GeneMark

Version 2.5p (09.08.06) *Copyright 1993 - M. Borodovsky, J. McIninch*

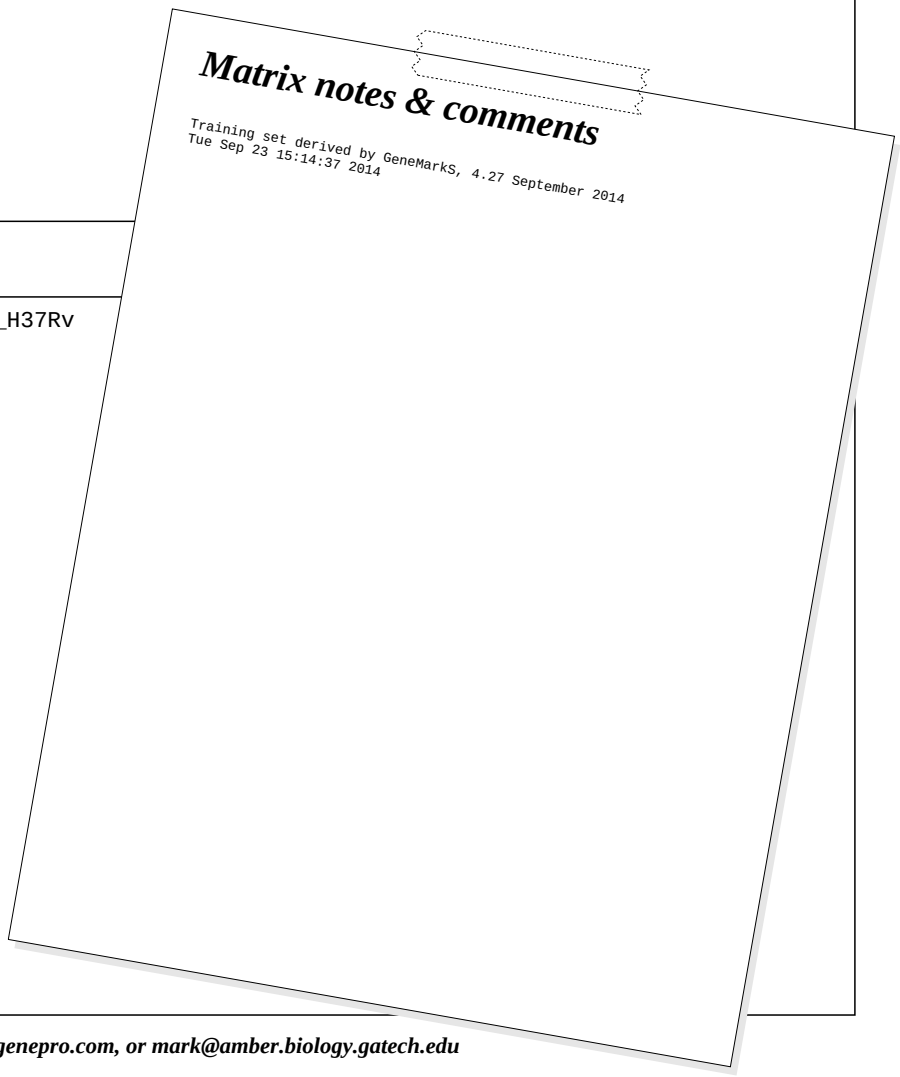
PROGRAM INFORMATION

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Analysis Date    :  3/29/24 at 0:58:35
Pages            :  32
Sequence Length  :  64606 bp
GC Content       :   59.69%

Window Length    :  96 bp
Window Step      :  12 bp
Threshold Value  :  0.500

PS-Version       :  1.2

GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

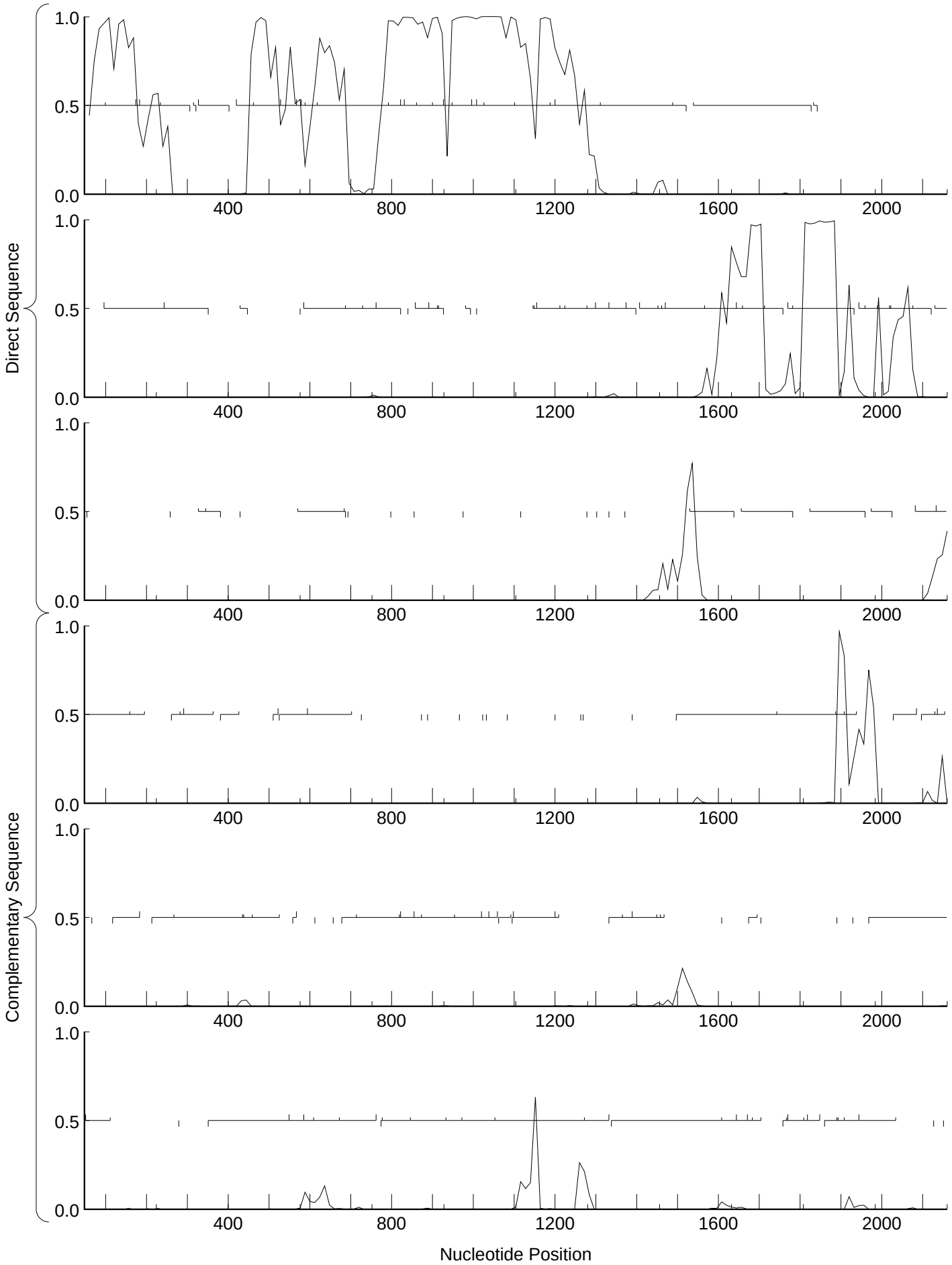
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Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

Order      :  4
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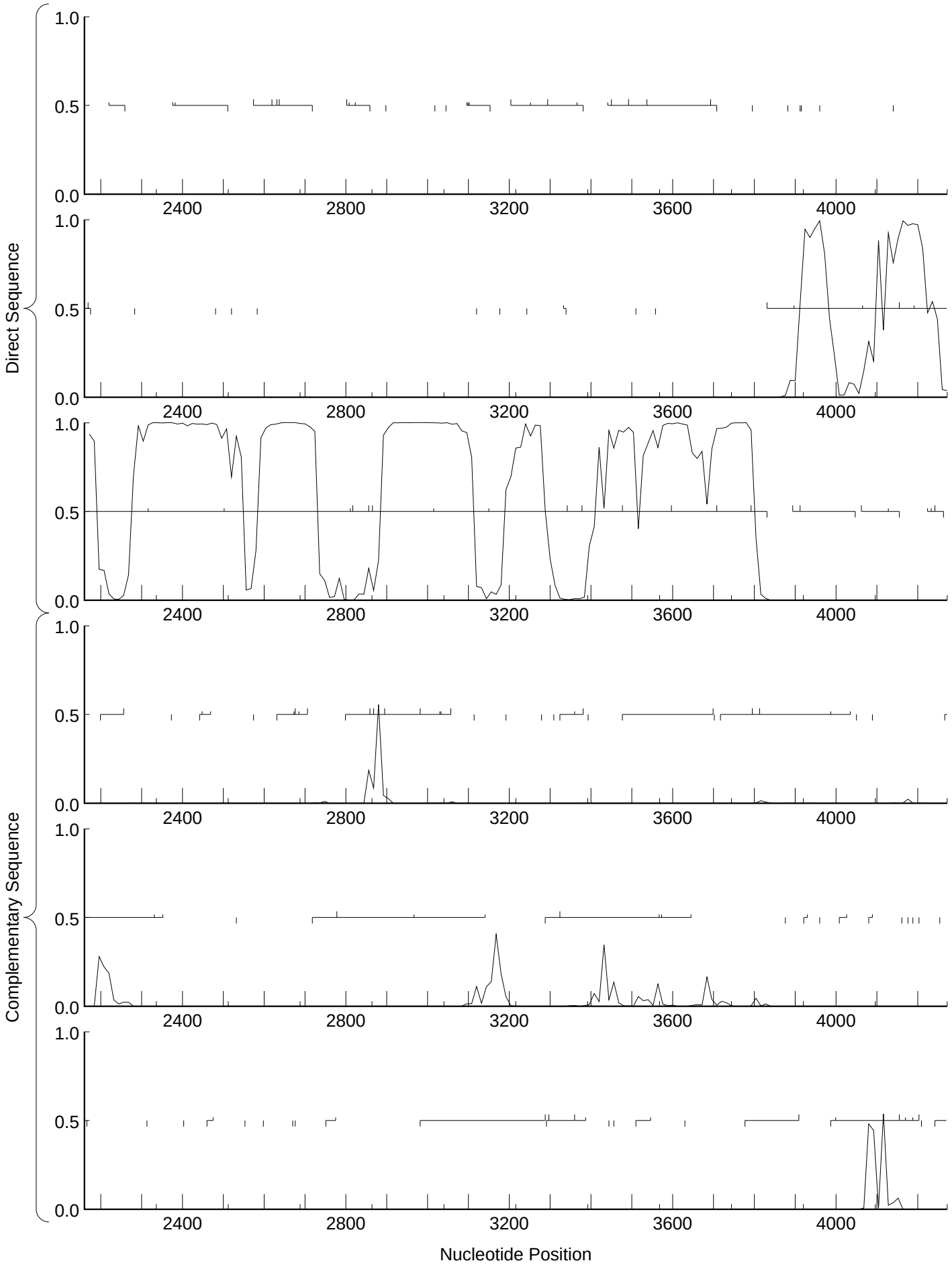

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

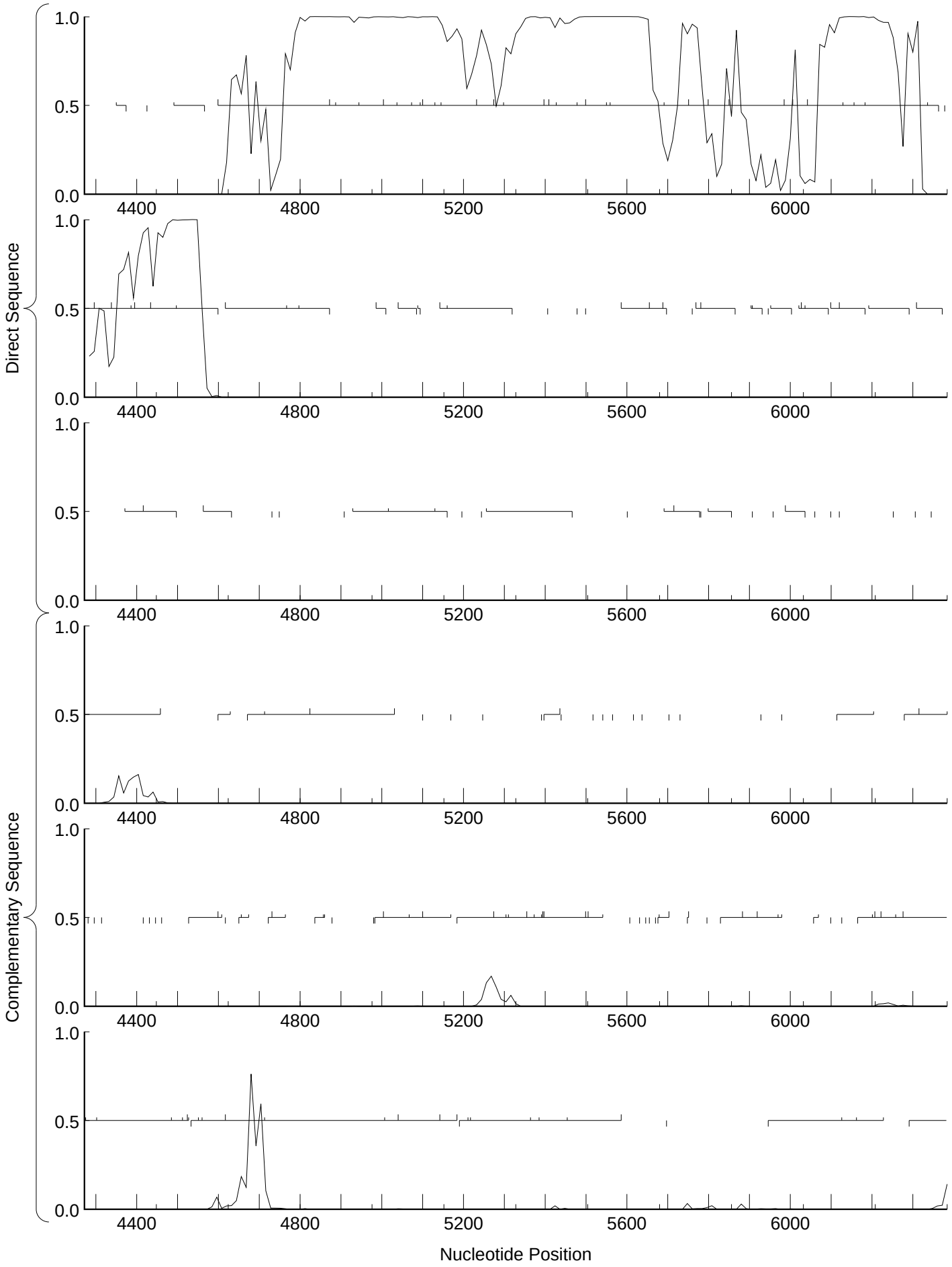
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 2/32



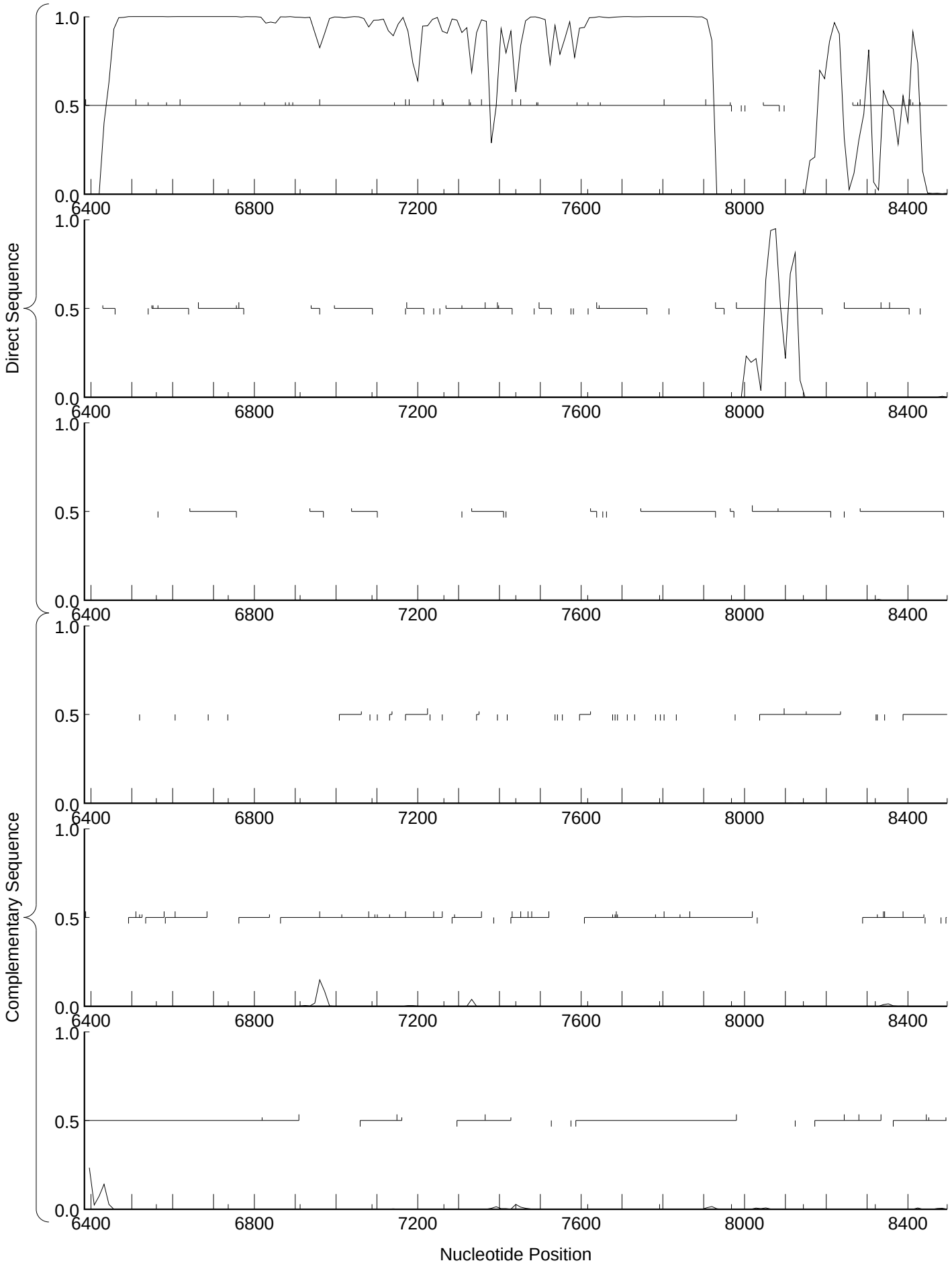
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 3/32



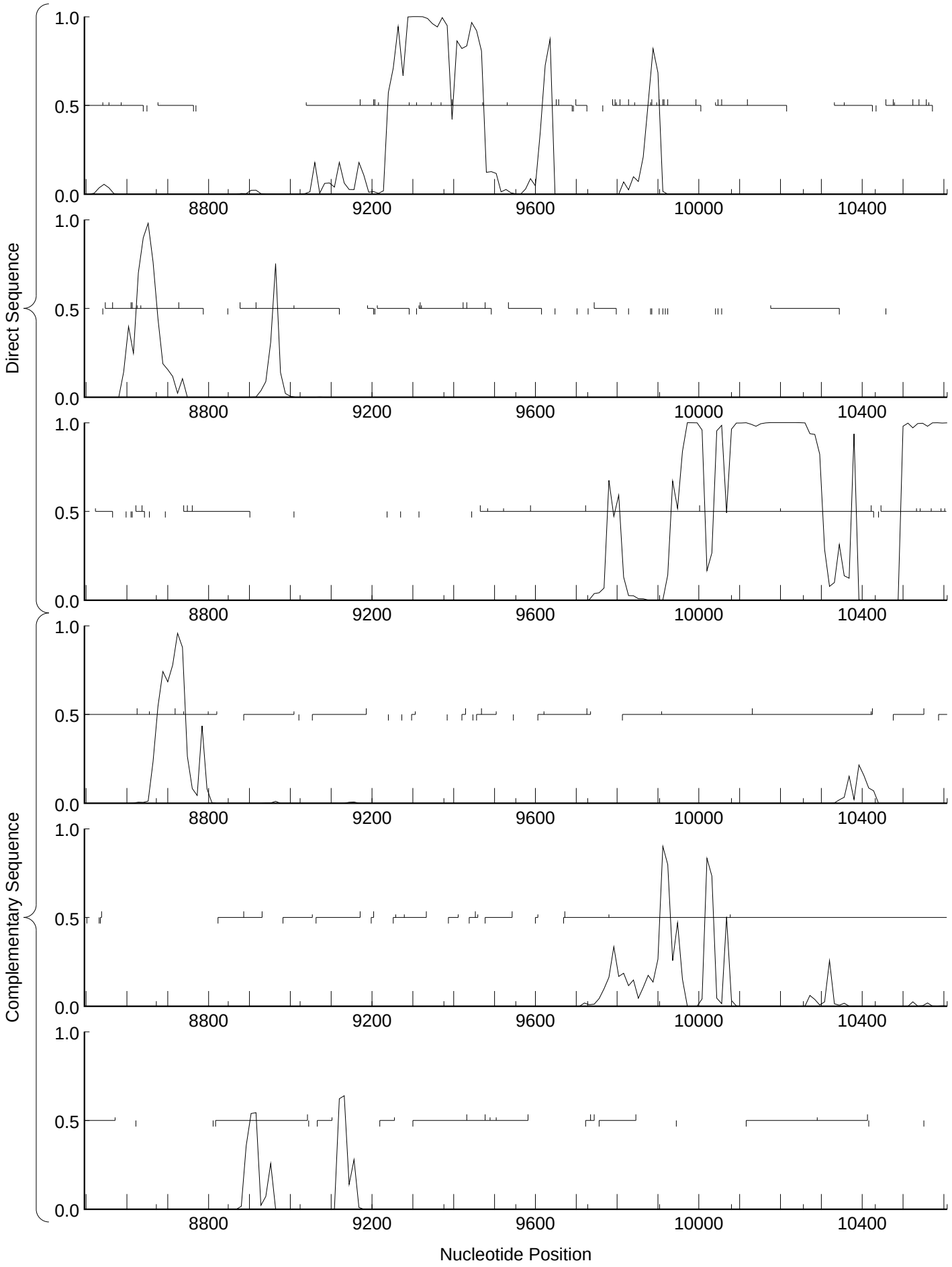
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 4/32



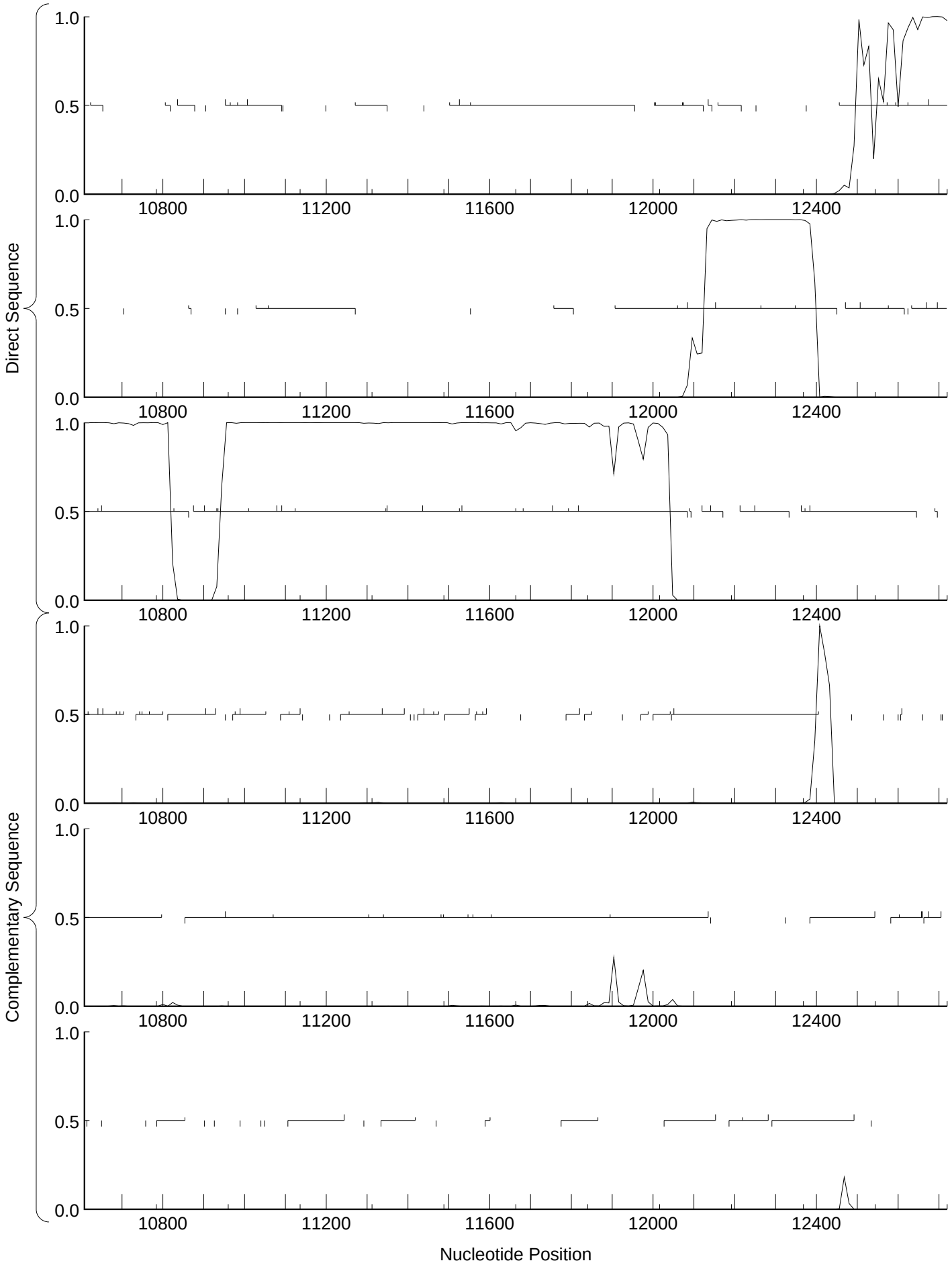
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 5/32



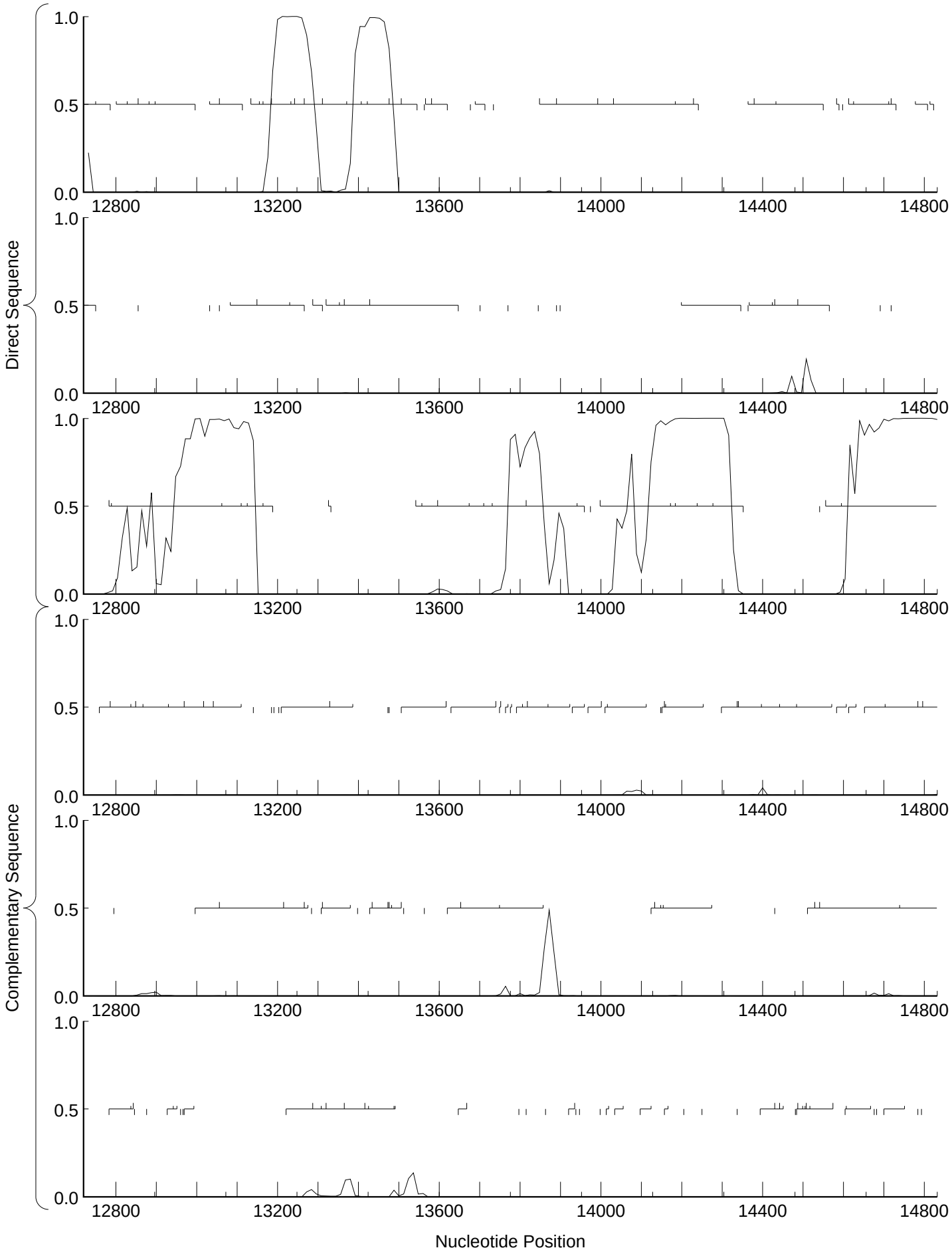
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 6/32



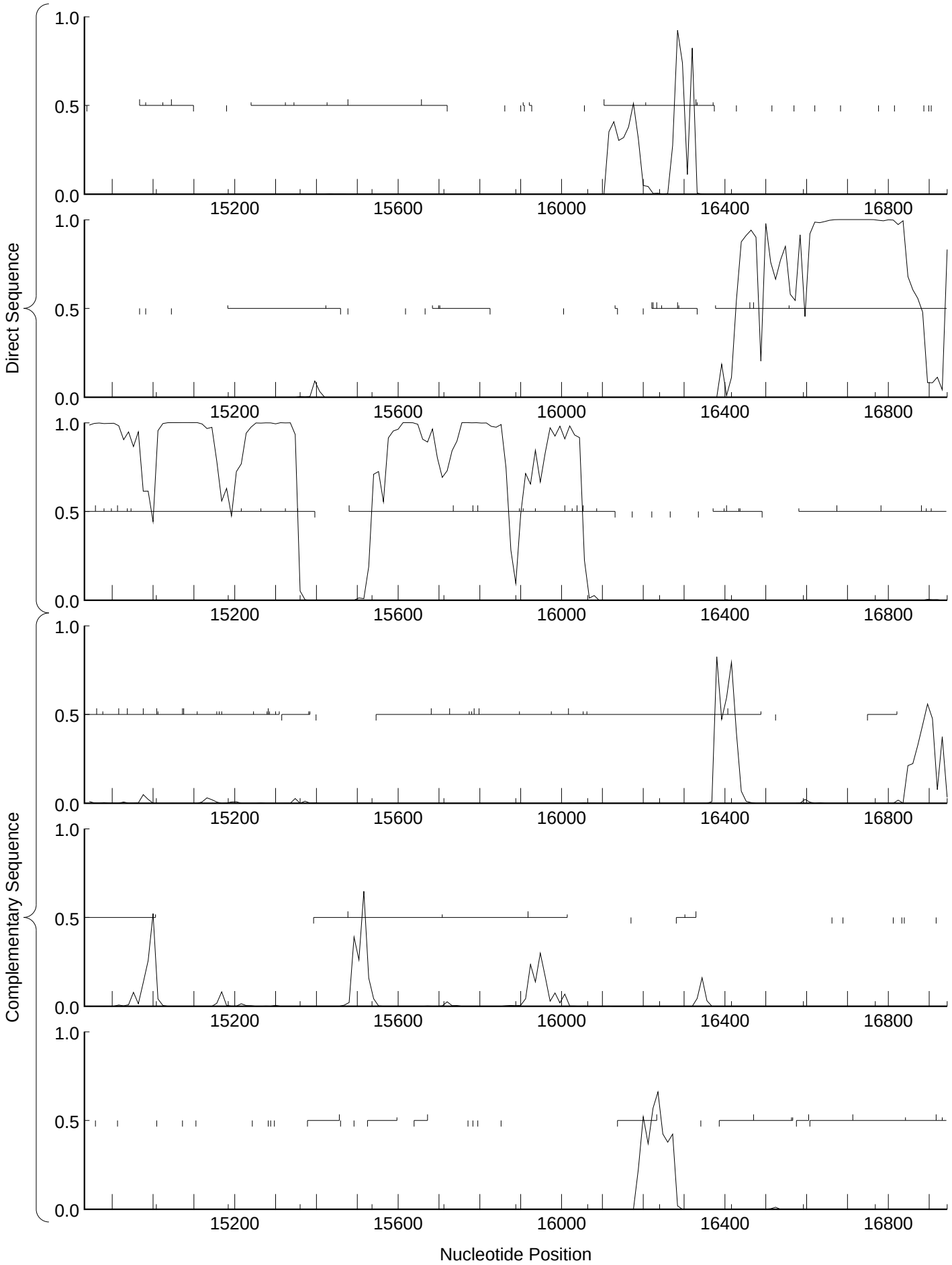
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 7/32



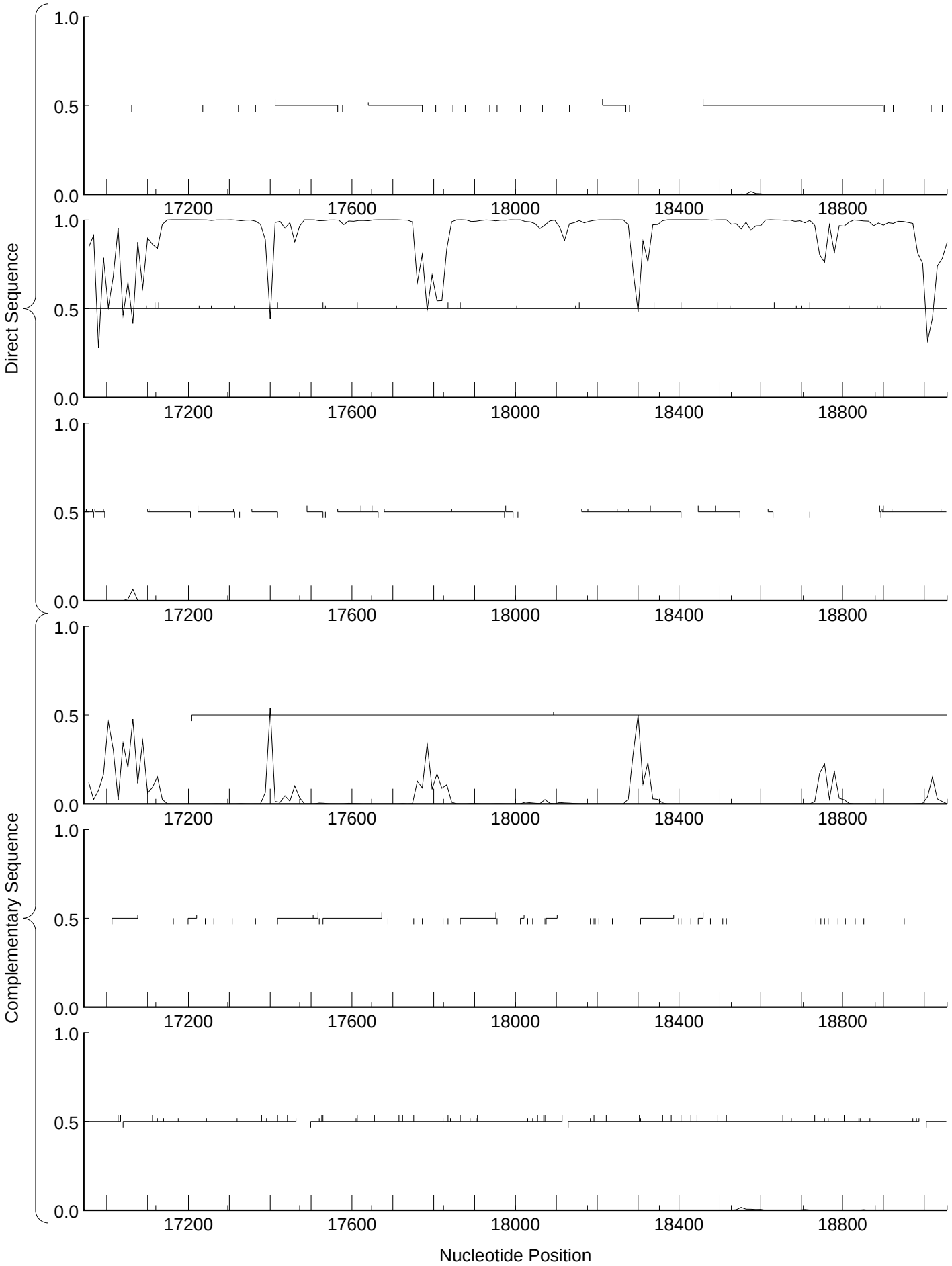
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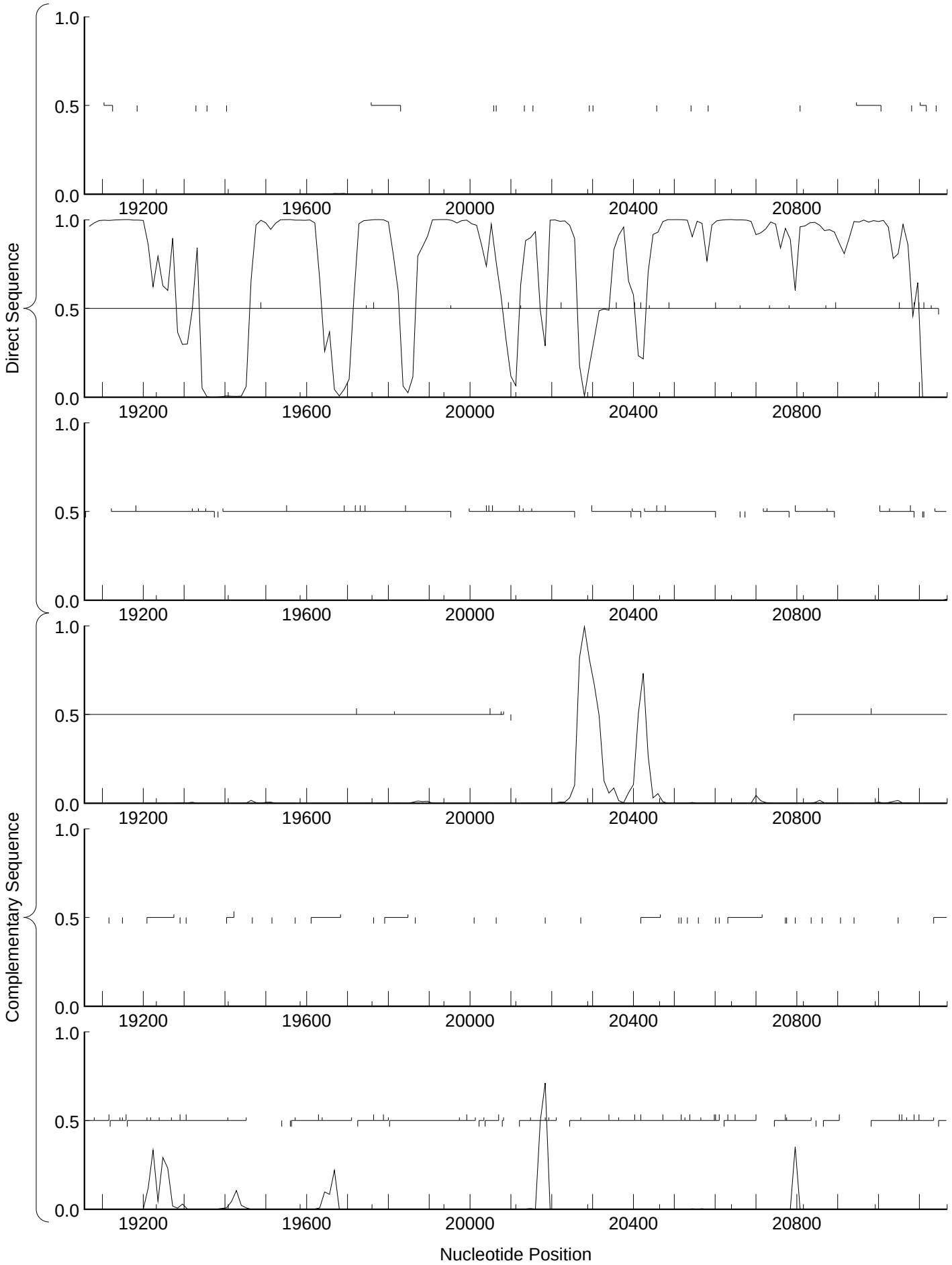
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 9/32



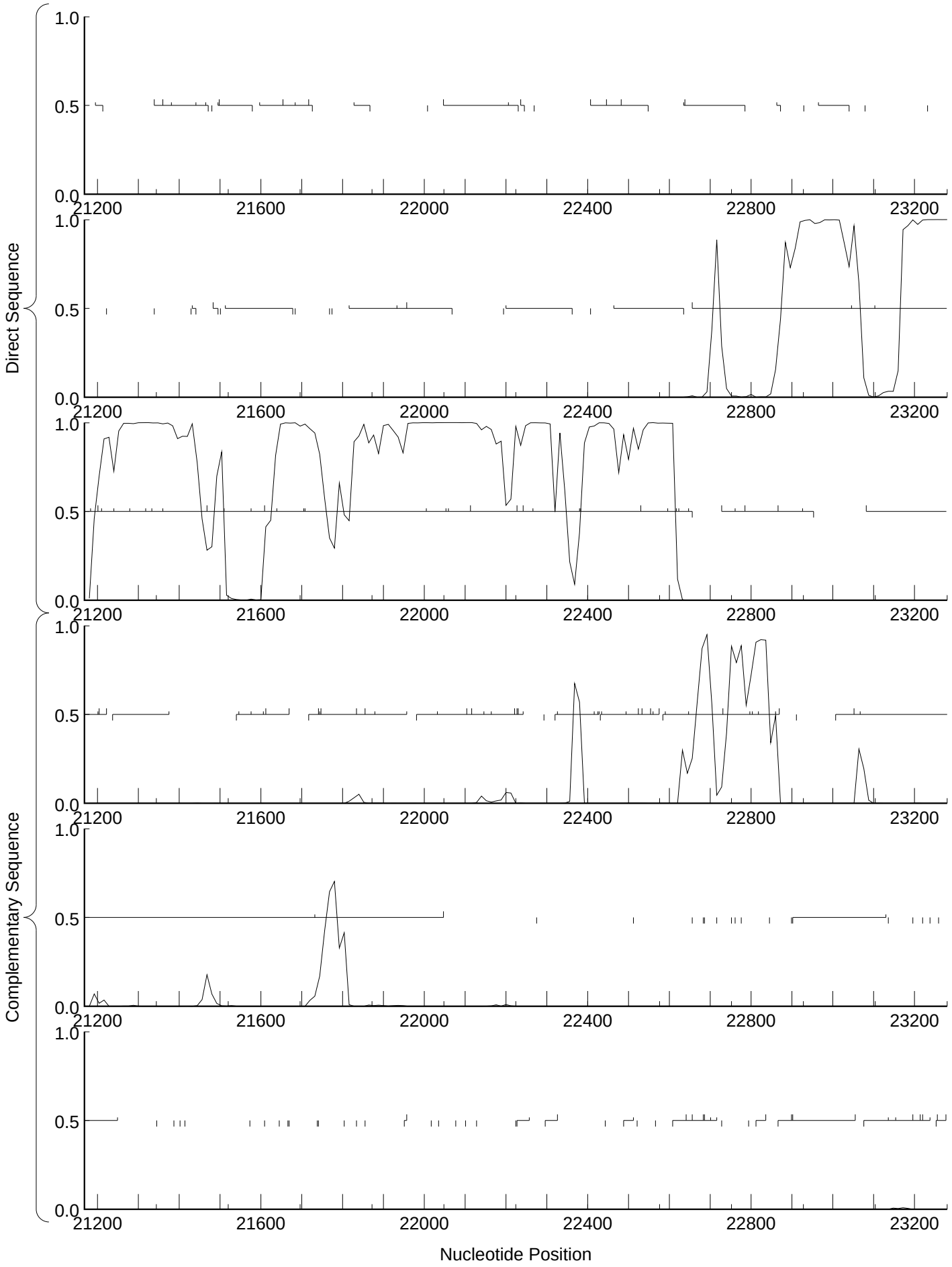
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 10/32



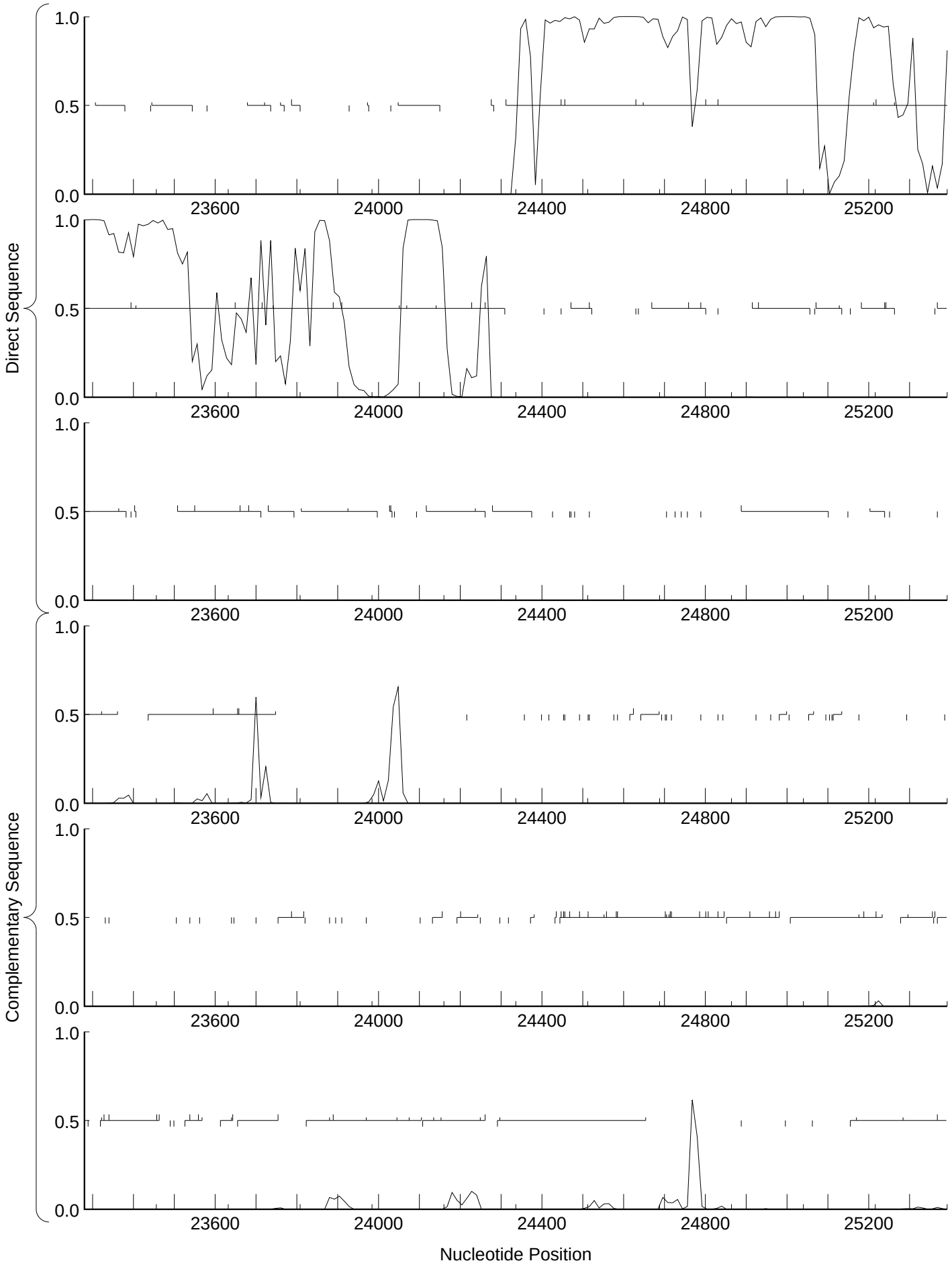
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 11/32



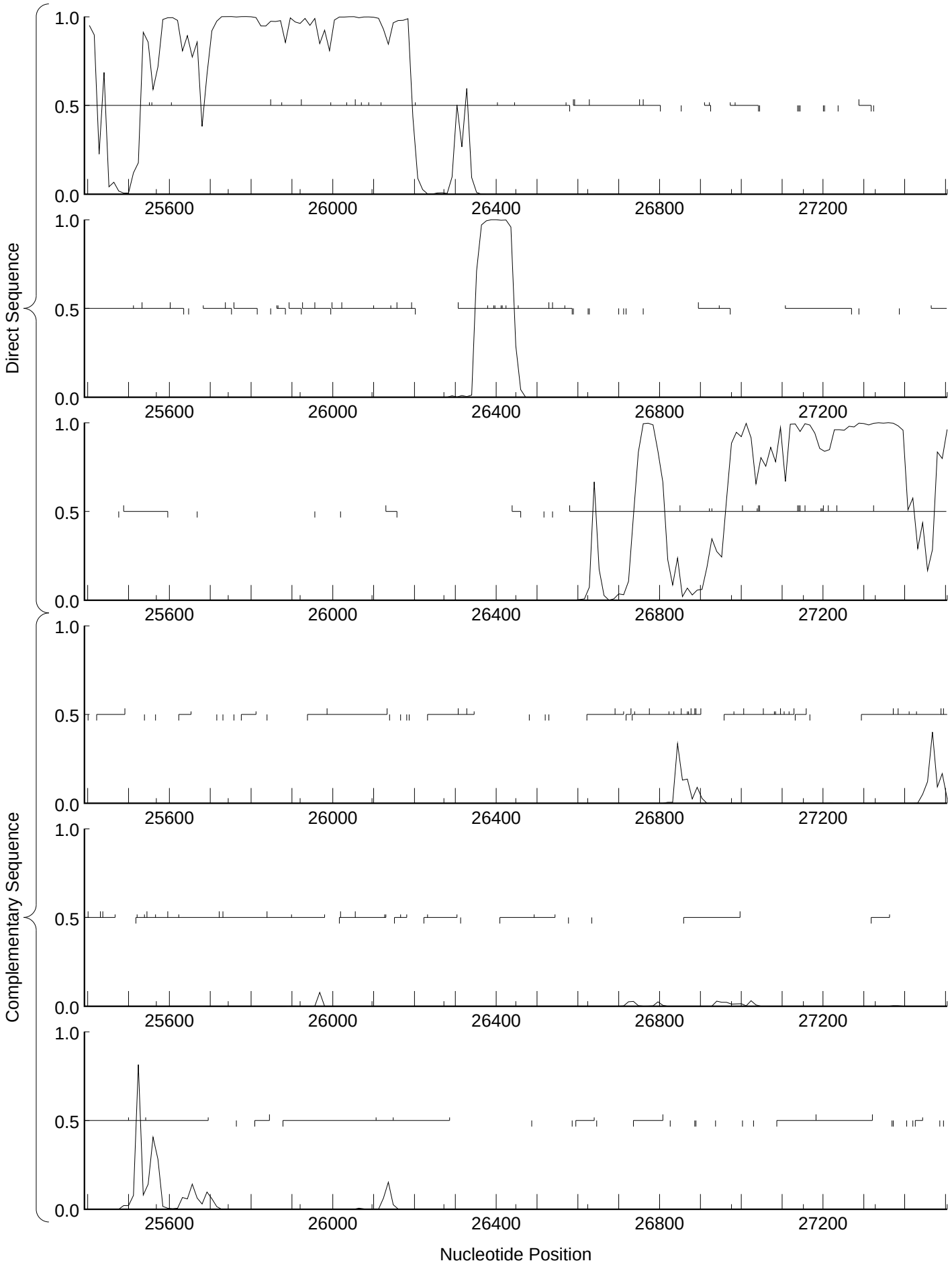
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 12/32



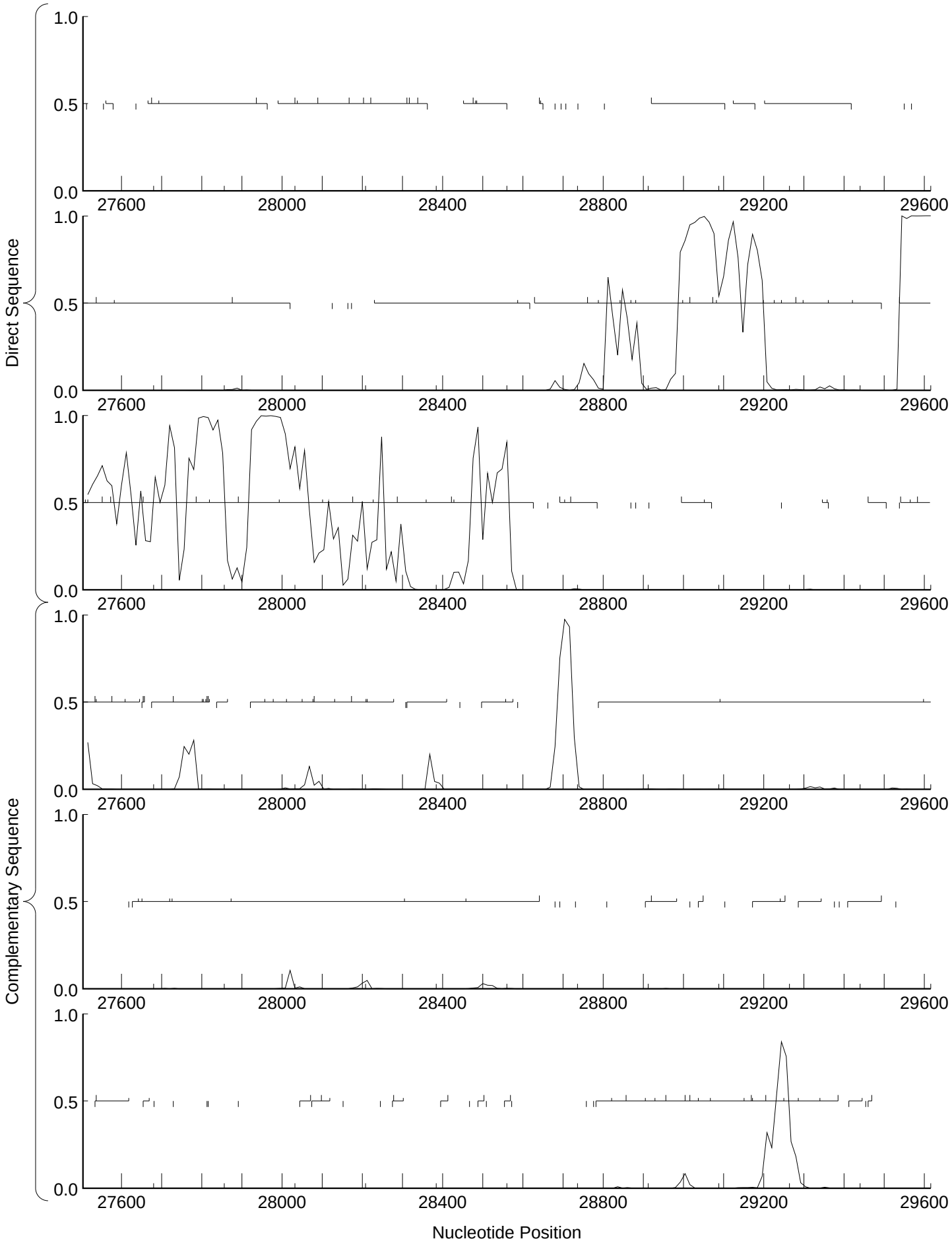
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 13/32



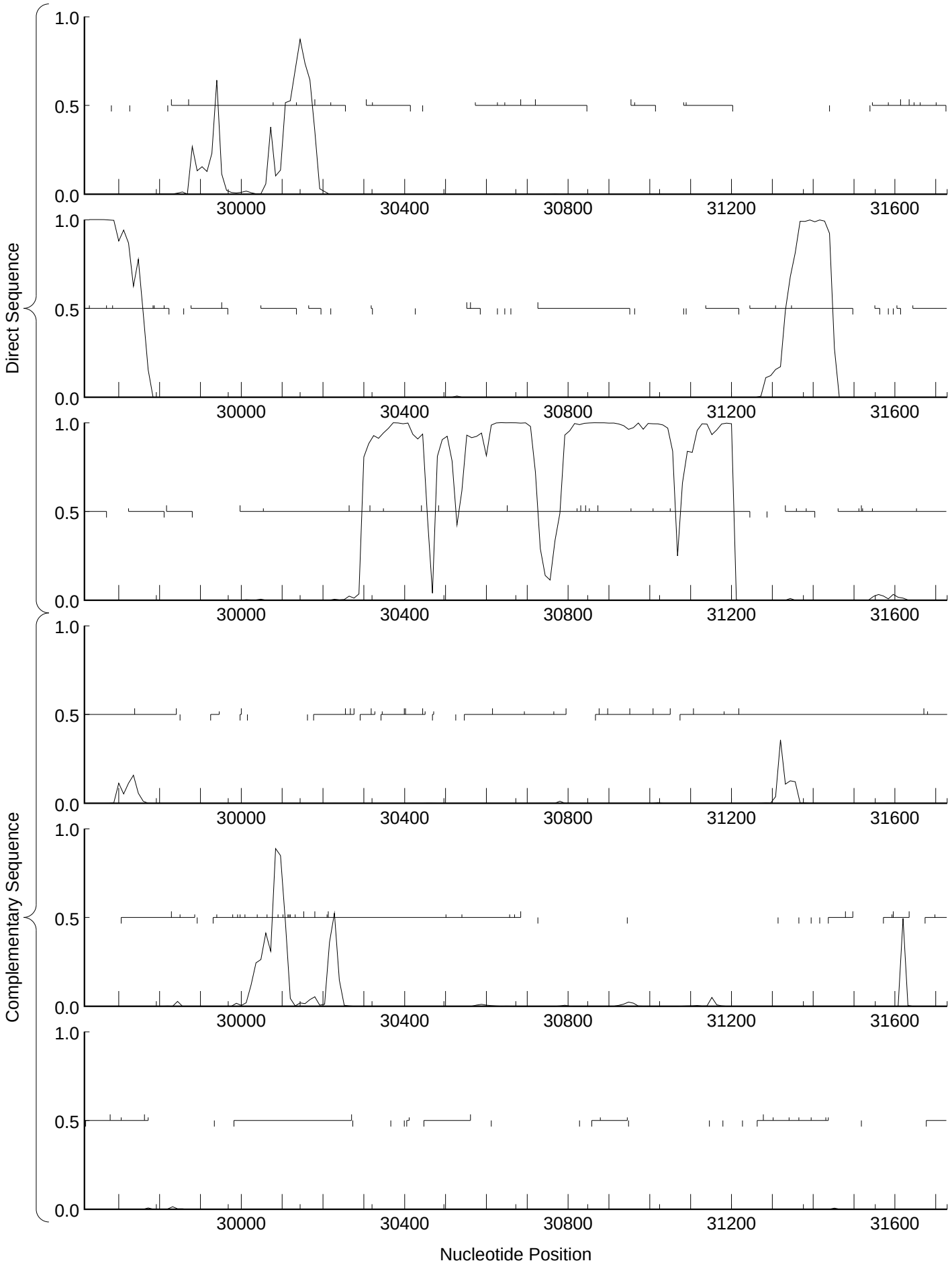
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 14/32



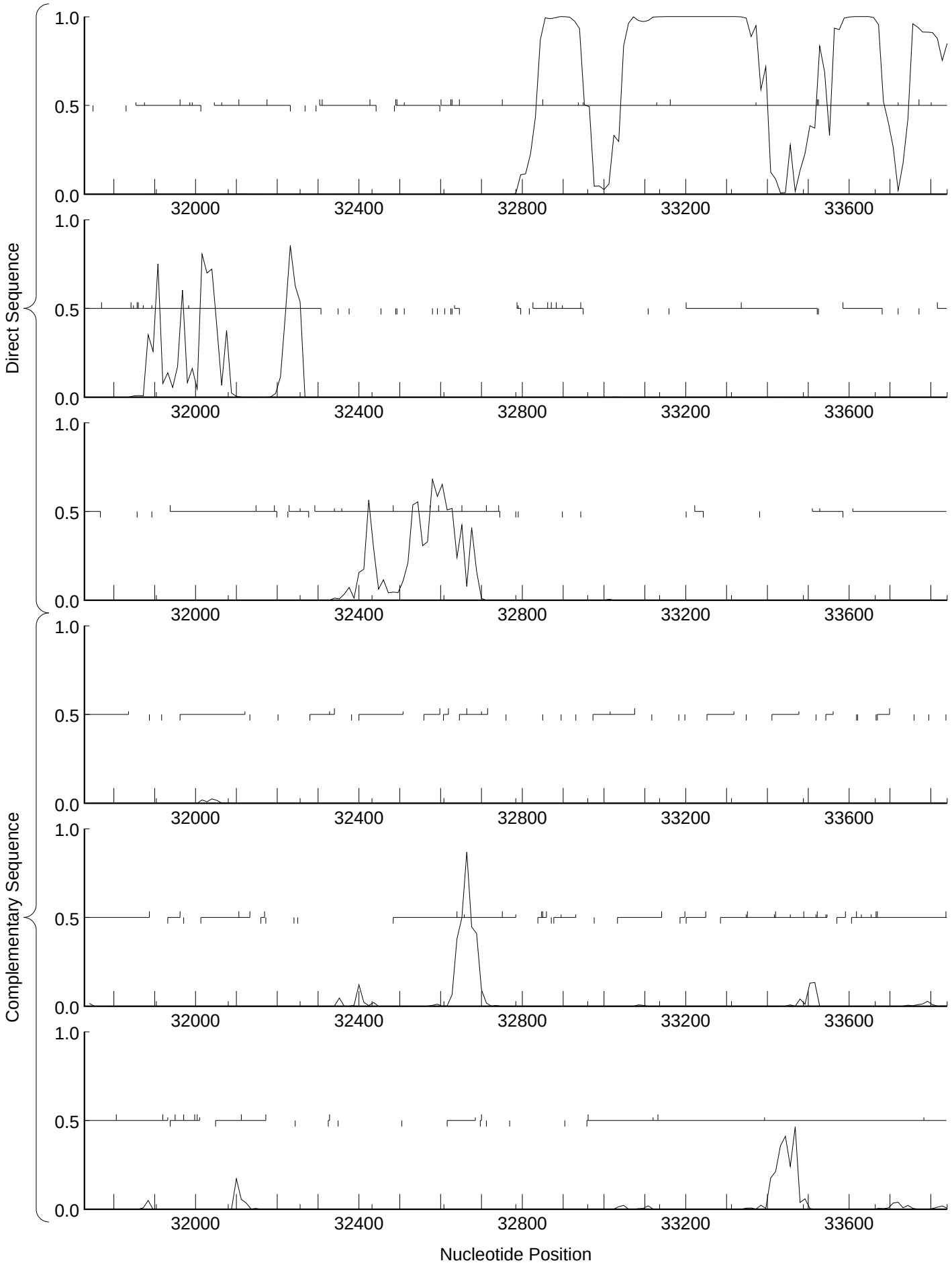
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 15/32



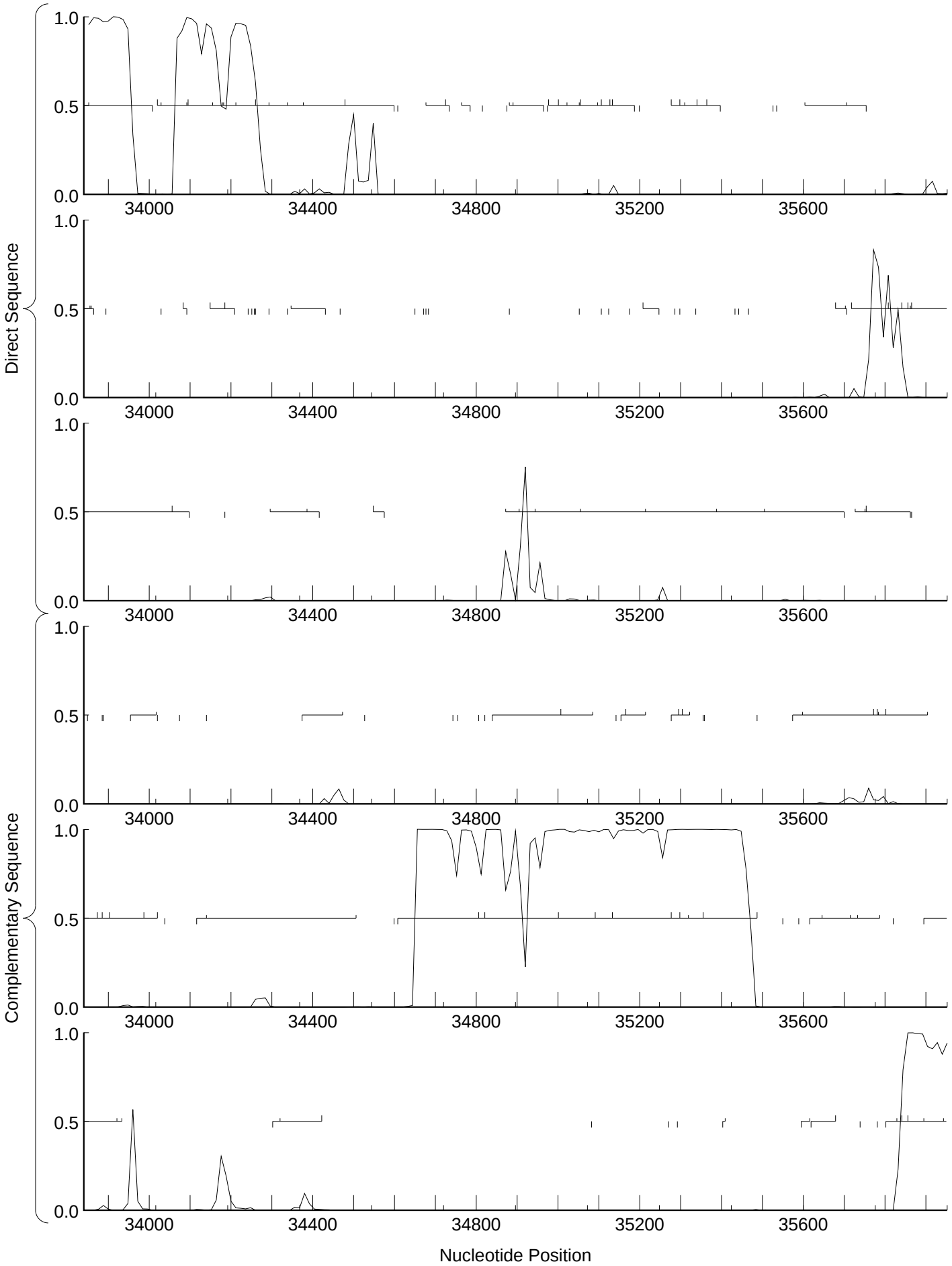
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 16/32



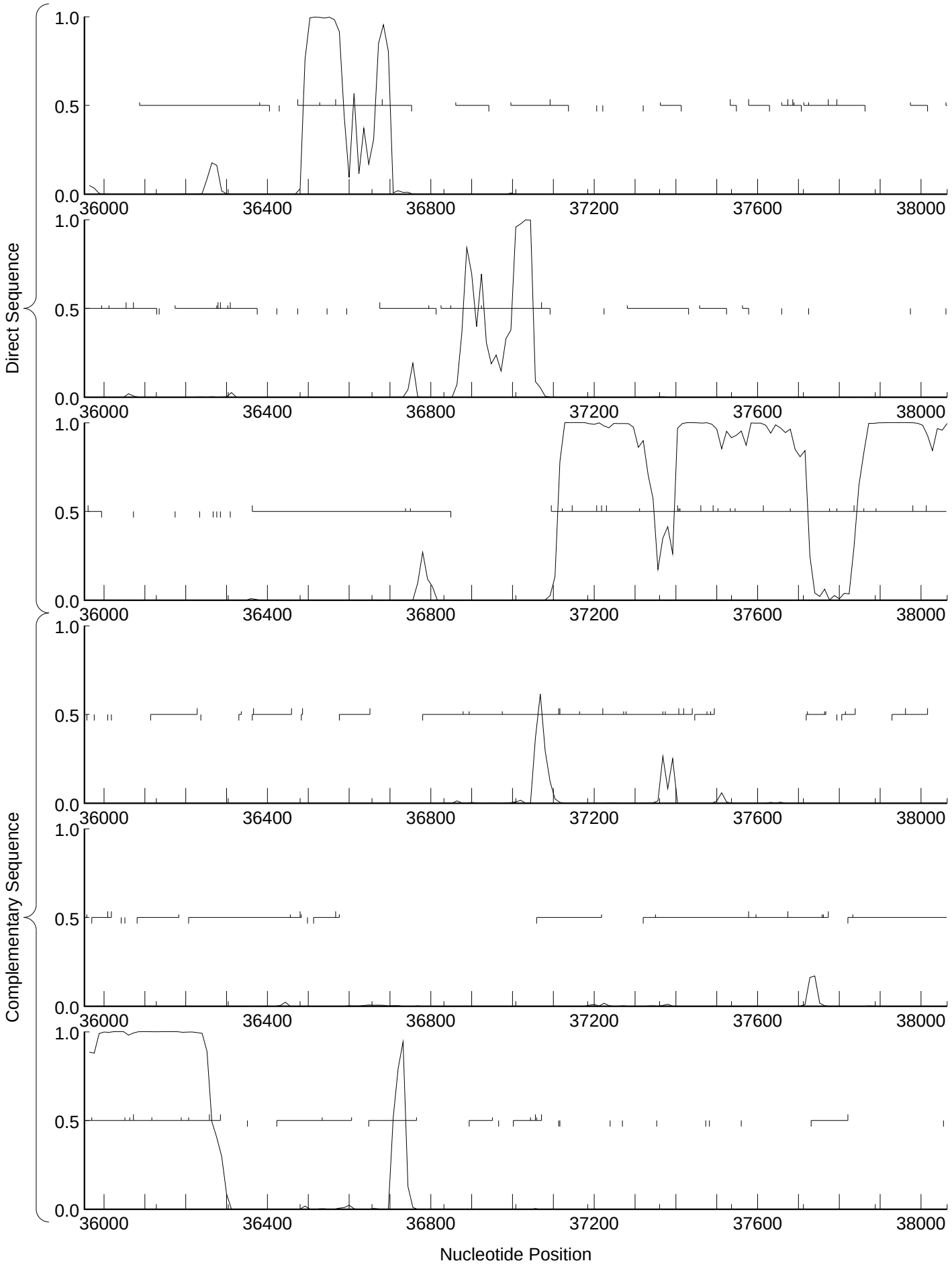
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 17/32



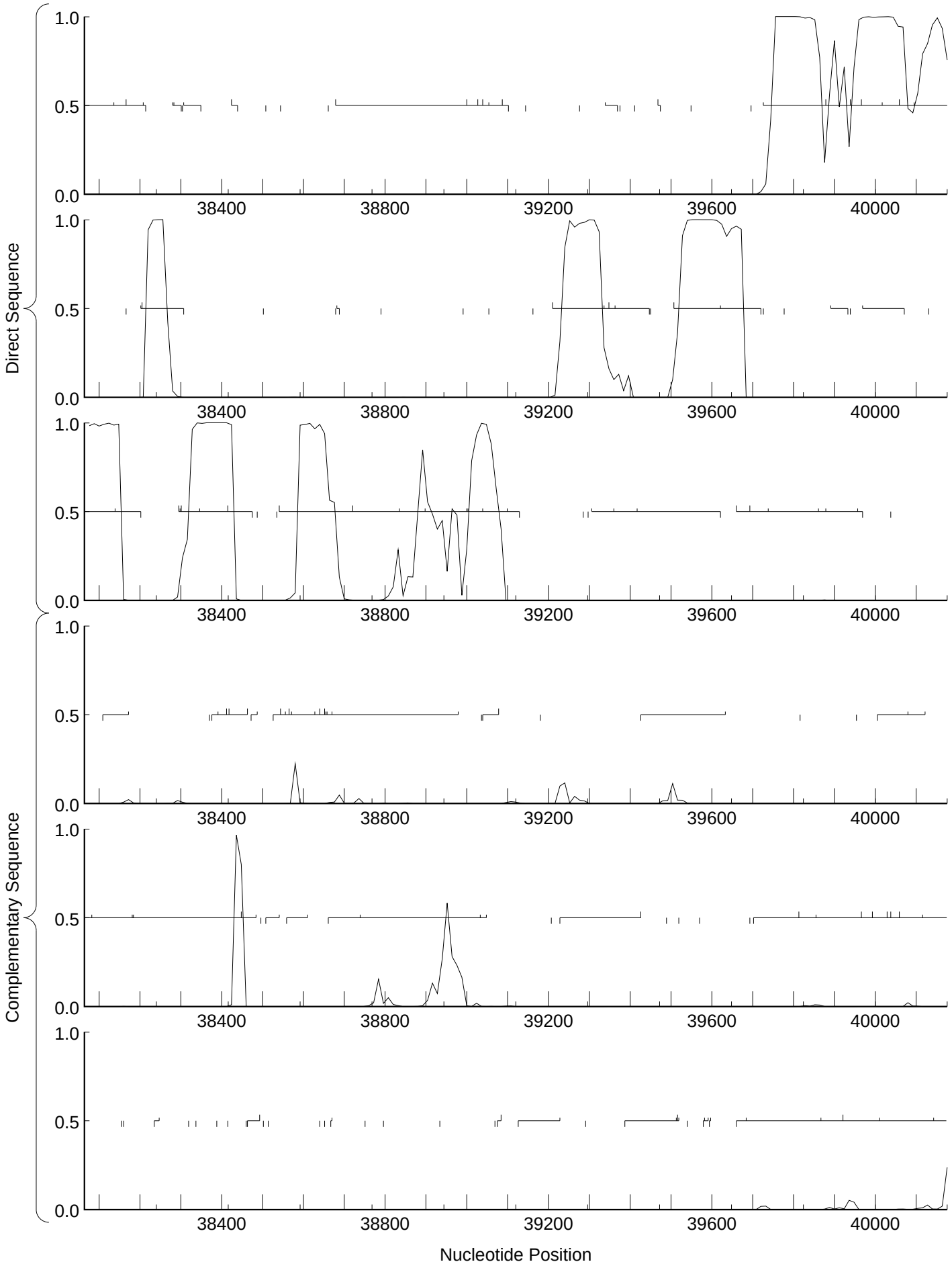
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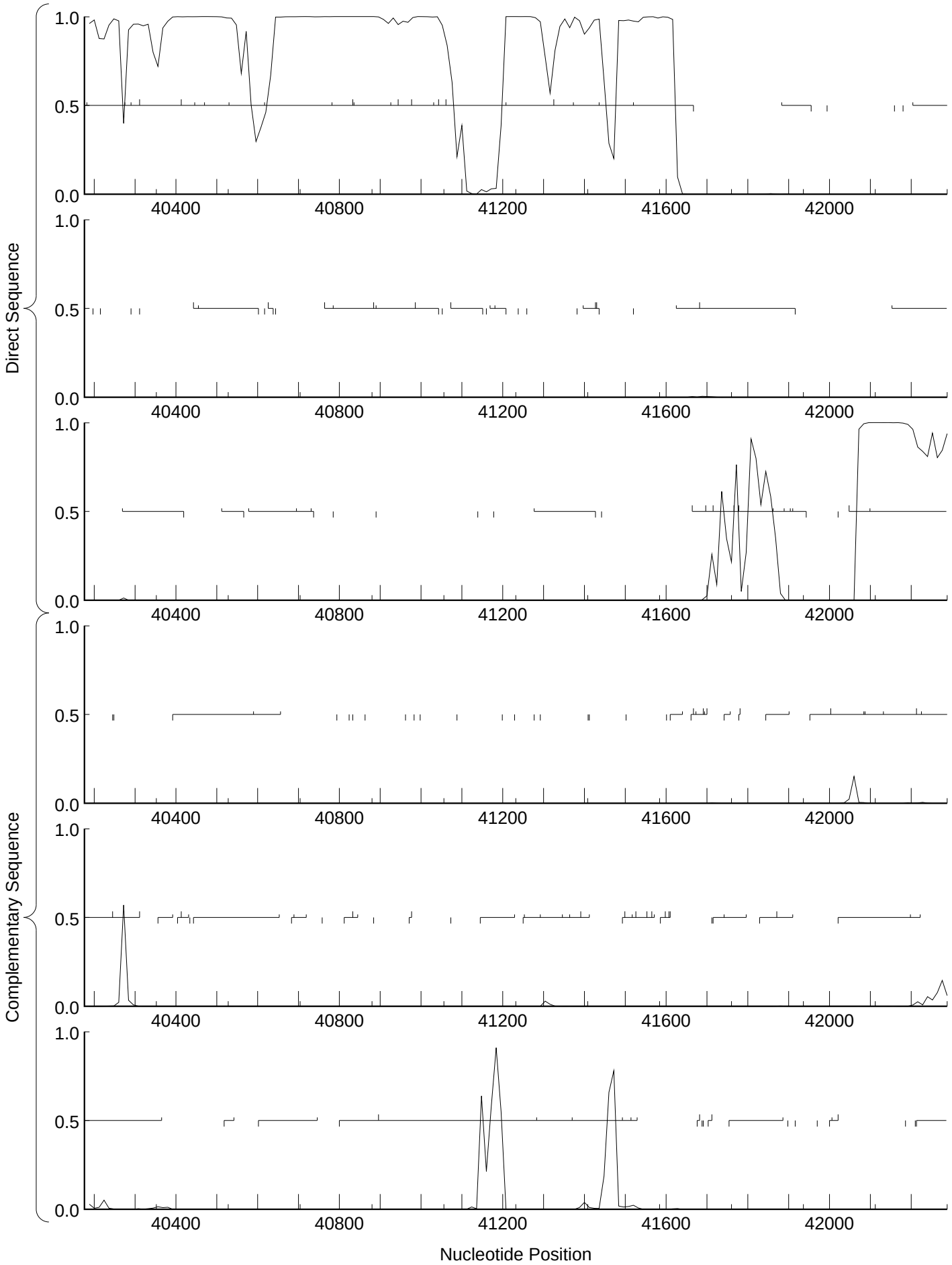
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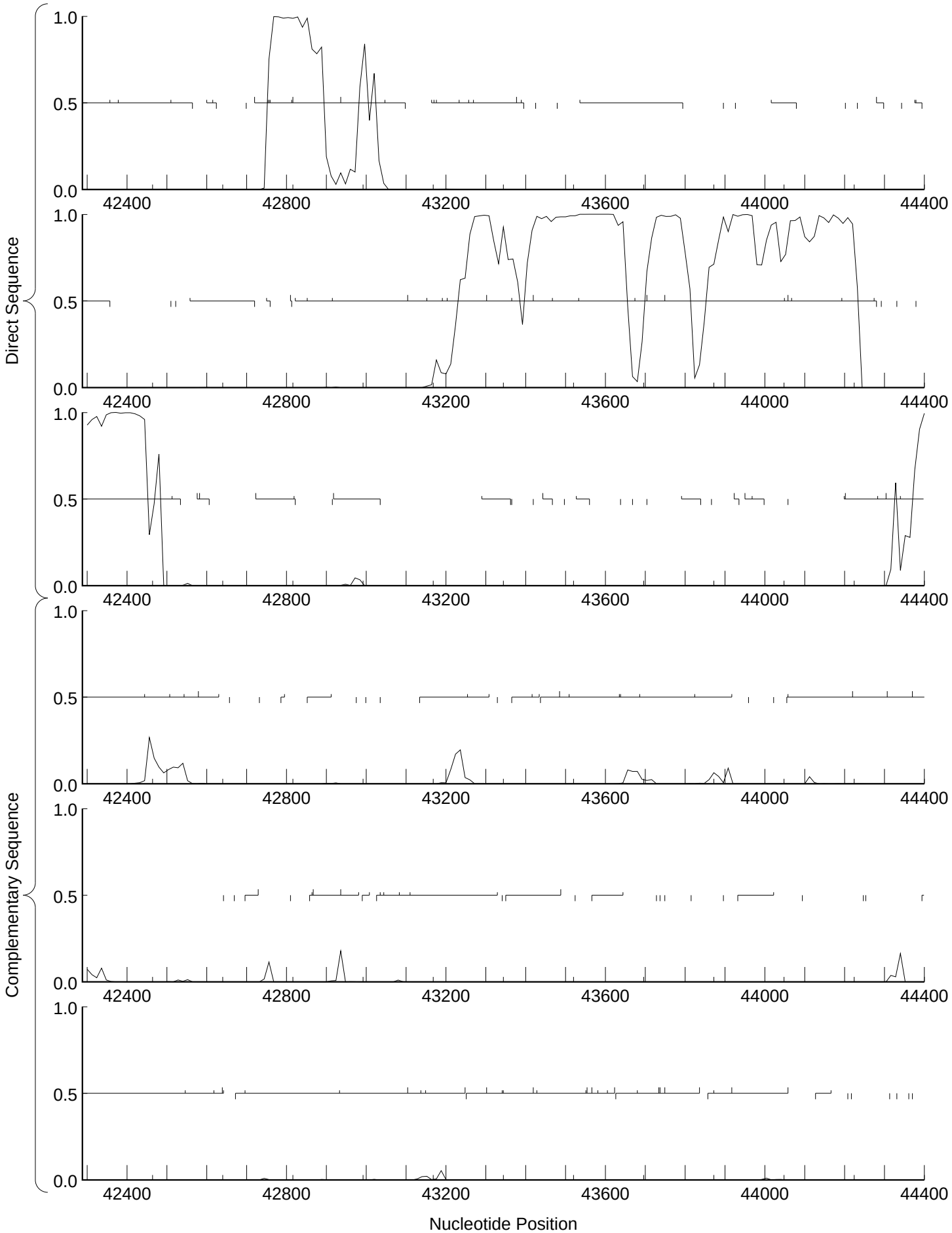
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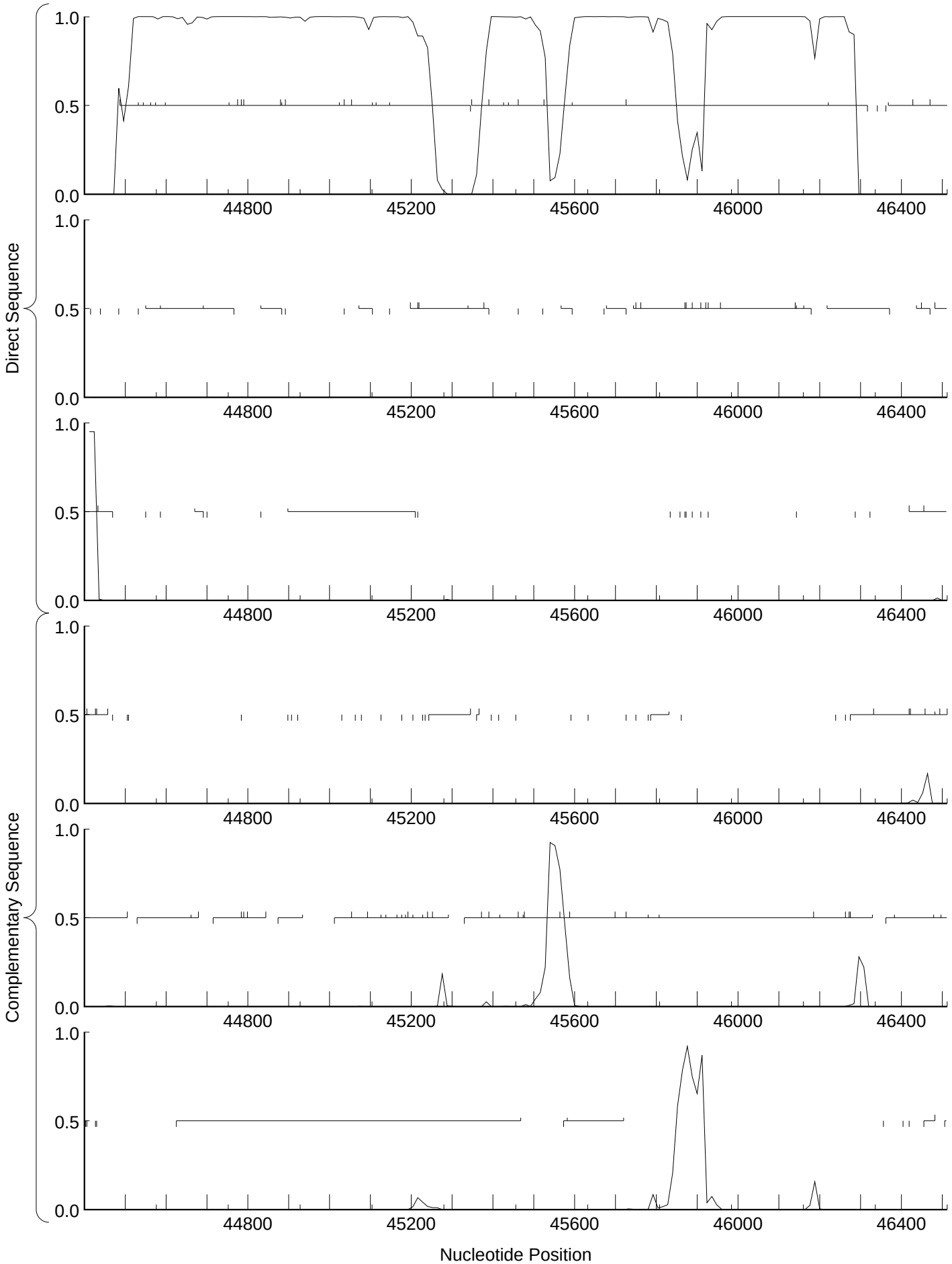
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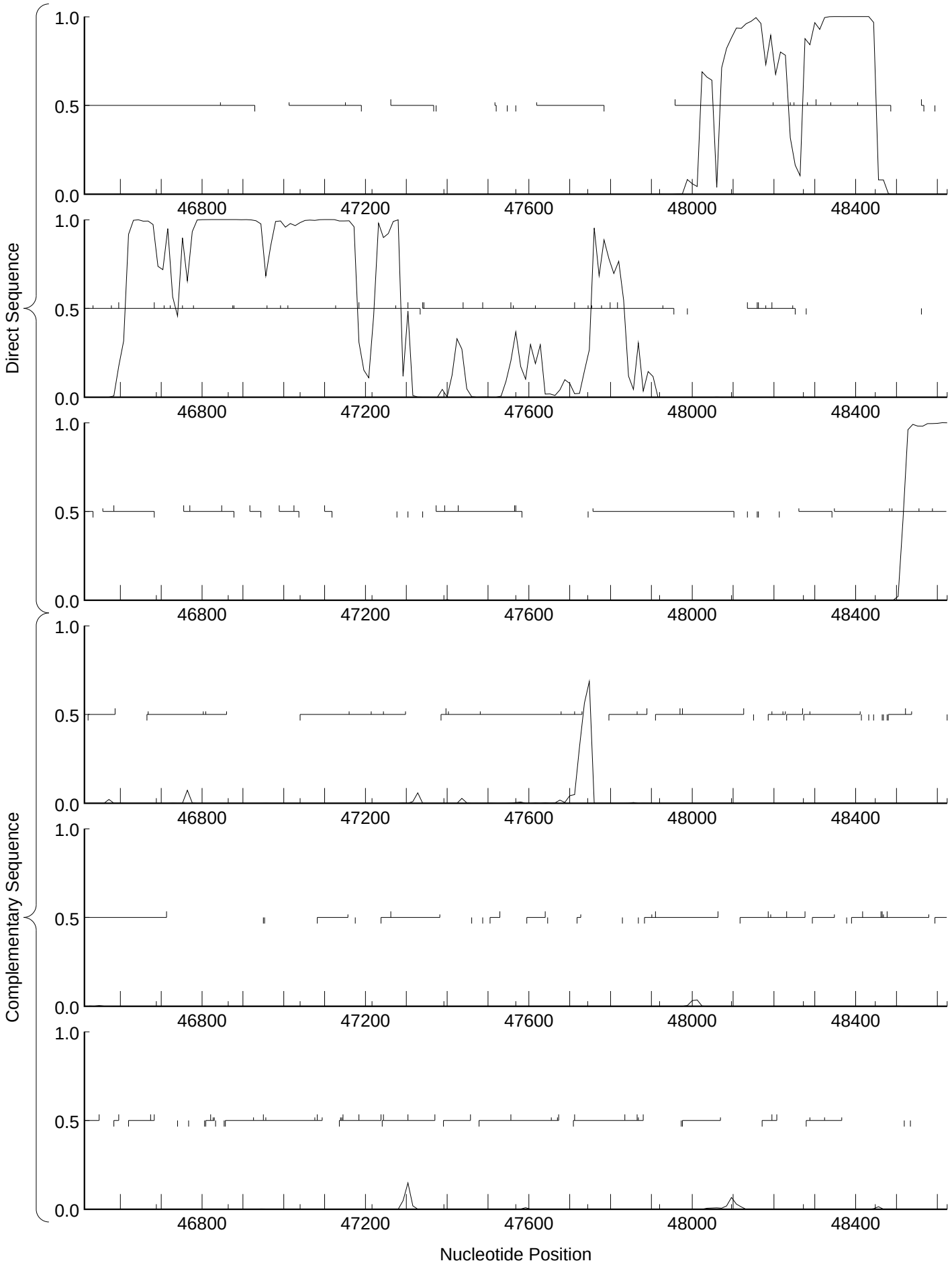
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 22/32



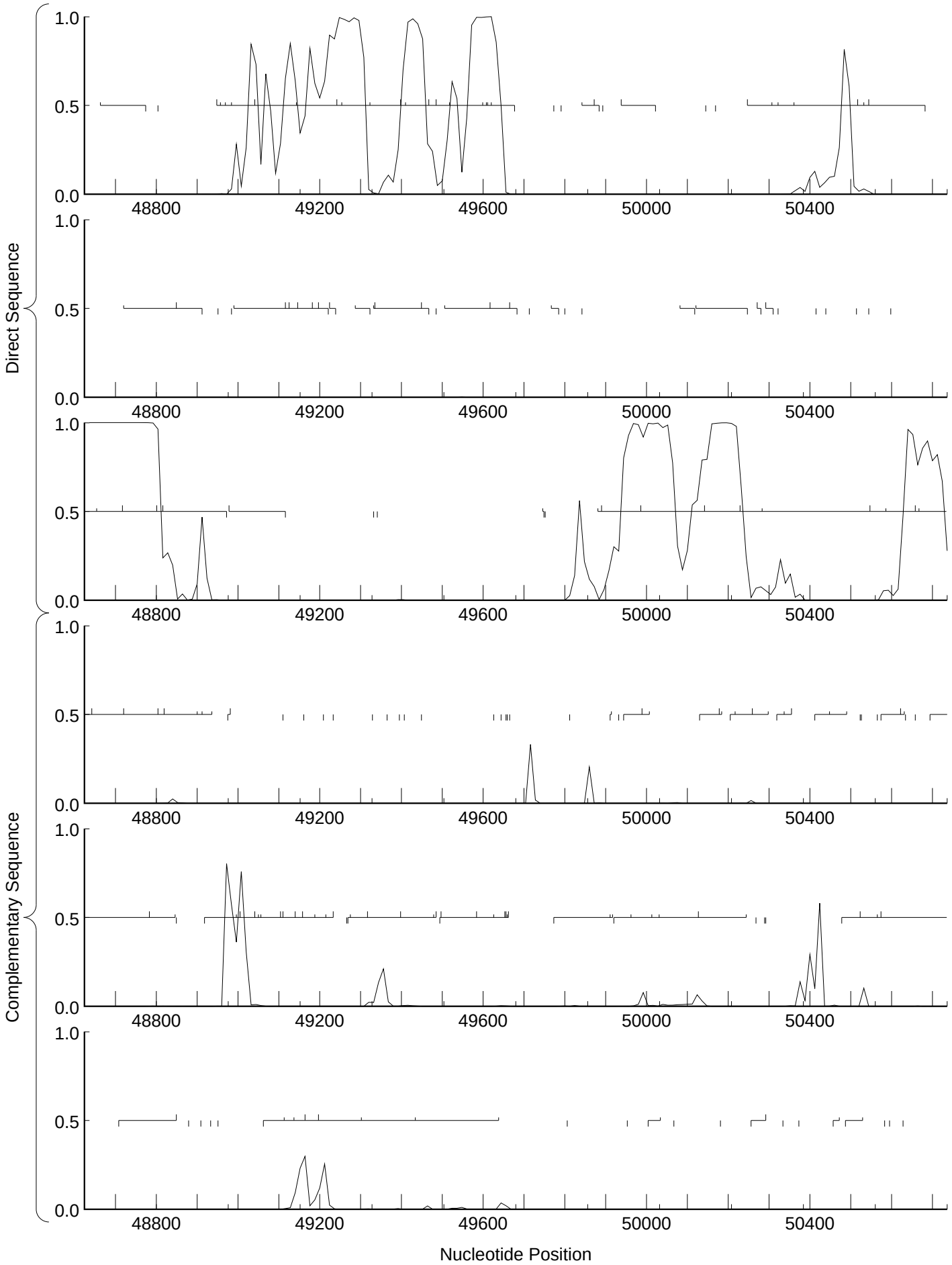
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 23/32



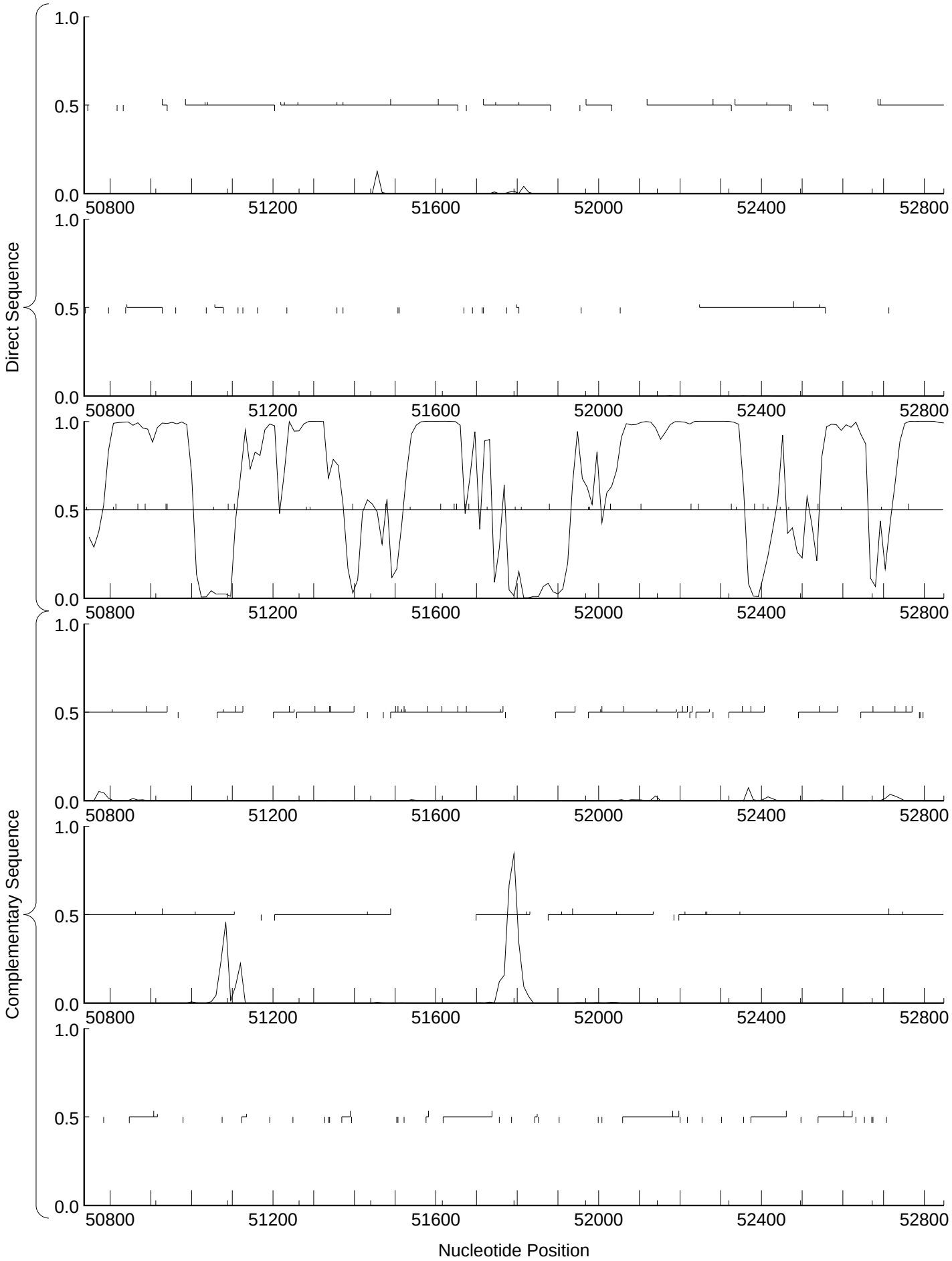
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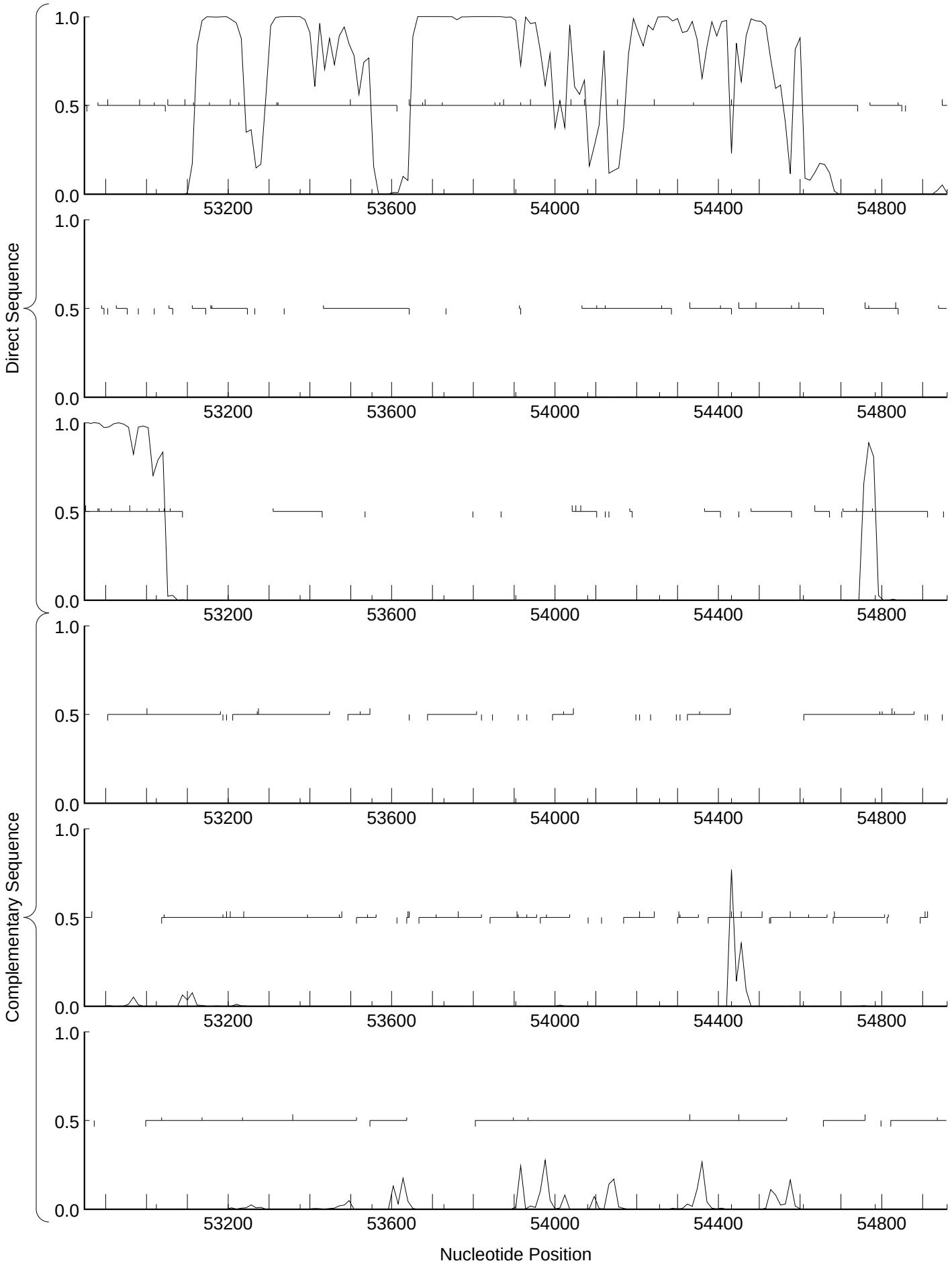
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 25/32



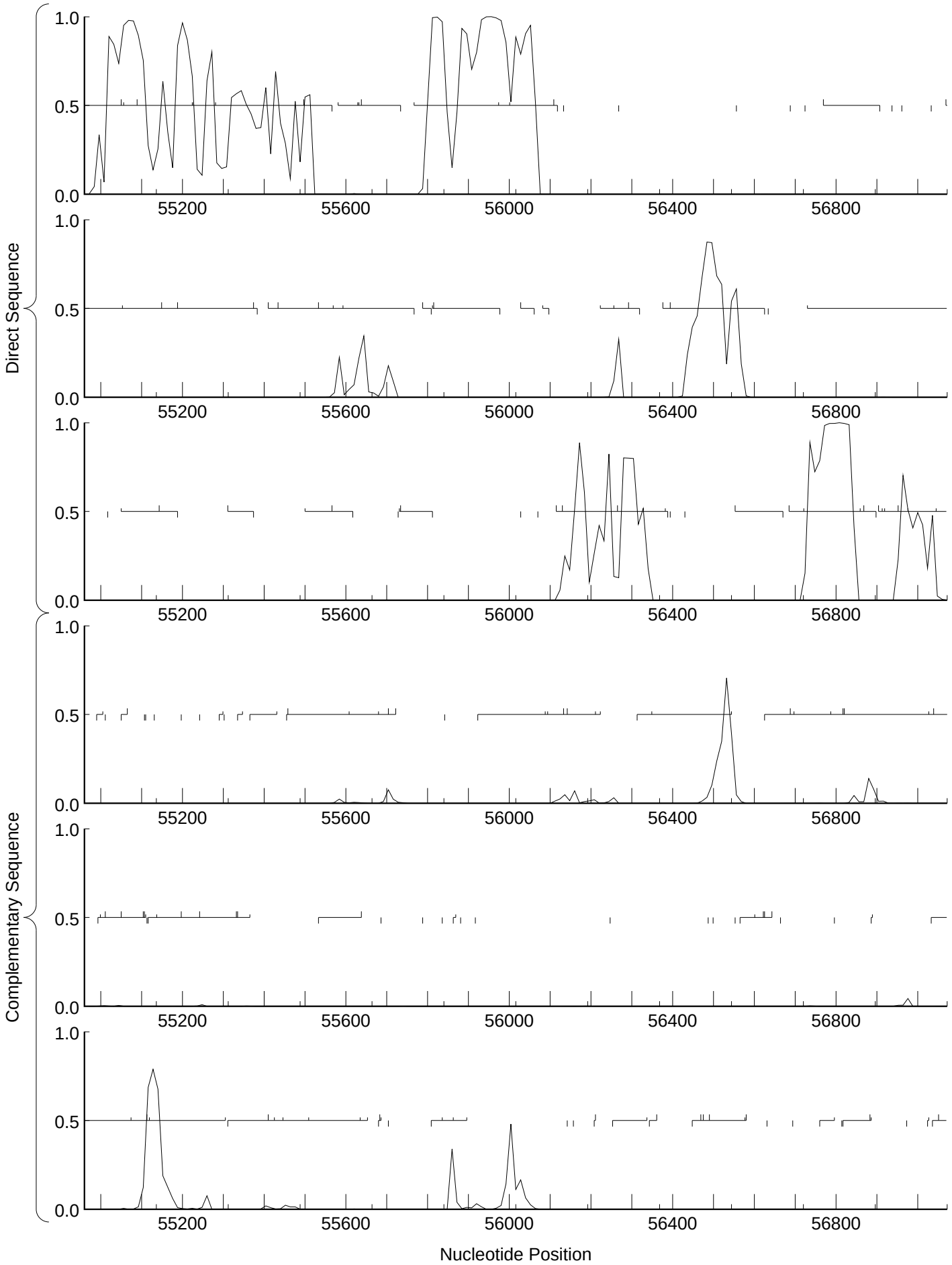
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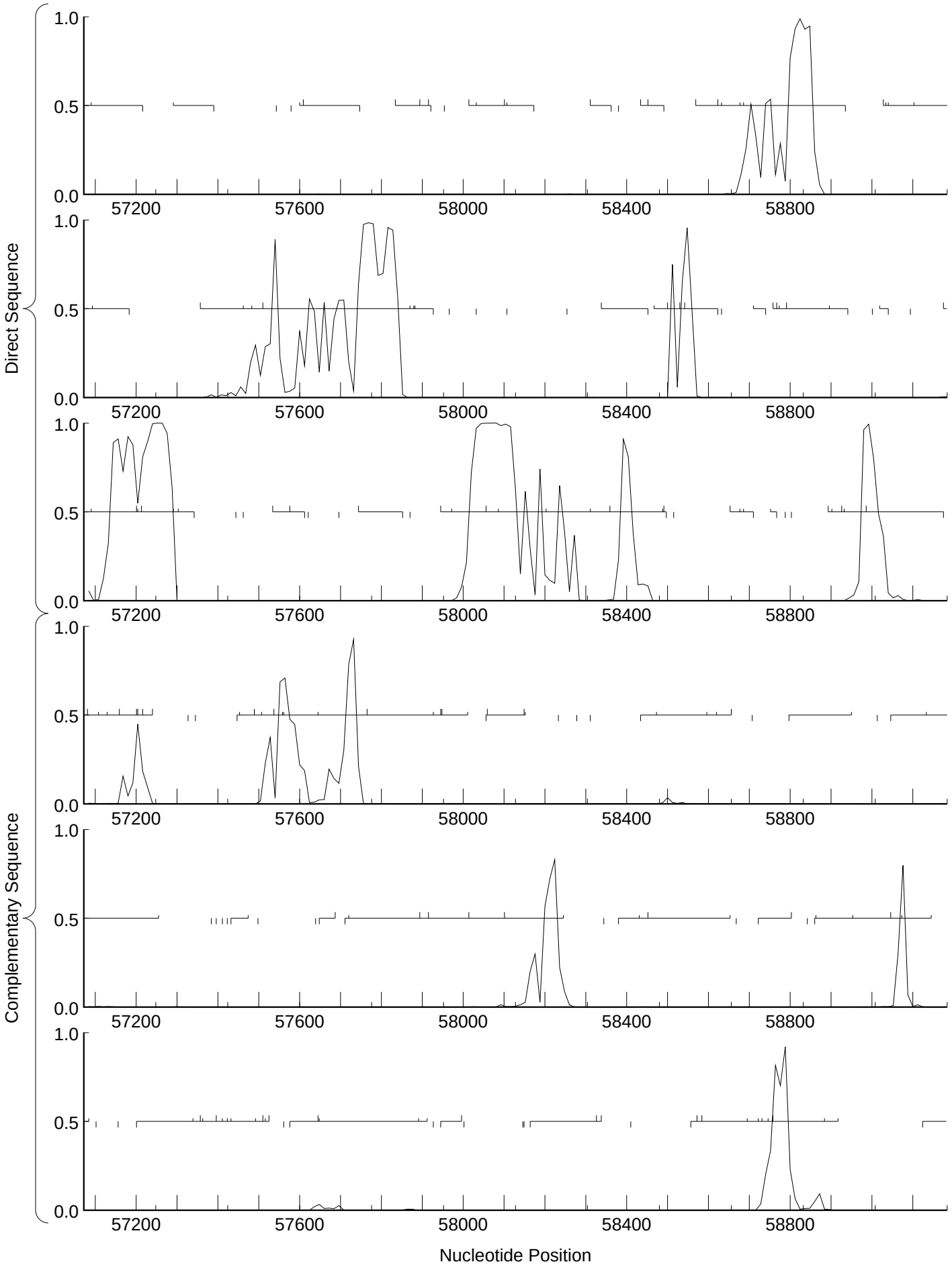
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 27/32



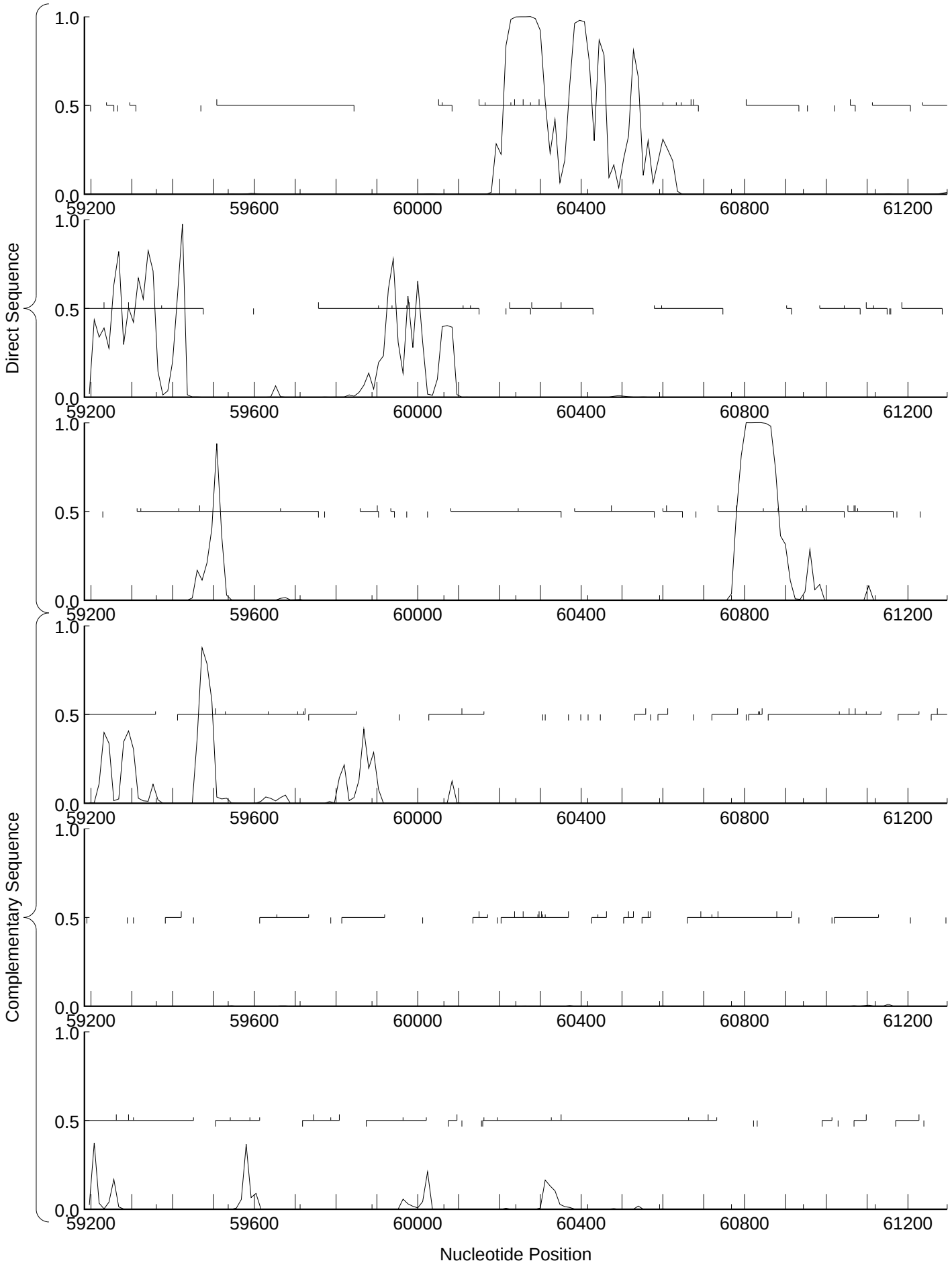
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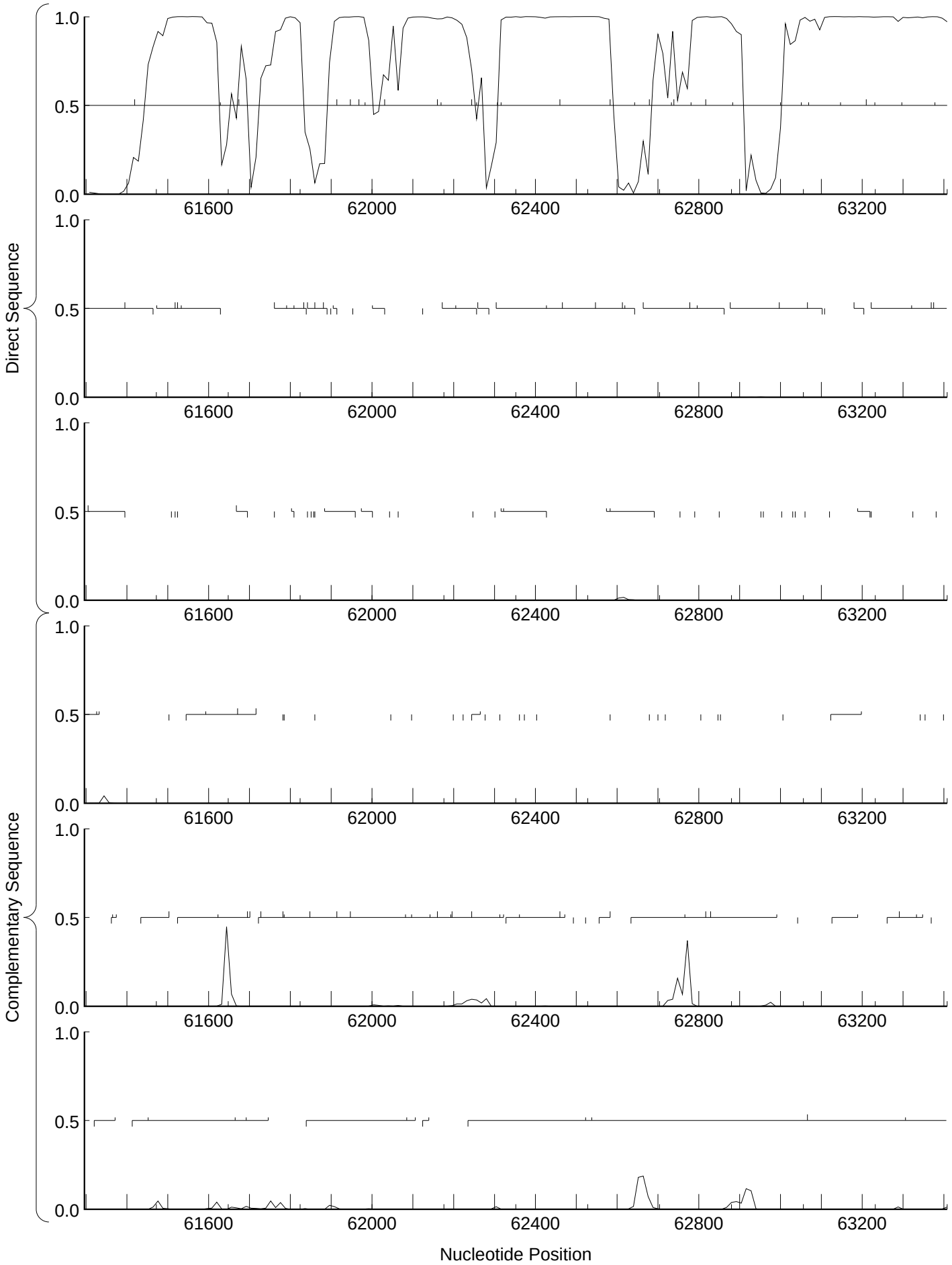
Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 29/32



Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 30/32



Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 31/32



Mycobacterium phage Thoth complete sequence, 64606 bp, circularly permuted, Cluster D1, Order 4, Window 96, Step 12, 32/32

