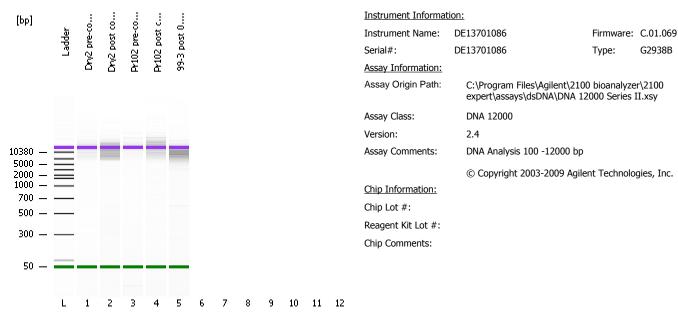
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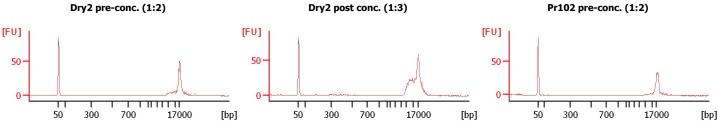
Assay Class: DNA 12000 5/16/2014 9:20:12 AM Created: C:\... bioanalyzer\2100 expert\Data\2014-05-16\2014-05-16_001.xad Modified: 5/16/2014 10:05:31 AM Data Path:

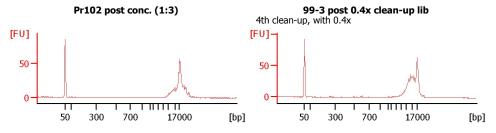
G2938B

Printed: 5/16/2014 10:05:53 AM

Electrophoresis File Run Summary







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Assay Class: 5/16/2014 9:20:12 AM 5/16/2014 10:05:31 AM DNA 12000 Created: C:\... bioanalyzer\2100 expert\Data\2014-05-16\2014-05-16_001.xad Data Path: Modified: **Electrophoresis File Run Summary (Chip Summary) Sample Name** Sample Comment Rest. Digest Status Observation **Result Label Result Color** Dry2 pre-conc. (1:2) Dry2 post conc. (1:3) Pr102 pre-conc. (1:2) Pr102 post conc. (1:3) 99-3 post 0.4x clean-up lib 4th clean-up, with 0.4x Ladder Chip Lot # Reagent Kit Lot #

Chip Comments:

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Assay Class: DNA 12000 Created: 5/16/2014 9:20:12 AM Data Path: C:\... bioanalyzer\2100 expert\Data\2014-05-16\2014-05-16_001.xad Modified: 5/16/2014 10:05:31 AM

Electrophoresis Assay Details

General Analysis Settings

Number of Available Sample and Ladder Wells (Max.): 13

Minimum Visible Range [s]: 20
Maximum Visible Range [s]: 99
Start Analysis Time Range [s]: 20
End Analysis Time Range [s]: 98.95
Ladder Concentration [ng/µl]: 44
Uses Standard Area for Ladder Fragments
Lower Marker Concentration [ng/µl]: 8.3
Upper Marker Concentration [ng/µl]: 4.2
Used Upper Marker for Quantitation
This is a Qualitative Assay Only
Standard Curve Fit is Point to Point
Show Data Aligned to Lower and Upper Marker

Integrator Settings

Integration Start Time [s]: 20 Integration End Time [s]: 98.95

Slope Threshold: 0.8
Height Threshold [FU]: 20
Area Threshold: 0.1
Width Threshold [s]: 0.5
Baseline Plateau [s]: 0.5

Filter Settings

Filter Width [s]: 0.5 Polynomial Order: 4

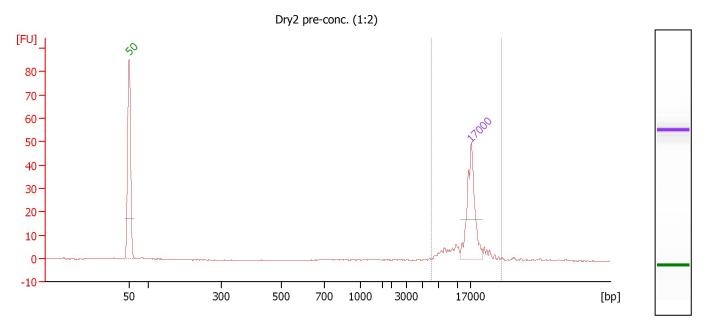
Ladder

| Size | Area |
|-------|---|
| 50 | 120 |
| 100 | 48 |
| 300 | 61 |
| 500 | 77 |
| 700 | 81 |
| 1000 | 86 |
| 1500 | 92 |
| 2000 | 92 |
| 3000 | 96 |
| 5000 | 100 |
| 7000 | 98 |
| 10380 | 102 |
| 17000 | 110 |
| | 50 100 300 500 700 1000 1500 2000 3000 5000 7000 10380 |

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Assay Class: DNA 12000 Created: 5/16/2014 9:20:12 AM Data Path: C:\... bioanalyzer\2100 expert\Data\2014-05-16\2014-05-16_001.xad Modified: 5/16/2014 10:05:31 AM

Electropherogram Summary



Overall Results for sample 1: <u>Dry2 pre-conc. (1:2)</u>

Number of peaks found: 0 Area 1: 22.8

Peak table for sample 1 : <u>Dry2 pre-conc. (1:2)</u>

 Peak
 Size [bp]
 Conc. [ng/μl]
 Molarity [nmol/l]
 Observations

 1
 4
 50
 8.30
 251.5
 Lower Marker

 2
 17,000
 4.20
 0.4
 Upper Marker

Region table for sample 1: <u>Dry2 pre-conc. (1:2)</u>

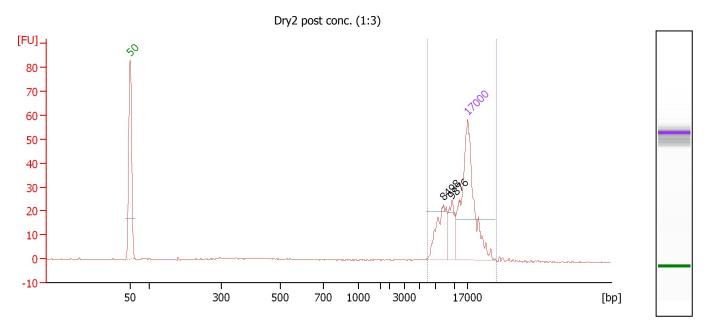
 From [bp]
 To [bp]
 Average Size [bp]
 Conc. [ng/μl] Col Area or [w]
 % of Total [w]
 Size distribution in CV [w]

 6,163
 32,194
 16,253
 1.66
 22.8
 76
 29.3

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Assay Class: DNA 12000 Created: 5/16/2014 9:20:12 AM Data Path: C:\... bioanalyzer\2100 expert\Data\2014-05-16\2014-05-16_001.xad Modified: 5/16/2014 10:05:31 AM

Electropherogram Summary Continued ...



Overall Results for sample 2: <u>Dry2 post conc. (1:3)</u>

Number of peaks found: 2 Area 1: 63.3

| Peak table for sample 2: | Dry2 post conc. (| 1:3) |
|----------------------------|-------------------|------|
| i can table for sample 2 i | Dig post colici (| |

| Peak | | Size [bp] | Conc. [ng/µl] | Molarity [nmol/l] | Observations |
|------|---|-----------|---------------|-------------------|--------------|
| 1 | 4 | 50 | 8.30 | 251.5 | Lower Marker |
| 2 | | 8,498 | 1.44 | 0.3 | |
| 3 | | 9,876 | 0.84 | 0.1 | |
| 4 | | 17,000 | 4.20 | 0.4 | Upper Marker |

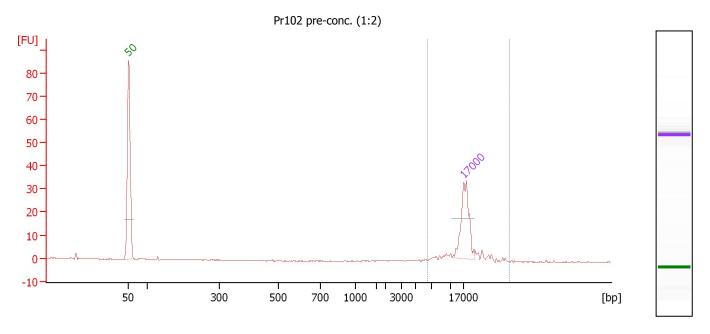
Region table for sample 2: <u>Dry2 post conc. (1:3)</u>

| From [bp] To [bp] | | Average Size Conc. [ng/µl] Col Are | | ıl] Col Area | % of Total | Size distribution in C |
|-------------------|--------|------------------------------------|------|--------------|------------|------------------------|
| | | [bp] | | or | | [%] |
| 6.024 | 31 602 | 14 490 | 2 42 | 63.3 | 77 | 37.0 |

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Assay Class: DNA 12000 Created: 5/16/2014 9:20:12 AM Data Path: C:\... bioanalyzer\2100 expert\Data\2014-05-16\2014-05-16_001.xad Modified: 5/16/2014 10:05:31 AM

Electropherogram Summary Continued ...



Overall Results for sample 3: Pr102 pre-conc. (1:2)

Number of peaks found: 0 Area 1: 14.0

Peak table for sample 3 : <u>Pr102 pre-conc. (1:2)</u>

 Peak
 Size [bp]
 Conc. [ng/μl]
 Molarity [nmol/l]
 Observations

 1
 4
 50
 8.30
 251.5
 Lower Marker

 2
 17,000
 4.20
 0.4
 Upper Marker

Region table for sample 3: Pr102 pre-conc. (1:2)

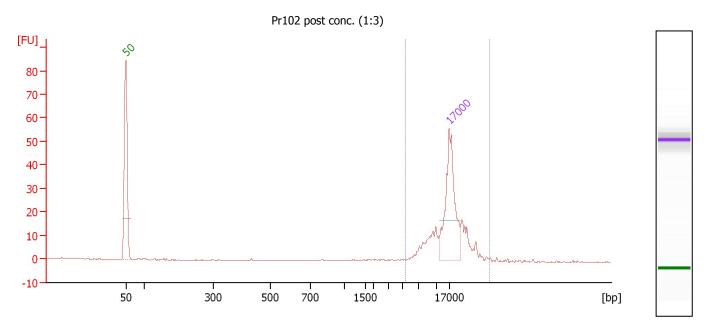
 From [bp]
 To [bp]
 Average Size [bp]
 Conc. [ng/μl] Col Area or [w]
 % of Total [w]
 Size distribution in CV [w]

 6,490
 40,101
 17,938
 1.37
 14.0
 80
 29.3

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Assay Class: DNA 12000 Created: 5/16/2014 9:20:12 AM Data Path: C:\... bioanalyzer\2100 expert\Data\2014-05-16\2014-05-16_001.xad Modified: 5/16/2014 10:05:31 AM

Electropherogram Summary Continued ...



Overall Results for sample 4: Pr102 post conc. (1:3)

Number of peaks found: 0 Area 1: 57.4

Peak table for sample 4 : Pr102 post conc. (1:3)

 Peak
 Size [bp]
 Conc. [ng/μl]
 Molarity [nmol/l]
 Observations

 1
 4
 50
 8.30
 251.5
 Lower Marker

 2
 17,000
 4.20
 0.4
 Upper Marker

Region table for sample 4: Pr102 post conc. (1:3)

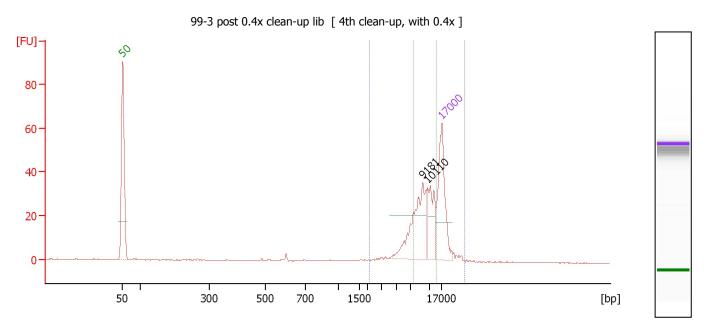
 From [bp]
 To [bp]
 Average Size [bp]
 Conc. [ng/μl] Col Area or [w]
 % of Total [w]
 Size distribution in CV [w]

 5,285
 38,398
 17,525
 2.98
 57.4
 89
 36.4

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Assay Class: DNA 12000 Created: 5/16/2014 9:20:12 AM Data Path: C:\... bioanalyzer\2100 expert\Data\2014-05-16\2014-05-16_001.xad Modified: 5/16/2014 10:05:31 AM

Electropherogram Summary Continued ...



Overall Results for sample 5 : 99-3 post 0.4x clean-up lib

Number of peaks found: 2 Area 2: 91.3

Area 1: 128.6

| Peak table for sample 5: | 99-3 post 0.4x clean-up lib |
|---------------------------|-----------------------------|
| Peak table for sample 5 : | 99-3 DOST U.4X Clean-UD IID |

| Peak | | Size [bp] | Conc. [ng/µl] | Molarity [nmol/l] | Observations |
|------|---|-----------|---------------|-------------------|--------------|
| 1 | 4 | 50 | 8.30 | 251.5 | Lower Marker |
| 2 | | 9,181 | 4.63 | 0.8 | |
| 3 | | 10,110 | 2.21 | 0.3 | |
| 4 | | 17,000 | 4.20 | 0.4 | Upper Marker |

| Region table for sample 5 | : <u>99</u> | 9-3 post 0. | <u>4x clean-up lib</u> |
|---------------------------|-------------|-------------|------------------------|
|---------------------------|-------------|-------------|------------------------|

| From [bp] | To [bp] | Average Size [bp] | Conc. [ng/µl] | Col Area or | % of Total | Size distribution in CV [%] |
|-----------|---------|----------------------|---------------|----------------|------------|-----------------------------|
| 2,171 | 29,159 | 12,177 | 8.20 | 128.6 | 87 | 40.3 |
| 7.534 | 13,992 | 9.929 | 5.80 | 91.3 | 62 | 16.8 |

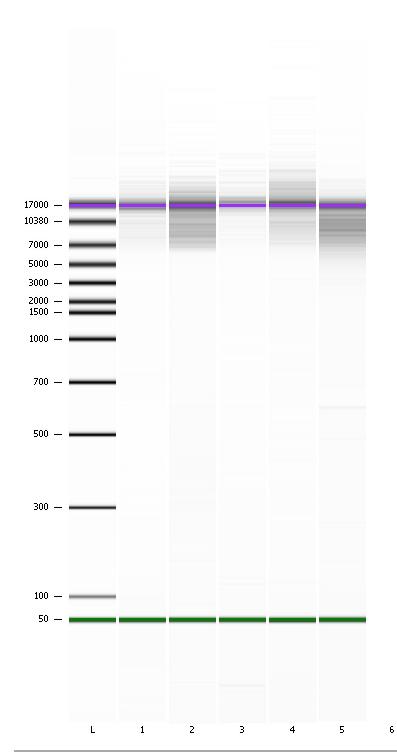
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Assay Class: DNA 12000 C:\... bioanalyzer\2100 expert\Data\2014-05-16\2014-05-16_001.xad

Created: 5/16/2014 9:20:12 AM Modified: 5/16/2014 10:05:31 AM

Gel Image

Dry2 pre-co...
Pr102 pre-co...



7

8

10

11

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12

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Assay Class: DNA 12000

5/16/2014 9:20:12 AM 5/16/2014 10:05:31 AM Created: C:\... bioanalyzer\2100 expert\Data\2014-05-16\2014-05-16_001.xad Data Path: Modified:

Invalid Samples

Sample 6 has not been run, no results available.

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.

Sample 12 has not been run, no results available.