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Instrument controller software run summary:

Filename and data path: C:\Users\aati\Desktop\Customer Data\Wipapat Kladwang\2020 12 08\

Ann_1_11232020 15-42-56\2020 12 08 15H 42M.raw

Created: Tuesday, December 08, 2020 4:16:51 PM

Number of capillaries: 96

Array serial number: 082118-06SFS

Effect length: 33 cm Array usage count: 329

Instrument type: 5300 Fragment Analyzer

Instrument controller software version: 1.2.0.11

Device serial number: 3414

Method Information

Method name: DNF-471-33 - SS Total RNA 15nt.mthds

Gel prime: No

Full conditioning: Yes
Gel prime to buffer: Yes
Gel selection: Gel 1

Perform prerun: 8.0 kV, 30 sec.

Rinse: No Marker 1: No

Rinse: Tray: 3, Row: A, Dip count: 2 Sample injection: 5.0 kV, 4 sec. Separation: 8.0 kV, 40.0 min. Tray name: Ann_1_11232020

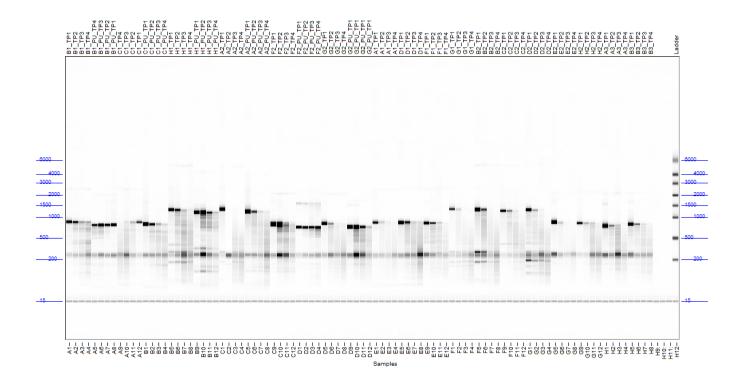
Analysis mode: RNA (Eukaryotic)

Notes

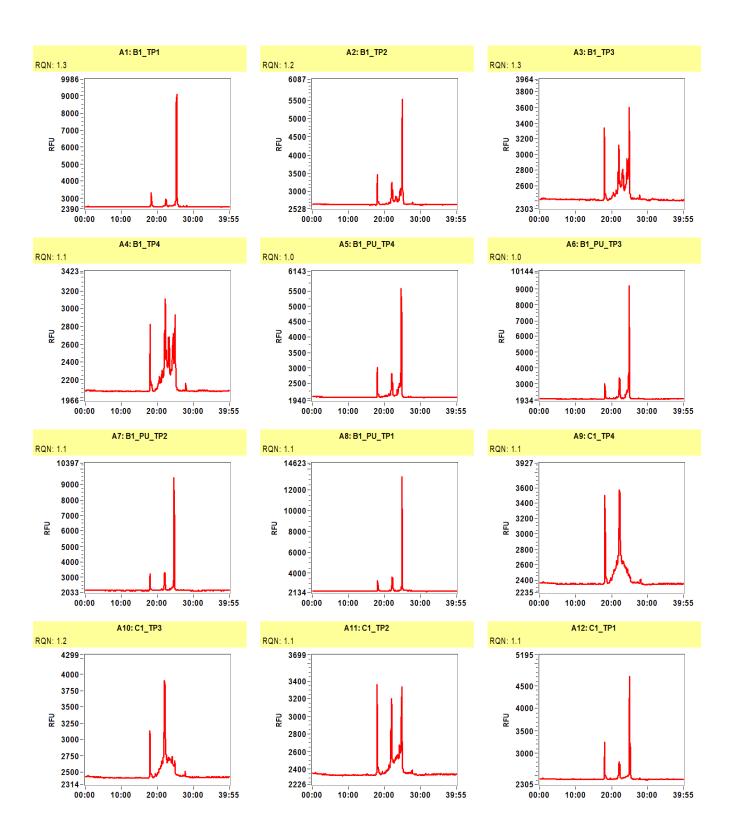
First Trial Run

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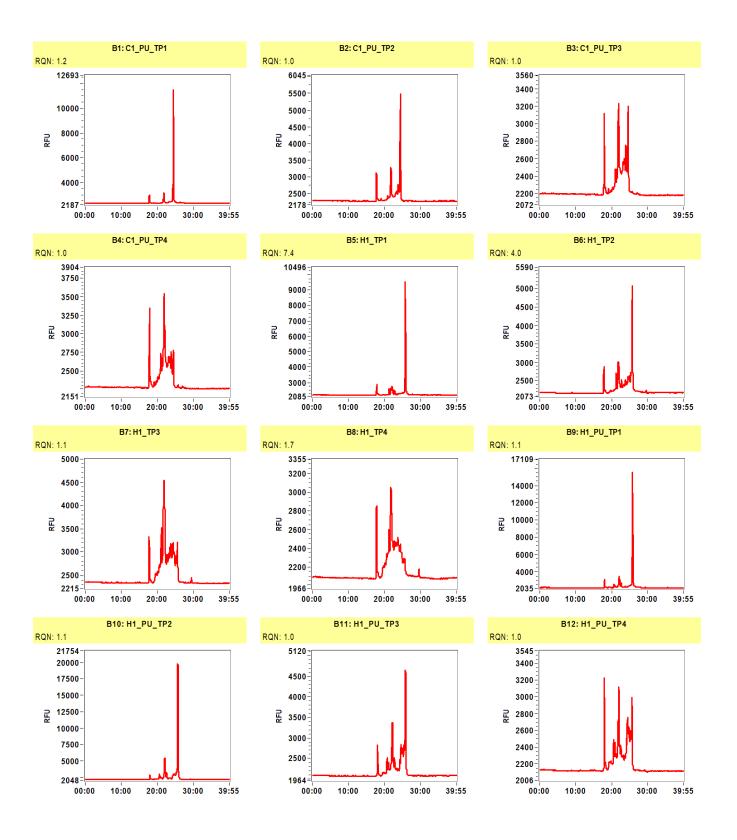
Gel Image



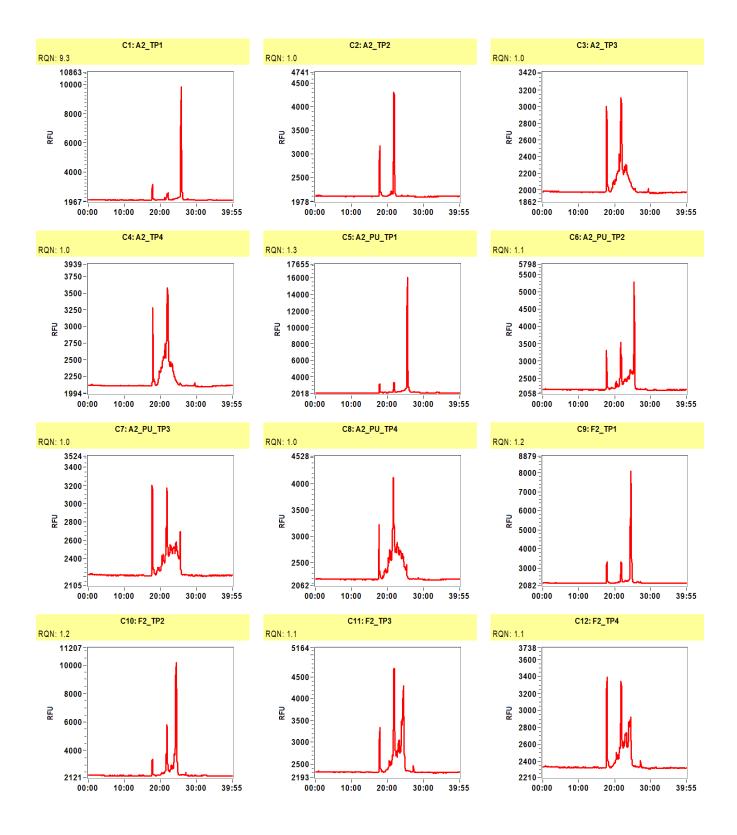
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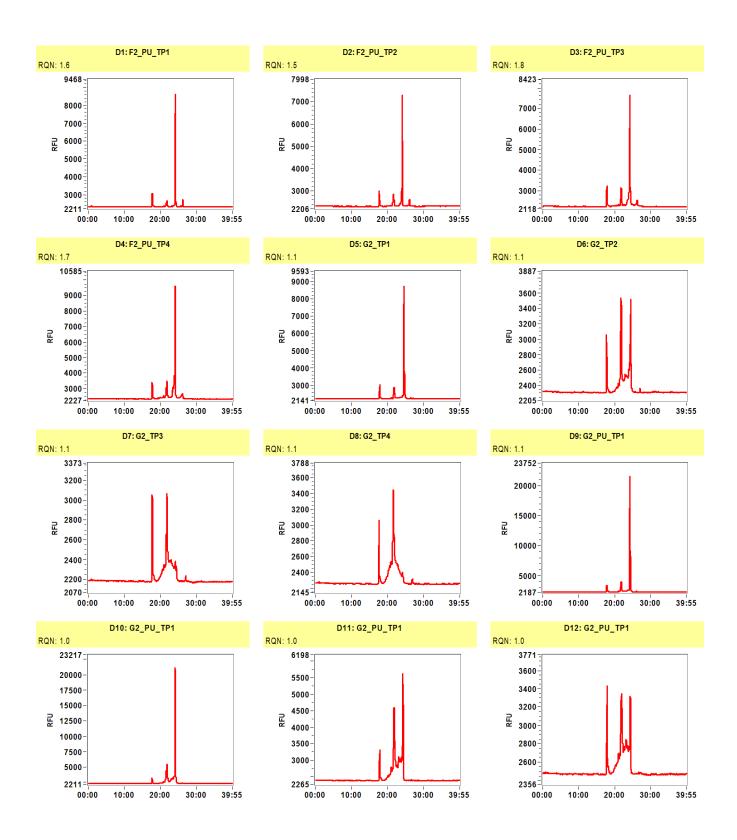
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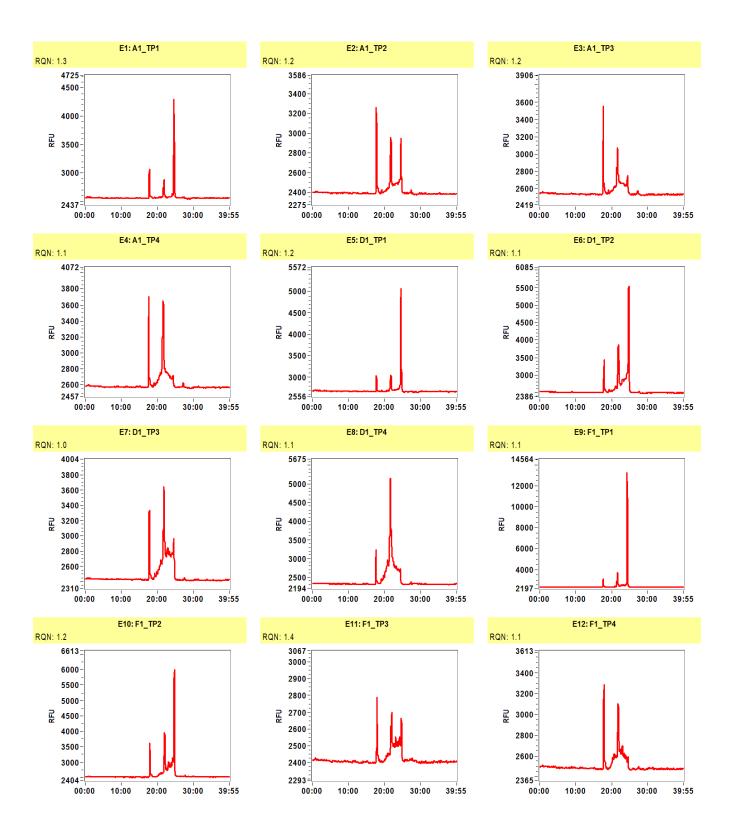
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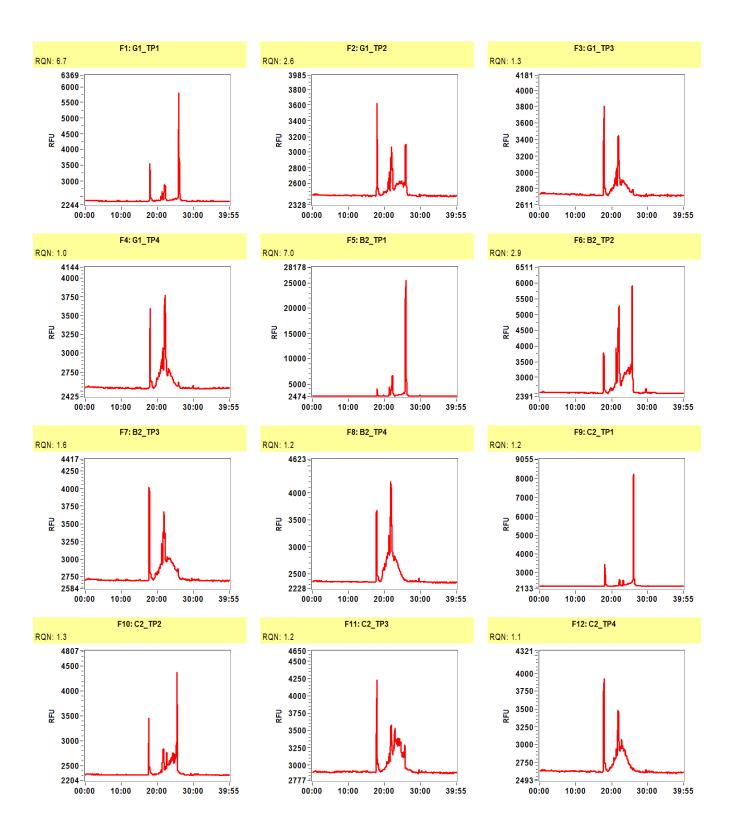
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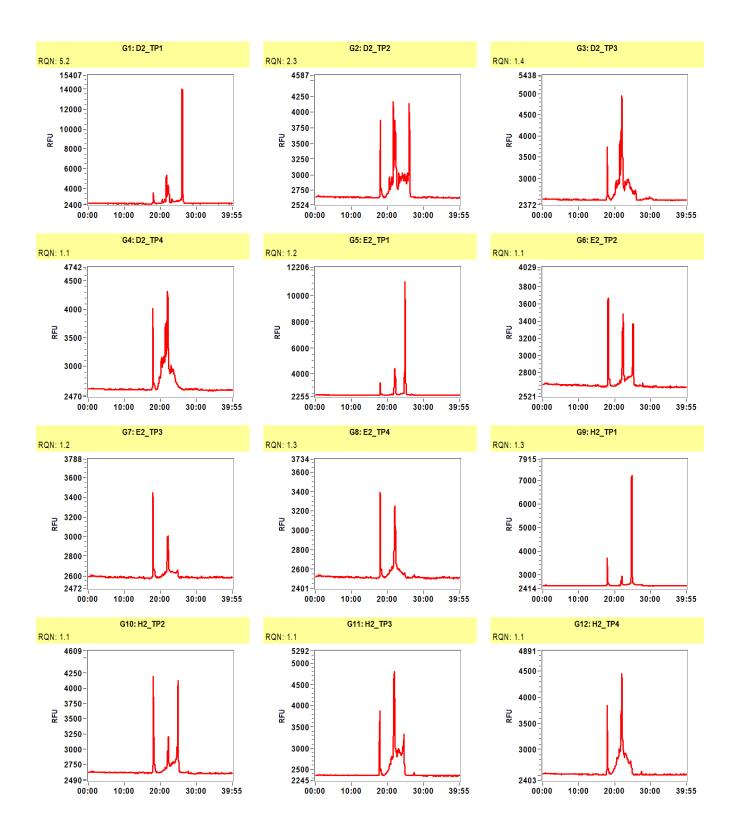
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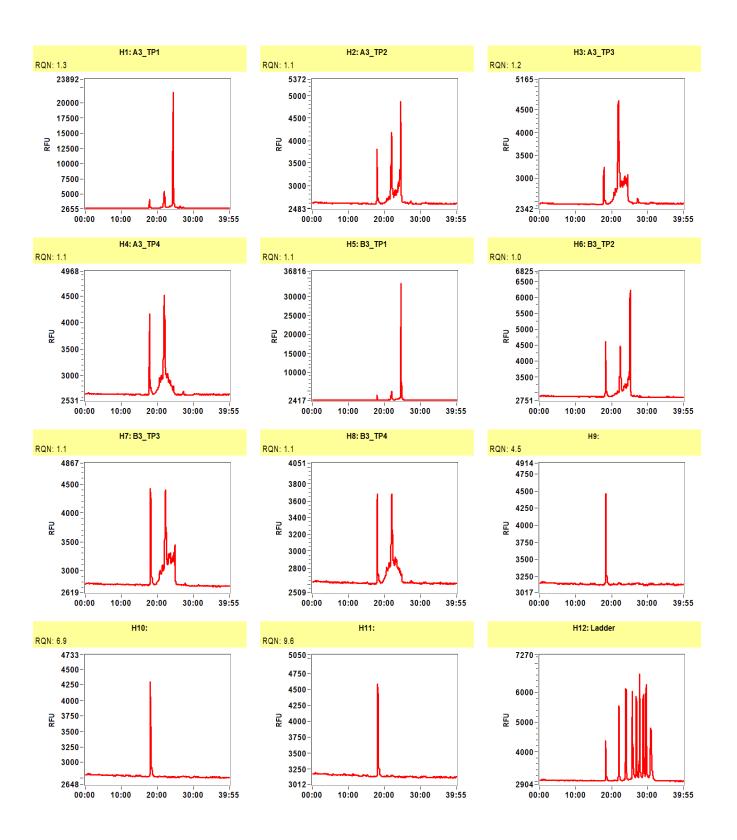
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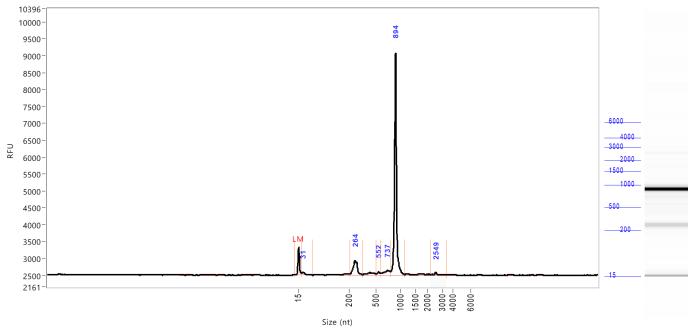


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Sample: B1_TP1 Well location: A1



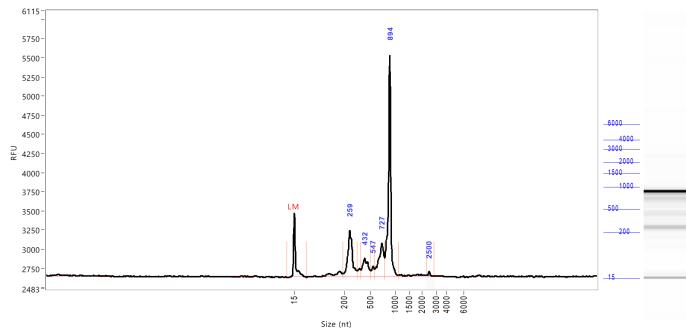
| | | | | | 31 |
|---|-----|----------------------|----------|---------|------|
| Р | eak | Concentration | From | То | RFU |
| | | (ng/uL) | (nt) | (nt) | |
| 1 | | 0.4293 | 1 | 25 | 820 |
| 2 | | 1.1378 | 25 | 66 | 81 |
| 3 | | 6.3699 | 204 | 350 | 436 |
| 4 | | 0.8719 | 501 | 598 | 107 |
| 5 | | 2.5165 | 598 | 797 | 138 |
| 6 | | 39.8194 | 797 | 1130 | 6574 |
| 7 | | 0.8683 | 2194 | 3348 | 80 |
| | | | | | |
| | | TIC. | E4 E040 | m m/u l | |
| | | TIC: | 51.5840 | ng/uL | |
| | | TIM: | 330.7559 | nmole/L | |
| | | Total concentration: | 56.4316 | ng/uL | |
| | | 28s/18s: | 0.0 | | |
| | | RQN | 1.3 | | |
| | | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                            Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: B1_TP2 Well location: A2

Created: Tuesday, December 08, 2020 4:16:51 PM

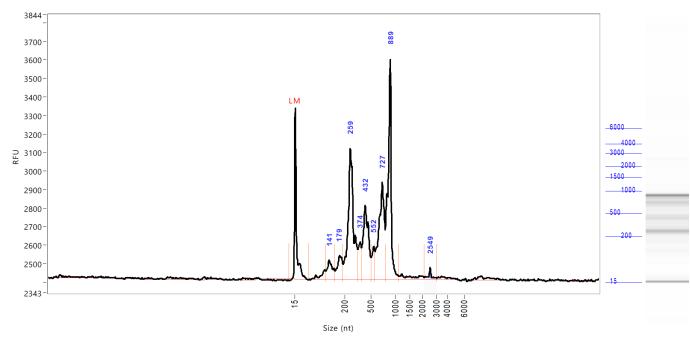


| | | | | SIZ |
|------|----------------------|----------|---------|------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 59 | 822 |
| 2 | 7.6623 | 196 | 345 | 599 |
| 3 | 3.5063 | 390 | 500 | 236 |
| 4 | 0.9431 | 500 | 575 | 131 |
| 5 | 5.5960 | 575 | 783 | 433 |
| 6 | 17.8036 | 783 | 1114 | 2890 |
| 7 | 0.3043 | 2323 | 2887 | 64 |
| | | | | |
| | | | | |
| | TIC: | 35.8156 | ng/uL | |
| | TIM: | 209.5143 | nmole/L | |
| | Total concentration: | 39.1348 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.2 | | |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: B1_TP3 Well location: A3



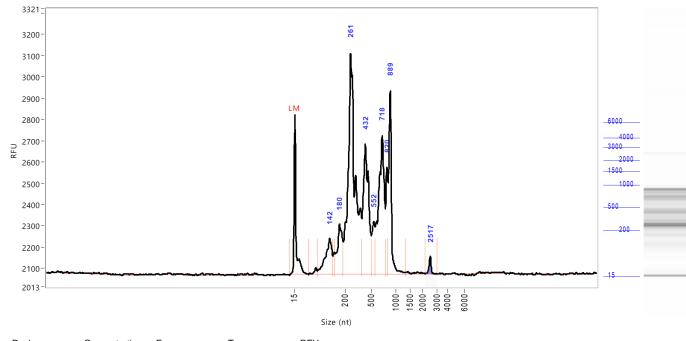
| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 64 | 922 |
| 2 | 1.4248 | 128 | 162 | 101 |
| 3 | 1.6712 | 162 | 192 | 125 |
| 4 | 9.6593 | 192 | 348 | 702 |
| 5 | 1.4882 | 348 | 390 | 199 |
| 6 | 5.1628 | 390 | 501 | 396 |
| 7 | 1.2548 | 501 | 580 | 176 |
| 8 | 6.4111 | 580 | 787 | 522 |
| 9 | 8.6765 | 787 | 1114 | 1187 |
| 10 | 0.3744 | 2114 | 3023 | 64 |
| | | | | |
| | TIC: | 36.1231 | ng/uL | |
| | TIM: | 289.3398 | nmole/L | |
| | Total concentration: | 37.4155 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.3 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: B1_TP4 Well location: A4

Created: Tuesday, December 08, 2020 4:16:51 PM



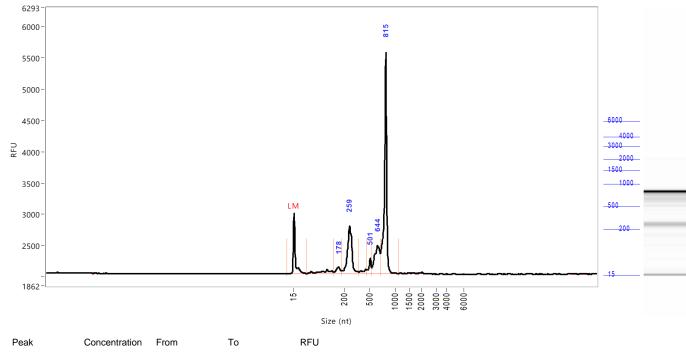
| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 65 | 748 |
| 2 | 3.5349 | 98 | 155 | 167 |
| 3 | 3.5934 | 163 | 193 | 232 |
| 4 | 22.0614 | 193 | 392 | 1037 |
| 5 | 9.5161 | 392 | 501 | 609 |
| 6 | 2.1872 | 501 | 580 | 245 |
| 7 | 9.9905 | 580 | 783 | 649 |
| 8 | 2.9552 | 783 | 843 | 502 |
| 9 | 6.1390 | 843 | 1355 | 860 |
| 10 | 0.3782 | 2162 | 3023 | 82 |
| | | | | |
| | TIC: | 60.3559 | ng/uL | |
| | TIM: | 546.0193 | nmole/L | |
| | Total concentration: | 60.5247 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: B1_PU_TP4 Well location: A5

Created: Tuesday, December 08, 2020 4:16:51 PM



| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 60 | 962 |
| 2 | 1.0982 | 161 | 189 | 105 |
| 3 | 8.1196 | 189 | 369 | 760 |
| 4 | 1.1669 | 463 | 529 | 248 |
| 5 | 5.1901 | 529 | 704 | 452 |
| 6 | 17.3410 | 704 | 1122 | 3531 |
| | | | | |
| | | | | |
| | TIC: | 32.9158 | ng/uL | |
| | TIM: | 217.6502 | nmole/L | |
| | Total concentration: | 35.3622 | ng/uL | |
| | 28s/18s: | 0.0 | | |

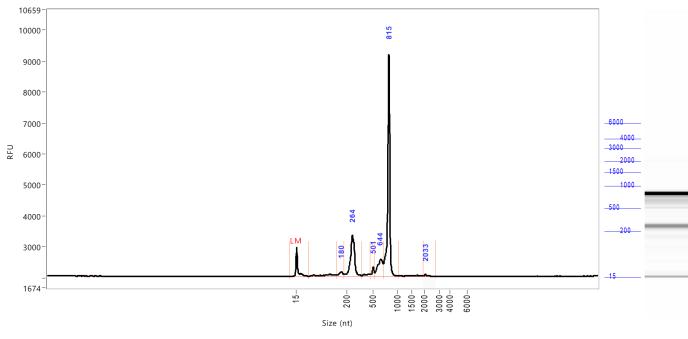
1.0

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

RQN

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Sample: B1_PU_TP3 **Well location:** A6



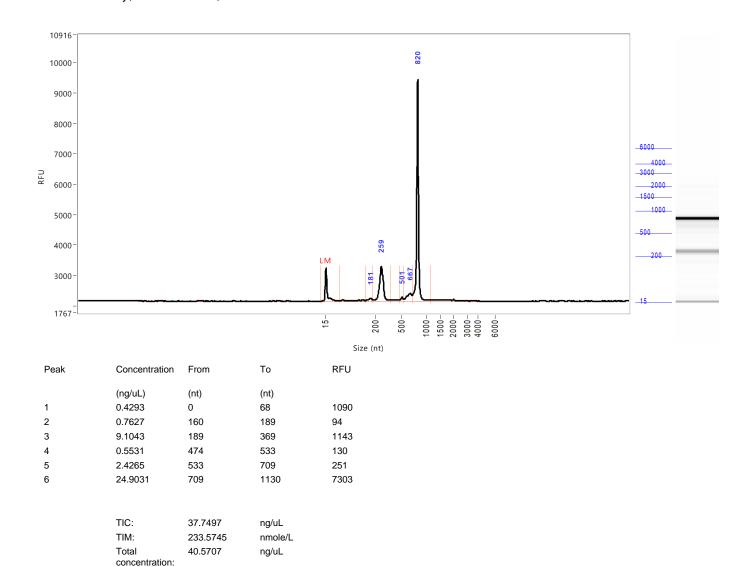
| Peak | Concentration | From | То | RFU |
|------|----------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 62 | 922 |
| 2 | 1.5453 | 162 | 190 | 146 |
| 3 | 13.8847 | 190 | 369 | 1314 |
| 4 | 1.3869 | 471 | 533 | 297 |
| 5 | 6.6196 | 533 | 704 | 554 |
| 6 | 33.8245 | 704 | 1009 | 7174 |
| 7 | 0.2458 | 1935 | 2791 | 59 |
| | | | | |
| | TIC: | 57.5068 | ng/uL | |
| | TIM: | 365.9356 | nmole/L | |
| | Total | 60.3305 | ng/uL | |
| | concentration: | 00.3303 | rig/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |
| | | | | |

```
Sample baseline V to V?: Y
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: B1_PU_TP2 Well location: A7

Created: Tuesday, December 08, 2020 4:16:51 PM



```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                           Sample baseline V to V?: Y
                                                                                         Sample baseline V to V points: 3
Sample filter: Binomial
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Manual baseline start (min): 15
                                   Manual baseline end (min): 38
Marker peak width (sec): 6
                             Marker min peak height: 100
                                                           Marker baseline V to V?: Y
                                                                                         Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                                                                  Dilution factor: 12.0
Quantification using: Ladder
                                     Final concentration (ng/uL): 8.0000
Minimum RFU for data processing: 2
```

28s/18s:

RQN

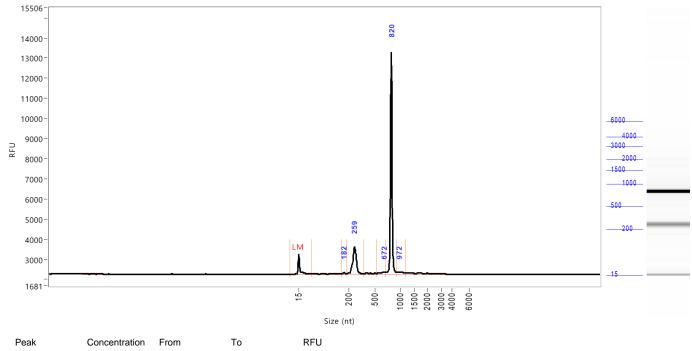
0.0

1.1

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Sample: B1_PU_TP1 Well location: A8

Created: Tuesday, December 08, 2020 4:16:51 PM



| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|-------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 62 | 991 |
| 2 | 0.5568 | 172 | 191 | 87 |
| 3 | 11.6952 | 191 | 369 | 1352 |
| 4 | 1.1521 | 524 | 704 | 103 |
| 5 | 37.0148 | 704 | 935 | 11044 |
| 6 | 1.2525 | 935 | 1194 | 95 |
| | | | | |
| | TIC: | 51.6715 | ng/uL | |
| | TIM: | 300.9556 | nmole/L | |
| | | | | |
| | Total concentration: | 56.4610 | ng/uL | |
| | 28s/18s: | 0.0 | | |

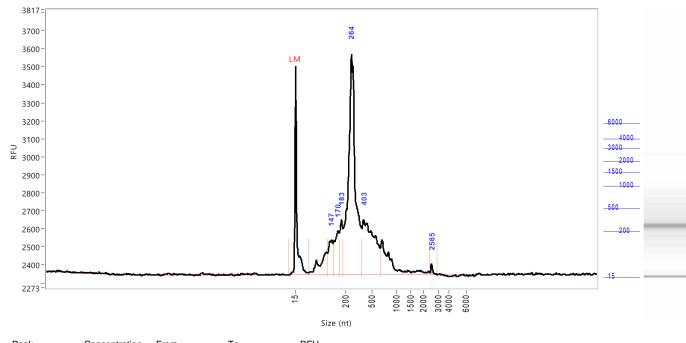
Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

RQN

1.1

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Sample: C1_TP4 Well location: A9



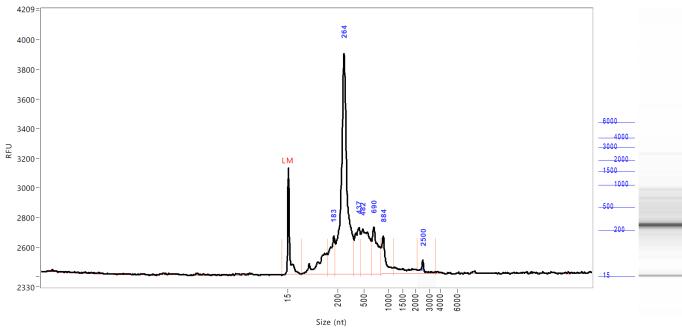
| Peak | Concentration | From | То | RFU |
|------|----------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 63 | 1152 |
| 2 | 1.7315 | 133 | 153 | 188 |
| 3 | 2.2538 | 153 | 175 | 237 |
| 4 | 2.0048 | 175 | 190 | 302 |
| 5 | 17.1241 | 190 | 382 | 1220 |
| 6 | 6.7711 | 382 | 663 | 303 |
| 7 | 0.1652 | 2404 | 2968 | 58 |
| | | | | |
| | TIC: | 30.0505 | ng/uL | |
| | TIM: | 355.1213 | nmole/L | |
| | Total | 34.7816 | ng/uL | |
| | concentration: | 04.7010 | Hg/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: C1_TP3 Well location: A10

Created: Tuesday, December 08, 2020 4:16:51 PM

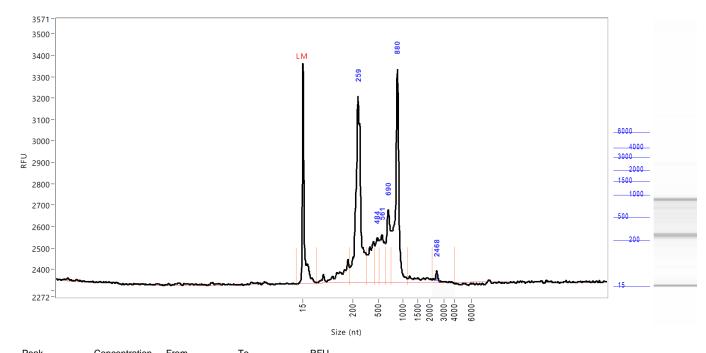


| | | | | 31Ze (1 |
|------|----------------------|----------|---------|---------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 63 | 717 |
| 2 | 4.2232 | 161 | 189 | 254 |
| 3 | 28.4402 | 189 | 382 | 1490 |
| 4 | 4.7920 | 382 | 455 | 308 |
| 5 | 7.6426 | 455 | 644 | 299 |
| 6 | 5.3043 | 644 | 838 | 315 |
| 7 | 3.0398 | 838 | 1162 | 256 |
| 8 | 0.7363 | 2114 | 3587 | 92 |
| | | | | |
| | TIC: | E4 4704 | n m/ul | |
| | TIC: | 54.1784 | ng/uL | |
| | TIM: | 518.0010 | nmole/L | |
| | Total concentration: | 60.0010 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.2 | | |

Sample baseline V to V?: Y Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Quantification using: Ladder Final concentration (ng/uL): 8.0000 Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: C1_TP2 Well location: A11



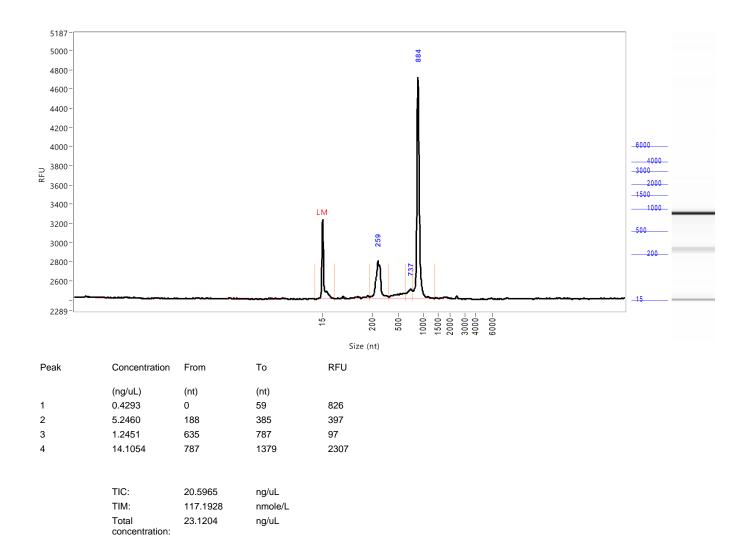
| Peak | Concentration | From | То | RFU |
|------|----------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 65 | 1025 |
| 2 | 10.0870 | 188 | 361 | 867 |
| 3 | 1.5335 | 455 | 501 | 208 |
| 4 | 2.4524 | 501 | 640 | 223 |
| 5 | 2.6653 | 640 | 750 | 337 |
| 6 | 7.2761 | 750 | 1154 | 993 |
| 7 | 0.2570 | 2146 | 4001 | 53 |
| | | | | |
| | | | | |
| | TIC: | 24.2712 | ng/uL | |
| | TIM: | 181.3849 | nmole/L | |
| | Total | 29.1705 | ng/uL | |
| | concentration: | | | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: C1_TP1 Well location: A12

Created: Tuesday, December 08, 2020 4:16:51 PM



```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                           Sample baseline V to V?: Y
                                                                                         Sample baseline V to V points: 3
Sample filter: Binomial
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Manual baseline start (min): 15
                                   Manual baseline end (min): 38
Marker peak width (sec): 6
                             Marker min peak height: 100
                                                           Marker baseline V to V?: Y
                                                                                         Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                                                                  Dilution factor: 12.0
Quantification using: Ladder
                                     Final concentration (ng/uL): 8.0000
Minimum RFU for data processing: 2
```

28s/18s:

RQN

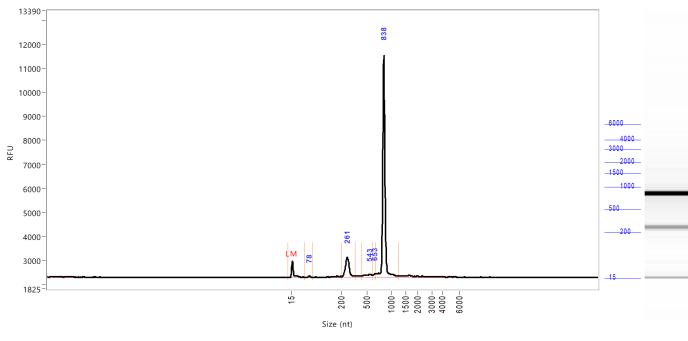
0.0

1.1

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Sample: C1_PU_TP1 Well location: B1

Created: Tuesday, December 08, 2020 4:16:51 PM



| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 61 | 674 |
| 2 | 0.5040 | 61 | 93 | 50 |
| 3 | 11.5586 | 200 | 361 | 847 |
| 4 | 2.7273 | 434 | 598 | 110 |
| 5 | 1.0328 | 598 | 658 | 149 |
| 6 | 65.3493 | 658 | 1234 | 9241 |
| | | | | |
| | TIC: | 81.1719 | ng/uL | |
| | TIM: | 420.8450 | nmole/L | |
| | Total concentration: | 87.9970 | ng/uL | |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

28s/18s:

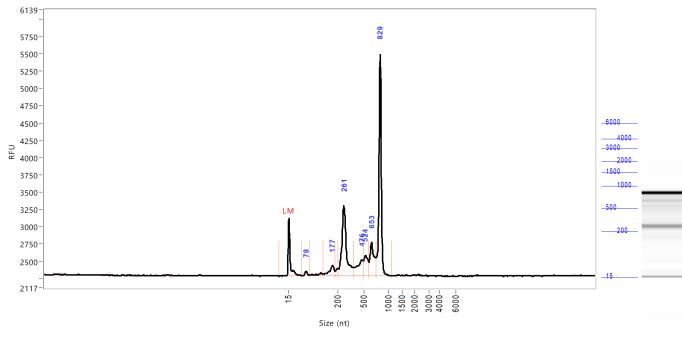
RQN

0.0

1.2

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Sample: C1_PU_TP2 Well location: B2



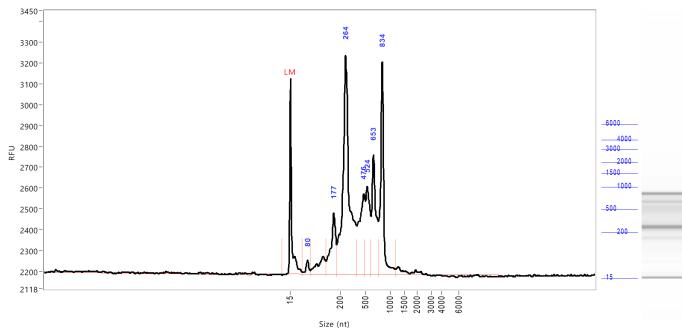
| | | | | () |
|------|----------------------|----------|---------|------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 63 | 835 |
| 2 | 0.4989 | 63 | 92 | 65 |
| 3 | 2.0632 | 143 | 187 | 144 |
| 4 | 13.0994 | 199 | 377 | 1010 |
| 5 | 3.6736 | 377 | 492 | 229 |
| 6 | 3.3384 | 492 | 598 | 293 |
| 7 | 5.0780 | 598 | 741 | 483 |
| 8 | 18.9689 | 741 | 1114 | 3202 |
| | | | | |
| | TIC: | 46.7204 | ng/uL | |
| | TIM: | 348.0854 | nmole/L | |
| | Total concentration: | 47.8440 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: C1_PU_TP3 Well location: B3

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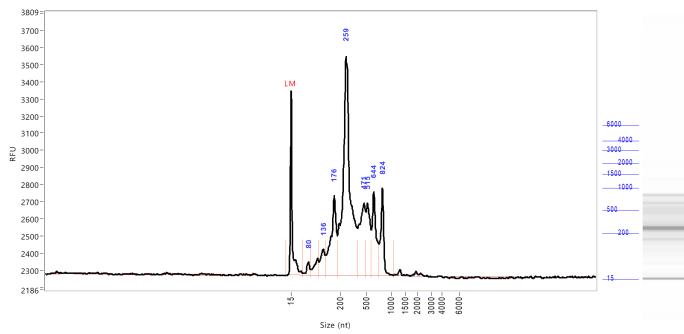


| | | | | * * |
|------|-----------------------------------|----------|---------|------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 60 | 932 |
| 2 | 0.5185 | 60 | 89 | 64 |
| 3 | 3.7069 | 148 | 188 | 292 |
| 4 | 16.7123 | 188 | 395 | 1047 |
| 5 | 5.2479 | 395 | 495 | 385 |
| 6 | 4.1506 | 495 | 607 | 418 |
| 7 | 5.2406 | 607 | 755 | 571 |
| 8 | 6.9494 | 755 | 1178 | 1015 |
| | | | | |
| | | | | |
| | TIC: | 42.5261 | ng/uL | |
| | TIM: | 386.0443 | nmole/L | |
| | Total concentration: | 44.3753 | ng/uL | |
| | 28s/18s· | 0.0 | | |
| | | | | |
| | concentration: 28s/18s: RQN | 0.0 | | |

Sample baseline V to V?: Y Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Upper marker selection: Last peak > 100 RFU Lower marker selection: First peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: C1_PU_TP4 Well location: B4

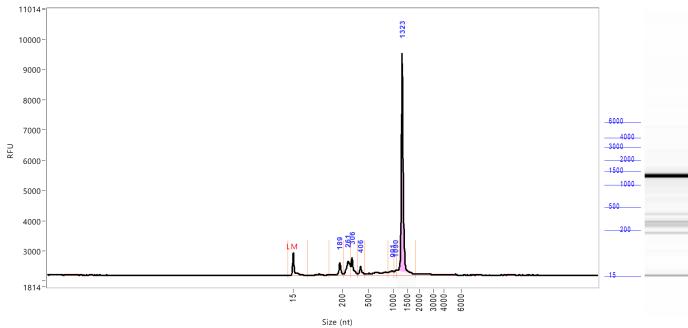


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 59 | 1073 |
| 2 | 0.5850 | 59 | 88 | 74 |
| 3 | 1.5480 | 120 | 143 | 151 |
| 4 | 5.7856 | 143 | 187 | 464 |
| 5 | 20.0390 | 187 | 395 | 1278 |
| 6 | 5.2094 | 395 | 492 | 421 |
| 7 | 3.5250 | 492 | 598 | 421 |
| 8 | 3.9599 | 598 | 755 | 486 |
| 9 | 2.9895 | 755 | 1073 | 508 |
| | | | | |
| | | | | |
| | TIC: | 43.6414 | ng/uL | |
| | TIM: | 481.9074 | nmole/L | |
| | Total concentration: | 44.5071 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: H1_TP1 Well location: B5

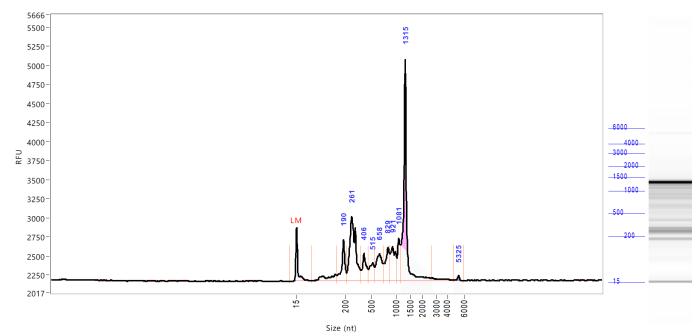


| | | | | 3 |
|------|----------------------|----------|---------|------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 68 | 724 |
| 2 | 3.2055 | 150 | 206 | 408 |
| 3 | 5.5084 | 206 | 290 | 467 |
| 4 | 4.4185 | 290 | 374 | 576 |
| 5 | 2.3790 | 374 | 463 | 298 |
| 6 | 1.8254 | 903 | 1033 | 152 |
| 7 | 0.9688 | 1033 | 1122 | 167 |
| 8 | 40.4910 | 1122 | 1843 | 7350 |
| | | | | |
| | TIC: | 58.7966 | ng/uL | |
| | TIM: | 284.8129 | nmole/L | |
| | Total concentration: | 66.3850 | ng/uL | |
| | | | | |
| | 28s/18s: | 0.0 | | |
| | RQN | 7.4 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: H1_TP2 Well location: B6



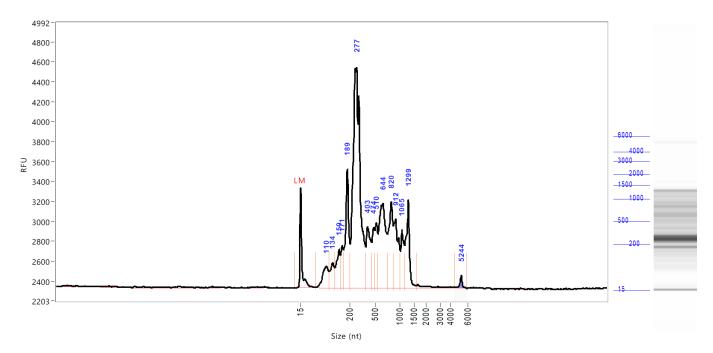
| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 70 | 698 |
| 2 | 5.6896 | 165 | 204 | 538 |
| 3 | 16.9734 | 204 | 371 | 841 |
| 4 | 4.5119 | 371 | 458 | 355 |
| 5 | 3.3729 | 458 | 552 | 227 |
| 6 | 6.9229 | 552 | 750 | 349 |
| 7 | 4.8513 | 750 | 866 | 441 |
| 8 | 6.8586 | 866 | 1017 | 449 |
| 9 | 4.3434 | 1017 | 1146 | 552 |
| 10 | 23.5212 | 1146 | 2646 | 2903 |
| 11 | 0.3138 | 4730 | 5946 | 59 |
| | TIC: | 77.3590 | ng/uL | |
| | TIM: | 481.3533 | nmole/L | |
| | Total concentration: | 79.9096 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 4.0 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: H1_TP3 Well location: B7

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| Peak | Concentration | From | То | RFU |
|------|----------------------|-----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 72 | 999 |
| 2 | 3.2121 | 72 | 121 | 210 |
| 3 | 2.6340 | 121 | 143 | 249 |
| 4 | 3.5857 | 143 | 163 | 384 |
| 5 | 2.7622 | 163 | 176 | 422 |
| 6 | 9.4550 | 176 | 201 | 1185 |
| 7 | 36.5302 | 201 | 382 | 2210 |
| 8 | 6.0839 | 382 | 453 | 611 |
| 9 | 3.2937 | 453 | 487 | 603 |
| 10 | 4.1168 | 487 | 543 | 649 |
| 11 | 12.1620 | 543 | 741 | 842 |
| 12 | 7.2563 | 741 | 857 | 860 |
| 13 | 7.0957 | 857 | 1001 | 686 |
| 14 | 3.5620 | 1001 | 1154 | 582 |
| 15 | 6.0419 | 1154 | 1584 | 880 |
| 16 | 0.4968 | 4406 | 5892 | 128 |
| | | | | |
| | | | | |
| | TIC: | 108.2882 | ng/uL | |
| | TIM: | 1071.7787 | nmole/L | |
| | Total concentration: | 106.6652 | ng/uL | |

0.0

1.1

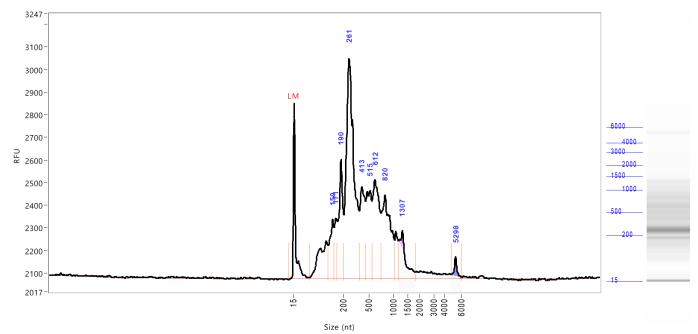
```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

28s/18s:

RQN

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Sample: H1_TP4 Well location: B8

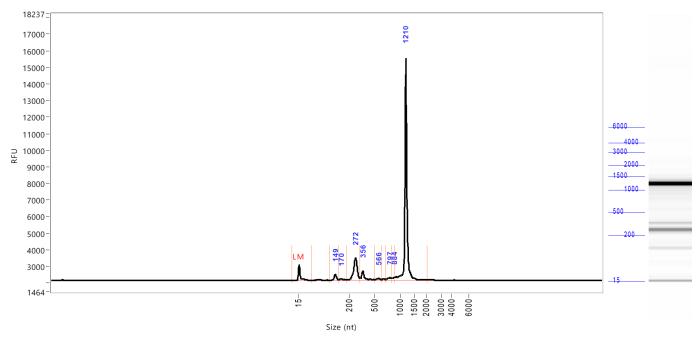


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|-----|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 71 | 773 |
| 2 | 3.1752 | 142 | 164 | 259 |
| 3 | 2.3899 | 164 | 177 | 264 |
| 4 | 6.0390 | 177 | 201 | 523 |
| 5 | 21.8567 | 201 | 385 | 972 |
| 6 | 5.5610 | 385 | 458 | 403 |
| 7 | 5.5176 | 458 | 552 | 385 |
| 8 | 7.9030 | 552 | 741 | 436 |
| 9 | 8.0731 | 741 | 1025 | 367 |
| 10 | 3.1482 | 1170 | 1806 | 209 |
| 11 | 0.5962 | 4757 | 6053 | 94 |
| | | | | |
| | | | | |
| | TIC: | 64.2599 | ng/uL | |
| | TIM: | 599.5573 | nmole/L | |
| | Total concentration: | 70.9224 | ng/uL | |
| | 28s/18s: | 1.1 | | |
| | RQN | 1.7 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: H1_PU_TP1 Well location: B9



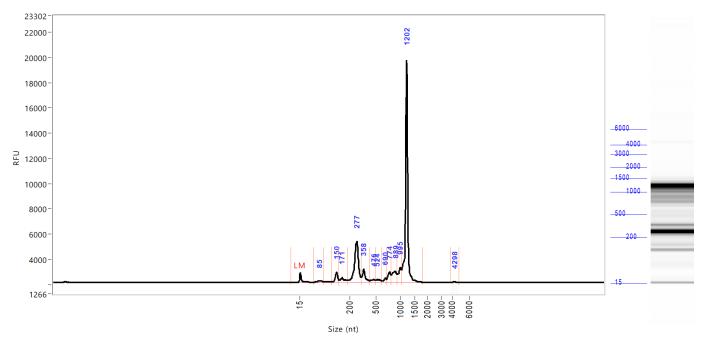
| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|-------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 61 | 940 |
| 2 | 2.3408 | 128 | 160 | 349 |
| 3 | 1.1540 | 160 | 190 | 97 |
| 4 | 12.6841 | 190 | 324 | 1343 |
| 5 | 4.5362 | 324 | 495 | 562 |
| 6 | 1.0815 | 495 | 626 | 116 |
| 7 | 1.3191 | 709 | 829 | 157 |
| 8 | 0.8839 | 829 | 889 | 183 |
| 9 | 58.5678 | 889 | 2017 | 13408 |
| | | | | |
| | TIO: | 00 5074 | | |
| | TIC: | 82.5674 | ng/uL | |
| | TIM: | 422.4011 | nmole/L | |
| | Total concentration: | 84.2716 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: H1_PU_TP2 Well location: B10

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| Peak | Concentration | From | То | RFU |
|------|---------------|-----------|---------|-------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 63 | 765 |
| 2 | 1.9665 | 63 | 102 | 110 |
| 3 | 6.7668 | 132 | 159 | 813 |
| 4 | 5.2639 | 159 | 192 | 375 |
| 5 | 39.5006 | 192 | 327 | 3221 |
| 6 | 9.3111 | 327 | 424 | 1030 |
| 7 | 2.7240 | 424 | 497 | 231 |
| 8 | 2.6453 | 497 | 612 | 212 |
| 9 | 2.4972 | 612 | 714 | 345 |
| 10 | 6.6186 | 714 | 806 | 798 |
| 11 | 11.0149 | 806 | 935 | 875 |
| 12 | 9.2835 | 935 | 1033 | 1170 |
| 13 | 94.3242 | 1033 | 1824 | 17613 |
| 14 | 0.3139 | 3891 | 4811 | 58 |
| | | | | |
| | | | | |
| | TIC: | 192.2305 | ng/uL | |
| | TIM: | 1232.5111 | nmole/L | |

192.5250

0.0

1.1

ng/uL

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

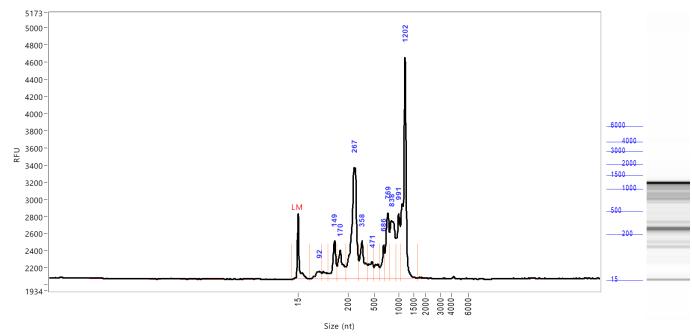
Total

RQN

concentration: 28s/18s:

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Sample: H1_PU_TP3 Well location: B11

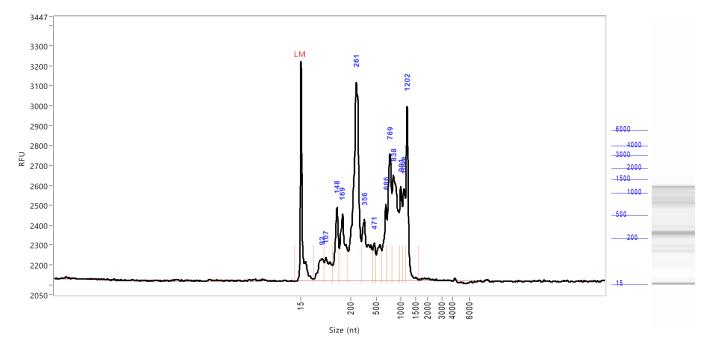


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 58 | 754 |
| 2 | 1.6475 | 58 | 102 | 83 |
| 3 | 4.6967 | 126 | 158 | 444 |
| 4 | 4.9913 | 158 | 191 | 332 |
| 5 | 20.5096 | 191 | 322 | 1297 |
| 6 | 5.7821 | 322 | 424 | 440 |
| 7 | 2.6436 | 424 | 492 | 200 |
| 8 | 3.0768 | 607 | 709 | 389 |
| 9 | 6.9822 | 709 | 806 | 762 |
| 10 | 9.2202 | 806 | 931 | 678 |
| 11 | 6.3888 | 931 | 1033 | 753 |
| 12 | 19.7792 | 1033 | 1677 | 2579 |
| | | | | |
| | | | | |
| | TIC: | 85.7181 | ng/uL | |
| | TIM: | 711.8967 | nmole/L | |
| | Total concentration: | 88.0745 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                            Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                  Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: H1_PU_TP4 Well location: B12

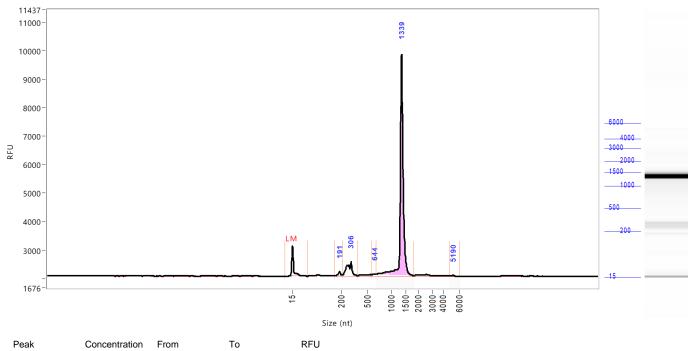


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 61 | 1099 |
| 2 | 1.4494 | 61 | 101 | 108 |
| 3 | 1.5191 | 101 | 132 | 113 |
| 4 | 2.7291 | 132 | 157 | 365 |
| 5 | 3.6819 | 157 | 190 | 331 |
| 6 | 11.4210 | 190 | 322 | 993 |
| 7 | 4.0906 | 322 | 453 | 305 |
| 8 | 1.0163 | 453 | 492 | 187 |
| 9 | 2.2985 | 603 | 709 | 382 |
| 10 | 4.0987 | 709 | 806 | 634 |
| 11 | 5.3767 | 806 | 958 | 526 |
| 12 | 2.2337 | 958 | 1033 | 473 |
| 13 | 1.9863 | 1033 | 1138 | 459 |
| 14 | 3.7687 | 1138 | 1630 | 874 |
| | | | | |
| | TIC: | 45.6700 | ng/uL | |
| | TIM: | 466.5893 | nmole/L | |
| | Total concentration: | 46.3314 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                         Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A2_TP1 Well location: C1

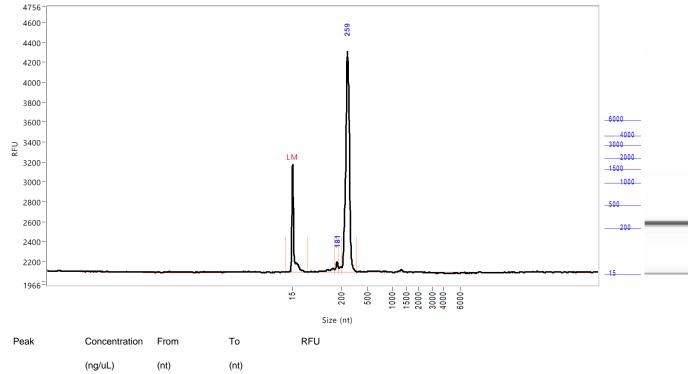


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 70 | 1049 |
| 2 | 1.0118 | 171 | 204 | 172 |
| 3 | 5.8104 | 204 | 379 | 500 |
| 4 | 0.5083 | 584 | 667 | 74 |
| 5 | 44.7802 | 667 | 1806 | 7794 |
| 6 | 0.1714 | 4703 | 6027 | 47 |
| | | | | |
| | TIC: | 52.2820 | ng/uL | |
| | TIM: | 191.4086 | nmole/L | |
| | Total concentration: | 55.9423 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 9.3 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A2_TP2 Well location: C2

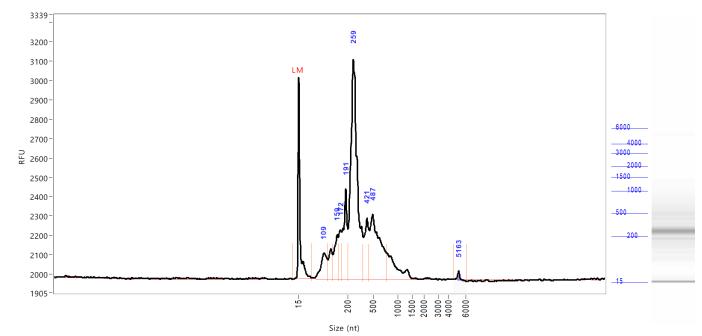


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 70 | 1079 |
| 2 | 0.4885 | 172 | 187 | 103 |
| 3 | 17.6292 | 199 | 371 | 2214 |
| | | | | |
| | | | | |
| | TIC: | 18.1177 | ng/uL | |
| | TIM: | 221.2334 | nmole/L | |
| | Total concentration: | 19.1056 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                         Sample baseline V to V?: Y
                                                                                       Sample baseline V to V points: 3
                            Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                    Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A2_TP3 Well location: C3

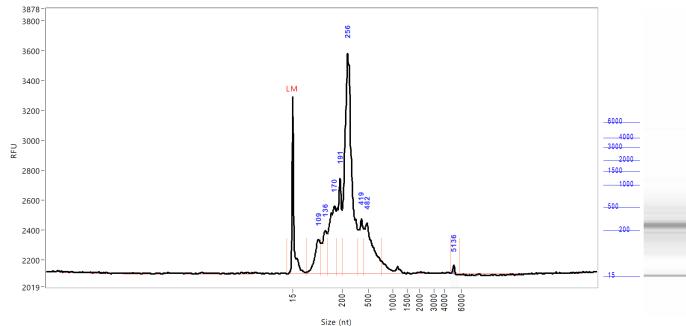


| Peak | Concentration | From | То | RFU | |
|------|----------------------|----------|---------|------|--|
| | (ng/uL) | (nt) | (nt) | | |
| 1 | 0.4293 | 0 | 63 | 1037 | |
| 2 | 2.0887 | 63 | 125 | 132 | |
| 3 | 2.0143 | 143 | 162 | 228 | |
| 4 | 1.7697 | 162 | 176 | 252 | |
| 5 | 4.0760 | 176 | 200 | 461 | |
| 6 | 16.8035 | 200 | 379 | 1132 | |
| 7 | 2.9623 | 379 | 450 | 314 | |
| 8 | 6.9811 | 450 | 769 | 332 | |
| 9 | 0.1217 | 4487 | 6160 | 41 | |
| | TIC: | 36.8172 | ng/uL | | |
| | TIM: | 452.9326 | nmole/L | | |
| | Total concentration: | 40.1275 | ng/uL | | |
| | 28s/18s: | 0.0 | | | |
| | RQN | 1.0 | | | |
| | | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A2_TP4 Well location: C4

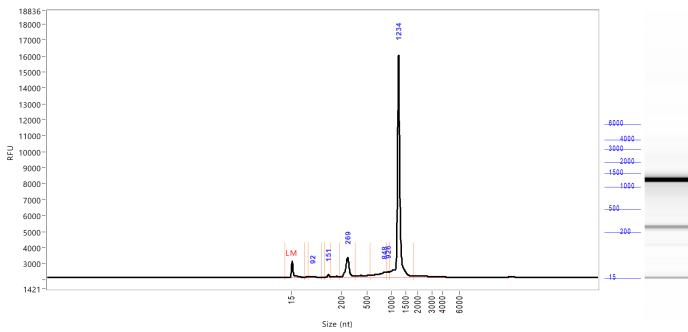


| | | | | Size (nt) |
|------|----------------------|----------|---------|-----------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 65 | 1178 |
| 2 | 2.7963 | 65 | 120 | 222 |
| 3 | 2.8578 | 120 | 142 | 283 |
| 4 | 6.0932 | 142 | 176 | 447 |
| 5 | 5.1602 | 176 | 198 | 633 |
| 6 | 20.2717 | 198 | 377 | 1470 |
| 7 | 3.3282 | 377 | 453 | 363 |
| 8 | 5.1819 | 453 | 769 | 333 |
| 9 | 0.1215 | 4730 | 5784 | 51 |
| | | | | |
| | | | | |
| | TIC: | 45.8109 | ng/uL | |
| | TIM: | 642.4351 | nmole/L | |
| | Total concentration: | 46.0897 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                         Sample baseline V to V?: Y
                                                                                       Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
                                                          Upper marker selection: Last peak > 100 RFU
Lower marker selection: First peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A2_PU_TP1 Well location: C5

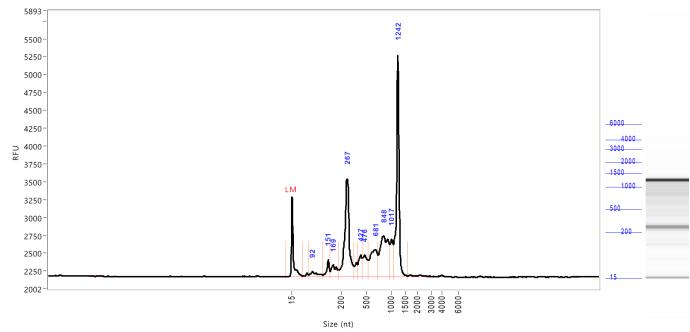


| Peak | Concentration | From | То | RFU |
|------|----------------|----------|---------|-------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 62 | 1039 |
| 2 | 0.6918 | 76 | 125 | 55 |
| 3 | 0.8887 | 136 | 159 | 169 |
| 4 | 10.5080 | 192 | 358 | 1237 |
| 5 | 5.8247 | 556 | 889 | 321 |
| 6 | 1.7962 | 889 | 949 | 351 |
| 7 | 66.3784 | 949 | 1824 | 13921 |
| | | | | |
| | TIC: | 86.0877 | ng/uL | |
| | TIM: | 361.4707 | nmole/L | |
| | Total | 93.5159 | ng/uL | |
| | concentration: | | | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.3 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A2_PU_TP2 Well location: C6

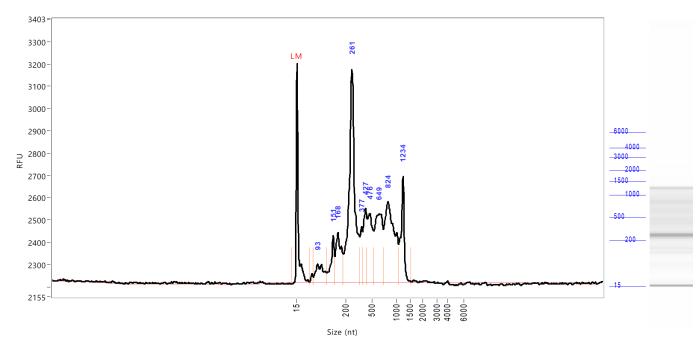


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 56 | 1118 |
| 2 | 1.1128 | 77 | 131 | 72 |
| 3 | 1.3932 | 131 | 159 | 241 |
| 4 | 1.7857 | 159 | 188 | 164 |
| 5 | 13.1692 | 188 | 350 | 1372 |
| 6 | 2.0933 | 392 | 445 | 303 |
| 7 | 2.8599 | 445 | 529 | 299 |
| 8 | 5.3022 | 529 | 732 | 378 |
| 9 | 9.0559 | 732 | 972 | 568 |
| 10 | 2.9578 | 972 | 1081 | 525 |
| 11 | 14.4047 | 1081 | 1593 | 3099 |
| | TIC: | 54.1347 | ng/uL | |
| | TIM: | 390.2806 | nmole/L | |
| | Total concentration: | 55.1776 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A2_PU_TP3 Well location: C7

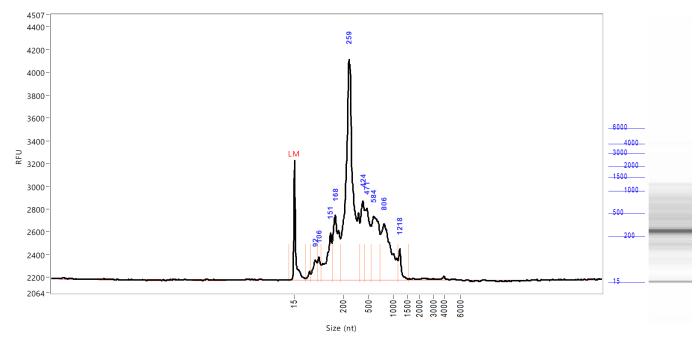


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|-----|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 62 | 984 |
| 2 | 1.7598 | 76 | 126 | 82 |
| 3 | 1.8698 | 126 | 157 | 209 |
| 4 | 2.9737 | 157 | 187 | 224 |
| 5 | 13.4106 | 187 | 356 | 955 |
| 6 | 1.4819 | 356 | 392 | 252 |
| 7 | 2.7845 | 392 | 445 | 331 |
| 8 | 3.5631 | 445 | 533 | 308 |
| 9 | 5.1876 | 533 | 727 | 307 |
| 10 | 7.6958 | 727 | 1073 | 361 |
| 11 | 3.2582 | 1073 | 1529 | 476 |
| | | | | |
| | | | | |
| | TIC: | 43.9850 | ng/uL | |
| | TIM: | 424.1686 | nmole/L | |
| | Total concentration: | 43.6480 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |
| | | | | |

```
Sample baseline V to V?: Y
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                                                       Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A2_PU_TP4 Well location: C8

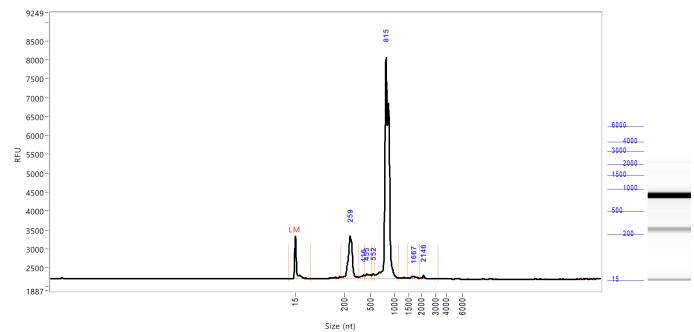


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 57 | 1044 |
| 2 | 1.6742 | 77 | 100 | 170 |
| 3 | 1.5442 | 100 | 116 | 198 |
| 4 | 4.9257 | 116 | 157 | 411 |
| 5 | 7.4426 | 157 | 187 | 568 |
| 6 | 32.1336 | 187 | 390 | 1936 |
| 7 | 5.4156 | 390 | 445 | 693 |
| 8 | 6.6979 | 445 | 533 | 623 |
| 9 | 8.5534 | 533 | 727 | 556 |
| 10 | 9.7041 | 727 | 1162 | 488 |
| 11 | 1.3998 | 1162 | 1538 | 271 |
| | | | | |
| | TIO | 70.4044 | | |
| | TIC: | 79.4911 | ng/uL | |
| | TIM: | 877.5856 | nmole/L | |
| | Total concentration: | 79.0091 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: F2_TP1 Well location: C9



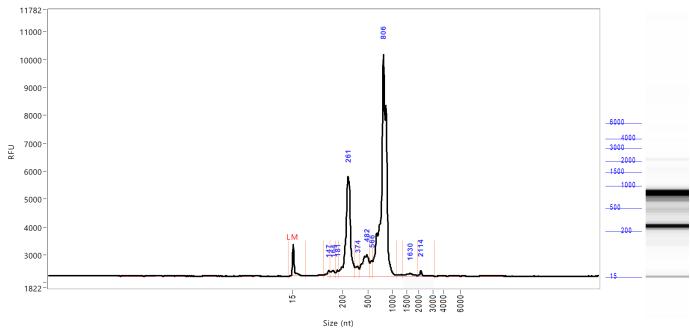
| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 72 | 1121 |
| 2 | 10.0741 | 187 | 364 | 1131 |
| 3 | 0.7658 | 364 | 432 | 99 |
| 4 | 1.2244 | 432 | 515 | 129 |
| 5 | 0.6776 | 515 | 589 | 134 |
| 6 | 43.3604 | 589 | 1138 | 5872 |
| 7 | 0.7176 | 1436 | 1917 | 64 |
| 8 | 0.4324 | 1917 | 3240 | 98 |
| | | | | |
| | | | | |
| | TIC: | 57.2523 | ng/uL | |
| | TIM: | 303.4922 | nmole/L | |
| | Total concentration: | 58.3521 | ng/uL | |
| | 28s/18s: | 0.7 | | |
| | RQN | 1.2 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: F2_TP2 Well location: C10

Created: Tuesday, December 08, 2020 4:16:51 PM

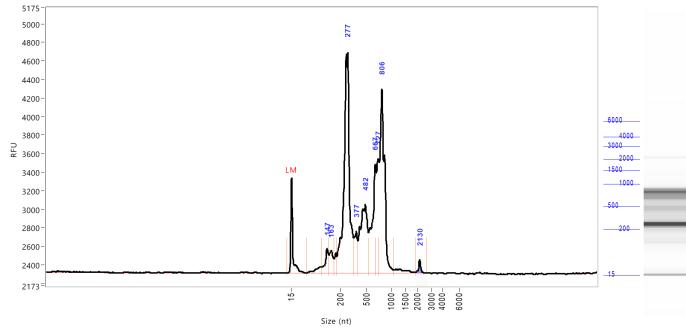


| Peak | Concentration | From | То | RFU |
|------|----------------|----------|------------------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 63 | 1123 |
| 2 | 1.5219 | 127 | 155 | 198 |
| 3 | 1.5917 | 155 | 173 | 192 |
| 4 | 1.0175 | 173 | 185 | 214 |
| 5 | 35.0361 | 185 | 345 | 3567 |
| 6 | 2.4060 | 345 | 392 | 349 |
| 7 | 10.0140 | 392 | 524 | 763 |
| 8 | 2.8069 | 524 | 584 | 549 |
| 9 | 77.9570 | 584 | 1138 | 7950 |
| 10 | 1.2143 | 1355 | 1926 | 85 |
| 11 | 0.8050 | 1926 | 3131 | 212 |
| | | | | |
| | TIC: | 134.3704 | ng/uL | |
| | TIM: | 912.3014 | nmole/L | |
| | Total | 135.0059 | nmoie/L ng/uL | |
| | concentration: | 133.0059 | rig/uL | |
| | 28s/18s: | 1.2 | | |
| | RQN | 1.2 | | |
| | | | | |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: F2_TP3 Well location: C11

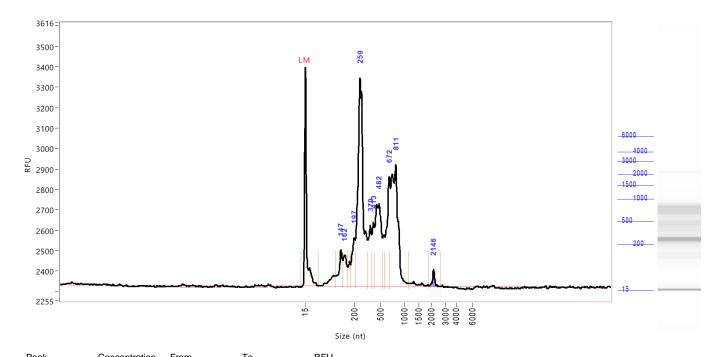


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 70 | 1022 |
| 2 | 2.2439 | 126 | 154 | 261 |
| 3 | 2.1755 | 154 | 173 | 238 |
| 4 | 29.5894 | 183 | 343 | 2383 |
| 5 | 3.4337 | 343 | 395 | 450 |
| 6 | 11.0727 | 395 | 524 | 740 |
| 7 | 10.0588 | 524 | 690 | 1169 |
| 8 | 6.6755 | 690 | 750 | 1237 |
| 9 | 17.7978 | 750 | 1090 | 1984 |
| 10 | 0.5775 | 1917 | 2646 | 149 |
| | | | | |
| | | | | |
| | TIC: | 83.6248 | ng/uL | |
| | TIM: | 698.9614 | nmole/L | |
| | Total concentration: | 84.9708 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: F2_TP4 Well location: C12



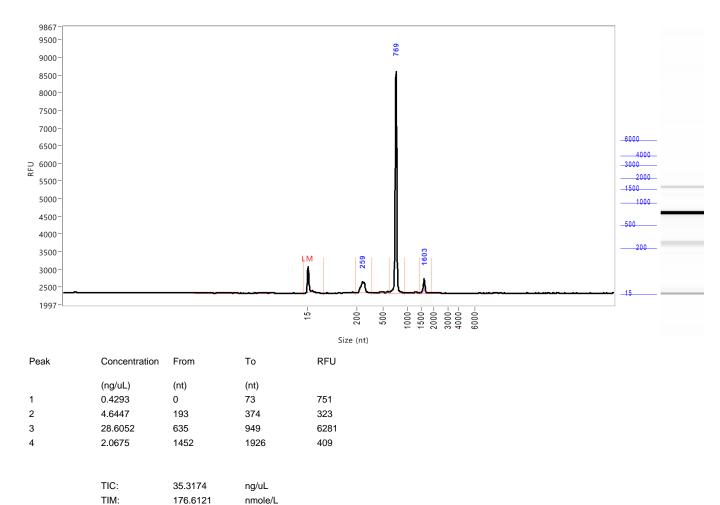
| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 64 | 1069 |
| 2 | 1.4741 | 126 | 155 | 175 |
| 3 | 1.3394 | 155 | 173 | 150 |
| 4 | 1.7946 | 184 | 204 | 236 |
| 5 | 11.9271 | 204 | 350 | 1015 |
| 6 | 1.9602 | 350 | 395 | 297 |
| 7 | 1.5517 | 395 | 427 | 311 |
| 8 | 5.0717 | 427 | 533 | 404 |
| 9 | 3.6157 | 589 | 695 | 533 |
| 10 | 7.9392 | 695 | 1114 | 593 |
| 11 | 0.3098 | 1898 | 2678 | 82 |
| | | | | |
| | | | | |
| | TIC: | 36.9835 | ng/uL | |
| | TIM: | 336.0659 | nmole/L | |
| | Total concentration: | 38.8941 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: F2_PU_TP1 Well location: D1

Created: Tuesday, December 08, 2020 4:16:51 PM

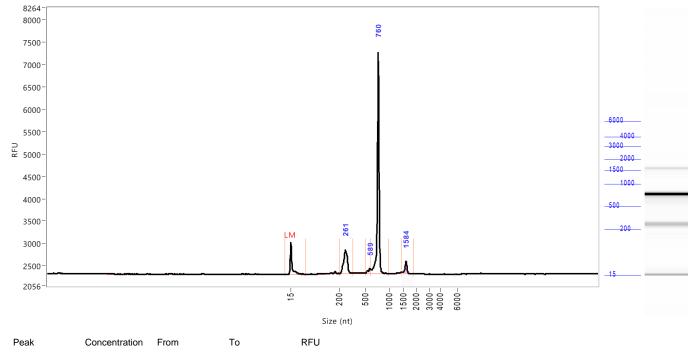


```
Total 37.3207 ng/uL concentration: 28s/18s: 0.0 RQN 1.6
```

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Sample filter: Binomial Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Dilution factor: 12.0 Quantification using: Ladder Final concentration (ng/uL): 8.0000 Minimum RFU for data processing: 2

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Sample: F2_PU_TP2 Well location: D2

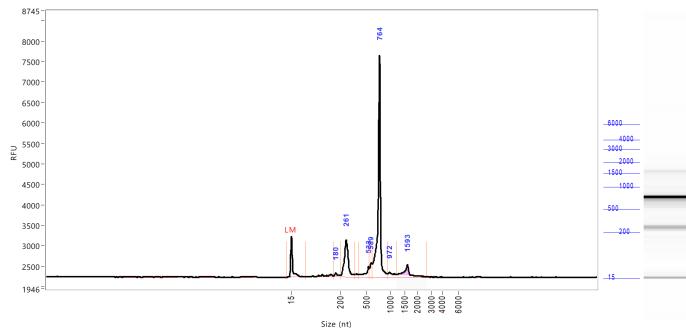


| | | | | 312 |
|------|----------------------|----------|---------|------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 73 | 703 |
| 2 | 7.1679 | 198 | 353 | 541 |
| 3 | 1.2244 | 501 | 617 | 124 |
| 4 | 27.0966 | 617 | 977 | 4953 |
| 5 | 1.8476 | 1428 | 1898 | 282 |
| | | | | |
| | TIC: | 37.3365 | ng/uL | |
| | TIM: | 208.0810 | nmole/L | |
| | Total concentration: | 39.5939 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.5 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: F2_PU_TP3 Well location: D3



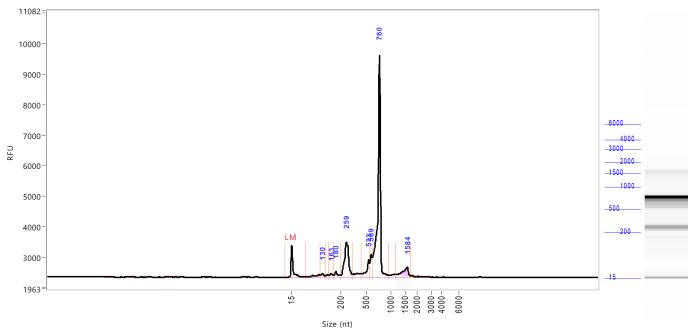
| | | | | • , , |
|------|----------------------|----------|---------|-------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 69 | 985 |
| 2 | 0.8448 | 172 | 199 | 102 |
| 3 | 8.5915 | 199 | 361 | 903 |
| 4 | 2.1419 | 406 | 556 | 259 |
| 5 | 1.6647 | 556 | 612 | 355 |
| 6 | 27.3657 | 612 | 935 | 5427 |
| 7 | 1.3621 | 935 | 1202 | 124 |
| 8 | 3.7502 | 1202 | 2678 | 302 |
| | | | | |
| | TIC: | 45.7209 | ng/uL | |
| | TIM: | 264.6649 | nmole/L | |
| | Total concentration: | 47.7004 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.8 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: F2_PU_TP4 Well location: D4

Created: Tuesday, December 08, 2020 4:16:51 PM



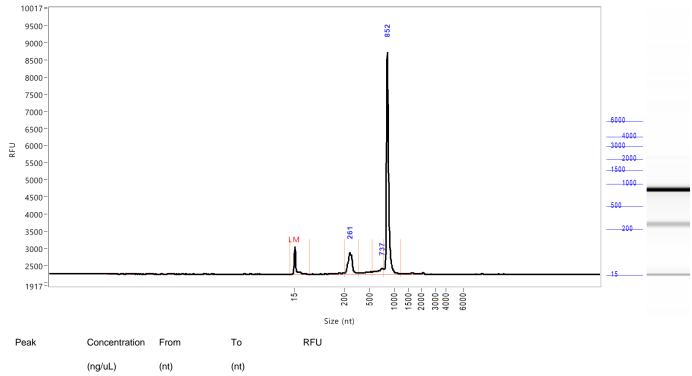
| | | | | 3 |
|------|----------------|----------|----------|------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 69 | 1039 |
| 2 | 0.8515 | 122 | 140 | 122 |
| 3 | 0.9057 | 153 | 172 | 117 |
| 4 | 1.3665 | 172 | 199 | 178 |
| 5 | 11.1540 | 199 | 332 | 1153 |
| 6 | 3.4869 | 437 | 552 | 579 |
| 7 | 3.2952 | 552 | 612 | 743 |
| 8 | 37.3718 | 612 | 949 | 7283 |
| 9 | 4.0366 | 1162 | 1732 | 335 |
| | | | | |
| | TIC: | 62.4681 | ng/uL | |
| | TIM: | 399.7671 | nmole/L | |
| | Total | 67.0849 | ng/uL | |
| | concentration: | | <i>y</i> | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.7 | | |
| | | | | |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: G2_TP1 Well location: D5

Created: Tuesday, December 08, 2020 4:16:51 PM



| (ng/uL) | (nt) | (nt) | |
|----------------------|----------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 0.4293 | 0 | 70 | 789 |
| 8.1681 | 200 | 364 | 629 |
| 2.8101 | 538 | 774 | 170 |
| 36.3057 | 774 | 1194 | 6469 |
| | | | |
| | | | |
| TIC: | 47.2839 | ng/uL | |
| TIM: | 241.0374 | nmole/L | |
| Total concentration: | 50.9038 | ng/uL | |
| 28s/18s: | 0.0 | | |
| | 0.4293 8.1681 2.8101 36.3057 TIC: TIM: Total concentration: | 0.4293 0 8.1681 200 2.8101 538 36.3057 774 TIC: 47.2839 TIM: 241.0374 Total 50.9038 concentration: | 0.4293 0 70 8.1681 200 364 2.8101 538 774 36.3057 774 1194 TIC: 47.2839 ng/uL TIM: 241.0374 nmole/L Total 50.9038 ng/uL concentration: |

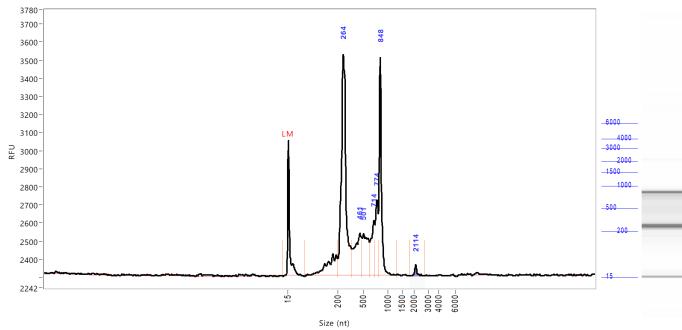
Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

RQN

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Sample: G2_TP2 Well location: D6

Created: Tuesday, December 08, 2020 4:16:51 PM



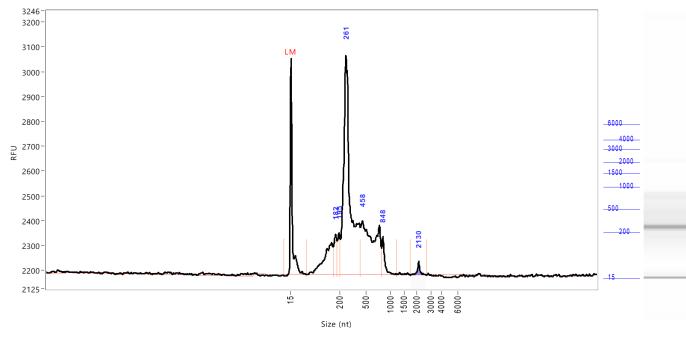
| | | | | - |
|------|----------------|----------|---------|------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 75 | 743 |
| 2 | 17.9195 | 200 | 364 | 1221 |
| 3 | 5.0057 | 364 | 487 | 229 |
| 4 | 3.7856 | 487 | 621 | 225 |
| 5 | 3.3212 | 621 | 737 | 300 |
| 6 | 2.9956 | 737 | 801 | 413 |
| 7 | 7.7005 | 801 | 1307 | 1203 |
| 8 | 0.2211 | 1834 | 2774 | 54 |
| | | | | |
| | TIC: | 40.9492 | ng/uL | |
| | TIM: | 321.2073 | nmole/L | |
| | Total | 44.1345 | ng/uL | |
| | concentration: | 77.1070 | rig/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: G2_TP3 Well location: D7

Created: Tuesday, December 08, 2020 4:16:51 PM



| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|-----|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 73 | 867 |
| 2 | 1.4684 | 172 | 188 | 161 |
| 3 | 1.2383 | 188 | 200 | 166 |
| 4 | 16.1596 | 200 | 434 | 882 |
| 5 | 7.4951 | 434 | 820 | 215 |
| 6 | 0.9868 | 820 | 1234 | 152 |
| 7 | 0.2373 | 1750 | 2710 | 51 |
| | | | | |
| | TIC: | 27.5854 | ng/uL | |
| | TIM: | 258.9176 | nmole/L | |
| | Total concentration: | 30.5696 | ng/uL | |
| | 28s/18s: | 0.0 | | |

Sample baseline V to V?: Y Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Upper marker selection: Last peak > 100 RFU Lower marker selection: First peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

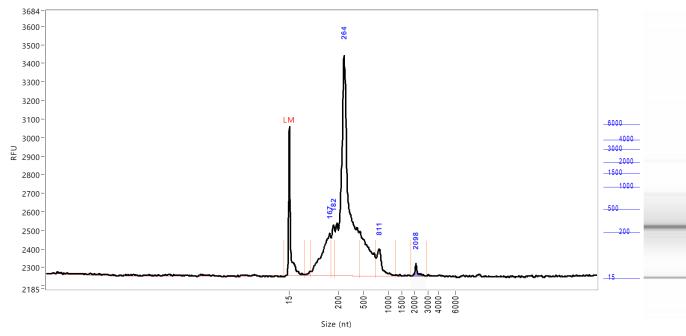
RQN

1.1

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Sample: G2_TP4 Well location: D8

Created: Tuesday, December 08, 2020 4:16:51 PM



| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 73 | 801 |
| 2 | 6.2088 | 98 | 172 | 227 |
| 3 | 2.6968 | 172 | 187 | 270 |
| 4 | 26.1930 | 187 | 448 | 1186 |
| 5 | 1.8194 | 750 | 1259 | 142 |
| 6 | 0.3259 | 1871 | 2887 | 64 |
| | | | | |
| | TIC: | 37.2440 | ng/uL | |
| | TIM: | 469.4200 | nmole/L | |
| | Total concentration: | 43.1488 | ng/uL | |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

28s/18s:

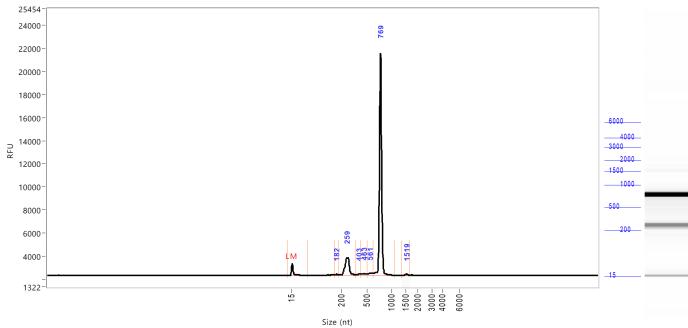
RQN

0.0

1.1

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Sample: G2_PU_TP1 **Well location:** D9

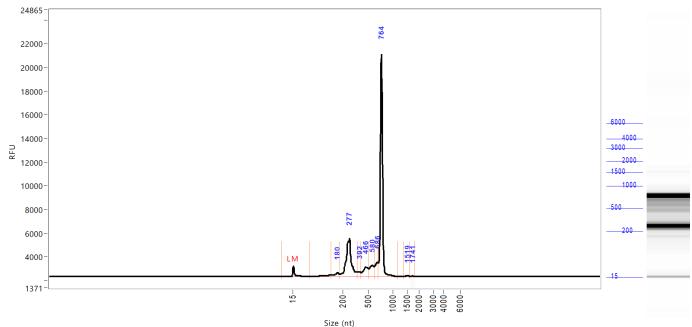


| F | Peak | Concentration | From | То | RFU |
|---|------|----------------------|----------|---------|-------|
| | | (ng/uL) | (nt) | (nt) | |
| 1 | 1 | 0.4293 | 0 | 73 | 1047 |
| 2 | 2 | 0.6366 | 172 | 190 | 100 |
| 3 | 3 | 16.0597 | 190 | 366 | 1566 |
| 4 | 4 | 0.8113 | 366 | 421 | 111 |
| 5 | 5 | 1.5983 | 421 | 495 | 154 |
| 6 | 6 | 1.8750 | 495 | 612 | 191 |
| 7 | 7 | 84.5748 | 612 | 1114 | 19290 |
| 8 | 8 | 0.5488 | 1363 | 1667 | 105 |
| | | | | | |
| | | | | | |
| | | TIC: | 106.1046 | ng/uL | |
| | | TIM: | 575.7487 | nmole/L | |
| | | Total concentration: | 107.1061 | ng/uL | |
| | | 28s/18s: | 0.0 | | |
| | | RQN | 1.1 | | |
| | | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: G2_PU_TP1 Well location: D10



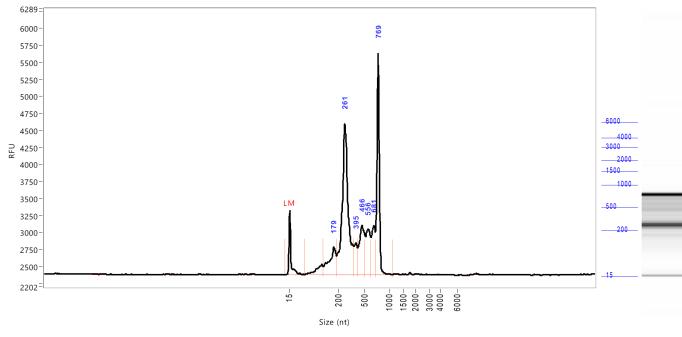
| | | | | 312 |
|------|----------------------|-----------|---------|-------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 75 | 880 |
| 2 | 3.8853 | 156 | 188 | 312 |
| 3 | 43.6088 | 188 | 371 | 3185 |
| 4 | 3.0096 | 371 | 413 | 384 |
| 5 | 10.0523 | 413 | 501 | 788 |
| 6 | 9.9272 | 501 | 617 | 924 |
| 7 | 9.4682 | 617 | 704 | 1211 |
| 8 | 94.6420 | 704 | 1162 | 18779 |
| 9 | 0.4082 | 1387 | 1621 | 72 |
| 10 | 0.1848 | 1621 | 1834 | 51 |
| | | | | |
| | TIC: | 175.1862 | ng/uL | |
| | TIM: | 1153.5587 | nmole/L | |
| | Total concentration: | 176.2612 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: G2_PU_TP1 Well location: D11

Created: Tuesday, December 08, 2020 4:16:51 PM

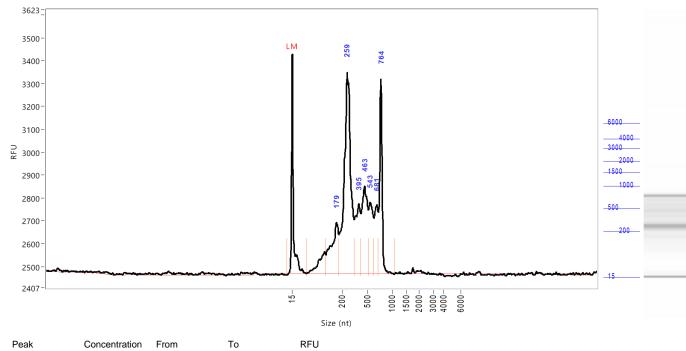


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 70 | 941 |
| 2 | 6.9332 | 141 | 190 | 399 |
| 3 | 33.6718 | 190 | 371 | 2217 |
| 4 | 3.6741 | 371 | 416 | 466 |
| 5 | 9.1908 | 416 | 506 | 722 |
| 6 | 7.0258 | 506 | 621 | 663 |
| 7 | 6.0725 | 621 | 714 | 714 |
| 8 | 17.5620 | 714 | 1098 | 3251 |
| | | | | |
| | TIC: | 04 4202 | n a/ul | |
| | TIC: | 84.1302 | ng/uL | |
| | TIM: | 745.0142 | nmole/L | |
| | Total concentration: | 85.6007 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                            Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: G2_PU_TP1 Well location: D12



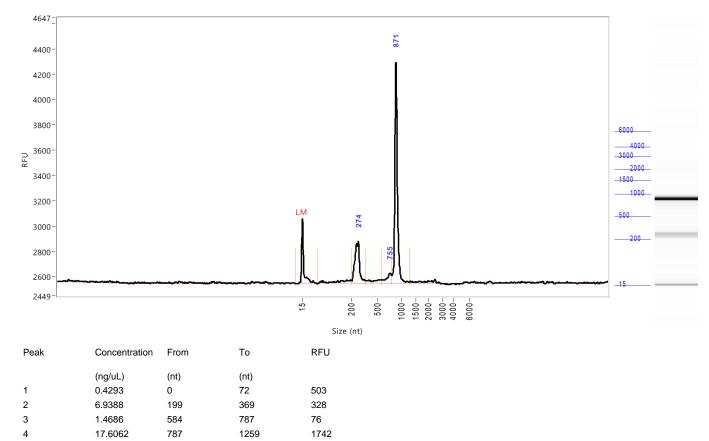
| | | | | 31. |
|------|----------------|----------|---------|-----|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 69 | 956 |
| 2 | 4.0590 | 140 | 188 | 221 |
| 3 | 14.1211 | 188 | 343 | 875 |
| 4 | 3.5490 | 343 | 416 | 301 |
| 5 | 4.9118 | 416 | 506 | 379 |
| 6 | 3.2416 | 506 | 626 | 309 |
| 7 | 2.4054 | 626 | 714 | 297 |
| 8 | 4.6779 | 714 | 1065 | 846 |
| | | | | |
| | TIC: | 36.9657 | ng/uL | |
| | TIM: | 355.5205 | nmole/L | |
| | Total | 38.2111 | ng/uL | |
| | concentration: | | - | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A1_TP1 Well location: E1

Created: Tuesday, December 08, 2020 4:16:51 PM

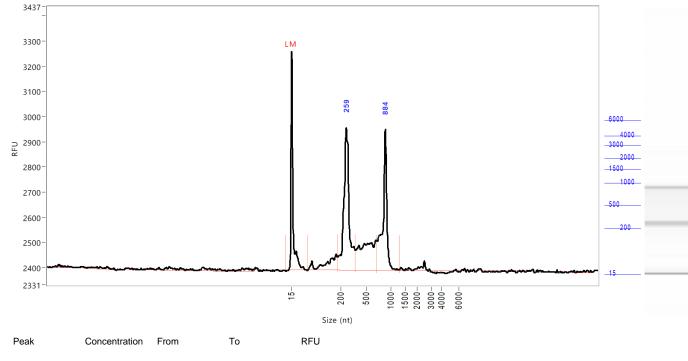


| TIC: | 26.0136 | ng/uL |
|----------------------|----------|---------|
| TIM: | 152.2598 | nmole/L |
| Total concentration: | 29.1971 | ng/uL |
| 28s/18s: | 0.0 | |
| RQN | 1.3 | |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: A1_TP2 Well location: E2



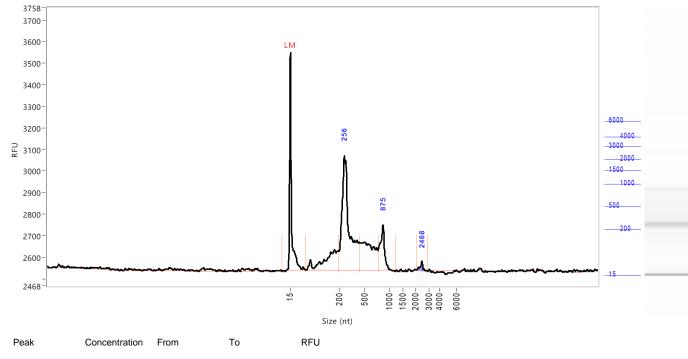
| Peak | Concentration | From | То | RFU |
|------|----------------------|------------|---------|-----|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 74 | 868 |
| 2 | 8.1215 | 187 | 366 | 565 |
| 3 | 5.1549 | 718 | 1283 | 559 |
| | | | | |
| | | | | |
| | TIC: | 13.2764 | ng/uL | |
| | TIM: | 115.0132 | nmole/L | |
| | Total concentration: | 20.0380 | ng/uL | |
| | 28s/18s: RQN | 0.0 1.2 | | |
| | 113(11 | | | |

```
Sample baseline V to V?: Y
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                                                       Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
                                                          Upper marker selection: Last peak > 100 RFU
Lower marker selection: First peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A1_TP3 Well location: E3

Created: Tuesday, December 08, 2020 4:16:51 PM



| Peak | Concentration | From | То | RFU |
|------|---------------|------|------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 75 | 1008 |
| 2 | 9.0785 | 199 | 442 | 529 |
| 3 | 1.7637 | 783 | 1218 | 208 |
| 4 | 0.1697 | 2049 | 2903 | 40 |
| | | | | |
| | | | | |

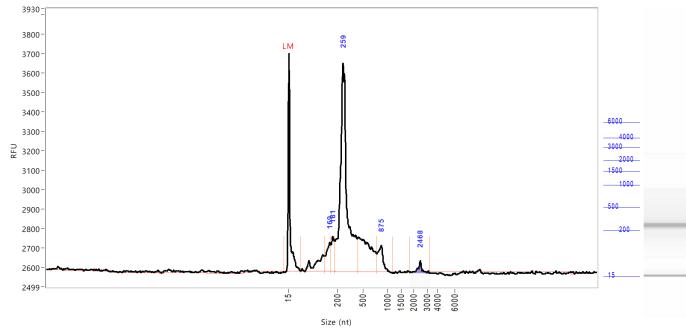
| 11.0119 | ng/uL |
|----------|----------------------------|
| 102.8242 | nmole/L |
| 17.9499 | ng/uL |
| 0.0 | |
| 1.2 | |
| | 102.8242 17.9499 0.0 |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: A1_TP4 Well location: E4

Created: Tuesday, December 08, 2020 4:16:51 PM



| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 59 | 1121 |
| 2 | 1.3656 | 151 | 173 | 147 |
| 3 | 1.2399 | 173 | 188 | 178 |
| 4 | 15.4723 | 188 | 437 | 1068 |
| 5 | 1.2765 | 778 | 1186 | 132 |
| 6 | 0.2015 | 1843 | 3218 | 53 |
| | | | | |
| | TIO | 40 5557 | | |
| | TIC: | 19.5557 | ng/uL | |
| | TIM: | 223.7922 | nmole/L | |
| | Total concentration: | 25.4948 | ng/uL | |
| | 28s/18s: | 0.0 | | |

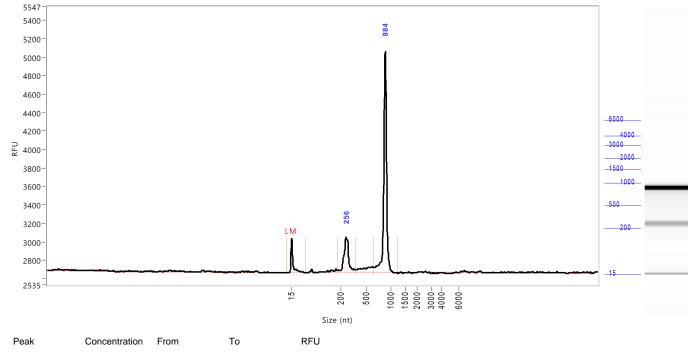
```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                       Sample baseline V to V points: 3
                            Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

RQN

1.1

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Sample: D1_TP1 Well location: E5

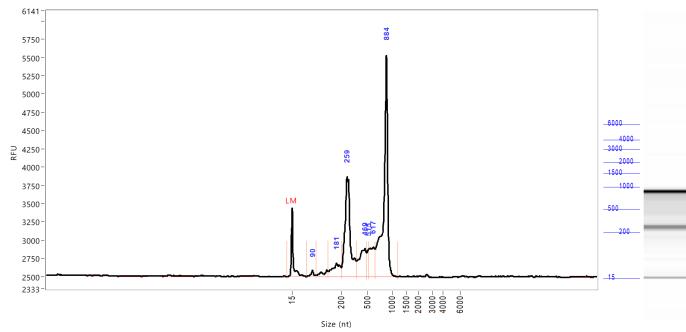


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 69 | 358 |
| 2 | 12.2021 | 187 | 379 | 377 |
| 3 | 40.5485 | 649 | 1242 | 2392 |
| | | | | |
| | | | | |
| | TIC: | 52.7506 | ng/uL | |
| | TIM: | 291.6027 | nmole/L | |
| | Total concentration: | 58.7175 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.2 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: D1_TP2 Well location: E6

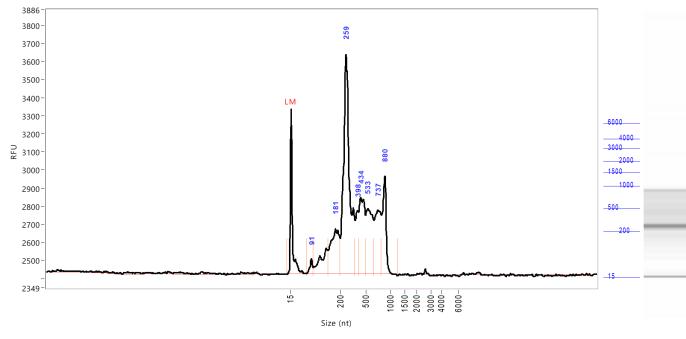


| | | | | 31 |
|------|----------------------|----------|---------|------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 69 | 931 |
| 2 | 0.5647 | 69 | 106 | 80 |
| 3 | 3.5902 | 150 | 198 | 181 |
| 4 | 18.2933 | 198 | 369 | 1365 |
| 5 | 6.4891 | 369 | 490 | 378 |
| 6 | 1.6147 | 490 | 519 | 368 |
| 7 | 4.8885 | 519 | 649 | 399 |
| 8 | 27.0409 | 649 | 1186 | 3029 |
| | | | | |
| | 710 | | | |
| | TIC: | 62.4815 | ng/uL | |
| | TIM: | 477.0452 | nmole/L | |
| | Total concentration: | 63.6183 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: D1_TP3 Well location: E7



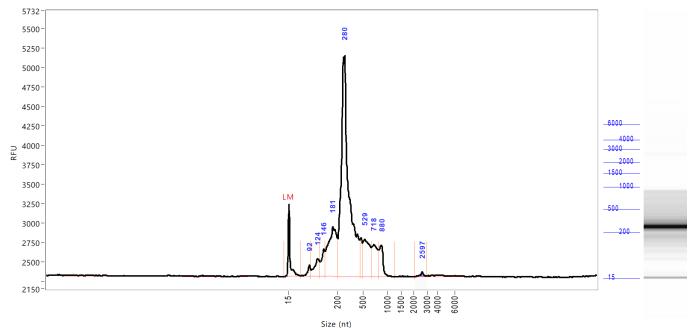
| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 71 | 907 |
| 2 | 0.5736 | 71 | 98 | 80 |
| 3 | 5.6179 | 153 | 198 | 249 |
| 4 | 19.0264 | 198 | 369 | 1212 |
| 5 | 2.5863 | 369 | 411 | 347 |
| 6 | 5.7638 | 411 | 490 | 420 |
| 7 | 5.9705 | 490 | 653 | 360 |
| 8 | 5.0380 | 653 | 806 | 351 |
| 9 | 4.5749 | 806 | 1234 | 539 |
| | | | | |
| | TIO | 40.454.4 | | |
| | TIC: | 49.1514 | ng/uL | |
| | TIM: | 467.4662 | nmole/L | |
| | Total concentration: | 51.2738 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.0 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: D1_TP4 Well location: E8

Created: Tuesday, December 08, 2020 4:16:51 PM

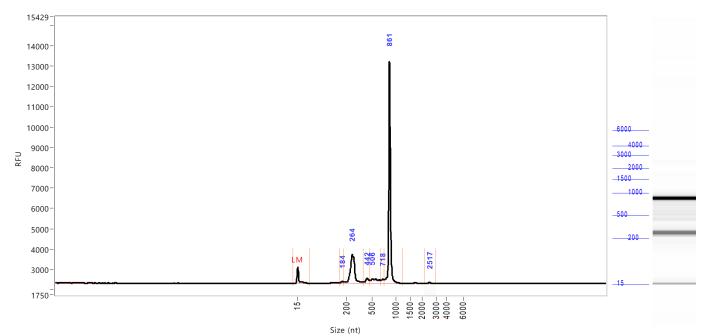


| | | | | Size (III) |
|------|----------------------|-----------|---------|------------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 59 | 925 |
| 2 | 1.2079 | 59 | 99 | 149 |
| 3 | 3.2978 | 99 | 132 | 228 |
| 4 | 3.3030 | 132 | 151 | 351 |
| 5 | 14.3436 | 151 | 199 | 641 |
| 6 | 54.8337 | 199 | 463 | 2848 |
| 7 | 7.5970 | 492 | 663 | 477 |
| 8 | 5.6197 | 663 | 815 | 410 |
| 9 | 3.6509 | 815 | 1250 | 398 |
| 10 | 0.3551 | 2082 | 2952 | 62 |
| | | | | |
| | TIC: | 94.2086 | ng/uL | |
| | TIM: | 1099.2977 | nmole/L | |
| | Total concentration: | 96.2756 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: F1_TP1 Well location: E9

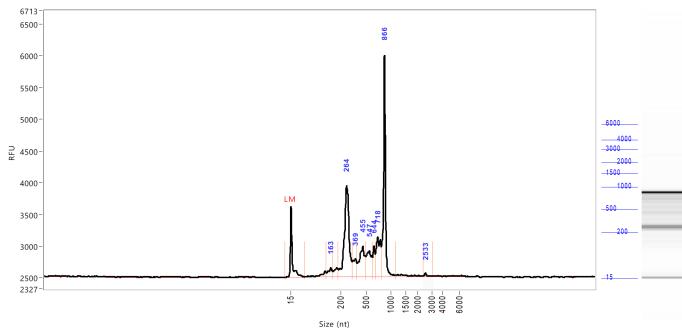


| | | | | , |
|------|----------------------|----------|------------------|-------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 59 | 812 |
| 2 | 0.7001 | 174 | 190 | 89 |
| 3 | 18.3798 | 190 | 392 | 1410 |
| 4 | 2.3705 | 392 | 471 | 242 |
| 5 | 4.5295 | 471 | 672 | 230 |
| 6 | 1.4260 | 672 | 746 | 203 |
| 7 | 48.2881 | 746 | 1234 | 10930 |
| 8 | 0.2598 | 2130 | 2984 | 50 |
| | | | | |
| | TIC: | 75.0520 | ng/ul | |
| | | 75.9539 | ng/uL nmole/L | |
| | TIM: | 448.3749 | | |
| | Total concentration: | 76.8270 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: F1_TP2 Well location: E10



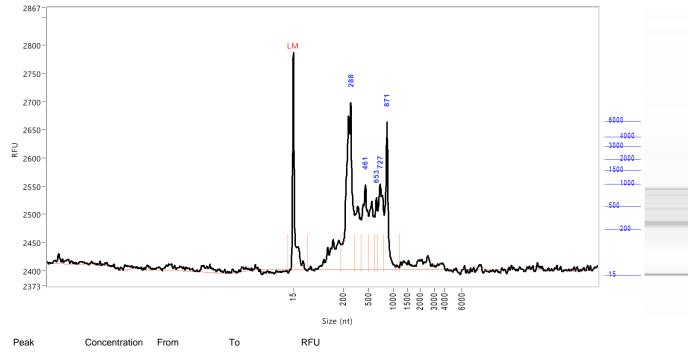
| Peak | Concentration | From | То | RFU |
|------|----------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 66 | 1115 |
| 2 | 1.1814 | 146 | 170 | 148 |
| 3 | 13.4658 | 191 | 330 | 1444 |
| 4 | 2.2577 | 330 | 387 | 286 |
| 5 | 5.0351 | 387 | 492 | 490 |
| 6 | 3.8094 | 492 | 617 | 417 |
| 7 | 1.8634 | 617 | 672 | 484 |
| 8 | 5.1654 | 672 | 797 | 633 |
| 9 | 12.3874 | 797 | 1138 | 3499 |
| 10 | 0.3042 | 2372 | 3088 | 62 |
| | | | | |
| | TIC: | 45.4698 | ng/uL | |
| | TIM: | 335.4366 | nmole/L | |
| | Total | 48.4827 | ng/uL | |
| | concentration: | 40.4027 | rig/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.2 | | |
| | | | | |

```
Sample baseline V to V?: Y
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100
                                                          Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: F1_TP3 Well location: E11

Created: Tuesday, December 08, 2020 4:16:51 PM



| Peak | Concentration | From | То | RFU | |
|------|---------------|----------|---------|-----|--|
| | (ng/uL) | (nt) | (nt) | | |
| 1 | 0.4293 | 0 | 67 | 384 | |
| 2 | 9.5253 | 194 | 335 | 294 | |
| 3 | 3.8388 | 413 | 500 | 149 | |
| 4 | 1.5254 | 621 | 681 | 125 | |
| 5 | 3.4542 | 681 | 811 | 150 | |
| 6 | 3.6072 | 811 | 1210 | 260 | |
| | | | | | |
| | | | | | |
| | TIC: | 21.9510 | ng/uL | | |
| | TIM: | 171.9840 | nmole/L | | |

ng/uL

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

Total

RQN

concentration: 28s/18s:

32.5411

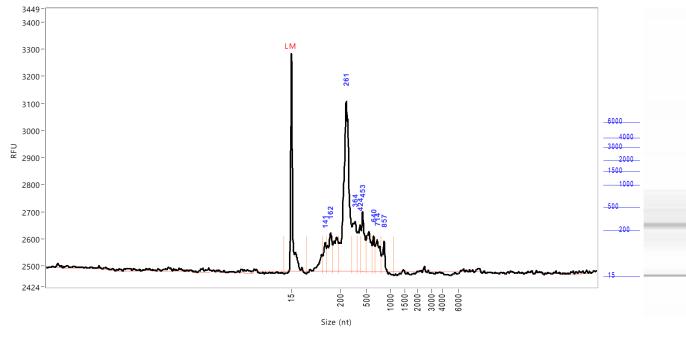
0.0

1.4

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Sample: F1_TP4 Well location: E12

Created: Tuesday, December 08, 2020 4:16:51 PM

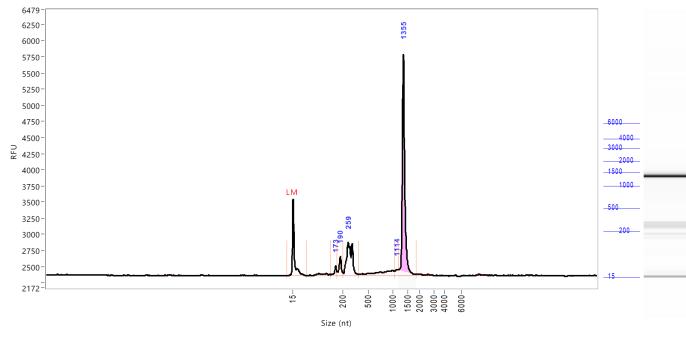


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|-----|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 71 | 801 |
| 2 | 0.8754 | 132 | 146 | 104 |
| 3 | 1.8307 | 146 | 169 | 138 |
| 4 | 10.2802 | 192 | 327 | 625 |
| 5 | 2.2148 | 327 | 387 | 182 |
| 6 | 1.6159 | 387 | 437 | 169 |
| 7 | 1.9206 | 437 | 492 | 218 |
| 8 | 0.7554 | 612 | 667 | 128 |
| 9 | 1.4083 | 667 | 801 | 115 |
| 10 | 0.6865 | 801 | 1106 | 110 |
| | | | | |
| | TIC: | 21.5877 | ng/uL | |
| | TIM: | 236.5006 | nmole/L | |
| | Total concentration: | 25.7191 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: G1_TP1 Well location: F1

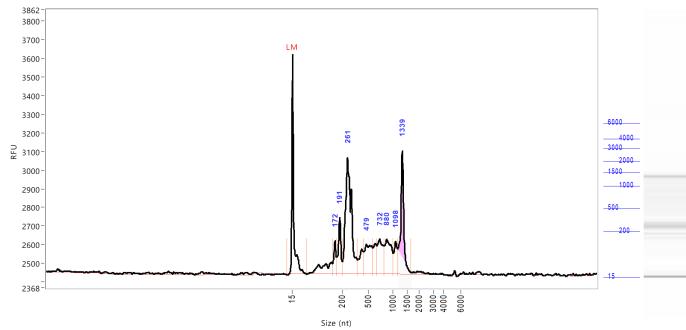


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 65 | 1180 |
| 2 | 0.6435 | 156 | 179 | 152 |
| 3 | 1.3255 | 179 | 206 | 293 |
| 4 | 5.6376 | 206 | 387 | 513 |
| 5 | 0.4935 | 1033 | 1178 | 93 |
| 6 | 13.9780 | 1178 | 1852 | 3430 |
| | | | | |
| | TIC: | 22.0781 | ng/uL | |
| | TIM: | 130.0640 | nmole/L | |
| | Total concentration: | 25.6912 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 6.7 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: G1_TP2 Well location: F2



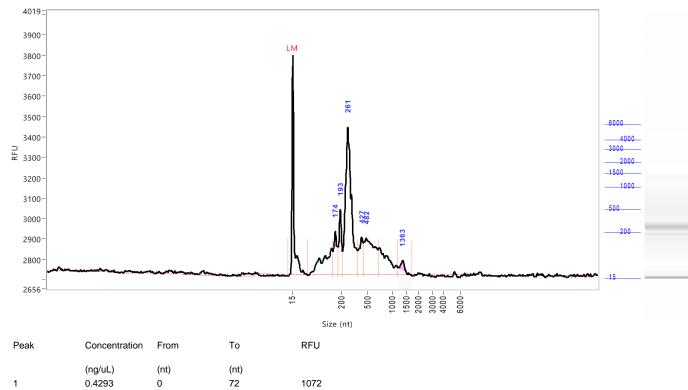
| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 68 | 1175 |
| 2 | 0.8366 | 163 | 178 | 174 |
| 3 | 1.5698 | 178 | 200 | 298 |
| 4 | 7.3376 | 200 | 377 | 621 |
| 5 | 2.1497 | 448 | 593 | 156 |
| 6 | 1.9075 | 677 | 824 | 184 |
| 7 | 2.2058 | 824 | 1017 | 182 |
| 8 | 0.9704 | 1017 | 1178 | 171 |
| 9 | 3.5513 | 1178 | 1677 | 657 |
| | | | | |
| | TIC: | 20.5287 | ng/uL | |
| | TIM: | 164.4352 | nmole/L | |
| | Total concentration: | 23.9770 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 2.6 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: G1_TP3 Well location: F3

Created: Tuesday, December 08, 2020 4:16:51 PM

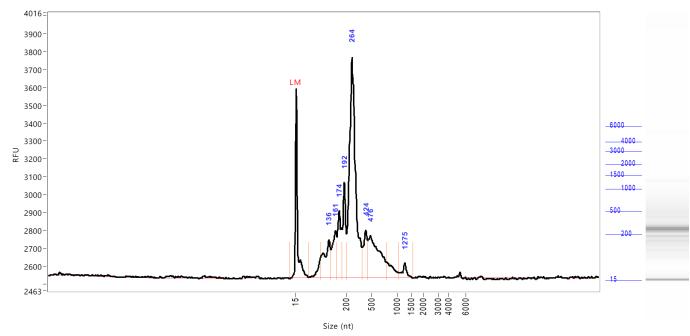


| | (ng/uL) | (nt) | (nt) | |
|---|----------------------|----------|---------|------|
| 1 | 0.4293 | 0 | 72 | 1072 |
| 2 | 1.5559 | 166 | 184 | 208 |
| 3 | 1.8308 | 184 | 204 | 316 |
| 4 | 9.4471 | 204 | 382 | 719 |
| 5 | 1.5038 | 382 | 445 | 181 |
| 6 | 3.9937 | 445 | 718 | 176 |
| 7 | 0.5815 | 1194 | 1732 | 67 |
| | | | | |
| | TIC: | 18.9128 | ng/uL | |
| | TIM: | | · · | |
| | I IIVI. | 198.5329 | nmole/L | |
| | Total concentration: | 24.1460 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.3 | | |
| | | | | |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: G1_TP4 Well location: F4

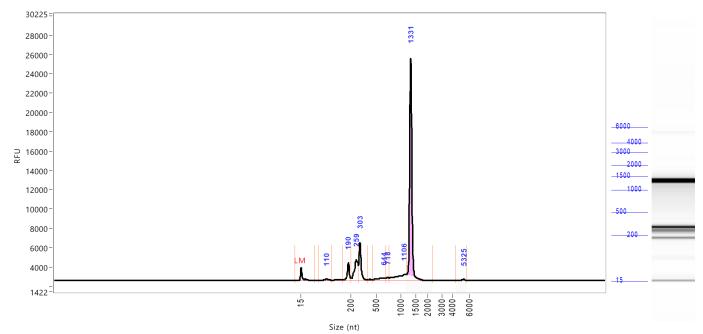


| Concentration | From | То | RFU |
|----------------------|------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| (ng/uL) | (nt) | (nt) | |
| 0.4293 | 0 | 61 | 1051 |
| 3.0746 | 105 | 144 | 210 |
| 2.5530 | 144 | 166 | 260 |
| 3.1032 | 166 | 185 | 371 |
| 3.1509 | 185 | 204 | 529 |
| 16.0288 | 204 | 382 | 1229 |
| 2.2105 | 382 | 448 | 264 |
| 5.2712 | 448 | 801 | 233 |
| 0.5596 | 1065 | 1566 | 79 |
| 710 | | | |
| | | - | |
| | | | |
| Total concentration: | 37.2018 | ng/uL | |
| 28s/18s: | 0.0 | | |
| RQN | 1.0 | | |
| | (ng/uL) 0.4293 3.0746 2.5530 3.1032 3.1509 16.0288 2.2105 5.2712 0.5596 TIC: TIM: Total concentration: 28s/18s: | (ng/uL) (nt) 0.4293 0 3.0746 105 2.5530 144 3.1032 166 3.1509 185 16.0288 204 2.2105 382 5.2712 448 0.5596 1065 TIC: 35.9517 TIM: 460.6989 Total 37.2018 concentration: 28s/18s: 0.0 | (ng/uL) (nt) (nt) 0.4293 0 61 3.0746 105 144 2.5530 144 166 3.1032 166 185 3.1509 185 204 16.0288 204 382 2.2105 382 448 5.2712 448 801 0.5596 1065 1566 TIC: 35.9517 ng/uL Total 37.2018 ng/uL concentration: 28s/18s: 0.0 |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: B2_TP1 Well location: F5



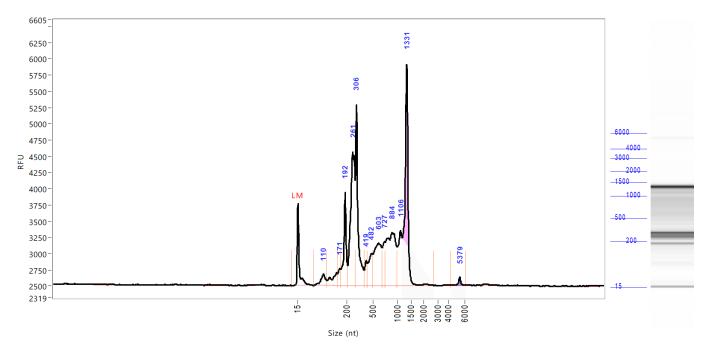
| Peak | Concentration | From | То | RFU |
|------|----------------|----------|---------|-------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 65 | 1326 |
| 2 | 1.0962 | 83 | 130 | 153 |
| 3 | 6.9573 | 172 | 206 | 1816 |
| 4 | 13.9392 | 206 | 288 | 2132 |
| 5 | 13.7758 | 288 | 390 | 3976 |
| 6 | 3.1483 | 453 | 672 | 241 |
| 7 | 1.4805 | 672 | 750 | 286 |
| 8 | 9.4516 | 750 | 1170 | 642 |
| 9 | 76.8771 | 1170 | 2291 | 23025 |
| 10 | 0.4395 | 4352 | 5649 | 147 |
| | | | | |
| | TIC: | 127.1656 | ng/uL | |
| | TIM: | 688.2629 | nmole/L | |
| | Total | 128.8276 | ng/uL | |
| | concentration: | 120.0270 | rig/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 7.0 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: B2_TP2 Well location: F6

Created: Tuesday, December 08, 2020 4:16:51 PM



| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 73 | 1261 |
| 2 | 1.6643 | 73 | 125 | 173 |
| 3 | 1.3095 | 163 | 176 | 260 |
| 4 | 6.8027 | 176 | 201 | 1428 |
| 5 | 16.3945 | 201 | 290 | 2056 |
| 6 | 12.6301 | 290 | 395 | 2777 |
| 7 | 1.7894 | 395 | 434 | 387 |
| 8 | 3.3038 | 434 | 495 | 489 |
| 9 | 8.2542 | 495 | 681 | 650 |
| 10 | 3.3545 | 681 | 750 | 680 |
| 11 | 12.0897 | 750 | 995 | 811 |
| 12 | 5.1440 | 995 | 1170 | 851 |
| 13 | 16.1833 | 1170 | 2726 | 3411 |
| 14 | 0.5022 | 4325 | 6134 | 134 |
| | | | | |
| | | | | |
| | TIC: | 89.4222 | ng/uL | |
| | TIM: | 696.3689 | nmole/L | |
| | Total concentration: | 90.2689 | ng/uL | |
| | 28s/18s: | 0.0 | | |

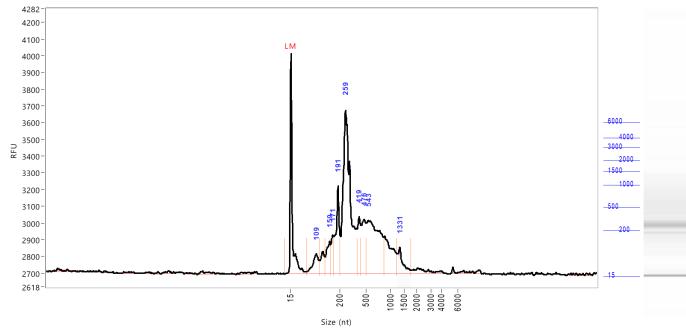
2.9

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                            Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

RQN

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Sample: B2_TP3 Well location: F7

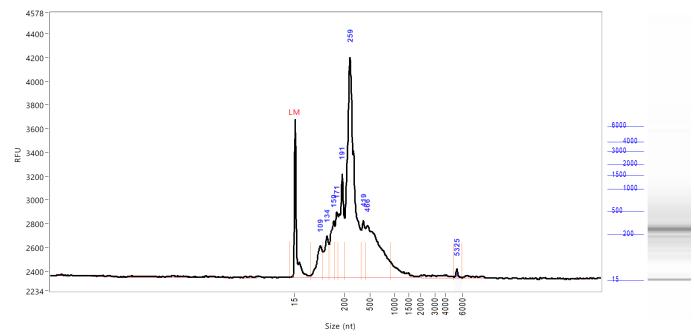


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 73 | 1314 |
| 2 | 1.3112 | 73 | 124 | 114 |
| 3 | 1.3539 | 142 | 163 | 188 |
| 4 | 1.0936 | 163 | 176 | 224 |
| 5 | 3.1654 | 176 | 200 | 519 |
| 6 | 12.2768 | 200 | 395 | 974 |
| 7 | 1.9485 | 395 | 445 | 338 |
| 8 | 2.0544 | 445 | 500 | 319 |
| 9 | 6.6863 | 500 | 871 | 317 |
| 10 | 1.1515 | 1234 | 1769 | 156 |
| | | | | |
| | | | | |
| | TIC: | 31.0414 | ng/uL | |
| | TIM: | 335.0150 | nmole/L | |
| | Total concentration: | 34.9373 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.6 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: B2_TP4 Well location: F8

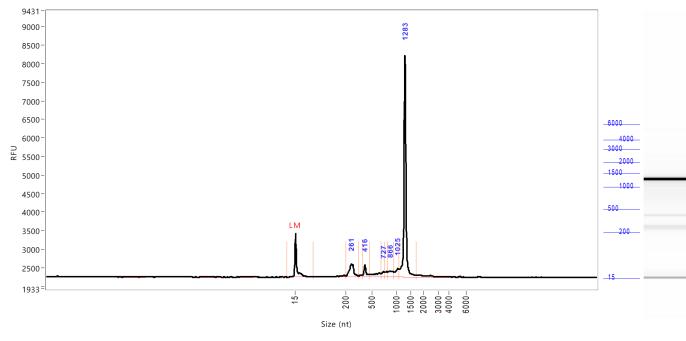


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|----------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 71 | 1327 |
| 2 | 2.7179 | 71 | 118 | 260 |
| 3 | 2.7869 | 118 | 141 | 343 |
| 4 | 3.6028 | 141 | 163 | 470 |
| 5 | 2.8284 | 163 | 176 | 547 |
| 6 | 6.1378 | 176 | 200 | 869 |
| 7 | 22.0315 | 200 | 392 | 1853 |
| 8 | 2.7925 | 392 | 445 | 479 |
| 9 | 9.7982 | 445 | 912 | 432 |
| 10 | 0.1796 | 5000 | 5973 | 70 |
| | | | | |
| | TIC: | E0 07E7 | m er/u l | |
| | | 52.8757 | ng/uL | |
| | TIM: | 692.4393 | nmole/L | |
| | Total concentration: | 54.3629 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.2 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                         Sample baseline V to V?: Y
                                                                                       Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
                                                          Upper marker selection: Last peak > 100 RFU
Lower marker selection: First peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: C2_TP1 Well location: F9

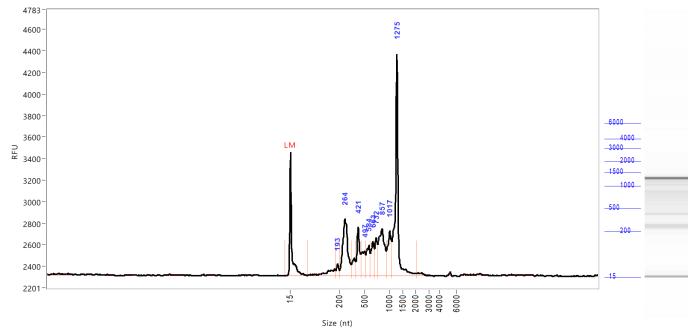


| Р | Peak | Concentration | From | То | RFU |
|---|------|----------------------|----------|---------|------|
| | | (ng/uL) | (nt) | (nt) | |
| 1 | | 0.4293 | 0 | 80 | 1173 |
| 2 | 2 | 3.1944 | 200 | 345 | 350 |
| 3 | 3 | 1.5813 | 387 | 482 | 329 |
| 4 | 1 | 0.7225 | 677 | 755 | 144 |
| 5 | 5 | 1.3655 | 815 | 945 | 159 |
| 6 | 6 | 1.3659 | 945 | 1073 | 232 |
| 7 | , | 19.4541 | 1073 | 1714 | 5984 |
| | | | | | |
| | | TIO: | 07.0000 | | |
| | | TIC: | 27.6836 | ng/uL | |
| | | TIM: | 109.1140 | nmole/L | |
| | | Total concentration: | 31.3861 | ng/uL | |
| | | 28s/18s: | 0.0 | | |
| | | RQN | 1.2 | | |
| | | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: C2_TP2 Well location: F10



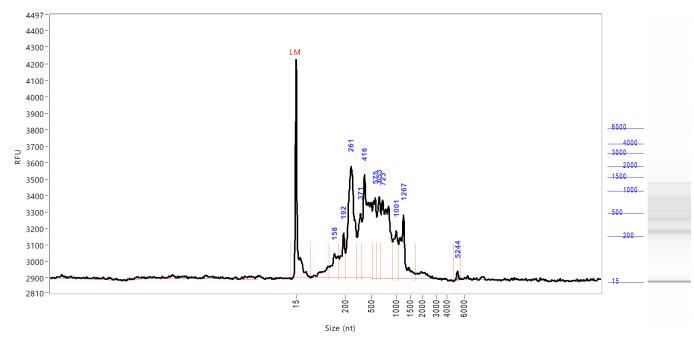
| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 79 | 1138 |
| 2 | 0.4801 | 187 | 201 | 104 |
| 3 | 5.1696 | 201 | 340 | 521 |
| 4 | 2.3422 | 390 | 453 | 445 |
| 5 | 1.6850 | 453 | 515 | 226 |
| 6 | 1.6992 | 515 | 607 | 282 |
| 7 | 1.8534 | 607 | 695 | 316 |
| 8 | 1.4935 | 695 | 760 | 350 |
| 9 | 4.4866 | 760 | 940 | 433 |
| 10 | 2.6935 | 940 | 1090 | 415 |
| 11 | 9.3152 | 1090 | 2049 | 2056 |
| | | | | |
| | TIO | 04.0400 | | |
| | TIC: | 31.2183 | ng/uL | |
| | TIM: | 168.5023 | nmole/L | |
| | Total concentration: | 33.4617 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.3 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: C2_TP3 Well location: F11

Created: Tuesday, December 08, 2020 4:16:51 PM



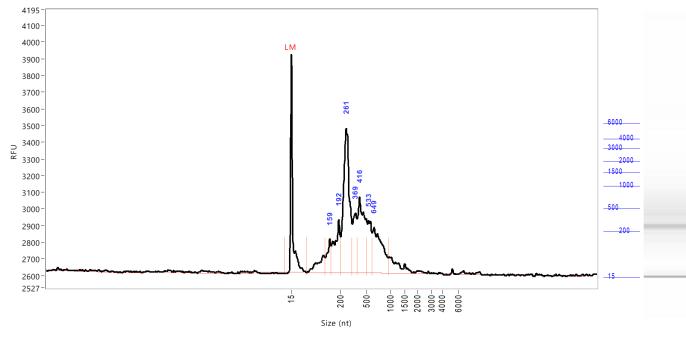
| Peak | Concentration | From | То | RFU |
|------|---------------|------|------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 70 | 1322 |
| 2 | 1.6340 | 139 | 176 | 144 |
| 3 | 1.6088 | 176 | 198 | 270 |
| 4 | 7.2006 | 198 | 332 | 675 |
| 5 | 2.4873 | 332 | 390 | 389 |
| 6 | 6.6548 | 390 | 510 | 624 |
| 7 | 2.7765 | 510 | 603 | 484 |
| 8 | 2.6395 | 603 | 690 | 491 |
| 9 | 5.8123 | 690 | 931 | 475 |
| 10 | 1.9400 | 931 | 1081 | 288 |
| 11 | 2.7443 | 1081 | 1714 | 386 |
| 12 | 0.0960 | 4757 | 5568 | 45 |
| | | | | |
| | | | | |

TIC: 35.5942 ng/uL
TIM: 275.2287 nmole/L
Total 36.4131 ng/uL
concentration: 28s/18s: 0.0
RQN 1.2

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Dilution factor: 12.0 Quantification using: Ladder Final concentration (ng/uL): 8.0000 Minimum RFU for data processing: 2

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Sample: C2_TP4 Well location: F12



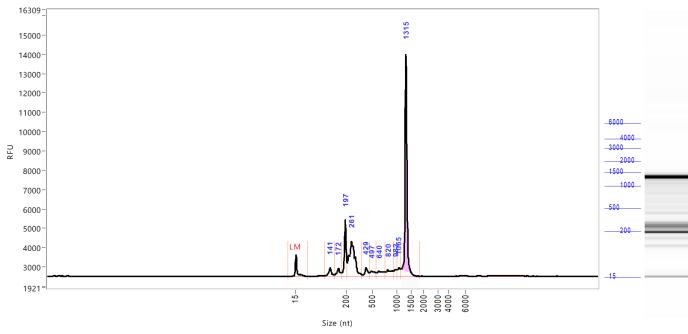
| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 71 | 1306 |
| 2 | 1.2795 | 141 | 163 | 198 |
| 3 | 3.0958 | 163 | 199 | 314 |
| 4 | 9.0242 | 199 | 330 | 862 |
| 5 | 2.3834 | 330 | 387 | 357 |
| 6 | 5.2520 | 387 | 510 | 453 |
| 7 | 1.9209 | 510 | 607 | 312 |
| 8 | 4.1768 | 607 | 958 | 274 |
| | | | | |
| | TIC: | 27.1325 | ng/uL | |
| | TIM: | 271.3295 | nmole/L | |
| | Total concentration: | 29.7625 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: D2_TP1 Well location: G1

Created: Tuesday, December 08, 2020 4:16:51 PM



| Peak | Concentration | From | То | RFU |
|------|----------------|----------|---------|-------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 58 | 1098 |
| 2 | 3.1088 | 121 | 156 | 441 |
| 3 | 3.3716 | 156 | 186 | 422 |
| 4 | 11.9796 | 186 | 211 | 2915 |
| 5 | 22.2056 | 211 | 387 | 1823 |
| 6 | 2.8244 | 387 | 466 | 447 |
| 7 | 2.6385 | 466 | 580 | 264 |
| 8 | 3.4723 | 580 | 764 | 265 |
| 9 | 3.6239 | 764 | 931 | 337 |
| 10 | 2.0809 | 931 | 1009 | 340 |
| 11 | 2.4043 | 1009 | 1138 | 459 |
| 12 | 41.3831 | 1138 | 1815 | 11495 |
| | | | | |
| | | | | |
| | TIC: | 99.0929 | ng/uL | |
| | TIM: | 753.0867 | nmole/L | |
| | Total | 99.8957 | ng/uL | |
| | concentration: | | | |
| | | | | |

0.0

5.2

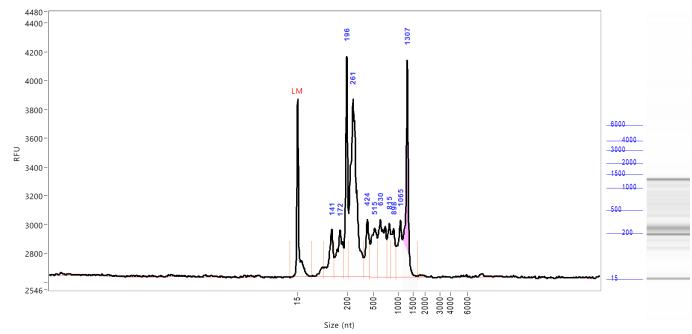
Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

28s/18s:

RQN

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Sample: D2_TP2 Well location: G2

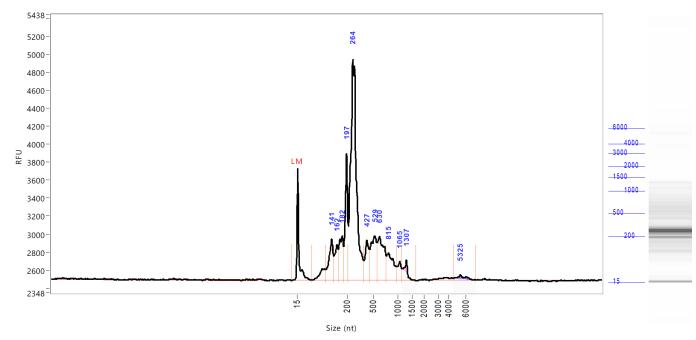


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 66 | 1233 |
| 2 | 2.6638 | 111 | 149 | 329 |
| 3 | 3.4503 | 149 | 185 | 322 |
| 4 | 6.0868 | 185 | 206 | 1530 |
| 5 | 14.0622 | 206 | 382 | 1232 |
| 6 | 2.6713 | 382 | 461 | 396 |
| 7 | 3.2294 | 461 | 570 | 335 |
| 8 | 4.5753 | 570 | 764 | 396 |
| 9 | 1.9763 | 764 | 848 | 368 |
| 10 | 2.1722 | 848 | 954 | 333 |
| 11 | 2.5059 | 954 | 1138 | 392 |
| 12 | 6.0356 | 1138 | 1695 | 1502 |
| | | | | |
| | | | | |
| | TIC: | 49.4291 | ng/uL | |
| | TIM: | 483.0879 | nmole/L | |
| | Total concentration: | 49.5714 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 2.3 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: D2_TP3 Well location: G3



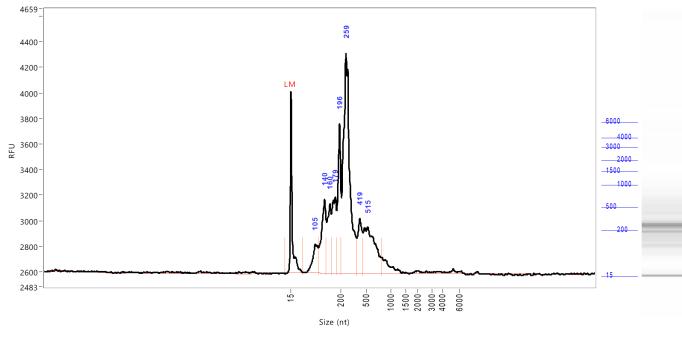
| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 67 | 1231 |
| 2 | 3.8459 | 119 | 150 | 454 |
| 3 | 2.5407 | 150 | 166 | 389 |
| 4 | 4.0514 | 166 | 187 | 482 |
| 5 | 6.6394 | 187 | 209 | 1402 |
| 6 | 27.1314 | 209 | 395 | 2450 |
| 7 | 3.0395 | 395 | 458 | 437 |
| 8 | 4.9813 | 458 | 580 | 490 |
| 9 | 5.4515 | 580 | 769 | 480 |
| 10 | 3.4780 | 769 | 986 | 299 |
| 11 | 1.1984 | 986 | 1154 | 203 |
| 12 | 1.2899 | 1154 | 1677 | 223 |
| 13 | 0.5978 | 4514 | 7175 | 60 |
| | | | | |
| | | | | |
| | TIC: | 64.2452 | ng/uL | |
| | TIM: | 719.2025 | nmole/L | |
| | Total concentration: | 65.0488 | ng/uL | |
| | 28s/18s: | 1.0 | | |
| | RQN | 1.4 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: D2_TP4 Well location: G4

Created: Tuesday, December 08, 2020 4:16:51 PM

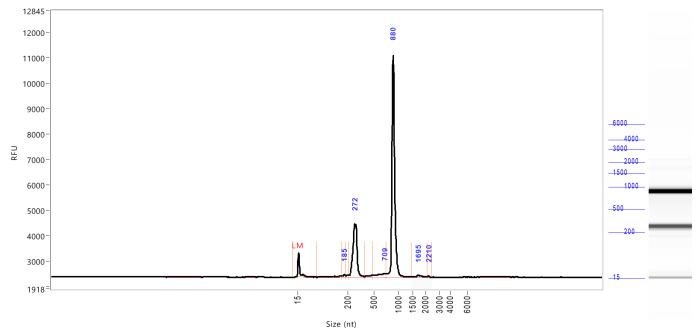


| | | | | Size (iii) |
|------|----------------------|----------|---------|------------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 60 | 1418 |
| 2 | 2.1732 | 60 | 117 | 222 |
| 3 | 4.7021 | 117 | 146 | 573 |
| 4 | 3.4942 | 146 | 165 | 541 |
| 5 | 4.5270 | 165 | 185 | 595 |
| 6 | 5.5886 | 185 | 206 | 1167 |
| 7 | 18.0442 | 206 | 382 | 1718 |
| 8 | 3.0899 | 382 | 458 | 425 |
| 9 | 6.1937 | 458 | 792 | 359 |
| | | | | |
| | TIC: | 47.8130 | ng/uL | |
| | TIM: | 677.2305 | nmole/L | |
| | Total concentration: | 48.8988 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

Sample baseline V to V?: Y Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Upper marker selection: Last peak > 100 RFU Lower marker selection: First peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: E2_TP1 Well location: G5

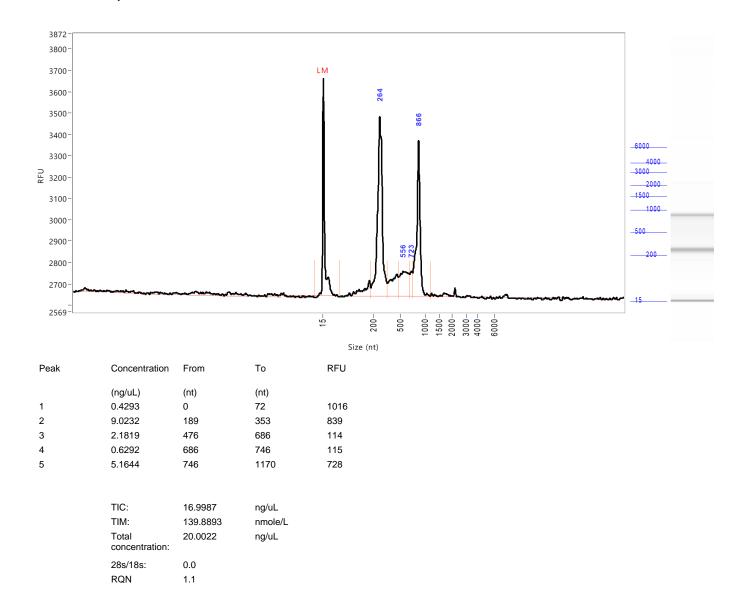


| | | | | 31 |
|------|----------------------|----------|---------|------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 83 | 959 |
| 2 | 0.6269 | 175 | 191 | 106 |
| 3 | 20.3429 | 204 | 385 | 2087 |
| 4 | 2.4437 | 482 | 737 | 137 |
| 5 | 49.3994 | 737 | 1444 | 8725 |
| 6 | 0.8060 | 1444 | 2098 | 77 |
| 7 | 0.1858 | 2098 | 2436 | 61 |
| | | | | |
| | TIO | 70.0040 | | |
| | TIC: | 73.8046 | ng/uL | |
| | TIM: | 431.1071 | nmole/L | |
| | Total concentration: | 76.5664 | ng/uL | |
| | 28s/18s: | 0.3 | | |
| | RQN | 1.2 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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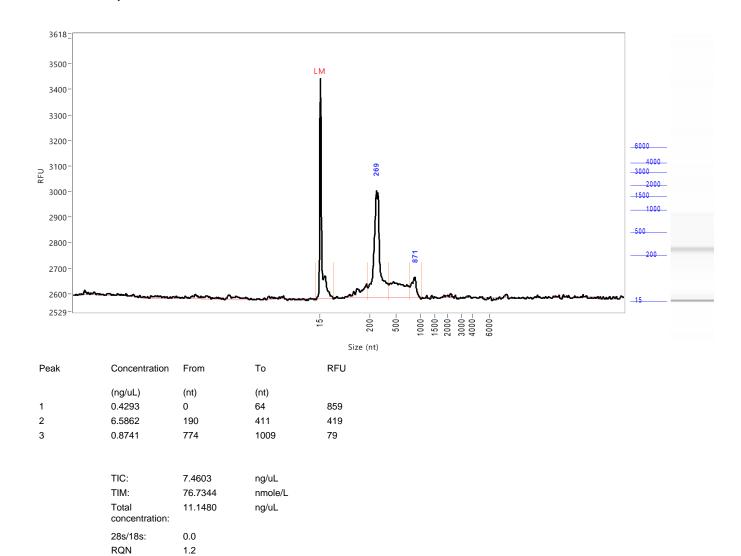
Sample: E2_TP2 Well location: G6



```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                           Sample baseline V to V?: Y
                                                                                         Sample baseline V to V points: 3
Sample filter: Binomial
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Manual baseline start (min): 15
                                   Manual baseline end (min): 38
                             Marker min peak height: 100
Marker peak width (sec): 6
                                                           Marker baseline V to V?: Y
                                                                                         Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                                                                  Dilution factor: 12.0
Quantification using: Ladder
                                     Final concentration (ng/uL): 8.0000
Minimum RFU for data processing: 2
```

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Sample: E2_TP3 Well location: G7

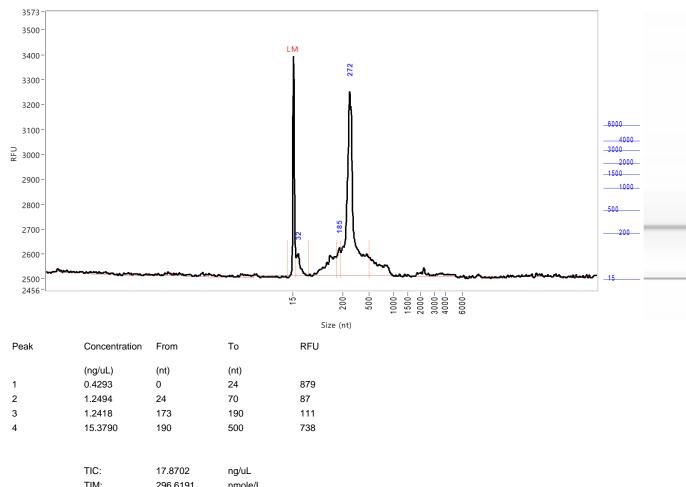


```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                         Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                   Manual baseline end (min): 38
Marker peak width (sec): 6
                            Marker min peak height: 100
                                                           Marker baseline V to V?: Y
                                                                                        Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
Quantification using: Ladder
                                     Final concentration (ng/uL): 8.0000
                                                                                  Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: E2_TP4 Well location: G8

Created: Tuesday, December 08, 2020 4:16:51 PM



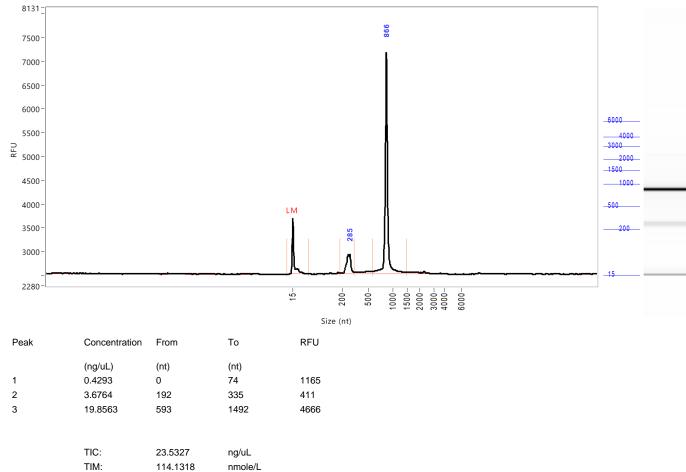
```
TIM: 296.6191 nmole/L
Total 23.4499 ng/uL

28s/18s: 0.0
RQN 1.3
```

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Sample filter: Binomial Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Manual baseline start (min): 15 Manual baseline end (min): 38 Marker min peak height: 100 Marker peak width (sec): 6 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Dilution factor: 12.0 Quantification using: Ladder Final concentration (ng/uL): 8.0000 Minimum RFU for data processing: 2

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Sample: H2_TP1 Well location: G9



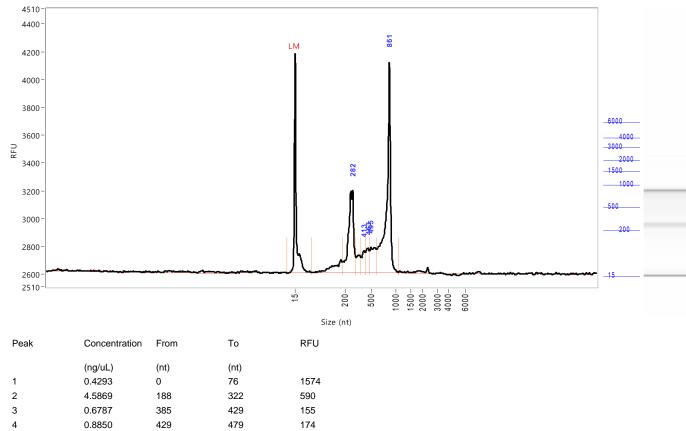
```
Total 26.3041 ng/uL concentration: 28s/18s: 0.0 RQN 1.3
```

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                           Sample baseline V to V?: Y
                                                                                         Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                   Manual baseline end (min): 38
Marker peak width (sec): 6
                             Marker min peak height: 100
                                                           Marker baseline V to V?: Y
                                                                                         Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                                                                  Dilution factor: 12.0
Quantification using: Ladder
                                     Final concentration (ng/uL): 8.0000
Minimum RFU for data processing: 2
```

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Sample: H2_TP2 Well location: G10

Created: Tuesday, December 08, 2020 4:16:51 PM



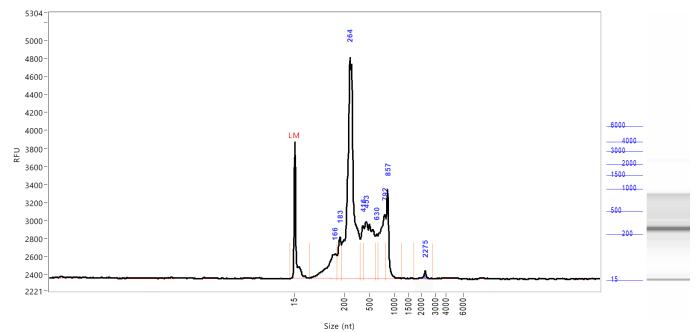
182 1513

| 4 | 0.8850 | 429 | 479 |
|---|----------------|----------|---------|
| 5 | 1.4267 | 479 | 603 |
| 6 | 7.6283 | 603 | 1073 |
| | | | |
| | | | |
| | TIC: | 15.2057 | ng/uL |
| | TIM: | 104.7543 | nmole/L |
| | Total | 17.1843 | ng/uL |
| | concentration: | | |
| | 28s/18s: | 0.0 | |
| | RQN | 1.1 | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                            Marker min peak height: 100
                                                          Marker baseline V to V?: Y
                                                                                        Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                                                                  Dilution factor: 12.0
Quantification using: Ladder
                                     Final concentration (ng/uL): 8.0000
Minimum RFU for data processing: 2
```

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Sample: H2_TP3 Well location: G11

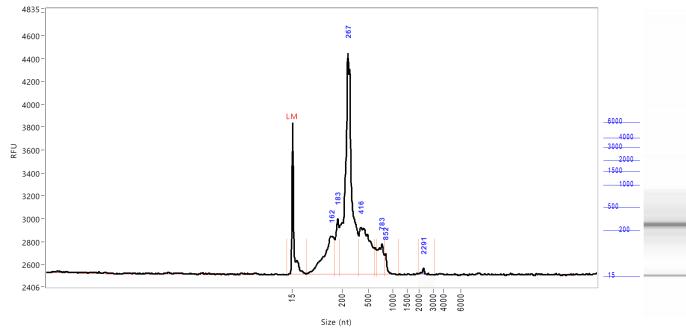


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 69 | 1508 |
| 2 | 4.3117 | 69 | 172 | 269 |
| 3 | 2.2667 | 172 | 188 | 456 |
| 4 | 24.0645 | 188 | 385 | 2451 |
| 5 | 2.6163 | 385 | 429 | 583 |
| 6 | 7.9886 | 429 | 607 | 627 |
| 7 | 1.4603 | 607 | 653 | 490 |
| 8 | 5.7293 | 653 | 820 | 706 |
| 9 | 3.2684 | 820 | 1234 | 981 |
| 10 | 0.2309 | 1723 | 2839 | 84 |
| | | | | |
| | | | | |
| | TIC: | 51.9368 | ng/uL | |
| | TIM: | 520.6578 | nmole/L | |
| | Total concentration: | 51.1266 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                       Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: H2_TP4 Well location: G12

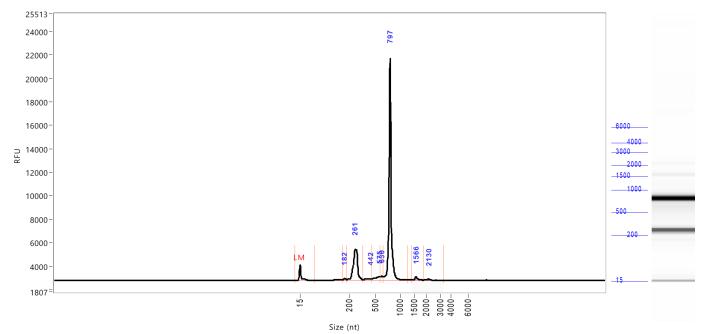


| | | | | (, |
|------|----------------------|----------|---------|------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 65 | 1314 |
| 2 | 6.6048 | 65 | 172 | 324 |
| 3 | 2.8896 | 172 | 189 | 478 |
| 4 | 23.0941 | 189 | 390 | 1927 |
| 5 | 7.5526 | 390 | 626 | 406 |
| 6 | 2.3626 | 677 | 834 | 260 |
| 7 | 0.6156 | 834 | 1202 | 175 |
| 8 | 0.1434 | 1963 | 3110 | 51 |
| | | | | |
| | | | | |
| | TIC: | 43.2627 | ng/uL | |
| | TIM: | 520.0760 | nmole/L | |
| | Total concentration: | 43.5803 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A3_TP1 Well location: H1



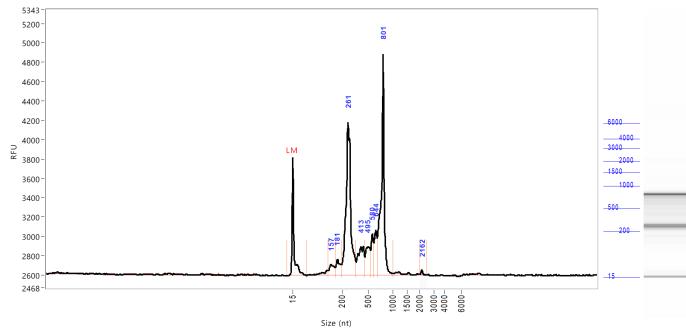
| Peak | Concentration | From | То | RFU |
|------|----------------|----------------|---------|-------|
| | (ng/ul.) | (nt) | (nt) | |
| | (ng/uL) | (nt) | (nt) | 1010 |
| 1 | 0.4293 | 0 | 69 | 1316 |
| 2 | 0.6762 | 174 | 188 | 162 |
| 3 | 20.7918 | 188 | 356 | 2688 |
| 4 | 1.4495 | 356 | 455 | 150 |
| 5 | 2.7777 | 455 | 598 | 308 |
| 6 | 1.4538 | 598 | 658 | 351 |
| 7 | 62.4317 | 658 | 1283 | 18949 |
| 8 | 1.4803 | 1411 | 1871 | 297 |
| 9 | 0.6245 | 1871 | 3262 | 137 |
| | | | | |
| | | | | |
| | TIC: | 91.6856 | ng/uL | |
| | TIM: | 541.8409 | nmole/L | |
| | Total | 92.8744 | ng/uL | |
| | concentration: | | | |
| | 28s/18s: | 0.4 | | |
| | RQN | 1.3 | | |
| | | · - | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A3_TP2 Well location: H2

Created: Tuesday, December 08, 2020 4:16:51 PM

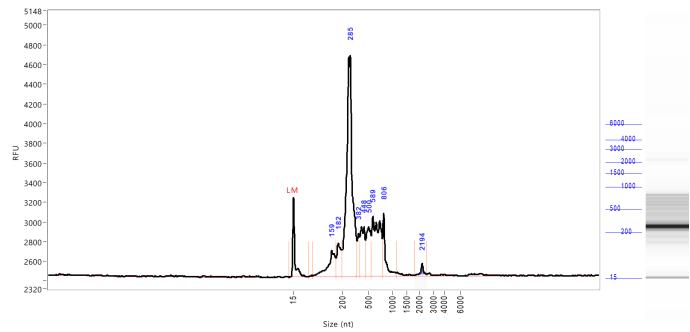


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 66 | 1206 |
| 2 | 1.0679 | 146 | 173 | 107 |
| 3 | 1.2464 | 173 | 196 | 161 |
| 4 | 14.5516 | 196 | 356 | 1575 |
| 5 | 3.2788 | 356 | 458 | 292 |
| 6 | 2.4808 | 458 | 543 | 293 |
| 7 | 1.8571 | 543 | 607 | 421 |
| 8 | 2.2563 | 607 | 677 | 461 |
| 9 | 11.8345 | 677 | 1009 | 2282 |
| 10 | 0.1540 | 2001 | 2517 | 51 |
| | | | | |
| | TIC: | 38.7274 | ng/uL | |
| | TIM: | 321.4461 | nmole/L | |
| | Total concentration: | 39.3530 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

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Sample: A3_TP3 Well location: H3

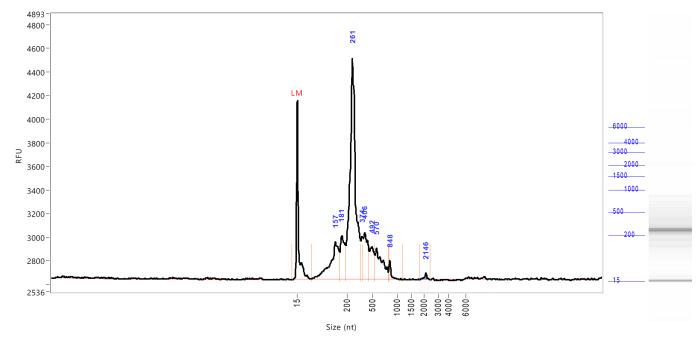


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 71 | 802 |
| 2 | 5.7062 | 88 | 174 | 262 |
| 3 | 4.4597 | 174 | 197 | 335 |
| 4 | 36.5180 | 197 | 364 | 2248 |
| 5 | 2.8159 | 364 | 395 | 433 |
| 6 | 6.4792 | 395 | 466 | 504 |
| 7 | 5.9122 | 466 | 552 | 505 |
| 8 | 12.6758 | 552 | 783 | 610 |
| 9 | 4.3560 | 783 | 1122 | 636 |
| 10 | 0.8706 | 1787 | 2500 | 125 |
| | | | | |
| | | | | |
| | TIC: | 79.7935 | ng/uL | |
| | TIM: | 790.7545 | nmole/L | |
| | Total concentration: | 80.4767 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.2 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: A3_TP4 Well location: H4

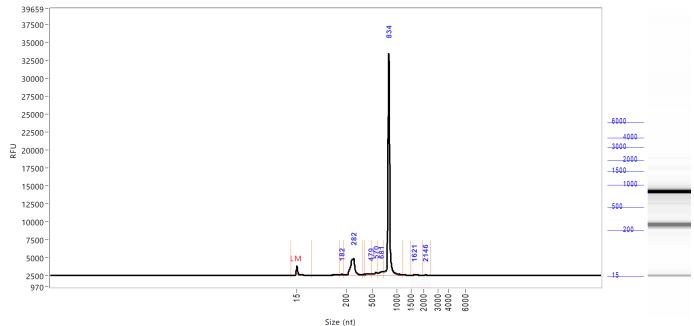


| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 66 | 1514 |
| 2 | 4.3241 | 66 | 172 | 313 |
| 3 | 2.5580 | 172 | 195 | 367 |
| 4 | 15.8464 | 195 | 358 | 1869 |
| 5 | 1.2004 | 358 | 387 | 360 |
| 6 | 2.6111 | 387 | 461 | 392 |
| 7 | 1.7074 | 461 | 538 | 269 |
| 8 | 2.4852 | 538 | 815 | 257 |
| 9 | 0.3559 | 829 | 1170 | 153 |
| 10 | 0.1129 | 1834 | 2452 | 48 |
| | | | | |
| | TIO | 04 0045 | | |
| | TIC: | 31.2015 | ng/uL | |
| | TIM: | 375.7154 | nmole/L | |
| | Total concentration: | 30.9647 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: B3_TP1 Well location: H5



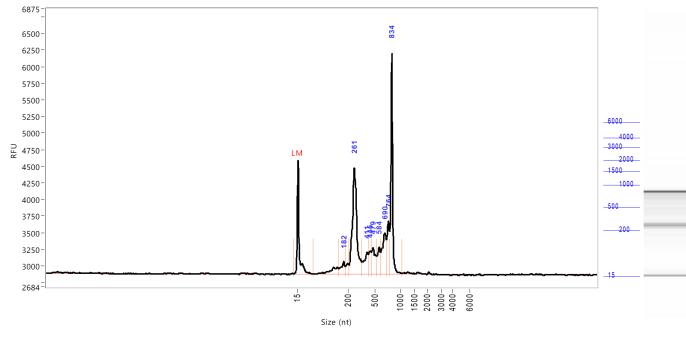
| | | | | Size (nt) |
|------|----------------------|----------|---------|-----------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 72 | 1349 |
| 2 | 0.6204 | 173 | 188 | 159 |
| 3 | 18.9666 | 188 | 385 | 2352 |
| 4 | 1.9322 | 416 | 492 | 231 |
| 5 | 2.4743 | 492 | 607 | 448 |
| 6 | 3.8226 | 607 | 732 | 536 |
| 7 | 79.4421 | 732 | 1178 | 30932 |
| 8 | 0.7573 | 1468 | 1945 | 127 |
| 9 | 0.3018 | 1945 | 2517 | 82 |
| | | | | |
| | | | | |
| | TIC: | 108.3175 | ng/uL | |
| | TIM: | 573.6045 | nmole/L | |
| | Total concentration: | 110.1243 | ng/uL | |
| | 28s/18s: | 0.3 | | |
| | RQN | 1.1 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                            Marker min peak height: 100
                                                          Marker baseline V to V?: Y
                                                                                        Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                  Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: B3_TP2 Well location: H6

Created: Tuesday, December 08, 2020 4:16:51 PM



| Peak | Concentration | From | То | RFU |
|------|----------------------|----------|---------|------|
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 70 | 1713 |
| 2 | 0.9675 | 163 | 187 | 186 |
| 3 | 9.5938 | 201 | 353 | 1604 |
| 4 | 1.6944 | 353 | 424 | 330 |
| 5 | 1.2215 | 424 | 461 | 343 |
| 6 | 1.9505 | 461 | 538 | 395 |
| 7 | 1.4076 | 538 | 612 | 403 |
| 8 | 3.1043 | 612 | 732 | 617 |
| 9 | 2.2254 | 732 | 792 | 797 |
| 10 | 7.1371 | 792 | 1057 | 3329 |
| | | | | |
| | TIC | 20.2022 | ng/ul | |
| | TIC: | 29.3022 | ng/uL | |
| | TIM: | 220.9710 | nmole/L | |
| | Total concentration: | 30.7722 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | | | | |

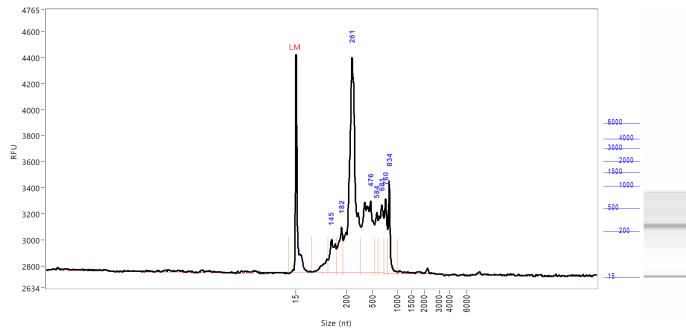
Sample peak width (sec): 6 Sample min peak height: 50 Sample baseline V to V?: Y Sample baseline V to V points: 3 Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40 Sample filter: Binomial Manual baseline start (min): 15 Manual baseline end (min): 38 Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y Marker baseline V to V points: 3 Lower marker selection: First peak > 100 RFU Upper marker selection: Last peak > 100 RFU Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000 Final concentration (ng/uL): 8.0000Quantification using: Ladder Dilution factor: 12.0 Minimum RFU for data processing: 2

RQN

1.0

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Sample: B3_TP3 Well location: H7



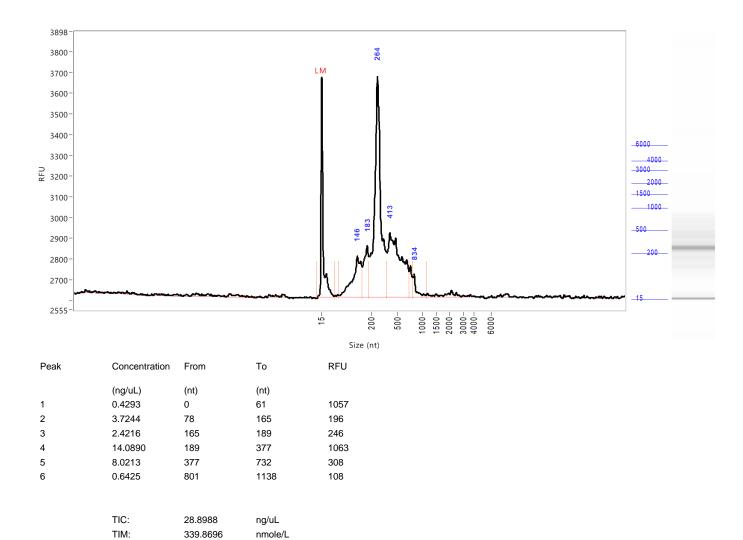
| | | | | () |
|------|----------------------|----------|---------|------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 73 | 1668 |
| 2 | 2.0009 | 133 | 164 | 255 |
| 3 | 2.1408 | 164 | 188 | 345 |
| 4 | 14.3479 | 188 | 369 | 1646 |
| 5 | 7.0217 | 369 | 538 | 551 |
| 6 | 1.7878 | 538 | 612 | 466 |
| 7 | 2.8992 | 612 | 732 | 522 |
| 8 | 1.8764 | 732 | 806 | 566 |
| 9 | 1.6889 | 806 | 995 | 706 |
| | TIC: | 33.7637 | ng/uL | |
| | TIM: | 332.2534 | nmole/L | |
| | Total concentration: | 34.3995 | ng/uL | |
| | 28s/18s: | 0.0 | | |
| | RQN | 1.1 | | |
| | | | | |

```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6 Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                      Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                          Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: B3_TP4 Well location: H8

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```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                           Sample baseline V to V?: Y
                                                                                         Sample baseline V to V points: 3
Sample filter: Binomial
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Manual baseline start (min): 15
                                   Manual baseline end (min): 38
                             Marker min peak height: 100
Marker peak width (sec): 6
                                                           Marker baseline V to V?: Y
                                                                                         Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                                                                  Dilution factor: 12.0
Quantification using: Ladder
                                     Final concentration (ng/uL): 8.0000
Minimum RFU for data processing: 2
```

Total

RQN

concentration: 28s/18s:

29.9544

0.0

1.1

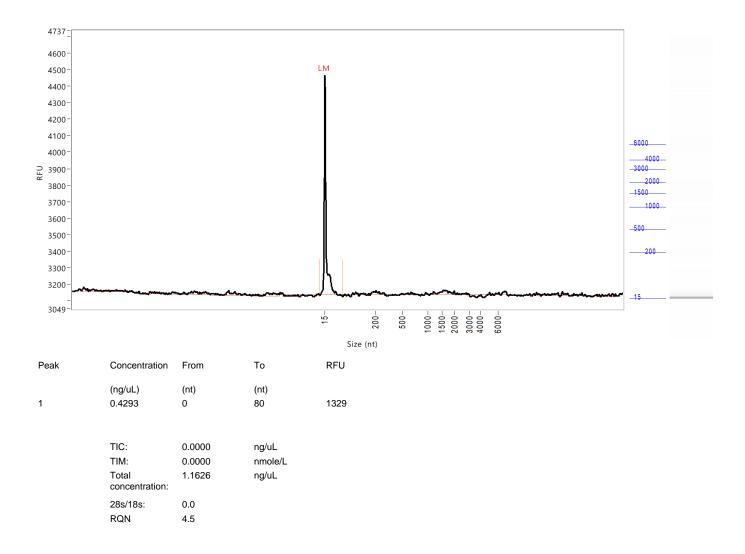
ng/uL

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Sample: H9:

Well location: Tuesday, December 08, 2020 4:16:51 PM

Created: 96



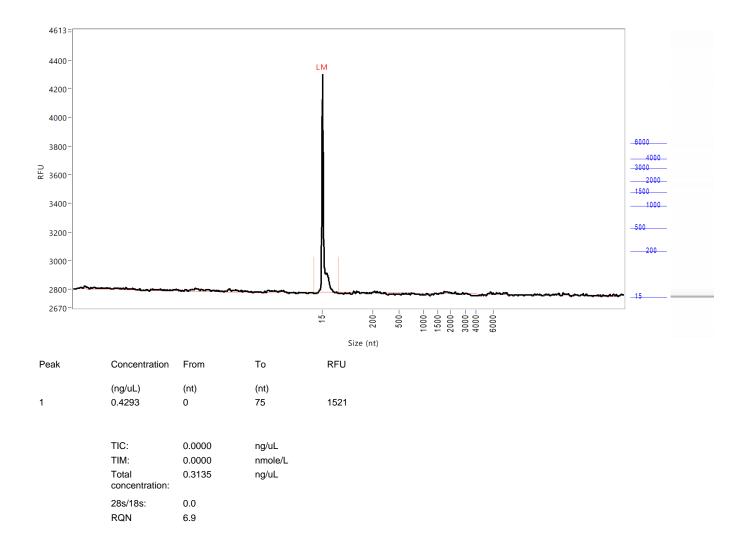
```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100
                                                          Marker baseline V to V?: Y
                                                                                        Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: H10:

Well location: Tuesday, December 08, 2020 4:16:51 PM

Created: 96



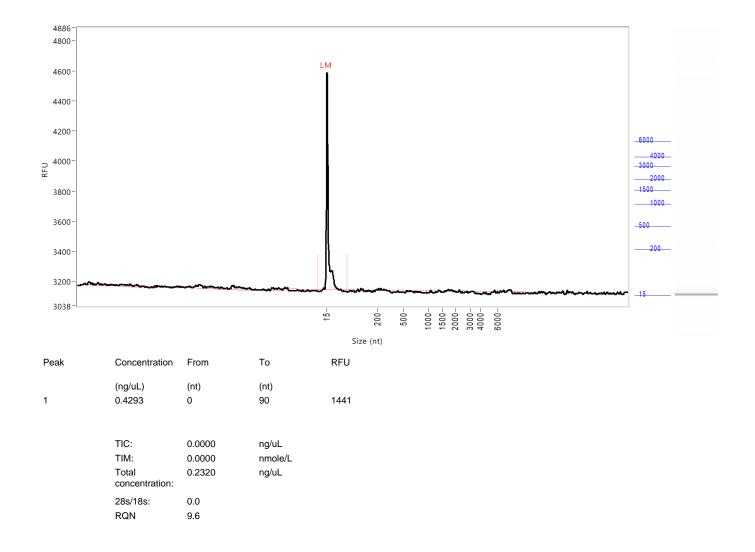
```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100
                                                          Marker baseline V to V?: Y
                                                                                        Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: H11:

Well location: Tuesday, December 08, 2020 4:16:51 PM

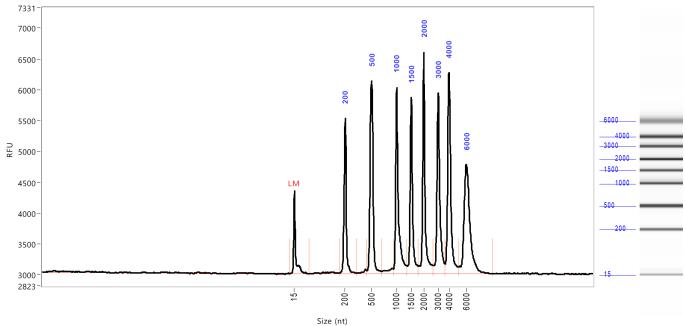
Created: 96



```
Sample peak width (sec): 6
                             Sample min peak height: 50
                                                          Sample baseline V to V?: Y
                                                                                        Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 15
                                  Manual baseline end (min): 38
Marker peak width (sec): 6
                           Marker min peak height: 100 Marker baseline V to V?: Y
                                                                                        Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                           Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                     Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                 Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: Ladder Well location: H12



| | | | | 5.25 (11) |
|------|----------------------|----------|---------|-----------|
| Peak | Concentration | From | То | RFU |
| | (ng/uL) | (nt) | (nt) | |
| 1 | 0.4293 | 0 | 69 | 1343 |
| 2 | 9.2998 | 180 | 335 | 2521 |
| 3 | 14.8581 | 448 | 700 | 3117 |
| 4 | 12.2400 | 931 | 1363 | 3017 |
| 5 | 9.8182 | 1363 | 1778 | 2859 |
| 6 | 11.6404 | 1778 | 2646 | 3590 |
| 7 | 11.2380 | 2646 | 3631 | 2926 |
| 8 | 12.9825 | 3631 | 5055 | 3254 |
| 9 | 12.9498 | 5055 | 9099 | 1773 |
| | 710 | | | |
| | TIC: | 95.0269 | ng/uL | |
| | TIM: | 337.7375 | nmole/L | |
| | Total concentration: | 96.0000 | ng/uL | |

```
Sample peak width (sec): 6
                              Sample min peak height: 200 \,\, Sample baseline V to V?: Y \,\,
                                                                                         Sample baseline V to V points: 3
                             Number of points for filter: 9 Sample start region (min): 0 Sample end region (min): 40
Sample filter: Binomial
Manual baseline start (min): 18
                                   Manual baseline end (min): 38
Marker peak width (sec): 6
                            Marker min peak height: 100
                                                           Marker baseline V to V?: Y
                                                                                         Marker baseline V to V points: 3
Lower marker selection: First peak > 100 RFU
                                                            Upper marker selection: Last peak > 100 RFU
Ladder size (nt)15, 200, 500, 1000, 1500, 2000, 3000, 4000, 6000
                                      Final concentration (ng/uL): 8.0000
Quantification using: Ladder
                                                                                   Dilution factor: 12.0
Minimum RFU for data processing: 2
```

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Sample: Ladder Well location: H12

Created: Tuesday, December 08, 2020 4:16:51 PM

Fit type: Point to point

