GeneMark

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

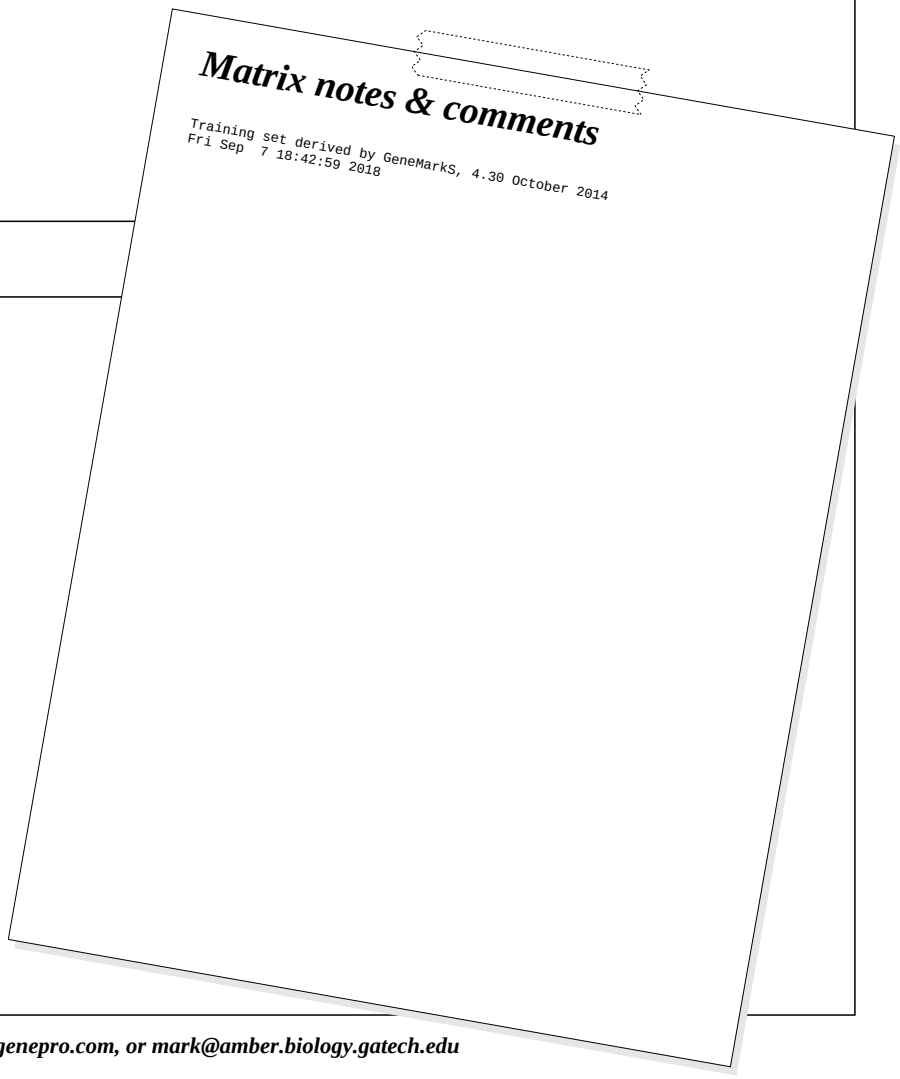
PROGRAM INFORMATION

```
Sequence         :  Microbacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6
Analysis Date    :  3/28/24 at 9:25:06
Pages            :  21
Sequence Length  :  41099 bp
GC Content       :   62.77%

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,
```



MATRIX INFORMATION

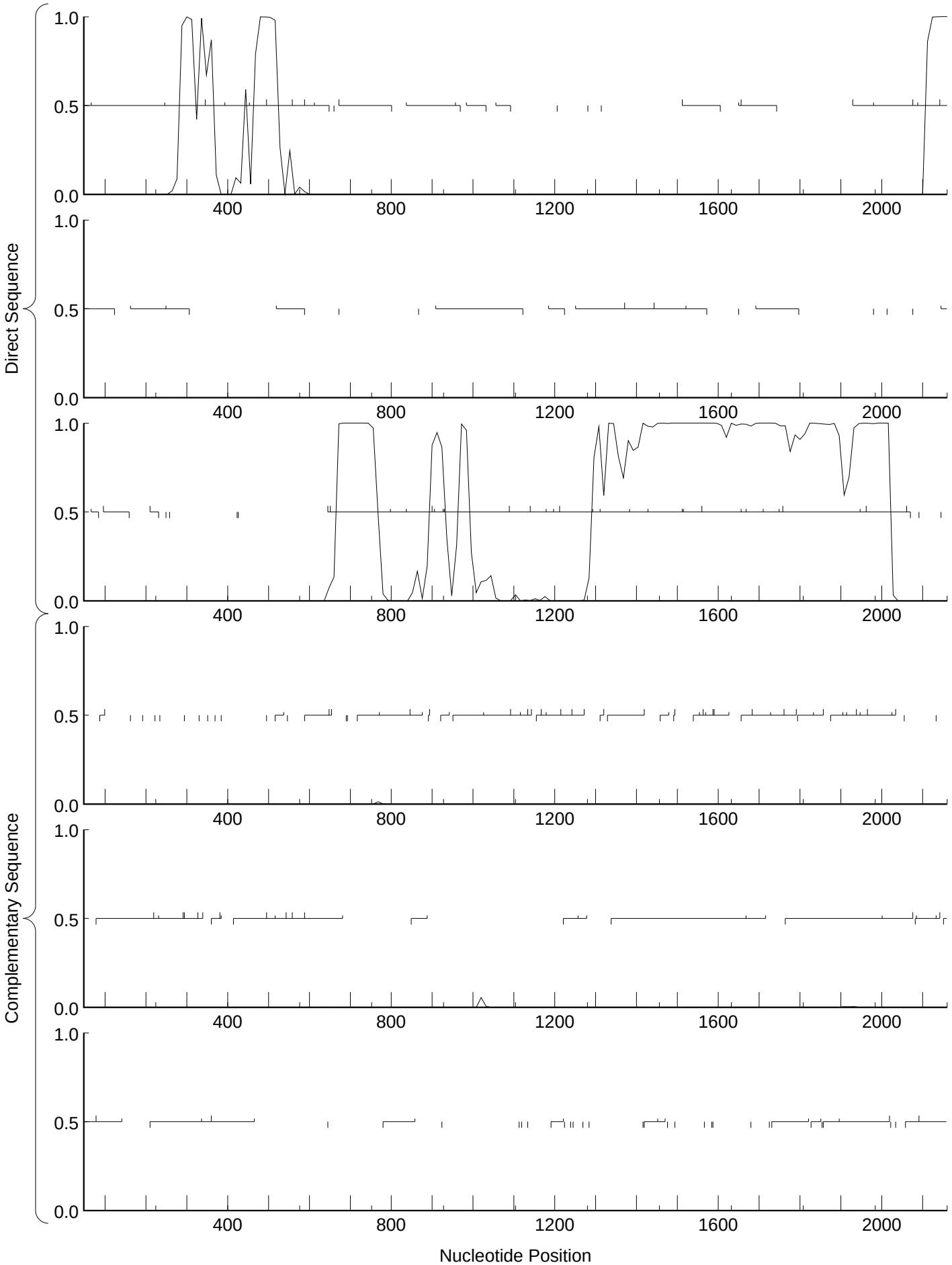
```
Matrix     :  -
Author     :  -

Order      :  4
```

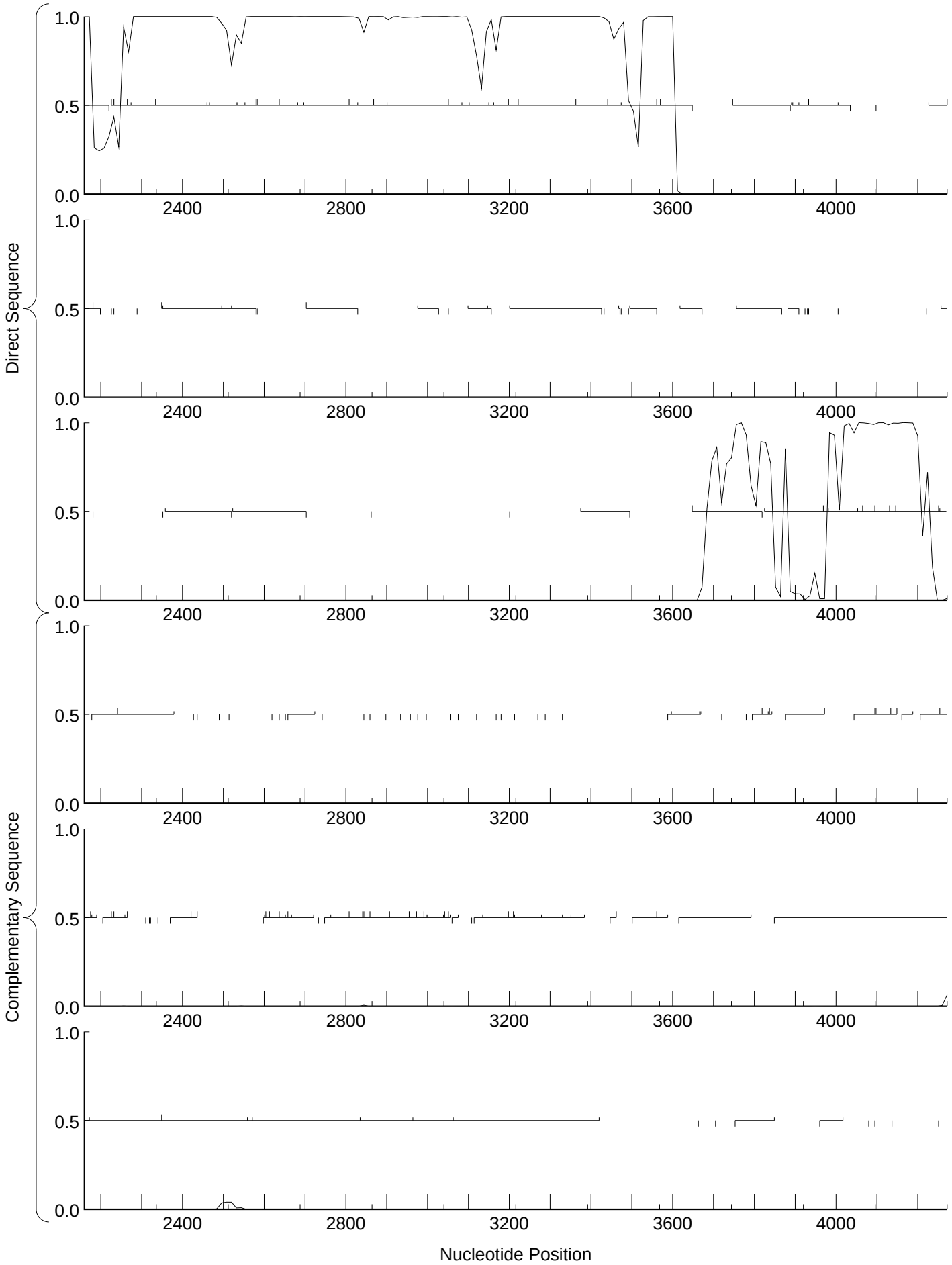

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

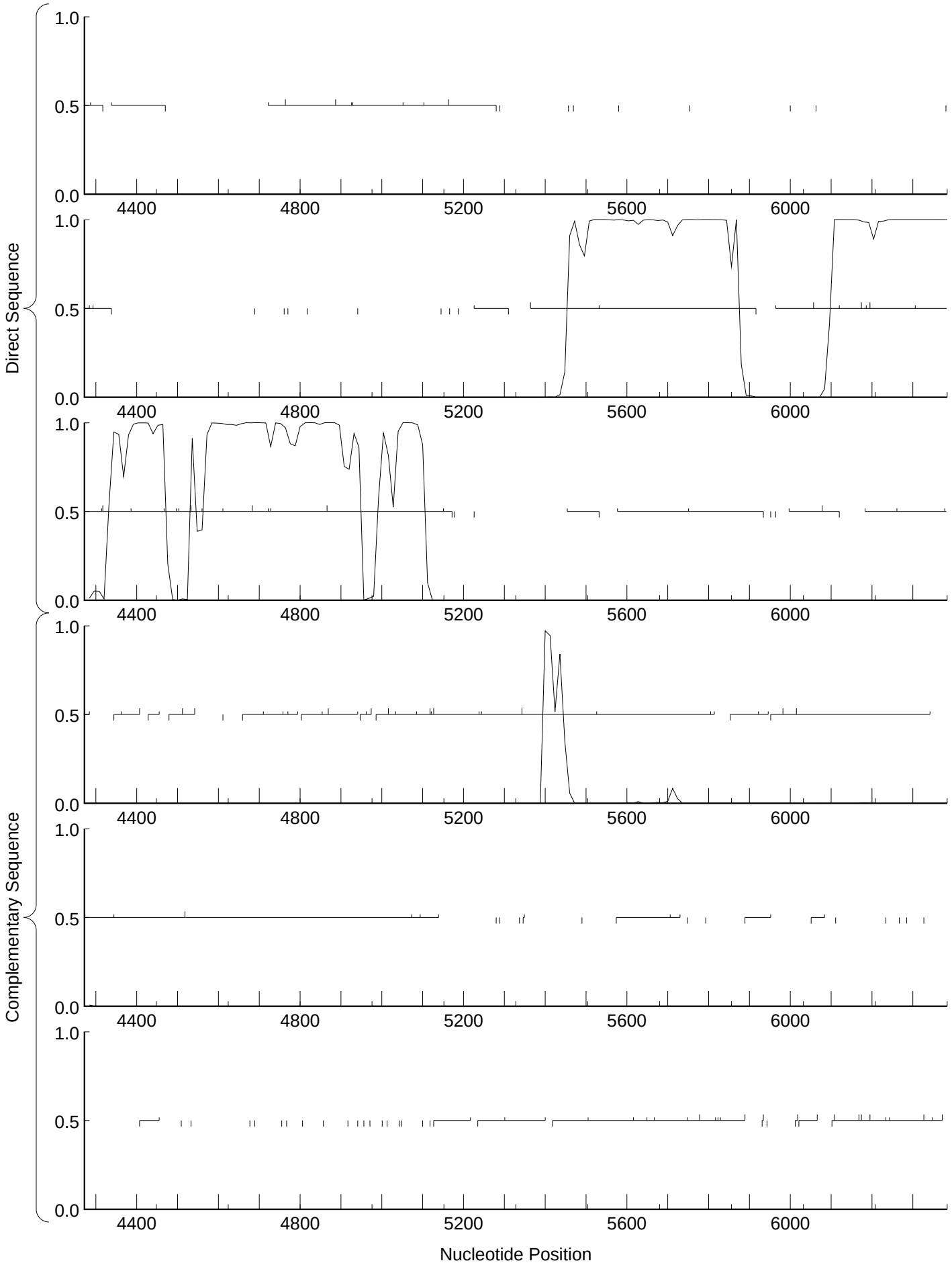
robacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 2/21

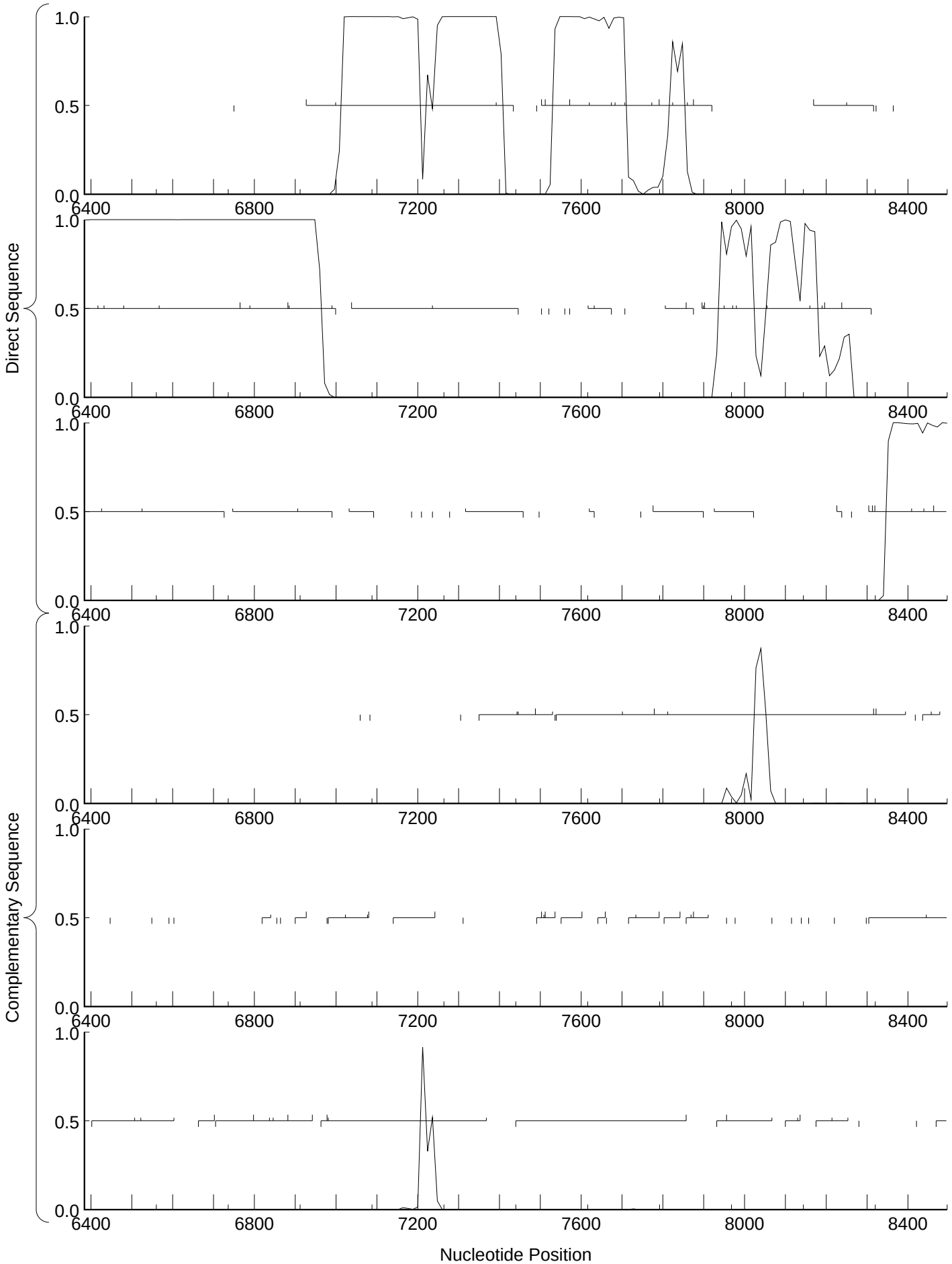


robacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 3/21

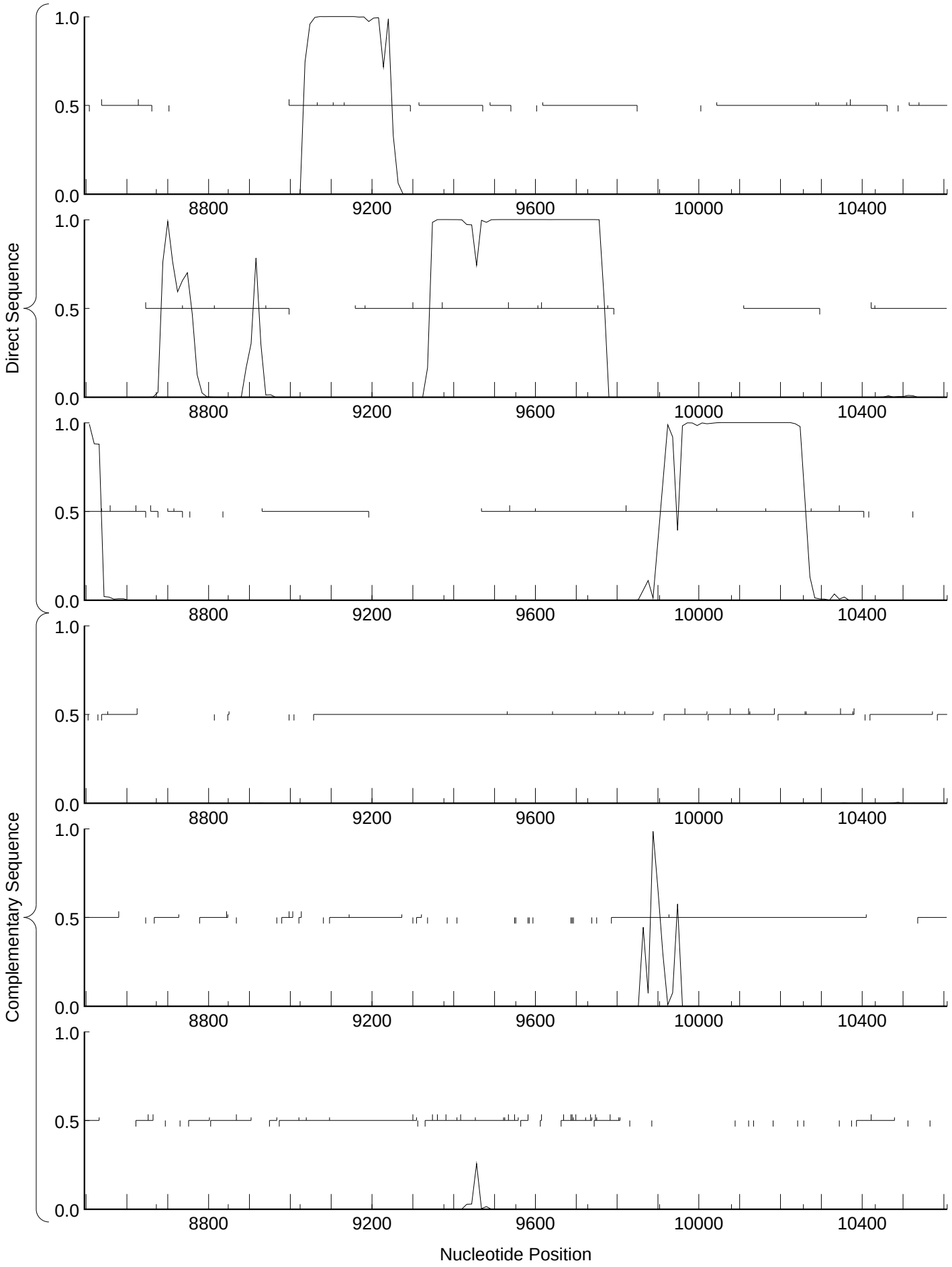


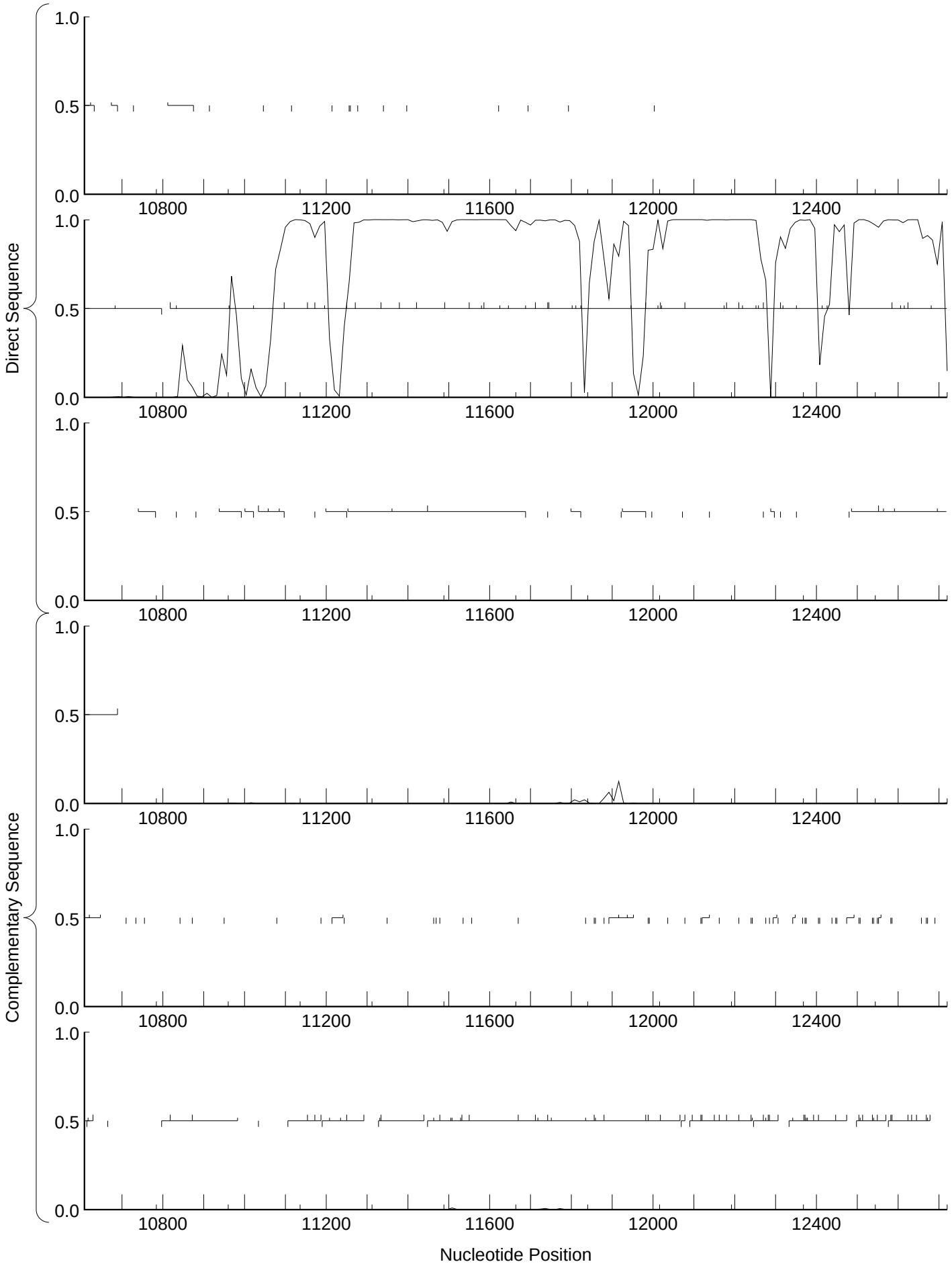
robacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 4/21



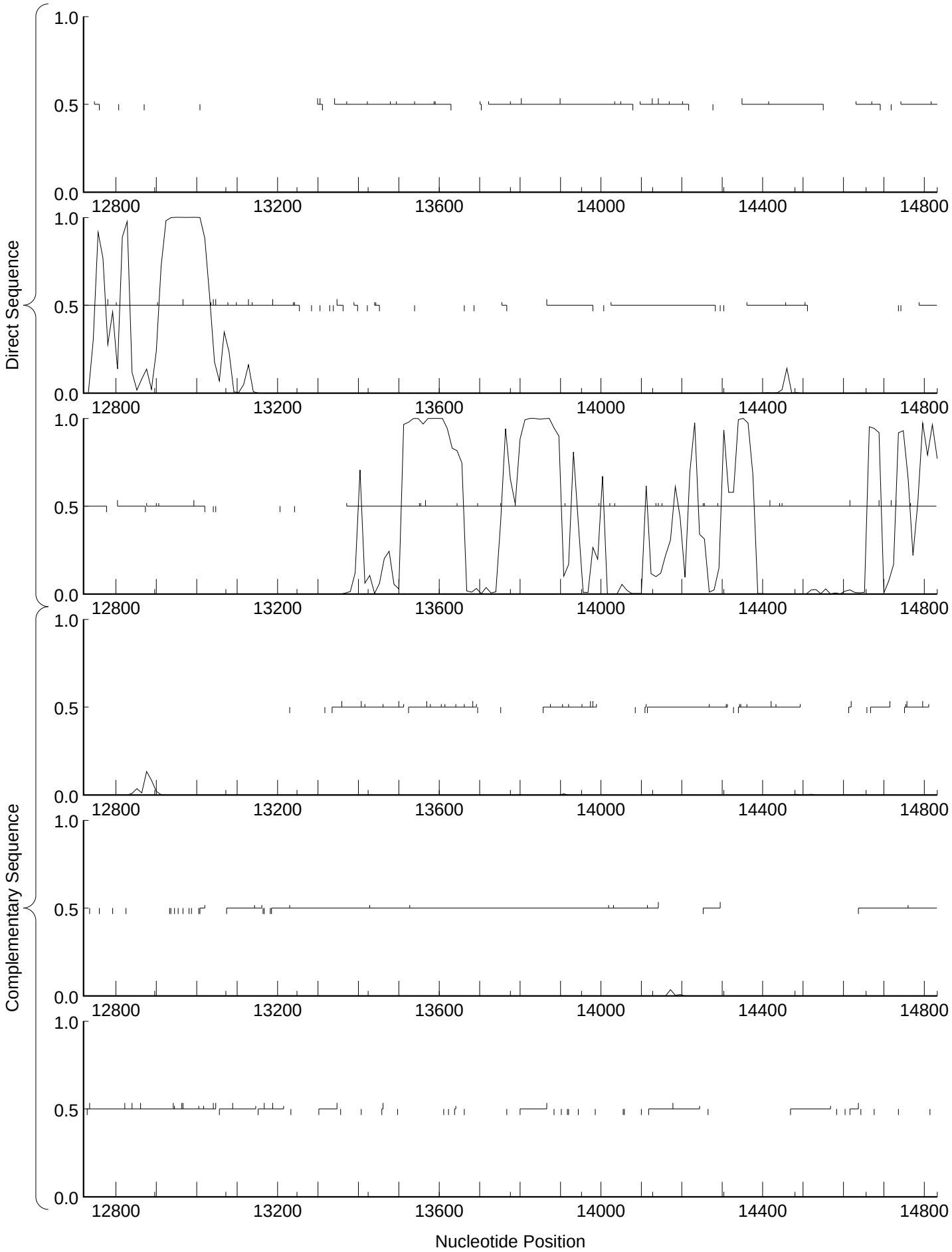

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
6400
6800
7200
7600
8000
8400
Nucleotide Position

robacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 6/21

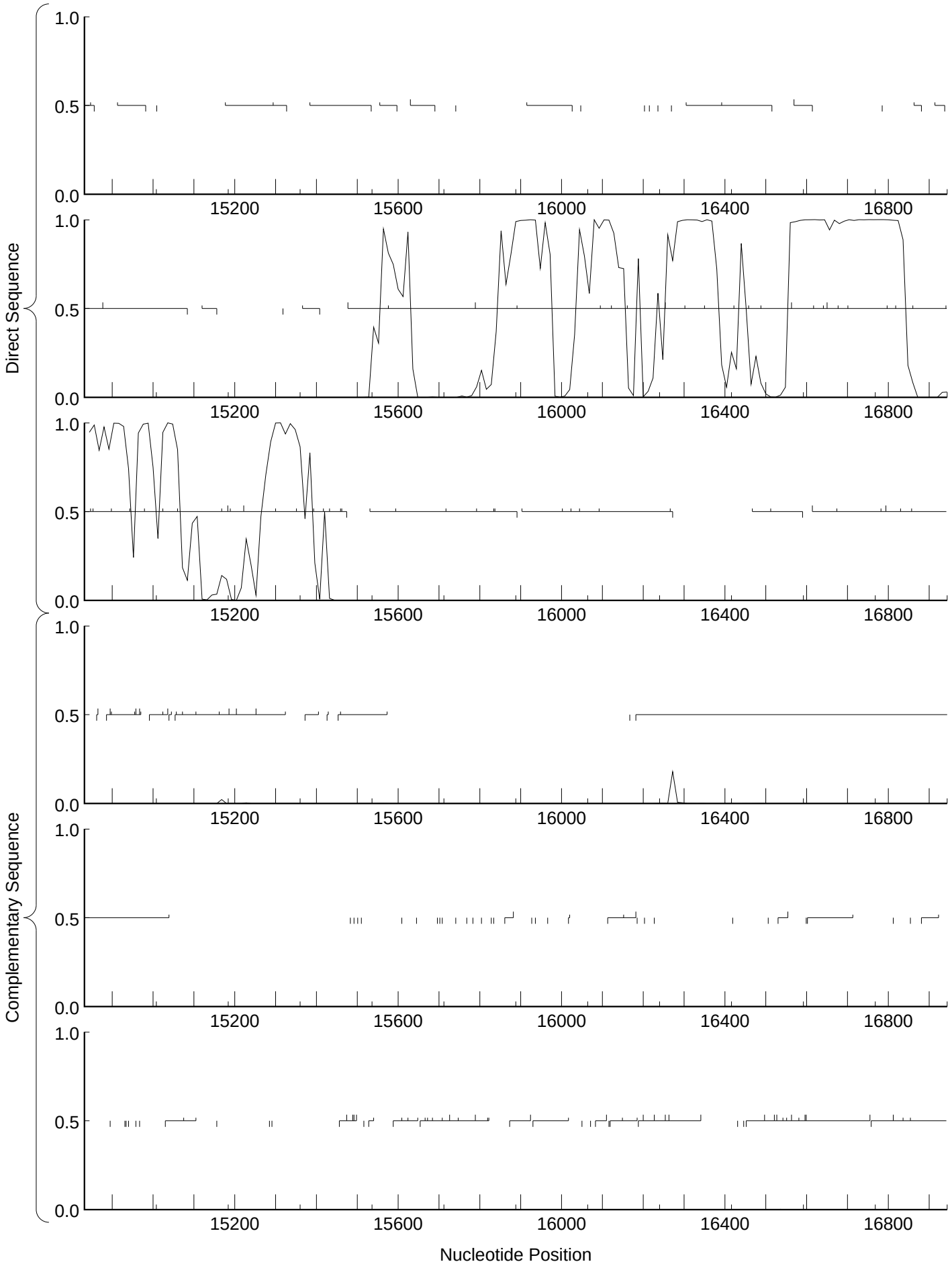



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
10800
11200
11600
12000
12400
Nucleotide Position

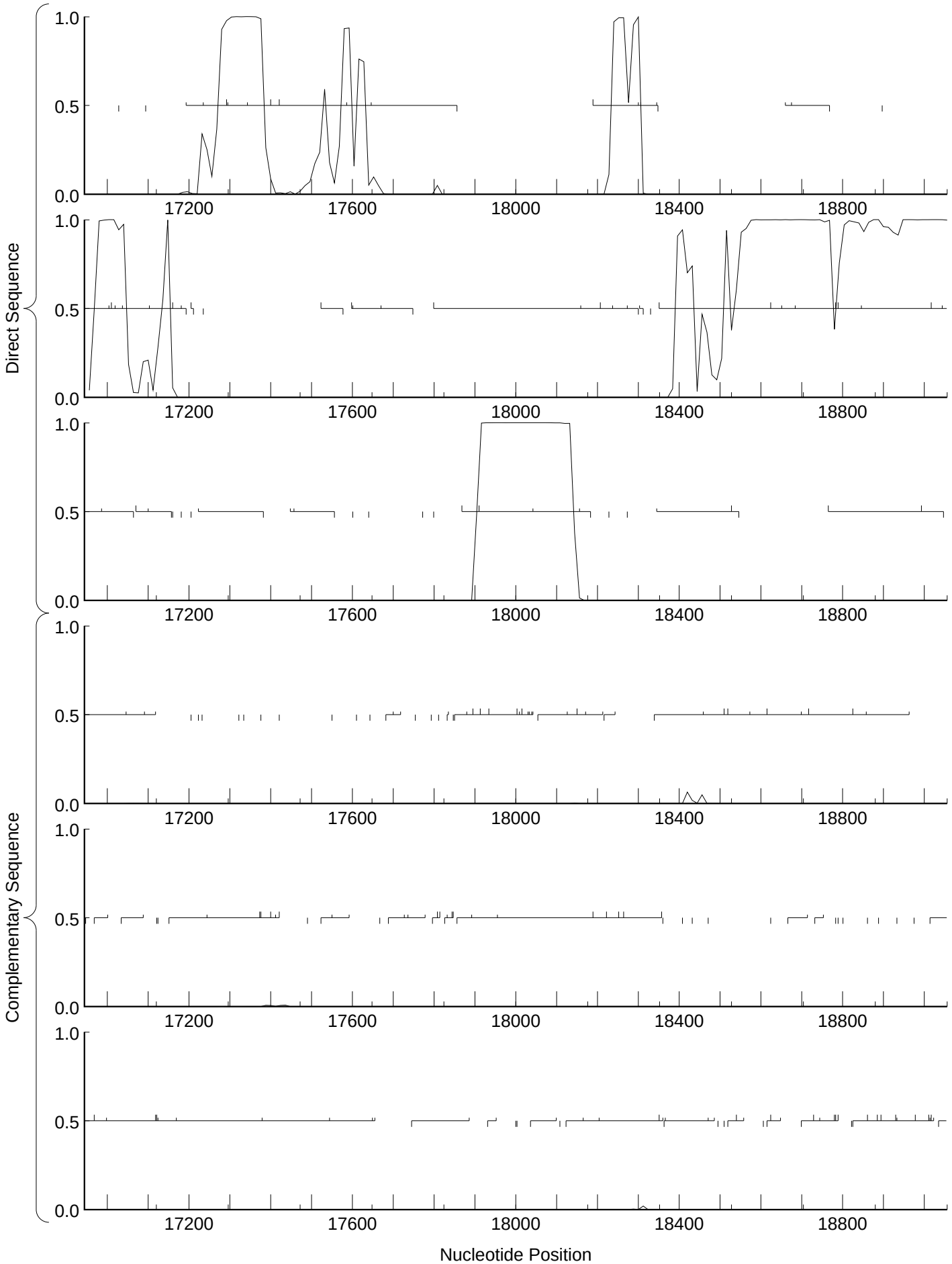
robacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 8/21



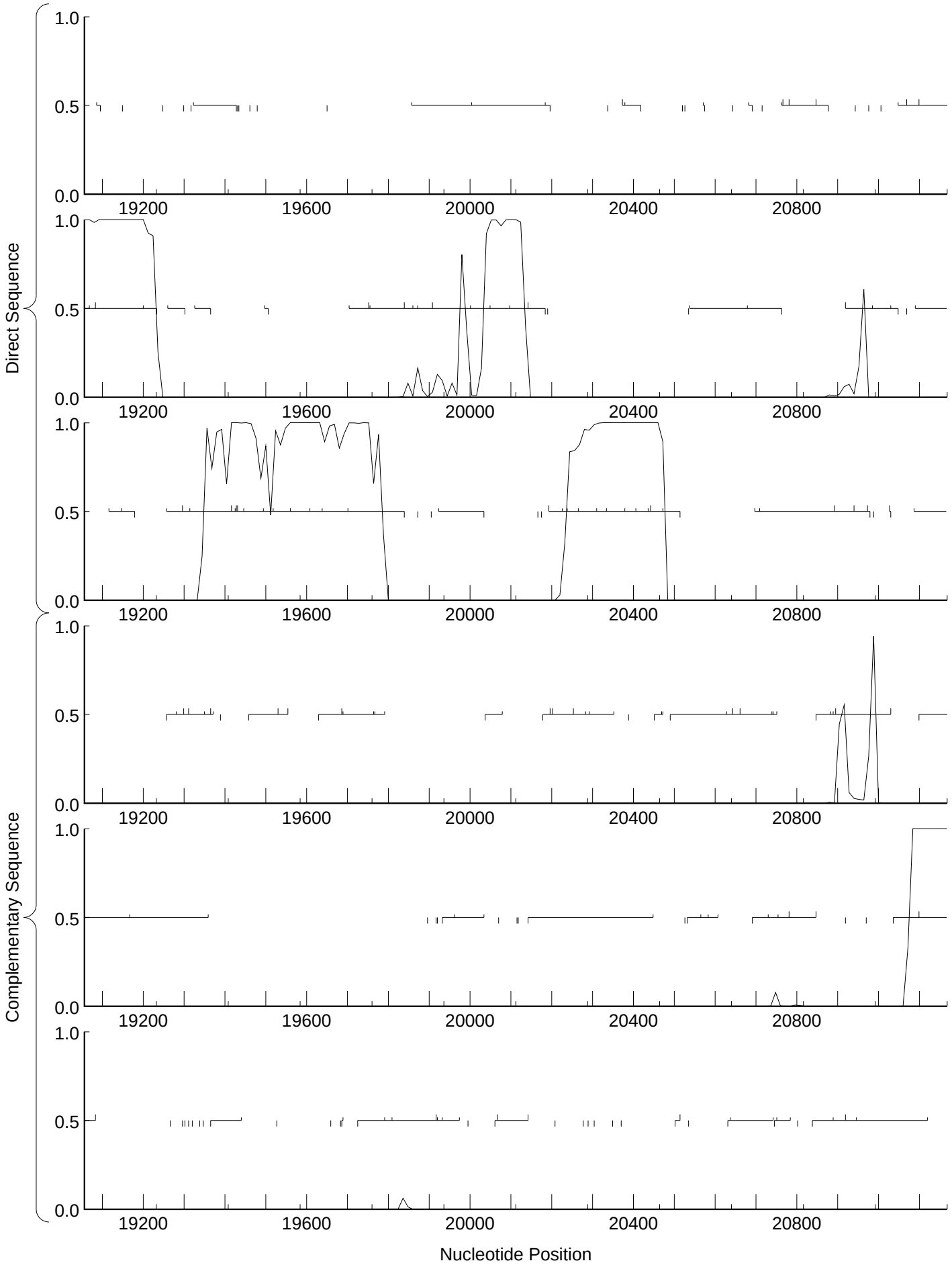
robacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 9/21



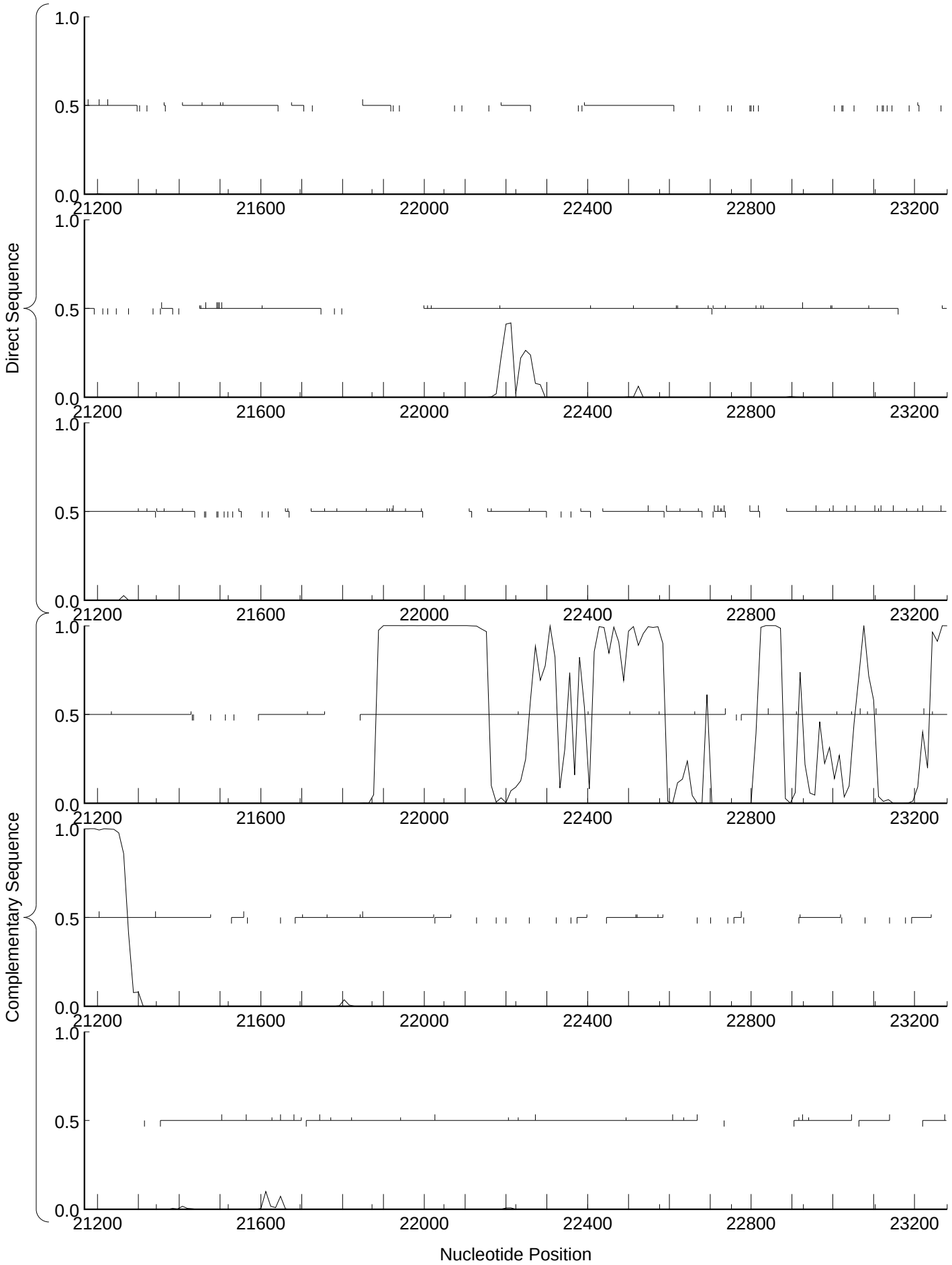
obacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 10/21



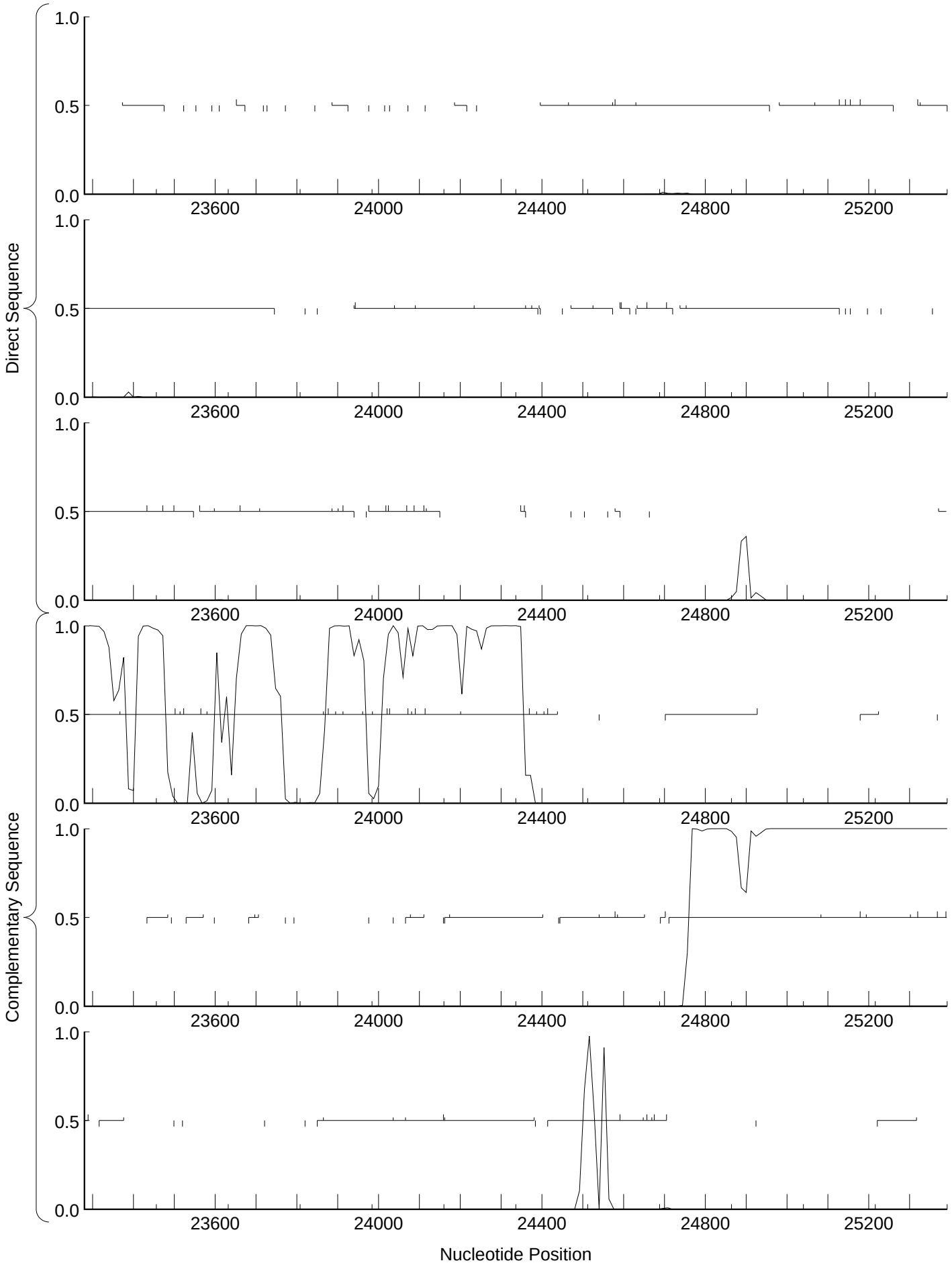
obacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 11/21



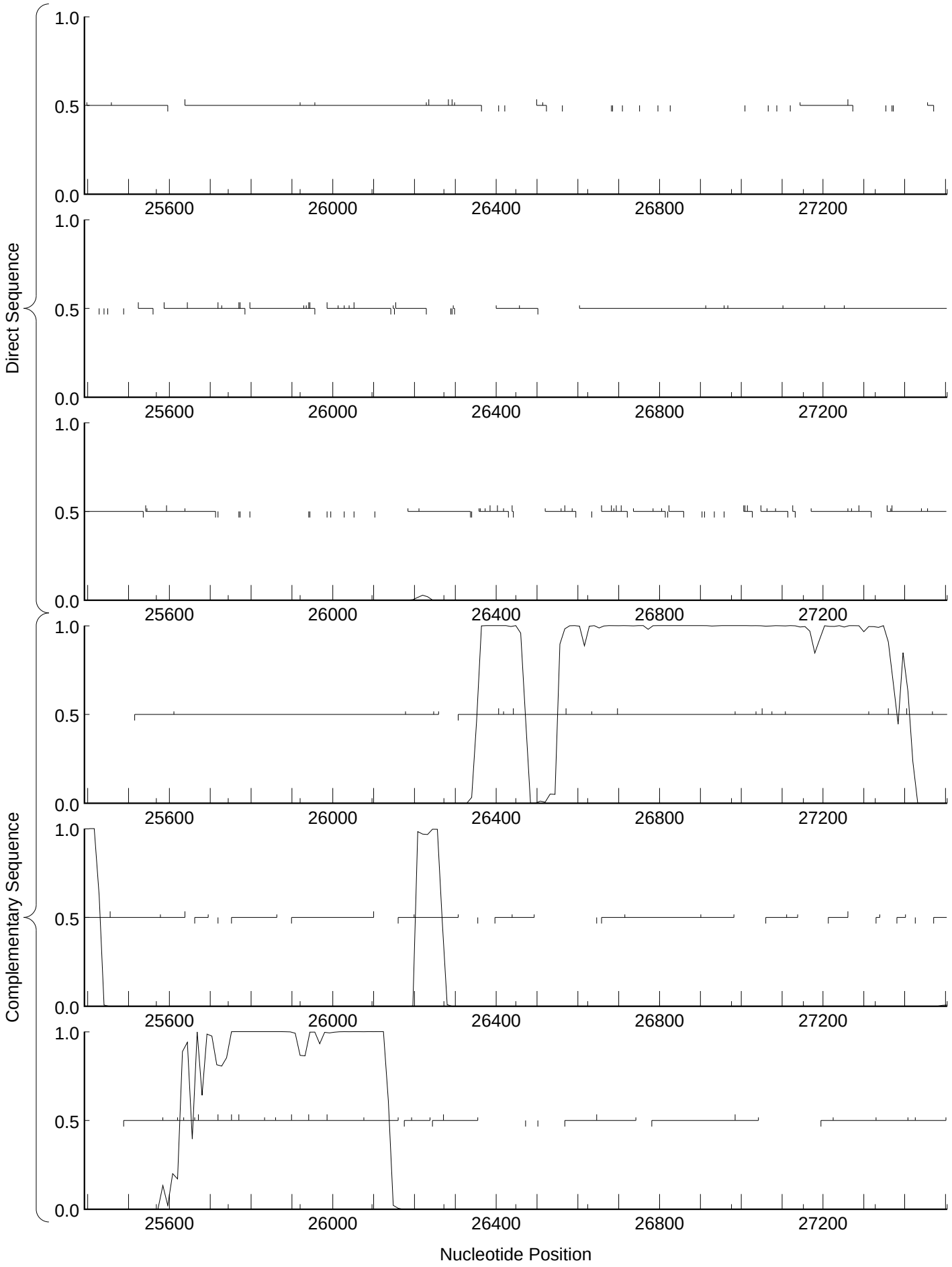
obacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 12/21



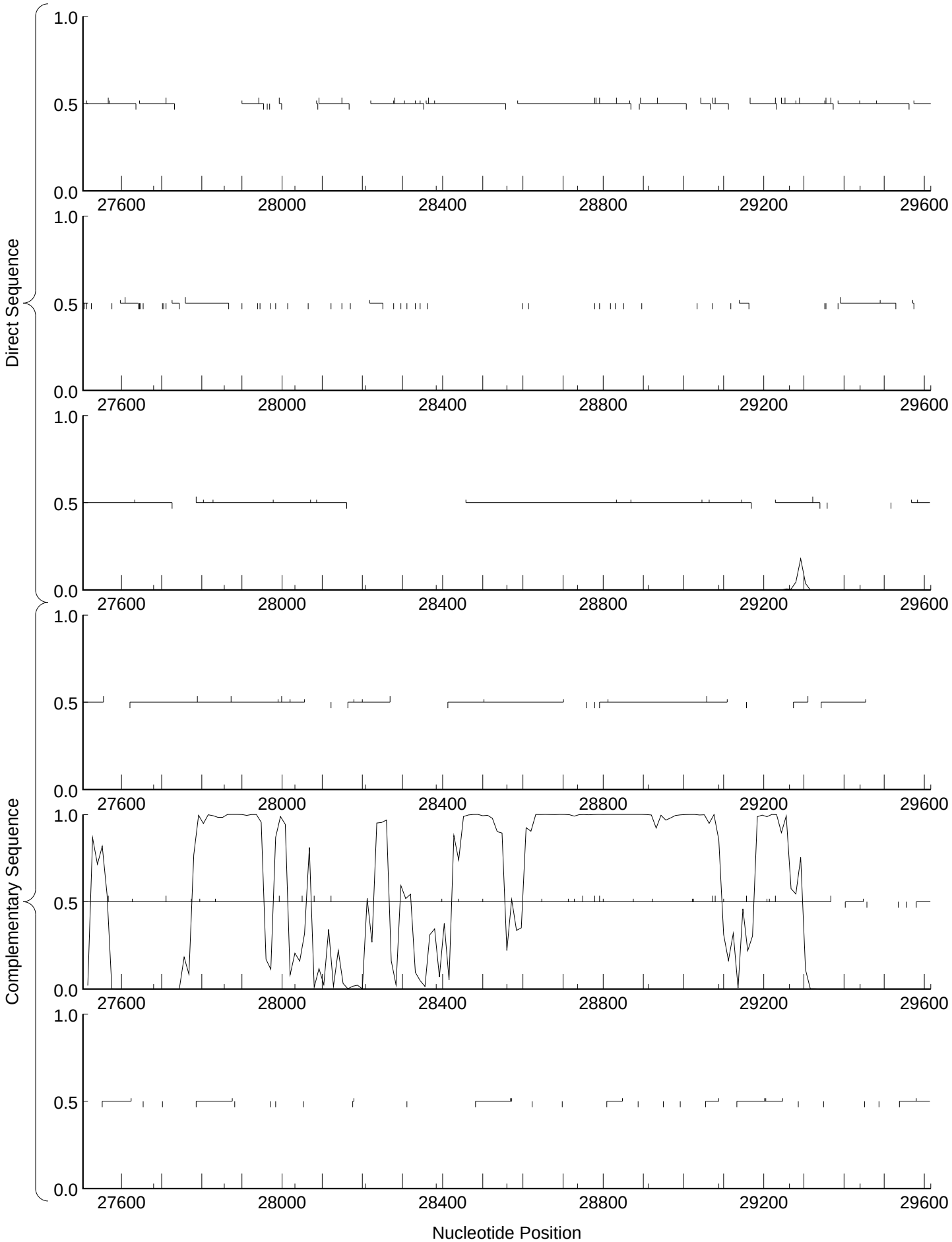
obacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 13/21



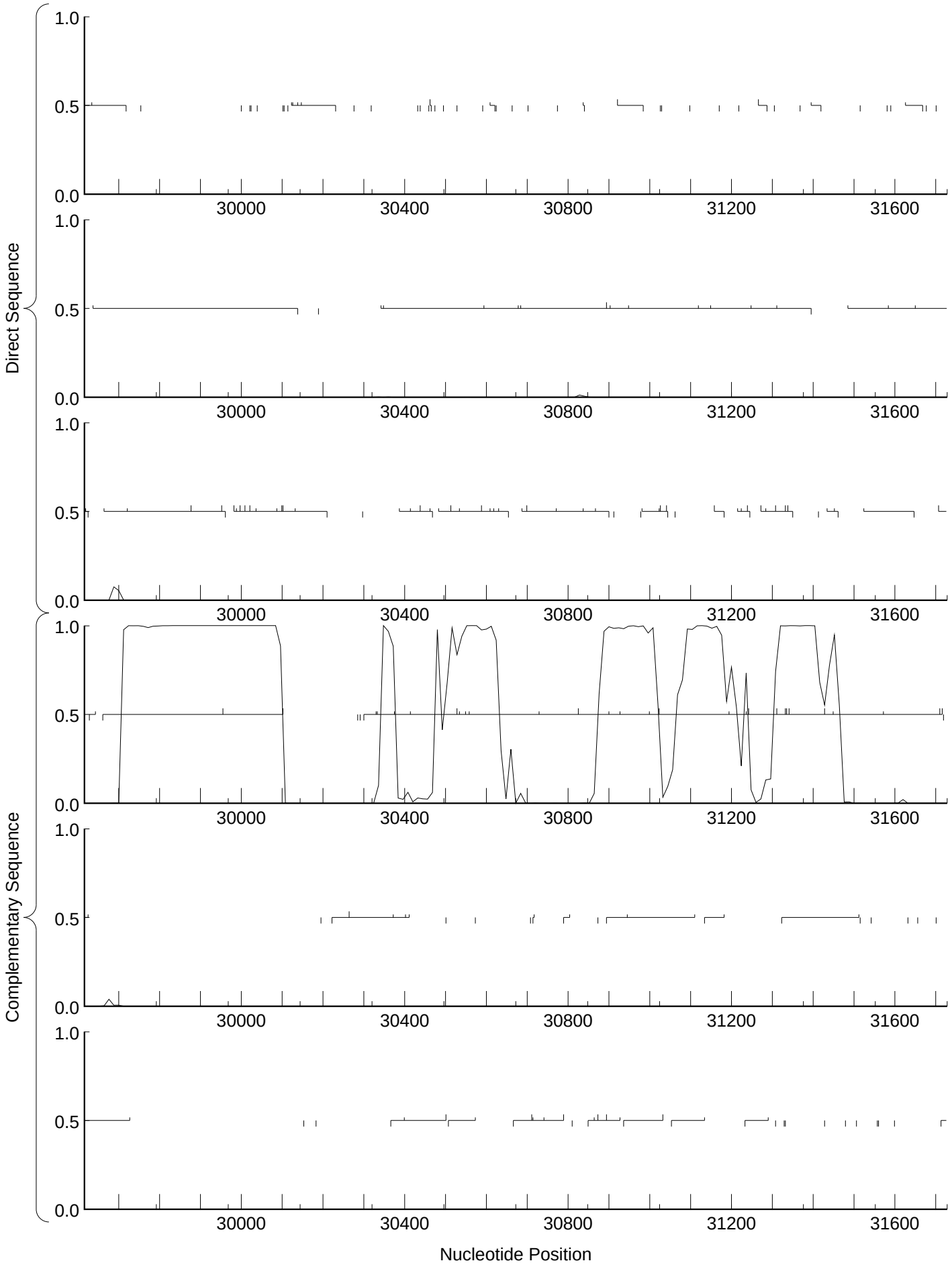
obacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 14/21



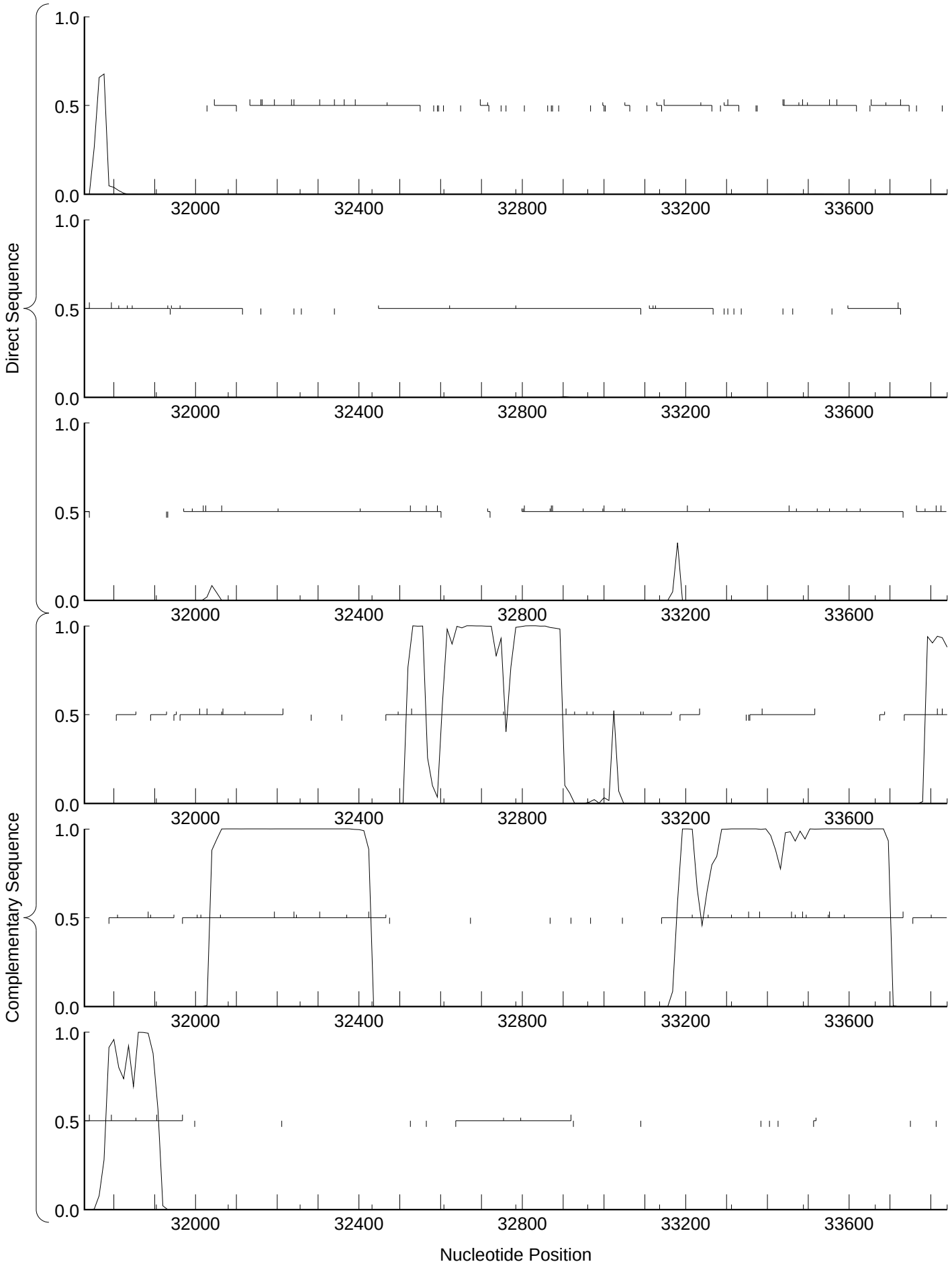
obacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 15/21



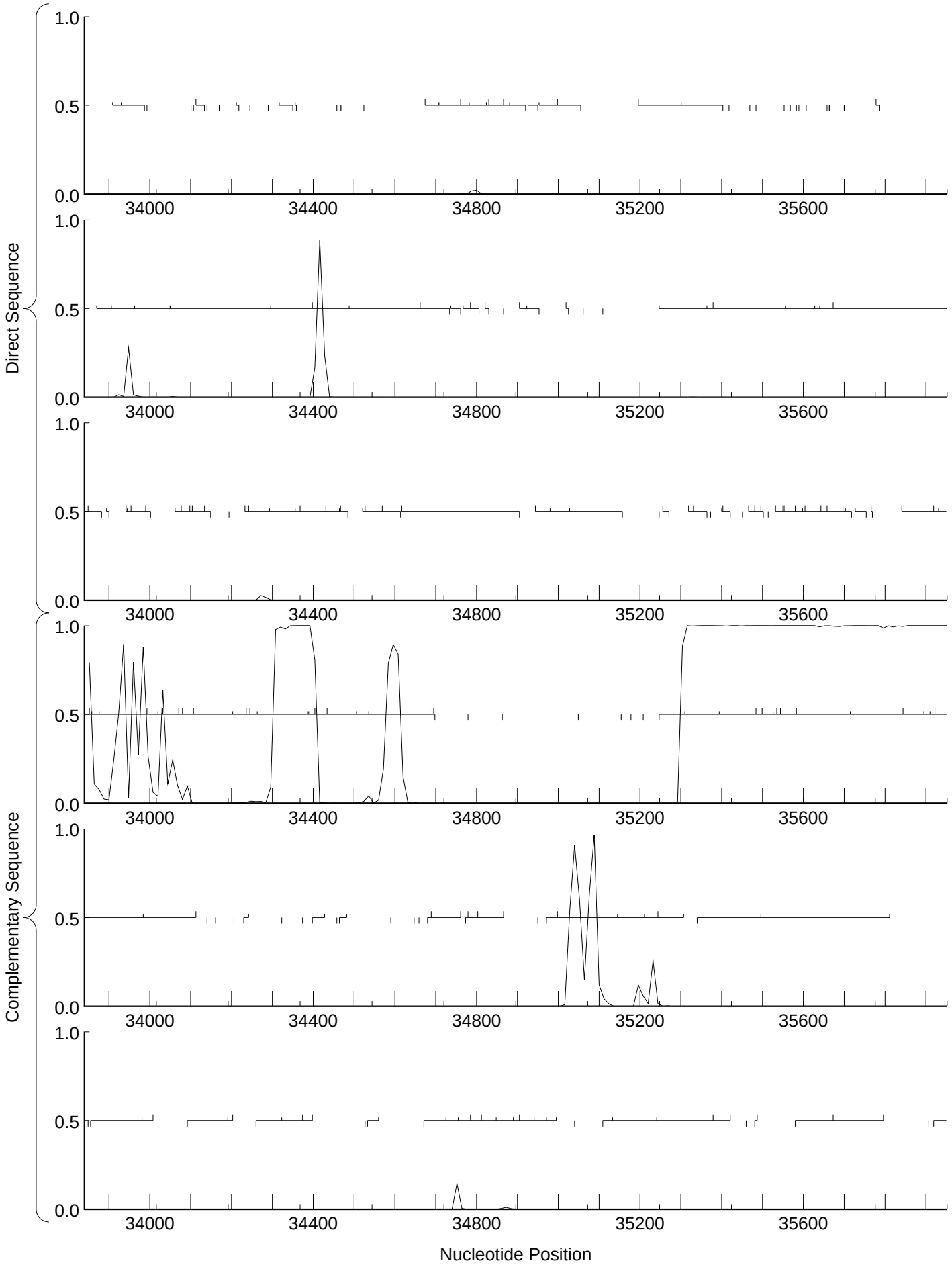
obacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 16/21



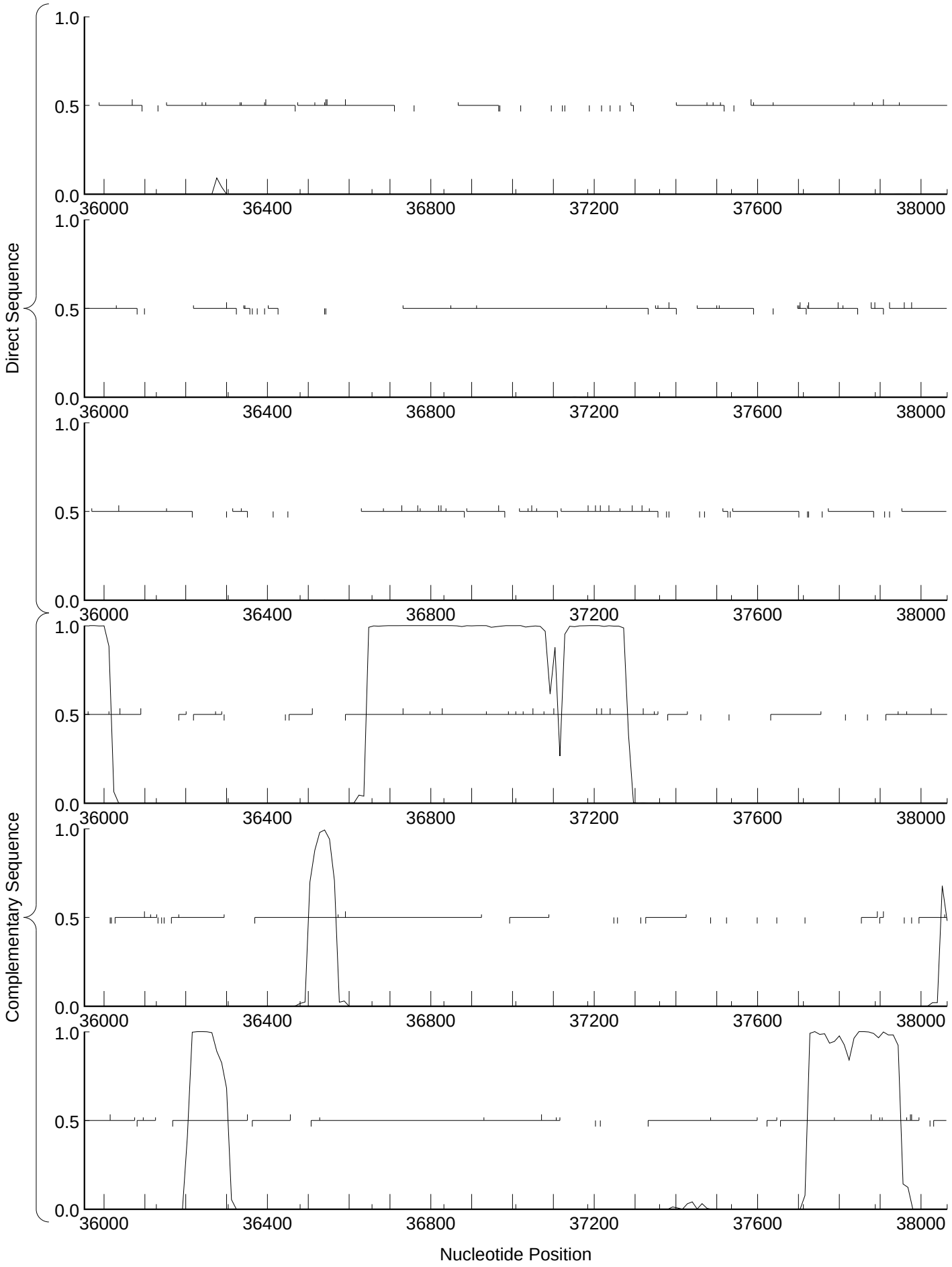
obacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 17/21



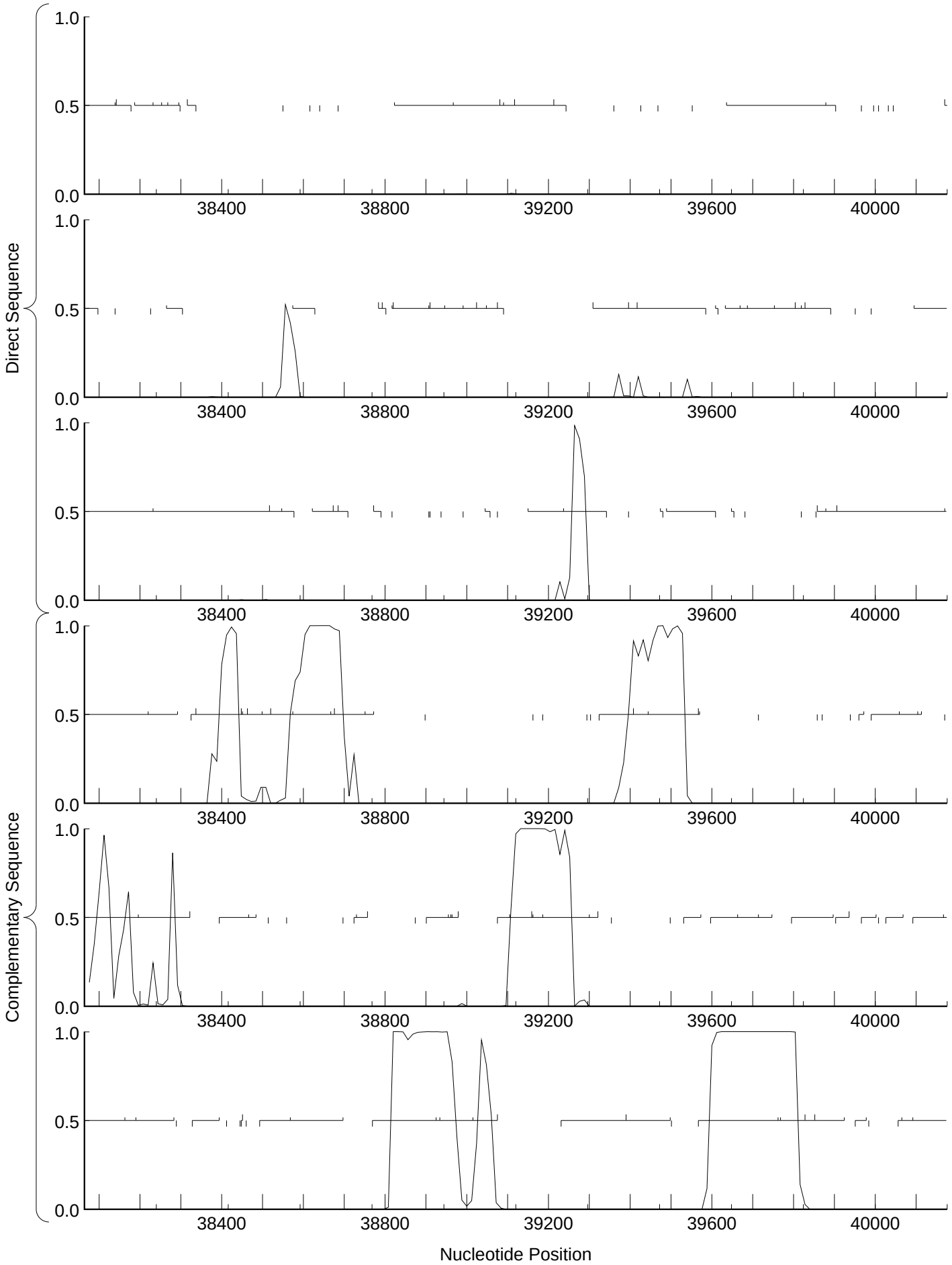
obacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 18/21



obacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 19/21



obacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 20/21



obacterium phage Jemerald complete sequence, 41099 bp, circularly permuted, Cluster EA6, Order 4, Window 96, Step 12, 21/21

