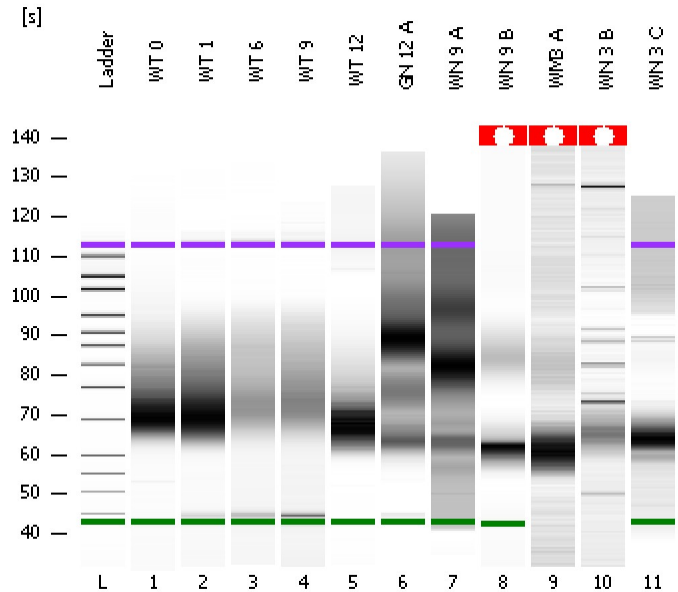


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
Modified: 7/19/2016 2:22:30 PM

Electrophoresis File Run Summary



Instrument Information:

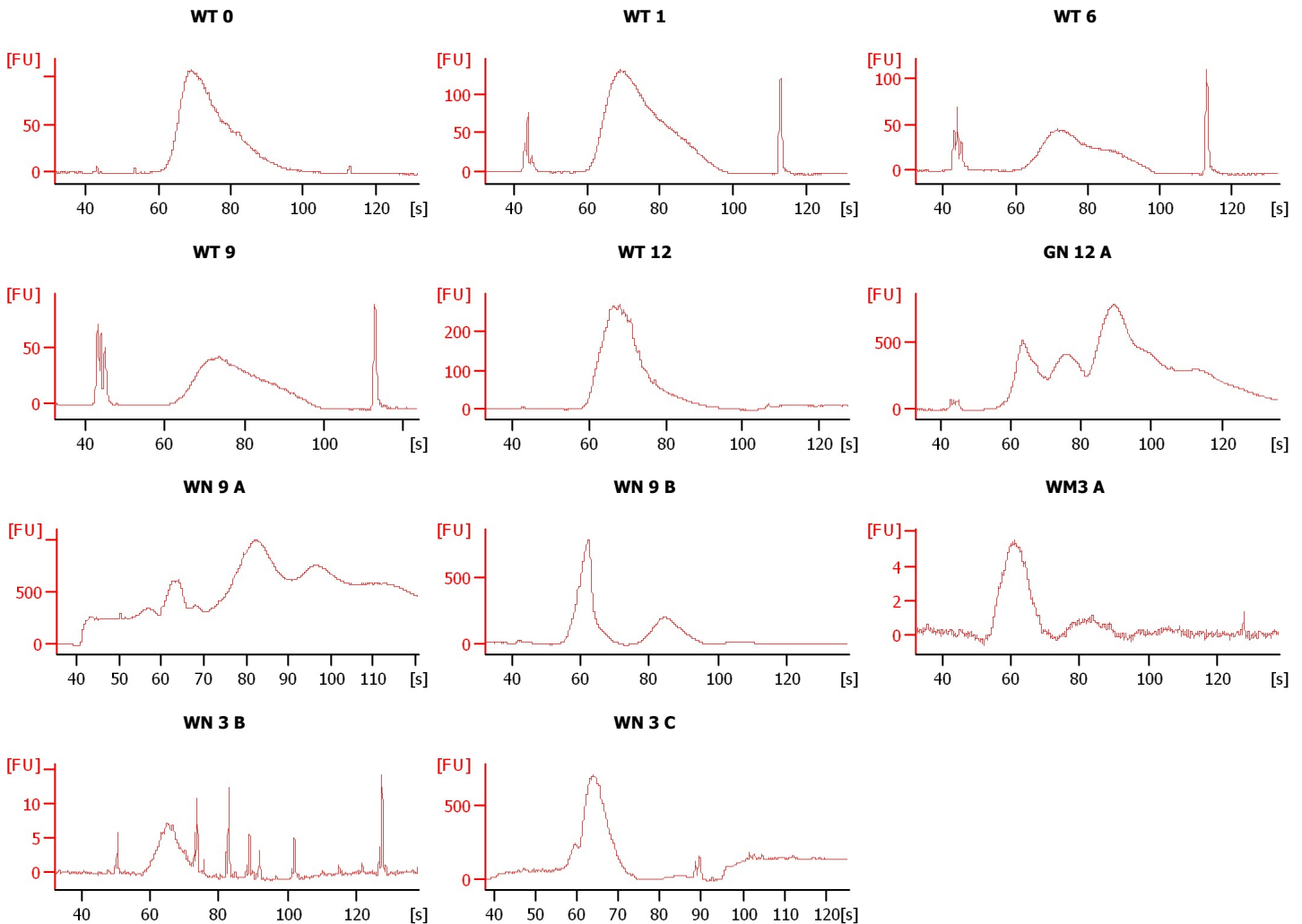
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
Modified: 7/19/2016 2:22:30 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
WT 0		<input type="checkbox"/>	✓			
WT 1		<input type="checkbox"/>	✓			
WT 6		<input type="checkbox"/>	✓			
WT 9		<input type="checkbox"/>	✓			
WT 12		<input type="checkbox"/>	✓			
GN 12 A		<input type="checkbox"/>	✓			
WN 9 A		<input type="checkbox"/>	✓			
WN 9 B		<input type="checkbox"/>	✓			
WM3 A		<input type="checkbox"/>	✓			
WN 3 B		<input type="checkbox"/>	✓			
WN 3 C		<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\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
Modified: 7/19/2016 2:22:30 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

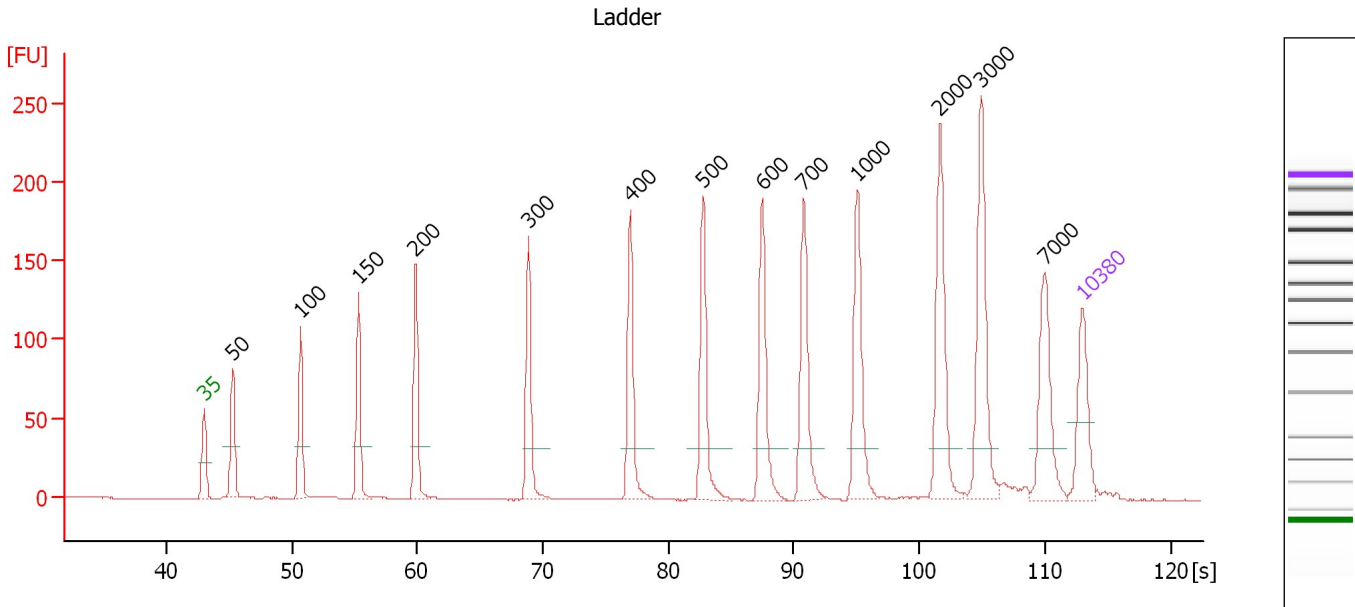
Ladder

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\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
 Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

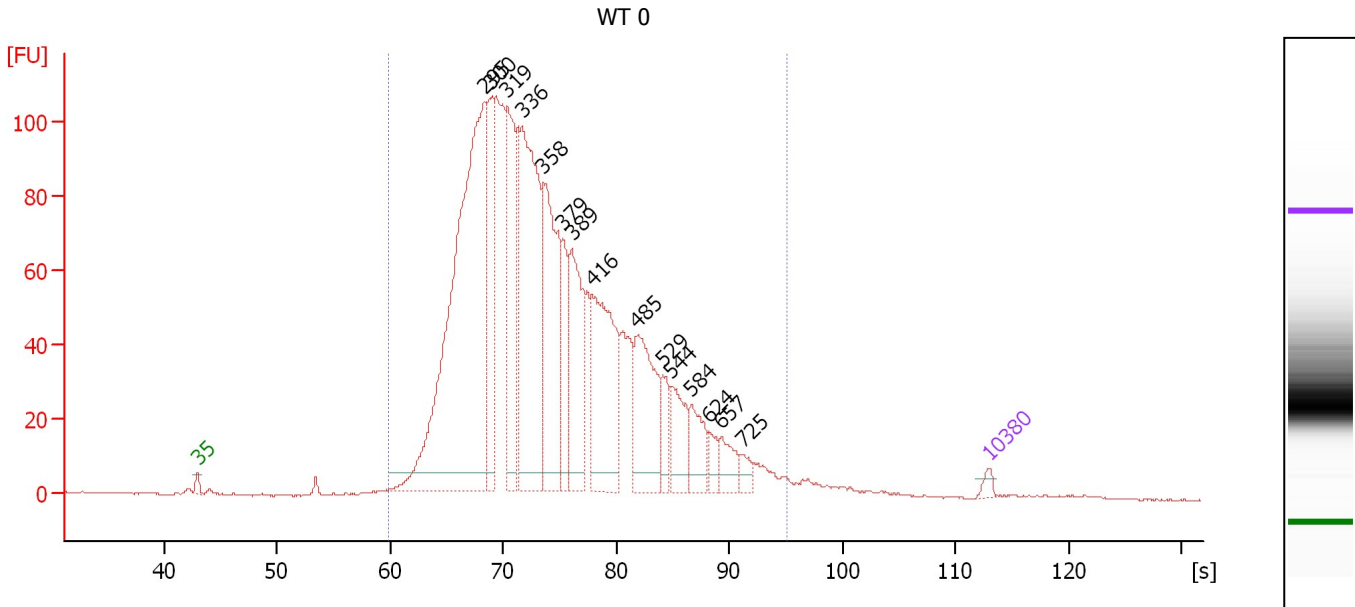
Peak table for Ladder

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	50.70
4	150	150.00	1,515.2	Ladder Peak	55.32
5	200	150.00	1,136.4	Ladder Peak	59.86
6	300	150.00	757.6	Ladder Peak	68.84
7	400	150.00	568.2	Ladder Peak	76.97
8	500	150.00	454.5	Ladder Peak	82.75
9	600	150.00	378.8	Ladder Peak	87.46
10	700	150.00	324.7	Ladder Peak	90.79
11	1,000	150.00	227.3	Ladder Peak	95.07
12	2,000	150.00	113.6	Ladder Peak	101.62
13	3,000	150.00	75.8	Ladder Peak	104.91
14	7,000	150.00	32.5	Ladder Peak	110.00
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

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 Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : WT 0

Number of peaks found: 15 Corr. Area 1: 2,198.4
 Noise: 0.2

Peak table for sample 1 : WT 0

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	295	9,578.29	49,171.3		68.41
3	300	2,004.52	10,107.9		68.88
4	319	2,313.31	10,981.8		70.40
5	336	4,442.76	20,062.6		71.73
6	358	2,986.54	12,629.1		73.58
7	379	899.32	3,592.1		75.29
8	389	1,899.46	7,393.3		76.10
9	416	2,498.09	9,095.6		77.91
10	485	1,745.51	5,450.8		81.89
11	529	431.11	1,234.2		84.13
12	544	779.57	2,169.7		84.84
13	584	586.00	1,521.0		86.69
14	624	232.29	564.0		88.26
15	657	385.73	889.9		89.35
16	725	187.78	392.2		91.15
17	10,380	75.00	10.9	Upper Marker	113.00

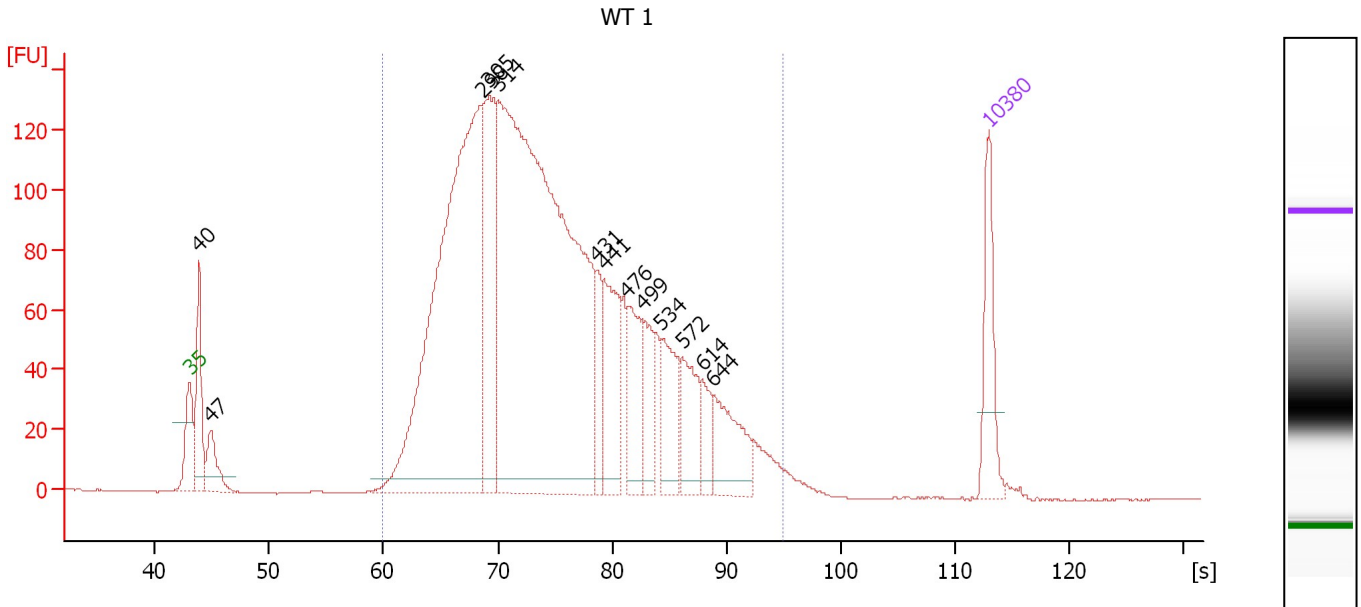
Region table for sample 1 : WT 0

From [s]	To [s]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
59.86	95.07	383	2,198.4	164,808.3	37,575.87	97	31.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
 Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : WT 1

Number of peaks found: 13 Corr. Area 1: 3,199.3
 Noise: 0.2

Peak table for sample 2 : WT 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	40	141.23	5,324.1		43.80
3	47	73.26	2,351.0		44.88
4	298	975.35	4,963.7		68.64
5	305	278.93	1,385.6		69.25
6	314	1,458.74	7,033.1		70.00
7	431	58.90	207.1		78.75
8	441	137.99	474.6		79.32
9	476	100.68	320.2		81.39
10	499	72.56	220.2		82.70
11	534	94.52	268.2		84.35
12	572	82.15	217.6		86.14
13	614	36.00	88.8		87.93
14	644	95.26	224.2		88.91
15	10,380	75.00	10.9	Upper Marker	113.00

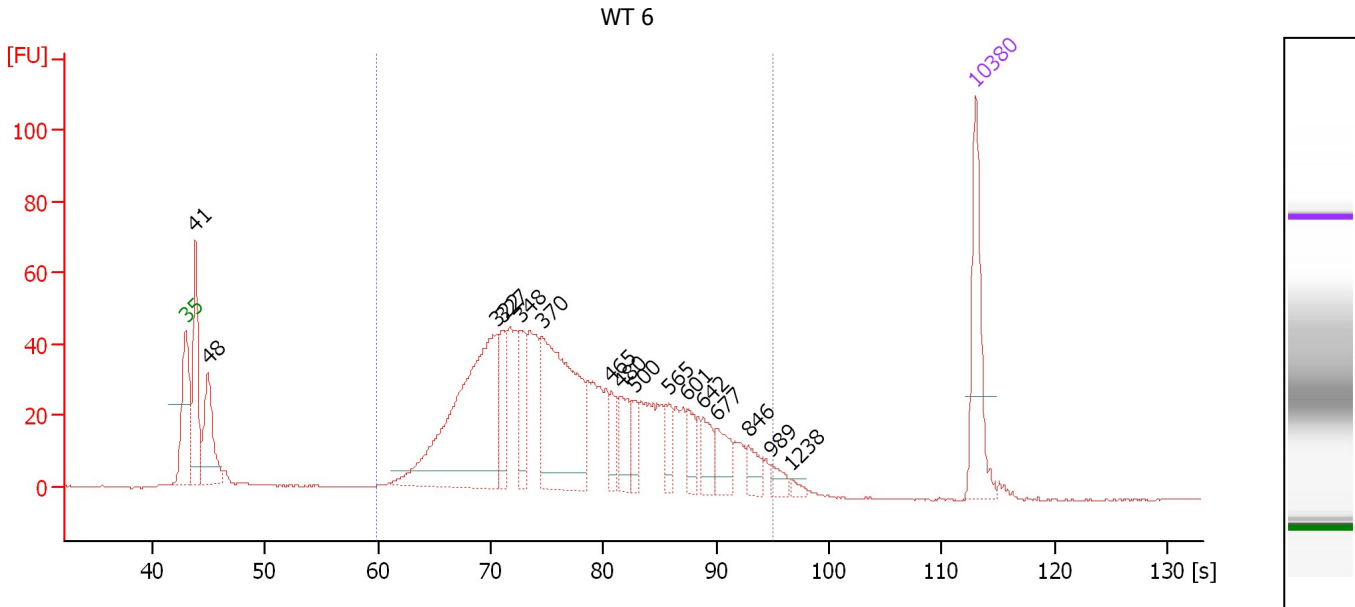
Region table for sample 2 : WT 1

From [s]	To [s]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
59.86	95.07	393	3,199.3	15,005.7	3,429.78	95	33.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

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 Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : WT 6

Number of peaks found: 16 Corr. Area 1: 1,164.7
 Noise: 0.2

Peak table for sample 3 : WT 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	41	140.70	5,255.9		43.86
3	48	112.46	3,574.7		44.95
4	322	341.63	1,609.5		70.60
5	327	52.05	241.3		71.03
6	348	53.19	231.6		72.74
7	370	236.48	967.9		74.55
8	465	23.98	78.1		80.74
9	480	40.19	126.9		81.59
10	500	23.53	71.3		82.73
11	565	20.52	55.0		85.83
12	601	24.36	61.4		87.49
13	642	27.17	64.1		88.87
14	677	32.46	72.7		90.02
15	846	20.18	36.2		92.87
16	989	11.23	17.2		94.92
17	1,238	5.19	6.3		96.63
18	10,380	75.00	10.9	Upper Marker	113.00

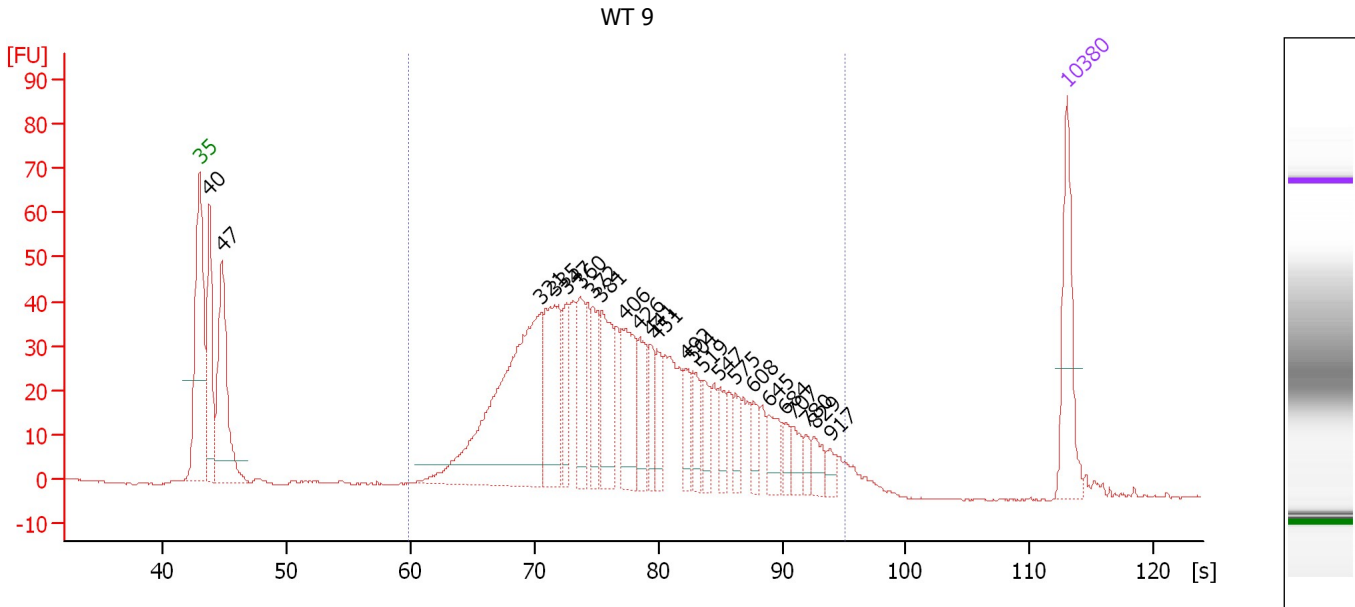
Region table for sample 3 : WT 6

From [s]	To [s]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
59.86	95.07	436	1,164.7	5,427.5	1,358.06	85	34.4

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

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 Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : WT 9

Number of peaks found: 24 Corr. Area 1: 1,107.0
 Noise: 0.2

Peak table for sample 4 : WT 9

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	172.25	6,514.0		43.78
3	47	228.33	7,389.3		44.82
4	321	403.18	1,901.9		70.57
5	335	143.59	650.4		71.65
6	347	56.92	248.7		72.65
7	360	73.21	308.5		73.69
8	372	48.82	198.6		74.73
9	381	87.00	346.0		75.42
10	406	92.43	344.8		77.33
11	426	54.51	193.7		78.50
12	441	35.31	121.2		79.37
13	451	36.34	122.0		79.93
14	492	28.89	88.9		82.31
15	501	31.78	96.1		82.79
16	519	30.71	89.6		83.66
17	547	29.41	81.5		84.96
18	575	23.57	62.2		86.26
19	608	19.66	49.0		87.73
20	645	33.25	78.2		88.94
21	684	16.57	36.7		90.24
22	707	20.92	44.8		90.89
23	780	13.04	25.3		91.93
24	829	19.11	34.9		92.63
25	917	12.17	20.1		93.89

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...**... Peak table for sample 4 : WT 9**

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

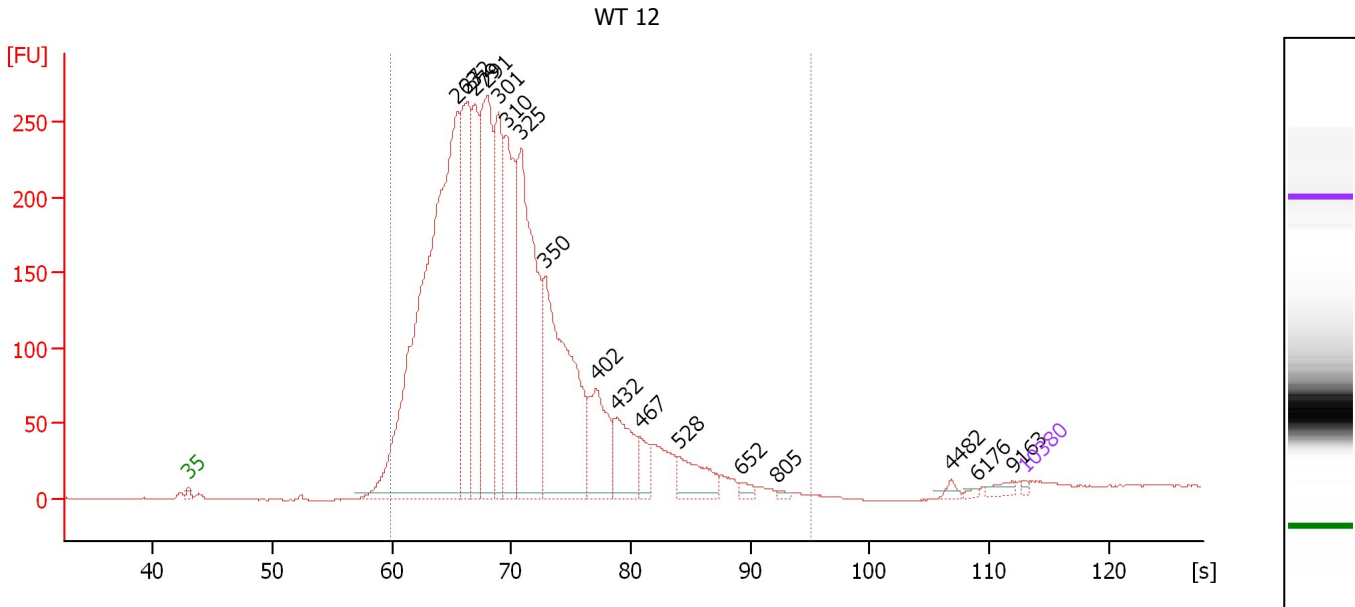
Region table for sample 4 : WT 9

From [s]	To [s]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/ μ l]	% of Total	Size distribution in CV [%]
59.86	95.07	436	1,107.0	6,626.2	1,677.29	84	33.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
 Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : WT 12

Number of peaks found: 17 Corr. Area 1: 4,881.9
 Noise: 0.2

Peak table for sample 5 : WT 12

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	263	38,705.17	223,069.5		65.51
3	272	10,390.90	57,903.5		66.32
4	279	7,709.41	41,882.5		66.95
5	291	12,154.84	63,308.8		68.03
6	301	7,891.57	39,724.9		68.92
7	310	9,614.87	47,018.4		69.64
8	325	15,121.99	70,551.3		70.86
9	350	13,431.88	58,208.5		72.88
10	402	4,096.66	15,431.9		77.10
11	432	3,072.11	10,780.3		78.81
12	467	1,084.13	3,519.0		80.83
13	528	2,137.24	6,133.5		84.07
14	652	364.68	847.6		89.19
15	805	144.72	272.5		92.29
16	4,482	209.54	70.8		106.80
17	6,176	109.36	26.8		108.96
18	9,163	277.25	45.8		111.92
19	10,380	75.00	10.9	Upper Marker	113.00

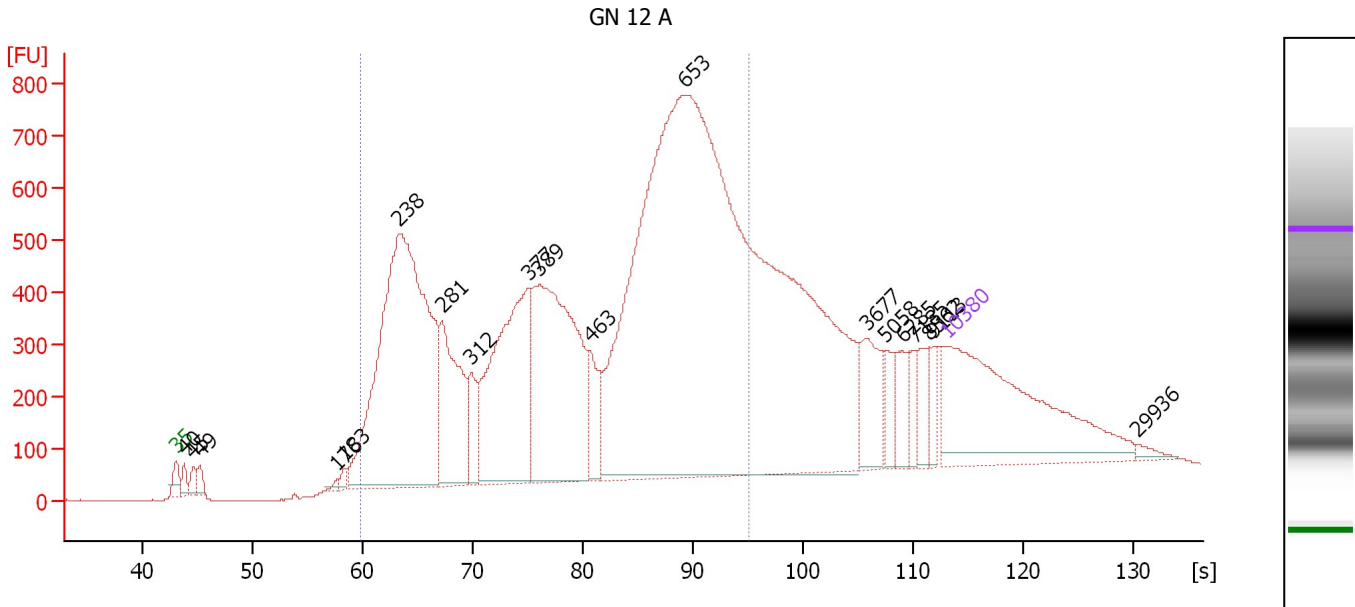
Region table for sample 5 : WT 12

From [s]	To [s]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
59.86	95.07	314	4,881.9	635,269.4	123,673.45	99	24.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
 Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : GN 12 A

Number of peaks found: 19 Corr. Area 1: 18,257.7
 Noise: 0.8

Peak table for sample 6 : GN 12 A

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	5.62	214.1		43.73
3	45	5.73	192.4		44.56
4	49	5.69	176.0		45.15
5	176	1.51	13.0		57.64
6	183	3.11	25.7		58.33
7	238	253.14	1,609.0		63.31
8	281	58.53	315.3		67.16
9	312	16.80	81.5		69.85
10	377	102.31	411.6		75.07
11	389	119.53	466.0		76.05
12	463	17.89	58.5		80.64
13	653	579.05	1,343.4		89.23
14	3,677	19.01	7.8		105.78
15	5,058	6.42	1.9		107.53
16	6,285	9.94	2.4		109.09
17	7,185	6.33	1.3		110.17
18	8,562	8.51	1.5		111.39
19	9,113	5.01	0.8		111.88
20	10,380	75.00	10.9	Upper Marker	113.00
21	29,936	0.00	0.0		130.33

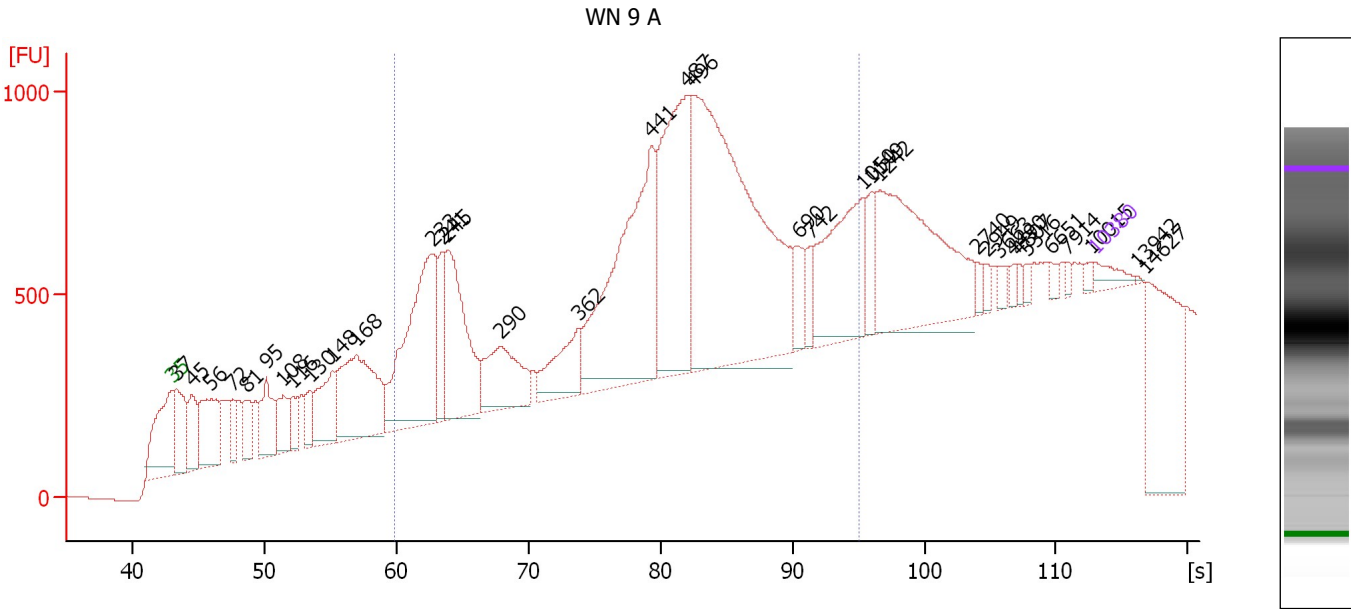
Region table for sample 6 : GN 12 A

From [s]	To [s]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
59.86	95.07	508	18,257.7	3,770.4	960.66	76	40.6

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
 Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : WN 9 A

Number of peaks found: 35 Corr. Area 1: 19,251.6
 Noise: 1.4

Peak table for sample 7 : WN 9 A

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	37	660.62	26,788.8		43.36
3	45	464.30	15,639.3		44.54
4	56	777.75	20,915.6		45.99
5	72	225.45	4,762.3		47.65
6	81	286.18	5,370.8		48.62
7	95	526.95	8,404.9		50.16
8	108	360.38	5,047.9		51.46
9	116	195.51	2,552.6		52.18
10	130	206.29	2,403.3		53.48
11	148	588.23	6,003.9		55.18
12	168	1,335.24	12,007.1		57.00
13	233	1,836.51	11,940.6		62.83
14	241	354.20	2,229.7		63.51
15	245	1,225.14	7,584.6		63.88
16	290	667.76	3,486.1		67.97
17	362	438.24	1,834.9		73.87
18	441	2,060.72	7,081.7		79.34
19	487	1,635.86	5,088.1		82.01
20	496	3,543.08	10,818.1		82.53
21	690	184.84	405.8		90.46
22	742	117.88	240.7		91.39
23	1,050	829.79	1,197.5		95.40
24	1,149	187.33	247.1		96.05
25	1,242	1,270.34	1,550.3		96.65

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
 Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...

... Peak table for sample 7 : WN 9 A

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	2,740	41.69	23.1		104.06
27	2,949	35.92	18.5		104.75
28	3,663	52.76	21.8		105.76
29	4,490	31.56	10.7		106.81
30	4,807	30.69	9.7		107.21
31	5,316	32.10	9.1		107.86
32	6,651	36.76	8.4		109.56
33	7,914	24.16	4.6		110.82
34	10,015	27.53	4.2		112.68
35	10,380	75.00	10.9	Upper Marker	113.00
36	13,942	0.00	0.0		116.16
37	14,627	0.00	0.0		116.76

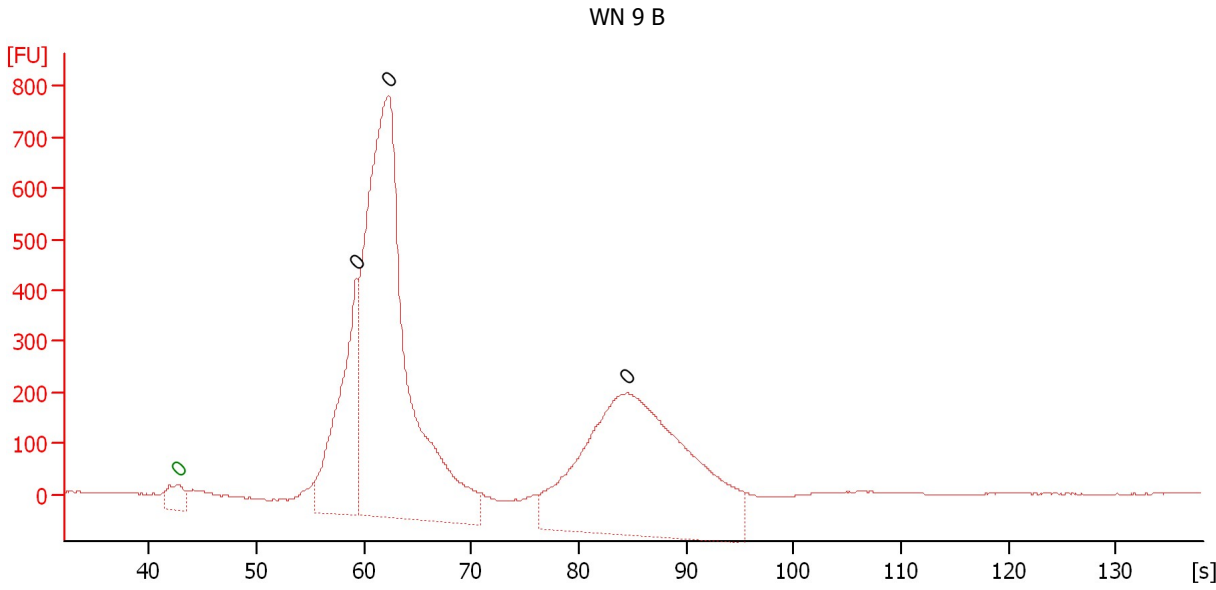
Region table for sample 7 : WN 9 A

From [s]	To [s]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
59.86	95.07	482	19,251.6	58,440.8	14,581.12	61	38.3

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : WN 9 B

Number of peaks found: 3 Noise: 0.2

Peak table for sample 8 : WN 9 B

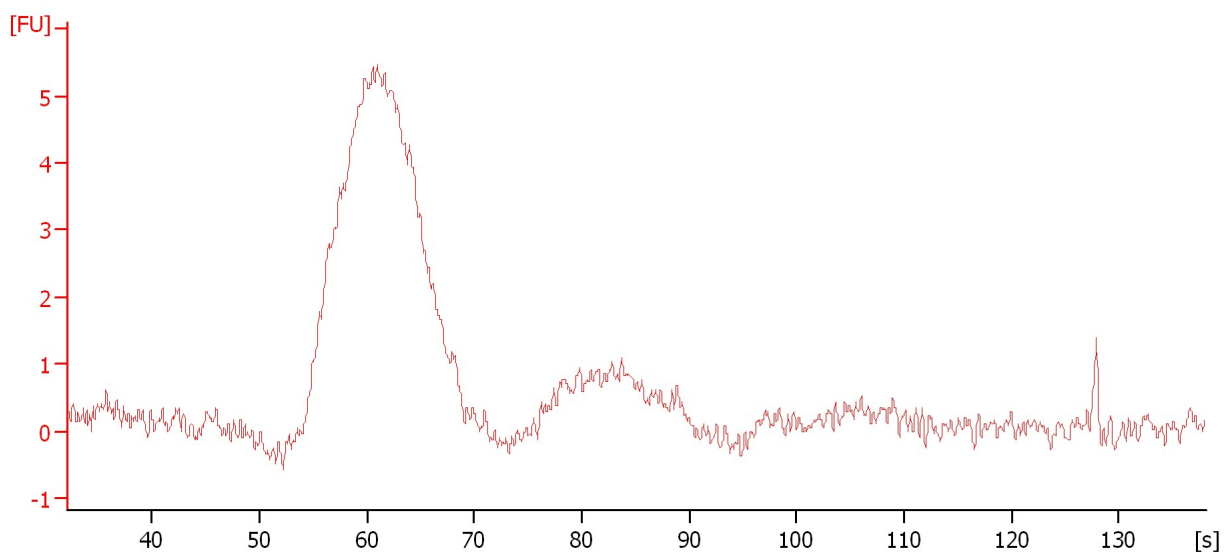
Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	0	0.00	0.0	Lower Marker	42.70
2	0	0.00	0.0		59.30
3	0	0.00	0.0		62.30
4	0	0.00	0.0		84.50

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...

WM3 A



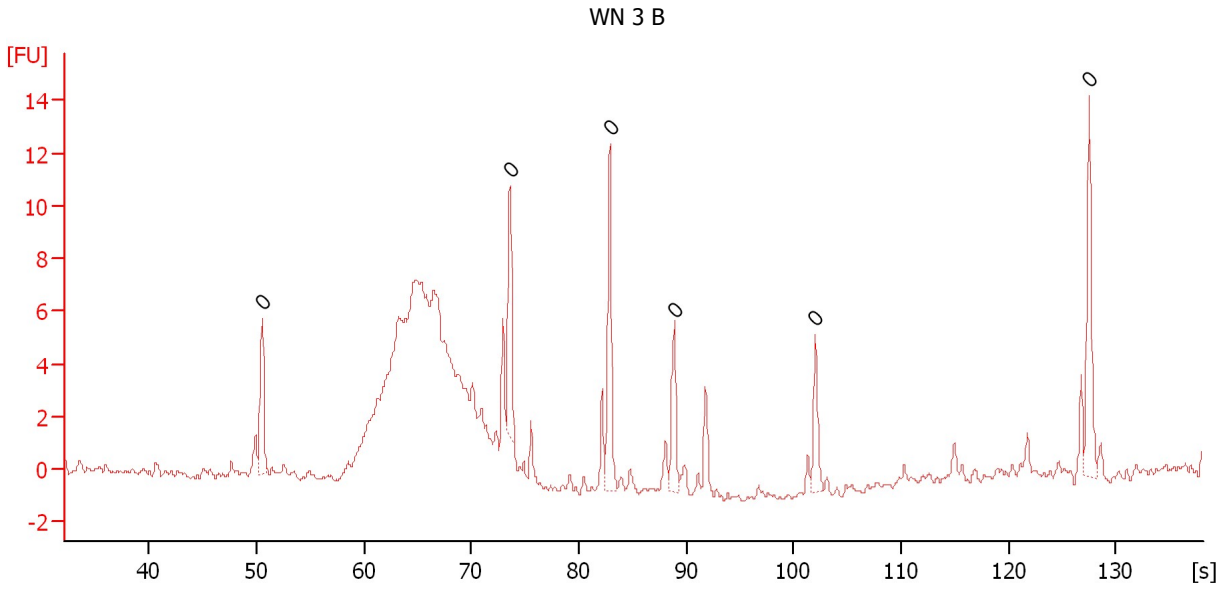
Overall Results for sample 9 : WM3 A

Noise: 0.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : WN 3 B

Number of peaks found: 0 Noise: 0.2

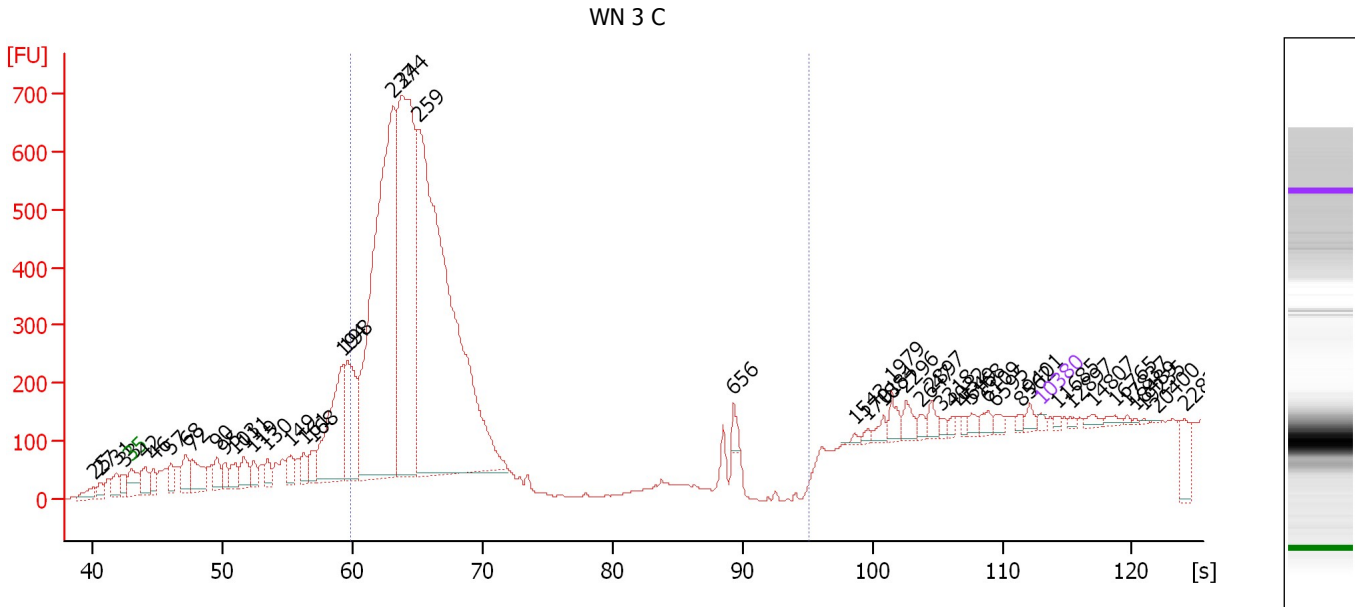
Peak table for sample 10 : WN 3 B

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	0	0.00	0.0		50.55
2	0	0.00	0.0		73.60
3	0	0.00	0.0		82.90
4	0	0.00	0.0		88.85
5	0	0.00	0.0		102.00
6	0	0.00	0.0		127.55

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

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 Modified: 7/19/2016 2:22:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : WN 3 C

Number of peaks found: 48 Corr. Area 1: 7,819.9
 Noise: 1.2

Peak table for sample 11 : WN 3 C

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	25	0.00	0.0		40.11
2	27	0.00	0.0		40.69
3	31	0.00	0.0		41.84
4	33	0.00	0.0		42.42
5	35	125.00	5,411.3	Lower Marker	43.00
6	42	771.21	27,840.8		44.07
7	46	465.12	15,233.8		44.73
8	57	566.17	15,072.3		46.06
9	68	888.11	19,894.1		47.21
10	72	1,152.94	24,183.2		47.71
11	90	623.99	10,522.1		49.61
12	95	411.84	6,553.6		50.19
13	103	578.27	8,477.2		51.01
14	111	688.74	9,443.5		51.67
15	119	392.96	5,022.3		52.42
16	130	455.18	5,298.4		53.49
17	149	515.12	5,224.8		55.27
18	161	447.37	4,216.7		56.30
19	168	480.73	4,346.5		56.92
20	194	3,580.07	28,028.3		59.27
21	198	2,610.28	20,012.1		59.64
22	237	14,863.14	94,967.4		63.19
23	244	11,079.83	68,794.1		63.81
24	259	19,202.13	112,248.3		65.18
25	656	208.29	480.8		89.34

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
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Electropherogram Summary Continued ...

... Peak table for sample 11 : WN 3 C

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	1,543	63.53	62.4		98.63
27	1,701	73.17	65.2		99.66
28	1,884	162.43	130.6		100.86
29	1,979	253.15	193.9		101.48
30	2,296	241.46	159.4		102.59
31	2,647	121.33	69.5		103.75
32	2,897	191.02	99.9		104.58
33	3,318	82.21	37.5		105.32
34	4,032	84.49	31.7		106.23
35	4,649	72.70	23.7		107.01
36	5,168	115.05	33.7		107.67
37	6,109	158.78	39.4		108.87
38	6,595	113.90	26.2		109.49
39	8,562	69.83	12.4		111.39
40	9,401	129.23	20.8		112.13
41	10,380	75.00	10.9	Upper Marker	113.00
42	11,685	0.00	0.0		114.16
43	12,897	0.00	0.0		115.23
44	14,807	0.00	0.0		116.92
45	16,765	0.00	0.0		118.66
46	17,837	0.00	0.0		119.61
47	18,489	0.00	0.0		120.19
48	19,235	0.00	0.0		120.85
49	20,400	0.00	0.0		121.88
50	22,824	0.00	0.0		124.03

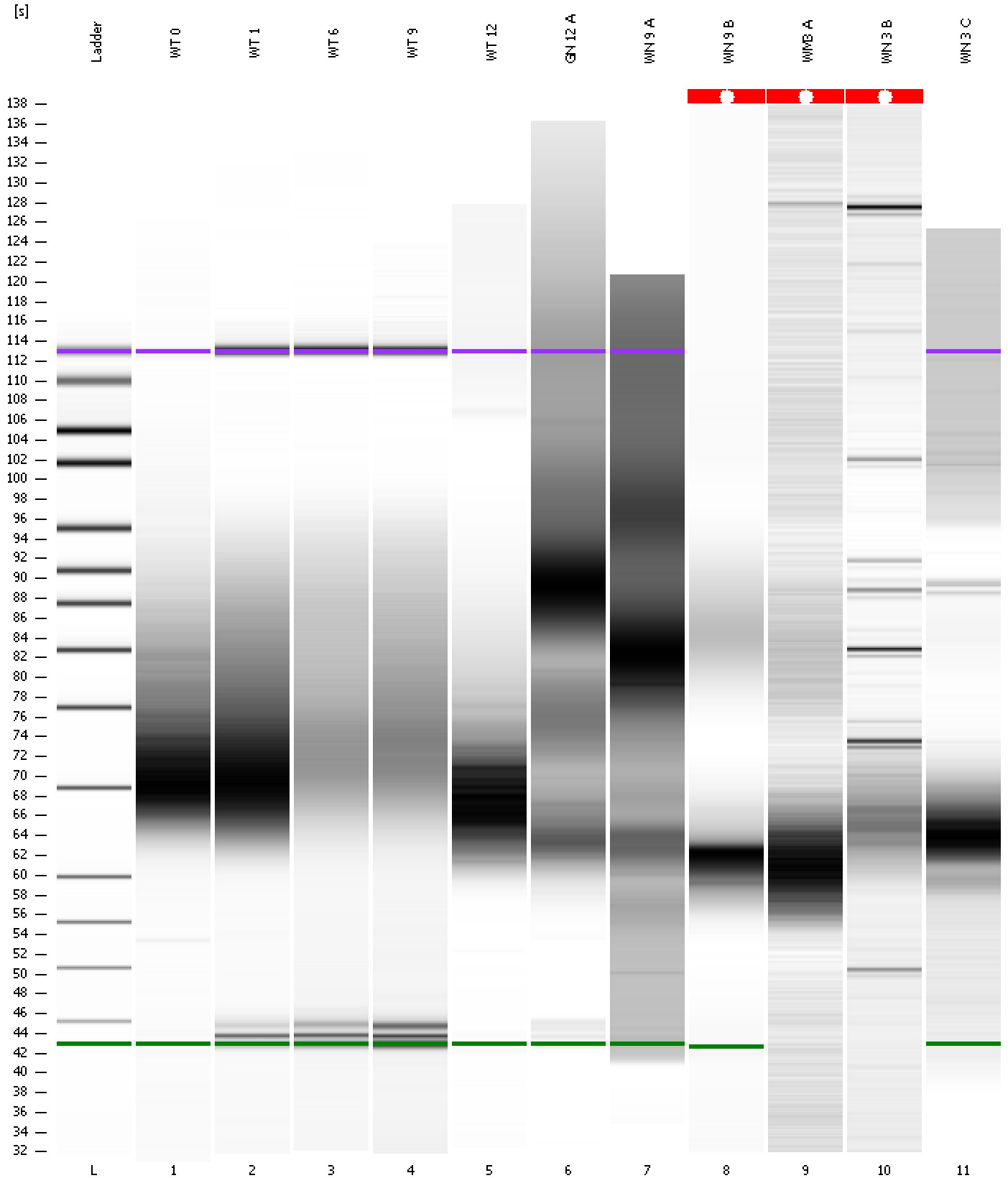
Region table for sample 11 : WN 3 C

From [s]	To [s]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
59.86	95.07	258	7,819.9	273,942.8	45,813.35	71	19.2

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
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Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-07-19\2016-07-19_003.xad

Created: 7/19/2016 1:14:22 PM
 Modified: 7/19/2016 2:22:30 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		7/19/2016 1:55:42 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\2016-07-19\2016-07-19_003.xad)		Instrument	Run		7/19/2016 1:14:28 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		7/19/2016 1:14:28 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		7/19/2016 1:14:28 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		7/19/2016 1:14:28 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		7/19/2016 1:14:28 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		7/19/2016 1:14:28 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		7/19/2016 1:14:28 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1