GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ
Analysis Date    :  3/28/24 at 18:58:47
Pages            :  23
Sequence Length  :  45507 bp
GC Content       :   64.50%

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

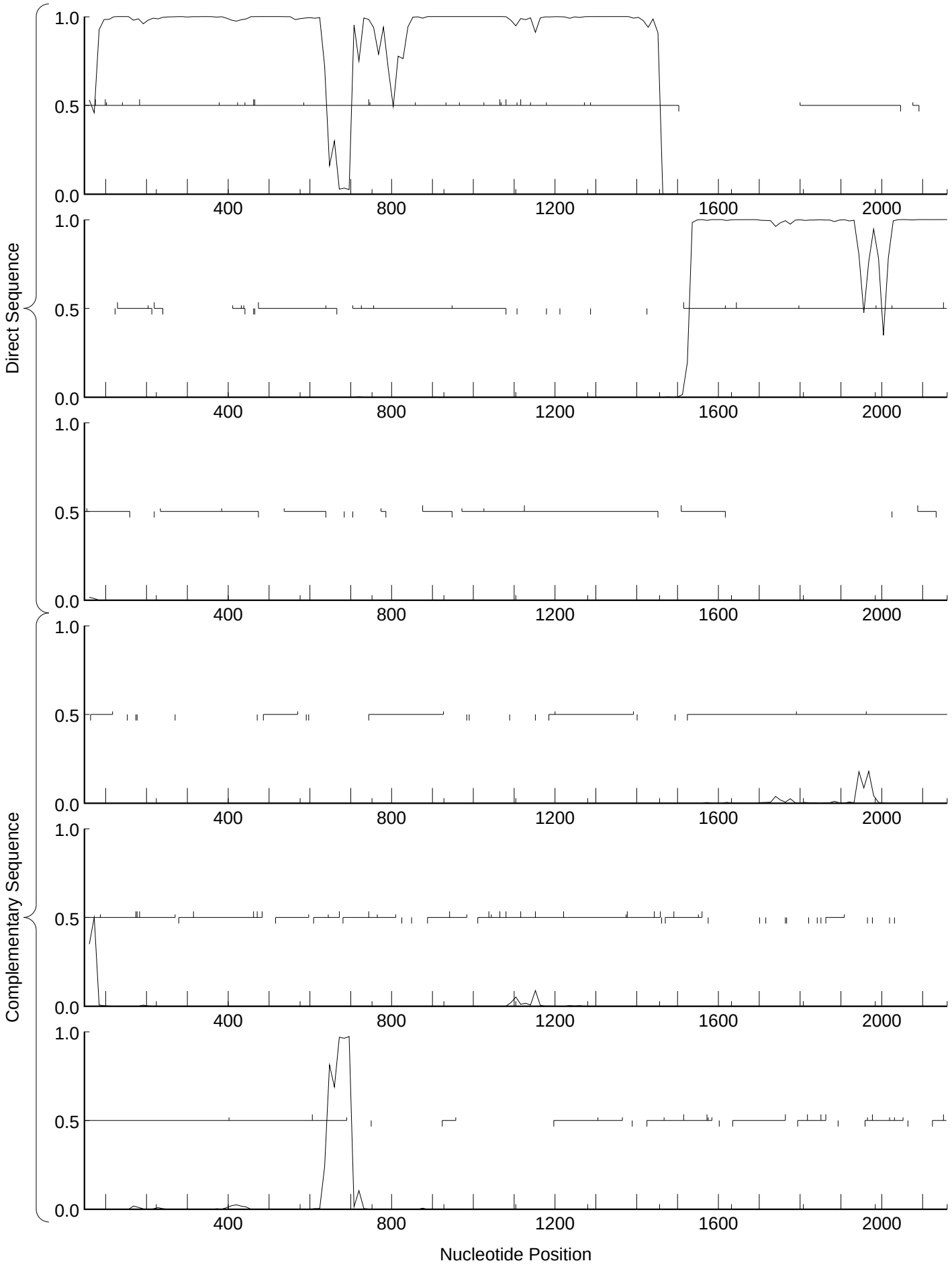
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Order      :  4
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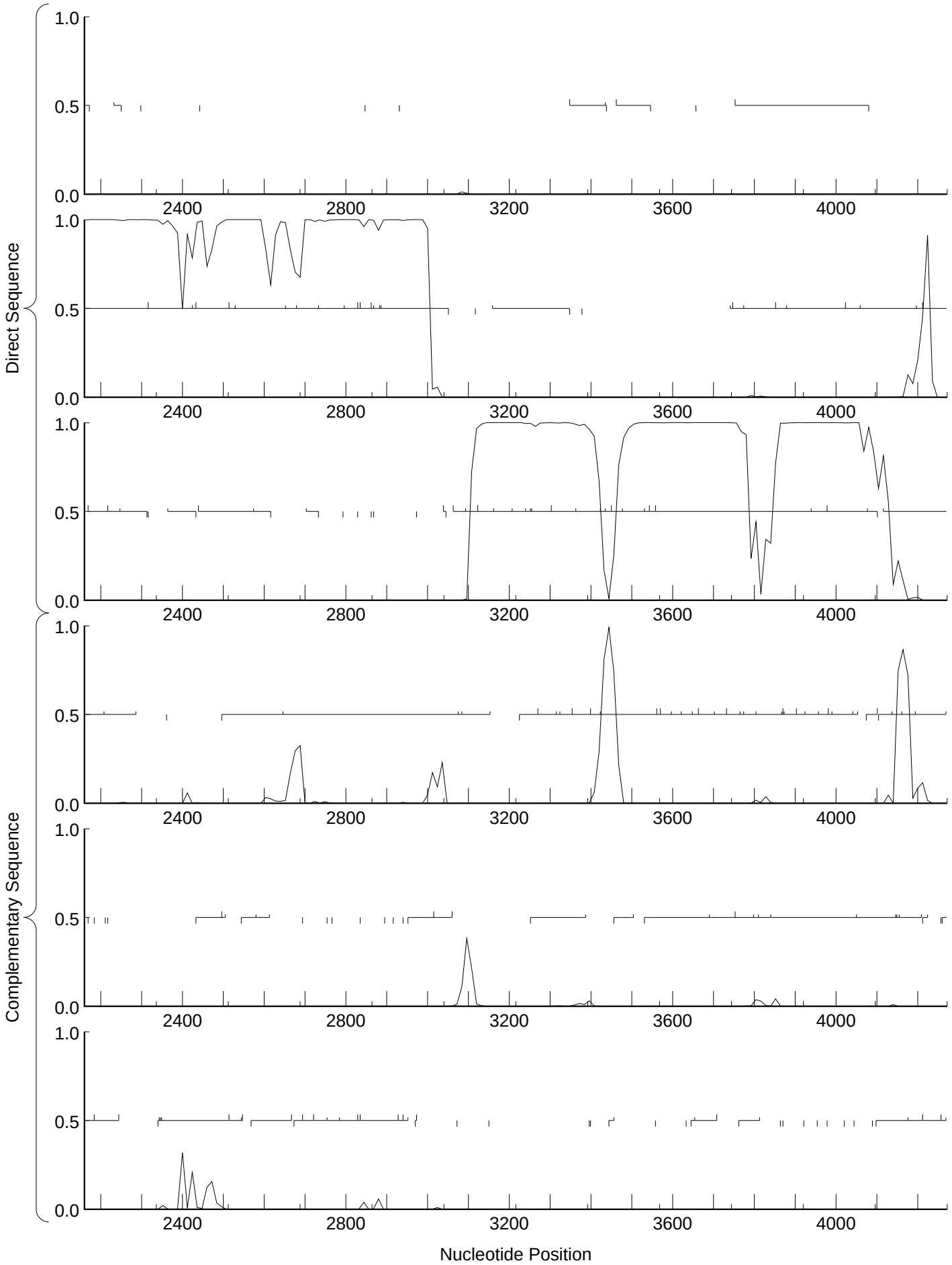

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

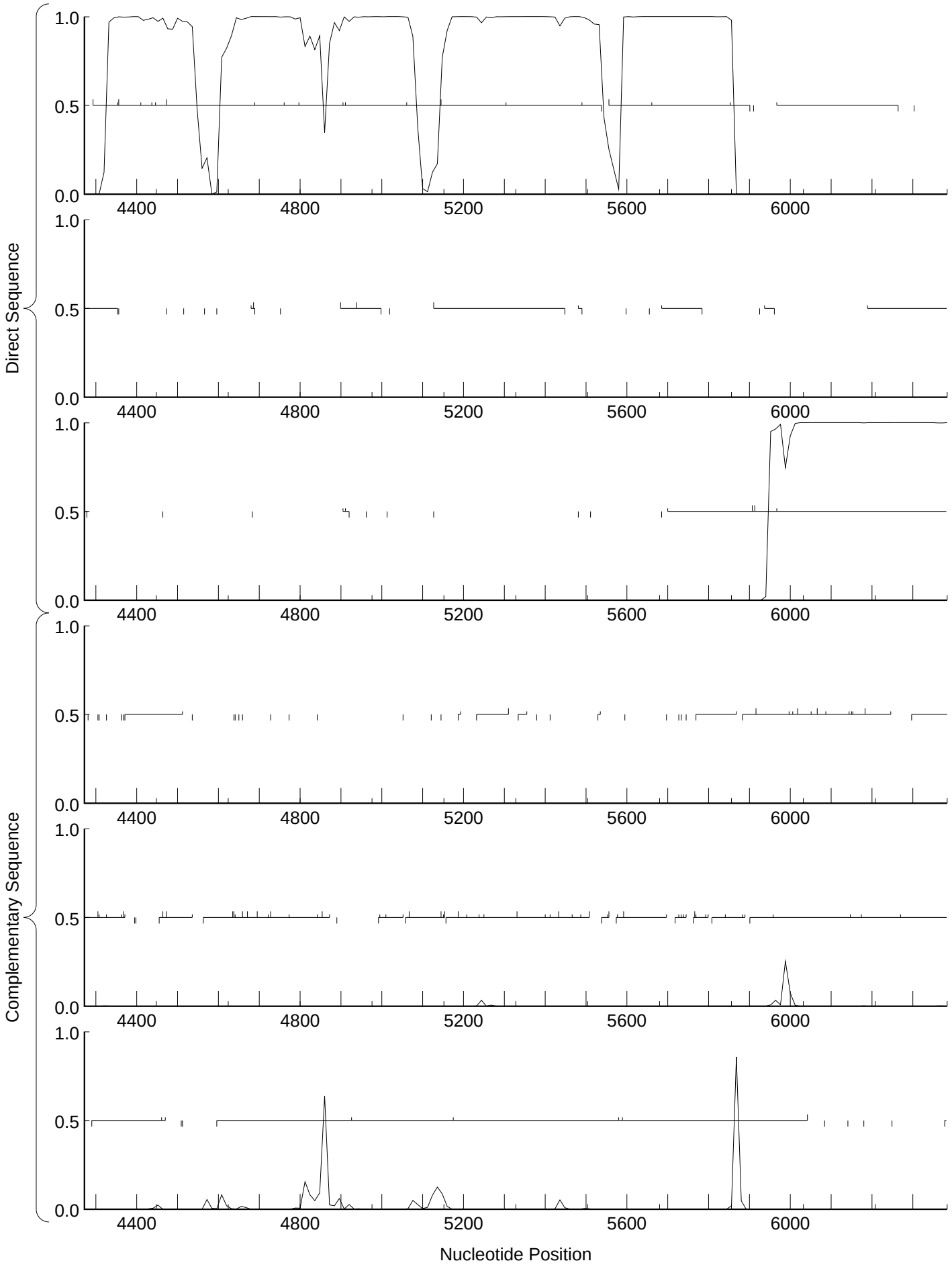
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 2/23



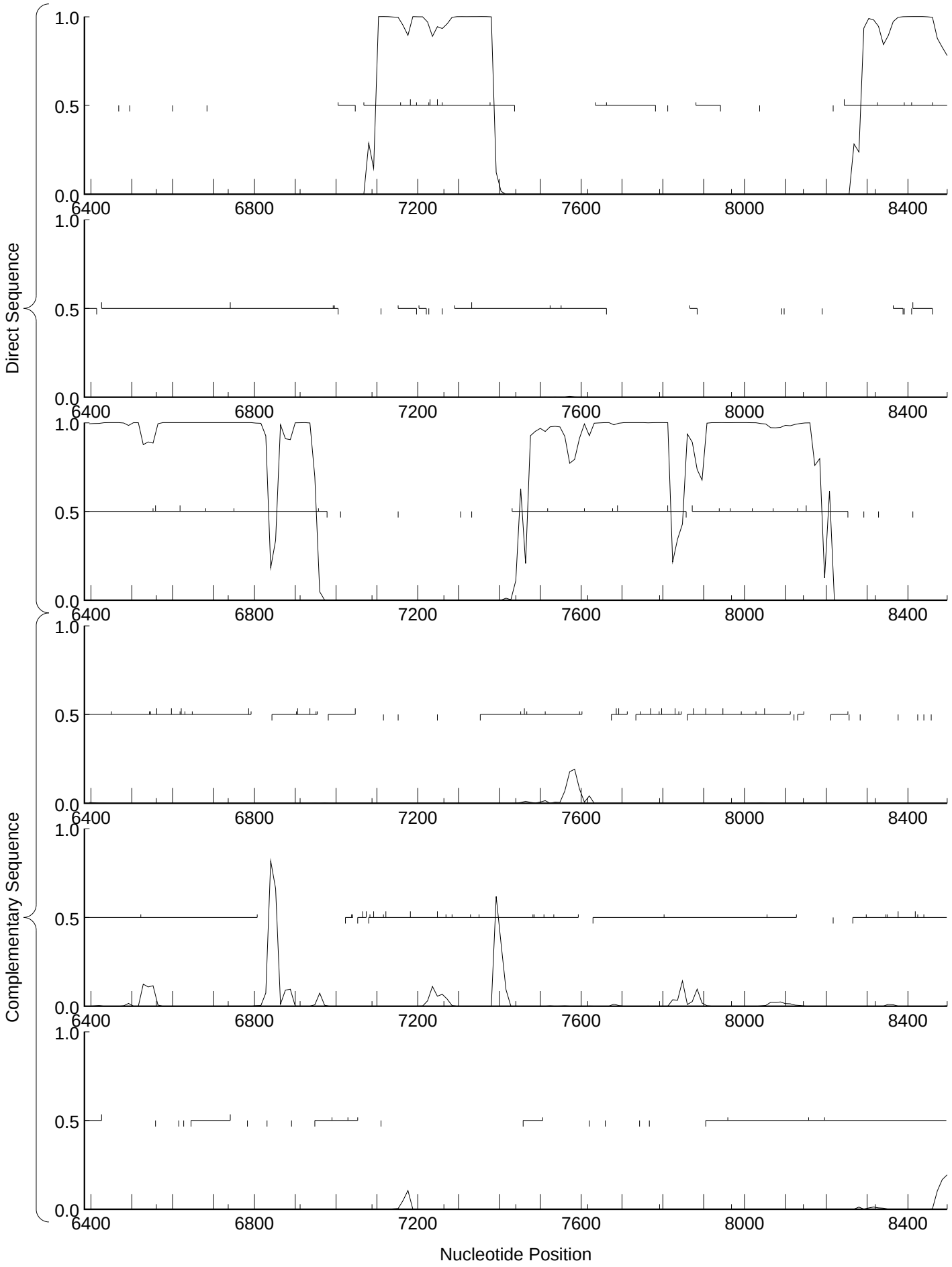
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 3/23



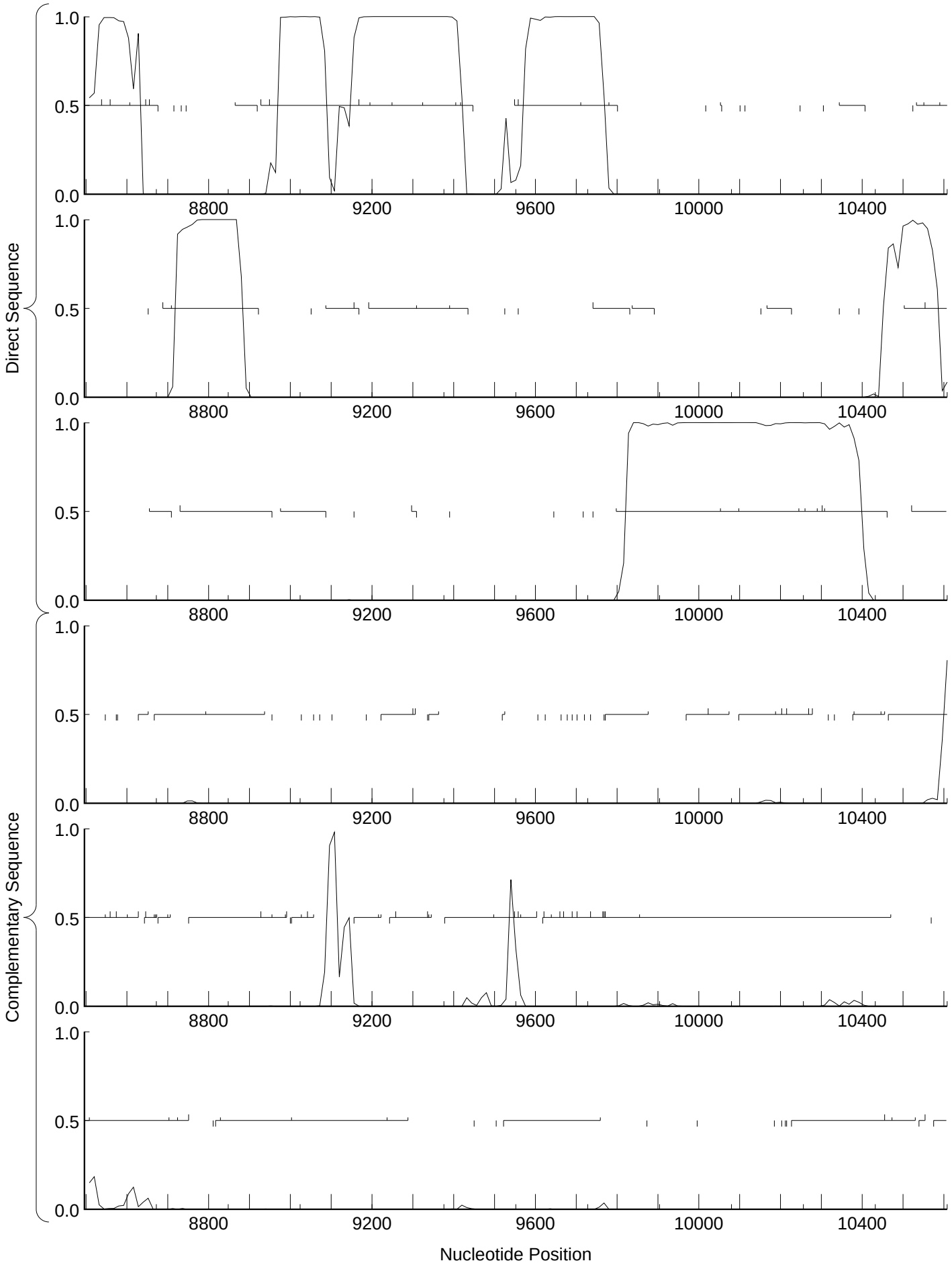
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 4/23



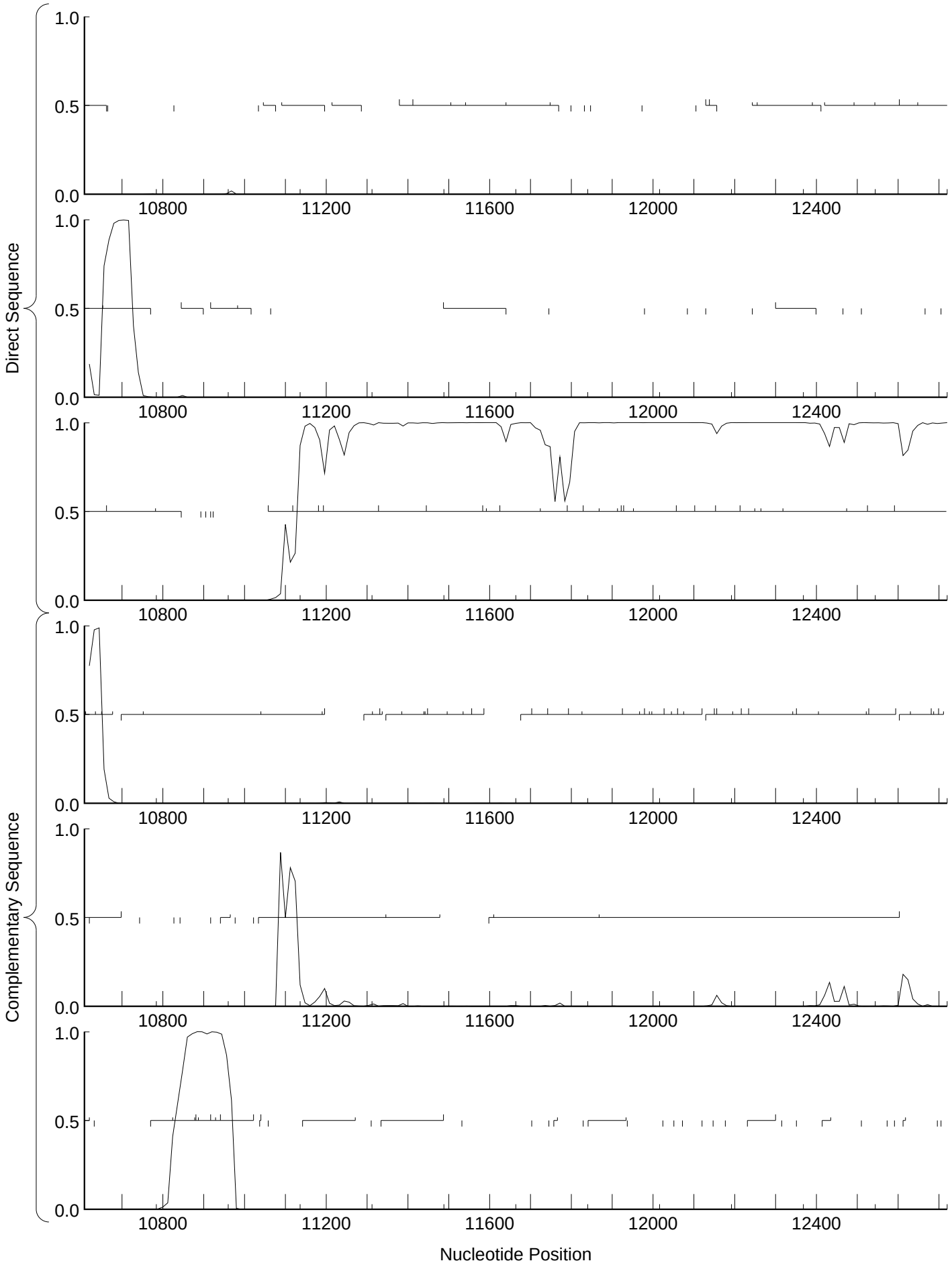
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 5/23



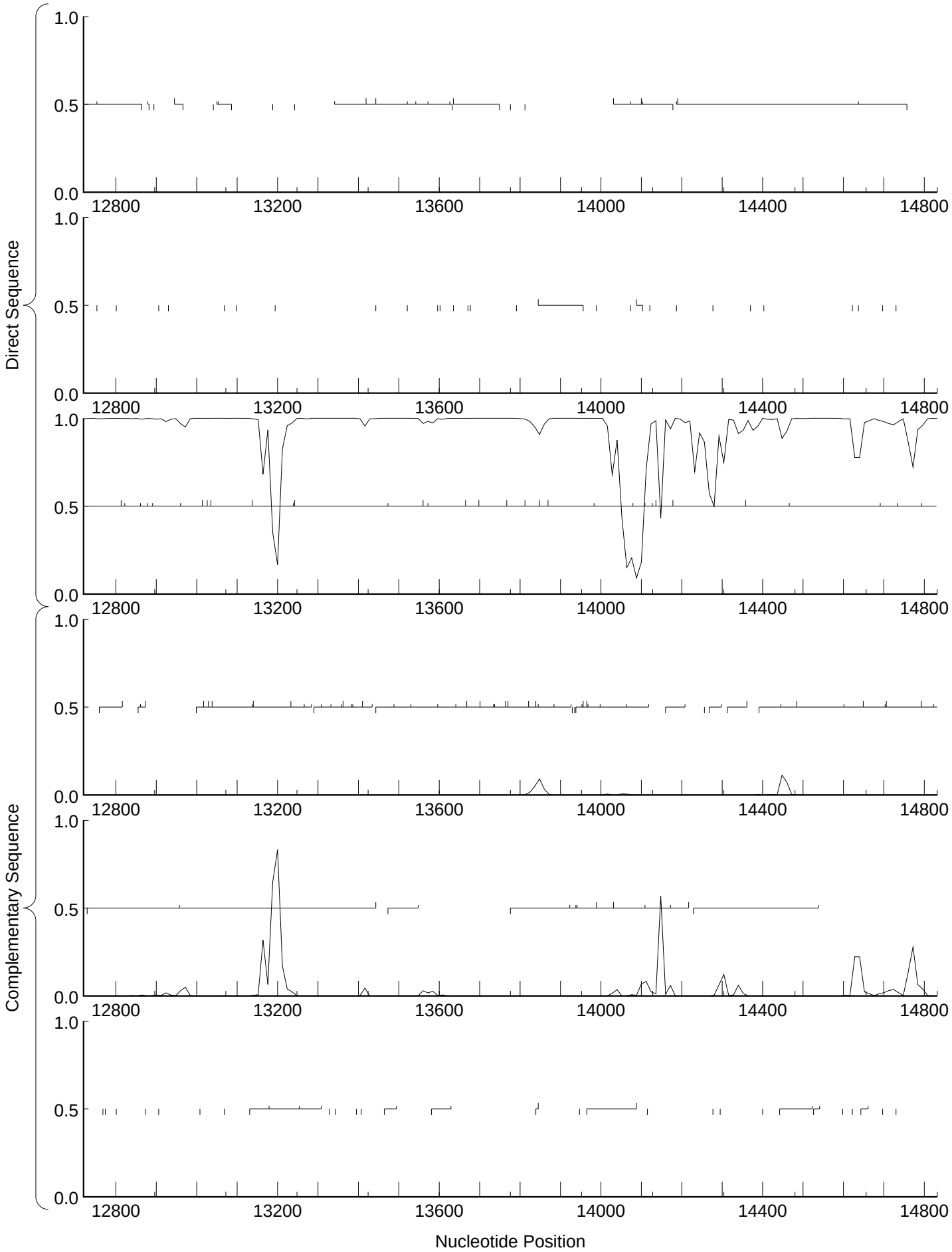
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 6/23



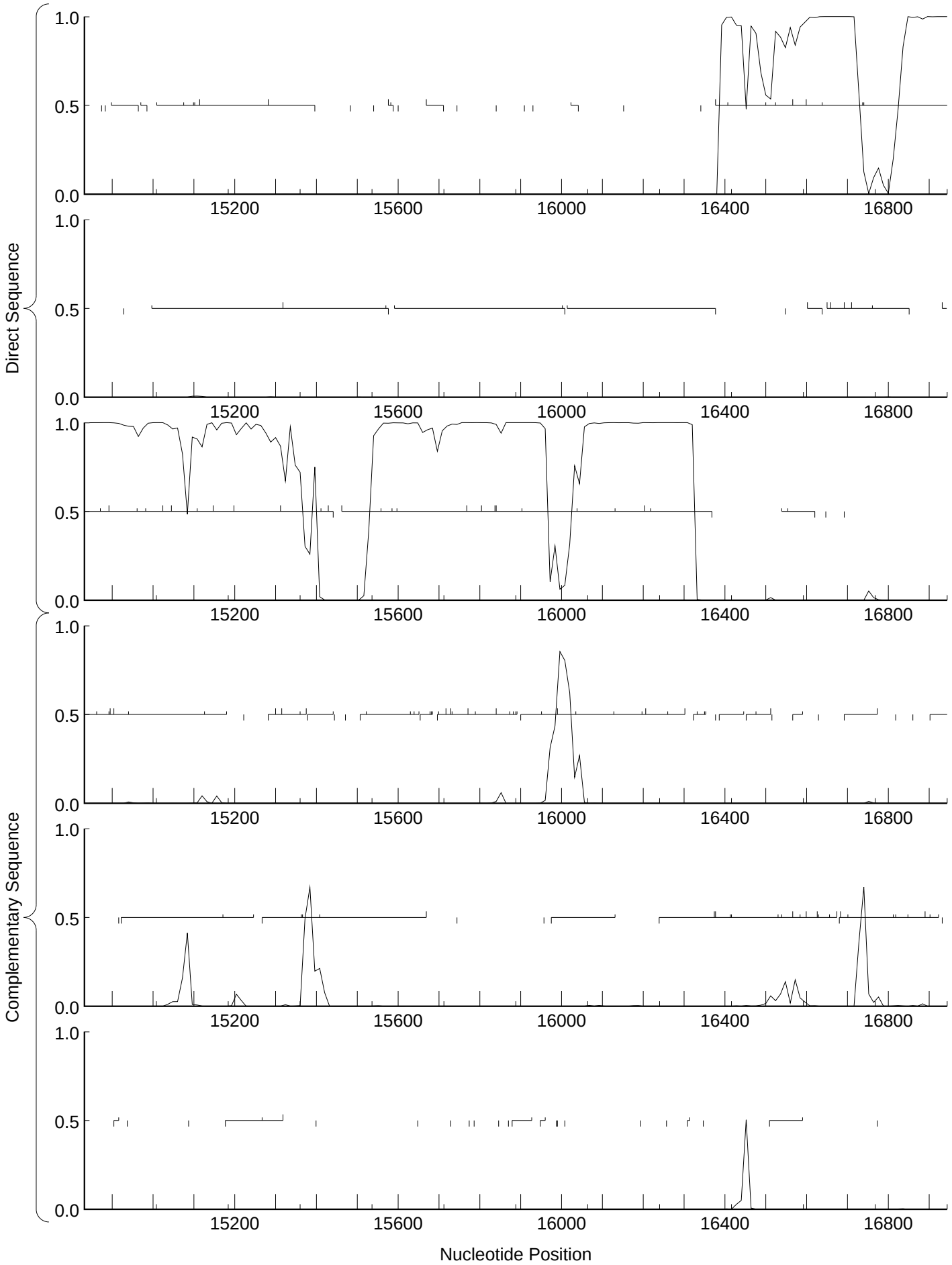
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 7/23



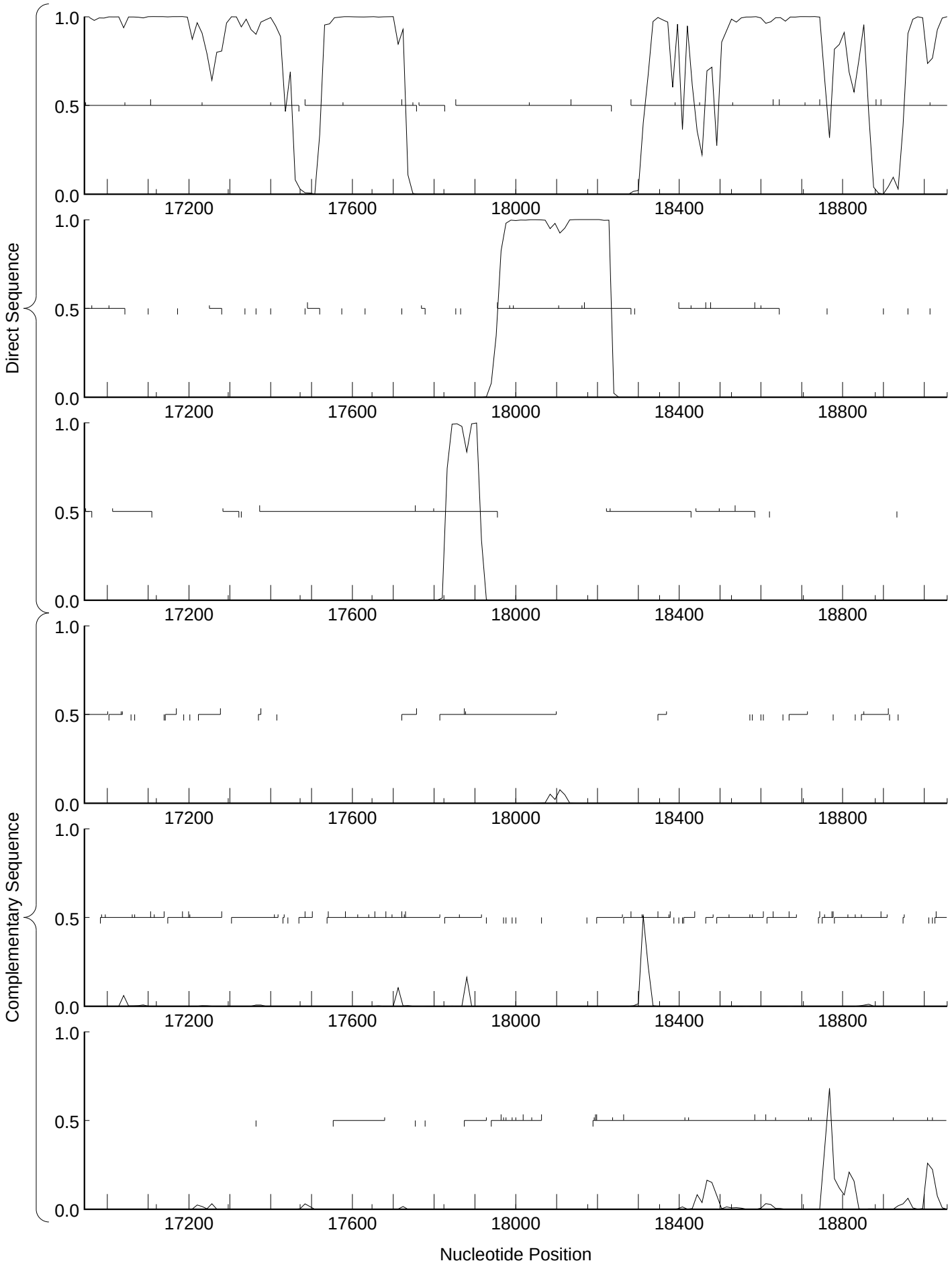
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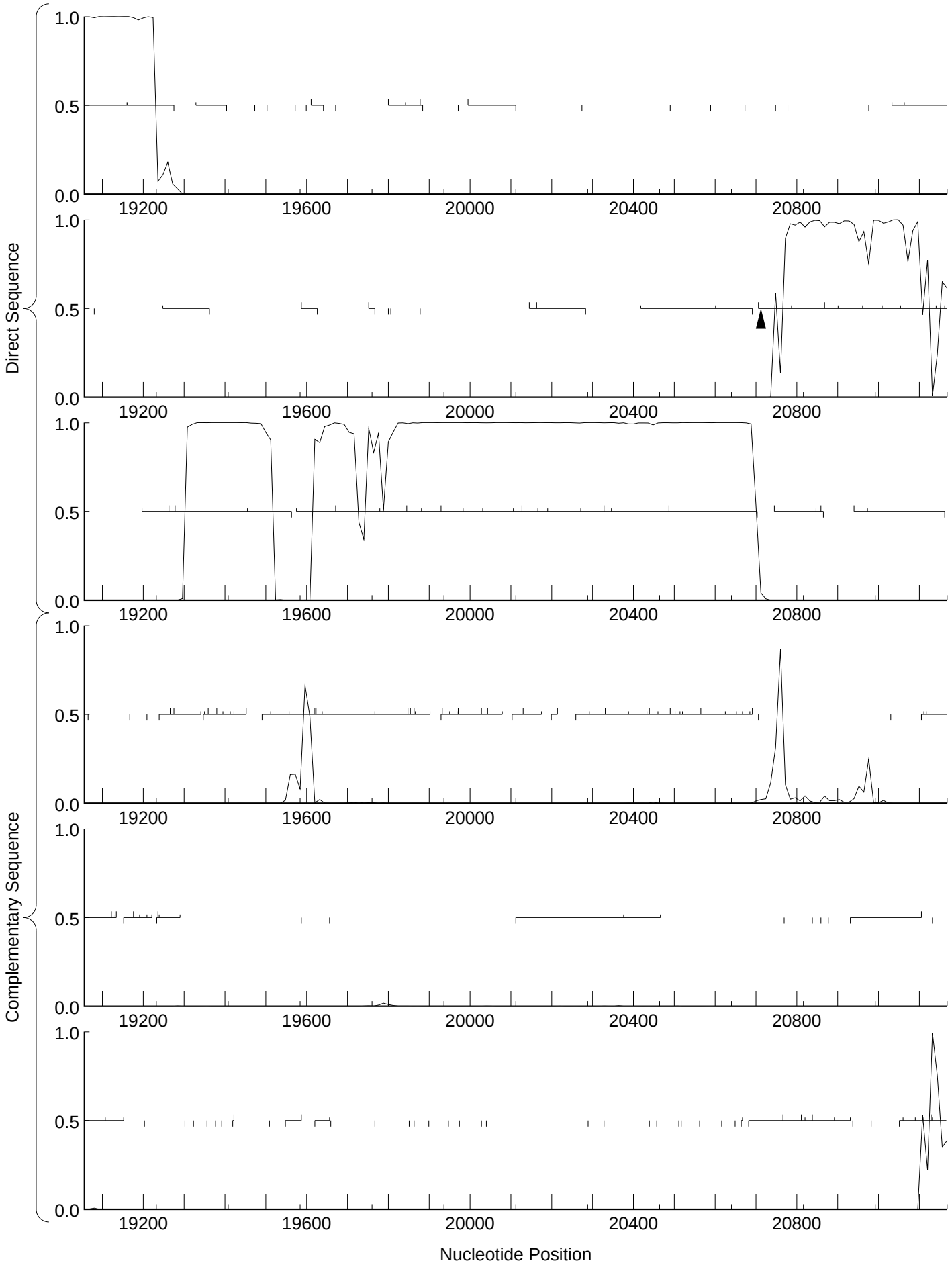
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 9/23



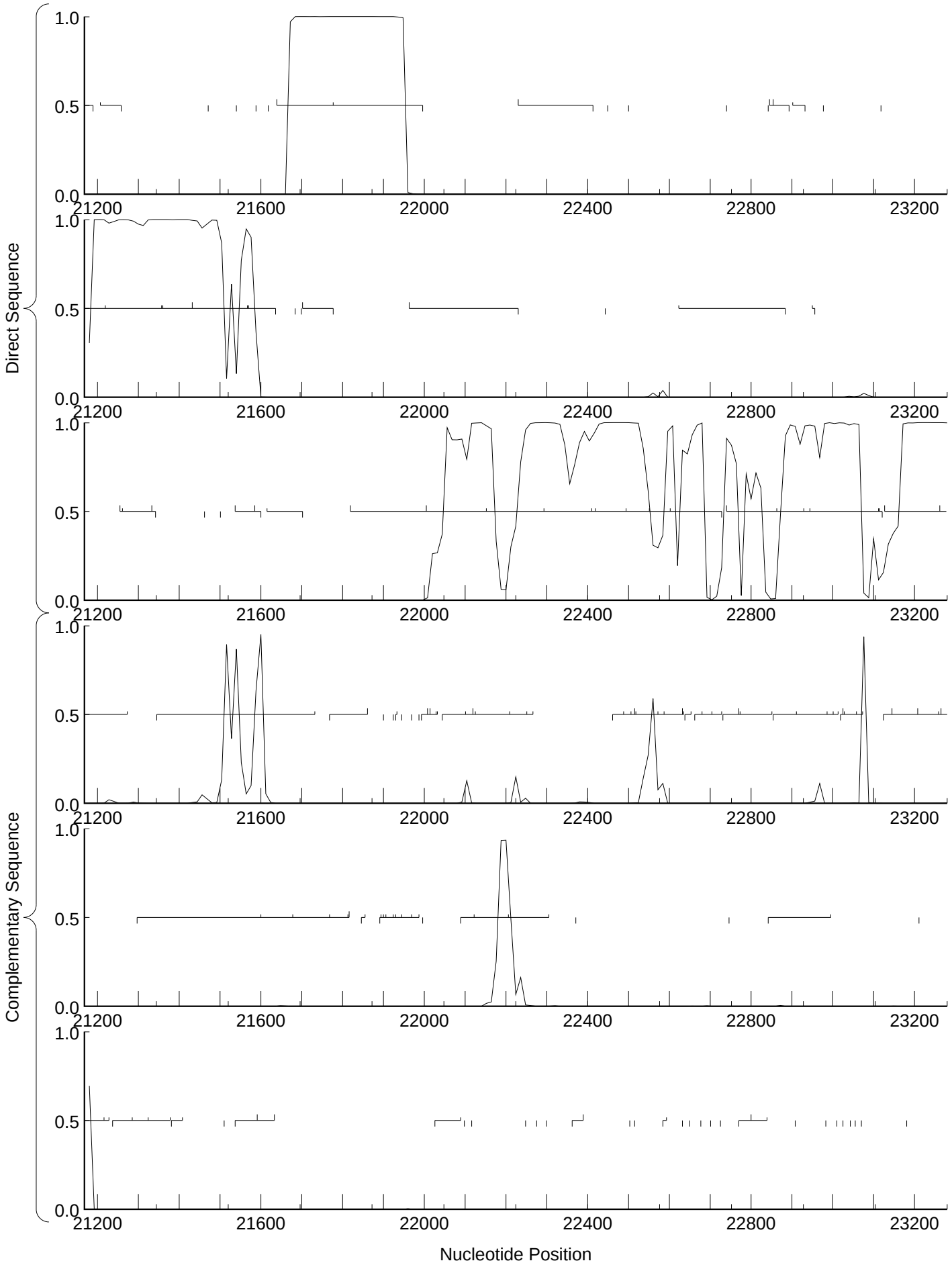
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 10/23



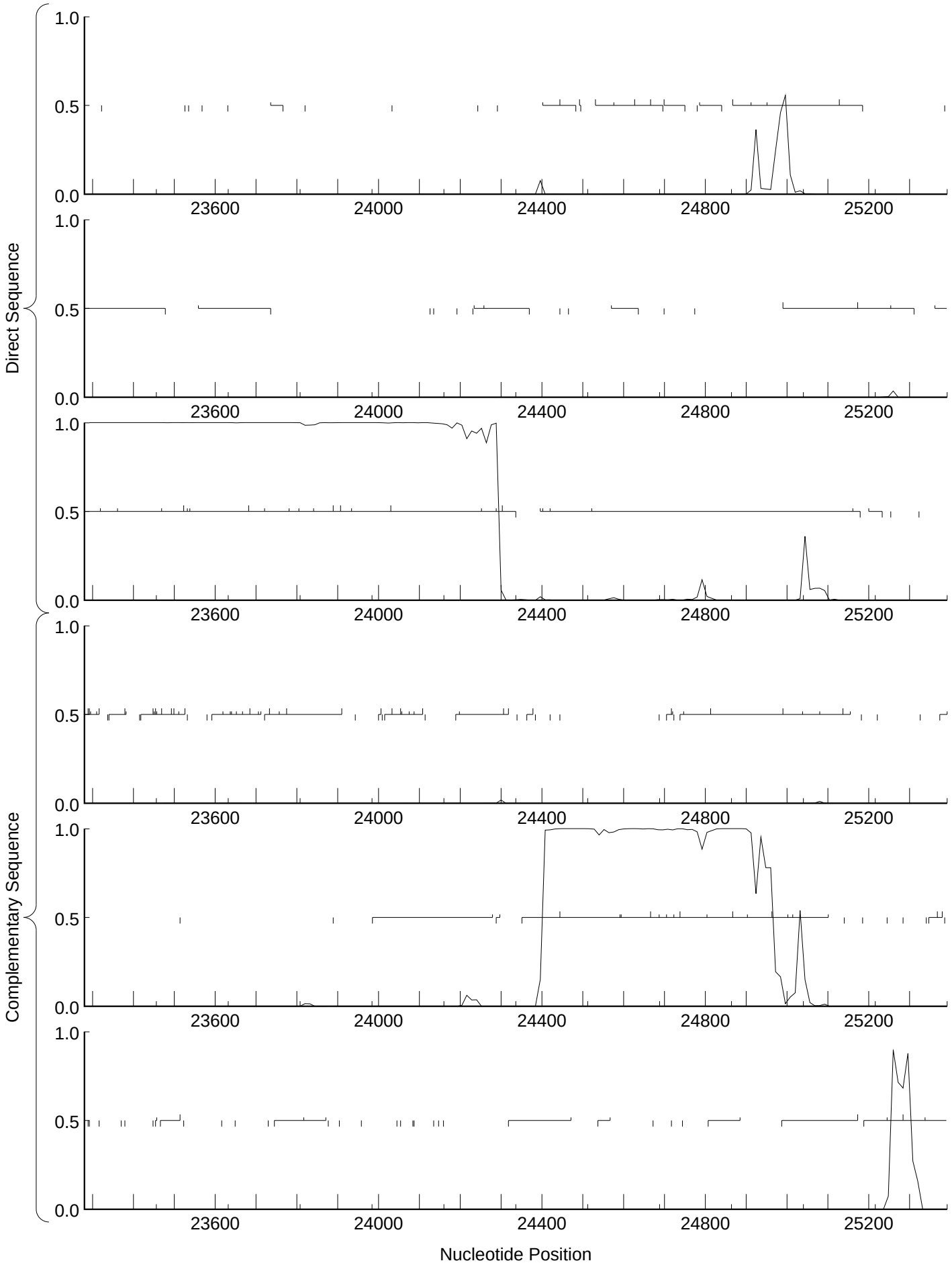
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 11/23



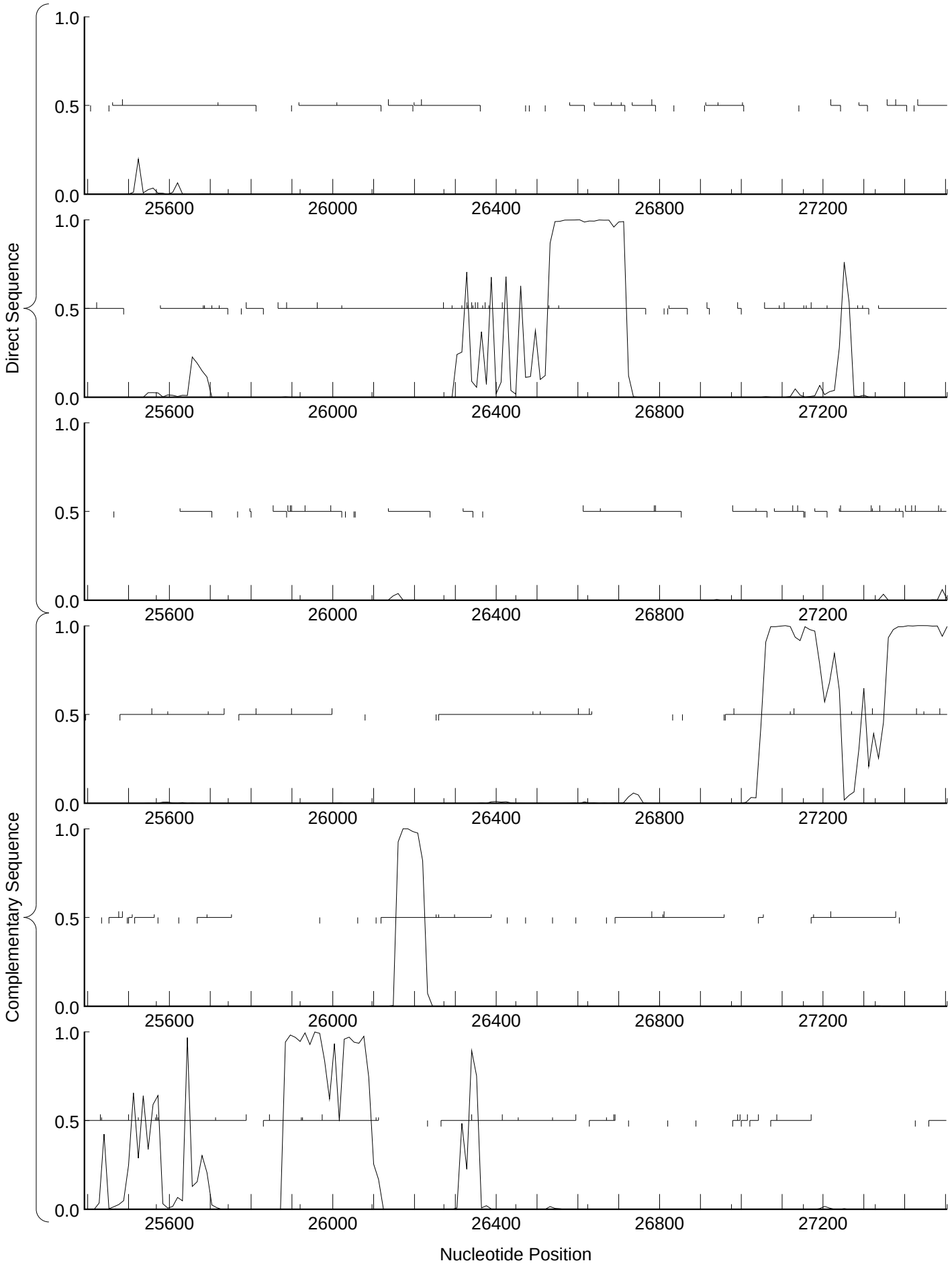
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 12/23



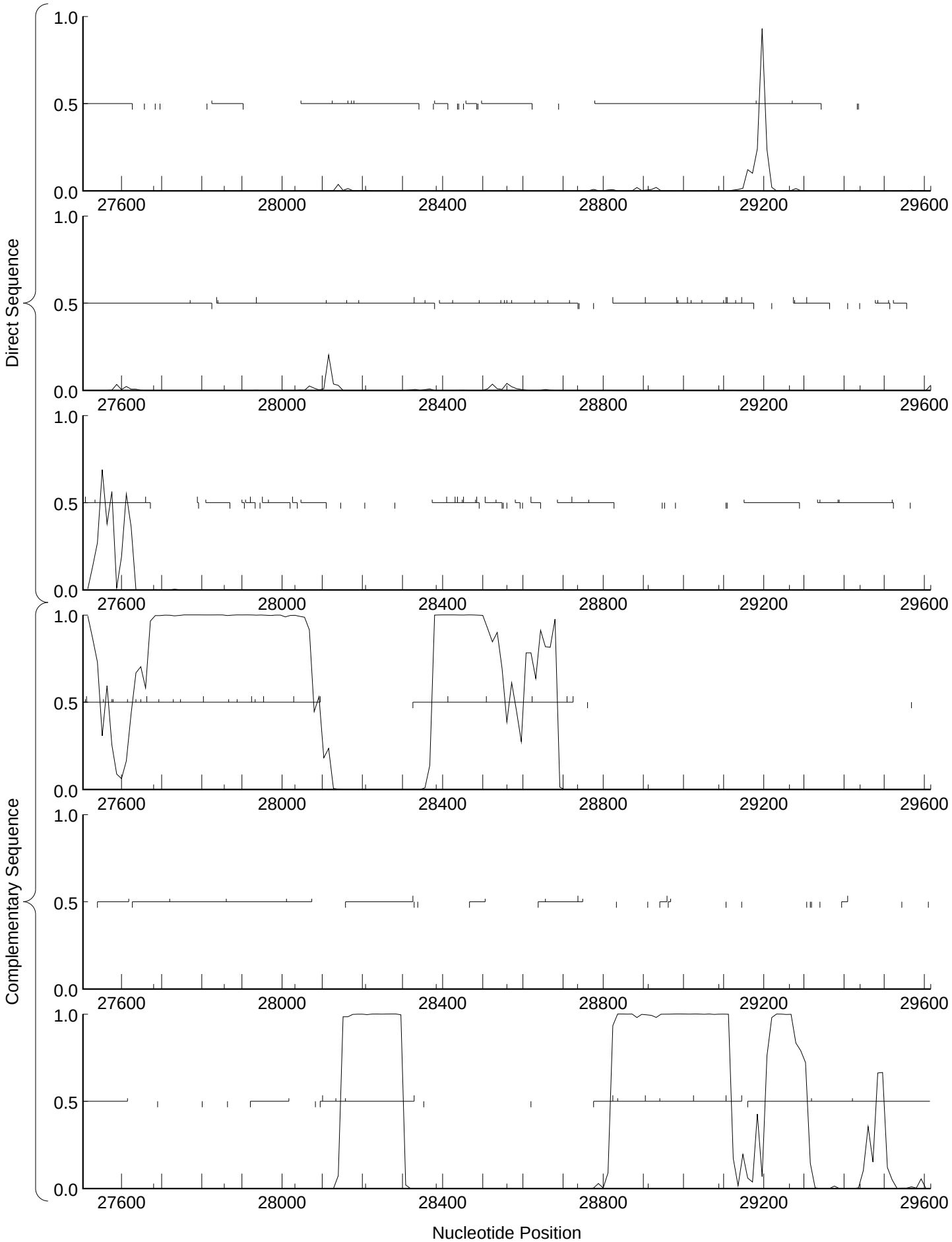
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 13/23



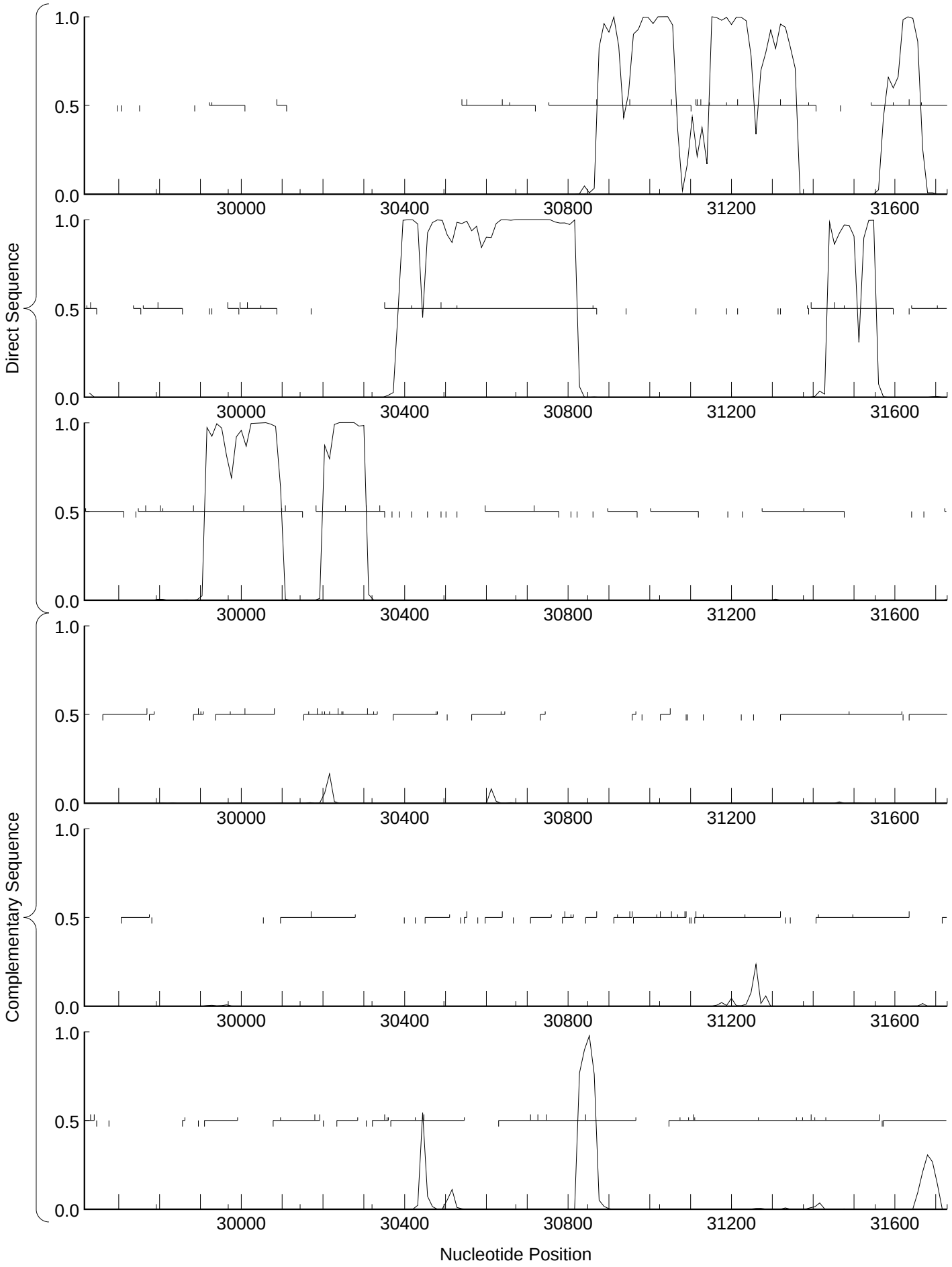
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 14/23



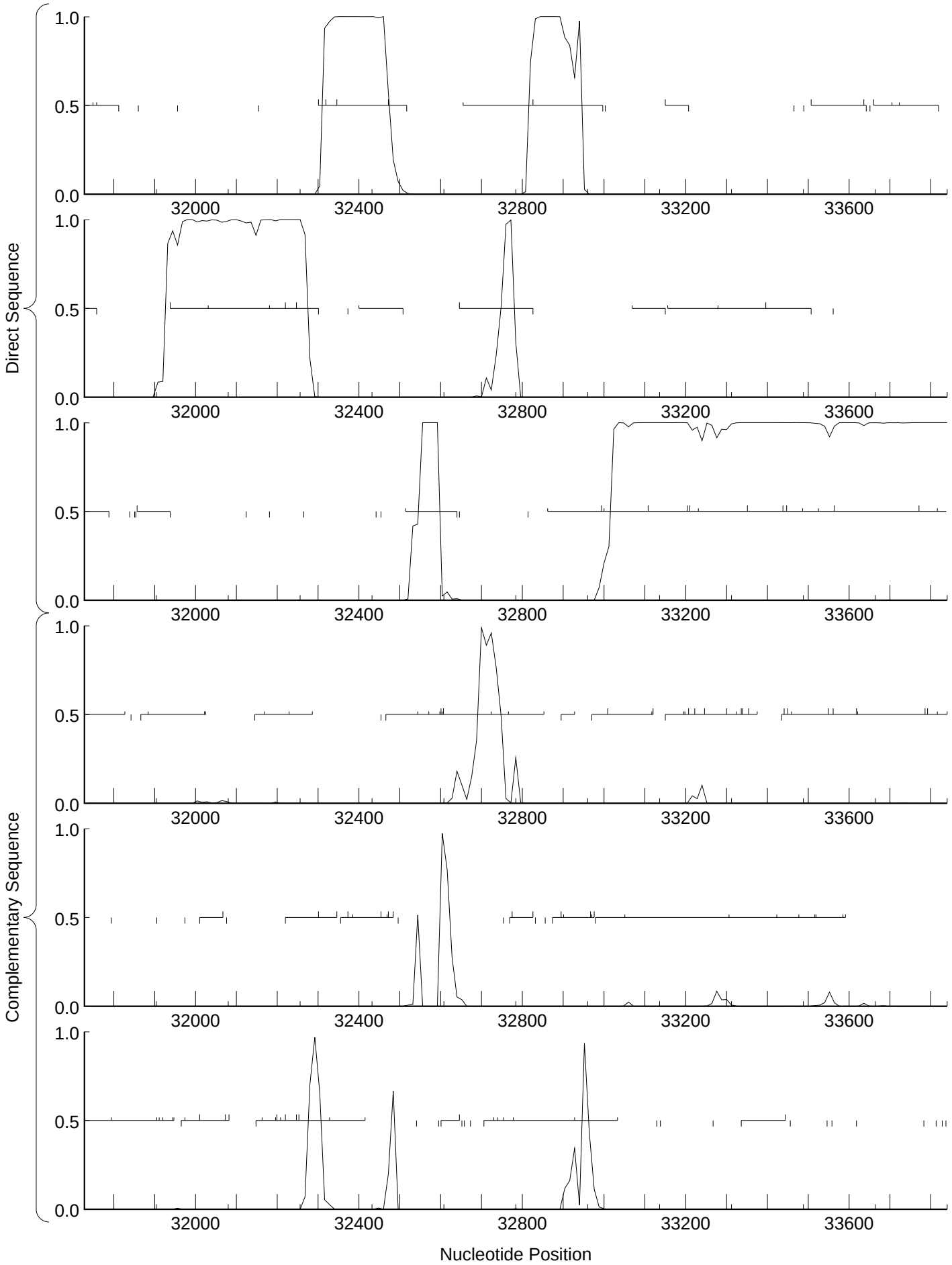
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 15/23



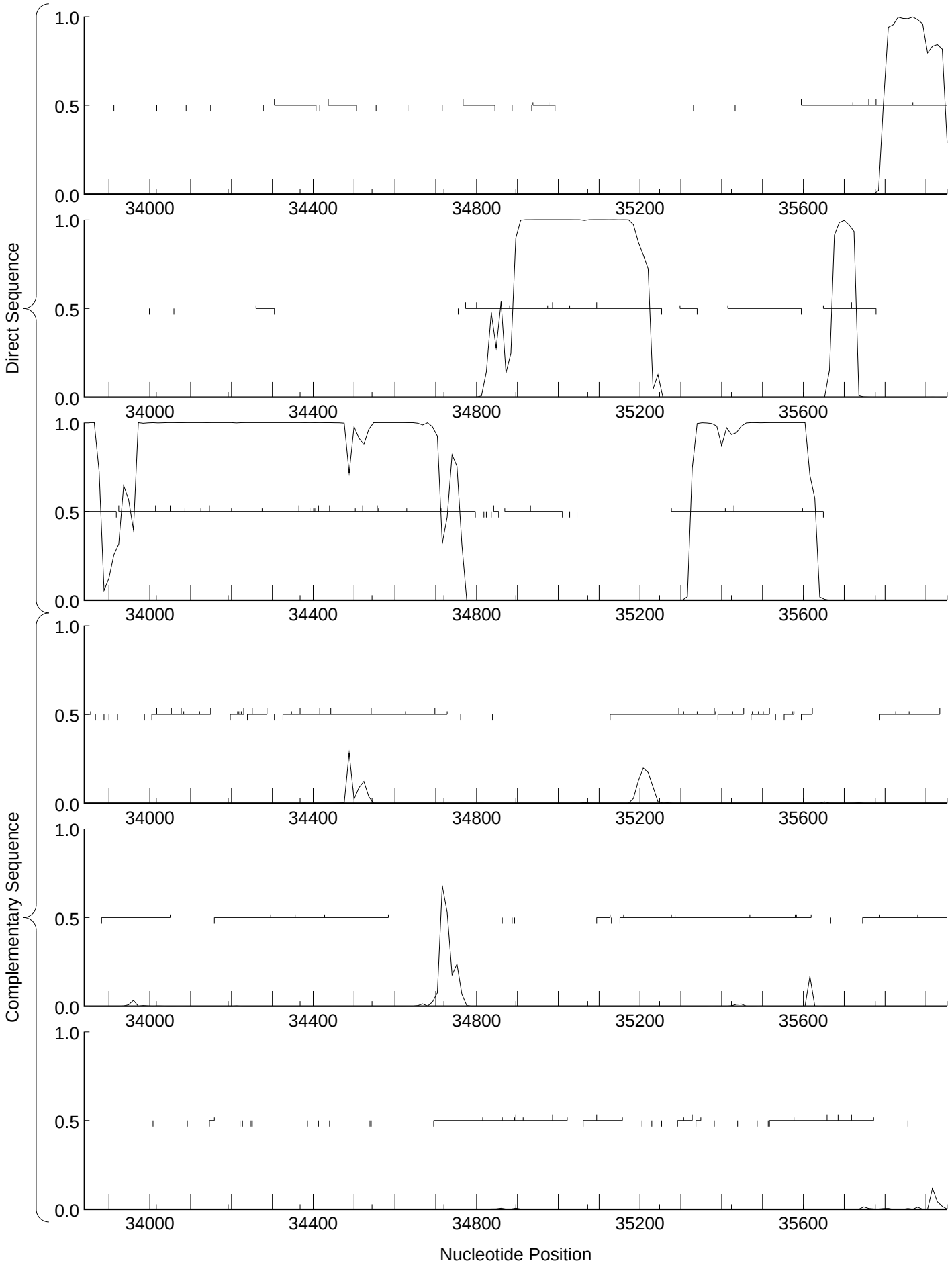
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 16/23



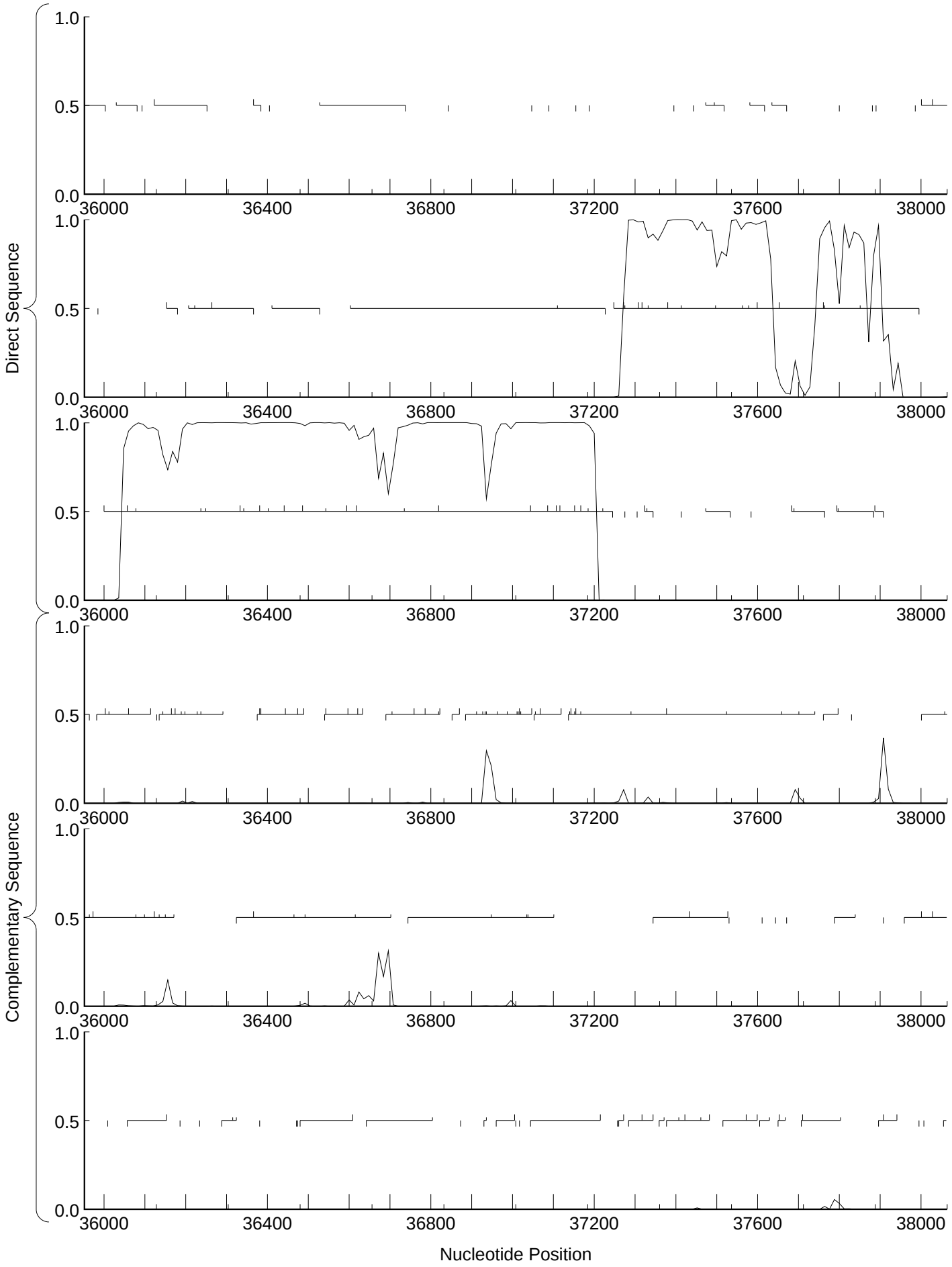
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 17/23



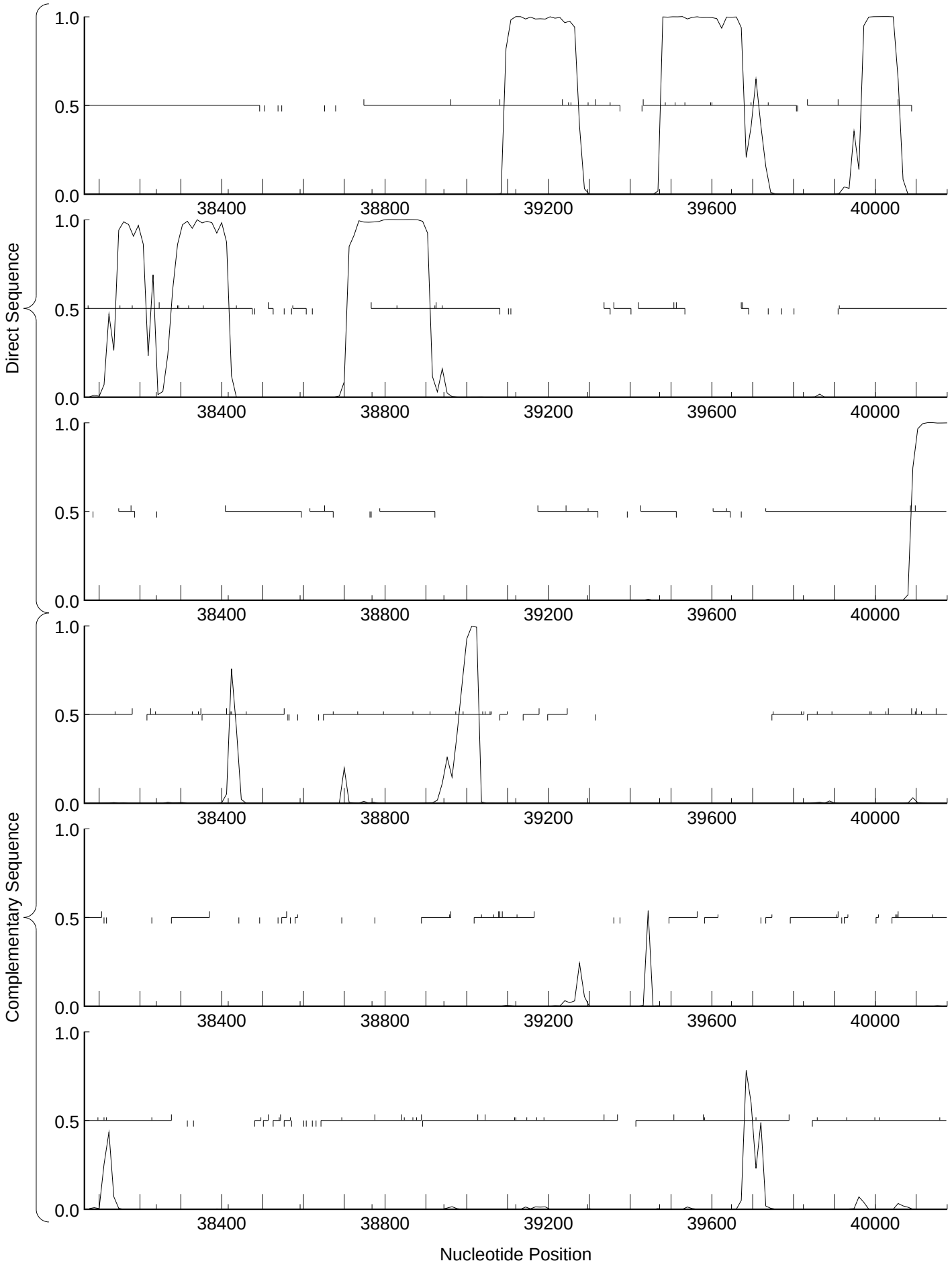
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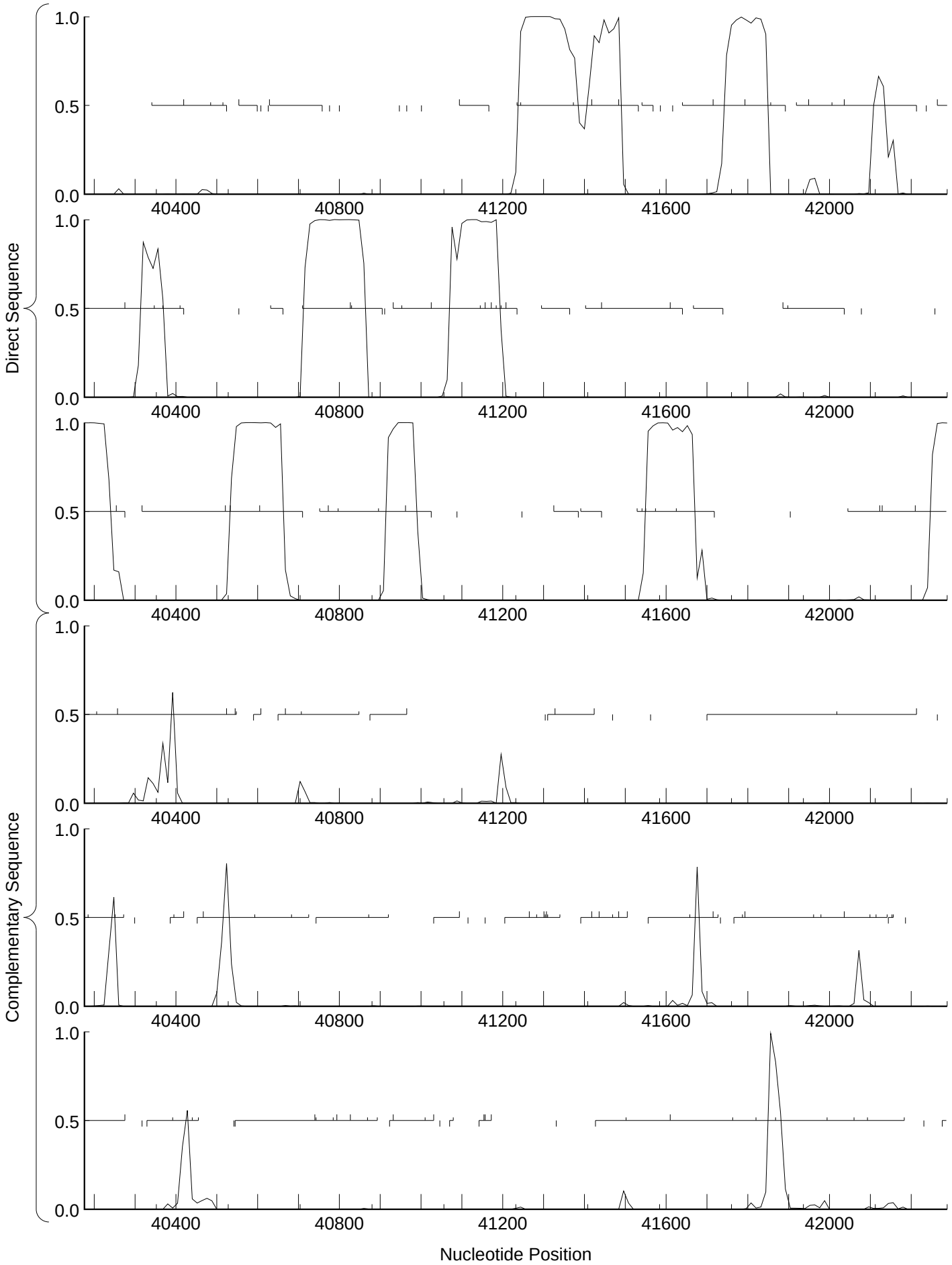
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 19/23



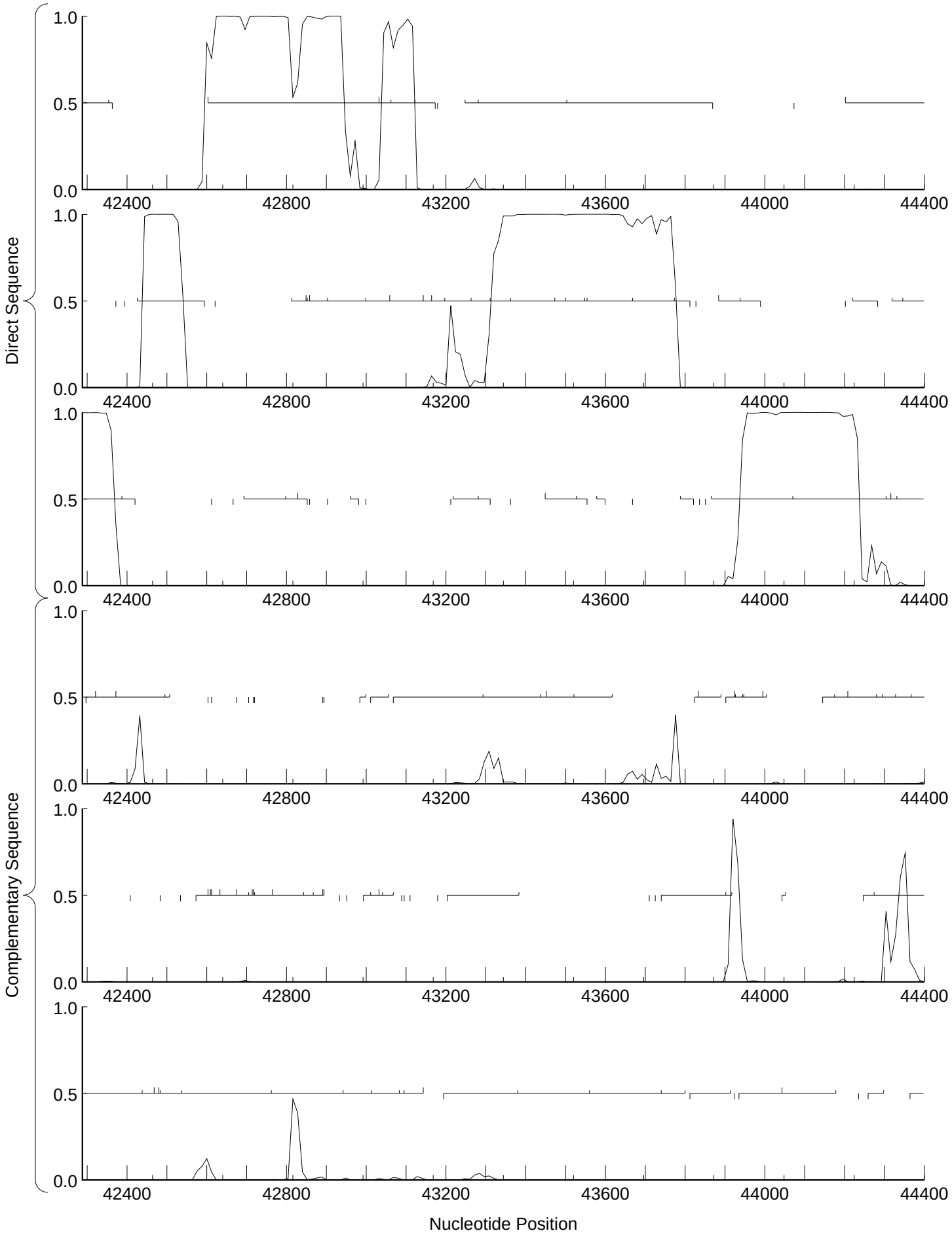
Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 20/23



Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 21/23



Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 22/23



Arthrobacter phage TripleJ complete sequence, 45507 bp, circularly permuted, Cluster FJ, Order 4, Window 96, Step 12, 23/23

