GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA
Analysis Date    :  3/29/24 at 10:30:33
Pages            :  20
Sequence Length  :  38684 bp
GC Content       :   68.04%

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

PS-Version       :  1.2

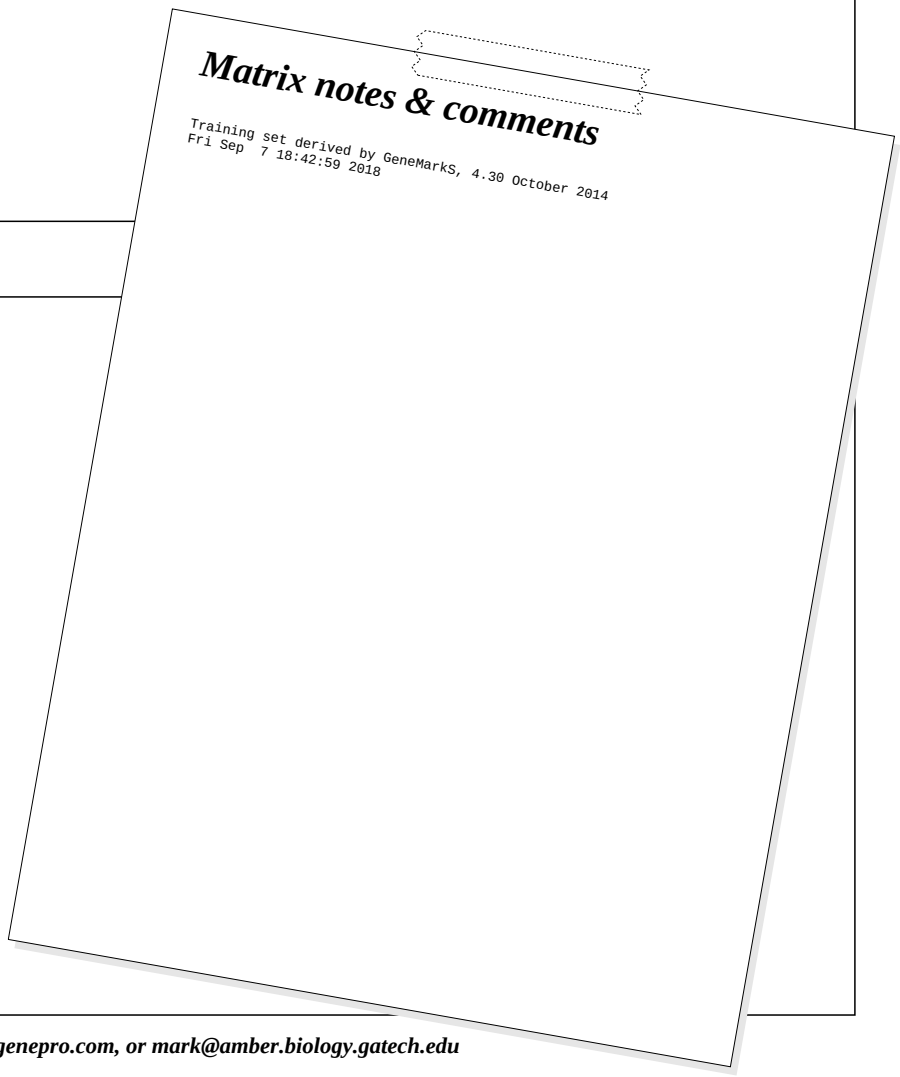
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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                    List regions and/or splice sites,
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MATRIX INFORMATION

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Author     :  -

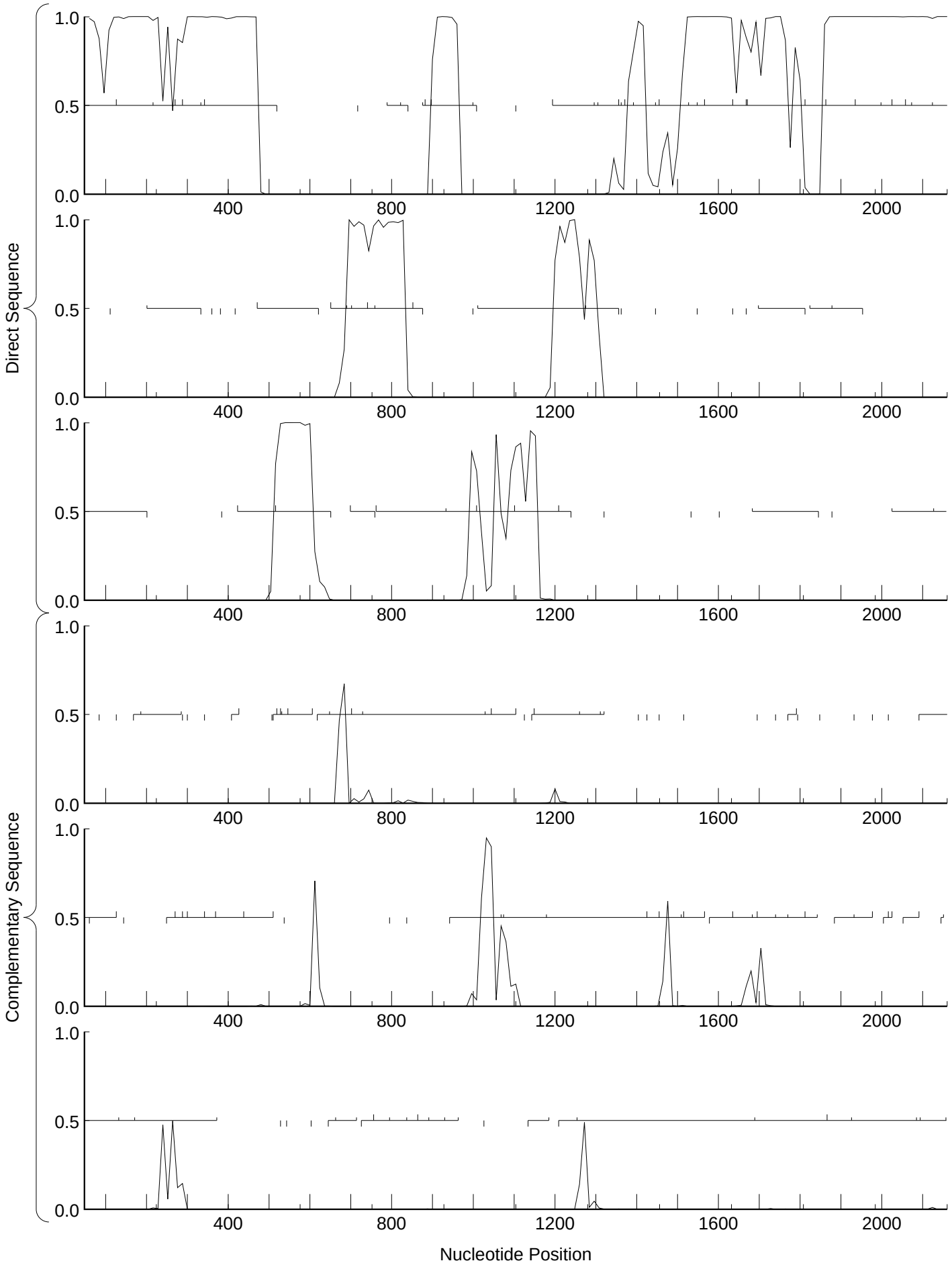
Order      :  4
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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

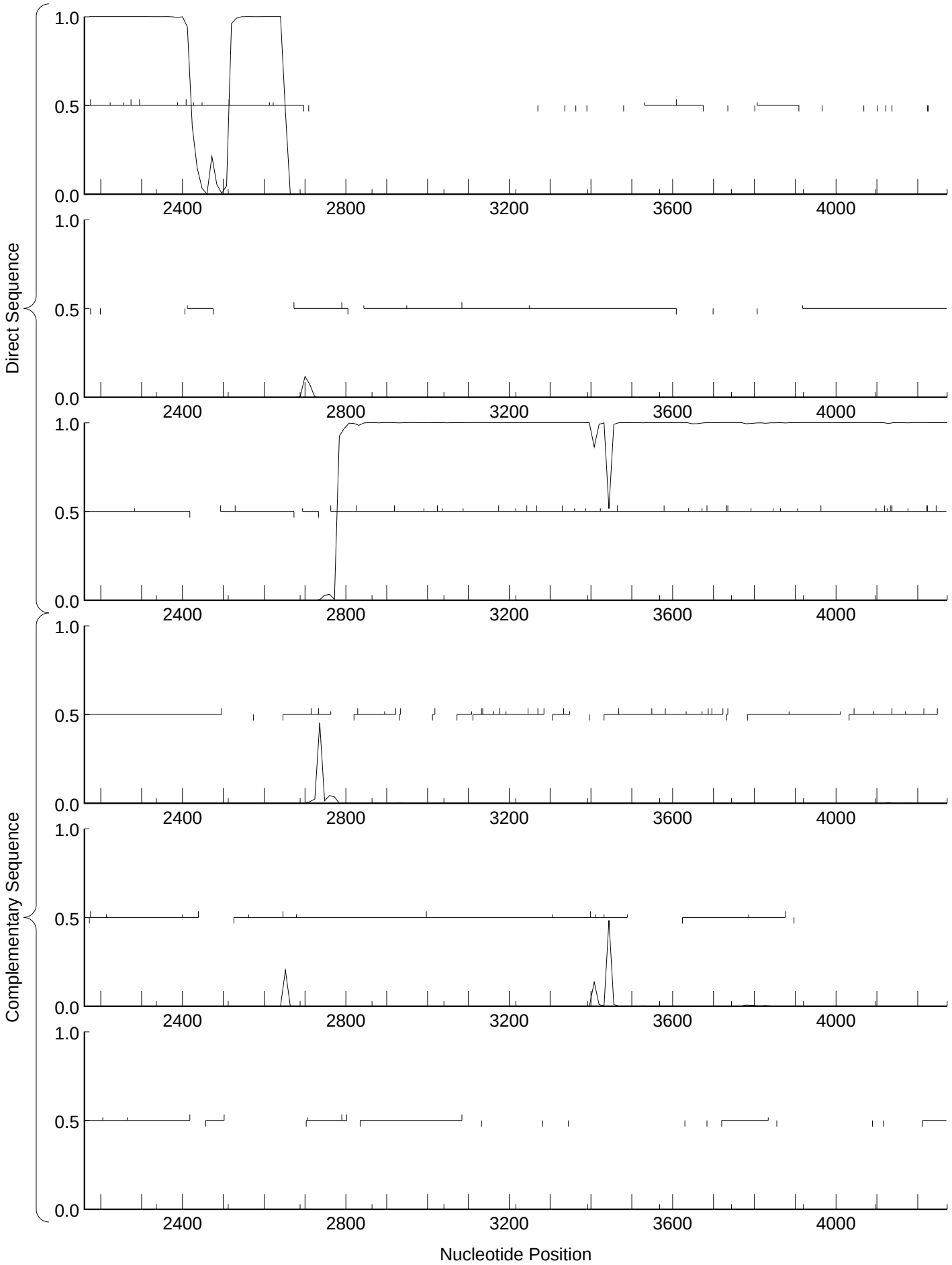



For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

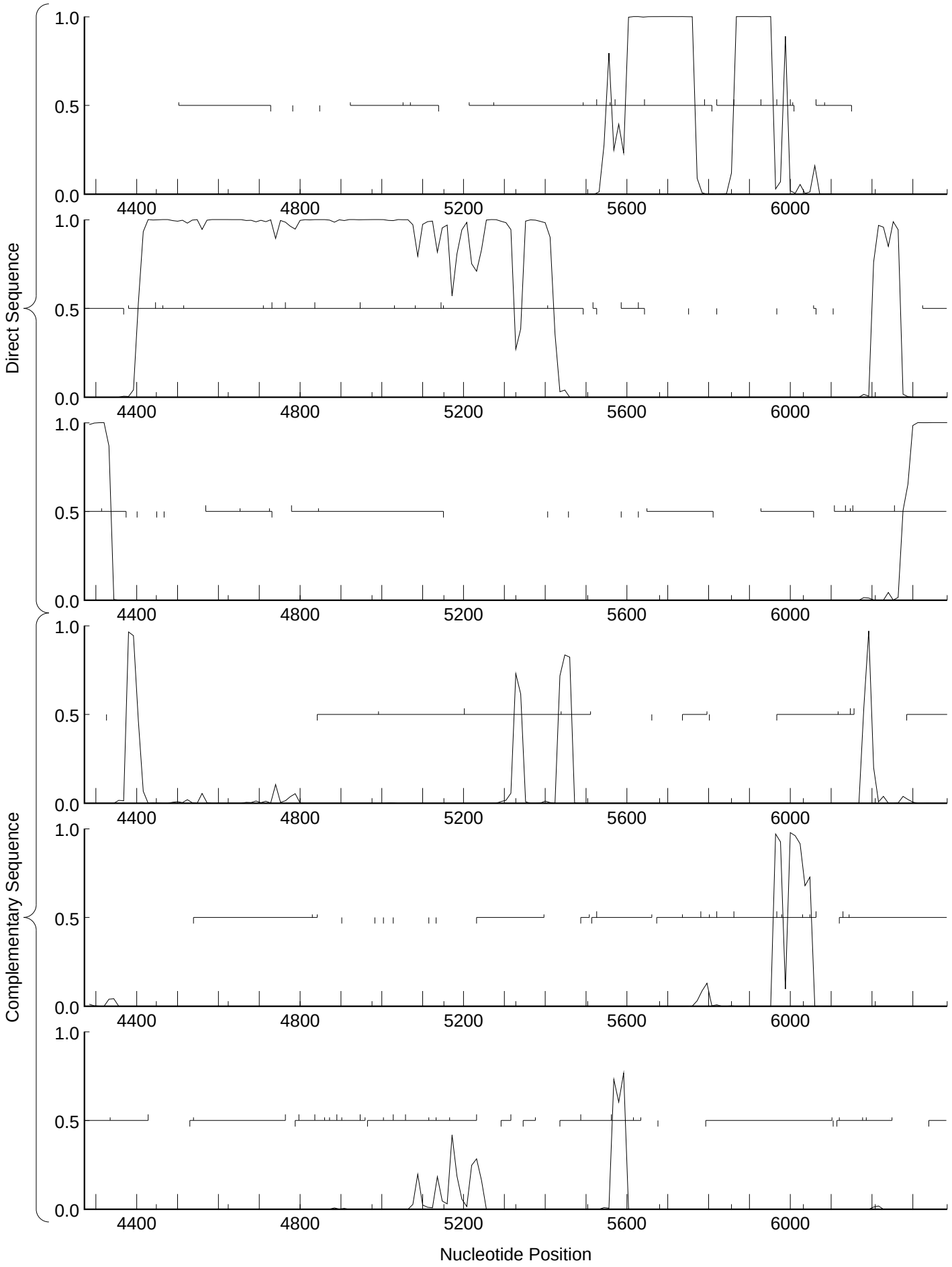
Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 2/20



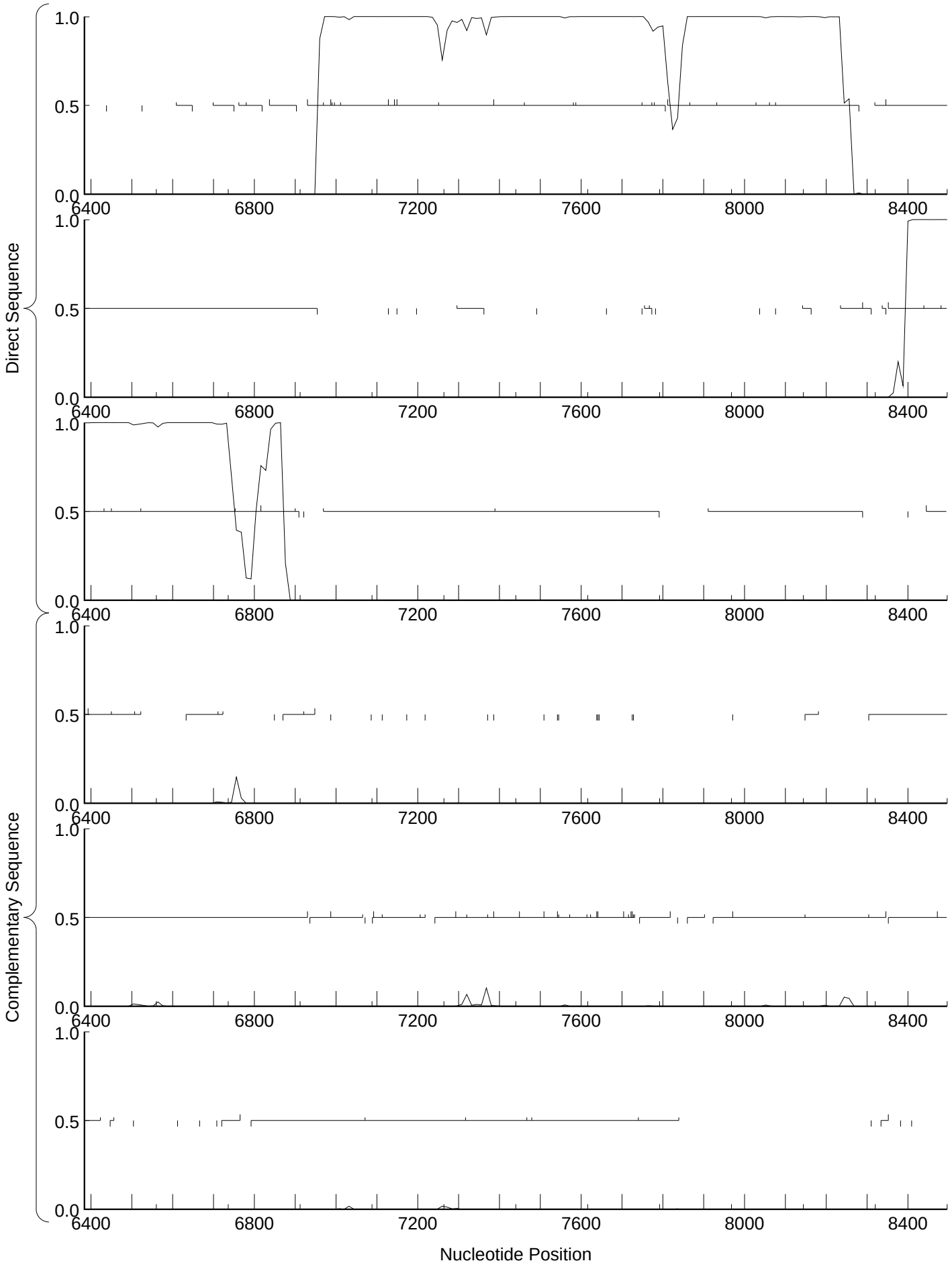
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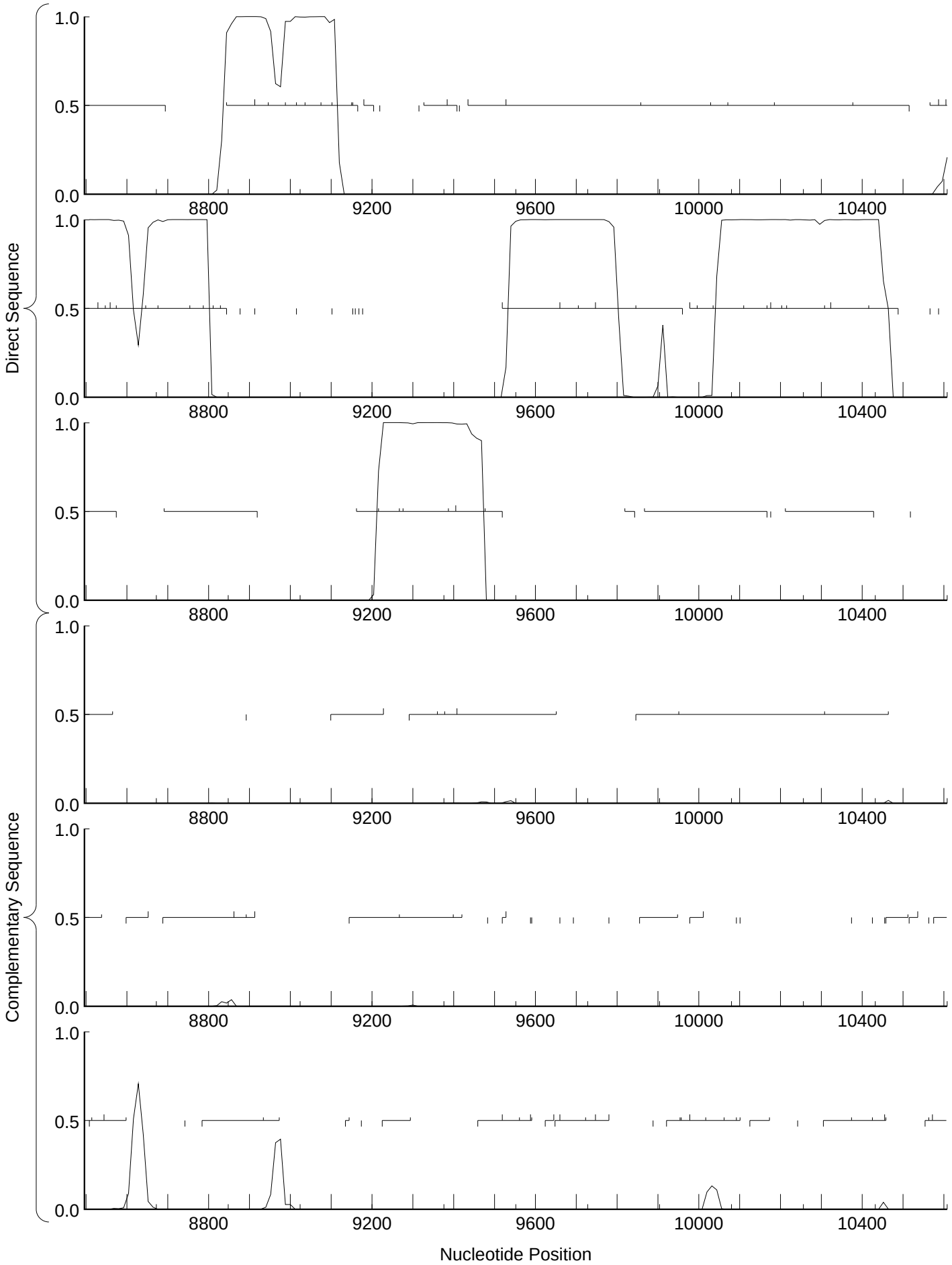
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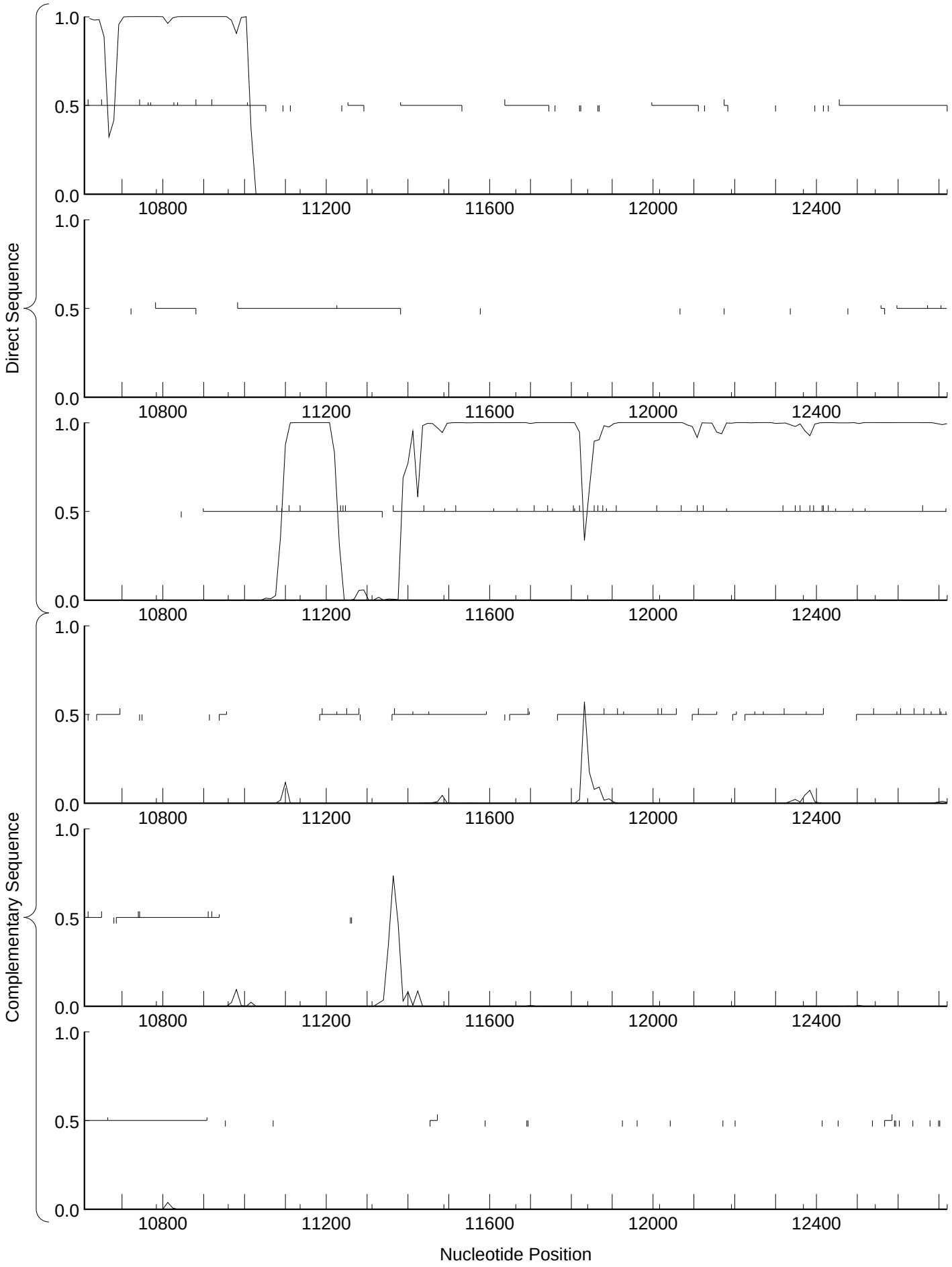
Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 5/20



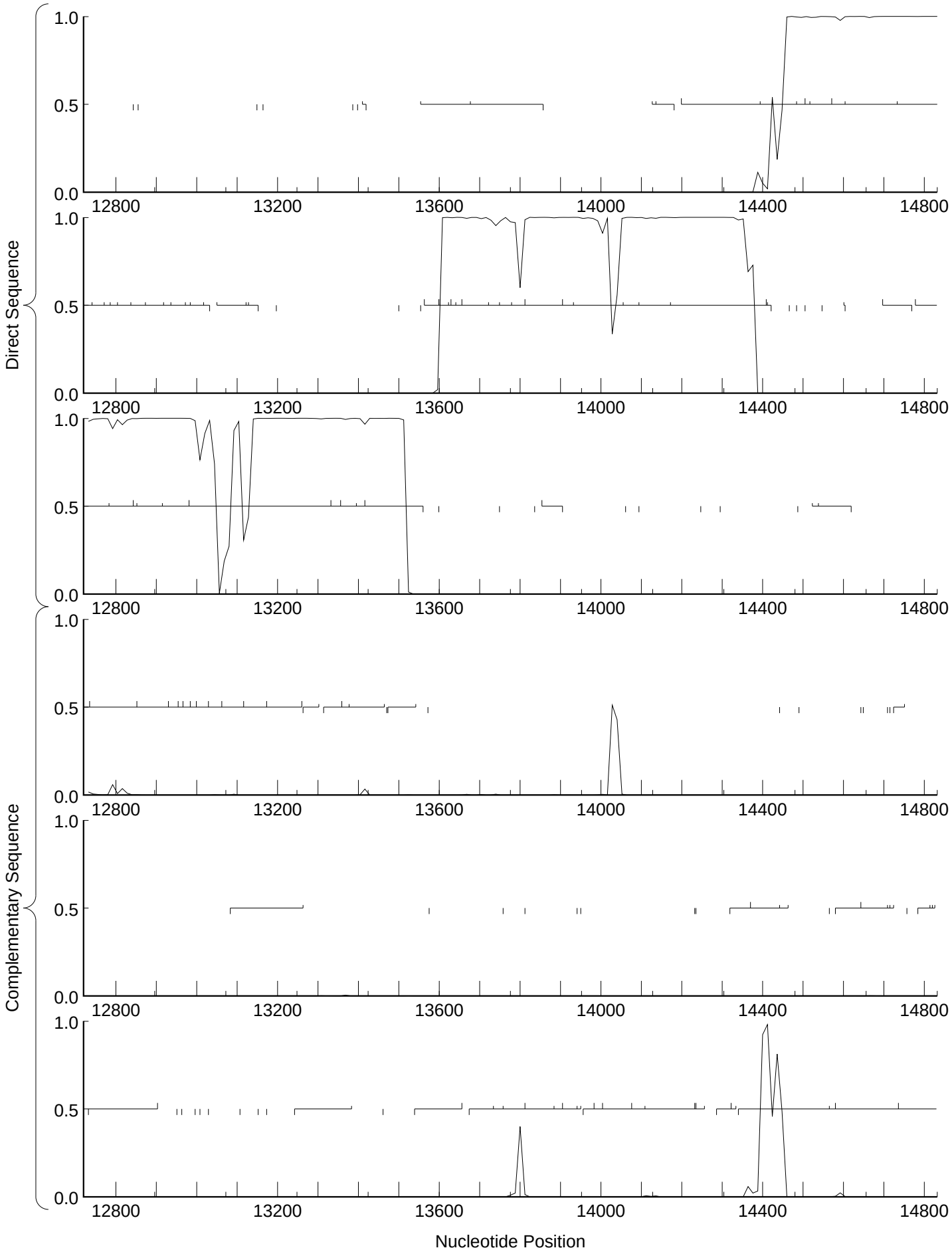
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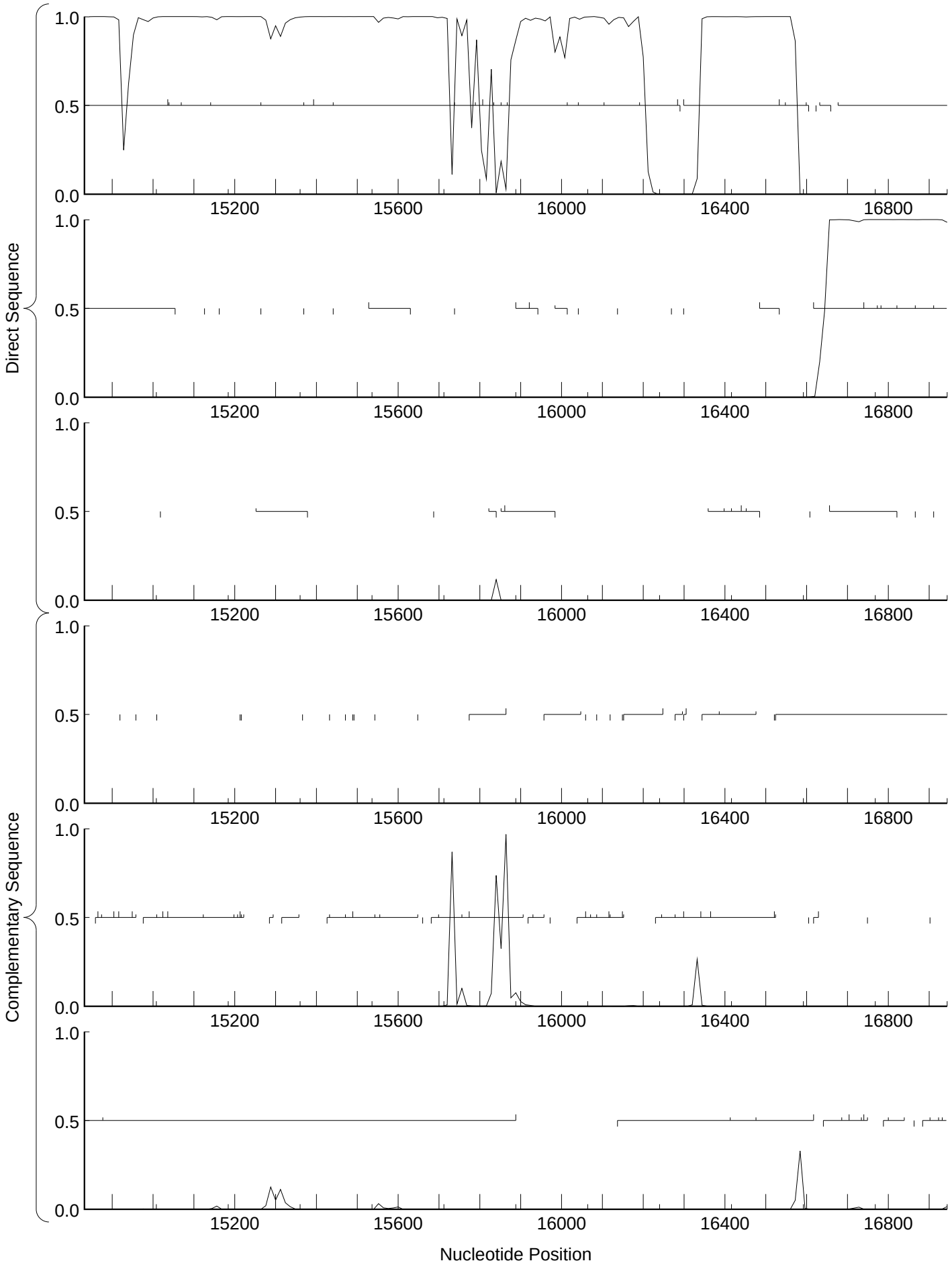
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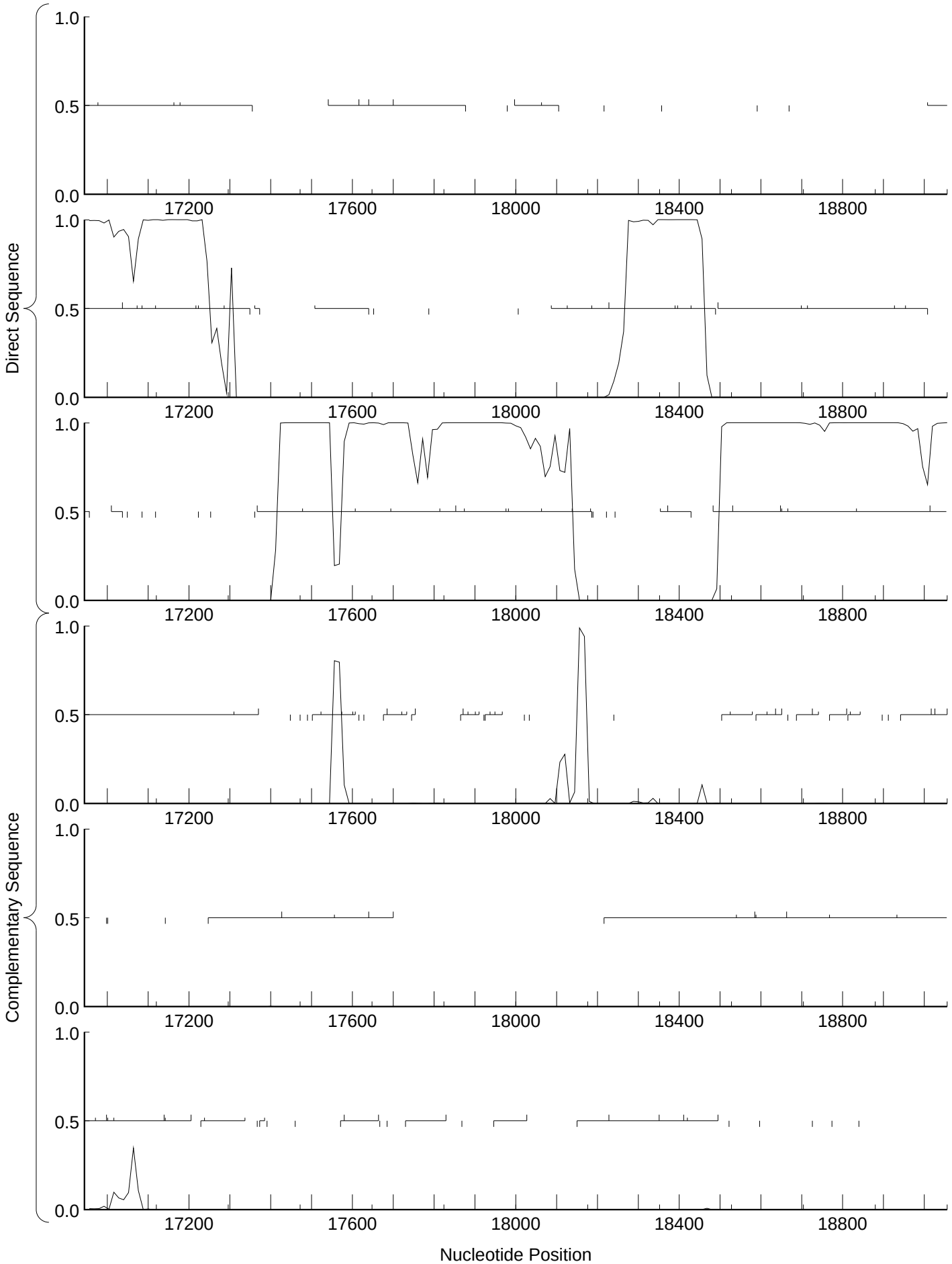
Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 8/20



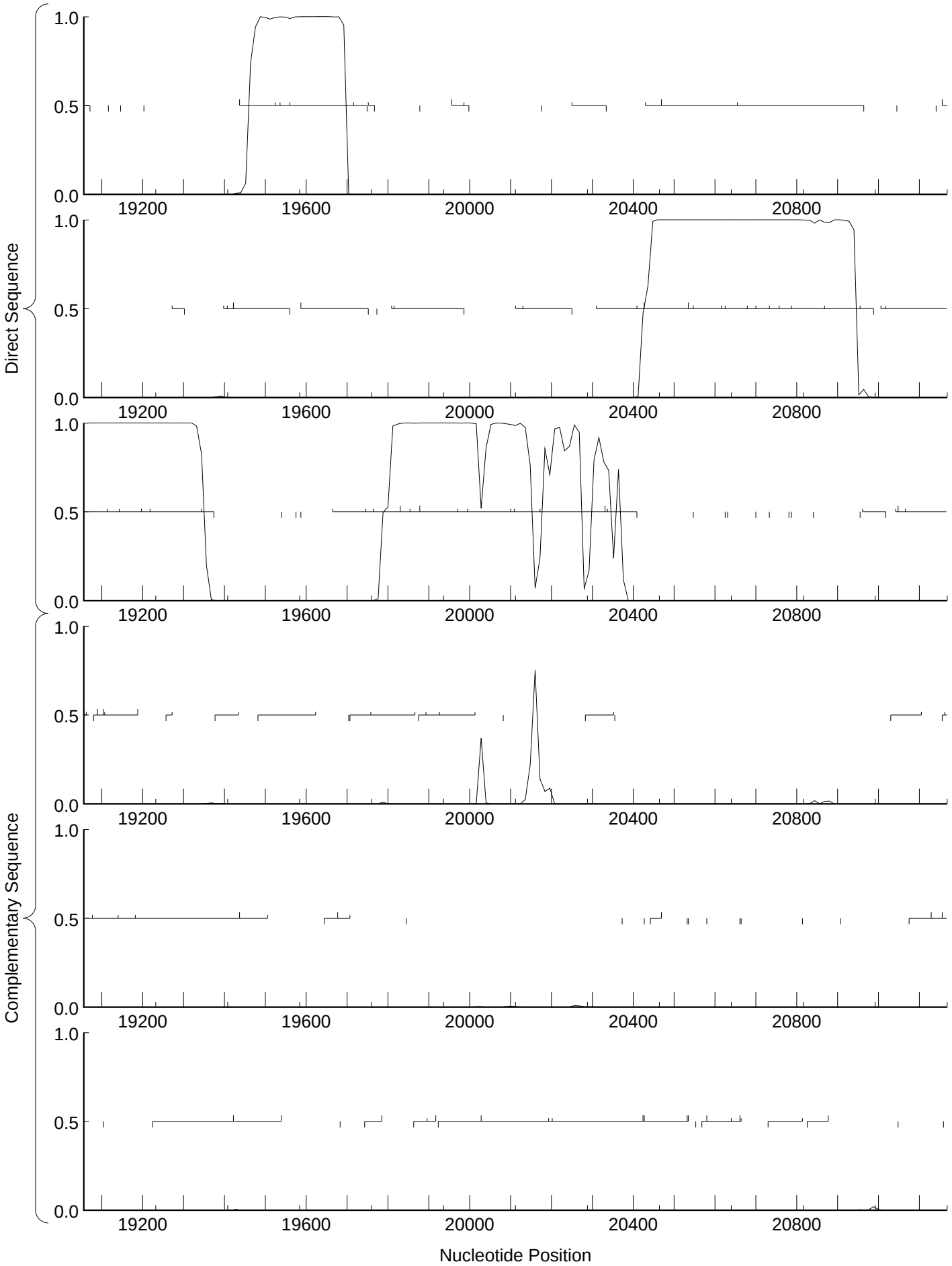
Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 9/20



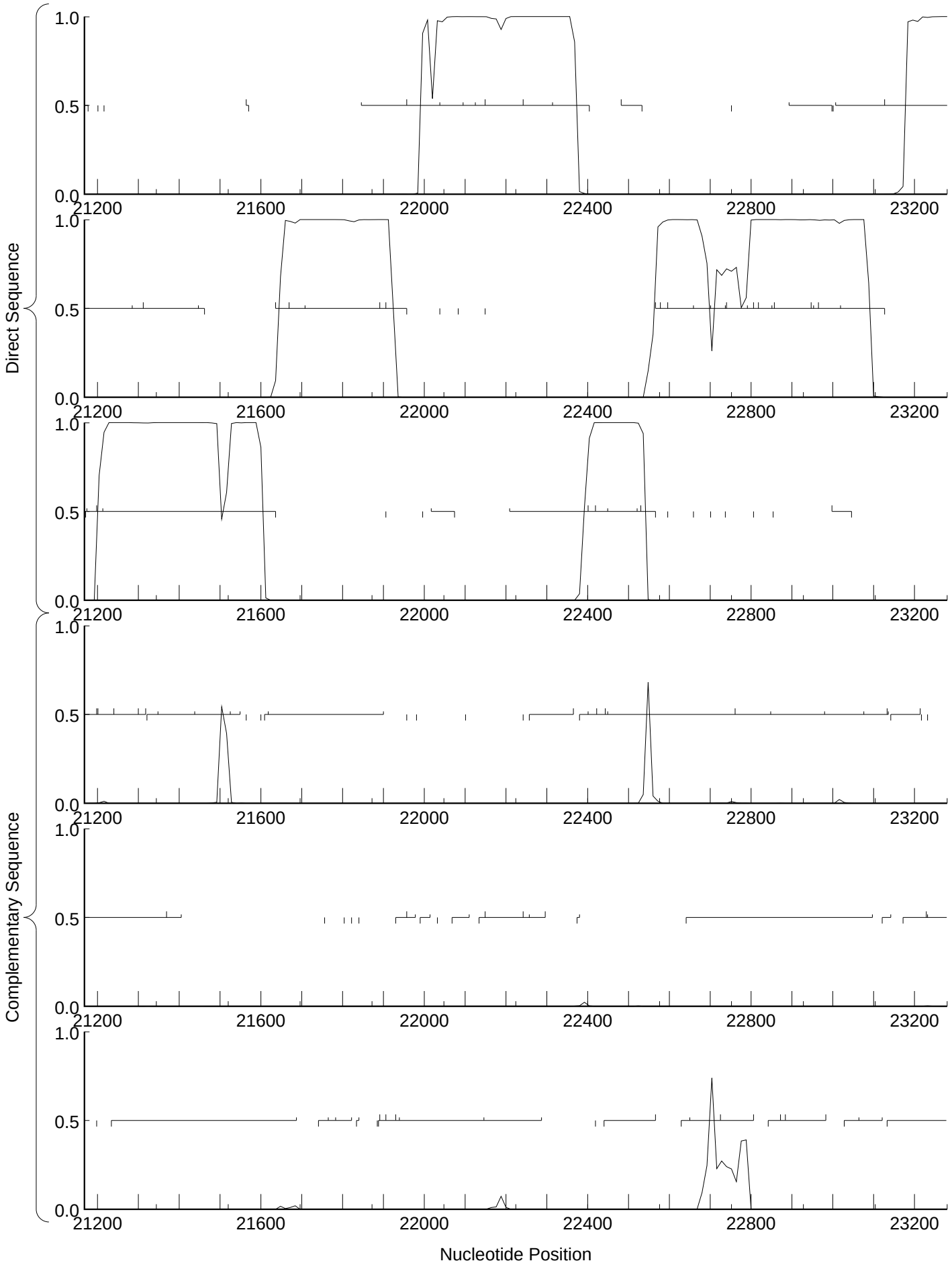
Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 10/20



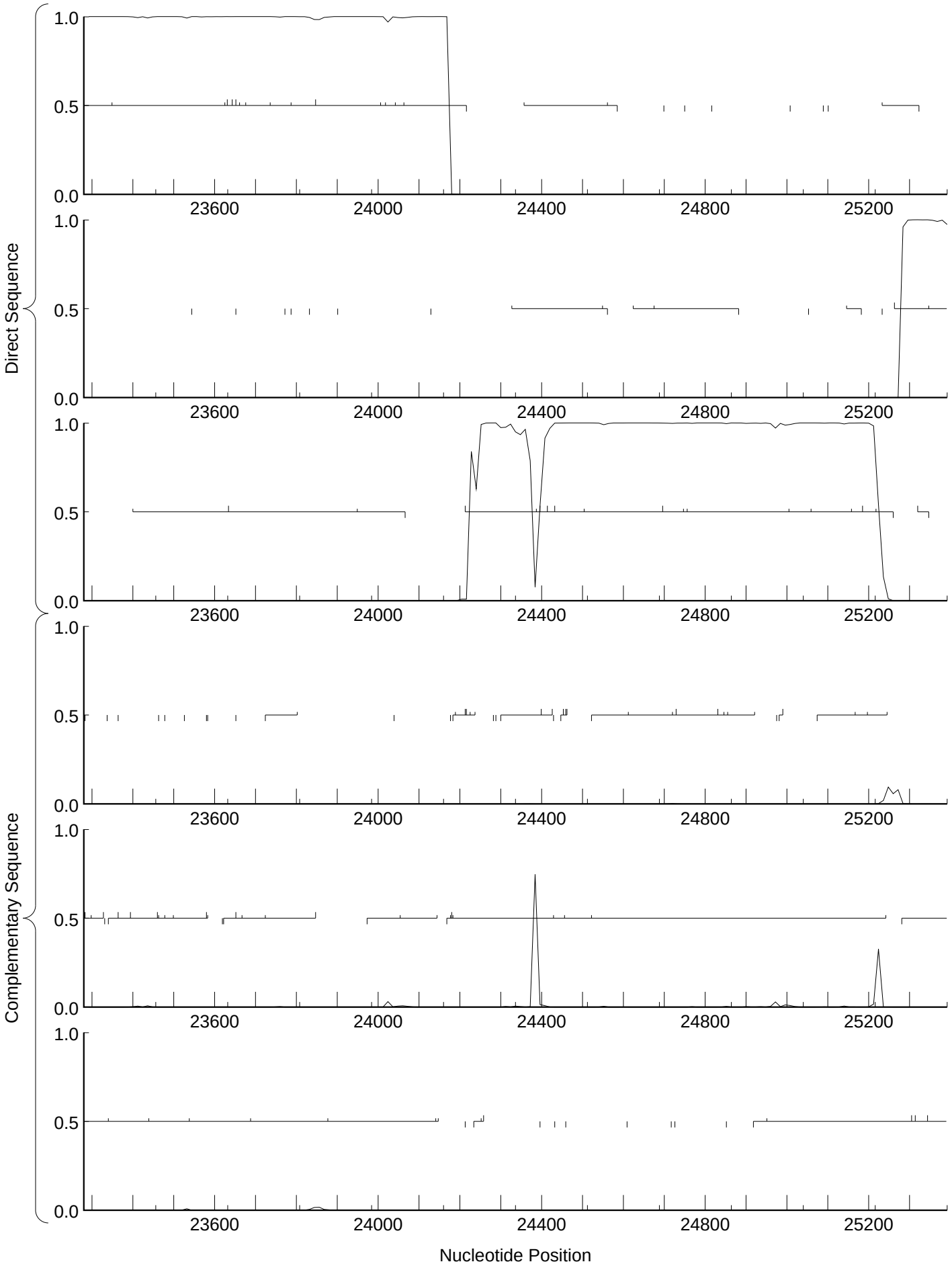
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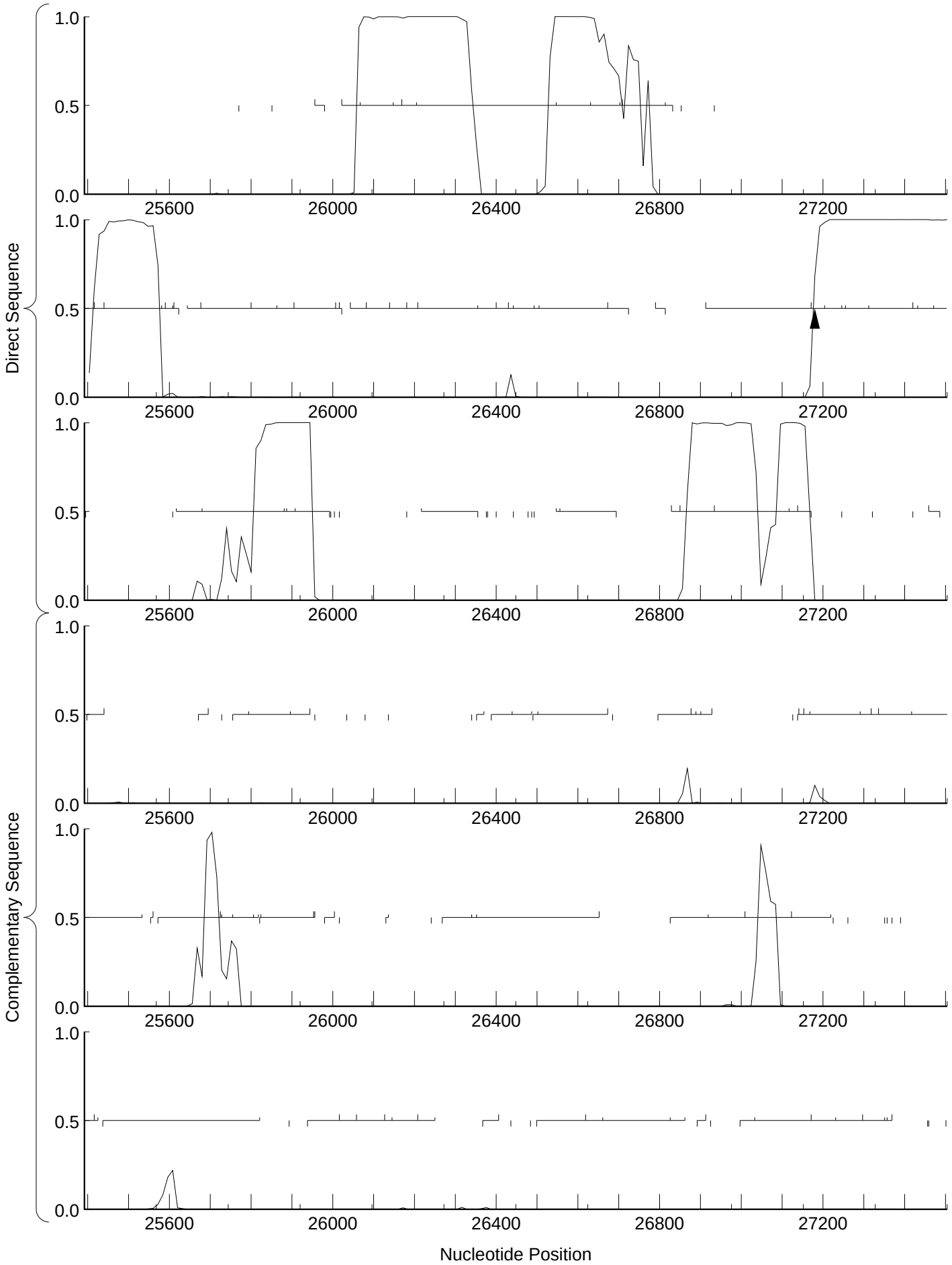
Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 12/20



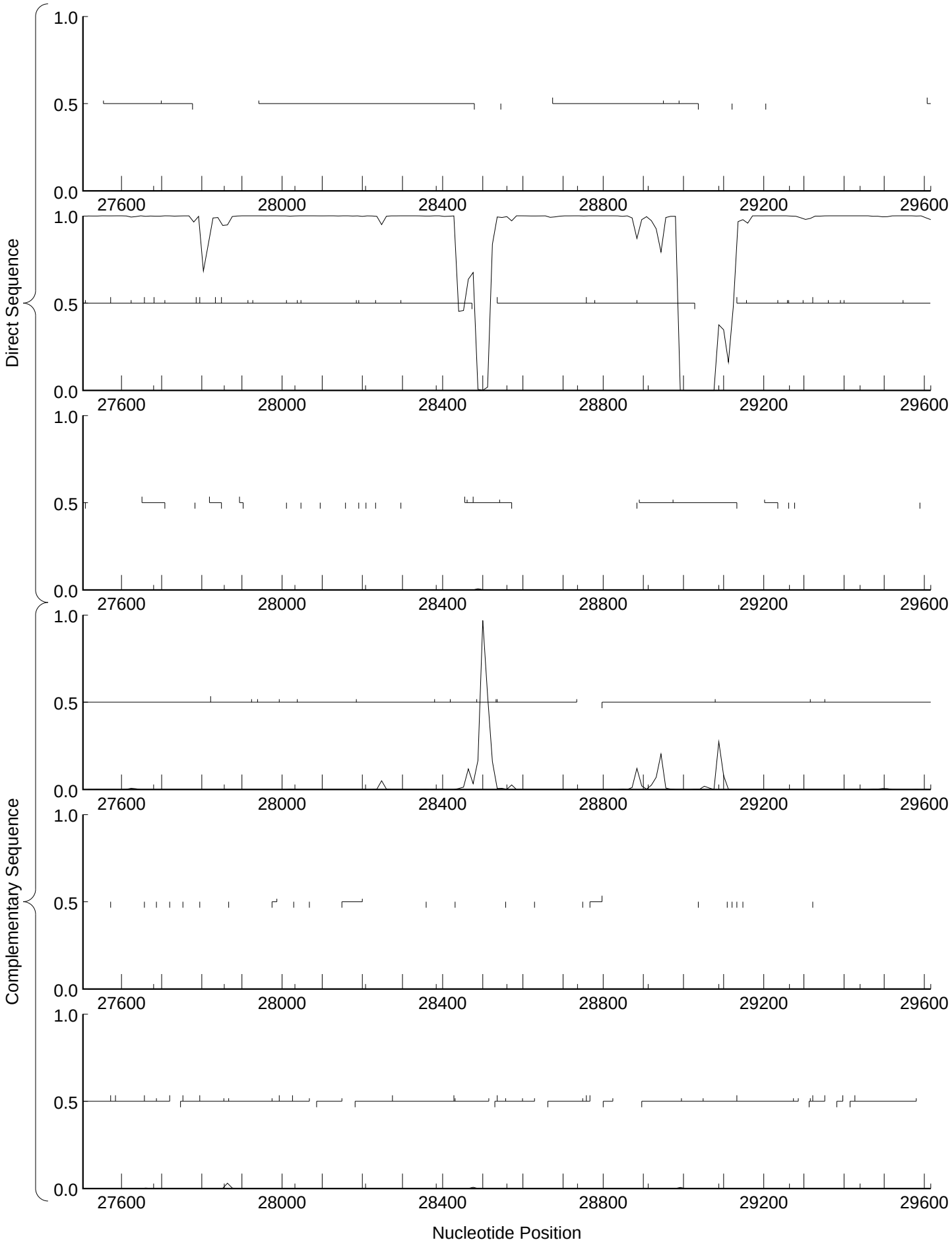
Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 13/20



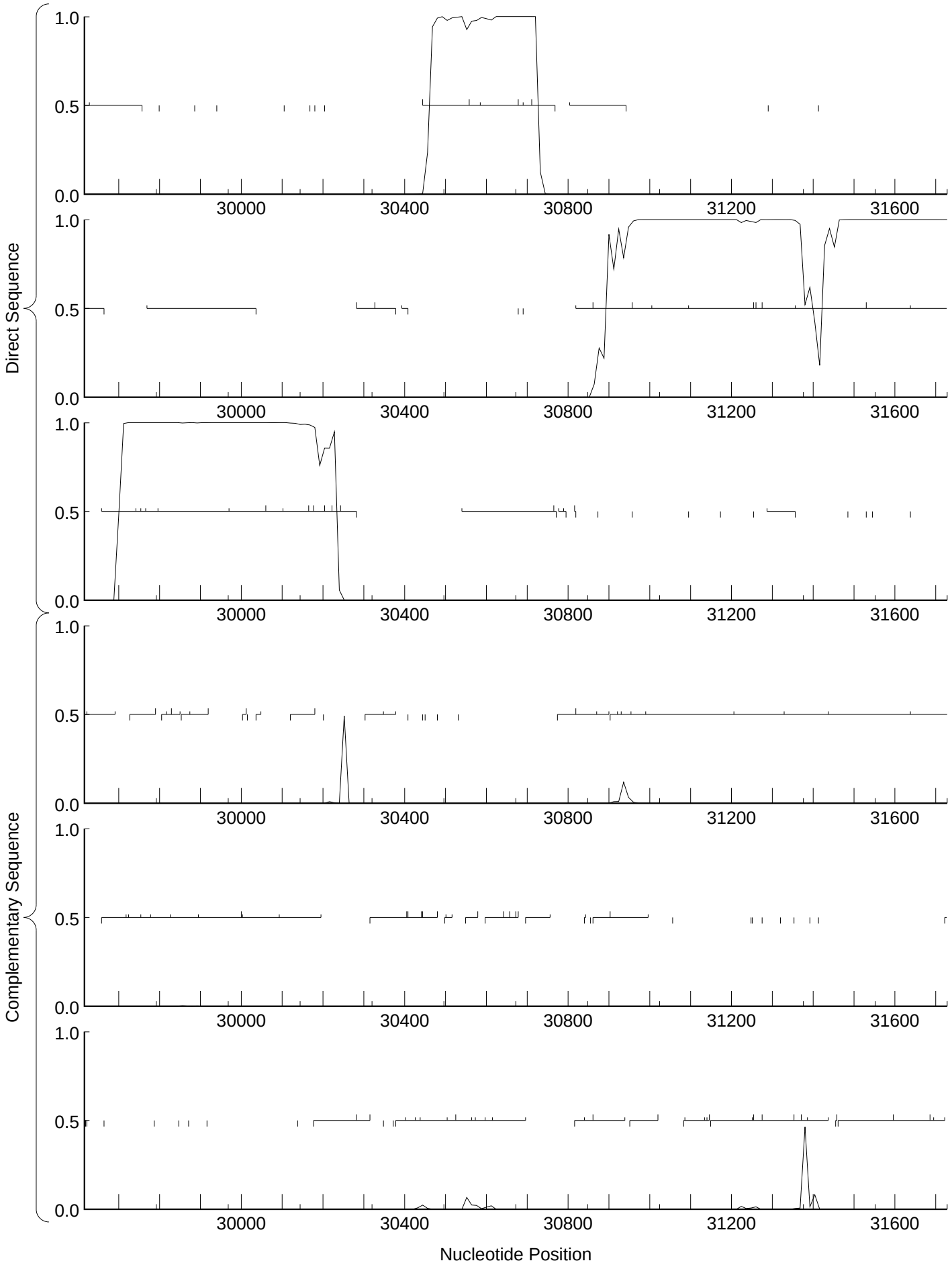
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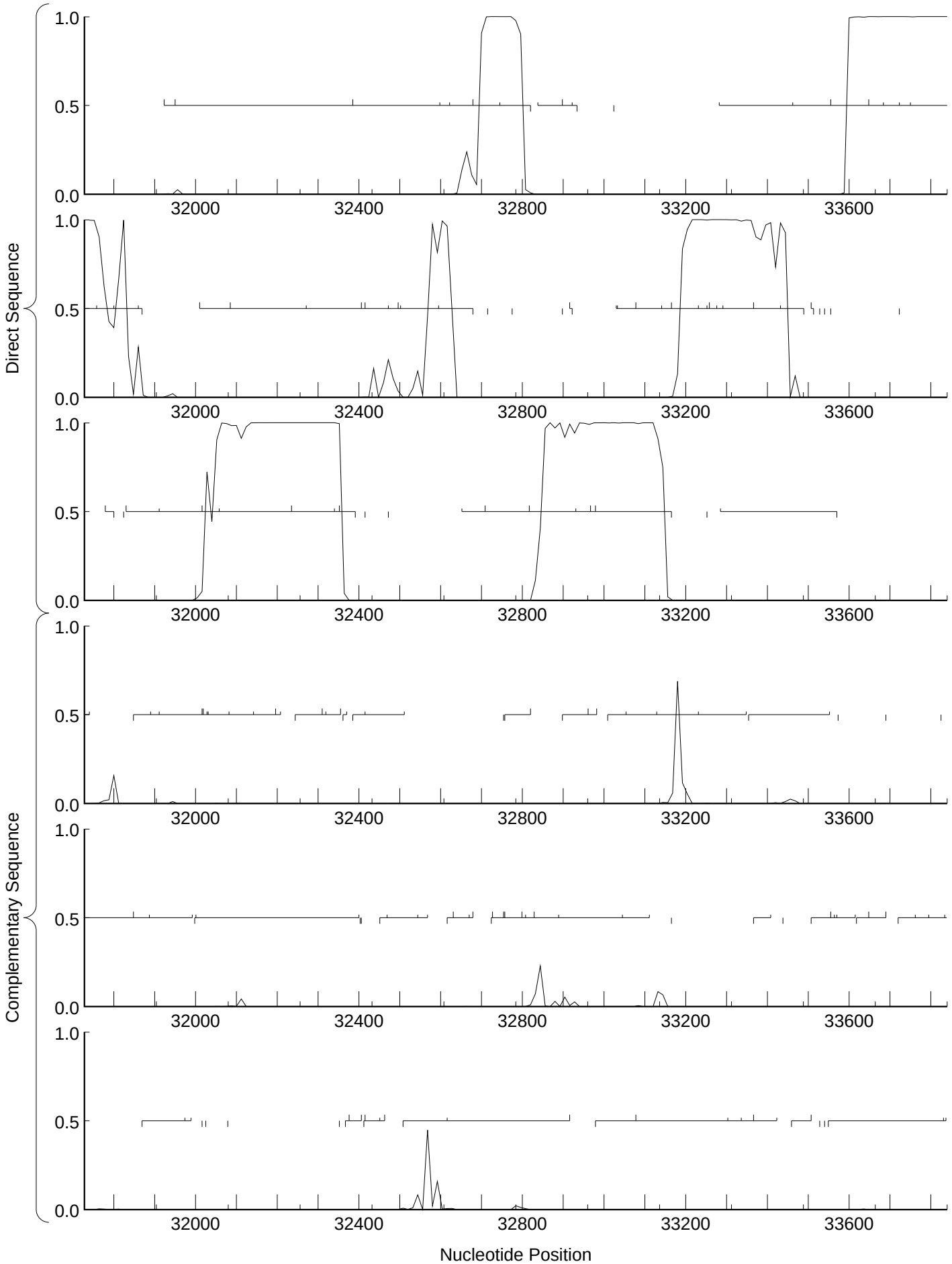
Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 15/20



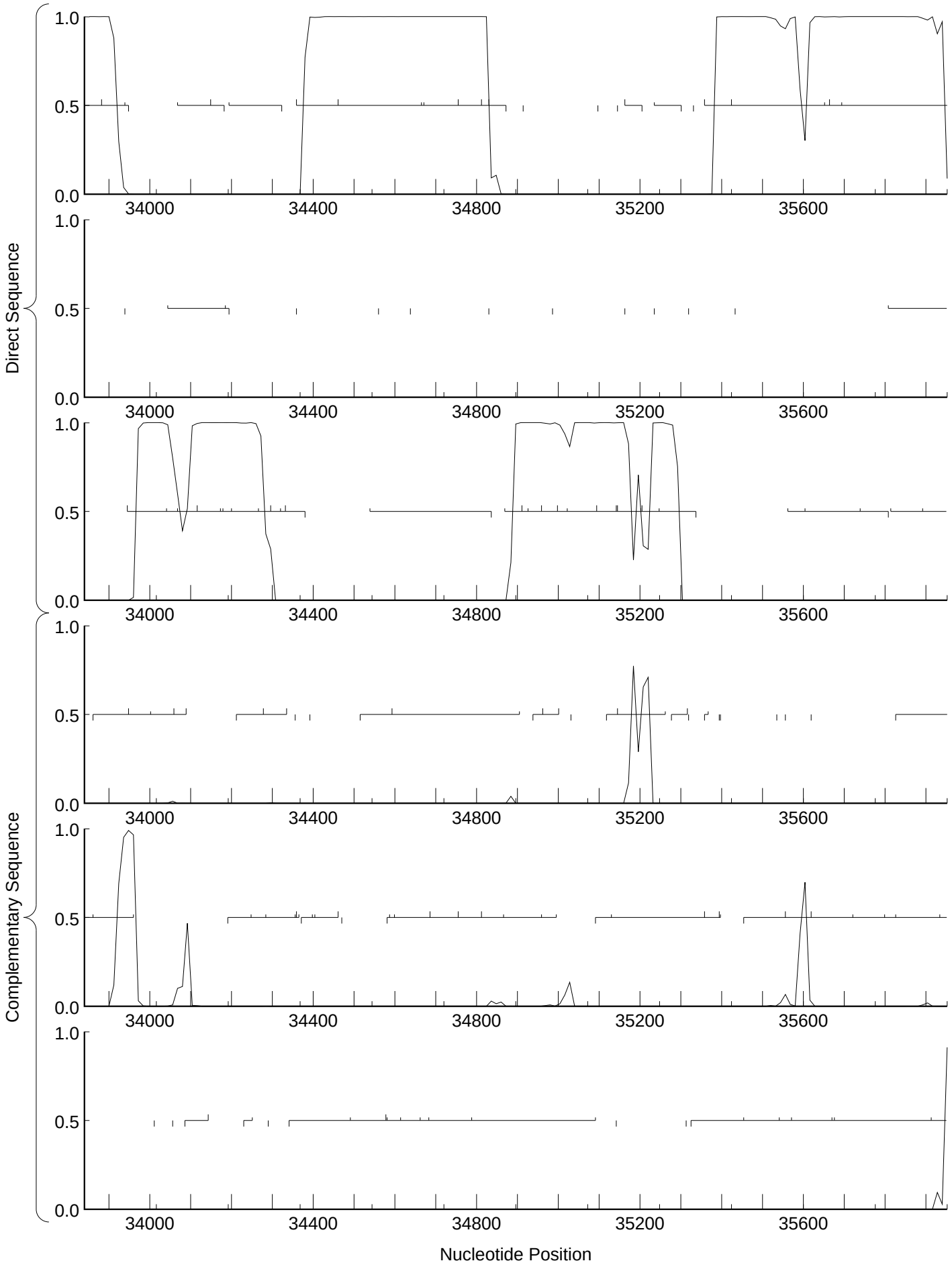
Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 16/20



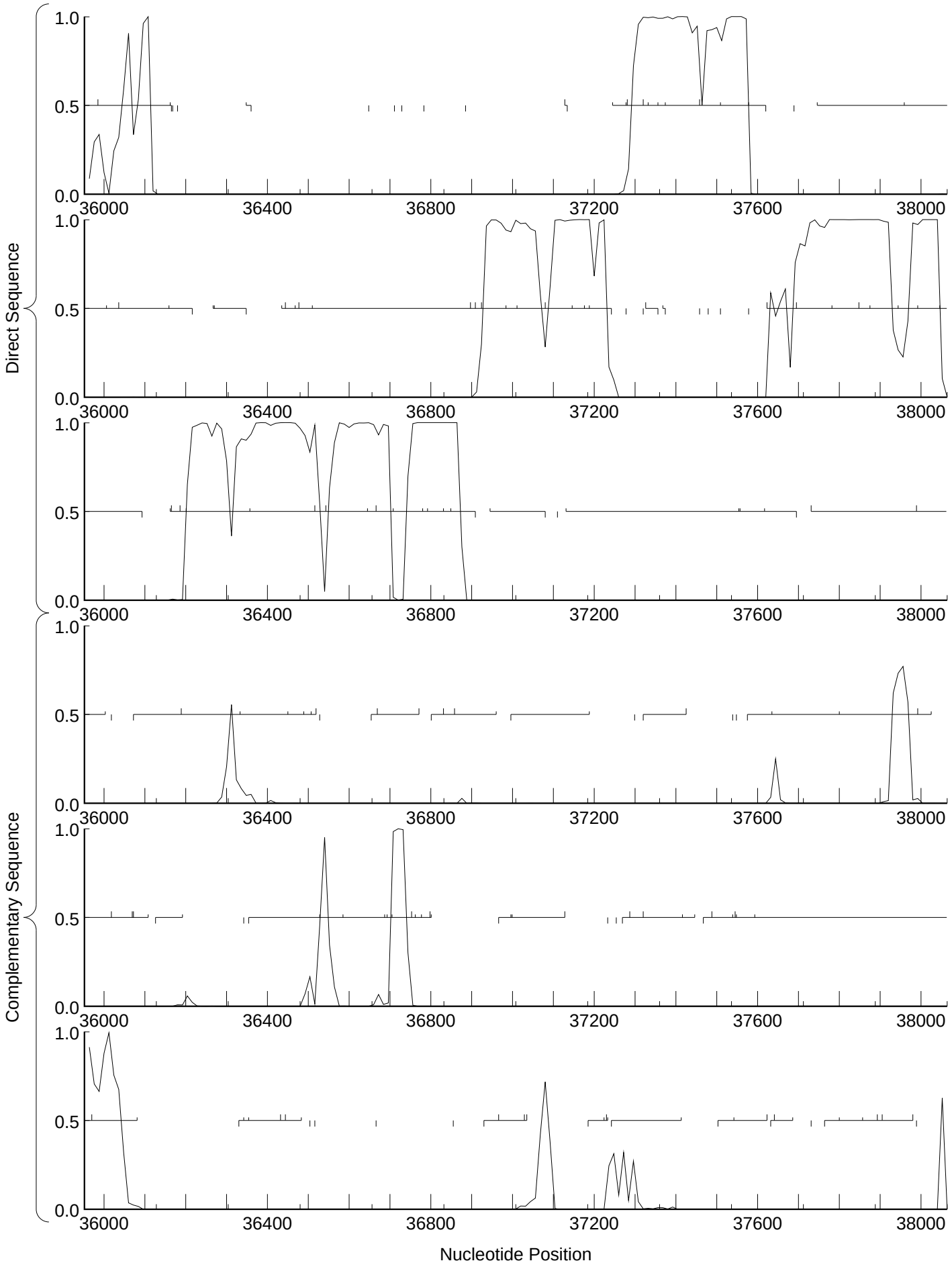
Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 17/20



Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 18/20



Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 19/20



Microbacterium phage Appa complete sequence, 38684 bp, circularly permuted, Cluster GA, Order 4, Window 96, Step 12, 20/20

