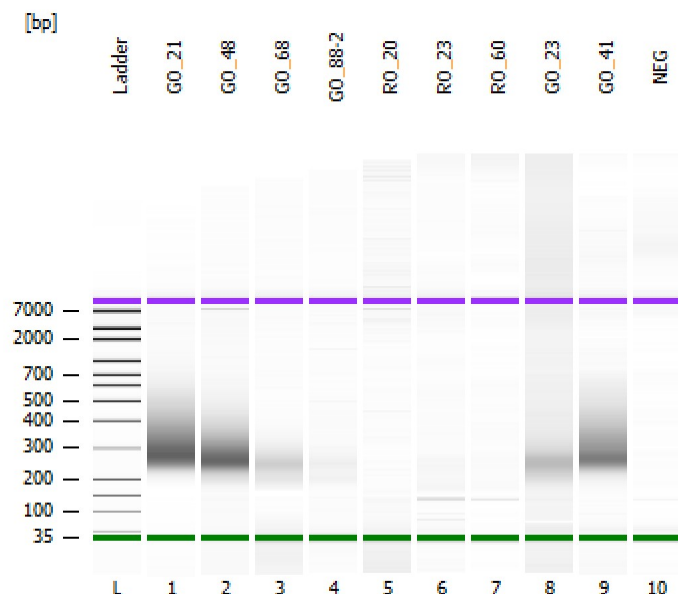


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ioanalyzer\2100 expert\data\2016-05-20\CNelson_1-9and36.xad

Created: 5/20/2016 1:32:18 PM
Modified: 5/20/2016 3:28:28 PM

Electrophoresis File Run Summary



Instrument Information:

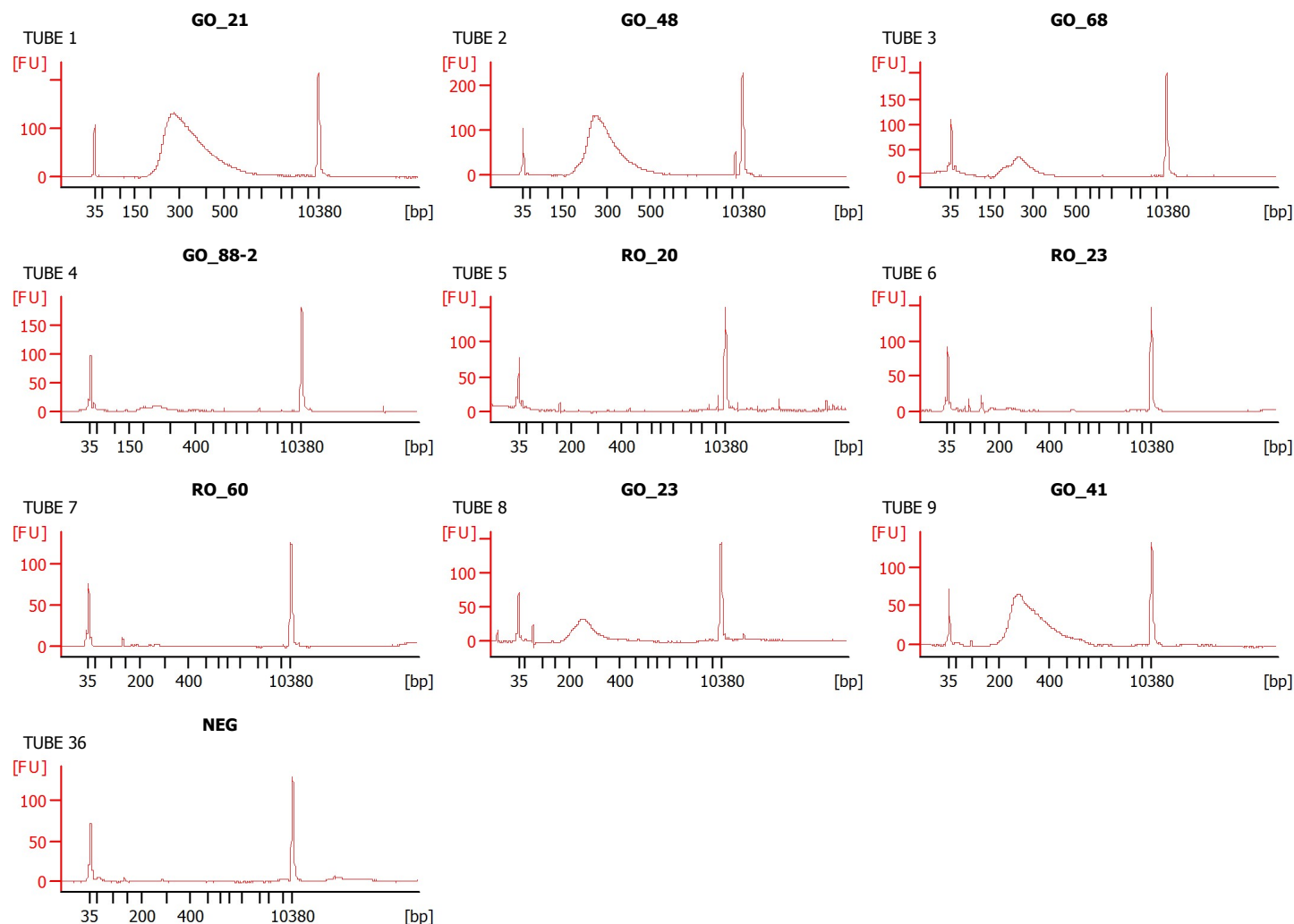
Instrument Name: DE13804163 Firmware: C.01.069
Serial#: DE13804163 Type: G2939A

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:\...ioanalyzer\2100 expert\data\2016-05-20\CNelson_1-9and36.xad

Created: 5/20/2016 1:32:18 PM
Modified: 5/20/2016 3:28:28 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
GO_21	TUBE 1	<input type="checkbox"/>	✓			
GO_48	TUBE 2	<input type="checkbox"/>	✓			
GO_68	TUBE 3	<input type="checkbox"/>	✓			
GO_88-2	TUBE 4	<input type="checkbox"/>	✓			
RO_20	TUBE 5	<input type="checkbox"/>	✓			
RO_23	TUBE 6	<input type="checkbox"/>	✓			
RO_60	TUBE 7	<input type="checkbox"/>	✓			
GO_23	TUBE 8	<input type="checkbox"/>	✓			
GO_41	TUBE 9	<input type="checkbox"/>	✓			
NEG	TUBE 36	<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

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

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ioanalyzer\2100 expert\data\2016-05-20\CNelson_1-9and36.xad

Created: 5/20/2016 1:32:18 PM
Modified: 5/20/2016 3:28:28 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

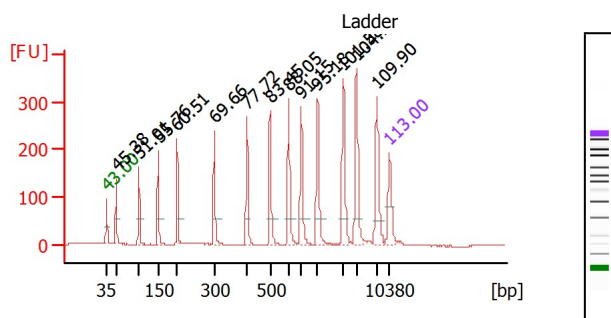
Ladder

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:\...ioanalyzer\2100 expert\data\2016-05-20\CNelson_1-9and36.xad

Created: 5/20/2016 1:32:18 PM
 Modified: 5/20/2016 3:28:28 PM

Electropherogram Summary



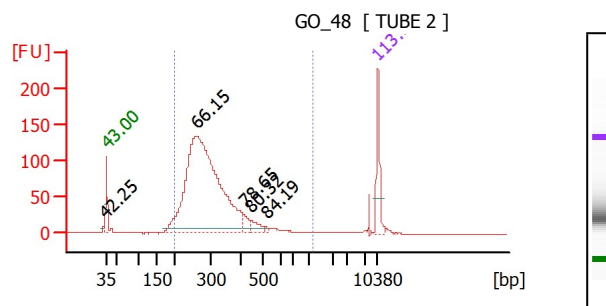
Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ioanalyzer\2100 expert\data\2016-05-20\CNelson_1-9and36.xad

Created: 5/20/2016 1:32:18 PM
 Modified: 5/20/2016 3:28:28 PM

Electropherogram Summary Continued ...

Peak table for sample 1 : GO_21

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	32	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	275	478.13	2,630.5	
4	282	1,036.89	5,575.9	
5	416	151.80	553.5	
6	510	62.17	184.9	
7	635	4.66	11.1	
8	669	2.81	6.4	
9	10,380	75.00	10.9	Upper Marker
10	12,569	0.00	0.0	



Overall Results for sample 2 : GO_48

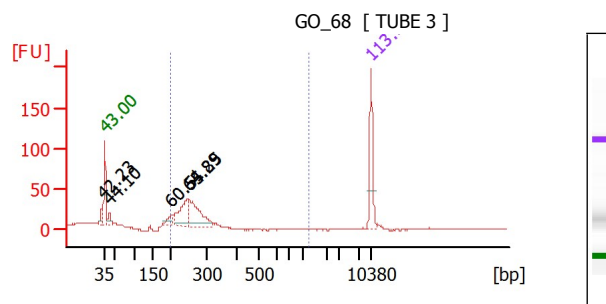
Number of peaks found: 5
 Noise: 0.5
 Corr. Area 1: 2,042.0

Region table for sample 2 : GO_48

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,200	2,042.0	95	309	27.2	1,303.50	6,830.4	Blue

Peak table for sample 2 : GO_48

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	32	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	262	1,310.98	7,590.4	
4	416	25.17	91.6	
5	445	32.98	112.2	
6	516	4.66	13.7	
7	10,380	75.00	10.9	Upper Marker



Overall Results for sample 3 : GO_68

Number of peaks found: 5
 Noise: 0.6
 Corr. Area 1: 315.4

Region table for sample 3 : GO_68

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,200	315.4	68	254	11.7	245.60	1,471.3	Blue

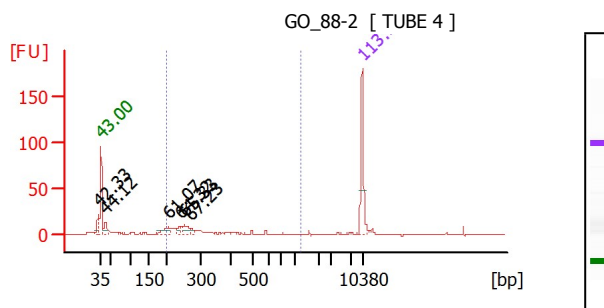
Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ioanalyzer\2100 expert\data\2016-05-20\CNelson_1-9and36.xad

Created: 5/20/2016 1:32:18 PM
 Modified: 5/20/2016 3:28:28 PM

Electropherogram Summary Continued ...

Peak table for sample 3 : GO_68

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	32	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	42	28.67	1,036.1	
4	200	27.64	209.4	
5	247	102.16	625.6	
6	252	121.09	727.4	
7	10,380	75.00	10.9	Upper Marker



Overall Results for sample 4 : GO_88-2

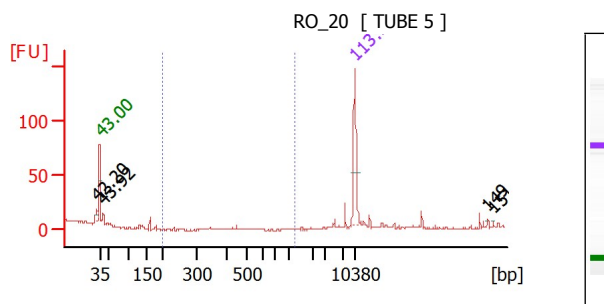
Number of peaks found: 6
 Noise: 0.4
 Corr. Area 1: 119.4

Region table for sample 4 : GO_88-2

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,200	119.4	49	287	33.0	117.86	675.6	Blue

Peak table for sample 4 : GO_88-2

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	33	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	42	35.68	1,285.3	
4	206	32.83	241.3	
5	242	13.48	84.5	
6	253	26.62	159.7	
7	273	7.77	43.0	
8	10,380	75.00	10.9	Upper Marker



Overall Results for sample 5 : RO_20

Number of peaks found: 4
 Noise: 0.6
 Corr. Area 1: 0.0

Region table for sample 5 : RO_20

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,200	0.0	0	0	0.0	0.00	0.0	Blue

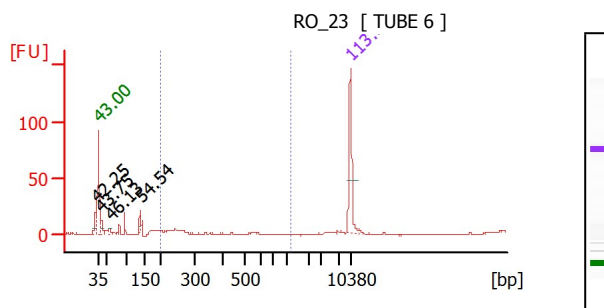
Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ioanalyzer\2100 expert\data\2016-05-20\CNelson_1-9and36.xad

Created: 5/20/2016 1:32:18 PM
 Modified: 5/20/2016 3:28:28 PM

Electropherogram Summary Continued ...

Peak table for sample 5 : RO_20

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	32	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	41	11.97	444.6	
4	10,380	75.00	10.9	Upper Marker
5	50,107	0.00	0.0	
6	51,919	0.00	0.0	



Overall Results for sample 6 : RO_23

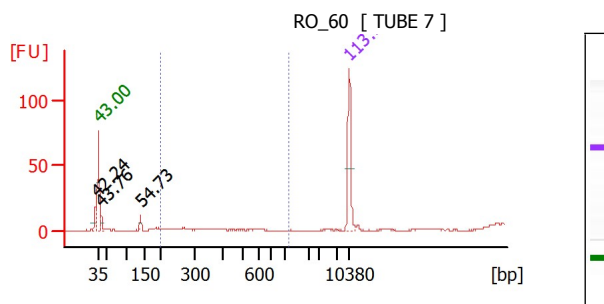
Number of peaks found: 4
 Noise: 0.3
 Corr. Area 1: 19.4

Region table for sample 6 : RO_23

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,200	19.4	16	239	8.4	21.08	133.7	Blue

Peak table for sample 6 : RO_23

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	32	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	40	22.37	852.5	
4	57	8.84	236.1	
5	137	20.73	228.9	
6	10,380	75.00	10.9	Upper Marker



Overall Results for sample 7 : RO_60

Number of peaks found: 3
 Noise: 0.4
 Corr. Area 1: 2.4

Region table for sample 7 : RO_60

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,200	2.4	4	251	8.4	2.74	16.5	Blue

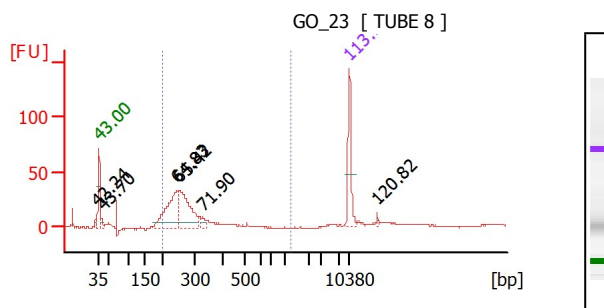
Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ioanalyzer\2100 expert\data\2016-05-20\CNelson_1-9and36.xad

Created: 5/20/2016 1:32:18 PM
 Modified: 5/20/2016 3:28:28 PM

Electropherogram Summary Continued ...

Peak table for sample 7 : RO_60

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	32	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	40	18.06	687.8	
4	139	12.59	137.1	
5	10,380	75.00	10.9	Upper Marker



Overall Results for sample 8 : GO_23

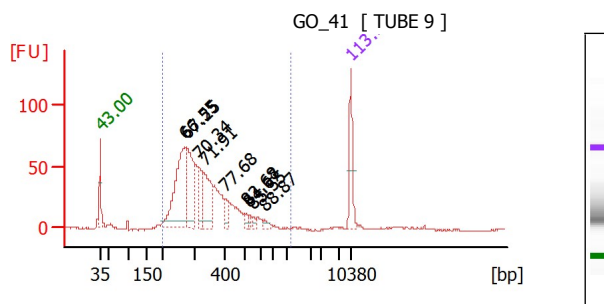
Number of peaks found: 6
 Noise: 0.5
 Corr. Area 1: 363.5

Region table for sample 8 : GO_23

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,200	363.5	80	270	18.6	343.58	1,984.5	Blue

Peak table for sample 8 : GO_23

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	32	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	39	11.18	429.5	
4	247	175.05	1,072.6	
5	254	183.07	1,093.5	
6	328	12.24	56.6	
7	10,380	75.00	10.9	Upper Marker
8	18,911	0.00	0.0	



Overall Results for sample 9 : GO_41

Number of peaks found: 9
 Noise: 0.6
 Corr. Area 1: 1,119.1

Region table for sample 9 : GO_41

From [bp]	To [bp]	Corr. Area	% of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200	1,200	1,119.1	96	336	28.5	1,154.05	5,670.8	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ioanalyzer\2100 expert\data\2016-05-20\CNelson_1-9and36.xad

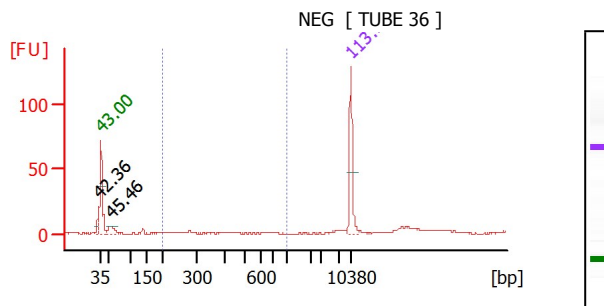
Created: 5/20/2016 1:32:18 PM
 Modified: 5/20/2016 3:28:28 PM

Electropherogram Summary Continued ...

Peak table for sample 9 :

GO 41

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	266	356.21	2,029.1	
3	274	217.77	1,205.8	
4	308	93.53	459.5	
5	328	159.29	735.9	
6	400	39.16	148.5	
7	505	10.09	30.3	
8	527	8.85	25.5	
9	546	10.83	30.1	
10	627	10.20	24.7	
11	10,380	75.00	10.9	Upper Marker



Overall Results for sample 10 : NEG

Number of peaks found: 2
 Noise: 0.4
 Corr. Area 1: 3.1

Region table for sample 10 : NEG

From To [bp]	Area	Corr. % of Total	Average Size [bp]	Size distribution in CV [%]	Conc. [pg/μl]	Molarity [pmol/l]	Color
200 1,000	3.1	4	382	42.5	3.48	16.8	Blue

Peak table for sample 10 :

NEG

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	33	0.00	0.0	
2	35	125.00	5,411.3	Lower Marker
3	51	33.34	995.9	
4	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ioanalyzer\2100 expert\data\2016-05-20\CNelson_1-9and36.xad

Created: 5/20/2016 1:32:18 PM
Modified: 5/20/2016 3:28:28 PM

Gel Image