GeneMark

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

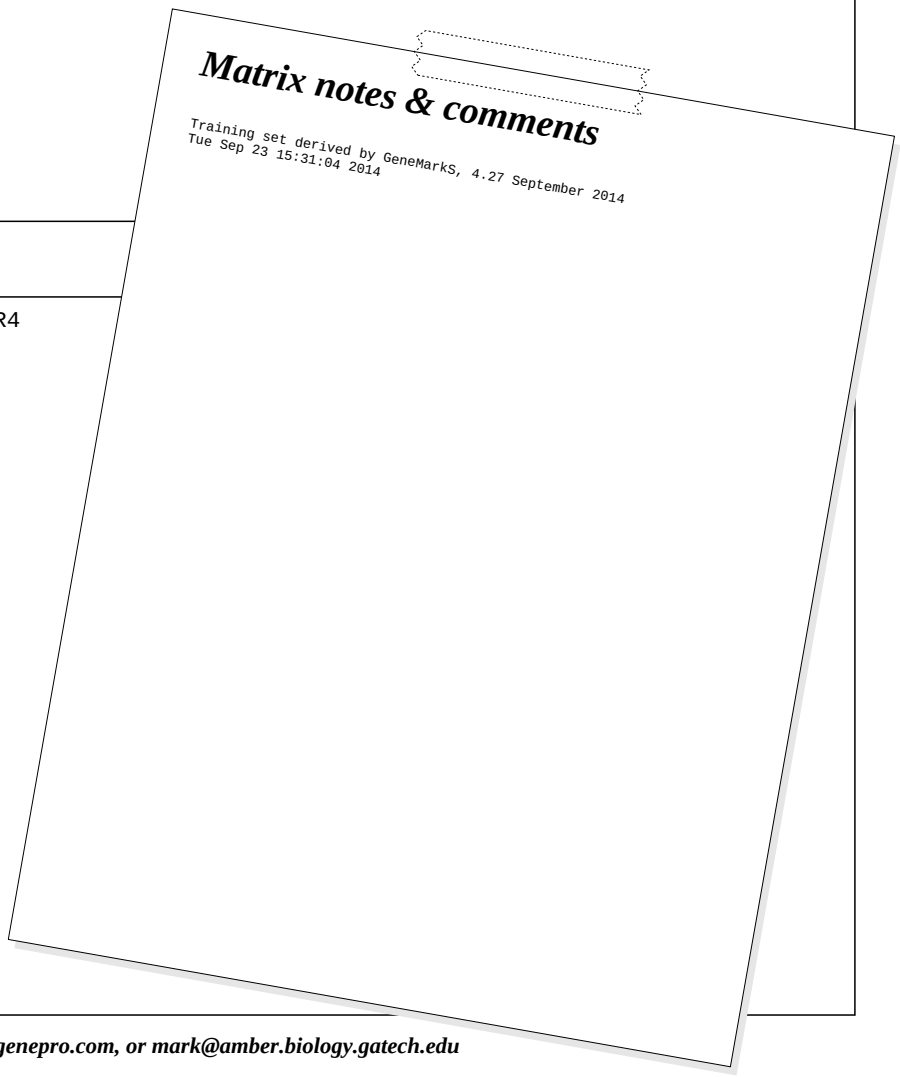
PROGRAM INFORMATION

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Analysis Date    :  3/18/24 at 21:42:37
Pages            :  27
Sequence Length  :  53912 bp
GC Content       :   66.98%

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

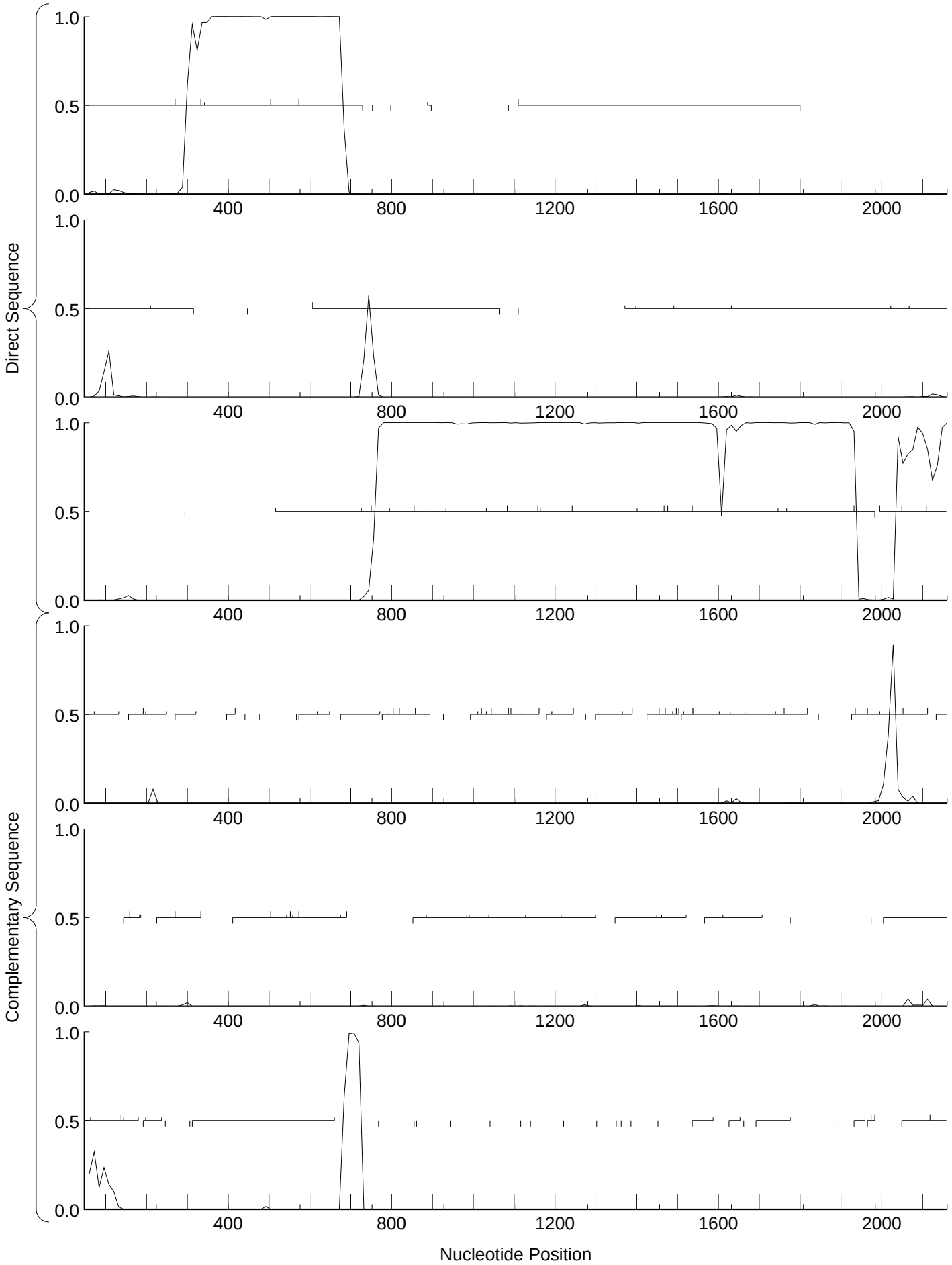
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Matrix     :  Rhodococcus_erythropolis_PR4
Author     :  -

Order      :  4
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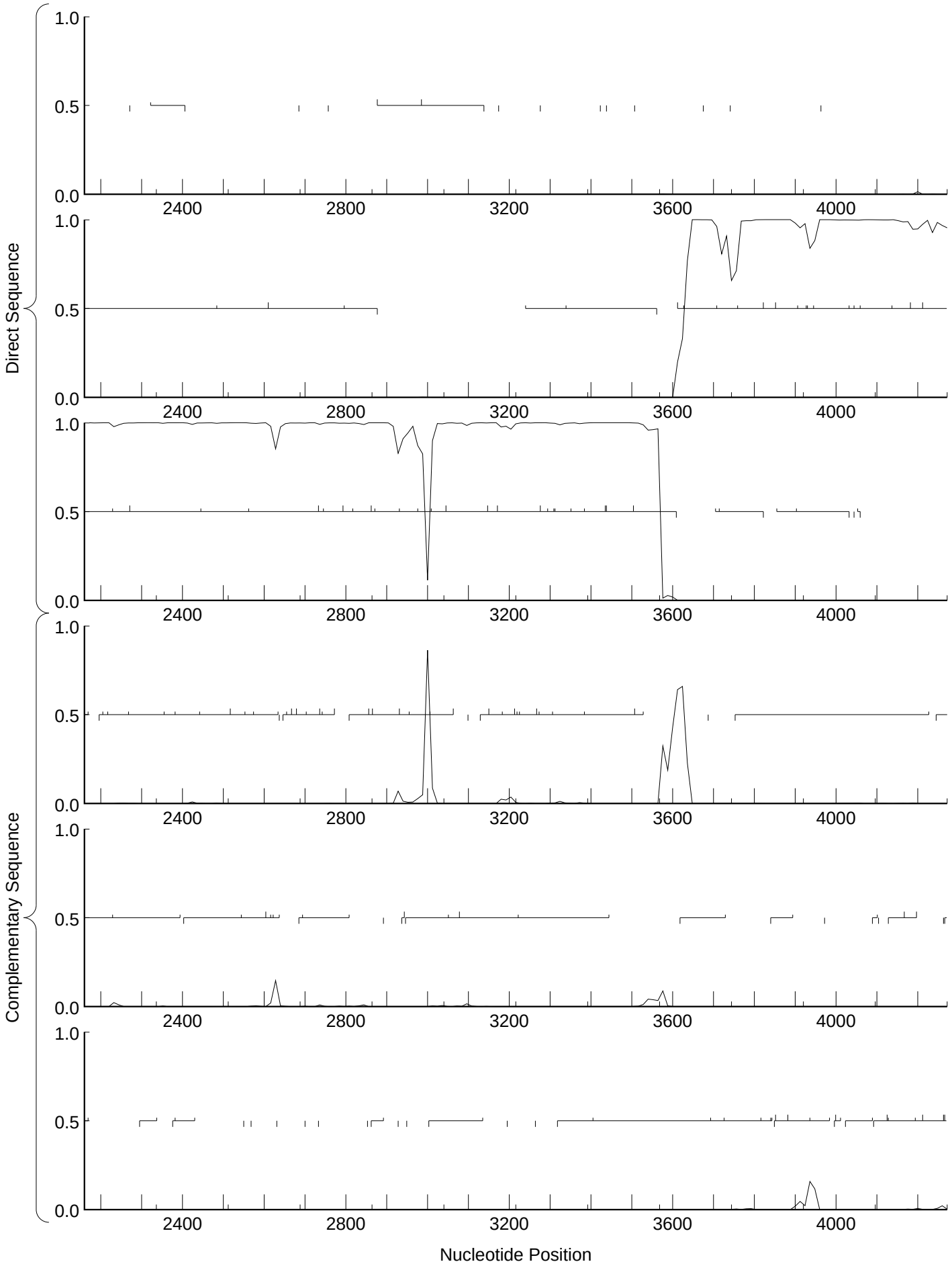

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

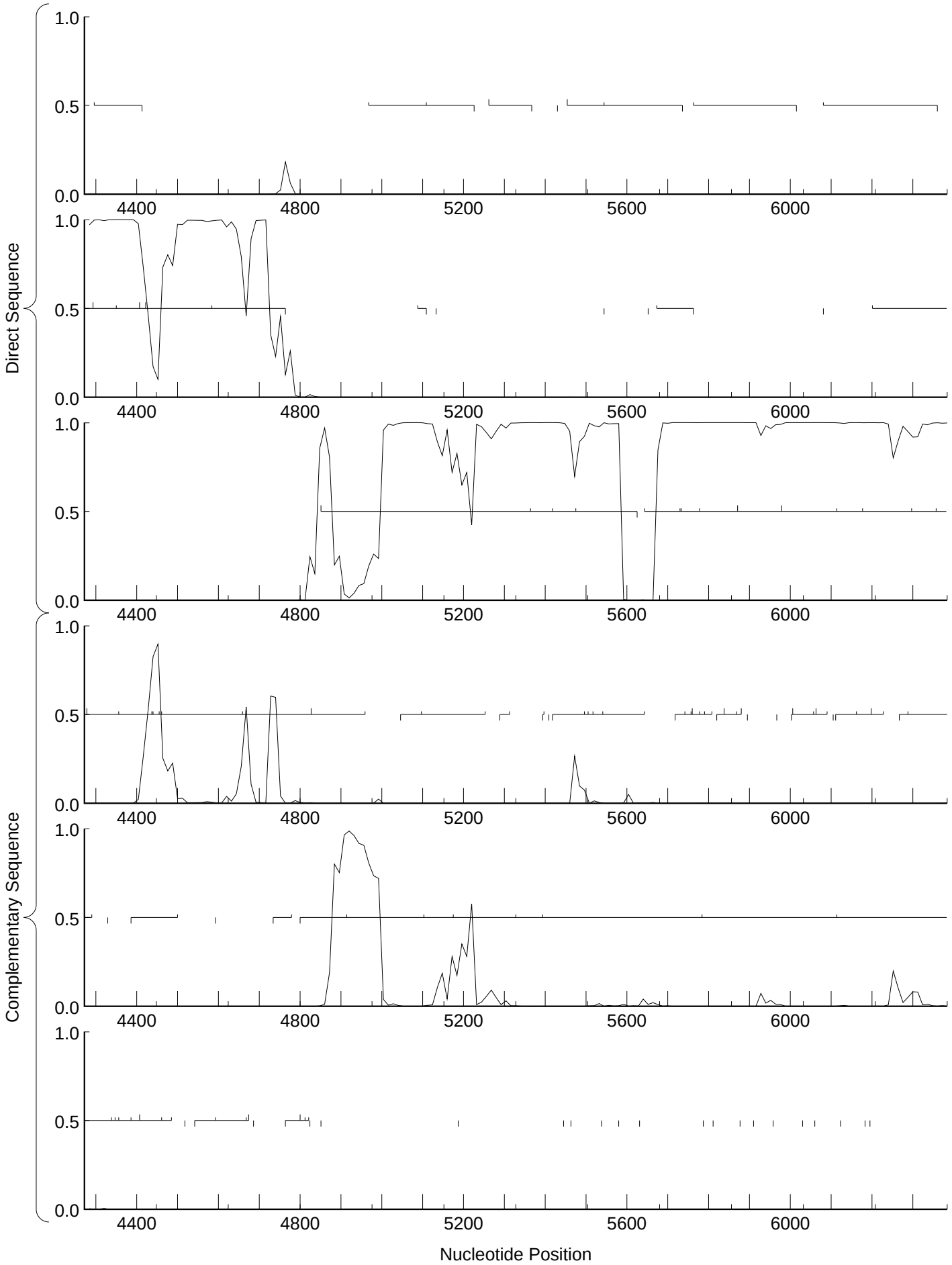
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 2/27



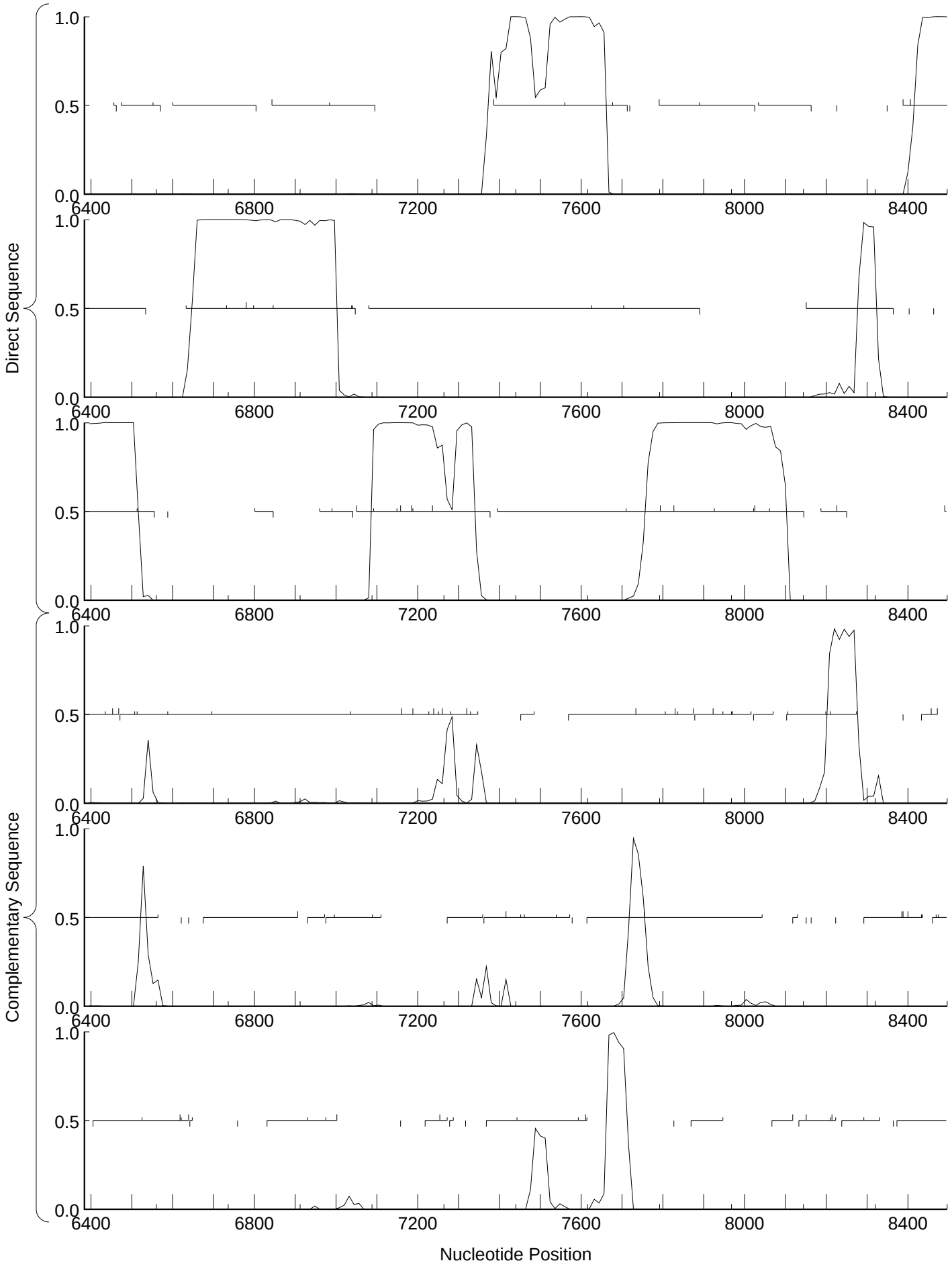
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 3/27



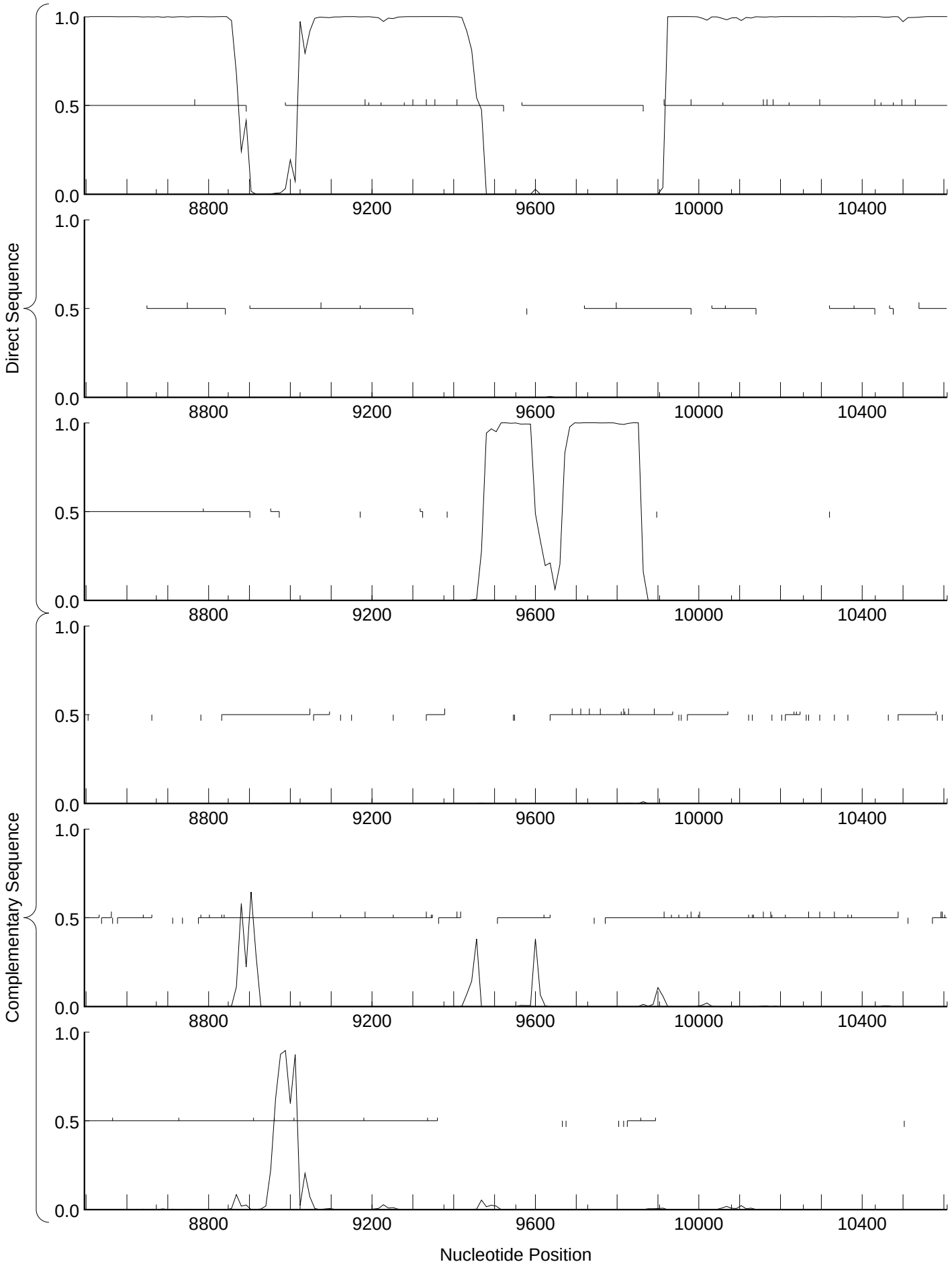
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 4/27

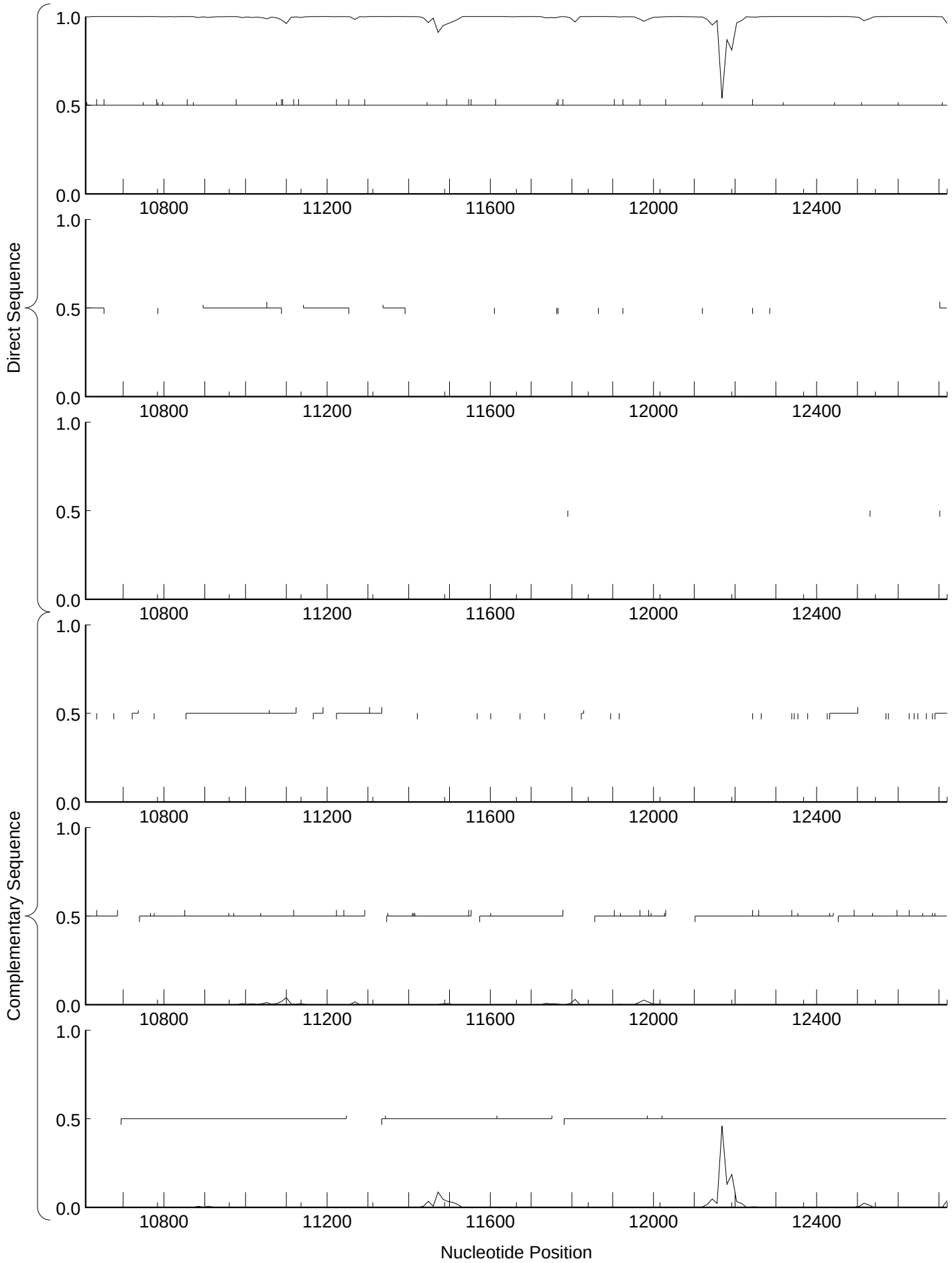


Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 5/27

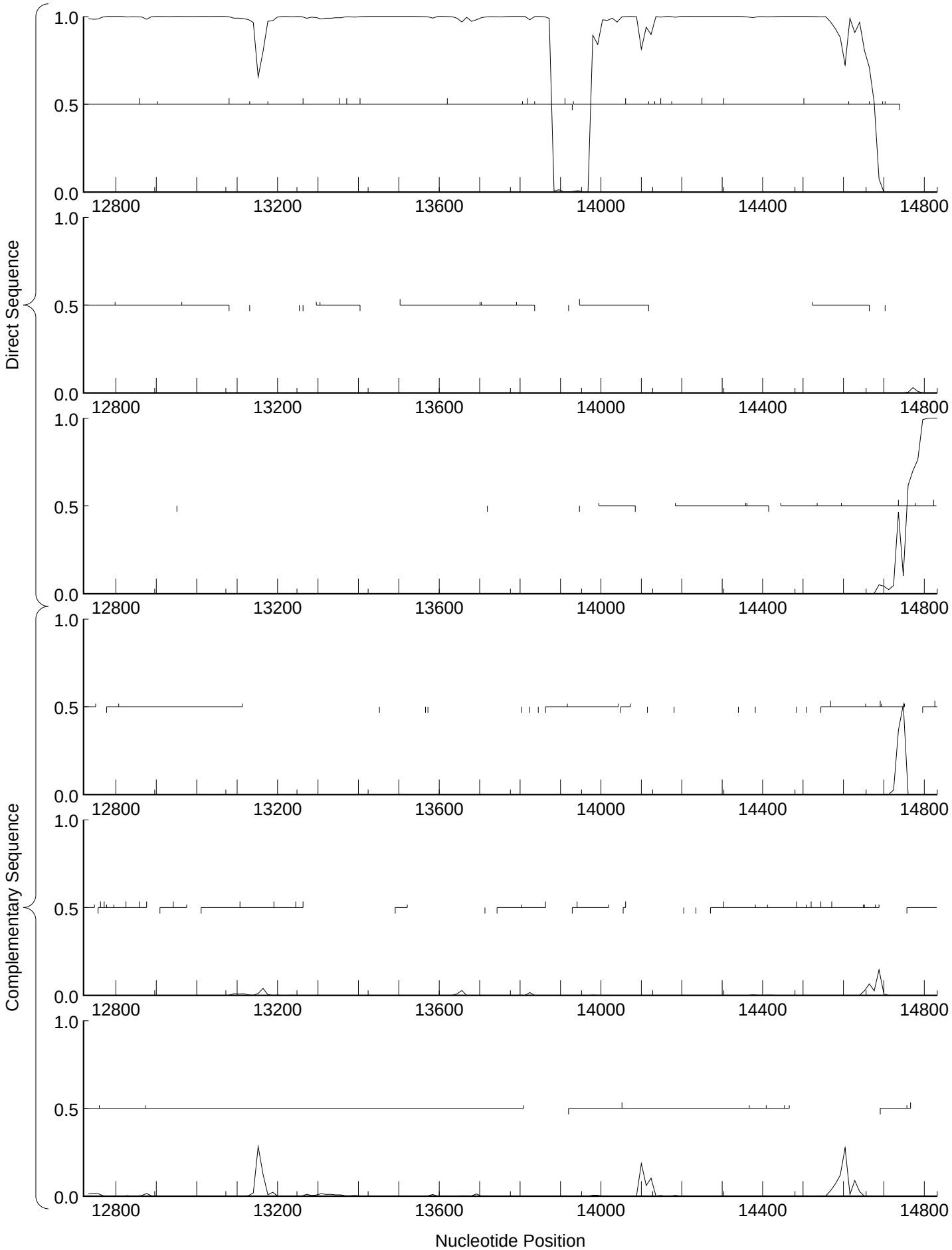


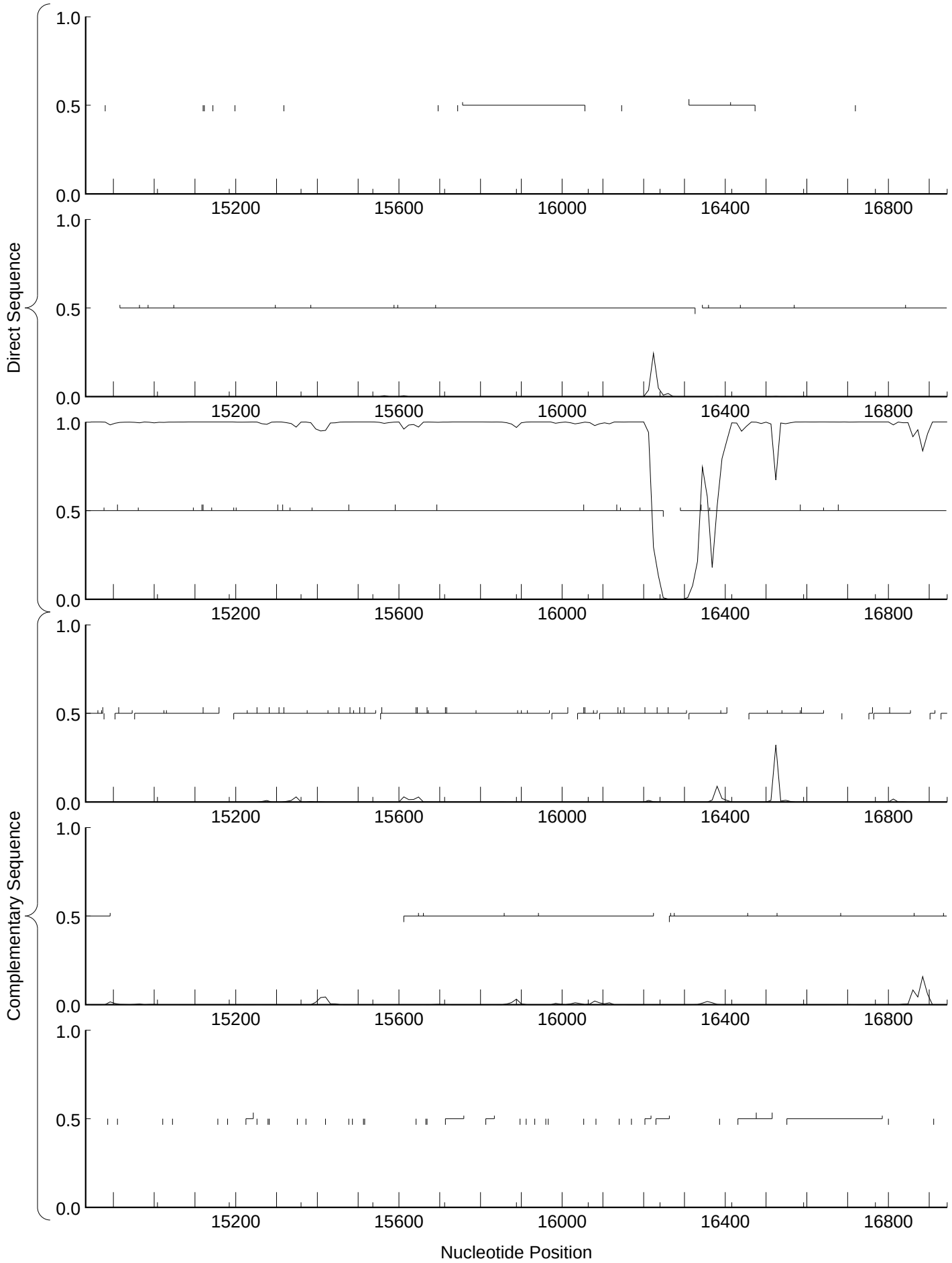
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 6/27



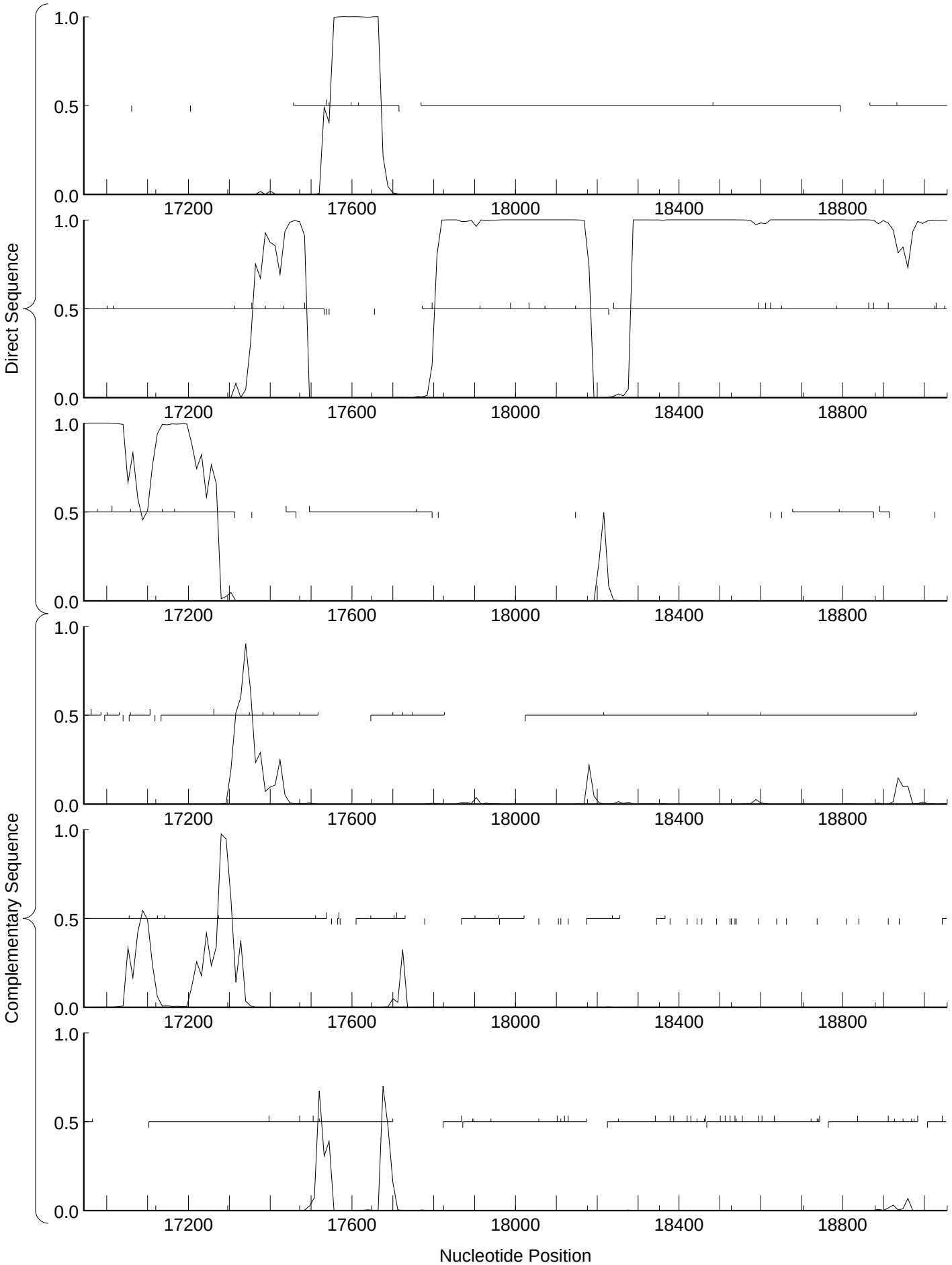

Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 7/27
Direct Sequence
Complementary Sequence
1.0
0.5
0.0
10800
11200
11600
12000
12400
Nucleotide Position

Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 8/27

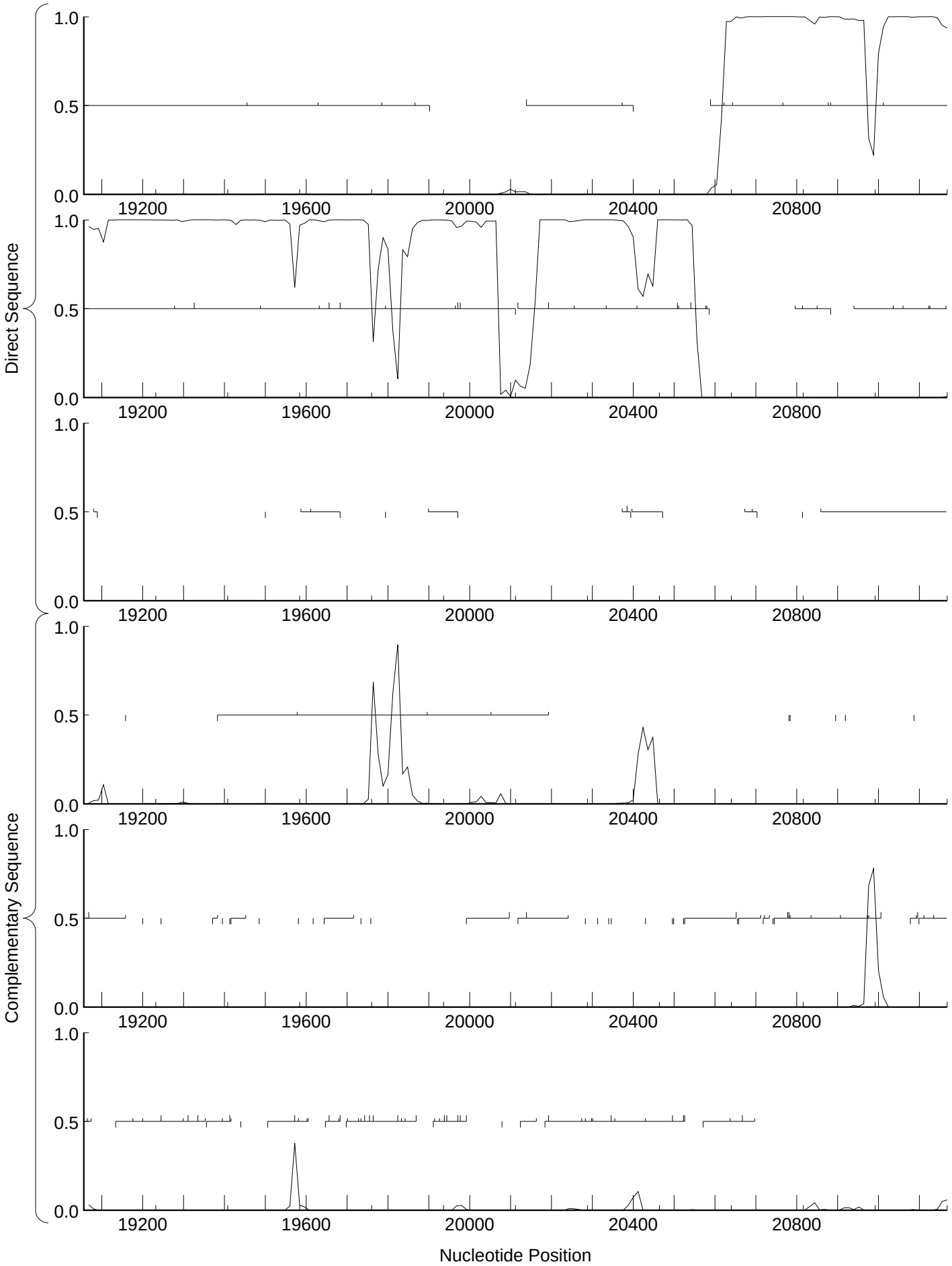



Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 9/27
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Complementary Sequence
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0.0
15200
15600
16000
16400
16800
Nucleotide Position

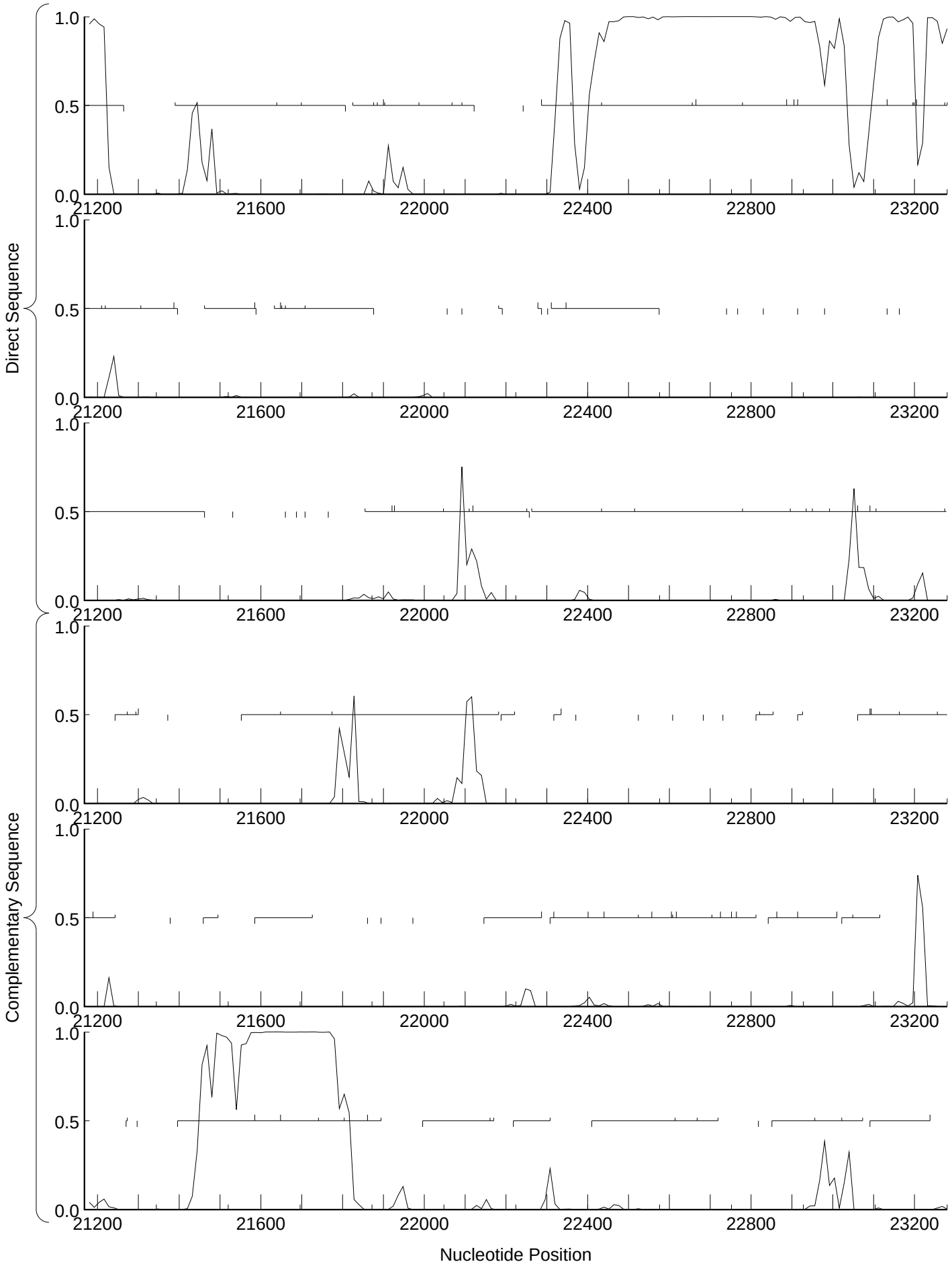
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 10/27

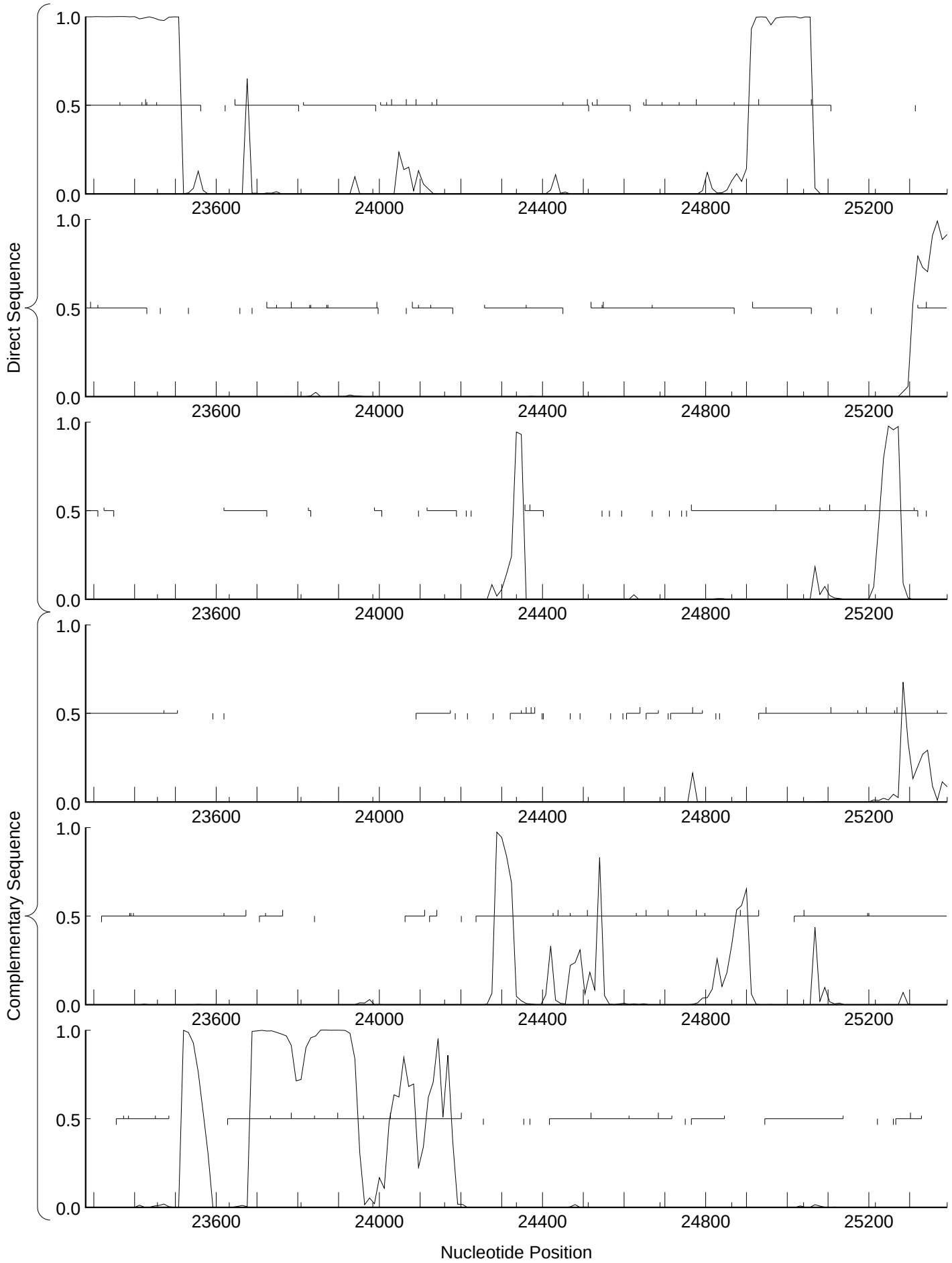


Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 11/27

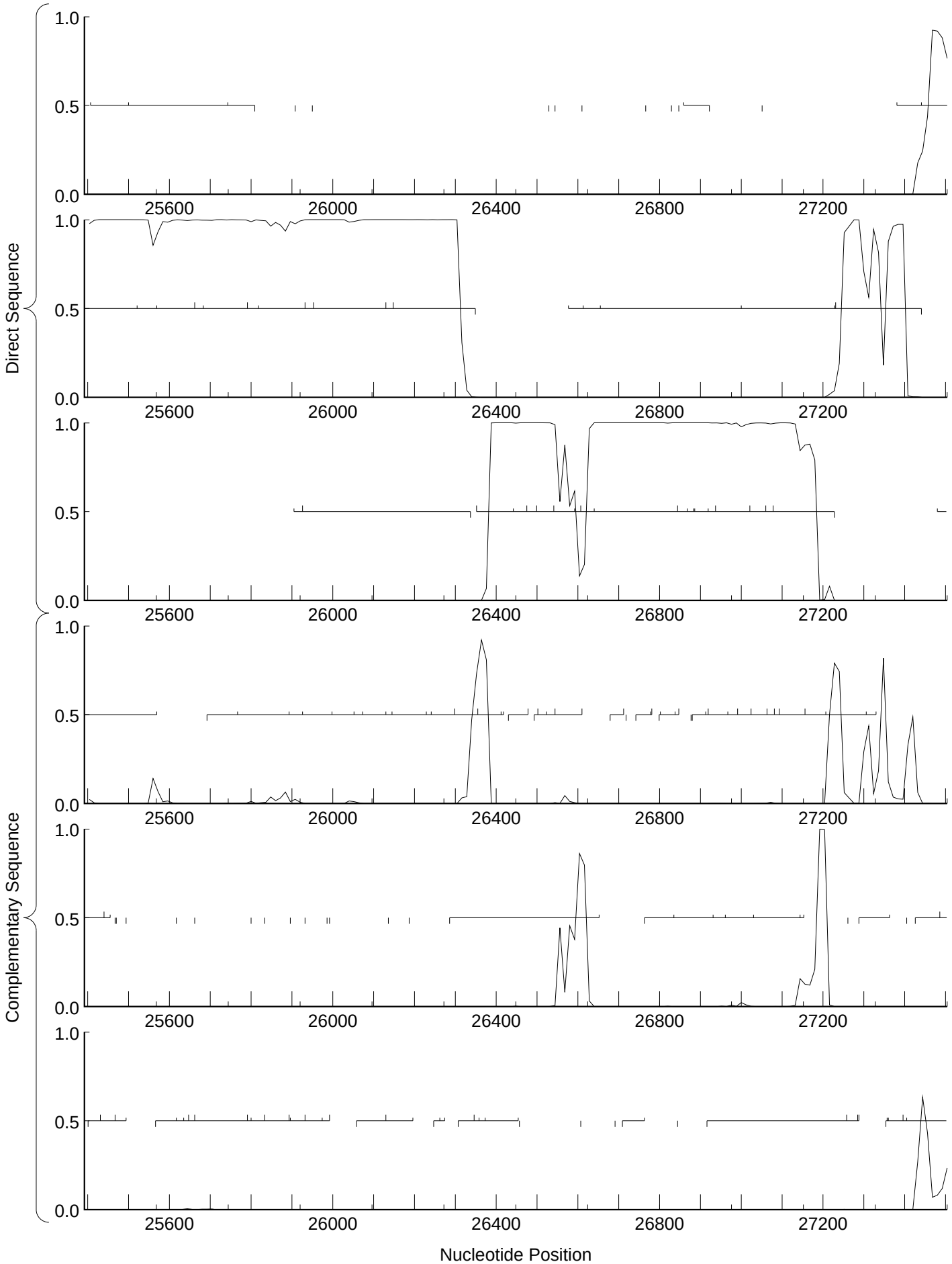


Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 12/27

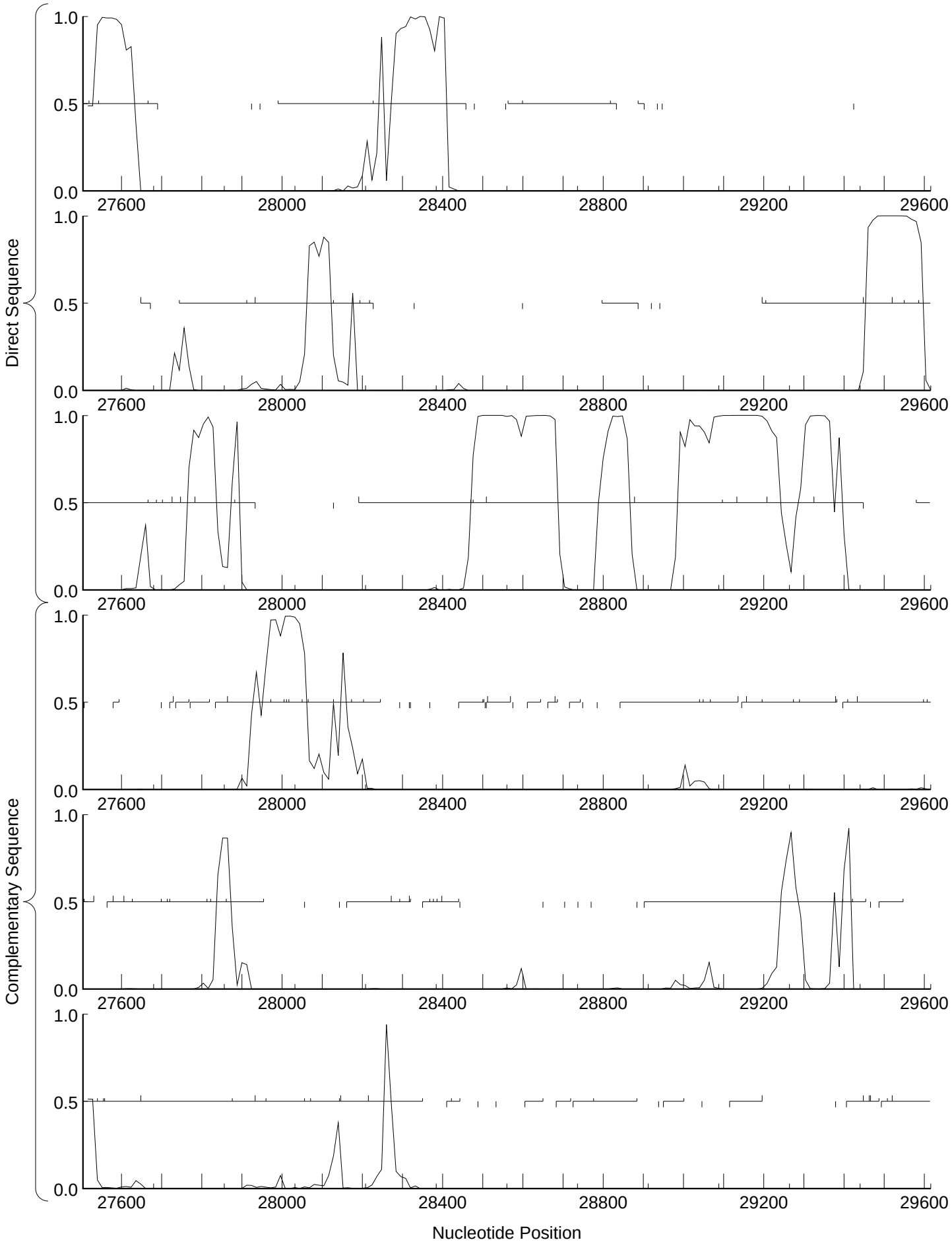



Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 13/27
Direct Sequence
Complementary Sequence
1.0
0.5
0.0
23600
24000
24400
24800
25200
Nucleotide Position

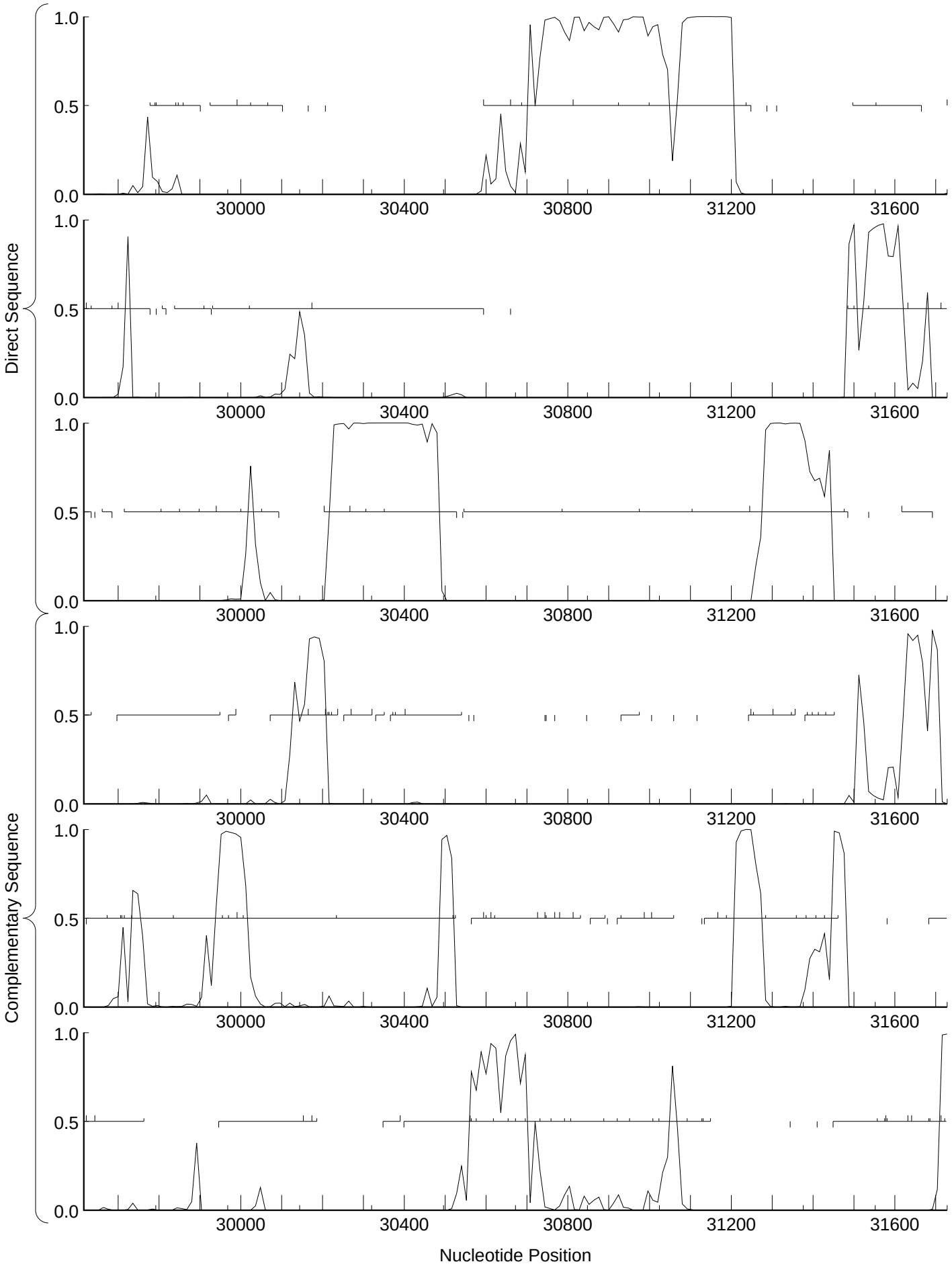
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 14/27



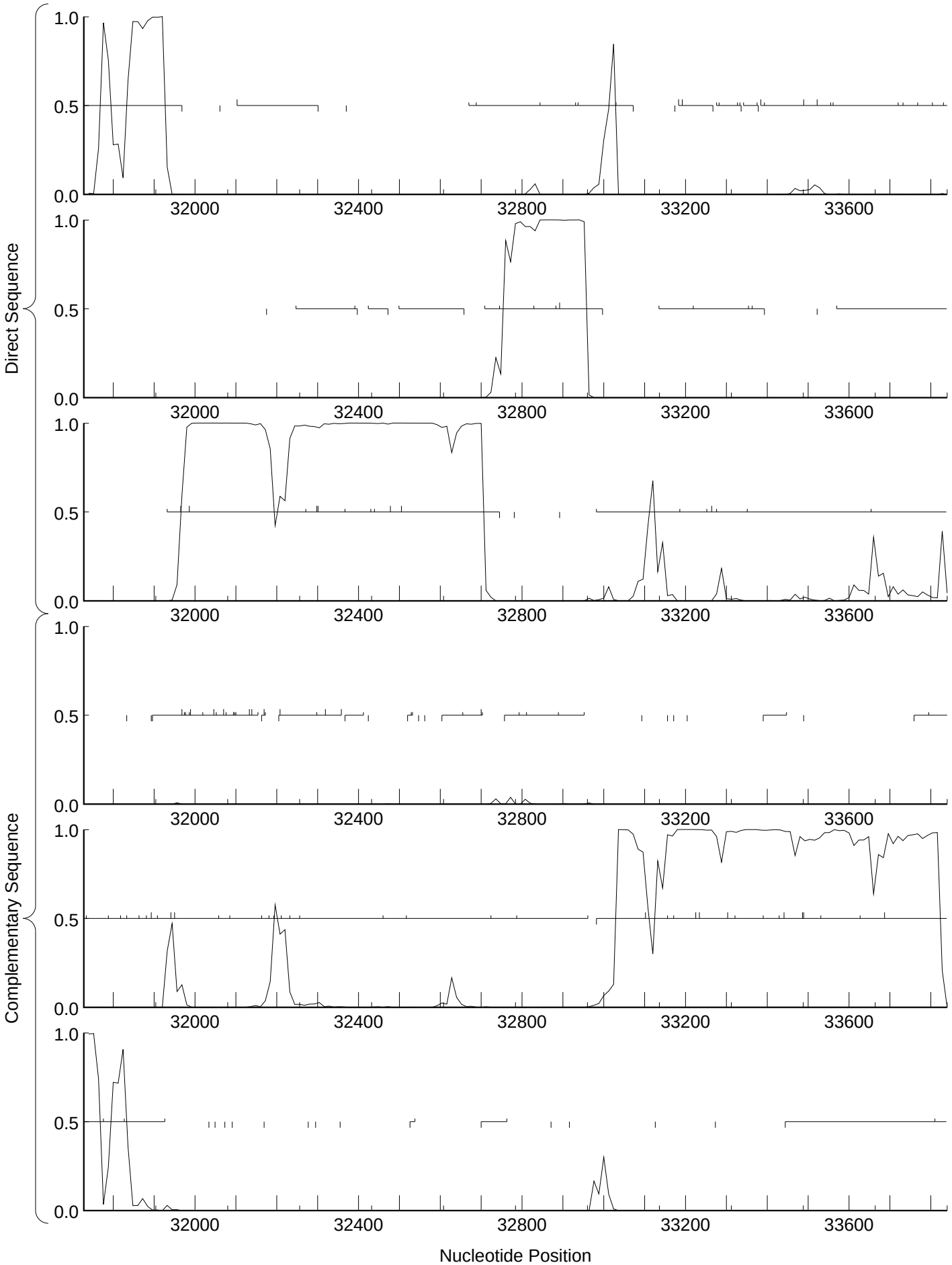
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 15/27



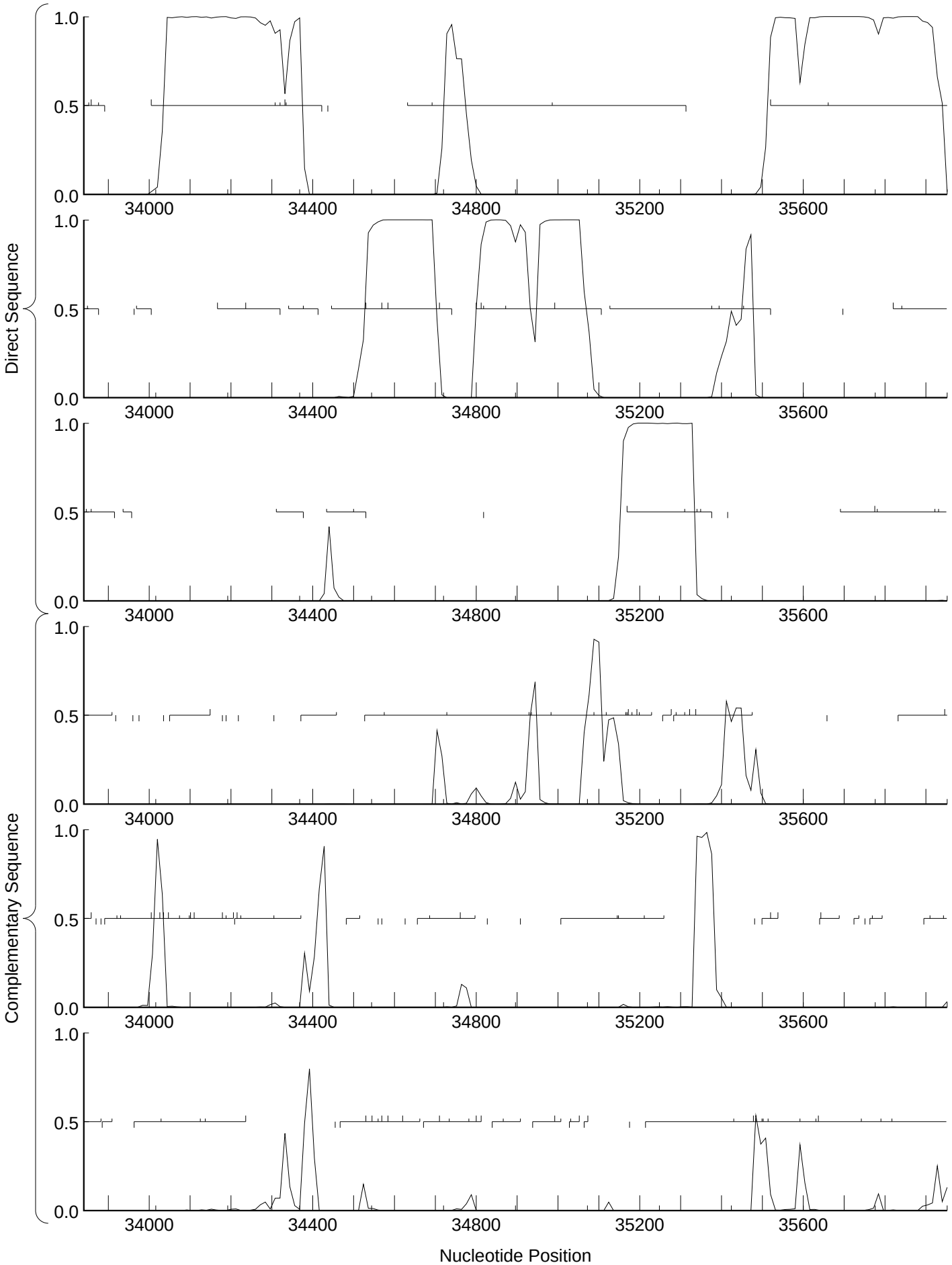
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 16/27



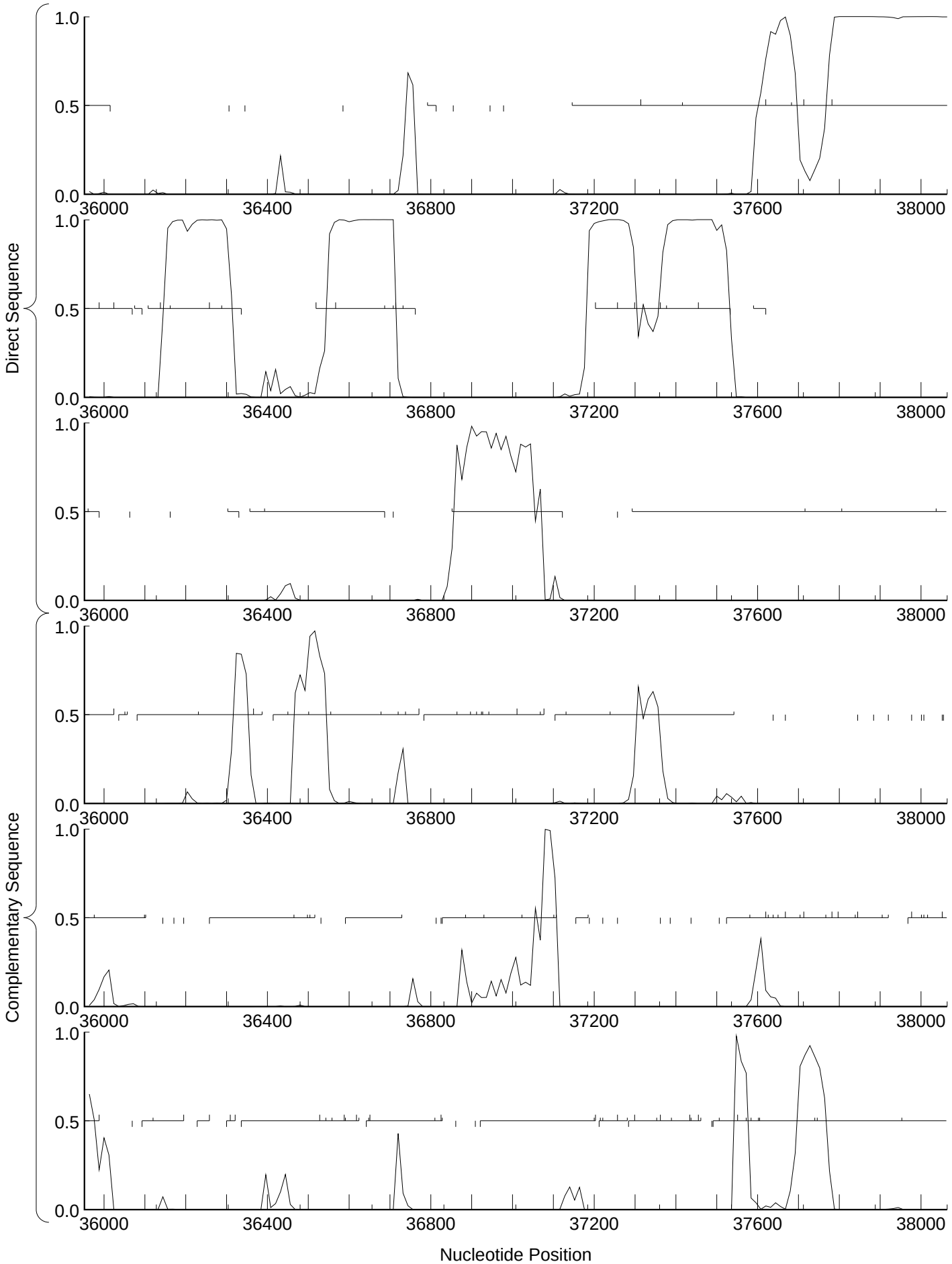
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 17/27



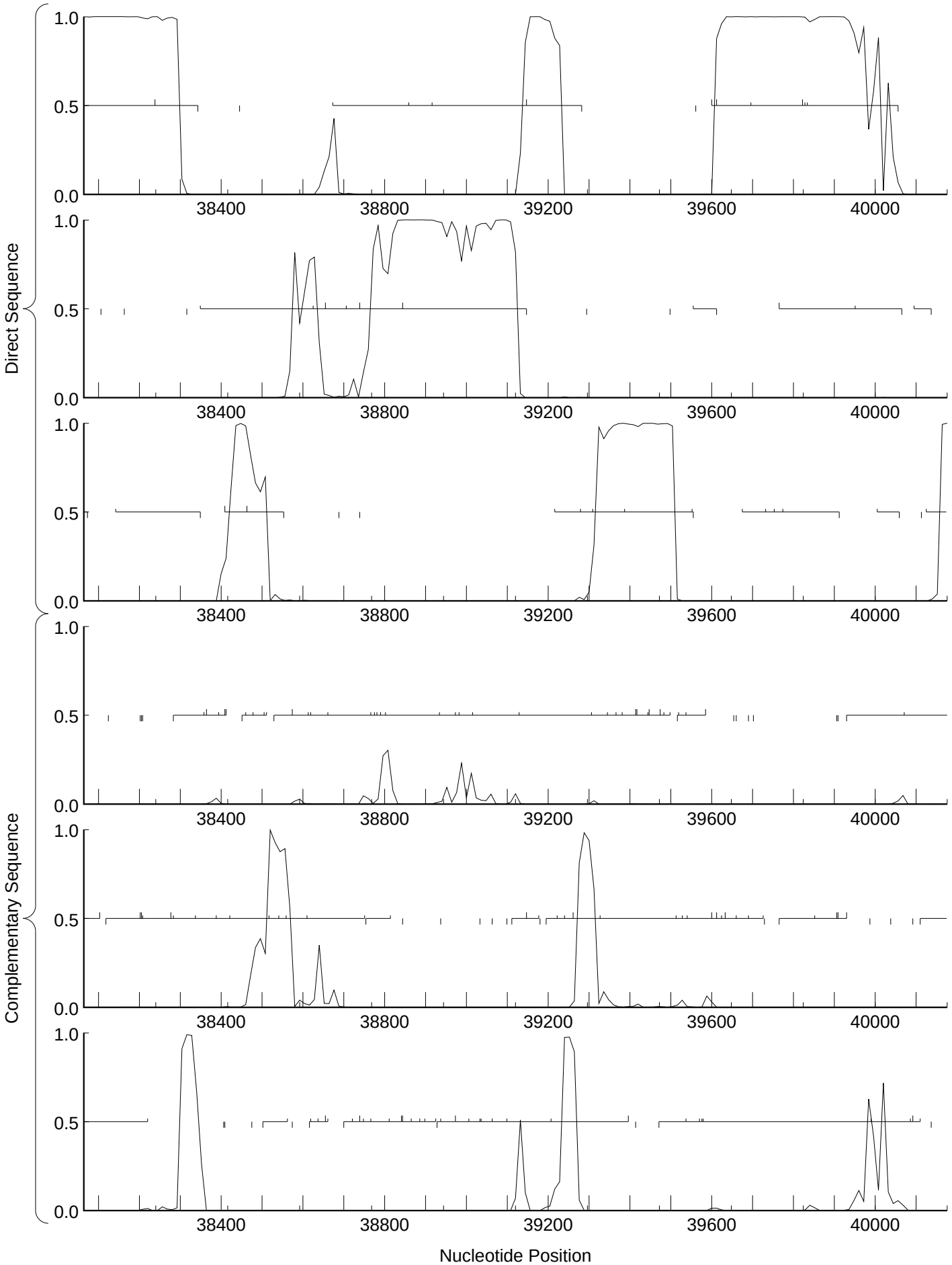
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 18/27



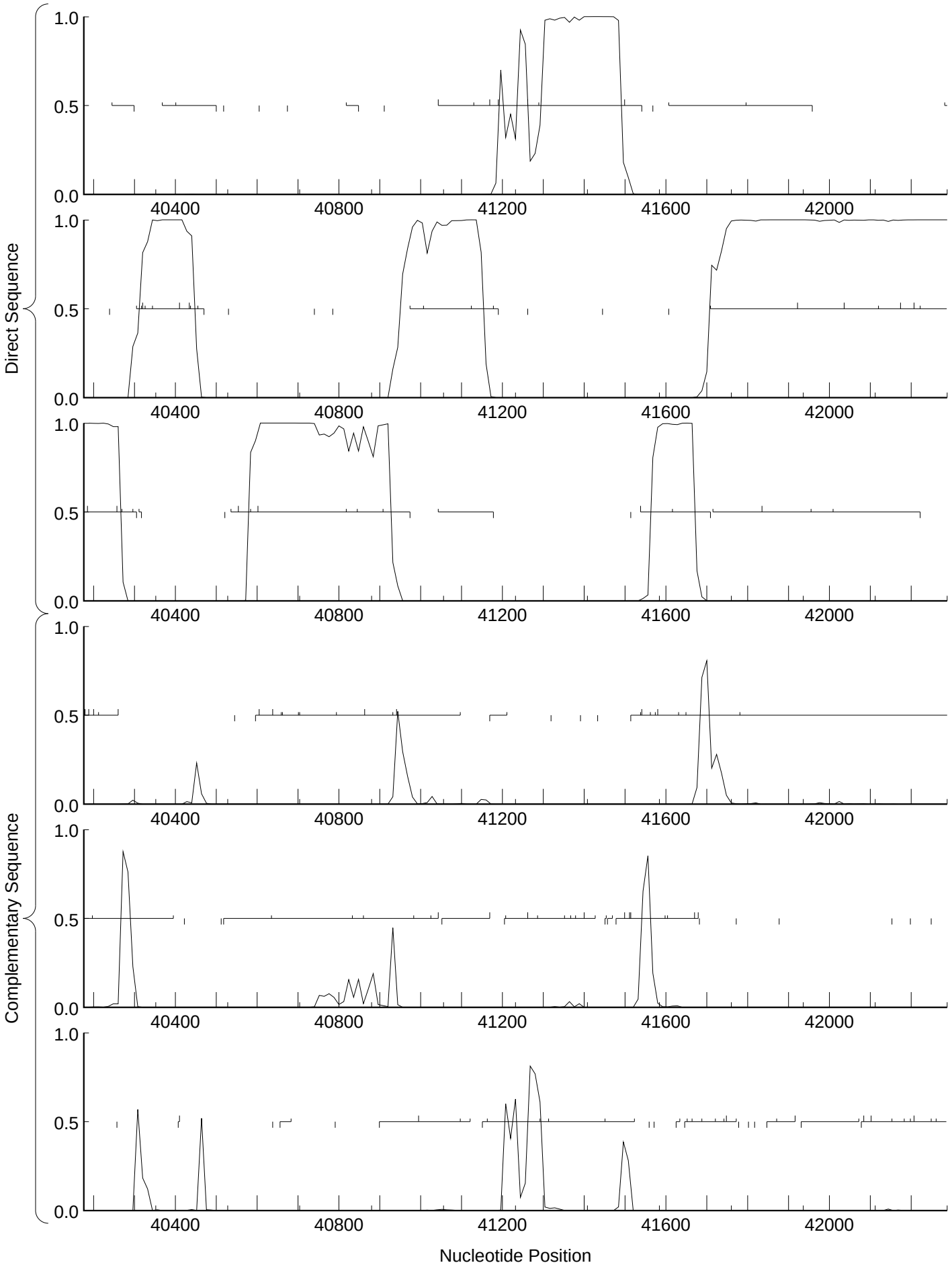
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 19/27



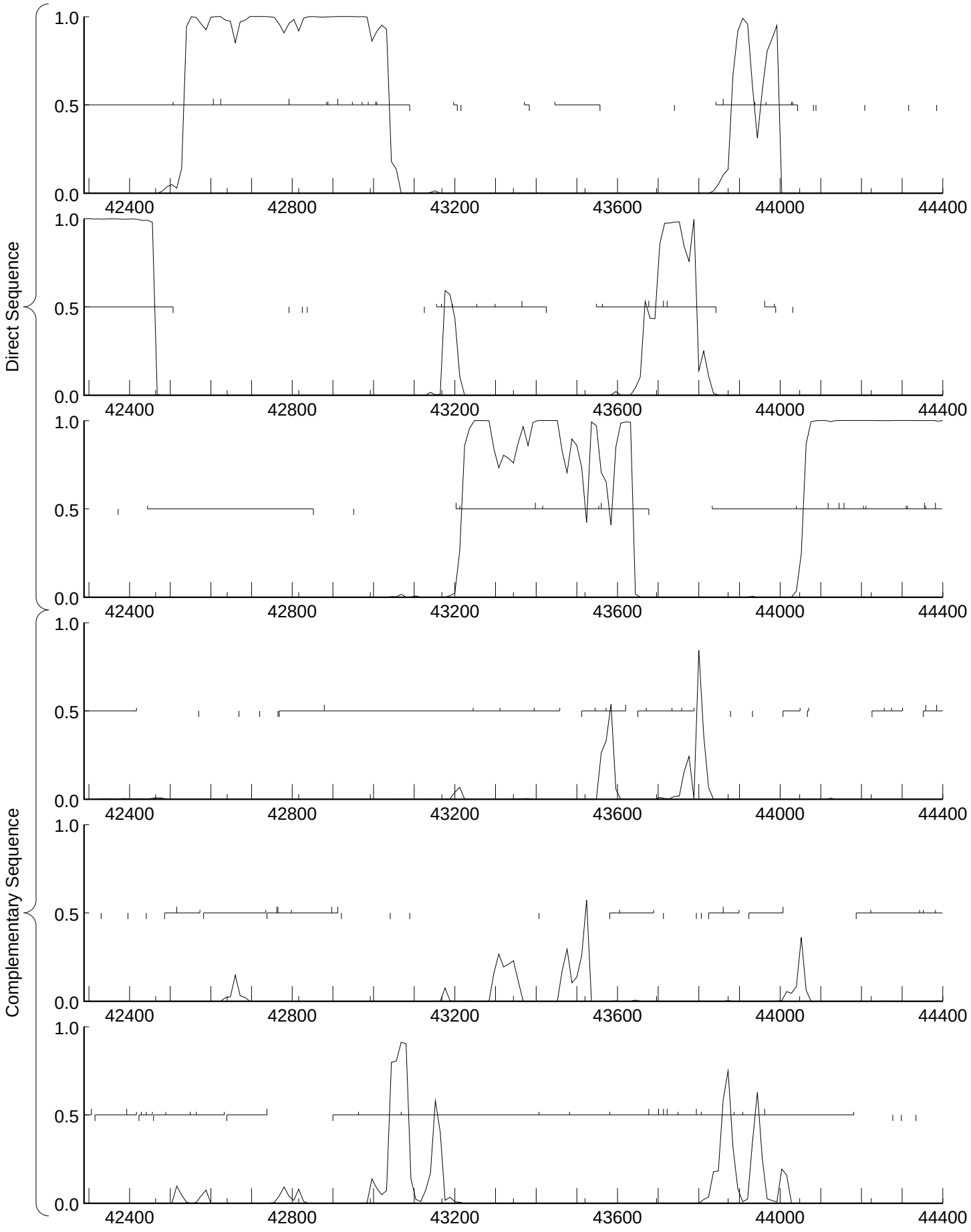
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 20/27



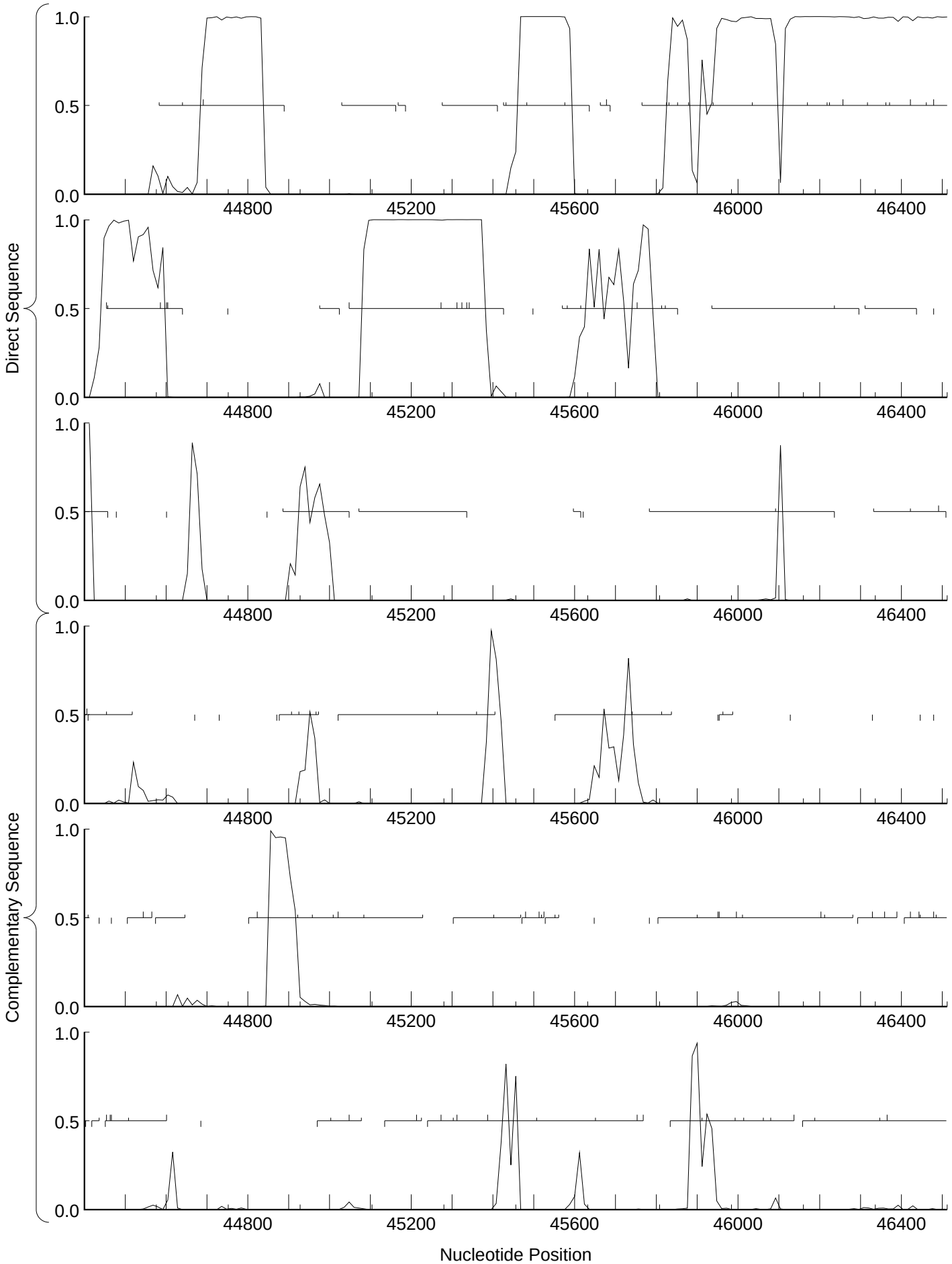
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 21/27



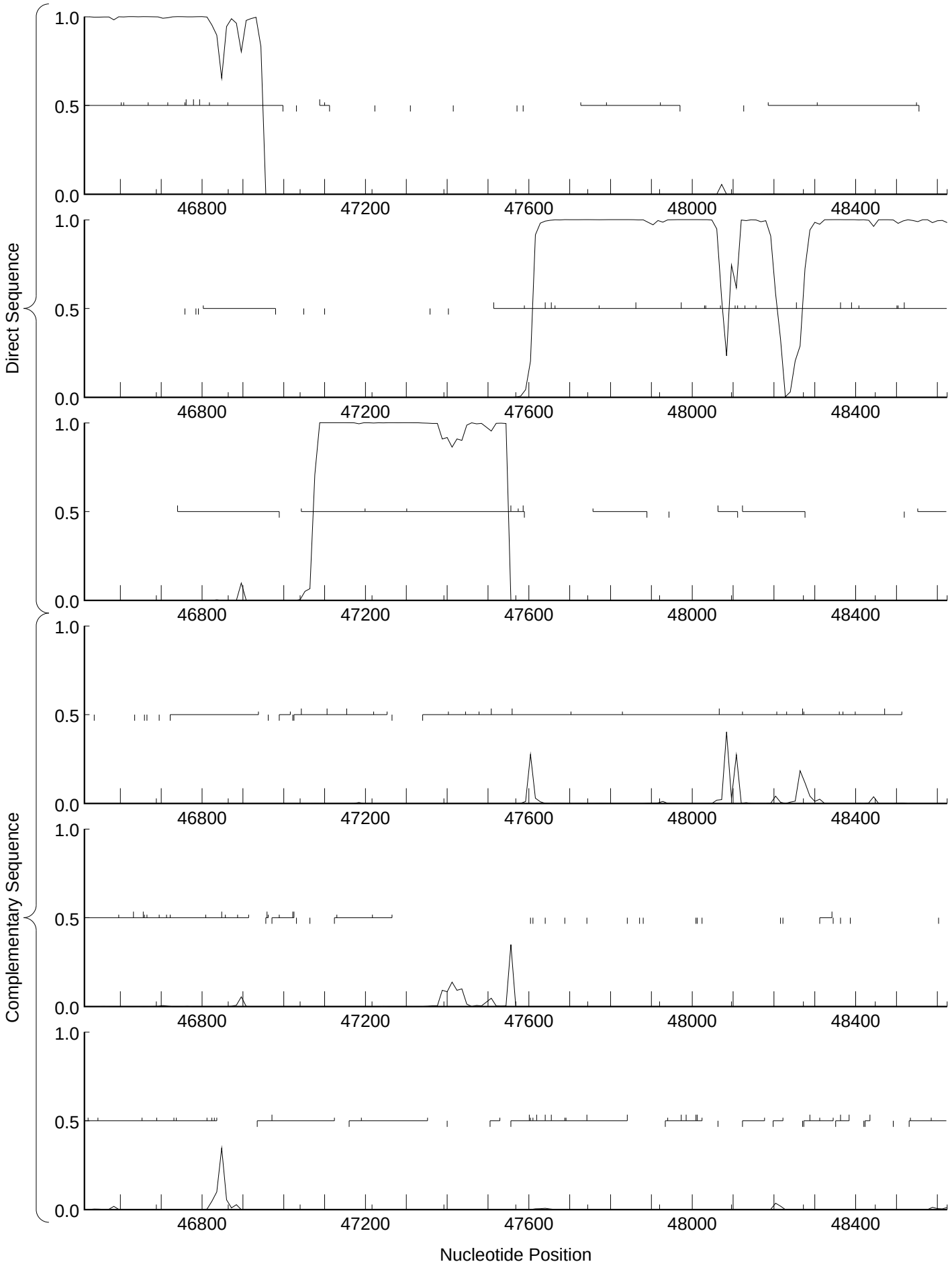
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 22/27



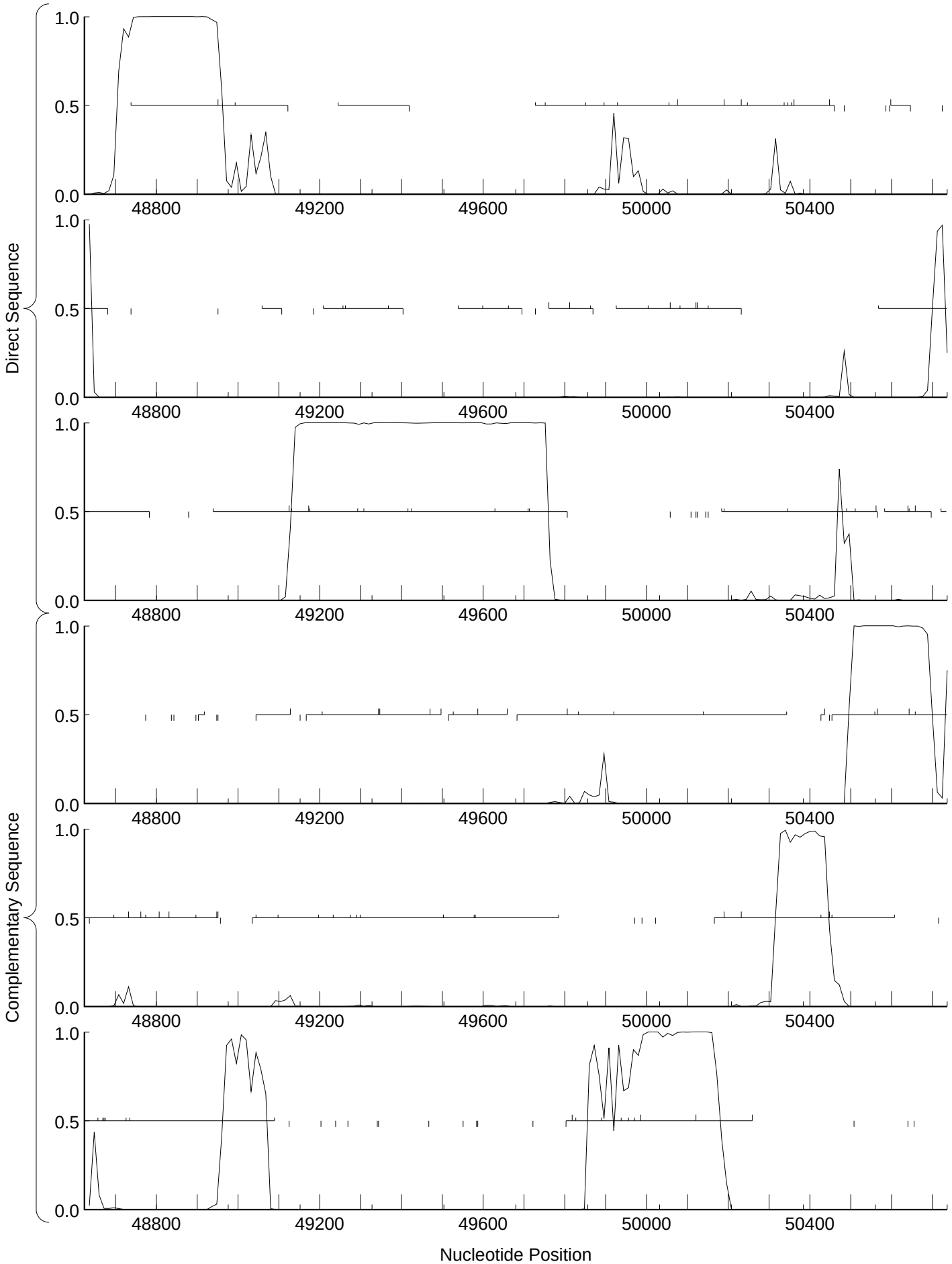
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 23/27



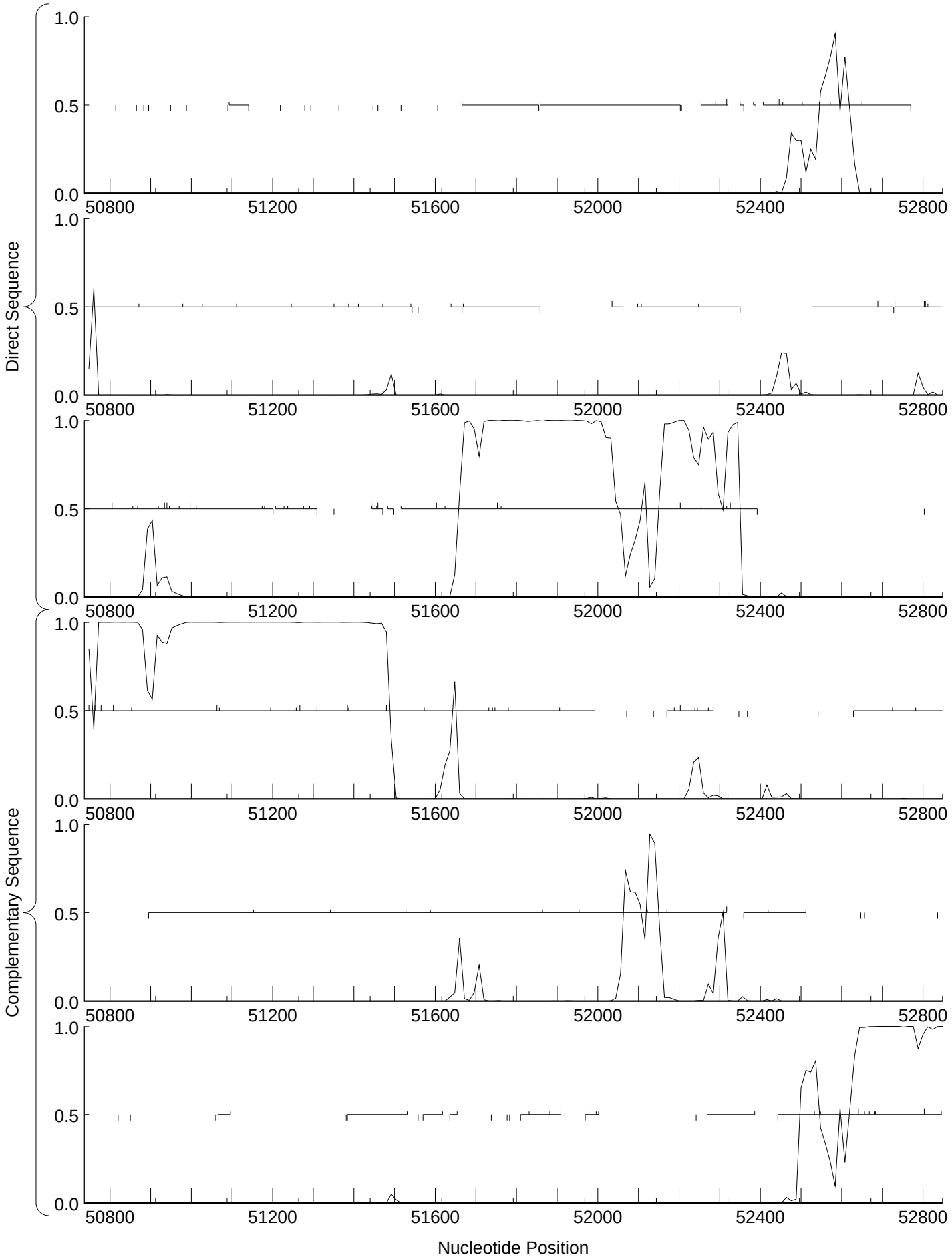
Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 24/27



Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 25/27



Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 26/27



Rhodococcus phage Jace complete genome, 53912 bp, circularly permuted, Order 4, Window 96, Step 12, 27/27

