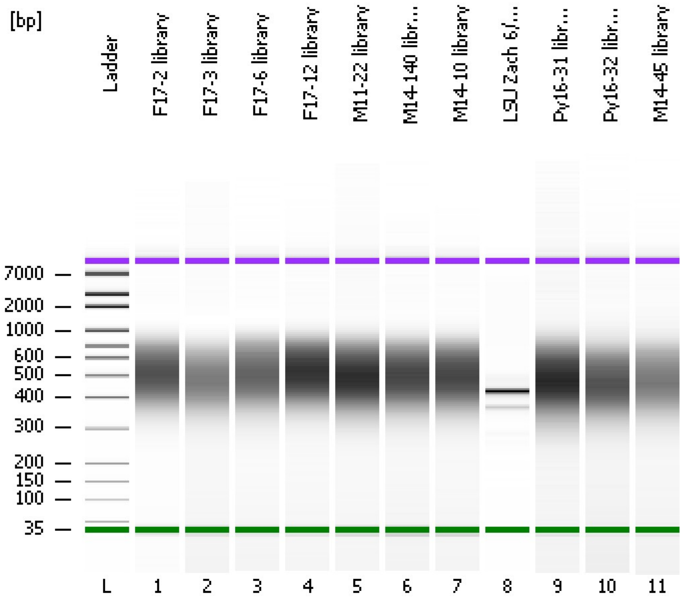
| | | | |
|---|---|---|---|
| Assay Class: | High Sensitivity DNA Assay | Created: | 6/20/2017 10:12:23 AM |
| Data Path: | C:\...ibraries9-15_HiSeq995-Libraries12-14_LSUZach6-9-Library.xad | Modified: | 6/20/2017 11:03:49 AM |

## 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:

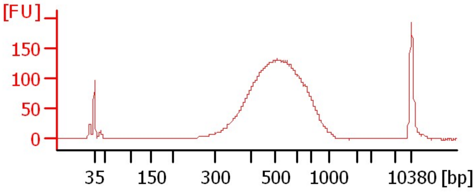
Chip Lot #:

Reagent Kit Lot #:

Chip Comments:

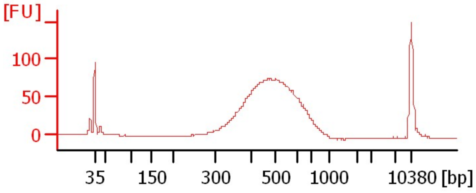
**F17-2 library**

HiSeq_992



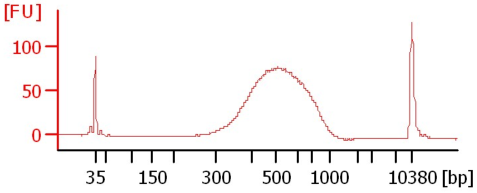
**F17-3 library**

HiSeq_992



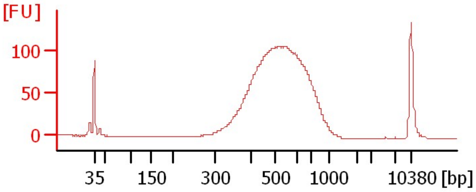
**F17-6 library**

HiSeq_992



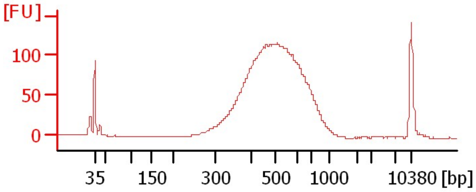
**F17-12 library**

HiSeq_992



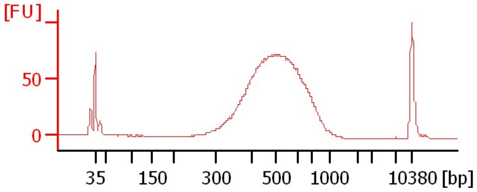
**M11-22 library**

HiSeq_992



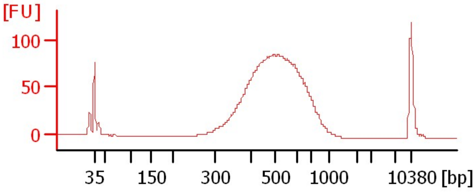
**M14-140 library**

HiSeq_992

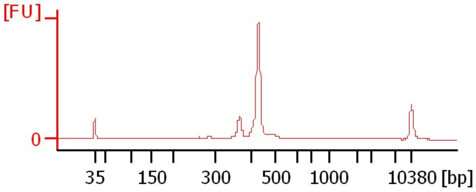


**M14-10 library**

HiSeq_992

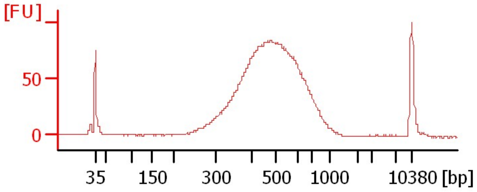


**LSU Zach 6/9 libary**



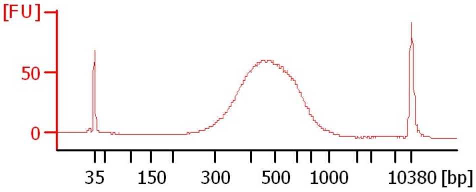
**Py16-31 library**

HiSeq_995



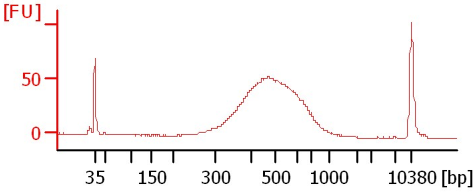
**Py16-32 library**

HiSeq_995



**M14-45 library**

HiSeq_995



| | | | |
|---|---|---|---|
| Assay Class: | High Sensitivity DNA Assay | Created: | 6/20/2017 10:12:23 AM |
| Data Path: | C:\...ibraries9-15_HiSeq995-Libraries12-14_LSUZach6-9-Library.xad | Modified: | 6/20/2017 11:03:49 AM |

## Electrophoresis File Run Summary (Chip Summary)

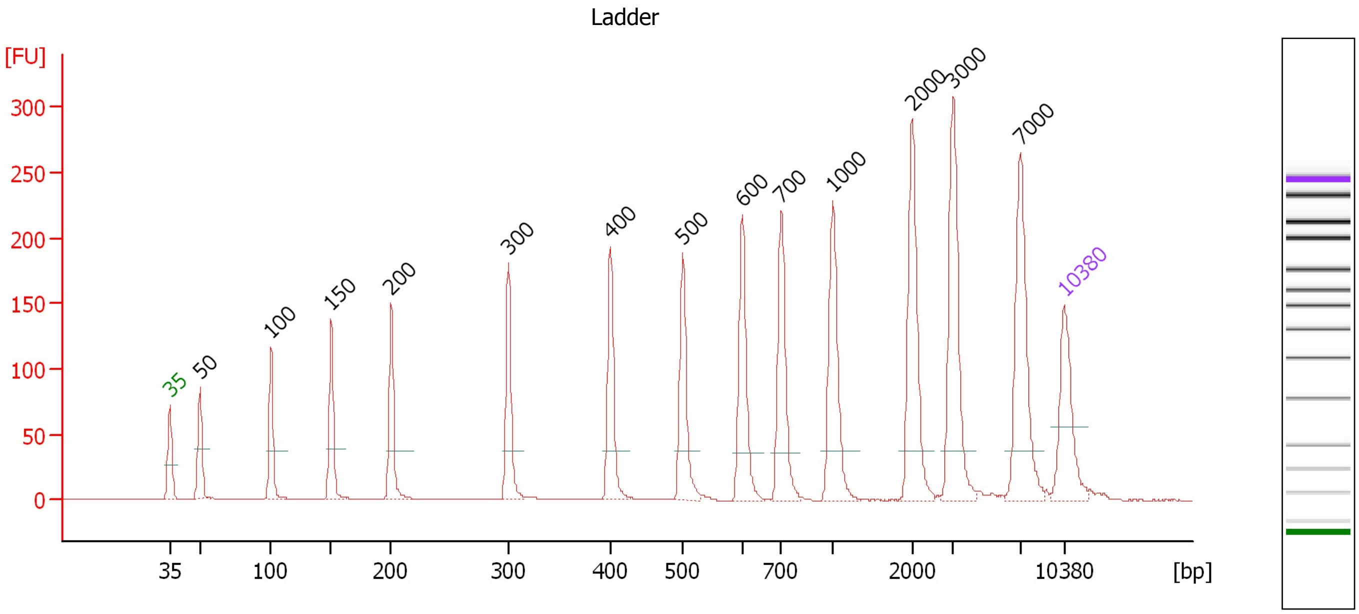
| Sample Name | Sample Comment | Rest. Digest | Status | Observation | Result Label | Result Color |
|---|---|---|---|---|---|---|
| F17-2 library | HiSeq_992 | ☐ | ✓ | | | |
| F17-3 library | HiSeq_992 | ☐ | ✓ | | | |
| F17-6 library | HiSeq_992 | ☐ | ✓ | | | |
| F17-12 library | HiSeq_992 | ☐ | ✓ | | | |
| M11-22 library | HiSeq_992 | ☐ | ✓ | | | |
| M14-140 library | HiSeq_992 | ☐ | ✓ | | | |
| M14-10 library | HiSeq_992 | ☐ | ✓ | | | |
| LSU Zach 6/9 libary | | ☐ | ✓ | | | |
| Py16-31 library | HiSeq_995 | ☐ | ✓ | | | |
| Py16-32 library | HiSeq_995 | ☐ | ✓ | | | |
| M14-45 library | HiSeq_995 | ☐ | ✓ | | | |
| Ladder | | ☐ | ✓ | | | |

**Chip Lot #** **Reagent Kit Lot #**

**Chip Comments :**

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ibraries9-15_HiSeq995-Libraries12-14_LSUZach6-9-Library.xad
Created: 6/20/2017 10:12:23 AM
Modified: 6/20/2017 11:03:49 AM

## Electropherogram Summary



### Overall Results for Ladder

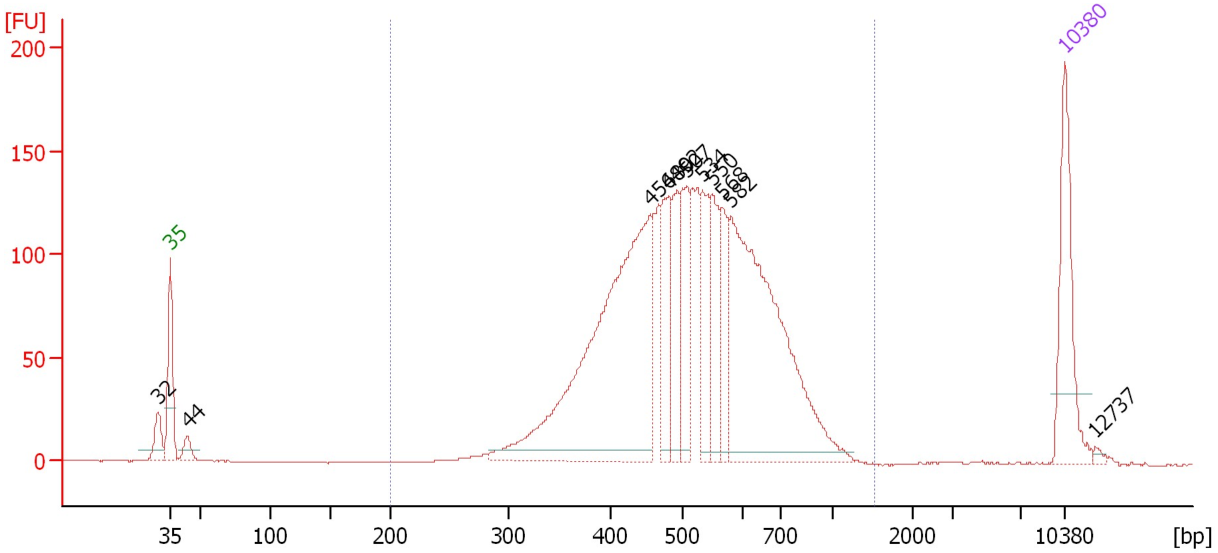
Noise: 0.2

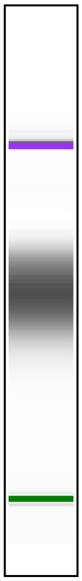
### Peak table for Ladder

| 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 | L | 50 | 150.00 | 4,545.5 | Ladder Peak | 45.34 |
| 3 | L | 100 | 150.00 | 2,272.7 | Ladder Peak | 50.90 |
| 4 | L | 150 | 150.00 | 1,515.2 | Ladder Peak | 55.62 |
| 5 | L | 200 | 150.00 | 1,136.4 | Ladder Peak | 60.30 |
| 6 | L | 300 | 150.00 | 757.6 | Ladder Peak | 69.49 |
| 7 | L | 400 | 150.00 | 568.2 | Ladder Peak | 77.47 |
| 8 | L | 500 | 150.00 | 454.5 | Ladder Peak | 83.12 |
| 9 | L | 600 | 150.00 | 378.8 | Ladder Peak | 87.80 |
| 10 | L | 700 | 150.00 | 324.7 | Ladder Peak | 90.83 |
| 11 | L | 1,000 | 150.00 | 227.3 | Ladder Peak | 94.87 |
| 12 | L | 2,000 | 150.00 | 113.6 | Ladder Peak | 101.07 |
| 13 | L | 3,000 | 150.00 | 75.8 | Ladder Peak | 104.28 |
| 14 | L | 7,000 | 150.00 | 32.5 | Ladder Peak | 109.51 |
| 15 | ▶ | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

**Electropherogram Summary Continued ...**

F17-2 library [ HiSeq_992 ]





## Overall Results for sample 1 : F17-2 library

| | | | |
|---|---|---|---|
| Number of peaks found: | 11 | Corr. Area 1: | 2,425.9 |
| Noise: | 0.2 | | |

## Peak table for sample 1 : F17-2 library

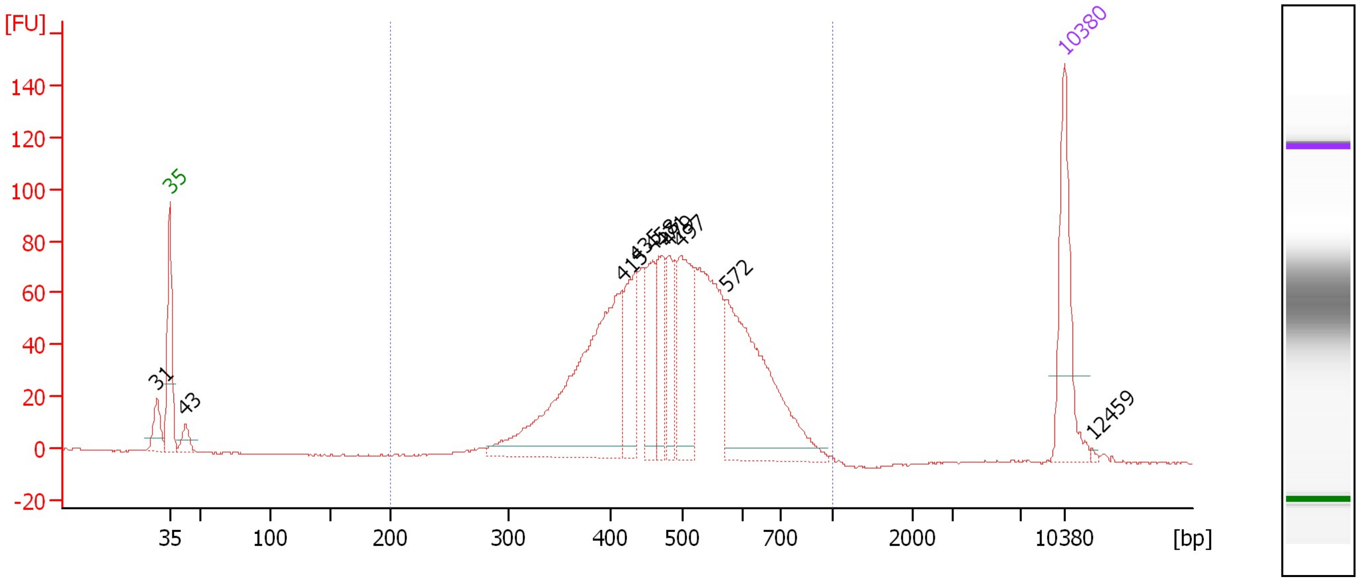
| Peak | | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|---|---|---|---|---|---|---|
| 1 | | 32 | 0.00 | 0.0 | | 42.06 |
| 2 | ◀ | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | | 44 | 17.08 | 592.2 | | 44.36 |
| 4 | | 456 | 456.58 | 1,517.9 | | 80.62 |
| 5 | | 480 | 72.62 | 229.3 | | 81.98 |
| 6 | | 492 | 64.70 | 199.2 | | 82.68 |
| 7 | | 507 | 69.13 | 206.7 | | 83.43 |
| 8 | | 534 | 65.22 | 185.2 | | 84.69 |
| 9 | | 550 | 58.14 | 160.3 | | 85.44 |
| 10 | | 568 | 47.01 | 125.5 | | 86.28 |
| 11 | | 582 | 342.59 | 892.5 | | 86.94 |
| 12 | ▶ | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 13 | | 12,737 | 0.00 | 0.0 | | 115.43 |

## Region table for sample 1 : F17-2 library

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Color | Conc. [pg/µl] | % of Total | Size distribution in CV [%] |
|---|---|---|---|---|---|---|---|---|
| 200 | 1,528 | 525 | 2,425.9 | 4,257.6 | 🟦 | 1,365.84 | 97 | 25.1 |

## Electropherogram Summary Continued ...

F17-3 library [ HiSeq_992 ]



### Overall Results for sample 2 : F17-3 library

Number of peaks found: 10 Corr. Area 1: 1,348.2

Noise: 0.3

### Peak table for sample 2 : F17-3 library

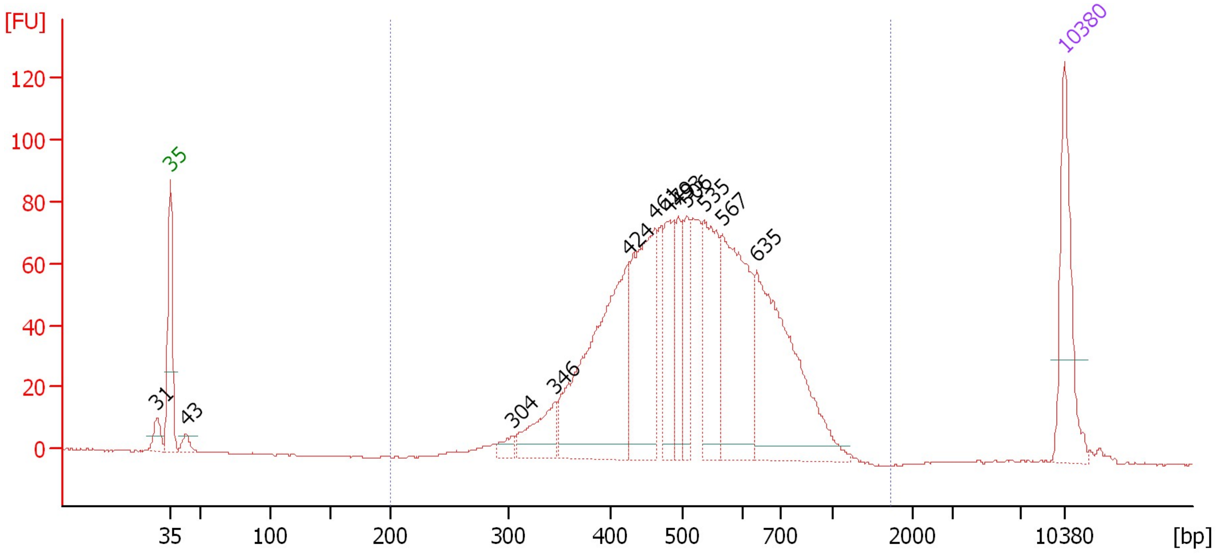
| Peak | | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|---|---|---|---|---|---|---|
| 1 | | 31 | 0.00 | 0.0 | | 41.95 |
| 2 | ◀ | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | | 43 | 20.46 | 726.9 | | 44.19 |
| 4 | | 415 | 292.92 | 1,068.8 | | 78.33 |
| 5 | | 435 | 79.57 | 277.4 | | 79.43 |
| 6 | | 458 | 60.52 | 200.0 | | 80.77 |
| 7 | | 471 | 58.16 | 187.1 | | 81.48 |
| 8 | | 479 | 46.92 | 148.3 | | 81.96 |
| 9 | | 497 | 105.20 | 320.6 | | 82.96 |
| 10 | | 572 | 209.08 | 553.8 | | 86.49 |
| 11 | ▶ | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 12 | | 12,459 | 0.00 | 0.0 | | 115.15 |

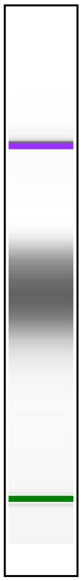
### Region table for sample 2 : F17-3 library

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Color | Conc. [pg/µl] | % of Total | Size distribution in CV [%] |
|---|---|---|---|---|---|---|---|---|
| 200 | 1,000 | 492 | 1,348.2 | 3,412.4 | 🟦 | 1,044.10 | 97 | 21.4 |

## Electropherogram Summary Continued ...

F17-6 library [ HiSeq_992 ]





### Overall Results for sample 3 : F17-6 library

| | | | |
|---|---|---|---|
| Number of peaks found: | 12 | Corr. Area 1: | 1,451.3 |
| Noise: | 0.3 | | |

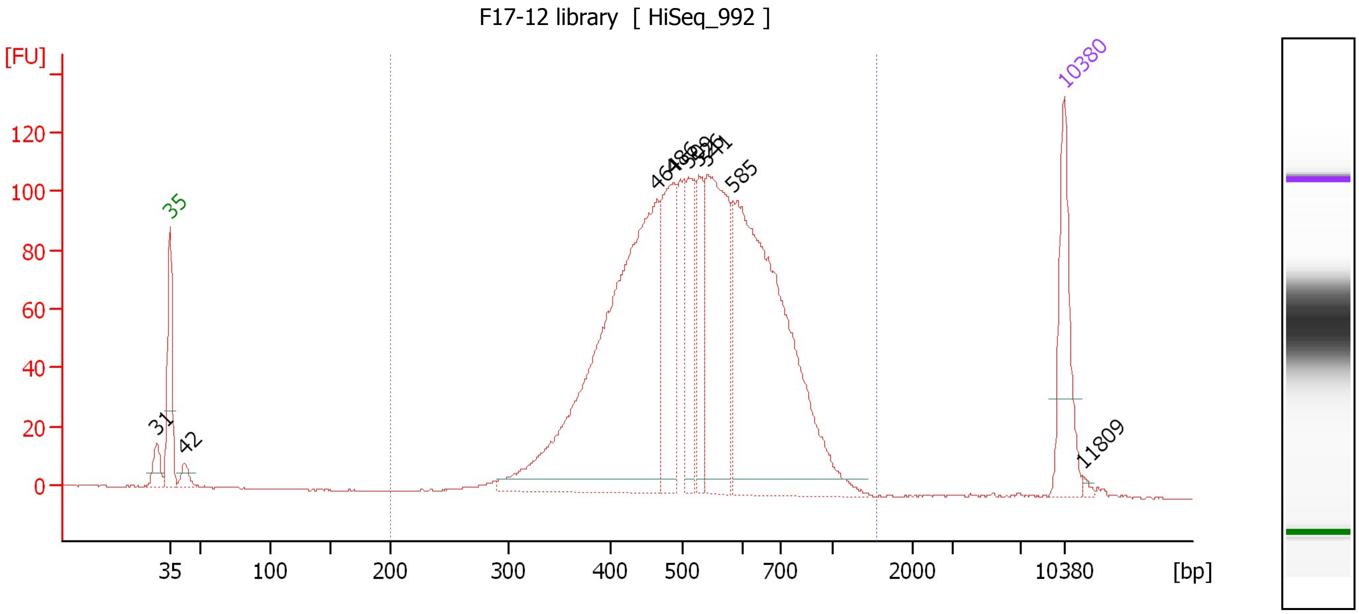
### Peak table for sample 3 : F17-6 library

| Peak | | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|---|---|---|---|---|---|---|
| 1 | | 31 | 0.00 | 0.0 | | 41.98 |
| 2 | ◀ | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | | 43 | 13.68 | 484.9 | | 44.21 |
| 4 | | 304 | 13.56 | 67.6 | | 69.79 |
| 5 | | 346 | 54.69 | 239.7 | | 73.13 |
| 6 | | 424 | 264.21 | 943.3 | | 78.85 |
| 7 | | 461 | 178.13 | 585.1 | | 80.93 |
| 8 | | 479 | 75.96 | 240.1 | | 81.95 |
| 9 | | 493 | 59.30 | 182.3 | | 82.72 |
| 10 | | 506 | 54.48 | 163.1 | | 83.40 |
| 11 | | 535 | 107.87 | 305.5 | | 84.76 |
| 12 | | 567 | 176.29 | 471.0 | | 86.26 |
| 13 | | 635 | 218.22 | 520.3 | | 88.88 |
| 14 | ▶ | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

### Region table for sample 3 : F17-6 library

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Color | Conc. [pg/µl] | % of Total | Size distribution in CV [%] |
|---|---|---|---|---|---|---|---|---|
| 200 | 1,726 | 527 | 1,451.3 | 4,043.2 | 🟦 | 1,299.46 | 98 | 25.2 |

**Electropherogram Summary Continued ...**



### Overall Results for sample 4 : F17-12 library

| | | | |
|---|---|---|---|
| Number of peaks found: | 9 | Corr. Area 1: | 1,966.9 |
| Noise: | 0.2 | | |

### Peak table for sample 4 : F17-12 library

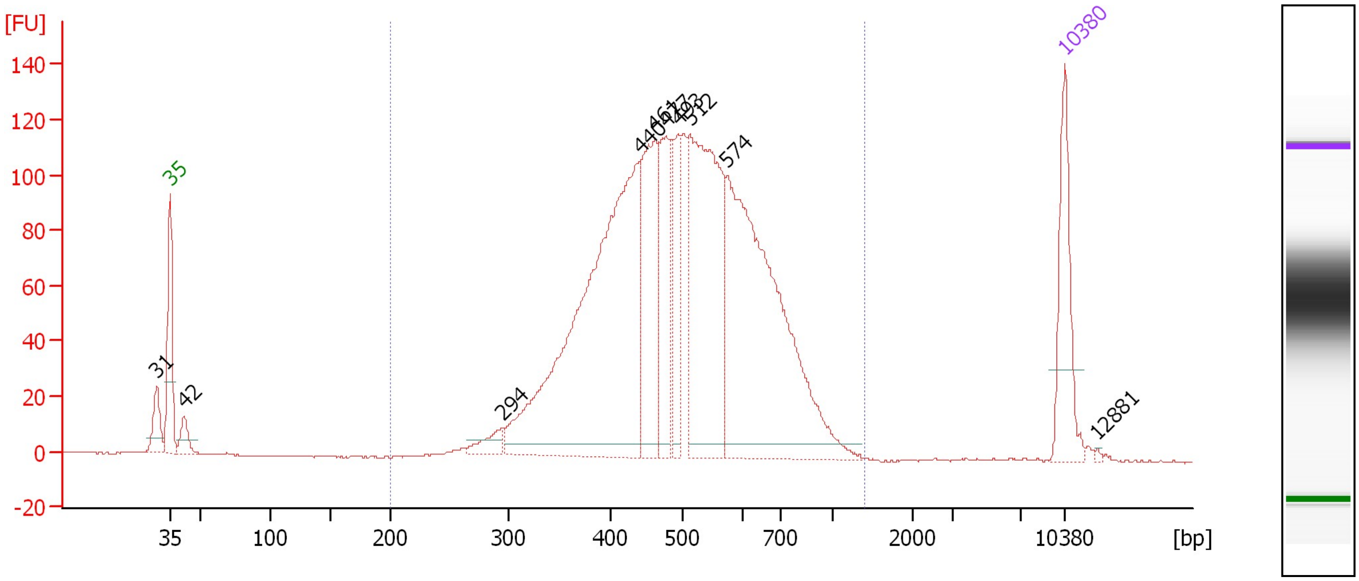
| Peak | | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|---|---|---|---|---|---|---|
| 1 | | 31 | 0.00 | 0.0 | | 41.97 |
| 2 | ◀ | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | | 42 | 17.95 | 643.7 | | 44.13 |
| 4 | | 464 | 592.03 | 1,932.4 | | 81.10 |
| 5 | | 486 | 149.68 | 466.7 | | 82.33 |
| 6 | | 509 | 94.97 | 282.5 | | 83.55 |
| 7 | | 526 | 71.15 | 204.9 | | 84.34 |
| 8 | | 541 | 218.85 | 613.1 | | 85.03 |
| 9 | | 585 | 461.32 | 1,195.0 | | 87.09 |
| 10 | ▶ | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 11 | | 11,809 | 0.00 | 0.0 | | 114.47 |

### Region table for sample 4 : F17-12 library

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Color | Conc. [pg/µl] | % of Total | Size distribution in CV [%] |
|---|---|---|---|---|---|---|---|---|
| 200 | 1,548 | 538 | 1,966.9 | 5,141.4 | 🟦 | 1,691.46 | 98 | 25.3 |

## Electropherogram Summary Continued ...

M11-22 library [ HiSeq_992 ]



### Overall Results for sample 5 : M11-22 library

| | | | |
|---|---|---|---|
| Number of peaks found: | 10 | Corr. Area 1: | 2,237.4 |
| Noise: | 0.2 | | |

### Peak table for sample 5 : M11-22 library

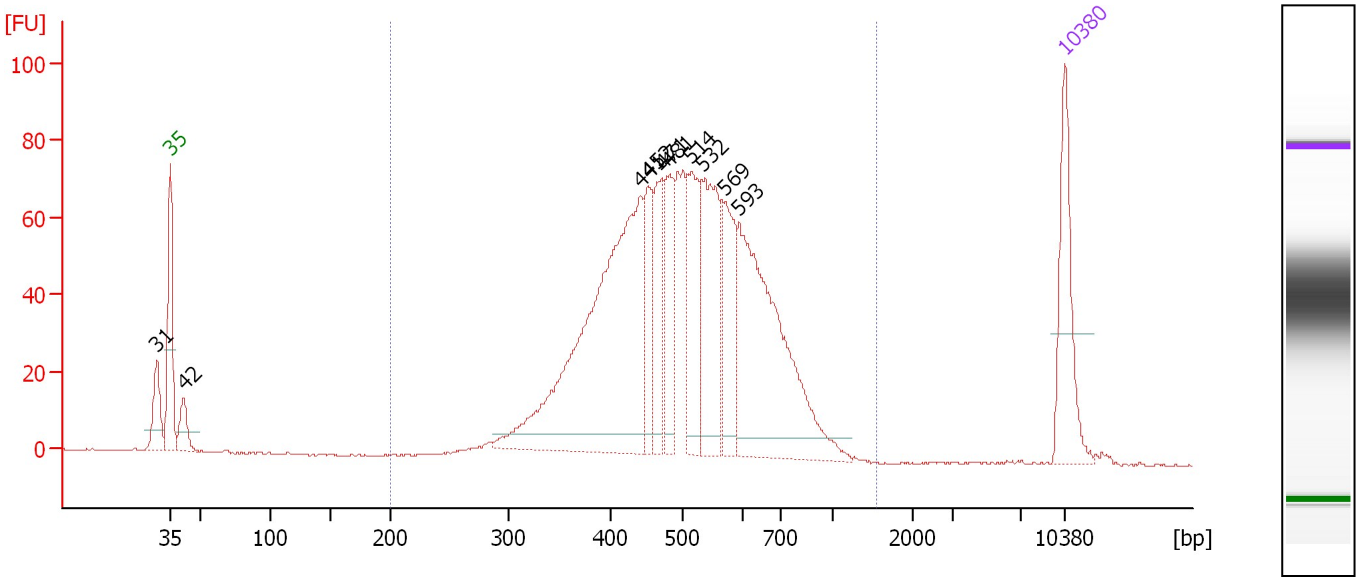
| Peak | | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|---|---|---|---|---|---|---|
| 1 | | 31 | 0.00 | 0.0 | | 41.96 |
| 2 | ◀ | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | | 42 | 27.79 | 1,002.4 | | 44.09 |
| 4 | | 294 | 23.43 | 120.6 | | 68.96 |
| 5 | | 440 | 599.71 | 2,064.5 | | 79.74 |
| 6 | | 461 | 170.98 | 561.7 | | 80.93 |
| 7 | | 477 | 108.05 | 343.2 | | 81.82 |
| 8 | | 493 | 77.64 | 238.7 | | 82.72 |
| 9 | | 512 | 323.16 | 957.1 | | 83.66 |
| 10 | | 574 | 452.93 | 1,195.3 | | 86.59 |
| 11 | ▶ | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |
| 12 | | 12,881 | 0.00 | 0.0 | | 115.58 |

### Region table for sample 5 : M11-22 library

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Color | Conc. [pg/µl] | % of Total | Size distribution in CV [%] |
|---|---|---|---|---|---|---|---|---|
| 200 | 1,391 | 511 | 2,237.4 | 6,012.6 | ■ | 1,870.69 | 97 | 25.7 |

### Electropherogram Summary Continued ...

M14-140 library [ HiSeq_992 ]



**Overall Results for sample 6 :** **M14-140 library**

| | | | |
|---|---|---|---|
| Number of peaks found: | 10 | Corr. Area 1: | 1,356.4 |
| Noise: | 0.2 | | |

**Peak table for sample 6 :** **M14-140 library**

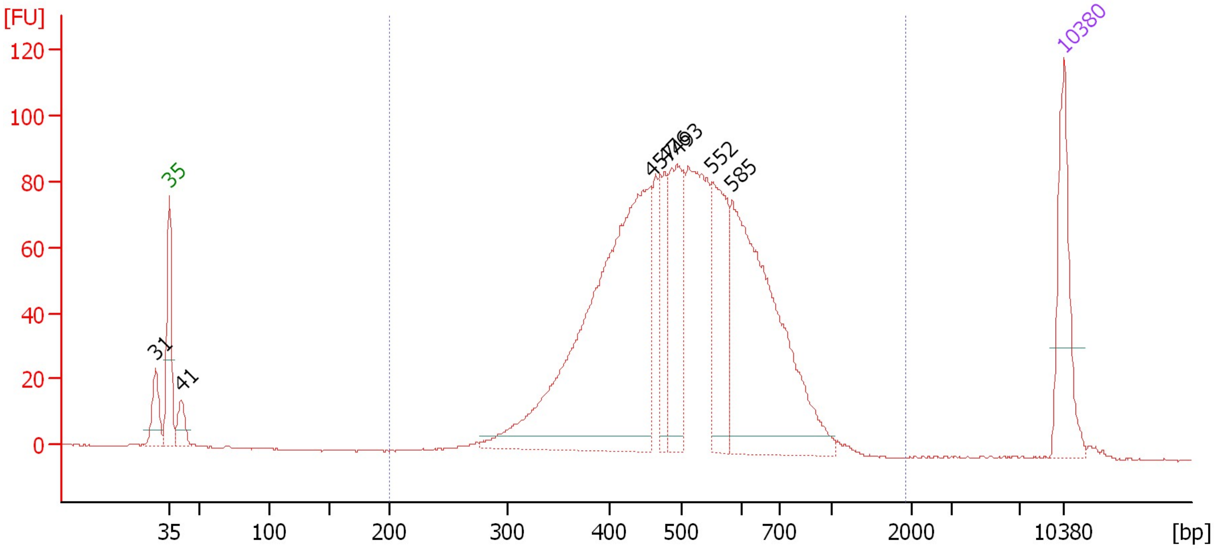
| Peak | | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|---|---|---|---|---|---|---|
| 1 | | 31 | 0.00 | 0.0 | | 41.95 |
| 2 | ◀ | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | | 42 | 35.82 | 1,301.1 | | 44.05 |
| 4 | | 441 | 459.27 | 1,577.4 | | 79.79 |
| 5 | | 452 | 65.58 | 219.9 | | 80.39 |
| 6 | | 471 | 75.82 | 243.8 | | 81.49 |
| 7 | | 481 | 67.52 | 212.7 | | 82.04 |
| 8 | | 514 | 107.49 | 316.7 | | 83.78 |
| 9 | | 532 | 135.65 | 386.1 | | 84.63 |
| 10 | | 569 | 79.33 | 211.4 | | 86.33 |
| 11 | | 593 | 278.37 | 711.2 | | 87.47 |
| 12 | ▶ | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

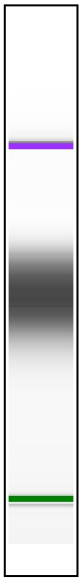
**Region table for sample 6 :** **M14-140 library**

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Color | Conc. [pg/µl] | % of Total | Size distribution in CV [%] |
|---|---|---|---|---|---|---|---|---|
| 200 | 1,553 | 513 | 1,356.4 | 4,650.3 | 🟦 | 1,461.43 | 96 | 24.4 |

## Electropherogram Summary Continued ...

M14-10 library [ HiSeq_992 ]





**Overall Results for sample 7 :** **M14-10 library**

| | | | |
|---|---|---|---|
| Number of peaks found: | 7 | Corr. Area 1: | 1,596.8 |
| Noise: | 0.2 | | |

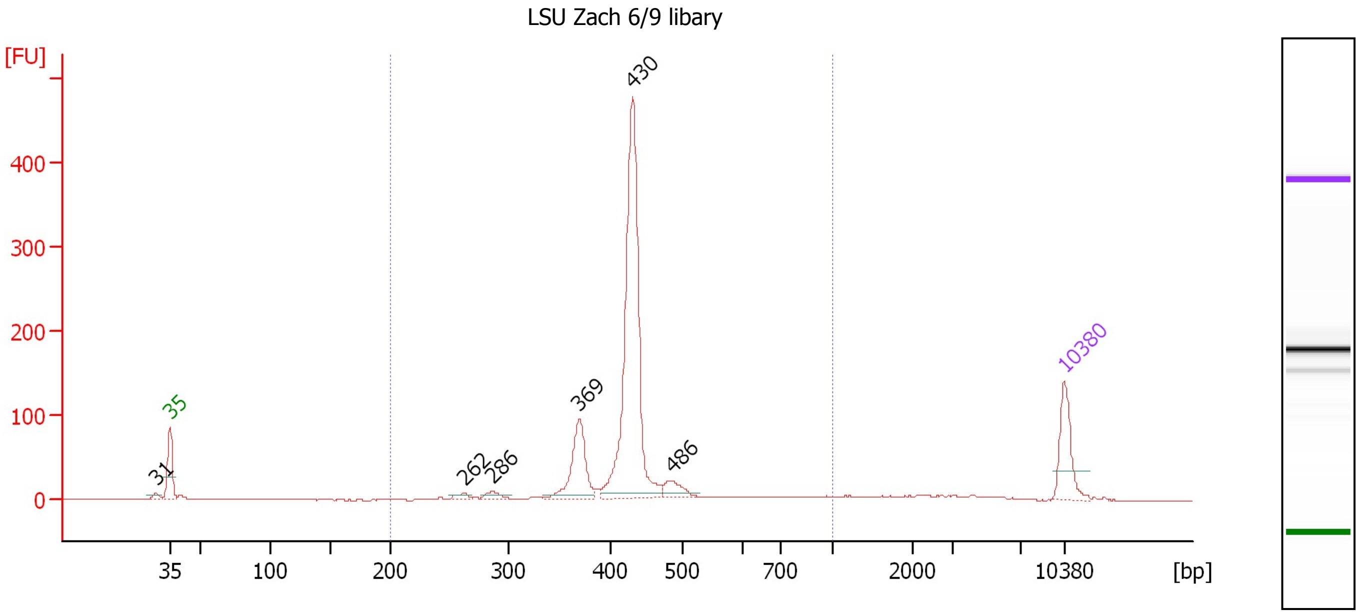
**Peak table for sample 7 :** **M14-10 library**

| Peak | | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|---|---|---|---|---|---|---|
| 1 | | 31 | 0.00 | 0.0 | | 41.95 |
| 2 | ◀ | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | | 41 | 31.72 | 1,169.5 | | 43.95 |
| 4 | | 457 | 547.34 | 1,815.6 | | 80.68 |
| 5 | | 476 | 64.88 | 206.4 | | 81.78 |
| 6 | | 493 | 130.68 | 401.5 | | 82.73 |
| 7 | | 552 | 124.13 | 341.0 | | 85.53 |
| 8 | | 585 | 332.72 | 862.3 | | 87.08 |
| 9 | ▶ | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

**Region table for sample 7 :** **M14-10 library**

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Color | Conc. [pg/µl] | % of Total | Size distribution in CV [%] |
|---|---|---|---|---|---|---|---|---|
| 200 | 1,942 | 516 | 1,596.8 | 4,771.6 | ■ | 1,506.31 | 96 | 24.7 |

**Electropherogram Summary Continued ...**



**Overall Results for sample 8 :** **LSU Zach 6/9 libary**

| | | | |
|---|---|---|---|
| Number of peaks found: | 6 | Corr. Area 1: | 1,046.4 |
| Noise: | 0.2 | | |

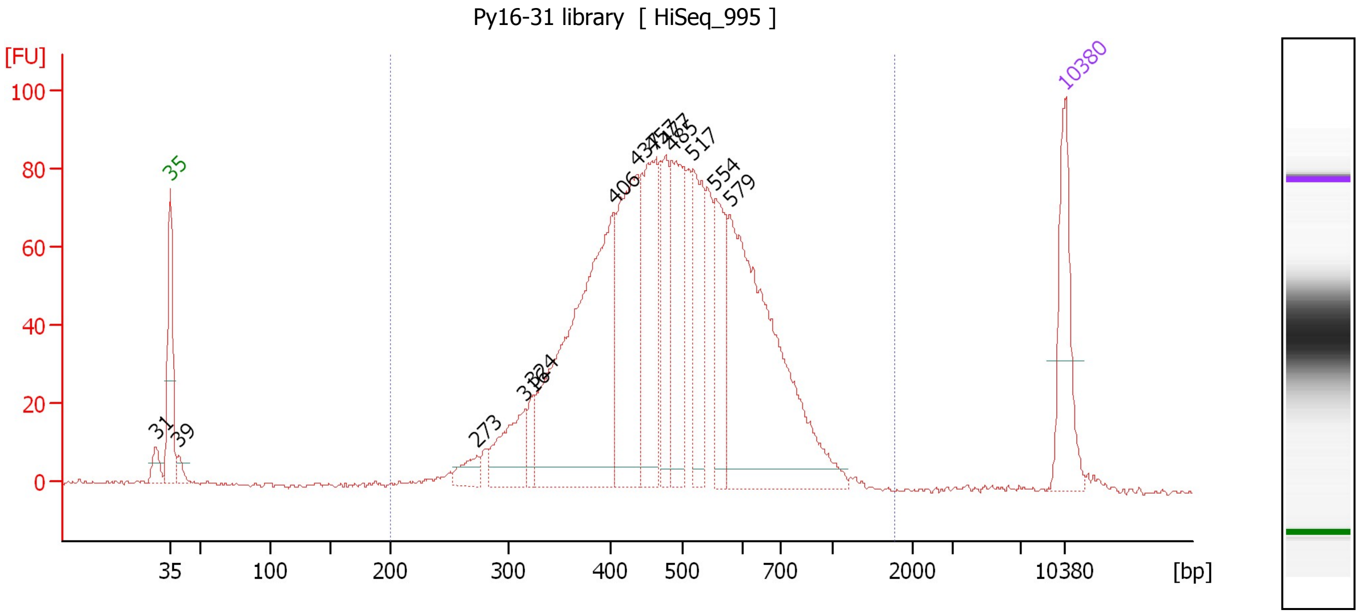
**Peak table for sample 8 :** **LSU Zach 6/9 libary**

| Peak | | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|---|---|---|---|---|---|---|
| 1 | | 31 | 0.00 | 0.0 | | 41.94 |
| 2 | ◀ | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | | 262 | 13.41 | 77.4 | | 66.03 |
| 4 | | 286 | 17.53 | 92.9 | | 68.20 |
| 5 | | 369 | 141.12 | 579.3 | | 75.00 |
| 6 | | 430 | 621.53 | 2,188.4 | | 79.18 |
| 7 | | 486 | 35.47 | 110.7 | | 82.31 |
| 8 | ▶ | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

**Region table for sample 8 :** **LSU Zach 6/9 libary**

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Color | Conc. [pg/µl] | % of Total | Size distribution in CV [%] |
|---|---|---|---|---|---|---|---|---|
| 200 | 1,000 | 431 | 1,046.4 | 3,258.0 | ■ | 894.67 | 93 | 18.3 |

**Electropherogram Summary Continued ...**



Py16-31 library [ HiSeq_995 ]

**Overall Results for sample 9 :** **Py16-31 library**

| | | | |
|---|---|---|---|
| Number of peaks found: | 13 | Corr. Area 1: | 1,700.0 |
| Noise: | 0.5 | | |

**Peak table for sample 9 :** **Py16-31 library**

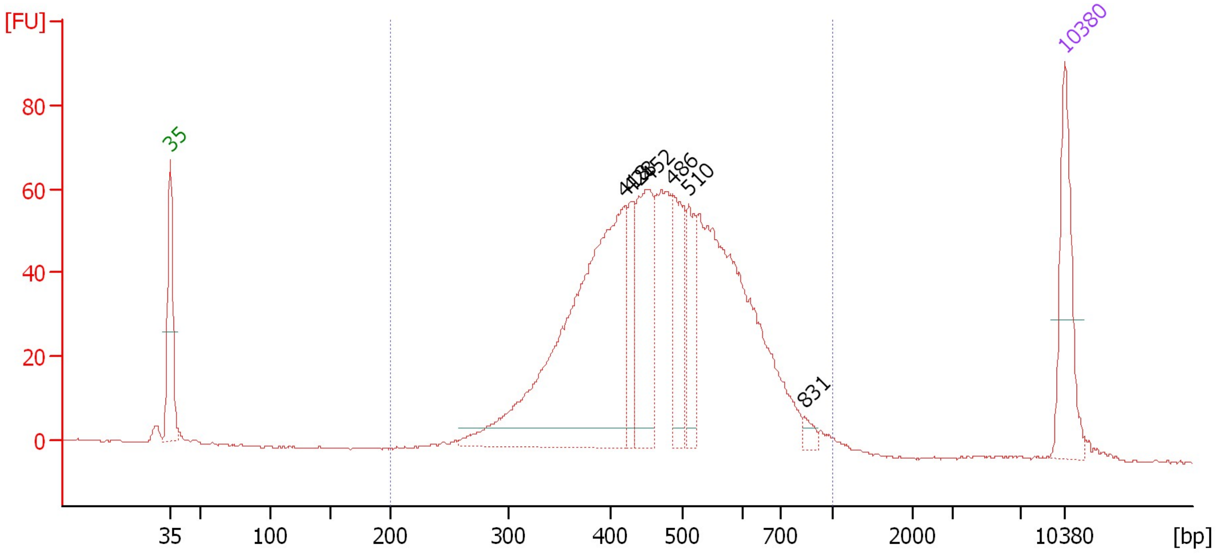
| Peak | | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|---|---|---|---|---|---|---|
| 1 | | 31 | 0.00 | 0.0 | | 41.88 |
| 2 | ◀ | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | | 39 | 14.32 | 553.3 | | 43.66 |
| 4 | | 273 | 24.16 | 133.9 | | 67.04 |
| 5 | | 316 | 75.29 | 361.2 | | 70.75 |
| 6 | | 324 | 24.62 | 115.1 | | 71.41 |
| 7 | | 406 | 412.37 | 1,539.8 | | 77.80 |
| 8 | | 437 | 217.92 | 755.2 | | 79.57 |
| 9 | | 457 | 162.62 | 539.2 | | 80.69 |
| 10 | | 477 | 87.20 | 277.1 | | 81.80 |
| 11 | | 485 | 131.34 | 410.5 | | 82.26 |
| 12 | | 517 | 110.01 | 322.1 | | 83.93 |
| 13 | | 554 | 85.85 | 234.7 | | 85.66 |
| 14 | | 579 | 373.66 | 977.5 | | 86.83 |
| 15 | ▶ | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

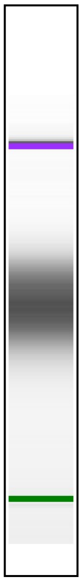
**Region table for sample 9 :** **Py16-31 library**

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Co Conc. lor [pg/µl] | % of Total | Size distribution in CV [%] |
|---|---|---|---|---|---|---|---|
| 200 | 1,777 | 498 | 1,700.0 | 6,386.8 | ■ 1,901.59 | 97 | 28.8 |

## Electropherogram Summary Continued ...

Py16-32 library [ HiSeq_995 ]





### Overall Results for sample 10 : Py16-32 library

| | | | |
|---|---|---|---|
| Number of peaks found: | 6 | Corr. Area 1: | 1,169.7 |
| Noise: | 0.2 | | |

### Peak table for sample 10 : Py16-32 library

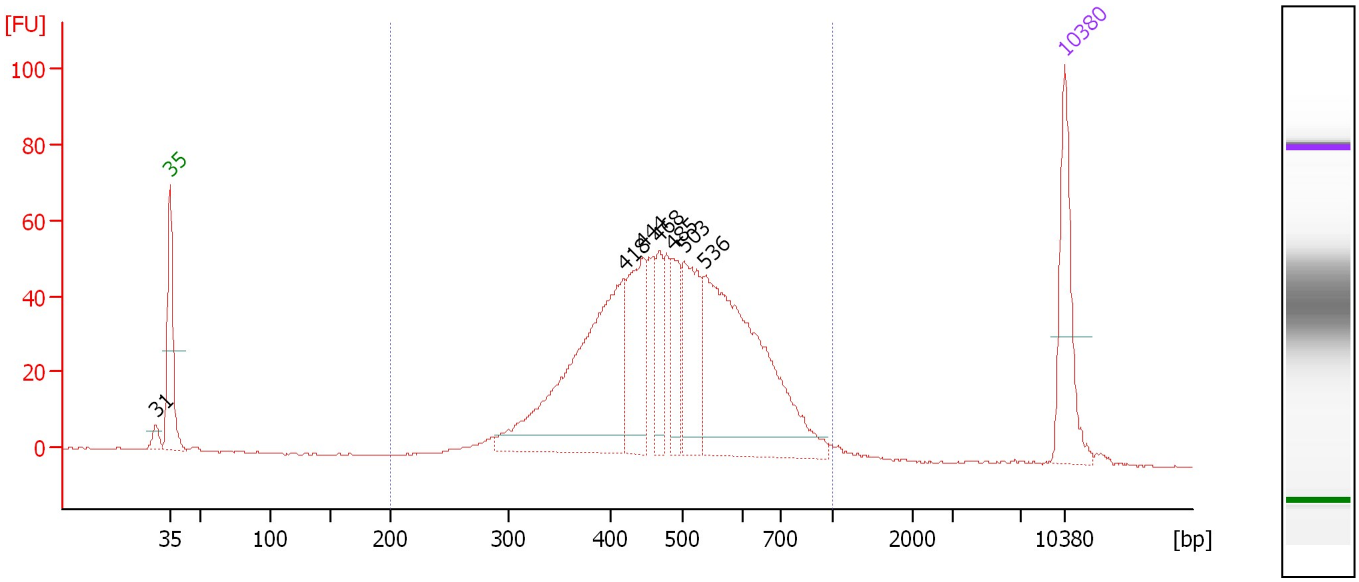
| Peak | | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|---|---|---|---|---|---|---|
| 1 | ◀ | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 2 | | 418 | 475.38 | 1,723.2 | | 78.49 |
| 3 | | 428 | 59.03 | 209.0 | | 79.05 |
| 4 | | 452 | 138.51 | 464.7 | | 80.38 |
| 5 | | 486 | 87.54 | 272.9 | | 82.33 |
| 6 | | 510 | 65.47 | 194.7 | | 83.56 |
| 7 | | 831 | 8.49 | 15.5 | | 92.59 |
| 8 | ▶ | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

### Region table for sample 10 : Py16-32 library

| From [bp] | To [bp] | Average Size [bp] | Corr. Area | Molarity [pmol/l] | Color | Conc. [pg/µl] | % of Total | Size distribution in CV [%] |
|---|---|---|---|---|---|---|---|---|
| 200 | 1,000 | 482 | 1,169.7 | 4,740.1 | 🟦 | 1,401.15 | 98 | 23.8 |

## Electropherogram Summary Continued ...

M14-45 library [ HiSeq_995 ]



**Overall Results for sample 11 :** **M14-45 library**

| | | | |
|---|---|---|---|
| Number of peaks found: | 7 | Corr. Area 1: | 985.4 |
| Noise: | 0.2 | | |

**Peak table for sample 11 :** **M14-45 library**

| Peak | | Size [bp] | Conc. [pg/µl] | Molarity [pmol/l] | Observations | Aligned Migration Time [s] |
|---|---|---|---|---|---|---|
| 1 | | 31 | 0.00 | 0.0 | | 41.86 |
| 2 | ◀ | 35 | 125.00 | 5,411.3 | Lower Marker | 43.00 |
| 3 | | 418 | 305.03 | 1,105.4 | | 78.49 |
| 4 | | 444 | 111.56 | 380.8 | | 79.95 |
| 5 | | 468 | 48.05 | 155.6 | | 81.30 |
| 6 | | 485 | 53.09 | 165.7 | | 82.29 |
| 7 | | 503 | 100.41 | 302.2 | | 83.27 |
| 8 | | 536 | 325.63 | 921.3 | | 84.78 |
| 9 | ▶ | 10,380 | 75.00 | 10.9 | Upper Marker | 113.00 |

**Region table for sample 11 :** **M14-45 library**

| 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 | 494 | 985.4 | 3,594.4 | ■ 1,085.56 | 98 | 24.2 |

Assay Class: High Sensitivity DNA Assay
Data Path: C:\...ibraries9-15_HiSeq995-Libraries12-14_LSUZach6-9-Library.xad

Created: 6/20/2017 10:12:23 AM
Modified: 6/20/2017 11:03:49 AM

## Gel Image