GeneMark

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

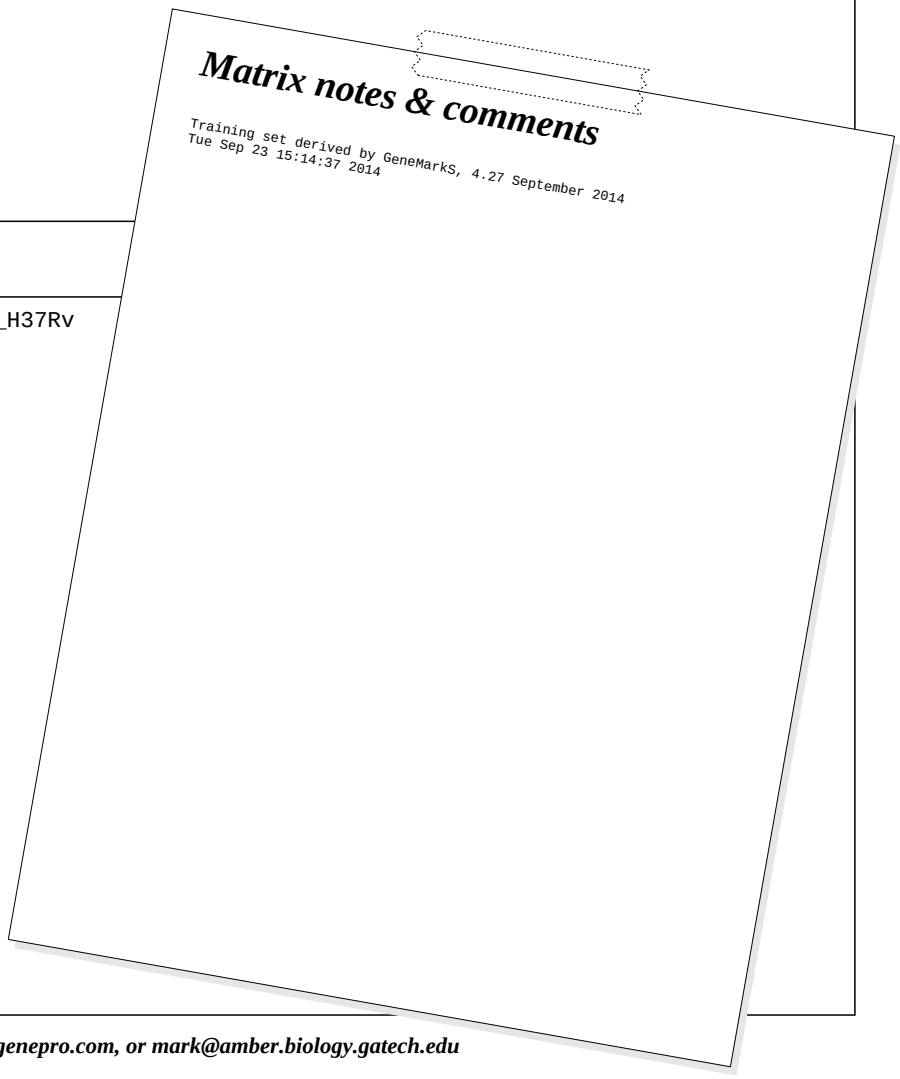
PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Terror complete sequence, 41077 bp including 11-base 3' overhang (CCCCA
Analysis Date    :  3/29/24 at 6:35:45
Pages            :  21
Sequence Length  :  41077 bp
GC Content       :   66.43%

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

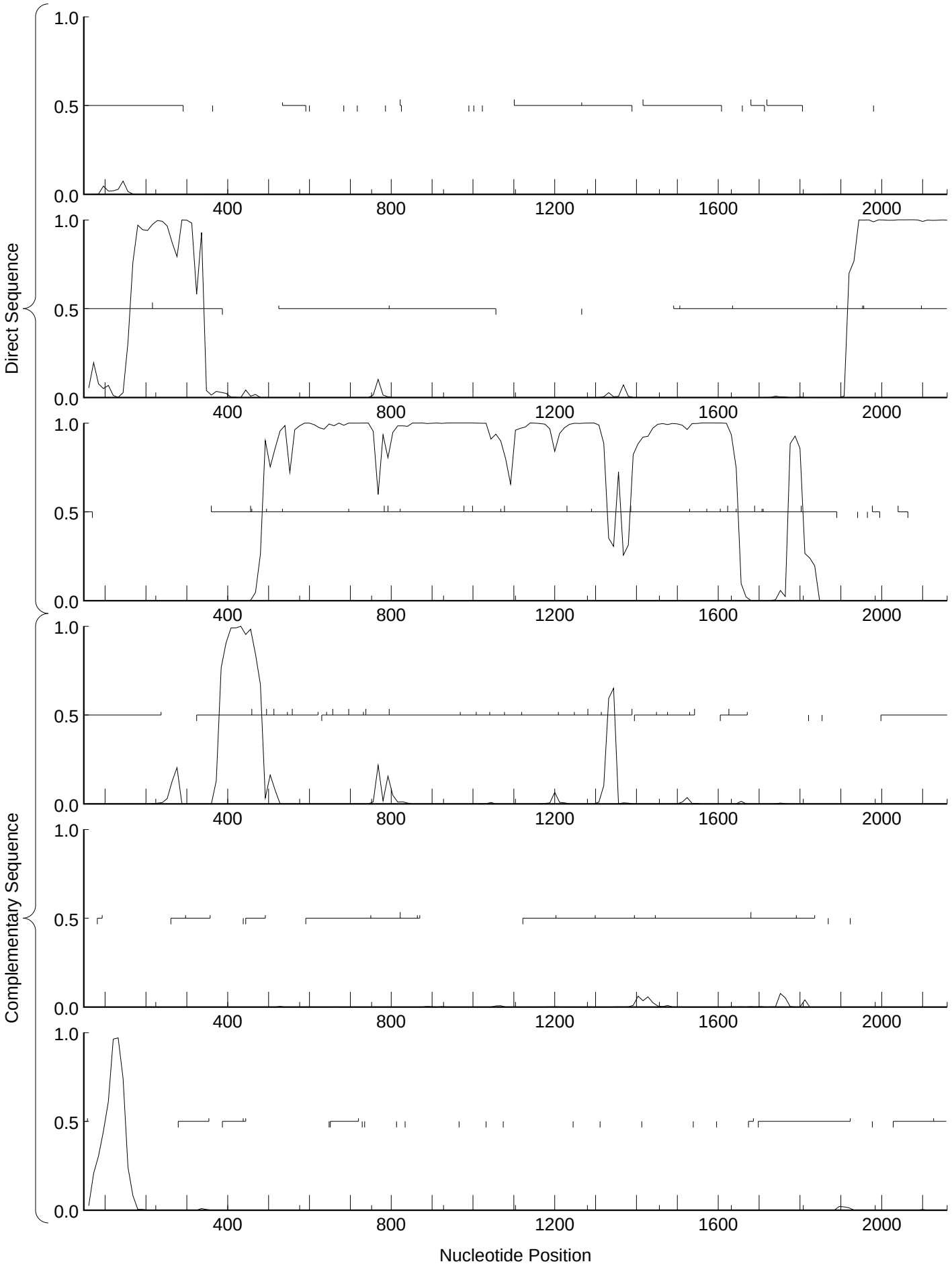
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Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

Order      :  4

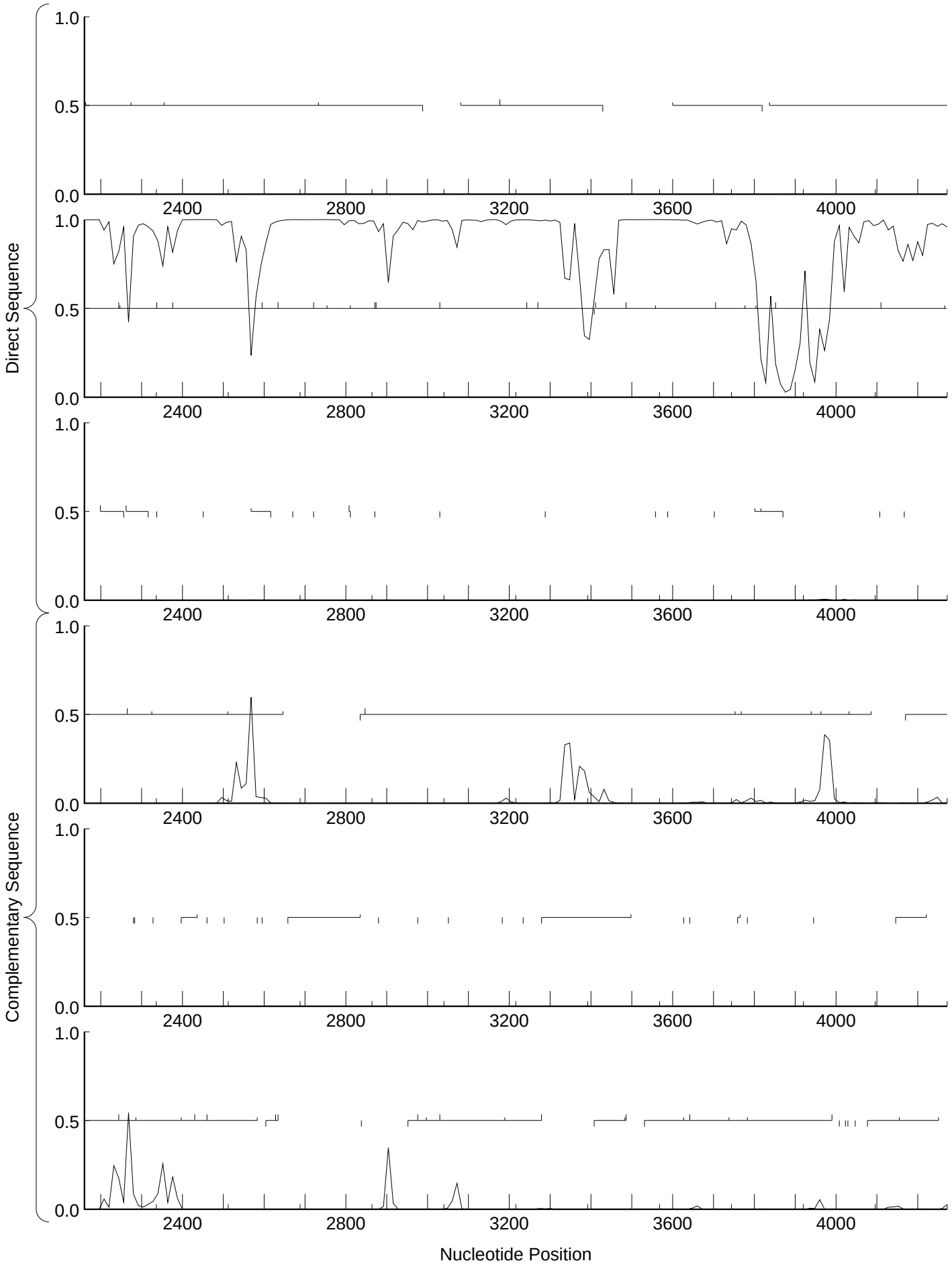

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

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

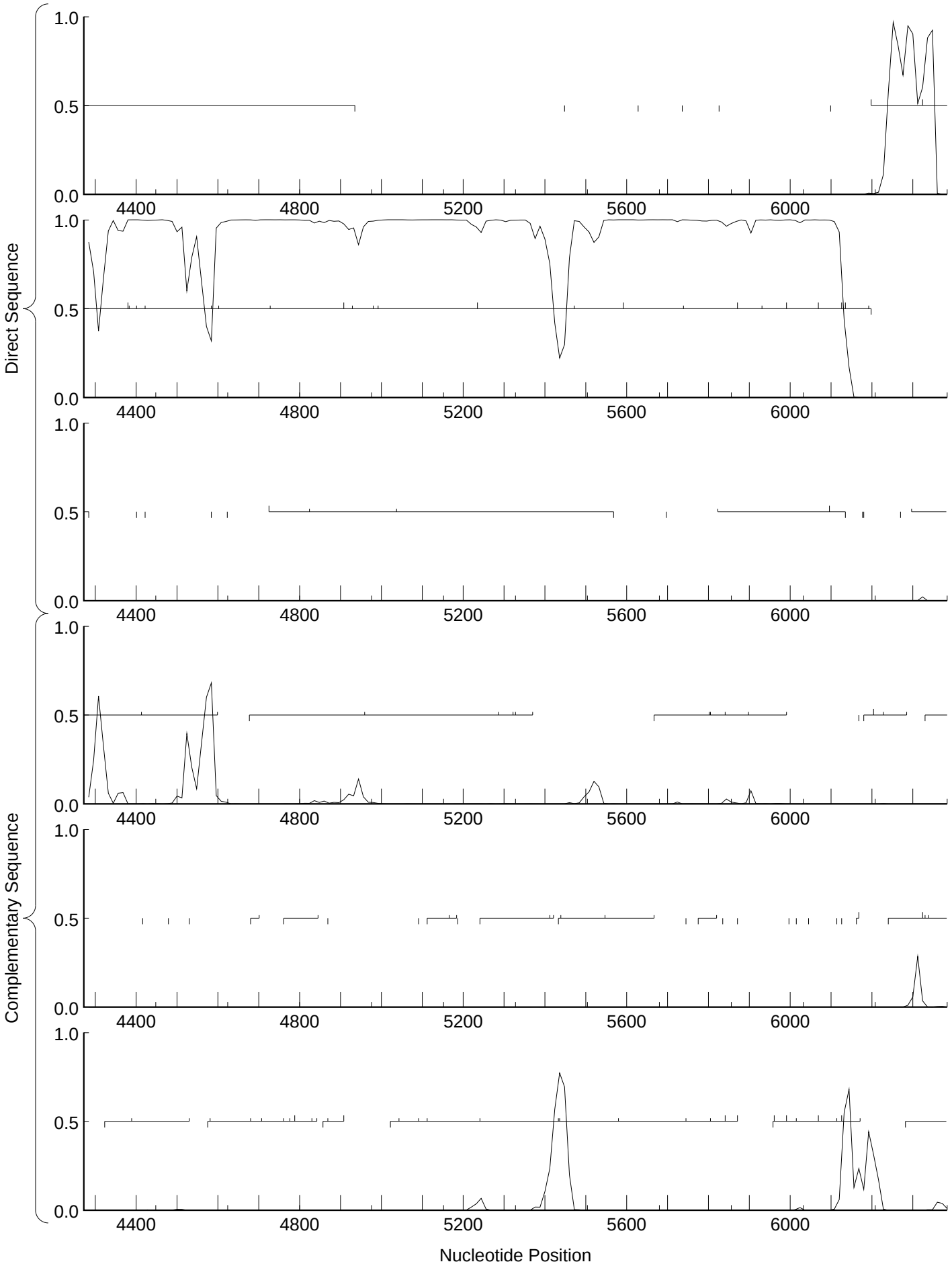
complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 2/21



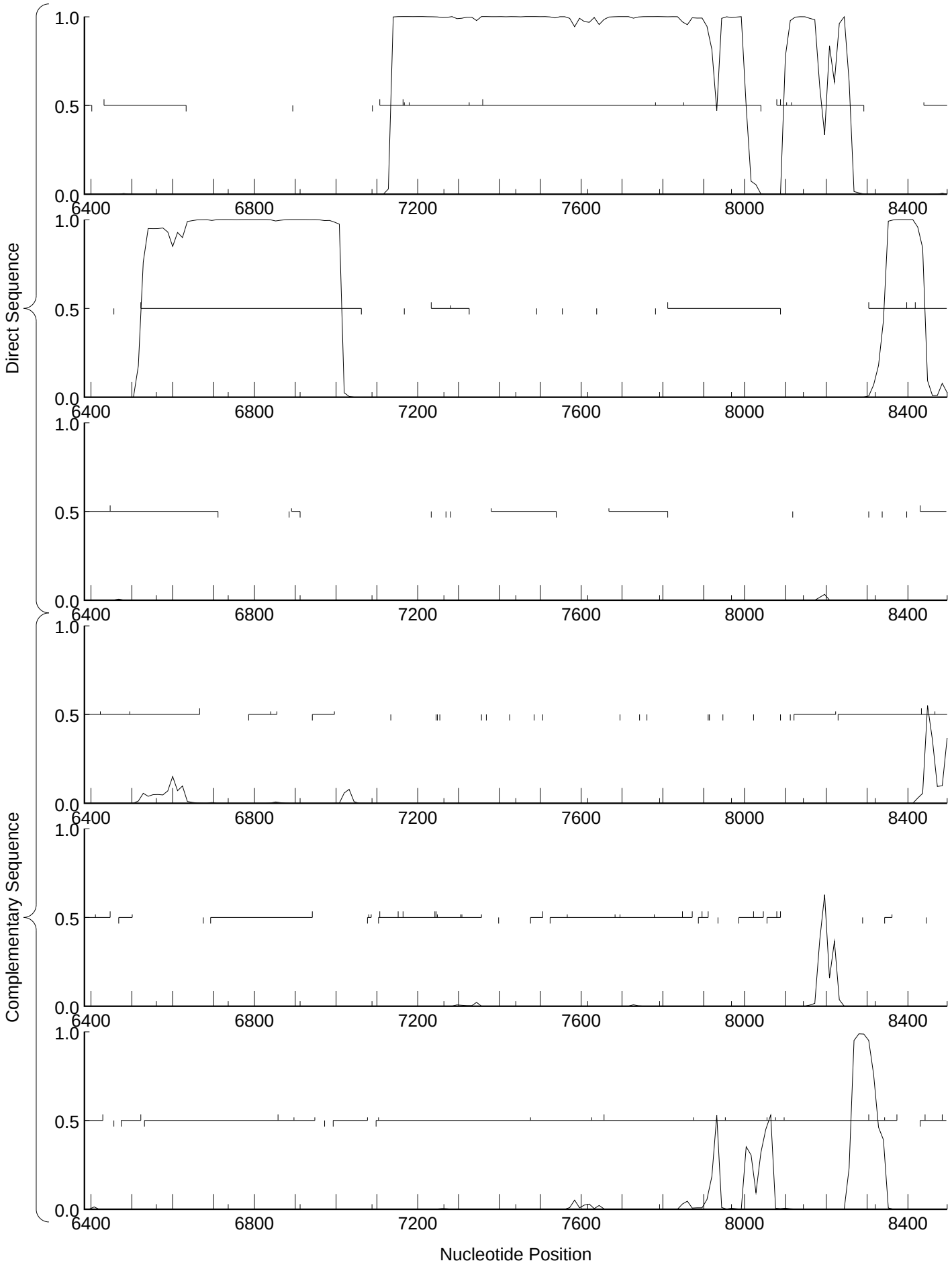
complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 3/21



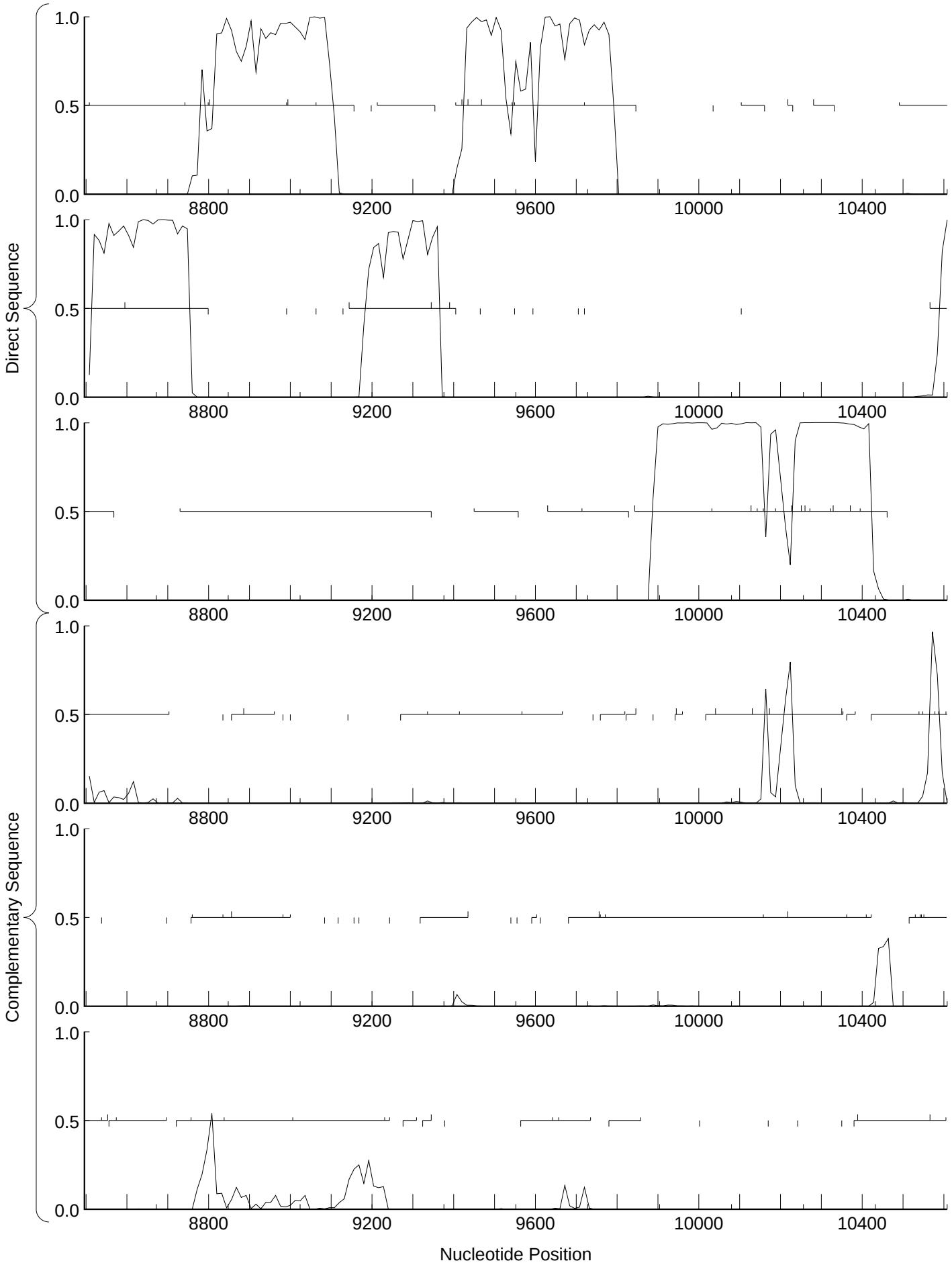
complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 4/21



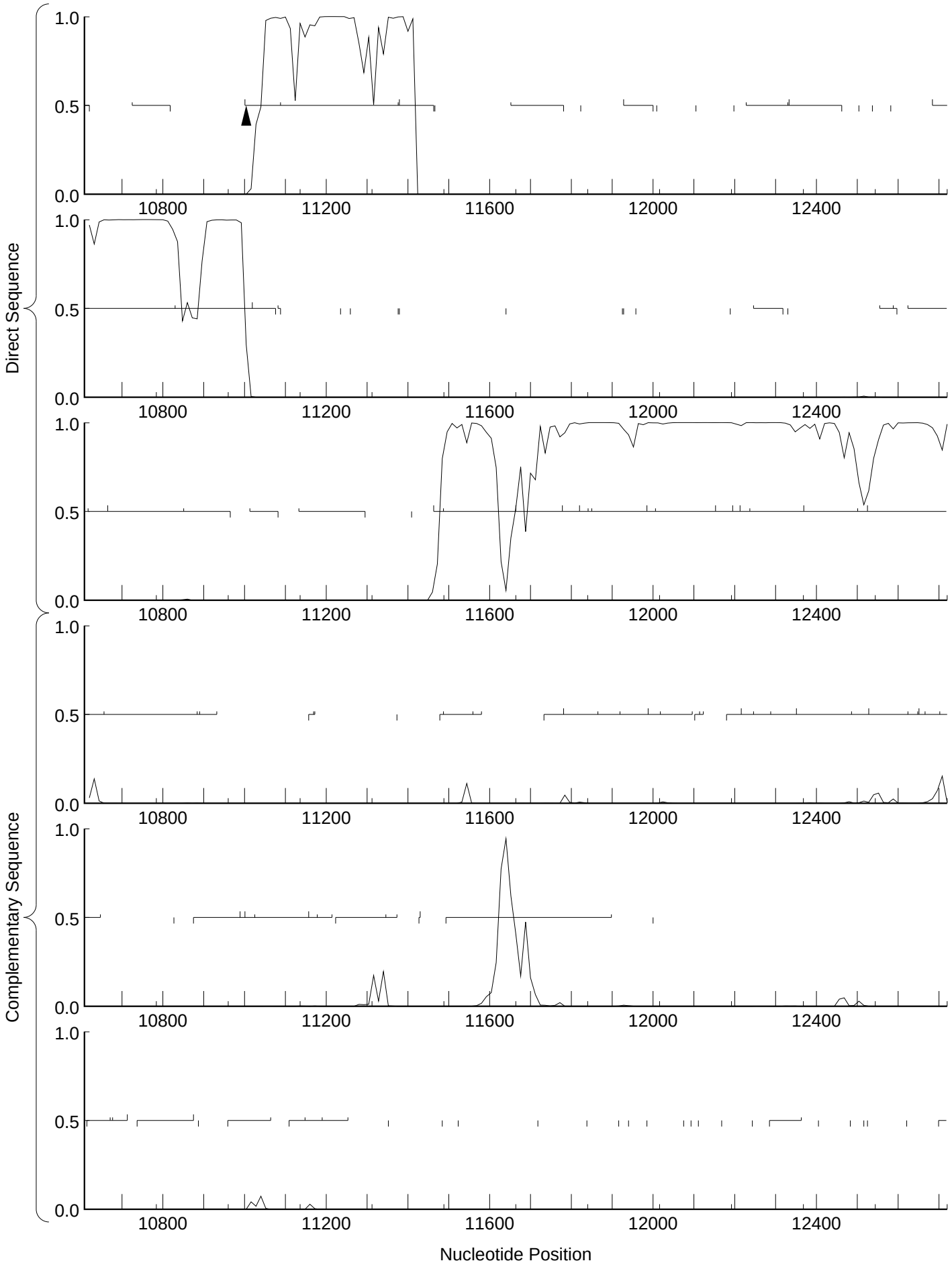
complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 5/21



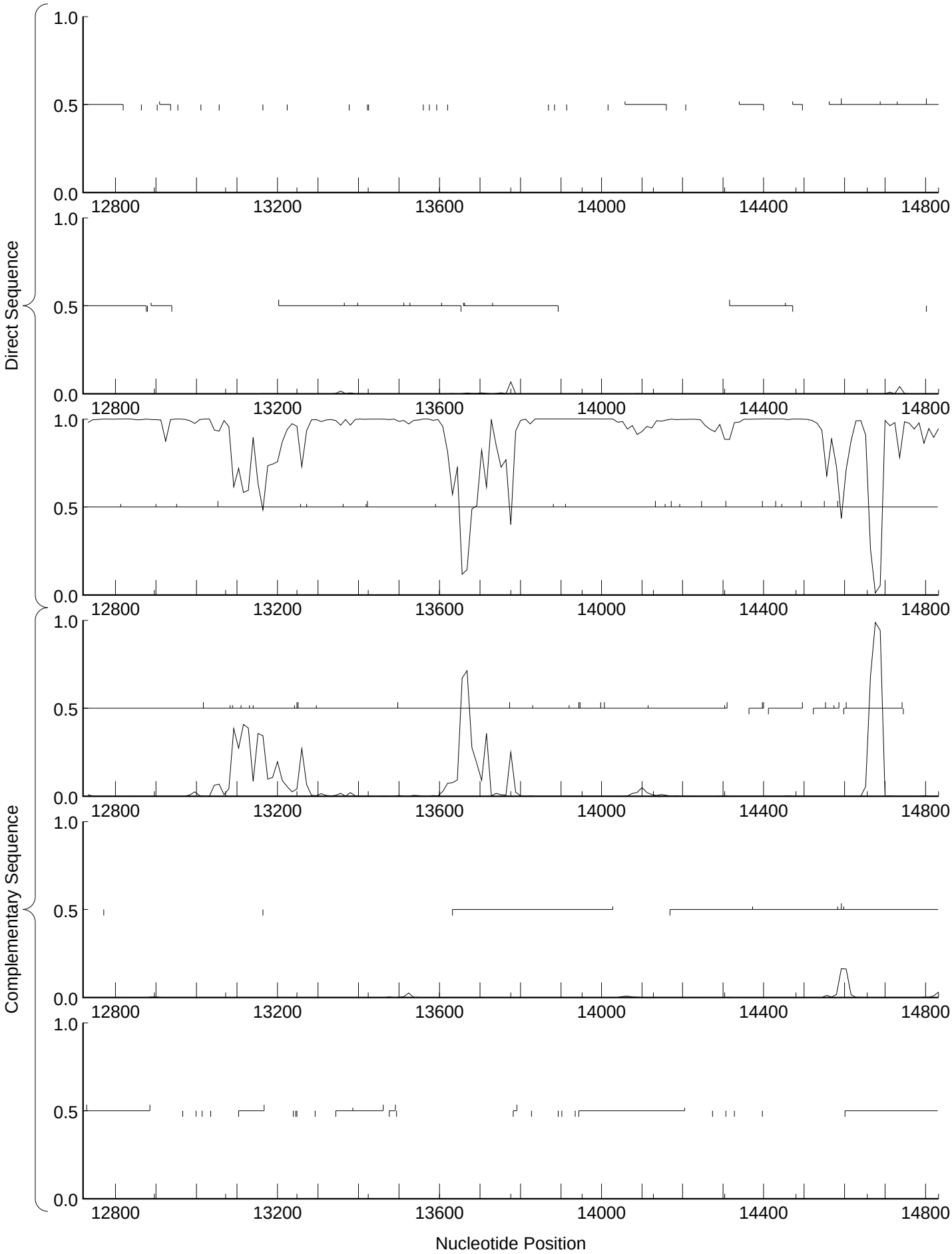
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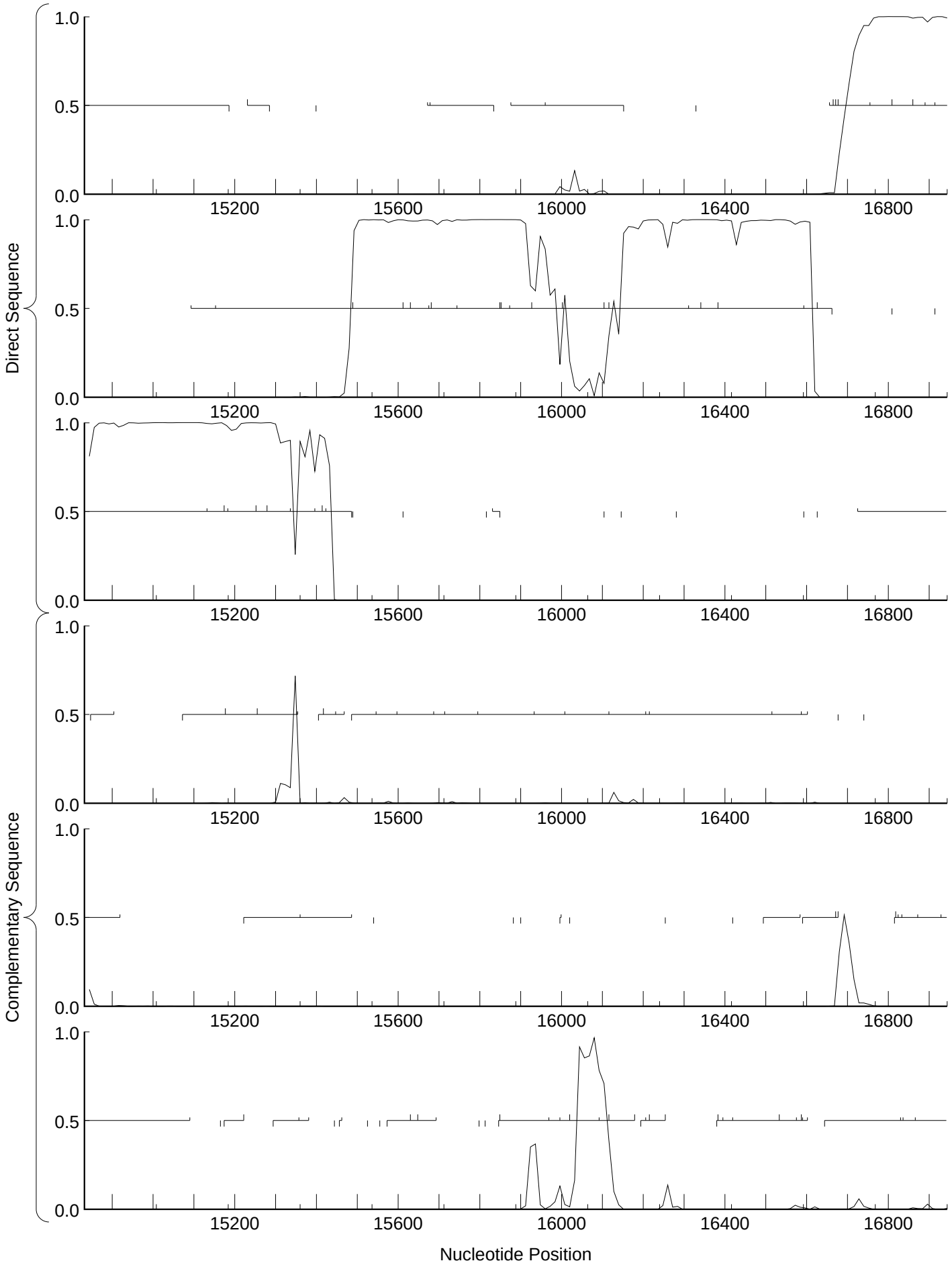
complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 7/21



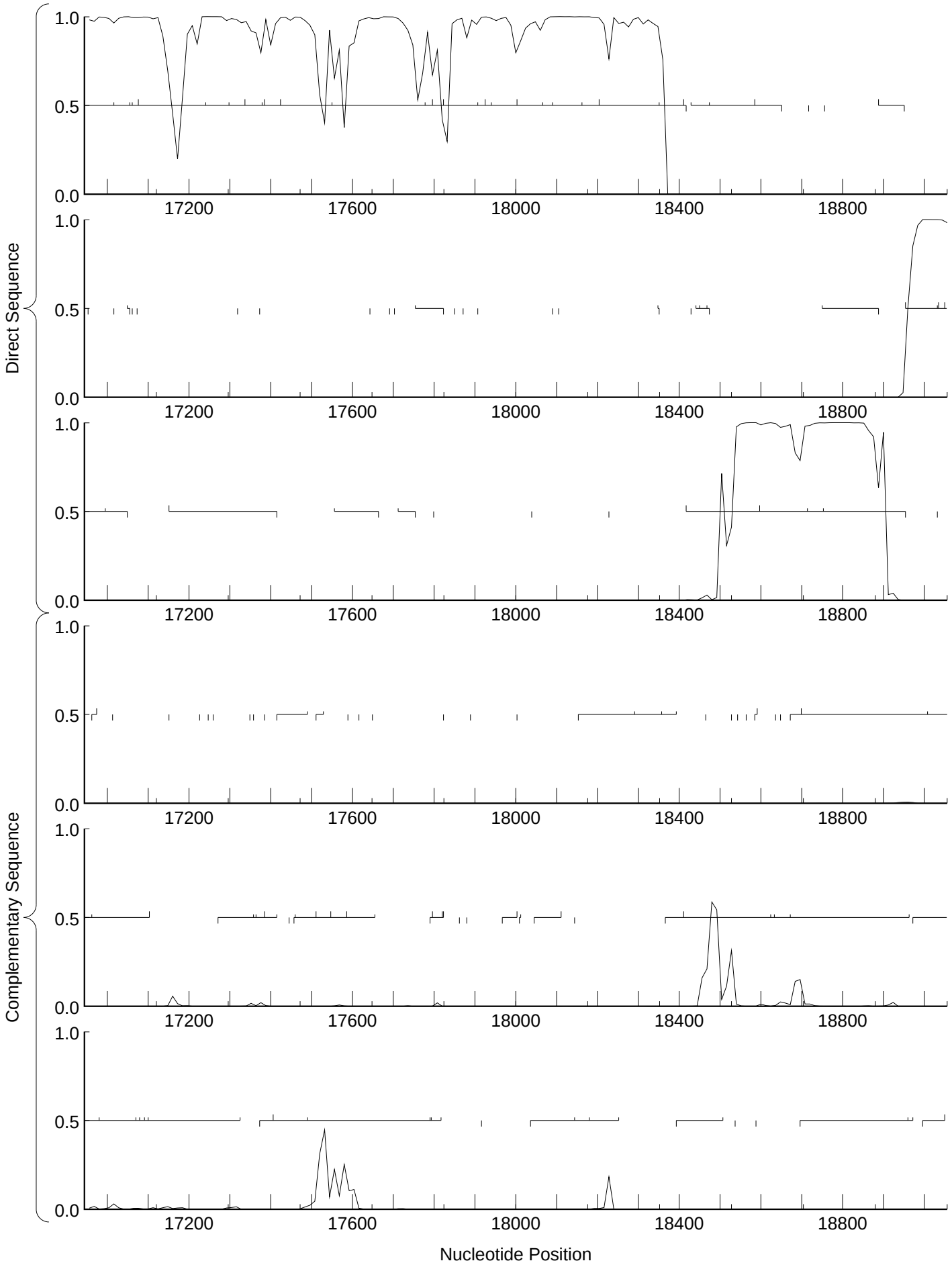
complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 8/21



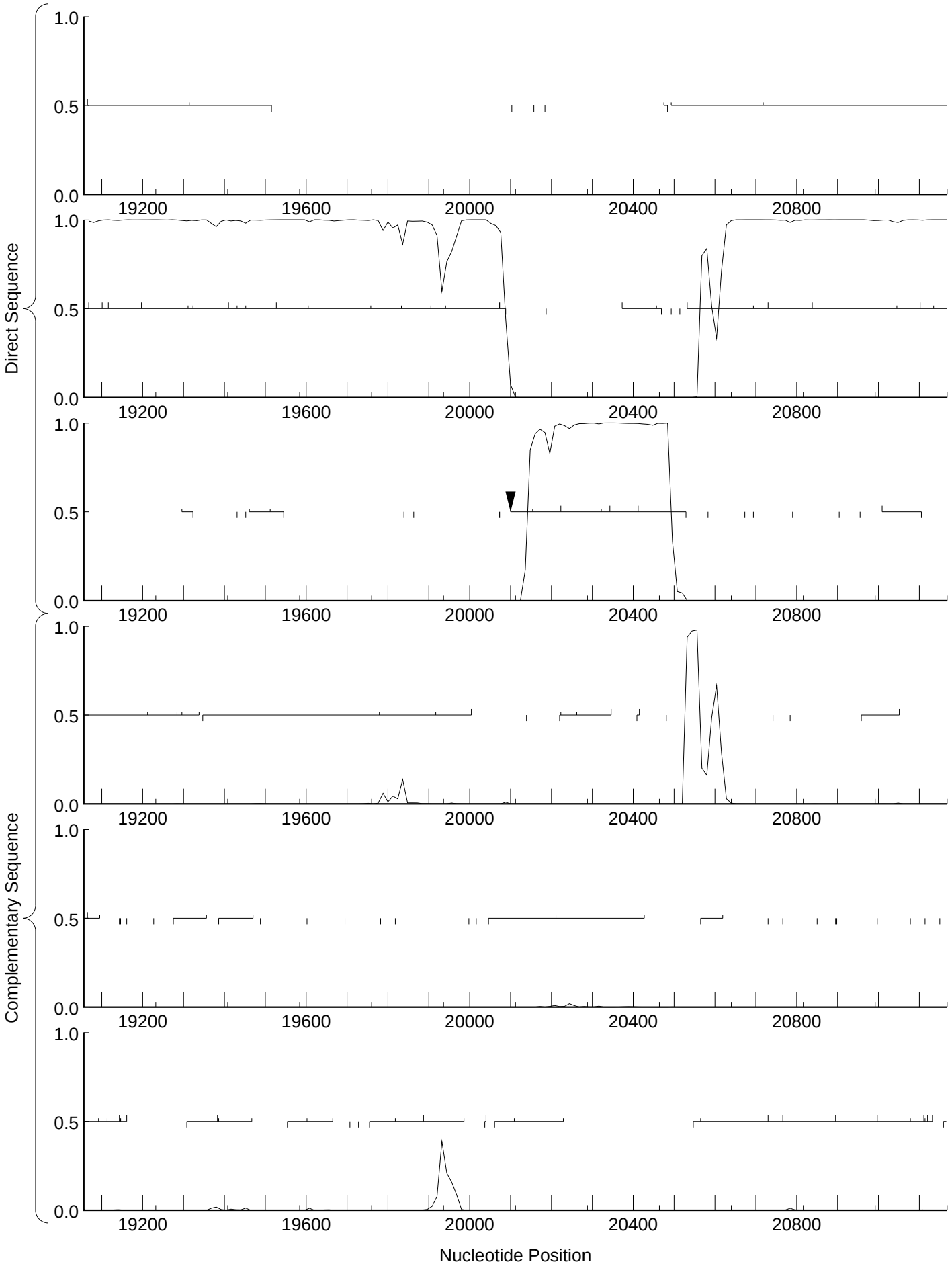
complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 9/21



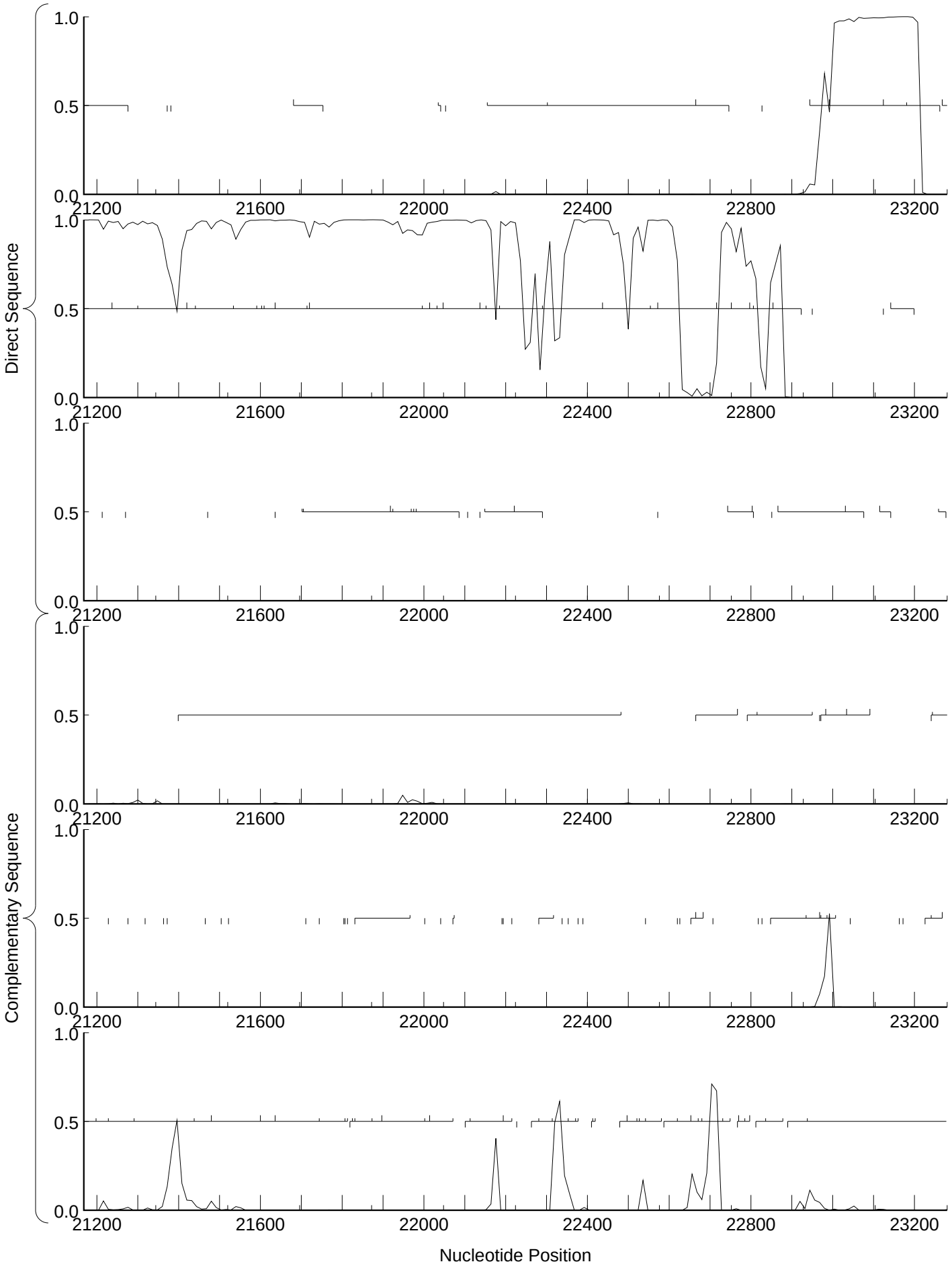
complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 10/21



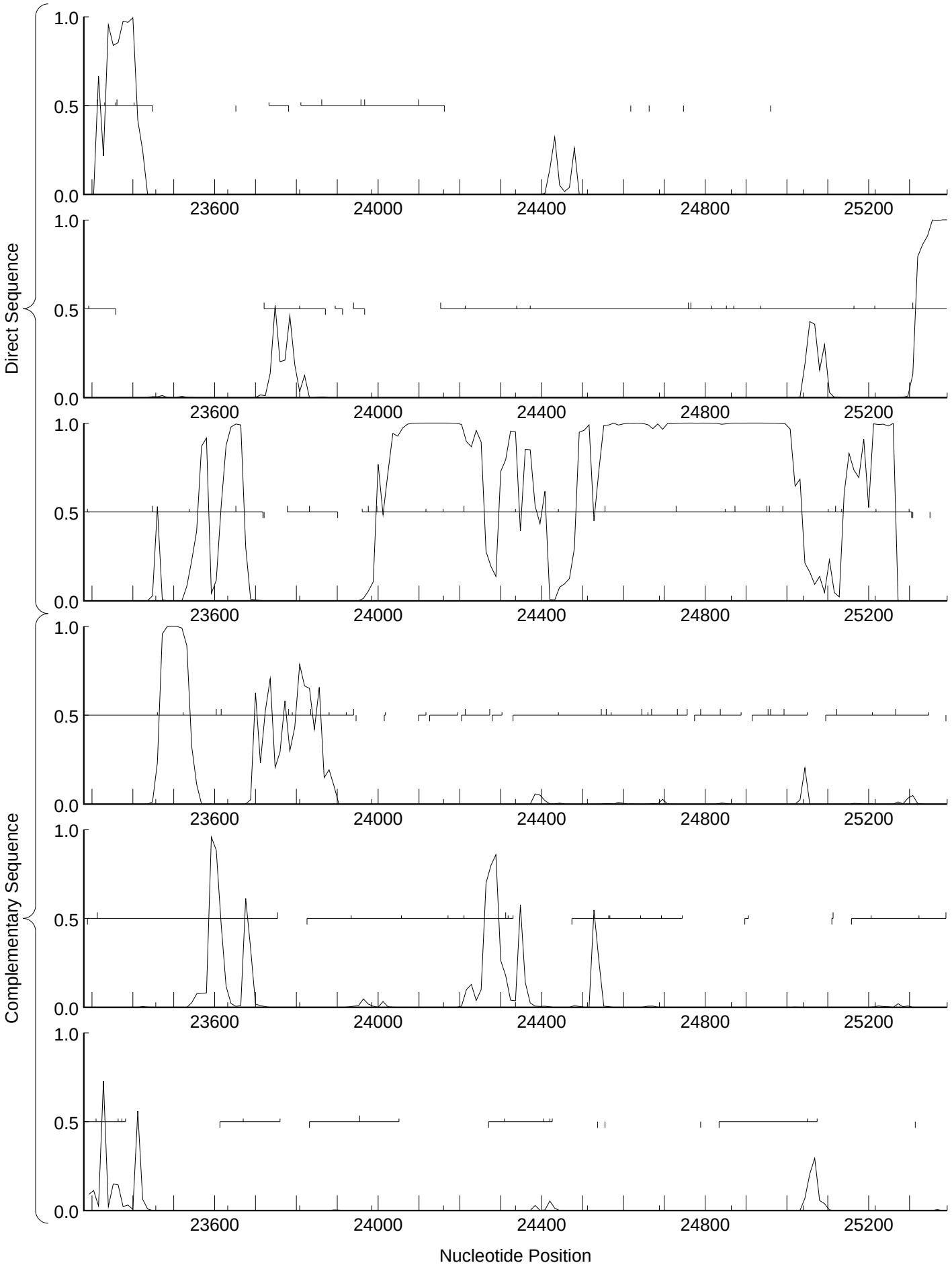
complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 11/21

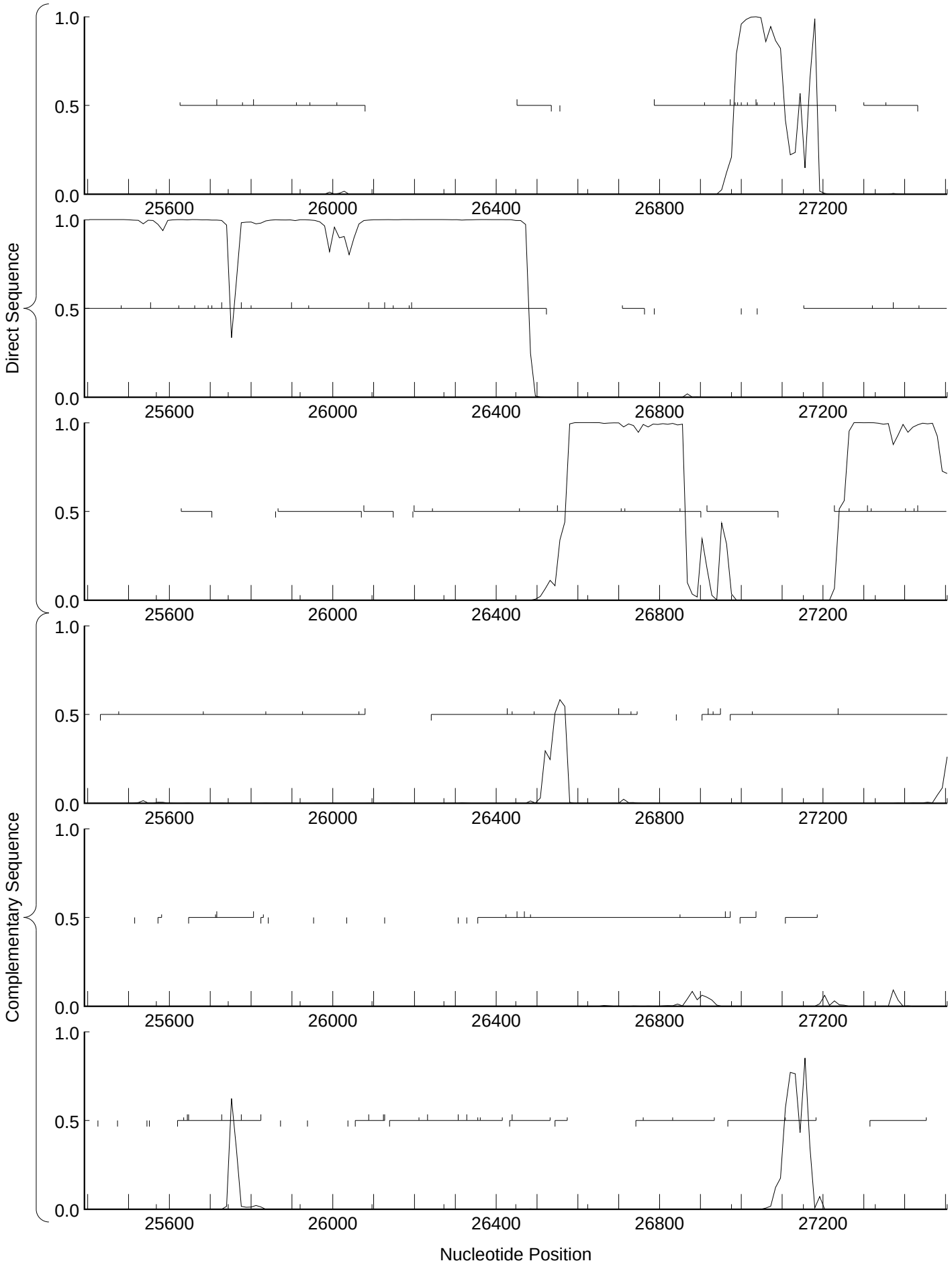


complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 12/21

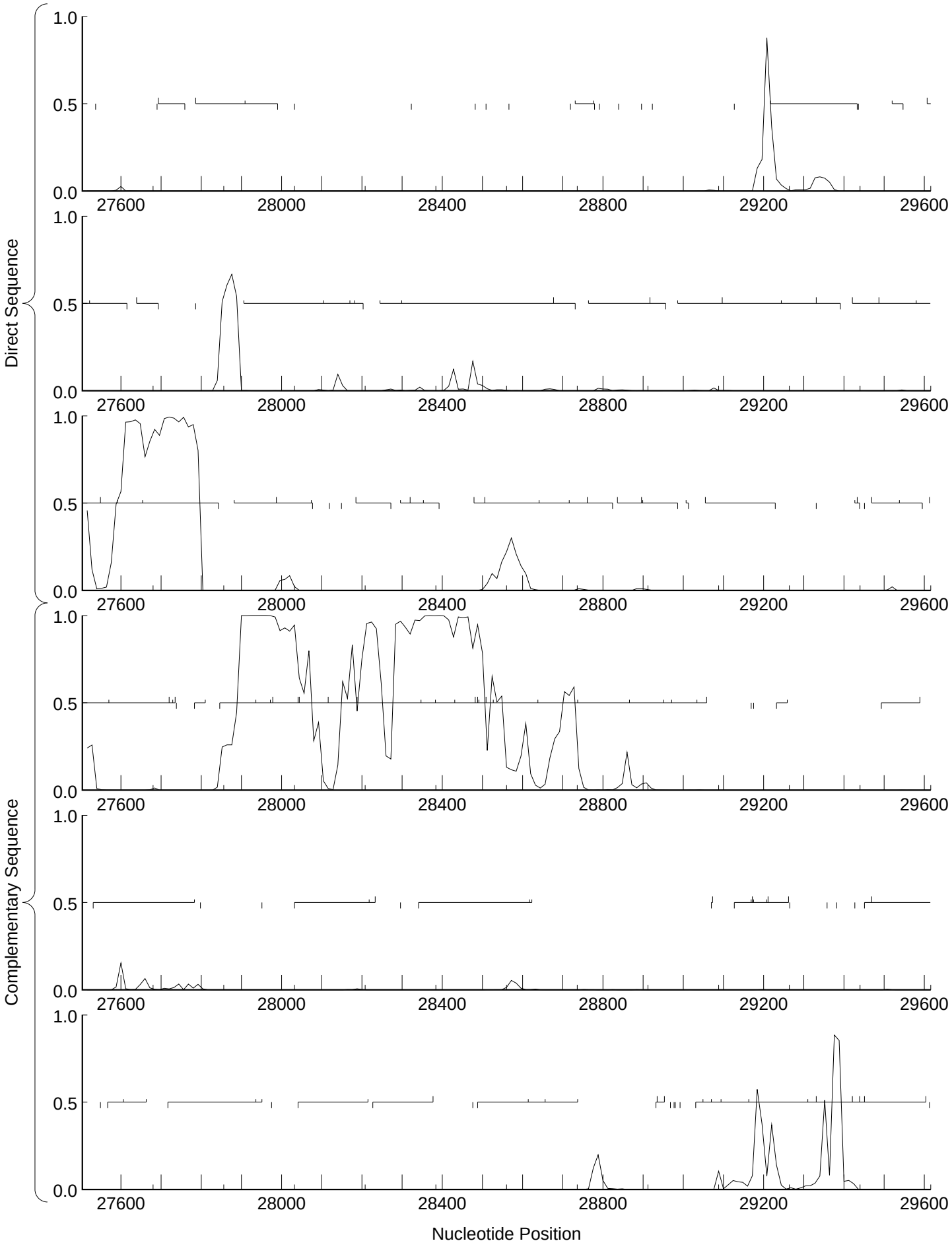


complete sequence, 41077 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 13/21

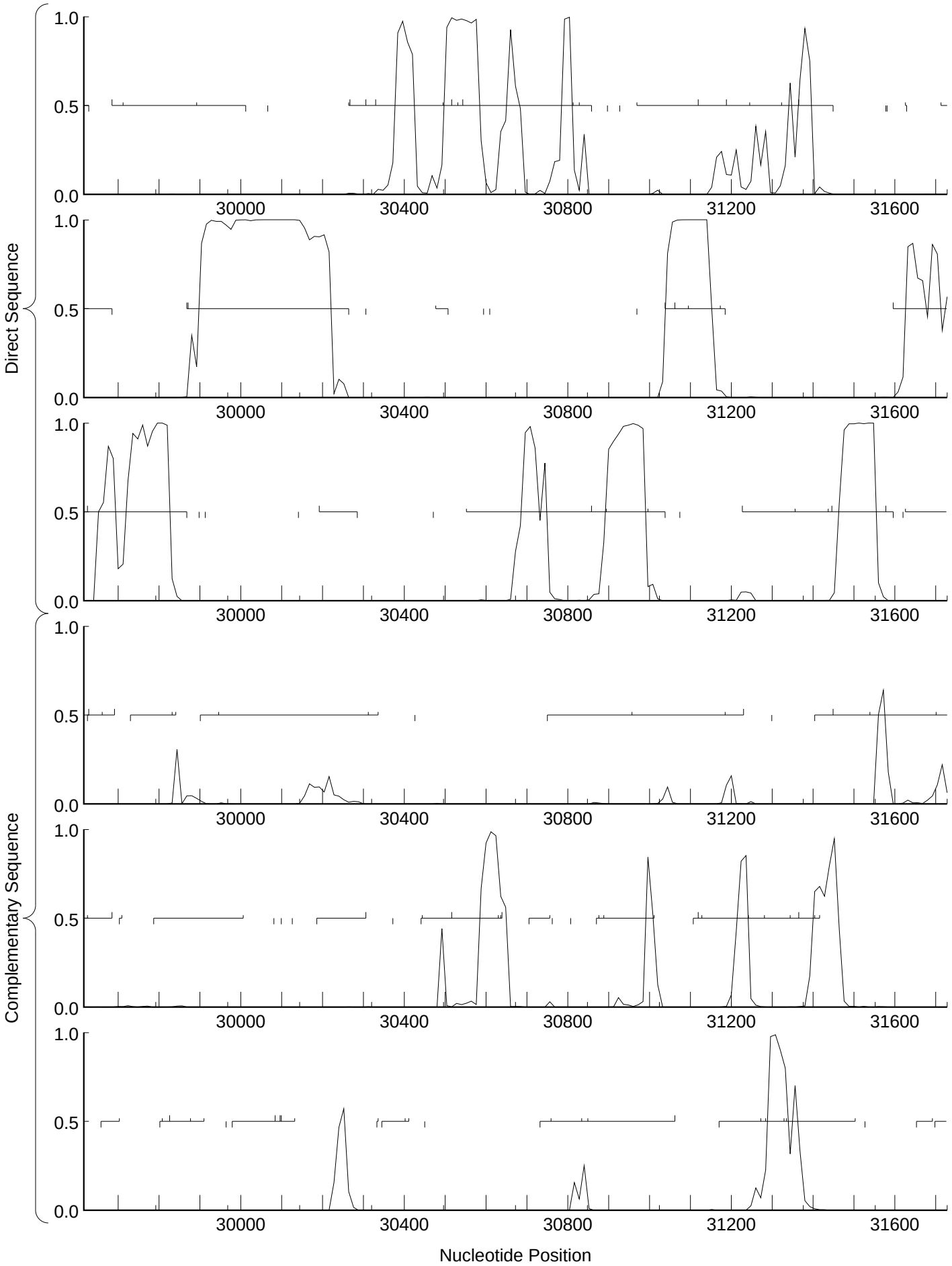



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
25600
26000
26400
26800
27200
Nucleotide Position

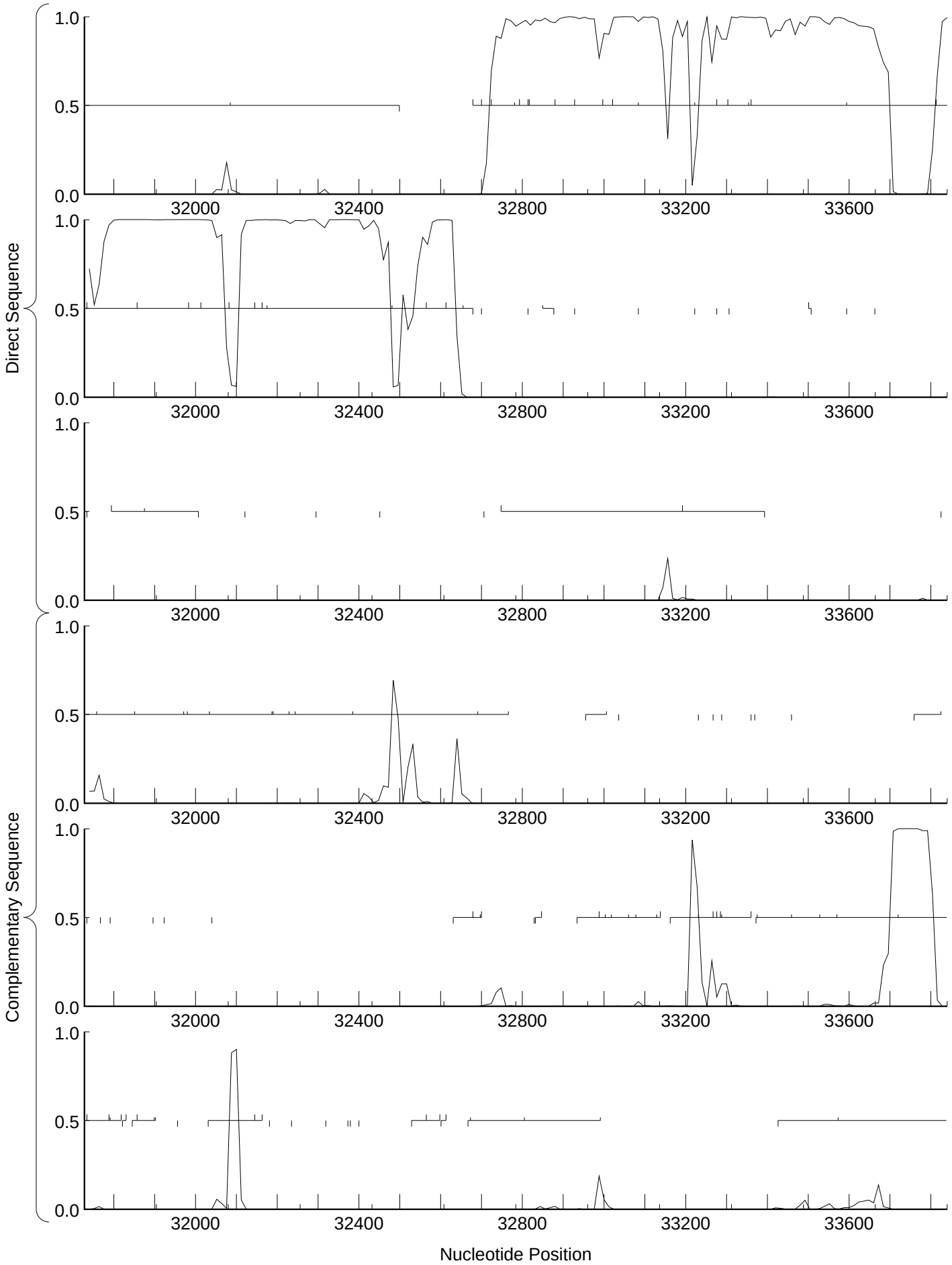
complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 15/21



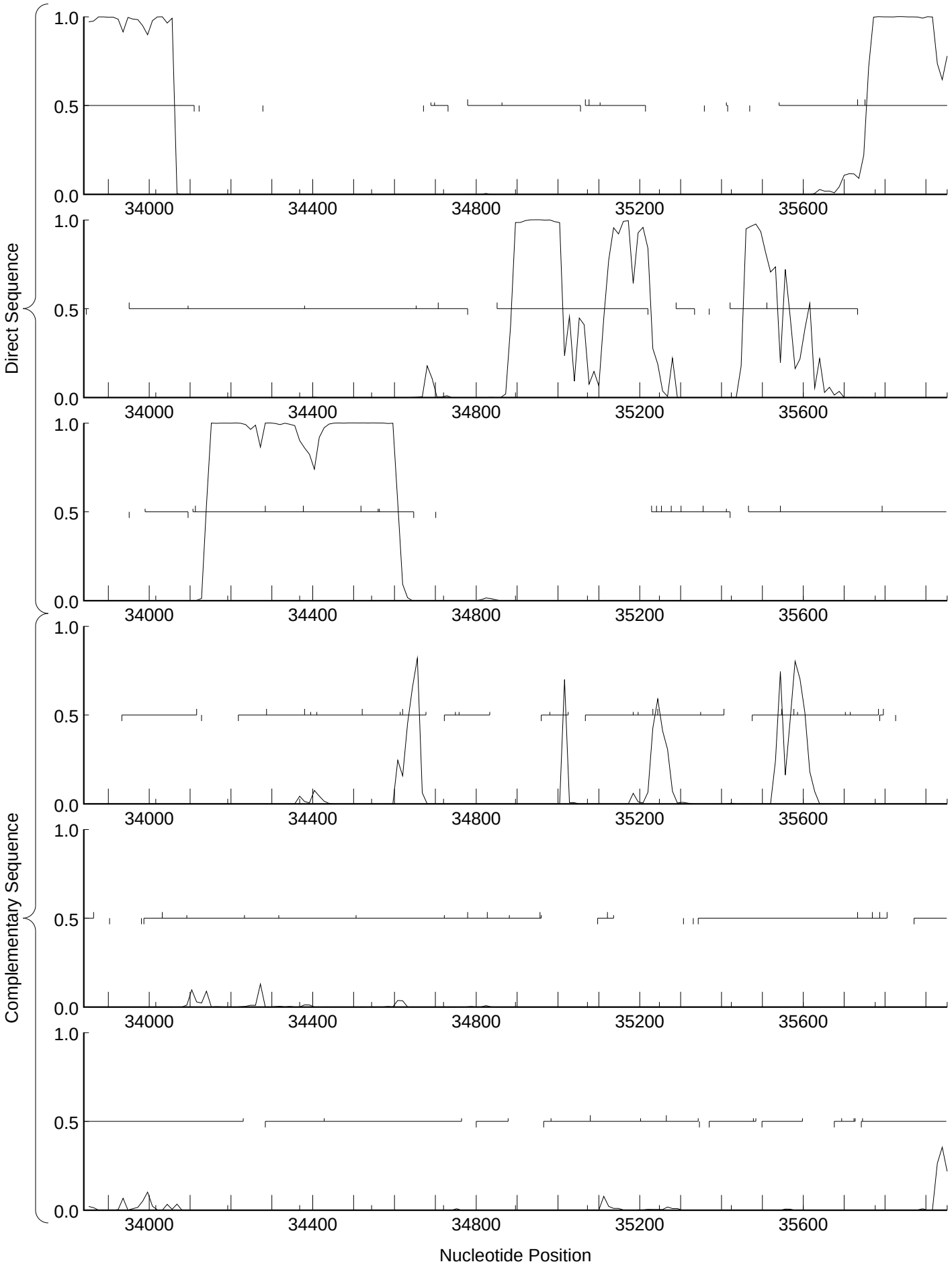
complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 16/21



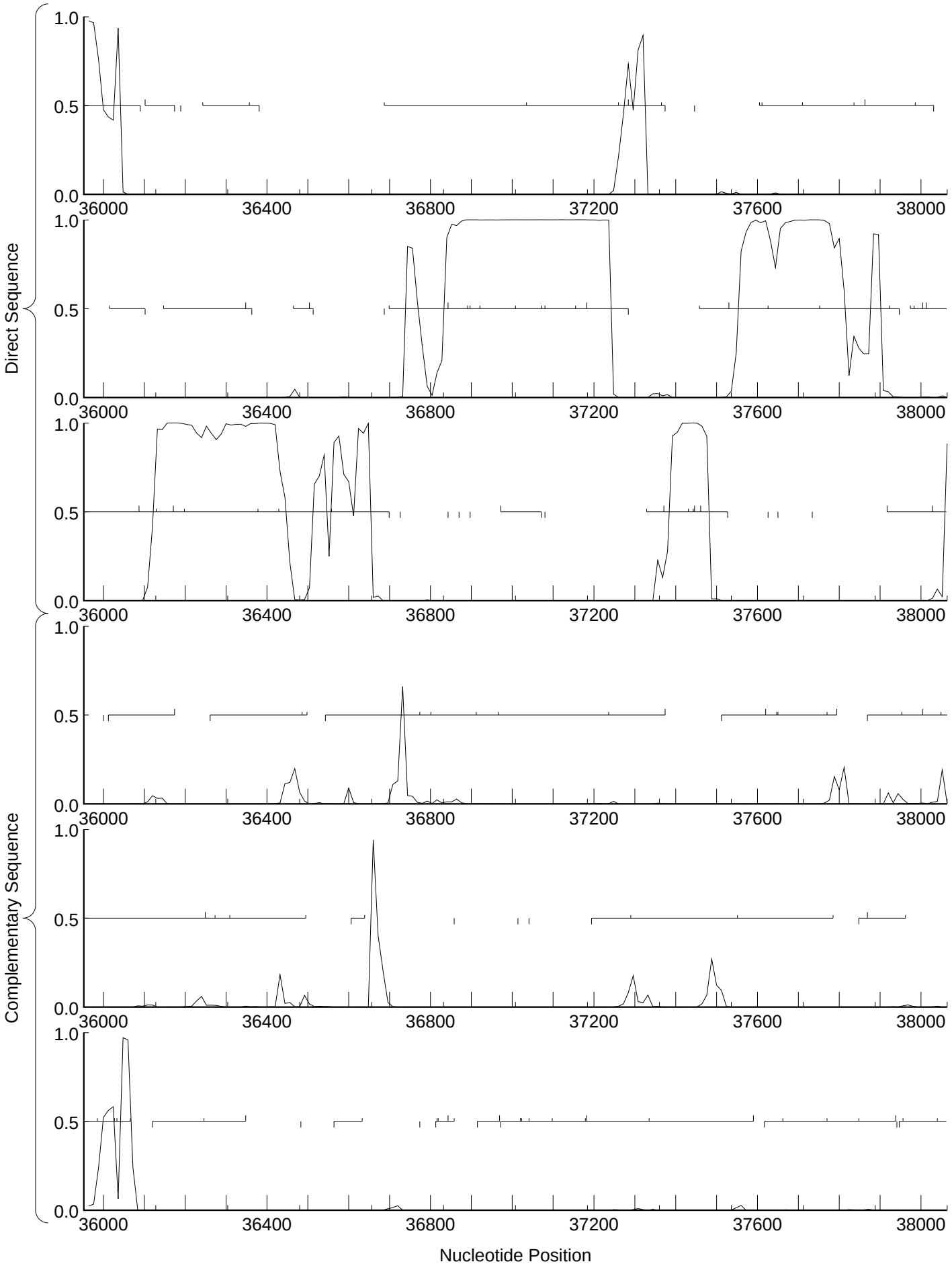
complete sequence, 41077 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 17/21



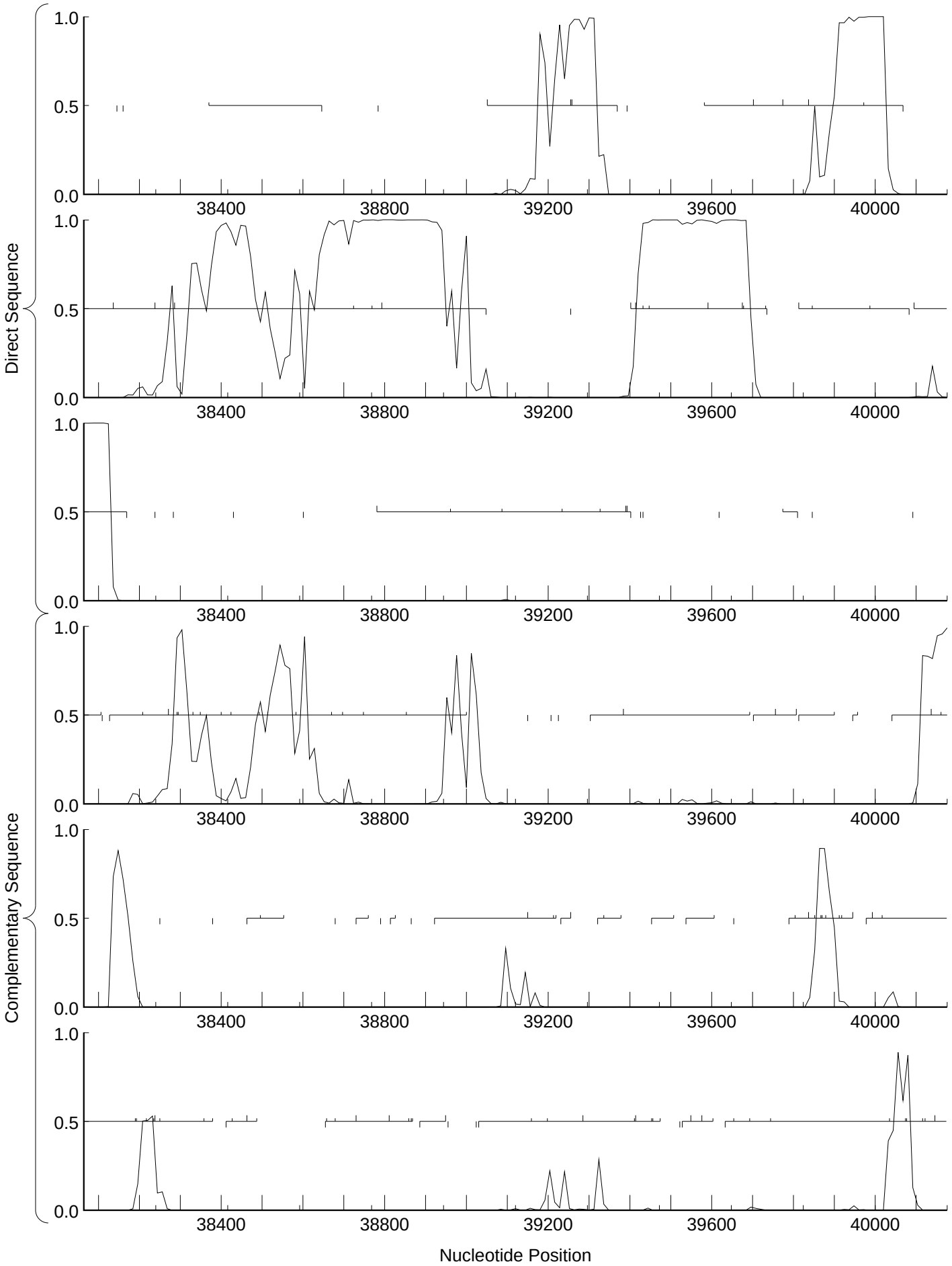
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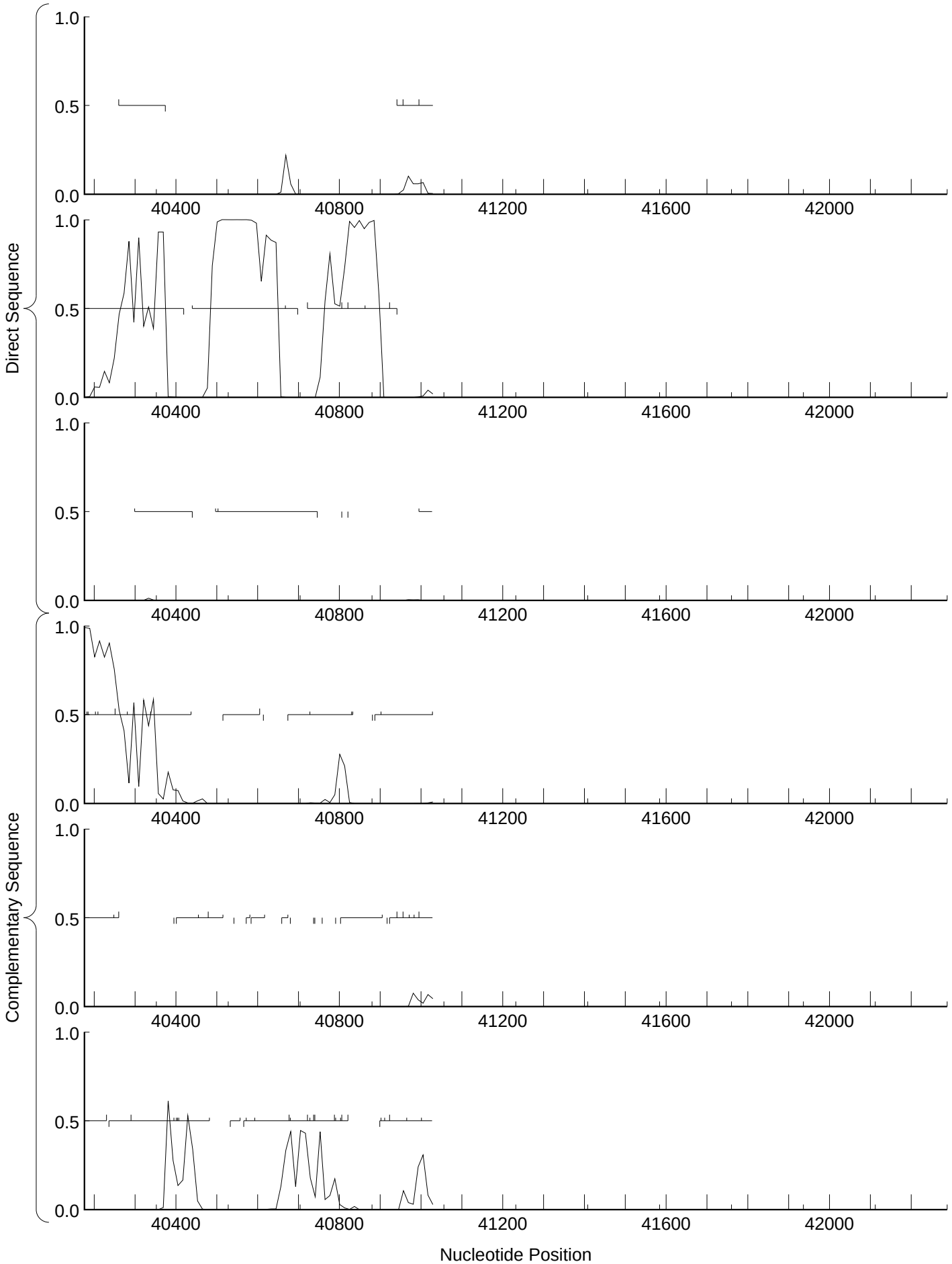


complete sequence, 41077 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 19/21



complete sequence, 41077 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 20/21




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