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    :  3/29/24 at 11:11:35
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

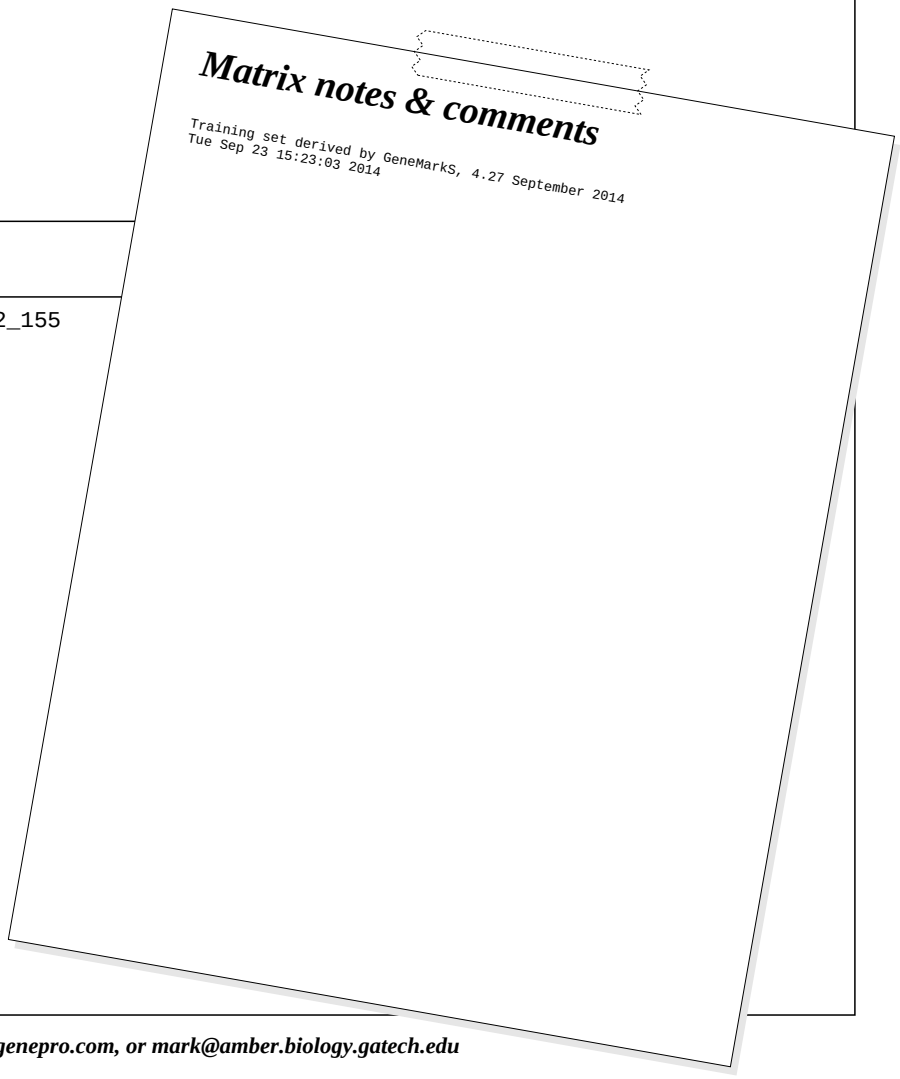
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
```

MATRIX INFORMATION

```
Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

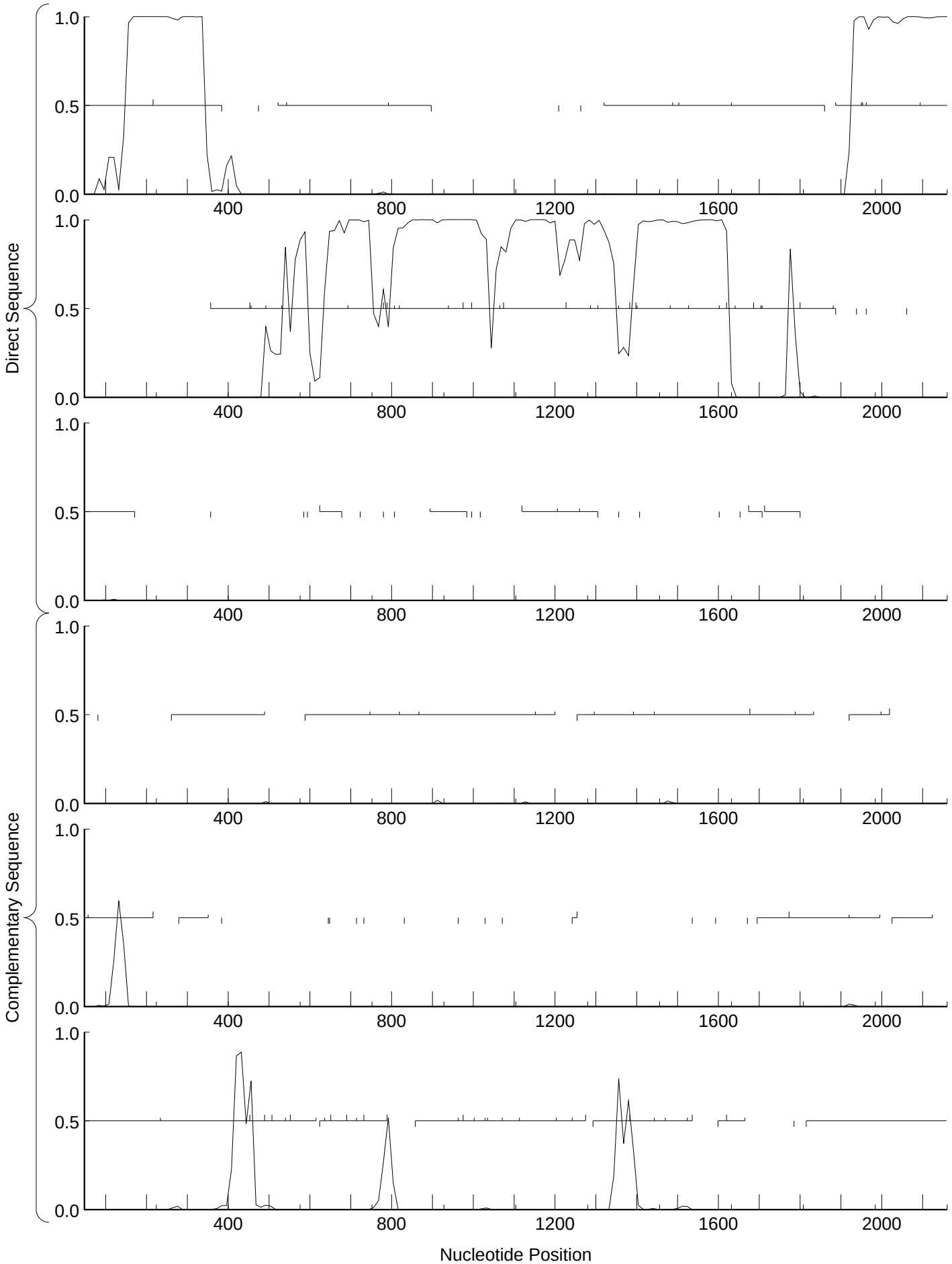
Order      :  4


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

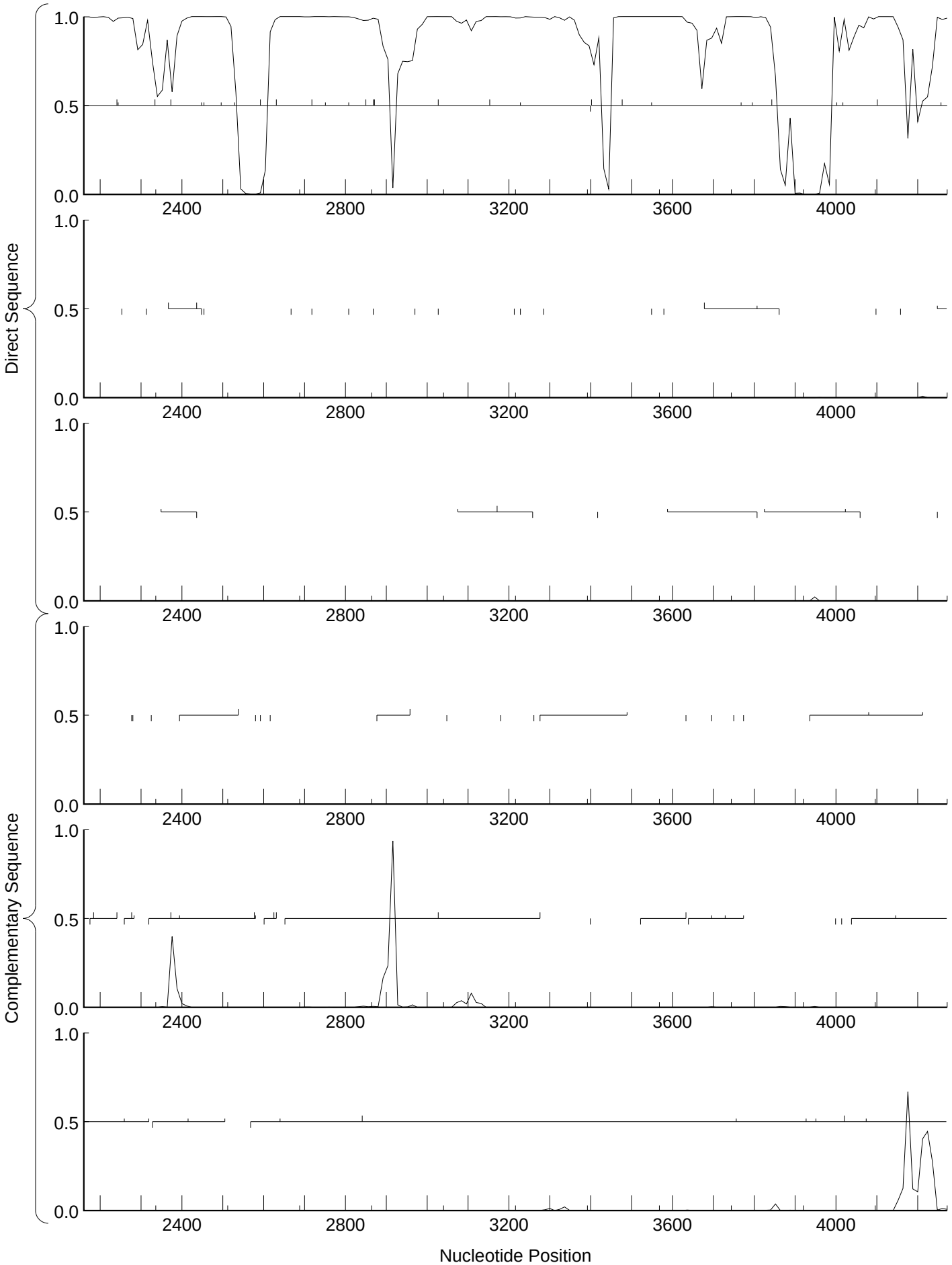


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

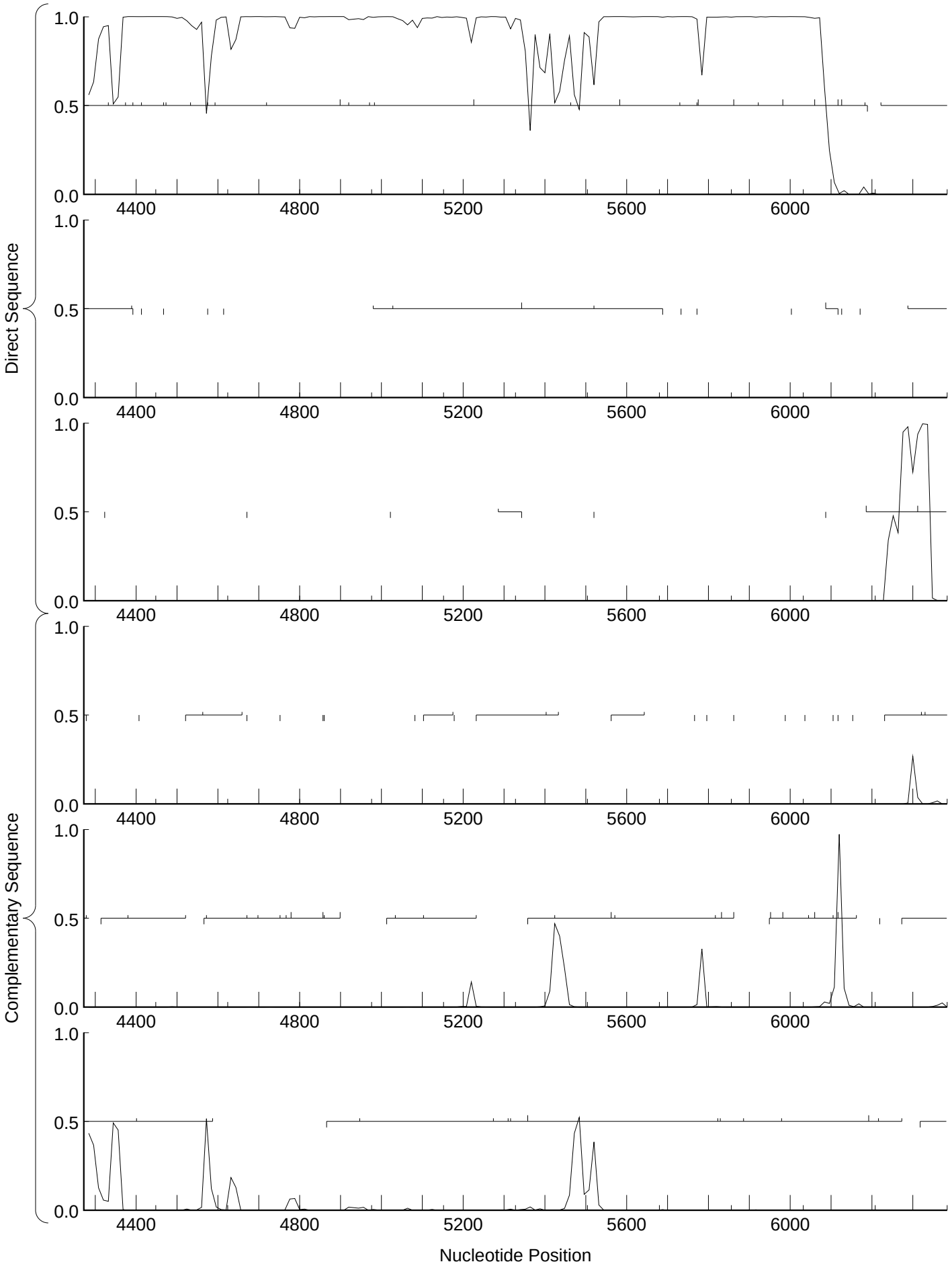
or complete sequence, 41893 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 2/21



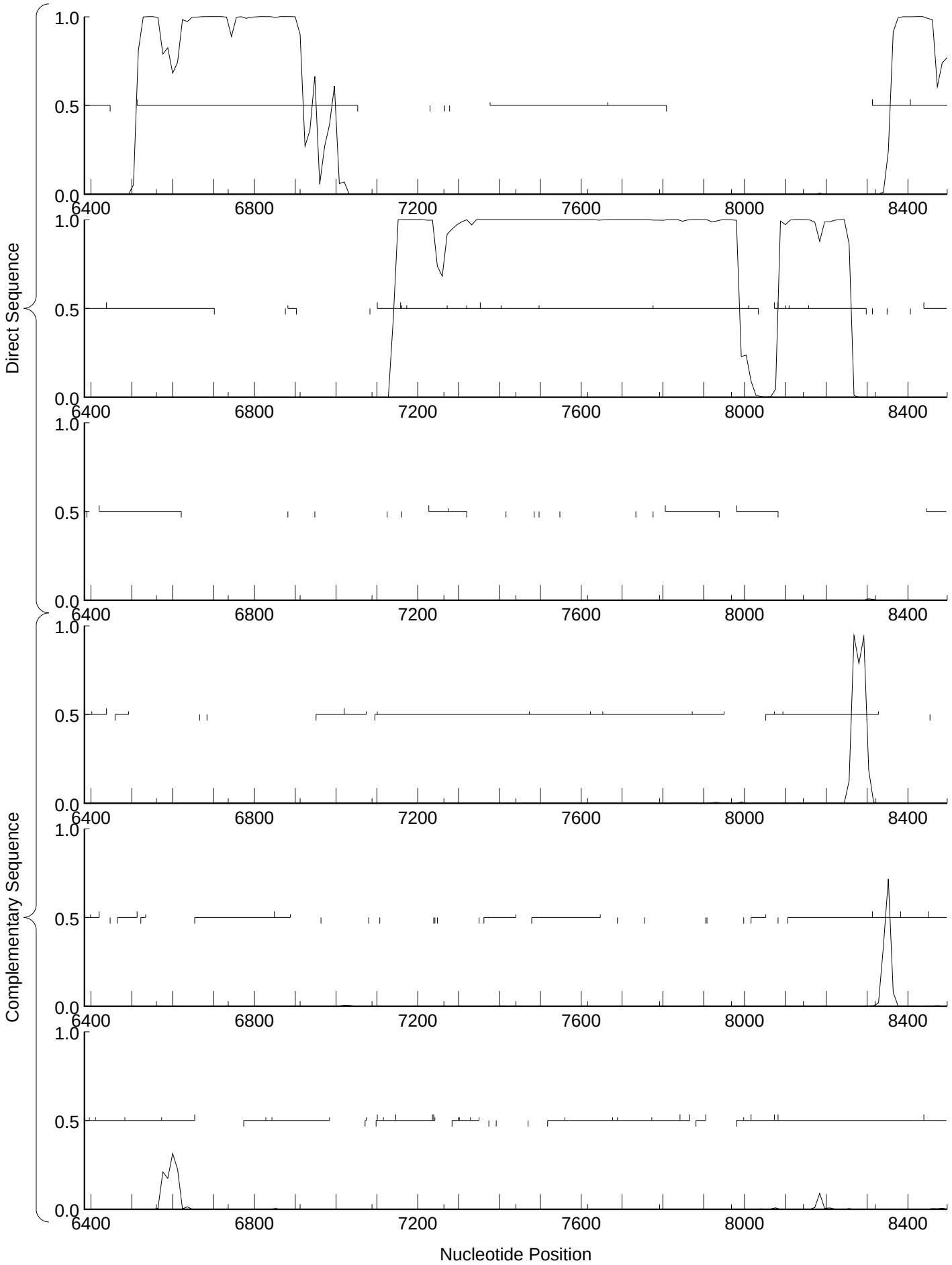
or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 3/21



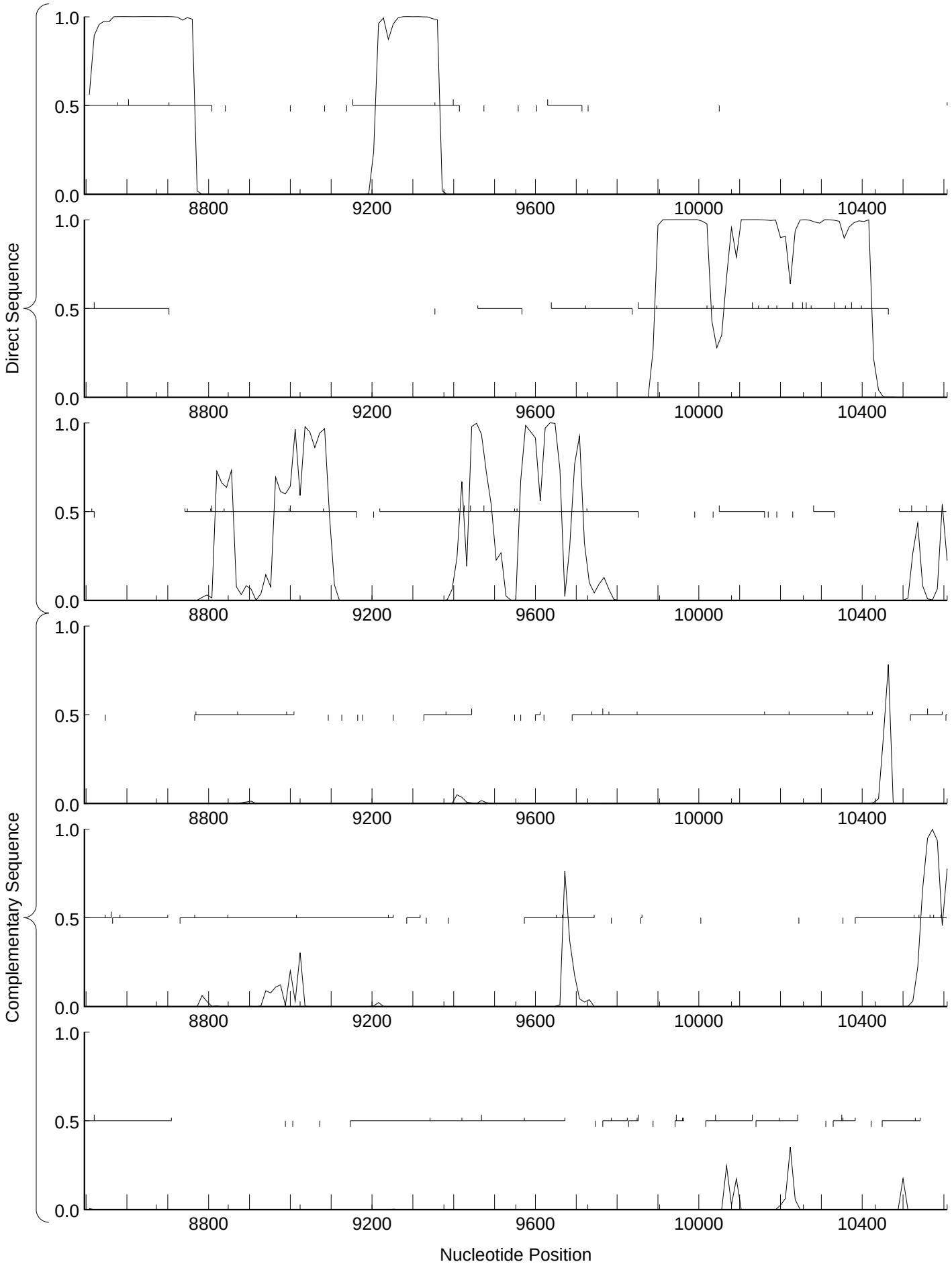
or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 4/21



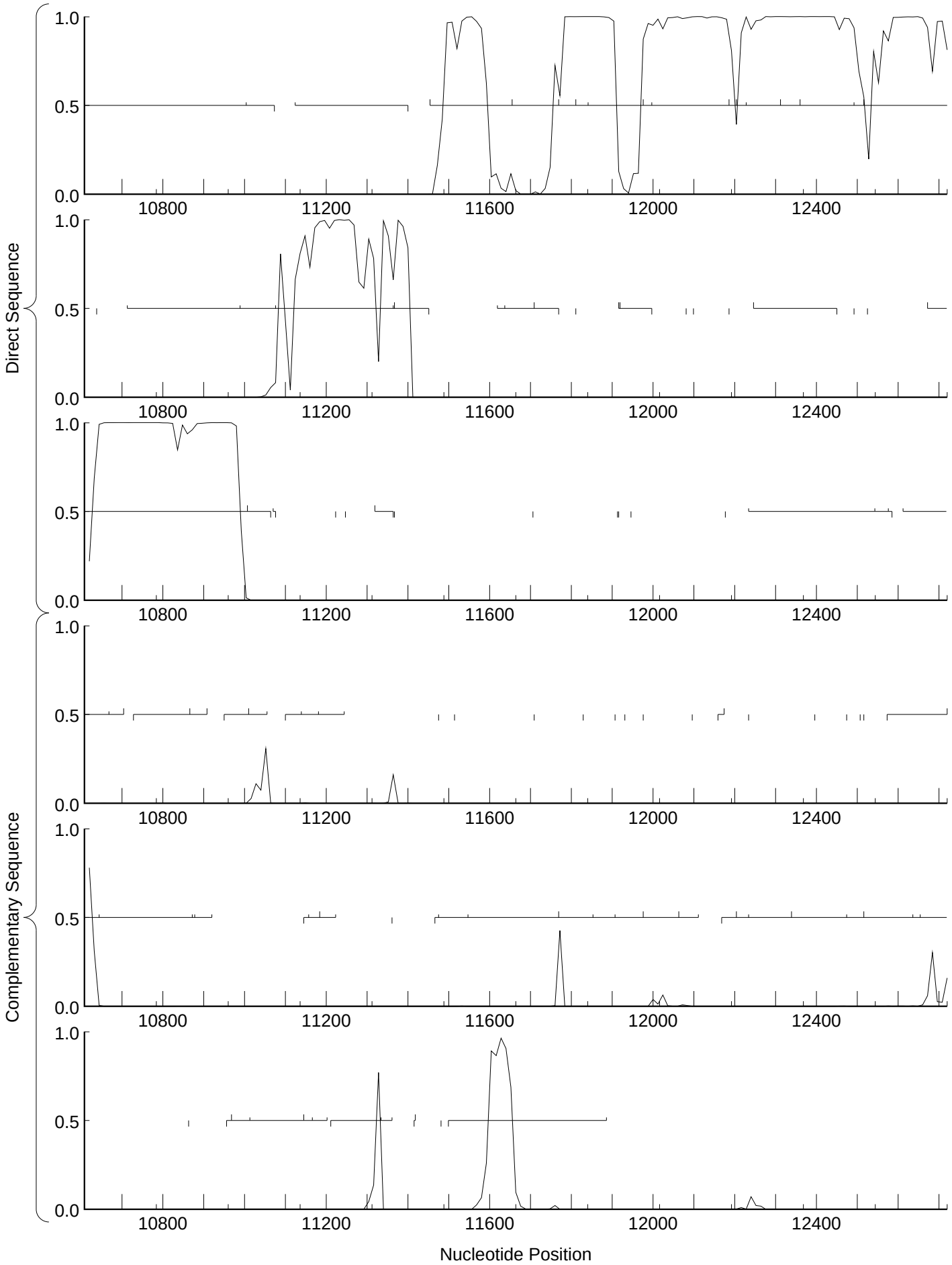
or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 5/21



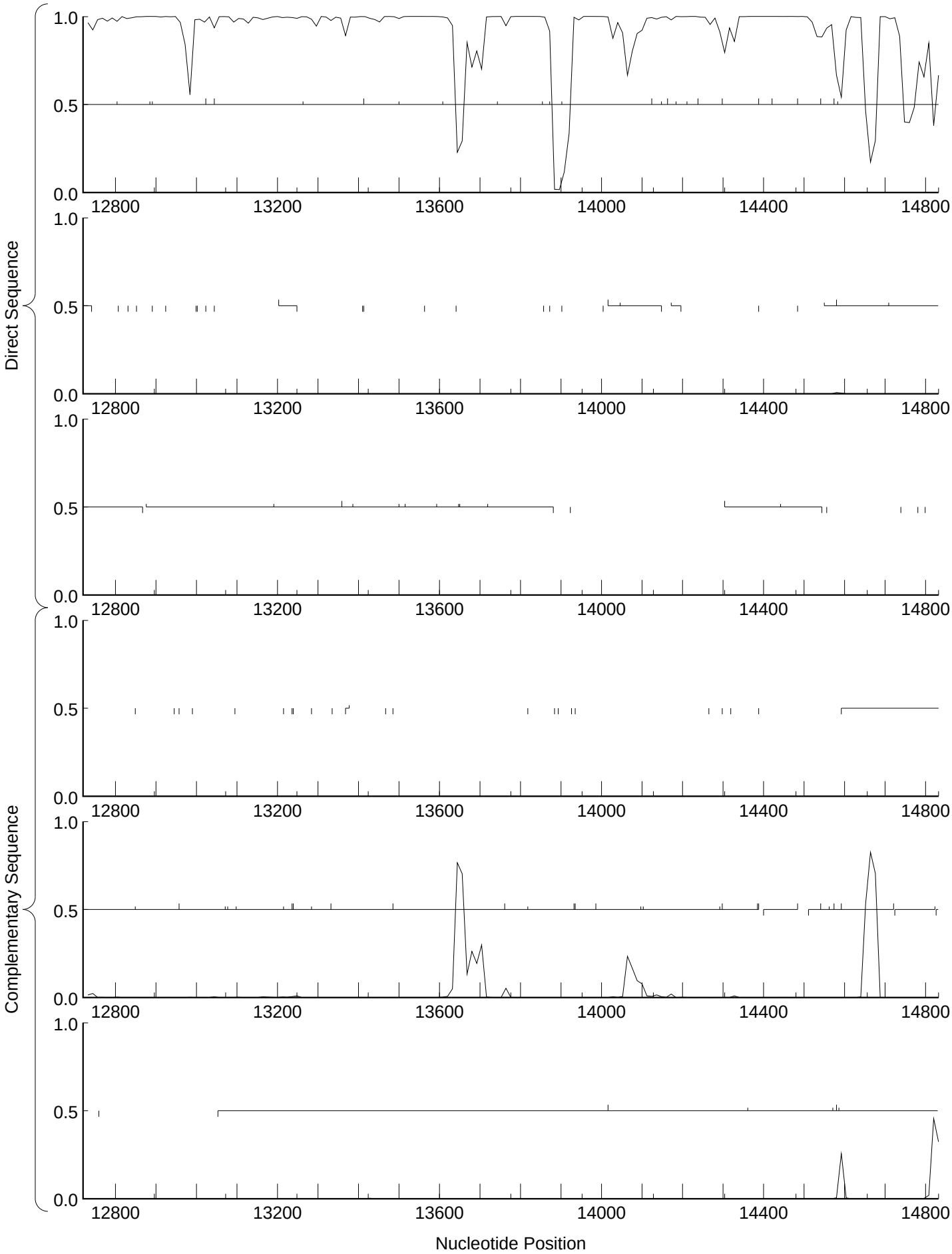
or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 6/21



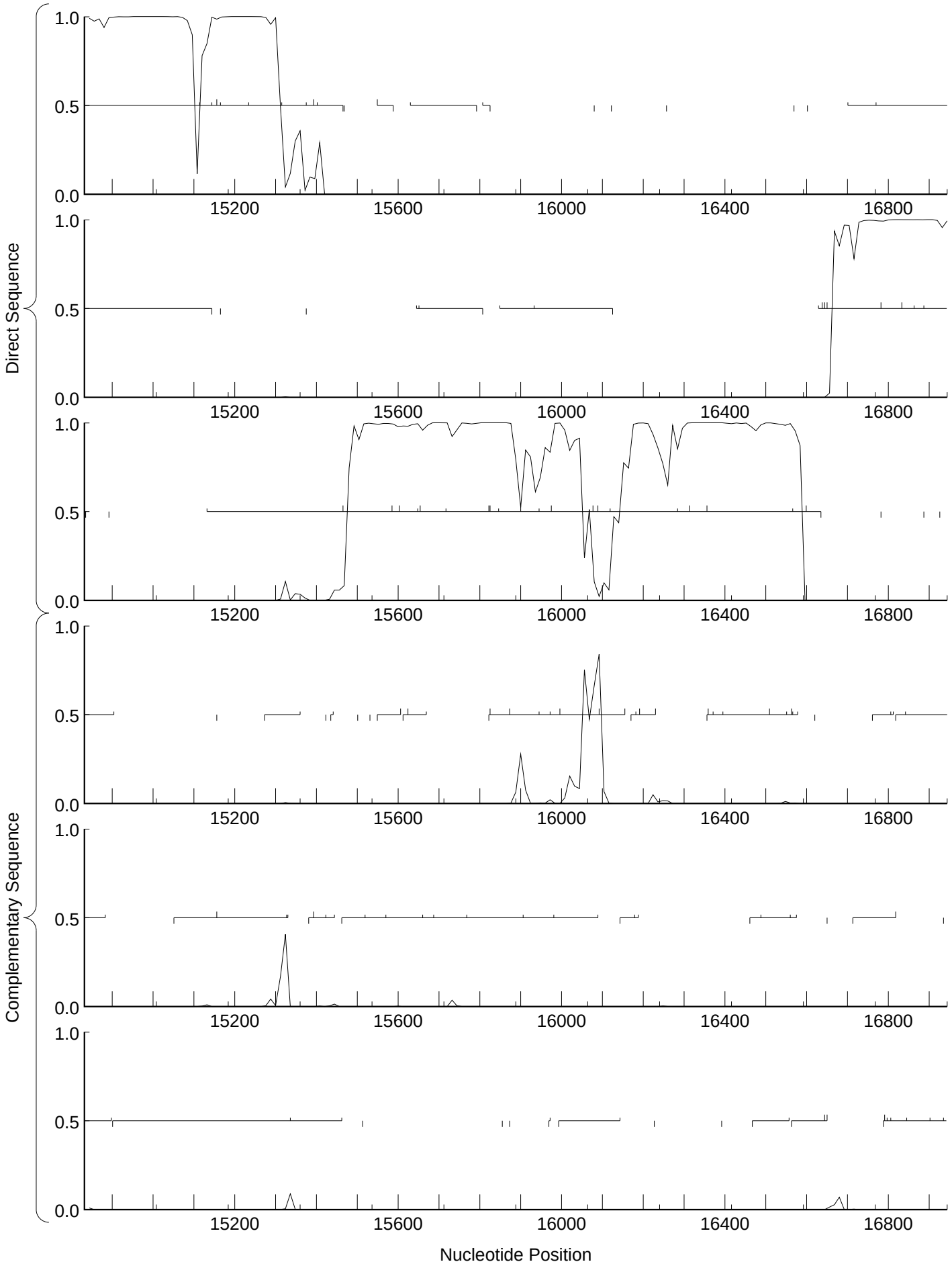
or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 7/21



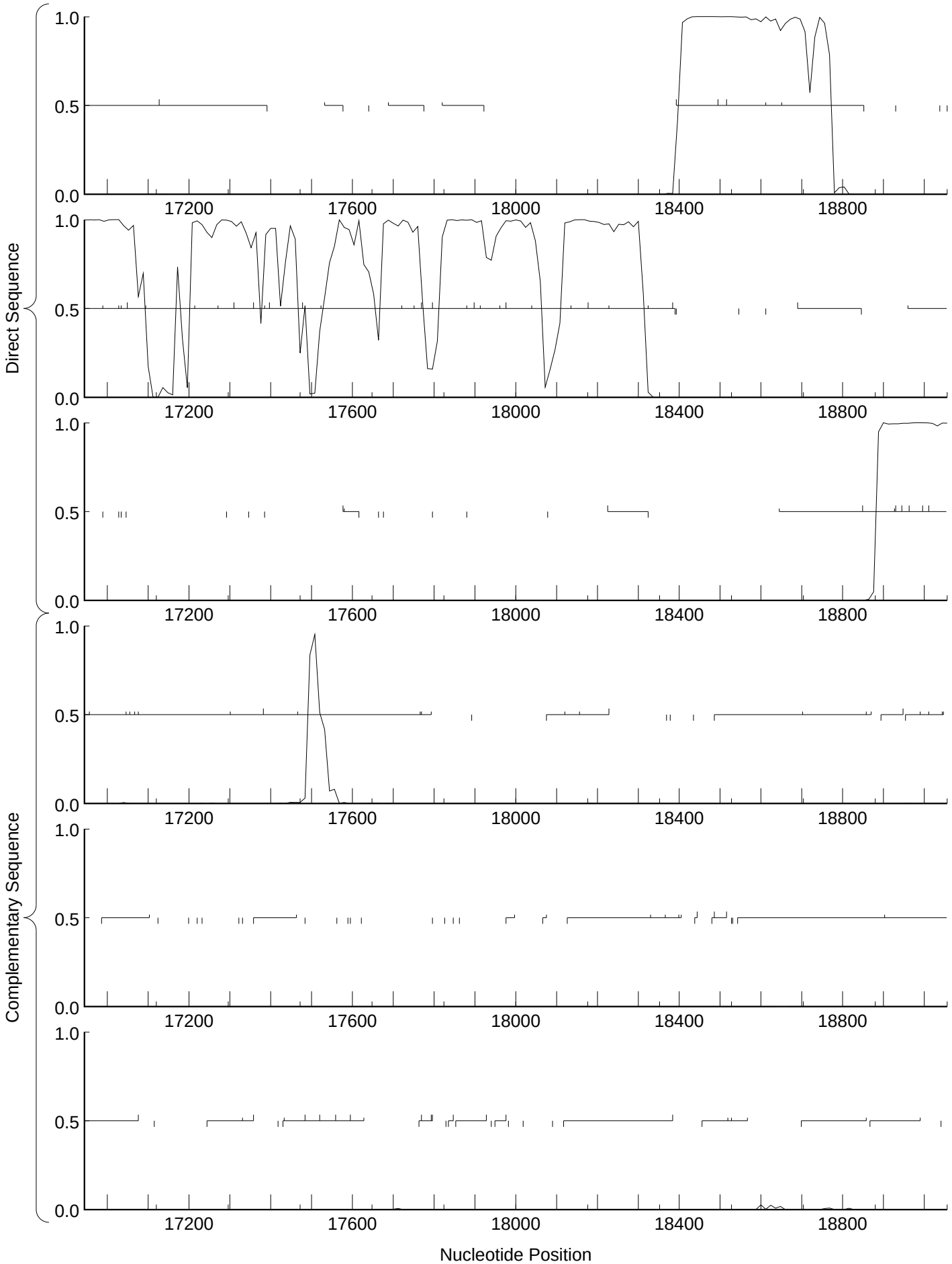
or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 8/21



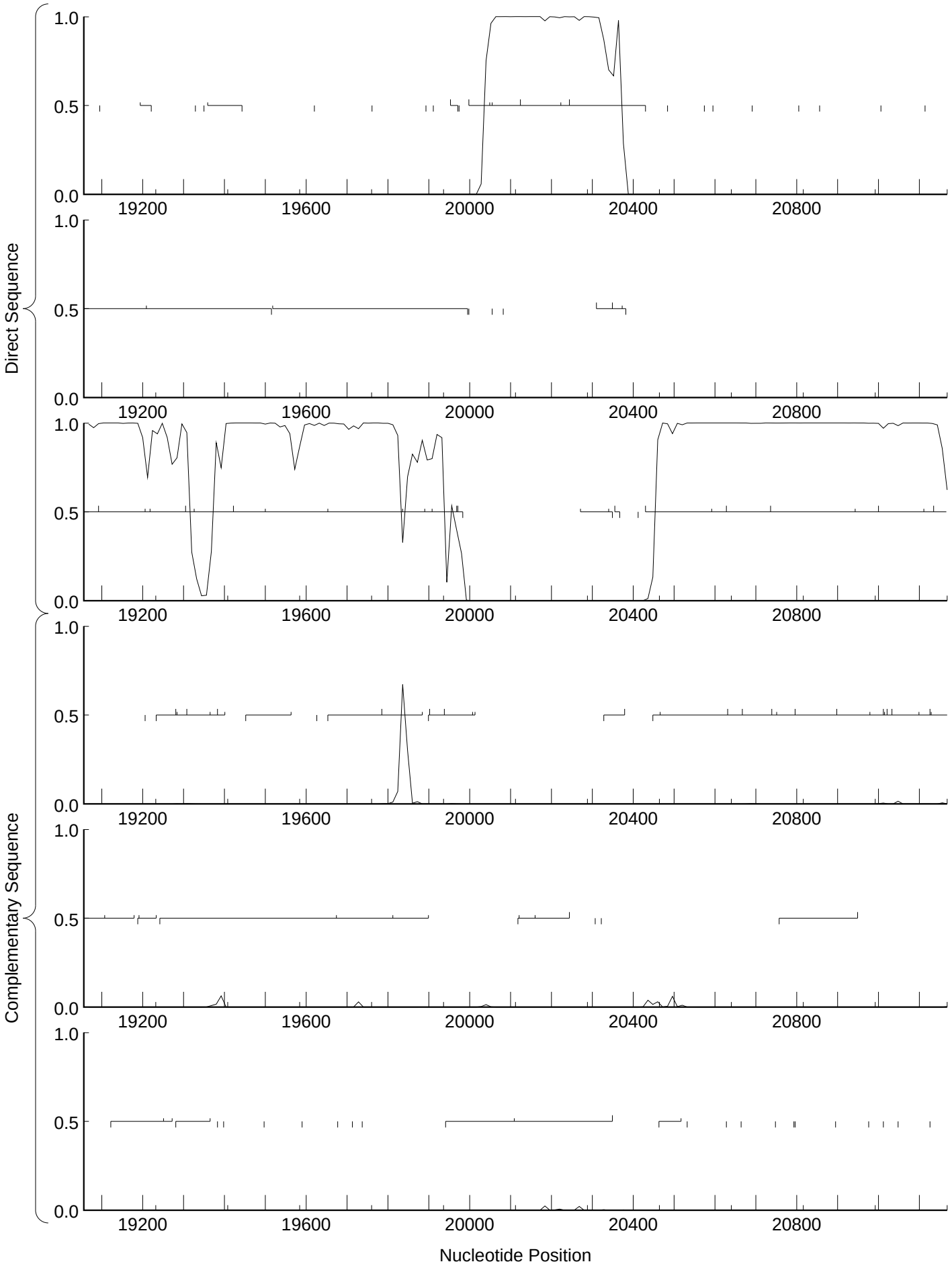
or complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 9/21



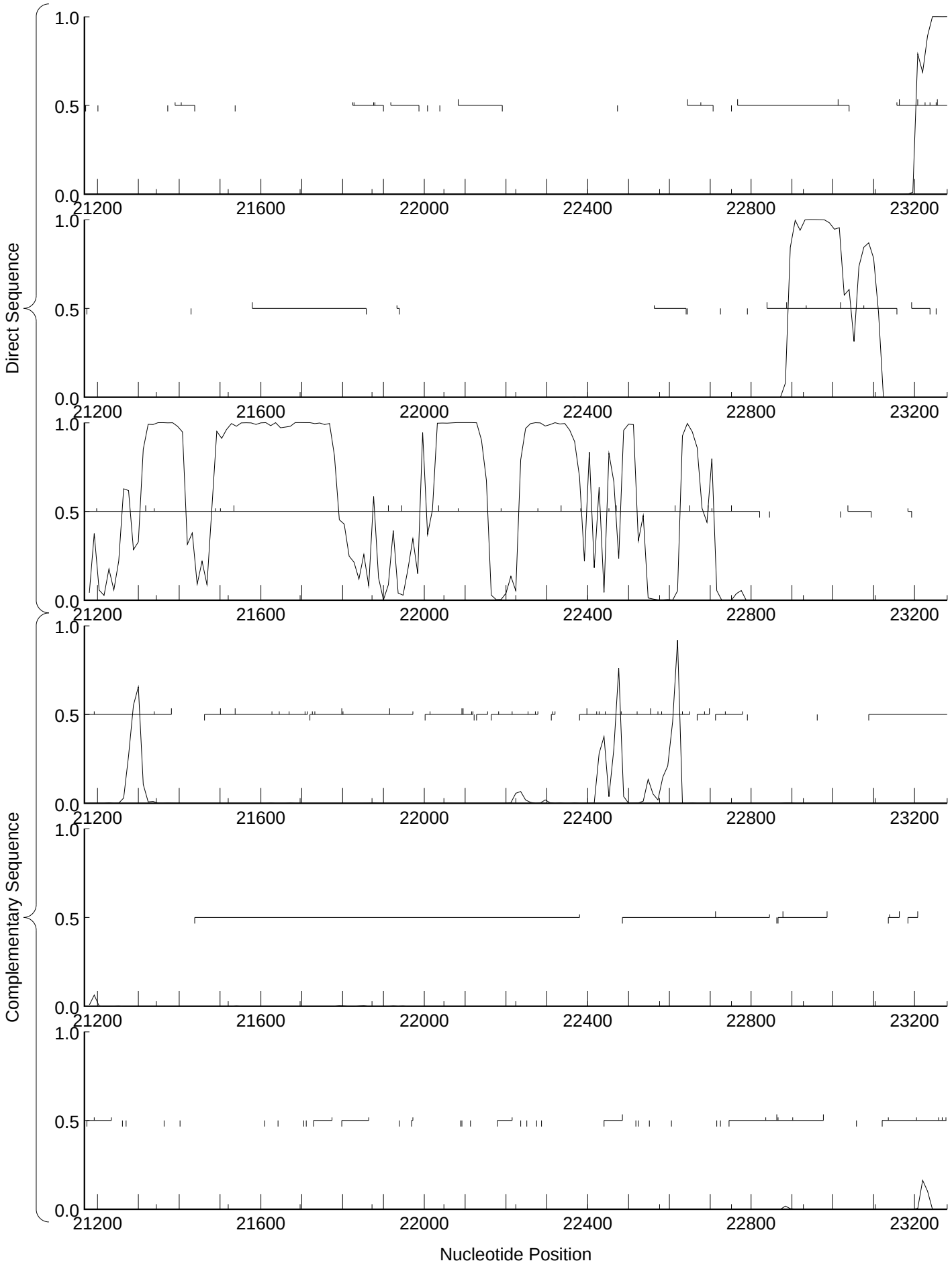
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 10/21



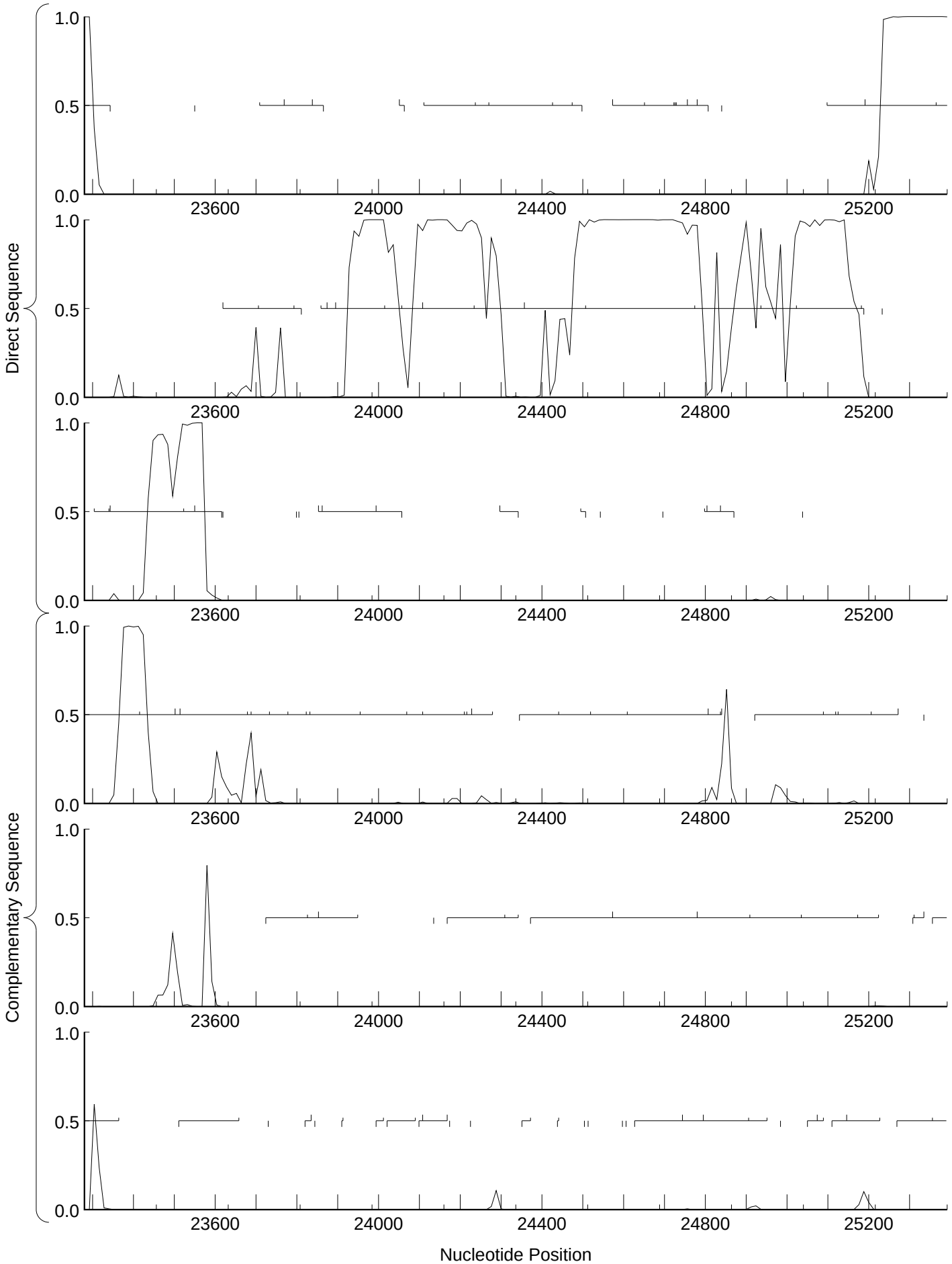
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 11/21



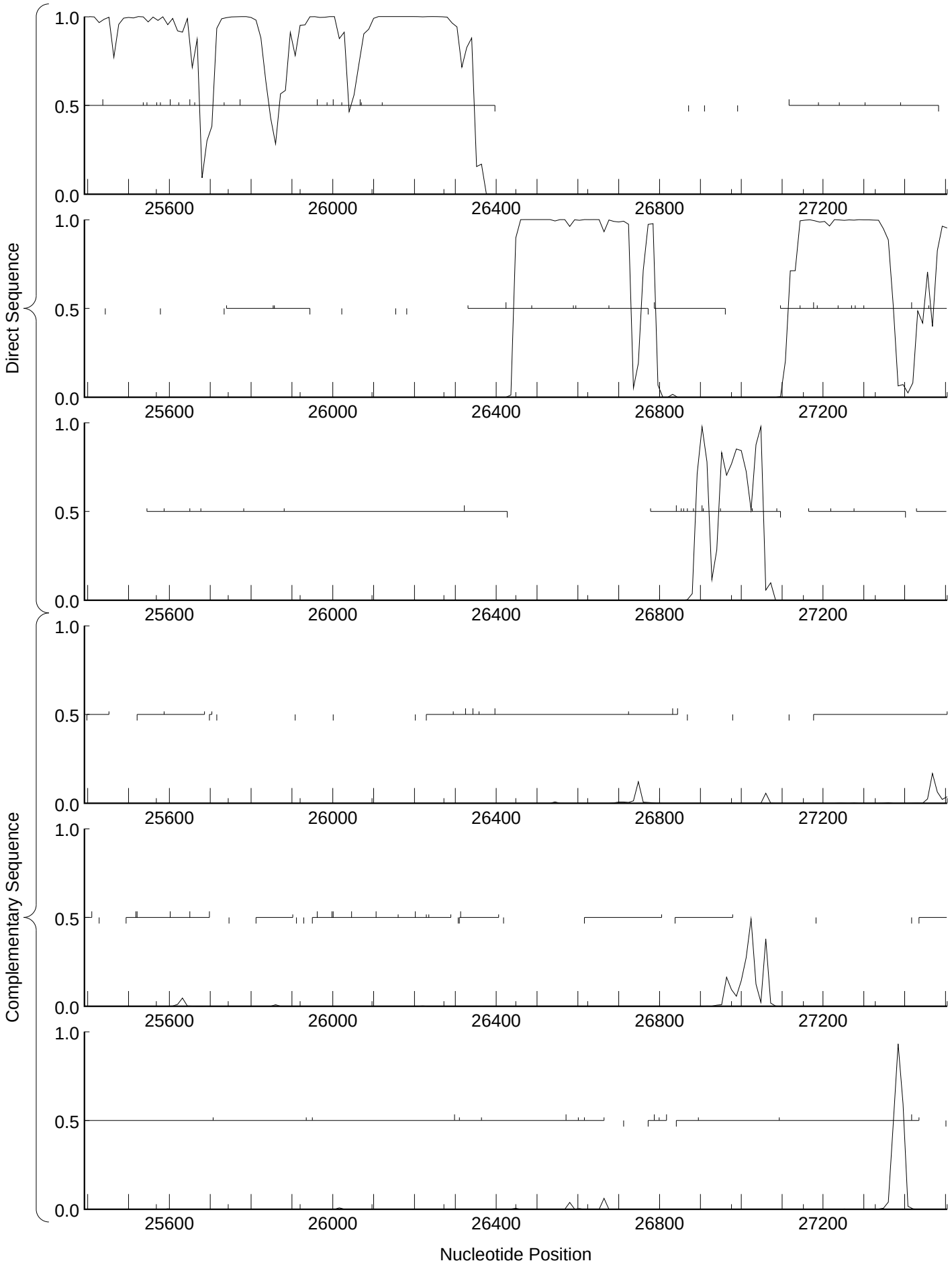
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 12/21



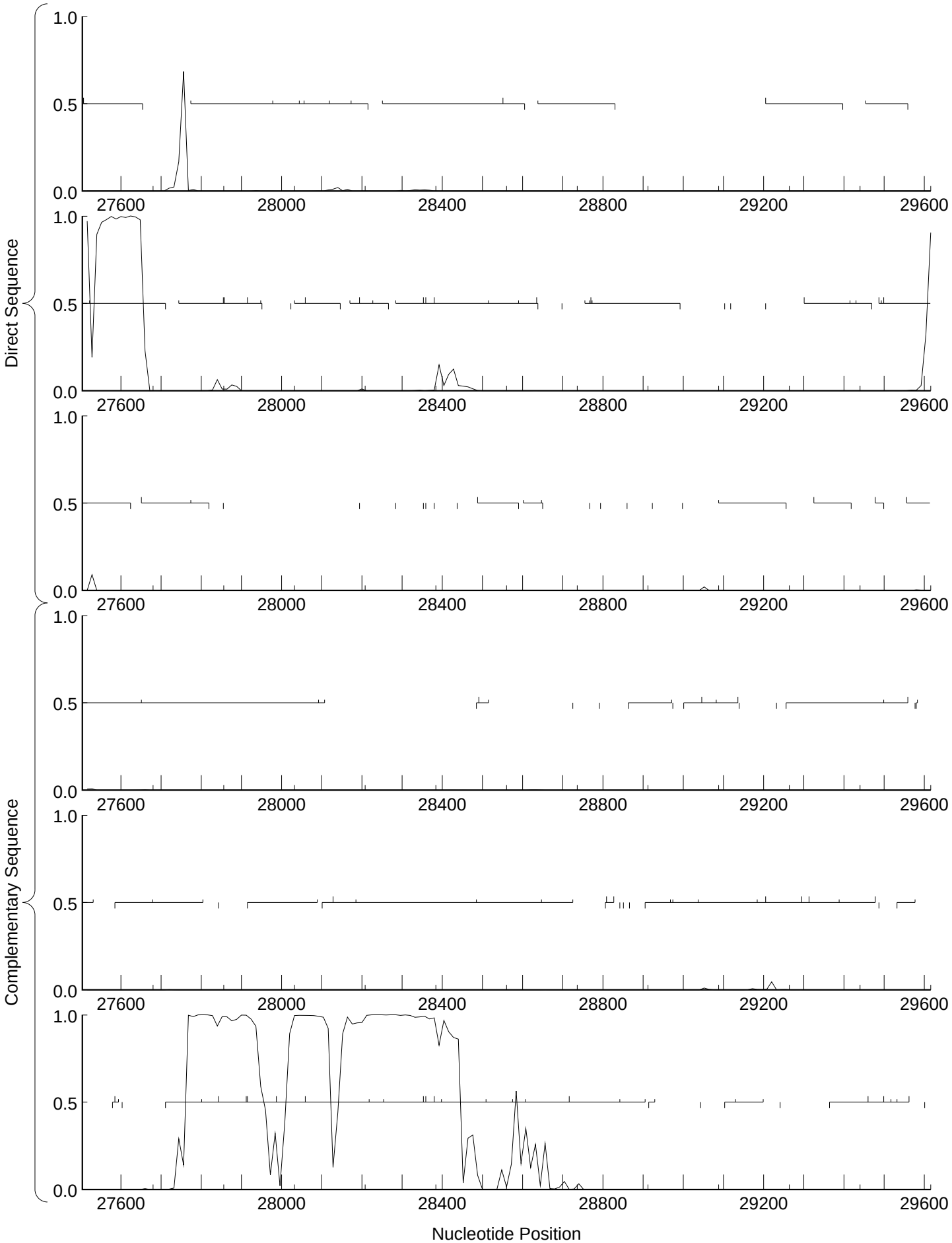
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 13/21



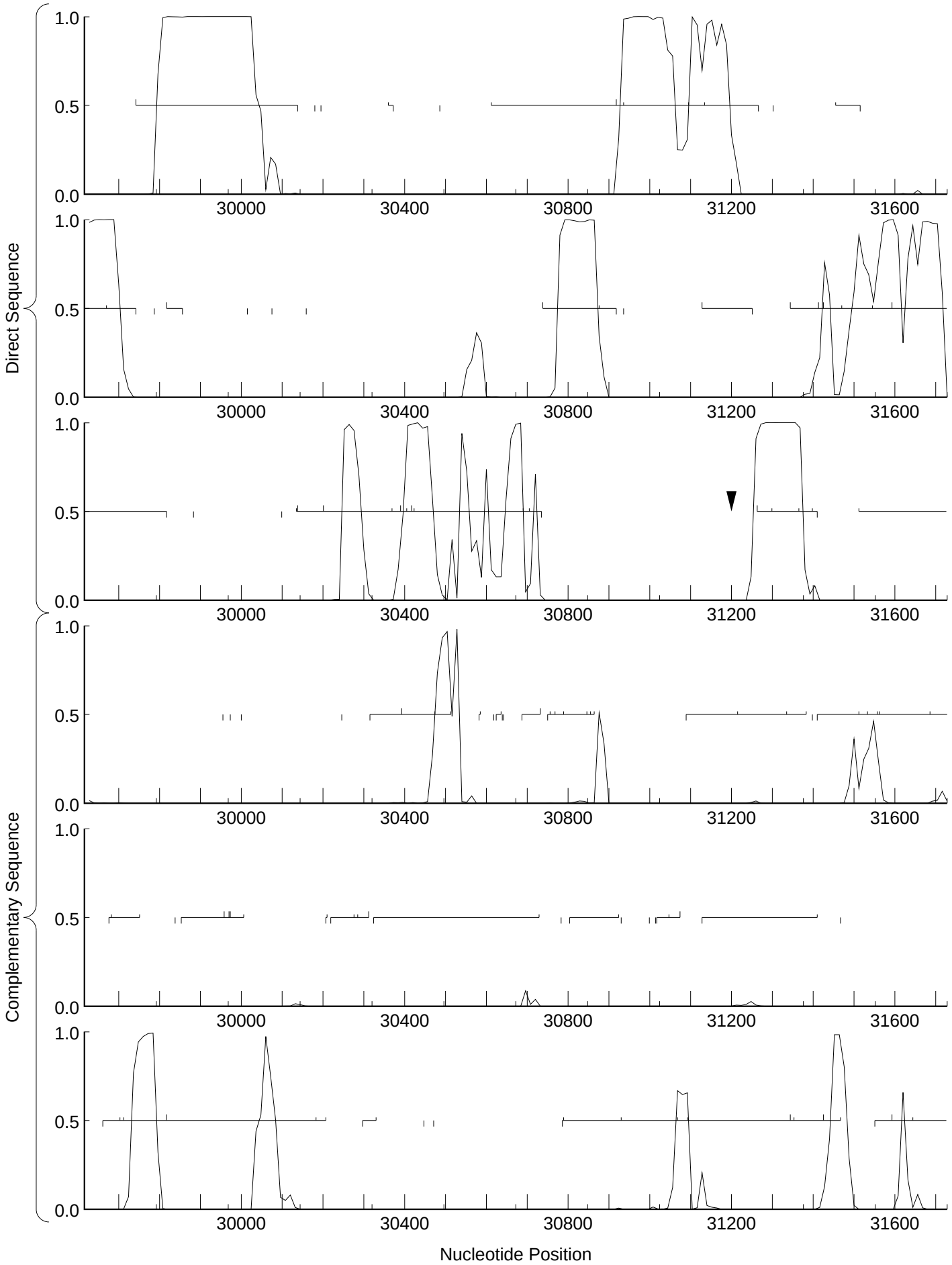
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 14/21



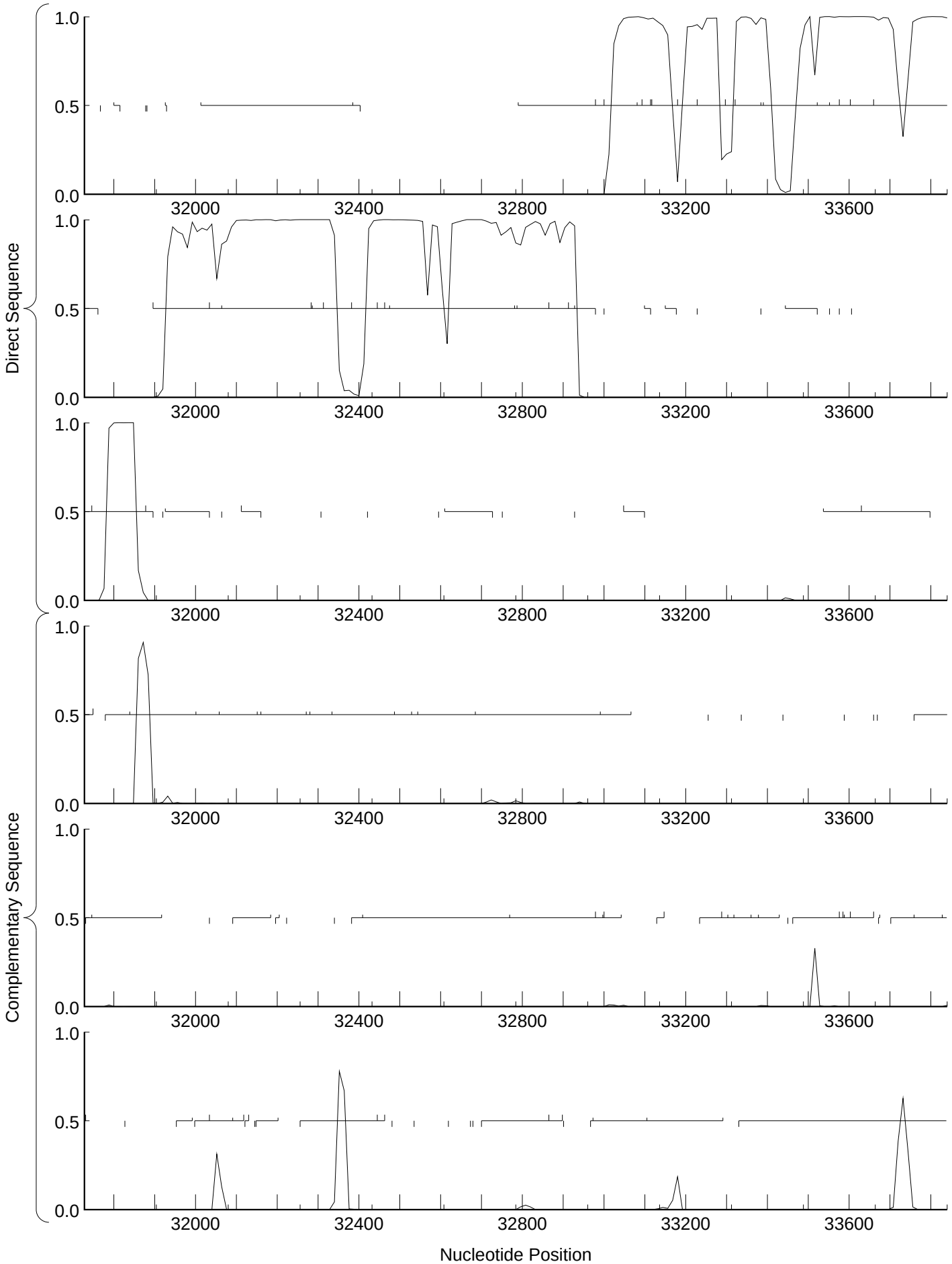
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 15/21



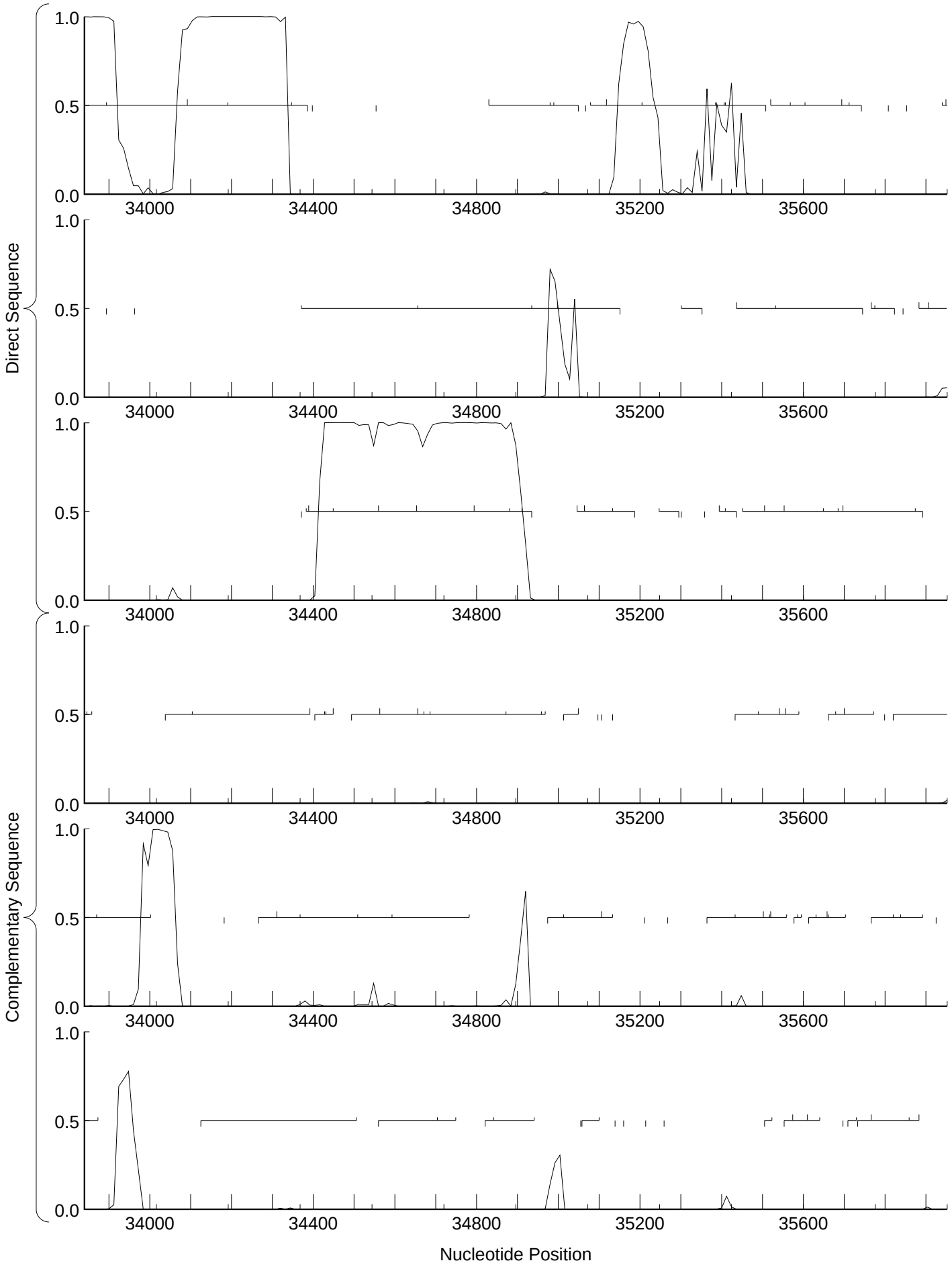
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 16/21



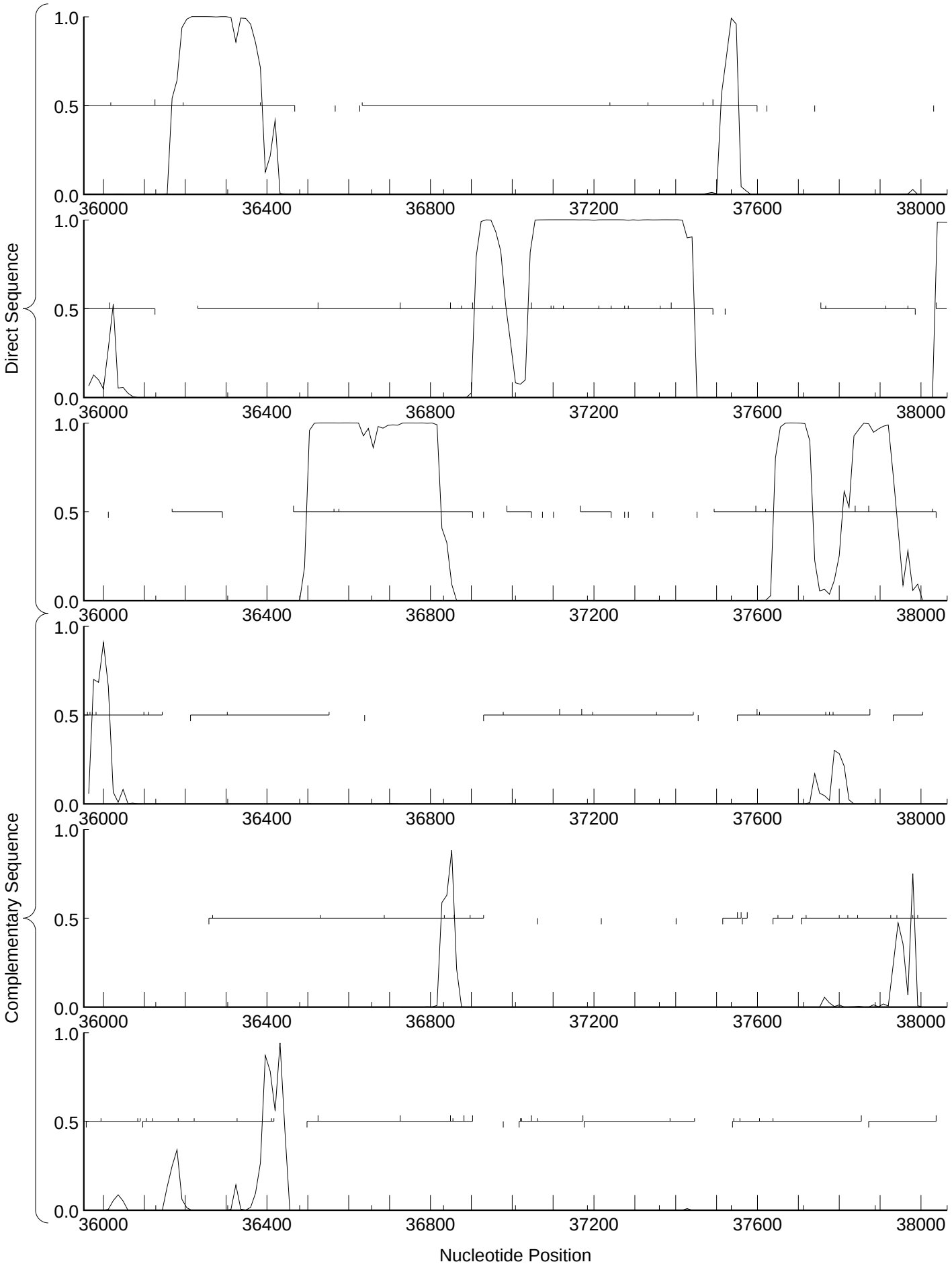
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 17/21



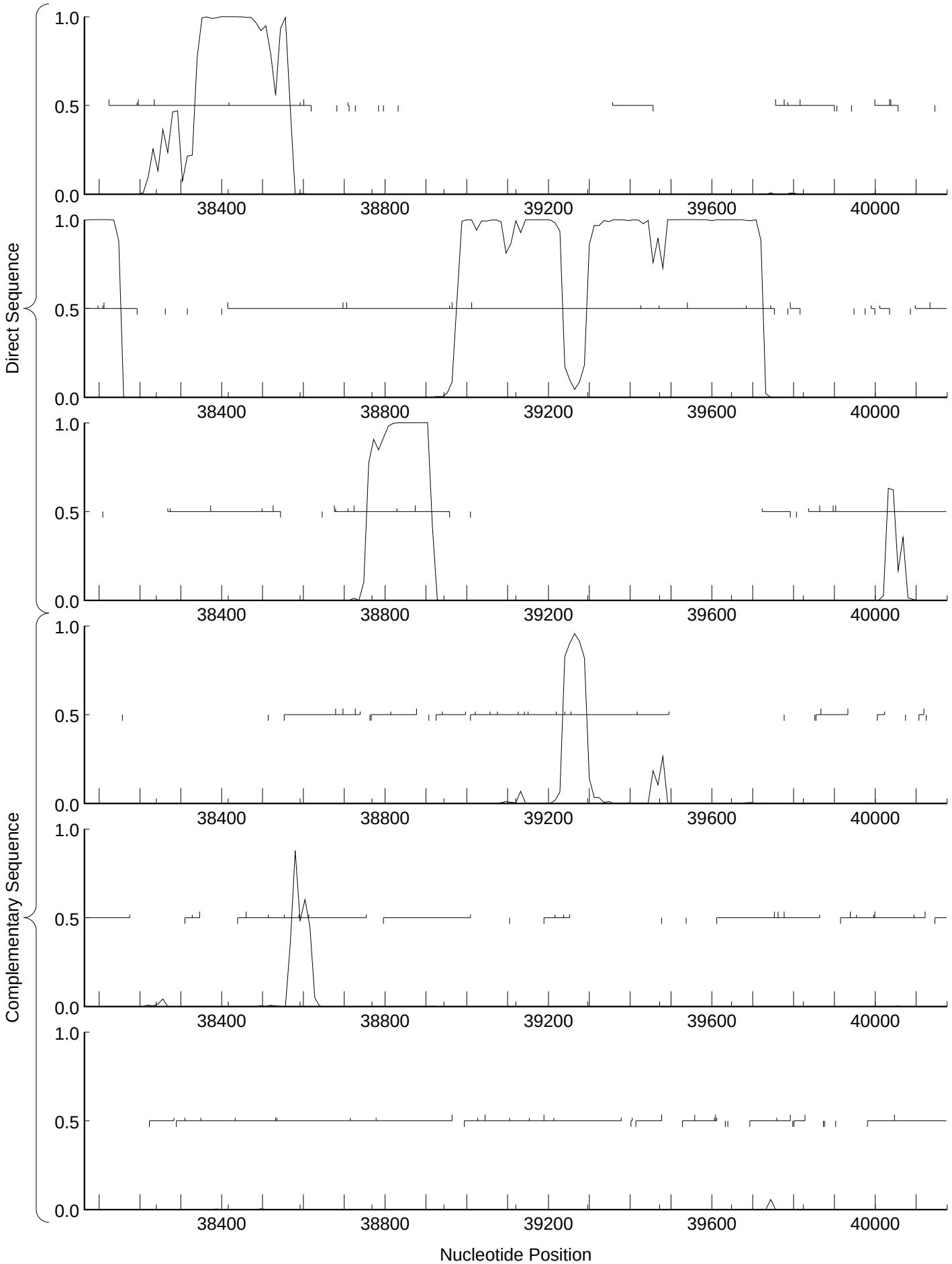
complete sequence, 41893 bp including 11-base 3’ overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 18/21



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

