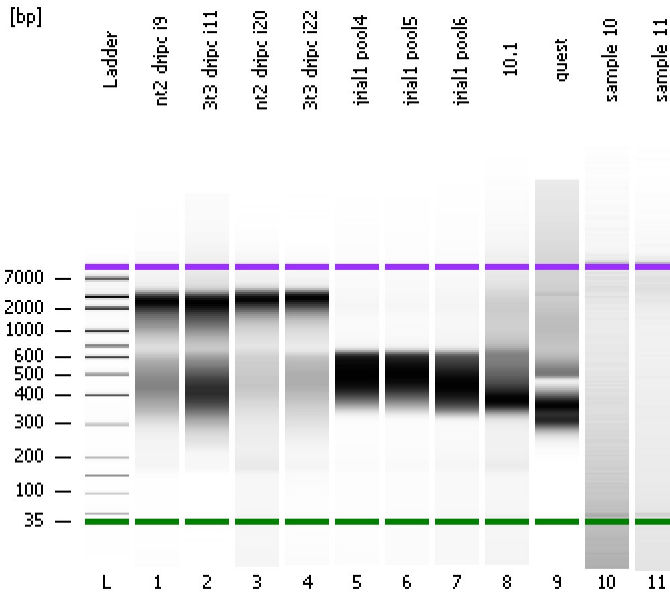


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
Modified: 11/10/2015 3:51:43 PM

### Electrophoresis File Run Summary



#### Instrument Information:

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

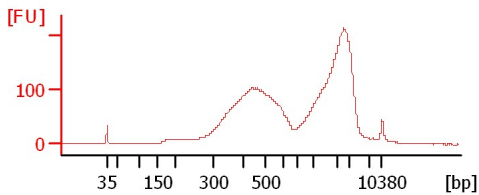
#### Assay Information:

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

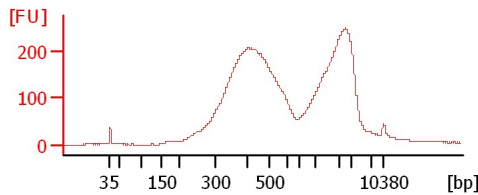
#### Chip Information:

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

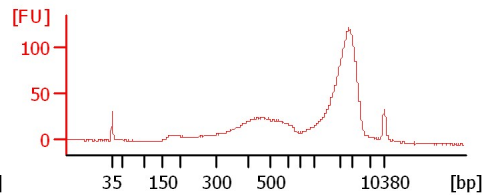
nt2 dripc i9



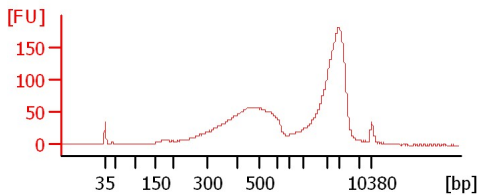
3t3 dripc i11



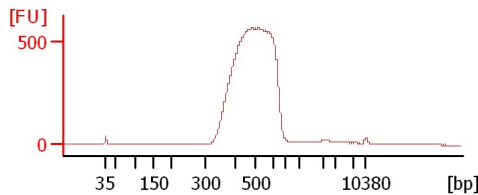
nt2 dripc i20



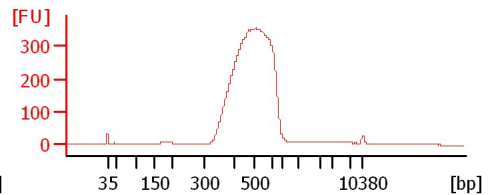
3t3 dripc i22



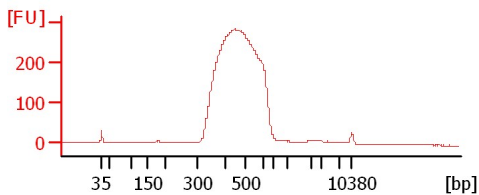
jrial1 pool4



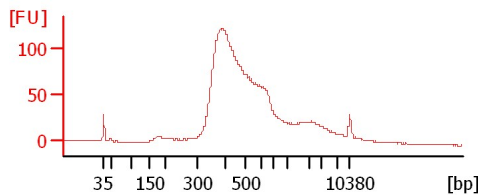
jrial1 pool5



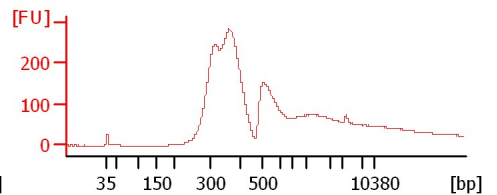
jrial1 pool6



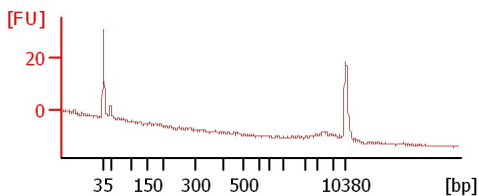
10.1



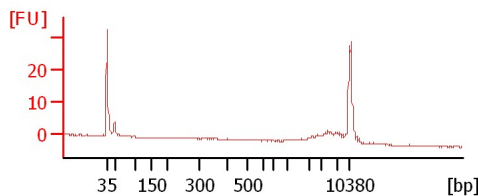
quest



sample 10



sample 11



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
 Modified: 11/10/2015 3:51:43 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
nt2 dripc i9		<input type="checkbox"/>	✓			
3t3 dripc i11		<input type="checkbox"/>	✓			
nt2 dripc i20		<input type="checkbox"/>	✓			
3t3 dripc i22		<input type="checkbox"/>	✓			
jrial1 pool4		<input type="checkbox"/>	✓			
jrial1 pool5		<input type="checkbox"/>	✓			
jrial1 pool6		<input type="checkbox"/>	✓			
10.1		<input type="checkbox"/>	✓			
guest		<input type="checkbox"/>	✓			
sample 10		<input type="checkbox"/>	✓			
sample 11		<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\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
Modified: 11/10/2015 3:51:43 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

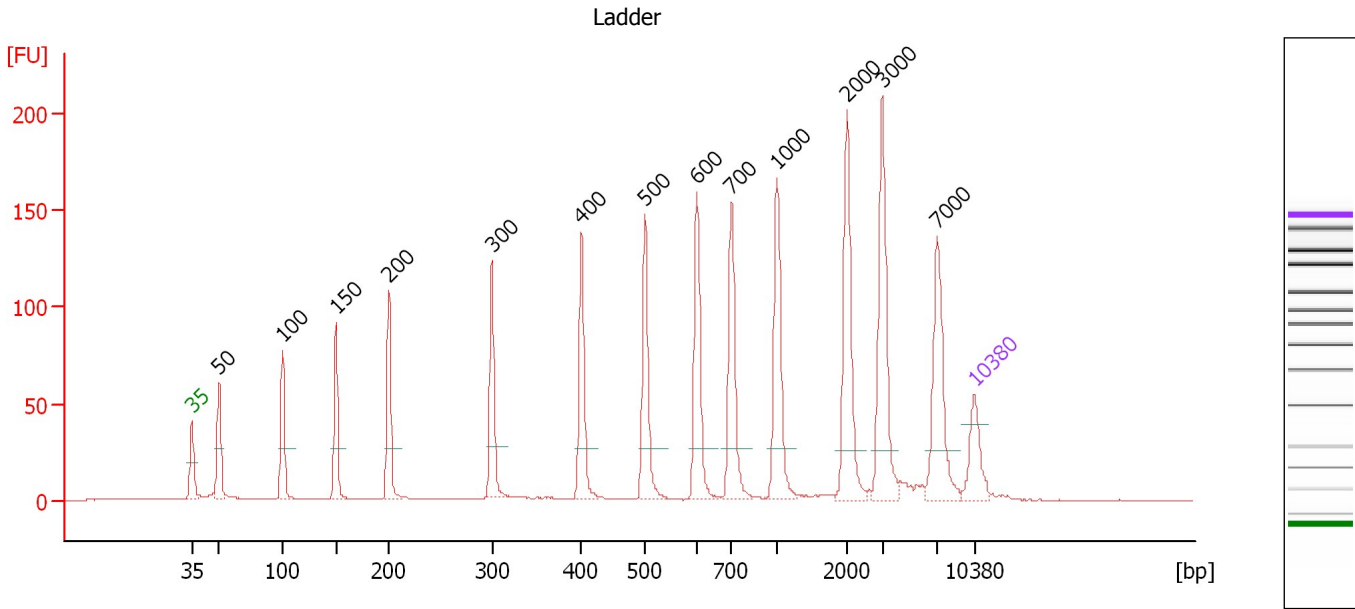
**Ladder**

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\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
 Modified: 11/10/2015 3:51:43 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.1

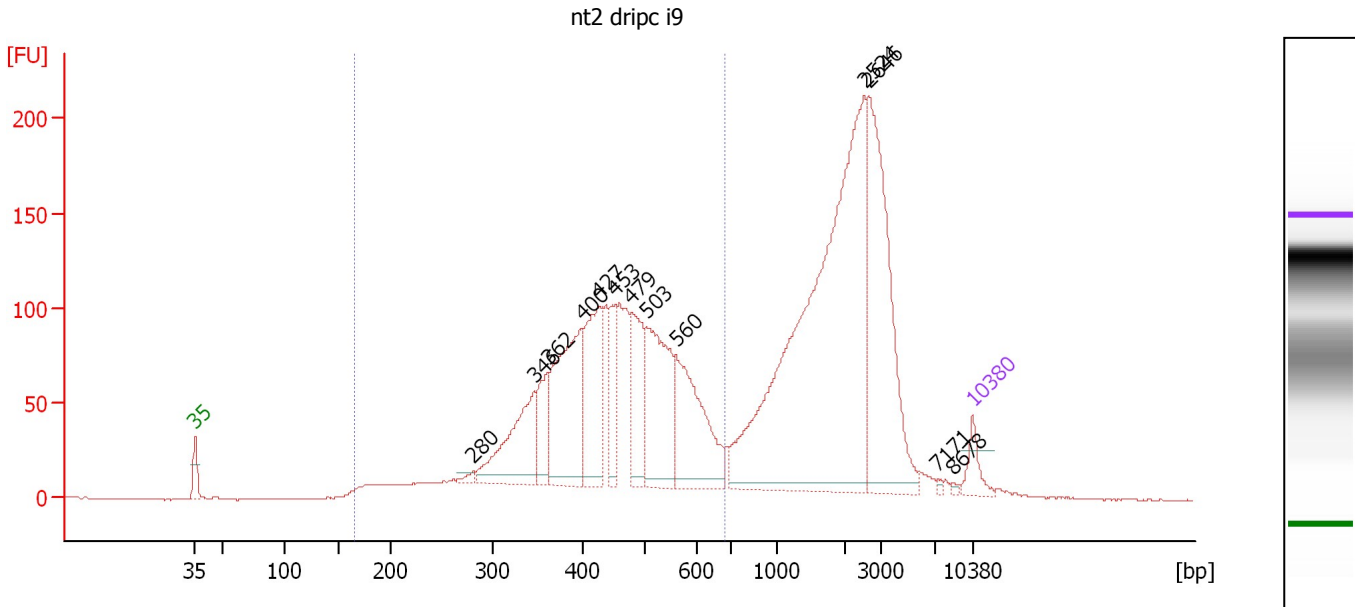
**Peak table for Ladder**

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	45.43
3	100	150.00	2,272.7	Ladder Peak	51.11
4	150	150.00	1,515.2	Ladder Peak	55.92
5	200	150.00	1,136.4	Ladder Peak	60.64
6	300	150.00	757.6	Ladder Peak	69.85
7	400	150.00	568.2	Ladder Peak	77.86
8	500	150.00	454.5	Ladder Peak	83.53
9	600	150.00	378.8	Ladder Peak	88.16
10	700	150.00	324.7	Ladder Peak	91.26
11	1,000	150.00	227.3	Ladder Peak	95.31
12	2,000	150.00	113.6	Ladder Peak	101.60
13	3,000	150.00	75.8	Ladder Peak	104.75
14	7,000	150.00	32.5	Ladder Peak	109.66
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
 Modified: 11/10/2015 3:51:43 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : nt2 dripc i9**

Number of peaks found: 13                      Corr. Area 1: 2,014.0  
 Noise: 0.2

**Peak table for sample 1 : nt2 dripc i9**

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	280	28.53	154.3		68.01
3	346	510.33	2,234.4		73.53
4	362	221.89	929.7		74.78
5	400	722.88	2,738.3		77.86
6	427	549.83	1,950.7		79.39
7	453	190.60	637.1		80.88
8	479	373.46	1,182.1		82.32
9	503	609.31	1,835.8		83.67
10	560	584.03	1,580.2		86.31
11	2,524	2,243.08	1,346.3		103.25
12	2,646	896.80	513.4		103.64
13	7,171	8.99	1.9		109.83
14	8,678	8.53	1.5		111.32
15	10,380	75.00	10.9	Upper Marker	113.00

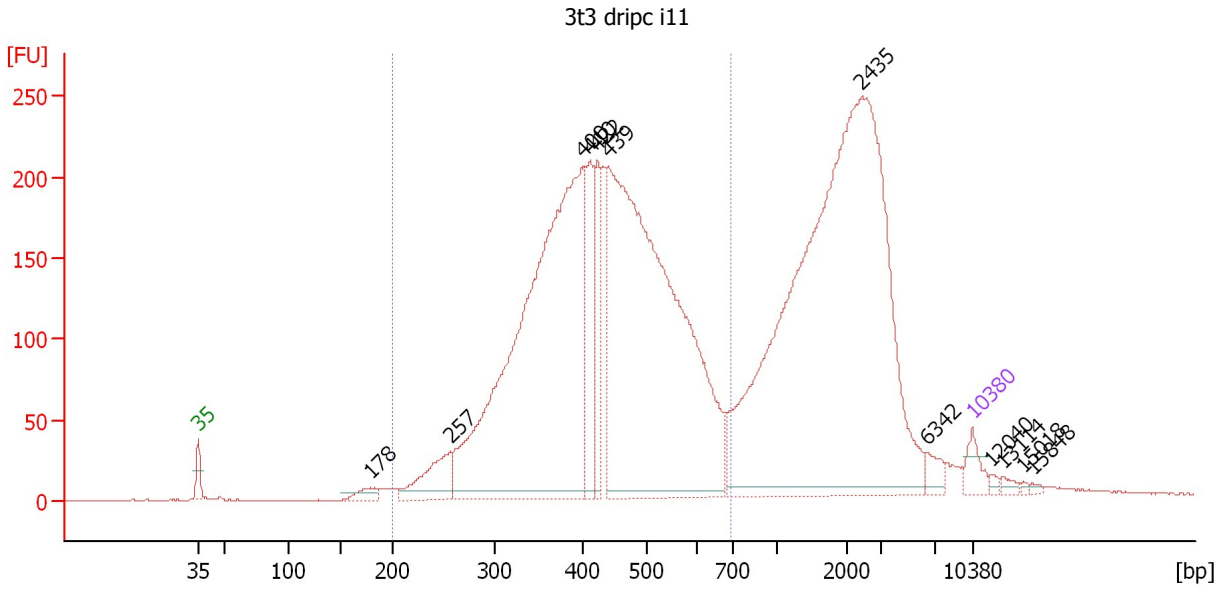
**Region table for sample 1 : nt2 dripc i9**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
166	681	442	2,014.0	20,056.0	5,290.01	52	23.1

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

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 Modified: 11/10/2015 3:51:43 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : 3t3 dripc i11**

Number of peaks found: 12                      Corr. Area 1: 4,167.3  
 Noise: 0.2

**Peak table for sample 2 : 3t3 dripc i11**

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	178	79.00	672.2		58.57
3	257	327.78	1,935.5		65.85
4	400	3,650.62	13,828.8		77.86
5	410	459.57	1,697.6		78.43
6	422	347.60	1,247.8		79.11
7	439	3,837.10	13,241.6		80.07
8	2,435	3,685.24	2,293.1		102.97
9	6,342	62.03	14.8		108.85
10	10,380	75.00	10.9	Upper Marker	113.00
11	12,040	0.00	0.0		114.64
12	13,114	0.00	0.0		115.70
13	15,018	0.00	0.0		117.58
14	15,848	0.00	0.0		118.40

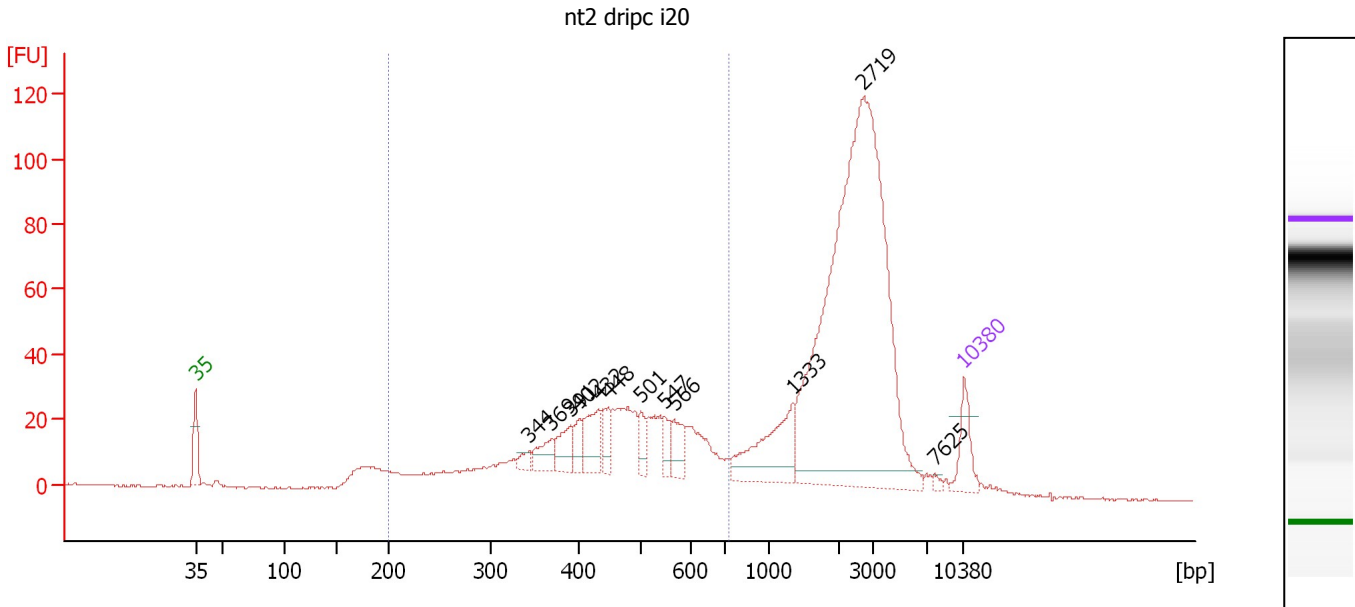
**Region table for sample 2 : 3t3 dripc i11**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	697	433	4,167.3	34,516.9	9,102.02	62	22.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
 Modified: 11/10/2015 3:51:43 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : nt2 dripc i20**

Number of peaks found: 12                      Corr. Area 1: 557.5  
 Noise: 0.3

**Peak table for sample 3 : nt2 dripc i20**

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	344	28.92	127.2		73.41
3	369	66.51	272.8		75.40
4	391	85.63	331.7		77.15
5	402	51.89	195.5		77.98
6	432	117.30	411.4		79.68
7	448	64.01	216.3		80.60
8	501	47.81	144.7		83.56
9	547	42.36	117.4		85.70
10	566	69.05	184.9		86.57
11	1,333	207.24	235.5		97.41
12	2,719	1,602.70	893.0		103.87
13	7,625	8.18	1.6		110.28
14	10,380	75.00	10.9	Upper Marker	113.00

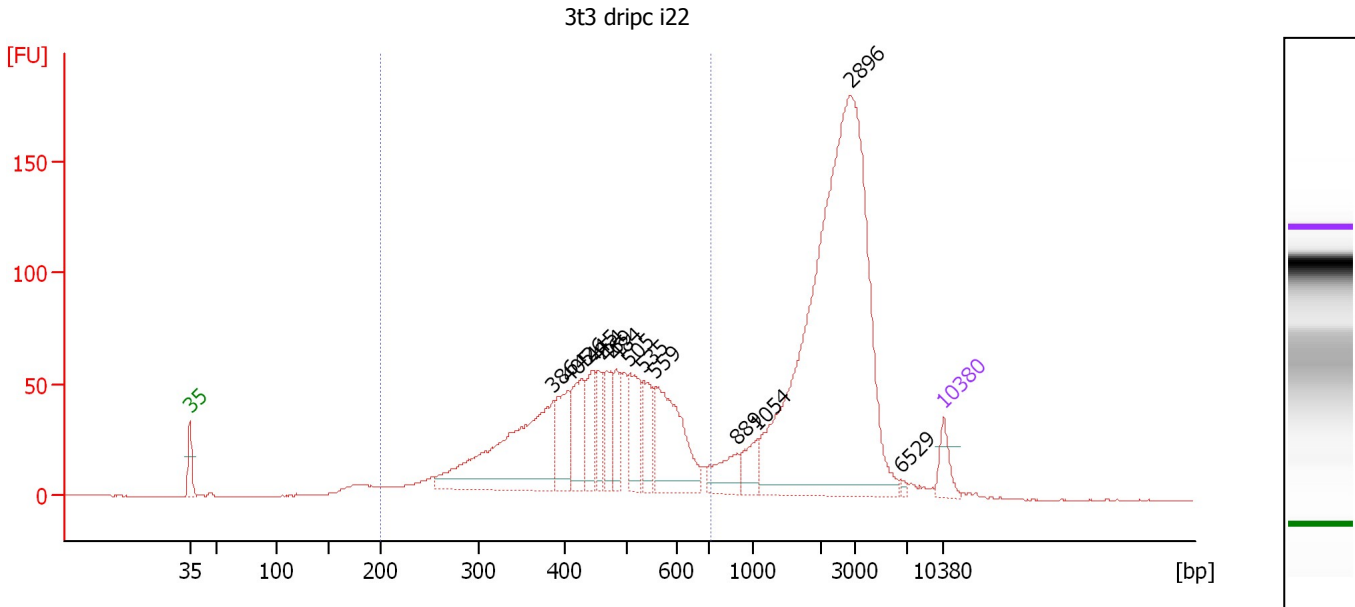
**Region table for sample 3 : nt2 dripc i20**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	723	453	557.5	6,731.5	1,787.11	38	25.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
 Modified: 11/10/2015 3:51:43 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : 3t3 dripc i22**

Number of peaks found: 14                      Corr. Area 1: 1,194.0  
 Noise: 0.2

**Peak table for sample 4 : 3t3 dripc i22**

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	386	911.85	3,576.3		76.76
3	405	240.77	900.4		78.15
4	426	247.88	881.4		79.34
5	445	196.35	668.0		80.43
6	454	140.26	468.0		80.92
7	469	158.08	510.8		81.77
8	484	137.02	429.1		82.61
9	505	230.90	693.1		83.75
10	535	183.32	519.4		85.14
11	559	458.56	1,242.0		86.28
12	889	127.66	217.5		93.81
13	1,054	95.74	137.6		95.65
14	2,896	2,376.46	1,243.3		104.42
15	6,529	9.63	2.2		109.08
16	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 4 : 3t3 dripc i22**

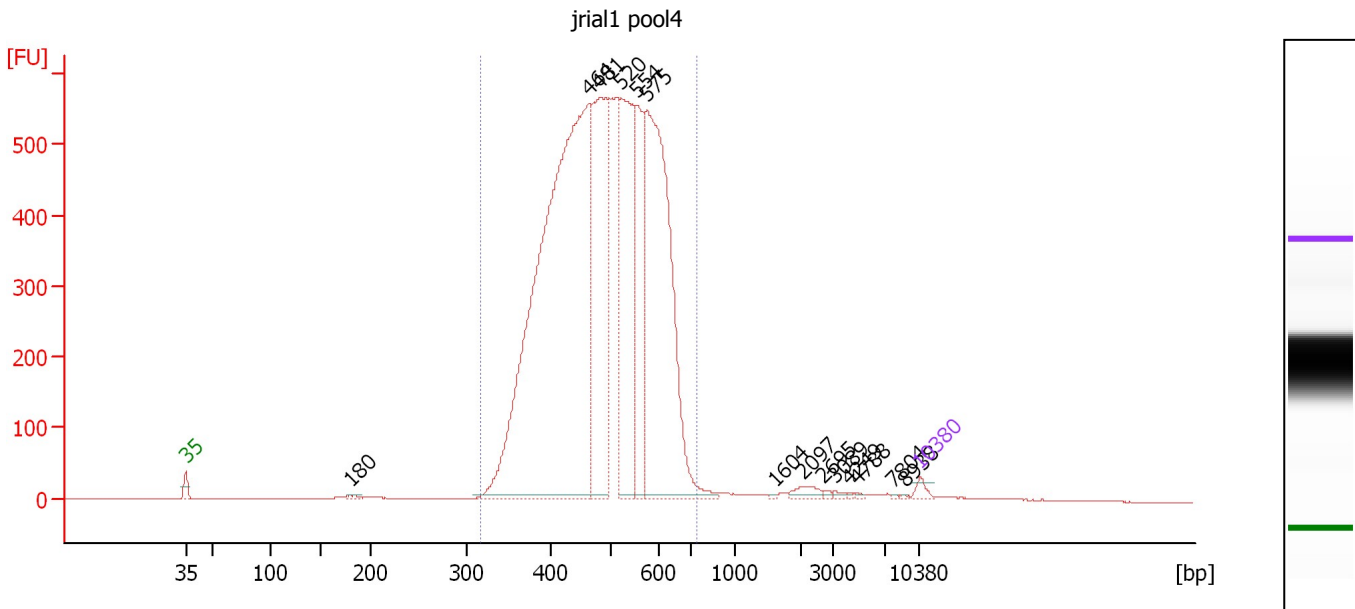
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	713	447	1,194.0	13,797.8	3,707.16	49	23.7



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
 Modified: 11/10/2015 3:51:43 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : jrial1 pool4**

Number of peaks found: 14                      Corr. Area 1: 9,086.2  
 Noise: 0.2

**Peak table for sample 5 : jrial1 pool4**

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	180	52.06	439.1		58.72
3	464	13,241.21	43,202.5		81.51
4	481	3,825.79	12,040.9		82.48
5	520	3,301.47	9,617.9		84.46
6	554	1,914.71	5,234.8		86.04
7	575	5,844.97	15,399.4		87.00
8	1,604	12.81	12.1		99.11
9	2,097	122.46	88.5		101.91
10	2,695	26.46	14.9		103.79
11	3,089	31.38	15.4		104.86
12	4,249	14.36	5.1		106.28
13	4,788	17.03	5.4		106.95
14	7,804	11.59	2.2		110.46
15	8,938	8.53	1.4		111.58
16	10,380	75.00	10.9	Upper Marker	113.00

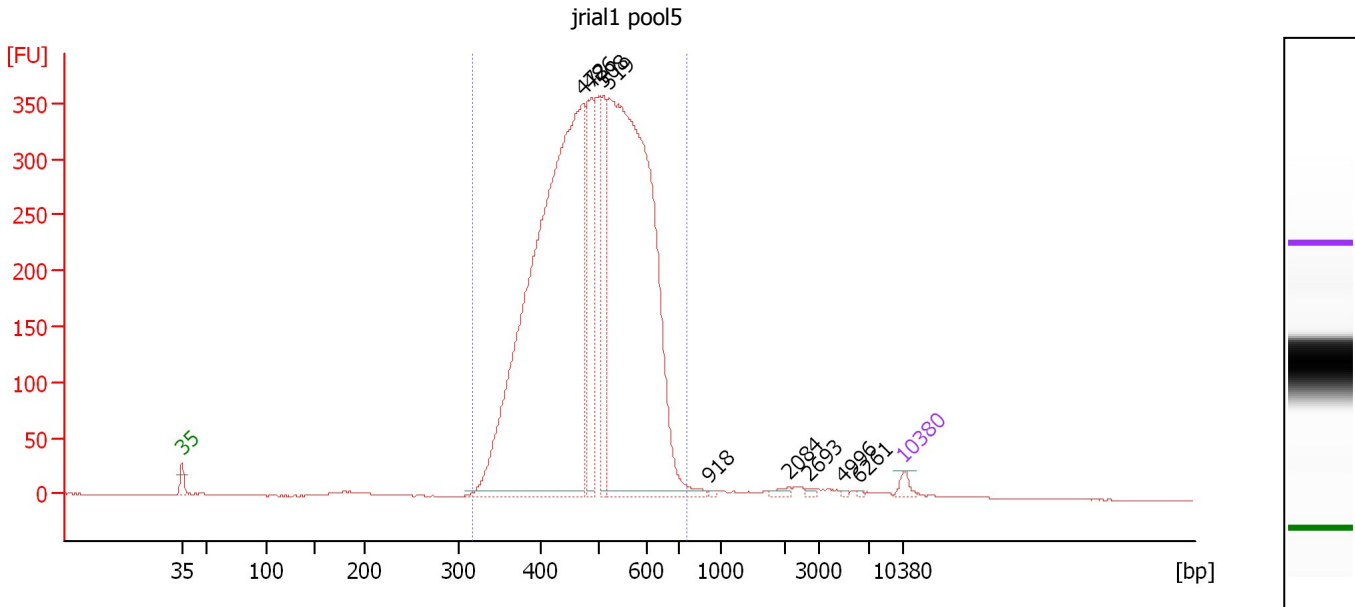
**Region table for sample 5 : jrial1 pool4**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
316	732	492	9,086.2	99,710.0	31,178.88	96	16.7

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
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**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : jrial1 pool5**

Number of peaks found: 9                      Corr. Area 1: 5,466.4  
 Noise: 0.2

**Peak table for sample 6 : jrial1 pool5**

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	472	11,038.80	35,445.3		81.93
3	486	1,475.65	4,604.7		82.71
4	508	1,259.79	3,757.3		83.90
5	519	9,164.43	26,743.2		84.42
6	918	17.26	28.5		94.21
7	2,084	50.27	36.5		101.87
8	2,693	24.97	14.1		103.78
9	4,996	12.22	3.7		107.20
10	6,261	14.59	3.5		108.75
11	10,380	75.00	10.9	Upper Marker	113.00

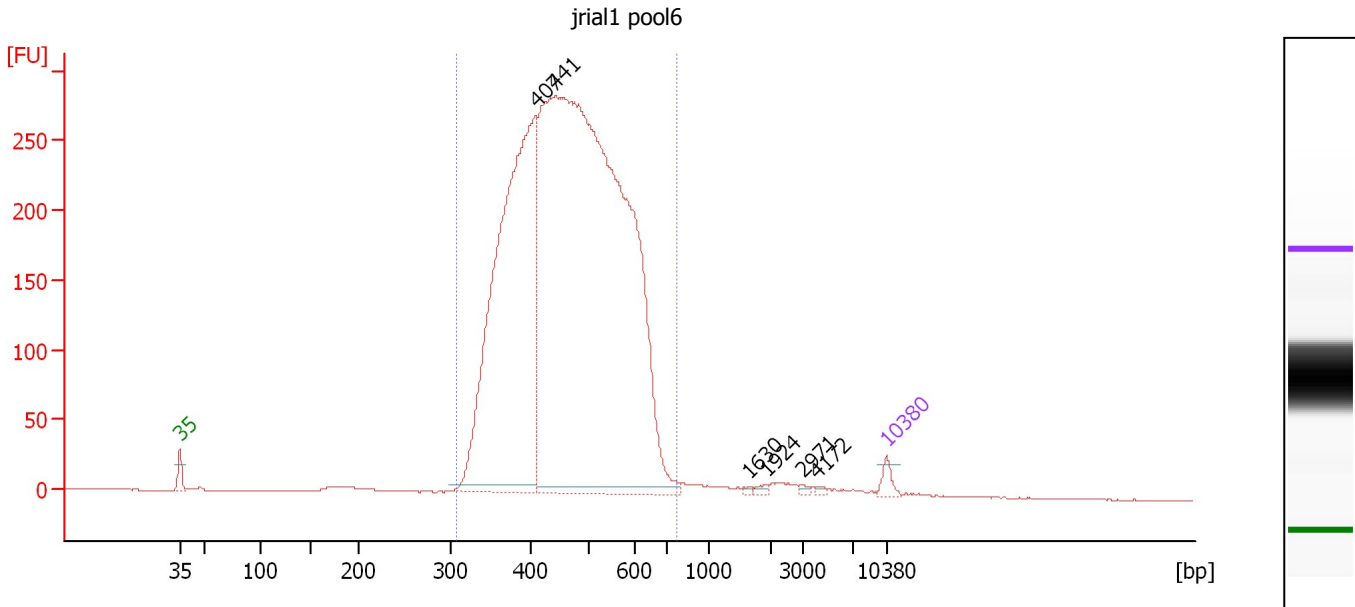
**Region table for sample 6 : jrial1 pool5**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
317	756	492	5,466.4	77,664.0	24,305.10	97	16.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
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**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : jrial1 pool6**

Number of peaks found: 6                      Corr. Area 1: 4,864.8  
 Noise: 0.1

**Peak table for sample 7 : jrial1 pool6**

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	407	6,039.58	22,474.2		78.26
3	441	13,488.79	46,378.9		80.16
4	1,630	15.89	14.8		99.27
5	1,924	29.77	23.4		101.12
6	2,971	22.70	11.6		104.66
7	4,172	20.15	7.3		106.19
8	10,380	75.00	10.9	Upper Marker	113.00

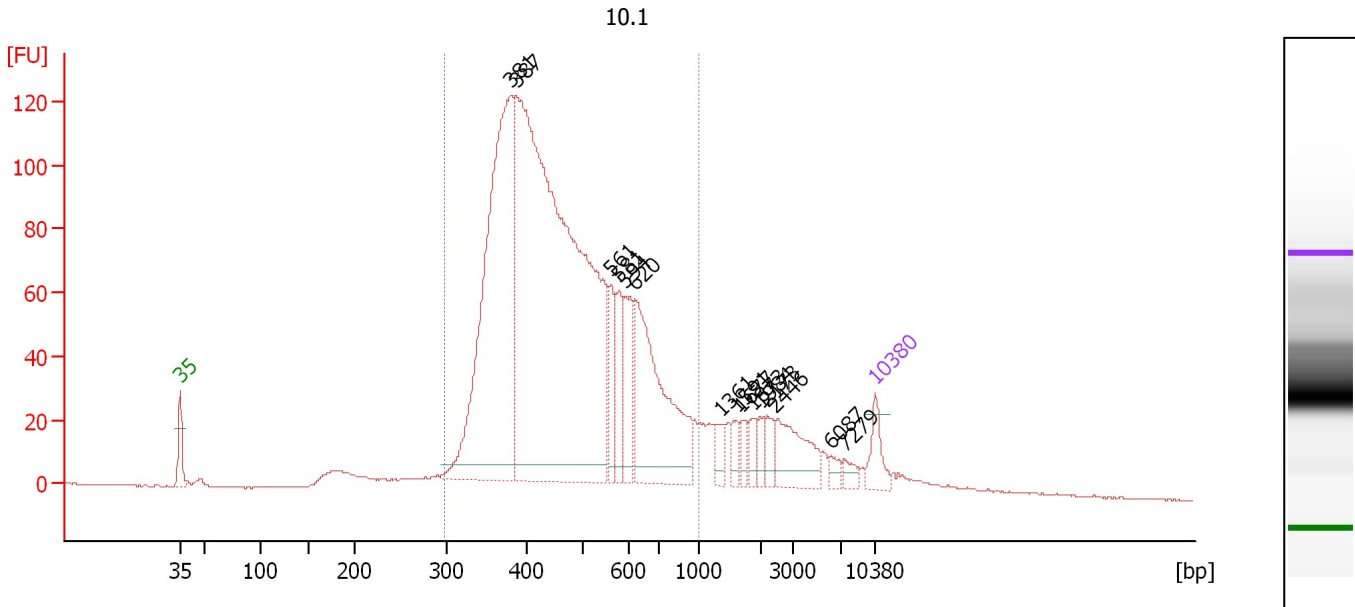
**Region table for sample 7 : jrial1 pool6**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
308	769	469	4,864.8	64,521.2	19,112.85	96	18.2

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
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**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : 10.1**

Number of peaks found: 15                      Corr. Area 1: 1,984.9  
 Noise: 0.2

**Peak table for sample 8 : 10.1**

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	381	1,754.15	6,982.1		76.31
3	387	3,629.38	14,196.0		76.84
4	561	154.77	418.0		86.35
5	581	169.18	441.4		87.27
6	594	194.80	497.3		87.86
7	620	698.38	1,707.2		88.77
8	1,361	51.12	56.9		97.58
9	1,591	45.69	43.5		99.03
10	1,677	37.26	33.7		99.57
11	1,873	51.15	41.4		100.81
12	1,984	36.95	28.2		101.50
13	2,173	50.32	35.1		102.15
14	2,446	186.92	115.8		103.01
15	6,087	29.94	7.5		108.54
16	7,279	32.58	6.8		109.94
17	10,380	75.00	10.9	Upper Marker	113.00

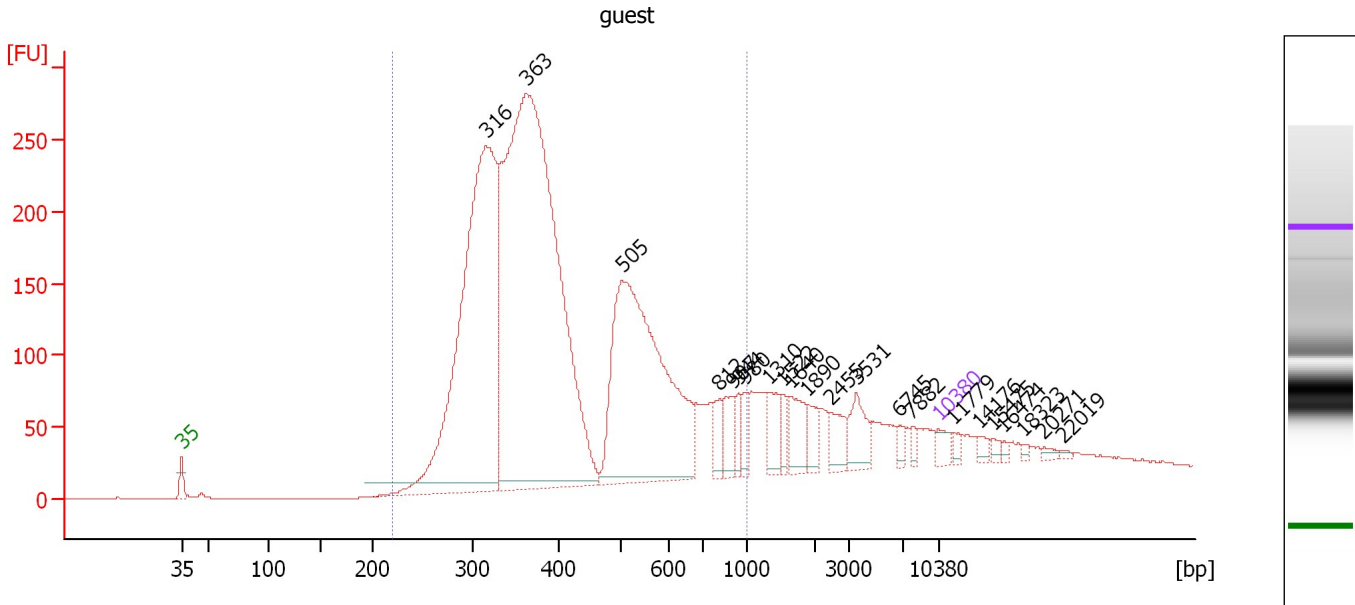
**Region table for sample 8 : 10.1**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
297	1,000	483	1,984.9	23,159.6	6,811.85	82	27.4

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : quest**

Number of peaks found: 22                      Corr. Area 1: 4,579.0  
 Noise: 0.1

**Peak table for sample 9 : quest**

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	316	4,915.41	23,531.2		71.17
3	363	7,396.71	30,900.6		74.87
4	505	3,116.70	9,357.3		83.75
5	812	156.19	291.3		92.77
6	907	187.30	312.8		94.06
7	944	111.28	178.6		94.55
8	980	110.47	170.7		95.04
9	1,310	185.82	214.8		97.26
10	1,522	100.72	100.3		98.60
11	1,640	230.63	213.1		99.34
12	1,890	130.37	104.5		100.91
13	2,455	149.17	92.1		103.04
14	3,531	211.72	90.8		105.40
15	6,745	49.93	11.2		109.35
16	7,882	40.13	7.7		110.53
17	10,380	75.00	10.9	Upper Marker	113.00
18	11,779	0.00	0.0		114.38
19	14,176	0.00	0.0		116.75
20	15,475	0.00	0.0		118.03
21	16,474	0.00	0.0		119.02
22	18,323	0.00	0.0		120.84
23	20,271	0.00	0.0		122.77
24	22,019	0.00	0.0		124.49

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
Modified: 11/10/2015 3:51:43 PM

**Electropherogram Summary Continued ...**

... Region table for sample 9 :

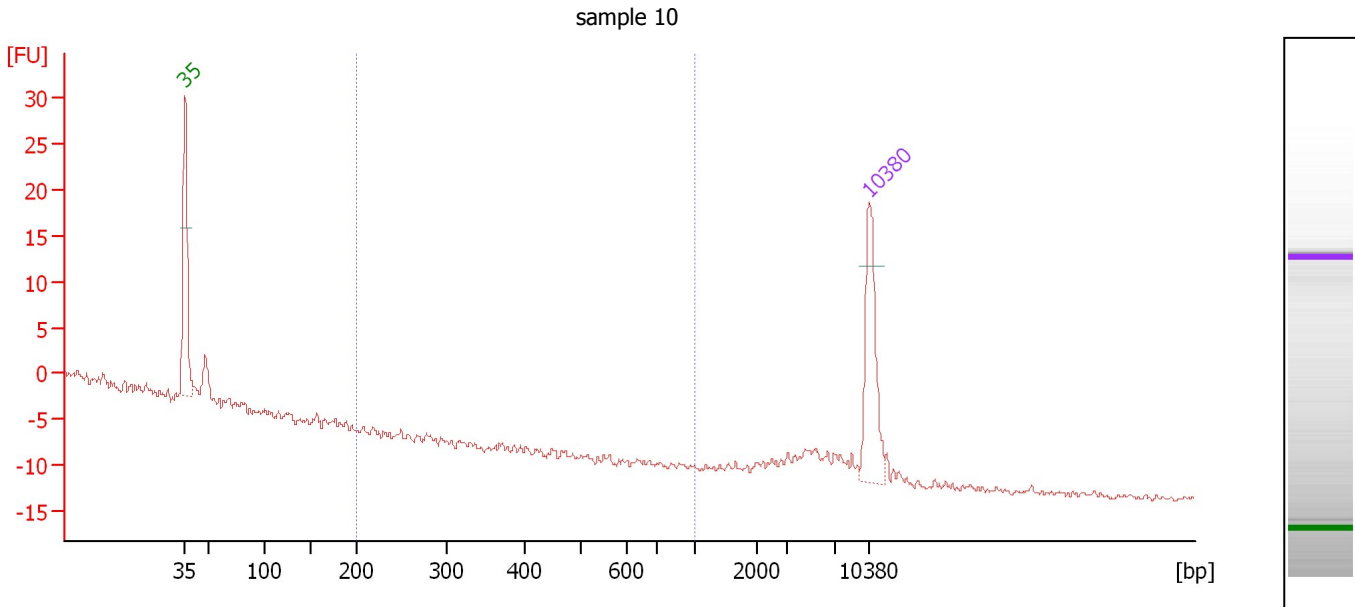
quest

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
221	1,000	430	4,579.0	62,667.1	15,771.91	82	34.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
 Modified: 11/10/2015 3:51:43 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : sample 10**

Number of peaks found: 0                      Corr. Area 1: 0.0  
 Noise: 0.4

**Peak table for sample 10 : sample 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	10,380	75.00	10.9	Upper Marker	113.00

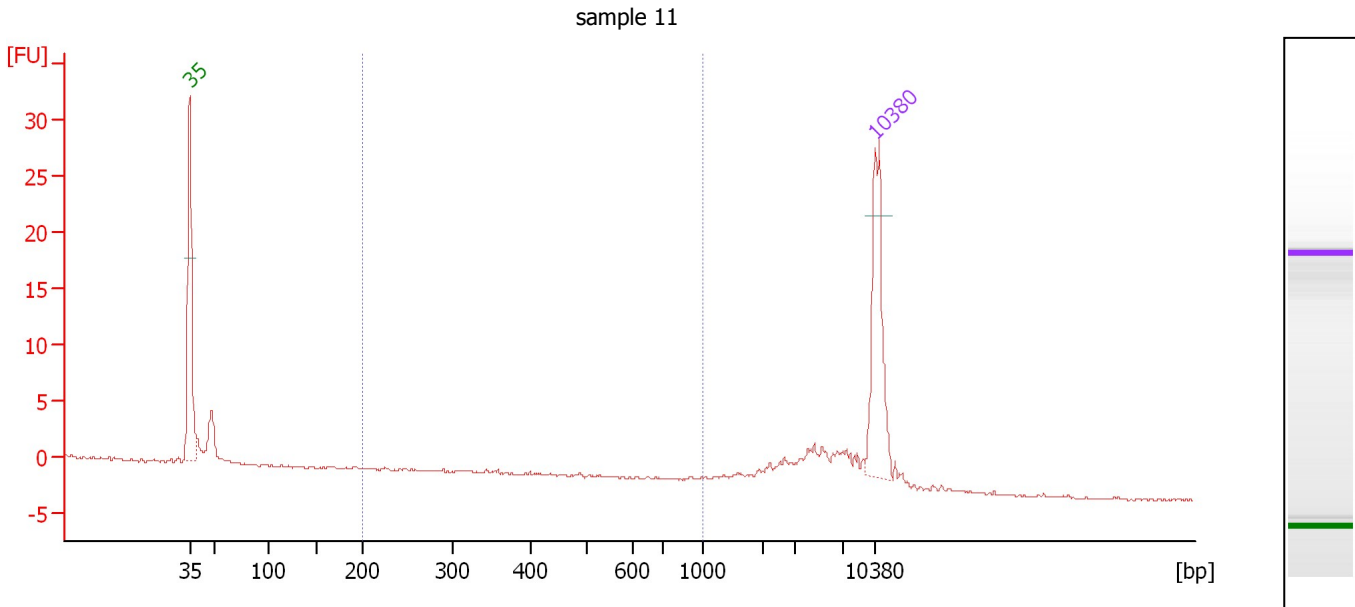
**Region table for sample 10 : sample 10**

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	0	0.0	0.0	0.00	0	0.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
 Modified: 11/10/2015 3:51:43 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 11 : sample 11**

Number of peaks found: 0                      Corr. Area 1: 3.7  
 Noise: 0.1

**Peak table for sample 11 : sample 11**

Pea k	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	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 11 : sample 11**

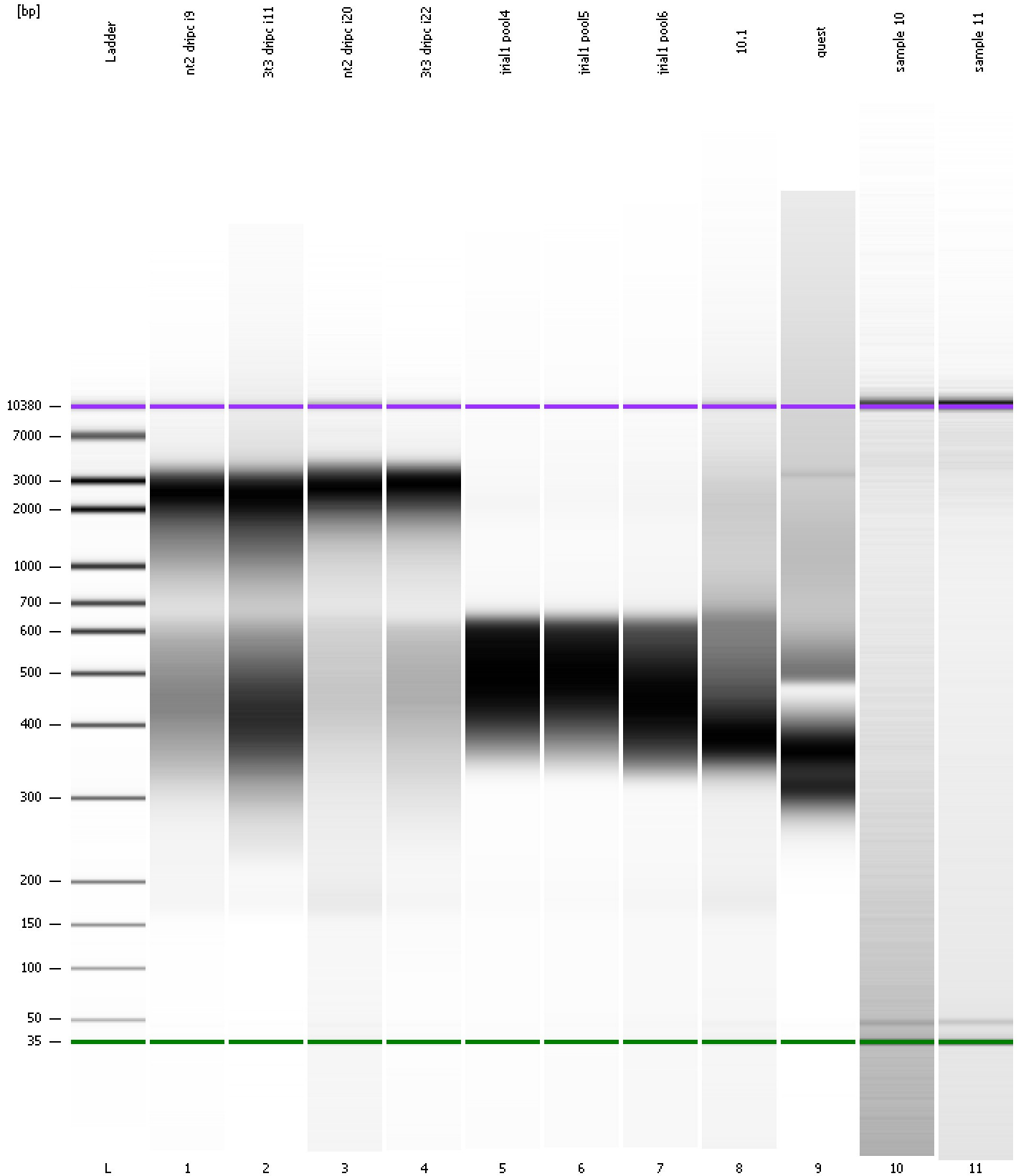
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	549	3.7	43.7	12.57	8	40.1



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
Modified: 11/10/2015 3:51:43 PM

**Gel Image**

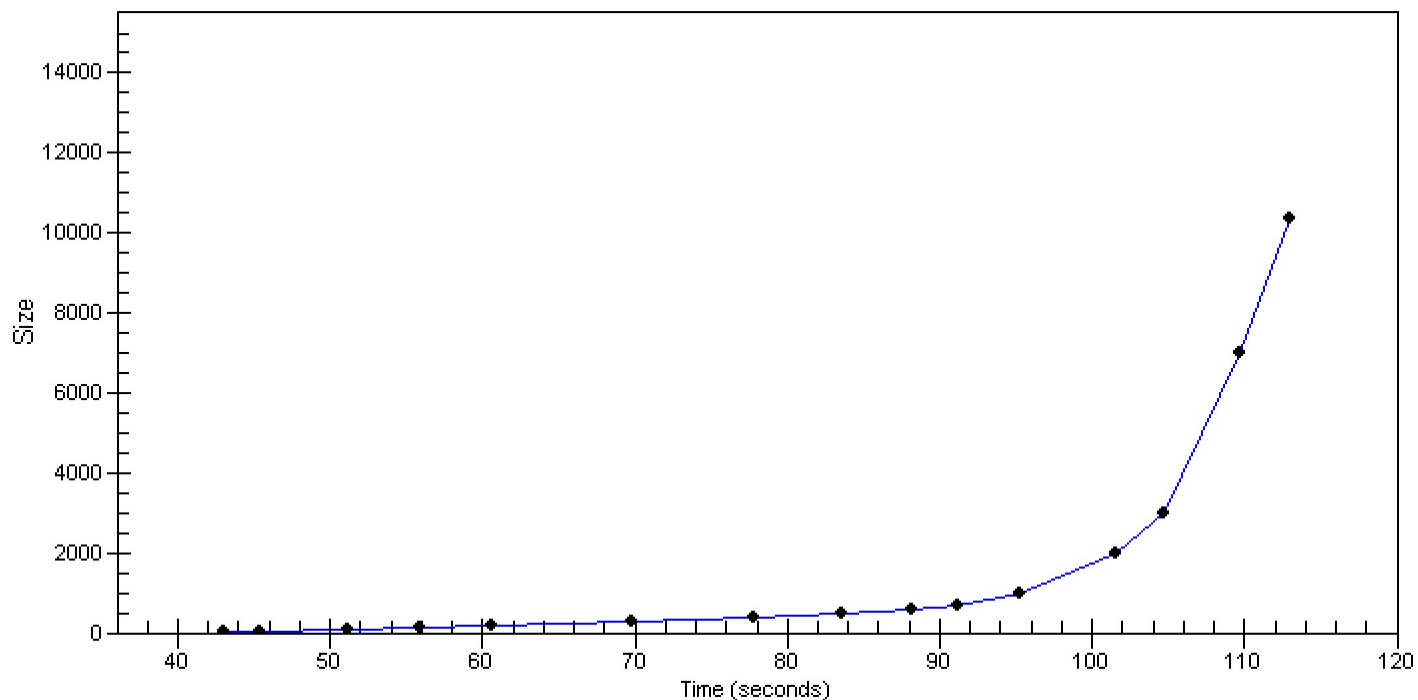


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
Modified: 11/10/2015 3:51:43 PM

**Curves**

**Standard Curve**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-11-10\2015-11-10\_003.xad

Created: 11/10/2015 3:10:26 PM  
 Modified: 11/10/2015 3:51:43 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		11/10/2015 3:51:42 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-11-10\2015-11-10_003.xad)		Instrument	Run		11/10/2015 3:10:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		11/10/2015 3:10:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		11/10/2015 3:10:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		11/10/2015 3:10:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		11/10/2015 3:10:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		11/10/2015 3:10:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		11/10/2015 3:10:31 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1