GeneMark

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

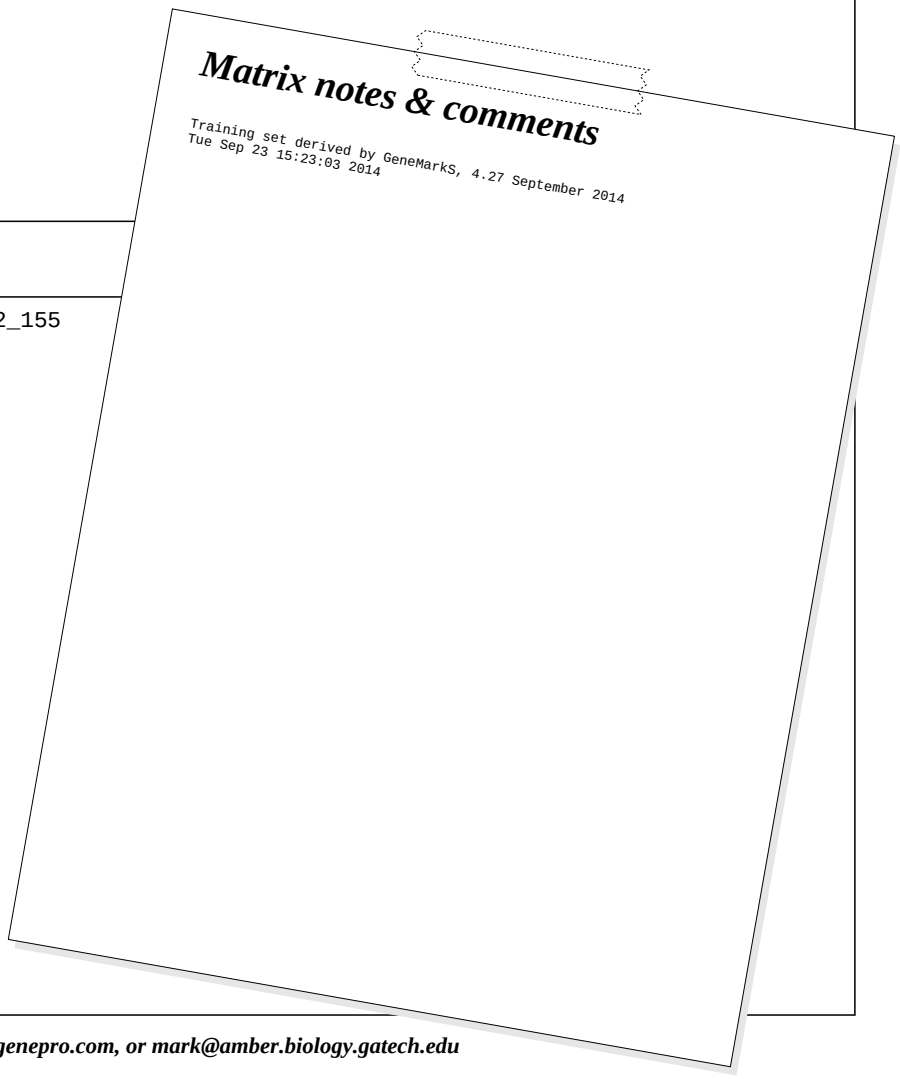
PROGRAM INFORMATION

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Analysis Date    :  3/29/24 at 4:05:44
Pages            :  26
Sequence Length  :  51094 bp
GC Content       :   64.96%

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

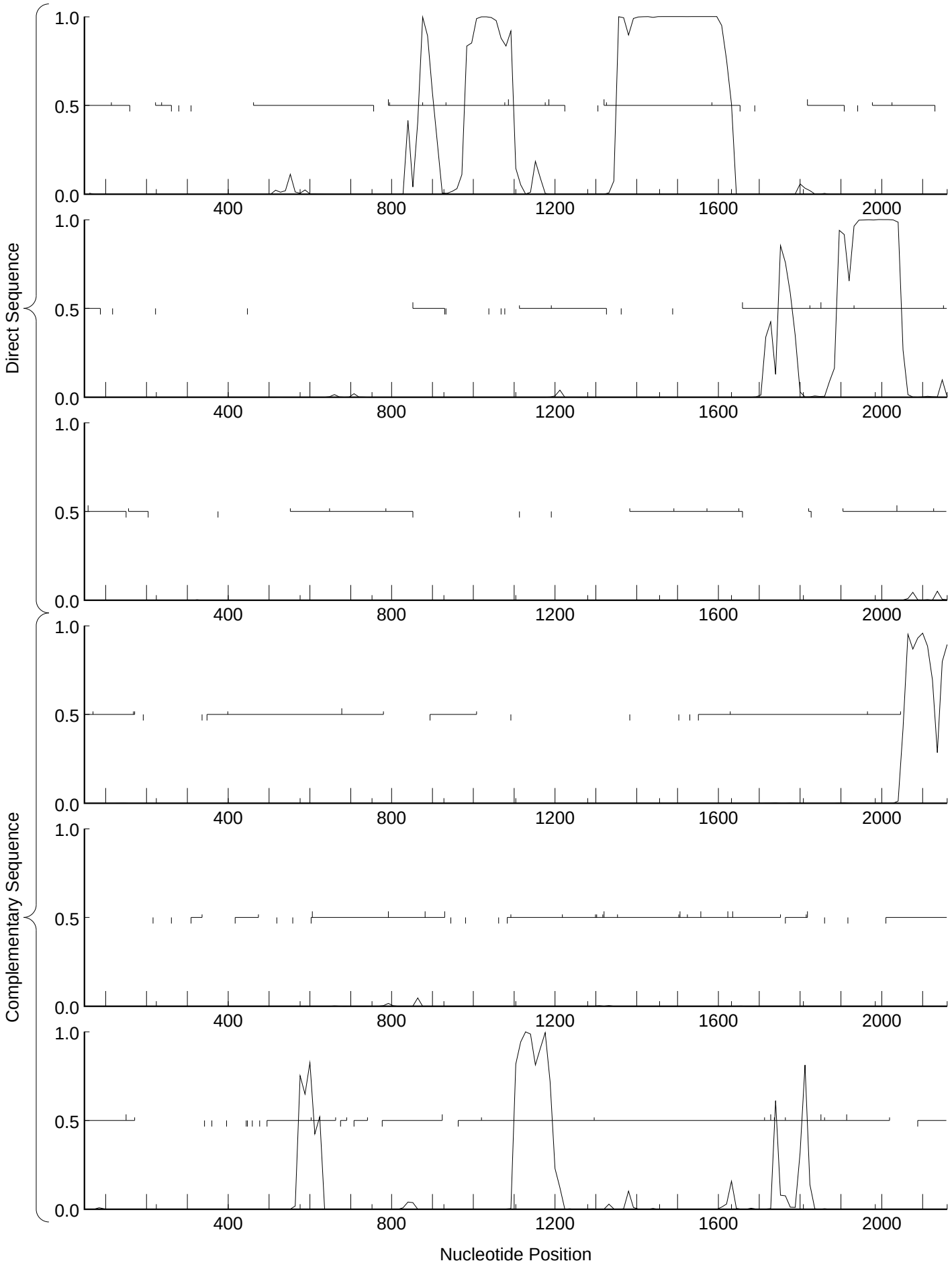
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Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

Order      :  4
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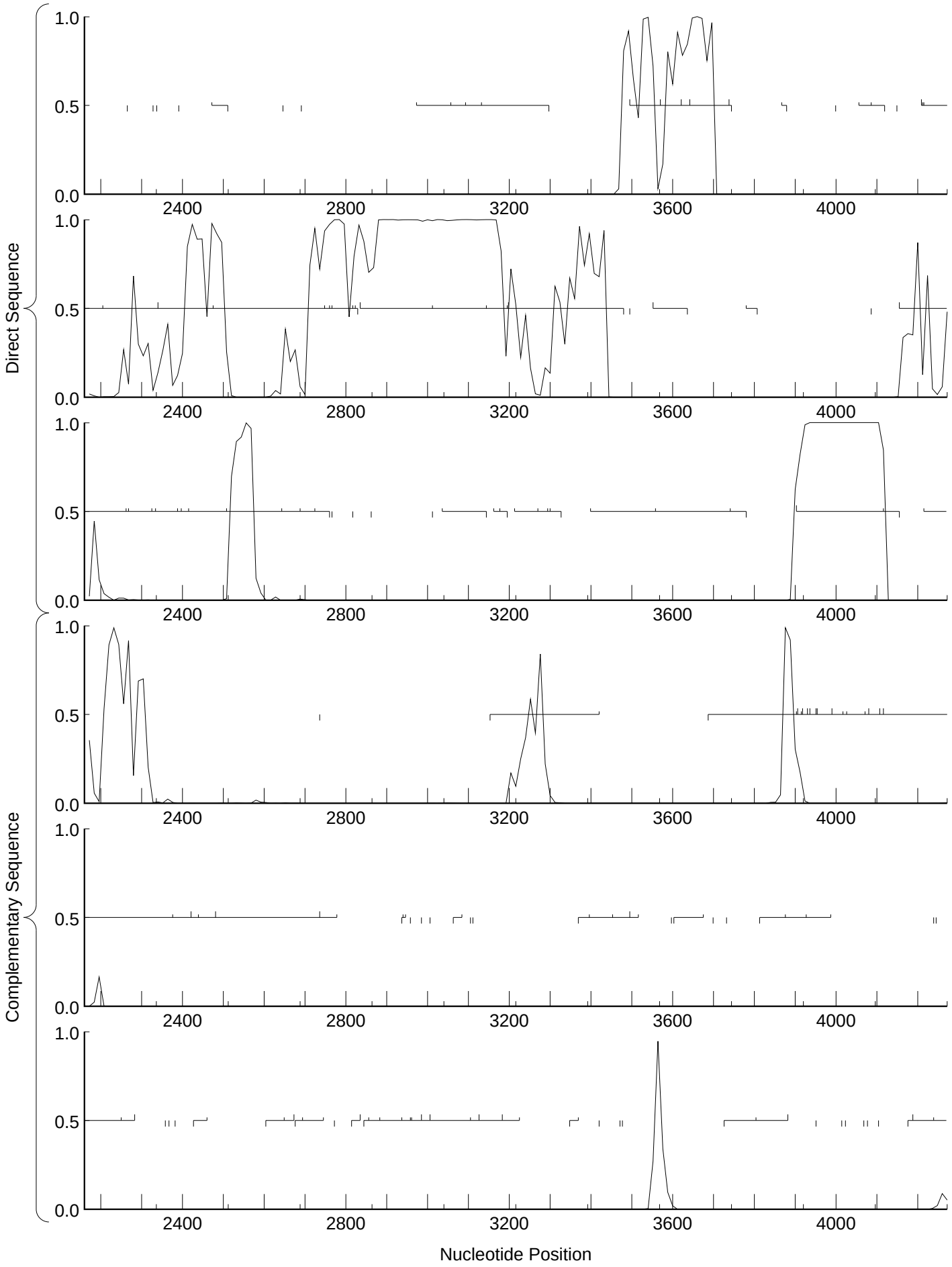

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

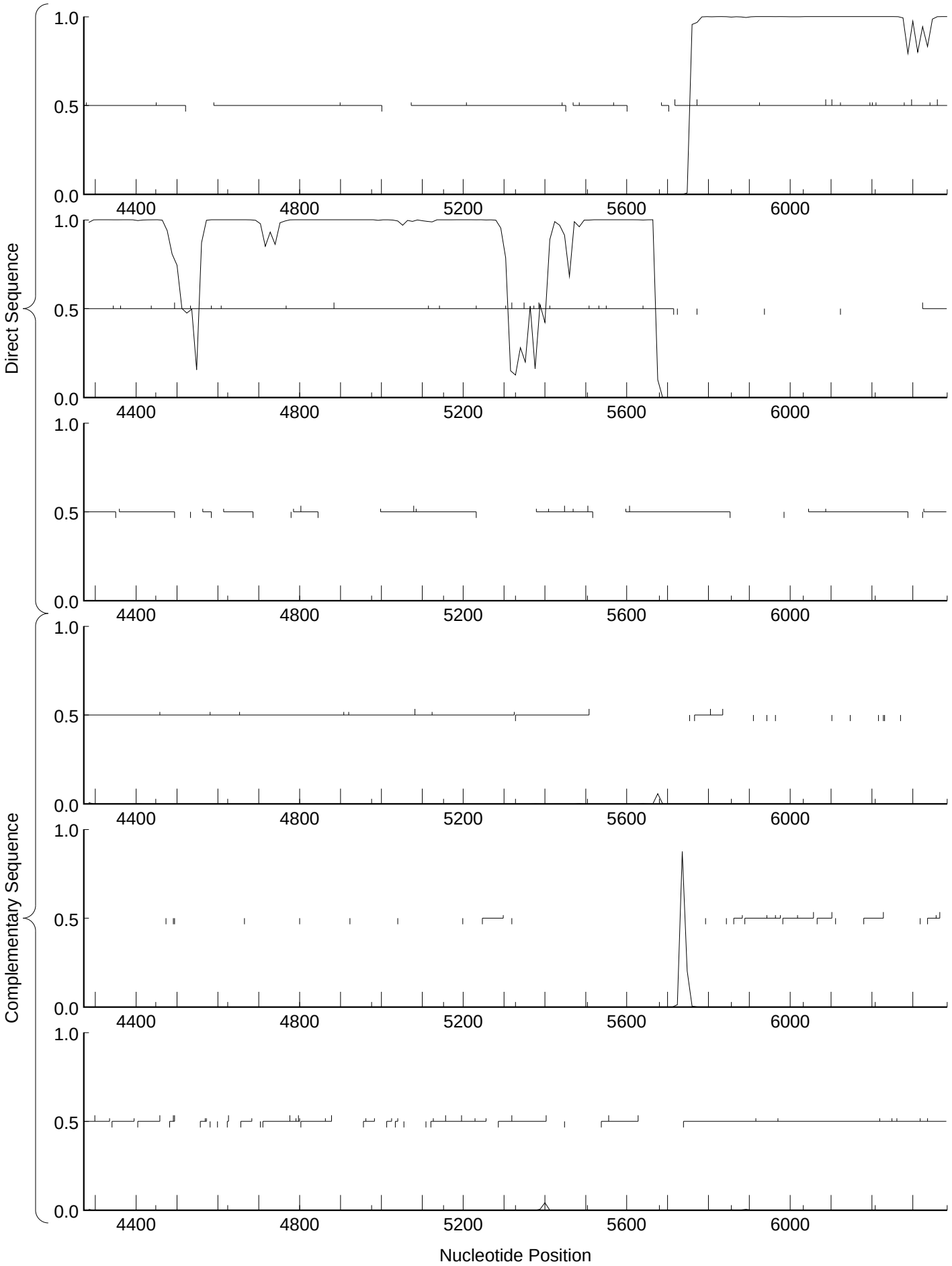
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 2/26



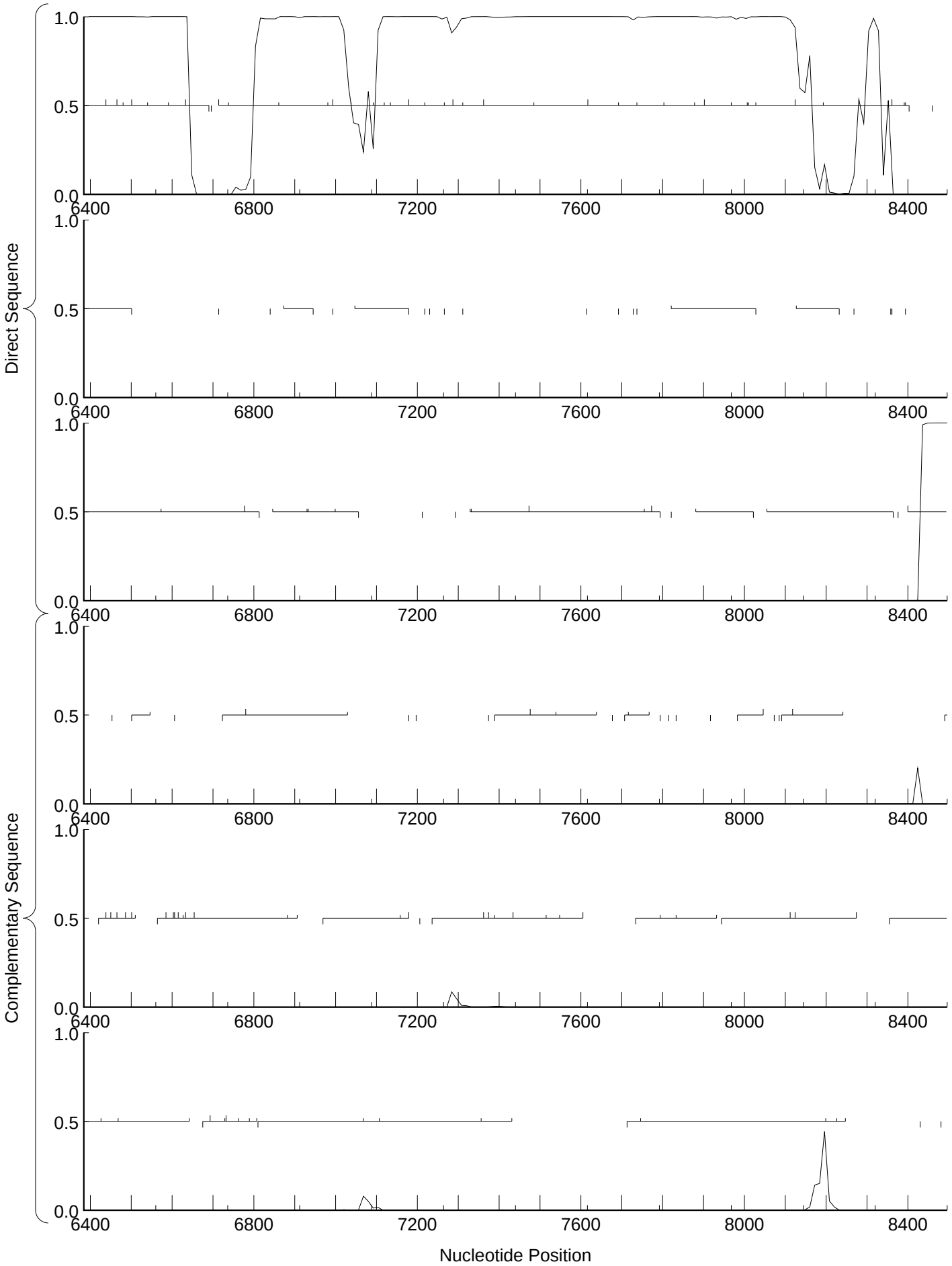
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 3/26



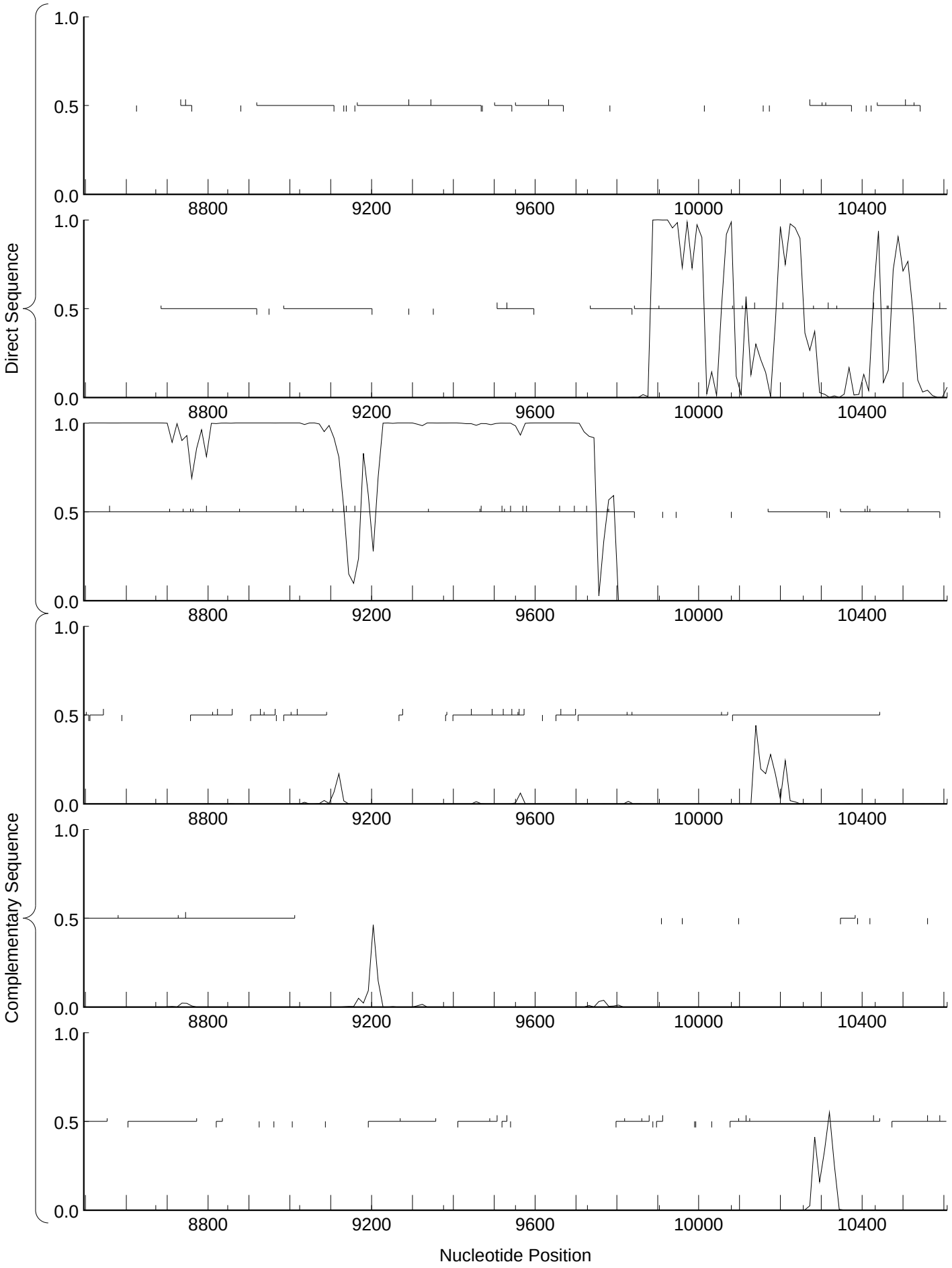
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 4/26



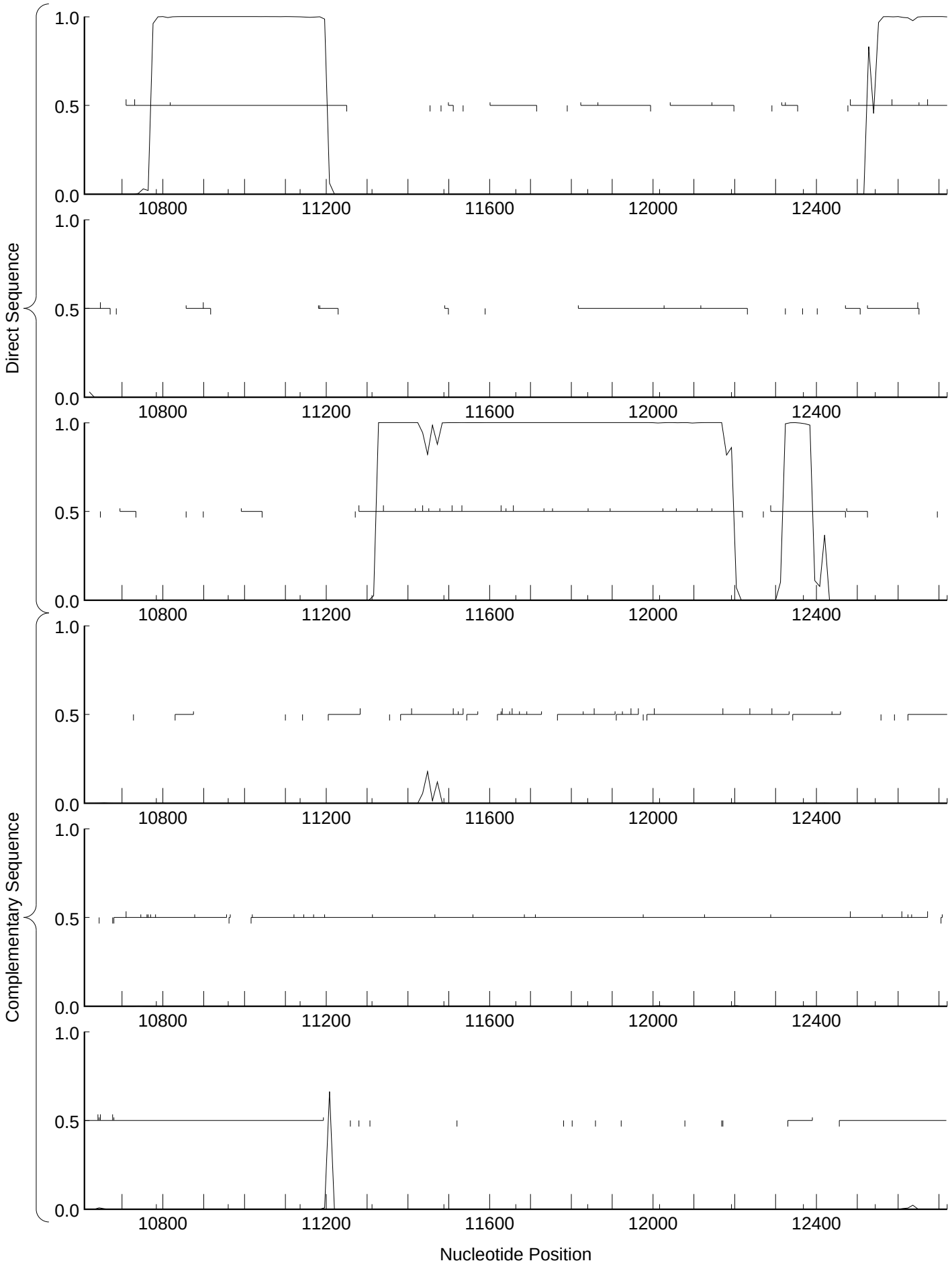
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 5/26



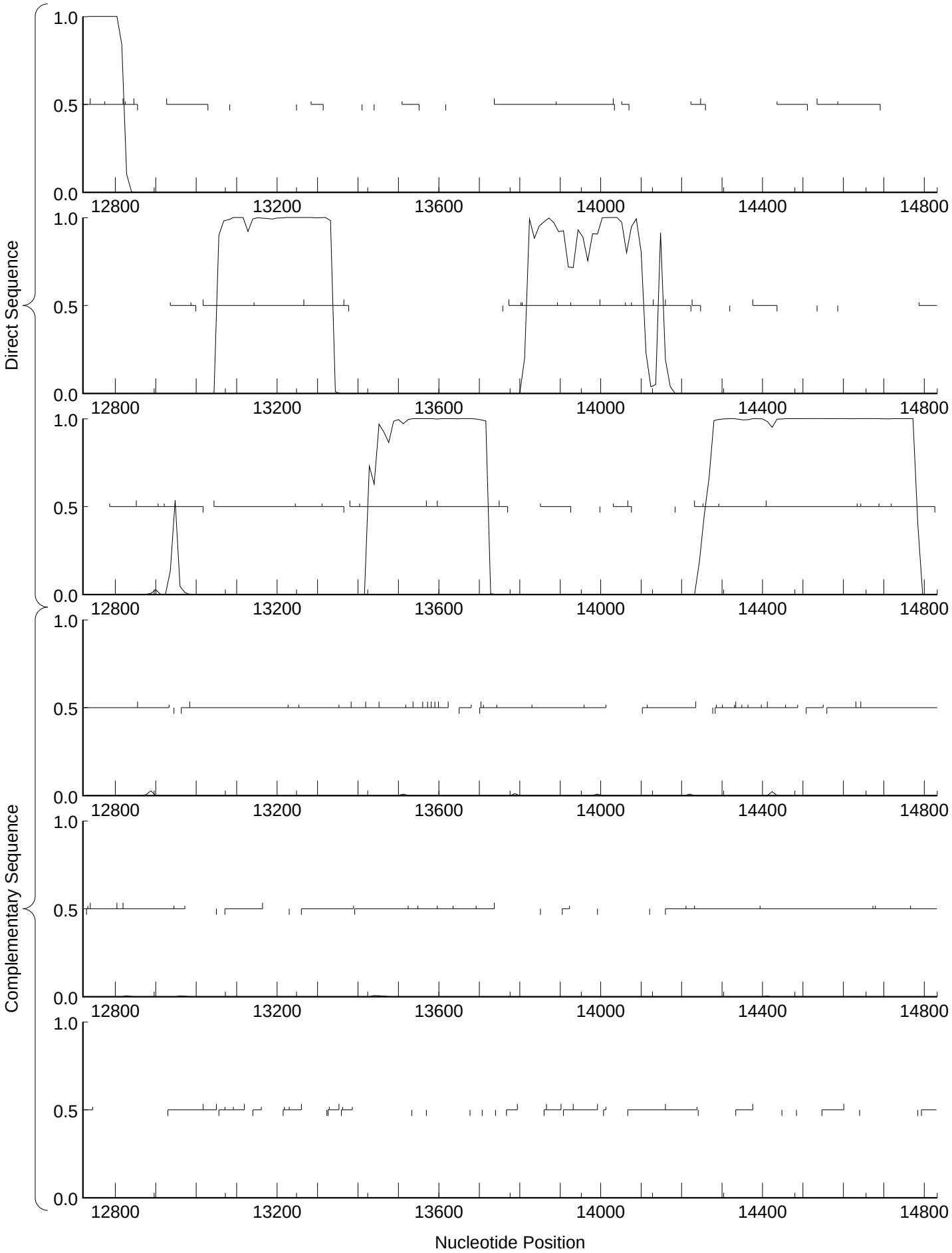
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 6/26



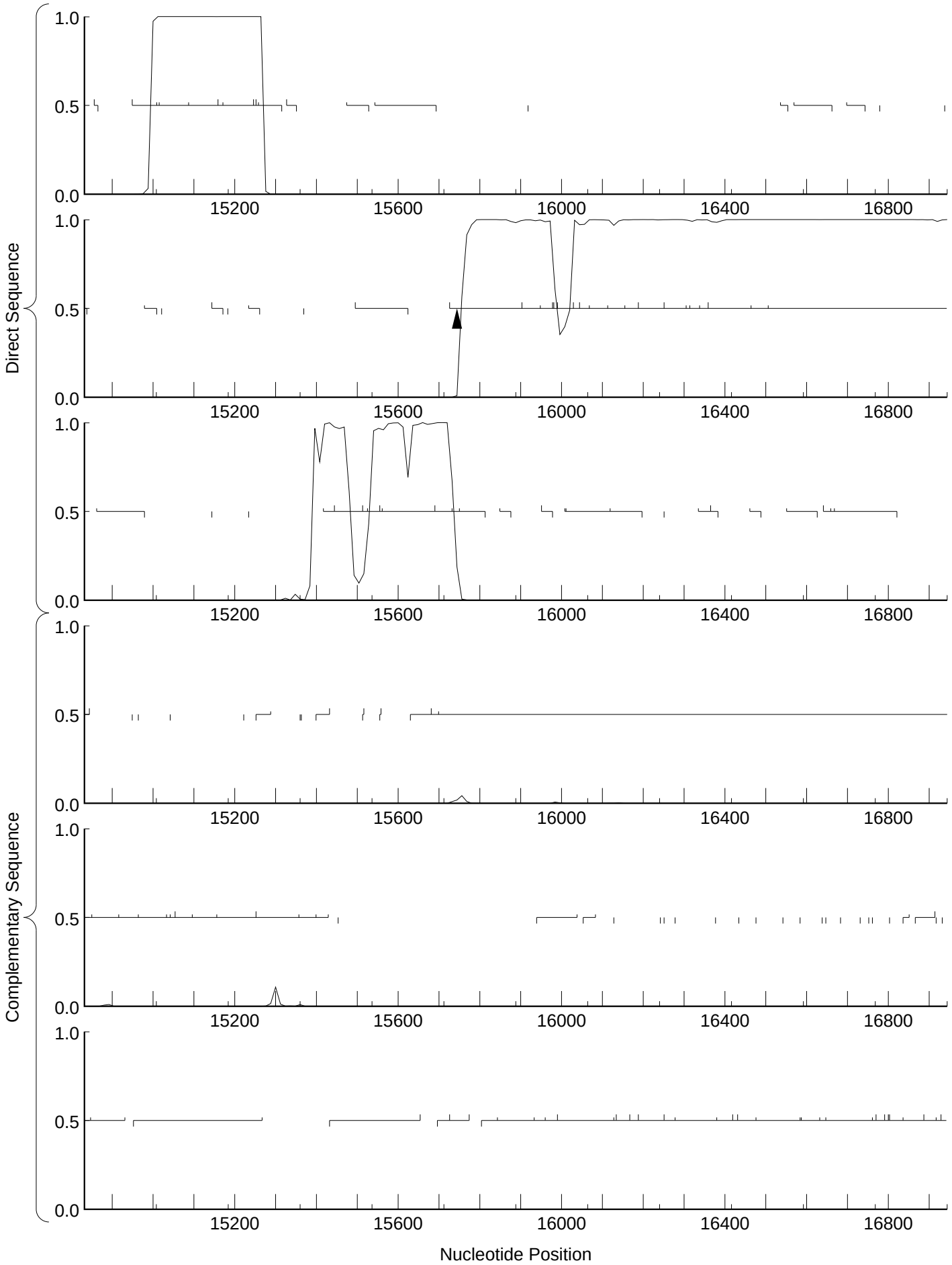
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 7/26



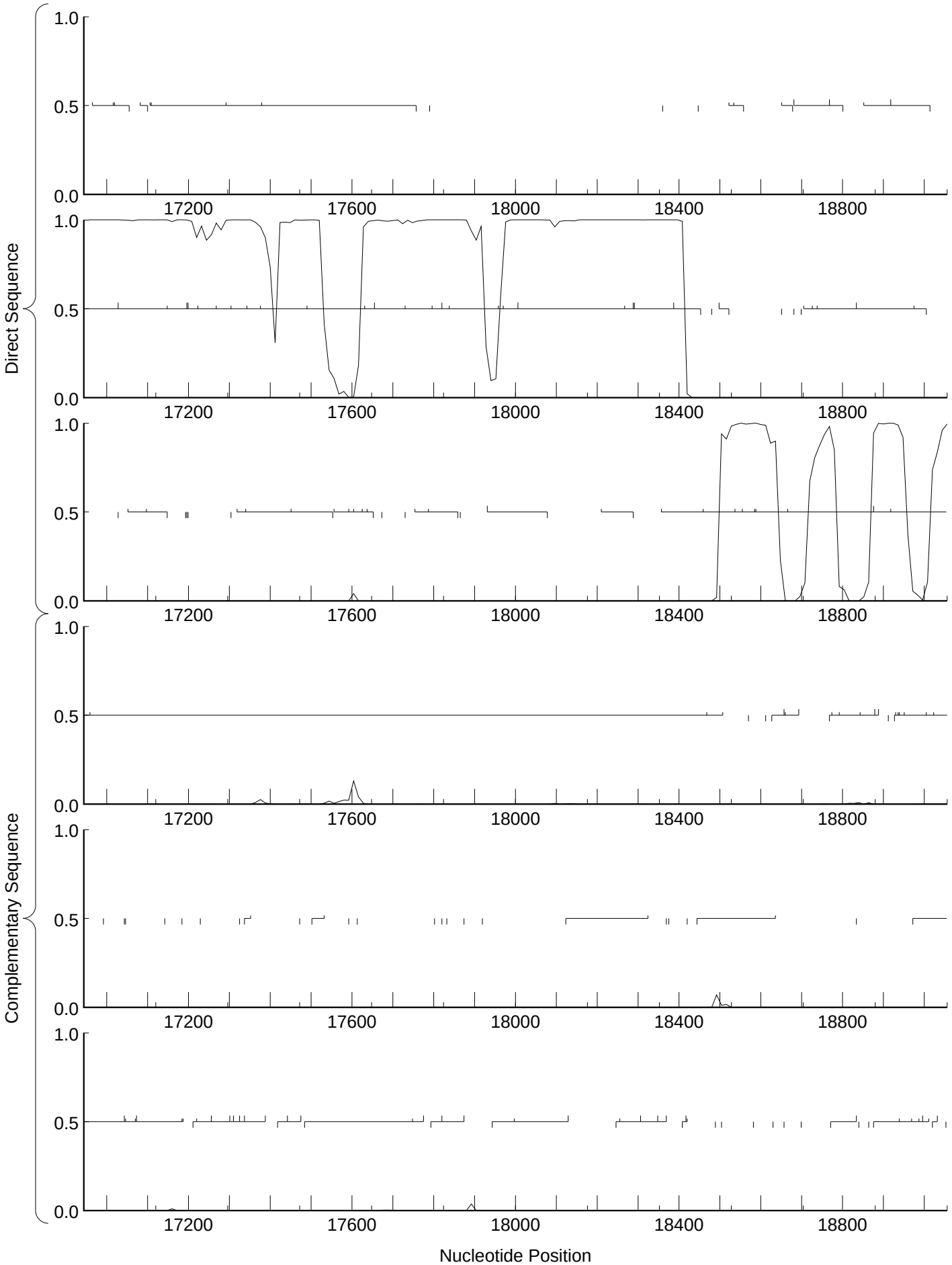
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 8/26



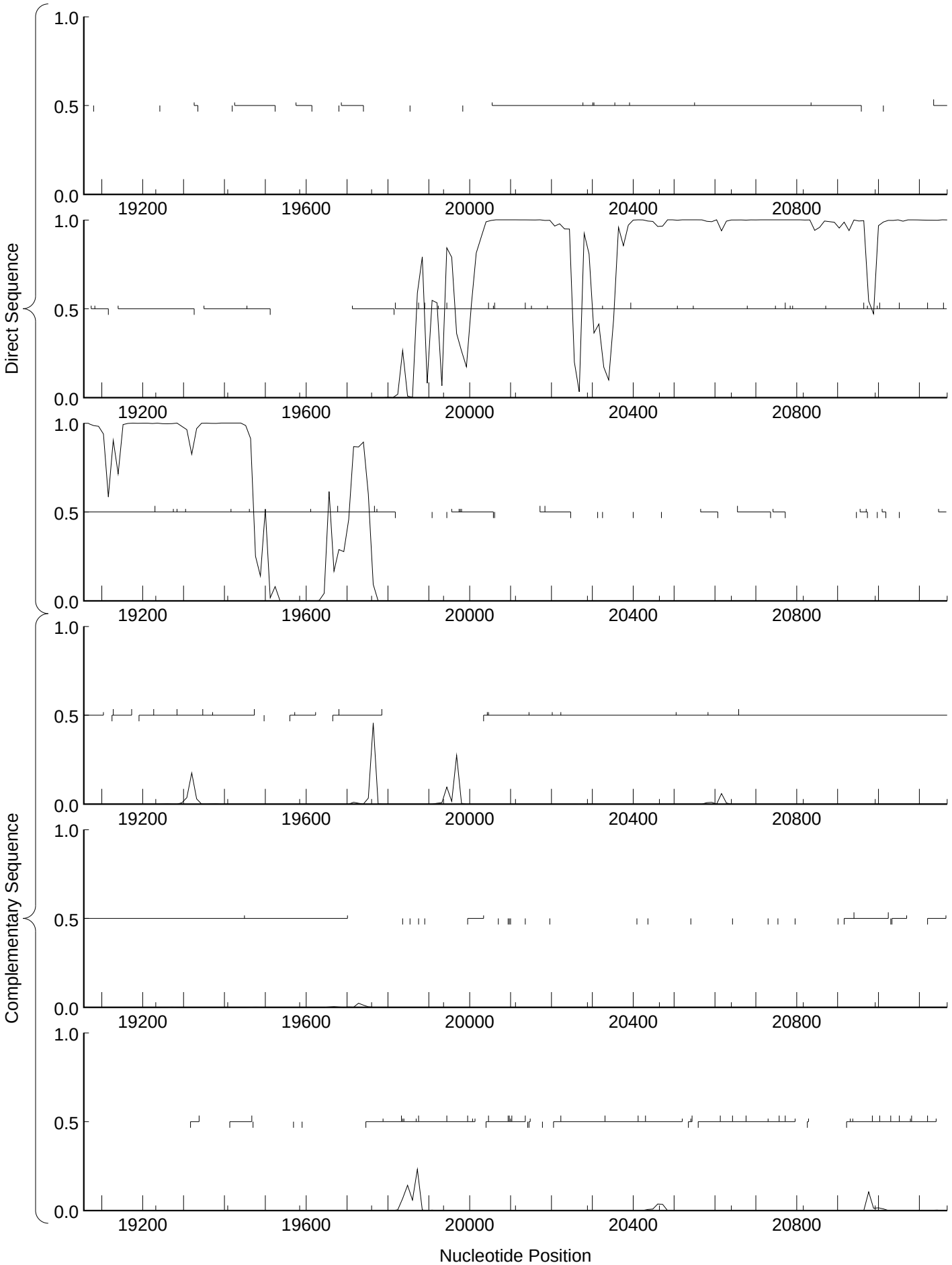
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 9/26



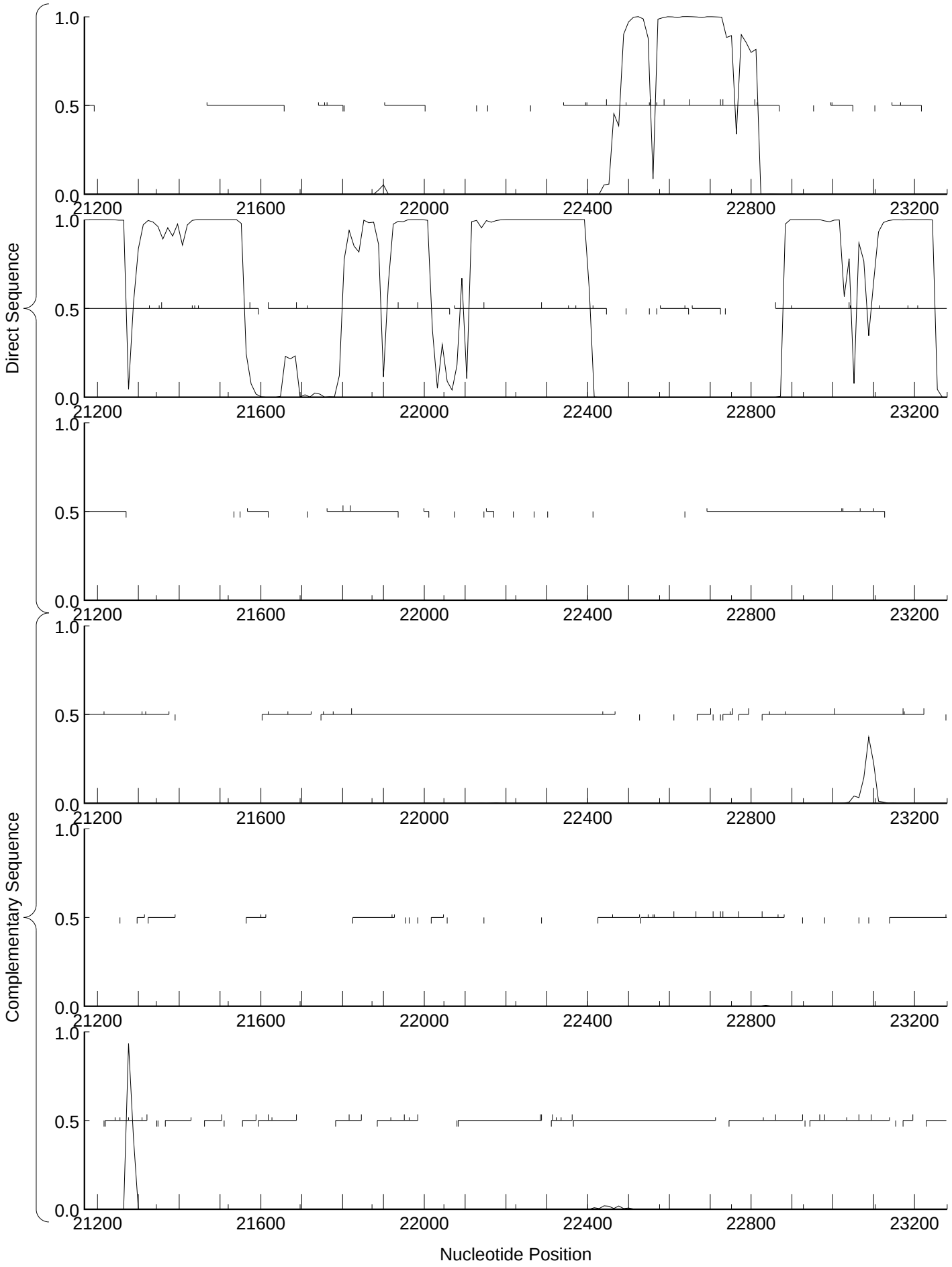
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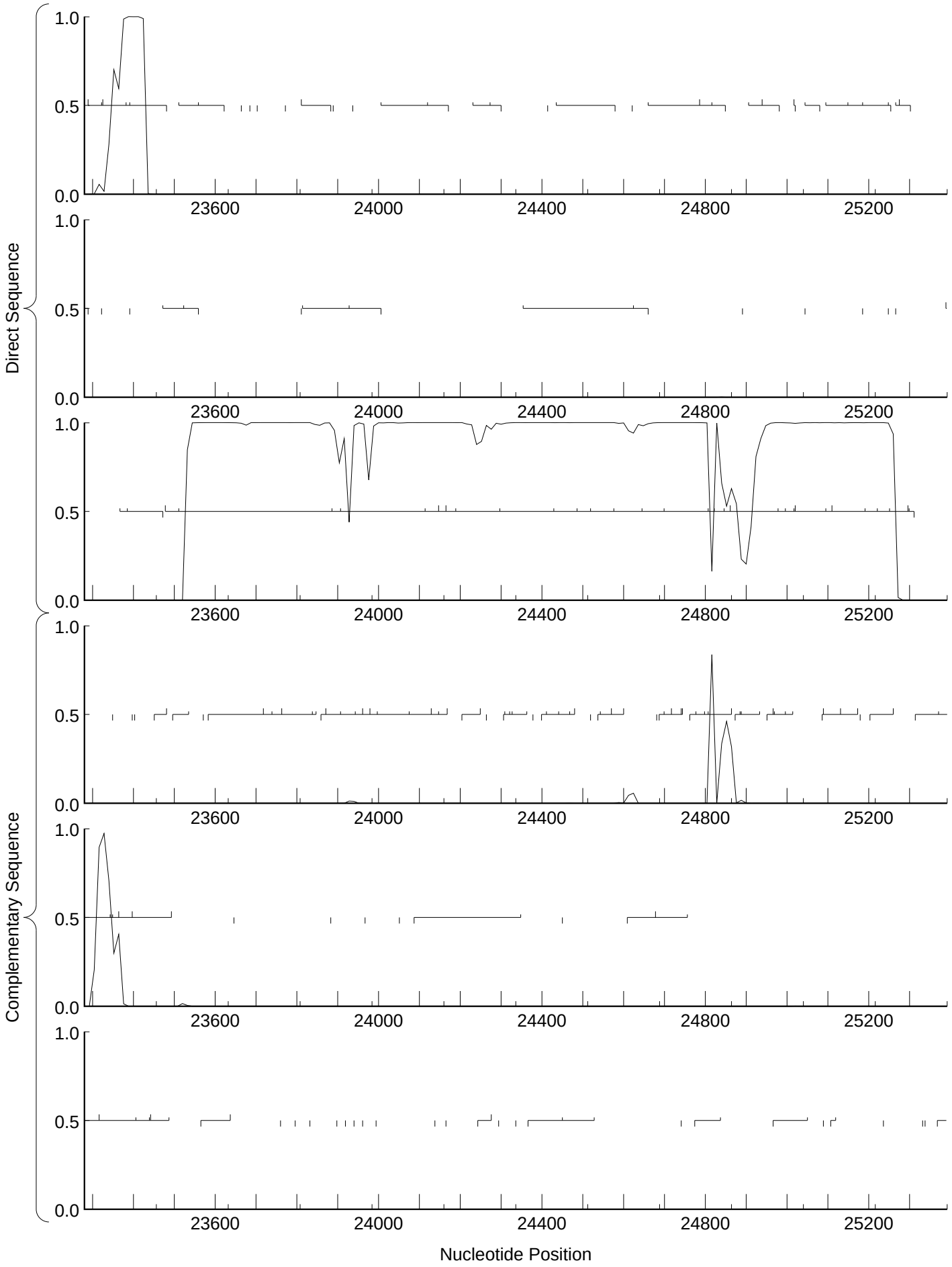
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 11/26

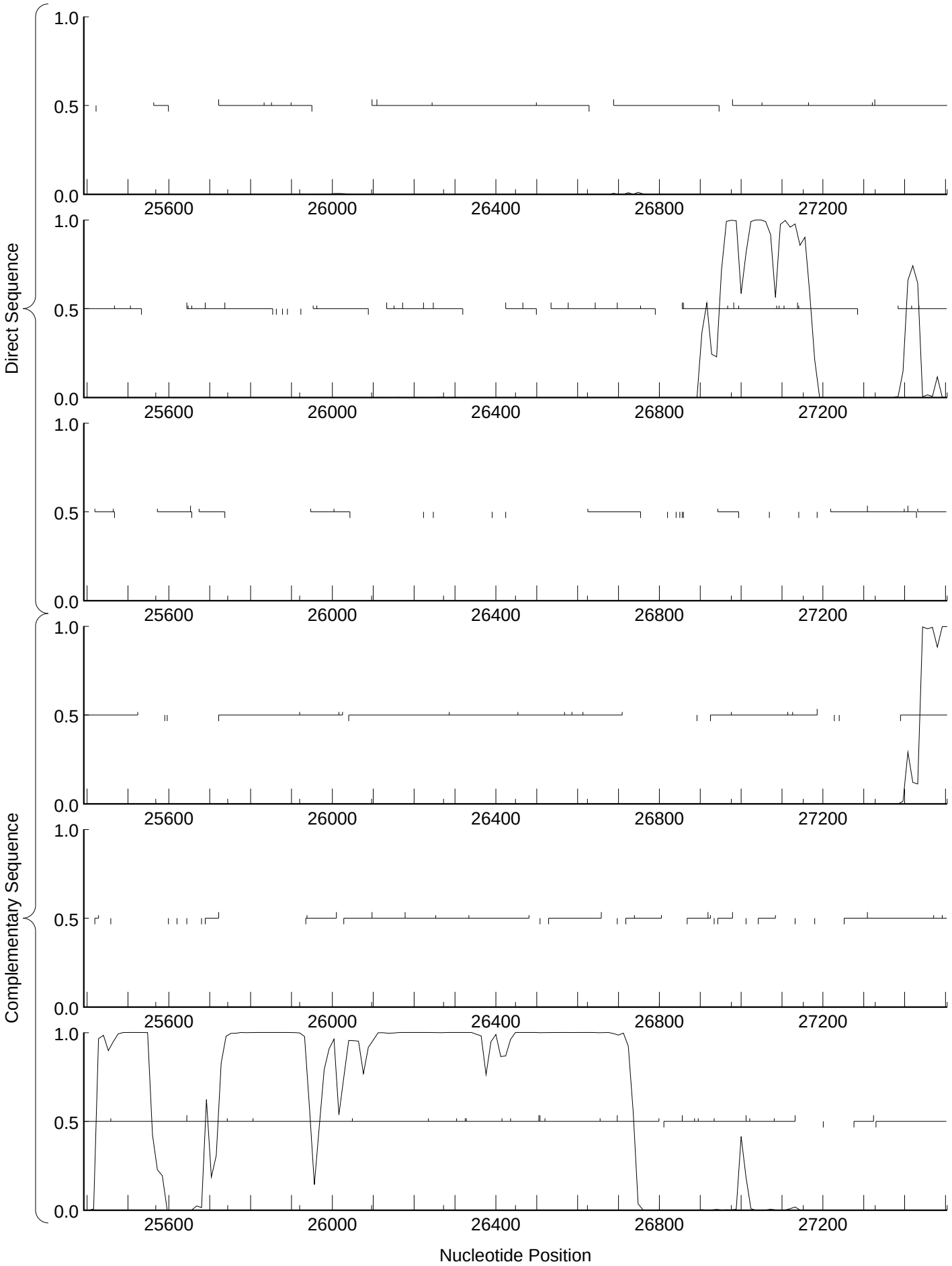


complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 12/26

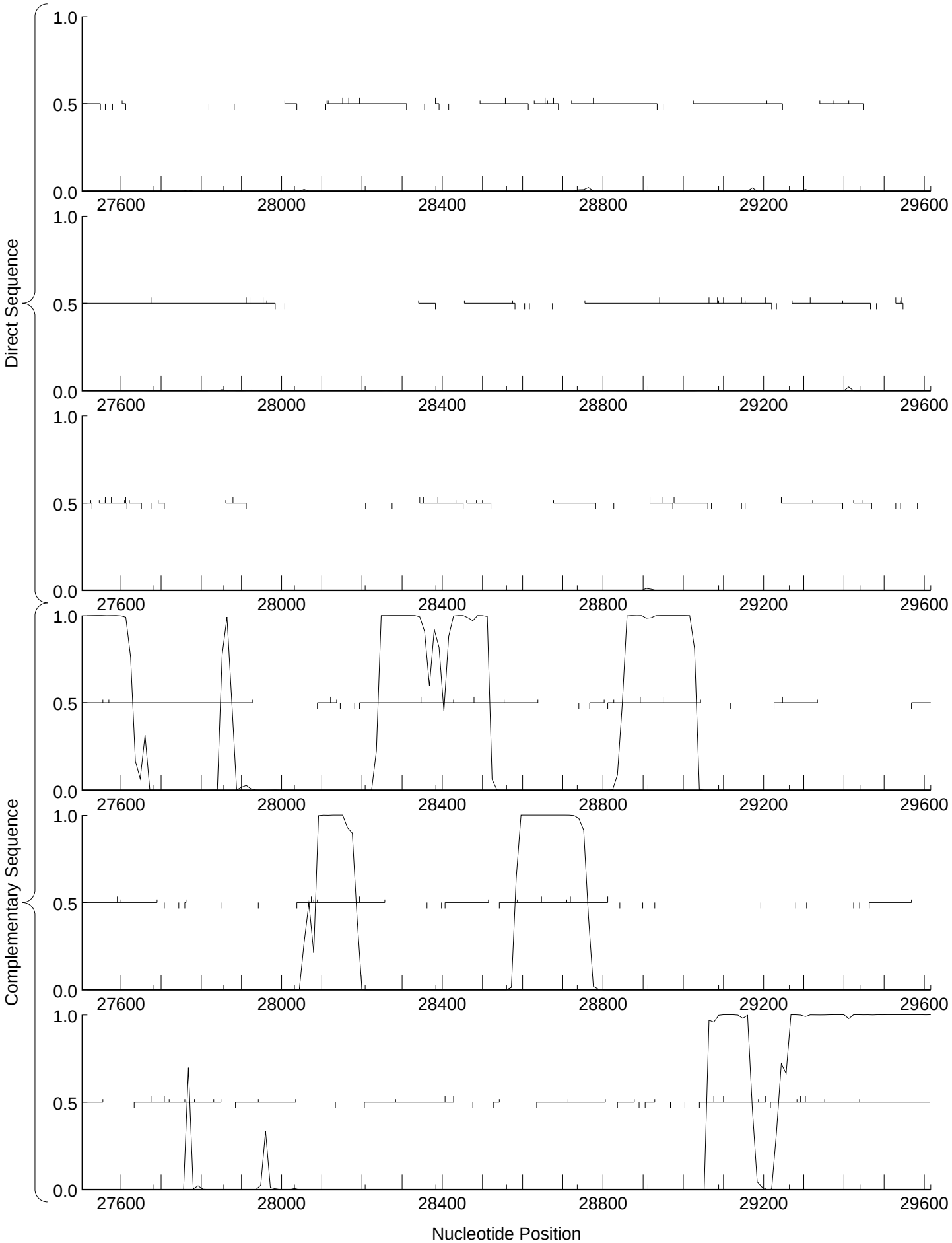


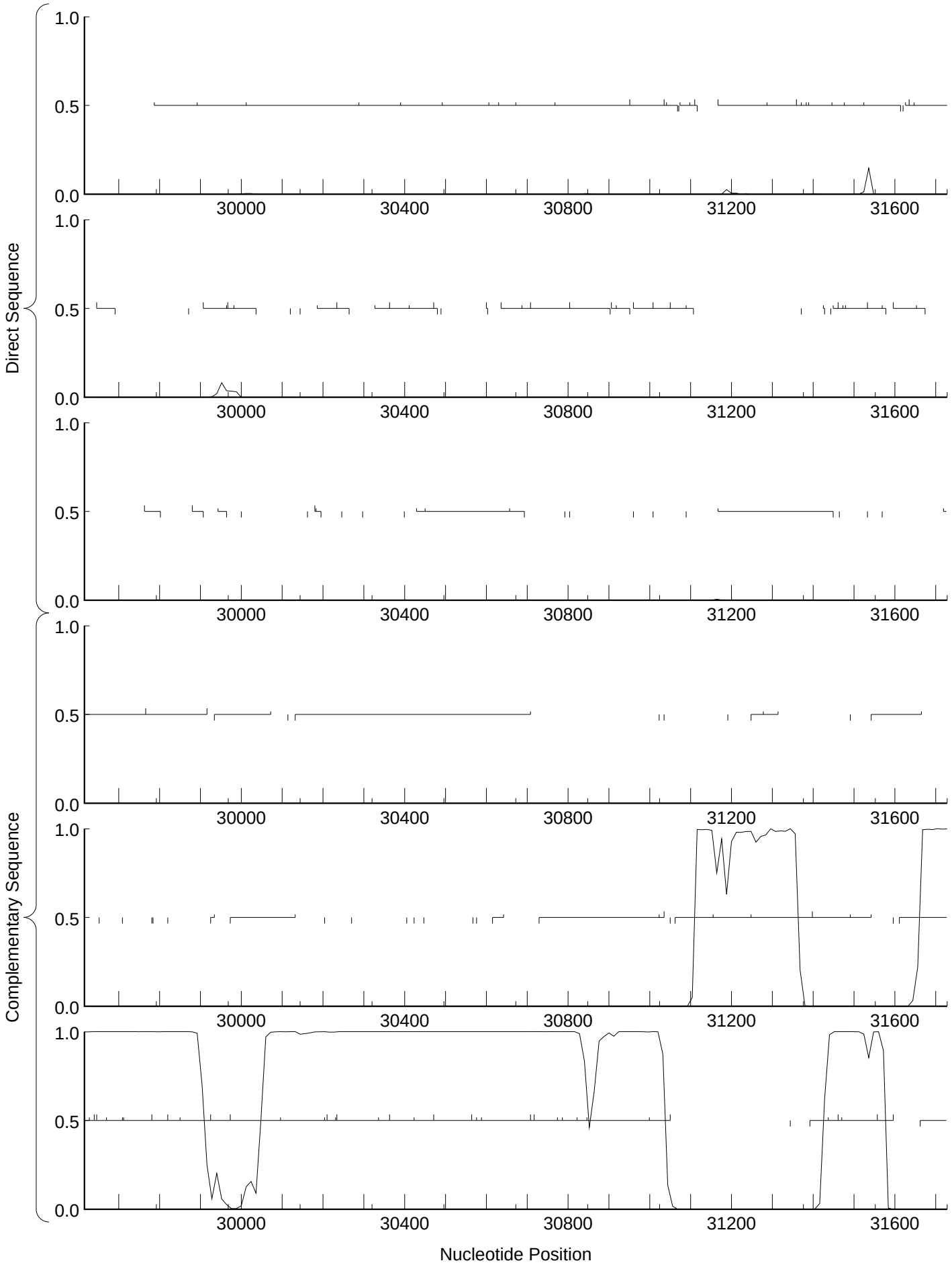
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 13/26



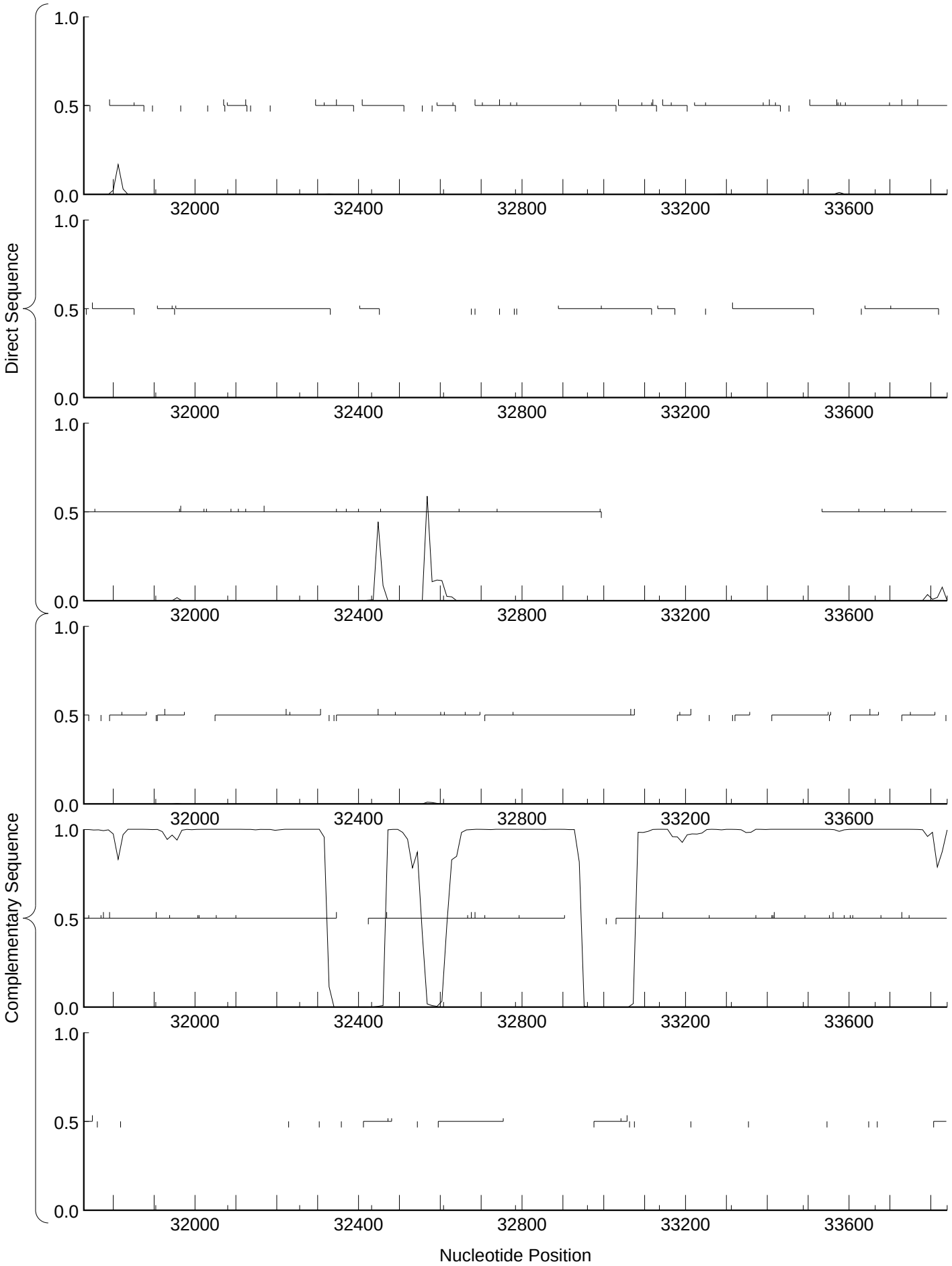

Direct Sequence
Complementary Sequence
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0.5
0.0
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26000
26400
26800
27200
Nucleotide Position

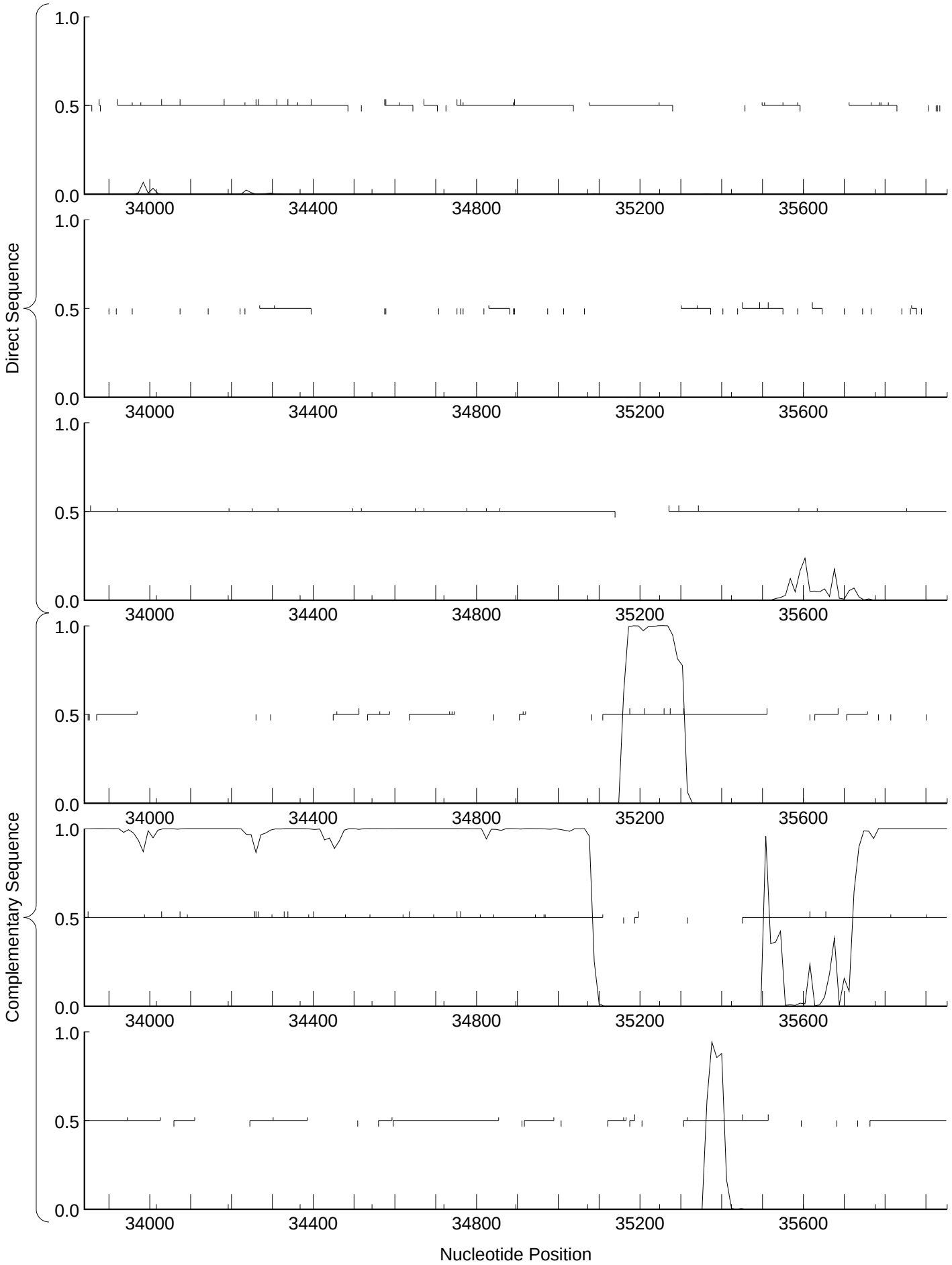
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 15/26



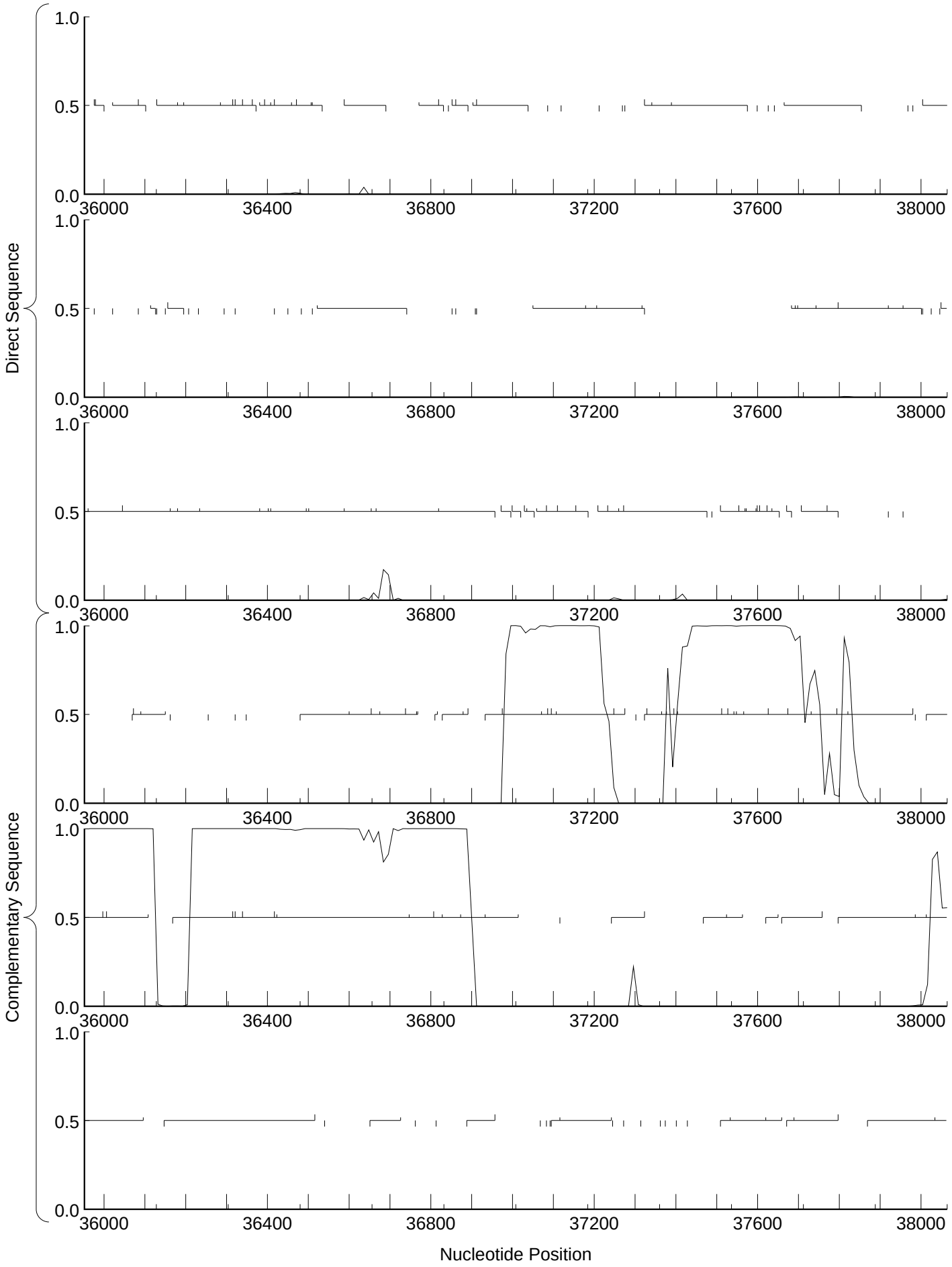

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
30000
30400
30800
31200
31600
Nucleotide Position

complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 17/26

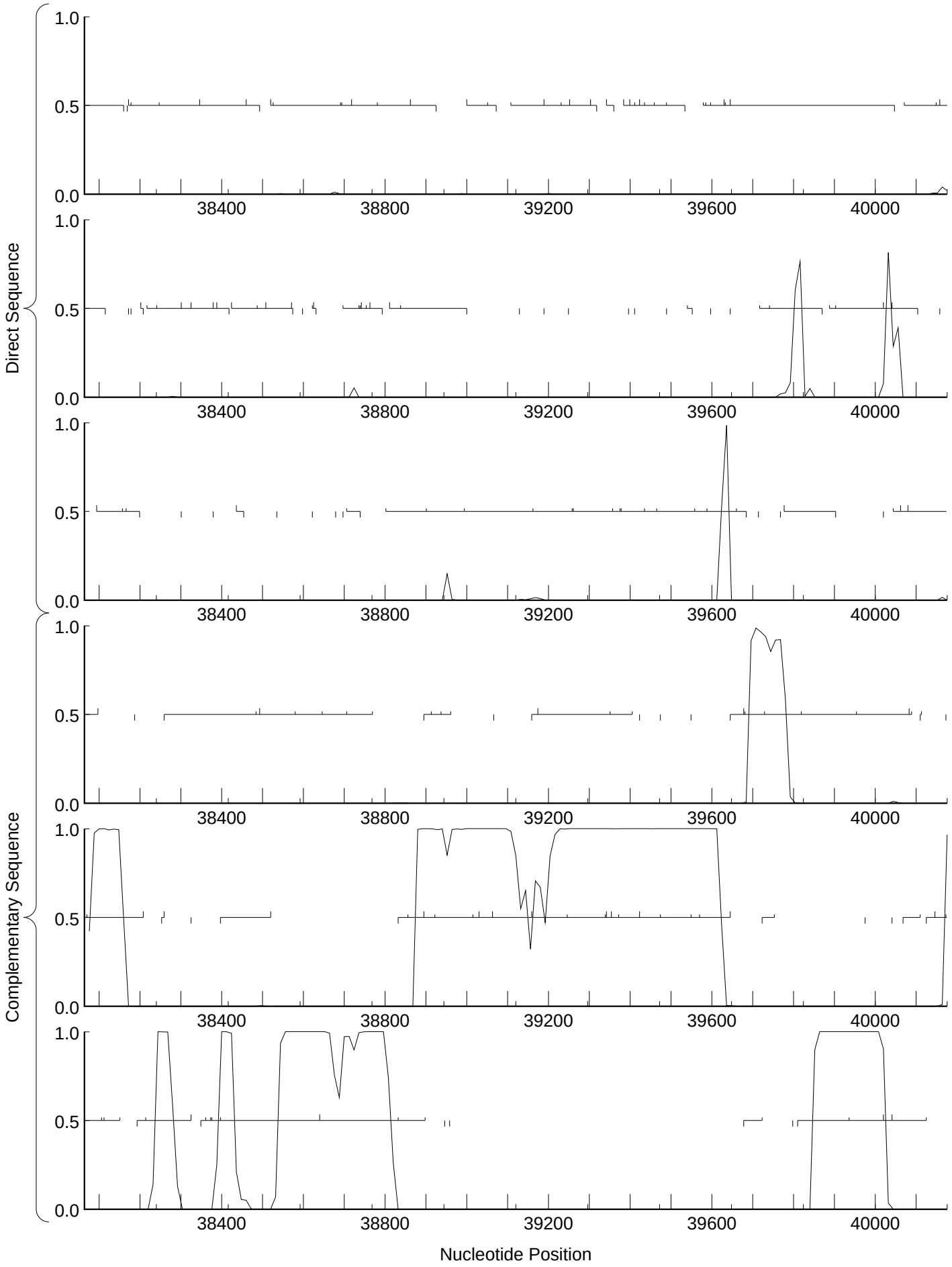


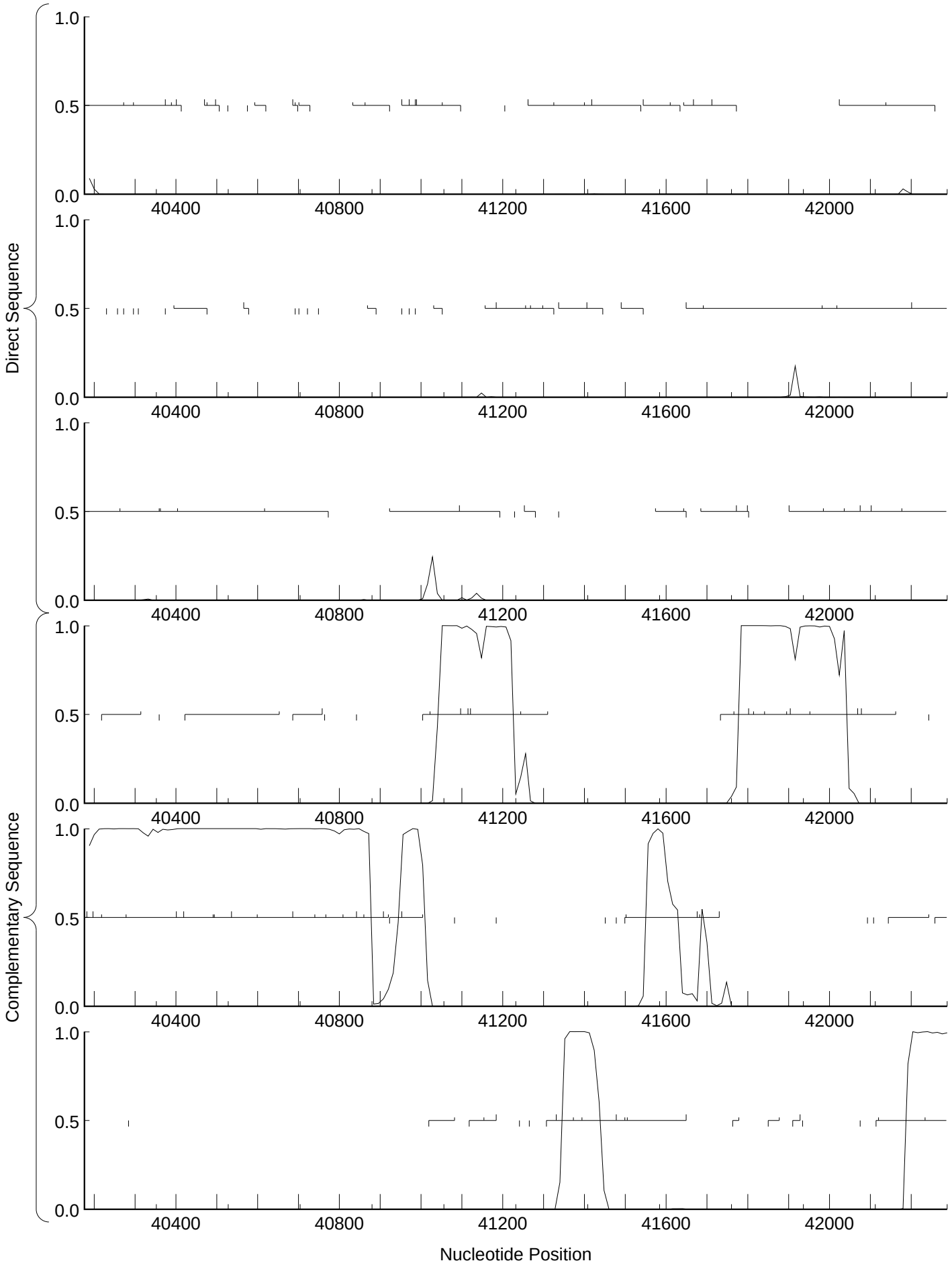

Direct Sequence
Complementary Sequence
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34000
34400
34800
35200
35600
Nucleotide Position

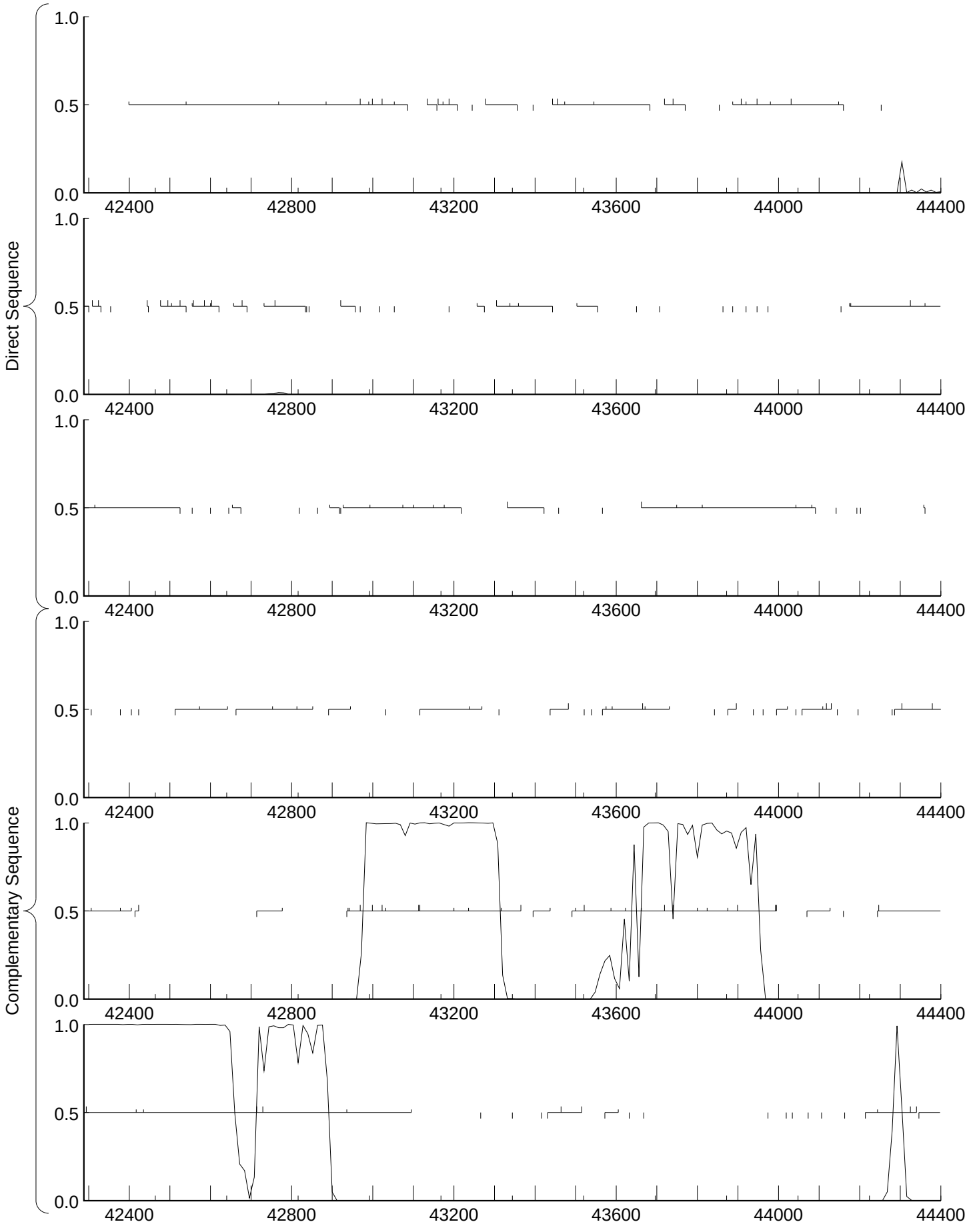
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 19/26



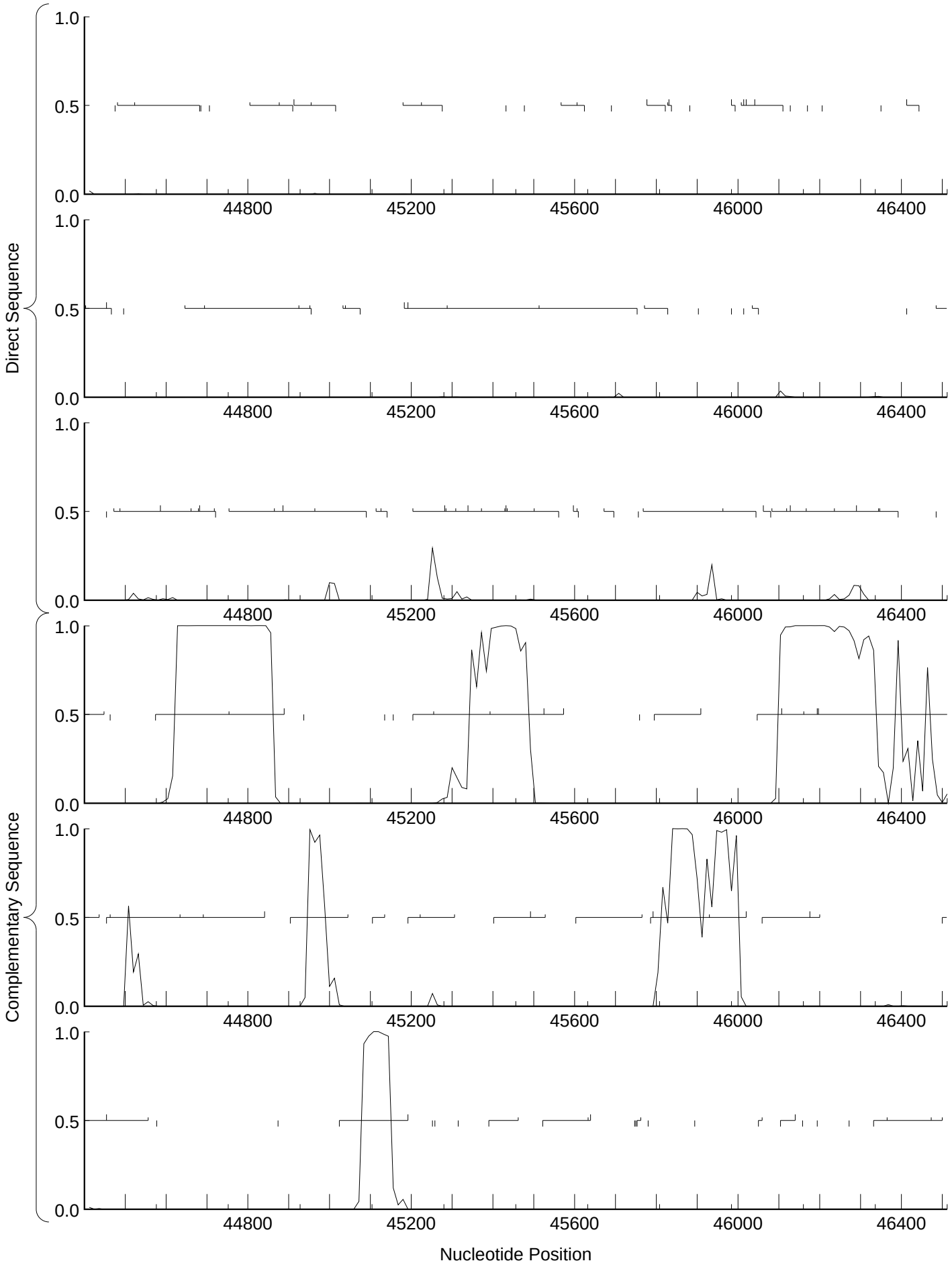
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 20/26



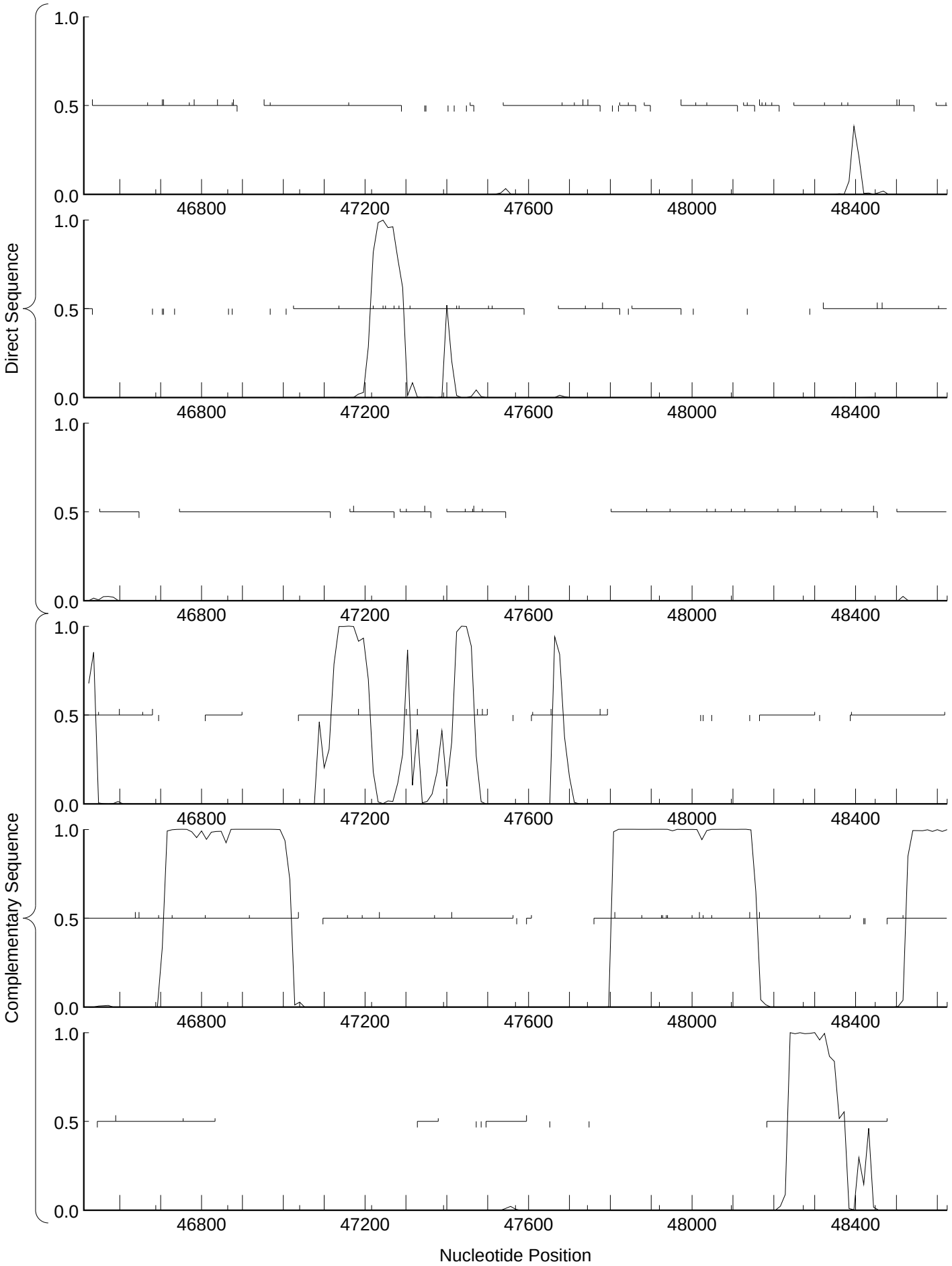

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
40400
40800
41200
41600
42000
Nucleotide Position


Direct Sequence
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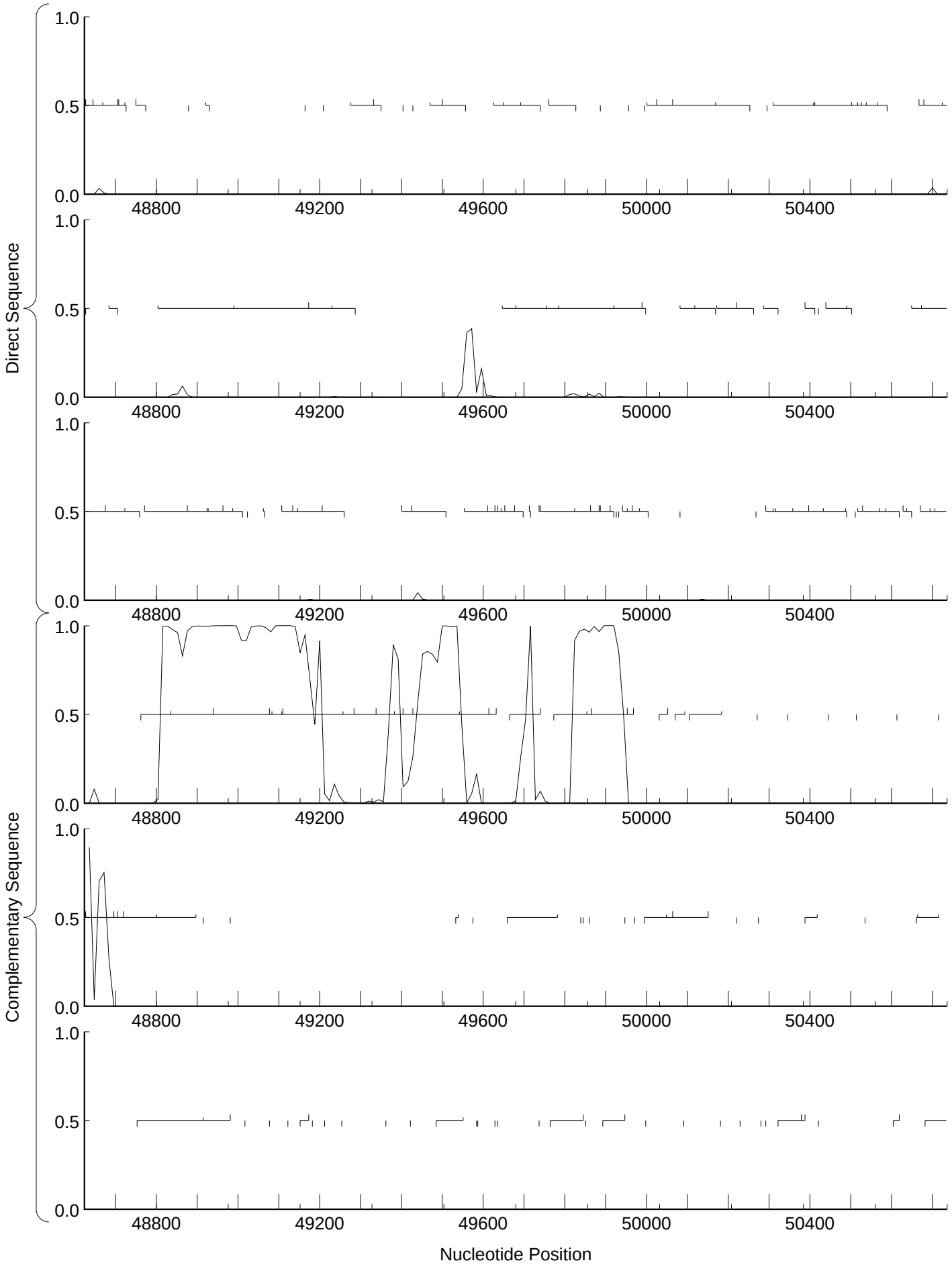
complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 23/26



complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 24/26



complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 25/26



complete sequence, 51094 bp including 10-base 3’ overhang (CGGCCGGTAA), Cluster A10, Order 4, Window 96, Step 12, 26/26

