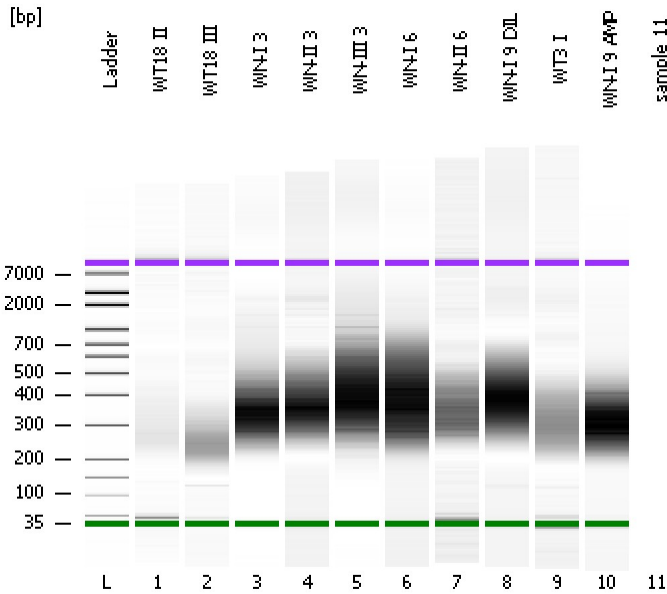


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
Modified: 7/29/2016 12:18:45 PM

**Electrophoresis File Run Summary**



Instrument Information:

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

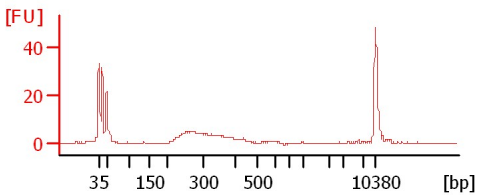
Assay Information:

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

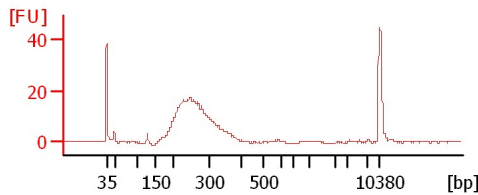
Chip Information:

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

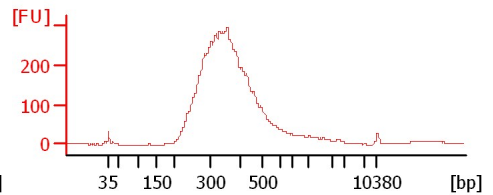
**WT18 II**



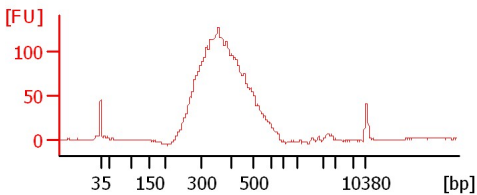
**WT18 III**



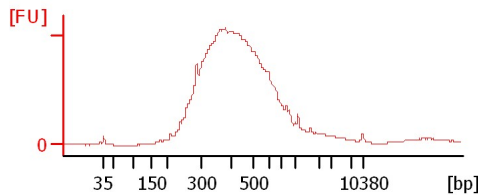
**WN-I 3**



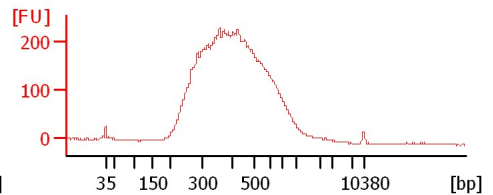
**WN-II 3**



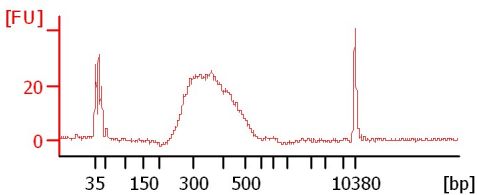
**WN-III 3**



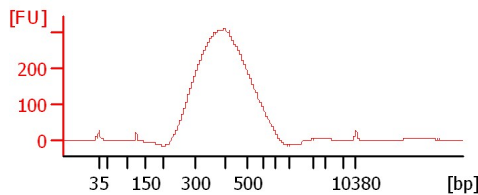
**WN-I 6**



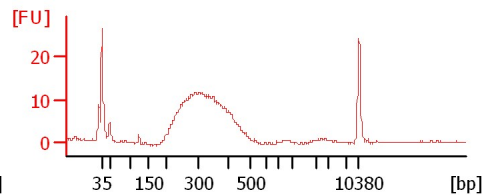
**WN-II 6**



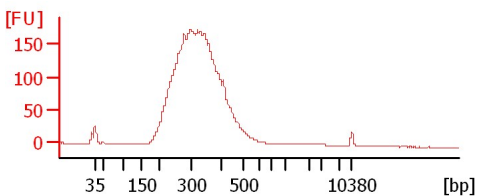
**WN-I 9 DIL**



**WN-I 9 AMP**



**WN-I 9 AMP**



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
Modified: 7/29/2016 12:18:45 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
WT18 II		<input type="checkbox"/>	✓			
WT18 III		<input type="checkbox"/>	✓			
WN-I 3		<input type="checkbox"/>	✓			
WN-II 3		<input type="checkbox"/>	✓			
WN-III 3		<input type="checkbox"/>	✓			
WN-I 6		<input type="checkbox"/>	✓			
WN-II 6		<input type="checkbox"/>	✓			
WN-I 9 DIL		<input type="checkbox"/>	✓			
WT3 I		<input type="checkbox"/>	✓			
WN-I 9 AMP		<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:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
Modified: 7/29/2016 12:18:45 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

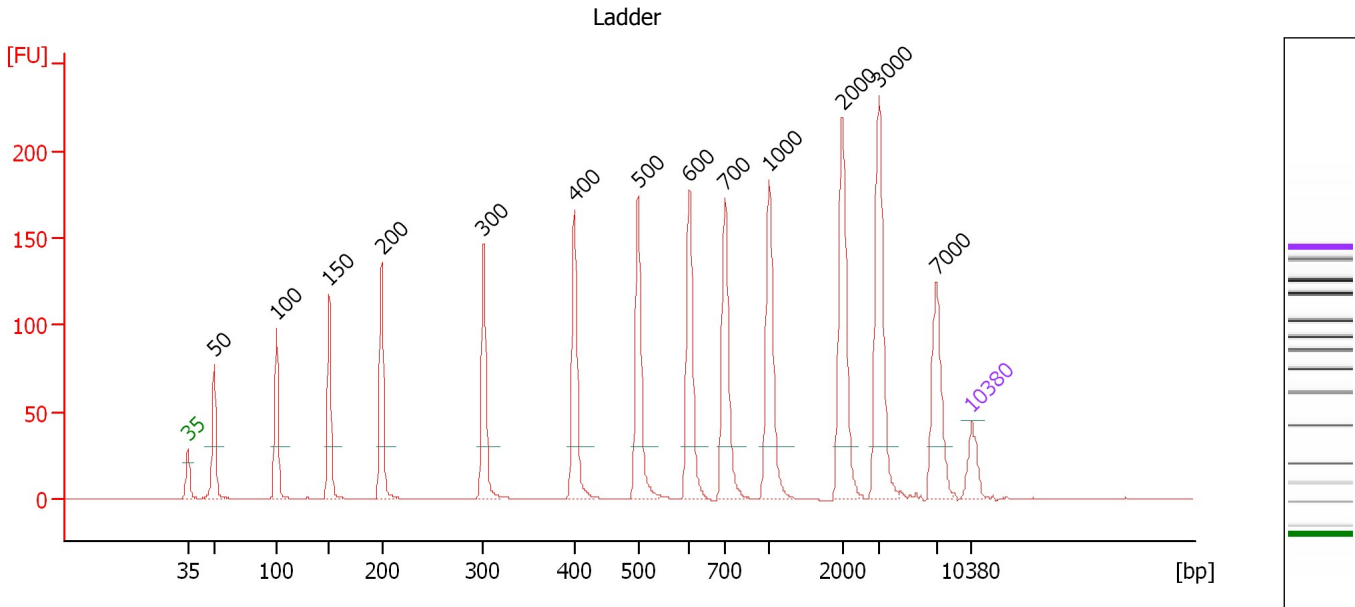
**Ladder**

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\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
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**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.2

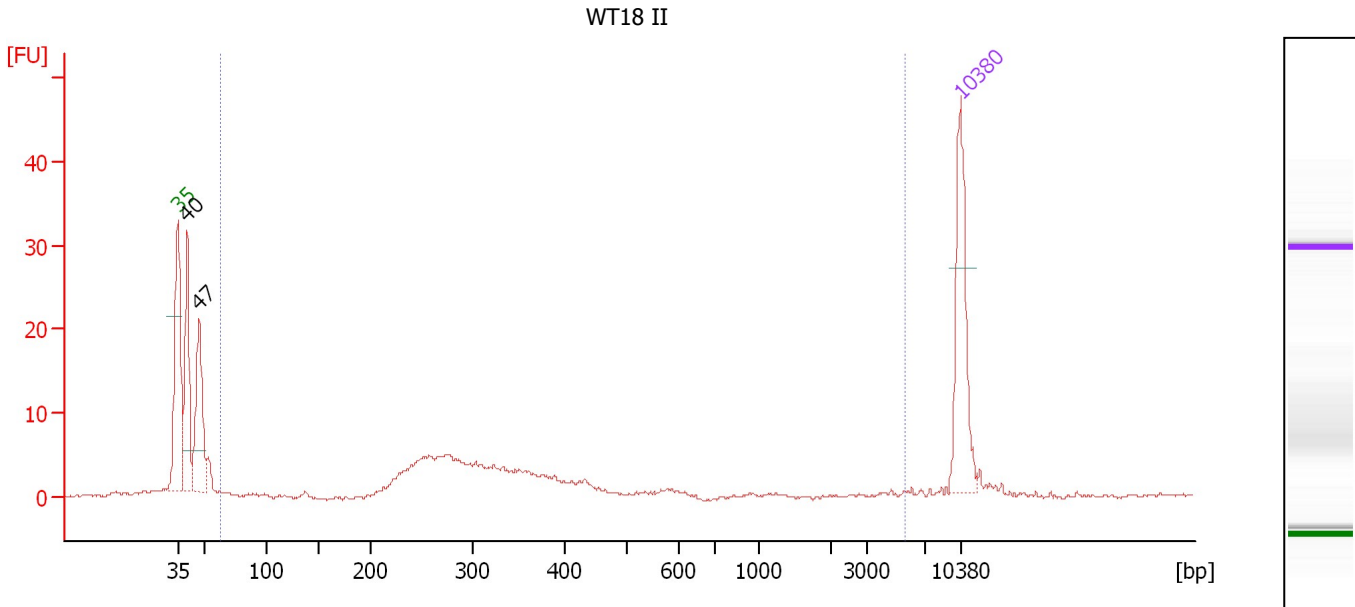
**Peak table for Ladder**

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	45.38
3	100	150.00	2,272.7	Ladder Peak	50.95
4	150	150.00	1,515.2	Ladder Peak	55.66
5	200	150.00	1,136.4	Ladder Peak	60.32
6	300	150.00	757.6	Ladder Peak	69.41
7	400	150.00	568.2	Ladder Peak	77.55
8	500	150.00	454.5	Ladder Peak	83.21
9	600	150.00	378.8	Ladder Peak	87.78
10	700	150.00	324.7	Ladder Peak	90.97
11	1,000	150.00	227.3	Ladder Peak	94.92
12	2,000	150.00	113.6	Ladder Peak	101.44
13	3,000	150.00	75.8	Ladder Peak	104.72
14	7,000	150.00	32.5	Ladder Peak	109.81
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
 Modified: 7/29/2016 12:18:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 1 : WT18 II**

Number of peaks found: 2                      Corr. Area 1: 95.8  
 Noise: 0.2

**Peak table for sample 1 : WT18 II**

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	40	123.22	4,654.3		43.81
3	47	119.15	3,838.4		44.91
4	10,380	75.00	10.9	Upper Marker	113.00

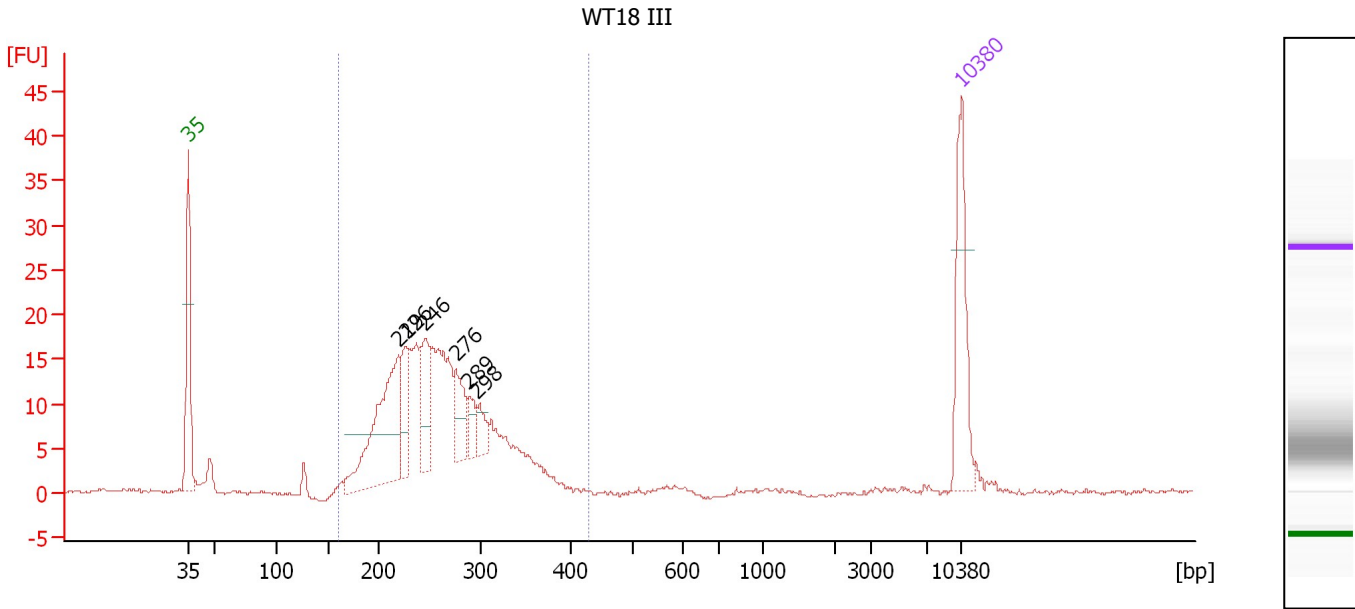
**Region table for sample 1 : WT18 II**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
63	5,645	368	95.8	1,454.1	261.49	52	100.0

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

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**Electropherogram Summary Continued ...**



**Overall Results for sample 2 : WT18 III**

Number of peaks found: 6                      Corr. Area 1: 271.9  
 Noise: 0.2

**Peak table for sample 2 : WT18 III**

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	219	191.78	1,329.5		62.01
3	226	51.17	343.1		62.68
4	246	69.78	430.5		64.47
5	276	49.49	271.8		67.22
6	289	22.39	117.3		68.42
7	298	20.92	106.5		69.20
8	10,380	75.00	10.9	Upper Marker	113.00

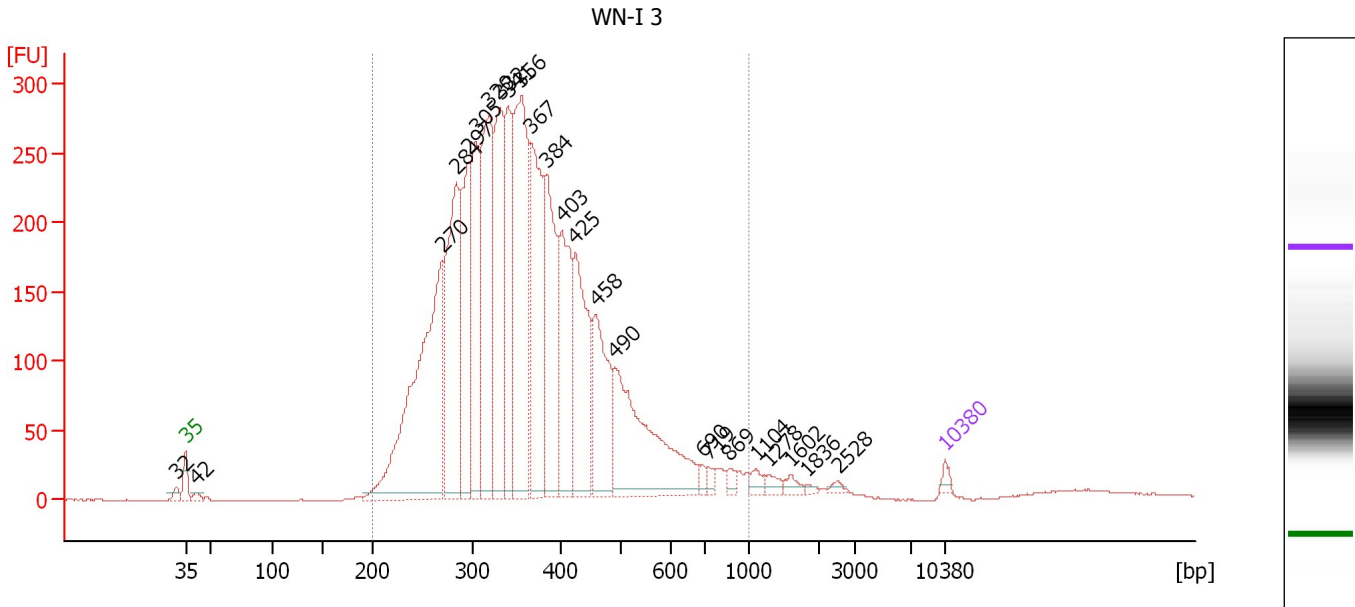
**Region table for sample 2 : WT18 III**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
160	428	260	271.9	5,159.3	853.12	94	18.2

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

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 Modified: 7/29/2016 12:18:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : WN-I 3**

Number of peaks found: 24                      Corr. Area 1: 5,879.1  
 Noise: 0.2

**Peak table for sample 3 : WN-I 3**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.16
2	35	125.00	5,411.3	Lower Marker	43.00
3	42	119.14	4,316.7		44.08
4	270	6,322.38	35,535.9		66.64
5	284	3,647.53	19,448.0		67.97
6	297	2,436.39	12,445.6		69.10
7	305	2,576.84	12,789.4		69.84
8	320	3,666.94	17,374.7		71.02
9	332	2,944.88	13,445.4		72.00
10	341	2,187.60	9,722.4		72.74
11	356	4,487.34	19,097.3		73.97
12	367	2,989.52	12,345.7		74.85
13	384	2,705.11	10,662.1		76.28
14	403	2,253.83	8,478.4		77.71
15	425	2,364.55	8,422.9		78.98
16	458	1,861.80	6,154.7		80.85
17	490	3,360.66	10,400.5		82.62
18	690	102.17	224.5		90.63
19	719	101.10	212.9		91.22
20	869	121.76	212.3		93.19
21	1,104	132.83	182.2		95.60
22	1,278	121.26	143.8		96.73
23	1,602	106.30	100.5		98.84
24	1,836	41.01	33.8		100.37
25	2,528	54.50	32.7		103.17

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
 Modified: 7/29/2016 12:18:45 PM

### Electropherogram Summary Continued ...

#### ... Peak table for sample 3 : WN-I 3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	10,380	75.00	10.9	Upper Marker	113.00

#### Region table for sample 3 : WN-I 3

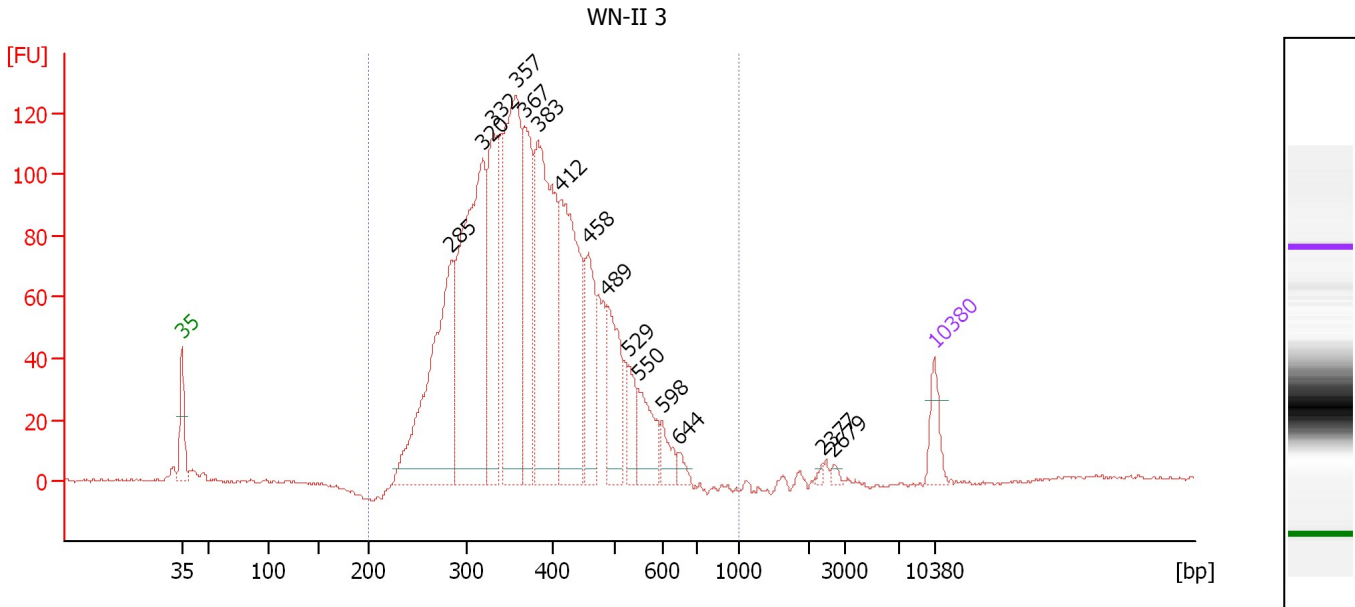
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	372	5,879.1	201,530.1	45,188.40	97	29.2



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
 Modified: 7/29/2016 12:18:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : WN-II 3**

Number of peaks found: 15                      Corr. Area 1: 2,298.5  
 Noise: 0.5

**Peak table for sample 4 : WN-II 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	285	973.96	5,173.2		68.07
3	320	1,246.72	5,911.7		71.00
4	332	522.63	2,387.1		71.99
5	357	942.90	3,997.8		74.08
6	367	493.01	2,038.1		74.82
7	383	932.98	3,691.1		76.16
8	412	754.57	2,772.5		78.25
9	458	297.74	985.1		80.83
10	489	302.56	938.2		82.57
11	529	115.15	329.6		84.55
12	550	175.61	483.7		85.50
13	598	68.73	174.2		87.68
14	644	27.58	64.9		89.17
15	2,377	8.15	5.2		102.67
16	2,679	10.97	6.2		103.67
17	10,380	75.00	10.9	Upper Marker	113.00

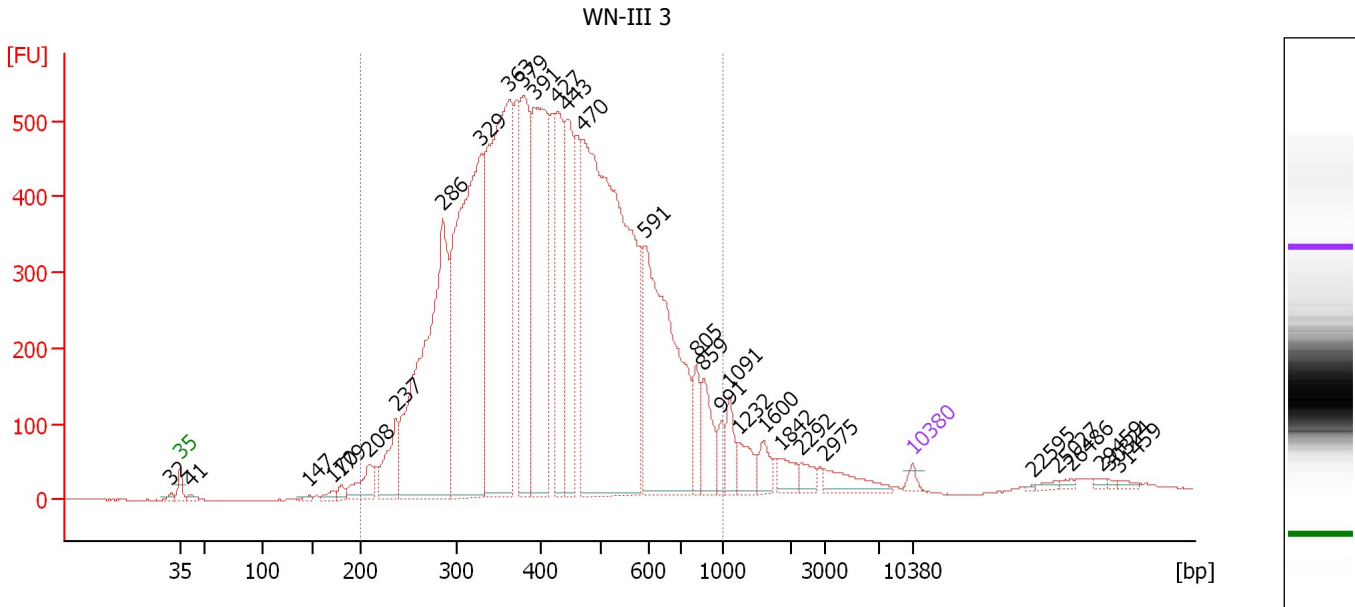
**Region table for sample 4 : WN-II 3**

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	381	2,298.5	30,730.2	7,275.50	98	21.5

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
 Modified: 7/29/2016 12:18:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : WN-III 3**

Number of peaks found: 31                      Corr. Area 1: 13,994.6  
 Noise: 0.4

**Peak table for sample 5 : WN-III 3**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.13
2	35	125.00	5,411.3	Lower Marker	43.00
3	41	86.50	3,188.3		43.97
4	147	53.25	550.3		55.34
5	170	109.46	975.2		57.53
6	179	105.34	889.8		58.40
7	208	511.38	3,725.2		61.05
8	237	859.28	5,502.7		63.65
9	286	5,945.66	31,502.6		68.13
10	329	6,642.61	30,551.3		71.81
11	363	6,519.27	27,238.4		74.51
12	379	2,668.04	10,668.2		75.83
13	391	4,042.17	15,645.3		76.85
14	427	1,968.37	6,978.9		79.10
15	443	1,997.23	6,836.4		79.96
16	470	9,397.17	30,316.1		81.49
17	591	4,047.26	10,380.8		87.36
18	805	455.63	857.4		92.35
19	859	590.41	1,040.9		93.07
20	991	208.42	318.6		94.80
21	1,091	335.78	466.2		95.51
22	1,232	330.70	406.7		96.43
23	1,600	231.29	219.1		98.83
24	1,842	214.65	176.6		100.41
25	2,292	158.78	105.0		102.40

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
 Modified: 7/29/2016 12:18:45 PM

### Electropherogram Summary Continued ...

#### ... Peak table for sample 5 : WN-III 3

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,975	299.79	152.7		104.64
27	10,380	75.00	10.9	Upper Marker	113.00
28	22,595	0.00	0.0		124.52
29	25,027	0.00	0.0		126.82
30	26,486	0.00	0.0		128.19
31	29,459	0.00	0.0		131.00
32	30,324	0.00	0.0		131.81
33	31,459	0.00	0.0		132.88

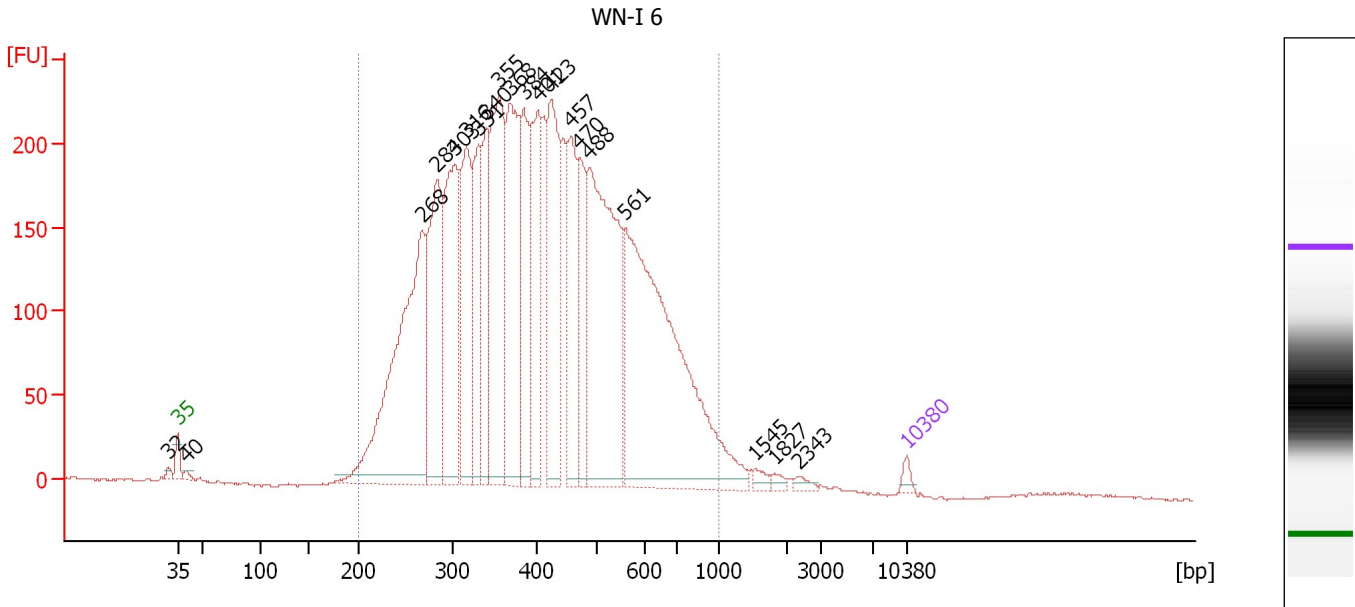
#### Region table for sample 5 : WN-III 3

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	439	13,994.6	192,588.3	49,121.80	94	31.9

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
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**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : WN-I 6**

Number of peaks found: 20                      Corr. Area 1: 6,628.2  
 Noise: 0.6

**Peak table for sample 6 : WN-I 6**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.13
2	35	125.00	5,411.3	Lower Marker	43.00
3	40	78.87	2,999.5		43.77
4	268	5,479.00	30,995.2		66.49
5	284	2,937.80	15,695.6		67.92
6	303	2,916.01	14,578.8		69.66
7	316	2,283.69	10,940.7		70.73
8	331	1,657.20	7,592.1		71.91
9	340	1,504.65	6,702.1		72.68
10	355	3,106.12	13,247.7		73.91
11	368	2,717.46	11,174.7		74.98
12	384	1,725.41	6,804.8		76.26
13	401	2,029.72	7,673.9		77.59
14	423	2,365.78	8,467.1		78.87
15	457	1,640.81	5,442.6		80.76
16	470	1,234.14	3,975.7		81.53
17	488	4,572.17	14,184.0		82.55
18	561	6,353.19	17,168.7		85.98
19	1,545	91.92	90.2		98.47
20	1,827	62.95	52.2		100.31
21	2,343	68.95	44.6		102.56
22	10,380	75.00	10.9	Upper Marker	113.00


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
Modified: 7/29/2016 12:18:45 PM

**Electropherogram Summary Continued ...**

... Region table for sample 6 :

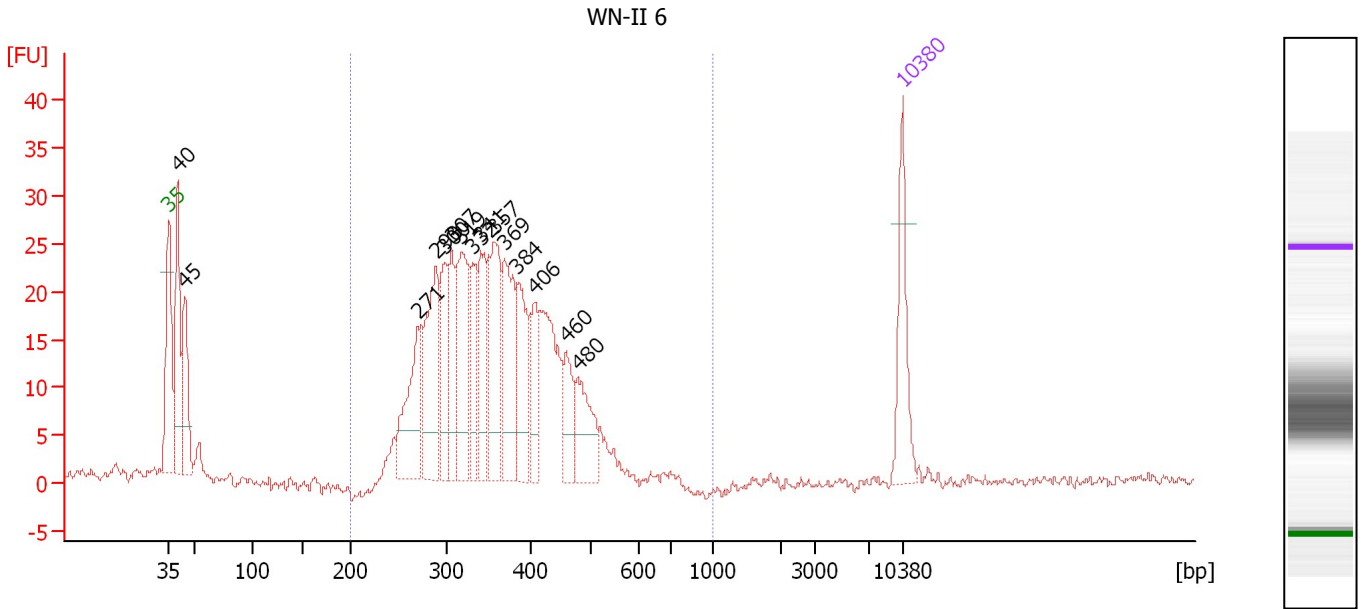
WN-I 6

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	427	6,628.2	180,544.5	 44,272.78	98	33.2

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
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**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : WN-II 6**

Number of peaks found: 15                      Corr. Area 1: 465.0  
 Noise: 0.6

**Peak table for sample 7 : WN-II 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	40	182.71	6,898.3		43.81
3	45	130.52	4,431.9		44.53
4	271	135.80	759.8		66.76
5	290	167.80	877.2		68.49
6	300	101.60	513.3		69.40
7	307	84.06	414.3		70.01
8	319	164.46	782.0		70.93
9	332	83.52	380.7		72.05
10	341	102.93	457.1		72.76
11	357	155.85	661.8		74.03
12	369	138.36	567.7		75.05
13	384	105.25	415.6		76.22
14	406	61.45	229.2		77.90
15	460	55.80	183.7		80.95
16	480	80.67	254.7		82.07
17	10,380	75.00	10.9	Upper Marker	113.00

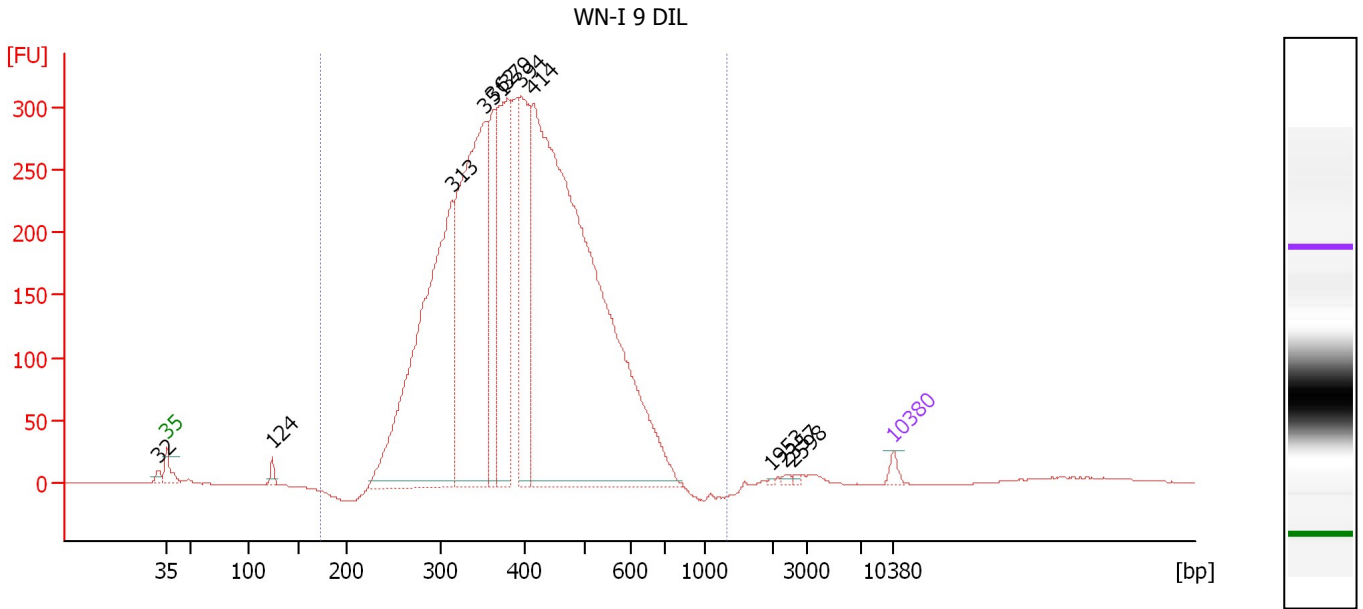
**Region table for sample 7 : WN-II 6**

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	363	465.0	7,548.2	1,716.54	85	19.8

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
 Modified: 7/29/2016 12:18:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : WN-I 9 DIL**

Number of peaks found: 11                      Corr. Area 1: 6,465.8  
 Noise: 0.2

**Peak table for sample 8 : WN-I 9 DIL**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.21
2	35	125.00	5,411.3	Lower Marker	43.00
3	124	115.68	1,416.8		53.18
4	313	6,602.86	31,944.4		70.48
5	351	5,978.70	25,776.8		73.60
6	362	1,356.46	5,680.6		74.44
7	379	2,783.41	11,137.6		75.81
8	394	2,304.91	8,858.9		77.08
9	414	13,058.34	47,786.3		78.34
10	1,953	12.54	9.7		101.13
11	2,357	30.11	19.4		102.61
12	2,598	20.63	12.0		103.40
13	10,380	75.00	10.9	Upper Marker	113.00

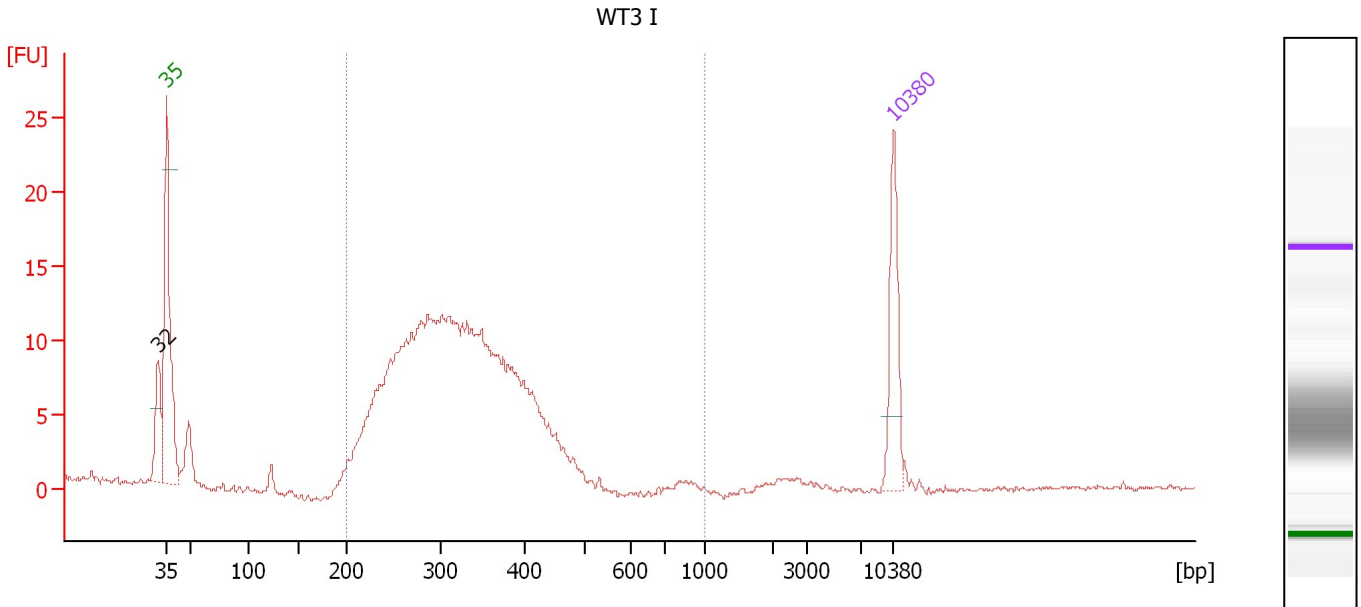
**Region table for sample 8 : WN-I 9 DIL**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
173	1,312	406	6,465.8	130,700.9	32,570.47	99	23.2

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
 Modified: 7/29/2016 12:18:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 9 : WT3 I**

Number of peaks found: 1                      Corr. Area 1: 221.5  
 Noise: 0.1

**Peak table for sample 9 : WT3 I**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.15
2	35	125.00	5,411.3	Lower Marker	43.00
3	10,380	75.00	10.9	Upper Marker	113.00

**Region table for sample 9 : WT3 I**

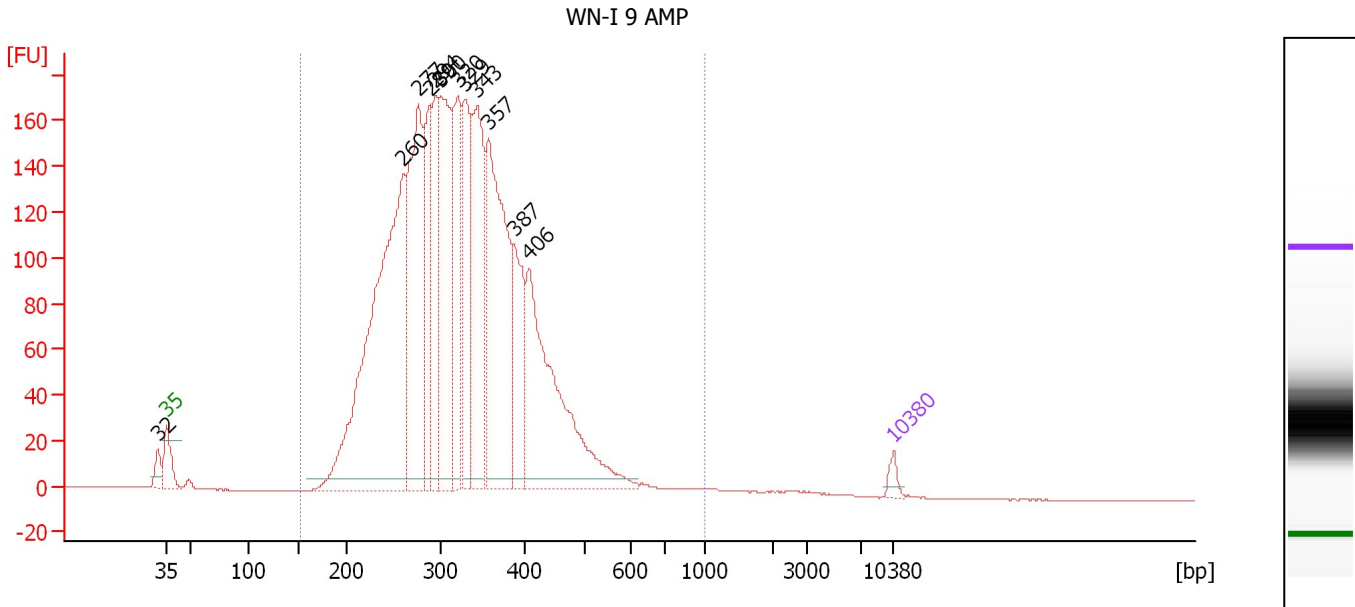
From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	322	221.5	6,677.9	1,346.61	91	20.6



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
 Modified: 7/29/2016 12:18:45 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 10 : WN-I 9 AMP**

Number of peaks found: 12                      Corr. Area 1: 3,537.3  
 Noise: 0.2

**Peak table for sample 10 : WN-I 9 AMP**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.15
2	35	125.00	5,411.3	Lower Marker	43.00
3	260	5,832.98	33,981.6		65.78
4	277	2,852.43	15,629.8		67.28
5	289	1,299.66	6,817.5		68.40
6	294	1,339.01	6,911.6		68.82
7	300	2,241.29	11,319.9		69.41
8	320	1,422.10	6,726.7		71.06
9	329	1,470.68	6,776.2		71.76
10	343	2,008.36	8,864.7		72.93
11	357	3,000.97	12,735.2		74.05
12	387	1,082.65	4,243.7		76.45
13	406	2,736.31	10,209.1		77.89
14	10,380	75.00	10.9	Upper Marker	113.00

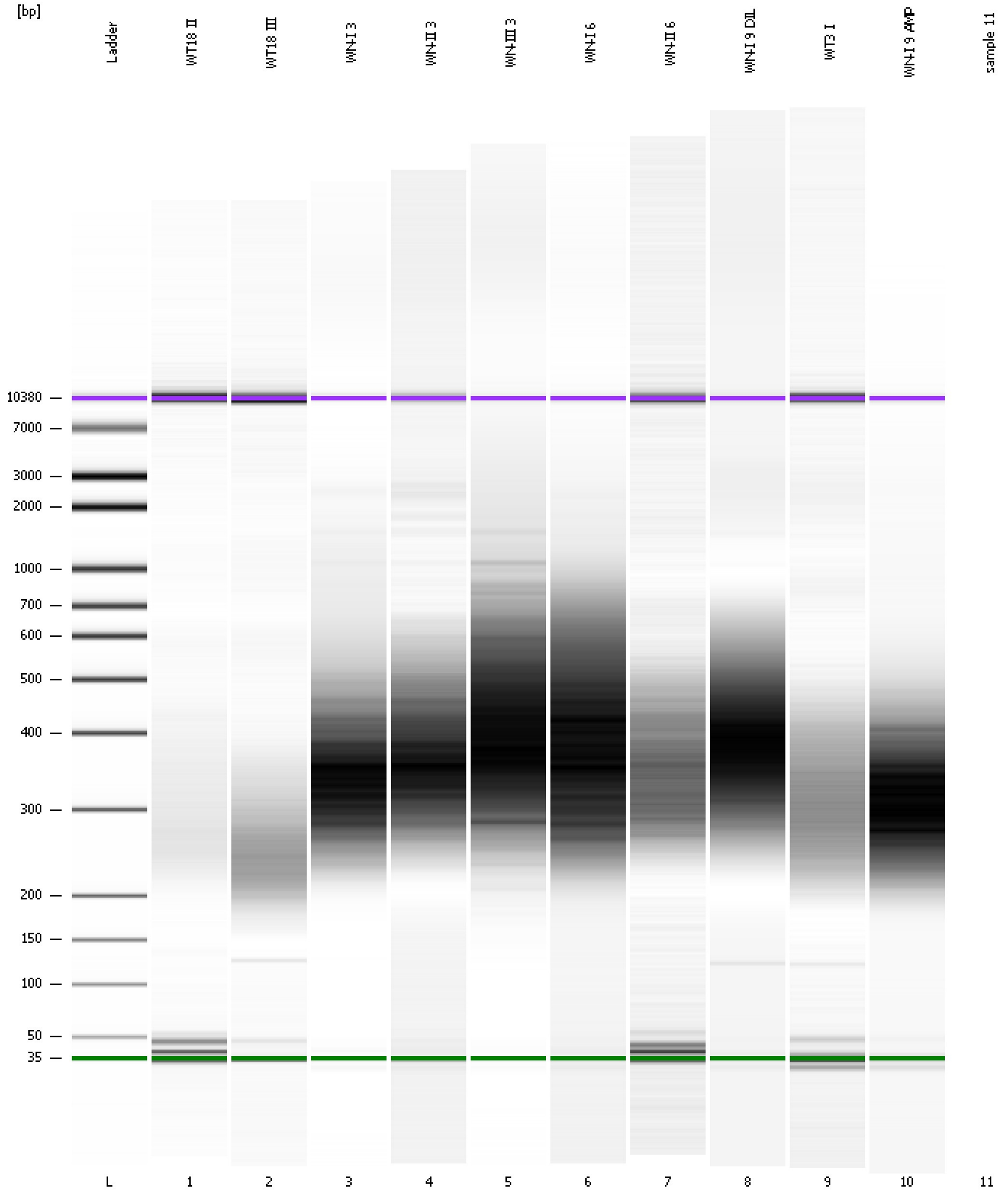
**Region table for sample 10 : WN-I 9 AMP**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
153	1,000	329	3,537.3	128,309.2	25,932.48	99	24.8

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
Modified: 7/29/2016 12:18:45 PM

**Gel Image**



Assay Class: High Sensitivity DNA Assay  
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
Modified: 7/29/2016 12:18:45 PM

**Invalid Samples**

Sample 11 has not been run, no results available.

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-29\2016-07-29\_001.xad

Created: 7/29/2016 11:29:26 AM  
 Modified: 7/29/2016 12:18:45 PM

**Run Logbook**

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 11)		Instrument	Run		7/29/2016 12:07:53 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\2016-07-29\2016-07-29_001.xad)		Instrument	Run		7/29/2016 11:29:32 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		7/29/2016 11:29:32 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		7/29/2016 11:29:32 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		7/29/2016 11:29:32 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		7/29/2016 11:29:32 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		7/29/2016 11:29:32 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		7/29/2016 11:29:32 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1