GeneMark

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

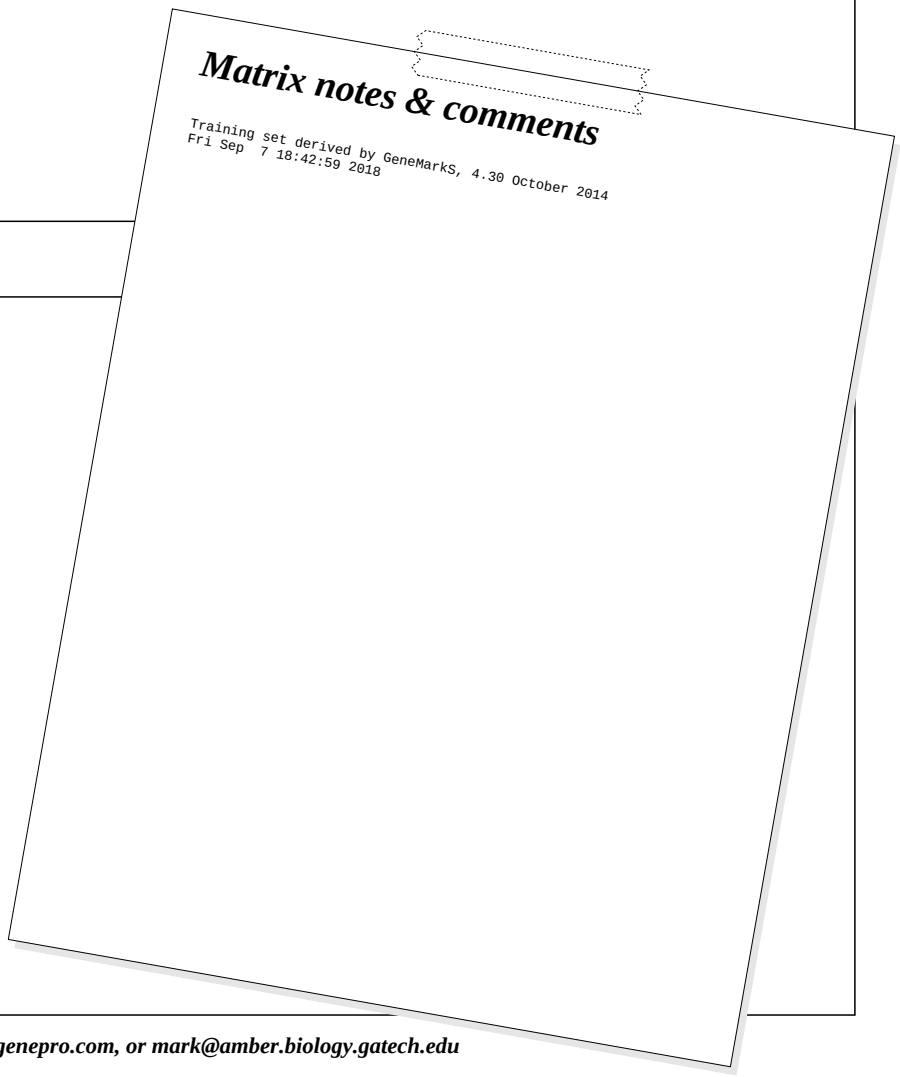
PROGRAM INFORMATION

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Analysis Date    :  3/29/24 at 3:03:09
Pages            :  28
Sequence Length  :  56618 bp
GC Content       :   63.75%

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

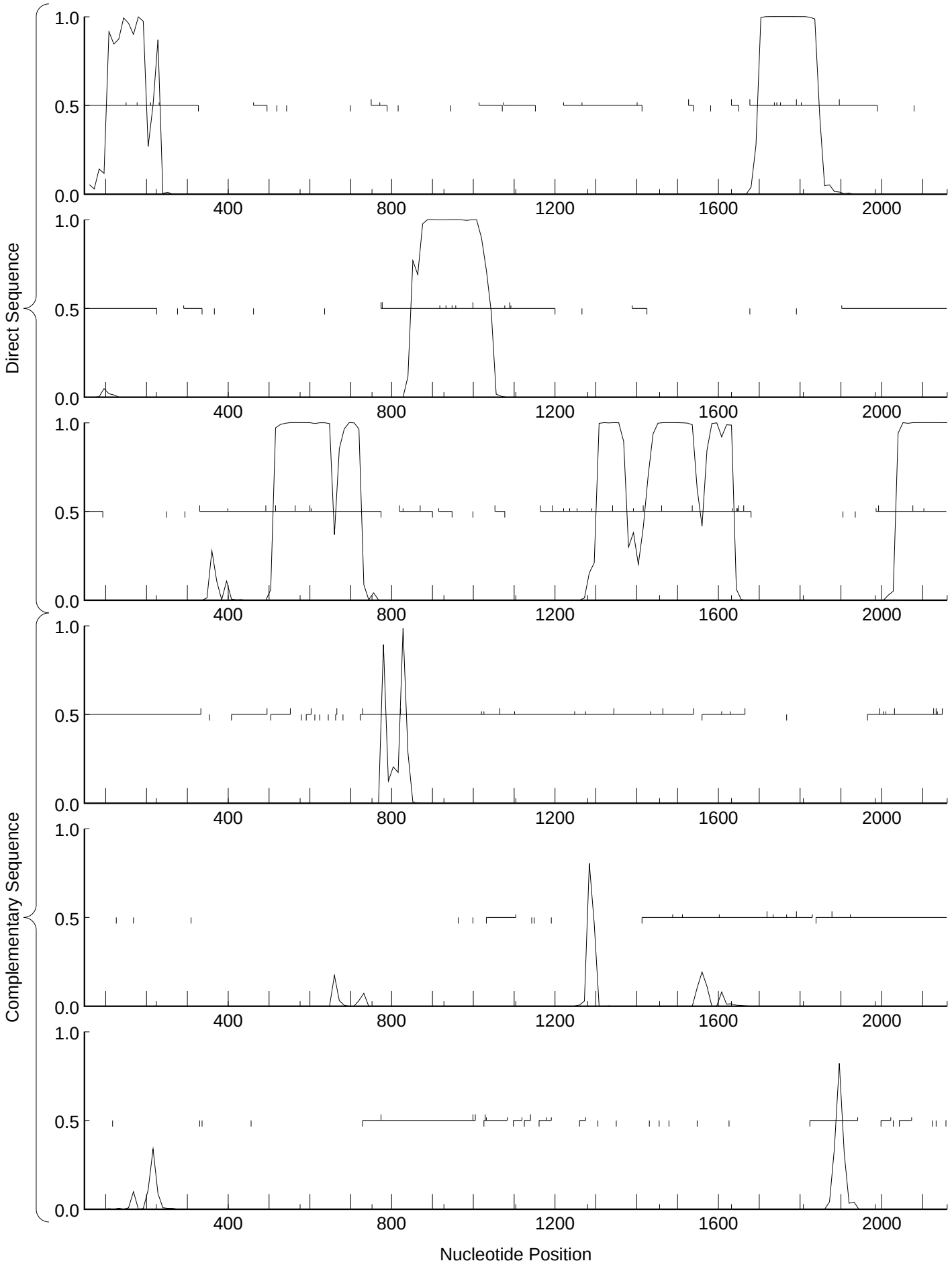
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Matrix     :  -
Author     :  -

Order      :  4
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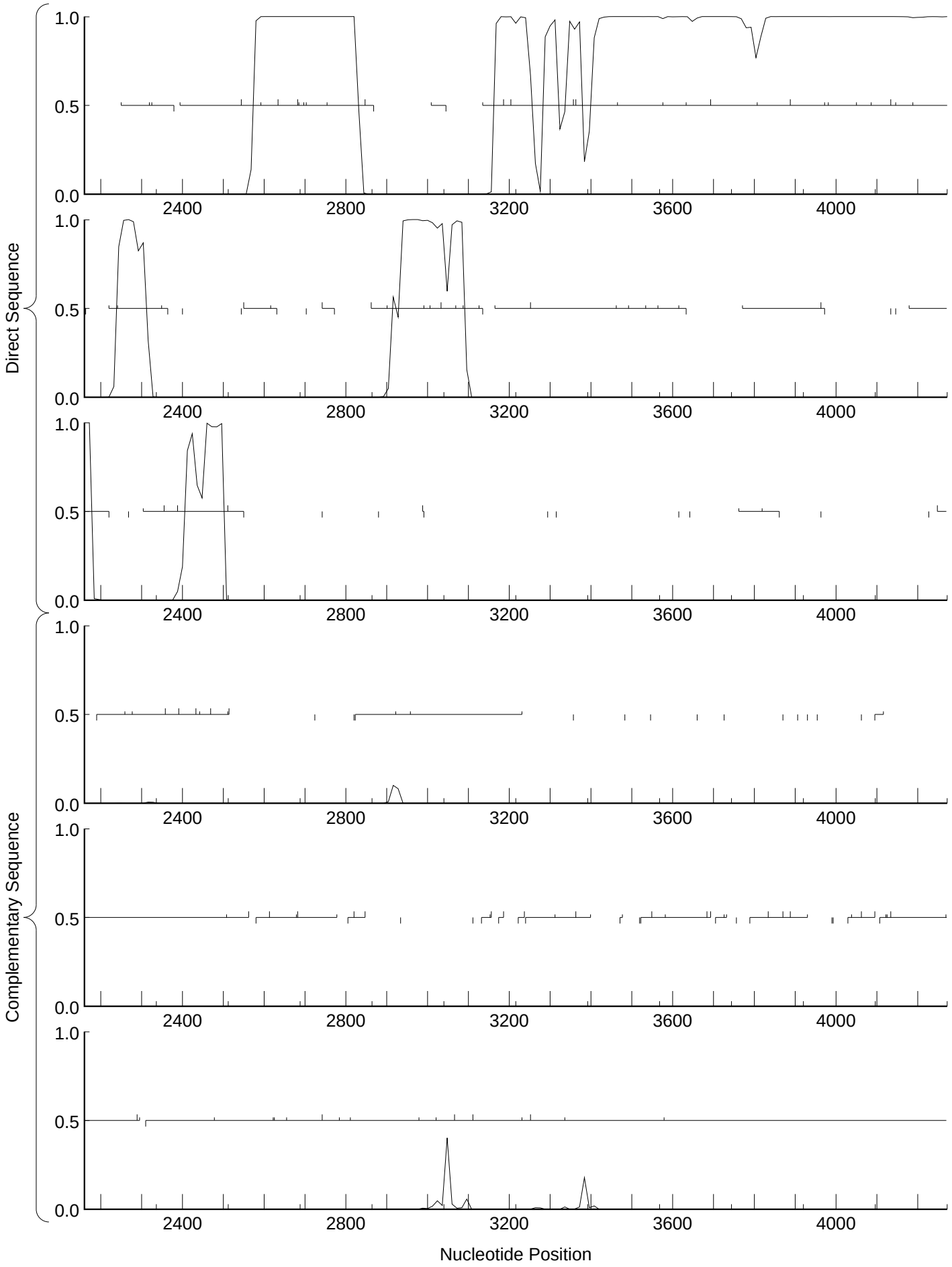

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

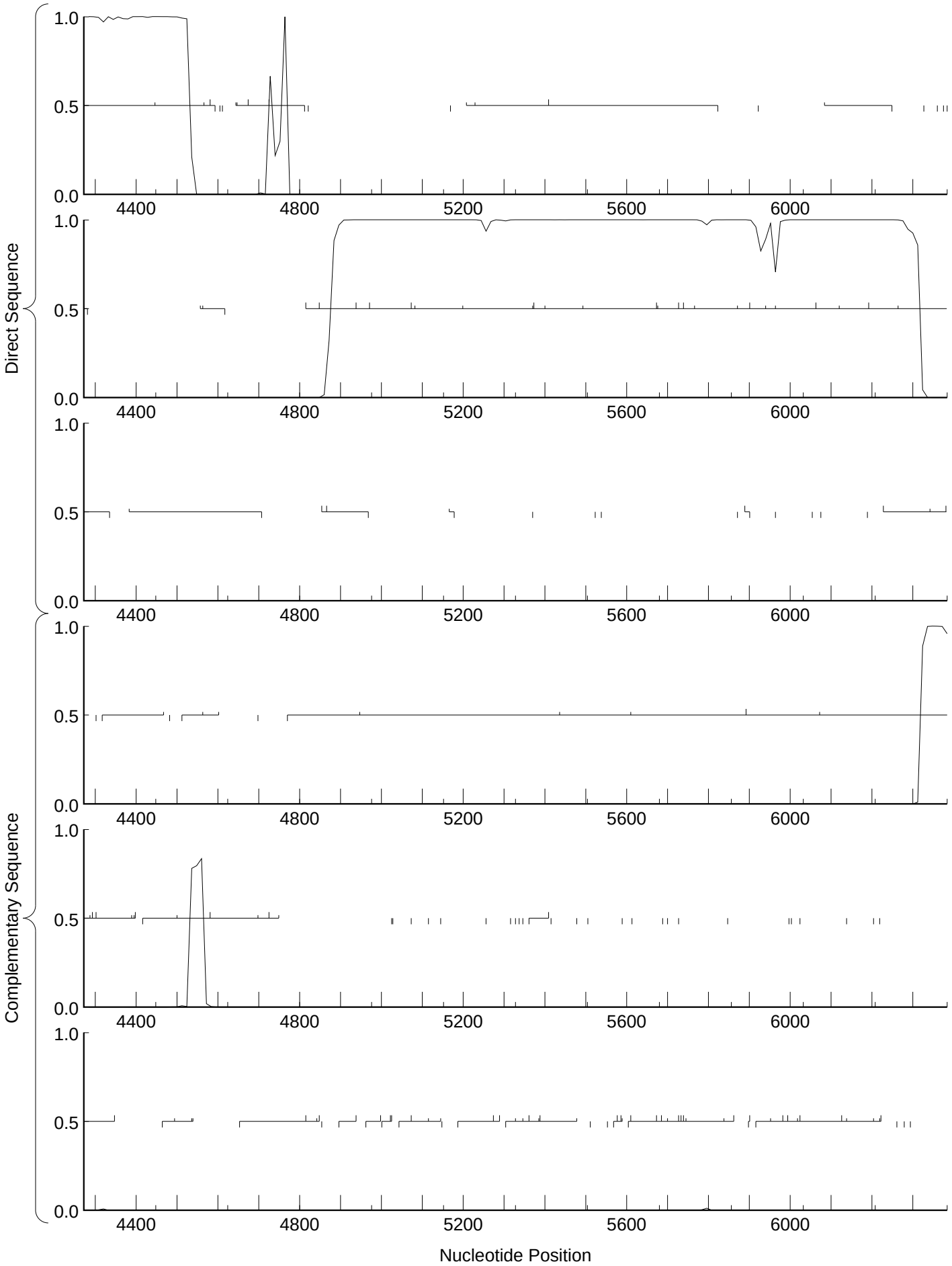
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 2/28



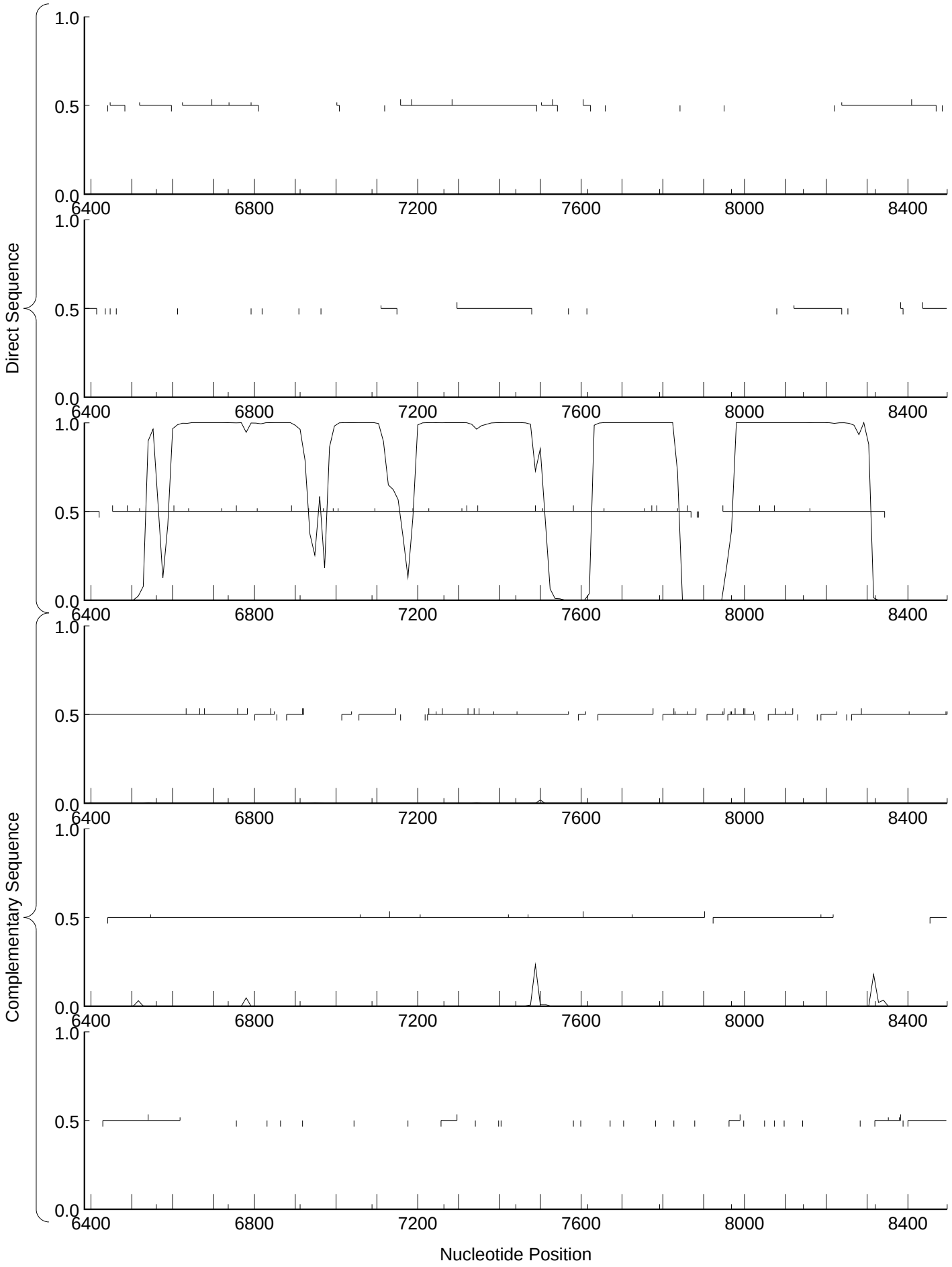
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 3/28



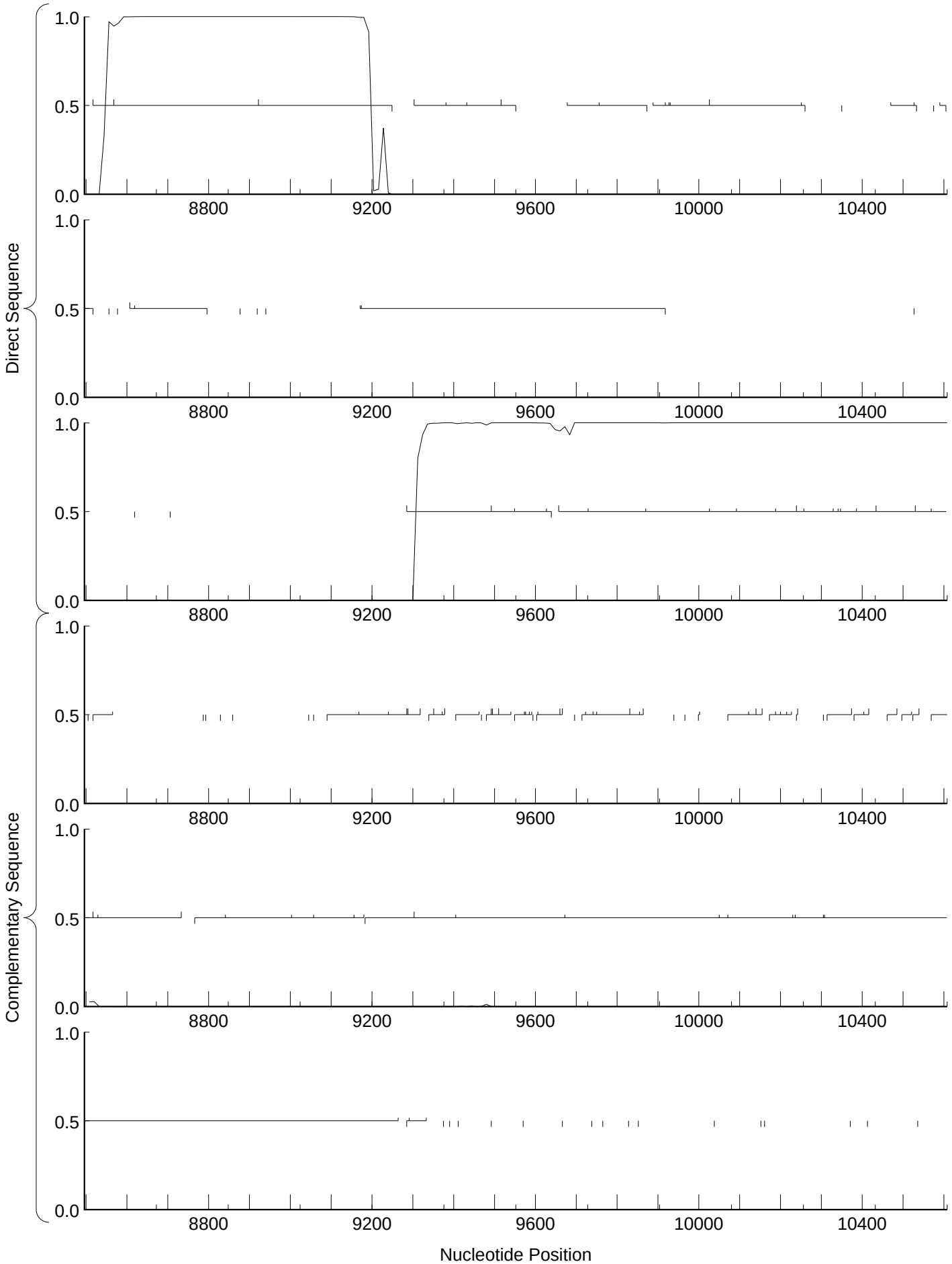
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 4/28



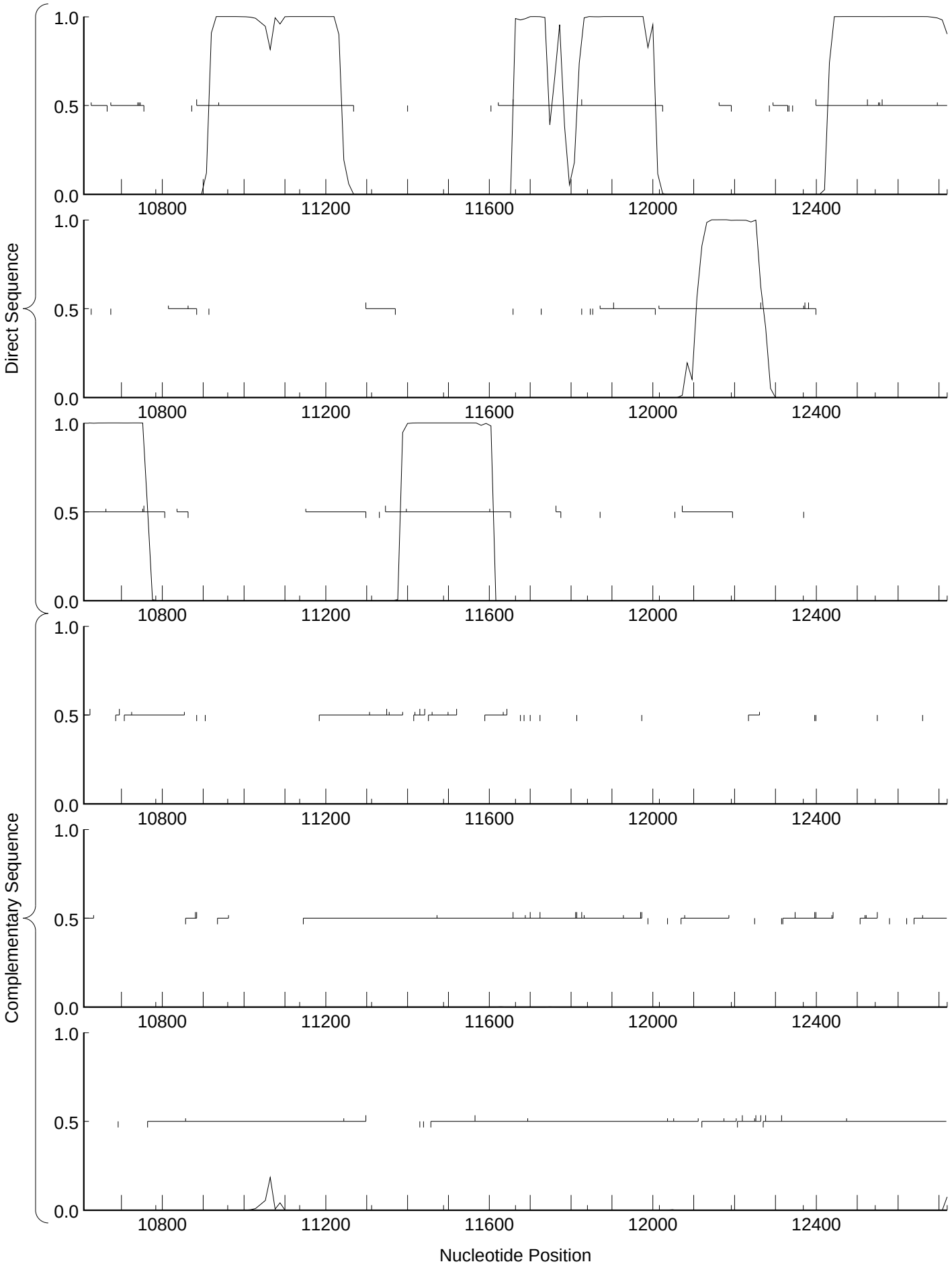
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 5/28



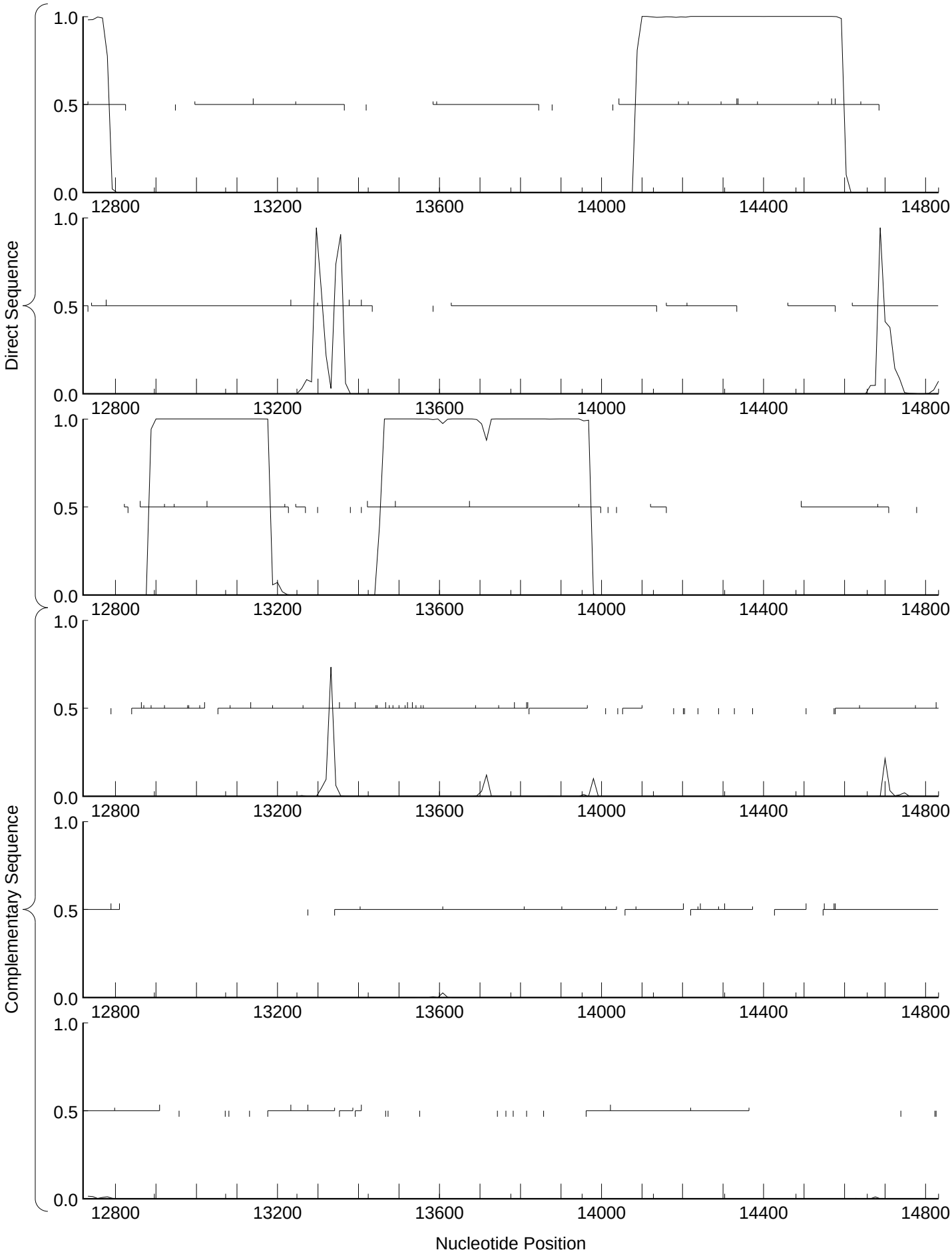
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 6/28



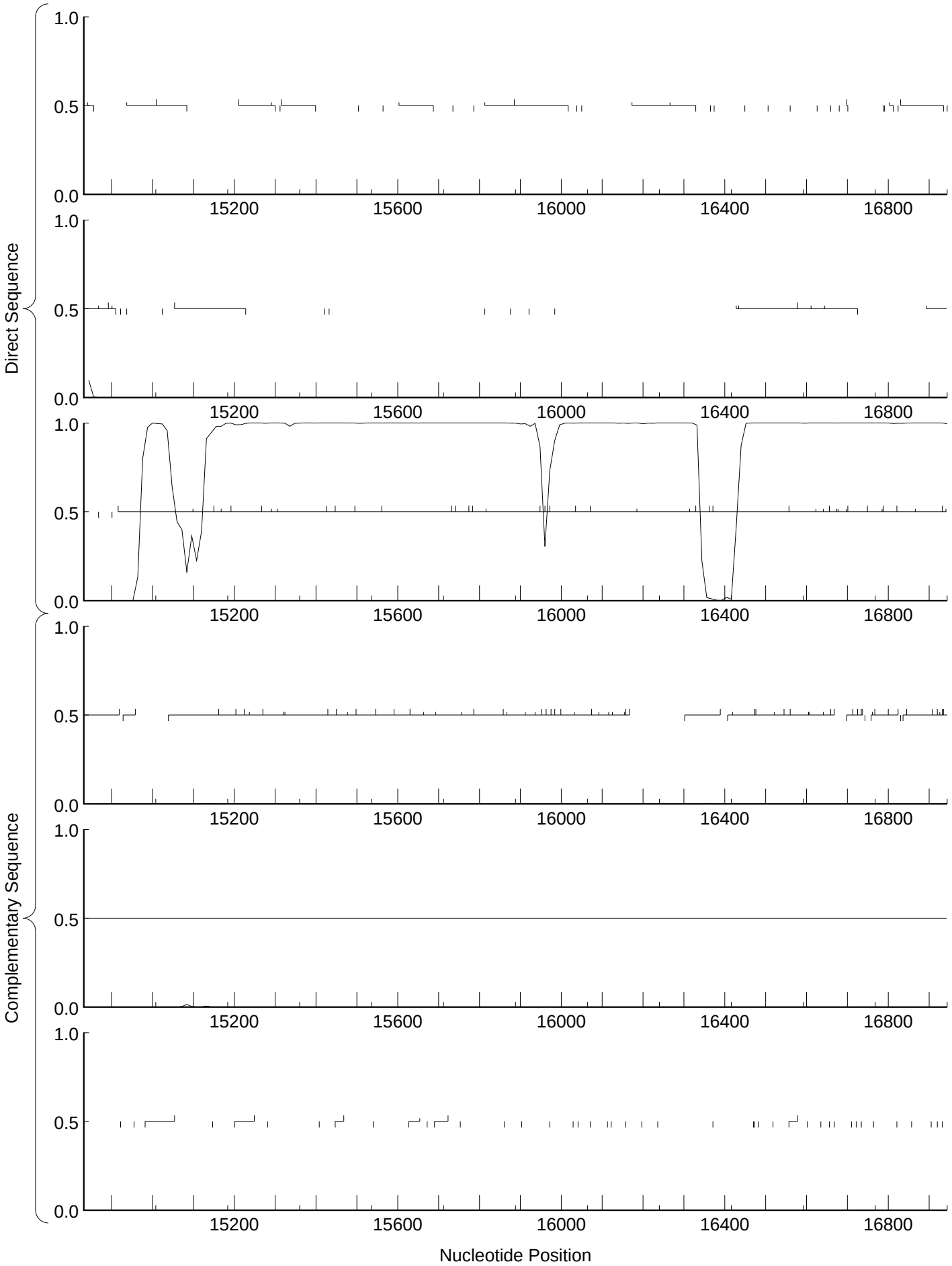
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 7/28



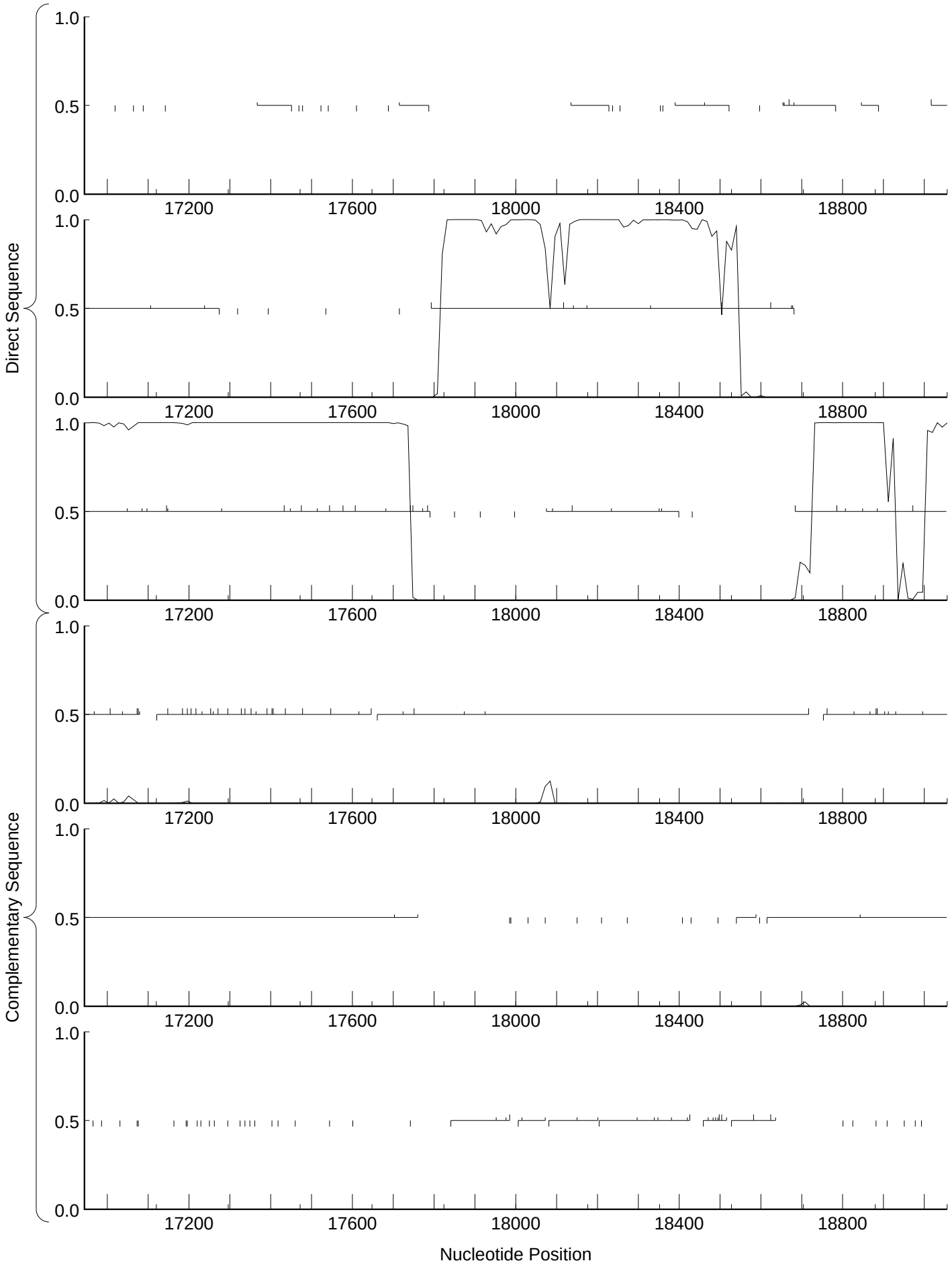
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 8/28



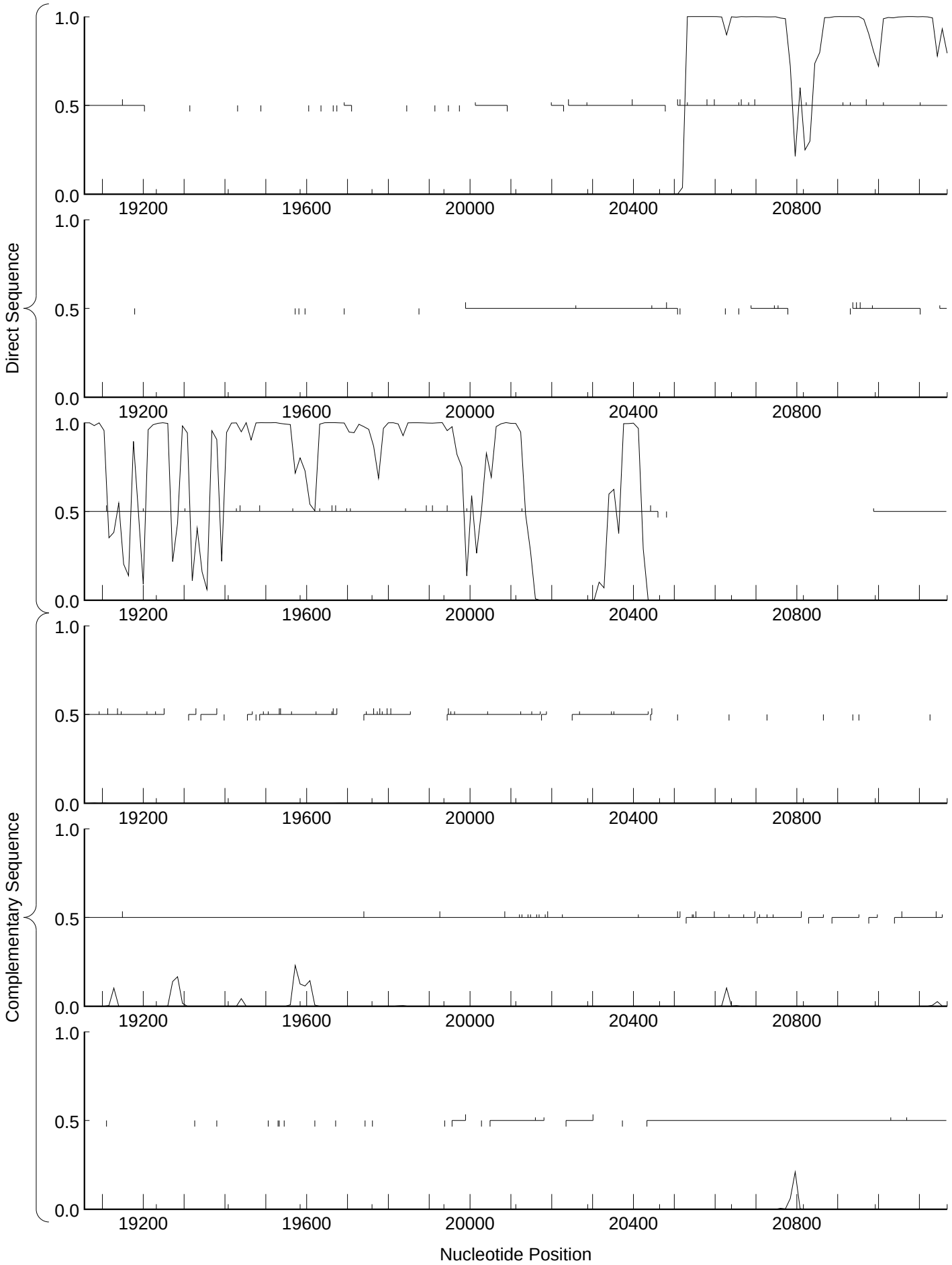
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 9/28



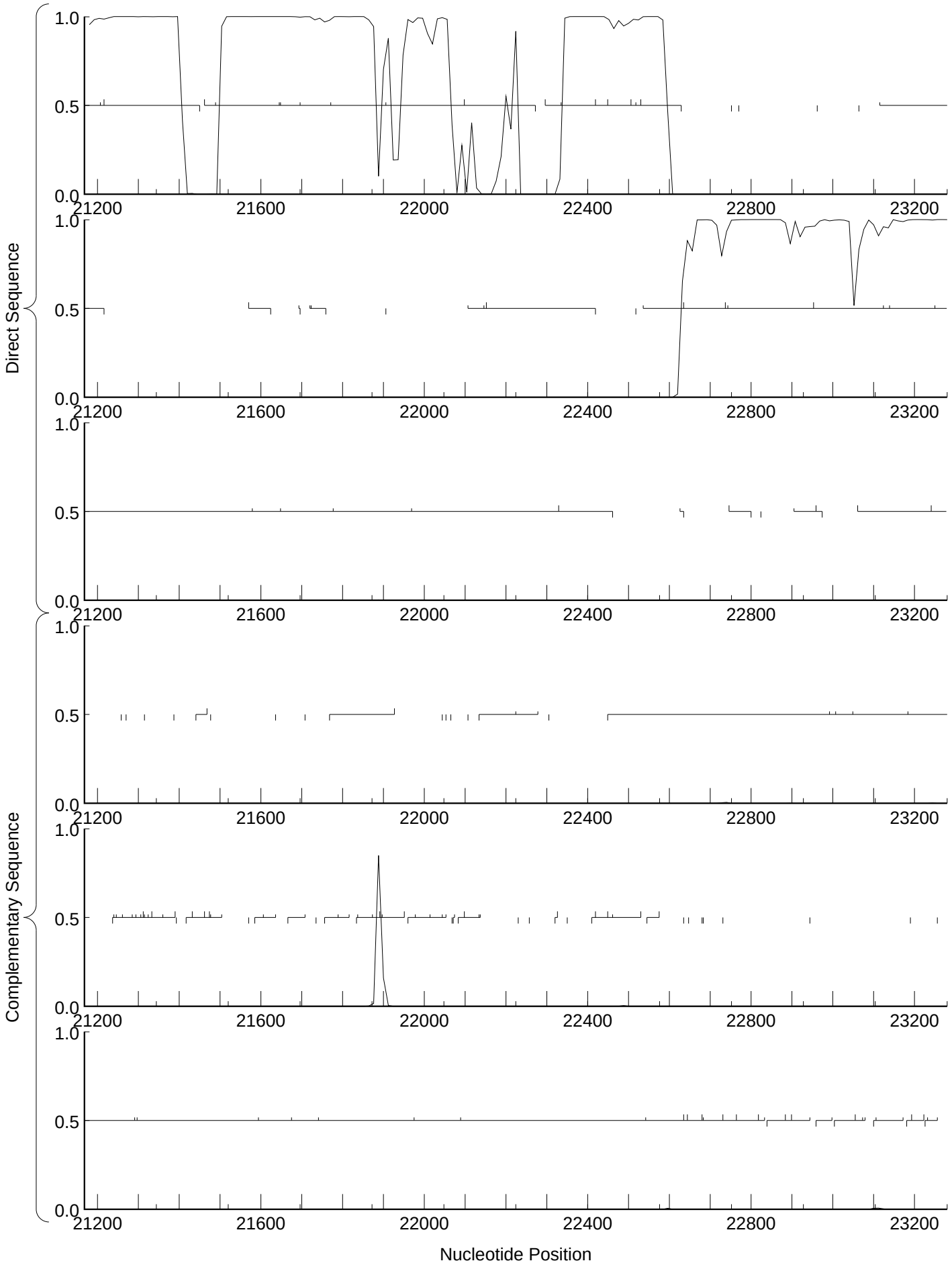
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 10/28



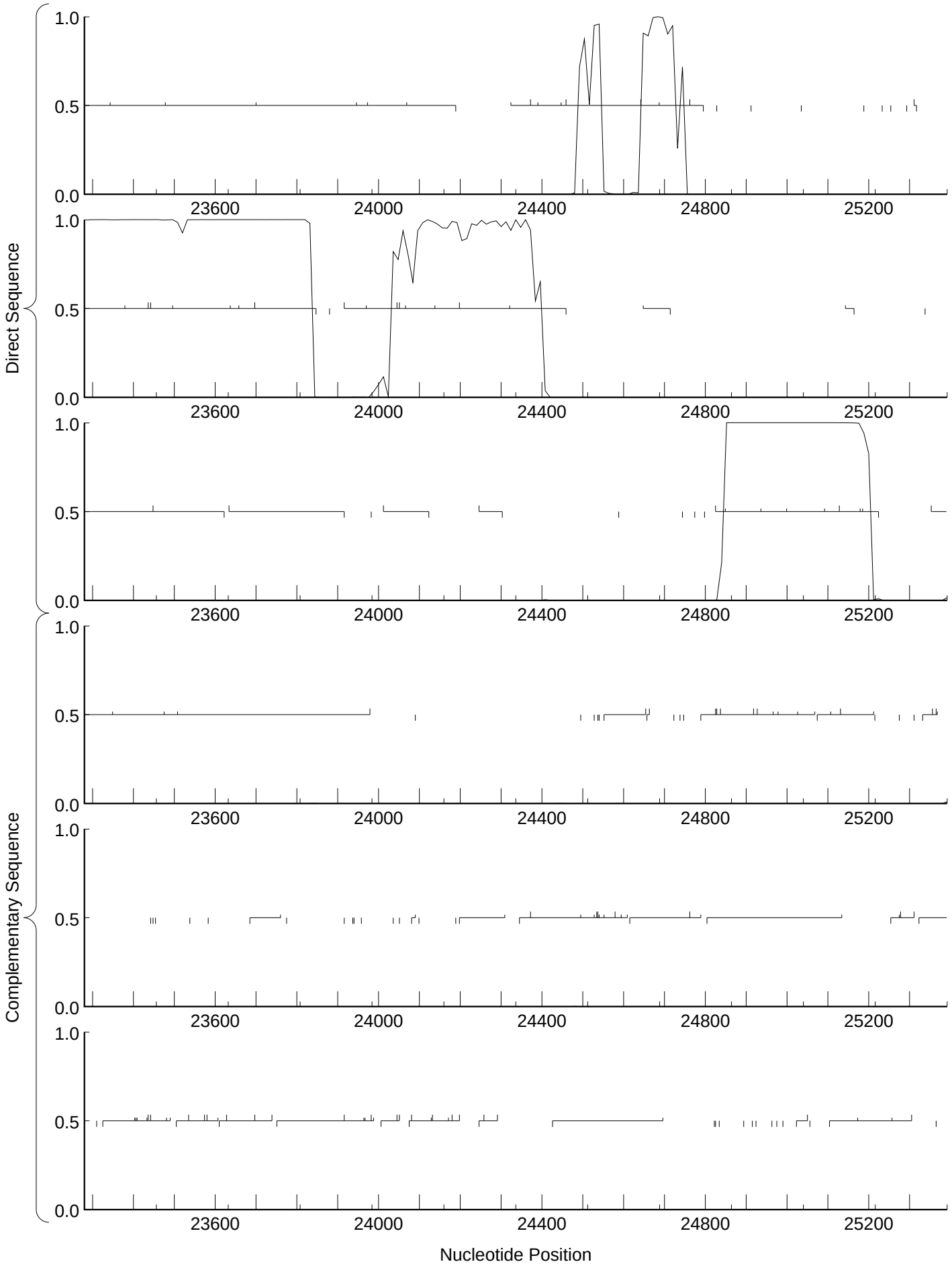
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 11/28



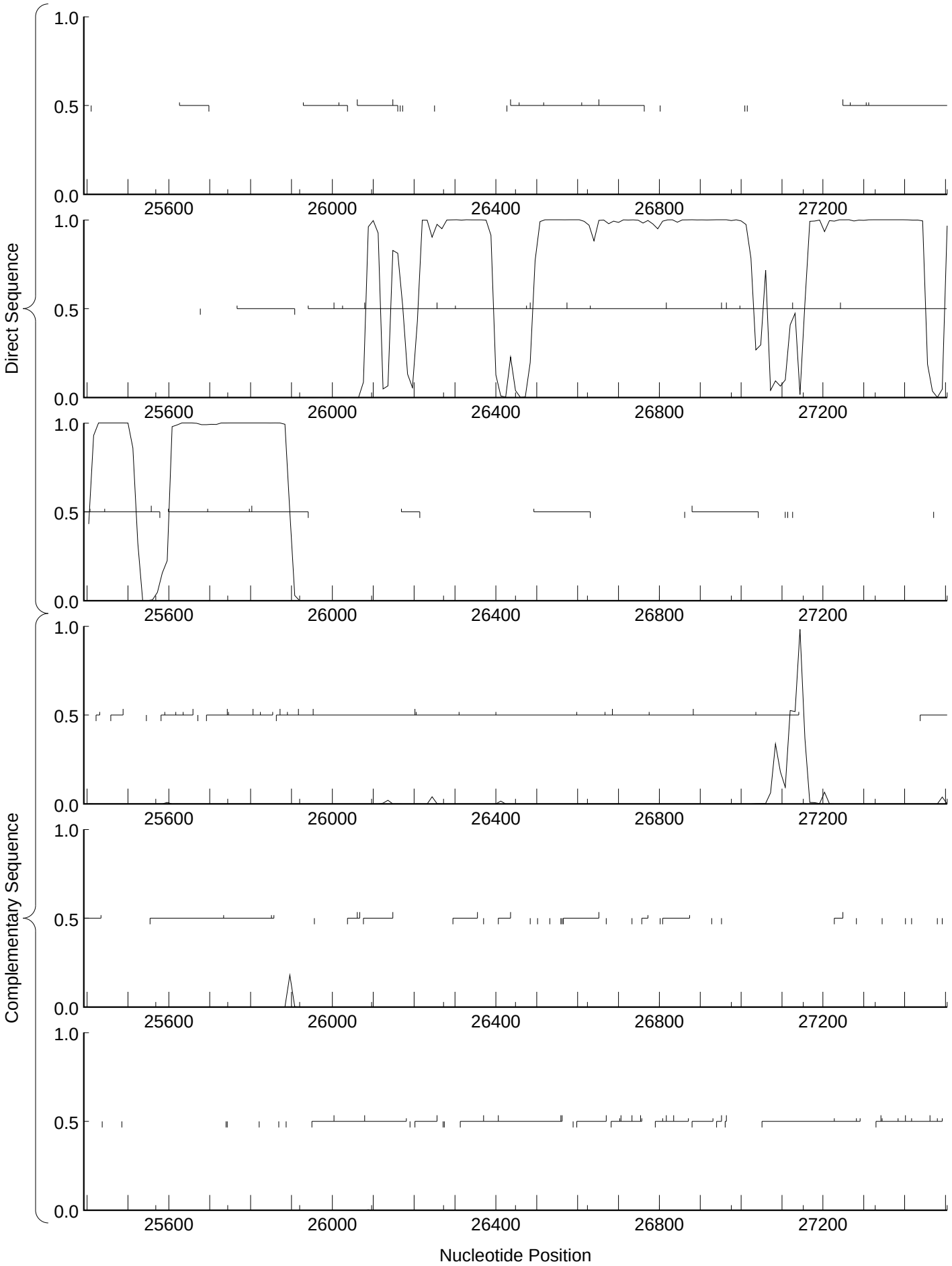
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 12/28



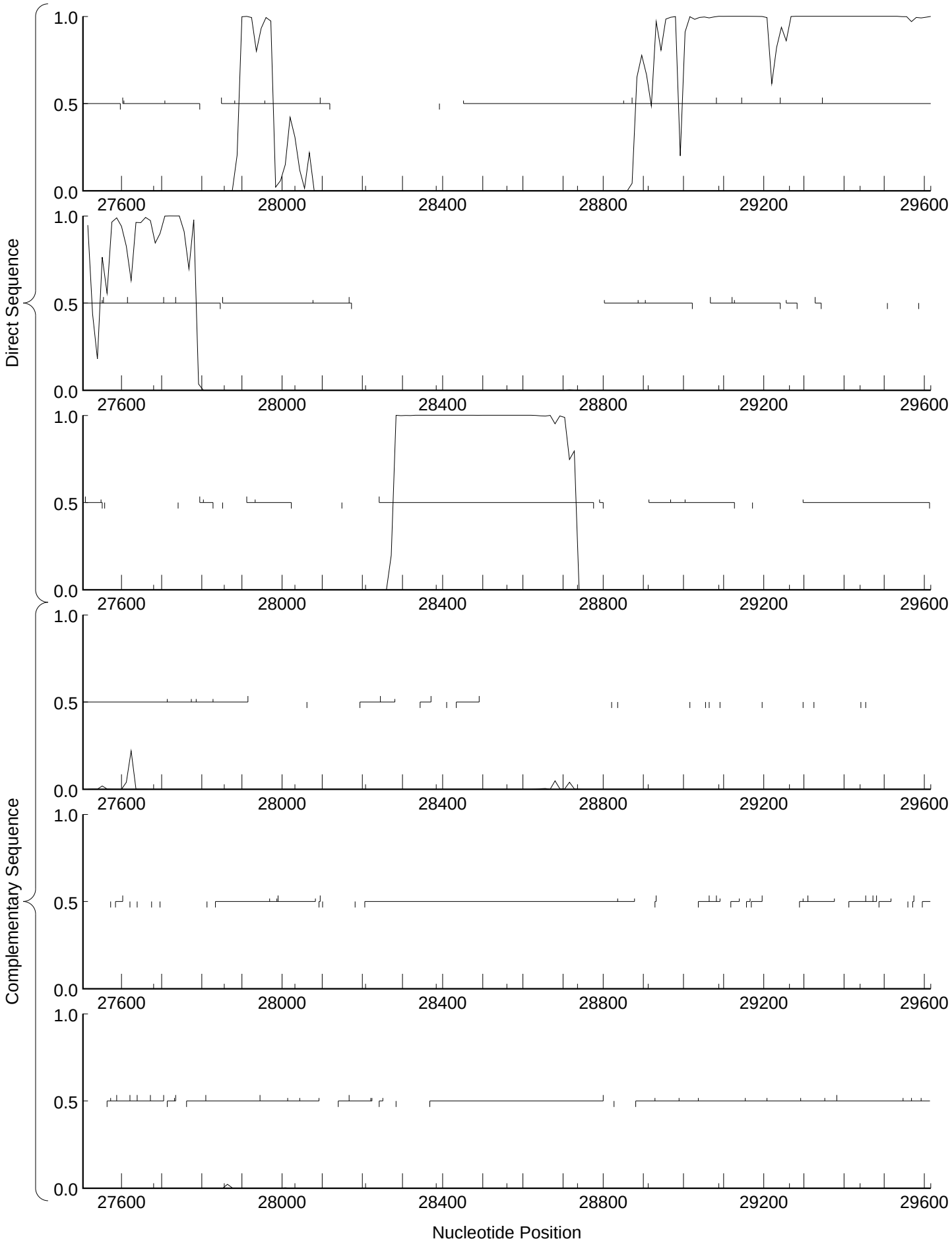
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 13/28



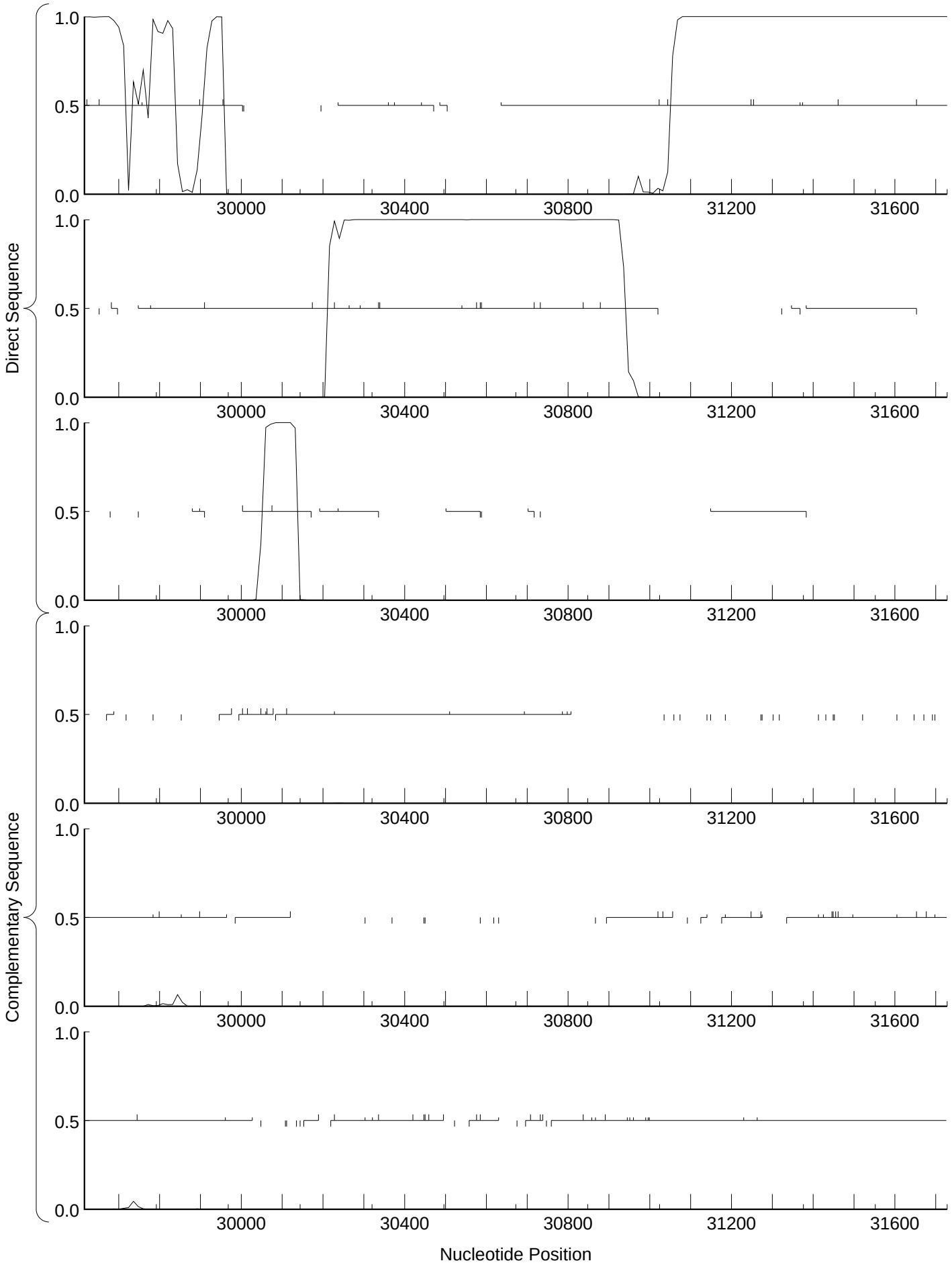
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 14/28



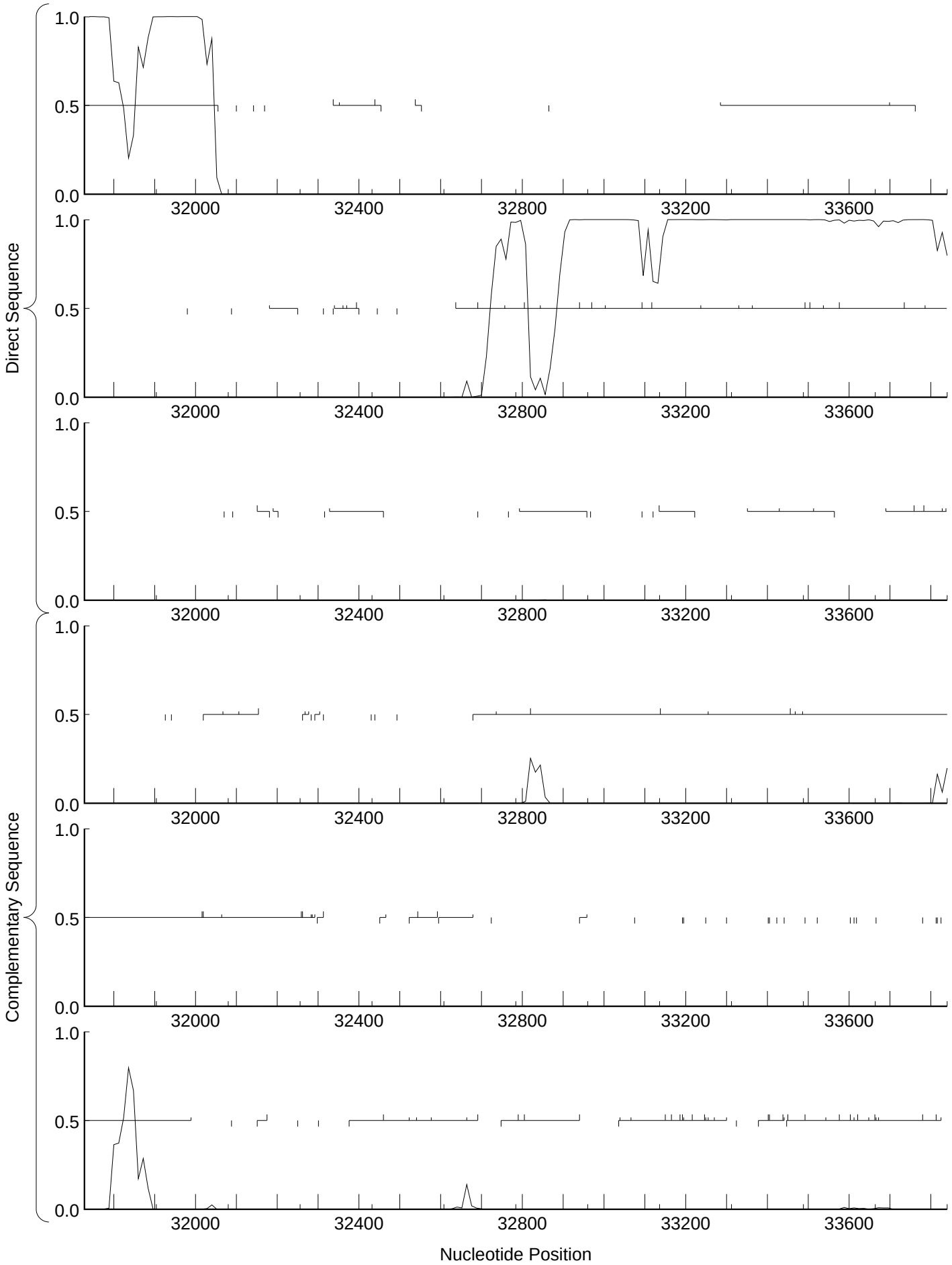
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 15/28



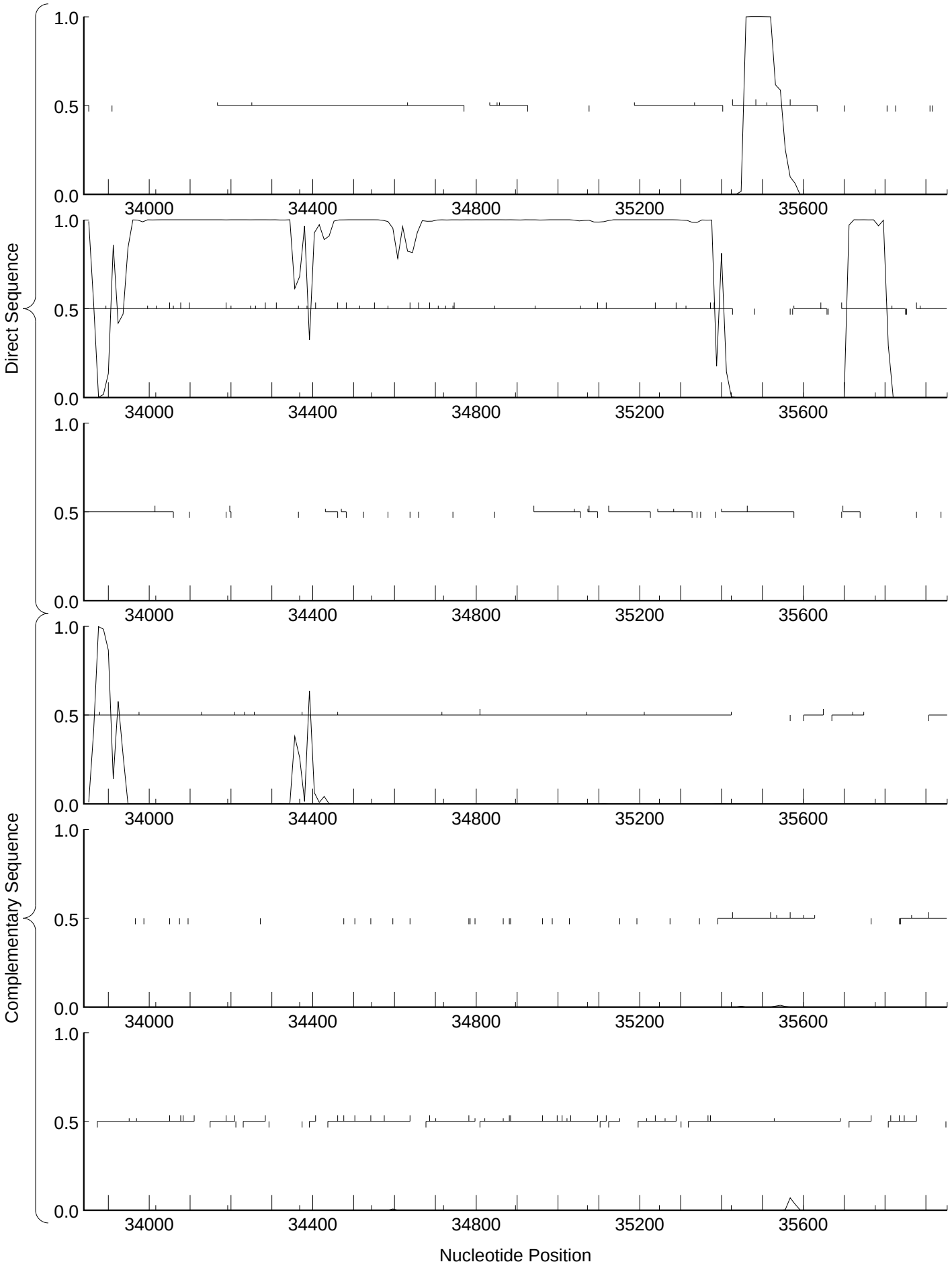
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 16/28



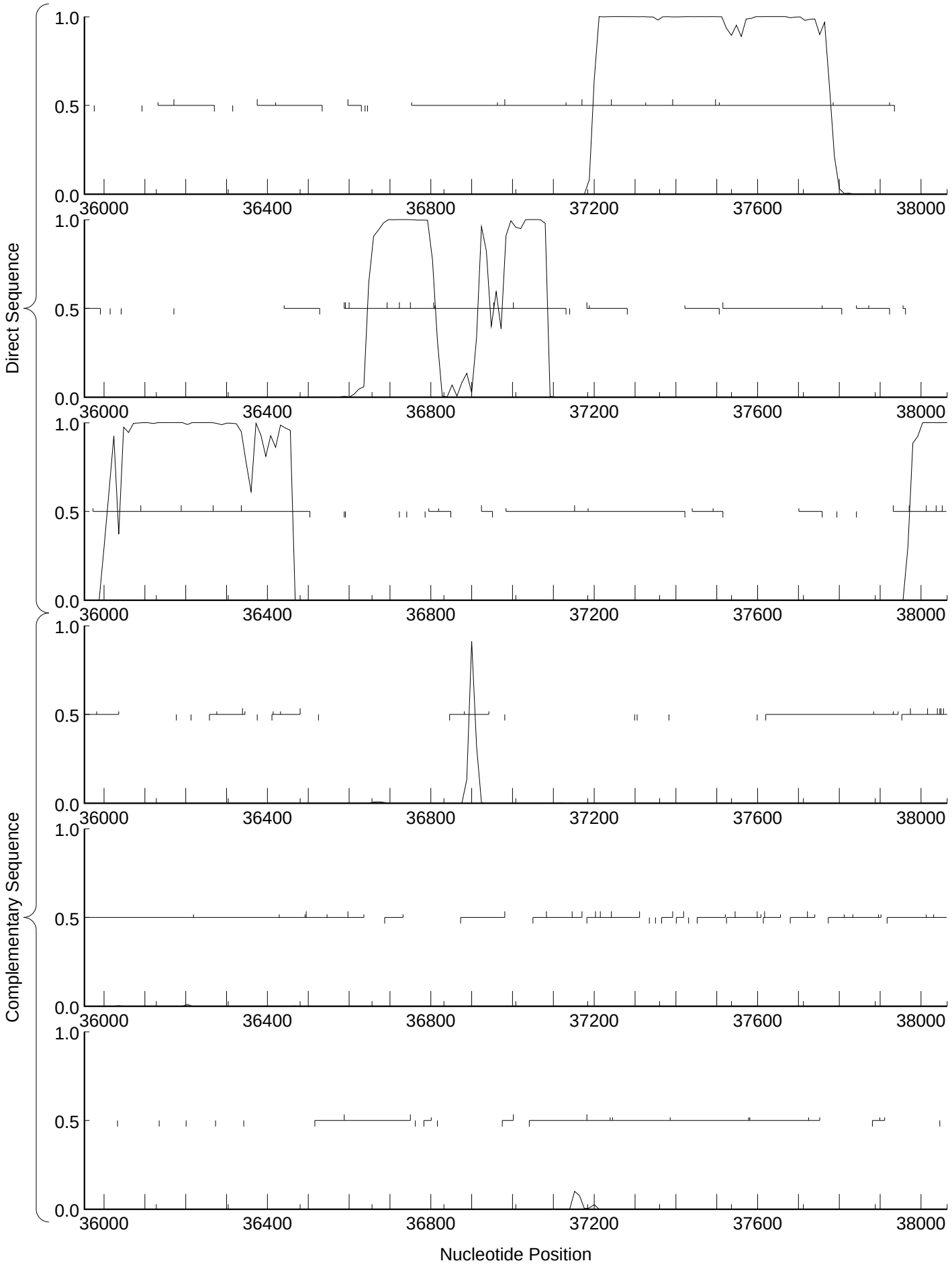
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 17/28



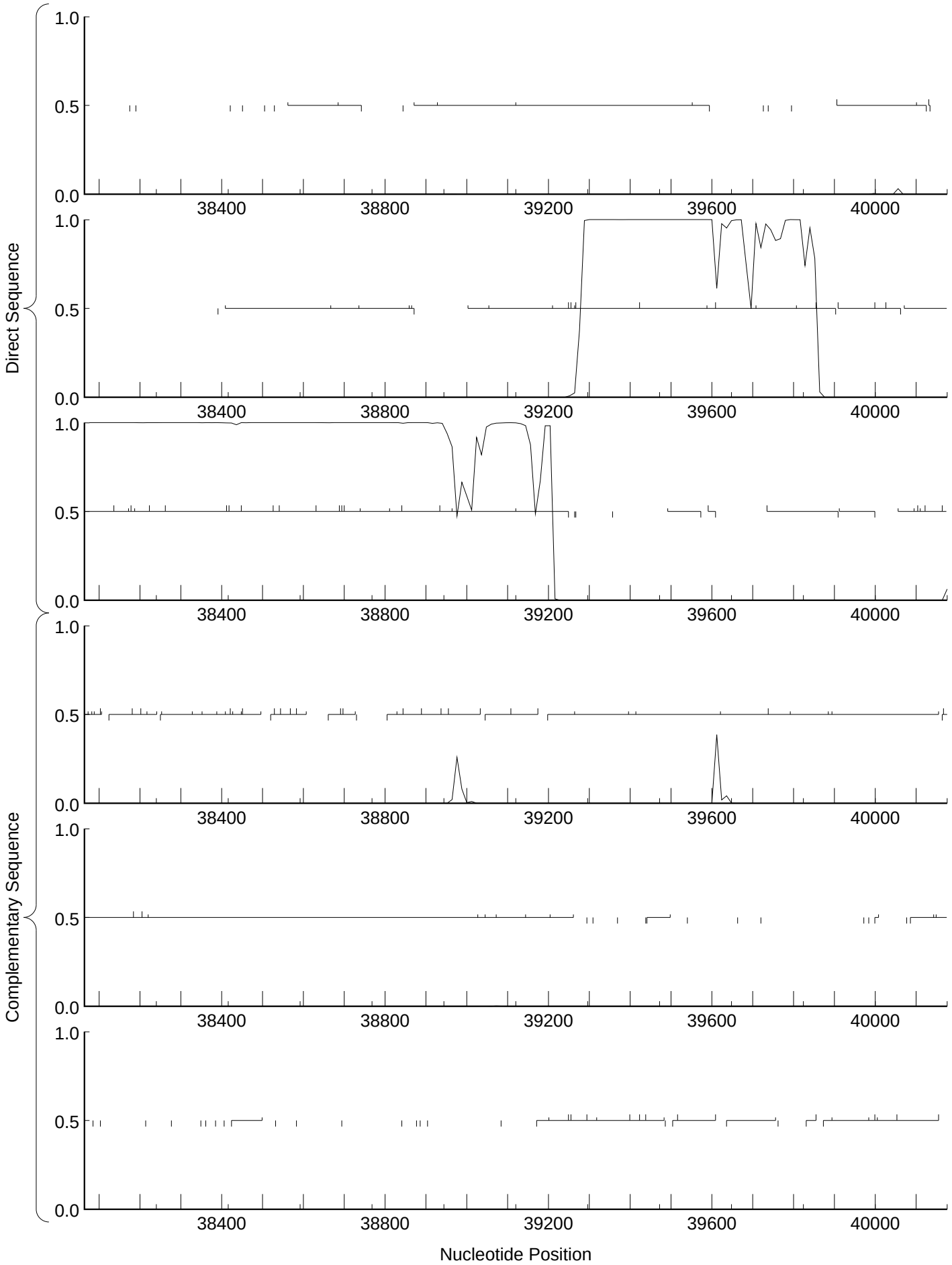
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 18/28



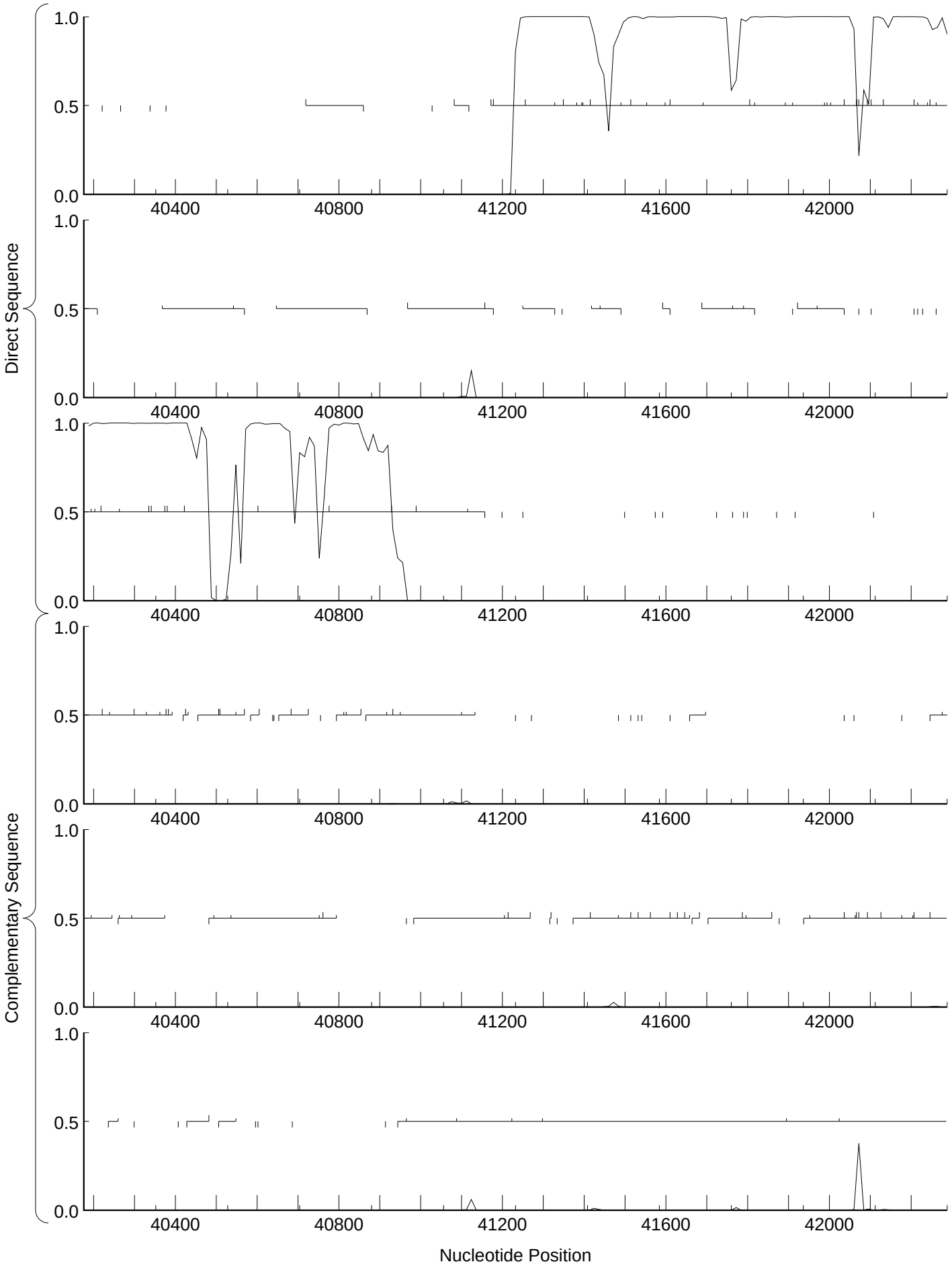
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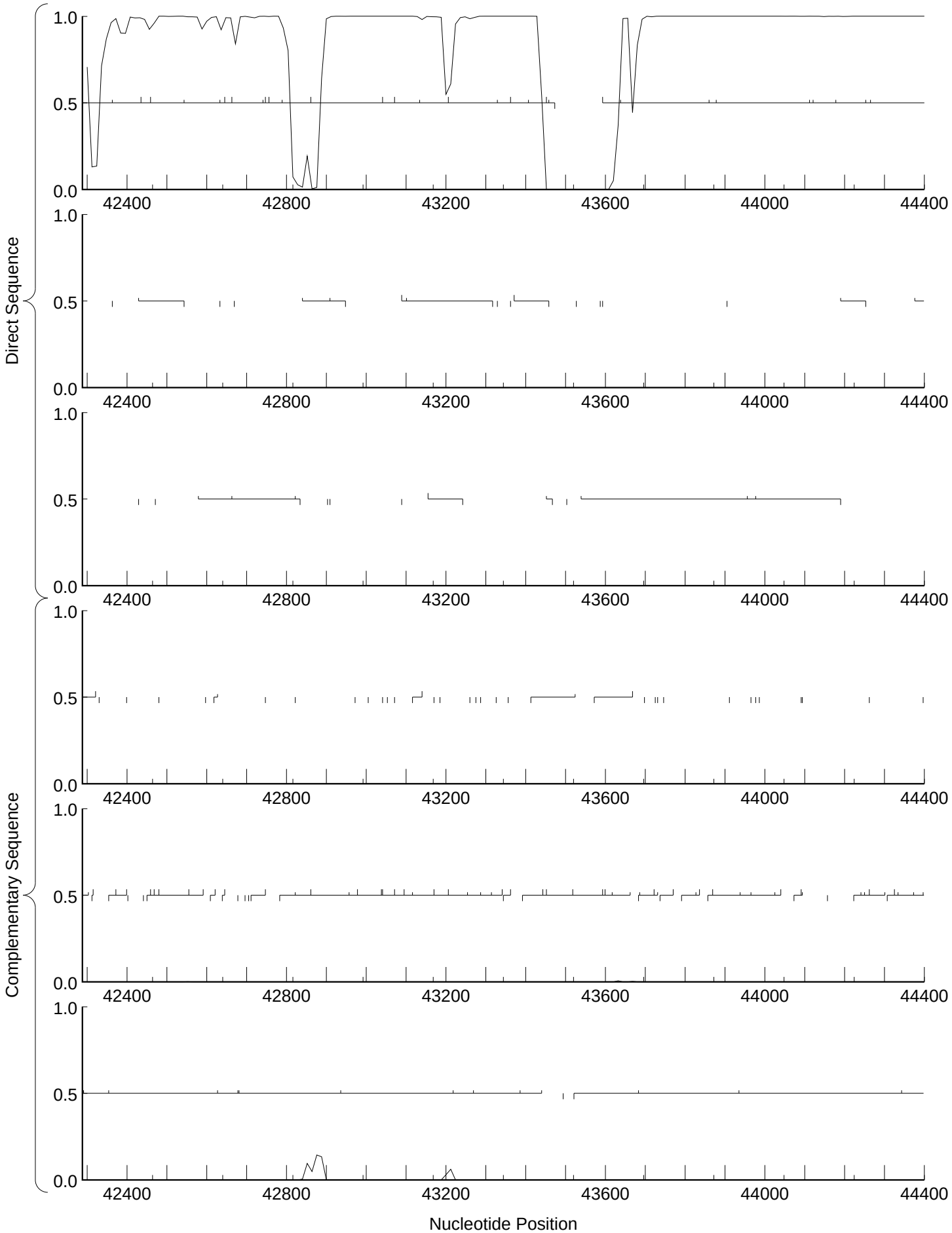
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 20/28



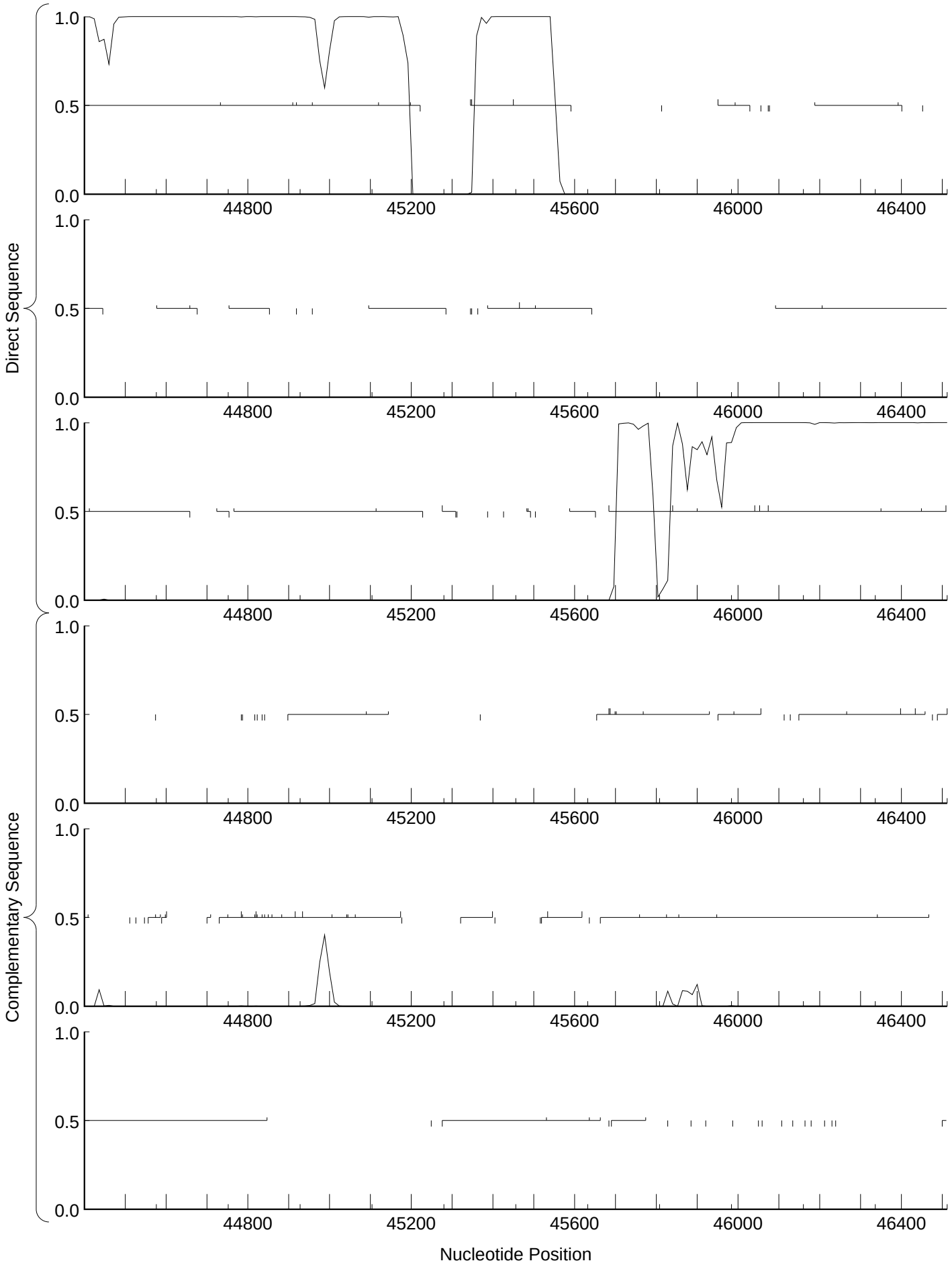
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 21/28



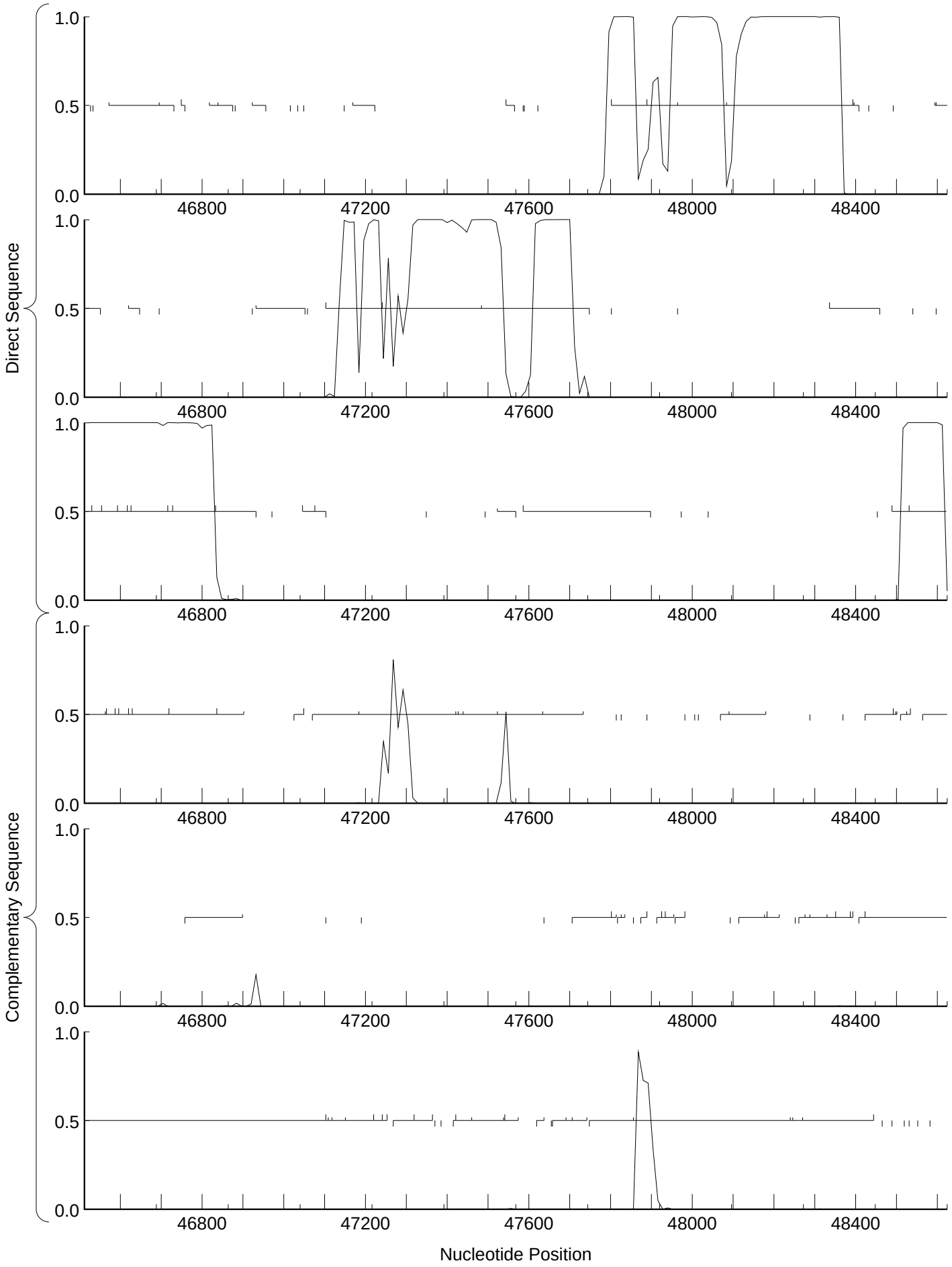
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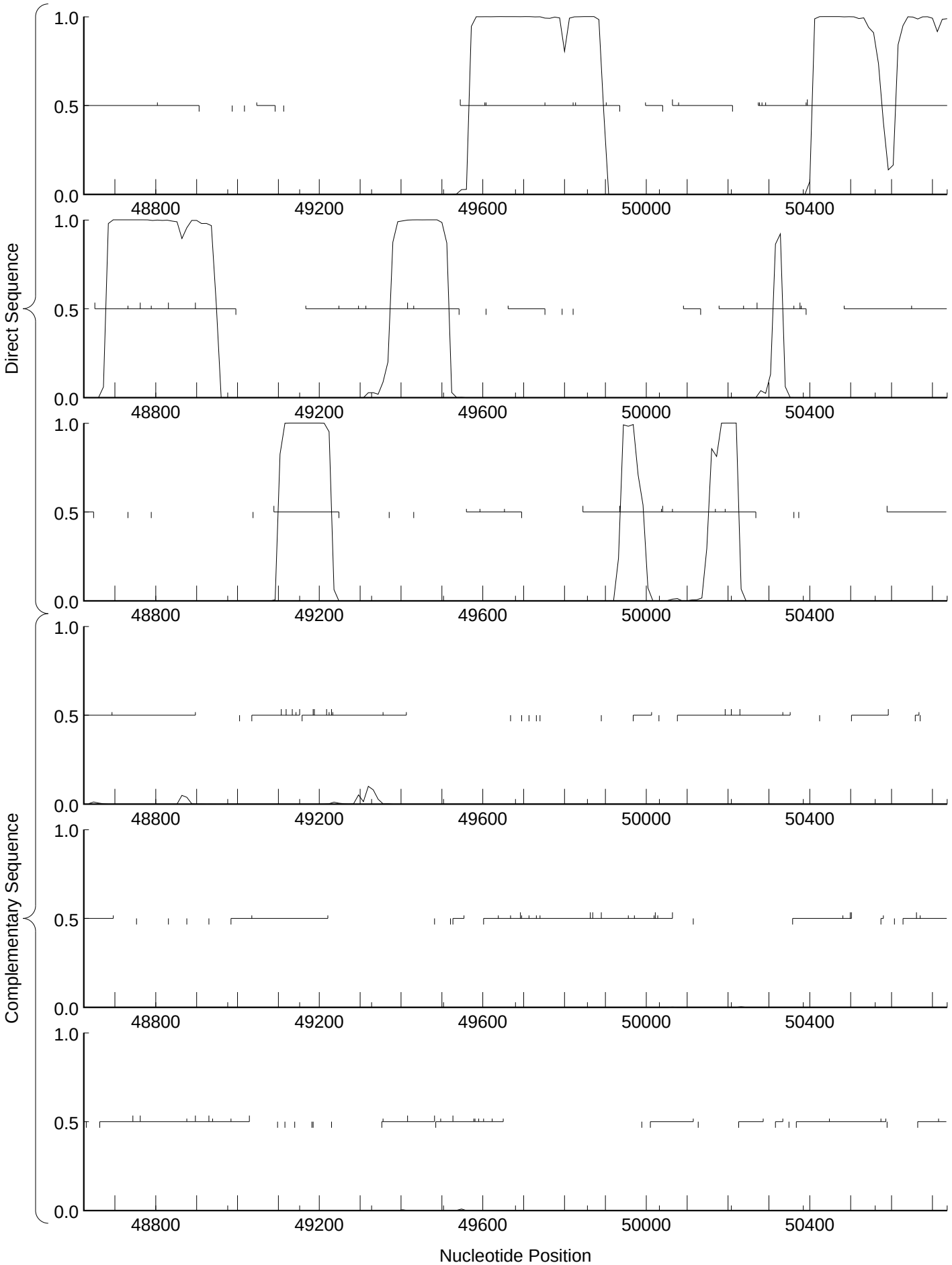
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 23/28



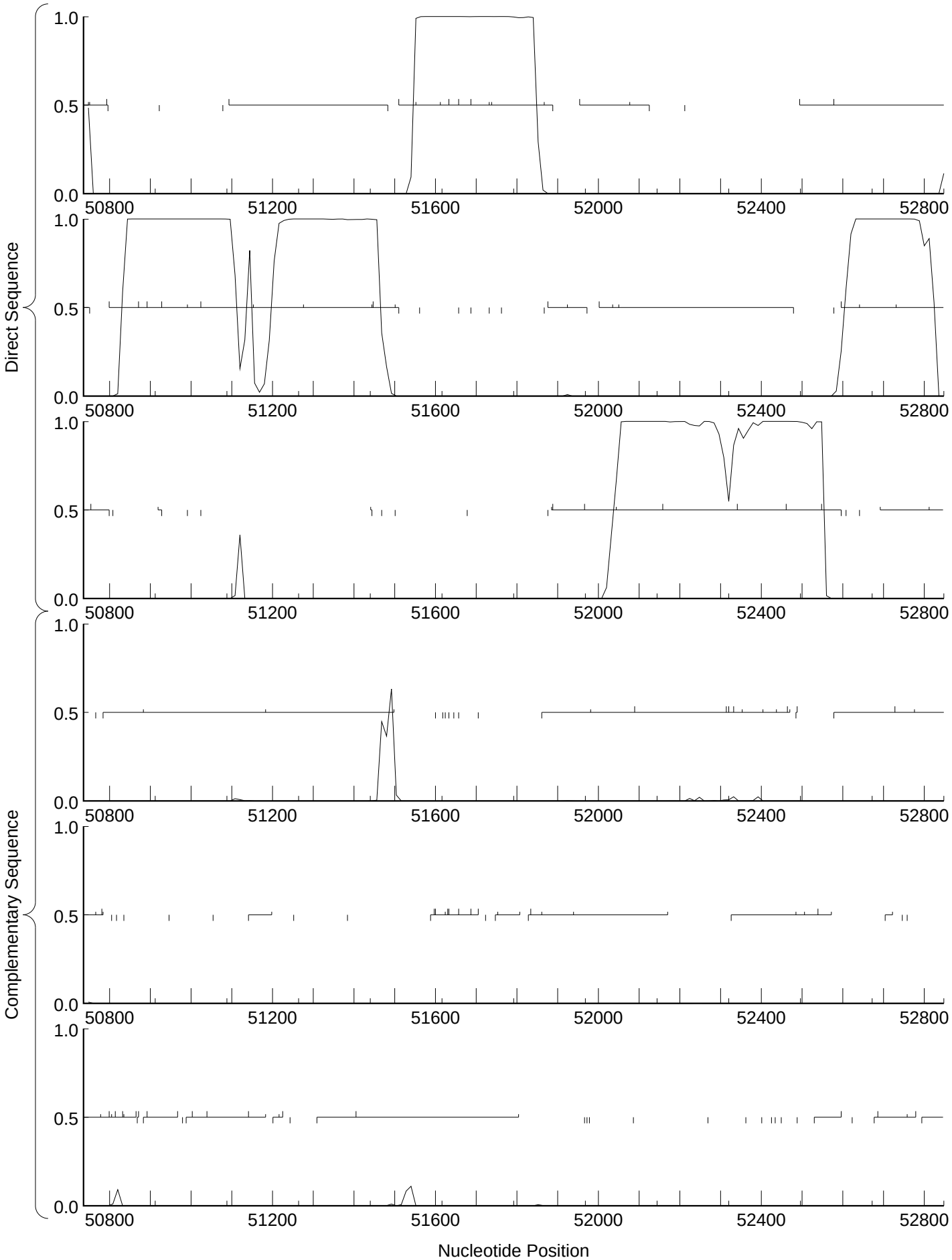
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 24/28



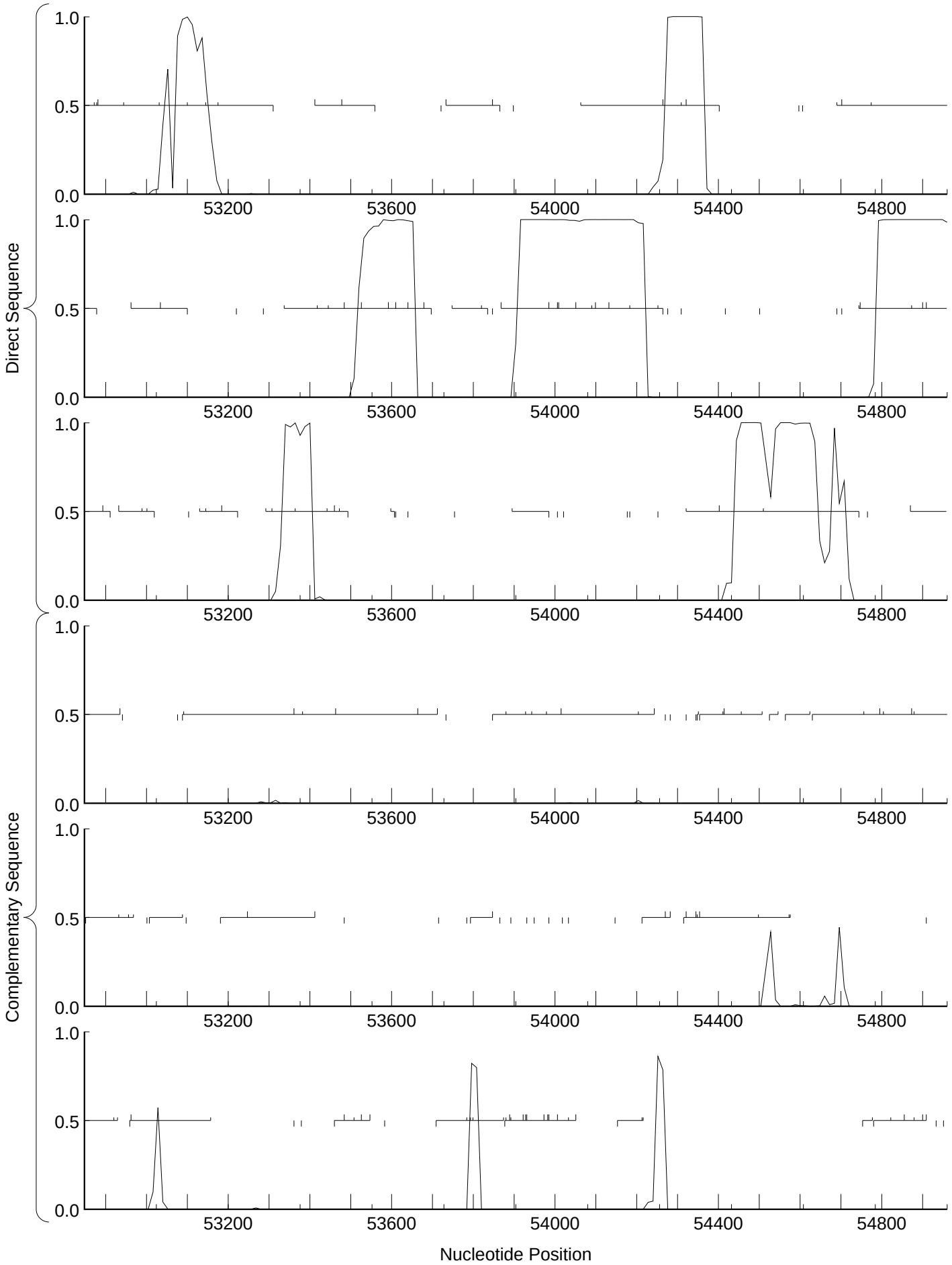
Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 25/28



Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 26/28



Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 27/28



Microbacterium phage Kate complete sequence, 56618 bp, circularly permuted, Cluster EF, Order 4, Window 96, Step 12, 28/28

