GeneMark

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

PROGRAM INFORMATION

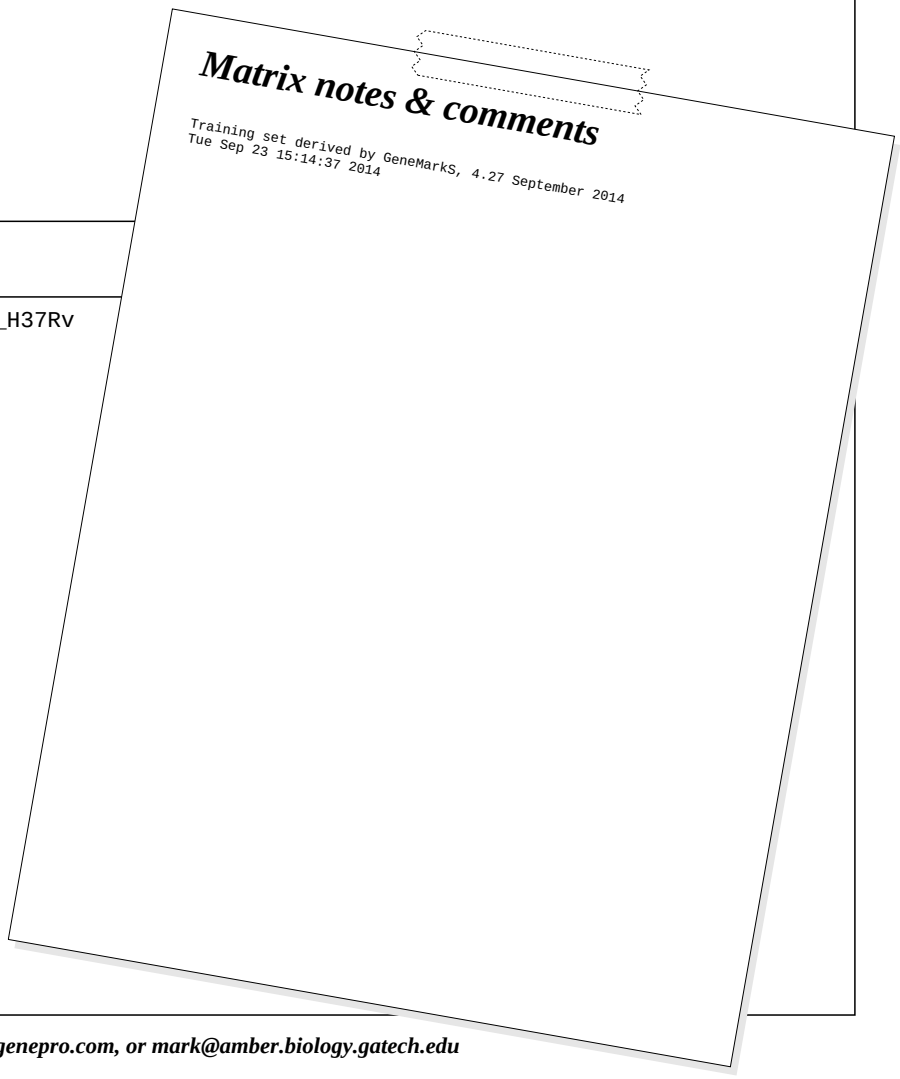
```
Sequence         :  Mycobacterium phage Annihilator complete sequence, 41893 bp including 11-base 3’ overhang (

Analysis Date    :  3/28/24 at 4:40:26
Pages            :  21
Sequence Length  :  41893 bp
GC Content       :   66.60%

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

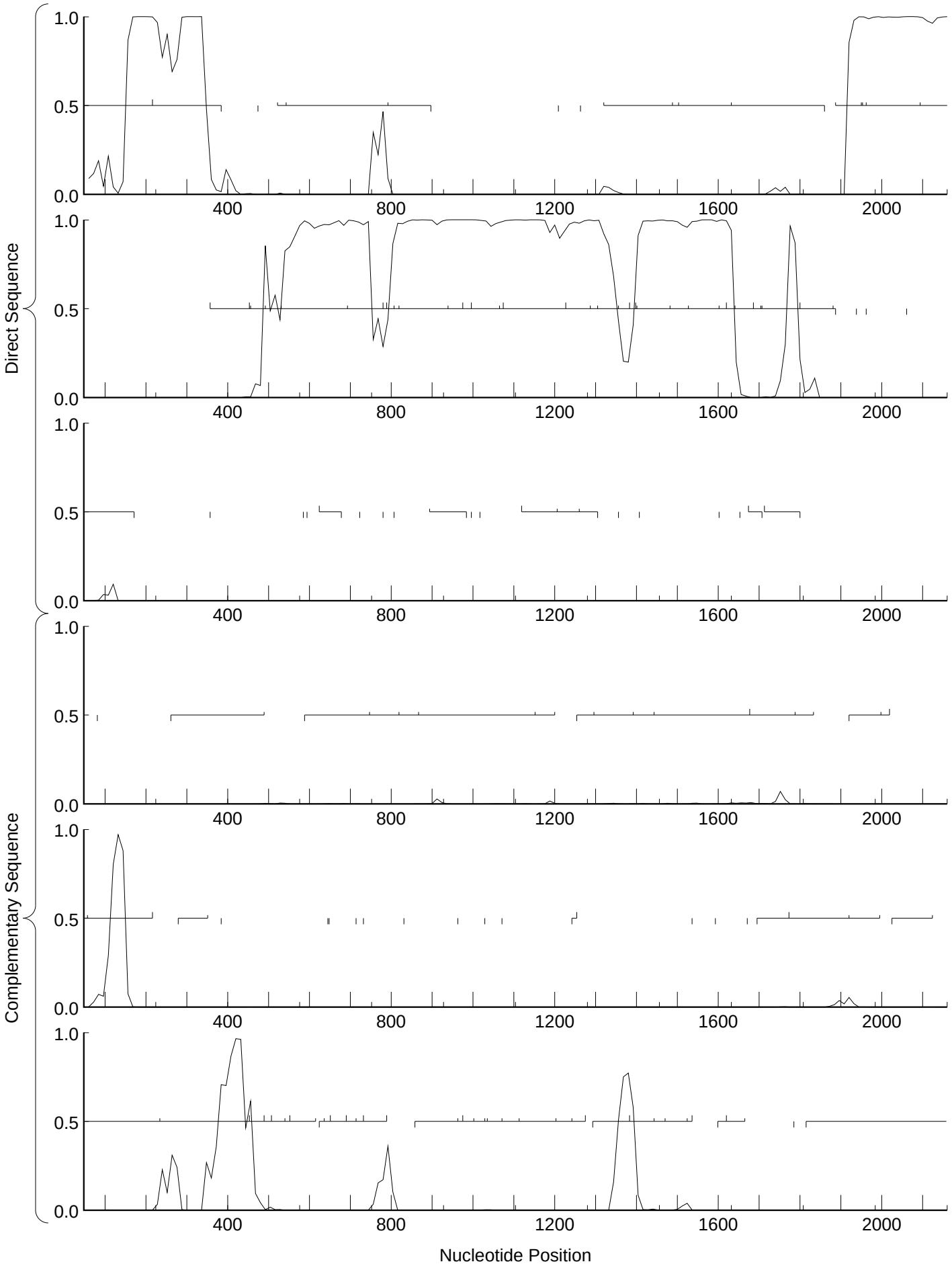
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Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

Order      :  4
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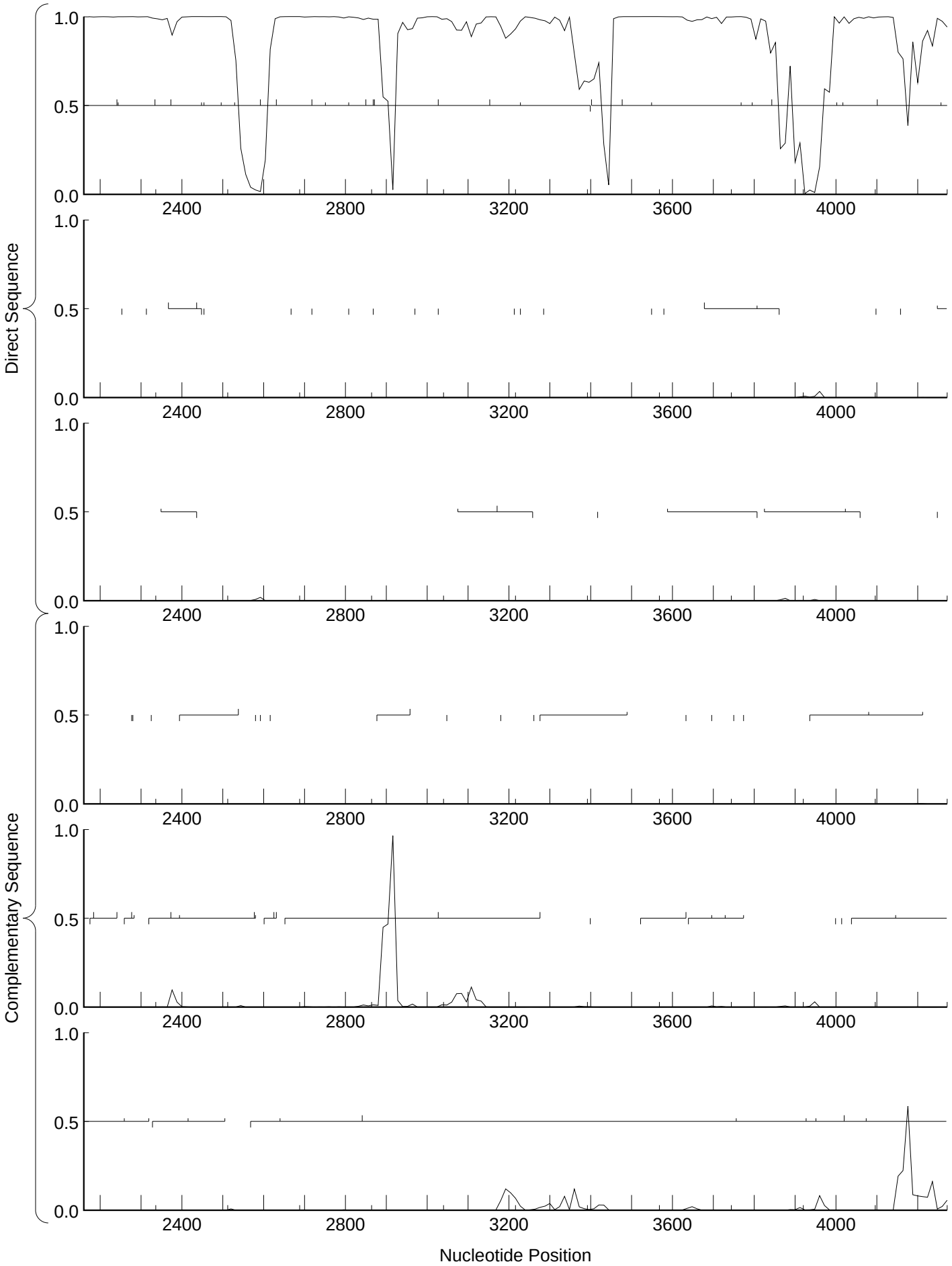

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

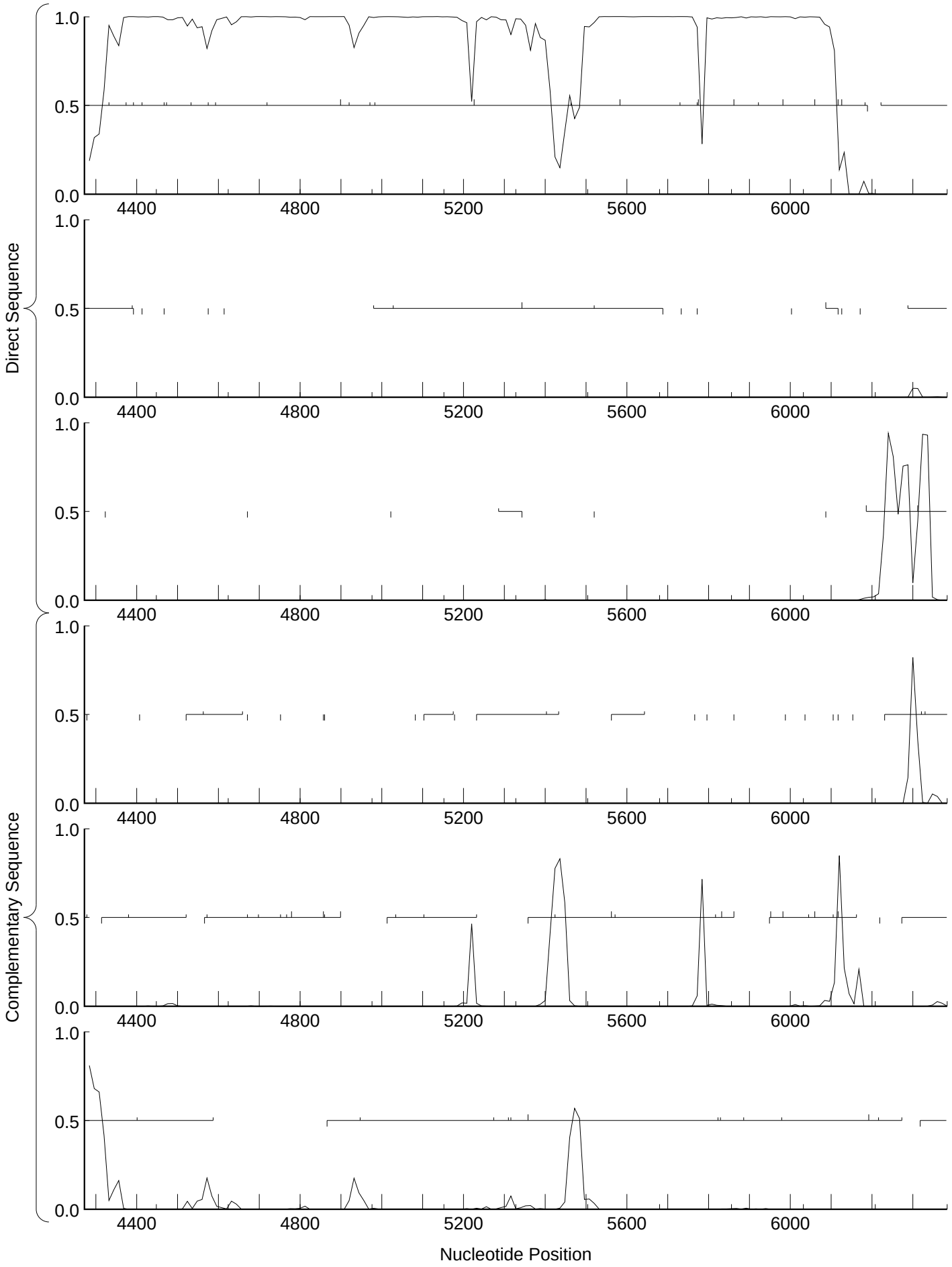
or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 2/21

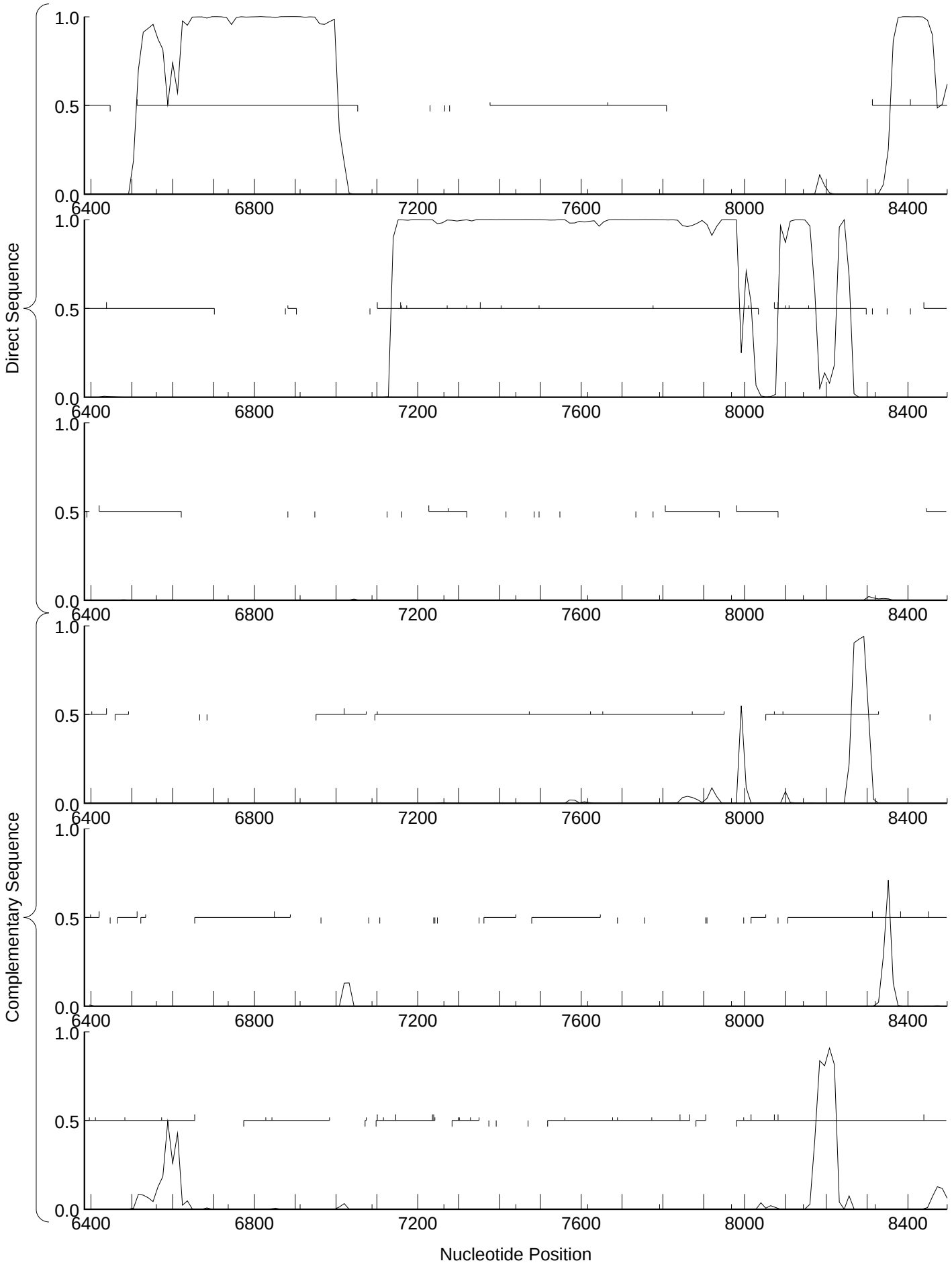


or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 3/21

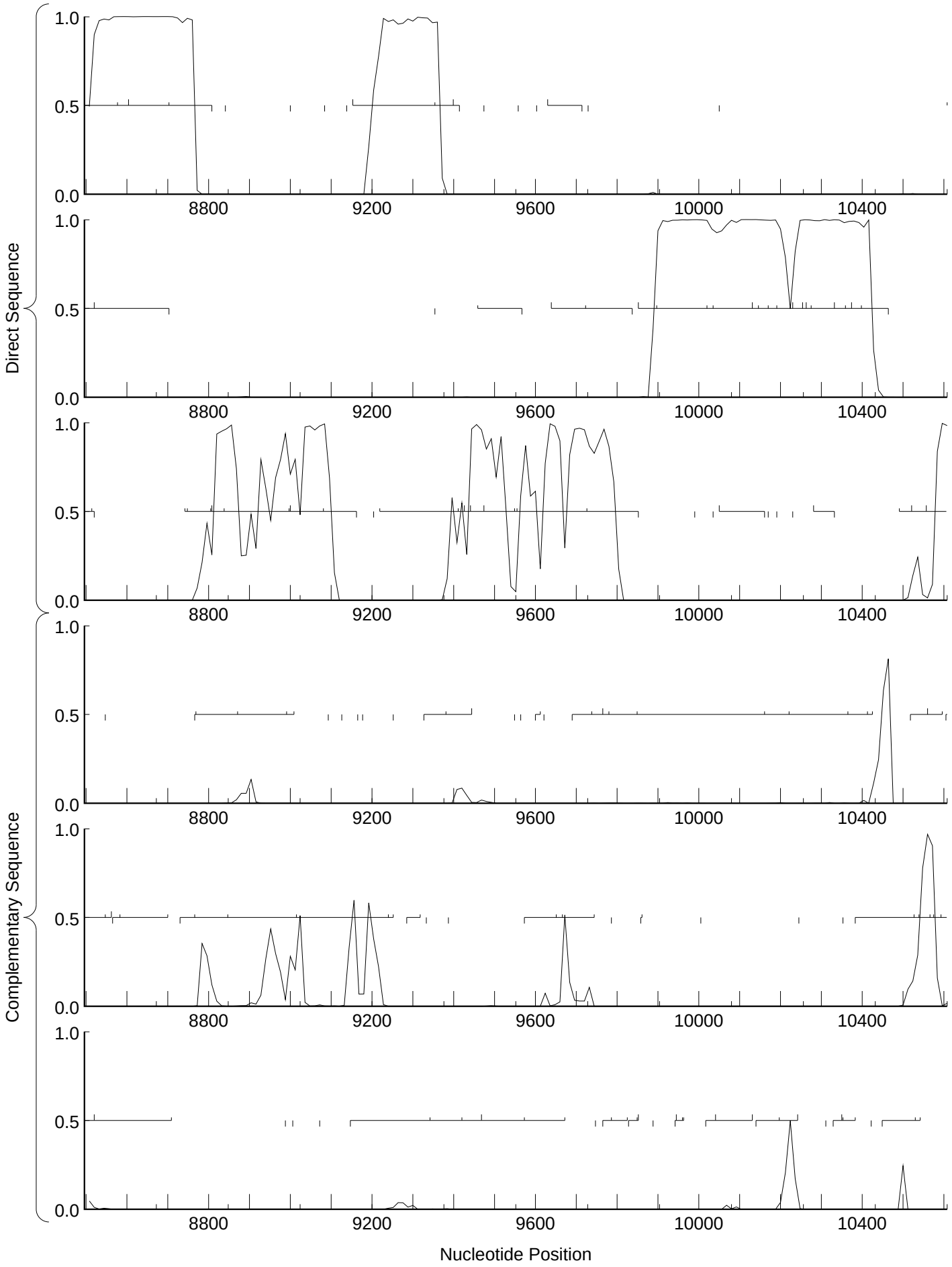


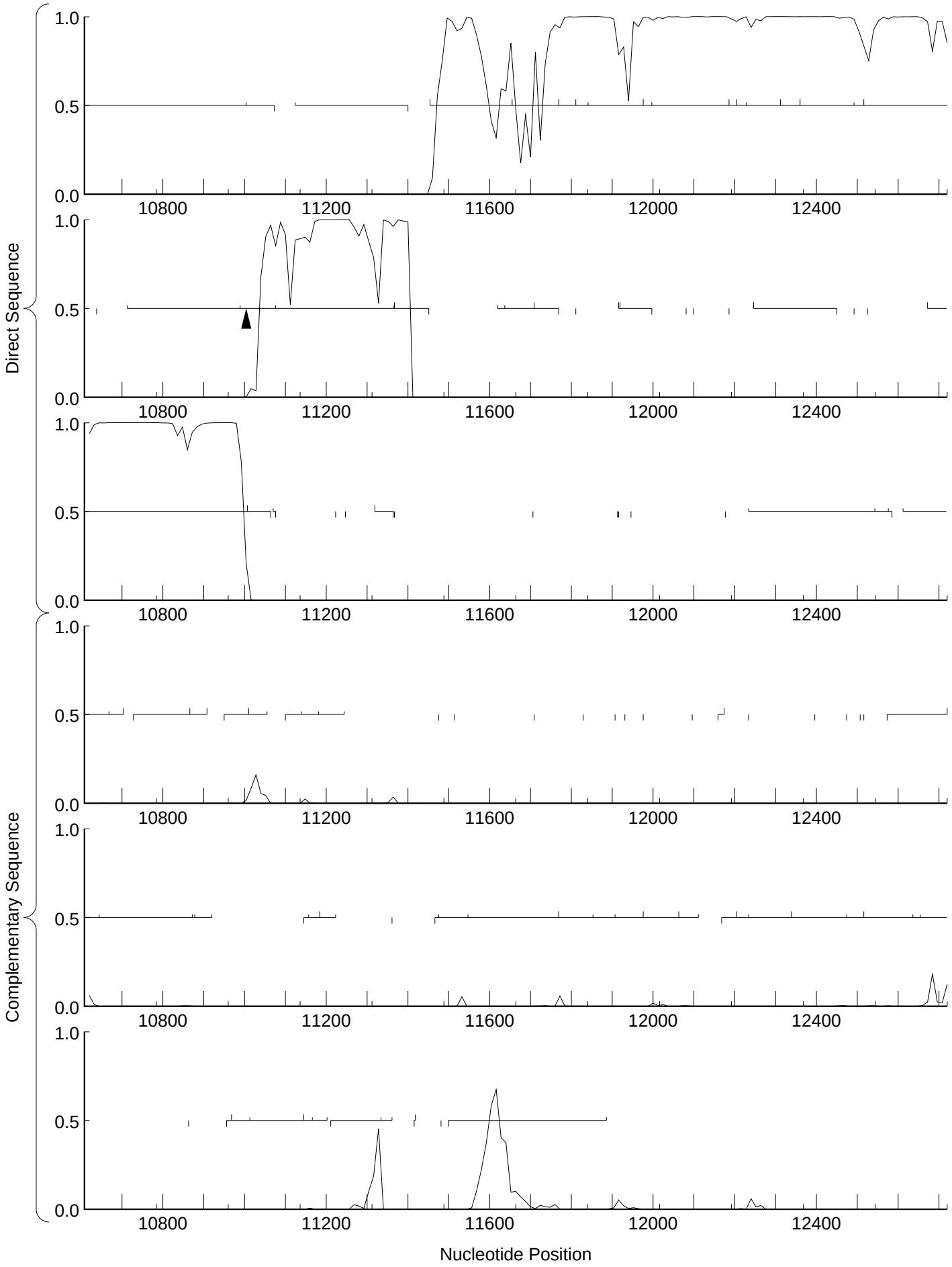
or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 4/21



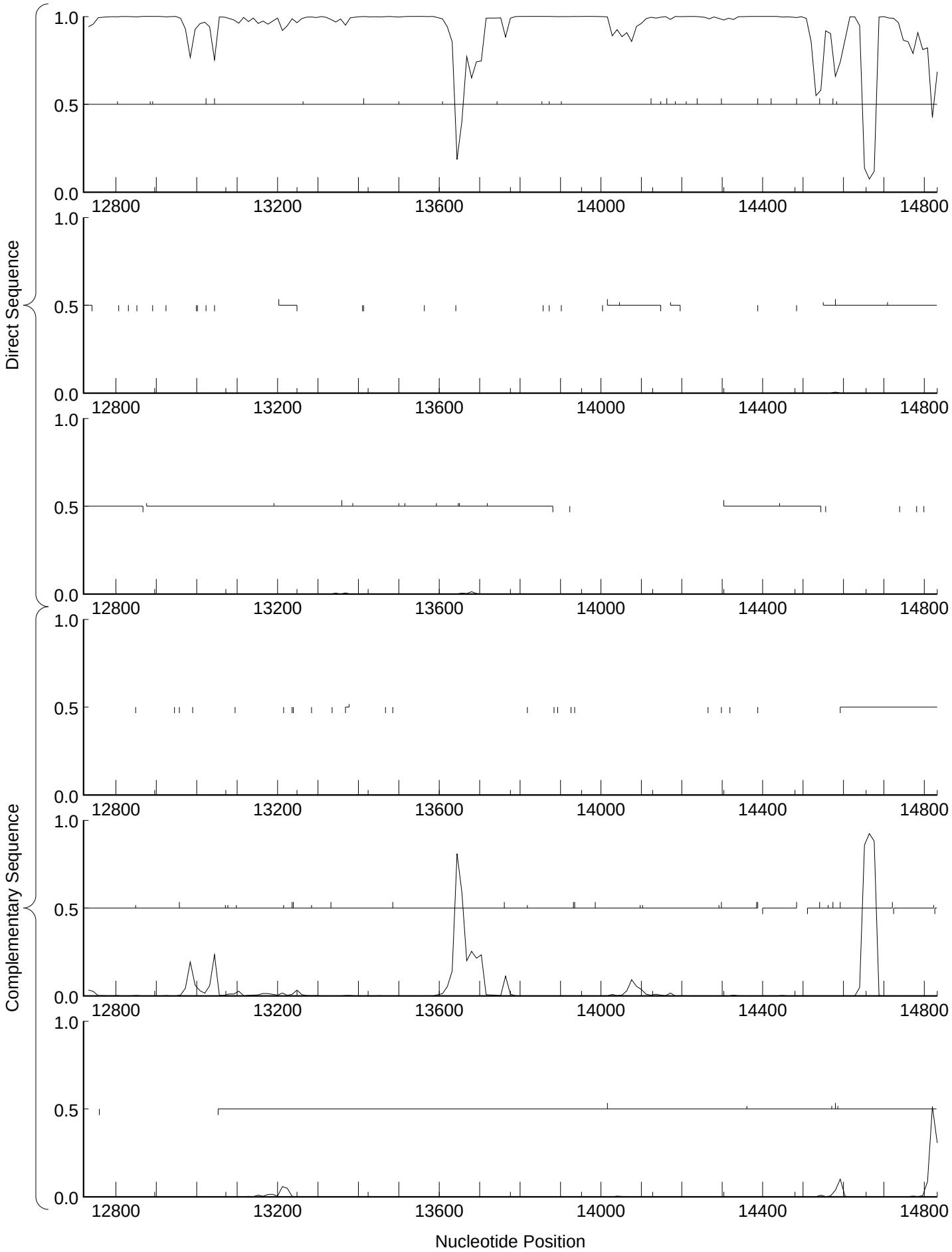

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
6400
6800
7200
7600
8000
8400
Nucleotide Position

or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 6/21

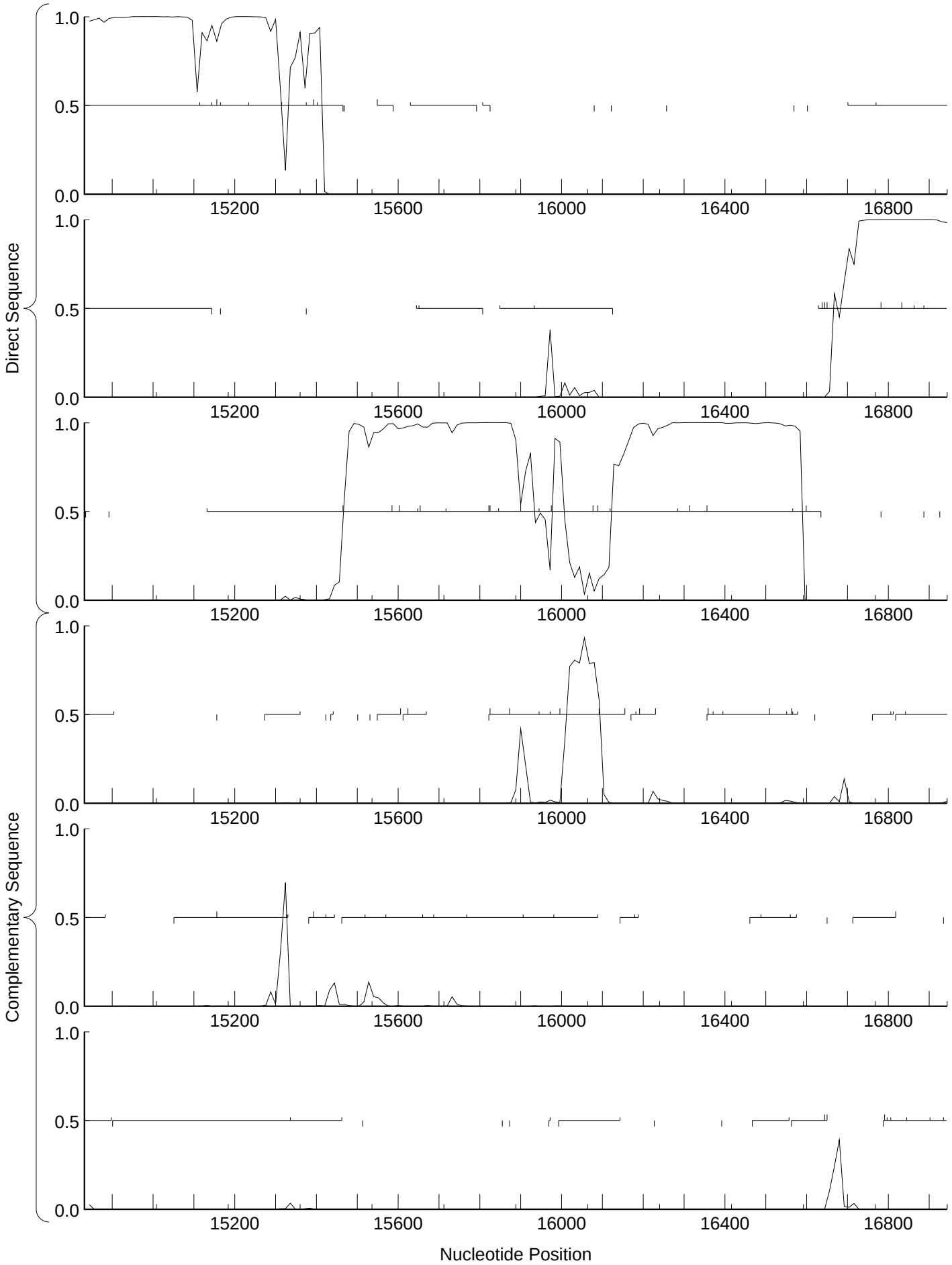



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
10800
11200
11600
12000
12400
Nucleotide Position

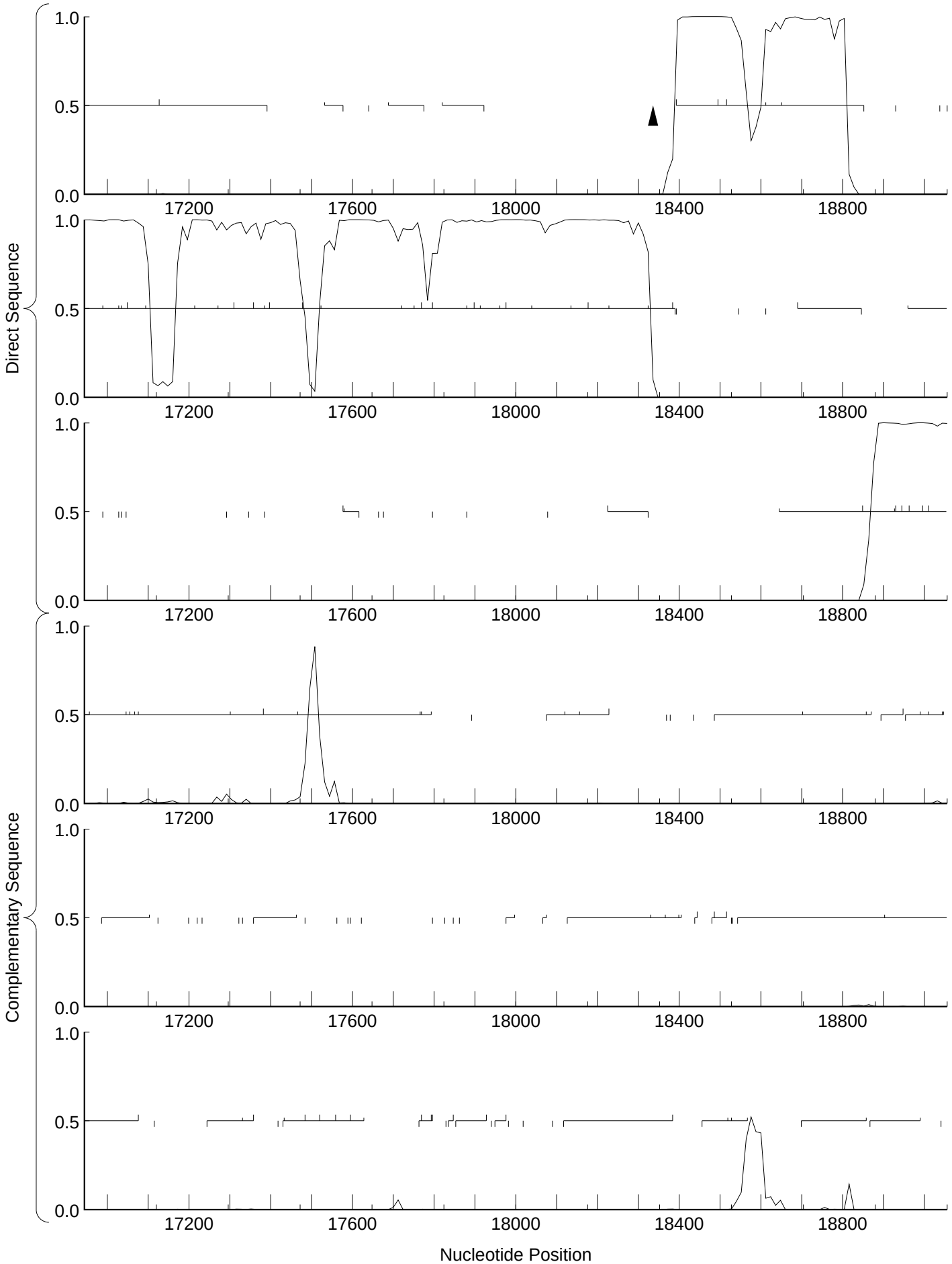
or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 8/21



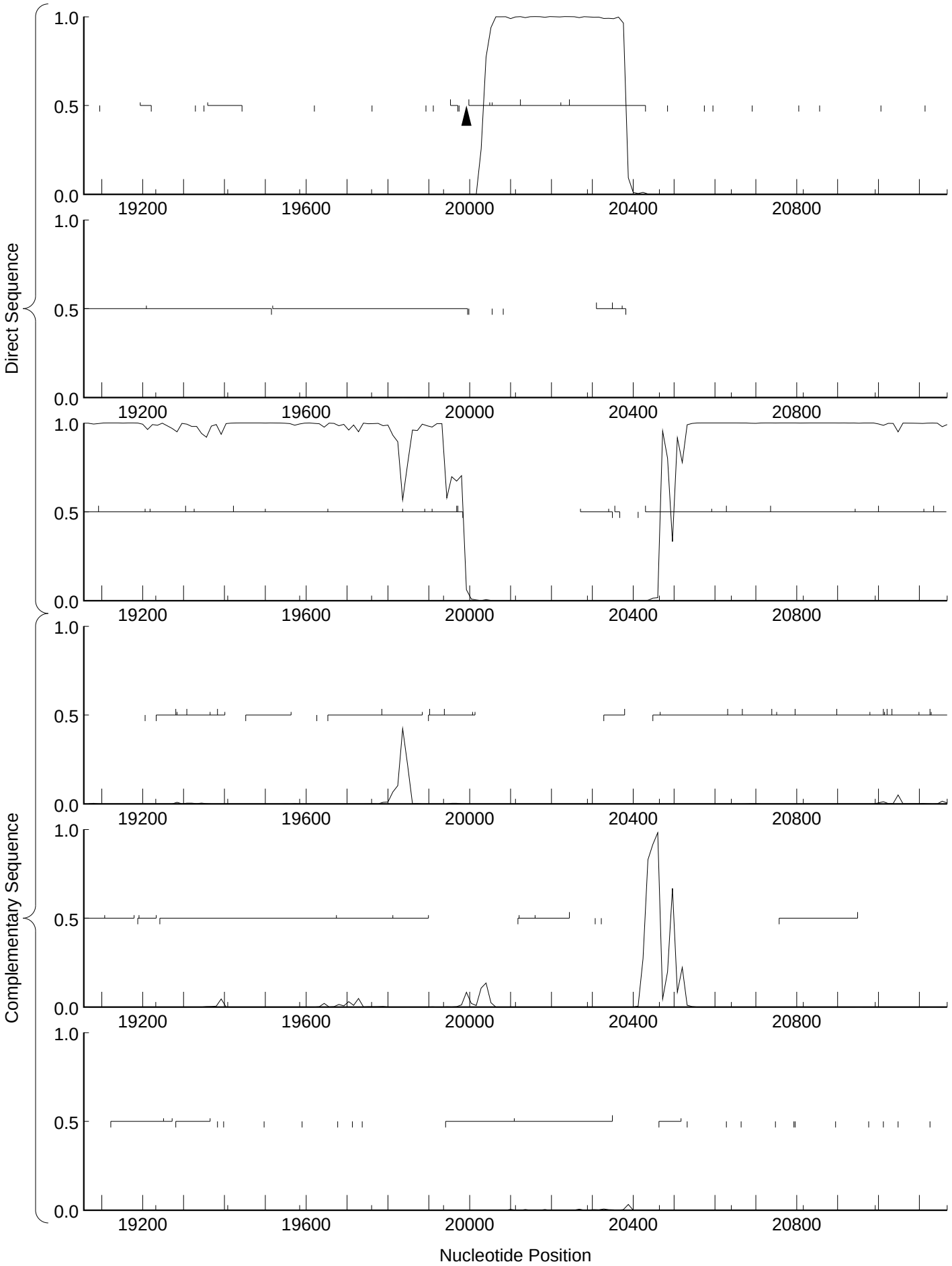
or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 9/21



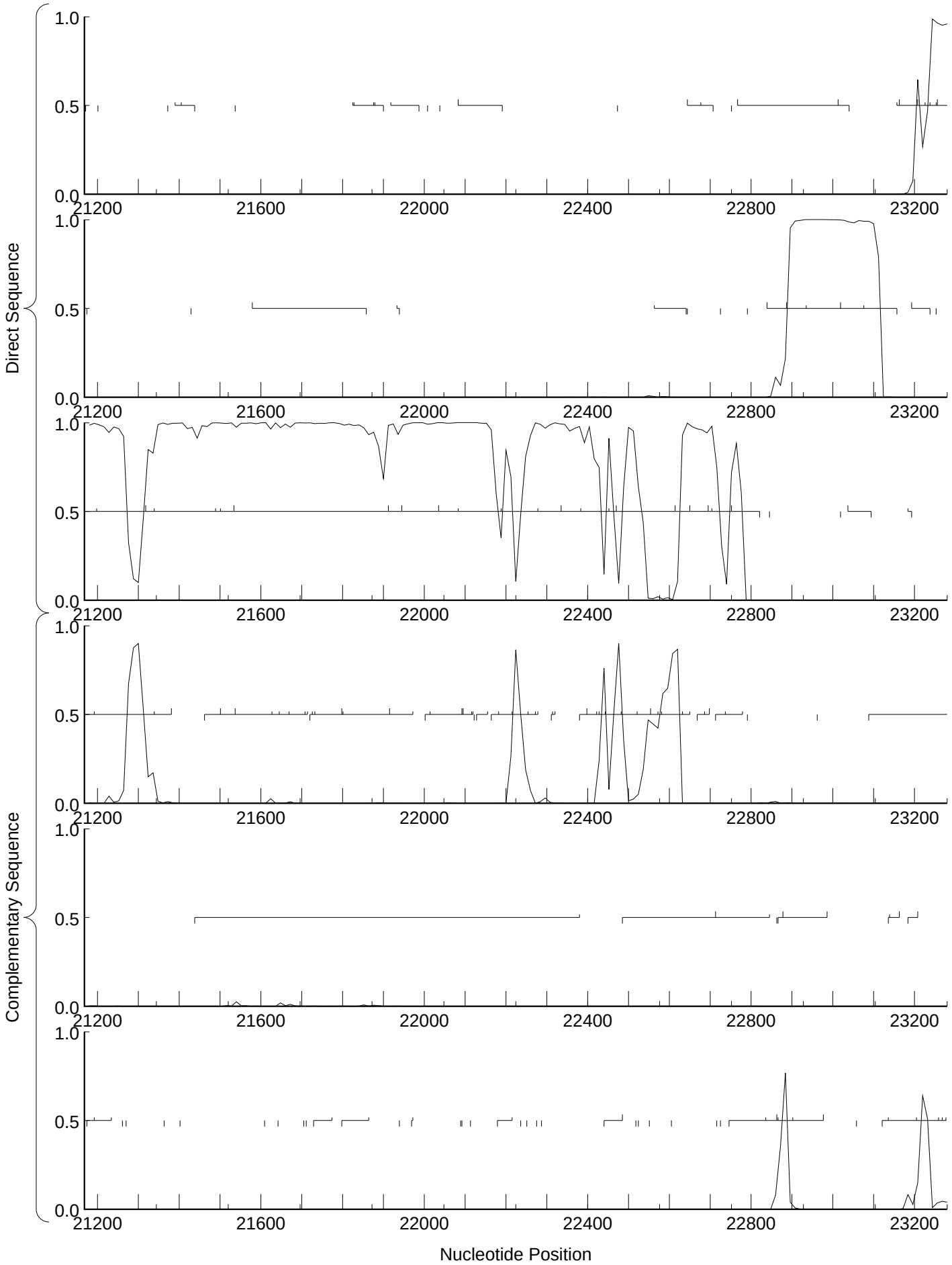
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 10/21



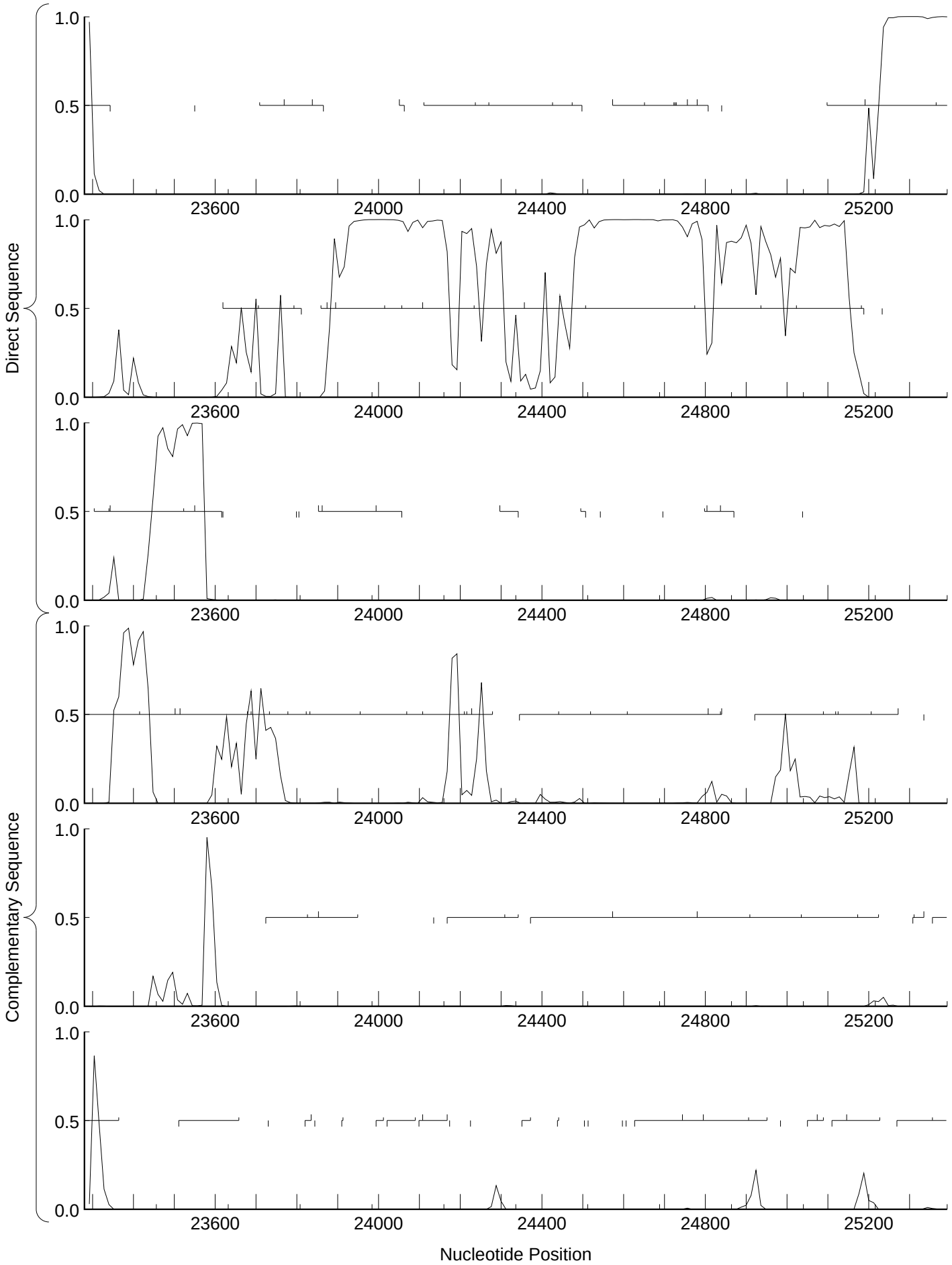
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 11/21



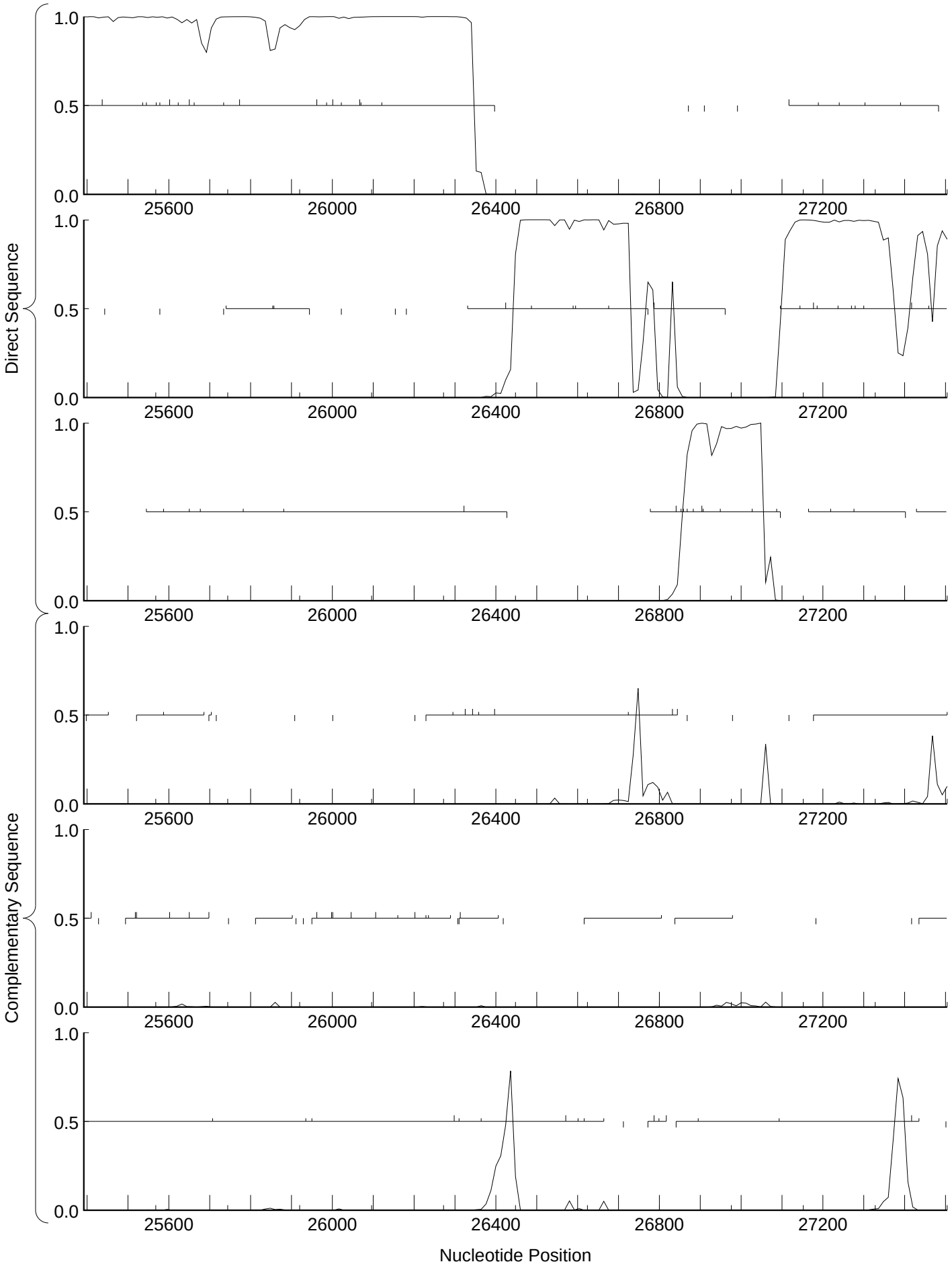
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 12/21



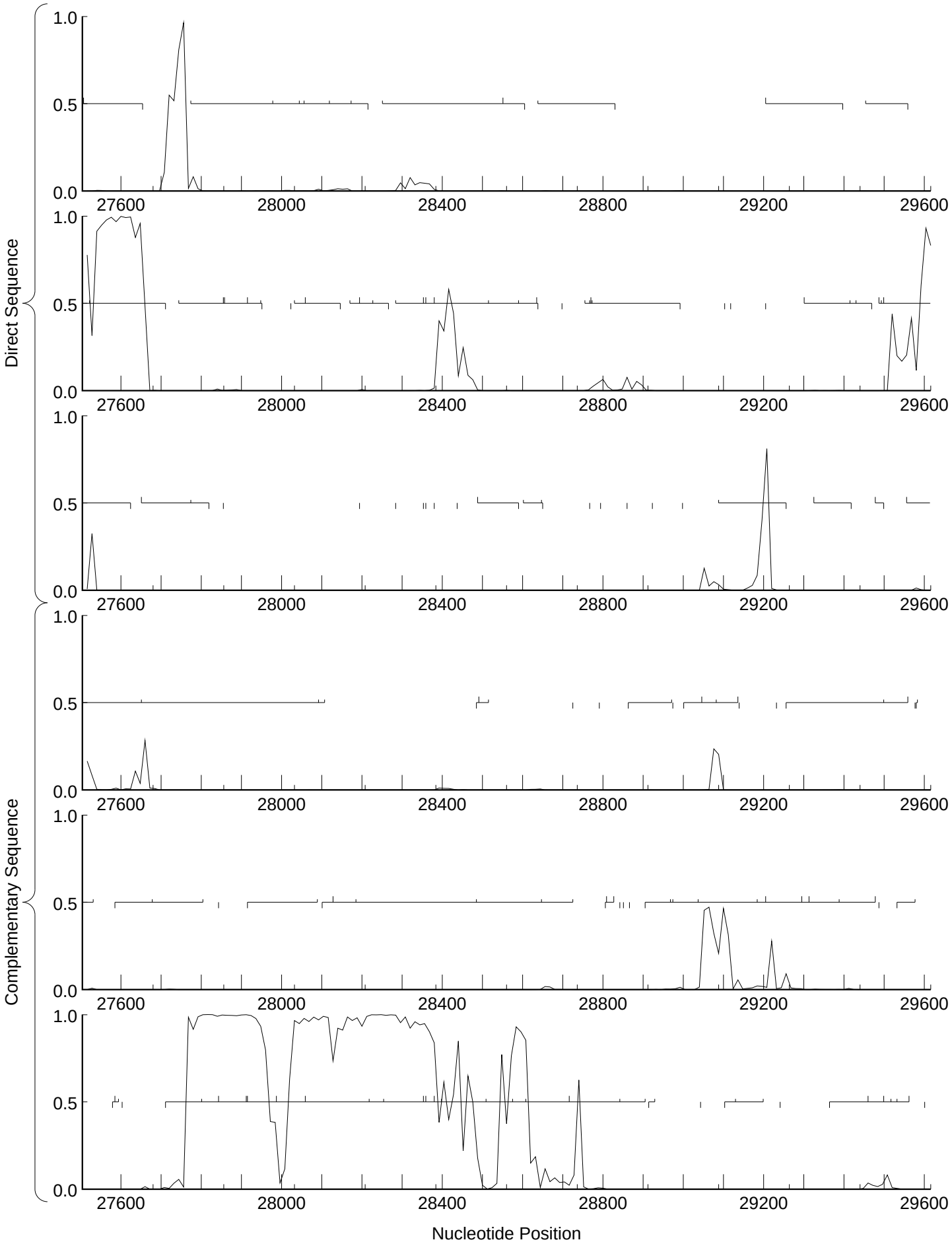
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 13/21



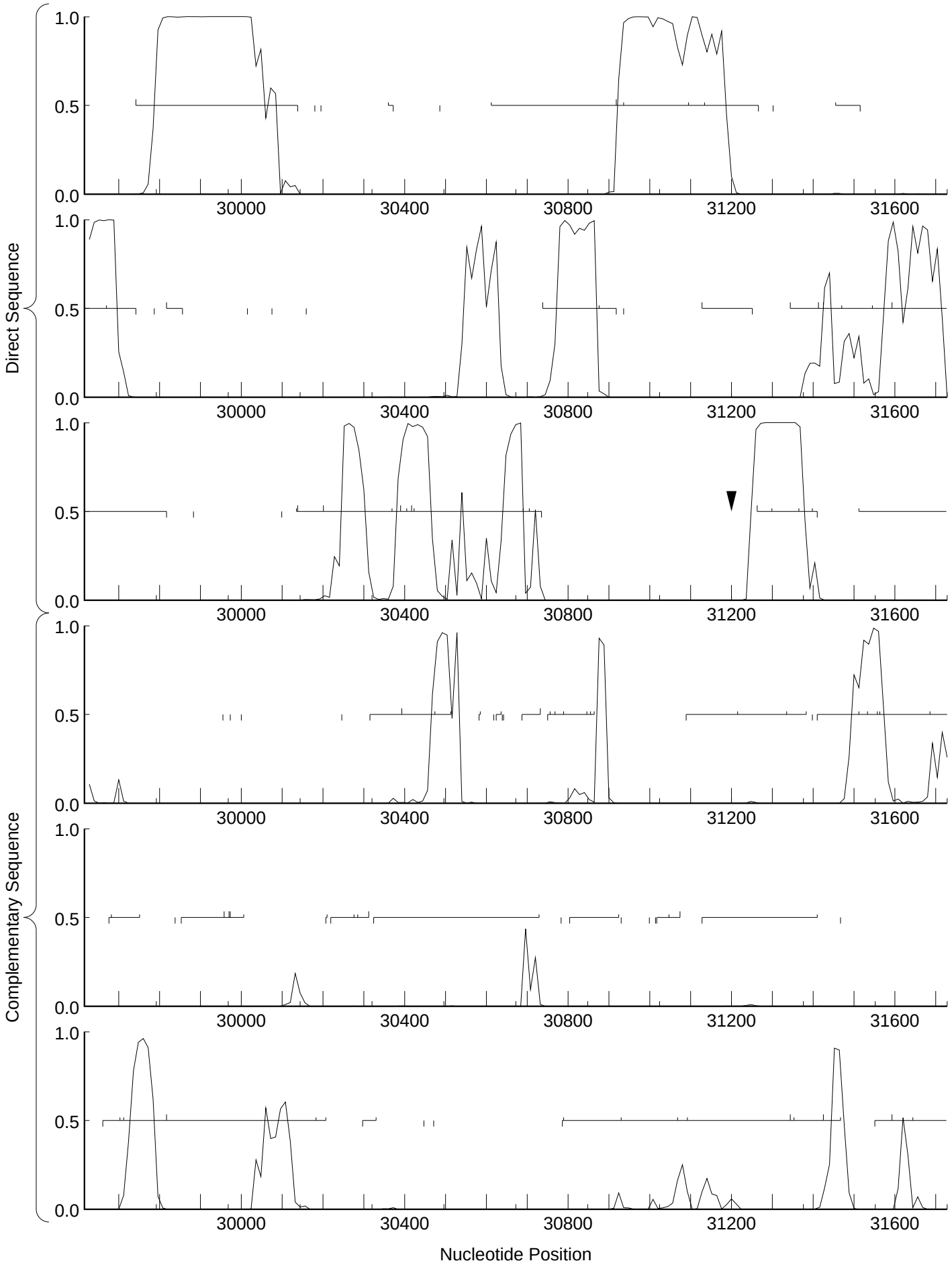
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 14/21



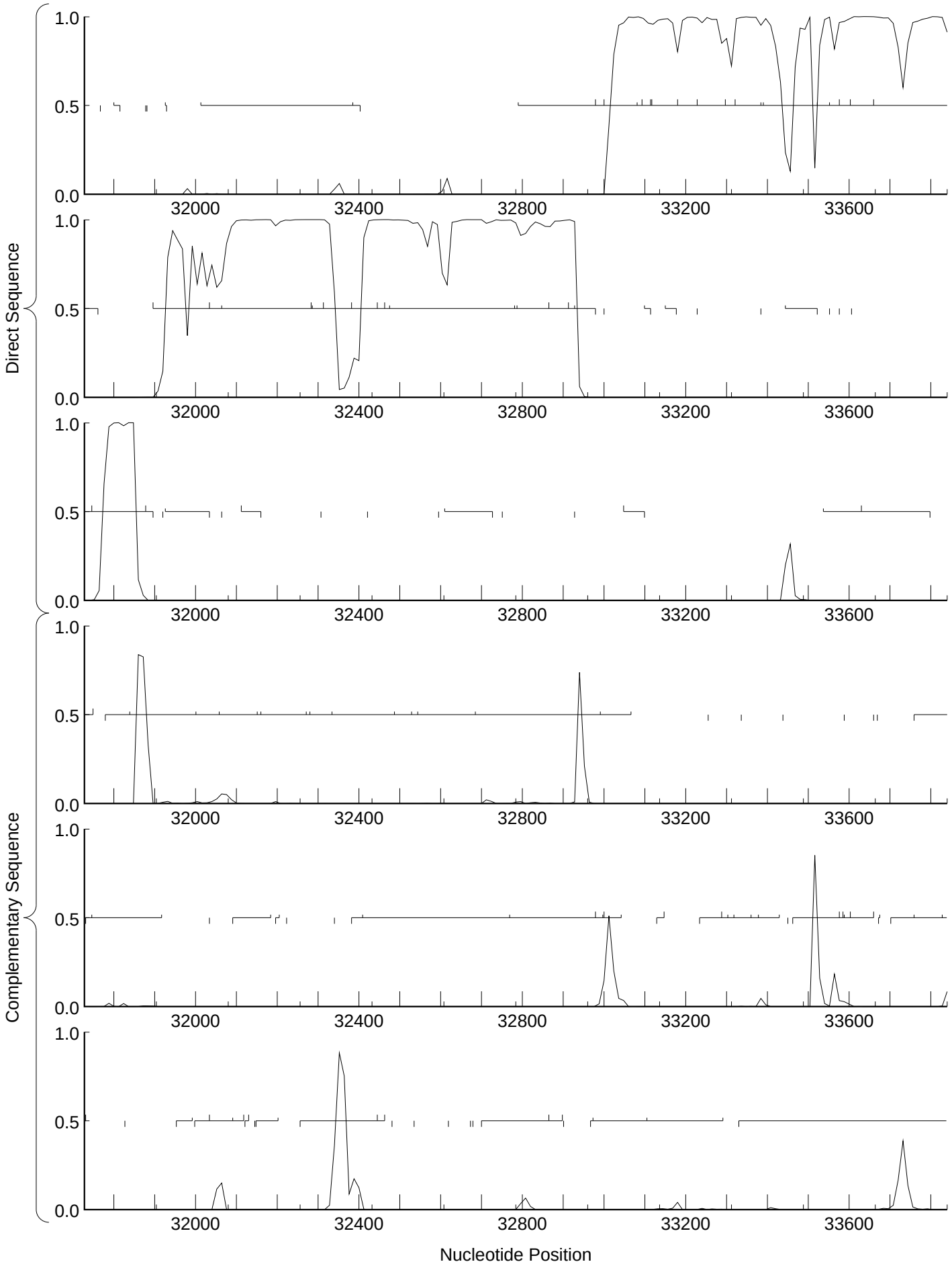
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 15/21



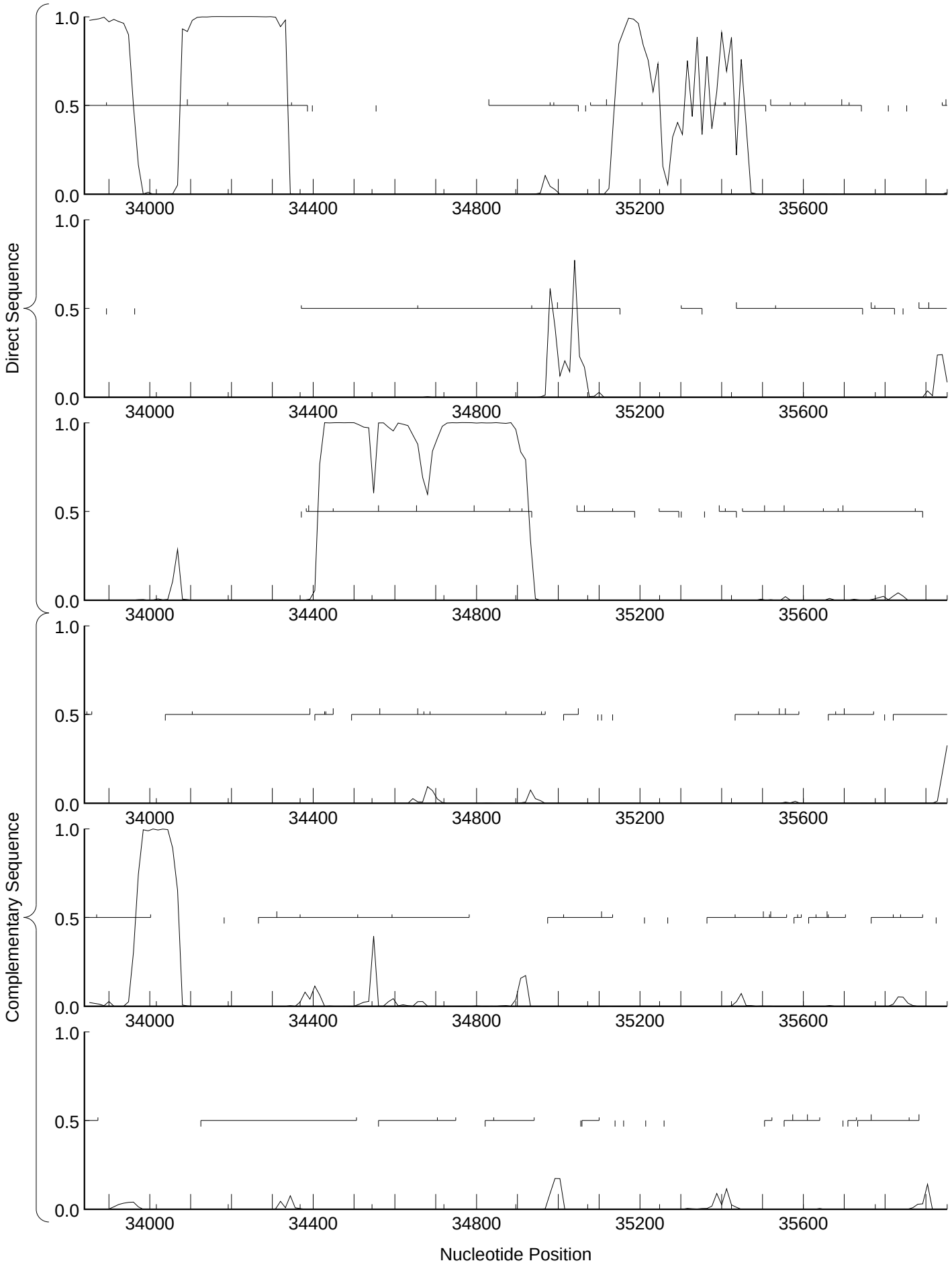
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 16/21



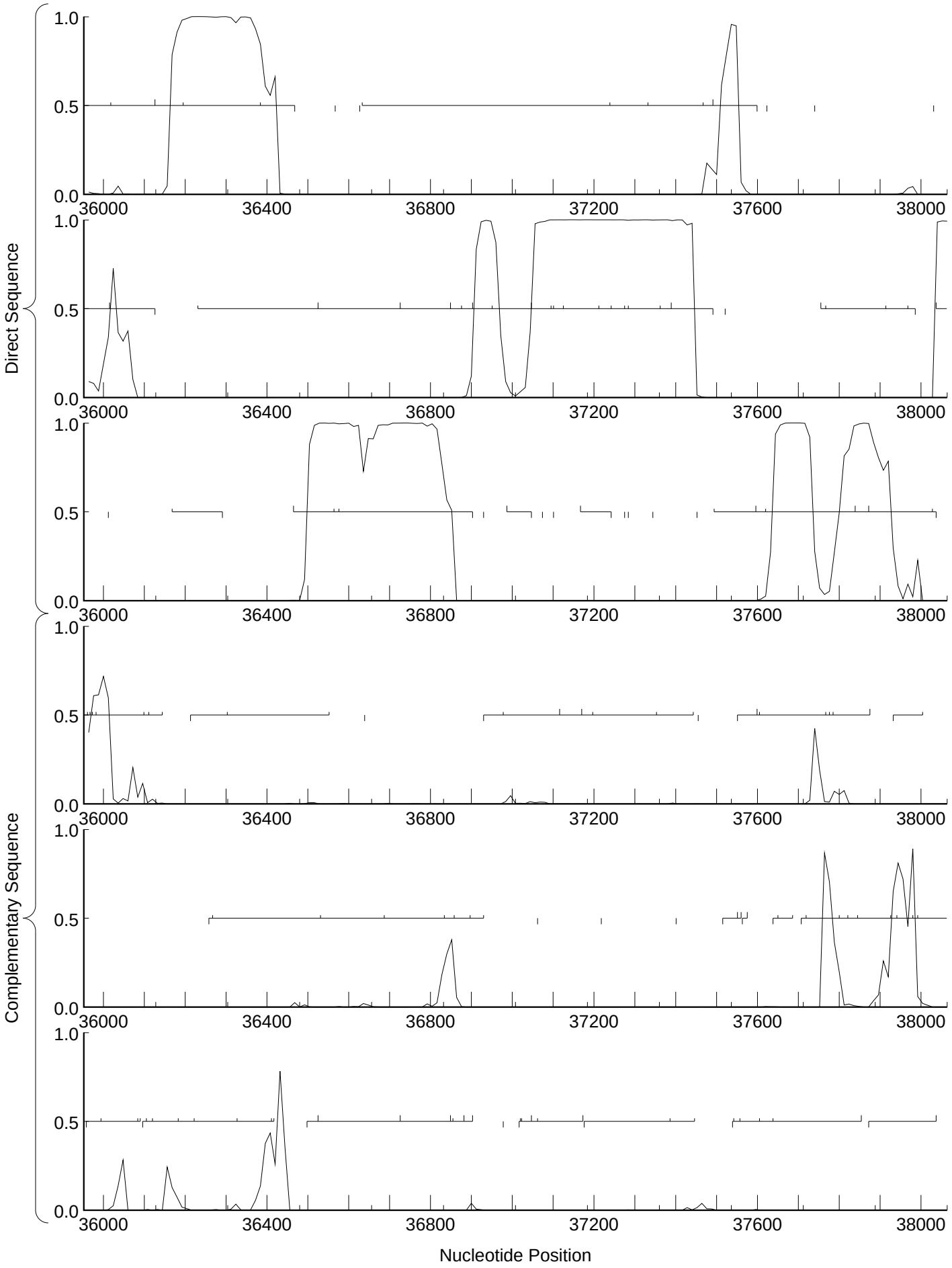
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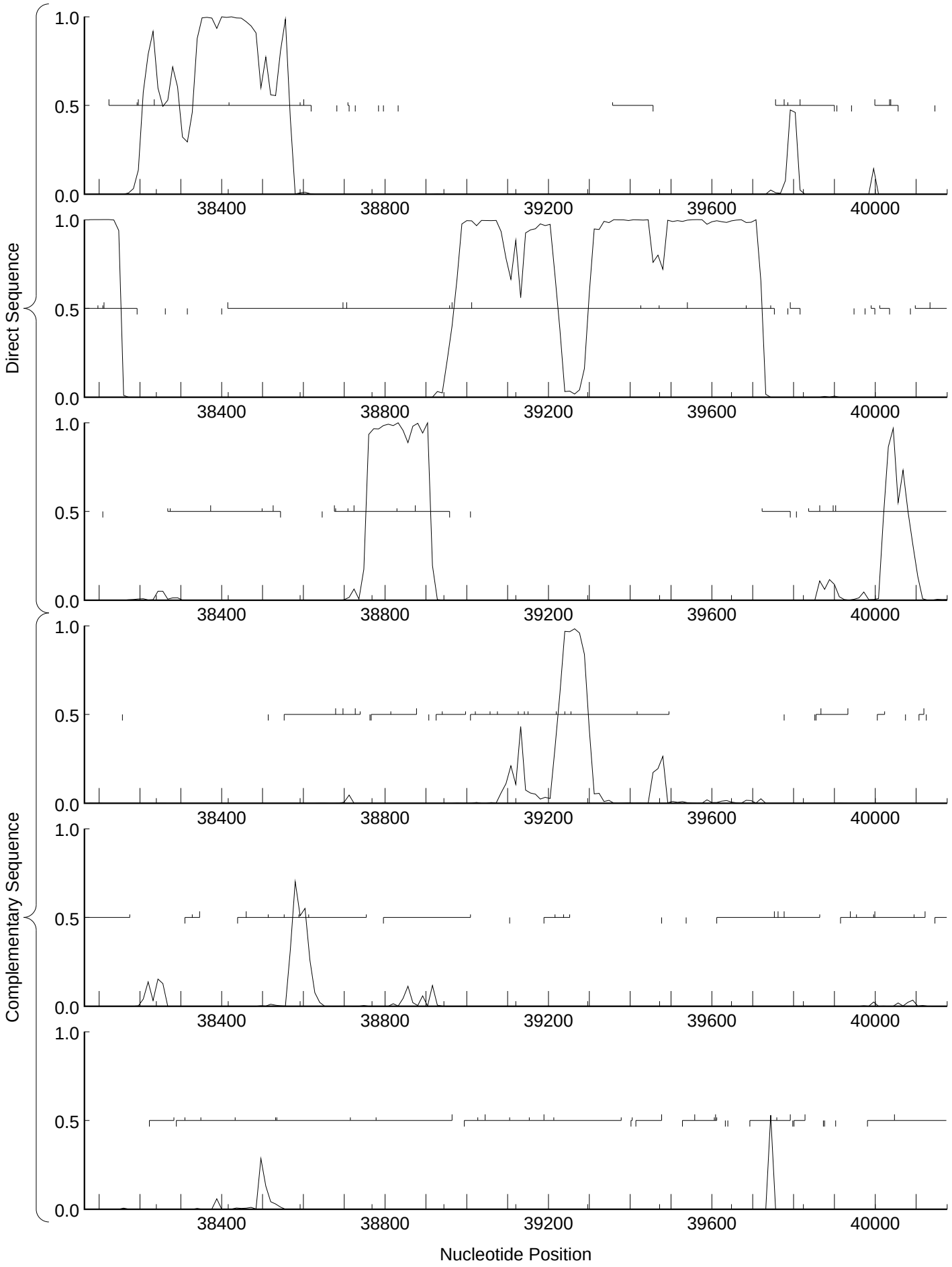
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 18/21



complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 19/21



complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 20/21



complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 21/21

