GeneMark

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

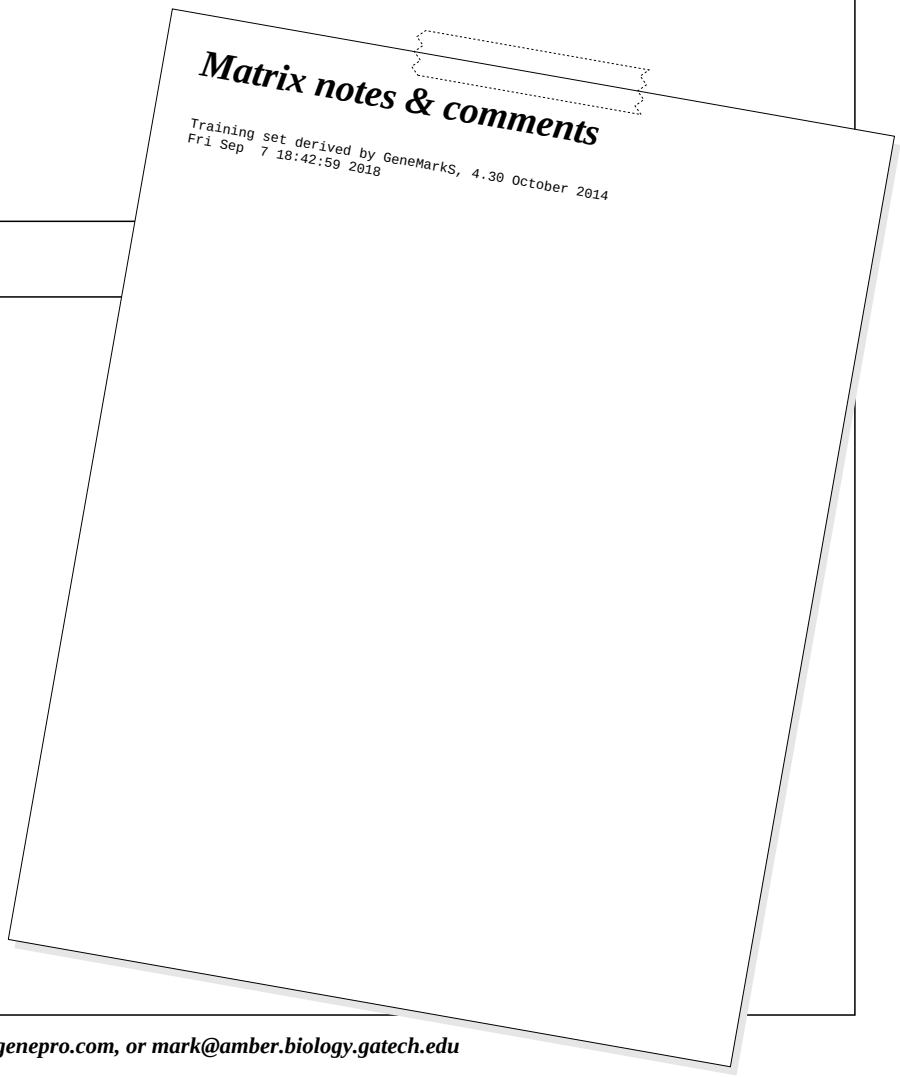
PROGRAM INFORMATION

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Analysis Date    :  3/29/24 at 3:39:41
Pages            :  23
Sequence Length  :  46365 bp
GC Content       :   68.31%

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

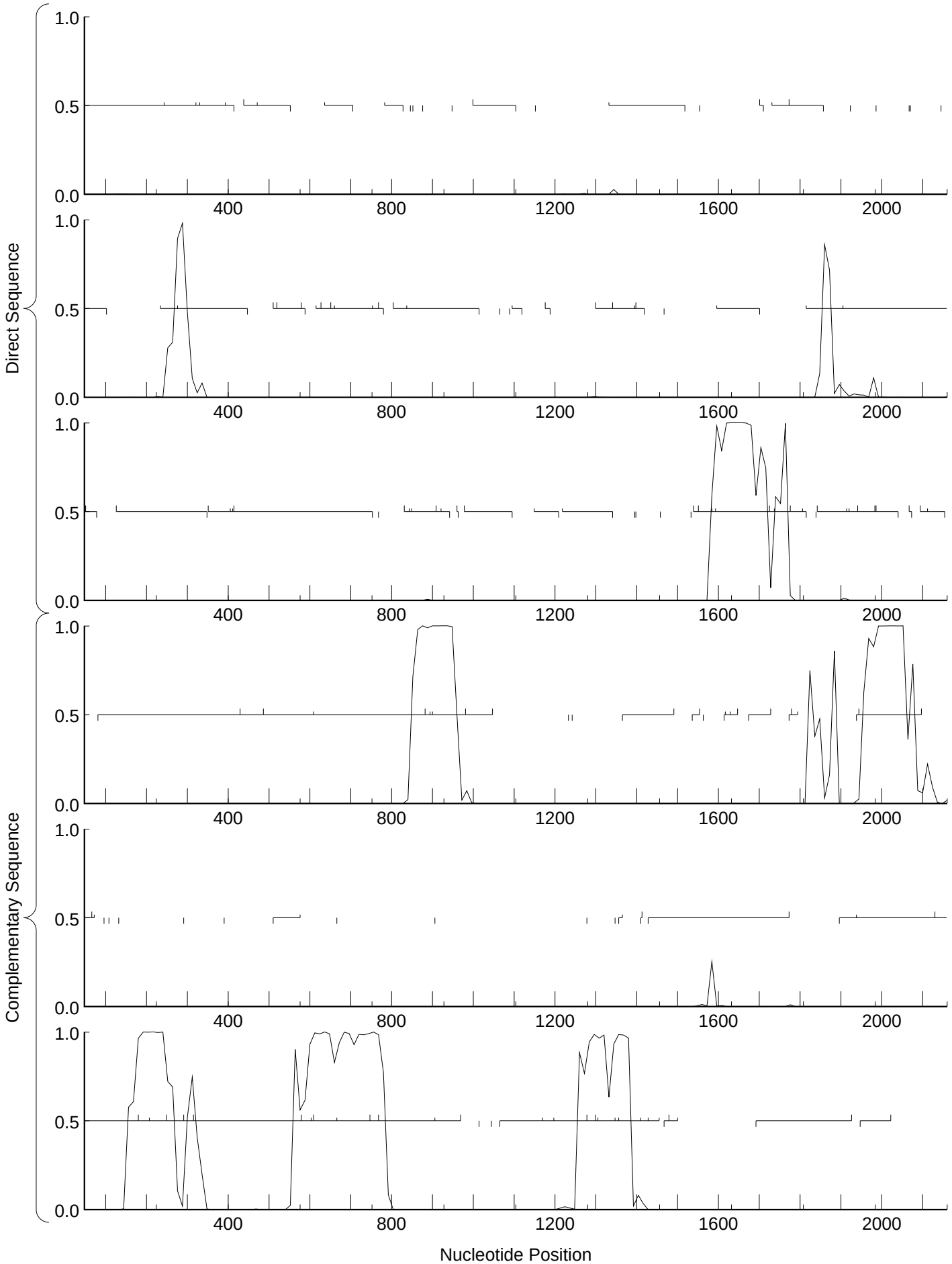
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Matrix     :  -
Author     :  -

Order      :  4
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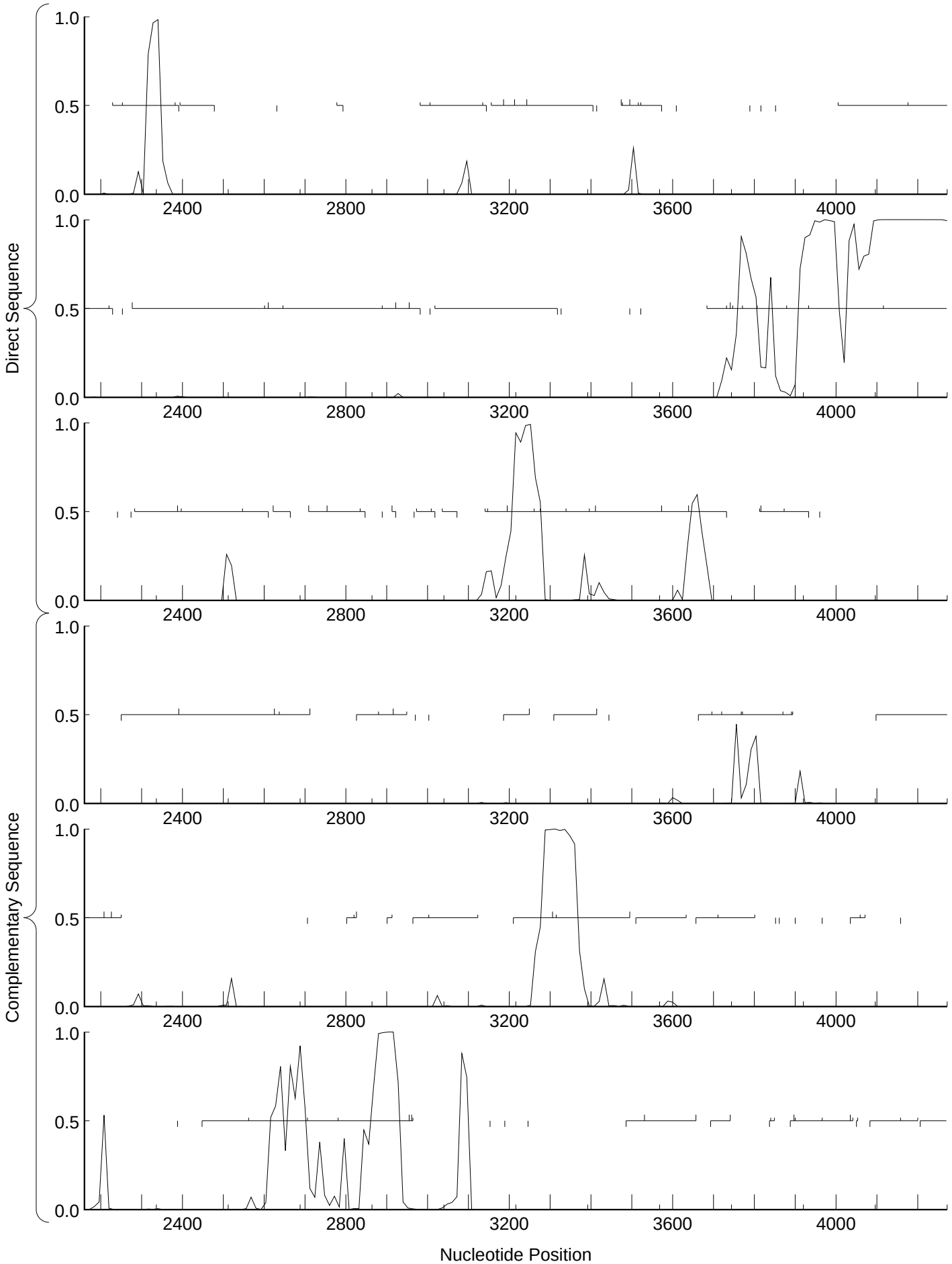

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

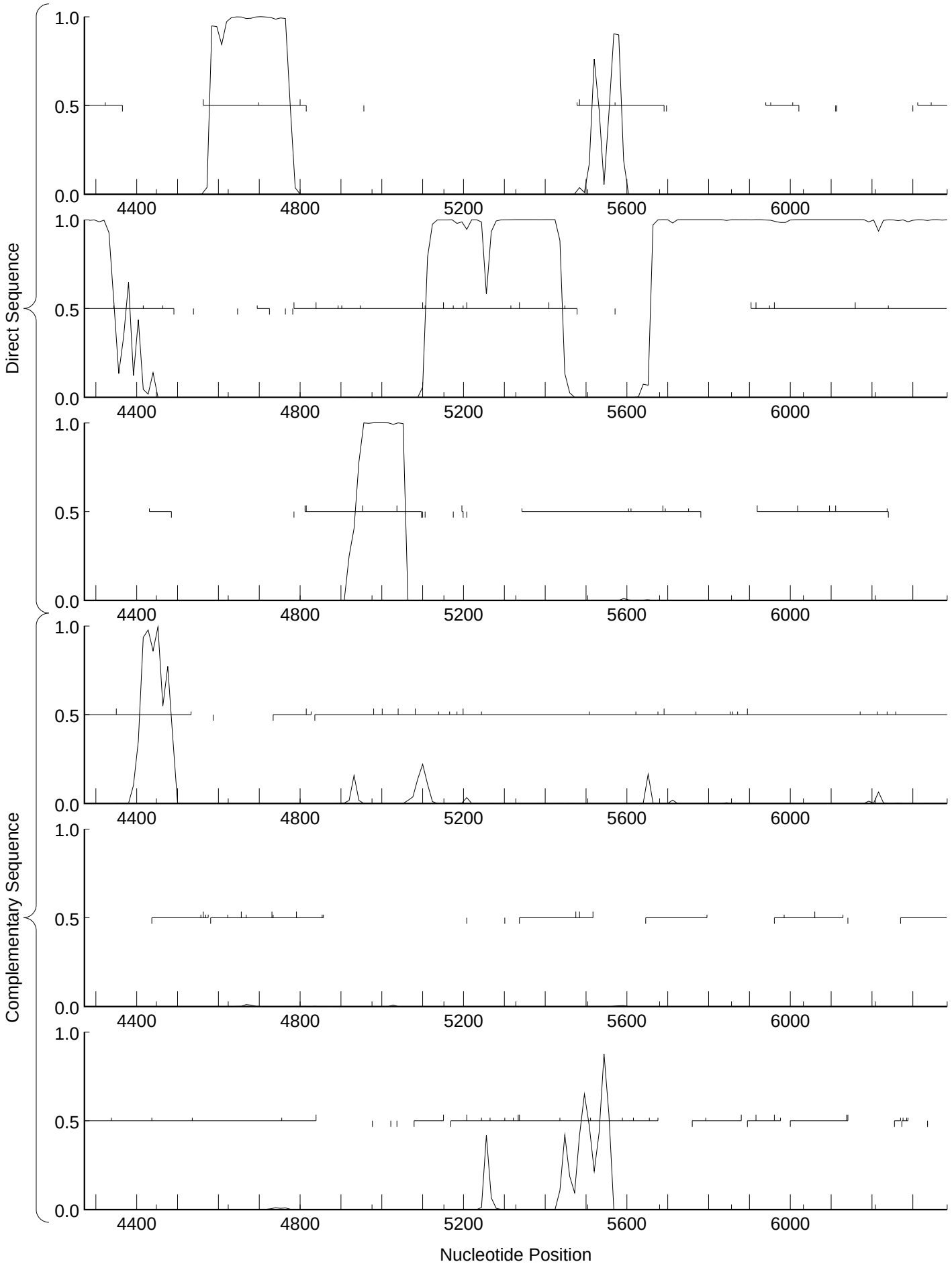
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 2/23



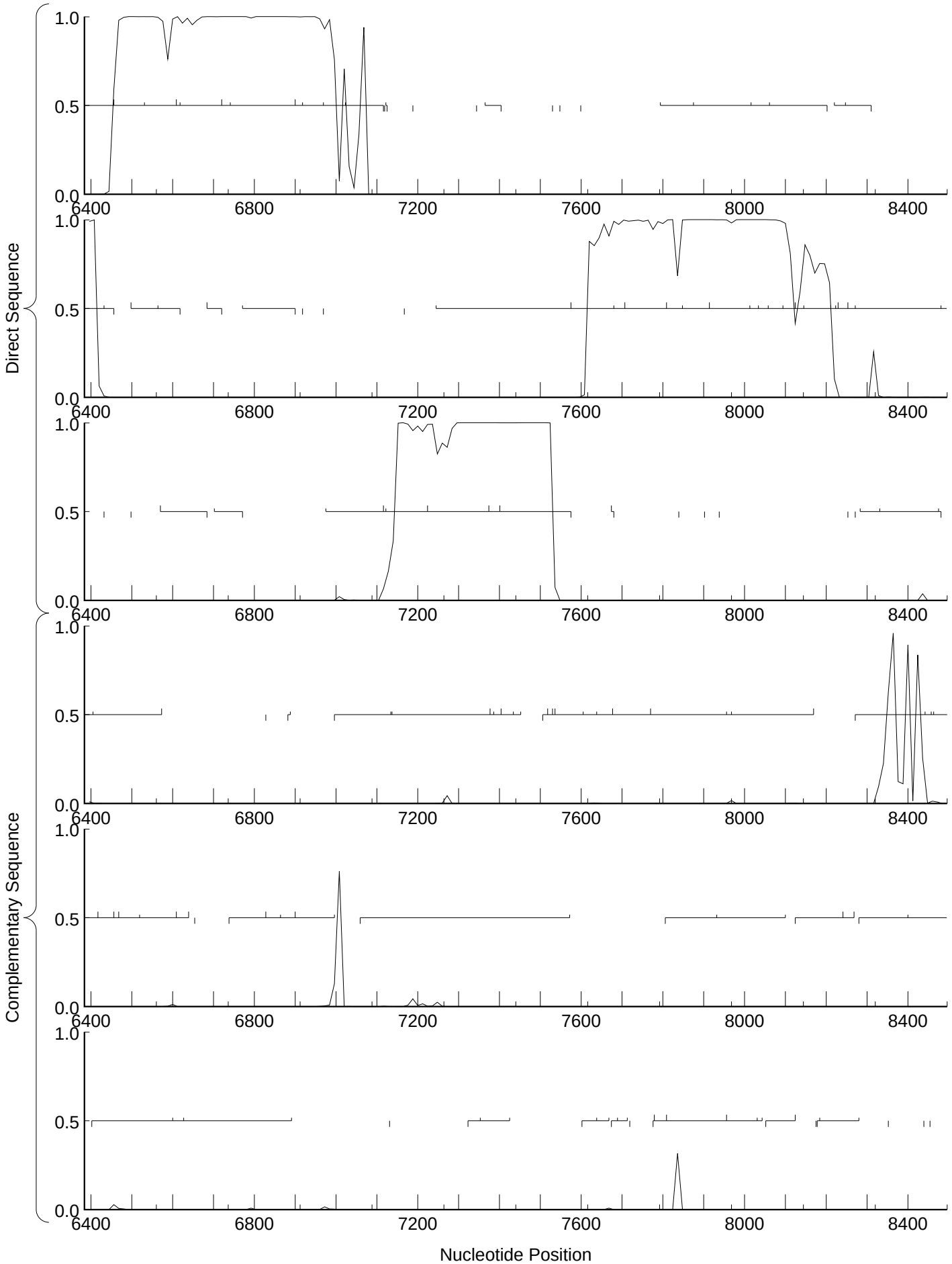
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 3/23



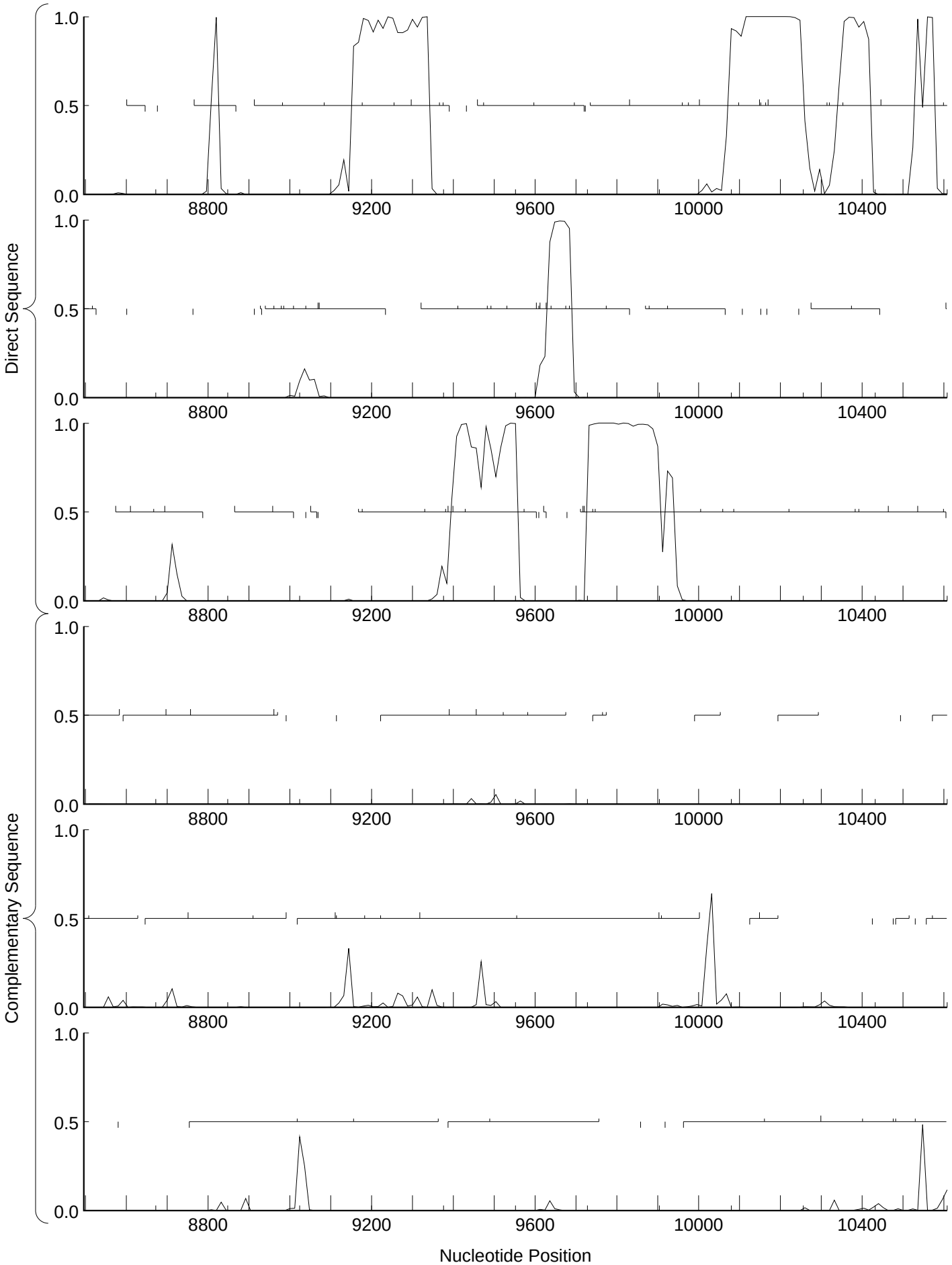
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 4/23



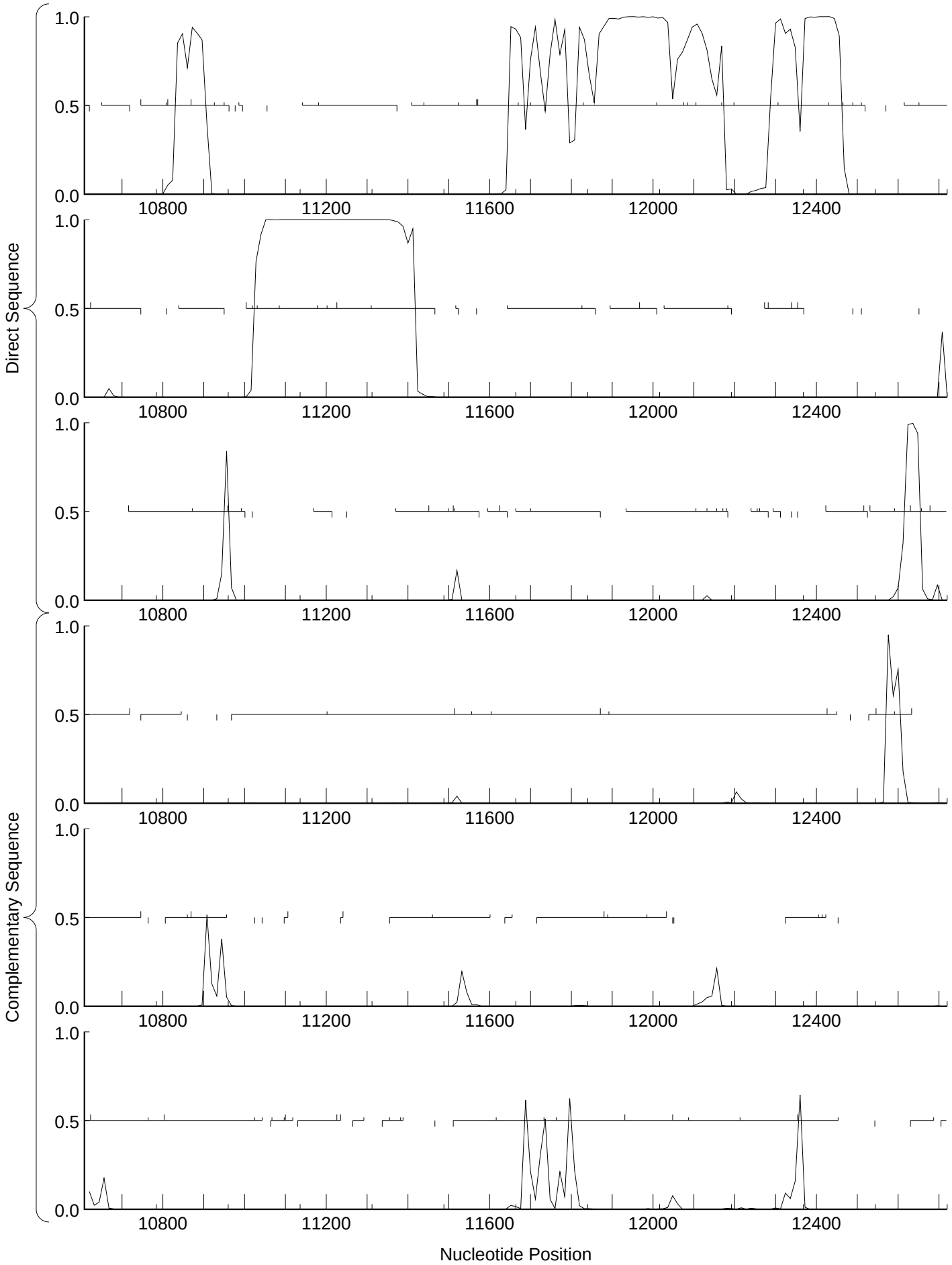
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 5/23



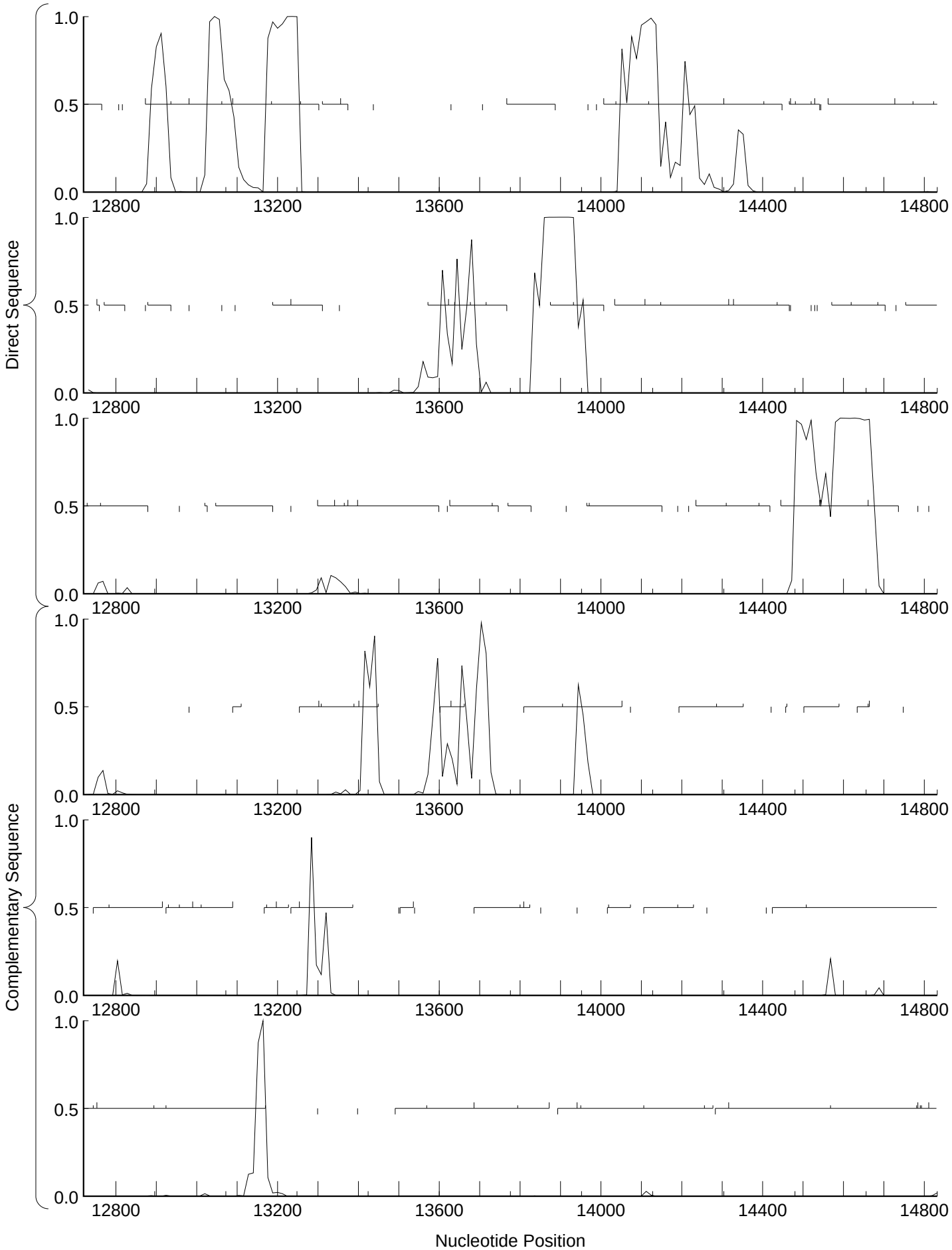
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 6/23



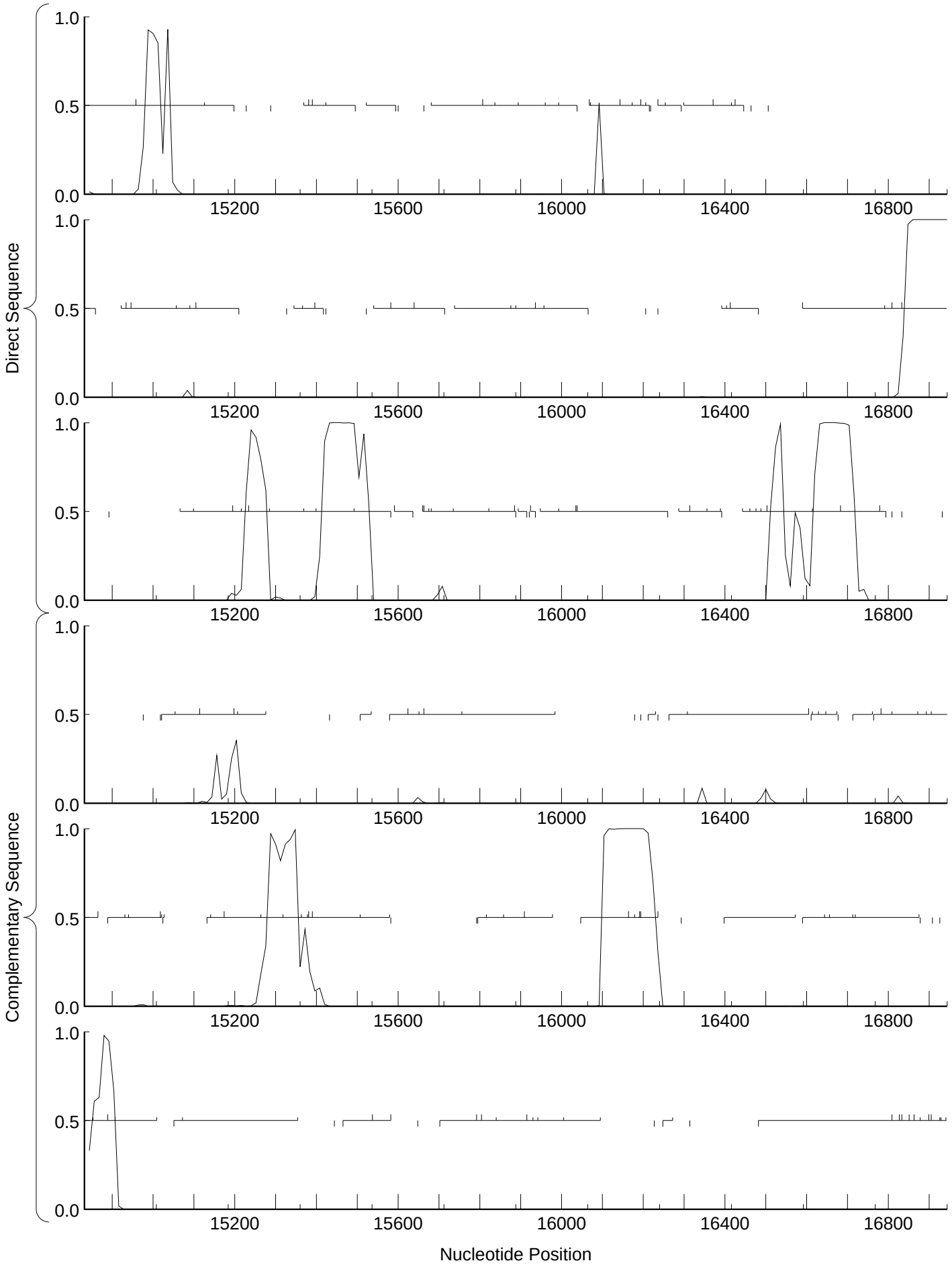
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 7/23



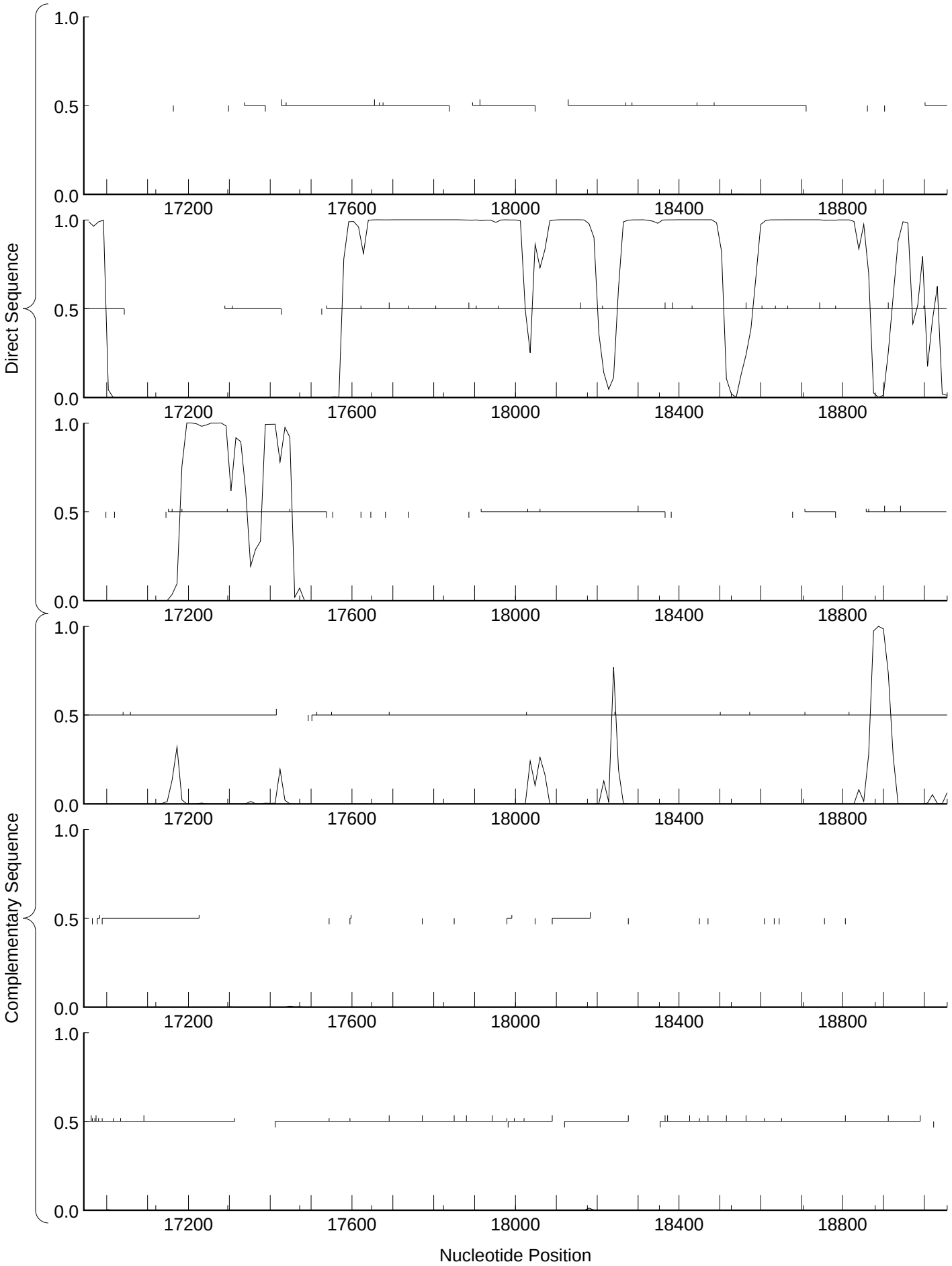
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 8/23



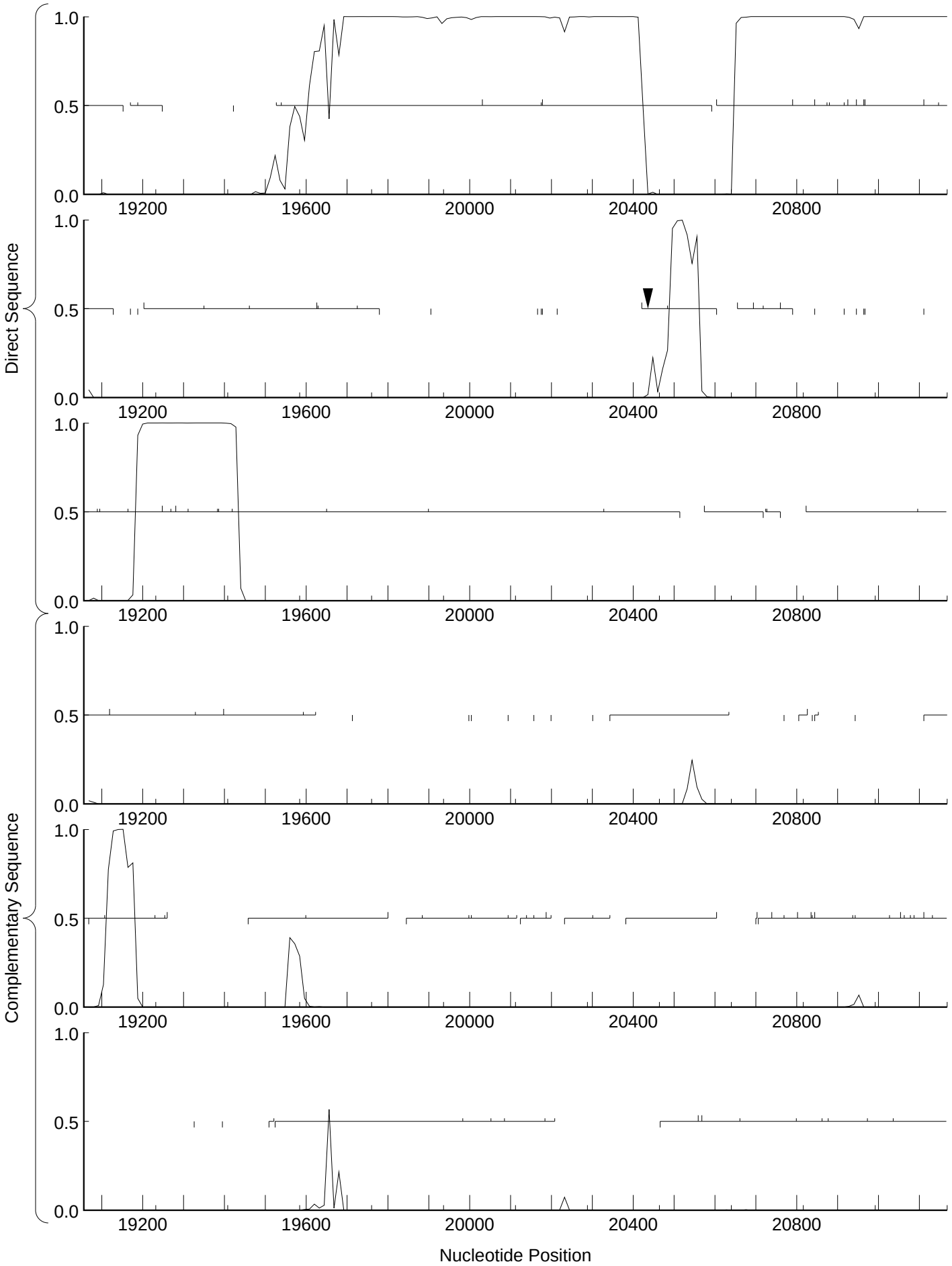
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 9/23



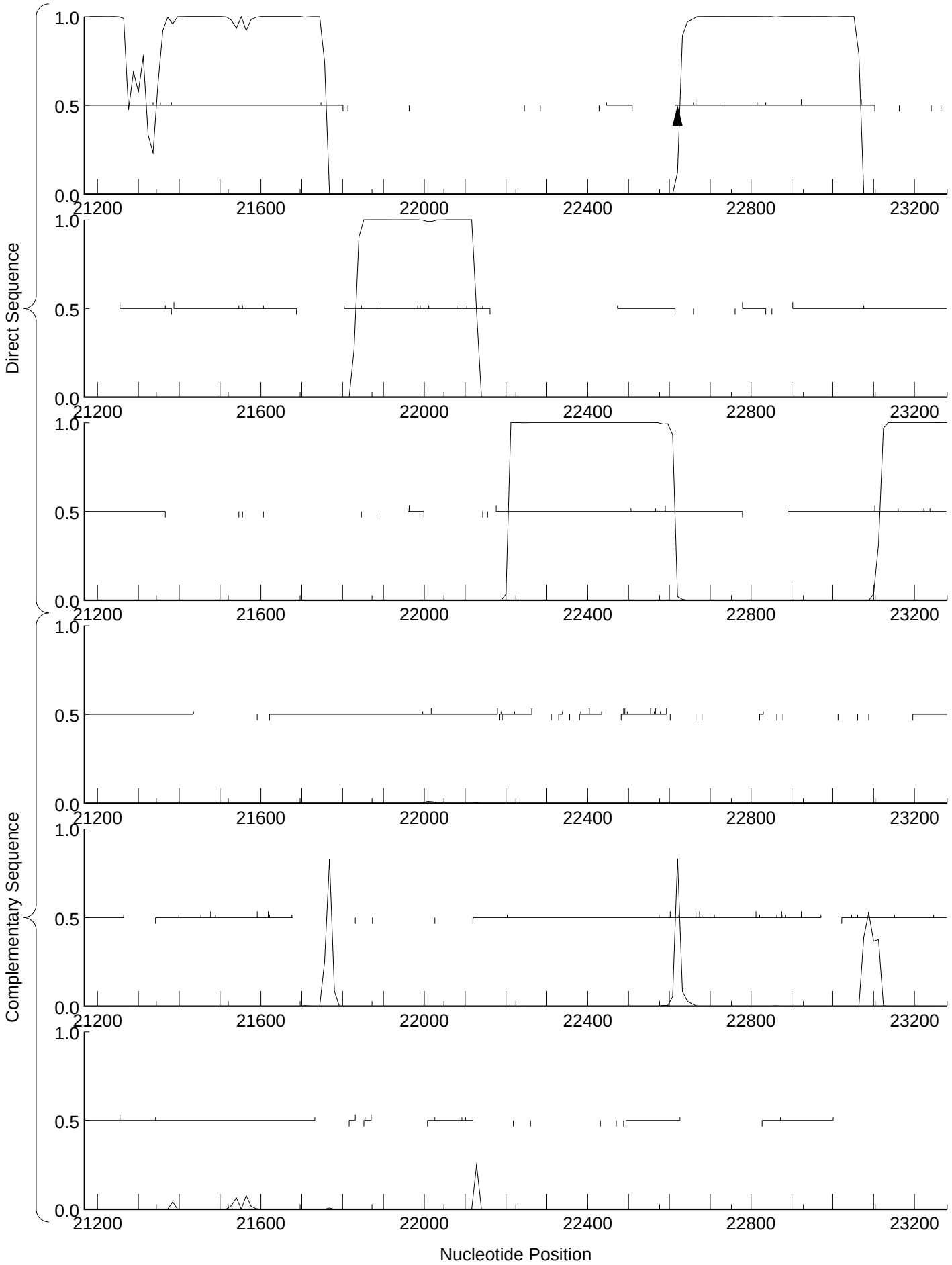
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 10/23



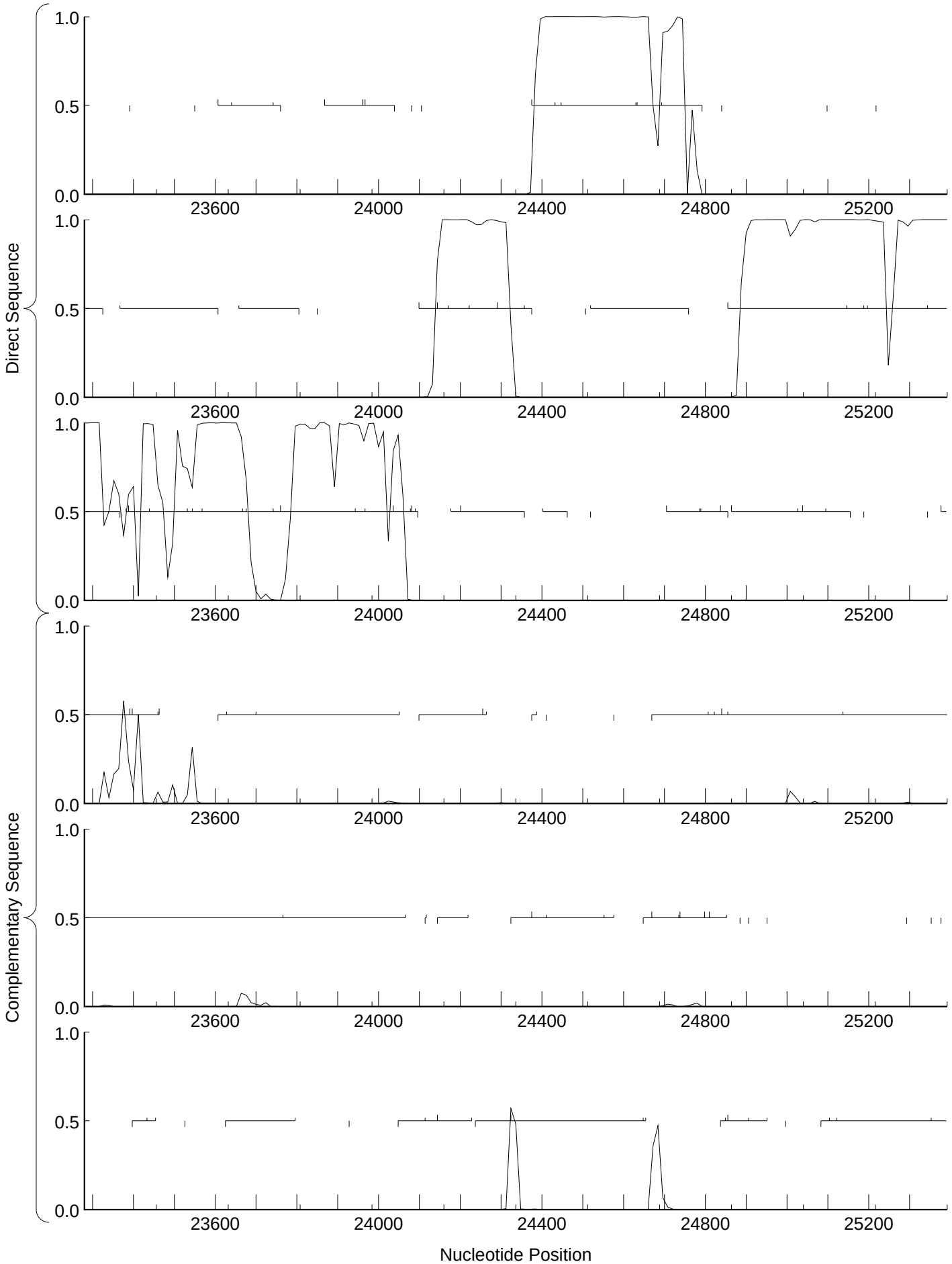
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 11/23



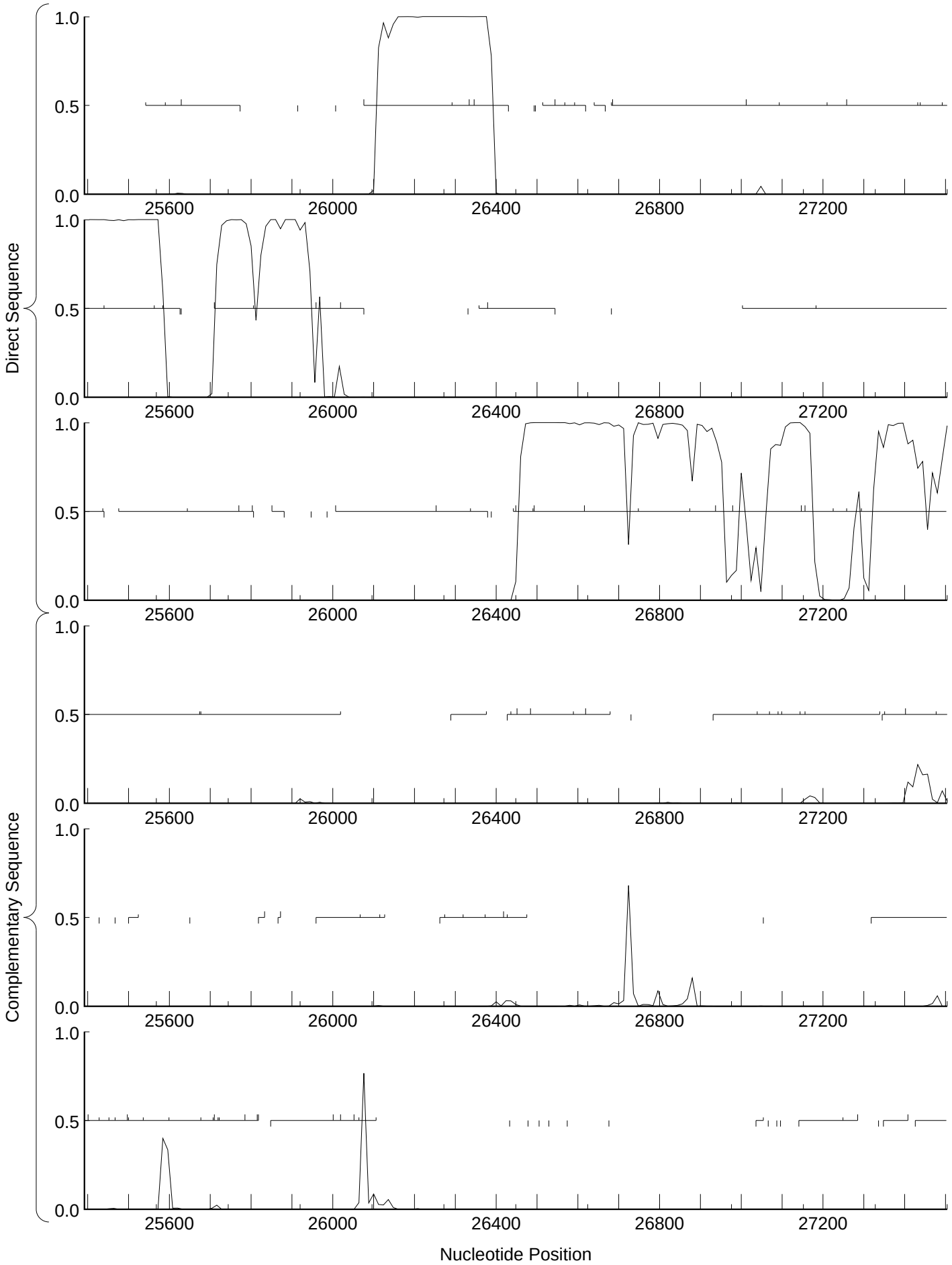
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 12/23



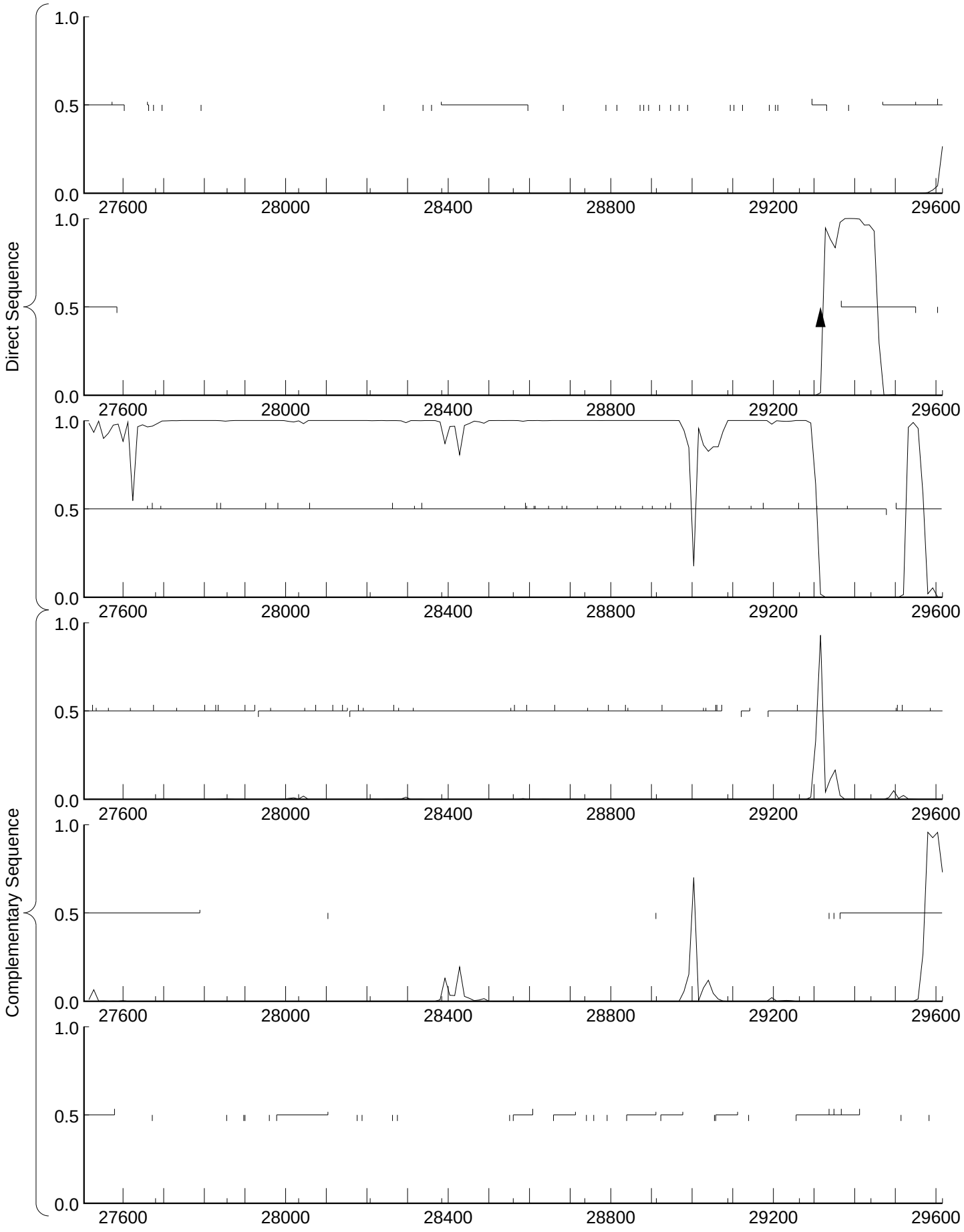
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 13/23



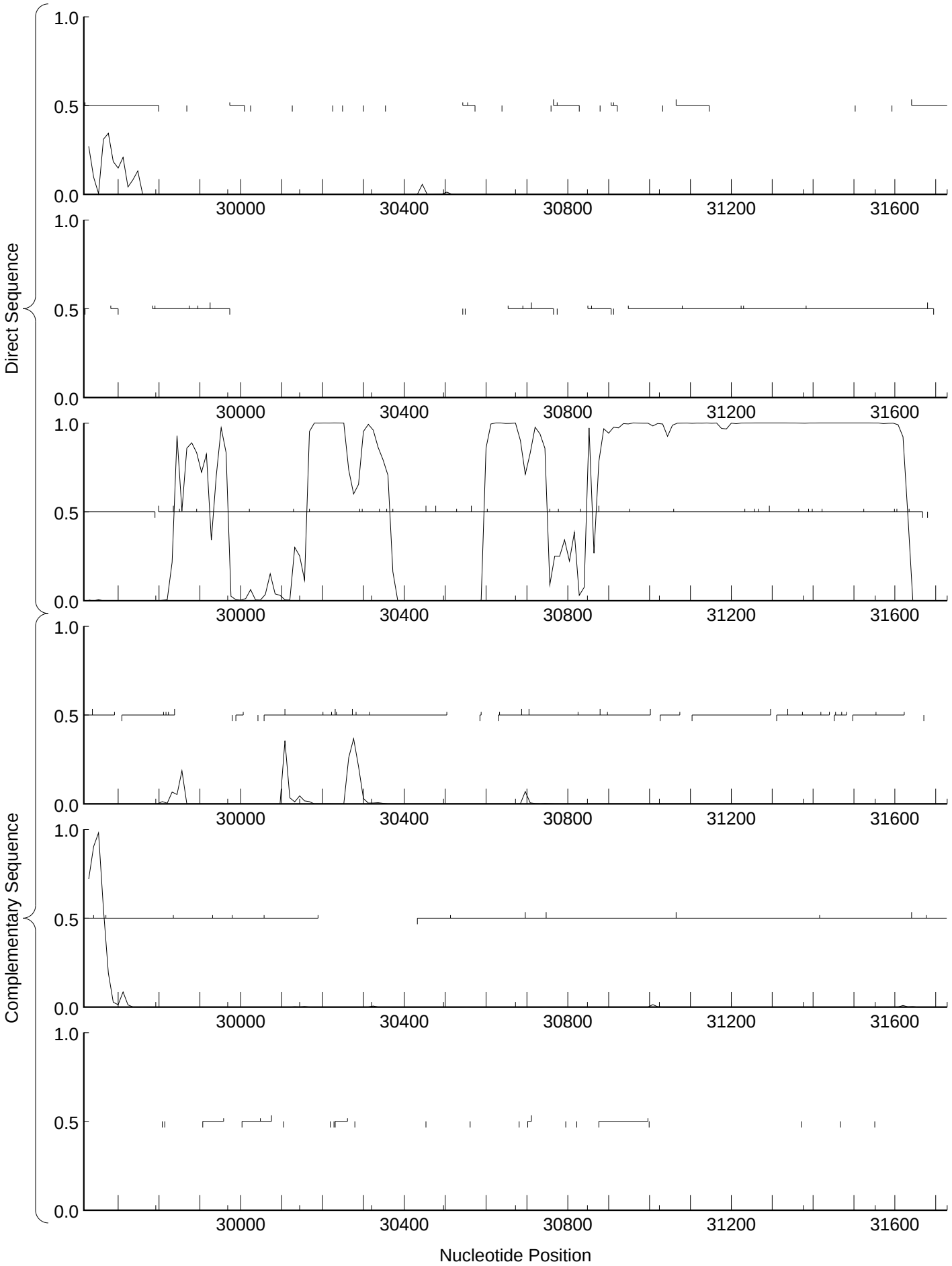
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 14/23



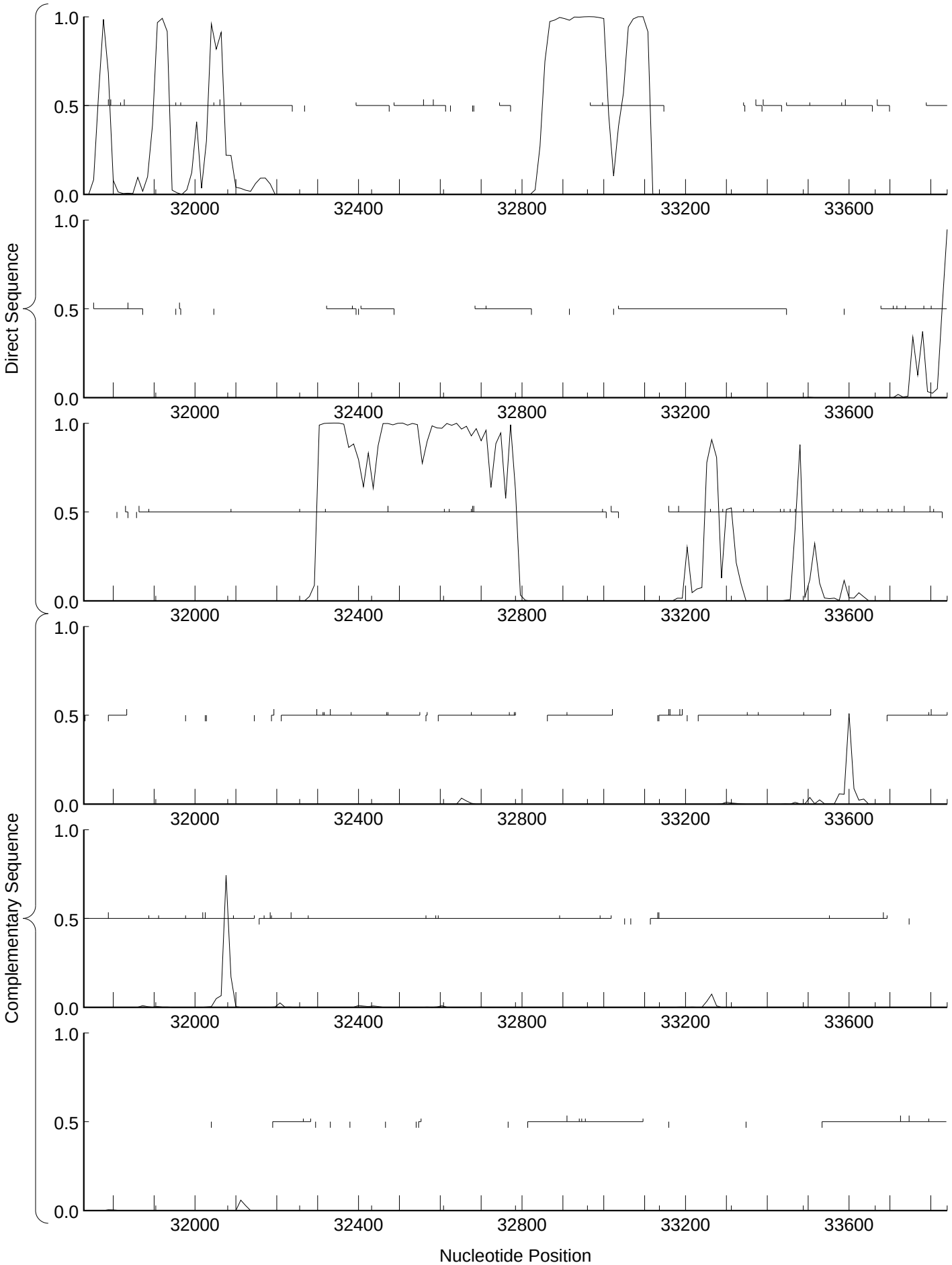
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 15/23



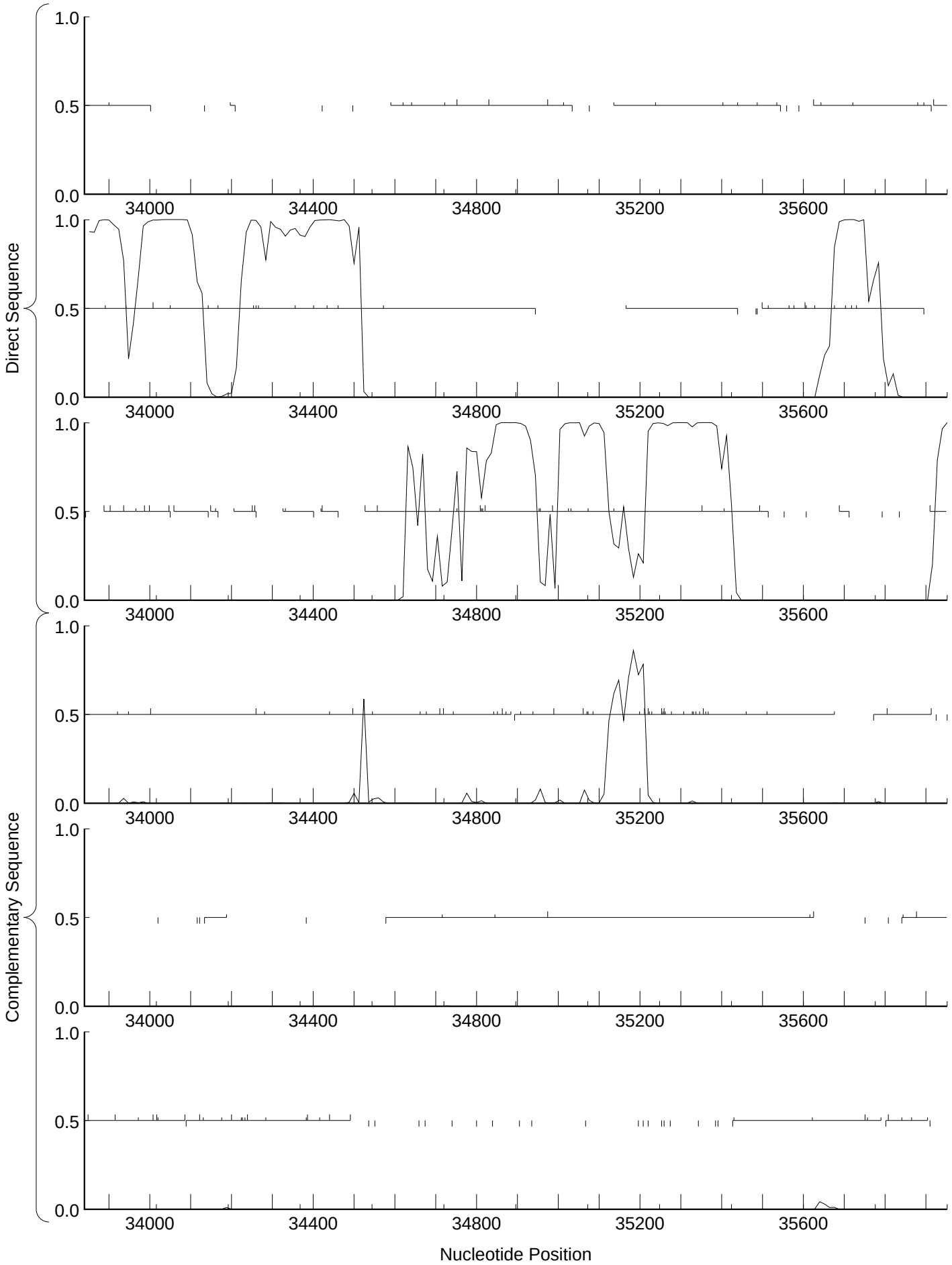
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 16/23



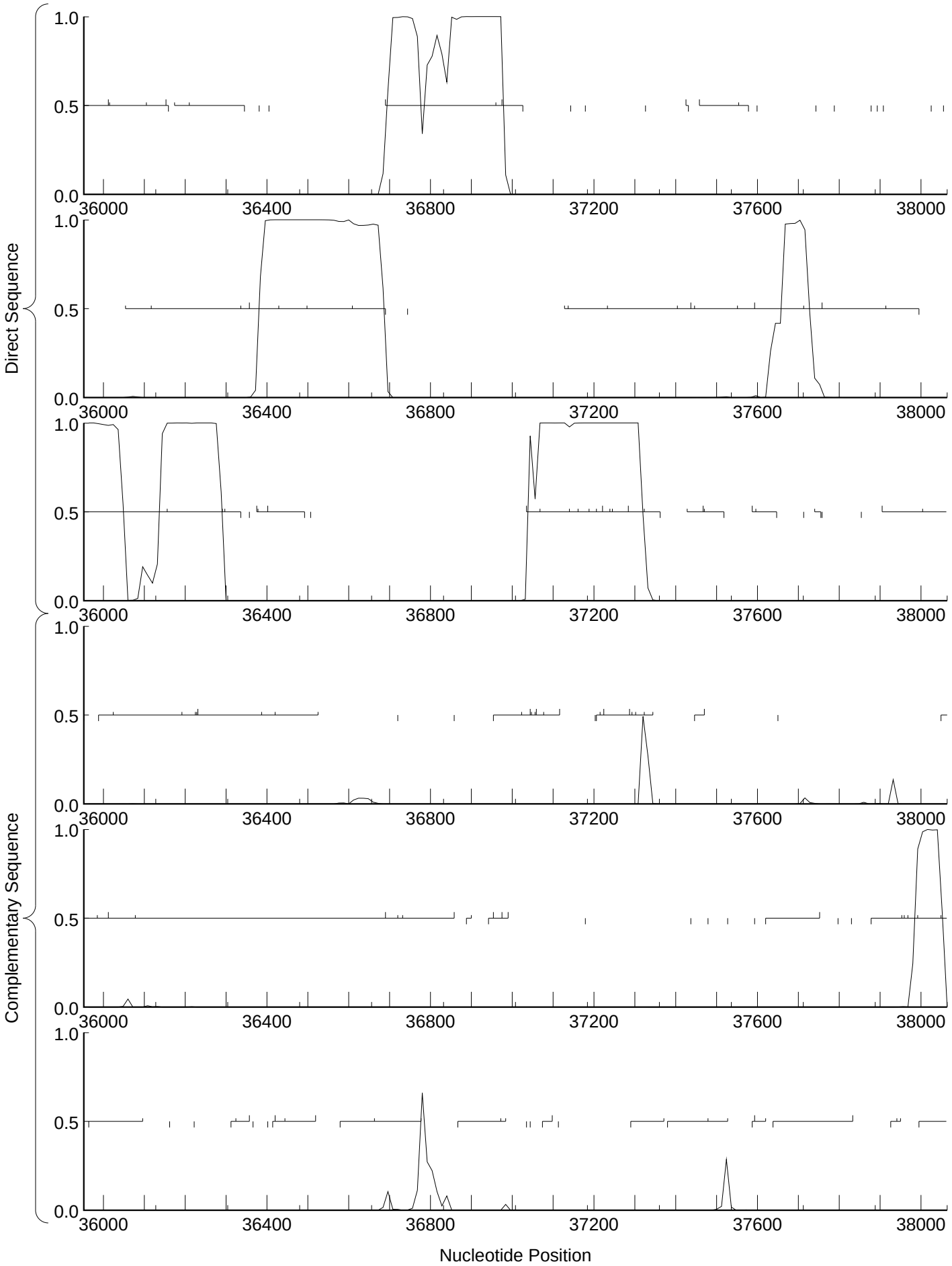
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 17/23



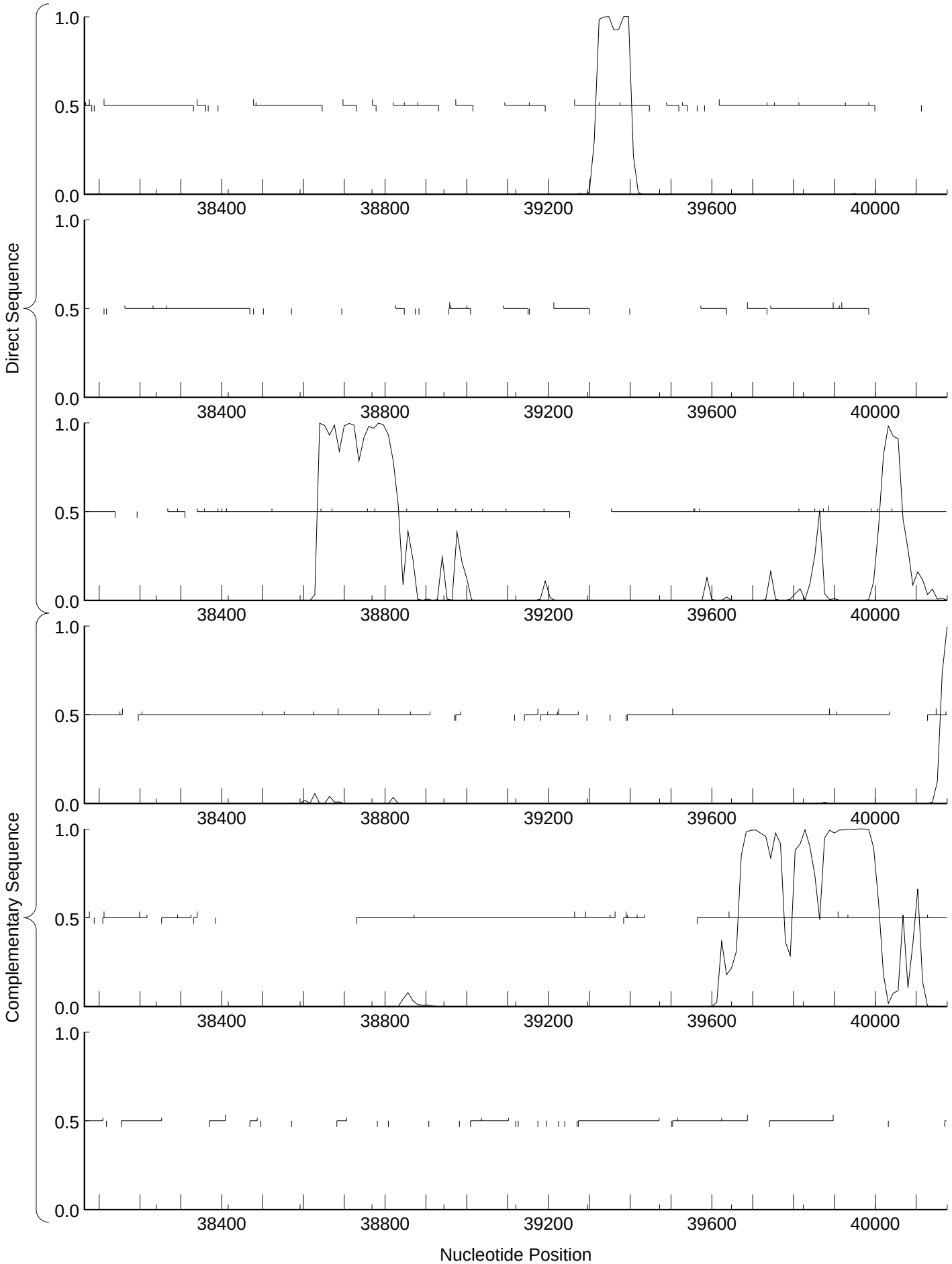
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 18/23



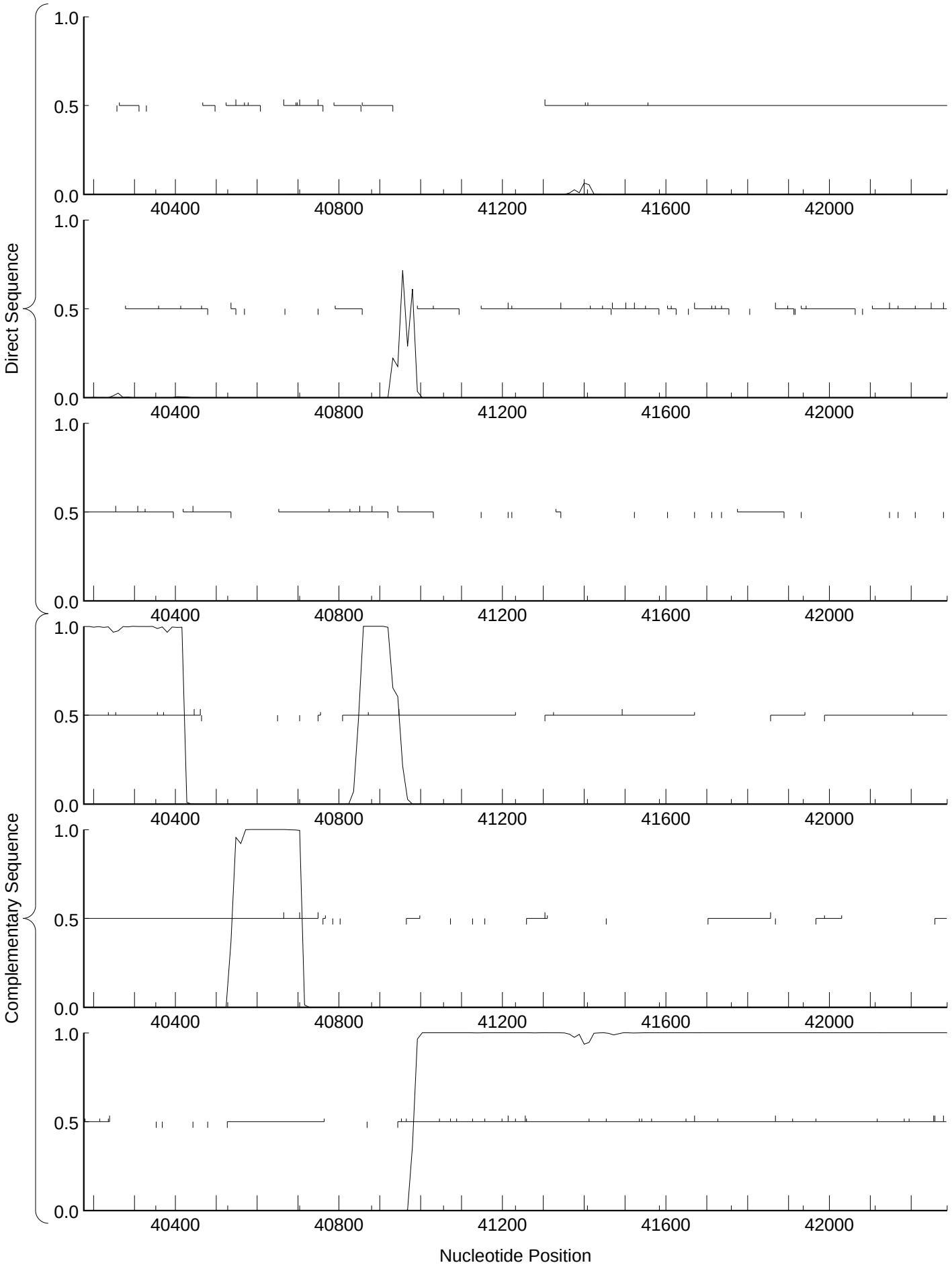
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 19/23



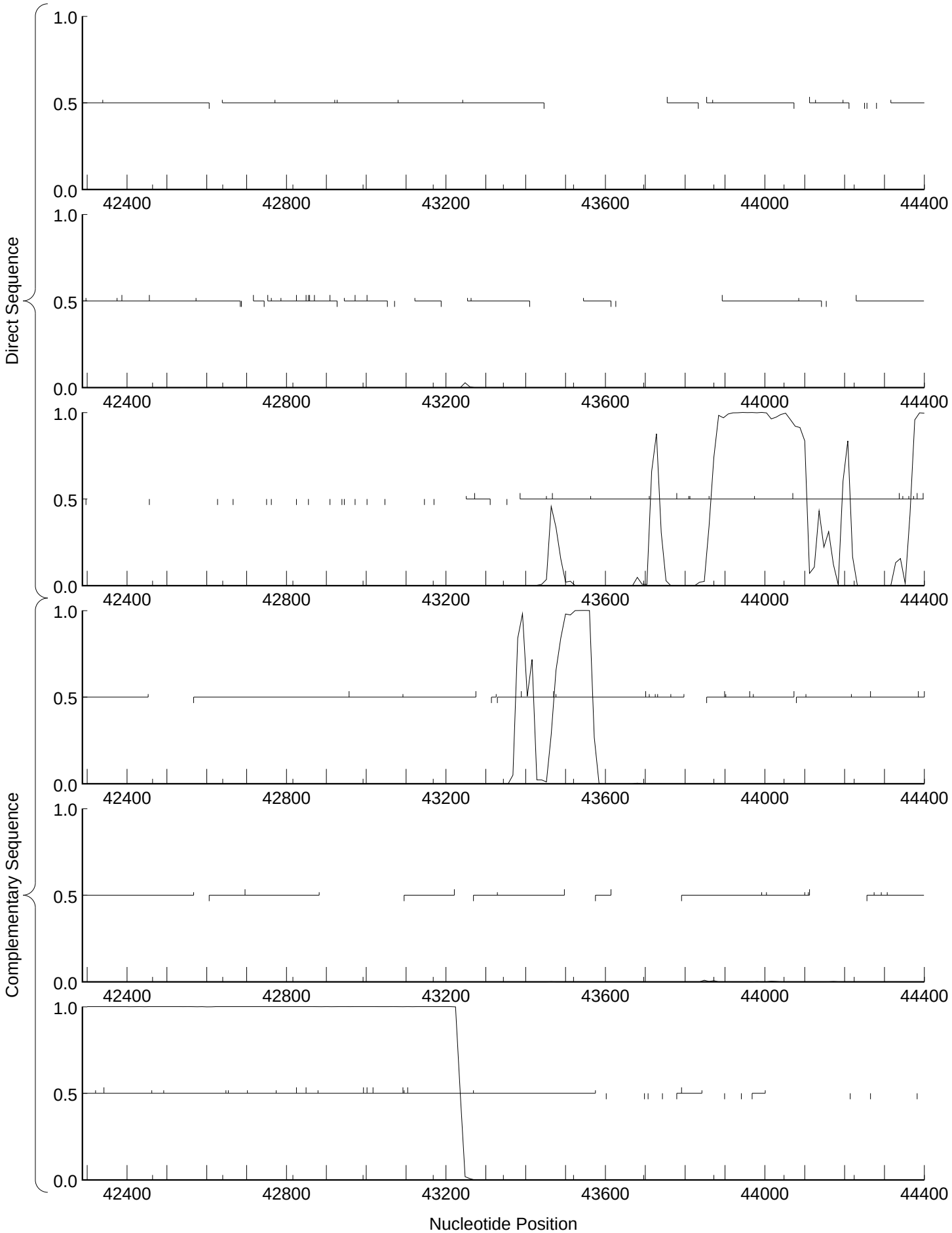
Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 20/23



Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 21/23



Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 22/23



Microbacterium phage Min1 complete genome, 46365 bp, circularly permuted, Order 4, Window 96, Step 12, 23/23

