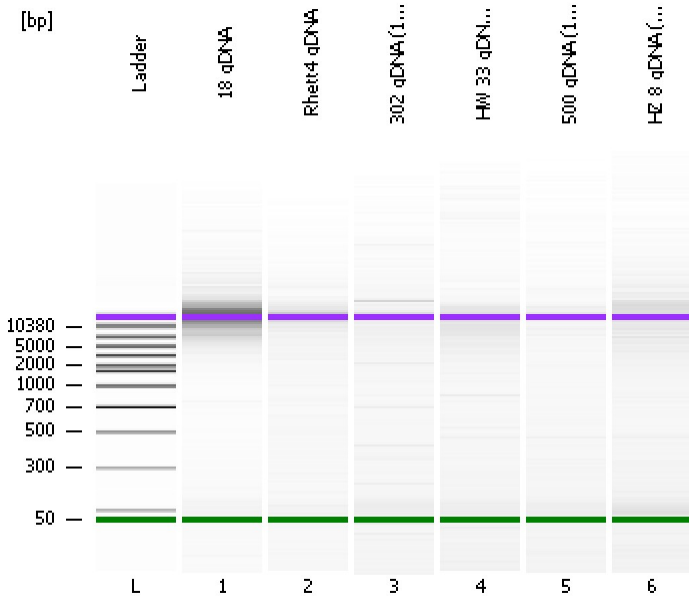


Assay Class: DNA 12000
Data Path: C:\...d Settings\Bioanalyzer\2014-09-09\2014-09-09_001_Coaker.xad

Created: 9/9/2014 9:44:03 AM
Modified: 9/9/2014 10:18:36 AM

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 12000 Series II.xsy

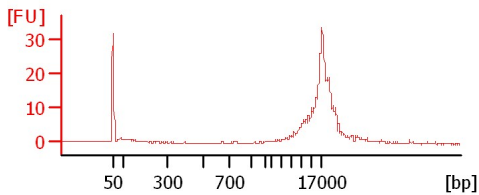
Assay Class: DNA 12000
Version: 2.4
Assay Comments: DNA Analysis 100 -12000 bp

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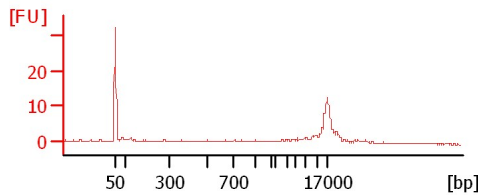
Chip Information:

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

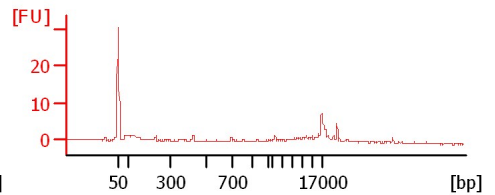
18 gDNA



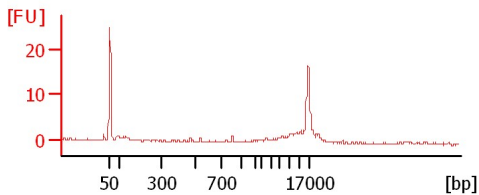
Rhett4 gDNA



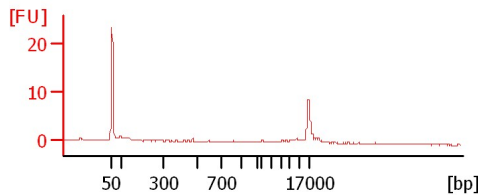
302 gDNA (1:2)



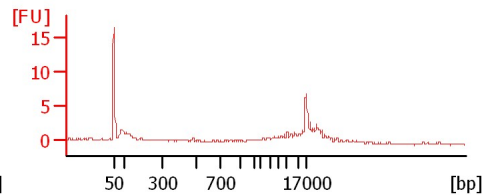
HW 33 gDNA (1:3)



500 gDNA (1:3)



HZ 8 gDNA (1:2)



Assay Class: DNA 12000
Data Path: C:\...d Settings\Bioanalyzer\2014-09-09\2014-09-09_001_Coaker.xad

Created: 9/9/2014 9:44:03 AM
Modified: 9/9/2014 10:18:36 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
18 gDNA		<input type="checkbox"/>	✓			
Rhett4 gDNA		<input type="checkbox"/>	✓			
302 gDNA (1:2)		<input type="checkbox"/>	✓			
HW 33 gDNA (1:3)		<input type="checkbox"/>	✓			
500 gDNA (1:3)		<input type="checkbox"/>	✓			
HZ 8 gDNA (1:2)		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

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

Assay Class: DNA 12000
Data Path: C:\...d Settings\Bioanalyzer\2014-09-09\2014-09-09_001_Coaker.xad

Created: 9/9/2014 9:44:03 AM
Modified: 9/9/2014 10:18:36 AM

Electrophoresis Assay Details

General Analysis Settings

Number of Available Sample and Ladder Wells (Max.) : 13
Minimum Visible Range [s] : 20
Maximum Visible Range [s] : 99
Start Analysis Time Range [s] : 20
End Analysis Time Range [s] : 98.95
Ladder Concentration [ng/μl] : 44
Uses Standard Area for Ladder Fragments
Lower Marker Concentration [ng/μl] : 8.3
Upper Marker Concentration [ng/μl] : 4.2
Used Upper Marker for Quantitation
This is a Qualitative Assay Only
Standard Curve Fit is Point to Point
Show Data Aligned to Lower and Upper Marker

Integrator Settings

Integration Start Time [s] : 20
Integration End Time [s] : 98.95
Slope Threshold : 0.8
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

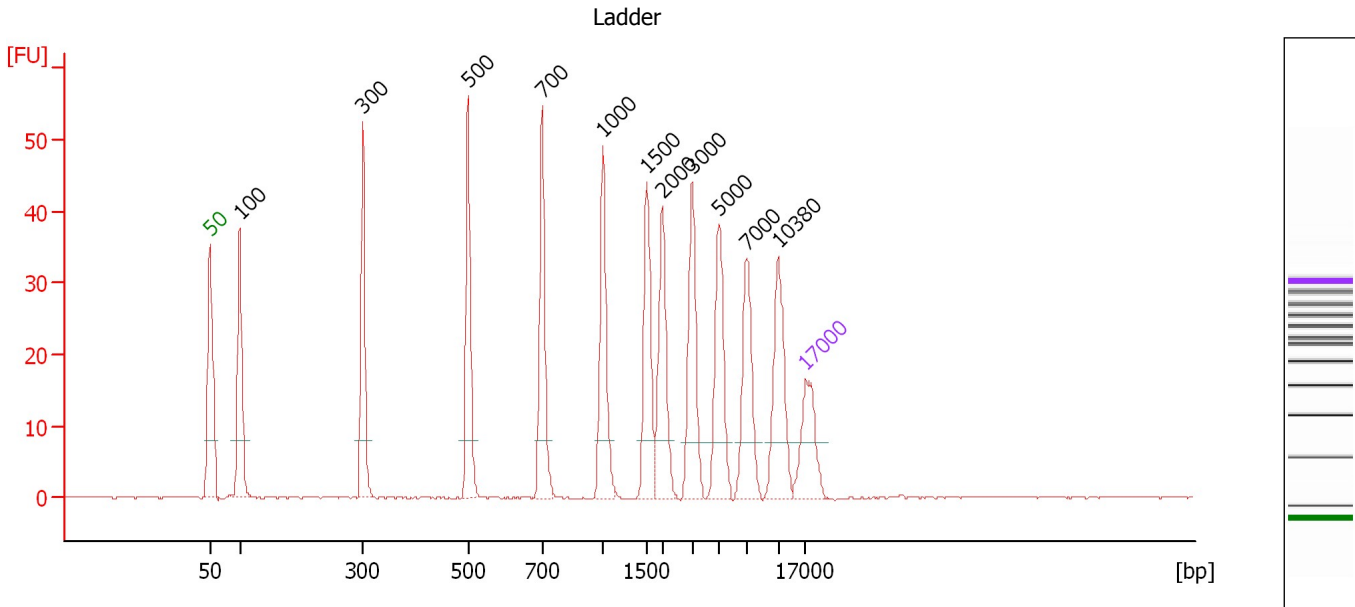
Ladder

Ladder Peak	Size	Area
1	50	120
2	100	48
3	300	61
4	500	77
5	700	81
6	1000	86
7	1500	92
8	2000	92
9	3000	96
10	5000	100
11	7000	98
12	10380	102
13	17000	110

Assay Class: DNA 12000
 Data Path: C:\...d Settings\Bioanalyzer\2014-09-09\2014-09-09_001_Coaker.xad

Created: 9/9/2014 9:44:03 AM
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Electropherogram Summary



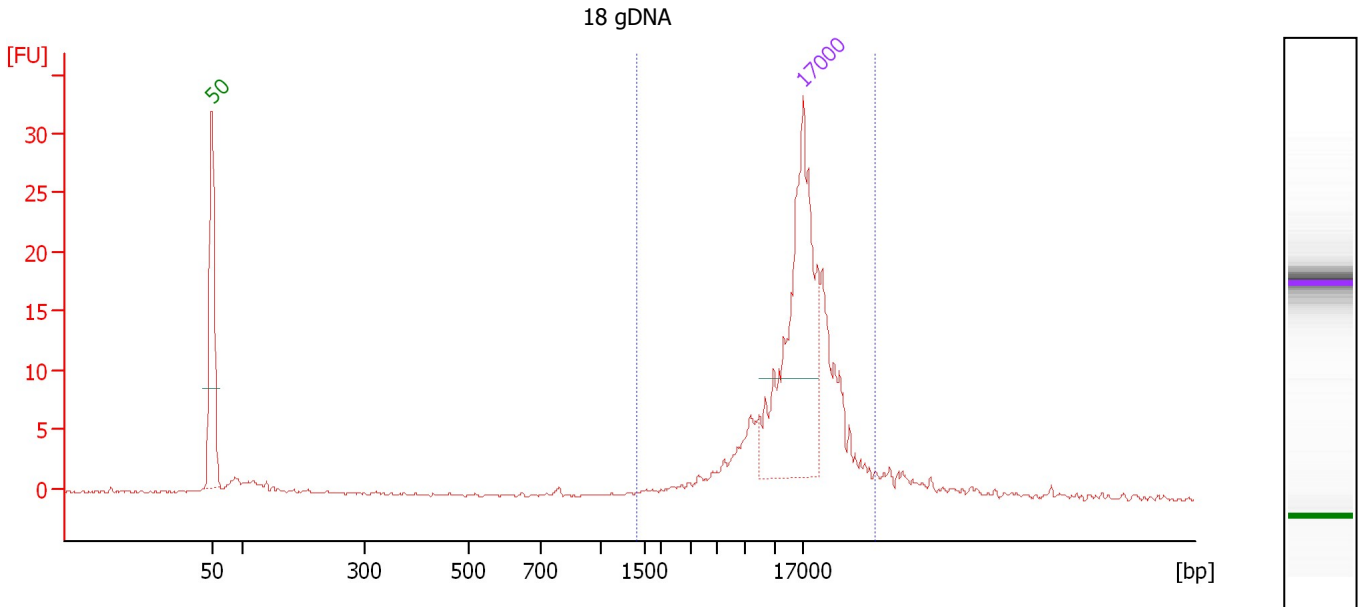
Peak table for Ladder

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	100	4.00	60.6	Ladder Peak
3	300	4.00	20.2	Ladder Peak
4	500	4.00	12.1	Ladder Peak
5	700	4.00	8.7	Ladder Peak
6	1,000	4.00	6.1	Ladder Peak
7	1,500	4.00	4.0	Ladder Peak
8	2,000	4.00	3.0	Ladder Peak
9	3,000	4.00	2.0	Ladder Peak
10	5,000	4.00	1.2	Ladder Peak
11	7,000	4.00	0.9	Ladder Peak
12	10,380	4.00	0.6	Ladder Peak
13	17,000	4.20	0.4	Upper Marker

Assay Class: DNA 12000
 Data Path: C:\...d Settings\Bioanalyzer\2014-09-09\2014-09-09_001_Coaker.xad

Created: 9/9/2014 9:44:03 AM
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Electropherogram Summary Continued ...



Overall Results for sample 1 : 18 gDNA

Number of peaks found: 0 Area 1: 49.5

Peak table for sample 1 : 18 gDNA

Peak	Size [bp]	Conc. [ng/µl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

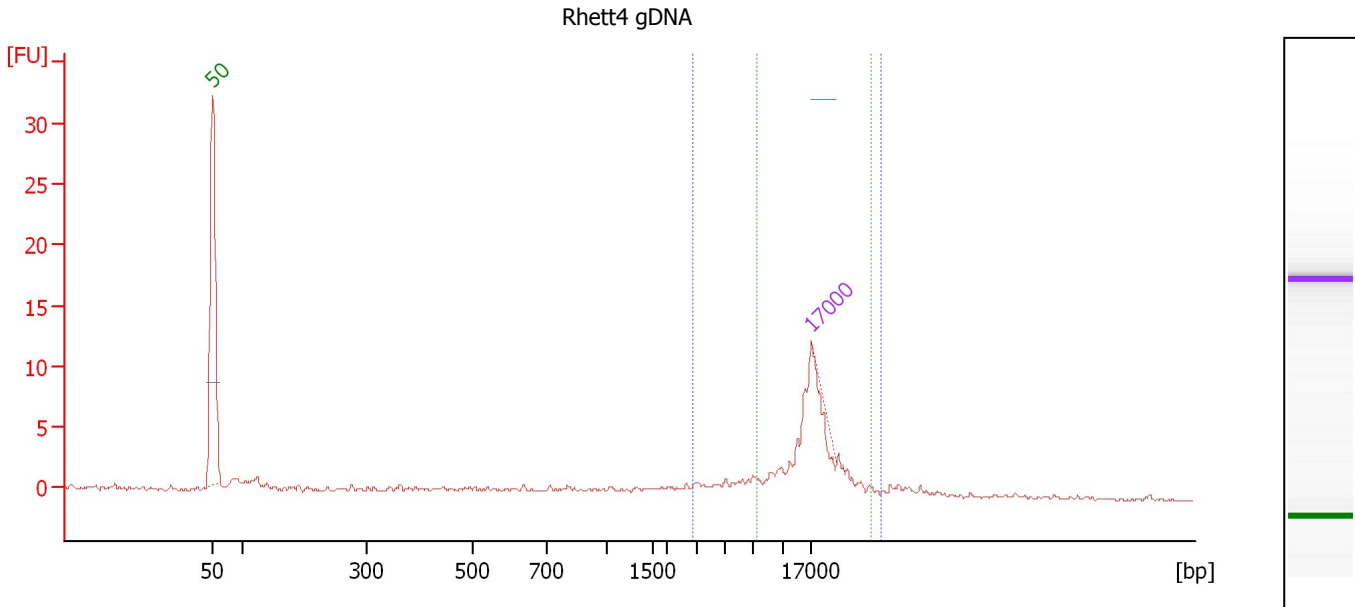
Region table for sample 1 : 18 gDNA

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/µl]	Area	% of Total	Size distribution in CV [%]	Color
1,393	34,379	16,382	3.42	49.5	72	41.5	Blue

Assay Class: DNA 12000
 Data Path: C:\...d Settings\Bioanalyzer\2014-09-09\2014-09-09_001_Coaker.xad

Created: 9/9/2014 9:44:03 AM
 Modified: 9/9/2014 10:18:36 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 : Rhett4 gDNA

Number of peaks found: 0 Area 2: 16.4
 Area 1: 21.5

Peak table for sample 2 : Rhett4 gDNA

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

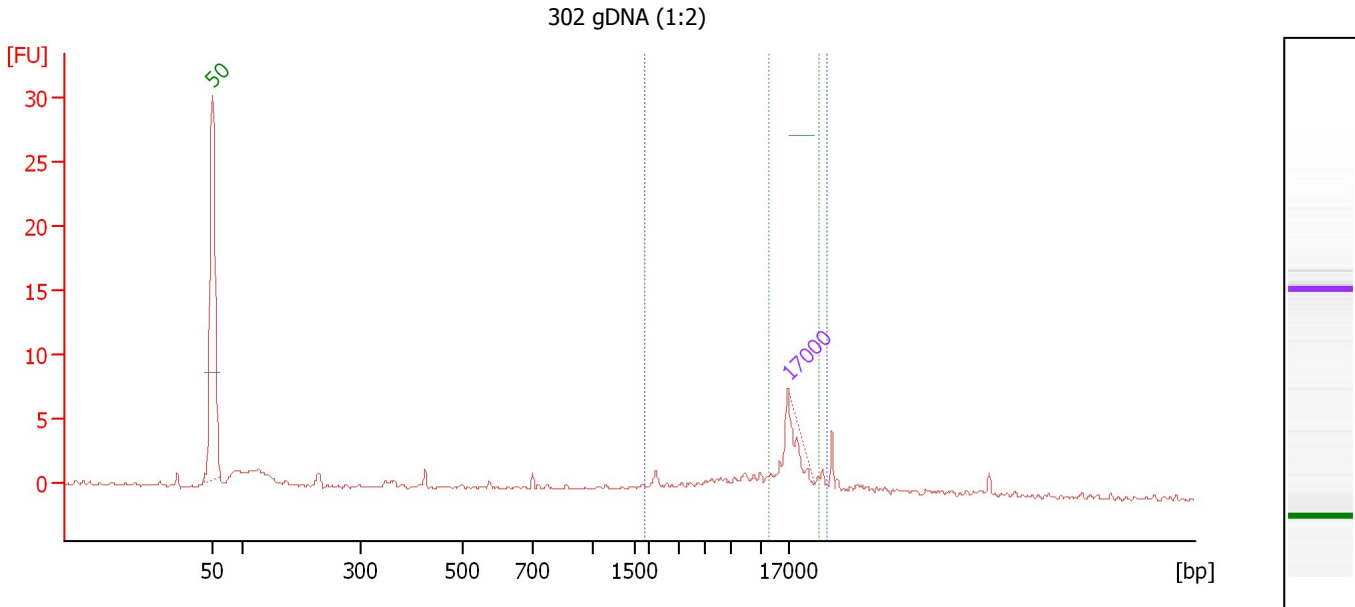
Region table for sample 2 : Rhett4 gDNA

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
2,836	33,599	15,804	118,606.87	21.5	54	40.5	Blue
7,532	31,140	17,336	89,385.31	16.4	41	28.2	Green

Assay Class: DNA 12000
 Data Path: C:\...d Settings\Bioanalyzer\2014-09-09\2014-09-09_001_Coaker.xad

Created: 9/9/2014 9:44:03 AM
 Modified: 9/9/2014 10:18:36 AM

Electropherogram Summary Continued ...



Overall Results for sample 3 : 302 gDNA (1:2)

Number of peaks found: 0 Area 2: 3.3
 Area 1: 10.7

Peak table for sample 3 : 302 gDNA (1:2)

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

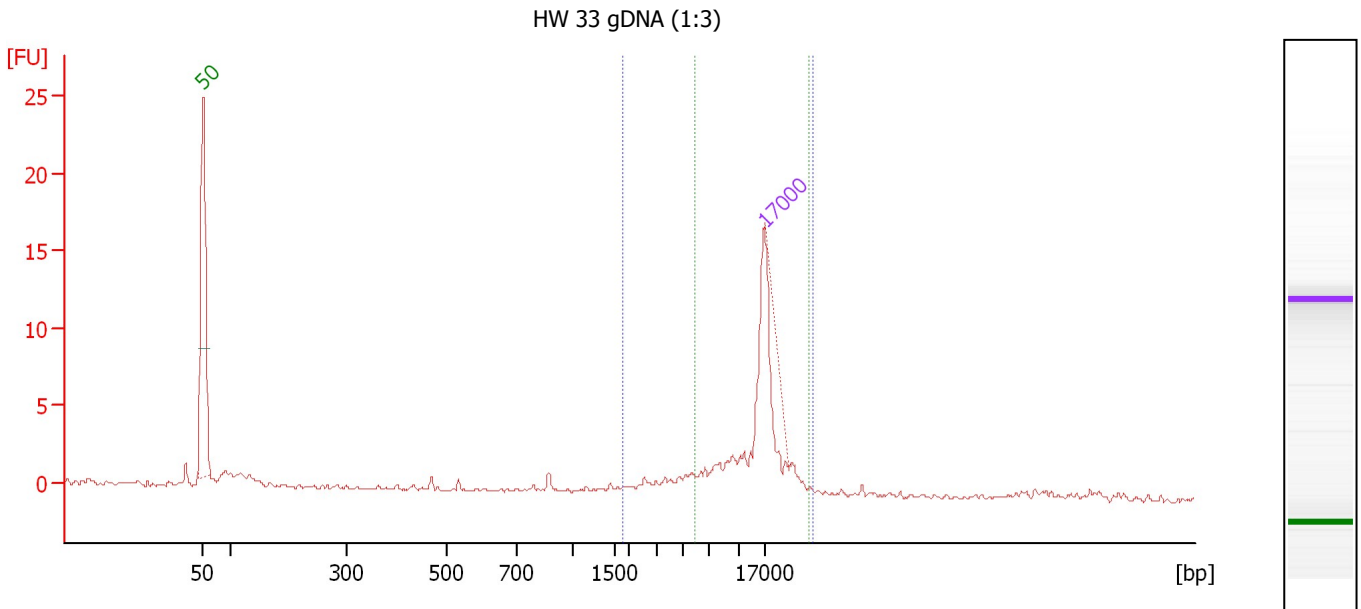
Region table for sample 3 : 302 gDNA (1:2)

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
1,836	26,771	13,589	0.00	10.7	49	49.0	Blue
12,433	24,723	17,899	0.00	3.3	15	14.7	Green

Assay Class: DNA 12000
 Data Path: C:\...d Settings\Bioanalyzer\2014-09-09\2014-09-09_001_Coaker.xad

Created: 9/9/2014 9:44:03 AM
 Modified: 9/9/2014 10:18:36 AM

Electropherogram Summary Continued ...



Overall Results for sample 4 : HW 33 gDNA (1:3)

Number of peaks found: 0 Area 2: 16.7
 Area 1: 19.9

Peak table for sample 4 : HW 33 gDNA (1:3)

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

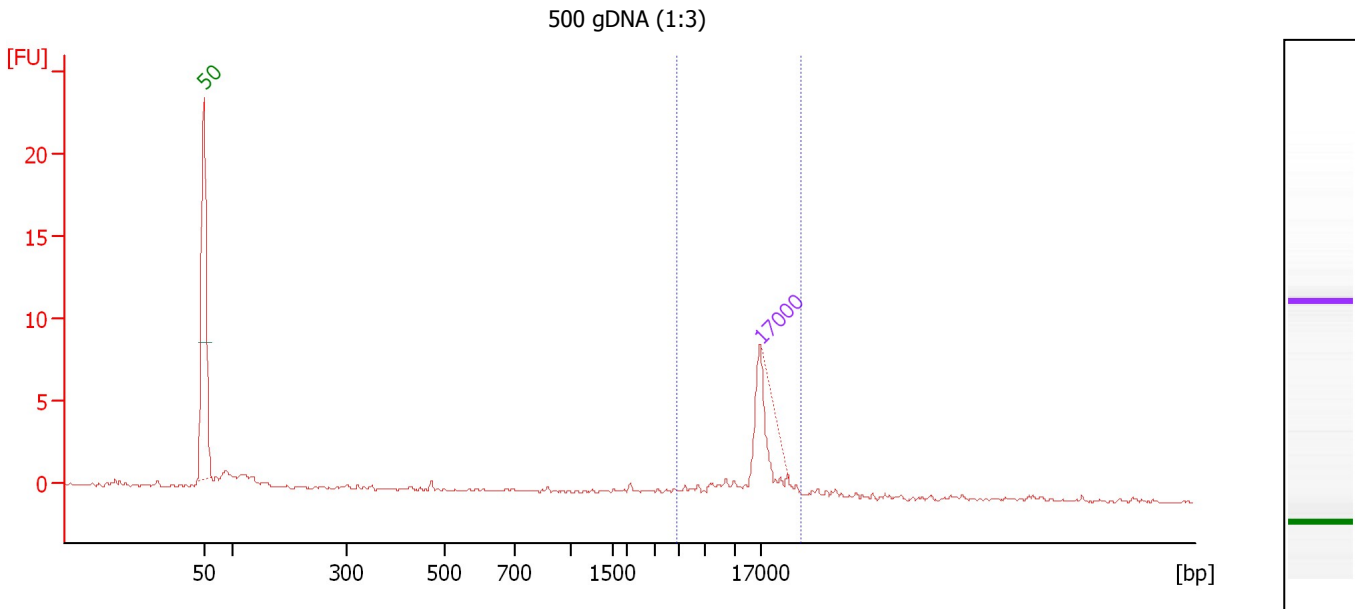
Region table for sample 4 : HW 33 gDNA (1:3)

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
1,795	29,085	14,264	0.00	19.9	73	40.1	Blue
5,858	28,037	15,455	0.00	16.7	62	29.8	Green

Assay Class: DNA 12000
 Data Path: C:\...d Settings\Bioanalyzer\2014-09-09\2014-09-09_001_Coaker.xad

Created: 9/9/2014 9:44:03 AM
 Modified: 9/9/2014 10:18:36 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : 500 gDNA (1:3)

Number of peaks found: 0 Area 1: 6.5

Peak table for sample 5 : 500 gDNA (1:3)

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

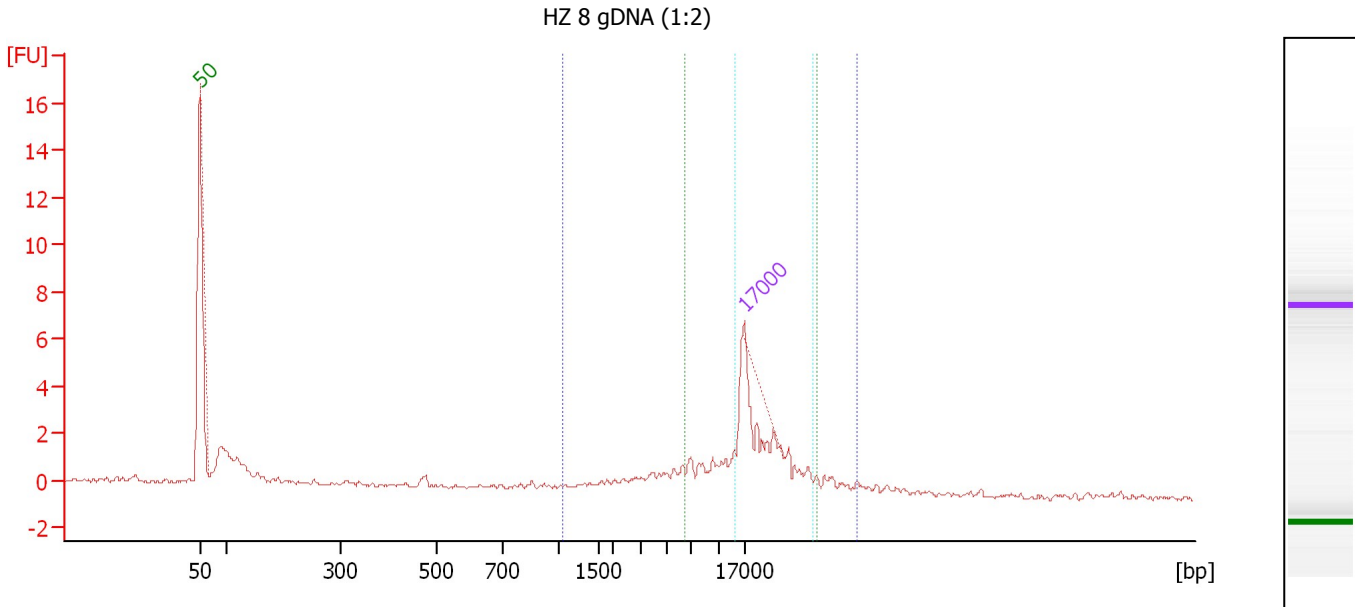
Region table for sample 5 : 500 gDNA (1:3)

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
4,836	27,111	16,025	0.00	6.5	52	27.9	Blue

Assay Class: DNA 12000
 Data Path: C:\...d Settings\Bioanalyzer\2014-09-09\2014-09-09_001_Coaker.xad

Created: 9/9/2014 9:44:03 AM
 Modified: 9/9/2014 10:18:36 AM

Electropherogram Summary Continued ...



Overall Results for sample 6 : HZ 8 gDNA (1:2)

Number of peaks found: 0 Area 2: 8.2
 Area 1: 13.5 Area 3: 3.9

Peak table for sample 6 : HZ 8 gDNA (1:2)

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	17,000	4.20	0.4	Upper Marker

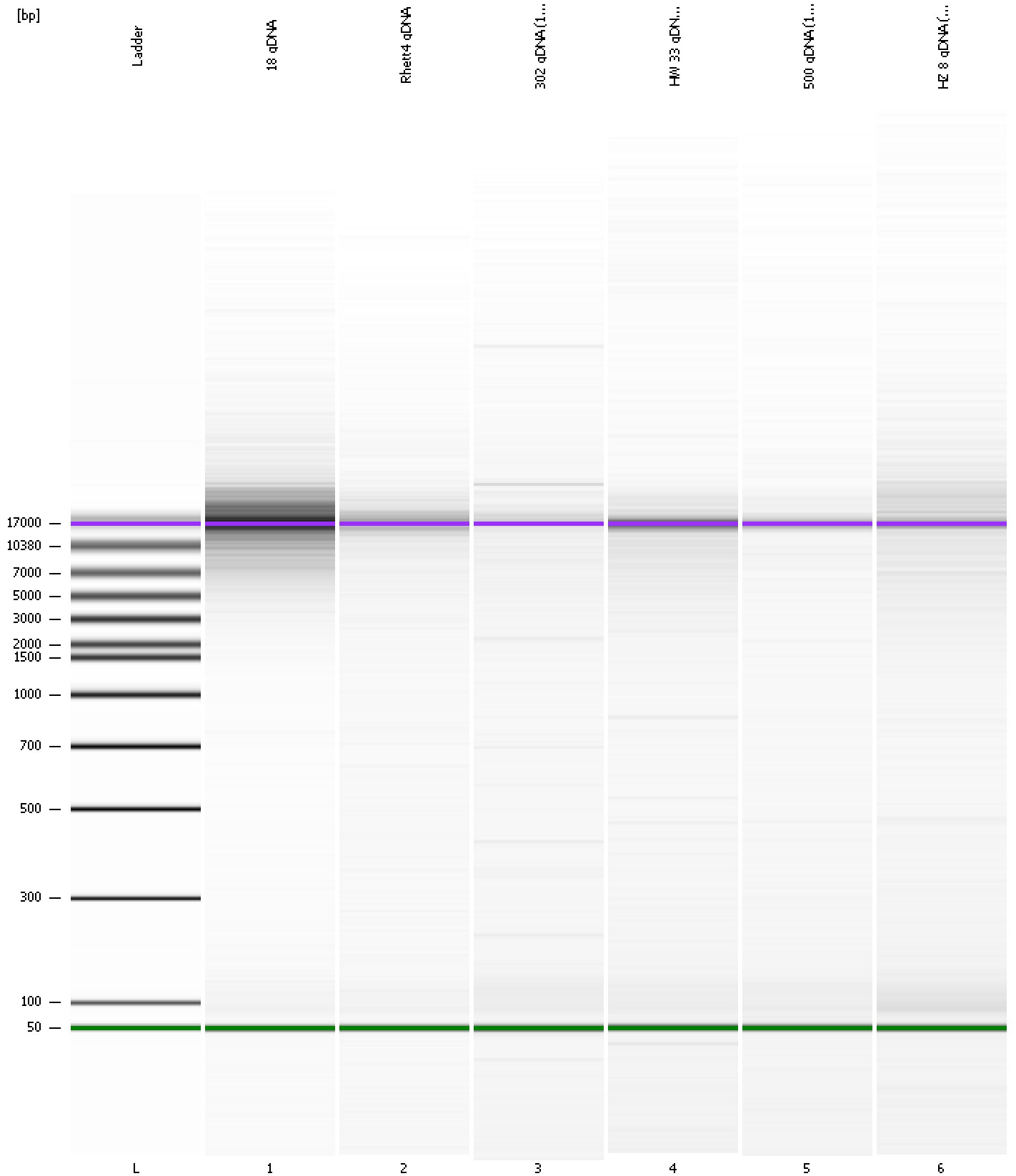
Region table for sample 6 : HZ 8 gDNA (1:2)

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
1,043	45,902	17,309	410.12	13.5	54	59.4	Blue
6,491	35,560	18,659	245.44	8.2	32	37.9	Green
14,333	34,472	21,363	113.35	3.9	15	24.6	Cyan

Assay Class: DNA 12000
Data Path: C:\...d Settings\Bioanalyzer\2014-09-09\2014-09-09_001_Coaker.xad

Created: 9/9/2014 9:44:03 AM
Modified: 9/9/2014 10:18:36 AM

Gel Image



Assay Class: DNA 12000
 Data Path: C:\...d Settings\Bioanalyzer\2014-09-09\2014-09-09_001_Coaker.xad

Created: 9/9/2014 9:44:03 AM
 Modified: 9/9/2014 10:18:36 AM

Run Logbook

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 8)		Instrument	Run		9/9/2014 10:08:28 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2014-09-09\2014-09-09_001.xad)		Instrument	Run		9/9/2014 9:44:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		9/9/2014 9:44:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		9/9/2014 9:44:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		9/9/2014 9:44:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		9/9/2014 9:44:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		9/9/2014 9:44:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		9/9/2014 9:44:09 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1