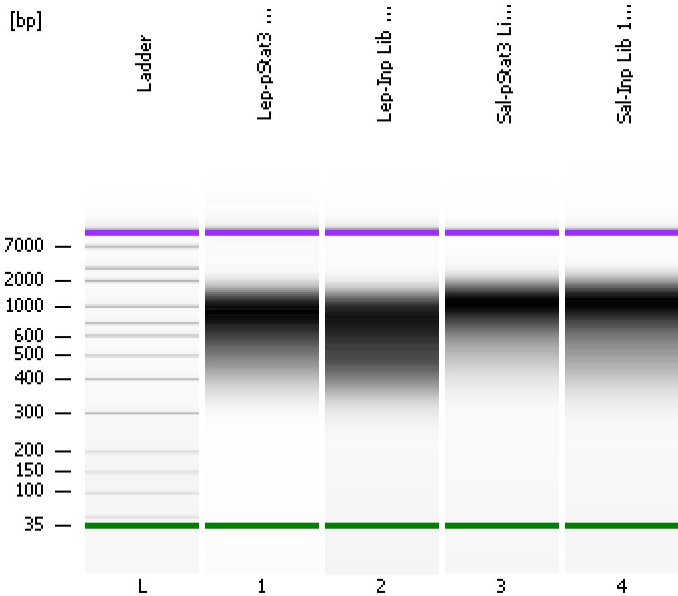


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...tings\Bioanalyzer\2013-07-12\2013-07-12_004_FI_ChIP_Lib.xad

Created: 7/12/2013 4:02:39 PM
Modified: 7/15/2013 4:09:33 PM

Electrophoresis File Run Summary



Instrument Information:

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

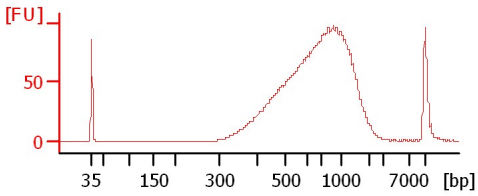
Assay Information:

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

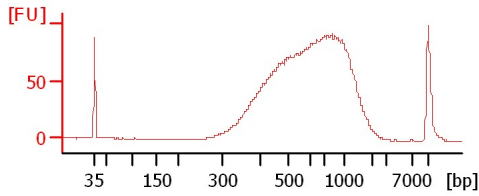
Chip Information:

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

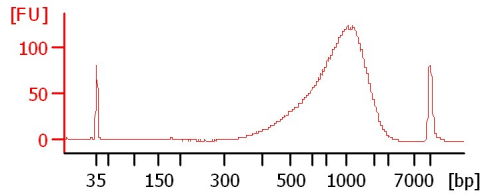
Lep-pStat3 Lib 1:50



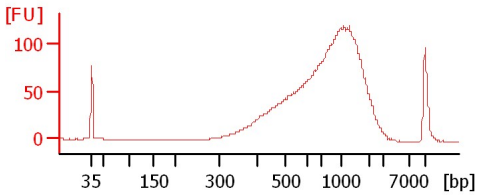
Lep-Inp Lib 1:17



Sal-pStat3 Lib 1:32



Sal-Inp Lib 1:11



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...tings\Bioanalyzer\2013-07-12\2013-07-12_004_FI_ChIP_Lib.xad

Created: 7/12/2013 4:02:39 PM
 Modified: 7/15/2013 4:09:33 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
Lep-pStat3 Lib 1:50		<input type="checkbox"/>	✓			
Lep-Inp Lib 1:17		<input type="checkbox"/>	✓			
Sal-pStat3 Lib 1:32		<input type="checkbox"/>	✓			
Sal-Inp Lib 1:11		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...tings\Bioanalyzer\2013-07-12\2013-07-12_004_FI_ChIP_Lib.xad

Created: 7/12/2013 4:02:39 PM
 Modified: 7/15/2013 4:09:33 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

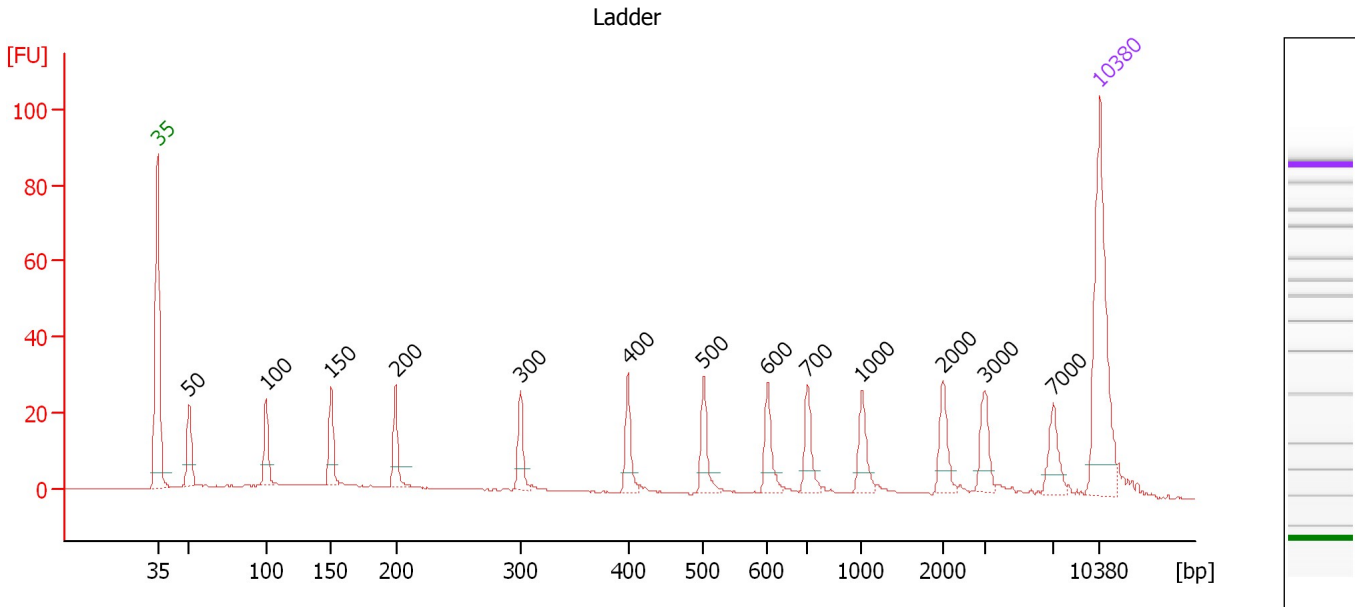
Ladder

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:\...tings\Bioanalyzer\2013-07-12\2013-07-12_004_FI_ChIP_Lib.xad

Created: 7/12/2013 4:02:39 PM
 Modified: 7/15/2013 4:09:33 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

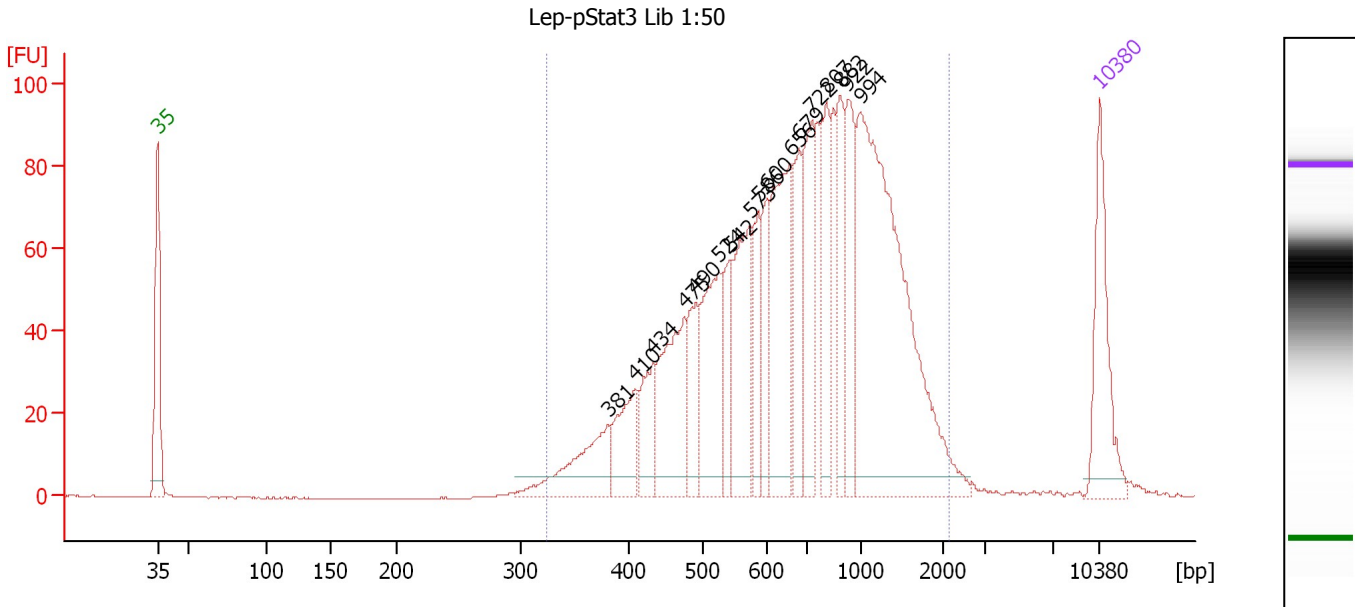
Peak table for Ladder

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	50	150.00	4,545.5	Ladder Peak
3	100	150.00	2,272.7	Ladder Peak
4	150	150.00	1,515.2	Ladder Peak
5	200	150.00	1,136.4	Ladder Peak
6	300	150.00	757.6	Ladder Peak
7	400	150.00	568.2	Ladder Peak
8	500	150.00	454.5	Ladder Peak
9	600	150.00	378.8	Ladder Peak
10	700	150.00	324.7	Ladder Peak
11	1,000	150.00	227.3	Ladder Peak
12	2,000	150.00	113.6	Ladder Peak
13	3,000	150.00	75.8	Ladder Peak
14	7,000	150.00	32.5	Ladder Peak
15	10,380	75.00	10.9	Upper Marker

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...tings\Bioanalyzer\2013-07-12\2013-07-12_004_FI_ChIP_Lib.xad

Created: 7/12/2013 4:02:39 PM
 Modified: 7/15/2013 4:09:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : Lep-pStat3 Lib 1:50

Number of peaks found: 17 Corr. Area 1: 1,750.3
 Noise: 0.1

Peak table for sample 1 : Lep-pStat3 Lib 1:50

Peak	Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	381	84.58	336.0	
3	410	63.97	236.3	
4	434	51.60	180.2	
5	475	122.55	391.3	
6	490	58.80	181.8	
7	524	116.82	337.9	
8	542	50.44	141.0	
9	573	113.18	299.4	
10	586	49.16	127.1	
11	600	54.66	138.1	
12	656	153.02	353.3	
13	679	64.24	143.3	
14	722	96.52	202.7	
15	807	72.95	137.0	
16	882	66.69	114.6	
17	922	73.56	120.8	
18	994	398.63	607.7	
19	10,380	75.00	10.9	Upper Marker

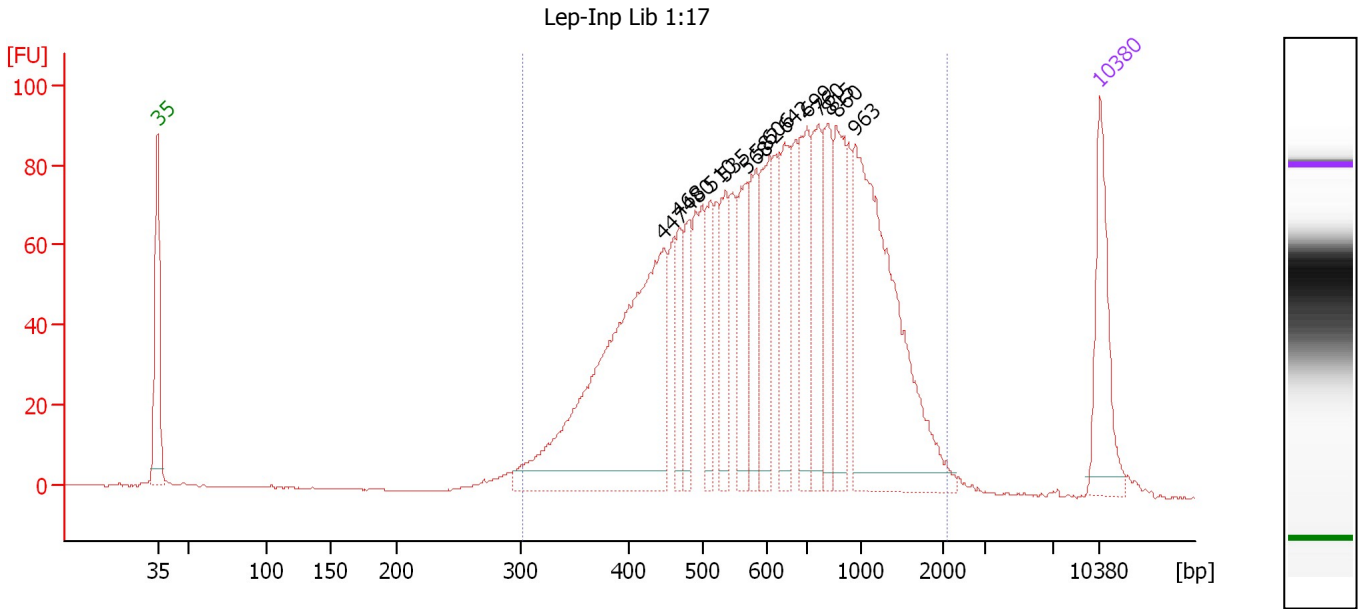
Region table for sample 1 : Lep-pStat3 Lib 1:50

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/µl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
324	2,163	804	4,242.1	1,807.16	1,750.3	97	45.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...tings\Bioanalyzer\2013-07-12\2013-07-12_004_FI_ChIP_Lib.xad

Created: 7/12/2013 4:02:39 PM
 Modified: 7/15/2013 4:09:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : Lep-Inp Lib 1:17

Number of peaks found: 14 Corr. Area 1: 2,079.8
 Noise: 0.2

Peak table for sample 2 : Lep-Inp Lib 1:17

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	447	471.02	1,596.7	
3	468	53.40	173.0	
4	480	54.46	171.9	
5	510	56.67	168.2	
6	535	68.79	194.9	
7	568	76.13	203.1	
8	582	77.51	201.6	
9	606	82.50	206.2	
10	642	87.28	206.1	
11	699	96.13	208.3	
12	760	84.48	168.4	
13	815	68.26	126.9	
14	860	100.12	176.4	
15	963	324.54	510.4	
16	10,380	75.00	10.9	Upper Marker

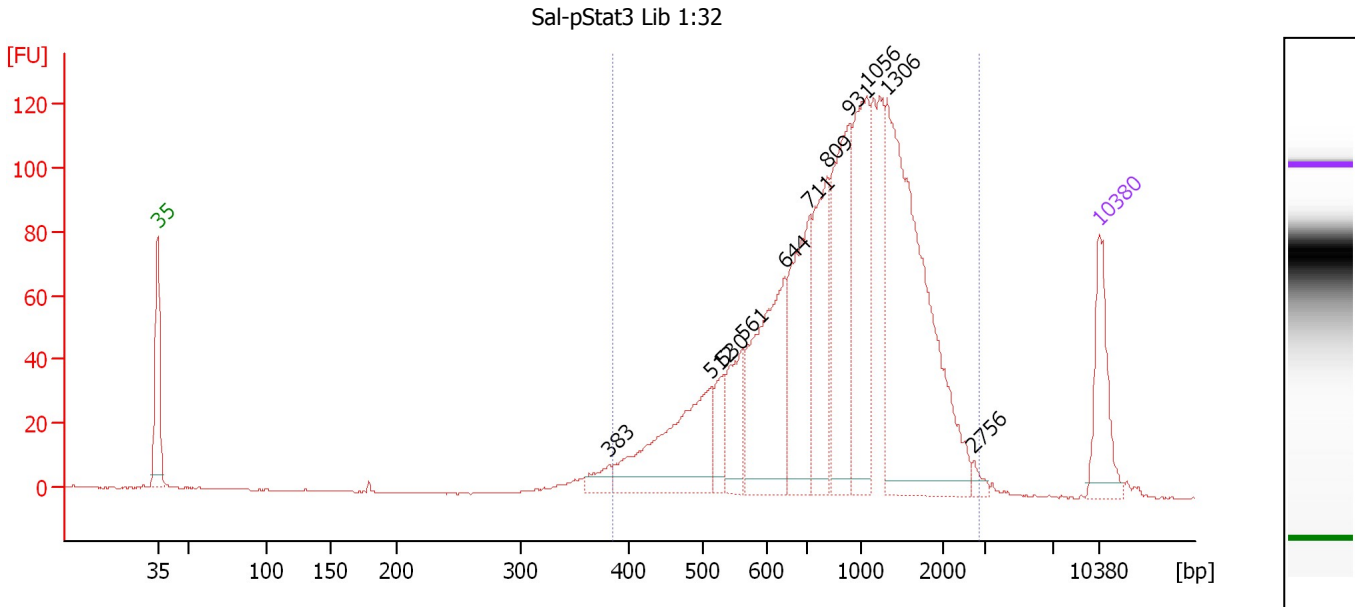
Region table for sample 2 : Lep-Inp Lib 1:17

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
302	2,122	711	5,595.1	2,118.49	2,079.8	97	46.6	Blue

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...tings\Bioanalyzer\2013-07-12\2013-07-12_004_FI_ChIP_Lib.xad

Created: 7/12/2013 4:02:39 PM
 Modified: 7/15/2013 4:09:33 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : Sal-pStat3 Lib 1:32

Number of peaks found: 11 Corr. Area 1: 1,748.6
 Noise: 0.2

Peak table for sample 3 : Sal-pStat3 Lib 1:32

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations
1	35	125.00	5,411.3	Lower Marker
2	383	23.03	91.2	
3	512	217.95	644.5	
4	530	45.18	129.2	
5	561	79.96	216.1	
6	644	240.79	566.6	
7	711	181.03	385.7	
8	809	162.61	304.7	
9	931	207.64	338.0	
10	1,056	187.08	268.5	
11	1,306	455.68	528.5	
12	2,756	8.99	4.9	
13	10,380	75.00	10.9	Upper Marker

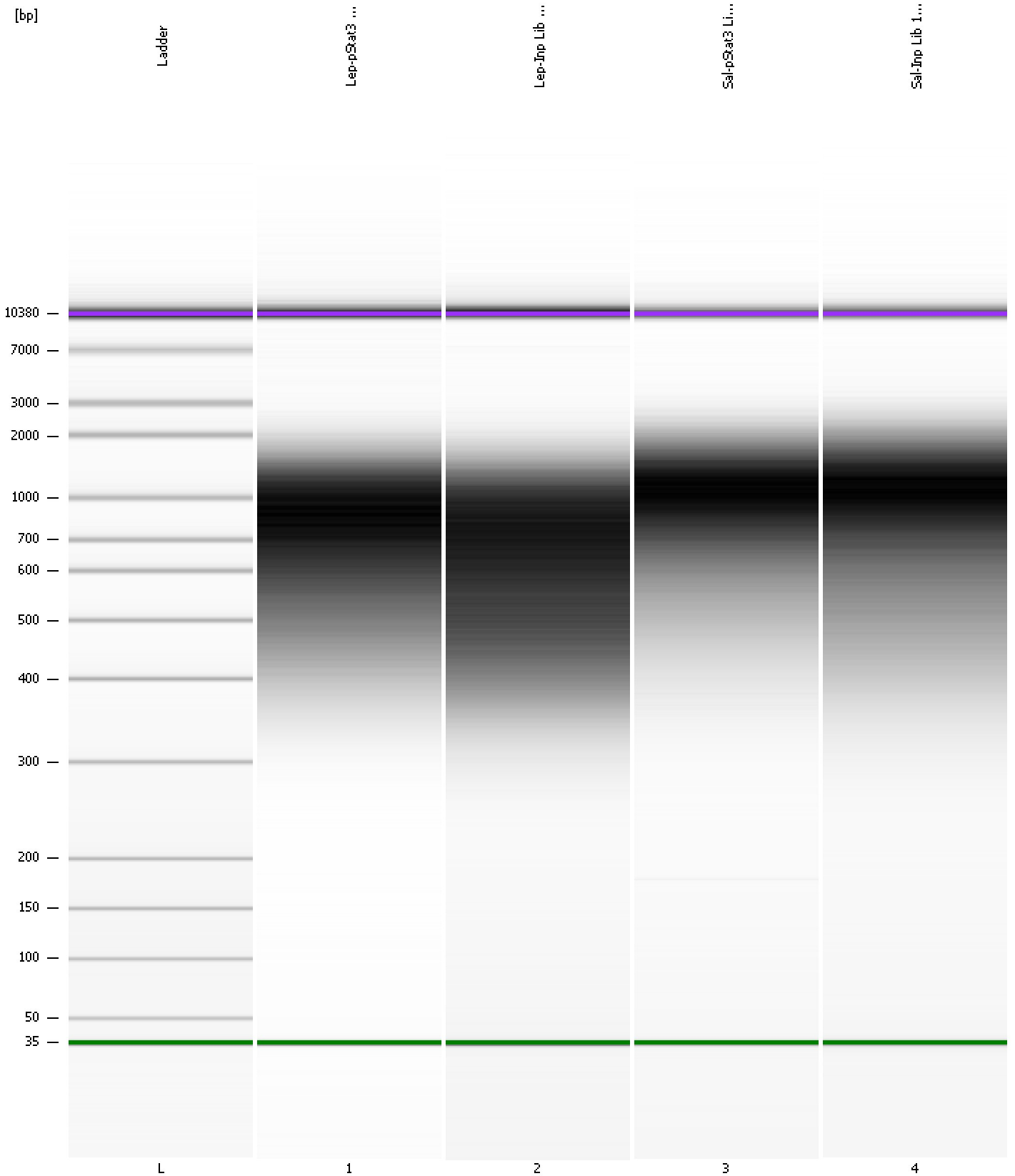
Region table for sample 3 : Sal-pStat3 Lib 1:32

From [bp]	To [bp]	Average Size [bp]	Molarity [pmol/l]	Conc. [pg/μl]	Corr. Area	% of Total	Size distribution in CV [%]	Color
385	2,883	1,020	3,717.3	1,956.97	1,748.6	97	46.8	Blue

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...tings\Bioanalyzer\2013-07-12\2013-07-12_004_FI_ChIP_Lib.xad

Created: 7/12/2013 4:02:39 PM
Modified: 7/15/2013 4:09:33 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...tings\Bioanalyzer\2013-07-12\2013-07-12_004_FI_ChIP_Lib.xad

Created: 7/12/2013 4:02:39 PM
 Modified: 7/15/2013 4:09:33 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		7/12/2013 4:43:57 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2013-07-12\2013-07-12_004.xad)		Instrument	Run		7/12/2013 4:02:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		7/12/2013 4:02:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		7/12/2013 4:02:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		7/12/2013 4:02:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		7/12/2013 4:02:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		7/12/2013 4:02:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		7/12/2013 4:02:44 PM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1