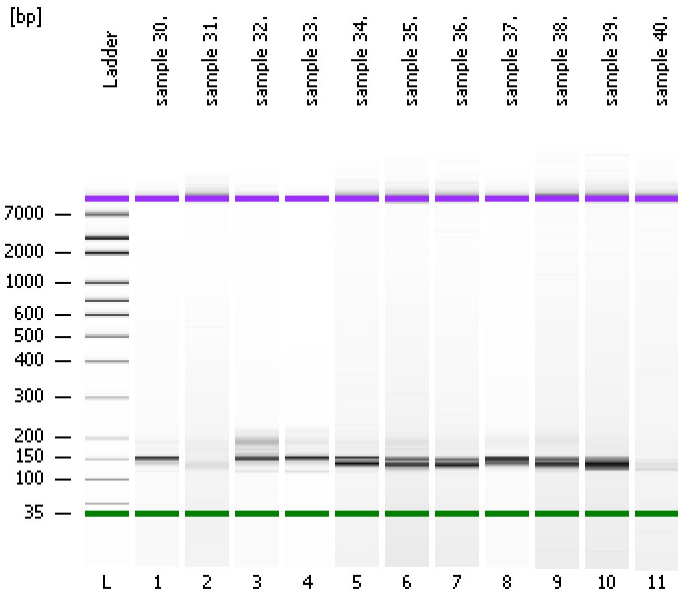


Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
Modified: 2/10/2015 1:56:57 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

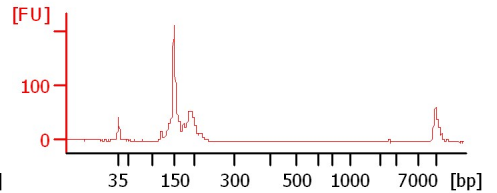
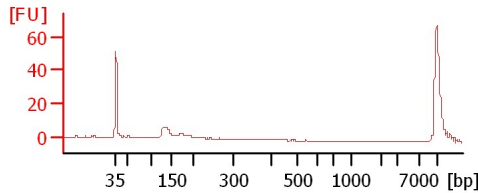
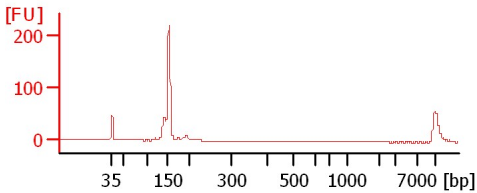
Chip Information:

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

sample 30.

sample 31.

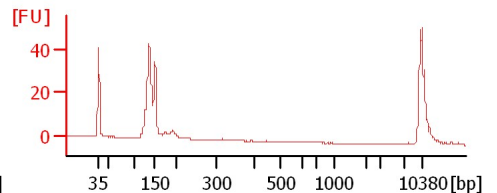
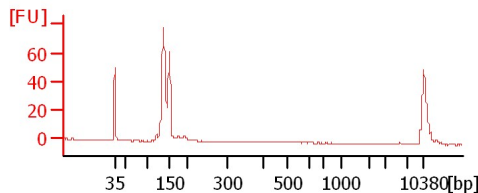
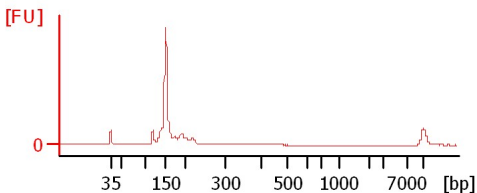
sample 32.



sample 33.

sample 34.

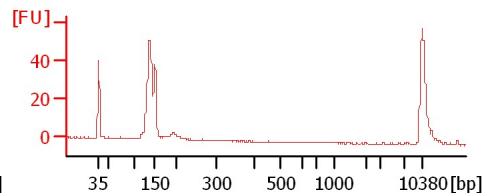
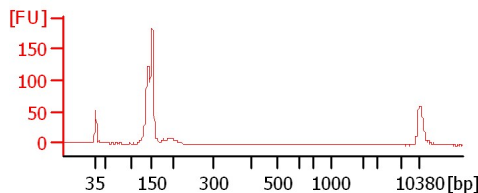
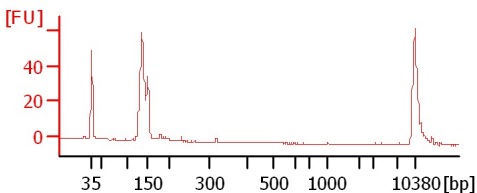
sample 35.



sample 36.

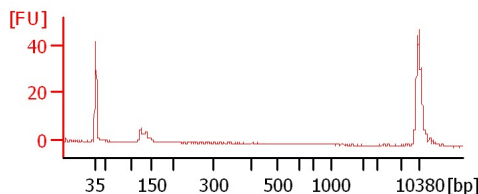
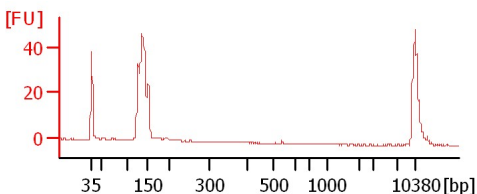
sample 37.

sample 38.



sample 39.

sample 40.



Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
Modified: 2/10/2015 1:56:57 PM

Electrophoresis File Run Summary (Chip Summary)

| Sample Name | Sample Comment | Rest. Digest | Status | Observation | Result Label | Result Color |
|-------------|----------------|--------------------------|--------|-------------|--------------|--------------|
| sample 30. | | <input type="checkbox"/> | ✓ | | | |
| sample 31. | | <input type="checkbox"/> | ✓ | | | |
| sample 32. | | <input type="checkbox"/> | ✓ | | | |
| sample 33. | | <input type="checkbox"/> | ✓ | | | |
| sample 34. | | <input type="checkbox"/> | ✓ | | | |
| sample 35. | | <input type="checkbox"/> | ✓ | | | |
| sample 36. | | <input type="checkbox"/> | ✓ | | | |
| sample 37. | | <input type="checkbox"/> | ✓ | | | |
| sample 38. | | <input type="checkbox"/> | ✓ | | | |
| sample 39. | | <input type="checkbox"/> | ✓ | | | |
| sample 40. | | <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\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
Modified: 2/10/2015 1:56:57 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

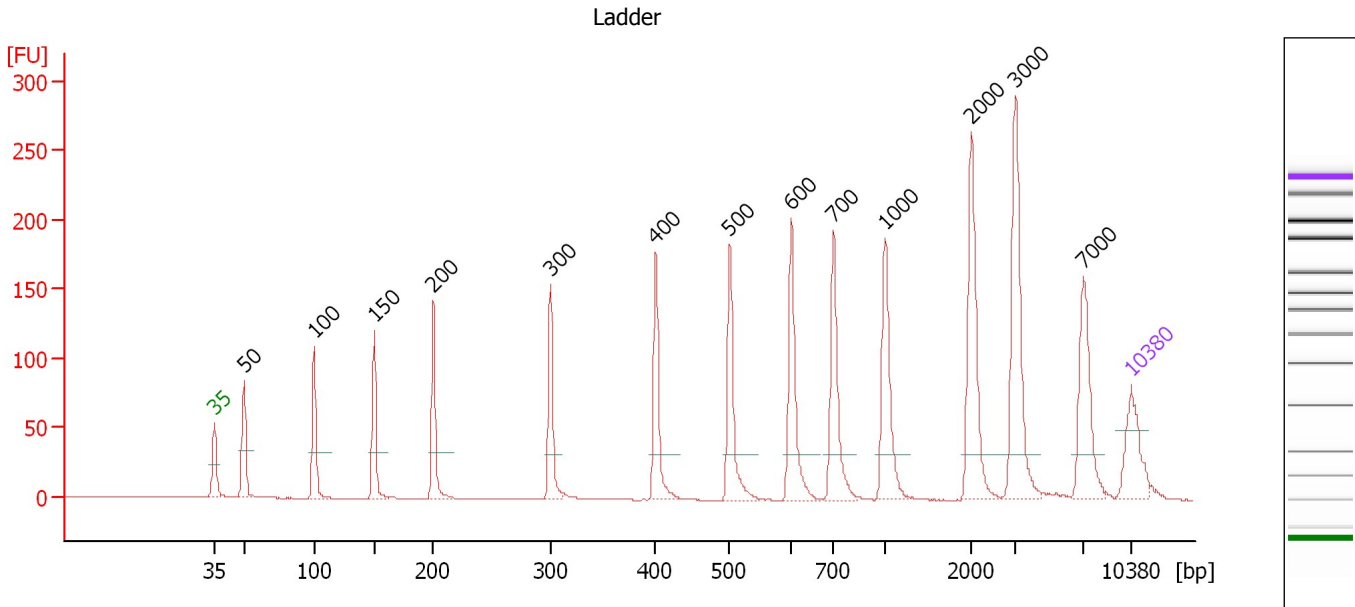
Ladder

| 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\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 PM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.1

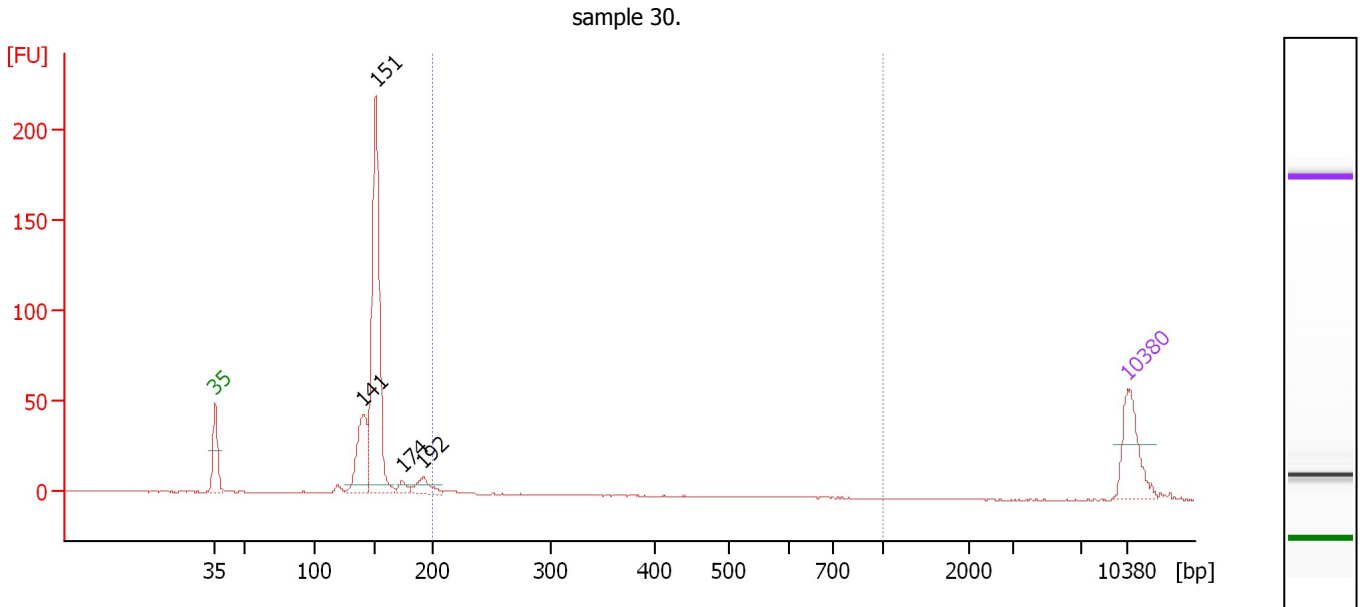
Peak table for Ladder

| 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 | 45.24 |
| 3 | 100 | 150.00 | 2,272.7 | Ladder Peak | 50.62 |
| 4 | 150 | 150.00 | 1,515.2 | Ladder Peak | 55.18 |
| 5 | 200 | 150.00 | 1,136.4 | Ladder Peak | 59.71 |
| 6 | 300 | 150.00 | 757.6 | Ladder Peak | 68.67 |
| 7 | 400 | 150.00 | 568.2 | Ladder Peak | 76.70 |
| 8 | 500 | 150.00 | 454.5 | Ladder Peak | 82.36 |
| 9 | 600 | 150.00 | 378.8 | Ladder Peak | 87.05 |
| 10 | 700 | 150.00 | 324.7 | Ladder Peak | 90.31 |
| 11 | 1,000 | 150.00 | 227.3 | Ladder Peak | 94.22 |
| 12 | 2,000 | 150.00 | 113.6 | Ladder Peak | 100.82 |
| 13 | 3,000 | 150.00 | 75.8 | Ladder Peak | 104.20 |
| 14 | 7,000 | 150.00 | 32.5 | Ladder Peak | 109.37 |
| 15 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 1 : sample 30.

Number of peaks found: 4 Corr. Area 1: 10.6
 Noise: 0.2

Peak table for sample 1 : sample 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 | 141 | 152.19 | 1,634.7 | | 54.37 |
| 3 | 151 | 540.00 | 5,404.9 | | 55.31 |
| 4 | 174 | 17.54 | 152.7 | | 57.35 |
| 5 | 192 | 38.24 | 302.4 | | 58.95 |
| 6 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

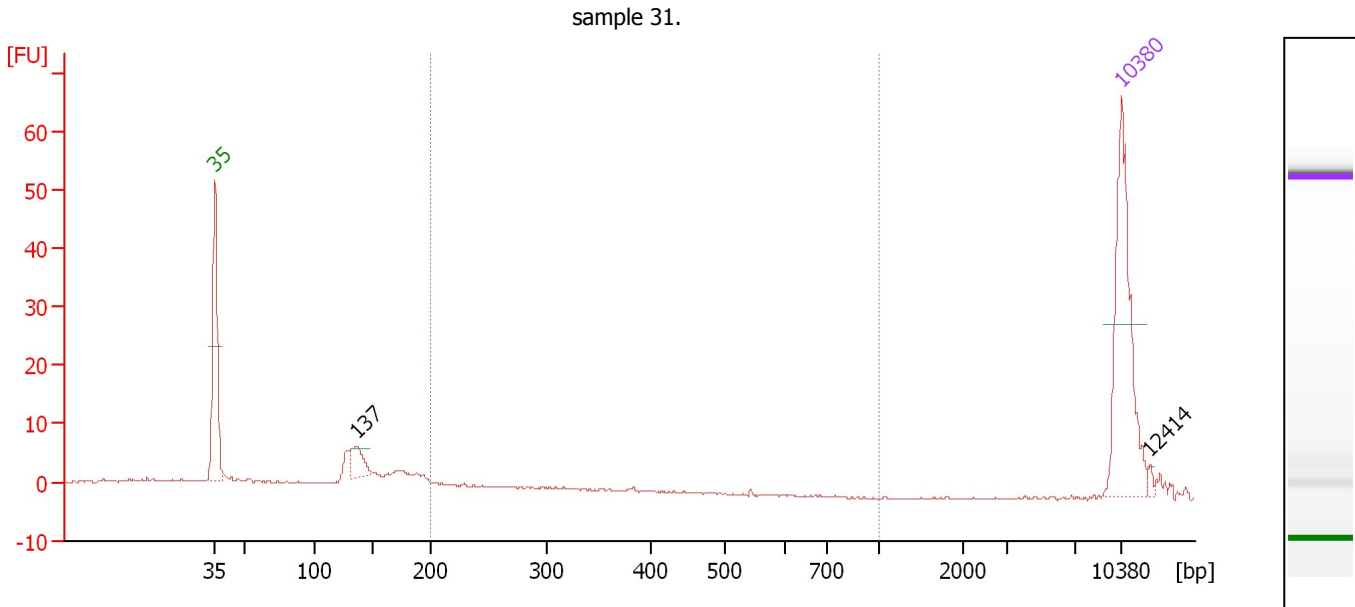
Region table for sample 1 : sample 30.

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 230 | 1,000 | 10.6 | 2 | 19.1 | 17.47 | 117.7 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 2 : sample 31.

Number of peaks found: 2 Corr. Area 1: 0.5
 Noise: 0.2

Peak table for sample 2 : sample 31.

| 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 | 137 | 20.84 | 231.3 | | 53.95 |
| 3 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 4 | 12,414 | 0.00 | 0.0 | | 115.18 |

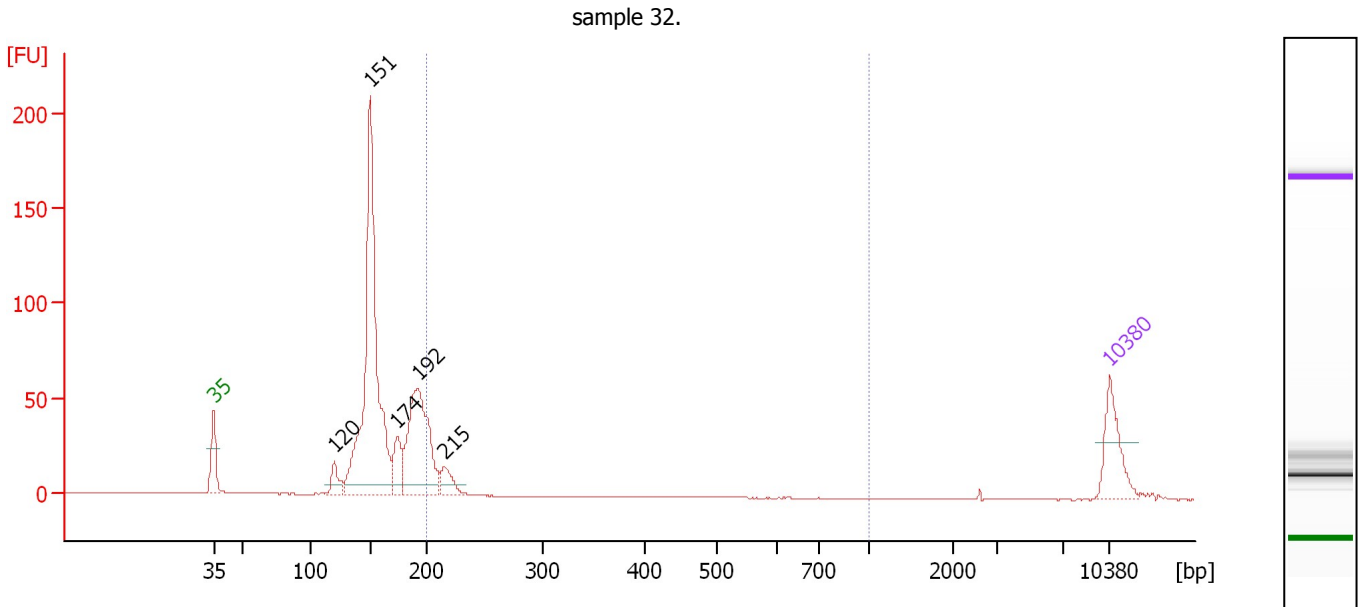
Region table for sample 2 : sample 31.

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 209 | 1,000 | 0.5 | 1 | 4.7 | 0.77 | 5.6 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 3 : sample 32.

Number of peaks found: 5 Corr. Area 1: 70.9
 Noise: 0.1

Peak table for sample 3 : sample 32.

| 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 | 120 | 44.48 | 563.3 | | 52.41 |
| 3 | 151 | 808.28 | 8,128.5 | | 55.24 |
| 4 | 174 | 77.50 | 676.1 | | 57.32 |
| 5 | 192 | 339.03 | 2,680.5 | | 58.95 |
| 6 | 215 | 49.13 | 346.6 | | 61.03 |
| 7 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

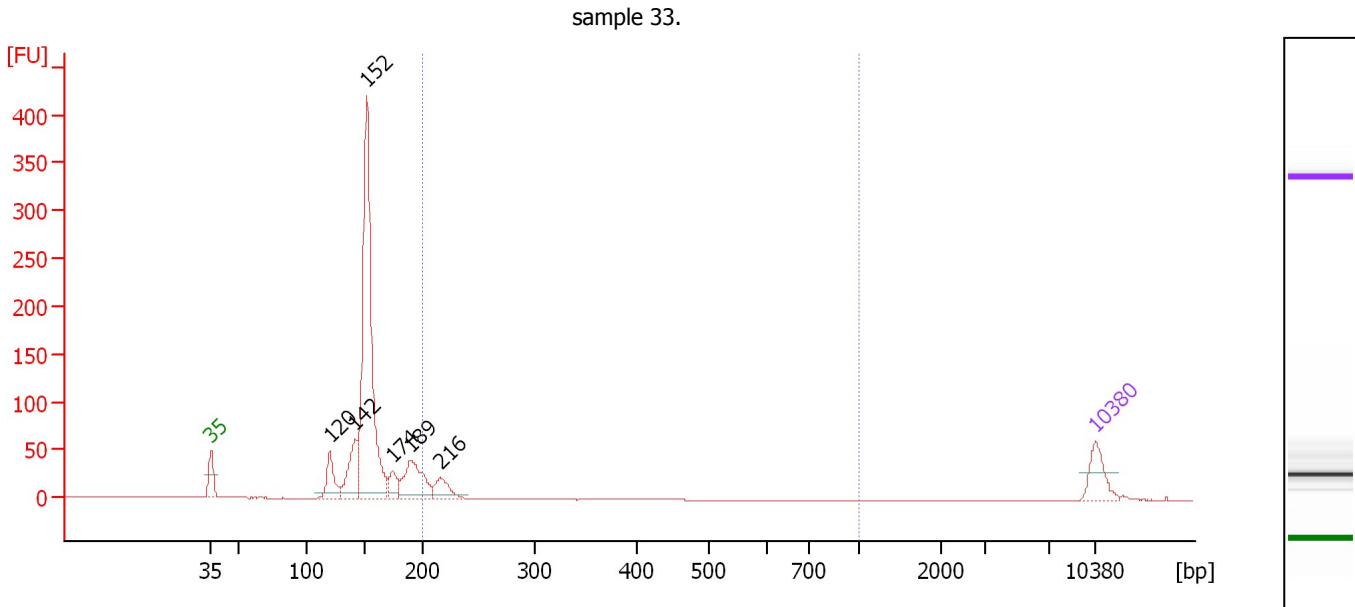
Region table for sample 3 : sample 32.

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 210 | 1,000 | 70.9 | 10 | 4.7 | 121.82 | 879.9 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 4 : sample 33.

Number of peaks found: 6 Corr. Area 1: 80.3
 Noise: 0.1

Peak table for sample 4 : sample 33.

| 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 | 120 | 136.83 | 1,733.2 | | 52.41 |
| 3 | 142 | 187.17 | 1,999.9 | | 54.43 |
| 4 | 152 | 1,248.89 | 12,488.1 | | 55.32 |
| 5 | 174 | 71.99 | 625.5 | | 57.39 |
| 6 | 189 | 225.90 | 1,808.0 | | 58.74 |
| 7 | 216 | 88.63 | 621.6 | | 61.14 |
| 8 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

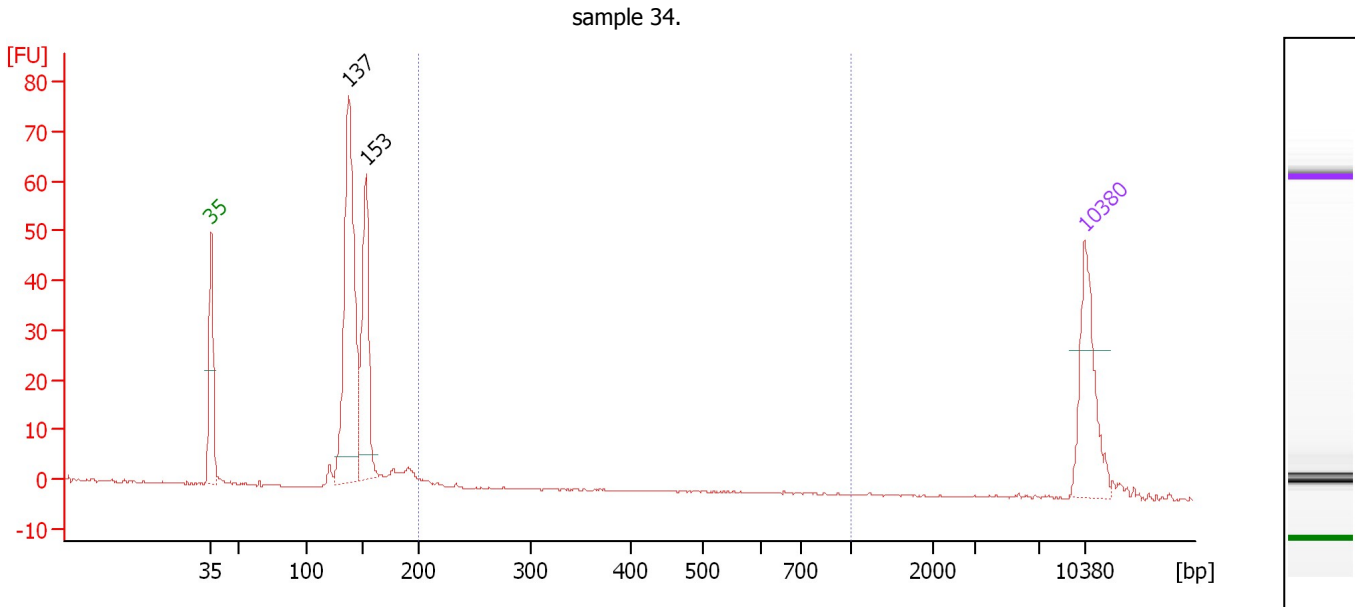
Region table for sample 4 : sample 33.

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 213 | 1,000 | 80.3 | 8 | 5.1 | 135.72 | 964.2 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 5 : sample 34.

Number of peaks found: 2 Corr. Area 1: 2.3
 Noise: 0.1

Peak table for sample 5 : sample 34.

| 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 | 137 | 338.44 | 3,734.2 | | 54.03 |
| 3 | 153 | 170.90 | 1,694.7 | | 55.44 |
| 4 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

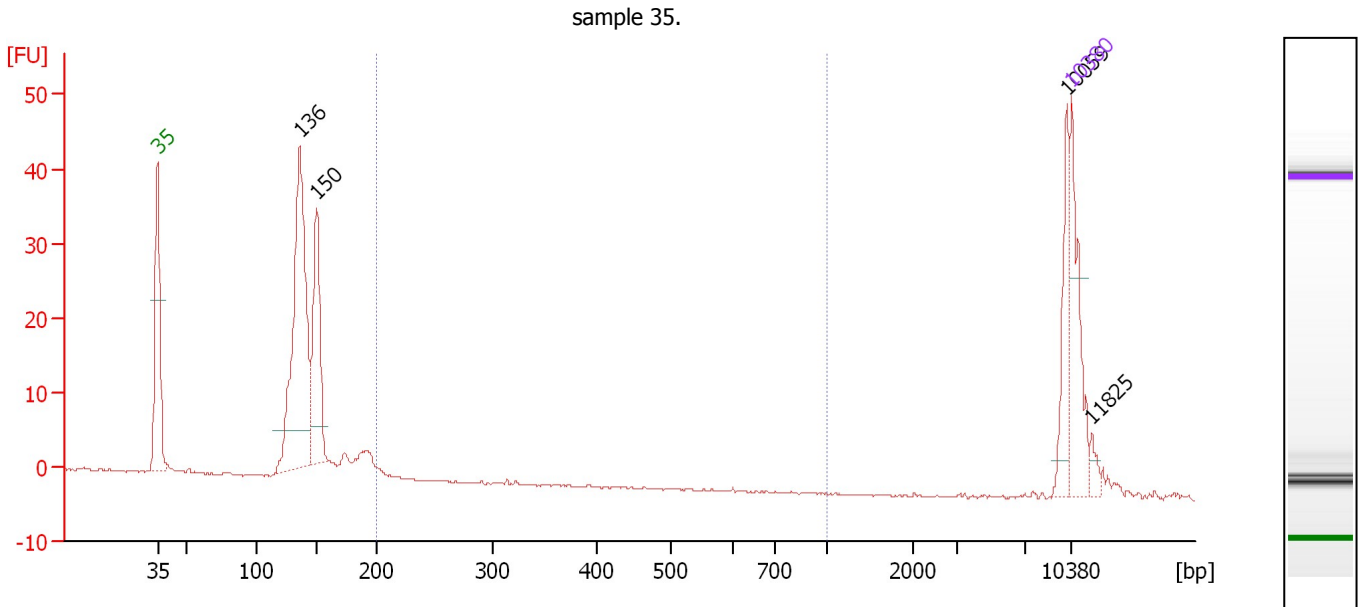
Region table for sample 5 : sample 34.

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 238 | 1,000 | 2.3 | 1 | 47.7 | 4.75 | 33.7 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 6 : sample 35.

Number of peaks found: 4 Corr. Area 1: 1.4
 Noise: 0.1

Peak table for sample 6 : sample 35.

| 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 | 136 | 366.65 | 4,089.2 | | 53.89 |
| 3 | 150 | 153.08 | 1,546.2 | | 55.18 |
| 4 | 10,059 | 52.36 | 7.9 | | 112.66 |
| 5 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 6 | 11,825 | 0.00 | 0.0 | | 114.55 |

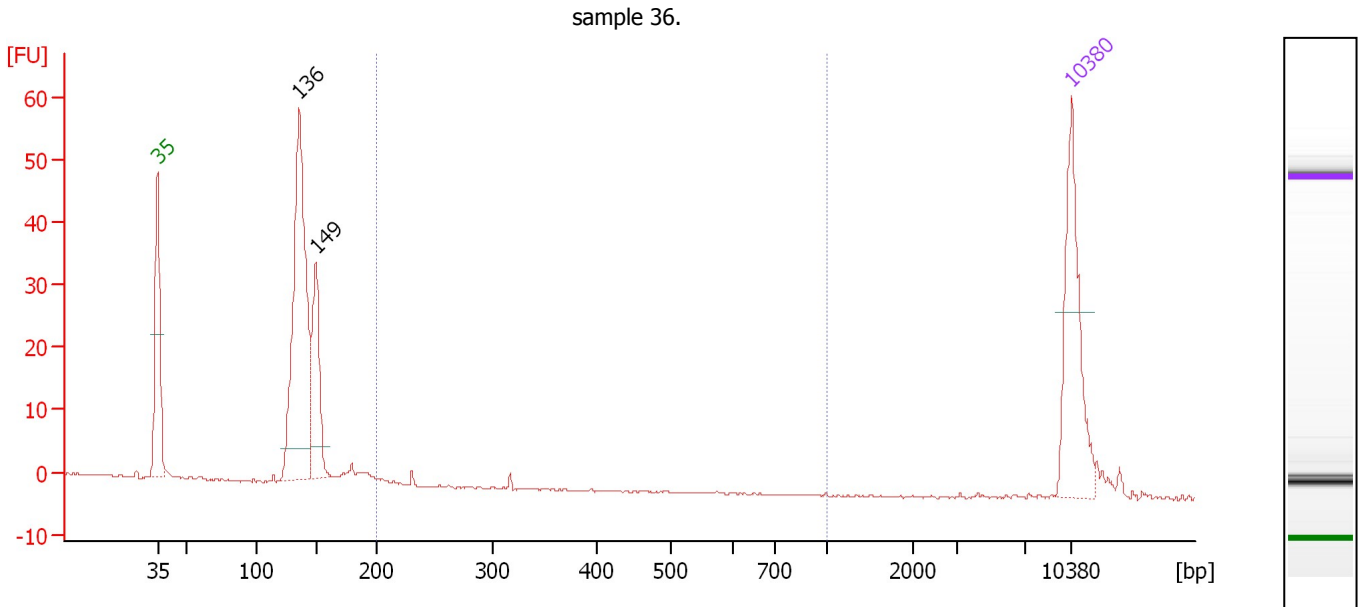
Region table for sample 6 : sample 35.

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 204 | 1,000 | 1.4 | 1 | 3.2 | 4.61 | 34.3 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 7 : sample 36.

Number of peaks found: 2 Corr. Area 1: 1.1
 Noise: 0.1

Peak table for sample 7 : sample 36.

| 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 | 136 | 265.37 | 2,967.3 | | 53.86 |
| 3 | 149 | 87.55 | 888.1 | | 55.12 |
| 4 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

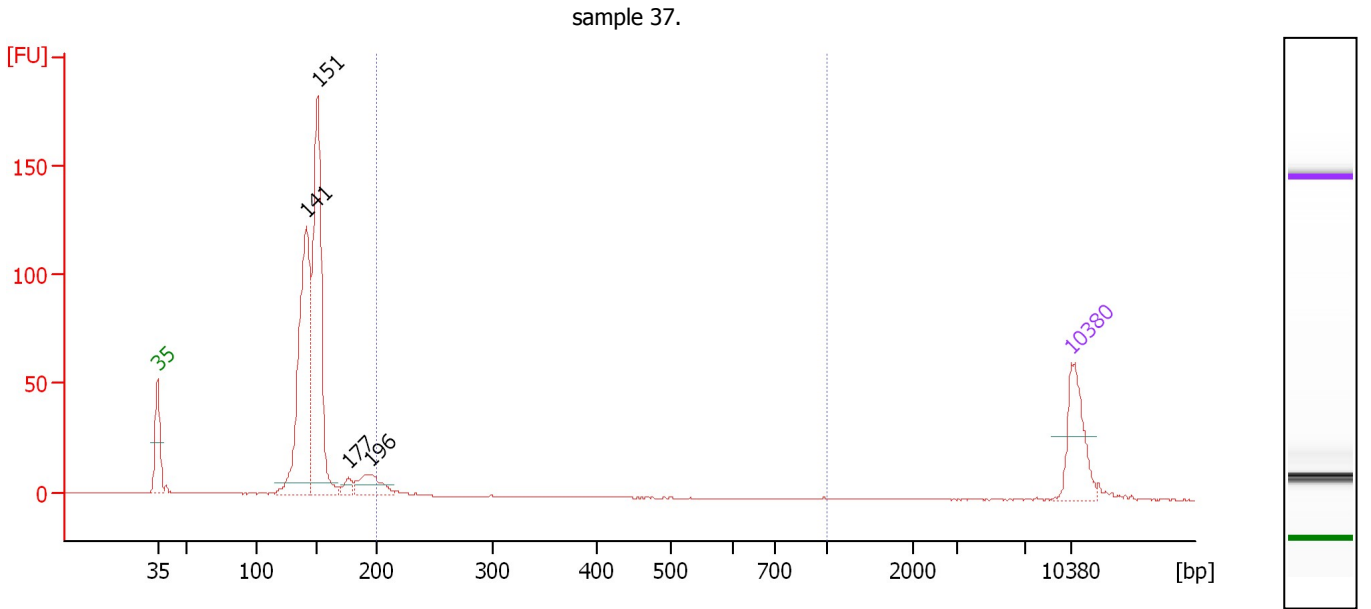
Region table for sample 7 : sample 36.

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 252 | 1,000 | 1.1 | 1 | 18.5 | 2.00 | 12.6 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 8 : sample 37.

Number of peaks found: 4 Corr. Area 1: 13.8
 Noise: 0.1

Peak table for sample 8 : sample 37.

| 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 | 141 | 446.65 | 4,792.7 | | 54.38 |
| 3 | 151 | 466.62 | 4,685.9 | | 55.26 |
| 4 | 177 | 18.04 | 154.2 | | 57.64 |
| 5 | 196 | 57.95 | 448.6 | | 59.32 |
| 6 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

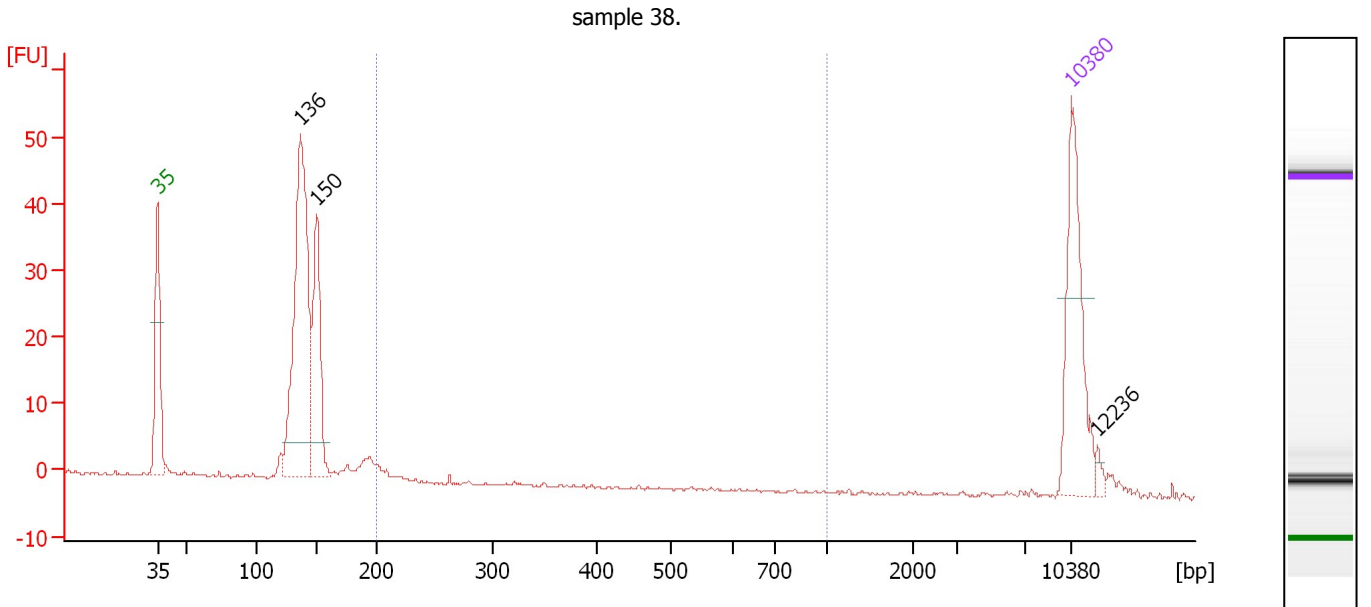
Region table for sample 8 : sample 37.

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 214 | 1,000 | 13.8 | 2 | 29.6 | 22.44 | 162.2 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 9 : sample 38.

Number of peaks found: 3 Corr. Area 1: 5.2
 Noise: 0.2

Peak table for sample 9 : sample 38.

| 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 | 136 | 254.47 | 2,829.5 | | 53.93 |
| 3 | 150 | 104.71 | 1,058.8 | | 55.17 |
| 4 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 5 | 12,236 | 0.00 | 0.0 | | 114.99 |

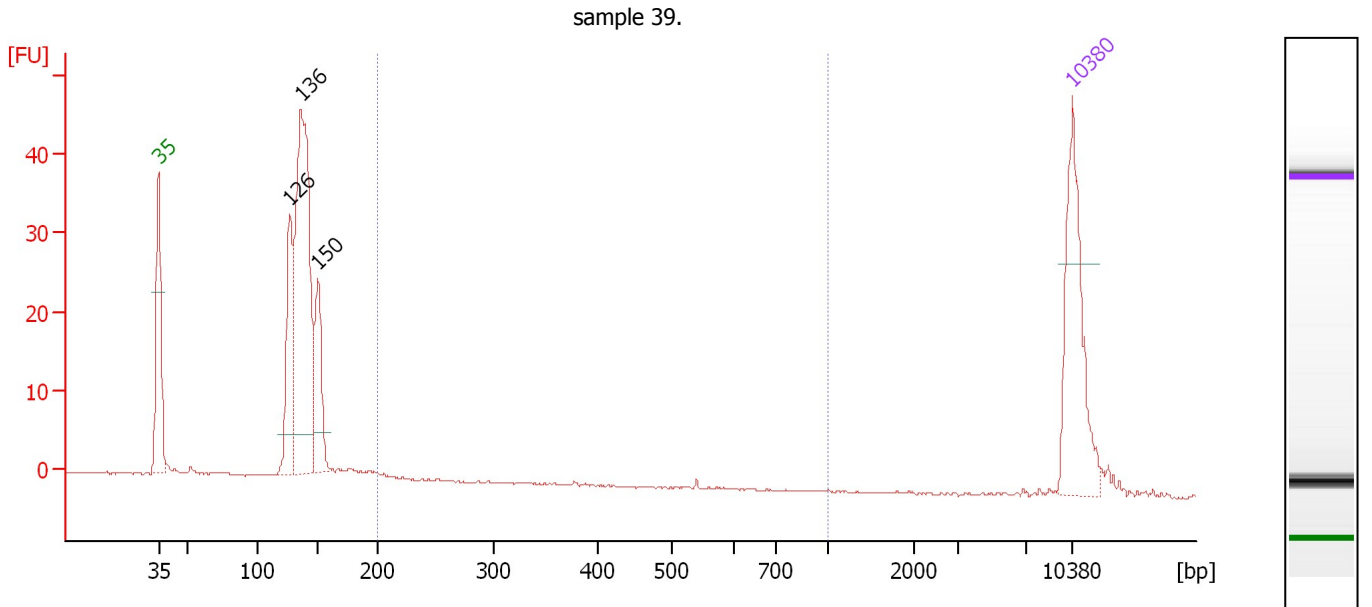
Region table for sample 9 : sample 38.

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 233 | 1,000 | 5.2 | 2 | 38.5 | 9.62 | 66.9 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 10 : sample 39.

Number of peaks found: 3 Corr. Area 1: 1.0
 Noise: 0.1

Peak table for sample 10 : sample 39.

| 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 | 126 | 88.63 | 1,062.8 | | 53.03 |
| 3 | 136 | 254.32 | 2,831.6 | | 53.91 |
| 4 | 150 | 65.44 | 660.2 | | 55.20 |
| 5 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

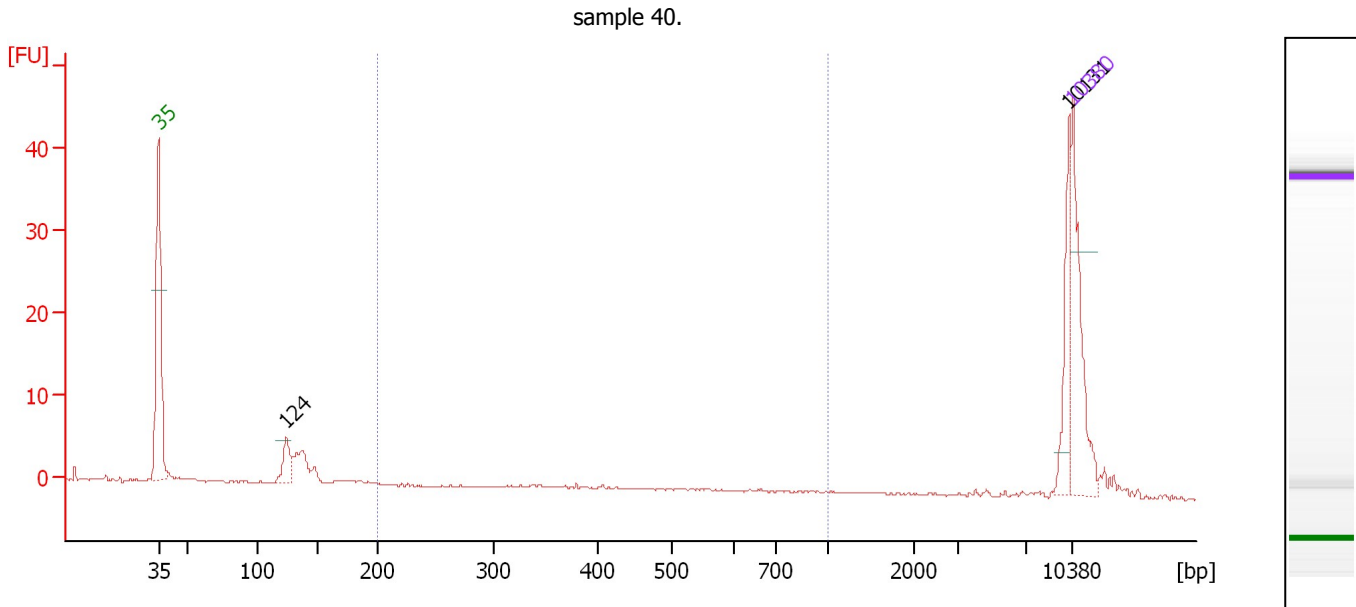
Region table for sample 10 : sample 39.

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 282 | 1,000 | 1.0 | 1 | 49.1 | 1.91 | 12.8 | Blue |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 PM

Electropherogram Summary Continued ...



Overall Results for sample 11 : sample 40.

Number of peaks found: 2 Corr. Area 1: 0.4
 Noise: 0.1

Peak table for sample 11 : sample 40.

| 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 | 124 | 28.11 | 343.9 | | 52.80 |
| 3 | 10,131 | 41.09 | 6.1 | | 112.73 |
| 4 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

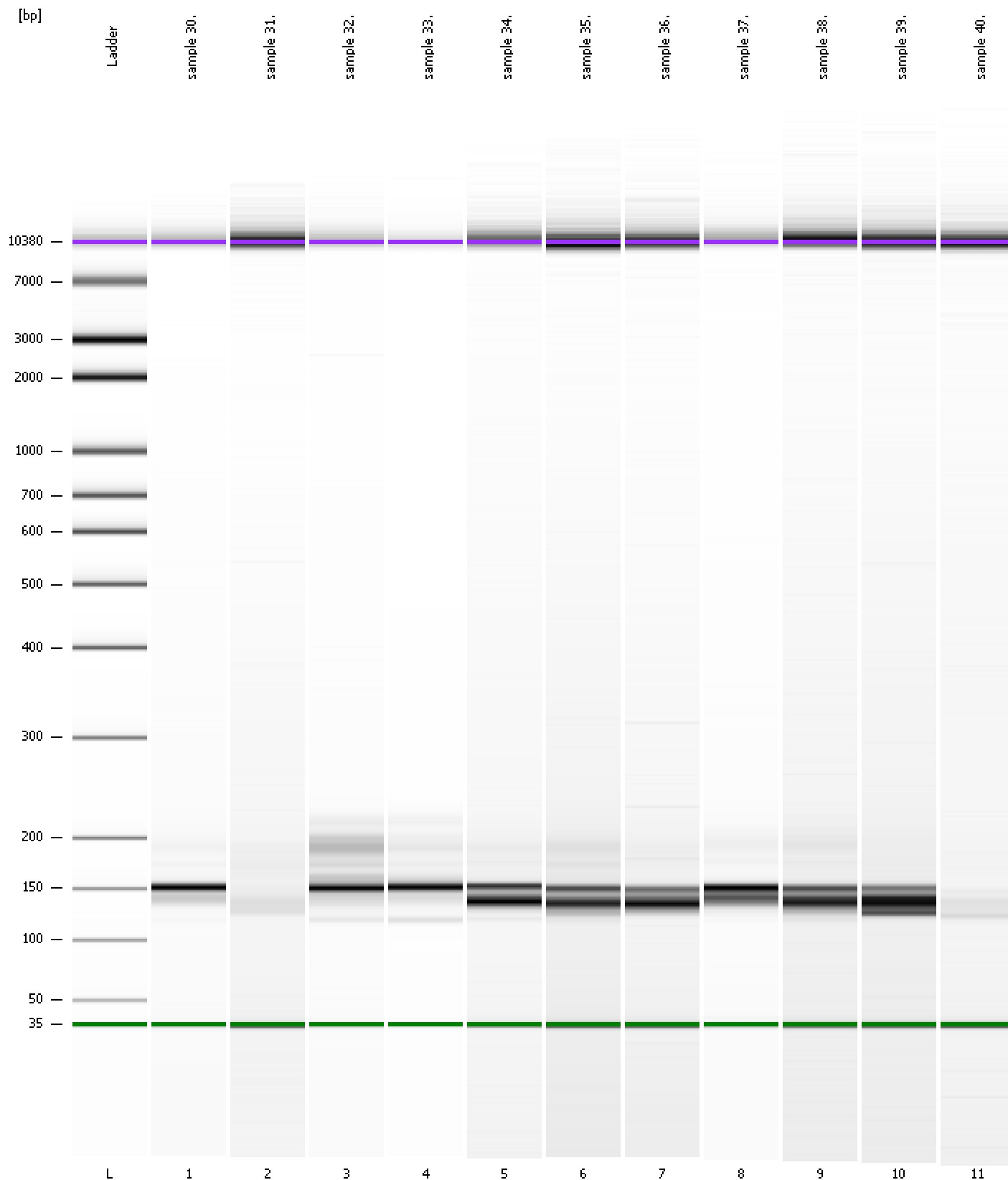
Region table for sample 11 : sample 40.

| From [bp] | Average Size [bp] | To [bp] | Corr. Area | % of Total | Size distribution in CV [%] | Conc. [pg/μl] | Molarity [pmol/l] | Color |
|-----------|-------------------|---------|------------|------------|-----------------------------|---------------|-------------------|-------|
| 200 | 486 | 1,000 | 0.4 | 1 | 38.0 | 1.26 | 4.5 | Blue |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
Modified: 2/10/2015 1:56:57 PM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\...ents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad

Created: 2/10/2015 1:16:27 PM
 Modified: 2/10/2015 1:56:57 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 | | 2/10/2015 1:56:54 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Run started on port 1 (File: C:\Documents and Settings\Bioanalyzer\2015-02-10\2015-02-10_002.xad) | | Instrument | Run | | 2/10/2015 1:16:32 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Product Number : G2938B | | Instrument | Run | | 2/10/2015 1:16:32 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Name : | | Instrument | Run | | 2/10/2015 1:16:32 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Vendor : Agilent Technologies | | Instrument | Run | | 2/10/2015 1:16:32 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Serial# : DE13701086 | | Instrument | Run | | 2/10/2015 1:16:32 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Firmware : C.01.069 | | Instrument | Run | | 2/10/2015 1:16:32 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Cartridge : Electrode | | Instrument | Run | | 2/10/2015 1:16:32 PM | (GMT --08:00) Pacific Standard Time | UC Davis | D8XSMGH1 |