GeneMark

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

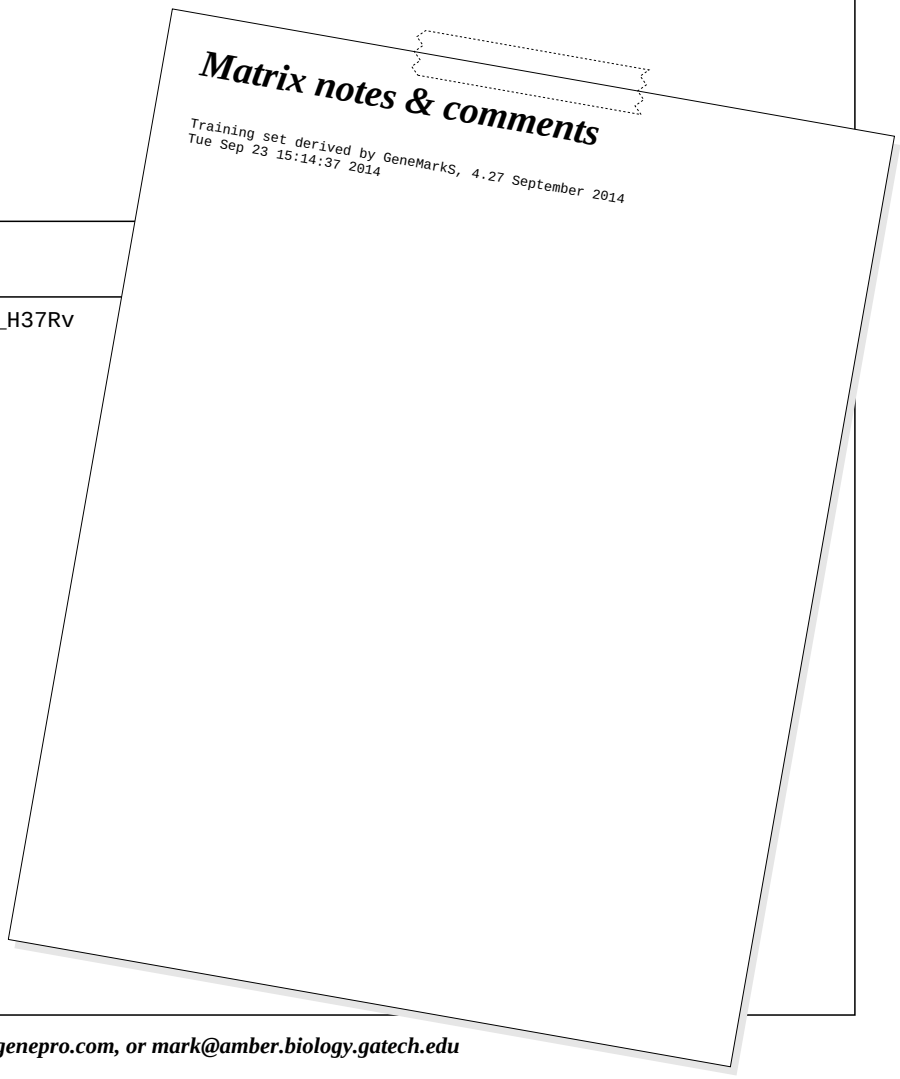
PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Marcoliusprime complete sequence, 58129 bp including 10-base 3' overhan
Analysis Date    :  3/29/24 at 10:13:15
Pages            :  29
Sequence Length  :  58129 bp
GC Content       :   68.25%

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

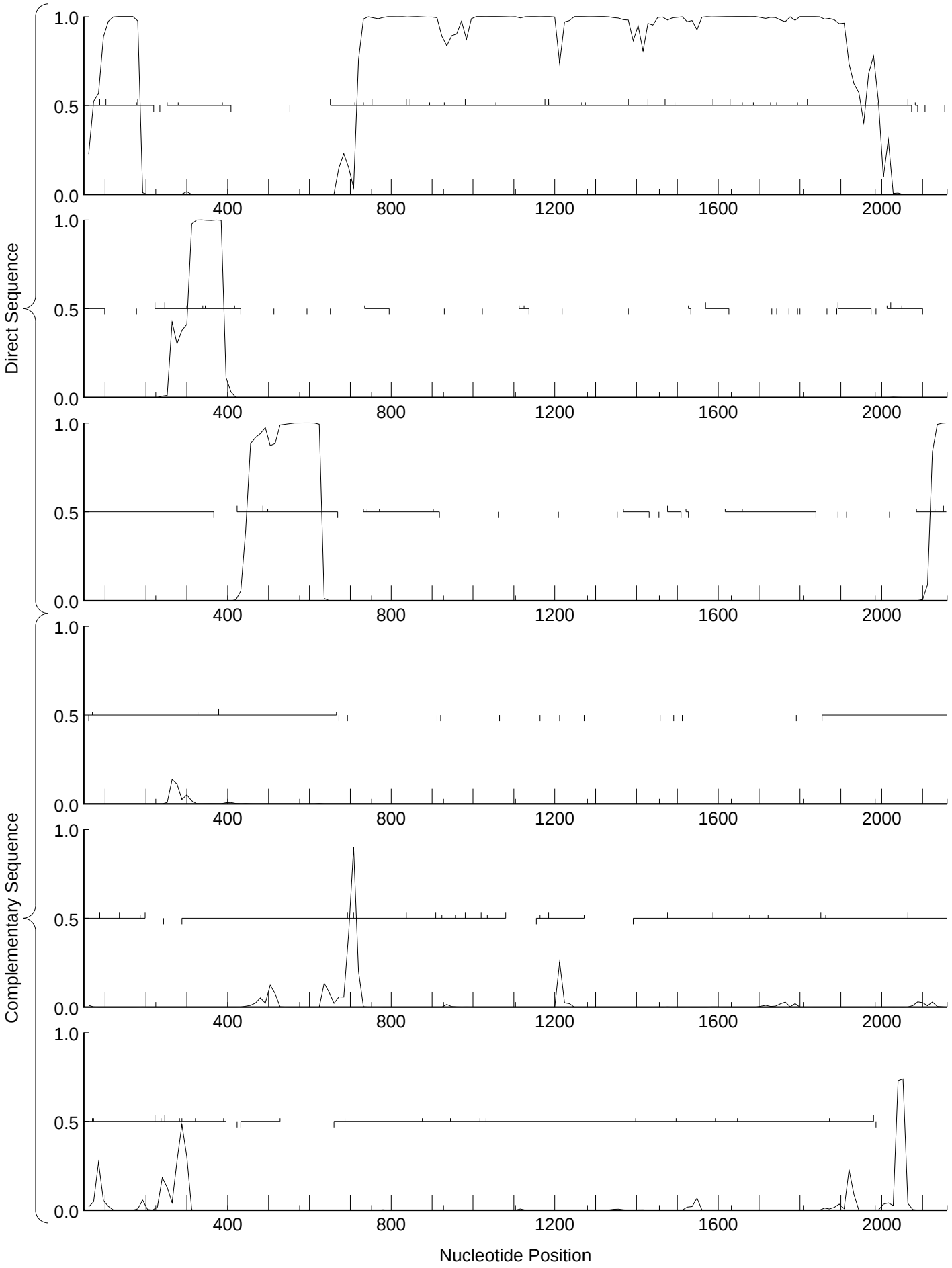
```
Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

Order      :  4
```

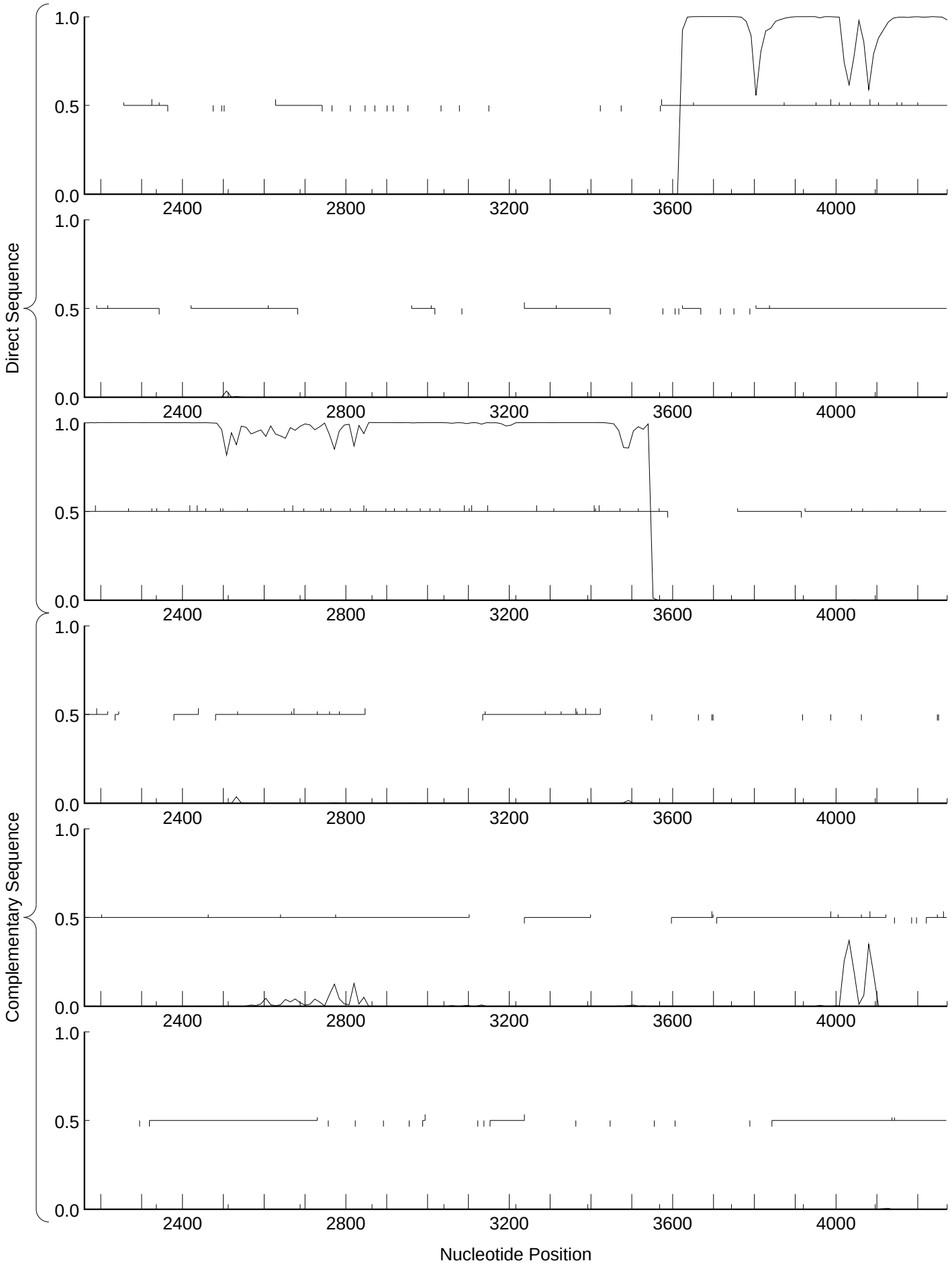

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

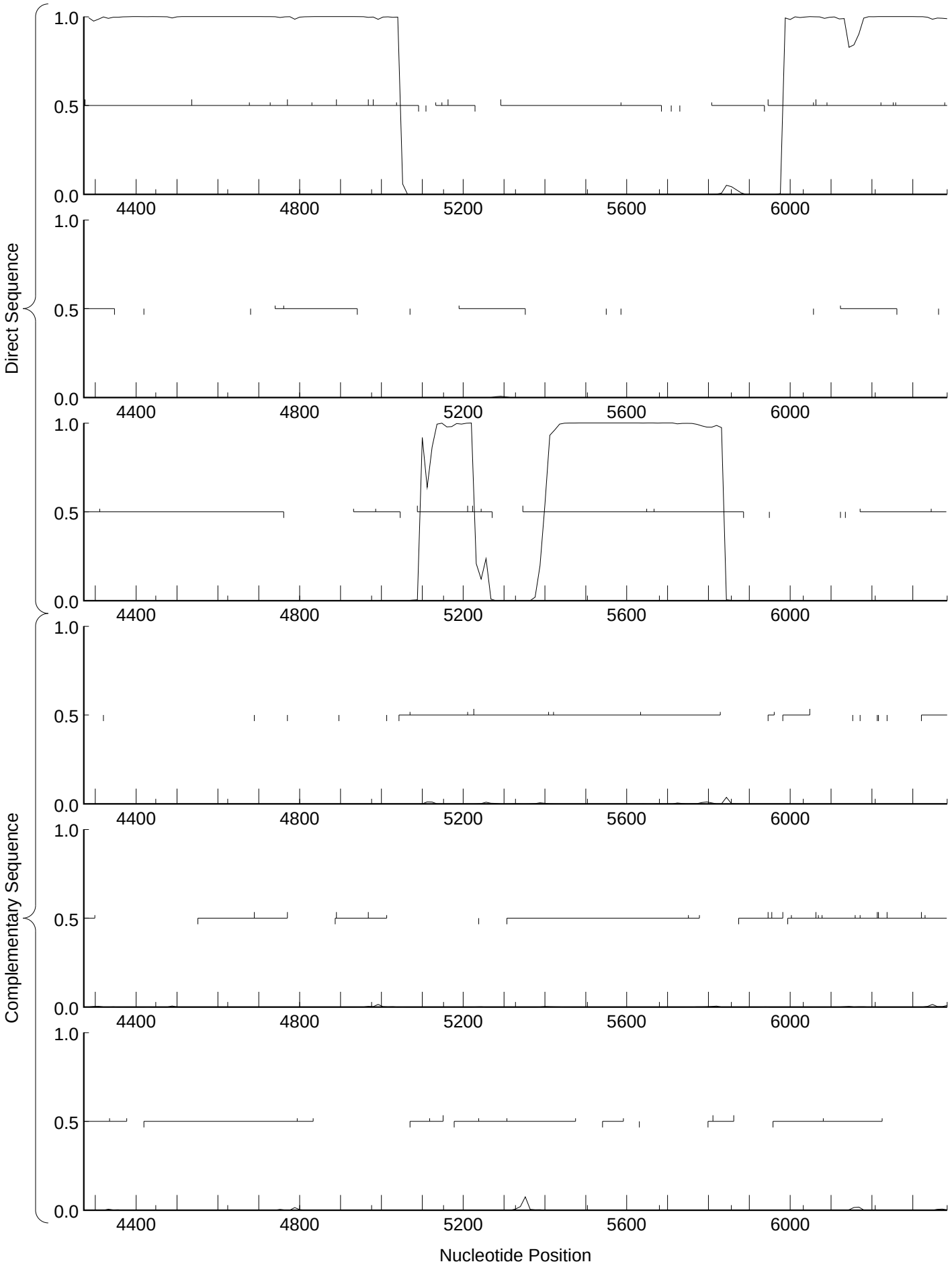
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 2/29



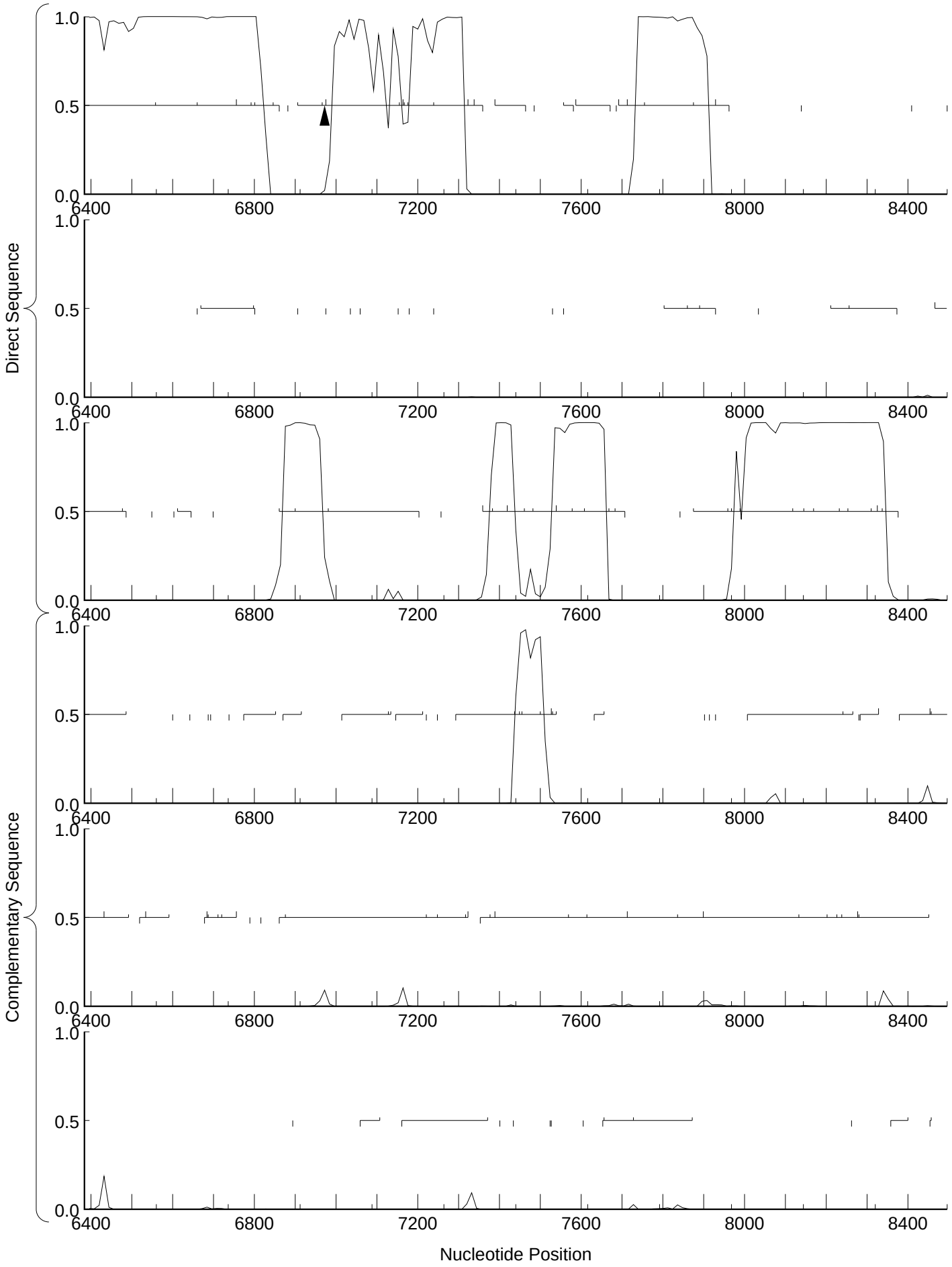
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 3/29



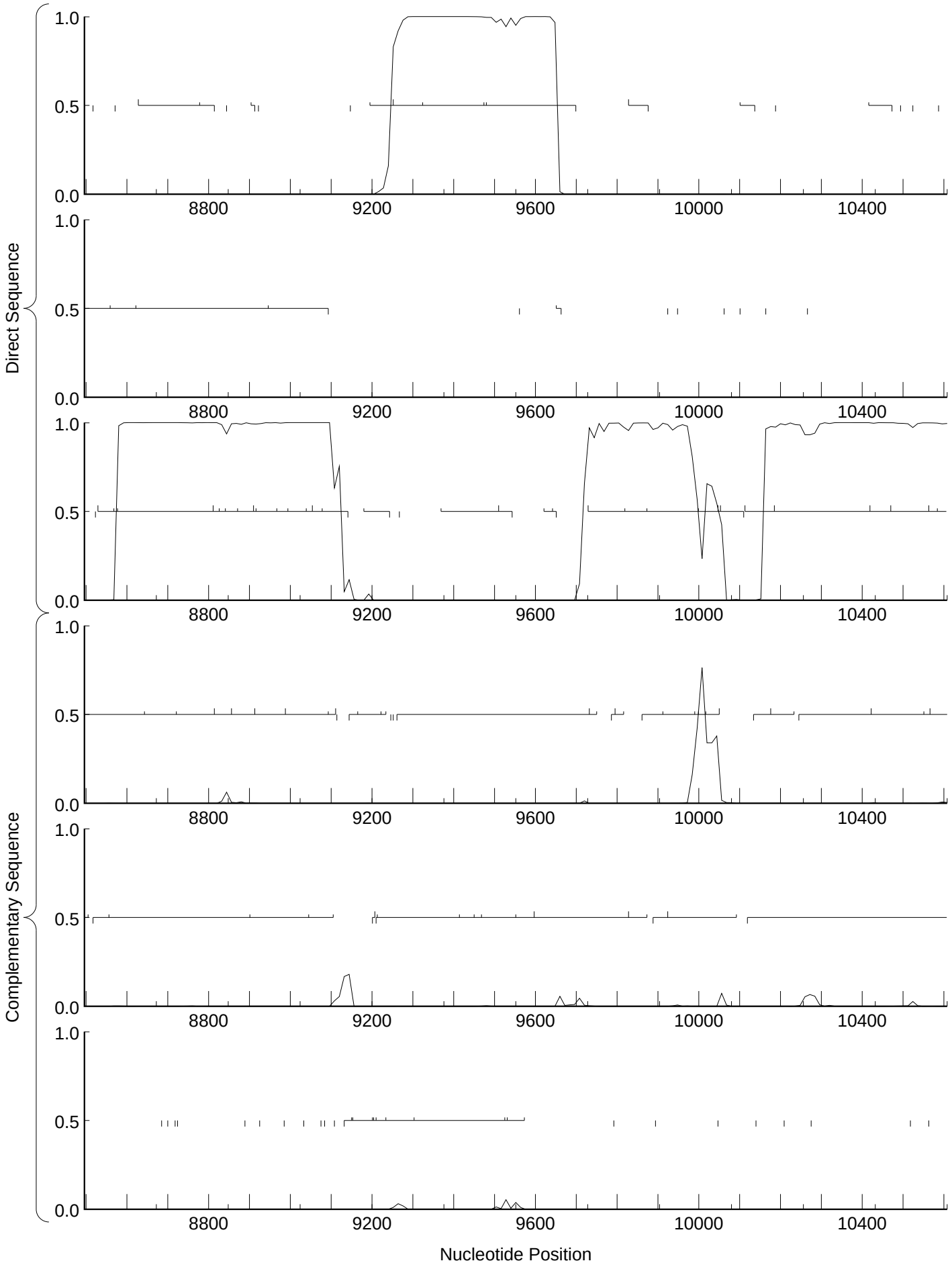
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 4/29



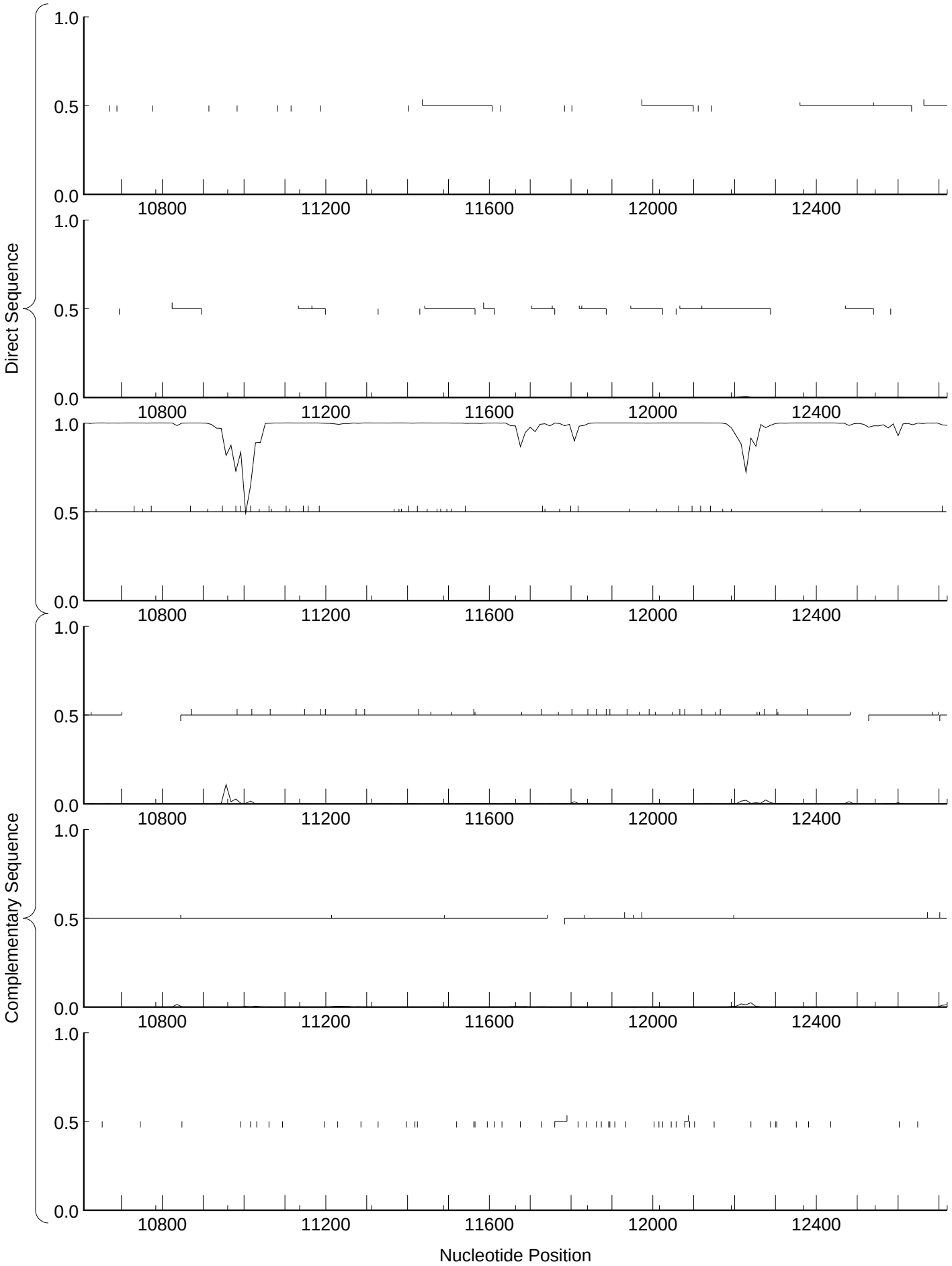
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 5/29



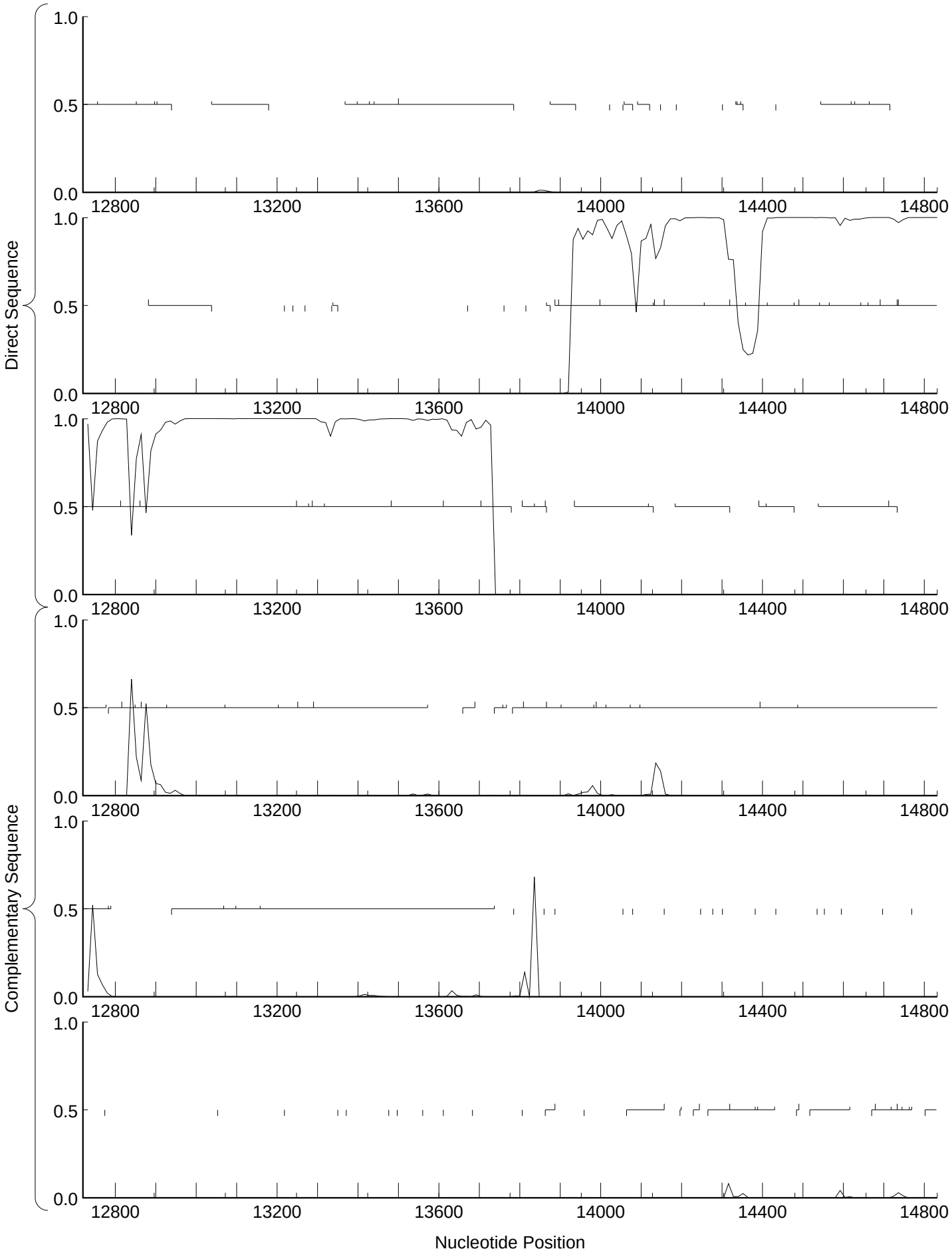
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 6/29



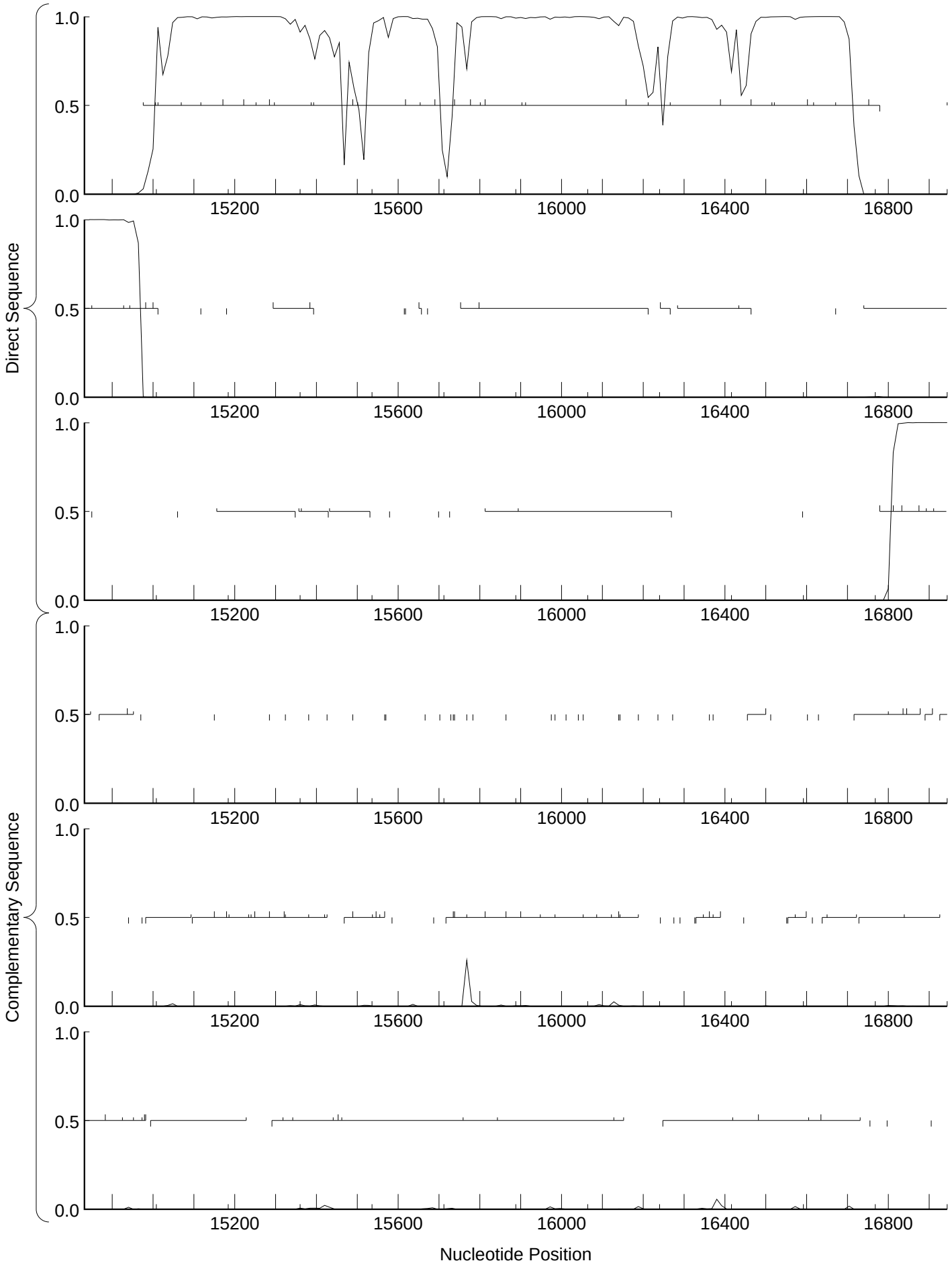
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 7/29



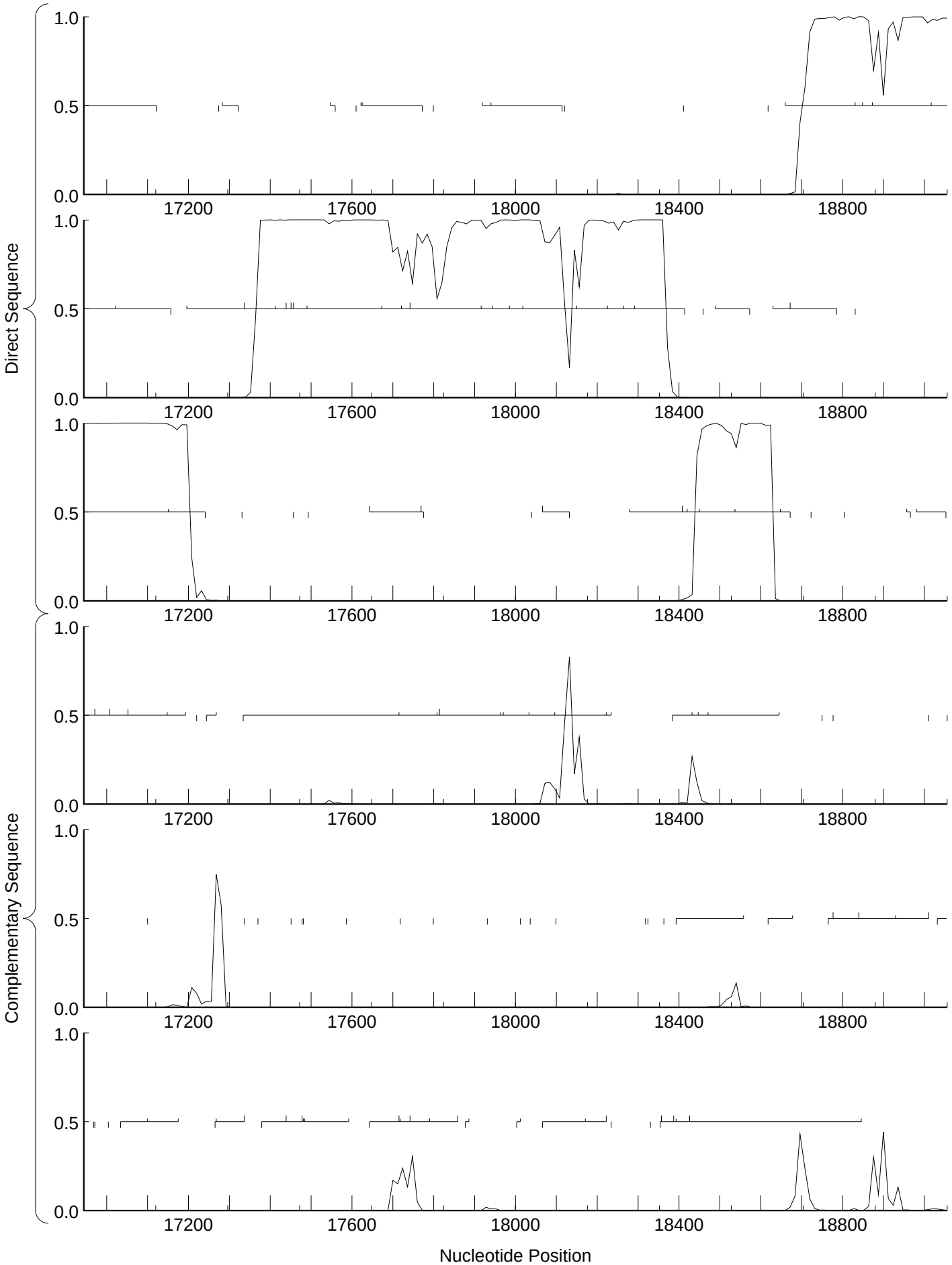
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 8/29



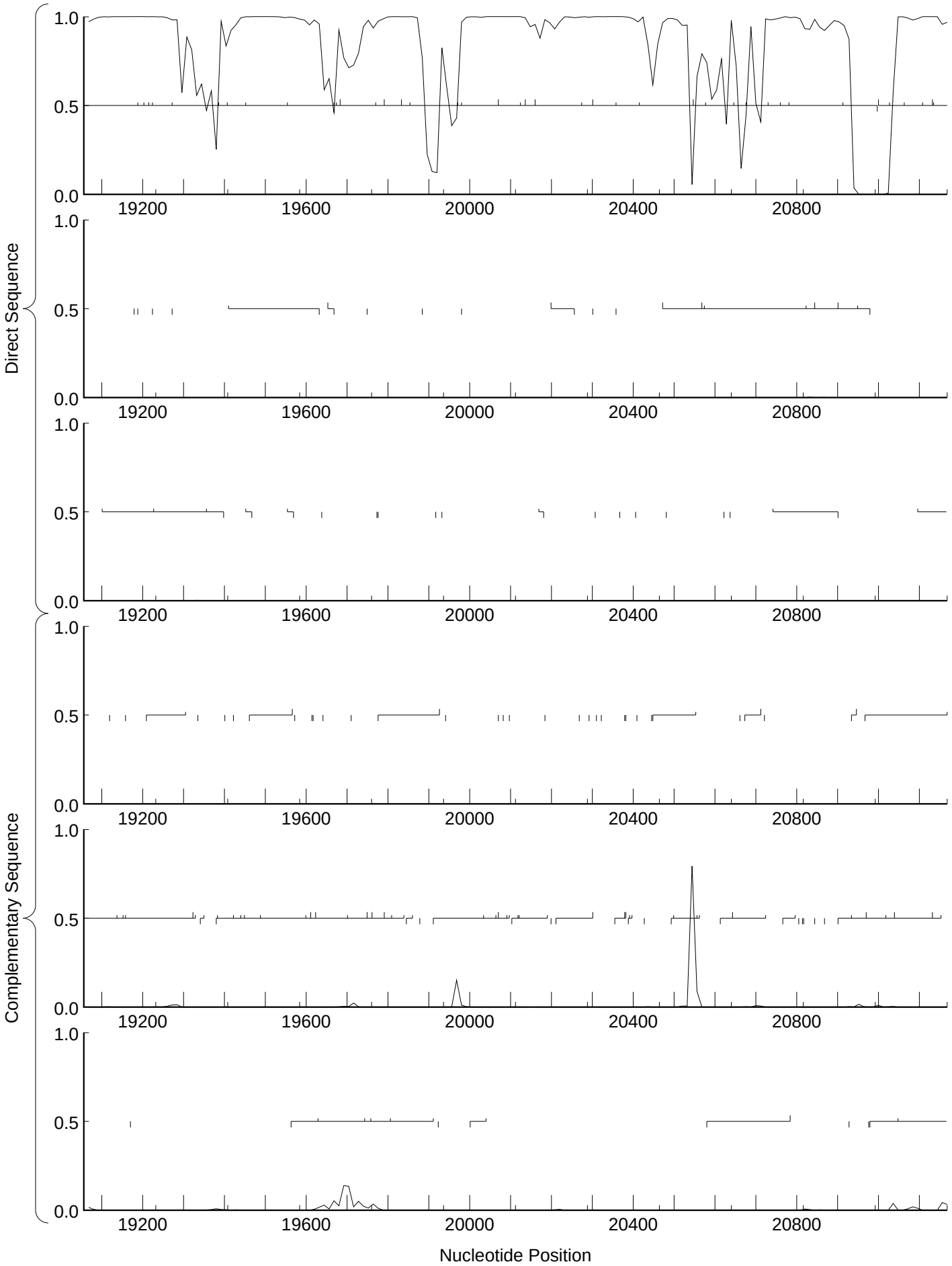
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 9/29



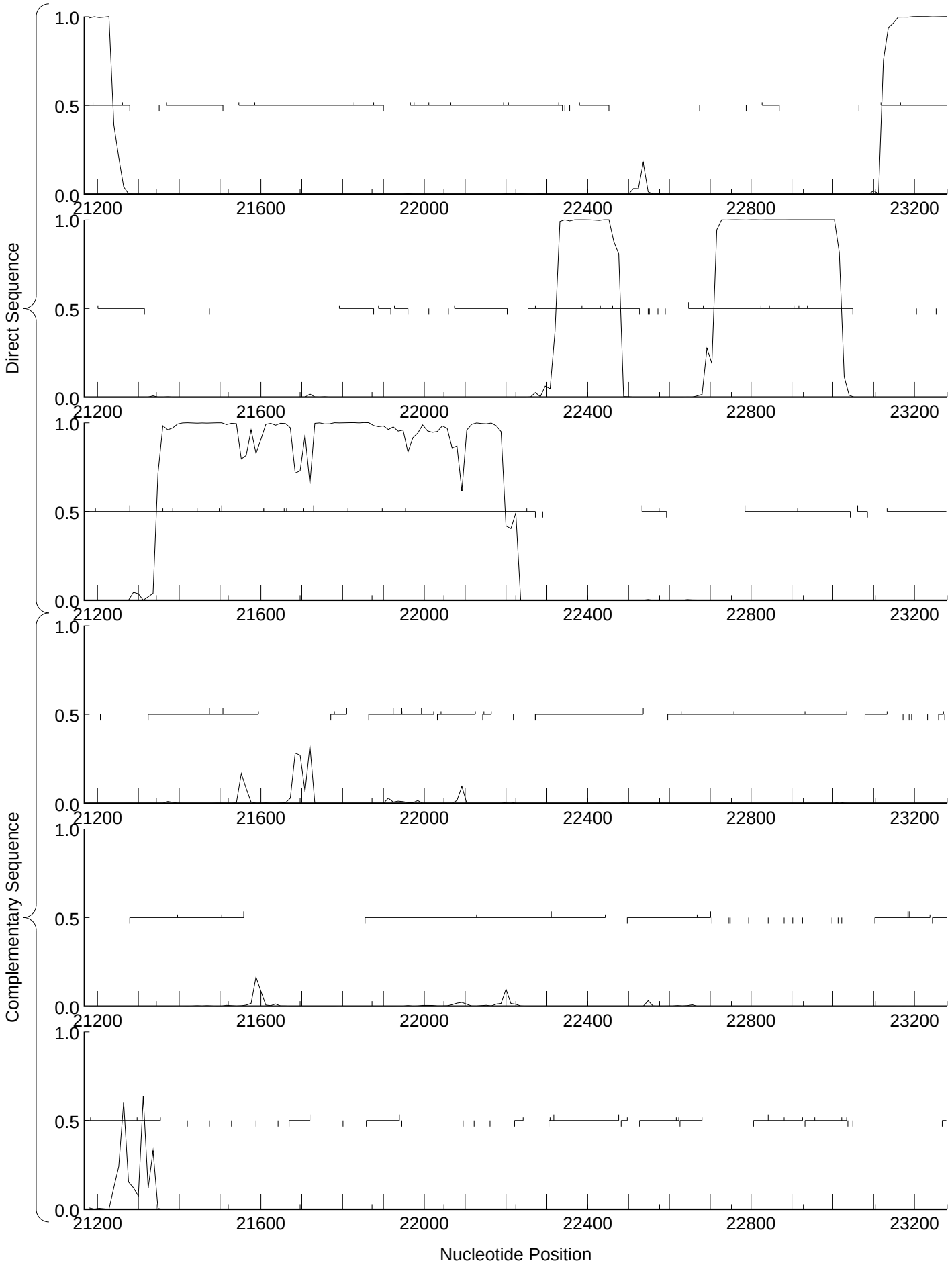
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 10/29



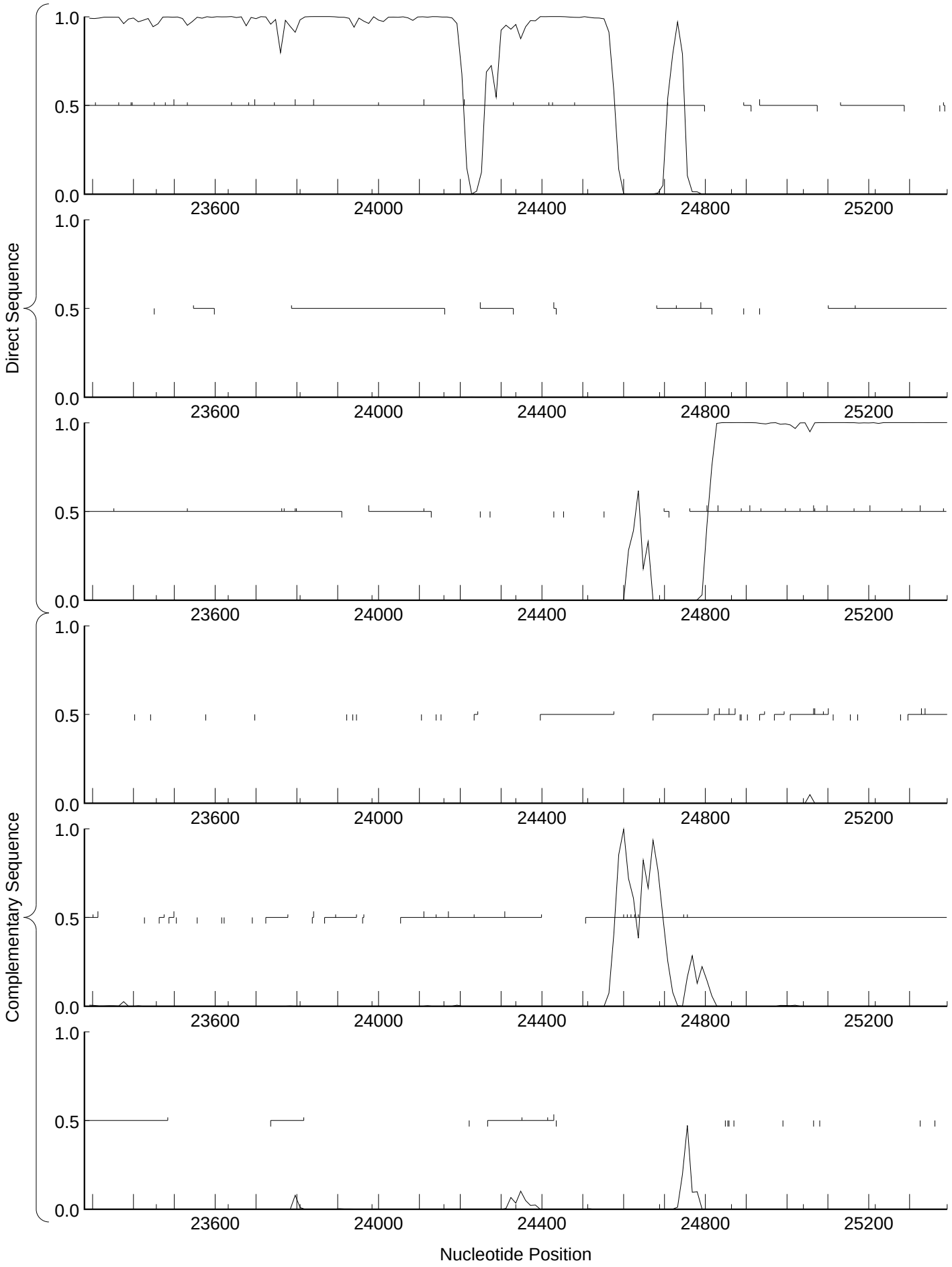
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 11/29



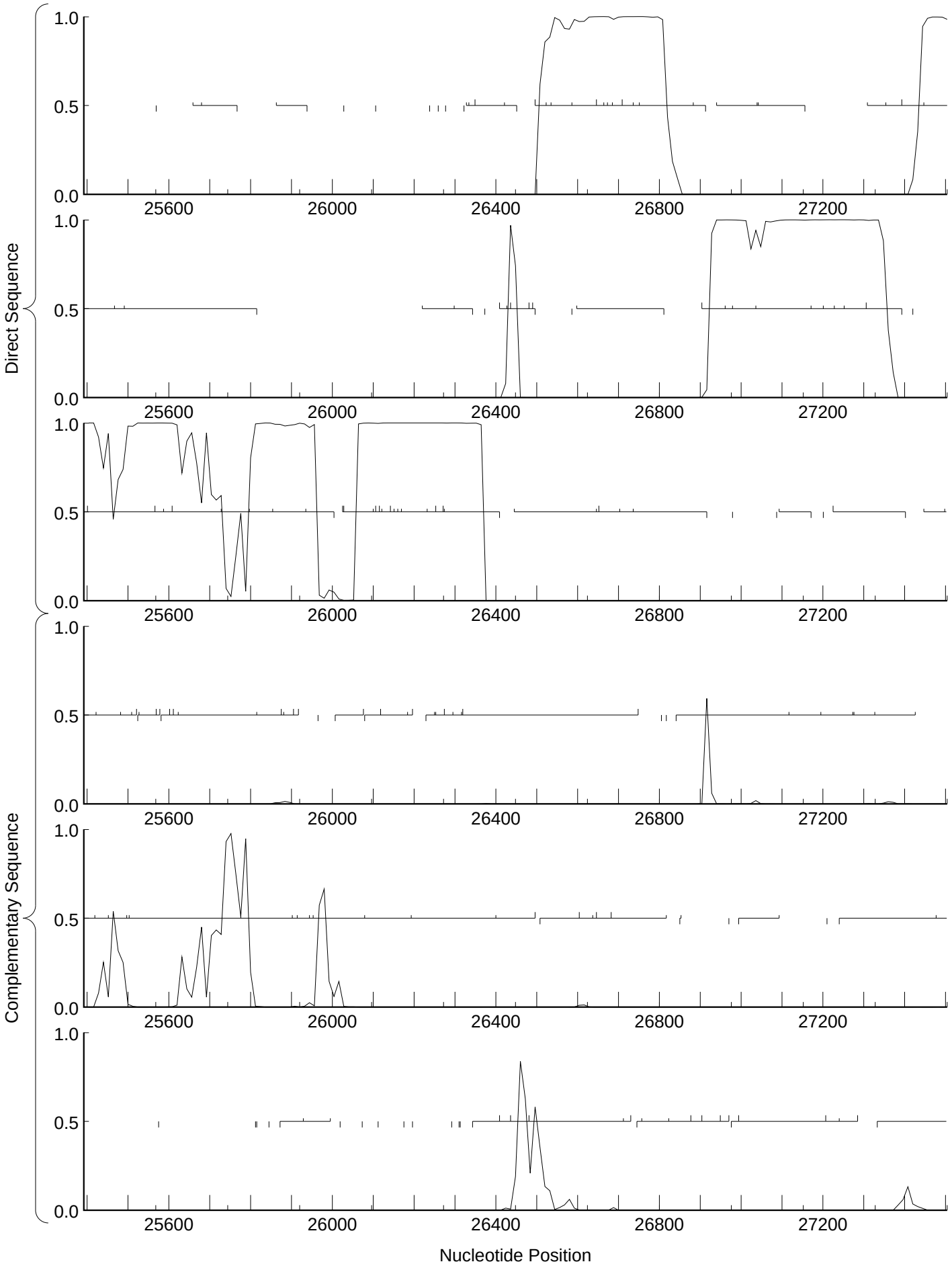
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 12/29



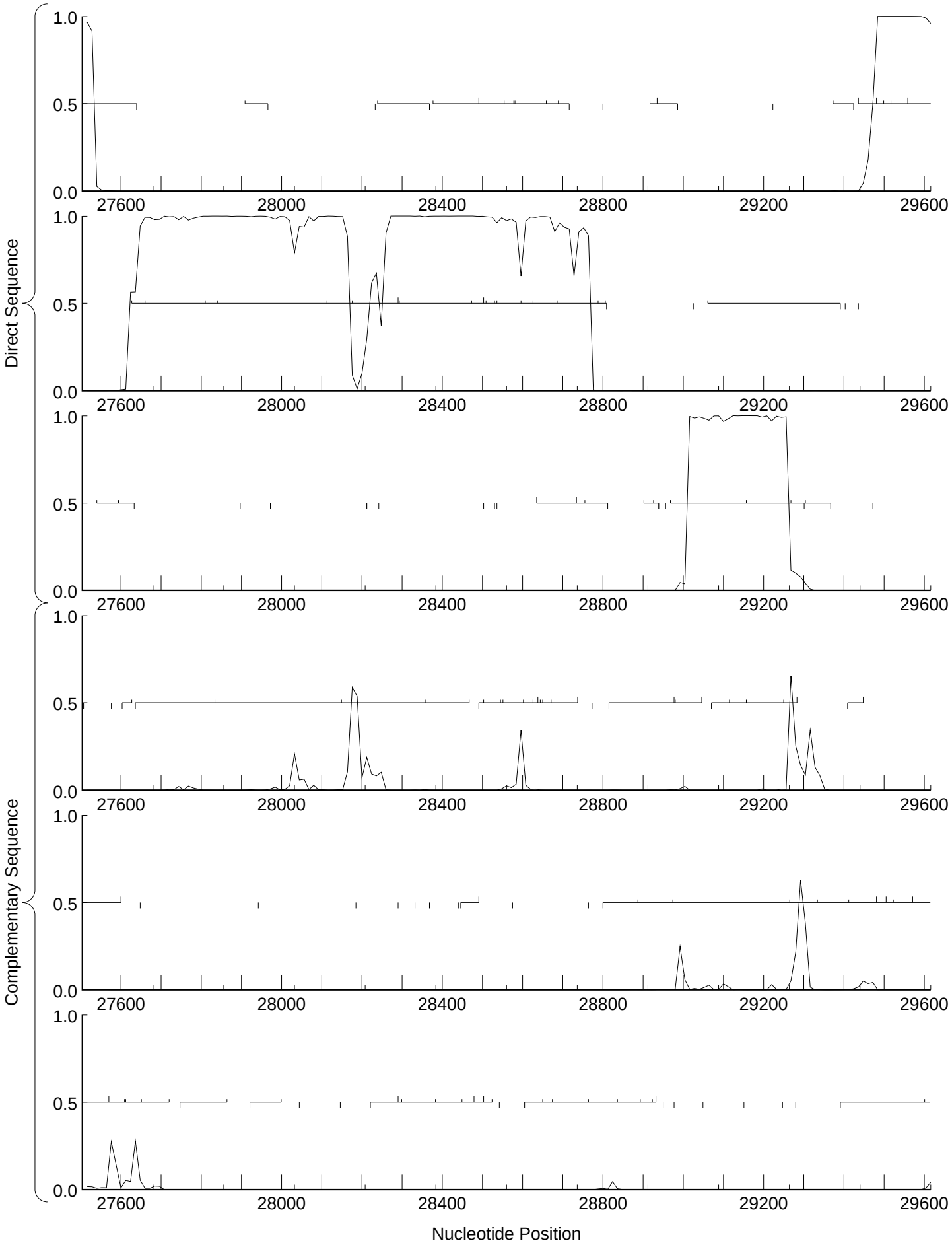
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 13/29



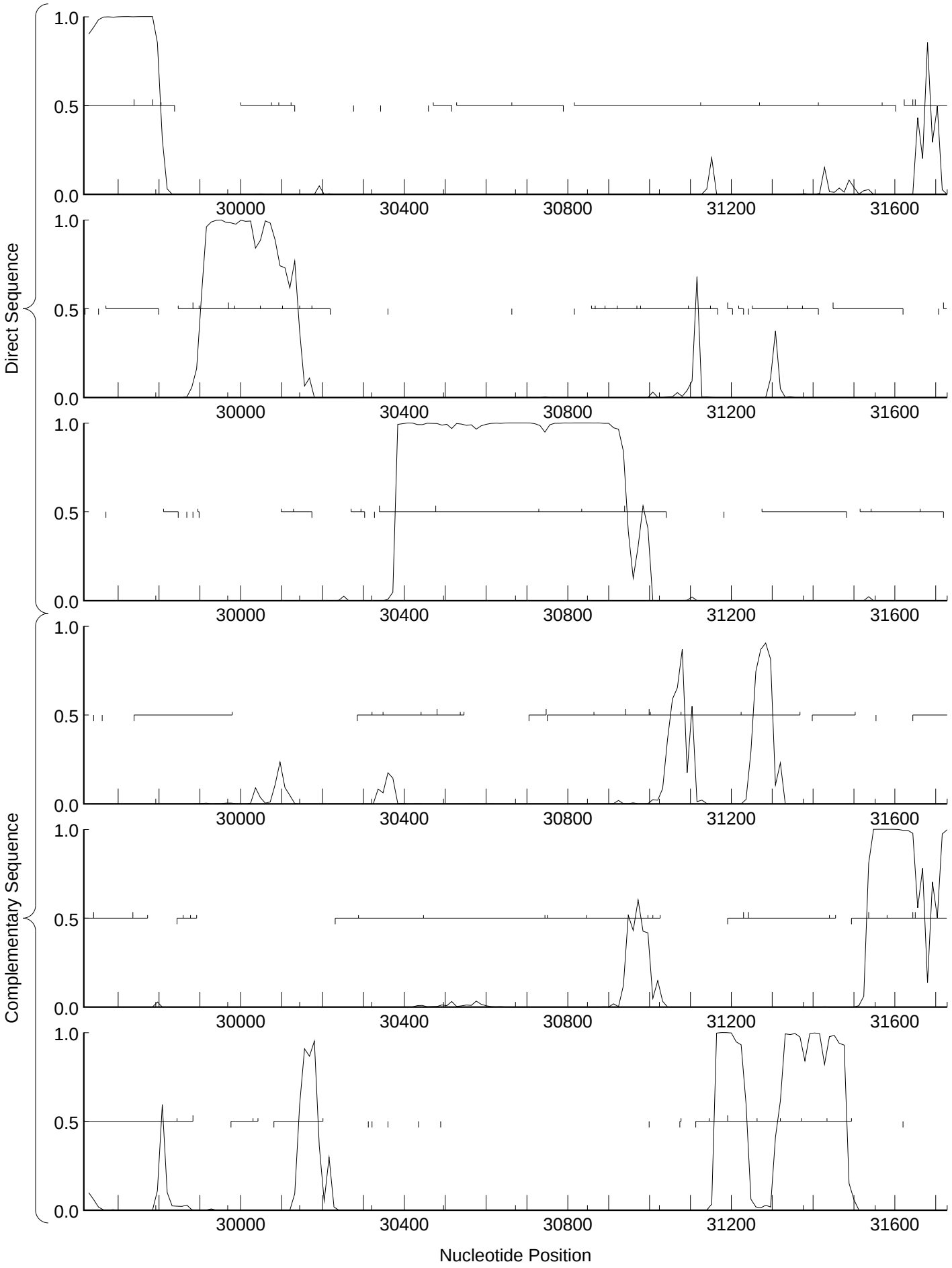
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 14/29



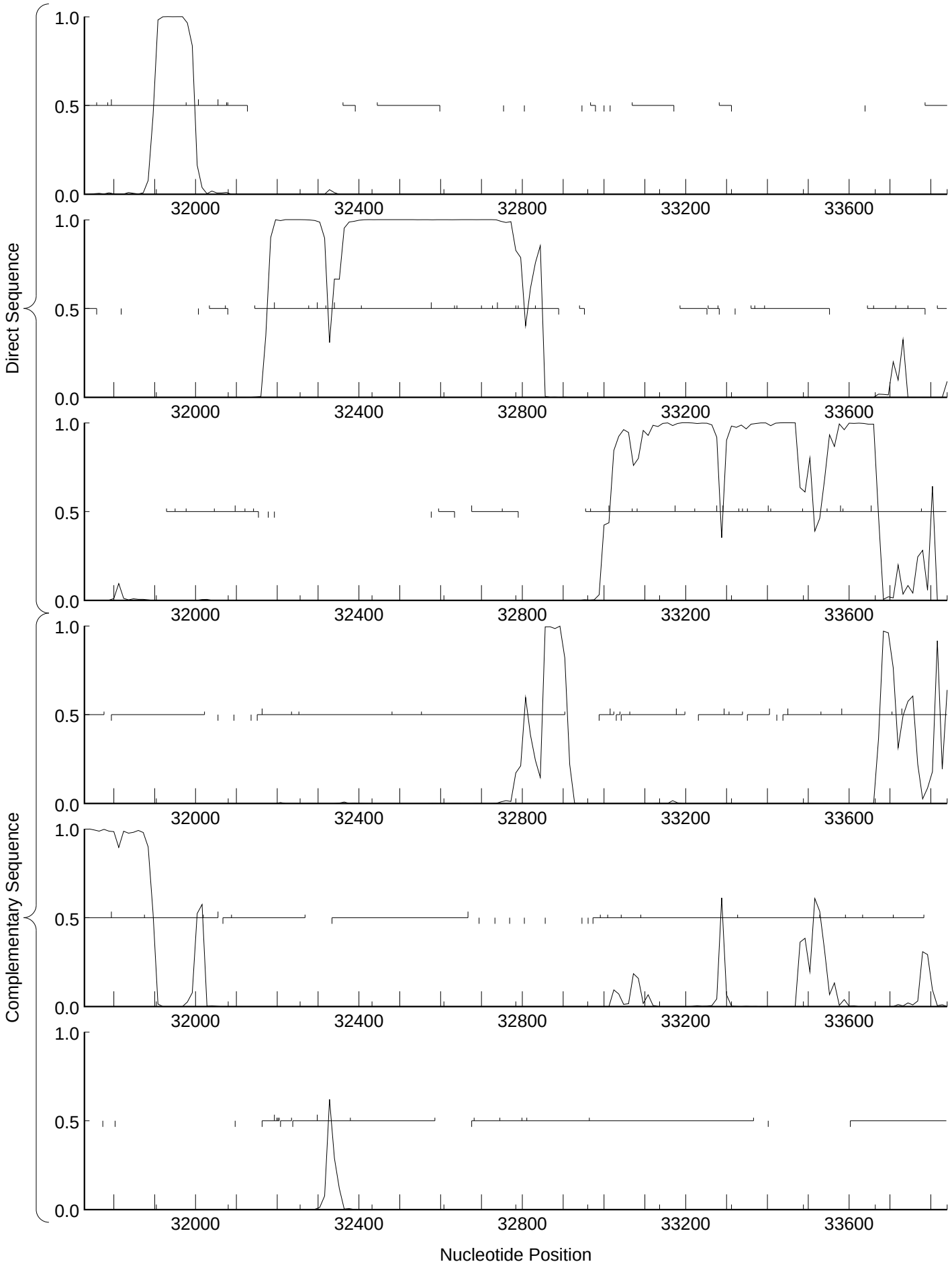
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 15/29



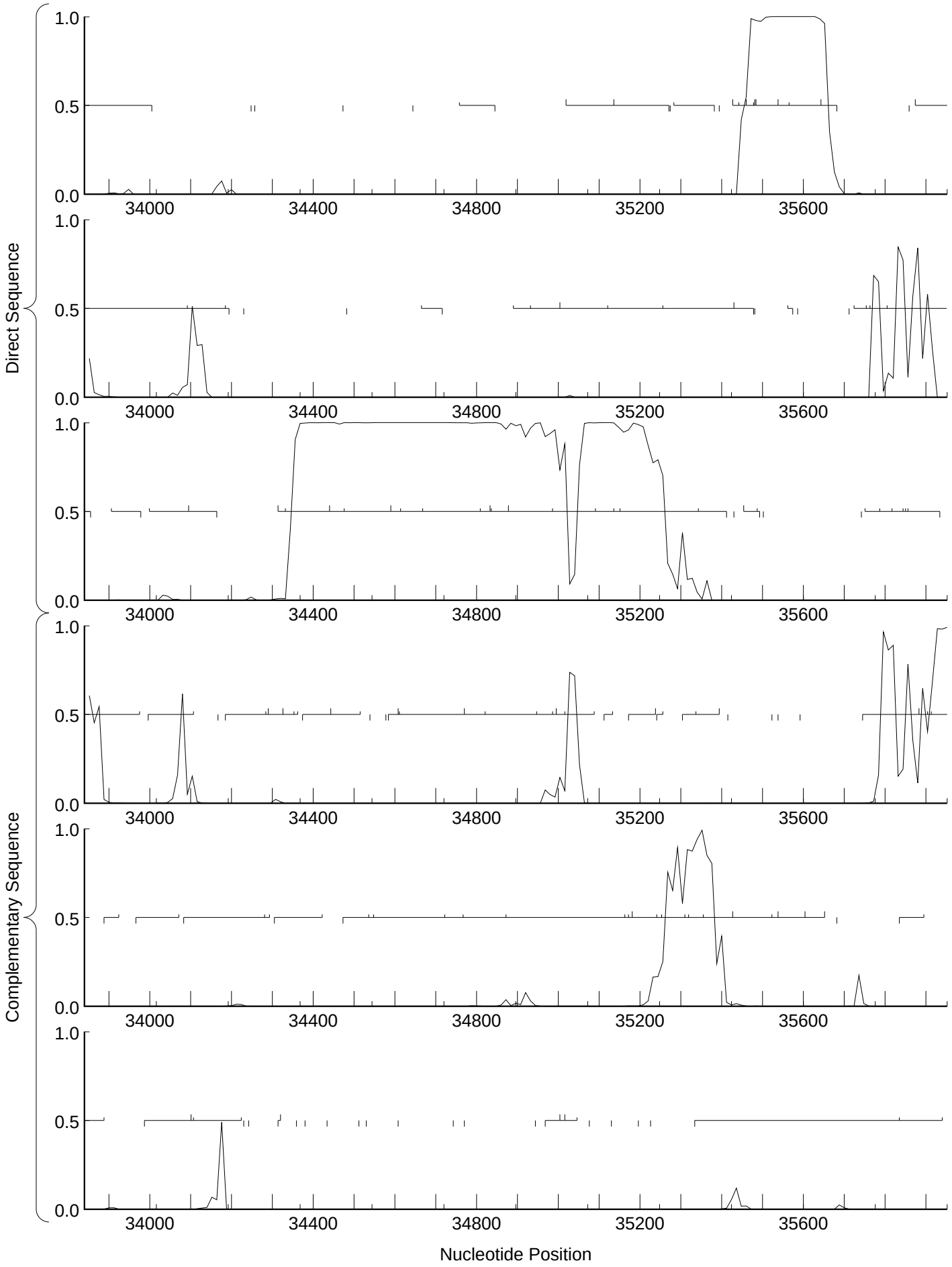
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 16/29



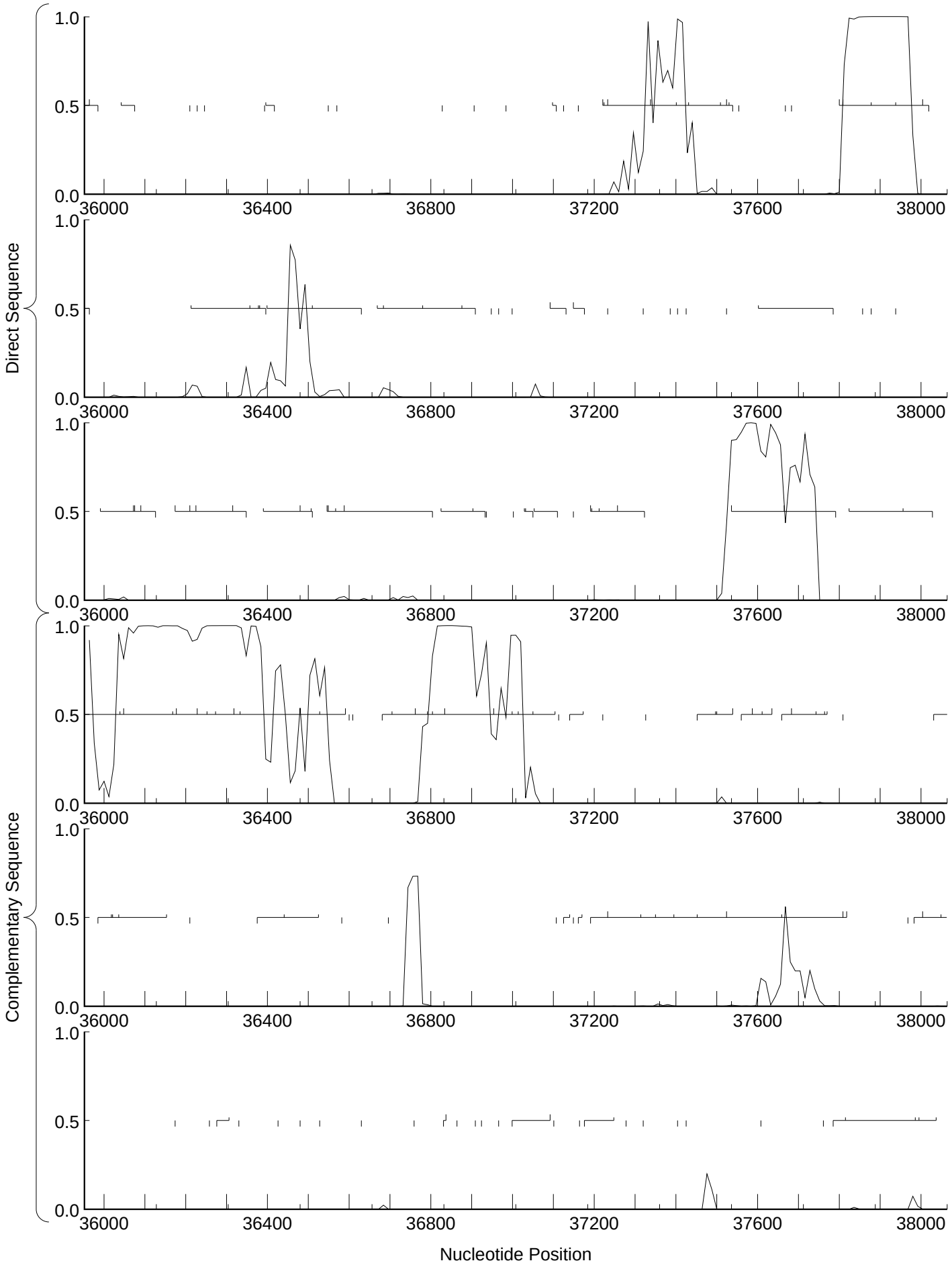
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 17/29



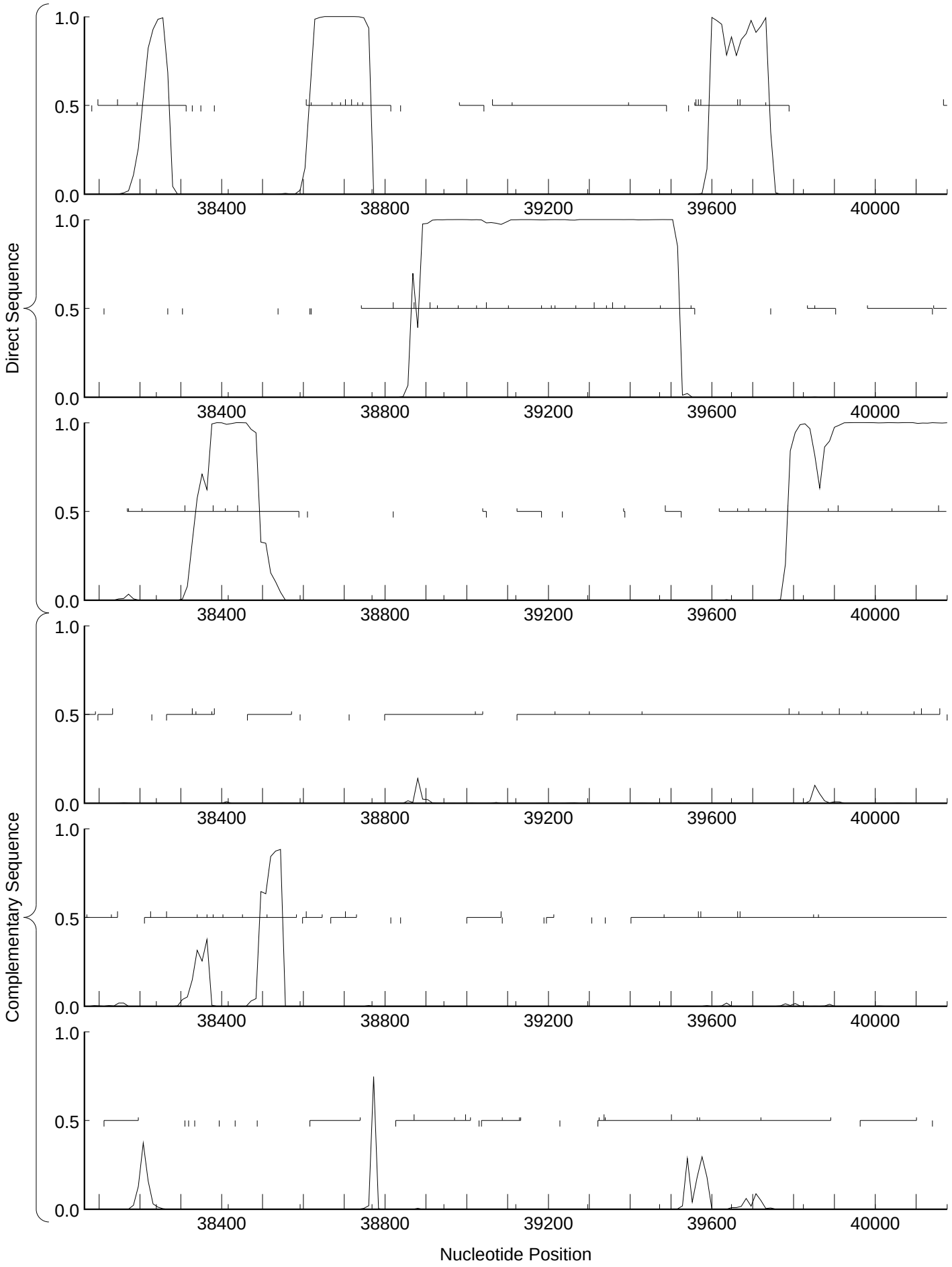
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 18/29



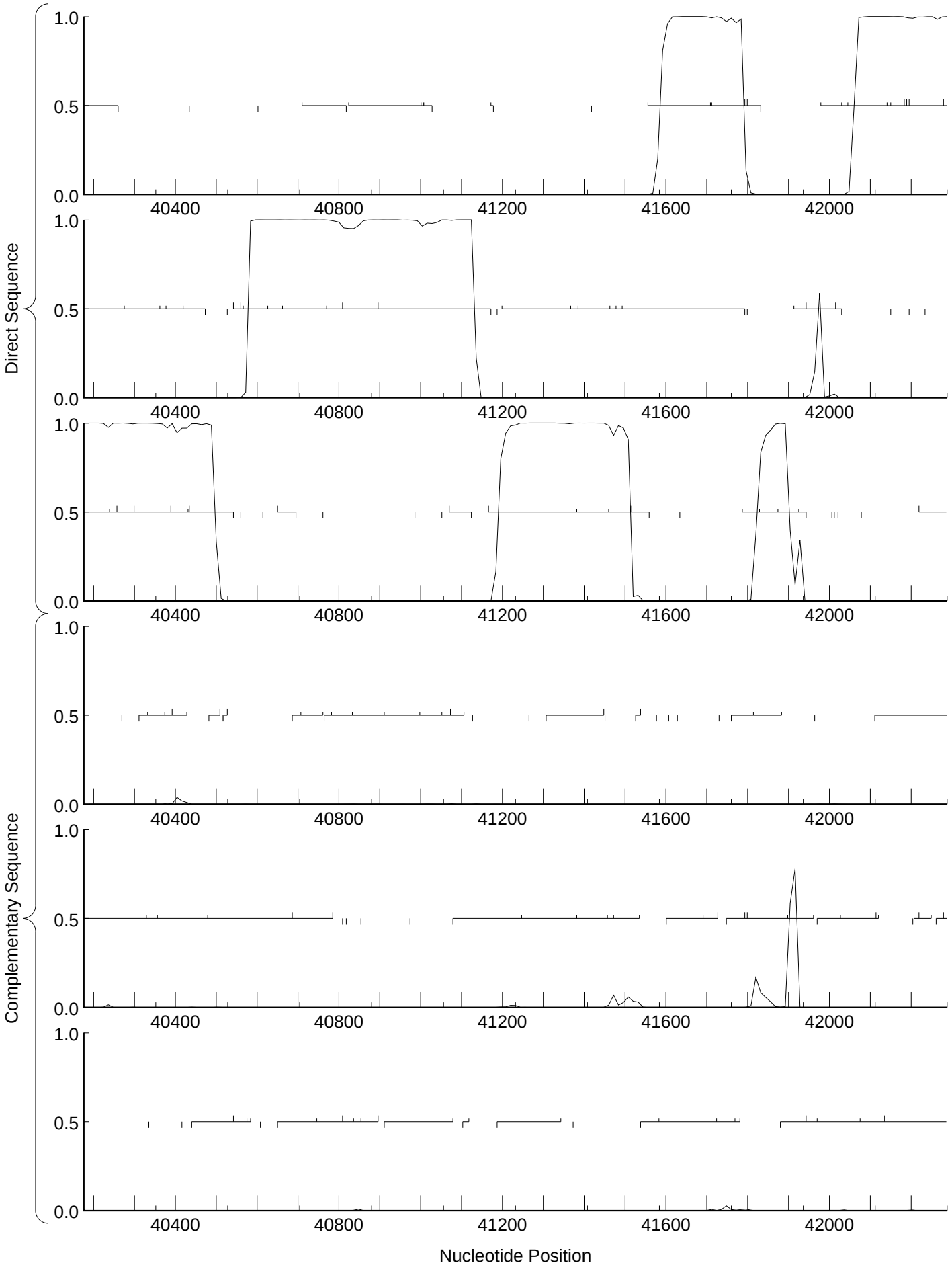
e complete sequence, 58129 bp including 10-base 3' overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 19/29



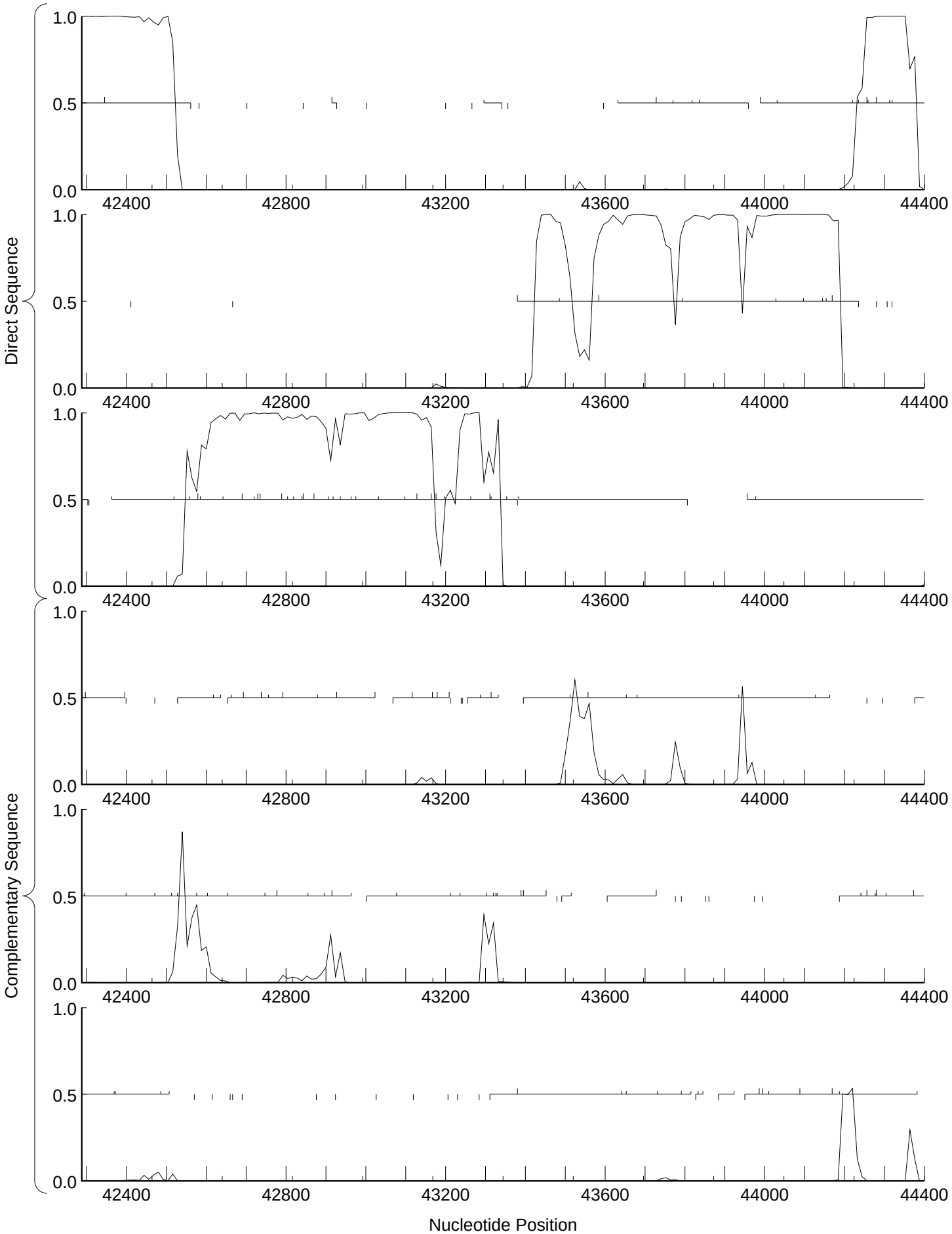
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 20/29



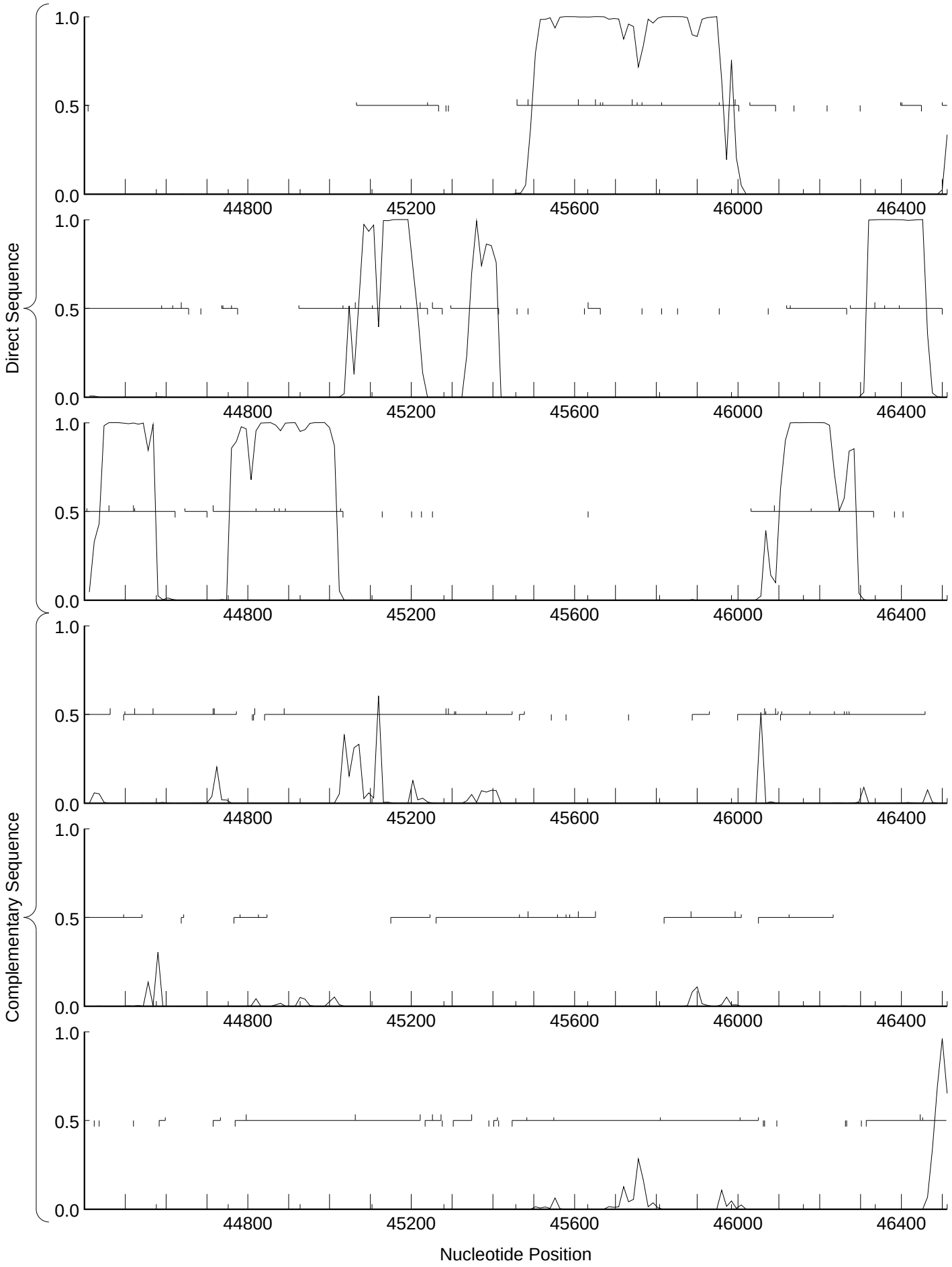
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 21/29



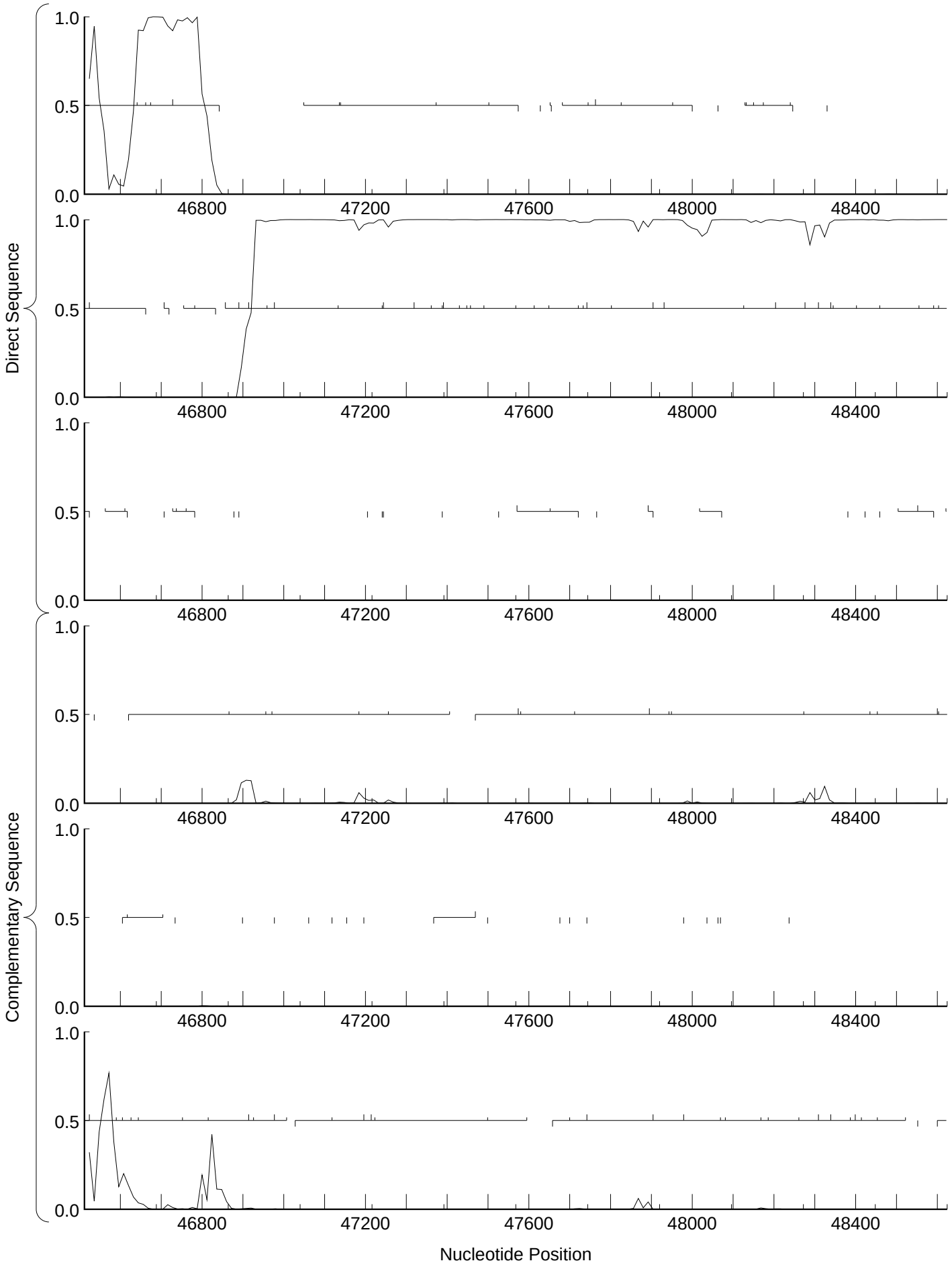
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 22/29



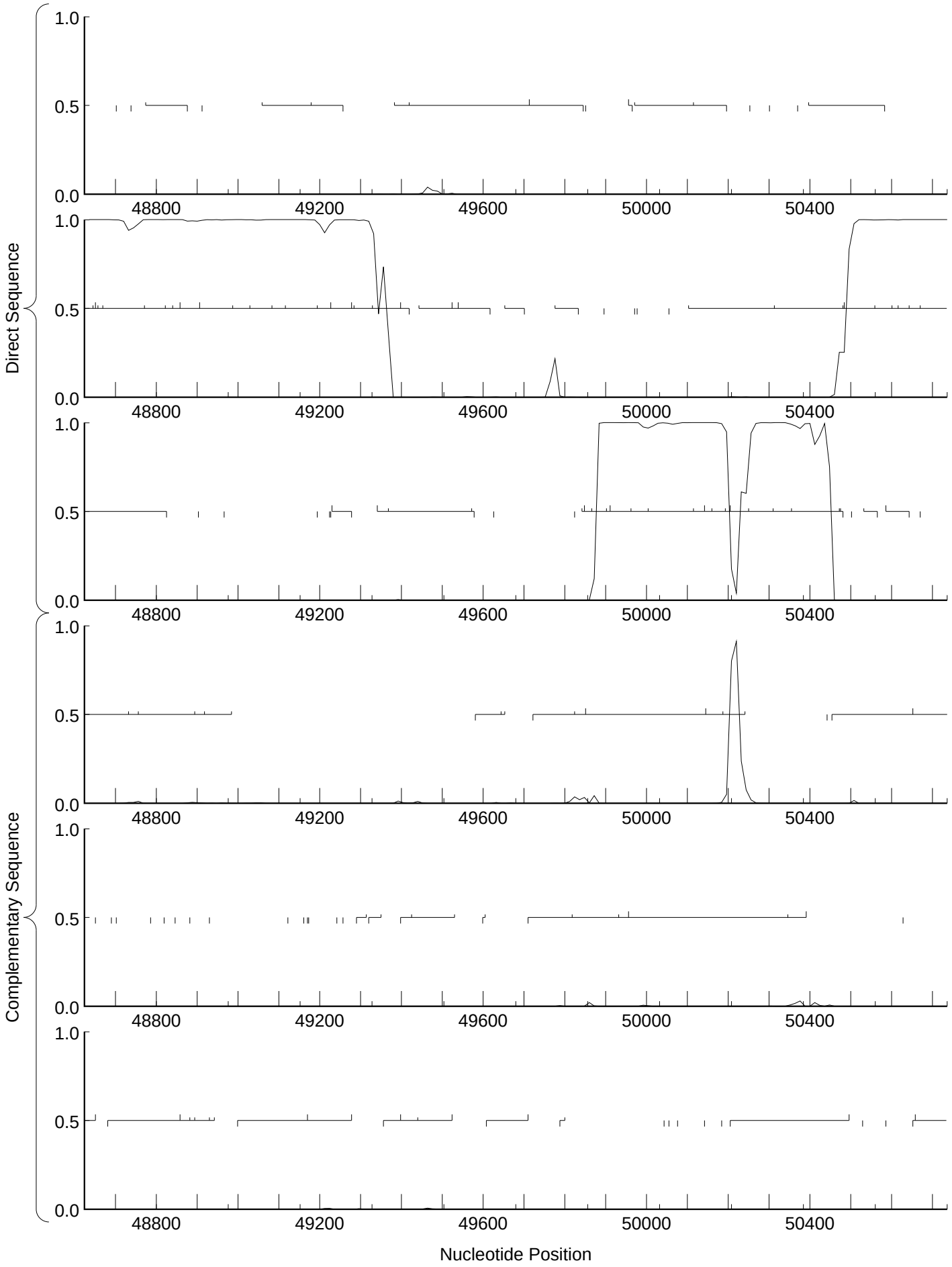
e complete sequence, 58129 bp including 10-base 3' overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 23/29



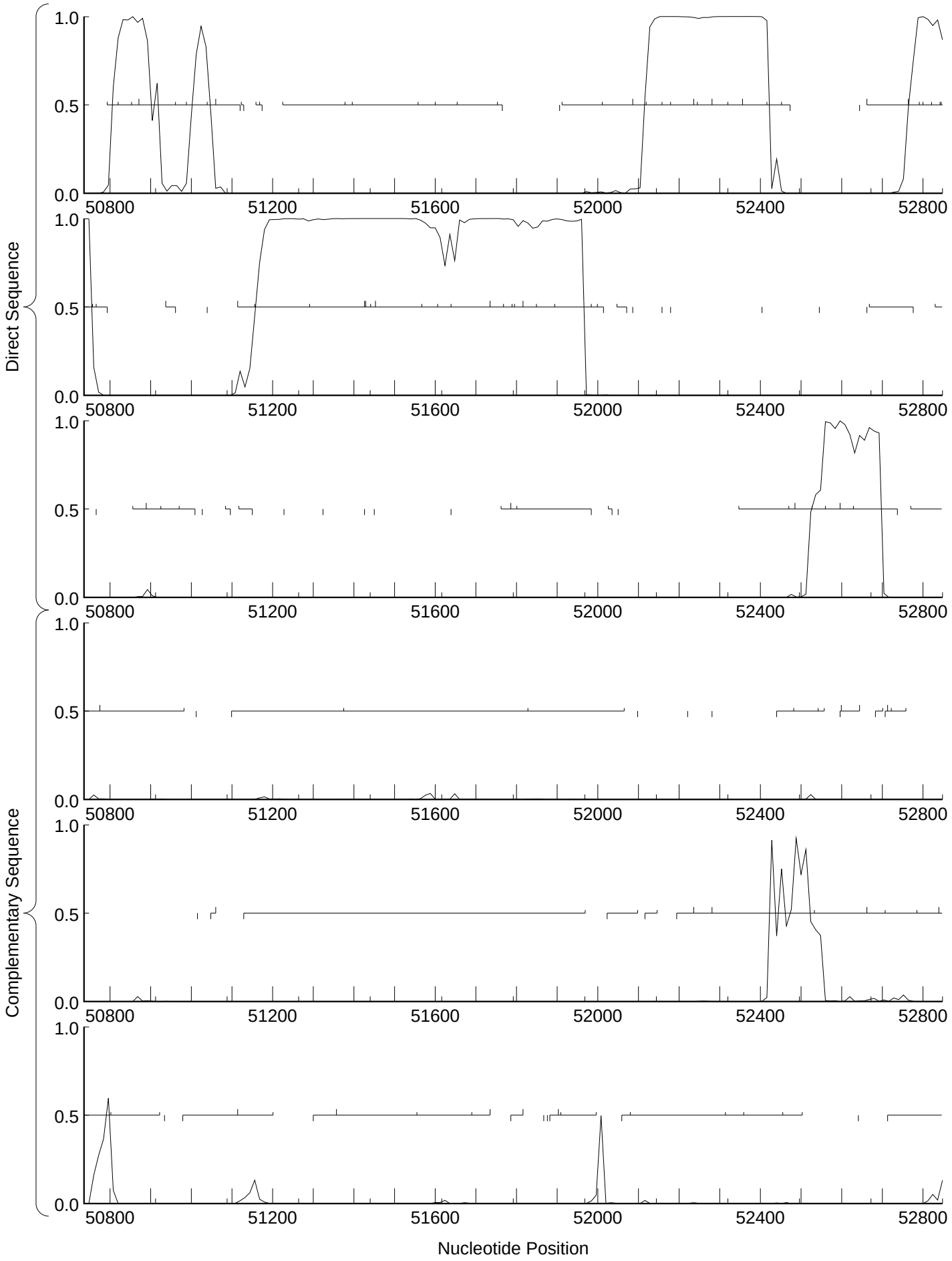
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 24/29



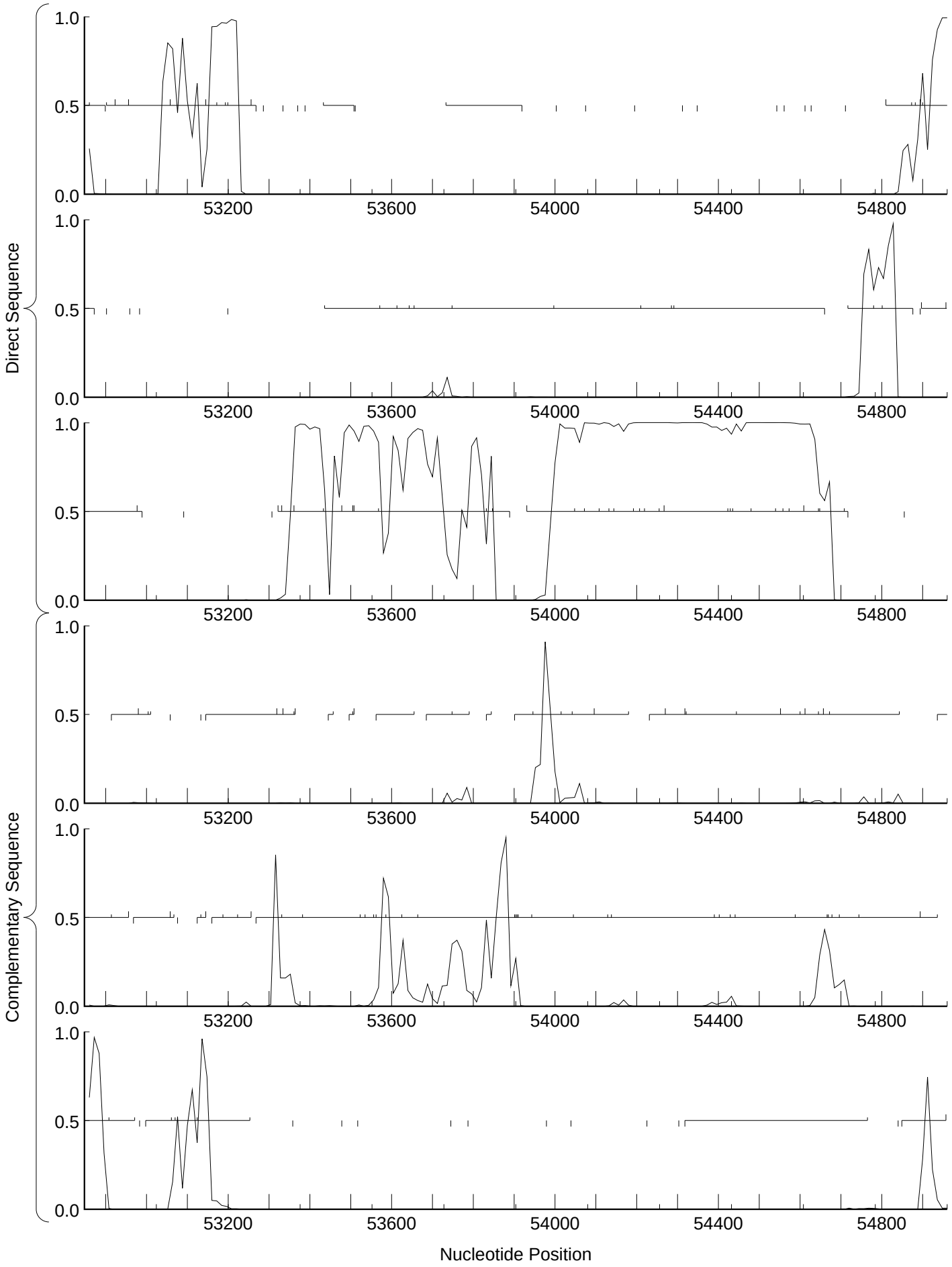
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 25/29



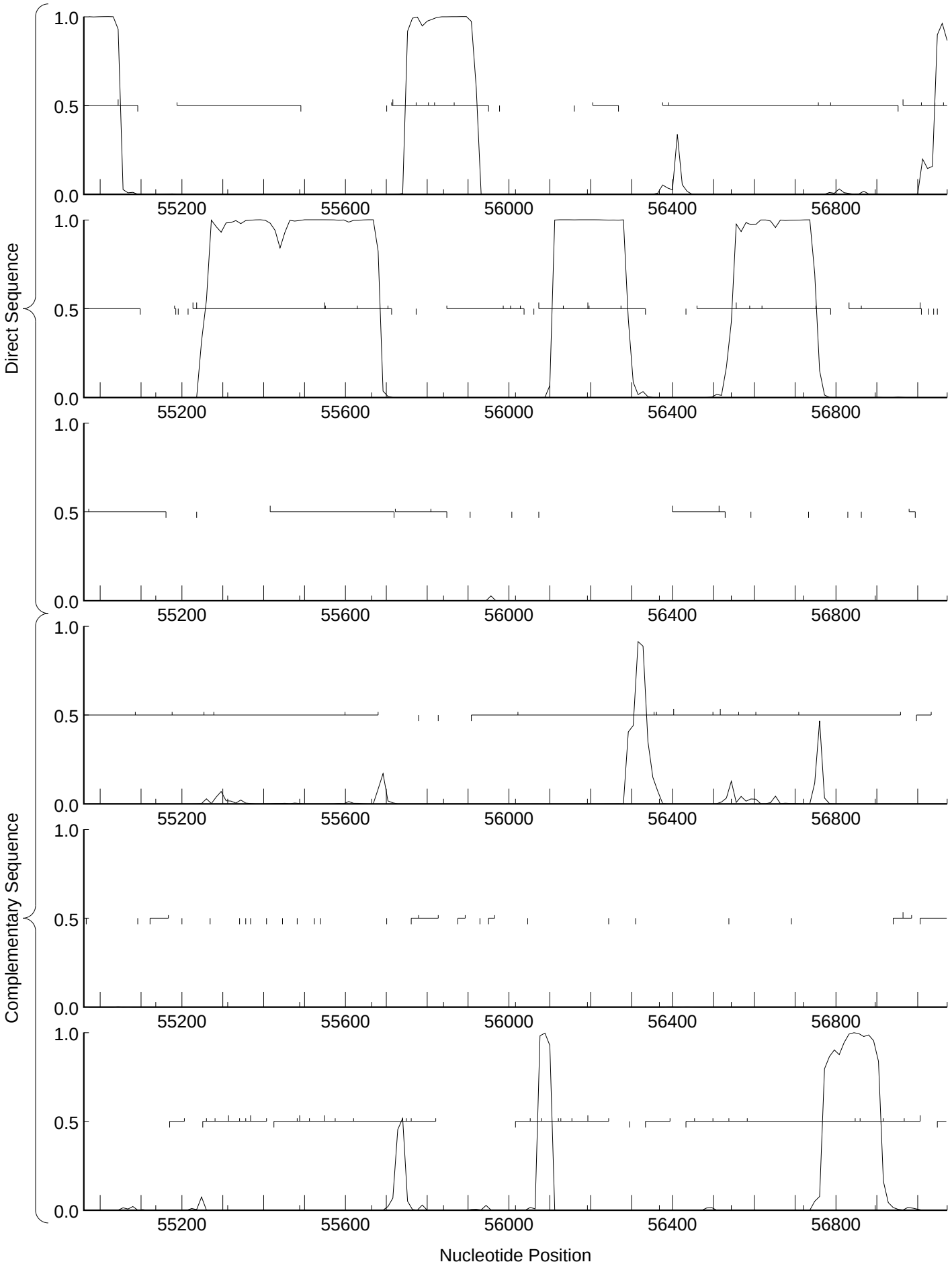
e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 26/29



e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 27/29



e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 28/29



e complete sequence, 58129 bp including 10-base 3’ overhang (CTCGCGGCAT), Cluster K2, Order 4, Window 96, Step 12, 29/29

