GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Antsirabe complete sequence, 44739 bp including 10-base 3’ overhang (CC
Analysis Date    :  3/28/24 at 13:16:00
Pages            :  23
Sequence Length  :  44739 bp
GC Content       :   69.01%

Window Length    :  96 bp
Window Step      :  12 bp
Threshold Value  :  0.500

PS-Version       :  1.2

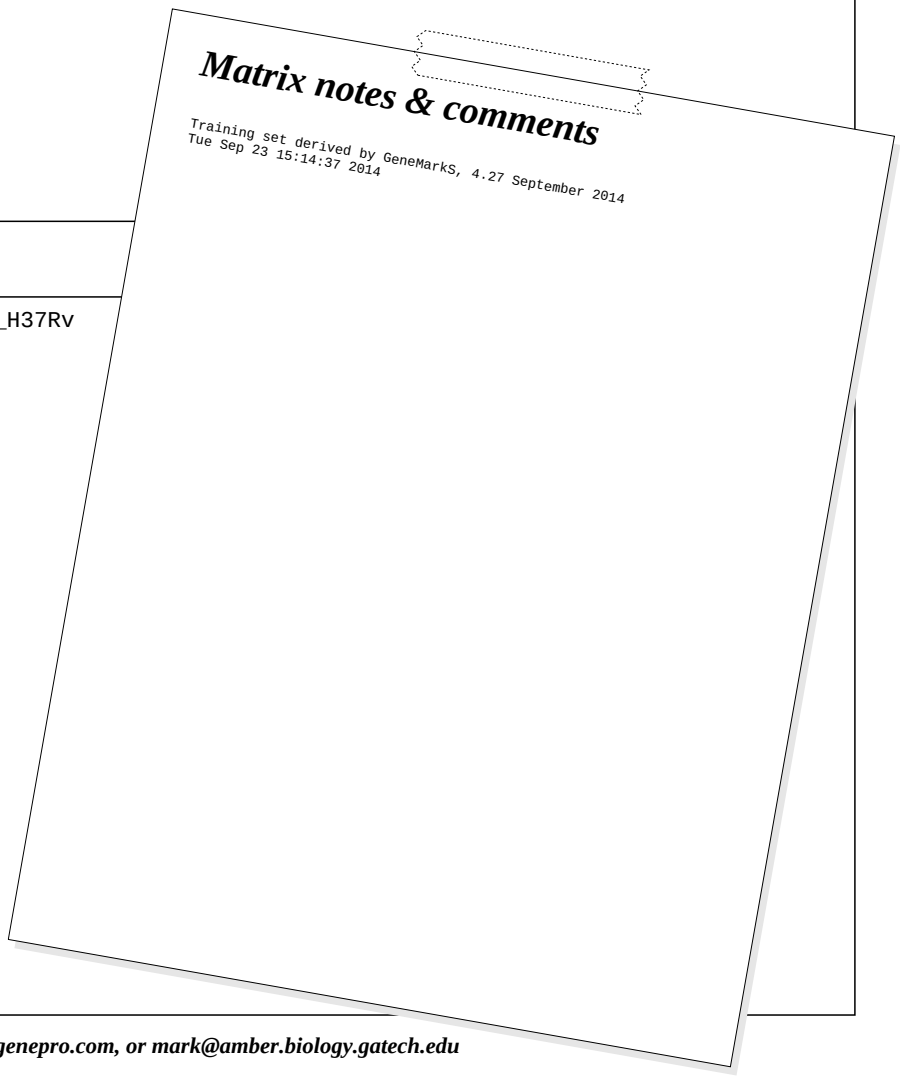
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
```

MATRIX INFORMATION

```
Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

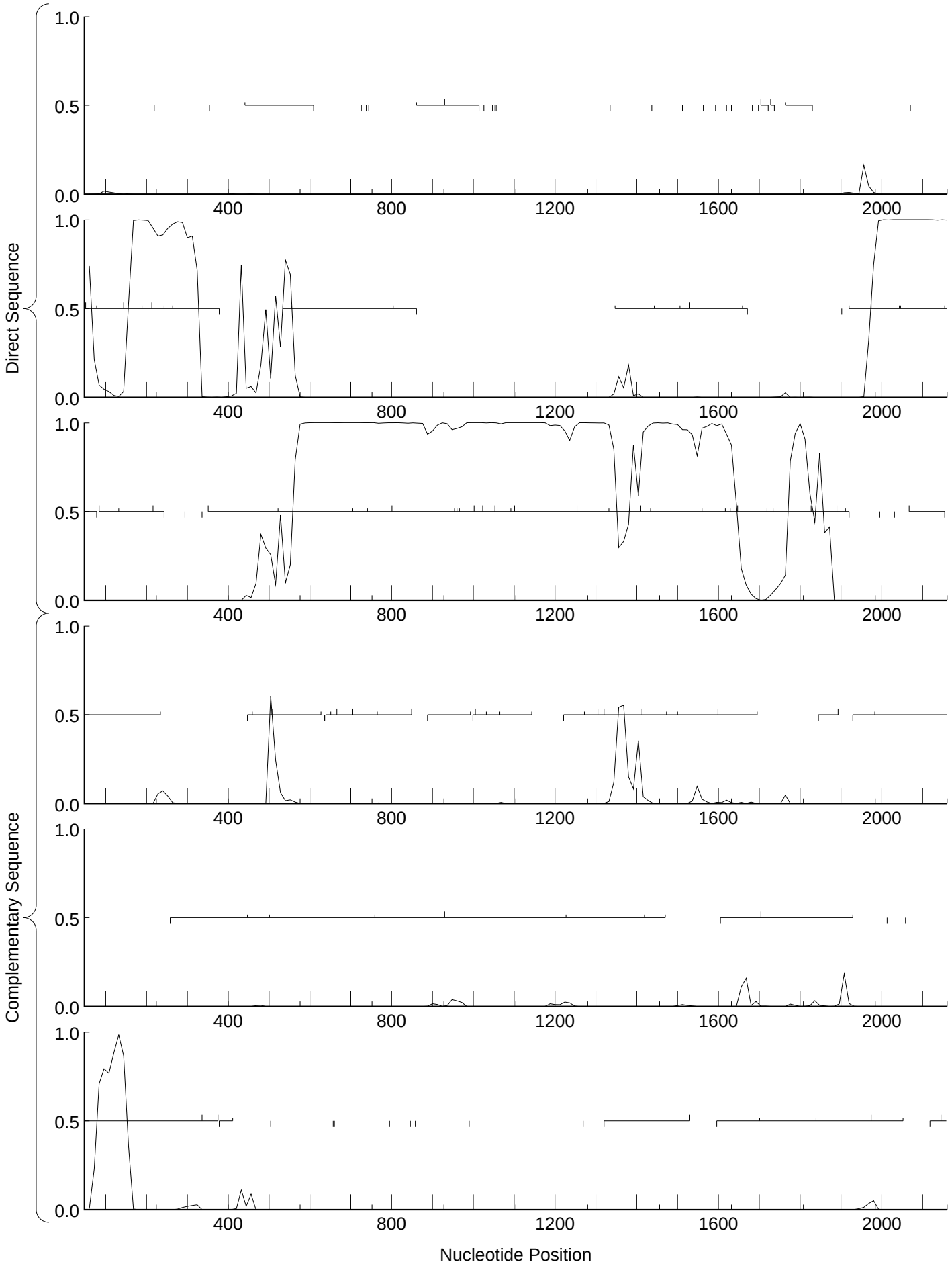
Order      :  4


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

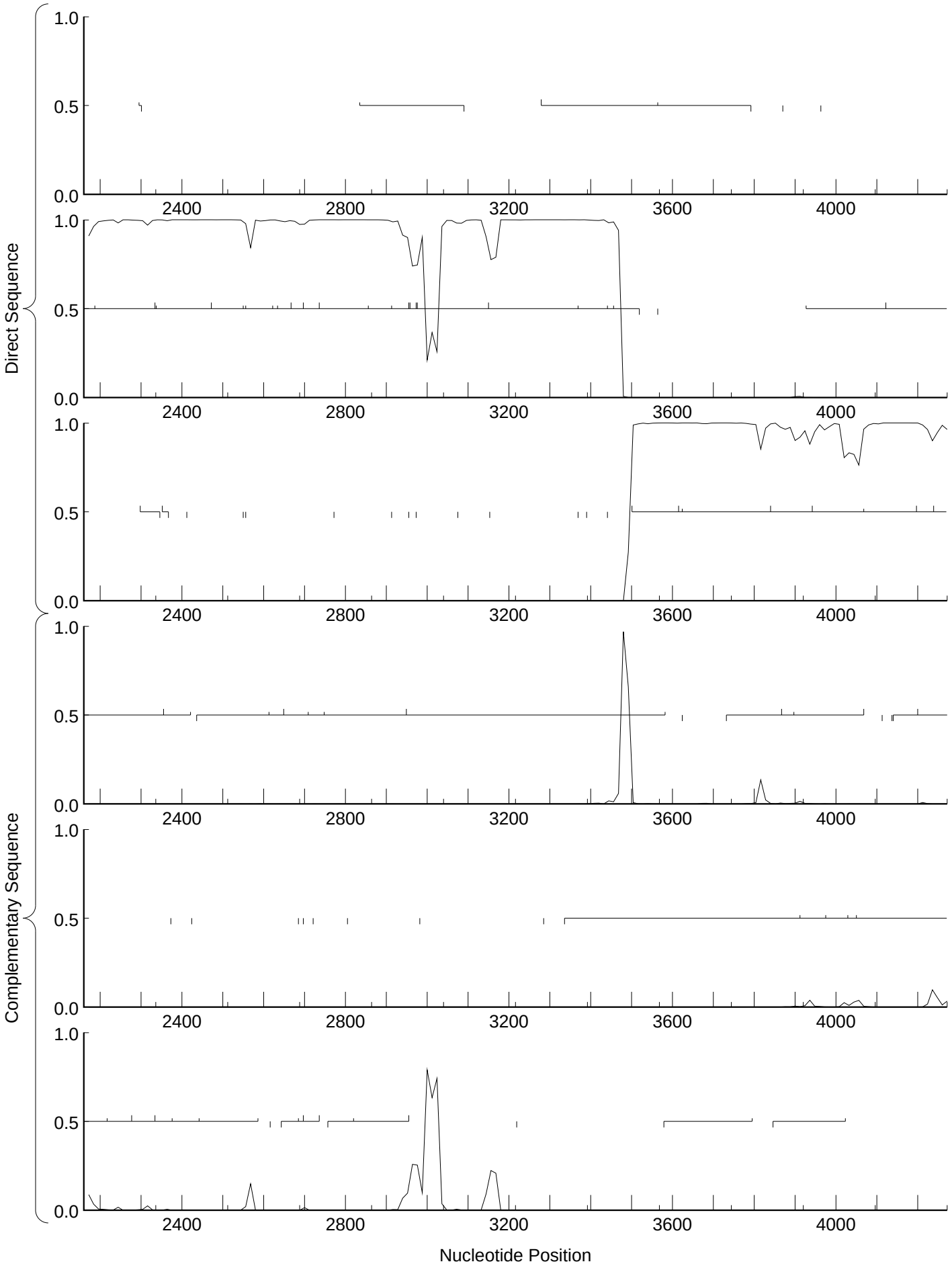


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

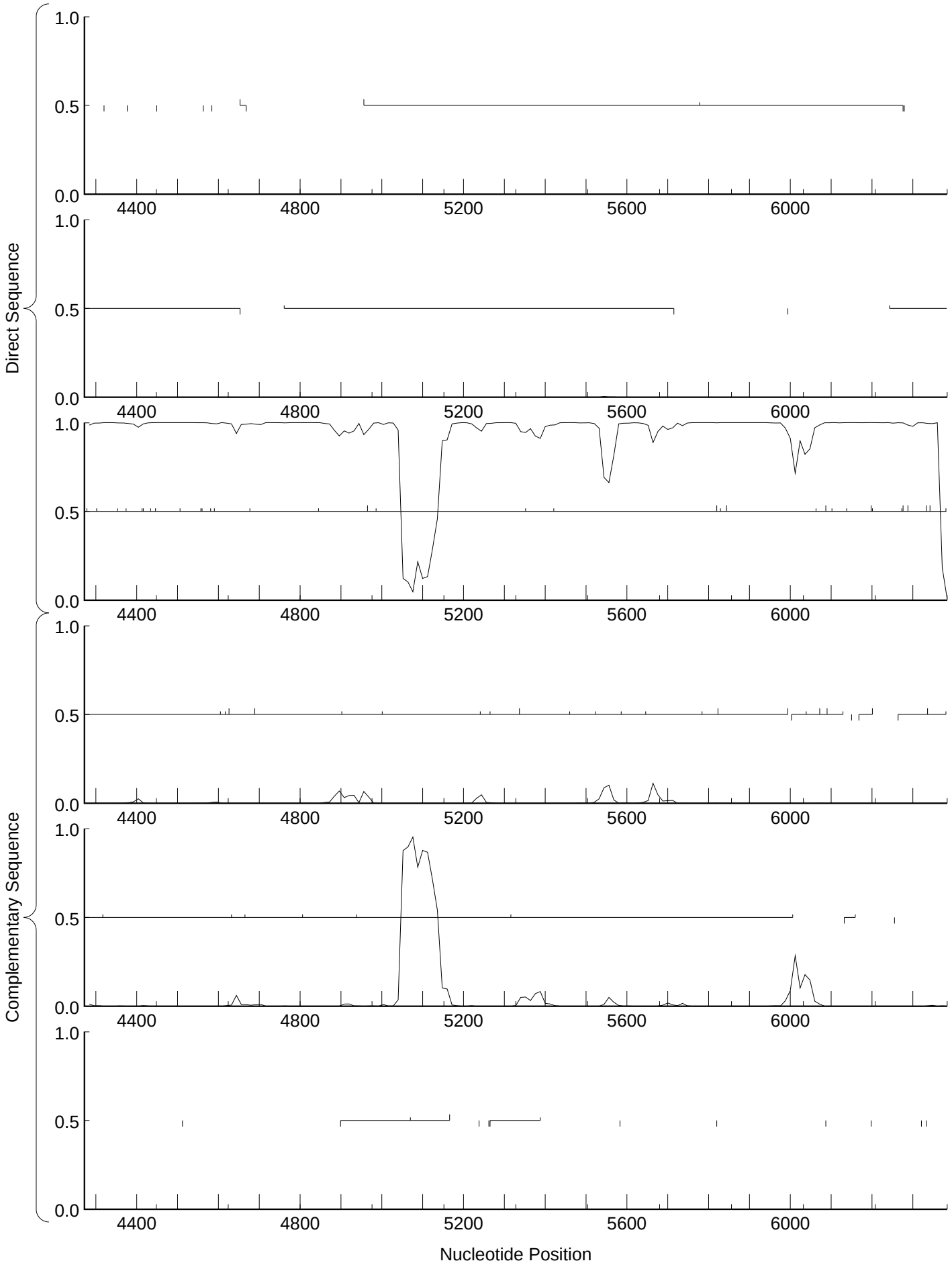
be complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 2/23



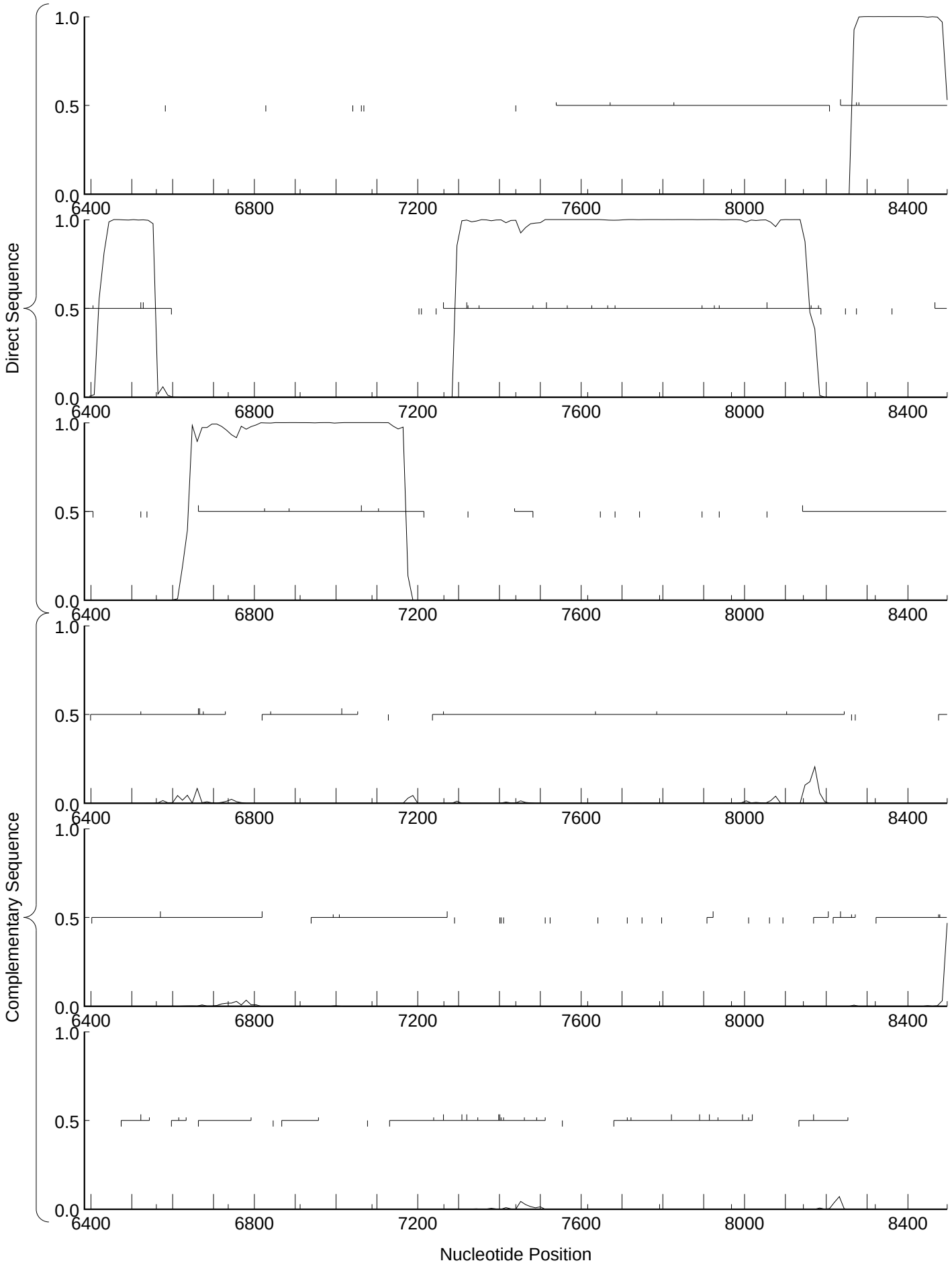
be complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 3/23



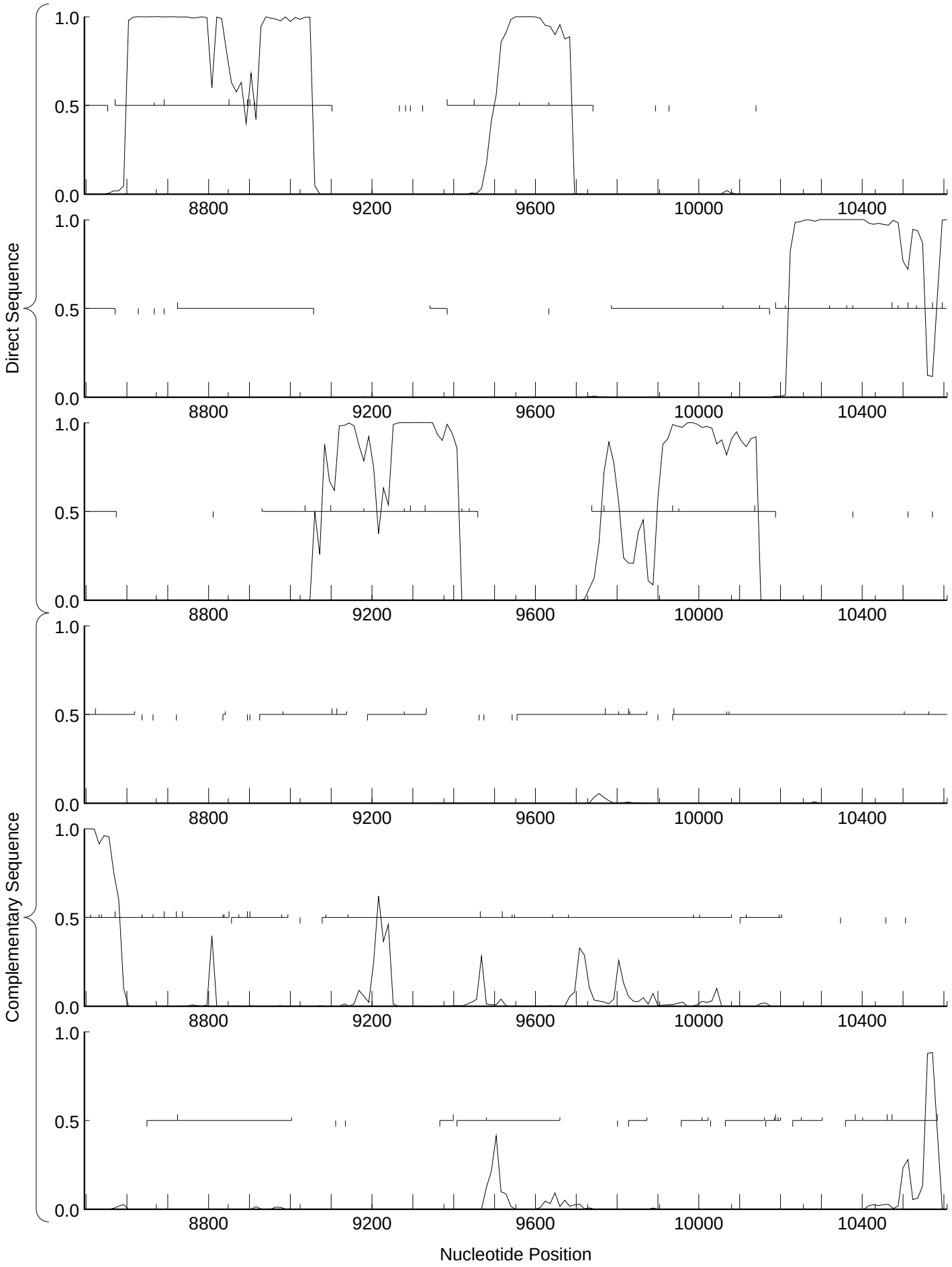
be complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 4/23



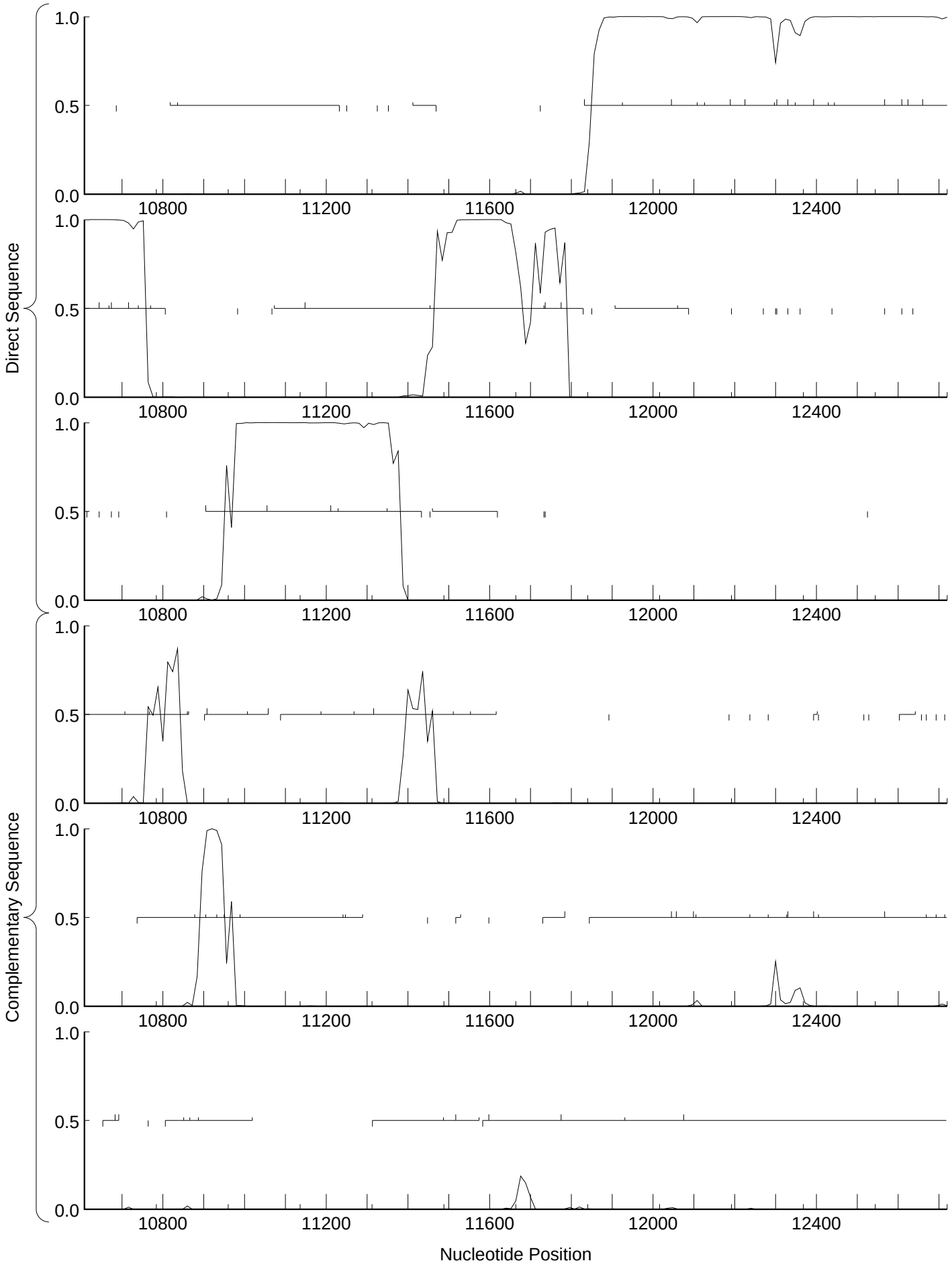
be complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 5/23



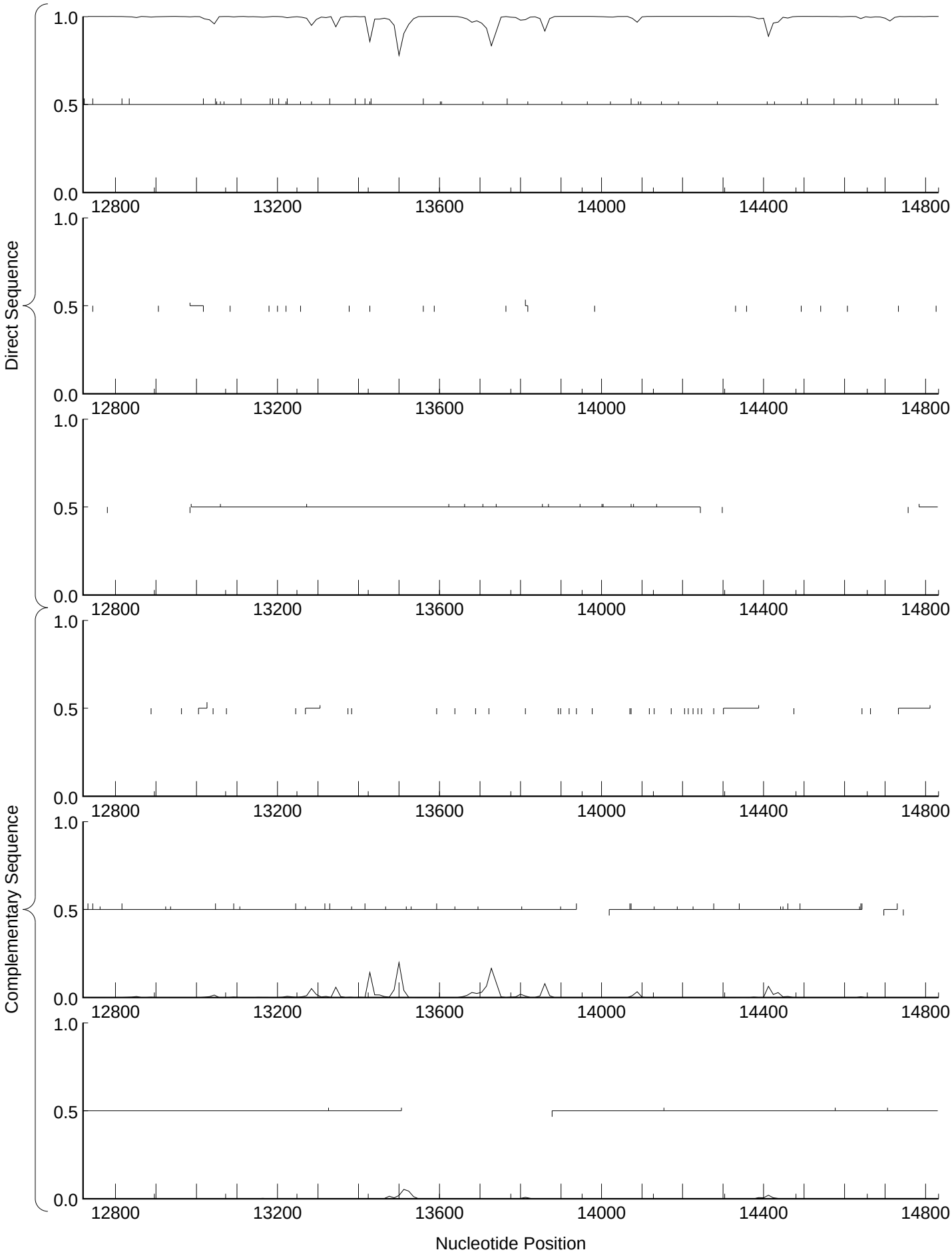
be complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 6/23



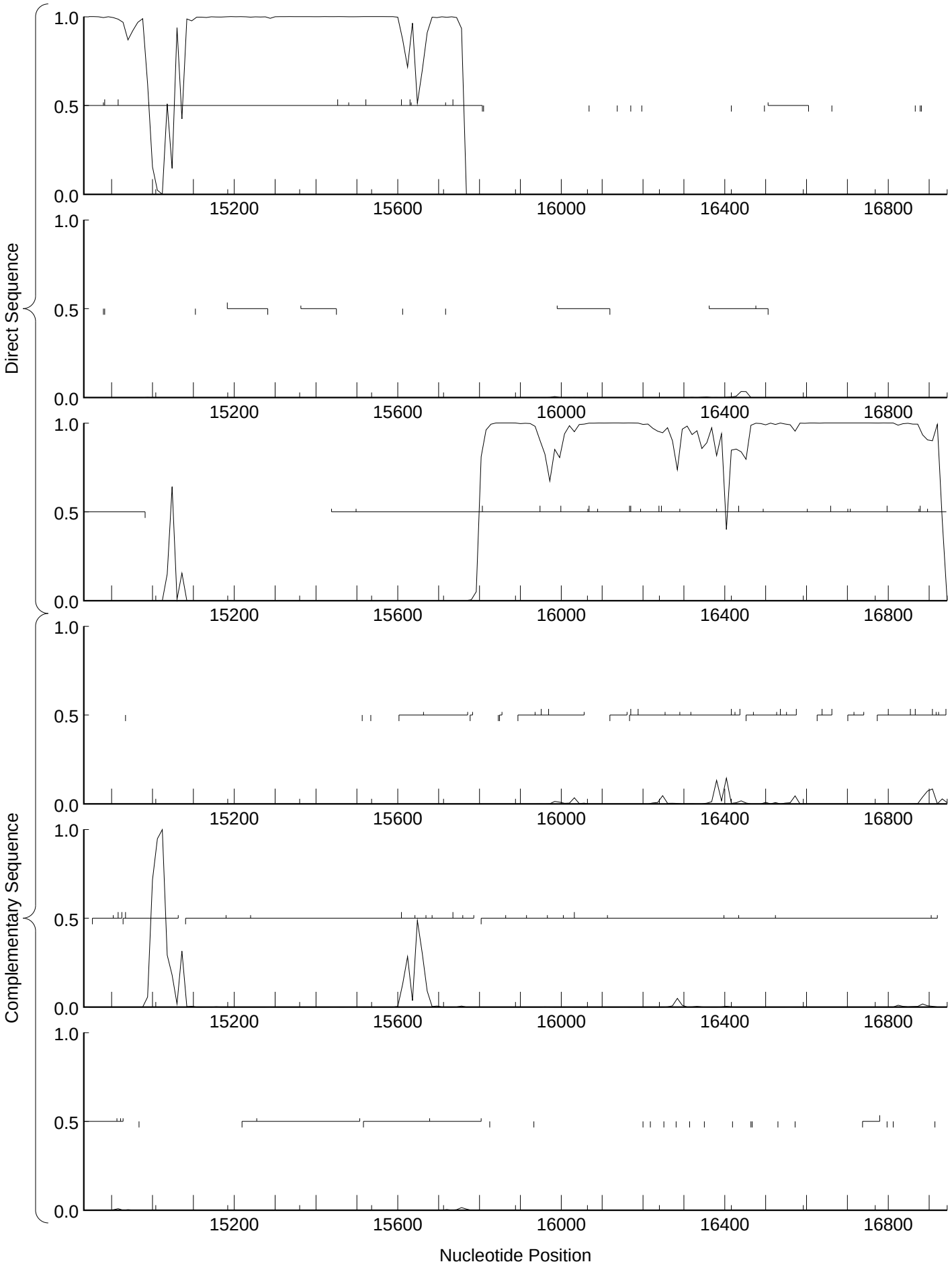
be complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 7/23



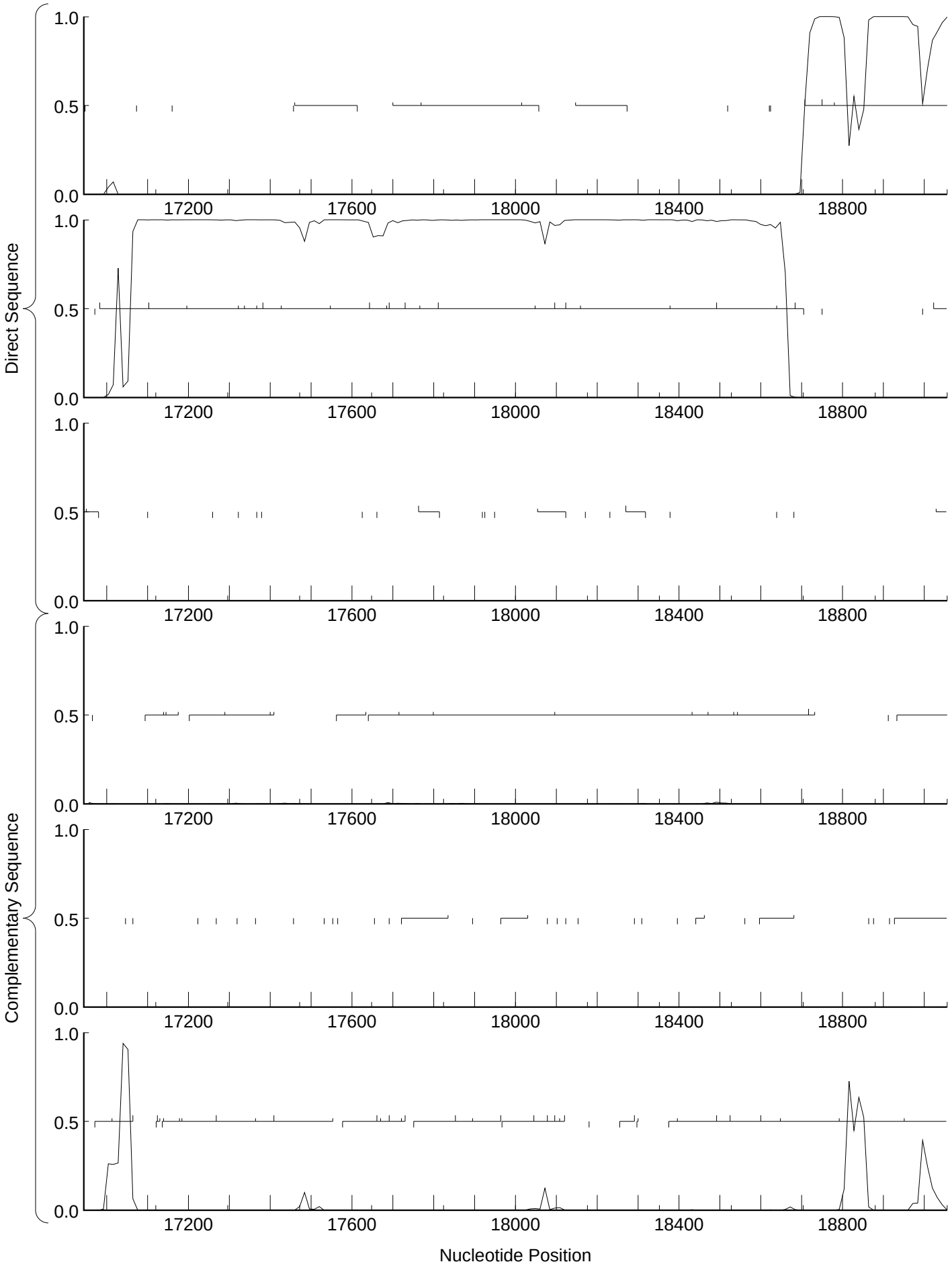
be complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 8/23



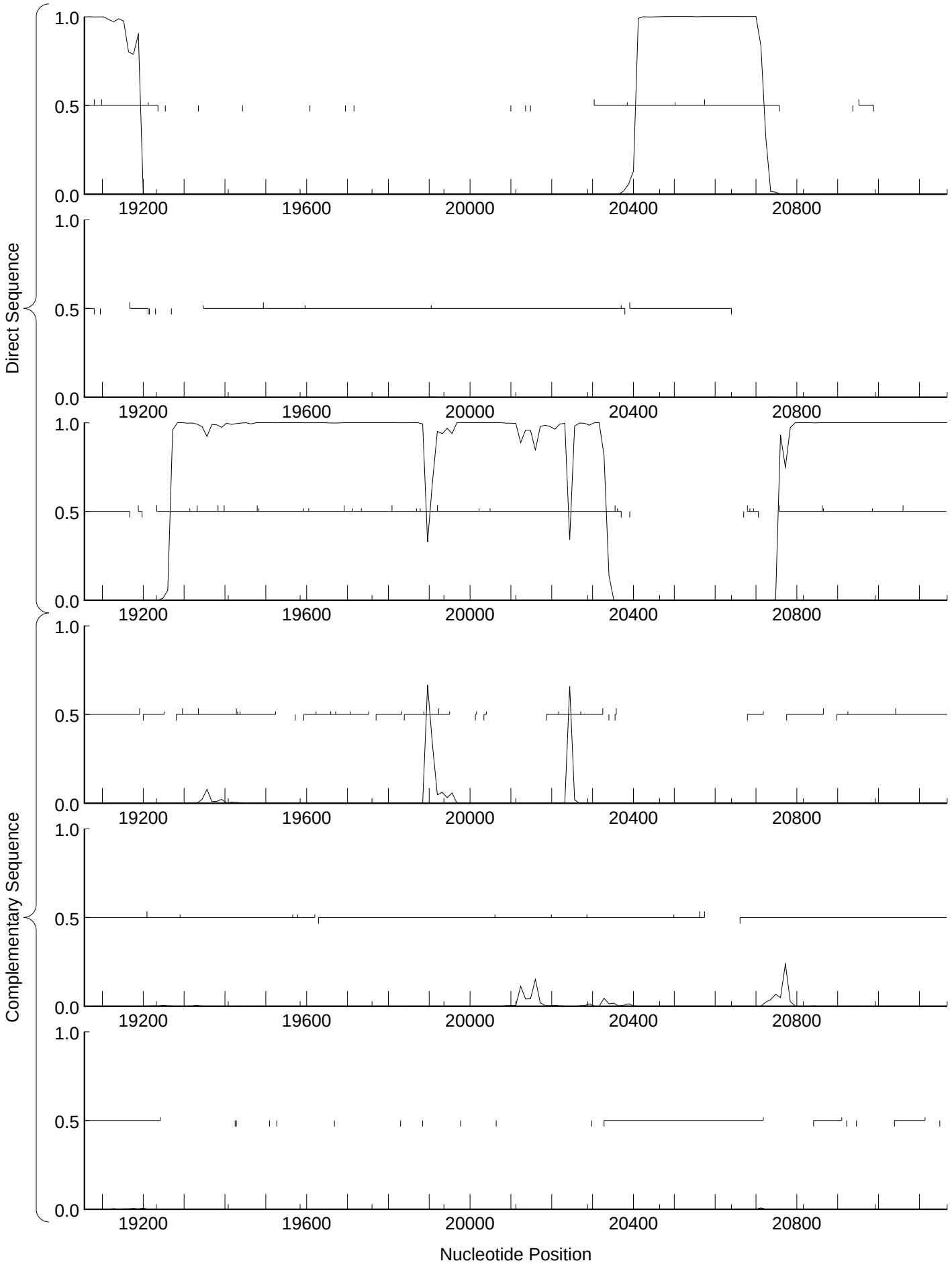
be complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 9/23



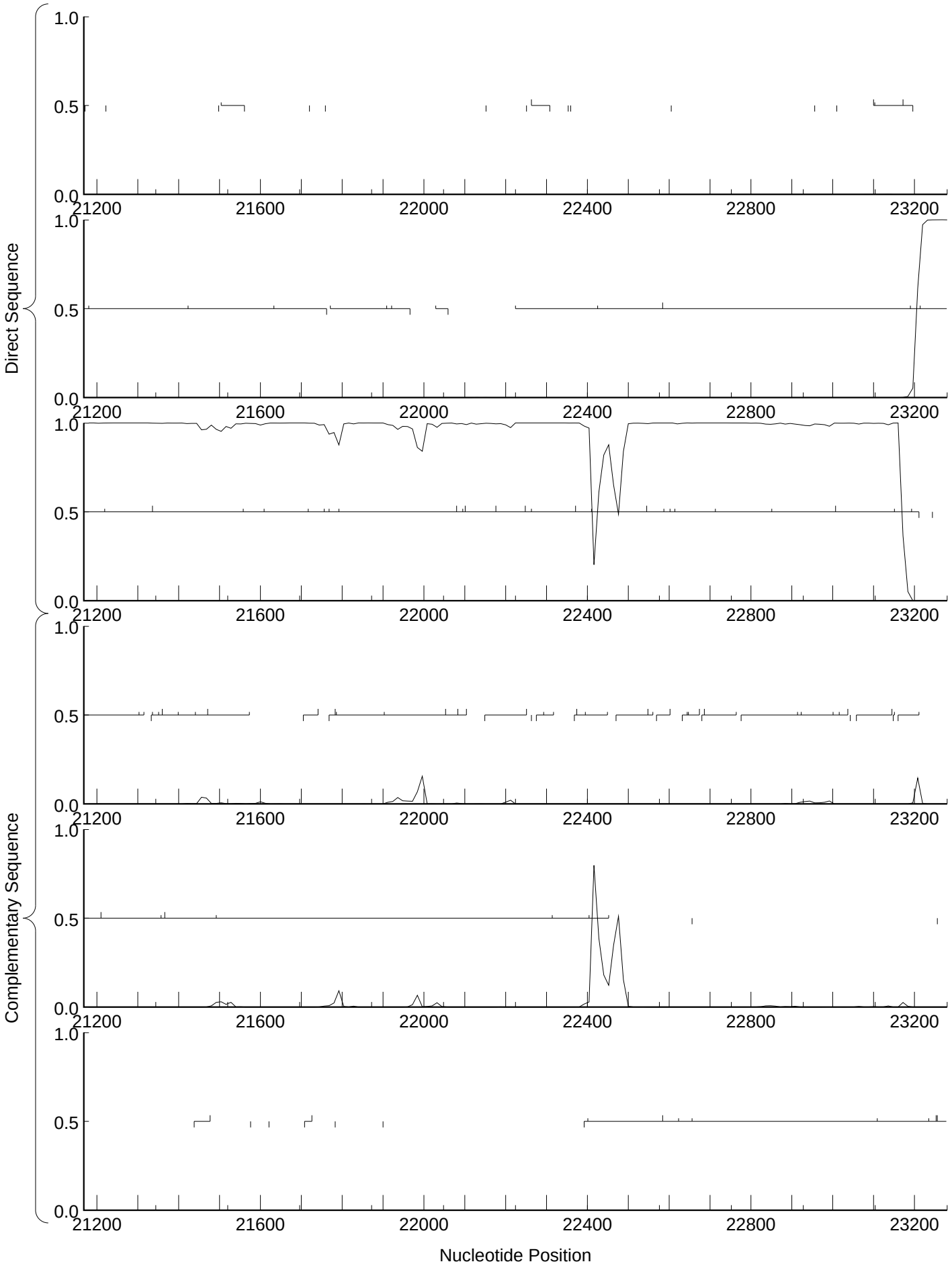
e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 10/23



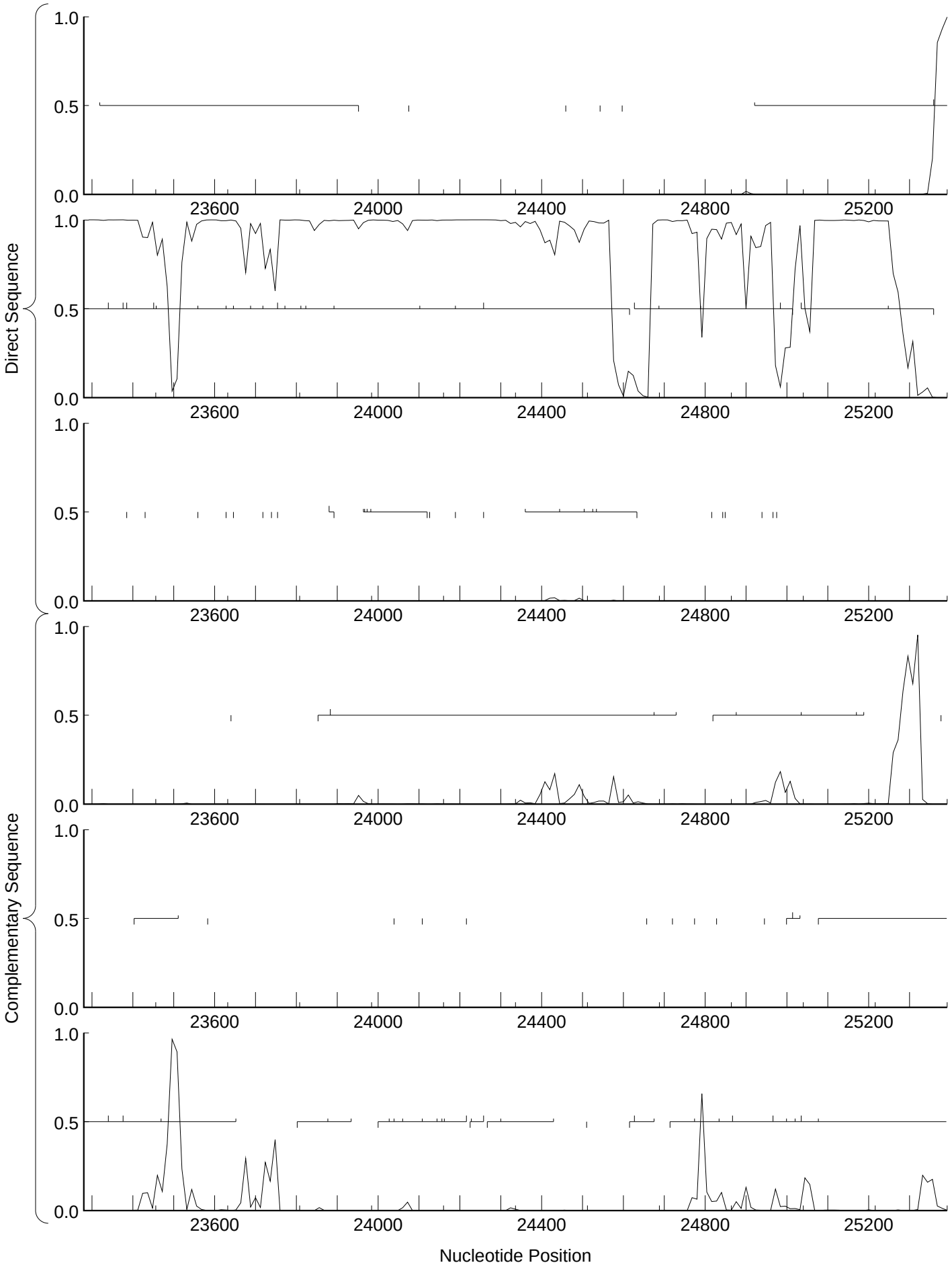
e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 11/23



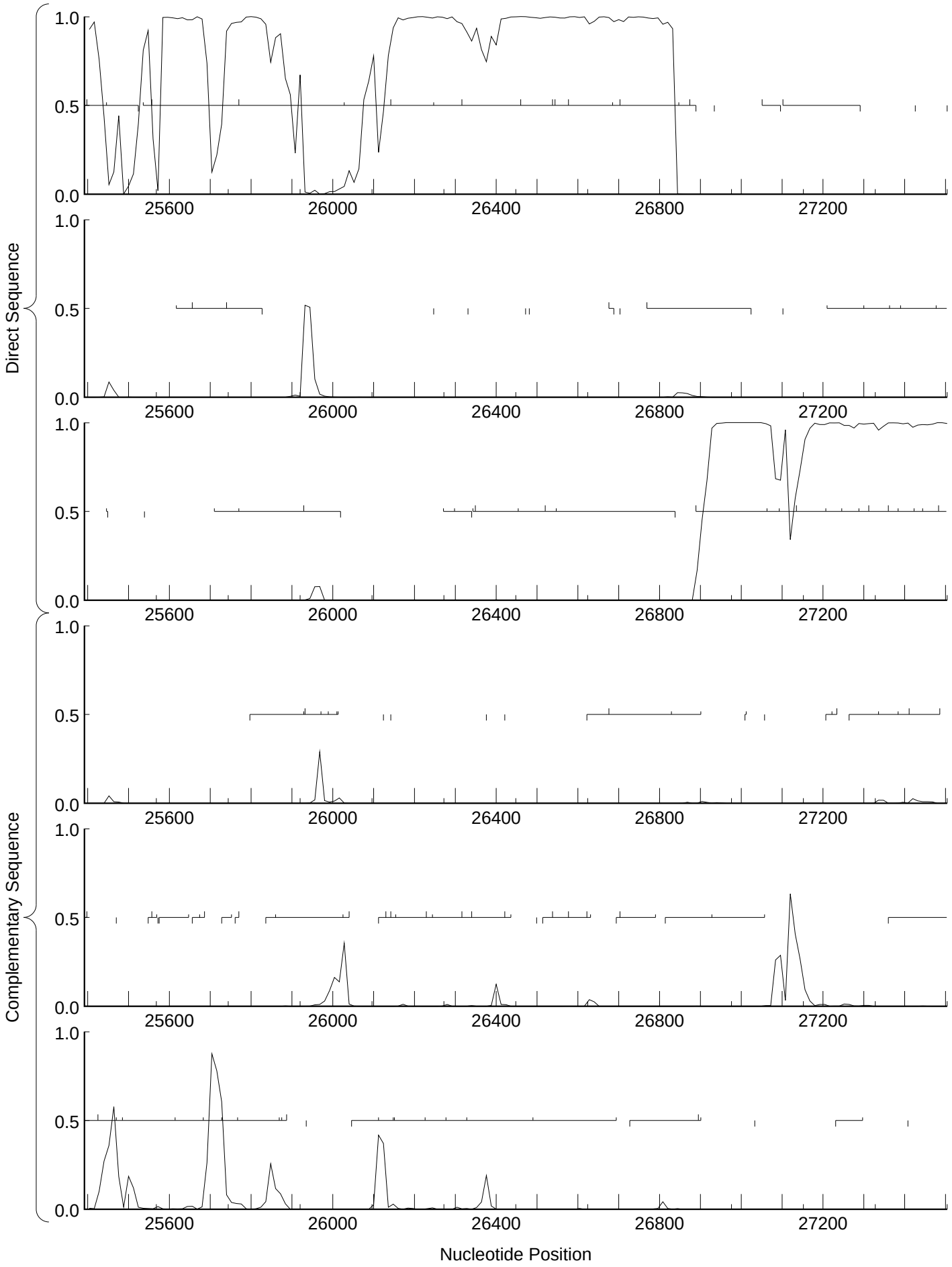
e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 12/23



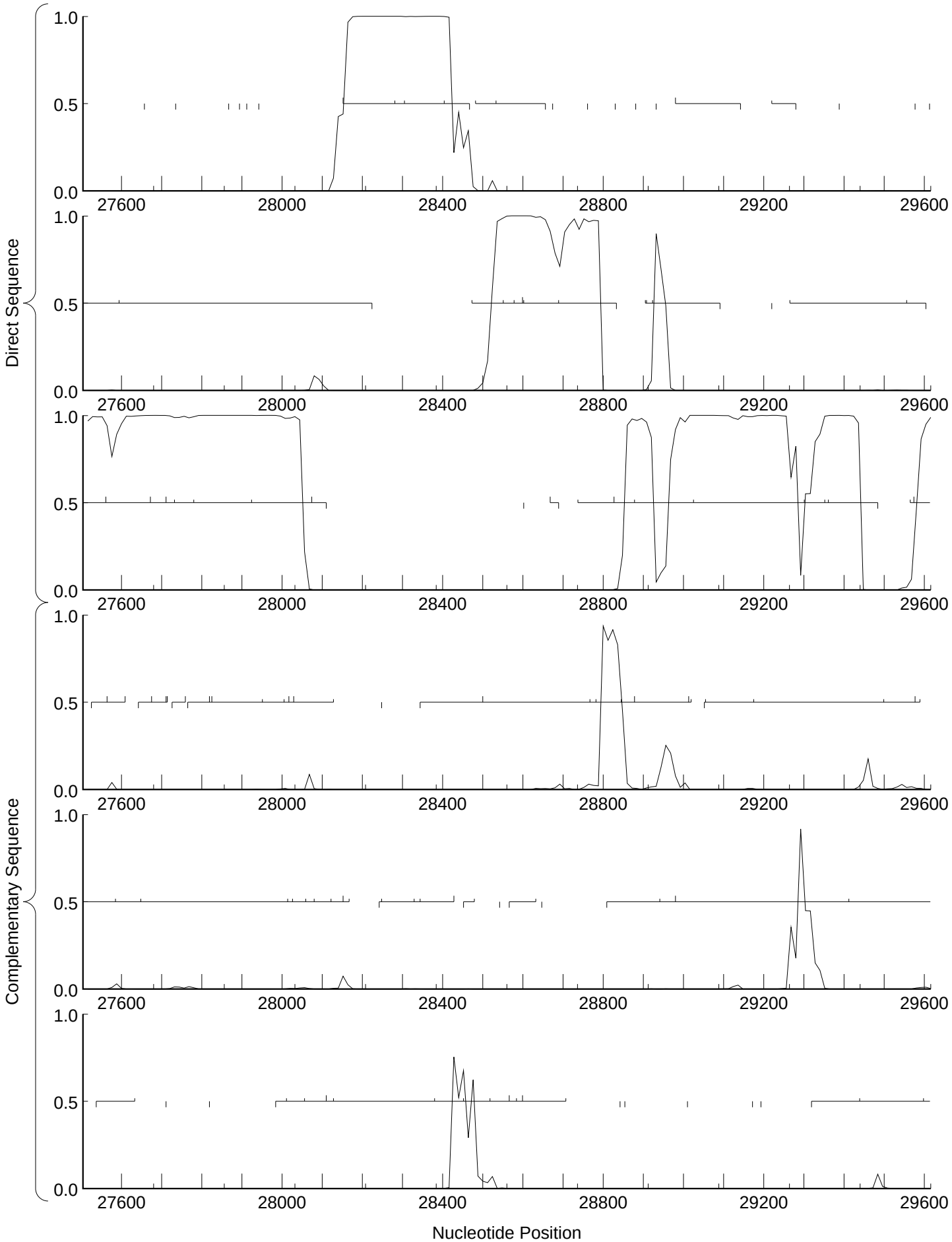
e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 13/23



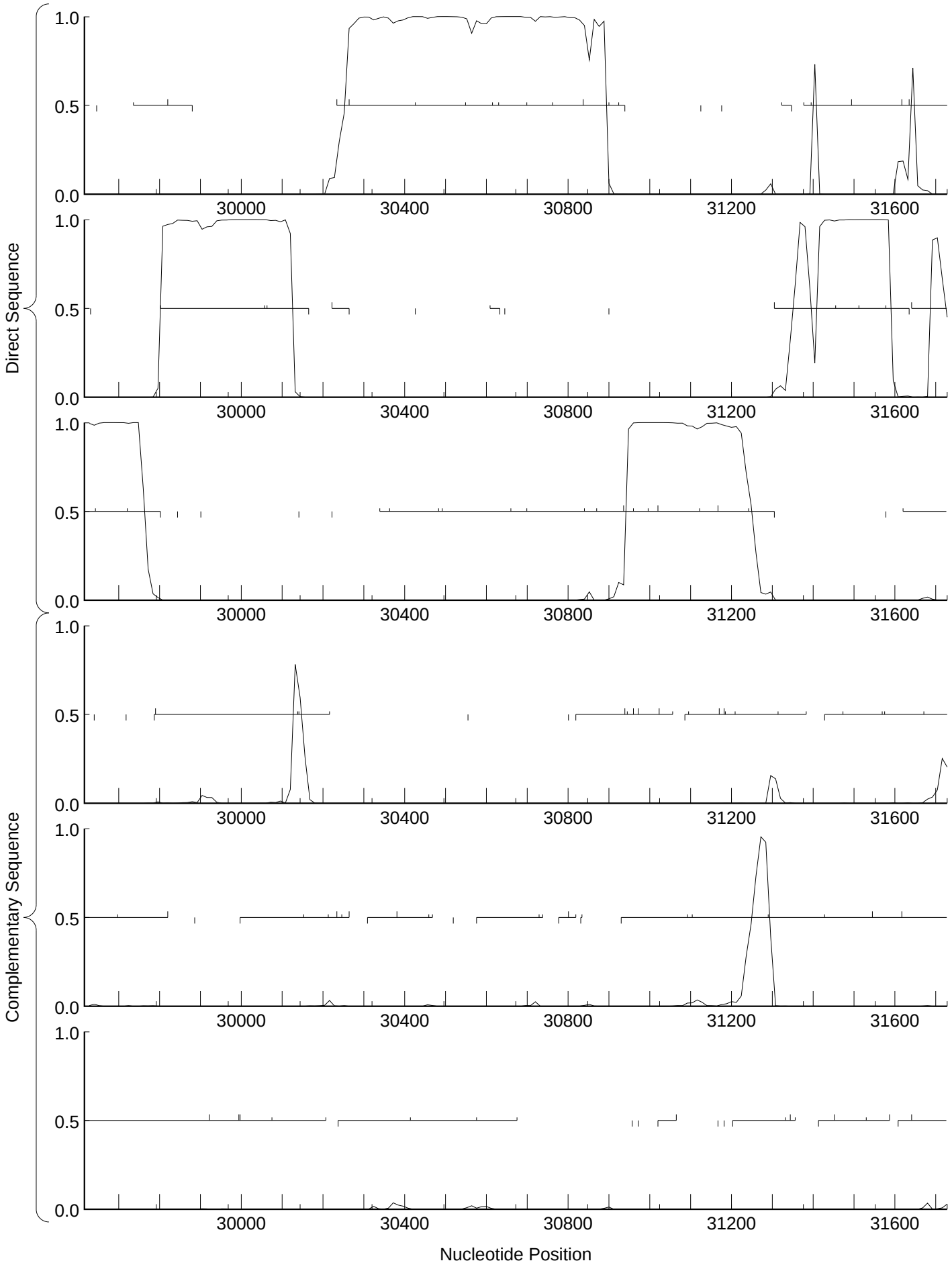
e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 14/23



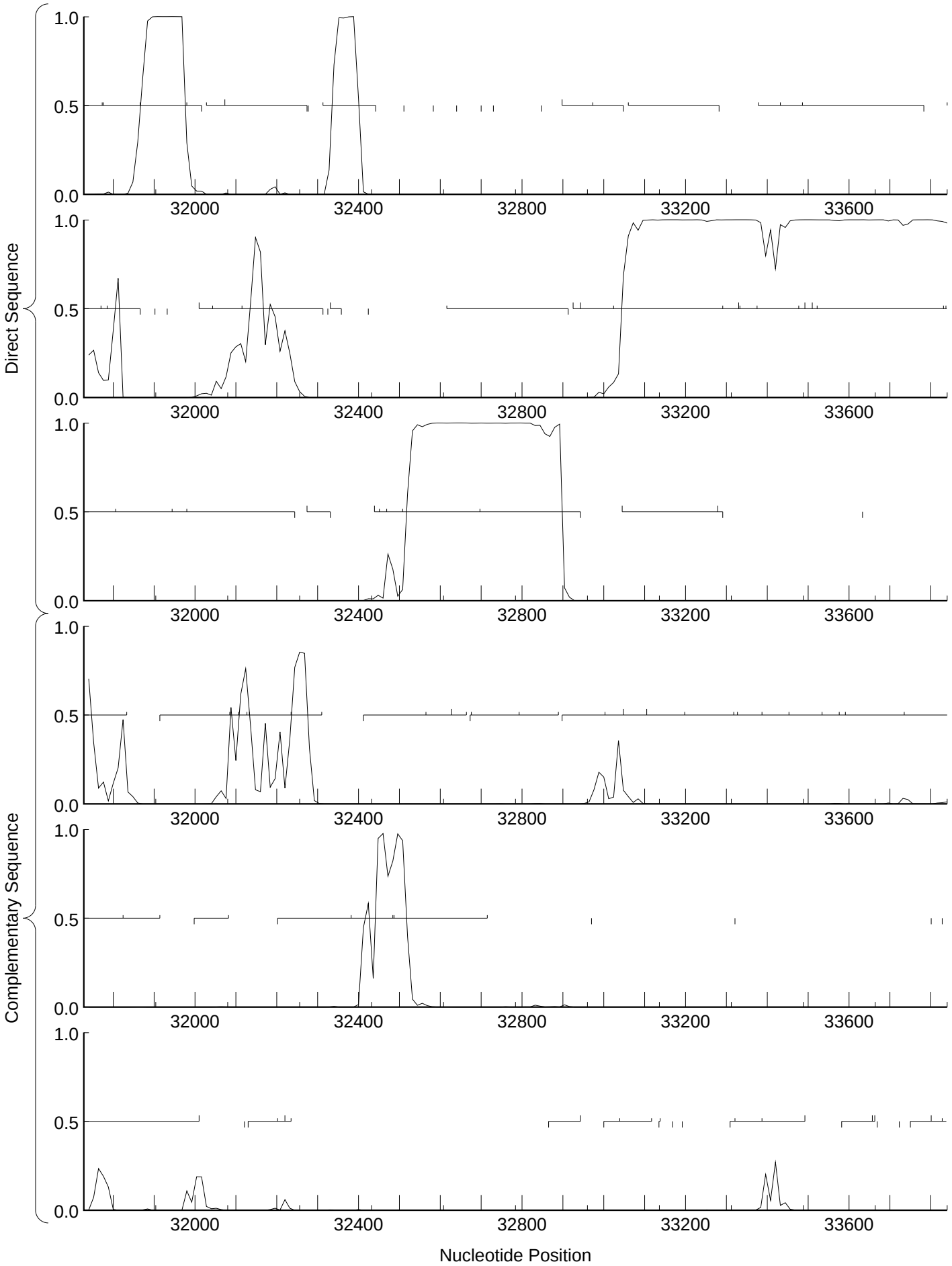
e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 15/23



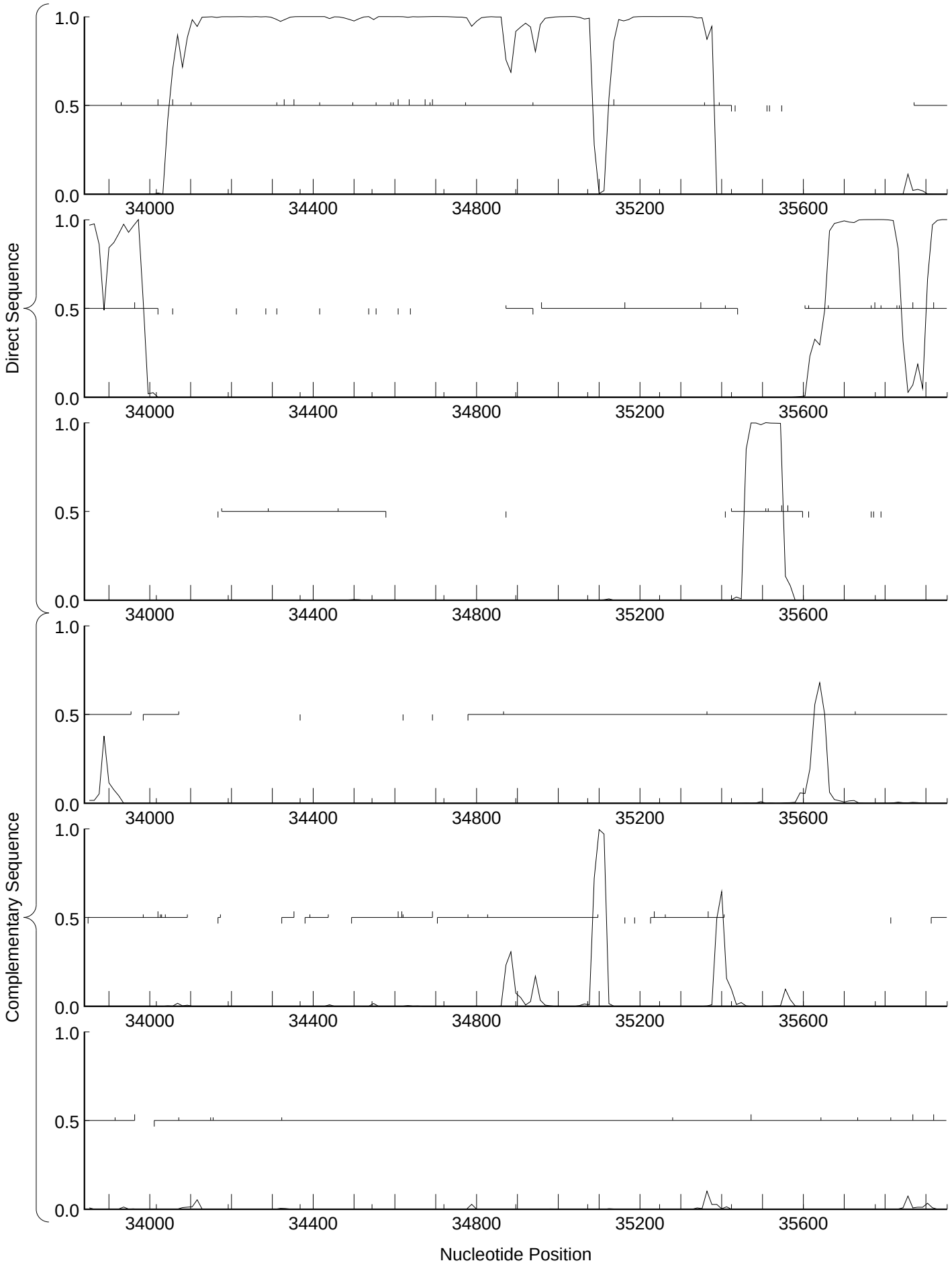
e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 16/23



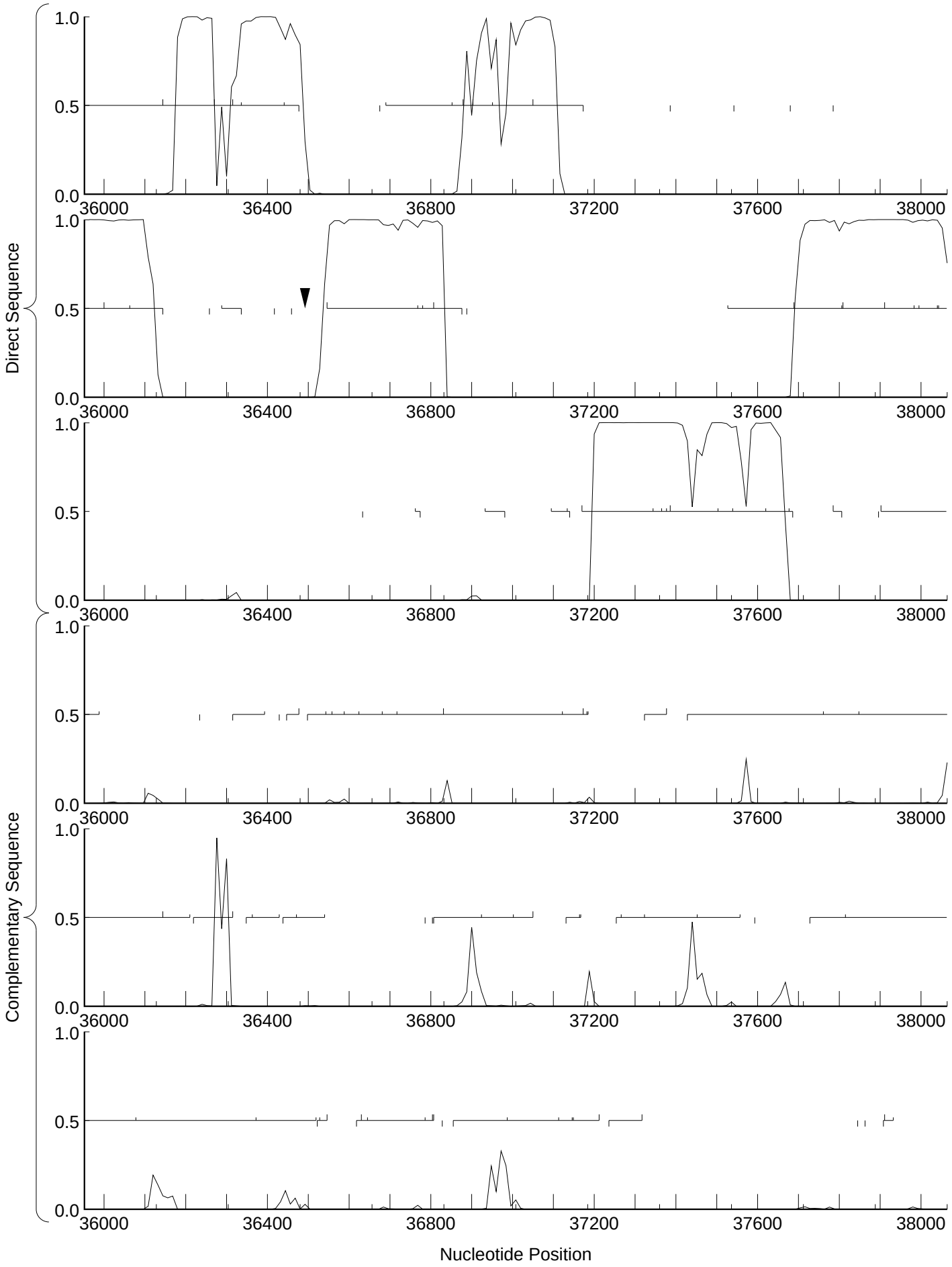
e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 17/23



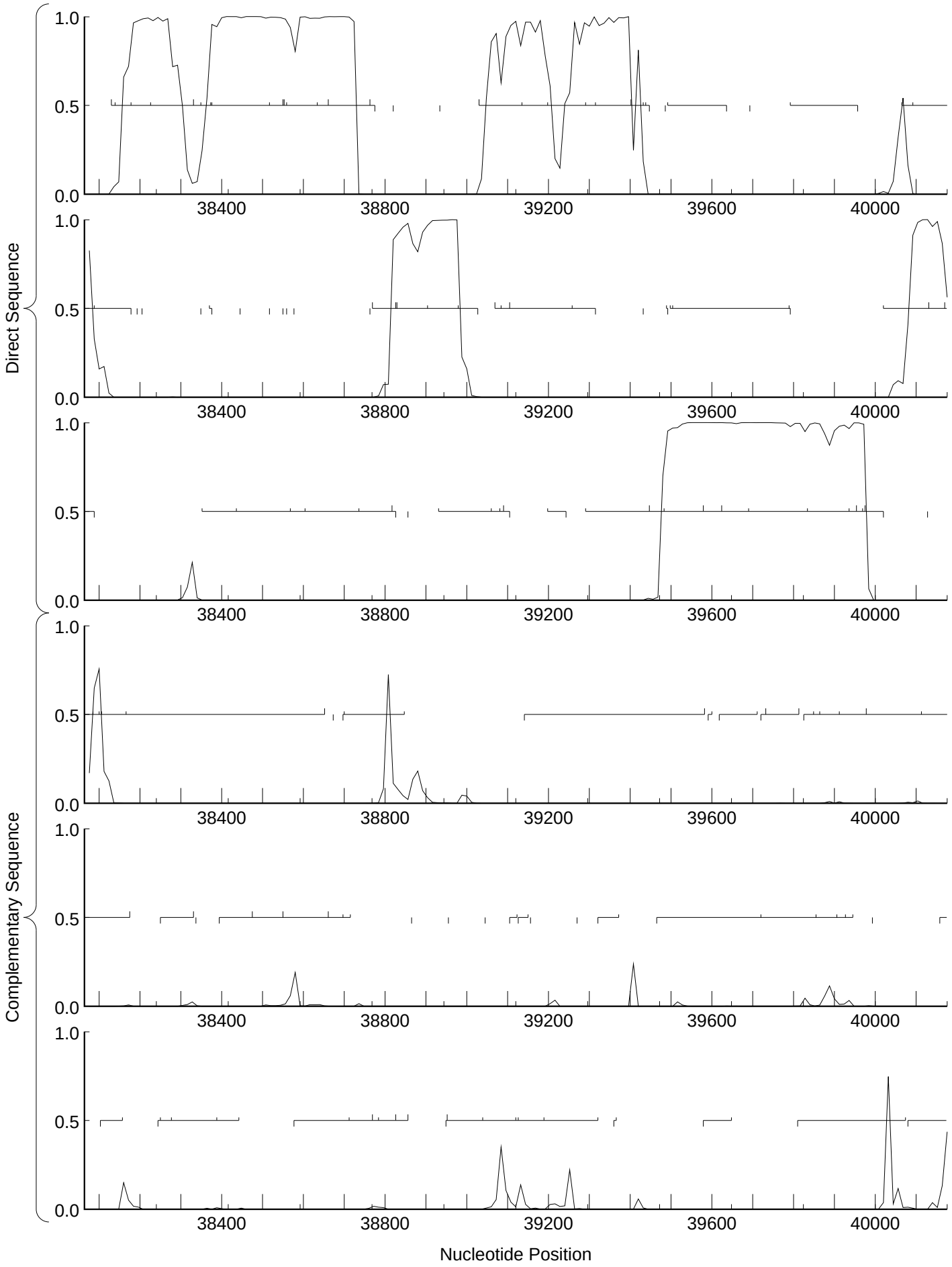
e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 18/23



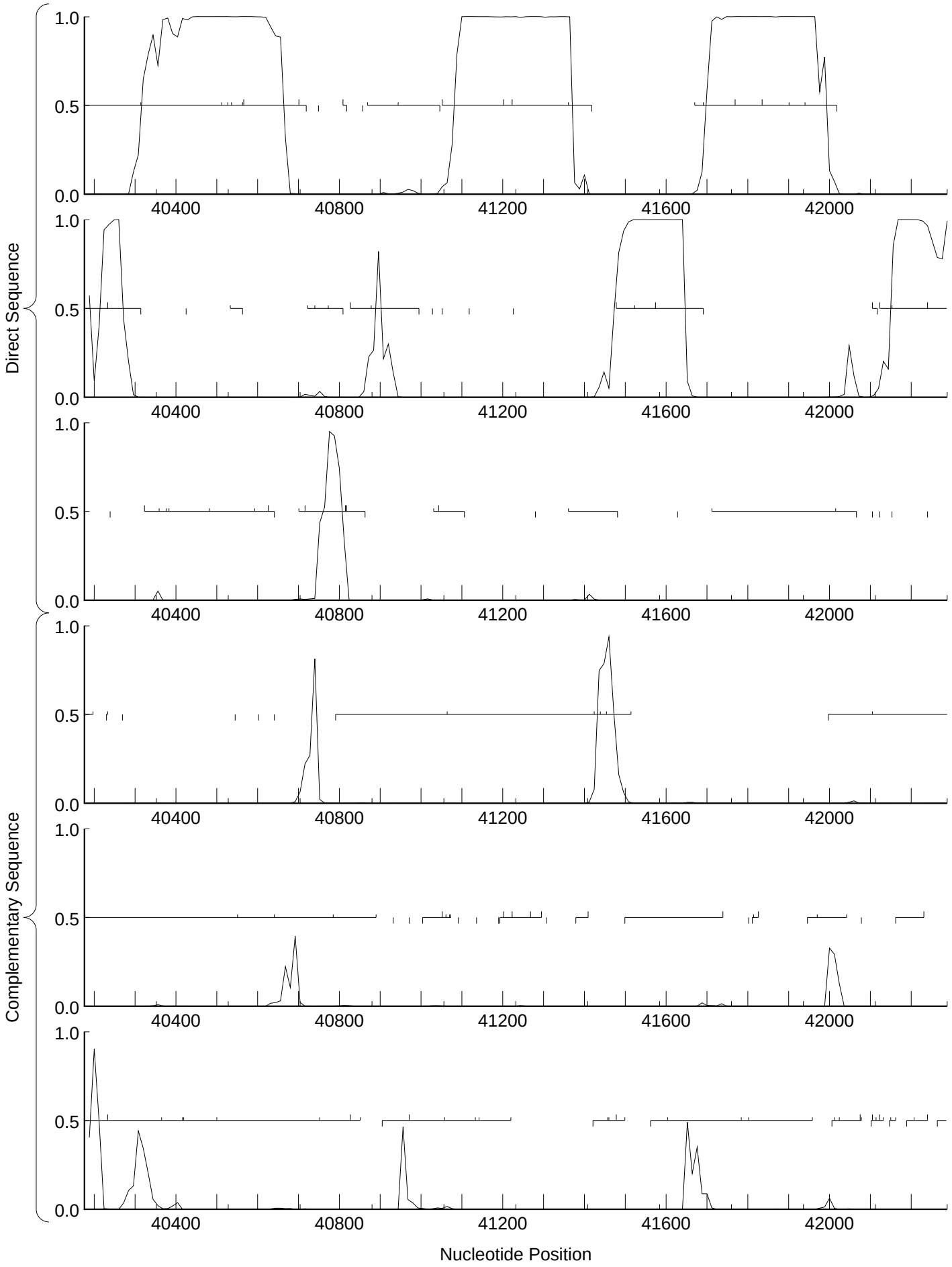
e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 19/23



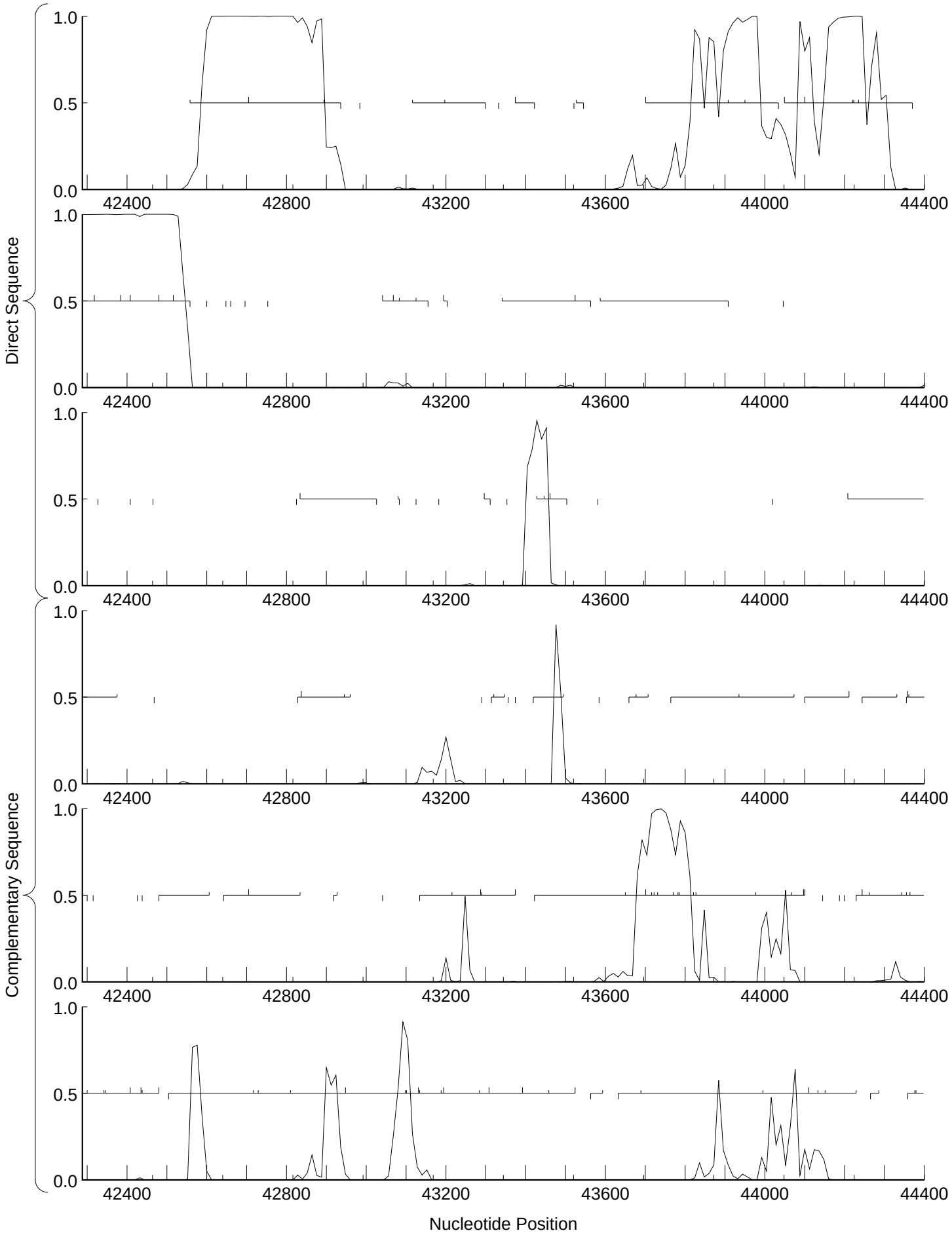
e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 20/23



e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 21/23



e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 22/23



e complete sequence, 44739 bp including 10-base 3’ overhang (CCCGTGGCAT), Cluster G, Order 4, Window 96, Step 12, 23/23

