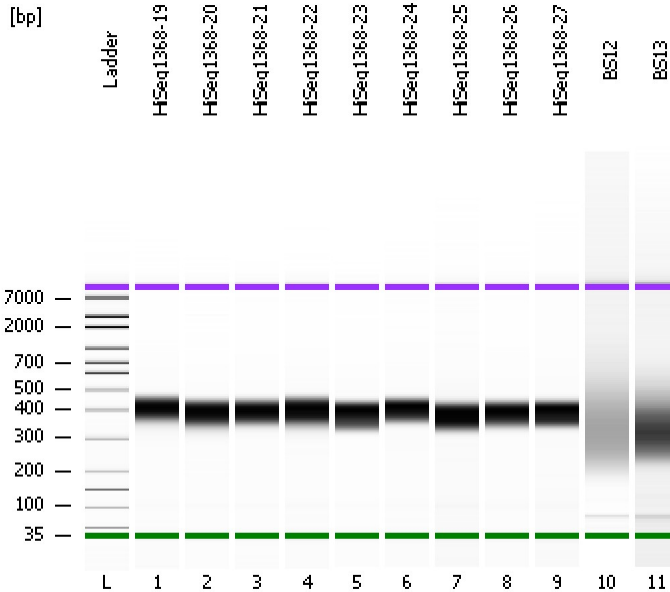


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
Modified: 5/24/2018 11:59:50 AM

Electrophoresis File Run Summary



Instrument Information:

Instrument Name: DE34903152 Firmware: C.01.069
Serial#: DE34903152 Type: G2938C

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

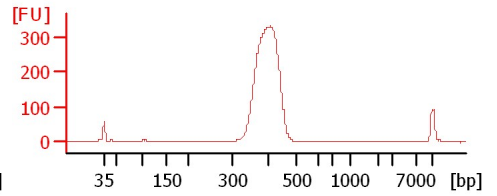
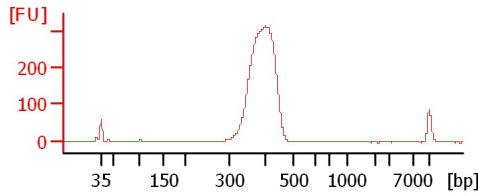
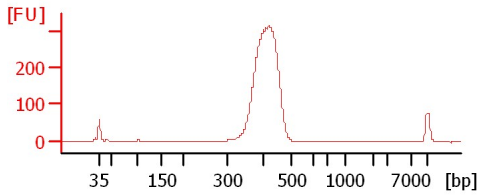
Chip Information:

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

HiSeq1368-19

HiSeq1368-20

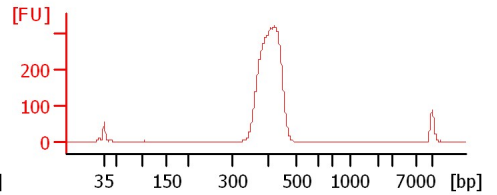
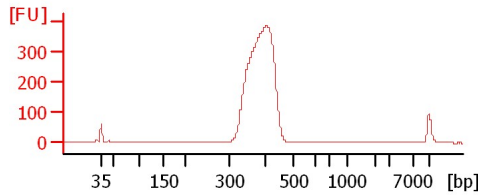
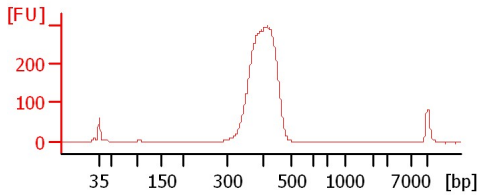
HiSeq1368-21



HiSeq1368-22

HiSeq1368-23

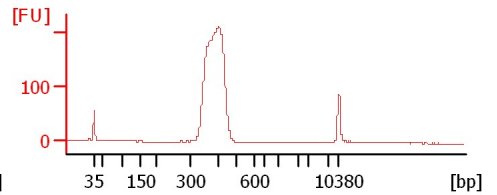
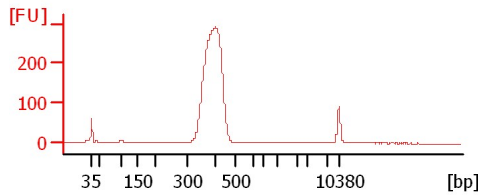
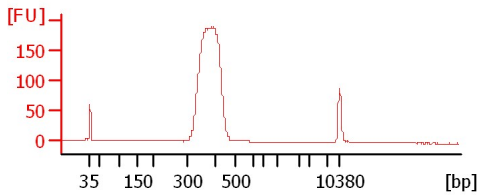
HiSeq1368-24



HiSeq1368-25

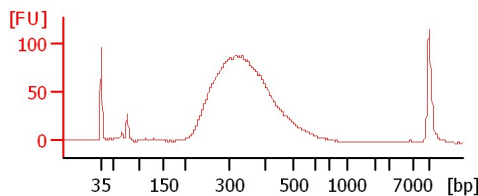
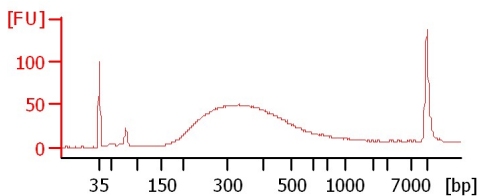
HiSeq1368-26

HiSeq1368-27



BS12

BS13



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
 Modified: 5/24/2018 11:59:50 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Rest. Digest	Status	Observation	Result Label	Result Color
HiSeq1368-19		<input type="checkbox"/>	✓			
HiSeq1368-20		<input type="checkbox"/>	✓			
HiSeq1368-21		<input type="checkbox"/>	✓			
HiSeq1368-22		<input type="checkbox"/>	✓			
HiSeq1368-23		<input type="checkbox"/>	✓			
HiSeq1368-24		<input type="checkbox"/>	✓			
HiSeq1368-25		<input type="checkbox"/>	✓			
HiSeq1368-26		<input type="checkbox"/>	✓			
HiSeq1368-27		<input type="checkbox"/>	✓			
BS12		<input type="checkbox"/>	✓			
BS13		<input type="checkbox"/>	✓			
Ladder		<input type="checkbox"/>	✓			

Chip Lot # **Reagent Kit Lot #**

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
Modified: 5/24/2018 11:59:50 AM

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

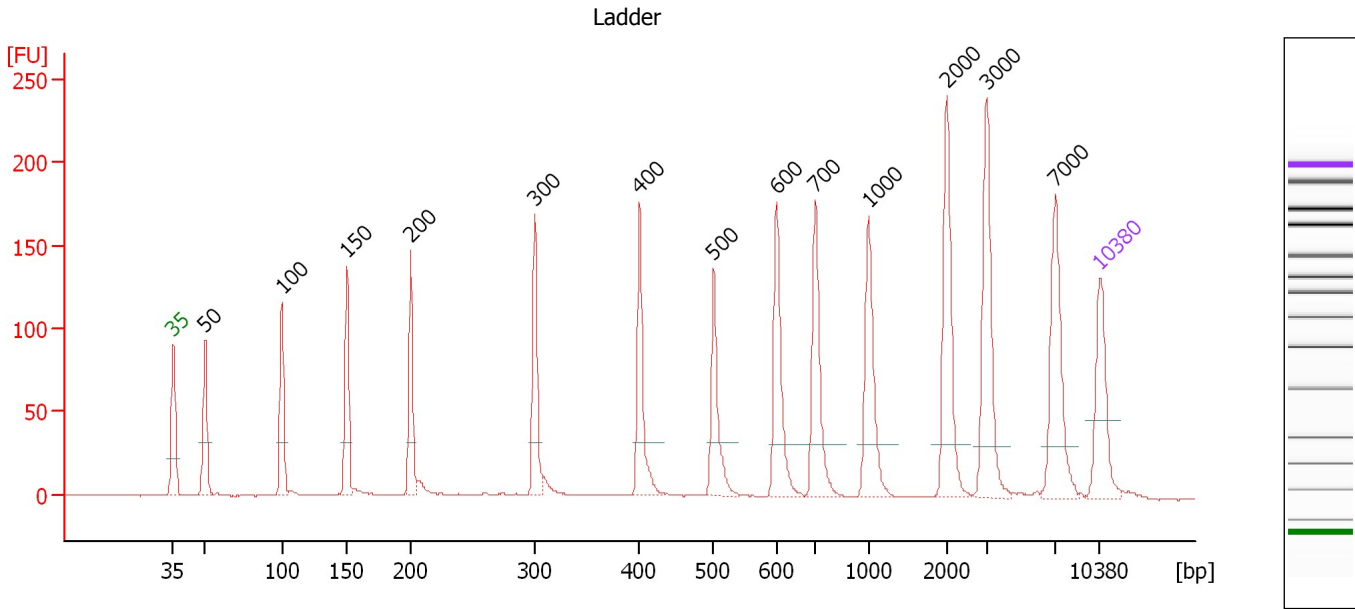
Ladder

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:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
 Modified: 5/24/2018 11:59:50 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.4

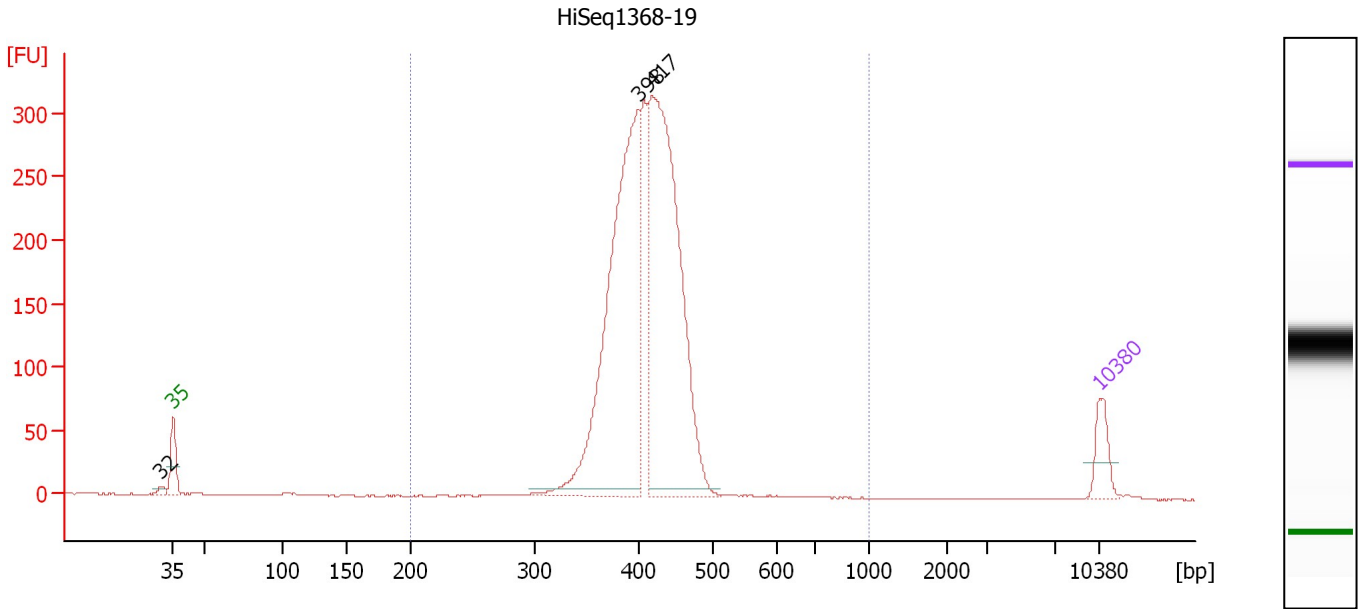
Peak table for Ladder

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	45.39
3	100	150.00	2,272.7	Ladder Peak	51.21
4	150	150.00	1,515.2	Ladder Peak	56.13
5	200	150.00	1,136.4	Ladder Peak	60.95
6	300	150.00	757.6	Ladder Peak	70.31
7	400	150.00	568.2	Ladder Peak	78.24
8	500	150.00	454.5	Ladder Peak	83.83
9	600	150.00	378.8	Ladder Peak	88.60
10	700	150.00	324.7	Ladder Peak	91.51
11	1,000	150.00	227.3	Ladder Peak	95.52
12	2,000	150.00	113.6	Ladder Peak	101.44
13	3,000	150.00	75.8	Ladder Peak	104.41
14	7,000	150.00	32.5	Ladder Peak	109.66
15	10,380	75.00	10.9	Upper Marker	113.00

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
 Modified: 5/24/2018 11:59:50 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : HiSeq1368-19

Number of peaks found: 3 Corr. Area 1: 2,459.8
 Noise: 0.4

Peak table for sample 1 : HiSeq1368-19

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.08
2	35	125.00	5,411.3	Lower Marker	43.00
3	398	1,406.94	5,353.2		78.10
4	417	1,445.24	5,256.9		79.16
5	10,380	75.00	10.9	Upper Marker	113.00

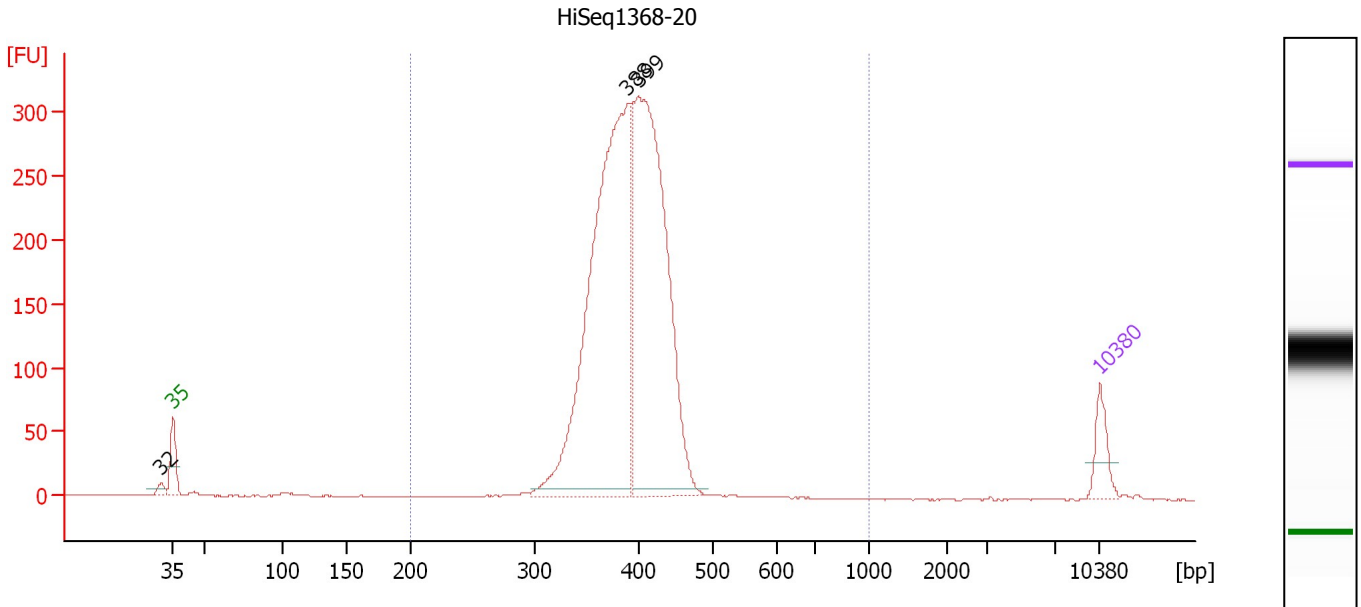
Region table for sample 1 : HiSeq1368-19

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	410	3,219.10	2,459.8	11,996.8	99	8.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
 Modified: 5/24/2018 11:59:50 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 : HiSeq1368-20

Number of peaks found: 3 Corr. Area 1: 2,710.1
 Noise: 0.3

Peak table for sample 2 : HiSeq1368-20

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.07
2	35	125.00	5,411.3	Lower Marker	43.00
3	388	1,796.50	7,014.9		77.29
4	399	1,553.92	5,898.6		78.17
5	10,380	75.00	10.9	Upper Marker	113.00

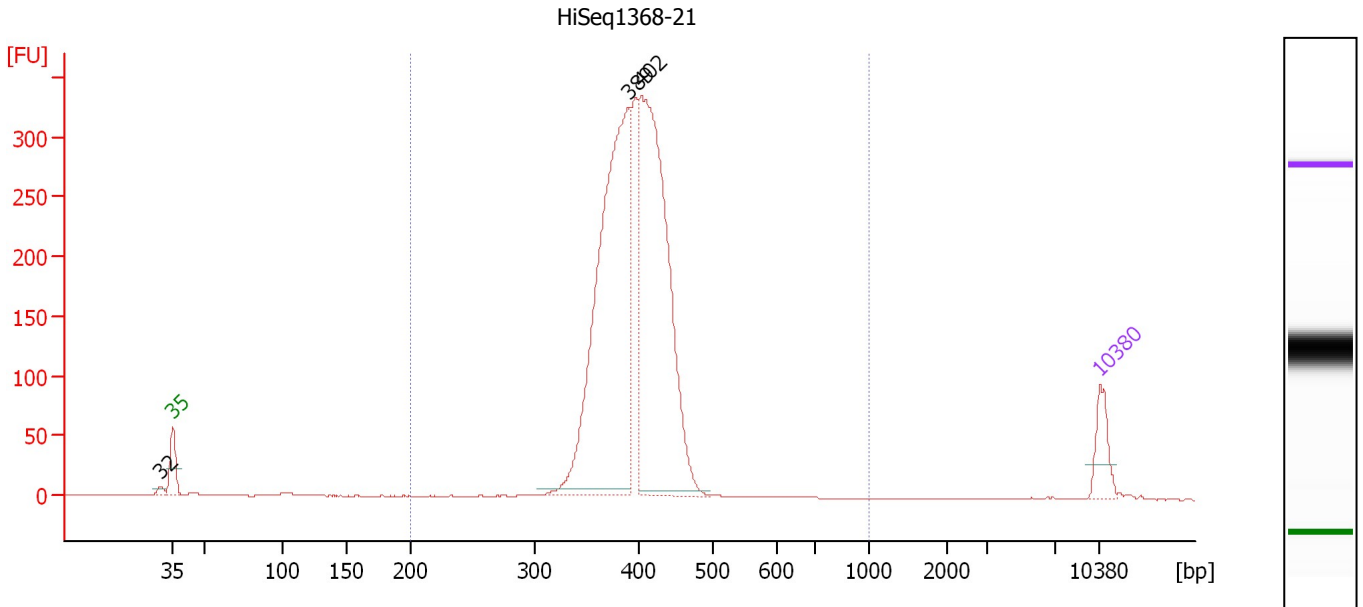
Region table for sample 2 : HiSeq1368-20

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	391	3,559.75	2,710.1	13,907.7	98	8.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
 Modified: 5/24/2018 11:59:50 AM

Electropherogram Summary Continued ...



Overall Results for sample 3 : HiSeq1368-21

Number of peaks found: 3 Corr. Area 1: 2,615.9
 Noise: 0.3

Peak table for sample 3 : HiSeq1368-21

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.06
2	35	125.00	5,411.3	Lower Marker	43.00
3	389	1,405.13	5,470.8		77.38
4	402	1,313.73	4,957.6		78.32
5	10,380	75.00	10.9	Upper Marker	113.00

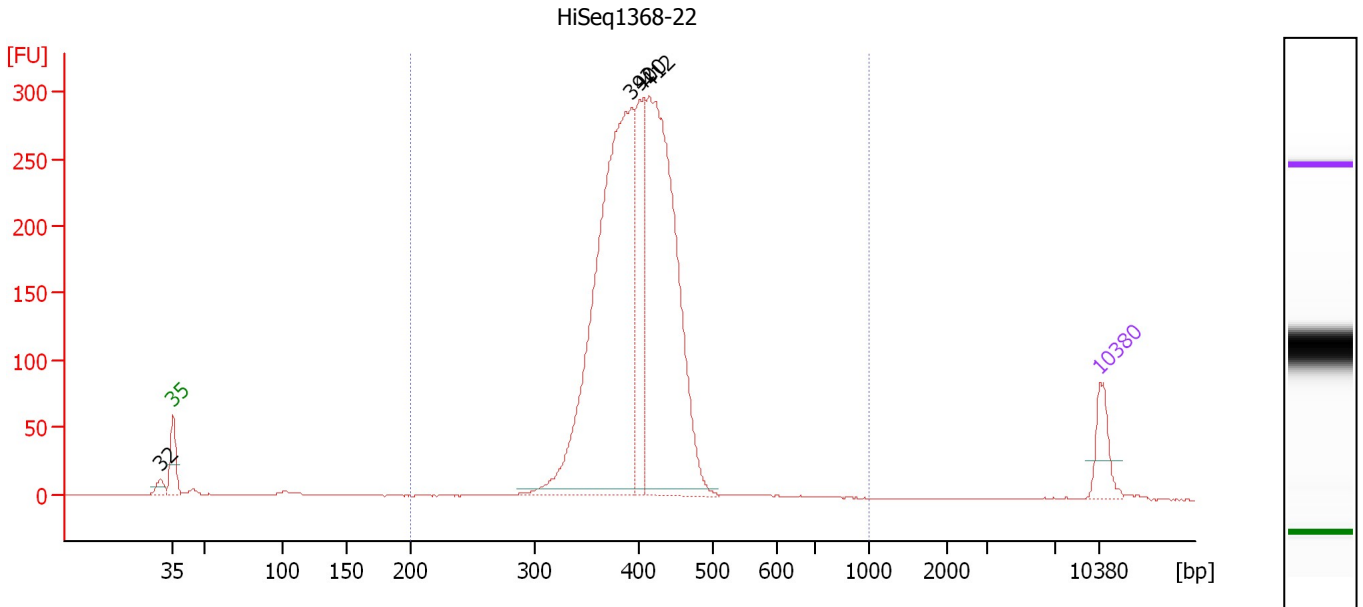
Region table for sample 3 : HiSeq1368-21

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	397	3,119.80	2,615.9	12,009.8	98	8.5

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
 Modified: 5/24/2018 11:59:50 AM

Electropherogram Summary Continued ...



Overall Results for sample 4 : HiSeq1368-22

Number of peaks found: 4 Corr. Area 1: 2,695.8
 Noise: 0.3

Peak table for sample 4 : HiSeq1368-22

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	32	0.00	0.0		42.04
2	35	125.00	5,411.3	Lower Marker	43.00
3	392	1,588.55	6,141.6		77.60
4	400	354.44	1,341.8		78.25
5	412	1,287.46	4,735.0		78.91
6	10,380	75.00	10.9	Upper Marker	113.00

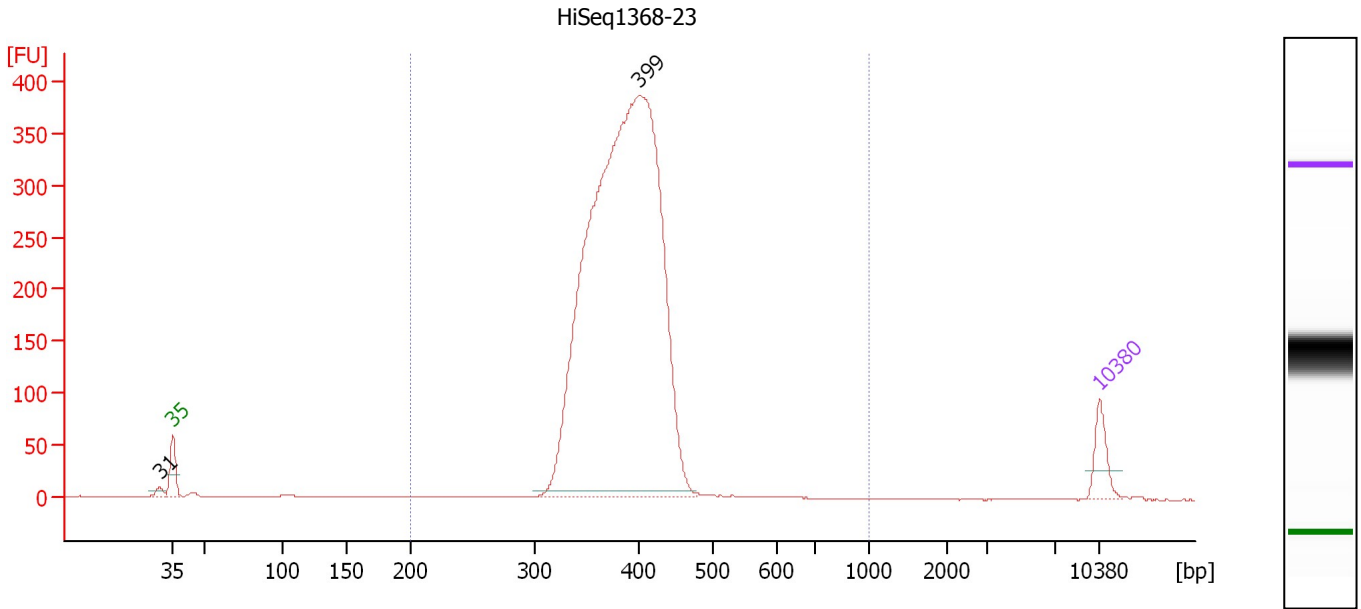
Region table for sample 4 : HiSeq1368-22

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	400	3,349.92	2,695.8	12,831.7	98	10.0

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
 Modified: 5/24/2018 11:59:50 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : HiSeq1368-23

Number of peaks found: 2 Corr. Area 1: 3,366.6
 Noise: 0.3

Peak table for sample 5 : HiSeq1368-23

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.97
2	35	125.00	5,411.3	Lower Marker	43.00
3	399	3,894.86	14,804.9		78.13
4	10,380	75.00	10.9	Upper Marker	113.00

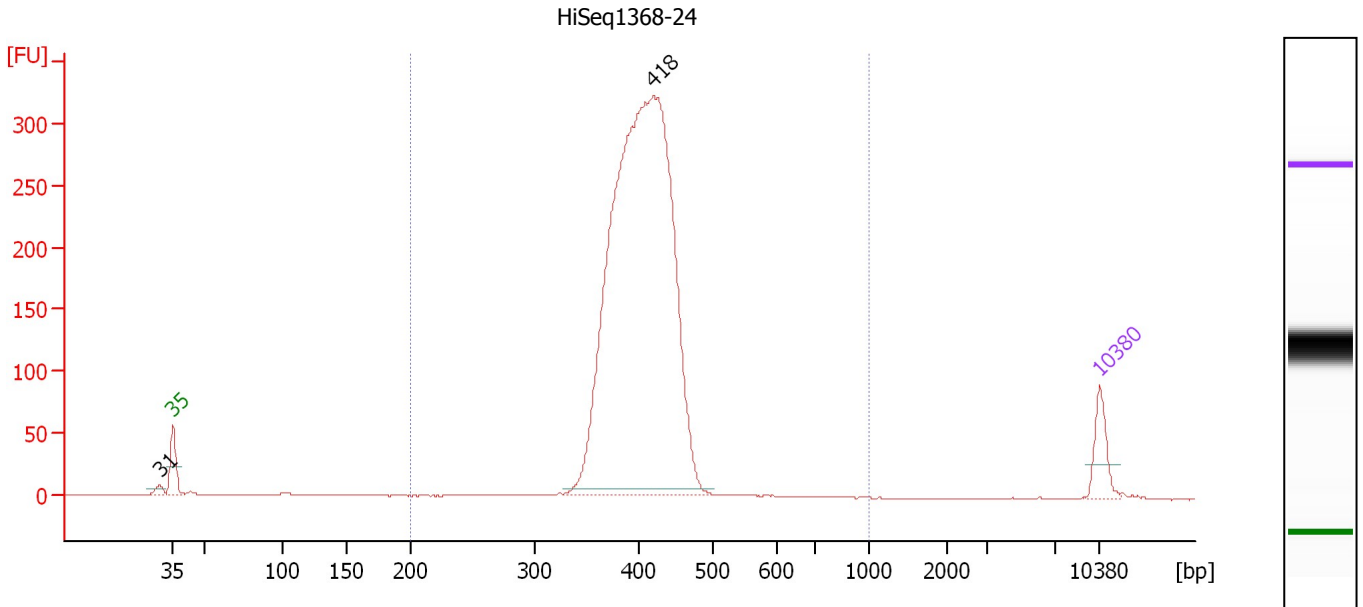
Region table for sample 5 : HiSeq1368-23

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	387	4,112.18	3,366.6	16,242.3	99	9.2

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
 Modified: 5/24/2018 11:59:50 AM

Electropherogram Summary Continued ...



Overall Results for sample 6 : HiSeq1368-24

Number of peaks found: 2 Corr. Area 1: 2,400.6
 Noise: 0.2

Peak table for sample 6 : HiSeq1368-24

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.95
2	35	125.00	5,411.3	Lower Marker	43.00
3	418	2,923.12	10,590.5		79.26
4	10,380	75.00	10.9	Upper Marker	113.00

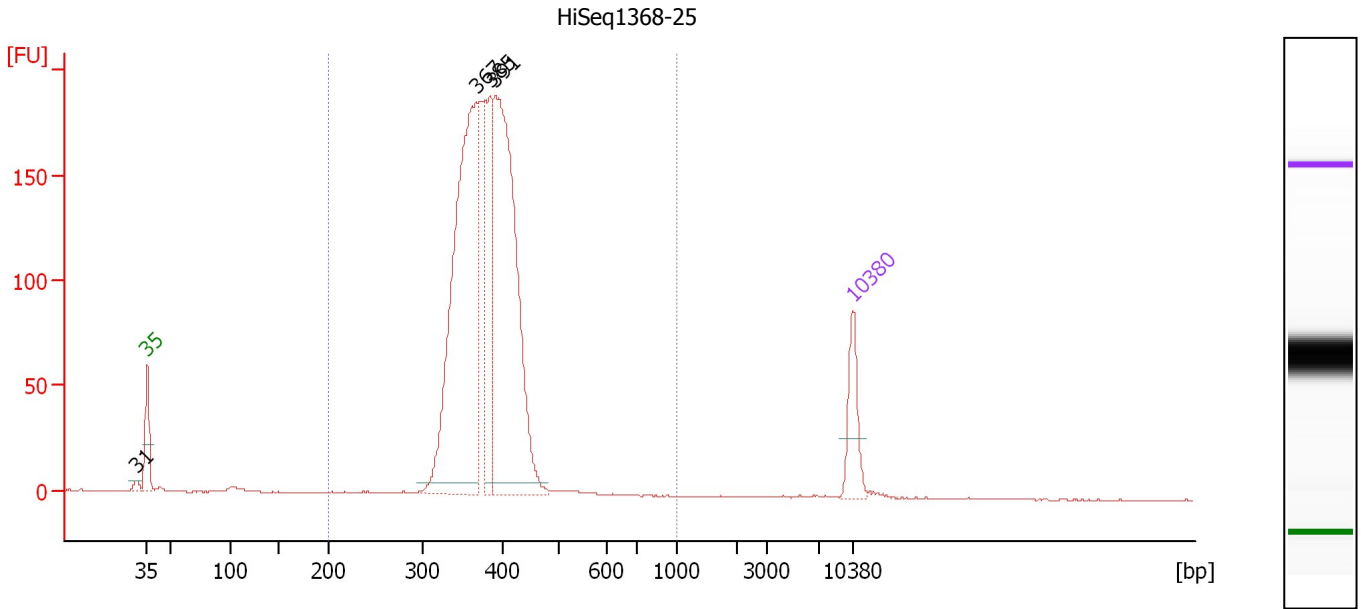
Region table for sample 6 : HiSeq1368-24

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	406	3,082.98	2,400.6	11,618.2	98	9.3

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
 Modified: 5/24/2018 11:59:50 AM

Electropherogram Summary Continued ...



Overall Results for sample 7 : HiSeq1368-25

Number of peaks found: 4 Corr. Area 1: 1,725.9
 Noise: 0.2

Peak table for sample 7 : HiSeq1368-25

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.89
2	35	125.00	5,411.3	Lower Marker	43.00
3	367	911.77	3,764.5		75.62
4	385	231.94	912.8		77.05
5	391	824.65	3,195.7		77.52
6	10,380	75.00	10.9	Upper Marker	113.00

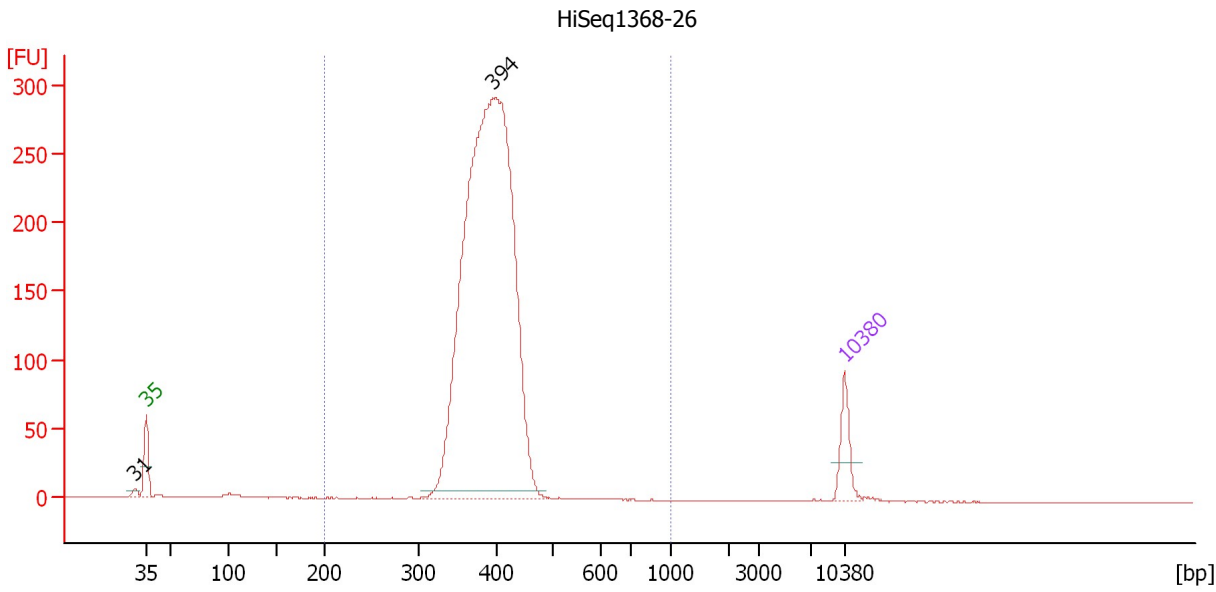
Region table for sample 7 : HiSeq1368-25

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	381	2,220.13	1,725.9	8,901.2	99	8.9

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
 Modified: 5/24/2018 11:59:50 AM

Electropherogram Summary Continued ...



Overall Results for sample 8 : HiSeq1368-26

Number of peaks found: 2 Corr. Area 1: 2,272.2
 Noise: 0.3

Peak table for sample 8 : HiSeq1368-26

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.88
2	35	125.00	5,411.3	Lower Marker	43.00
3	394	2,837.70	10,923.2		77.73
4	10,380	75.00	10.9	Upper Marker	113.00

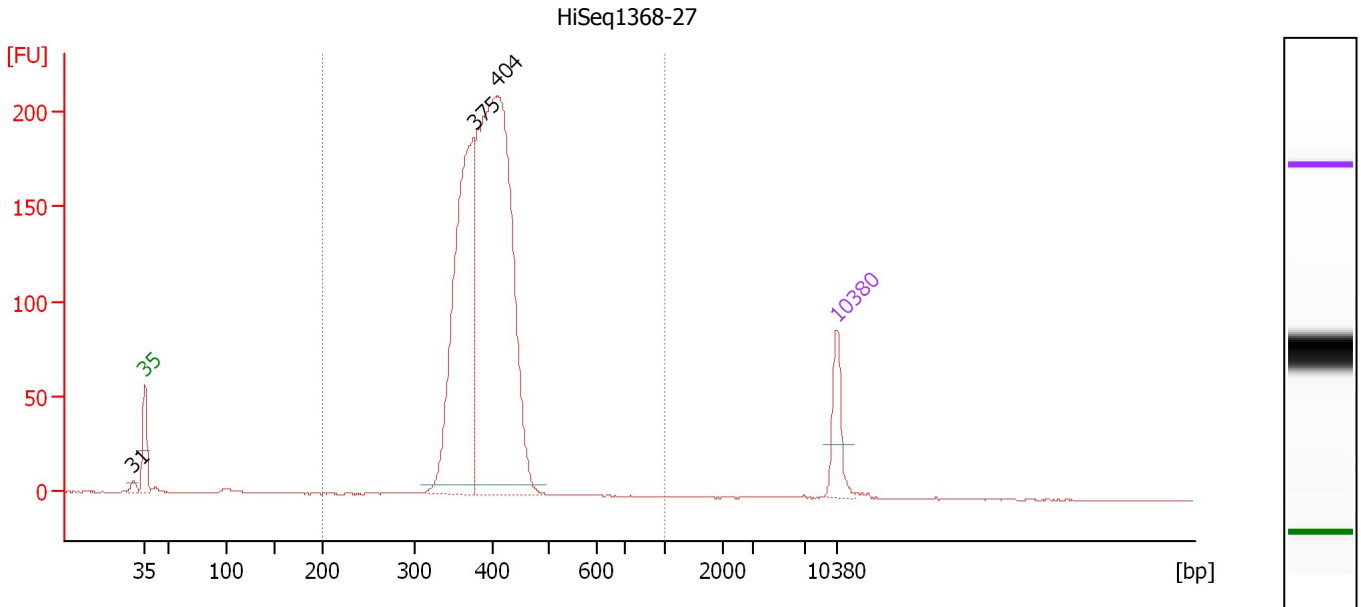
Region table for sample 8 : HiSeq1368-26

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of lor Total	Size distribution in CV [%]
200	1,000	390	2,927.13	2,272.2	11,437.6	98	8.1

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
 Modified: 5/24/2018 11:59:50 AM

Electropherogram Summary Continued ...



Overall Results for sample 9 : HiSeq1368-27

Number of peaks found: 3 Corr. Area 1: 1,690.4
 Noise: 0.2

Peak table for sample 9 : HiSeq1368-27

Peak	Size [bp]	Conc. [pg/μl]	Molarity [pmol/l]	Observations	Aligned Migration Time [s]
1	31	0.00	0.0		41.81
2	35	125.00	5,411.3	Lower Marker	43.00
3	375	758.94	3,067.3		76.25
4	404	1,392.75	5,224.3		78.46
5	10,380	75.00	10.9	Upper Marker	113.00

Region table for sample 9 : HiSeq1368-27

From [bp]	To [bp]	Average Size [bp]	Conc. [pg/μl]	Corr. Area	Molarity [pmol/l]	Co % of Ior Total	Size distribution in CV [%]
200	1,000	391	2,217.49	1,690.4	8,644.1	99	7.9

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-05-24\2018-05-24_002.xad

Created: 5/24/2018 11:14:07 AM
Modified: 5/24/2018 11:59:50 AM

Gel Image

