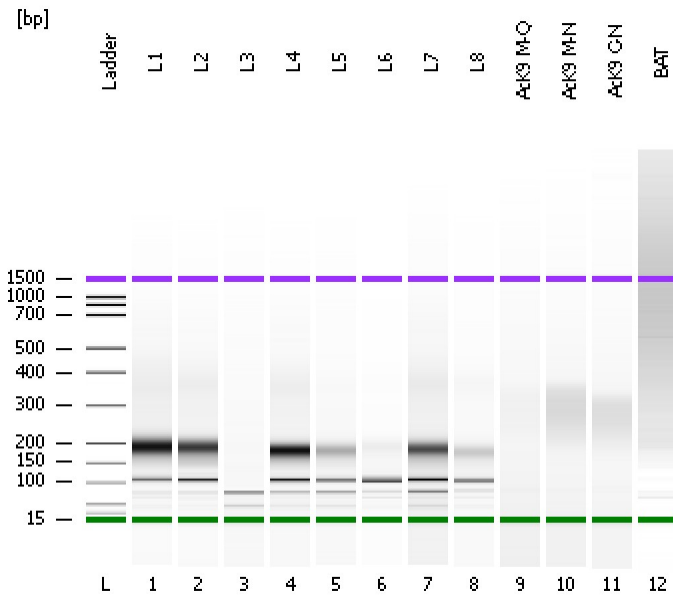


Assay Class: DNA 1000
Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
Modified: 6/22/2009 6:06:53 PM

Electrophoresis File Run Summary



Instrument Information:

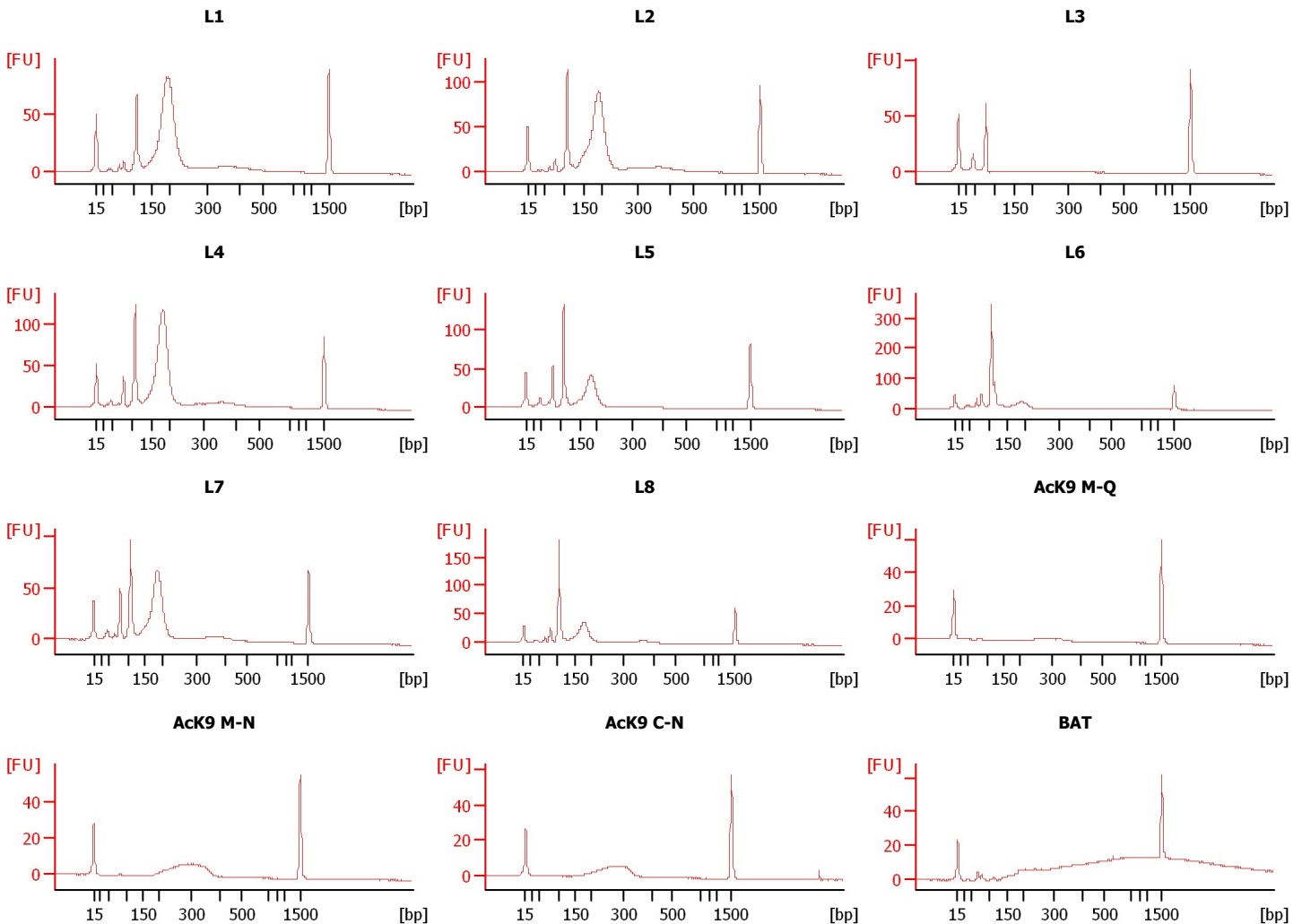
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

Assay Information:

Assay Origin Path: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\assays\dsDNA\DNA 1000 Series II.xsy
Title: DNA 1000 Series II
Version: 2.1
Assay Comments: © Copyright 2003-2007 Agilent Technologies

Chip Information:

Chip Lot:
Reagent Kit Lot:
Chip Comments:



Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
L1		<input type="checkbox"/>	✓			
L2		<input type="checkbox"/>	✓			
L3		<input type="checkbox"/>	✓			
L4		<input type="checkbox"/>	✓			
L5		<input type="checkbox"/>	✓			
L6		<input type="checkbox"/>	✓			
L7		<input type="checkbox"/>	✓			
L8		<input type="checkbox"/>	✓			
Ack9 M-Q		<input type="checkbox"/>	✓			
Ack9 M-N		<input type="checkbox"/>	✓			
Ack9 C-N		<input type="checkbox"/>	✓			
BAT		<input type="checkbox"/>	✓			
Chip Lot #				Reagent Kit Lot #		

Chip Comments :

Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electrophoresis Assay Details

General Analysis Settings

Number of Available Sample and Ladder Wells (Max.) : 13
 Minimum Visible Range [s] : 30
 Maximum Visible Range [s] : 129
 Start Analysis Time Range [s] : 30
 End Analysis Time Range [s] : 128.95
 Ladder Concentration [ng/μl] : 44
 Uses Standard Area for Ladder Fragments
 Lower Marker Concentration [ng/μl] : 4.2
 Upper Marker Concentration [ng/μl] : 2.1
 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] : 30
 Integration End Time [s] : 128.95
 Slope Threshold : 0.5
 Height Threshold [FU] : 20
 Area Threshold : 0.1
 Width Threshold [s] : 0.5
 Baseline Plateau [s] : 0.5

Filter Settings

Filter Width [s] : 0.5
 Polynomial Order : 4

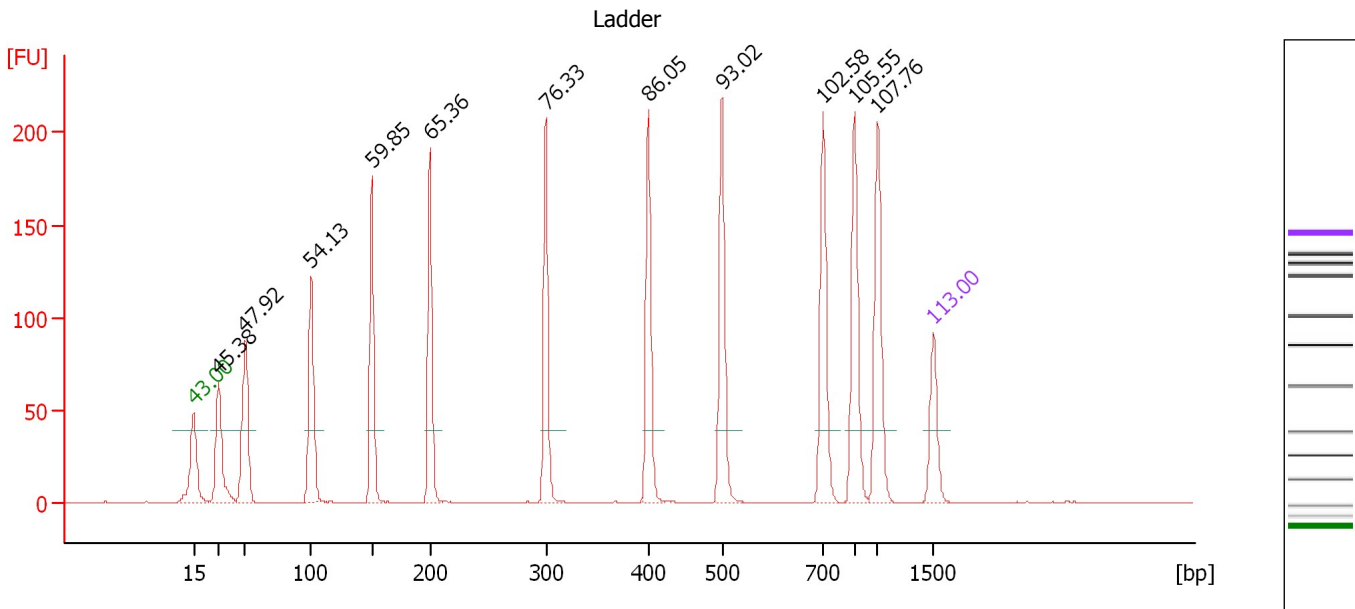
Ladder

Ladder Peak	Size	Area
1	15	25
2	25	26
3	50	34
4	100	41
5	150	45
6	200	52
7	300	63
8	400	76
9	500	83
10	700	88
11	850	86
12	1000	90
13	1500	52

Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary



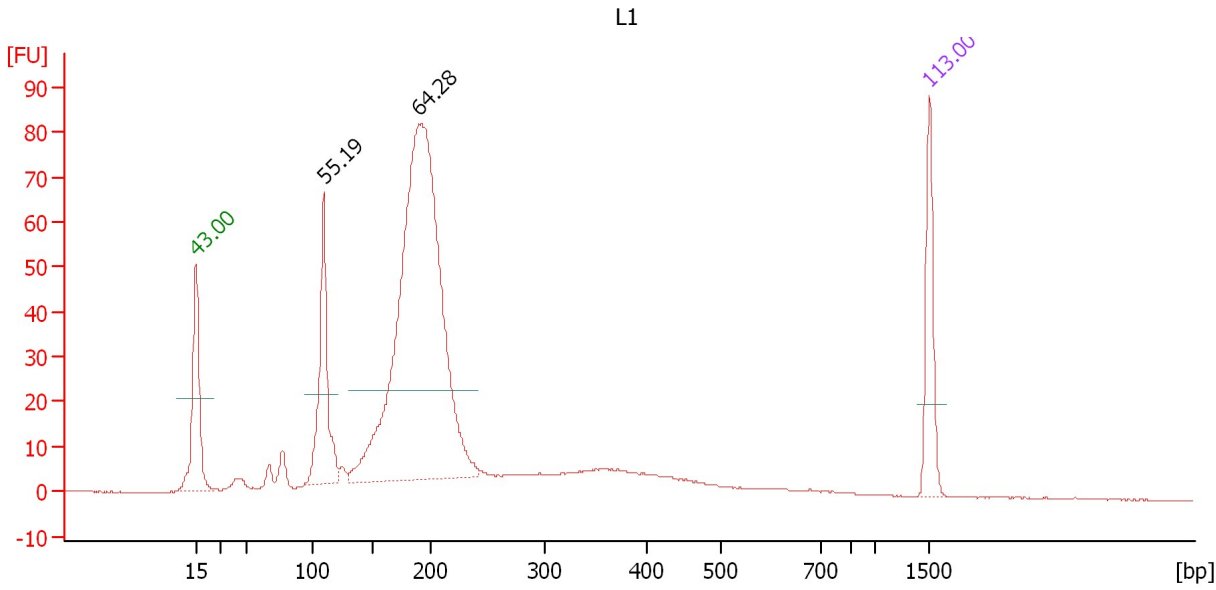
Peak table for Ladder

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	25	4.00	242.4	Ladder Peak
3	50	4.00	121.2	Ladder Peak
4	100	4.00	60.6	Ladder Peak
5	150	4.00	40.4	Ladder Peak
6	200	4.00	30.3	Ladder Peak
7	300	4.00	20.2	Ladder Peak
8	400	4.00	15.2	Ladder Peak
9	500	4.00	12.1	Ladder Peak
10	700	4.00	8.7	Ladder Peak
11	850	4.00	7.1	Ladder Peak
12	1,000	4.00	6.1	Ladder Peak
13	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : L1

Number of peaks found: 2

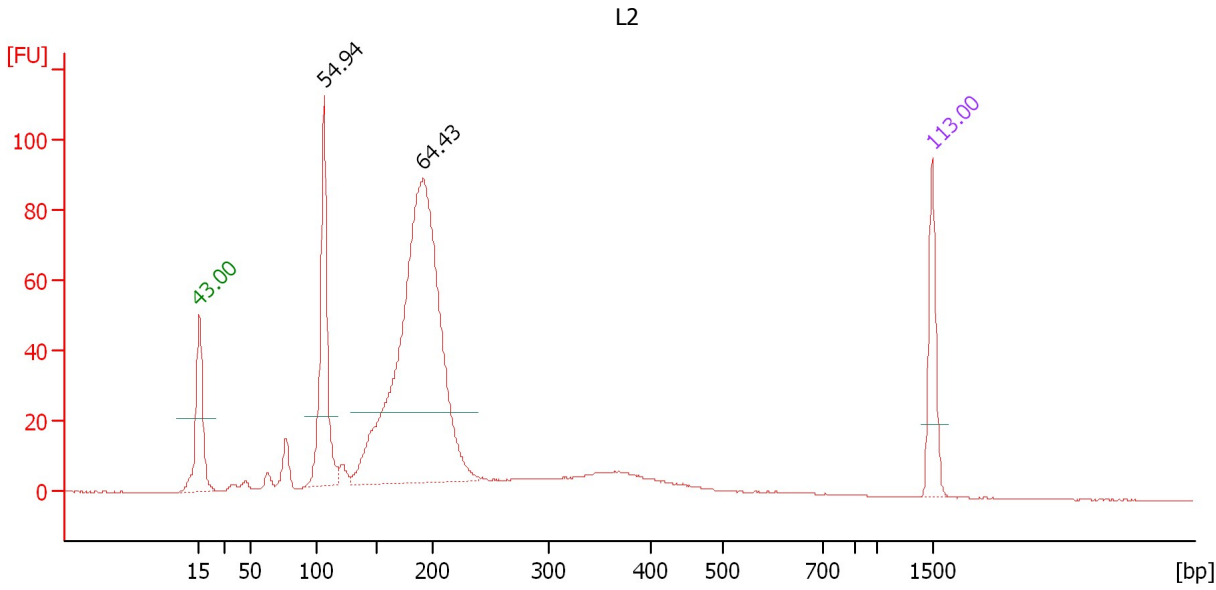
Peak table for sample 1 : L1

Peak	Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	109	3.85	53.3	
3	190	21.58	171.9	
4	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : L2

Number of peaks found: 2

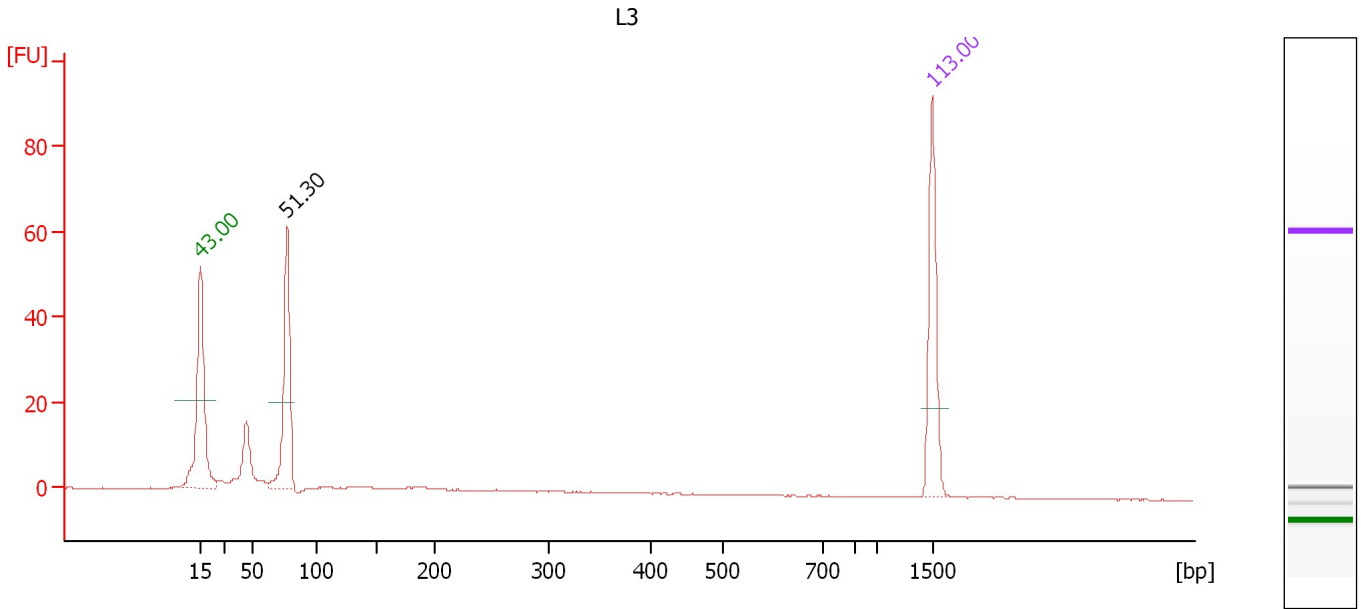
Peak table for sample 2 : L2

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	107	5.37	76.0	
3	192	21.08	166.8	
4	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : L3

Number of peaks found: 1

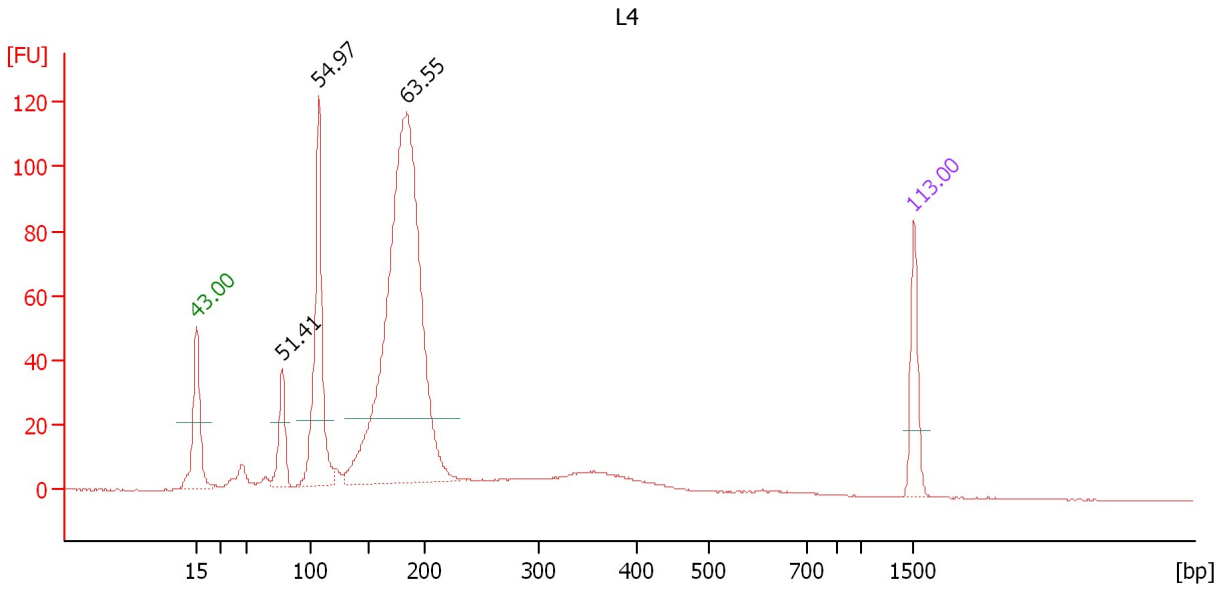
Peak table for sample 3 : L3

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	77	2.97	58.4	
3	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : L4

Number of peaks found: 3

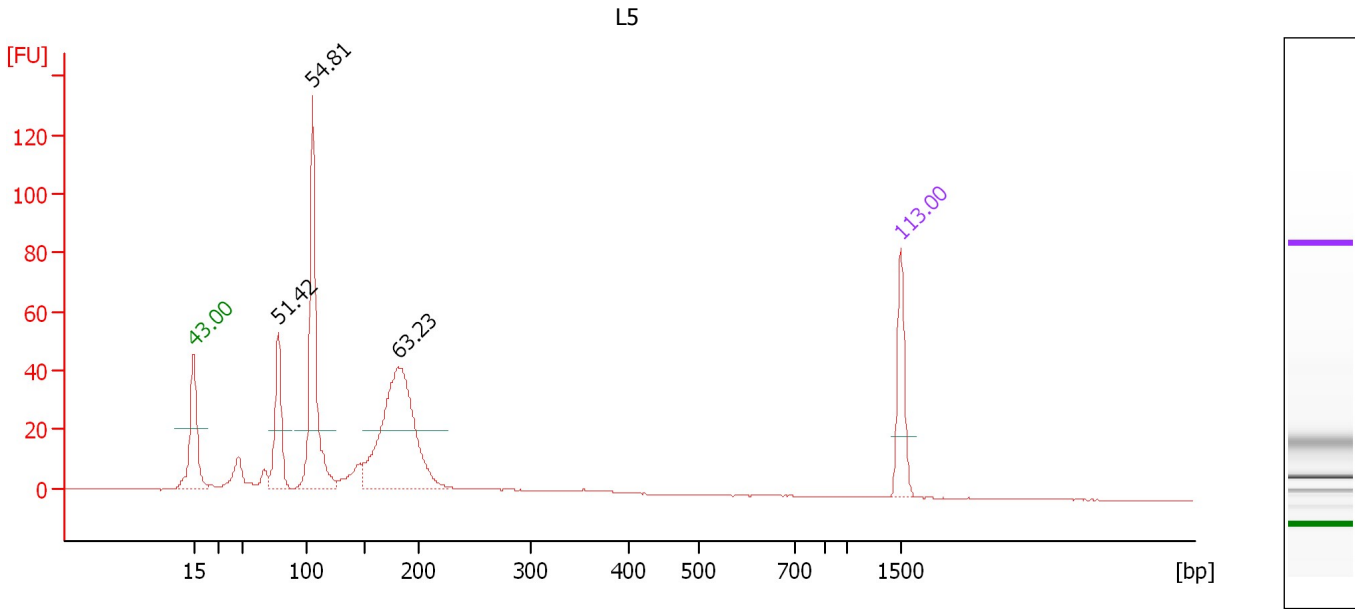
Peak table for sample 4 : L4

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	78	2.04	39.6	
3	107	7.00	98.7	
4	184	26.30	217.1	
5	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : L5

Number of peaks found: 3

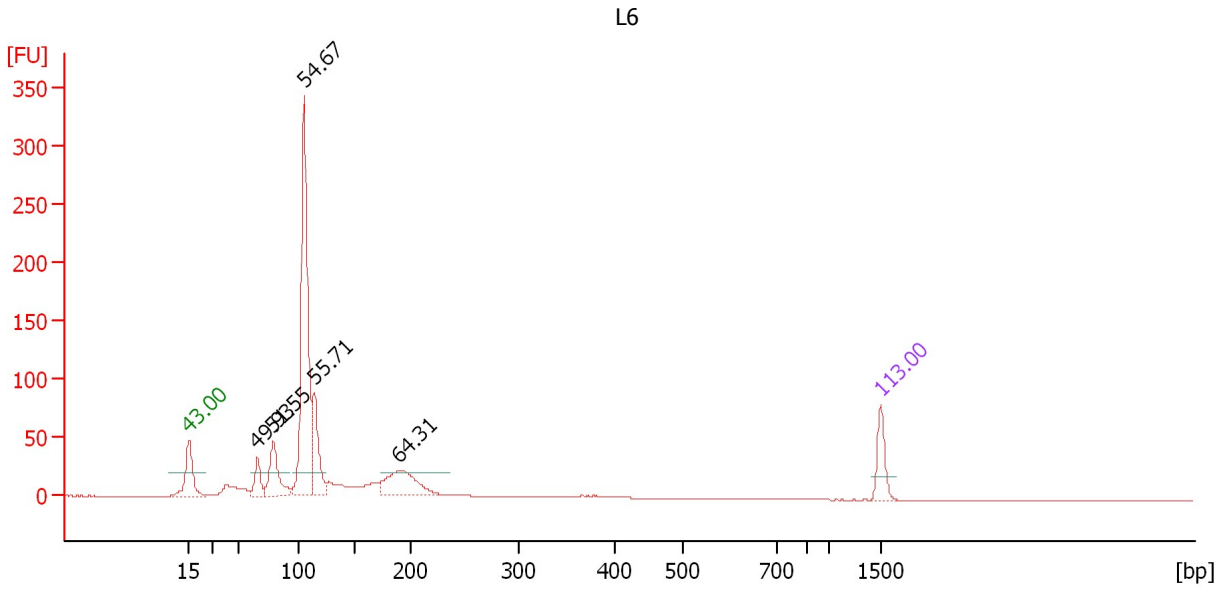
Peak table for sample 5 : L5

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	78	3.02	58.5	
3	106	6.92	99.0	
4	181	9.42	79.0	
5	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : L6

Number of peaks found: 5

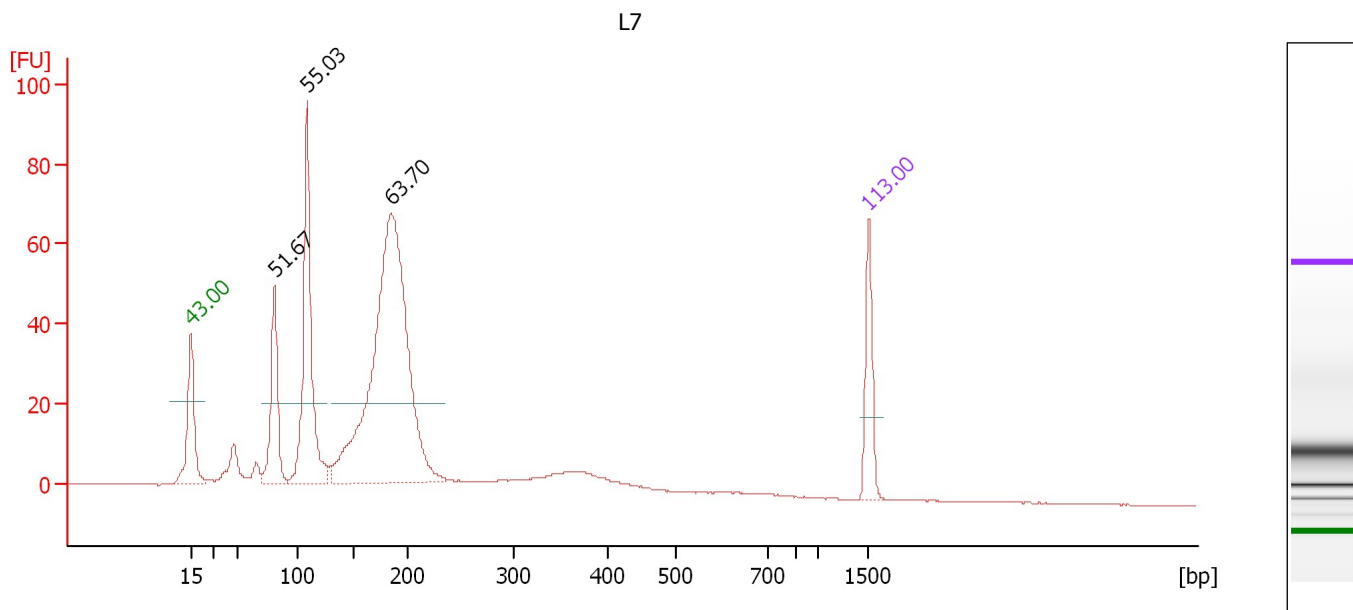
Peak table for sample 6 : L6

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	66	1.73	39.6	
3	79	3.85	73.6	
4	105	18.64	269.7	
5	114	4.43	58.9	
6	190	4.62	36.7	
7	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : L7

Number of peaks found: 3

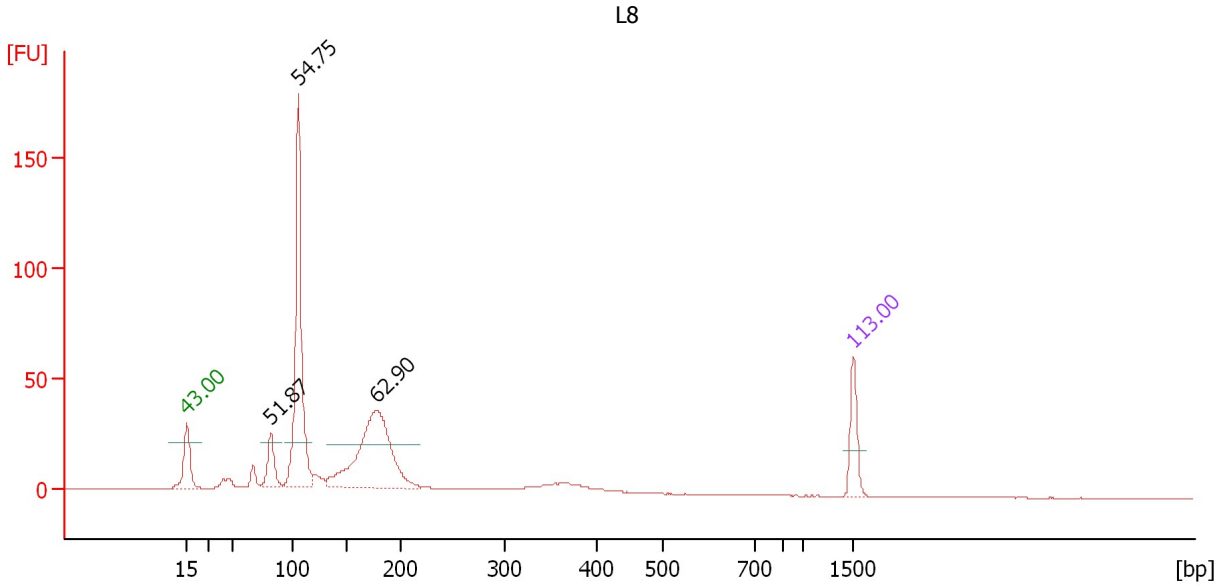
Peak table for sample 7 : L7

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	80	3.79	71.6	
3	108	7.60	106.7	
4	185	20.70	169.6	
5	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : L8

Number of peaks found: 3

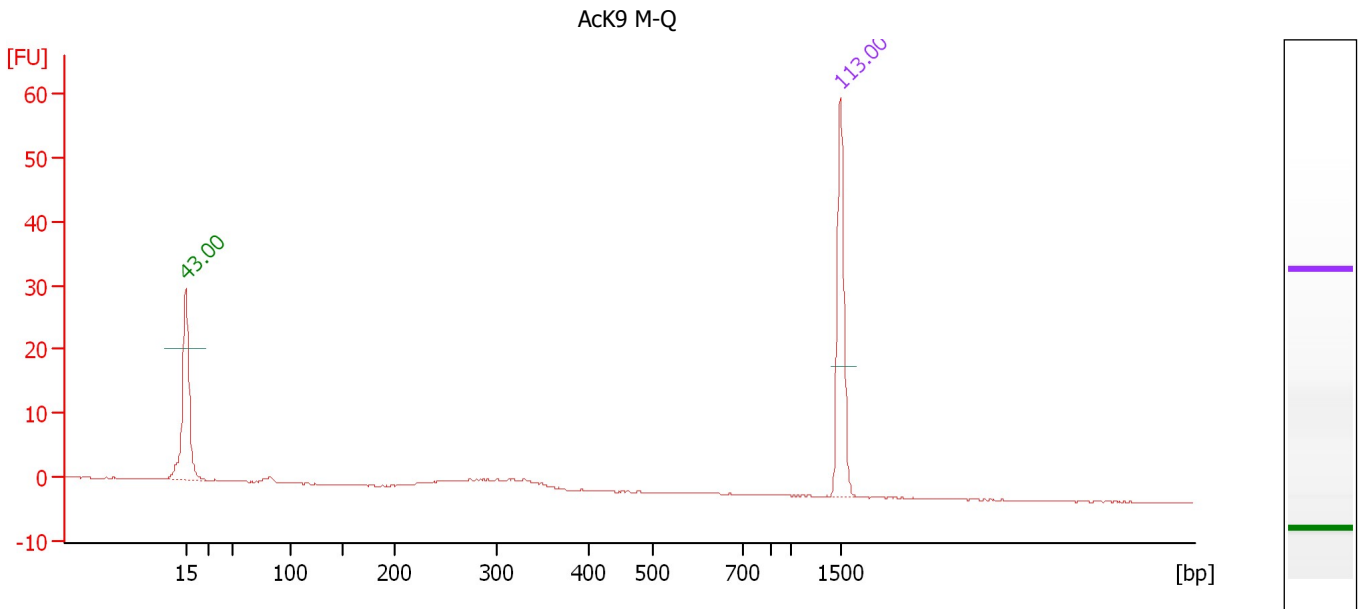
Peak table for sample 8 : L8

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	82	2.09	38.6	
3	105	12.25	176.0	
4	178	11.47	97.8	
5	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : Ack9 M-Q

Number of peaks found: 0

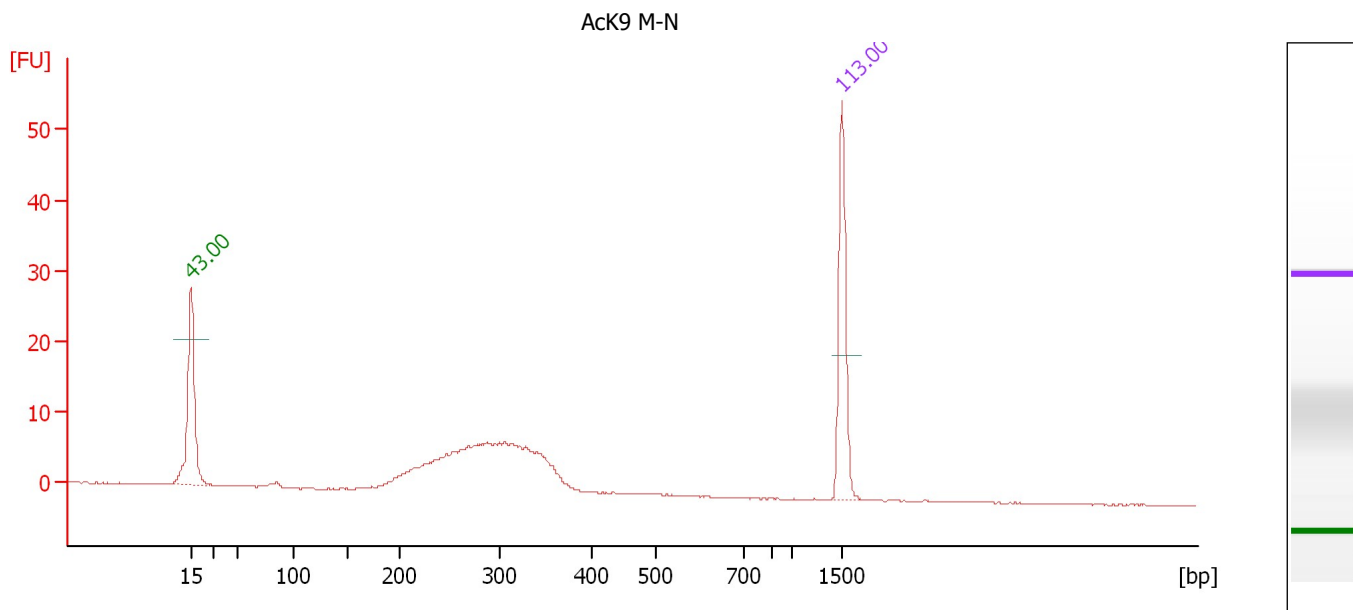
Peak table for sample 9 : Ack9 M-Q

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : Ack9 M-N

Number of peaks found: 0

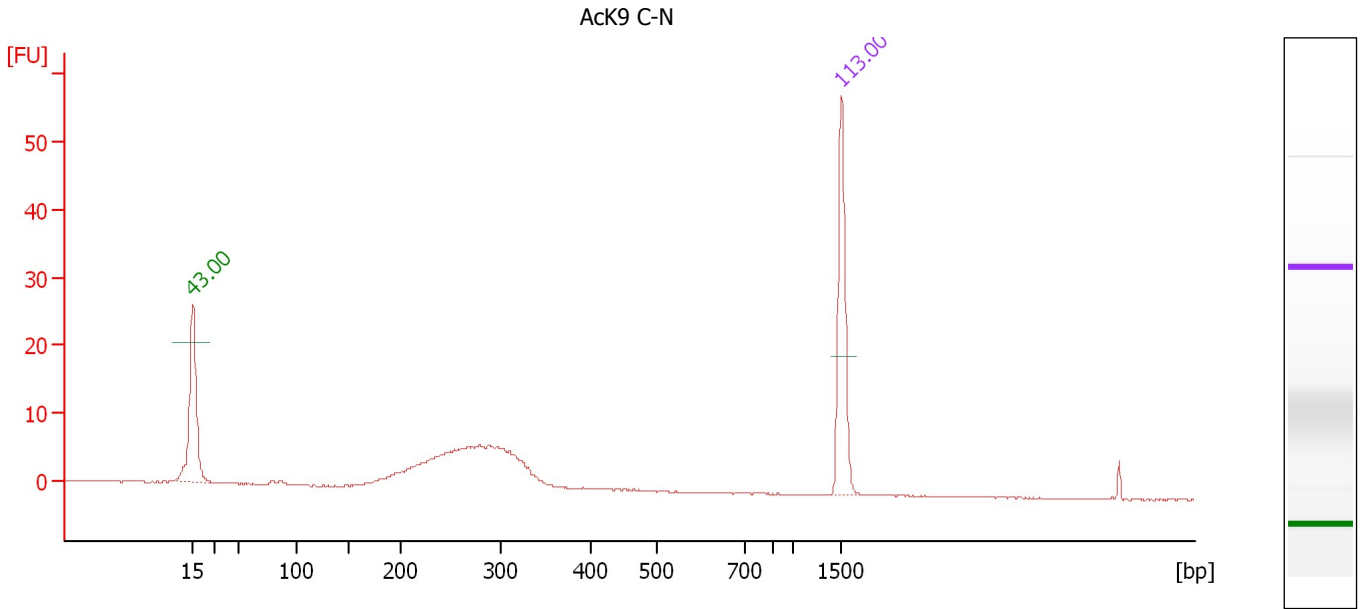
Peak table for sample 10 : Ack9 M-N

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : Ack9 C-N

Number of peaks found: 0

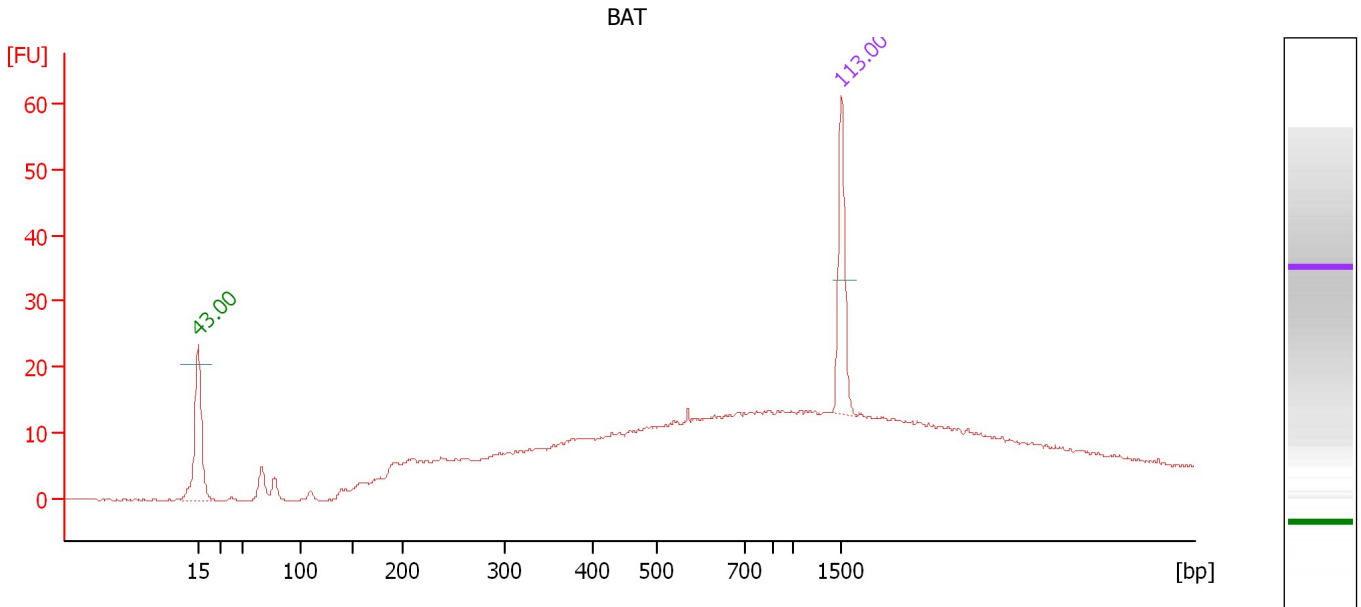
Peak table for sample 11 : Ack9 C-N

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
Modified: 6/22/2009 6:06:53 PM

Electropherogram Summary Continued ...



Overall Results for sample 12 : BAT

Number of peaks found: 0

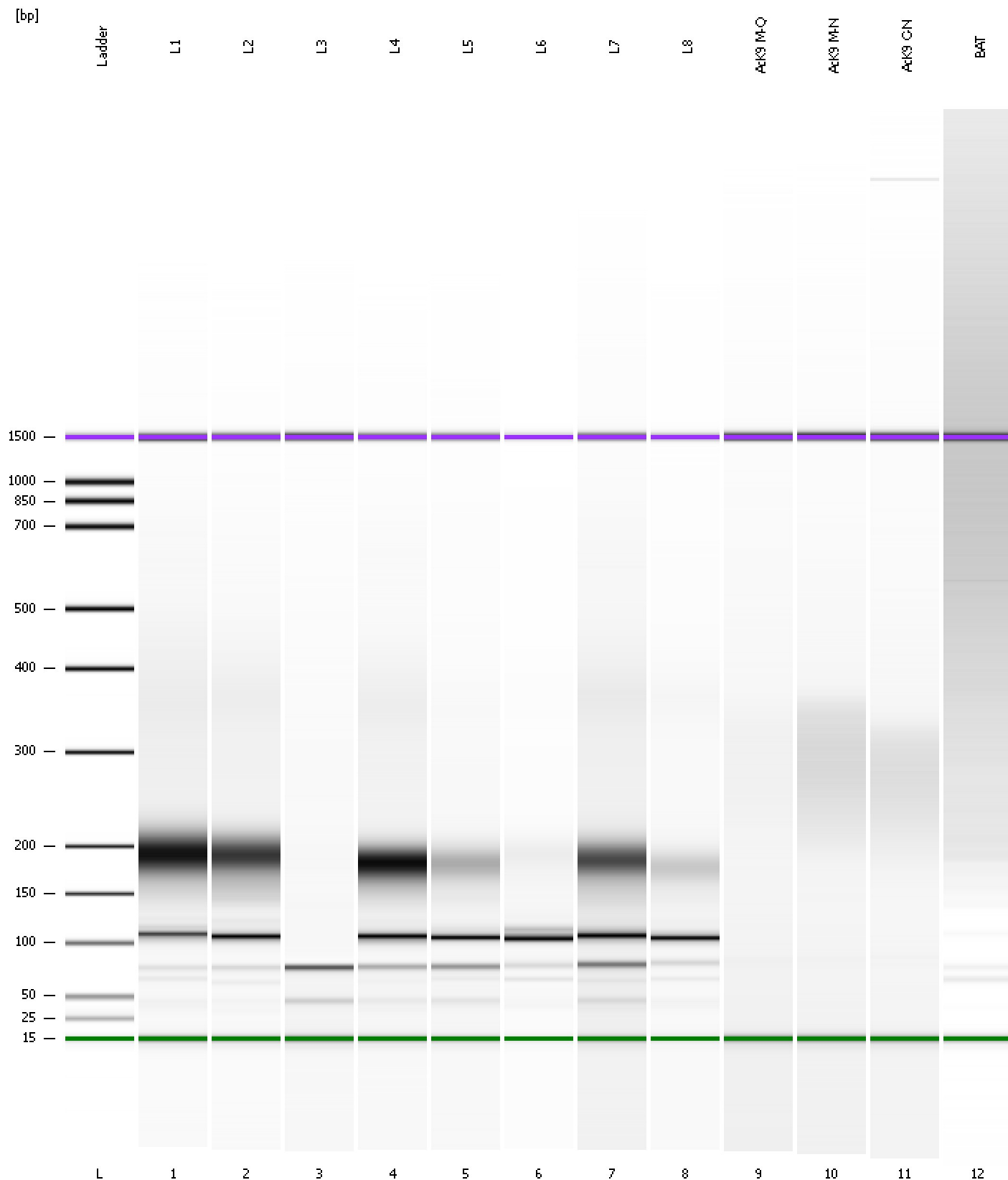
Peak table for sample 12 : BAT

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	15	4.20	424.2	Lower Marker
2	1,500	2.10	2.1	Upper Marker

Assay Class: DNA 1000
Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
Modified: 6/22/2009 6:06:53 PM

Gel Image

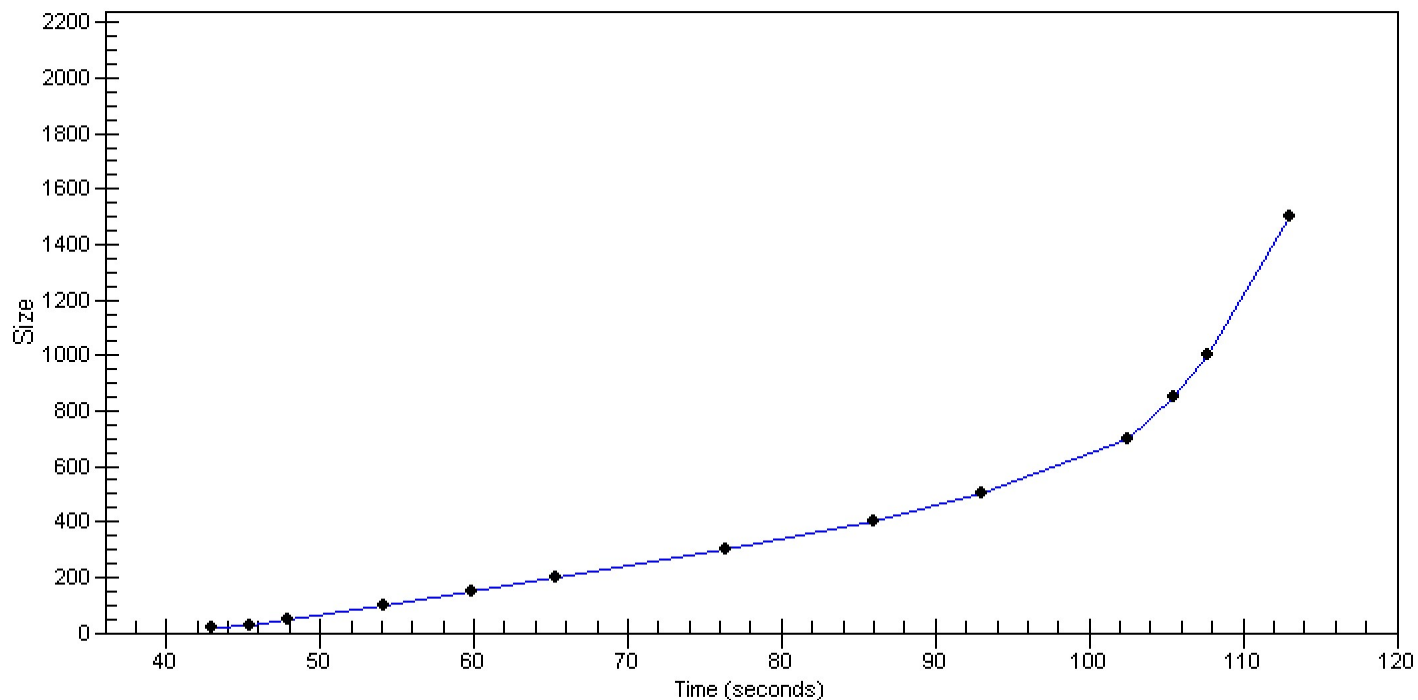


Assay Class: DNA 1000
Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
Modified: 6/22/2009 6:06:53 PM

Curves

Standard Curve



Assay Class: DNA 1000
 Data Path: C:\...Files\2009-06-22\Bioanalyzer_090622_facc_farn_sinha.xad.xad

Created: 6/22/2009 5:24:46 PM
 Modified: 6/22/2009 6:06:53 PM

Run Logbook

Description	Number	Source	Category	Sub Category	Time Stamp	User	Host
Run ended on port 4 (Number of wells acquired: 13)		Instrument	Run		6/22/2009 6:06:37 PM	cnicolet	NANODROP
Run Started on port 4 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\XAD Files\2009-06-22\Bioanalyzer_2009-06-22_17-24-46.xad)		Instrument	Run		6/22/2009 5:24:52 PM	cnicolet	NANODROP
Product Number : G2938B		Instrument	Run		6/22/2009 5:24:52 PM	cnicolet	NANODROP
Name :		Instrument	Run		6/22/2009 5:24:52 PM	cnicolet	NANODROP
Vendor : Agilent Technologies		Instrument	Run		6/22/2009 5:24:52 PM	cnicolet	NANODROP
Serial# : DE13701086		Instrument	Run		6/22/2009 5:24:52 PM	cnicolet	NANODROP
Firmware : C.01.069		Instrument	Run		6/22/2009 5:24:52 PM	cnicolet	NANODROP
Cartridge : Electrode		Instrument	Run		6/22/2009 5:24:52 PM	cnicolet	NANODROP