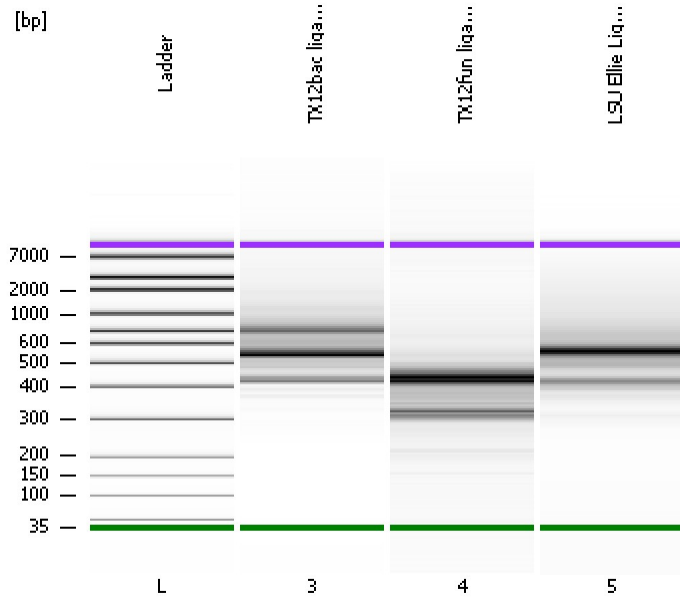


Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...16-10-20\2016-10-20_001_TX12bac_TX12fun_LSUelli_LigRxns.xad

Created: 10/20/2016 10:18:29 AM
 Modified: 10/20/2016 1:40:30 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

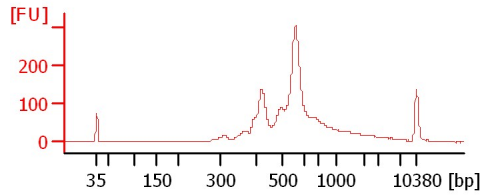
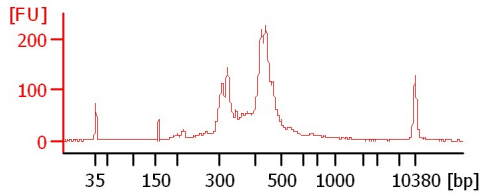
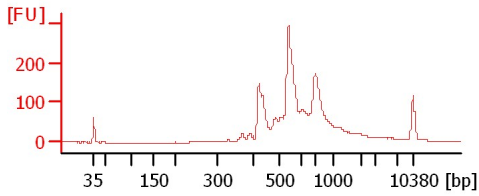
Chip Information:

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

TX12bac ligation rxn (1:3)

TX12fun ligation rxn (1:4)

LSU Ellie Ligation rxn (1:3)



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...16-10-20\2016-10-20_001_TX12bac_TX12fun_LSUElli_LigRxns.xad

Created: 10/20/2016 10:18:29 AM
 Modified: 10/20/2016 1:40:30 PM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
TX12bac ligation rxn (1:3)		<input type="checkbox"/>				
TX12fun ligation rxn (1:4)		<input type="checkbox"/>				
LSU Ellie Ligation rxn (1:3)		<input type="checkbox"/>				
Ladder		<input type="checkbox"/>				

Chip Lot # **Reagent Kit Lot #**

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...16-10-20\2016-10-20_001_TX12bac_TX12fun_LSUElli_LigRxns.xad

Created: 10/20/2016 10:18:29 AM
Modified: 10/20/2016 1:40:30 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

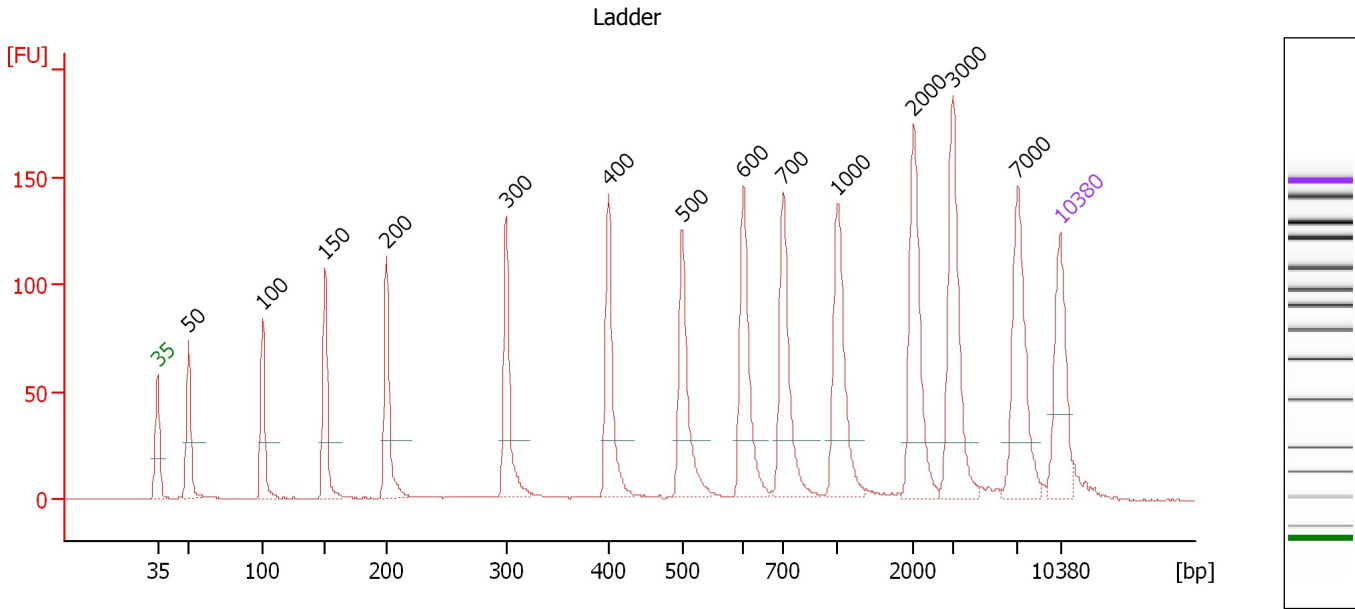
Ladder

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:\...16-10-20\2016-10-20_001_TX12bac_TX12fun_LSUElli_LigRxns.xad

Created: 10/20/2016 10:18:29 AM
 Modified: 10/20/2016 1:40:30 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

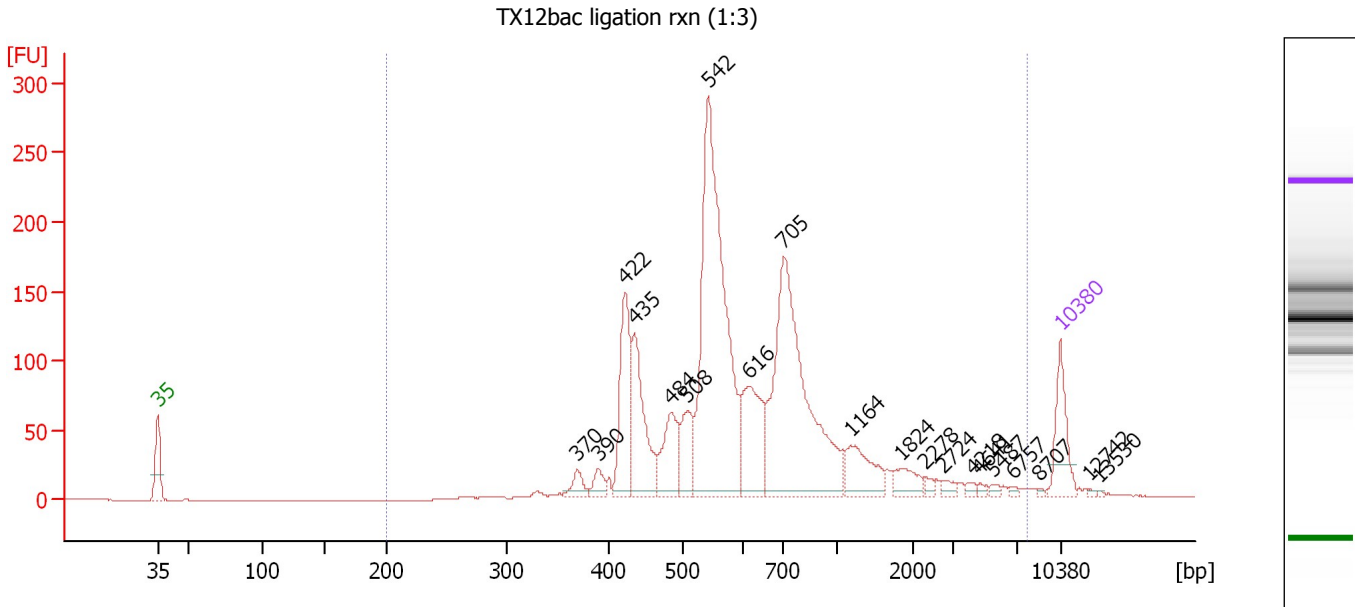
Peak table for Ladder

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	45.41
3	100	150.00	2,272.7	Ladder Peak	51.18
4	150	150.00	1,515.2	Ladder Peak	56.00
5	200	150.00	1,136.4	Ladder Peak	60.77
6	300	150.00	757.6	Ladder Peak	70.04
7	400	150.00	568.2	Ladder Peak	77.98
8	500	150.00	454.5	Ladder Peak	83.70
9	600	150.00	378.8	Ladder Peak	88.42
10	700	150.00	324.7	Ladder Peak	91.49
11	1,000	150.00	227.3	Ladder Peak	95.75
12	2,000	150.00	113.6	Ladder Peak	101.66
13	3,000	150.00	75.8	Ladder Peak	104.68
14	7,000	150.00	32.5	Ladder Peak	109.69
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...16-10-20\2016-10-20_001_TX12bac_TX12fun_LSUElli_LigRxns.xad

Created: 10/20/2016 10:18:29 AM
 Modified: 10/20/2016 1:40:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : TX12bac ligation rxn (1:3)

Number of peaks found: 20 Corr. Area 1: 2,253.7
 Noise: 0.2

Peak table for sample 3 : TX12bac ligation rxn (1:3)

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	370	32.92	134.9		75.57
3	390	28.39	110.4		77.16
4	422	192.03	689.7		79.23
5	435	188.29	656.5		79.95
6	484	113.04	353.7		82.79
7	508	87.15	259.7		84.09
8	542	723.32	2,021.8		85.68
9	616	164.44	404.5		88.91
10	705	586.43	1,260.8		91.56
11	1,164	84.49	110.0		96.72
12	1,824	36.63	30.4		100.62
13	2,278	9.71	6.5		102.50
14	2,724	11.35	6.3		103.85
15	4,218	7.55	2.7		106.21
16	4,641	6.11	2.0		106.74
17	5,487	6.11	1.7		107.80
18	6,757	4.27	1.0		109.39
19	8,707	2.96	0.5		111.36
20	10,380	75.00	10.9	Upper Marker	113.00
21	12,742	0.00	0.0		115.31
22	13,530	0.00	0.0		116.08

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...16-10-20\2016-10-20_001_TX12bac_TX12fun_LSUElli_LigRxns.xad

Created: 10/20/2016 10:18:29 AM
Modified: 10/20/2016 1:40:30 PM

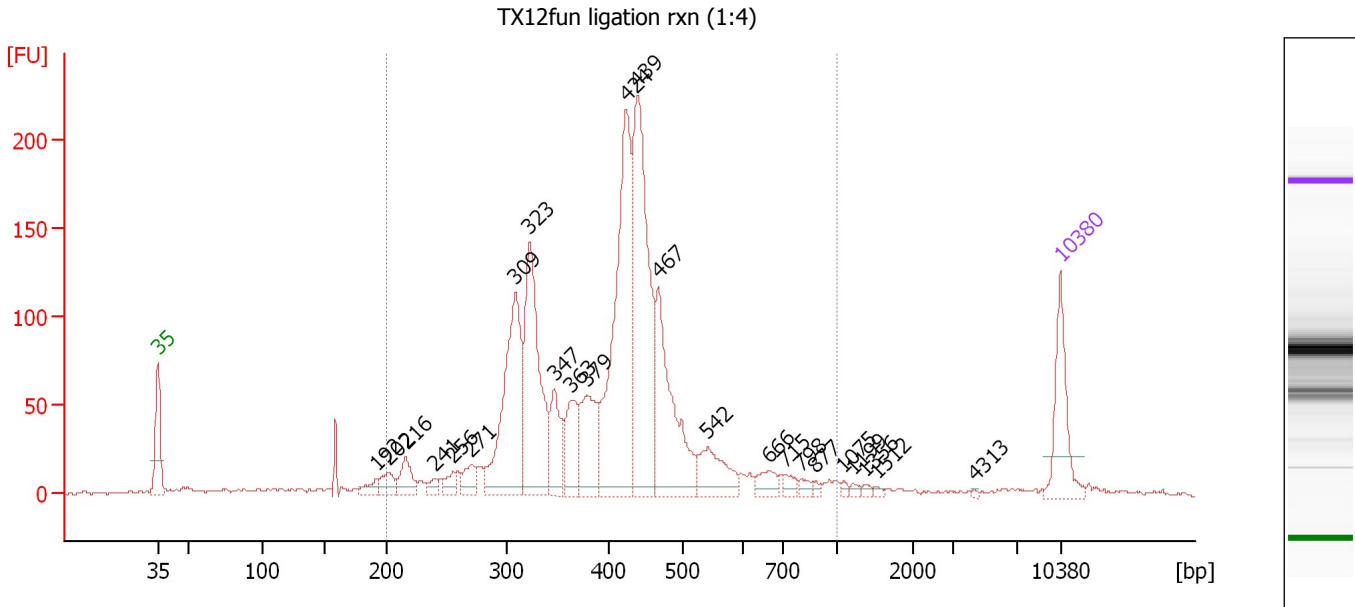
Electropherogram Summary Continued ...**... Region table for sample 3 : TX12bac ligation rxn (1:3)**

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	7,791	824	2,253.7	6,027.8	■ 2,345.50	99	100.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...16-10-20\2016-10-20_001_TX12bac_TX12fun_LSUElli_LigRxns.xad

Created: 10/20/2016 10:18:29 AM
 Modified: 10/20/2016 1:40:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : TX12fun ligation rxn (1:4)

Number of peaks found: 24 Corr. Area 1: 2,160.6
 Noise: 0.8

Peak table for sample 4 : TX12fun ligation rxn (1:4)

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	192	18.40	145.3		60.00
3	202	28.58	214.7		60.93
4	216	43.07	302.3		62.24
5	241	13.80	86.9		64.53
6	256	19.80	117.3		65.94
7	271	30.64	171.3		67.36
8	309	270.00	1,323.1		70.77
9	323	277.24	1,301.7		71.84
10	347	75.98	331.5		73.79
11	363	79.92	333.4		75.05
12	379	102.88	411.2		76.32
13	424	393.75	1,407.6		79.34
14	439	376.91	1,300.4		80.22
15	467	211.72	686.5		81.82
16	542	74.61	208.7		85.67
17	666	24.63	56.0		90.45
18	715	12.19	25.8		91.71
19	798	9.49	18.0		92.88
20	877	4.74	8.2		94.00
21	1,075	4.14	5.8		96.19
22	1,199	4.90	6.2		96.92
23	1,356	4.52	5.1		97.85
24	1,512	3.66	3.7		98.78
25	4,313	1.80	0.6		106.33

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...16-10-20\2016-10-20_001_TX12bac_TX12fun_LSUElli_LigRxns.xad

Created: 10/20/2016 10:18:29 AM
 Modified: 10/20/2016 1:40:30 PM

Electropherogram Summary Continued ...

... Peak table for sample 4 : TX12fun ligation rxn (1:4)

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
26	10,380	75.00	10.9	Upper Marker	113.00

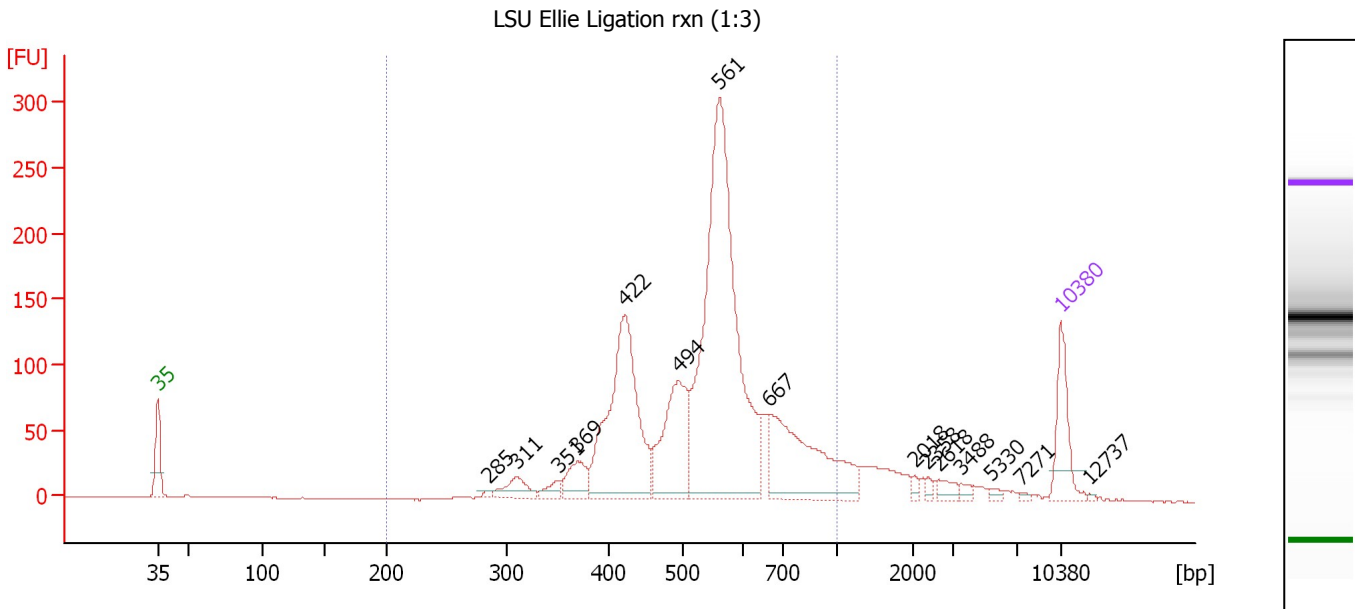
Region table for sample 4 : TX12fun ligation rxn (1:4)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	413	2,160.6	8,537.0	2,134.11	89	26.8

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...16-10-20\2016-10-20_001_TX12bac_TX12fun_LSUElli_LigRxns.xad

Created: 10/20/2016 10:18:29 AM
 Modified: 10/20/2016 1:40:30 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : LSU Ellie Ligation rxn (1:3)

Number of peaks found: 15 Corr. Area 1: 2,157.9
 Noise: 0.3

Peak table for sample 5 : LSU Ellie Ligation rxn (1:3)

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	285	5.70	30.3		68.67
3	311	44.97	219.3		70.88
4	351	20.78	89.6		74.12
5	369	57.05	234.0		75.55
6	422	405.21	1,455.3		79.23
7	494	209.83	643.6		83.35
8	561	813.67	2,196.6		86.59
9	667	264.69	601.6		90.47
10	2,018	7.40	5.6		101.71
11	2,358	6.69	4.3		102.74
12	2,618	15.93	9.2		103.53
13	3,488	8.16	3.5		105.29
14	5,330	5.78	1.6		107.60
15	7,271	3.36	0.7		109.96
16	10,380	75.00	10.9	Upper Marker	113.00
17	12,737	0.00	0.0		115.31

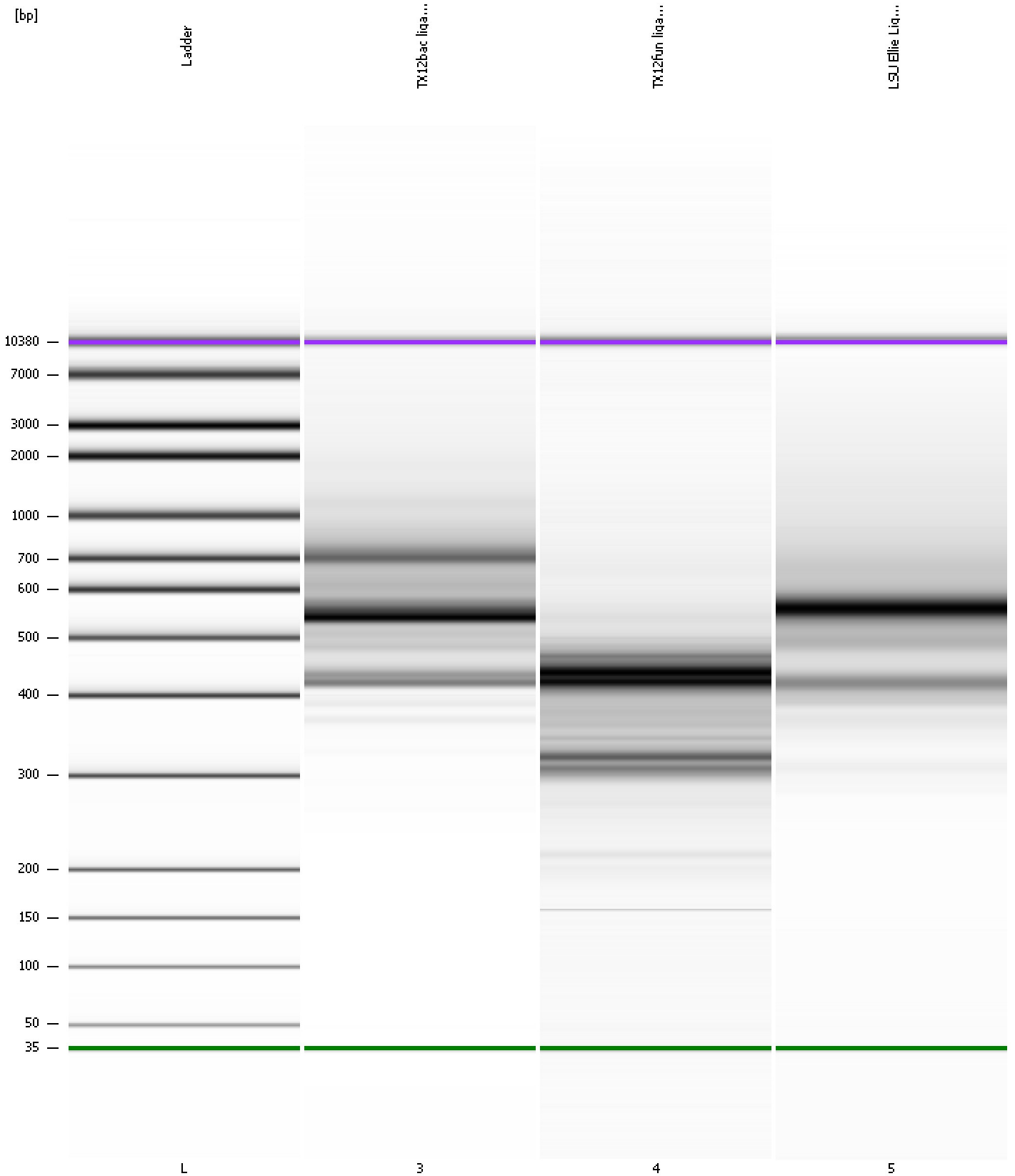
Region table for sample 5 : LSU Ellie Ligation rxn (1:3)

From [bp]	To [bp]	Average Size [bp]	Corr. Area	Molarity [pmol/l]	Co Conc. lor [pg/μl]	% of Total	Size distribution in CV [%]
200	1,000	547	2,157.9	5,428.8	1,823.67	89	23.5

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...16-10-20\2016-10-20_001_TX12bac_TX12fun_LSUElli_LigRxns.xad

Created: 10/20/2016 10:18:29 AM
Modified: 10/20/2016 1:40:30 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...16-10-20\2016-10-20_001_TX12bac_TX12fun_LSUElli_LigRxns.xad

Created: 10/20/2016 10:18:29 AM
 Modified: 10/20/2016 1:40:30 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		10/20/2016 10:59:48 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-10-20\2016-10-20_001.xad)		Instrument	Run		10/20/2016 10:18:34 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Product Number : G2938B		Instrument	Run		10/20/2016 10:18:34 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Name :		Instrument	Run		10/20/2016 10:18:34 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Vendor : Agilent Technologies		Instrument	Run		10/20/2016 10:18:34 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Serial# : DE13701086		Instrument	Run		10/20/2016 10:18:34 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Firmware : C.01.069		Instrument	Run		10/20/2016 10:18:34 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1
Cartridge : Electrode		Instrument	Run		10/20/2016 10:18:34 AM	(GMT --07:00) Pacific Standard Time	UC Davis	D8XSMGH1