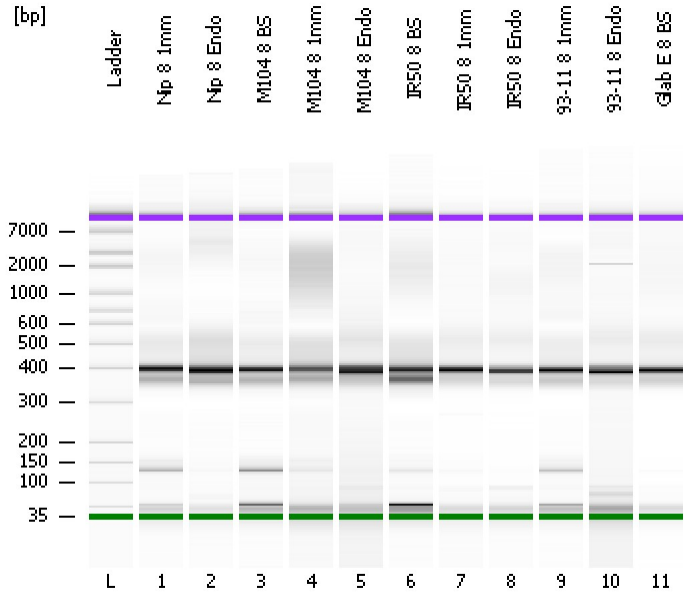
Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

## Electrophoresis File Run Summary




Instrument Information:

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

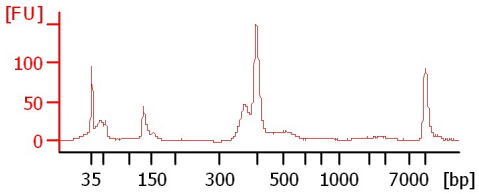
Assay Information:

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

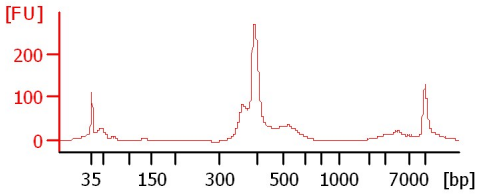
Chip Information:

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

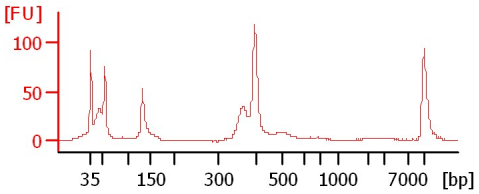
**Nip 8 1mm**



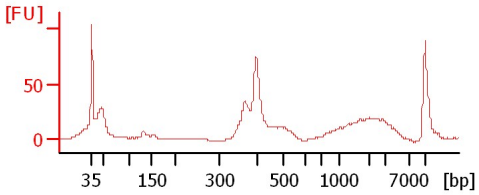

**Nip 8 Endo**



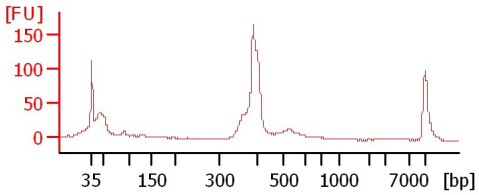

**M104 8 BS**



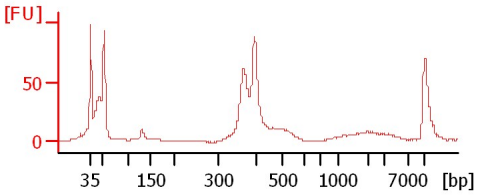

**M104 8 1mm**



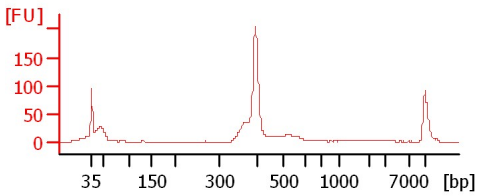

**M104 8 Endo**



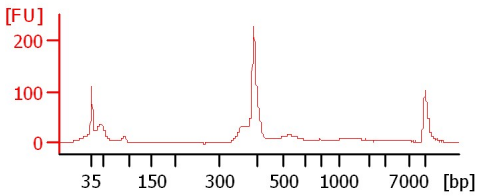

**IR50 8 BS**



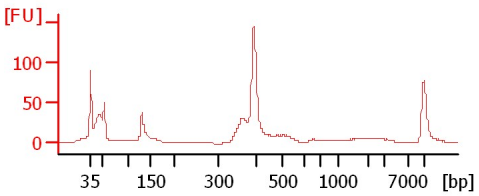

**IR50 8 1mm**



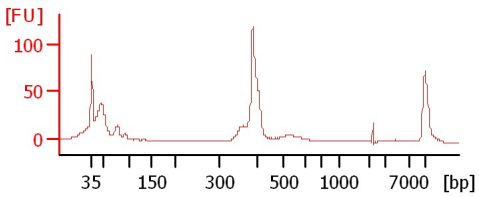

**IR50 8 Endo**



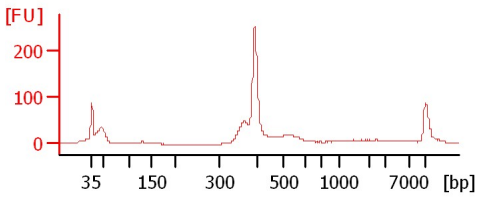

**93-11 8 1mm**




**93-11 8 Endo**




**Glab E 8 BS**



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

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

| Sample Name | Sample Comment | Rest. Digest | Status | Observation | Result Label | Result Color |
|---|---|---|---|---|---|---|
| Nip 8 1mm | | ☐ | ✔ | | | |
| Nip 8 Endo | | ☐ | ✔ | | | |
| M104 8 BS | | ☐ | ✔ | | | |
| M104 8 1mm | | ☐ | ✔ | | | |
| M104 8 Endo | | ☐ | ✔ | | | |
| IR50 8 BS | | ☐ | ✔ | | | |
| IR50 8 1mm | | ☐ | ✔ | | | |
| IR50 8 Endo | | ☐ | ✔ | | | |
| 93-11 8 1mm | | ☐ | ✔ | | | |
| 93-11 8 Endo | | ☐ | ✔ | | | |
| Glab E 8 BS | | ☐ | ✔ | | | |
| Ladder | | ☐ | ✔ | | | |

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

**Chip Comments :**

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 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

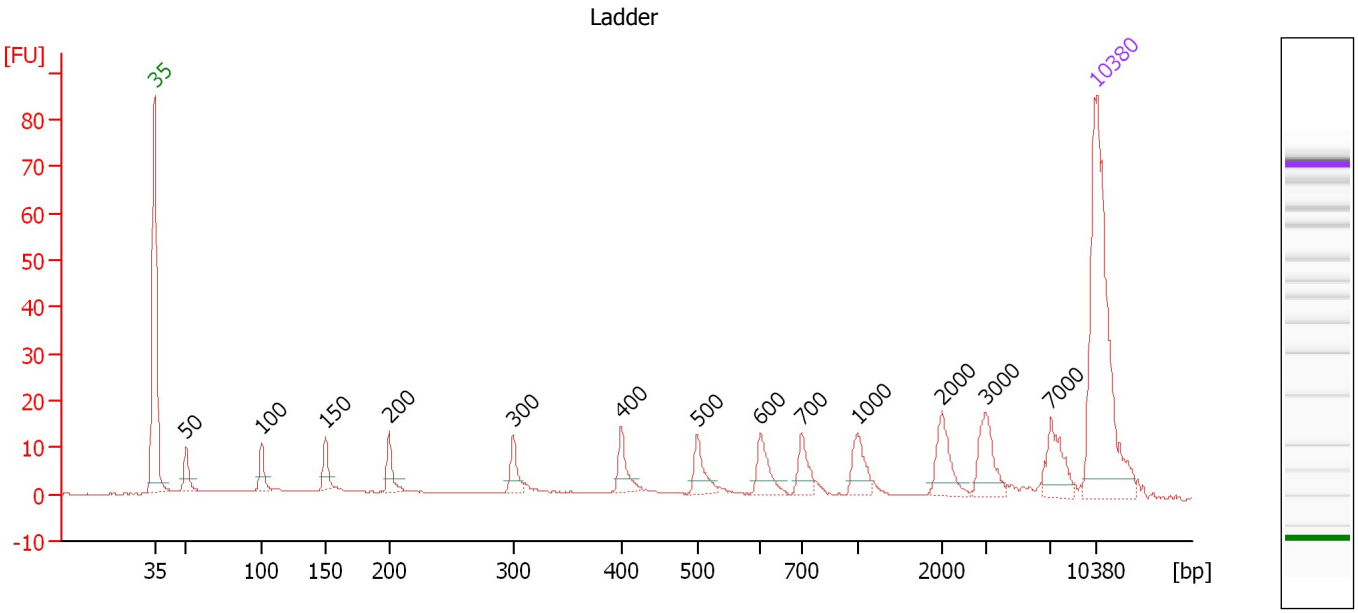
### Ladder

| 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:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

## Electropherogram Summary




## Overall Results for Ladder

Noise: 0.1

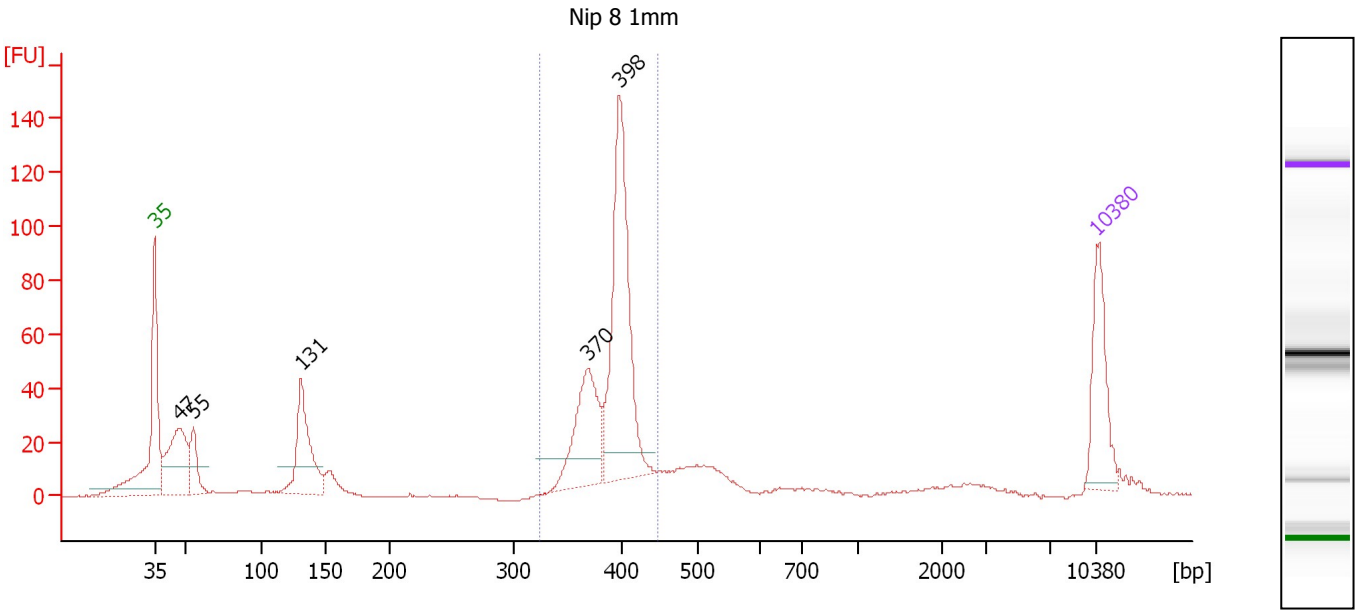
## Peak table for Ladder

| 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:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

**Electropherogram Summary Continued ...**




**Setpoint Deviations for sample 1 : Nip 8 1mm**

Height Threshold [FU] : 10

**Overall Results for sample 1 : Nip 8 1mm**

| | | | |
|---|---|---|---|
| Number of peaks found: | 5 | Corr. Area 1: | 444.1 |
| Noise: | 0.1 | | |

**Peak table for sample 1 : Nip 8 1mm**

| Peak | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations |
|---|---|---|---|---|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 47 | 157.39 | 5,070.1 | |
| 3 | 55 | 56.05 | 1,538.8 | |
| 4 | 131 | 142.40 | 1,649.0 | |
| 5 | 370 | 137.94 | 565.4 | |
| 6 | 398 | 296.86 | 1,129.2 | |
| 7 | 10,380 | 75.00 | 10.9 | Upper Marker |

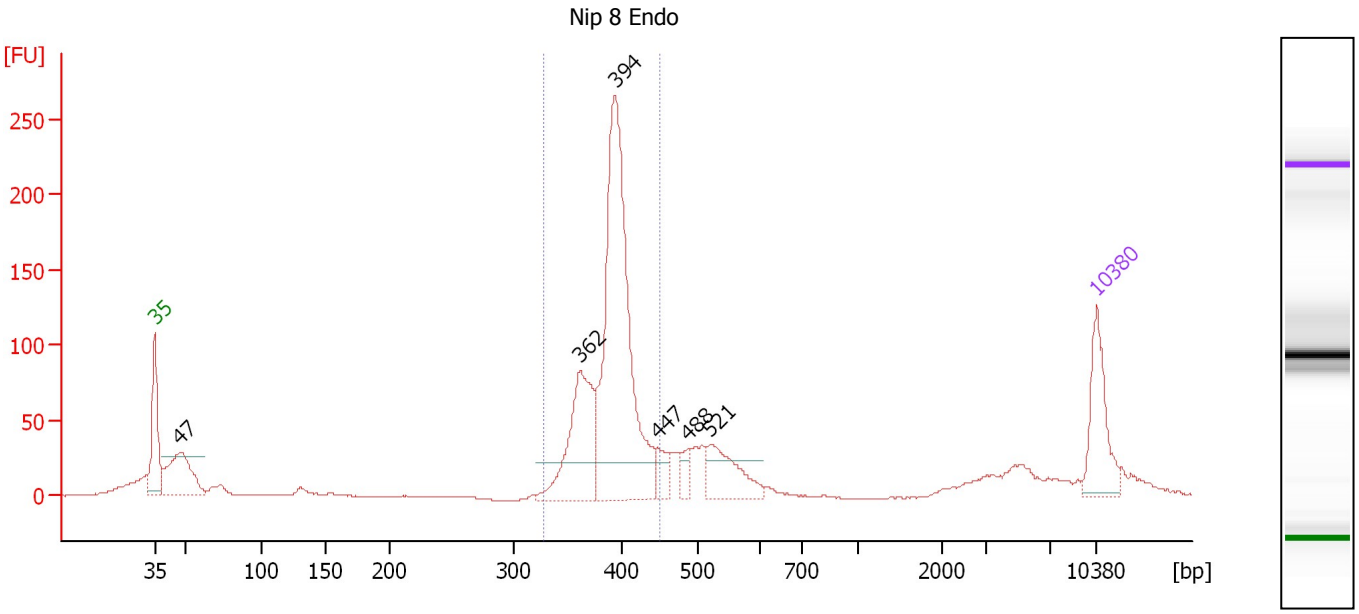
**Region table for sample 1 : Nip 8 1mm**

| From [bp] | To [bp] | Average Size [bp] | Molarity [pmol/l] | Conc. [pg/µl] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|---|---|---|---|---|---|---|---|---|
| 325 | 448 | 392 | 1,934.0 | 498.87 | 444.1 | 53 | 5.3 | |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

**Electropherogram Summary Continued ...**




**Setpoint Deviations for sample 2 : Nip 8 Endo**

Height Threshold [FU] : 25

**Overall Results for sample 2 : Nip 8 Endo**

| | | | |
|---|---|---|---|
| Number of peaks found: | 6 | Corr. Area 1: | 953.3 |
| Noise: | 0.3 | | |

**Peak table for sample 2 : Nip 8 Endo**

| Peak | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations |
|---|---|---|---|---|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 47 | 145.21 | 4,674.2 | |
| 3 | 362 | 188.37 | 788.0 | |
| 4 | 394 | 534.08 | 2,052.5 | |
| 5 | 447 | 29.00 | 98.2 | |
| 6 | 488 | 21.69 | 67.3 | |
| 7 | 521 | 86.20 | 250.6 | |
| 8 | 10,380 | 75.00 | 10.9 | Upper Marker |

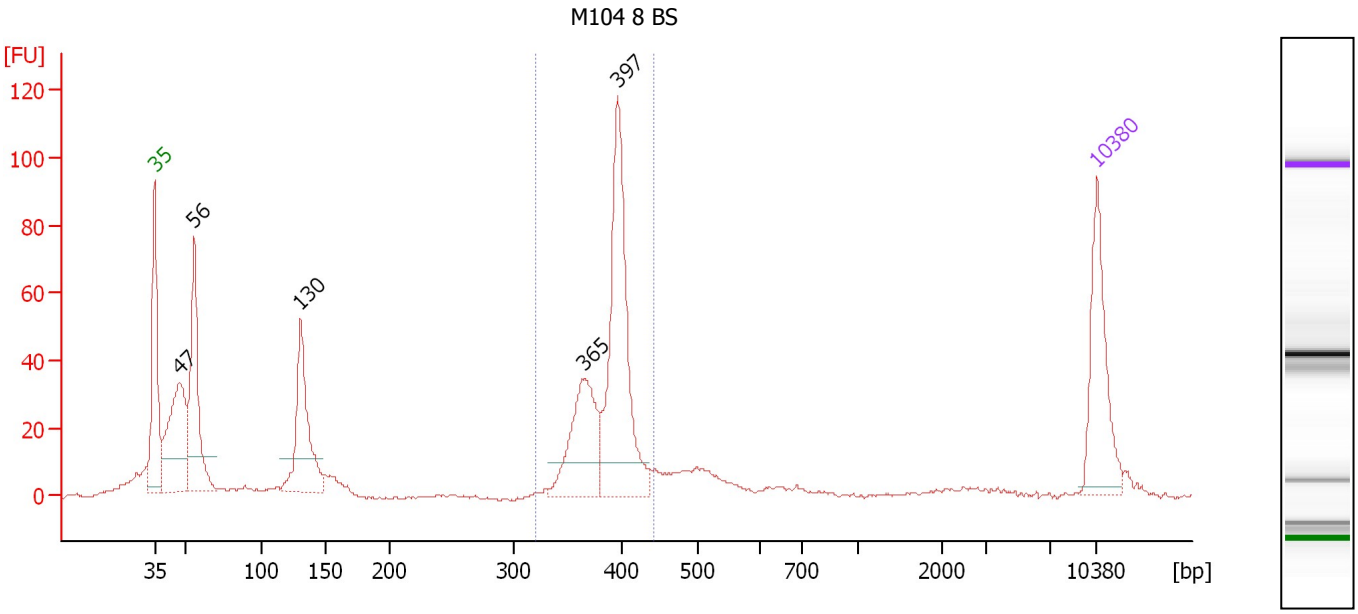
**Region table for sample 2 : Nip 8 Endo**

| From [bp] | To [bp] | Average Size [bp] | Molarity [pmol/l] | Conc. [pg/µl] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|---|---|---|---|---|---|---|---|---|
| 329 | 450 | 391 | 2,725.5 | 701.78 | 953.3 | 59 | 5.6 | |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

**Electropherogram Summary Continued ...**




**Setpoint Deviations for sample 3 : M104 8 BS**

Height Threshold [FU] : 10

**Overall Results for sample 3 : M104 8 BS**

| | | | |
|---|---|---|---|
| Number of peaks found: | 5 | Corr. Area 1: | 334.1 |
| Noise: | 0.4 | | |

**Peak table for sample 3 : M104 8 BS**

| Peak | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations |
|---|---|---|---|---|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 47 | 170.26 | 5,500.1 | |
| 3 | 56 | 164.28 | 4,483.5 | |
| 4 | 130 | 137.14 | 1,593.0 | |
| 5 | 365 | 110.96 | 460.2 | |
| 6 | 397 | 221.35 | 845.6 | |
| 7 | 10,380 | 75.00 | 10.9 | Upper Marker |

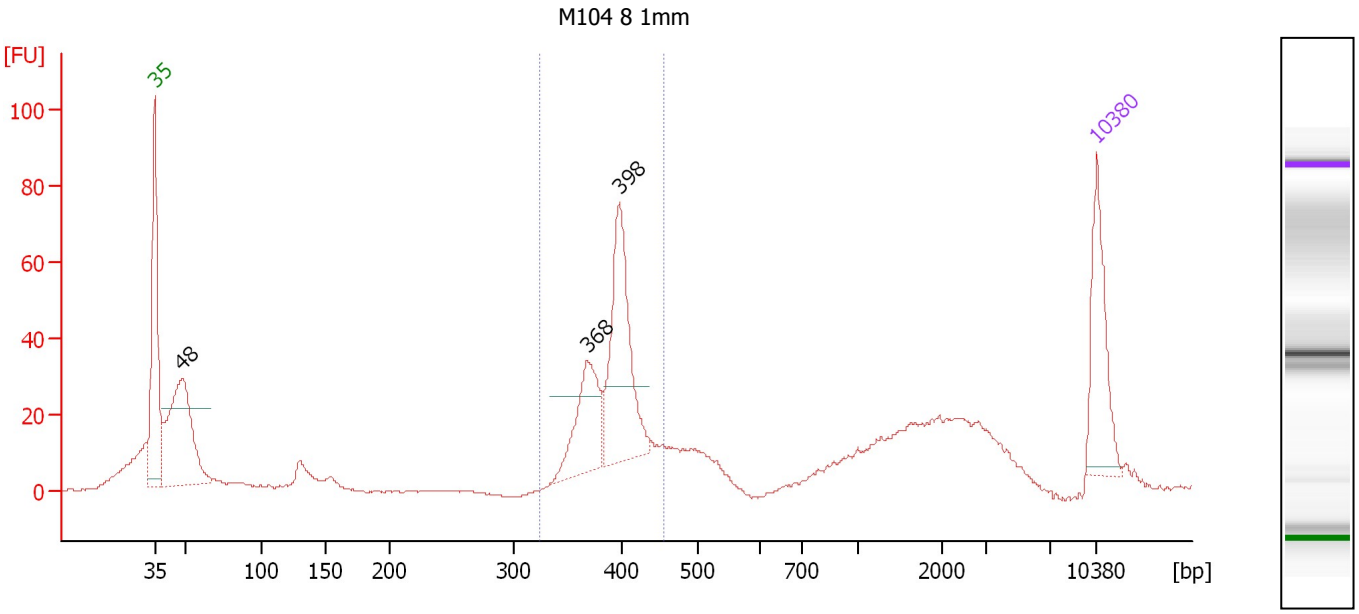
**Region table for sample 3 : M104 8 BS**

| From [bp] | To [bp] | Average Size [bp] | Molarity [pmol/l] | Conc. [pg/µl] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|---|---|---|---|---|---|---|---|---|
| 322 | 444 | 388 | 1,321.6 | 338.19 | 334.1 | 39 | 5.4 | |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

**Electropherogram Summary Continued ...**




**Setpoint Deviations for sample 4 : M104 8 1mm**

Height Threshold [FU] : 20

**Overall Results for sample 4 : M104 8 1mm**

| | | | |
|---|---|---|---|
| Number of peaks found: | 3 | Corr. Area 1: | 297.5 |
| Noise: | 0.2 | | |

**Peak table for sample 4 : M104 8 1mm**

| Peak | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations |
|---|---|---|---|---|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 48 | 238.33 | 7,487.7 | |
| 3 | 368 | 98.40 | 404.7 | |
| 4 | 398 | 175.17 | 666.1 | |
| 5 | 10,380 | 75.00 | 10.9 | Upper Marker |

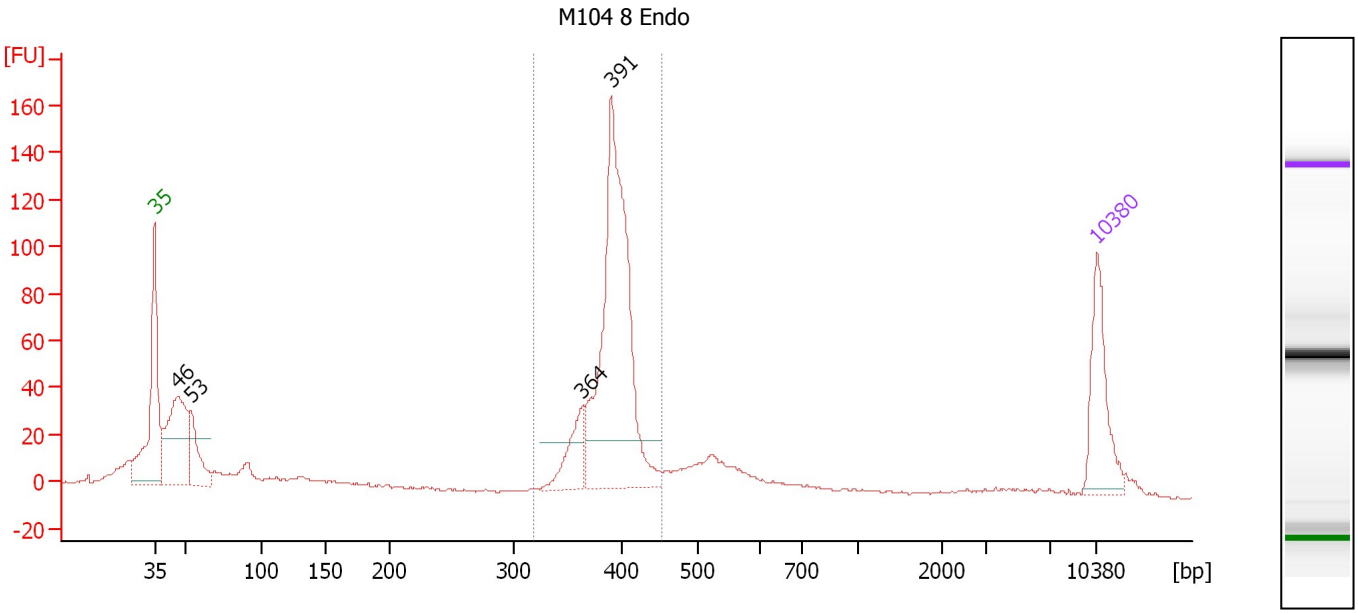
**Region table for sample 4 : M104 8 1mm**

| From [bp] | To [bp] | Average Size [bp] | Molarity [pmol/l] | Conc. [pg/µl] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|---|---|---|---|---|---|---|---|---|
| 326 | 455 | 393 | 1,420.0 | 367.25 | 297.5 | 36 | 6.2 | |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

**Electropherogram Summary Continued ...**




**Setpoint Deviations for sample 5 : M104 8 Endo**

Height Threshold [FU] : 20

**Overall Results for sample 5 : M104 8 Endo**

| | | | |
|---|---|---|---|
| Number of peaks found: | 4 | Corr. Area 1: | 552.3 |
| Noise: | 0.5 | | |

**Peak table for sample 5 : M104 8 Endo**

| Peak | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations |
|---|---|---|---|---|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 46 | 200.53 | 6,561.4 | |
| 3 | 53 | 75.67 | 2,150.4 | |
| 4 | 364 | 57.33 | 238.7 | |
| 5 | 391 | 431.45 | 1,672.6 | |
| 6 | 10,380 | 75.00 | 10.9 | Upper Marker |

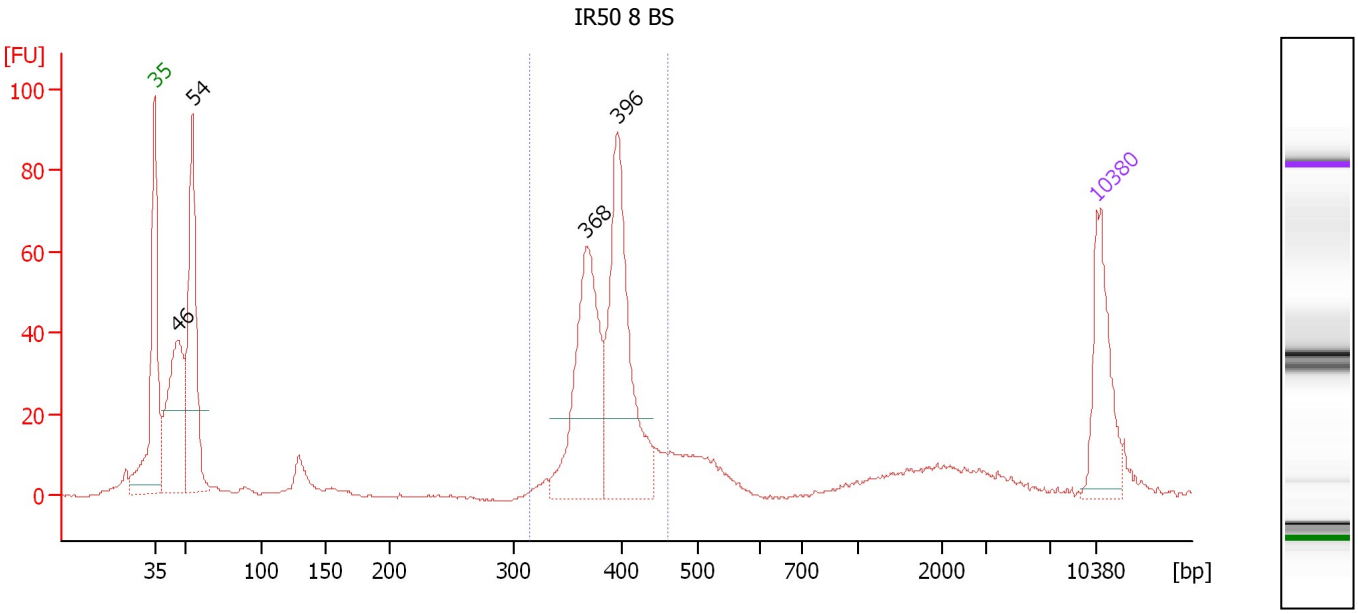
**Region table for sample 5 : M104 8 Endo**

| From [bp] | To [bp] | Average Size [bp] | Molarity [pmol/l] | Conc. [pg/µl] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|---|---|---|---|---|---|---|---|---|
| 320 | 453 | 392 | 1,934.5 | 499.20 | 552.3 | 53 | 5.1 | |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

**Electropherogram Summary Continued ...**




**Setpoint Deviations for sample 6 : IR50 8 BS**

Height Threshold [FU] : 20

**Overall Results for sample 6 : IR50 8 BS**

| | | | |
|---|---|---|---|
| Number of peaks found: | 4 | Corr. Area 1: | 396.1 |
| Noise: | 0.3 | | |

**Peak table for sample 6 : IR50 8 BS**

| Peak | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations |
|---|---|---|---|---|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 46 | 219.05 | 7,194.6 | |
| 3 | 54 | 247.43 | 6,894.2 | |
| 4 | 368 | 219.85 | 905.1 | |
| 5 | 396 | 246.41 | 941.9 | |
| 6 | 10,380 | 75.00 | 10.9 | Upper Marker |

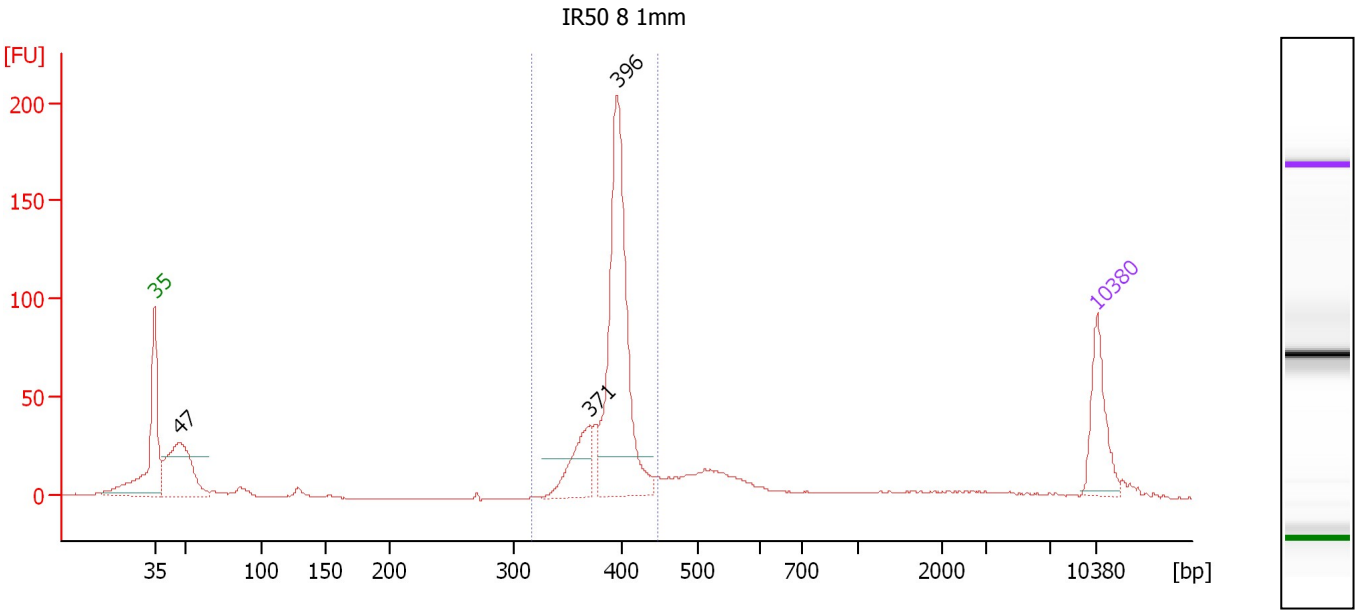
**Region table for sample 6 : IR50 8 BS**

| From [bp] | To [bp] | Average Size [bp] | Molarity [pmol/l] | Conc. [pg/µl] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|---|---|---|---|---|---|---|---|---|
| 315 | 462 | 388 | 1,863.2 | 475.58 | 396.1 | 45 | 6.6 | |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

**Electropherogram Summary Continued ...**




**Setpoint Deviations for sample 7 : IR50 8 1mm**

Height Threshold [FU] : 20

**Overall Results for sample 7 : IR50 8 1mm**

| | | | |
|---|---|---|---|
| Number of peaks found: | 3 | Corr. Area 1: | 520.7 |
| Noise: | 0.3 | | |

**Peak table for sample 7 : IR50 8 1mm**

| Peak | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations |
|---|---|---|---|---|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 47 | 222.66 | 7,164.1 | |
| 3 | 371 | 98.35 | 402.0 | |
| 4 | 396 | 441.47 | 1,687.8 | |
| 5 | 10,380 | 75.00 | 10.9 | Upper Marker |

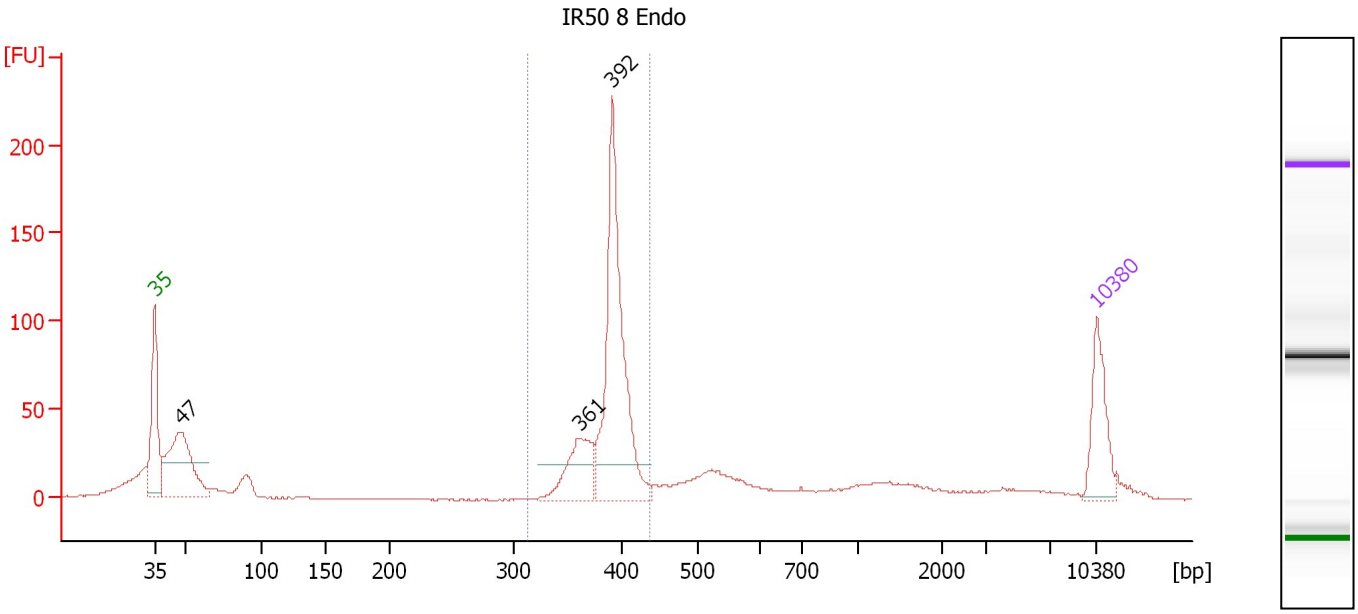
**Region table for sample 7 : IR50 8 1mm**

| From [bp] | To [bp] | Average Size [bp] | Molarity [pmol/l] | Conc. [pg/µl] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|---|---|---|---|---|---|---|---|---|
| 317 | 447 | 392 | 2,195.8 | 567.82 | 520.7 | 60 | 4.8 | |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

**Electropherogram Summary Continued ...**




**Setpoint Deviations for sample 8 : IR50 8 Endo**

Height Threshold [FU] : 20

**Overall Results for sample 8 : IR50 8 Endo**

| | | | |
|---|---|---|---|
| Number of peaks found: | 3 | Corr. Area 1: | 497.4 |
| Noise: | 0.3 | | |

**Peak table for sample 8 : IR50 8 Endo**

| Peak | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations |
|---|---|---|---|---|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 47 | 259.71 | 8,290.1 | |
| 3 | 361 | 108.84 | 456.5 | |
| 4 | 392 | 390.68 | 1,511.3 | |
| 5 | 10,380 | 75.00 | 10.9 | Upper Marker |

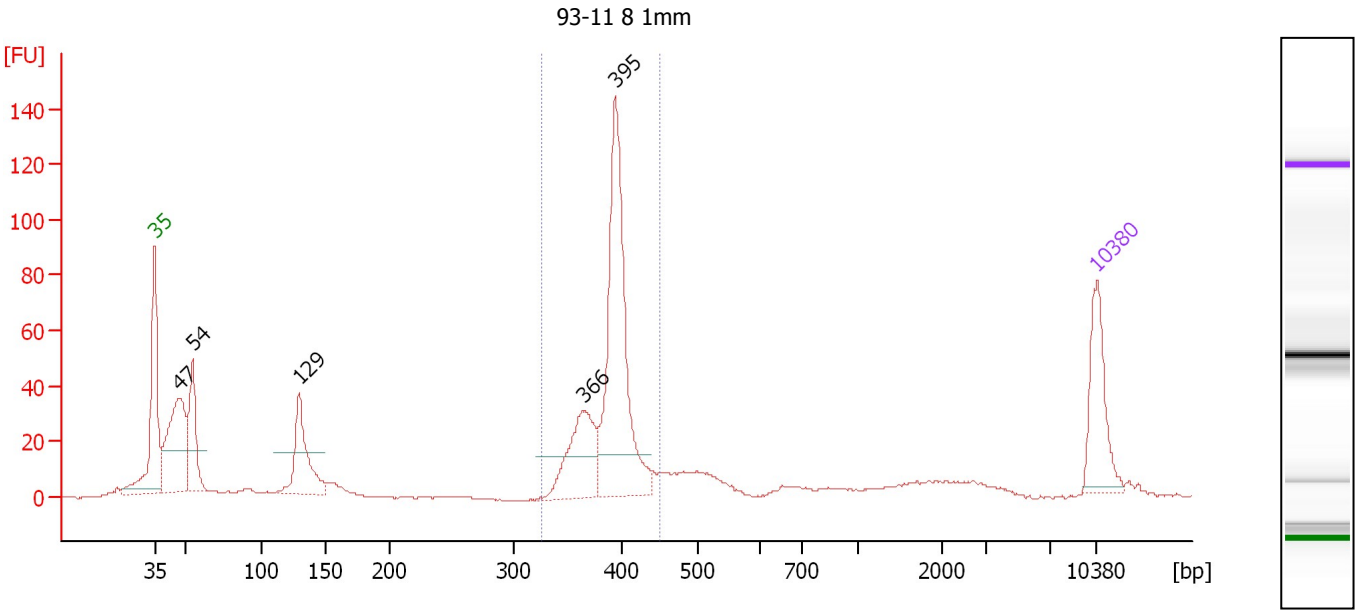
**Region table for sample 8 : IR50 8 Endo**

| From [bp] | To [bp] | Average Size [bp] | Molarity [pmol/l] | Conc. [pg/µl] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|---|---|---|---|---|---|---|---|---|
| 314 | 439 | 389 | 1,899.9 | 487.21 | 497.4 | 47 | 4.6 | |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

**Electropherogram Summary Continued ...**




**Setpoint Deviations for sample 9 : 93-11 8 1mm**

Height Threshold [FU] : 15

**Overall Results for sample 9 : 93-11 8 1mm**

| | | | |
|---|---|---|---|
| Number of peaks found: | 5 | Corr. Area 1: | 379.9 |
| Noise: | 0.2 | | |

**Peak table for sample 9 : 93-11 8 1mm**

| Peak | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations |
|---|---|---|---|---|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 47 | 216.88 | 7,009.6 | |
| 3 | 54 | 111.14 | 3,100.2 | |
| 4 | 129 | 124.09 | 1,456.1 | |
| 5 | 366 | 120.99 | 500.8 | |
| 6 | 395 | 333.46 | 1,279.7 | |
| 7 | 10,380 | 75.00 | 10.9 | Upper Marker |

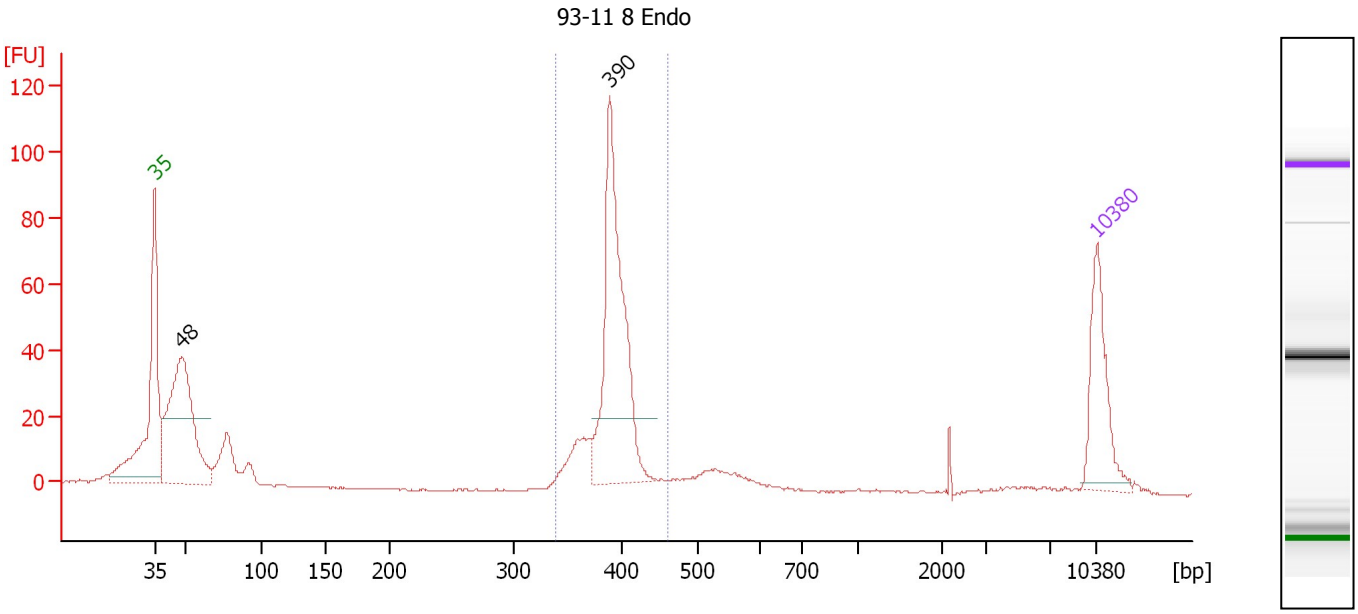
**Region table for sample 9 : 93-11 8 1mm**

| From [bp] | To [bp] | Average Size [bp] | Molarity [pmol/l] | Conc. [pg/µl] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|---|---|---|---|---|---|---|---|---|
| 327 | 451 | 390 | 1,809.8 | 465.29 | 379.9 | 45 | 5.3 | |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

**Electropherogram Summary Continued ...**




**Setpoint Deviations for sample 10 : 93-11 8 Endo**

Height Threshold [FU] : 20

**Overall Results for sample 10 : 93-11 8 Endo**

| | | | |
|---|---|---|---|
| Number of peaks found: | 2 | Corr. Area 1: | 291.1 |
| Noise: | 0.3 | | |

**Peak table for sample 10 : 93-11 8 Endo**

| Peak | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations |
|---|---|---|---|---|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 48 | 321.17 | 10,172.1 | |
| 3 | 390 | 282.36 | 1,098.2 | |
| 4 | 10,380 | 75.00 | 10.9 | Upper Marker |

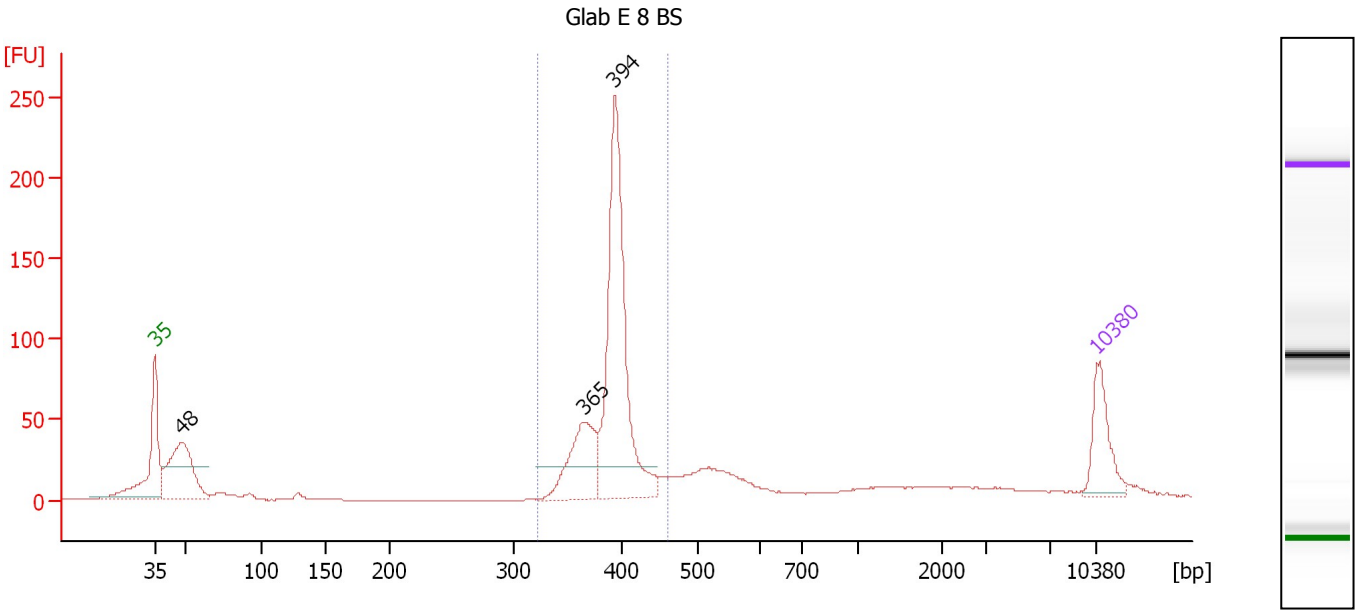
**Region table for sample 10 : 93-11 8 Endo**

| From [bp] | To [bp] | Average Size [bp] | Molarity [pmol/l] | Conc. [pg/µl] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|---|---|---|---|---|---|---|---|---|
| 339 | 462 | 392 | 1,349.6 | 348.47 | 291.1 | 48 | 4.7 | |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

**Electropherogram Summary Continued ...**




**Setpoint Deviations for sample 11 : Glab E 8 BS**

Height Threshold [FU] : 20

**Overall Results for sample 11 : Glab E 8 BS**

| | | | |
|---|---|---|---|
| Number of peaks found: | 3 | Corr. Area 1: | 647.7 |
| Noise: | 0.2 | | |

**Peak table for sample 11 : Glab E 8 BS**

| Peak | | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations |
|---|---|---|---|---|---|
| 1 | | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | | 48 | 254.18 | 8,030.1 | |
| 3 | | 365 | 159.33 | 661.5 | |
| 4 | | 394 | 495.76 | 1,906.0 | |
| 5 | | 10,380 | 75.00 | 10.9 | Upper Marker |

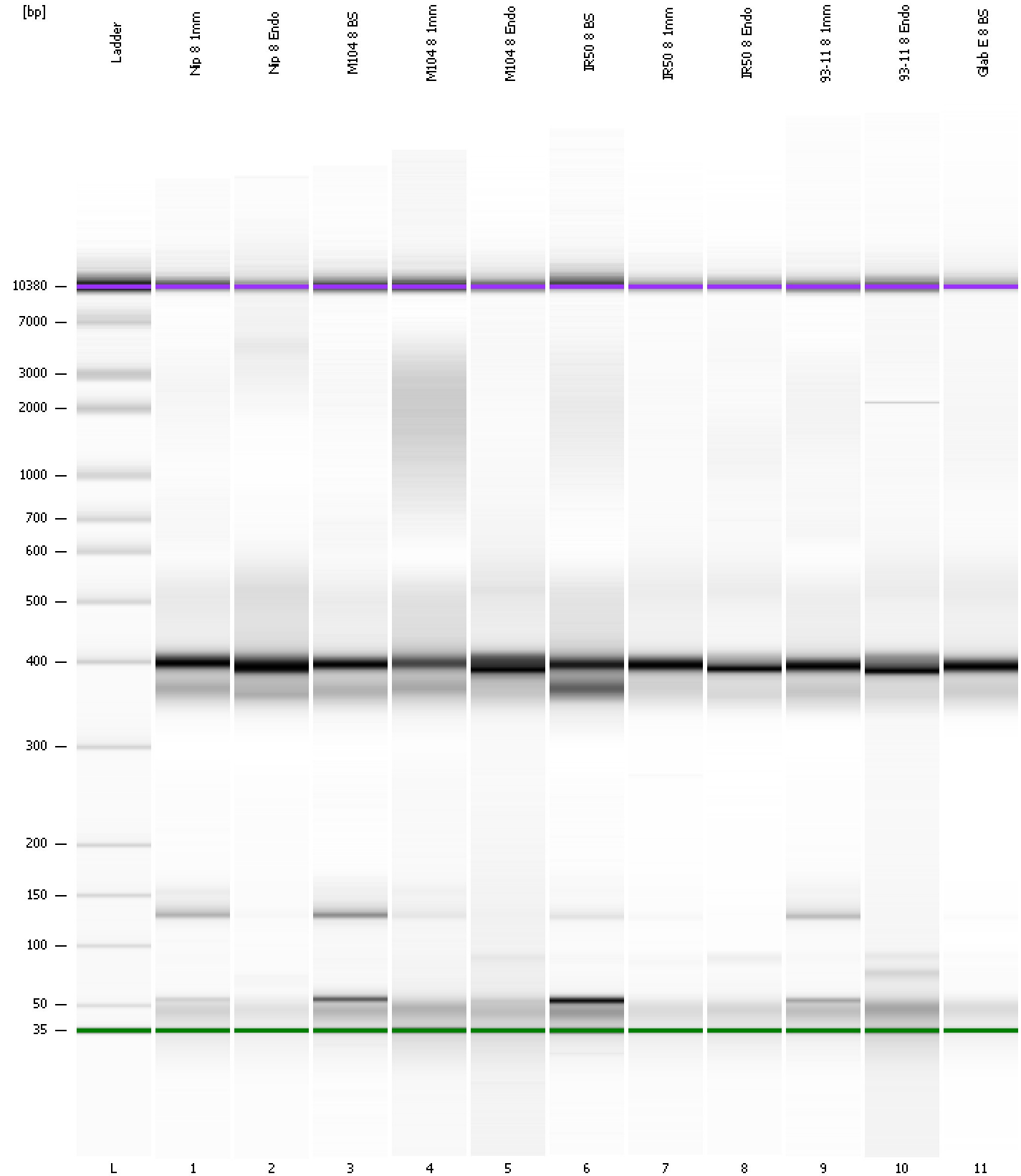
**Region table for sample 11 : Glab E 8 BS**

| From [bp] | To [bp] | Average Size [bp] | Molarity [pmol/l] | Conc. [pg/µl] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|---|---|---|---|---|---|---|---|---|
| 323 | 462 | 392 | 2,617.0 | 675.29 | 647.7 | 56 | 5.6 | |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 PM

**Gel Image**



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad

Created: 7/11/2013 1:32:58 PM
Modified: 7/11/2013 2:18:50 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 | | 7/11/2013 2:14:17 PM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2013-07-11\2013-07-11_003.xad) | | Instrument | Run | | 7/11/2013 1:33:04 PM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Product Number : G2938B | | Instrument | Run | | 7/11/2013 1:33:04 PM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Name : | | Instrument | Run | | 7/11/2013 1:33:04 PM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Vendor : Agilent Technologies | | Instrument | Run | | 7/11/2013 1:33:04 PM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Serial# : DE13701086 | | Instrument | Run | | 7/11/2013 1:33:04 PM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Firmware : C.01.069 | | Instrument | Run | | 7/11/2013 1:33:04 PM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Cartridge : Electrode | | Instrument | Run | | 7/11/2013 1:33:04 PM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |