GeneMark

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

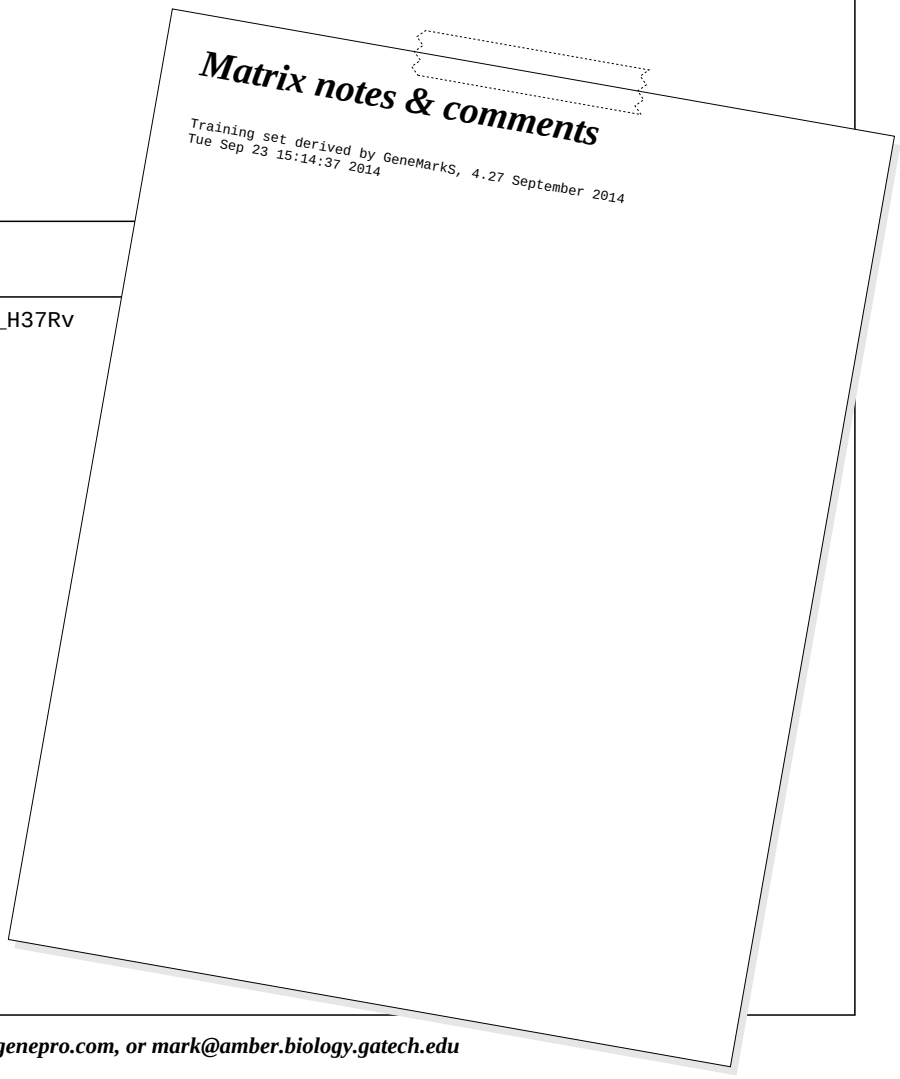
PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Newrala complete sequence, 52566 bp including 10-base 3’ overhang (CGGT
Analysis Date    :  3/29/24 at 4:20:46
Pages            :  26
Sequence Length  :  52566 bp
GC Content       :   61.58%

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

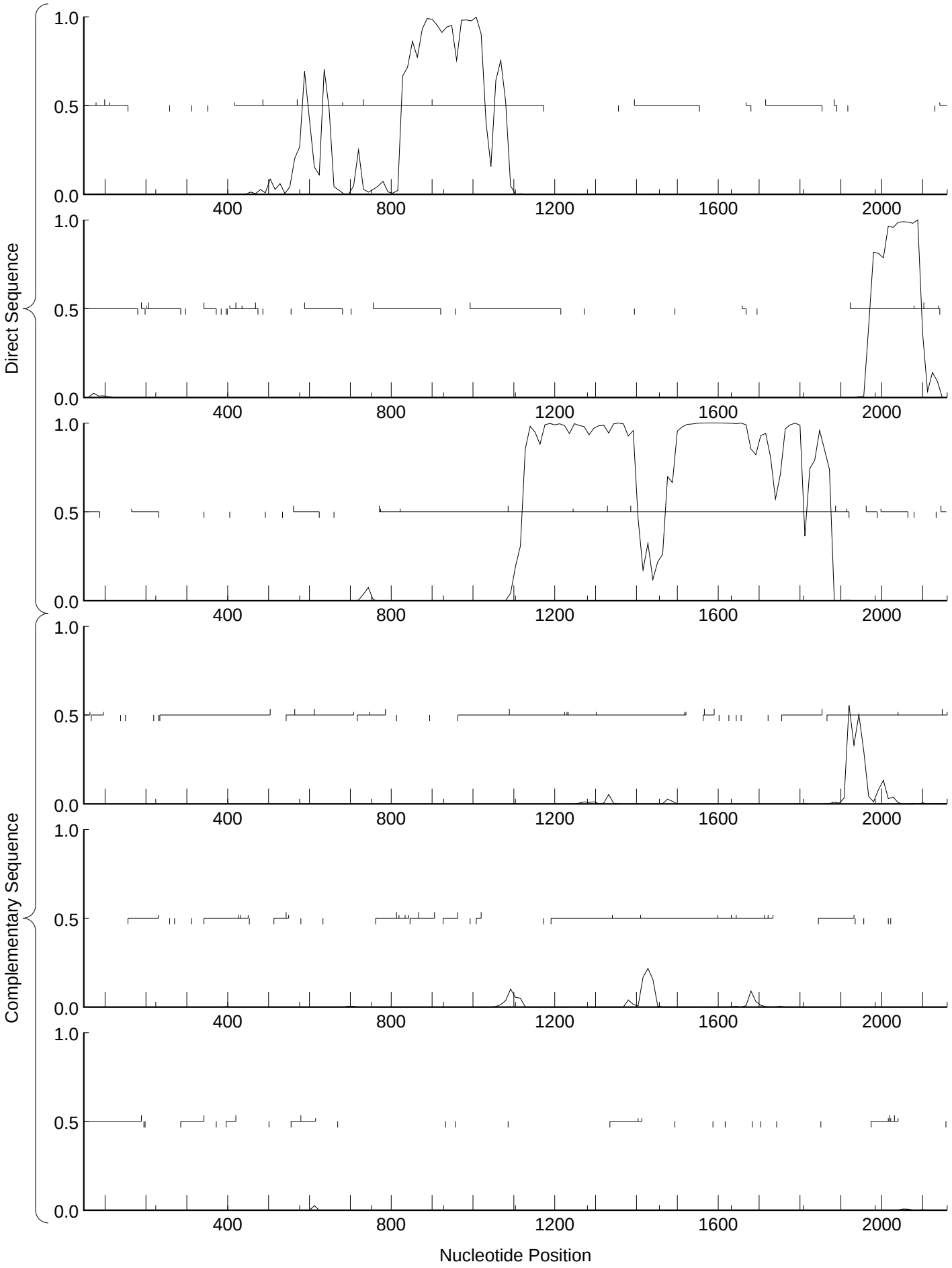
```
Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

Order      :  4

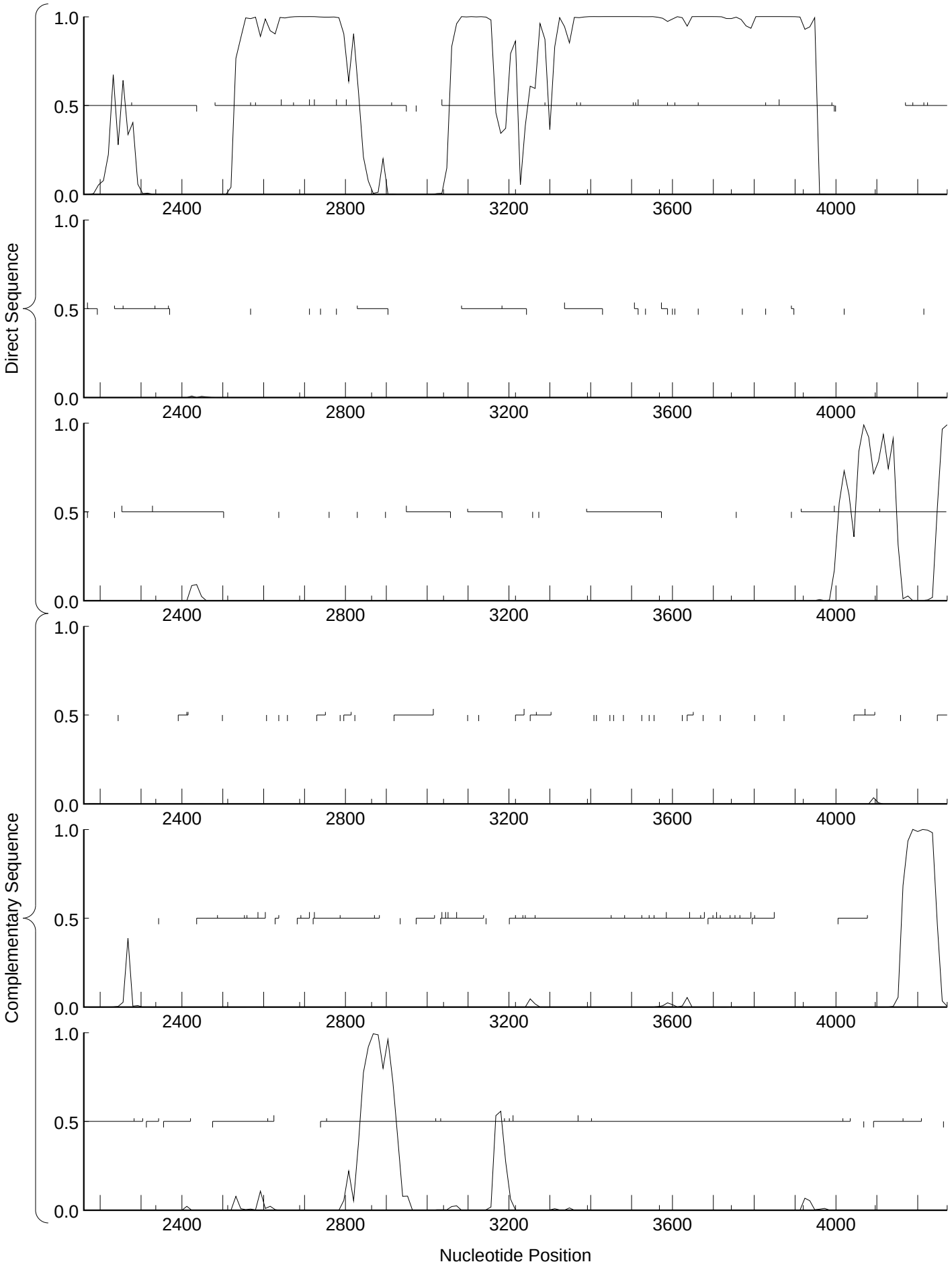

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

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

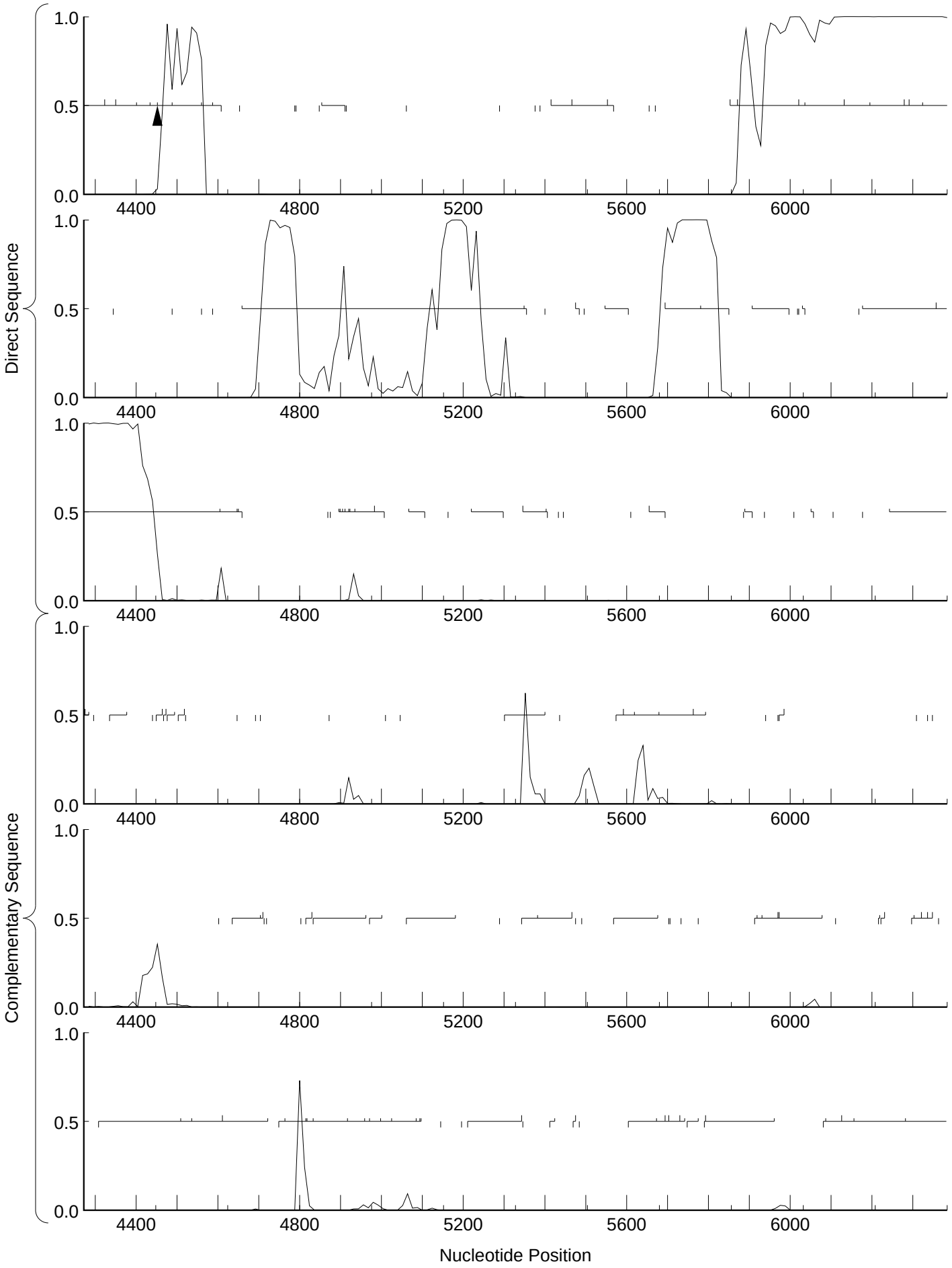
a complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 2/26



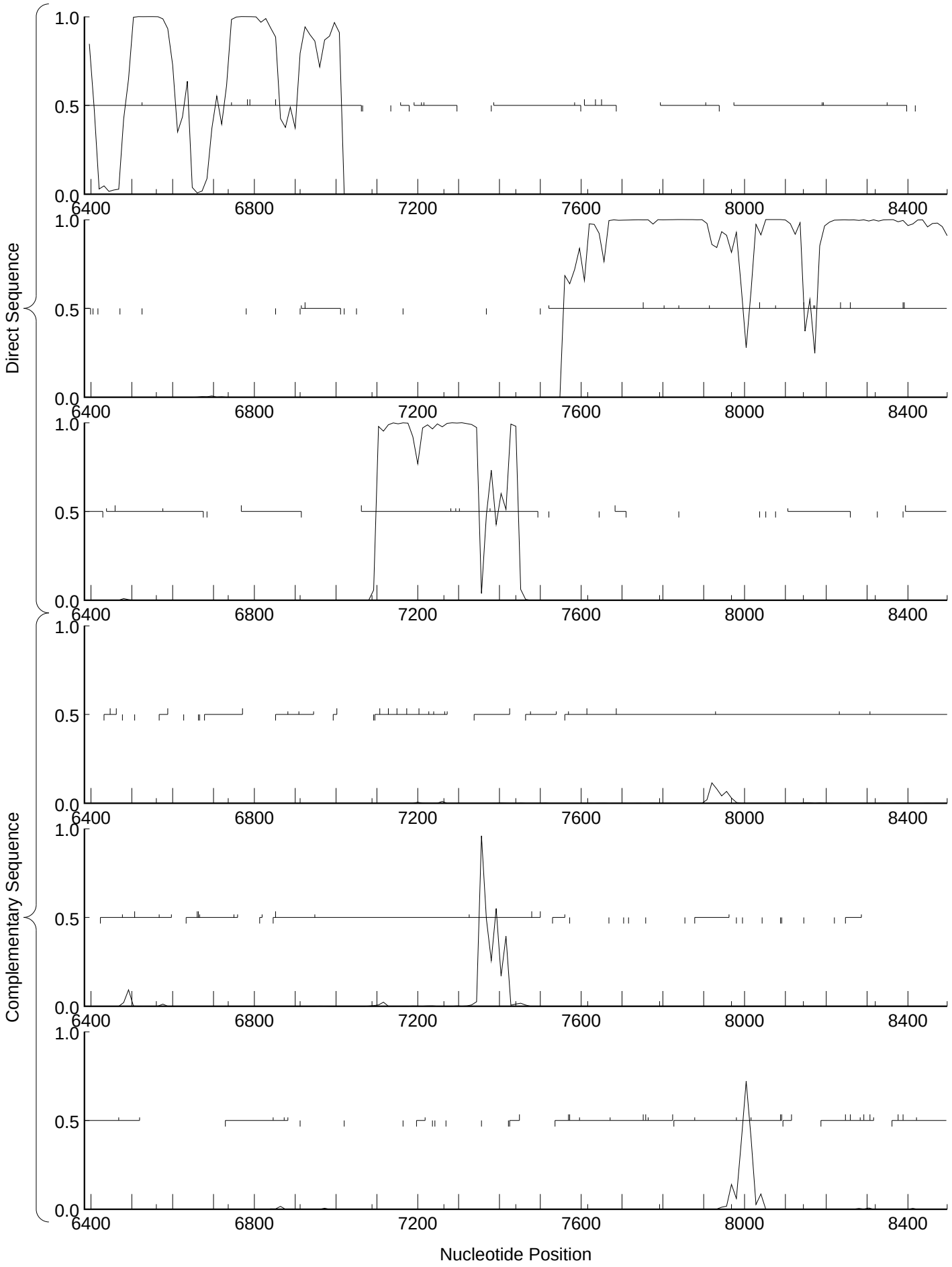
a complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 3/26



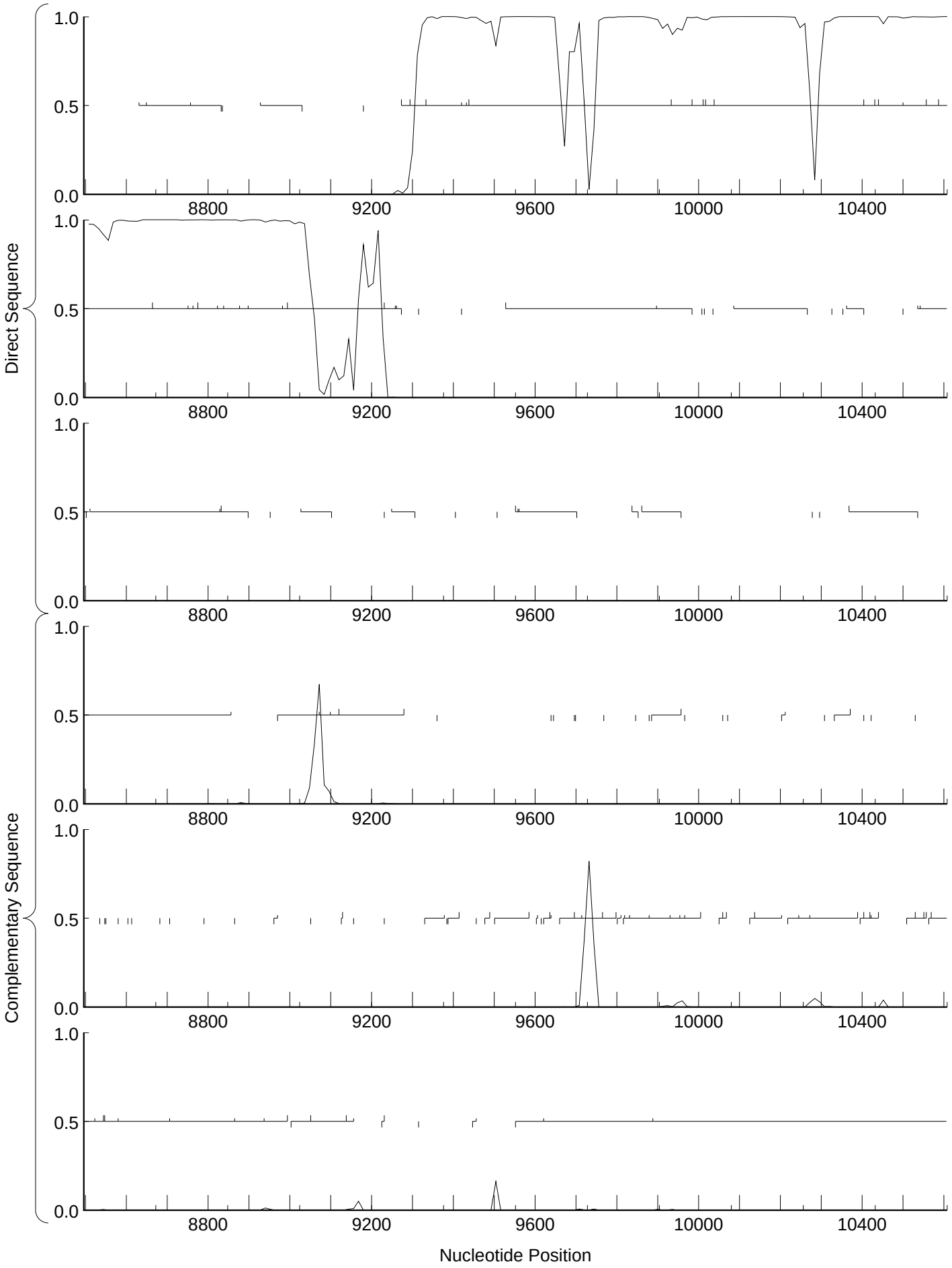
a complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 4/26



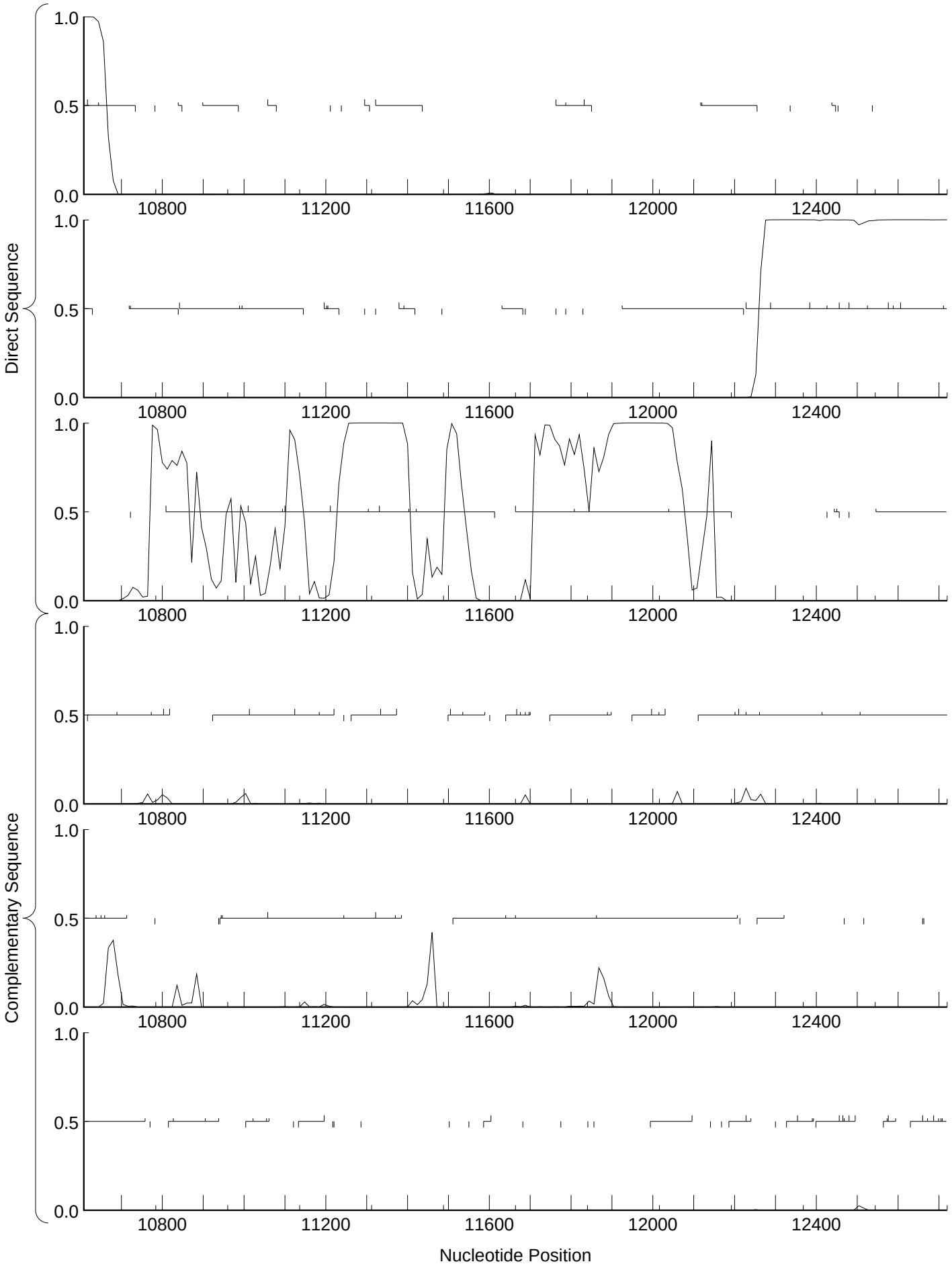
a complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 5/26



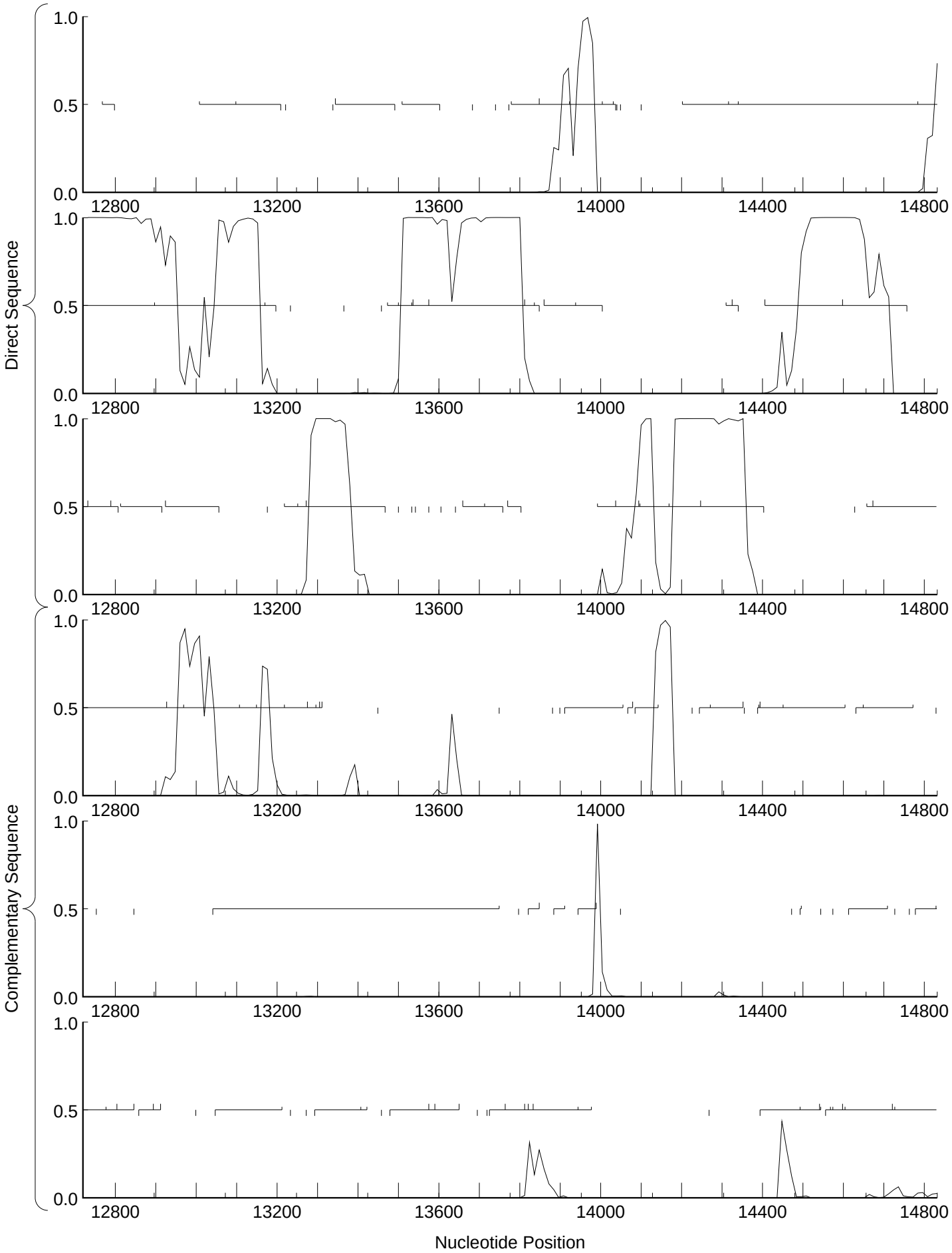
a complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 6/26



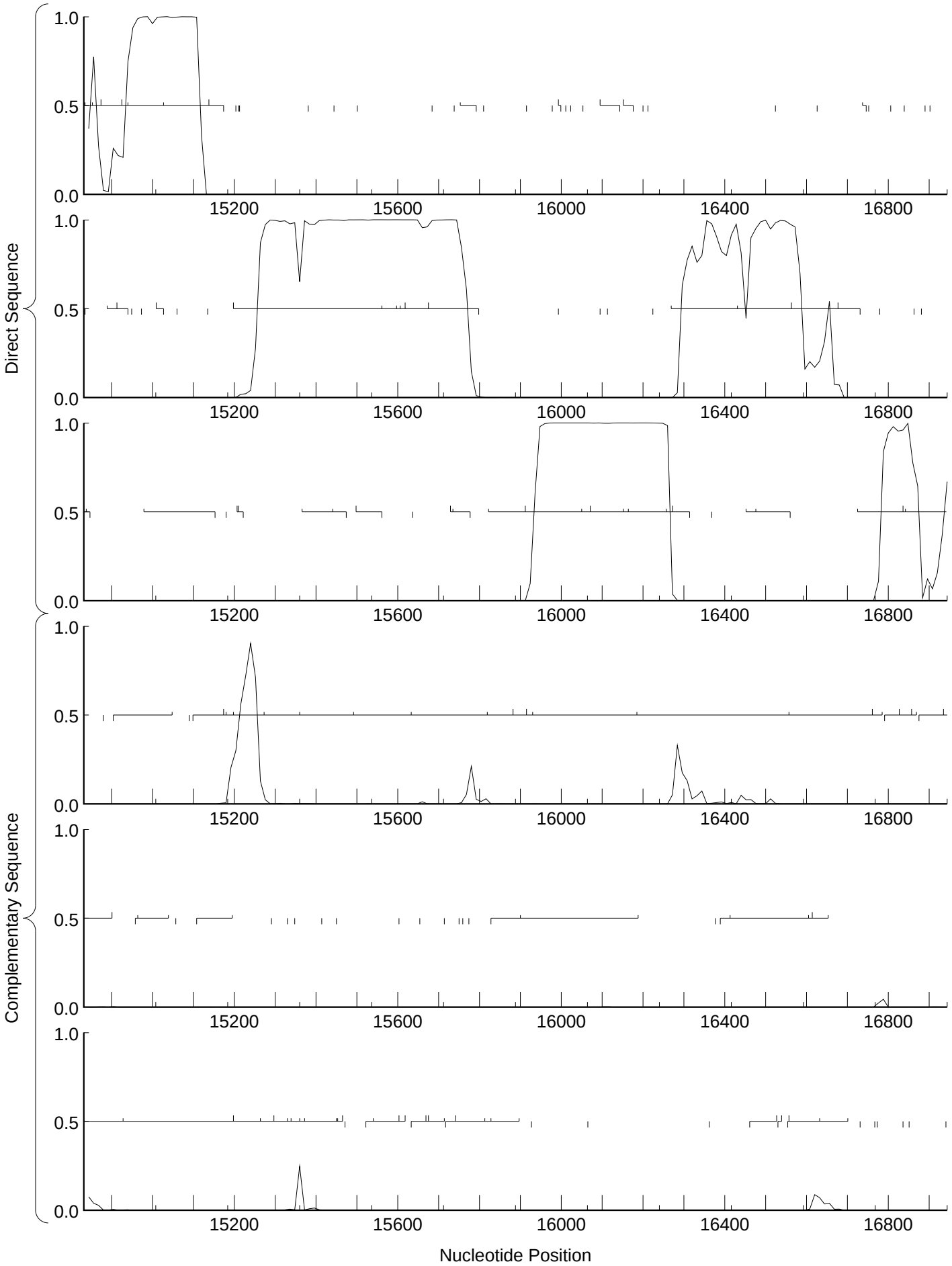
a complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 7/26



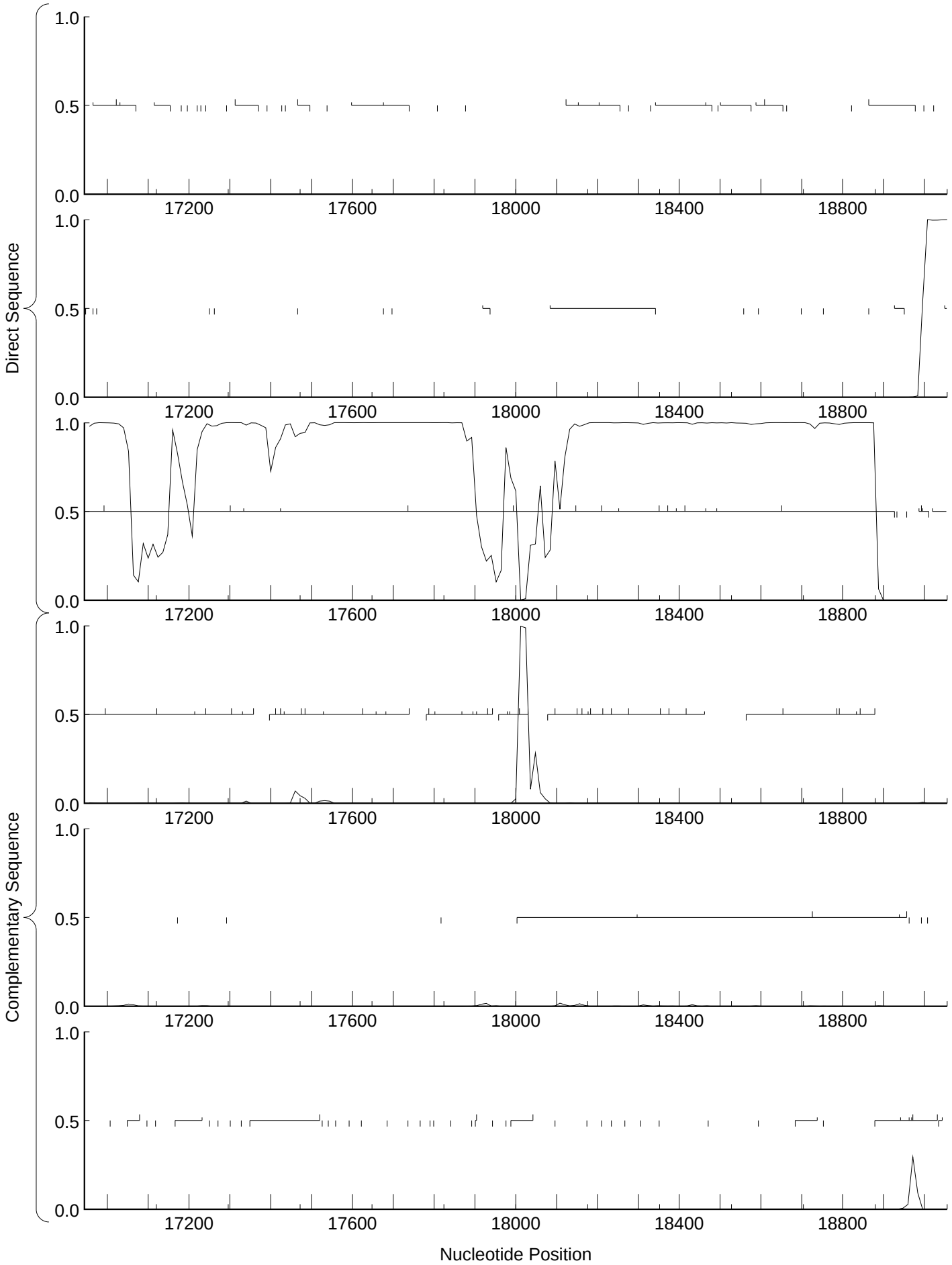
a complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 8/26



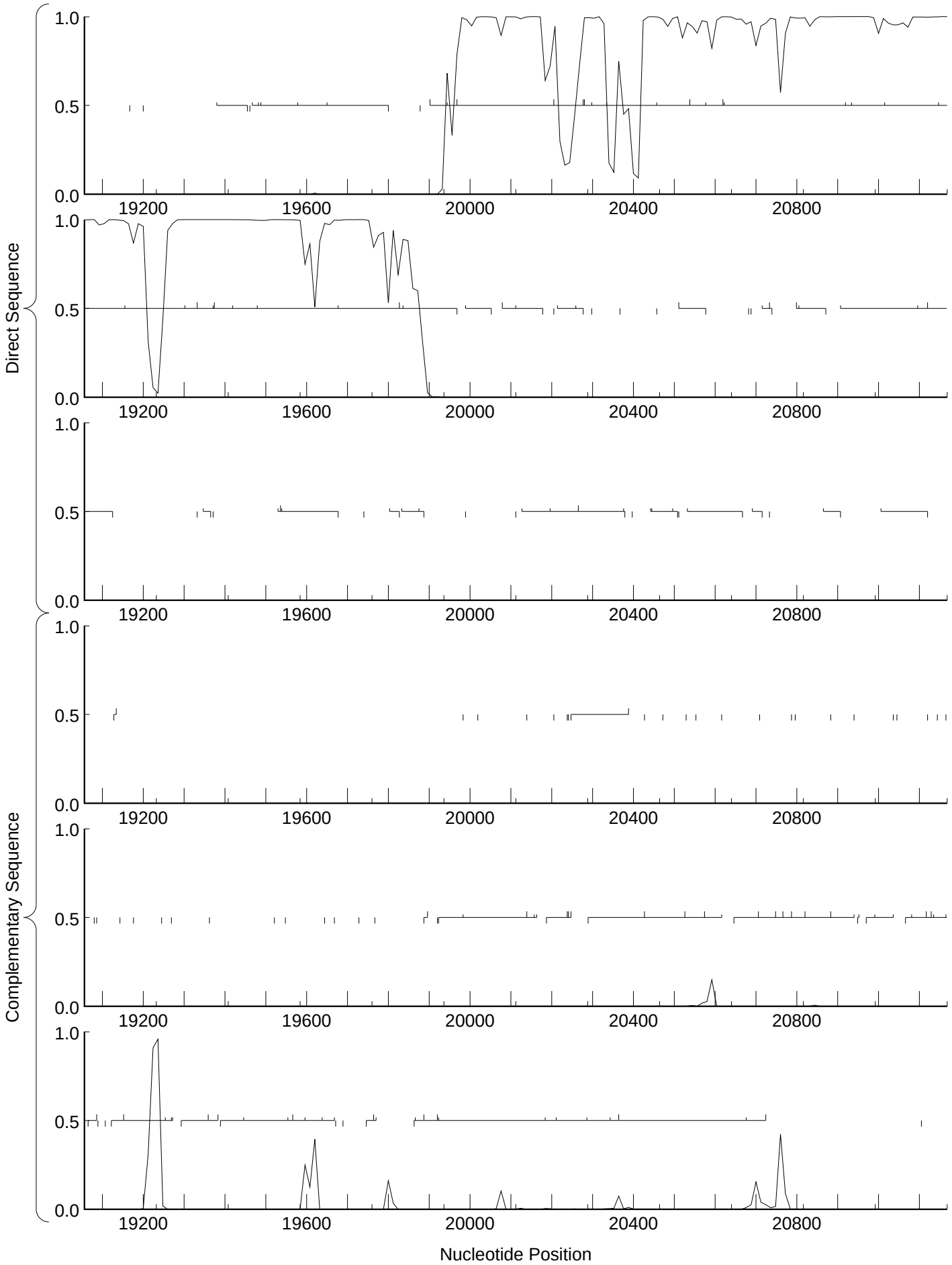
a complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 9/26



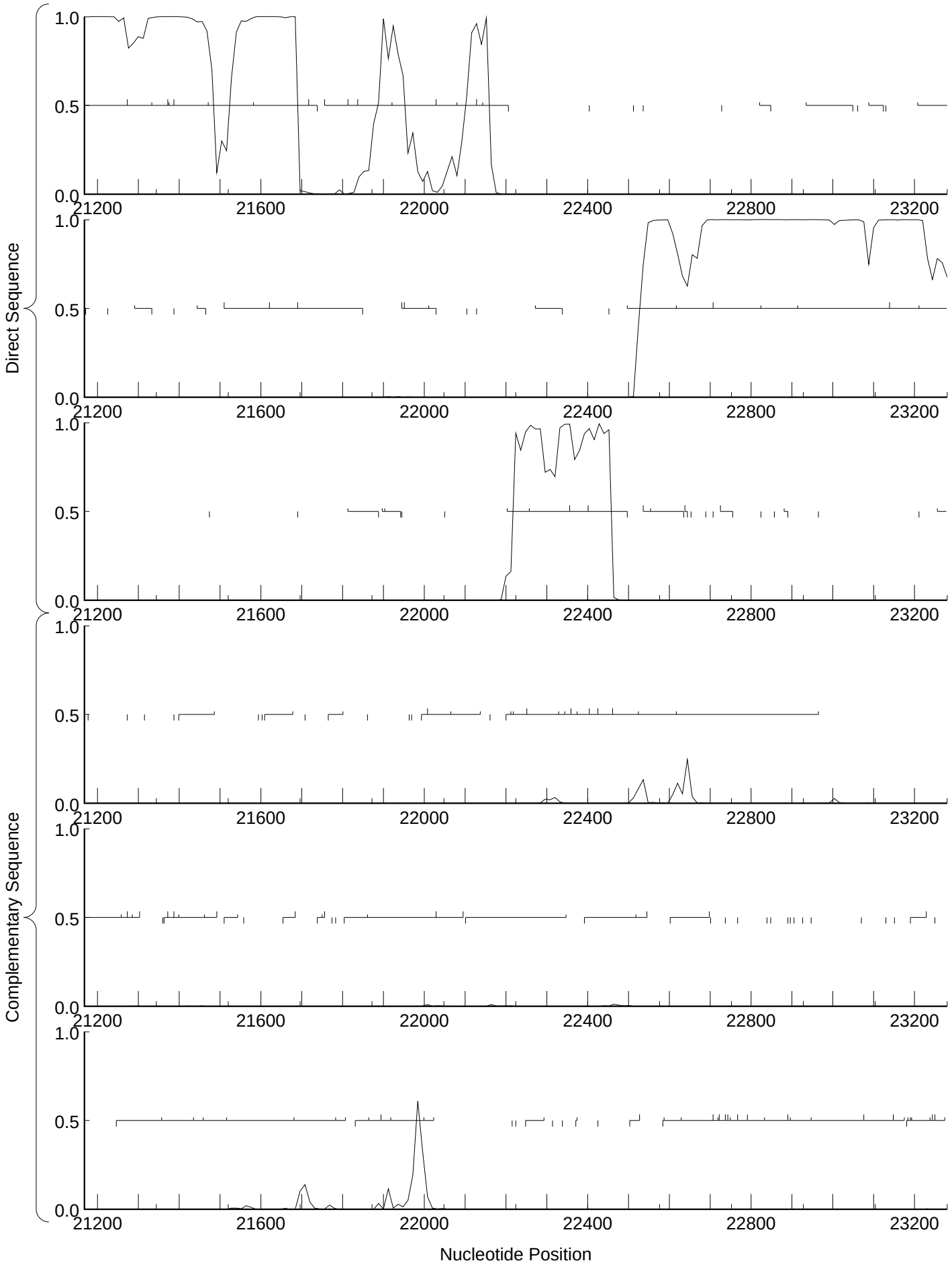
complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 10/26



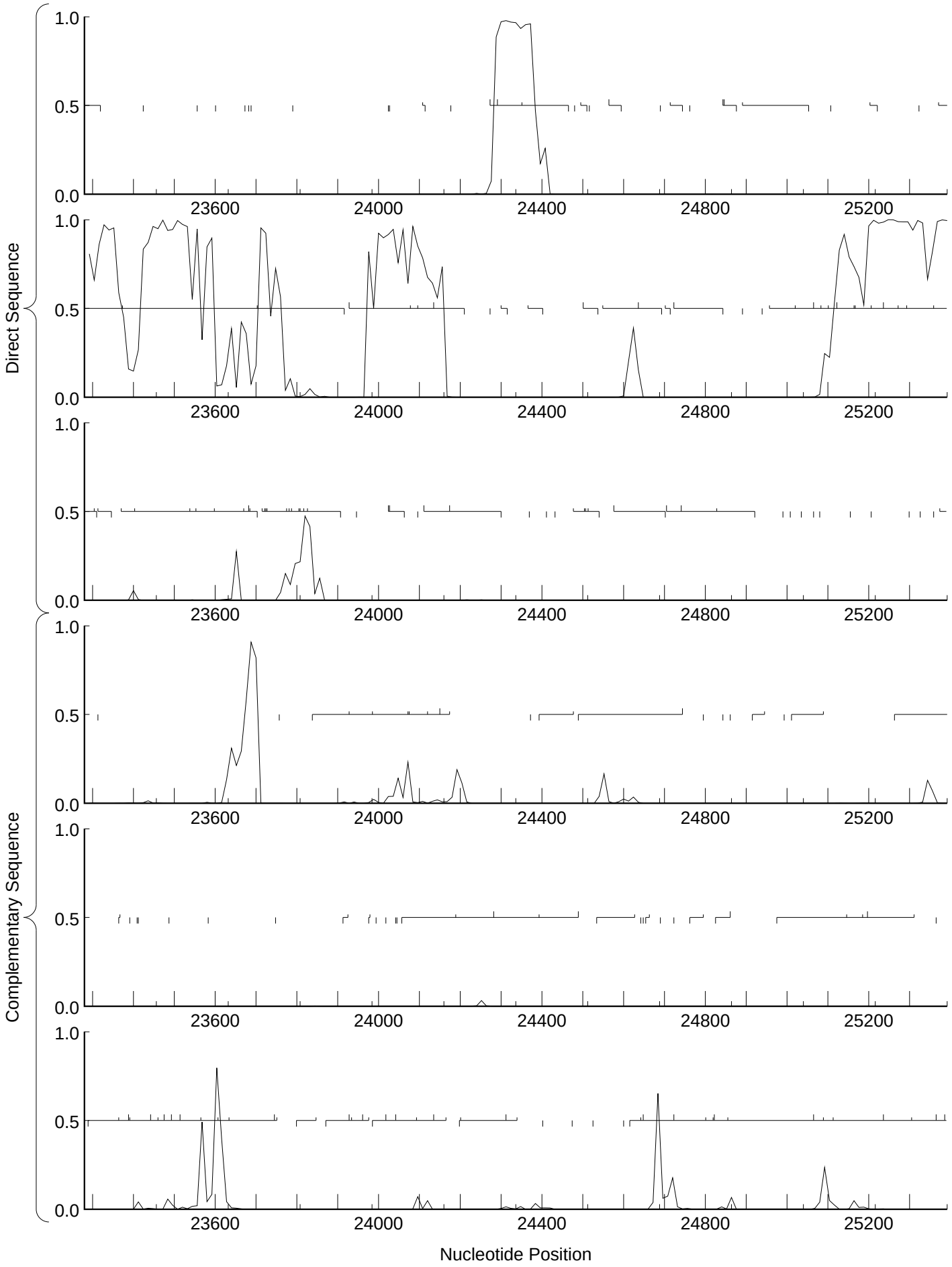
complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 11/26



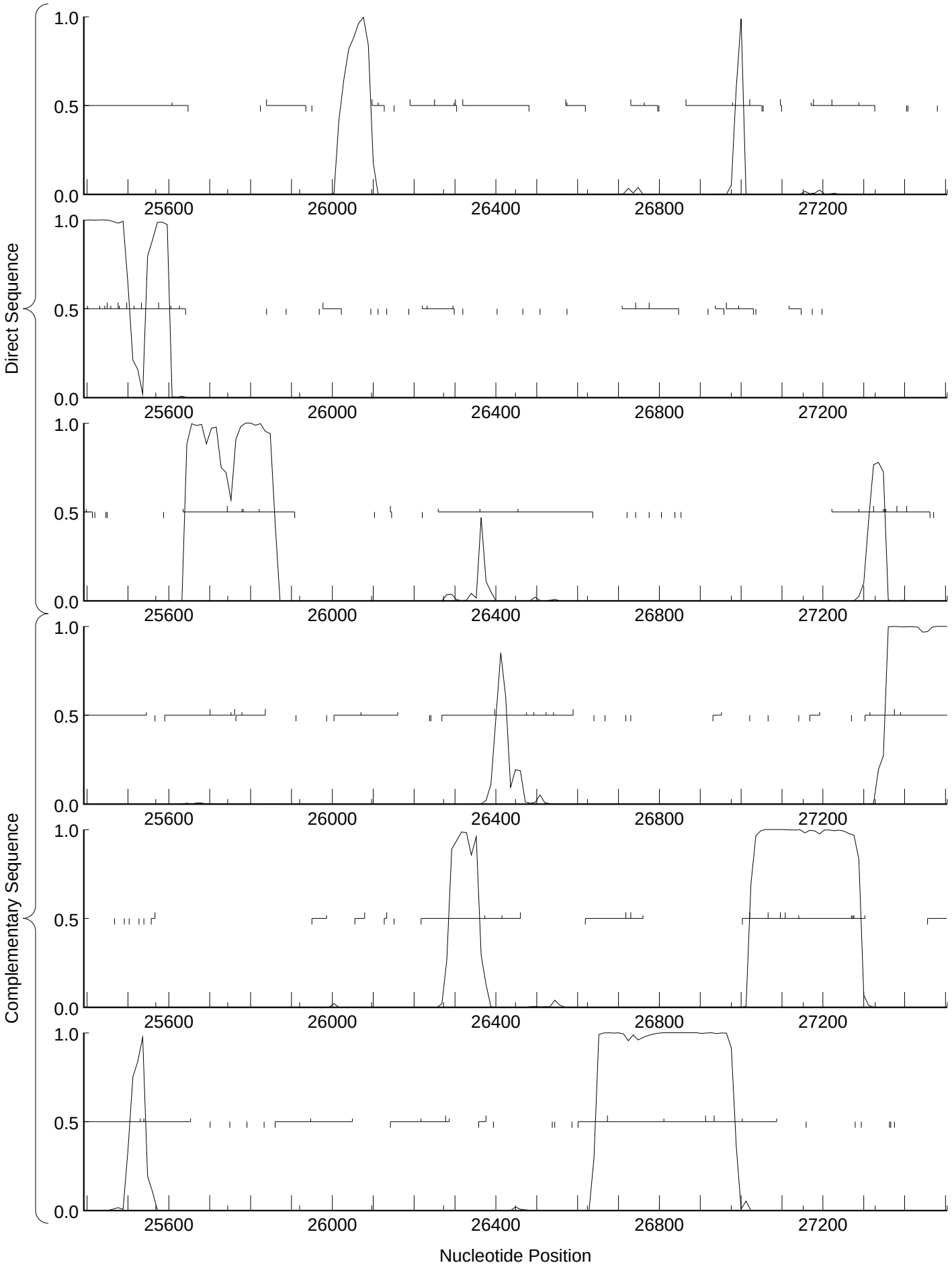
complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 12/26



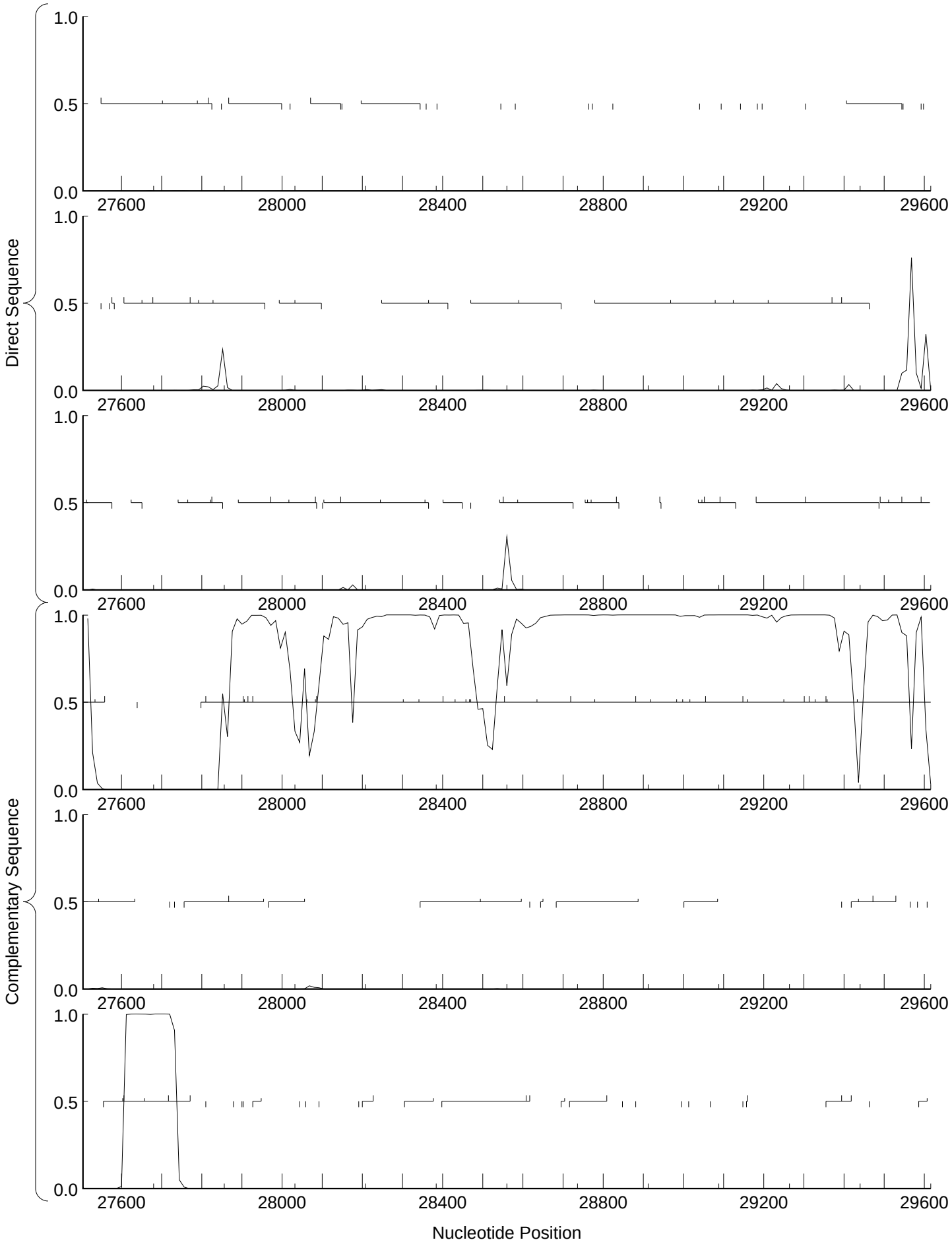
complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 13/26



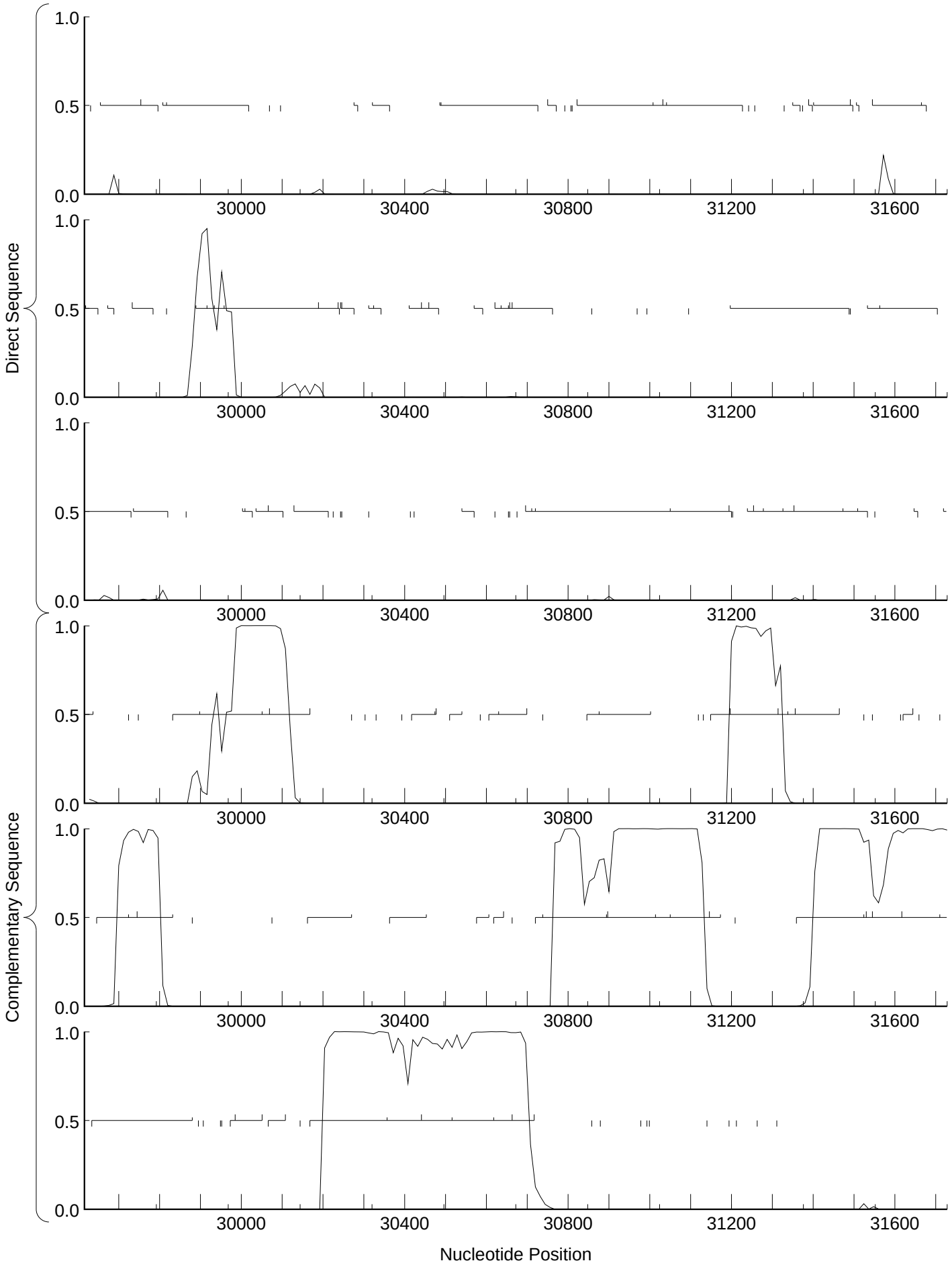
complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 14/26



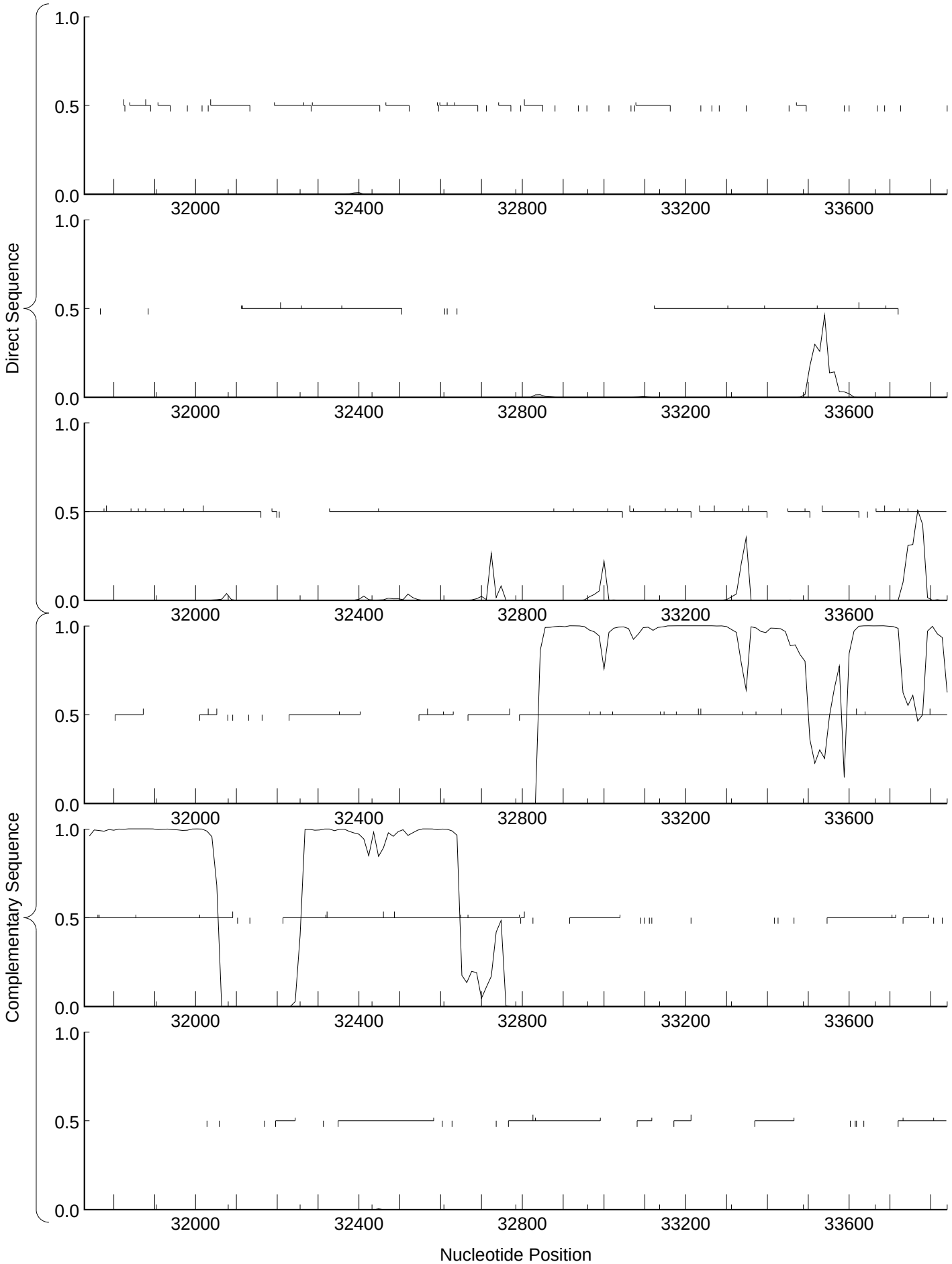
complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 15/26

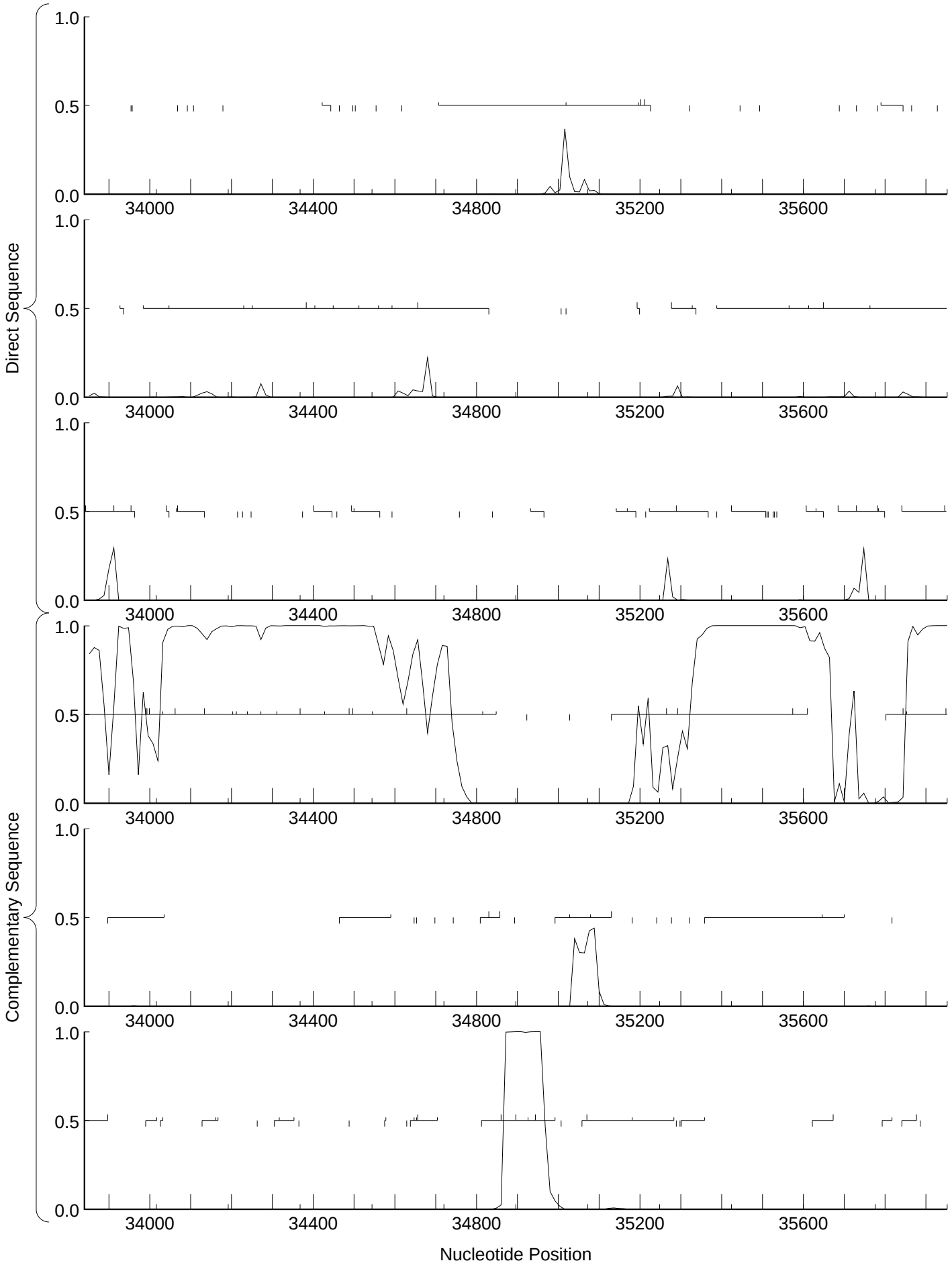


complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 16/26

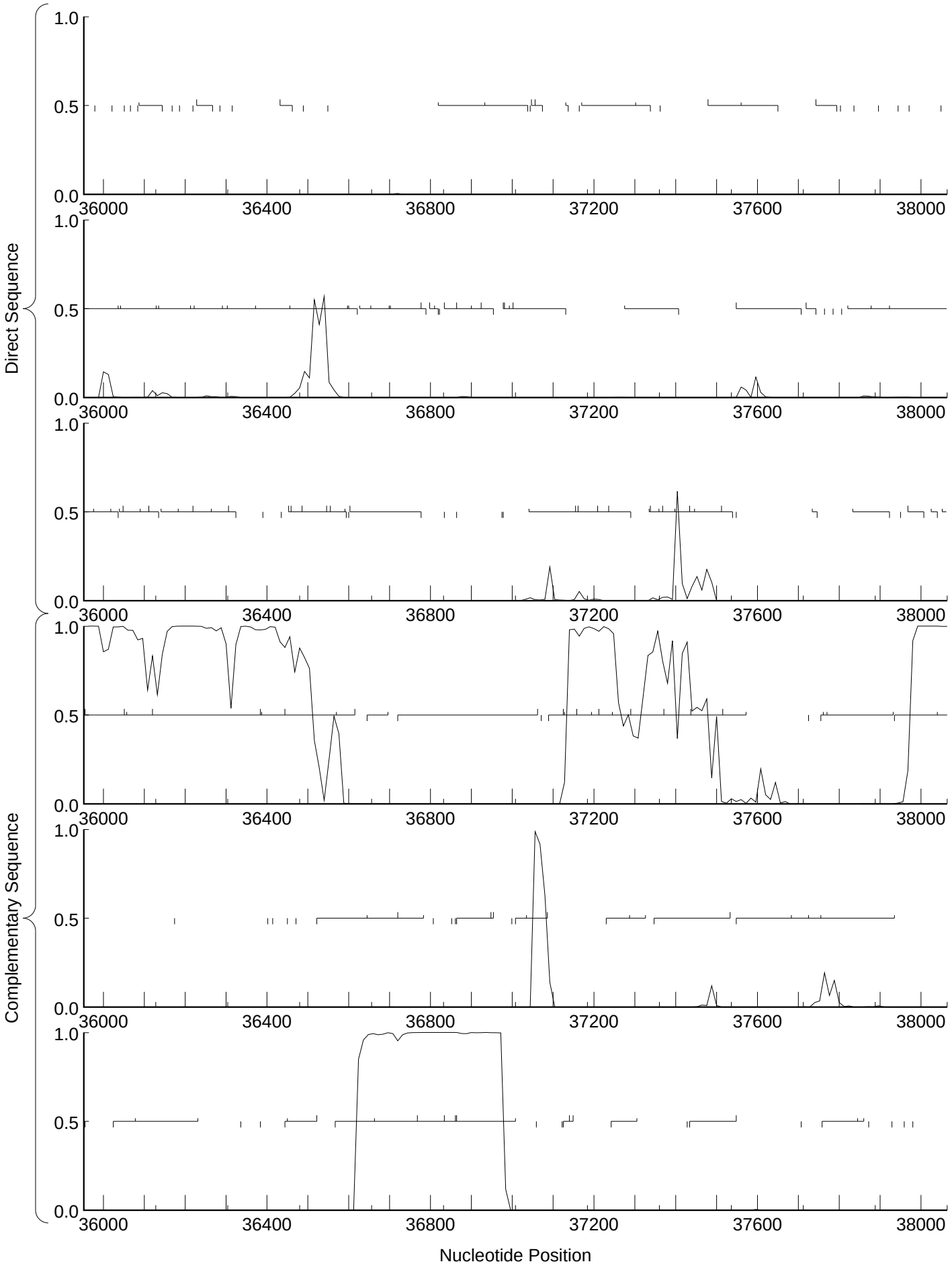


complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 17/26

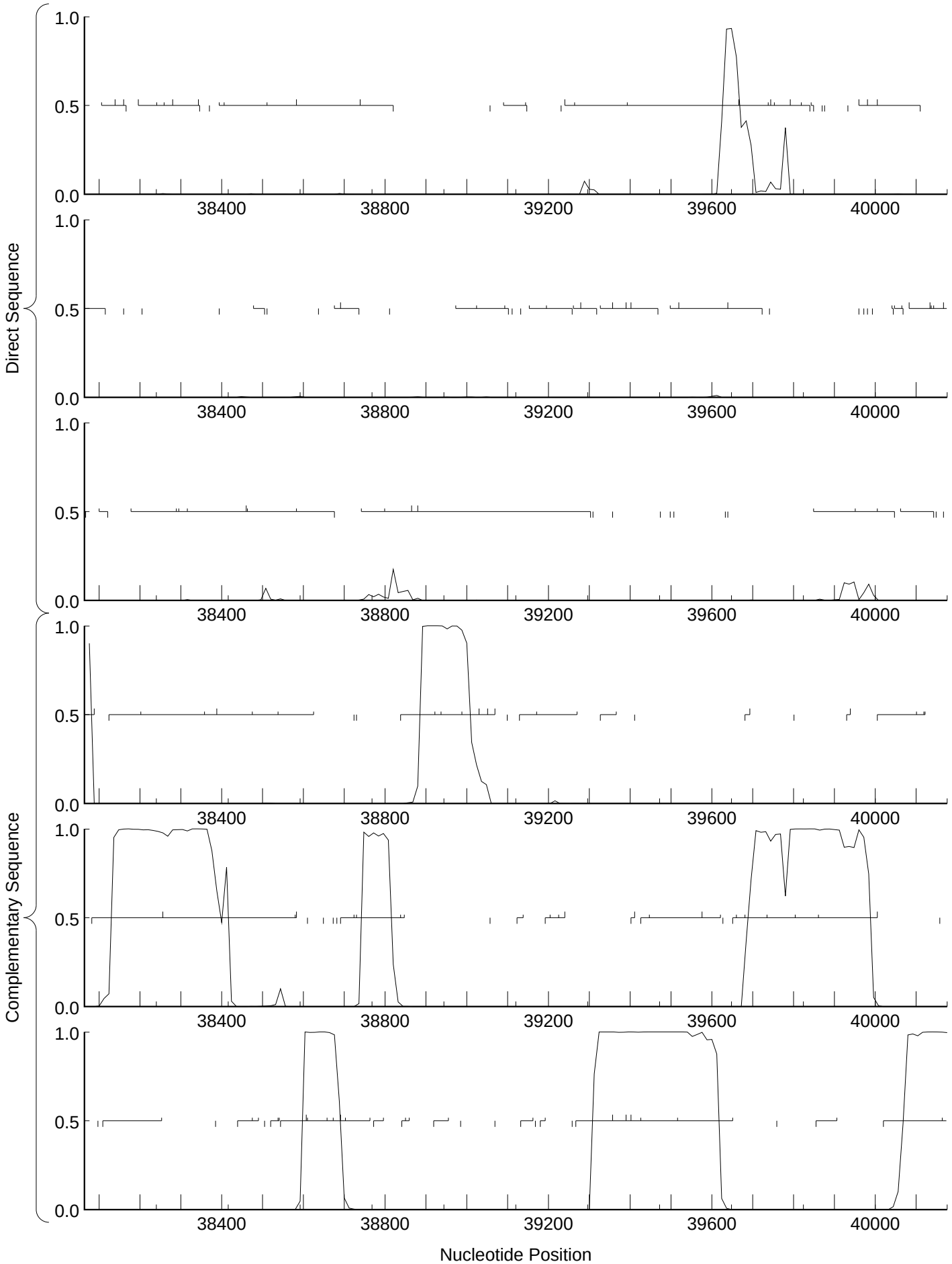



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

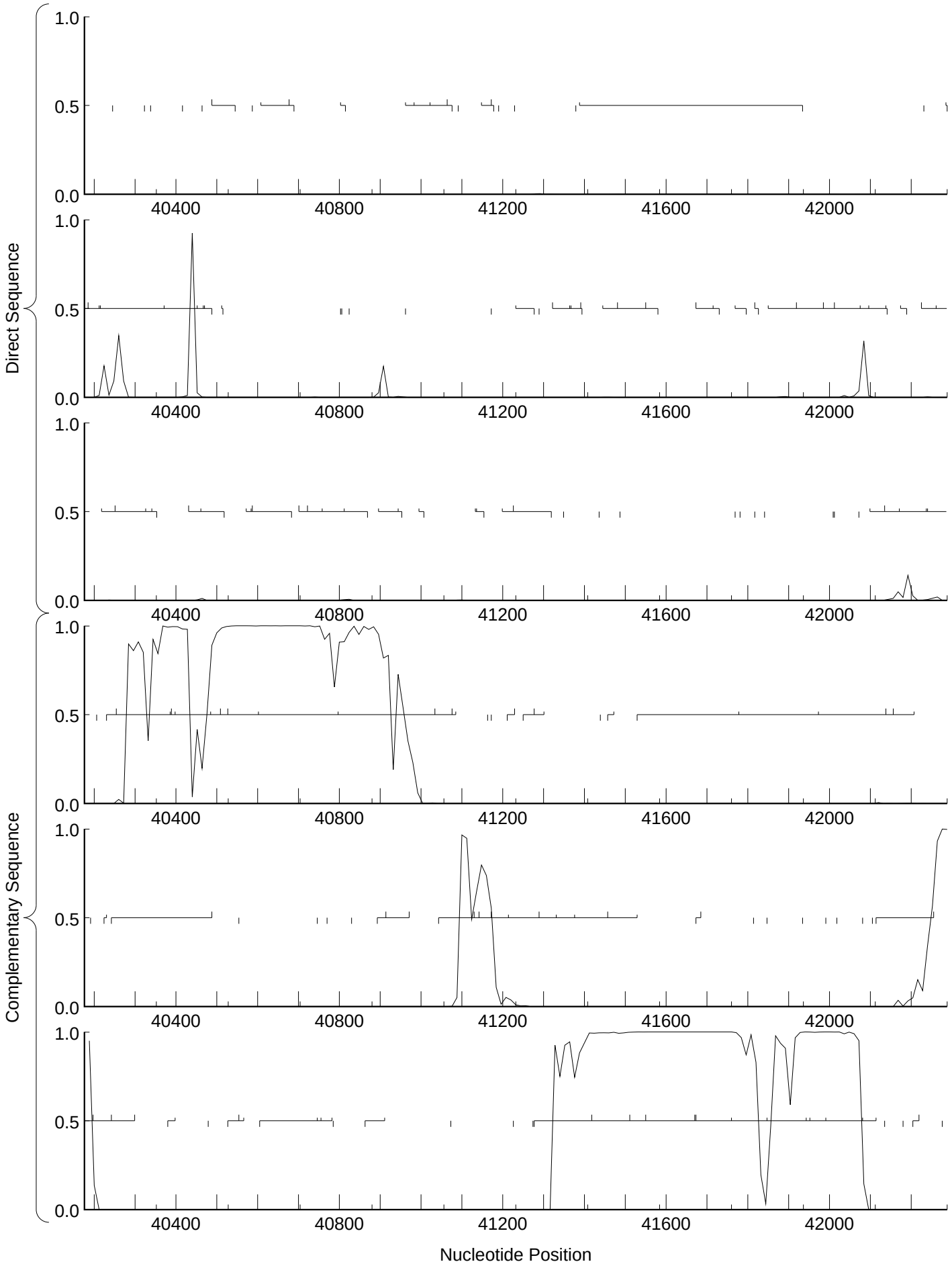
complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 19/26



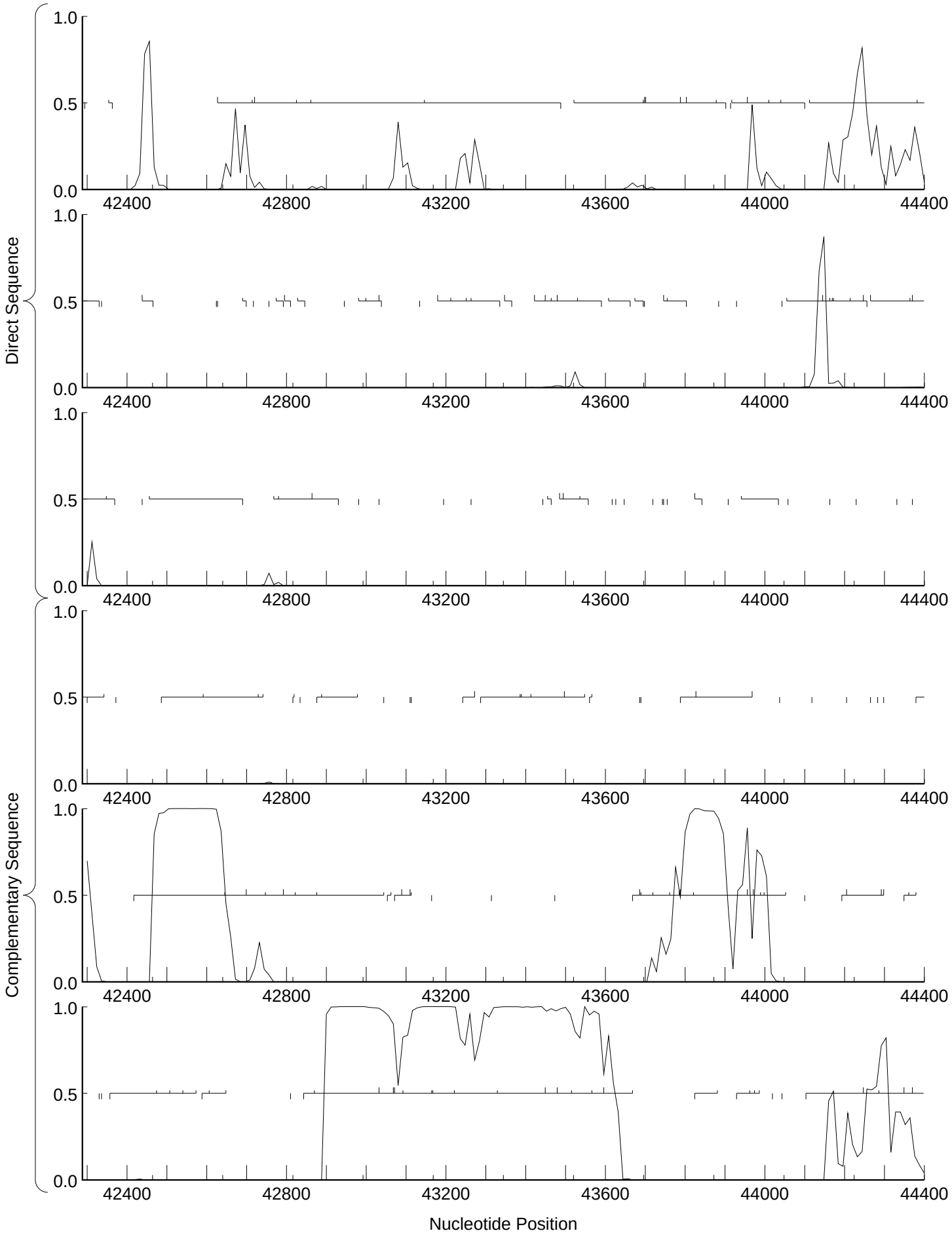
complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 20/26



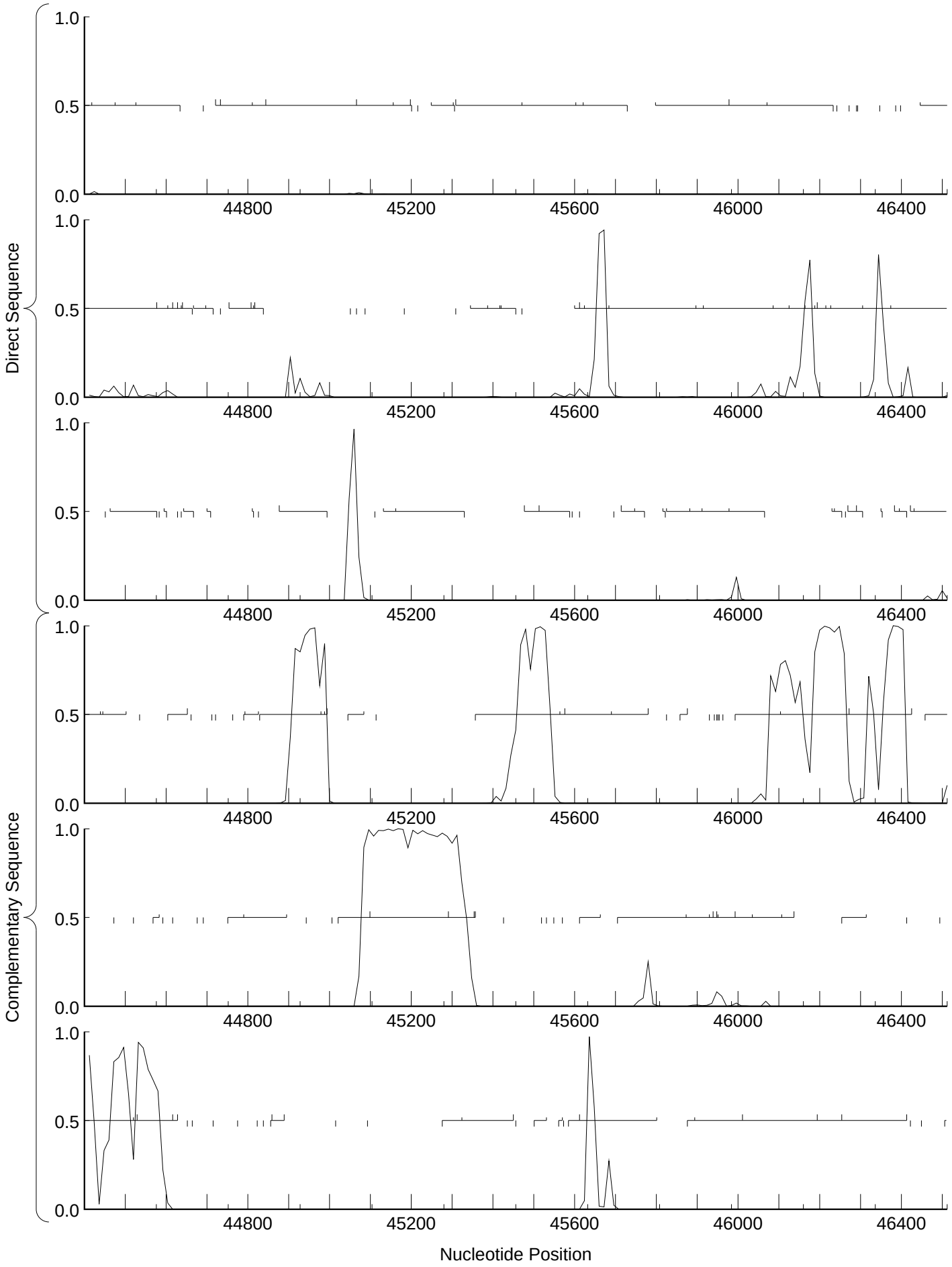
complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 21/26

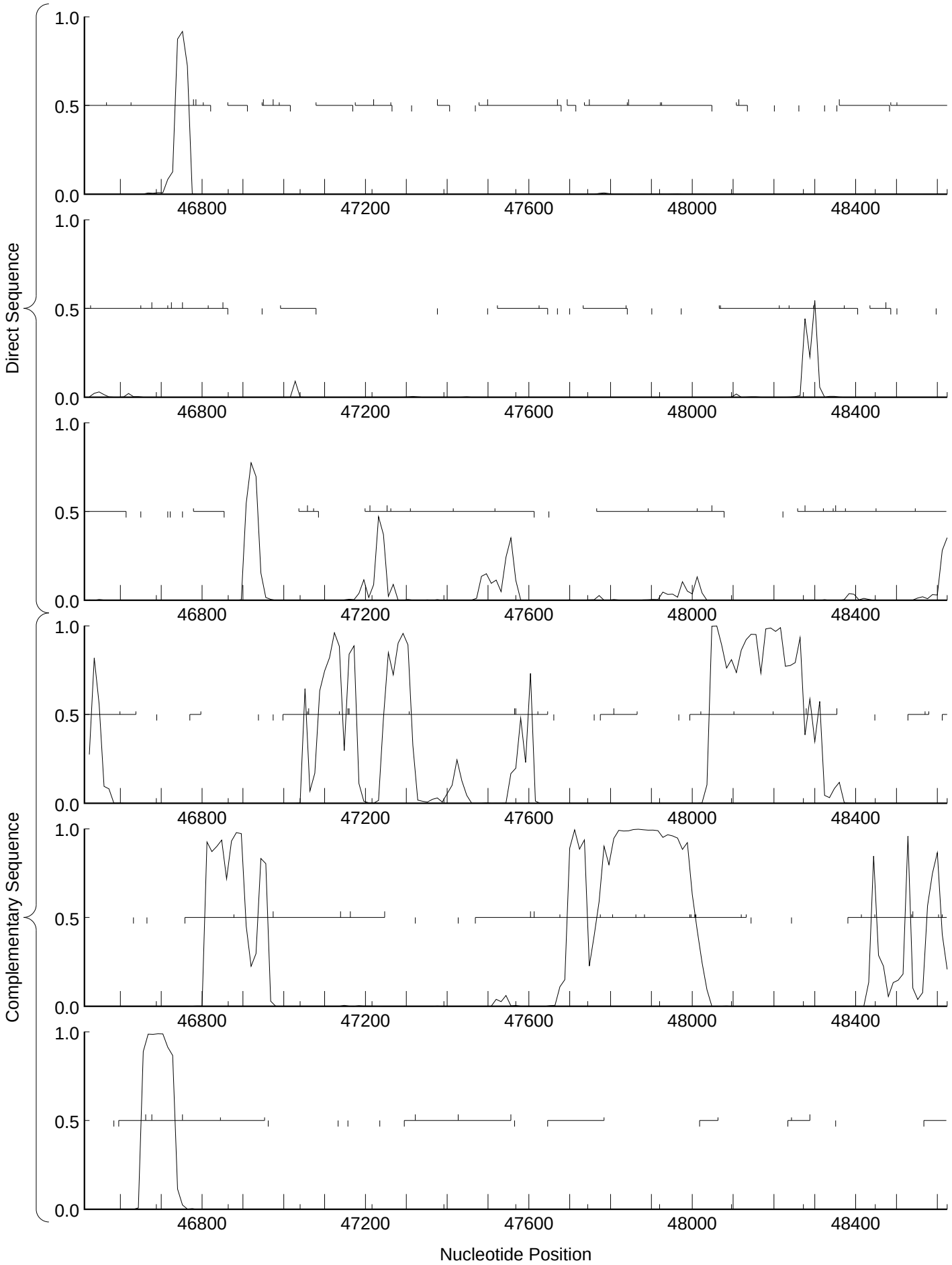


complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 22/26

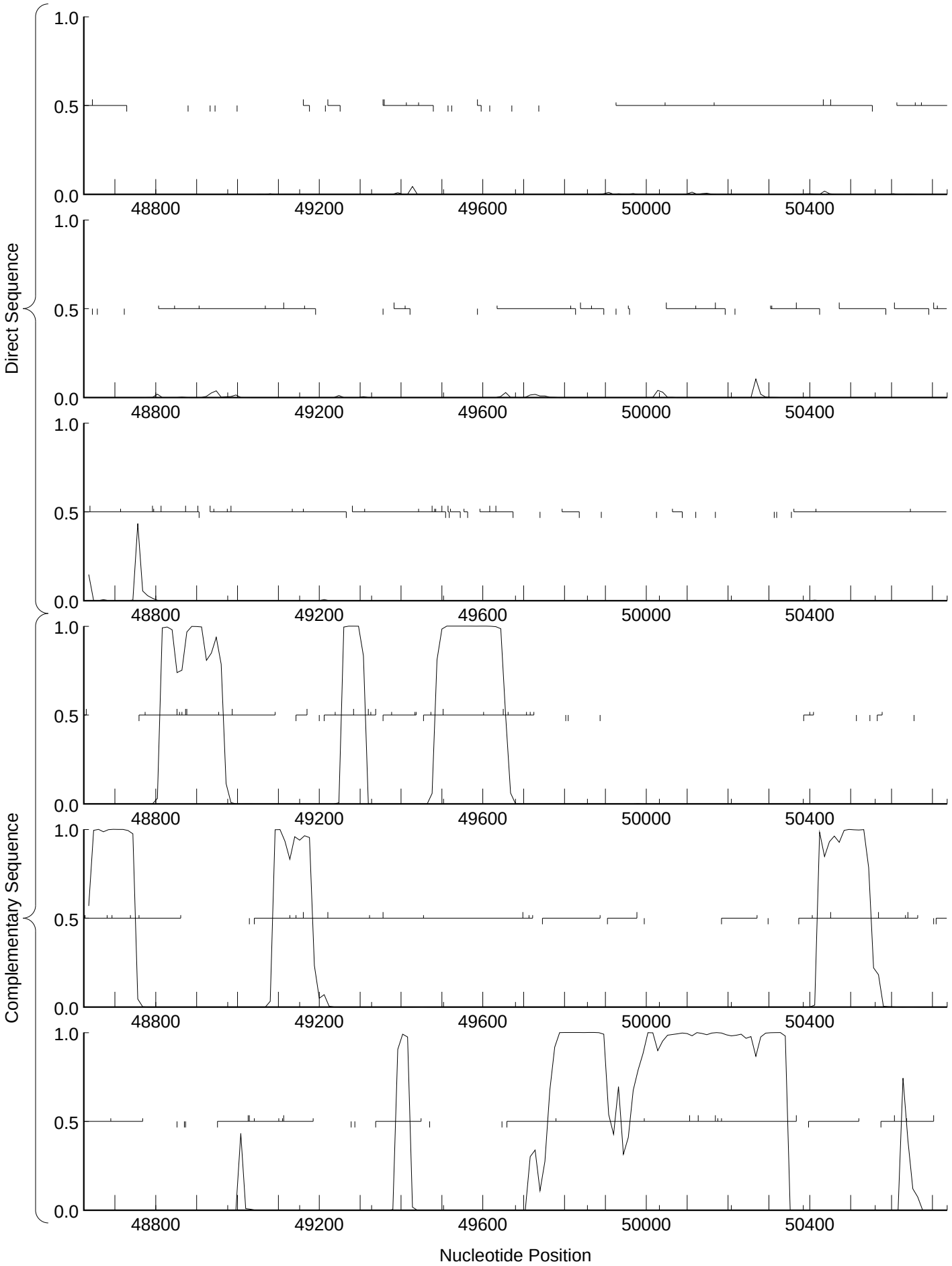


complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 23/26




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
46800
47200
47600
48000
48400
Nucleotide Position

complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 25/26



complete sequence, 52566 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 26/26

