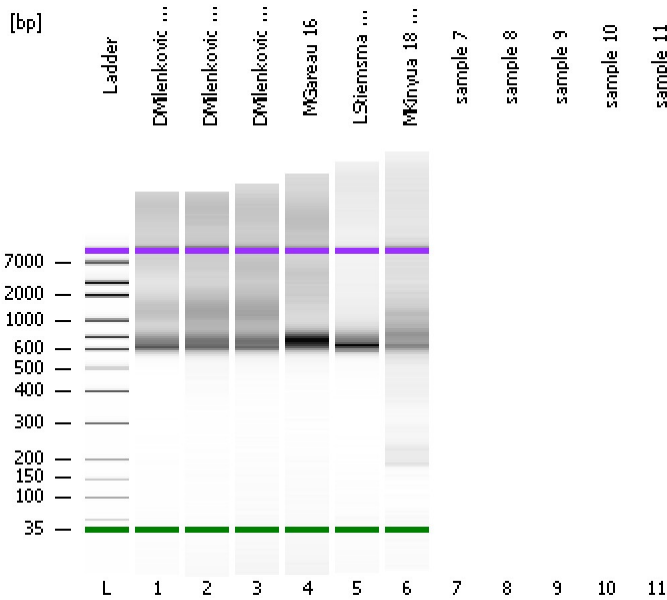


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
Modified: 10/16/2018 5:16:17 PM

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

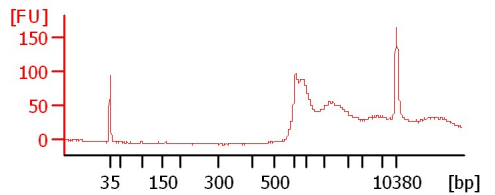
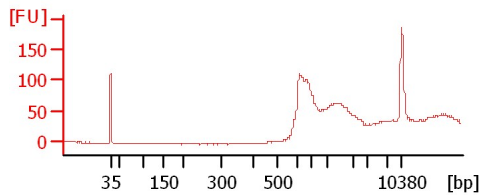
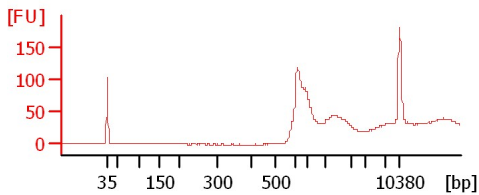
Chip Information:

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

DMilenkovic 1-48

DMilenkovic 49-80

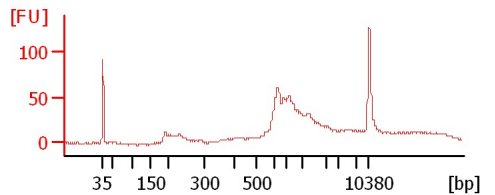
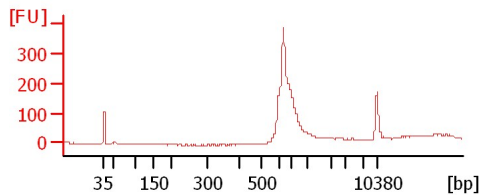
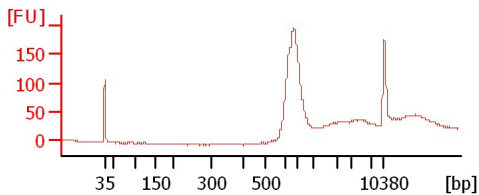
DMilenkovic 81-100



MGareau 16

LStiemsma 33 Nested

MKinyua 18 Arch 4F-534R



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
 Modified: 10/16/2018 5:16:17 PM

Electrophoresis File Run Summary (Chip Summary)

| Sample Name | Sample Comment | Rest. Digest | Status | Observation | Result Label | Result Color |
|-------------------------|----------------|--------------------------|--------|-------------|--------------|--------------|
| DMilenkovic 1-48 | | <input type="checkbox"/> | ✓ | | | |
| DMilenkovic 49-80 | | <input type="checkbox"/> | ✓ | | | |
| DMilenkovic 81-100 | | <input type="checkbox"/> | ✓ | | | |
| MGareau 16 | | <input type="checkbox"/> | ✓ | | | |
| LStiemsma 33 Nested | | <input type="checkbox"/> | ✓ | | | |
| MKinyua 18 Arch 4F-534R | | <input type="checkbox"/> | ✓ | | | |
| sample 7 | | <input type="checkbox"/> | | | | |
| sample 8 | | <input type="checkbox"/> | | | | |
| sample 9 | | <input type="checkbox"/> | | | | |
| sample 10 | | <input type="checkbox"/> | | | | |
| sample 11 | | <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\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
Modified: 10/16/2018 5:16:17 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

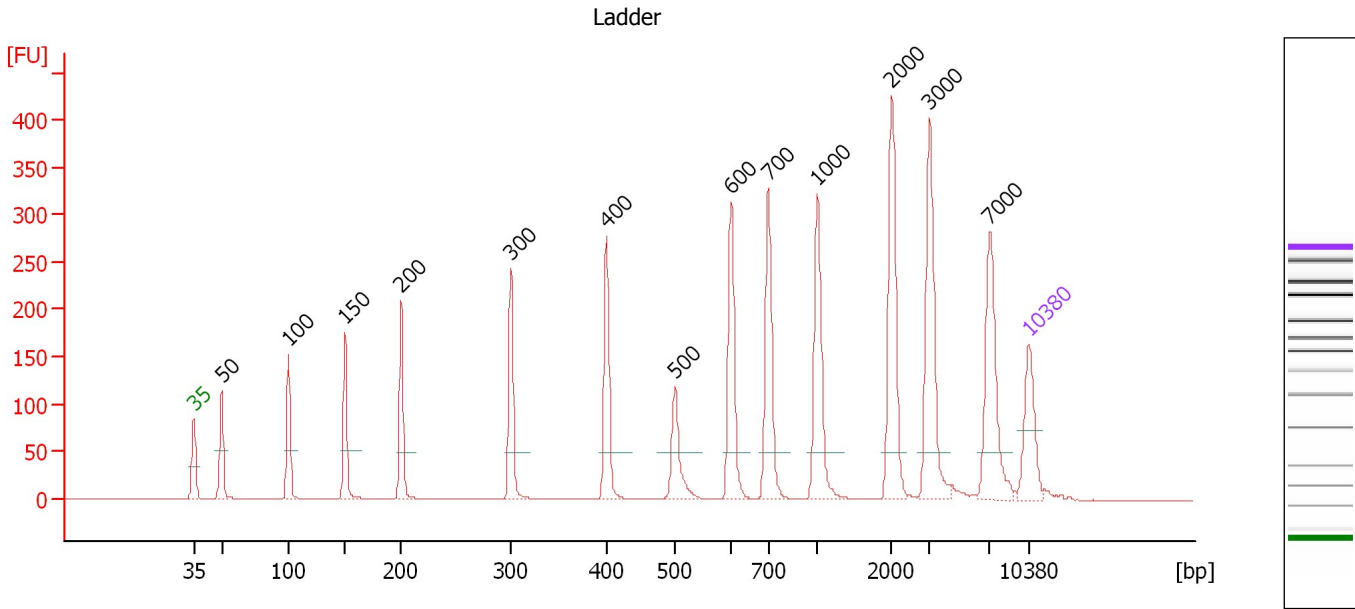
Ladder

| 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:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
 Modified: 10/16/2018 5:16:17 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

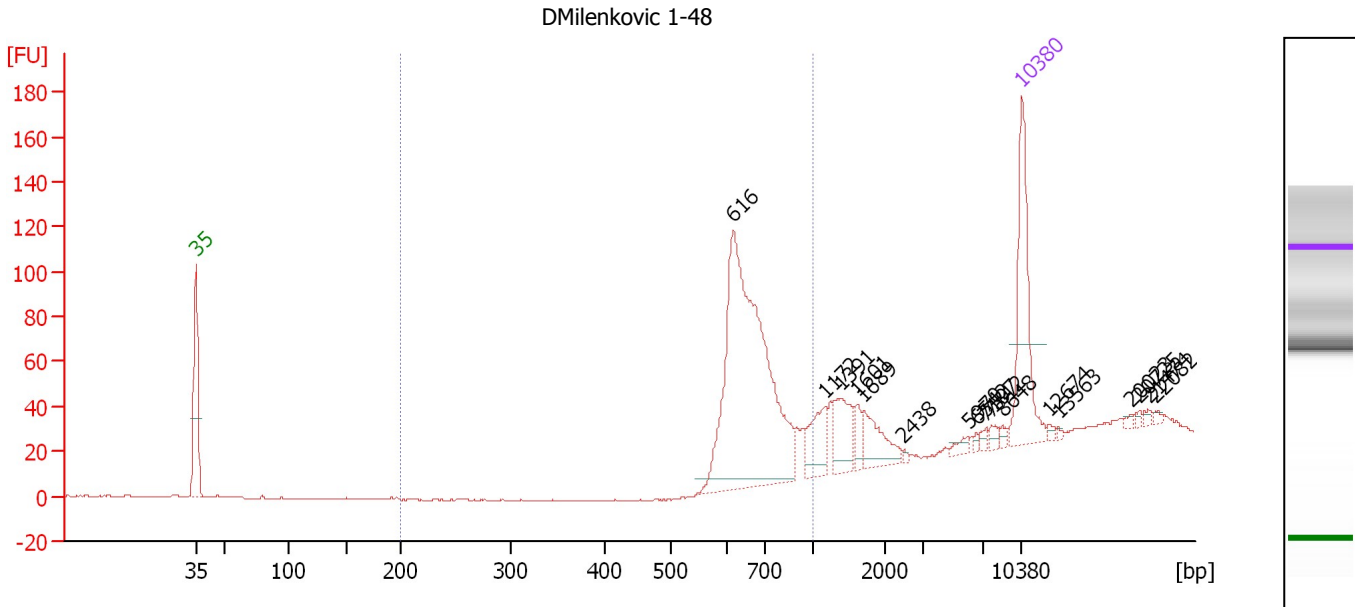
Peak table for Ladder

| 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 | 45.32 |
| 3 | 100 | 150.00 | 2,272.7 | Ladder Peak | 50.91 |
| 4 | 150 | 150.00 | 1,515.2 | Ladder Peak | 55.69 |
| 5 | 200 | 150.00 | 1,136.4 | Ladder Peak | 60.39 |
| 6 | 300 | 150.00 | 757.6 | Ladder Peak | 69.60 |
| 7 | 400 | 150.00 | 568.2 | Ladder Peak | 77.60 |
| 8 | 500 | 150.00 | 454.5 | Ladder Peak | 83.32 |
| 9 | 600 | 150.00 | 378.8 | Ladder Peak | 88.06 |
| 10 | 700 | 150.00 | 324.7 | Ladder Peak | 91.14 |
| 11 | 1,000 | 150.00 | 227.3 | Ladder Peak | 95.25 |
| 12 | 2,000 | 150.00 | 113.6 | Ladder Peak | 101.47 |
| 13 | 3,000 | 150.00 | 75.8 | Ladder Peak | 104.69 |
| 14 | 7,000 | 150.00 | 32.5 | Ladder Peak | 109.74 |
| 15 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
 Modified: 10/16/2018 5:16:17 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : DMilenkovic 1-48

Number of peaks found: 17 Corr. Area 1: 389.3
 Noise: 0.3

Peak table for sample 1 : DMilenkovic 1-48

| 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 | 616 | 332.05 | 816.7 | | 88.55 |
| 3 | 1,172 | 31.83 | 41.1 | | 96.32 |
| 4 | 1,391 | 32.06 | 34.9 | | 97.68 |
| 5 | 1,601 | 10.46 | 9.9 | | 98.99 |
| 6 | 1,689 | 29.33 | 26.3 | | 99.53 |
| 7 | 2,438 | 1.58 | 1.0 | | 102.88 |
| 8 | 5,970 | 4.99 | 1.3 | | 108.44 |
| 9 | 6,542 | 2.42 | 0.6 | | 109.16 |
| 10 | 7,197 | 3.41 | 0.7 | | 109.93 |
| 11 | 7,712 | 4.36 | 0.9 | | 110.42 |
| 12 | 8,648 | 3.25 | 0.6 | | 111.33 |
| 13 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 14 | 12,674 | 0.00 | 0.0 | | 115.21 |
| 15 | 13,563 | 0.00 | 0.0 | | 116.07 |
| 16 | 20,023 | 0.00 | 0.0 | | 122.31 |
| 17 | 20,725 | 0.00 | 0.0 | | 122.99 |
| 18 | 21,474 | 0.00 | 0.0 | | 123.71 |
| 19 | 22,082 | 0.00 | 0.0 | | 124.30 |

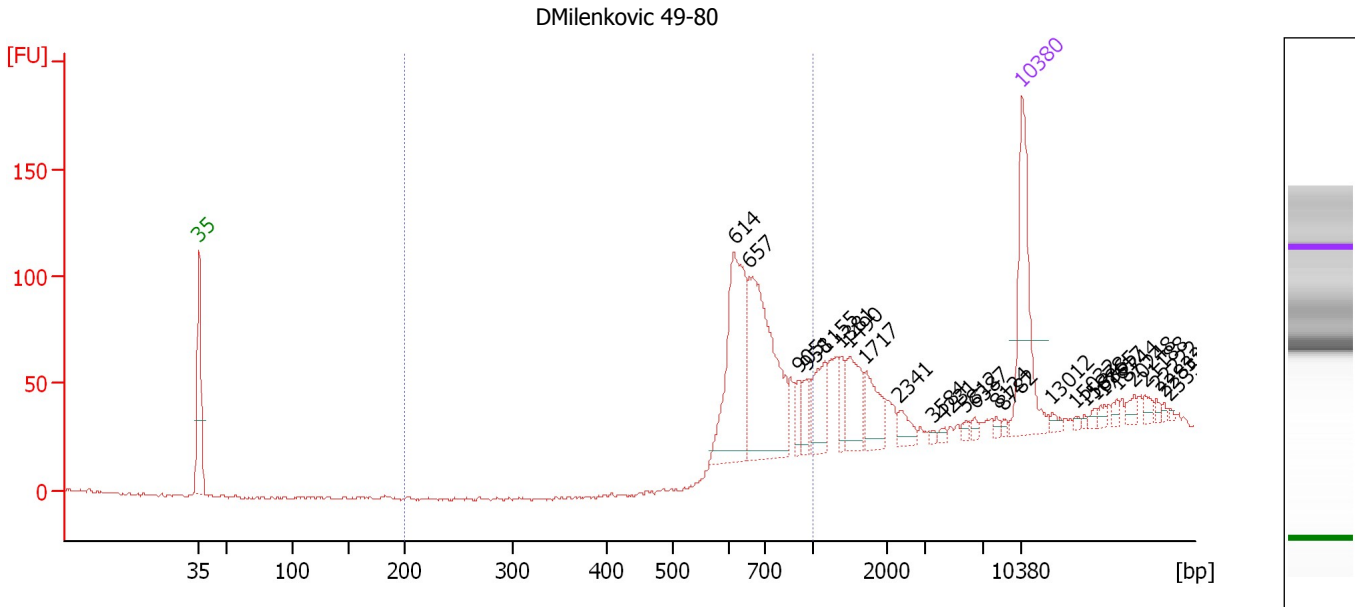
Region table for sample 1 : DMilenkovic 1-48

| 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 | 678 | 261.05 | 389.3 | 590.5 | 69 | 12.7 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
 Modified: 10/16/2018 5:16:17 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : DMilenkovic 49-80

Number of peaks found: 26 Corr. Area 1: 487.9
 Noise: 0.9

Peak table for sample 2 : DMilenkovic 49-80

| 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 | 614 | 131.37 | 324.0 | | 88.50 |
| 3 | 657 | 159.70 | 368.2 | | 89.82 |
| 4 | 905 | 13.40 | 22.4 | | 93.95 |
| 5 | 958 | 15.20 | 24.0 | | 94.67 |
| 6 | 1,155 | 32.51 | 42.7 | | 96.21 |
| 7 | 1,381 | 14.95 | 16.4 | | 97.62 |
| 8 | 1,490 | 35.56 | 36.1 | | 98.30 |
| 9 | 1,717 | 28.52 | 25.2 | | 99.71 |
| 10 | 2,341 | 11.11 | 7.2 | | 102.57 |
| 11 | 3,584 | 1.92 | 0.8 | | 105.42 |
| 12 | 4,231 | 2.41 | 0.9 | | 106.24 |
| 13 | 5,812 | 3.17 | 0.8 | | 108.24 |
| 14 | 6,387 | 2.76 | 0.7 | | 108.96 |
| 15 | 8,124 | 2.36 | 0.4 | | 110.82 |
| 16 | 8,782 | 2.10 | 0.4 | | 111.46 |
| 17 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 18 | 13,012 | 0.00 | 0.0 | | 115.54 |
| 19 | 15,032 | 0.00 | 0.0 | | 117.49 |
| 20 | 15,878 | 0.00 | 0.0 | | 118.31 |
| 21 | 16,865 | 0.00 | 0.0 | | 119.26 |
| 22 | 17,617 | 0.00 | 0.0 | | 119.99 |
| 23 | 18,744 | 0.00 | 0.0 | | 121.08 |
| 24 | 20,248 | 0.00 | 0.0 | | 122.53 |
| 25 | 21,188 | 0.00 | 0.0 | | 123.43 |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
Modified: 10/16/2018 5:16:17 PM

Electropherogram Summary Continued ...**... Peak table for sample 2 : DMilenkovic 49-80**

| Peak | Size [bp] | Conc. [pg/ μ l] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|------|-----------|---------------------|-------------------|--------------|----------------------------|
| 26 | 22,222 | 0.00 | 0.0 | | 124.43 |
| 27 | 22,833 | 0.00 | 0.0 | | 125.02 |
| 28 | 23,350 | 0.00 | 0.0 | | 125.52 |

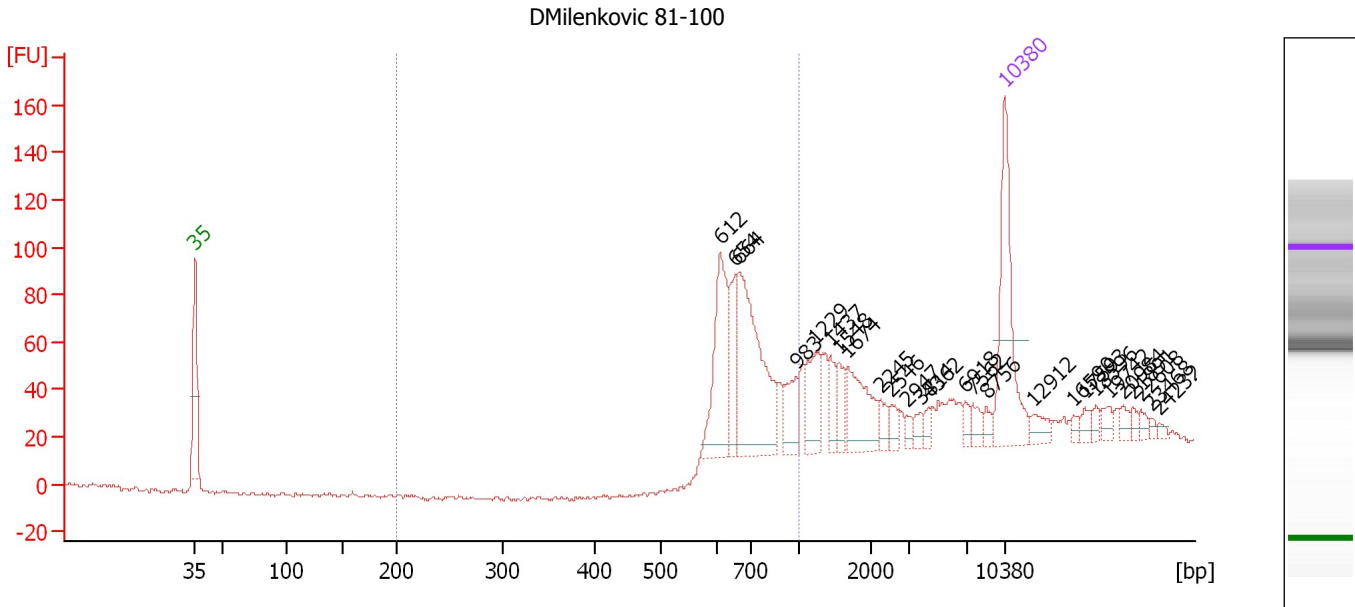
Region table for sample 2 : DMilenkovic 49-80

| From [bp] | To [bp] | Average Size [bp] | Conc. [pg/ μ l] | Corr. Area | Molarity [pmol/l] | Co % of lor Total | Size distribution in CV [%] |
|-----------|---------|-------------------|---------------------|------------|-------------------|--|-----------------------------|
| 200 | 1,000 | 708 | 308.54 | 487.9 | 673.6 |  56 | 15.3 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
 Modified: 10/16/2018 5:16:17 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : DMilenkovic 81-100

Number of peaks found: 26 Corr. Area 1: 465.0
 Noise: 0.9

Peak table for sample 3 : DMilenkovic 81-100

| 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 | 612 | 86.11 | 213.2 | | 88.42 |
| 3 | 654 | 38.73 | 89.8 | | 89.71 |
| 4 | 664 | 136.72 | 311.9 | | 90.04 |
| 5 | 983 | 25.37 | 39.1 | | 95.02 |
| 6 | 1,229 | 35.15 | 43.3 | | 96.68 |
| 7 | 1,437 | 15.61 | 16.5 | | 97.97 |
| 8 | 1,548 | 15.49 | 15.2 | | 98.66 |
| 9 | 1,674 | 43.43 | 39.3 | | 99.44 |
| 10 | 2,245 | 9.12 | 6.2 | | 102.26 |
| 11 | 2,546 | 8.28 | 4.9 | | 103.22 |
| 12 | 2,947 | 4.26 | 2.2 | | 104.52 |
| 13 | 3,814 | 6.20 | 2.5 | | 105.71 |
| 14 | 4,362 | 5.04 | 1.8 | | 106.41 |
| 15 | 6,918 | 5.89 | 1.3 | | 109.63 |
| 16 | 7,562 | 8.83 | 1.8 | | 110.28 |
| 17 | 8,756 | 5.83 | 1.0 | | 111.43 |
| 18 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 19 | 12,912 | 0.00 | 0.0 | | 115.44 |
| 20 | 16,590 | 0.00 | 0.0 | | 118.99 |
| 21 | 17,593 | 0.00 | 0.0 | | 119.96 |
| 22 | 18,596 | 0.00 | 0.0 | | 120.93 |
| 23 | 19,742 | 0.00 | 0.0 | | 122.04 |
| 24 | 20,984 | 0.00 | 0.0 | | 123.24 |
| 25 | 21,891 | 0.00 | 0.0 | | 124.11 |


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
Modified: 10/16/2018 5:16:17 PM

Electropherogram Summary Continued ...**... Peak table for sample 3 : DMilenkovic 81-100**

| Peak | Size [bp] | Conc. [pg/ μ l] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|------|-----------|---------------------|-------------------|--------------|----------------------------|
| 26 | 22,608 | 0.00 | 0.0 | | 124.81 |
| 27 | 23,468 | 0.00 | 0.0 | | 125.64 |
| 28 | 24,232 | 0.00 | 0.0 | | 126.37 |

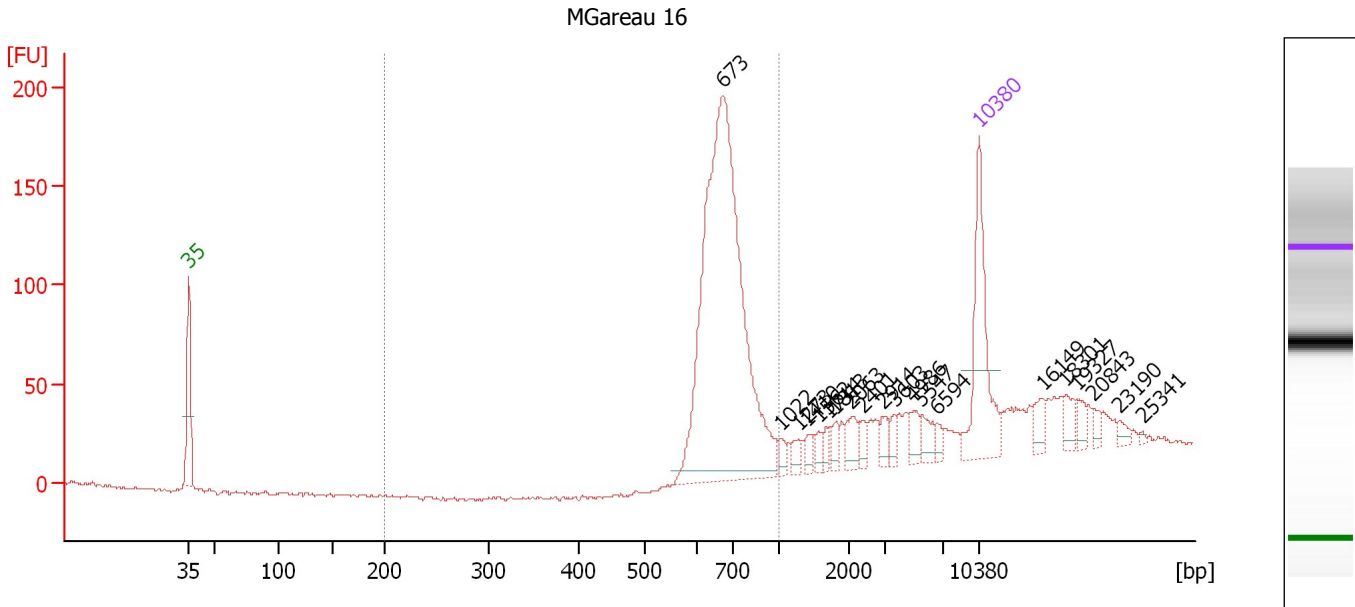
Region table for sample 3 : DMilenkovic 81-100

| From [bp] | To [bp] | Average Size [bp] | Conc. [pg/ μ l] | Corr. Area | Molarity [pmol/l] | Co % of lor Total | Size distribution in CV [%] |
|-----------|---------|-------------------|---------------------|------------|-------------------|--|-----------------------------|
| 200 | 1,000 | 712 | 301.81 | 465.0 | 656.1 |  46 | 15.4 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
 Modified: 10/16/2018 5:16:17 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : MGareau 16

Number of peaks found: 20 Corr. Area 1: 764.8
 Noise: 0.9

Peak table for sample 4 : MGareau 16


| 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 | 673 | 496.91 | 1,119.4 | | 90.30 |
| 3 | 1,022 | 7.74 | 11.5 | | 95.39 |
| 4 | 1,273 | 7.81 | 9.3 | | 96.95 |
| 5 | 1,410 | 6.77 | 7.3 | | 97.80 |
| 6 | 1,562 | 6.41 | 6.2 | | 98.75 |
| 7 | 1,714 | 6.29 | 5.6 | | 99.69 |
| 8 | 1,813 | 8.83 | 7.4 | | 100.30 |
| 9 | 2,063 | 13.90 | 10.2 | | 101.67 |
| 10 | 2,401 | 8.02 | 5.1 | | 102.76 |
| 11 | 2,914 | 9.01 | 4.7 | | 104.41 |
| 12 | 3,603 | 6.64 | 2.8 | | 105.45 |
| 13 | 4,986 | 10.59 | 3.2 | | 107.19 |
| 14 | 5,547 | 11.95 | 3.3 | | 107.90 |
| 15 | 6,594 | 5.96 | 1.4 | | 109.22 |
| 16 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 17 | 16,149 | 0.00 | 0.0 | | 118.57 |
| 18 | 18,301 | 0.00 | 0.0 | | 120.65 |
| 19 | 19,327 | 0.00 | 0.0 | | 121.64 |
| 20 | 20,843 | 0.00 | 0.0 | | 123.10 |
| 21 | 23,190 | 0.00 | 0.0 | | 125.37 |
| 22 | 25,341 | 0.00 | 0.0 | | 127.44 |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
Modified: 10/16/2018 5:16:17 PM

Electropherogram Summary Continued ...

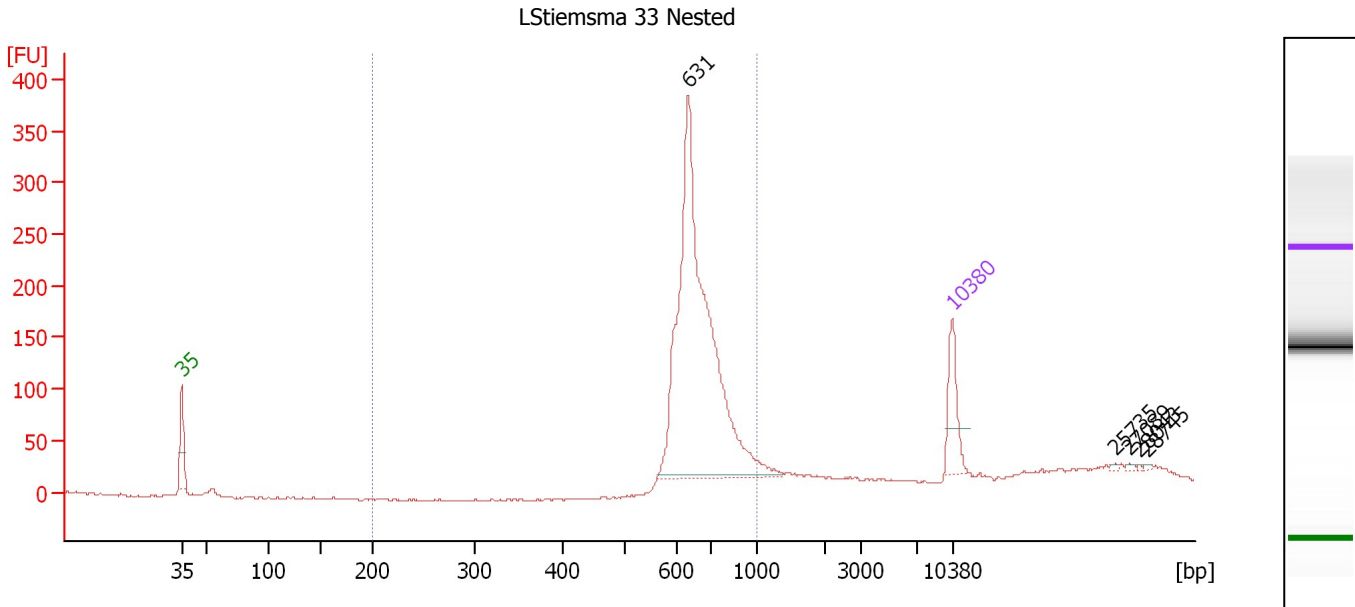
... Region table for sample 4 : MGareau 16

| From [bp] | To [bp] | Average Size [bp] | Conc. [pg/ μ l] | Corr. Area | Molarity [pmol/l] | Co % of lor Total | Size distribution in CV [%] |
|-----------|---------|-------------------|---------------------|------------|-------------------|--|-----------------------------|
| 200 | 1,000 | 690 | 429.52 | 764.8 | 952.4 |  64 | 10.9 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
 Modified: 10/16/2018 5:16:17 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : LStiemsma 33 Nested

Number of peaks found: 5 Corr. Area 1: 1,268.5
 Noise: 1.0

Peak table for sample 5 : LStiemsma 33 Nested

| 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 | 631 | 1,001.75 | 2,404.5 | | 89.02 |
| 3 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 4 | 25,735 | 0.00 | 0.0 | | 127.82 |
| 5 | 27,089 | 0.00 | 0.0 | | 129.13 |
| 6 | 28,043 | 0.00 | 0.0 | | 130.05 |
| 7 | 28,745 | 0.00 | 0.0 | | 130.73 |

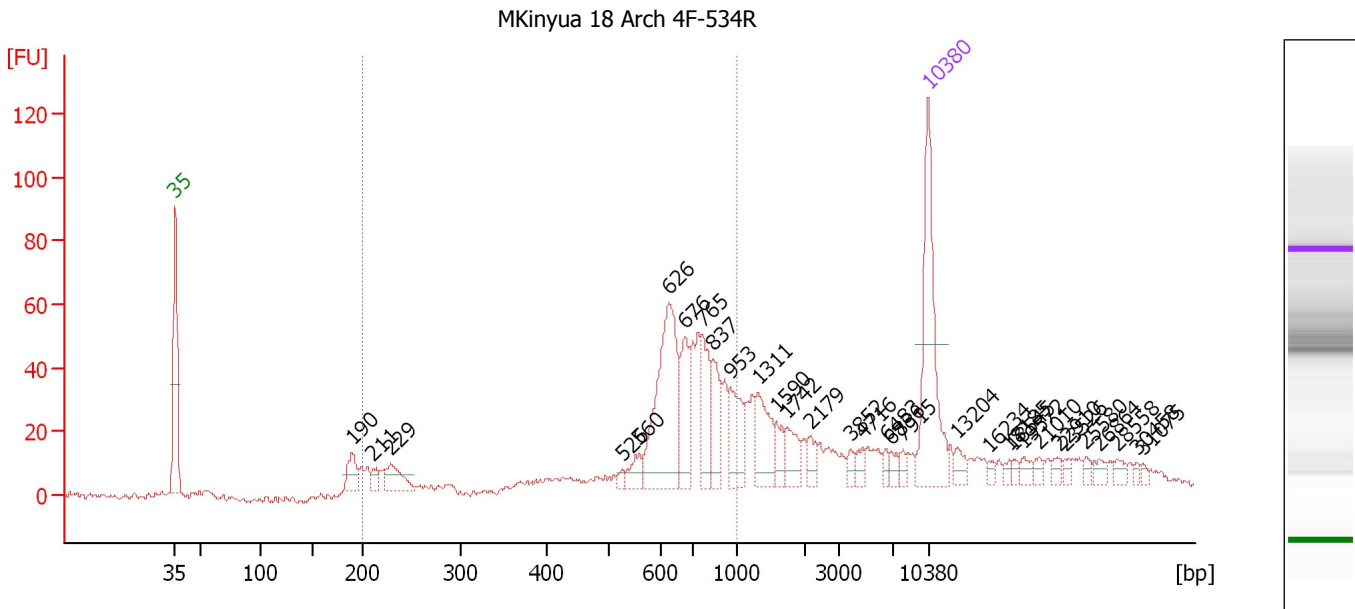
Region table for sample 5 : LStiemsma 33 Nested

| 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 | 676 | 1,010.11 | 1,268.5 | 2,294.5 | 87 | 12.5 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
 Modified: 10/16/2018 5:16:17 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : MKinyua 18 Arch 4F-534R

Number of peaks found: 32 Corr. Area 1: 479.3
 Noise: 0.9

Peak table for sample 6 : MKinyua 18 Arch 4F-534R

| 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 | 190 | 20.54 | 164.2 | | 59.41 |
| 3 | 211 | 8.97 | 64.4 | | 61.39 |
| 4 | 229 | 24.98 | 165.5 | | 63.03 |
| 5 | 526 | 3.88 | 11.2 | | 84.54 |
| 6 | 560 | 12.77 | 34.5 | | 86.18 |
| 7 | 626 | 111.79 | 270.6 | | 88.86 |
| 8 | 676 | 37.18 | 83.4 | | 90.39 |
| 9 | 765 | 35.29 | 69.9 | | 92.03 |
| 10 | 837 | 30.19 | 54.6 | | 93.02 |
| 11 | 953 | 32.68 | 52.0 | | 94.61 |
| 12 | 1,311 | 30.44 | 35.2 | | 97.19 |
| 13 | 1,590 | 11.89 | 11.3 | | 98.92 |
| 14 | 1,742 | 16.20 | 14.1 | | 99.86 |
| 15 | 2,179 | 8.85 | 6.2 | | 102.04 |
| 16 | 3,852 | 5.64 | 2.2 | | 105.76 |
| 17 | 4,716 | 5.67 | 1.8 | | 106.85 |
| 18 | 6,483 | 4.18 | 1.0 | | 109.08 |
| 19 | 6,836 | 5.08 | 1.1 | | 109.53 |
| 20 | 7,915 | 3.93 | 0.8 | | 110.62 |
| 21 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 22 | 13,204 | 0.00 | 0.0 | | 115.73 |
| 23 | 16,234 | 0.00 | 0.0 | | 118.65 |
| 24 | 18,185 | 0.00 | 0.0 | | 120.54 |
| 25 | 18,545 | 0.00 | 0.0 | | 120.88 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad


Created: 10/16/2018 4:49:15 PM
 Modified: 10/16/2018 5:16:17 PM

Electropherogram Summary Continued ...

... Peak table for sample 6 : MKinyua 18 Arch 4F-534R

| Peak | Size [bp] | Conc. [pg/ μ l] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|------|-----------|---------------------|-------------------|--------------|----------------------------|
| 26 | 19,572 | 0.00 | 0.0 | | 121.87 |
| 27 | 21,010 | 0.00 | 0.0 | | 123.26 |
| 28 | 22,910 | 0.00 | 0.0 | | 125.10 |
| 29 | 23,526 | 0.00 | 0.0 | | 125.69 |
| 30 | 25,580 | 0.00 | 0.0 | | 127.67 |
| 31 | 26,864 | 0.00 | 0.0 | | 128.91 |
| 32 | 28,558 | 0.00 | 0.0 | | 130.55 |
| 33 | 30,458 | 0.00 | 0.0 | | 132.38 |
| 34 | 31,075 | 0.00 | 0.0 | | 132.98 |

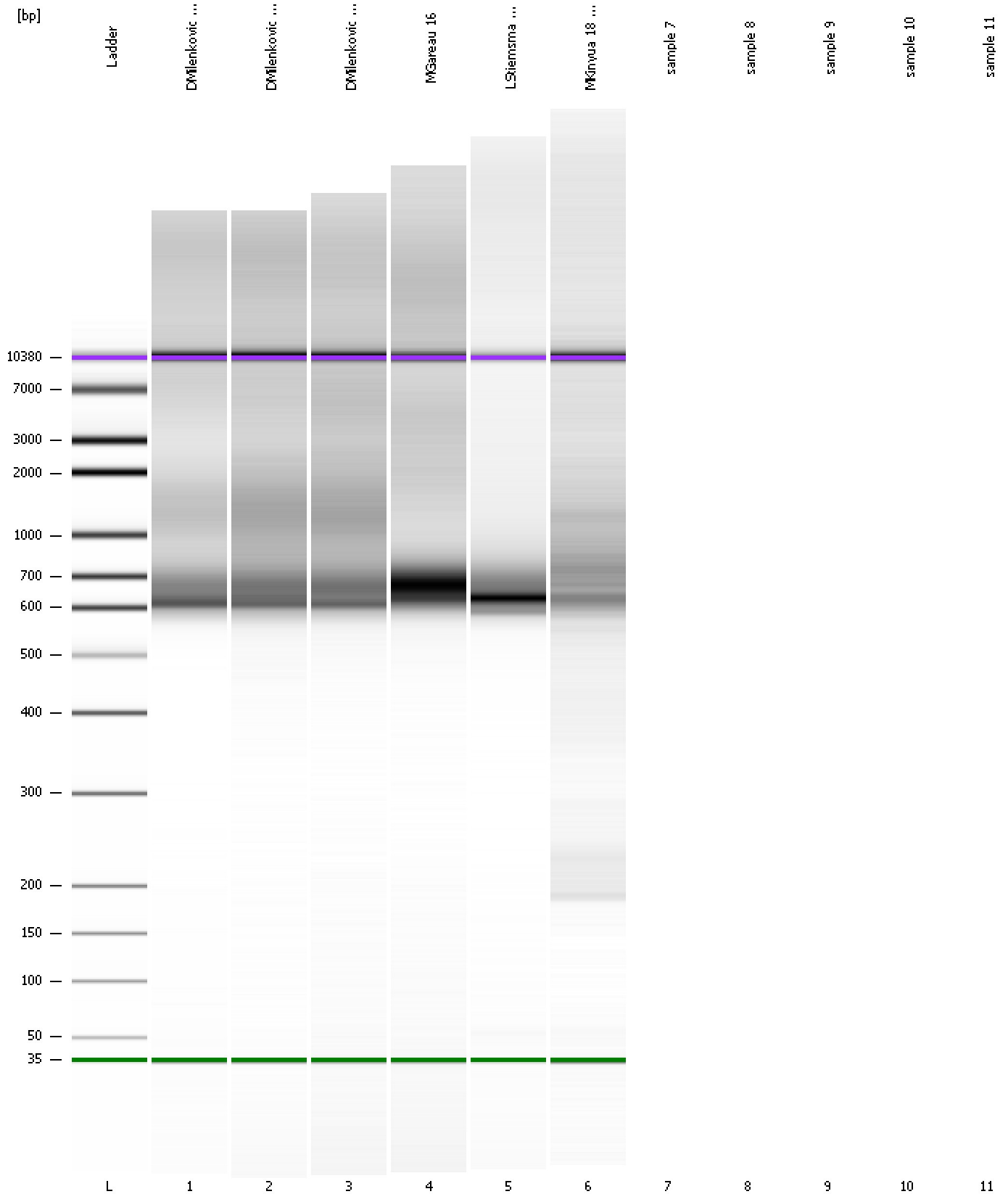
Region table for sample 6 : MKinyua 18 Arch 4F-534R

| From [bp] | To [bp] | Average Size [bp] | Conc. [pg/ μ l] | Corr. Area | Molarity [pmol/l] | Co % of lor Total | Size distribution in CV [%] |
|-----------|---------|-------------------|---------------------|------------|-------------------|--|-----------------------------|
| 200 | 1,000 | 665 | 380.00 | 479.3 | 1,084.7 |  55 | 27.2 |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
Modified: 10/16/2018 5:16:17 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2018-10-16\2018-10-16_001.xad

Created: 10/16/2018 4:49:15 PM
Modified: 10/16/2018 5:16:17 PM

Invalid Samples

Sample 7 has not been run, no results available.

Sample 8 has not been run, no results available.

Sample 9 has not been run, no results available.

Sample 10 has not been run, no results available.

Sample 11 has not been run, no results available.