# GeneMark

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

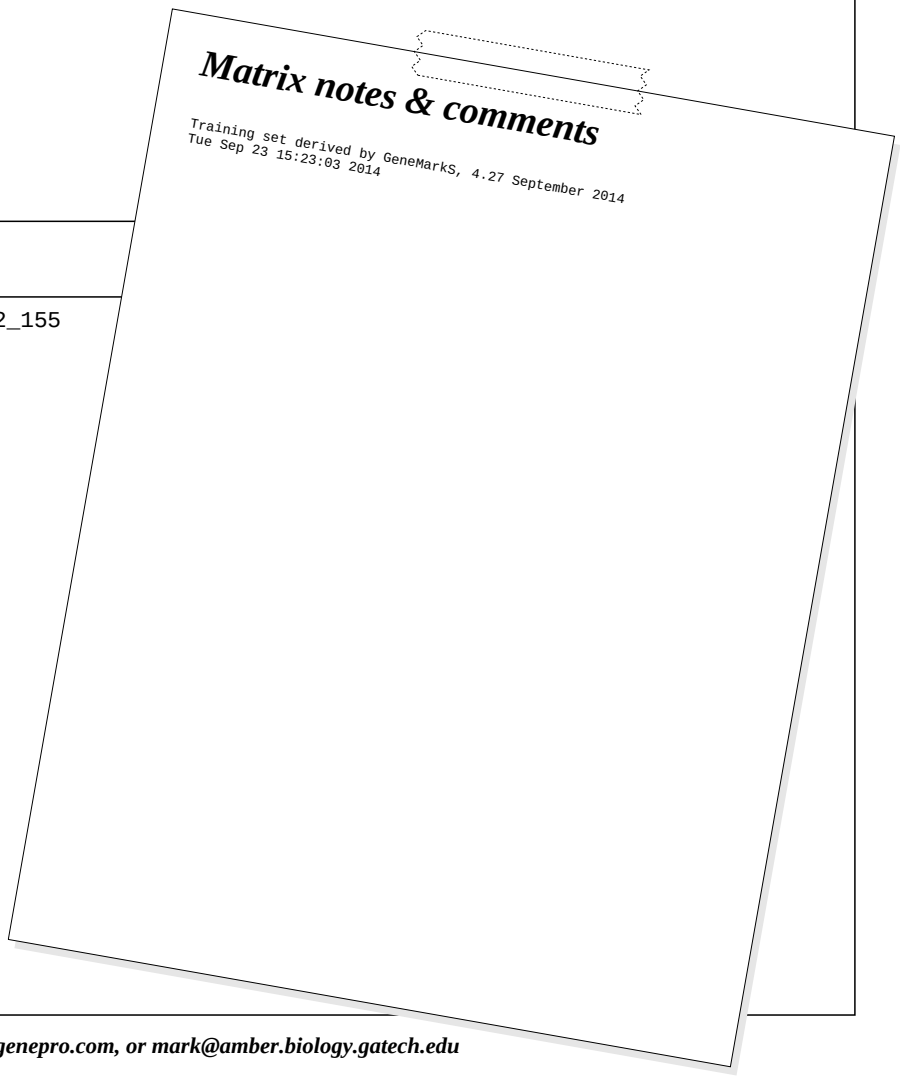
## PROGRAM INFORMATION

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Sequence         :  Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5
Analysis Date    :  3/28/24 at 17:10:35
Pages            :  35
Sequence Length  :  70270 bp
GC Content       :   67.64%

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

PS-Version       :  1.2

GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
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## MATRIX INFORMATION

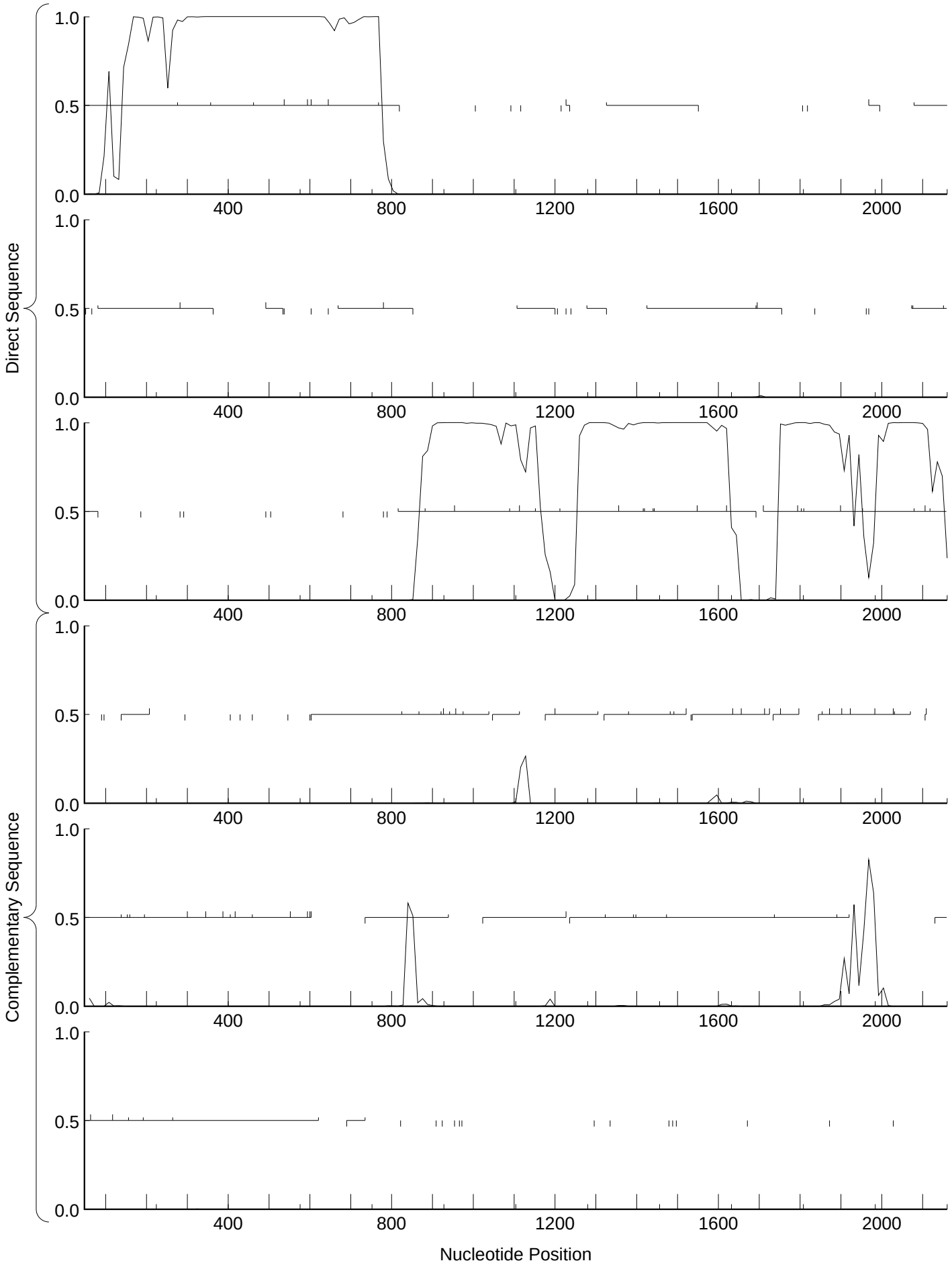
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Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

Order      :  4
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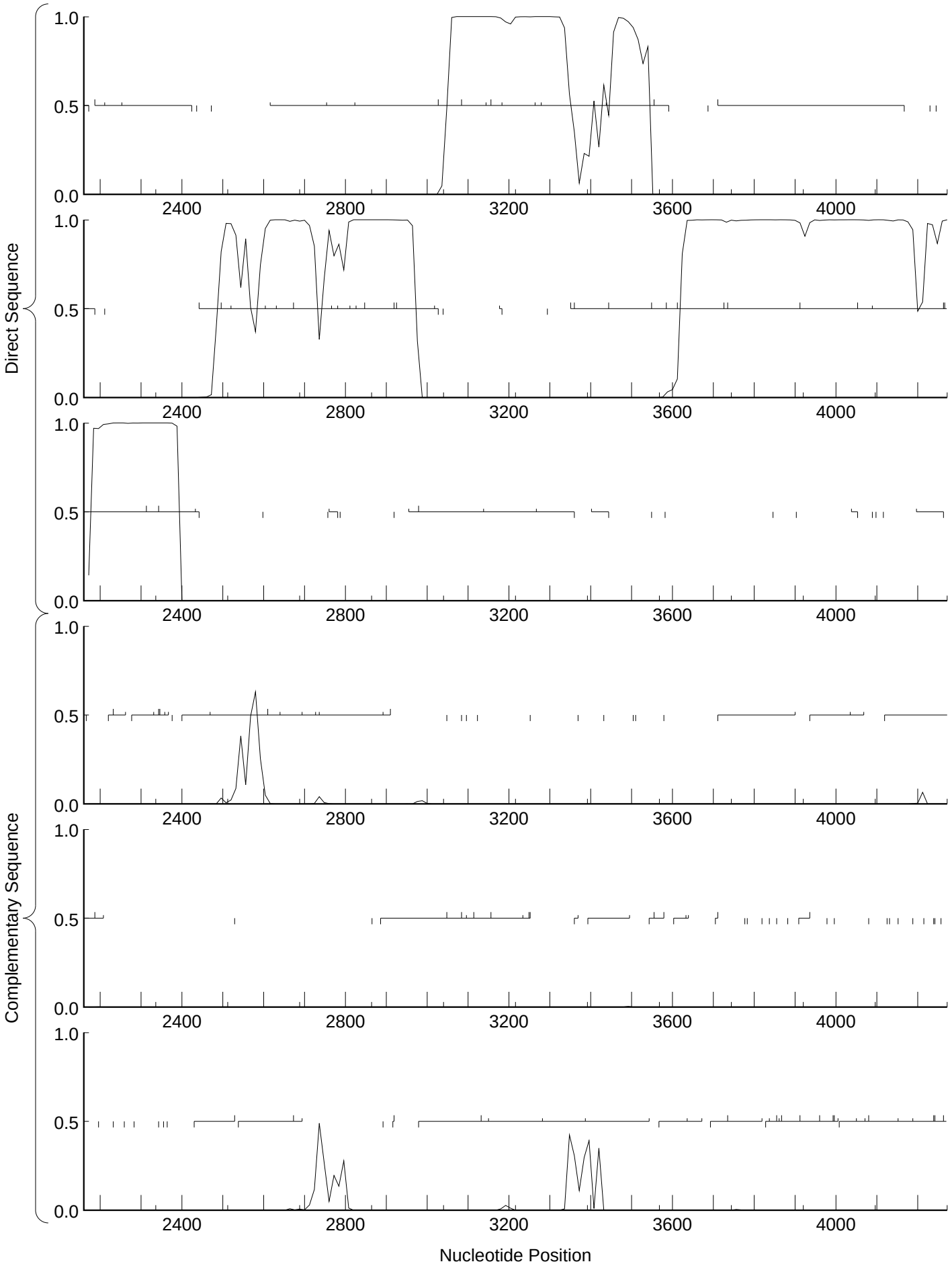

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

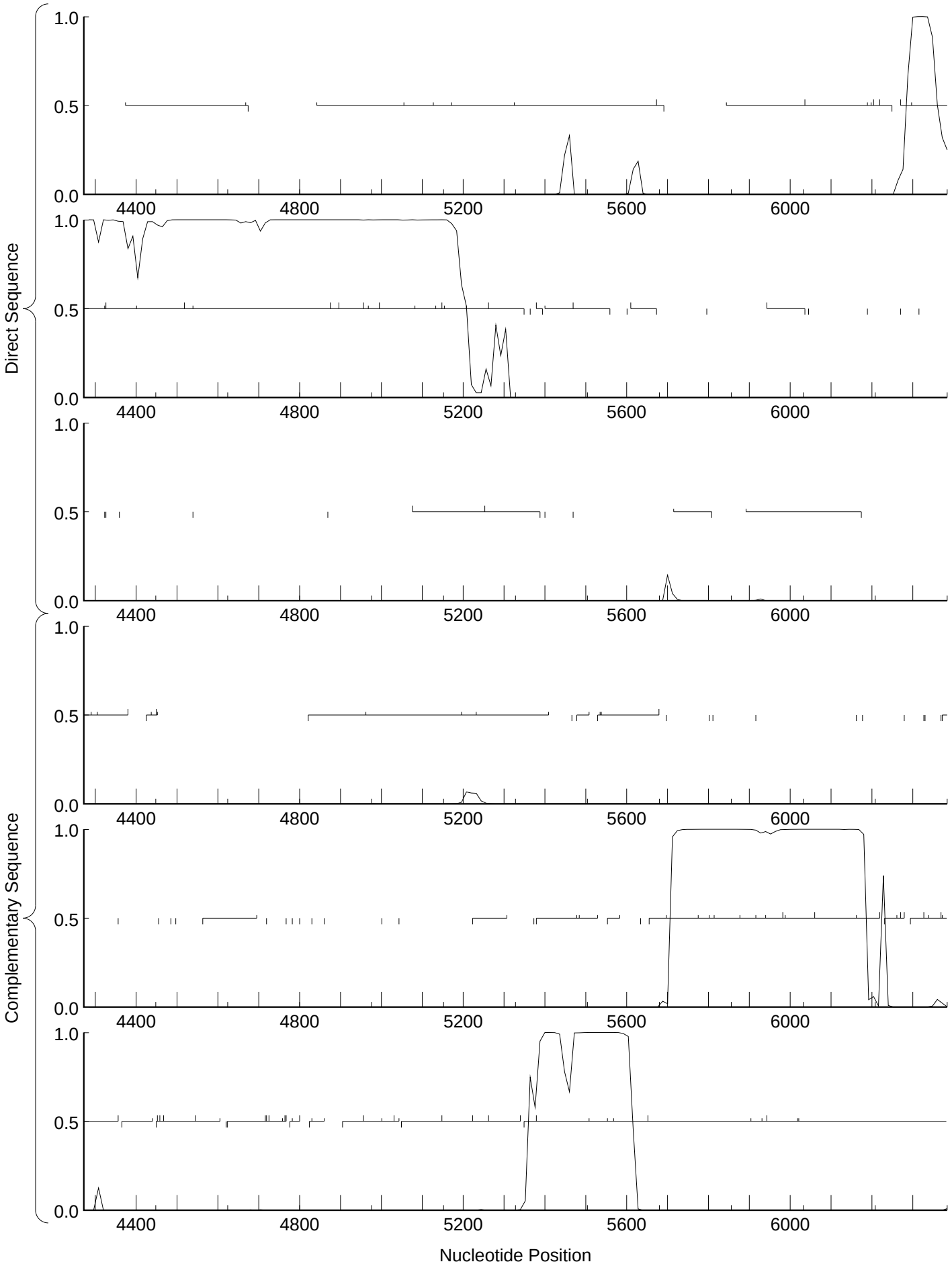
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 2/35



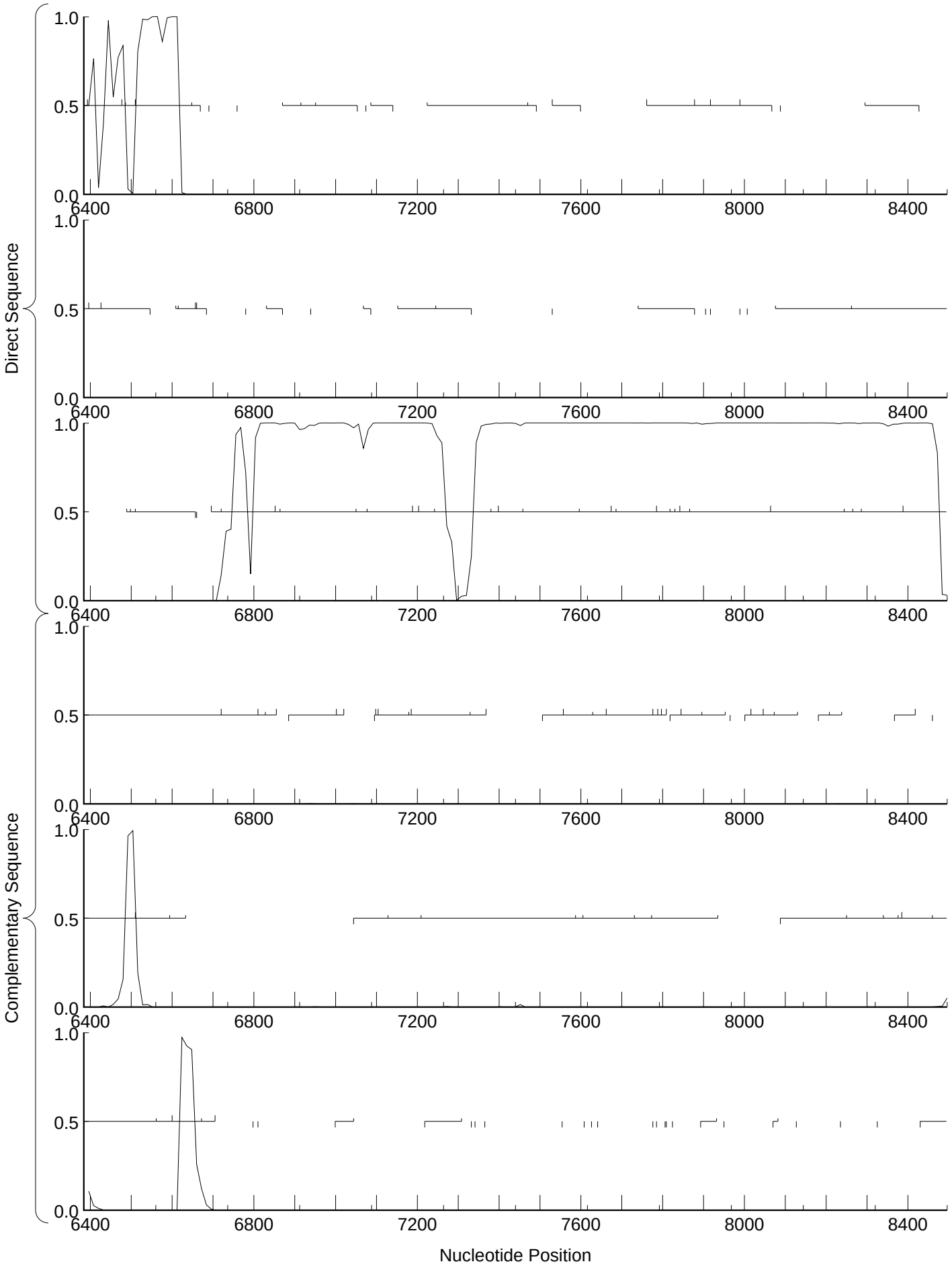
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 3/35



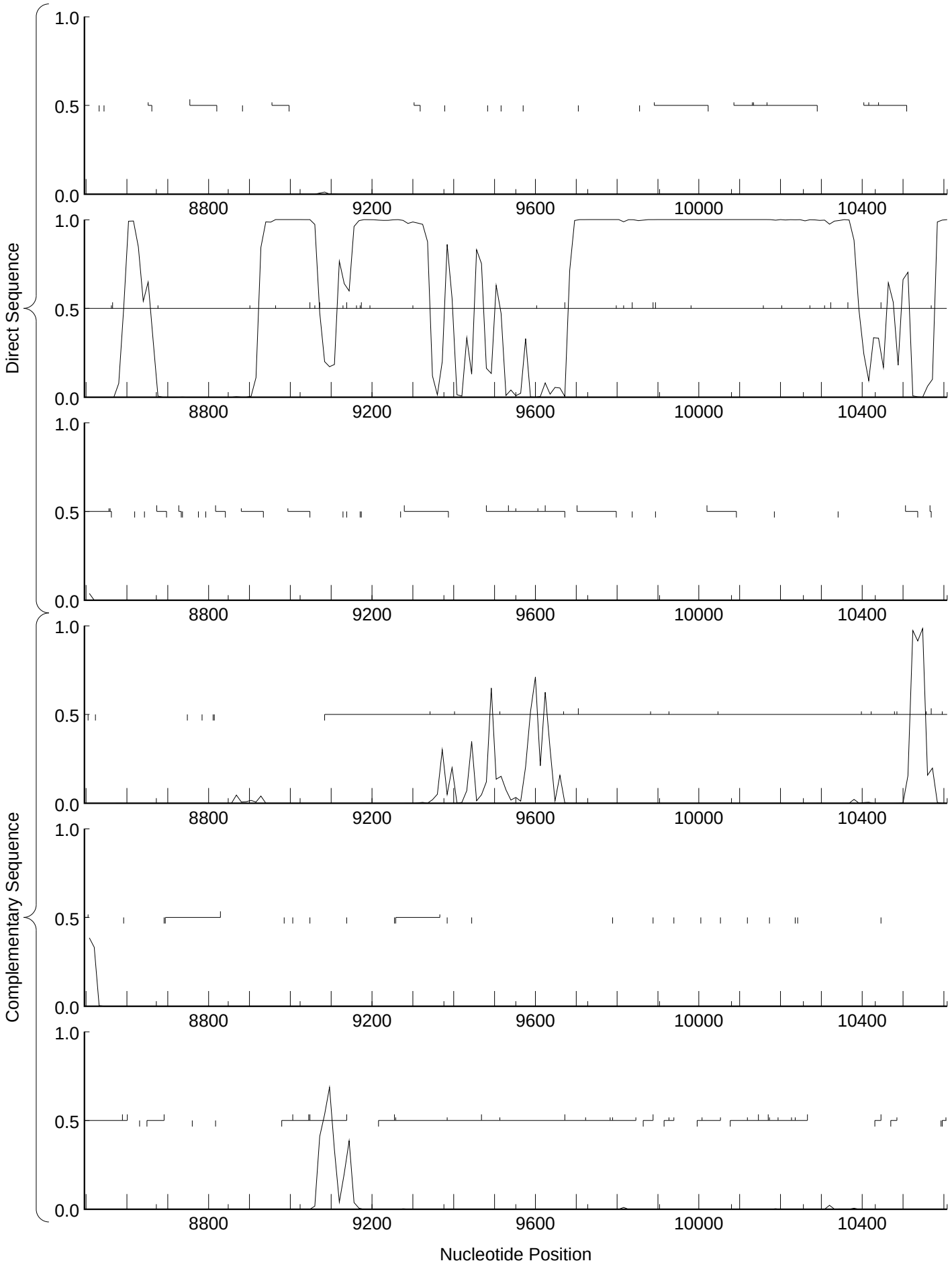
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 4/35



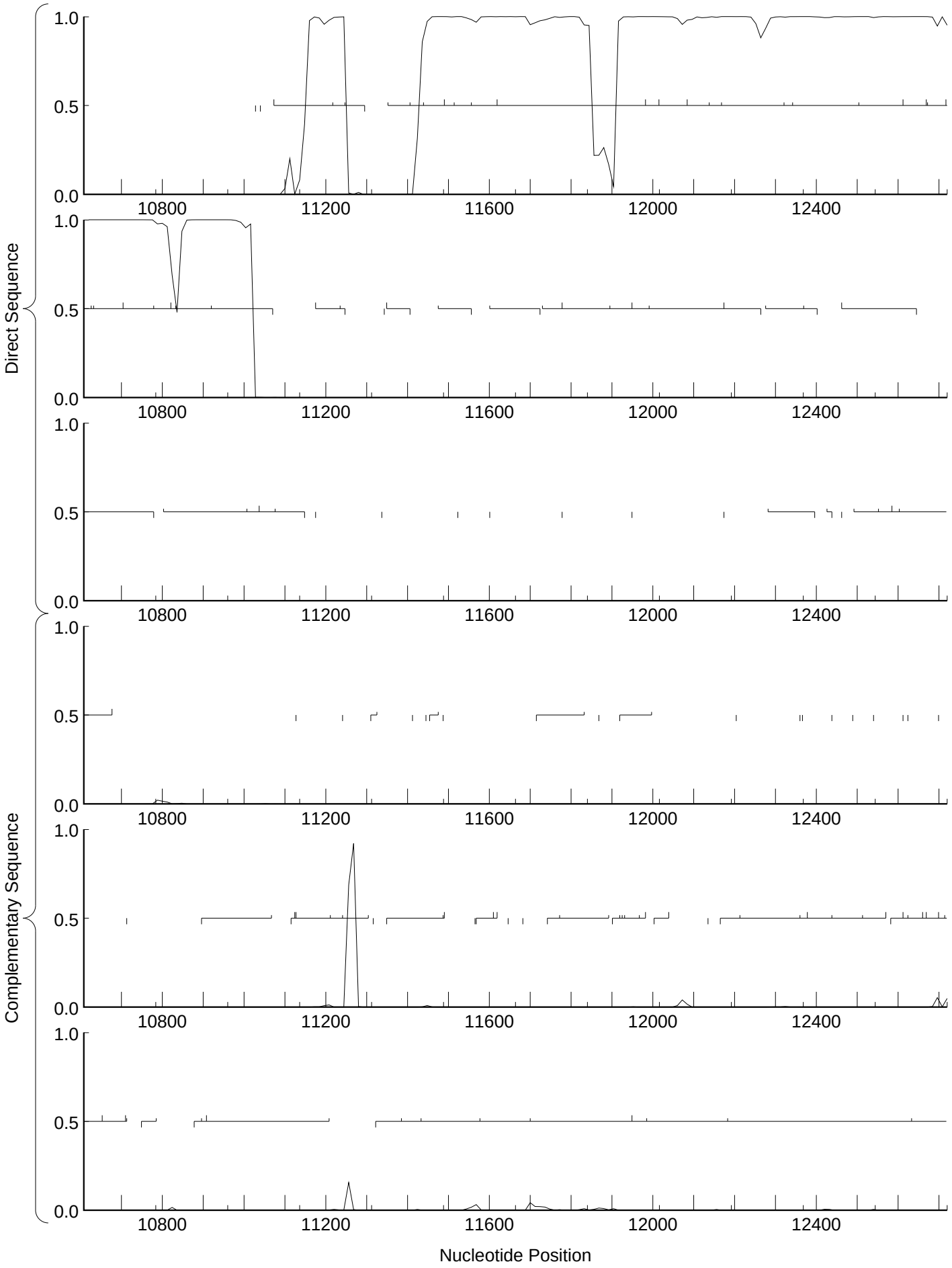
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 5/35



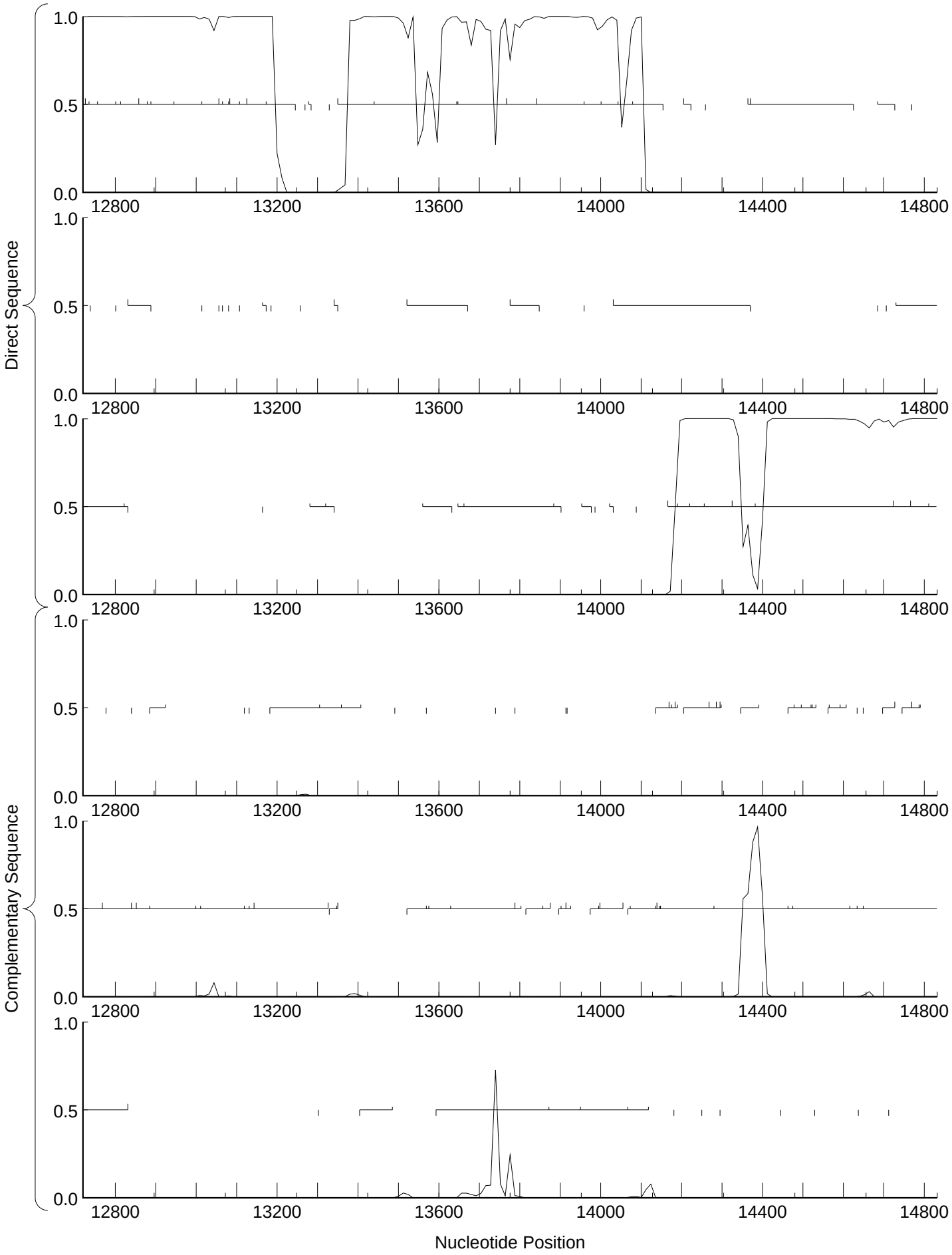
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 6/35



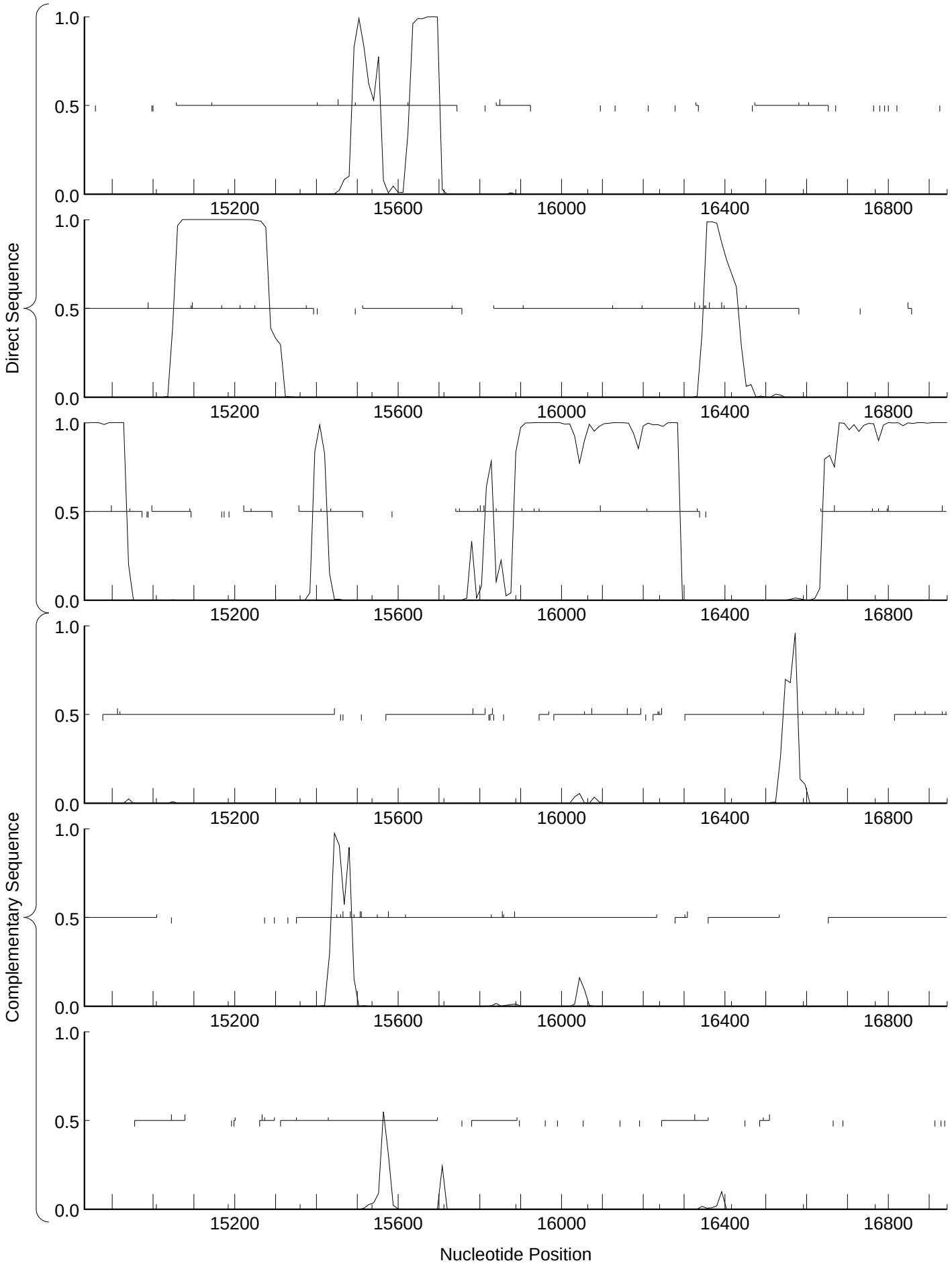
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 7/35



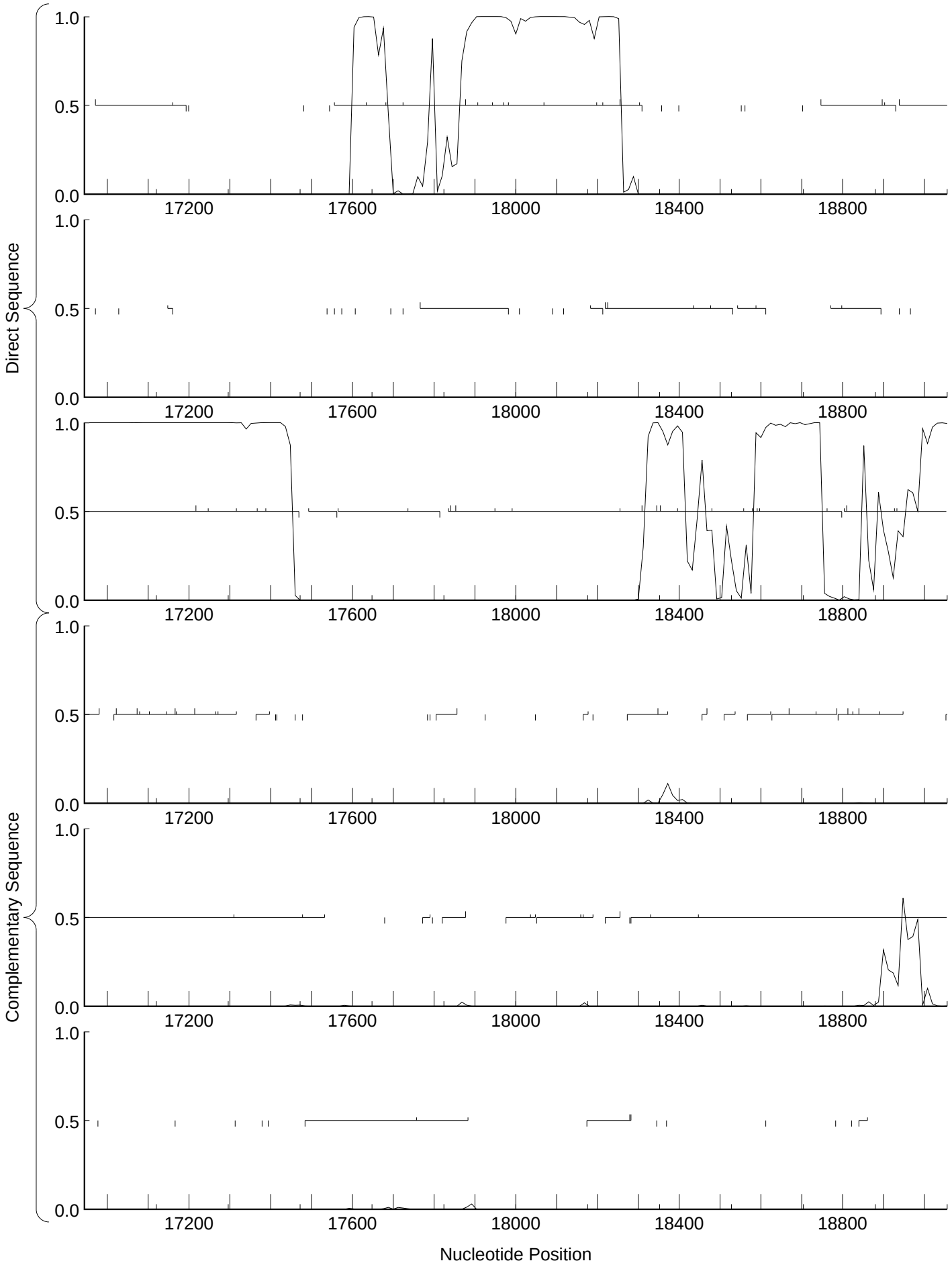
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 8/35



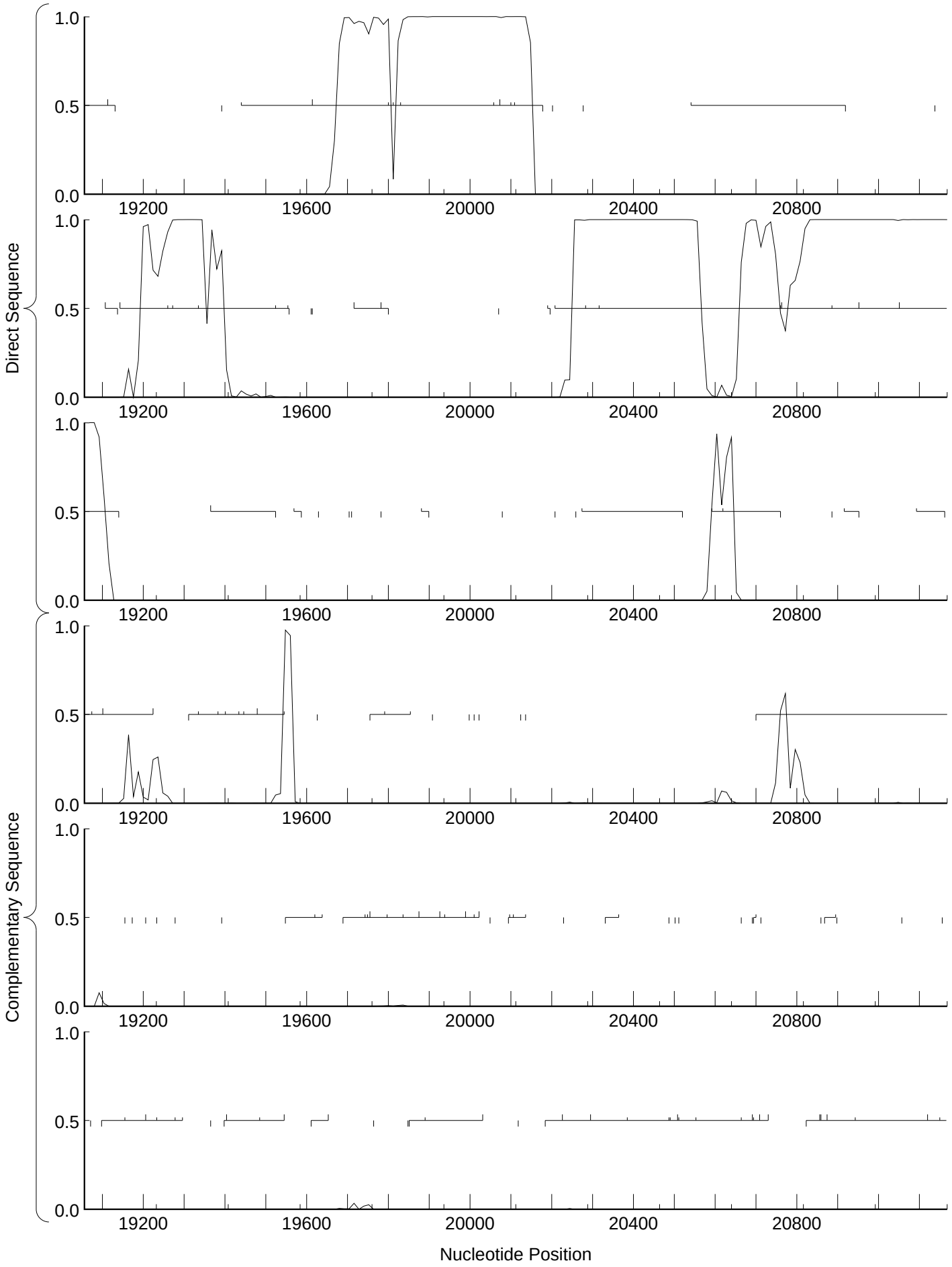
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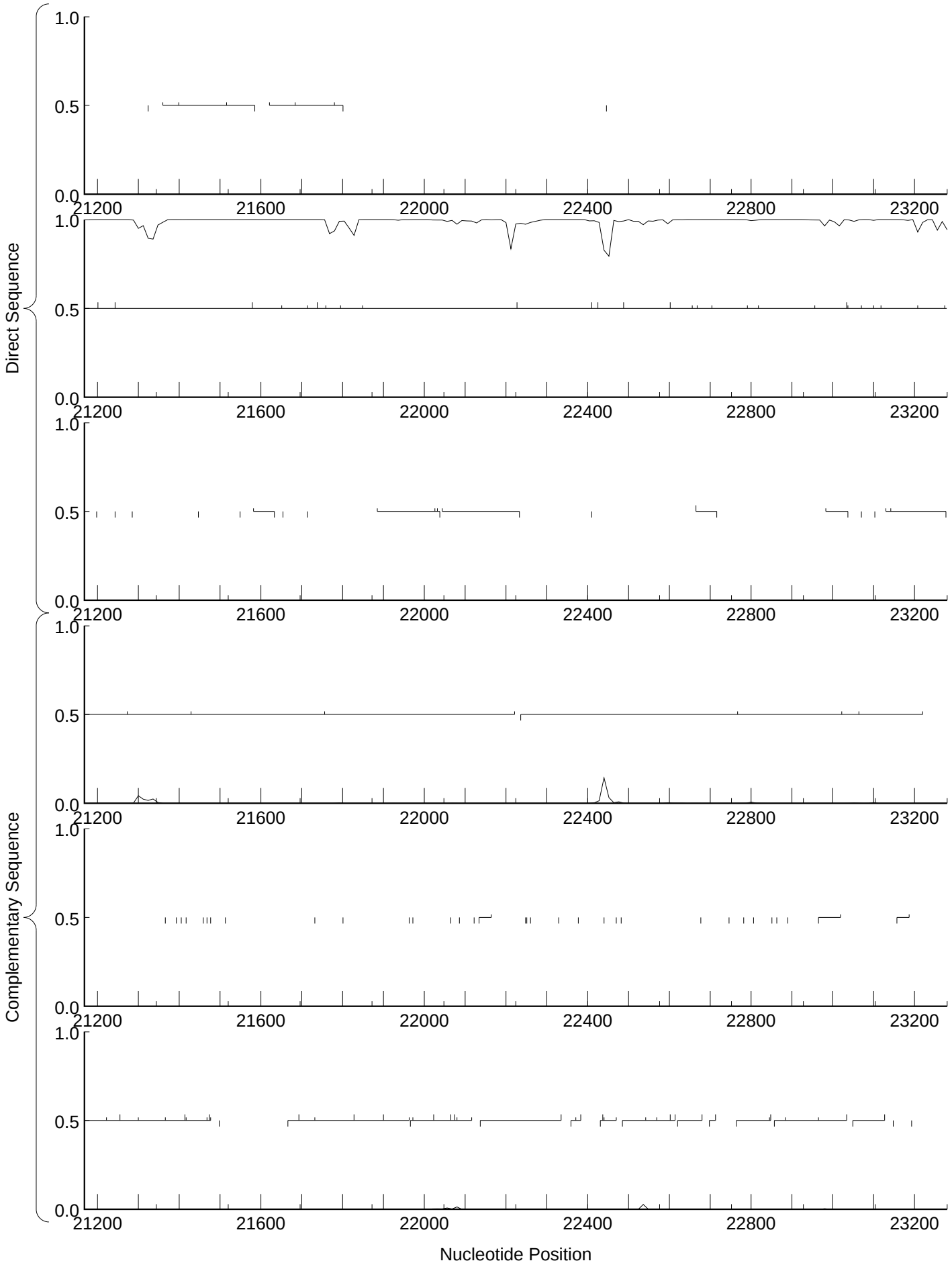
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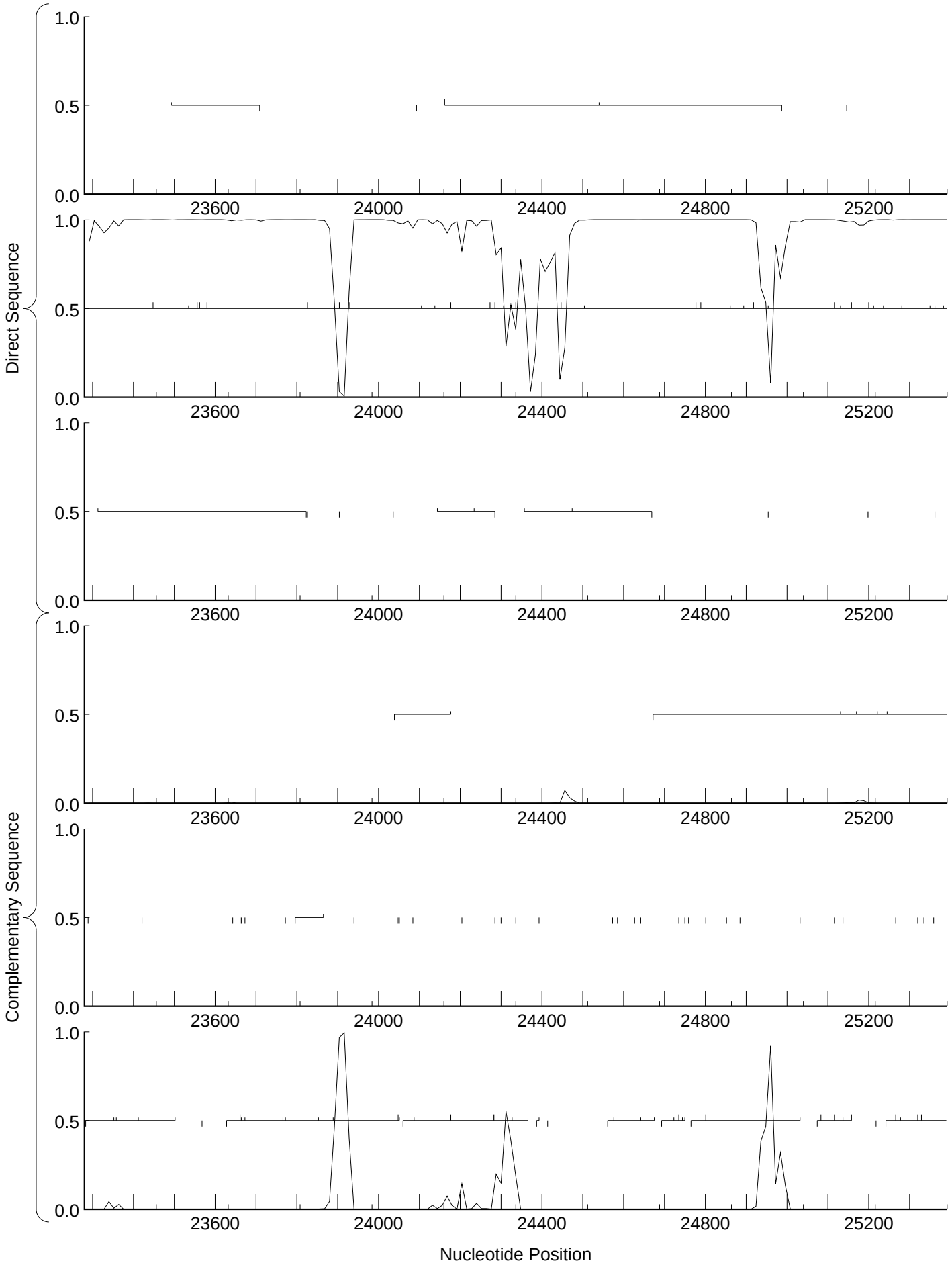
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 11/35



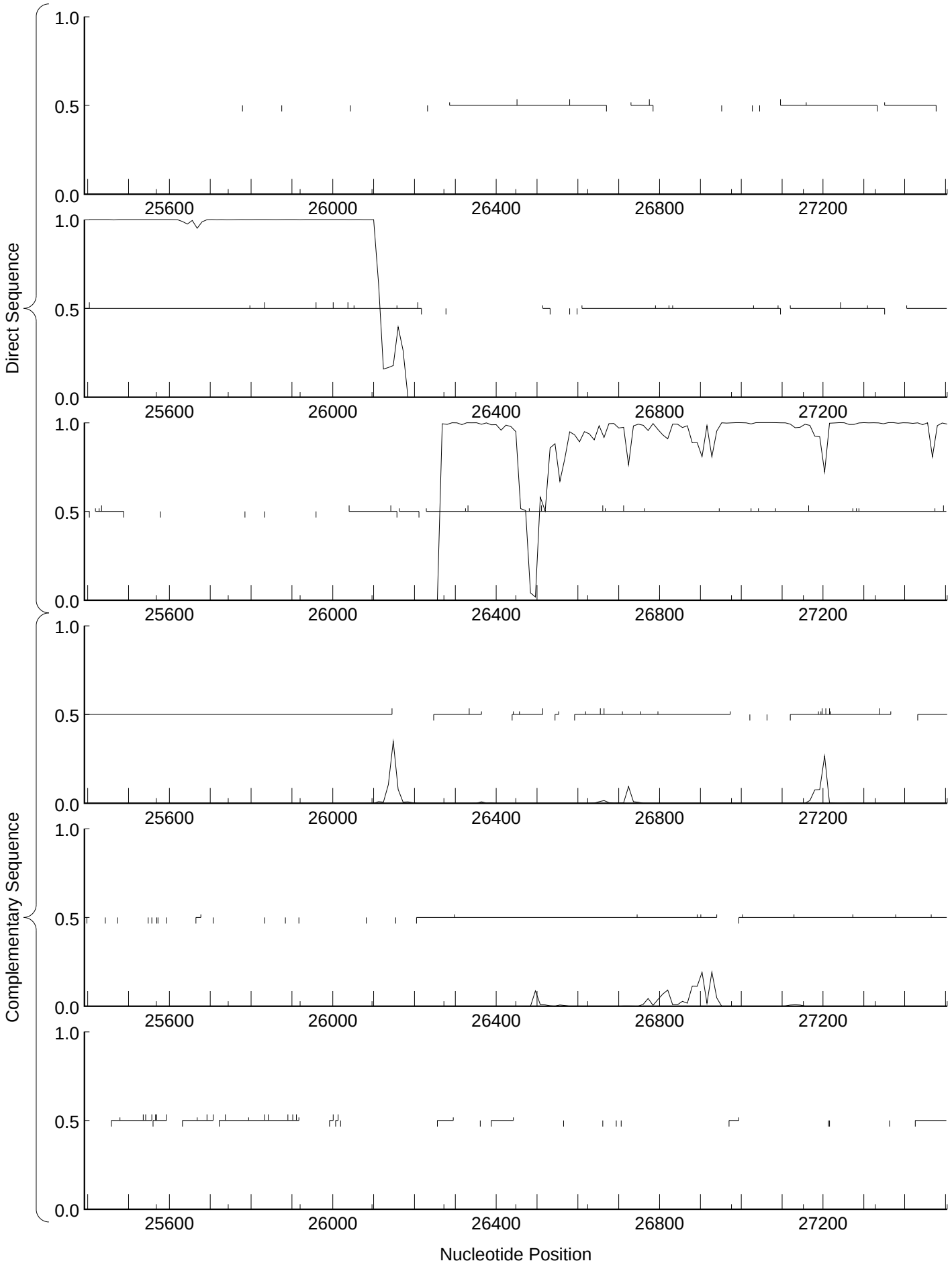
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 12/35



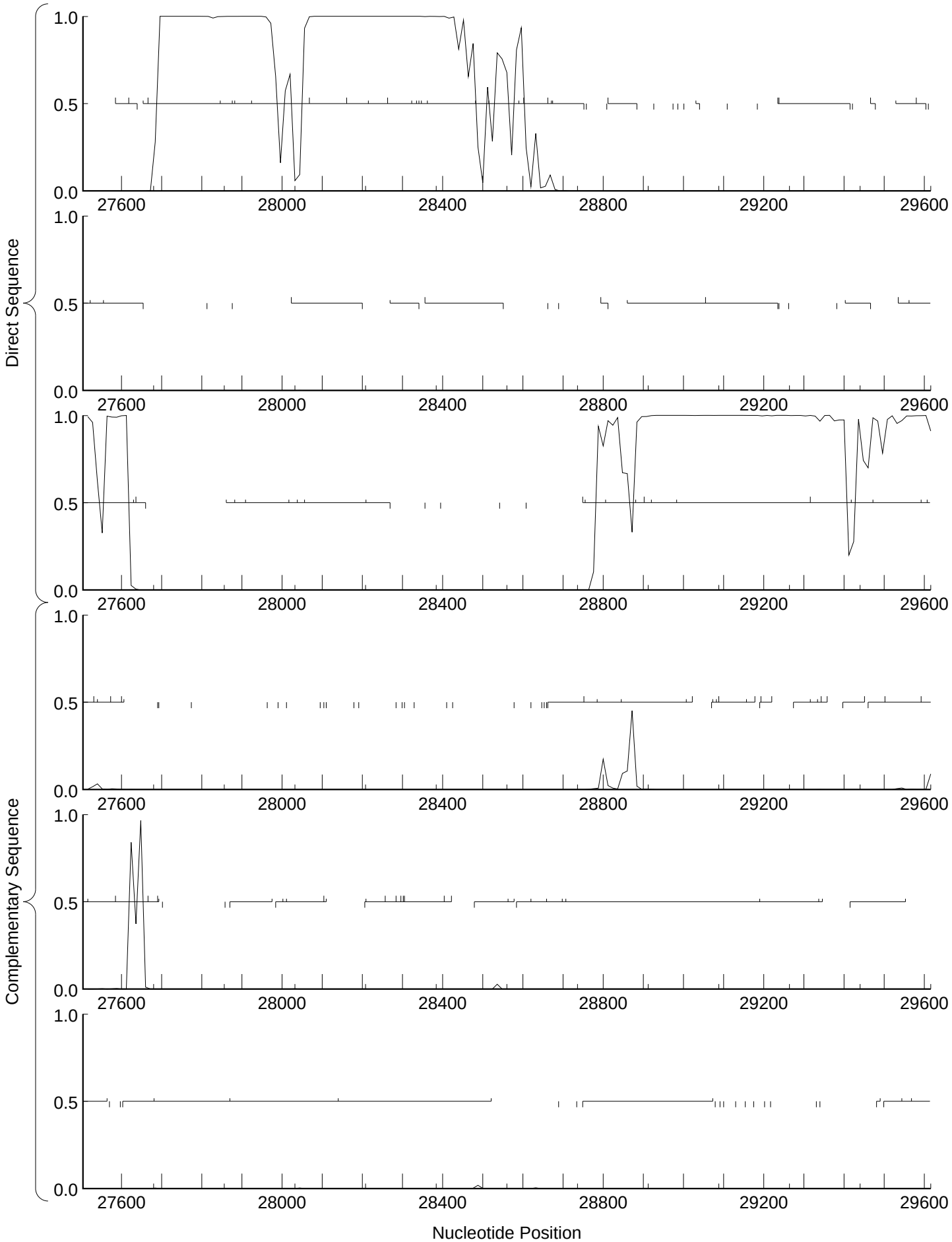
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 13/35



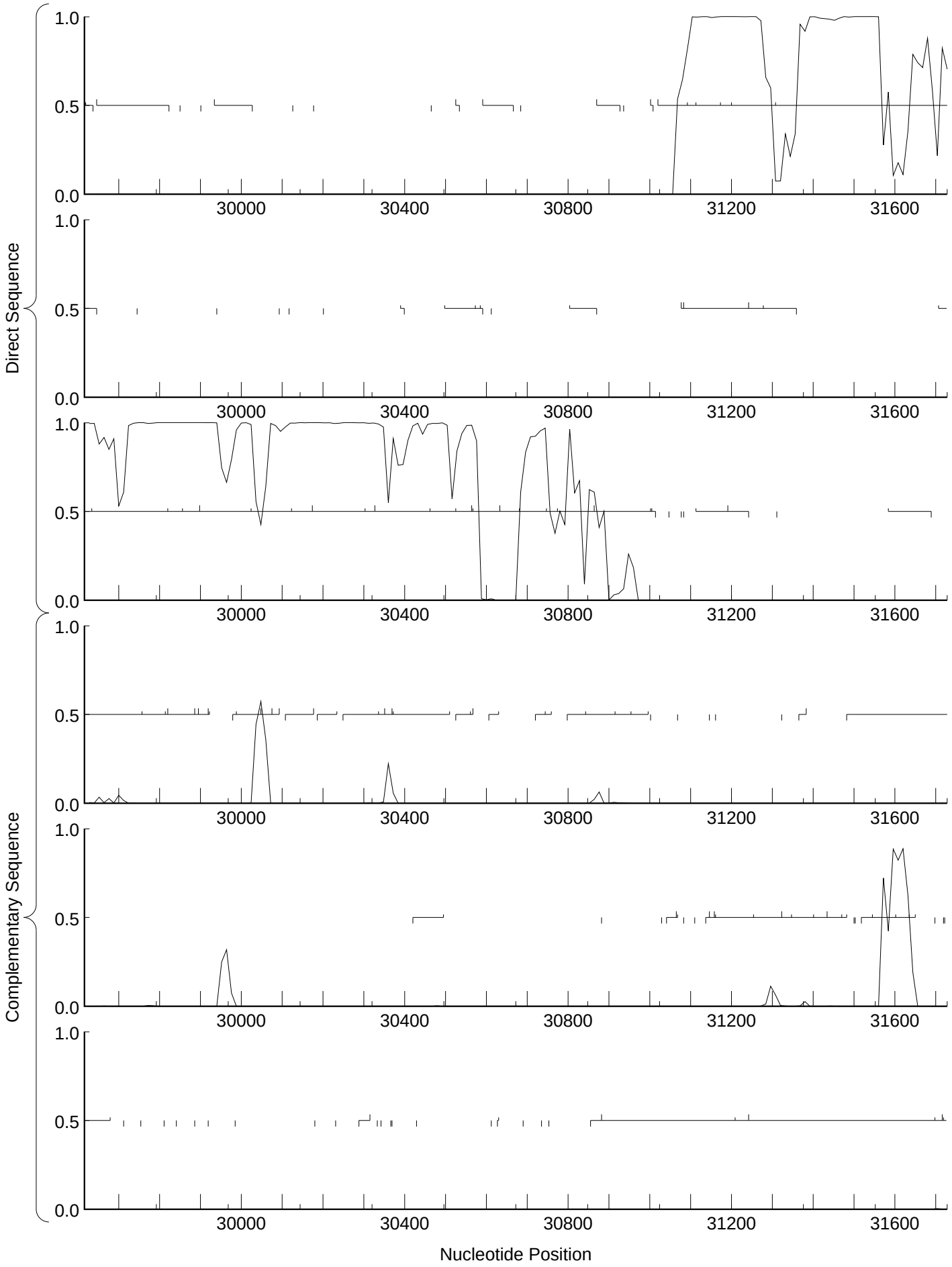
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 14/35



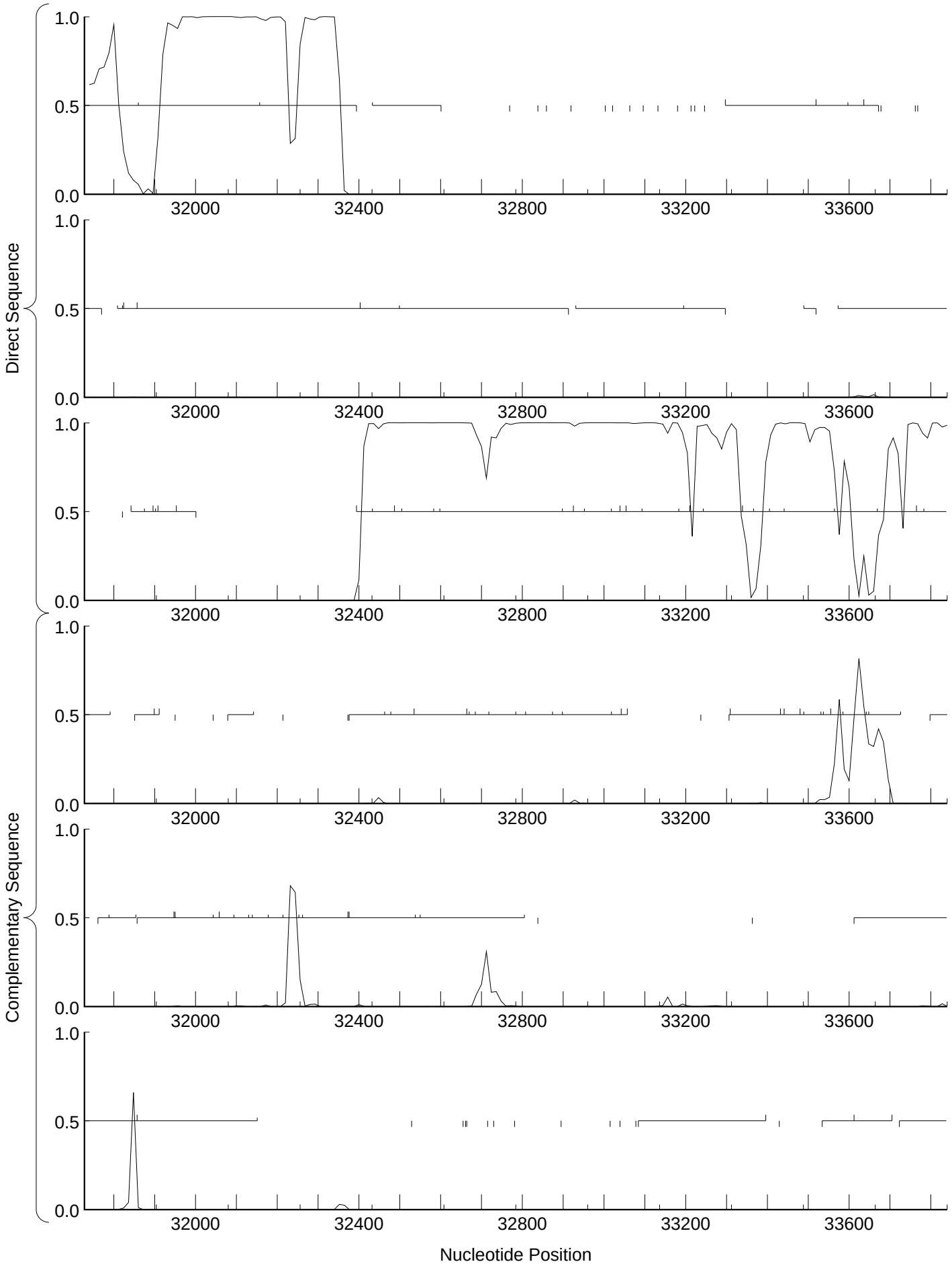
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 15/35



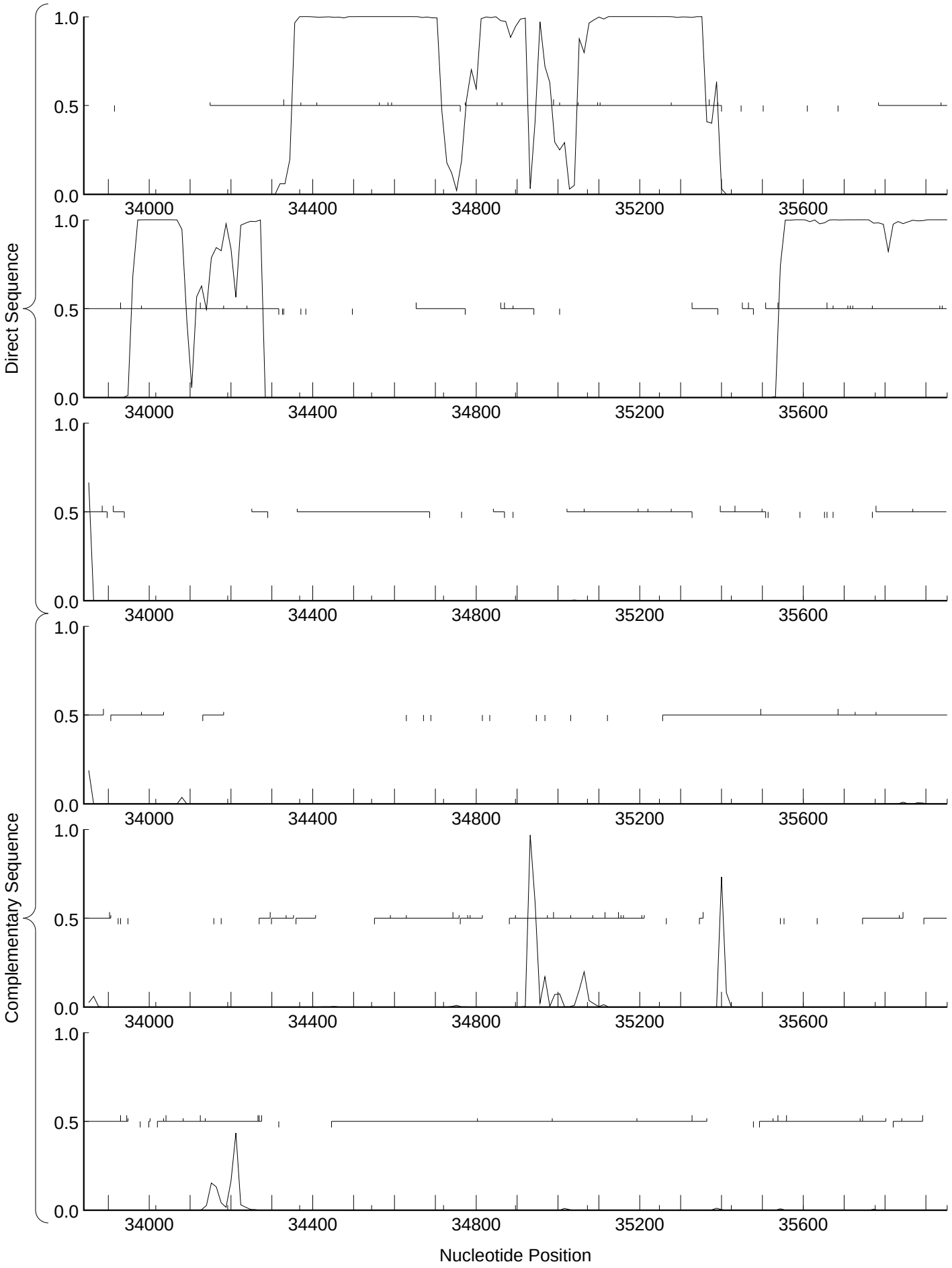
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 16/35



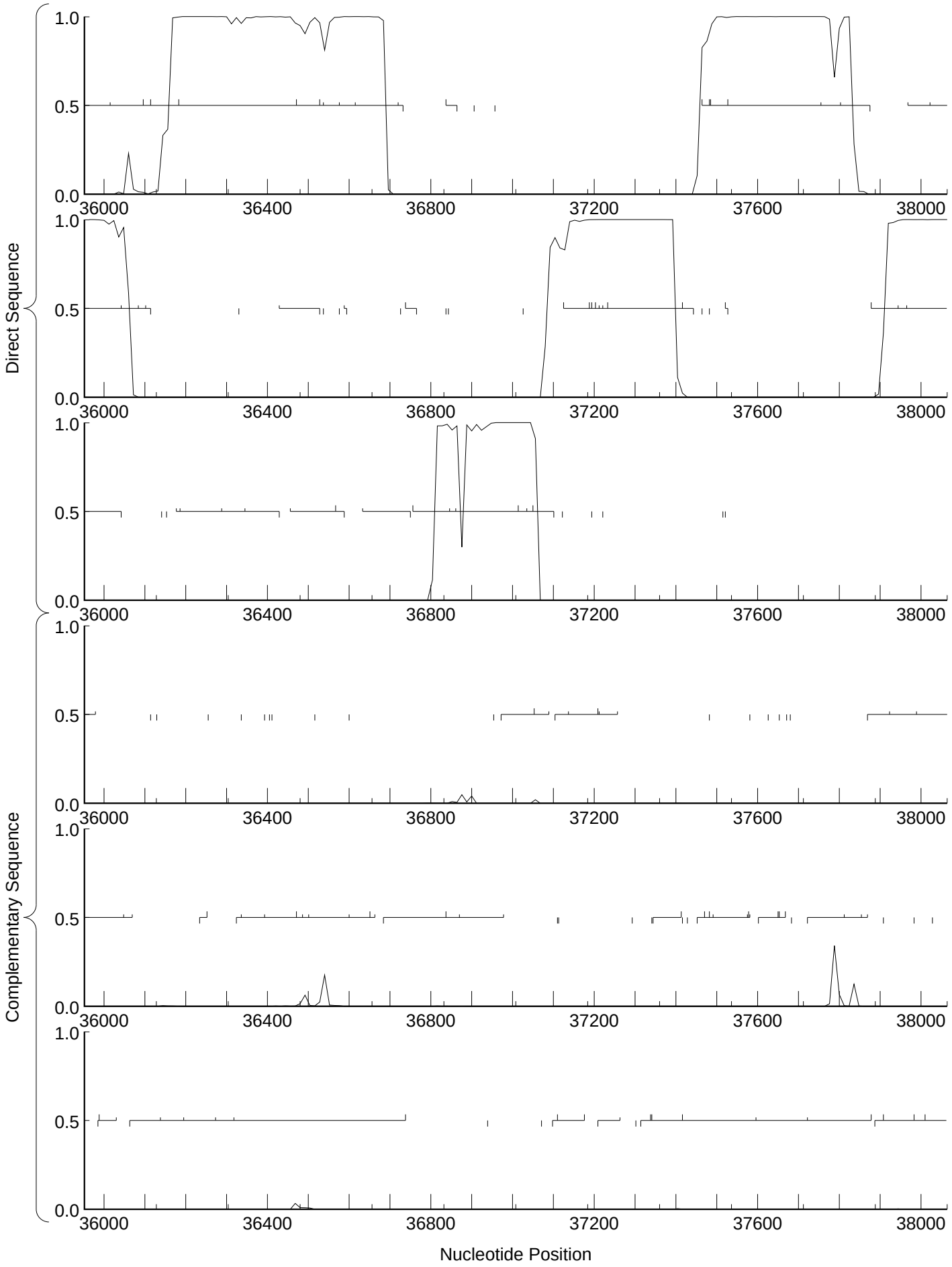
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 17/35



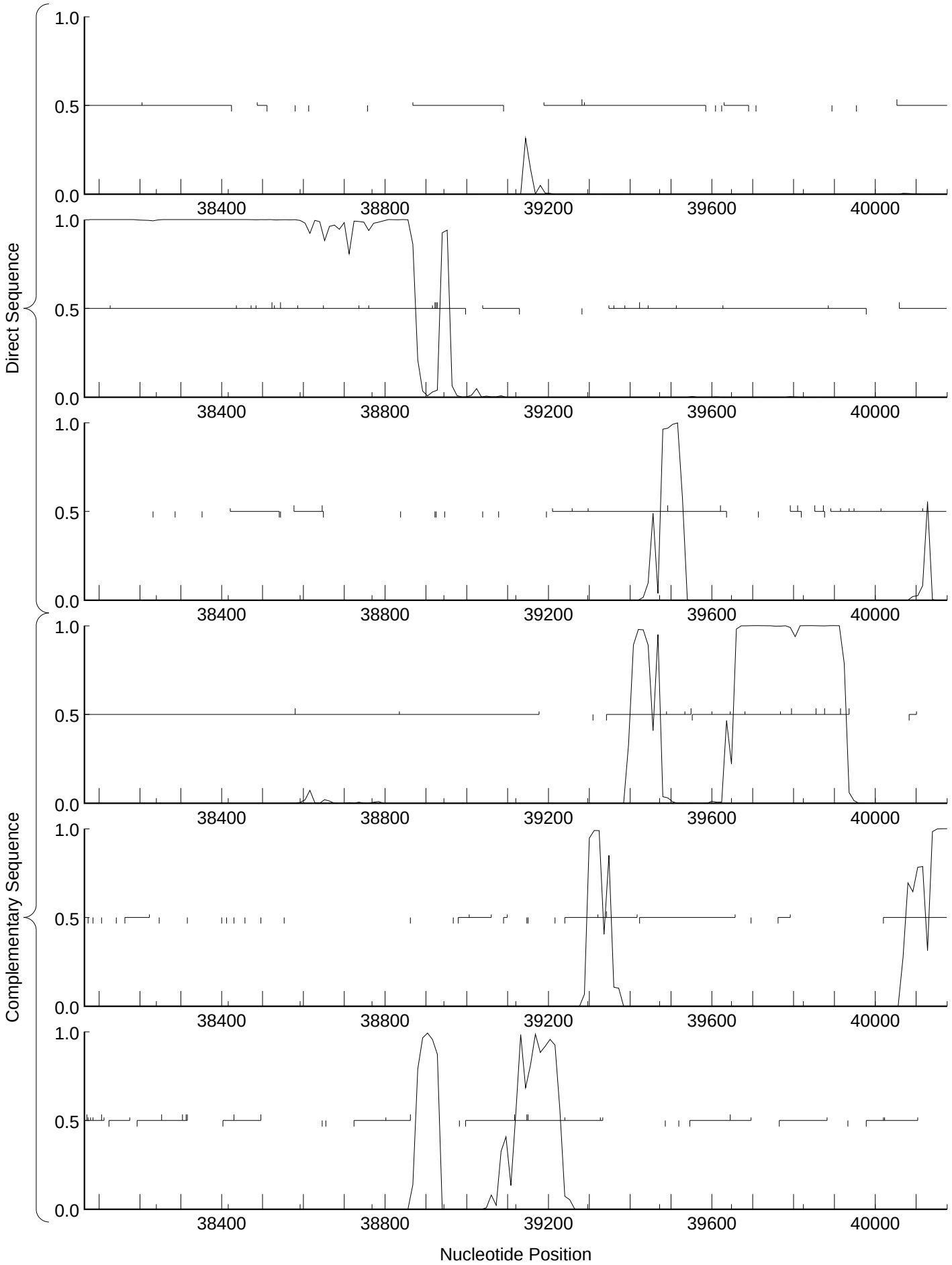
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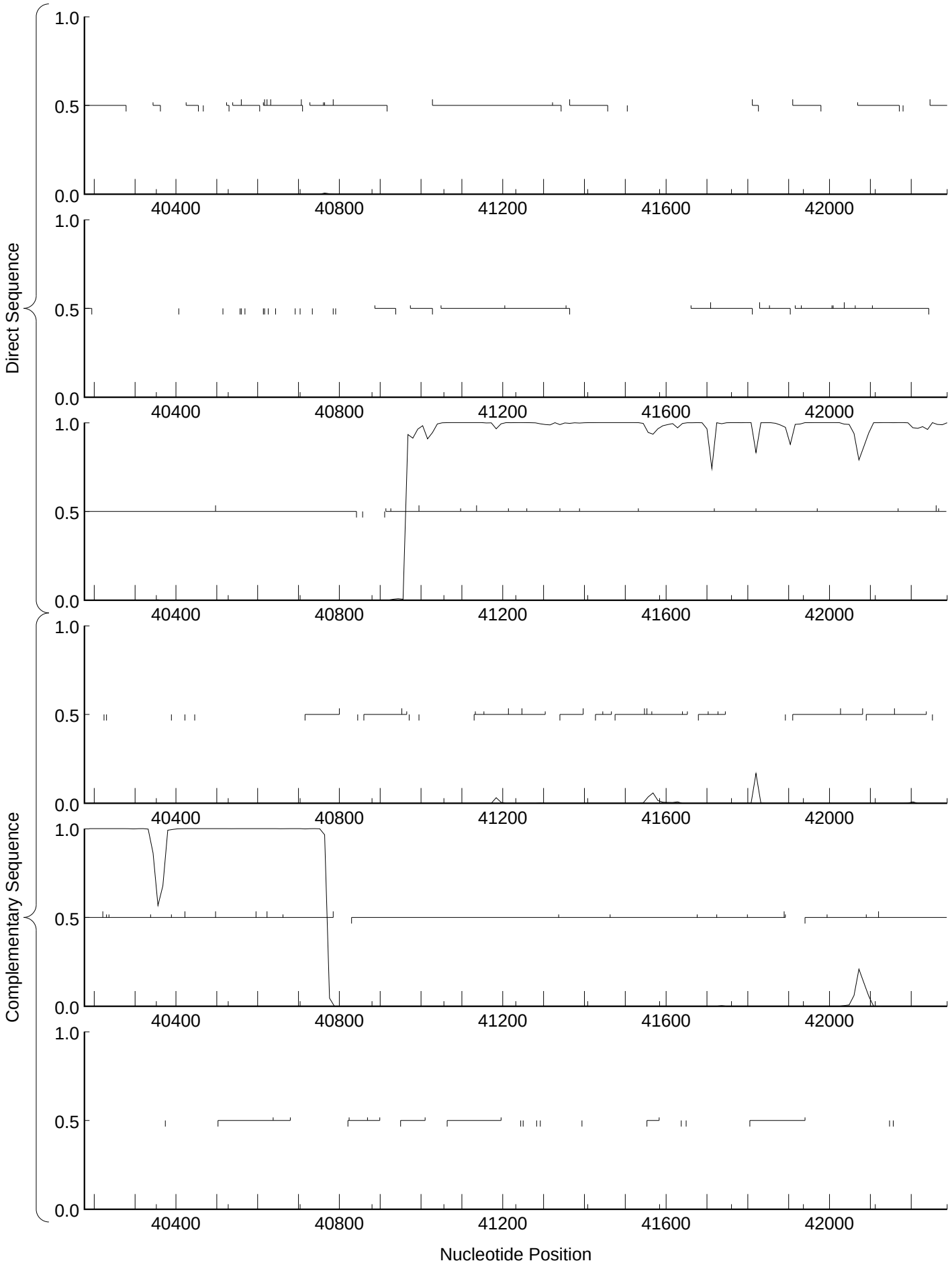
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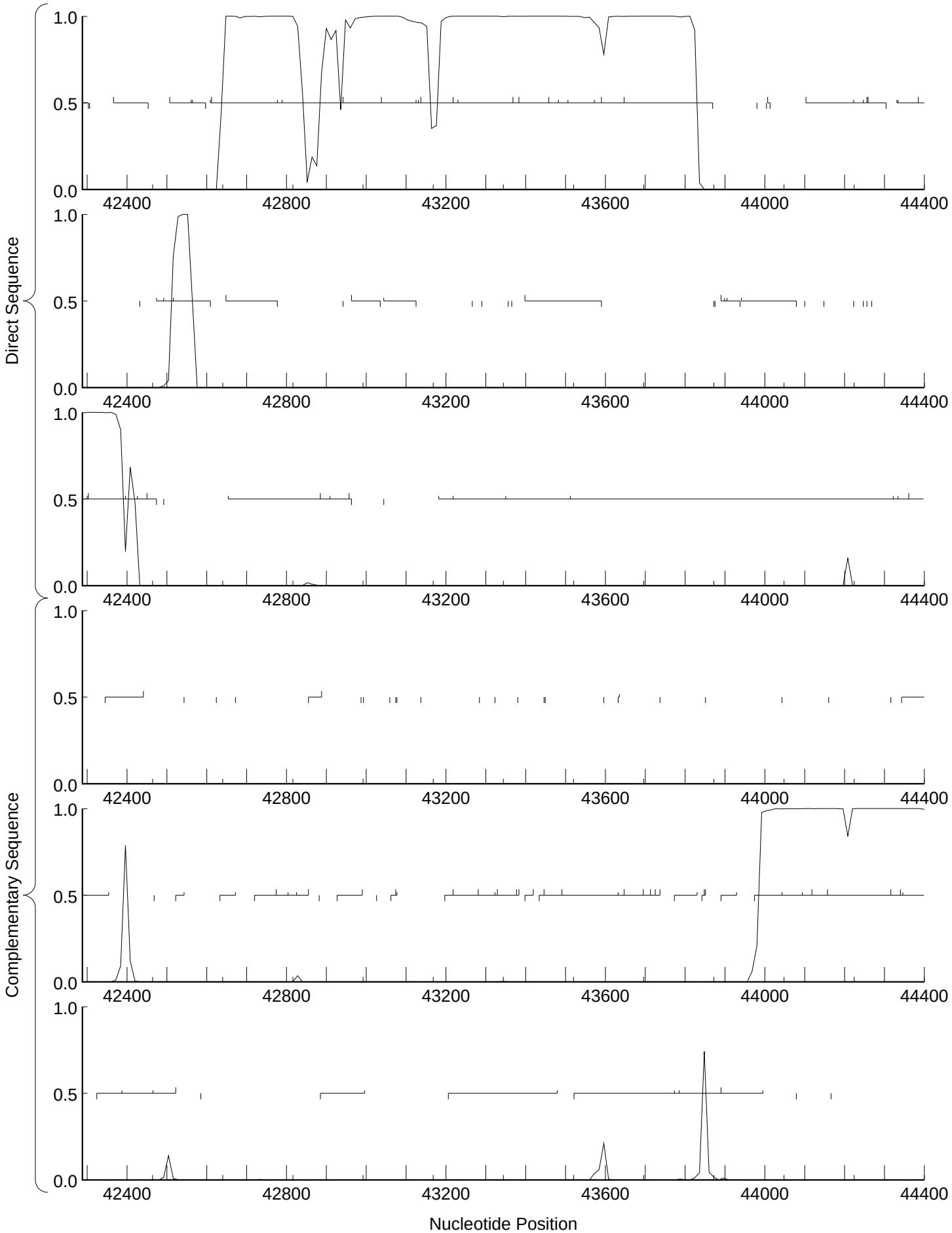
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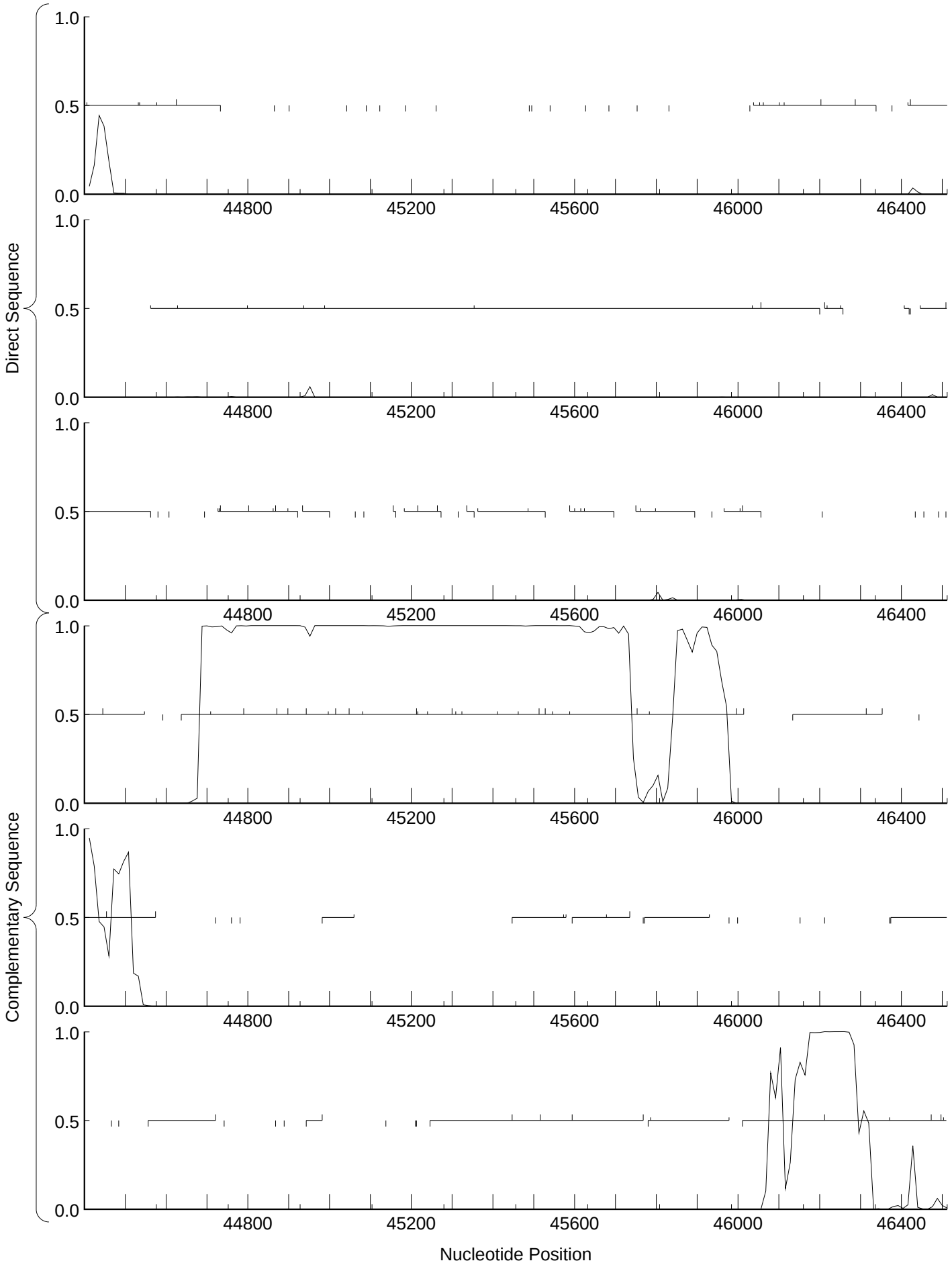
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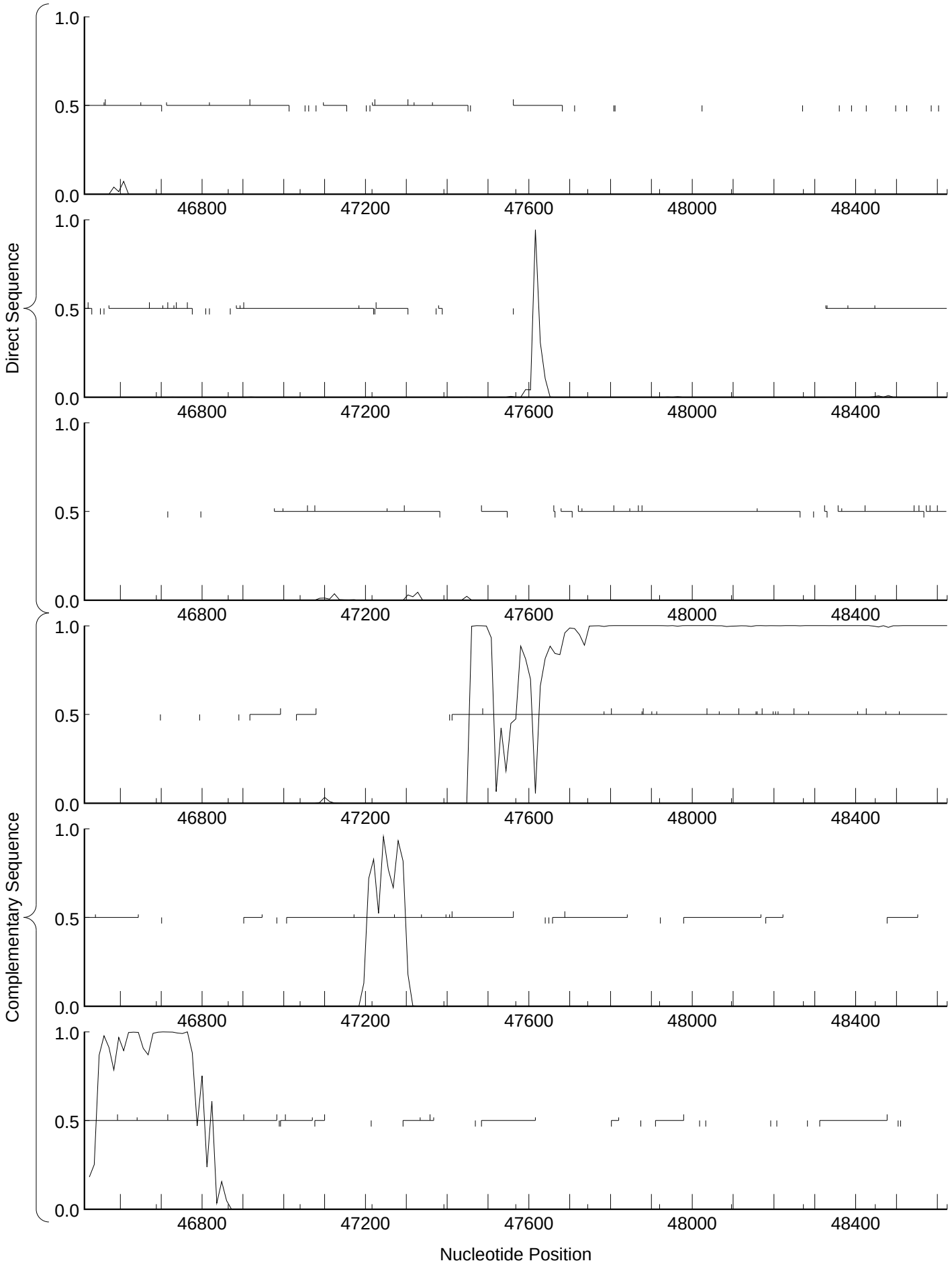
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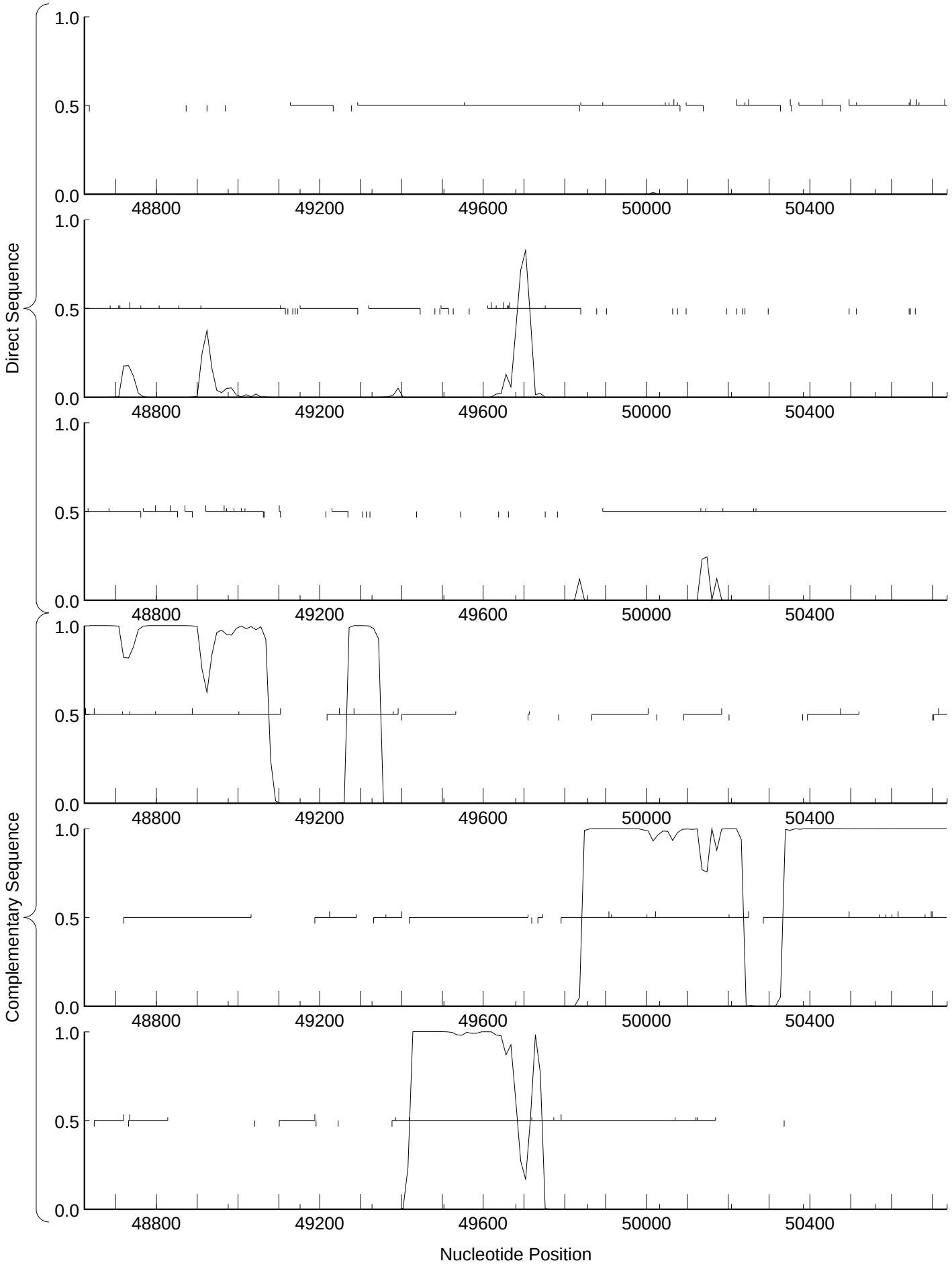
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 23/35



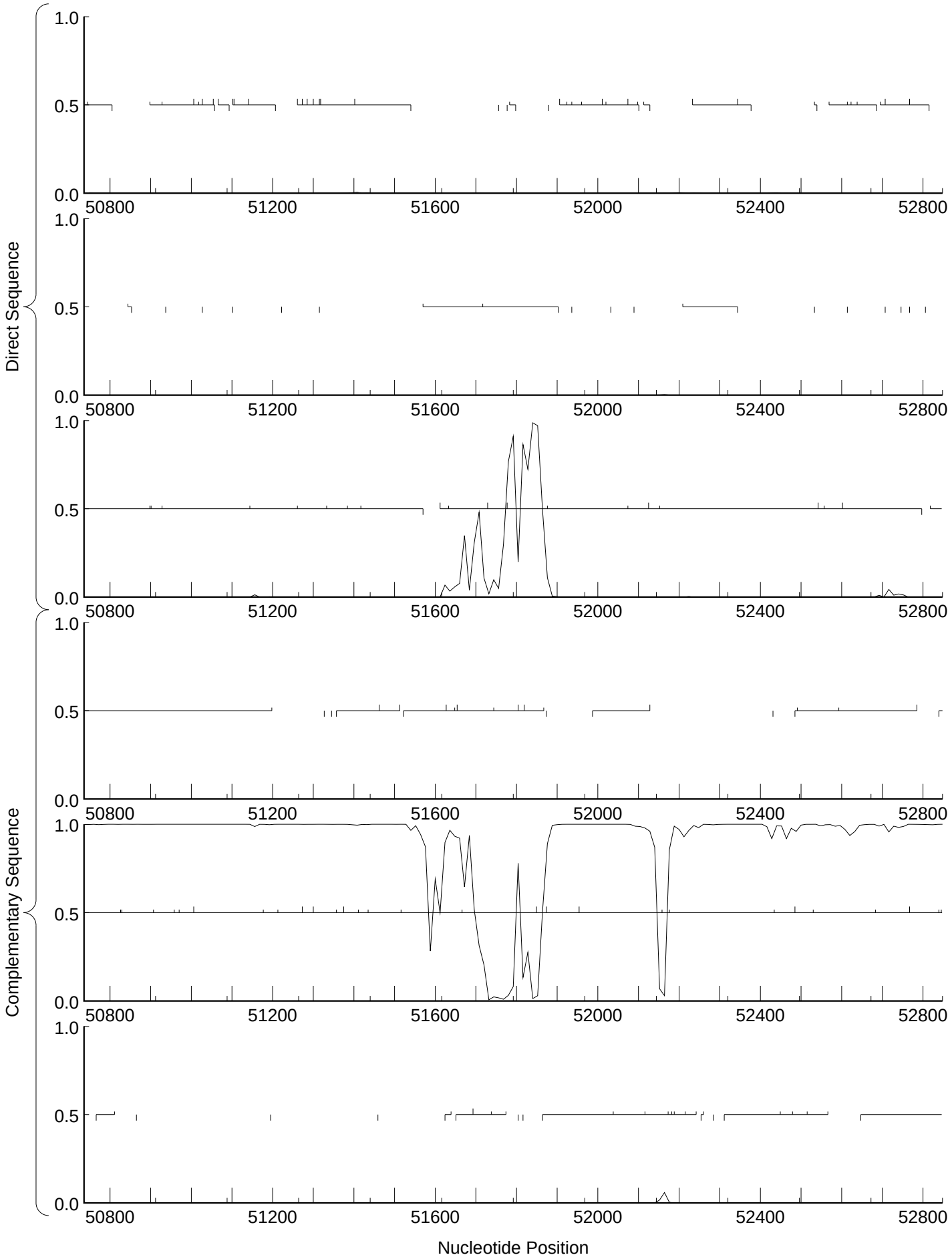
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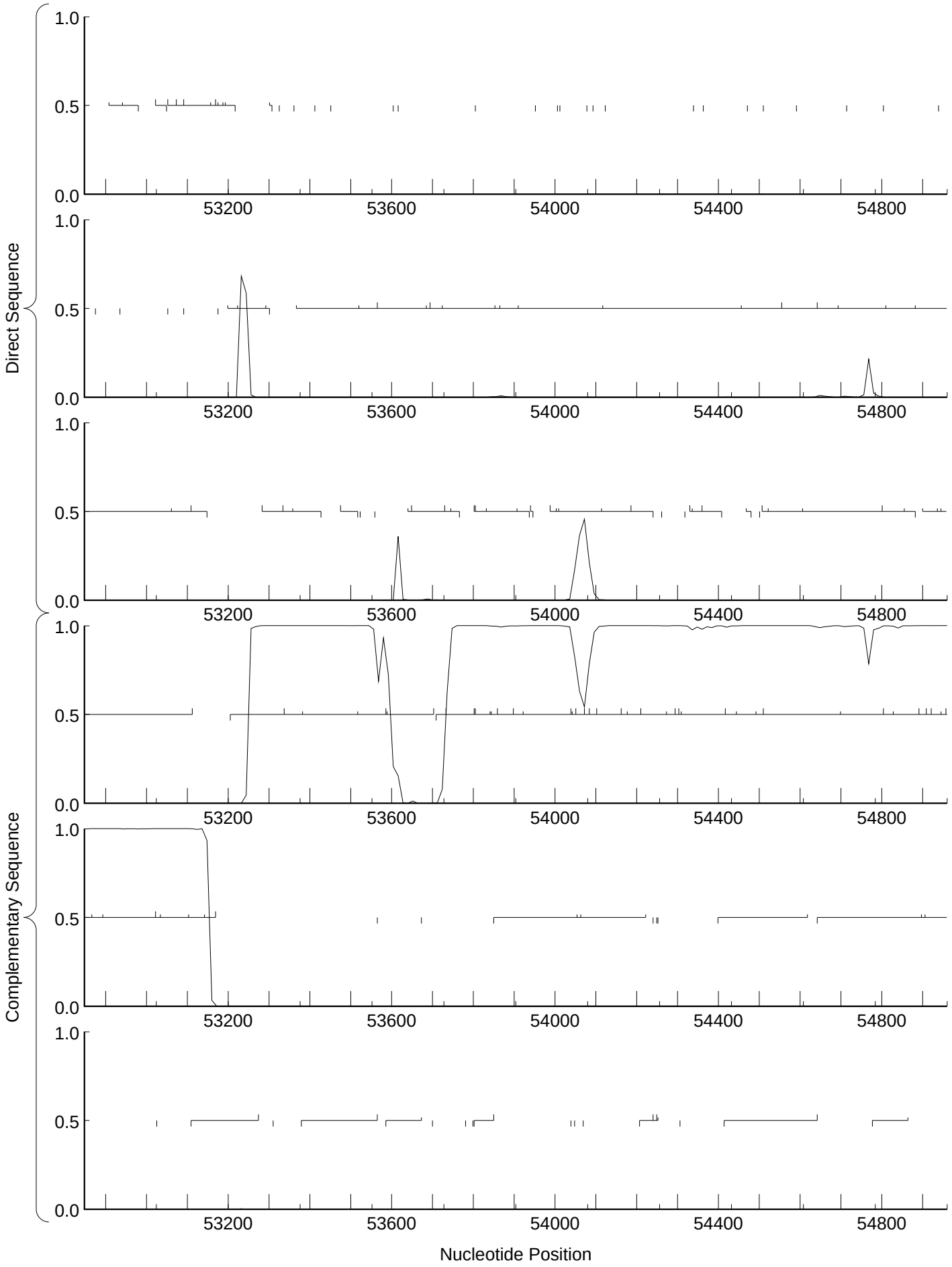
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 25/35



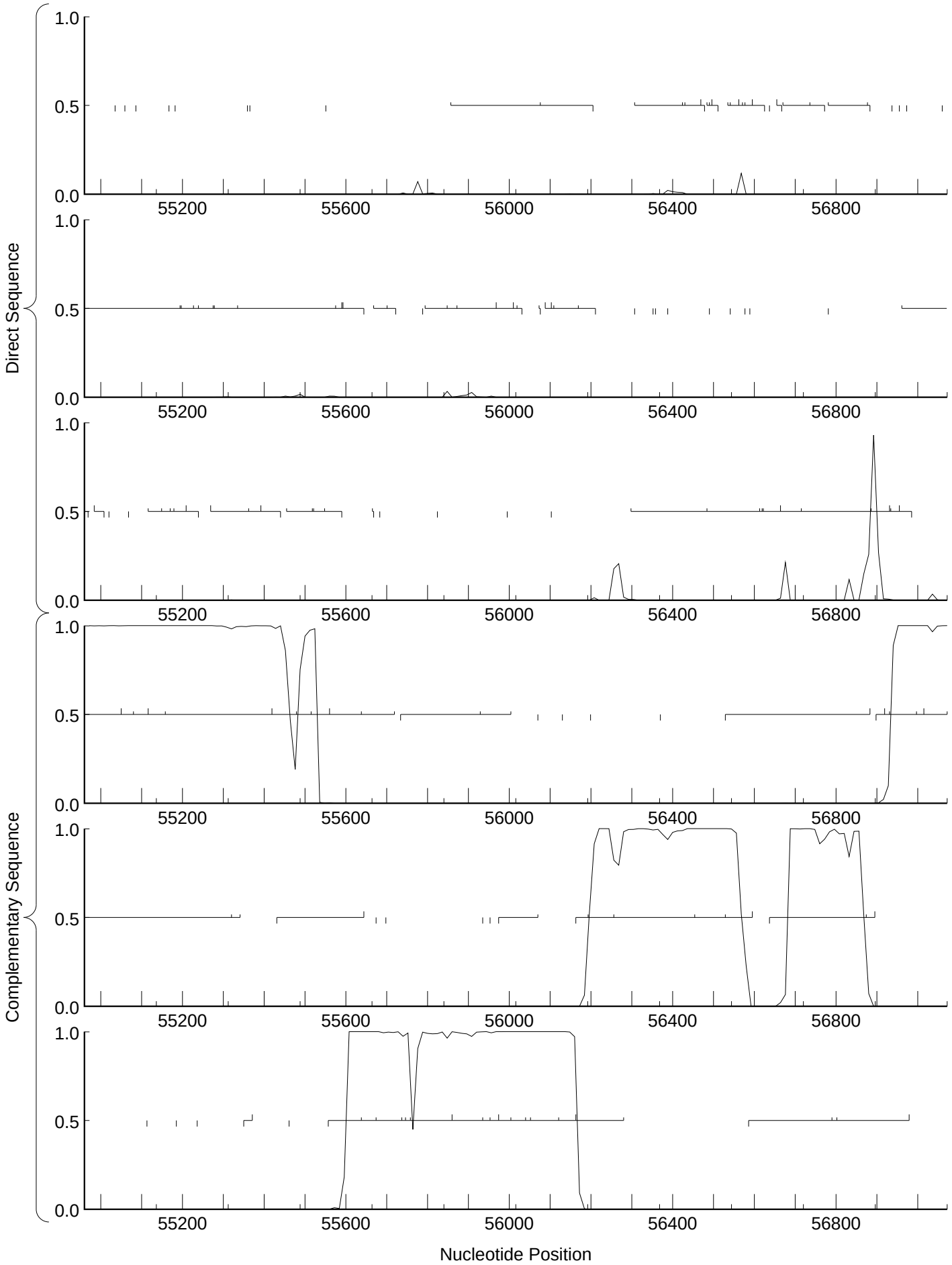
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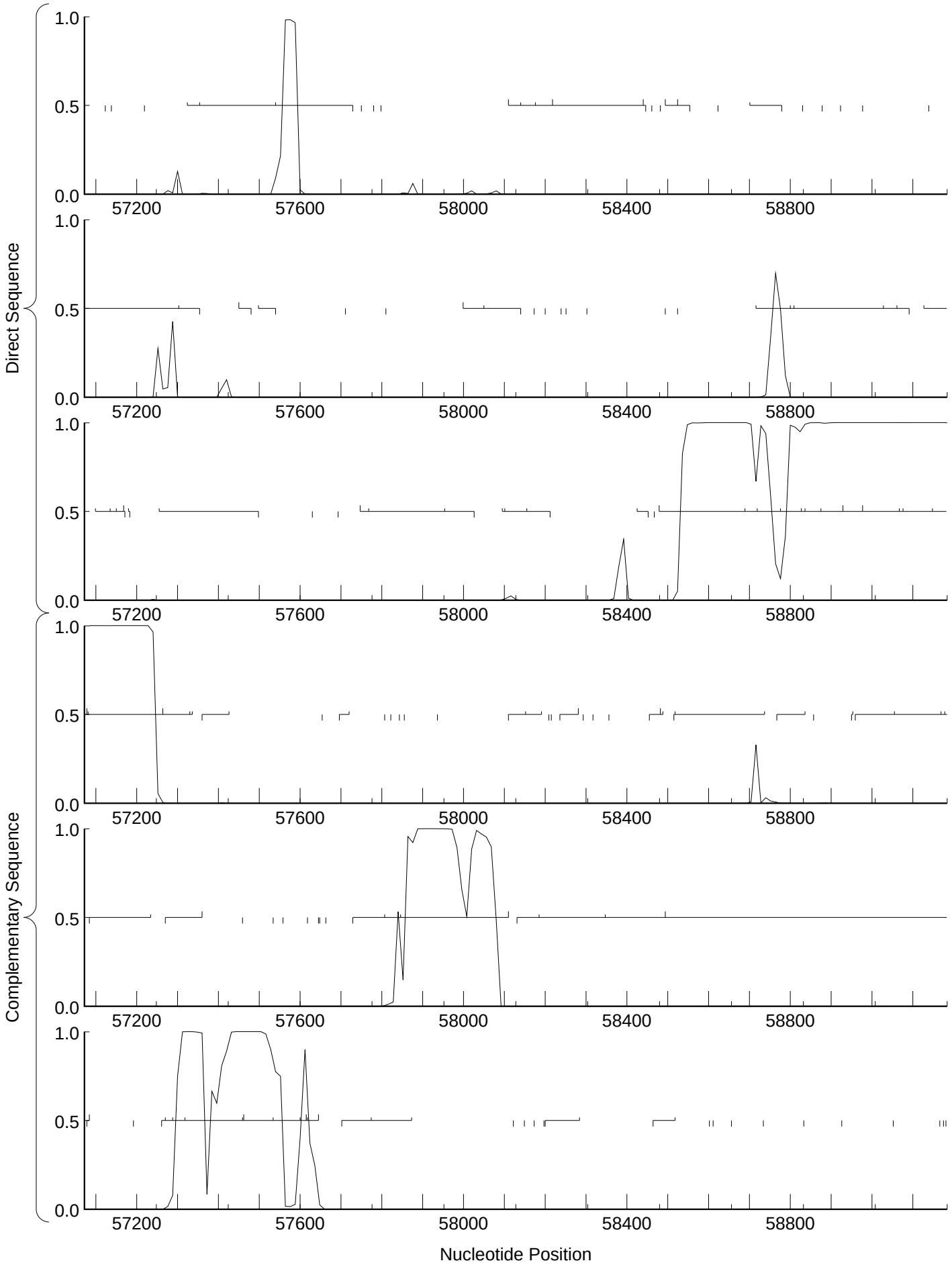
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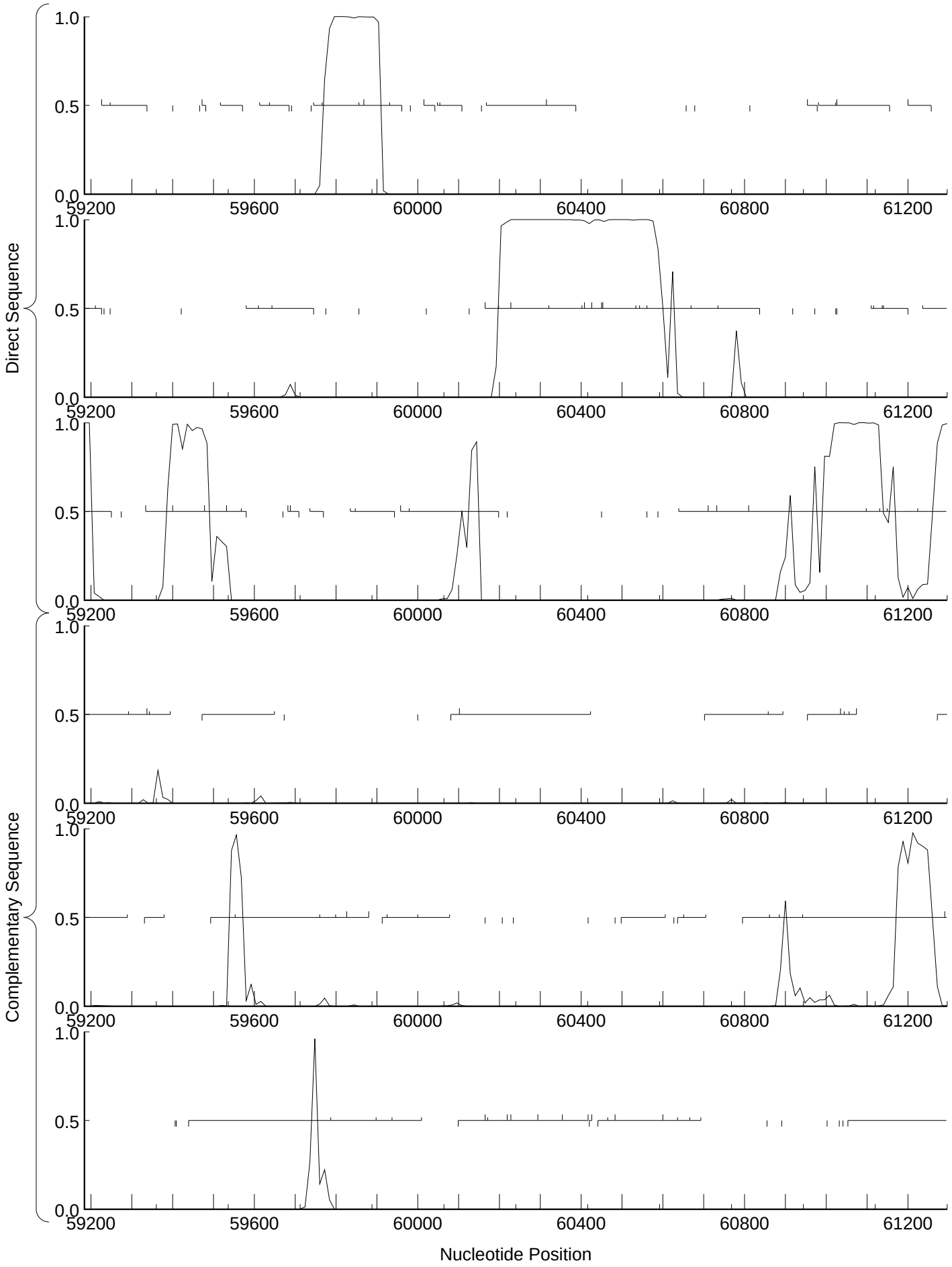
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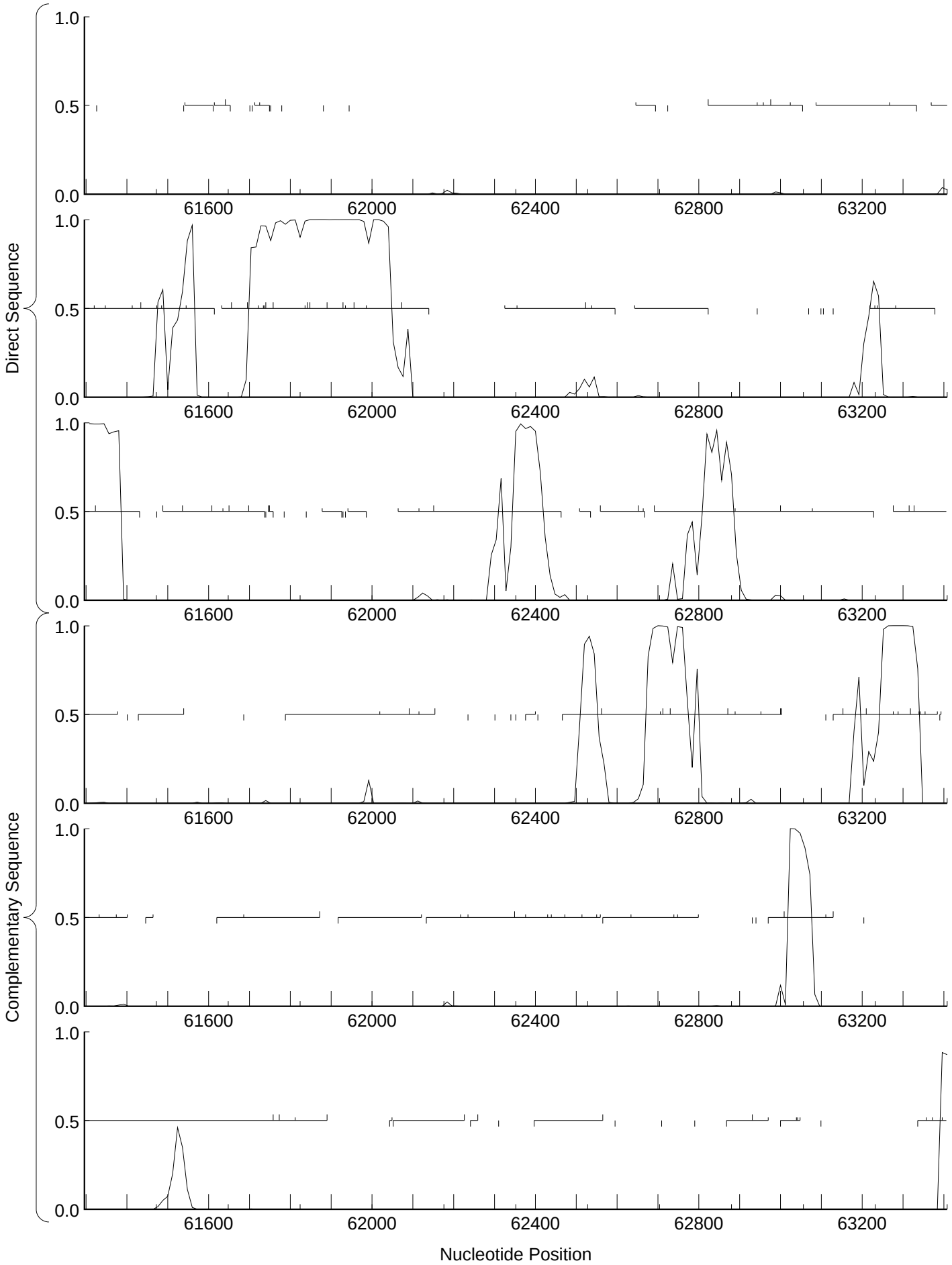
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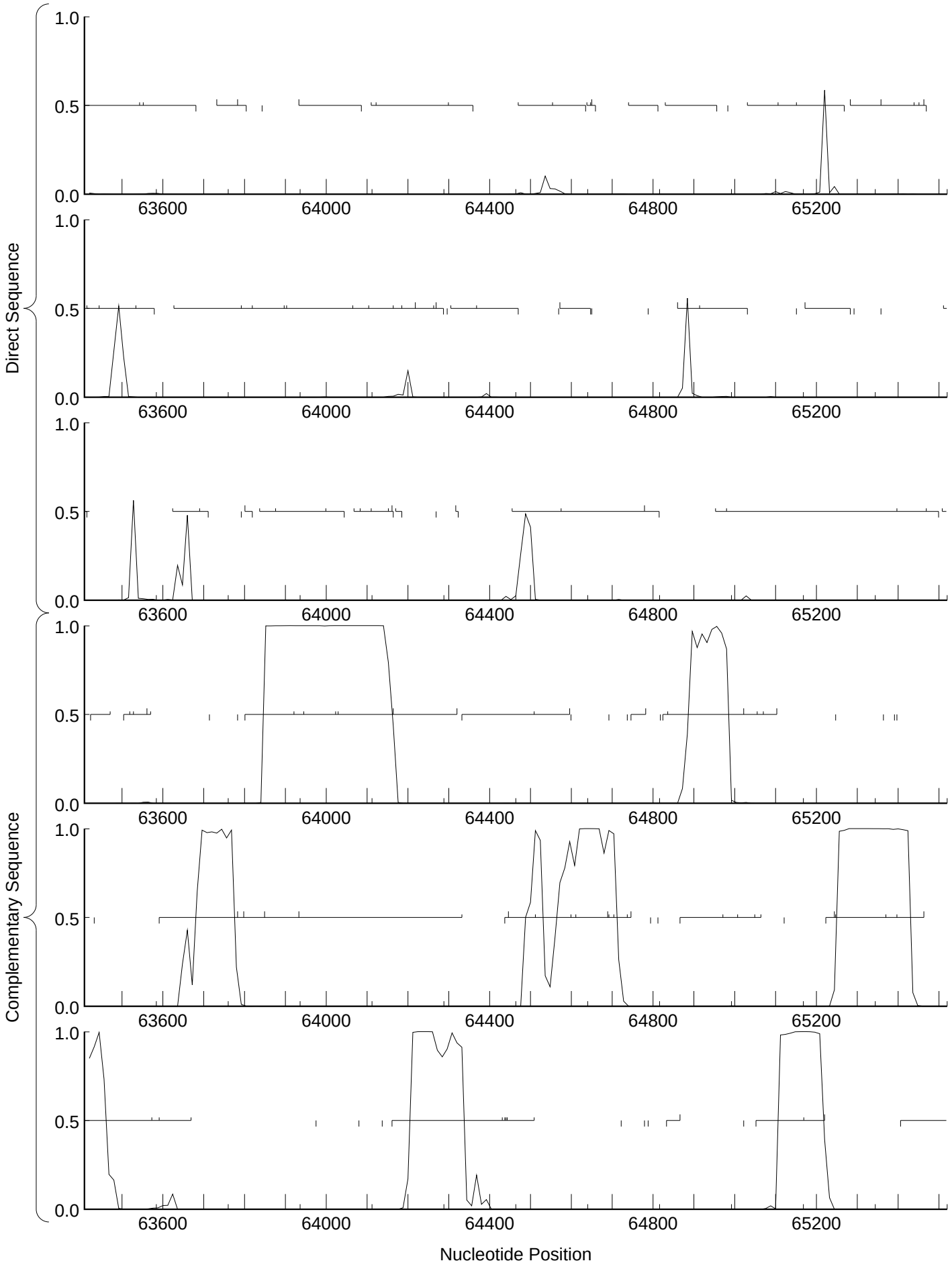
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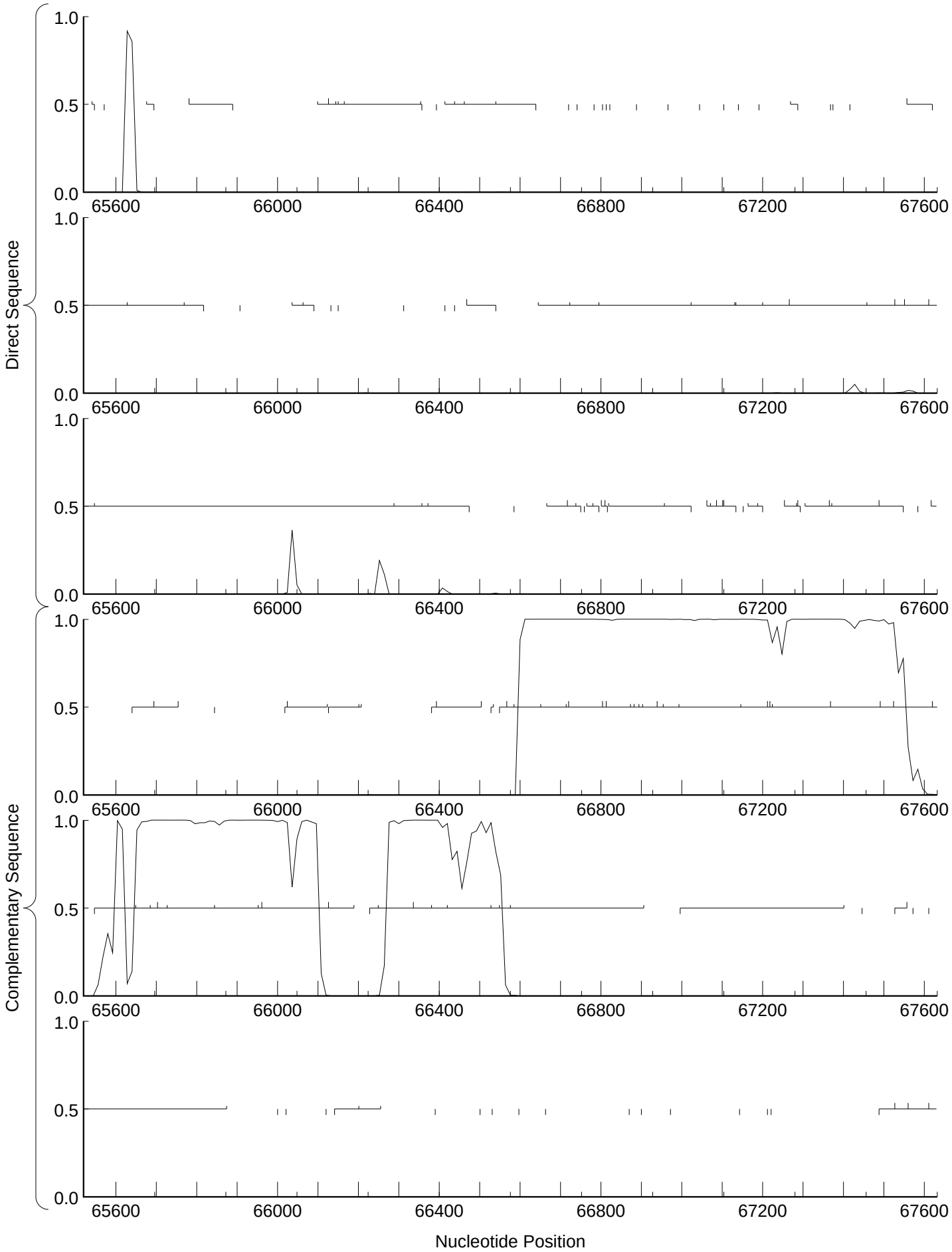
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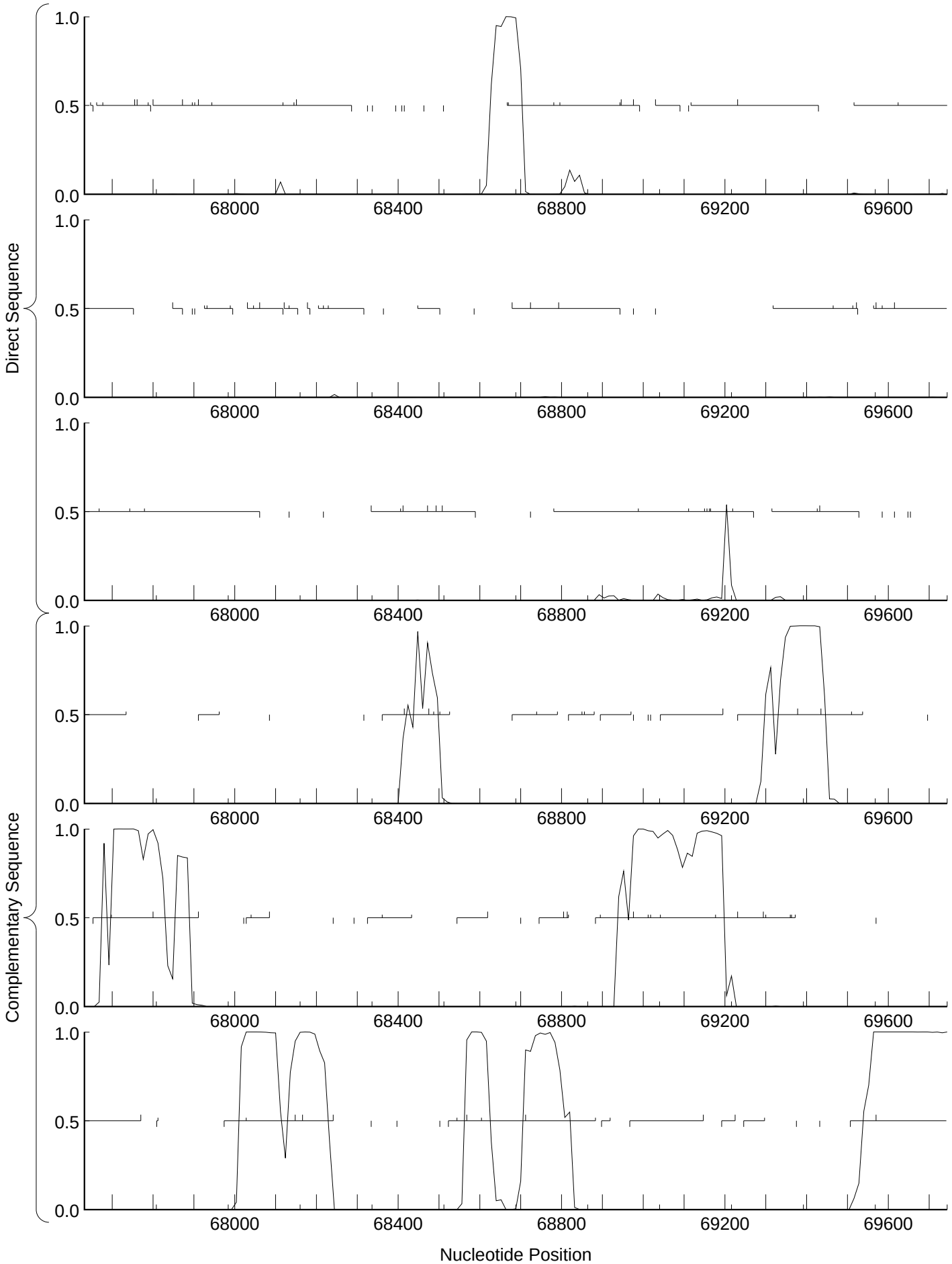
Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 32/35



Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 33/35



Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 34/35



Mycobacterium phage Baee complete sequence, 70270 bp, circularly permuted, Cluster B5, Order 4, Window 96, Step 12, 35/35

