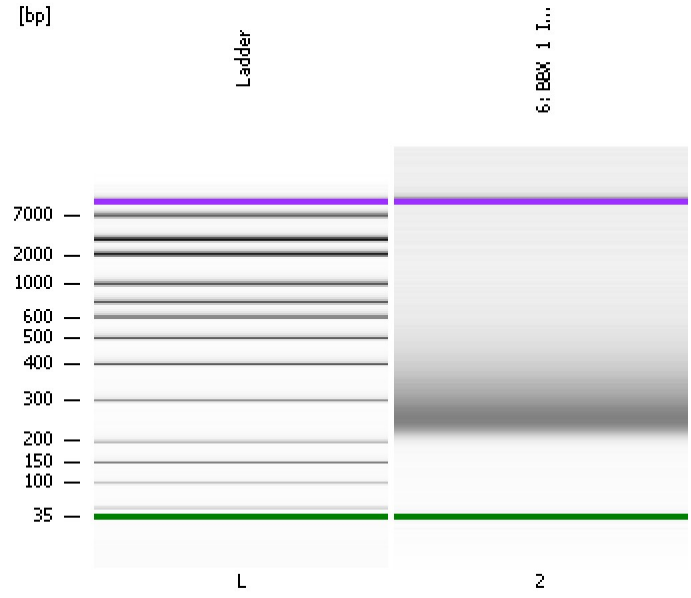
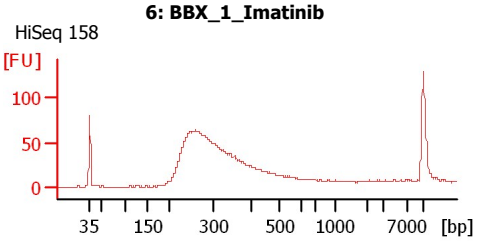
Assay Class: High Sensitivity DNA Assay
Data Path: C:\...analyzer\2015-08-07\2015-08-07_001_HiSeq158_Lib_Sample6.xad

Created: 8/7/2015 11:21:09 AM
Modified: 8/10/2015 10:27:09 AM

Electrophoresis File Run Summary



6: BBX_1_Imatinib



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:	

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...analyzer\2015-08-07\2015-08-07_001_HiSeq158_Lib_Sample6.xad

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Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
6: BBX_1_Imatinib	HiSeq 158	☐	✔			
Ladder		☐	✔			

Chip Lot # **Reagent Kit Lot #**

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...analyzer\2015-08-07\2015-08-07_001_HiSeq158_Lib_Sample6.xad

Created: 8/7/2015 11:21:09 AM
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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:\...analyzer\2015-08-07\2015-08-07_001_HiSeq158_Lib_Sample6.xad

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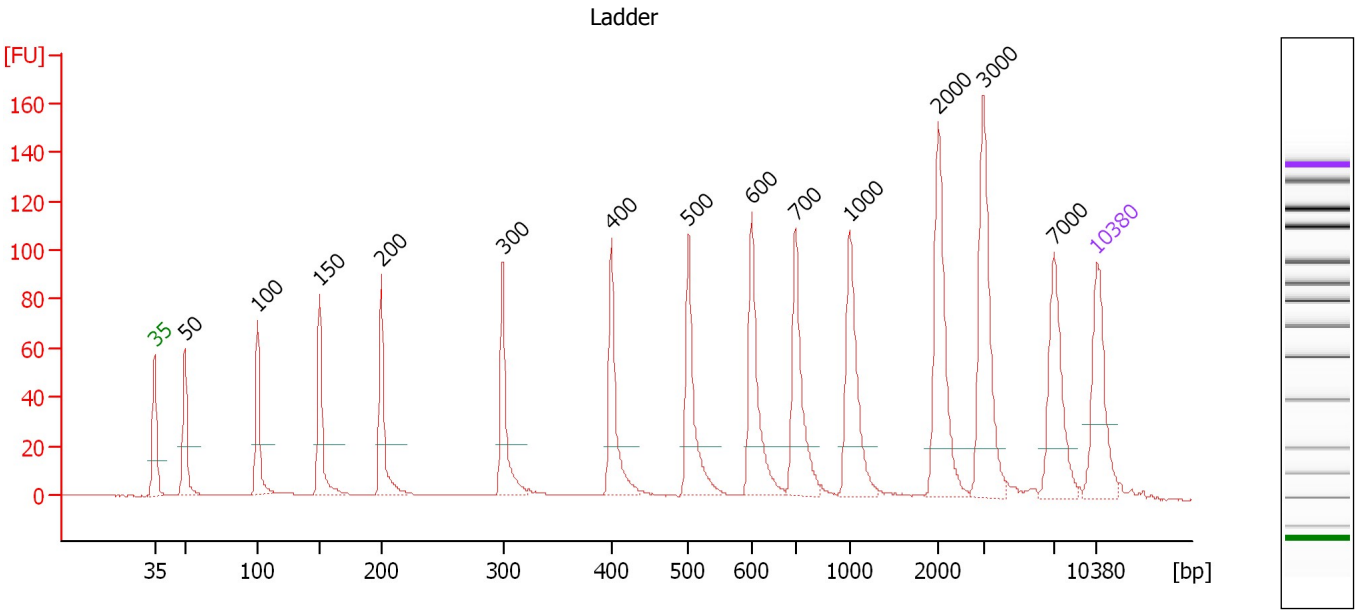
Electropherogram Summary



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	45.25
3		100	150.00	2,272.7	Ladder Peak	50.64
4		150	150.00	1,515.2	Ladder Peak	55.26
5		200	150.00	1,136.4	Ladder Peak	59.84
6		300	150.00	757.6	Ladder Peak	68.87
7		400	150.00	568.2	Ladder Peak	76.94
8		500	150.00	454.5	Ladder Peak	82.64
9		600	150.00	378.8	Ladder Peak	87.35
10		700	150.00	324.7	Ladder Peak	90.59
11		1,000	150.00	227.3	Ladder Peak	94.60
12		2,000	150.00	113.6	Ladder Peak	101.21
13		3,000	150.00	75.8	Ladder Peak	104.58
14		7,000	150.00	32.5	Ladder Peak	109.80
15		10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...analyzer\2015-08-07\2015-08-07_001_HiSeq158_Lib_Sample6.xad

Created: 8/7/2015 11:21:09 AM
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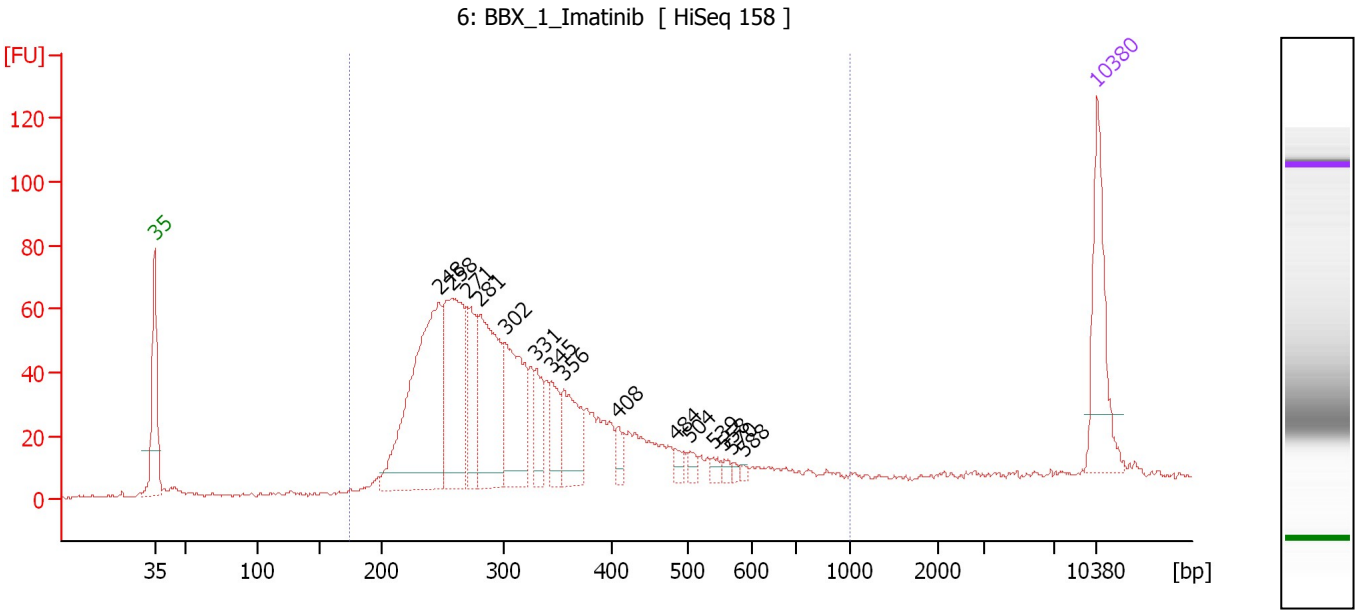
Electropherogram Summary Continued ...



Overall Results for sample 2 : 6: BBX_1_Imatinib

Number of peaks found:	15	Corr. Area 1:	1,203.8
Noise:	0.6		

Peak table for sample 2 : 6: BBX_1_Imatinib

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	248	295.00	1,805.6		64.13
3	258	188.22	1,106.0		65.06
4	271	79.40	443.7		66.26
5	281	157.96	850.5		67.19
6	302	122.06	612.9		69.01
7	331	38.47	176.2		71.35
8	345	39.25	172.4		72.51
9	356	67.54	287.4		73.39
10	408	13.09	48.6		77.42
11	484	8.39	26.3		81.72
12	504	7.76	23.3		82.83
13	539	6.56	18.5		84.47
14	558	5.46	14.8		85.35
15	570	3.60	9.6		85.93
16	588	3.19	8.2		86.77
17	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 2 : 6: BBX_1_Imatinib

From [bp]	Average Size [bp]	To [bp]	Corr. Area	Molarity [pmol/l]	Color % of Total	Size distribution in CV [%]	Conc. [pg/µl]
174	330	1,000	1,203.8	6,631.9	97	33.0	1,300.02

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Gel Image