GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Edugator complete sequence, 63344 including 11-base 3' overhang (CTCAGT
Analysis Date    :  3/28/24 at 18:23:53
Pages            :  31
Sequence Length  :  63344 bp
GC Content       :   65.33%

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

PS-Version       :  1.2

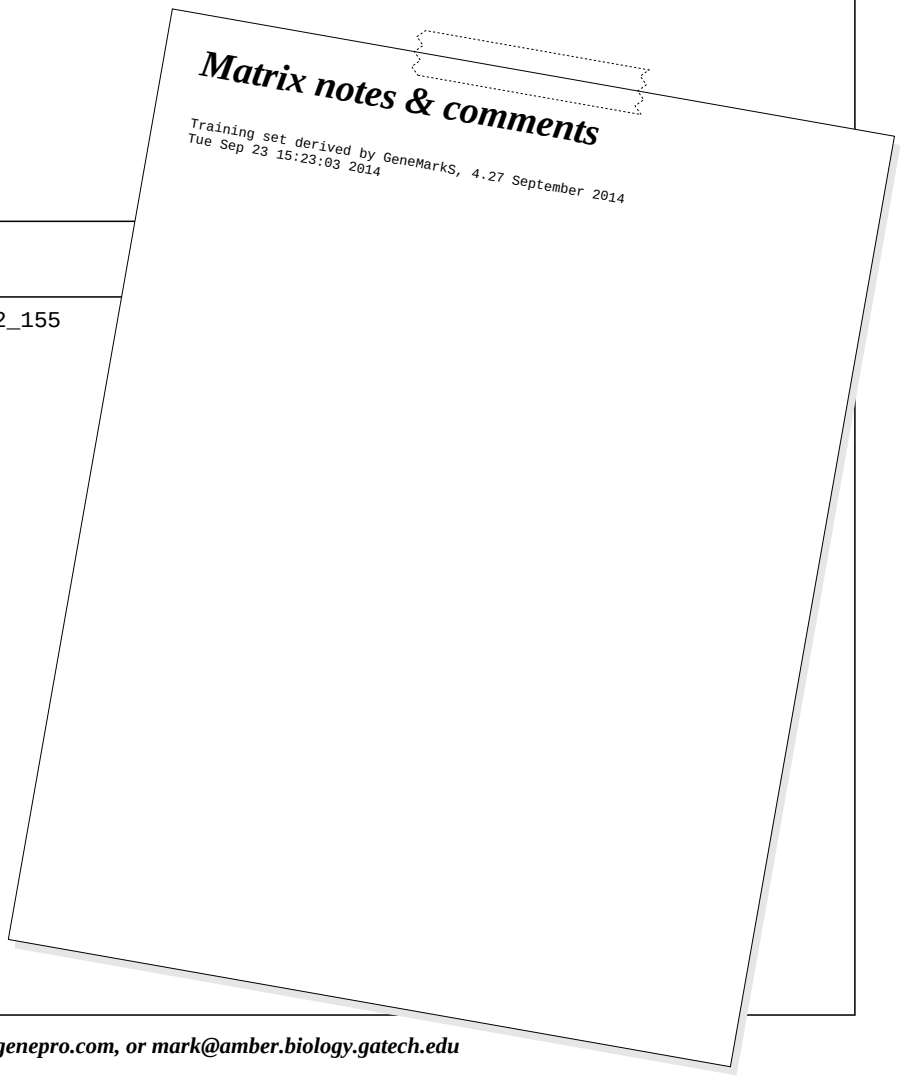
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
```

MATRIX INFORMATION

```
Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

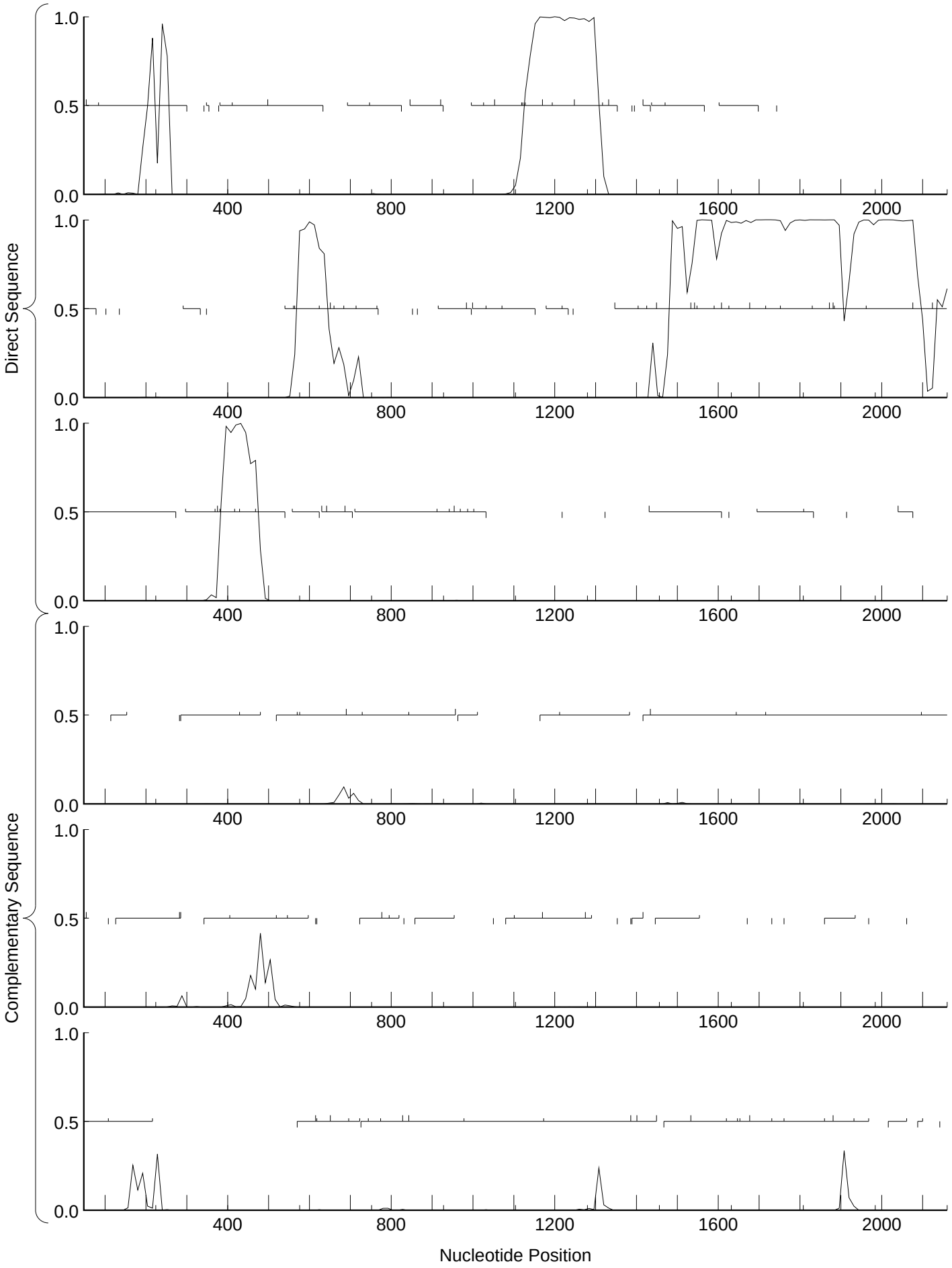
Order      :  4


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

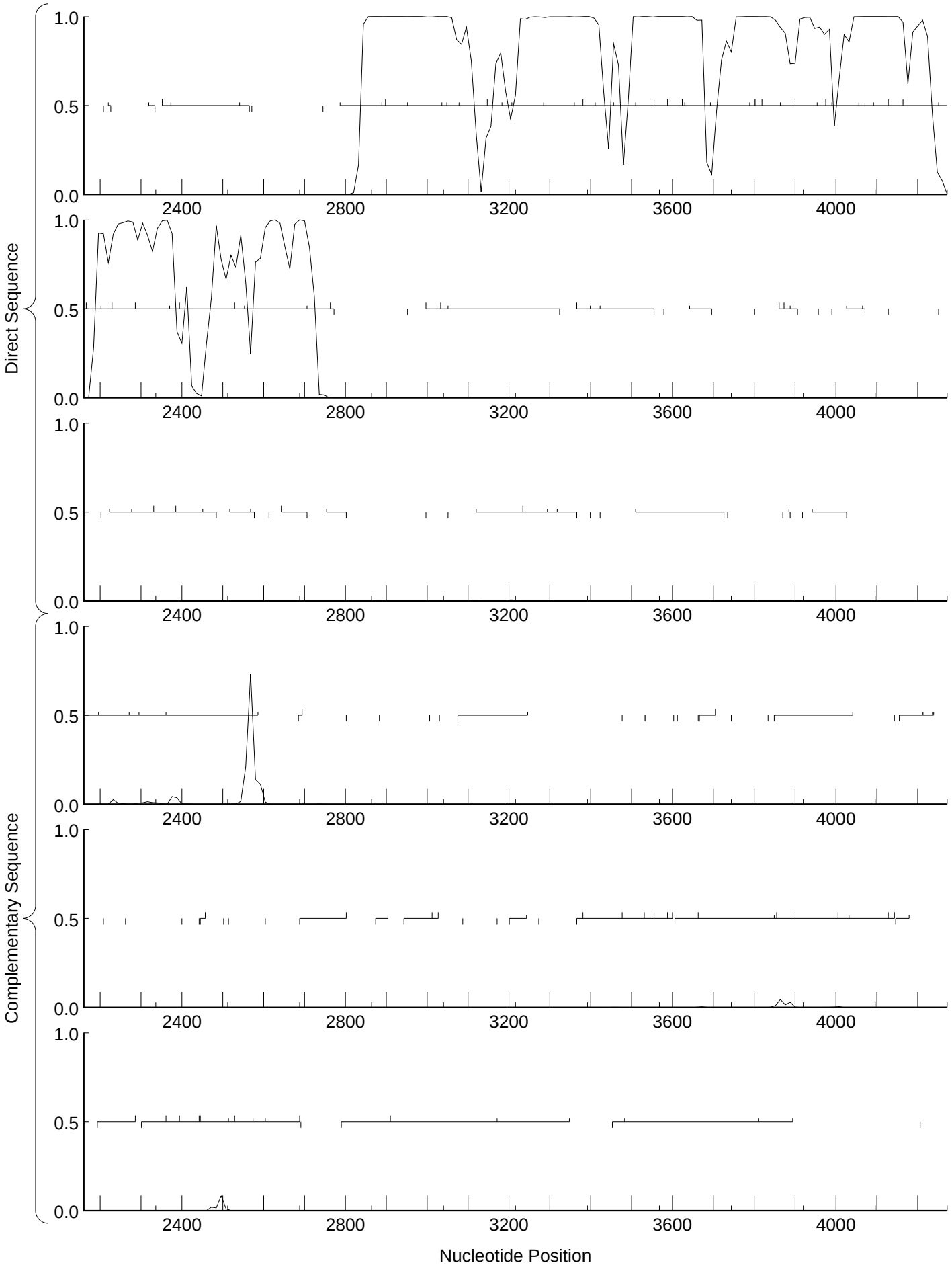


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

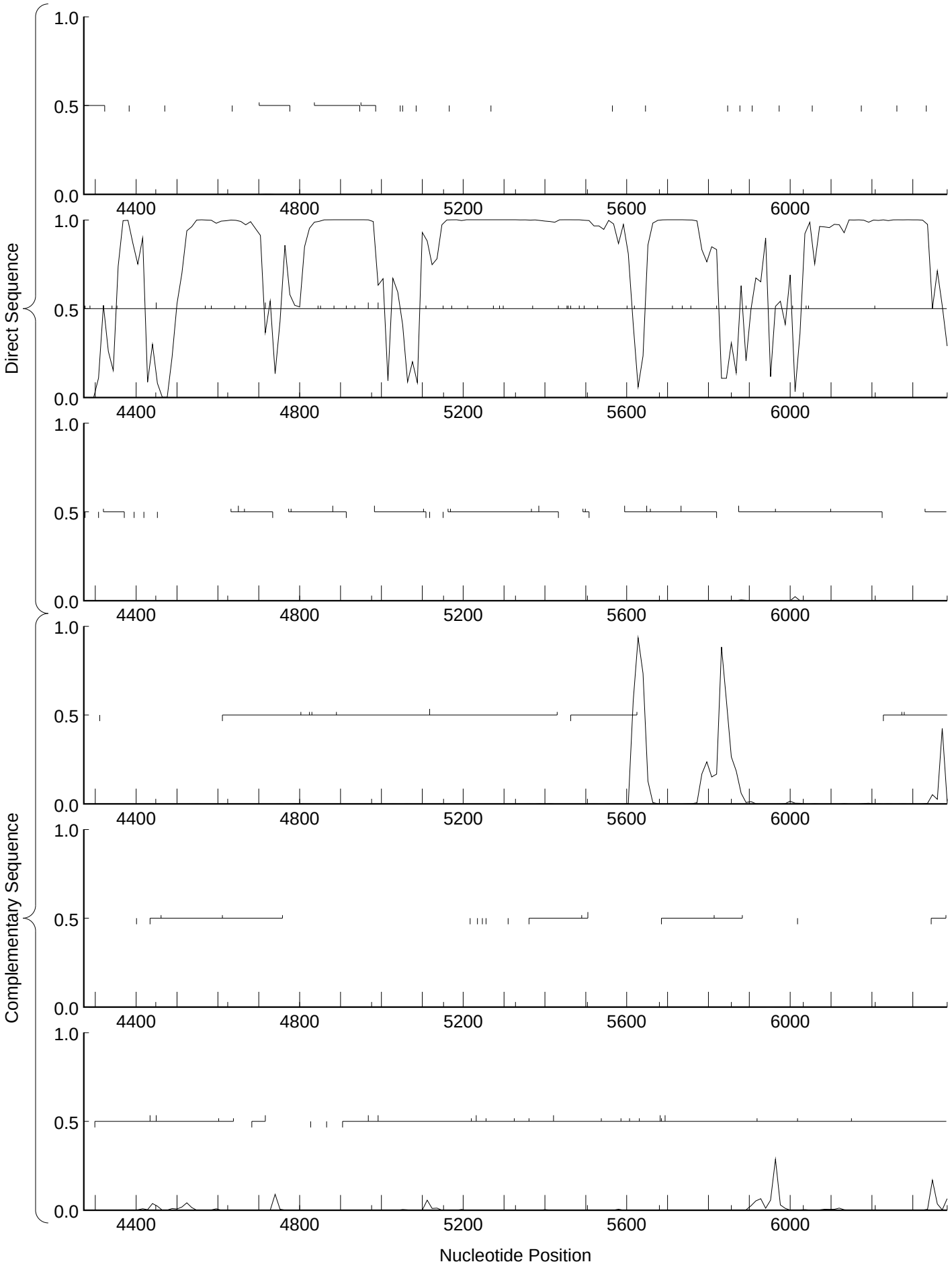
ator complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 2/31



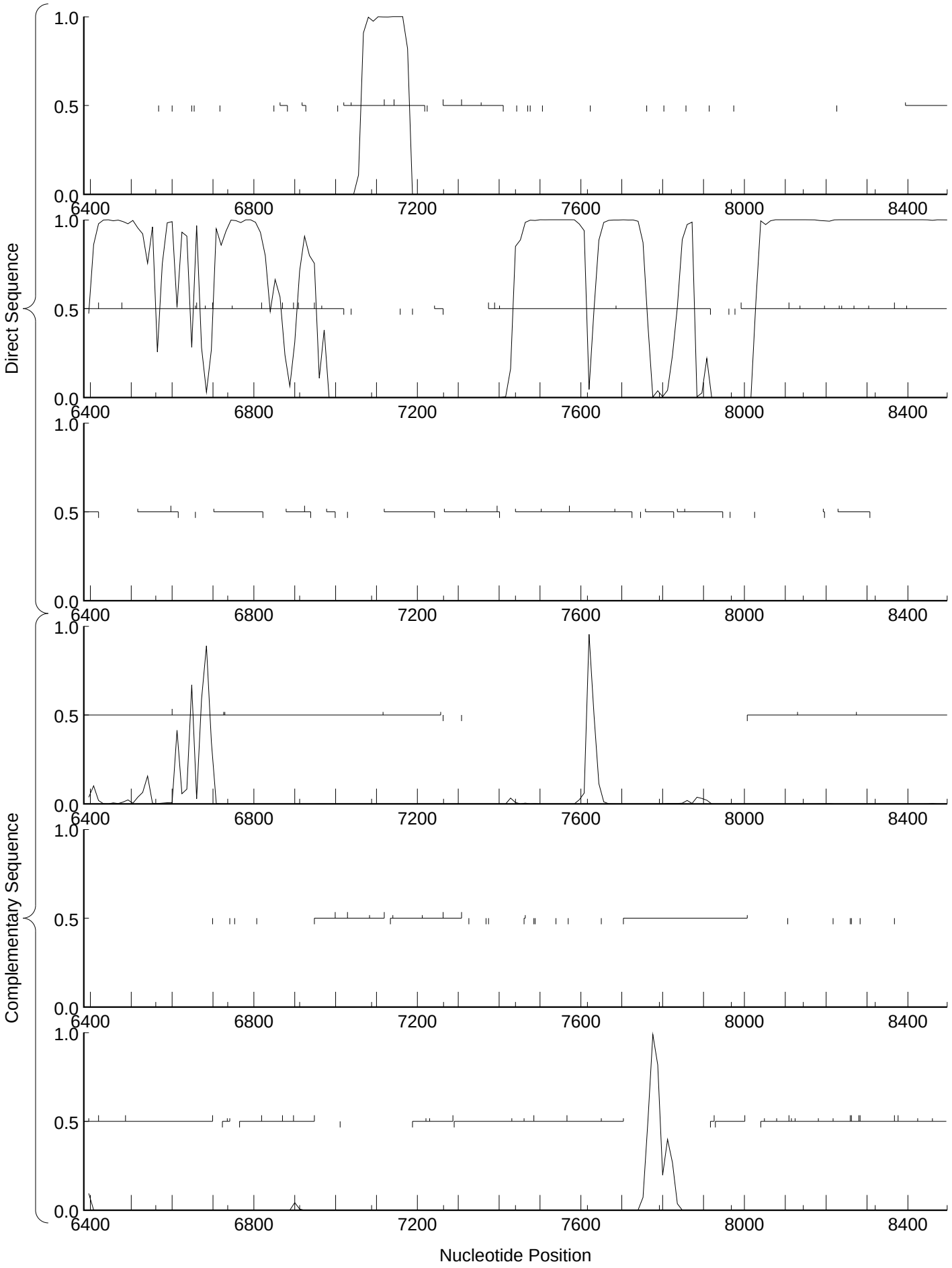
ator complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 3/31



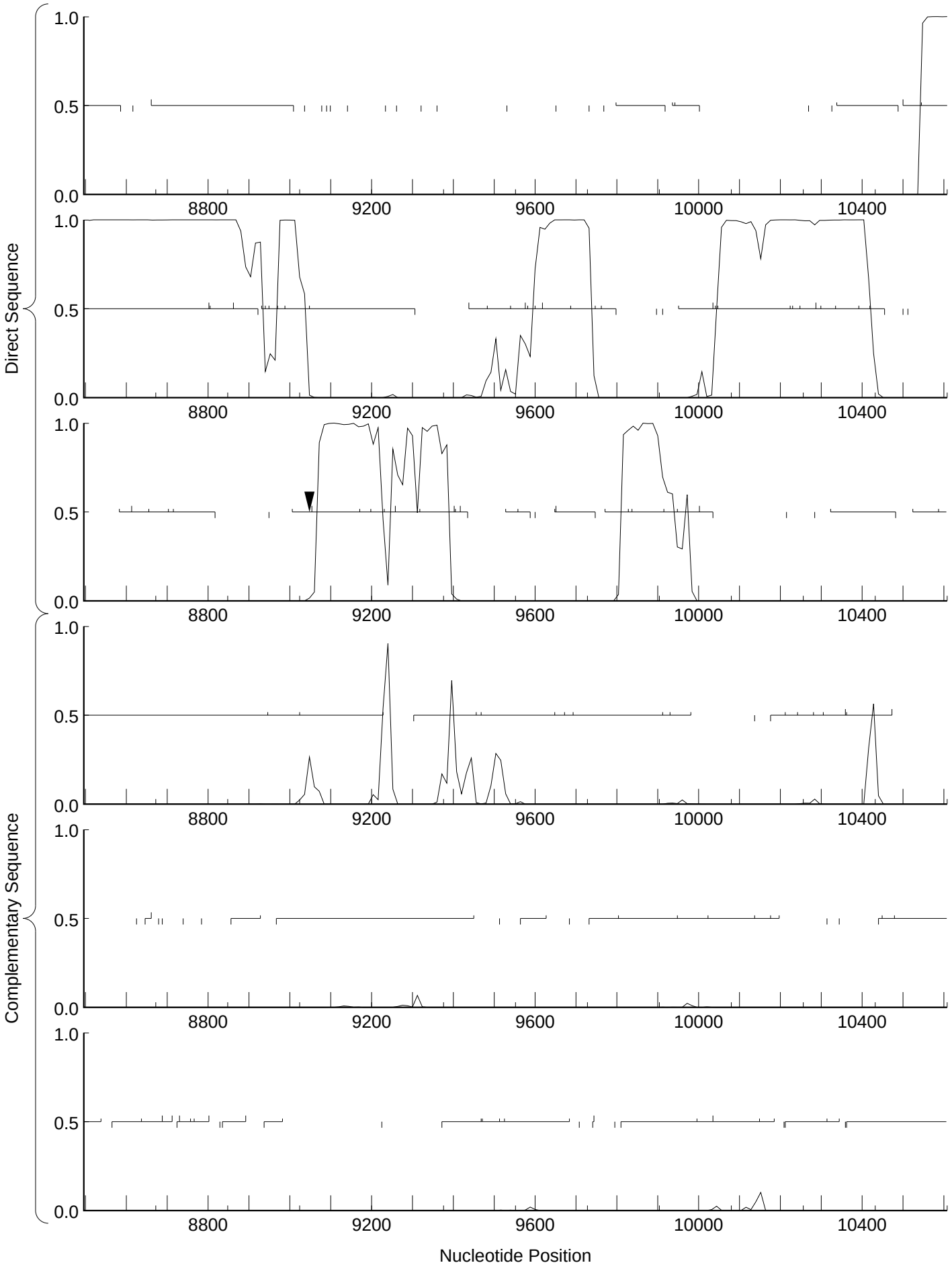
ator complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 4/31



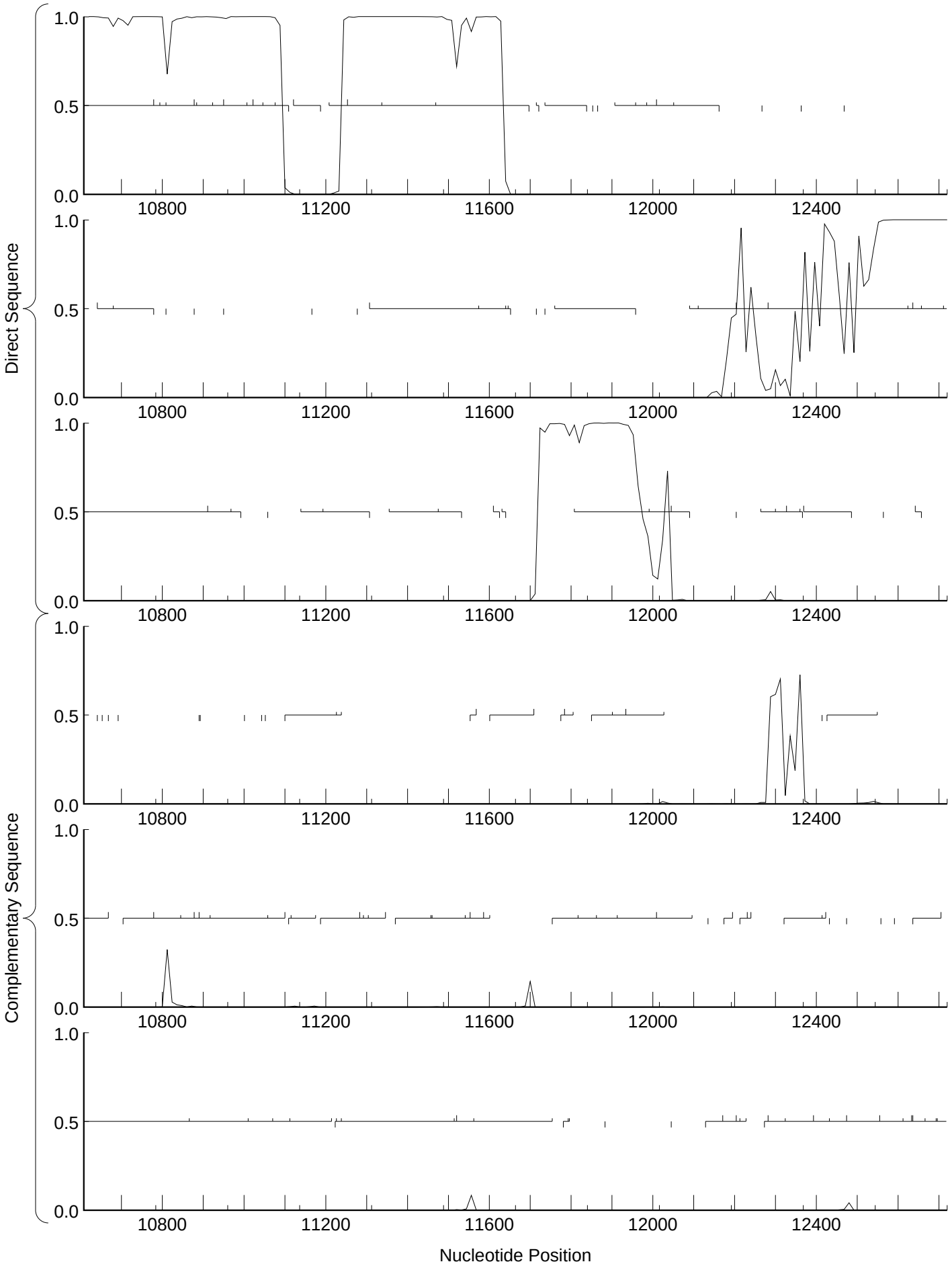
ator complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 5/31



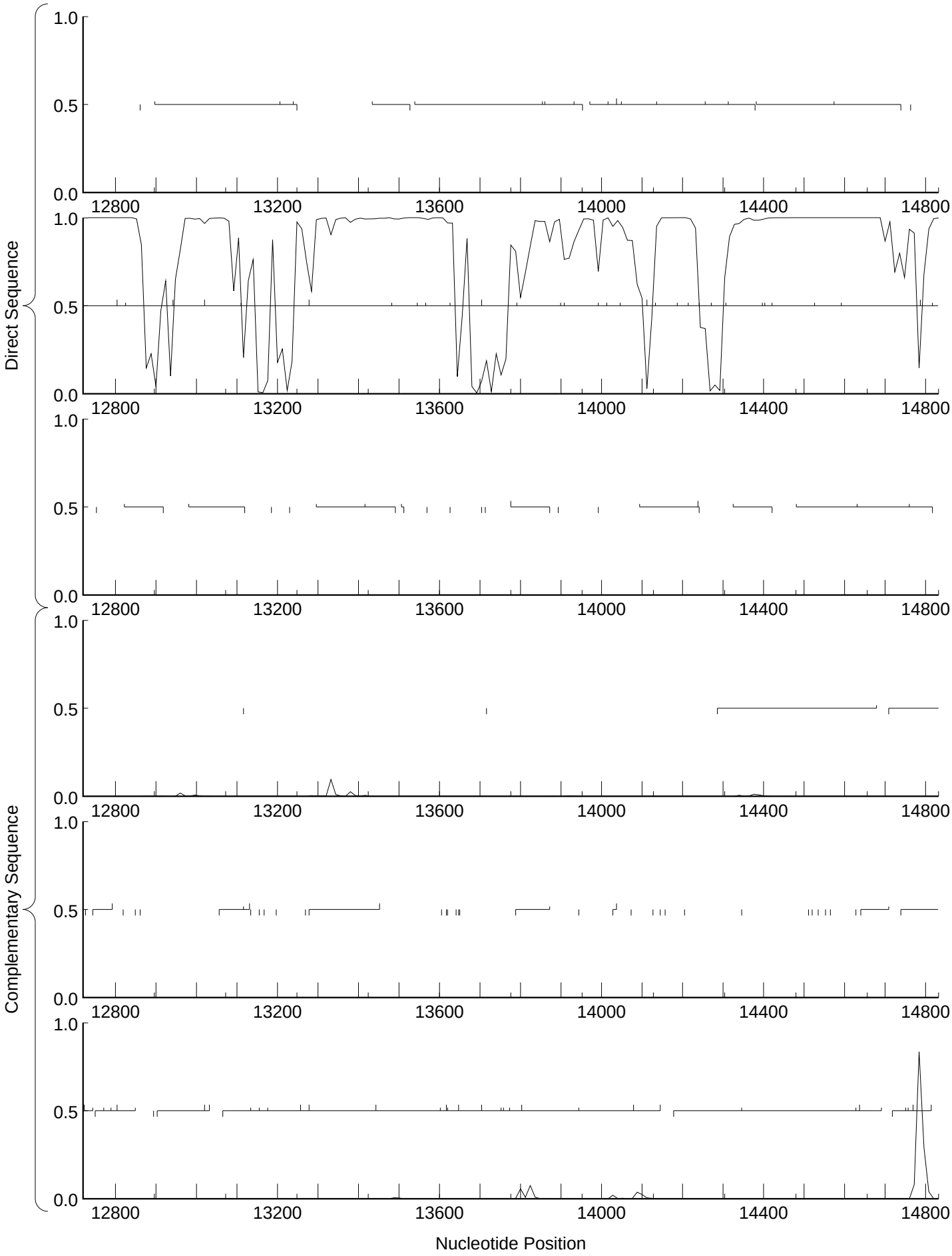
ator complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 6/31



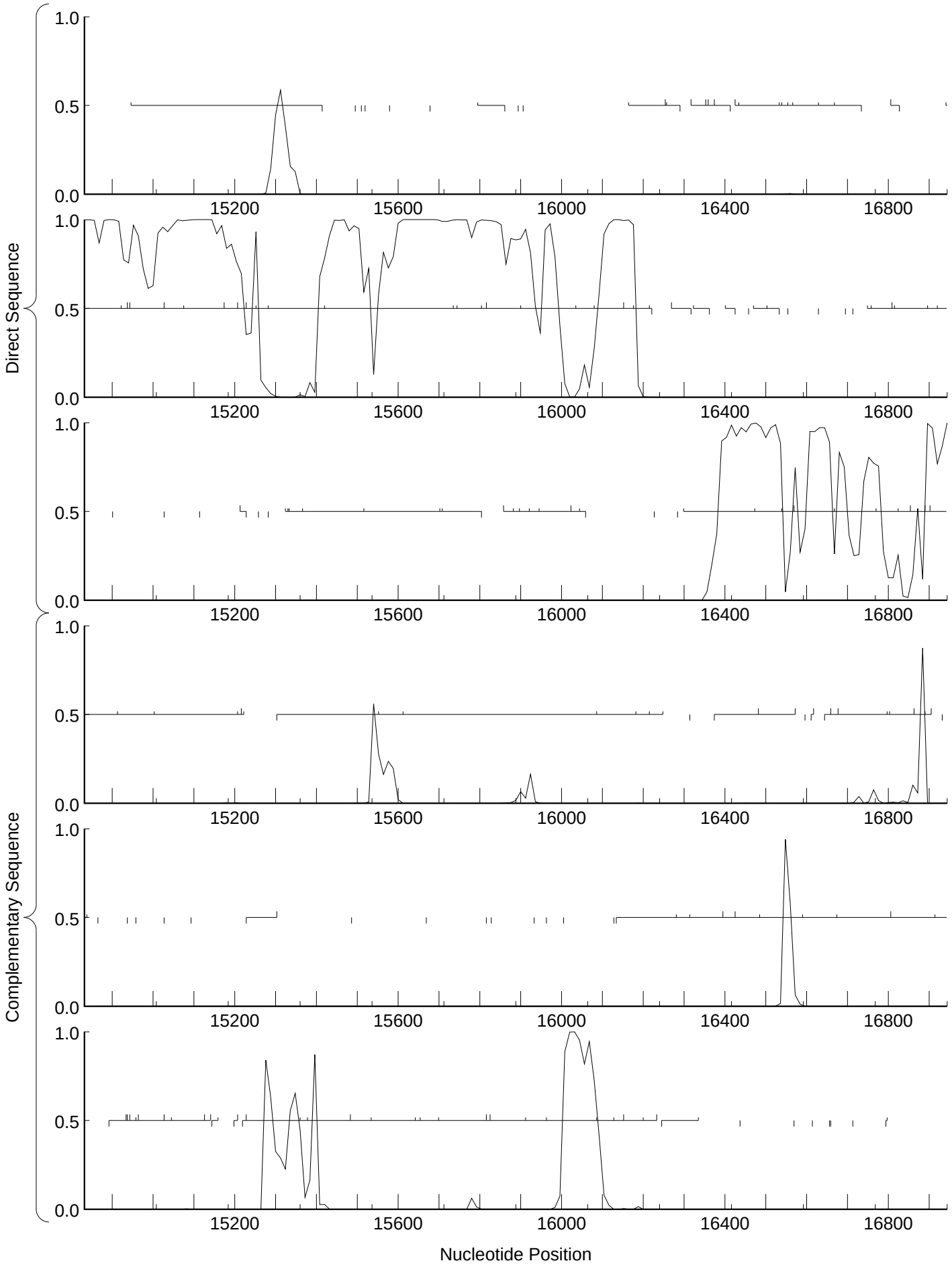
ator complete sequence, 63344 including 11-base 3' overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 7/31



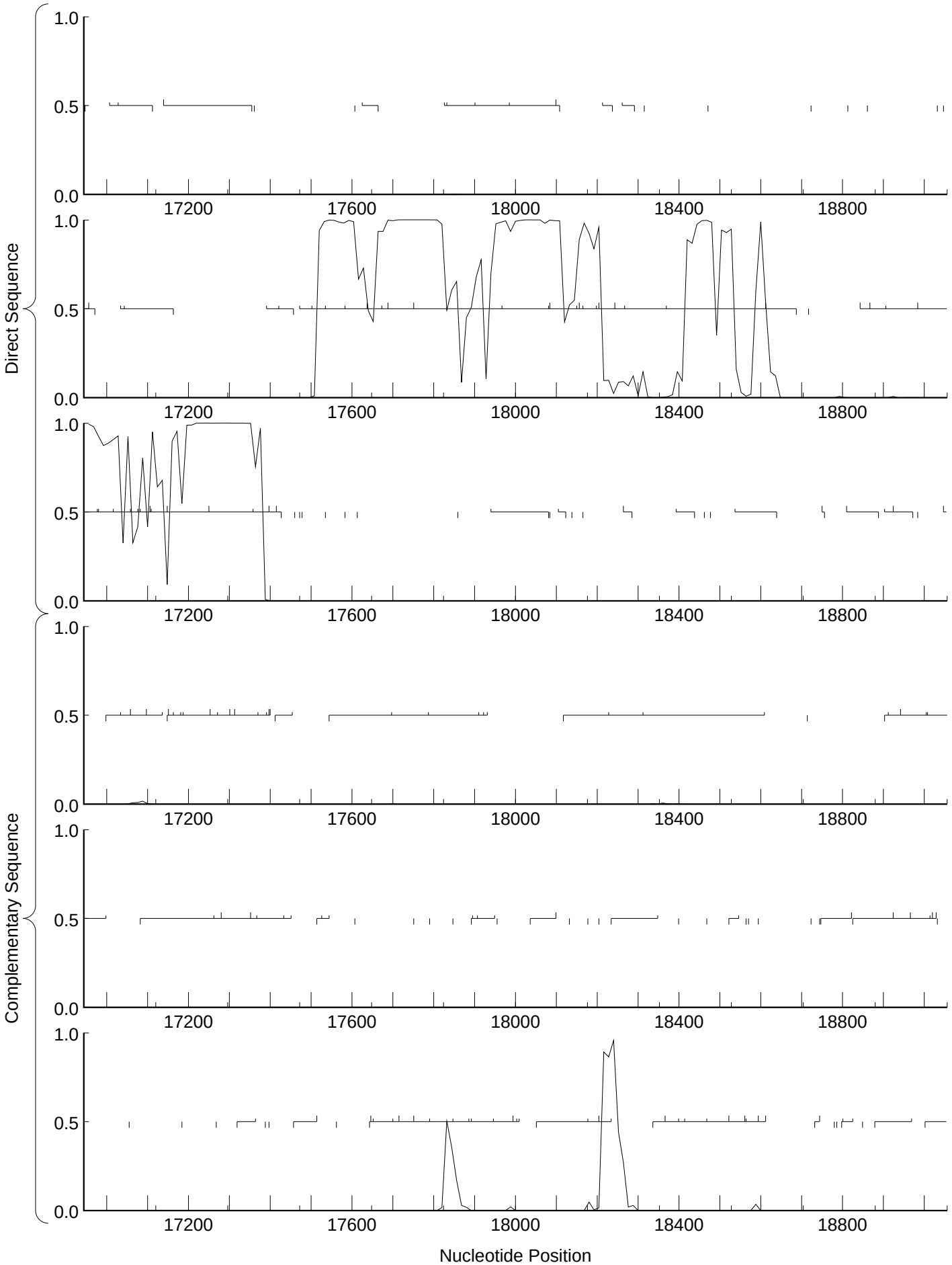
ator complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 8/31



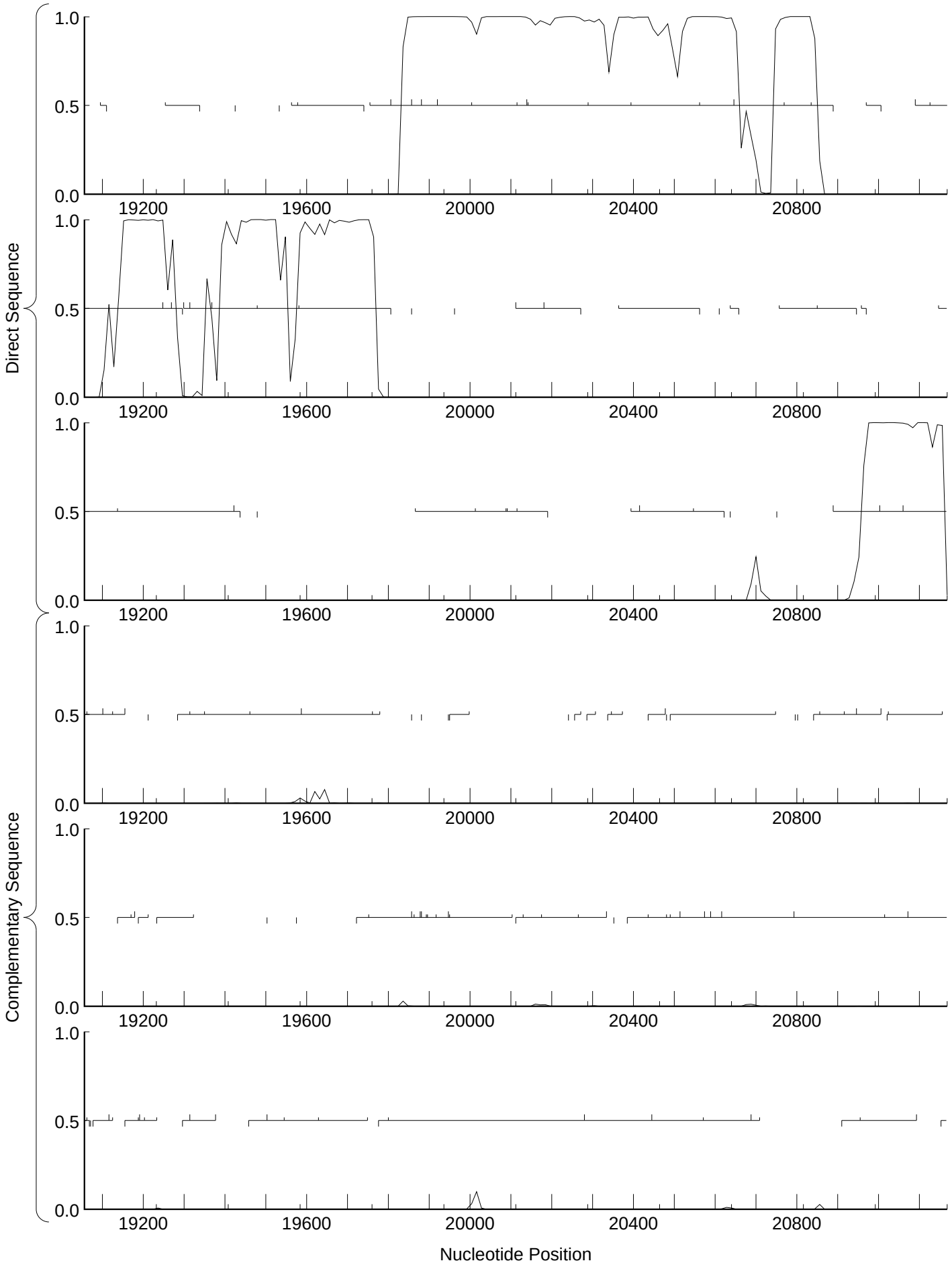
ator complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 9/31



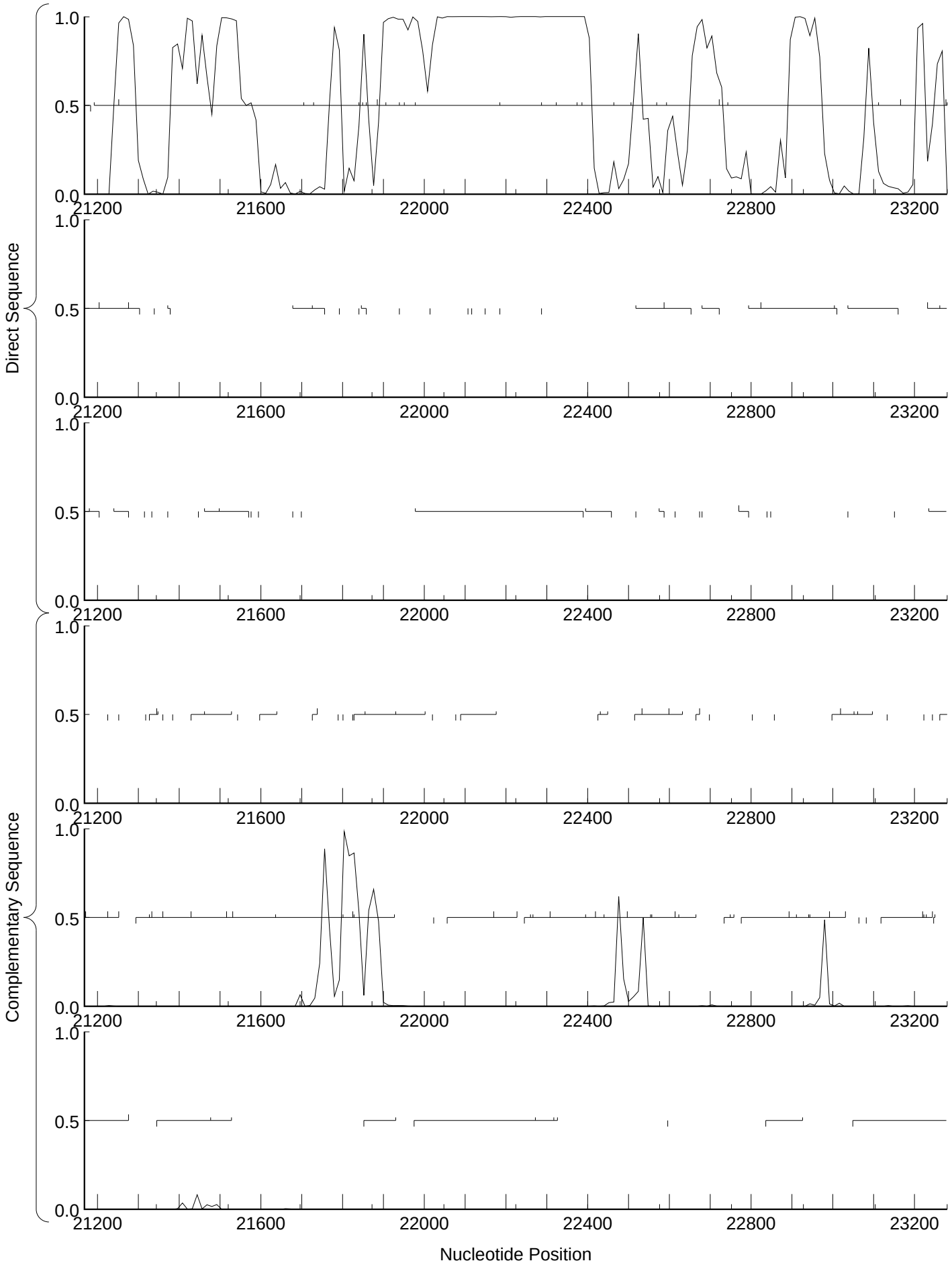
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 10/31



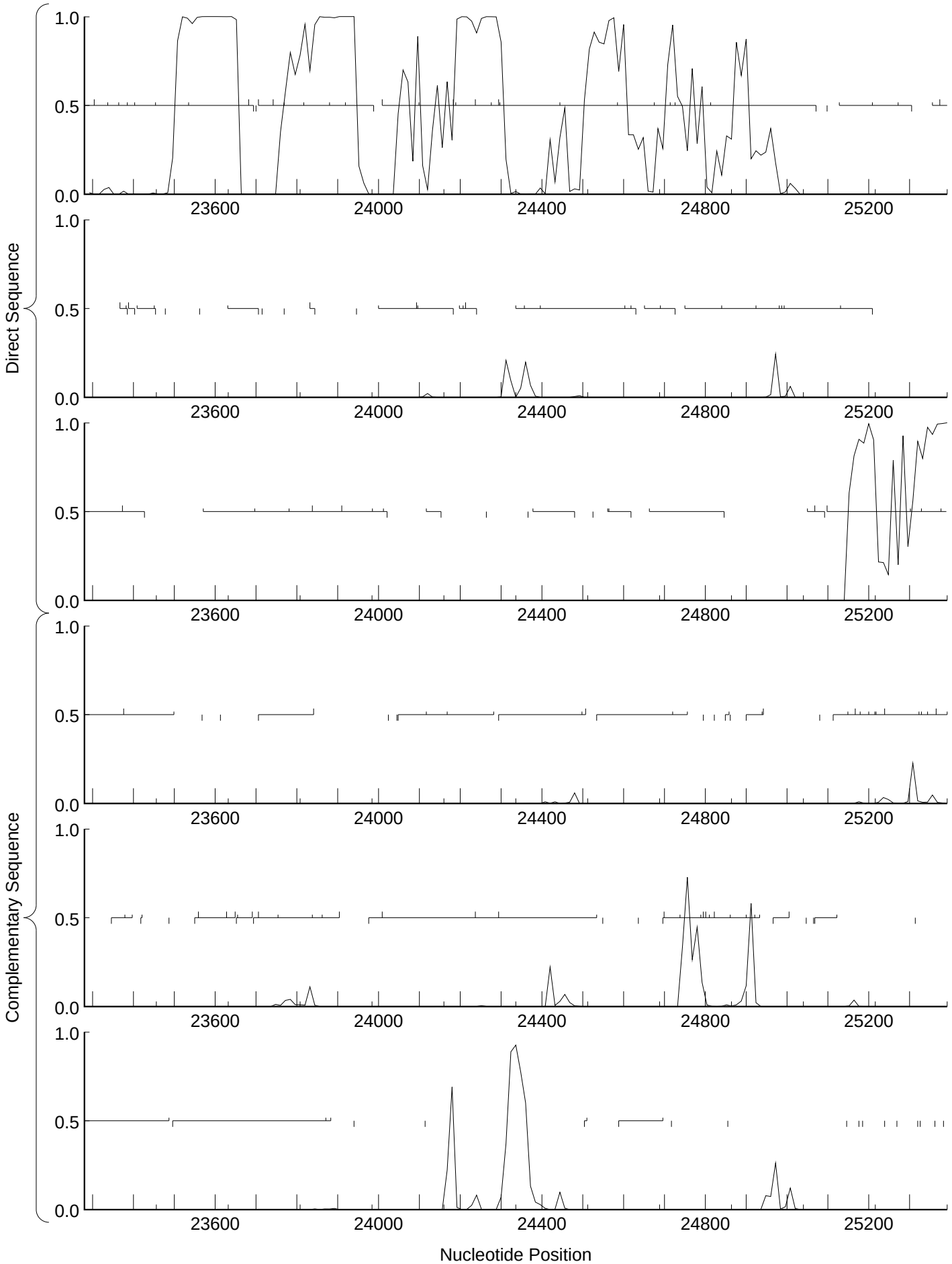
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 11/31



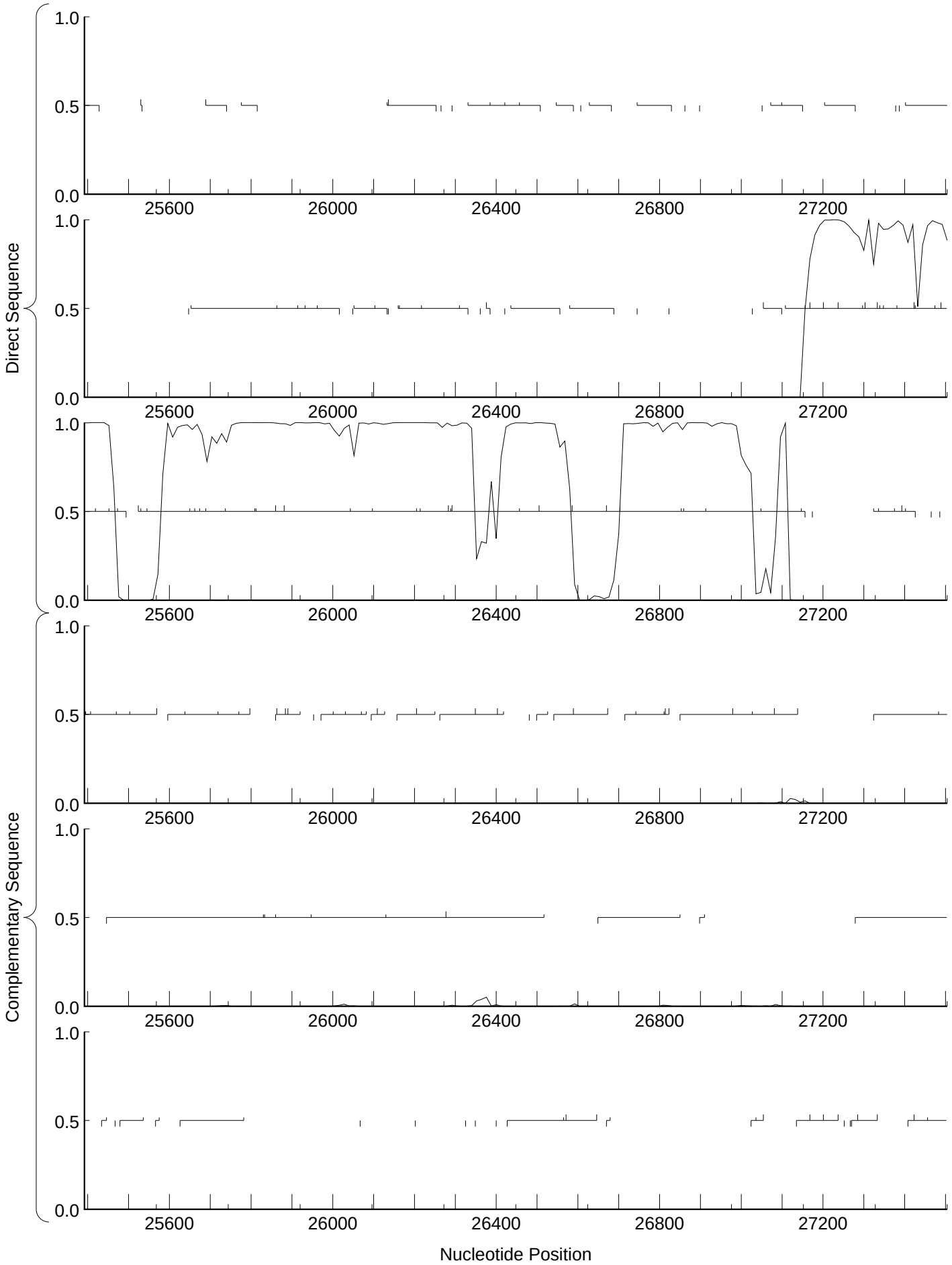
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 12/31



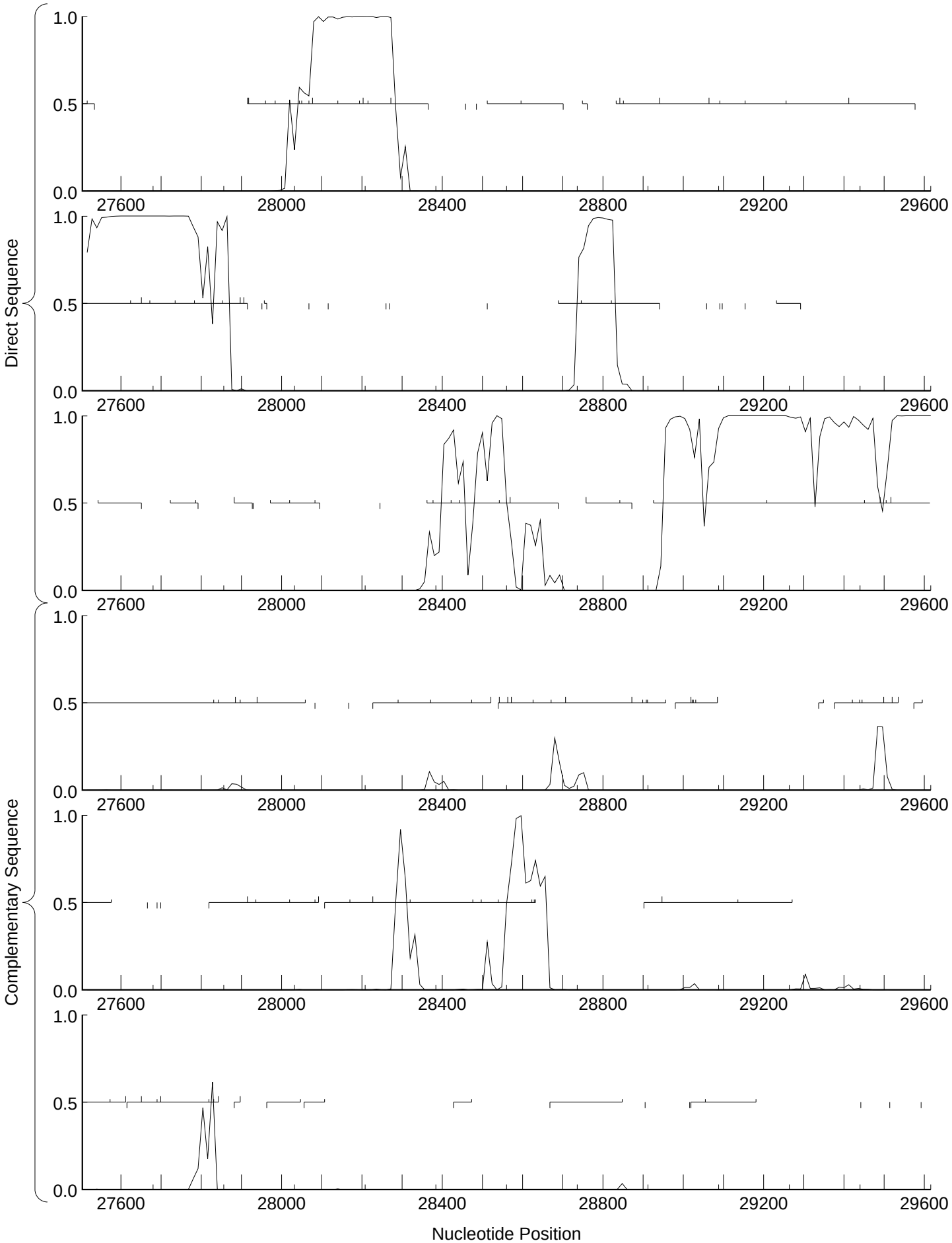
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 13/31



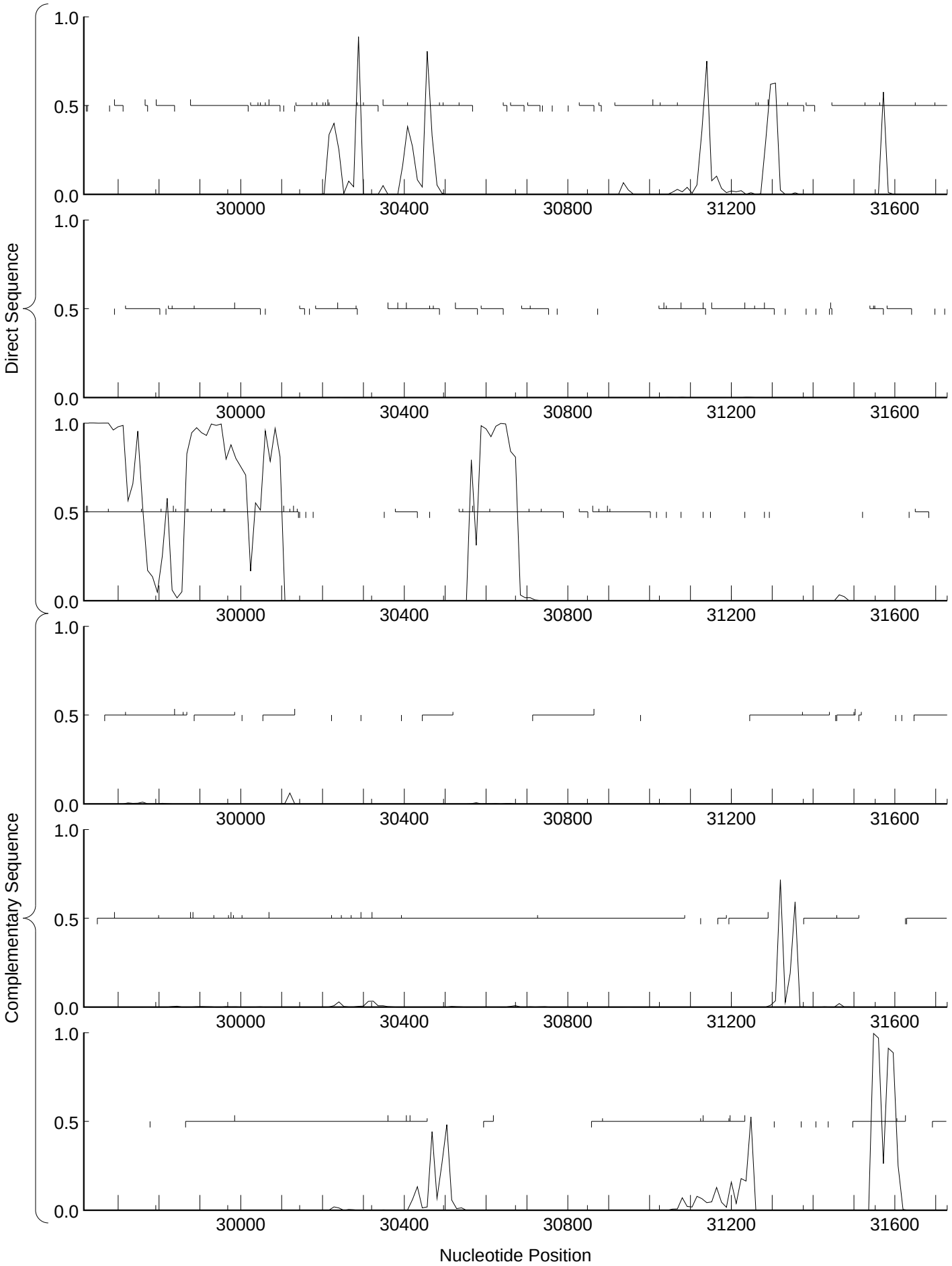
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 14/31



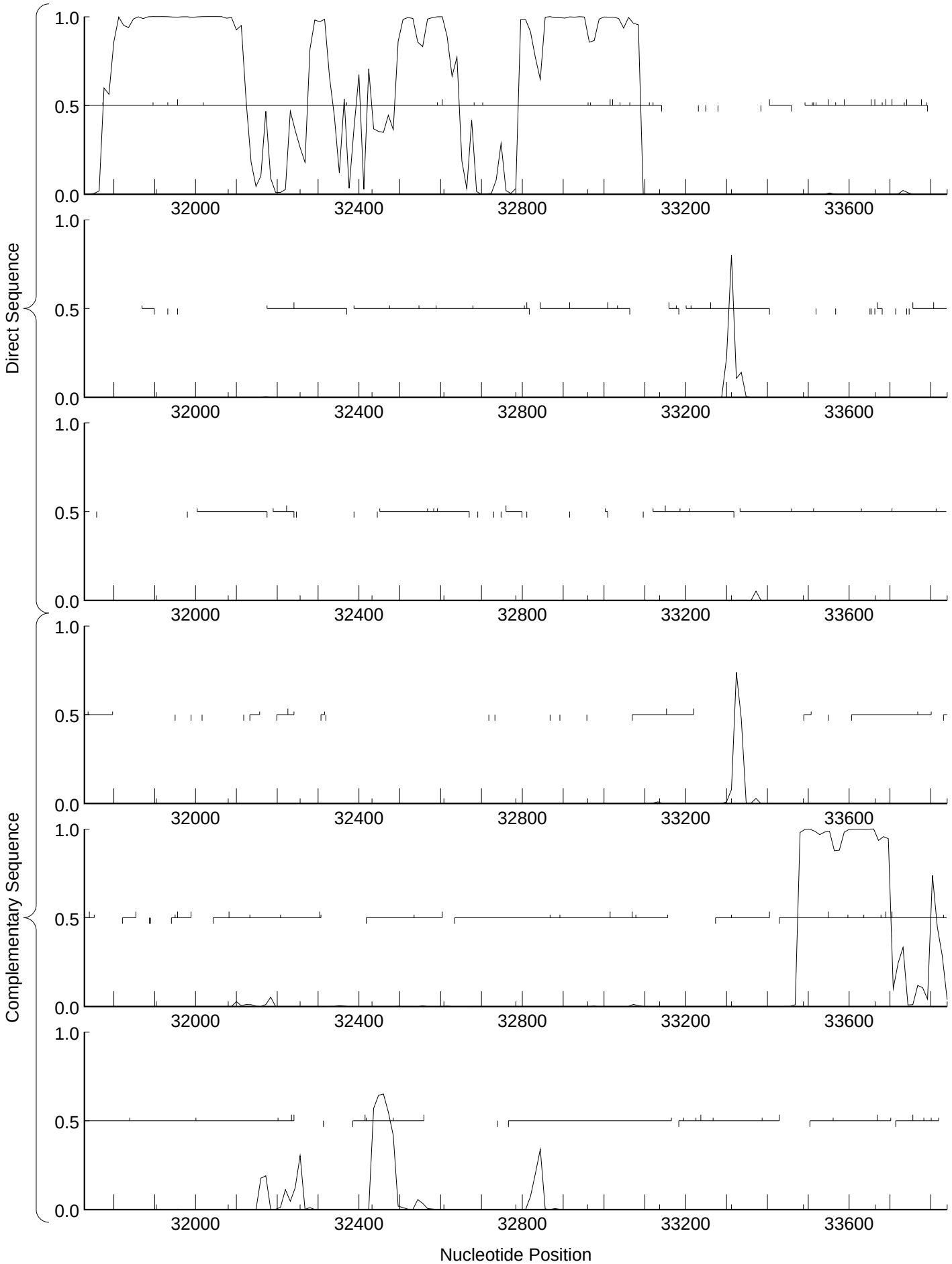
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 15/31



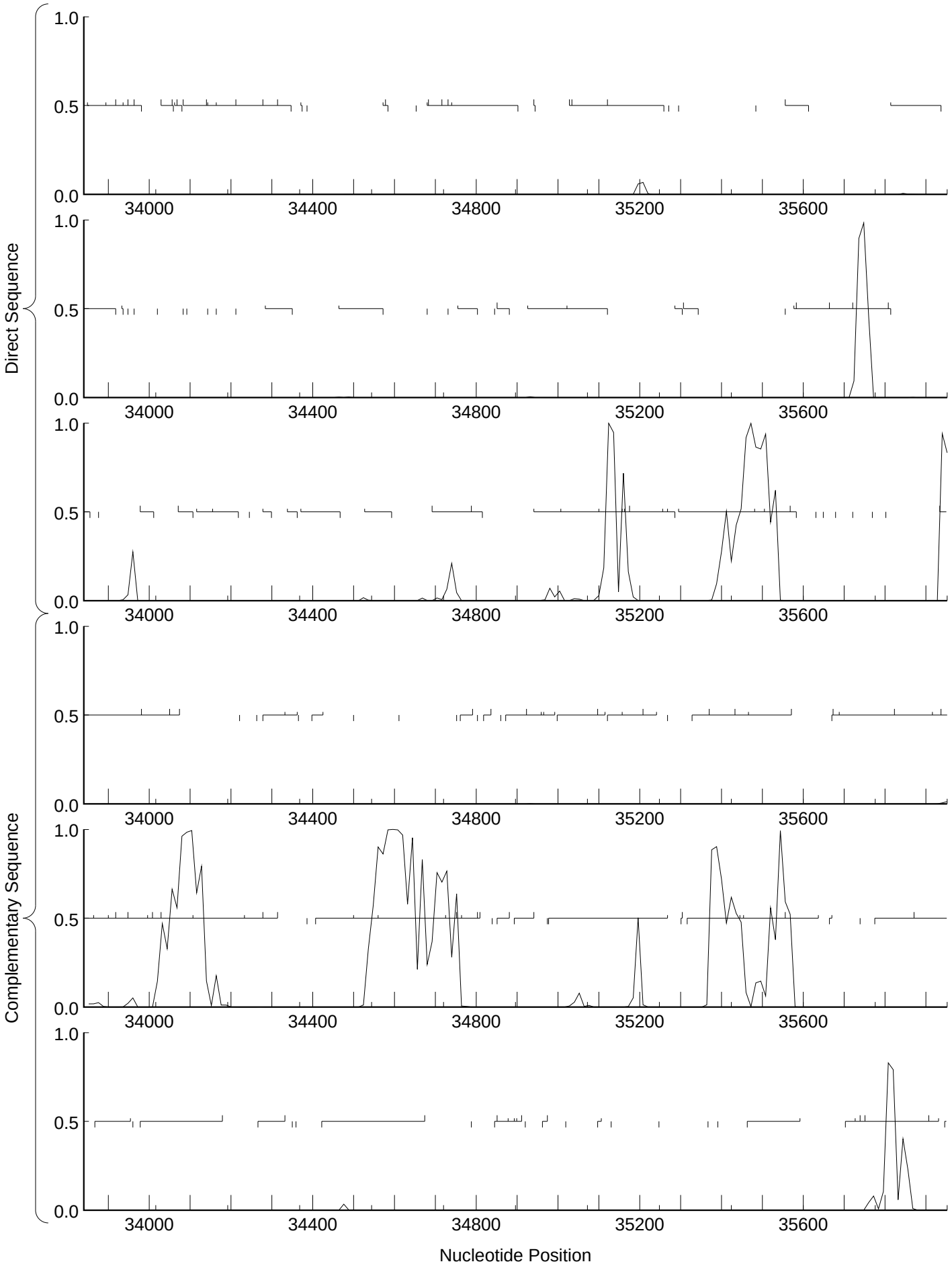
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 16/31



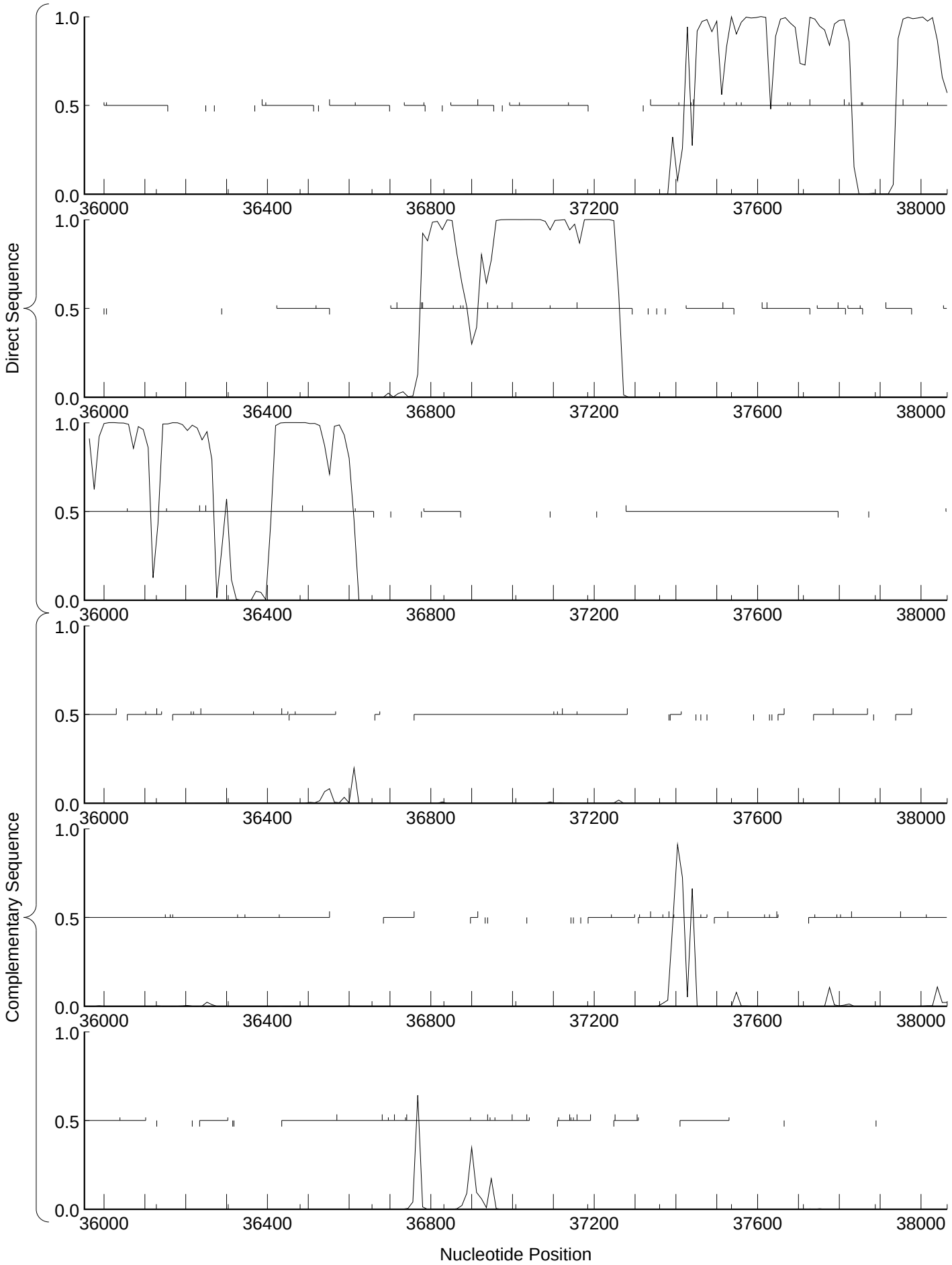
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 17/31



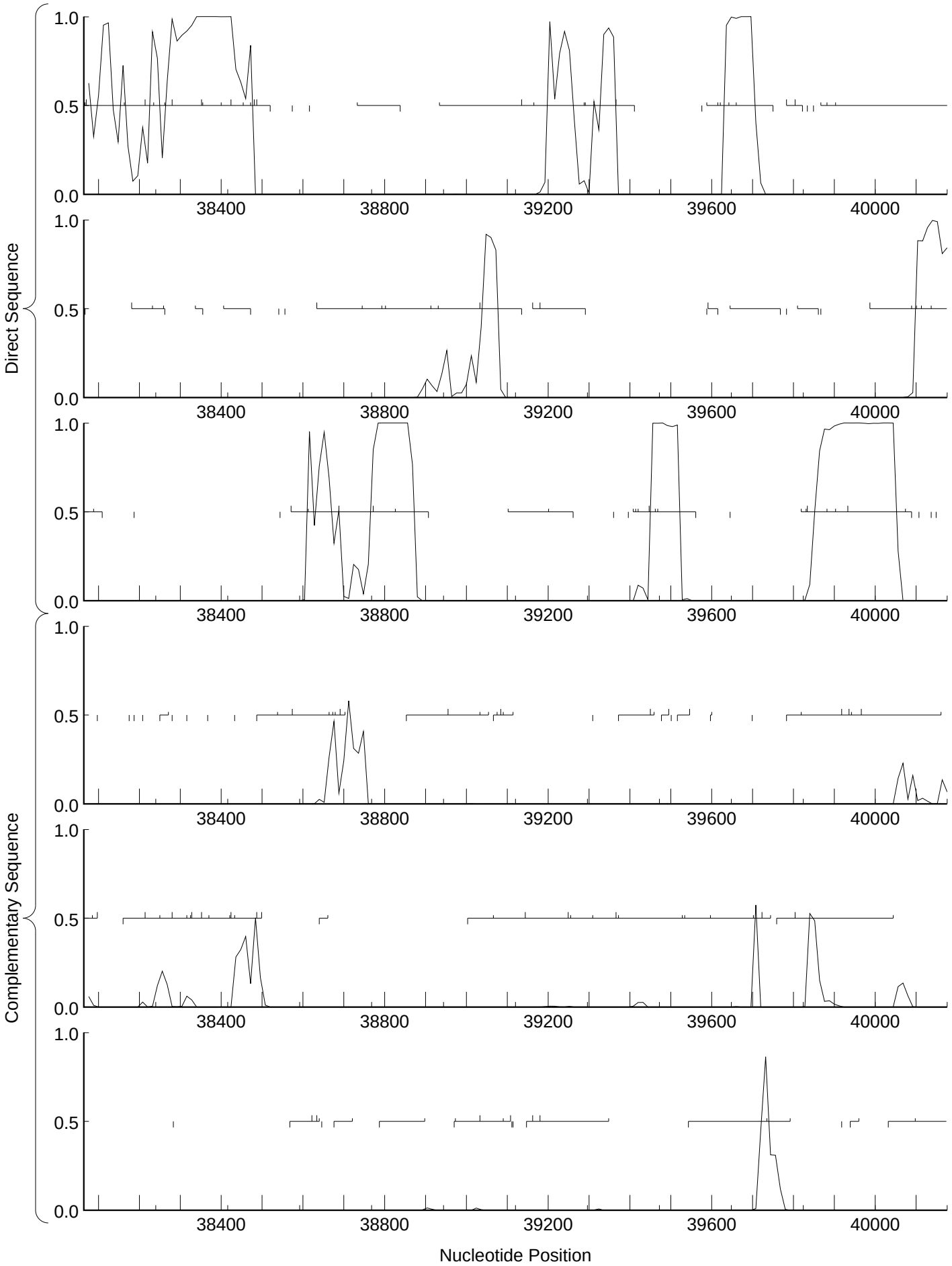
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 18/31



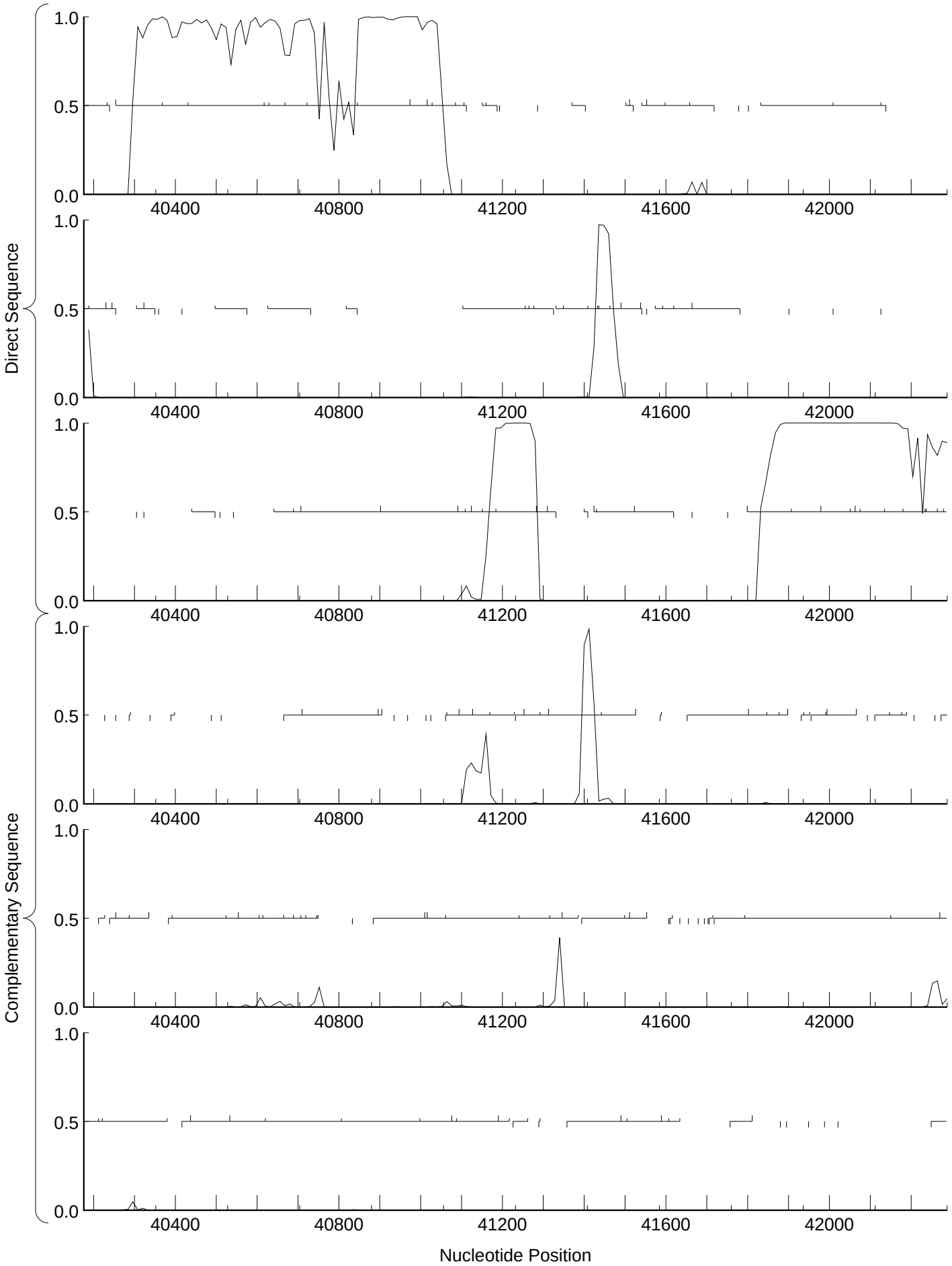
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 19/31



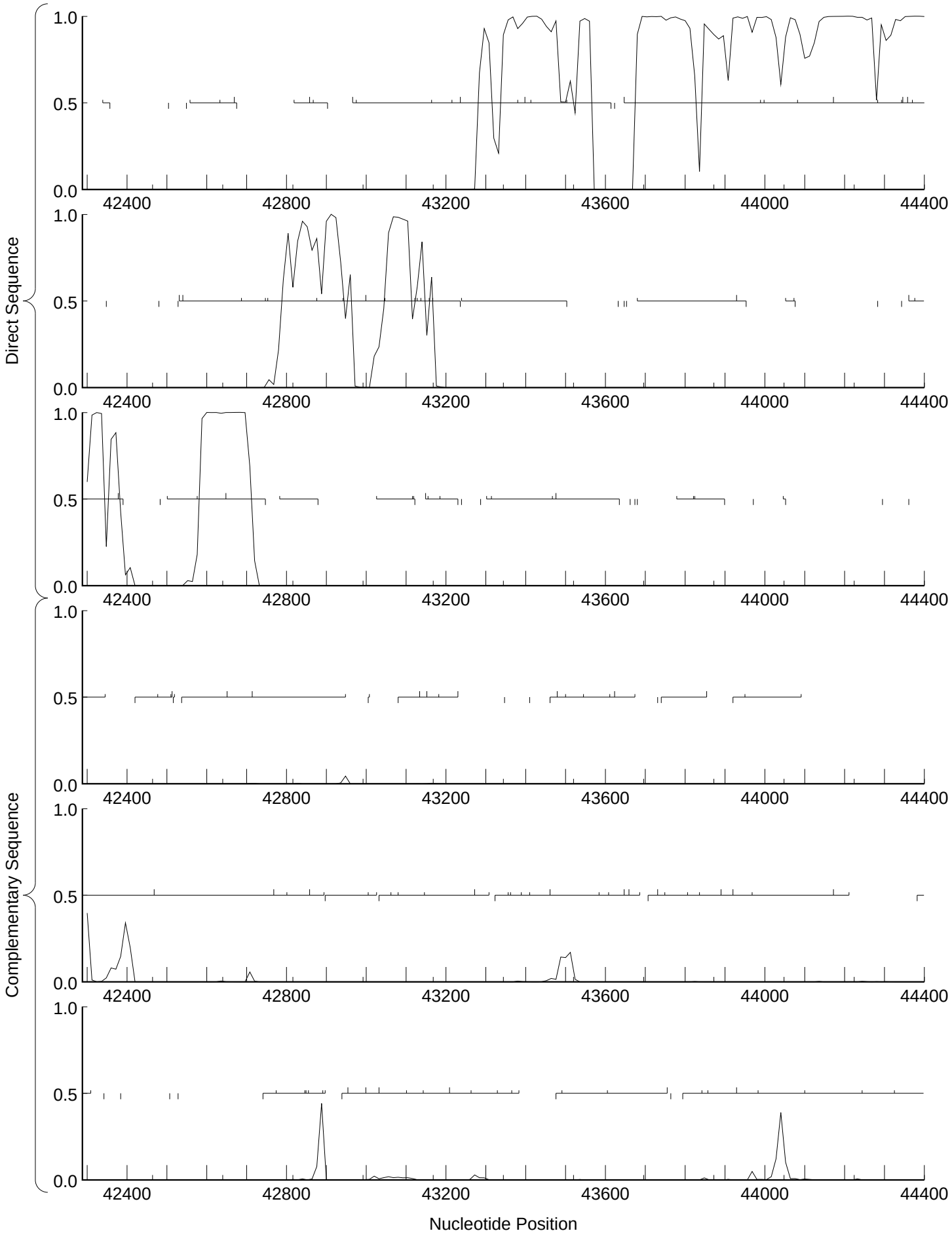
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 20/31



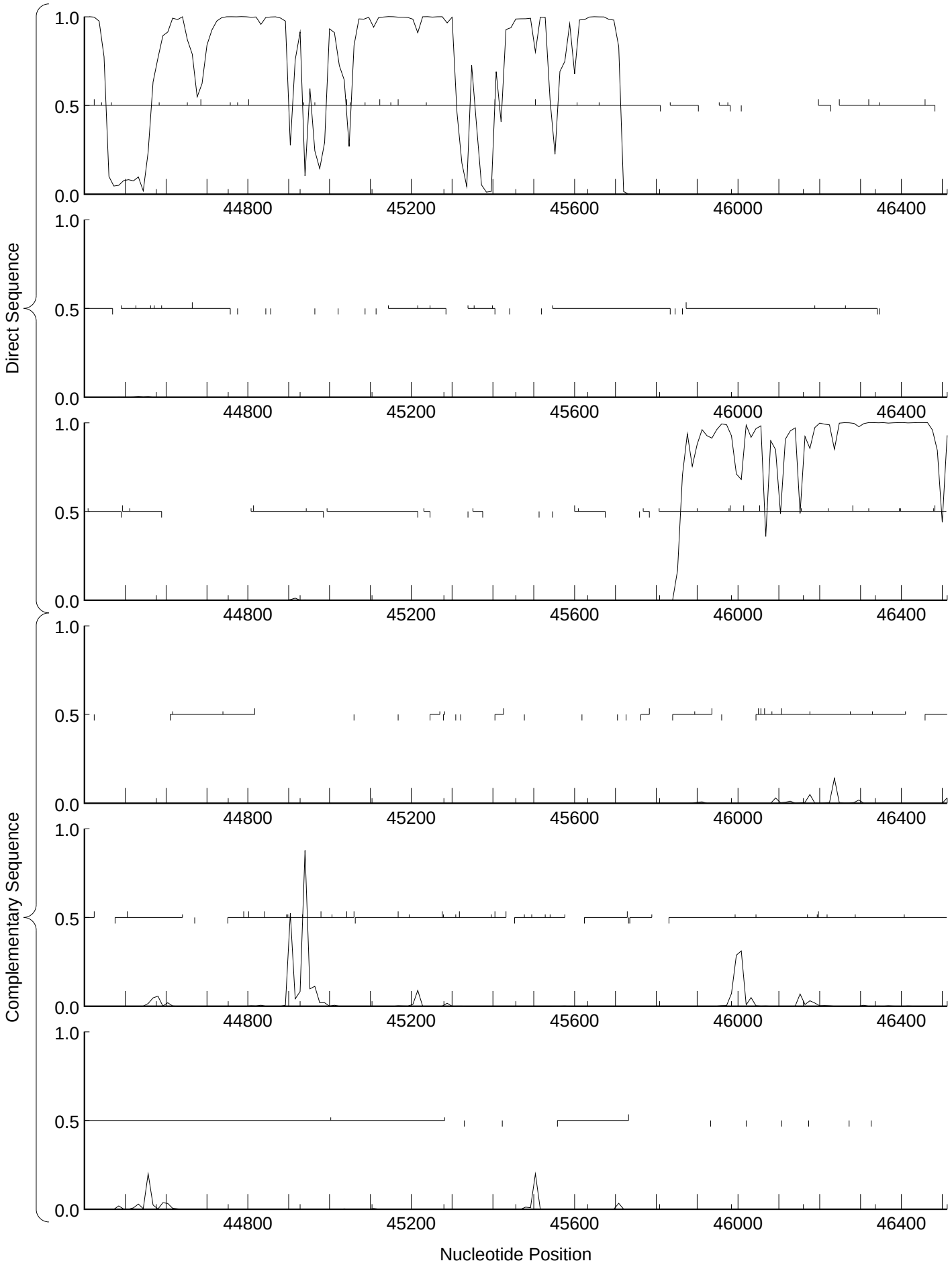
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 21/31



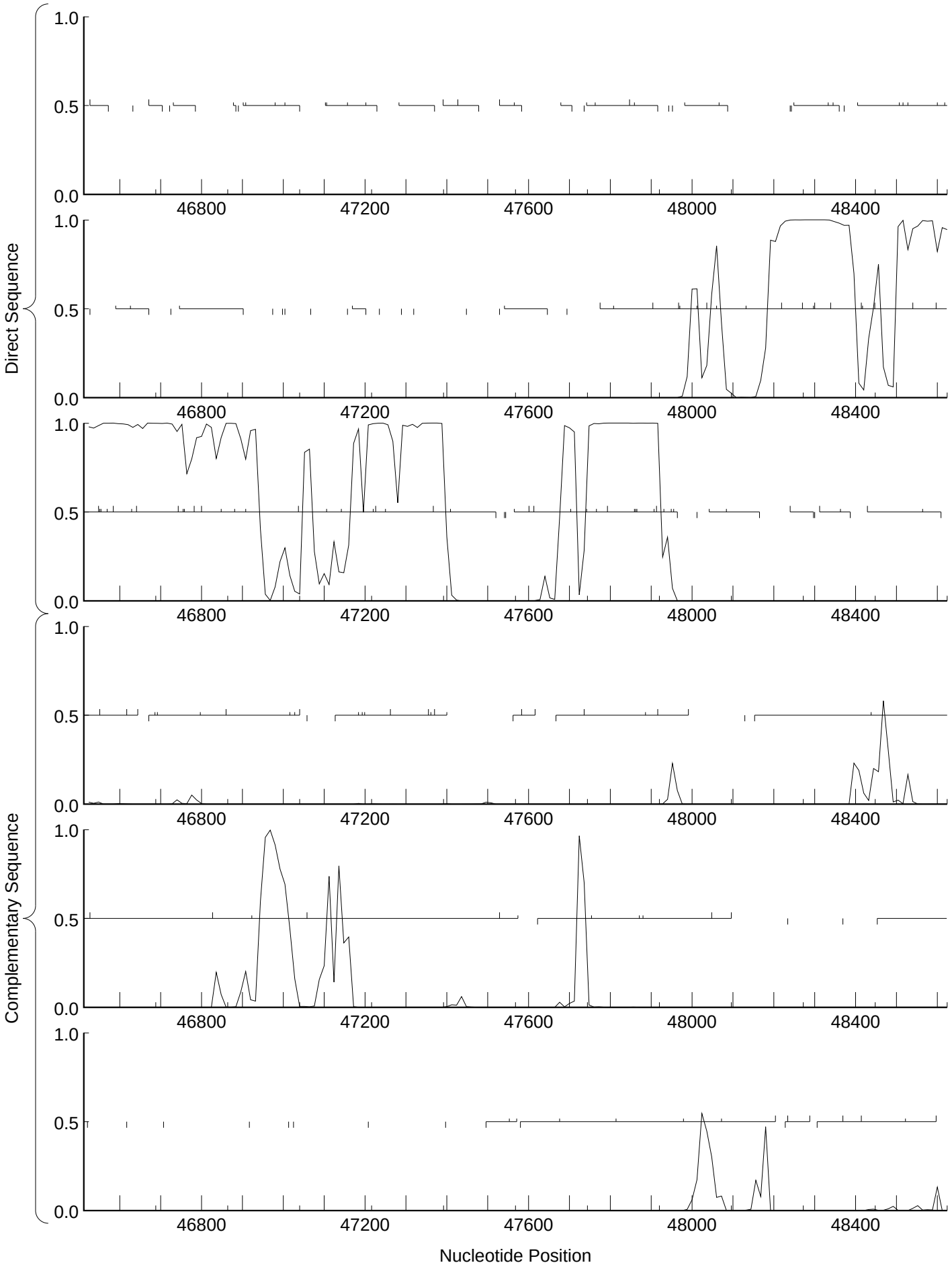
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 22/31



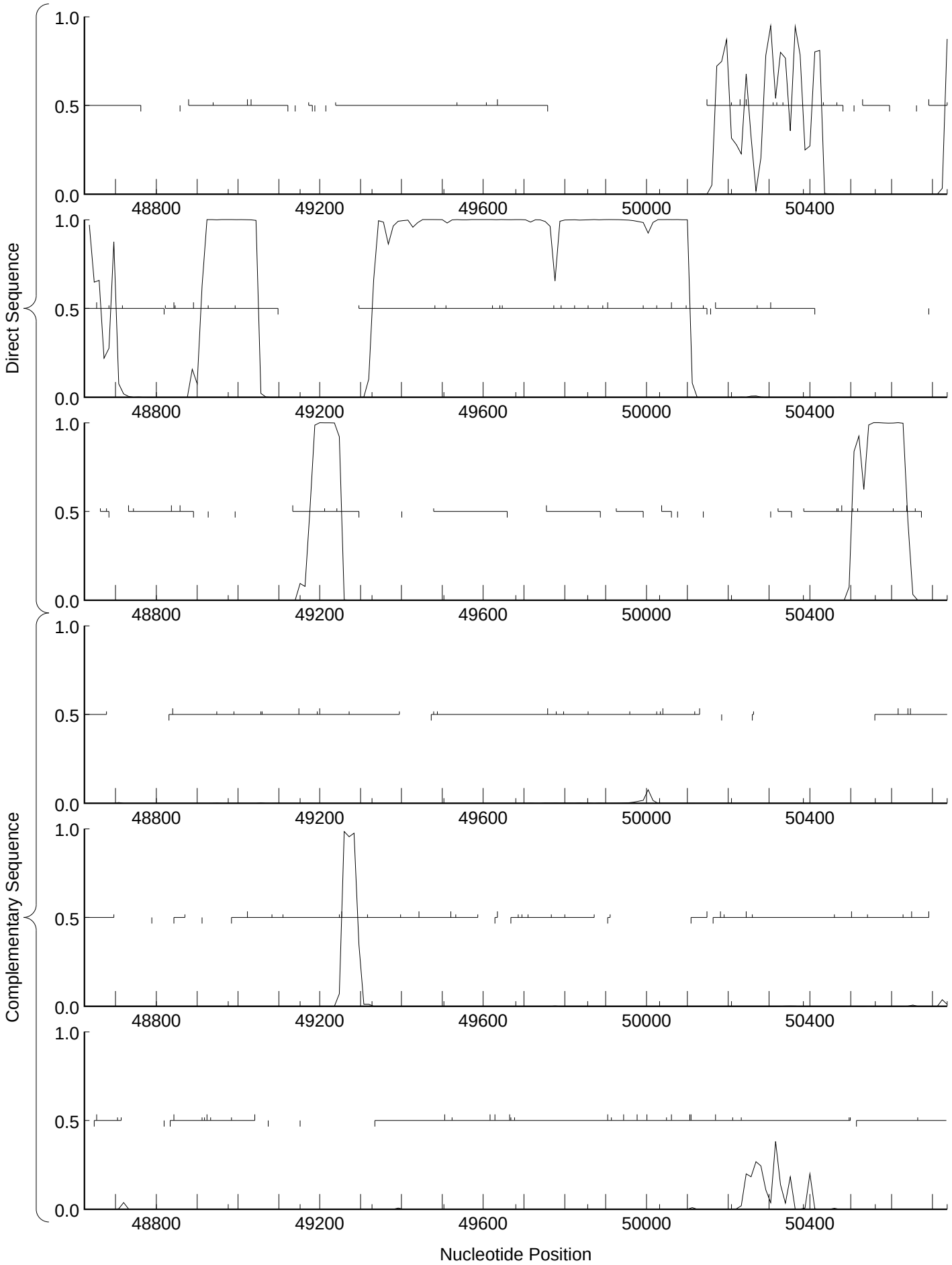
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 23/31



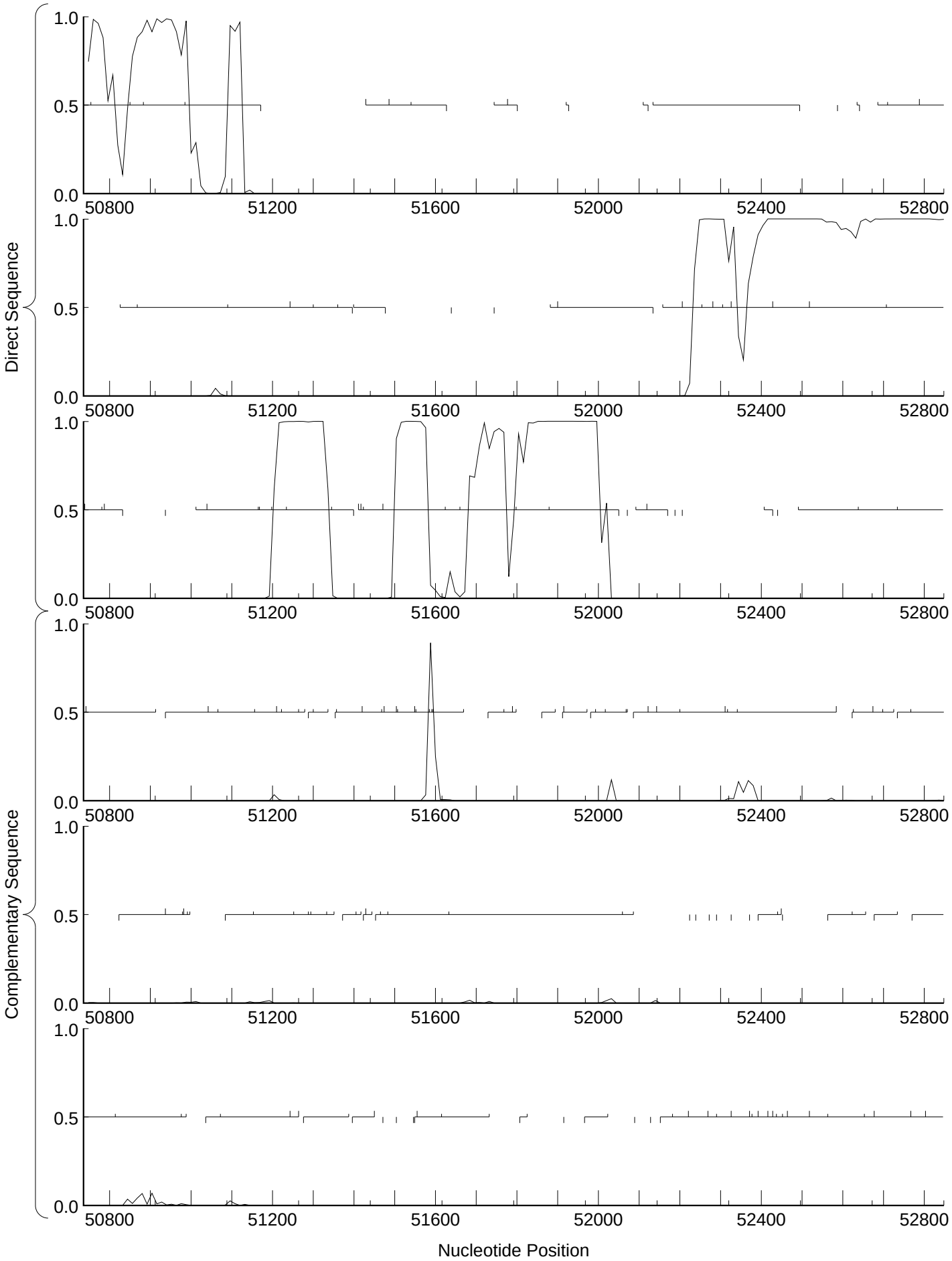
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 24/31



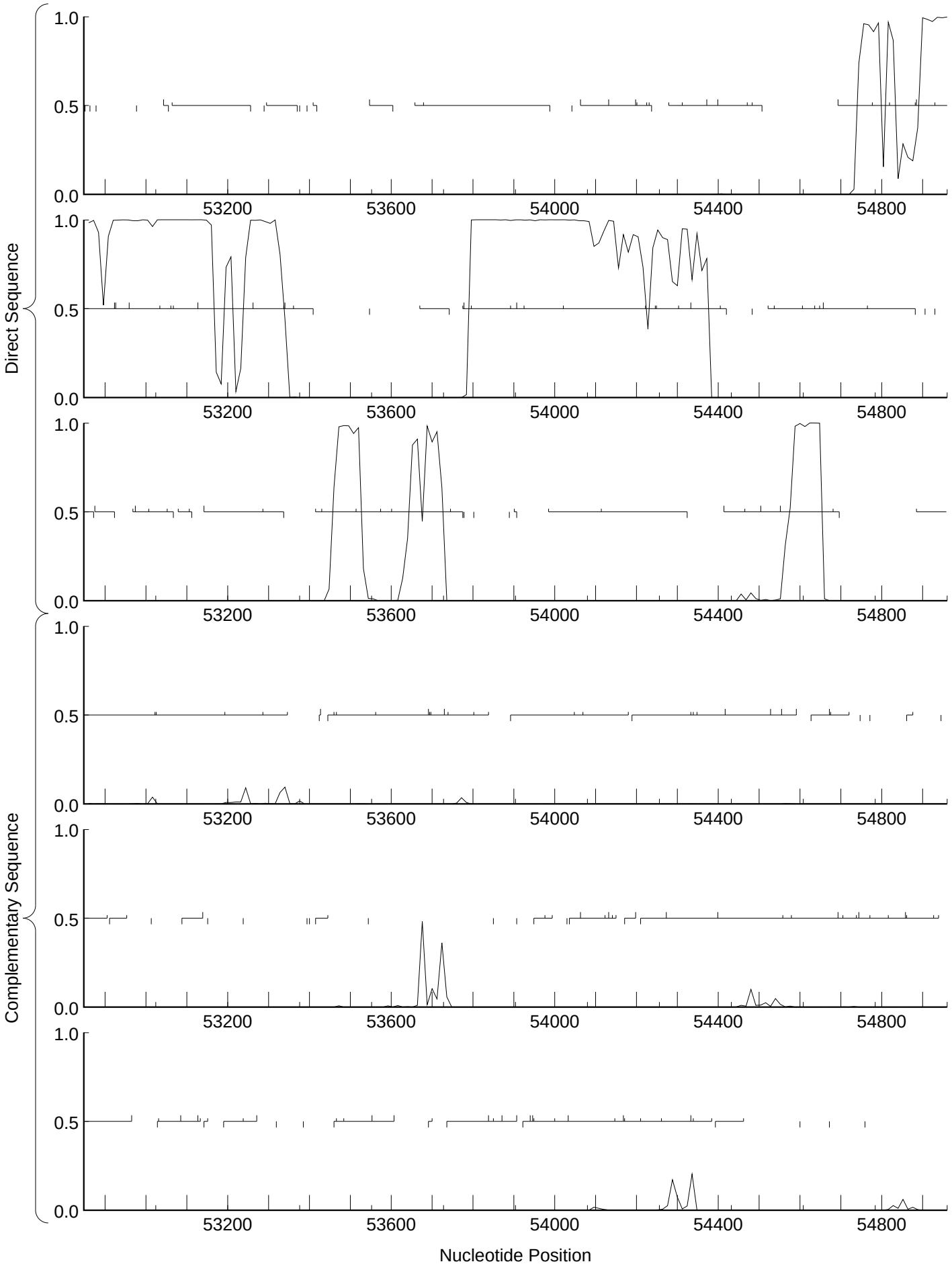
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 25/31



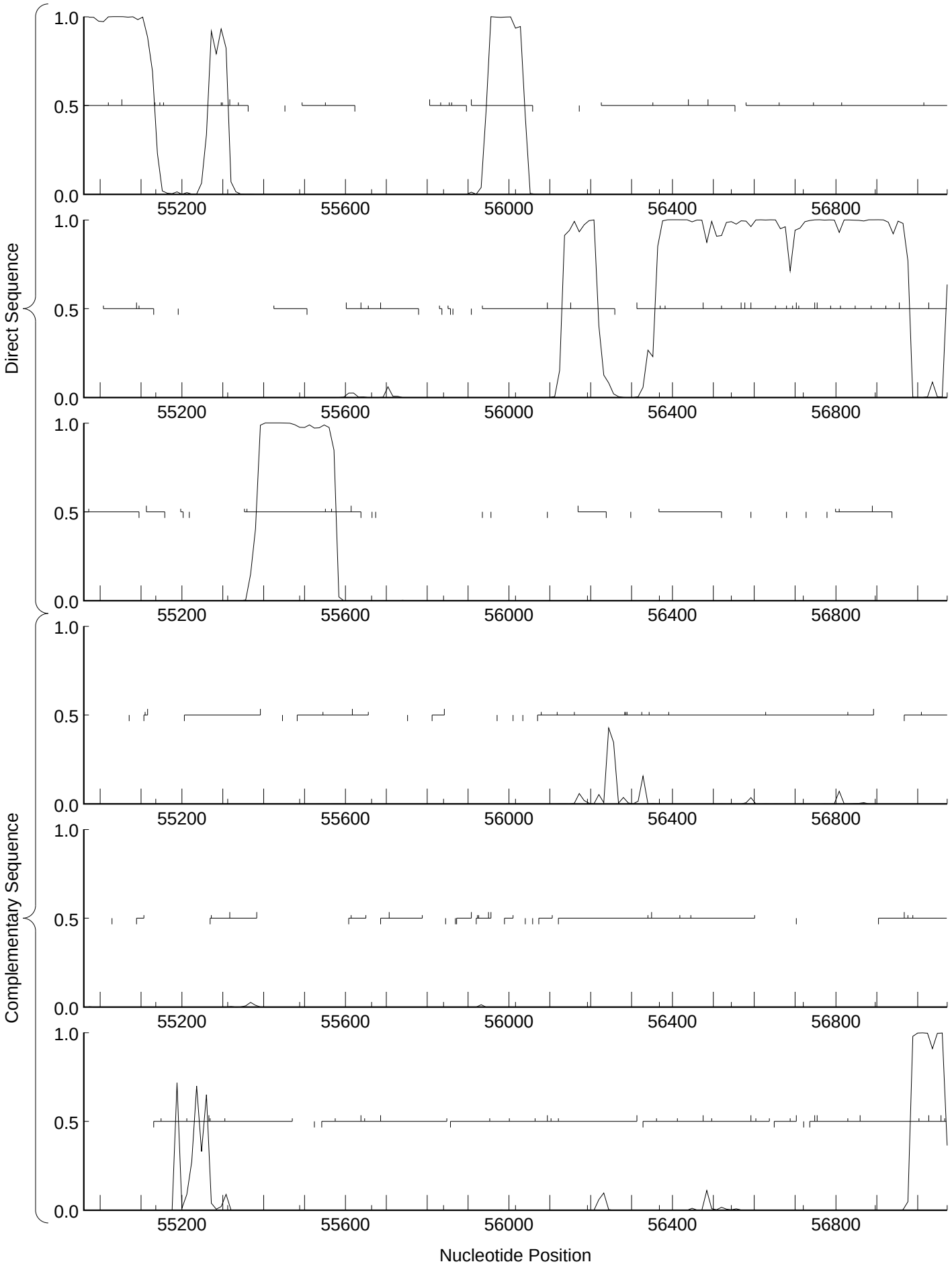
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 26/31



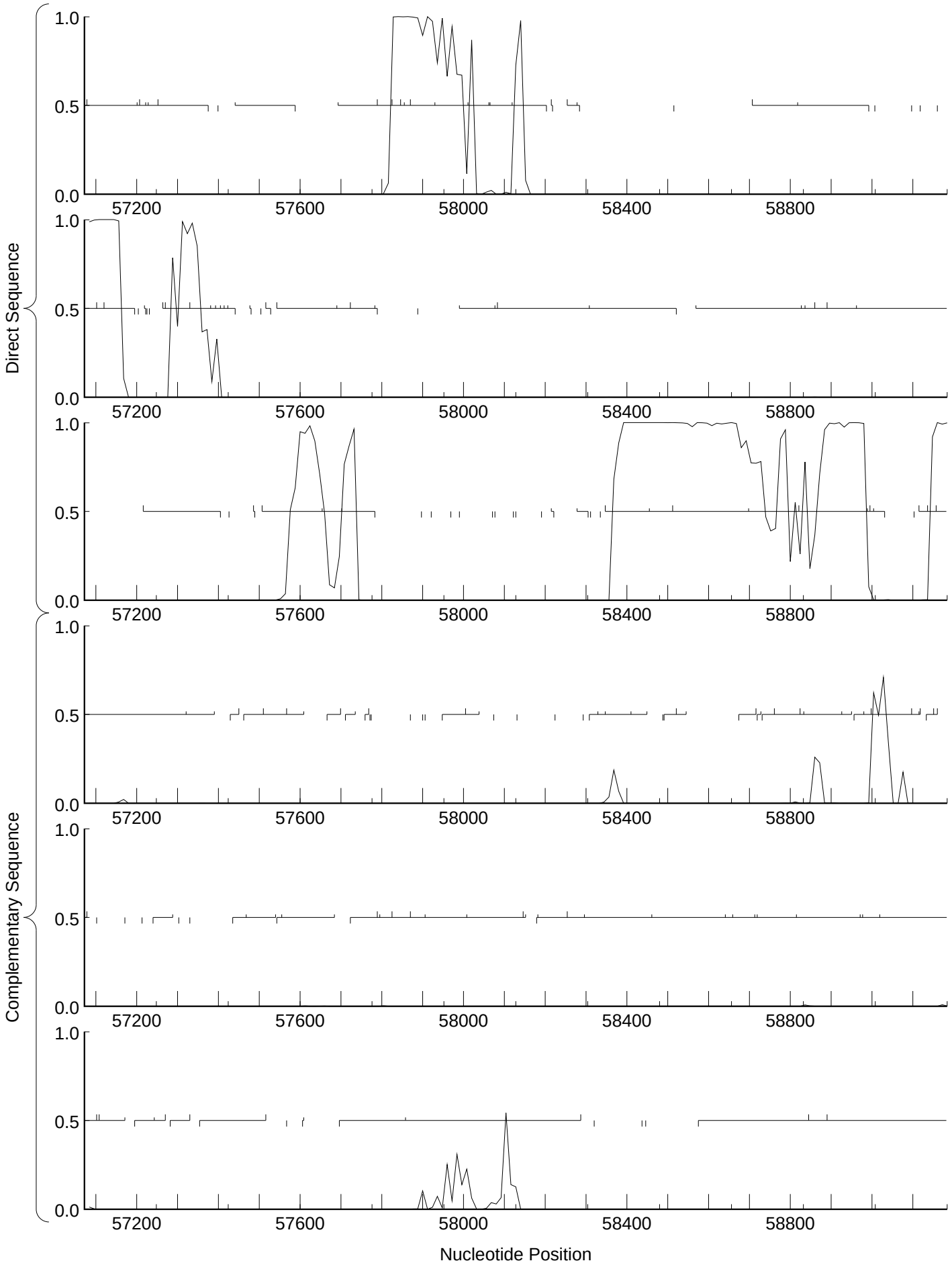
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 27/31



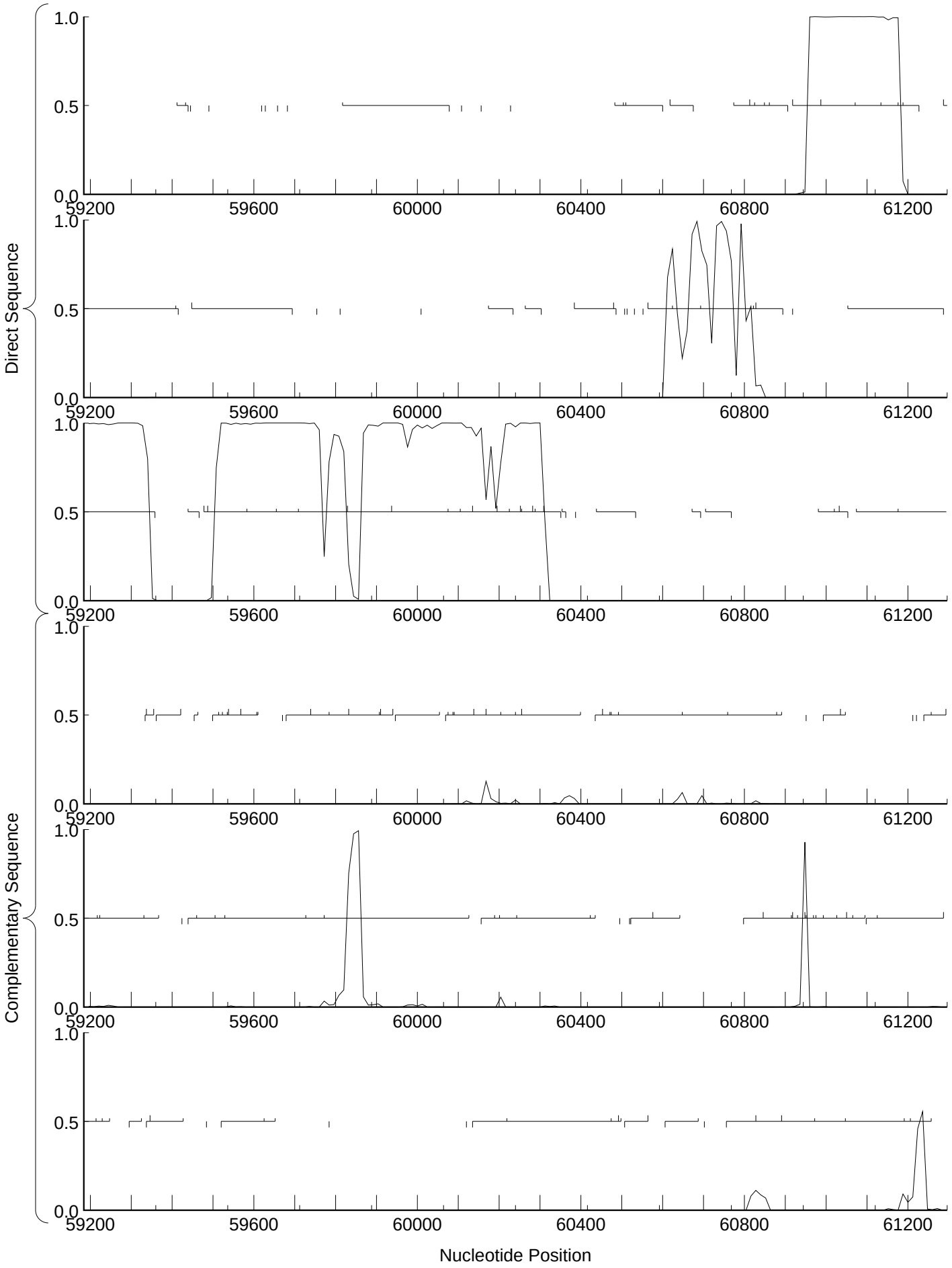
or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 28/31



or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 29/31



or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 30/31



or complete sequence, 63344 including 11-base 3’ overhang (CTCAGTGGCAT), Cluster K5, Order 4, Window 96, Step 12, 31/31

