GeneMark

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

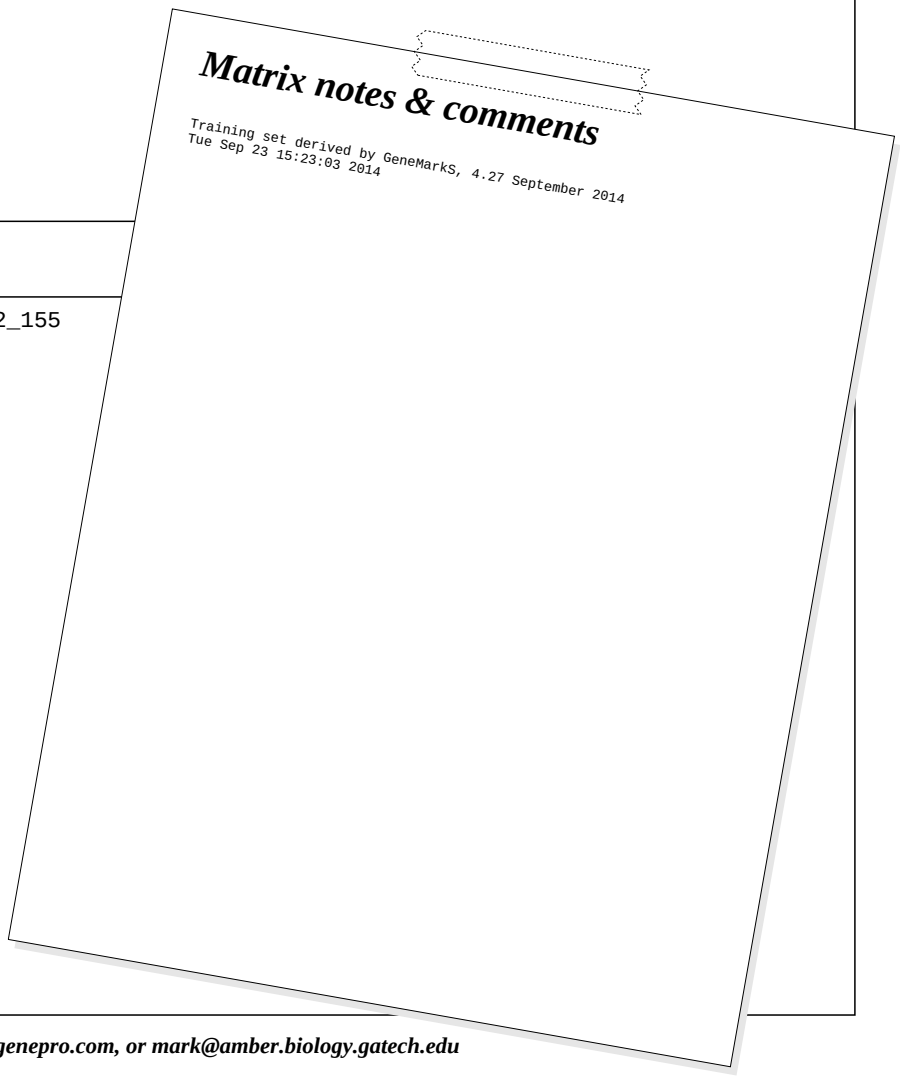
PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Paito complete sequence, 42311 bp including 11-base 3’ overhang (CCCCAT
Analysis Date    :  3/29/24 at 6:35:48
Pages            :  22
Sequence Length  :  42311 bp
GC Content       :   66.02%

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

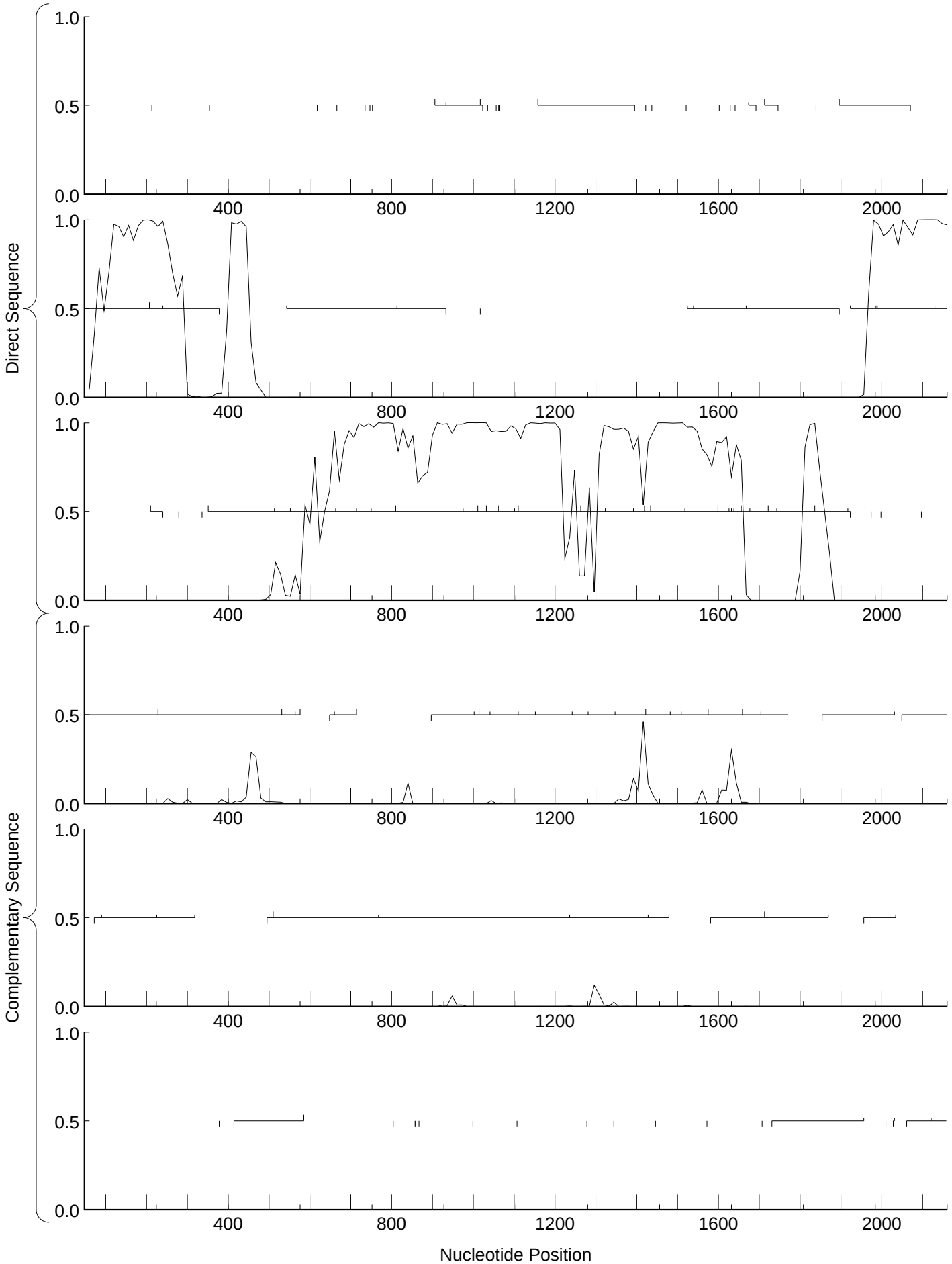
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Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

Order      :  4
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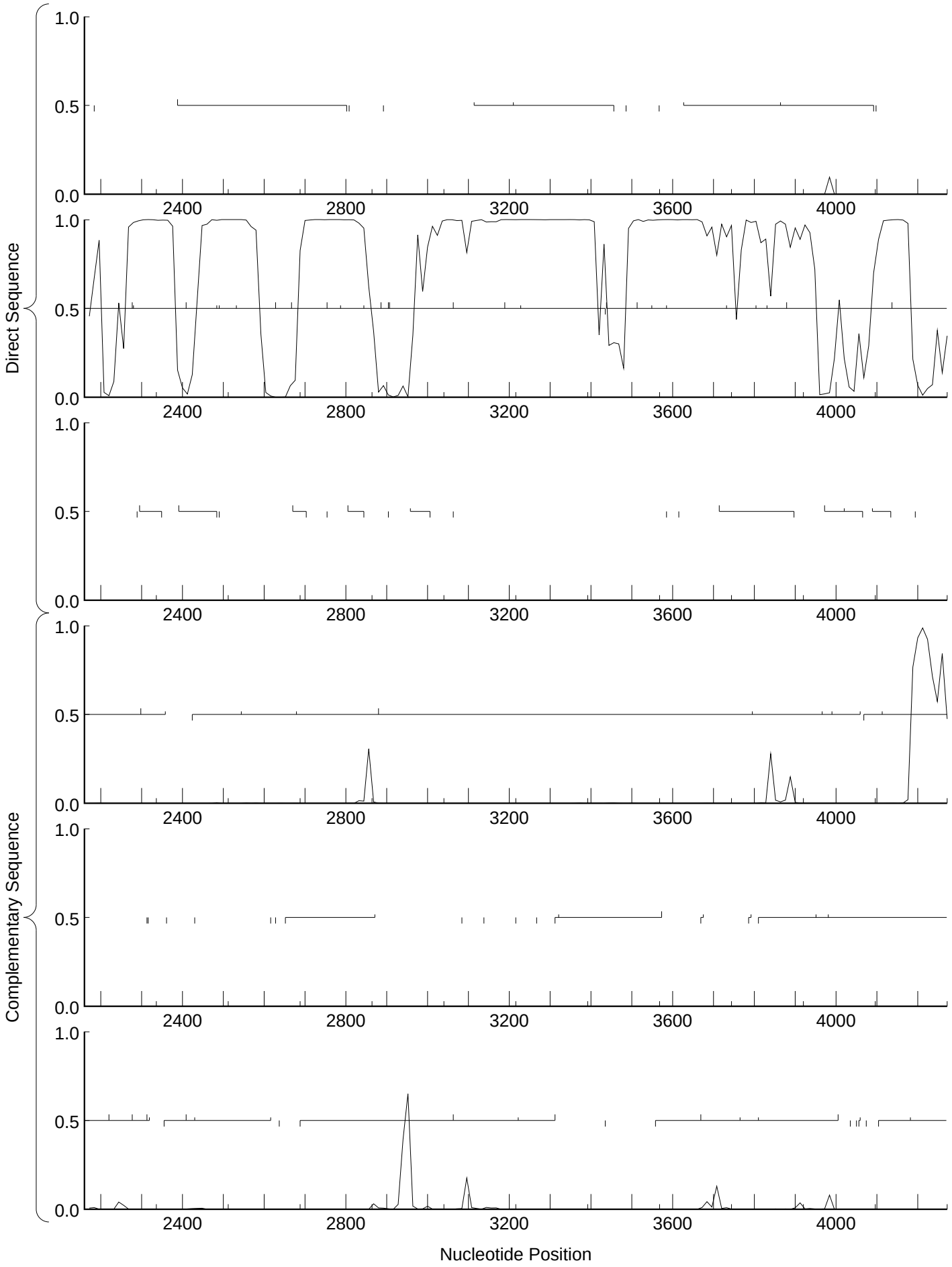

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

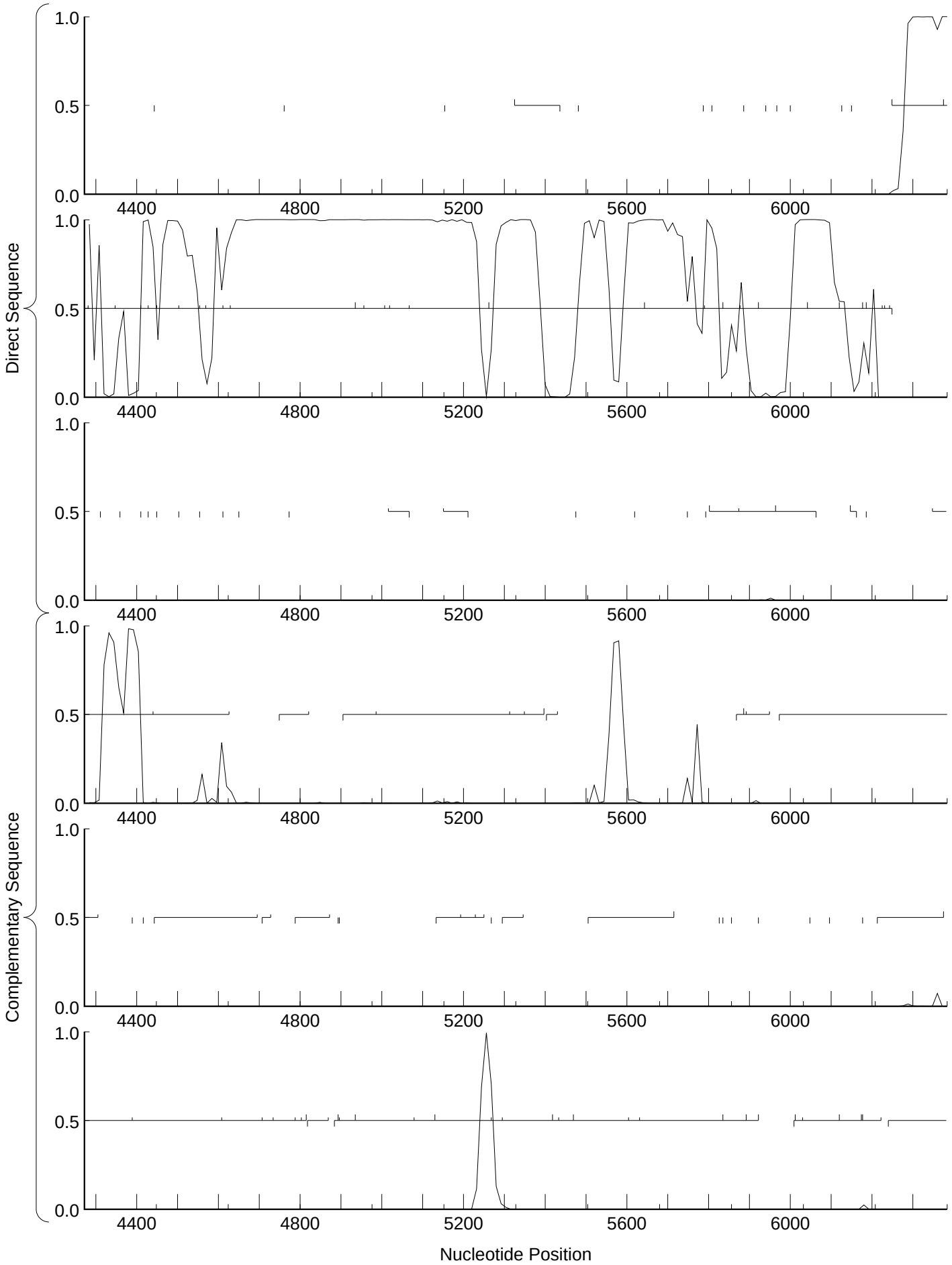
complete sequence, 42311 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 2/22



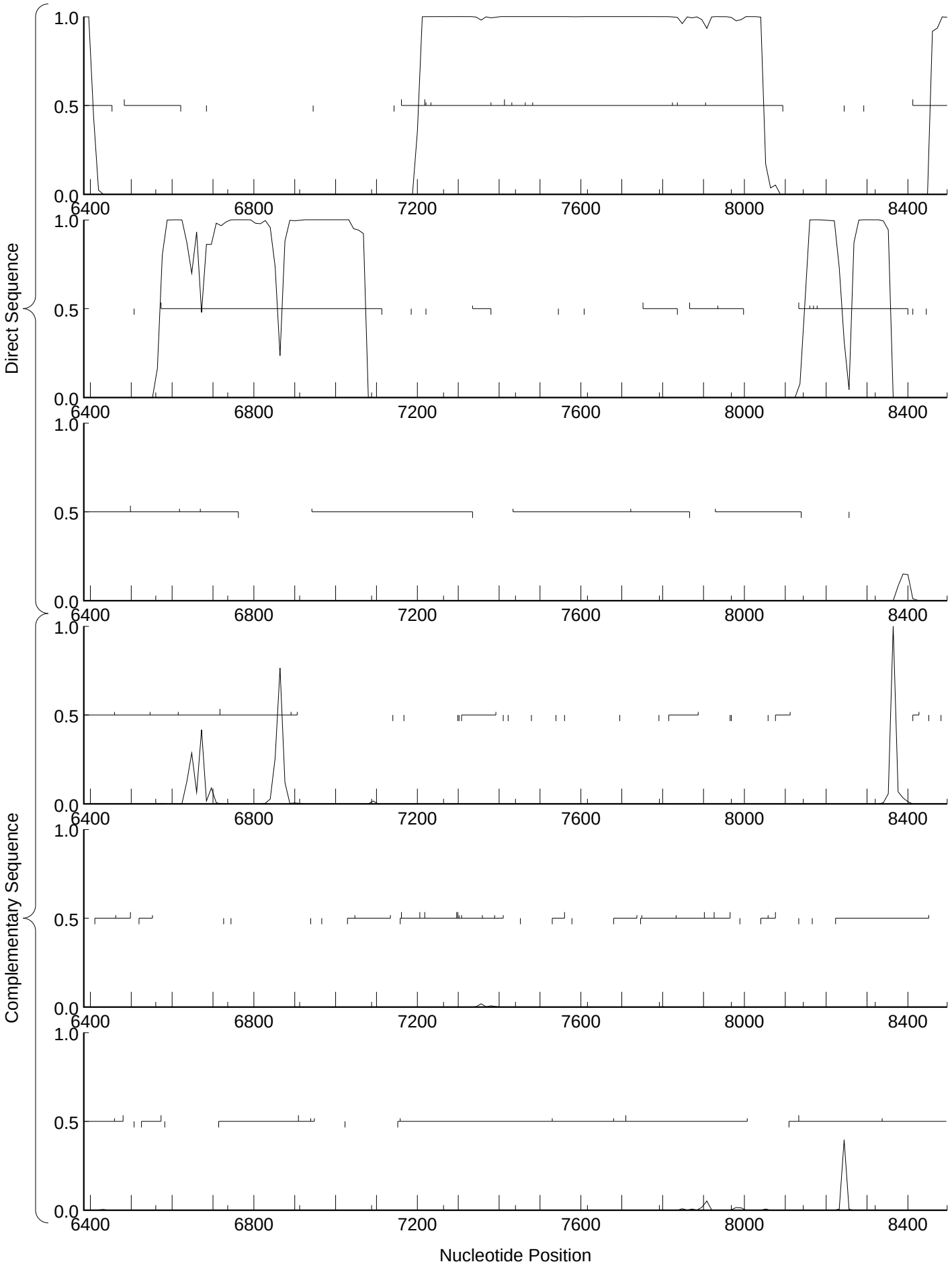
complete sequence, 42311 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 3/22



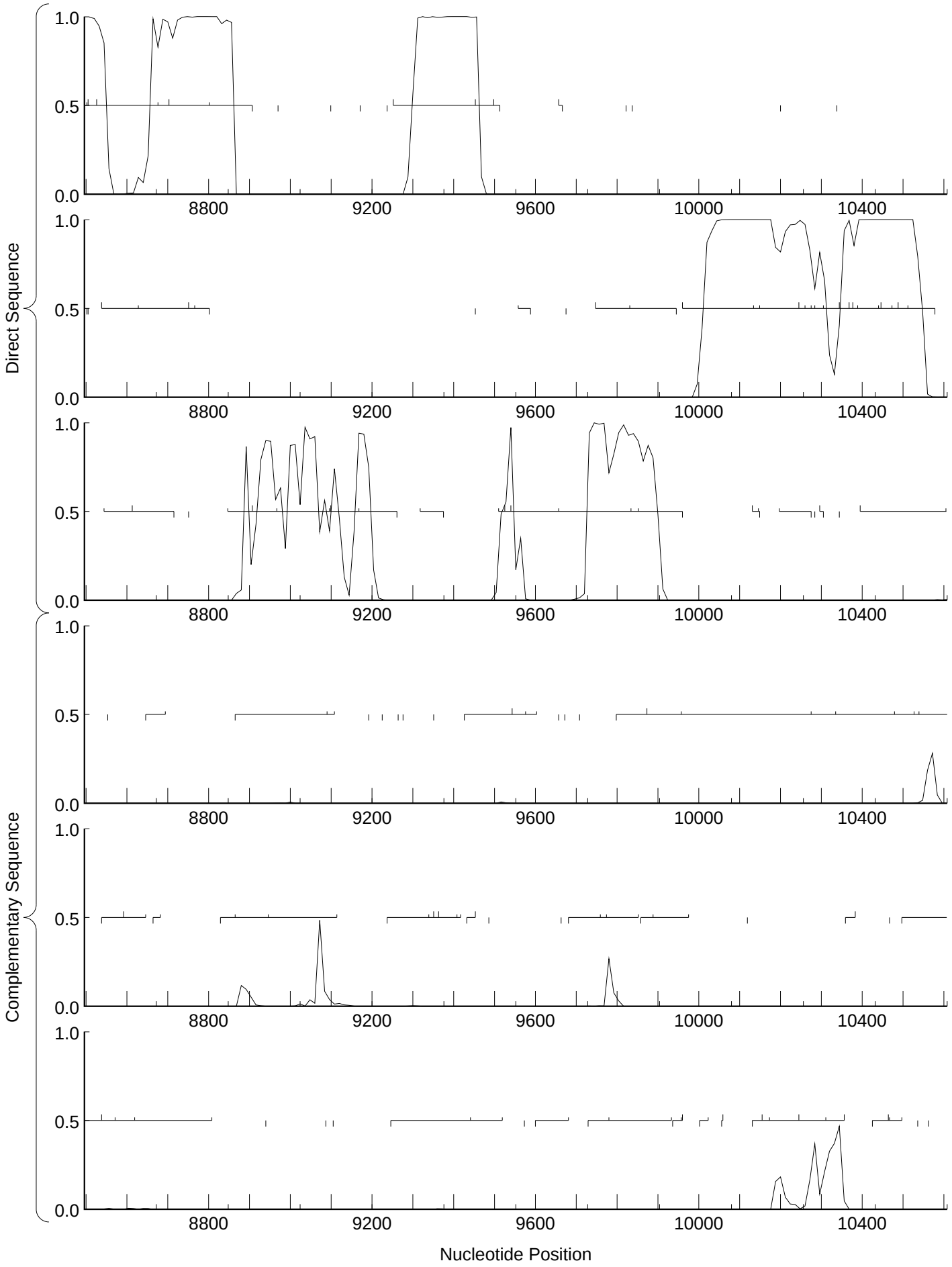
complete sequence, 42311 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 4/22



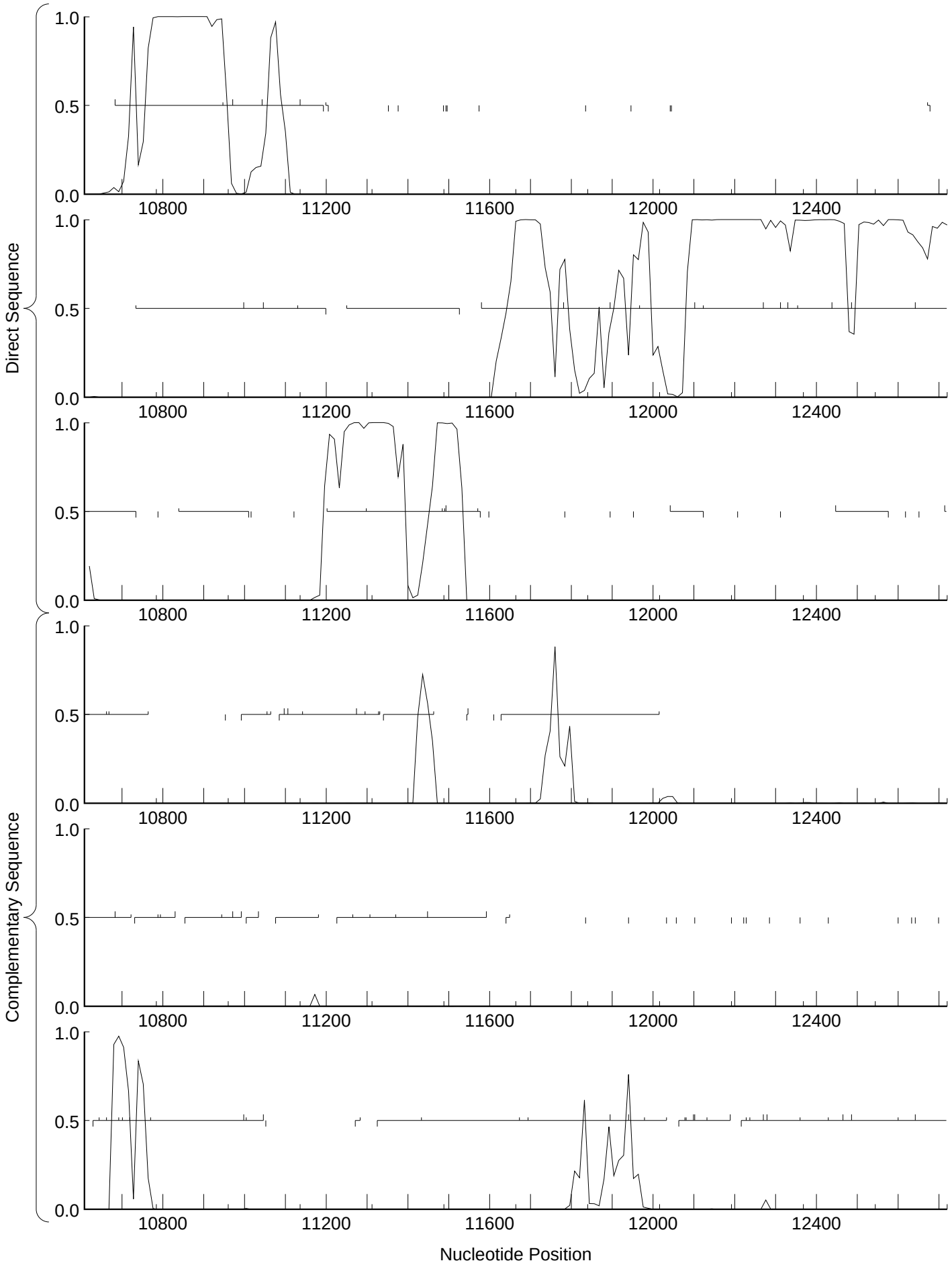
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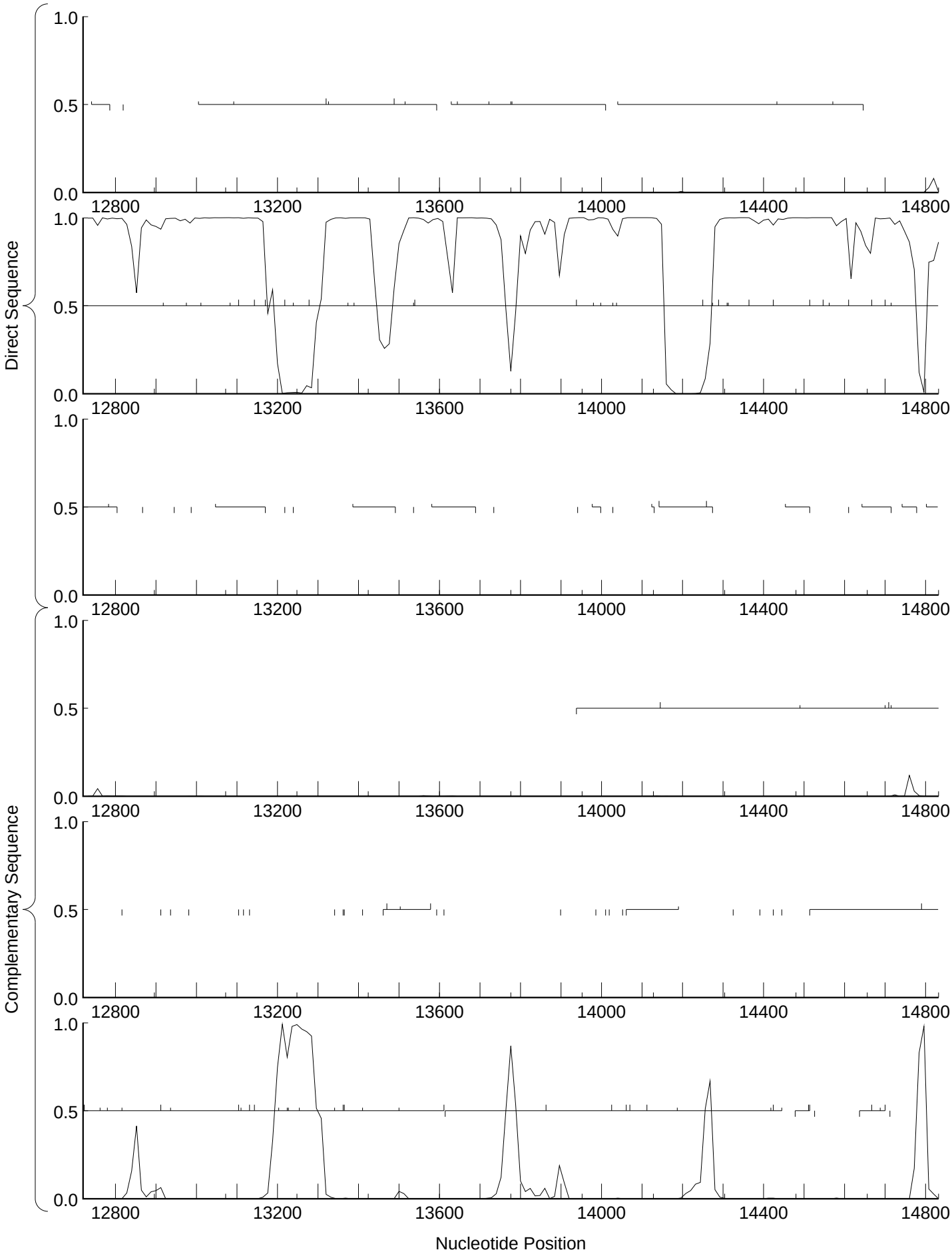
complete sequence, 42311 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 6/22



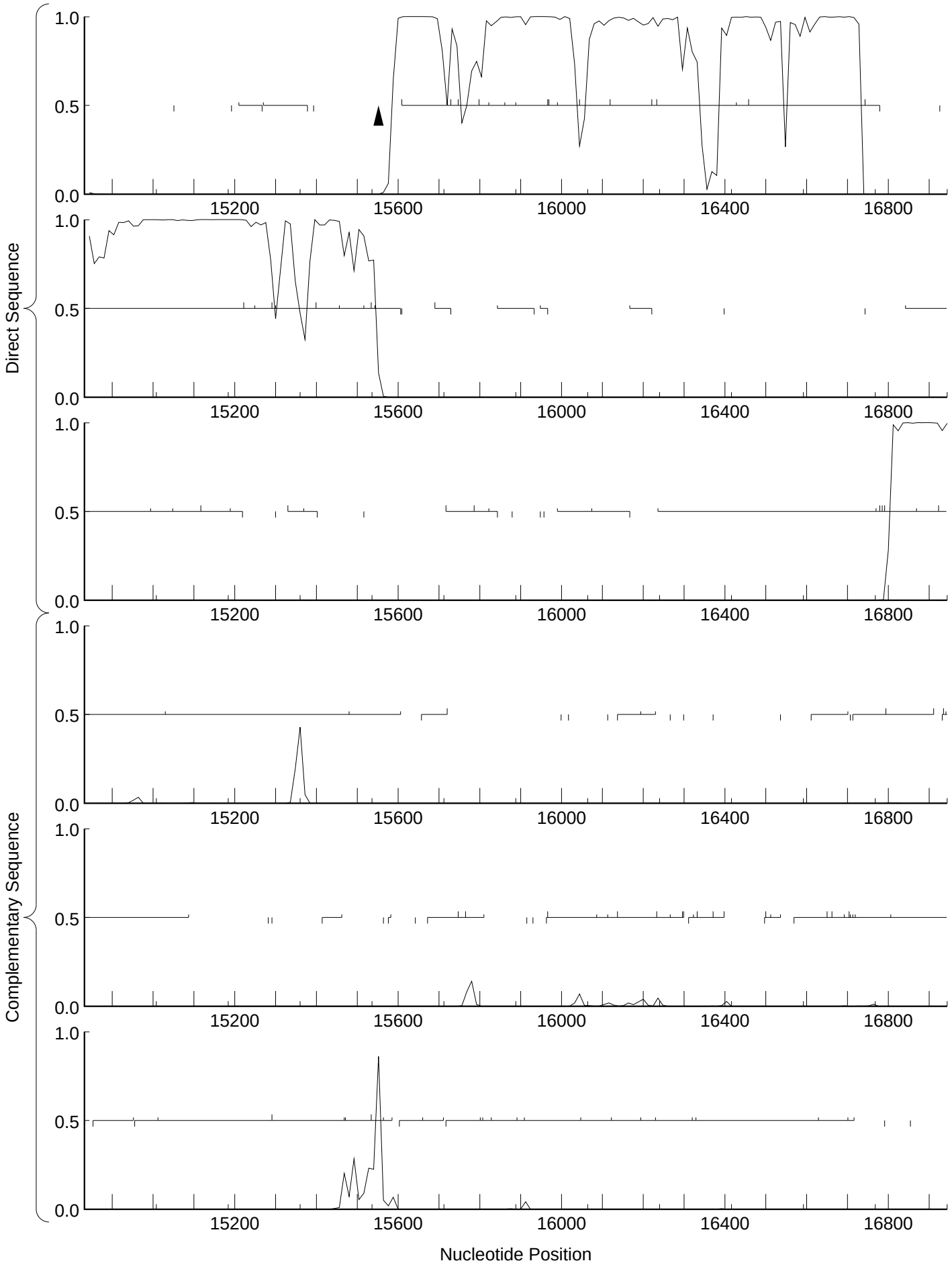
complete sequence, 42311 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 7/22



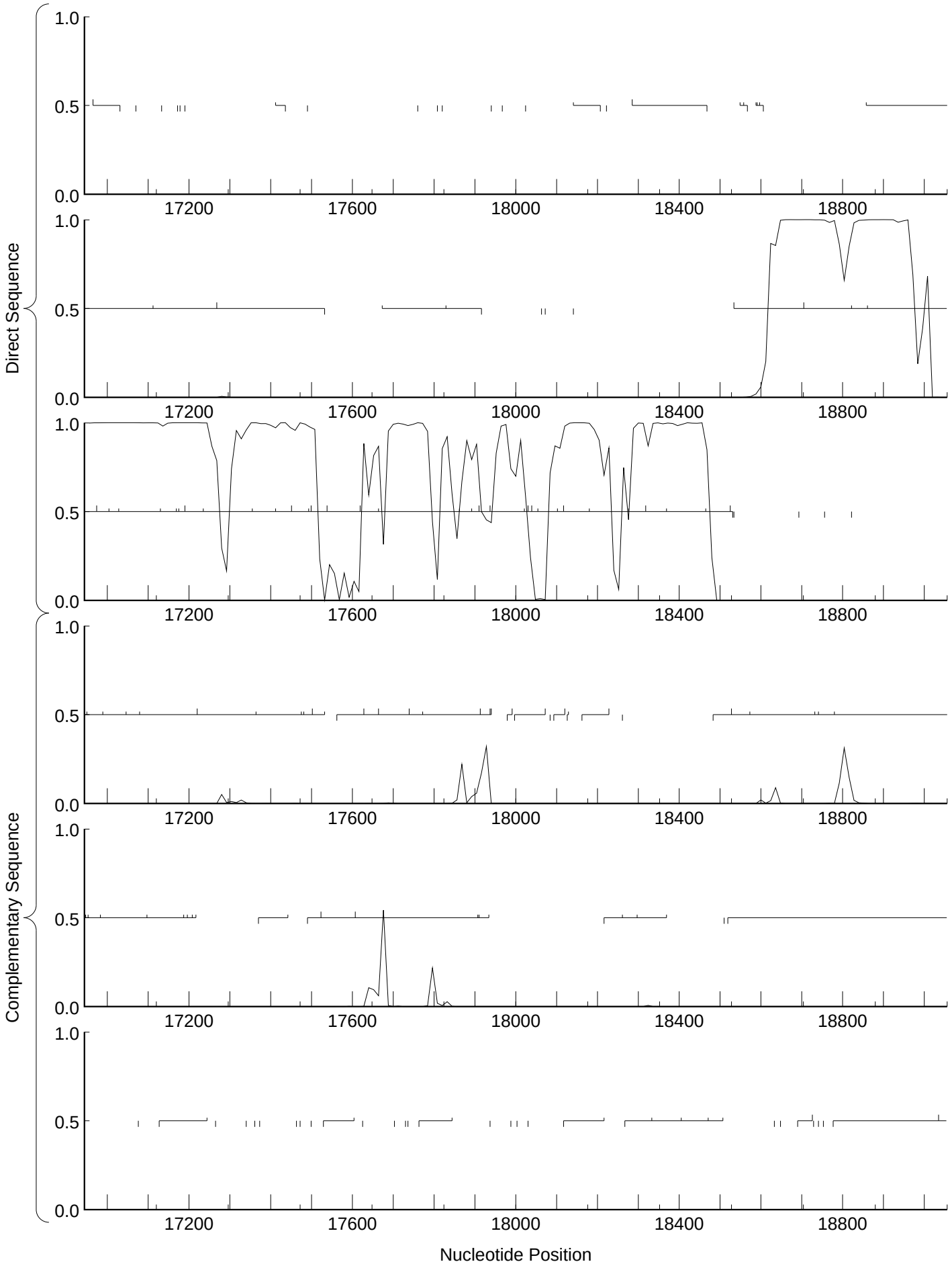
complete sequence, 42311 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 8/22



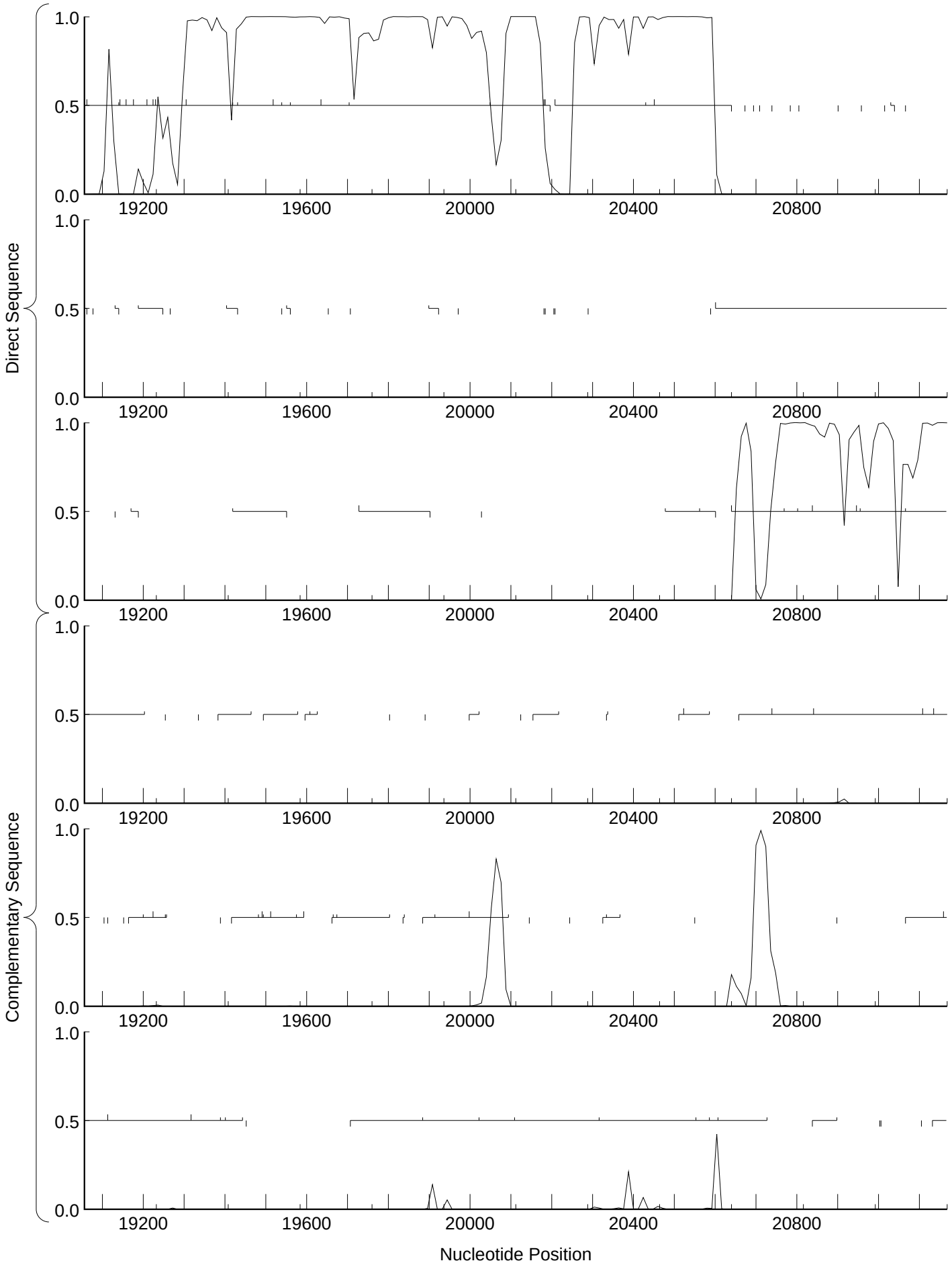
complete sequence, 42311 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 9/22



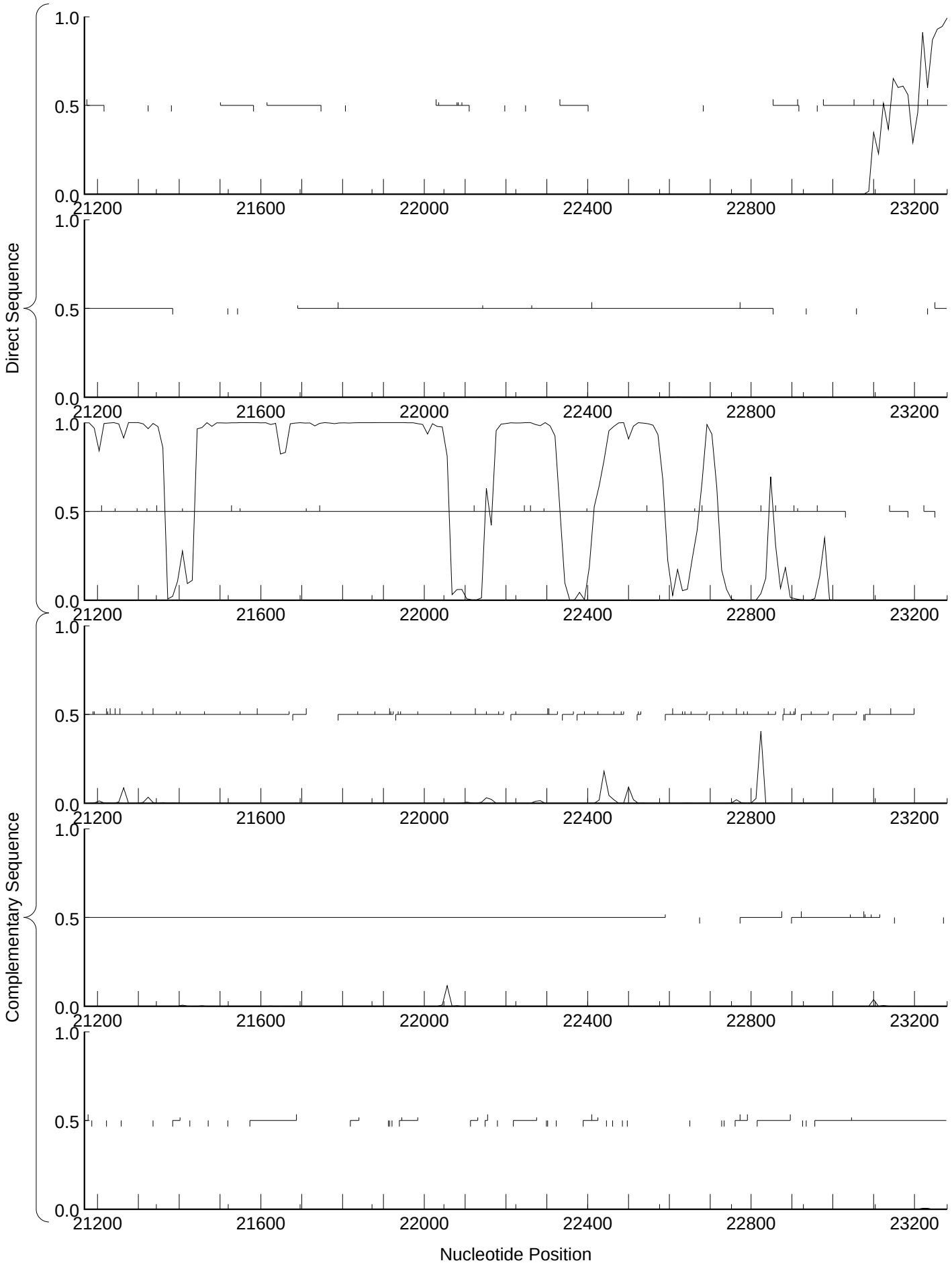
complete sequence, 42311 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 10/22



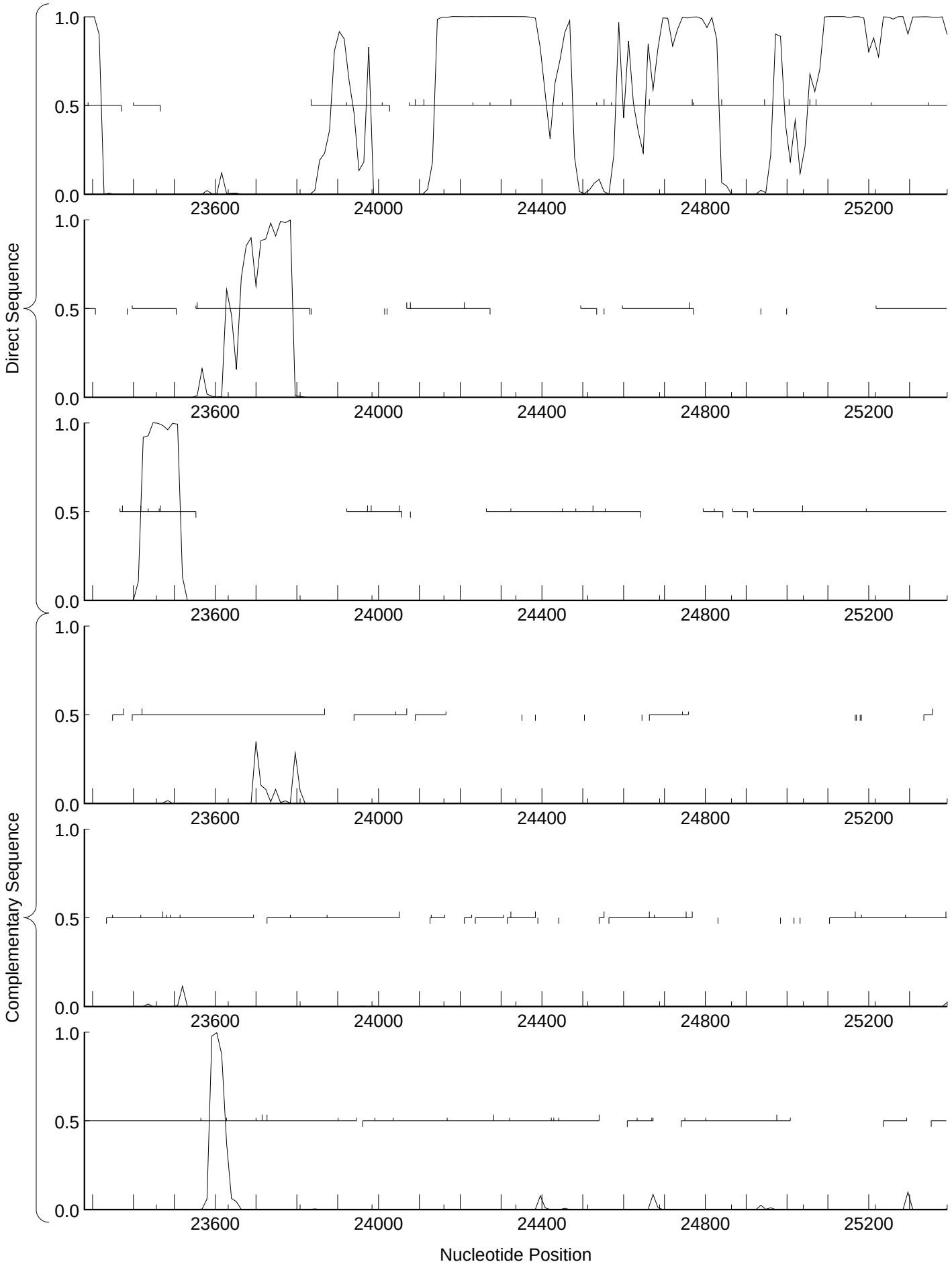
complete sequence, 42311 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 11/22



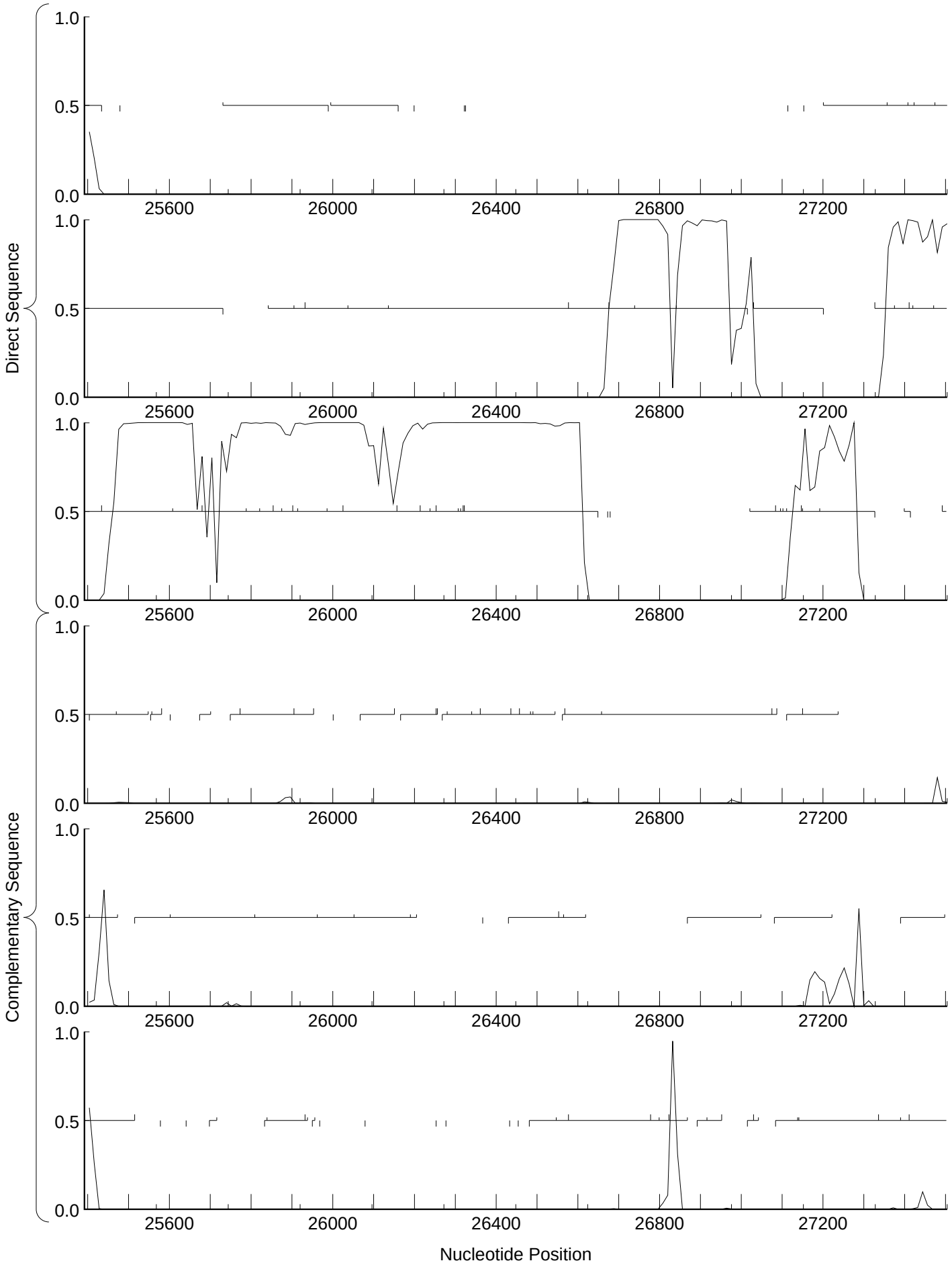
complete sequence, 42311 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 12/22



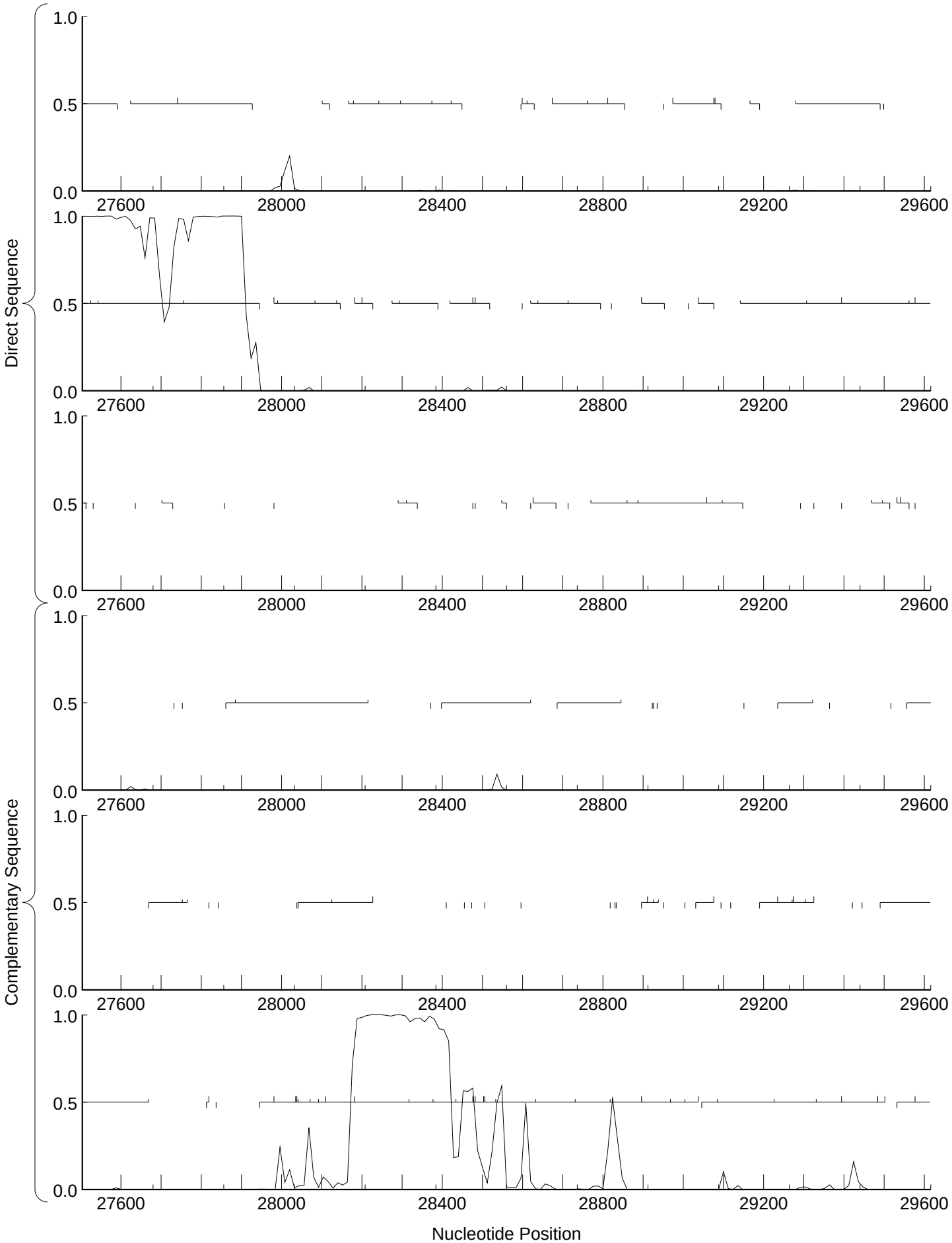
complete sequence, 42311 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 13/22



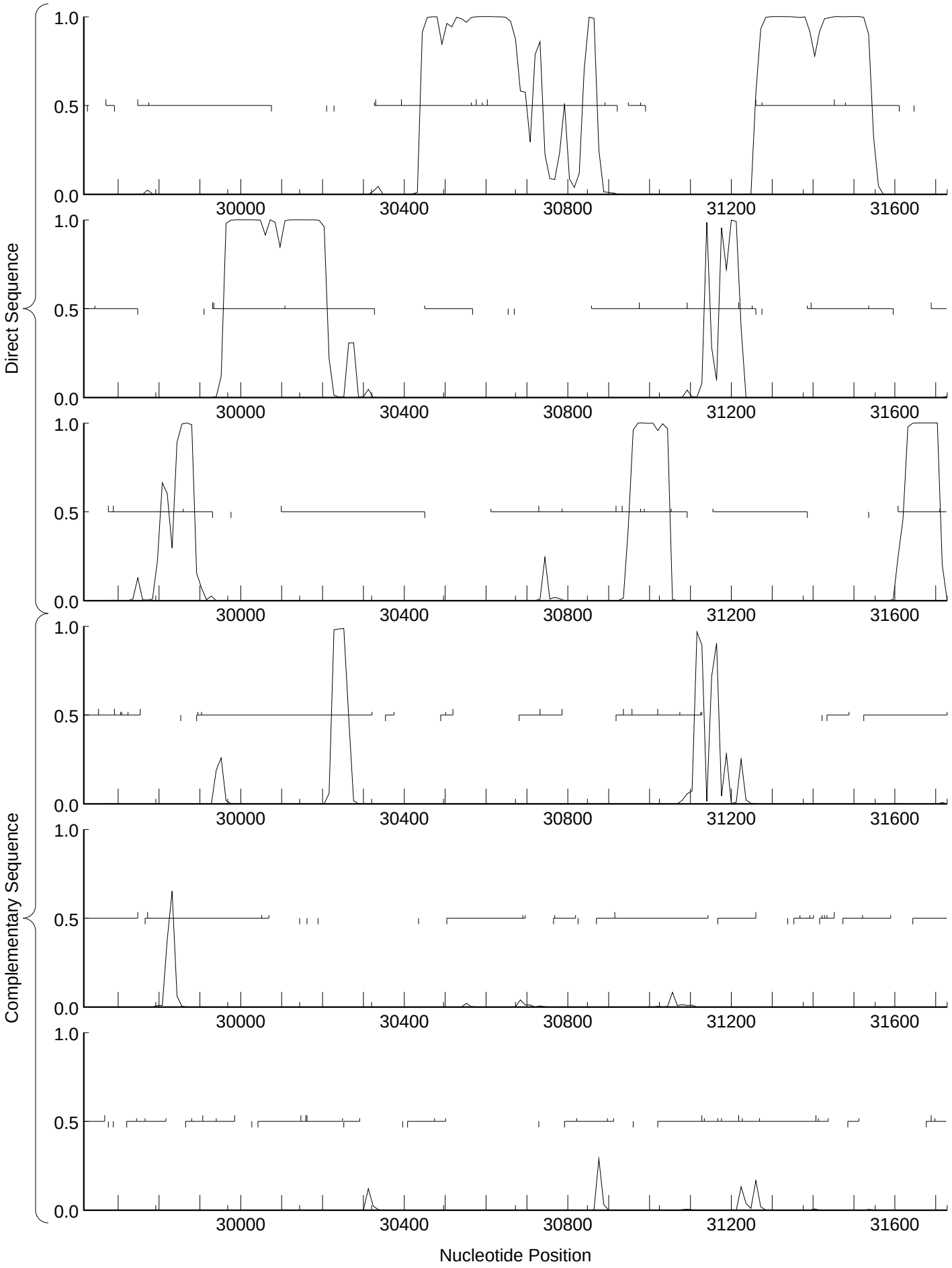
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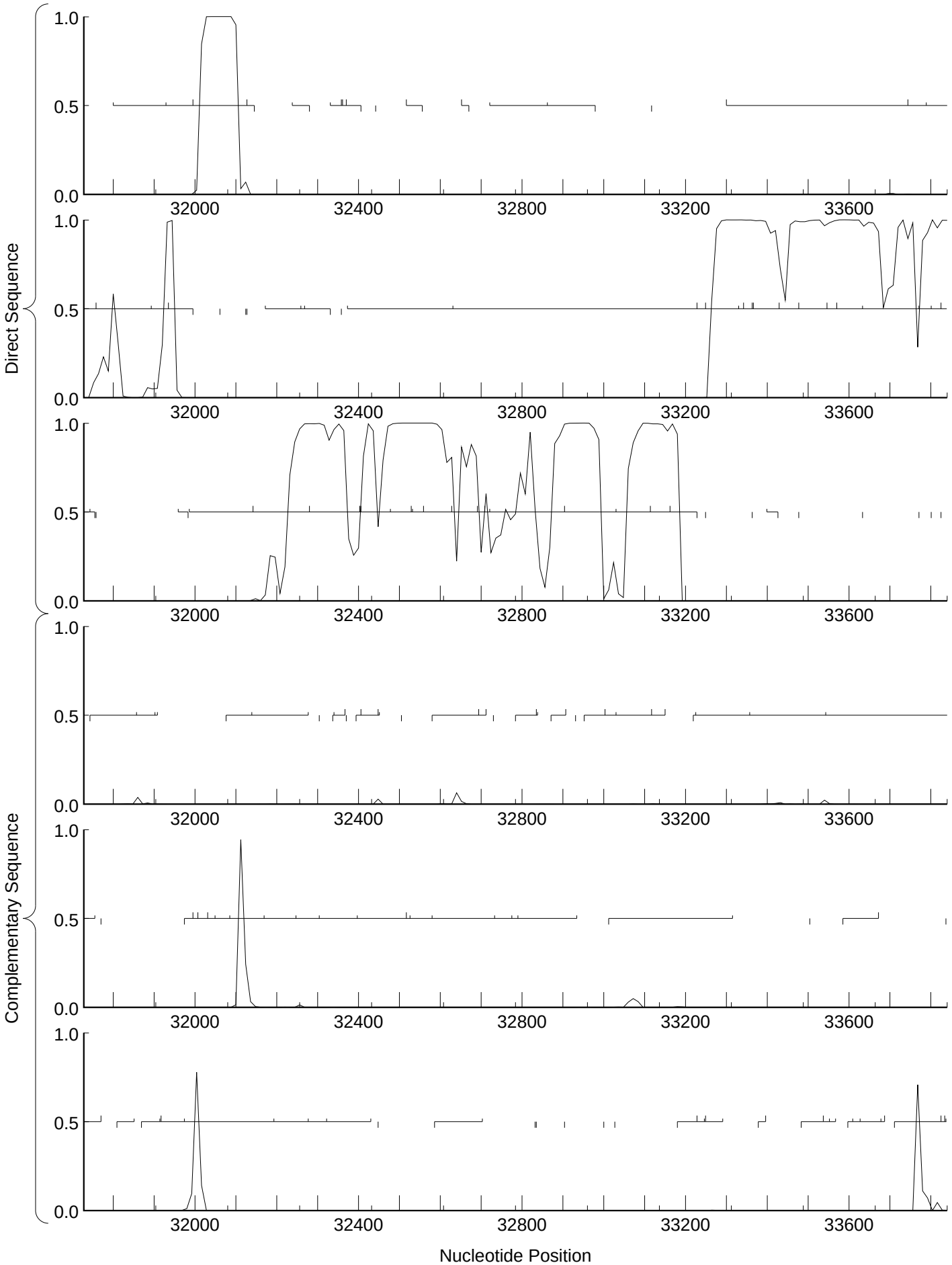
complete sequence, 42311 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 15/22

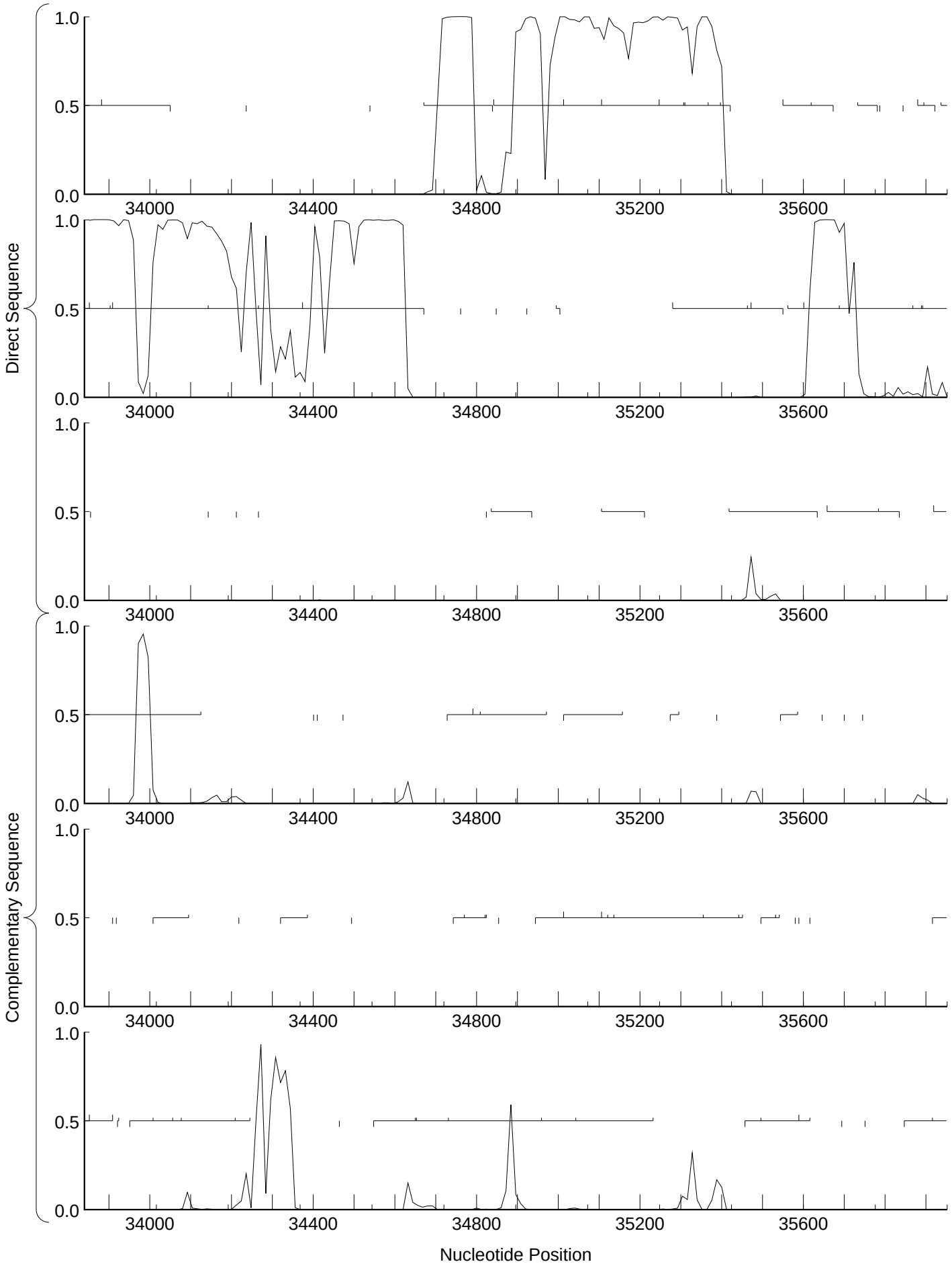


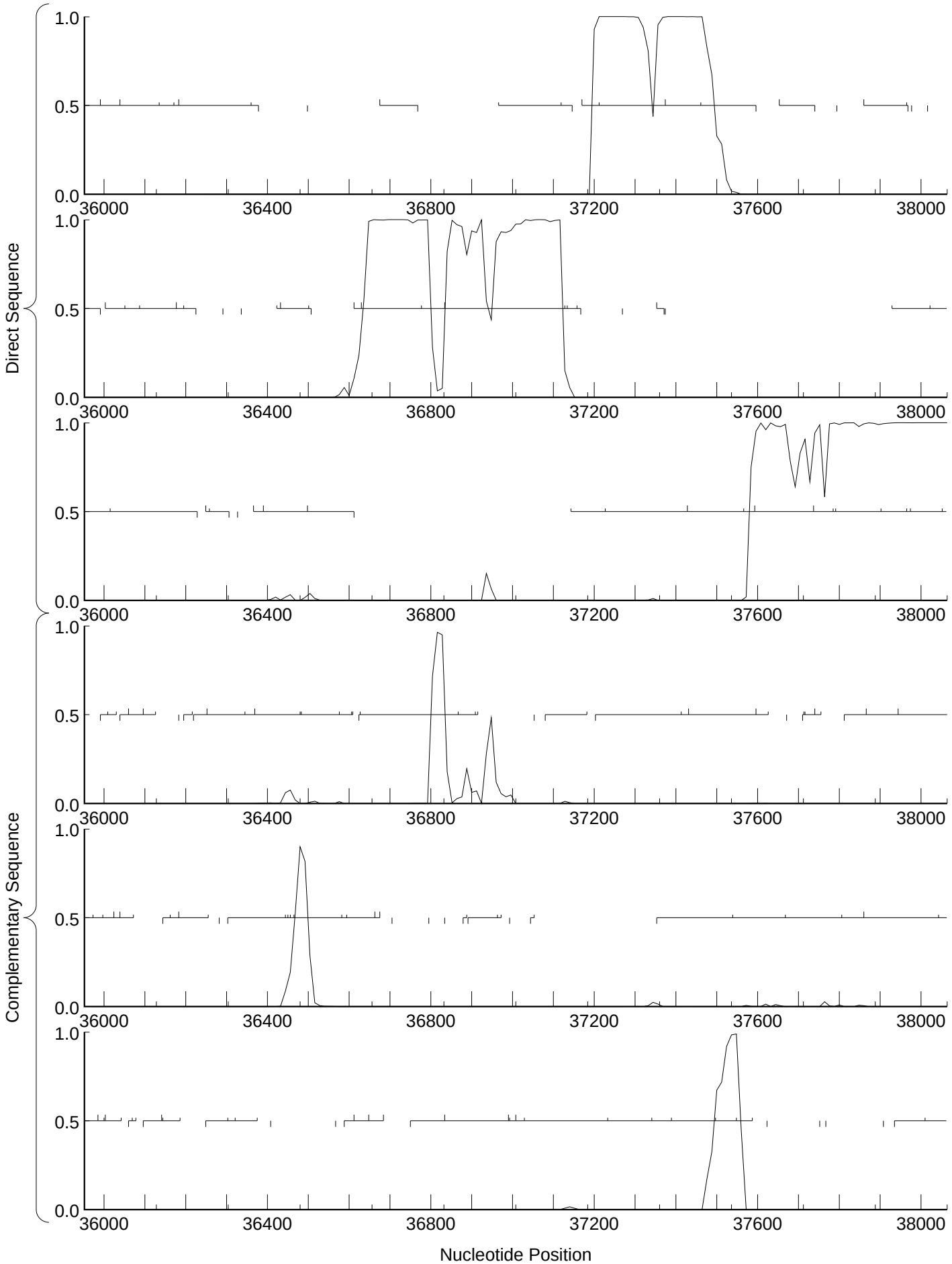
complete sequence, 42311 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 16/22



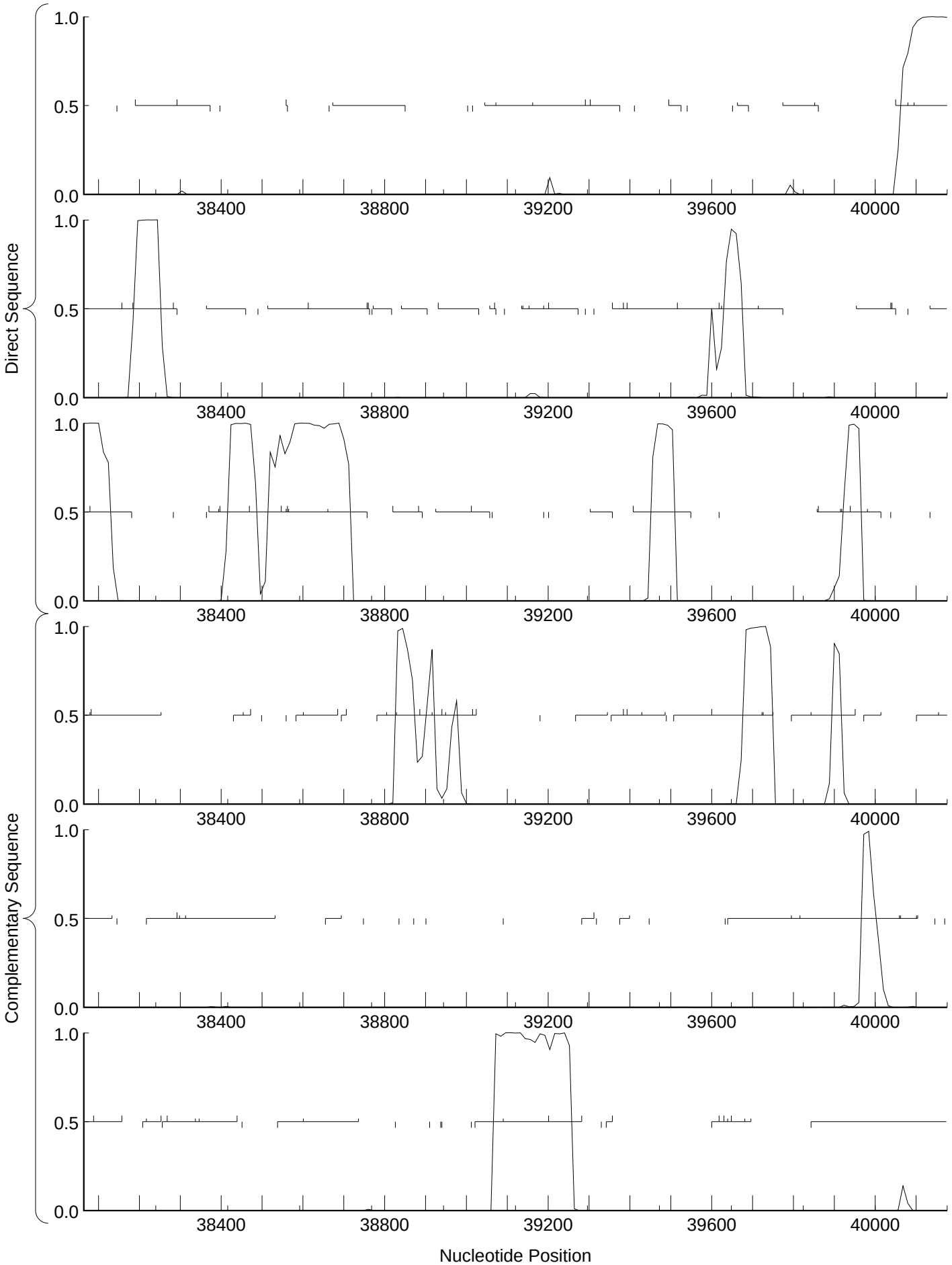
complete sequence, 42311 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 17/22




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


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
36000
36400
36800
37200
37600
38000
Nucleotide Position

complete sequence, 42311 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 20/22



complete sequence, 42311 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G1, Order 4, Window 96, Step 12, 21/22

