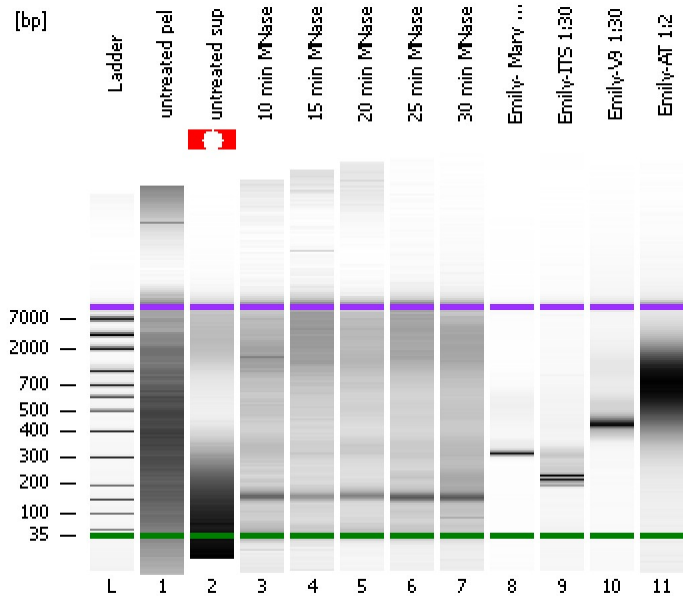


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
Modified: 2/26/2015 3:18:29 PM

Electrophoresis File Run Summary



Instrument Information:

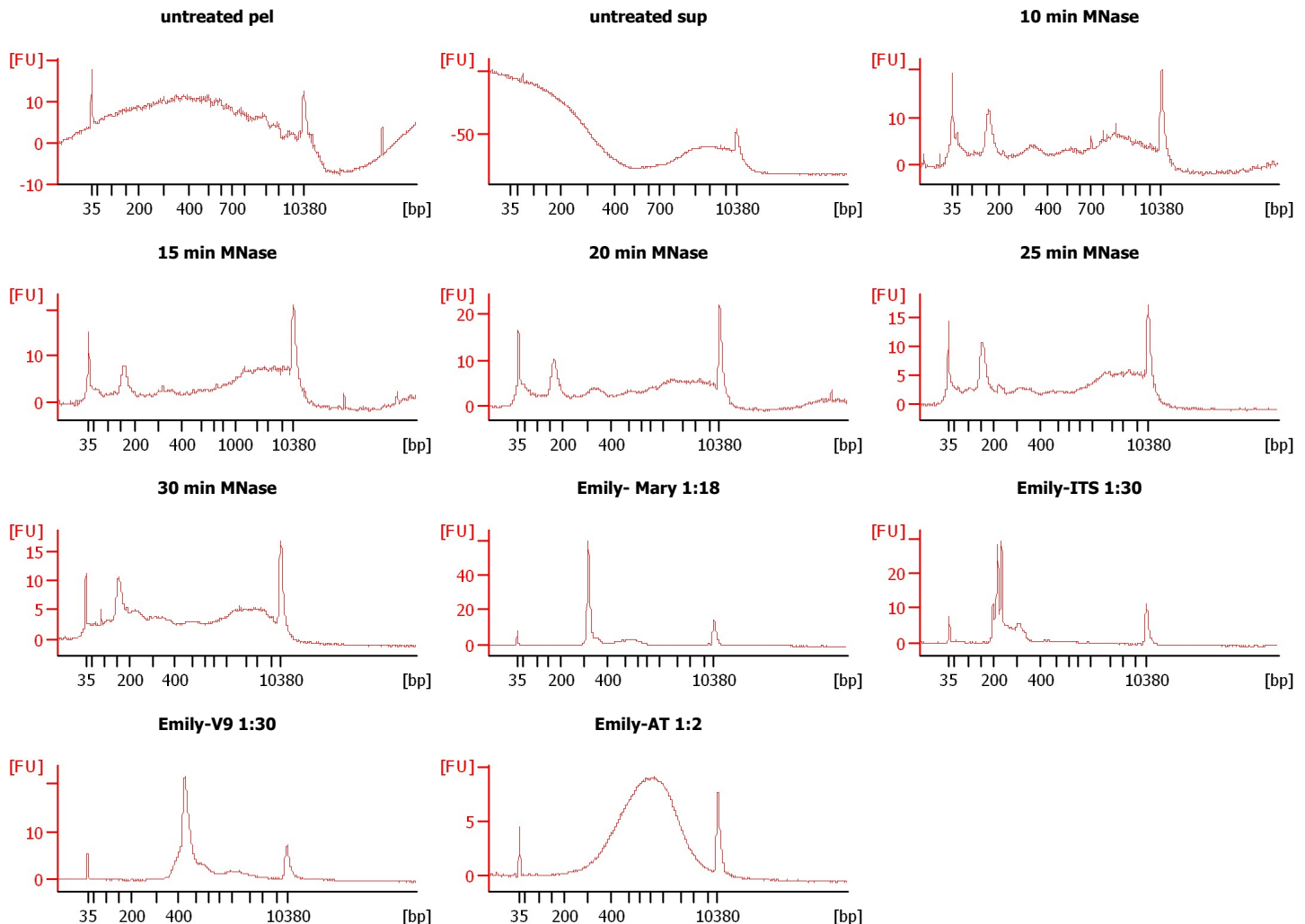
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electrophoresis File Run Summary (Chip Summary)

| Sample Name | Sample Comment | Rest. Digest | Status | Observation | Result Label | Result Color |
|------------------|----------------|--------------------------|--------|-------------|--------------|--------------|
| untreated pel | | <input type="checkbox"/> | | | | |
| untreated sup | | <input type="checkbox"/> | | | | |
| 10 min MNase | | <input type="checkbox"/> | | | | |
| 15 min MNase | | <input type="checkbox"/> | | | | |
| 20 min MNase | | <input type="checkbox"/> | | | | |
| 25 min MNase | | <input type="checkbox"/> | | | | |
| 30 min MNase | | <input type="checkbox"/> | | | | |
| Emily- Mary 1:18 | | <input type="checkbox"/> | | | | |
| Emily-ITS 1:30 | | <input type="checkbox"/> | | | | |
| Emily-V9 1:30 | | <input type="checkbox"/> | | | | |
| Emily-AT 1:2 | | <input type="checkbox"/> | | | | |
| Ladder | | <input type="checkbox"/> | | | | |

Chip Lot #

Reagent Kit Lot #

Chip Comments :

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
Modified: 2/26/2015 3:18:29 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

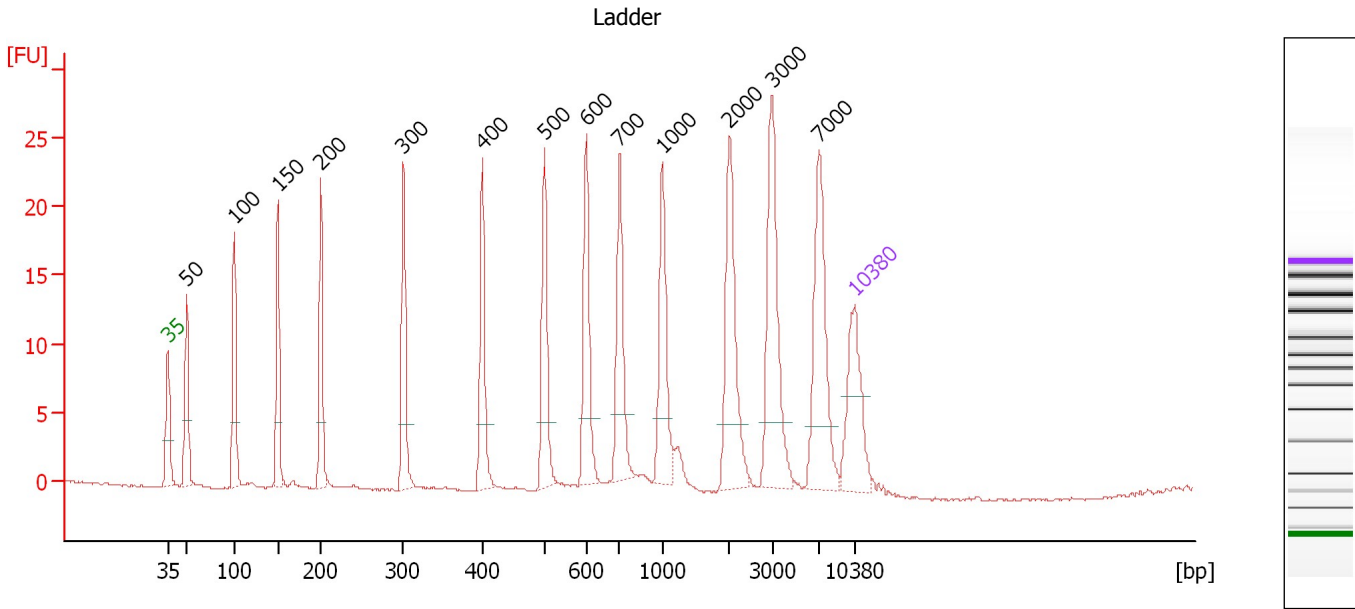
Ladder

| 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\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

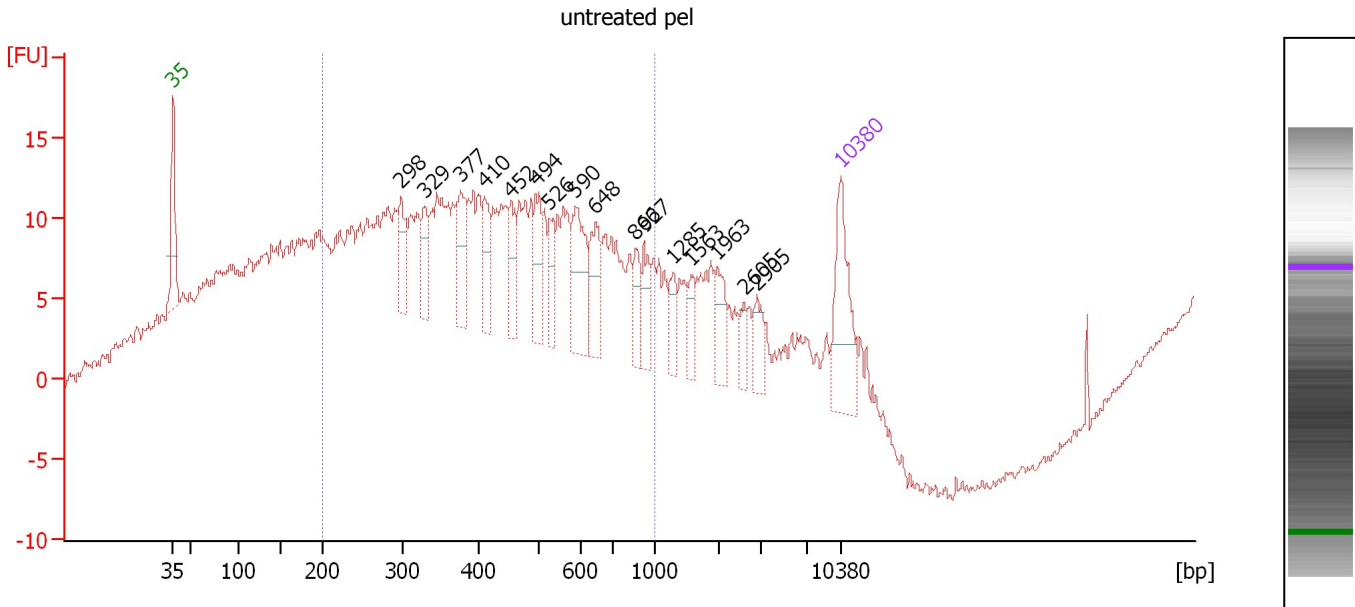
Peak table for Ladder

| 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 | 44.90 |
| 3 | 100 | 150.00 | 2,272.7 | Ladder Peak | 49.79 |
| 4 | 150 | 150.00 | 1,515.2 | Ladder Peak | 54.20 |
| 5 | 200 | 150.00 | 1,136.4 | Ladder Peak | 58.60 |
| 6 | 300 | 150.00 | 757.6 | Ladder Peak | 67.02 |
| 7 | 400 | 150.00 | 568.2 | Ladder Peak | 75.07 |
| 8 | 500 | 150.00 | 454.5 | Ladder Peak | 81.37 |
| 9 | 600 | 150.00 | 378.8 | Ladder Peak | 85.66 |
| 10 | 700 | 150.00 | 324.7 | Ladder Peak | 89.03 |
| 11 | 1,000 | 150.00 | 227.3 | Ladder Peak | 93.43 |
| 12 | 2,000 | 150.00 | 113.6 | Ladder Peak | 100.28 |
| 13 | 3,000 | 150.00 | 75.8 | Ladder Peak | 104.58 |
| 14 | 7,000 | 150.00 | 32.5 | Ladder Peak | 109.36 |
| 15 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : untreated pel

Number of peaks found: 16 Corr. Area 1: 354.4
 Noise: 0.3

Peak table for sample 1 : untreated pel

| 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 | 298 | 39.88 | 202.6 | | 66.87 |
| 3 | 329 | 35.79 | 164.6 | | 69.39 |
| 4 | 377 | 53.54 | 215.4 | | 73.19 |
| 5 | 410 | 44.21 | 163.3 | | 75.71 |
| 6 | 452 | 37.97 | 127.3 | | 78.34 |
| 7 | 494 | 46.14 | 141.4 | | 81.02 |
| 8 | 526 | 31.77 | 91.6 | | 82.47 |
| 9 | 590 | 75.65 | 194.4 | | 85.21 |
| 10 | 648 | 50.07 | 117.1 | | 87.28 |
| 11 | 866 | 26.55 | 46.4 | | 91.47 |
| 12 | 927 | 30.38 | 49.6 | | 92.37 |
| 13 | 1,285 | 18.70 | 22.0 | | 95.39 |
| 14 | 1,563 | 17.21 | 16.7 | | 97.29 |
| 15 | 1,963 | 29.68 | 22.9 | | 100.03 |
| 16 | 2,605 | 14.58 | 8.5 | | 102.88 |
| 17 | 2,905 | 21.38 | 11.2 | | 104.17 |
| 18 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

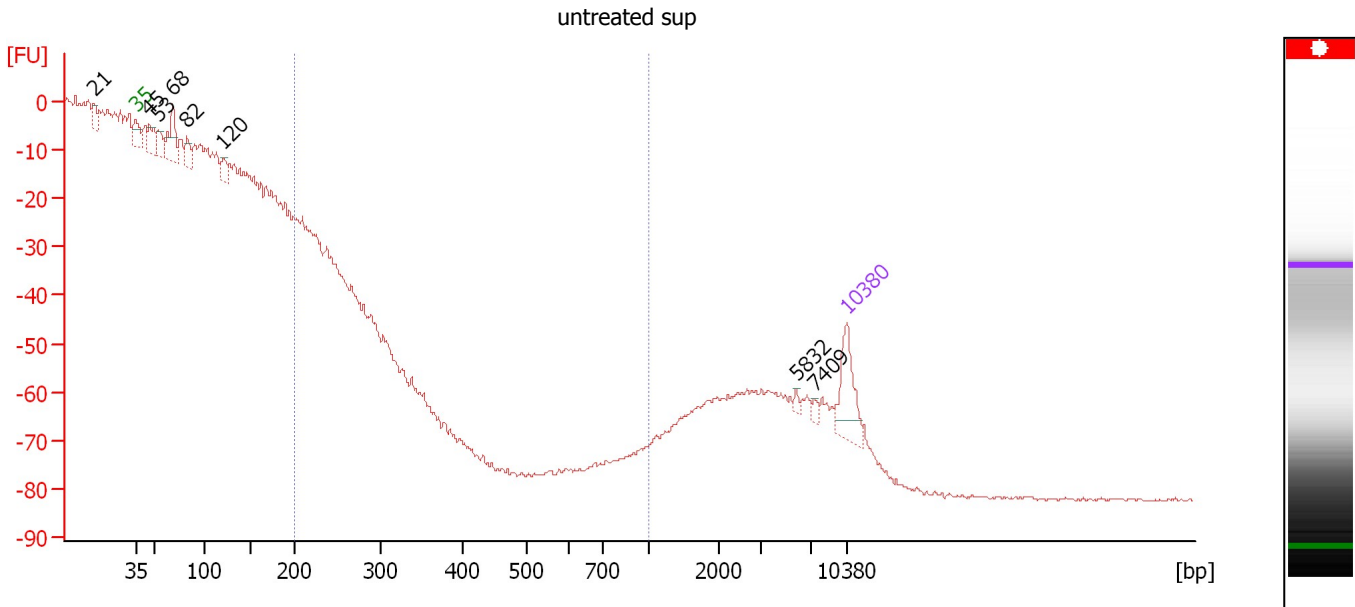
Region table for sample 1 : untreated pel

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 436 | 1,000 | 354.4 | 60 | 40.3 | 1,858.15 | 7,940.4 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : untreated sup

Number of peaks found: 8 Corr. Area 1: 0.0
 Noise: 0.4

Peak table for sample 2 : untreated sup

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|------|-----------|---------------|-------------------|--------------|----------------------------|
| 1 | 21 | 0.00 | 0.0 | | 38.91 |
| 2 | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | 45 | 55.96 | 1,869.8 | | 44.31 |
| 4 | 53 | 36.69 | 1,047.2 | | 45.20 |
| 5 | 68 | 93.25 | 2,091.0 | | 46.62 |
| 6 | 82 | 38.55 | 716.5 | | 47.99 |
| 7 | 120 | 32.87 | 415.0 | | 51.55 |
| 8 | 5,832 | 5.88 | 1.5 | | 107.96 |
| 9 | 7,409 | 7.53 | 1.5 | | 109.80 |
| 10 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

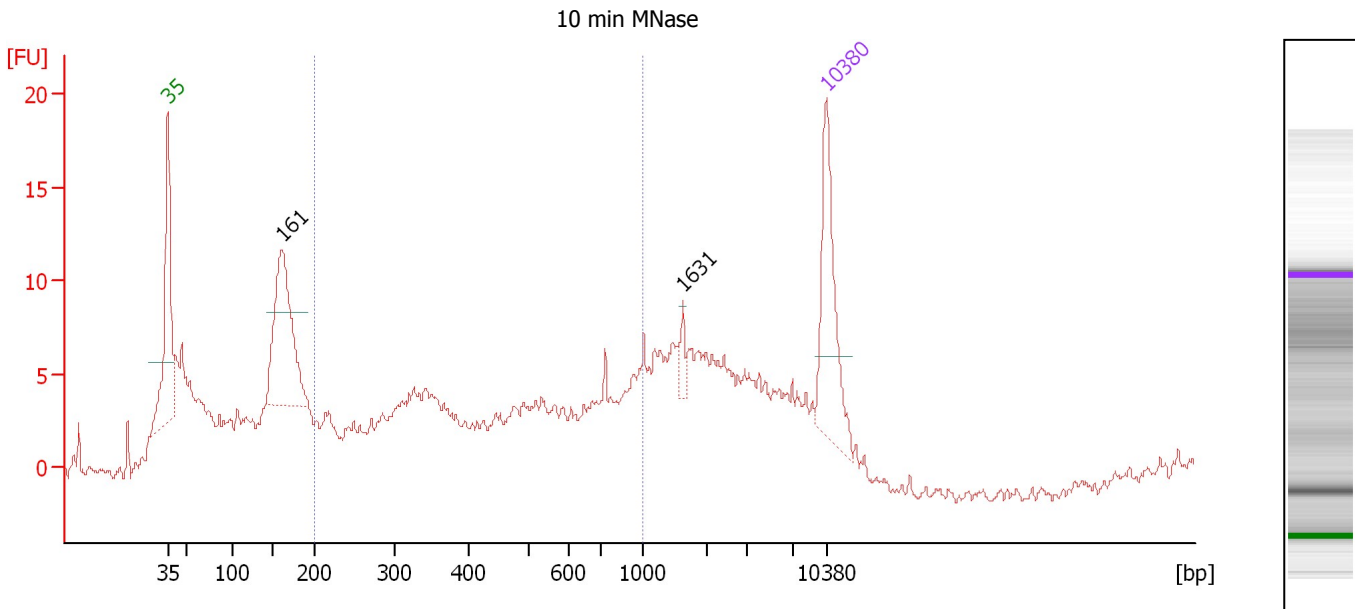
Region table for sample 2 : untreated sup

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 0 | 1,000 | 0.0 | 0 | 0.0 | 0.00 | 0.0 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : 10 min MNase

Number of peaks found: 2 Corr. Area 1: 113.3
 Noise: 0.3

Peak table for sample 3 : 10 min MNase

| 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 | 161 | 178.26 | 1,681.7 | | 55.13 |
| 3 | 1,631 | 9.08 | 8.4 | | 97.75 |
| 4 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

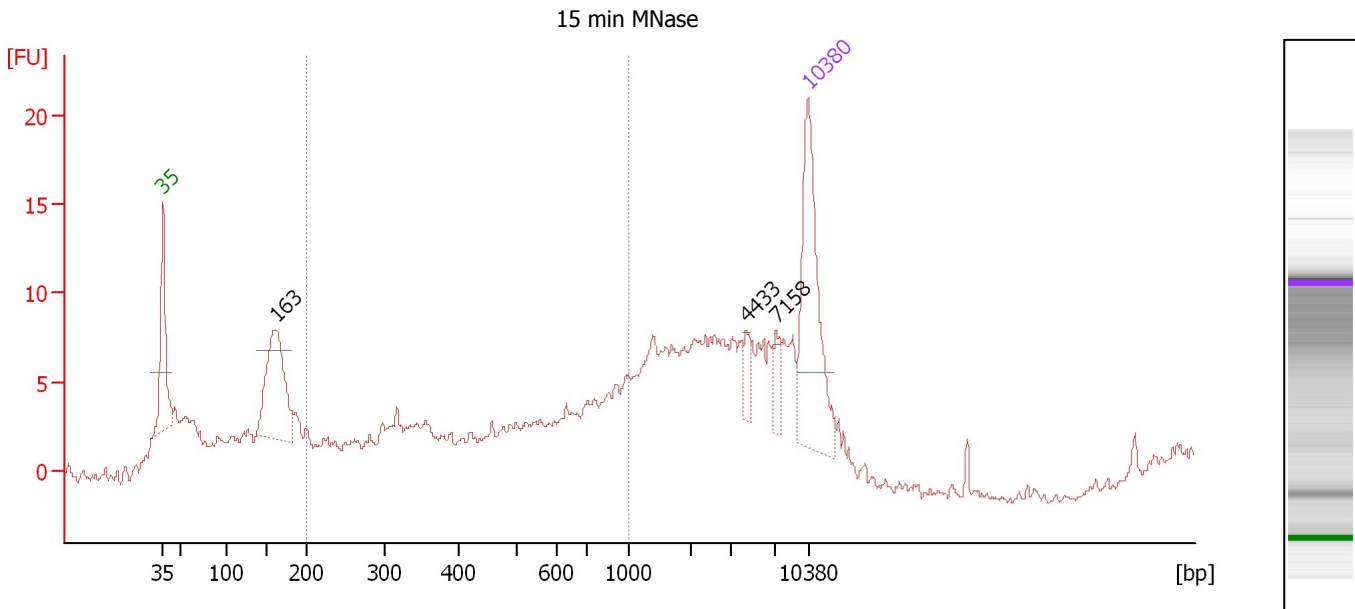
Region table for sample 3 : 10 min MNase

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 501 | 1,000 | 113.3 | 37 | 42.6 | 501.06 | 1,956.1 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : 15 min MNase

Number of peaks found: 3 Corr. Area 1: 98.6
 Noise: 0.2

Peak table for sample 4 : 15 min MNase

| 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 | 163 | 99.13 | 918.9 | | 55.38 |
| 3 | 4,433 | 7.61 | 2.6 | | 106.29 |
| 4 | 7,158 | 8.96 | 1.9 | | 109.53 |
| 5 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

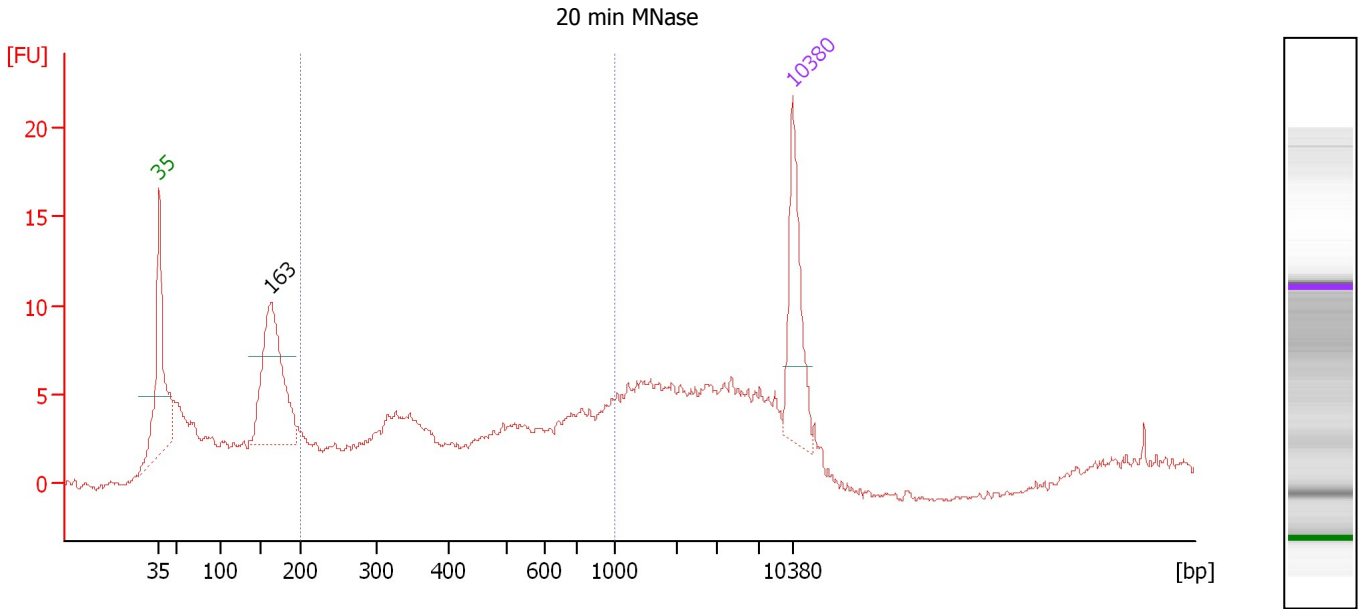
Region table for sample 4 : 15 min MNase

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 518 | 1,000 | 98.6 | 33 | 42.6 | 316.42 | 1,215.6 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : 20 min MNase

Number of peaks found: 1 Corr. Area 1: 105.1
 Noise: 0.2

Peak table for sample 5 : 20 min MNase

| 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 | 163 | 210.64 | 1,953.4 | | 55.37 |
| 3 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

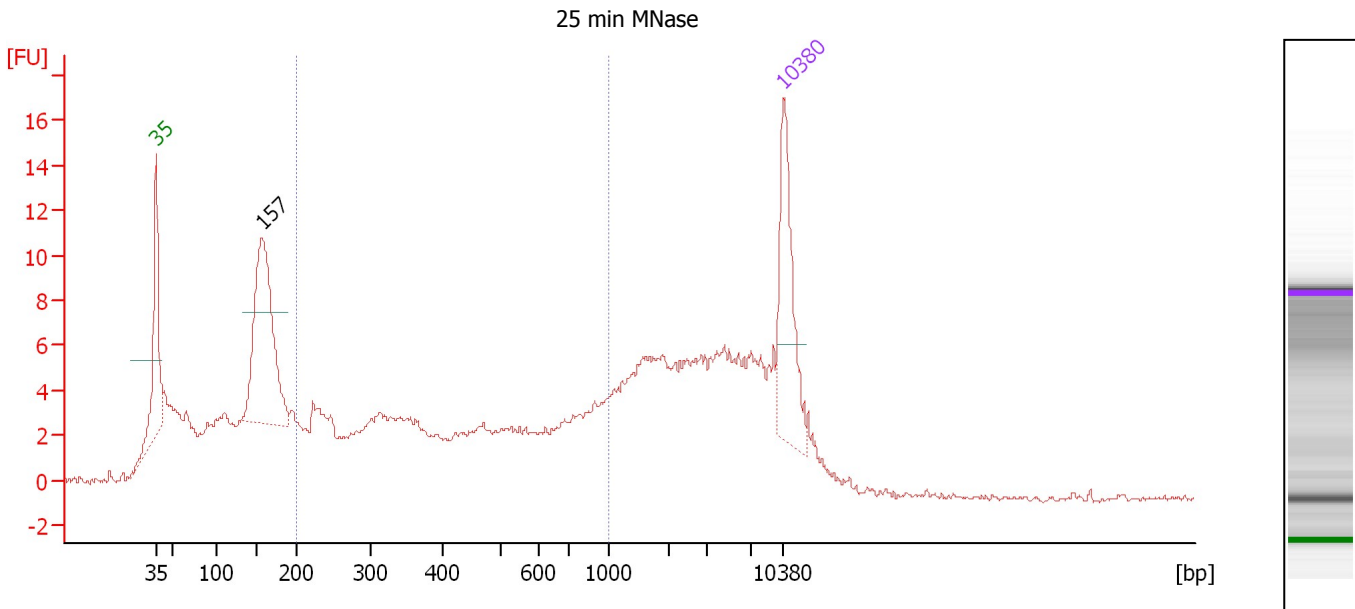
Region table for sample 5 : 20 min MNase

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 496 | 1,000 | 105.1 | 38 | 42.7 | 483.49 | 1,909.6 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : 25 min MNase

Number of peaks found: 1 Corr. Area 1: 115.0
 Noise: 0.1

Peak table for sample 6 : 25 min MNase

| 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 | 157 | 221.38 | 2,131.4 | | 54.85 |
| 3 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

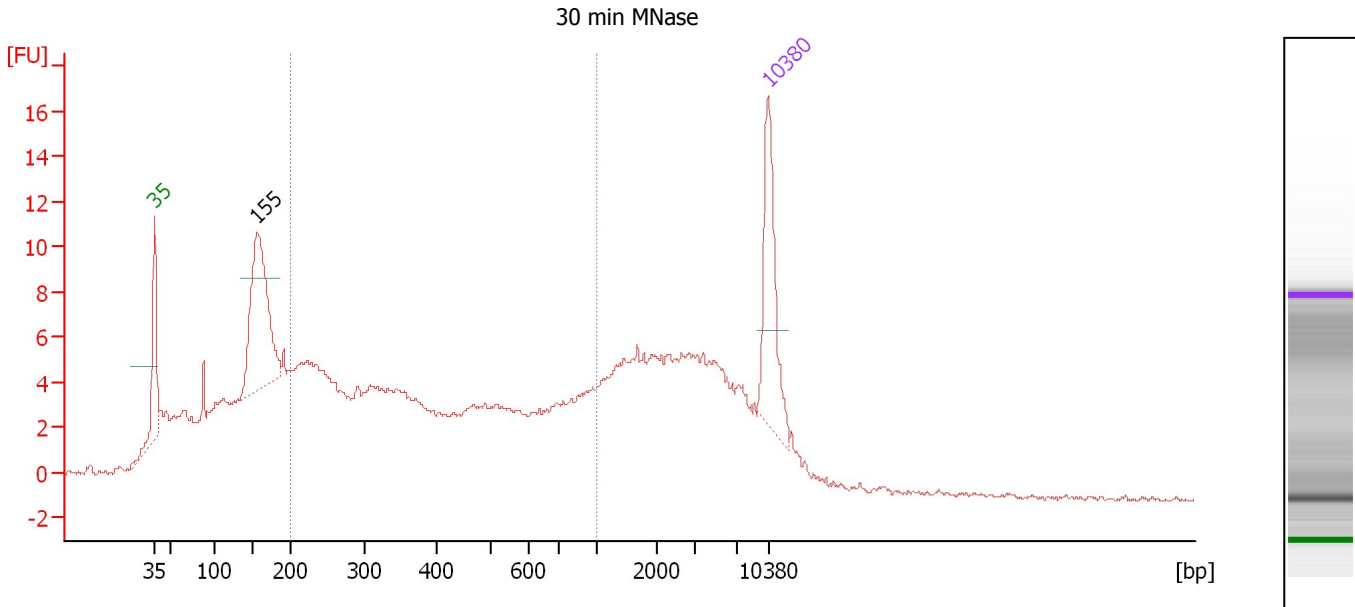
Region table for sample 6 : 25 min MNase

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 476 | 1,000 | 115.0 | 35 | 44.5 | 583.15 | 2,429.0 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : 30 min MNase

Number of peaks found: 1 Corr. Area 1: 161.9
 Noise: 0.1

Peak table for sample 7 : 30 min MNase

| 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 | 155 | 202.26 | 1,972.7 | | 54.67 |
| 3 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

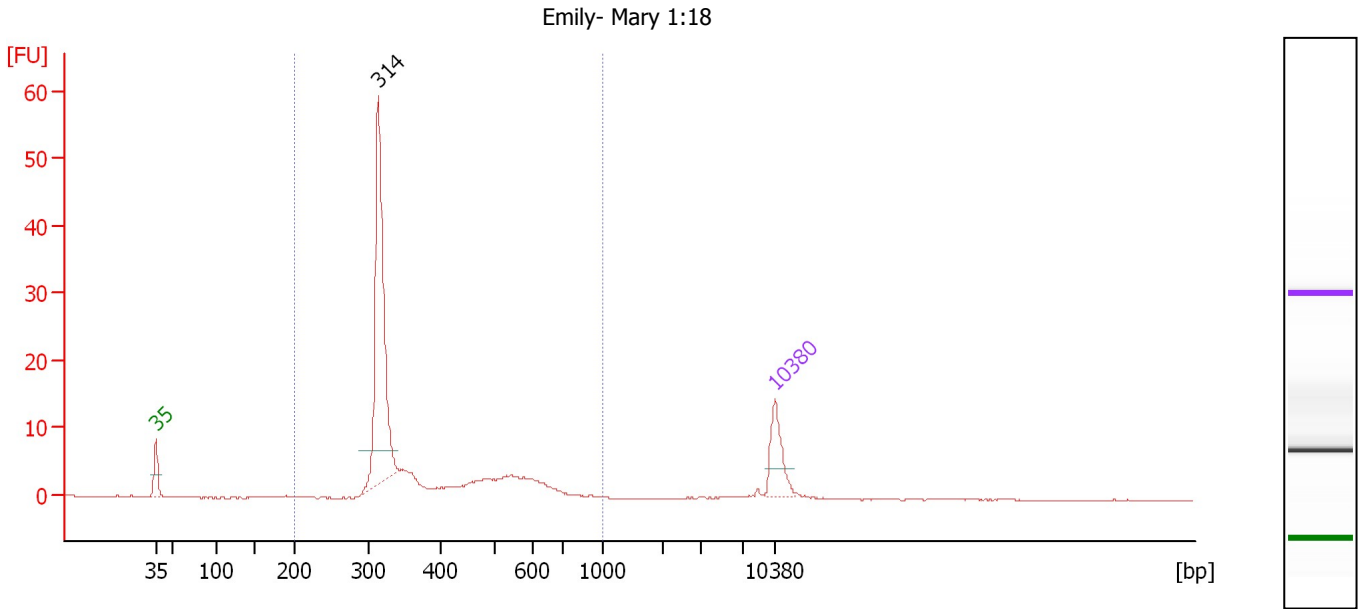
Region table for sample 7 : 30 min MNase

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 448 | 1,000 | 161.9 | 44 | 45.8 | 989.19 | 4,354.7 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : Emily- Mary 1:18

Number of peaks found: 1 Corr. Area 1: 148.7
 Noise: 0.1

Peak table for sample 8 : Emily- Mary 1:18

| 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 | 314 | 576.82 | 2,784.0 | | 68.14 |
| 3 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

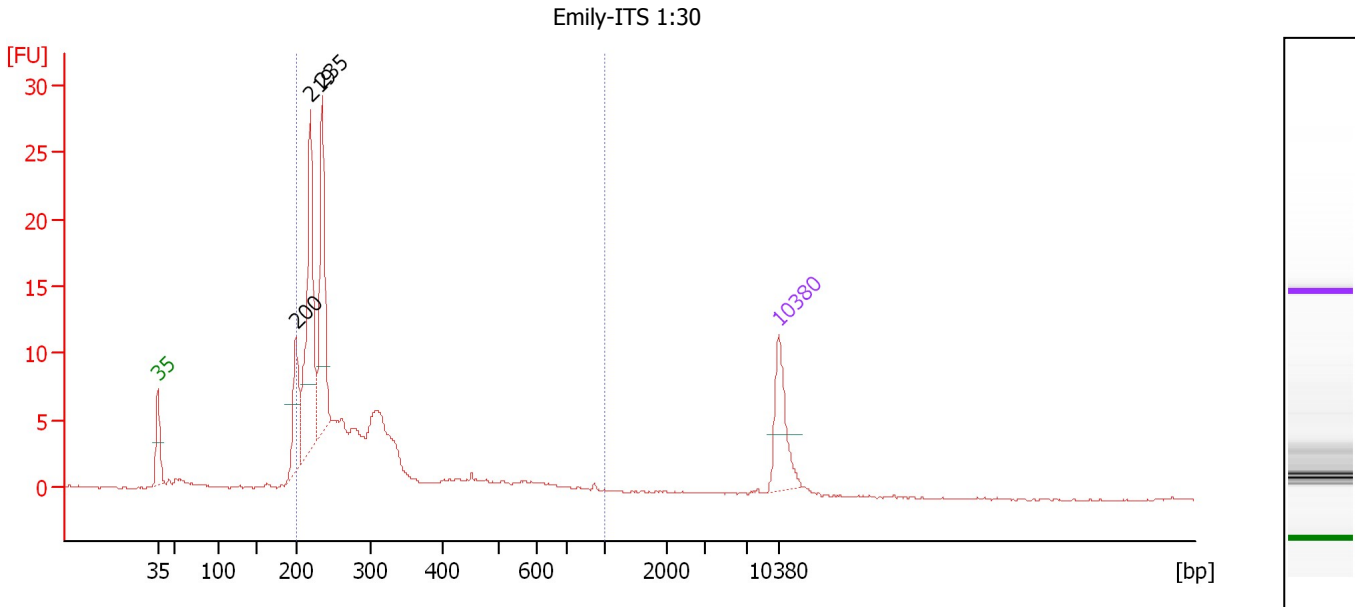
Region table for sample 8 : Emily- Mary 1:18

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 386 | 1,000 | 148.7 | 97 | 29.2 | 961.81 | 4,112.2 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : Emily-ITS 1:30

Number of peaks found: 3 Corr. Area 1: 153.6
 Noise: 0.1

Peak table for sample 9 : Emily-ITS 1:30

| 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 | 200 | 115.31 | 875.3 | | 58.56 |
| 3 | 219 | 322.34 | 2,231.8 | | 60.18 |
| 4 | 235 | 229.05 | 1,475.3 | | 61.57 |
| 5 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

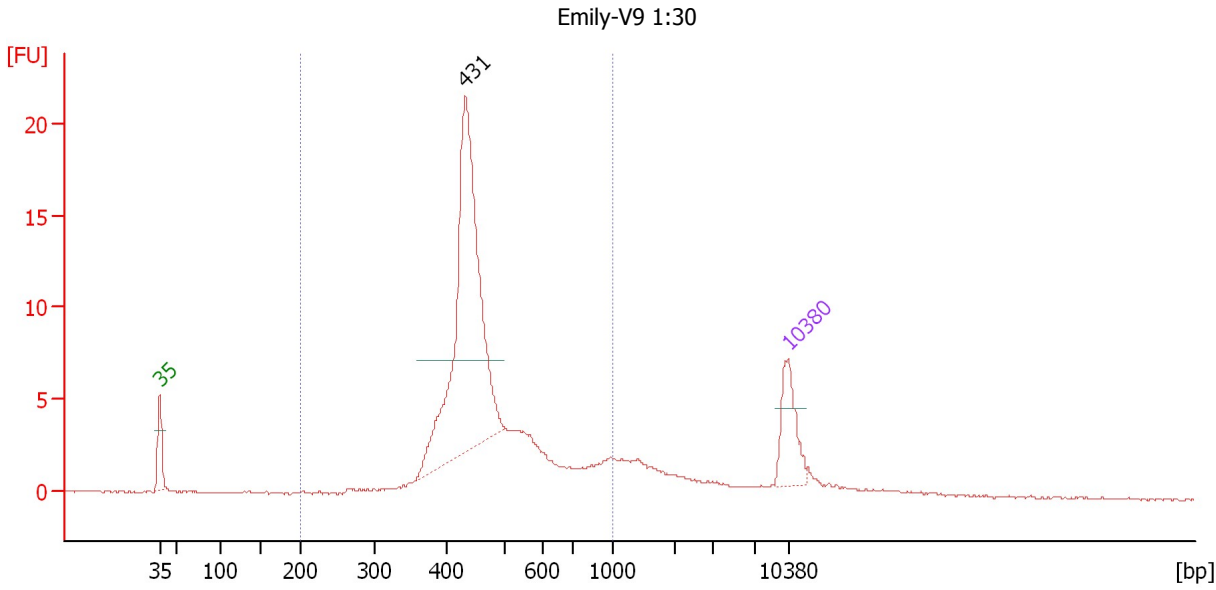
Region table for sample 9 : Emily-ITS 1:30

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 288 | 1,000 | 153.6 | 90 | 38.8 | 1,352.38 | 7,932.2 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : Emily-V9 1:30

Number of peaks found: 1 Corr. Area 1: 128.8
 Noise: 0.0

Peak table for sample 10 : Emily-V9 1:30

| 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 | 431 | 685.06 | 2,405.7 | | 77.05 |
| 3 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

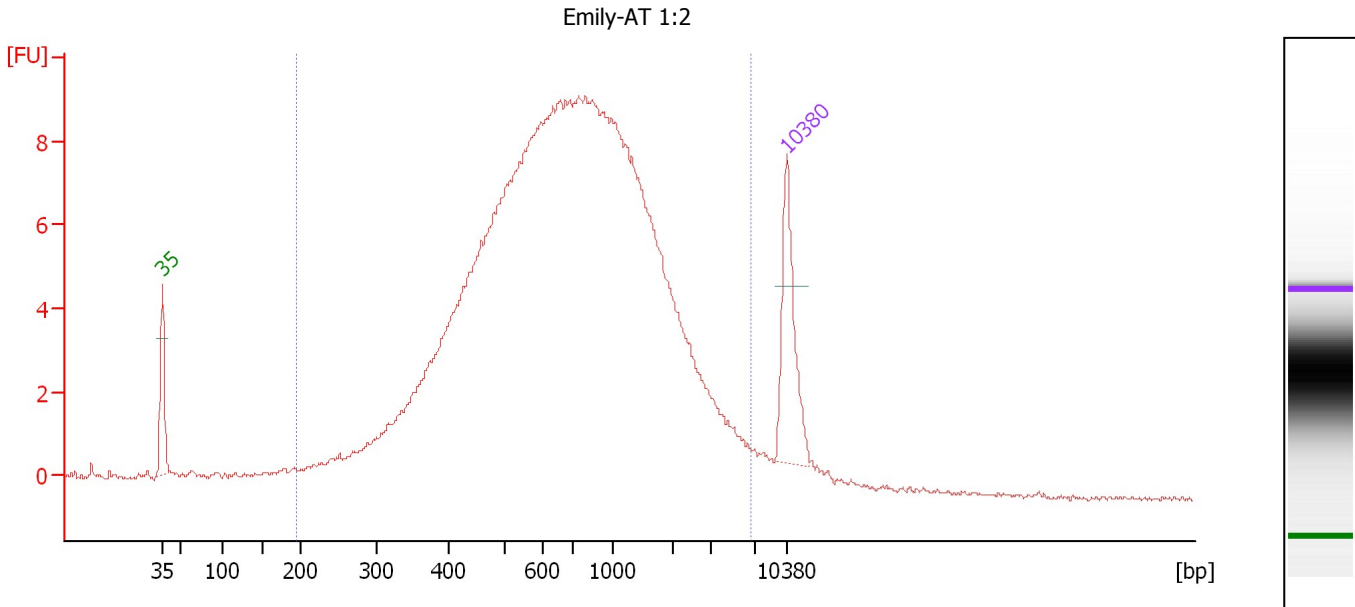
Region table for sample 10 : Emily-V9 1:30

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 485 | 1,000 | 128.8 | 83 | 25.3 | 1,222.49 | 4,053.0 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 PM

Electropherogram Summary Continued ...



Setpoint Deviations for sample 11 : Emily-AT 1:2

Height Threshold [FU] : 4

Overall Results for sample 11 : Emily-AT 1:2

Number of peaks found: 0 Corr. Area 1: 242.9
 Noise: 0.1

Peak table for sample 11 : Emily-AT 1:2

| 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 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

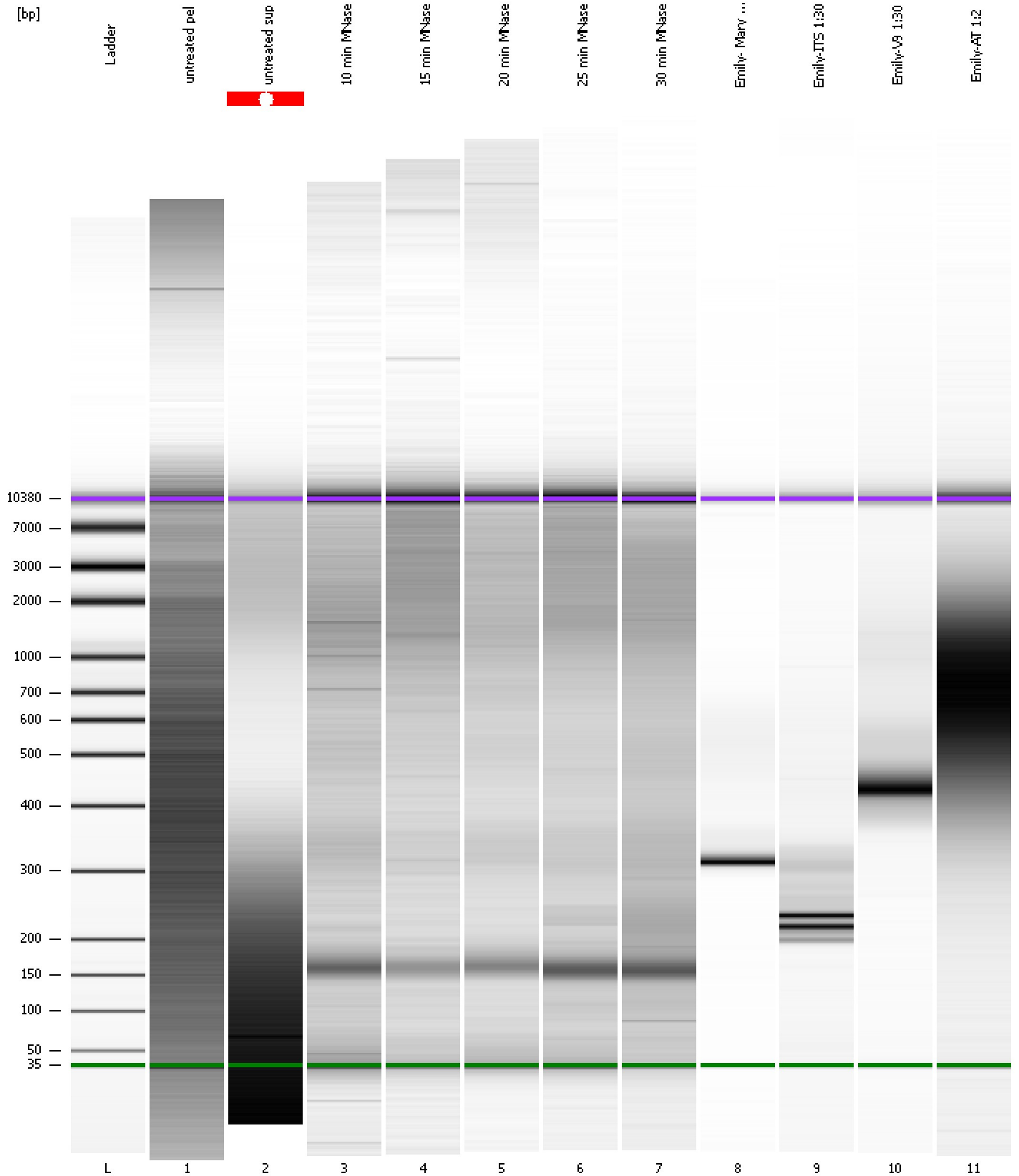
Region table for sample 11 : Emily-AT 1:2

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 193 | 1,005 | 6,806 | 242.9 | 98 | 86.5 | 2,437.89 | 6,033.8 | Blue |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
Modified: 2/26/2015 3:18:29 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad

Created: 2/26/2015 2:32:09 PM
 Modified: 2/26/2015 3:18:29 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 | | 2/26/2015 3:13:27 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-02-26\2015-02-26_002.xad) | | Instrument | Run | | 2/26/2015 2:32:14 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Product Number : G2938B | | Instrument | Run | | 2/26/2015 2:32:14 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Name : | | Instrument | Run | | 2/26/2015 2:32:14 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Vendor : Agilent Technologies | | Instrument | Run | | 2/26/2015 2:32:14 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Serial# : DE13701086 | | Instrument | Run | | 2/26/2015 2:32:14 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Firmware : C.01.069 | | Instrument | Run | | 2/26/2015 2:32:14 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Cartridge : Electrode | | Instrument | Run | | 2/26/2015 2:32:14 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |