GeneMark

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

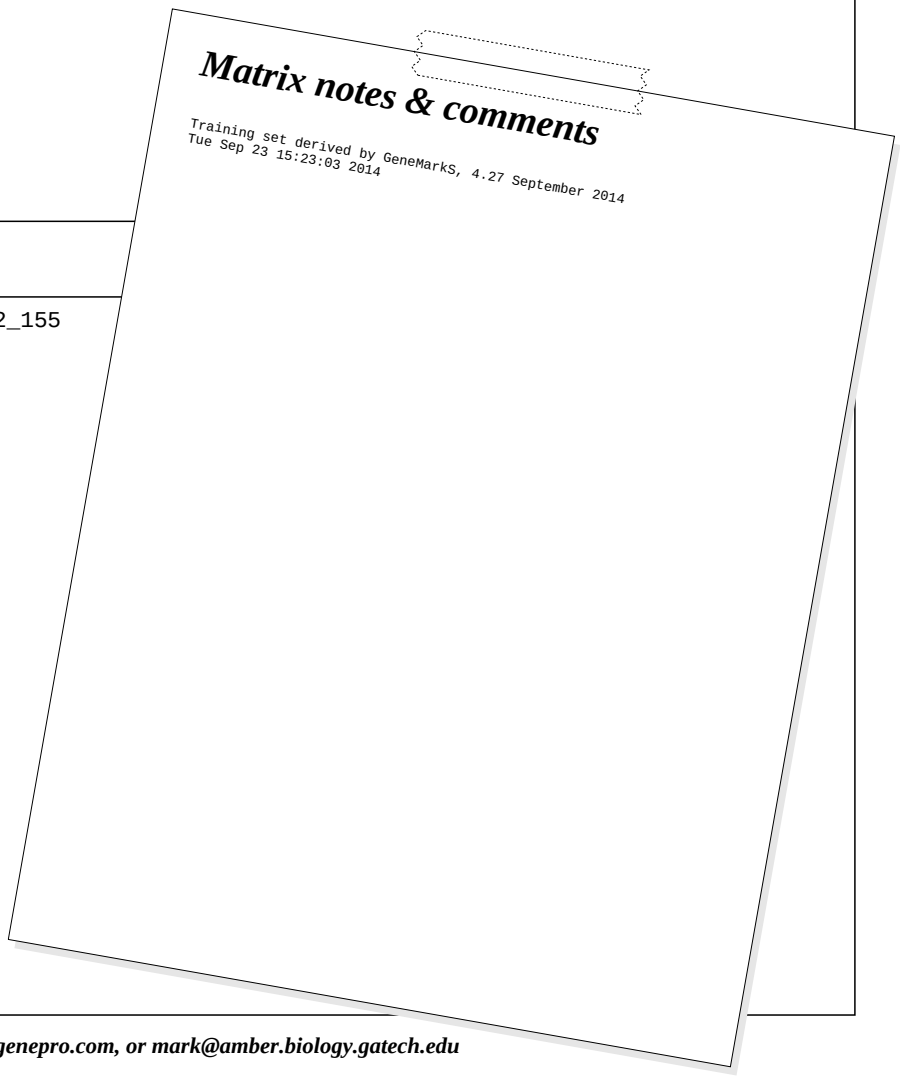
PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1
Analysis Date    :  3/29/24 at 2:53:26
Pages            :  34
Sequence Length  :  68791 bp
GC Content       :   66.49%

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

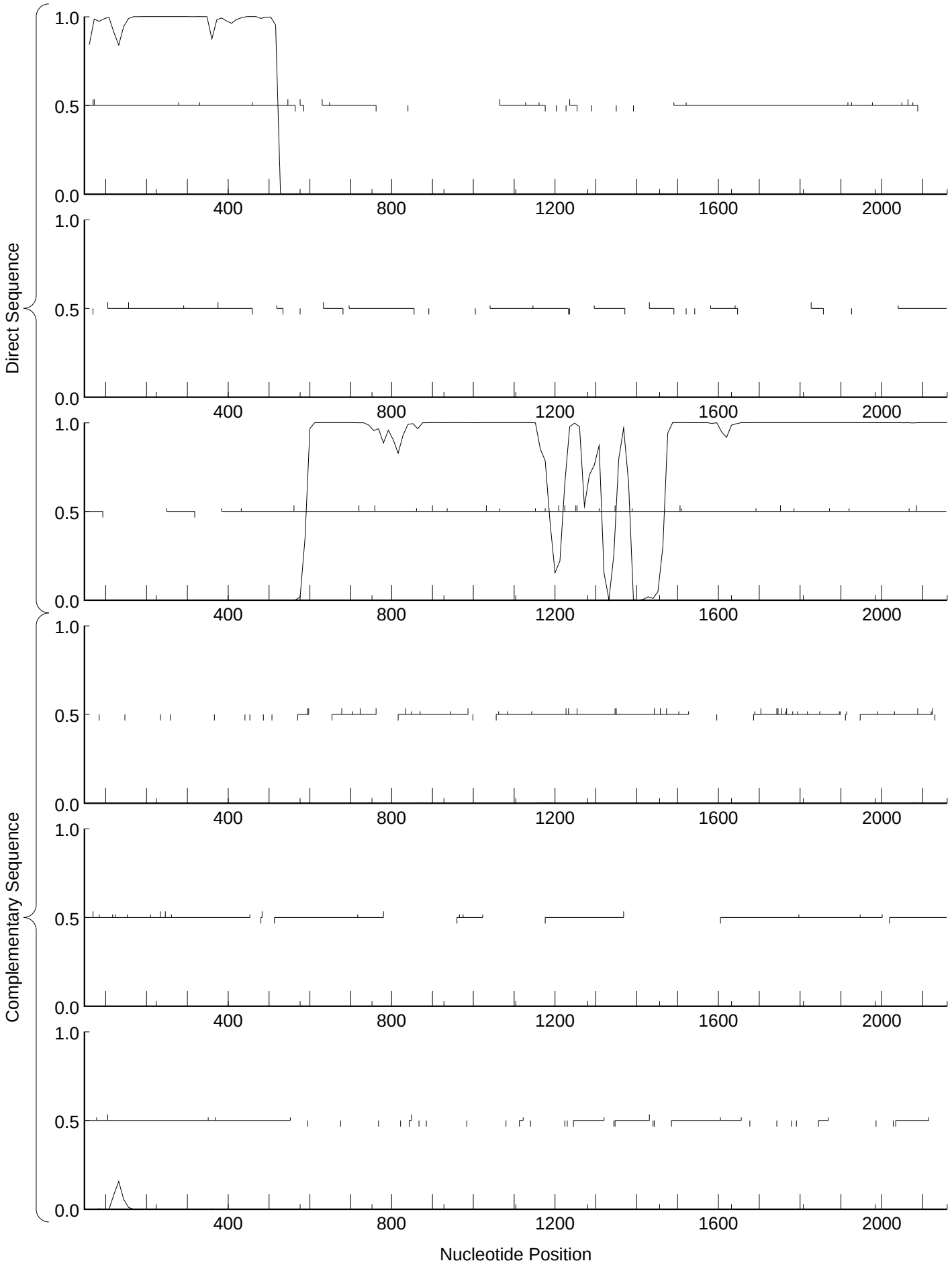
```
Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

Order      :  4
```

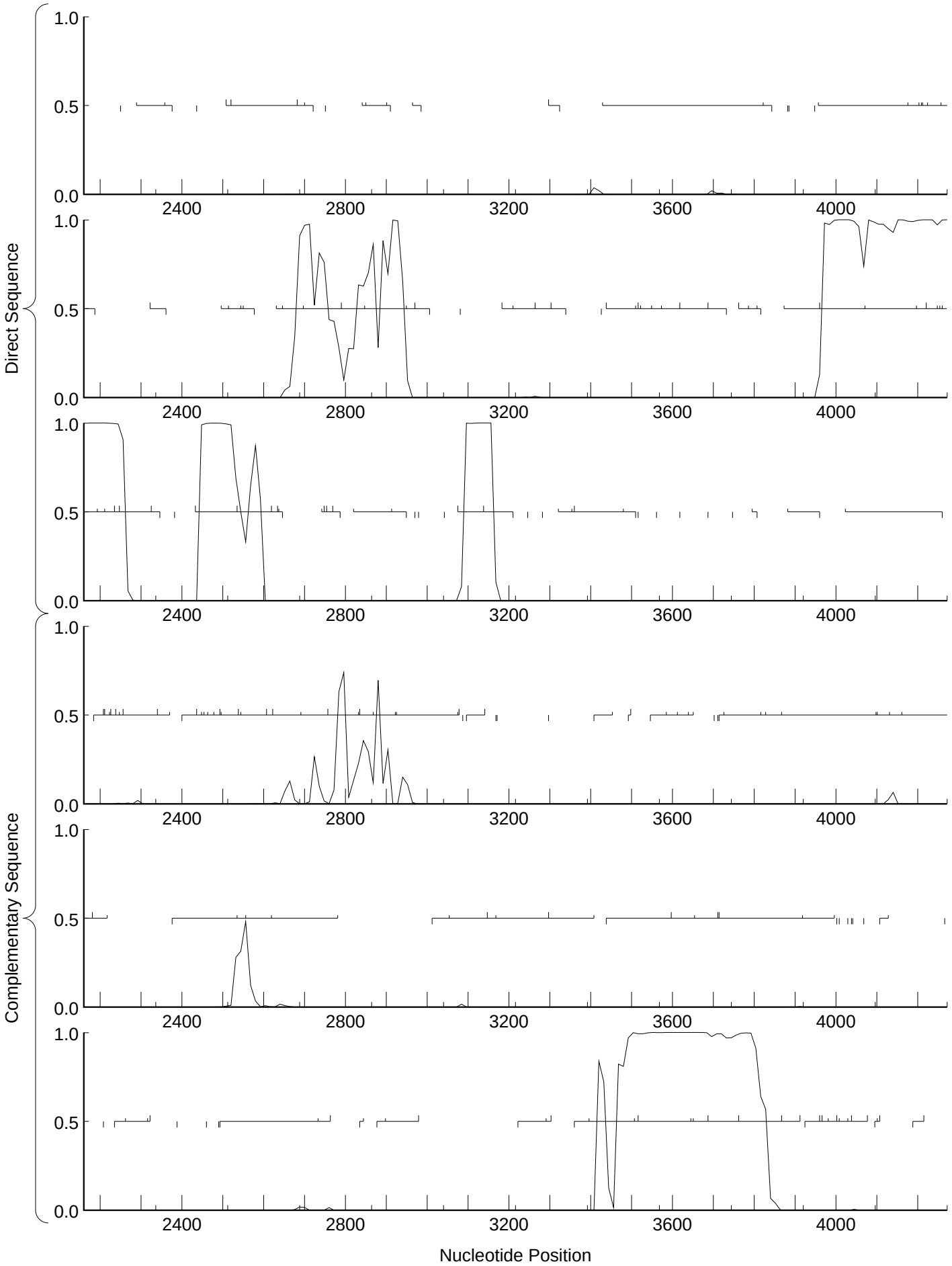

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

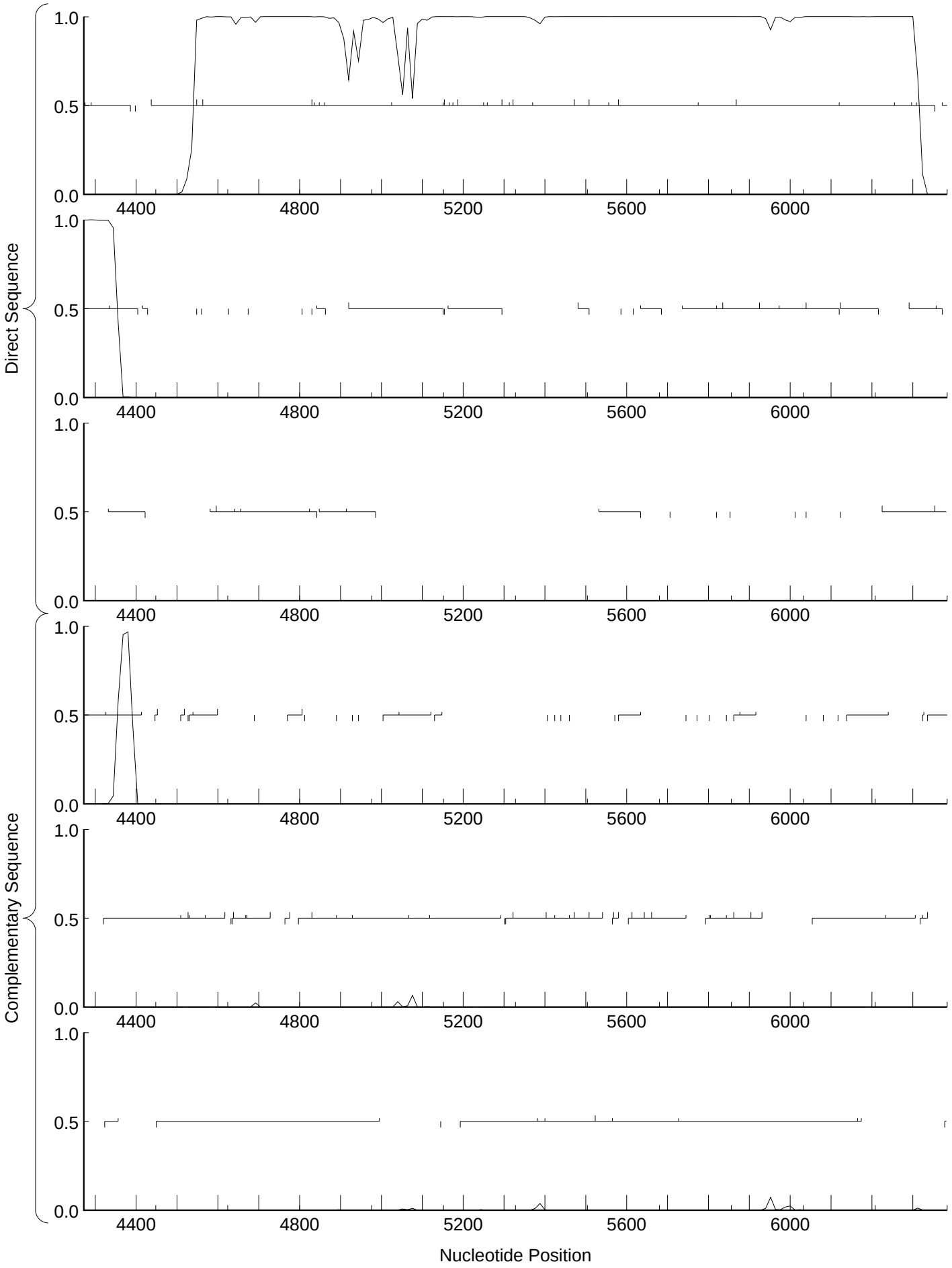
Mycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 2/34



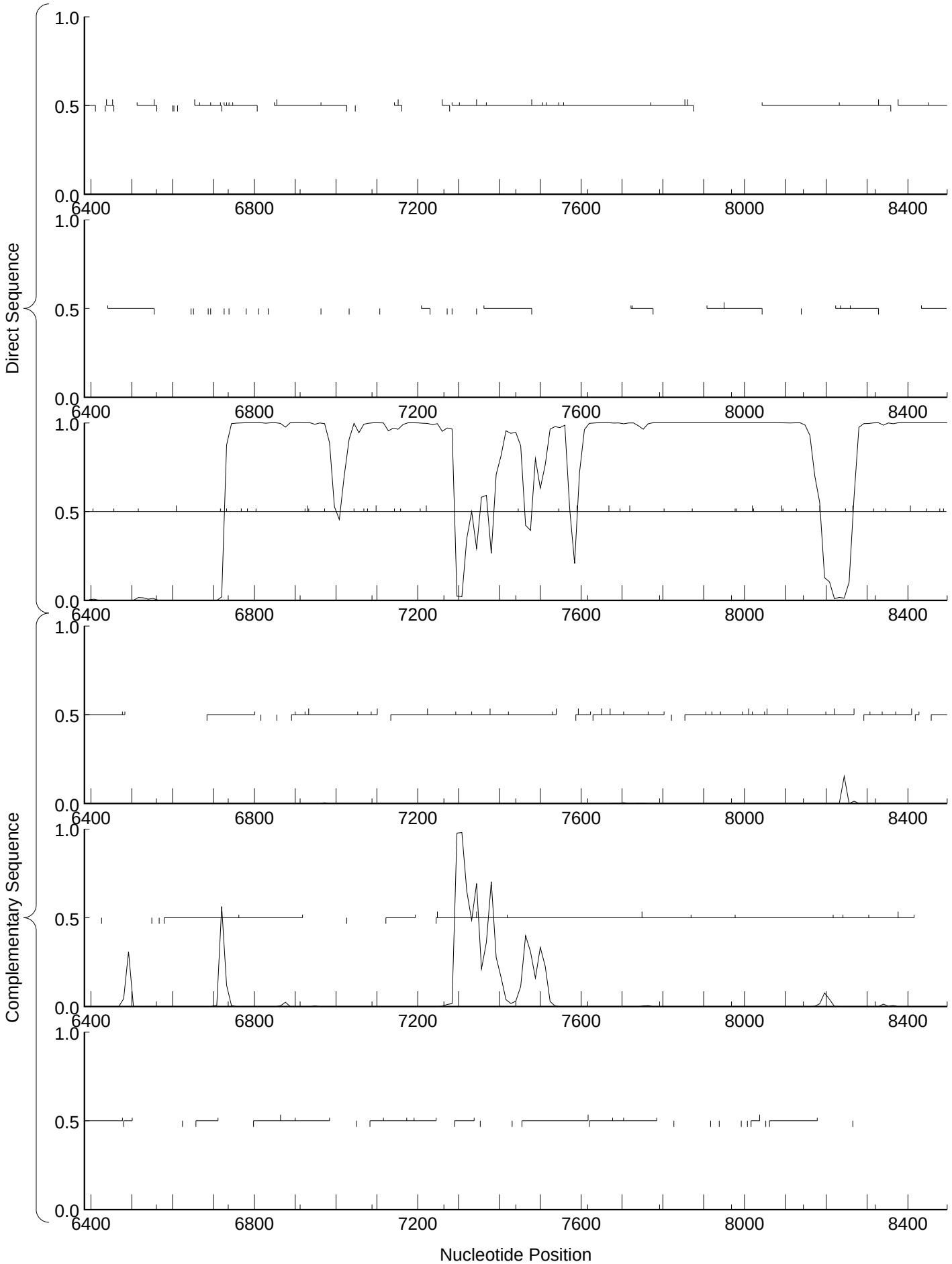
Mycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 3/34



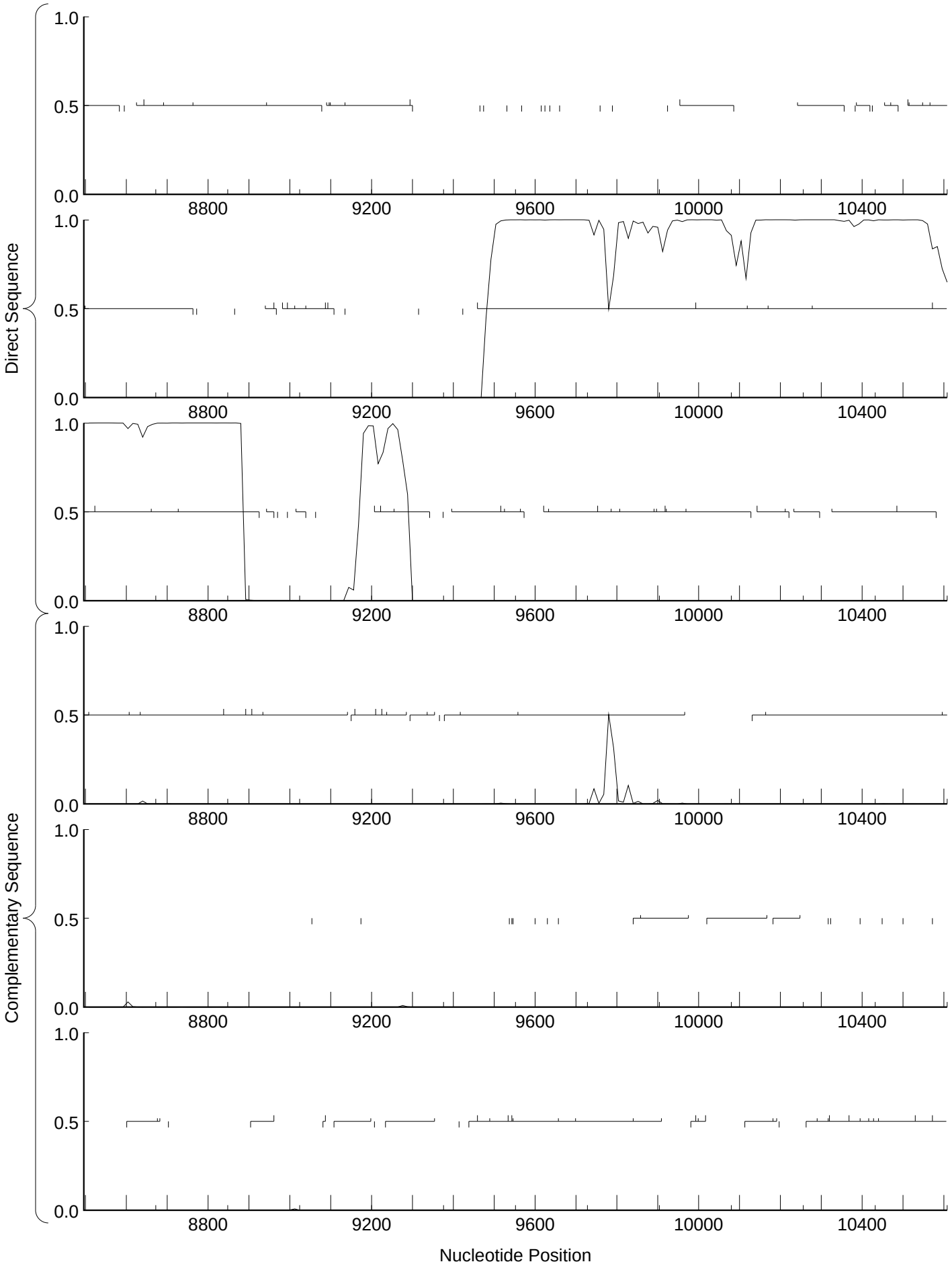
Mycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 4/34



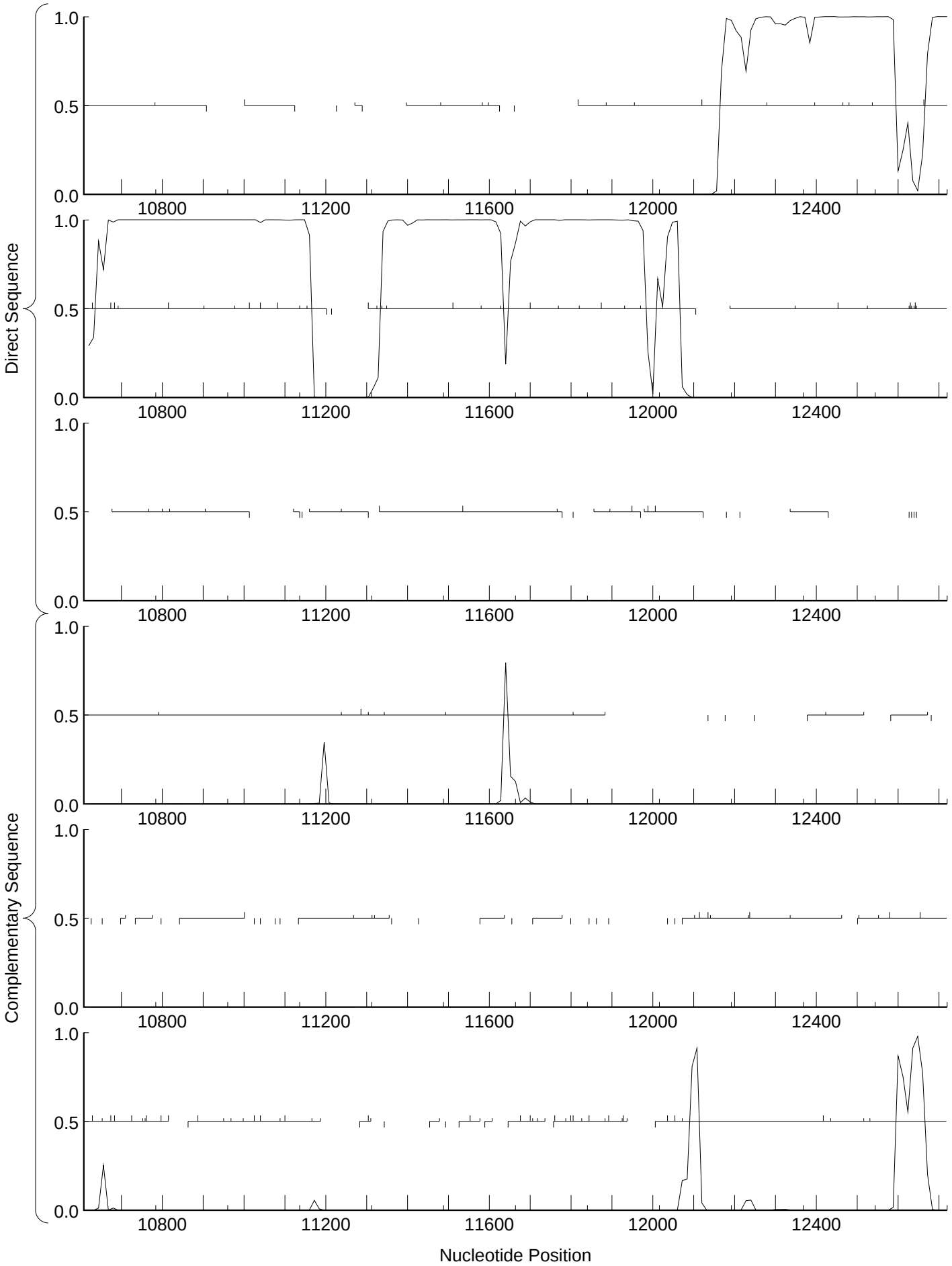
Mycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 5/34



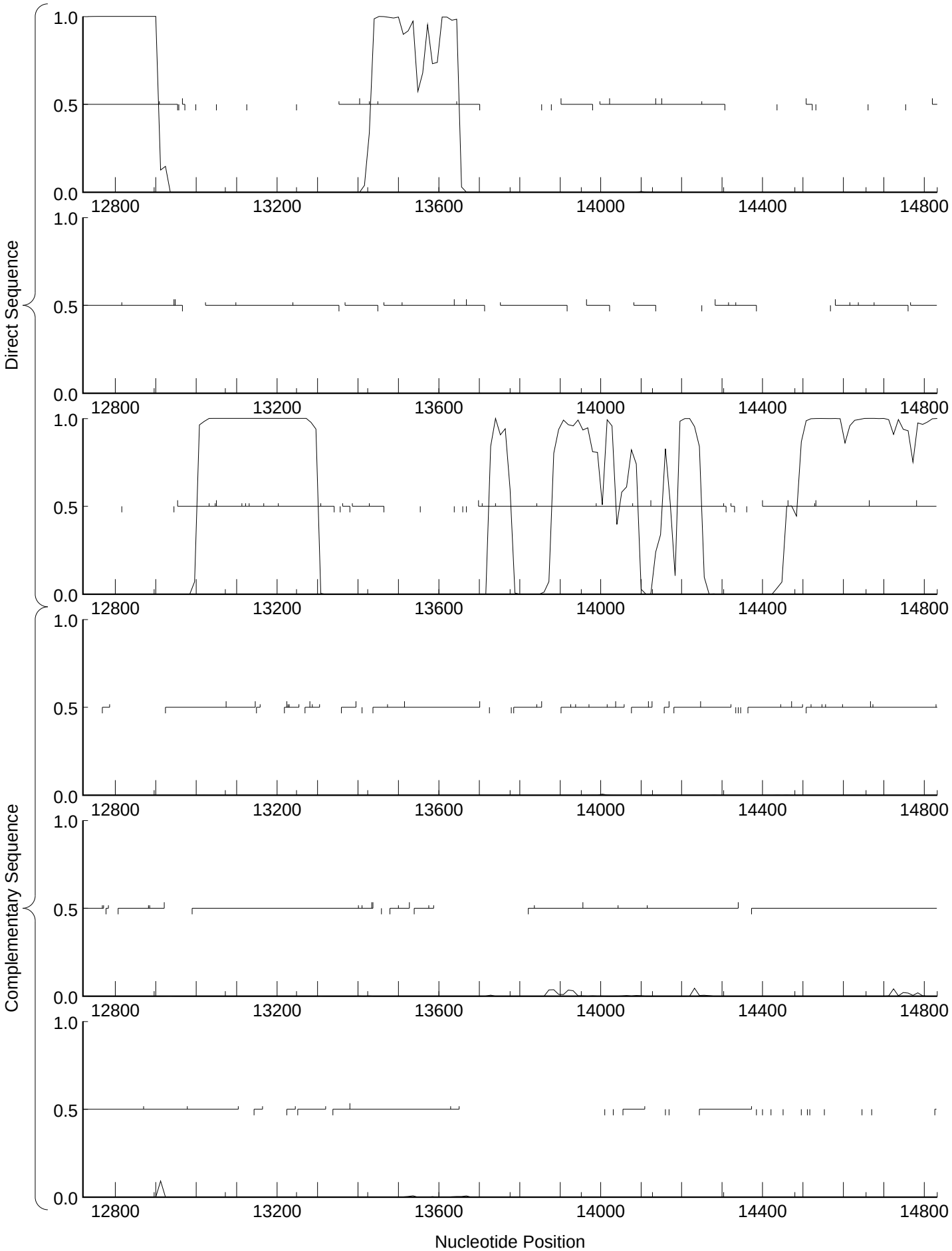
Mycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 6/34



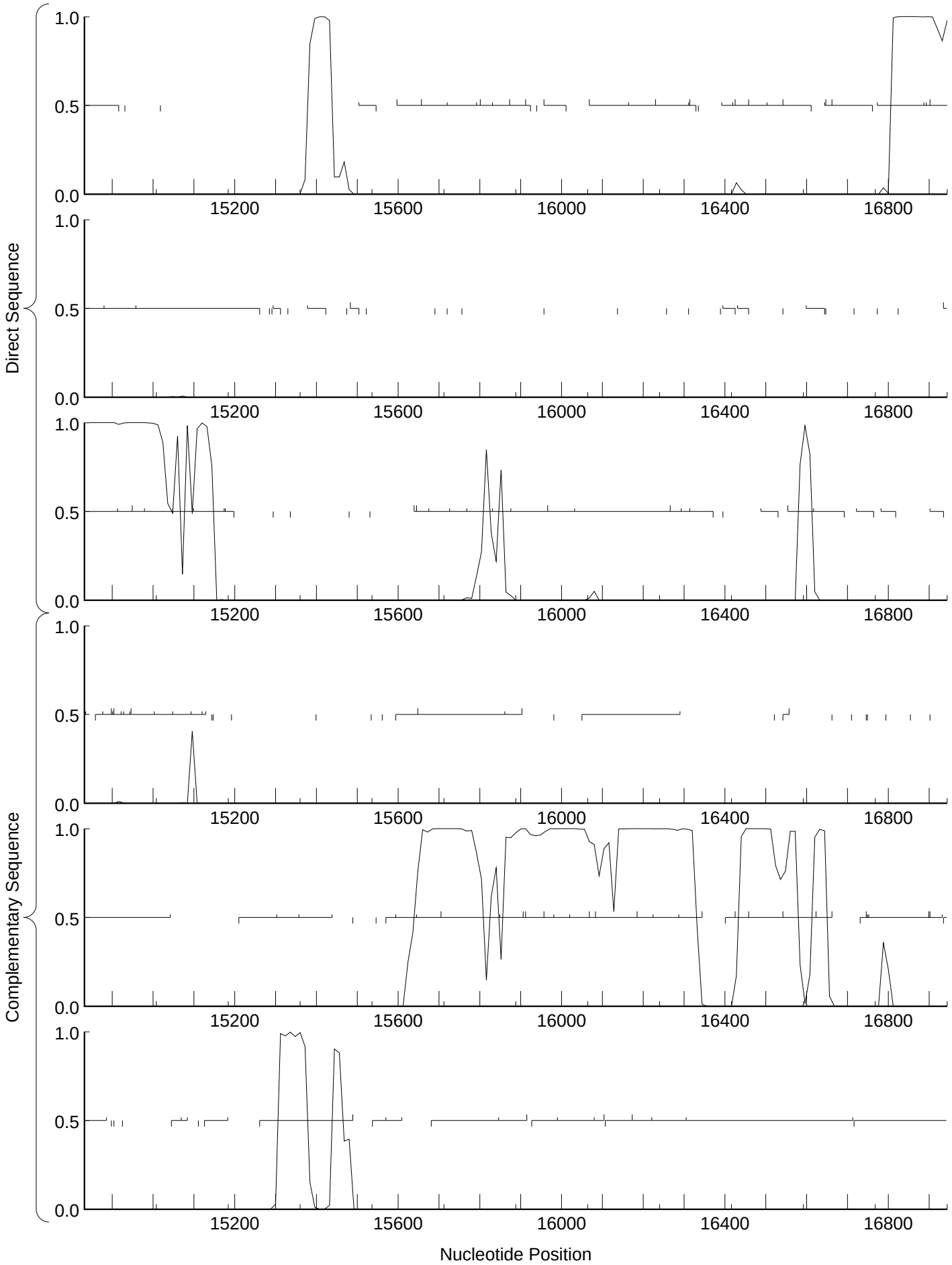
Mycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 7/34



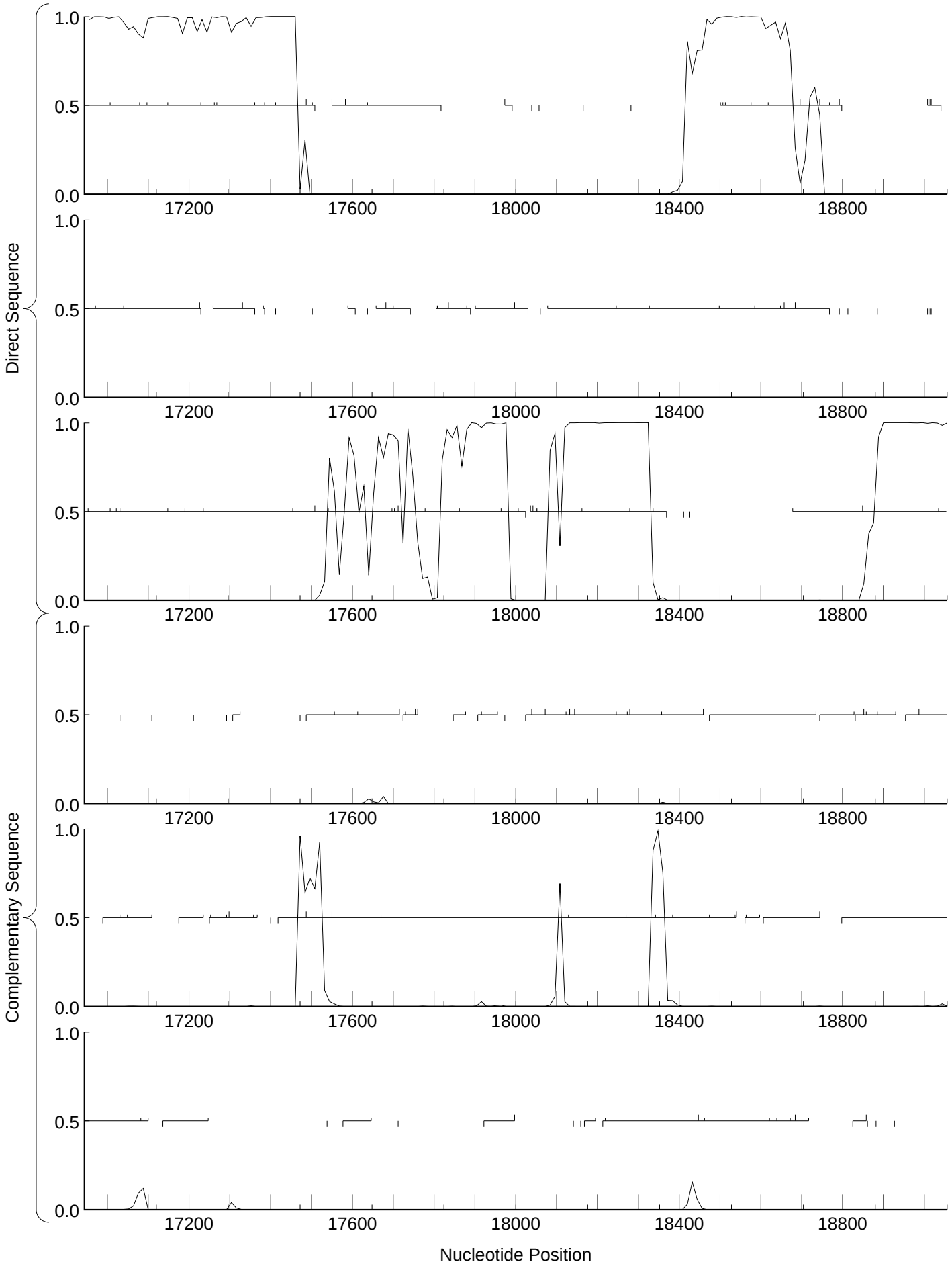
Mycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 8/34



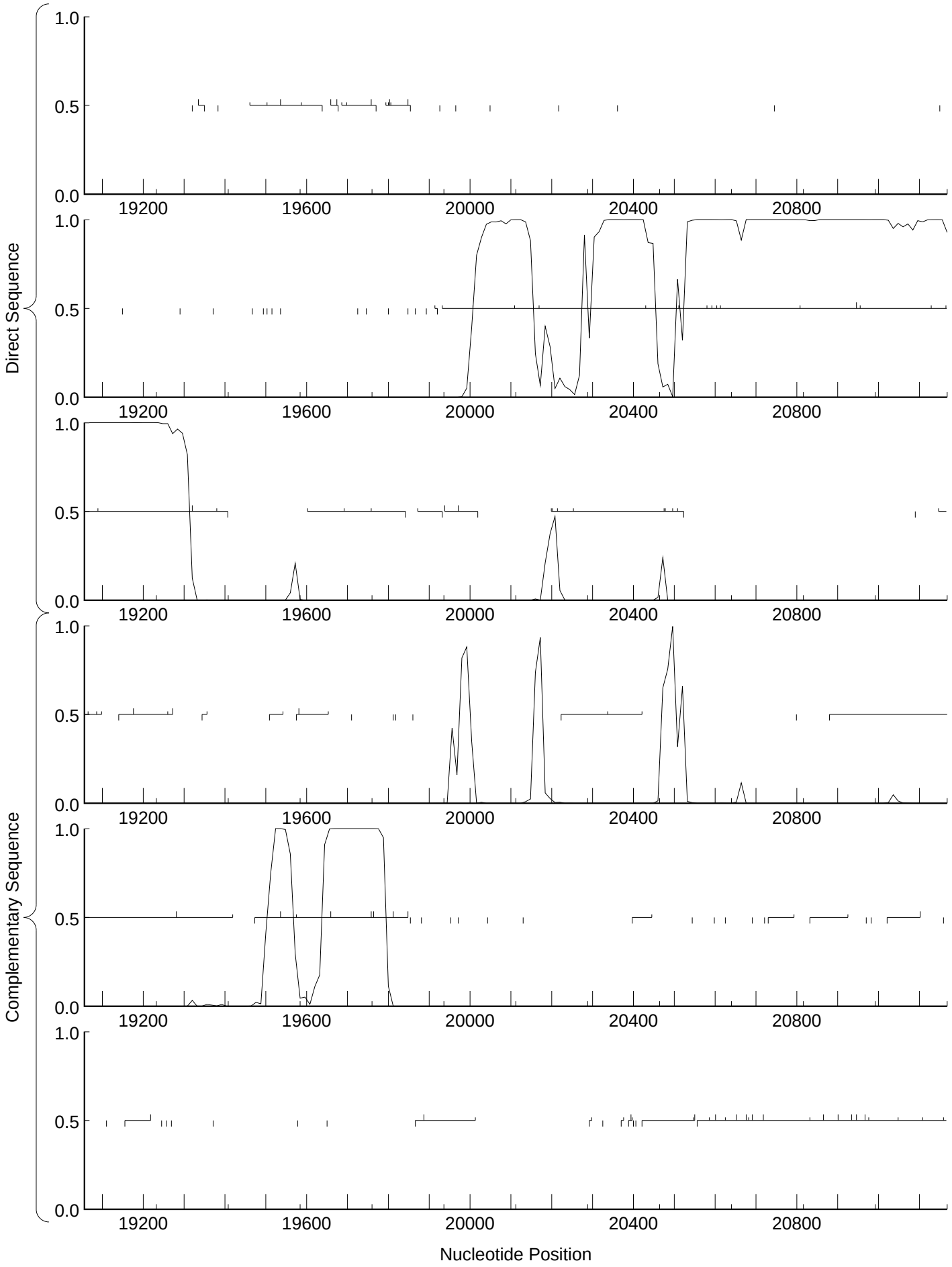
Mycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 9/34



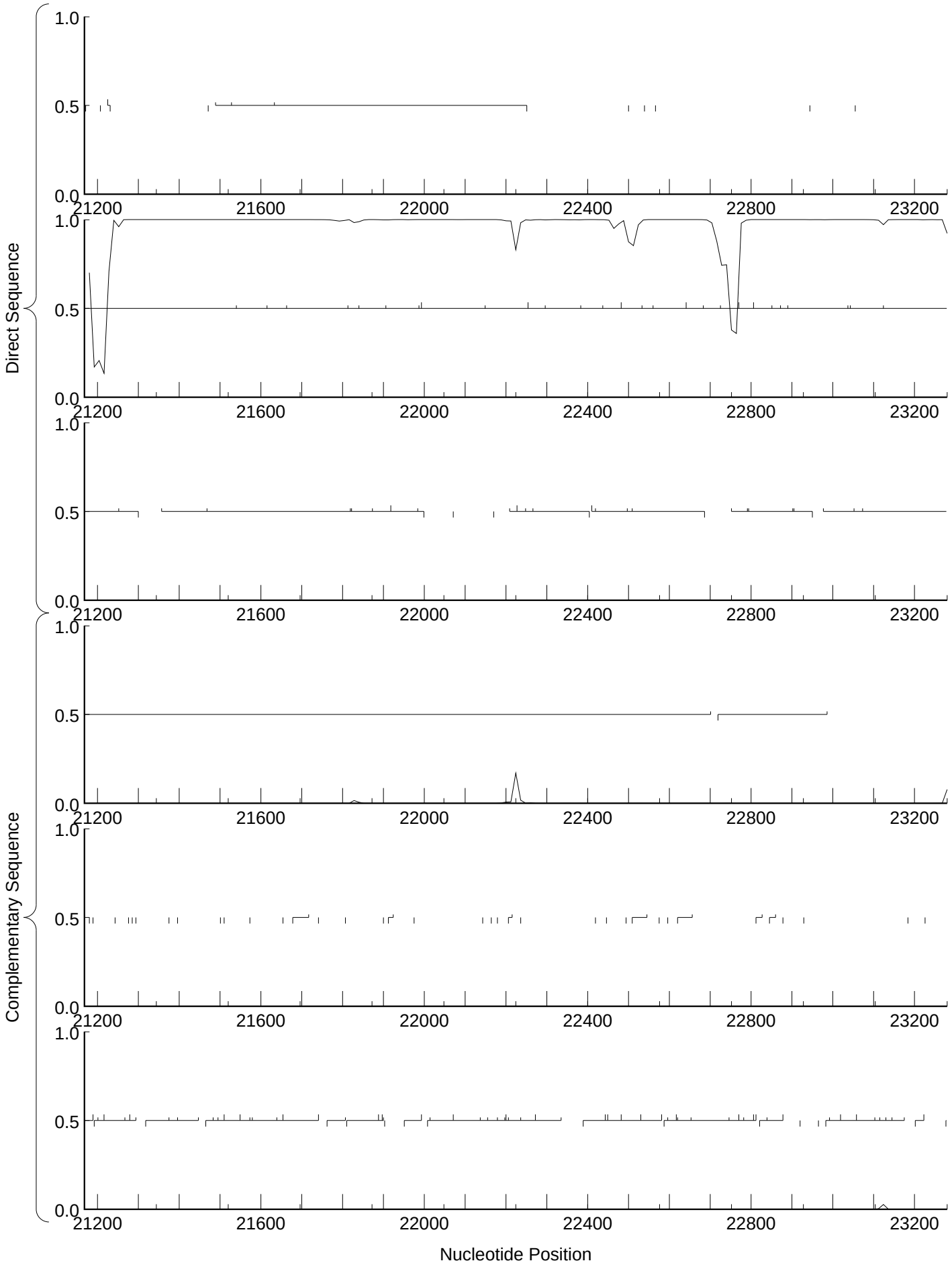
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 10/34



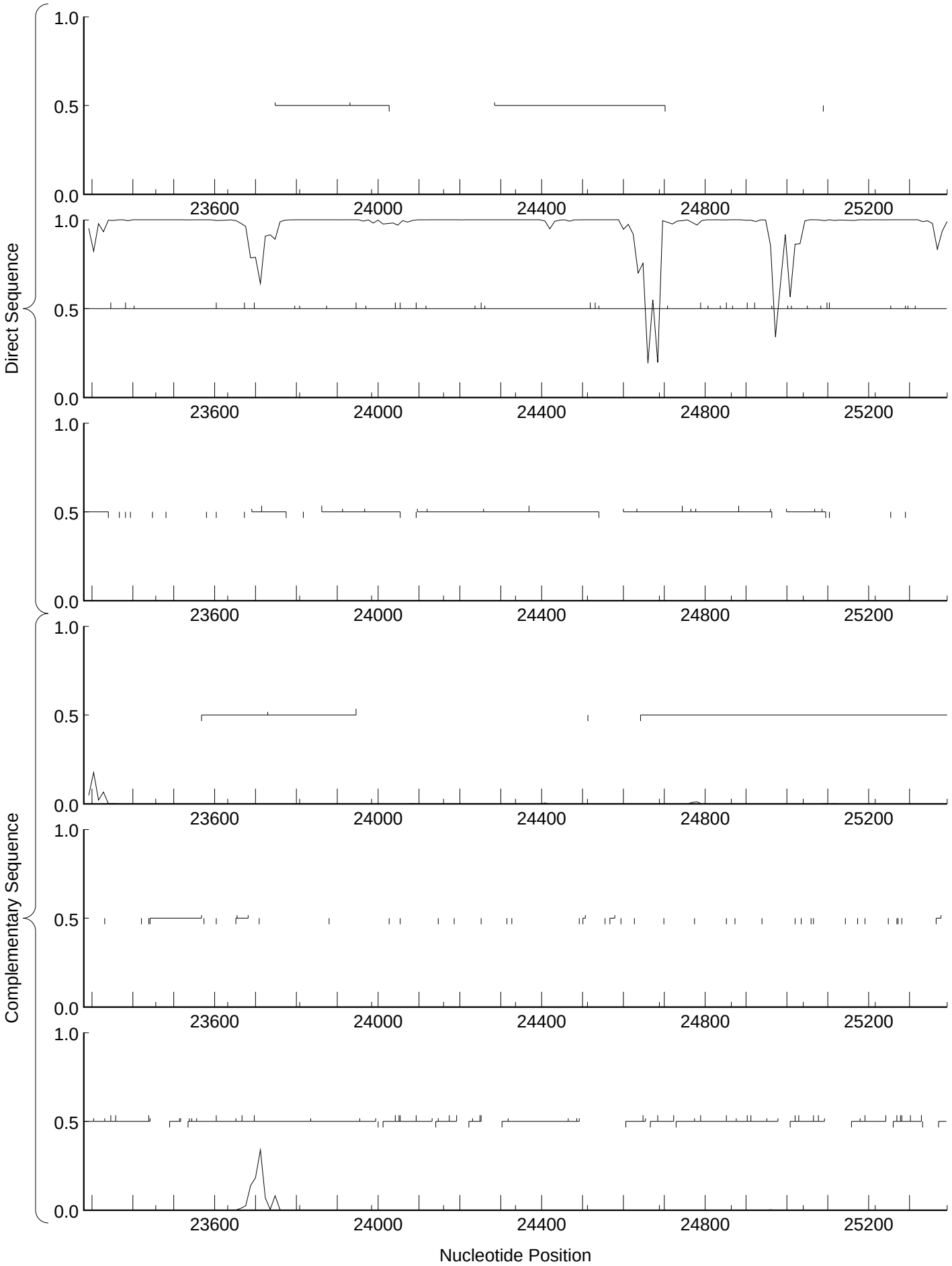
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 11/34



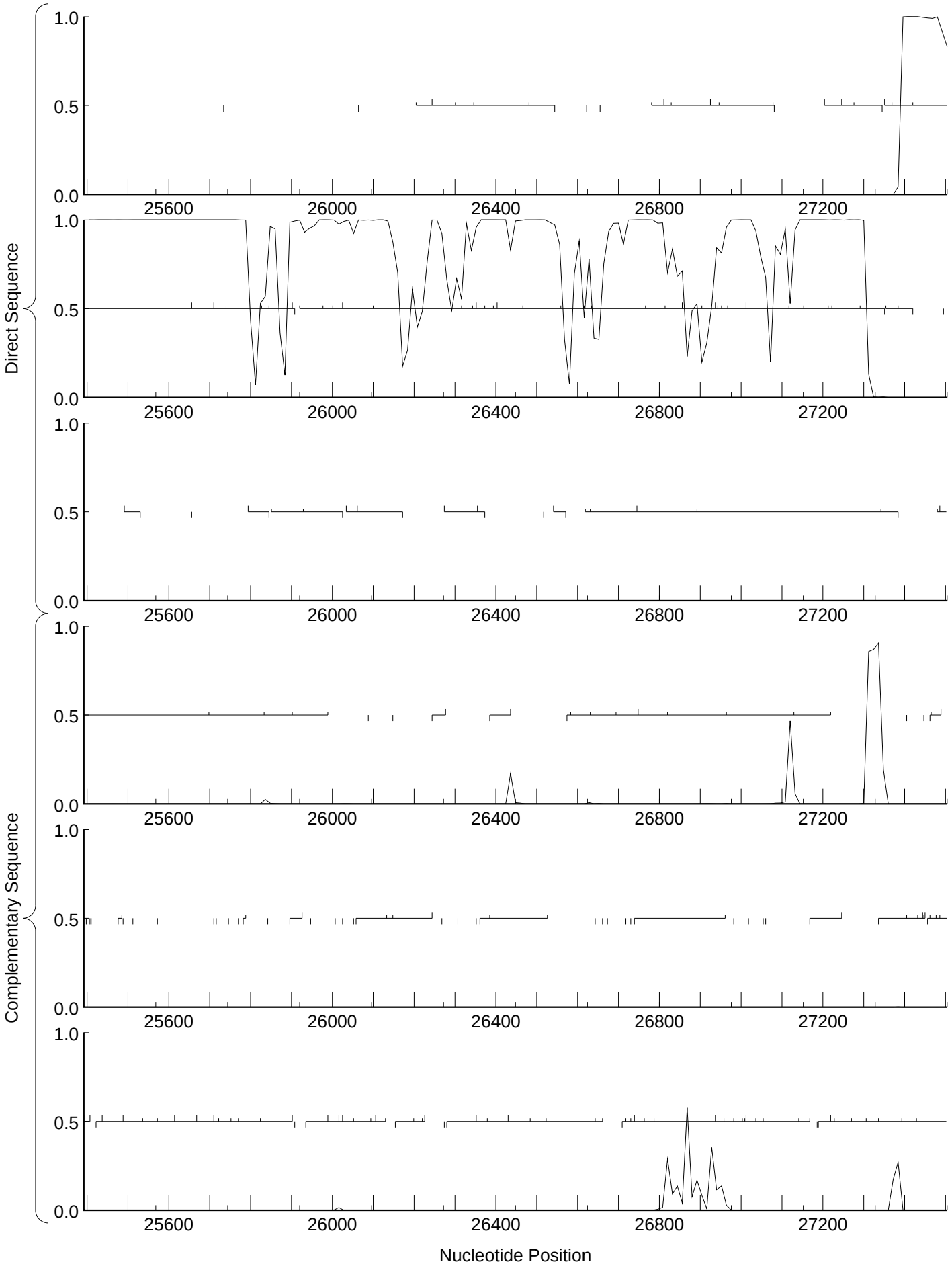
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 12/34



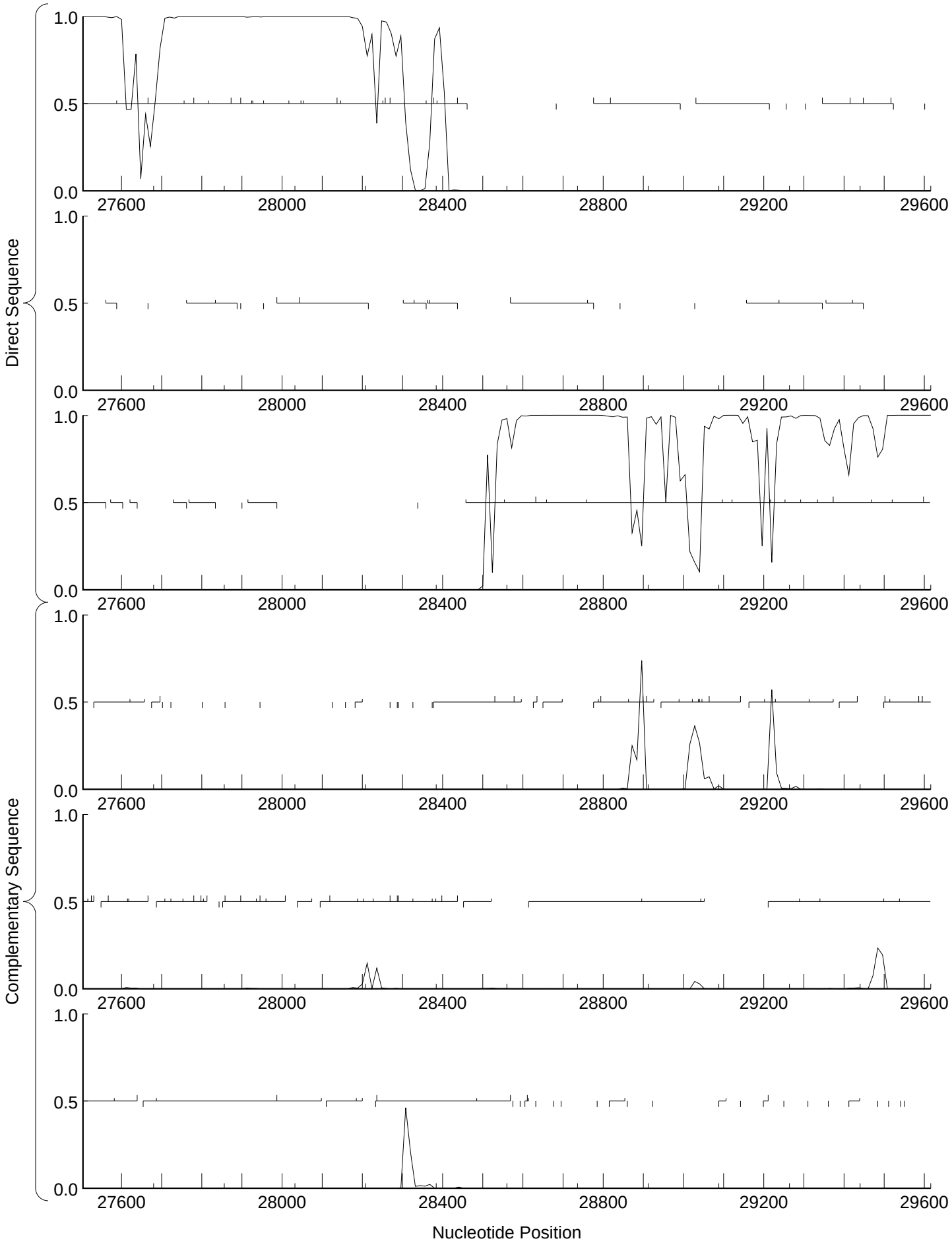
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 13/34



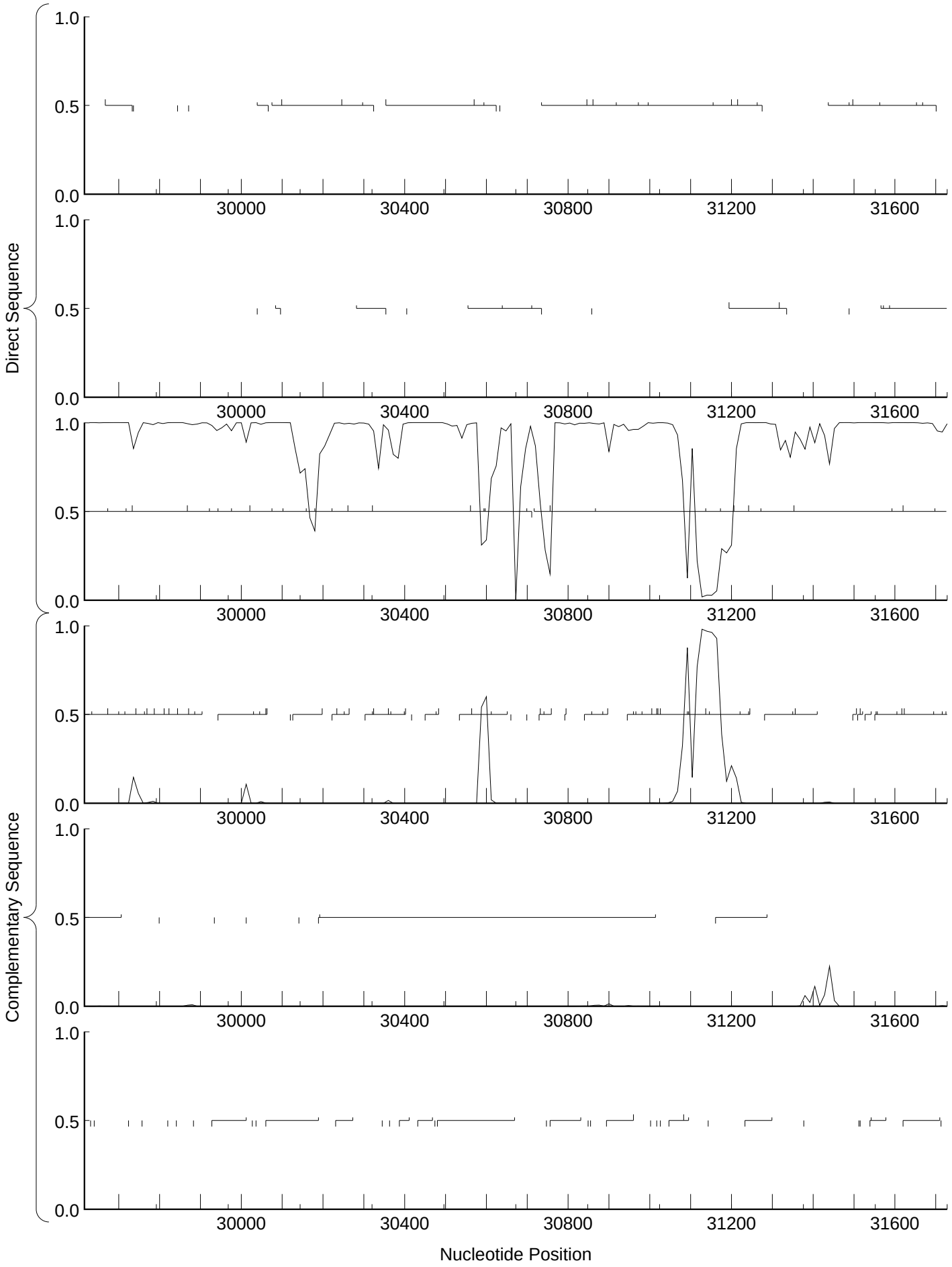
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 14/34



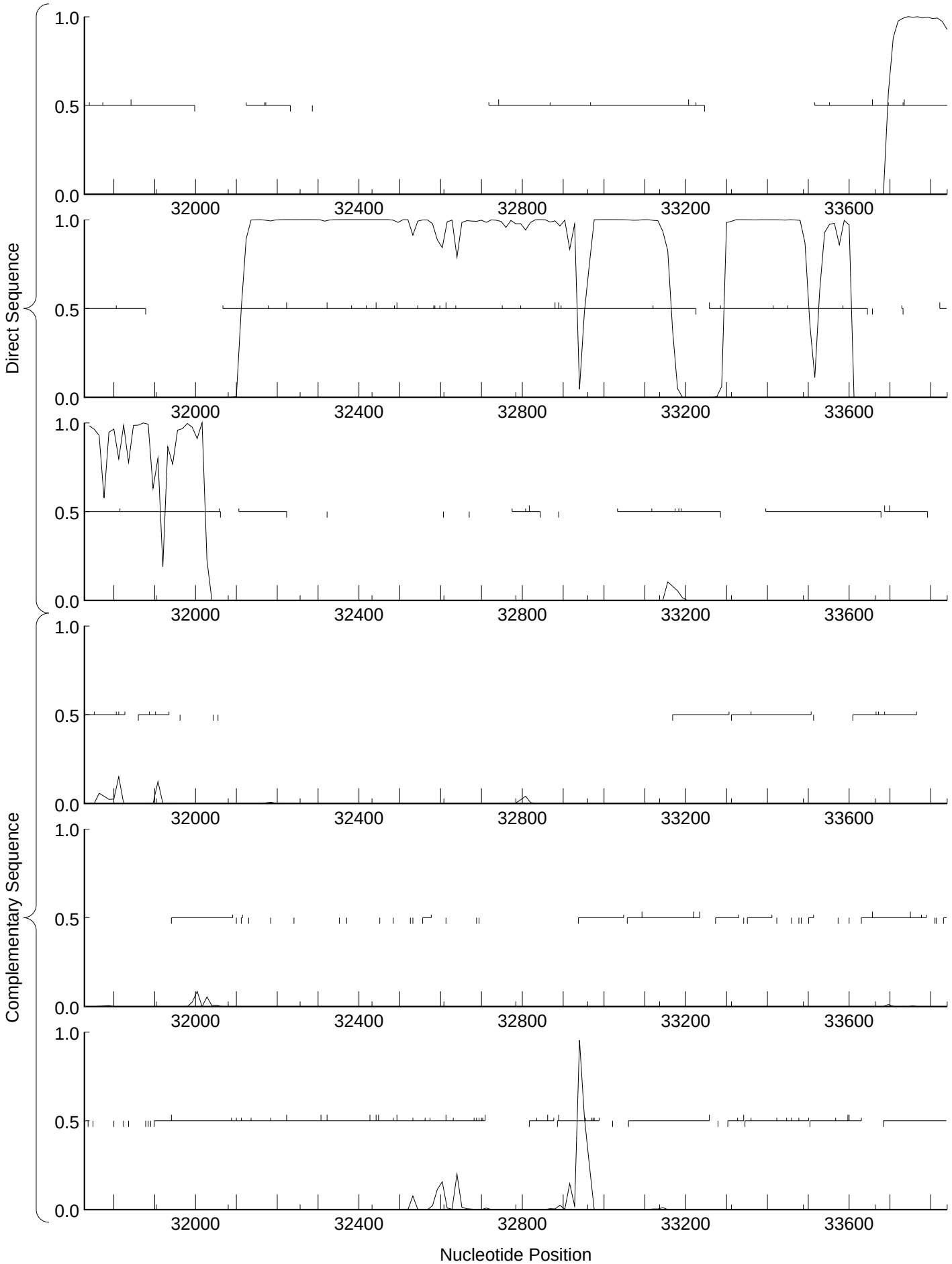
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 15/34



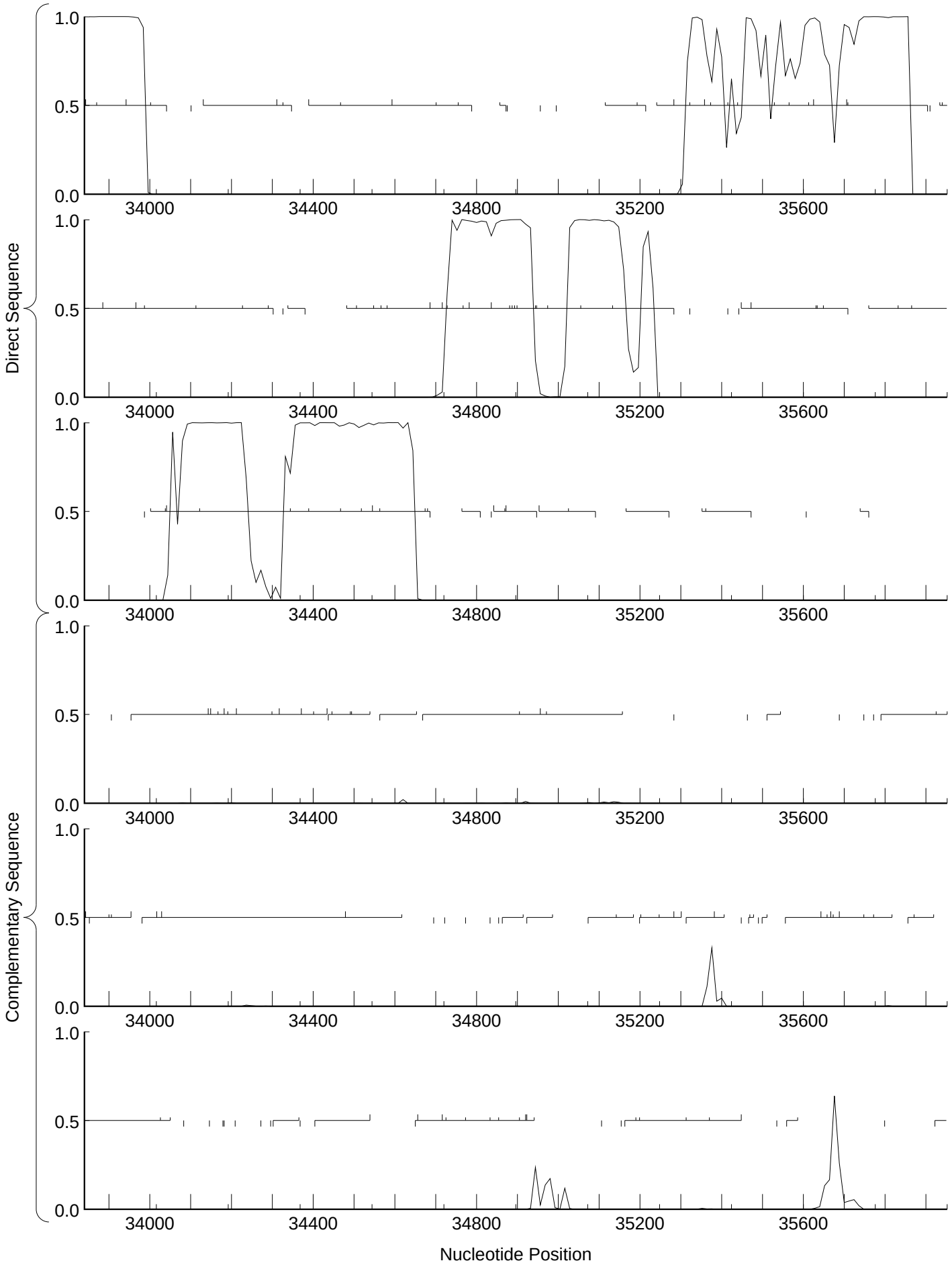
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 16/34



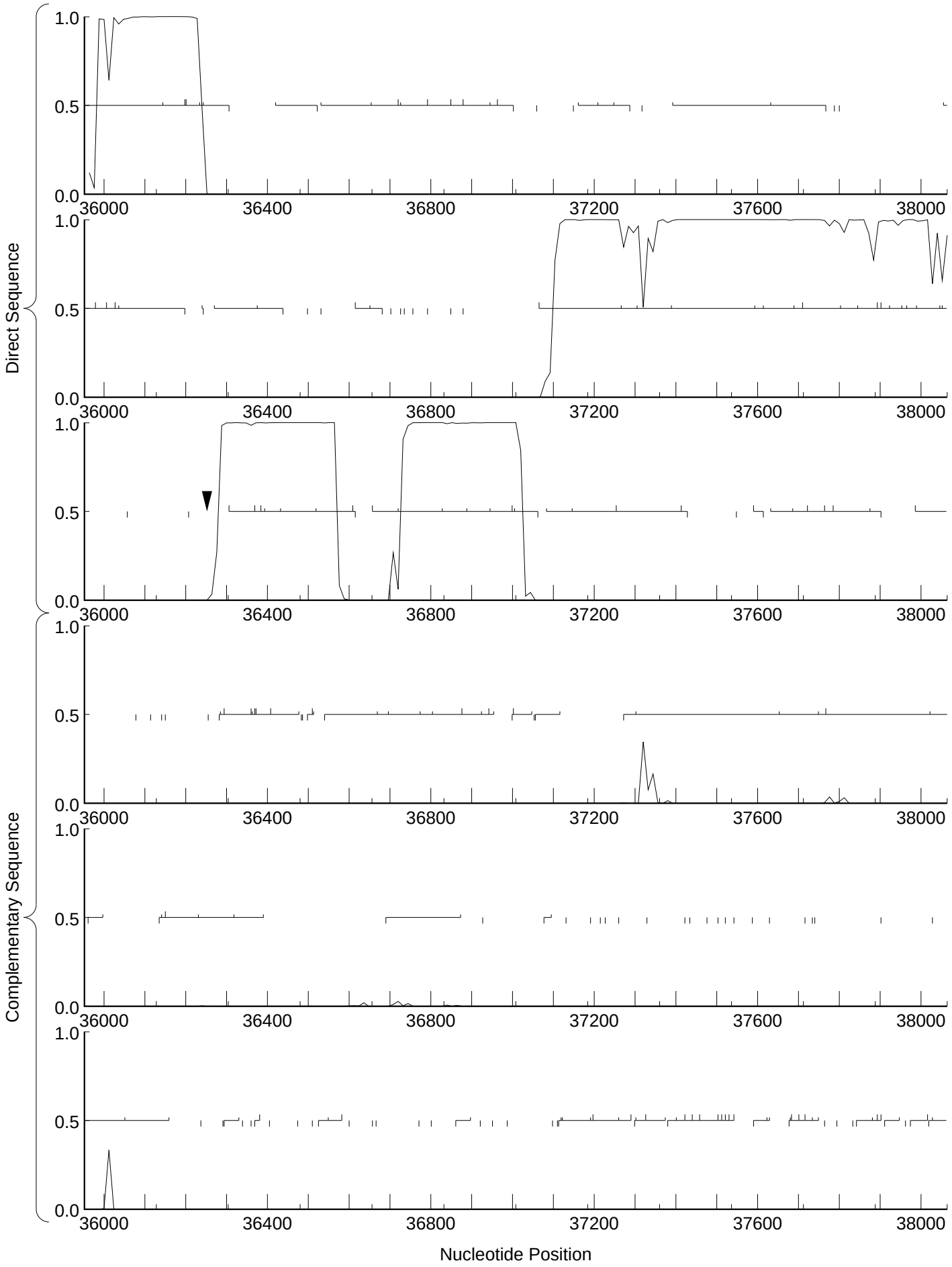
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 17/34



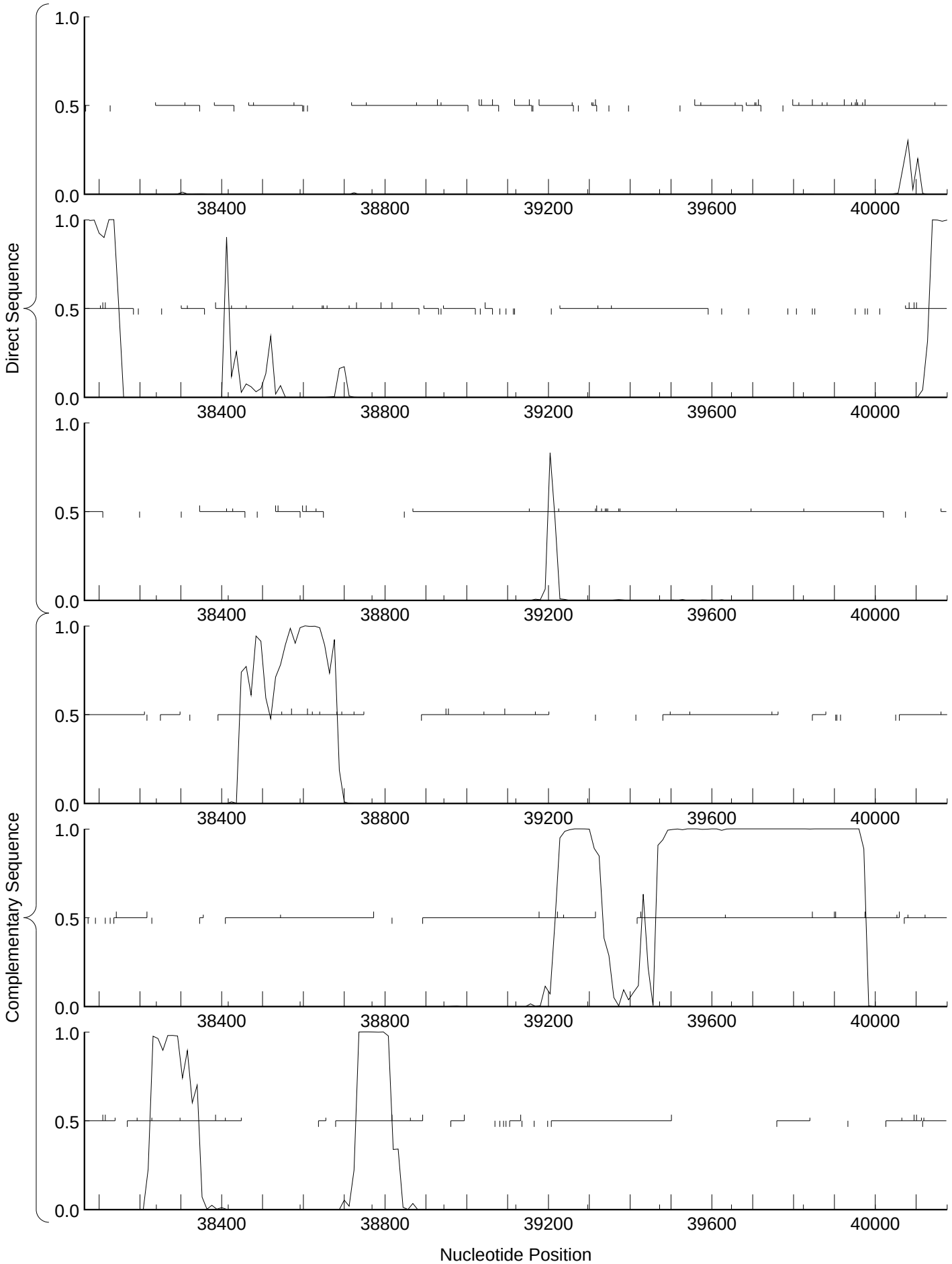
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 18/34



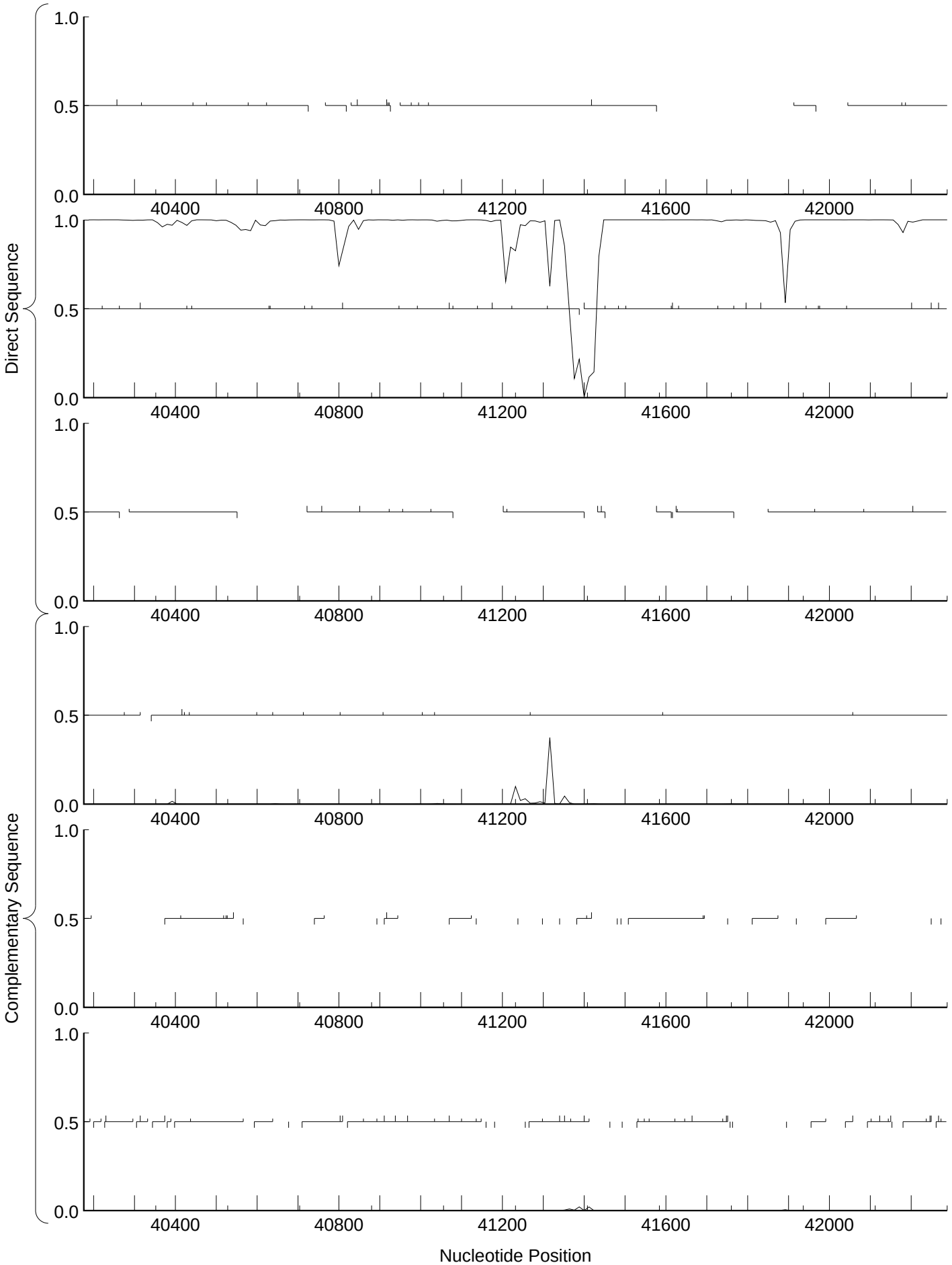
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 19/34



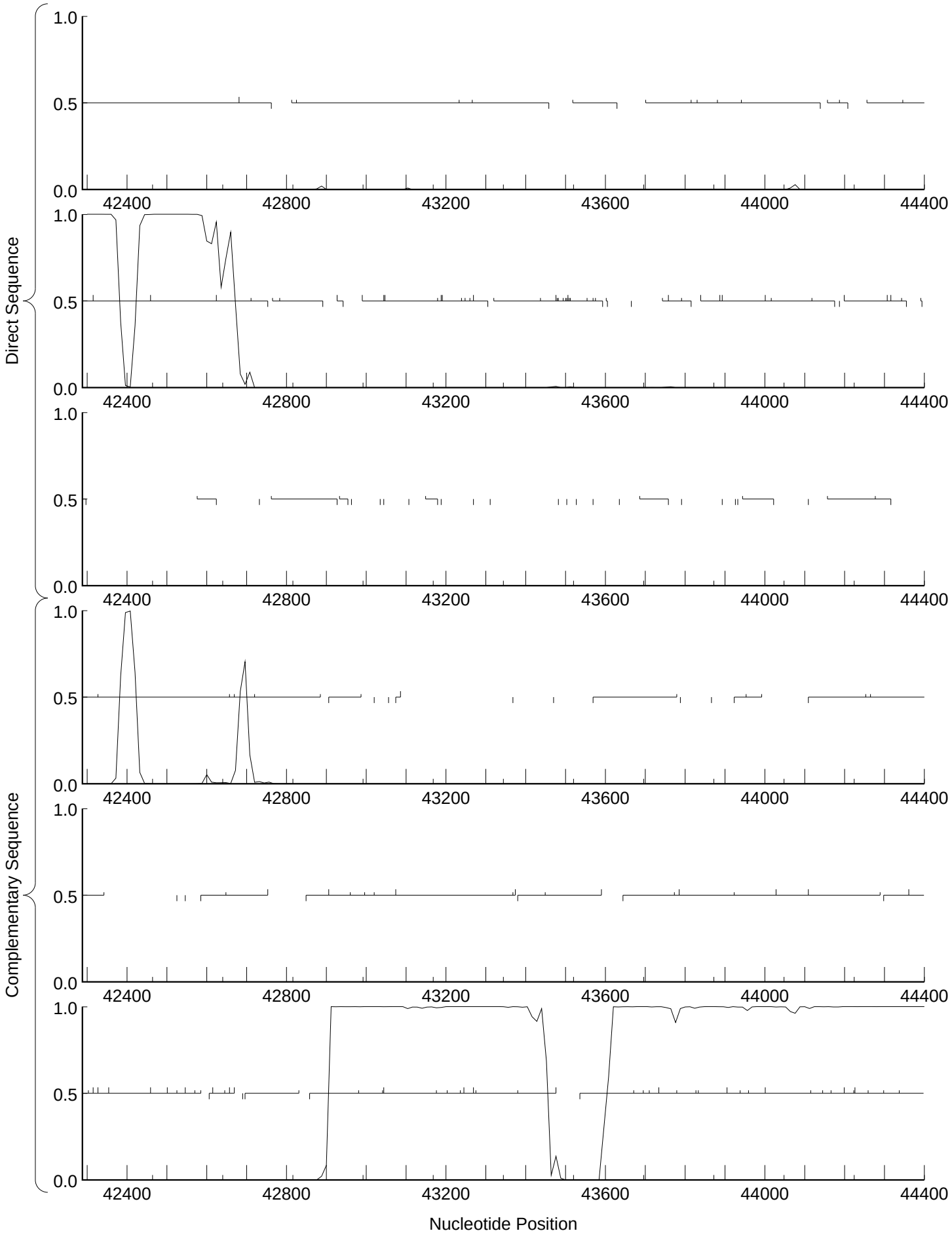
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 20/34



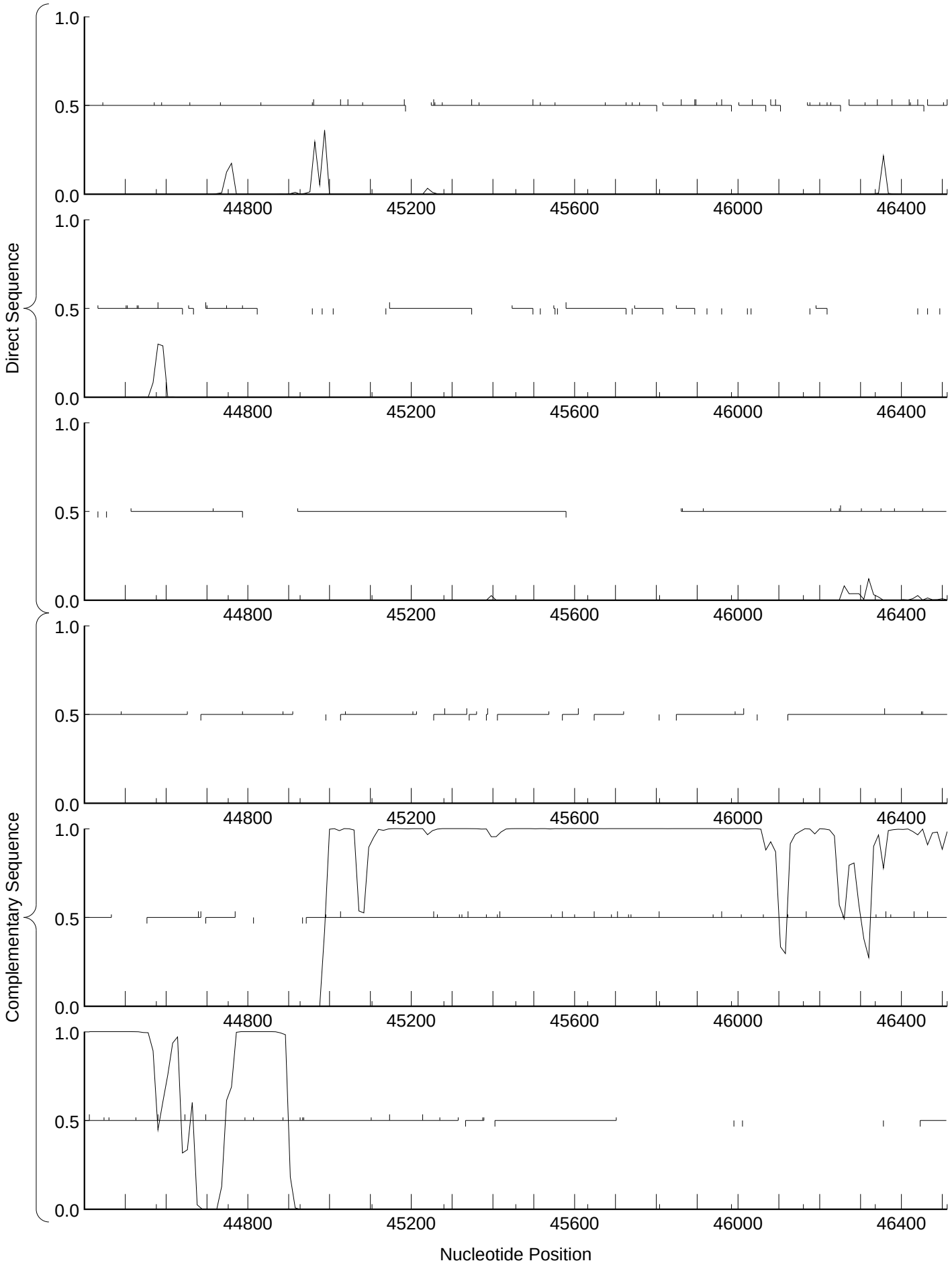
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 21/34



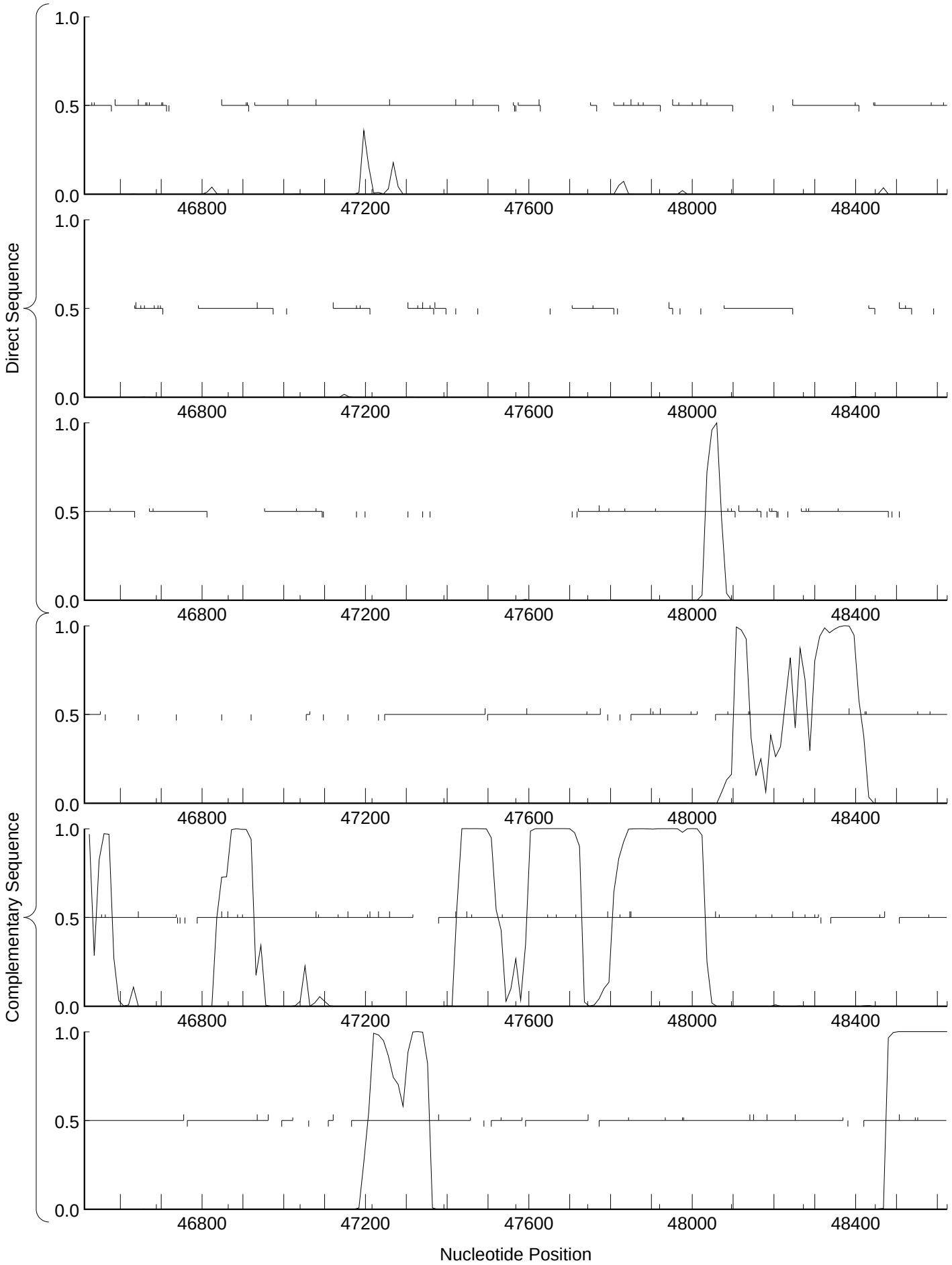
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 22/34



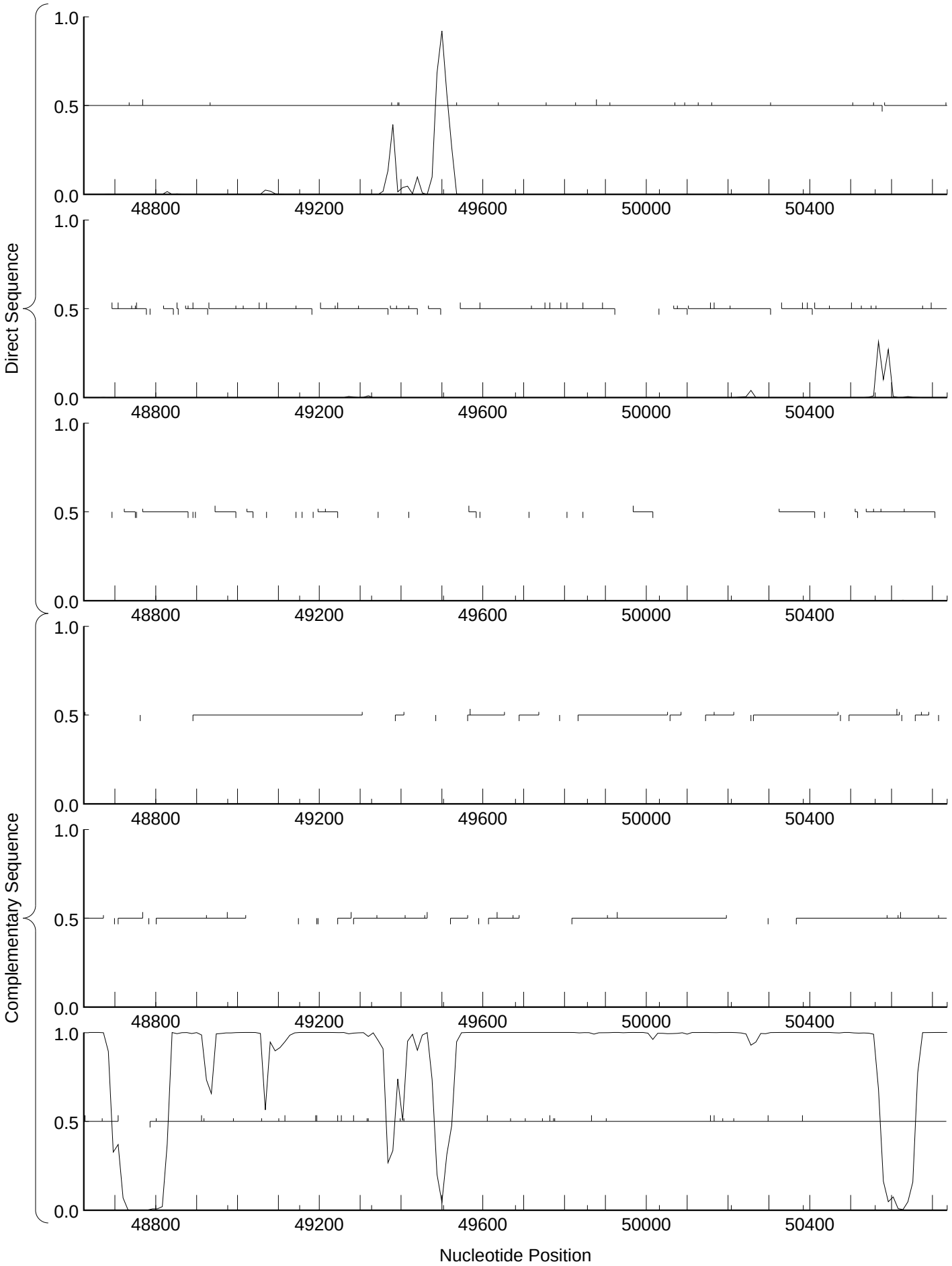
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 23/34



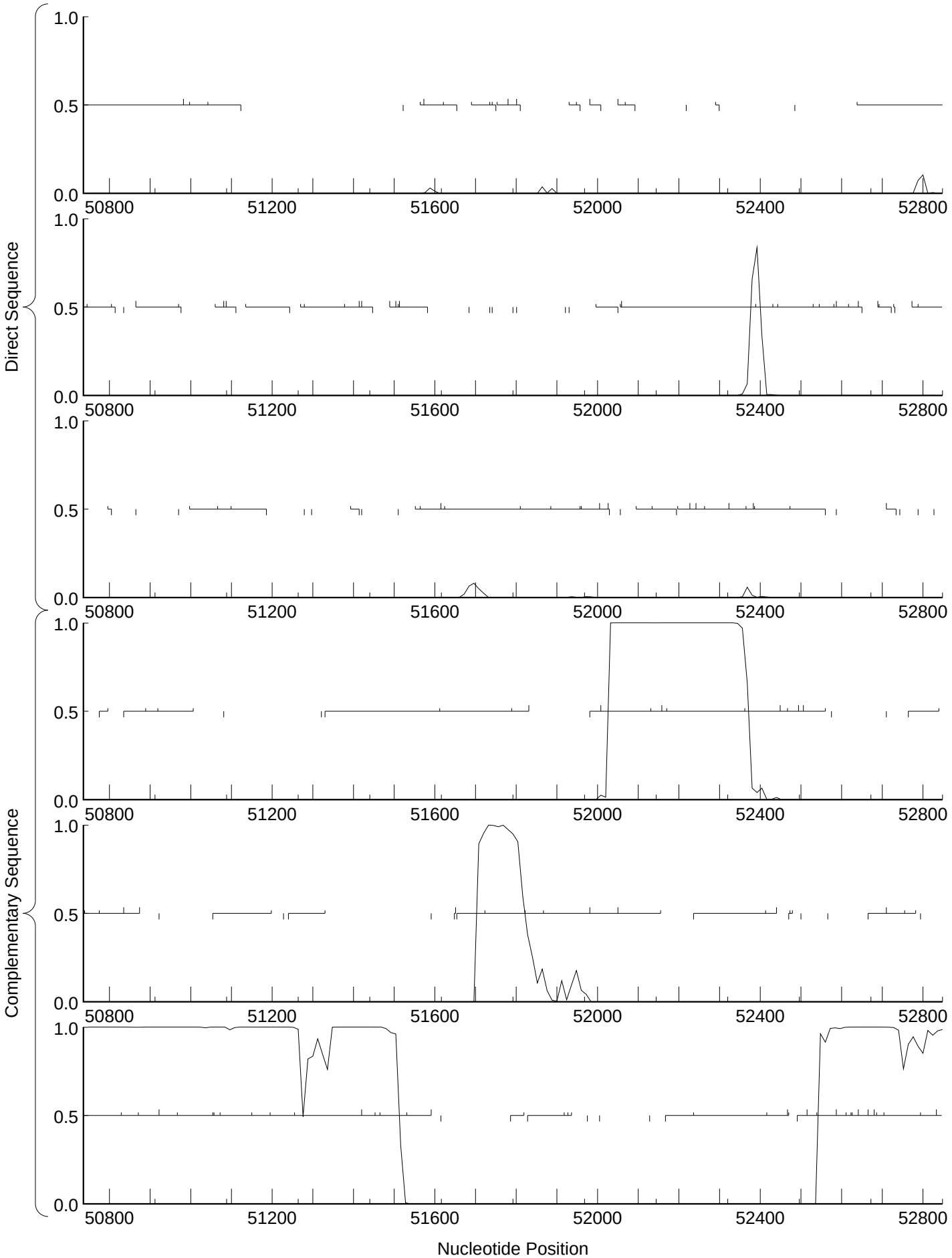
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 24/34



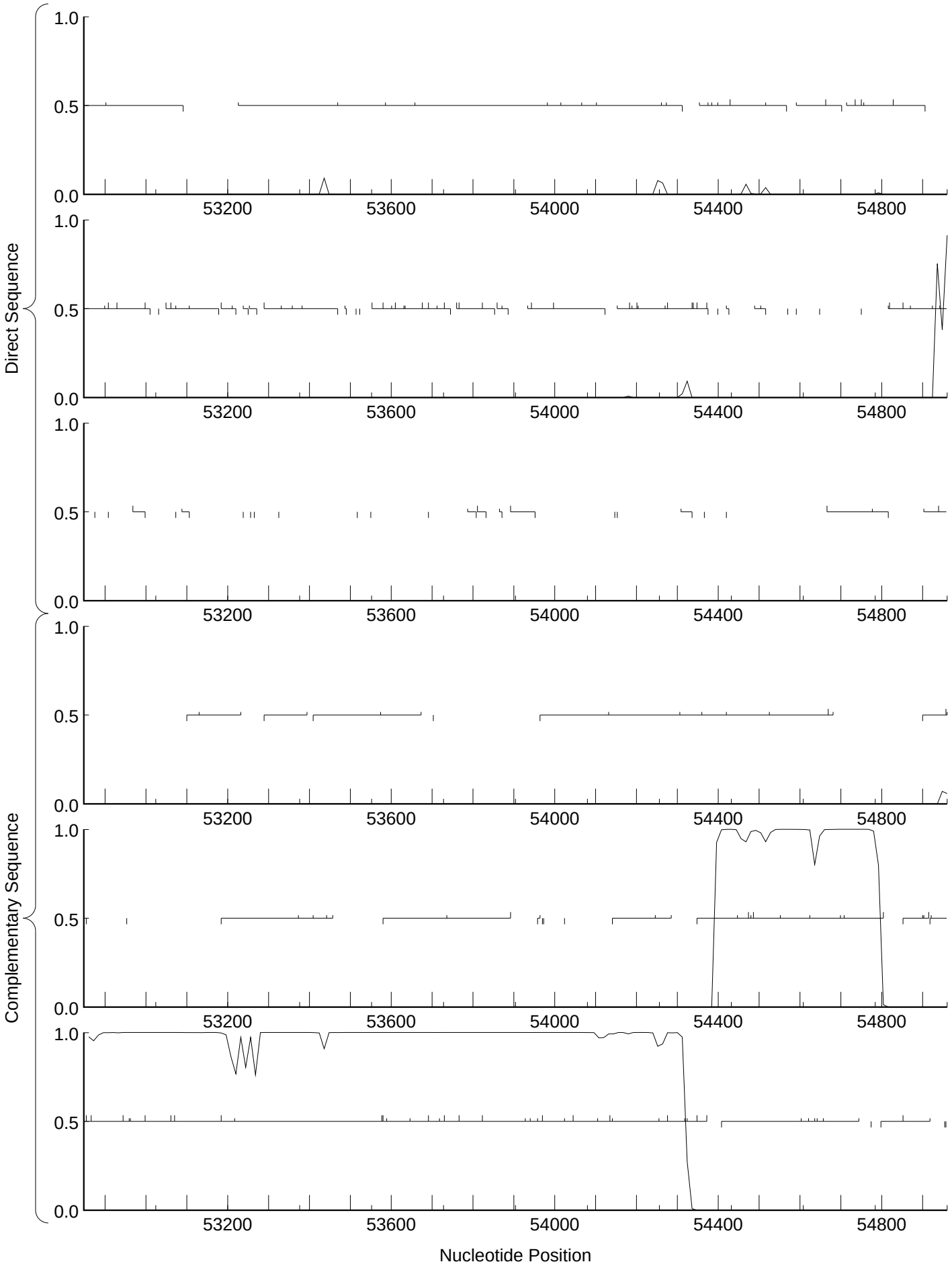
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 25/34



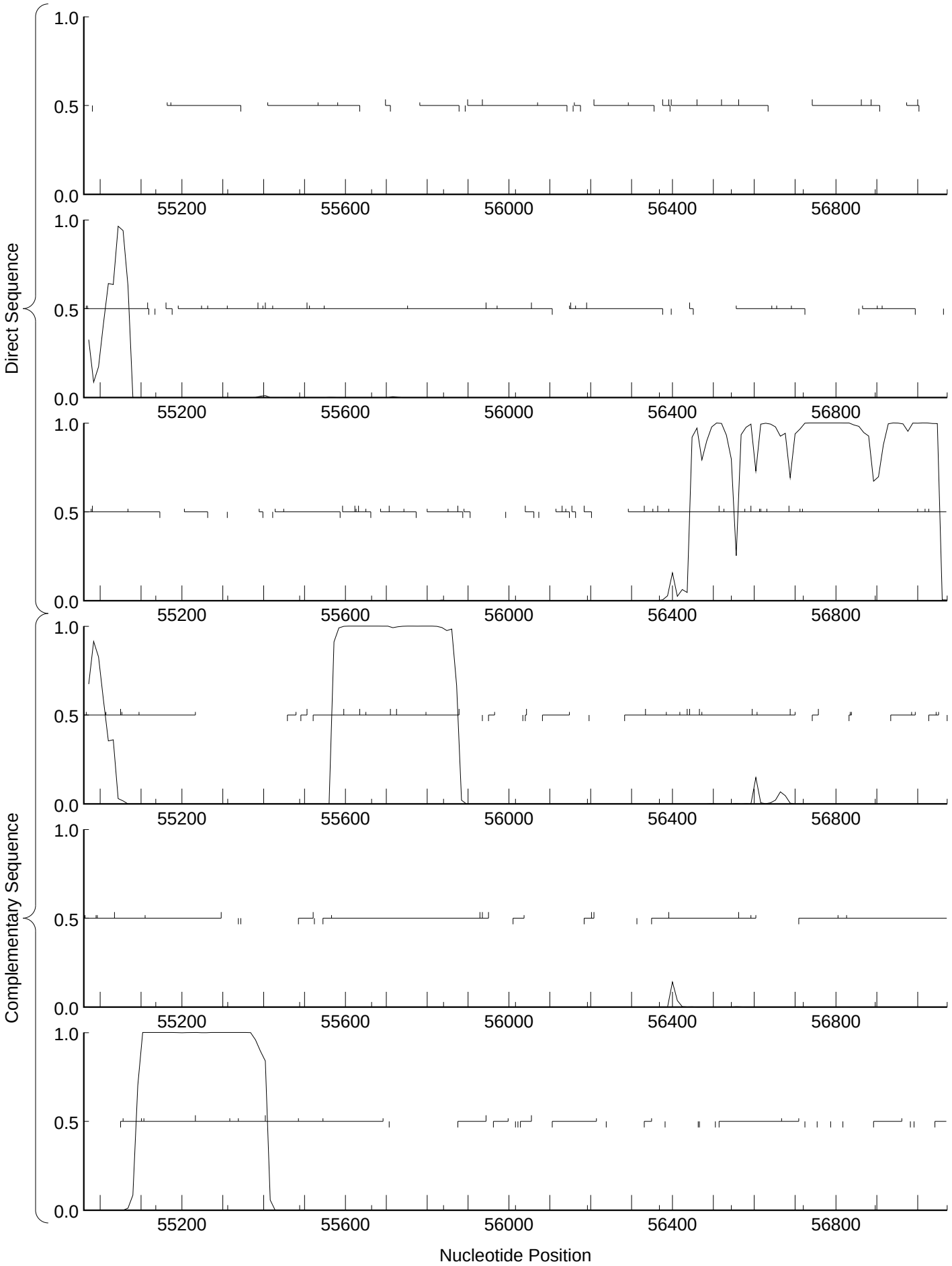
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 26/34



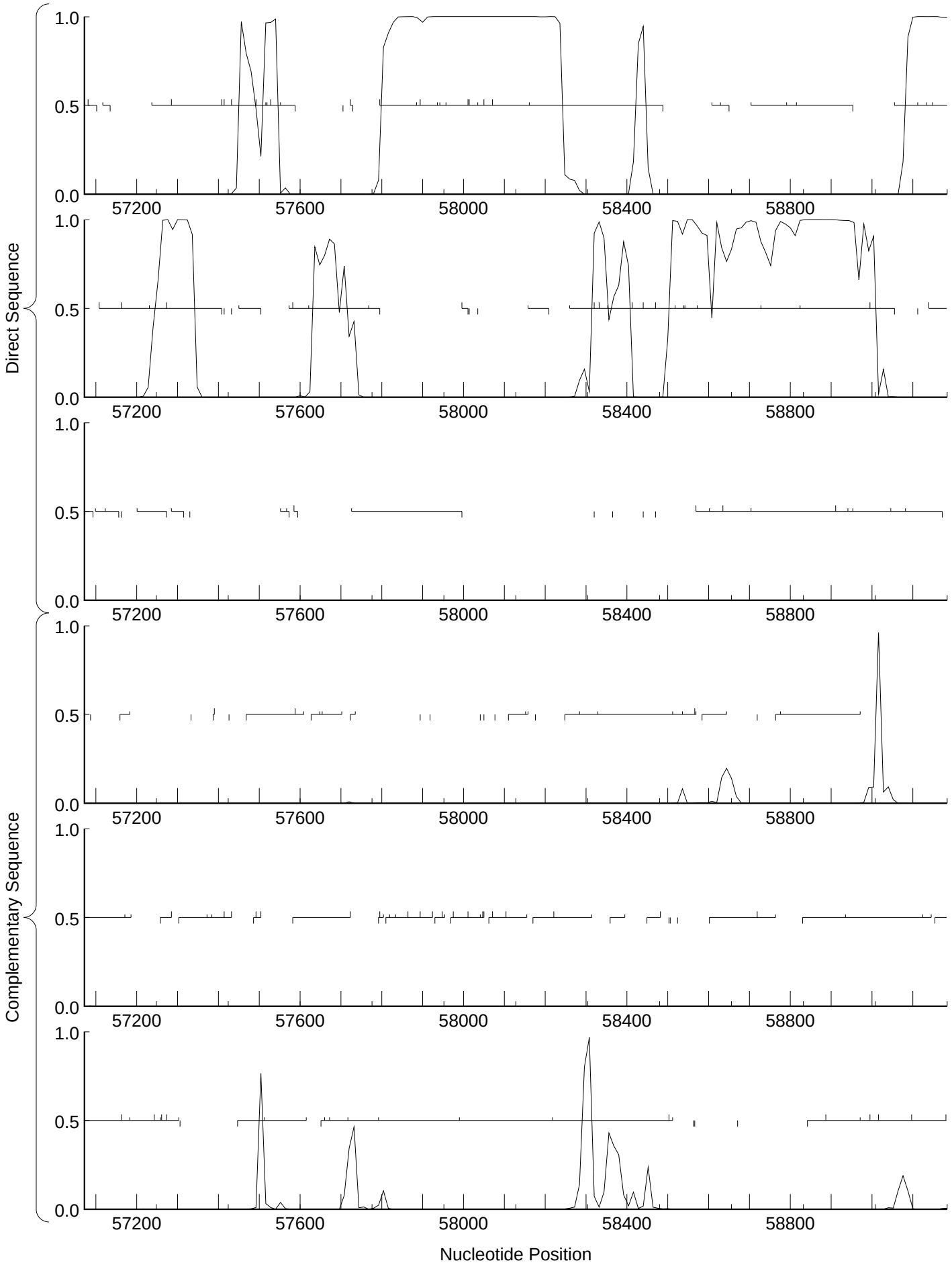
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 27/34



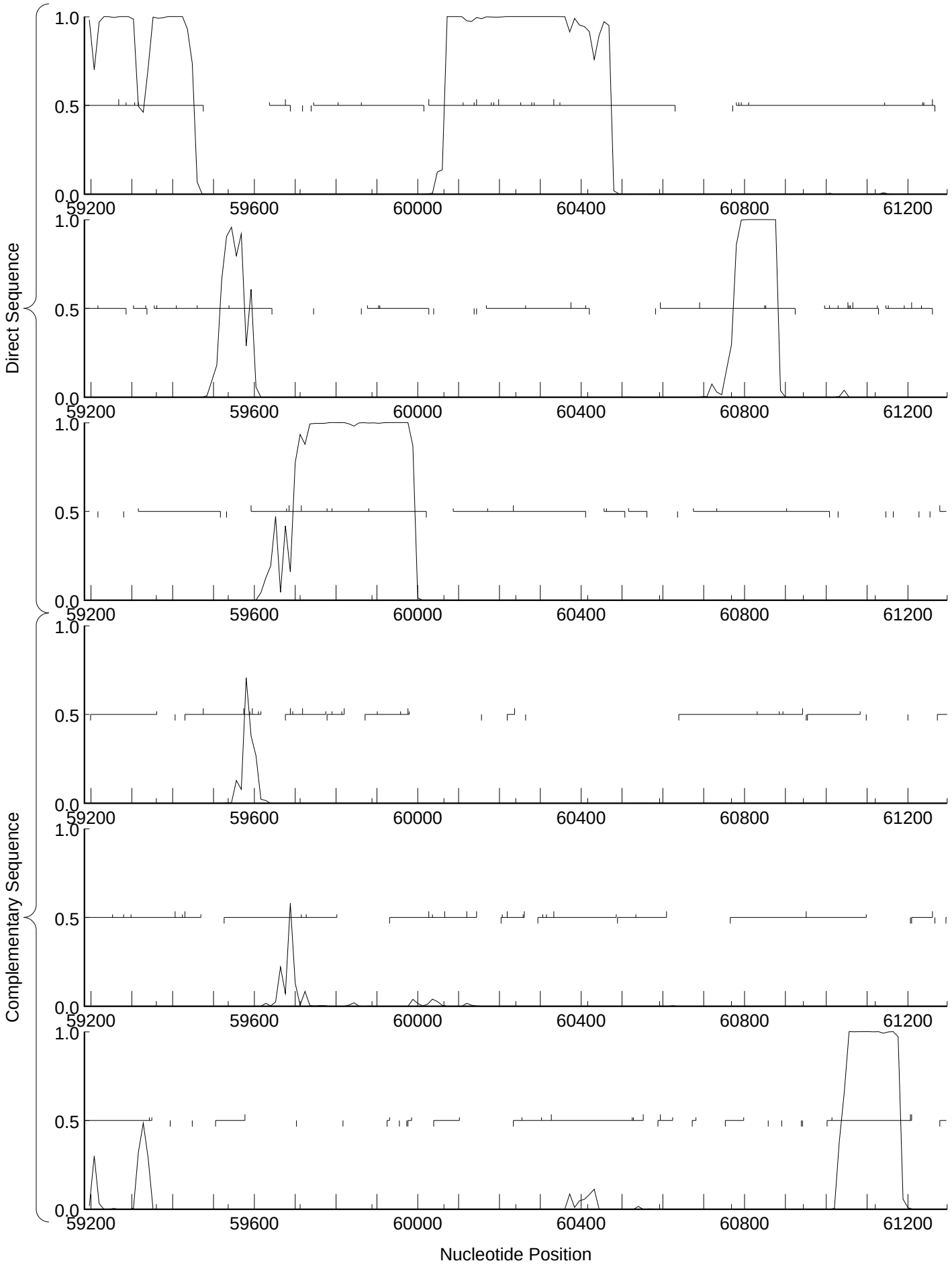
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 28/34



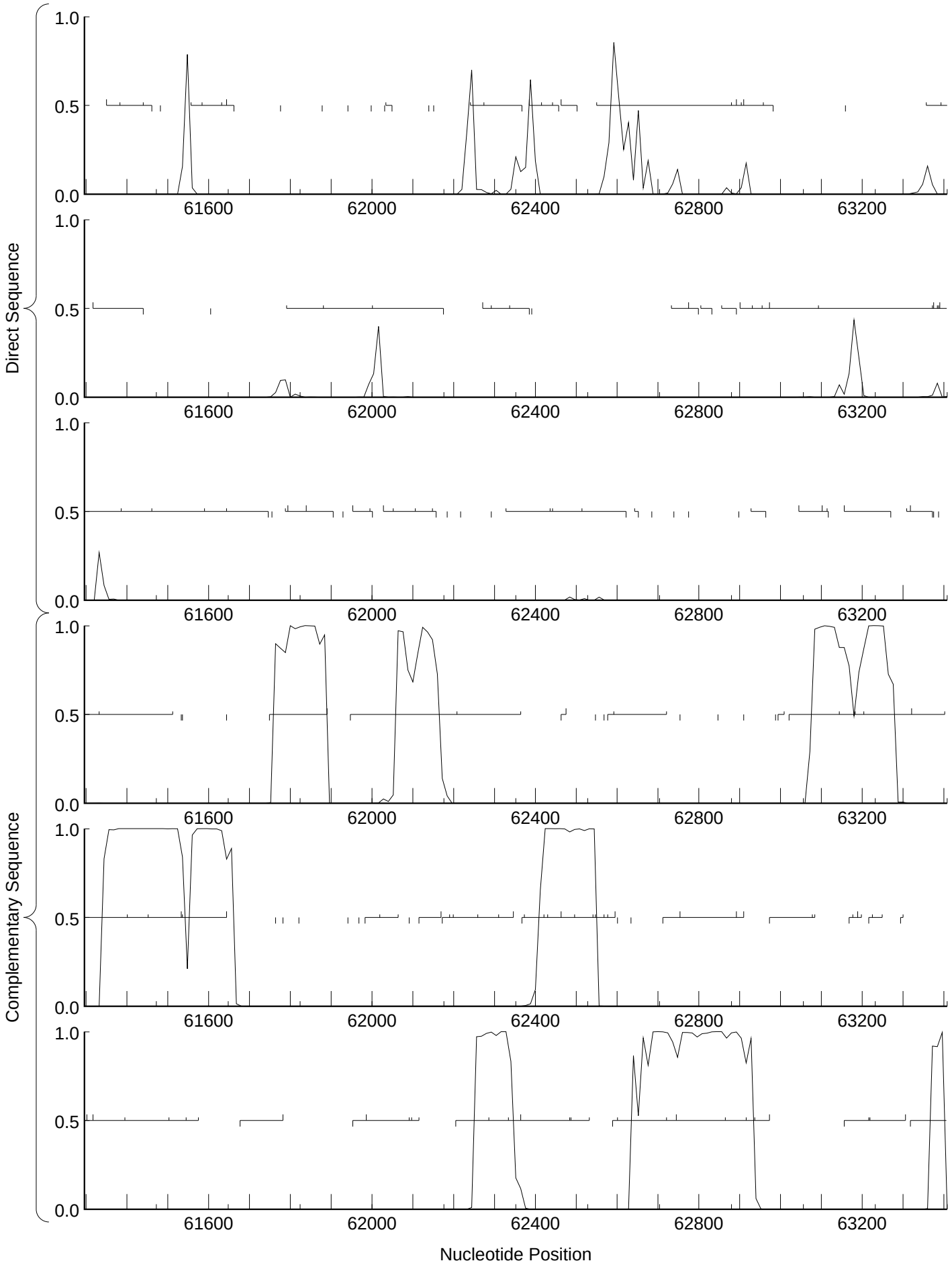
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 29/34



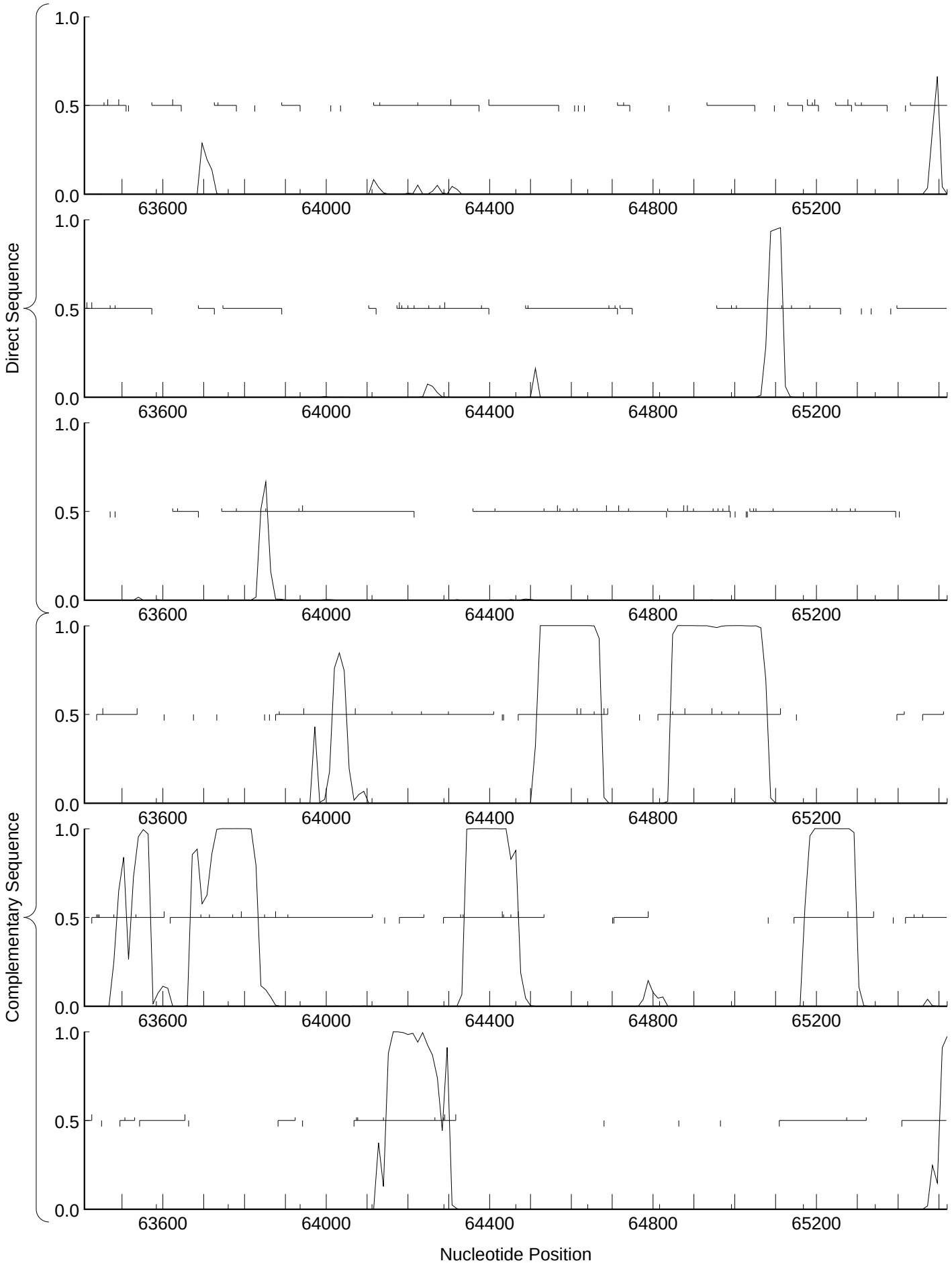
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 30/34



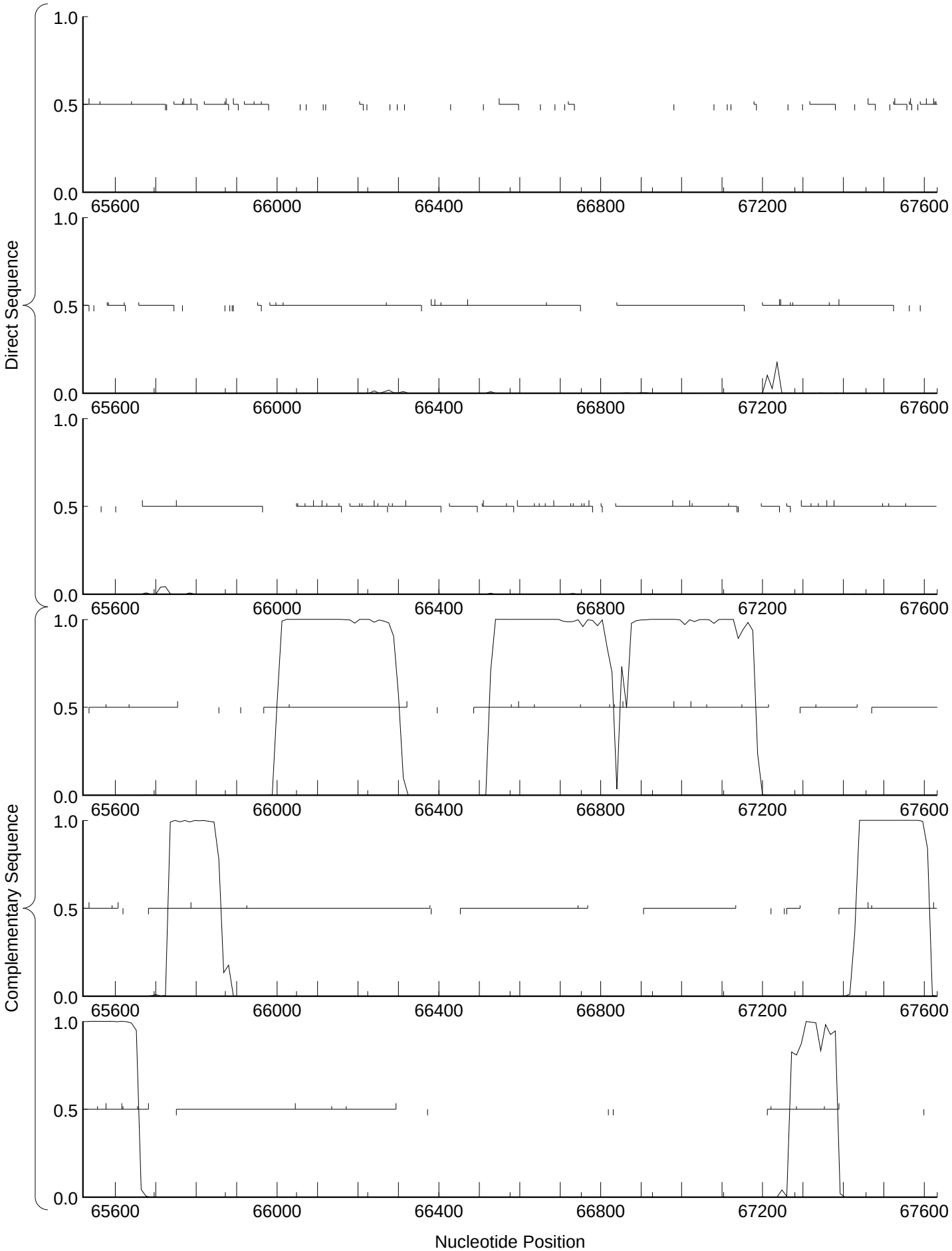
ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 31/34



ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 32/34



ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 33/34



ycobacterium phage MRabcd complete sequence, 68791 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 34/34

