GeneMark

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

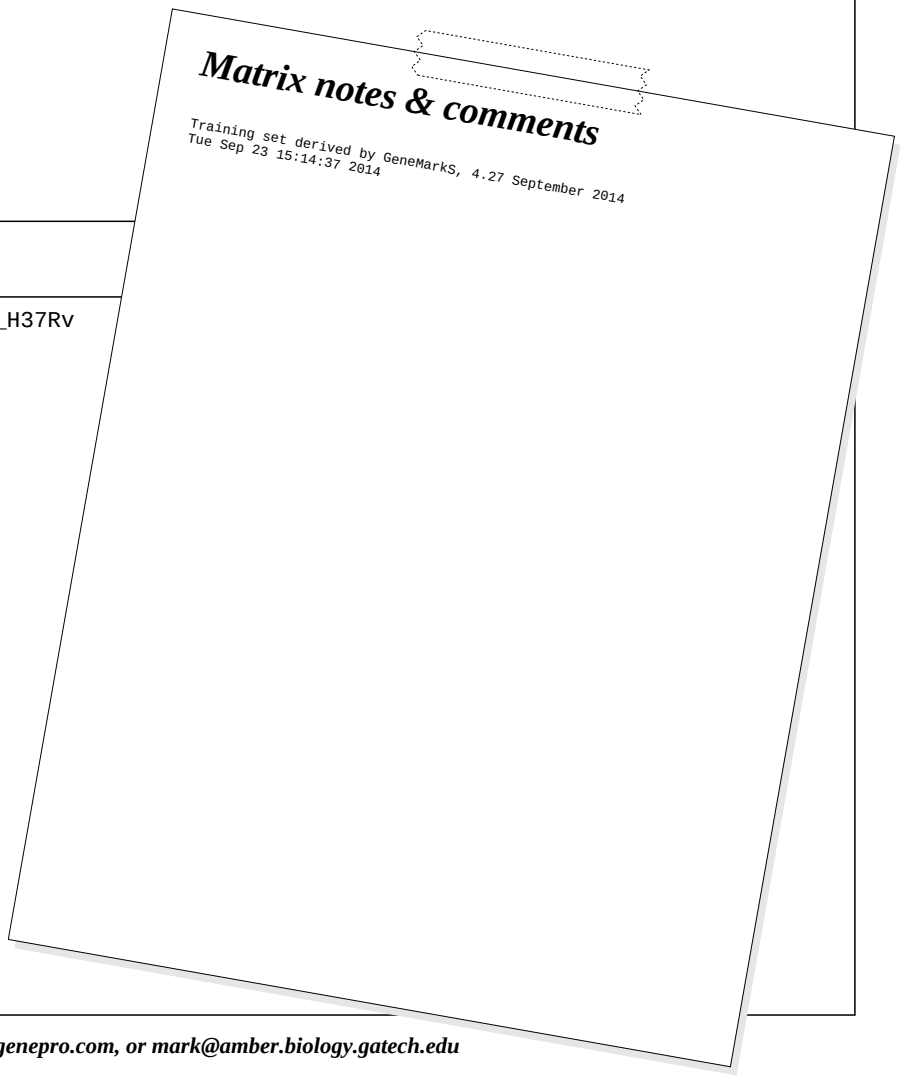
PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster
Analysis Date    :  3/29/24 at 9:05:52
Pages            :  75
Sequence Length  :  155658 bp
GC Content       :   64.67%

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

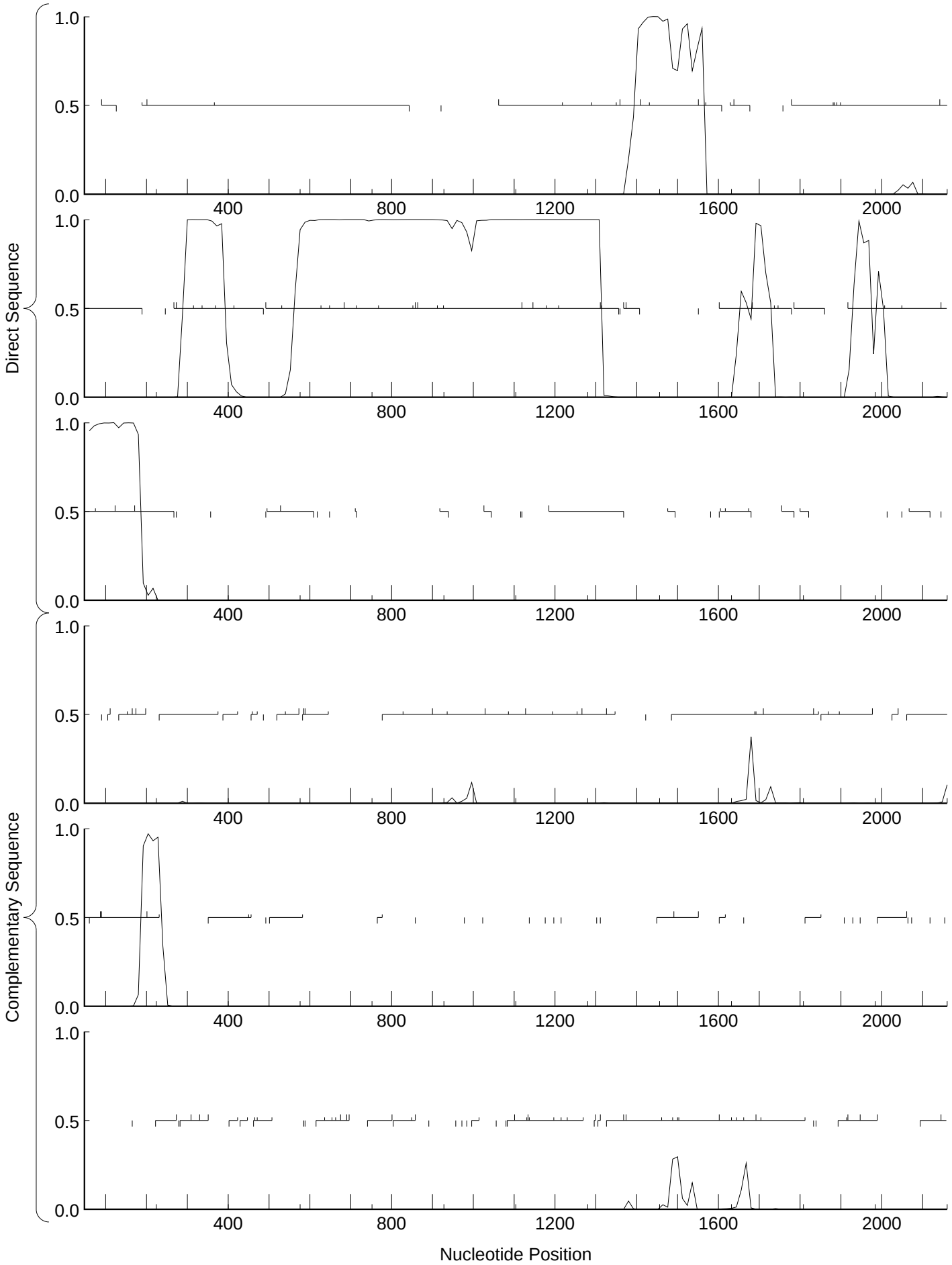
```
Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

Order      :  4
```

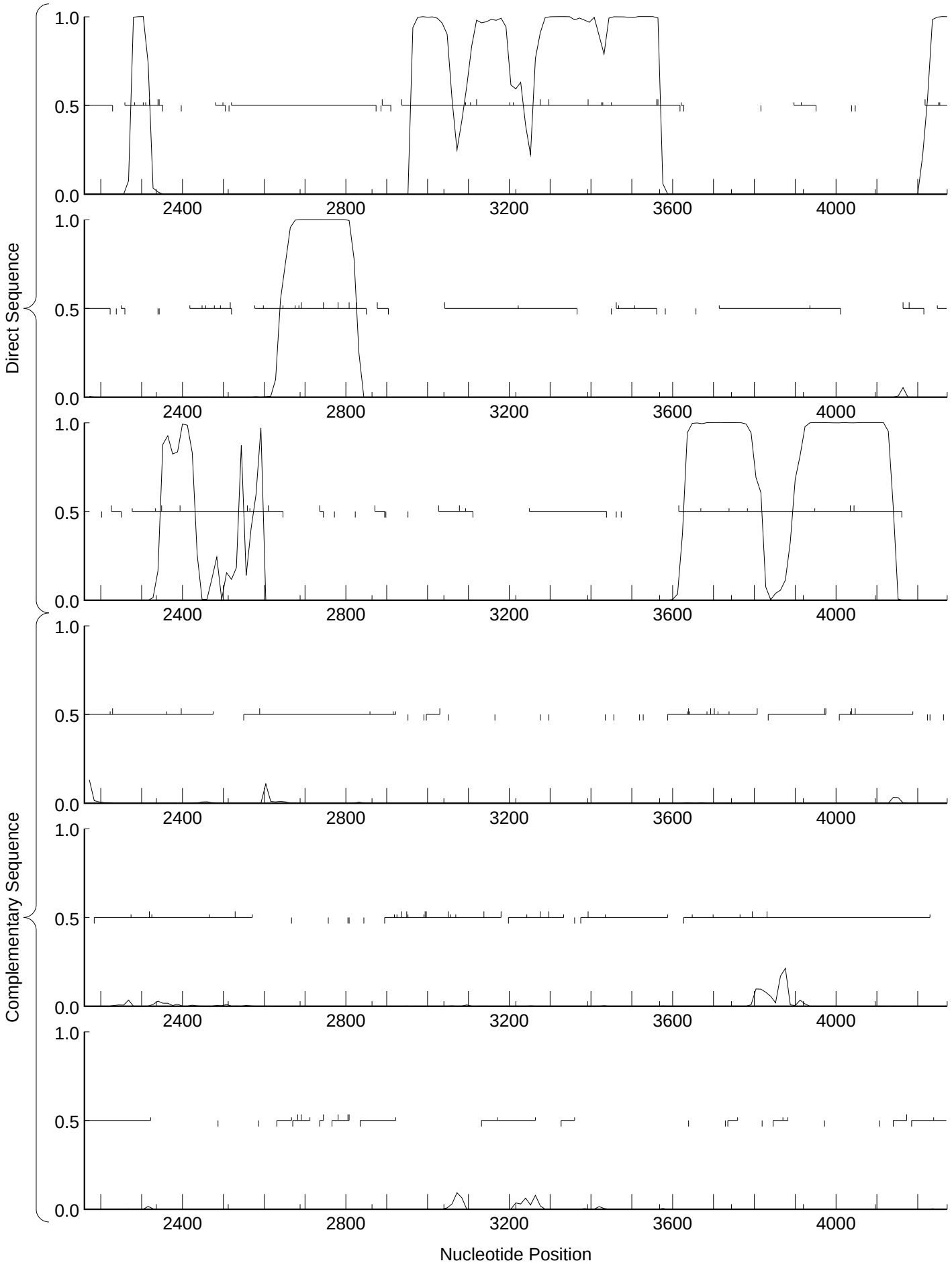

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

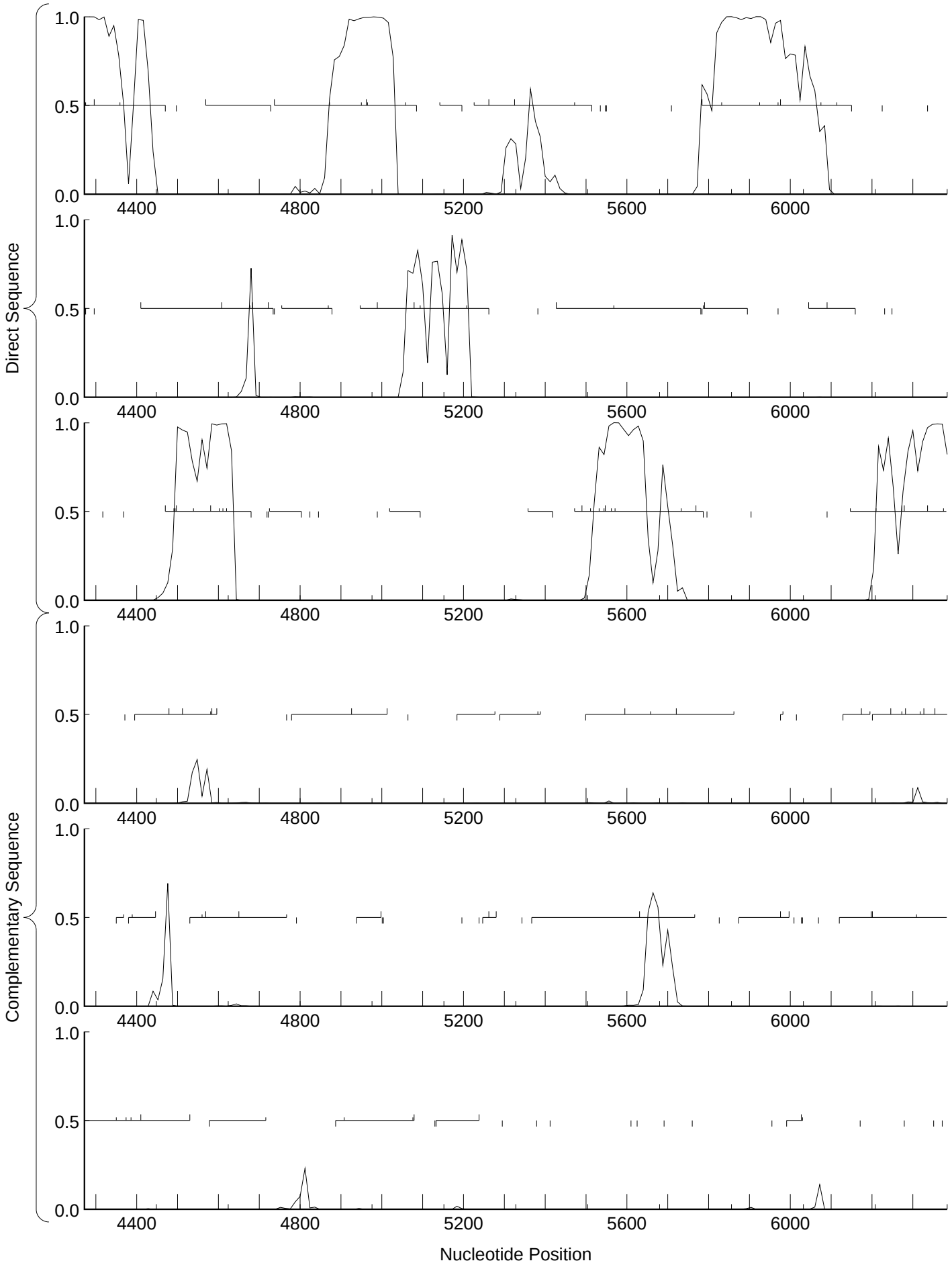
acterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 2/75



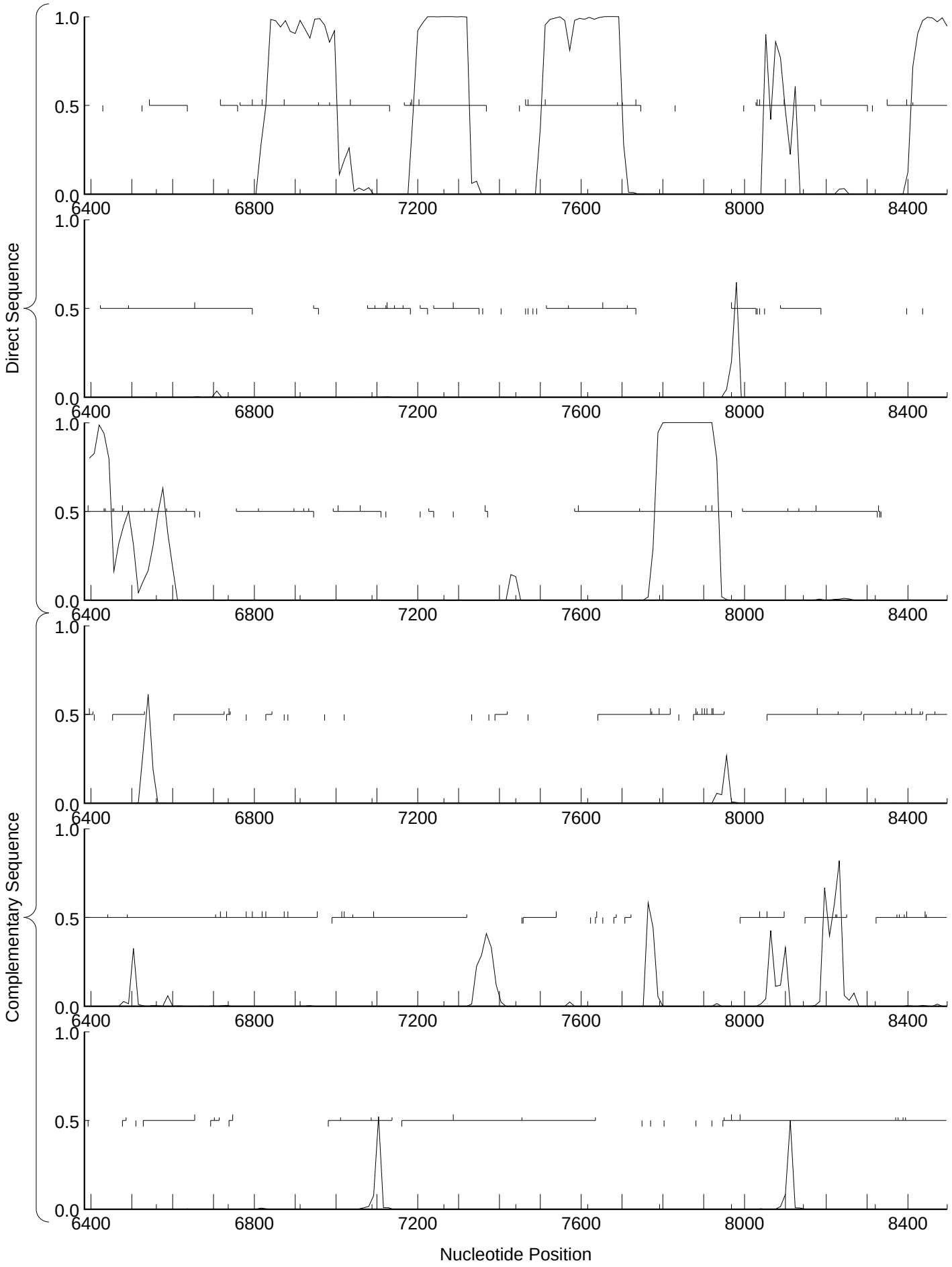
acterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 3/75



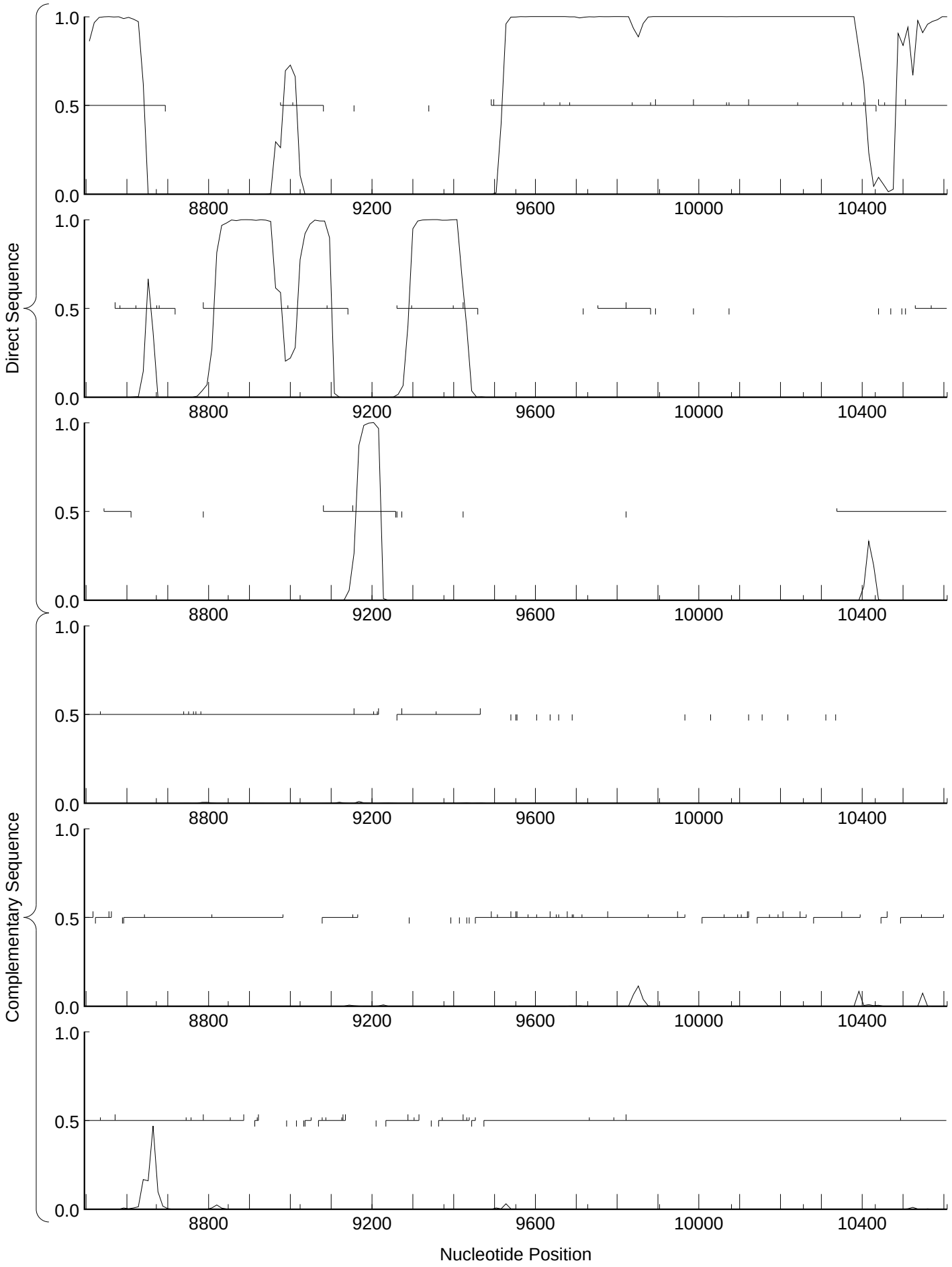
acterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 4/75



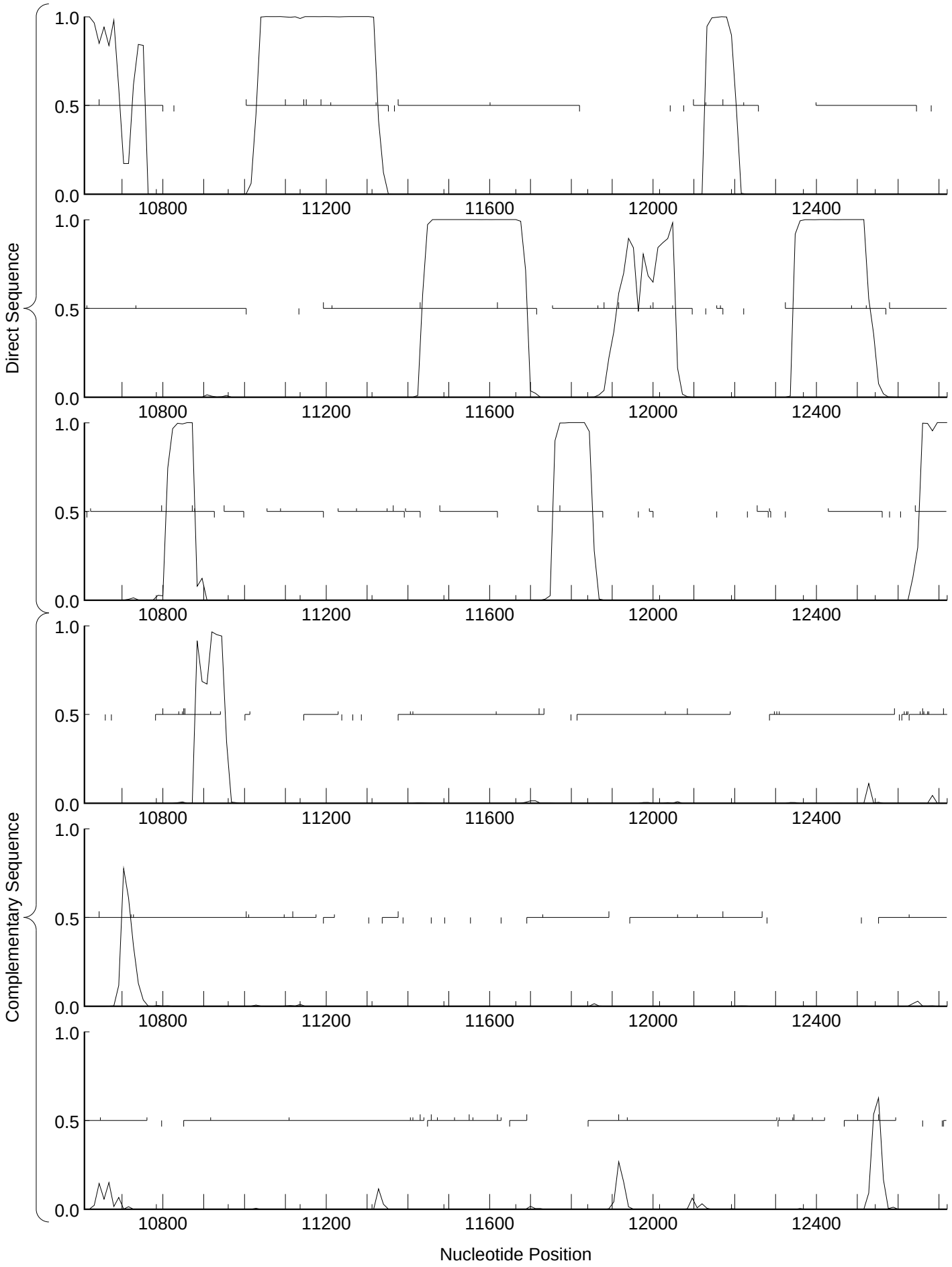
acterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 5/75



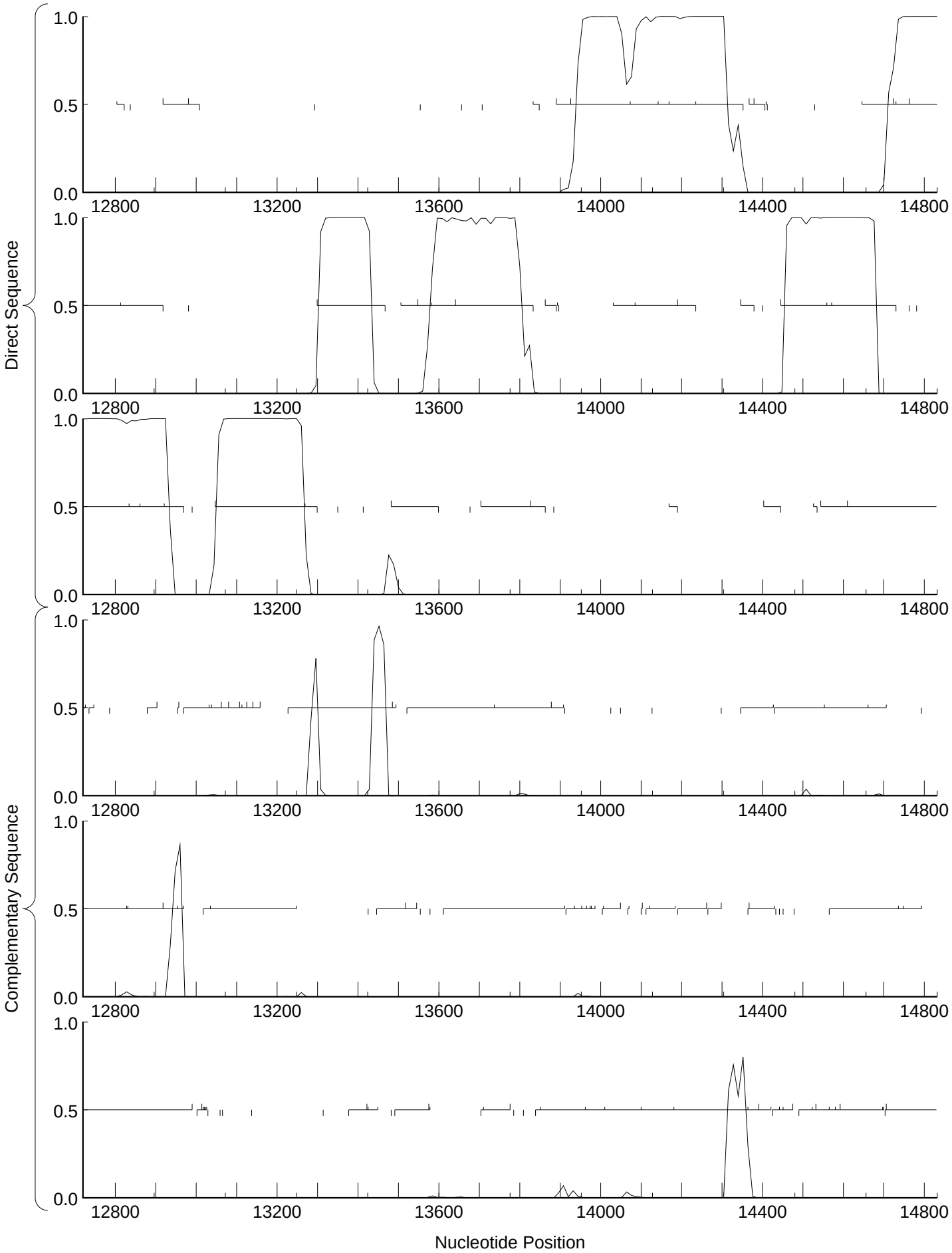
acterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 6/75



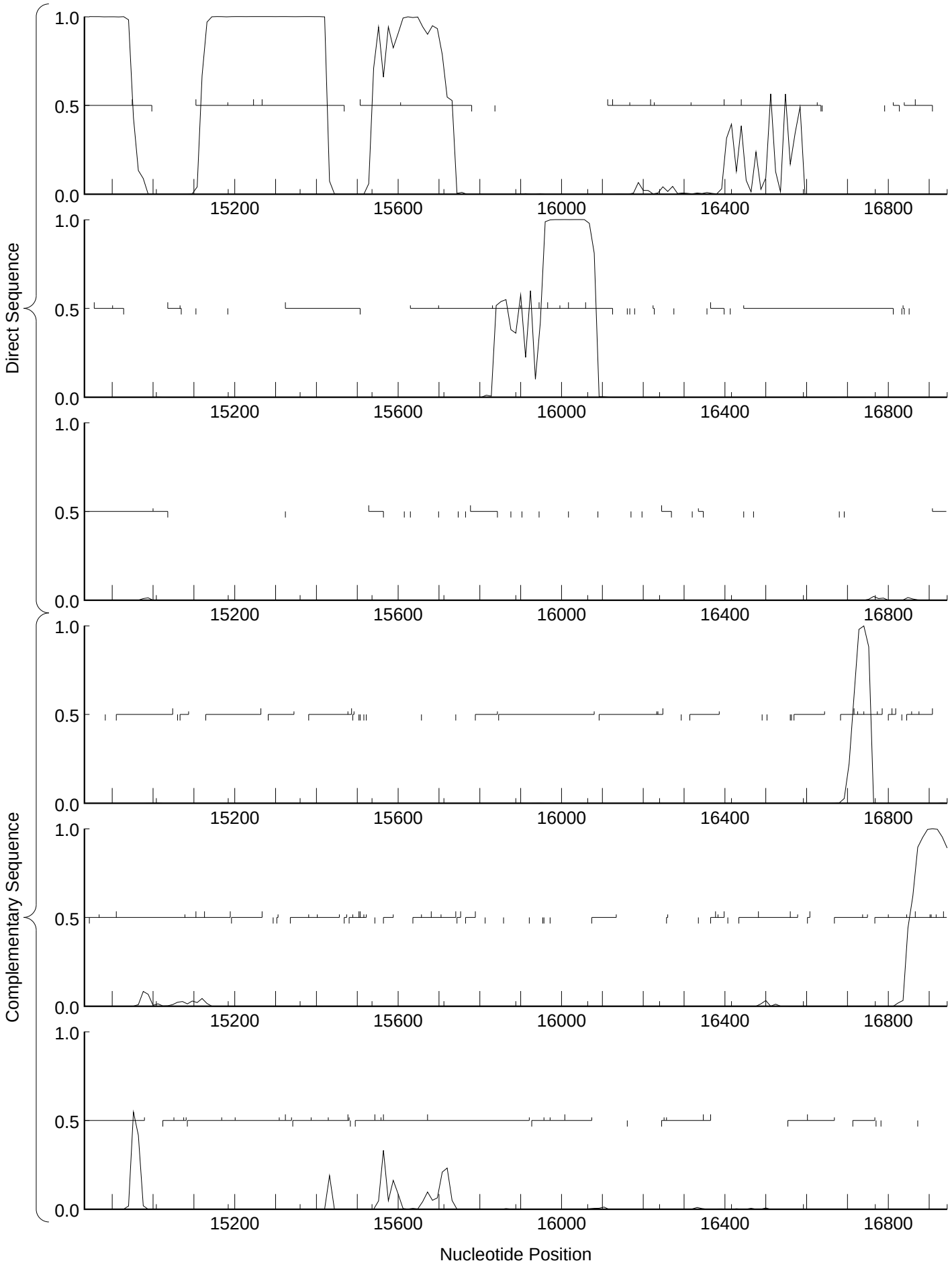
acterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 7/75



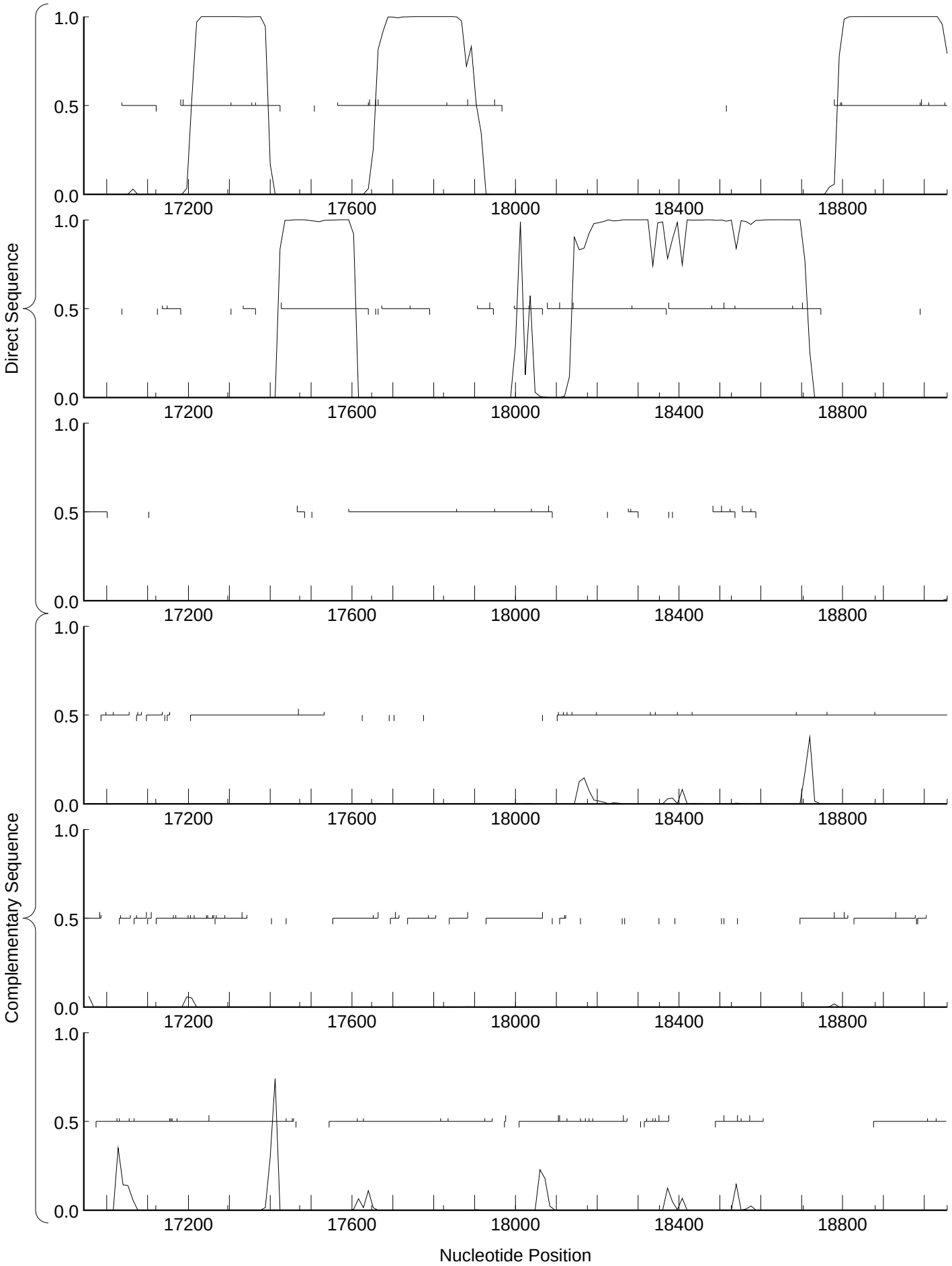
acterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 8/75



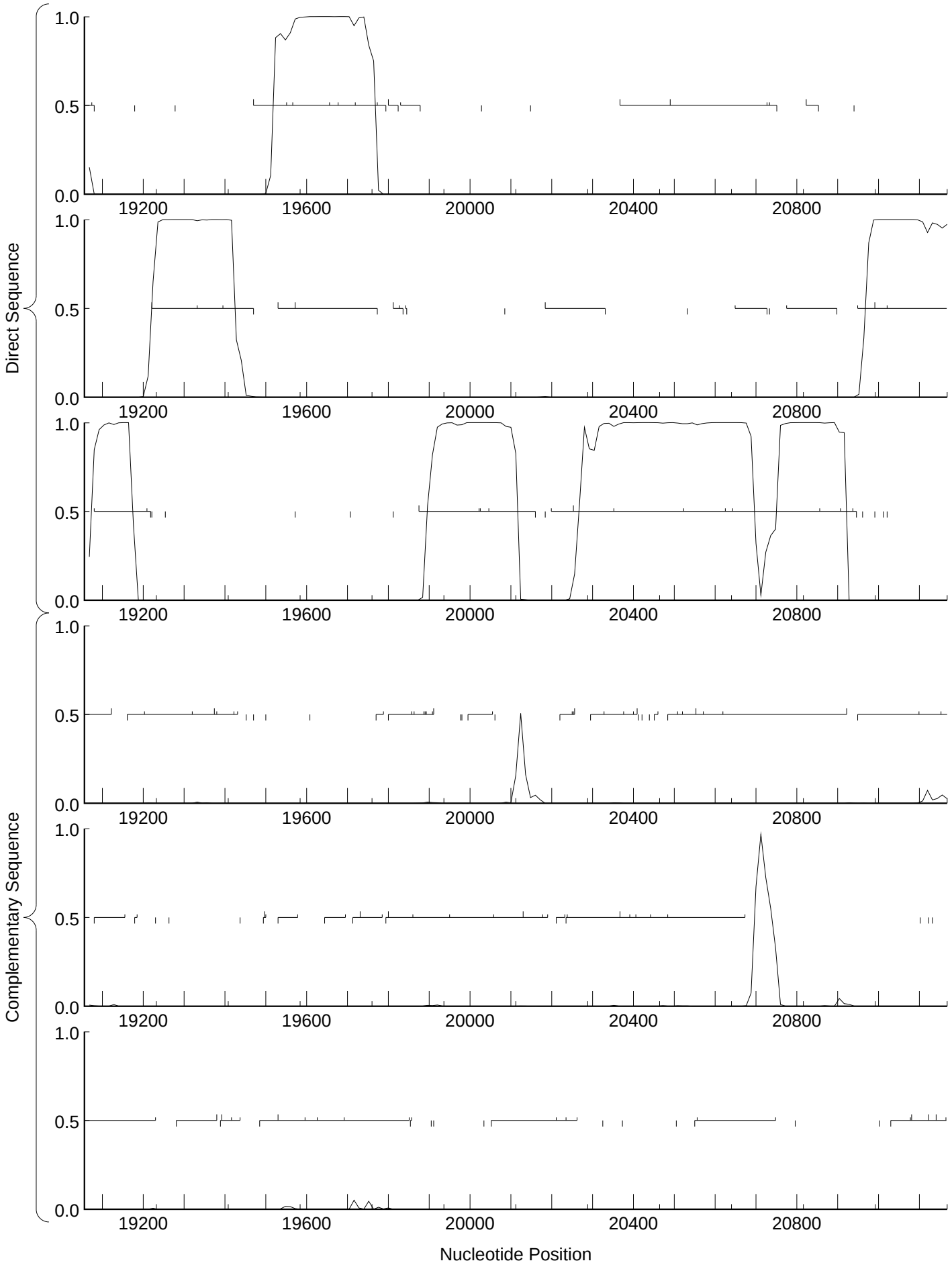
acterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 9/75



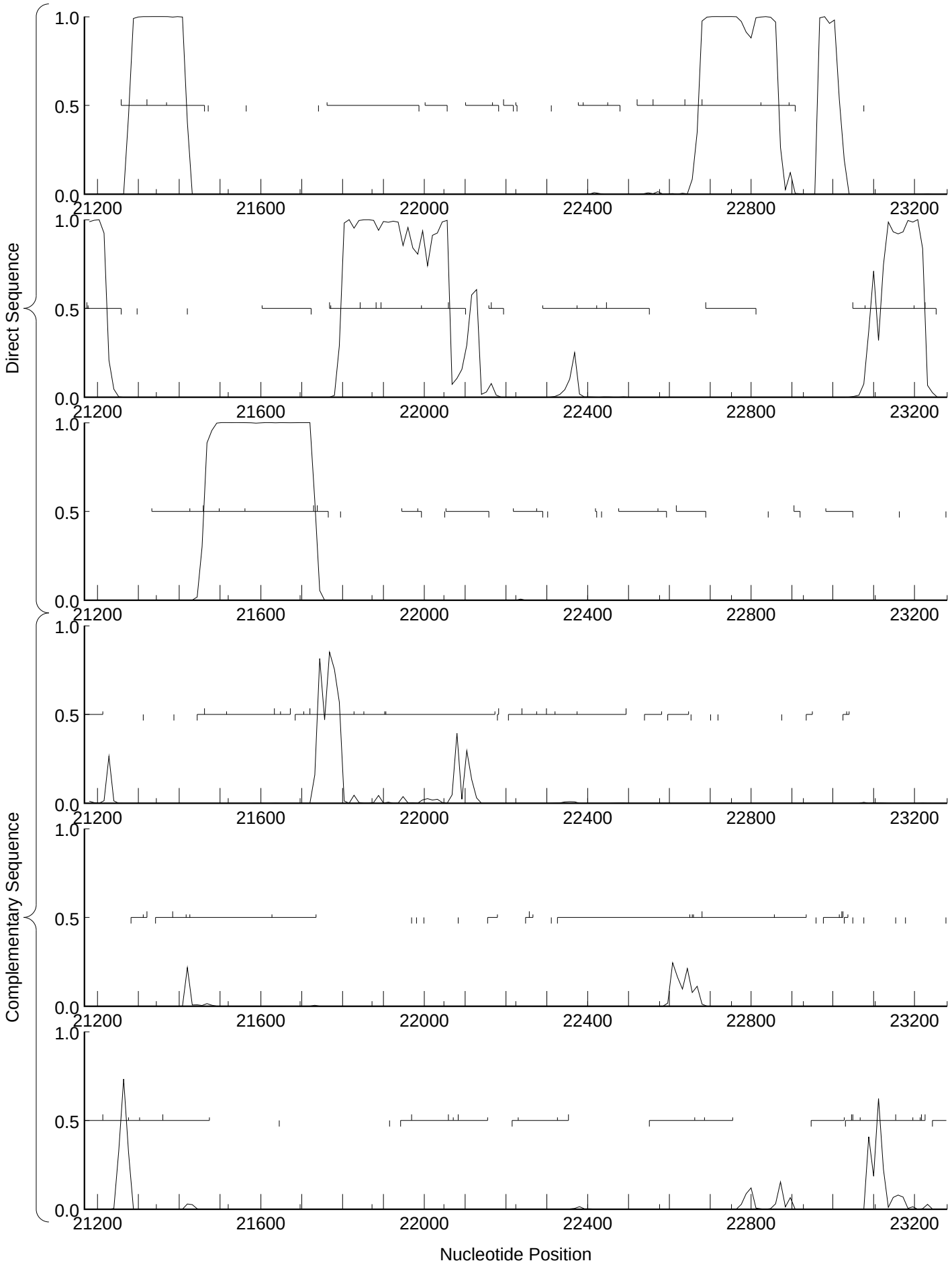
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 10/75



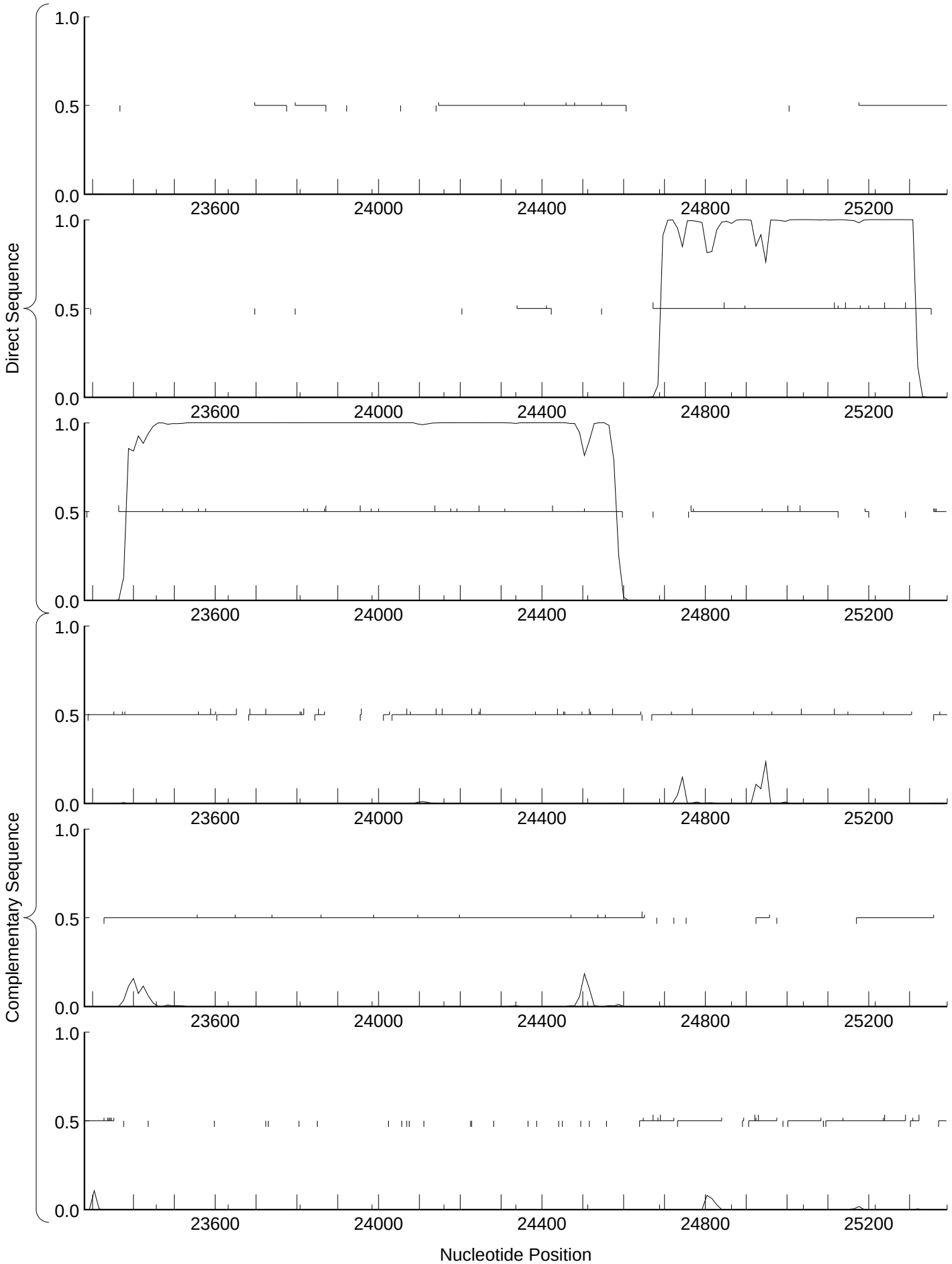
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 11/75



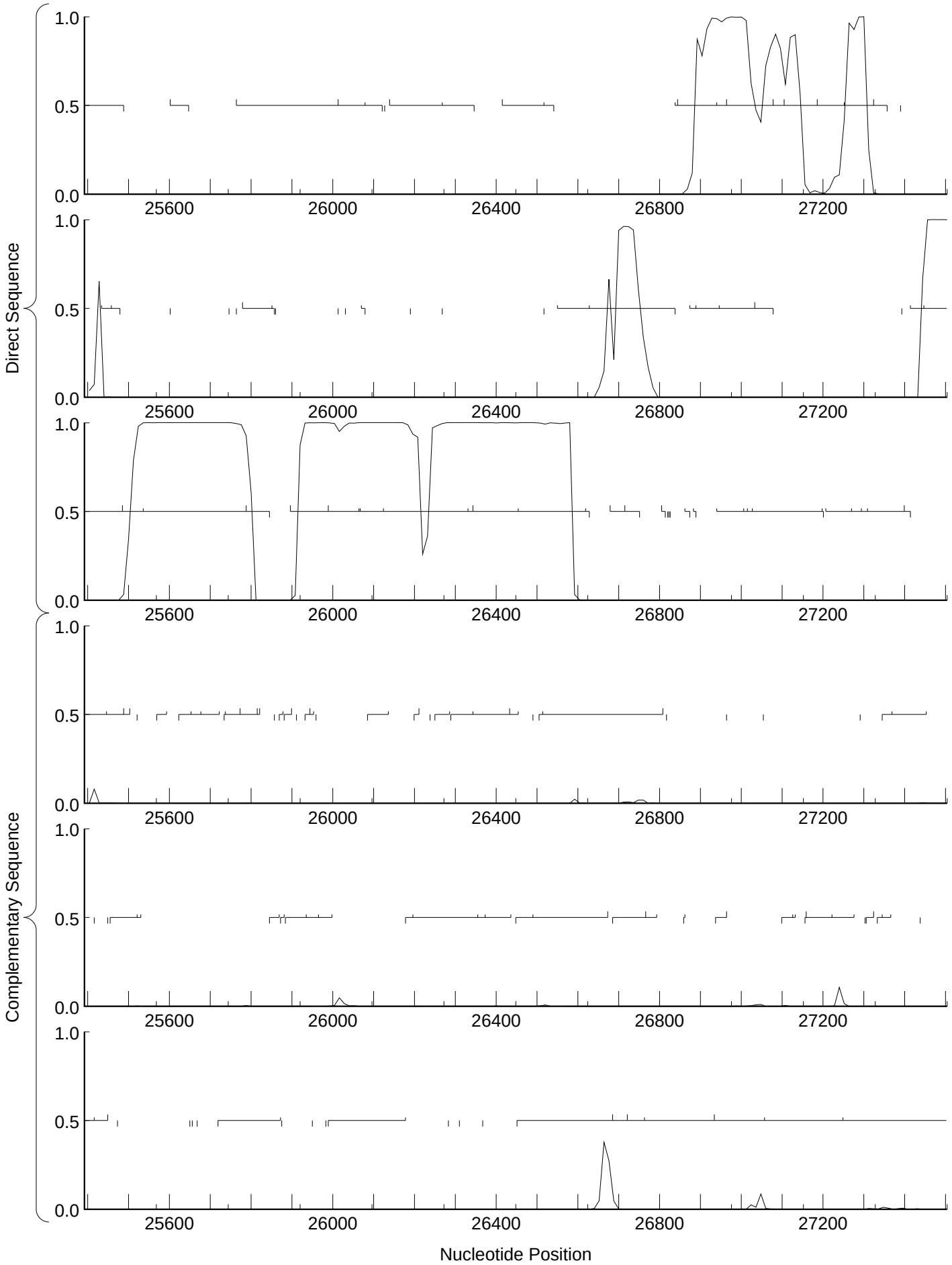
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 12/75



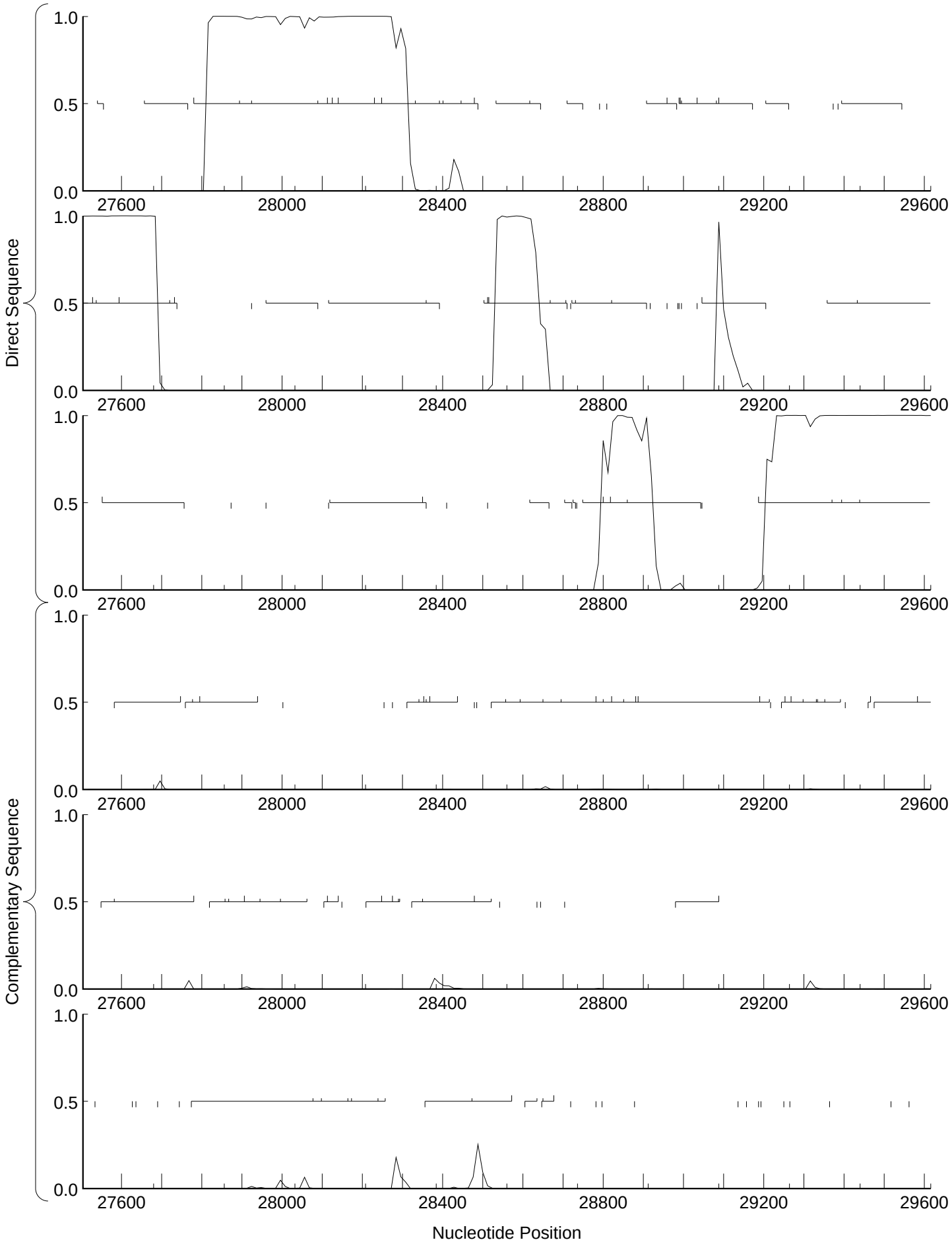
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 13/75



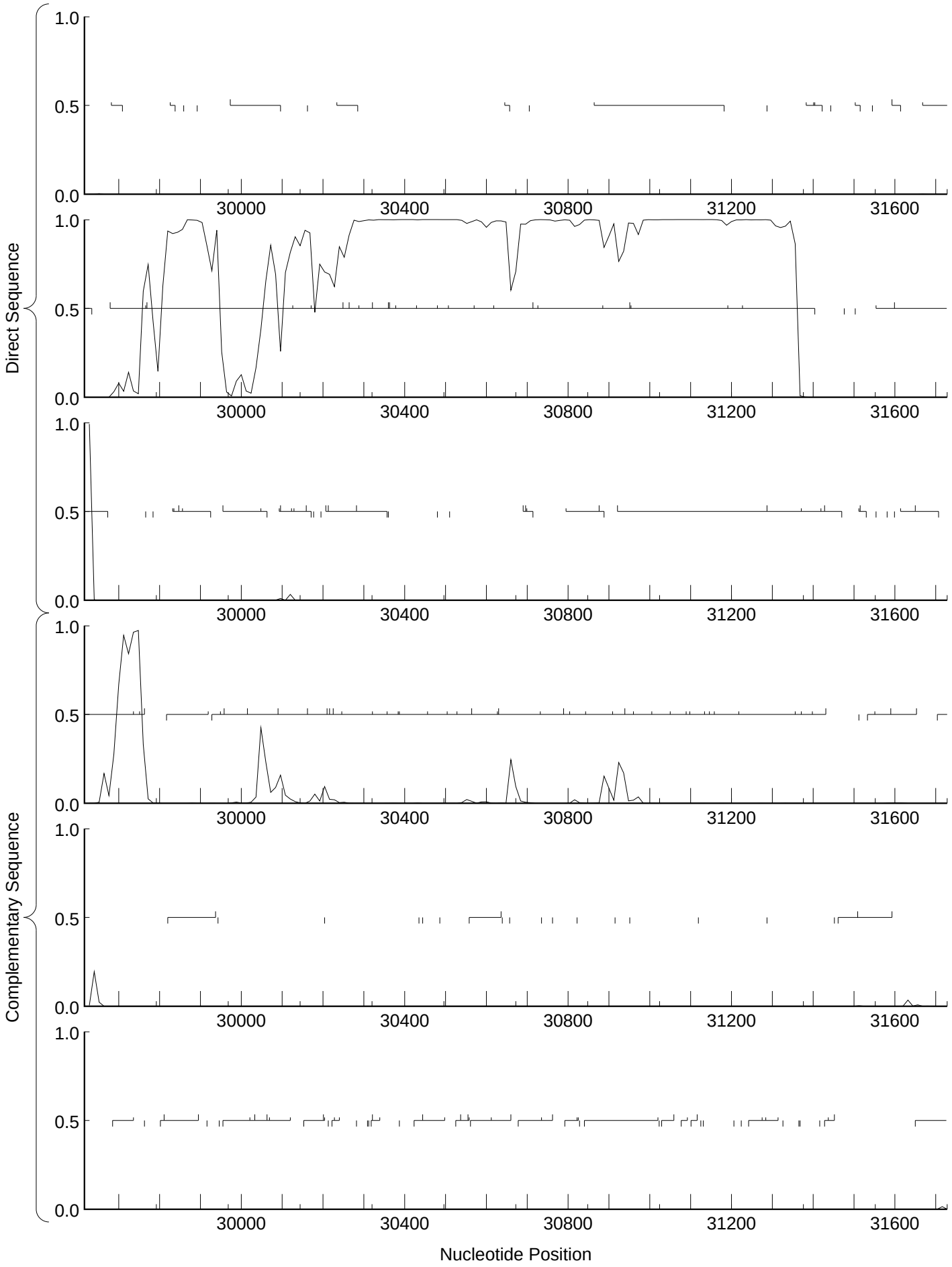
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 14/75



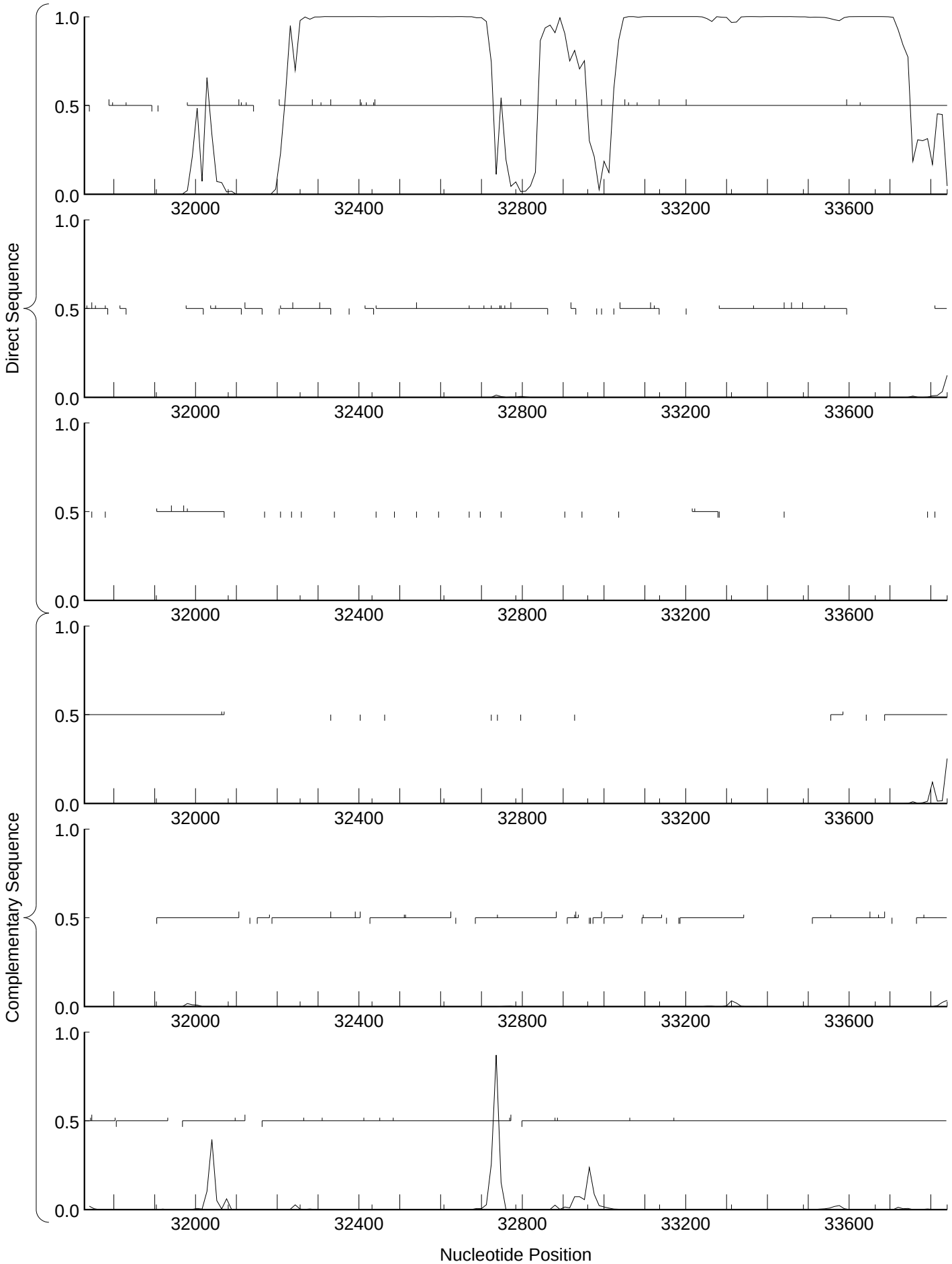
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 15/75



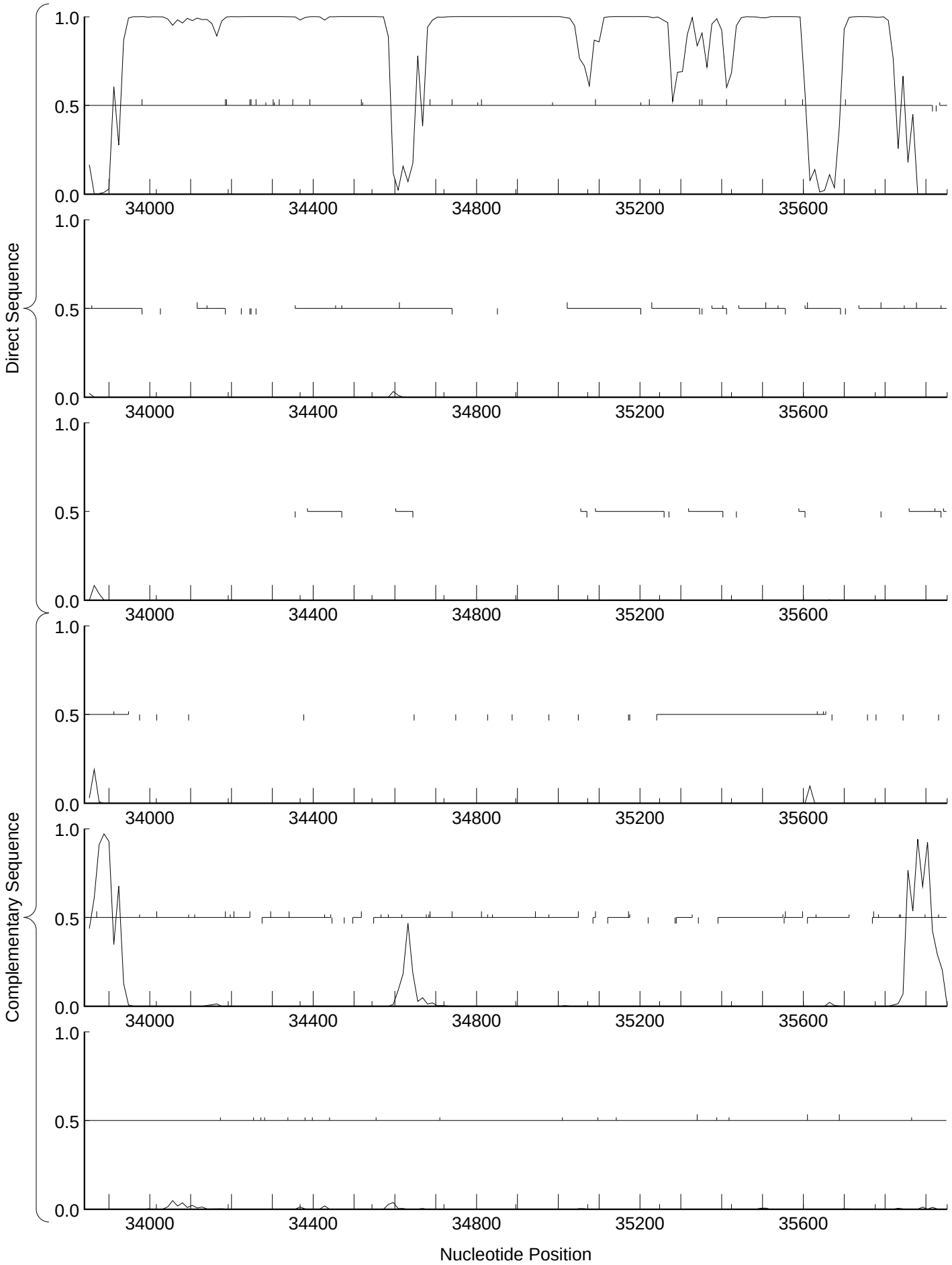
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 16/75



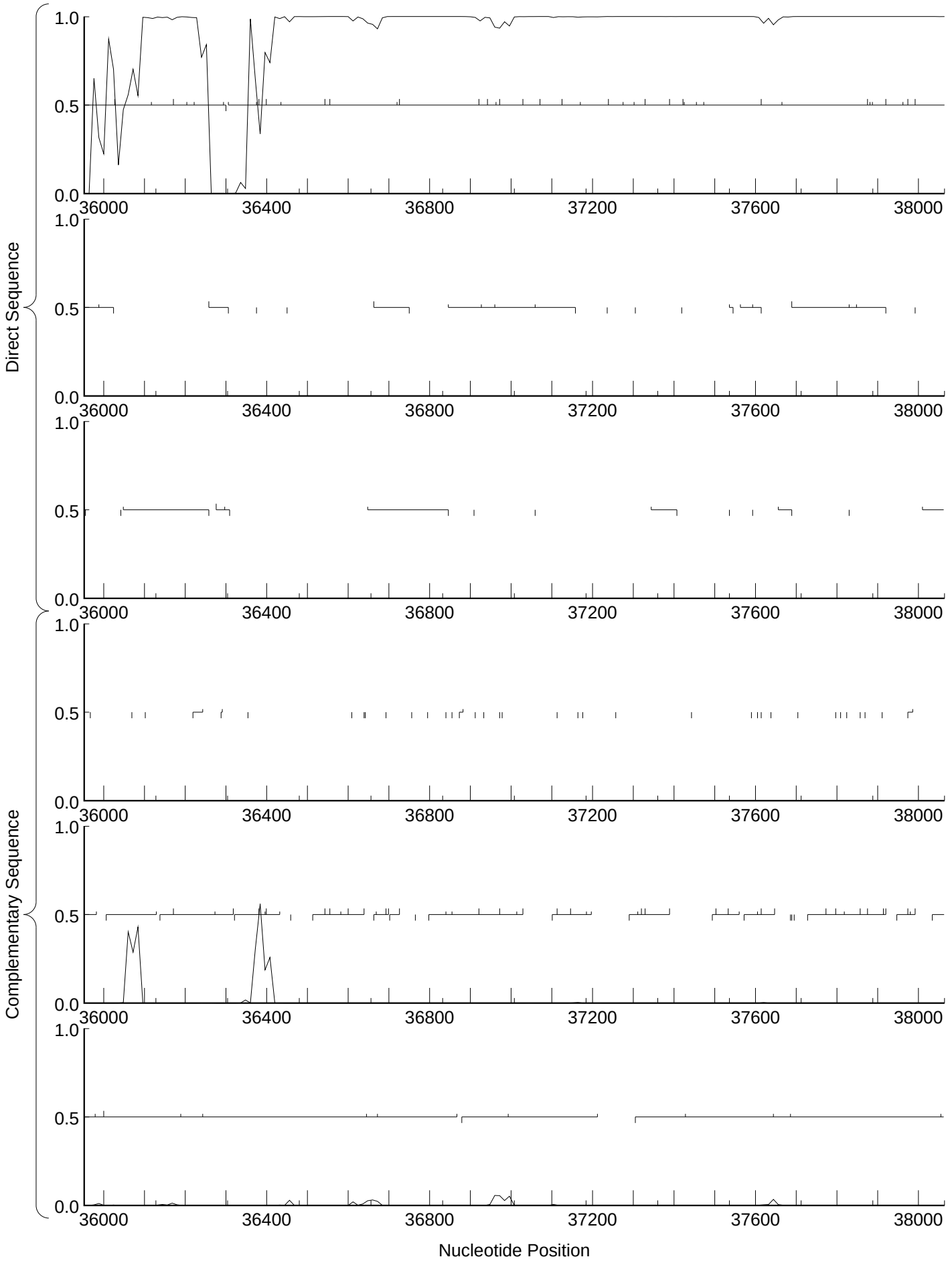
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 17/75



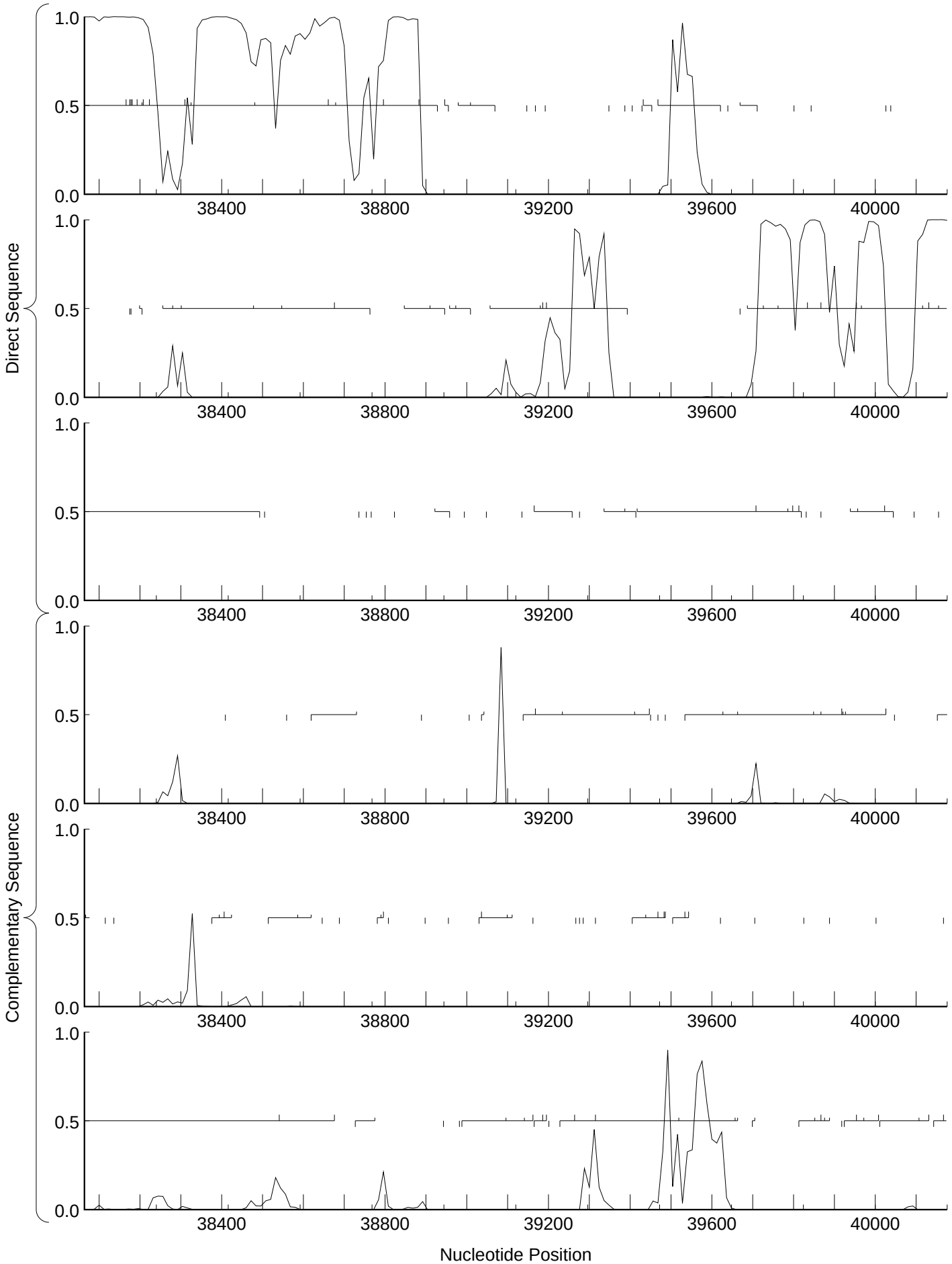
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 18/75



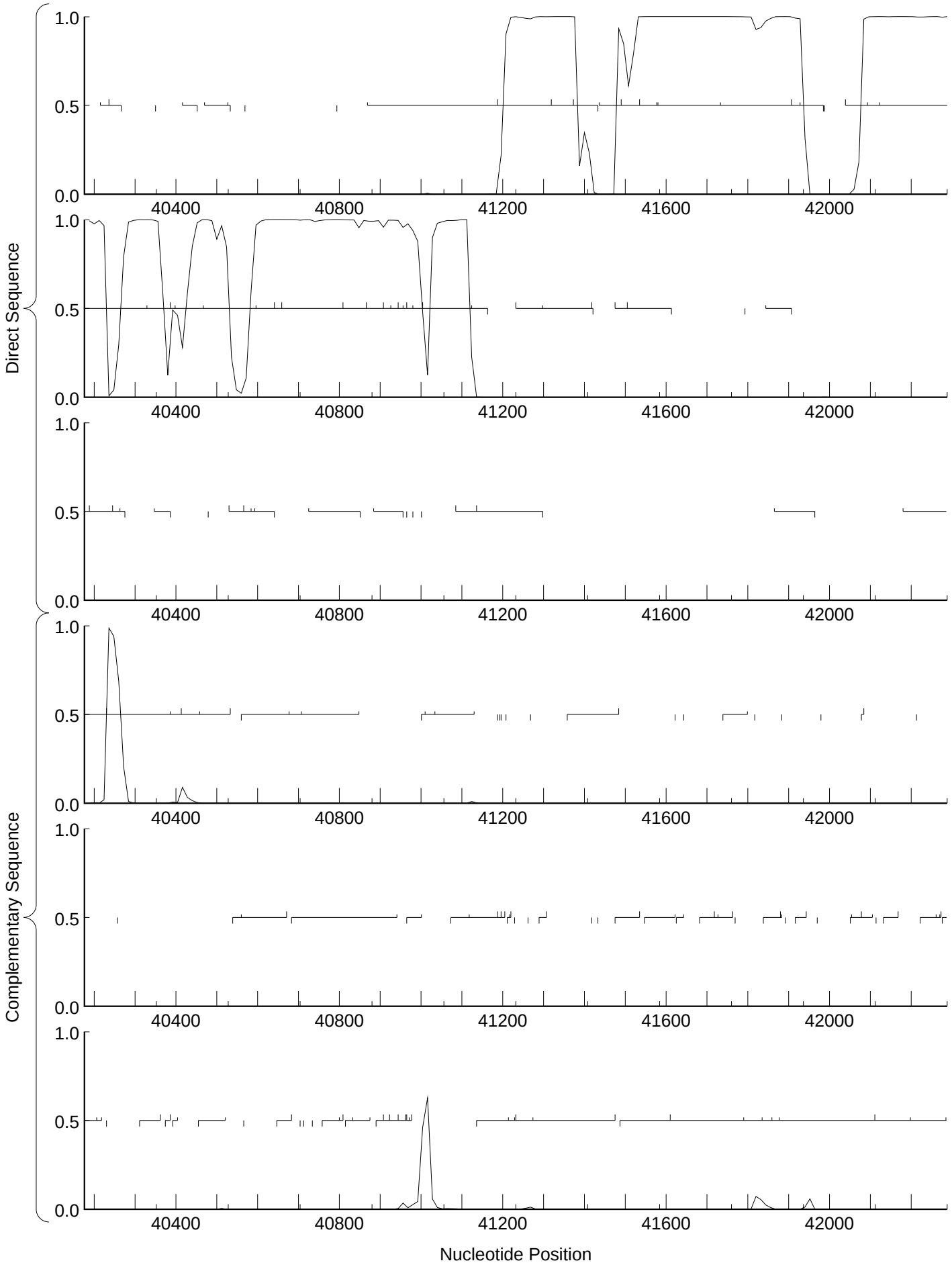
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 19/75



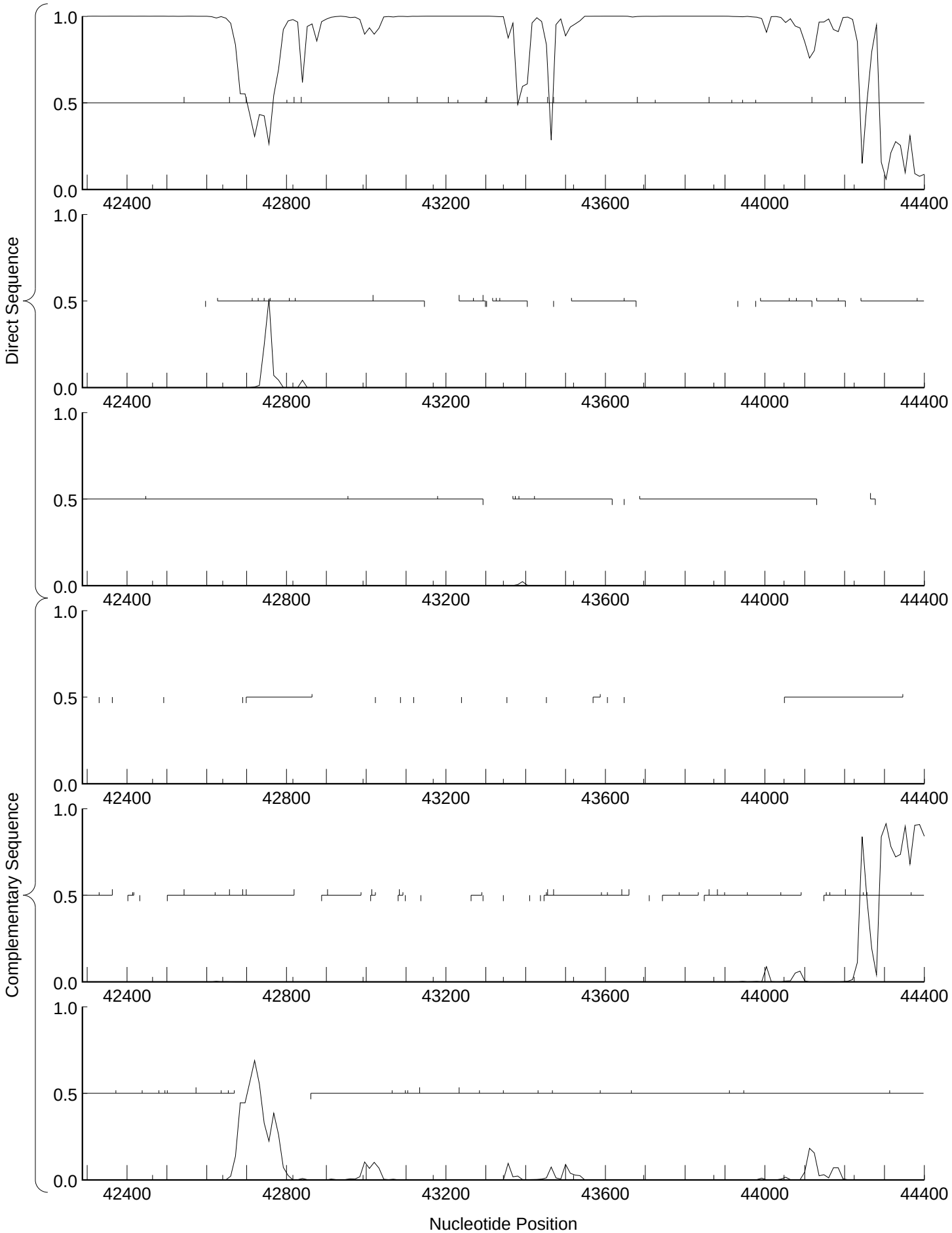
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 20/75



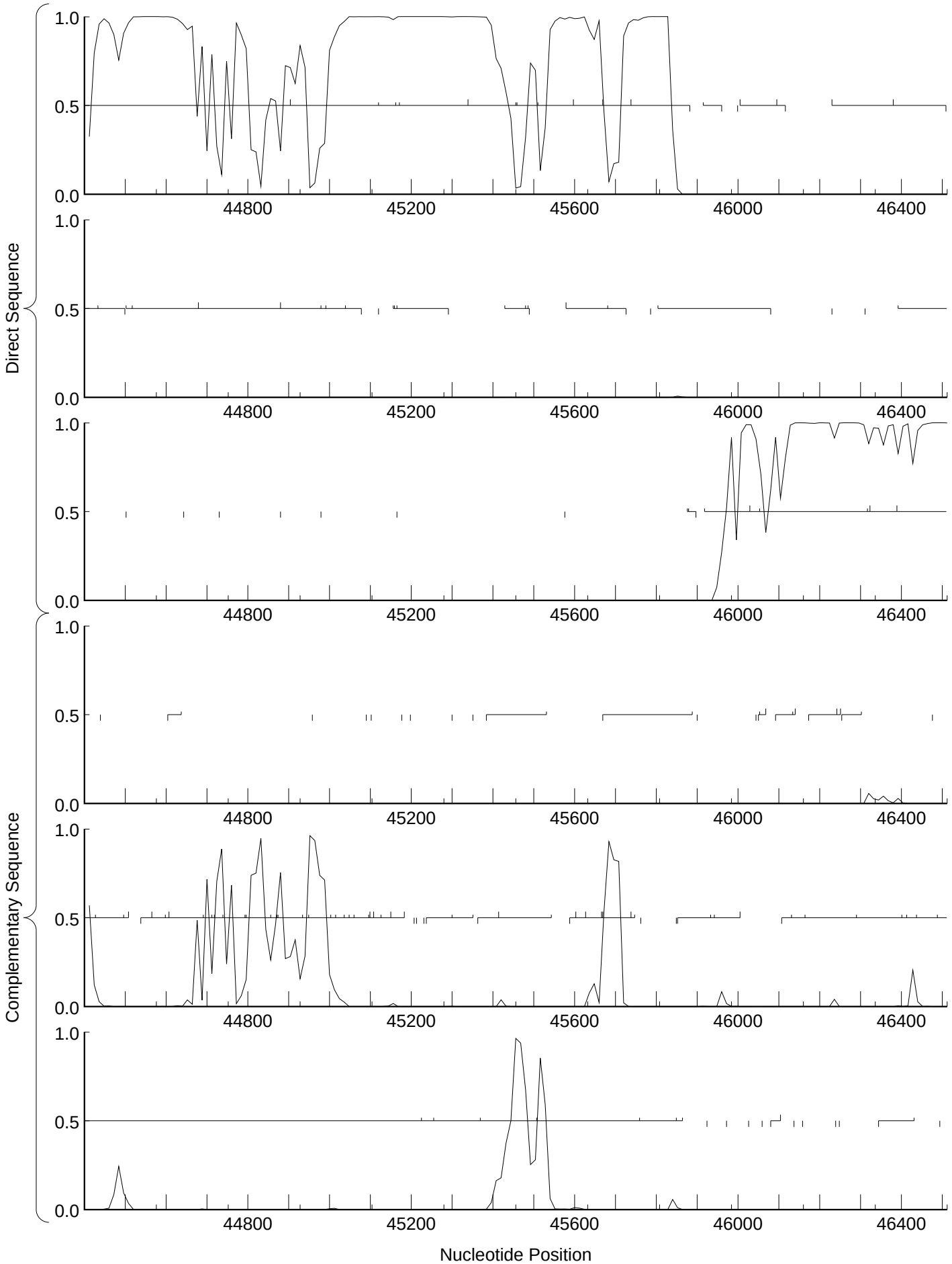
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 21/75



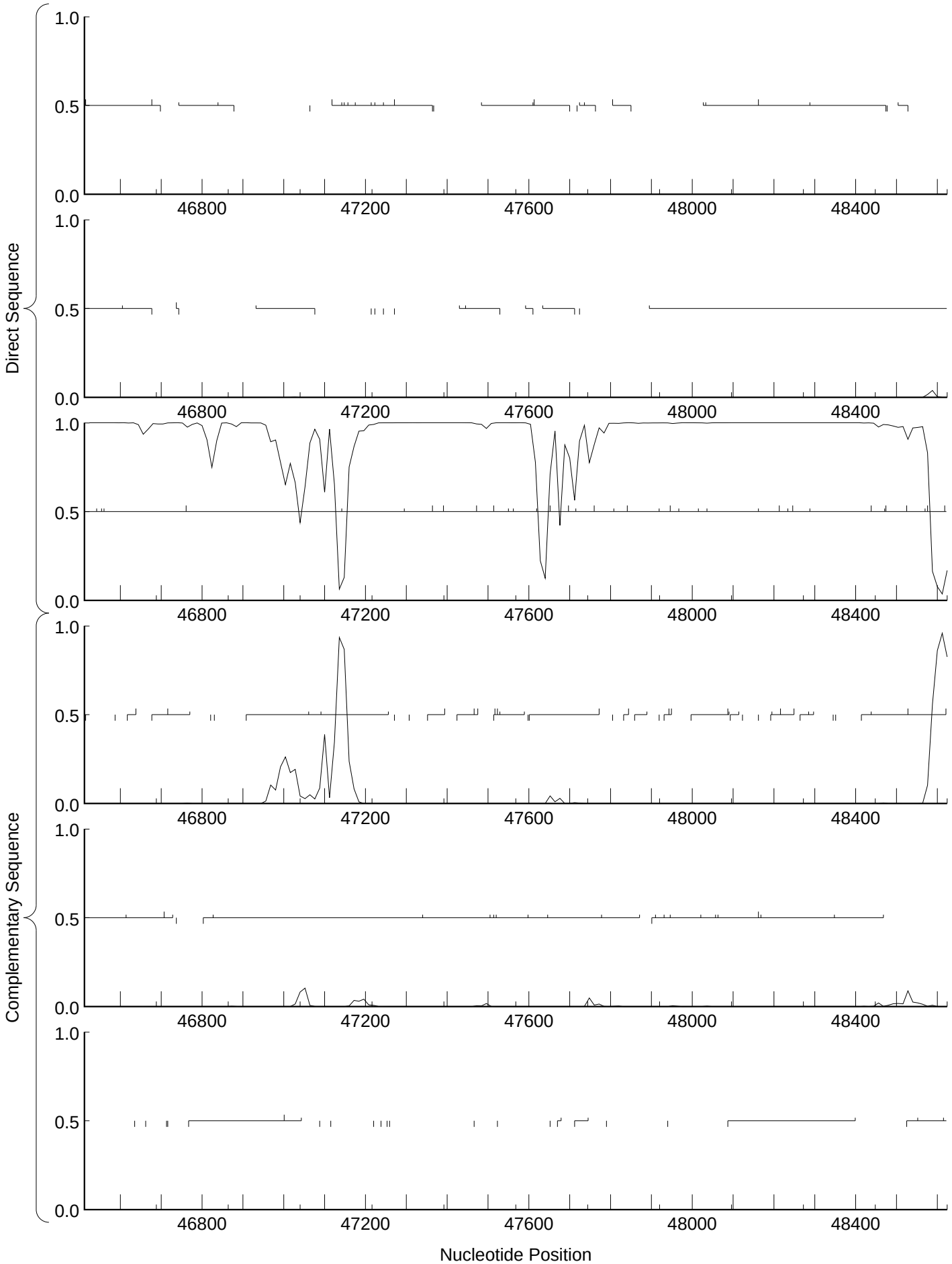
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 22/75



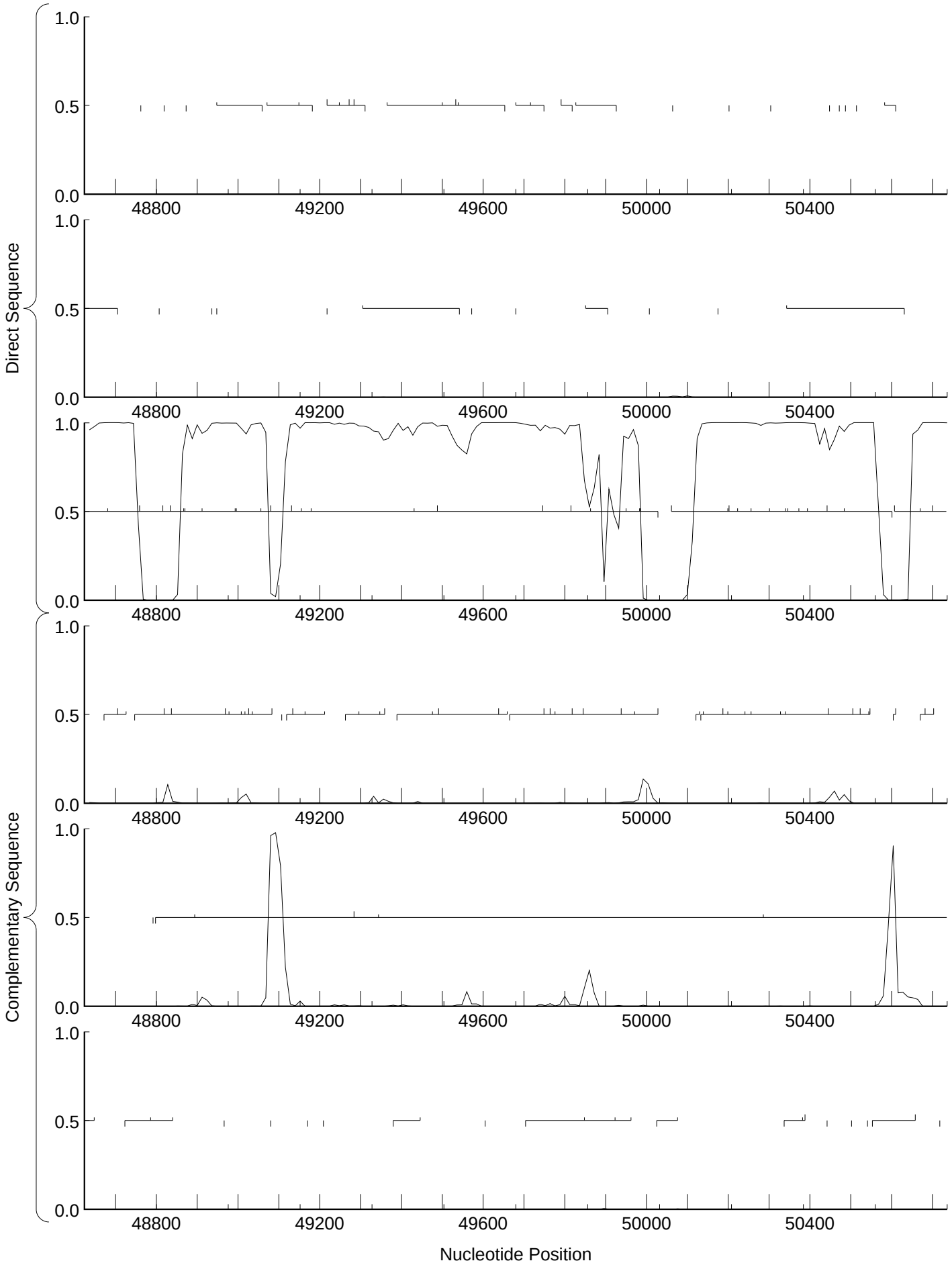
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 23/75



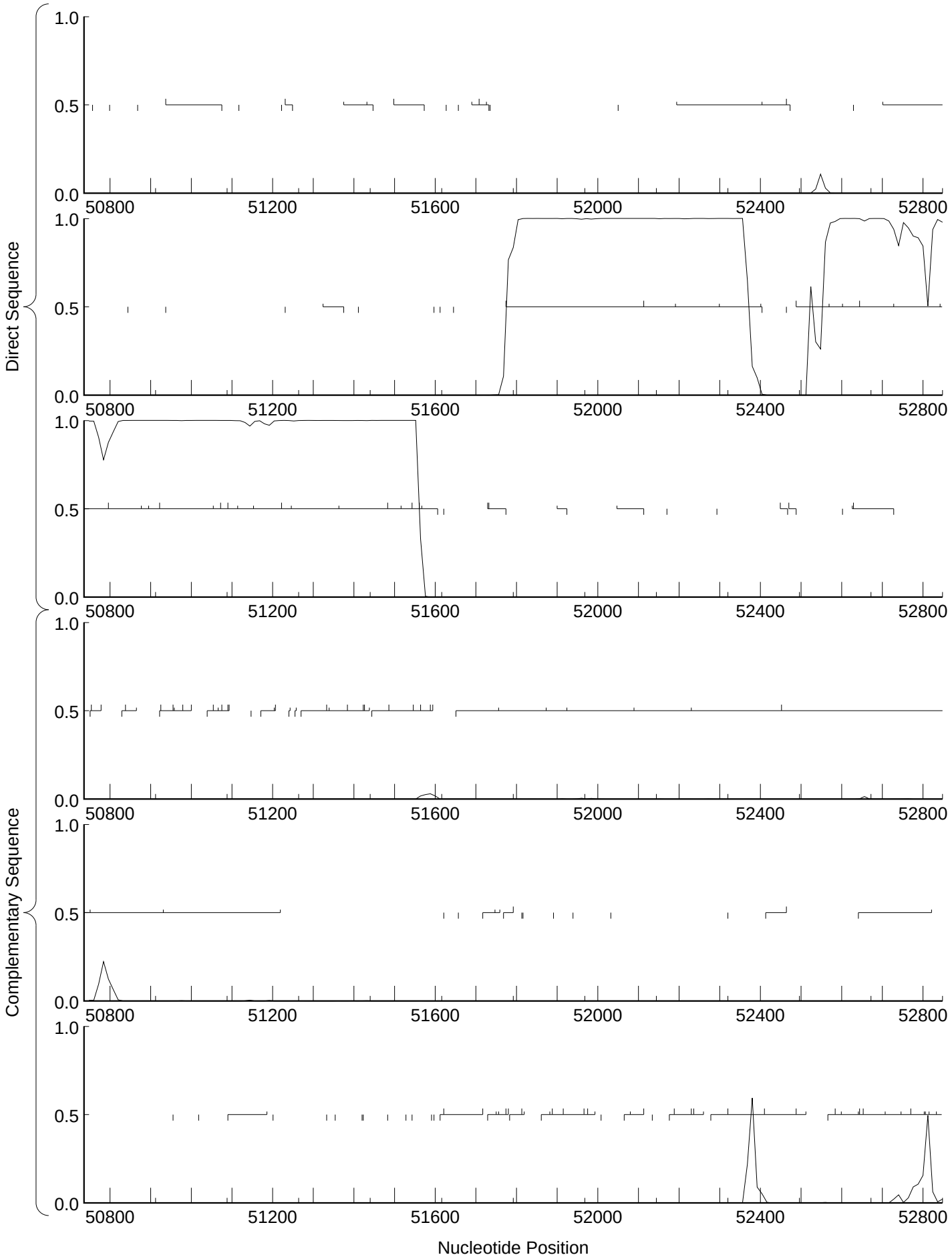
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 24/75



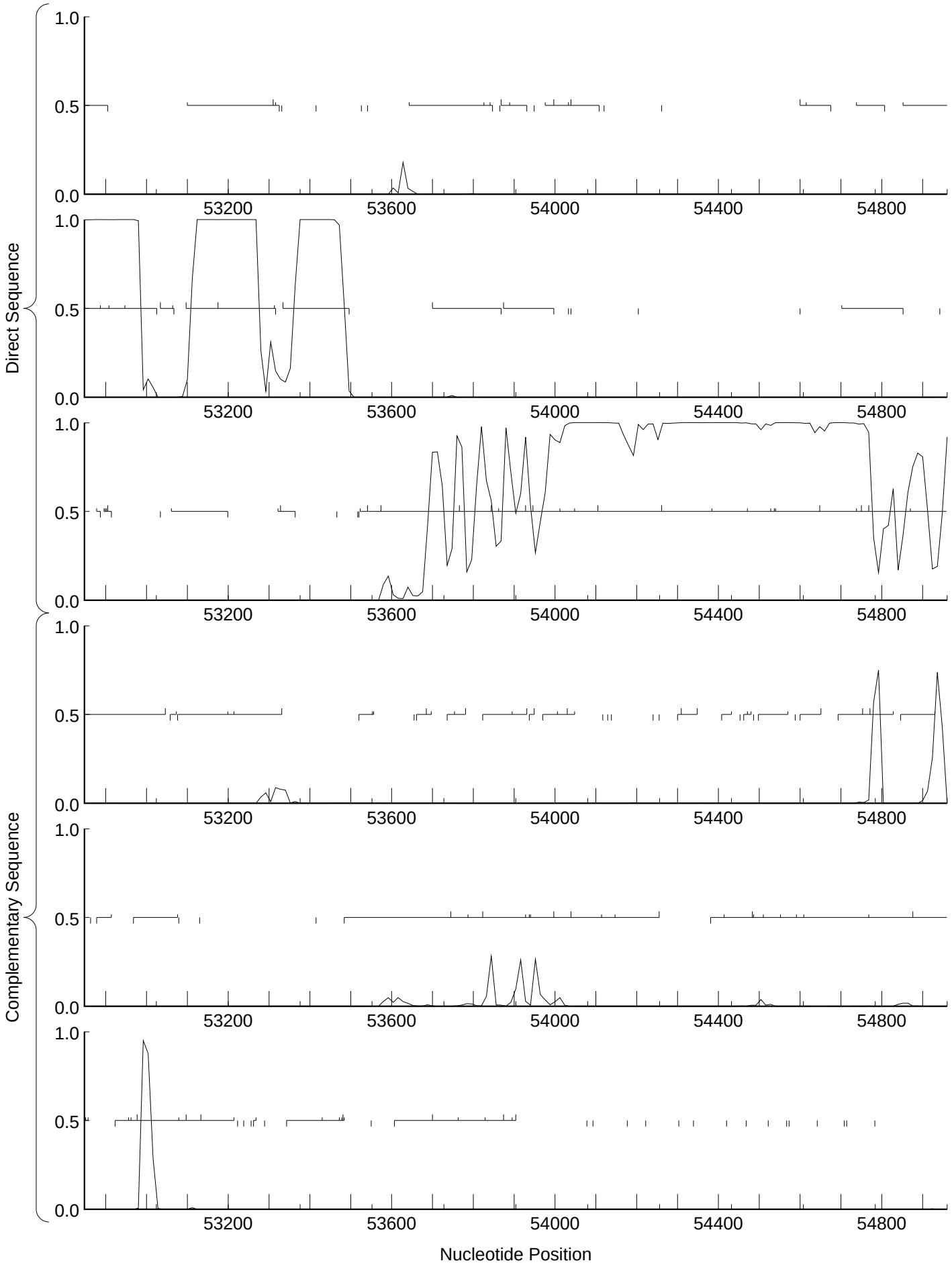
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 25/75



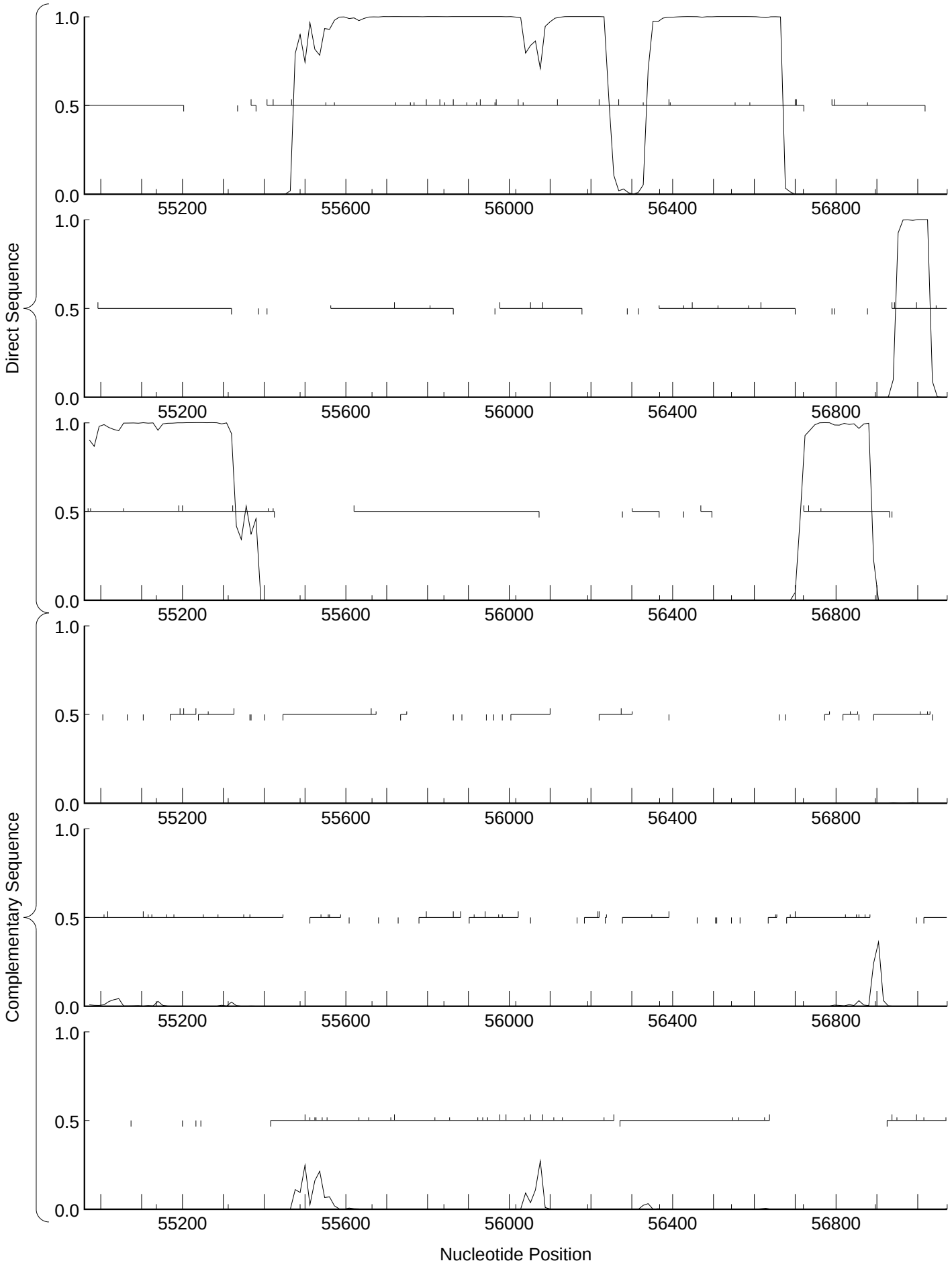
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 26/75



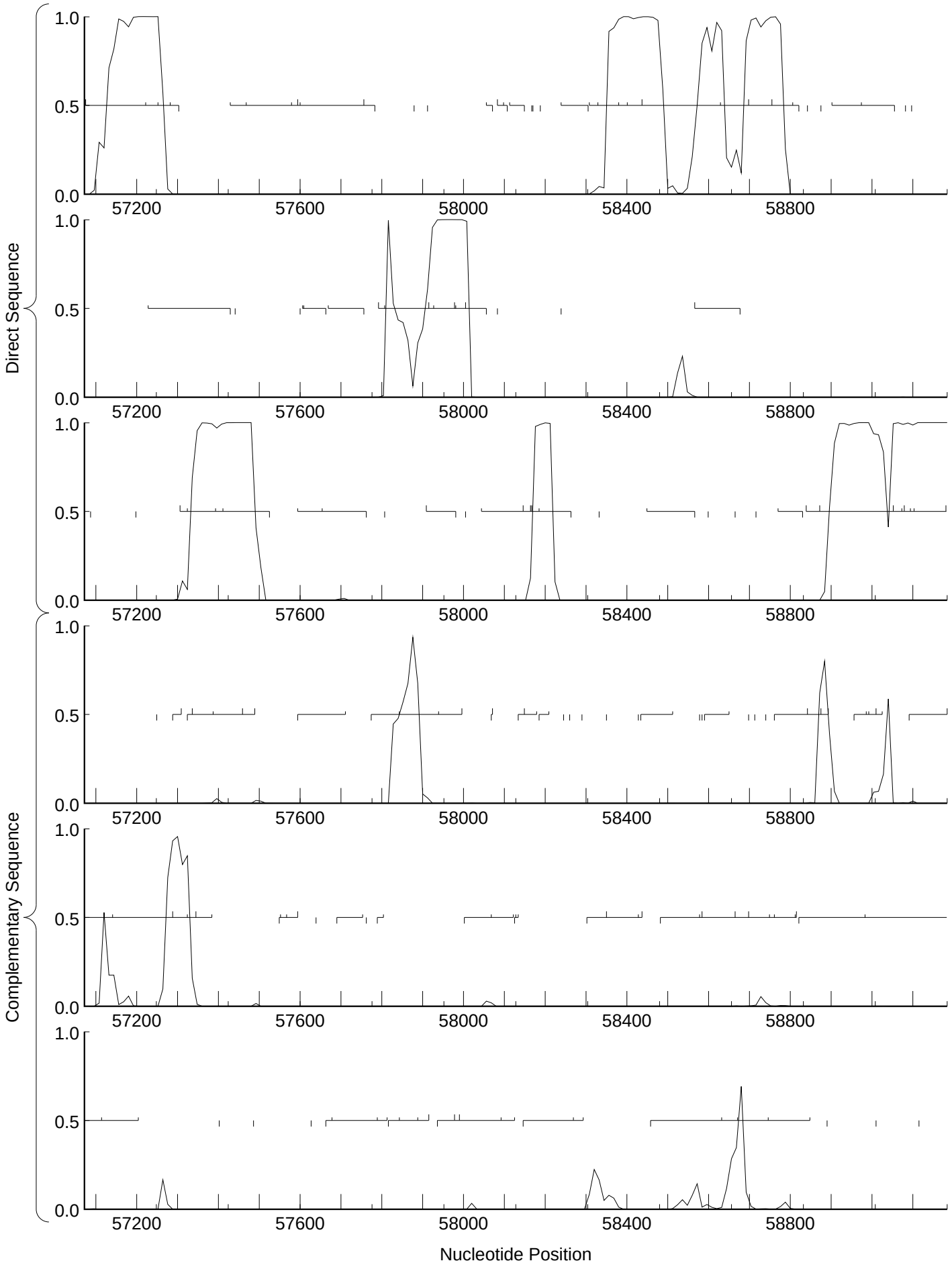
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 27/75



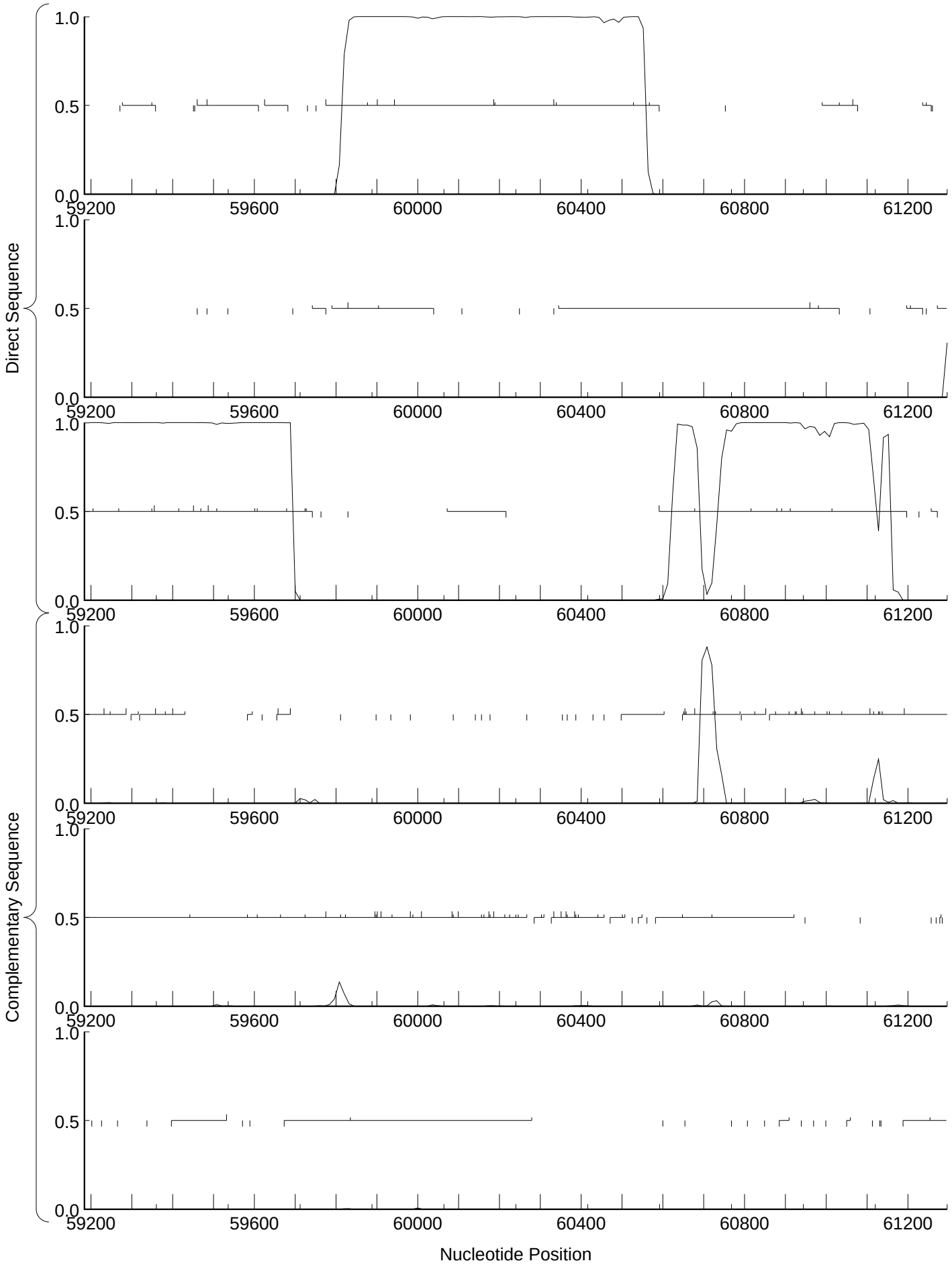
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 28/75



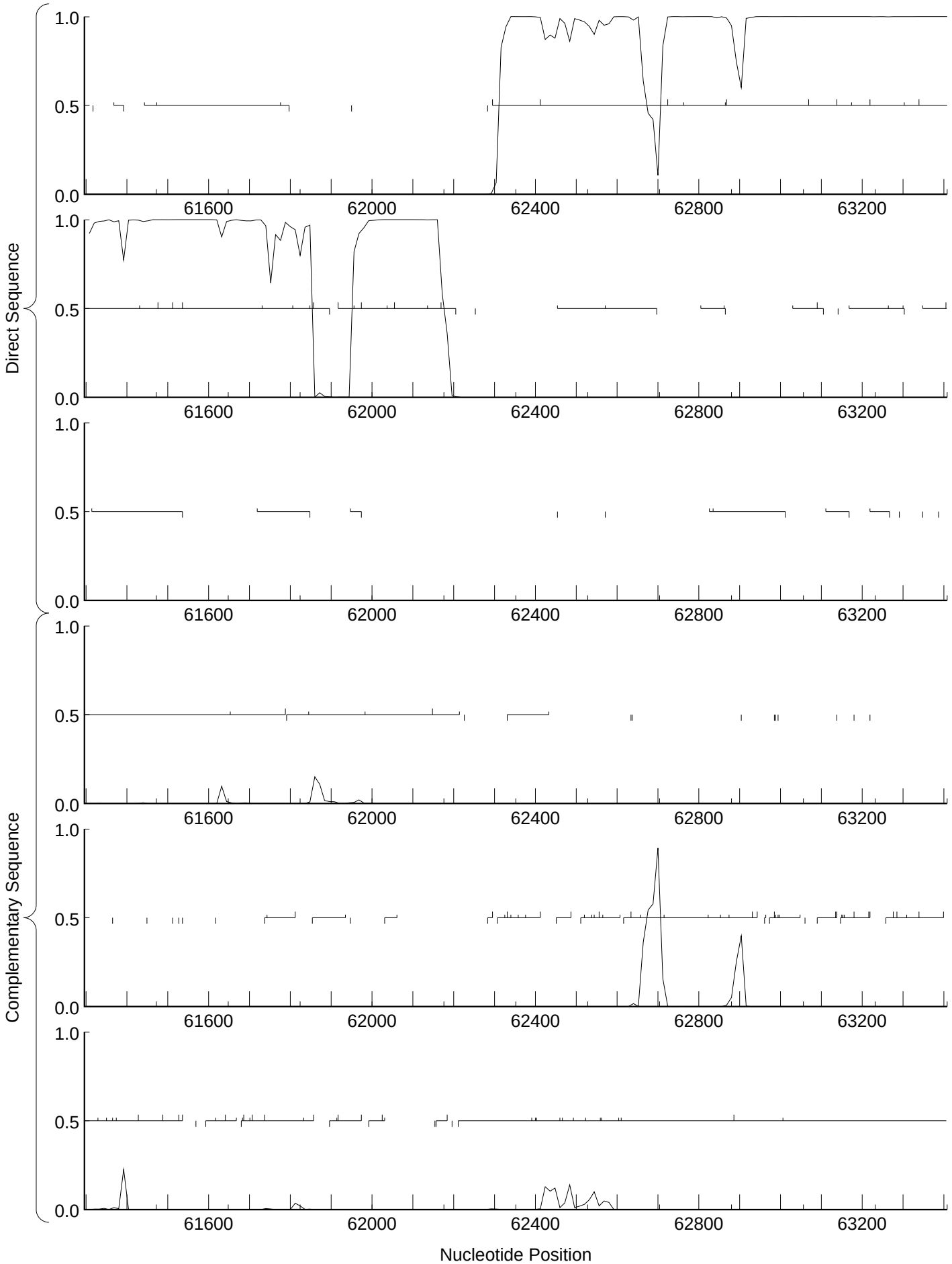
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 29/75



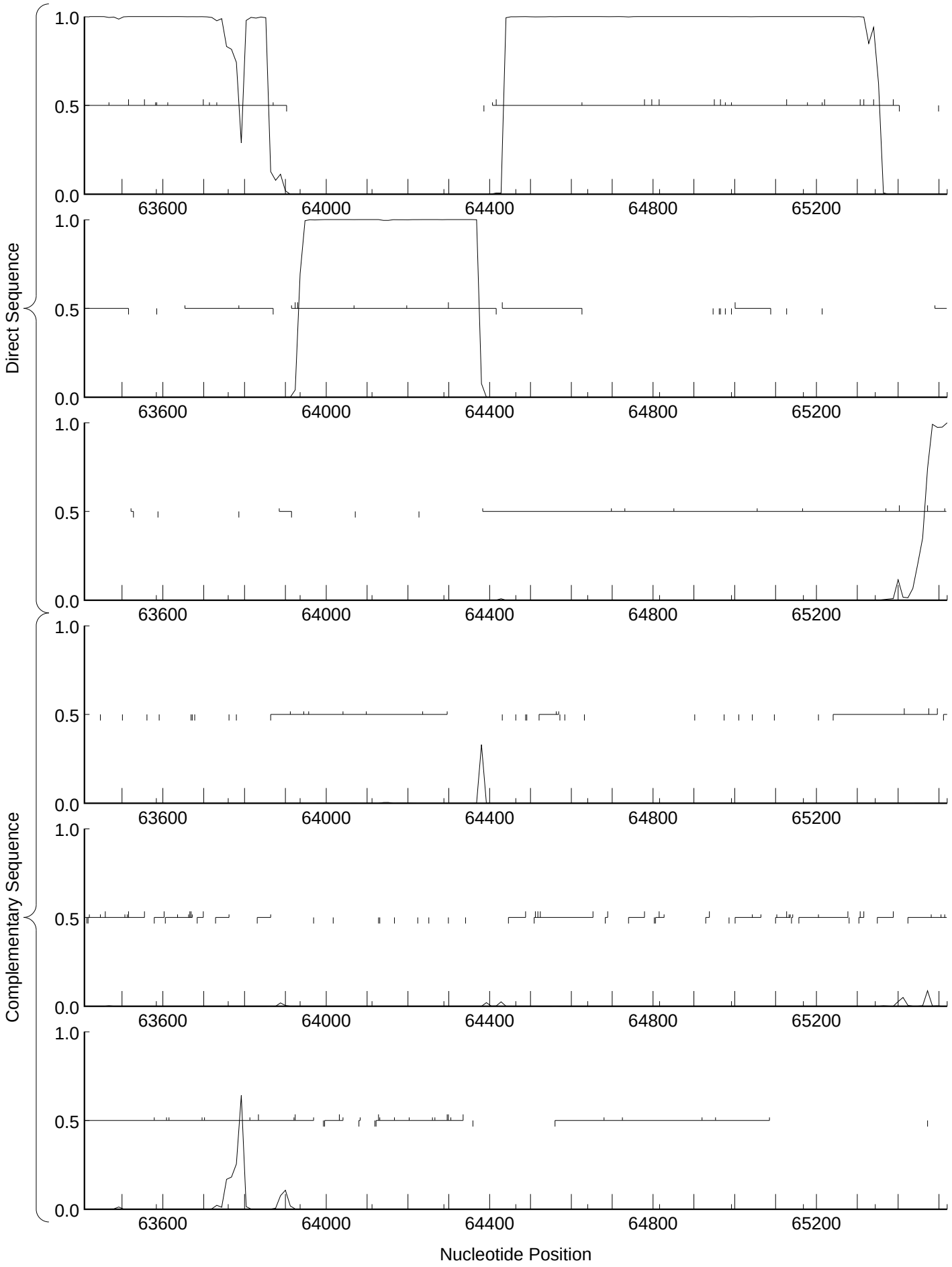
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 30/75



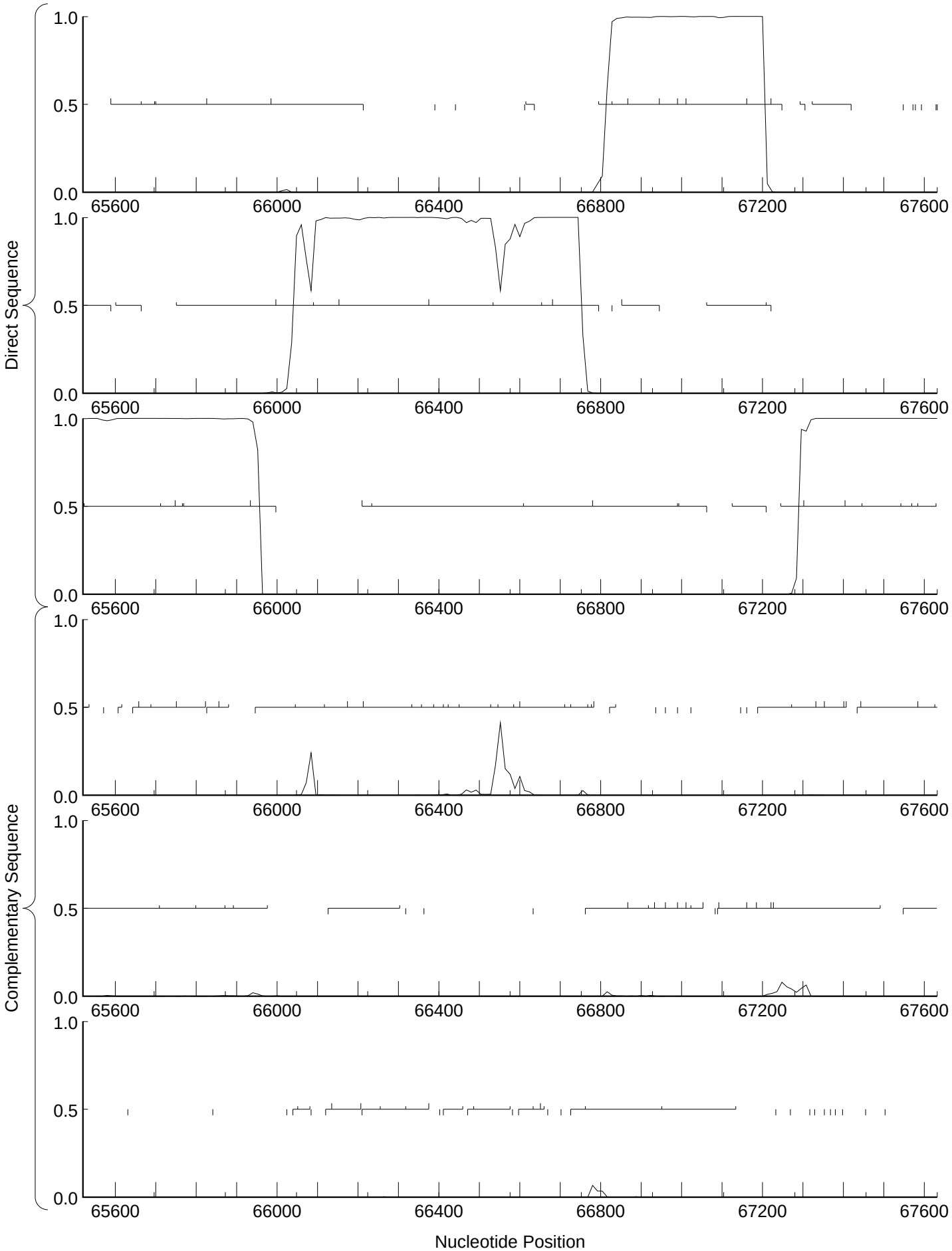
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 31/75



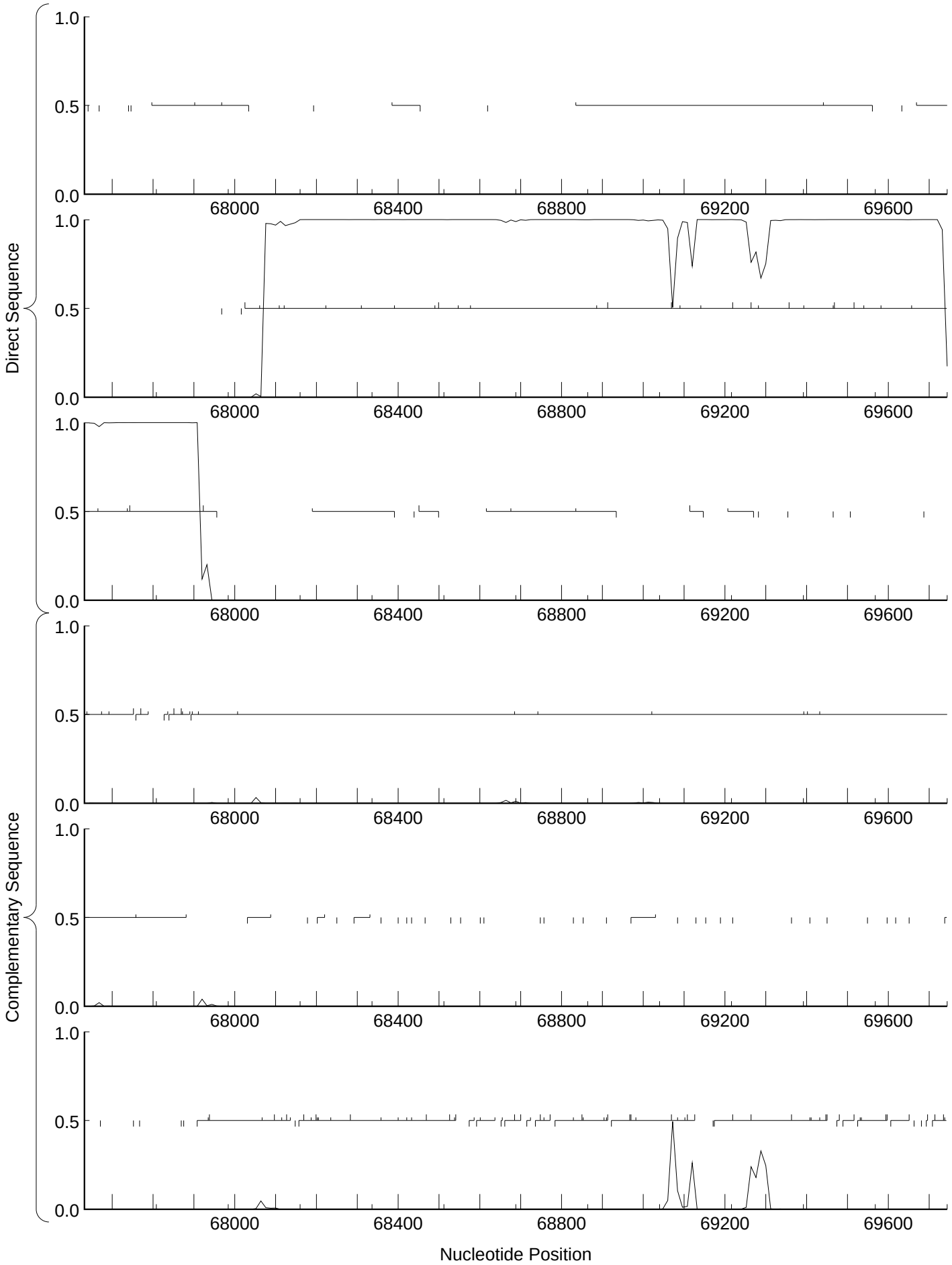
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 32/75



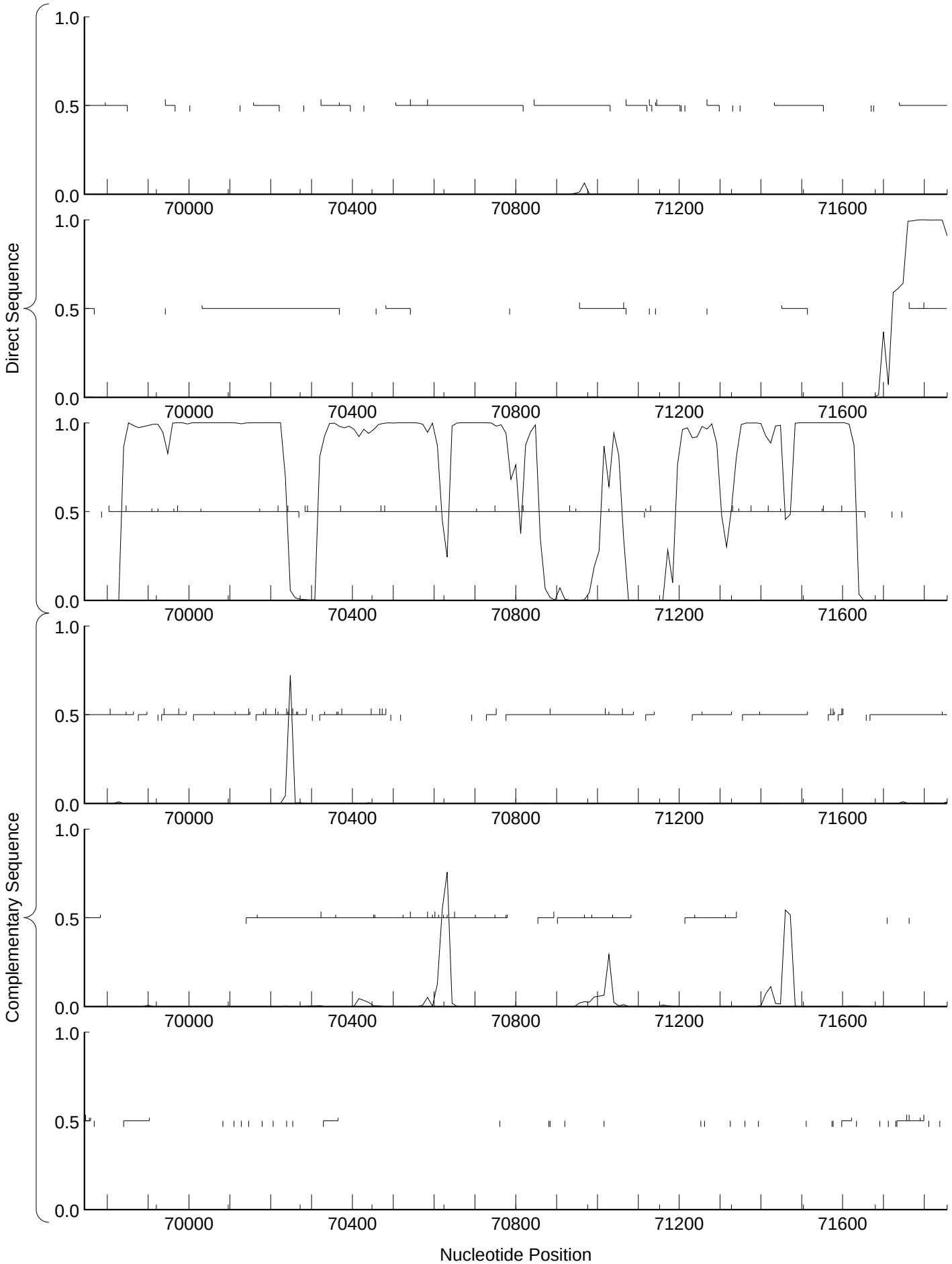
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 33/75



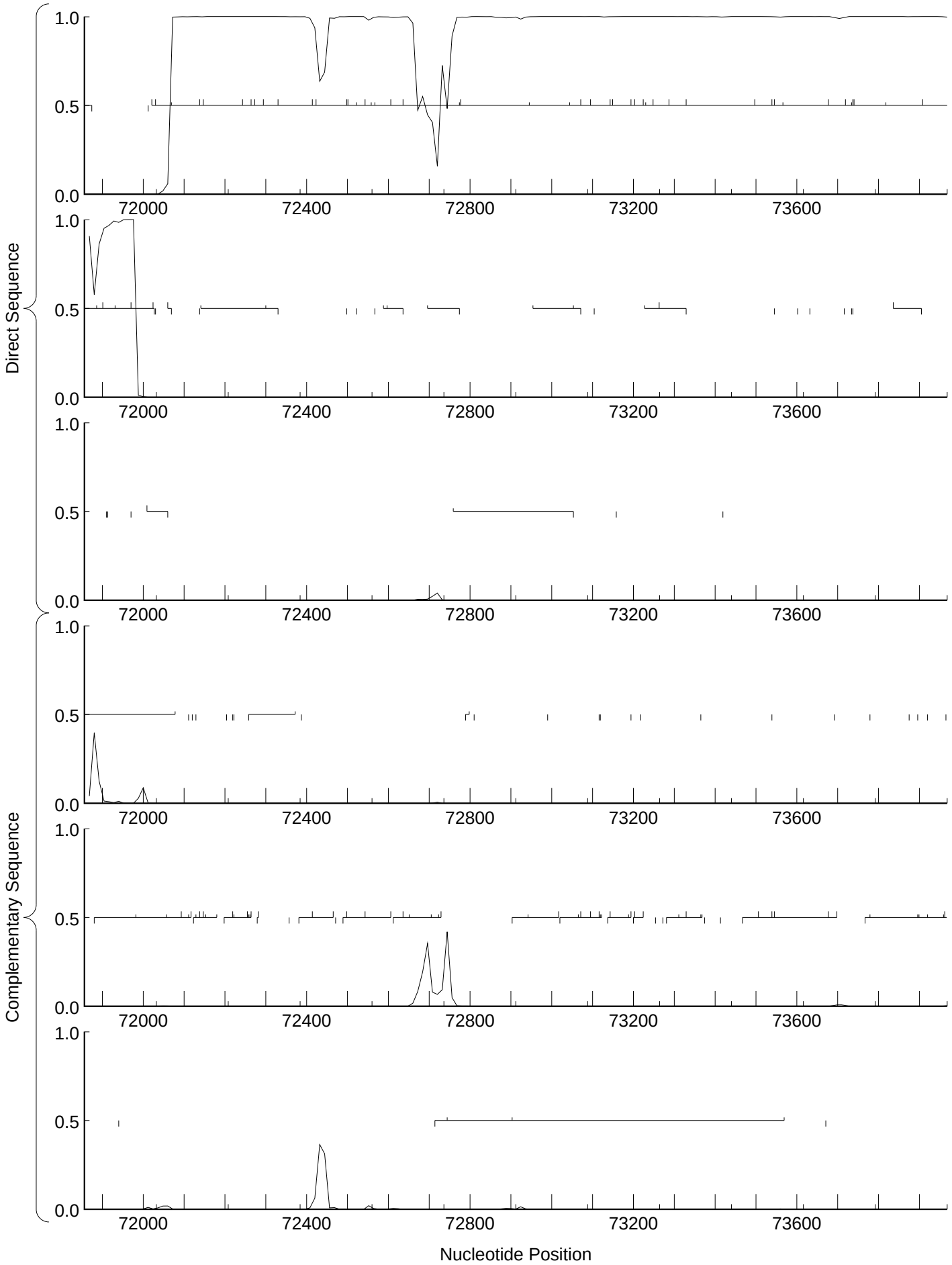
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 34/75



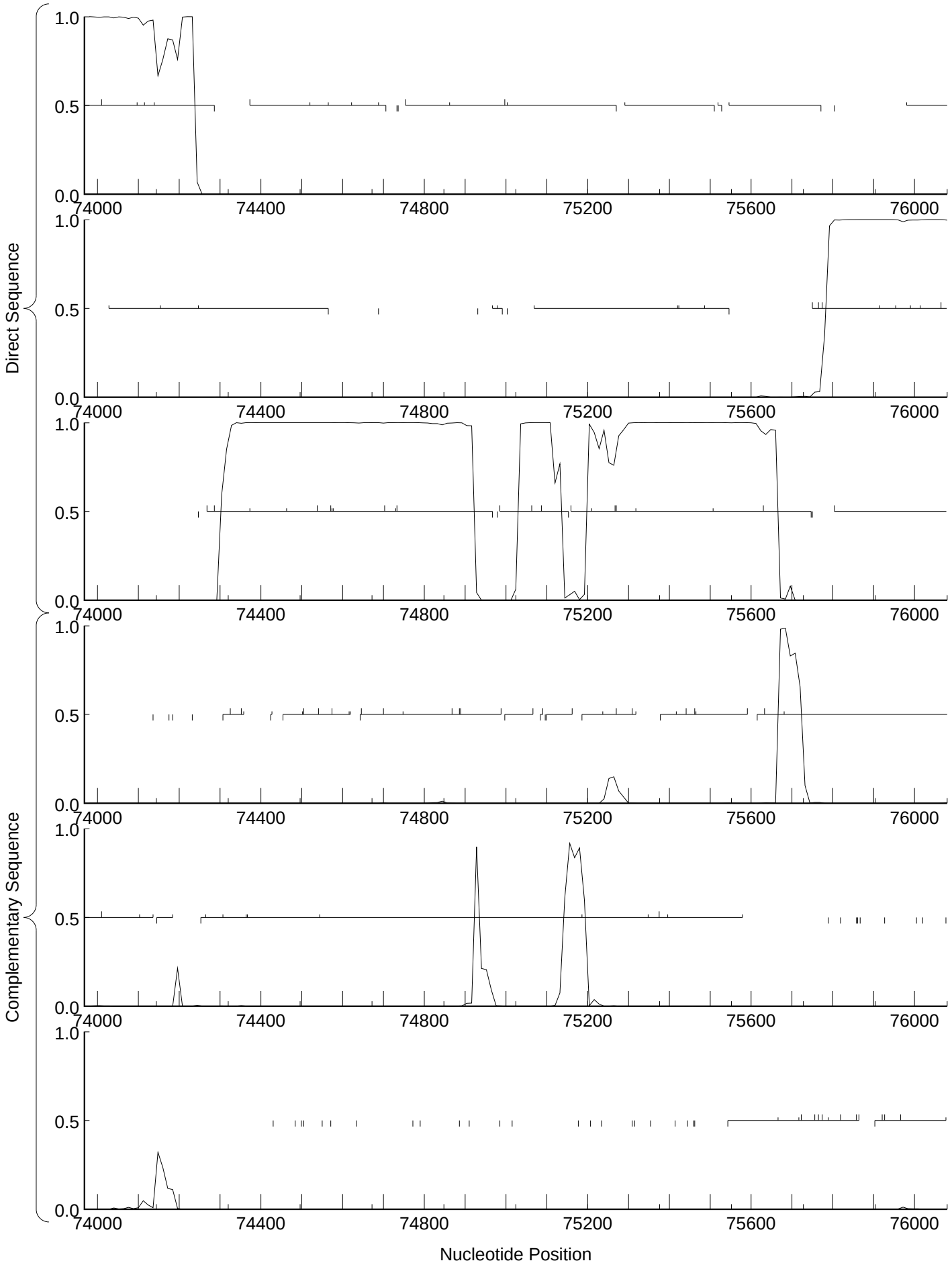
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 35/75



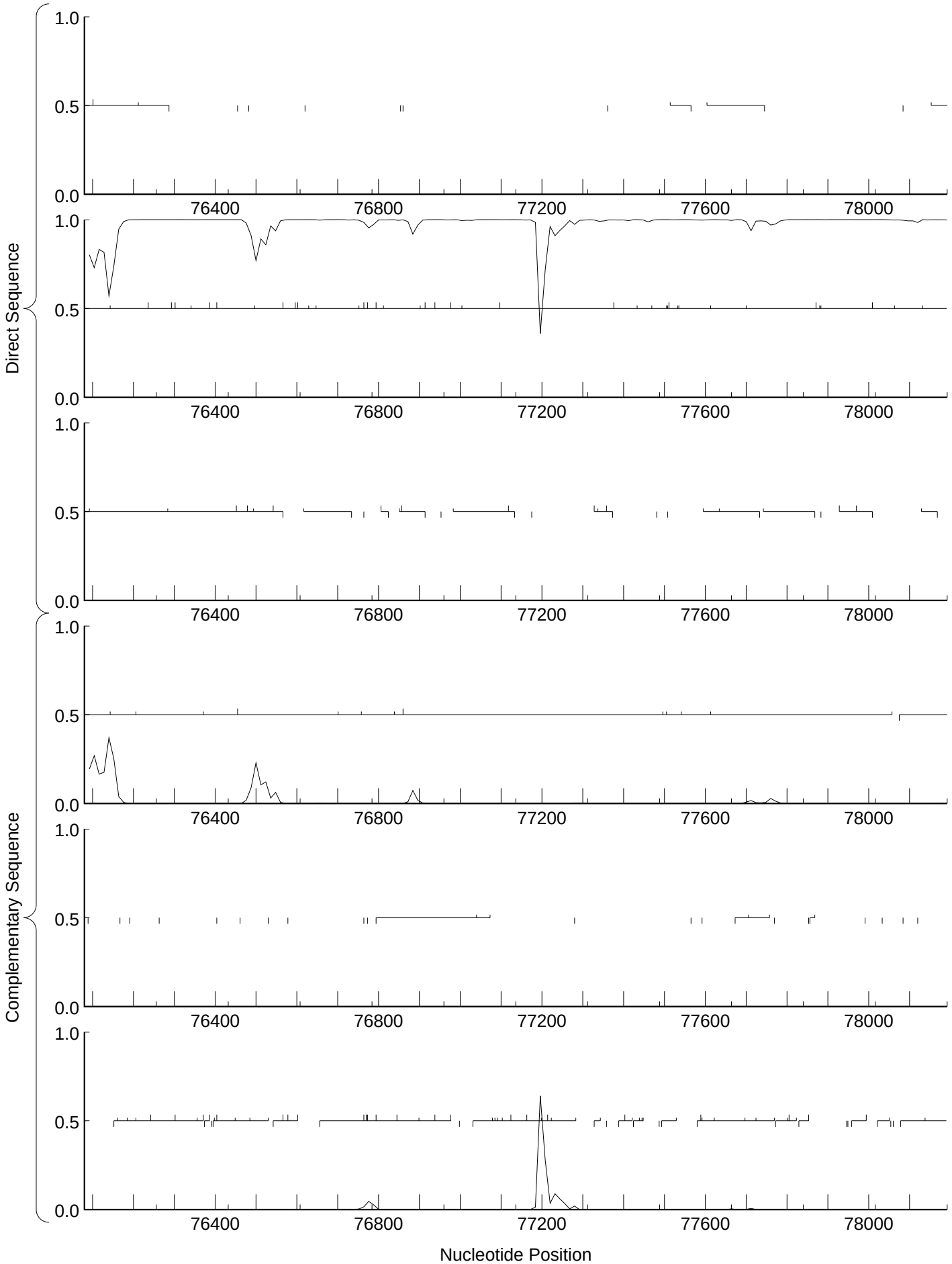
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 36/75



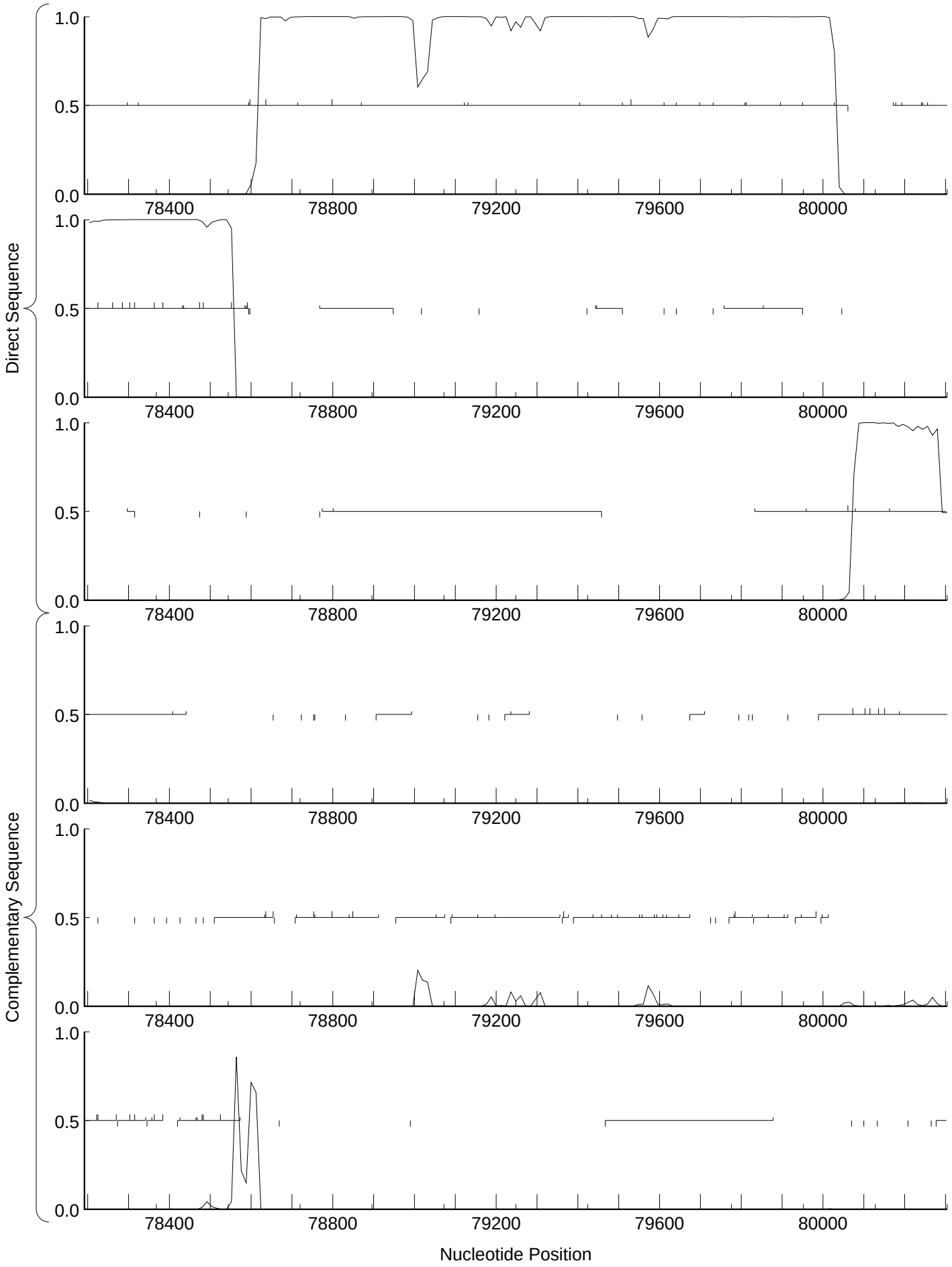
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 37/75



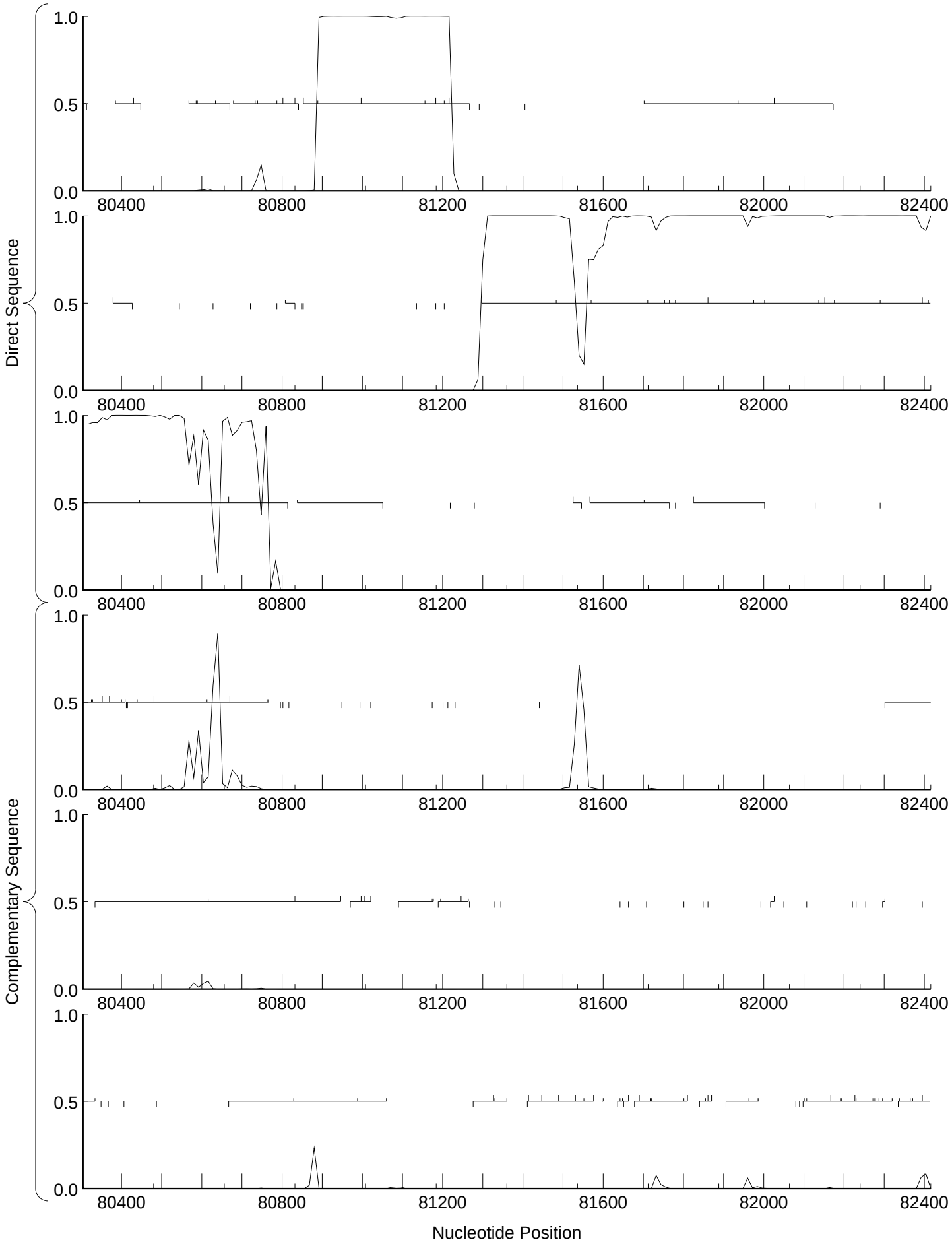
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 38/75



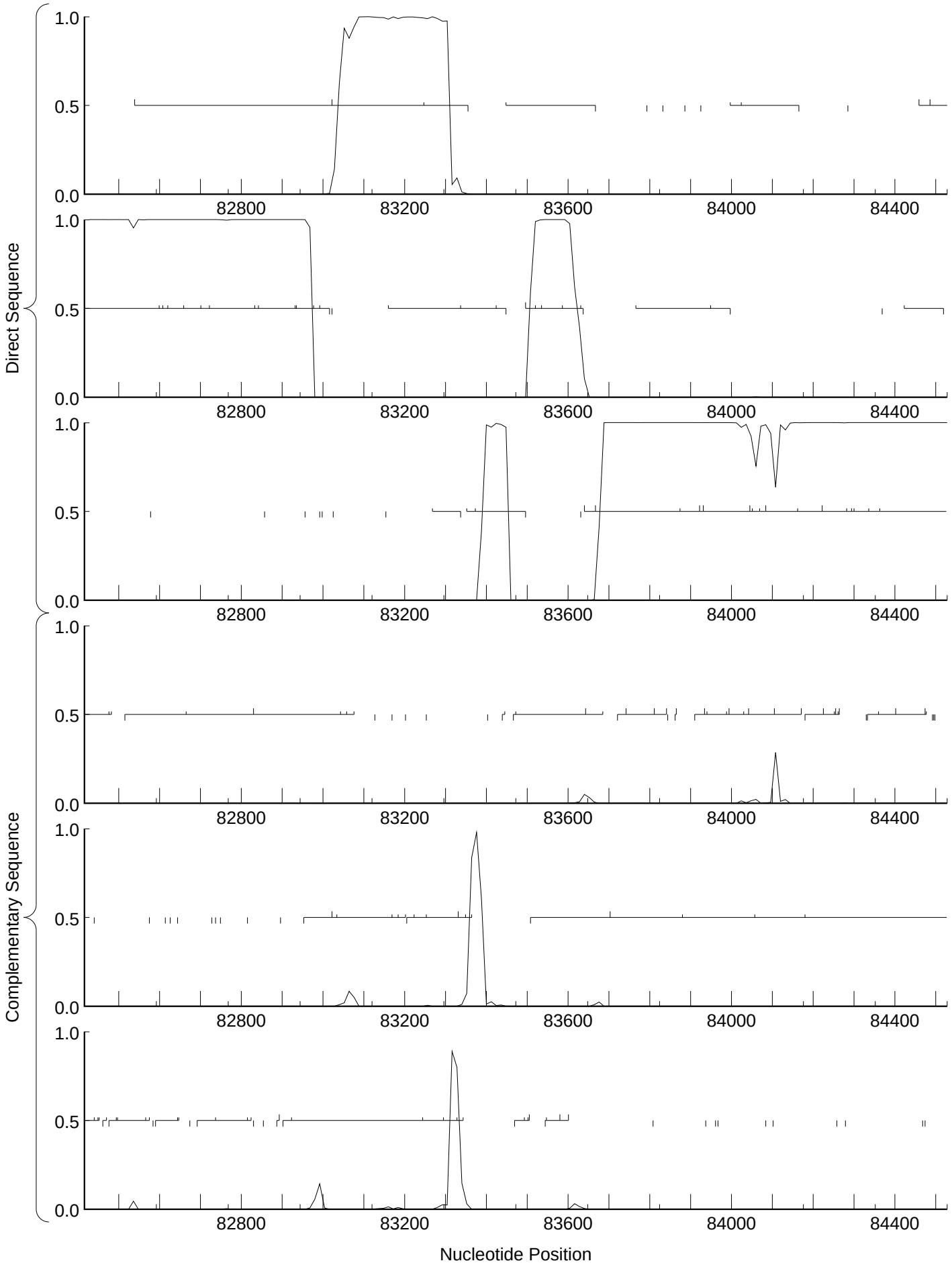
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 39/75



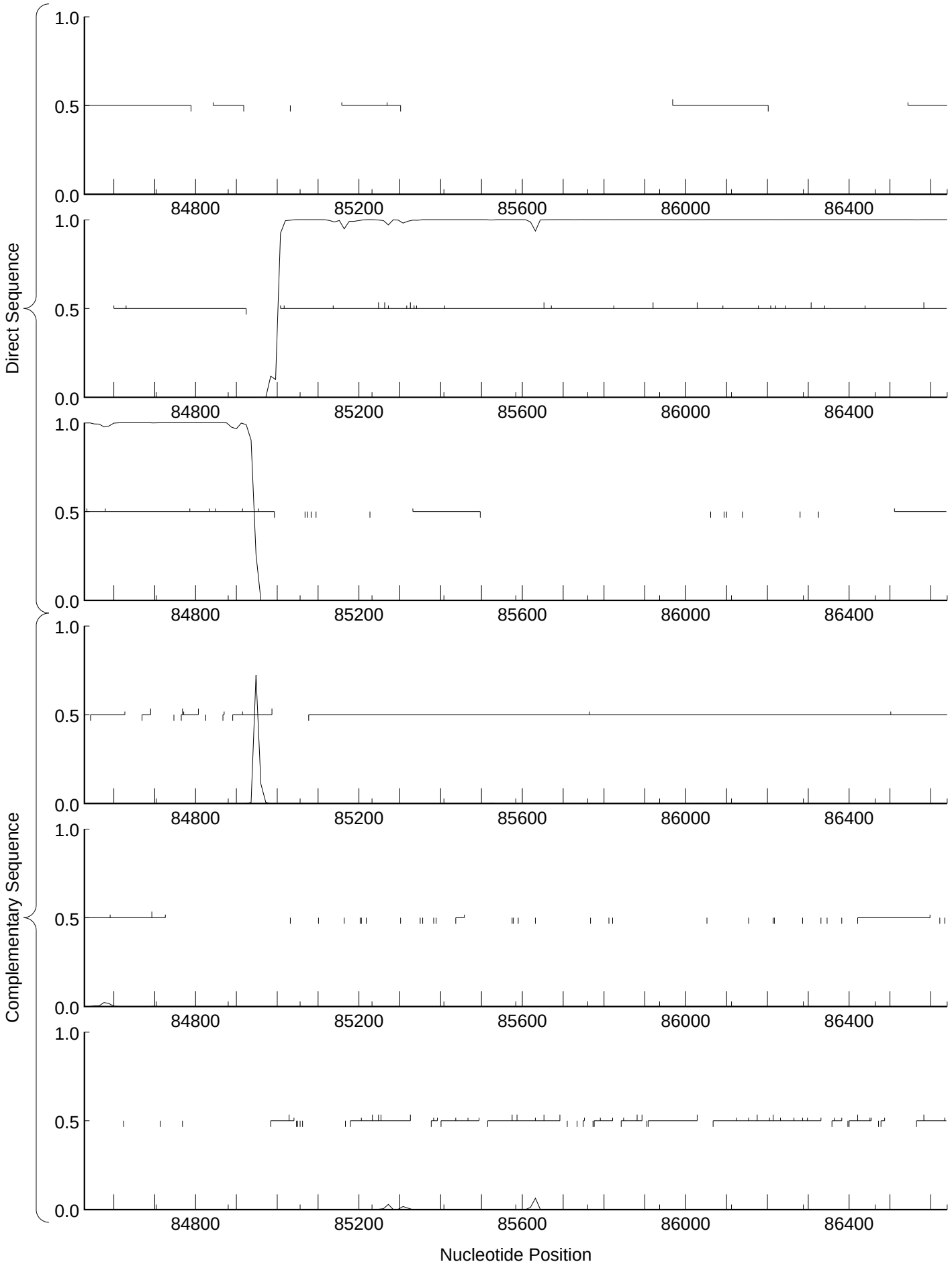
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 40/75



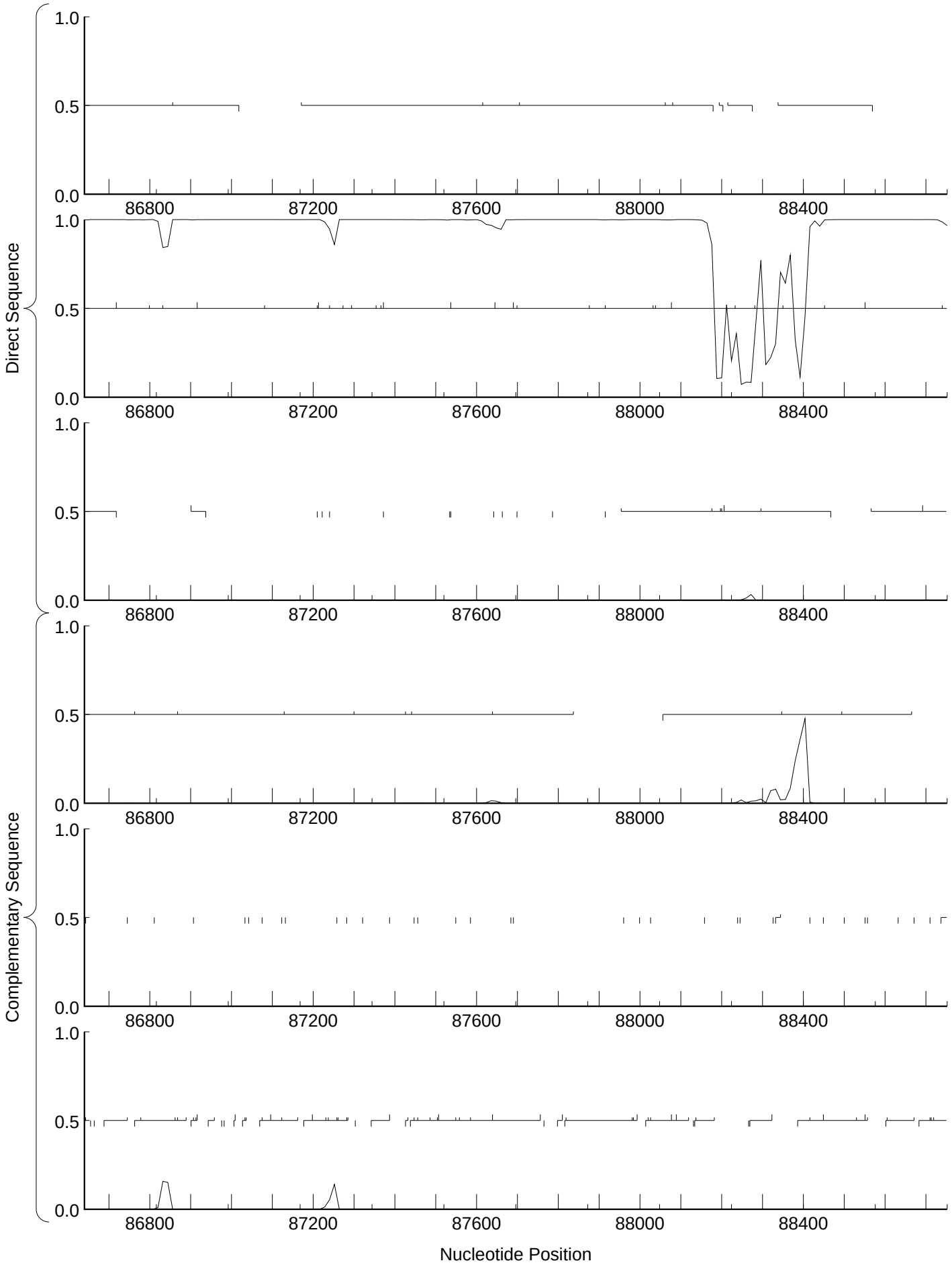
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 41/75



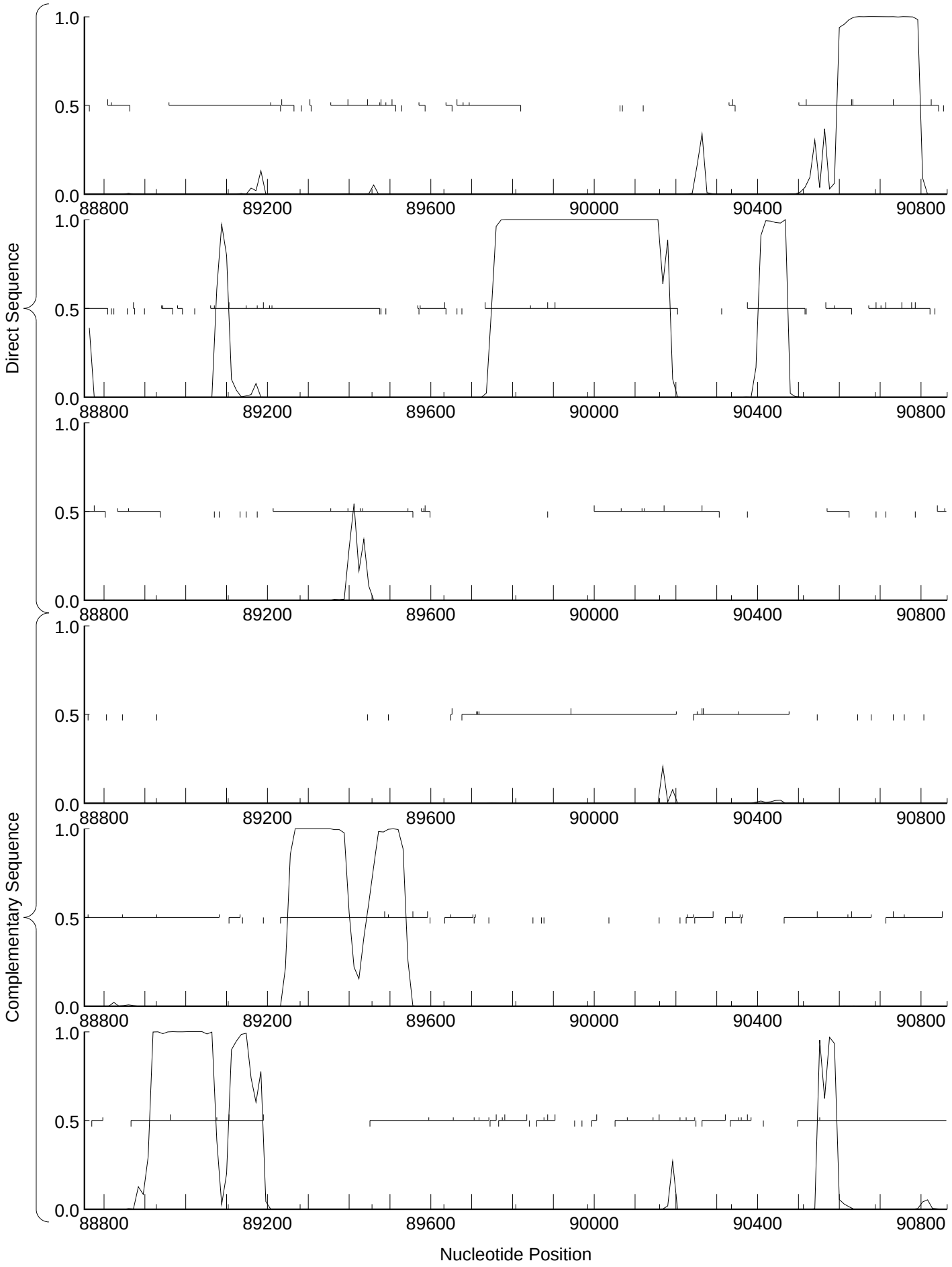
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 42/75



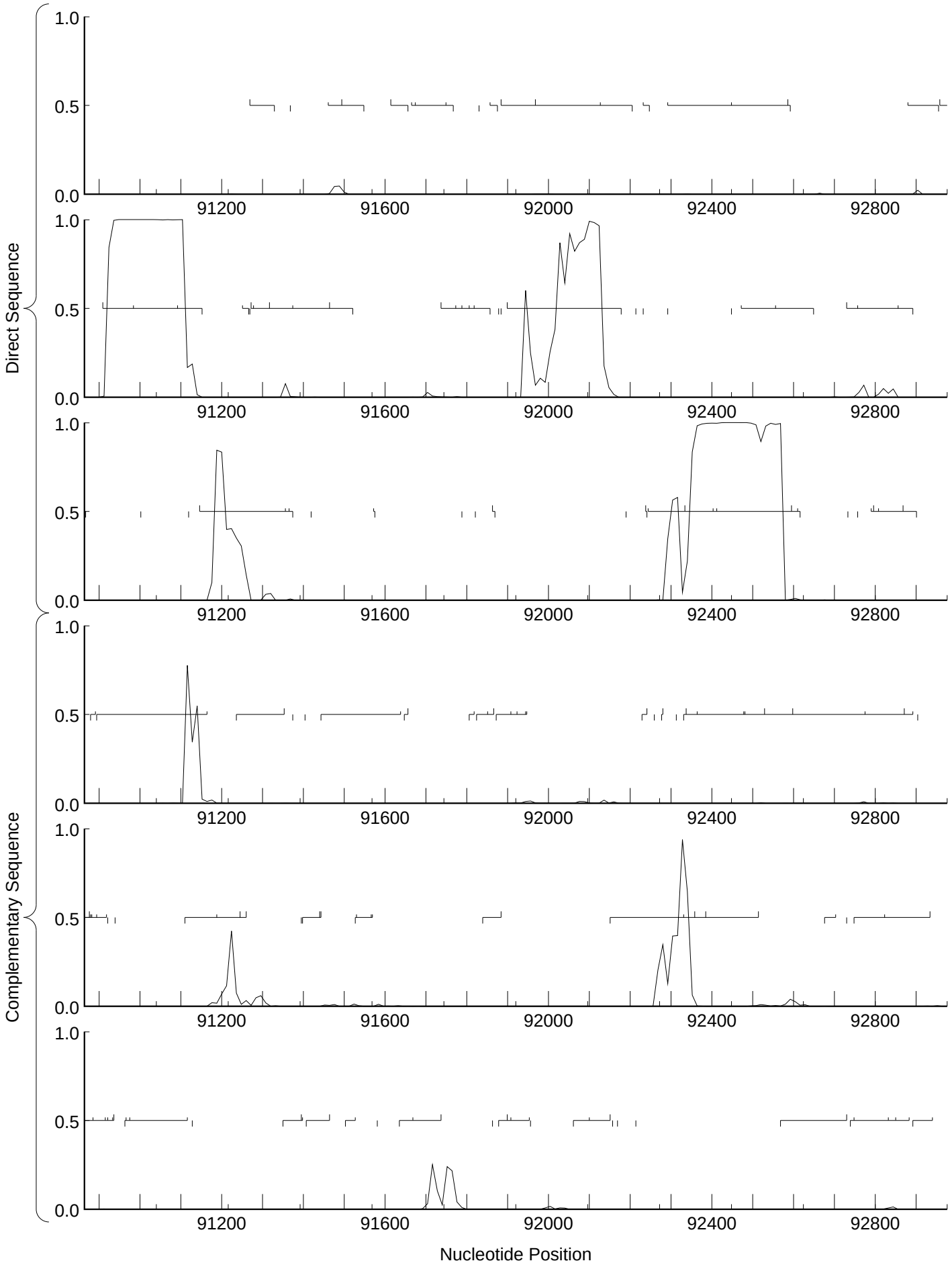
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 43/75



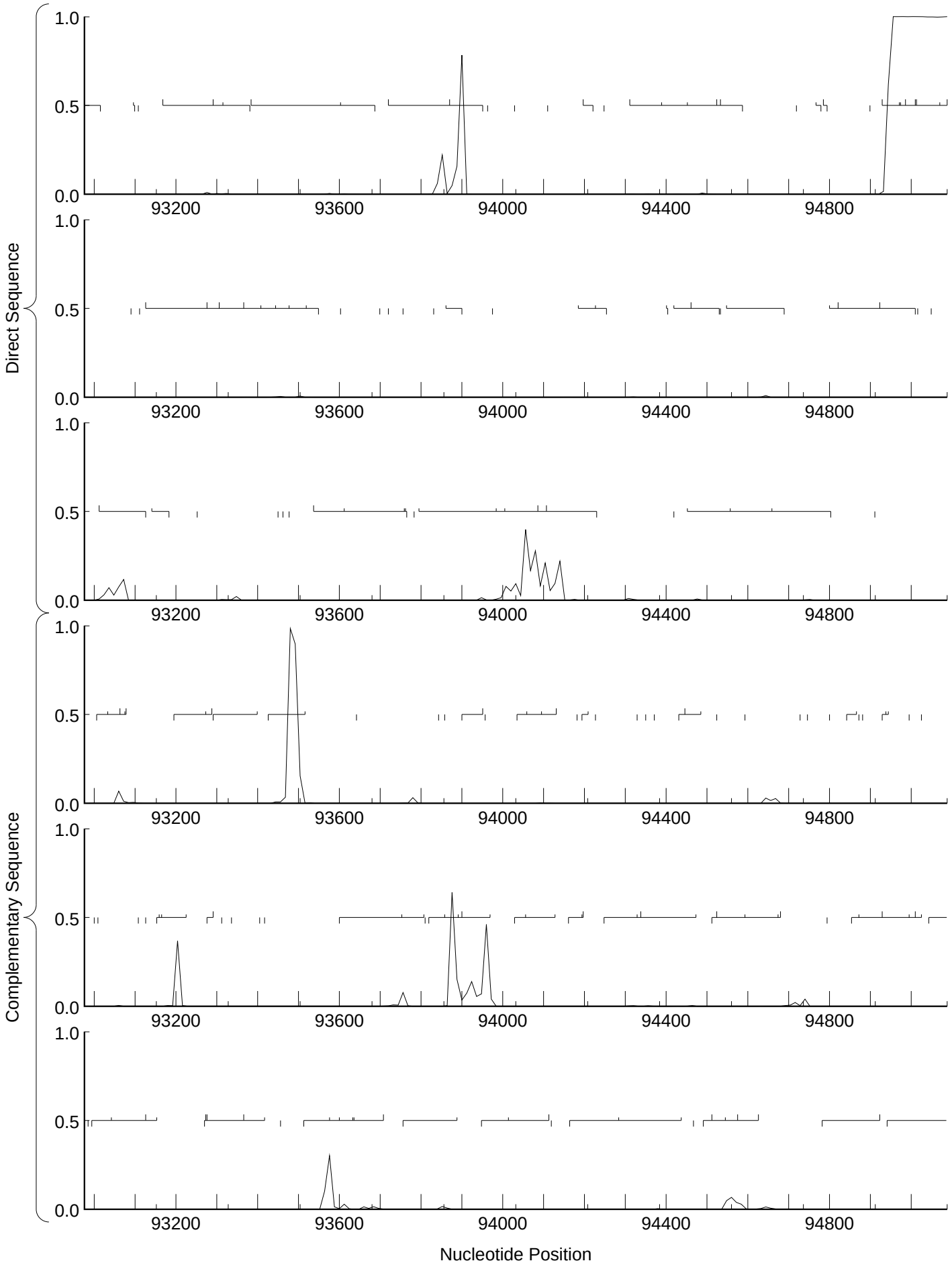
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 44/75



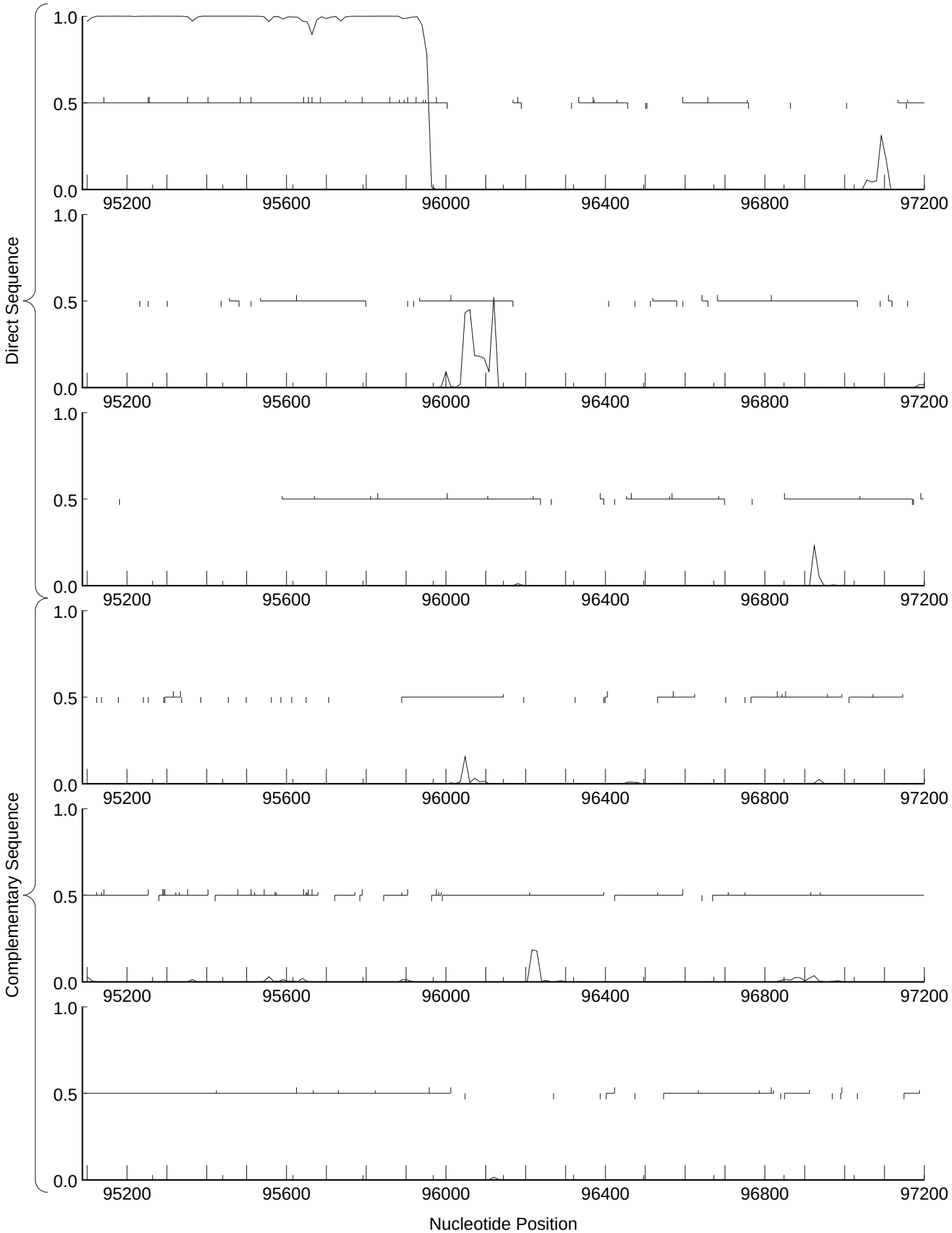
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 45/75



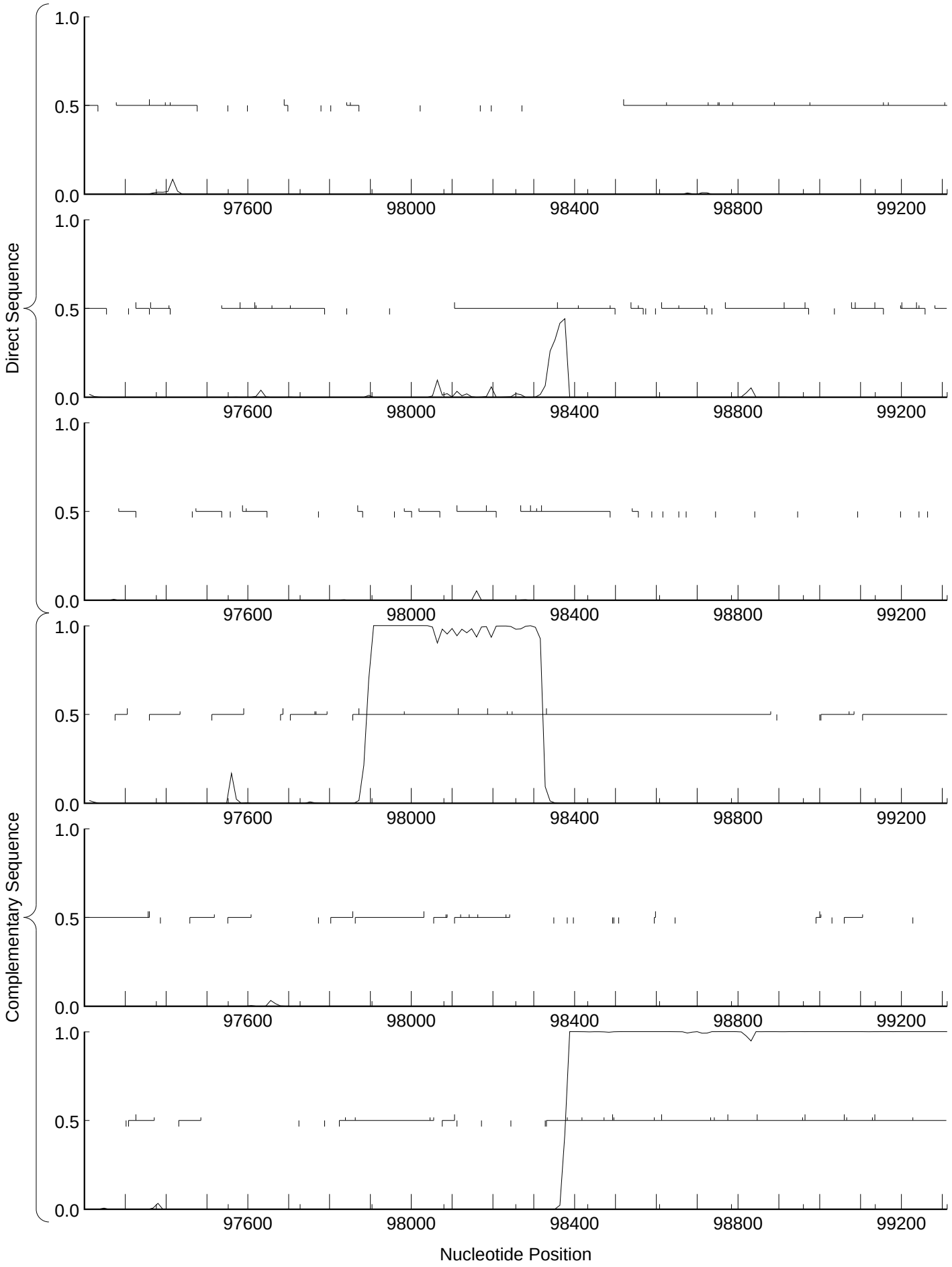
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 46/75



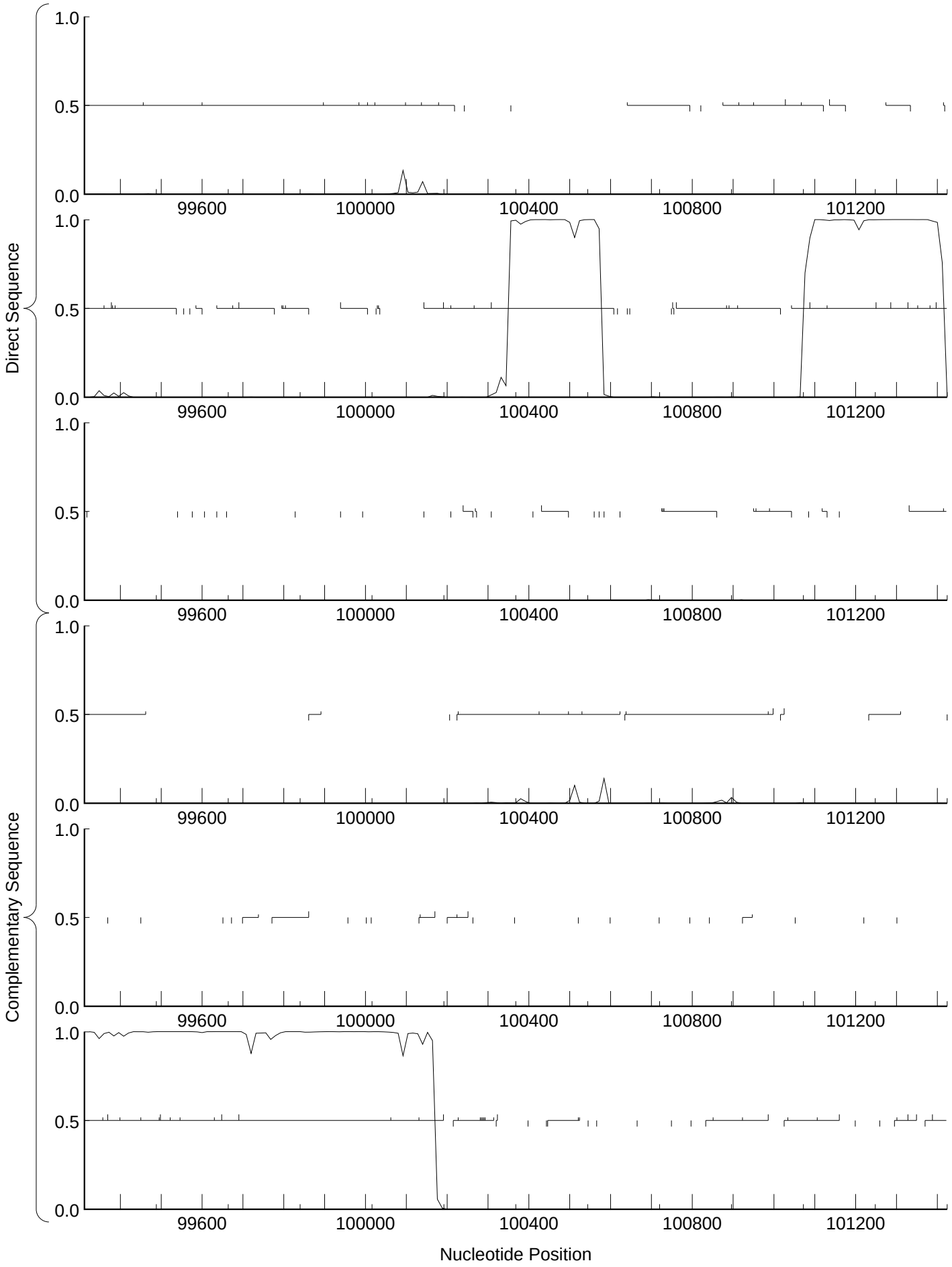
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 47/75



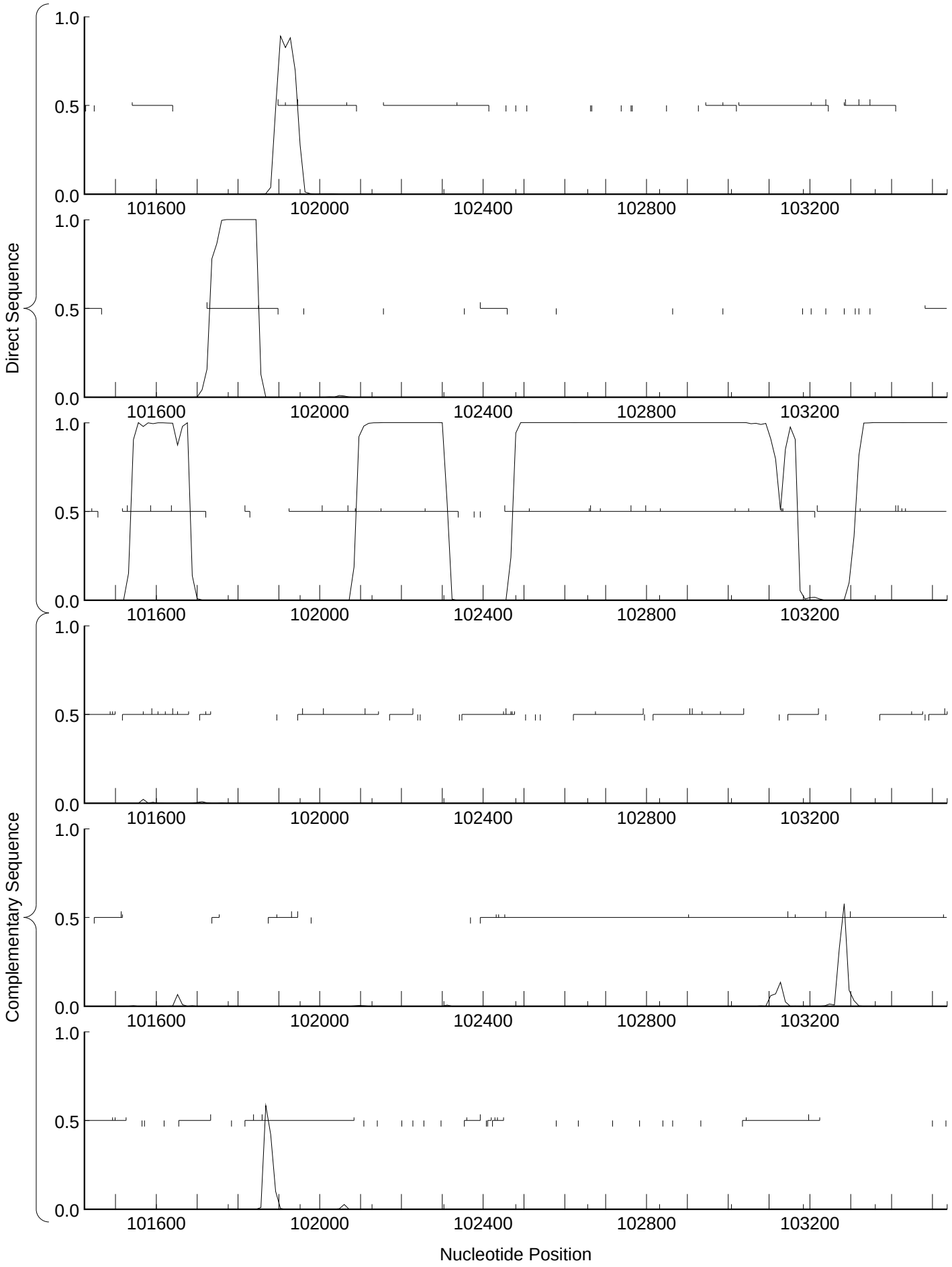
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 48/75



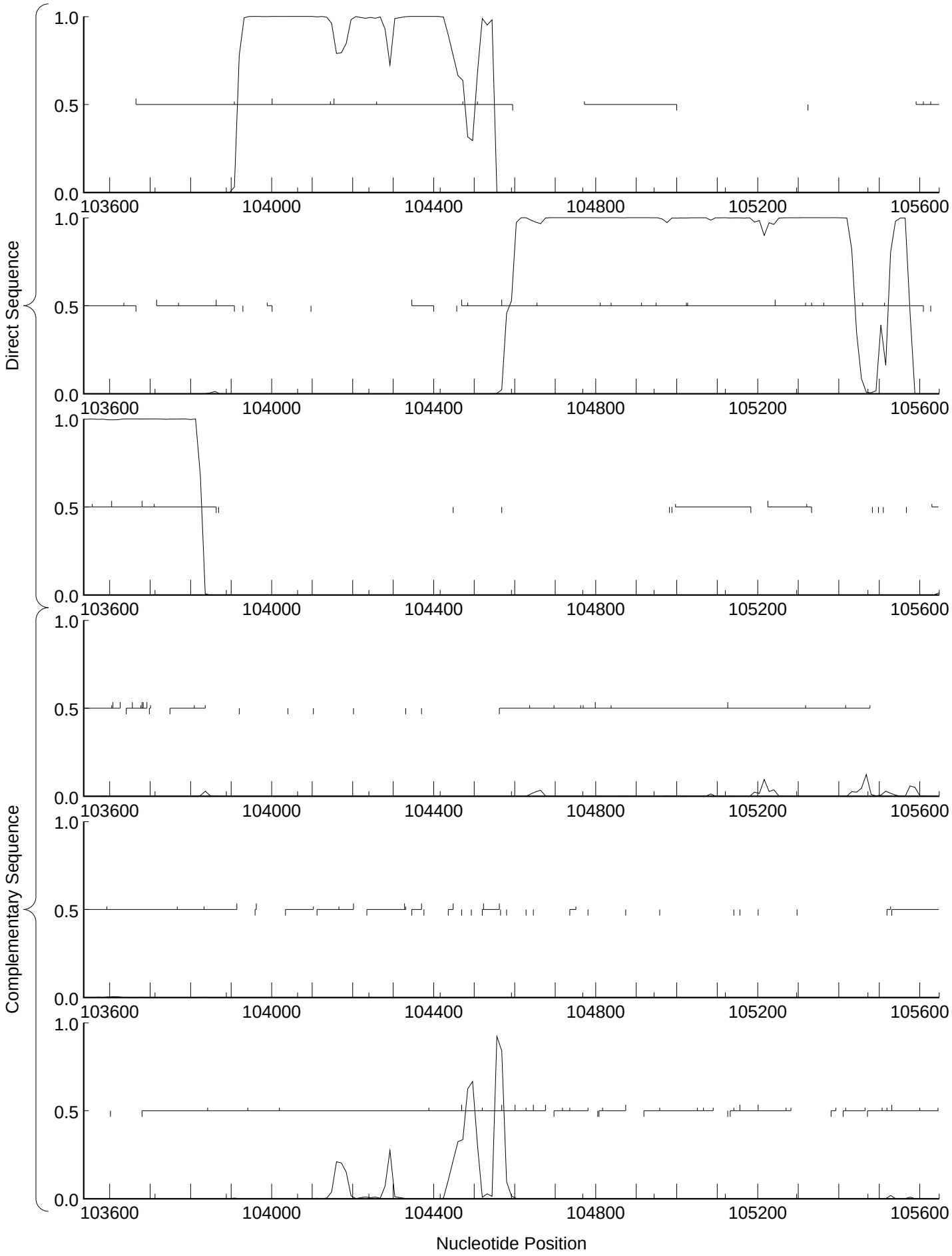
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 49/75



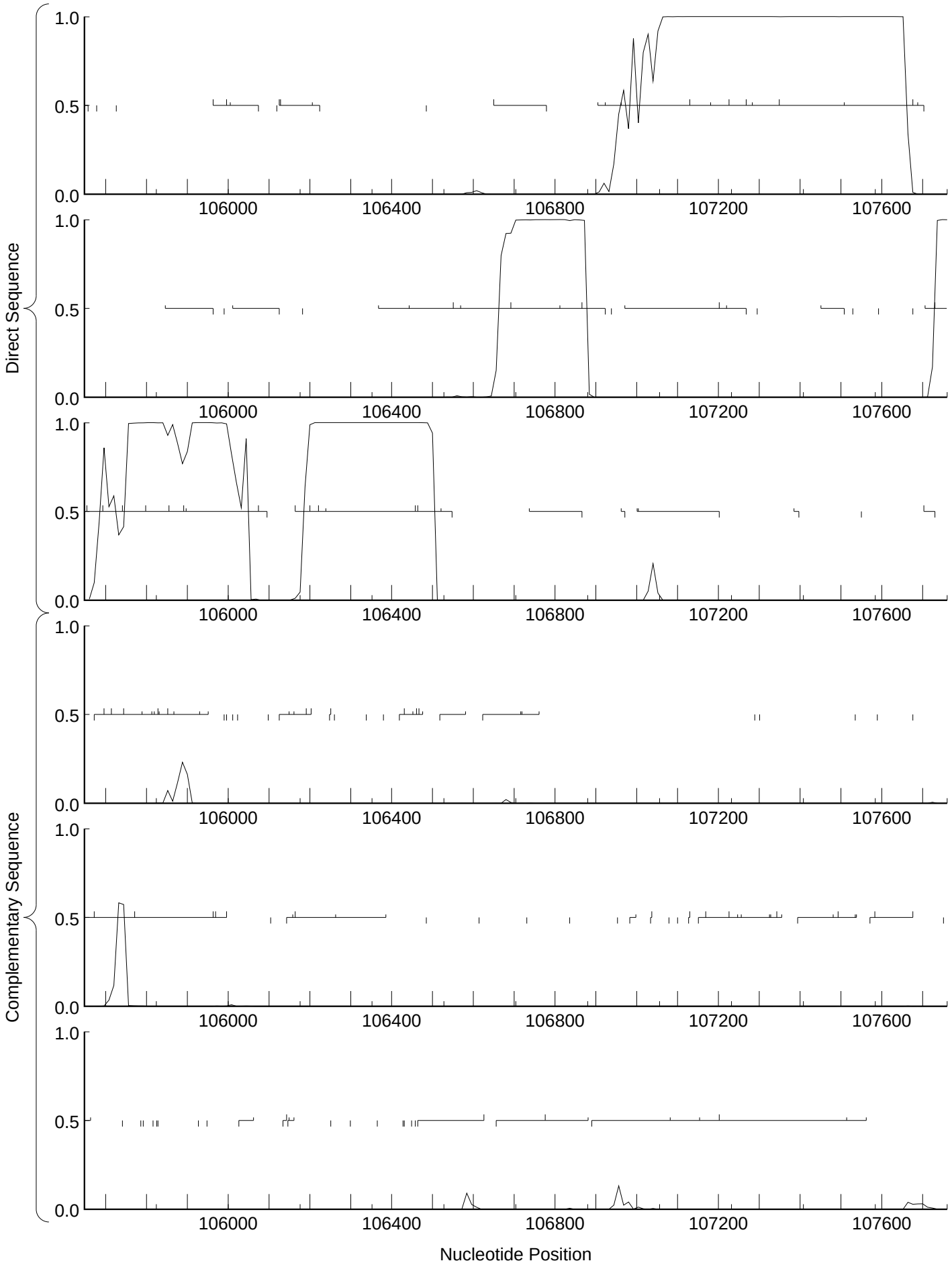
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 50/75



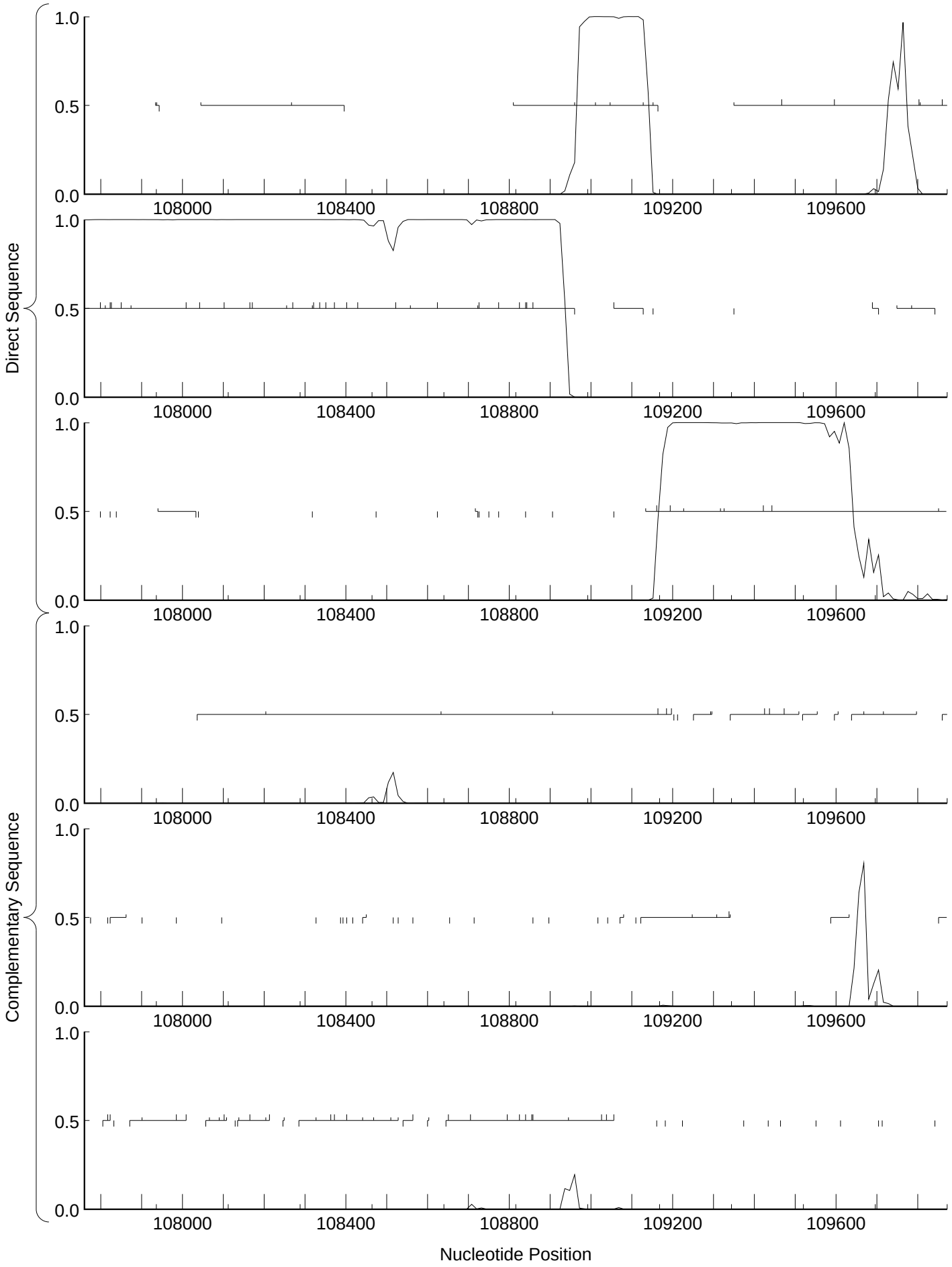
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 51/75



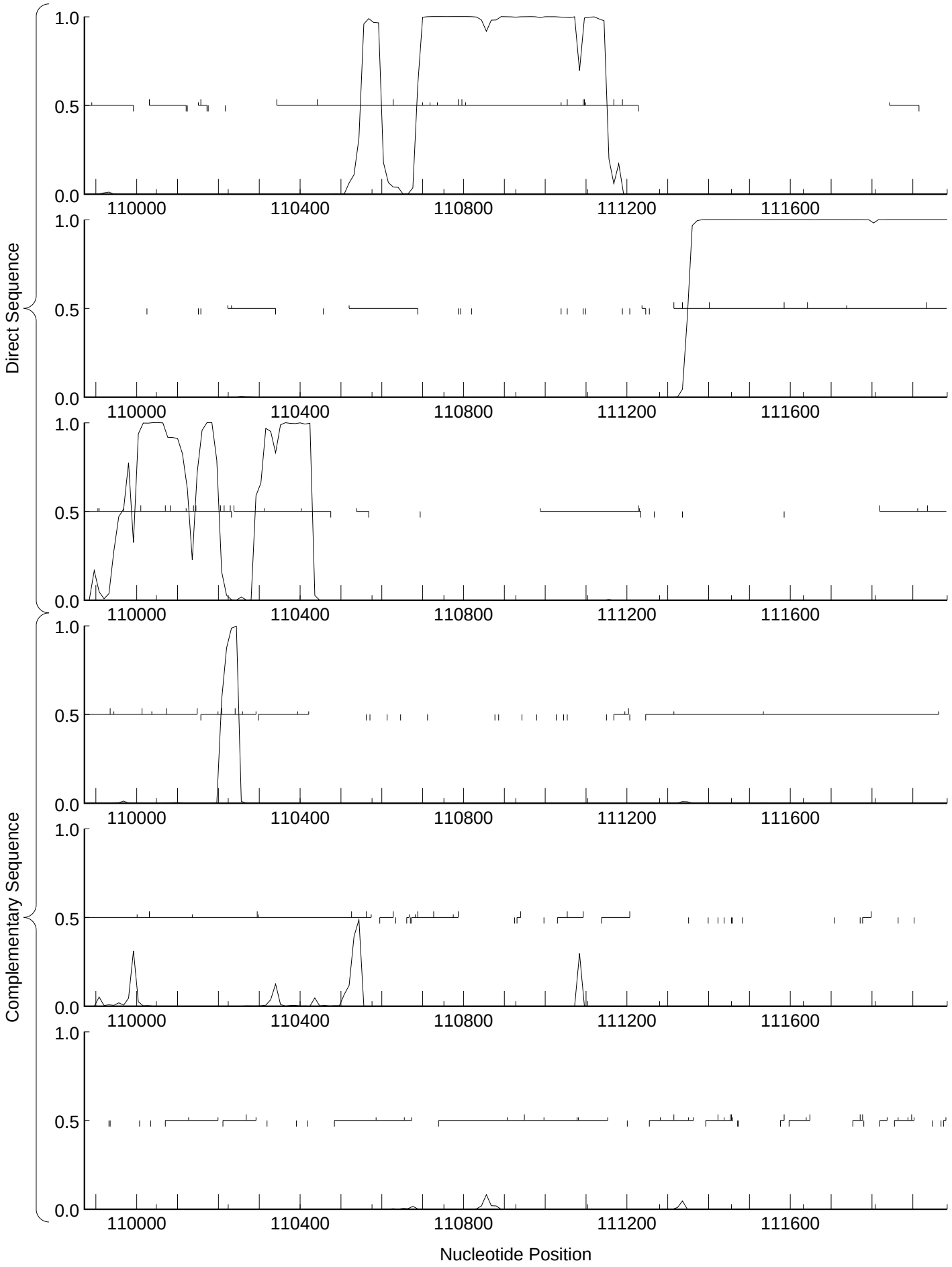
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 52/75



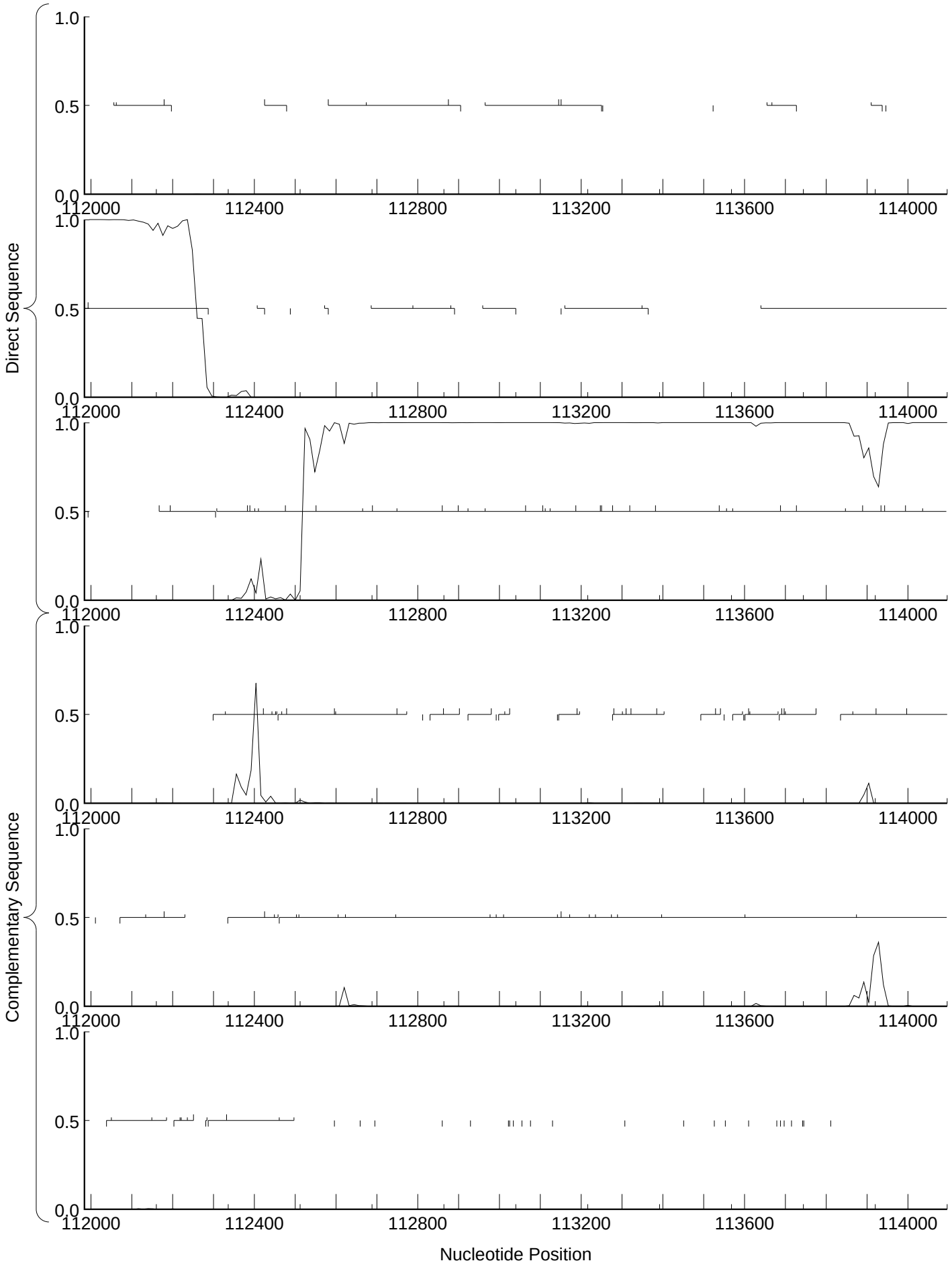
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 53/75



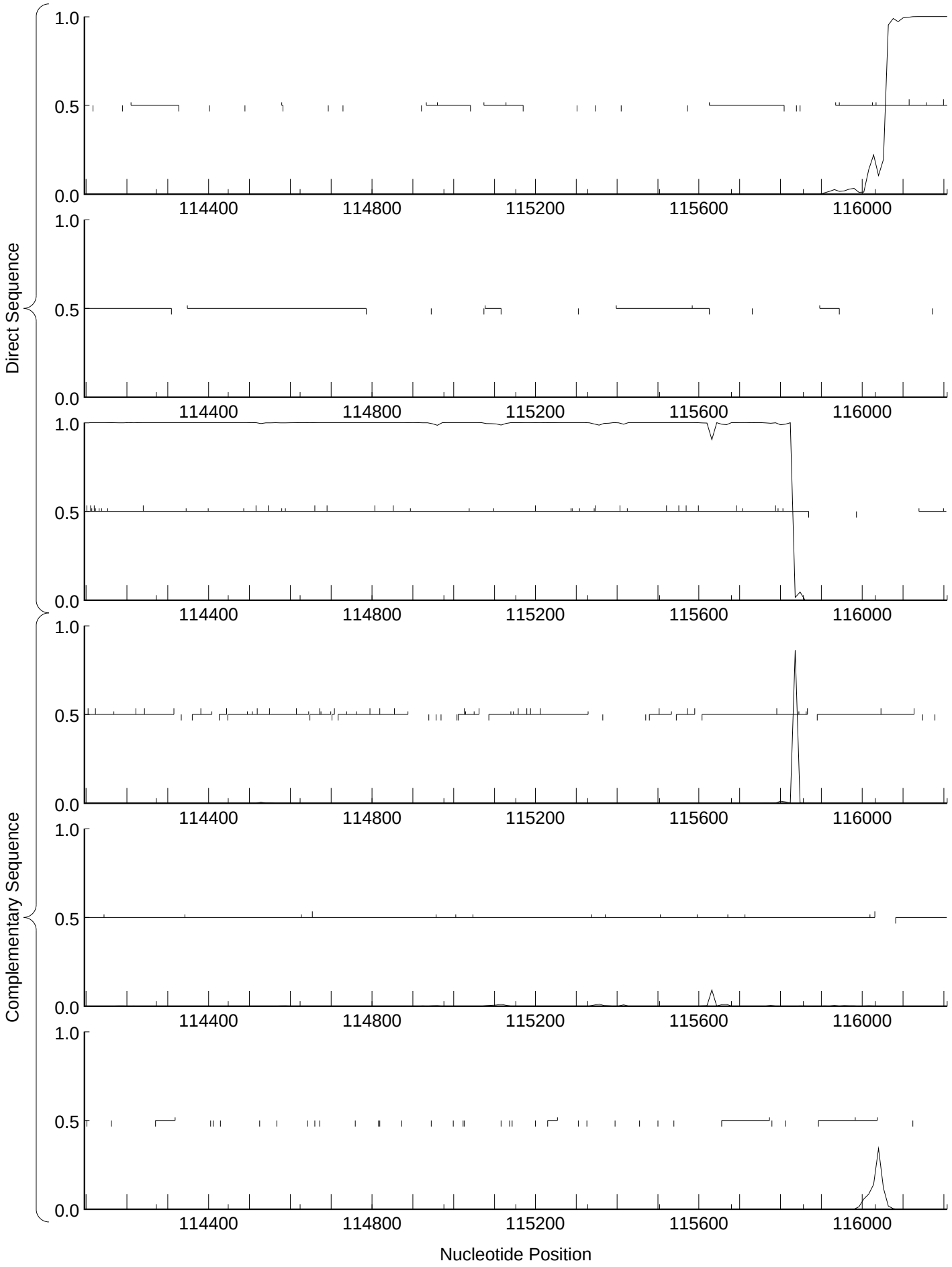
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 54/75



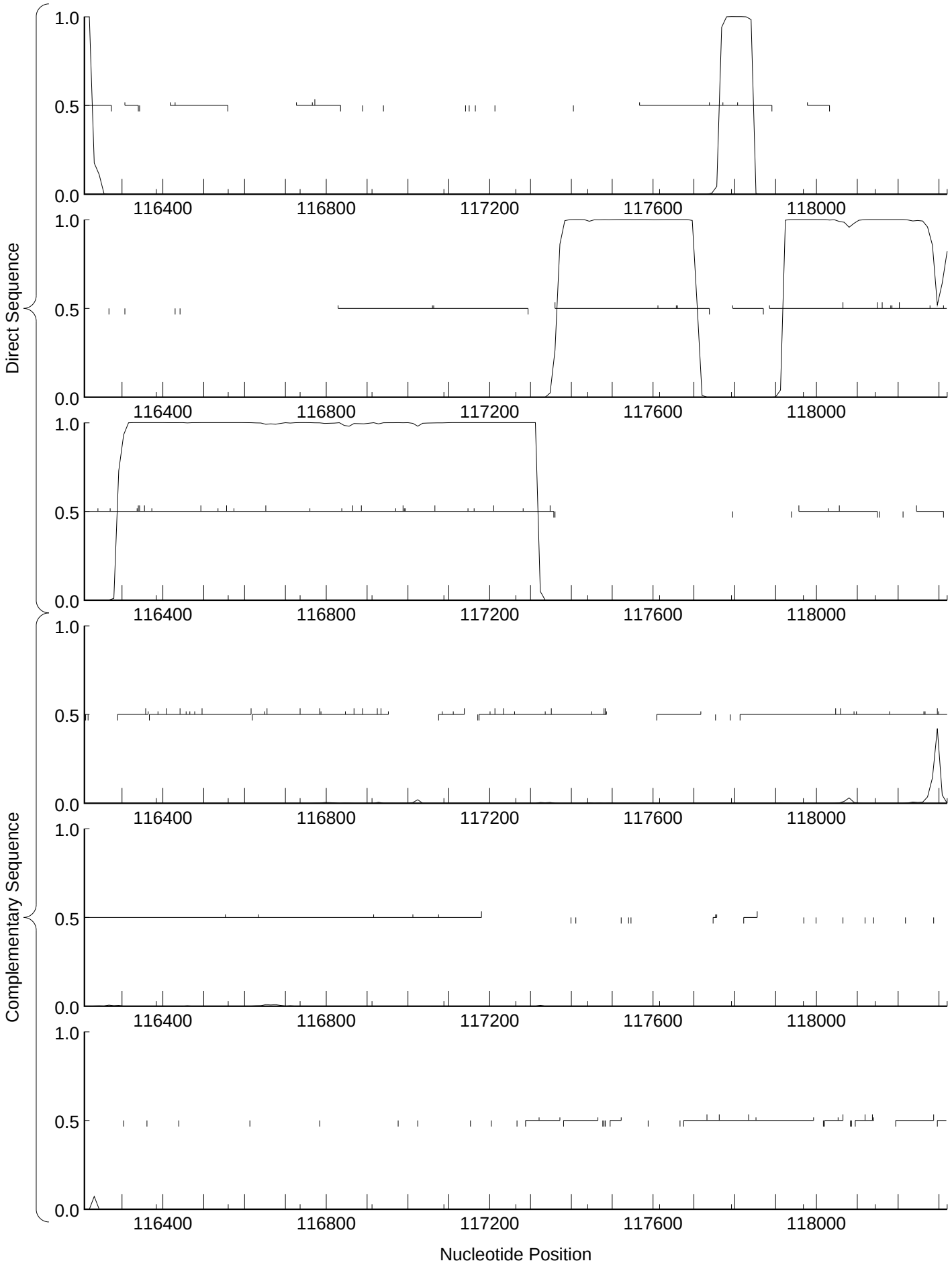
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 55/75



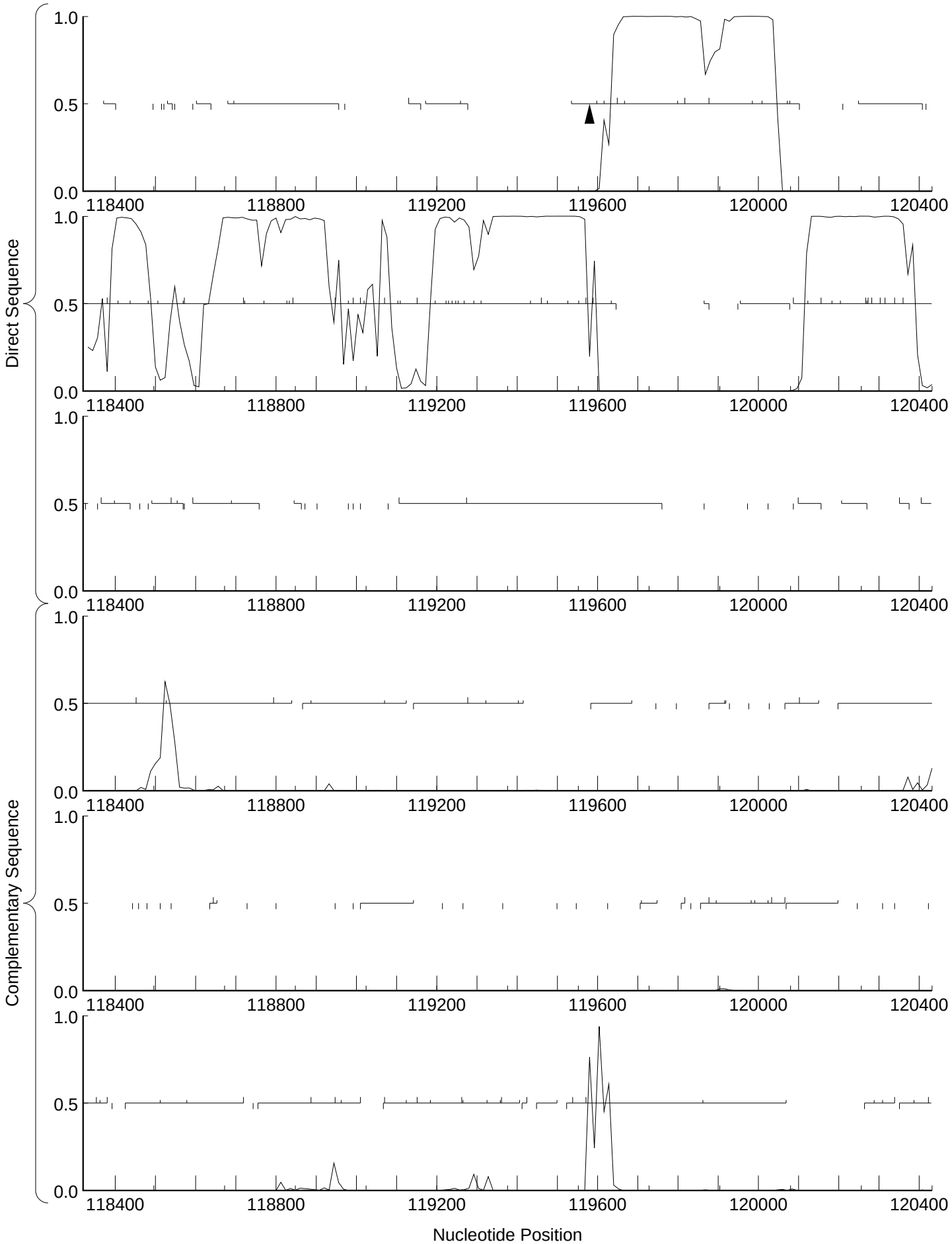
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 56/75



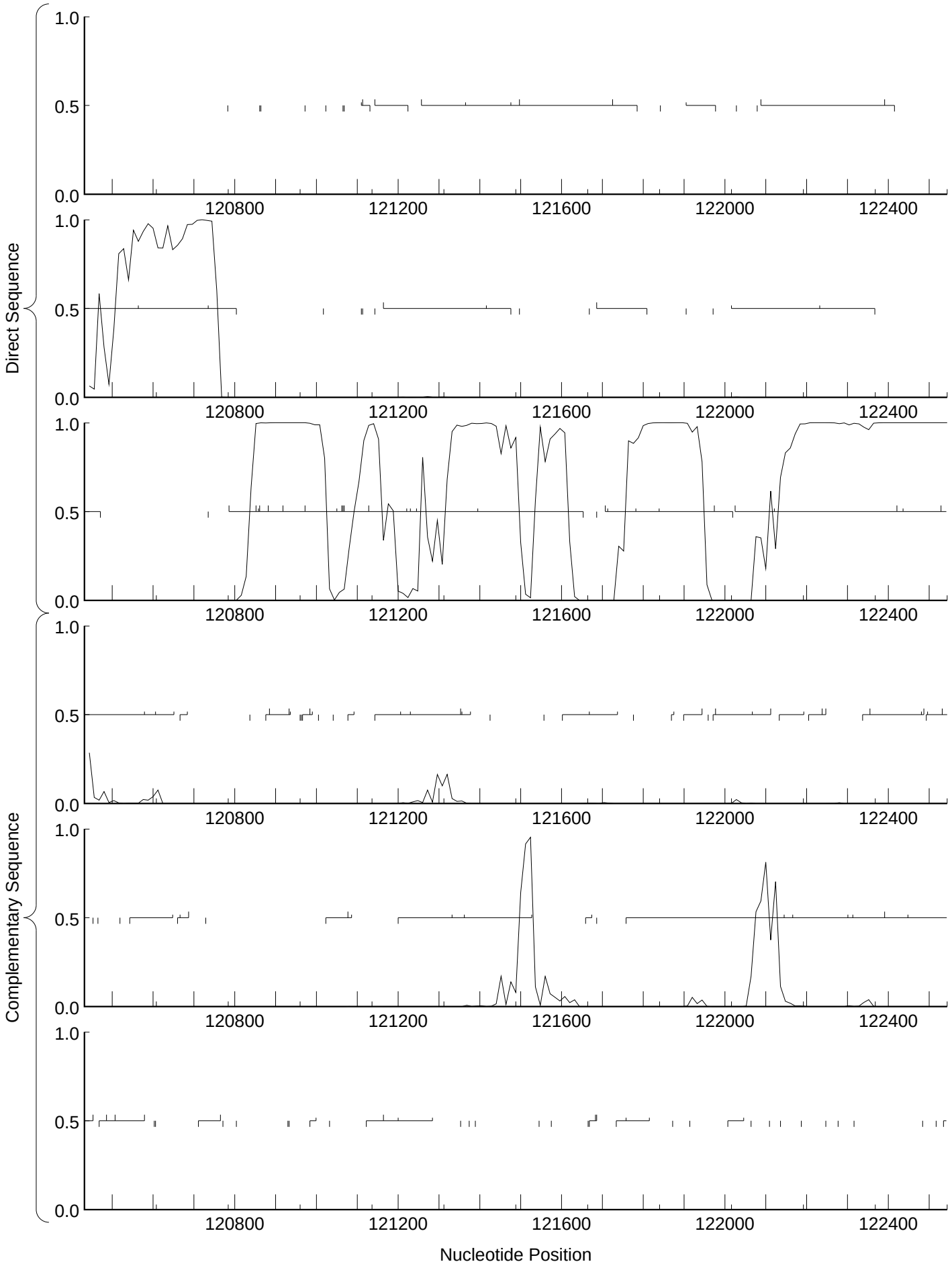
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 57/75



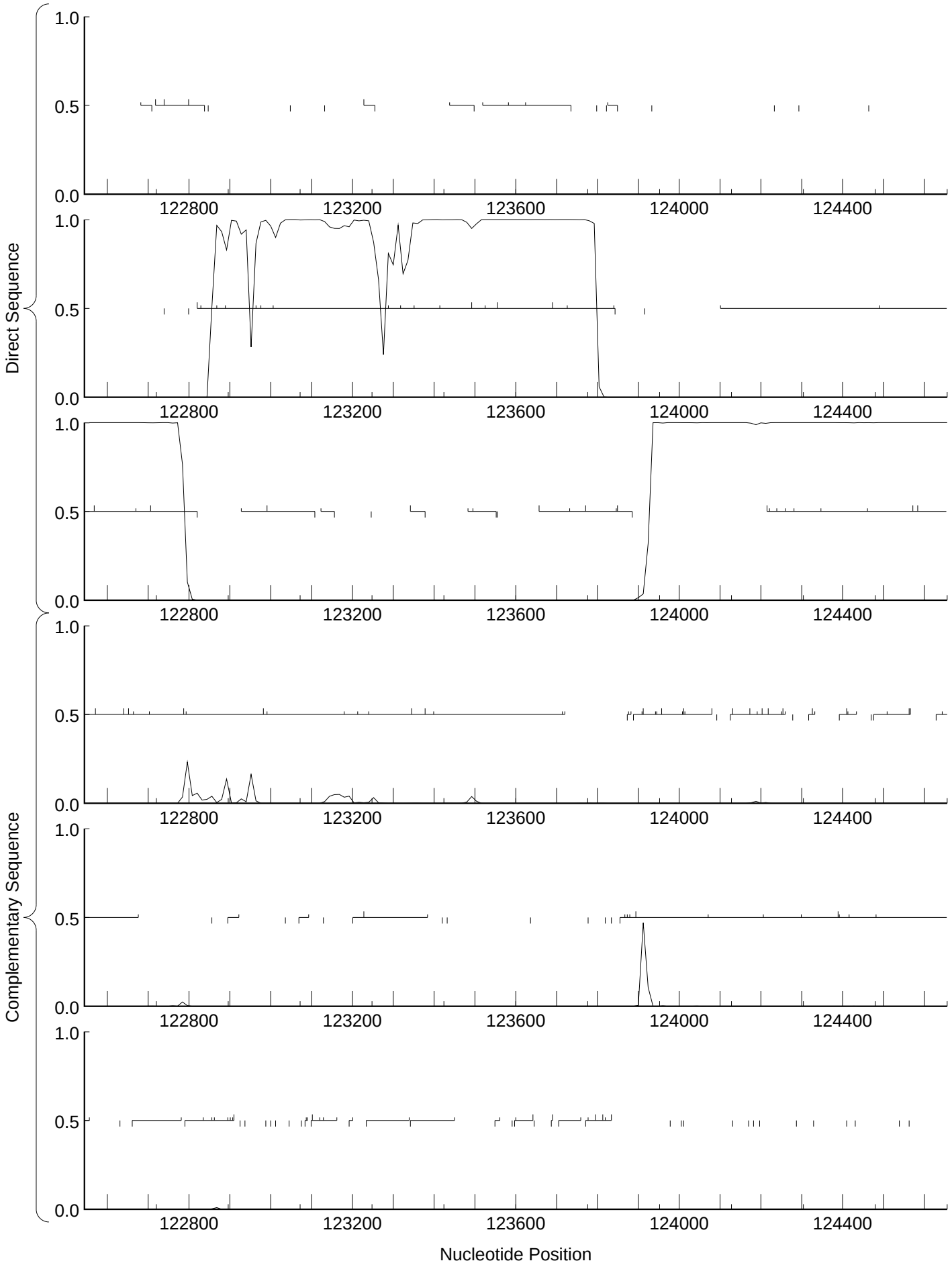
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 58/75



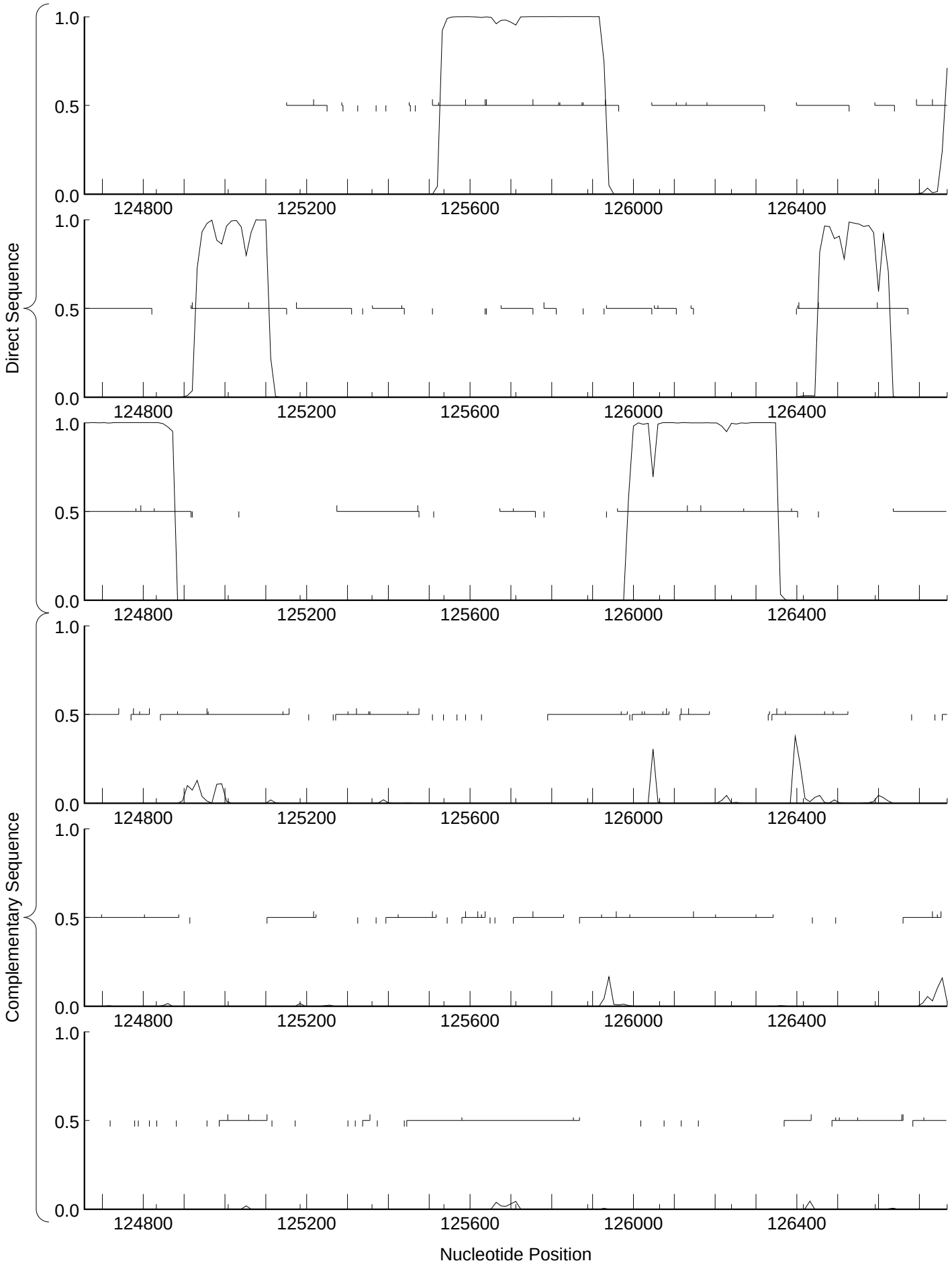
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 59/75



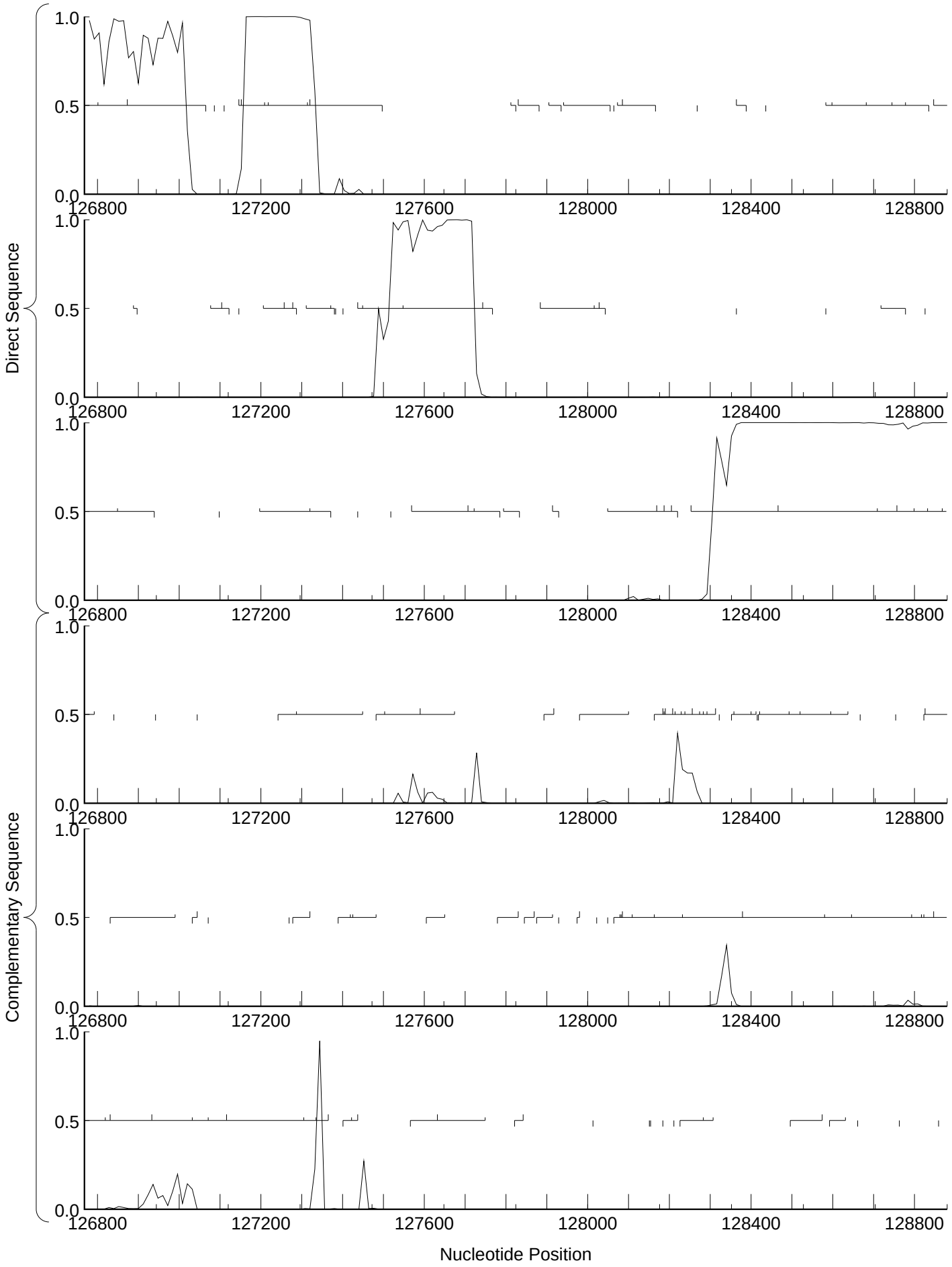
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 60/75



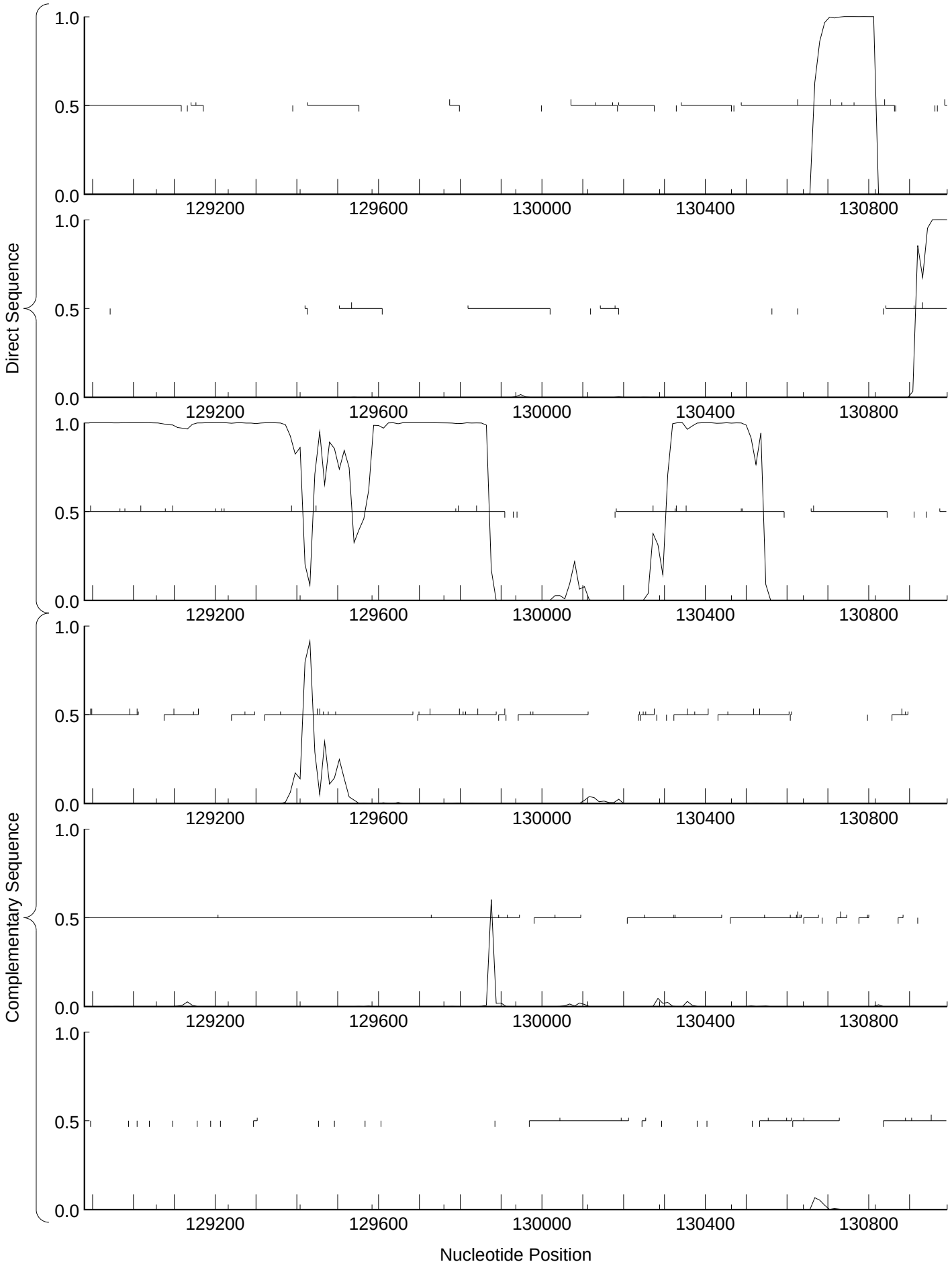
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 61/75



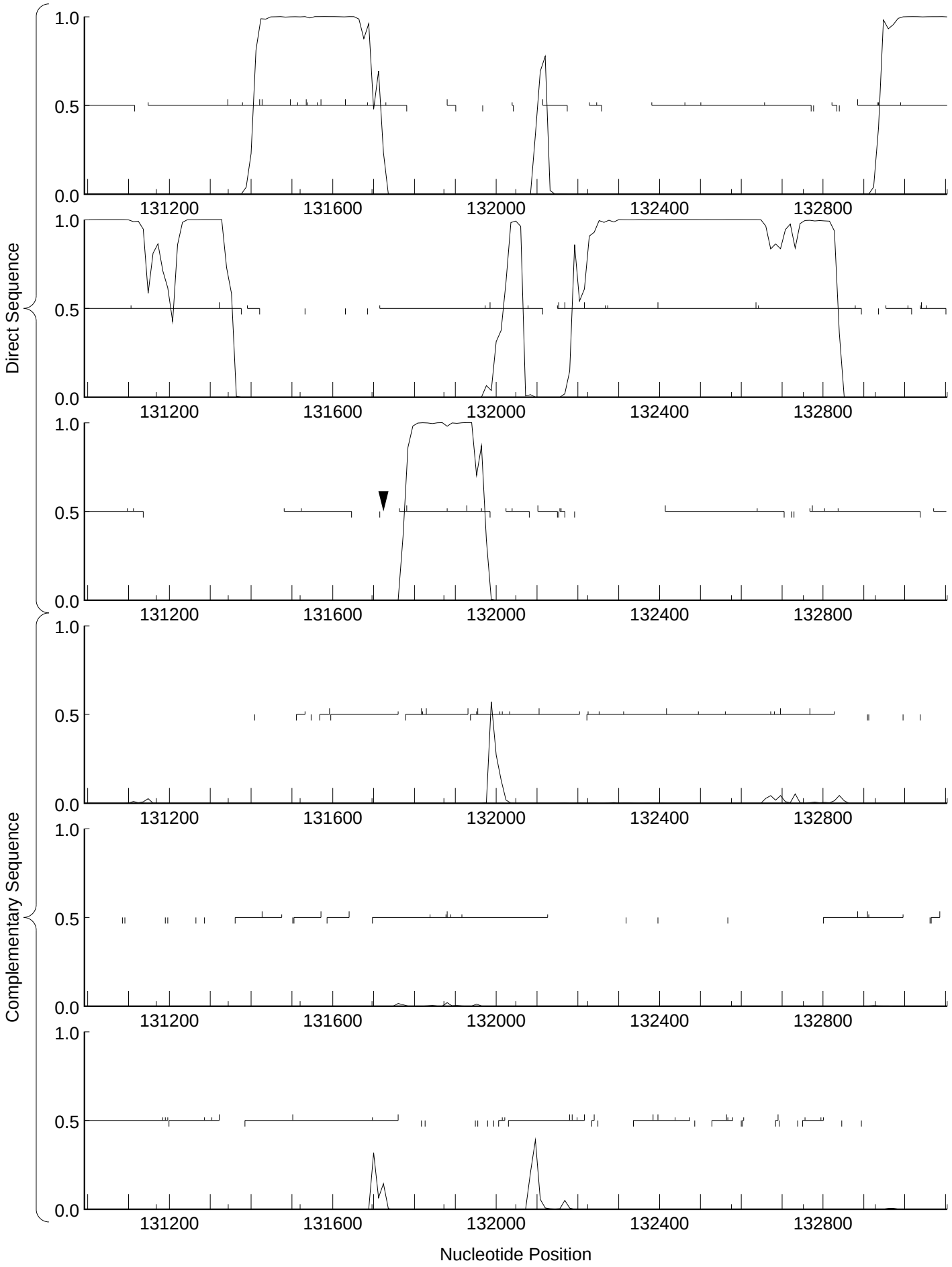
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 62/75



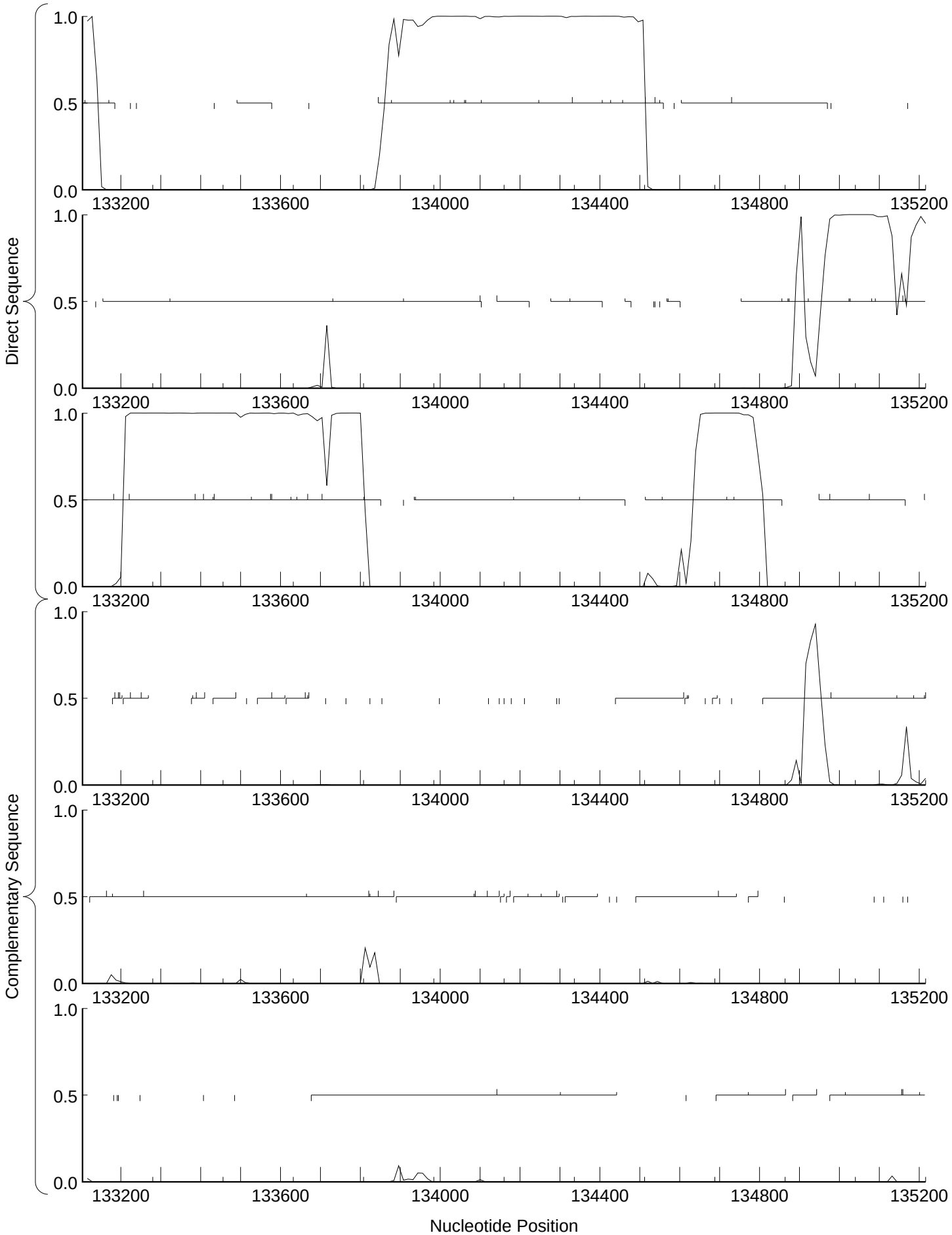
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 63/75



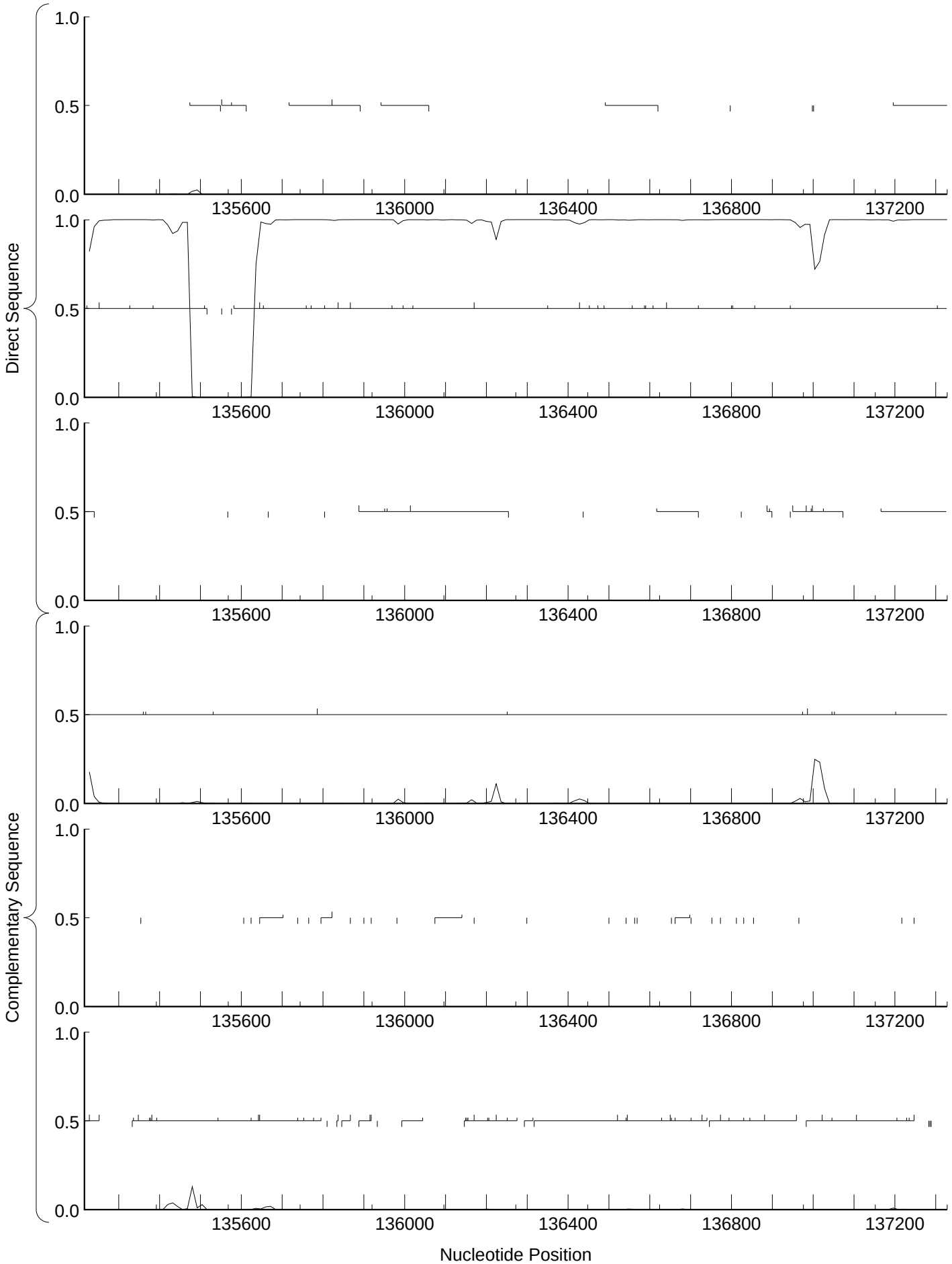
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 64/75



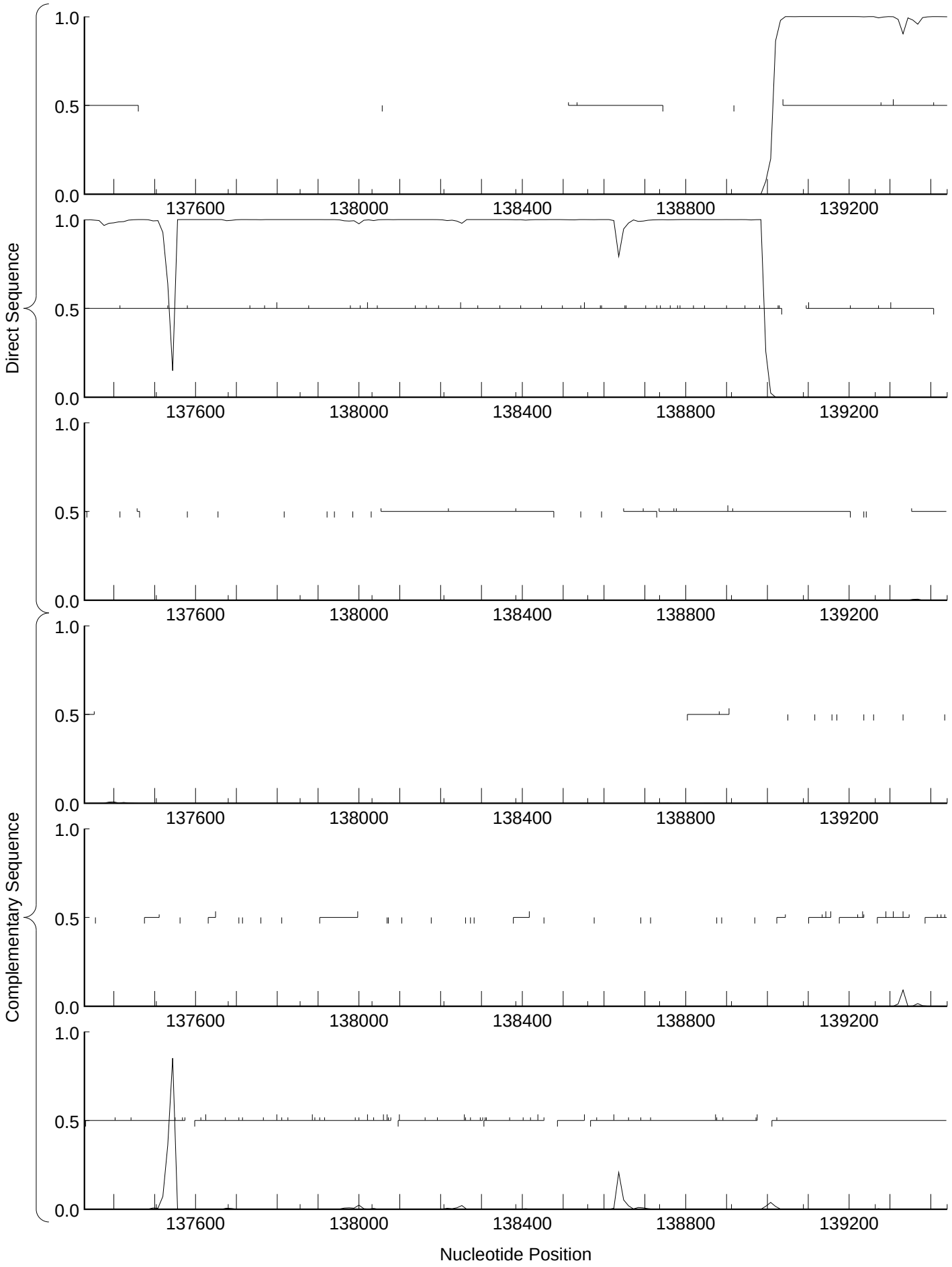
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 65/75



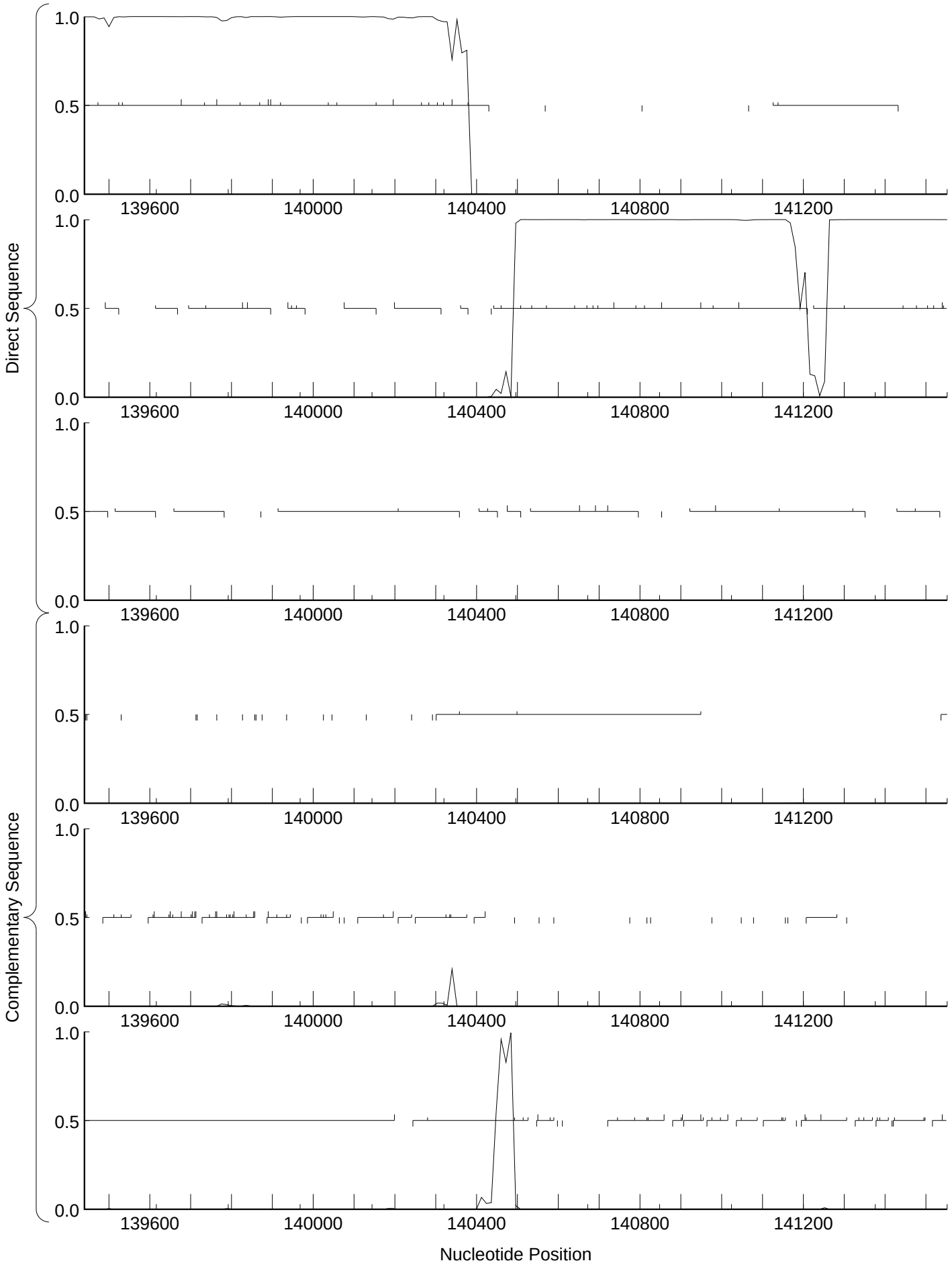
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 66/75



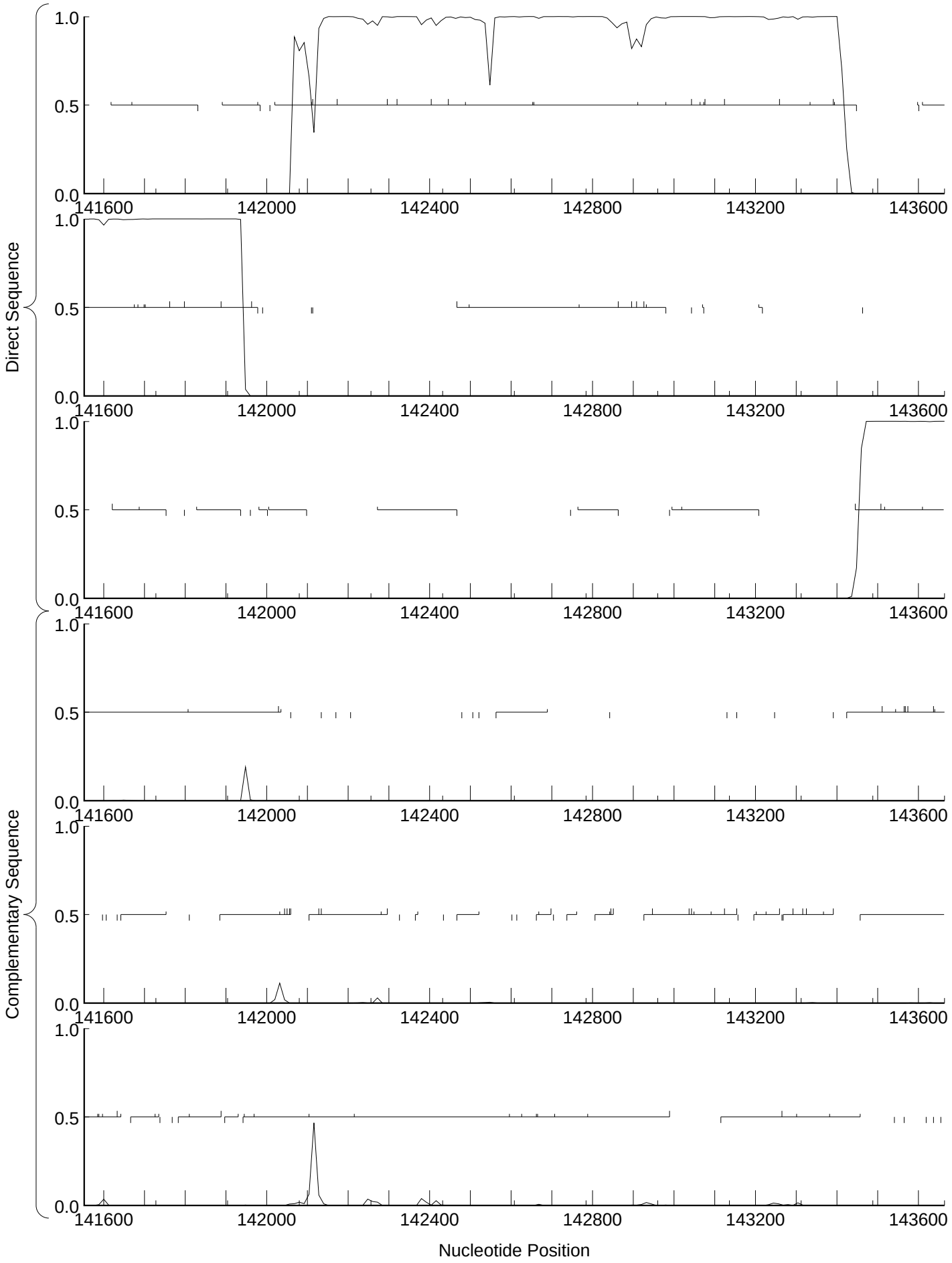
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 67/75



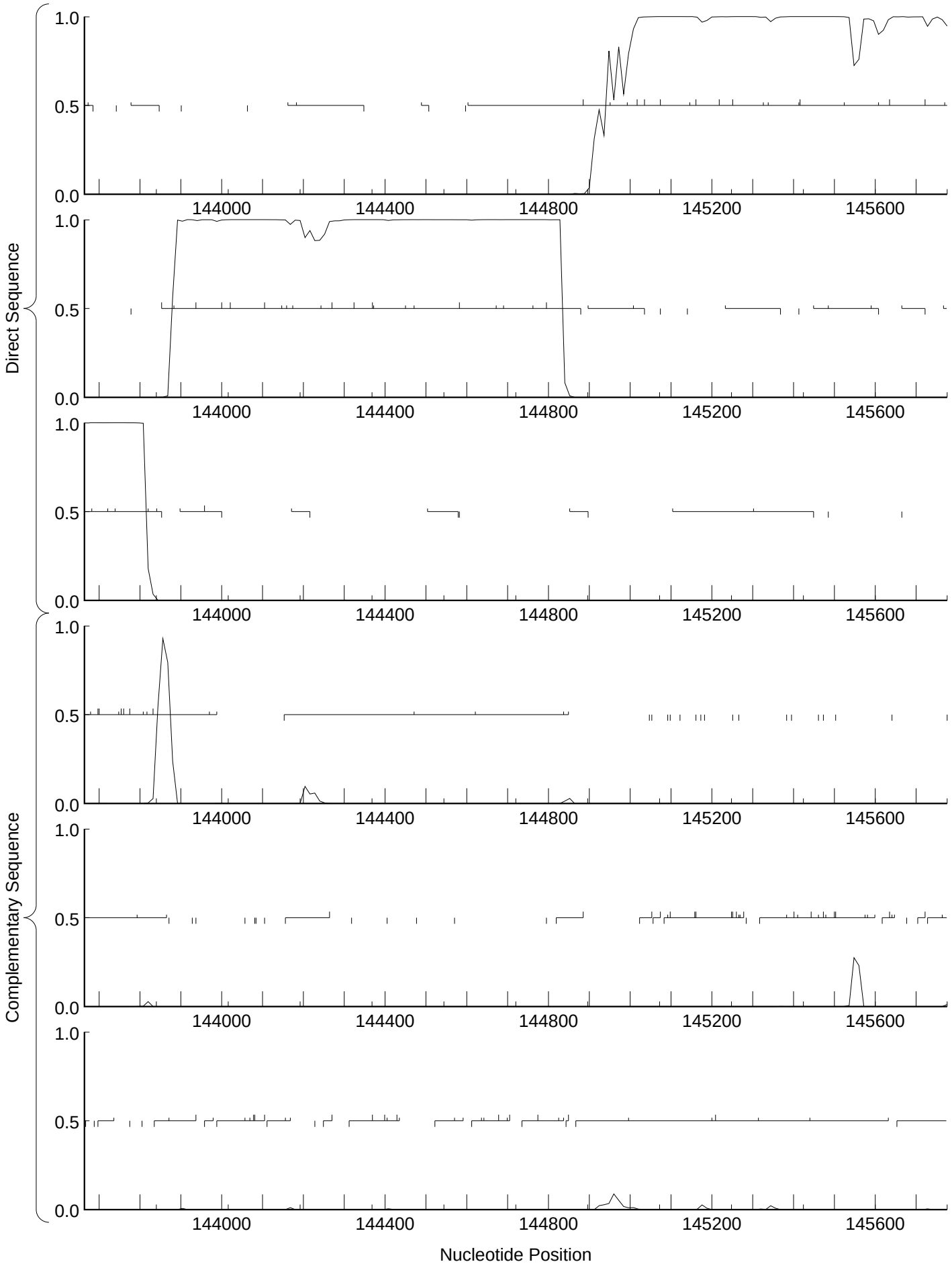
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 68/75



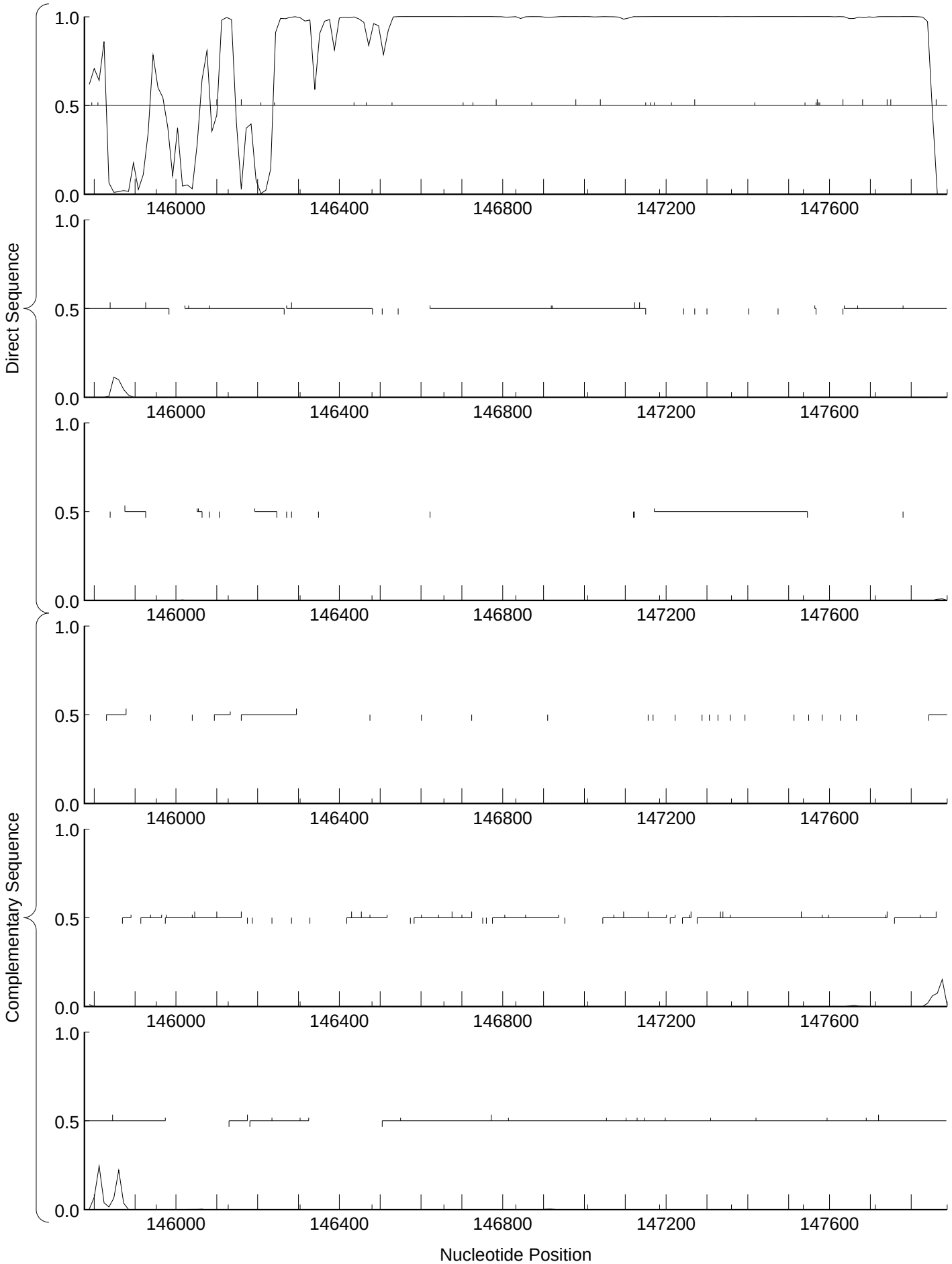
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 69/75



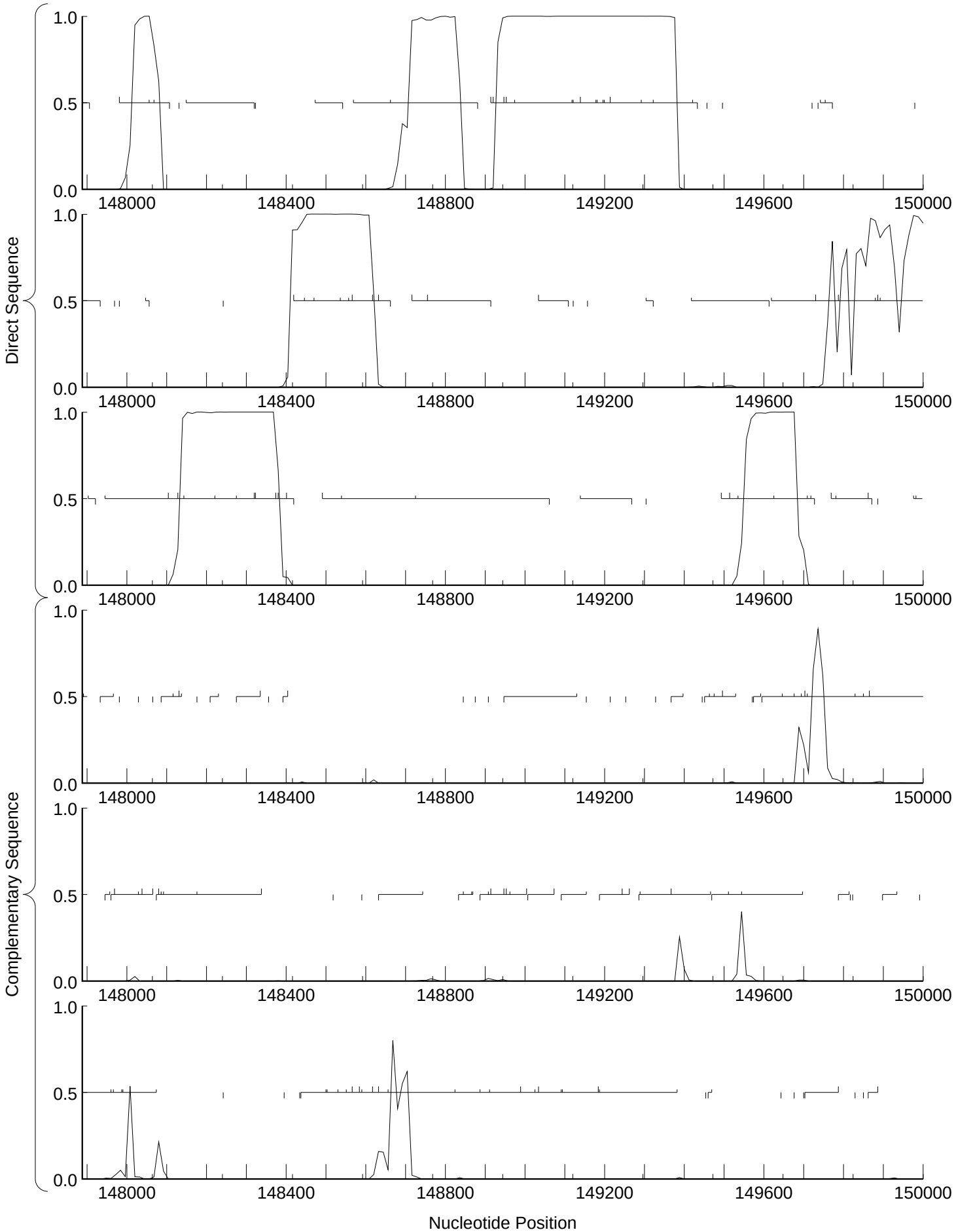
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 70/75



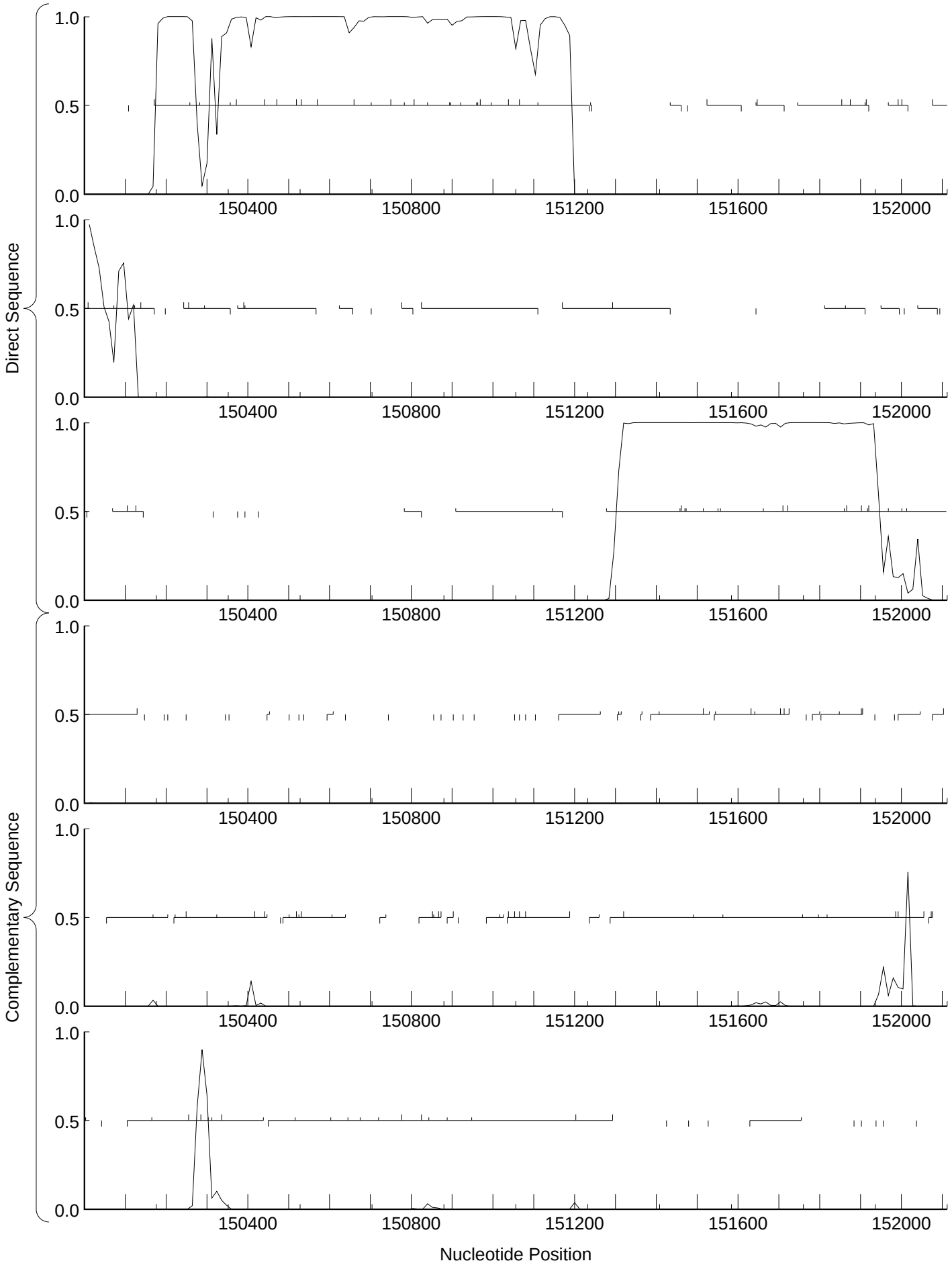
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 71/75



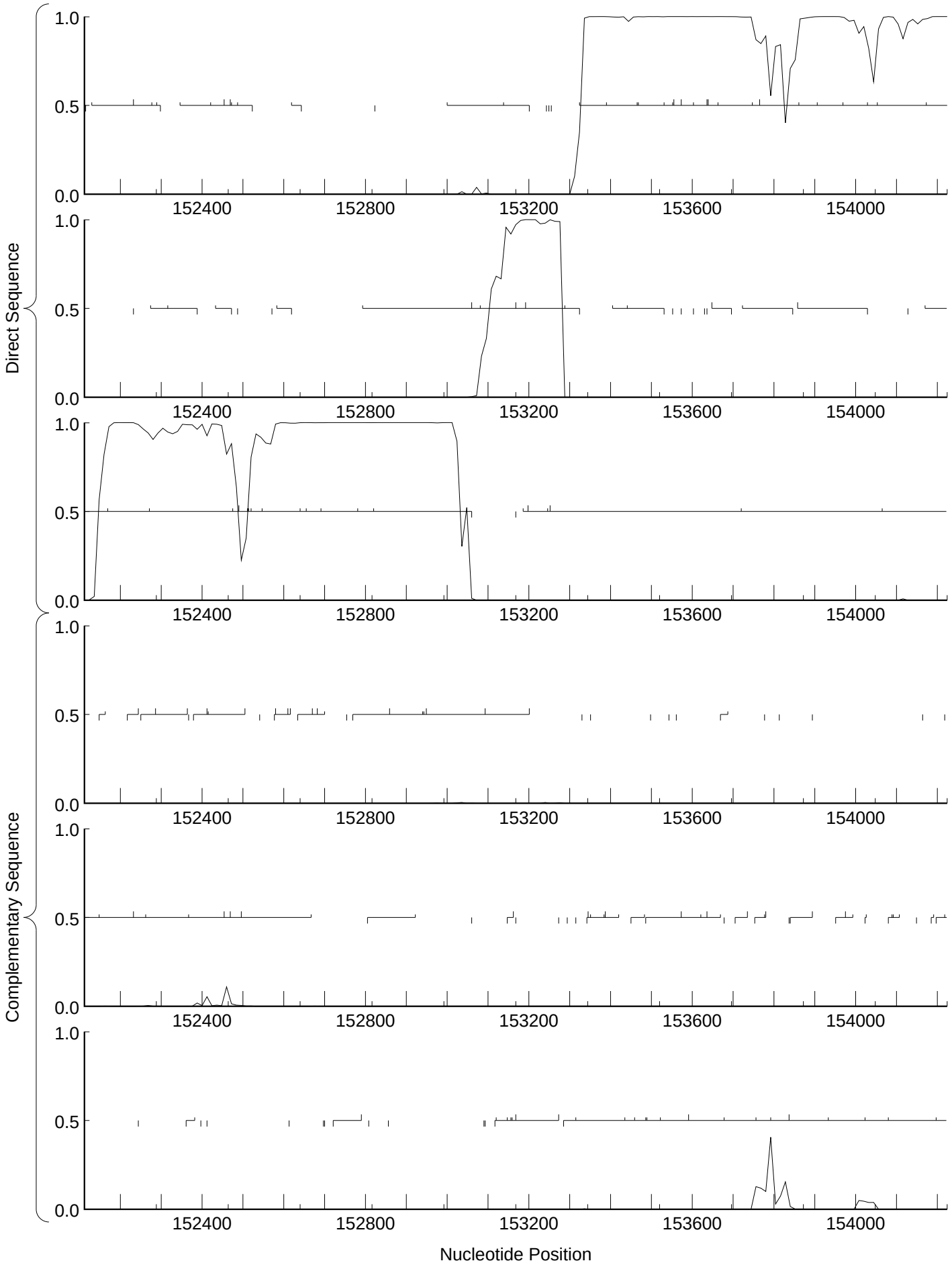
cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 72/75



cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 73/75



cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 74/75



cterium phage BadAgartude complete sequence, 155658 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 75/75

