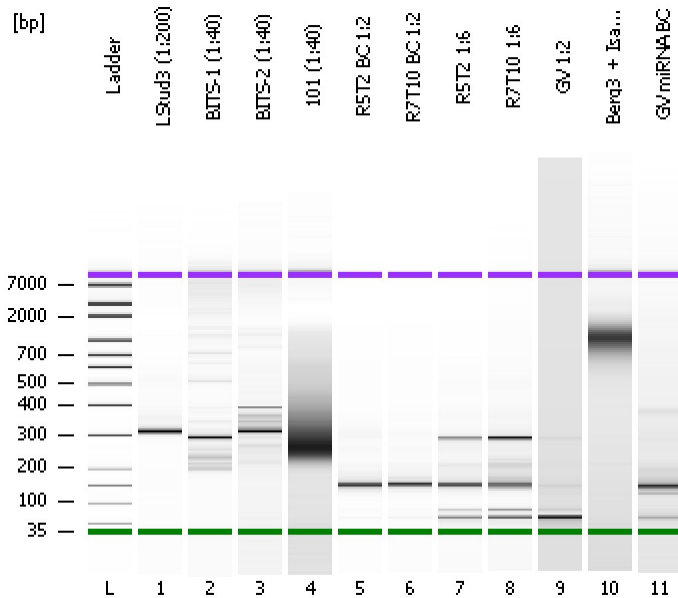


Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electrophoresis File Run Summary**



Instrument Information:

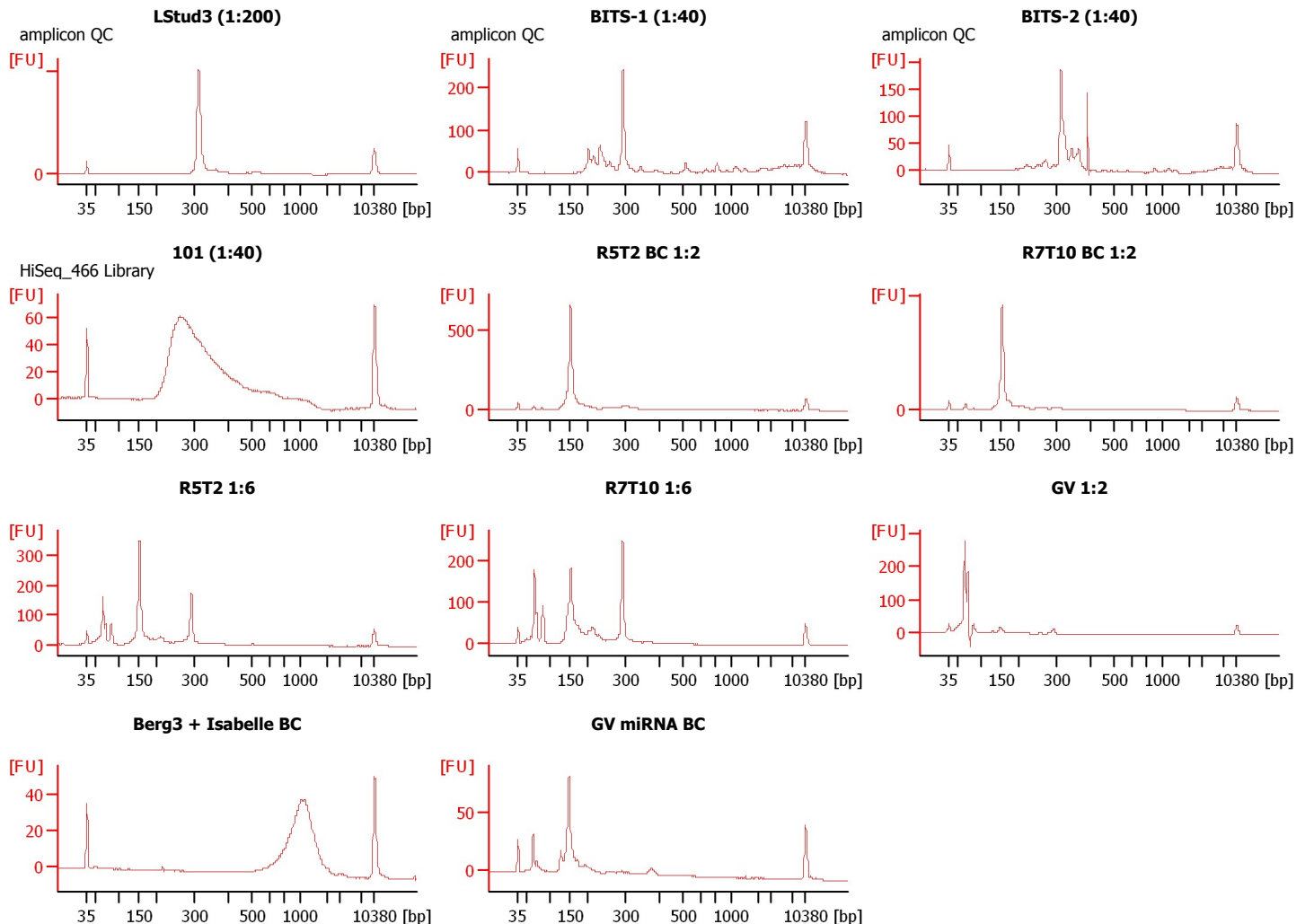
Instrument Name: DE13701086      Firmware: C.01.069  
 Serial#: DE13701086      Type: G2938B

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\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
Modified: 5/3/2016 1:30:04 PM

**Electrophoresis File Run Summary (Chip Summary)**

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
LStud3 (1:200)	amplicon QC	<input type="checkbox"/>	✓			
BITS-1 (1:40)	amplicon QC	<input type="checkbox"/>	✓			
BITS-2 (1:40)	amplicon QC	<input type="checkbox"/>	✓			
101 (1:40)	HiSeq_466 Library	<input type="checkbox"/>	✓			
R5T2 BC 1:2		<input type="checkbox"/>	✓			
R7T10 BC 1:2		<input type="checkbox"/>	✓			
R5T2 1:6		<input type="checkbox"/>	✓			
R7T10 1:6		<input type="checkbox"/>	✓			
GV 1:2		<input type="checkbox"/>	✓			
Berg3 + Isabelle BC		<input type="checkbox"/>	✓			
GV miRNA BC		<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\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
Modified: 5/3/2016 1:30:04 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

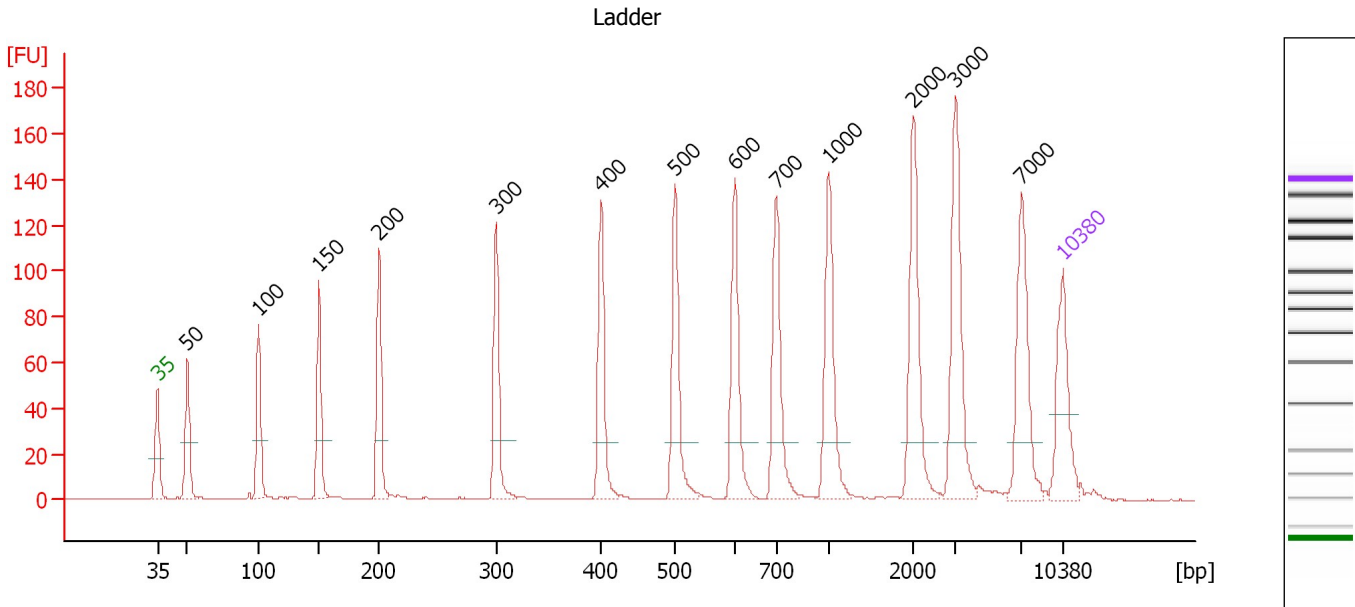
**Ladder**

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\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.2

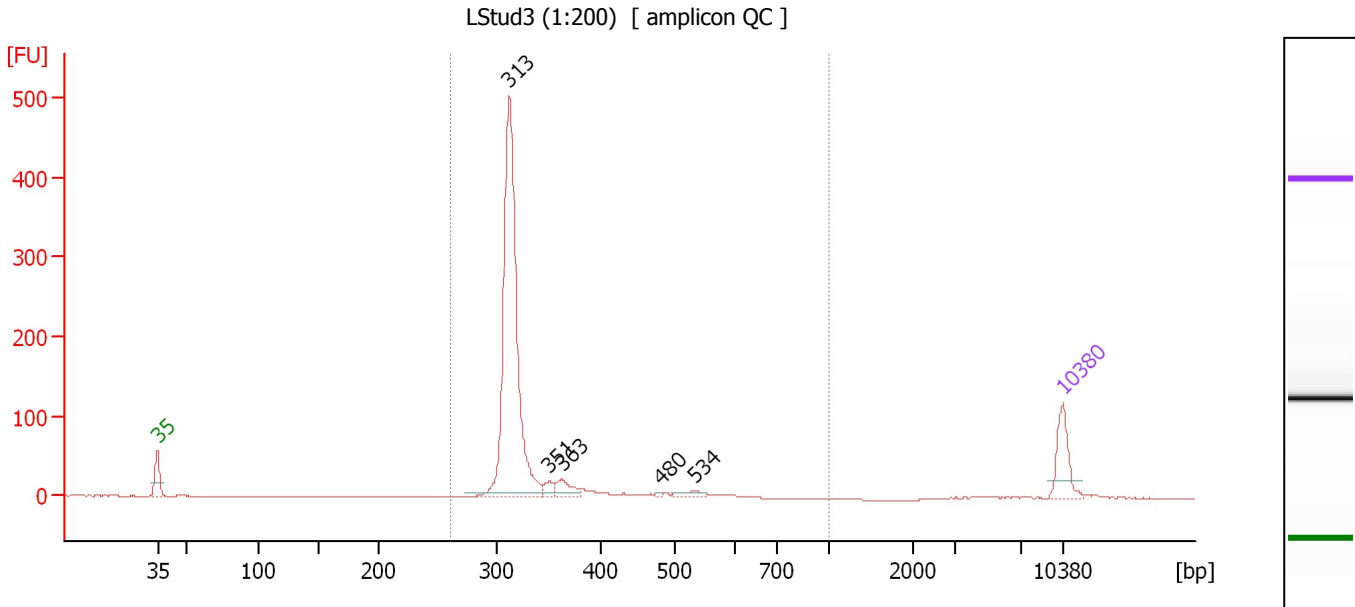
**Peak table for Ladder**

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	45.31
3	100	150.00	2,272.7	Ladder Peak	50.81
4	150	150.00	1,515.2	Ladder Peak	55.47
5	200	150.00	1,136.4	Ladder Peak	60.14
6	300	150.00	757.6	Ladder Peak	69.17
7	400	150.00	568.2	Ladder Peak	77.32
8	500	150.00	454.5	Ladder Peak	83.01
9	600	150.00	378.8	Ladder Peak	87.63
10	700	150.00	324.7	Ladder Peak	90.83
11	1,000	150.00	227.3	Ladder Peak	94.92
12	2,000	150.00	113.6	Ladder Peak	101.42
13	3,000	150.00	75.8	Ladder Peak	104.67
14	7,000	150.00	32.5	Ladder Peak	109.85
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : LStud3 (1:200)**

Number of peaks found: 5                      Corr. Area 1: 1,000.8  
 Noise: 0.2

**Peak table for sample 1 : LStud3 (1:200)**

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	313	943.25	4,567.4		70.22
3	351	24.22	104.7		73.30
4	363	43.35	181.0		74.29
5	480	4.01	12.7		81.85
6	534	19.25	54.6		84.59
7	10,380	75.00	10.9	Upper Marker	113.00

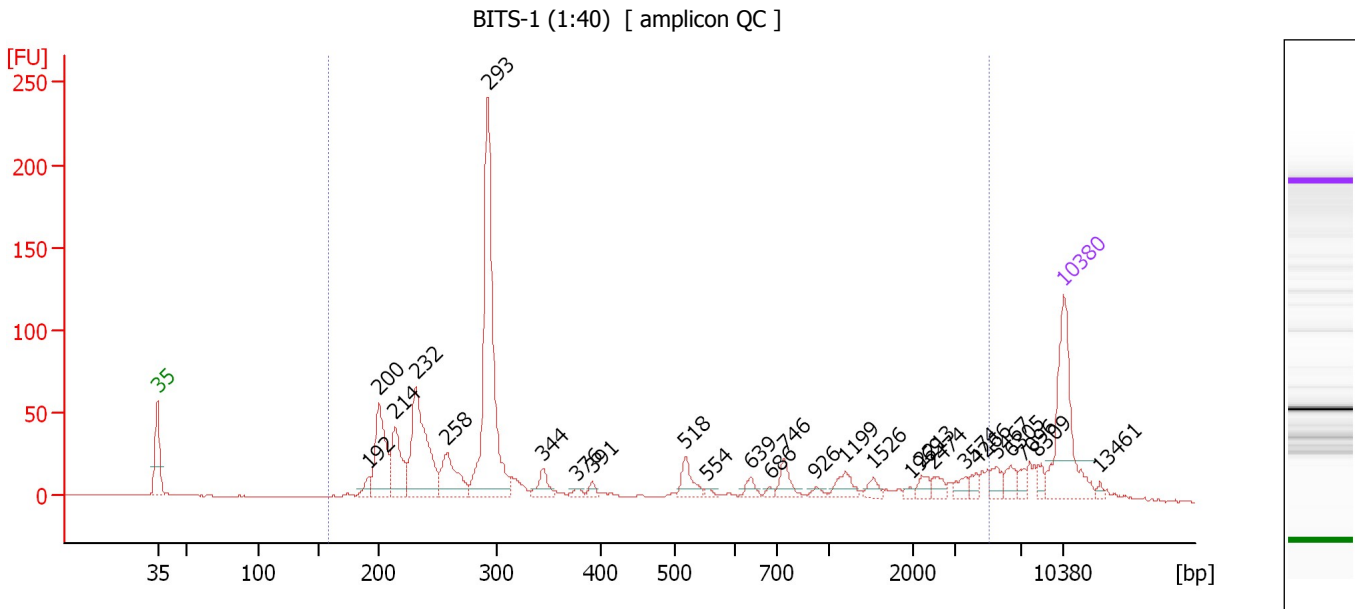
**Region table for sample 1 : LStud3 (1:200)**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
262	1,000	331	1,000.8	5,017.4	1,075.38	98	14.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : BITS-1 (1:40)**

Number of peaks found: 27                      Corr. Area 1: 983.4  
 Noise: 0.2

**Peak table for sample 2 : BITS-1 (1:40)**

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	192	12.82	101.3		59.36
3	200	83.31	632.0		60.11
4	214	62.37	442.0		61.38
5	232	131.57	860.9		62.99
6	258	51.87	304.5		65.39
7	293	263.48	1,362.1		68.55
8	344	16.94	74.5		72.79
9	376	5.21	21.0		75.38
10	391	5.91	22.9		76.61
11	518	21.36	62.5		83.82
12	554	3.34	9.1		85.52
13	639	8.58	20.3		88.87
14	686	3.77	8.3		90.37
15	746	17.06	34.7		91.46
16	926	4.43	7.3		93.91
17	1,199	14.80	18.7		96.22
18	1,526	7.15	7.1		98.34
19	1,969	2.76	2.1		101.22
20	2,213	7.61	5.2		102.11
21	2,474	7.71	4.7		102.96
22	3,574	6.93	2.9		105.41
23	4,266	5.82	2.1		106.31
24	5,467	9.06	2.5		107.86
25	6,305	10.51	2.5		108.95

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary Continued ...**

**... Peak table for sample 2 : BITS-1 (1:40)**

Peak	Size [bp]	Conc. [pg/ $\mu$ l]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	7,096	6.06	1.3		109.94
27	8,309	6.17	1.1		111.07
28	10,380	75.00	10.9	Upper Marker	113.00
29	13,461	0.00	0.0		115.88

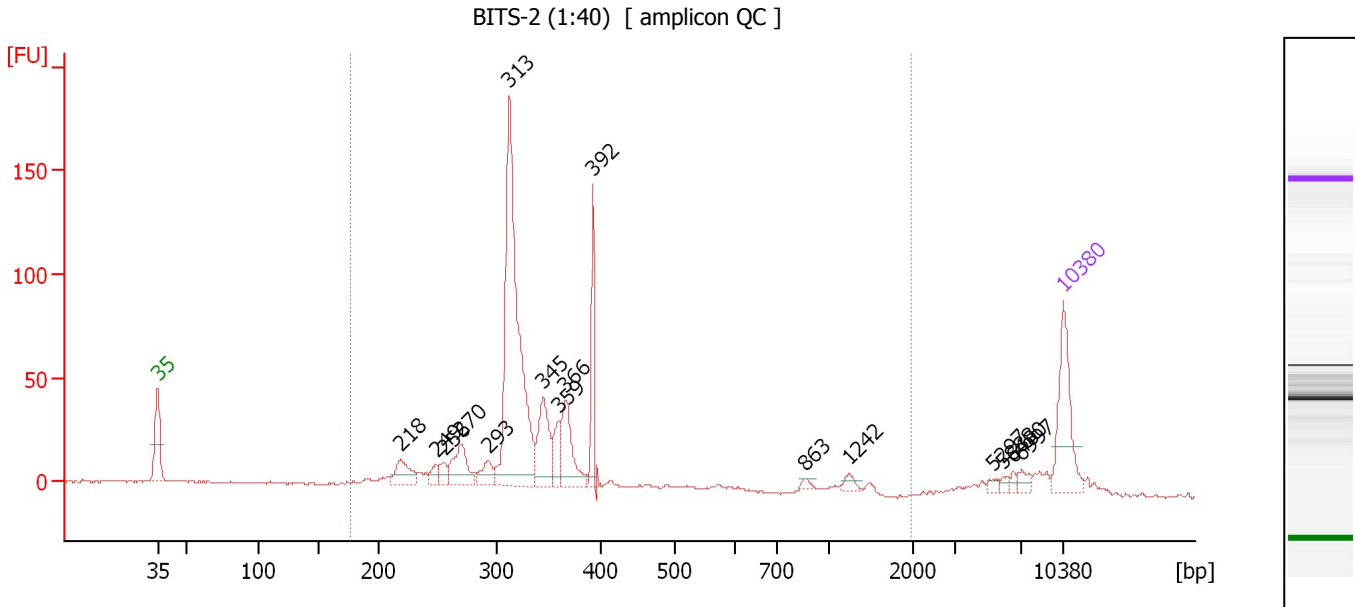
**Region table for sample 2 : BITS-1 (1:40)**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ $\mu$ l]	% of Total	Size distribution in CV [%]
159	5,090	747	983.4	4,041.3	808.31	91	100.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : BITS-2 (1:40)**

Number of peaks found: 16                      Corr. Area 1: 681.0  
 Noise: 0.4

**Peak table for sample 3 : BITS-2 (1:40)**

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	218	42.33	293.6		61.80
3	249	15.03	91.6		64.53
4	256	15.70	93.1		65.15
5	270	59.48	334.0		66.44
6	293	26.46	136.8		68.55
7	313	431.53	2,089.5		70.22
8	345	78.05	342.6		72.86
9	359	32.34	136.6		73.96
10	366	74.40	308.2		74.53
11	392	65.06	251.7		76.64
12	863	5.34	9.4		93.05
13	1,242	8.08	9.9		96.49
14	5,297	4.78	1.4		107.64
15	5,888	4.48	1.2		108.41
16	6,480	4.93	1.2		109.17
17	6,997	8.63	1.9		109.84
18	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 3 : BITS-2 (1:40)**

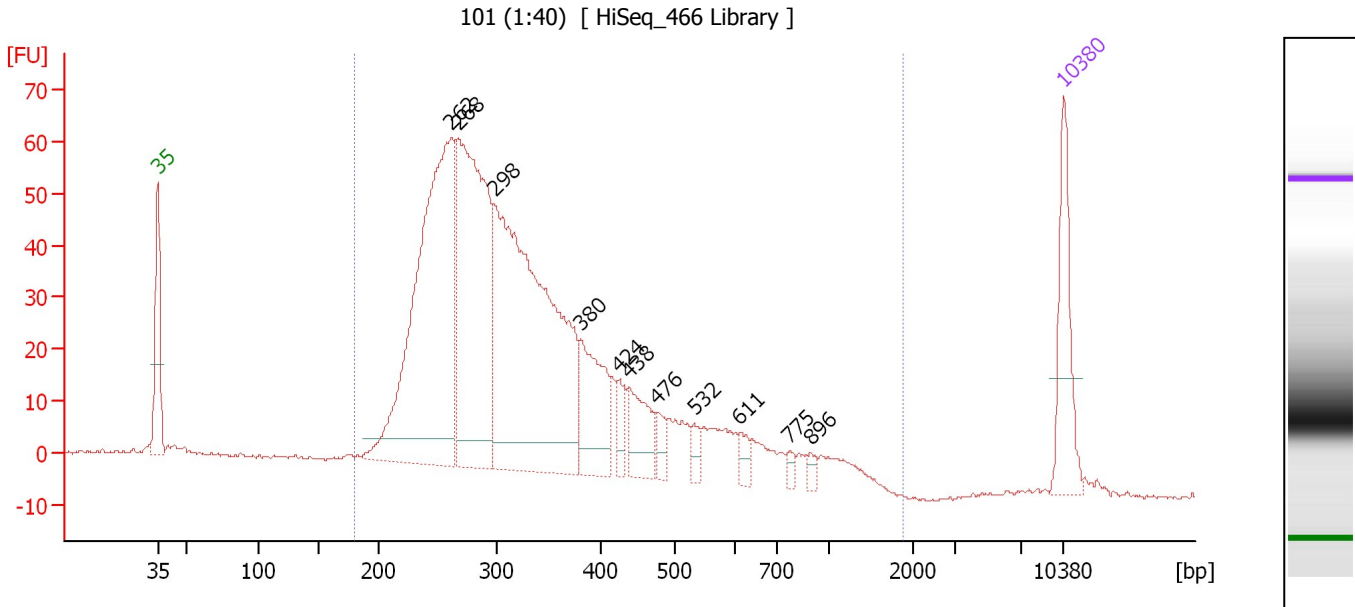
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
176	1,985	360	681.0	4,465.7	942.89	88	49.6



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : 101 (1:40)**

Number of peaks found: 11                      Corr. Area 1: 1,213.1  
 Noise: 0.4

**Peak table for sample 4 : 101 (1:40)**

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	262	587.84	3,403.0		65.71
3	268	452.39	2,560.7		66.25
4	298	637.38	3,235.5		69.04
5	380	110.91	441.7		75.73
6	424	24.48	87.4		78.71
7	438	57.37	198.4		79.49
8	476	18.48	58.9		81.64
9	532	13.19	37.6		84.47
10	611	13.75	34.1		87.99
11	775	8.02	15.7		91.85
12	896	7.75	13.1		93.51
13	10,380	75.00	10.9	Upper Marker	113.00

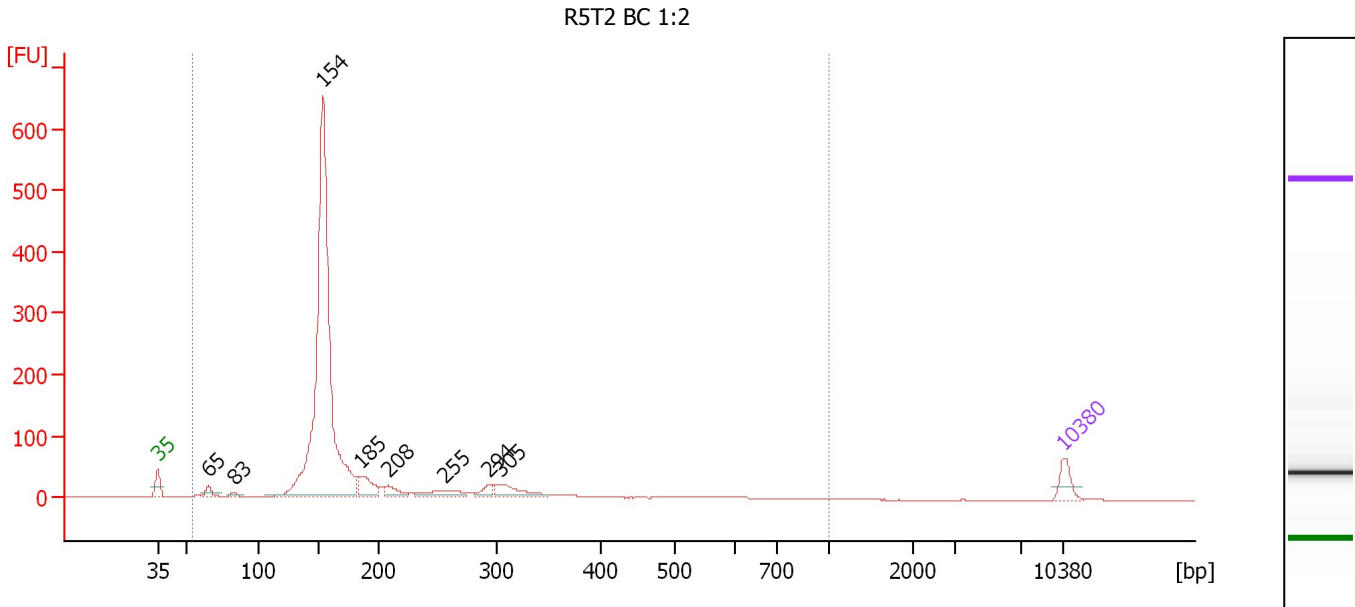
**Region table for sample 4 : 101 (1:40)**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
179	1,880	355	1,213.1	10,327.4	2,086.87	95	44.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : R5T2 BC 1:2**

Number of peaks found: 8                      Corr. Area 1: 1,830.4  
 Noise: 0.3

**Peak table for sample 5 : R5T2 BC 1:2**

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	65	51.08	1,196.7		46.92
3	83	24.71	450.3		48.96
4	154	3,160.22	31,177.2		55.81
5	185	149.17	1,221.7		58.74
6	208	74.01	540.1		60.82
7	255	107.90	640.1		65.14
8	294	56.40	290.8		68.62
9	305	134.40	668.2		69.56
10	10,380	75.00	10.9	Upper Marker	113.00

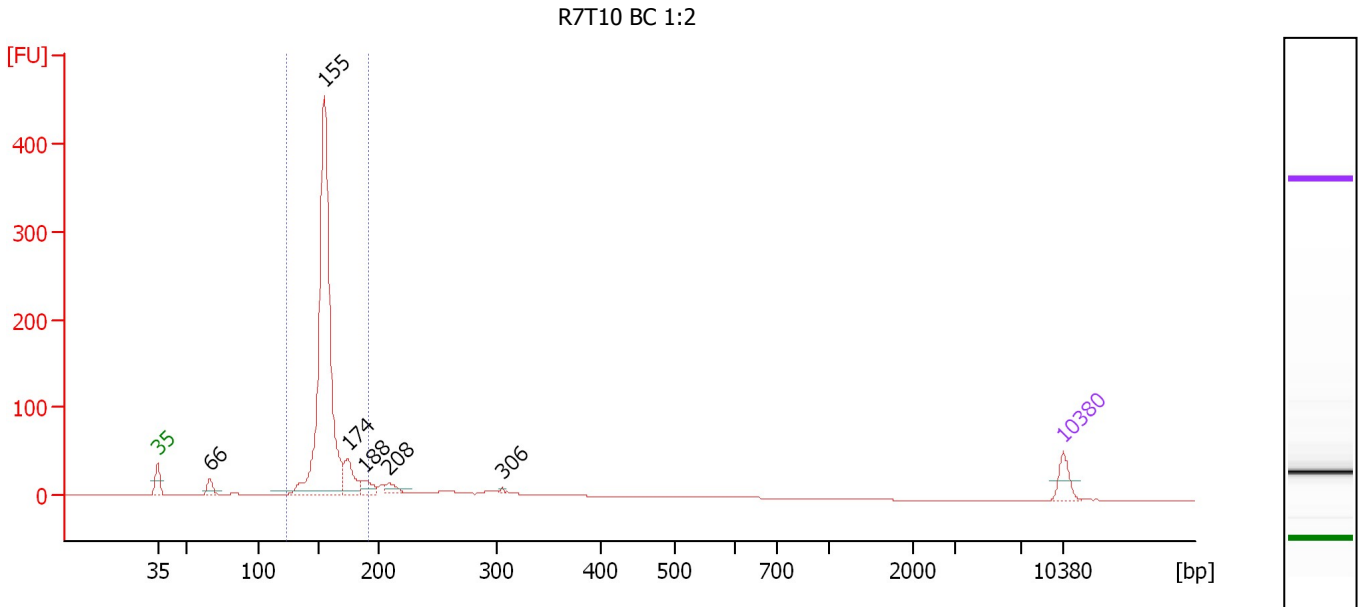
**Region table for sample 5 : R5T2 BC 1:2**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
54	1,000	190	1,830.4	38,547.6	4,096.81	99	47.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : R7T10 BC 1:2**

Number of peaks found: 6                      Corr. Area 1: 888.0  
 Noise: 0.2

**Peak table for sample 6 : R7T10 BC 1:2**

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	66	75.08	1,731.8		47.03
3	155	2,402.27	23,531.7		55.91
4	174	177.70	1,546.0		57.73
5	188	76.39	614.9		59.04
6	208	56.75	413.5		60.85
7	306	7.46	37.0		69.63
8	10,380	75.00	10.9	Upper Marker	113.00

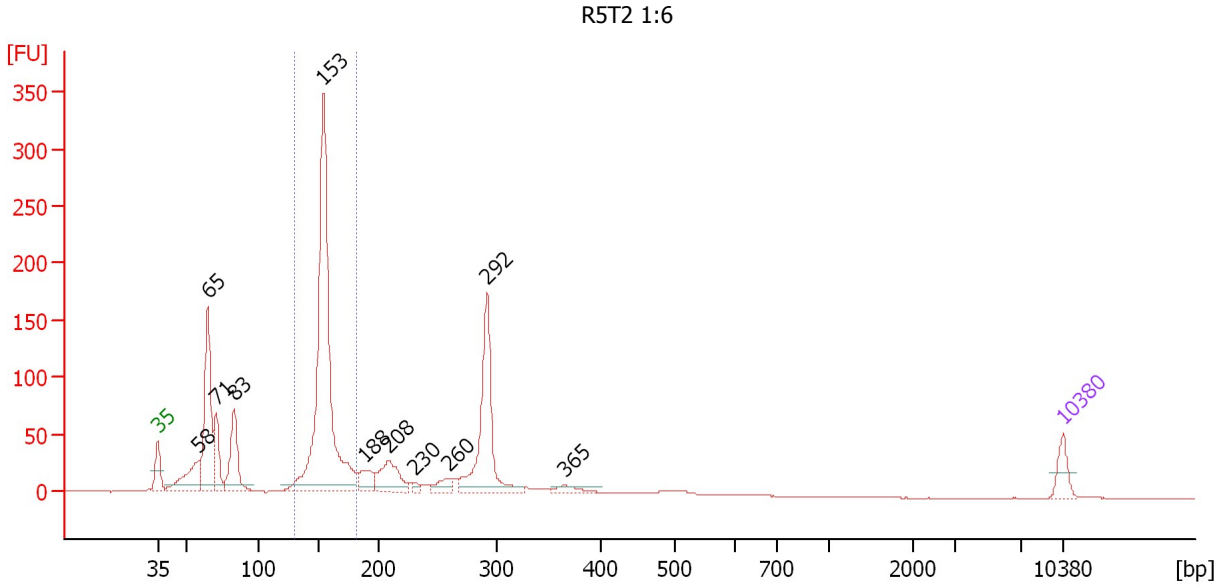
**Region table for sample 6 : R7T10 BC 1:2**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
123	191	157	888.0	26,121.8	2,698.37	80	6.3

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : R5T2 1:6**

Number of peaks found: 11                      Corr. Area 1: 641.7  
 Noise: 0.2

**Peak table for sample 7 : R5T2 1:6**

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	58	267.31	6,984.5		46.19
3	65	663.83	15,589.3		46.91
4	71	216.44	4,646.0		47.57
5	83	335.88	6,118.0		48.96
6	153	2,063.66	20,374.9		55.80
7	188	121.53	978.4		59.04
8	208	228.35	1,661.0		60.89
9	230	21.62	142.1		62.89
10	260	73.47	428.1		65.56
11	292	667.13	3,456.2		68.49
12	365	45.27	188.0		74.45
13	10,380	75.00	10.9	Upper Marker	113.00

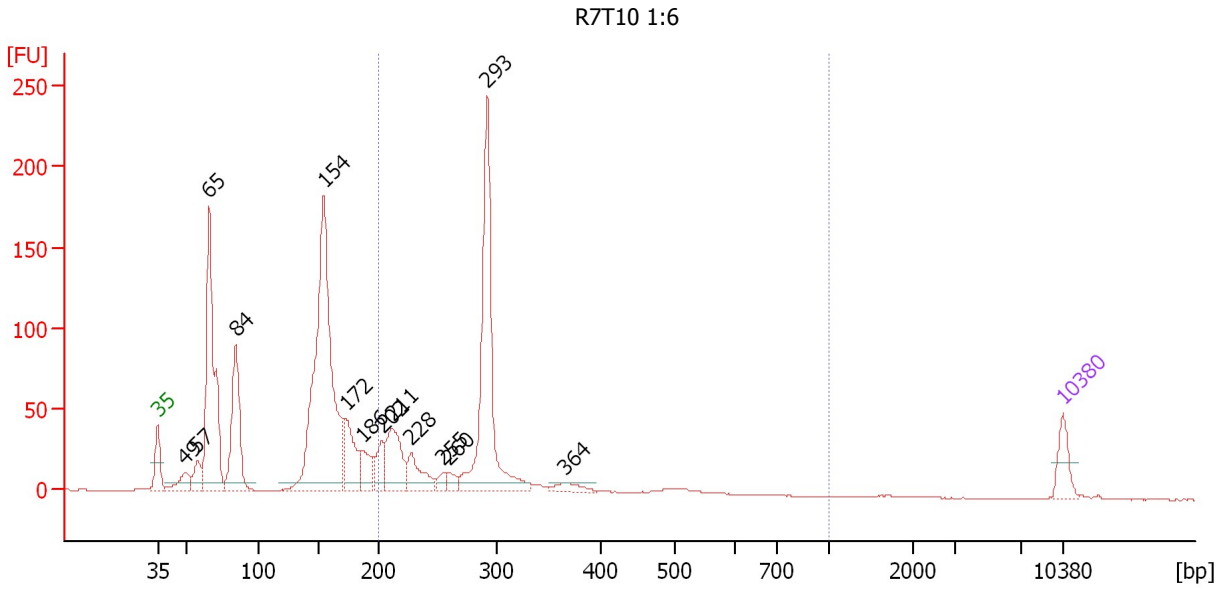
**Region table for sample 7 : R5T2 1:6**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
129	182	155	641.7	20,386.2	2,077.97	39	5.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : R7T10 1:6**

Number of peaks found: 14                      Corr. Area 1: 591.2  
 Noise: 0.2

**Peak table for sample 8 : R7T10 1:6**

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	49	102.69	3,178.1		45.15
3	57	96.97	2,594.7		46.04
4	65	967.64	22,492.5		46.98
5	84	475.89	8,564.0		49.07
6	154	1,636.21	16,062.3		55.88
7	172	255.09	2,242.9		57.55
8	186	129.39	1,051.9		58.86
9	202	113.29	849.1		60.33
10	211	280.35	2,009.1		61.17
11	228	154.47	1,028.2		62.63
12	255	39.47	234.6		65.09
13	260	42.39	247.5		65.51
14	293	961.44	4,979.9		68.50
15	364	48.35	201.0		74.41
16	10,380	75.00	10.9	Upper Marker	113.00

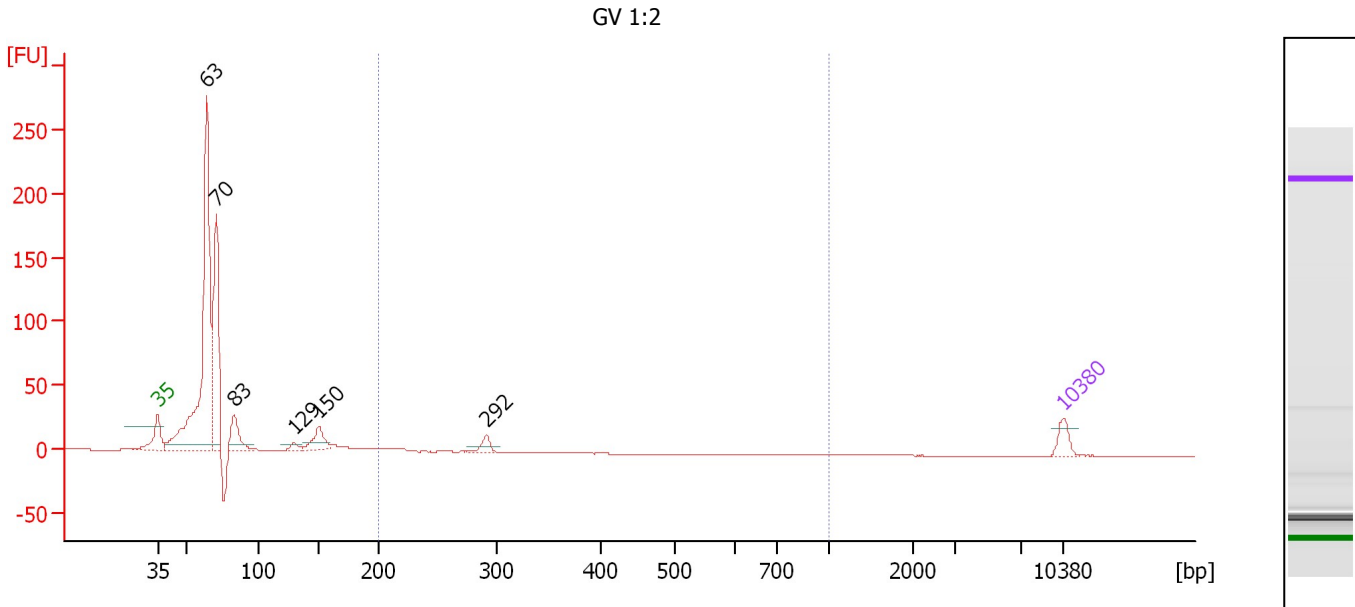
**Region table for sample 8 : R7T10 1:6**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	289	591.2	9,682.8	1,775.68	36	23.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : GV 1:2**

Number of peaks found: 6                      Corr. Area 1: 28.6  
 Noise: 0.1

**Peak table for sample 9 : GV 1:2**

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	63	2,596.87	62,217.3		46.77
3	70	1,090.91	23,458.1		47.56
4	83	260.77	4,761.6		48.94
5	129	50.08	586.5		53.55
6	150	193.74	1,958.8		55.46
7	292	74.19	384.9		68.45
8	10,380	75.00	10.9	Upper Marker	113.00

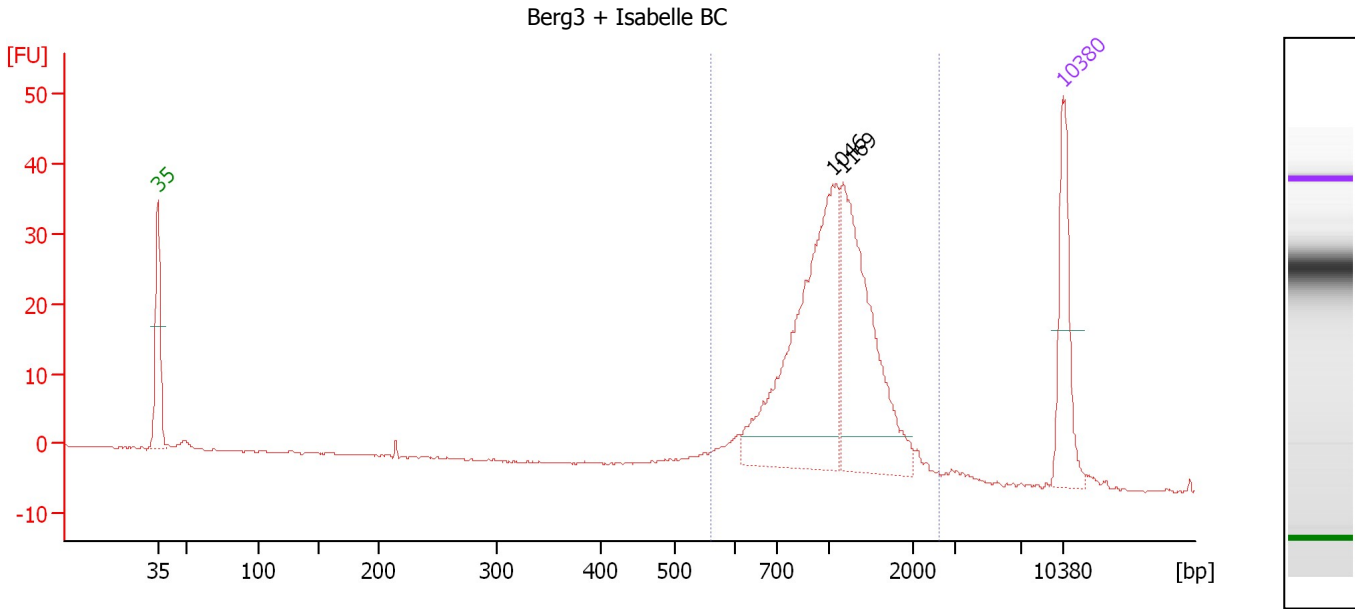
**Region table for sample 9 : GV 1:2**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	262	28.6	835.2	144.49	4	14.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : Berg3 + Isabelle BC**

Number of peaks found: 2                      Corr. Area 1: 302.5  
 Noise: 0.1

**Peak table for sample 10 : Berg3 + Isabelle BC**

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	1,046	312.70	453.1		95.22
3	1,169	226.34	293.4		96.02
4	10,380	75.00	10.9	Upper Marker	113.00

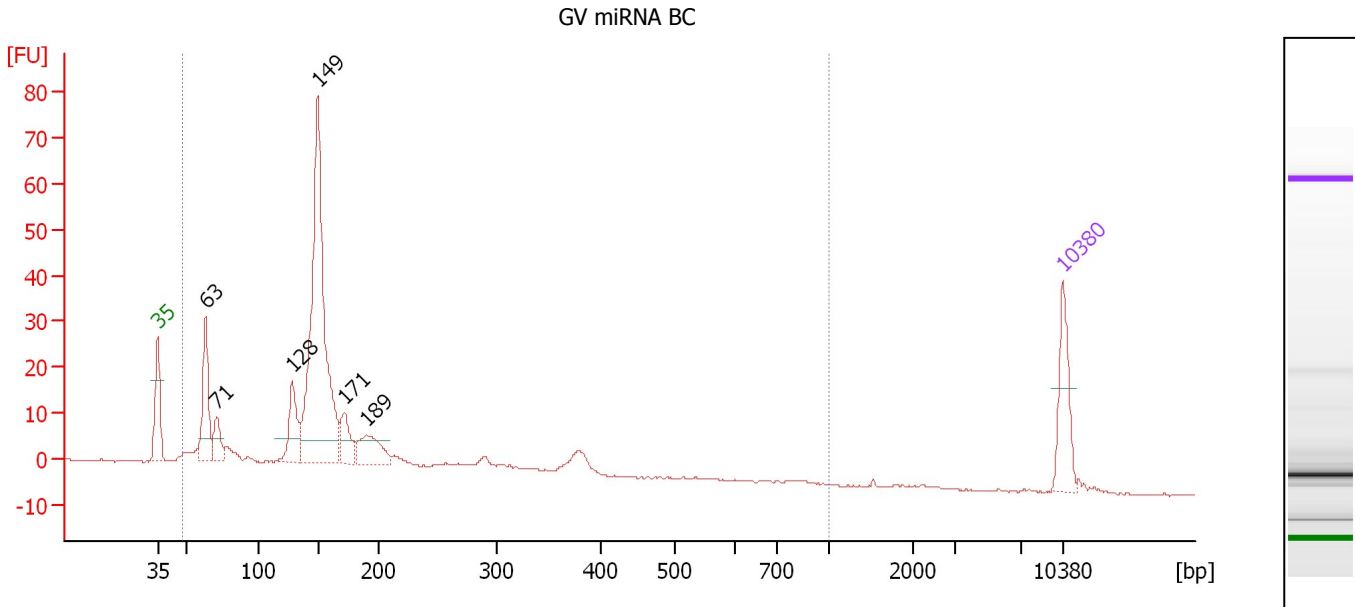
**Region table for sample 10 : Berg3 + Isabelle BC**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
559	2,619	1,110	302.5	902.9	597.89	93	31.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : GV miRNA BC**

Number of peaks found: 6                      Corr. Area 1: 336.4  
 Noise: 0.1

**Peak table for sample 11 : GV miRNA BC**

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	63	155.52	3,751.3		46.72
3	71	51.15	1,097.2		47.58
4	128	99.92	1,179.8		53.45
5	149	701.00	7,121.9		55.39
6	171	64.74	573.3		57.44
7	189	84.11	674.2		59.11
8	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 11 : GV miRNA BC**

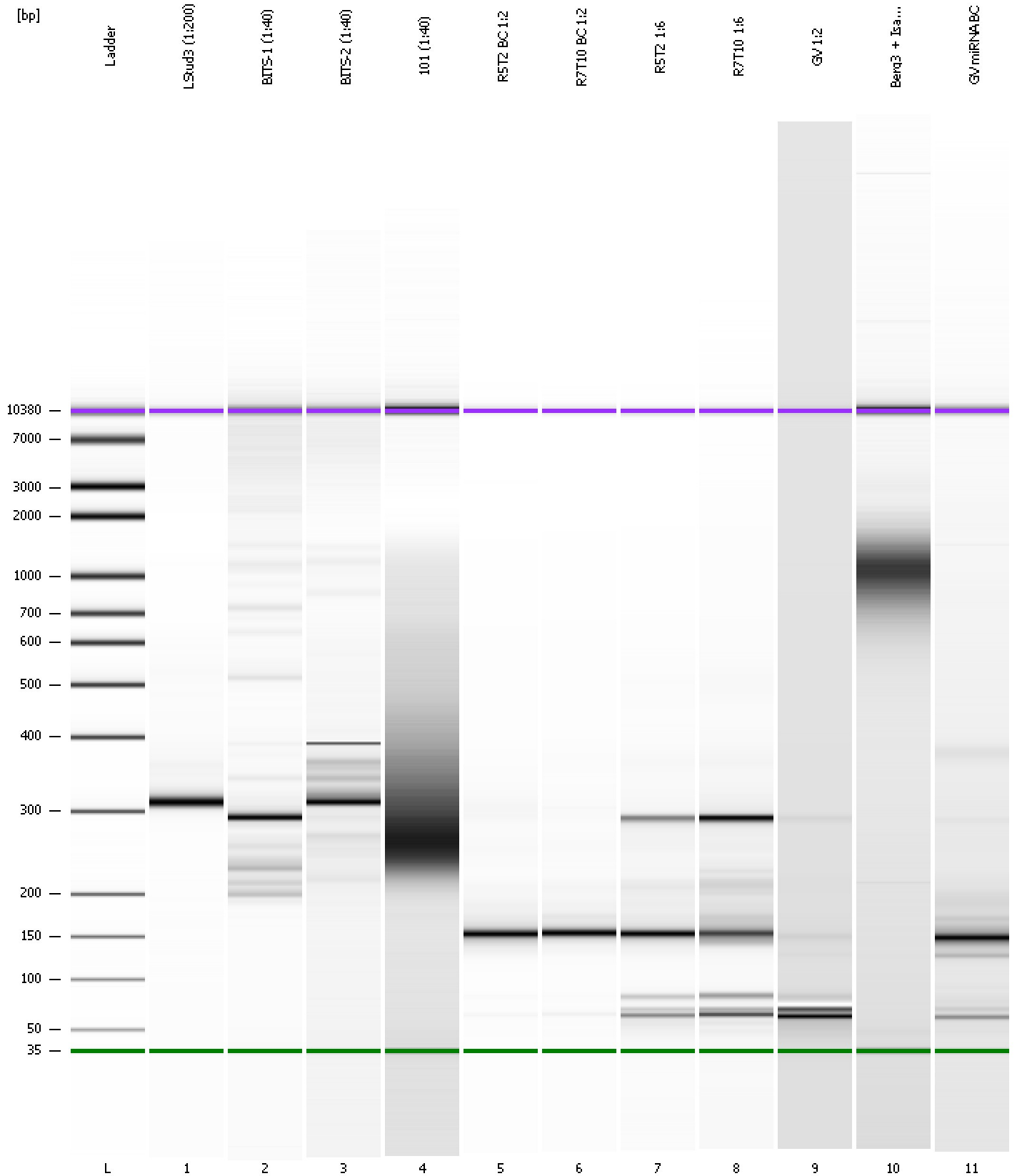
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
47	1,000	162	336.4	17,642.0	1,457.40	99	45.1



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
Modified: 5/3/2016 1:30:04 PM

**Gel Image**

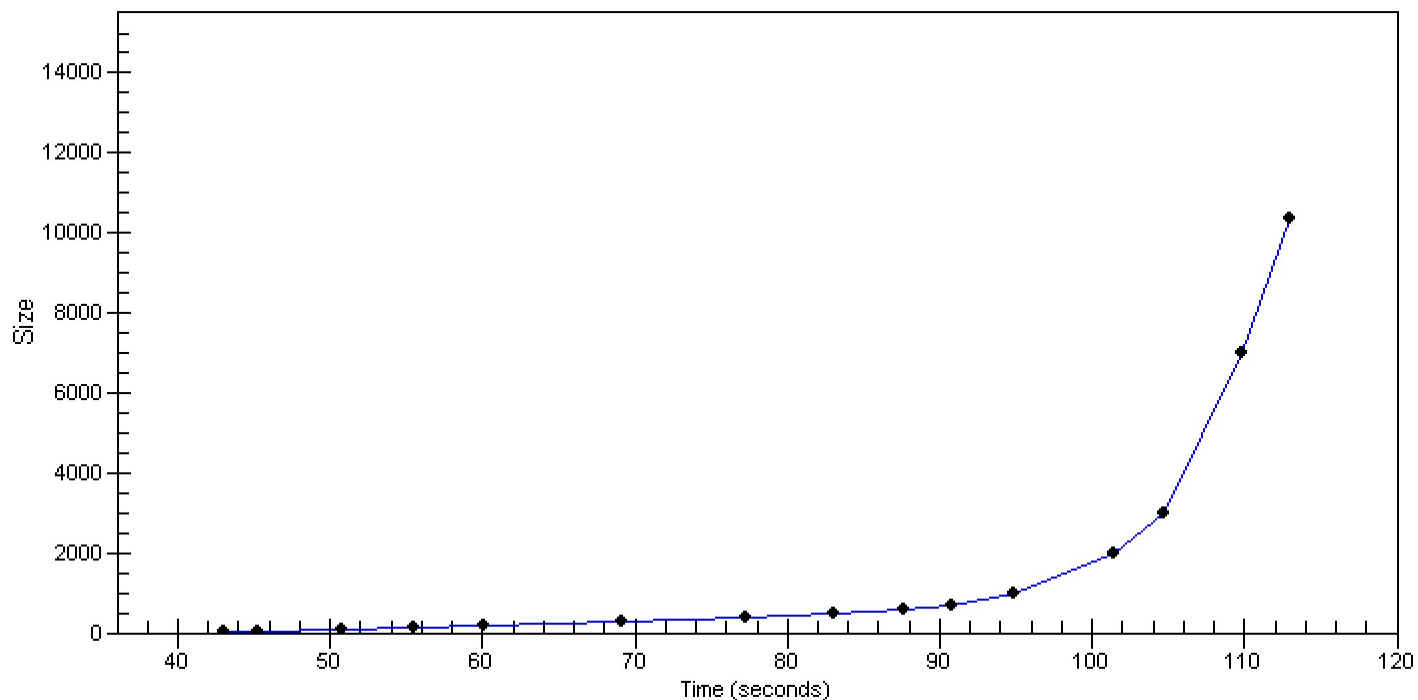


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
Modified: 5/3/2016 1:30:04 PM

**Curves**

**Standard Curve**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-03\2016-05-03\_002.xad

Created: 5/3/2016 12:39:01 PM  
 Modified: 5/3/2016 1:30:04 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		5/3/2016 1:20:20 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\2016-05-03\2016-05-03_002.xad)		Instrument	Run		5/3/2016 12:39:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		5/3/2016 12:39:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		5/3/2016 12:39:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		5/3/2016 12:39:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		5/3/2016 12:39:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		5/3/2016 12:39:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		5/3/2016 12:39:07 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1