# GeneMark

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

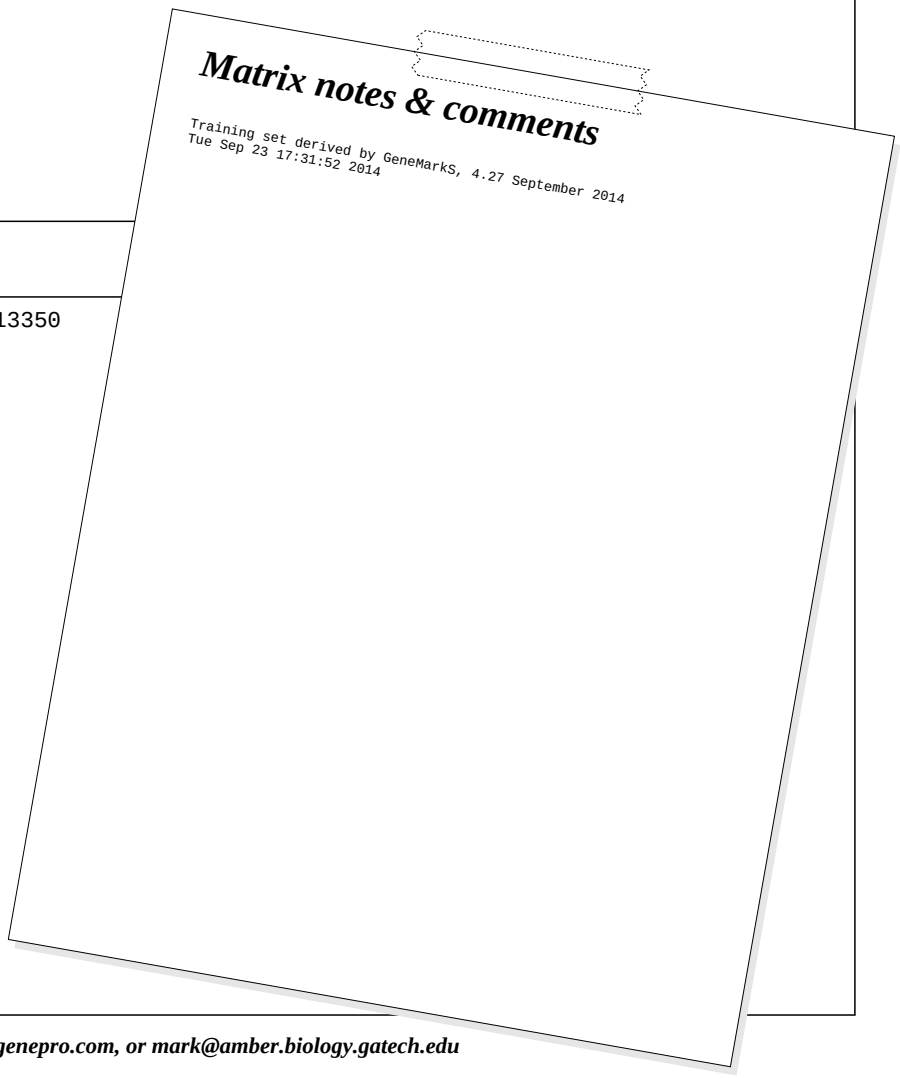
## PROGRAM INFORMATION

```
Sequence         :  Streptomyces phage Asten complete sequence, 51571 bp including 11-base 3' overhang (CGGCCAG
Analysis Date    :  3/29/24 at 4:24:36
Pages            :  26
Sequence Length  :  51571 bp
GC Content       :   65.67%

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

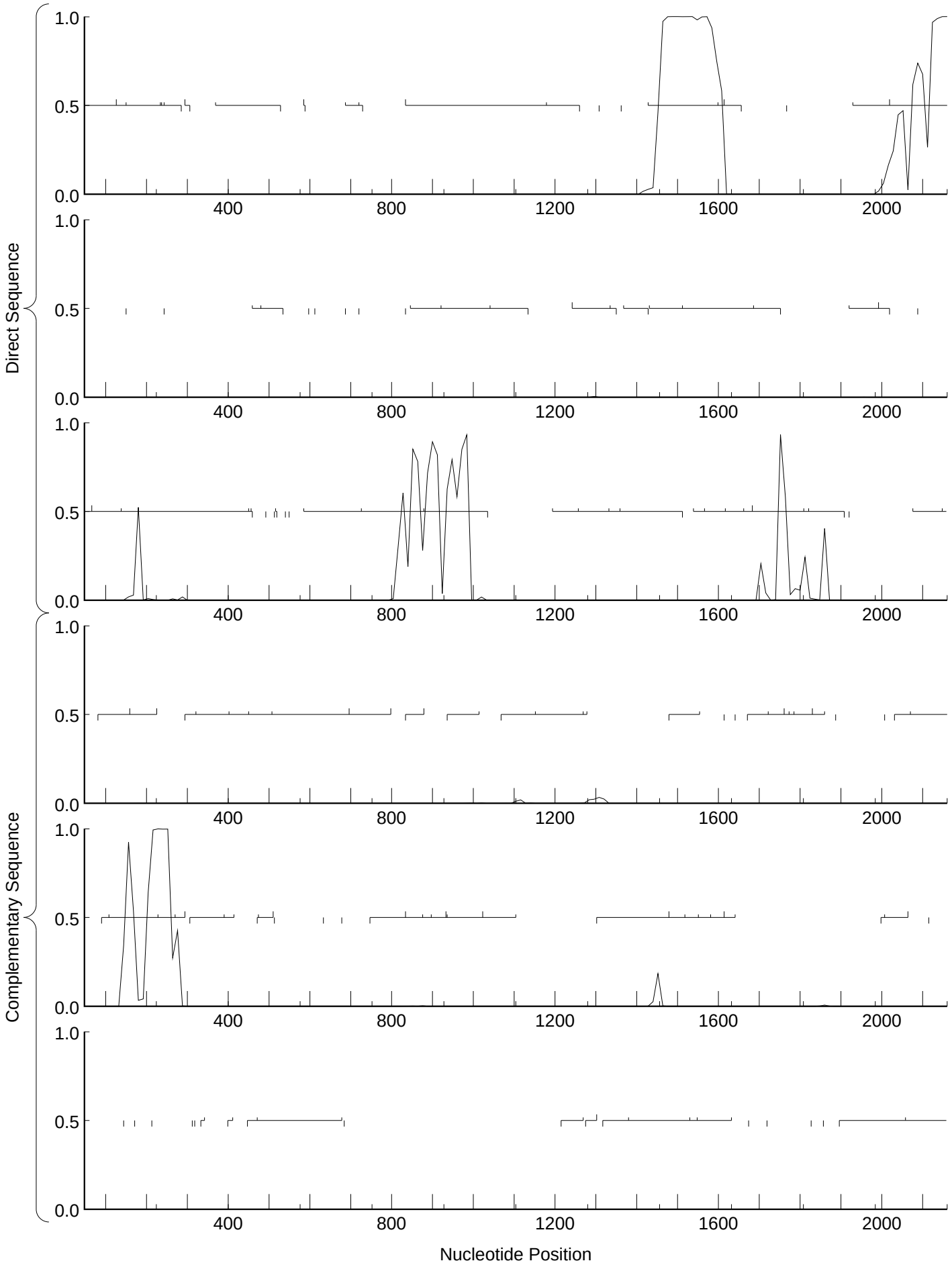
```
Matrix     :  Streptomyces_griseus_NBRC_13350
Author     :  -

Order      :  4

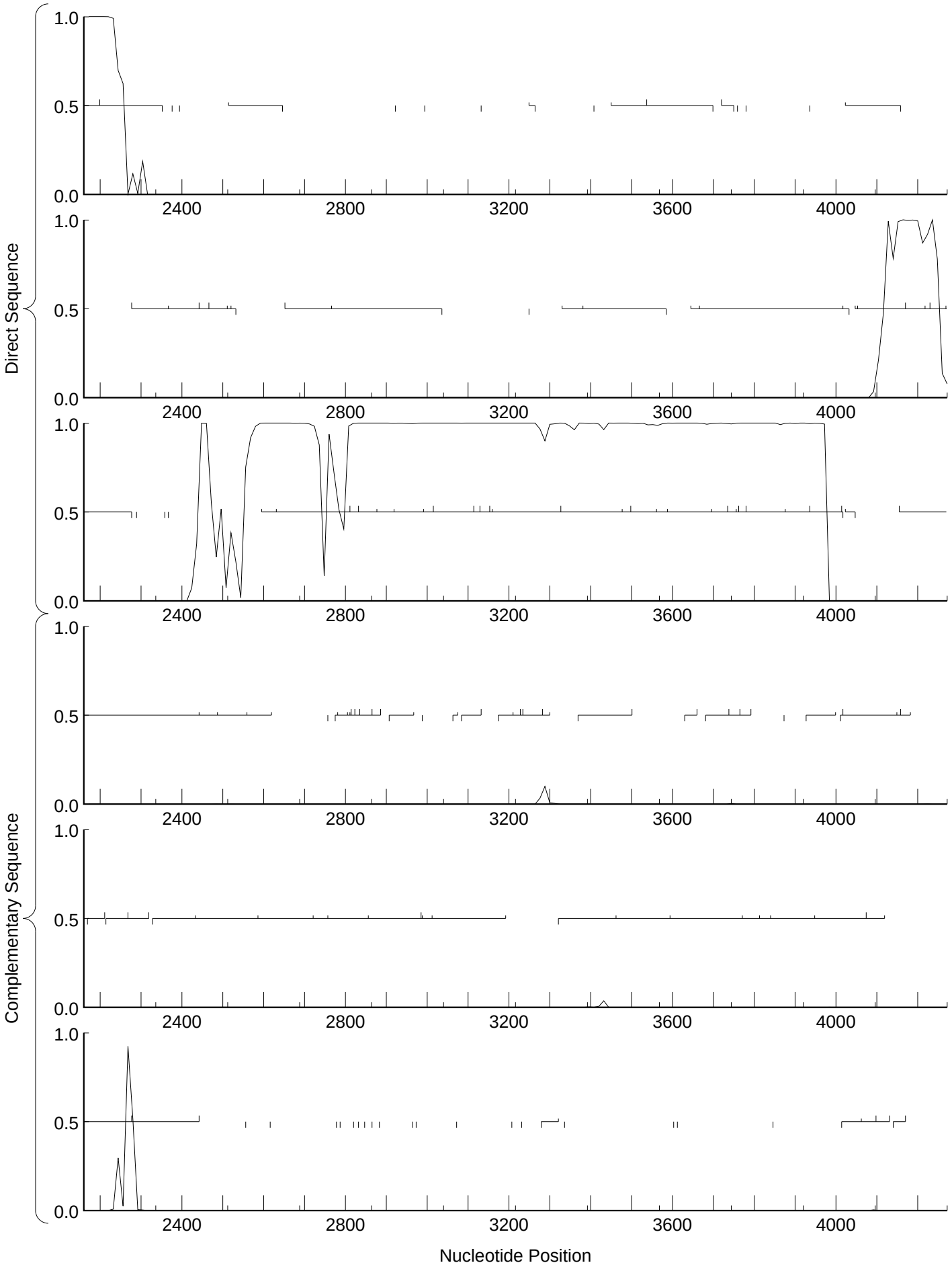

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

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

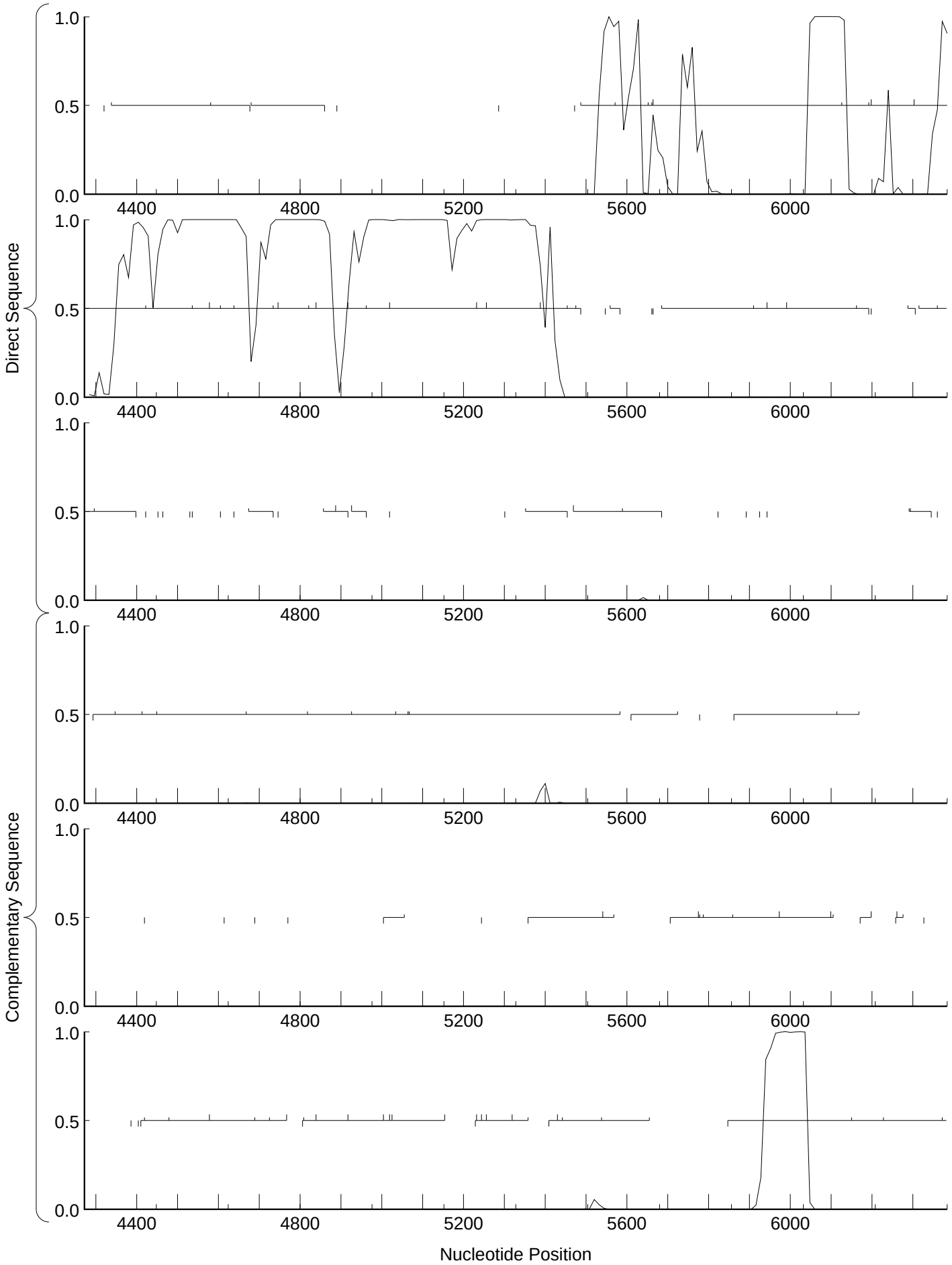
omplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 2/26



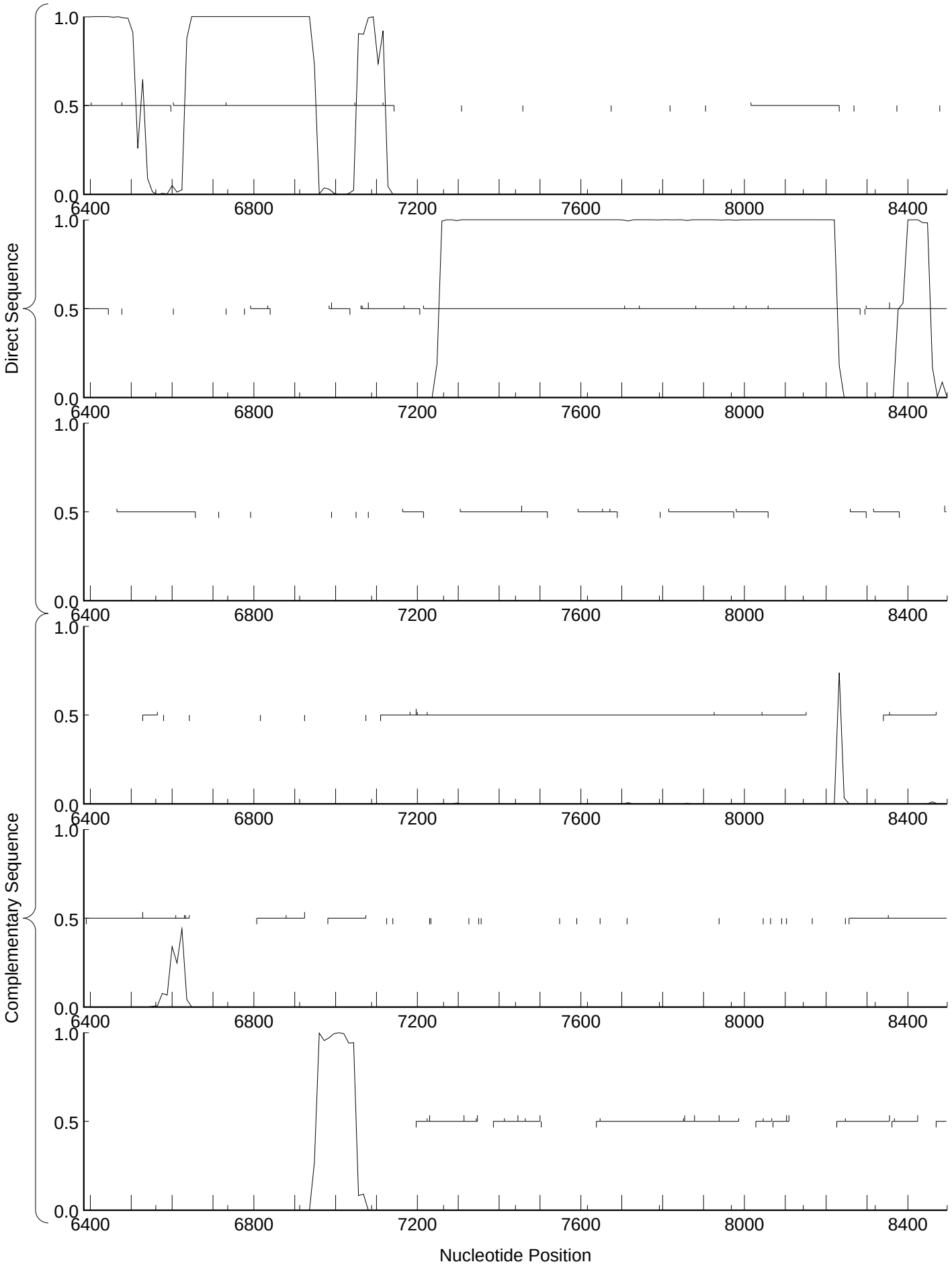
omplete sequence, 51571 bp including 11-base 3' overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 3/26



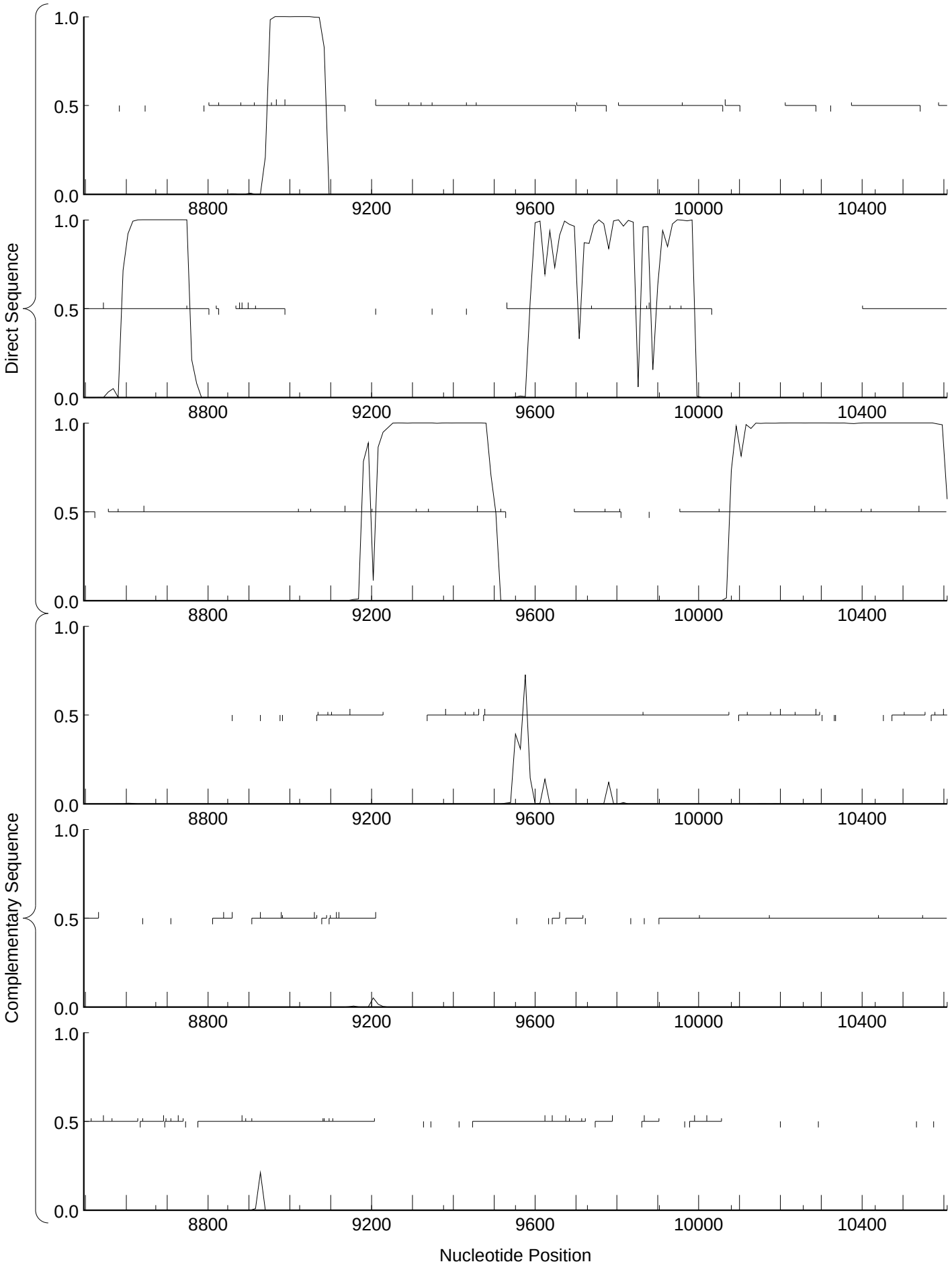
omplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 4/26



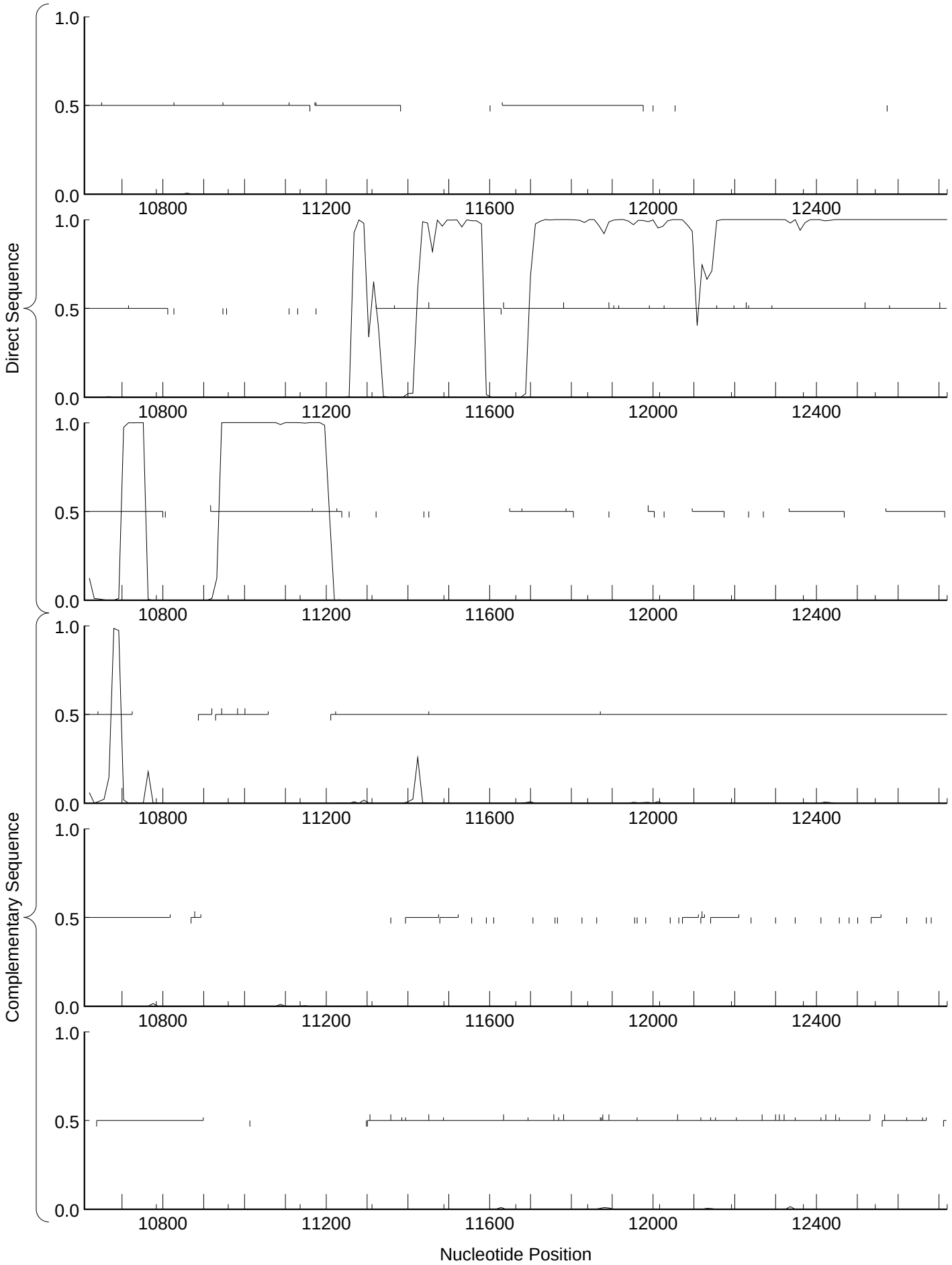
omplete sequence, 51571 bp including 11-base 3' overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 5/26



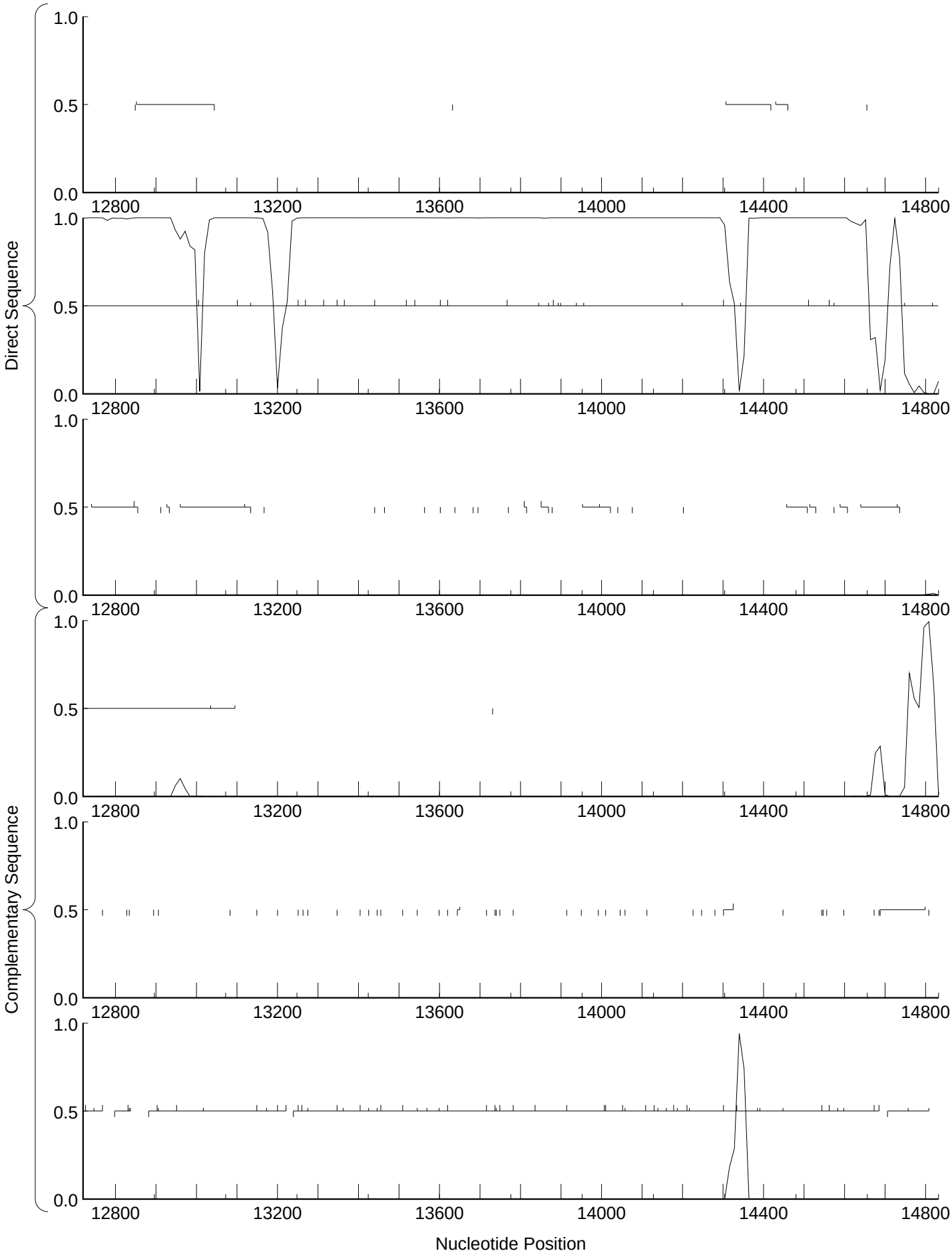
omplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 6/26



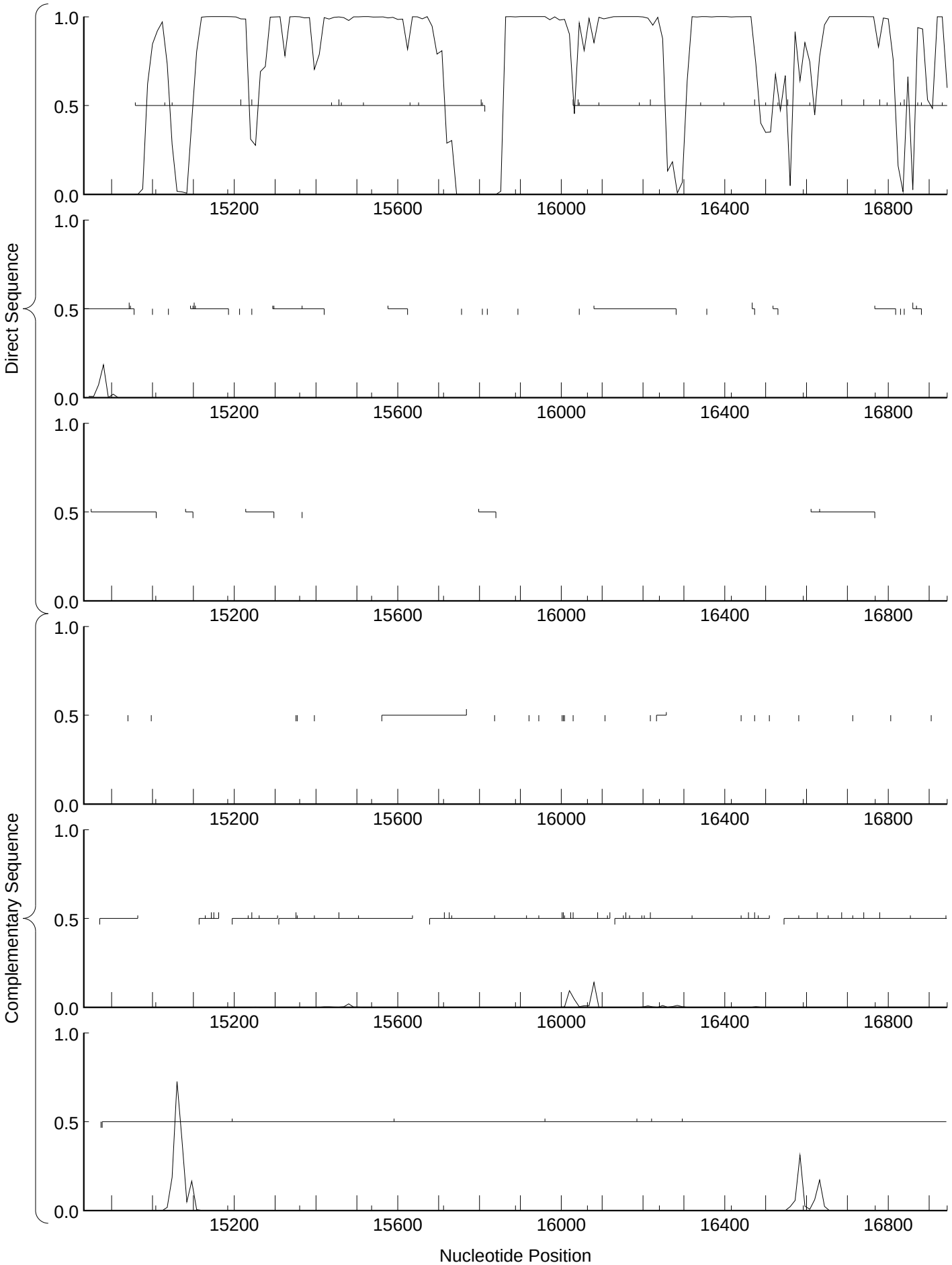
omplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 7/26

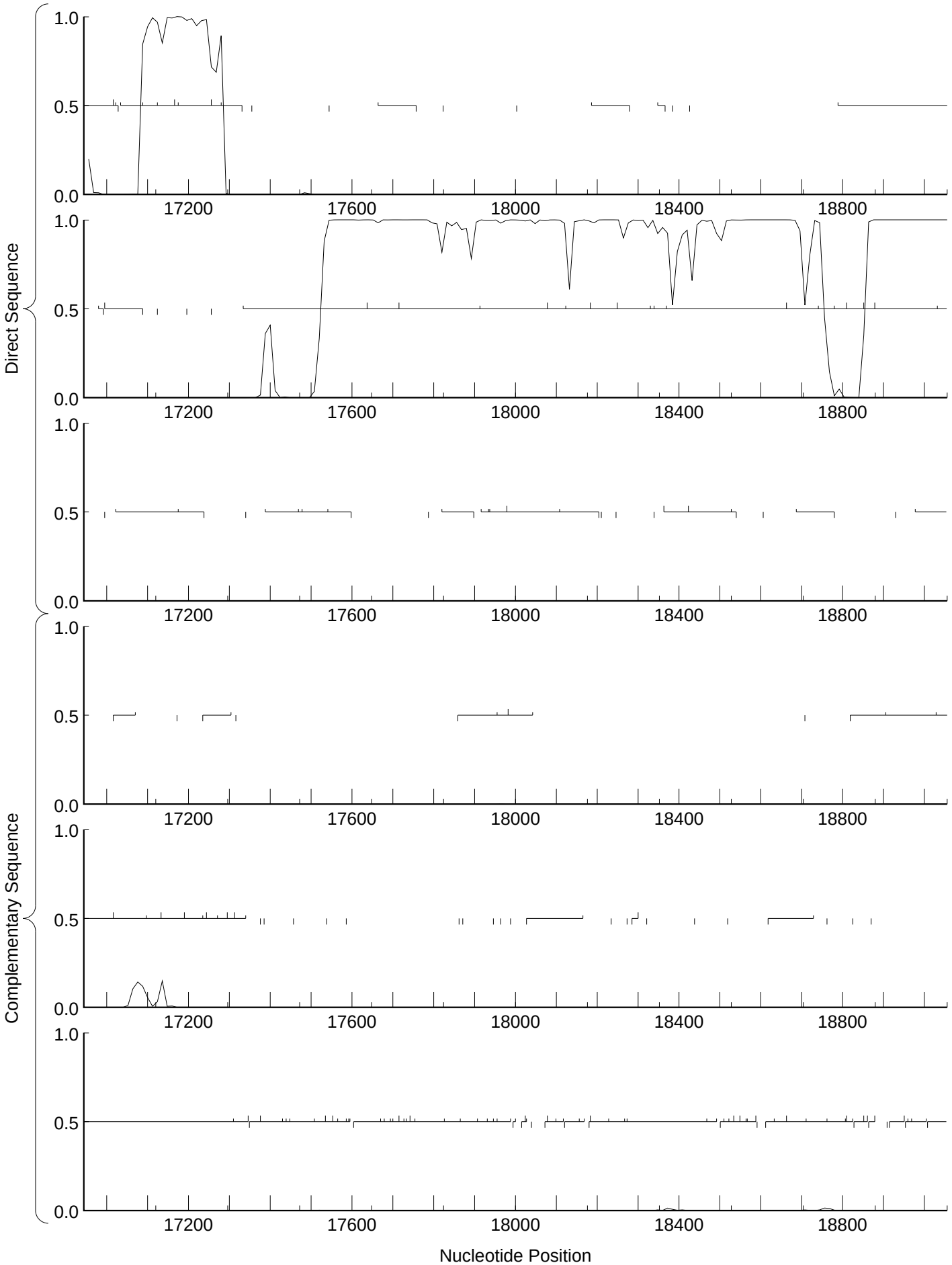


omplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 8/26

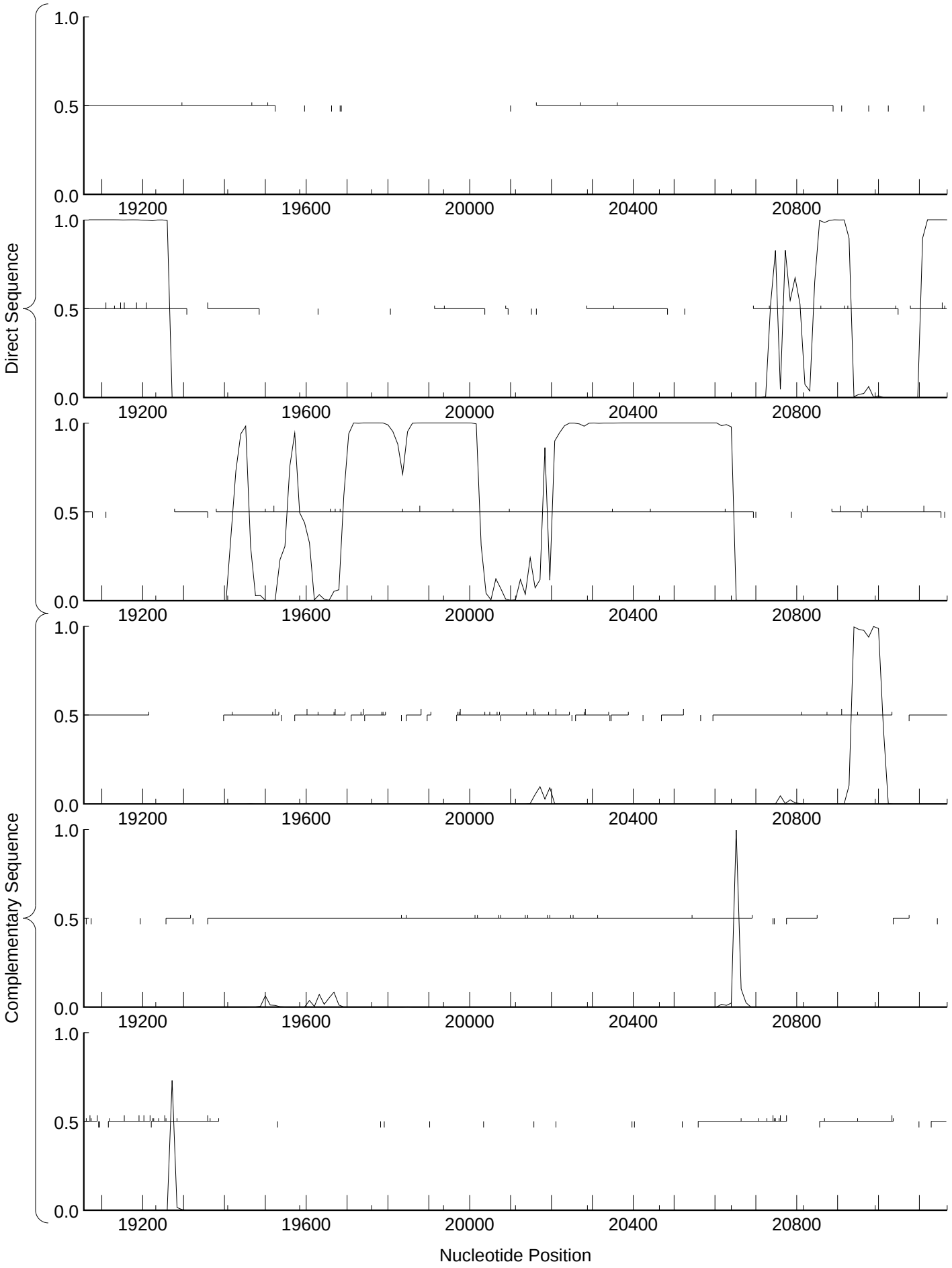


omplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 9/26

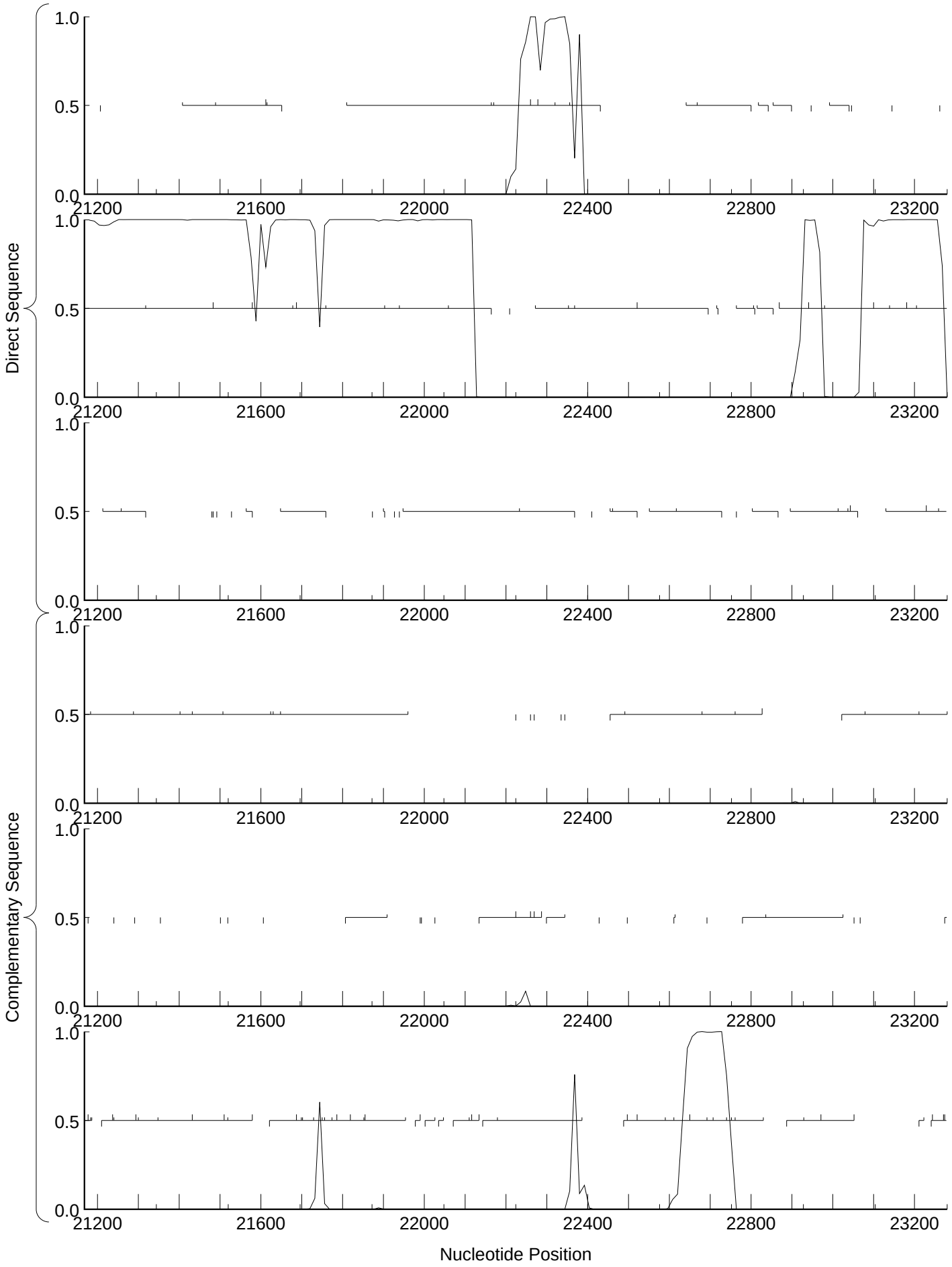



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
17200
17600
18000
18400
18800
Nucleotide Position

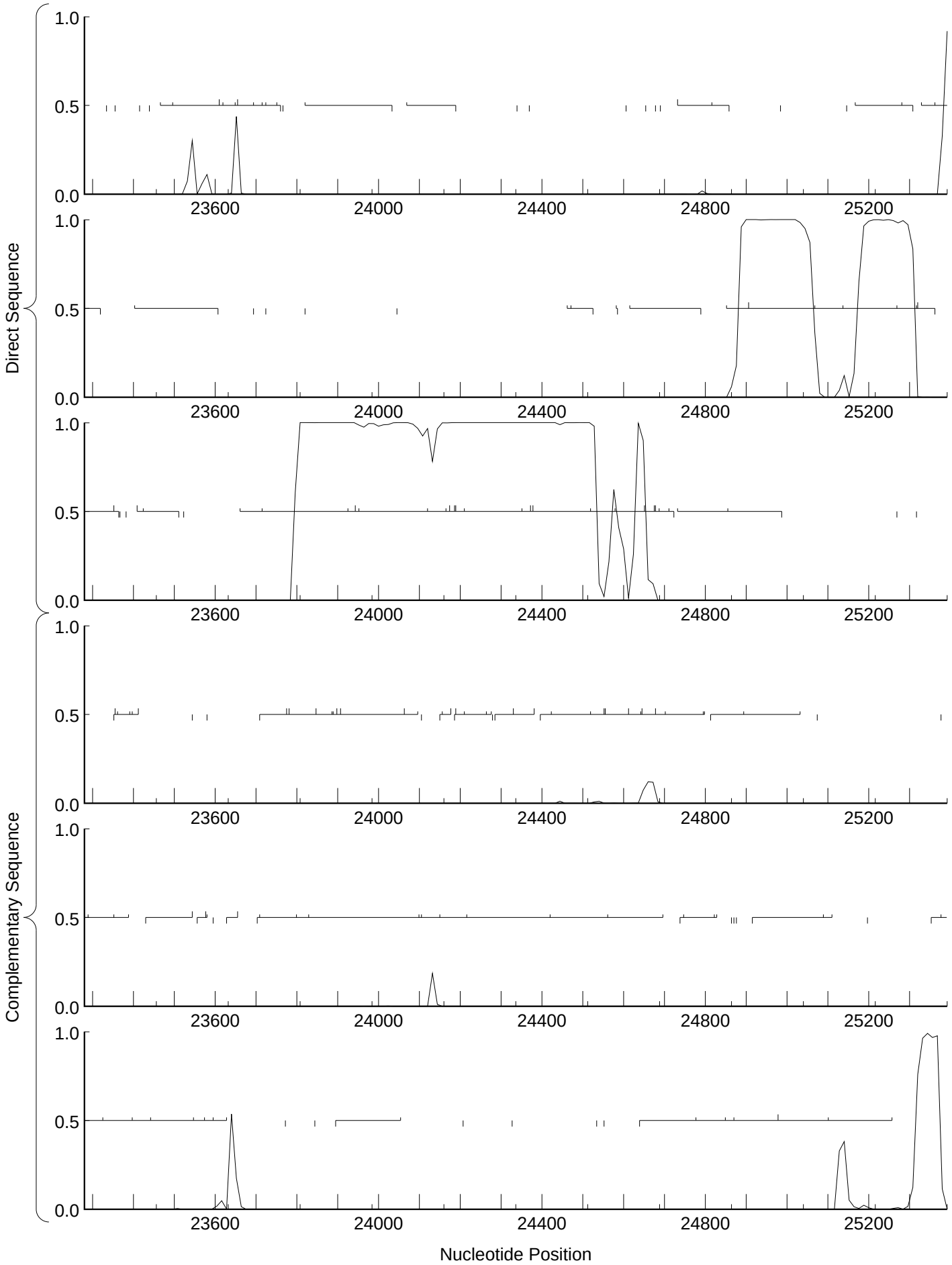
mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 11/26



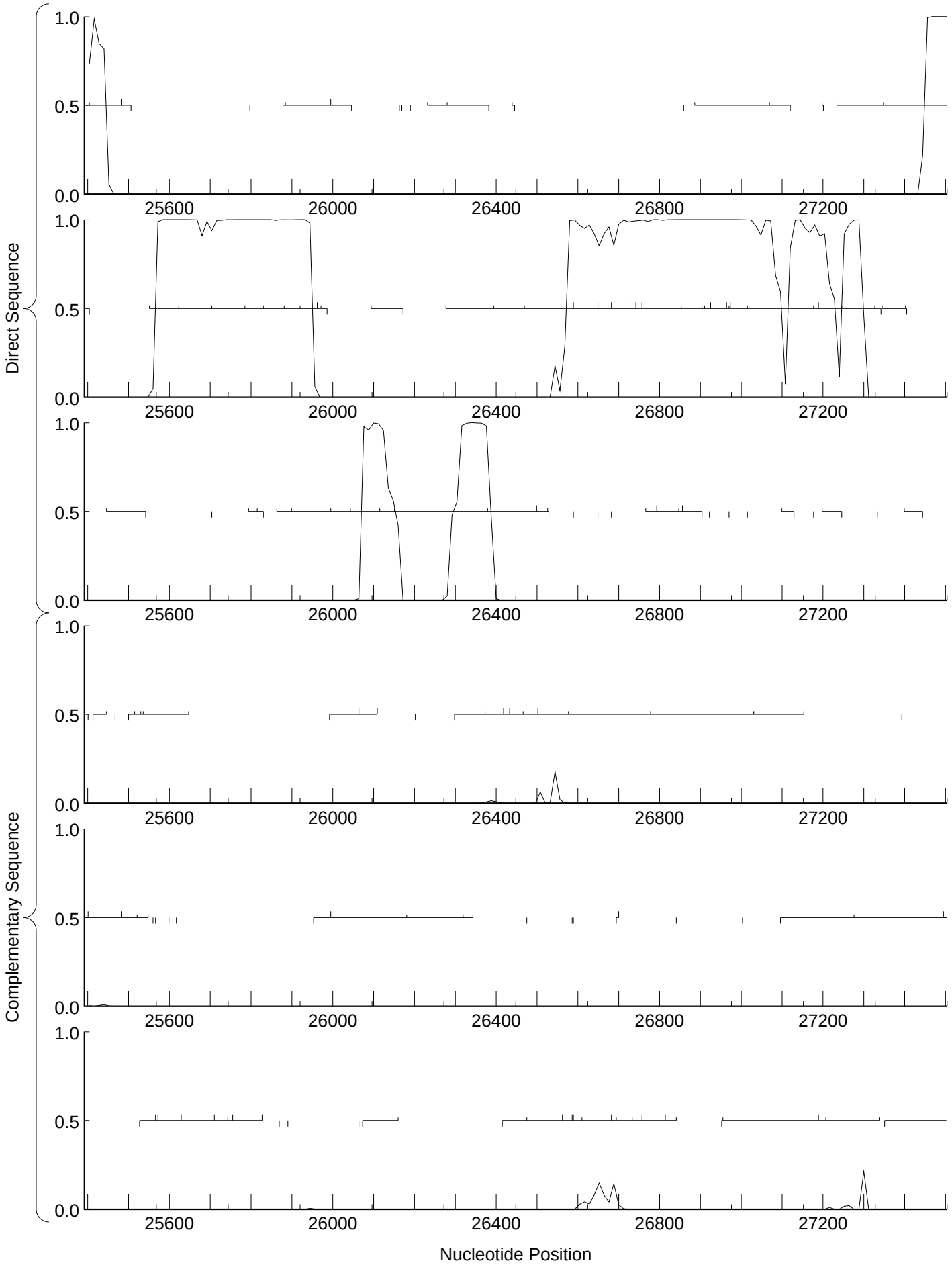
mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 12/26



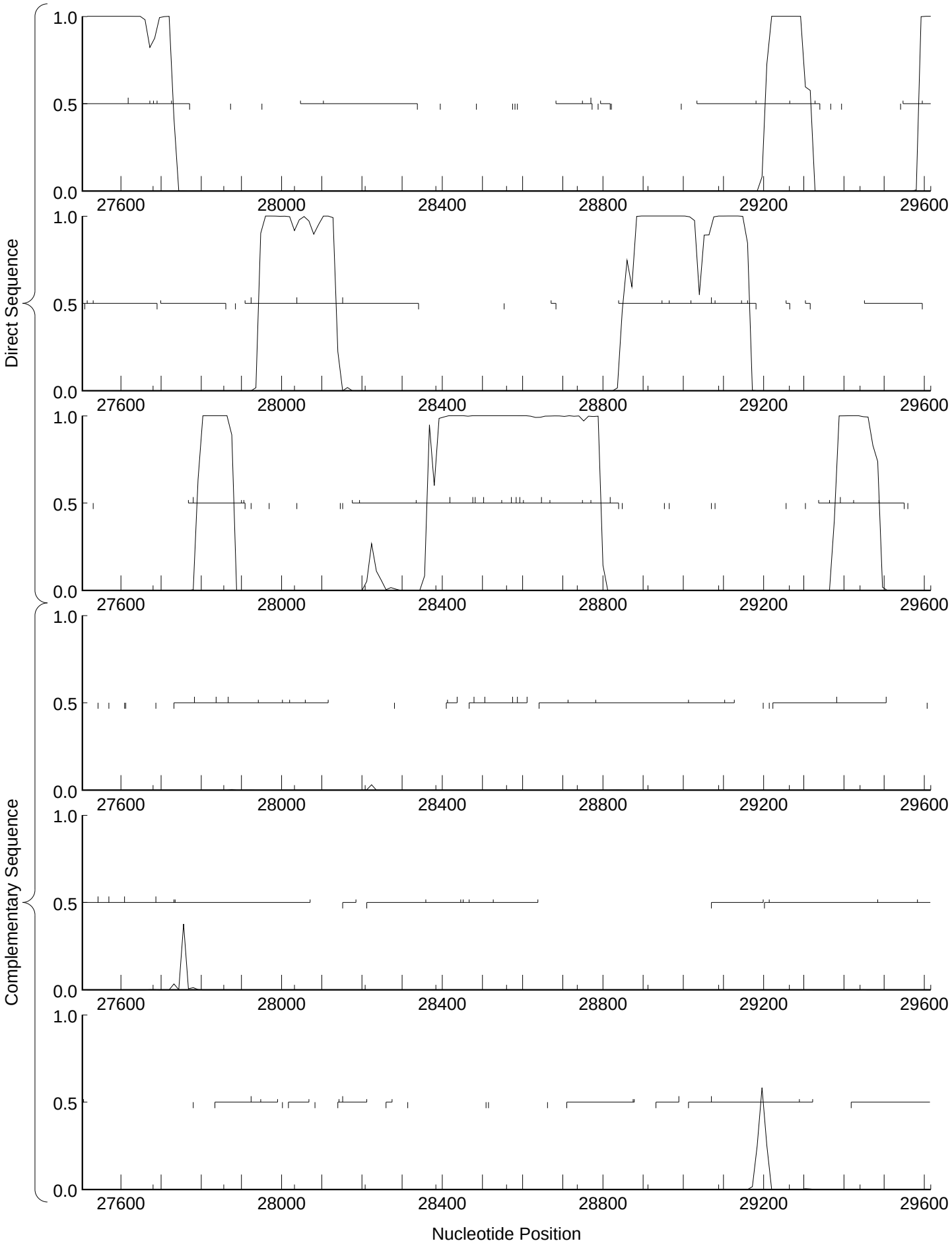
mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 13/26

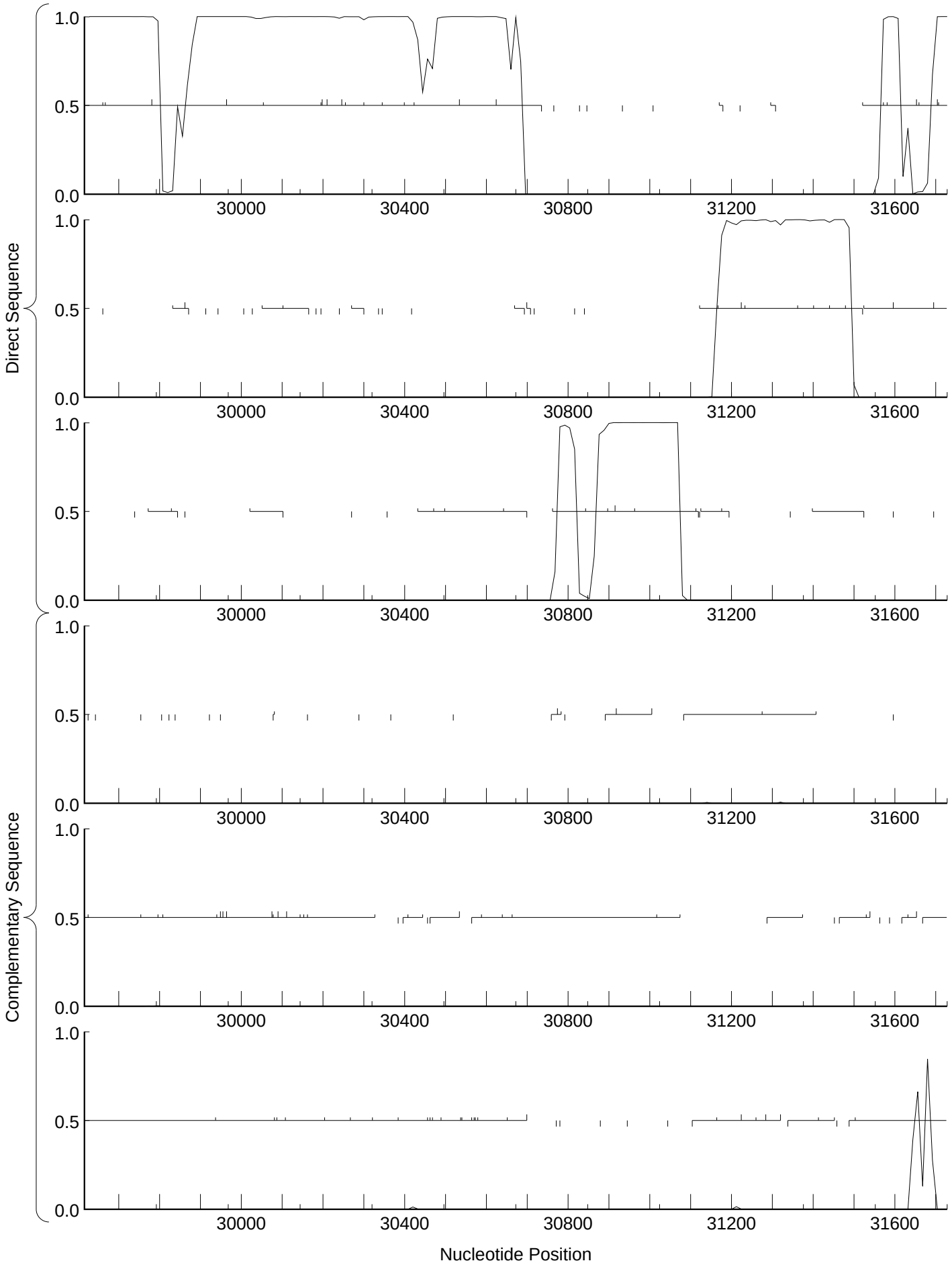


mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 14/26

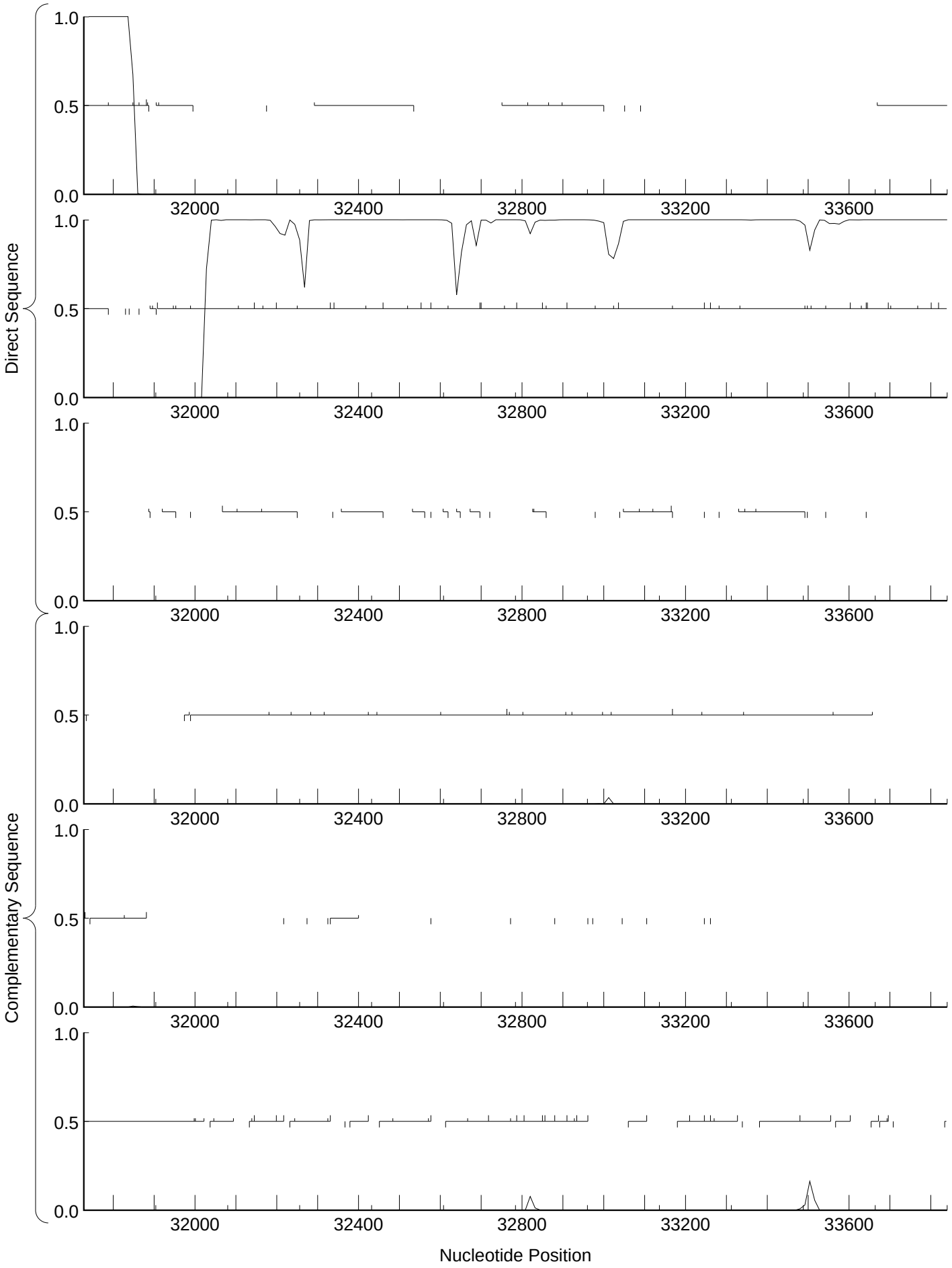


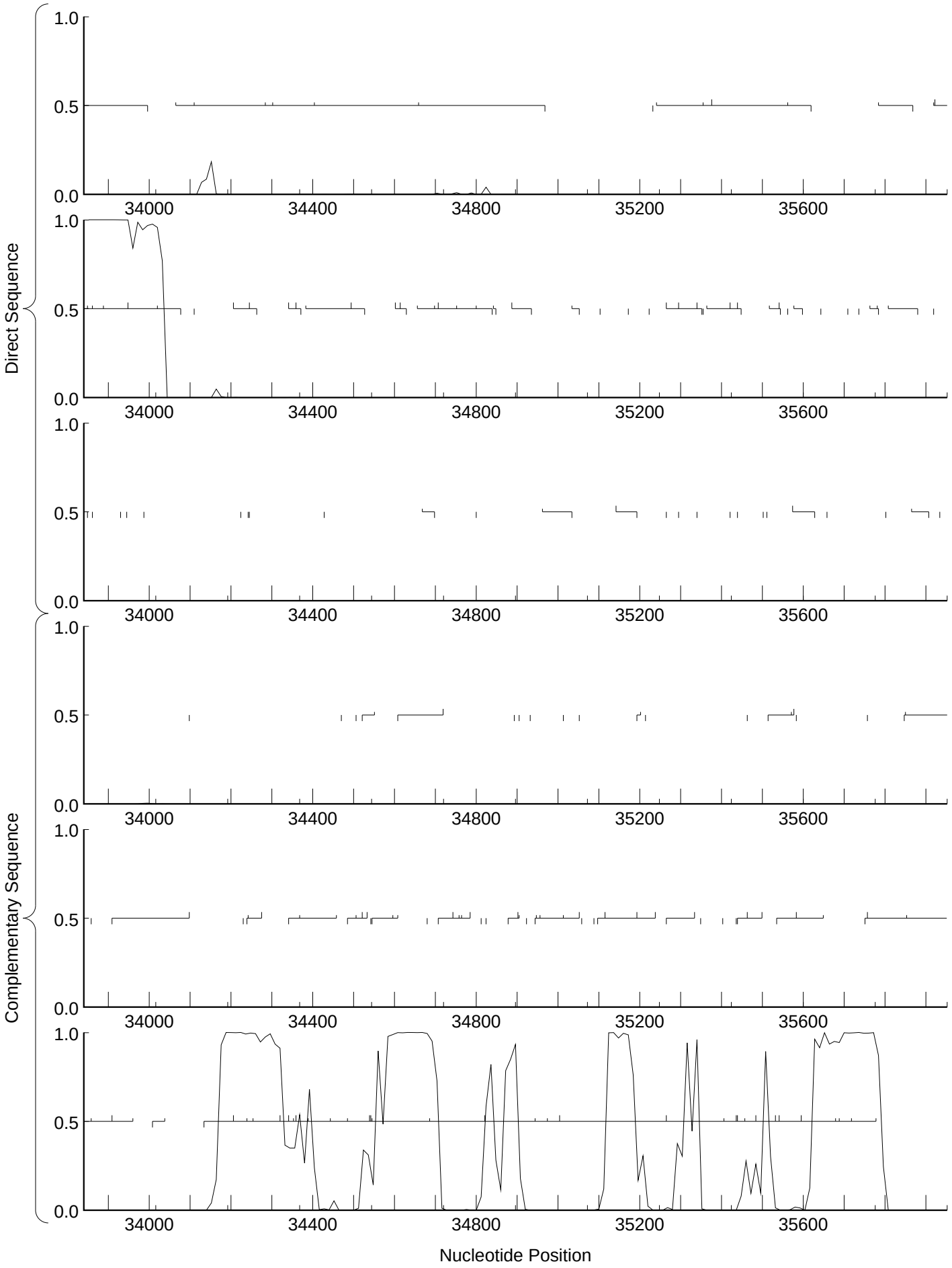
mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 15/26



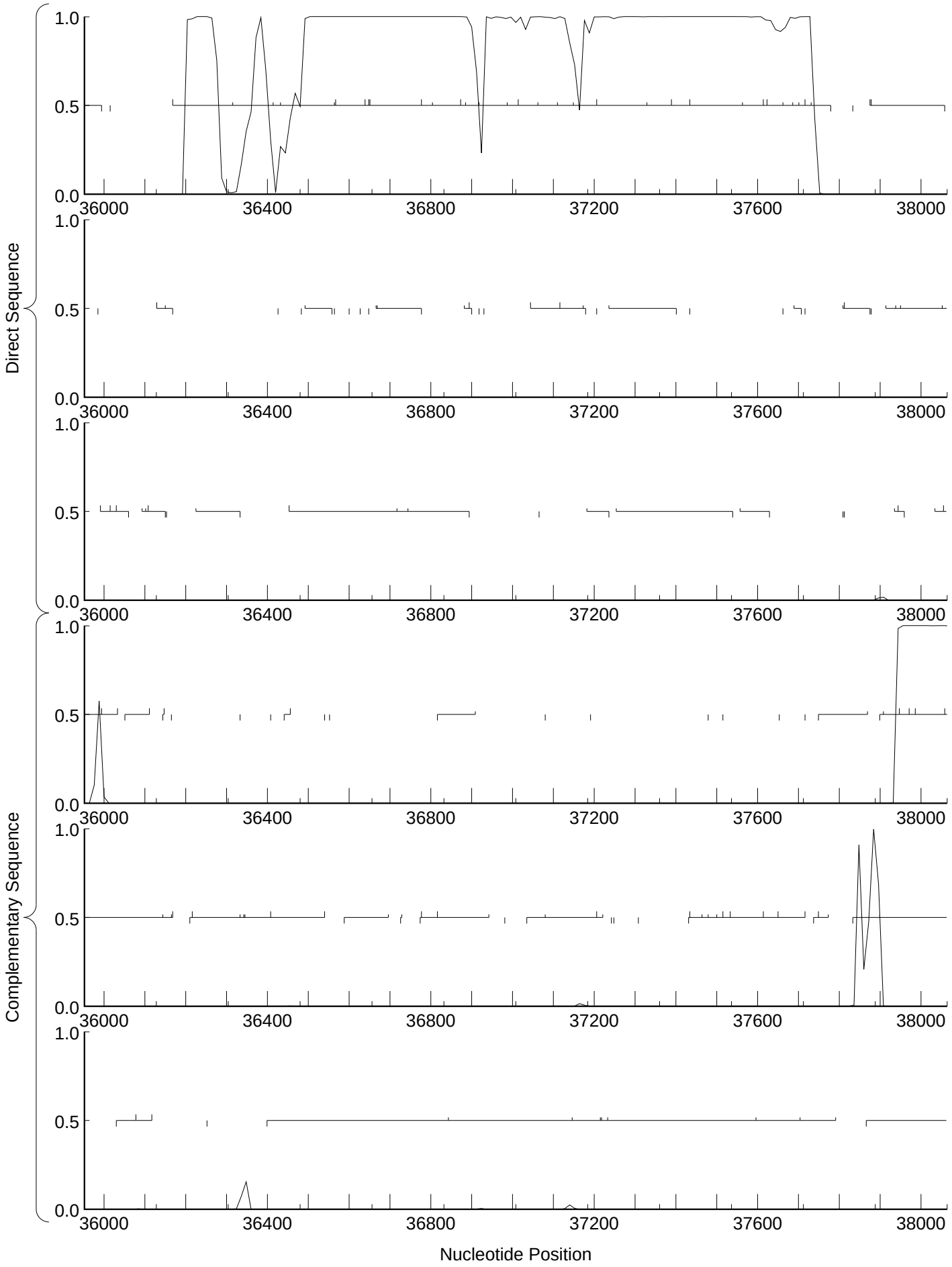

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
30000
30400
30800
31200
31600
Nucleotide Position

mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 17/26

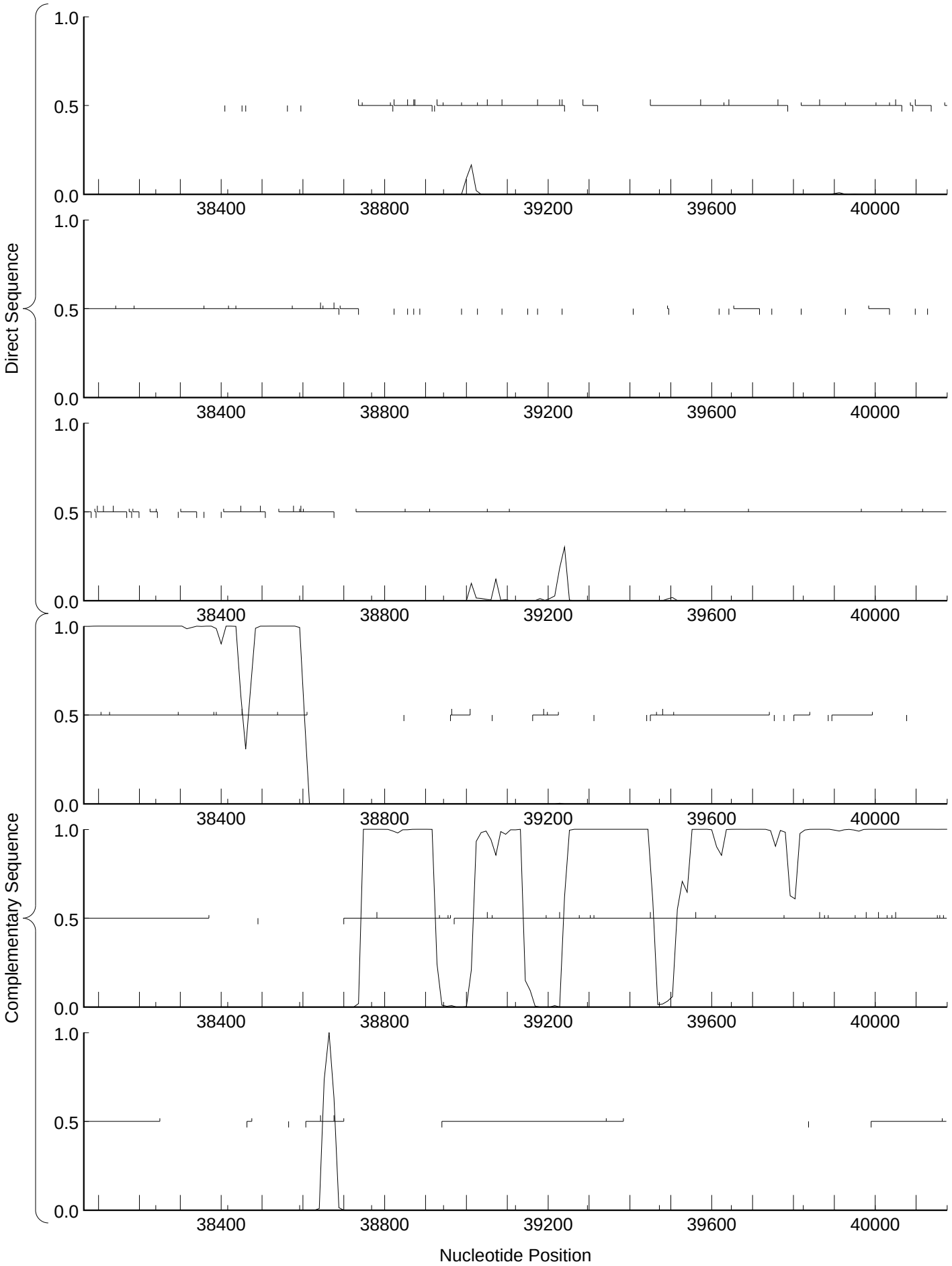



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

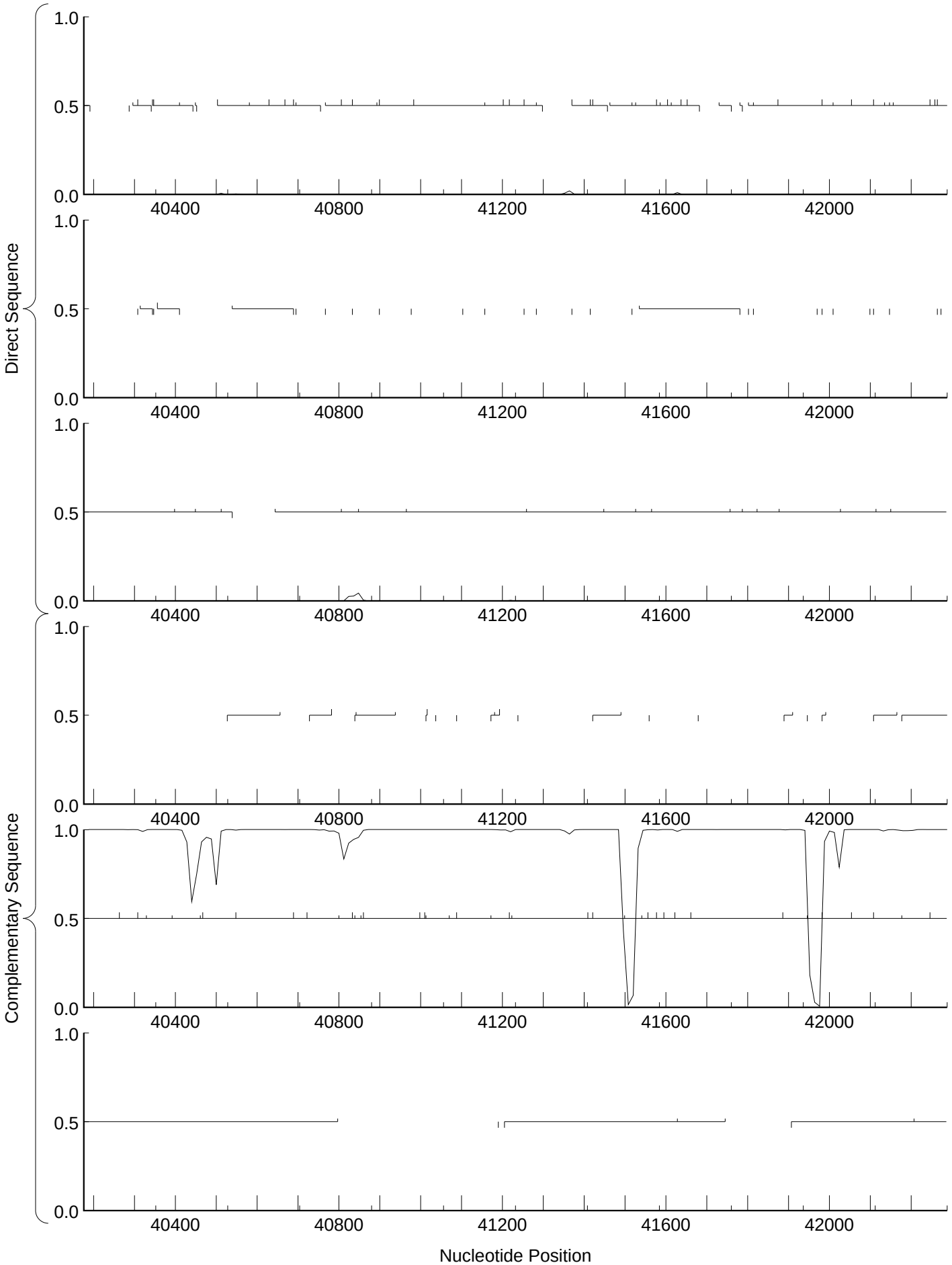
mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 19/26



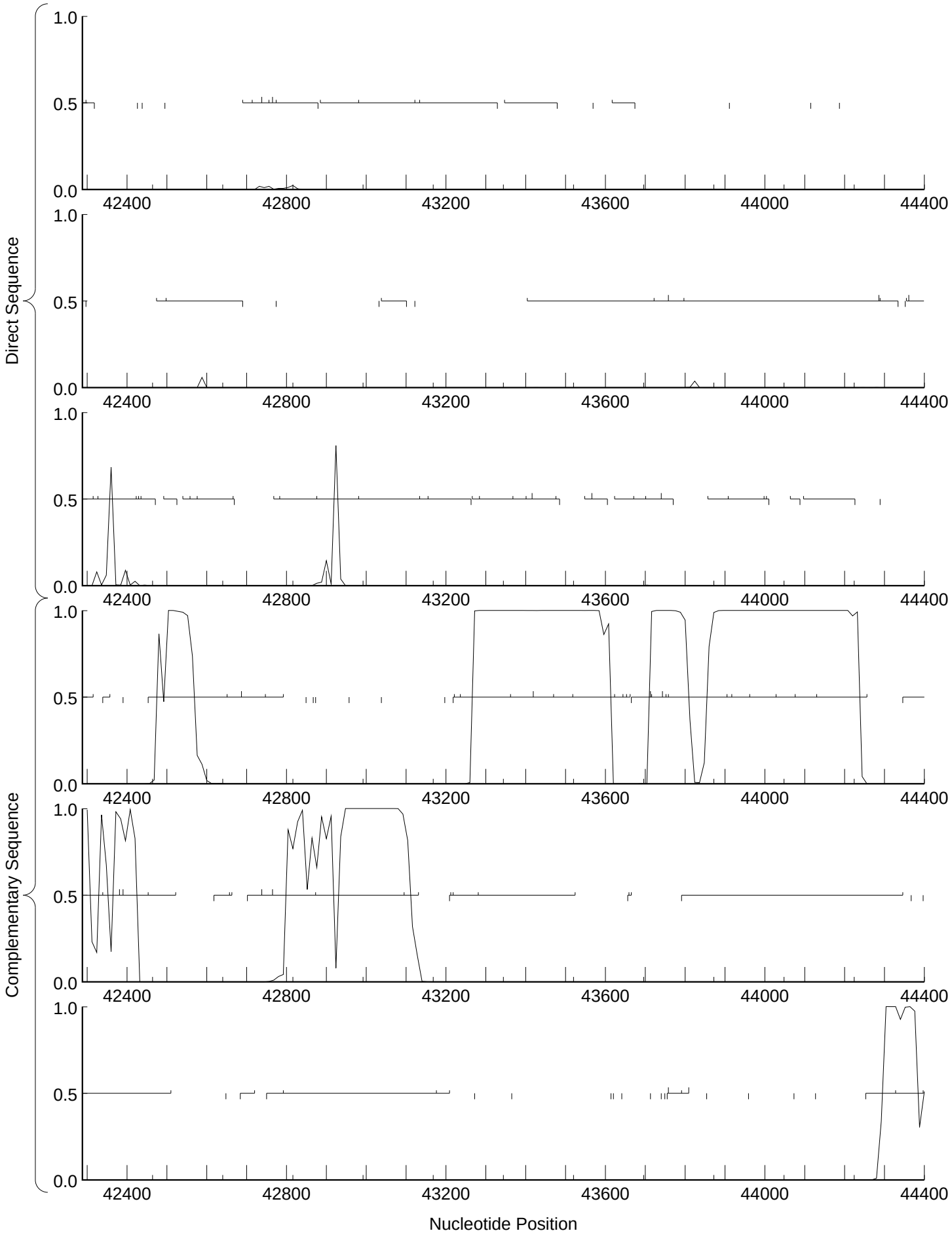
mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 20/26

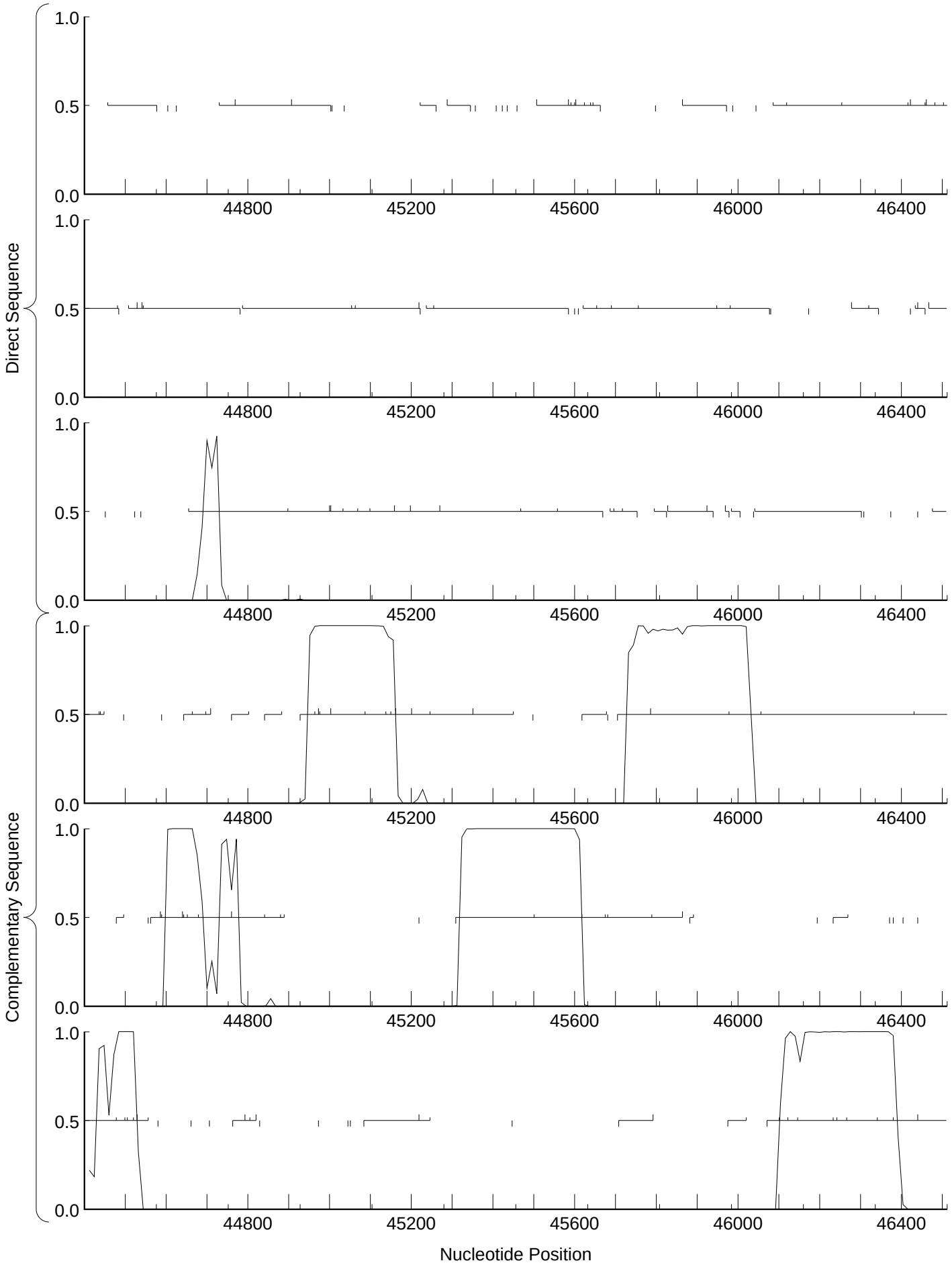


mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 21/26

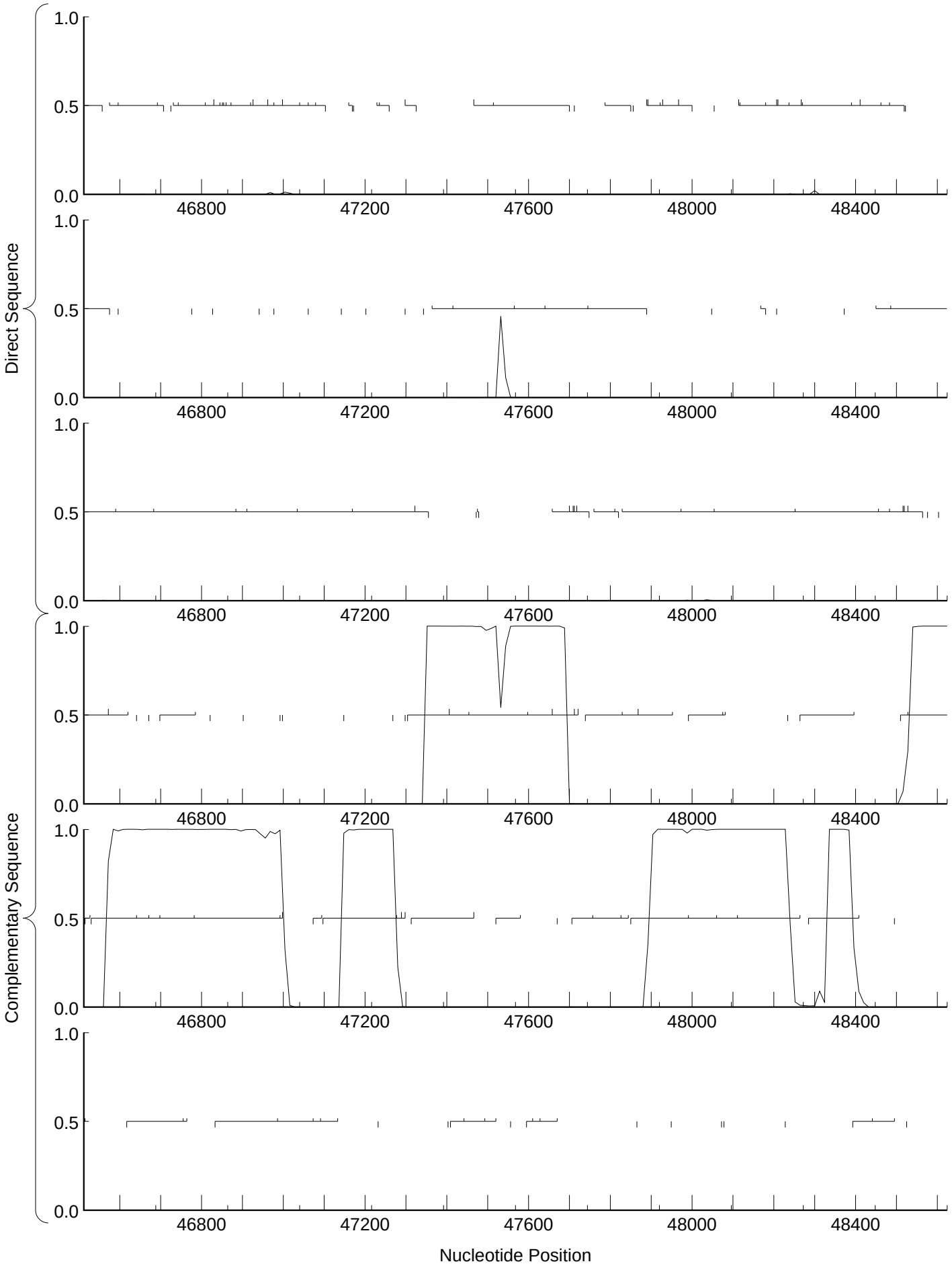


mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 22/26

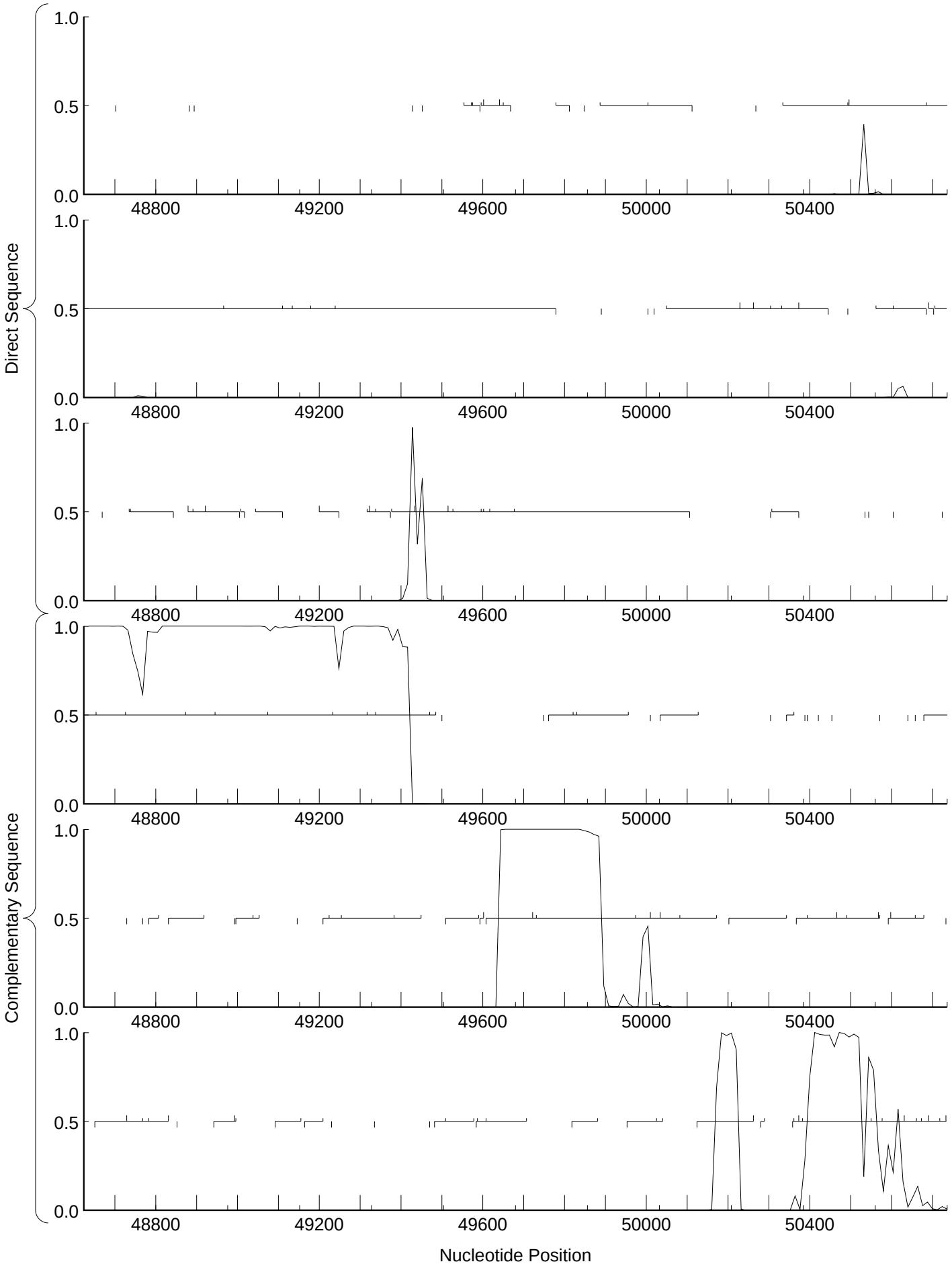



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
44800
45200
45600
46000
46400
Nucleotide Position

mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 24/26



mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 25/26



mplete sequence, 51571 bp including 11-base 3’ overhang (CGGCCAGTGAT), Cluster BD1, Order 4, Window 96, Step 12, 26/26

