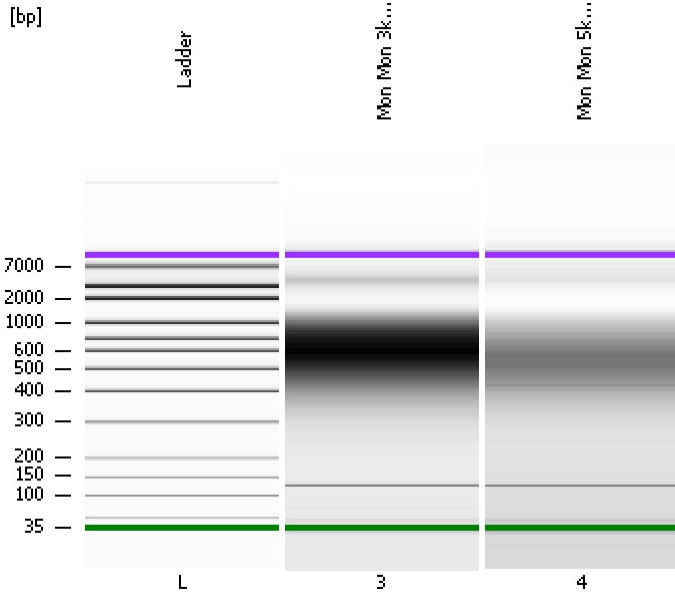


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...16-03-17\2016-03-17\_002\_Mon Mon\_MatePairLibraries\_3K\_5K.xad

Created: 3/17/2016 12:29:29 PM  
Modified: 3/17/2016 1:12:44 PM

**Electrophoresis File Run Summary**



Instrument Information:

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

Assay Information:

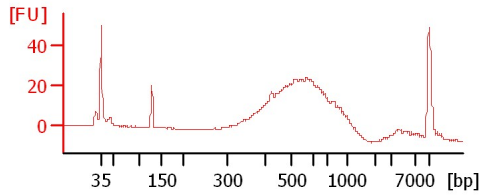
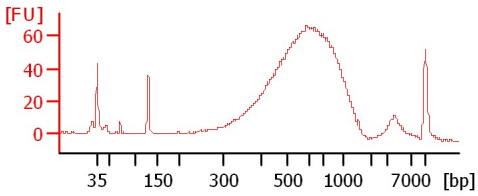
Assay Origin Path: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\assays\dsDNA\High Sensitivity DNA.xsy  
Assay Class: High Sensitivity DNA Assay  
Version: 1.03  
Assay Comments: Copyright © 2003-2010 Agilent Technologies

Chip Information:

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

**Mon Mon 3k Mate Pair Library**

**Mon Mon 5k Mate Pair Library**



Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...16-03-17\2016-03-17\_002\_Mon Mon\_MatePairLibraries\_3K\_5K.xad

Created: 3/17/2016 12:29:29 PM  
 Modified: 3/17/2016 1:12:44 PM

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

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Mon Mon 3k Mate Pair Library		<input type="checkbox"/>				
Mon Mon 5k Mate Pair Library		<input type="checkbox"/>				
Ladder		<input type="checkbox"/>				

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

**Chip Comments :**

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...16-03-17\2016-03-17\_002\_Mon Mon\_MatePairLibraries\_3K\_5K.xad

Created: 3/17/2016 12:29:29 PM  
 Modified: 3/17/2016 1:12:44 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

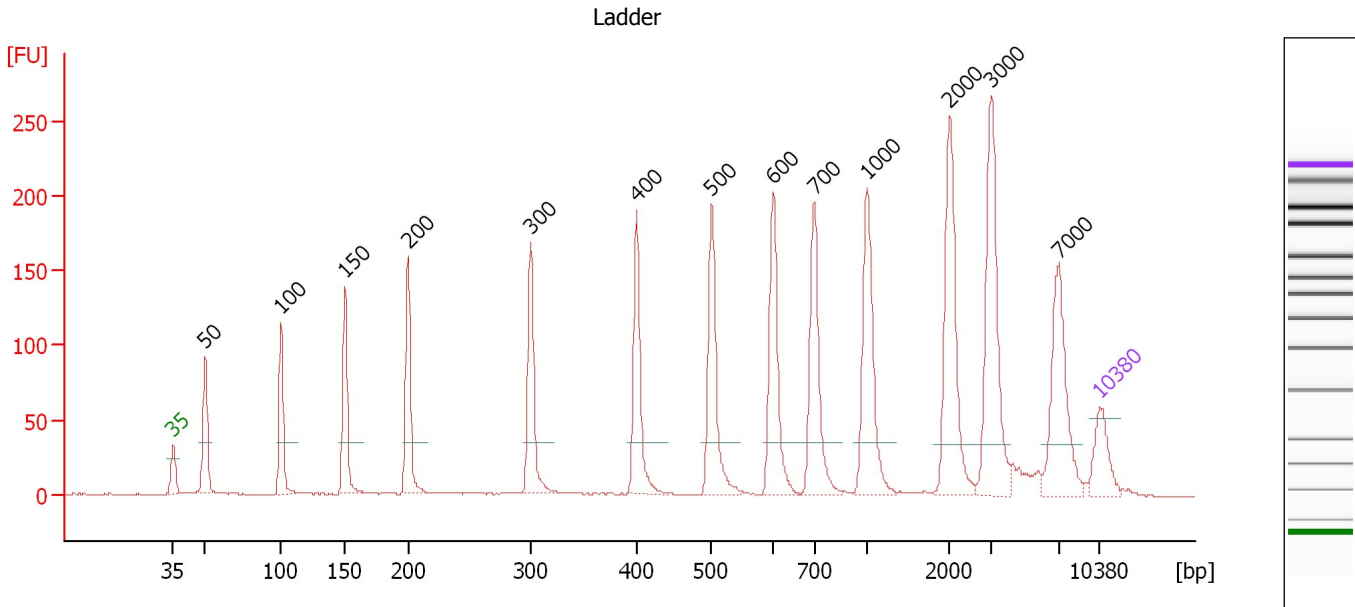
**Ladder**

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:\...16-03-17\2016-03-17\_002\_Mon Mon\_MatePairLibraries\_3K\_5K.xad

Created: 3/17/2016 12:29:29 PM  
 Modified: 3/17/2016 1:12:44 PM

**Electropherogram Summary**



**Overall Results for Ladder**

Noise: 0.3

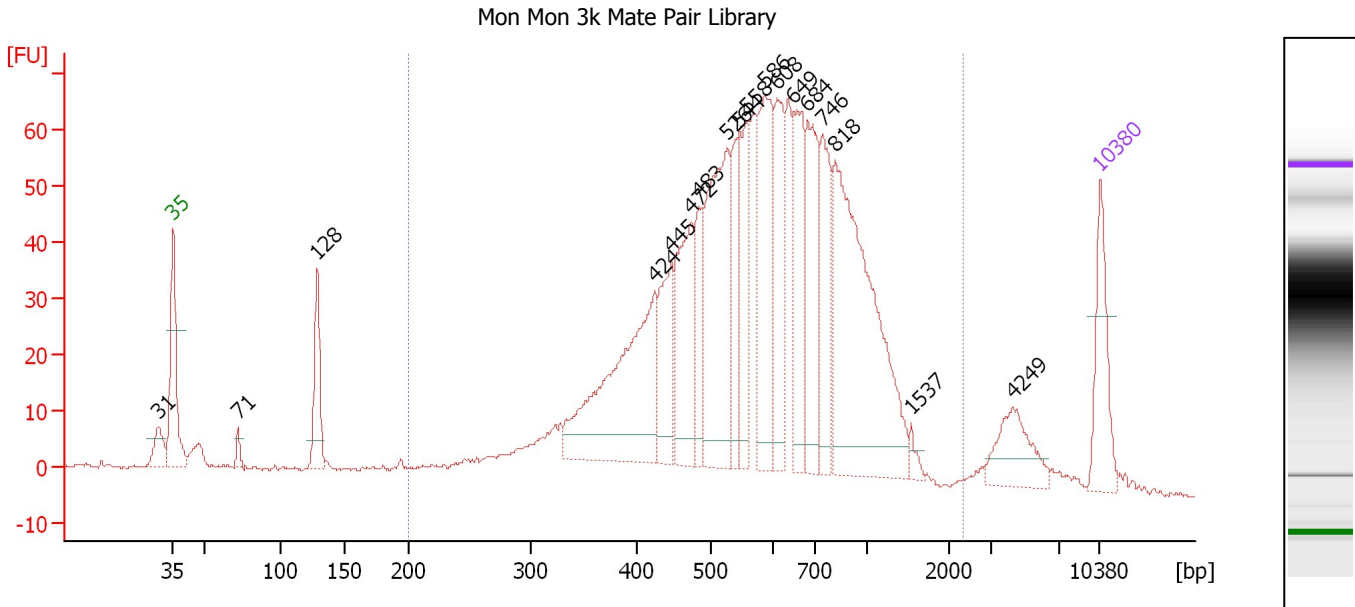
**Peak table for Ladder**

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	45.44
3	100	150.00	2,272.7	Ladder Peak	51.19
4	150	150.00	1,515.2	Ladder Peak	56.02
5	200	150.00	1,136.4	Ladder Peak	60.74
6	300	150.00	757.6	Ladder Peak	70.01
7	400	150.00	568.2	Ladder Peak	78.00
8	500	150.00	454.5	Ladder Peak	83.65
9	600	150.00	378.8	Ladder Peak	88.33
10	700	150.00	324.7	Ladder Peak	91.41
11	1,000	150.00	227.3	Ladder Peak	95.40
12	2,000	150.00	113.6	Ladder Peak	101.59
13	3,000	150.00	75.8	Ladder Peak	104.81
14	7,000	150.00	32.5	Ladder Peak	109.88
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...16-03-17\2016-03-17\_002\_Mon Mon\_MatePairLibraries\_3K\_5K.xad

Created: 3/17/2016 12:29:29 PM  
 Modified: 3/17/2016 1:12:44 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 3 : Mon Mon 3k Mate Pair Library**

Number of peaks found: 18                      Corr. Area 1: 1,336.1  
 Noise: 0.3

**Peak table for sample 3 : Mon Mon 3k Mate Pair Library**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.84
2	35	125.00	5,411.3	Lower Marker	43.00
3	71	13.06	277.4		47.89
4	128	100.37	1,187.9		53.89
5	424	315.08	1,125.7		79.36
6	445	104.52	355.5		80.57
7	472	162.95	522.8		82.09
8	483	79.19	248.4		82.69
9	526	280.95	809.7		84.86
10	544	94.16	262.2		85.72
11	558	96.15	261.0		86.37
12	586	183.98	475.6		87.68
13	608	127.74	318.1		88.59
14	649	132.10	308.2		89.85
15	684	141.70	313.9		90.91
16	746	124.46	252.7		92.02
17	818	416.25	770.9		92.98
18	1,537	10.49	10.3		98.73
19	4,249	63.86	22.8		106.39
20	10,380	75.00	10.9	Upper Marker	113.00

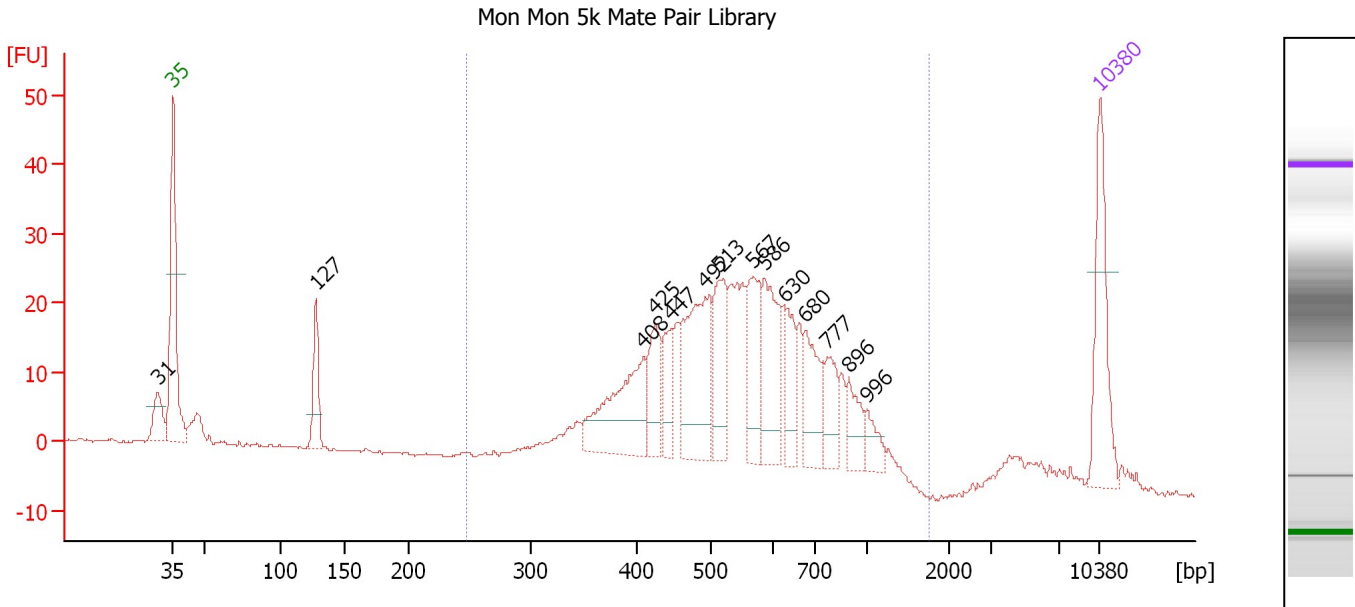
**Region table for sample 3 : Mon Mon 3k Mate Pair Library**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. [pg/μl]	% of Total	Size distribution in CV [%]
200	2,342	628	1,336.1	8,077.1	2,826.80	90	38.4

Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...16-03-17\2016-03-17\_002\_Mon Mon\_MatePairLibraries\_3K\_5K.xad

Created: 3/17/2016 12:29:29 PM  
 Modified: 3/17/2016 1:12:44 PM

**Electropherogram Summary Continued ...**



**Overall Results for sample 4 : Mon Mon 5k Mate Pair Library**

Number of peaks found: 14                      Corr. Area 1: 489.9  
 Noise: 0.2

**Peak table for sample 4 : Mon Mon 5k Mate Pair Library**

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.87
2	35	125.00	5,411.3	Lower Marker	43.00
3	127	56.24	670.8		53.80
4	408	119.63	444.1		78.46
5	425	53.51	190.6		79.43
6	447	37.32	126.5		80.66
7	492	136.97	421.5		83.22
8	513	69.83	206.4		84.24
9	567	69.88	186.7		86.80
10	586	78.43	202.8		87.67
11	630	42.44	102.1		89.26
12	680	57.50	128.1		90.79
13	777	38.94	75.9		92.43
14	896	28.74	48.6		94.02
15	996	19.28	29.3		95.35
16	10,380	75.00	10.9	Upper Marker	113.00

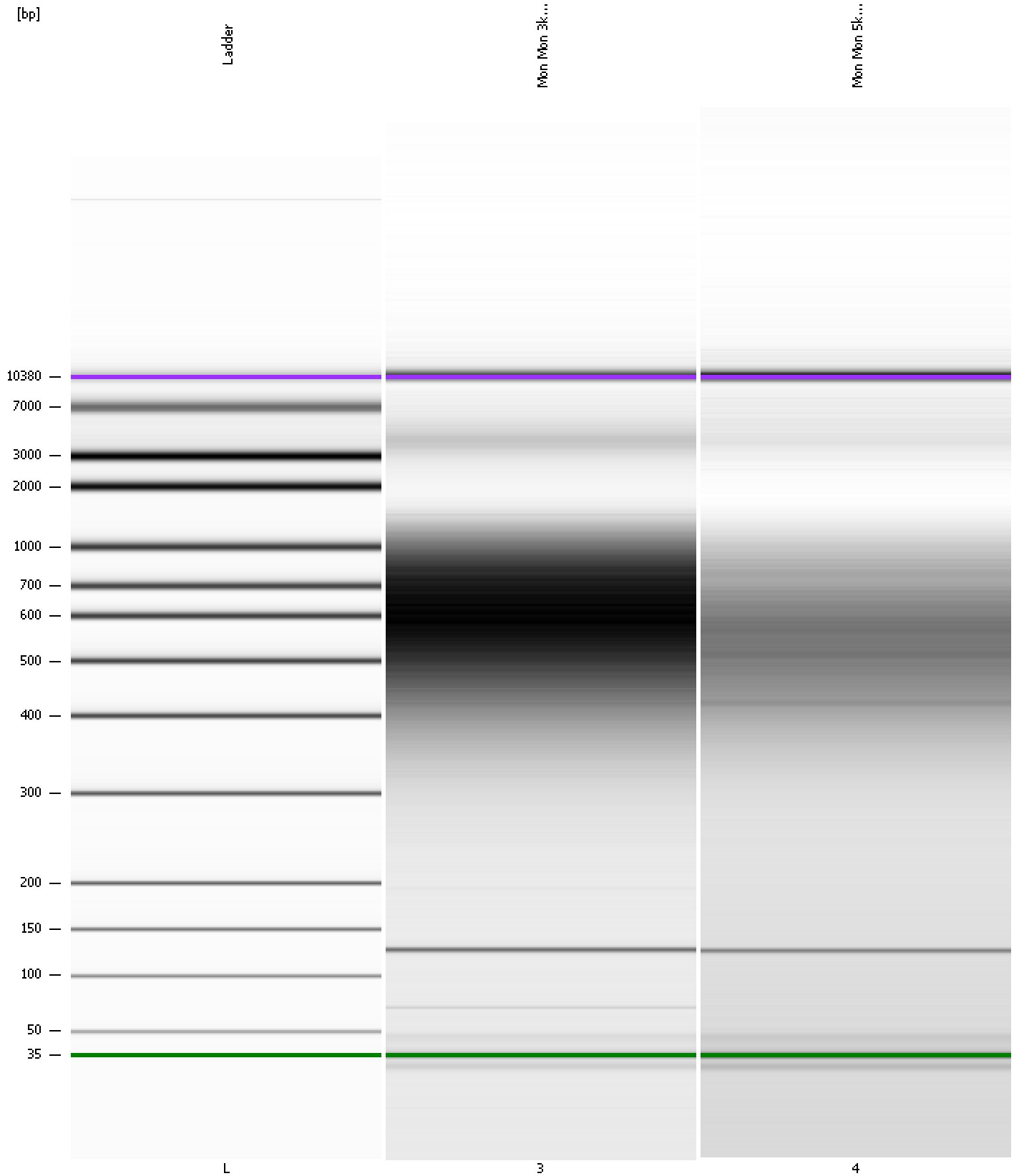
**Region table for sample 4 : Mon Mon 5k Mate Pair Library**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
248	1,763	566	489.9	3,067.3	1,033.71	89	30.1

Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...16-03-17\2016-03-17\_002\_Mon Mon\_MatePairLibraries\_3K\_5K.xad

Created: 3/17/2016 12:29:29 PM  
Modified: 3/17/2016 1:12:44 PM

**Gel Image**

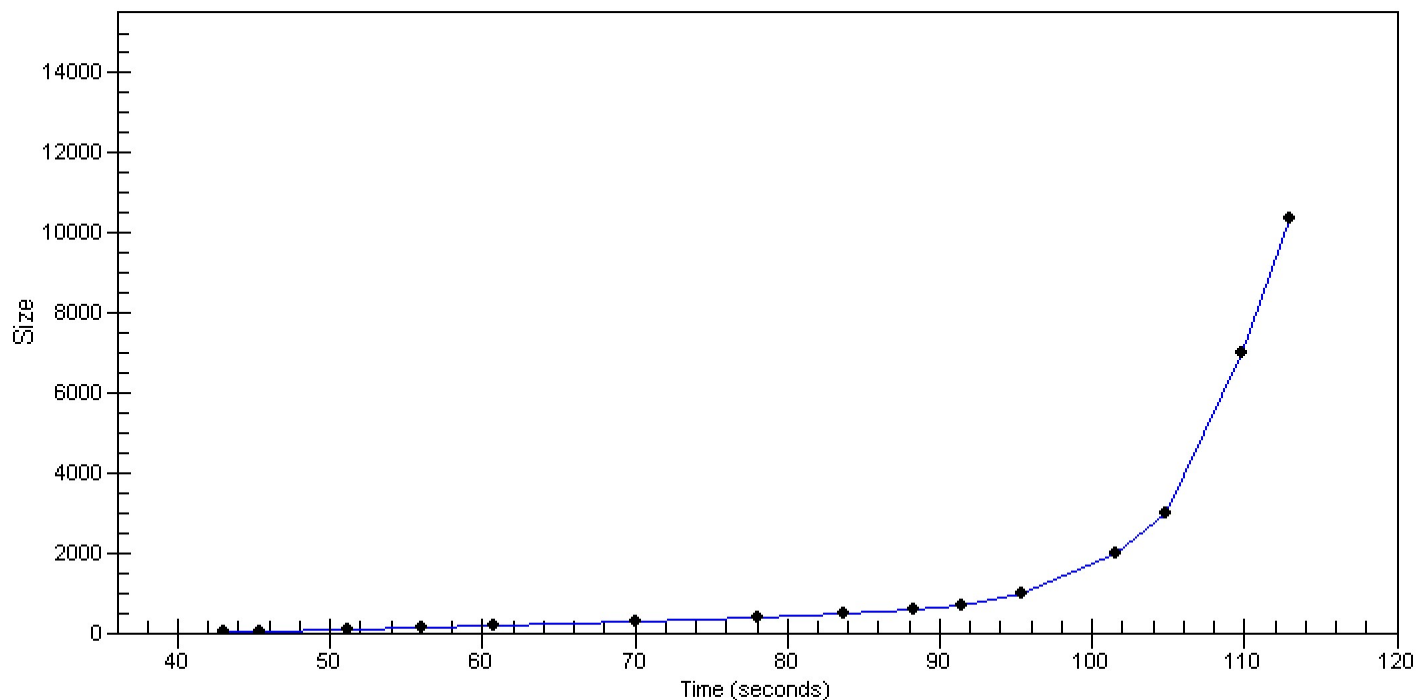


Assay Class: High Sensitivity DNA Assay  
Data Path: C:\...16-03-17\2016-03-17\_002\_Mon Mon\_MatePairLibraries\_3K\_5K.xad

Created: 3/17/2016 12:29:29 PM  
Modified: 3/17/2016 1:12:44 PM

**Curves**

**Standard Curve**





Assay Class: High Sensitivity DNA Assay  
 Data Path: C:\...16-03-17\2016-03-17\_002\_Mon Mon\_MatePairLibraries\_3K\_5K.xad

Created: 3/17/2016 12:29:29 PM  
 Modified: 3/17/2016 1:12:44 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		3/17/2016 1:10:46 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\2016-03-17\2016-03-17_002.xad)		Instrument	Run		3/17/2016 12:29:34 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		3/17/2016 12:29:34 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		3/17/2016 12:29:34 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		3/17/2016 12:29:34 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		3/17/2016 12:29:34 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		3/17/2016 12:29:34 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		3/17/2016 12:29:34 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1