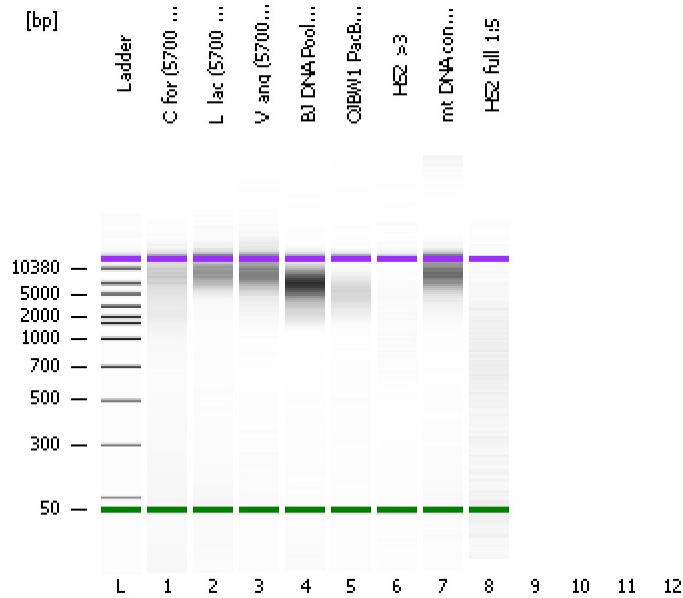


Assay Class: DNA 12000  
Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
Modified: 1/22/2013 4:04:46 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 12000 Series II.xsy

Assay Class: DNA 12000

Version: 2.4

Assay Comments: DNA Analysis 100 -12000 bp

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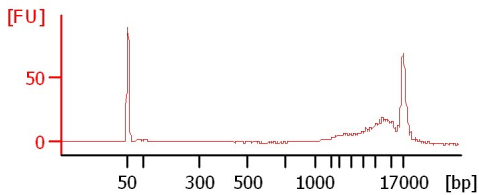
Chip Information:

Chip Lot #:

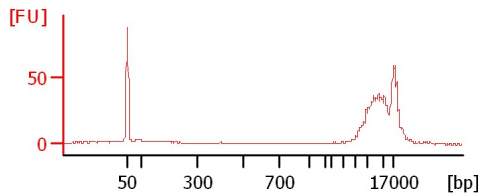
Reagent Kit Lot #:

Chip Comments:

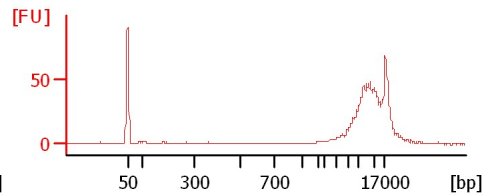
**C\_for (5700 RPM gtube)**



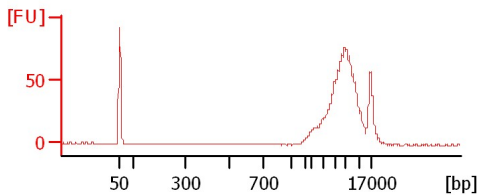
**L\_lac (5700 RPM gtube)**



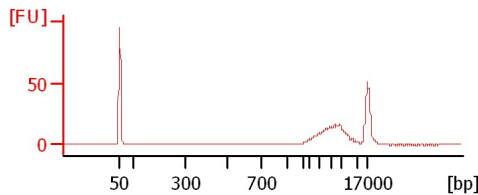
**V\_ang (5700 RPM gtube)**



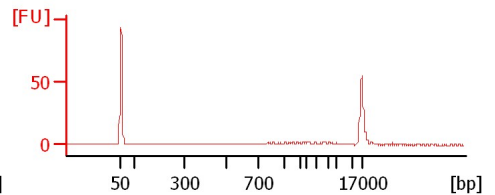
**BJ DNA Pool PacBio Lib**



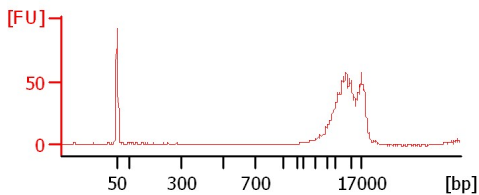
**OJBW1 PacBio Lib**



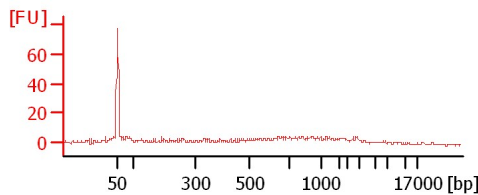
**HS2 >3**



**mt DNA conc 1:5**



**HS2 full 1:5**



Assay Class: DNA 12000  
Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
Modified: 1/22/2013 4:04:46 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
C_for (5700 RPM gtube		<input type="checkbox"/>	✓			
L_lac (5700 RPM gtube		<input type="checkbox"/>	✓			
V_ang (5700 RPM gtube		<input type="checkbox"/>	✓			
BJ DNA Pool PacBio Lib		<input type="checkbox"/>	✓			
OJBW1 PacBio Lib		<input type="checkbox"/>	✓			
HS2 >3		<input type="checkbox"/>	✓			
mt DNA conc 1:5		<input type="checkbox"/>	✓			
HS2 full 1:5		<input type="checkbox"/>	✓			
		<input type="checkbox"/>				
		<input type="checkbox"/>				
		<input type="checkbox"/>				
Ladder		<input type="checkbox"/>	✓			

**Chip Lot #** **Reagent Kit Lot #**

**Chip Comments :**

Assay Class: DNA 12000  
Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
Modified: 1/22/2013 4:04:46 PM

**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

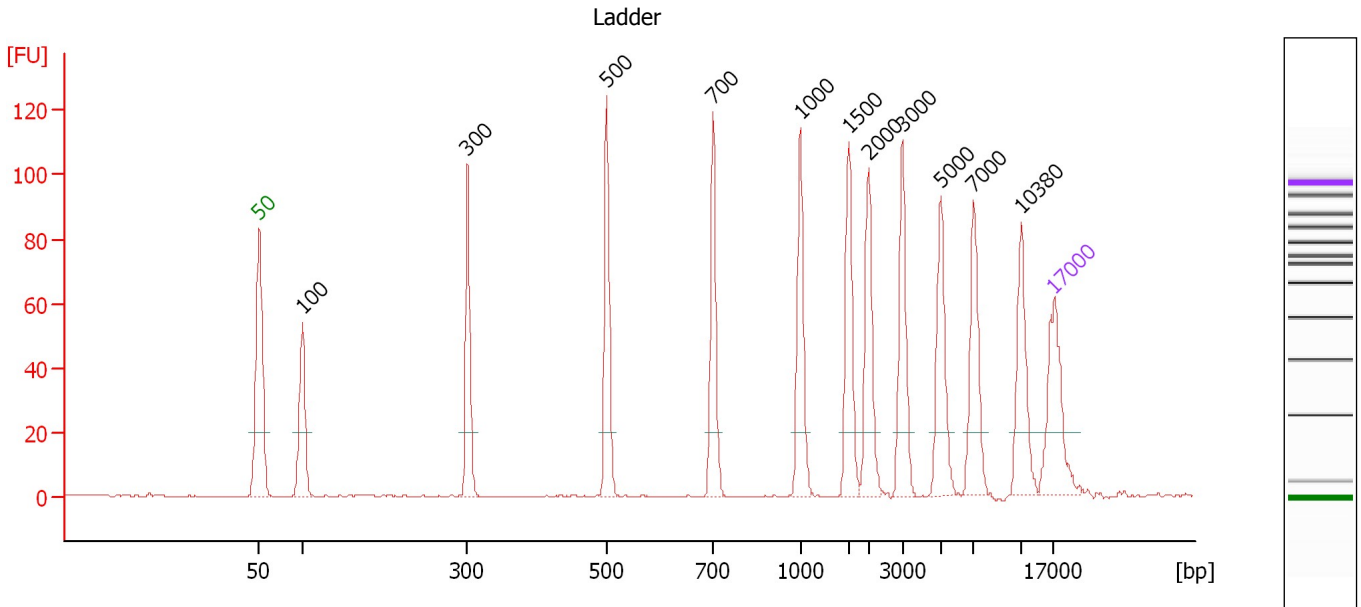
**Ladder**

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:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
 Modified: 1/22/2013 4:04:46 PM

**Electropherogram Summary**



**Peak table for Ladder**

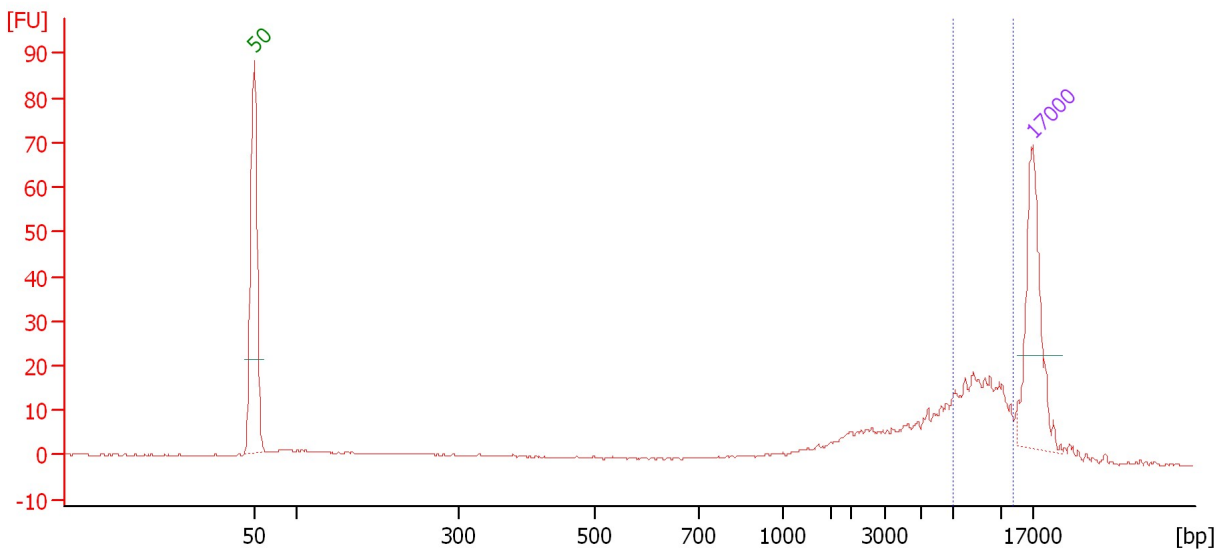
Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	100	4.00	60.6	Ladder Peak
3	300	4.00	20.2	Ladder Peak
4	500	4.00	12.1	Ladder Peak
5	700	4.00	8.7	Ladder Peak
6	1,000	4.00	6.1	Ladder Peak
7	1,500	4.00	4.0	Ladder Peak
8	2,000	4.00	3.0	Ladder Peak
9	3,000	4.00	2.0	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:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
 Modified: 1/22/2013 4:04:46 PM

**Electropherogram Summary Continued ...**

C\_for (5700 RPM gtube



**Overall Results for sample 1 : C\_for (5700 RPM gtube**

Number of peaks found: 0 Area 1: 70.3

**Peak table for sample 1 : C\_for (5700 RPM gtube**

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 : C\_for (5700 RPM gtube**

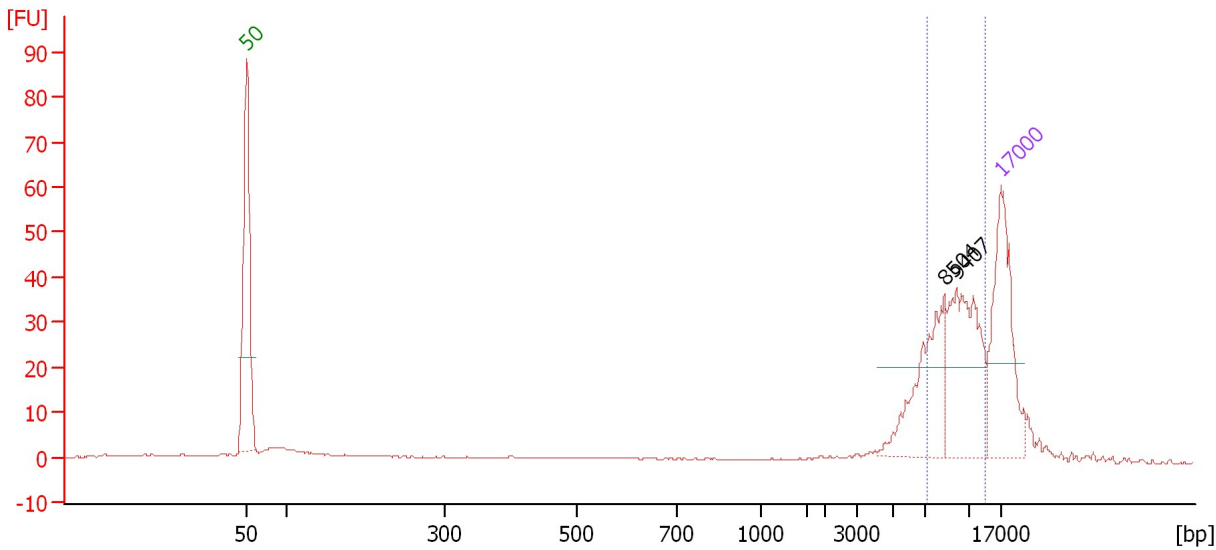
From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
6,967	12,923	9,155	3.90	70.3	36	15.5	Blue

Assay Class: DNA 12000  
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
 Modified: 1/22/2013 4:04:46 PM

**Electropherogram Summary Continued ...**

L\_lac (5700 RPM gtube)



**Overall Results for sample 2 : L\_lac (5700 RPM gtube)**

Number of peaks found: 2                      Area 1: 133.6

**Peak table for sample 2 : L\_lac (5700 RPM gtube)**

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	8,504	3.64	0.6	
3	9,407	4.53	0.7	
4	17,000	4.20	0.4	Upper Marker

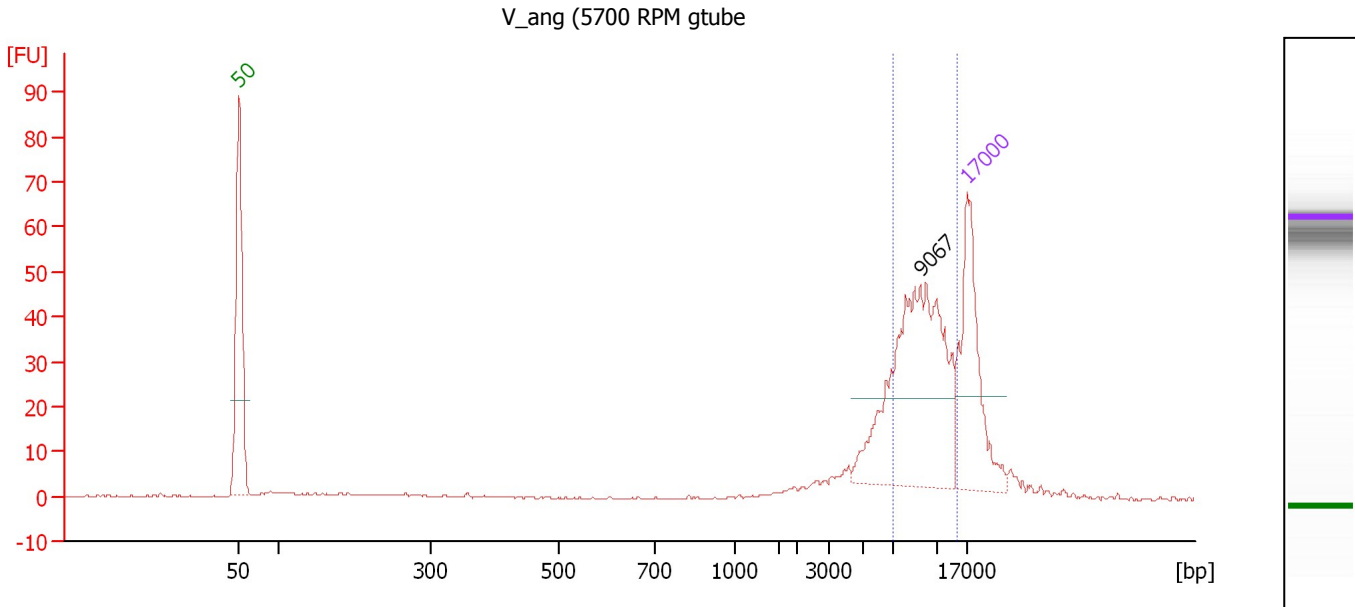
**Region table for sample 2 : L\_lac (5700 RPM gtube)**

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
7,195	13,548	9,528	6.48	133.6	59	16.5	Blue

Assay Class: DNA 12000  
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
 Modified: 1/22/2013 4:04:46 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : V\_ang (5700 RPM gtube)**

Number of peaks found: 1                      Area 1: 175.3

**Peak table for sample 3 : V\_ang (5700 RPM gtube)**

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	9,067	9.53	1.6	
3	17,000	4.20	0.4	Upper Marker

**Region table for sample 3 : V\_ang (5700 RPM gtube)**

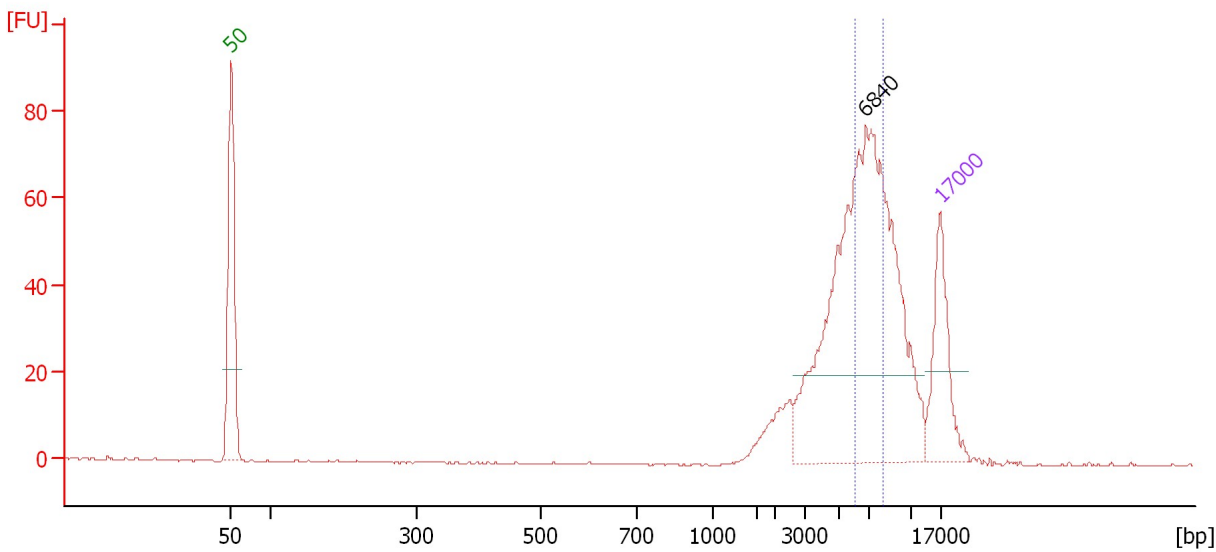
From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
6,976	14,673	9,660	8.25	175.3	62	19.6	Blue

Assay Class: DNA 12000  
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
 Modified: 1/22/2013 4:04:46 PM

**Electropherogram Summary Continued ...**

BJ DNA Pool PacBio Lib



**Overall Results for sample 4 : BJ DNA Pool PacBio Lib**

Number of peaks found: 1 Area: 140.0

**Peak table for sample 4 : BJ DNA Pool PacBio Lib**

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	6,840	26.31	5.8	
3	17,000	4.20	0.4	Upper Marker

**Region table for sample 4 : BJ DNA Pool PacBio Lib**

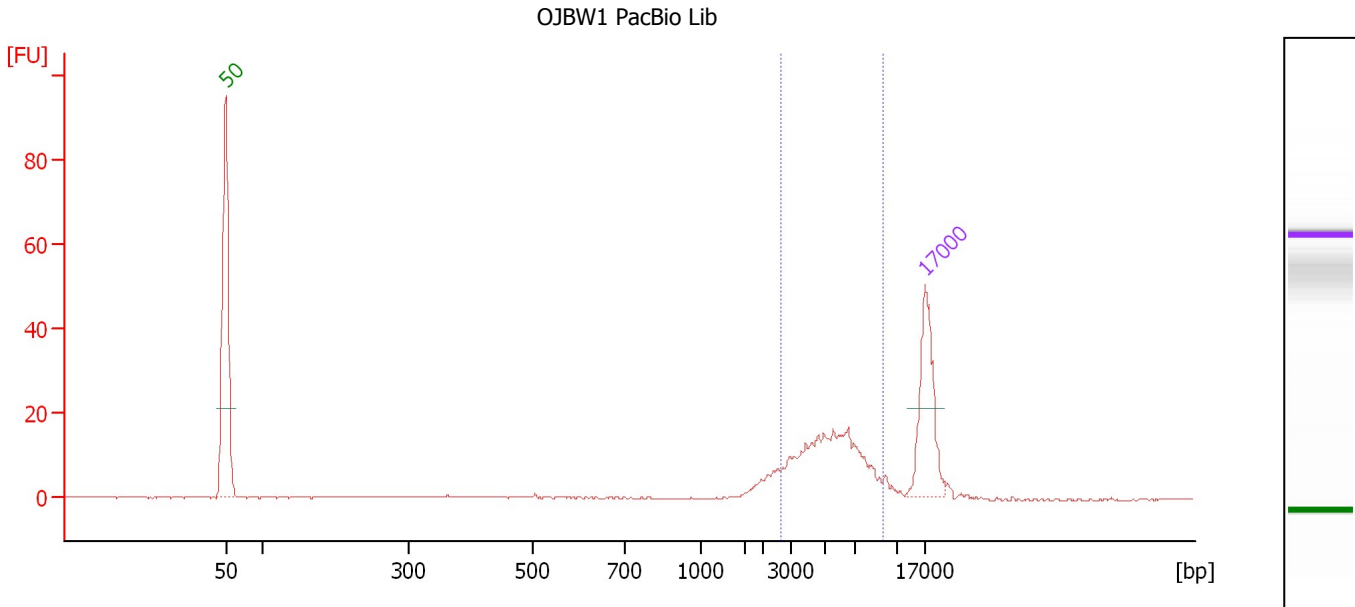
From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
6,092	8,177	7,049	9.26	140.0	33	8.6	Blue



Assay Class: DNA 12000  
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
 Modified: 1/22/2013 4:04:46 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 5 : OJBW1 PacBio Lib**

Number of peaks found: 0                      Area 1: 81.2

**Peak table for sample 5 : OJBW1 PacBio Lib**

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

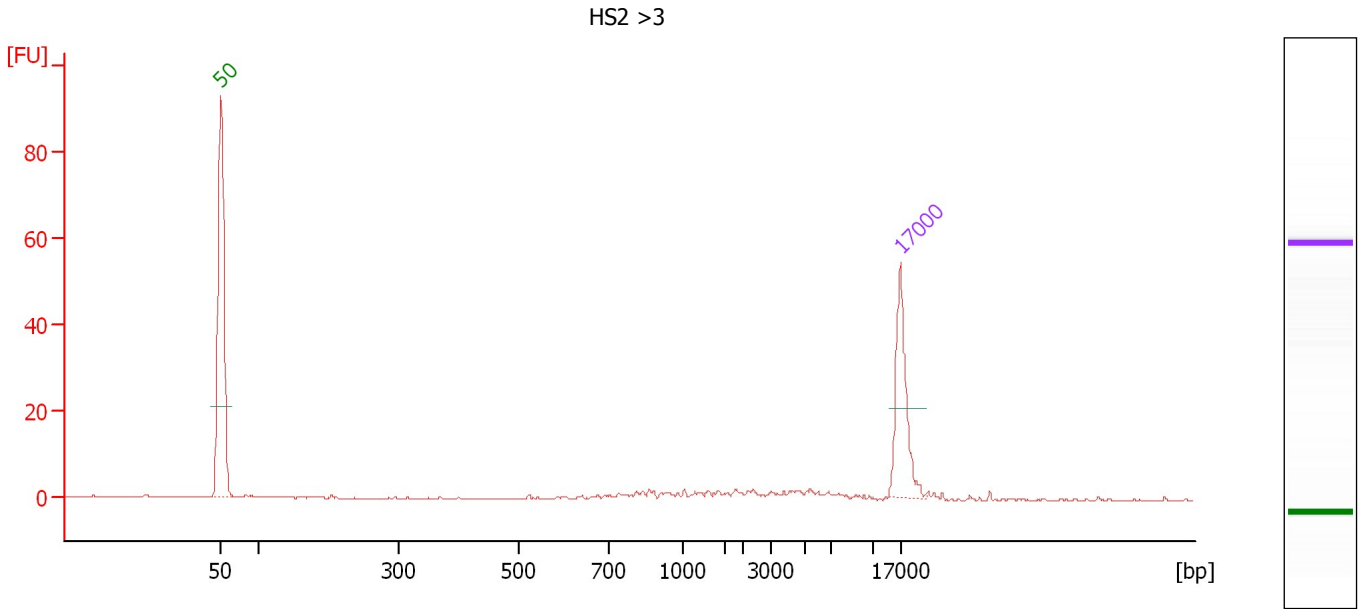
**Region table for sample 5 : OJBW1 PacBio Lib**

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
2,648	9,320	5,442	6.73	81.2	77	31.2	Blue

Assay Class: DNA 12000  
Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
Modified: 1/22/2013 4:04:46 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 6 : HS2 >3**

Number of peaks found: 0

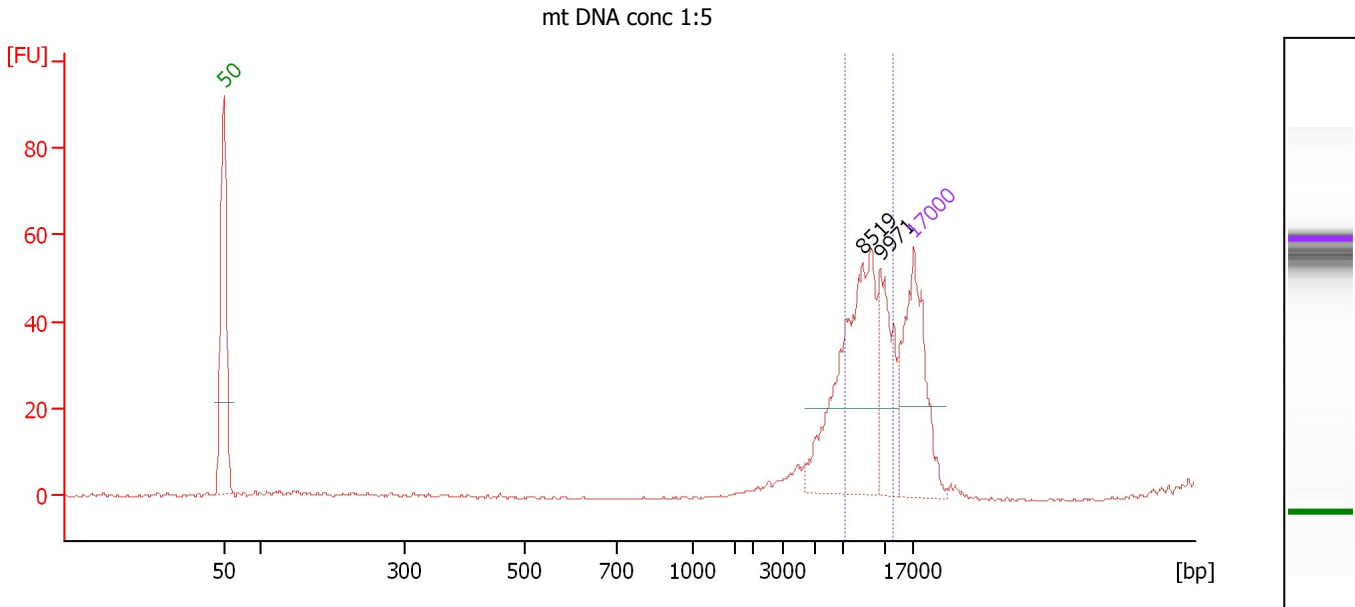
**Peak table for sample 6 : HS2 >3**

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

Assay Class: DNA 12000  
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
 Modified: 1/22/2013 4:04:46 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 7 : mt DNA conc 1:5**

Number of peaks found: 2                      Area 1: 152.0

**Peak table for sample 7 : mt DNA conc 1:5**

Peak	Size [bp]	Conc. [ng/μl]	Molarity [nmol/l]	Observations
1	50	8.30	251.5	Lower Marker
2	8,519	7.27	1.3	
3	9,971	2.54	0.4	
4	17,000	4.20	0.4	Upper Marker

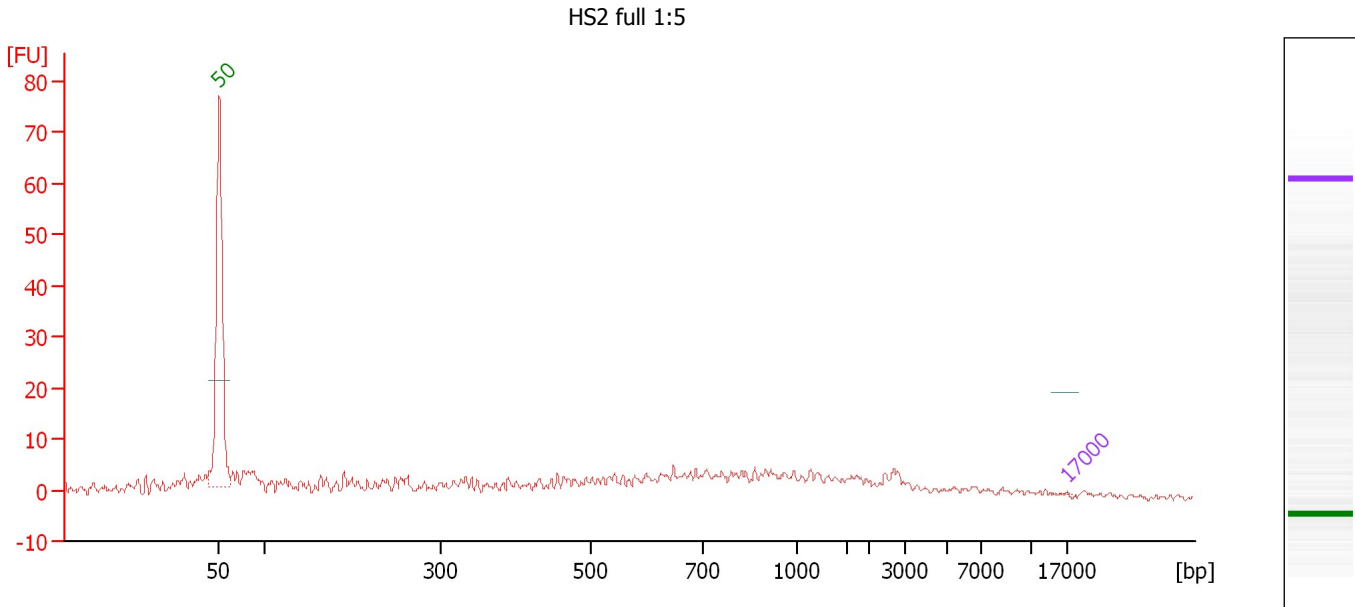
**Region table for sample 7 : mt DNA conc 1:5**

From [bp]	To [bp]	Average Size [bp]	Conc. [ng/μl]	Area	% of Total	Size distribution in CV [%]	Color
7,127	12,188	9,108	6.79	152.0	67	13.3	Blue

Assay Class: DNA 12000  
Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
Modified: 1/22/2013 4:04:46 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 : HS2 full 1:5**

Number of peaks found: 0

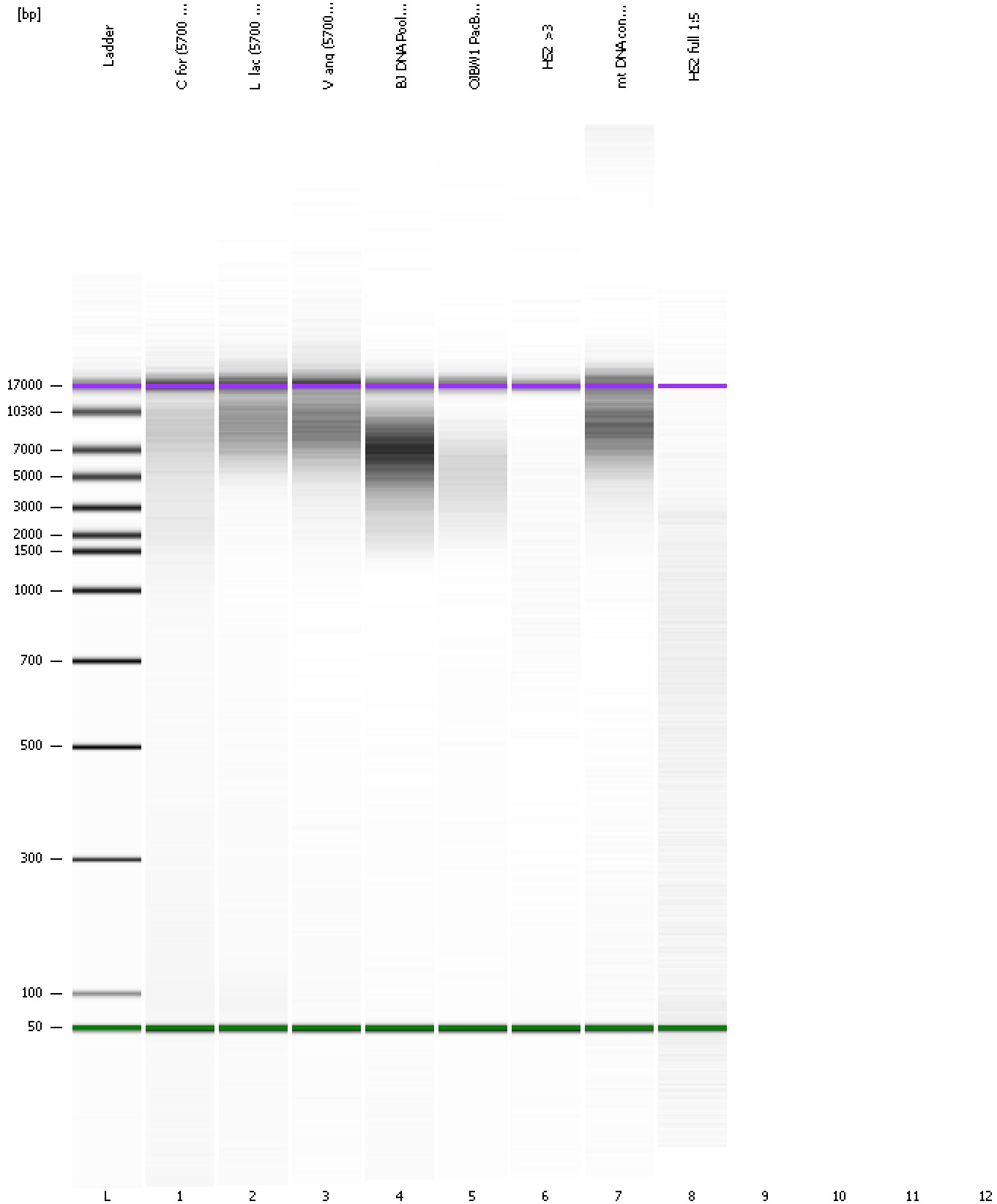
**Peak table for sample 8 : HS2 full 1:5**

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

Assay Class: DNA 12000  
Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
Modified: 1/22/2013 4:04:46 PM

**Gel Image**



Assay Class: DNA 12000  
Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
Modified: 1/22/2013 4:04:46 PM

**Invalid Samples**

Sample 9 has not been run, no results available.

Sample 10 has not been run, no results available.

Sample 11 has not been run, no results available.

Sample 12 has not been run, no results available.

Assay Class: DNA 12000  
 Data Path: C:\...ents and Settings\Bioanalyzer\2013-01-22\2013-01-22\_002.xad

Created: 1/22/2013 3:31:40 PM  
 Modified: 1/22/2013 4:04:46 PM

**Run Logbook**

Description	Number	Source	Category	Sub Category	Time	Time Zone	User	Host
Run ended on port 1 (Number of wells acquired: 9)		Instrument	Run		1/22/2013 3:58:11 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2013-01-22\2013-01-22_002.xad)		Instrument	Run		1/22/2013 3:31:45 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		1/22/2013 3:31:45 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		1/22/2013 3:31:45 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		1/22/2013 3:31:45 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		1/22/2013 3:31:45 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		1/22/2013 3:31:45 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		1/22/2013 3:31:45 PM	(GMT --08:00) Pacific Standard Time	UC Davis	D8XSMGH1