GeneMark

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

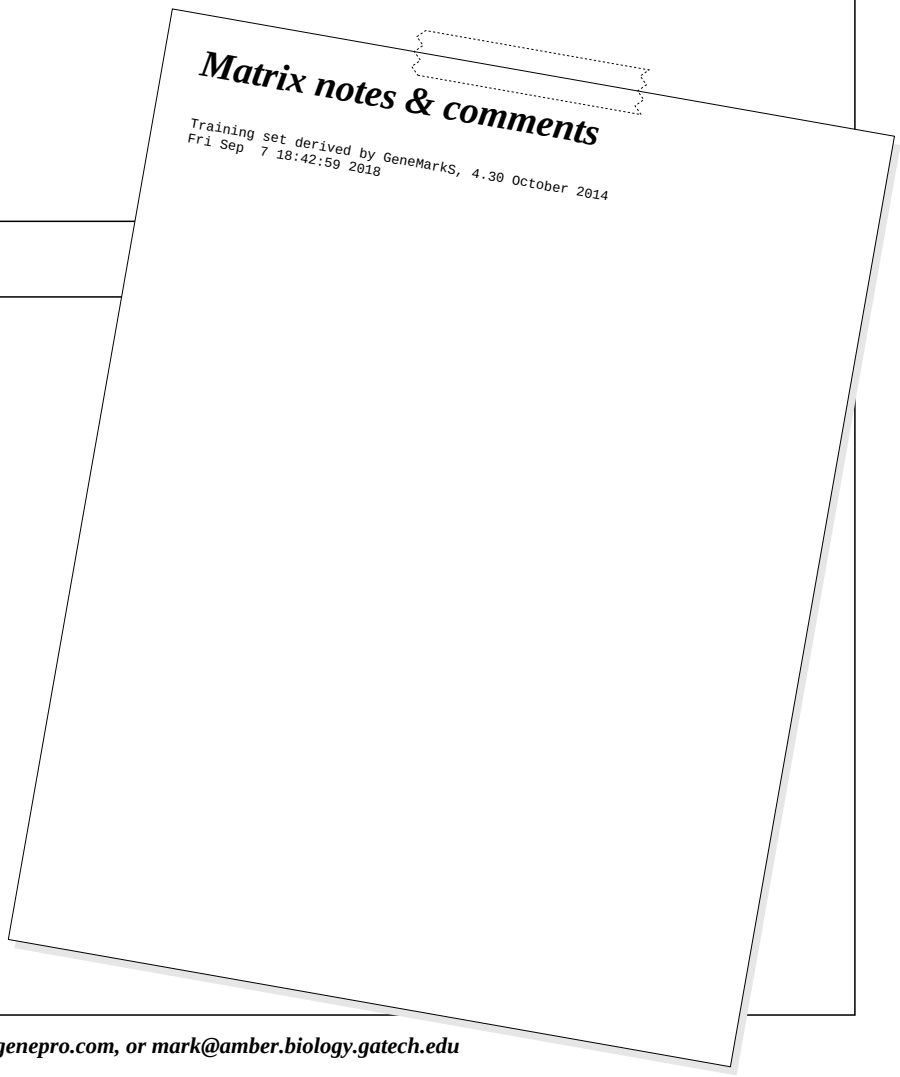
PROGRAM INFORMATION

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Sequence         :  Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1
Analysis Date    :  3/29/24 at 7:01:04
Pages            :  27
Sequence Length  :  53620 bp
GC Content       :   58.99%

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

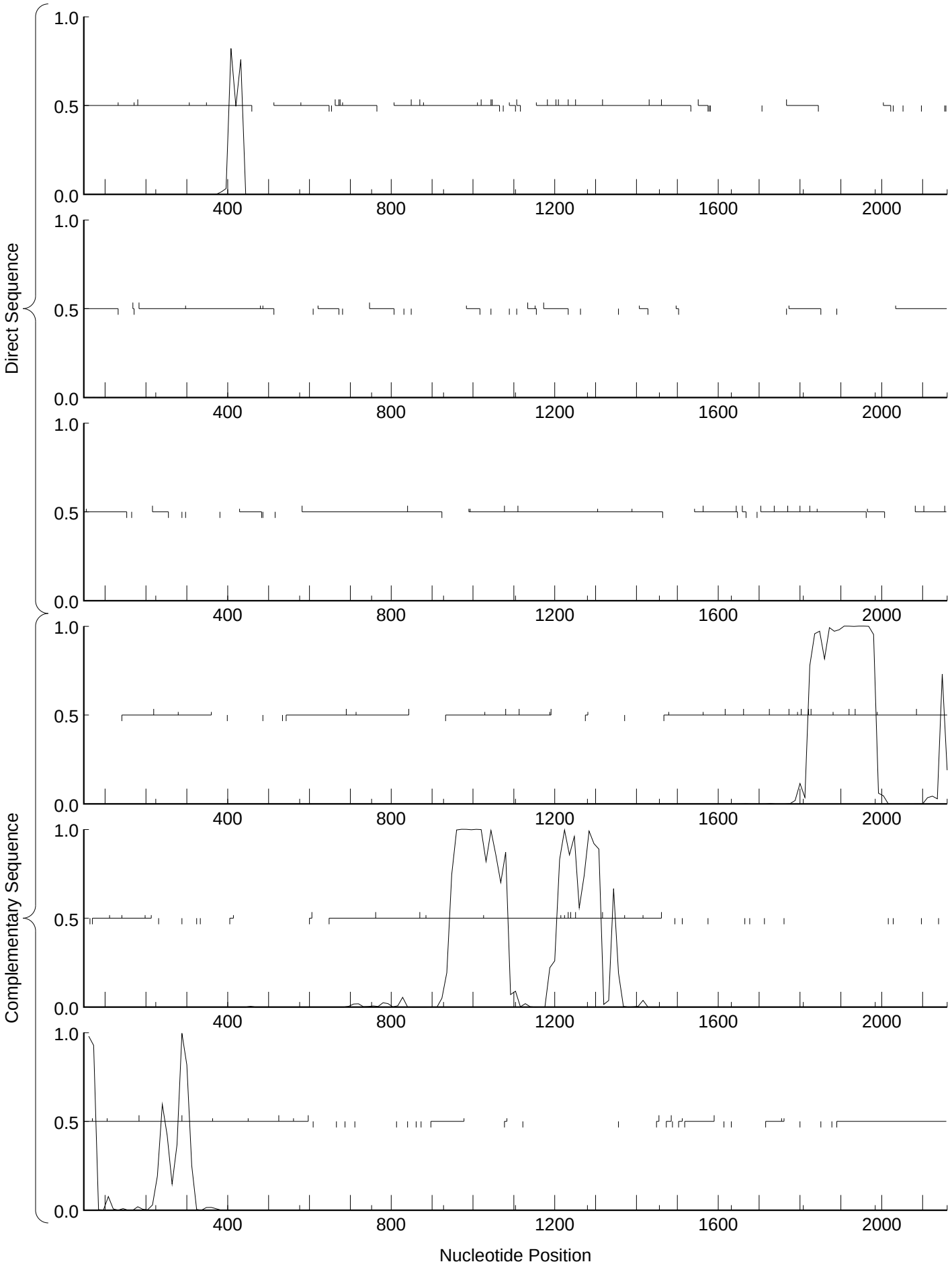
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Author     :  -

Order      :  4
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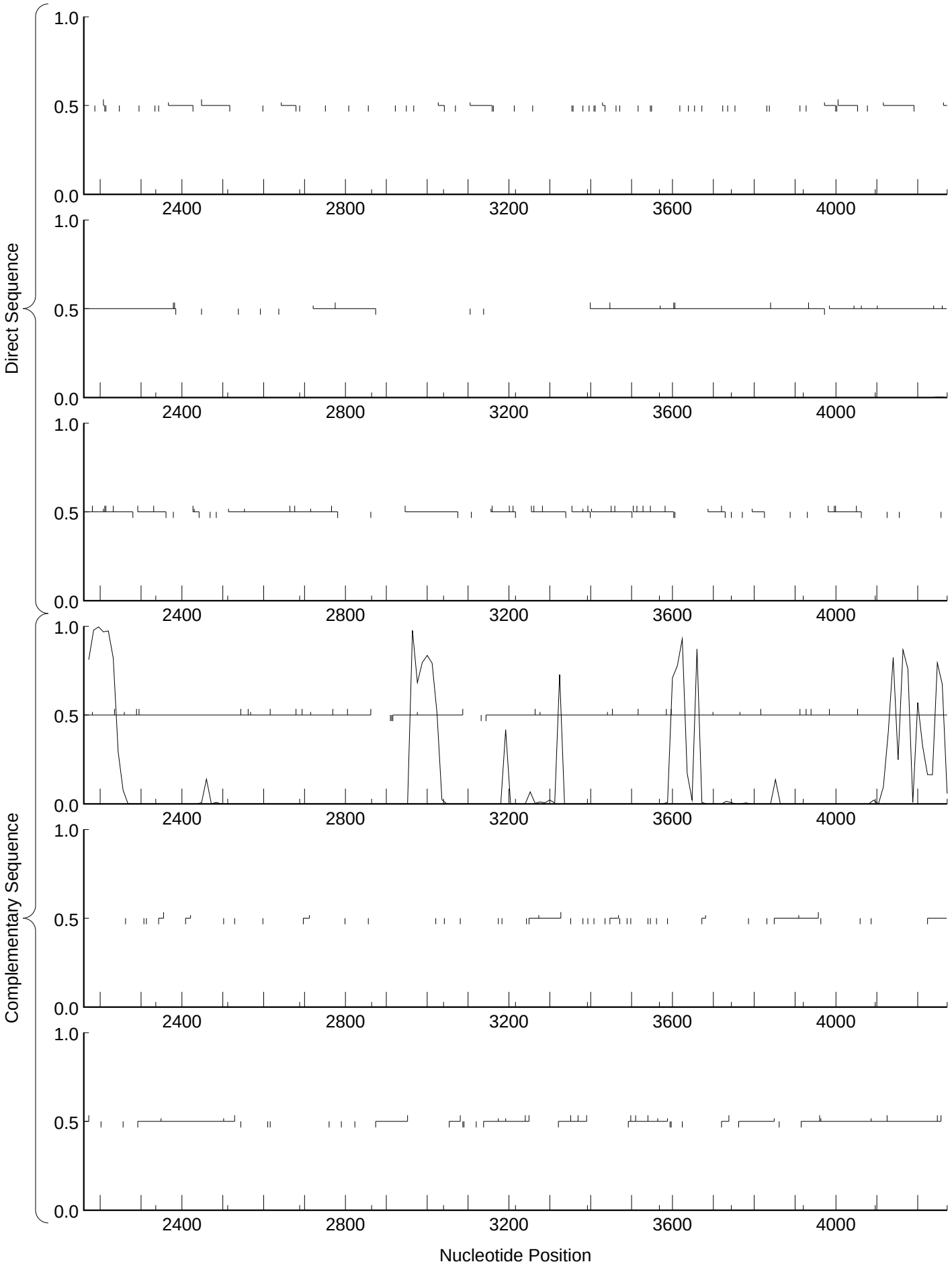

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

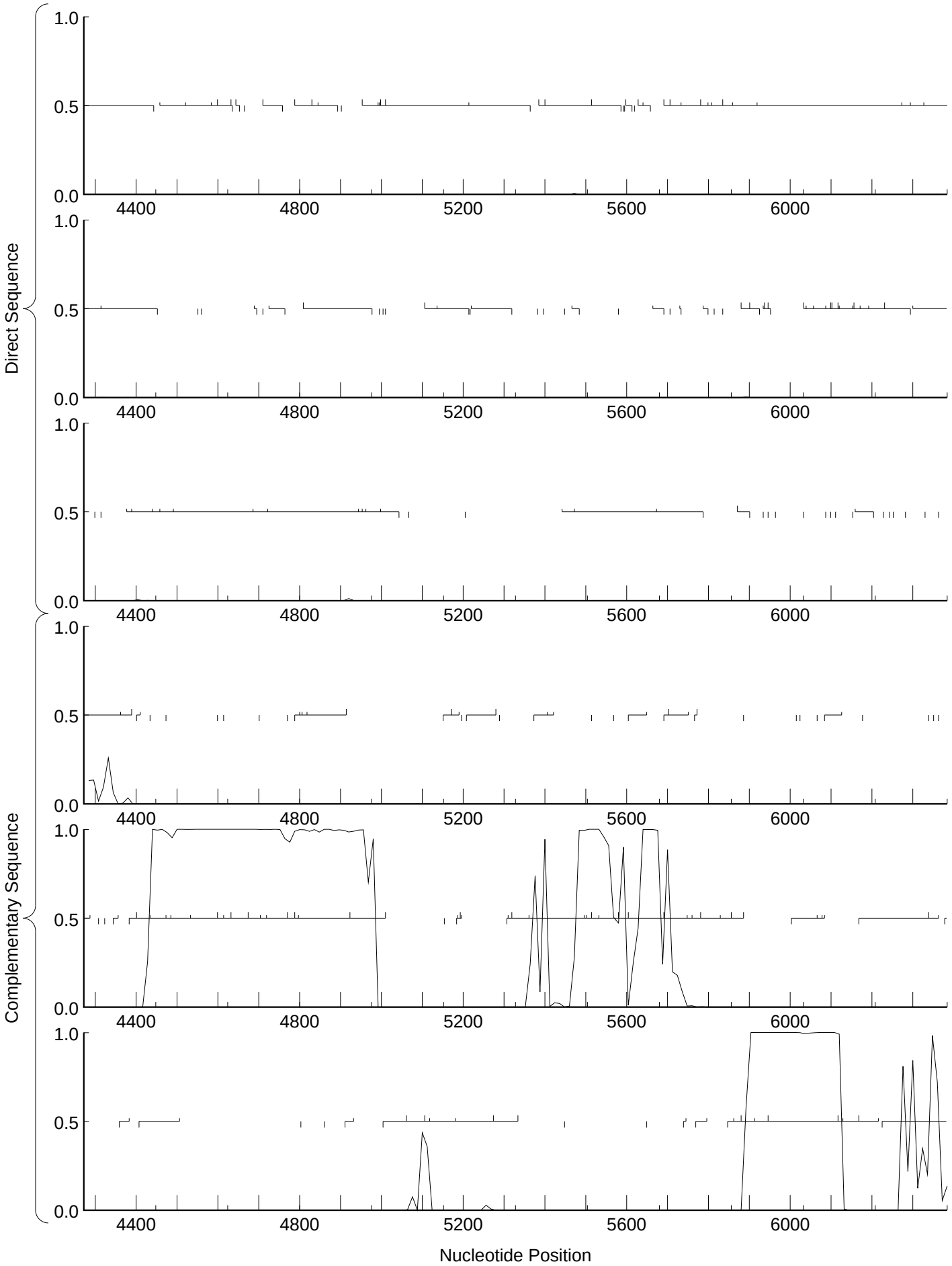
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 2/27



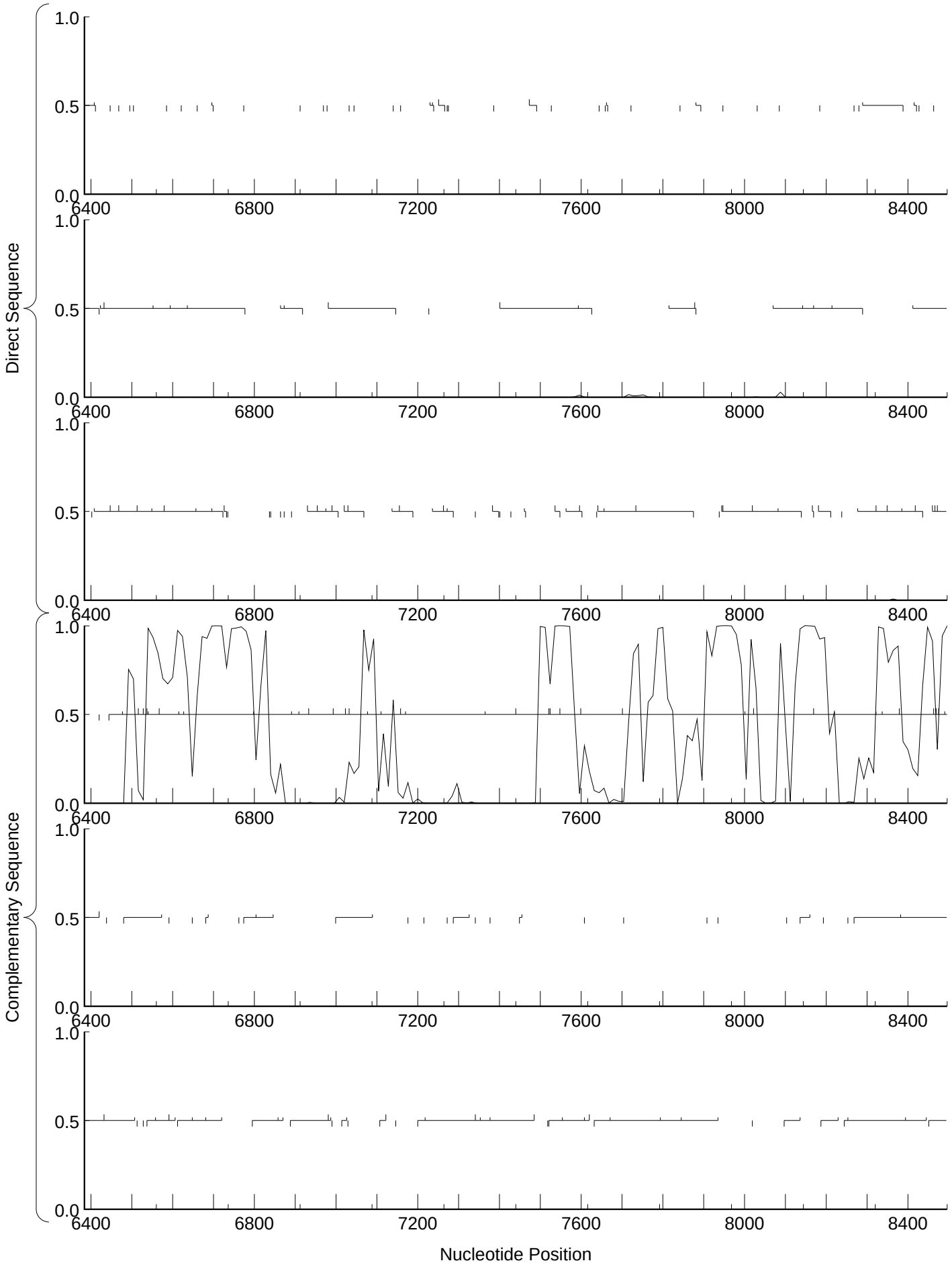
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 3/27



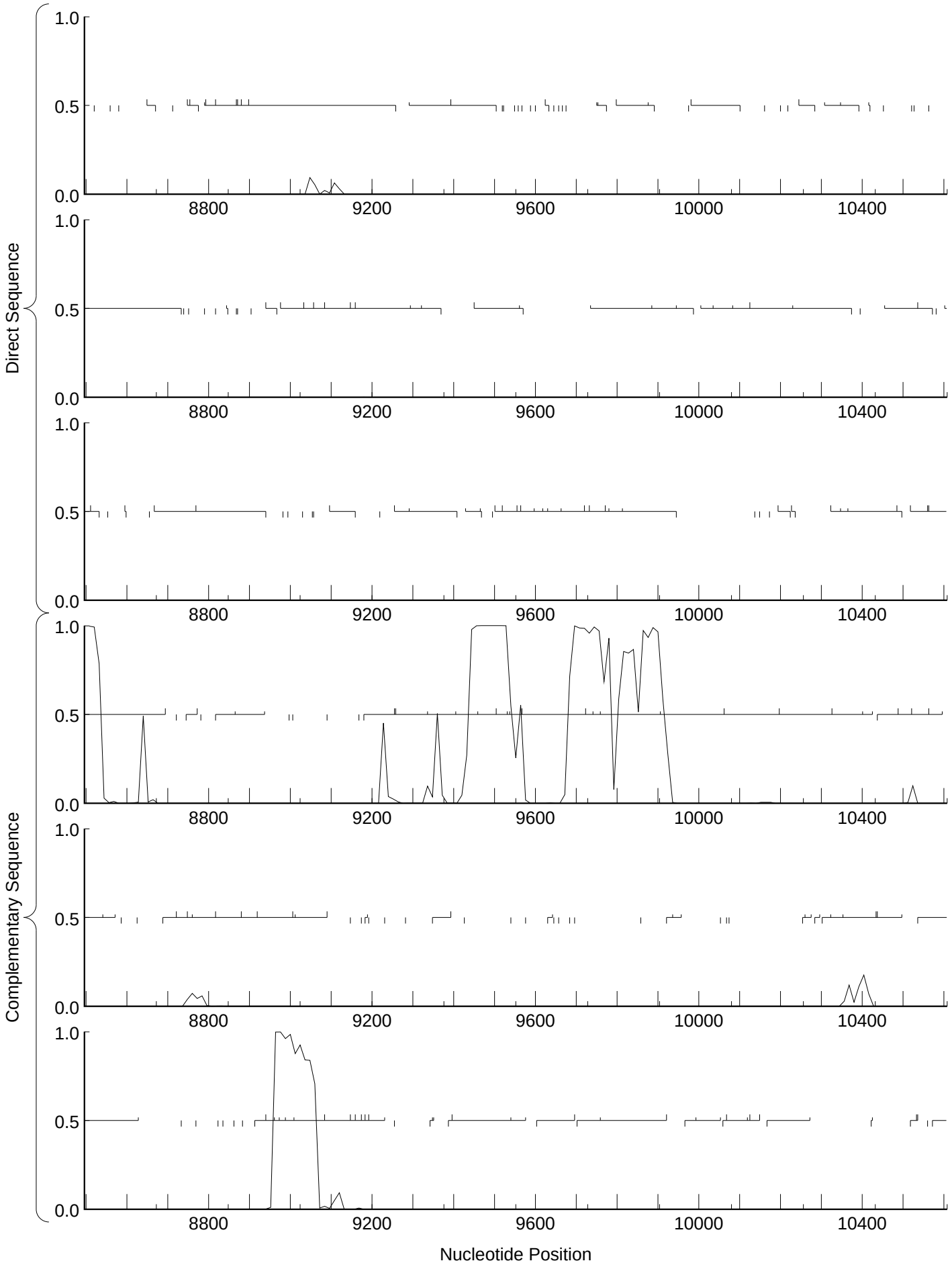
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 4/27



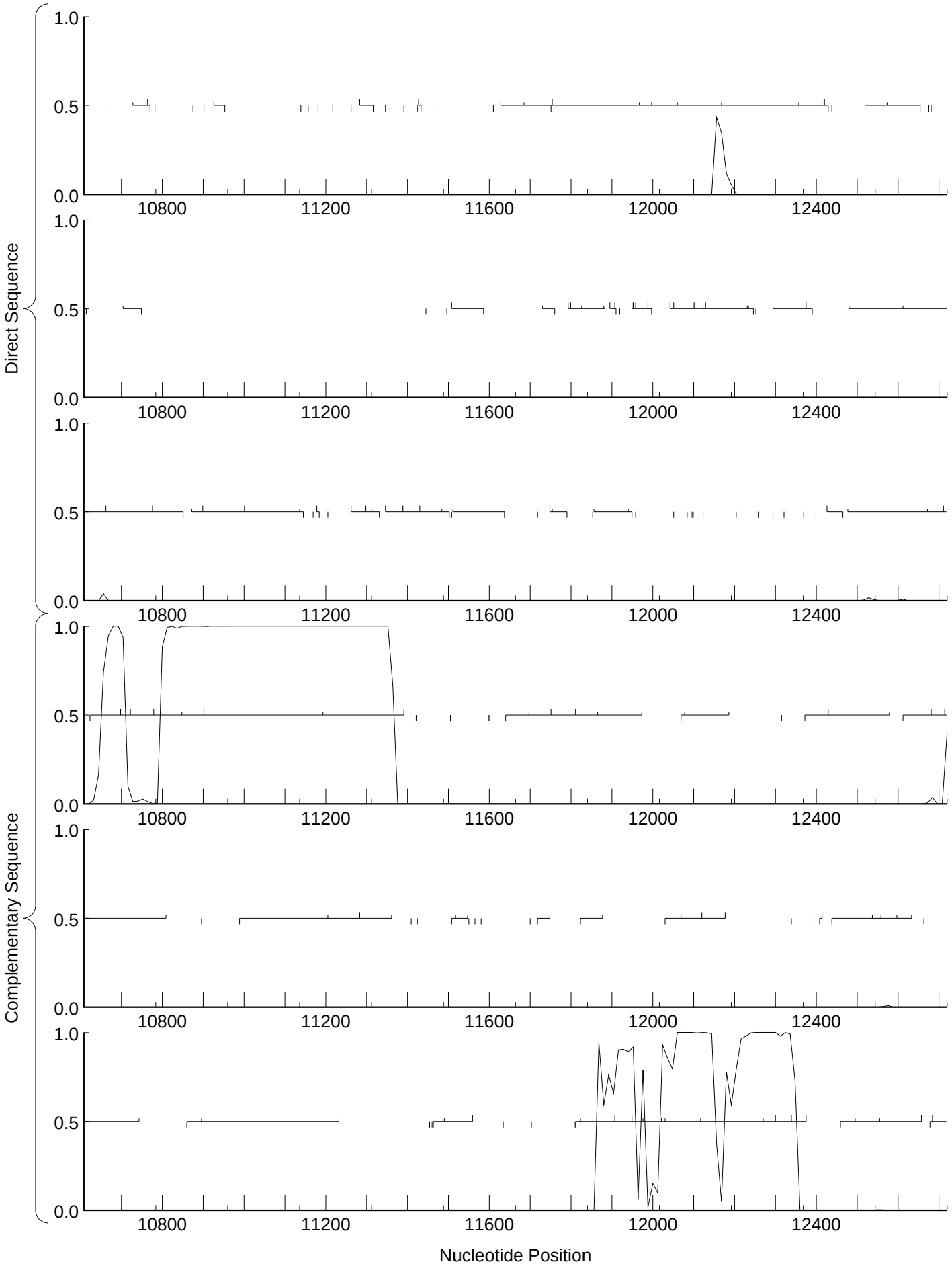
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 5/27



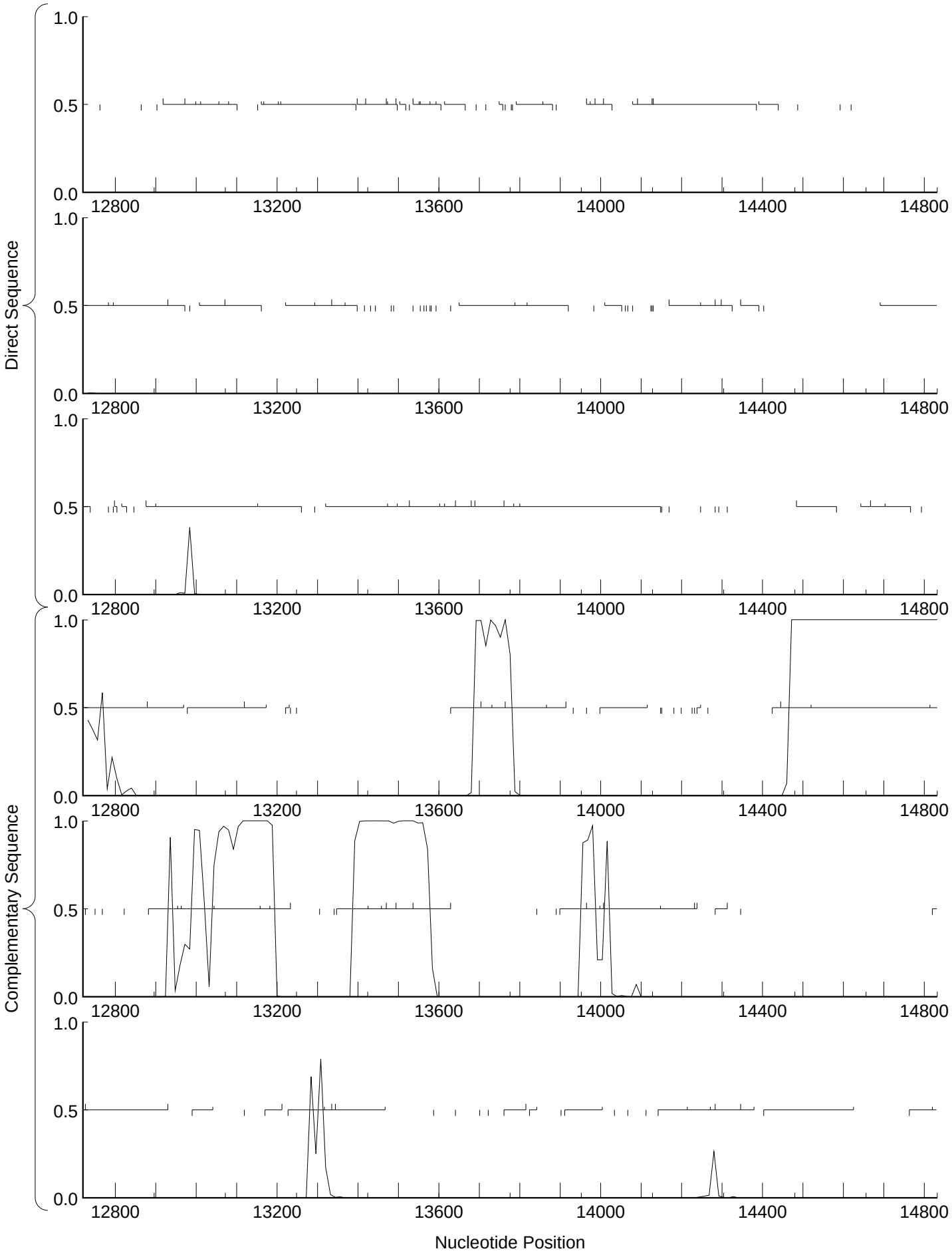
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 6/27



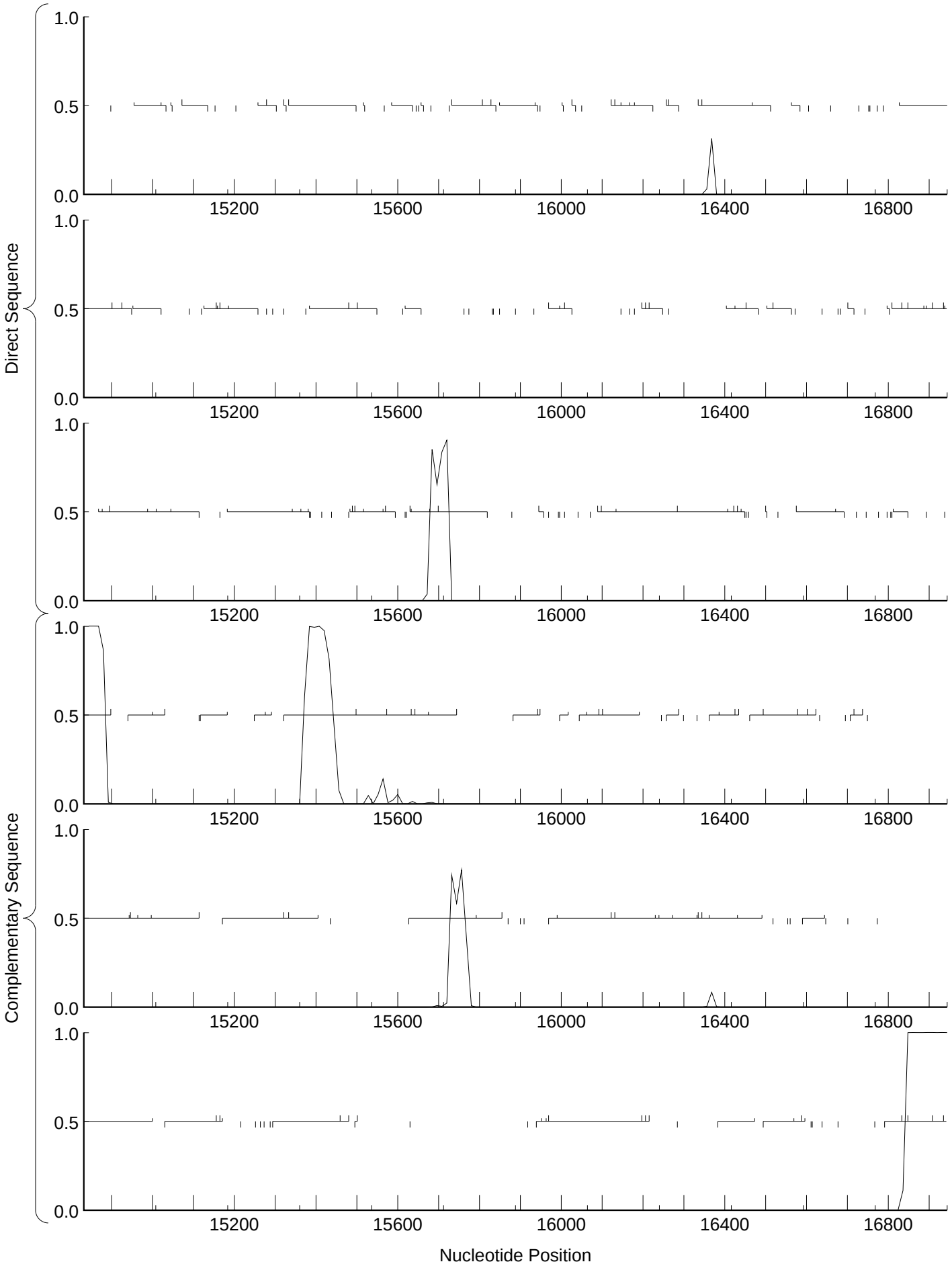
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 7/27



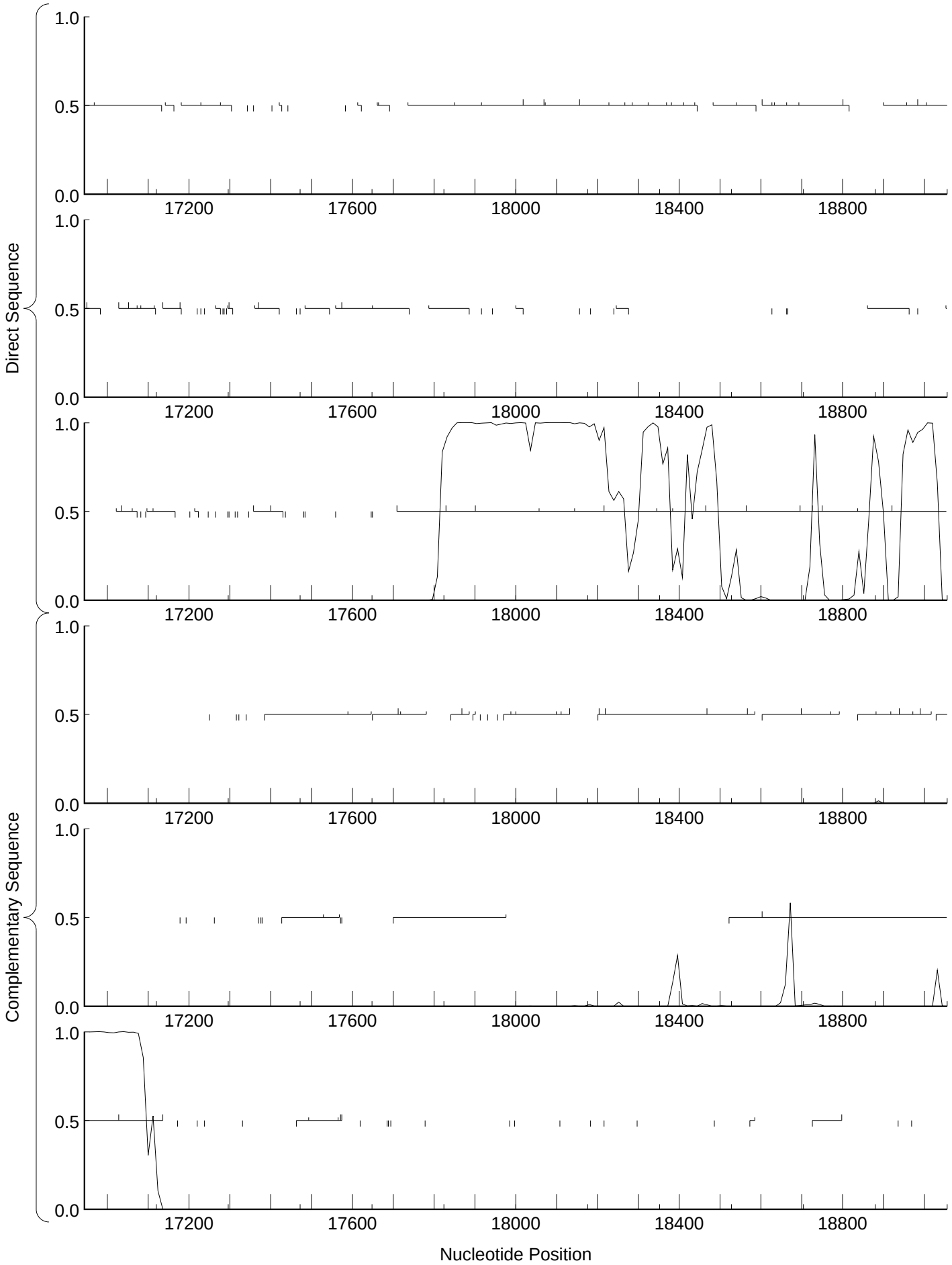
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 8/27



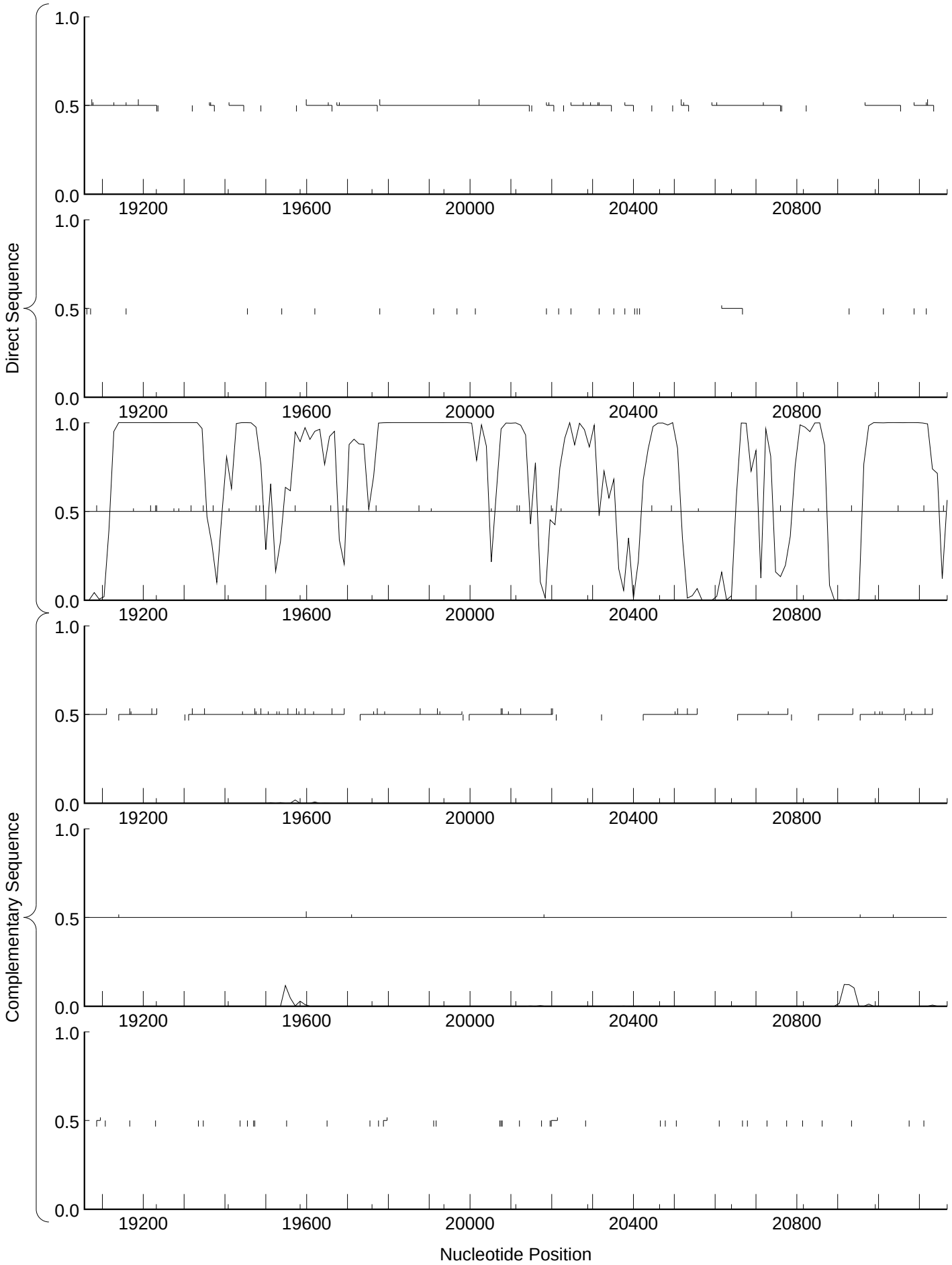
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 9/27



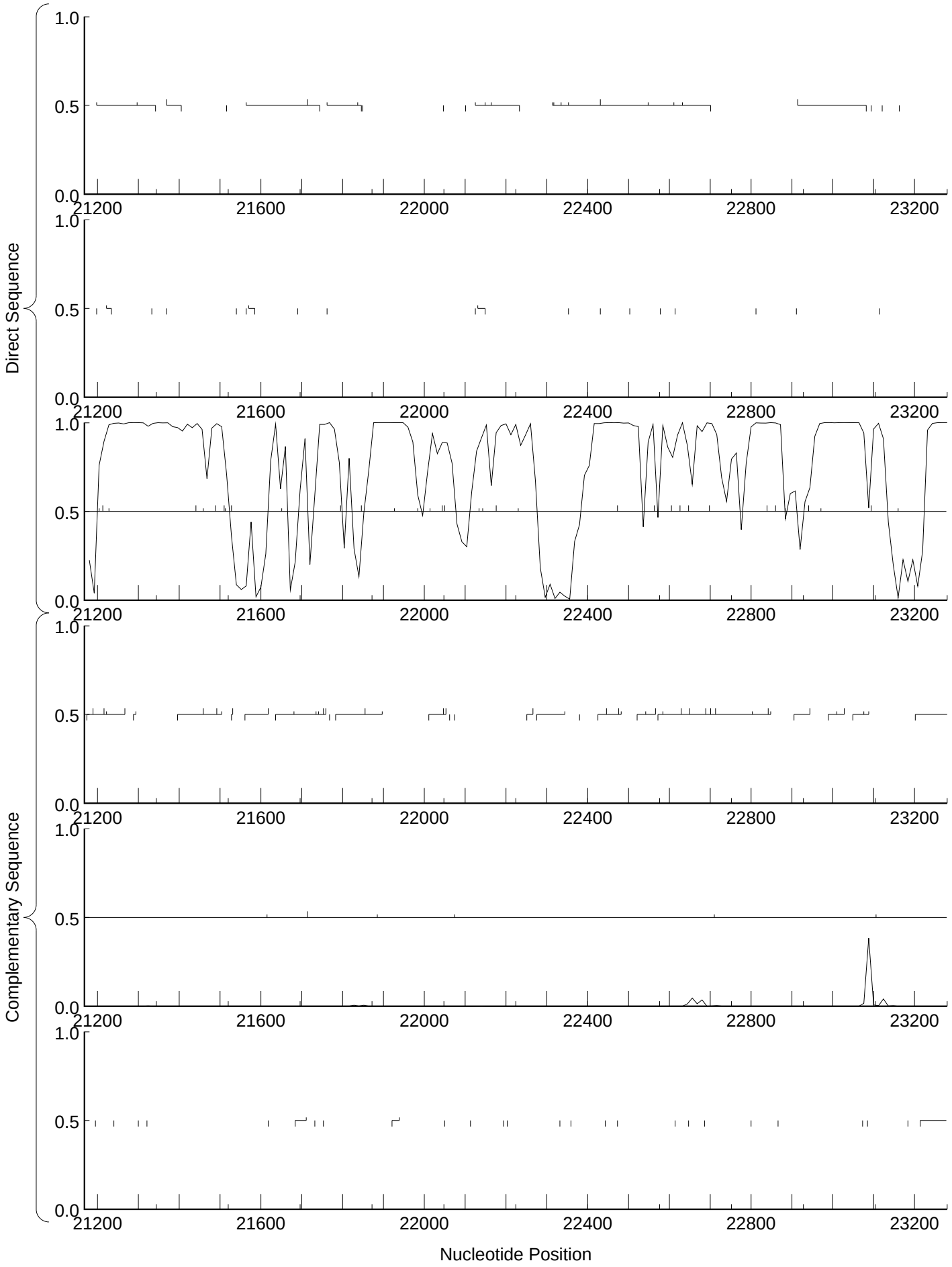
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 10/27



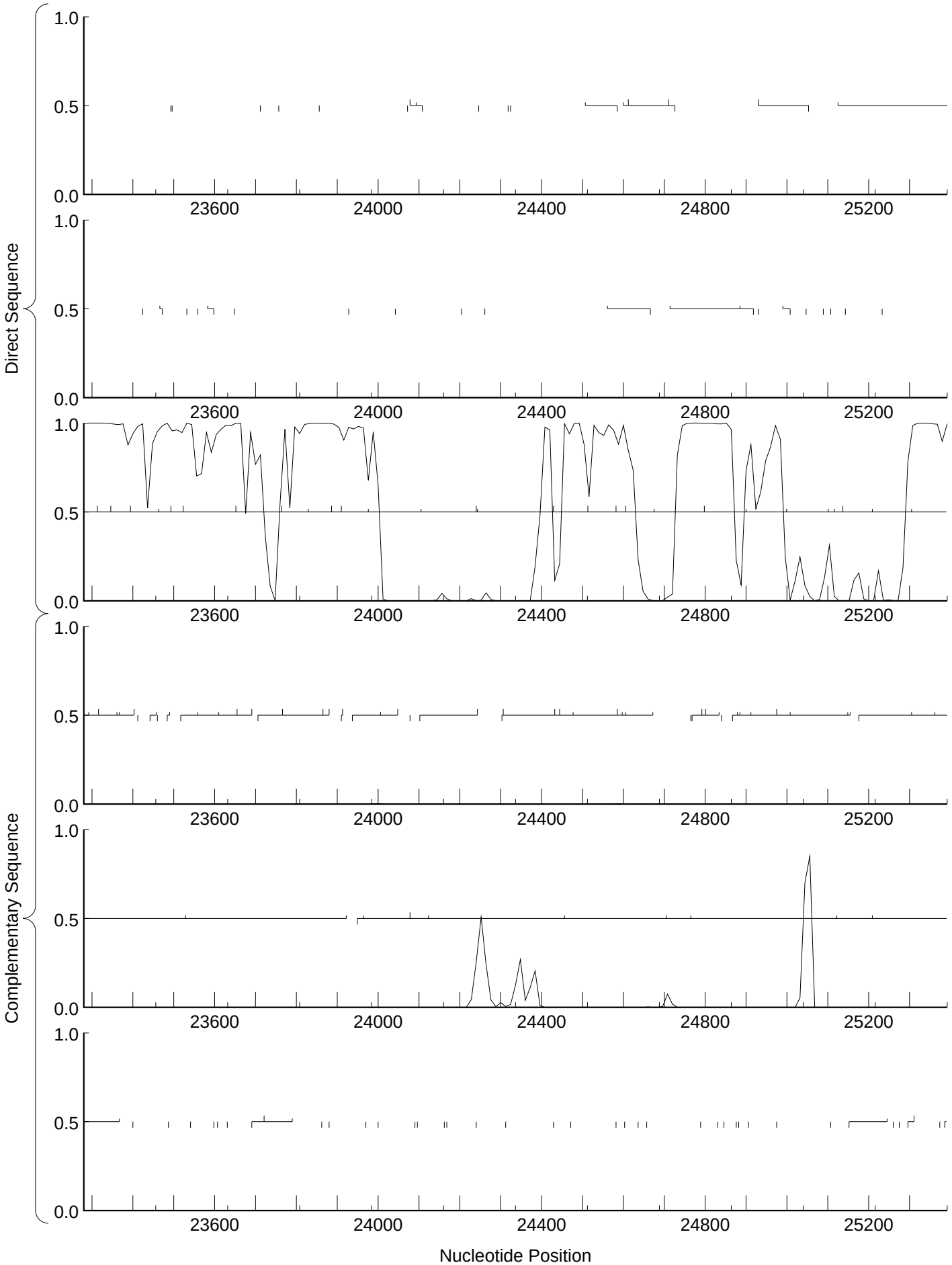
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 11/27



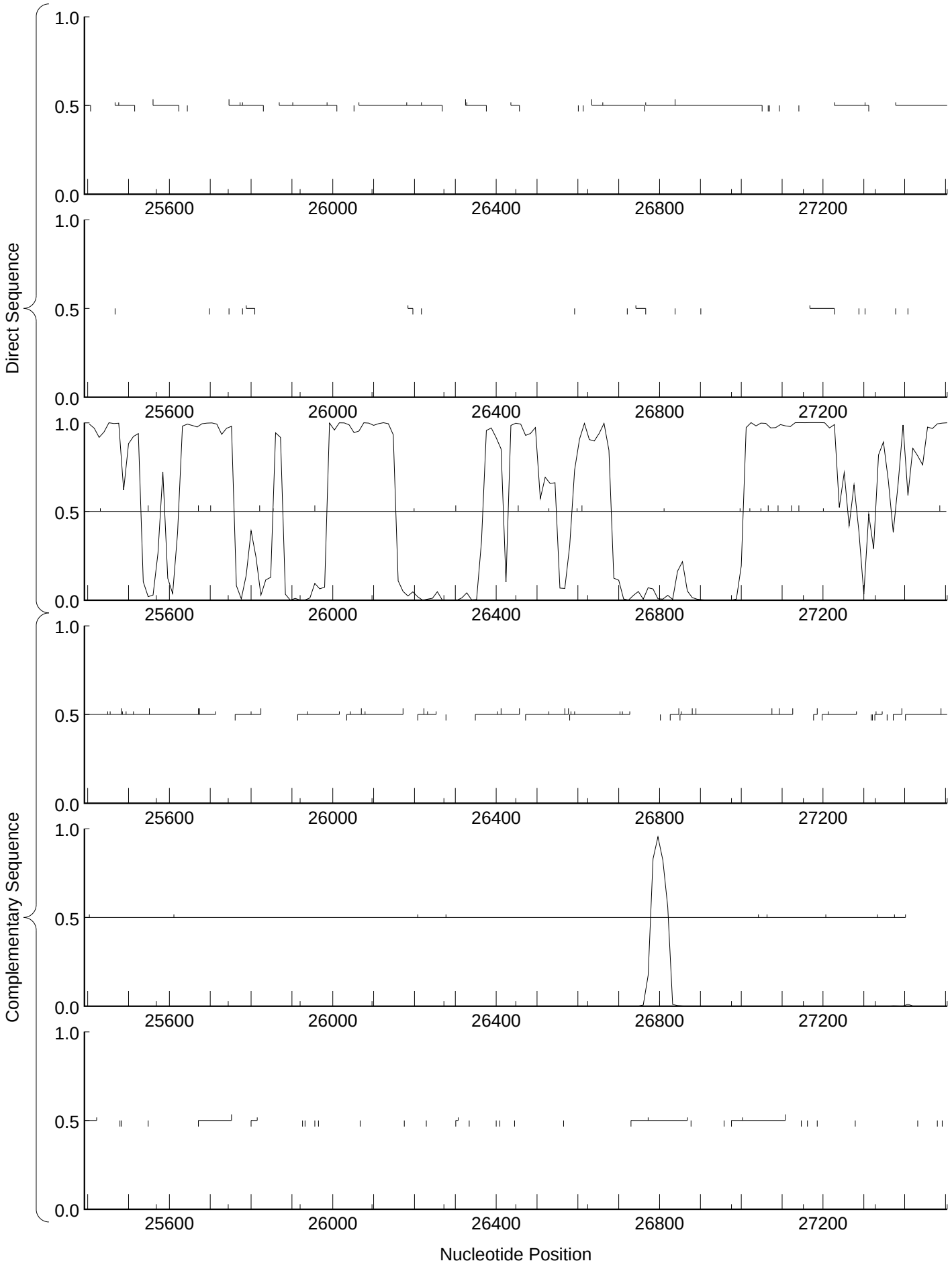
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 12/27



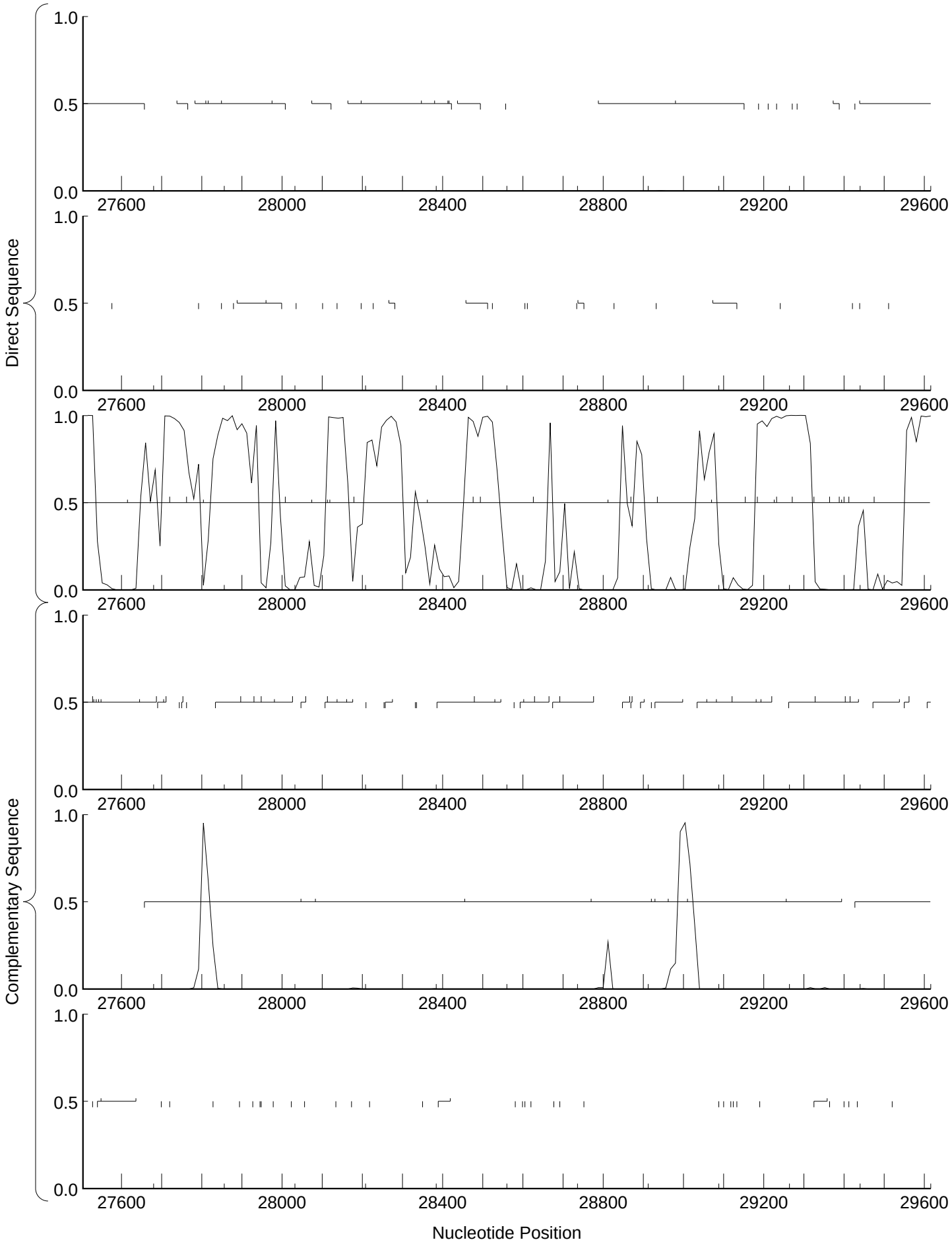
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 13/27



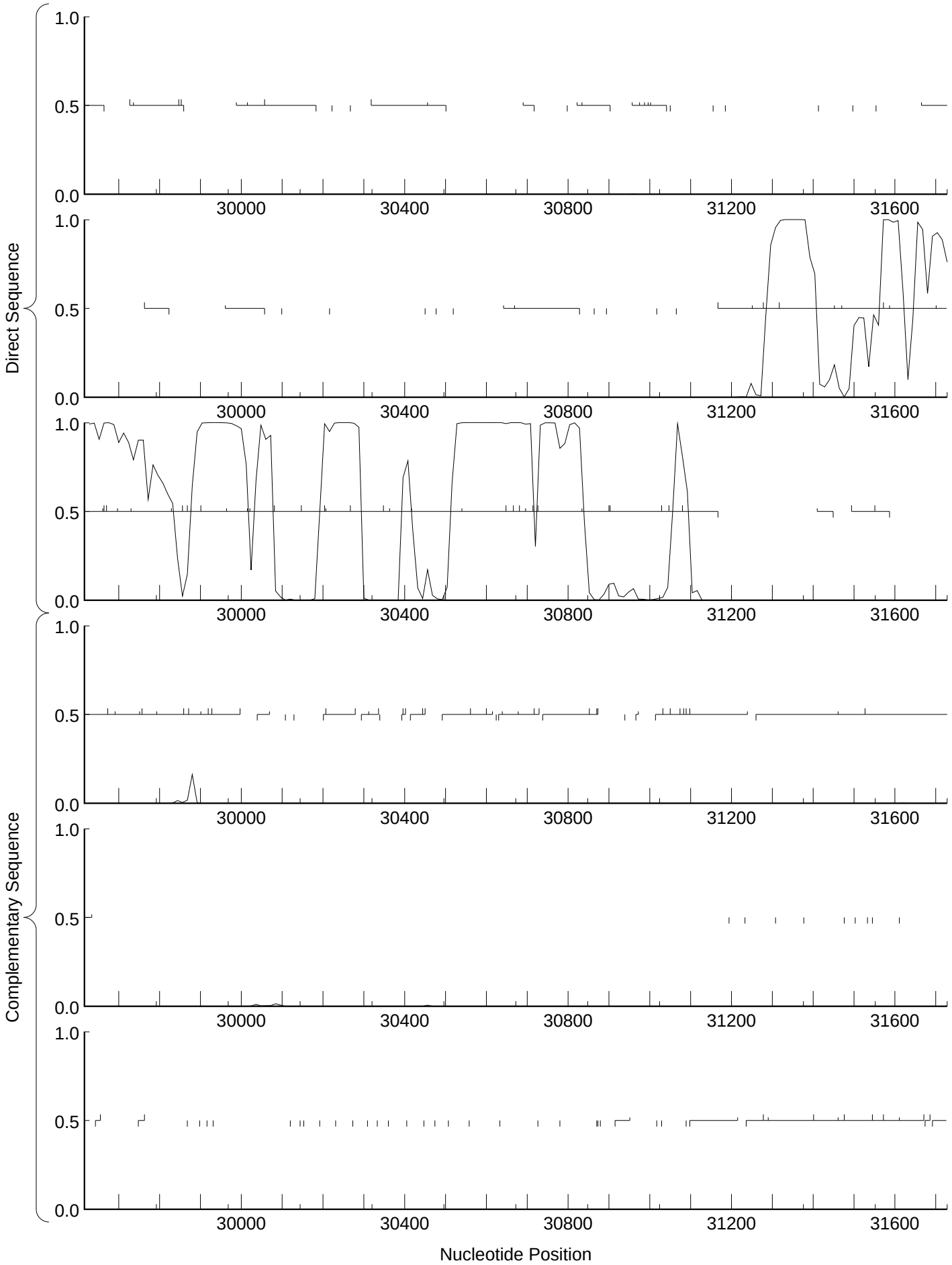
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 14/27



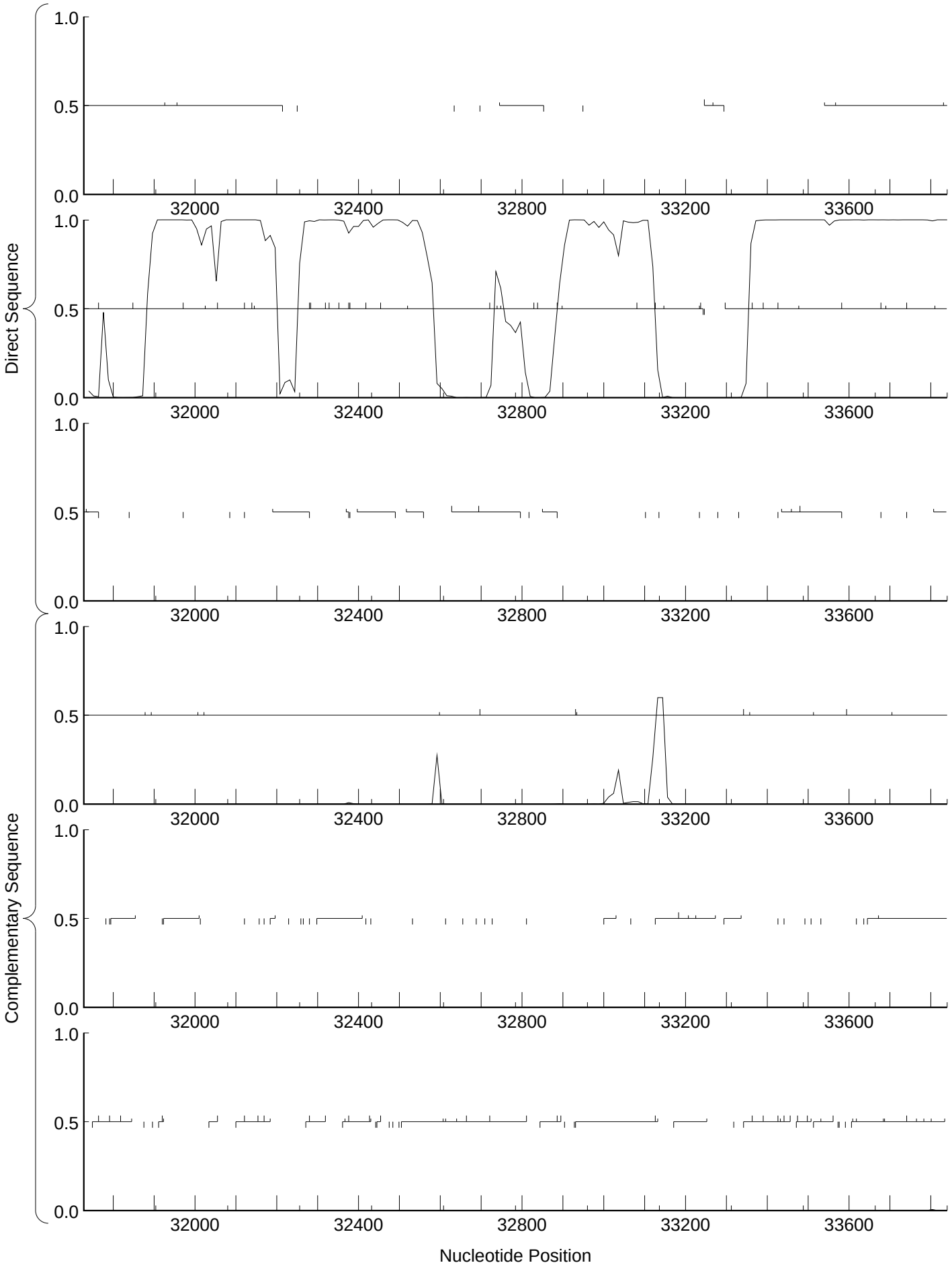
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 15/27

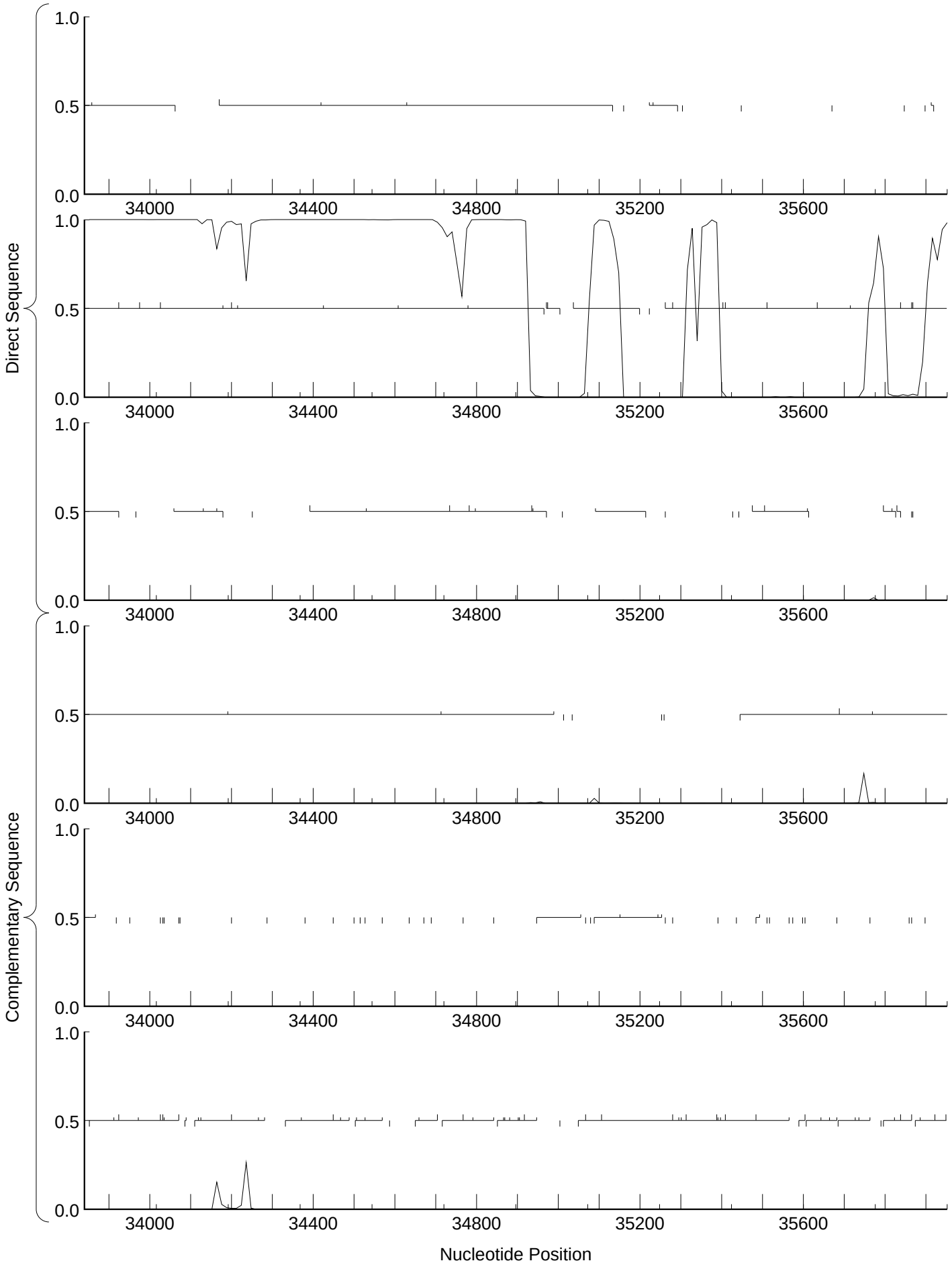


Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 16/27

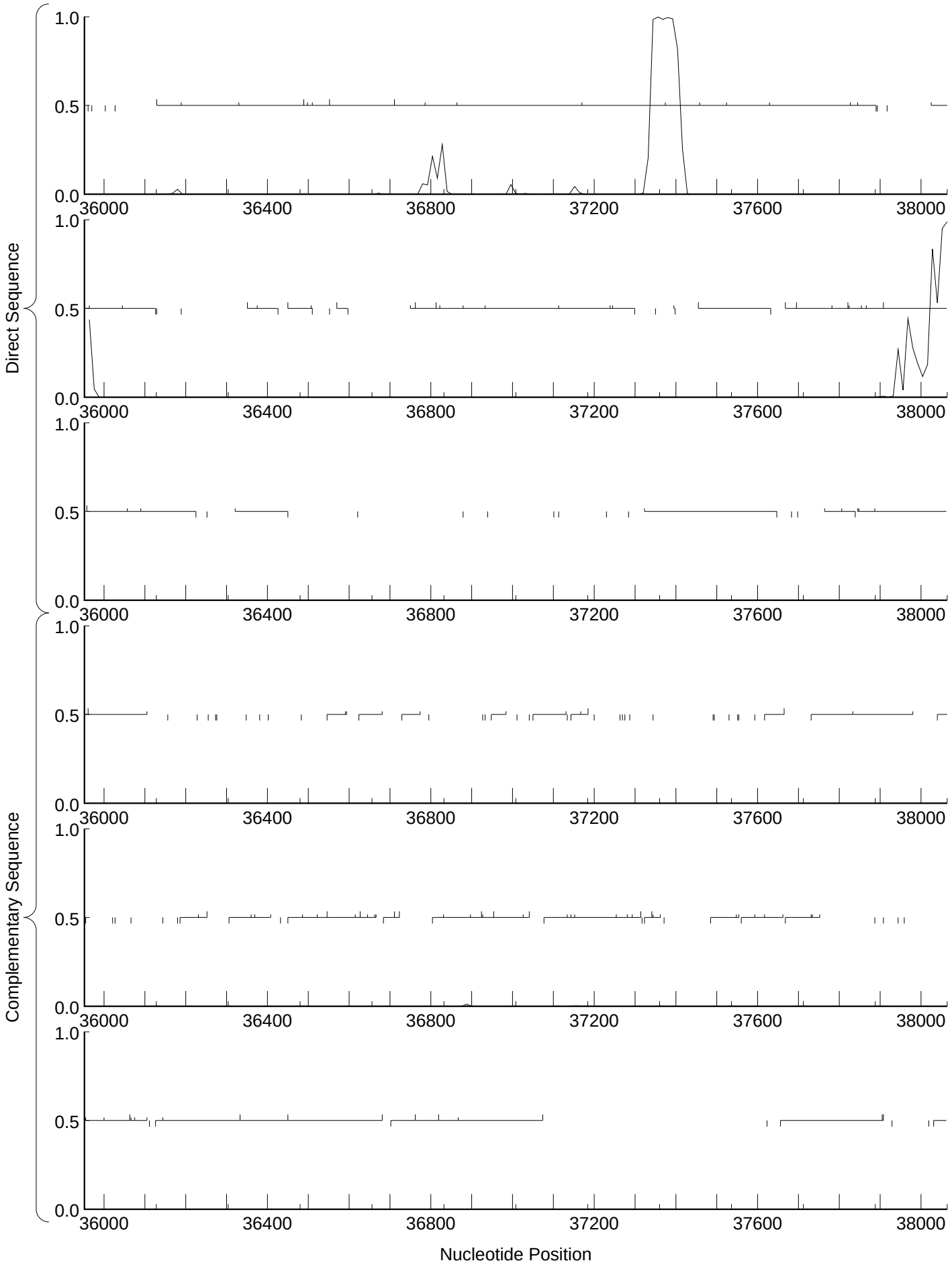


Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 17/27

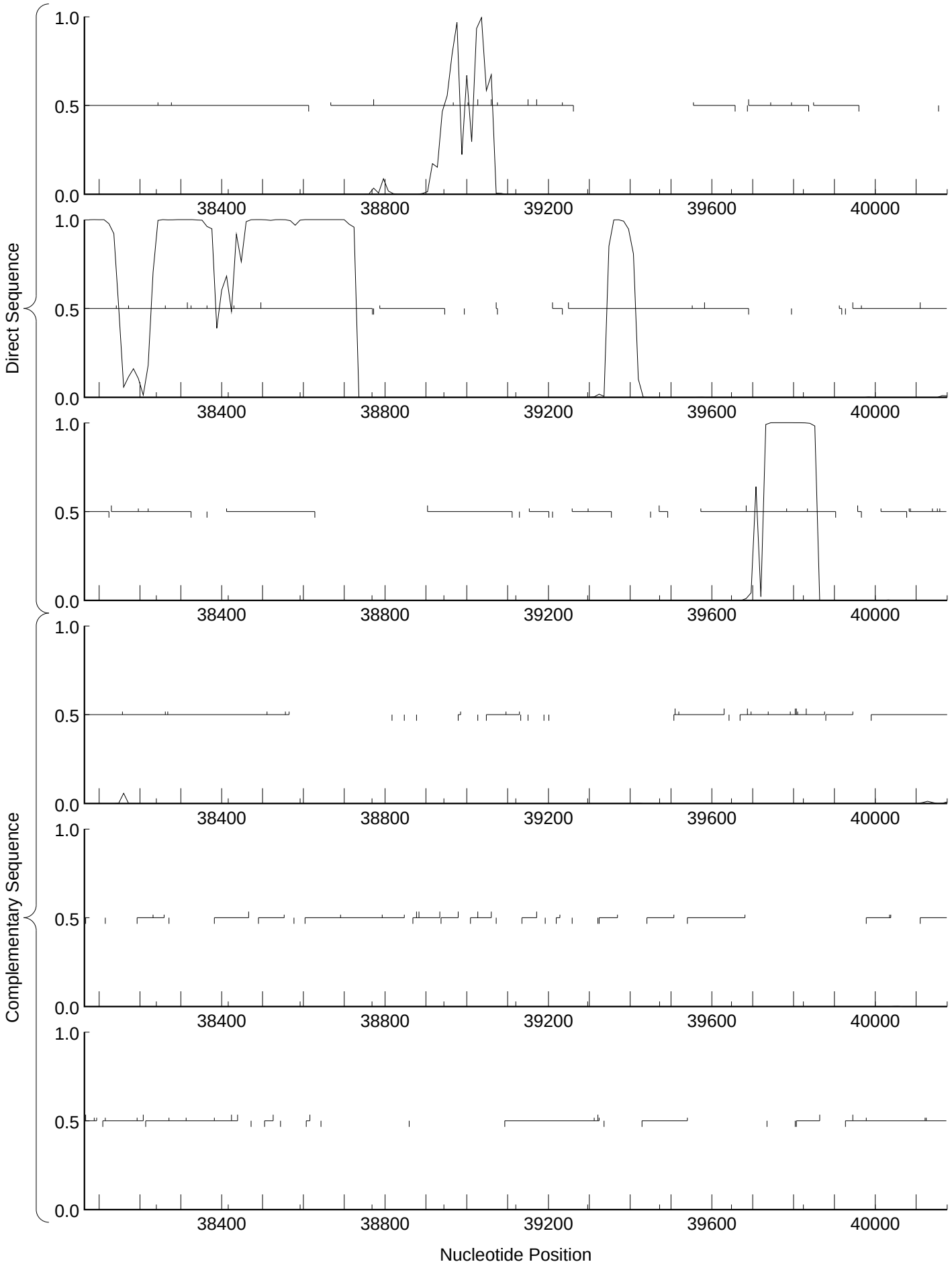



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

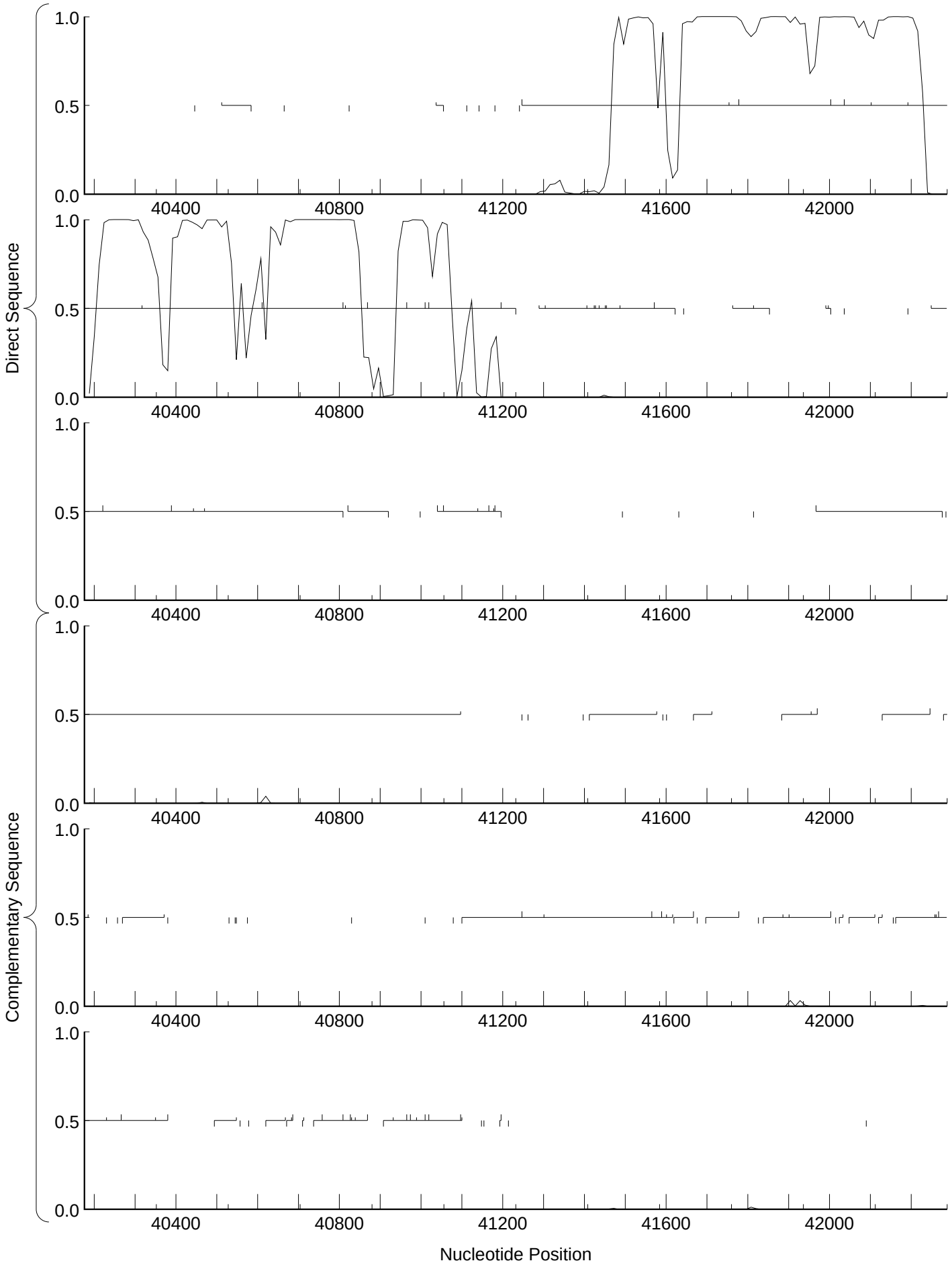
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 19/27



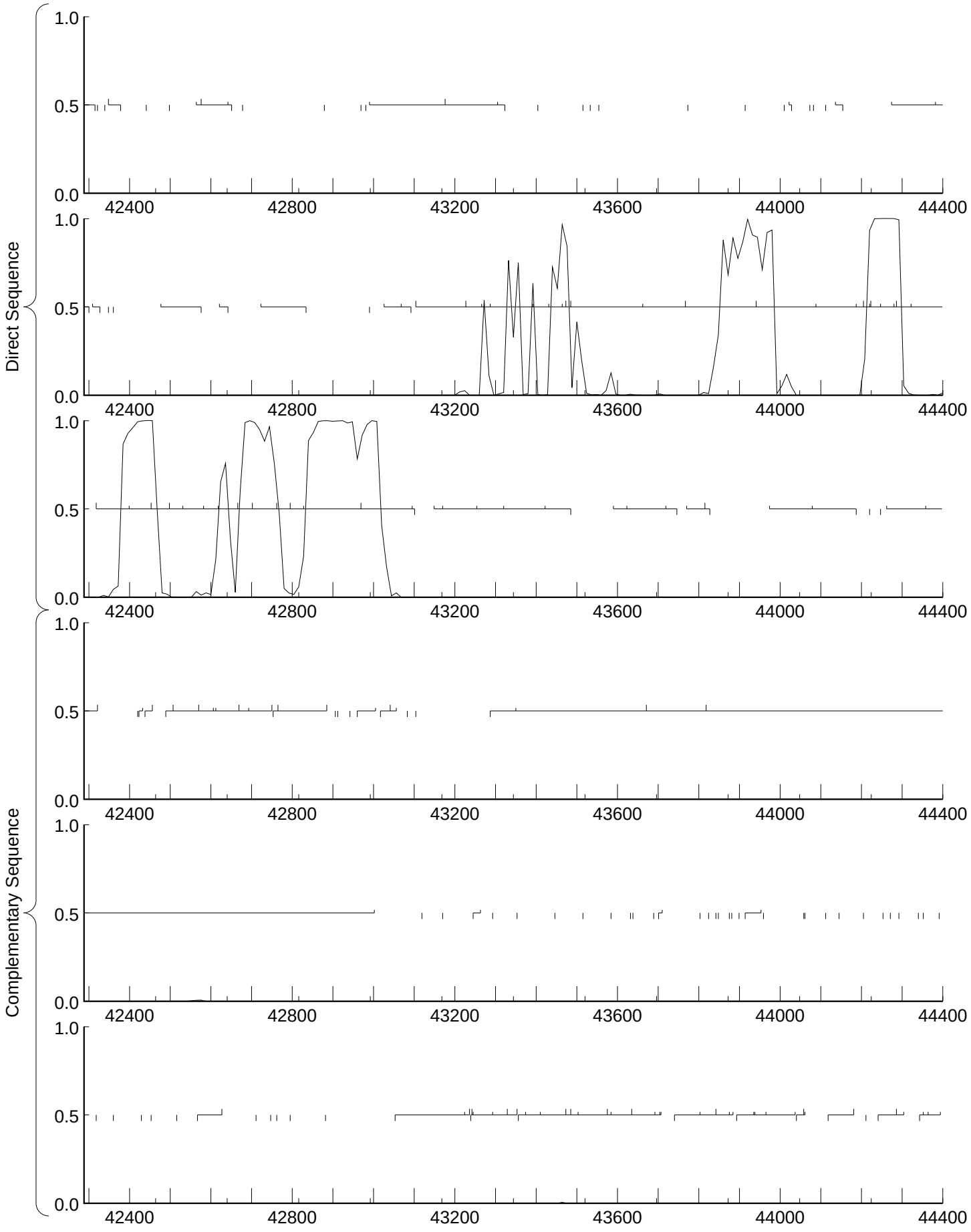
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 20/27



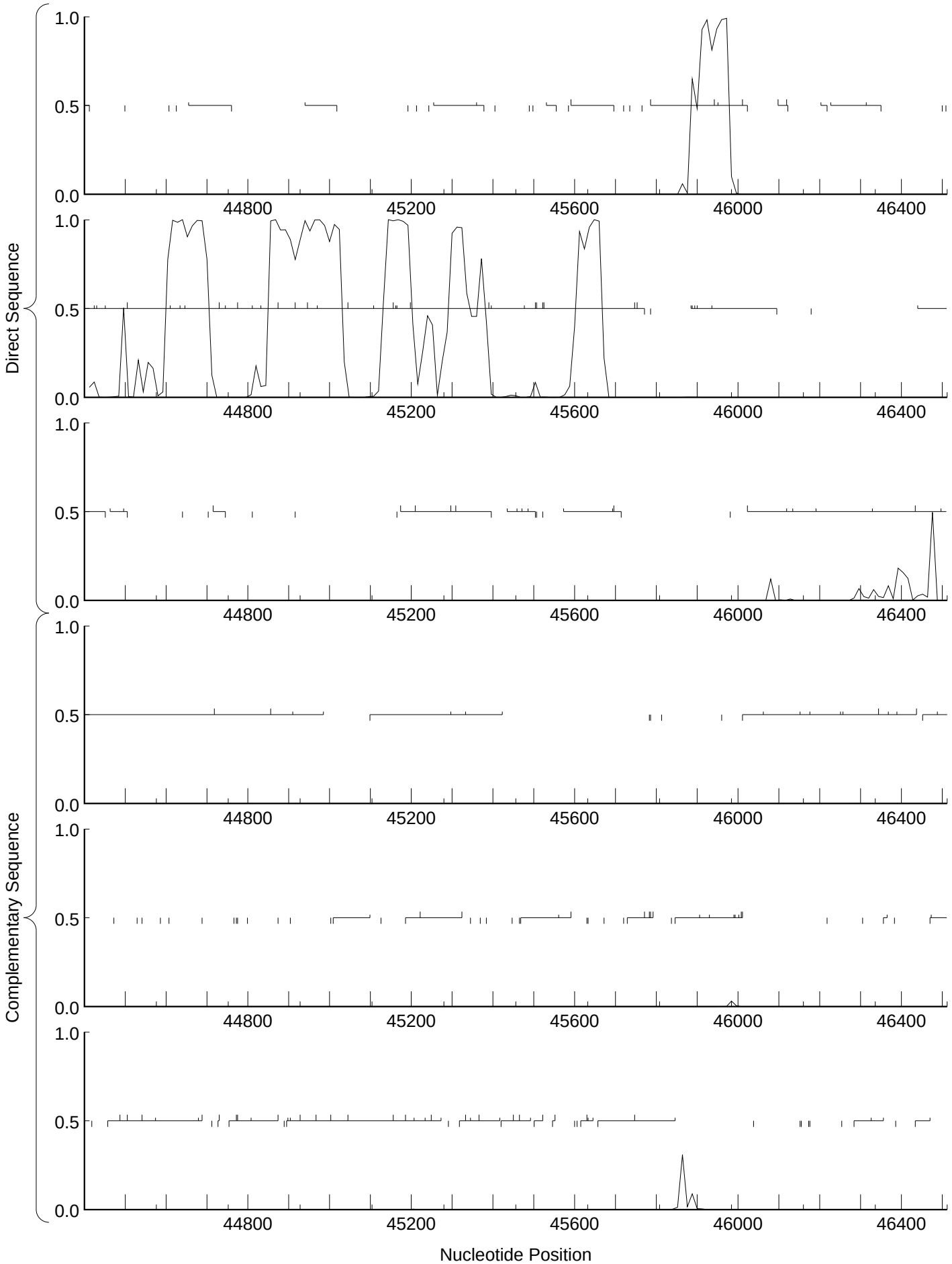
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 21/27



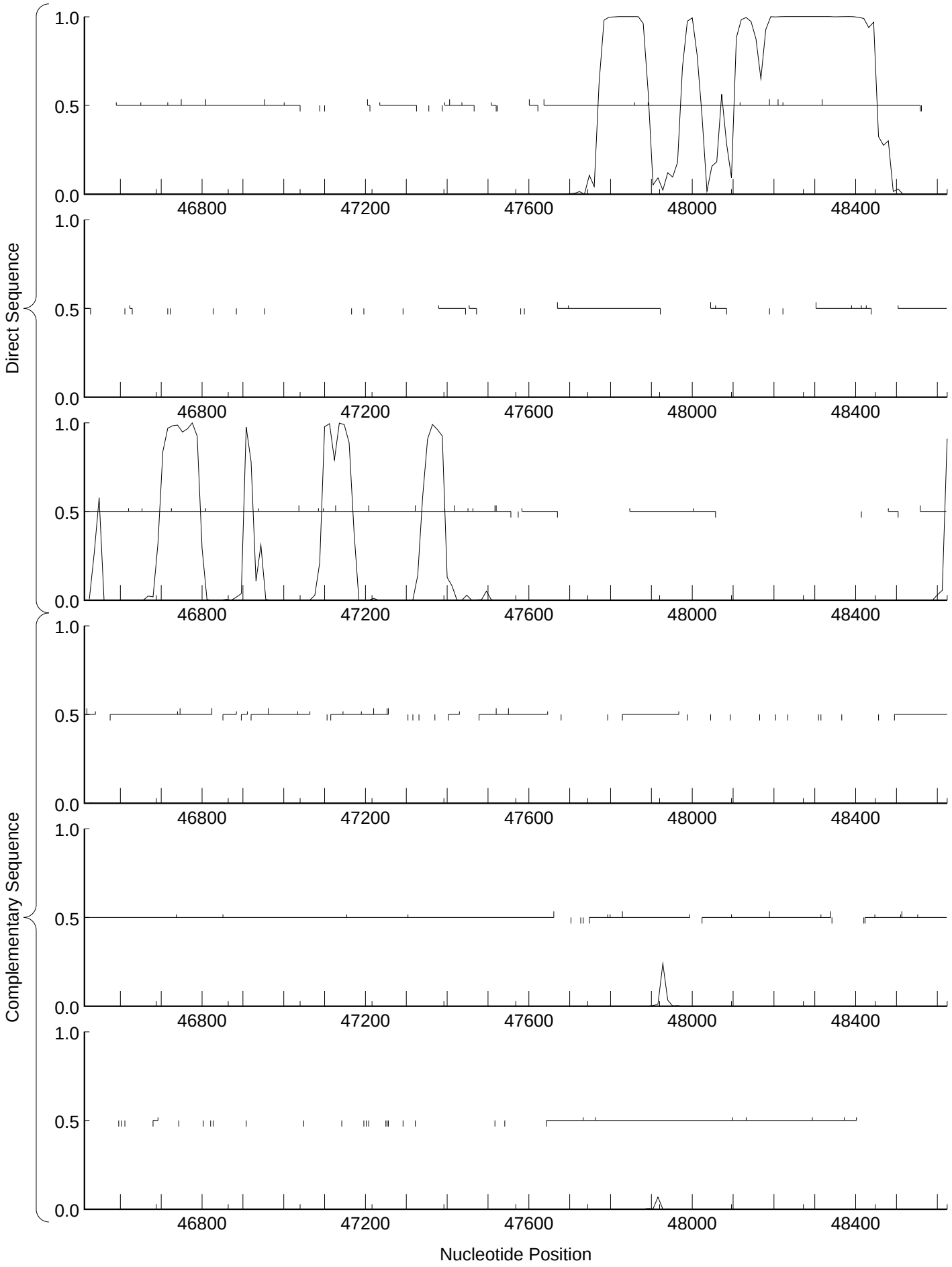
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 22/27



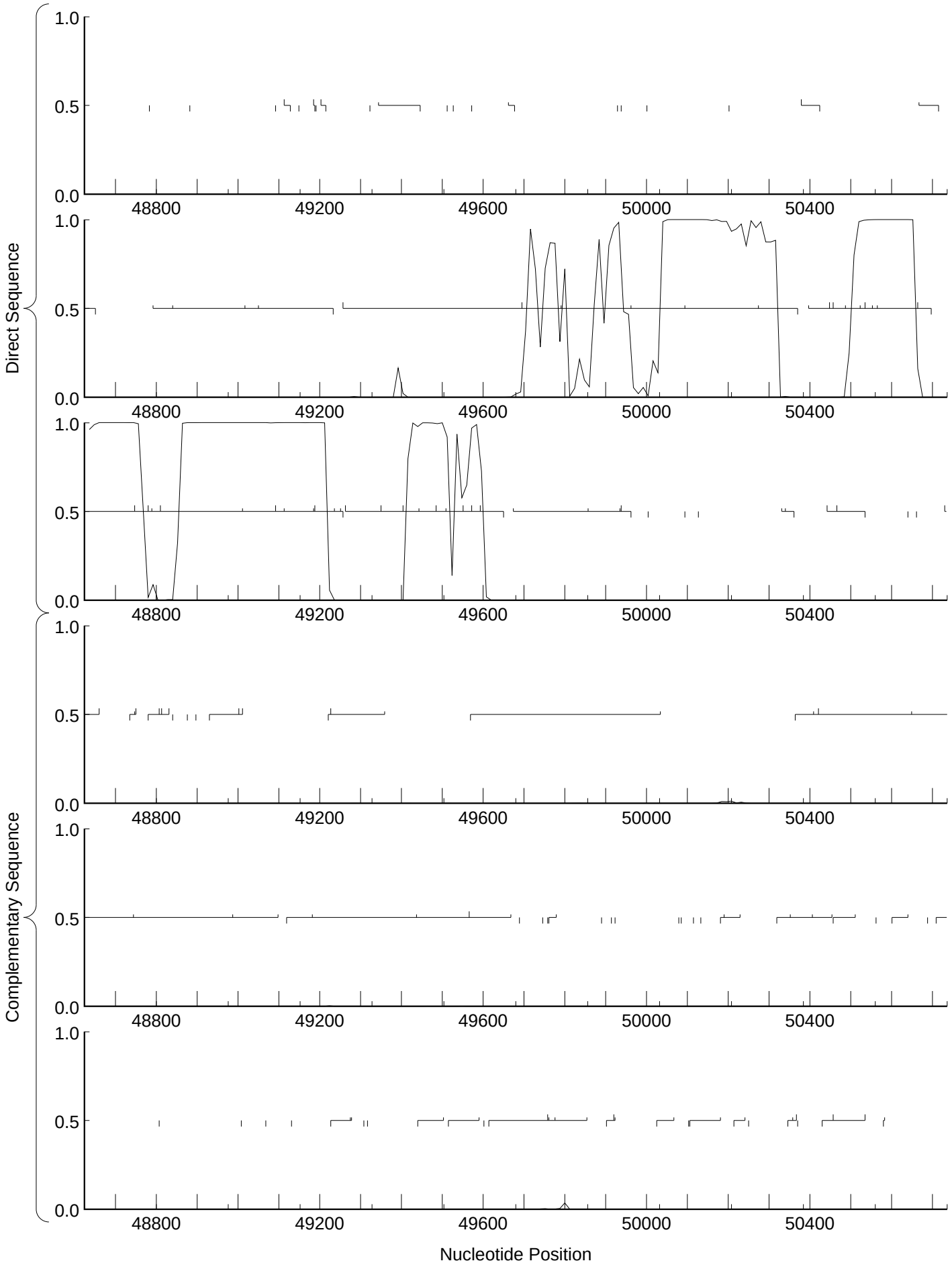
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 23/27



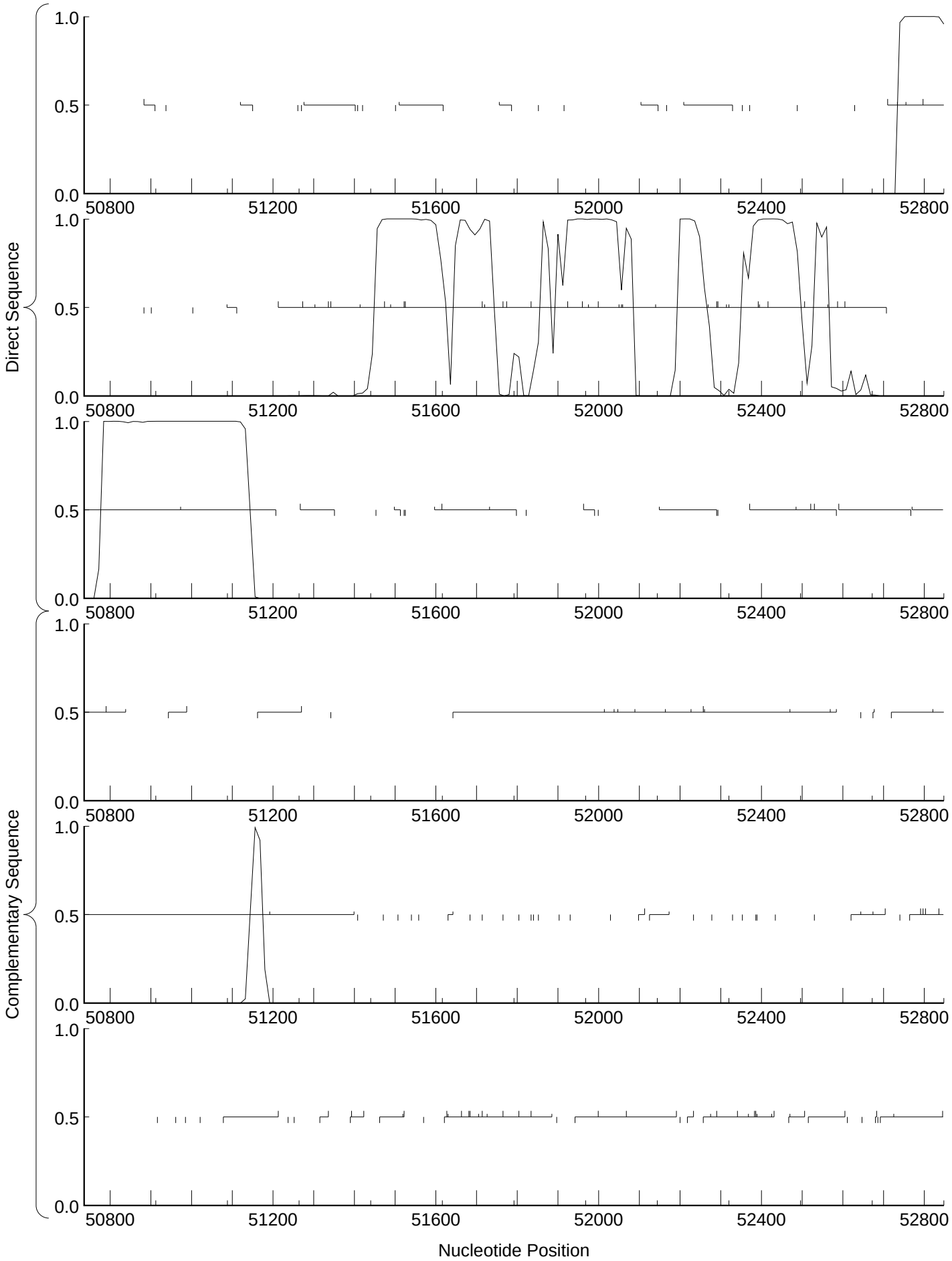
Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 24/27



Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 25/27



Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 26/27



Microbacterium phage Pabst complete sequence, 53620 bp, circularly permuted, Cluster EK1, Order 4, Window 96, Step 12, 27/27

