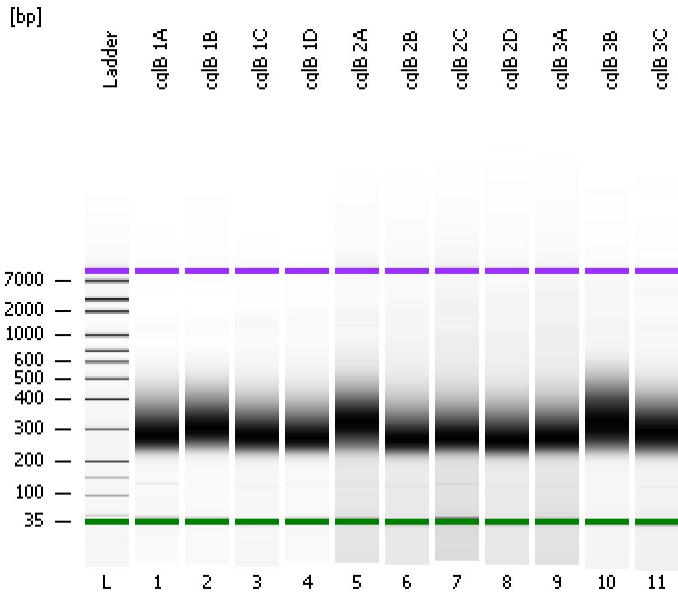


Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
Modified: 5/20/2016 10:09:22 AM

Electrophoresis File Run Summary



Instrument Information:

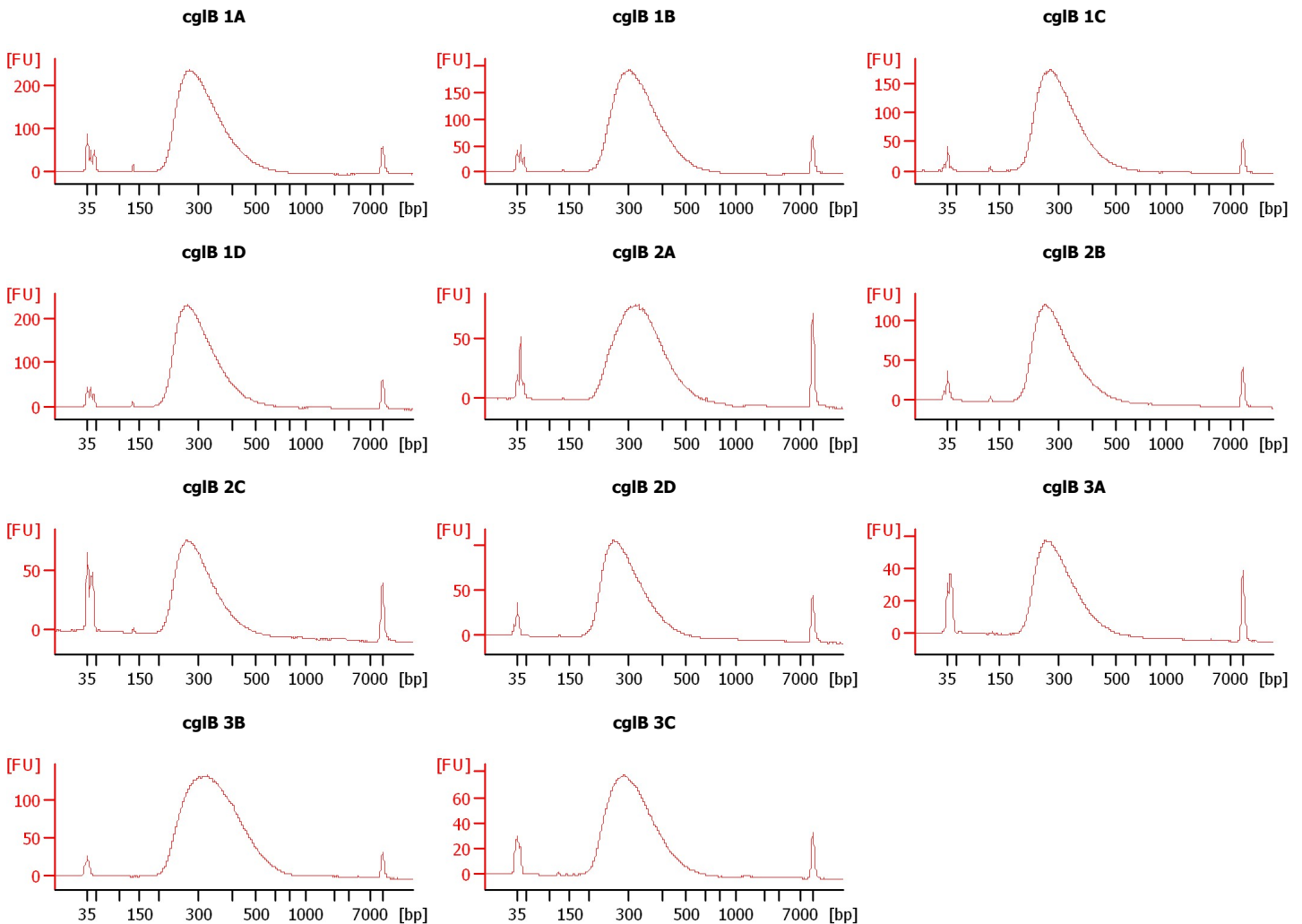
Instrument Name: DE13701086 Firmware: C.01.069
Serial#: DE13701086 Type: G2938B

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:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electrophoresis File Run Summary (Chip Summary)

| Sample Name | Sample Comment | Rest. Digest | Status | Observation | Result Label | Result Color |
|-------------|----------------|--------------------------|--------|-------------|--------------|--------------|
| cglB 1A | | <input type="checkbox"/> | ✓ | | | |
| cglB 1B | | <input type="checkbox"/> | ✓ | | | |
| cglB 1C | | <input type="checkbox"/> | ✓ | | | |
| cglB 1D | | <input type="checkbox"/> | ✓ | | | |
| cglB 2A | | <input type="checkbox"/> | ✓ | | | |
| cglB 2B | | <input type="checkbox"/> | ✓ | | | |
| cglB 2C | | <input type="checkbox"/> | ✓ | | | |
| cglB 2D | | <input type="checkbox"/> | ✓ | | | |
| cglB 3A | | <input type="checkbox"/> | ✓ | | | |
| cglB 3B | | <input type="checkbox"/> | ✓ | | | |
| cglB 3C | | <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\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
Modified: 5/20/2016 10:09:22 AM

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

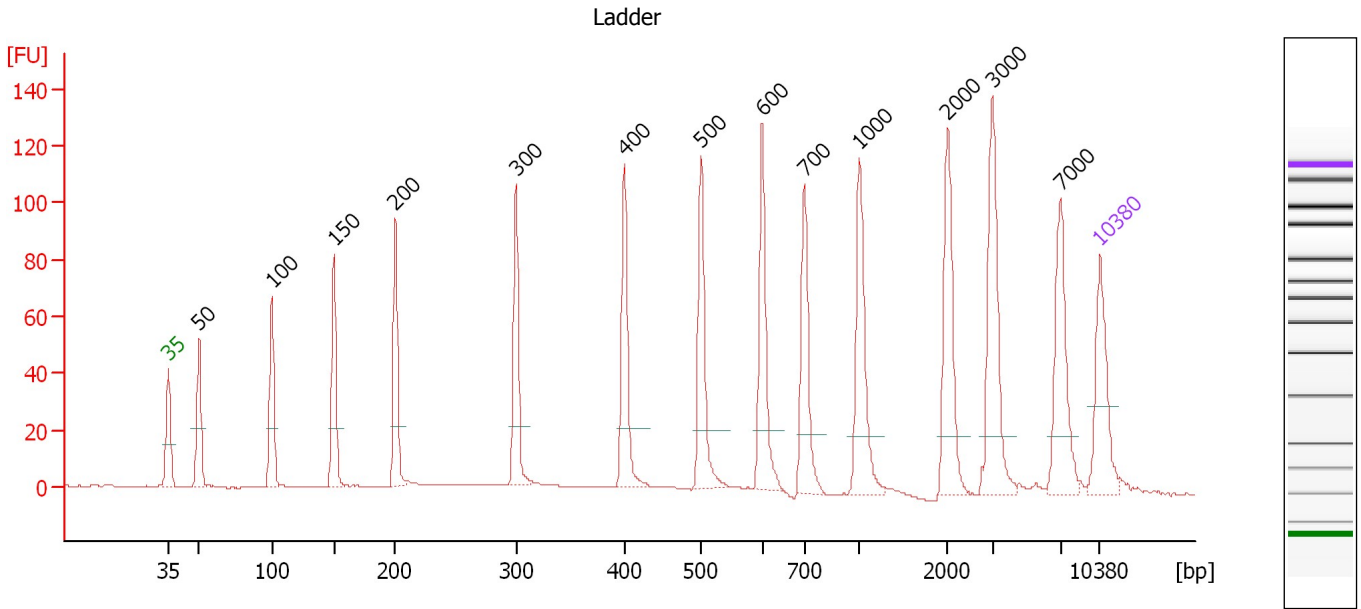
Ladder

| 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\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electropherogram Summary



Overall Results for Ladder

Noise: 0.2

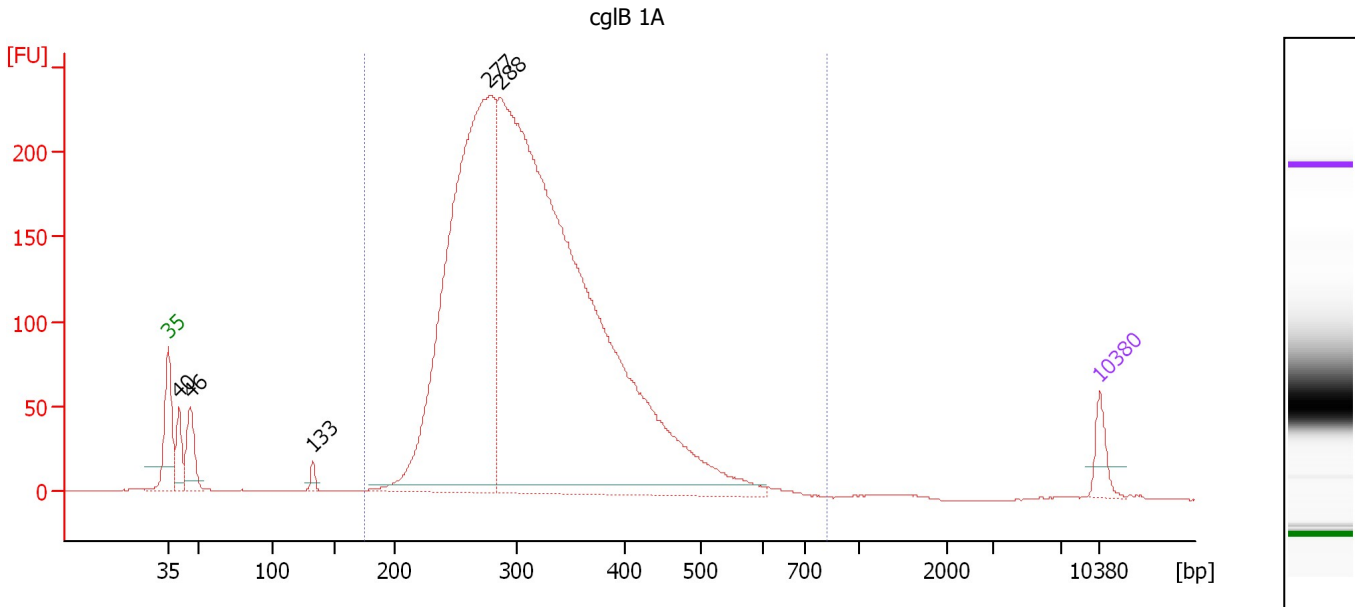
Peak table for Ladder

| 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 | 45.31 |
| 3 | 100 | 150.00 | 2,272.7 | Ladder Peak | 50.79 |
| 4 | 150 | 150.00 | 1,515.2 | Ladder Peak | 55.46 |
| 5 | 200 | 150.00 | 1,136.4 | Ladder Peak | 60.08 |
| 6 | 300 | 150.00 | 757.6 | Ladder Peak | 69.12 |
| 7 | 400 | 150.00 | 568.2 | Ladder Peak | 77.25 |
| 8 | 500 | 150.00 | 454.5 | Ladder Peak | 83.03 |
| 9 | 600 | 150.00 | 378.8 | Ladder Peak | 87.65 |
| 10 | 700 | 150.00 | 324.7 | Ladder Peak | 90.77 |
| 11 | 1,000 | 150.00 | 227.3 | Ladder Peak | 94.91 |
| 12 | 2,000 | 150.00 | 113.6 | Ladder Peak | 101.55 |
| 13 | 3,000 | 150.00 | 75.8 | Ladder Peak | 104.92 |
| 14 | 7,000 | 150.00 | 32.5 | Ladder Peak | 110.02 |
| 15 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electropherogram Summary Continued ...



Overall Results for sample 1 : cgIB 1A

Number of peaks found: 5 Corr. Area 1: 4,102.7
 Noise: 0.3

Peak table for sample 1 : cgIB 1A

| 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 | 40 | 163.37 | 6,140.9 | | 43.82 |
| 3 | 46 | 235.25 | 7,760.5 | | 44.68 |
| 4 | 133 | 37.99 | 433.1 | | 53.87 |
| 5 | 277 | 3,578.40 | 19,539.7 | | 67.09 |
| 6 | 288 | 6,437.72 | 33,918.4 | | 68.00 |
| 7 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

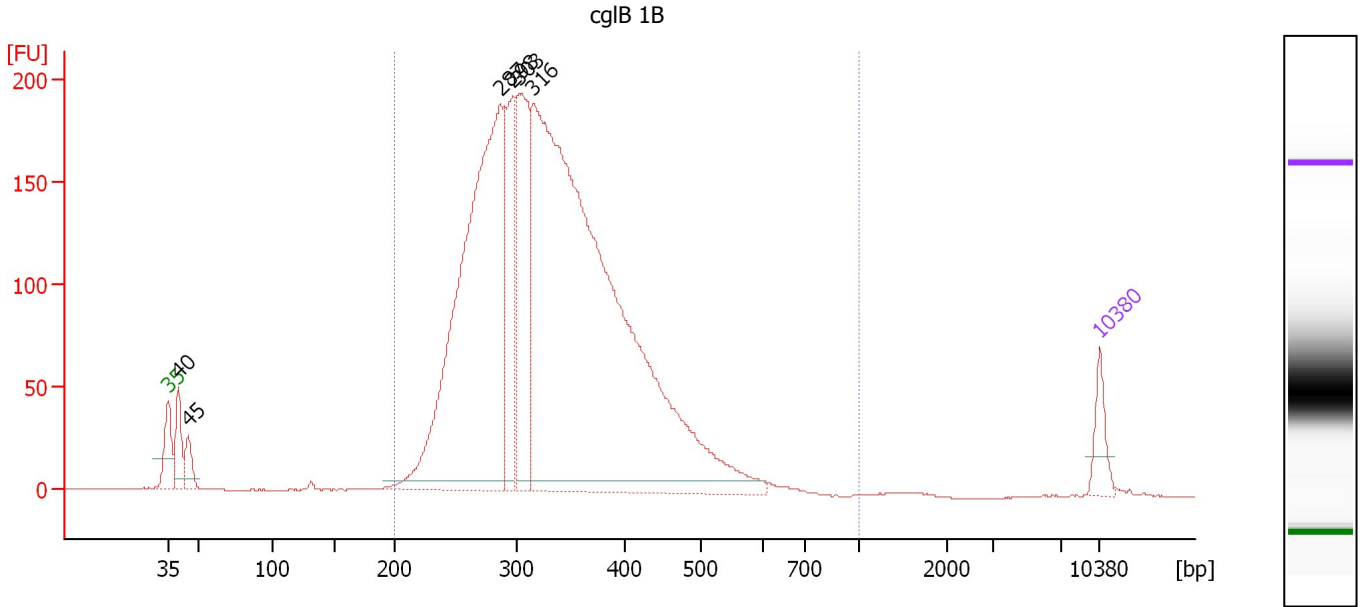
Region table for sample 1 : cgIB 1A

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Co Conc. [pg/μl] | % of Total | Size distribution in CV [%] |
|-----------|---------|-------------------|------------|-------------------|------------------|------------|-----------------------------|
| 175 | 829 | 321 | 4,102.7 | 47,487.7 | 9,501.33 | 95 | 22.0 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electropherogram Summary Continued ...



Overall Results for sample 2 : cgIB 1B

Number of peaks found: 6 Corr. Area 1: 3,461.4
 Noise: 0.2

Peak table for sample 2 : cgIB 1B

| 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 | 40 | 160.68 | 6,082.8 | | 43.77 |
| 3 | 45 | 109.52 | 3,709.3 | | 44.50 |
| 4 | 287 | 2,287.45 | 12,077.5 | | 67.94 |
| 5 | 298 | 470.72 | 2,396.1 | | 68.91 |
| 6 | 303 | 643.47 | 3,214.1 | | 69.40 |
| 7 | 316 | 4,165.83 | 19,985.4 | | 70.41 |
| 8 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

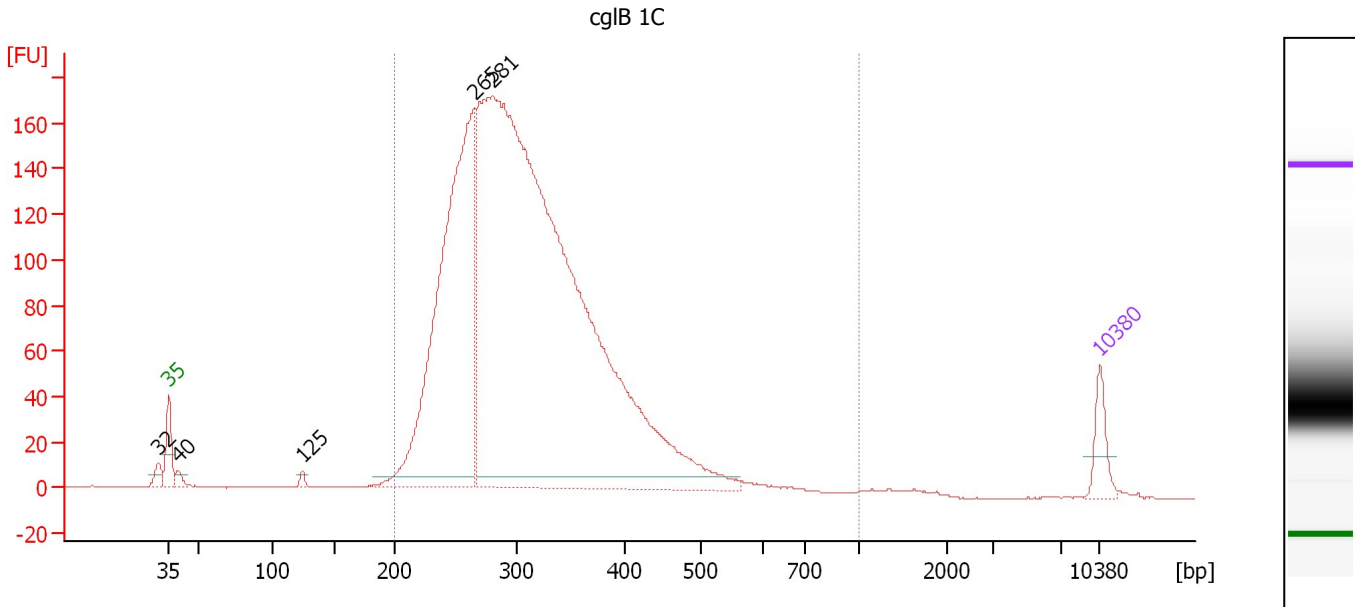
Region table for sample 2 : cgIB 1B

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Co Conc. lor [pg/μl] | % of Total | Size distribution in CV [%] |
|-----------|---------|-------------------|------------|-------------------|----------------------|------------|-----------------------------|
| 200 | 1,000 | 336 | 3,461.4 | 34,827.7 | 7,320.65 | 97 | 20.8 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electropherogram Summary Continued ...



Overall Results for sample 3 : cgIB 1C

Number of peaks found: 5 Corr. Area 1: 2,870.4
 Noise: 0.2

Peak table for sample 3 : cgIB 1C

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|------|-----------|---------------|-------------------|--------------|----------------------------|
| 1 | 32 | 0.00 | 0.0 | | 42.25 |
| 2 | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | 40 | 29.29 | 1,113.8 | | 43.75 |
| 4 | 125 | 16.11 | 195.8 | | 53.09 |
| 5 | 265 | 1,819.60 | 10,399.0 | | 65.97 |
| 6 | 281 | 5,309.87 | 28,680.9 | | 67.36 |
| 7 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

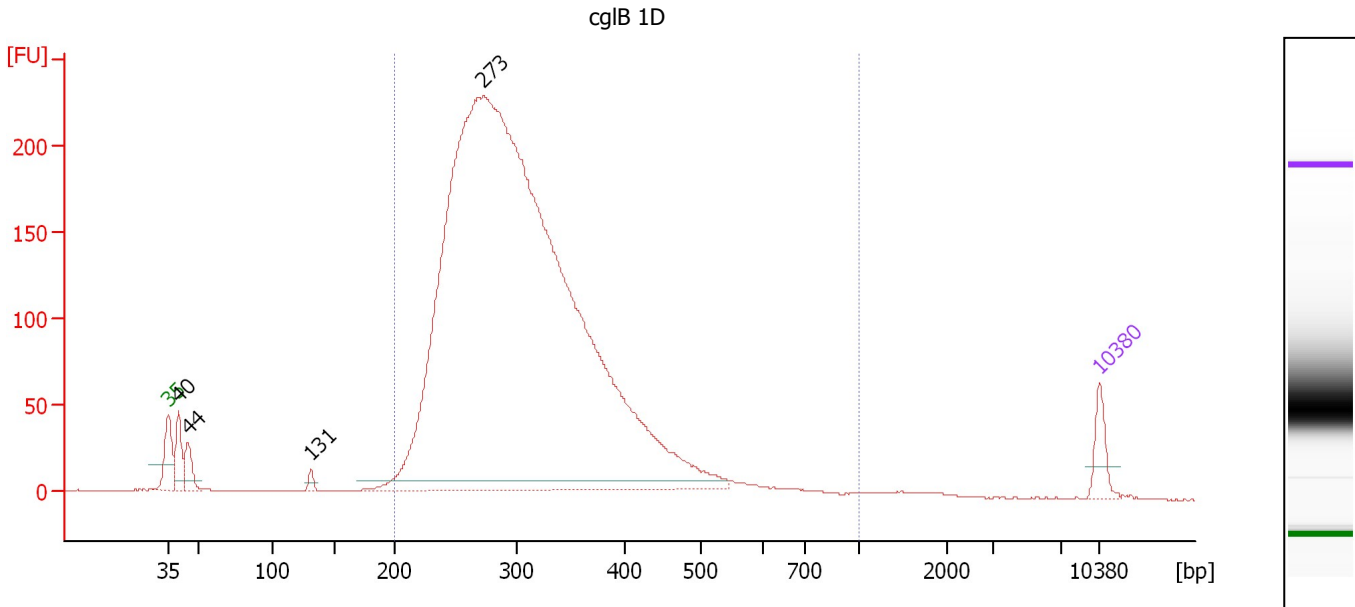
Region table for sample 3 : cgIB 1C

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Co Conc. [pg/μl] | % of Total | Size distribution in CV [%] |
|-----------|---------|-------------------|------------|-------------------|------------------|------------|-----------------------------|
| 200 | 1,000 | 316 | 2,870.4 | 34,942.7 | 6,891.16 | 97 | 23.3 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electropherogram Summary Continued ...



Overall Results for sample 4 : cgIB 1D

Number of peaks found: 4 Corr. Area 1: 3,725.9
 Noise: 0.2

Peak table for sample 4 : cgIB 1D

| 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 | 40 | 150.95 | 5,731.2 | | 43.76 |
| 3 | 44 | 116.64 | 3,972.7 | | 44.46 |
| 4 | 131 | 29.06 | 335.1 | | 53.73 |
| 5 | 273 | 8,375.26 | 46,506.4 | | 66.67 |
| 6 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

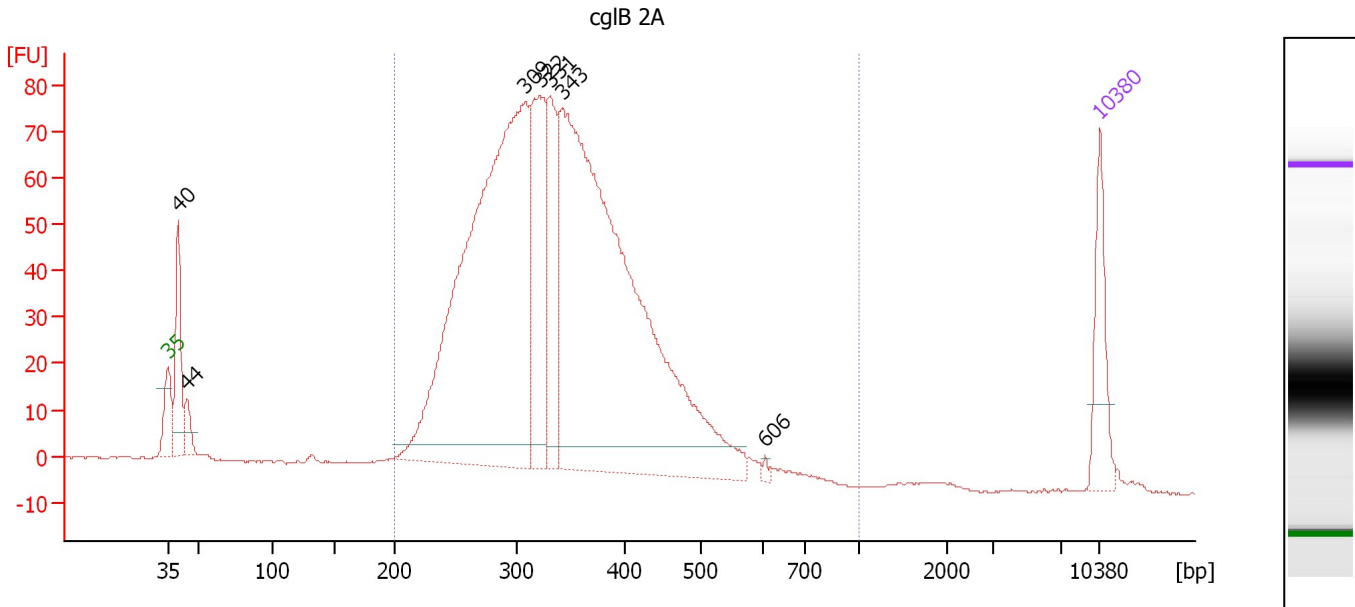
Region table for sample 4 : cgIB 1D

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Co Conc. lor [pg/μl] | % of Total | Size distribution in CV [%] |
|-----------|---------|-------------------|------------|-------------------|----------------------|------------|-----------------------------|
| 200 | 1,000 | 314 | 3,725.9 | 41,400.2 | 8,099.46 | 95 | 24.0 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : cgIB 2A

Number of peaks found: 7 Corr. Area 1: 1,549.0
 Noise: 0.2

Peak table for sample 5 : cgIB 2A

| 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 | 40 | 147.55 | 5,589.4 | | 43.77 |
| 3 | 44 | 36.42 | 1,245.0 | | 44.44 |
| 4 | 309 | 1,157.77 | 5,684.9 | | 69.82 |
| 5 | 322 | 251.06 | 1,179.7 | | 70.95 |
| 6 | 331 | 168.97 | 772.8 | | 71.67 |
| 7 | 343 | 1,304.22 | 5,767.4 | | 72.59 |
| 8 | 606 | 4.76 | 11.9 | | 87.82 |
| 9 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

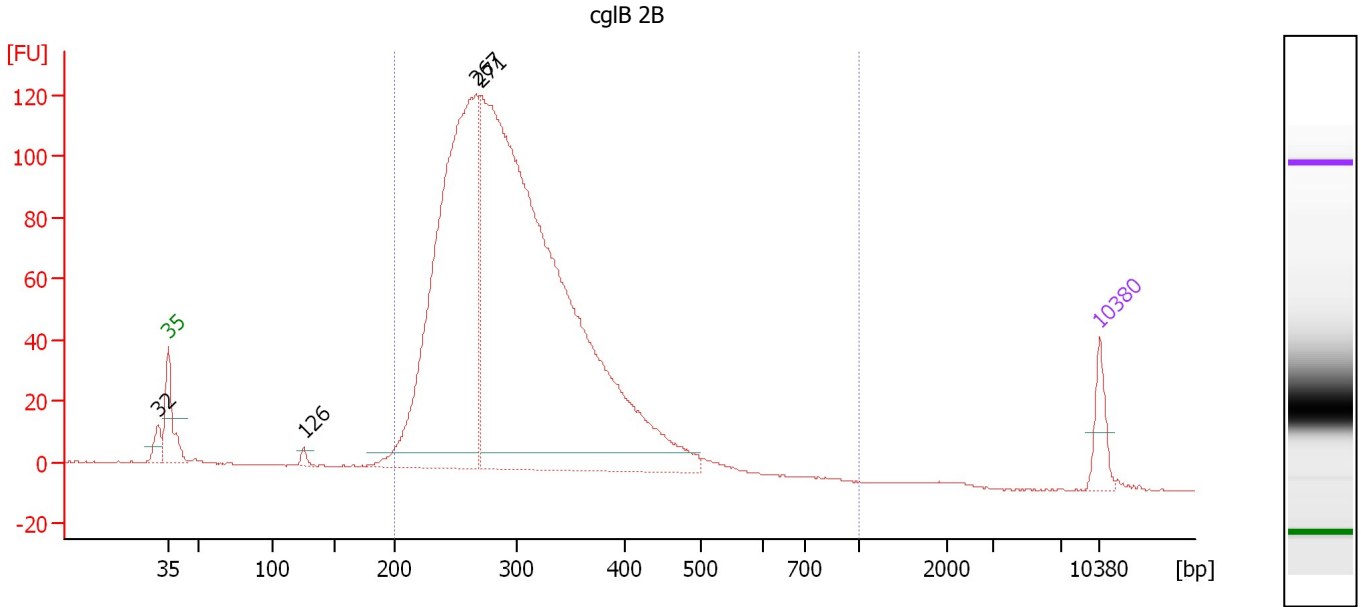
Region table for sample 5 : cgIB 2A

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Co Conc. [pg/μl] | % of Total | Size distribution in CV [%] |
|-----------|---------|-------------------|------------|-------------------|------------------|------------|-----------------------------|
| 200 | 1,000 | 343 | 1,549.0 | 13,721.9 | 2,932.57 | 94 | 21.7 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electropherogram Summary Continued ...



Overall Results for sample 6 : cgIB 2B

Number of peaks found: 4 Corr. Area 1: 1,927.3
 Noise: 0.2

Peak table for sample 6 : cgIB 2B

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|------|-----------|---------------|-------------------|--------------|----------------------------|
| 1 | 32 | 0.00 | 0.0 | | 42.21 |
| 2 | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | 126 | 21.69 | 261.2 | | 53.20 |
| 4 | 267 | 2,061.47 | 11,715.1 | | 66.10 |
| 5 | 271 | 3,960.06 | 22,116.7 | | 66.53 |
| 6 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

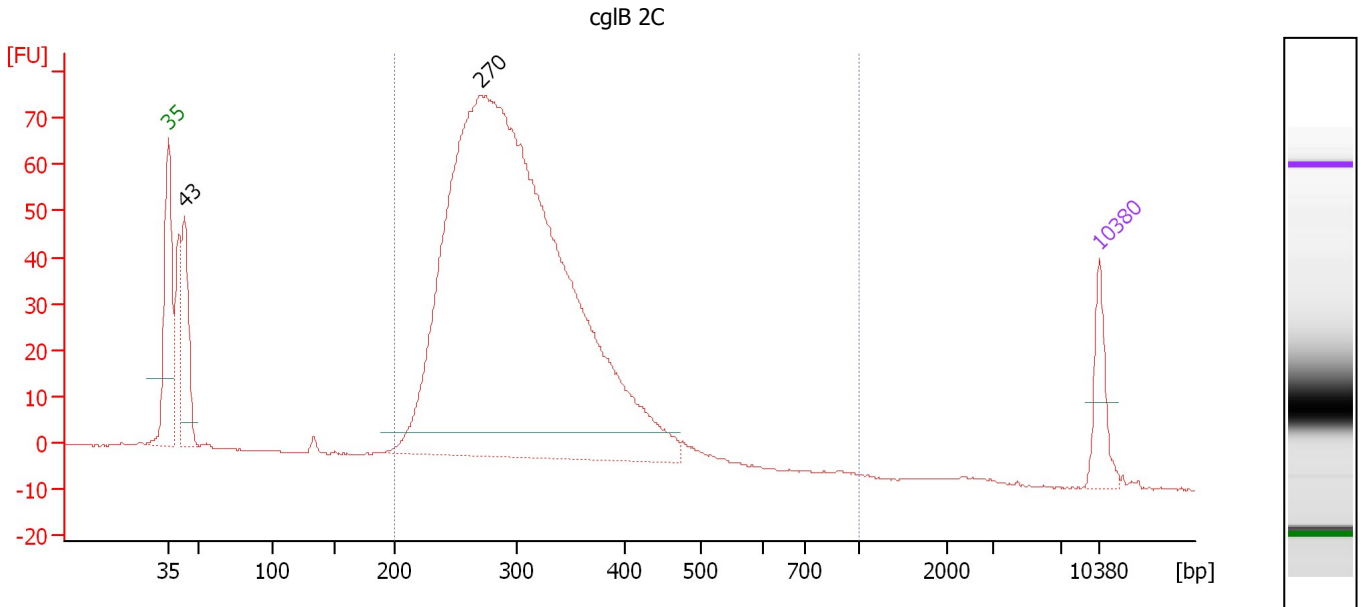
Region table for sample 6 : cgIB 2B

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Co Conc. lor [pg/μl] | % of Total | Size distribution in CV [%] |
|-----------|---------|-------------------|------------|-------------------|----------------------|------------|-----------------------------|
| 200 | 1,000 | 308 | 1,927.3 | 31,218.8 | 6,000.22 | 96 | 23.7 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electropherogram Summary Continued ...



Overall Results for sample 7 : cgIB 2C

Number of peaks found: 2 Corr. Area 1: 1,191.8
 Noise: 0.2

Peak table for sample 7 : cgIB 2C

| 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 | 43 | 258.67 | 9,120.6 | | 44.23 |
| 3 | 270 | 3,572.48 | 20,037.8 | | 66.42 |
| 4 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

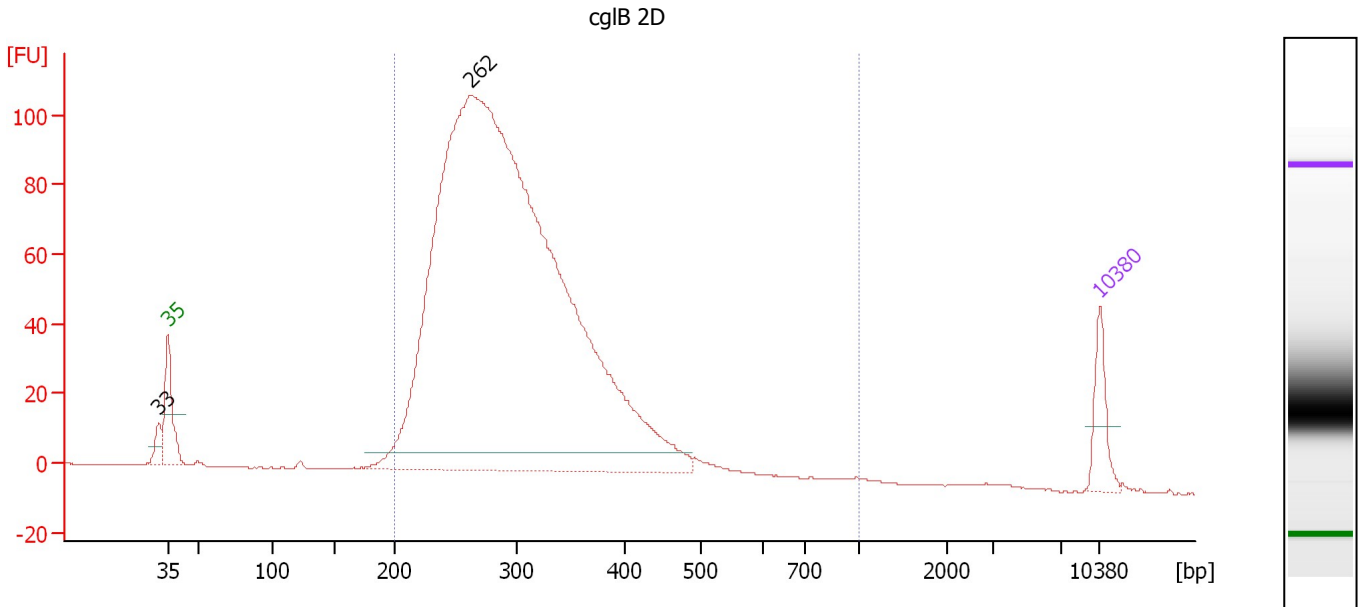
Region table for sample 7 : cgIB 2C

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Co Conc. lor [pg/μl] | % of Total | Size distribution in CV [%] |
|-----------|---------|-------------------|------------|-------------------|----------------------|------------|-----------------------------|
| 200 | 1,000 | 308 | 1,191.8 | 17,664.6 | 3,436.44 | 90 | 20.2 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electropherogram Summary Continued ...



Overall Results for sample 8 : cgIB 2D

Number of peaks found: 2 Corr. Area 1: 1,685.0
 Noise: 0.1

Peak table for sample 8 : cgIB 2D

| Peak | Size [bp] | Conc. [pg/μl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|------|-----------|---------------|-------------------|--------------|----------------------------|
| 1 | 33 | 0.00 | 0.0 | | 42.29 |
| 2 | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | 262 | 4,738.88 | 27,355.7 | | 65.73 |
| 4 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

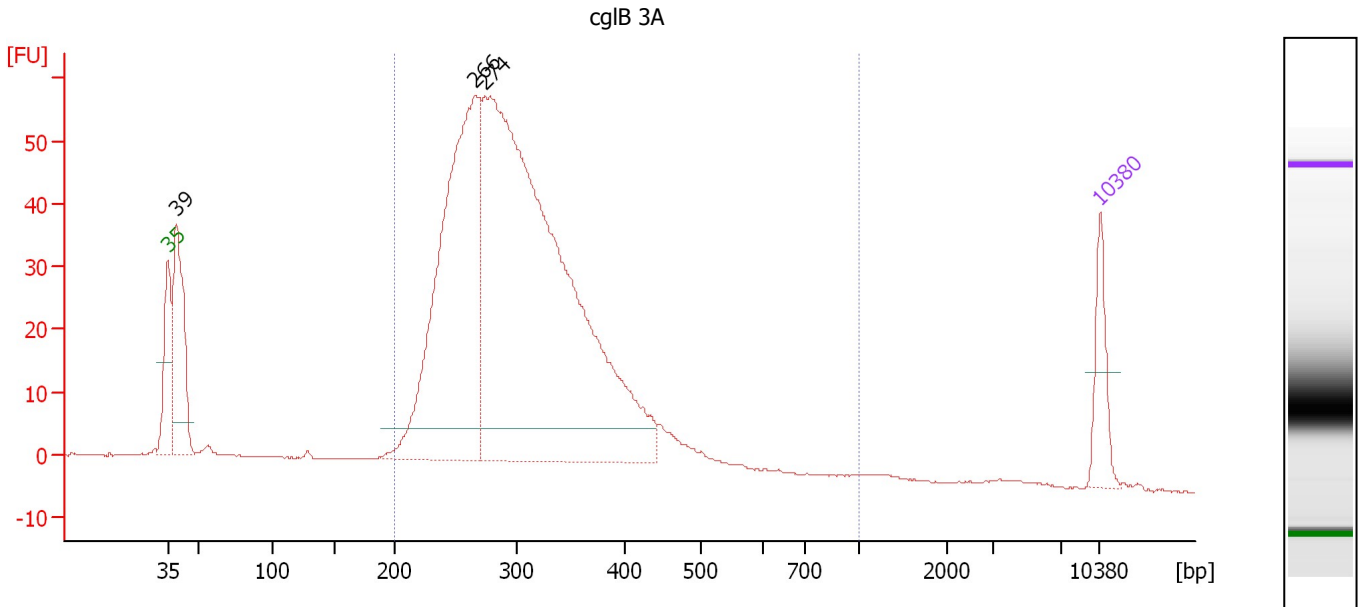
Region table for sample 8 : cgIB 2D

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Co Conc. lor [pg/μl] | % of Total | Size distribution in CV [%] |
|-----------|---------|-------------------|------------|-------------------|----------------------|------------|-----------------------------|
| 200 | 1,000 | 306 | 1,685.0 | 24,000.5 | 4,581.62 | 97 | 24.4 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electropherogram Summary Continued ...



Overall Results for sample 9 : cgIB 3A

Number of peaks found: 3 Corr. Area 1: 918.6
 Noise: 0.1

Peak table for sample 9 : cgIB 3A

| 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 | 39 | 337.66 | 13,148.4 | | 43.60 |
| 3 | 266 | 1,021.43 | 5,819.8 | | 66.04 |
| 4 | 274 | 2,068.99 | 11,449.9 | | 66.75 |
| 5 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

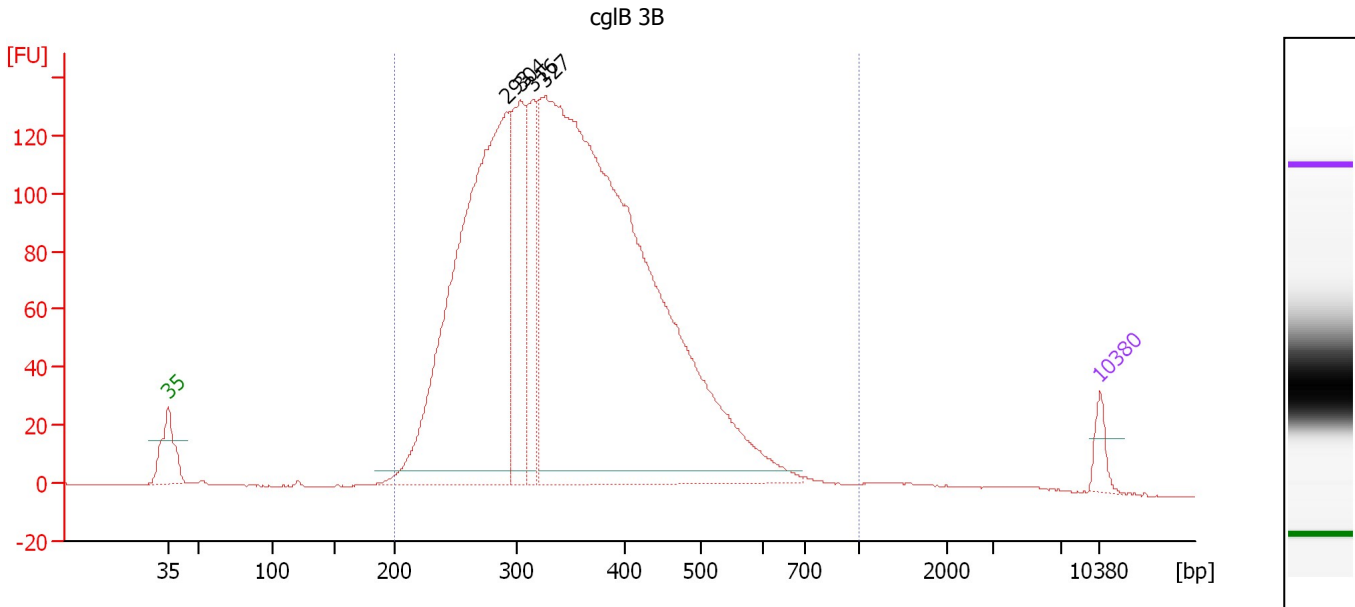
Region table for sample 9 : cgIB 3A

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Co Conc. lor [pg/μl] | % of Total | Size distribution in CV [%] |
|-----------|---------|-------------------|------------|-------------------|----------------------|------------|-----------------------------|
| 200 | 1,000 | 312 | 918.6 | 16,197.5 | 3,156.66 | 89 | 23.7 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electropherogram Summary Continued ...



Overall Results for sample 10 : cgIB 3B

Number of peaks found: 4 Corr. Area 1: 2,843.7
 Noise: 0.2

Peak table for sample 10 : cgIB 3B

| 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 | 293 | 3,613.39 | 18,710.9 | | 68.45 |
| 3 | 304 | 939.00 | 4,679.7 | | 69.45 |
| 4 | 316 | 587.95 | 2,822.7 | | 70.39 |
| 5 | 327 | 7,072.23 | 32,753.0 | | 71.33 |
| 6 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

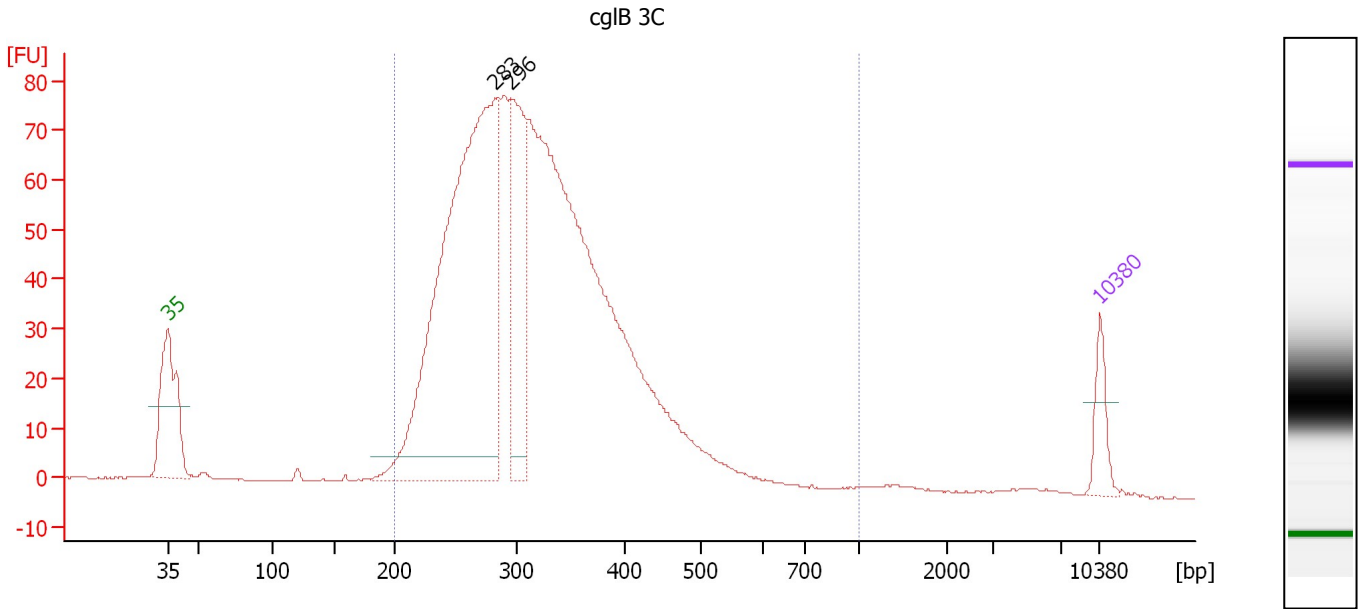
Region table for sample 10 : cgIB 3B

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Co Conc. lor [pg/μl] | % of Total | Size distribution in CV [%] |
|-----------|---------|-------------------|------------|-------------------|----------------------|------------|-----------------------------|
| 200 | 1,000 | 357 | 2,843.7 | 55,409.1 | 12,061.80 | 98 | 26.3 |

Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

Electropherogram Summary Continued ...



Overall Results for sample 11 : cgIB 3C

Number of peaks found: 2 Corr. Area 1: 1,376.0
 Noise: 0.1

Peak table for sample 11 : cgIB 3C

| 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 | 283 | 2,154.11 | 11,534.4 | | 67.58 |
| 3 | 296 | 526.09 | 2,695.1 | | 68.74 |
| 4 | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

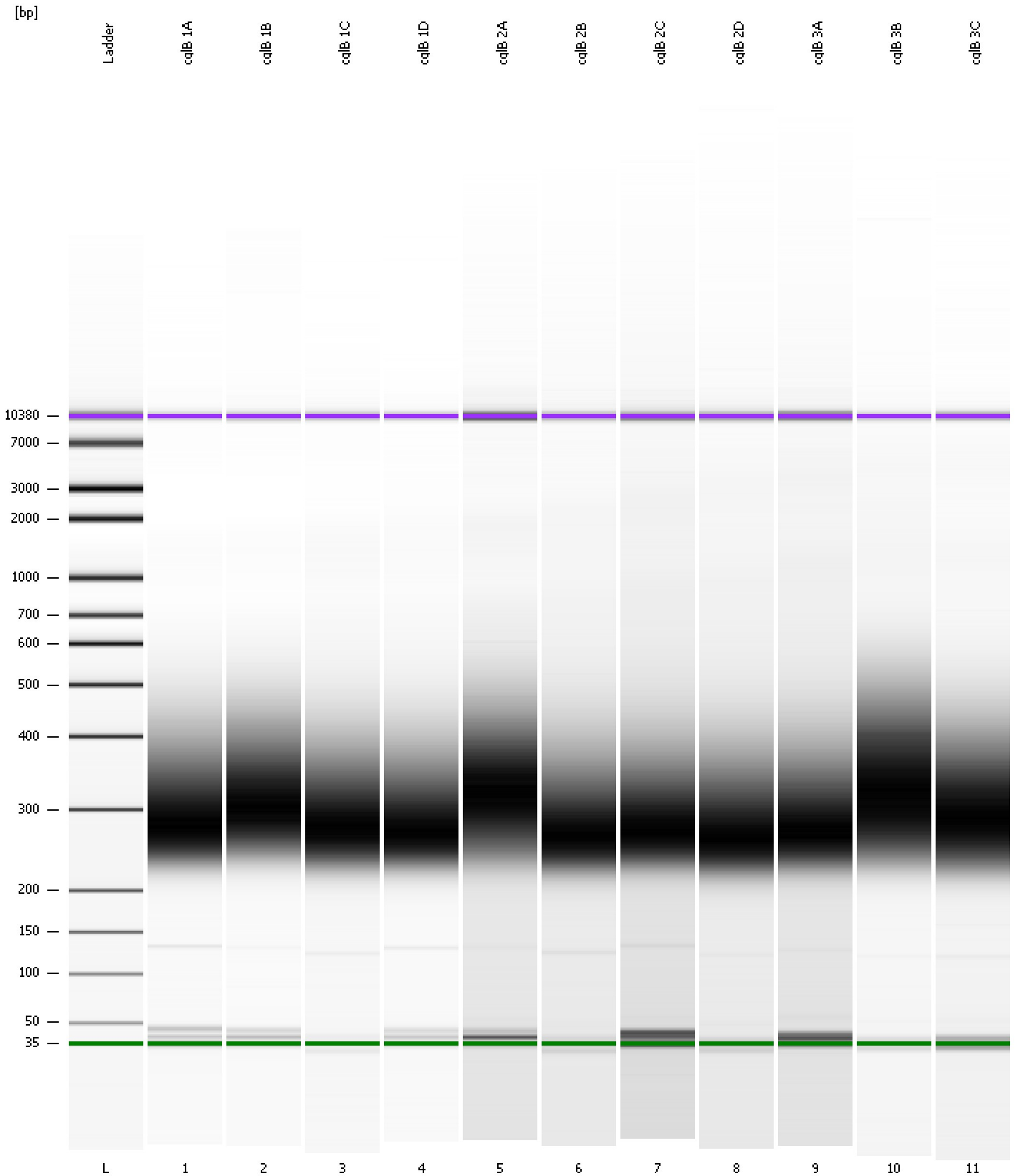
Region table for sample 11 : cgIB 3C

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Co Conc. lor [pg/μl] | % of Total | Size distribution in CV [%] |
|-----------|---------|-------------------|------------|-------------------|----------------------|------------|-----------------------------|
| 200 | 1,000 | 322 | 1,376.0 | 28,495.2 | 5,720.36 | 98 | 22.5 |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
Modified: 5/20/2016 10:09:22 AM

Gel Image



Assay Class: High Sensitivity DNA Assay
 Data Path: C:\... bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad

Created: 5/20/2016 9:26:23 AM
 Modified: 5/20/2016 10:09:22 AM

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 | | 5/20/2016 10:07:43 AM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Run started on port 1 (File: C:\Program Files\Agilent\2100 bioanalyzer\2100 expert\data\2016-05-20\2016-05-20_001.xad) | | Instrument | Run | | 5/20/2016 9:26:28 AM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Product Number : G2938B | | Instrument | Run | | 5/20/2016 9:26:28 AM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Name : | | Instrument | Run | | 5/20/2016 9:26:28 AM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Vendor : Agilent Technologies | | Instrument | Run | | 5/20/2016 9:26:28 AM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Serial# : DE13701086 | | Instrument | Run | | 5/20/2016 9:26:28 AM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Firmware : C.01.069 | | Instrument | Run | | 5/20/2016 9:26:28 AM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |
| Cartridge : Electrode | | Instrument | Run | | 5/20/2016 9:26:28 AM | (GMT --07:00) Pacific Standard Time | UC Davis | D8XSMGH1 |