# GeneMark

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

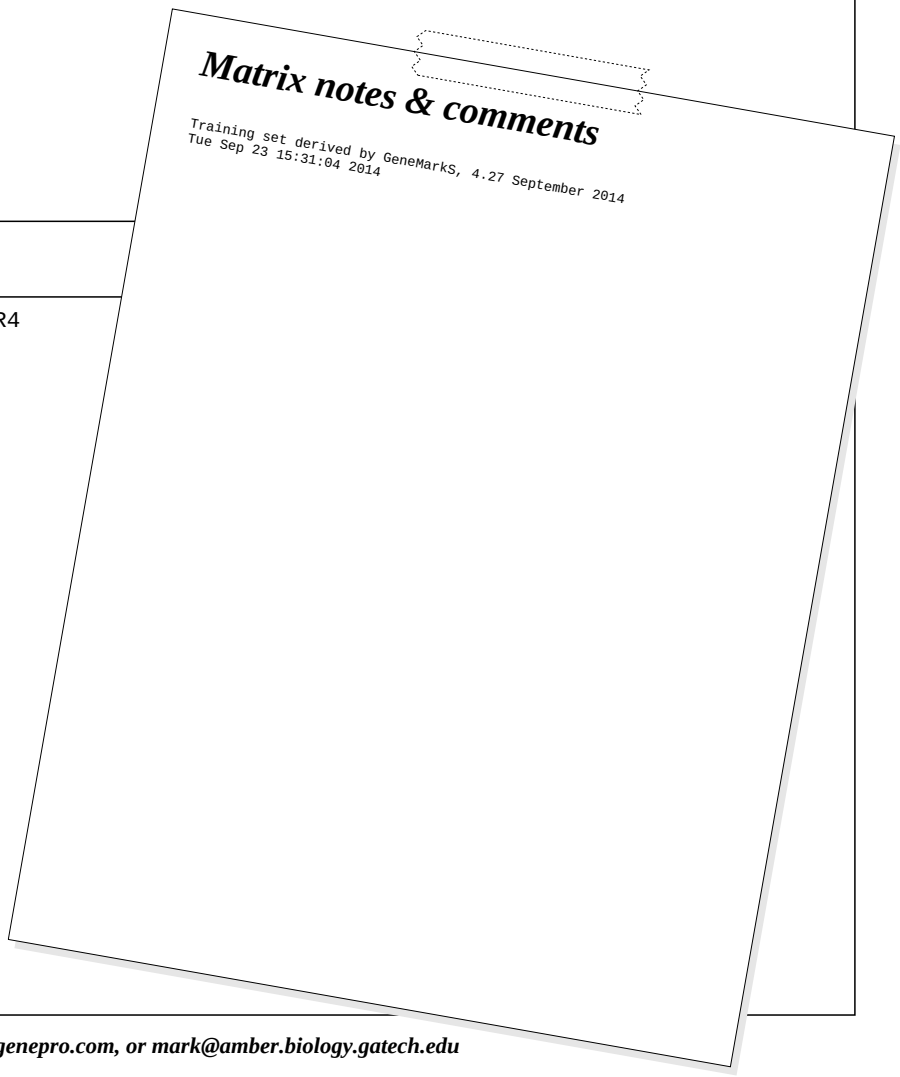
## PROGRAM INFORMATION

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Analysis Date    :  3/28/24 at 7:58:07
Pages            :  37
Sequence Length  :  75772 bp
GC Content       :   56.66%

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

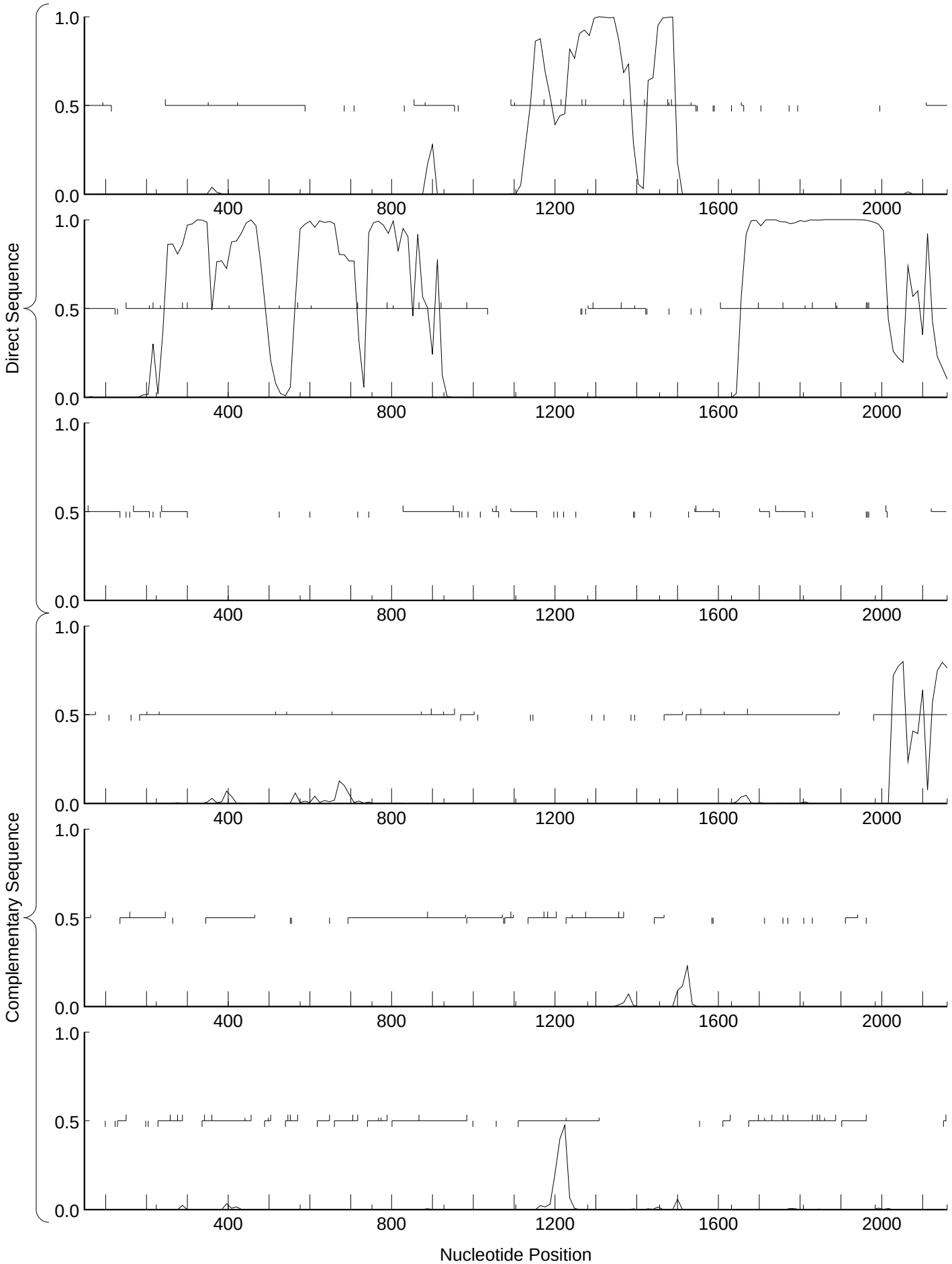
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Author     :  -

Order      :  4
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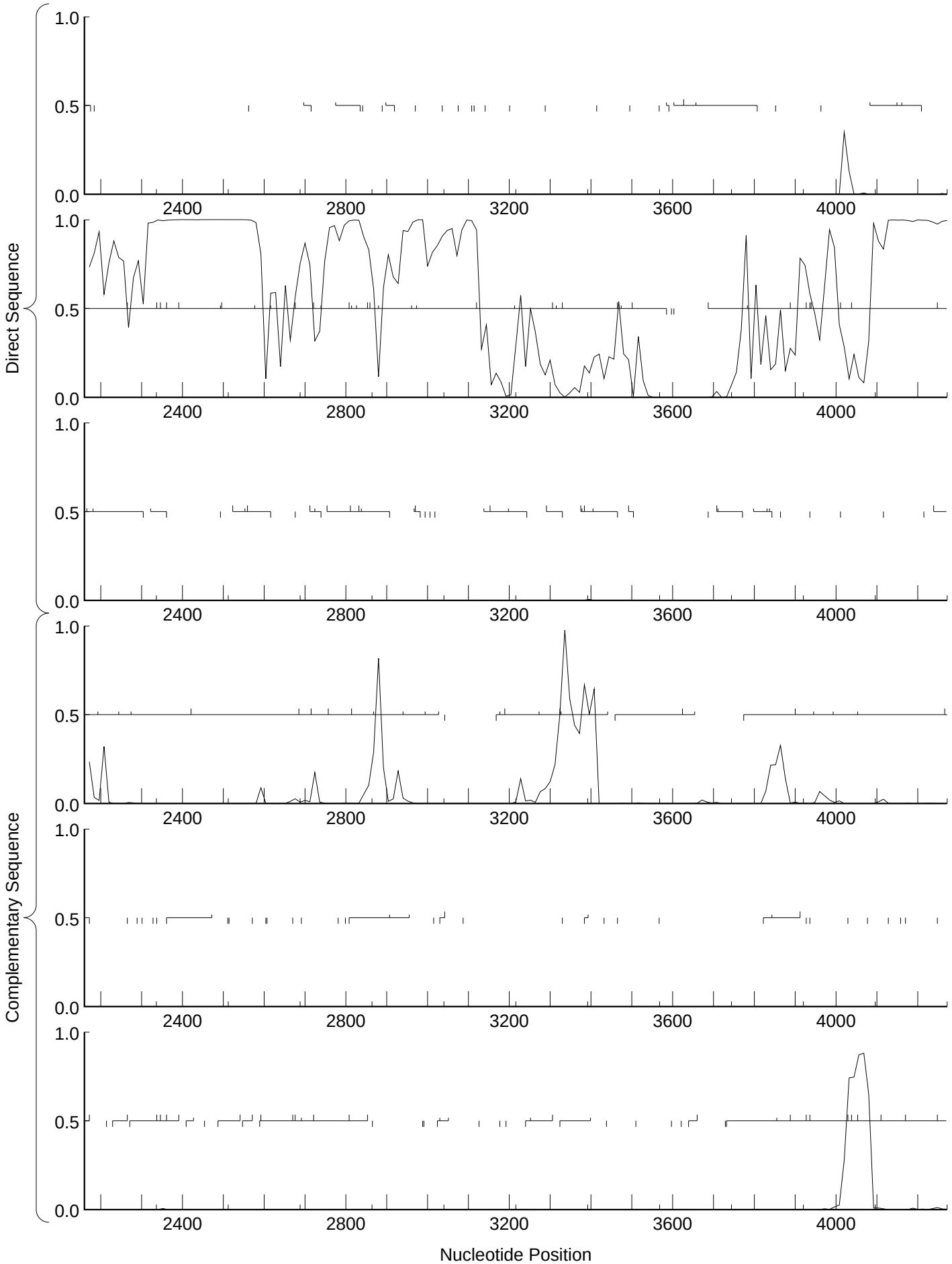

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

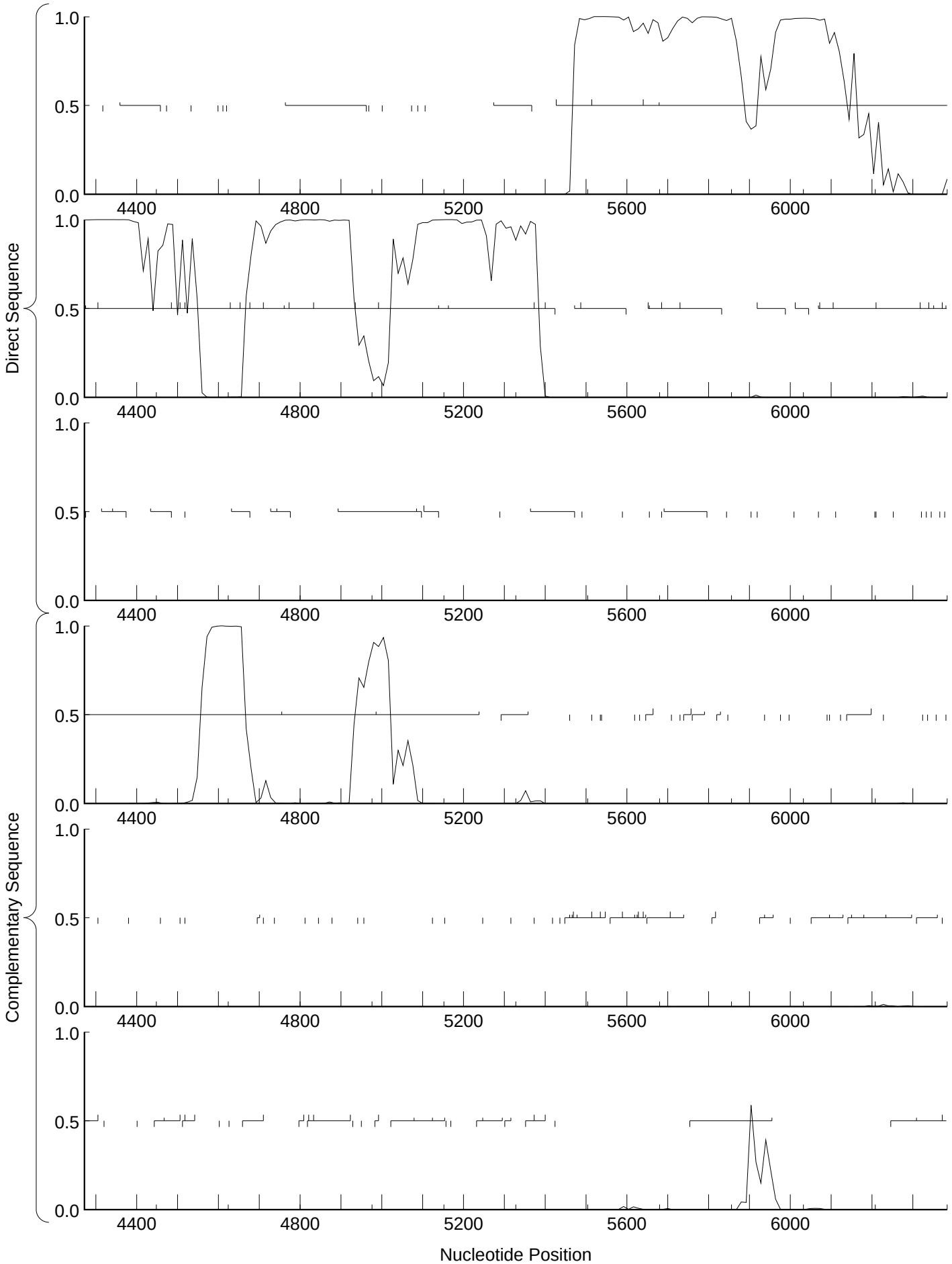
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 2/37



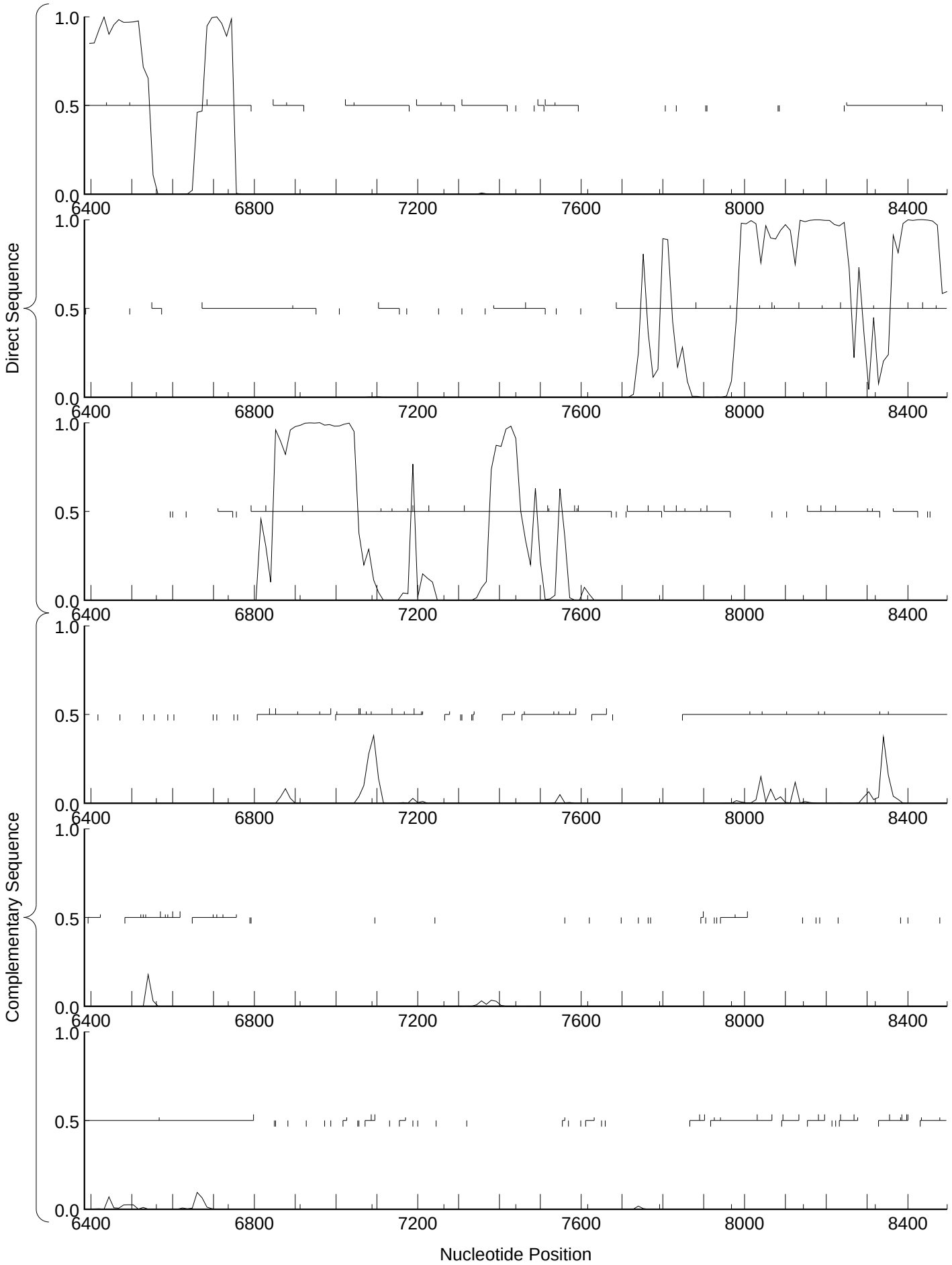
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 3/37



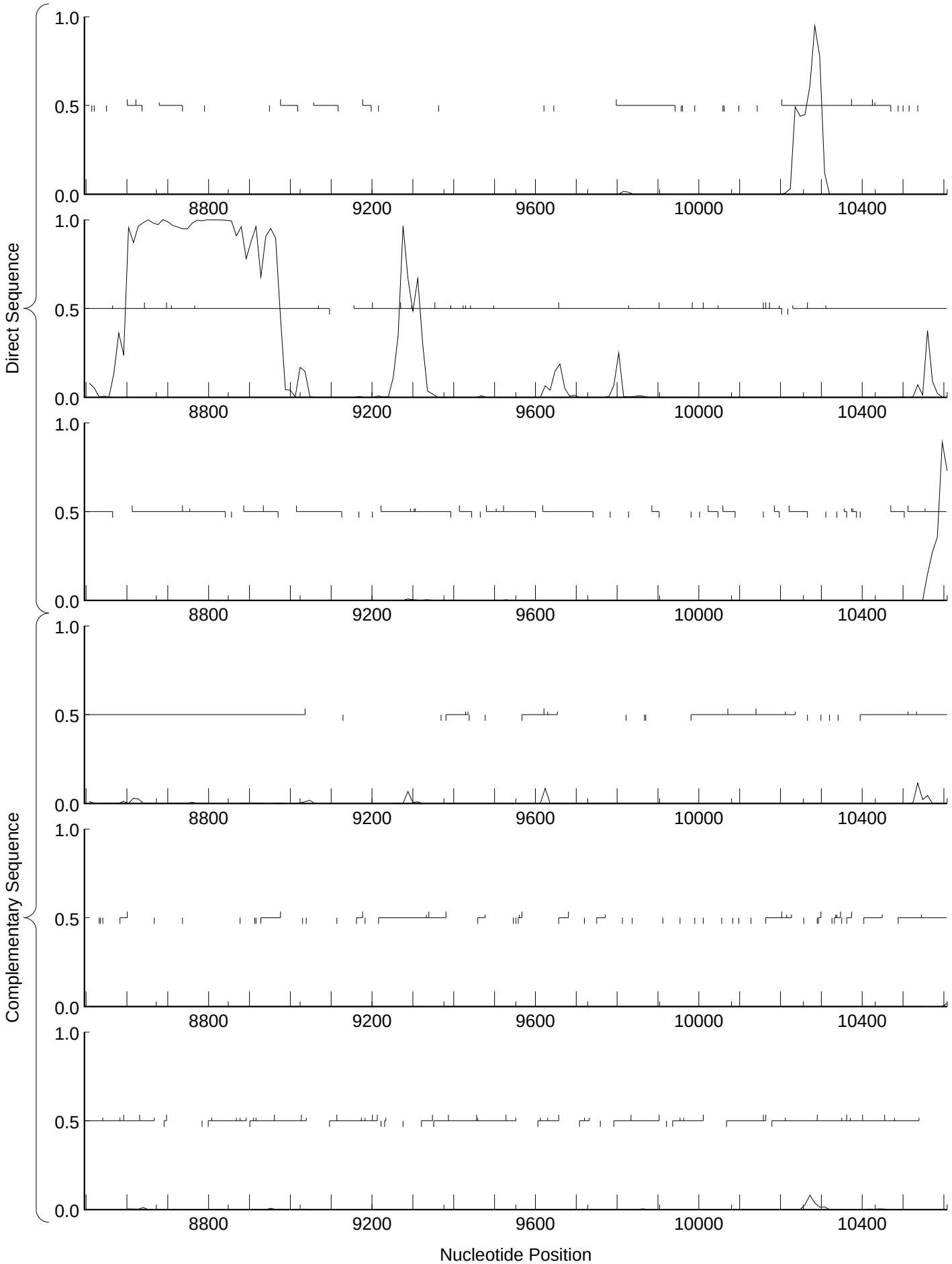
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 4/37



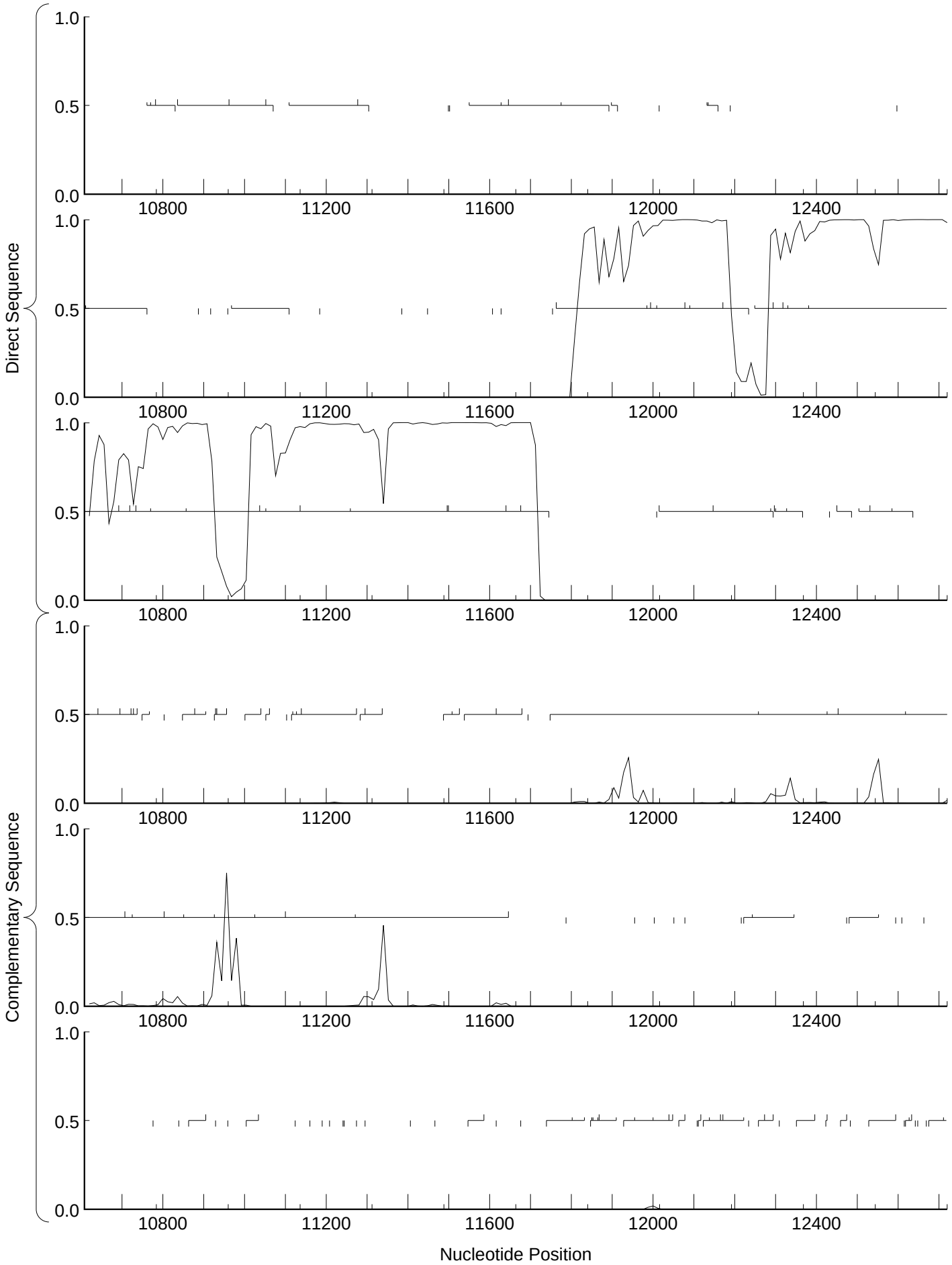
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 5/37



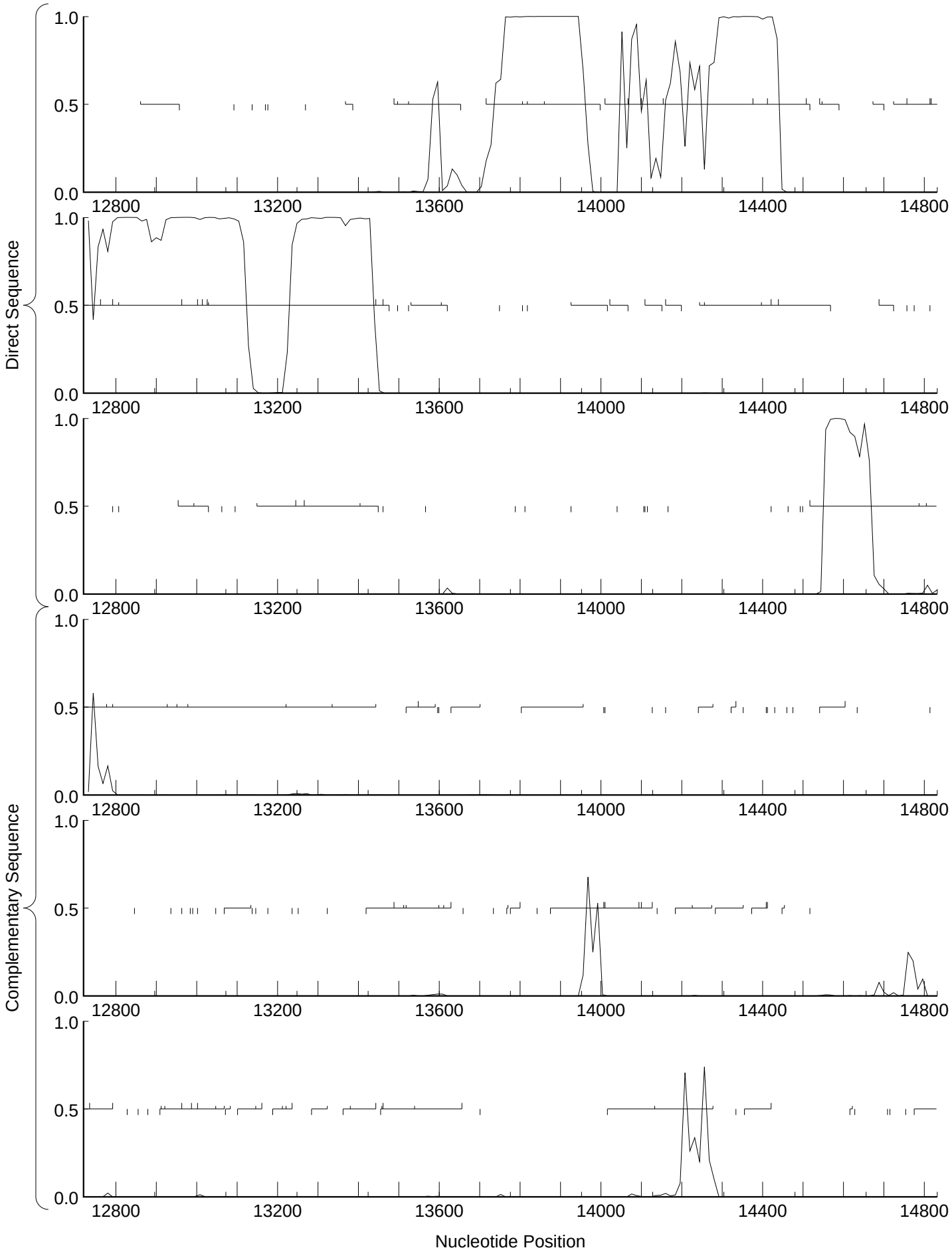
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 6/37



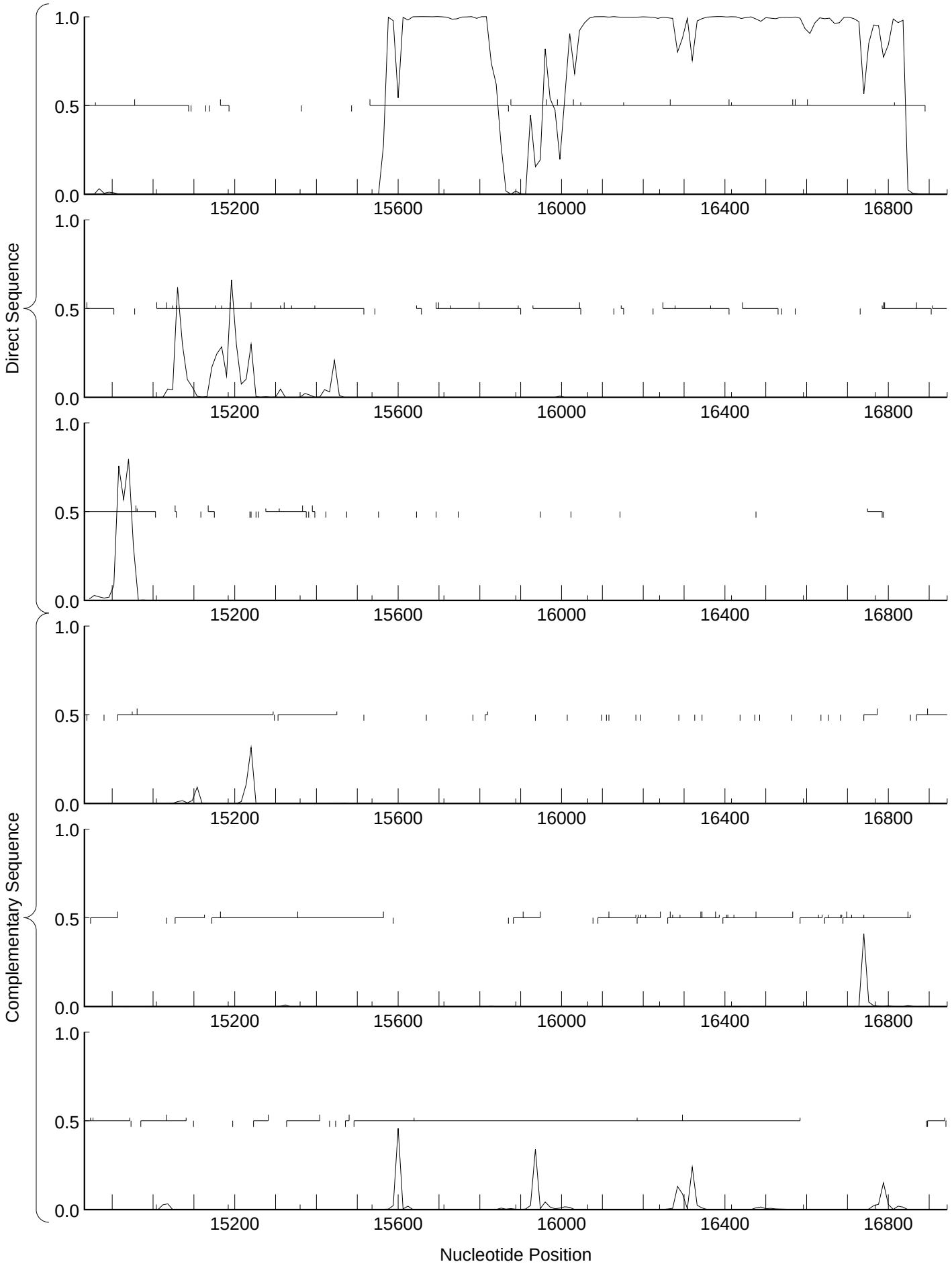
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 7/37



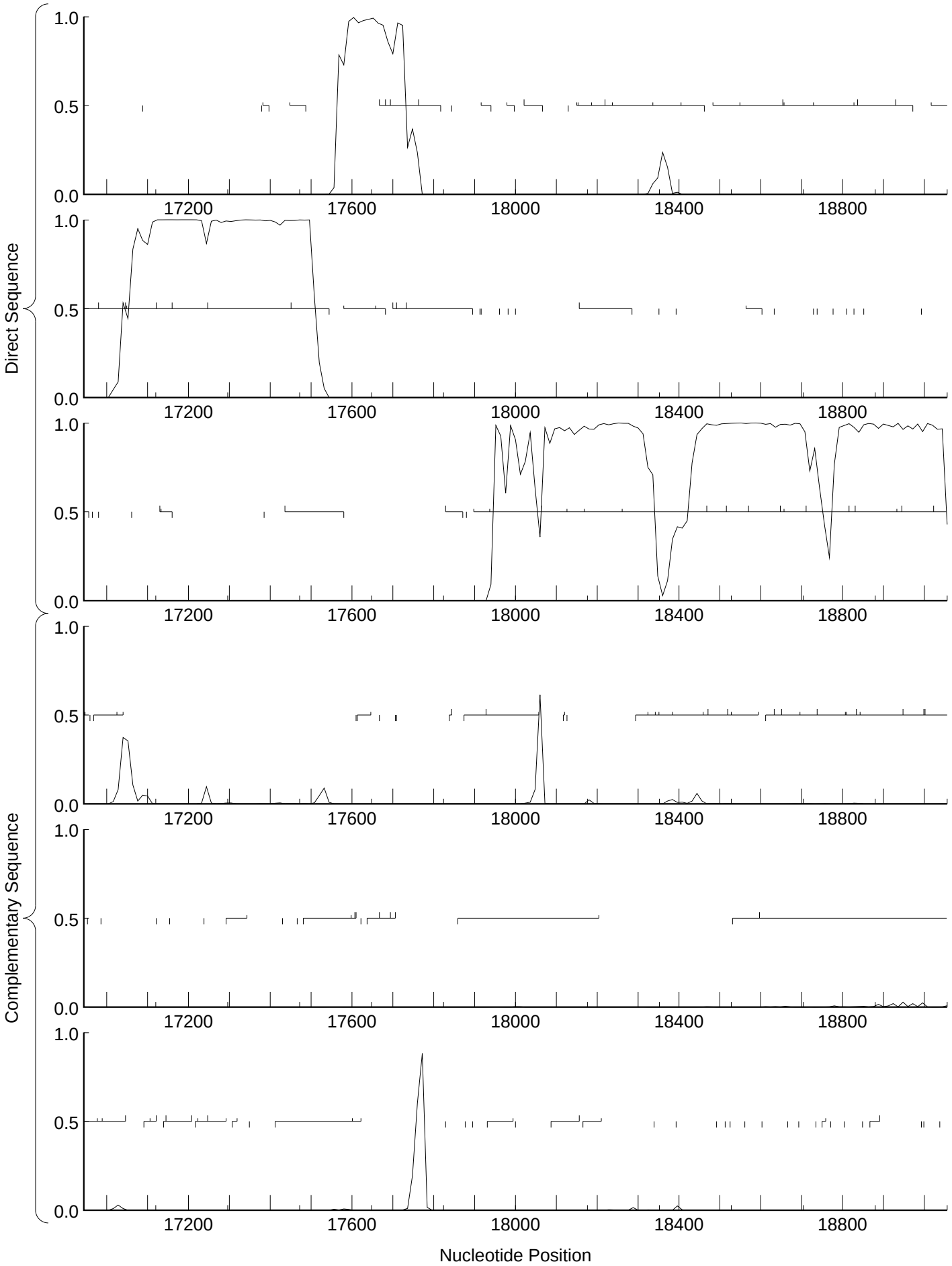
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 8/37



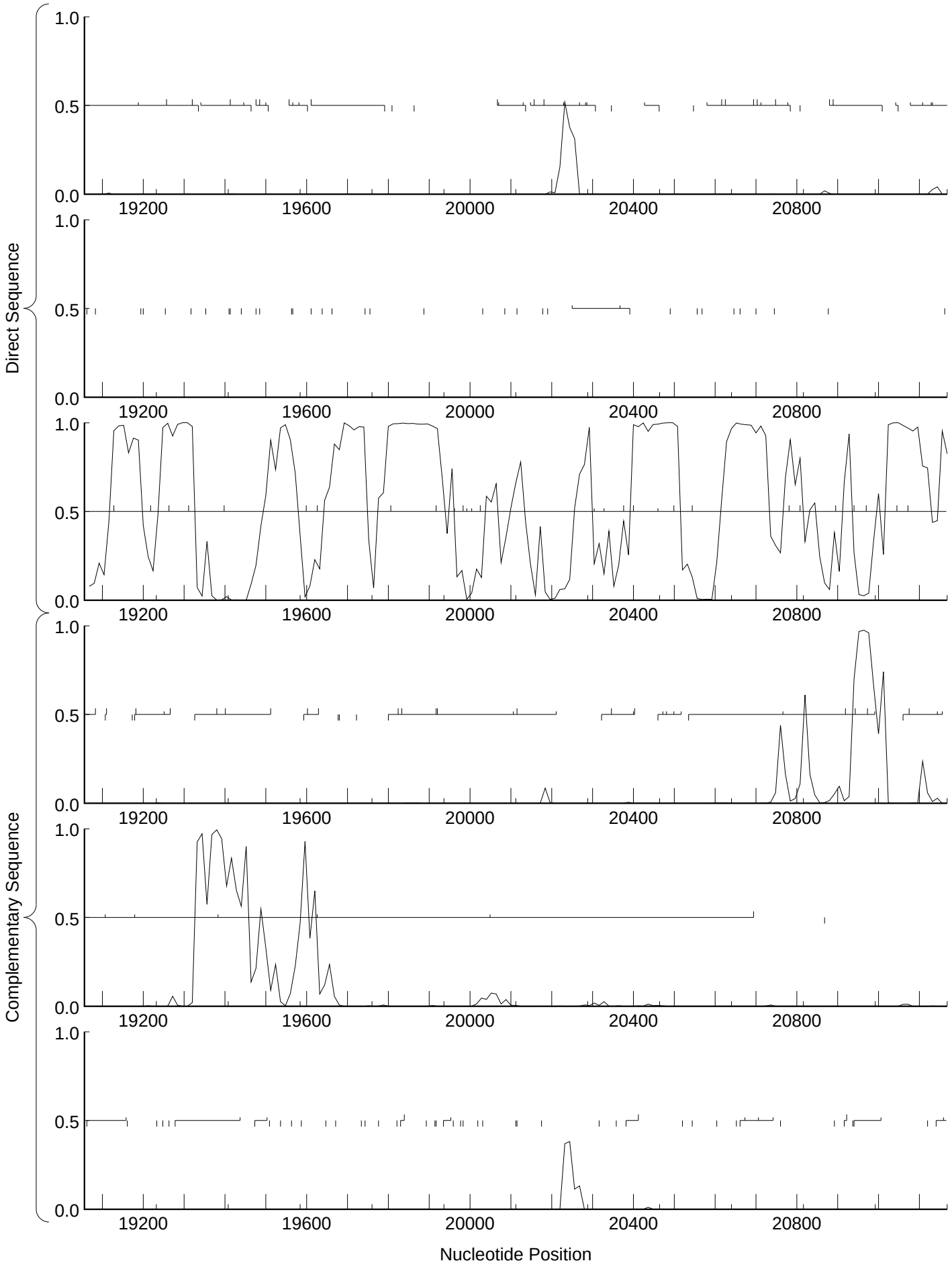
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 9/37



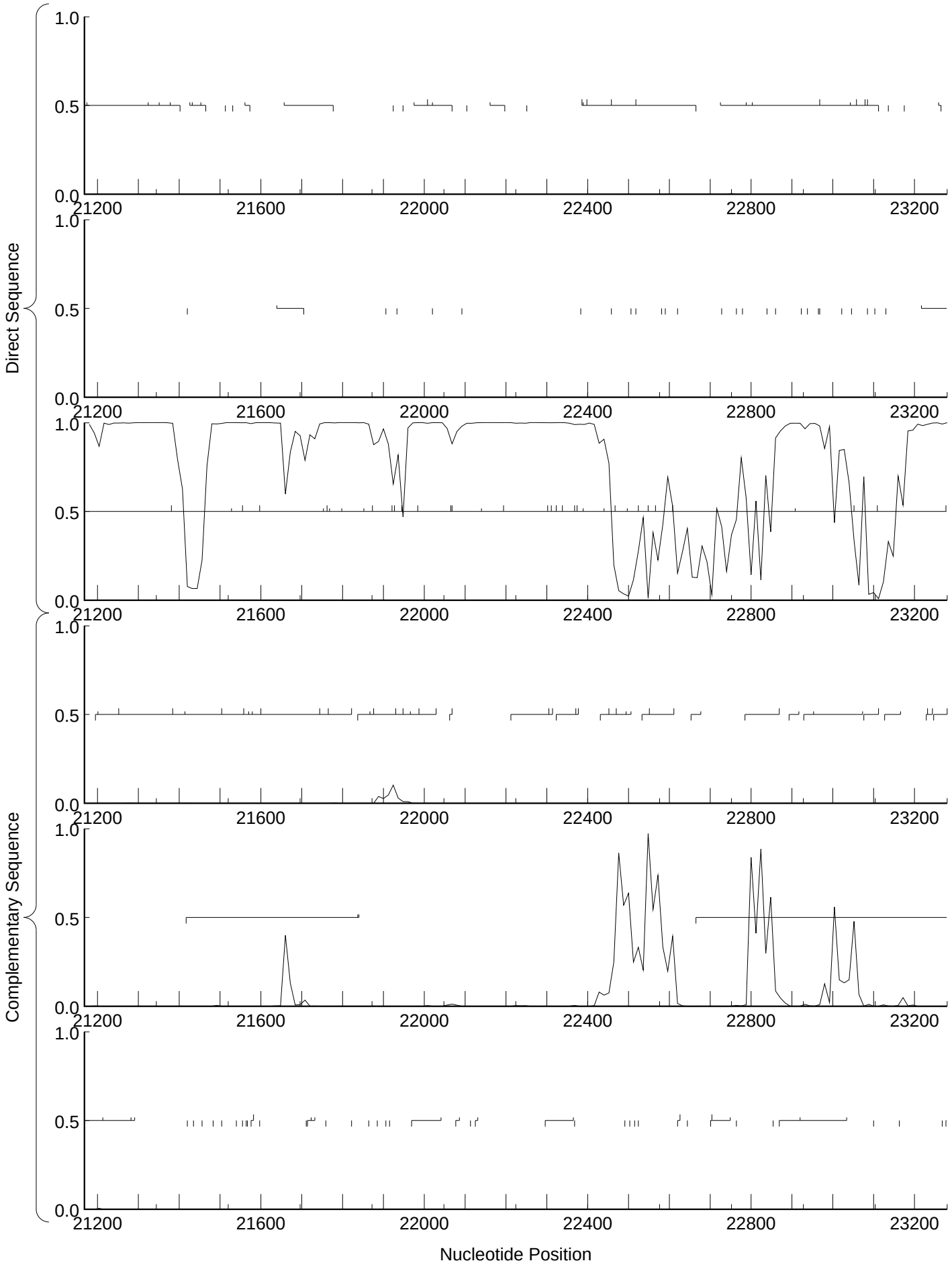
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 10/37



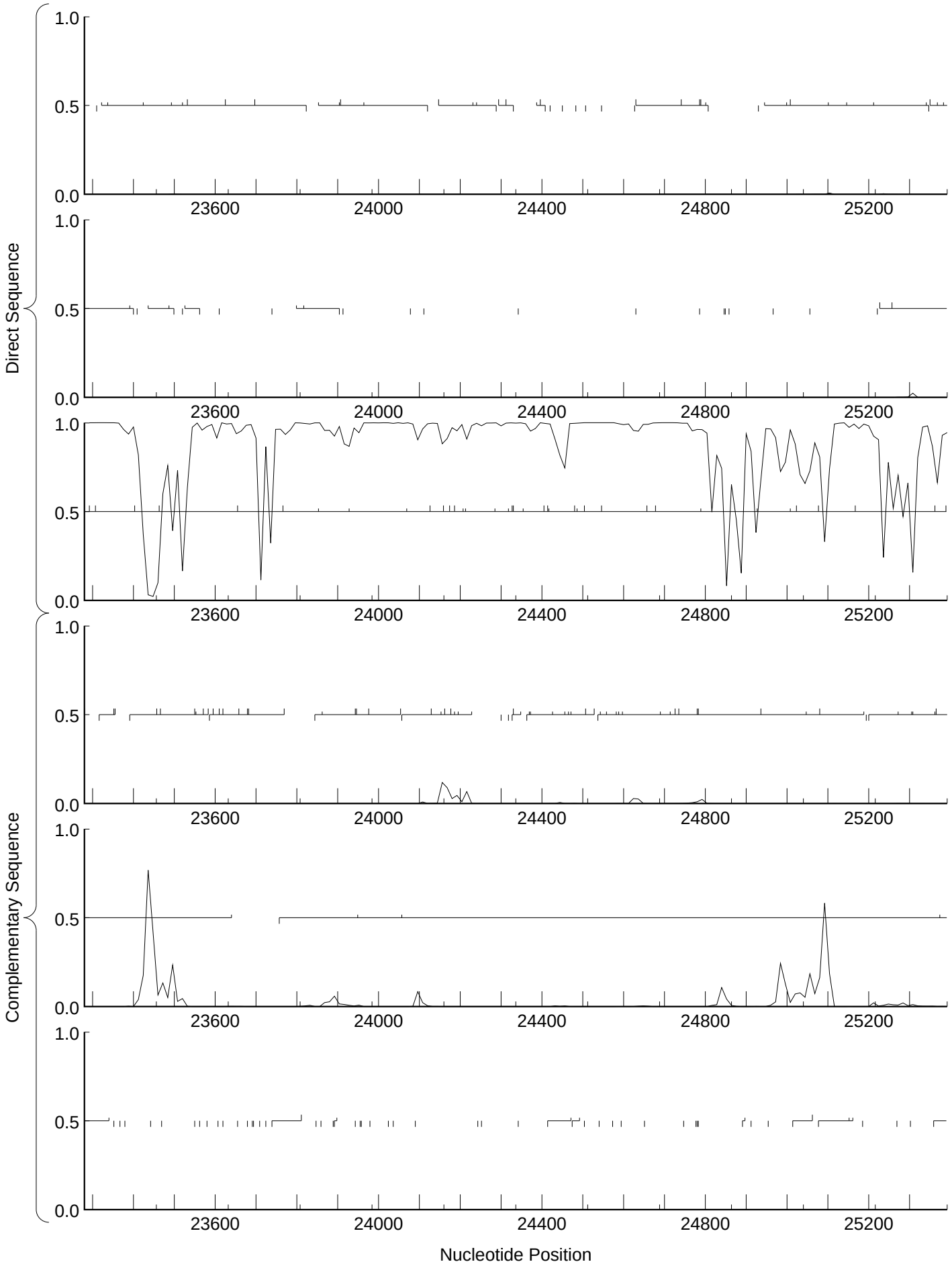
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 11/37



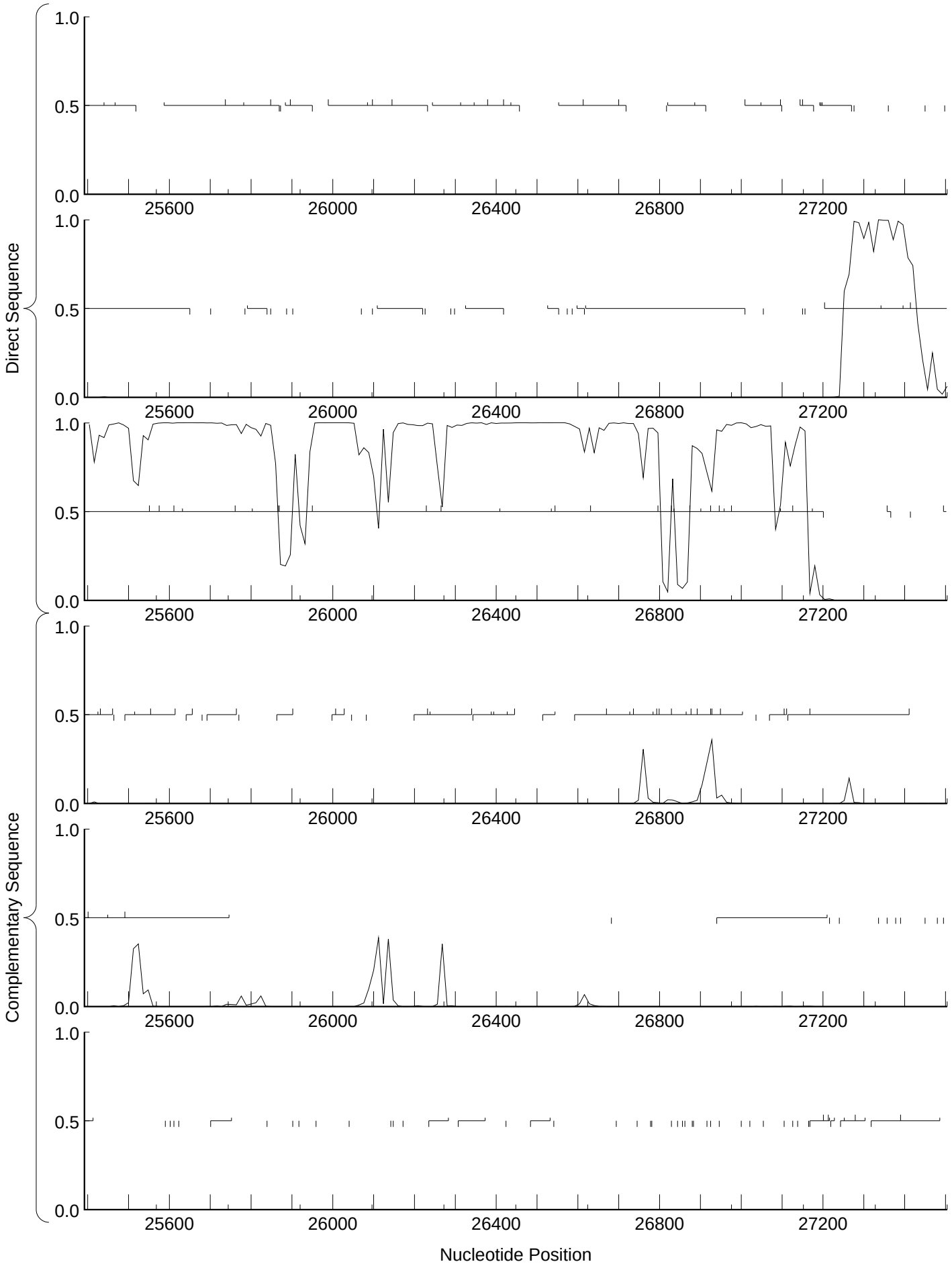
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 12/37



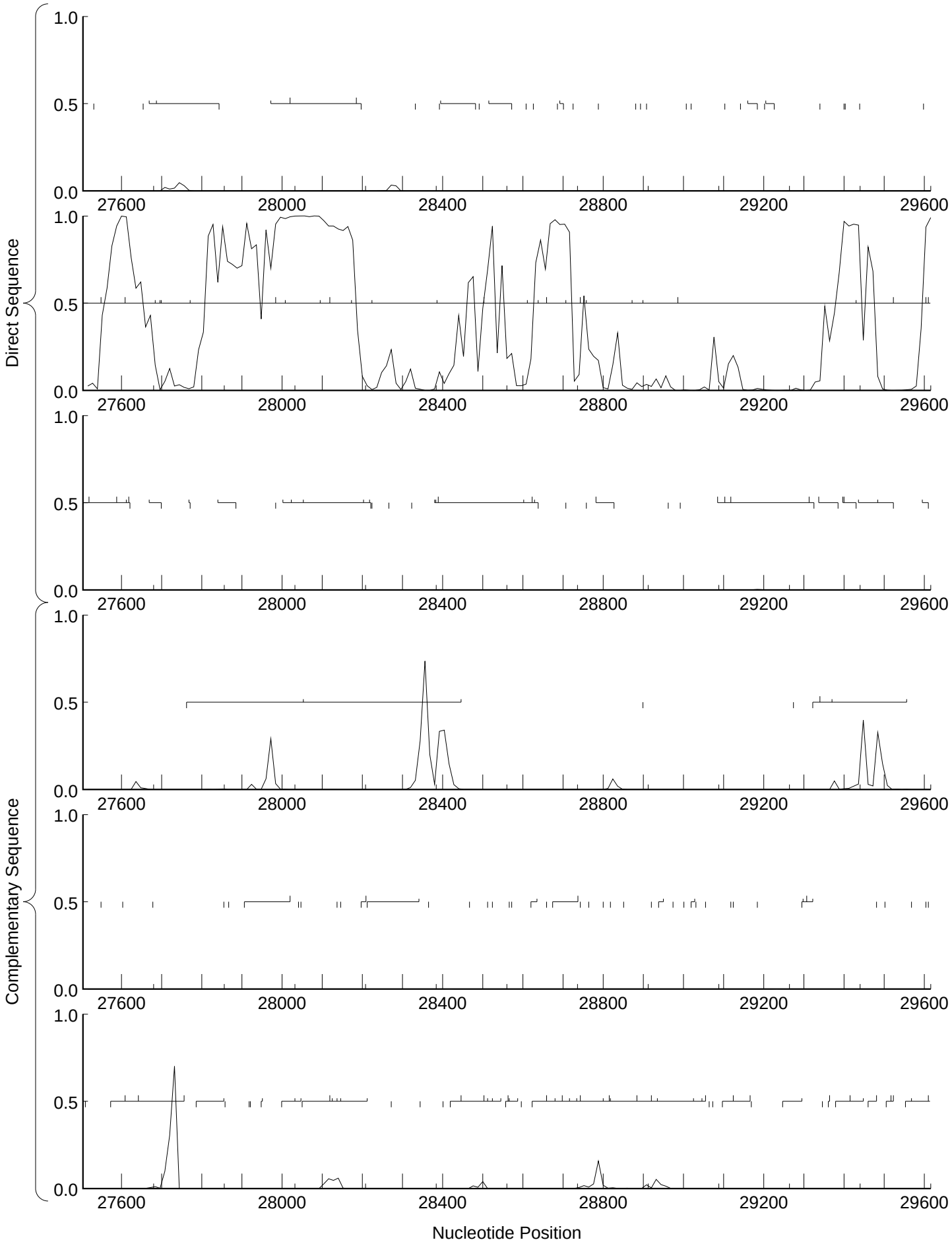
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 13/37

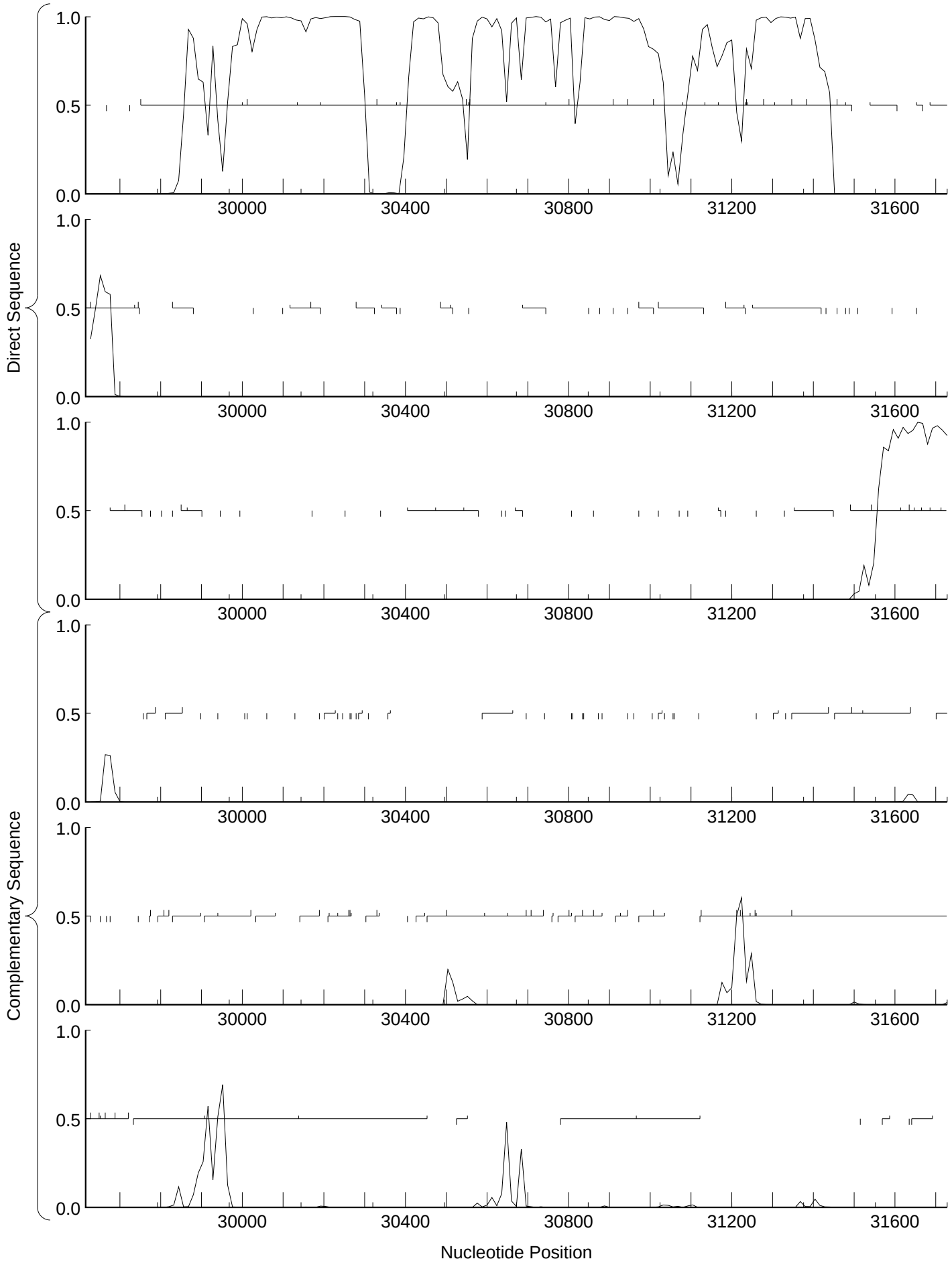


Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 14/37

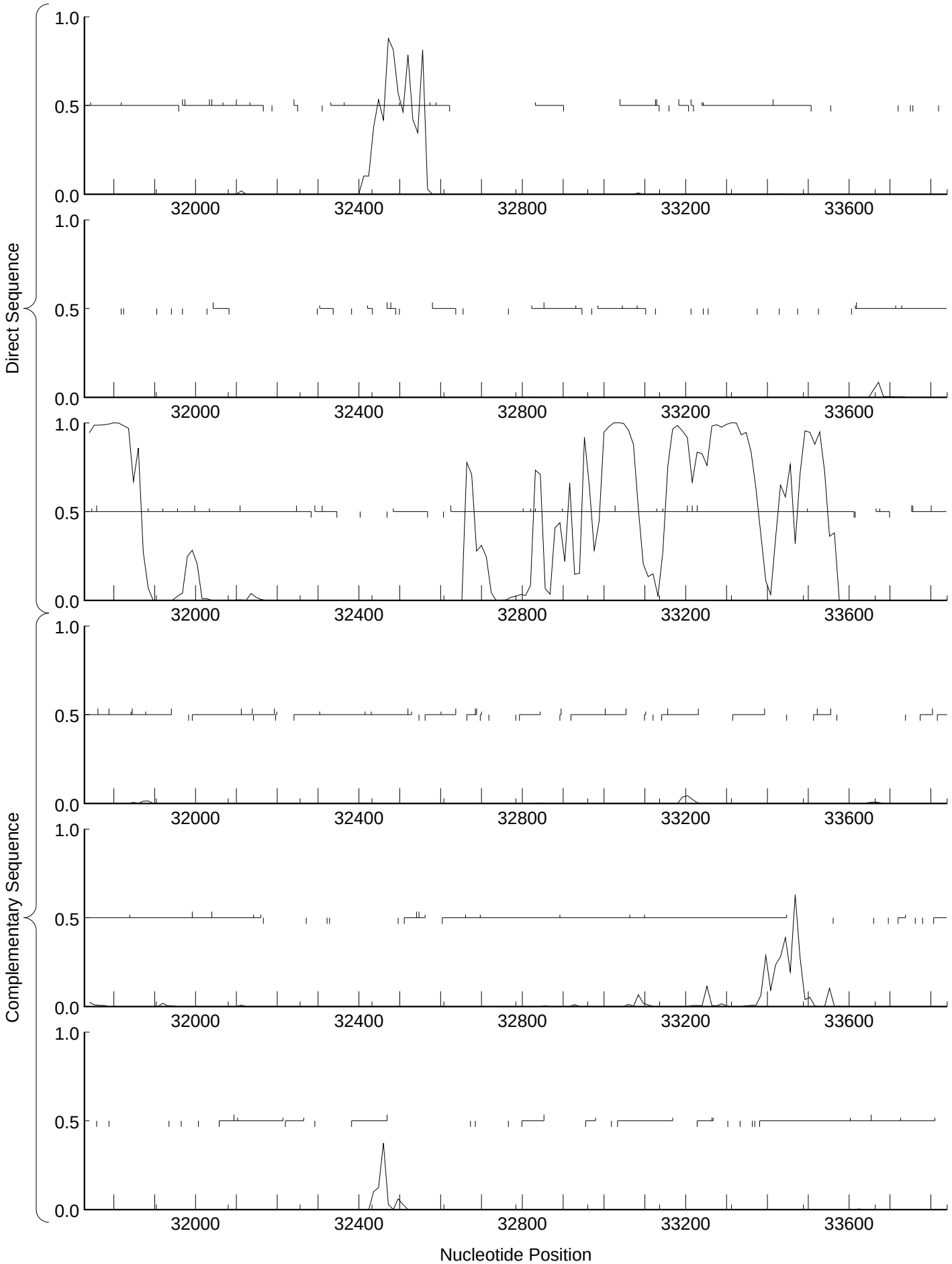


Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 15/37

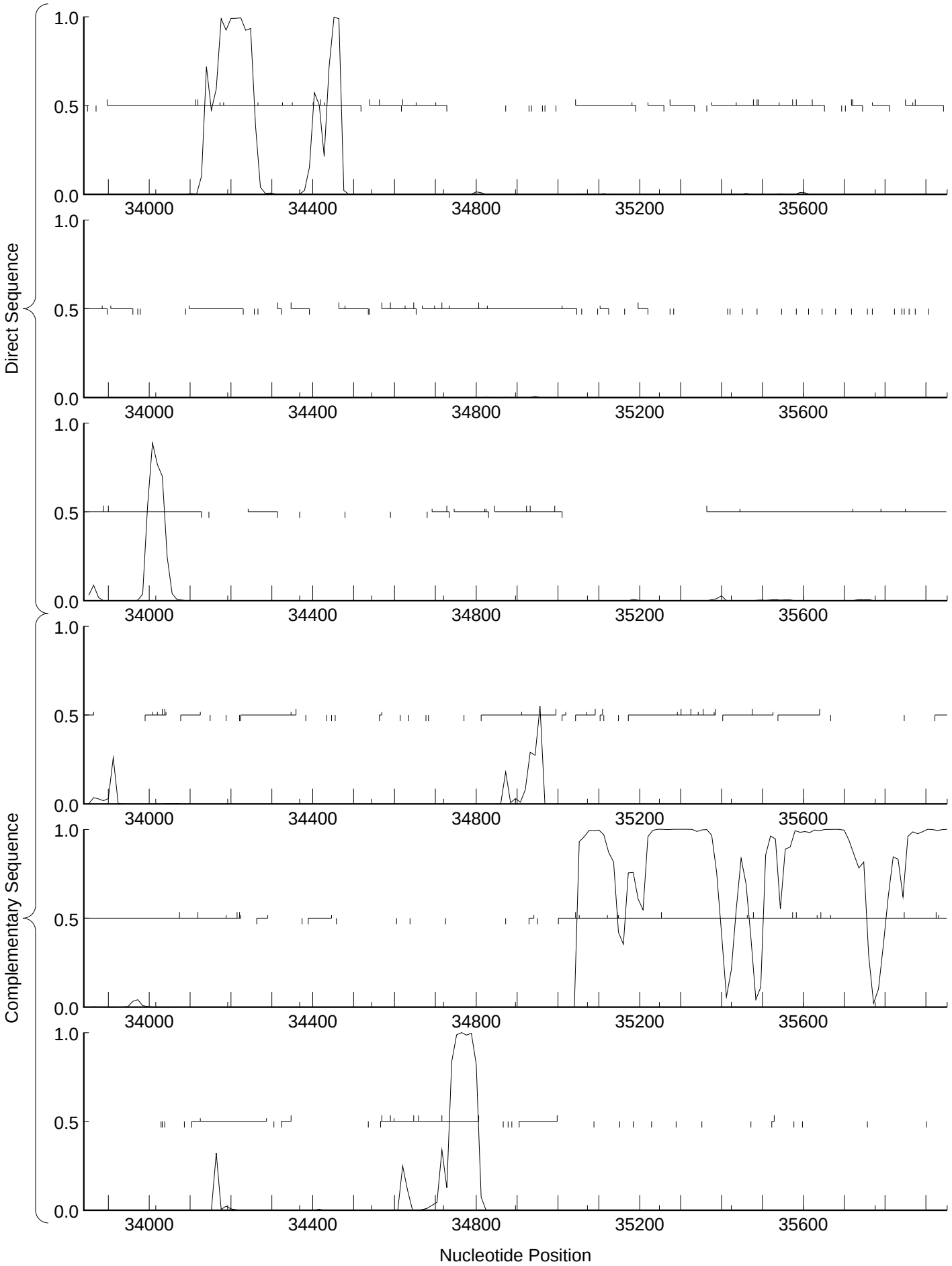



Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 16/37
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Complementary Sequence
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0.0
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30800
31200
31600
Nucleotide Position

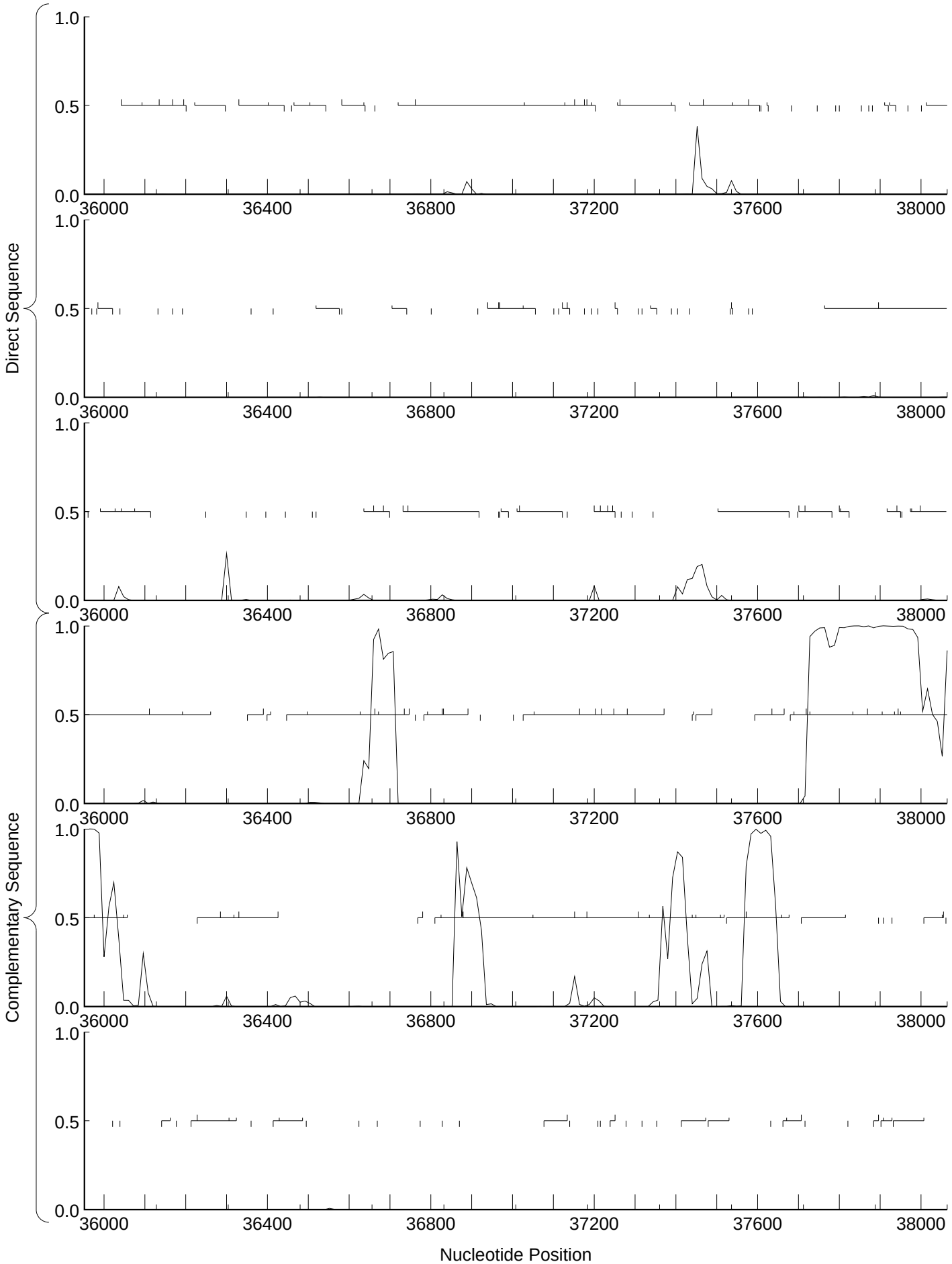
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 17/37



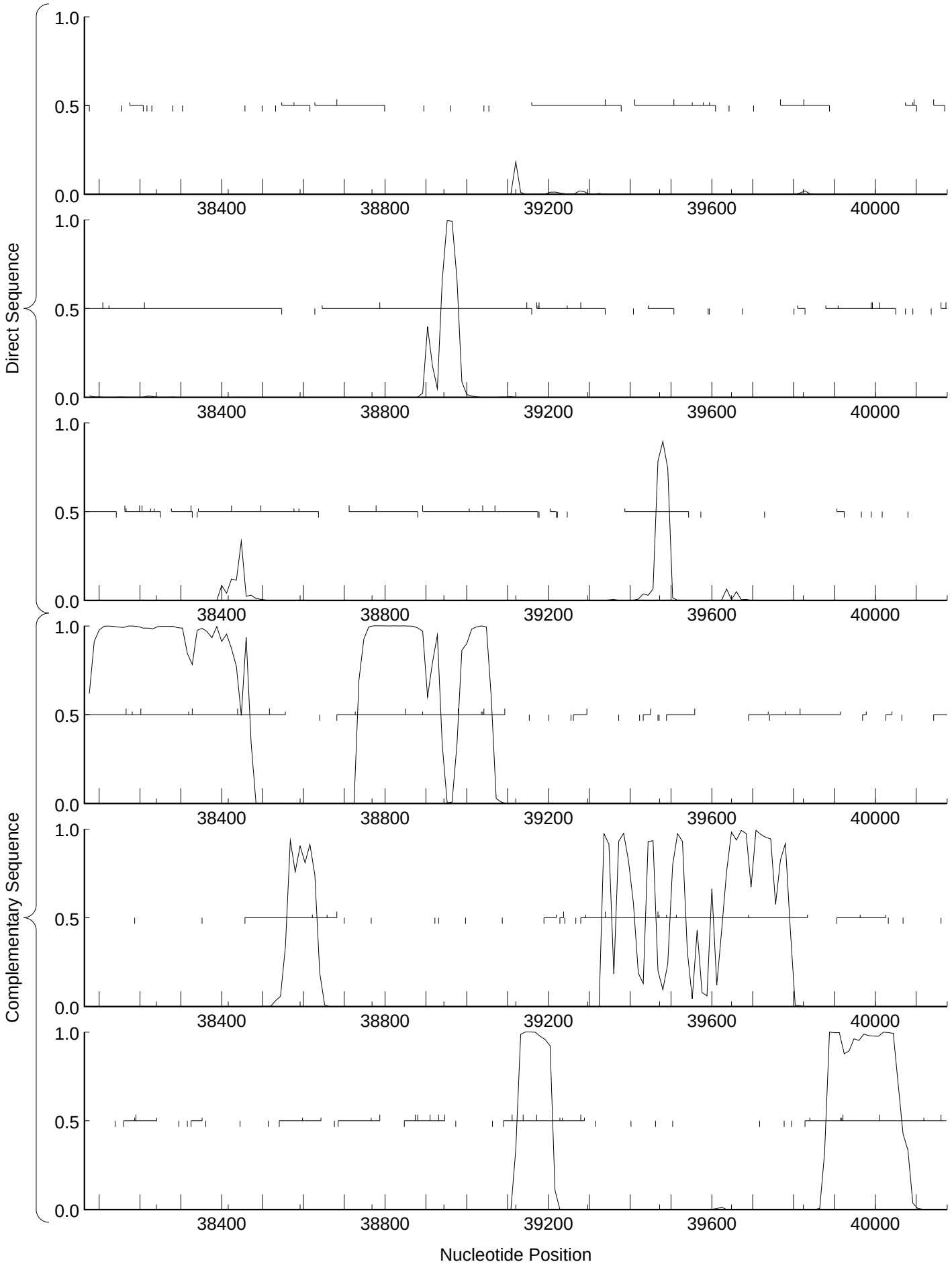
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 18/37



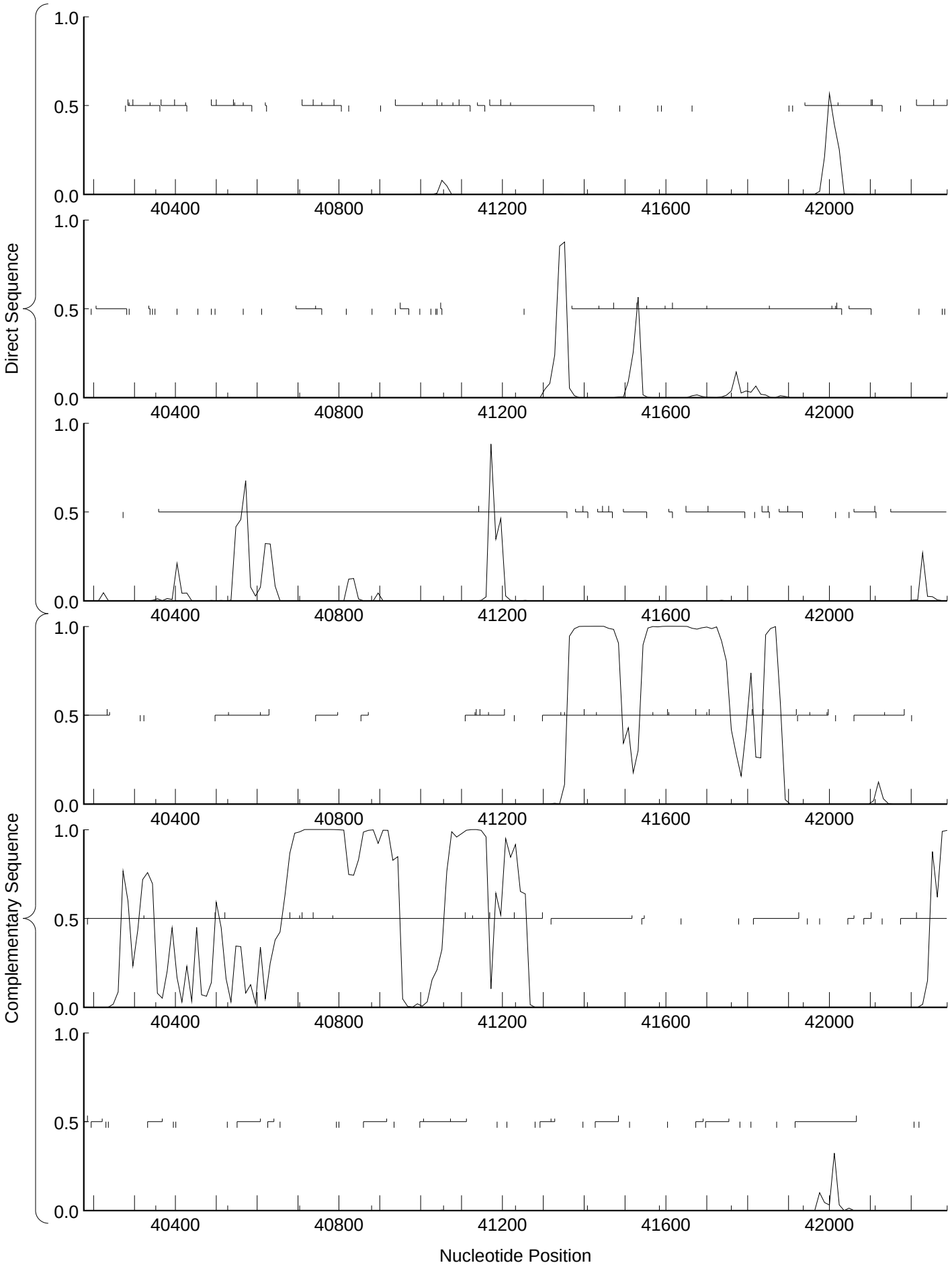
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 19/37



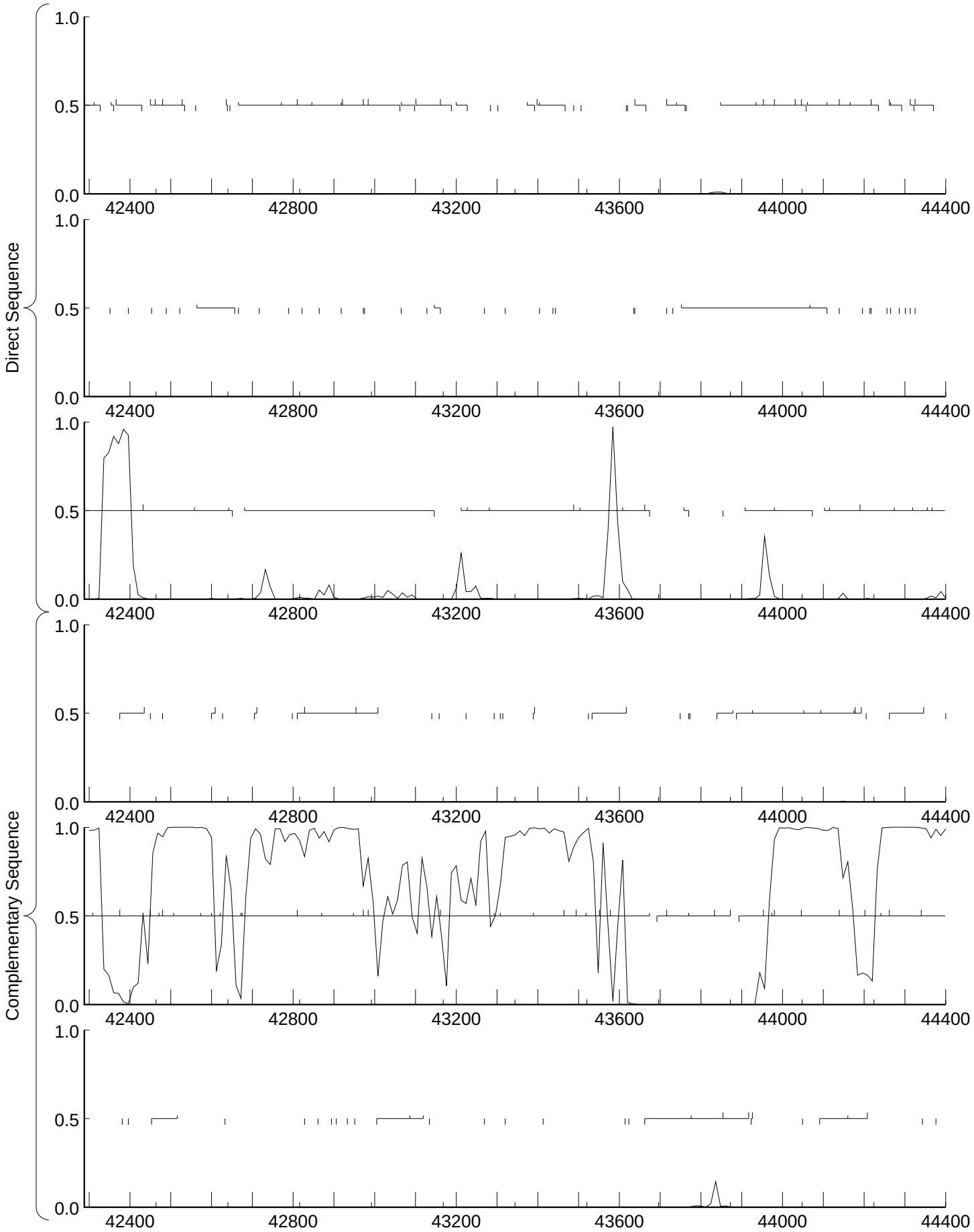
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 20/37



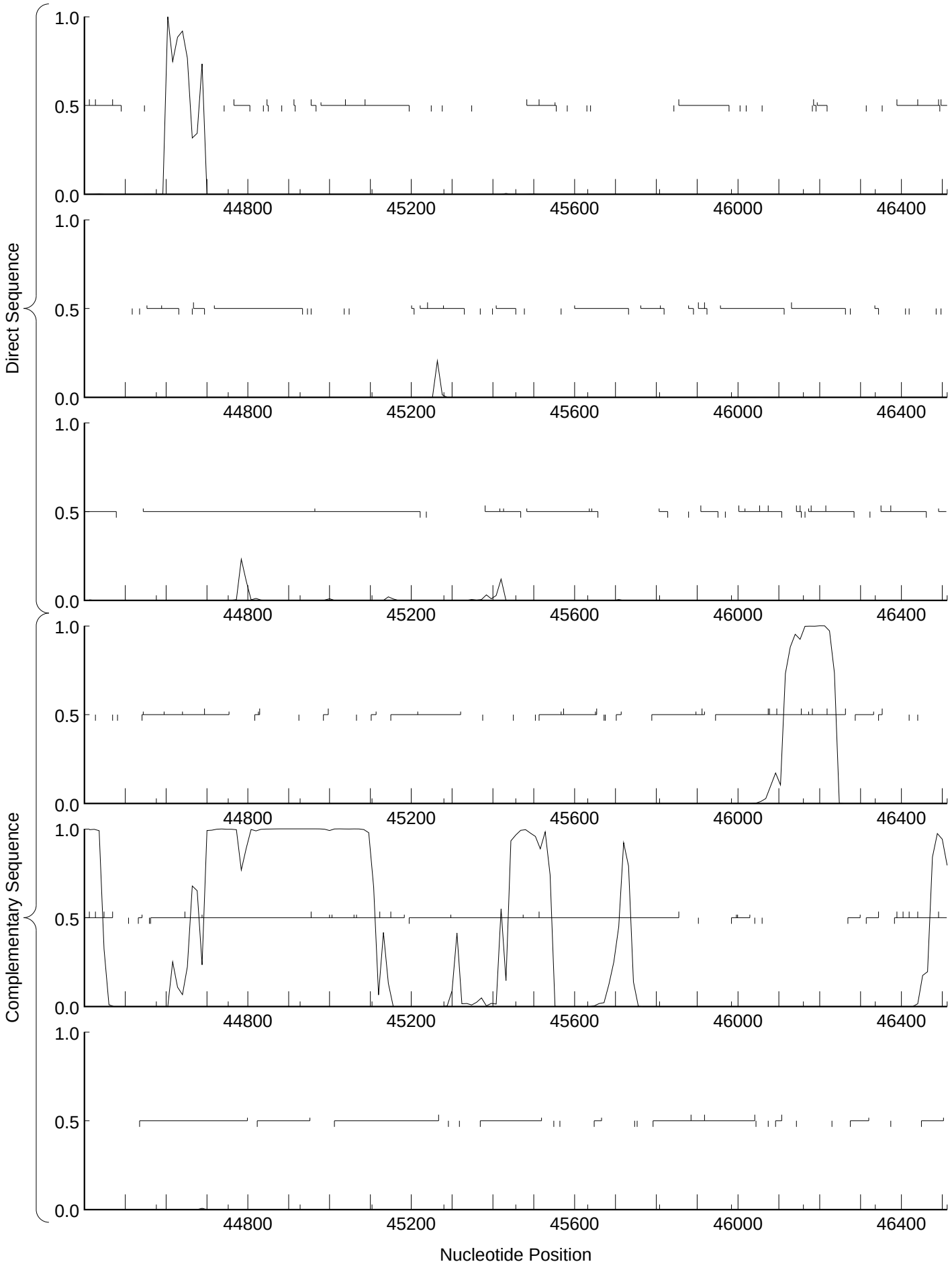
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 21/37



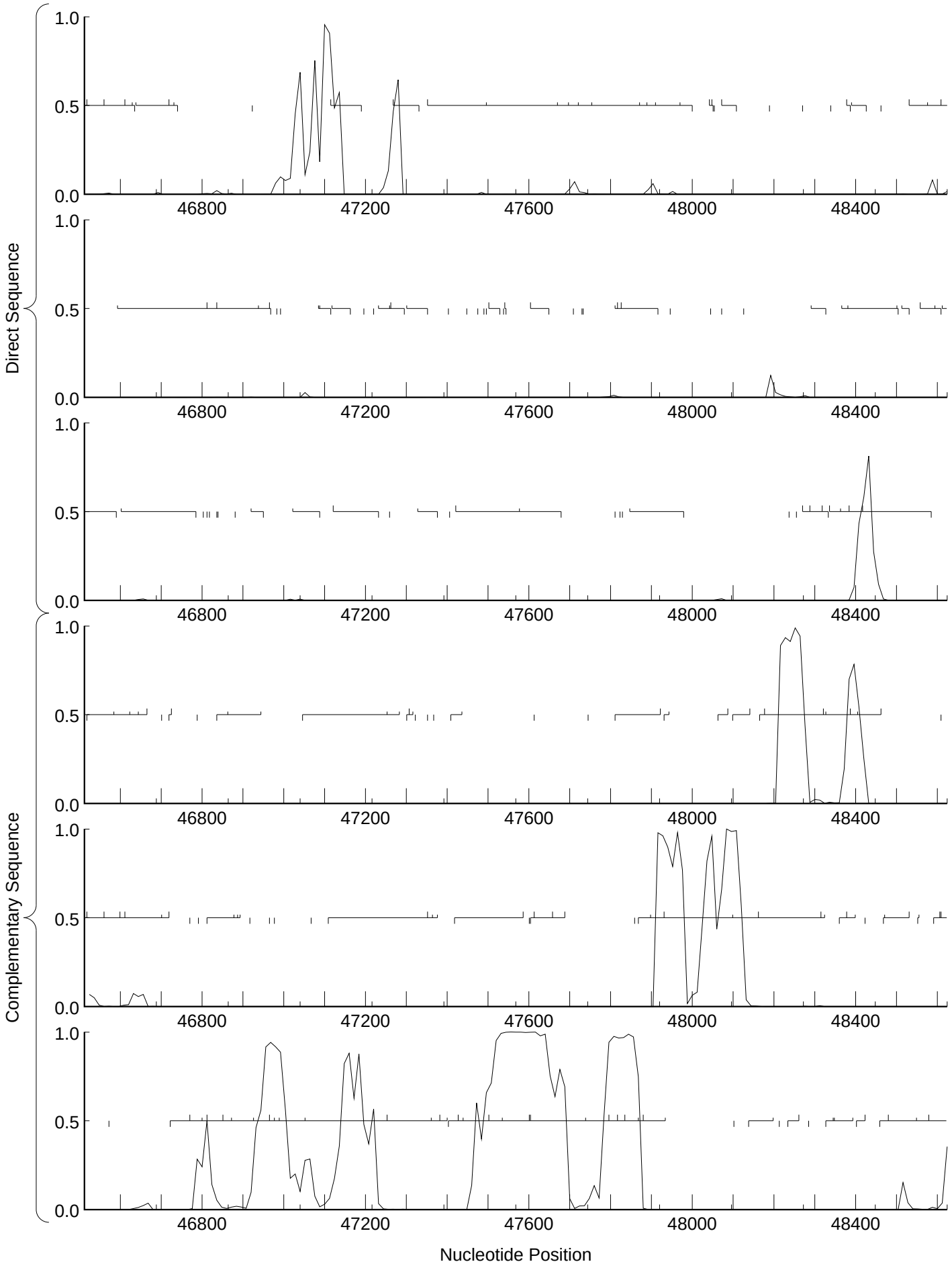
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 22/37



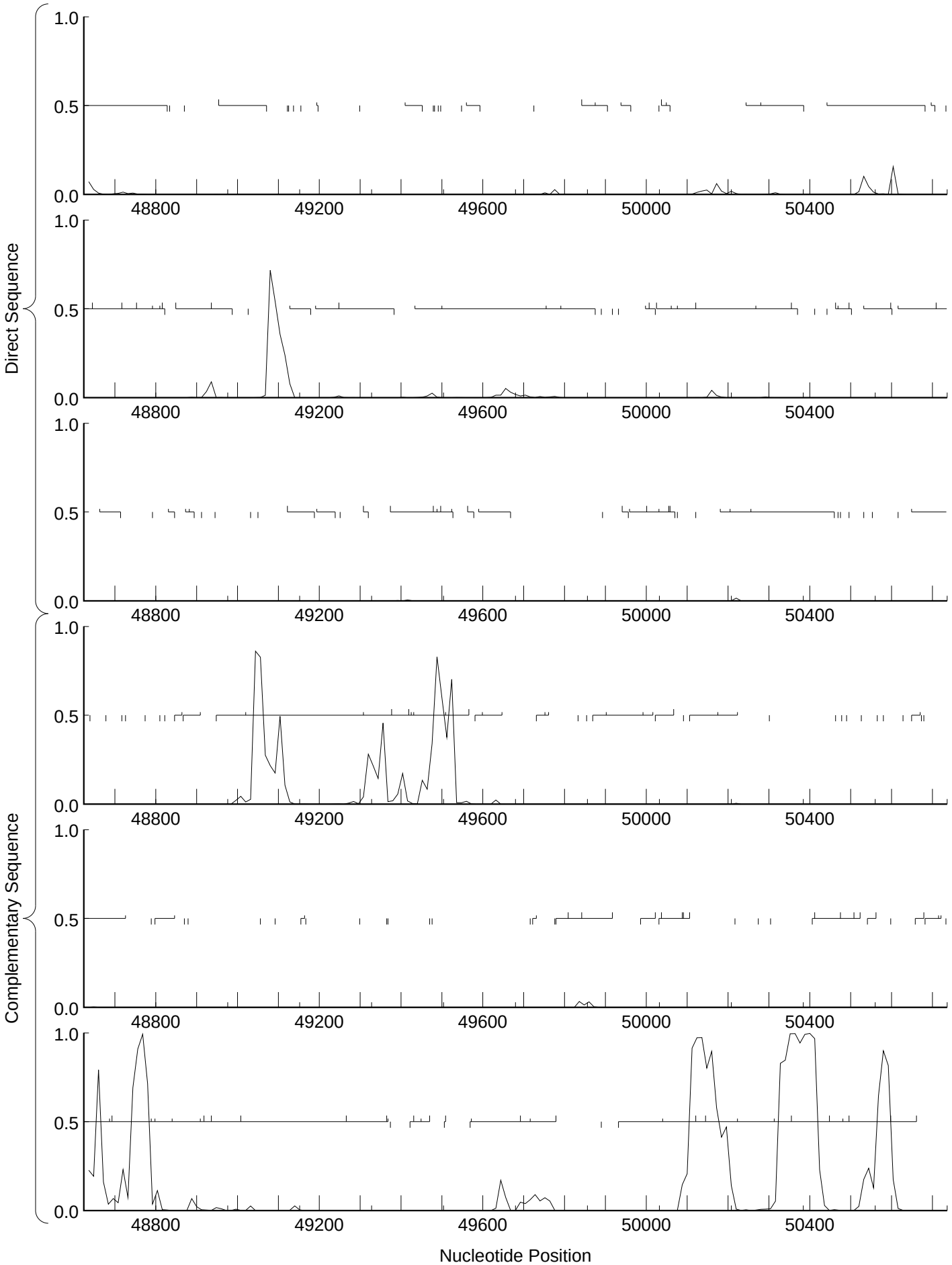
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 23/37



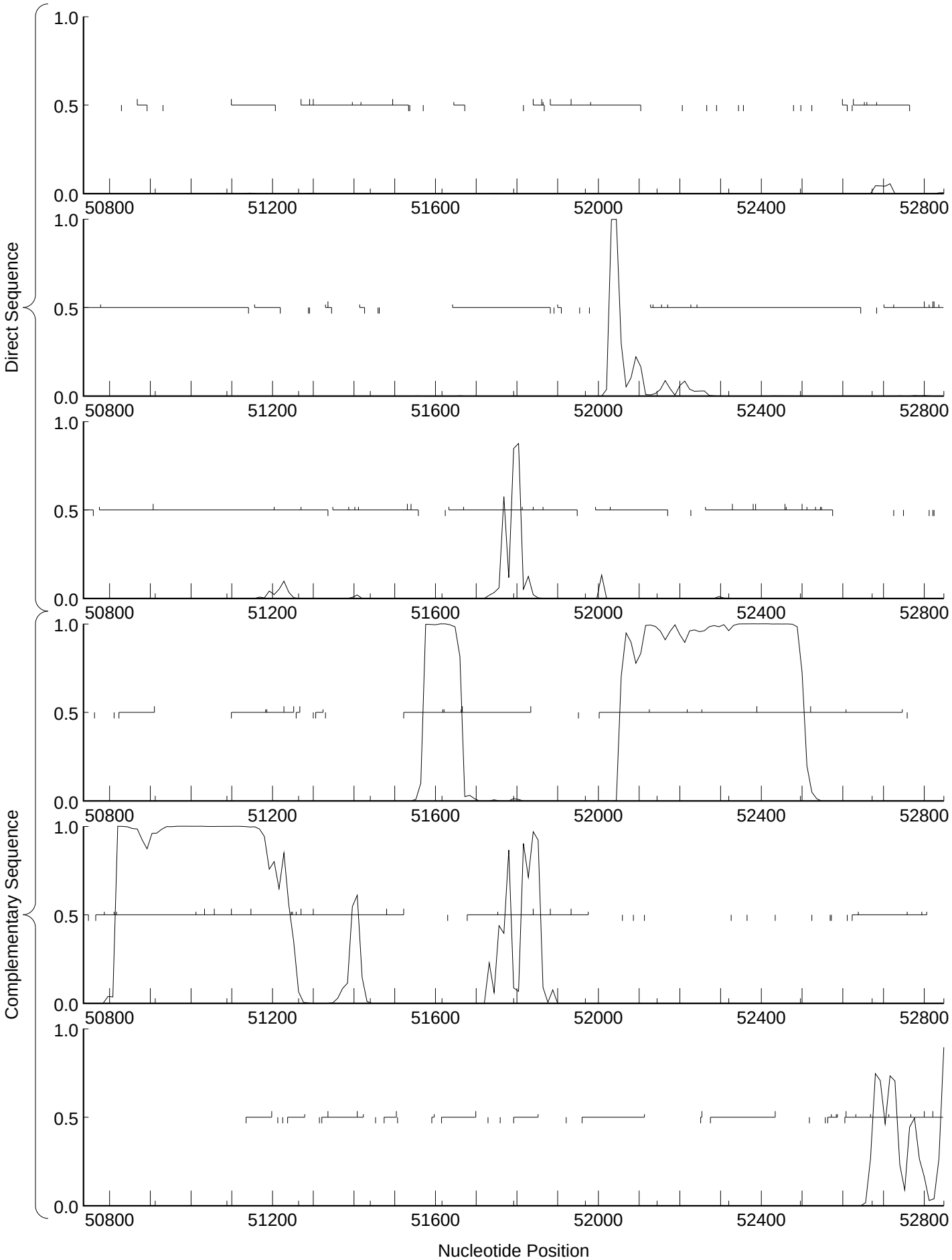
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 24/37



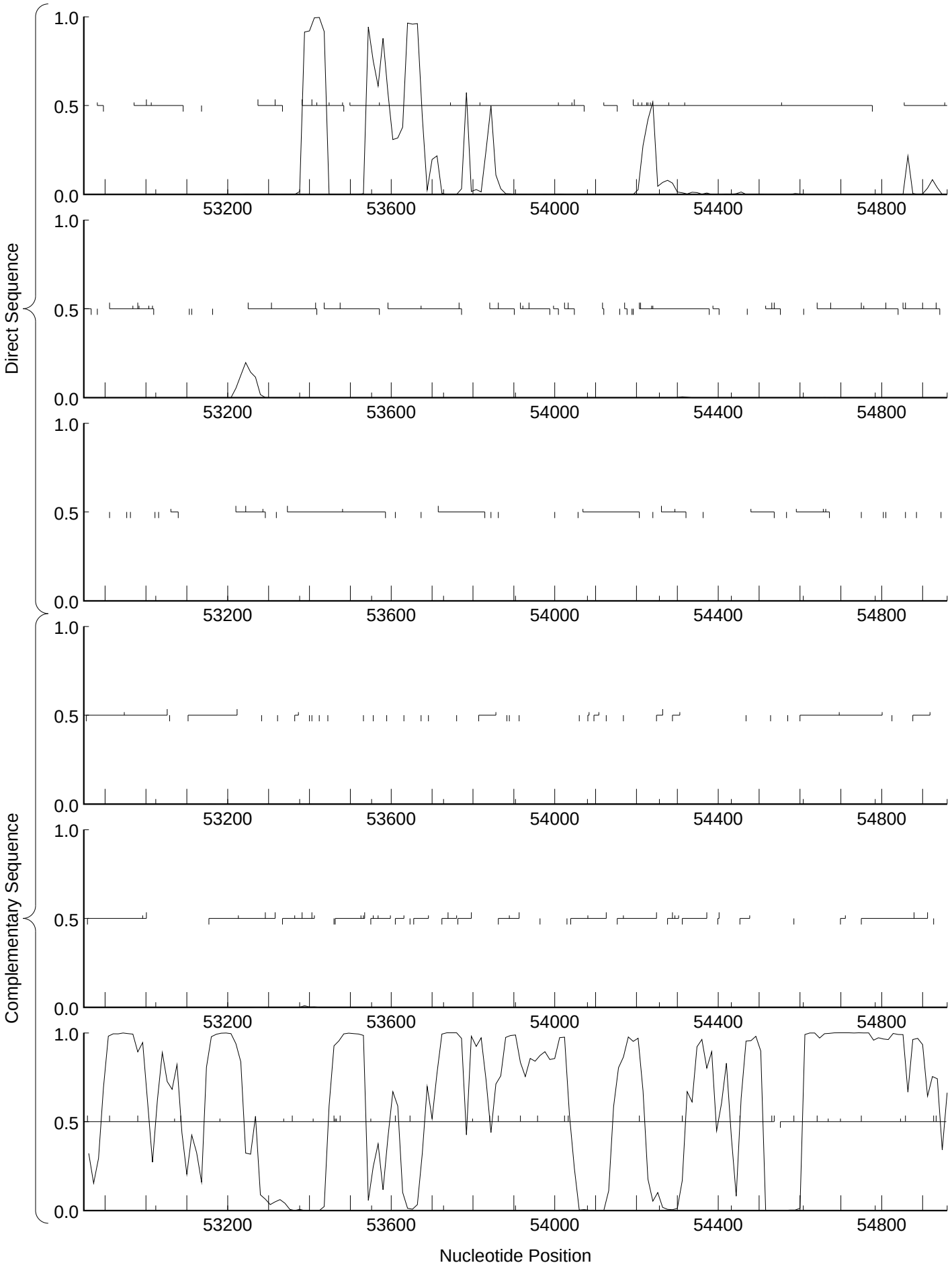
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 25/37



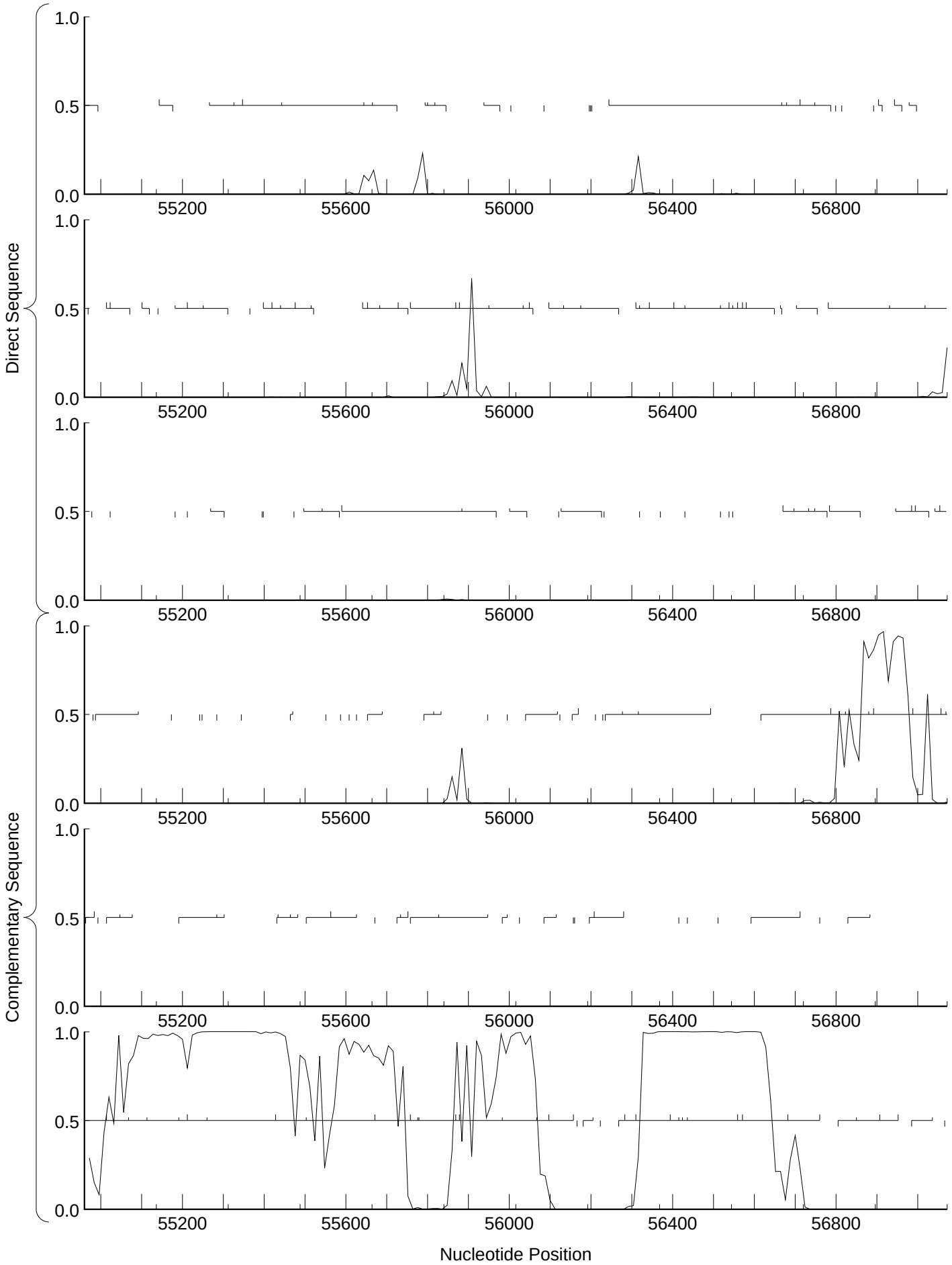
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 26/37



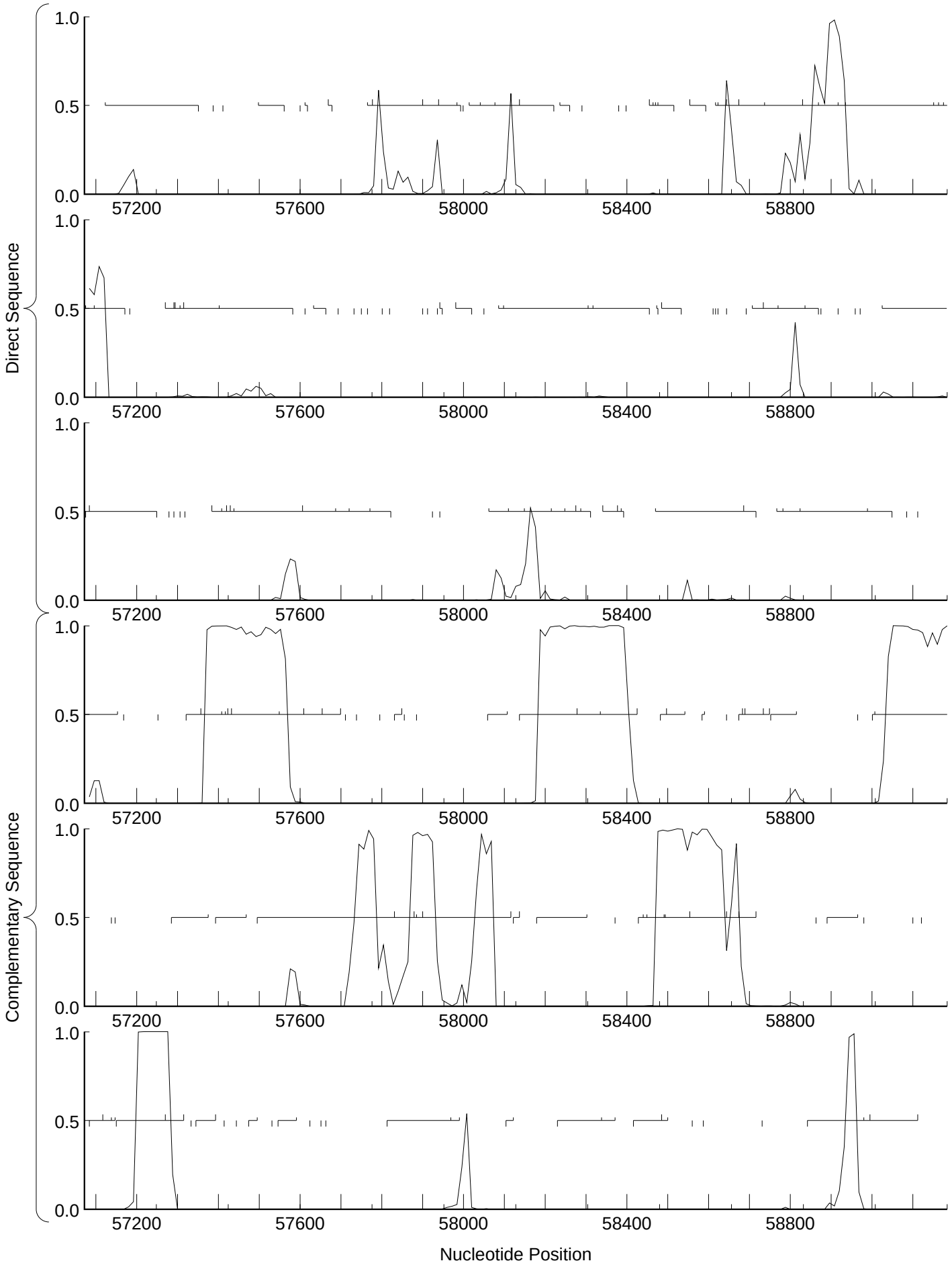
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 27/37

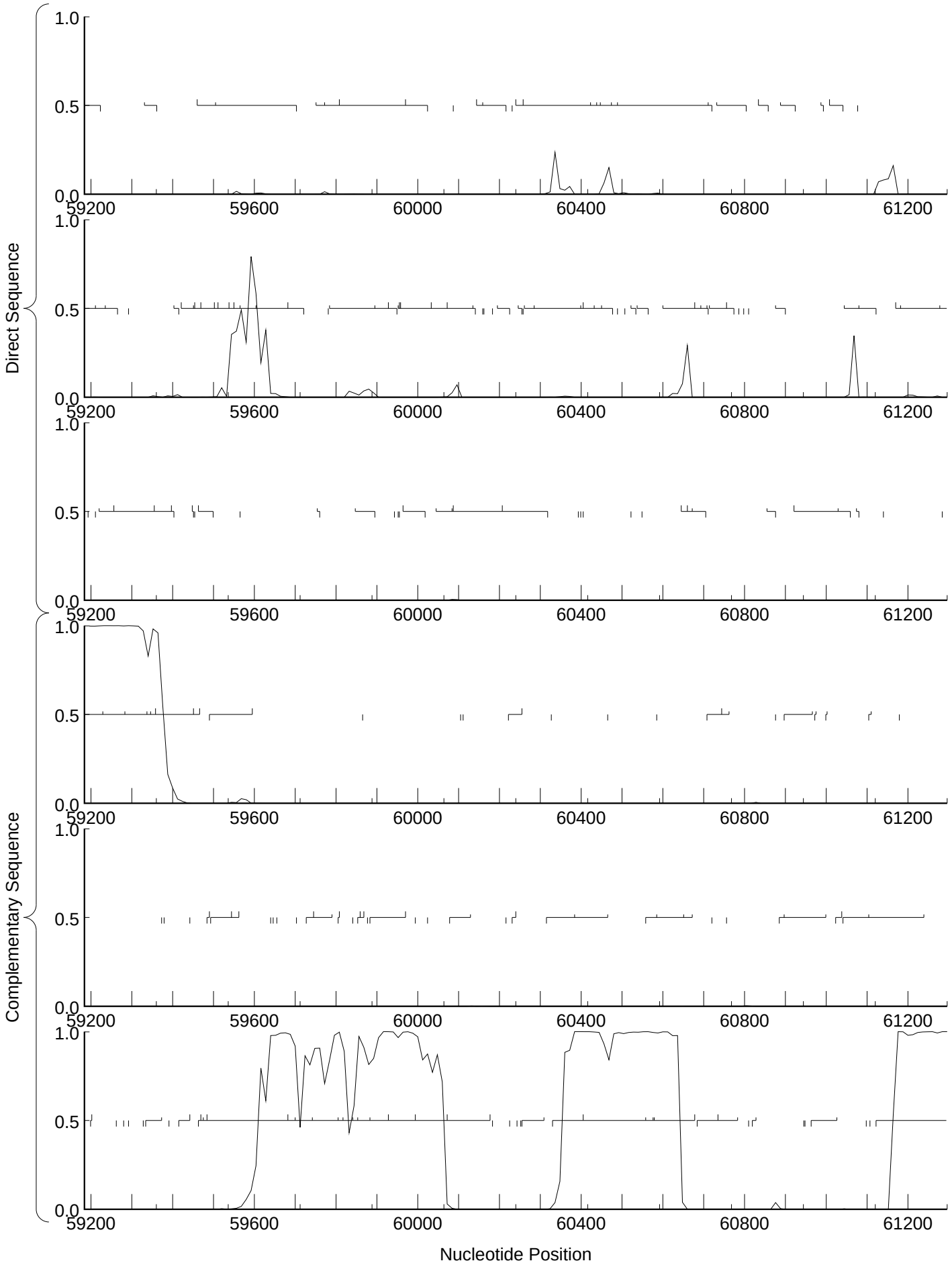


Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 28/37

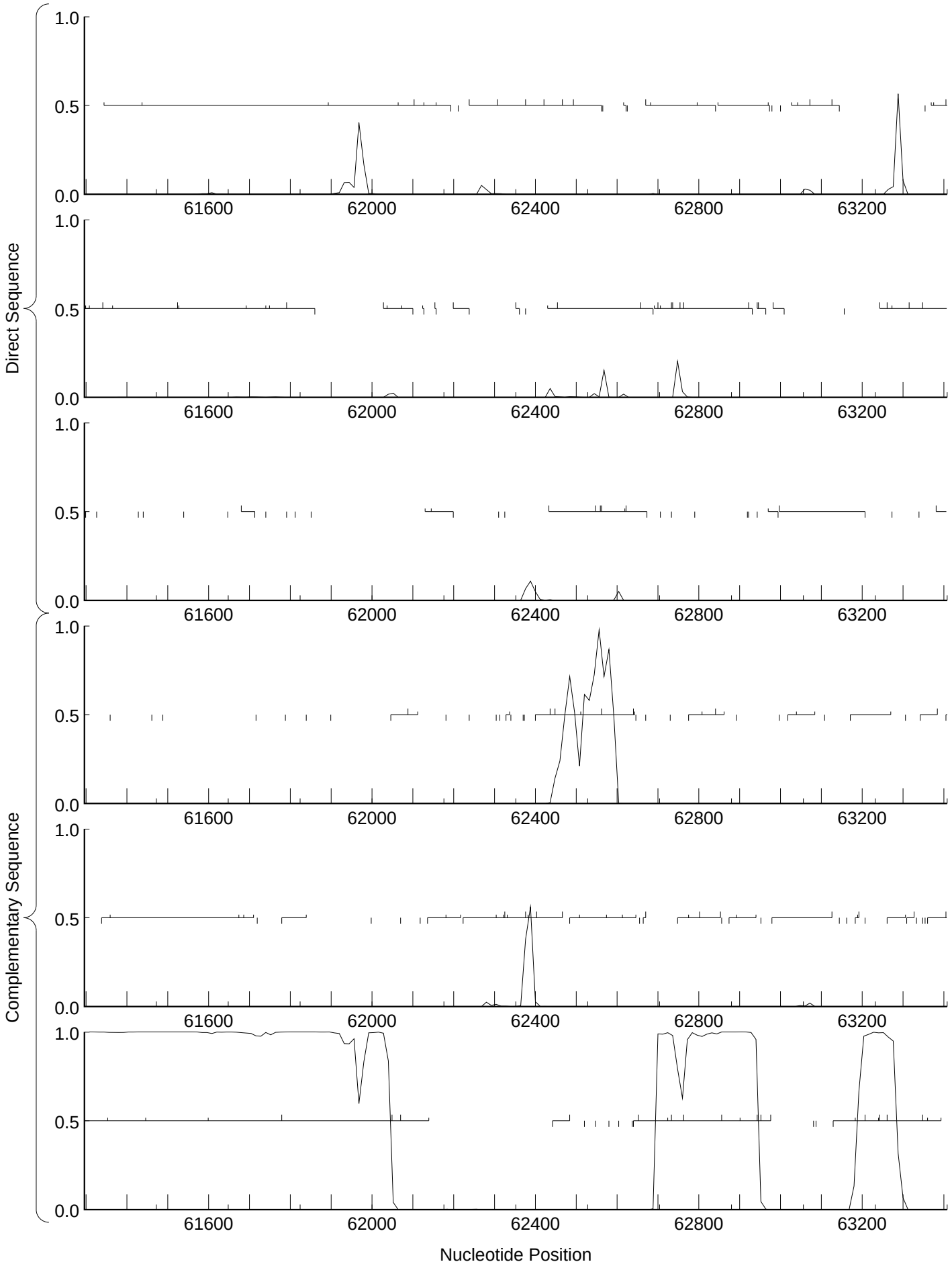


Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 29/37

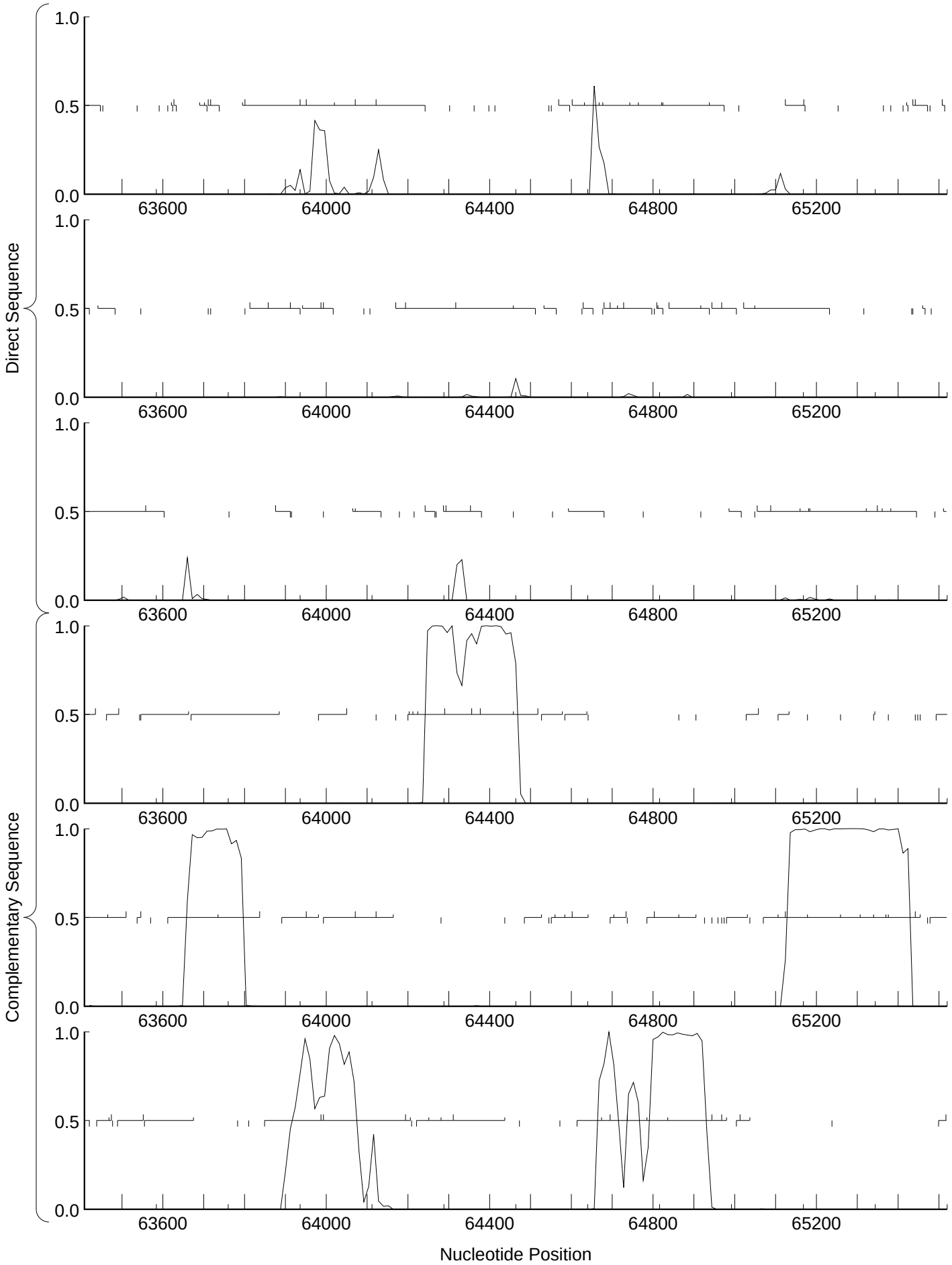



Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 30/37
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Complementary Sequence
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0.0
59200
59600
60000
60400
60800
61200
Nucleotide Position

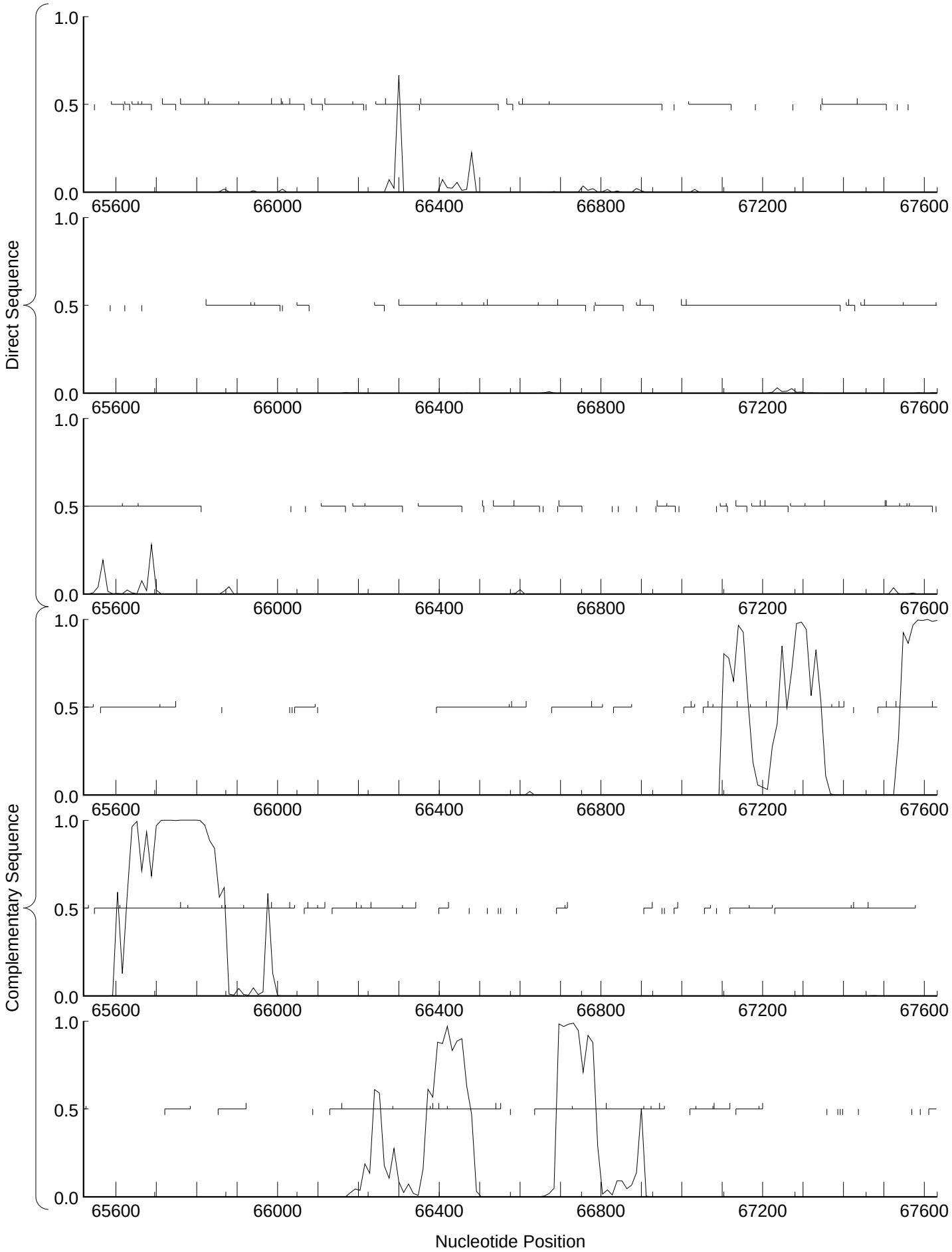
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 31/37



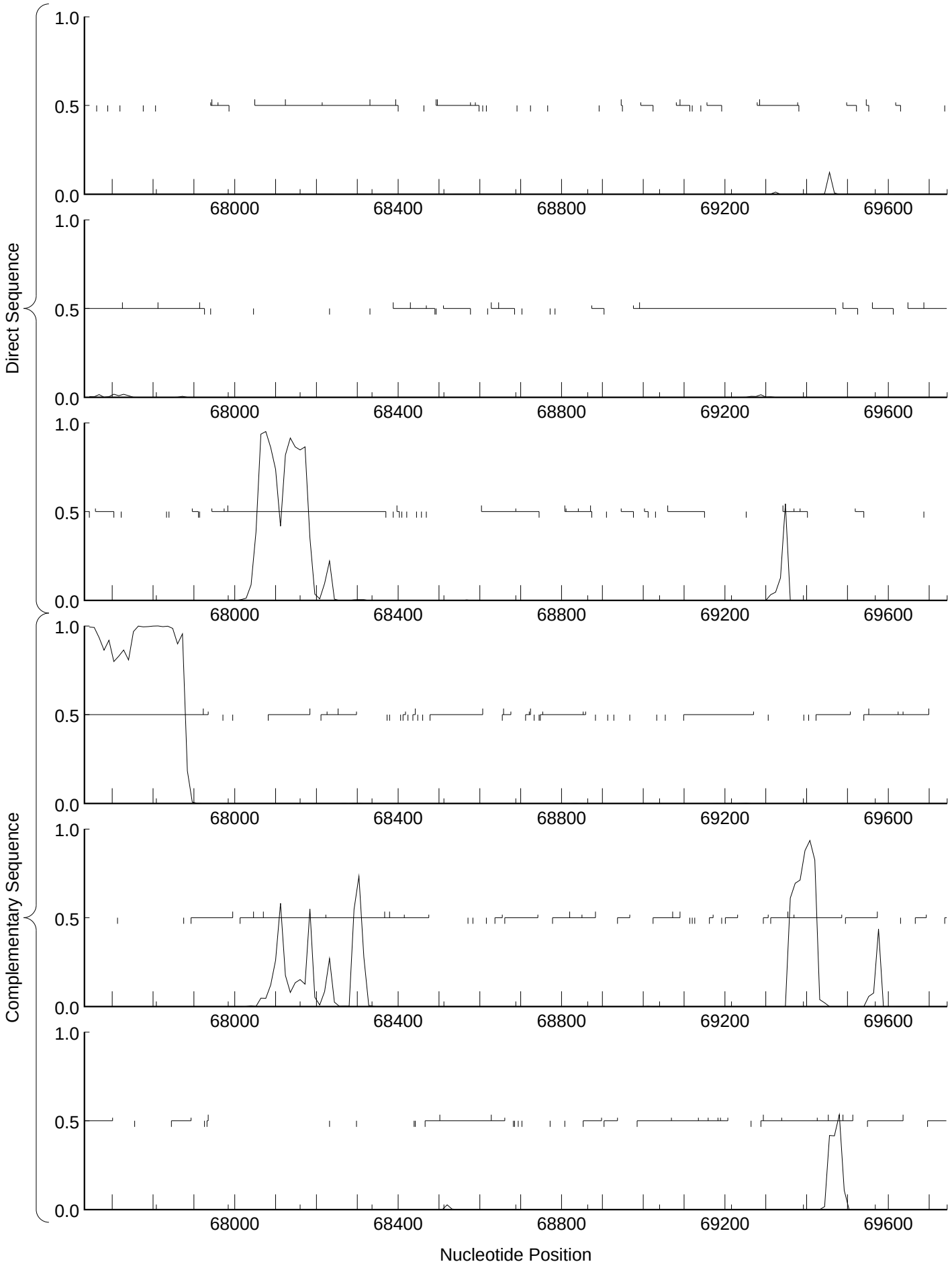
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 32/37



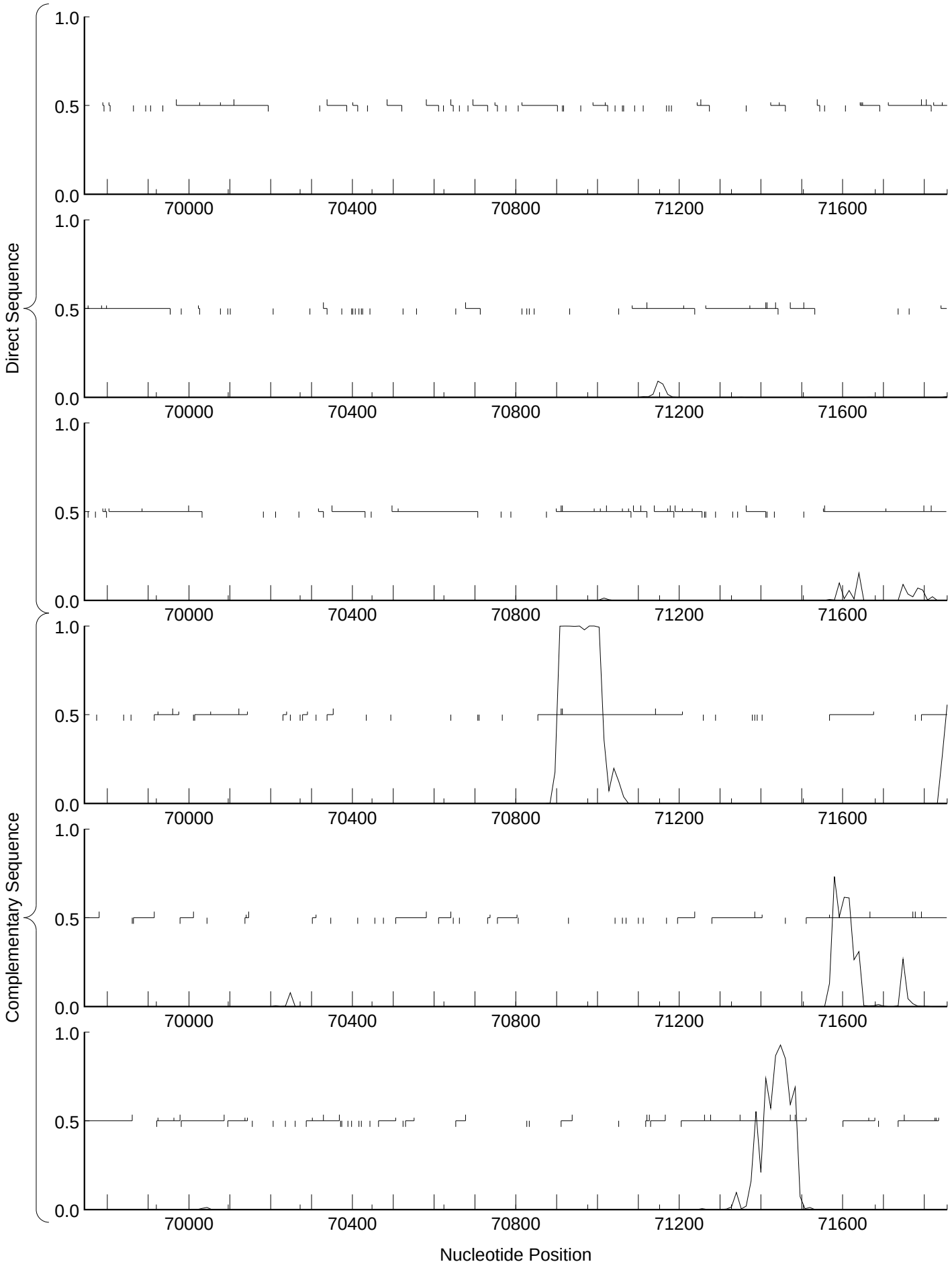
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 33/37



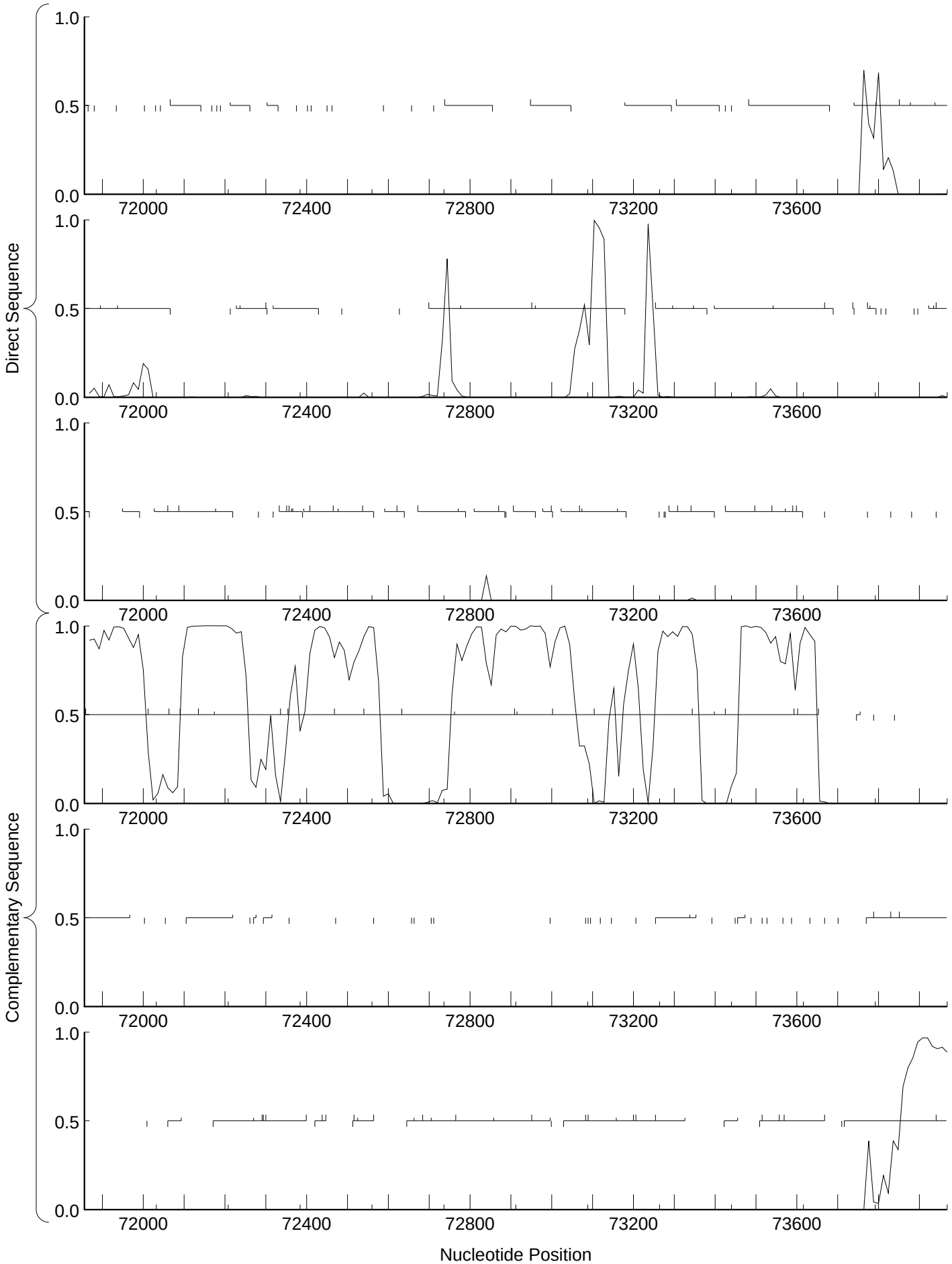
Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 34/37



Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 35/37



Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 36/37



Rhodococcus phage DocB7 complete genome, 75772 bp, circularly permuted, Order 4, Window 96, Step 12, 37/37

