GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage ExplosioNervosa complete sequence, 53014 bp including 10-base 3′ overha
Analysis Date    :  3/29/24 at 7:23:47
Pages            :  27
Sequence Length  :  53014 bp
GC Content       :   62.54%

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

PS-Version       :  1.2

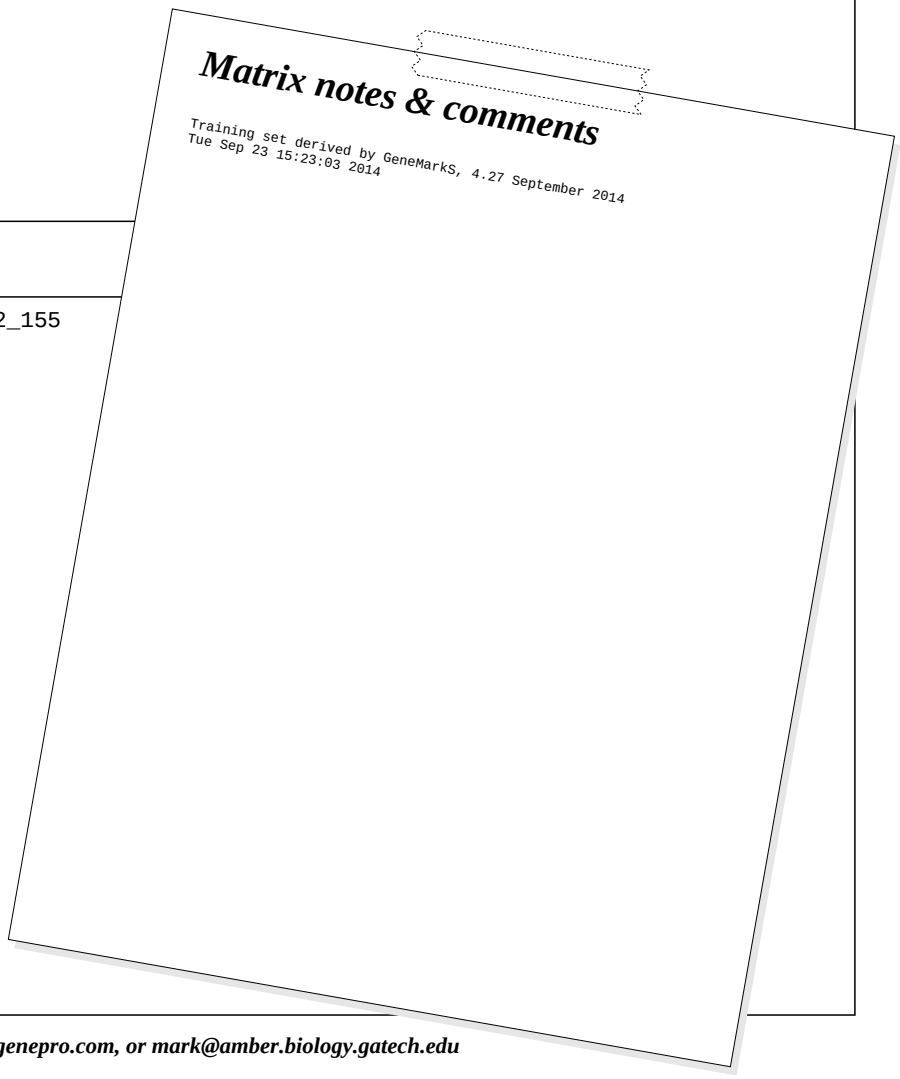
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
```

MATRIX INFORMATION

```
Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

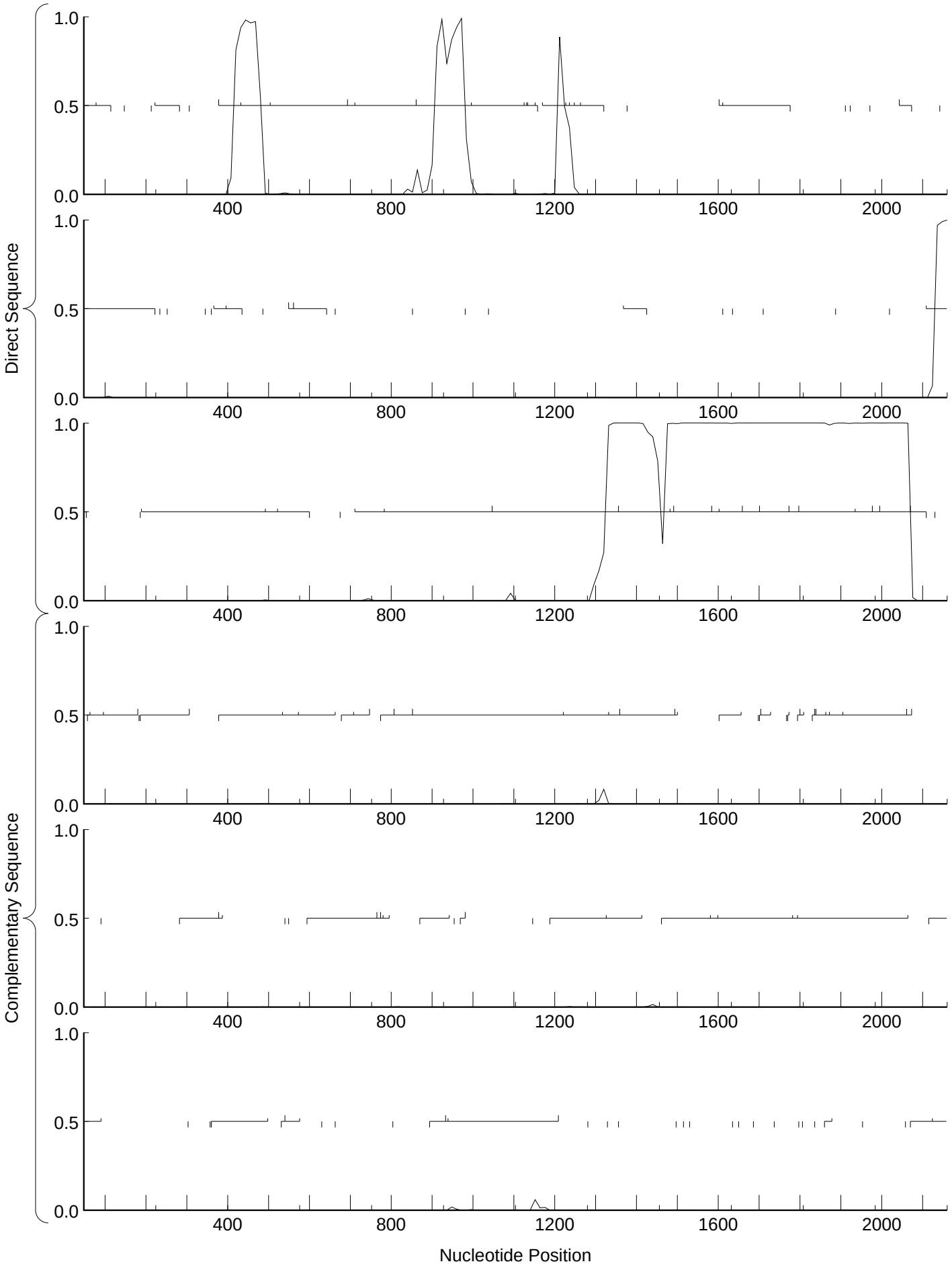
Order      :  4


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

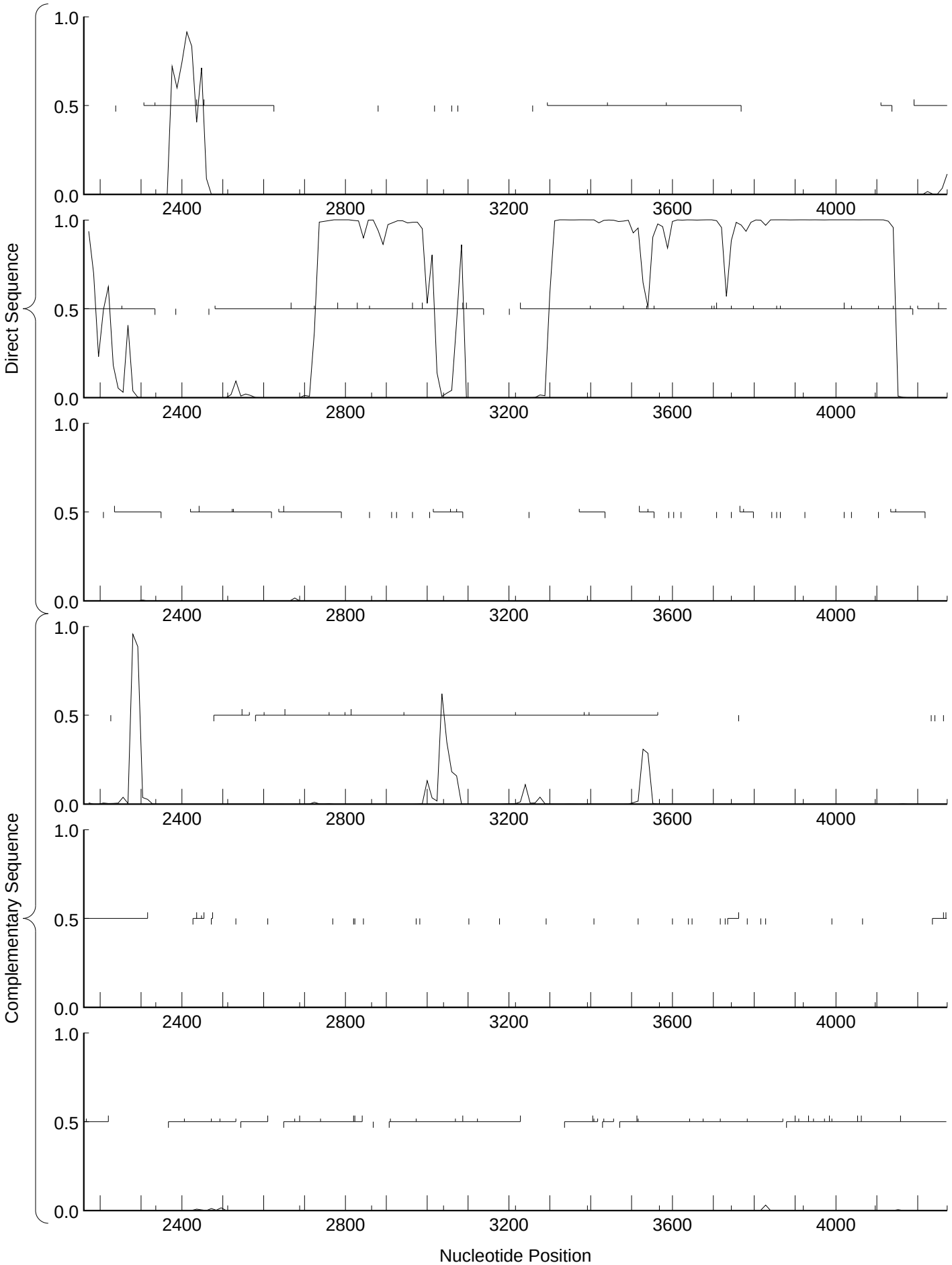


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

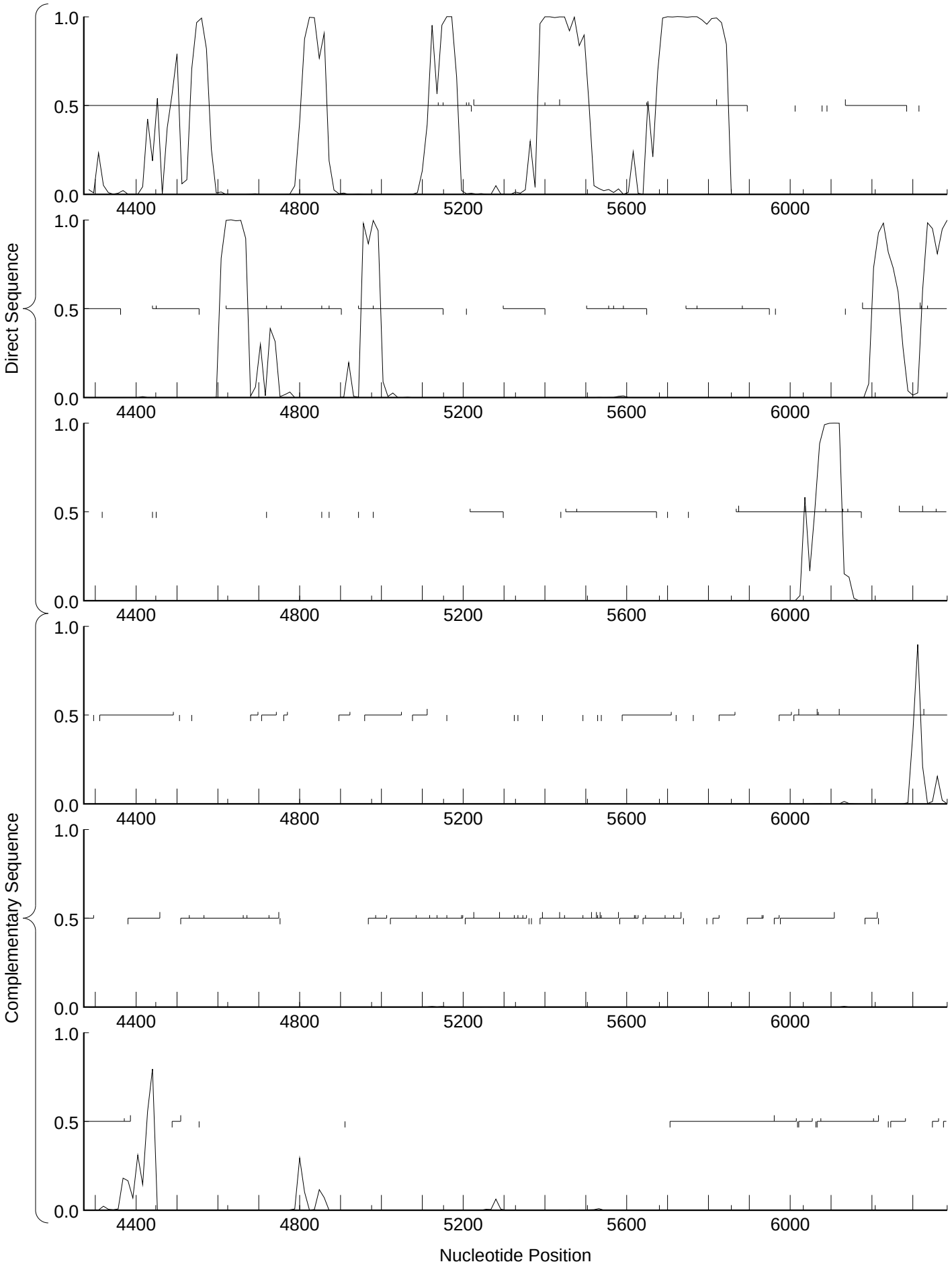
sa complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 2/27



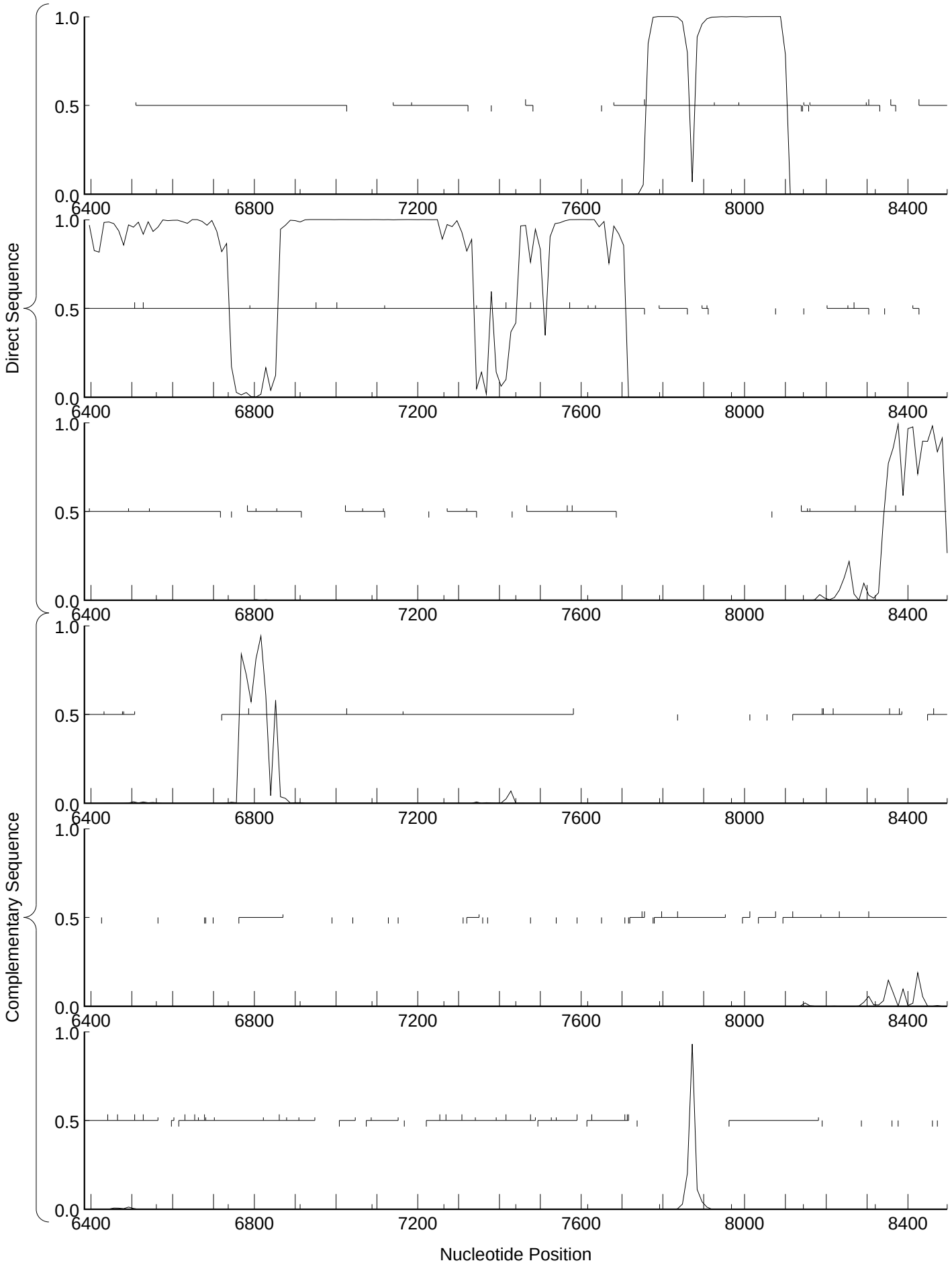
sa complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 3/27



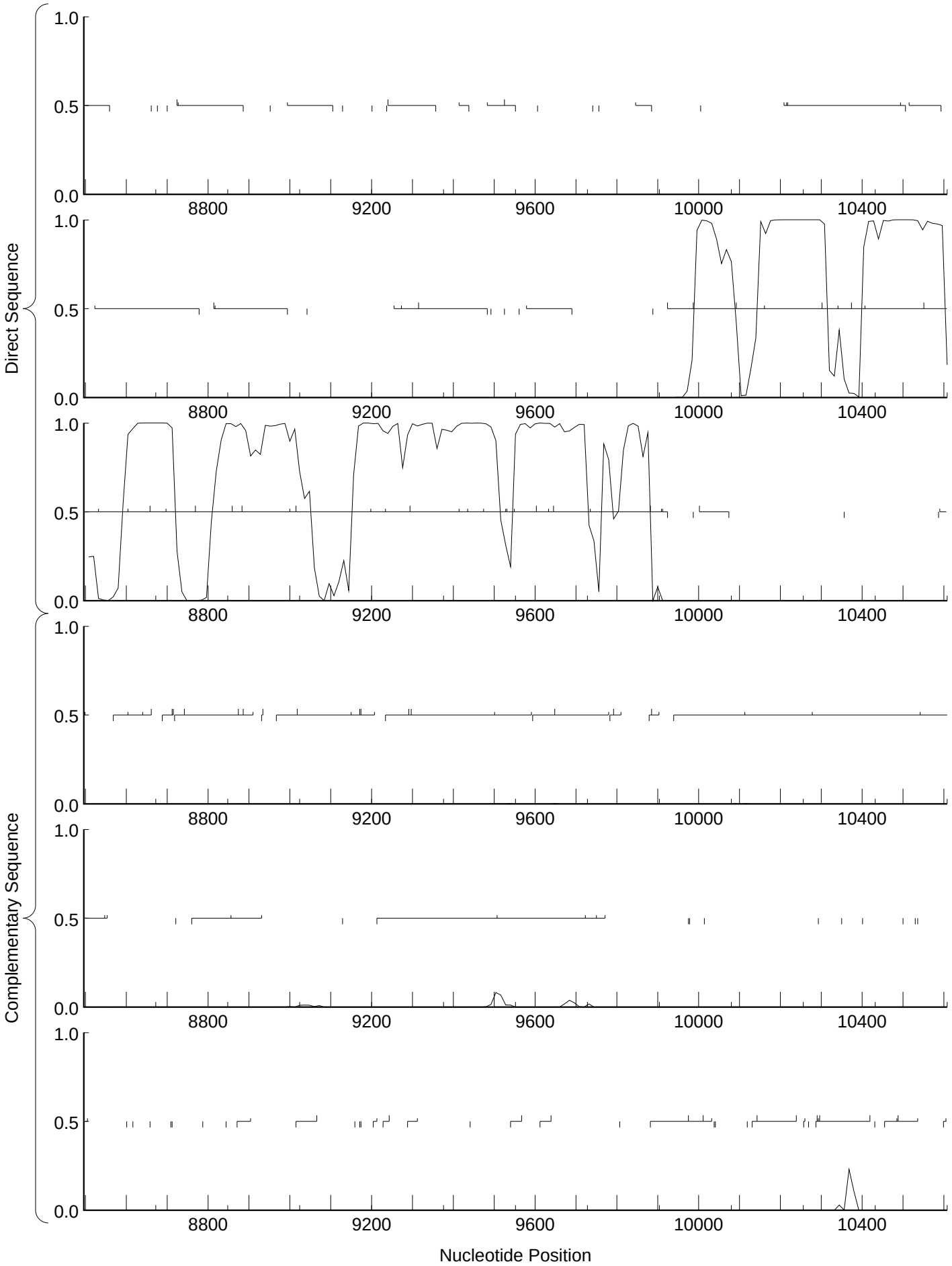
sa complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 4/27



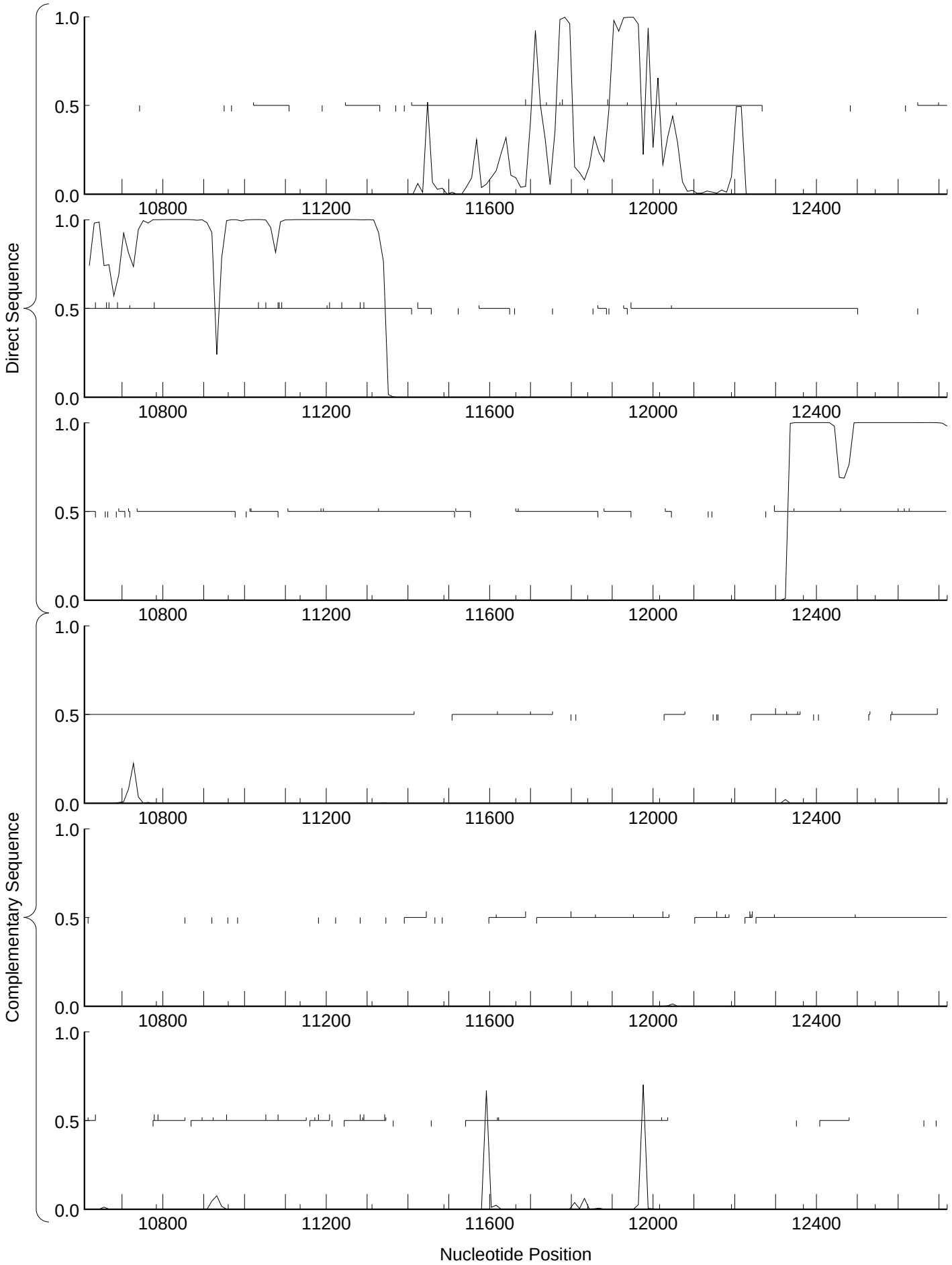
sa complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 5/27



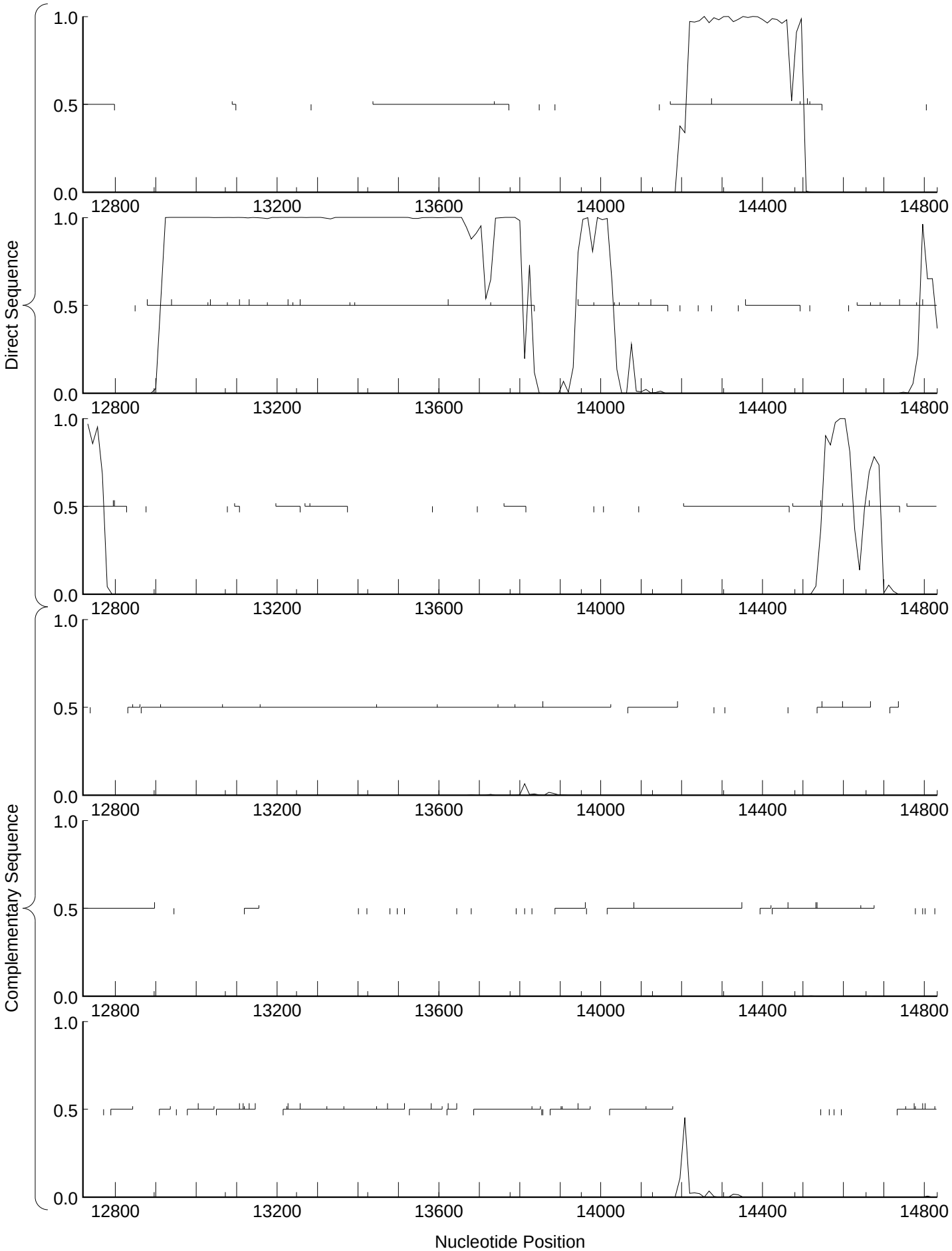
sa complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 6/27



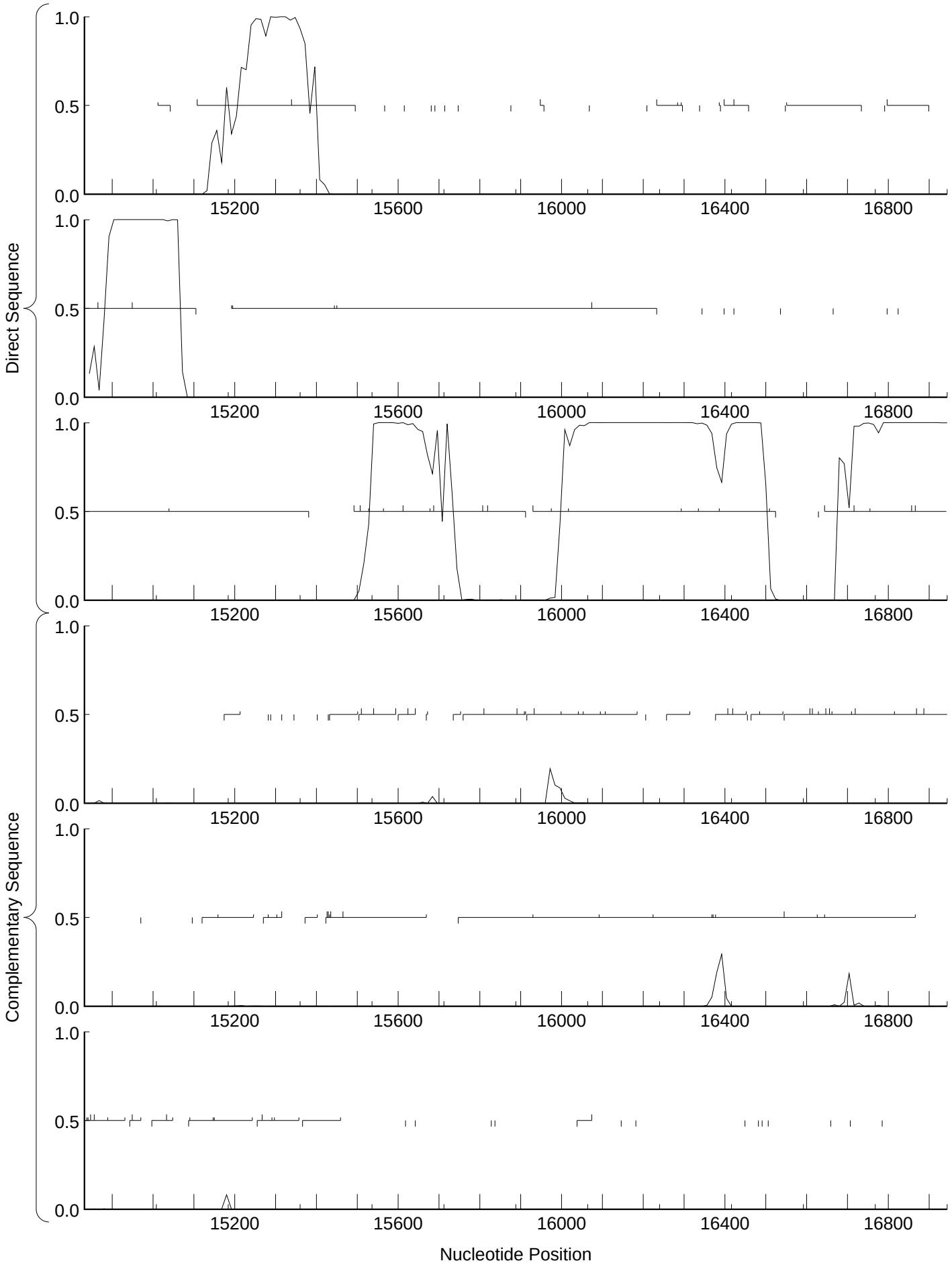
sa complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 7/27



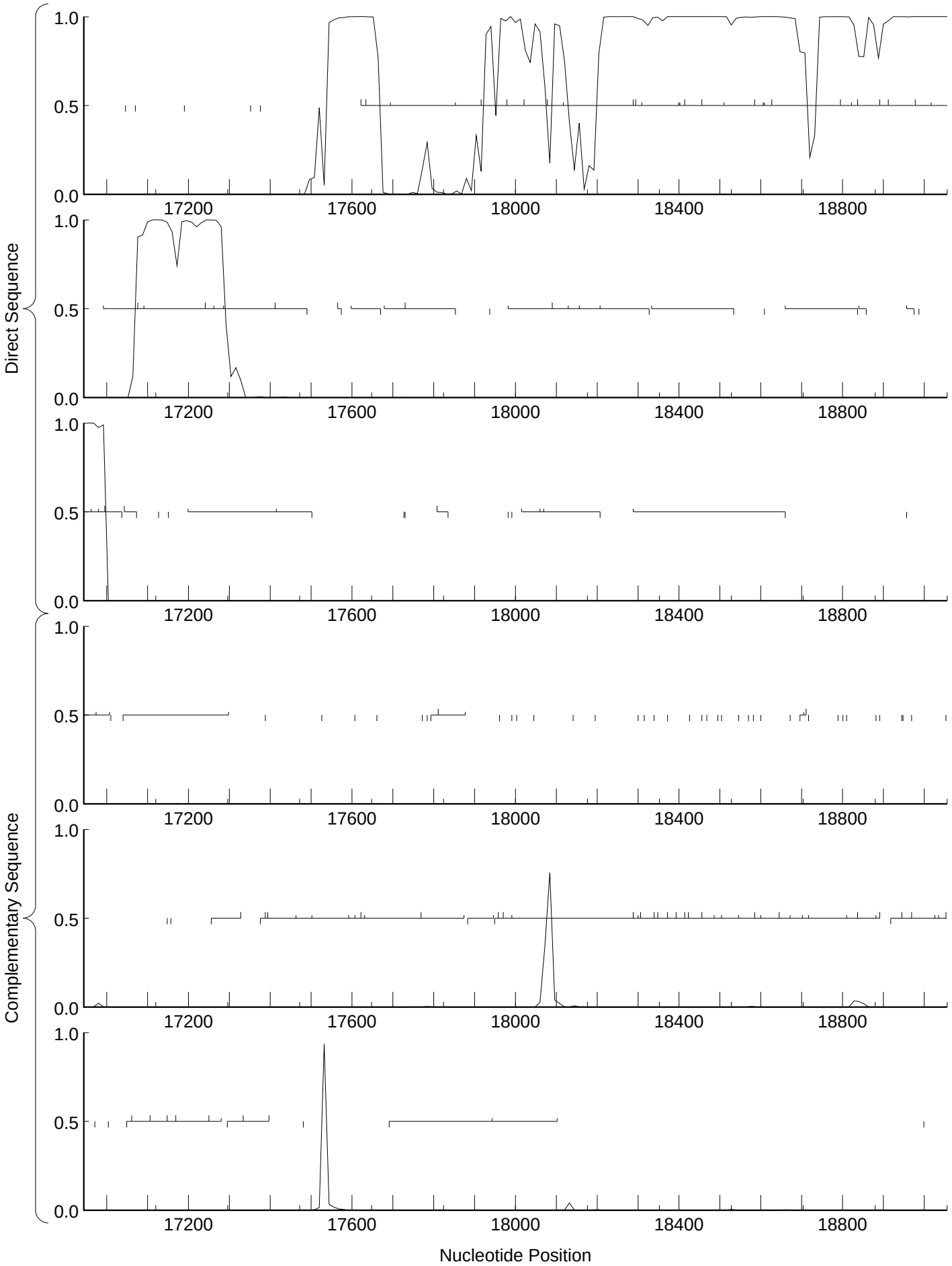
sa complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 8/27



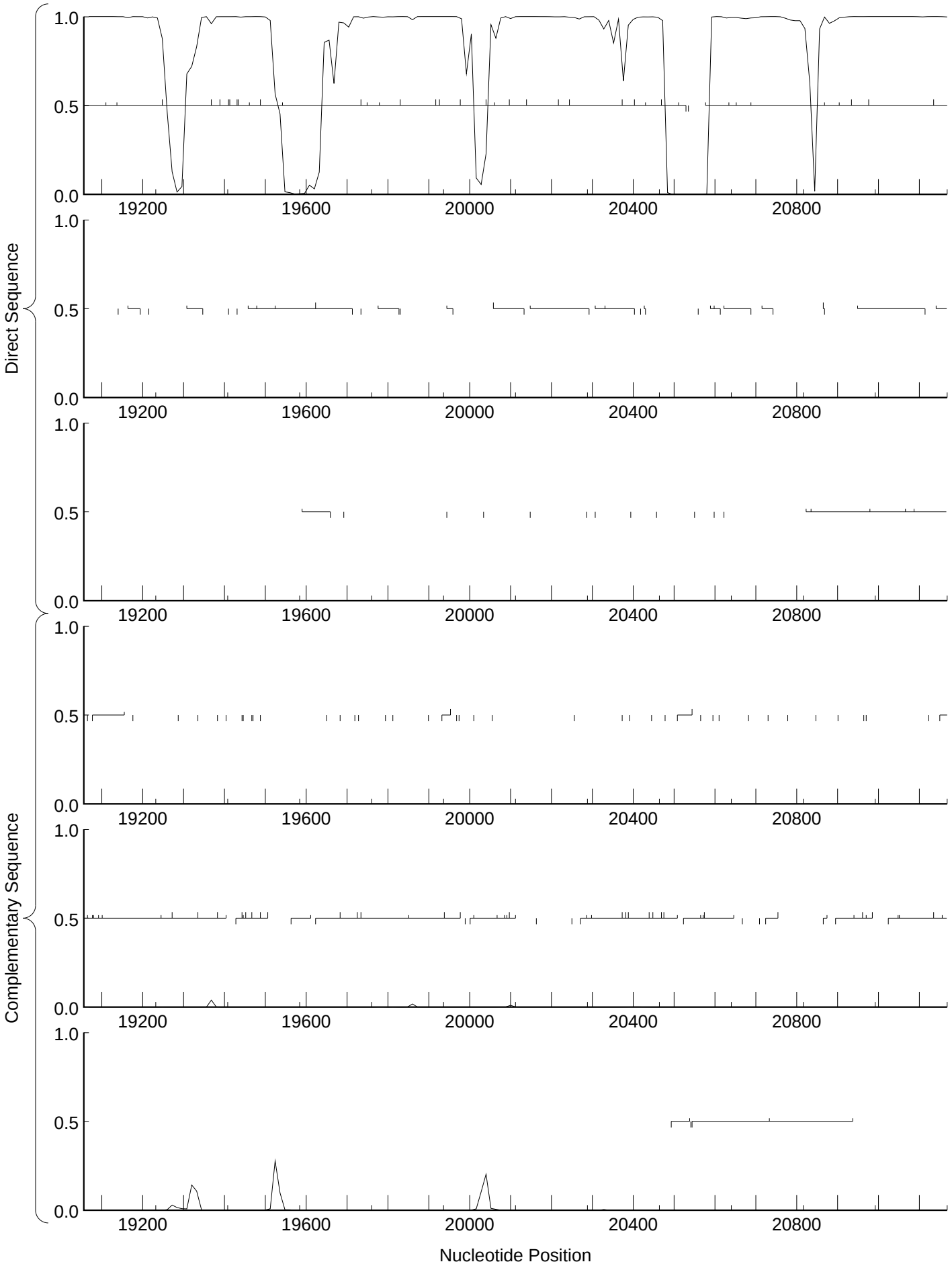
sa complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 9/27



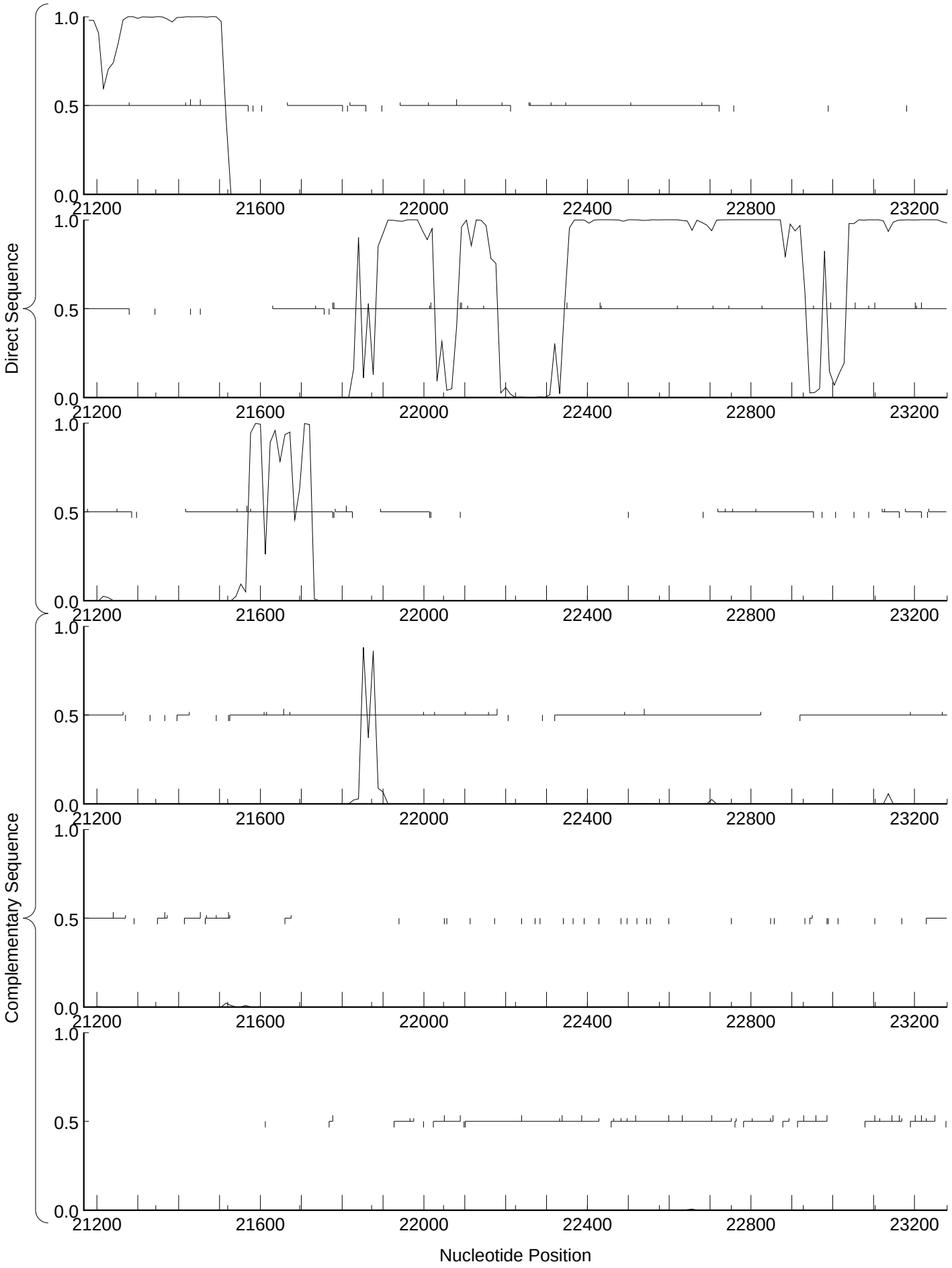
a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 10/27



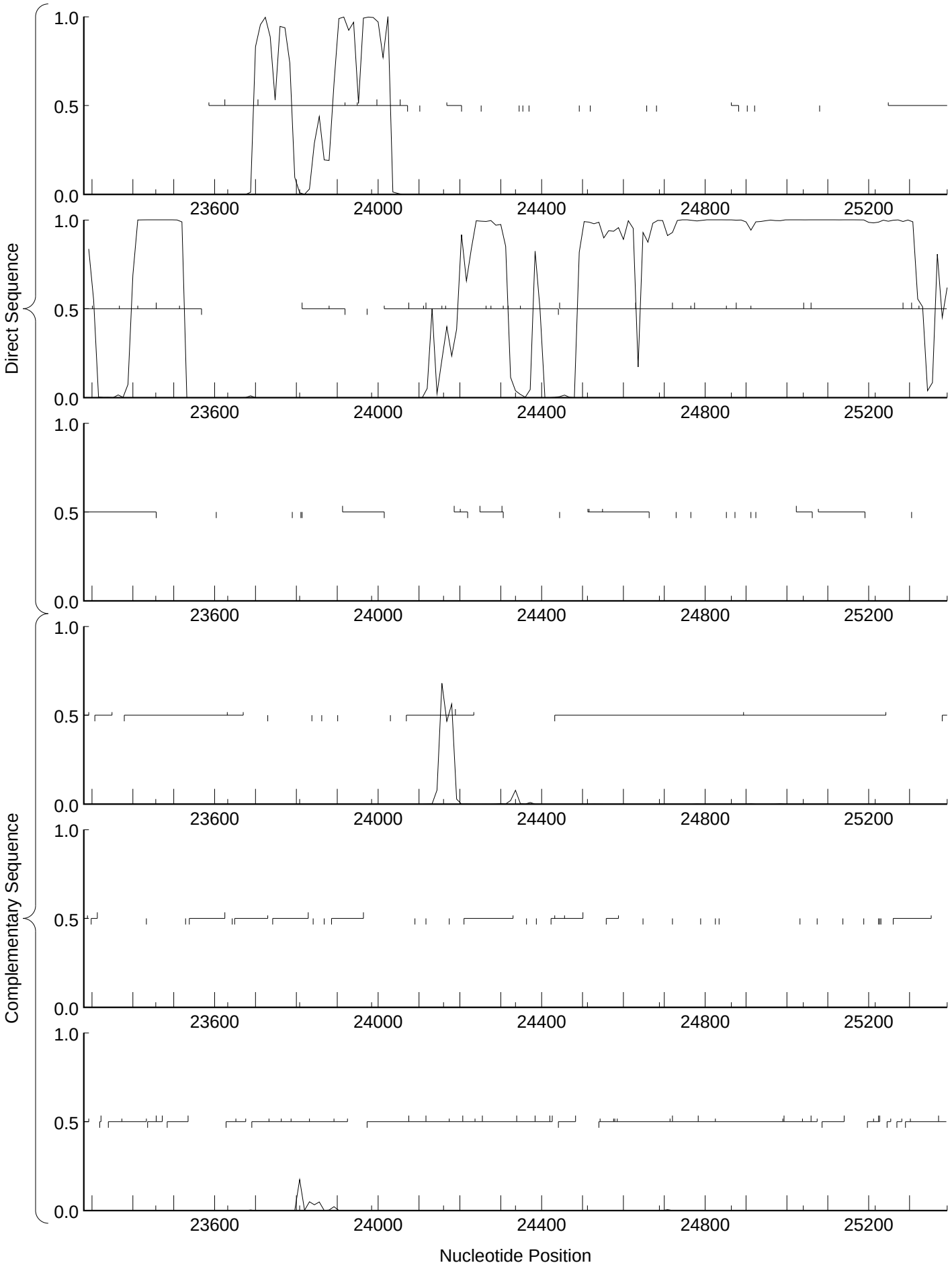
a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 11/27



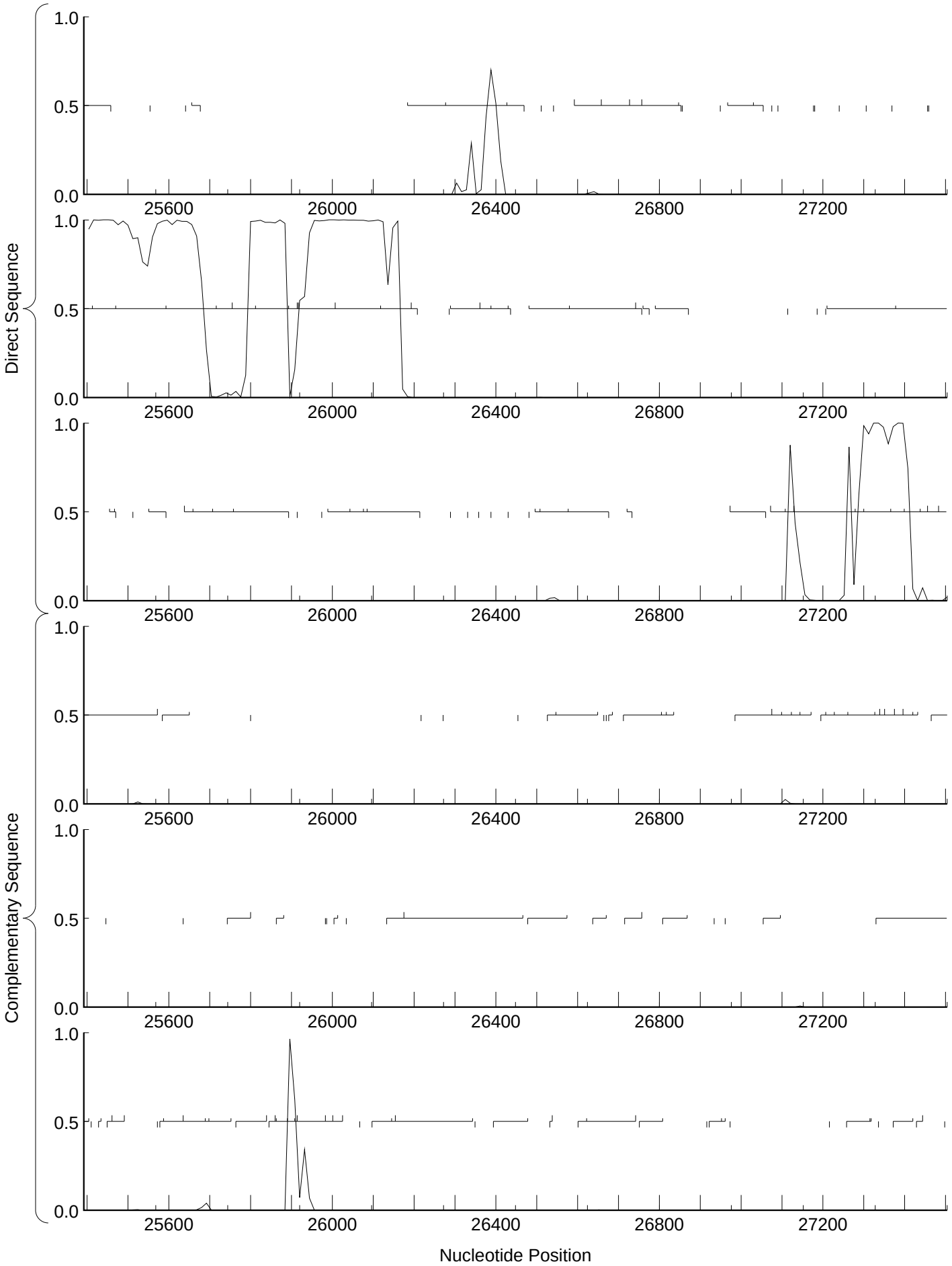
a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 12/27



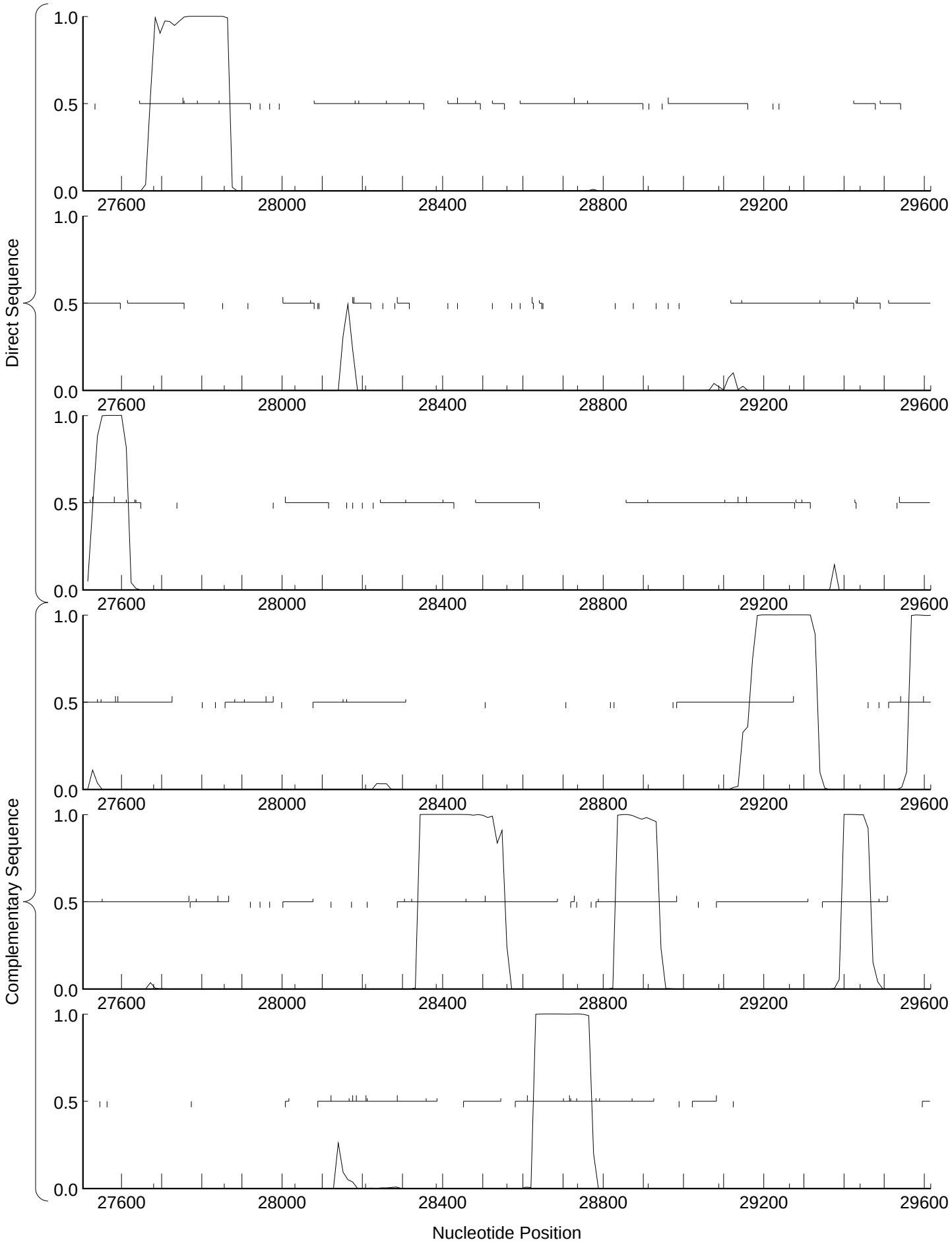
a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 13/27



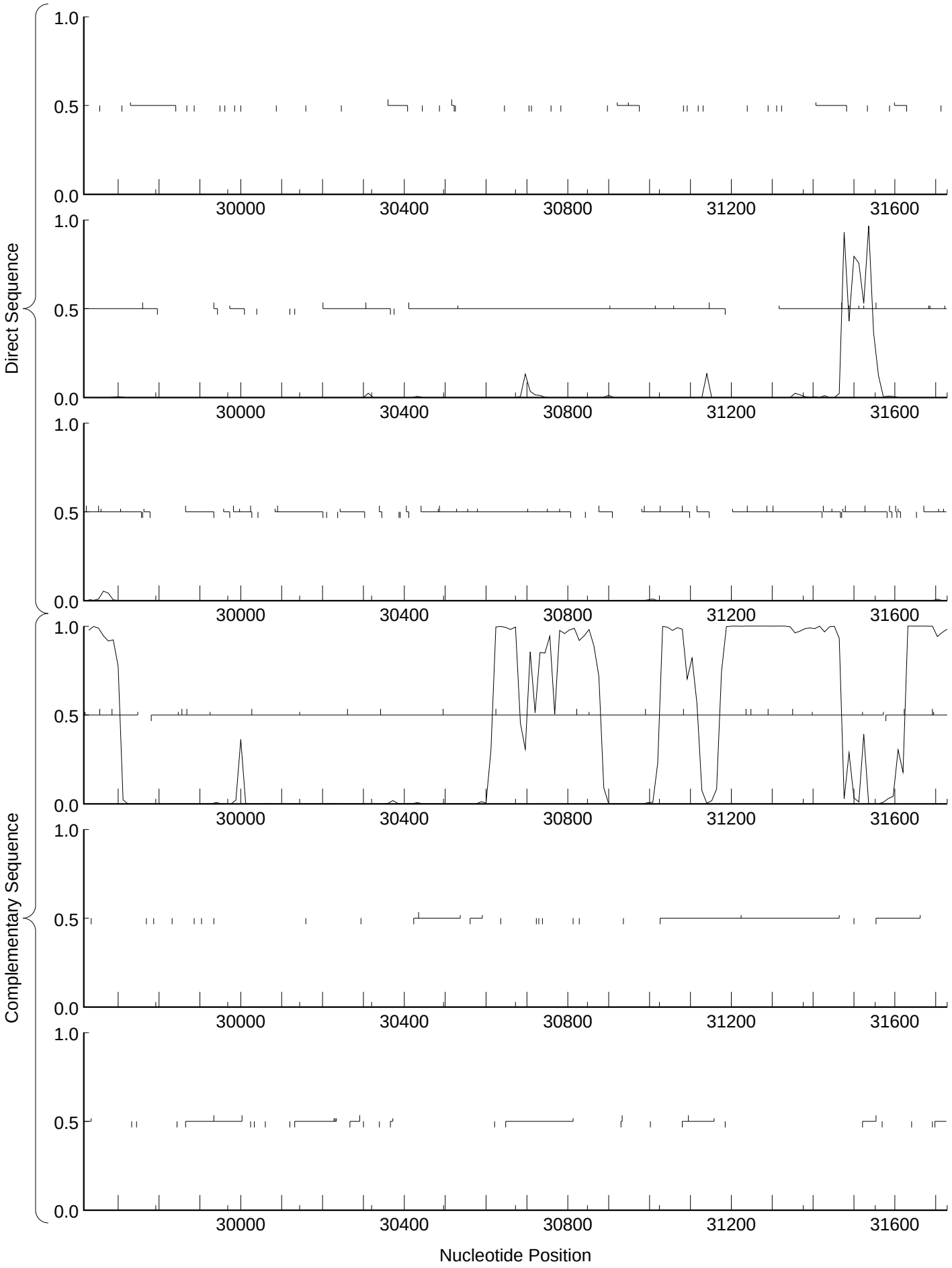
a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 14/27

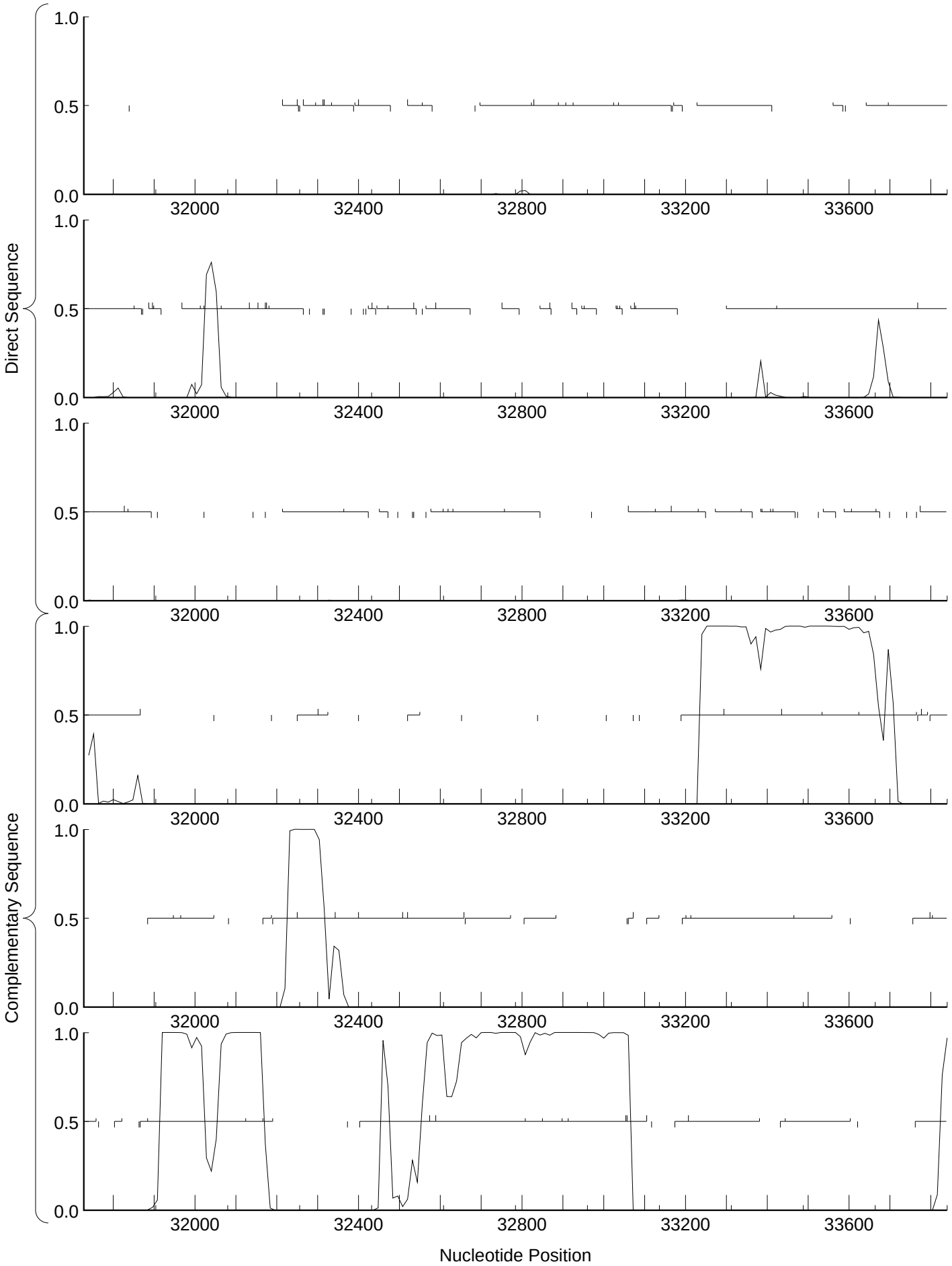


a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 15/27

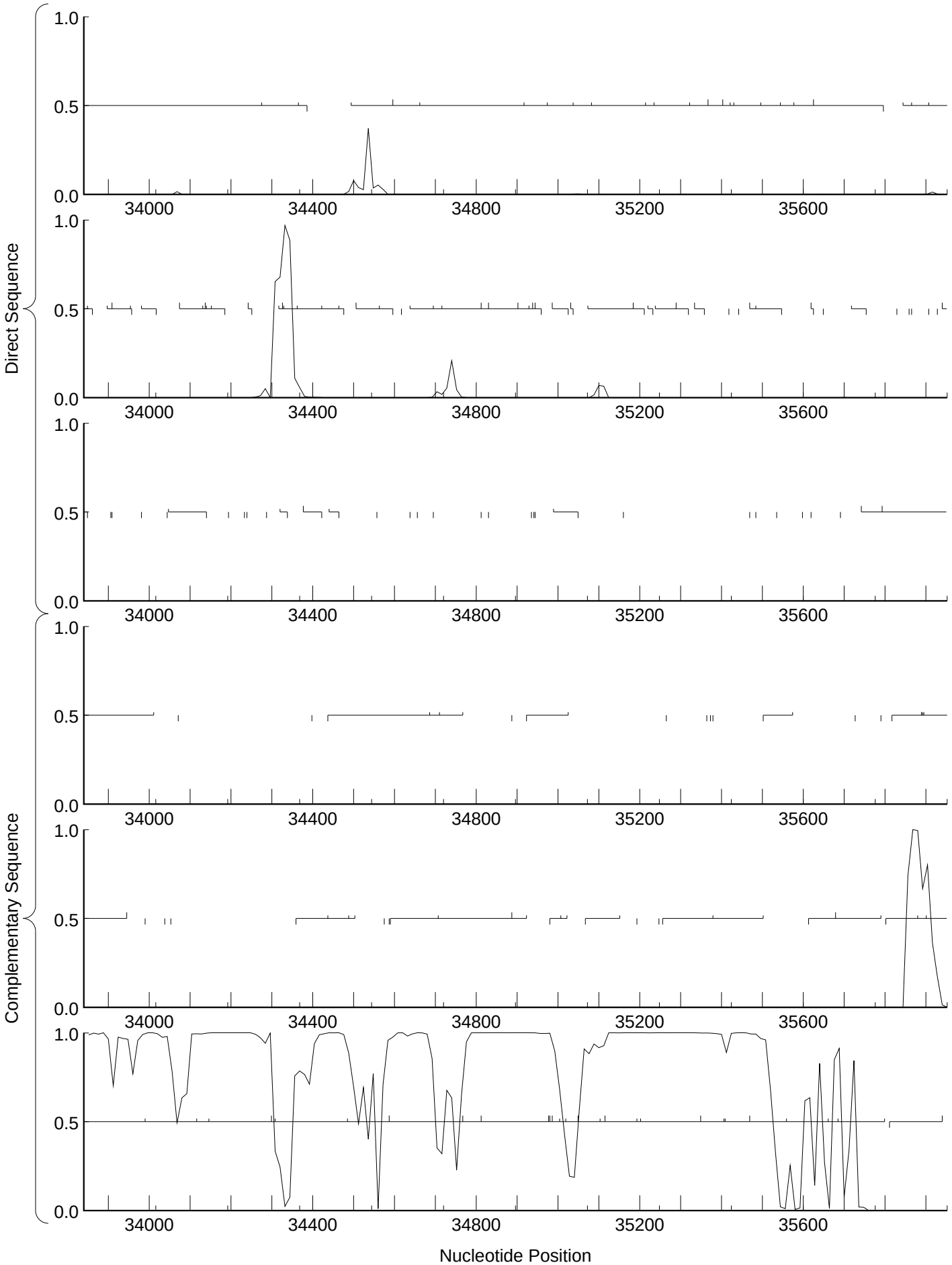


a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 16/27

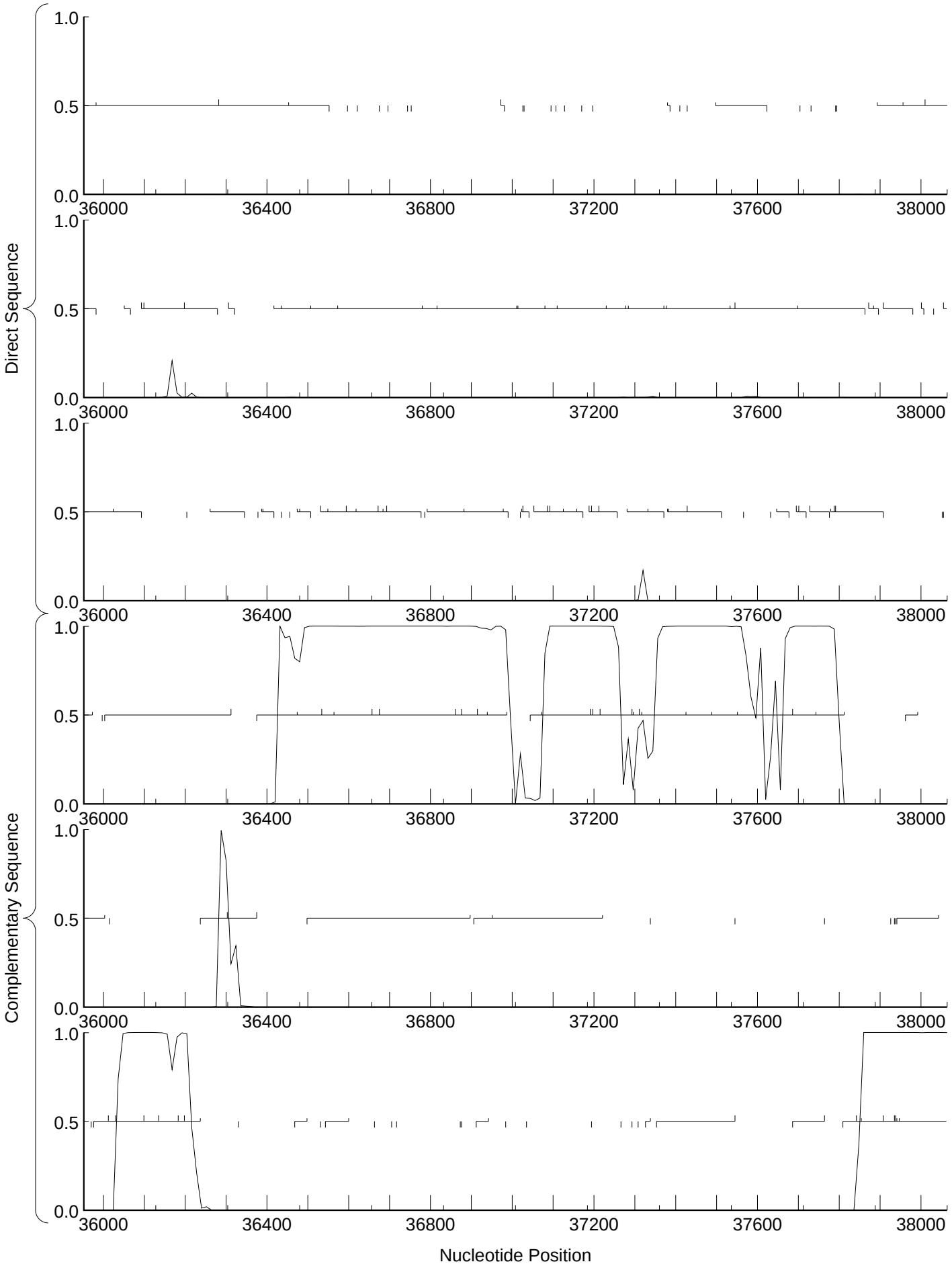



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
32000
32400
32800
33200
33600
Nucleotide Position

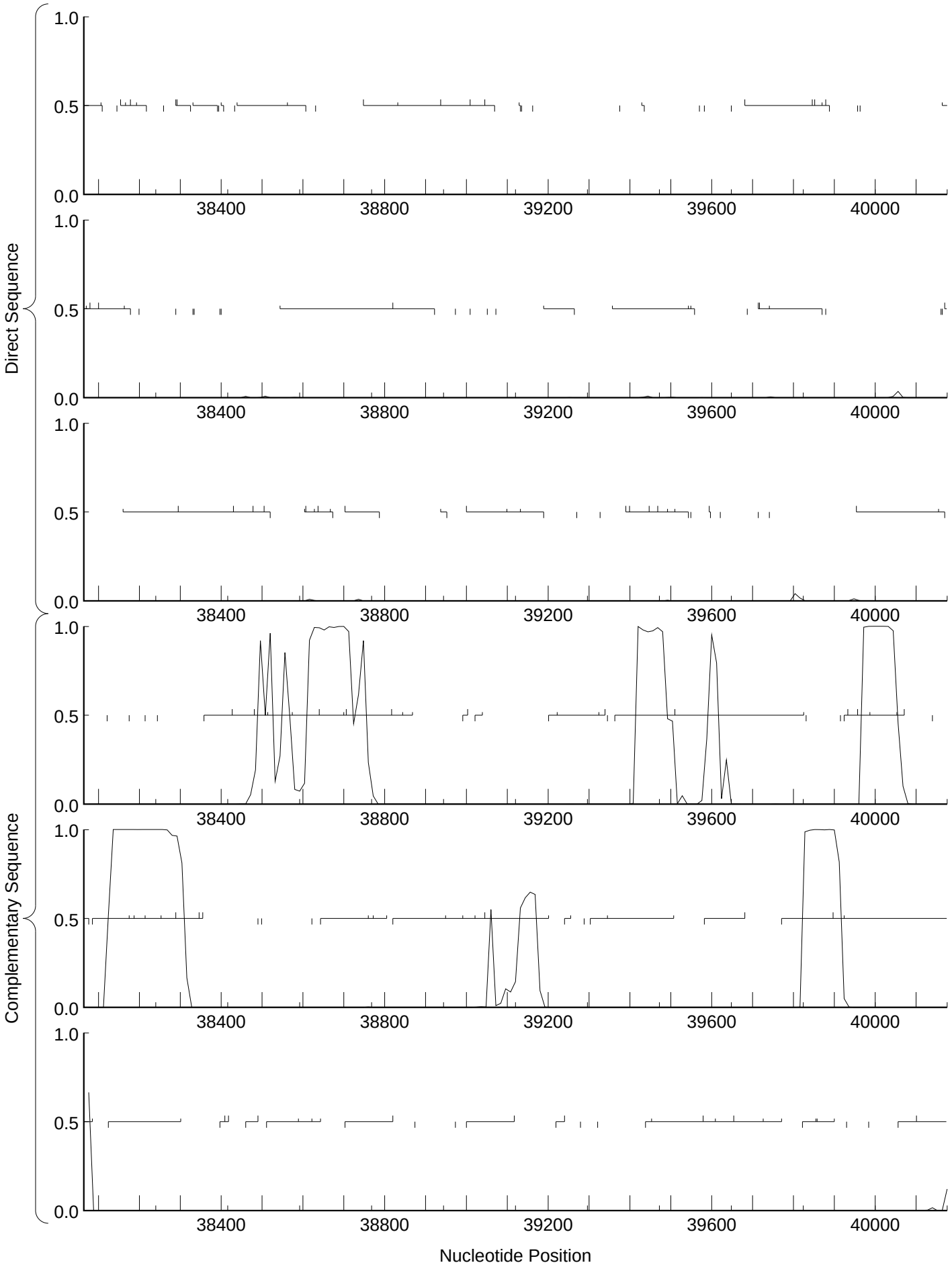
a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 18/27



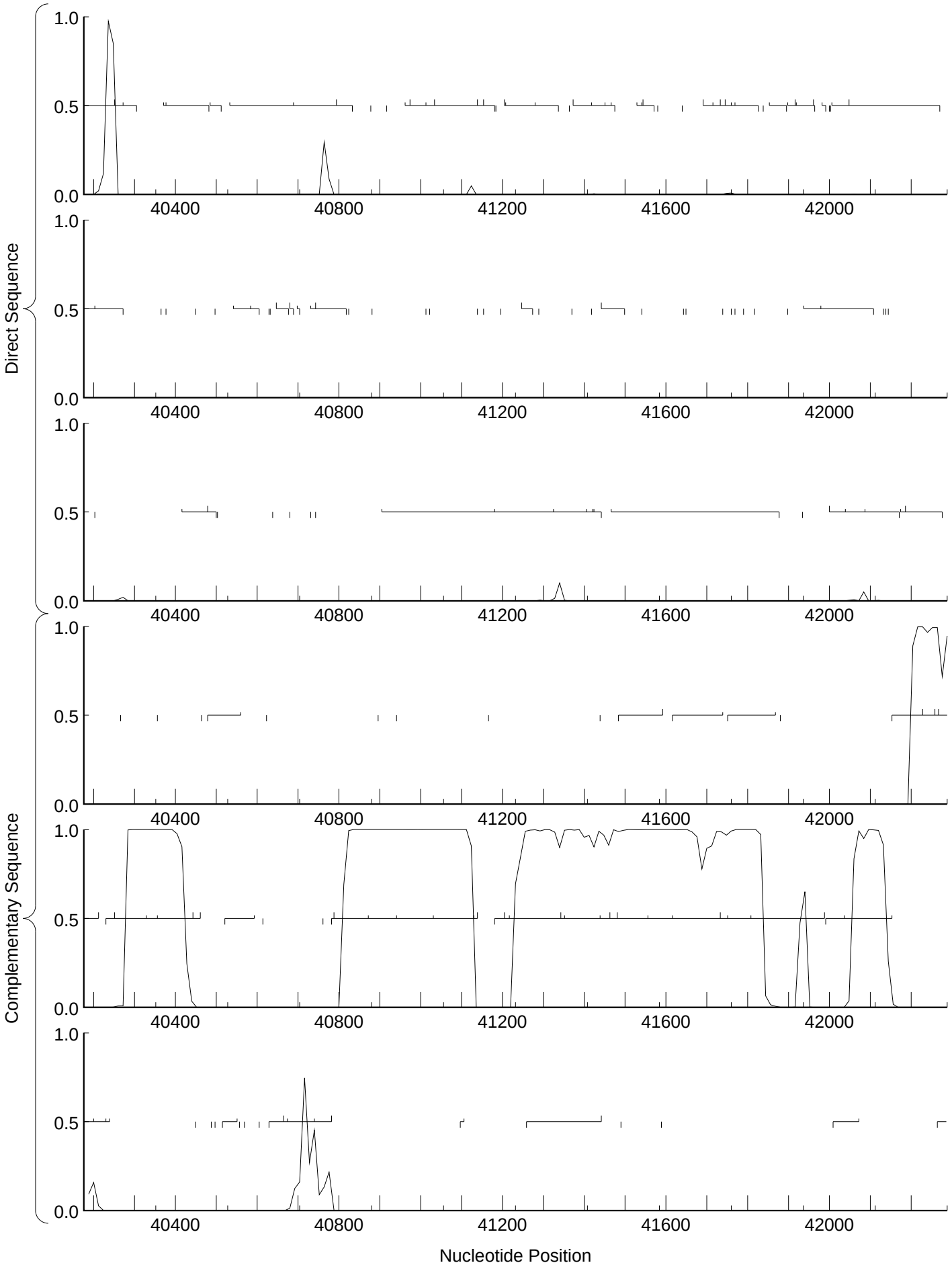
a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 19/27



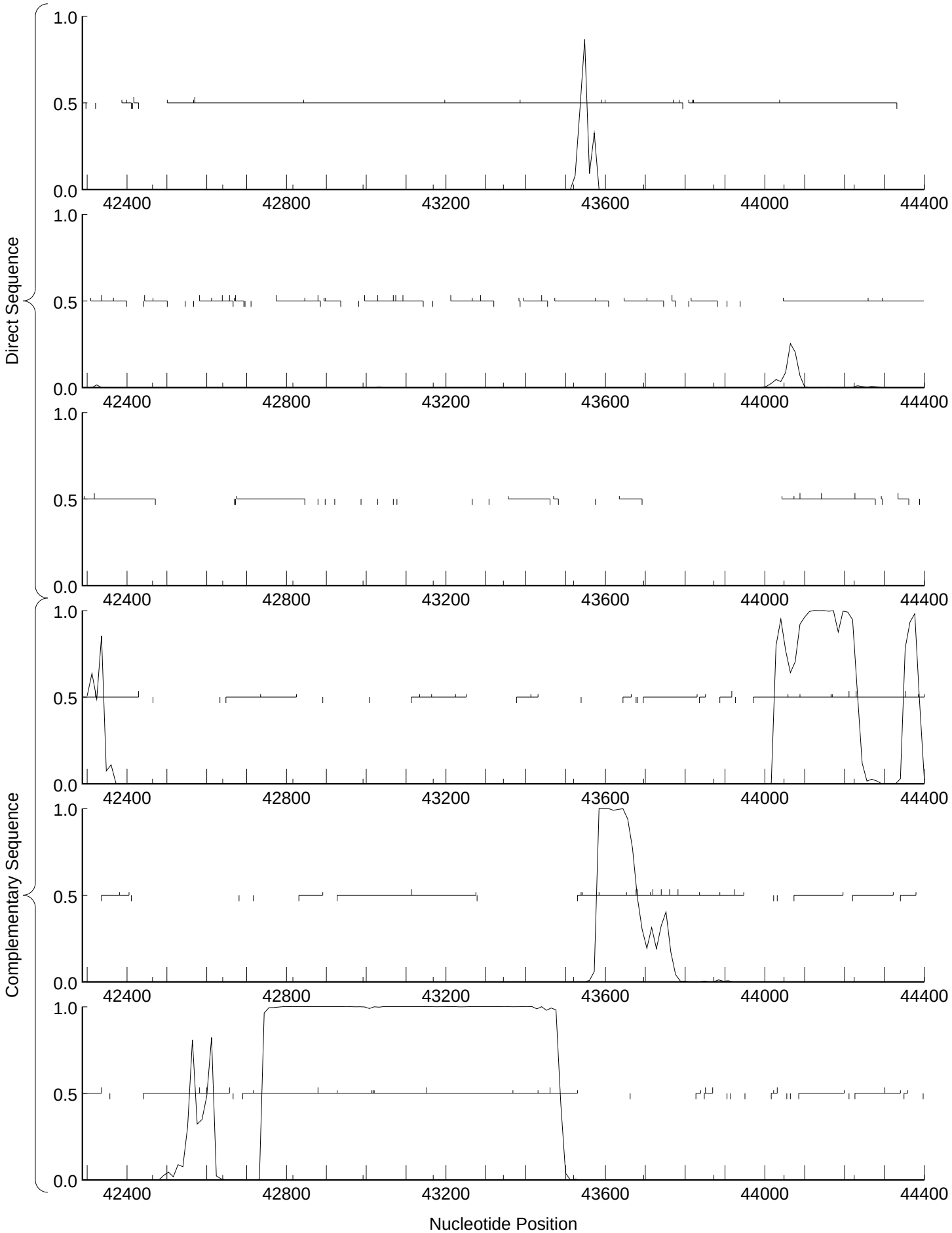
a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 20/27



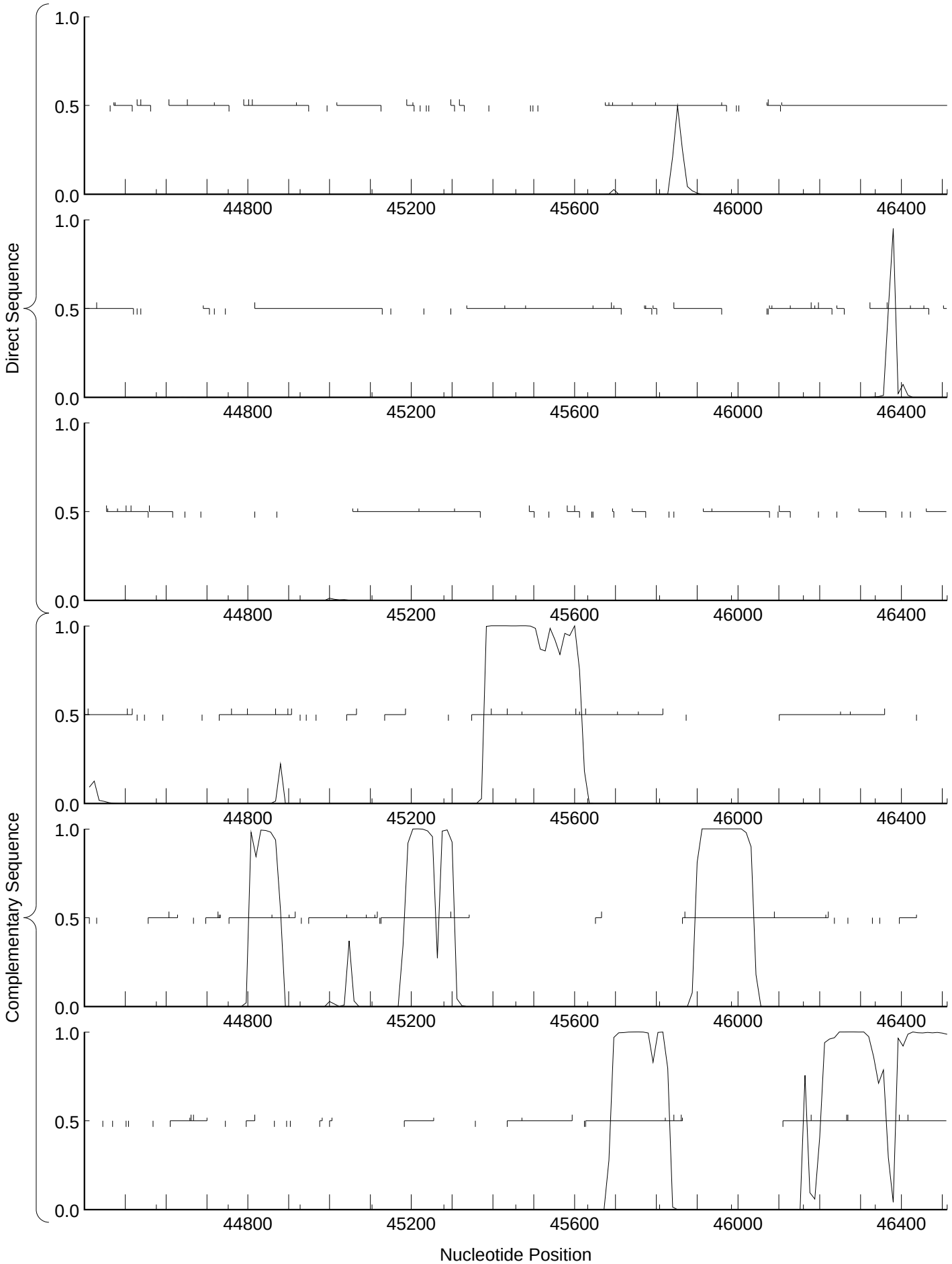
a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 21/27



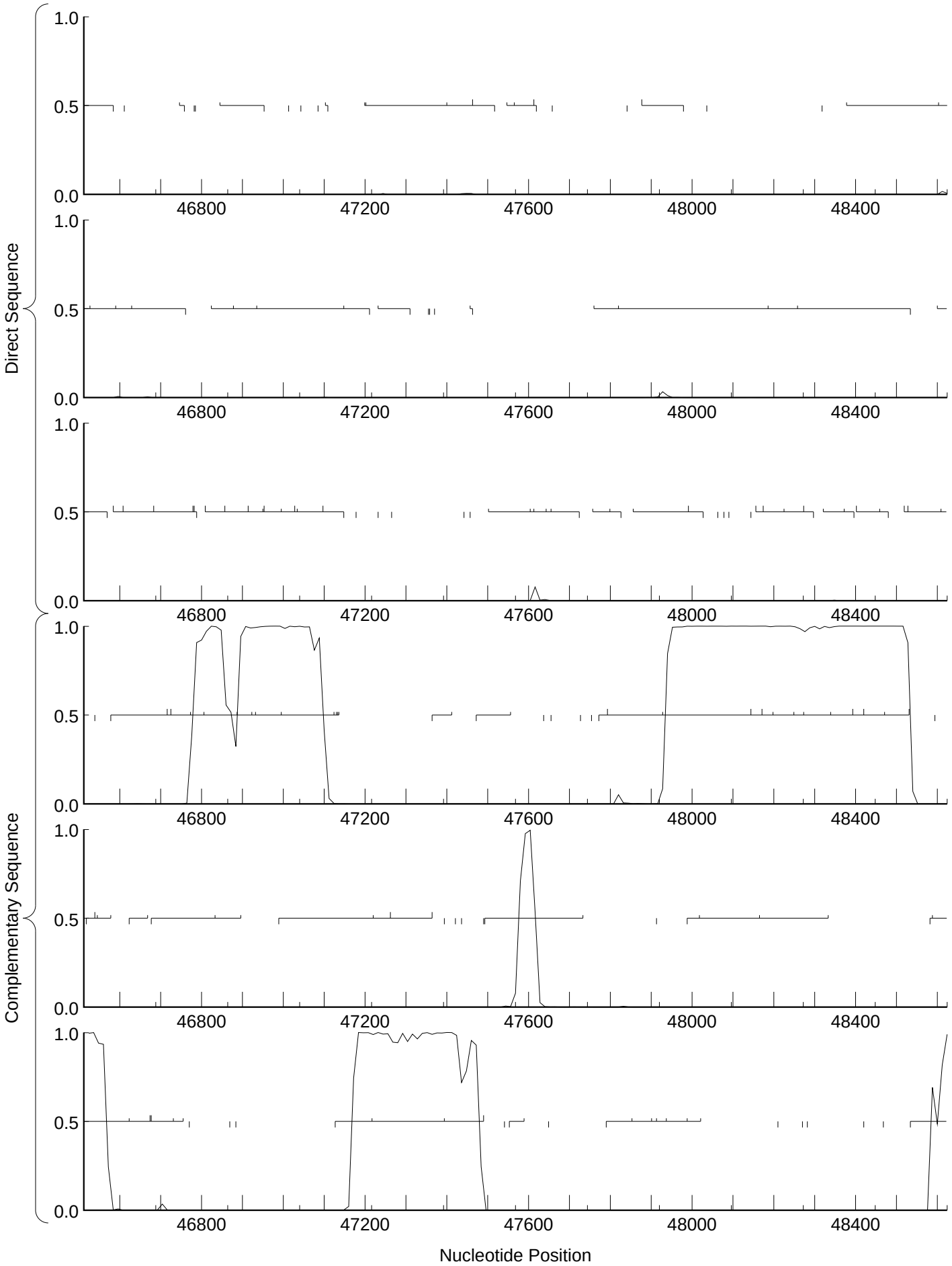
a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 22/27



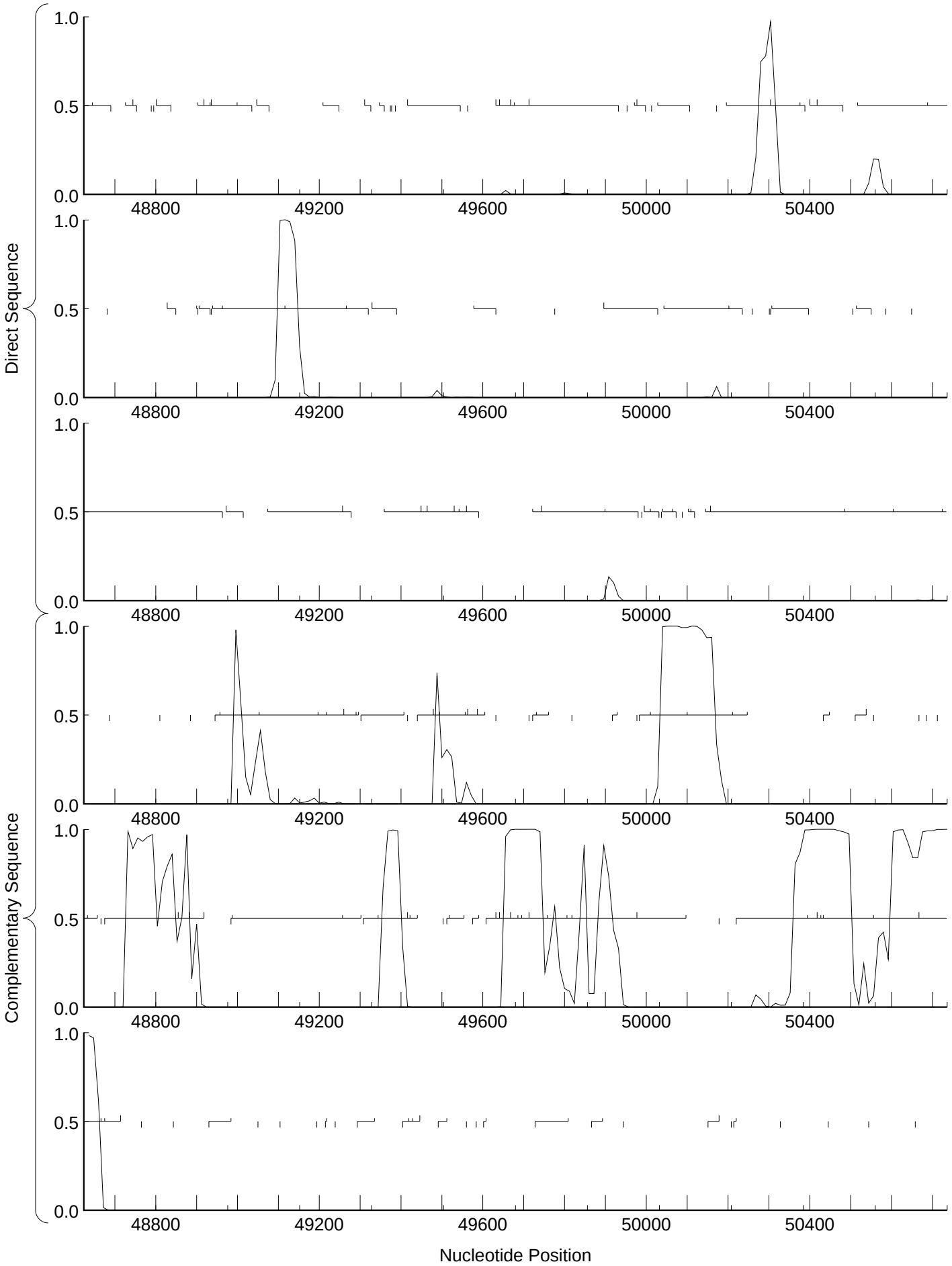
a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 23/27



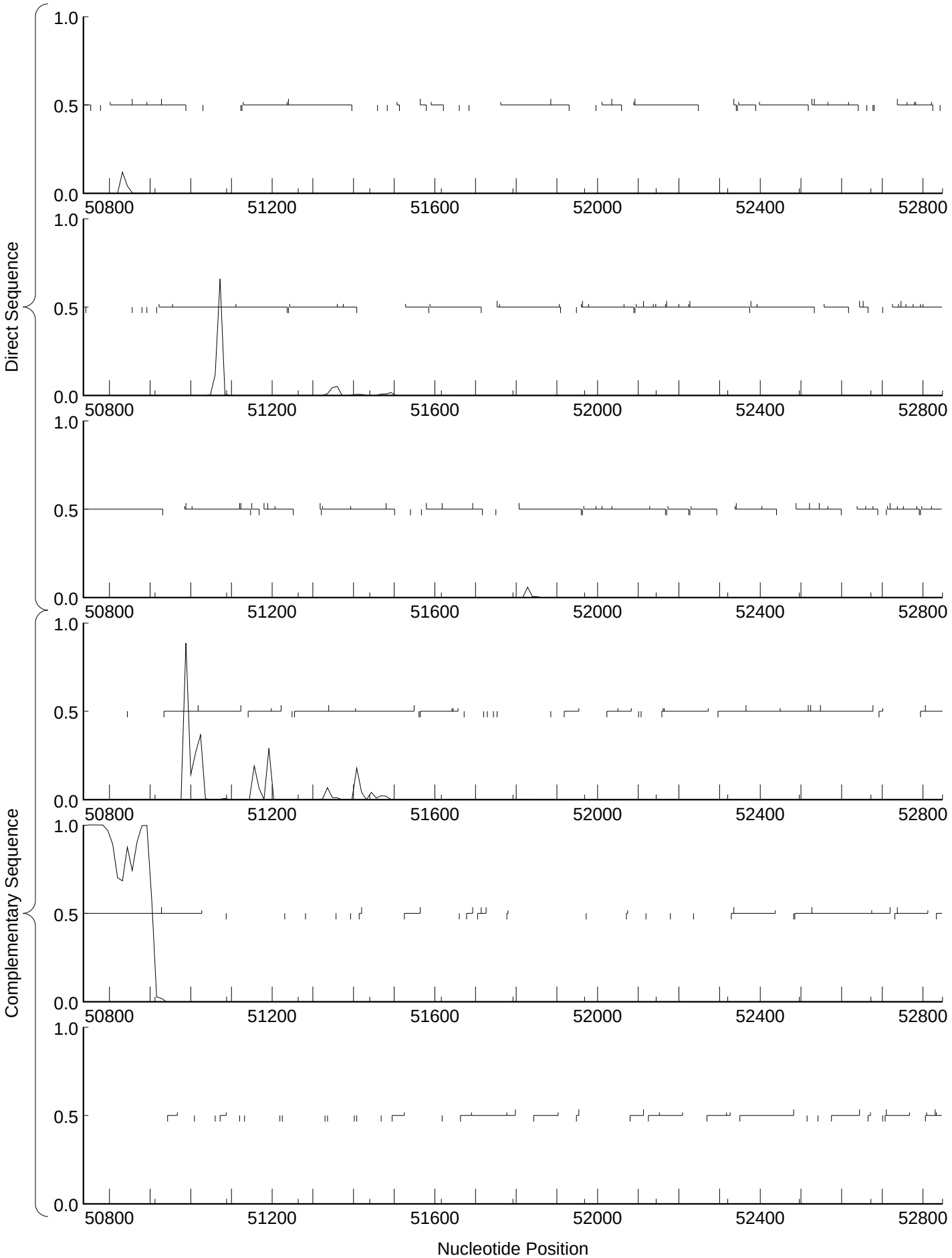
a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 24/27



a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 25/27



a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 26/27



a complete sequence, 53014 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 27/27

