GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Abbyshoes complete sequence, 51324 bp including 10-base 3’ overhang (CG
Analysis Date    :  3/28/24 at 21:48:36
Pages            :  26
Sequence Length  :  51324 bp
GC Content       :   63.74%

Window Length    :  96 bp
Window Step      :  12 bp
Threshold Value  :  0.500

PS-Version       :  1.2

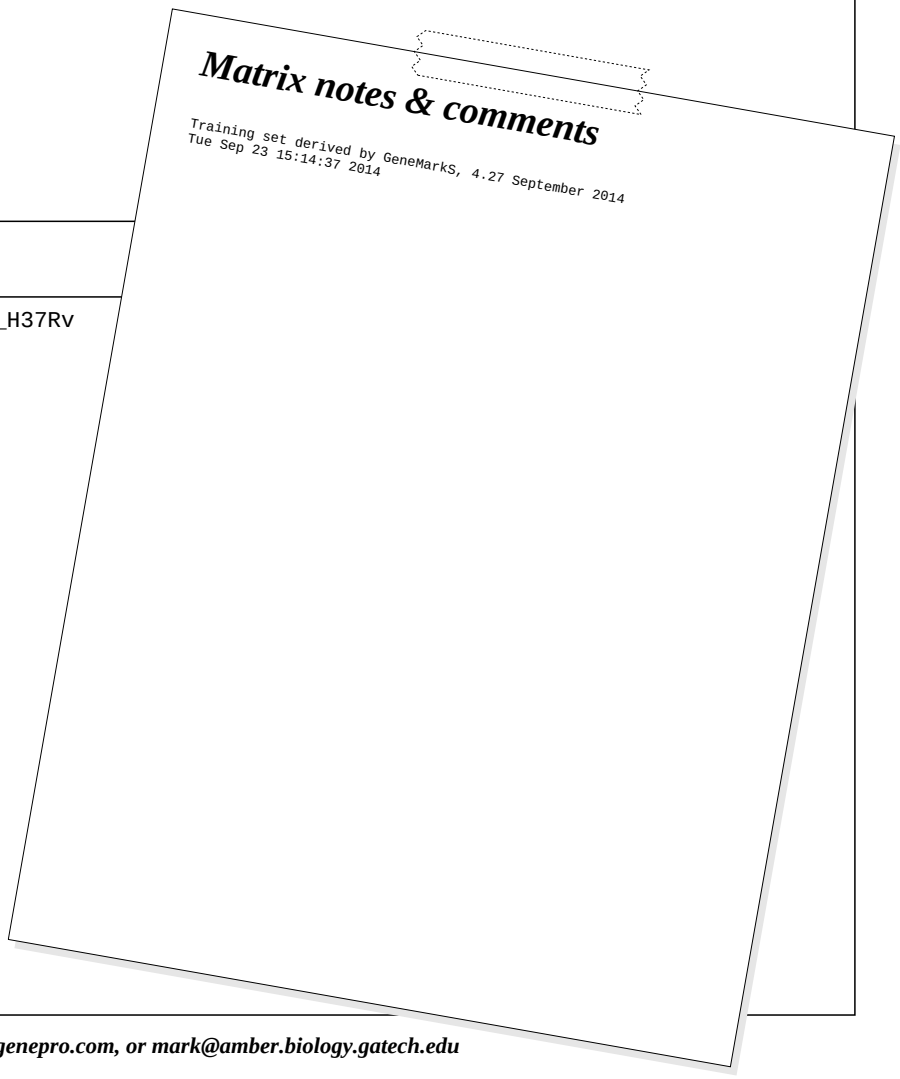
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
```

MATRIX INFORMATION

```
Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

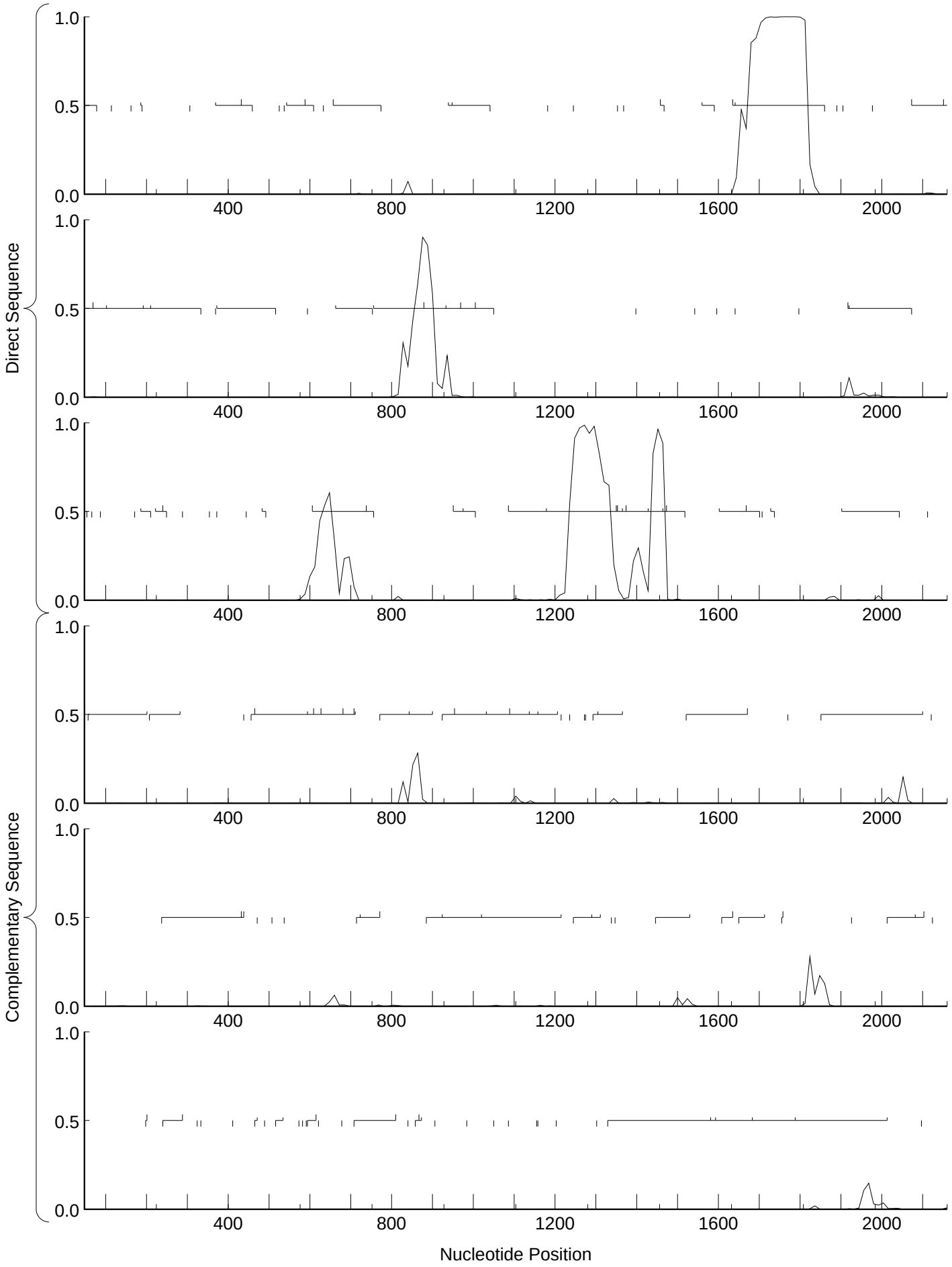
Order      :  4


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

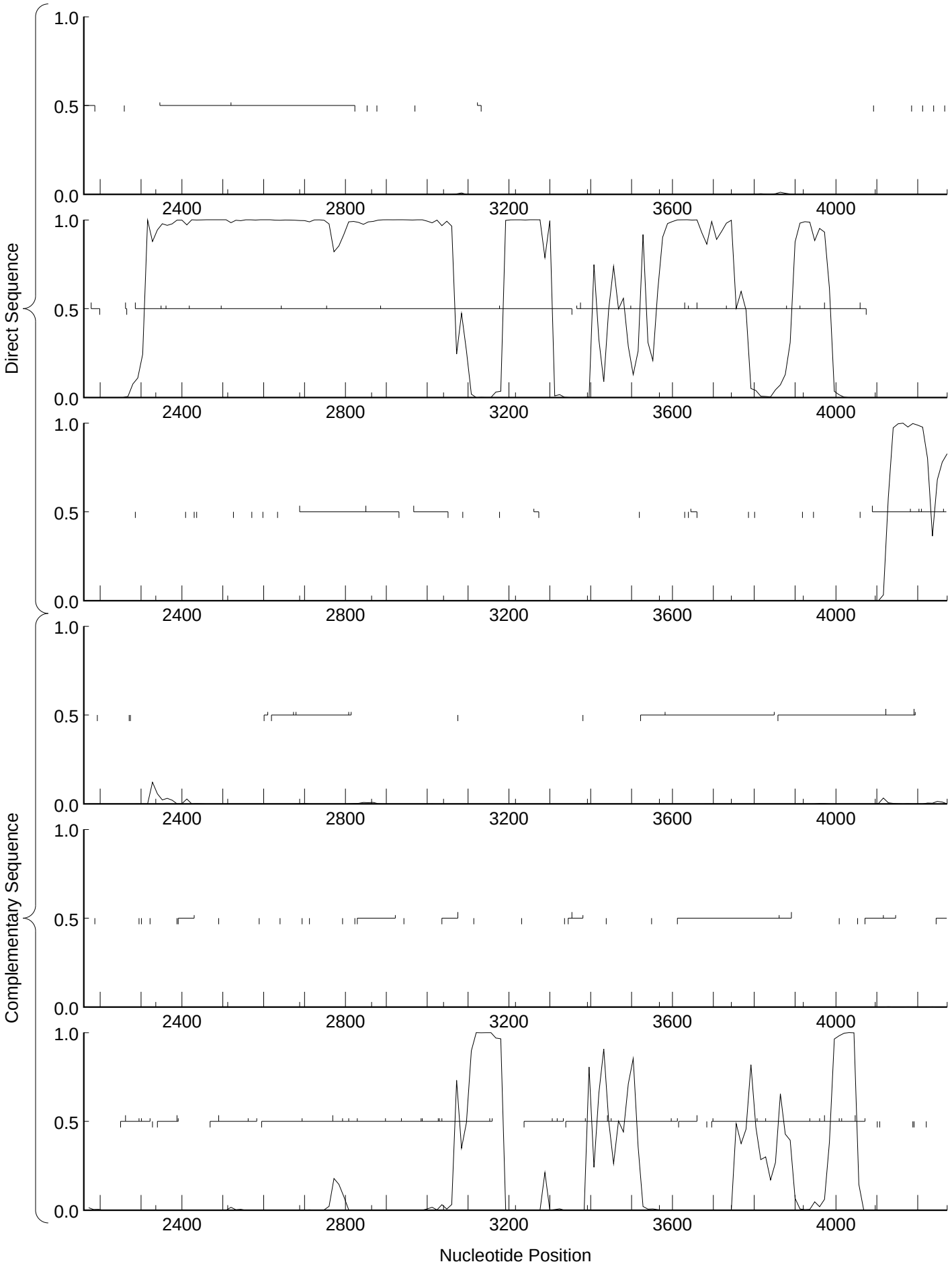


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

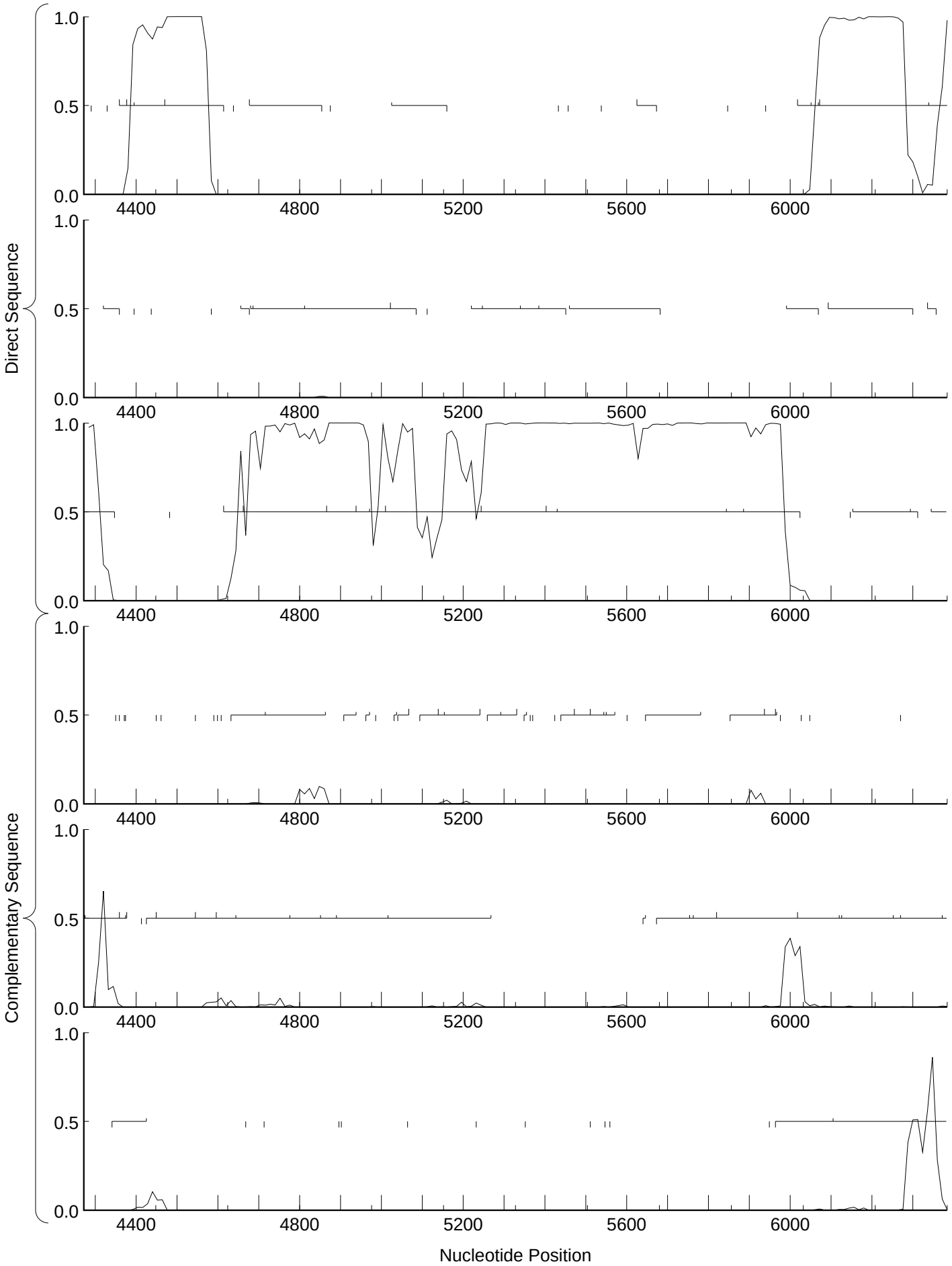
es complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 2/26



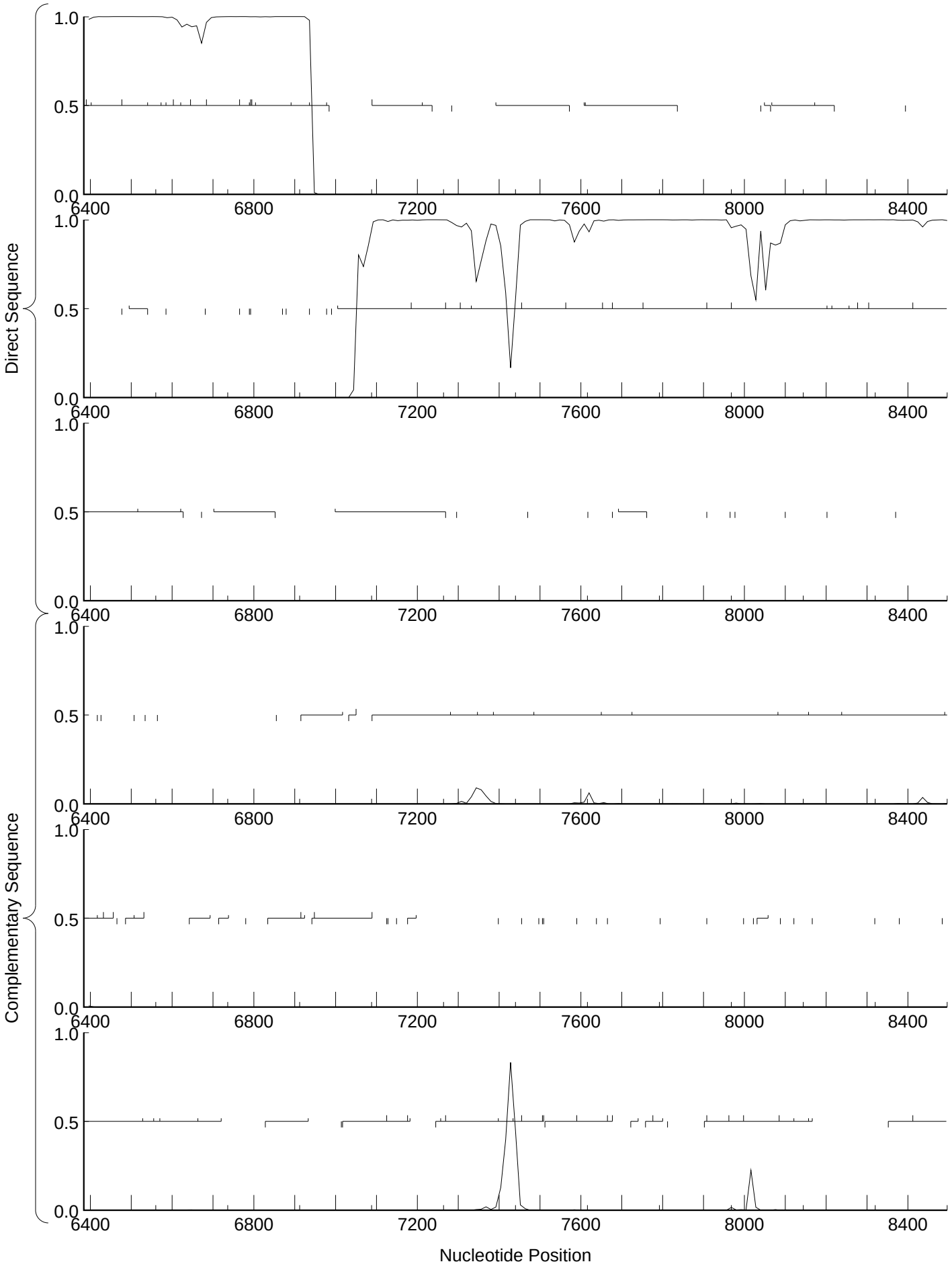
es complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 3/26



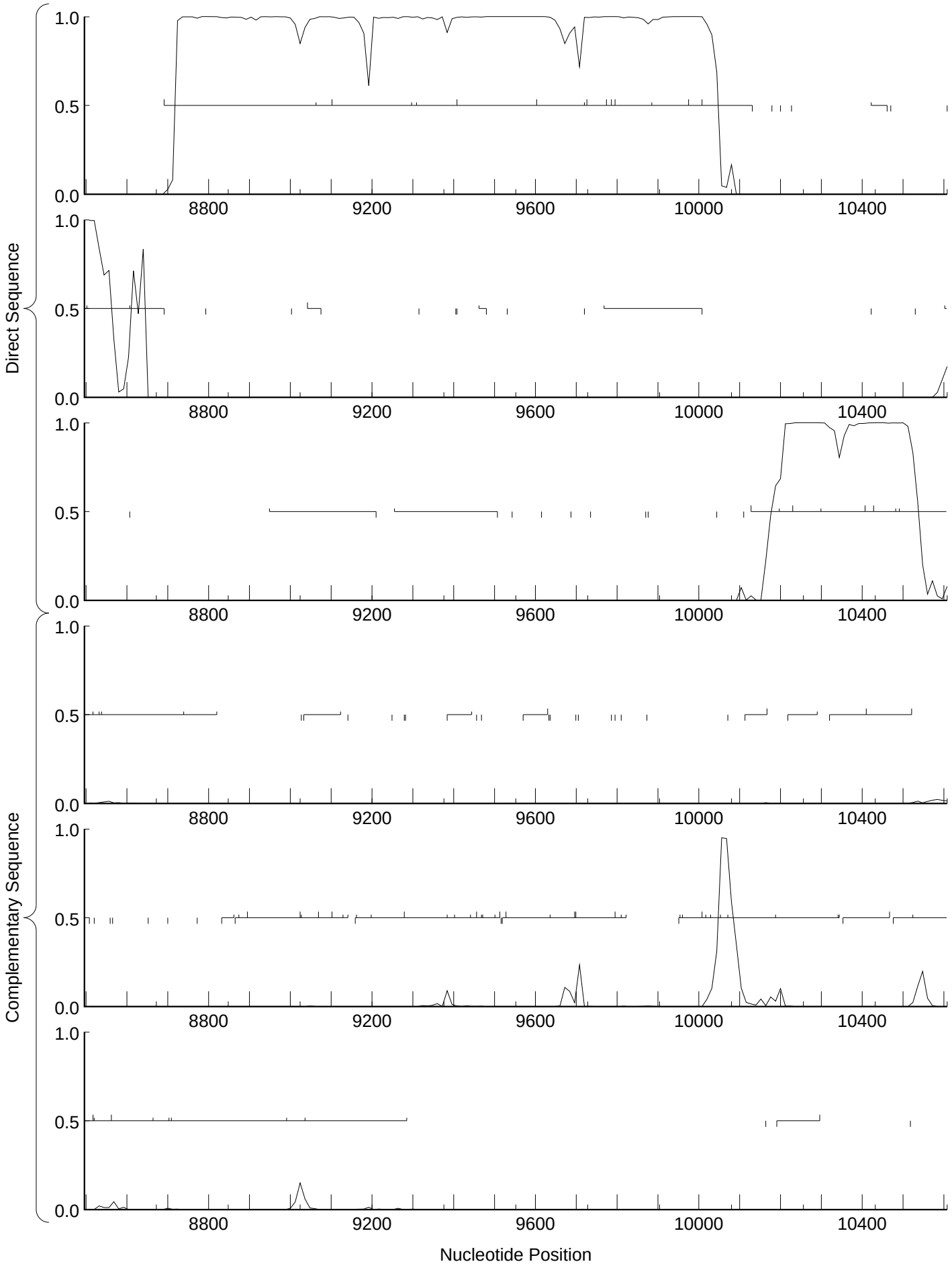
es complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 4/26



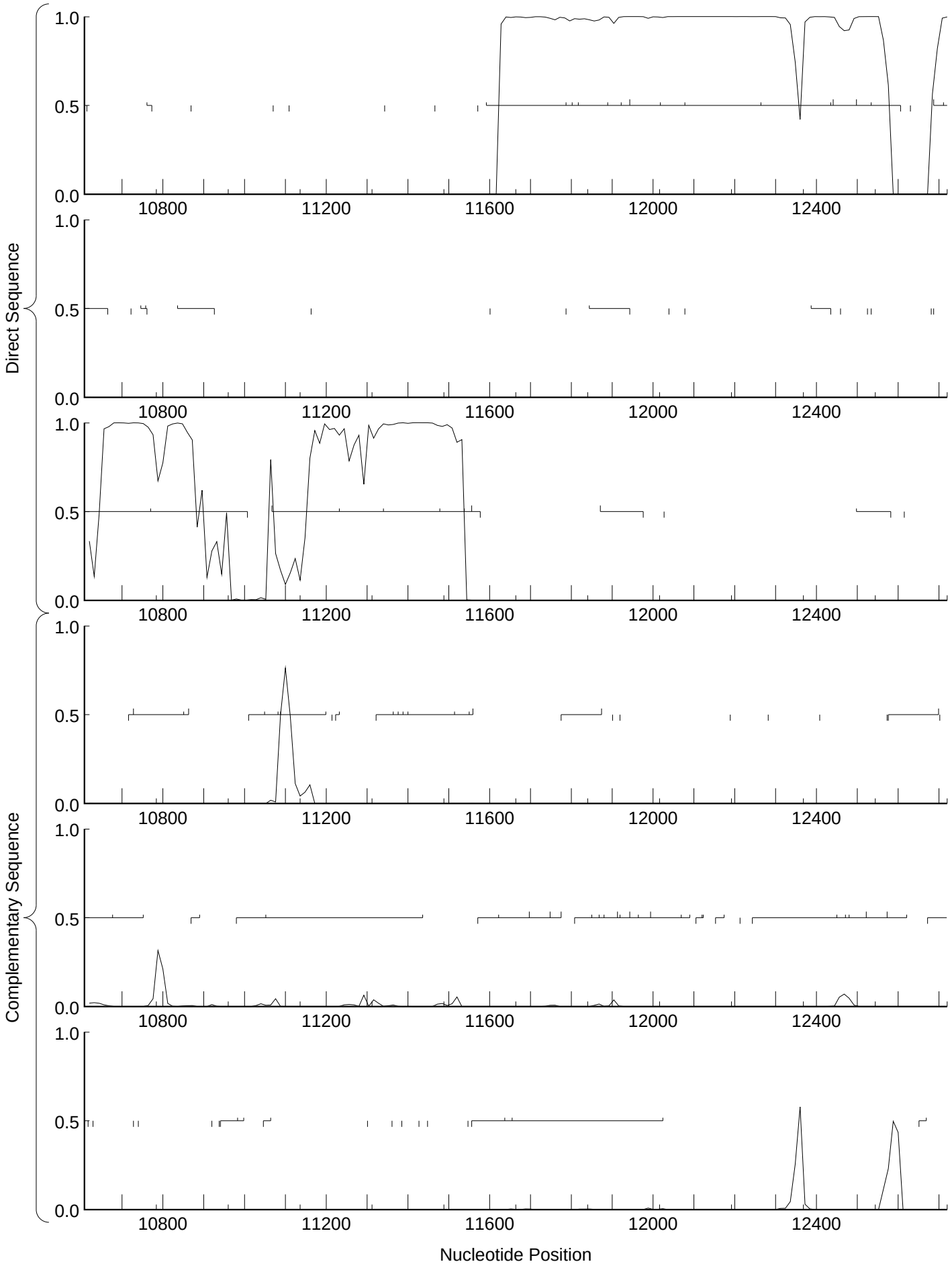
es complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 5/26



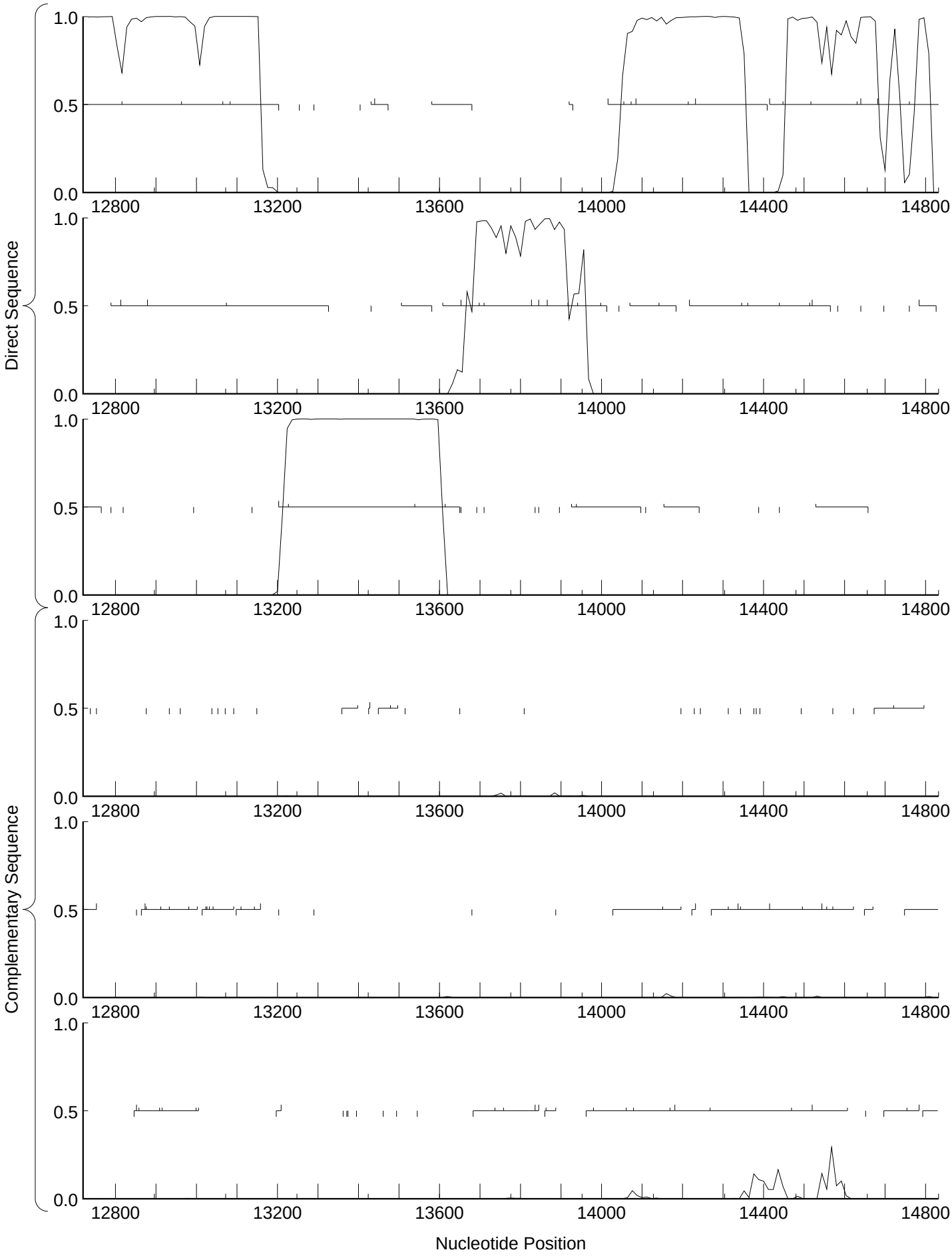
es complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 6/26



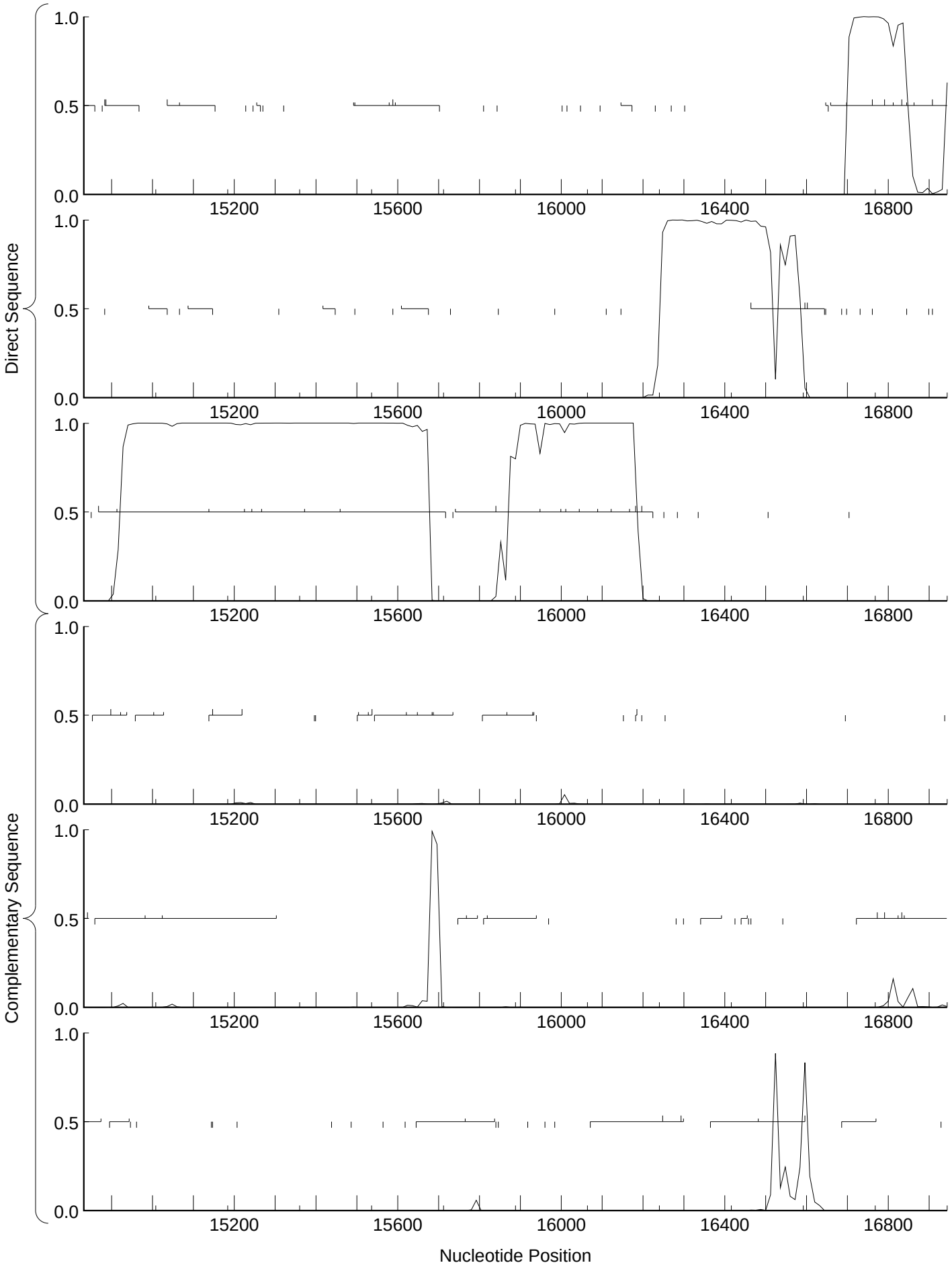
es complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 7/26



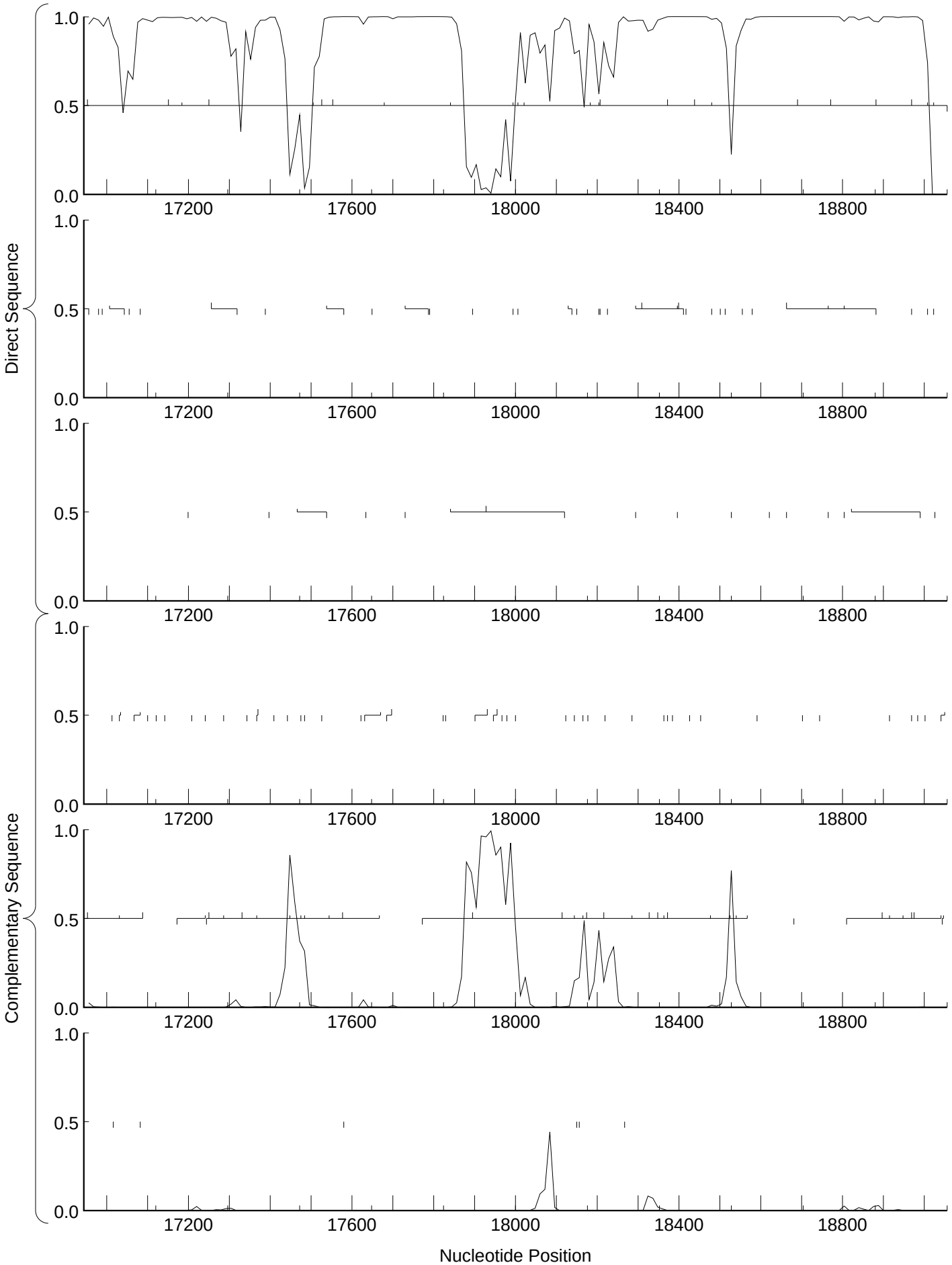
es complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 8/26



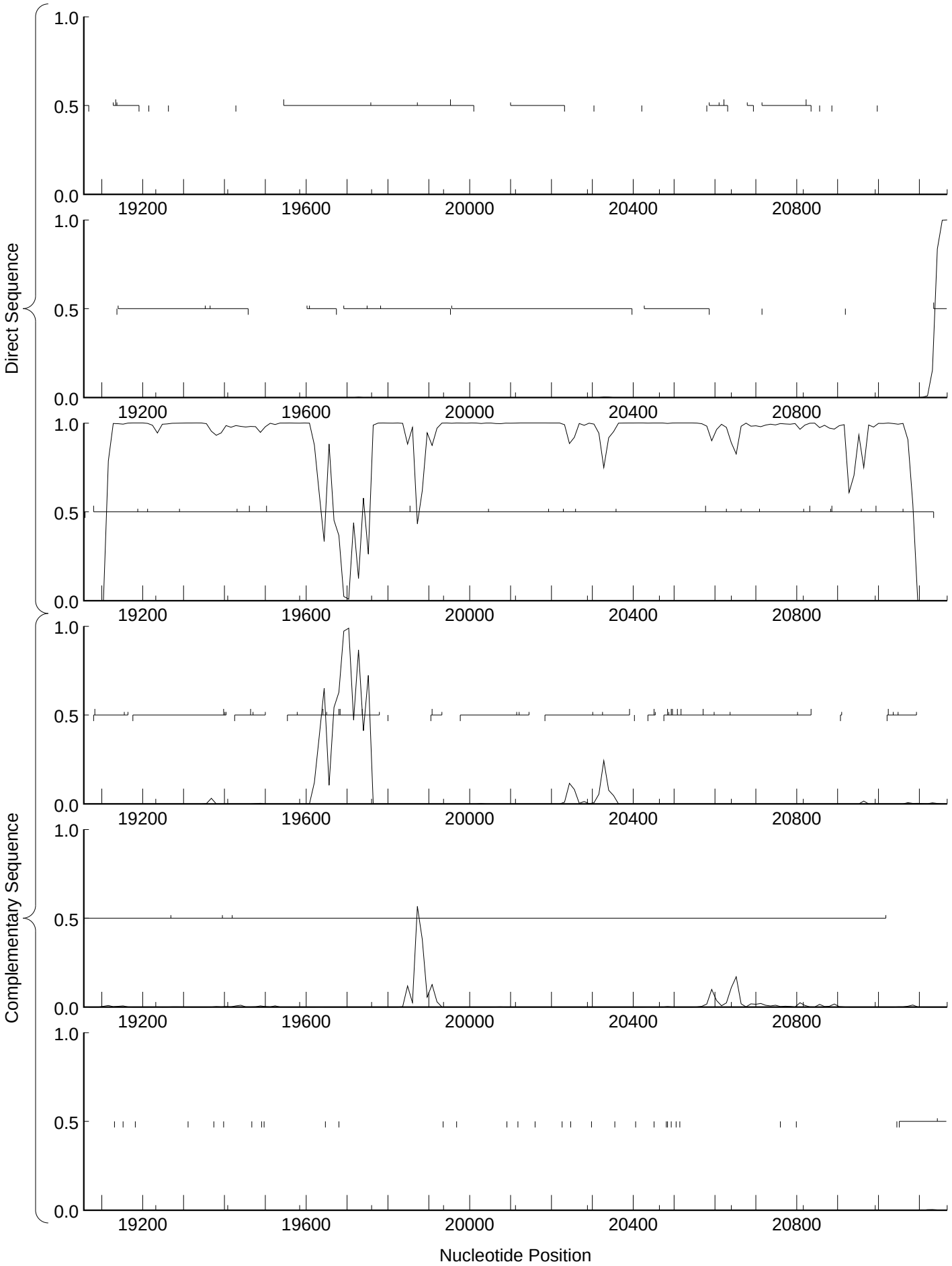
es complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 9/26

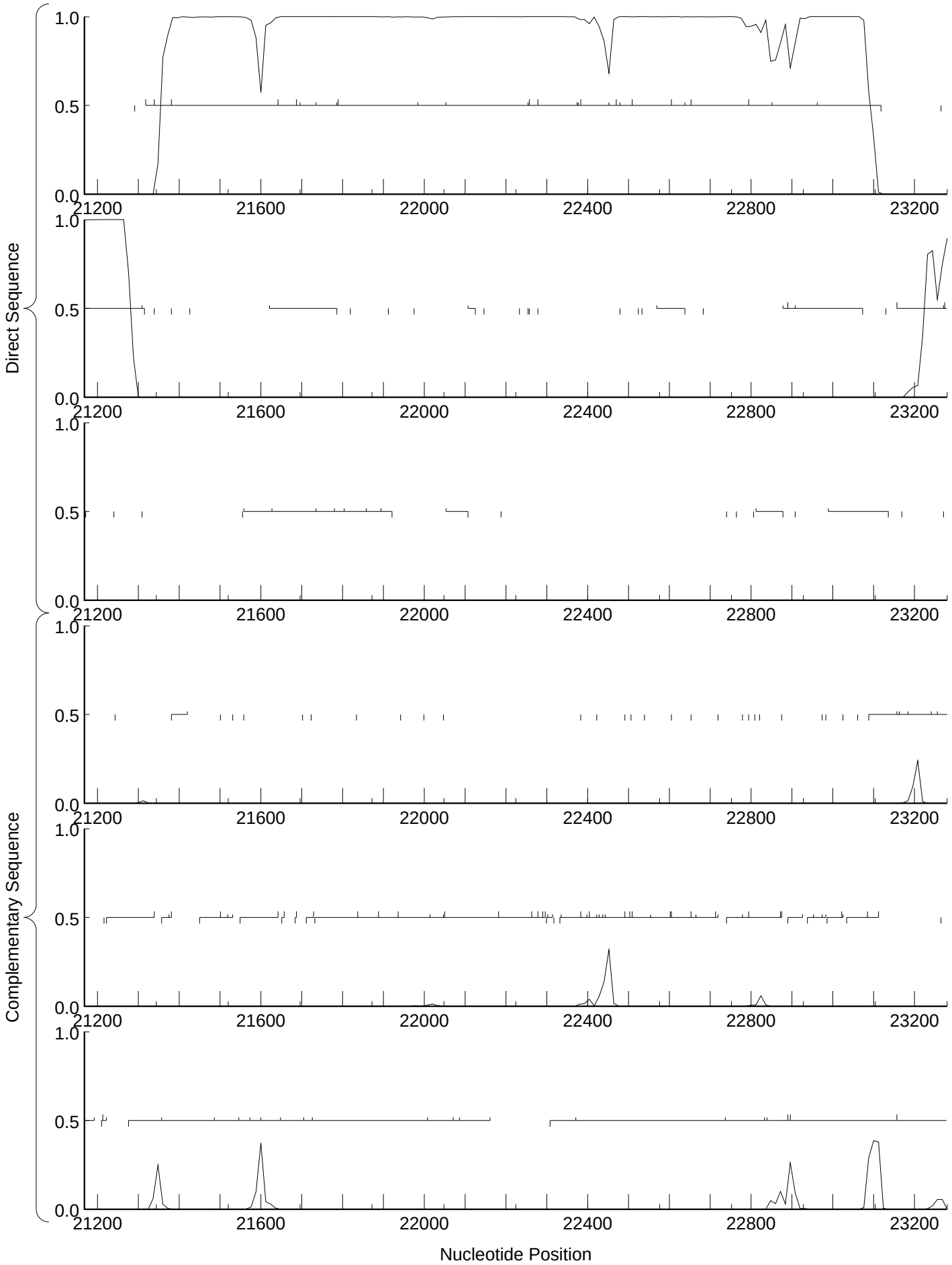


s complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 10/26

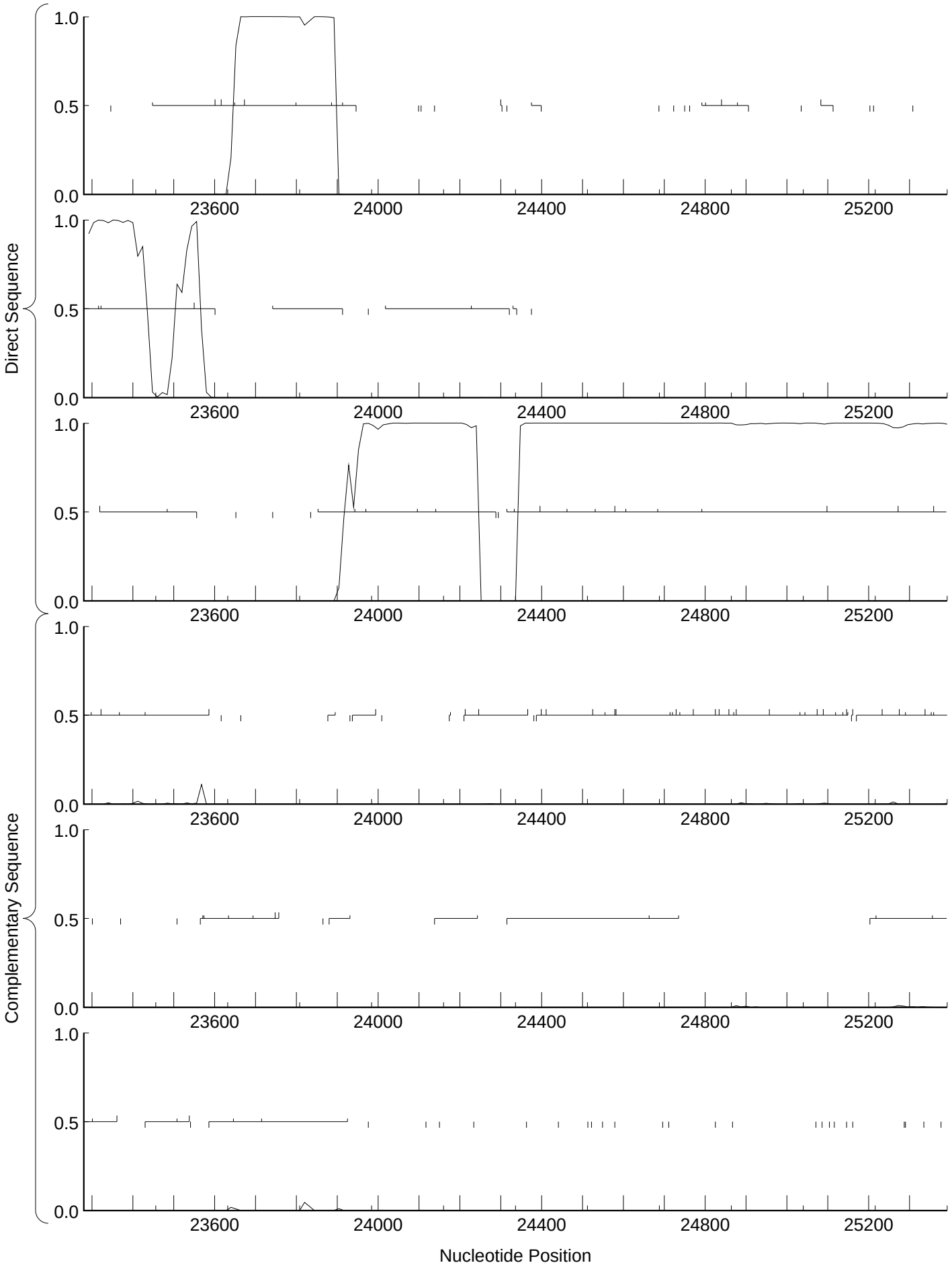


s complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 11/26

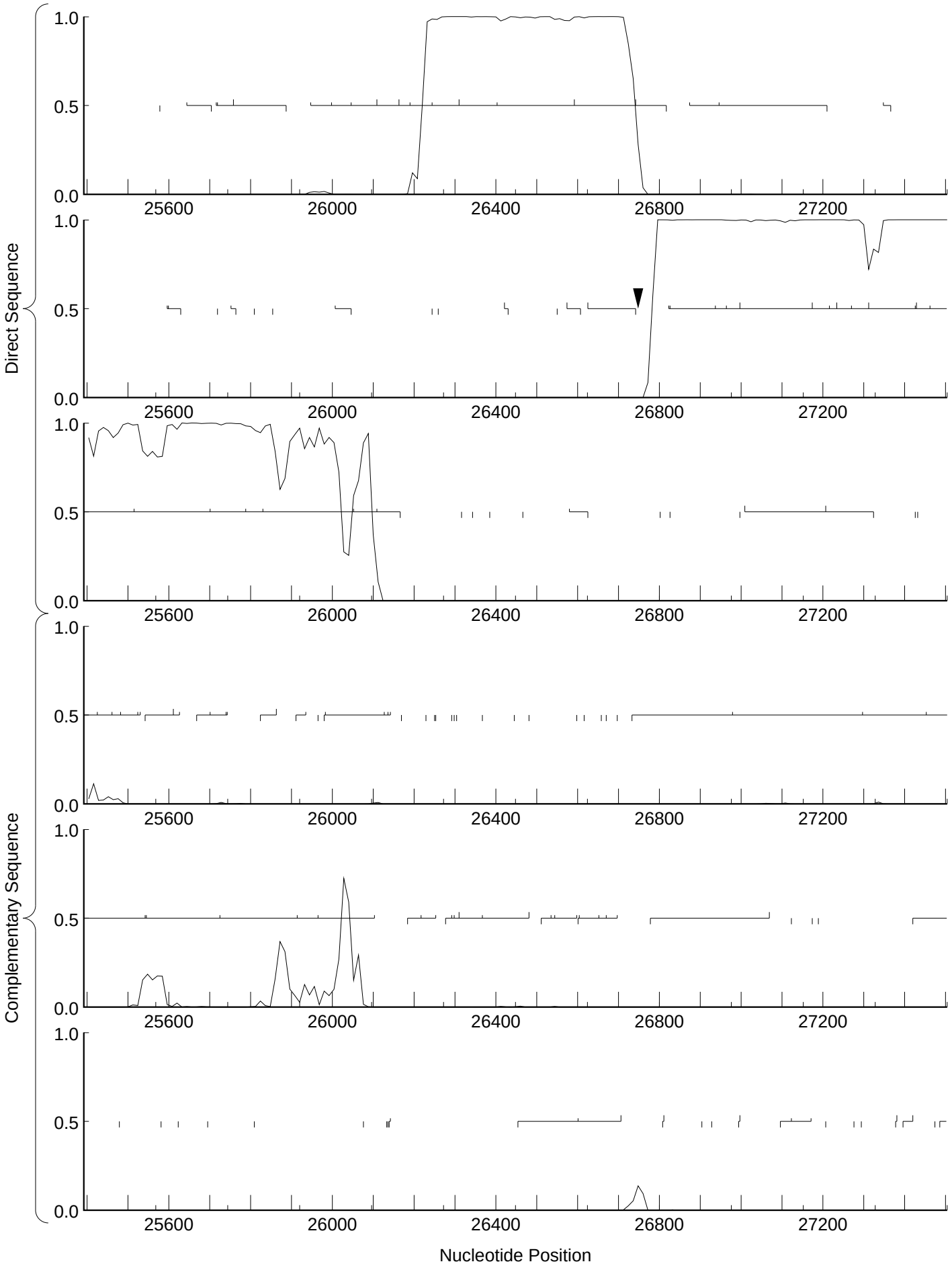



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
21200
21600
22000
22400
22800
23200
Nucleotide Position

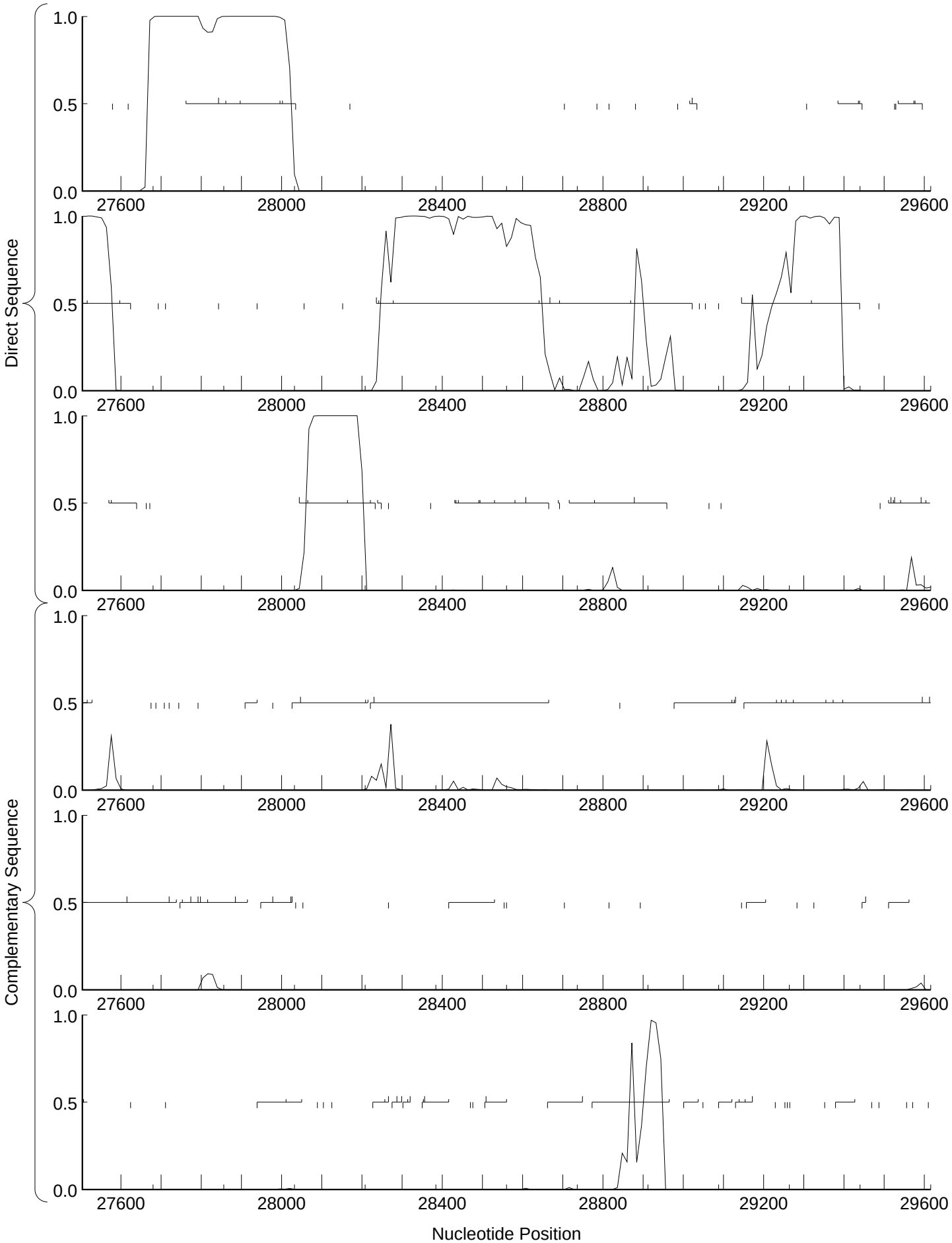
s complete sequence, 51324 bp including 10-base 3' overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 13/26



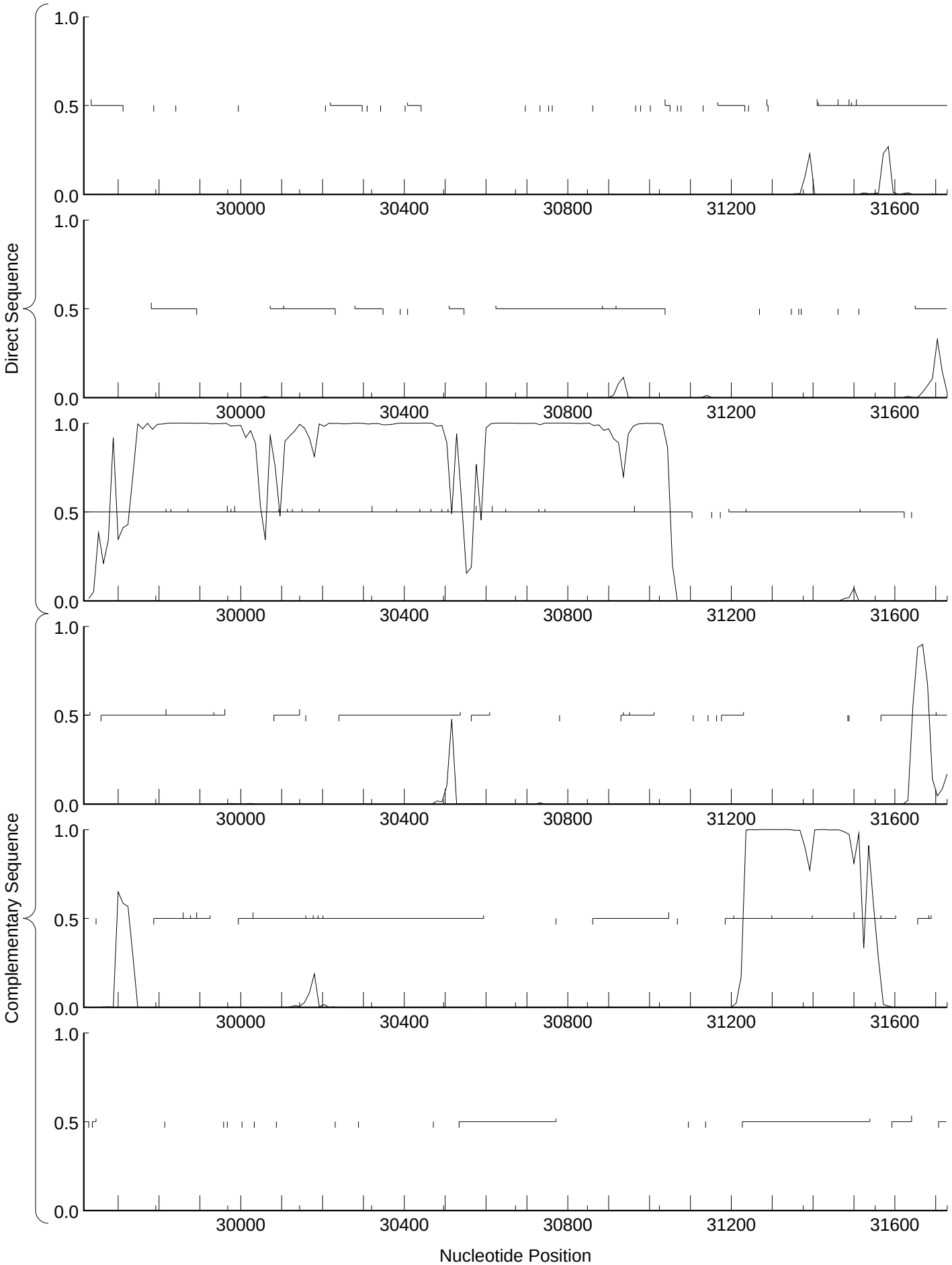
s complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 14/26



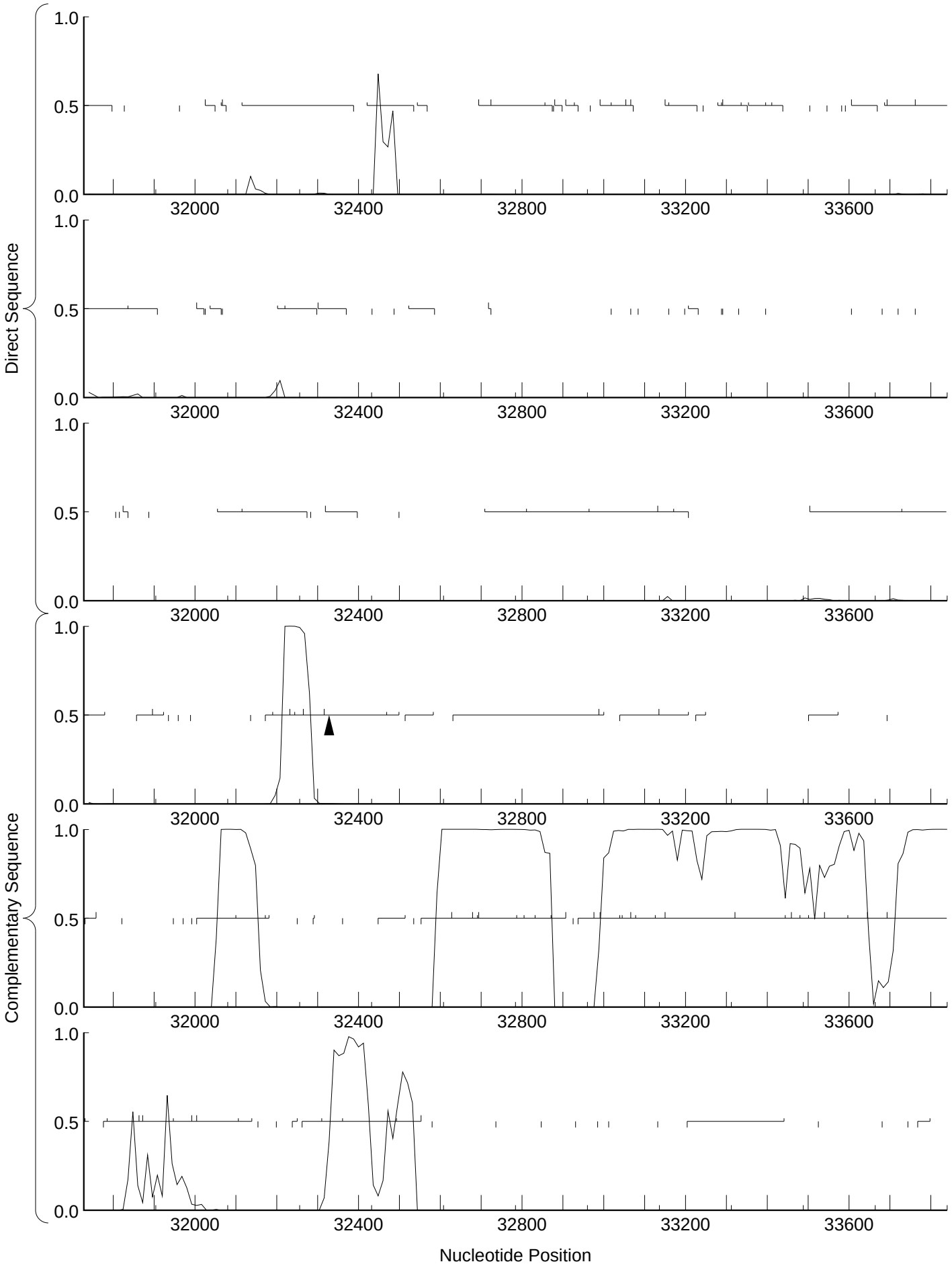
s complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 15/26



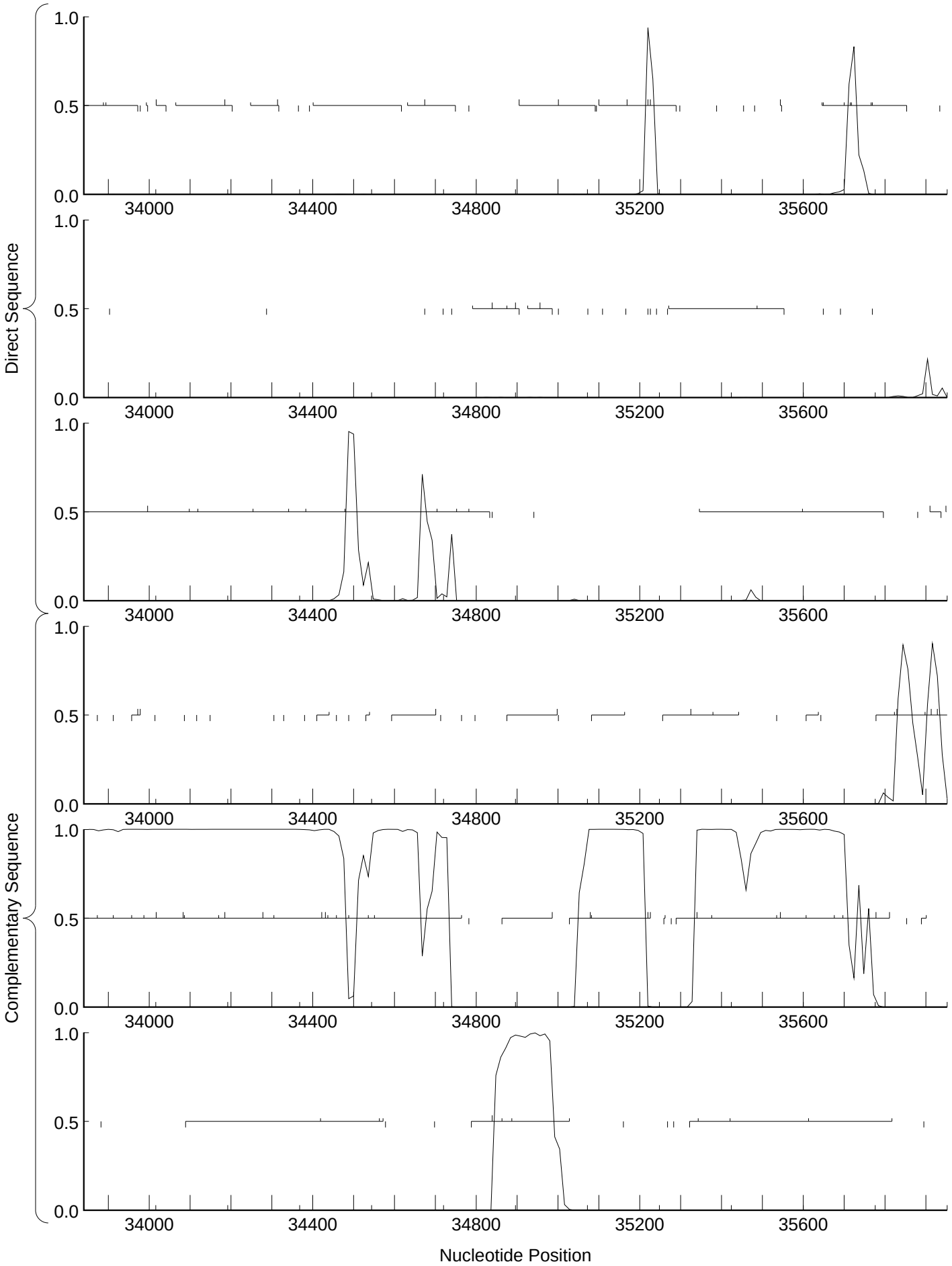
s complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 16/26

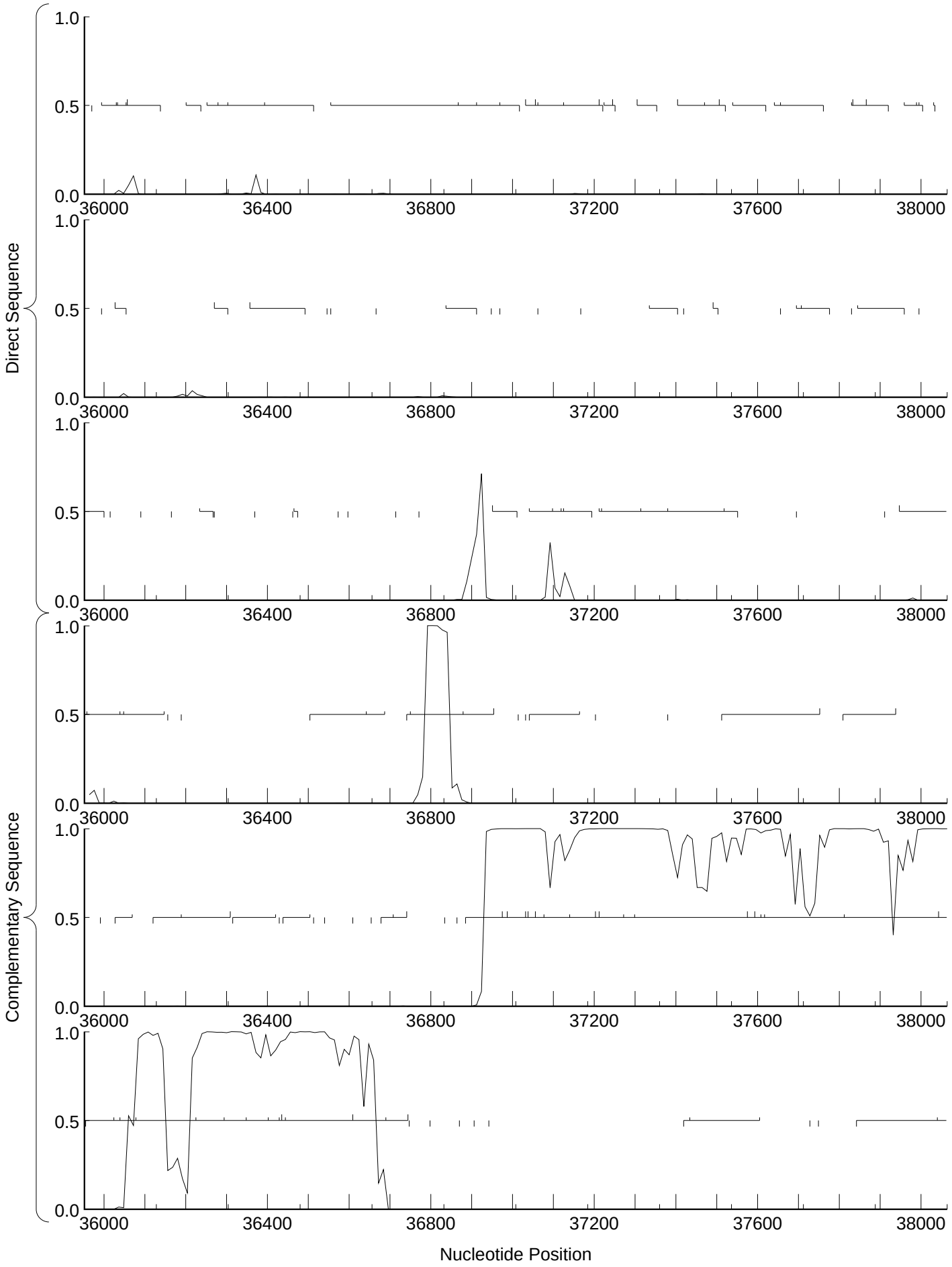


s complete sequence, 51324 bp including 10-base 3' overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 17/26

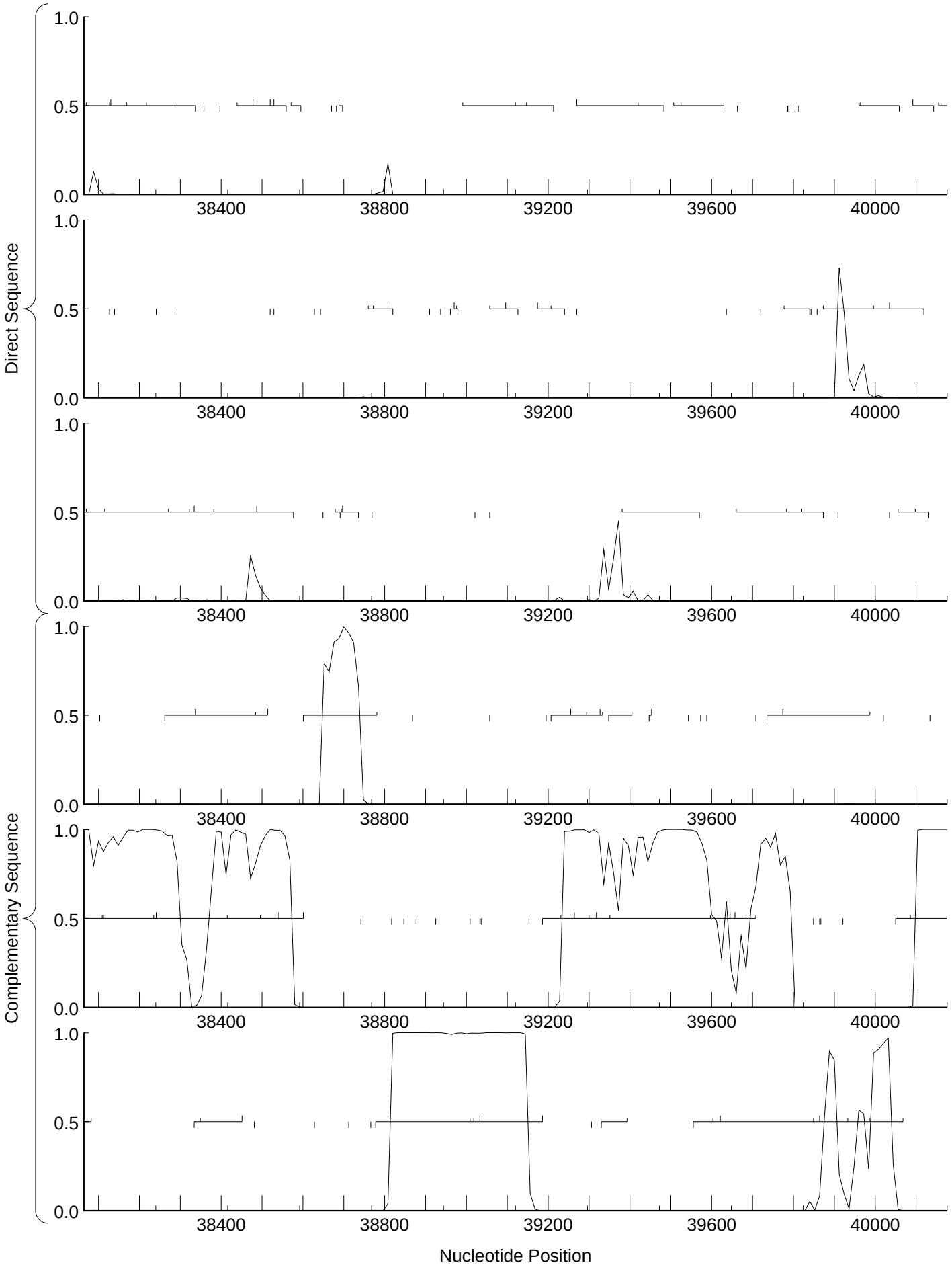


s complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 18/26

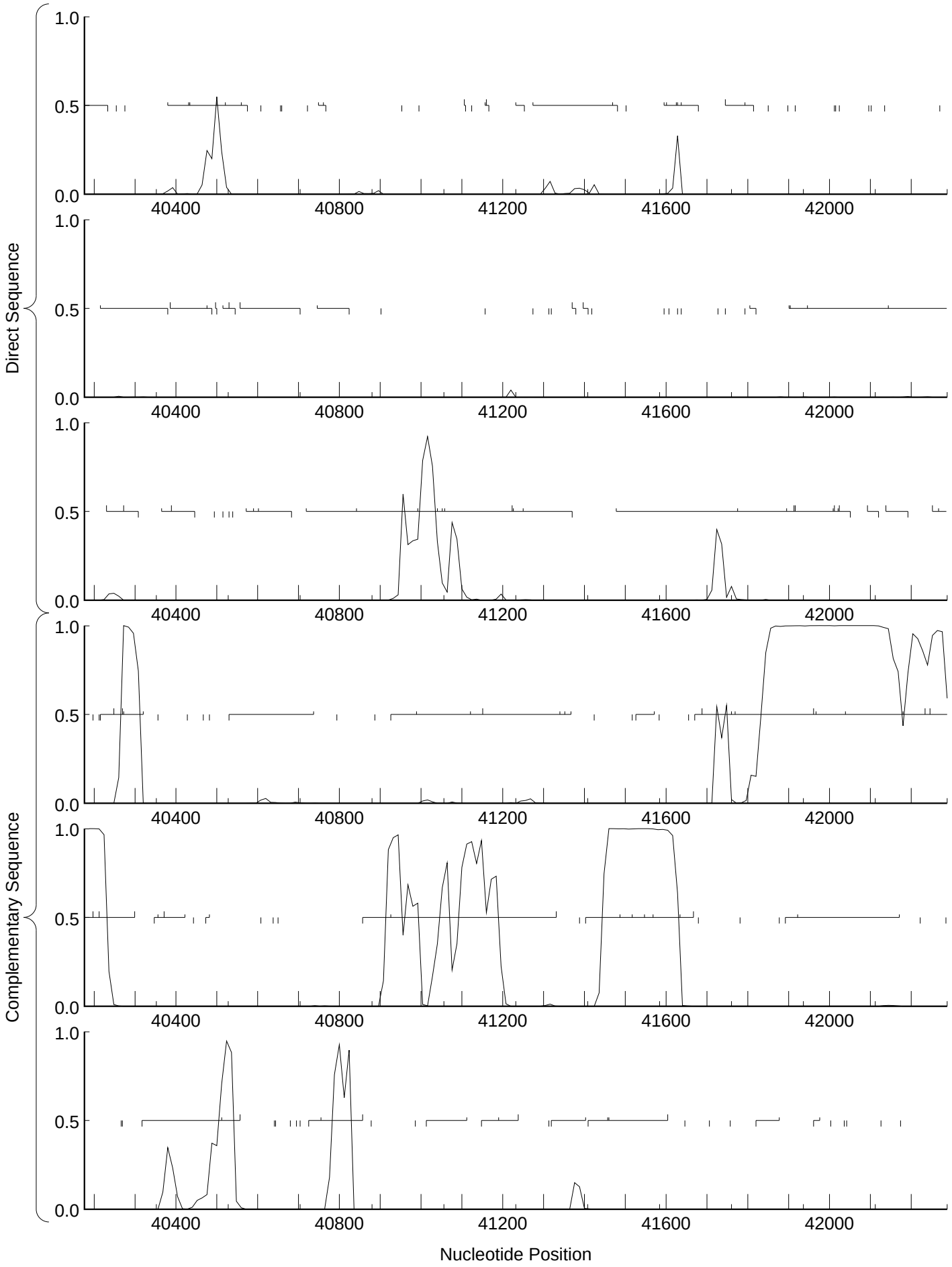



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
36000
36400
36800
37200
37600
38000
Nucleotide Position

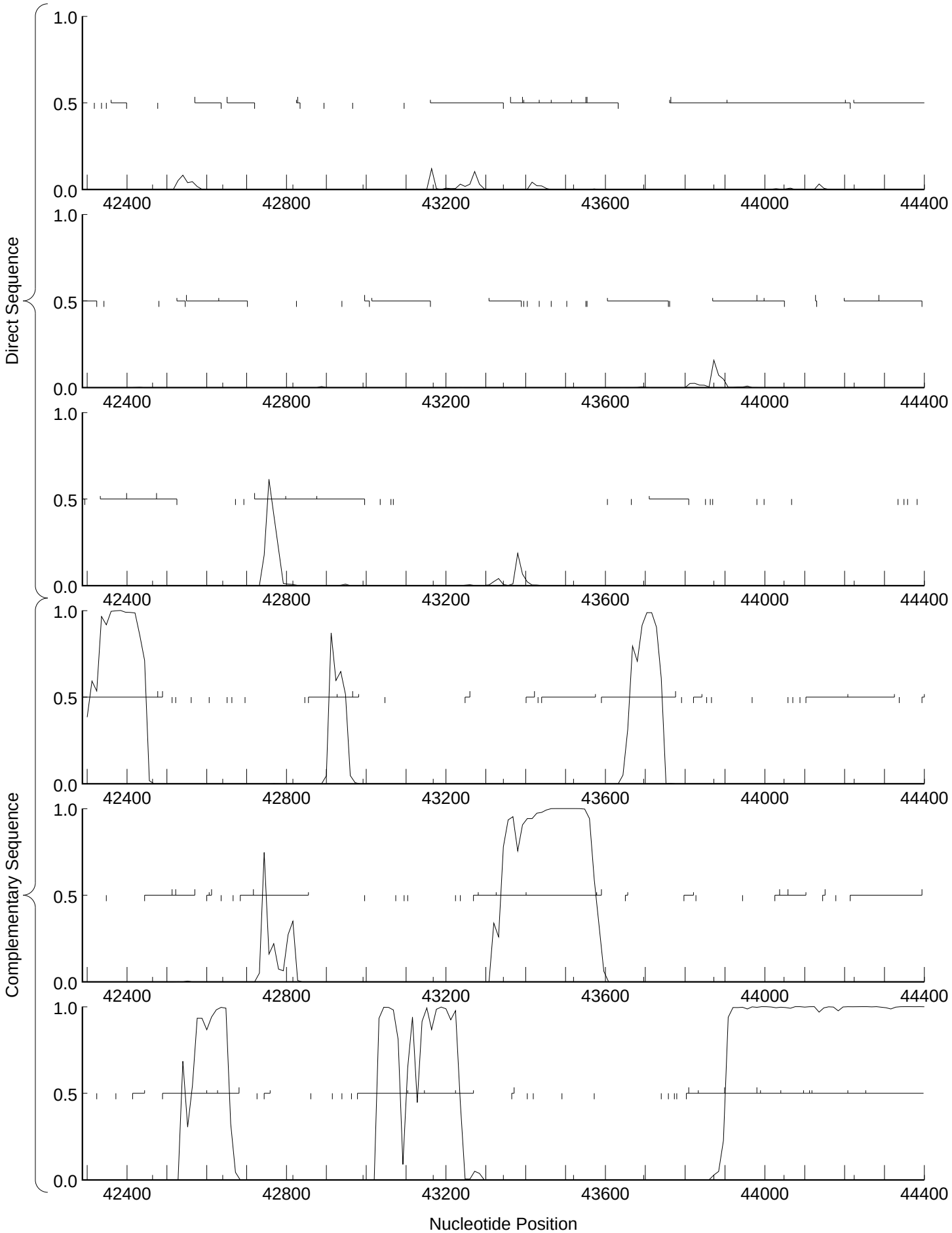
s complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 20/26



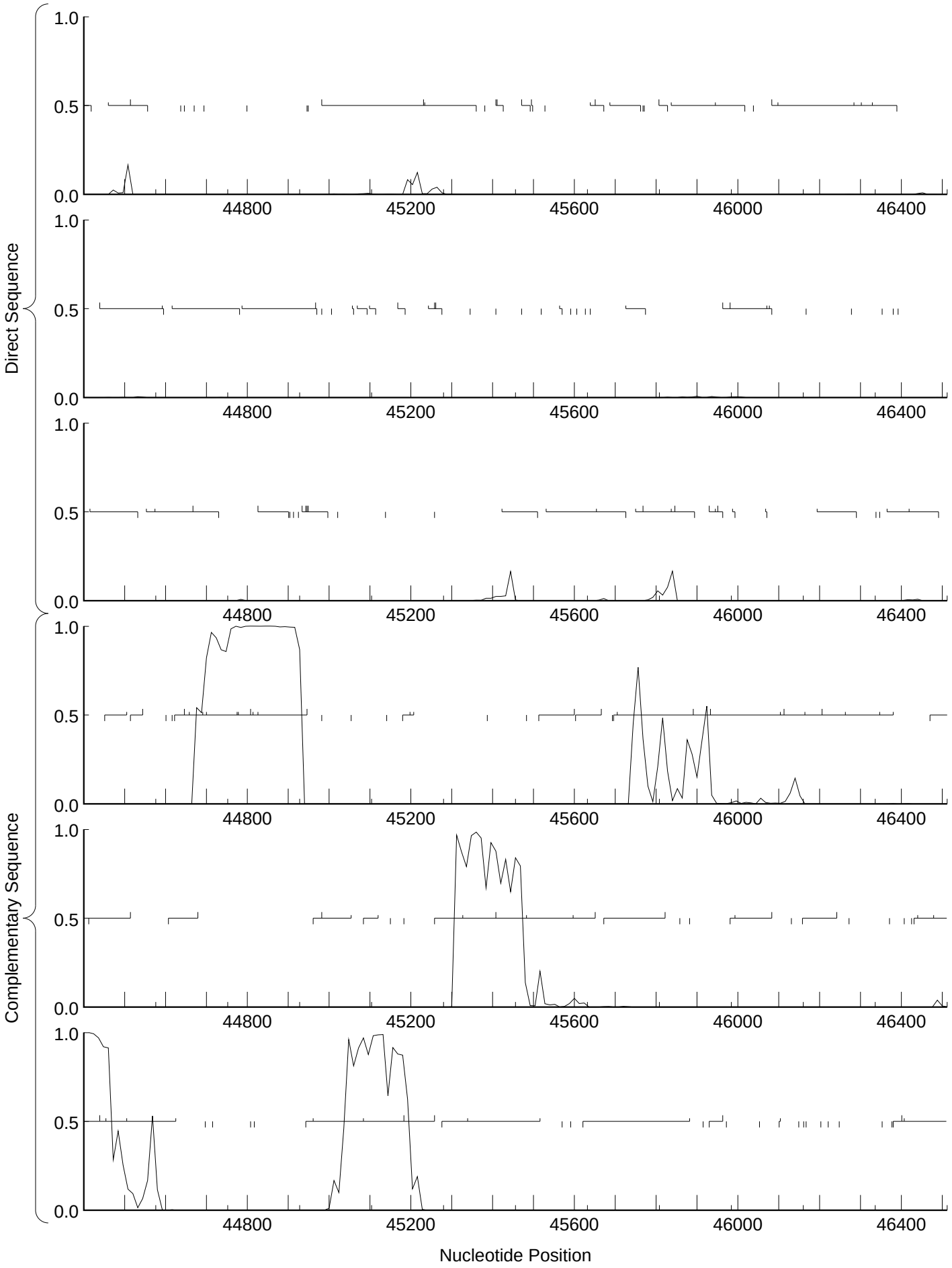
s complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 21/26



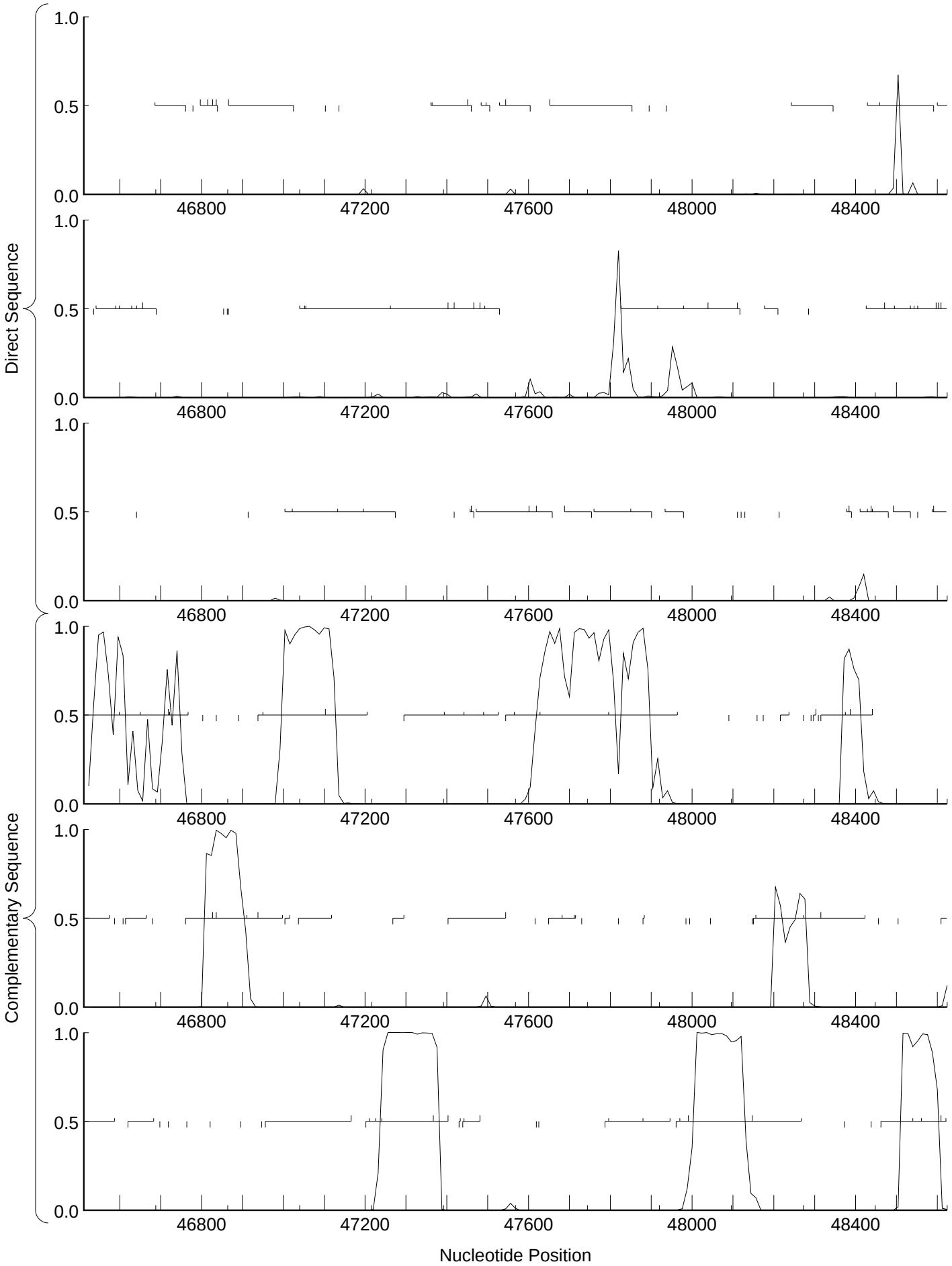
s complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 22/26

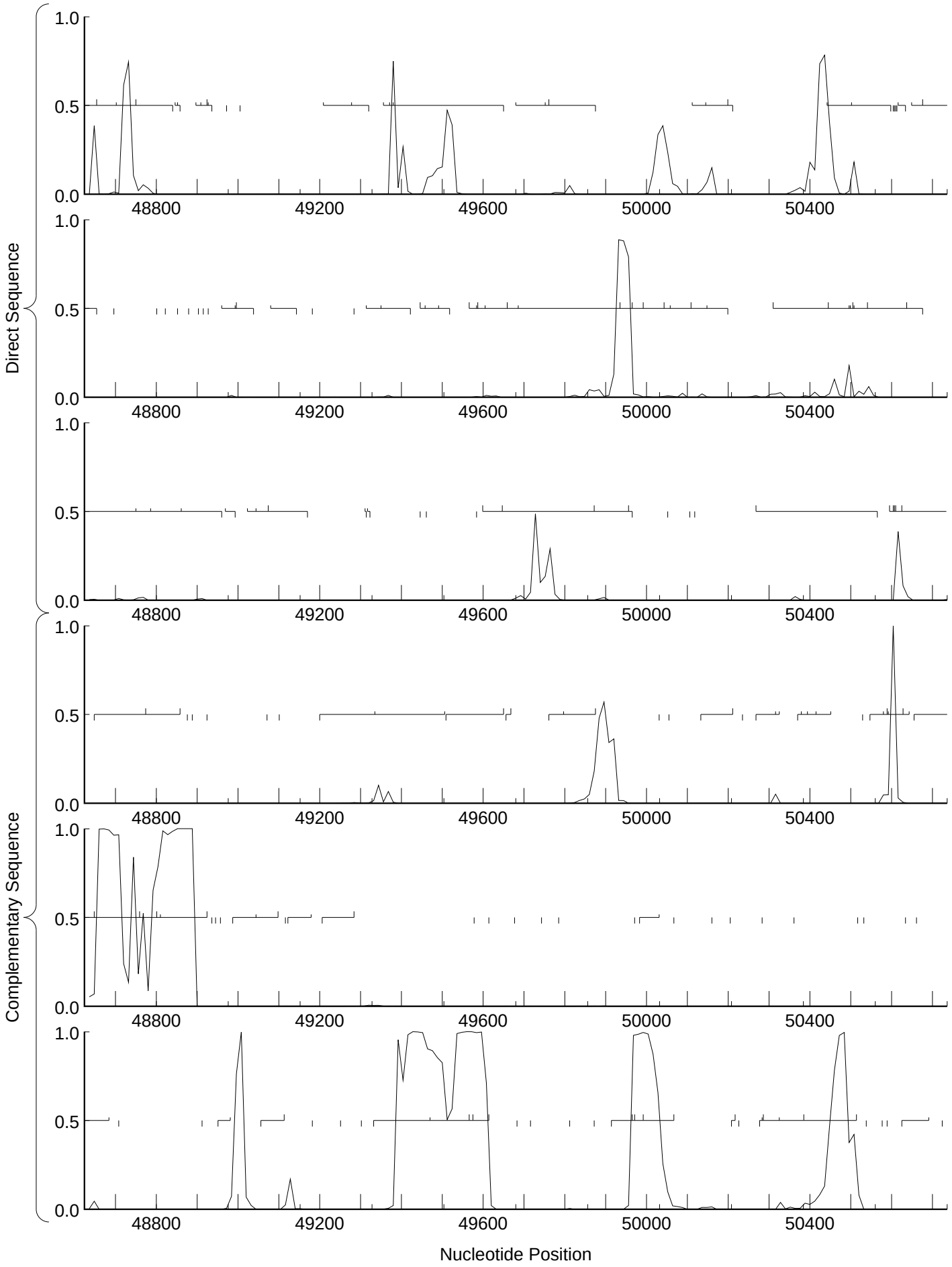


s complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 23/26



s complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 24/26




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
48800
49200
49600
50000
50400
Nucleotide Position

s complete sequence, 51324 bp including 10-base 3’ overhang (CGGATGGTAA) Cluster A1, Order 4, Window 96, Step 12, 26/26

