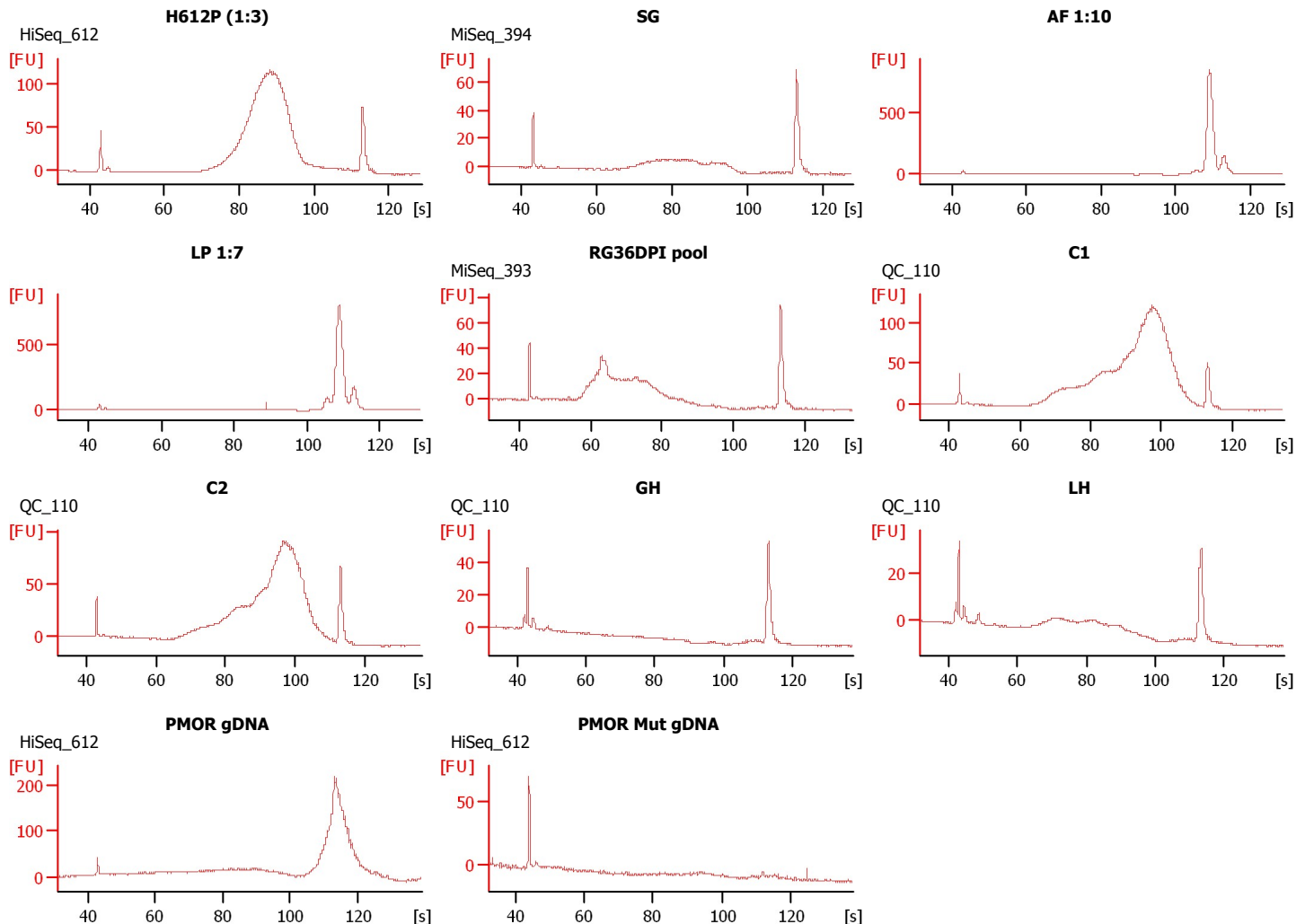
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
Modified: 8/11/2016 5:48:04 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
H612P (1:3)	HiSeq_612	<input type="checkbox"/>	✓			
SG	MiSeq_394	<input type="checkbox"/>	✓			
AF 1:10		<input type="checkbox"/>	✓			
LP 1:7		<input type="checkbox"/>	✓			
RG36DPI pool	MiSeq_393	<input type="checkbox"/>	✓			
C1	QC_110	<input type="checkbox"/>	✓			
C2	QC_110	<input type="checkbox"/>	✓			
GH	QC_110	<input type="checkbox"/>	✓			
LH	QC_110	<input type="checkbox"/>	✓			
PMOR gDNA	HiSeq_612	<input type="checkbox"/>	✓			
PMOR Mut gDNA	HiSeq_612	<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
Modified: 8/11/2016 5:48:04 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

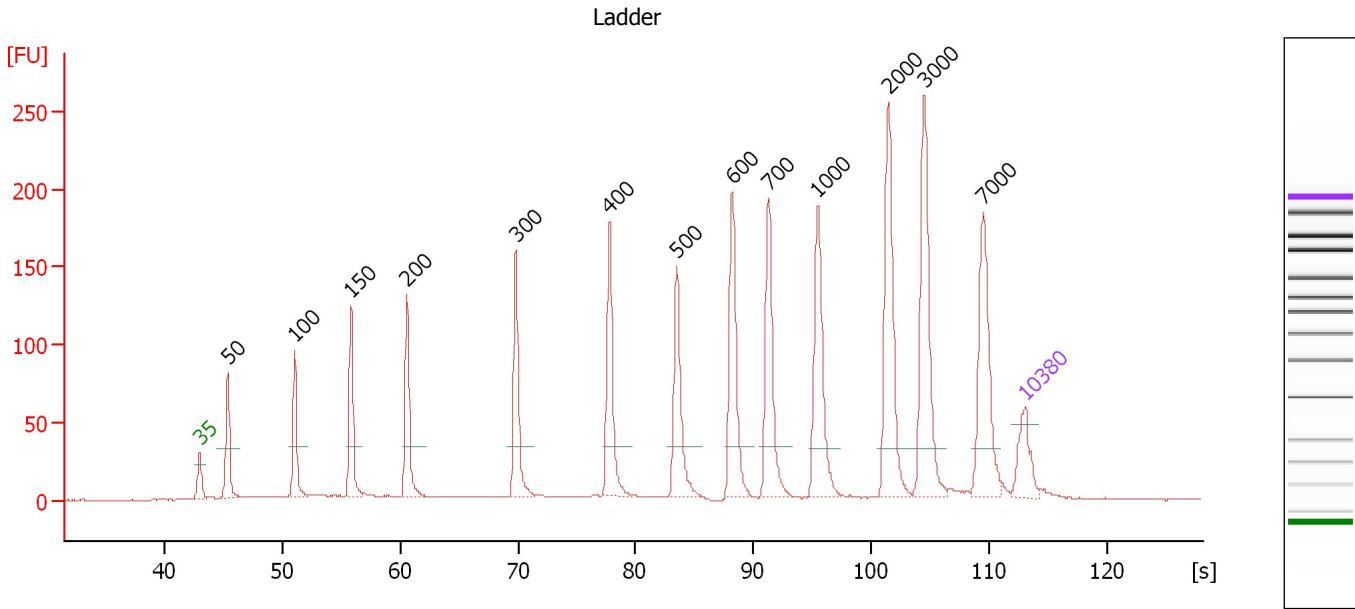
Ladder

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:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

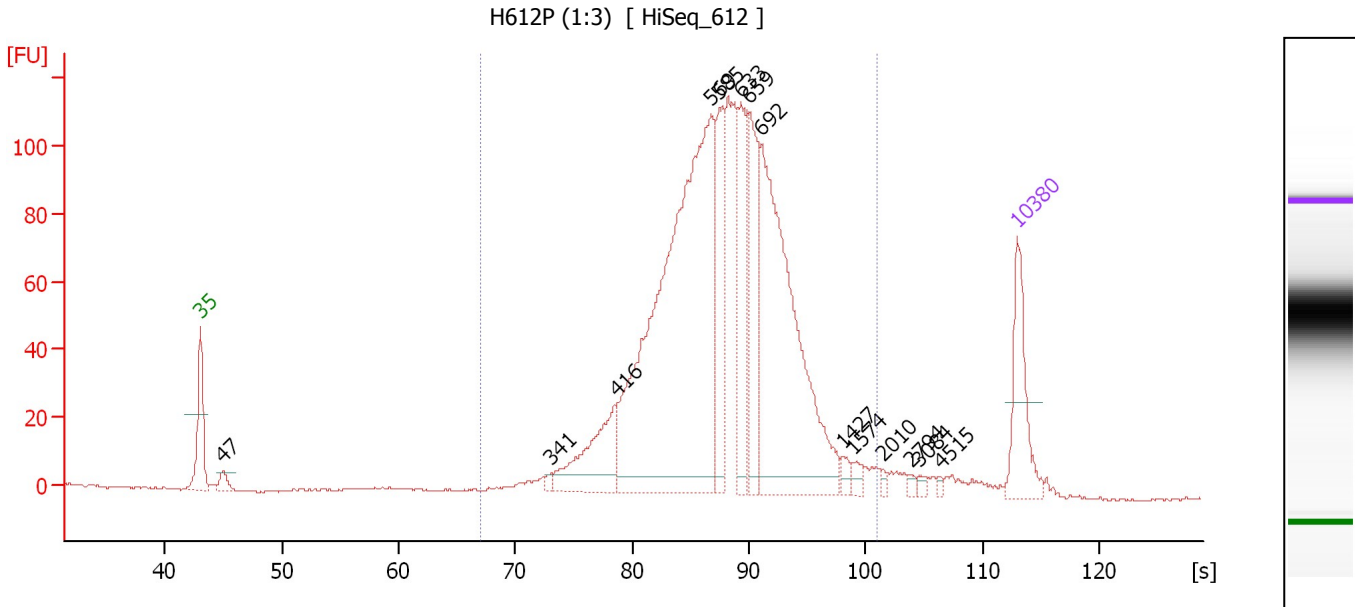
Peak table for Ladder

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	45.37
3	100	150.00	2,272.7	Ladder Peak	51.07
4	150	150.00	1,515.2	Ladder Peak	55.85
5	200	150.00	1,136.4	Ladder Peak	60.59
6	300	150.00	757.6	Ladder Peak	69.84
7	400	150.00	568.2	Ladder Peak	77.77
8	500	150.00	454.5	Ladder Peak	83.47
9	600	150.00	378.8	Ladder Peak	88.21
10	700	150.00	324.7	Ladder Peak	91.31
11	1,000	150.00	227.3	Ladder Peak	95.45
12	2,000	150.00	113.6	Ladder Peak	101.42
13	3,000	150.00	75.8	Ladder Peak	104.48
14	7,000	150.00	32.5	Ladder Peak	109.49
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : H612P (1:3)

Number of peaks found: 14 Corr. Area 1: 1,718.0
 Noise: 0.3

Peak table for sample 1 : H612P (1:3)

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	47	18.57	595.9		44.93
3	341	5.82	25.9		73.08
4	416	121.15	441.1		78.69
5	569	786.41	2,094.1		86.74
6	585	124.67	323.2		87.47
7	633	123.35	295.4		89.22
8	659	143.39	329.5		90.05
9	692	459.87	1,006.8		91.06
10	1,427	10.15	10.8		98.01
11	1,574	9.99	9.6		98.88
12	2,010	4.17	3.1		101.46
13	2,794	3.89	2.1		103.85
14	3,084	4.23	2.1		104.58
15	4,515	3.08	1.0		106.38
16	10,380	75.00	10.9	Upper Marker	113.00

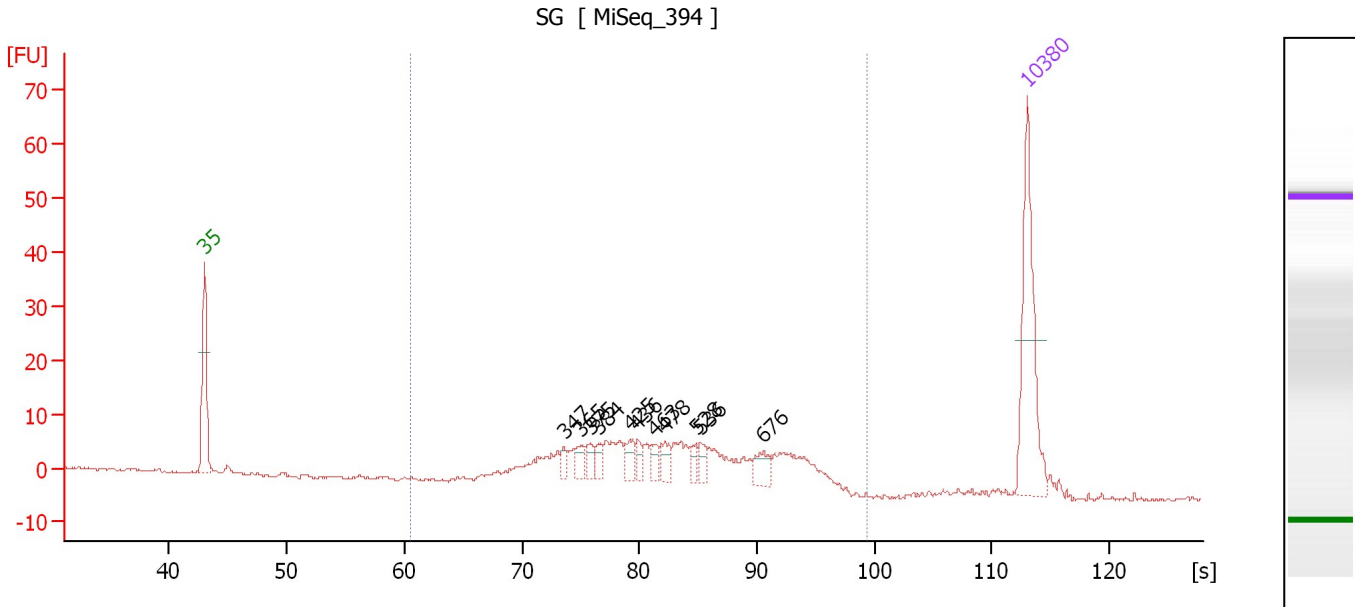
Region table for sample 1 : H612P (1:3)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
269	1,911	639	1,718.0	5,287.2	2,010.85	96	33.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : SG

Number of peaks found: 11 Corr. Area 1: 201.3
 Noise: 0.3

Peak table for sample 2 : SG

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	347	7.59	33.1		73.57
3	365	12.37	51.3		75.03
4	375	8.67	35.0		75.81
5	384	8.69	34.2		76.54
6	425	14.16	50.5		79.19
7	436	8.28	28.8		79.83
8	463	9.55	31.3		81.34
9	478	11.49	36.4		82.20
10	528	7.42	21.3		84.81
11	536	8.67	24.5		85.17
12	676	14.14	31.7		90.57
13	10,380	75.00	10.9	Upper Marker	113.00

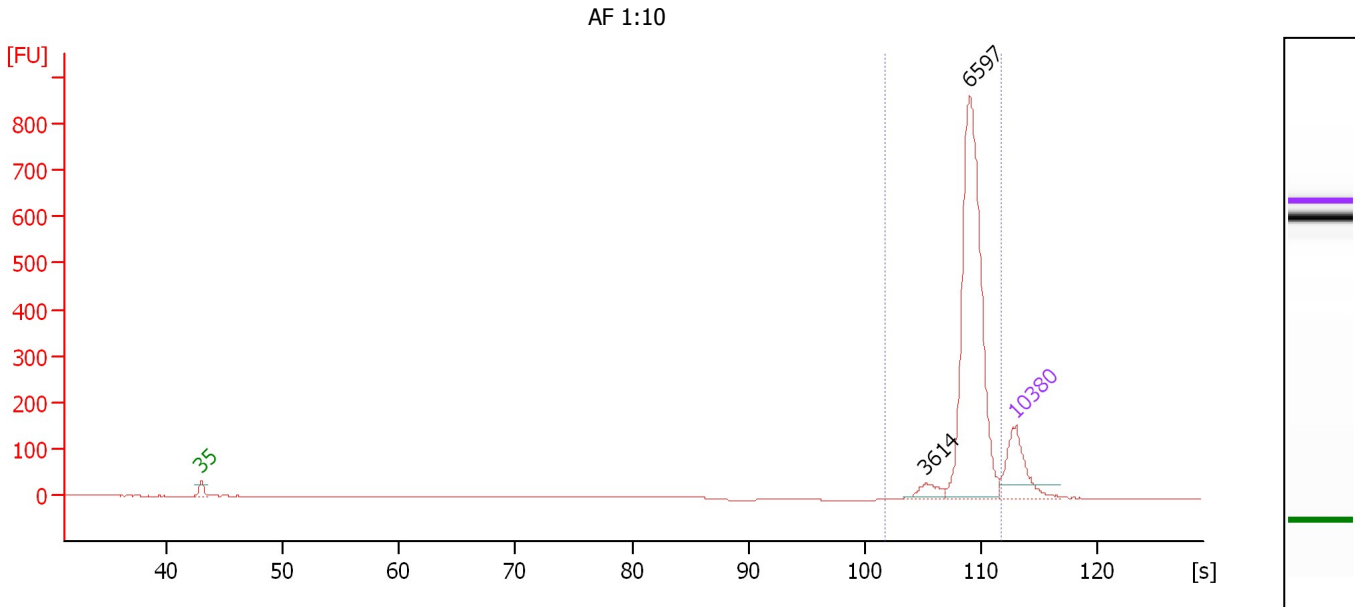
Region table for sample 2 : SG

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,658	524	201.3	1,020.7	306.52	97	35.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : AF 1:10

Number of peaks found: 2 Corr. Area 1: 1,572.2
 Noise: 0.2

Peak table for sample 3 : AF 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	3,614	19.81	8.3		105.25
3	6,597	462.69	106.3		108.99
4	10,380	75.00	10.9	Upper Marker	113.00

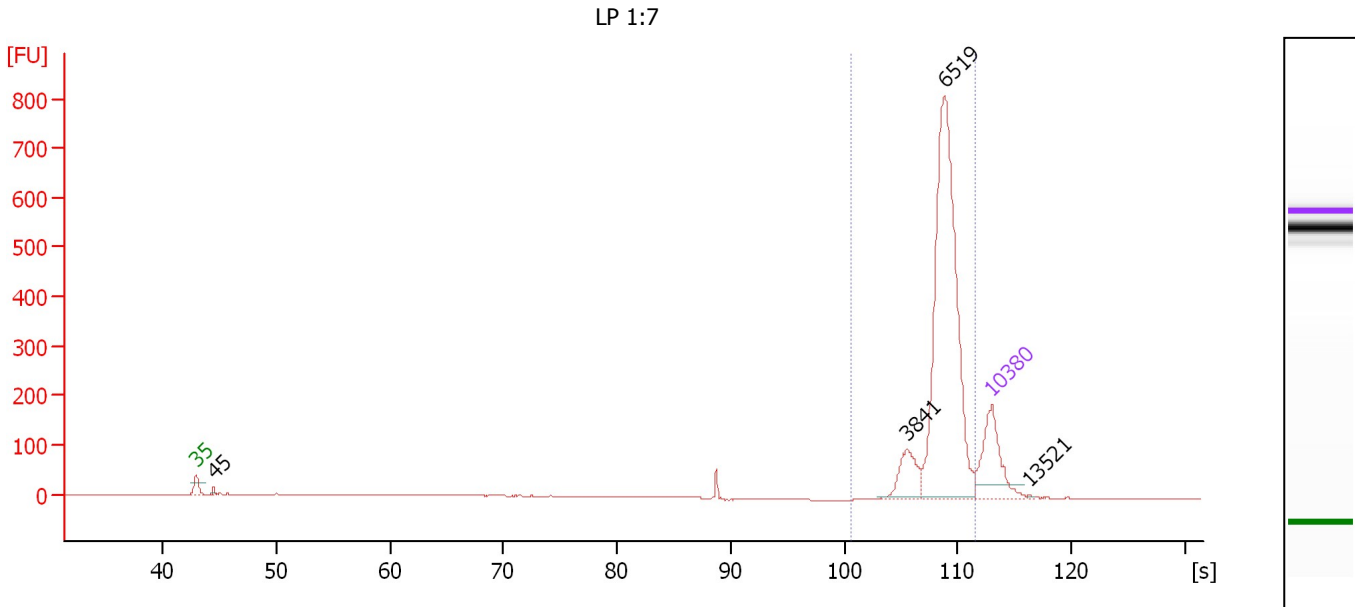
Region table for sample 3 : AF 1:10

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
2,106	9,178	6,745	1,572.2	107.8	475.18	99	12.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : LP 1:7

Number of peaks found: 4 Corr. Area 1: 1,788.0
 Noise: 0.4

Peak table for sample 4 : LP 1:7

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	3.87	131.4		44.51
3	3,841	45.58	18.0		105.53
4	6,519	438.83	102.0		108.89
5	10,380	75.00	10.9	Upper Marker	113.00
6	13,521	0.00	0.0		116.26

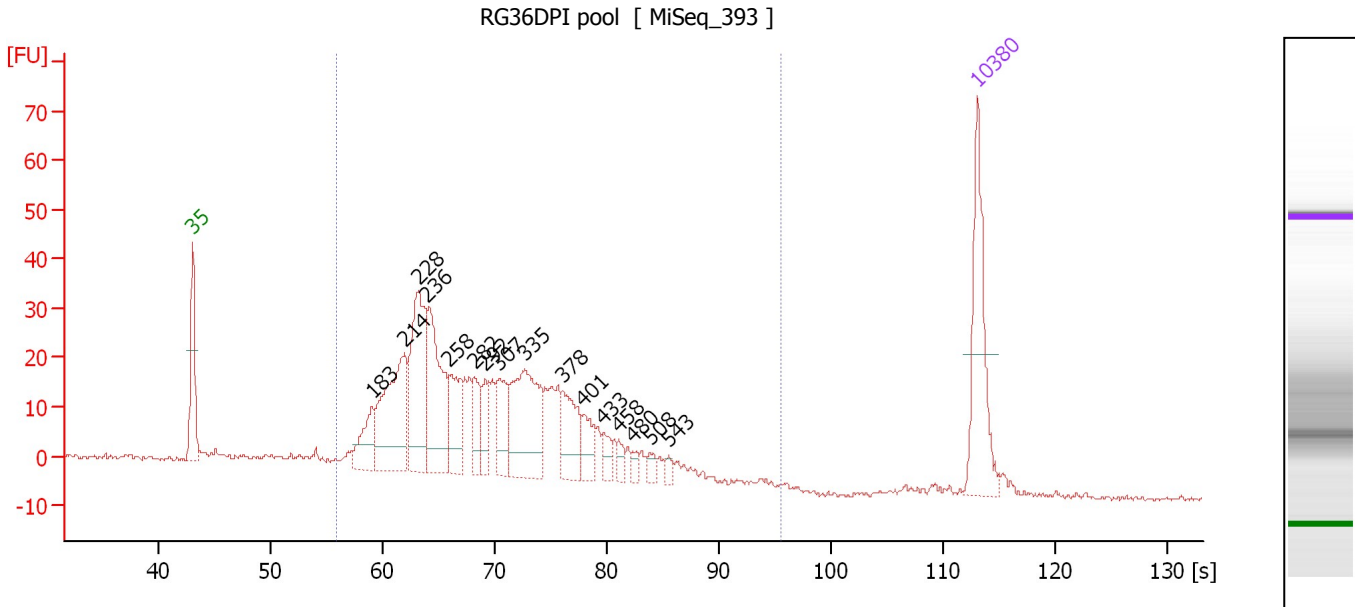
Region table for sample 4 : LP 1:7

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
1,868	9,029	6,395	1,788.0	114.8	478.35	97	16.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : RG36DPI pool

Number of peaks found: 16 Corr. Area 1: 660.9
 Noise: 0.4

Peak table for sample 5 : RG36DPI 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	183	48.41	400.0		59.01
3	214	140.34	993.5		61.89
4	228	134.62	896.5		63.14
5	236	121.19	777.0		63.95
6	258	60.24	354.3		65.92
7	282	36.83	198.0		68.17
8	292	25.76	133.7		69.08
9	307	45.17	223.1		70.38
10	335	118.45	535.5		72.63
11	378	48.97	196.2		76.03
12	401	26.05	98.3		77.86
13	433	13.89	48.5		79.68
14	458	9.46	31.3		81.07
15	480	6.73	21.2		82.32
16	508	8.20	24.5		83.85
17	543	5.15	14.4		85.53
18	10,380	75.00	10.9	Upper Marker	113.00

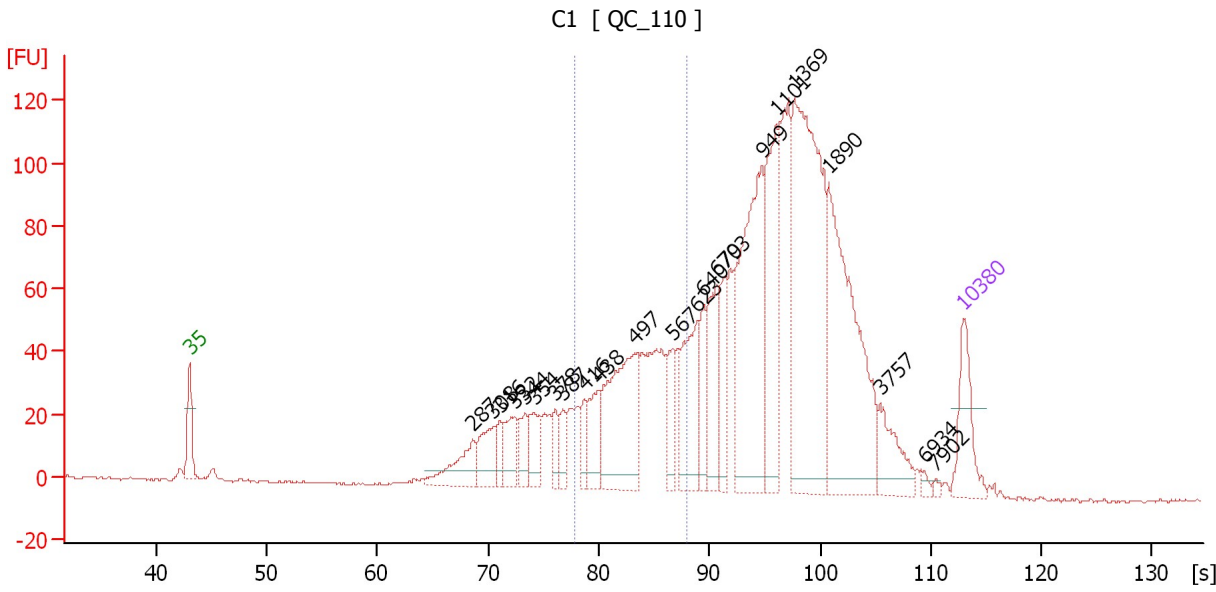
Region table for sample 5 : RG36DPI pool

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
151	1,000	305	660.9	5,590.5	998.21	93	30.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : C1

Number of peaks found: 23 Corr. Area 1: 467.6
 Noise: 0.4

Peak table for sample 6 : C1

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	287	86.17	455.1		68.63
3	308	77.82	382.8		70.48
4	316	34.14	163.7		71.11
5	332	64.79	295.8		72.37
6	344	45.60	200.8		73.34
7	354	61.17	261.9		74.12
8	378	32.35	129.5		76.06
9	387	32.28	126.4		76.74
10	416	34.93	127.3		78.68
11	438	71.63	247.7		79.94
12	497	250.31	763.3		83.29
13	567	62.62	167.4		86.64
14	623	145.72	354.4		88.92
15	640	59.50	140.8		89.46
16	679	120.74	269.3		90.67
17	703	75.61	162.9		91.35
18	949	340.38	543.5		94.75
19	1,101	197.23	271.4		96.06
20	1,369	481.84	533.1		97.66
21	1,890	331.69	265.9		100.77
22	3,757	68.13	27.5		105.43
23	6,934	6.61	1.4		109.41
24	7,902	4.20	0.8		110.43
25	10,380	75.00	10.9	Upper Marker	113.00


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...

... Region table for sample 6 :

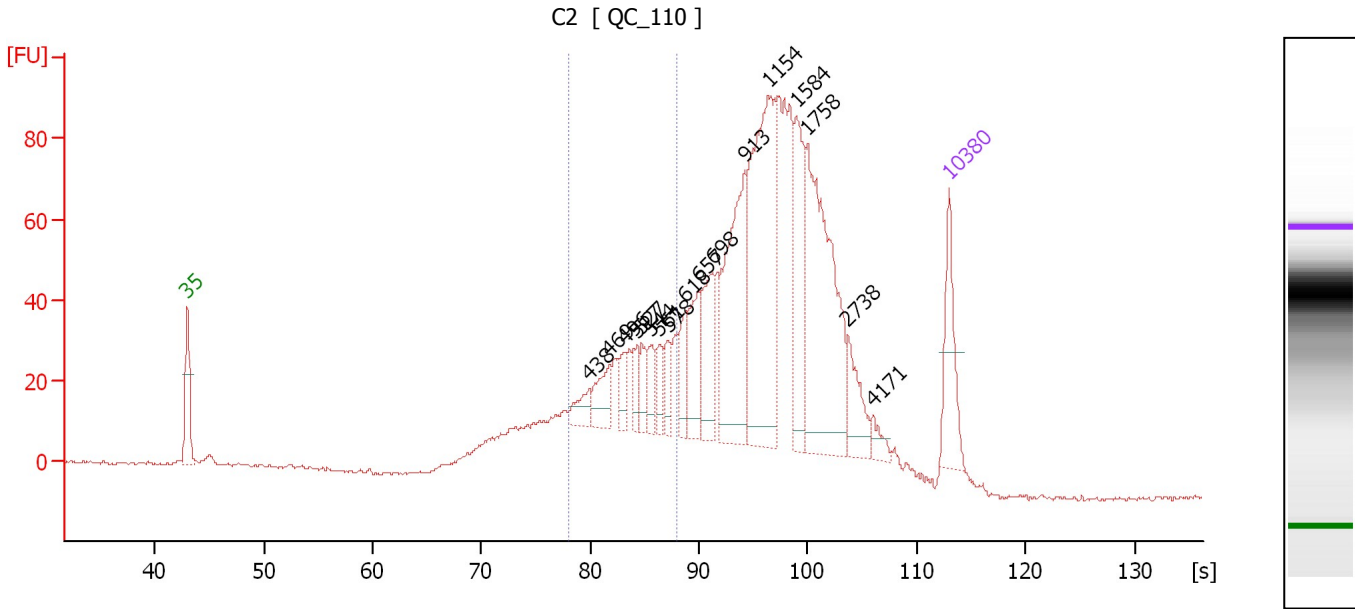
C1

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
401	594	501	467.6	2,206.0	 719.61	20	10.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : C2

Number of peaks found: 17 Corr. Area 1: 334.5
 Noise: 0.4

Peak table for sample 7 : C2

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	438	30.75	106.3		79.95
3	469	50.79	164.0		81.72
4	496	27.69	84.6		83.25
5	517	27.70	81.1		84.29
6	527	31.93	91.8		84.73
7	544	31.74	88.3		85.57
8	561	29.55	79.8		86.36
9	578	28.17	73.9		87.15
10	618	48.25	118.2		88.78
11	657	72.37	167.0		89.96
12	698	90.96	197.4		91.25
13	913	223.61	371.0		94.25
14	1,154	333.90	438.3		96.38
15	1,584	121.32	116.0		98.94
16	1,758	282.68	243.7		99.98
17	2,738	52.34	29.0		103.68
18	4,171	15.91	5.8		105.95
19	10,380	75.00	10.9	Upper Marker	113.00

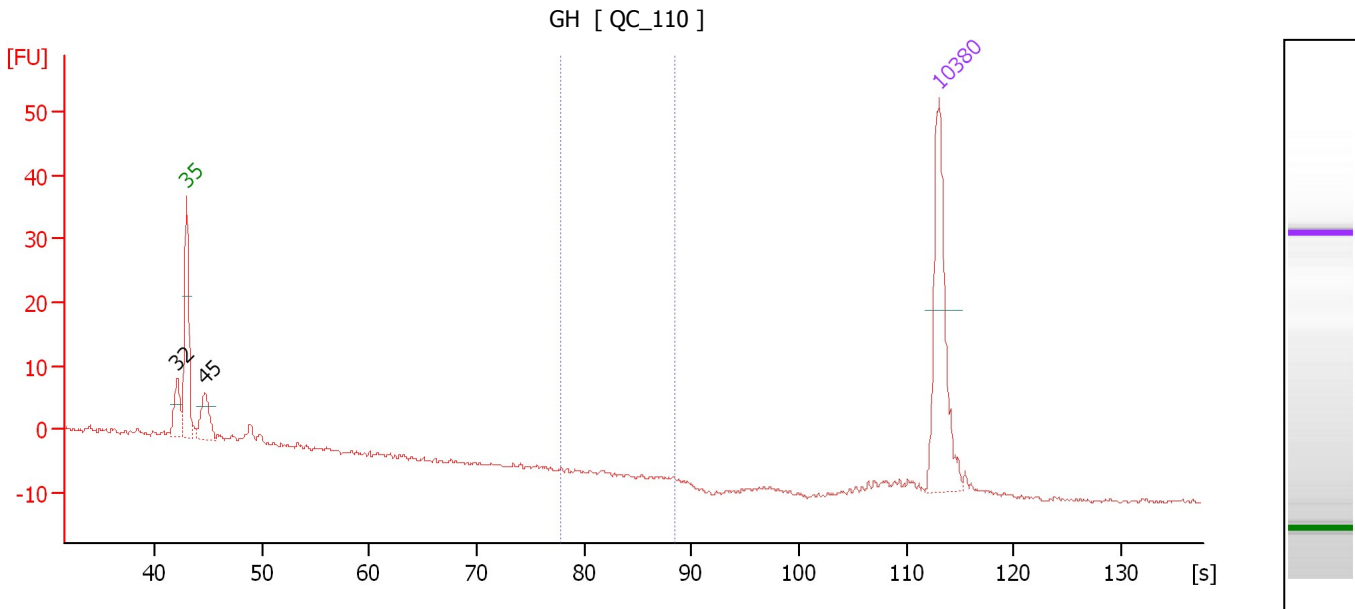
Region table for sample 7 : C2

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
404	595	505	334.5	1,780.4	585.43	20	10.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : GH

Number of peaks found: 2 Corr. Area 1: 0.0
 Noise: 0.3

Peak table for sample 8 : GH

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.10
2	35	125.00	5,411.3	Lower Marker	43.00
3	45	35.81	1,193.8		44.65
4	10,380	75.00	10.9	Upper Marker	113.00

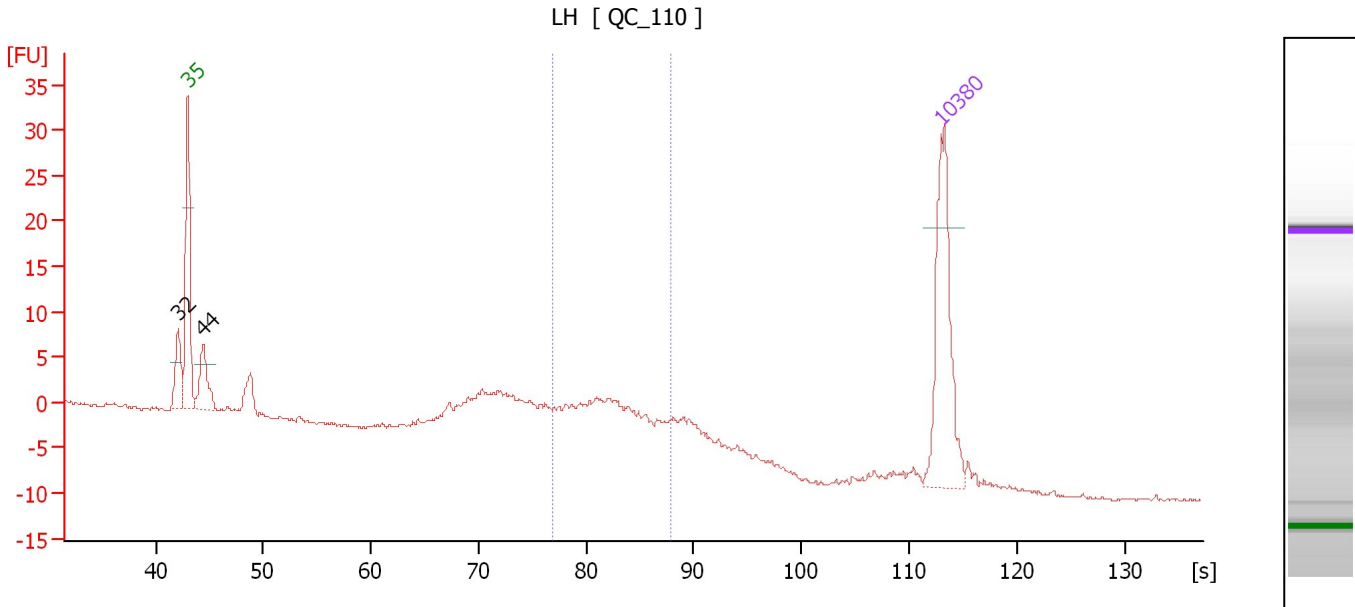
Region table for sample 8 : GH

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
402	607	0	0.0	0.0	0.00	0	0.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : LH

Number of peaks found: 2 Corr. Area 1: 59.6
 Noise: 0.2

Peak table for sample 9 : LH

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.10
2	35	125.00	5,411.3	Lower Marker	43.00
3	44	40.82	1,410.8		44.40
4	10,380	75.00	10.9	Upper Marker	113.00

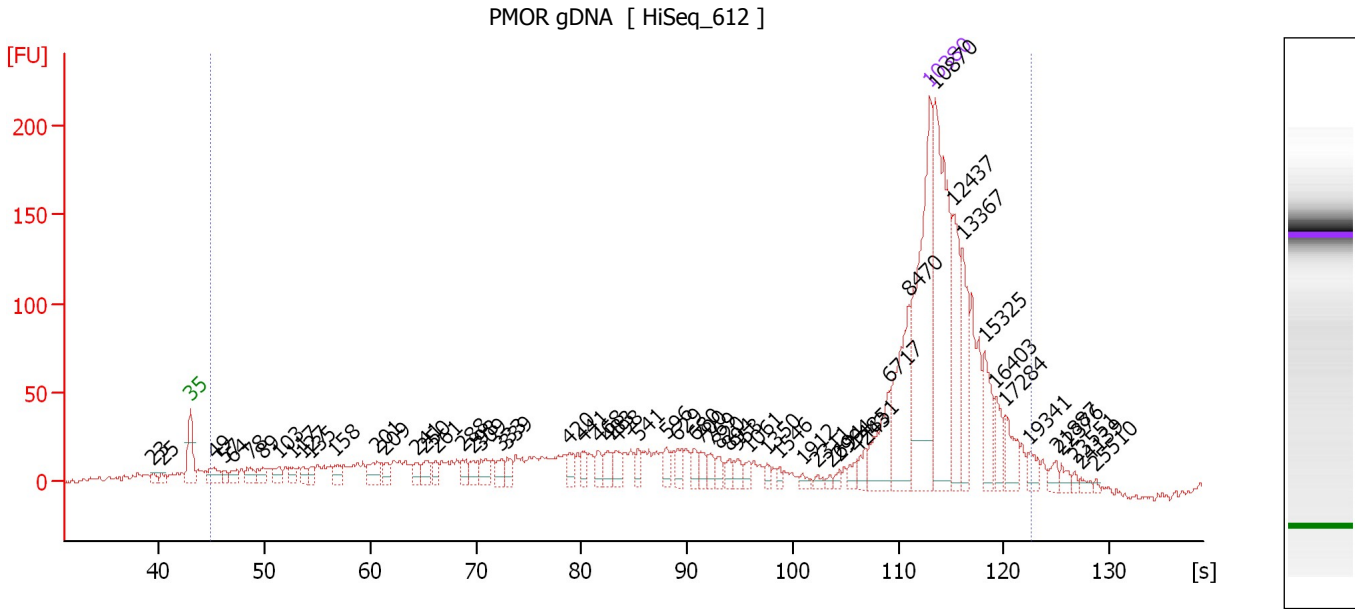
Region table for sample 9 : LH

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
389	592	479	59.6	373.2	116.11	29	11.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : PMOR gDNA

Number of peaks found: 59 Corr. Area 1: 1,923.6
 Noise: 1.5

Peak table for sample 10 : PMOR gDNA


Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	23	0.00	0.0		39.44
2	25	0.00	0.0		40.25
3	35	125.00	5,411.3	Lower Marker	43.00
4	49	11.25	349.1		45.19
5	57	4.61	122.9		46.15
6	64	6.89	161.9		47.02
7	78	8.84	171.1		48.59
8	89	7.97	135.7		49.81
9	103	7.70	113.4		51.34
10	117	5.97	77.2		52.71
11	127	5.91	70.4		53.68
12	135	5.56	62.3		54.44
13	158	7.92	75.9		56.62
14	201	12.12	91.3		60.69
15	209	6.04	43.7		61.45
16	241	6.36	40.0		64.40
17	250	7.56	45.8		65.21
18	261	5.23	30.3		66.23
19	288	6.70	35.2		68.77
20	298	6.67	33.9		69.69
21	309	8.69	42.6		70.55
22	333	6.14	28.0		72.43
23	339	5.93	26.5		72.94
24	420	6.65	24.0		78.89
25	441	5.52	19.0		80.11

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad


Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...

... Peak table for sample 10 : PMOR gDNA

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	468	6.20	20.1		81.63
27	483	7.49	23.5		82.50
28	498	7.61	23.1		83.36
29	541	5.30	14.8		85.40
30	596	5.04	12.8		88.04
31	629	5.89	14.2		89.11
32	680	5.48	12.2		90.68
33	700	5.85	12.7		91.29
34	769	4.96	9.8		92.26
35	820	4.55	8.4		92.97
36	894	3.76	6.4		93.99
37	953	4.42	7.0		94.80
38	1,061	3.58	5.1		95.82
39	1,350	2.60	2.9		97.55
40	1,546	2.05	2.0		98.72
41	1,912	1.93	1.5		100.90
42	2,311	1.89	1.2		102.38
43	2,694	1.27	0.7		103.54
44	2,994	1.73	0.9		104.46
45	4,243	3.32	1.2		106.04
46	4,851	5.01	1.6		106.80
47	6,717	19.26	4.3		109.14
48	8,470	34.76	6.2		111.02
49	 10,380	75.00	10.9	Upper Marker	113.00
50	10,870	0.00	0.0		113.51
51	12,437	0.00	0.0		115.14
52	13,367	0.00	0.0		116.10
53	15,325	0.00	0.0		118.13
54	16,403	0.00	0.0		119.25
55	17,284	0.00	0.0		120.17
56	19,341	0.00	0.0		122.30
57	21,887	0.00	0.0		124.95
58	22,376	0.00	0.0		125.45
59	23,551	0.00	0.0		126.67
60	24,139	0.00	0.0		127.28
61	25,510	0.00	0.0		128.71

Region table for sample 10 : PMOR gDNA

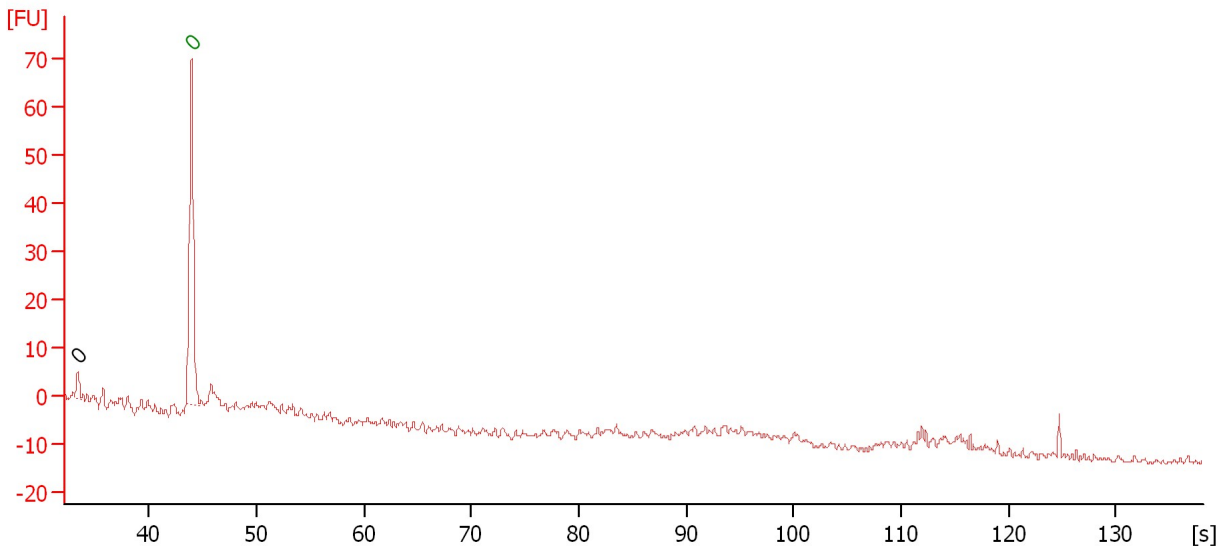
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
46	19,651	7,677	1,923.5	2,833.5	 623.31	93	75.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
Modified: 8/11/2016 5:48:04 PM

Electropherogram Summary Continued ...

PMOR Mut gDNA [HiSeq_612]



Overall Results for sample 11 : PMOR Mut gDNA

Number of peaks found: 1 Noise: 0.9

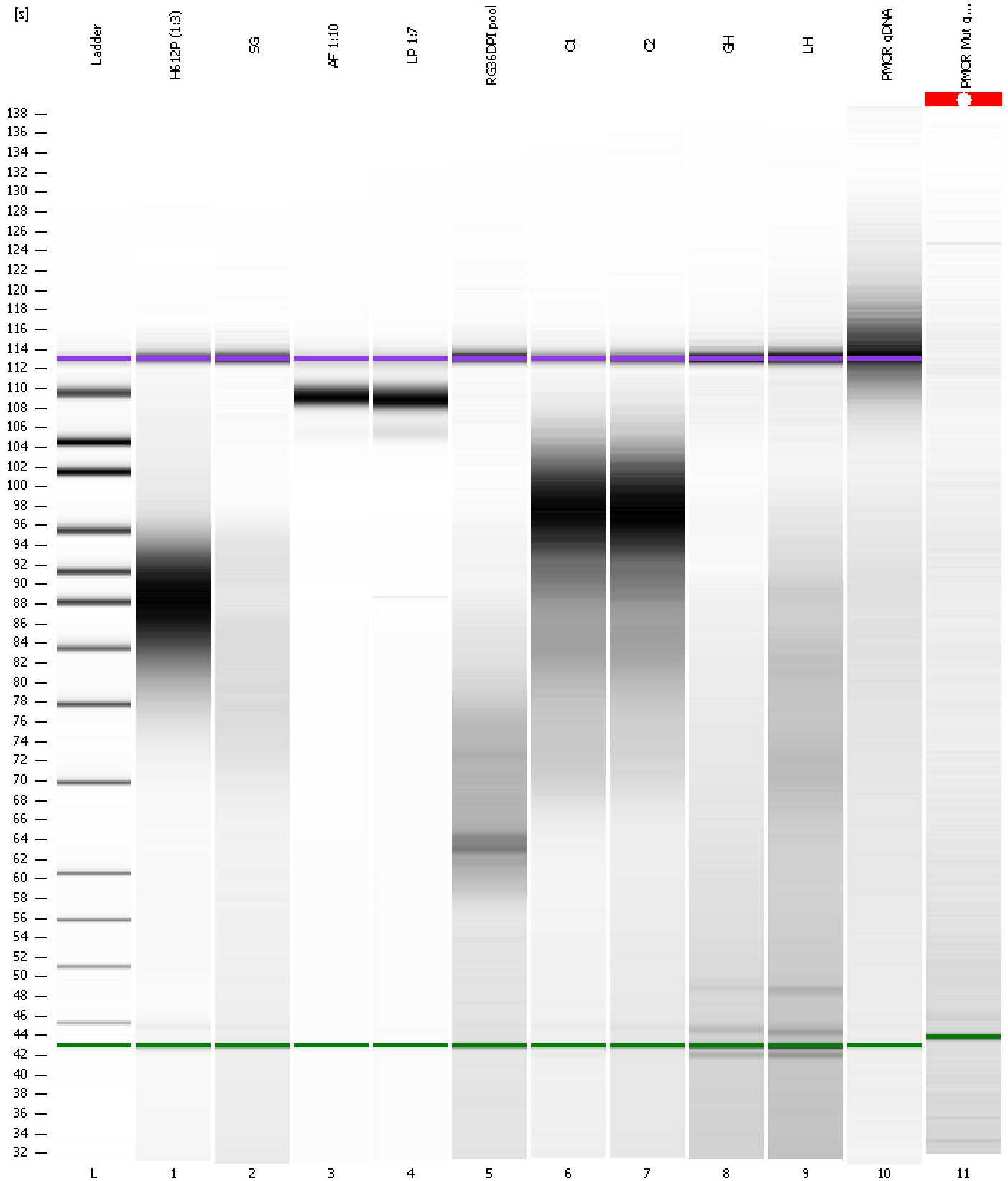
Peak table for sample 11 : PMOR Mut gDNA

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	0	0.00	0.0		33.35
2	0	0.00	0.0	Lower Marker	43.95

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
Modified: 8/11/2016 5:48:04 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad

Created: 8/11/2016 5:07:37 PM
 Modified: 8/11/2016 5:48:04 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		8/11/2016 5:48:03 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-08-11\2016-08-11_002.xad)		Instrument	Run		8/11/2016 5:07:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		8/11/2016 5:07:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		8/11/2016 5:07:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		8/11/2016 5:07:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		8/11/2016 5:07:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		8/11/2016 5:07:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		8/11/2016 5:07:43 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1