

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

Created: 3/6/2012 1:16:50 PM
Modified: 3/6/2012 2:00:32 PM

Electrophoresis File Run Summary

Instrument Information:

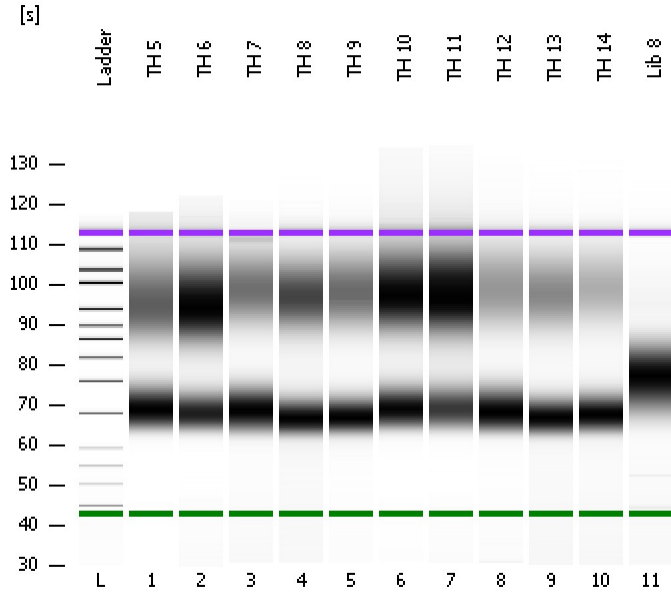
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:

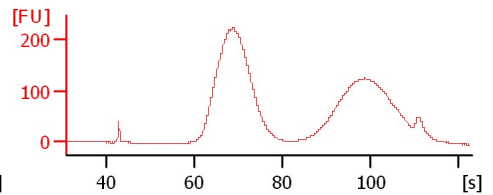
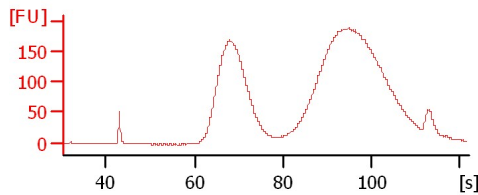
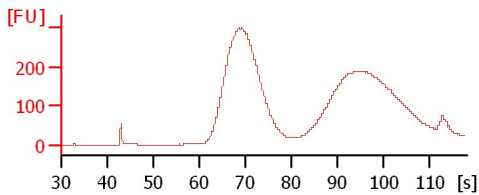
Chip Lot #: Reagent Kit Lot #: Chip Comments:



TH 5

TH 6

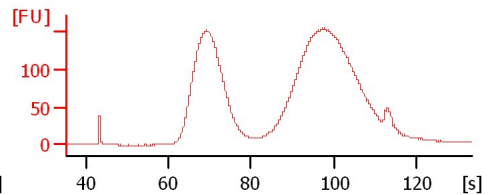
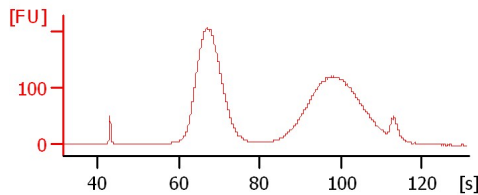
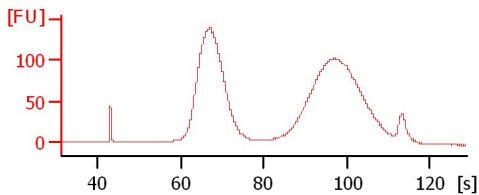
TH 7



TH 8

TH 9

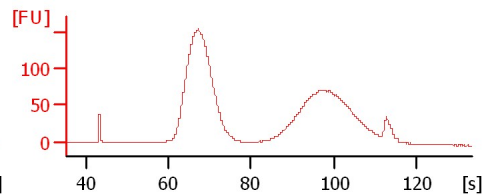
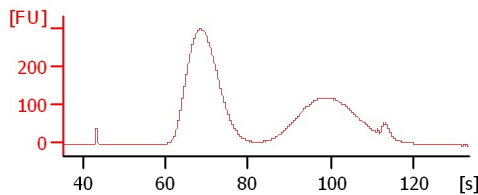
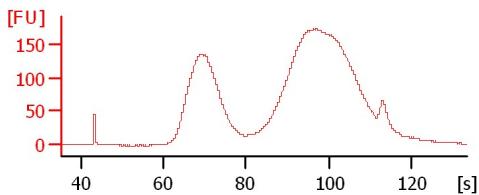
TH 10



TH 11

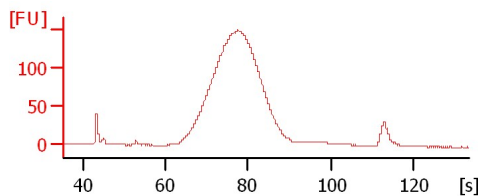
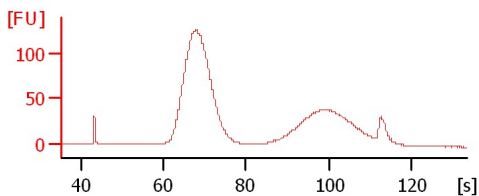
TH 12

TH 13



TH 14

Lib 8



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Electrophoresis File Run Summary (Chip Summary)

| Sample Name | Sample Comment | Rest. Digest | Status | Observation | Result Label | Result Color |
|-------------|----------------|--------------------------|--------|-------------|--------------|--------------|
| TH 5 | | <input type="checkbox"/> | ✓ | | | |
| TH 6 | | <input type="checkbox"/> | ✓ | | | |
| TH 7 | | <input type="checkbox"/> | ✓ | | | |
| TH 8 | | <input type="checkbox"/> | ✓ | | | |
| TH 9 | | <input type="checkbox"/> | ✓ | | | |
| TH 10 | | <input type="checkbox"/> | ✓ | | | |
| TH 11 | | <input type="checkbox"/> | ✓ | | | |
| TH 12 | | <input type="checkbox"/> | ✓ | | | |
| TH 13 | | <input type="checkbox"/> | ✓ | | | |
| TH 14 | | <input type="checkbox"/> | ✓ | | | |
| Lib 8 | | <input type="checkbox"/> | ✓ | | | |
| Ladder | | <input type="checkbox"/> | ✓ | | | |

Chip Lot #

Reagent Kit Lot #

Chip Comments :

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Created: 3/6/2012 1:16:50 PM
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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

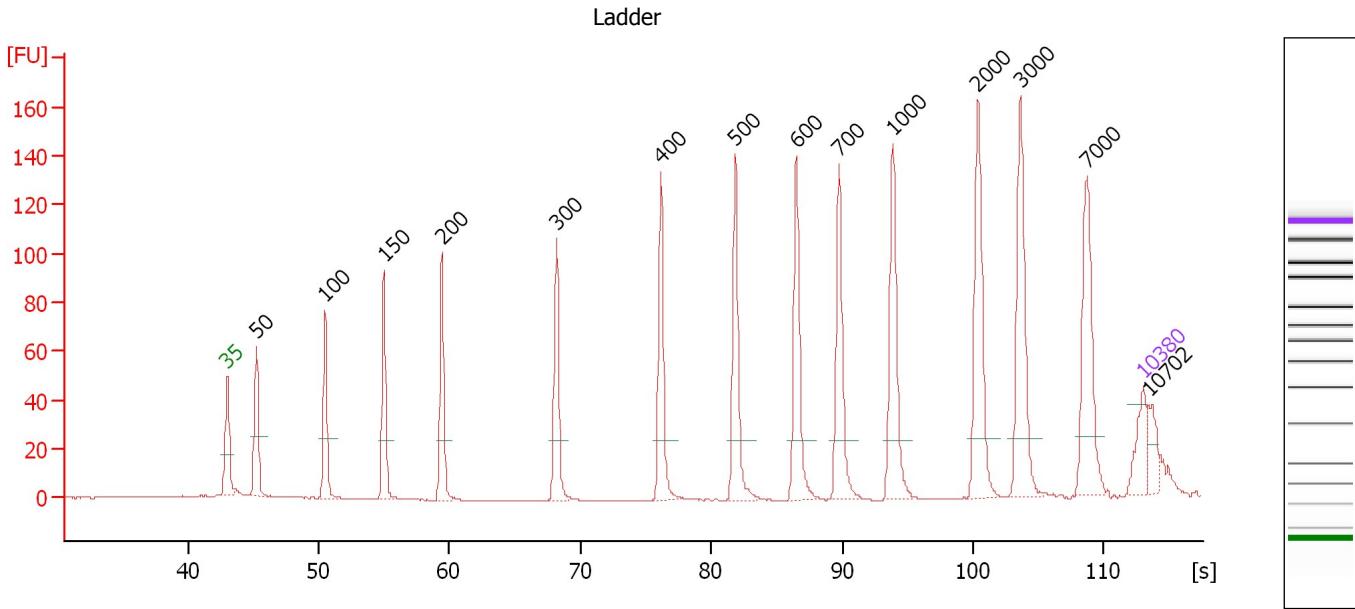
Ladder

| 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 |

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Created: 3/6/2012 1:16:50 PM
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Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

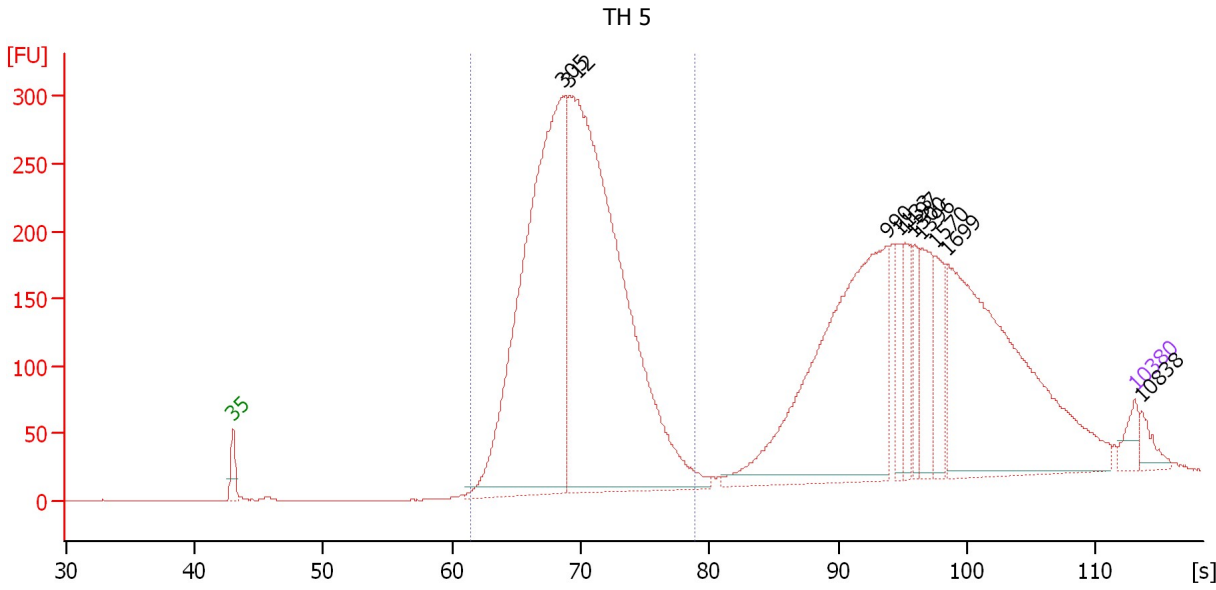
Peak table for Ladder

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations |
|------|-----------|---------------|-------------------|--------------|
| 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 |
| 16 | 10,702 | 0.00 | 0.0 | |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

Created: 3/6/2012 1:16:50 PM
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Electropherogram Summary Continued ...



Overall Results for sample 1 : TH 5

Number of peaks found: 10 Corr. Area 1: 3,736.3
 Noise: 0.2

Peak table for sample 1 : TH 5

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations |
|------|-----------|---------------|-------------------|--------------|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 305 | 3,678.55 | 18,268.6 | |
| 3 | 312 | 4,719.38 | 22,926.0 | |
| 4 | 990 | 1,949.15 | 2,984.0 | |
| 5 | 1,133 | 163.06 | 218.1 | |
| 6 | 1,197 | 197.82 | 250.4 | |
| 7 | 1,300 | 167.45 | 195.2 | |
| 8 | 1,396 | 299.16 | 324.6 | |
| 9 | 1,570 | 235.16 | 227.0 | |
| 10 | 1,699 | 1,656.22 | 1,477.4 | |
| 11 | 10,380 | 75.00 | 10.9 | Upper Marker |
| 12 | 10,838 | 0.00 | 0.0 | |

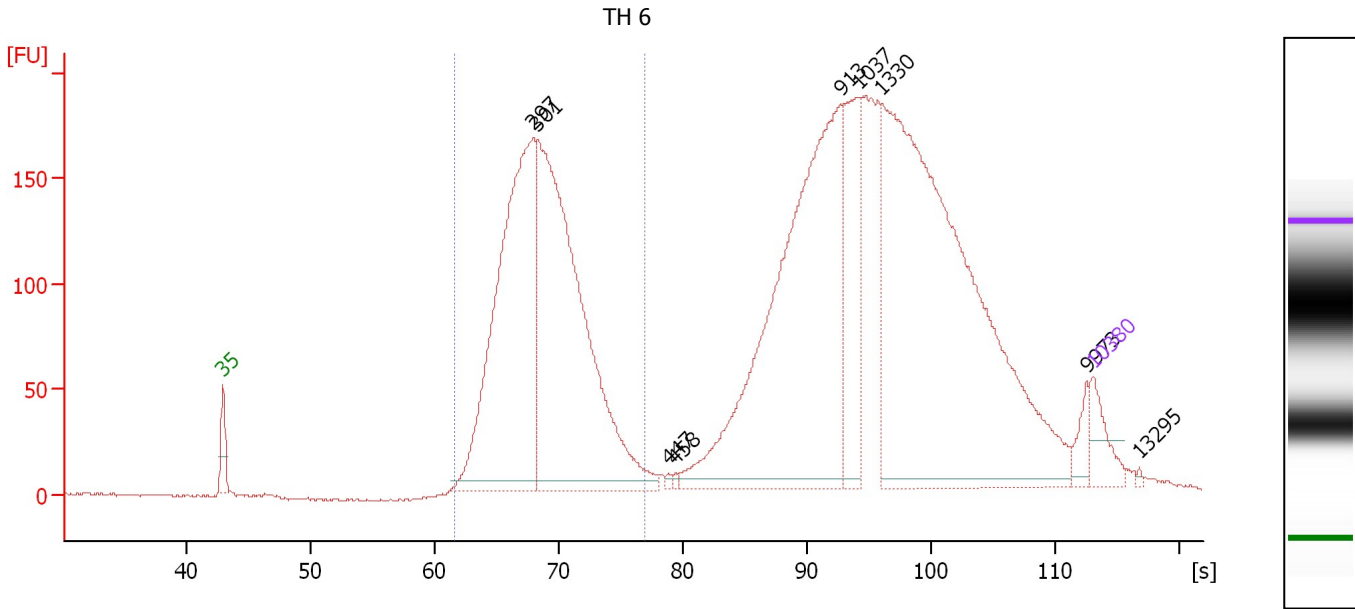
Region table for sample 1 : TH 5

| From [s] | Molarity [pmol/l] | Average Size [bp] | Conc. [pg/μl] | To [s] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|----------|-------------------|-------------------|---------------|--------|------------|------------|-----------------------------|-------|
| 61.44 | 38,302.1 | 319 | 7,914.15 | 78.85 | 3,736.3 | 54 | 12.4 | Blue |

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Electropherogram Summary Continued ...



Overall Results for sample 2 : TH 6

Number of peaks found: 9 Corr. Area 1: 1,948.4
 Noise: 0.6

Peak table for sample 2 : TH 6

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations |
|------|-----------|---------------|-------------------|--------------|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 297 | 1,428.70 | 7,287.9 | |
| 3 | 301 | 1,677.13 | 8,440.5 | |
| 4 | 447 | 7.38 | 25.0 | |
| 5 | 458 | 7.77 | 25.7 | |
| 6 | 913 | 1,458.67 | 2,420.3 | |
| 7 | 1,037 | 353.84 | 517.1 | |
| 8 | 1,330 | 1,794.07 | 2,043.9 | |
| 9 | 9,973 | 42.48 | 6.5 | |
| 10 | 10,380 | 75.00 | 10.9 | Upper Marker |
| 11 | 13,295 | 0.00 | 0.0 | |

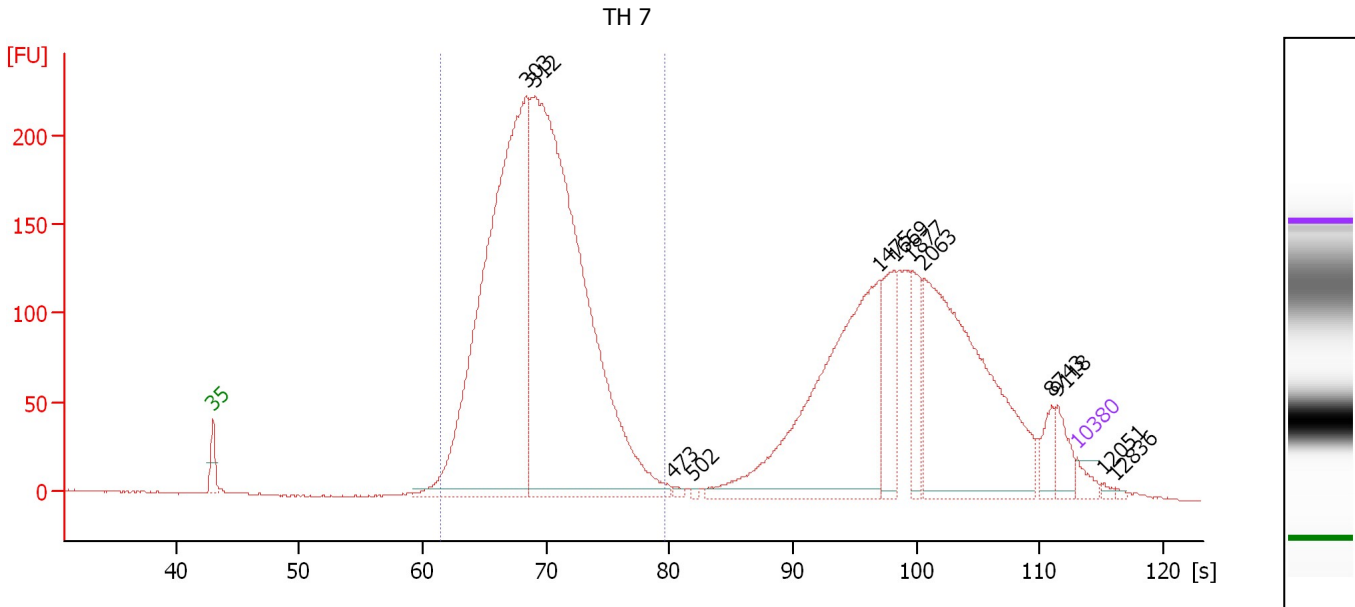
Region table for sample 2 : TH 6

| From [s] | Molarity [pmol/l] | Average Size [bp] | Conc. [pg/μl] | To [s] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|----------|-------------------|-------------------|---------------|--------|------------|------------|-----------------------------|-------|
| 61.67 | 15,240.1 | 308 | 3,048.81 | 77.02 | 1,948.4 | 36 | 11.6 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

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Electropherogram Summary Continued ...



Overall Results for sample 3 : TH 7

Number of peaks found: 12 Corr. Area 1: 3,050.8
 Noise: 0.5

Peak table for sample 3 : TH 7

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations |
|------|-----------|---------------|-------------------|--------------|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 303 | 6,401.56 | 32,007.6 | |
| 3 | 312 | 7,812.56 | 37,967.6 | |
| 4 | 473 | 32.66 | 104.7 | |
| 5 | 502 | 17.49 | 52.8 | |
| 6 | 1,475 | 2,486.30 | 2,554.6 | |
| 7 | 1,669 | 539.78 | 490.0 | |
| 8 | 1,877 | 381.10 | 307.6 | |
| 9 | 2,063 | 2,271.74 | 1,668.8 | |
| 10 | 8,743 | 147.82 | 25.6 | |
| 11 | 9,118 | 181.18 | 30.1 | |
| 12 | 10,380 | 75.00 | 10.9 | Upper Marker |
| 13 | 12,051 | 0.00 | 0.0 | |
| 14 | 12,836 | 0.00 | 0.0 | |

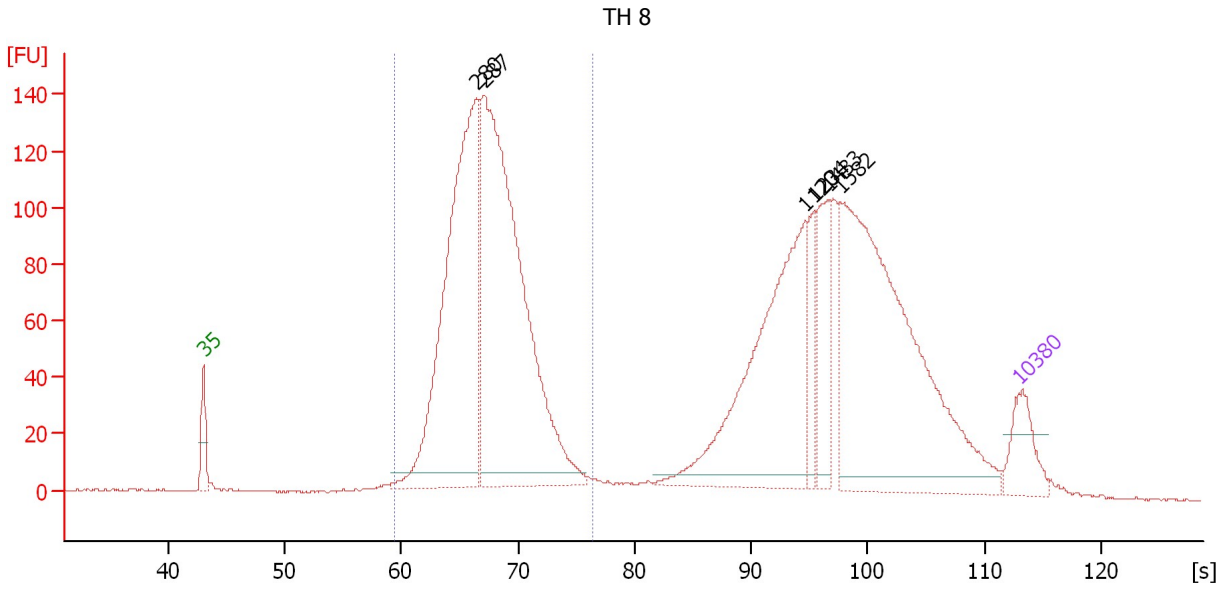
Region table for sample 3 : TH 7

| From [s] | Molarity [pmol/l] | Average Size [bp] | Conc. [pg/μl] | To [s] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|----------|-------------------|-------------------|---------------|--------|------------|------------|-----------------------------|-------|
| 61.49 | 67,407.4 | 315 | 13,728.41 | 79.65 | 3,050.8 | 59 | 13.4 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

Created: 3/6/2012 1:16:50 PM
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Electropherogram Summary Continued ...



Overall Results for sample 4 : TH 8

Number of peaks found: 6 Corr. Area 1: 1,597.1
 Noise: 0.3

Peak table for sample 4 : TH 8

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations |
|------|-----------|---------------|-------------------|--------------|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 280 | 1,089.09 | 5,899.8 | |
| 3 | 287 | 1,347.30 | 7,111.8 | |
| 4 | 1,120 | 580.85 | 785.6 | |
| 5 | 1,234 | 80.33 | 98.6 | |
| 6 | 1,433 | 134.35 | 142.0 | |
| 7 | 1,582 | 828.49 | 793.3 | |
| 8 | 10,380 | 75.00 | 10.9 | Upper Marker |

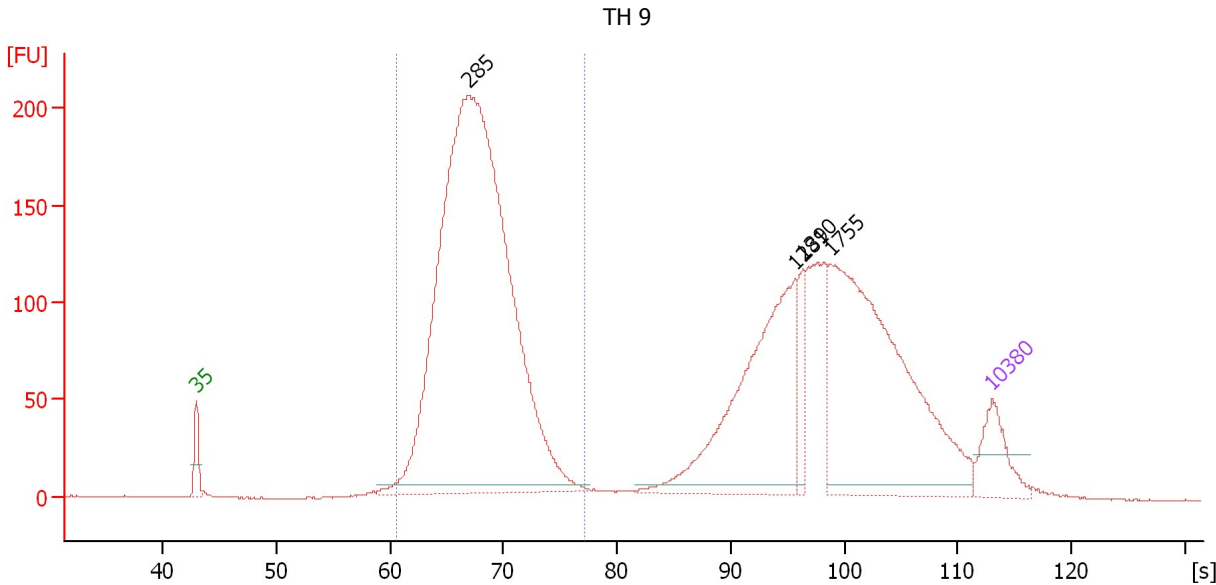
Region table for sample 4 : TH 8

| From [s] | Molarity [pmol/l] | Average Size [bp] | Conc. [pg/μl] | To [s] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|----------|-------------------|-------------------|---------------|--------|------------|------------|-----------------------------|-------|
| 59.41 | 13,344.3 | 292 | 2,530.78 | 76.38 | 1,597.1 | 48 | 12.1 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

Created: 3/6/2012 1:16:50 PM
 Modified: 3/6/2012 2:00:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : TH 9

Number of peaks found: 4 Corr. Area 1: 2,304.1
 Noise: 0.2

Peak table for sample 5 : TH 9

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations |
|------|-----------|---------------|-------------------|--------------|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 285 | 2,388.44 | 12,701.4 | |
| 3 | 1,281 | 469.17 | 554.9 | |
| 4 | 1,390 | 61.69 | 67.2 | |
| 5 | 1,755 | 650.10 | 561.3 | |
| 6 | 10,380 | 75.00 | 10.9 | Upper Marker |

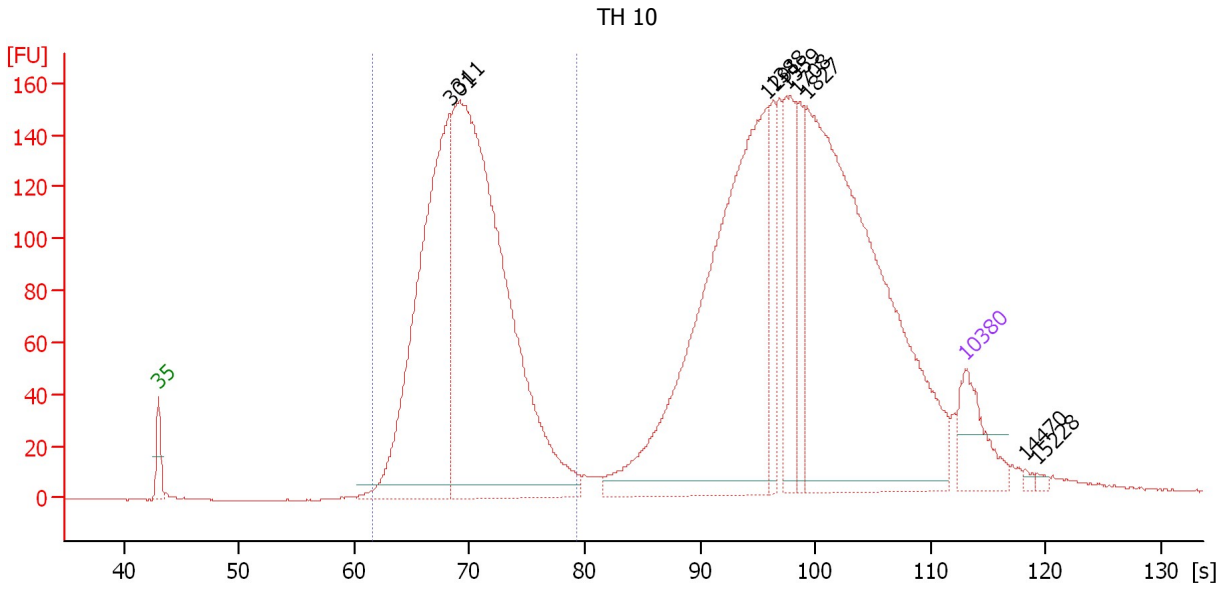
Region table for sample 5 : TH 9

| From [s] | Molarity [pmol/l] | Average Size [bp] | Conc. [pg/μl] | To [s] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|----------|-------------------|-------------------|---------------|--------|------------|------------|-----------------------------|-------|
| 60.54 | 12,402.2 | 296 | 2,386.62 | 77.16 | 2,304.1 | 53 | 11.8 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

Created: 3/6/2012 1:16:50 PM
 Modified: 3/6/2012 2:00:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : TH 10

Number of peaks found: 9 Corr. Area 1: 1,859.8
 Noise: 0.3

Peak table for sample 6 : TH 10

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations |
|------|-----------|---------------|-------------------|--------------|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 301 | 768.62 | 3,865.2 | |
| 3 | 311 | 1,296.02 | 6,314.1 | |
| 4 | 1,291 | 769.83 | 903.5 | |
| 5 | 1,388 | 75.20 | 82.1 | |
| 6 | 1,559 | 141.35 | 137.4 | |
| 7 | 1,708 | 70.50 | 62.5 | |
| 8 | 1,827 | 822.11 | 681.7 | |
| 9 | 10,380 | 75.00 | 10.9 | Upper Marker |
| 10 | 14,470 | 0.00 | 0.0 | |
| 11 | 15,228 | 0.00 | 0.0 | |

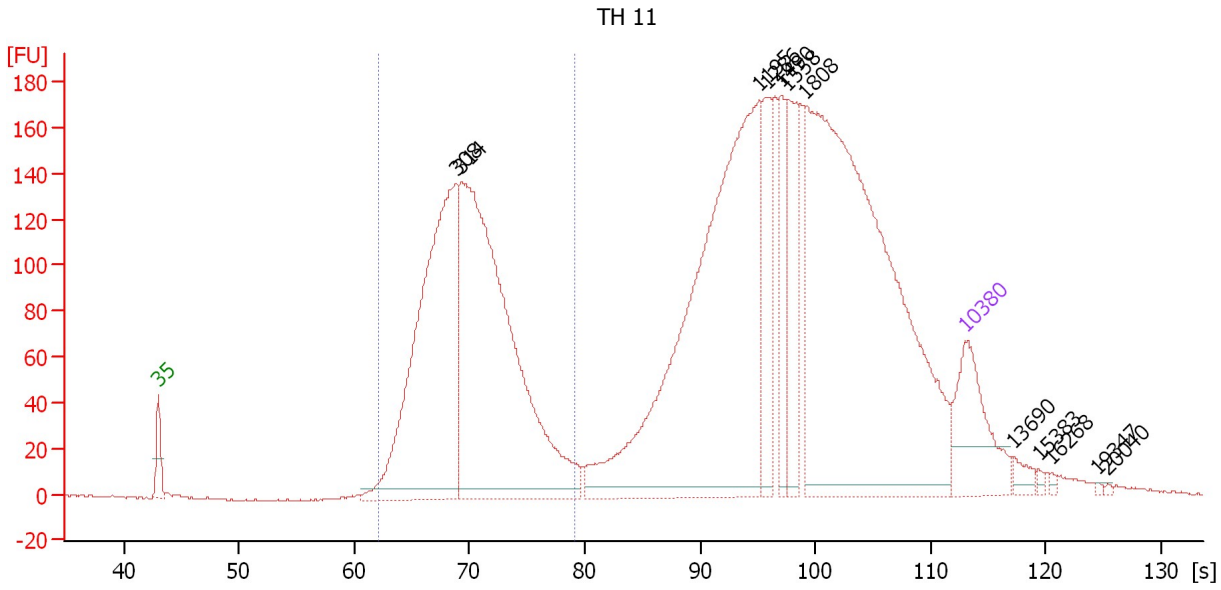
Region table for sample 6 : TH 10

| From [s] | Molarity [pmol/l] | Average Size [bp] | Conc. [pg/μl] | To [s] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|----------|-------------------|-------------------|---------------|--------|------------|------------|-----------------------------|-------|
| 61.61 | 9,477.7 | 321 | 1,966.68 | 79.29 | 1,859.8 | 41 | 12.8 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

Created: 3/6/2012 1:16:50 PM
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Electropherogram Summary Continued ...



Overall Results for sample 7 : TH 11

Number of peaks found: 12 Corr. Area 1: 1,739.5
 Noise: 0.5

Peak table for sample 7 : TH 11

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations |
|------|-----------|---------------|-------------------|--------------|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 308 | 501.53 | 2,470.0 | |
| 3 | 314 | 660.95 | 3,191.2 | |
| 4 | 1,195 | 575.51 | 729.6 | |
| 5 | 1,286 | 73.61 | 86.7 | |
| 6 | 1,490 | 55.19 | 56.1 | |
| 7 | 1,558 | 88.49 | 86.0 | |
| 8 | 1,808 | 622.40 | 521.6 | |
| 9 | 10,380 | 75.00 | 10.9 | Upper Marker |
| 10 | 13,690 | 0.00 | 0.0 | |
| 11 | 15,383 | 0.00 | 0.0 | |
| 12 | 16,268 | 0.00 | 0.0 | |
| 13 | 19,347 | 0.00 | 0.0 | |
| 14 | 20,040 | 0.00 | 0.0 | |

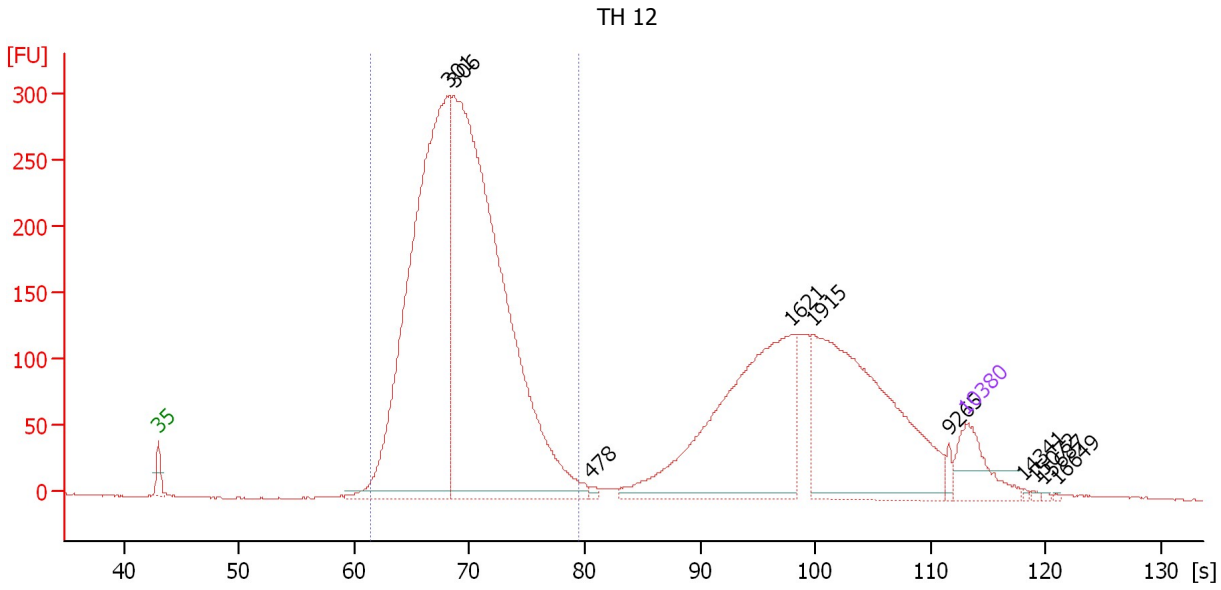
Region table for sample 7 : TH 11

| From [s] | Molarity [pmol/l] | Average Size [bp] | Conc. [pg/μl] | To [s] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|----------|-------------------|-------------------|---------------|--------|------------|------------|-----------------------------|-------|
| 62.04 | 5,240.6 | 323 | 1,092.56 | 79.11 | 1,739.5 | 34 | 13.3 | Blue |

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 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

Created: 3/6/2012 1:16:50 PM
 Modified: 3/6/2012 2:00:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : TH 12

Number of peaks found: 10 Corr. Area 1: 3,936.9
 Noise: 0.4

Peak table for sample 8 : TH 12

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations |
|------|-----------|---------------|-------------------|--------------|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 301 | 1,353.53 | 6,821.2 | |
| 3 | 306 | 1,745.52 | 8,654.9 | |
| 4 | 478 | 6.56 | 20.8 | |
| 5 | 1,621 | 515.81 | 482.3 | |
| 6 | 1,915 | 495.93 | 392.3 | |
| 7 | 9,265 | 9.84 | 1.6 | |
| 8 | 10,380 | 75.00 | 10.9 | Upper Marker |
| 9 | 14,341 | 0.00 | 0.0 | |
| 10 | 15,072 | 0.00 | 0.0 | |
| 11 | 15,687 | 0.00 | 0.0 | |
| 12 | 16,649 | 0.00 | 0.0 | |

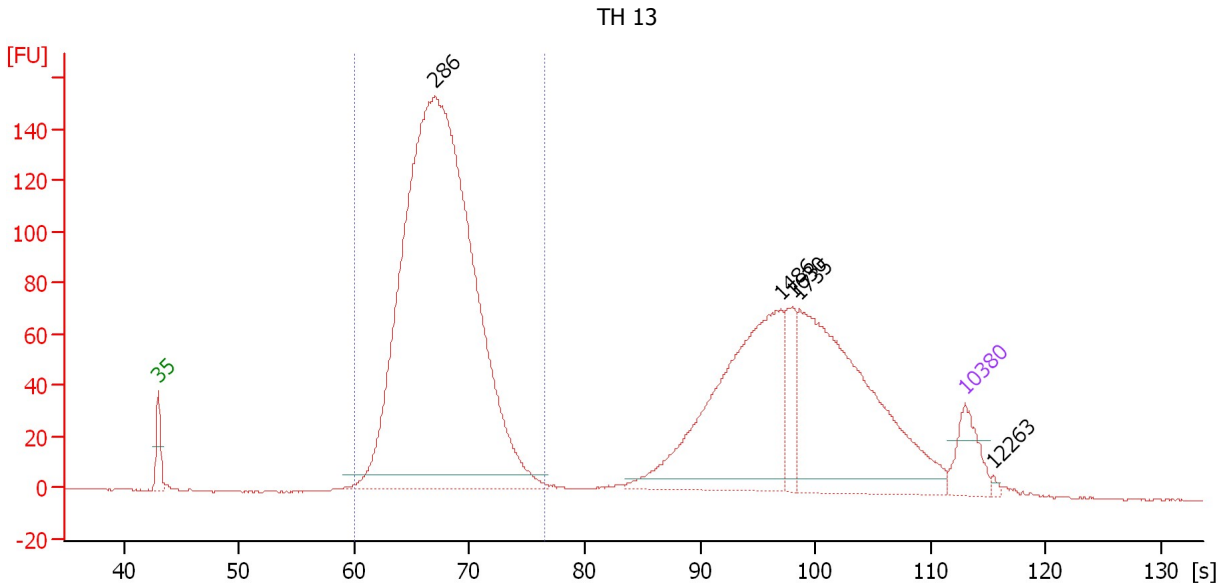
Region table for sample 8 : TH 12

| From [s] | Molarity [pmol/l] | Average Size [bp] | Conc. [pg/μl] | To [s] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|----------|-------------------|-------------------|---------------|--------|------------|------------|-----------------------------|-------|
| 61.34 | 14,748.2 | 314 | 2,992.78 | 79.47 | 3,936.9 | 64 | 13.6 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

Created: 3/6/2012 1:16:50 PM
 Modified: 3/6/2012 2:00:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : TH 13

Number of peaks found: 5 Corr. Area 1: 1,675.8
 Noise: 0.3

Peak table for sample 9 : TH 13

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations |
|------|-----------|---------------|-------------------|--------------|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 286 | 2,875.28 | 15,209.0 | |
| 3 | 1,486 | 584.31 | 595.7 | |
| 4 | 1,630 | 91.71 | 85.3 | |
| 5 | 1,735 | 617.99 | 539.5 | |
| 6 | 10,380 | 75.00 | 10.9 | Upper Marker |
| 7 | 12,263 | 0.00 | 0.0 | |

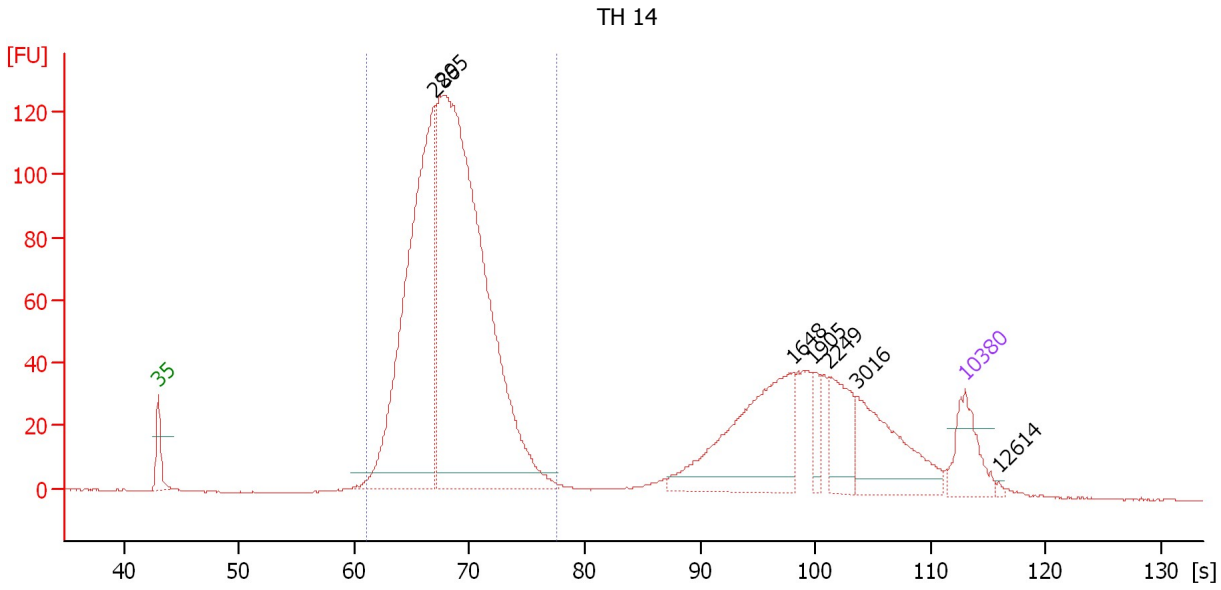
Region table for sample 9 : TH 13

| From [s] | Molarity [pmol/l] | Average Size [bp] | Conc. [pg/μl] | To [s] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|----------|-------------------|-------------------|---------------|--------|------------|------------|-----------------------------|-------|
| 60.05 | 15,179.7 | 293 | 2,897.65 | 76.56 | 1,675.8 | 58 | 11.7 | Blue |

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 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

Created: 3/6/2012 1:16:50 PM
 Modified: 3/6/2012 2:00:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : TH 14

Number of peaks found: 7 Corr. Area 1: 1,407.2
 Noise: 0.1

Peak table for sample 10 : TH 14

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations |
|------|-----------|---------------|-------------------|--------------|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 286 | 928.83 | 4,912.8 | |
| 3 | 295 | 1,583.73 | 8,138.5 | |
| 4 | 1,648 | 289.88 | 266.6 | |
| 5 | 1,905 | 33.35 | 26.5 | |
| 6 | 2,249 | 91.96 | 61.9 | |
| 7 | 3,016 | 160.48 | 80.6 | |
| 8 | 10,380 | 75.00 | 10.9 | Upper Marker |
| 9 | 12,614 | 0.00 | 0.0 | |

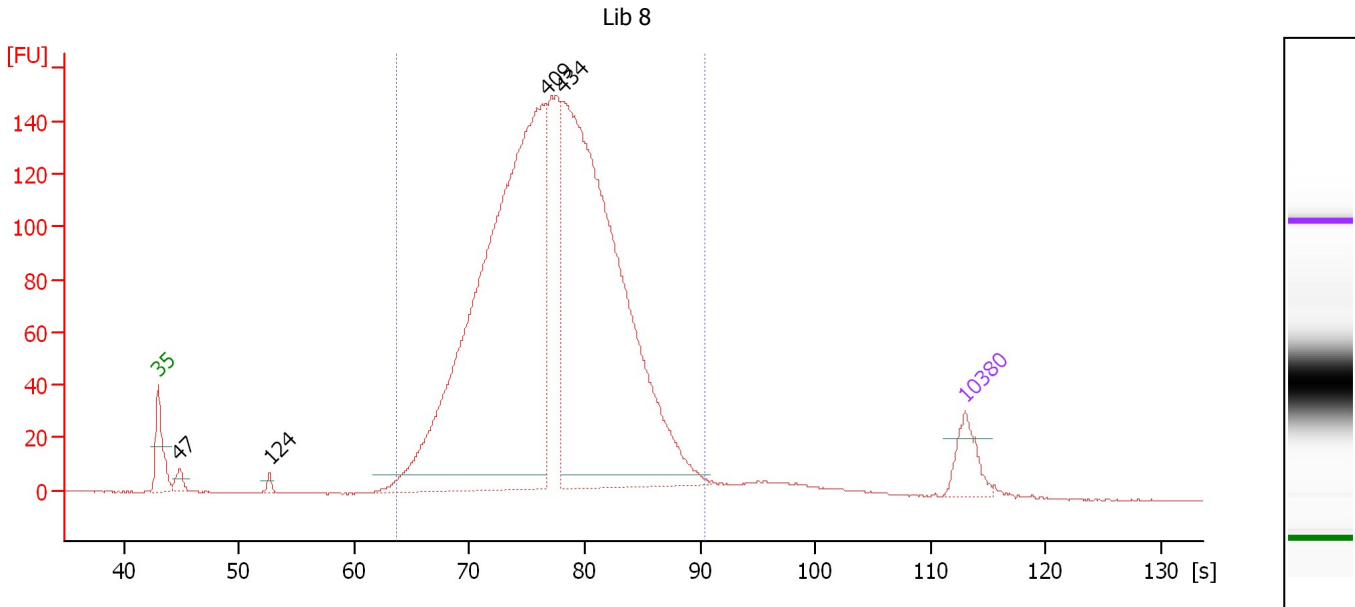
Region table for sample 10 : TH 14

| From [s] | Molarity [pmol/l] | Average Size [bp] | Conc. [pg/μl] | To [s] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|----------|-------------------|-------------------|---------------|--------|------------|------------|-----------------------------|-------|
| 61.02 | 12,831.9 | 303 | 2,523.02 | 77.56 | 1,407.2 | 69 | 11.9 | ■ |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

Created: 3/6/2012 1:16:50 PM
 Modified: 3/6/2012 2:00:32 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : Lib 8

Number of peaks found: 4 Corr. Area 1: 2,632.9
 Noise: 0.2

Peak table for sample 11 : Lib 8

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations |
|------|-----------|---------------|-------------------|--------------|
| 1 | 35 | 125.00 | 5,411.3 | Lower Marker |
| 2 | 47 | 38.89 | 1,250.3 | |
| 3 | 124 | 16.94 | 206.8 | |
| 4 | 409 | 2,302.12 | 8,535.6 | |
| 5 | 434 | 2,033.23 | 7,099.7 | |
| 6 | 10,380 | 75.00 | 10.9 | Upper Marker |

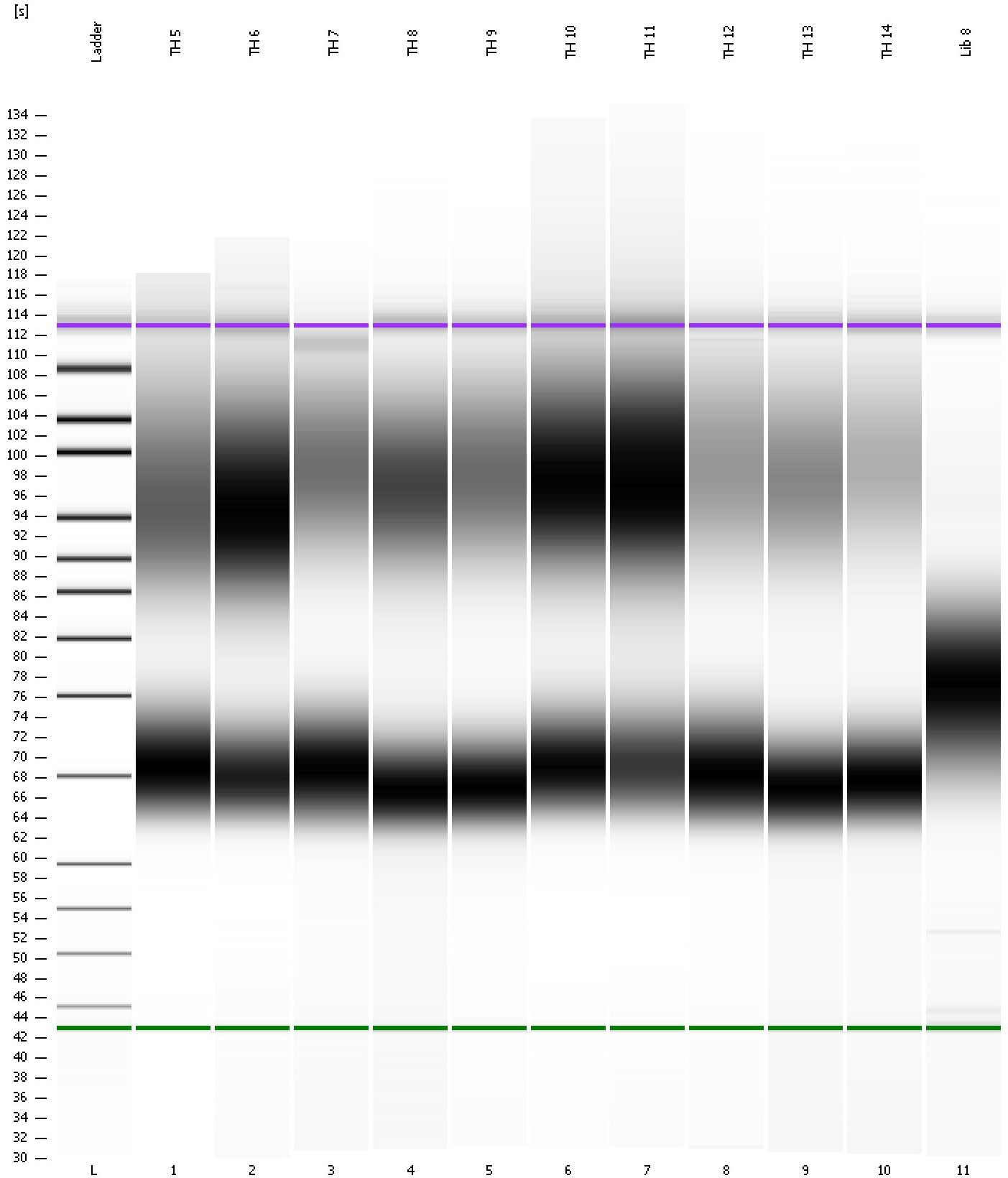
Region table for sample 11 : Lib 8

| From [s] | Molarity [pmol/l] | Average Size [bp] | Conc. [pg/μl] | To [s] | Corr. Area | % of Total | Size distribution in CV [%] | Color |
|----------|-------------------|-------------------|---------------|--------|------------|------------|-----------------------------|-------|
| 63.62 | 18,816.1 | 427 | 5,023.02 | 90.37 | 2,632.9 | 96 | 19.8 | Blue |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

Created: 3/6/2012 1:16:50 PM
Modified: 3/6/2012 2:00:32 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad

Created: 3/6/2012 1:16:50 PM
 Modified: 3/6/2012 2:00:32 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 | | 3/6/2012 1:58:08 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2012-03-06\2012-03-06_005.xad) | | Instrument | Run | | 3/6/2012 1:16:55 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Product Number : G2938B | | Instrument | Run | | 3/6/2012 1:16:55 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Name : | | Instrument | Run | | 3/6/2012 1:16:55 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Vendor : Agilent Technologies | | Instrument | Run | | 3/6/2012 1:16:55 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Serial# : DE13701086 | | Instrument | Run | | 3/6/2012 1:16:55 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Firmware : C.01.069 | | Instrument | Run | | 3/6/2012 1:16:55 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Cartridge : Electrode | | Instrument | Run | | 3/6/2012 1:16:55 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |