GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Tortellini complete sequence, 49658 bp including 12-base 3’ overhang (C
Analysis Date    :  3/28/24 at 6:27:11
Pages            :  25
Sequence Length  :  49658 bp
GC Content       :   65.82%

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

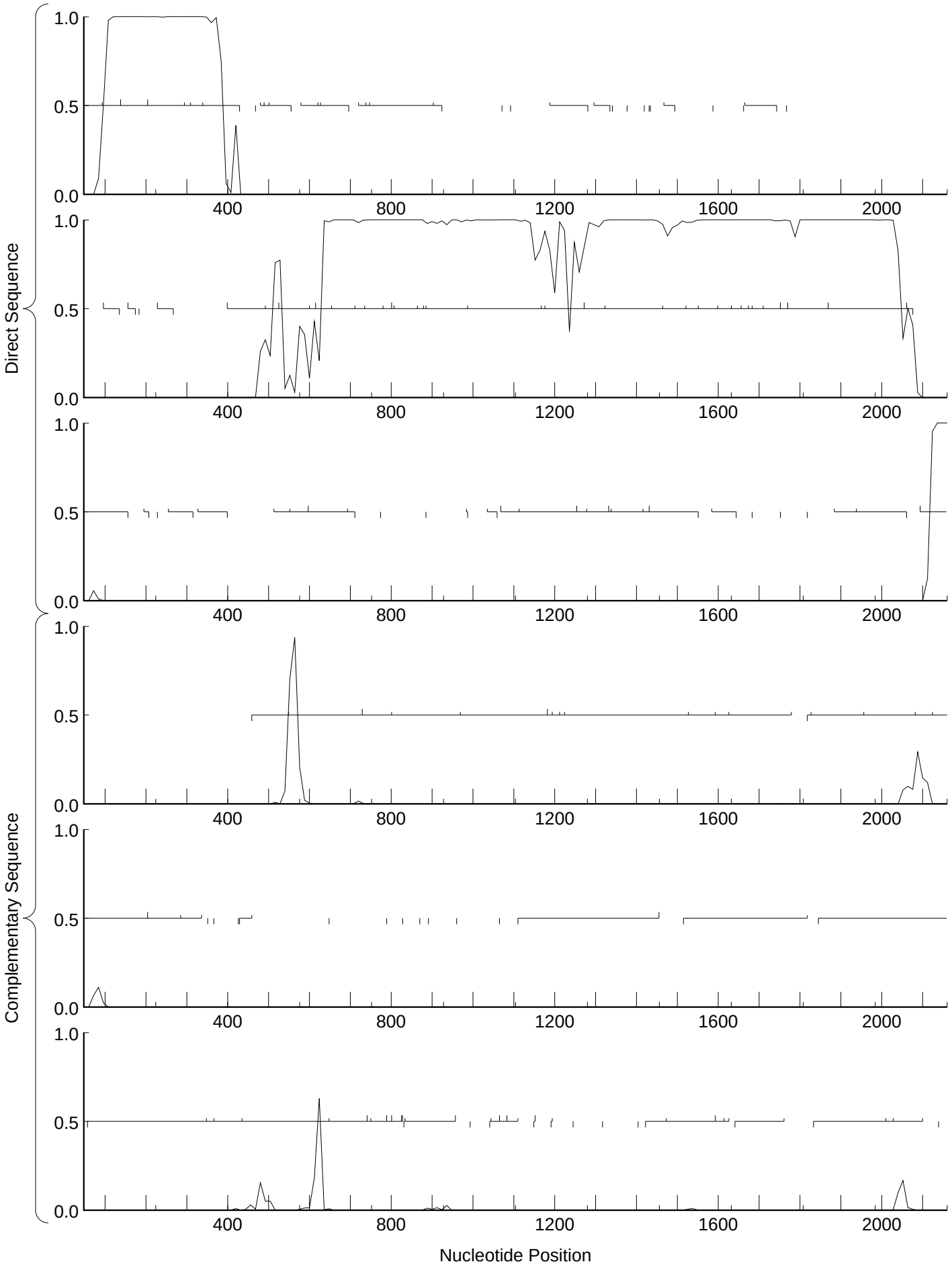

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

Matrix notes & comments

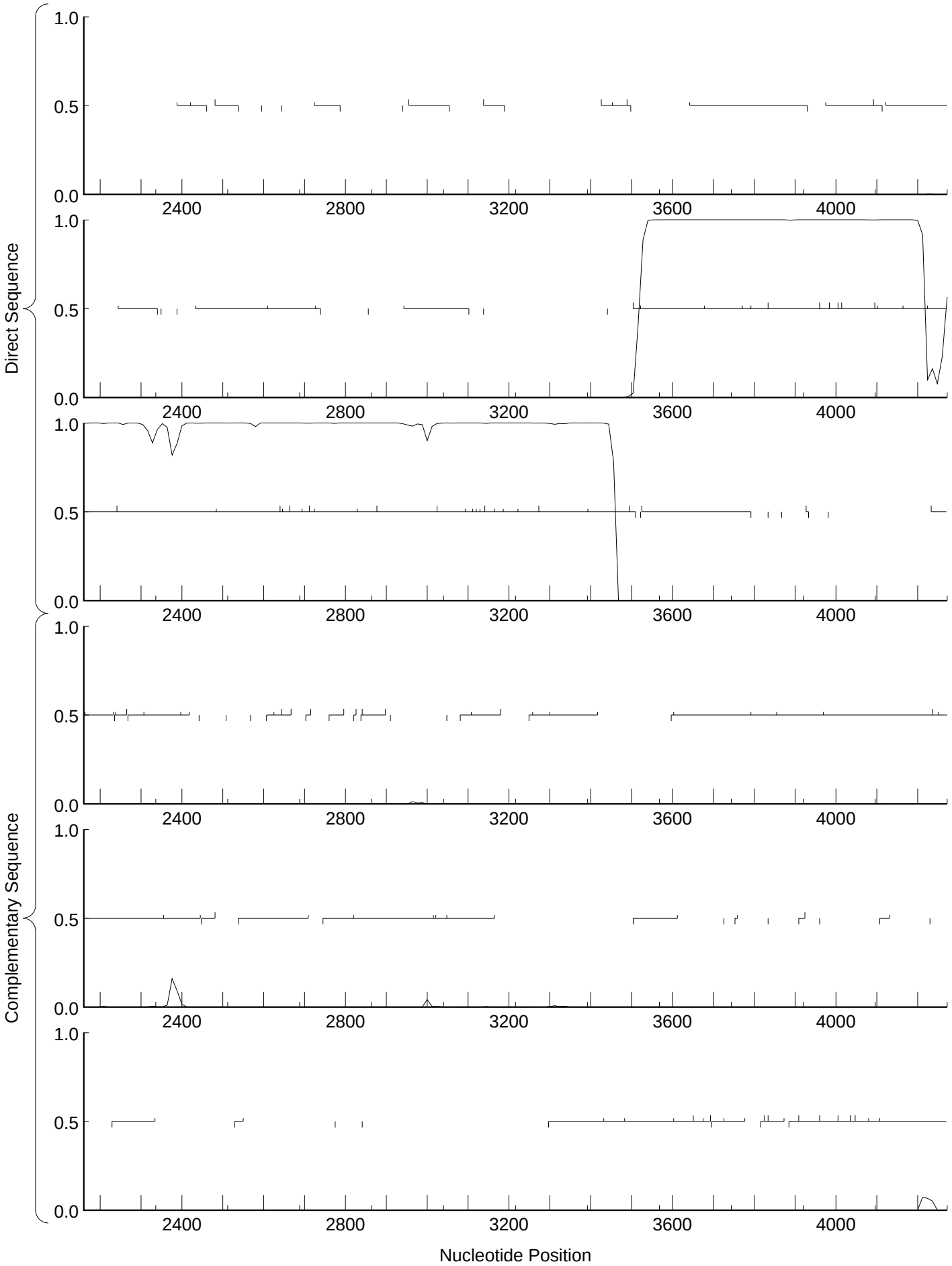
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:23:03 2014

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

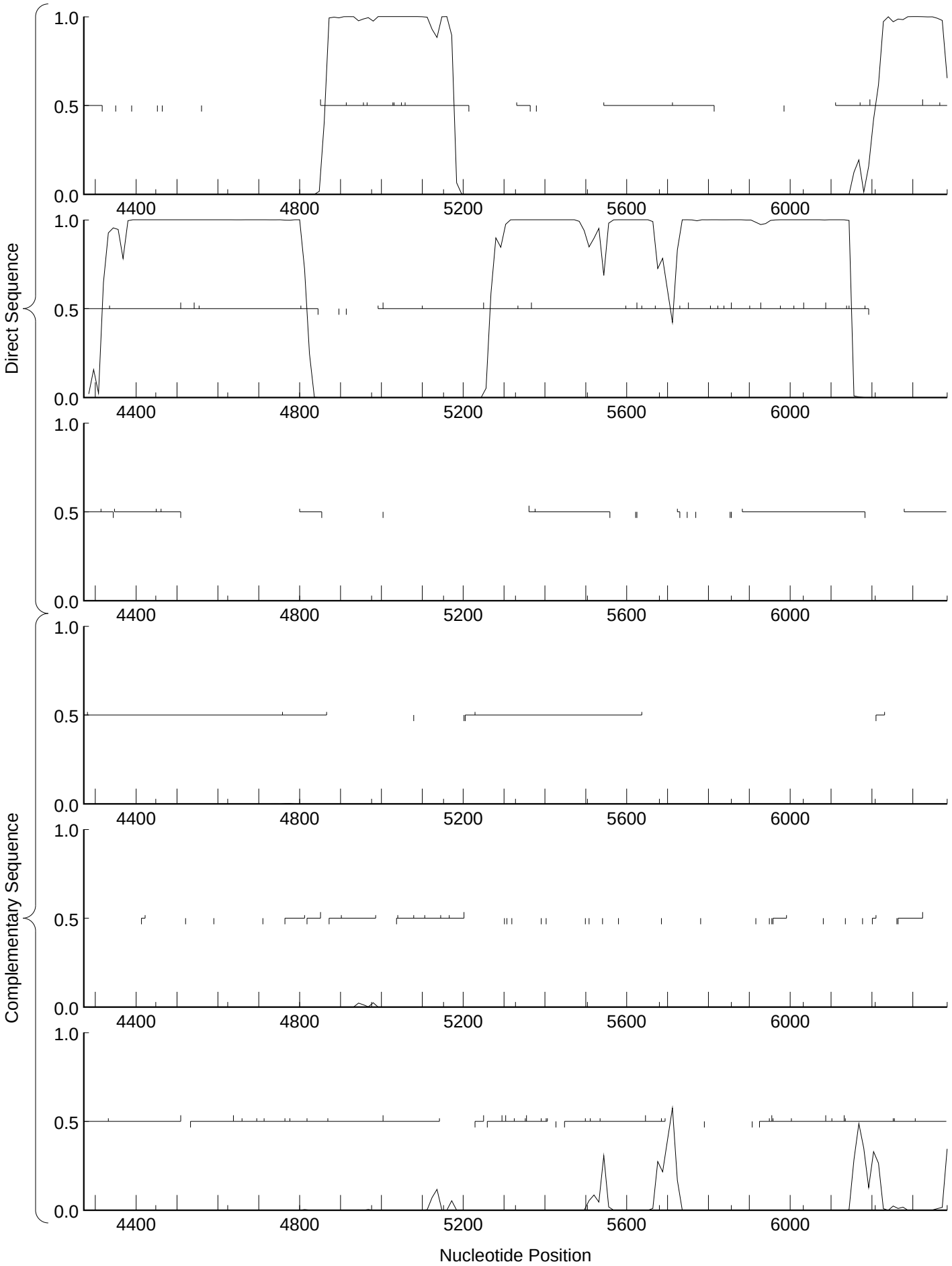
complete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 2/25



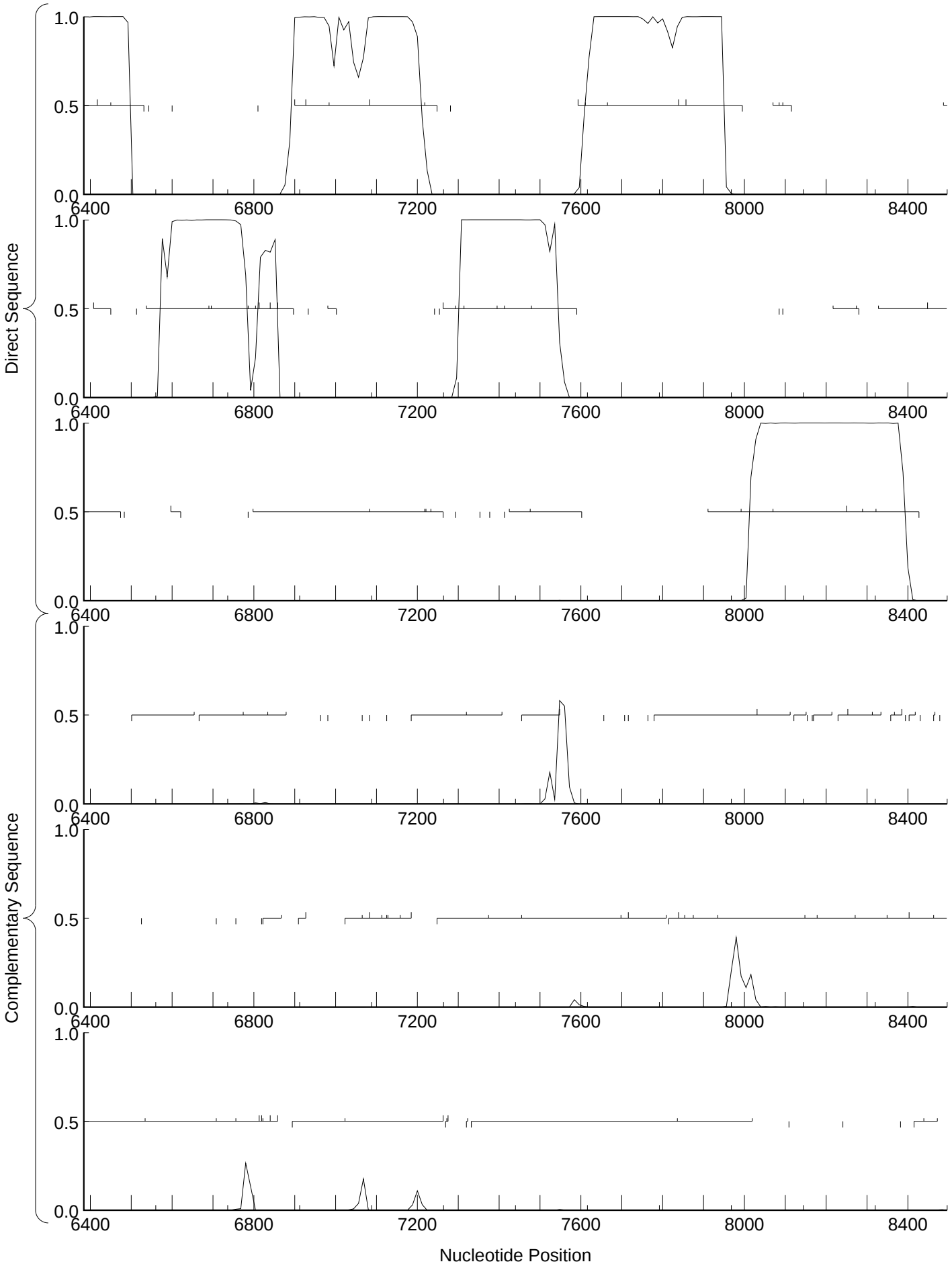
complete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 3/25



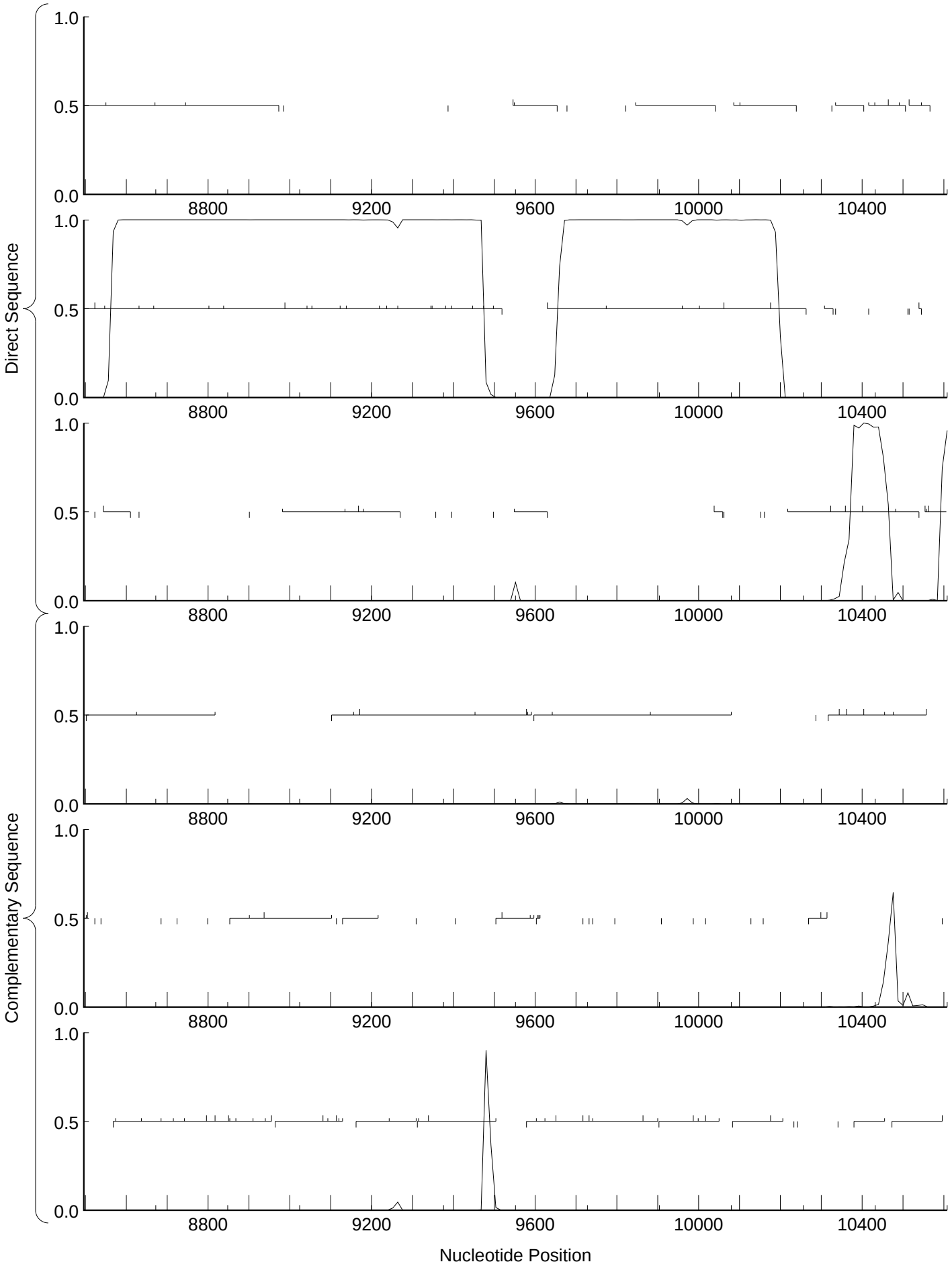
complete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 4/25



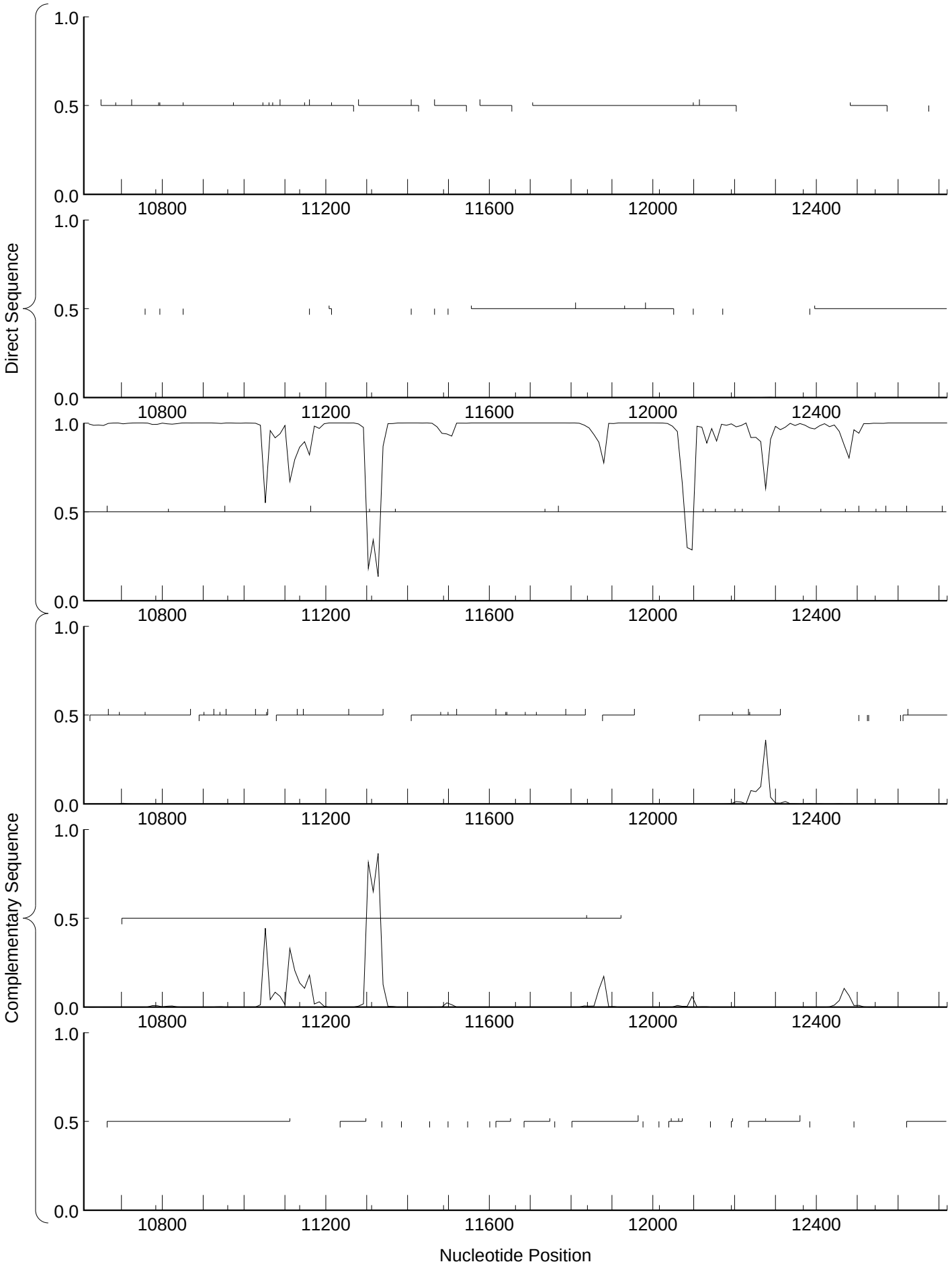
complete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 5/25



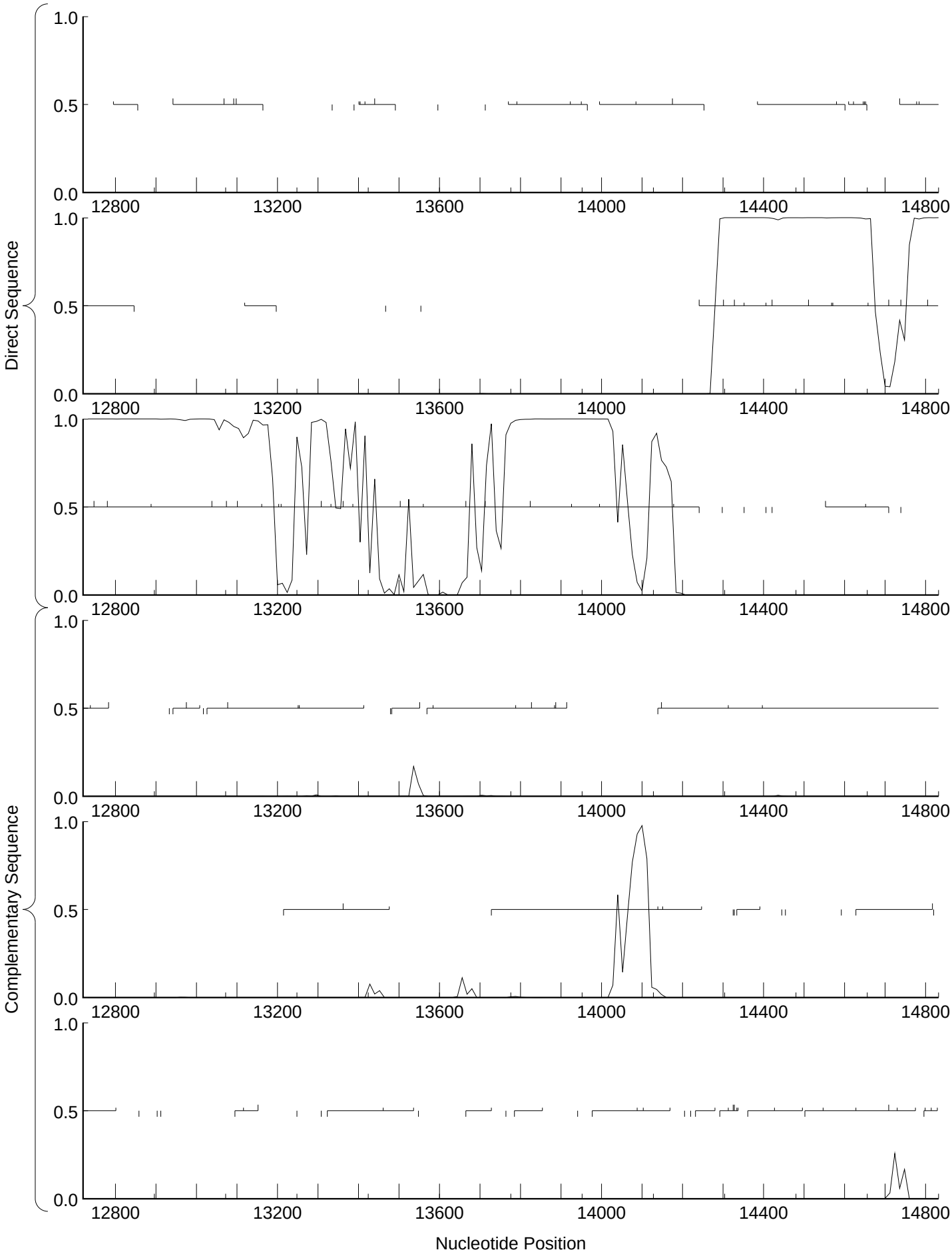
complete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 6/25



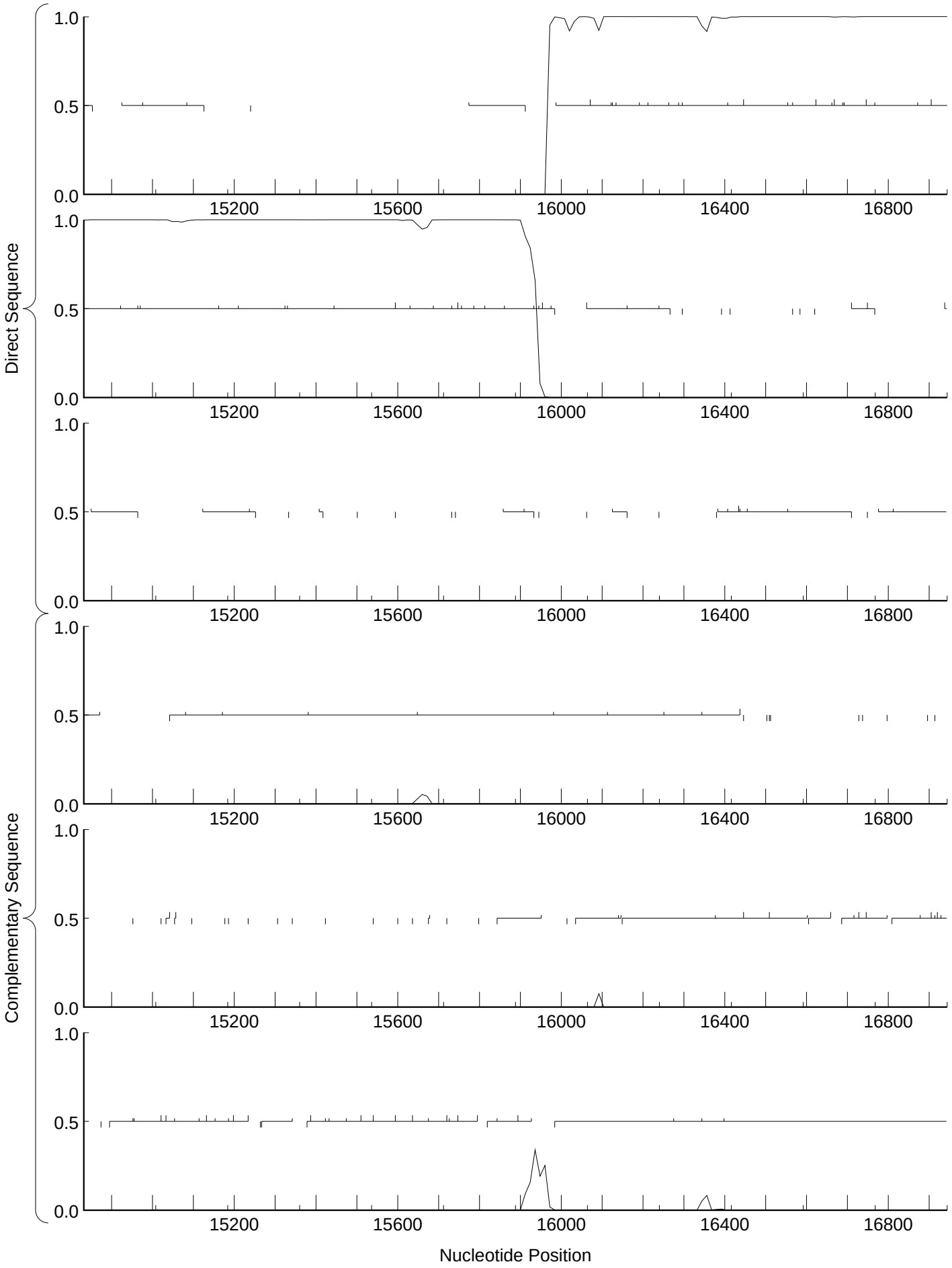
complete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 7/25



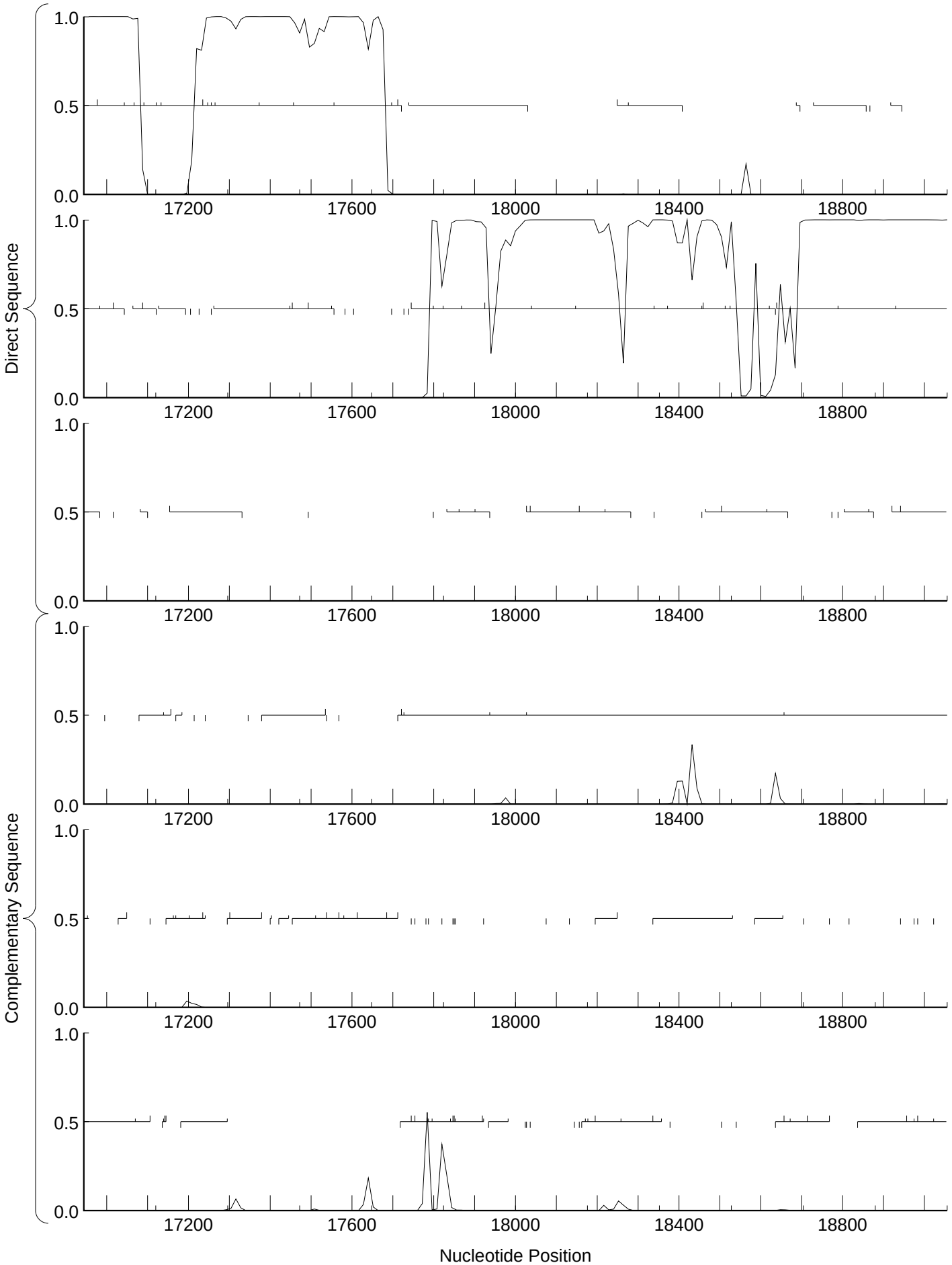
complete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 8/25



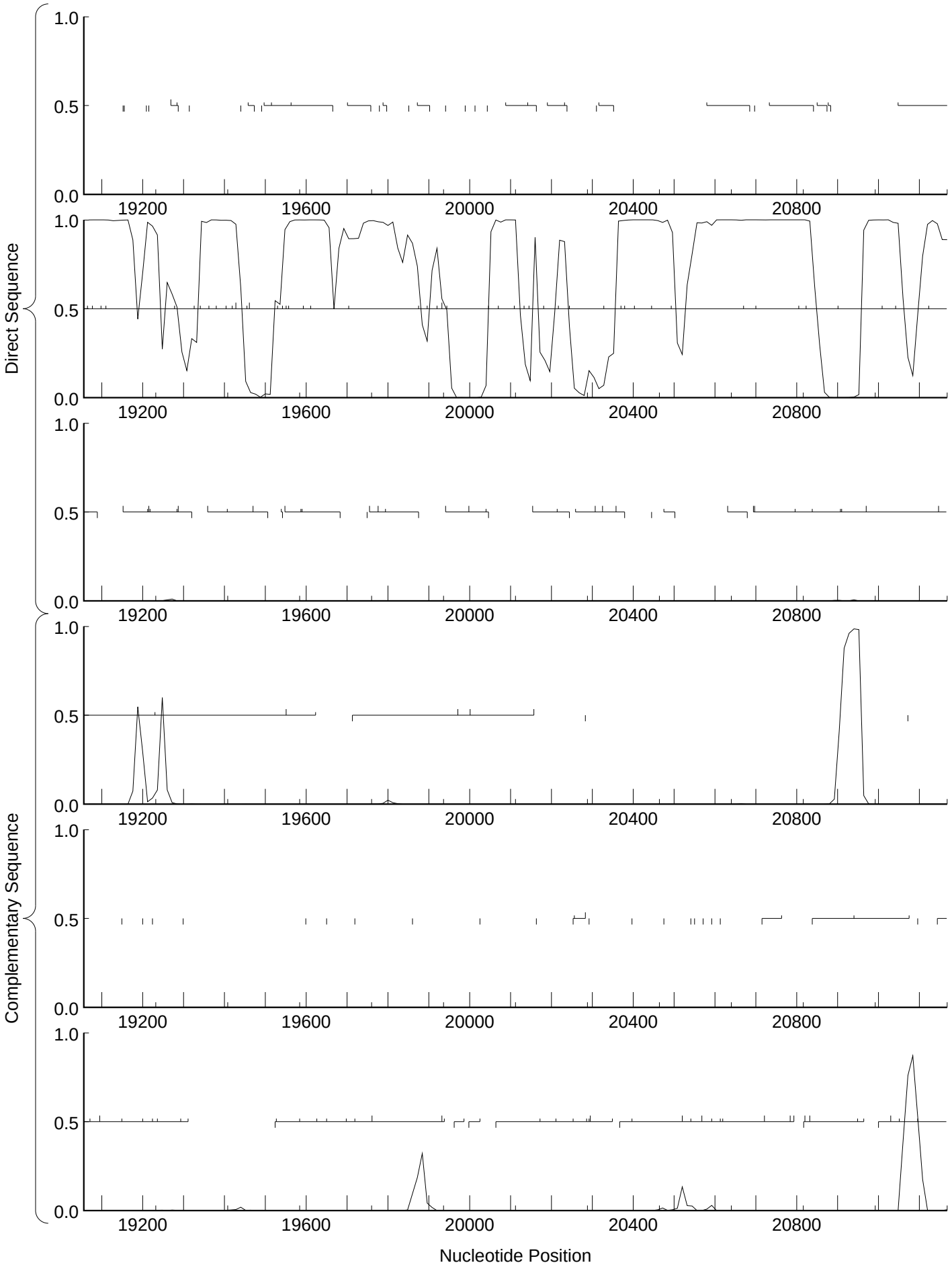
complete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 9/25



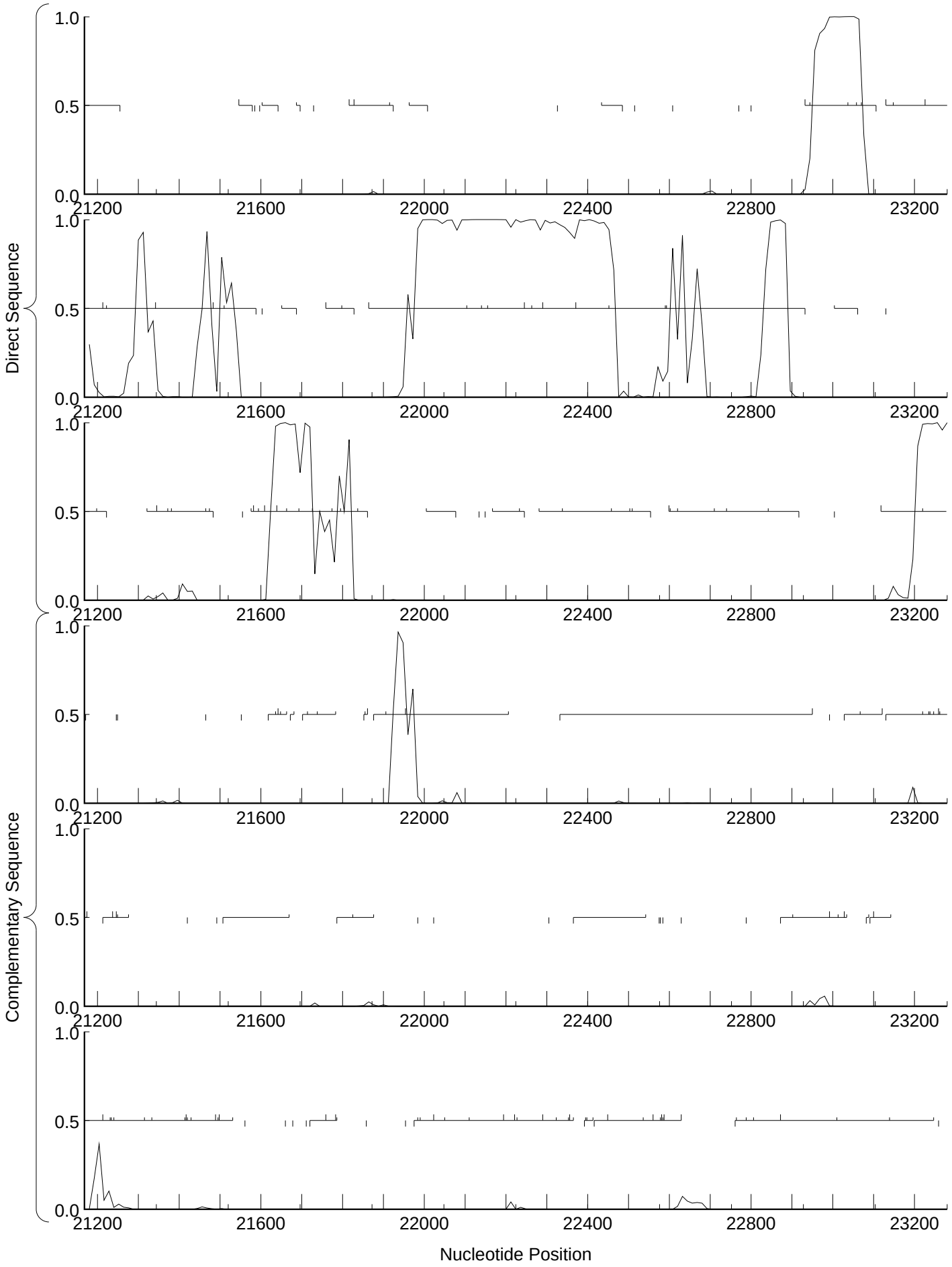
omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 10/25



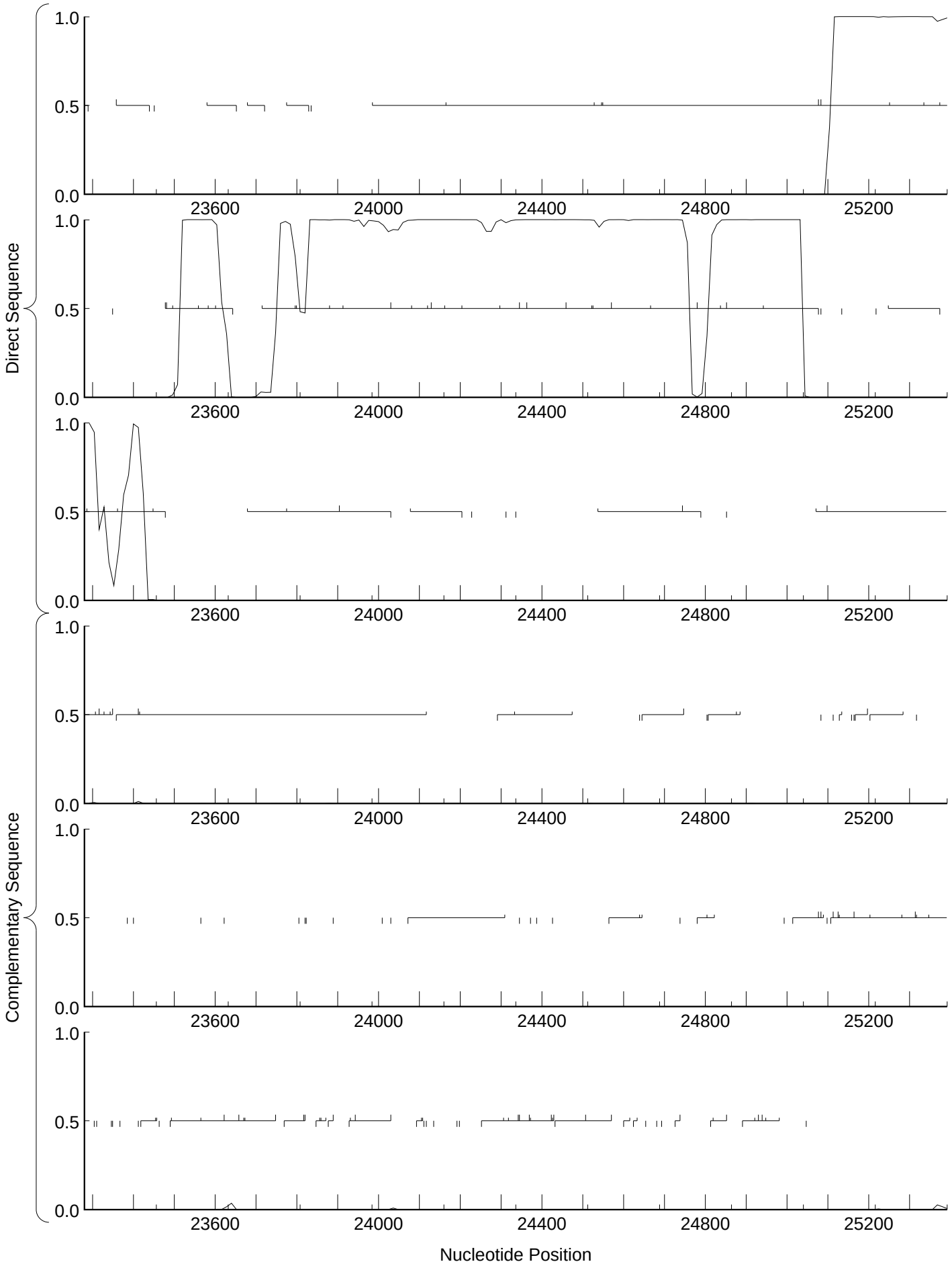
omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 11/25



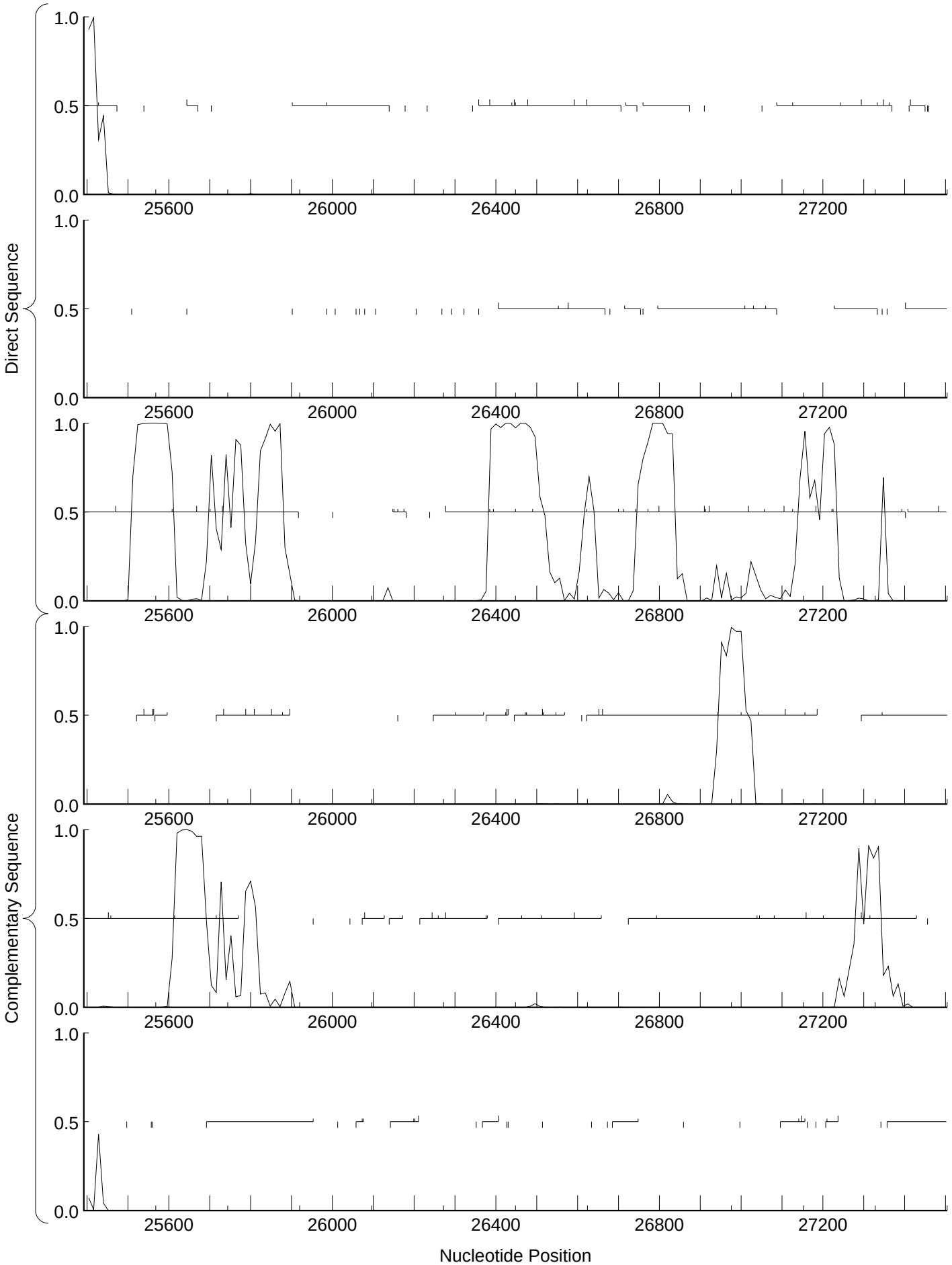
omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 12/25



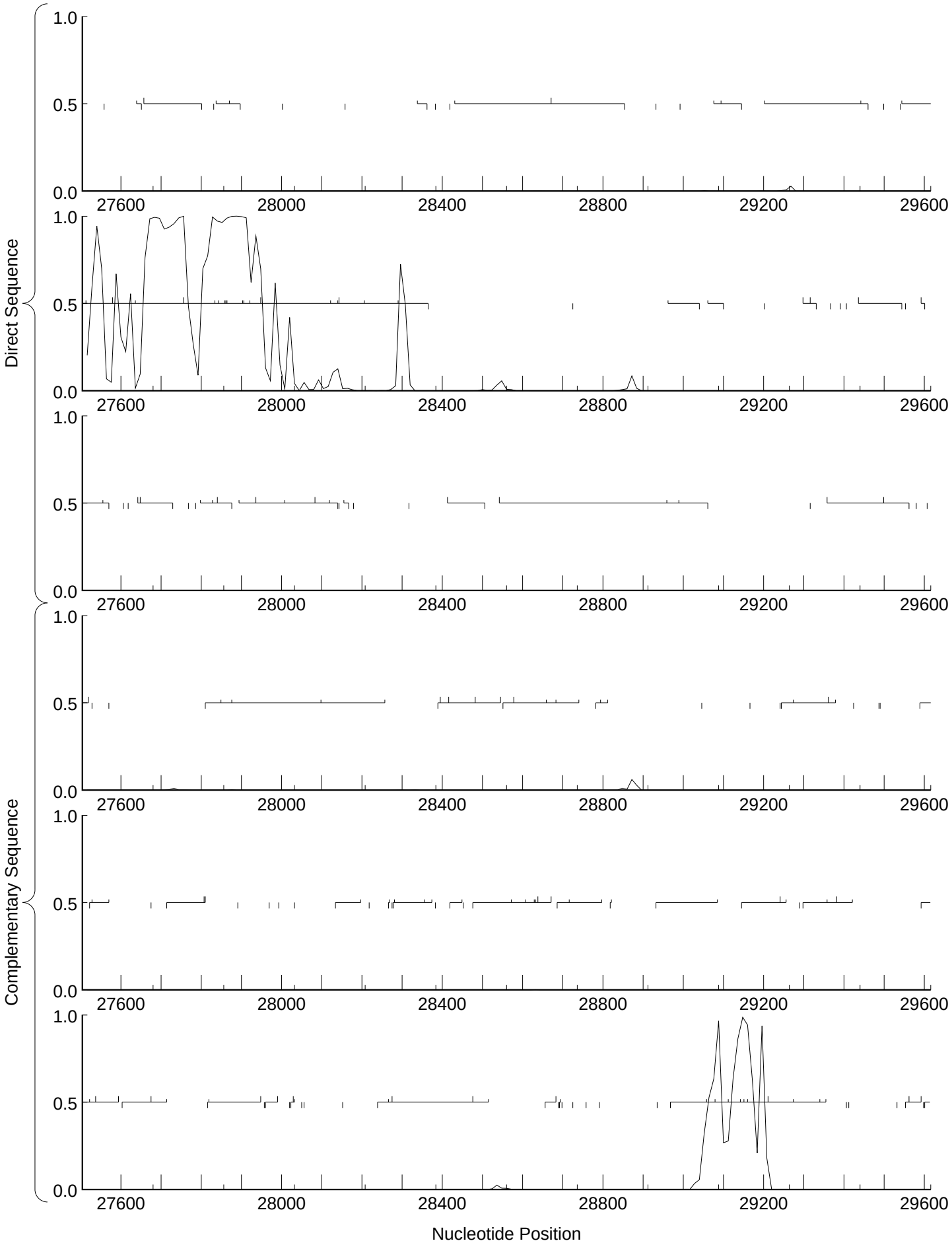
omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 13/25



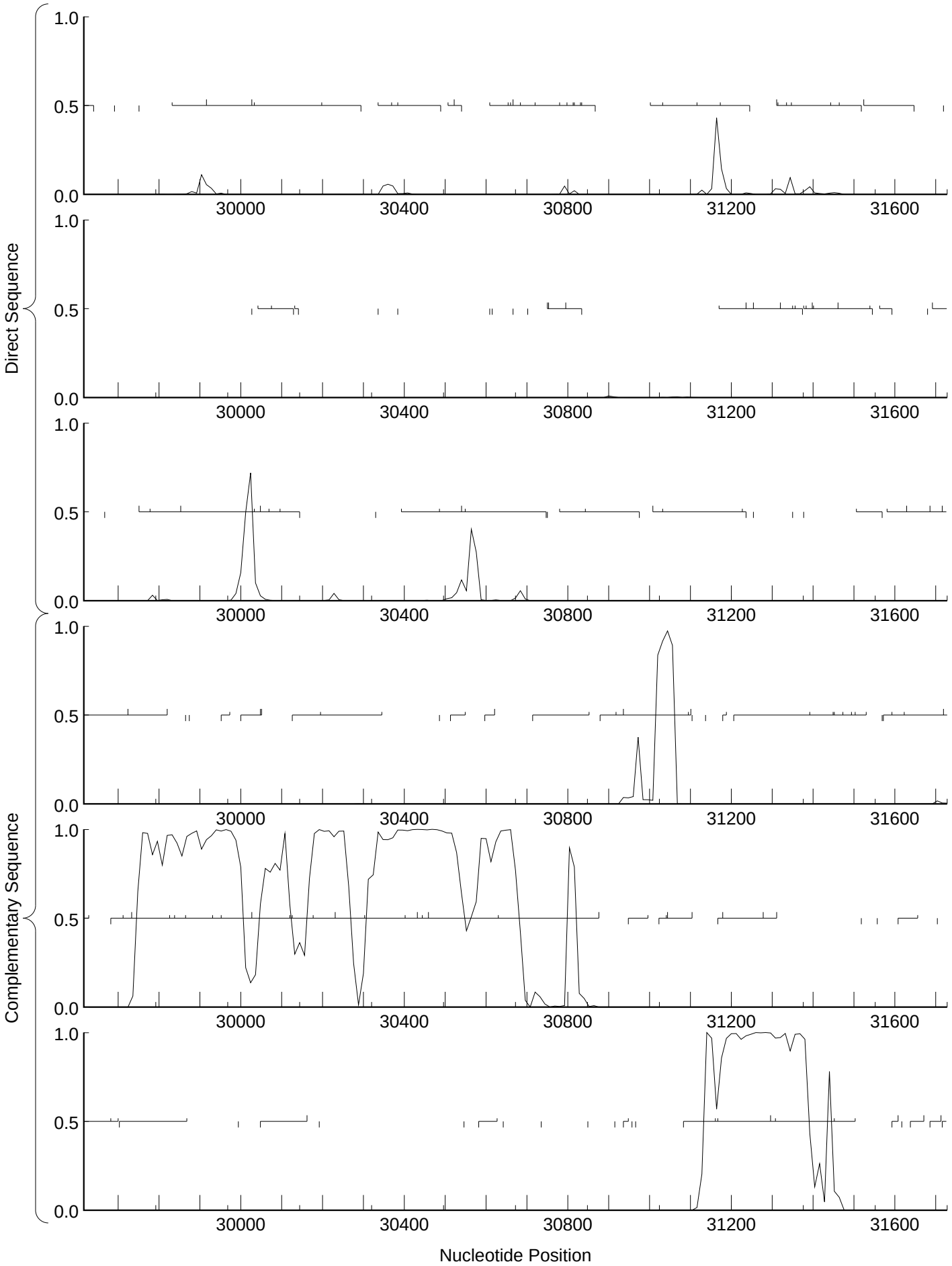
omplete sequence, 49658 bp including 12-base 3' overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 14/25



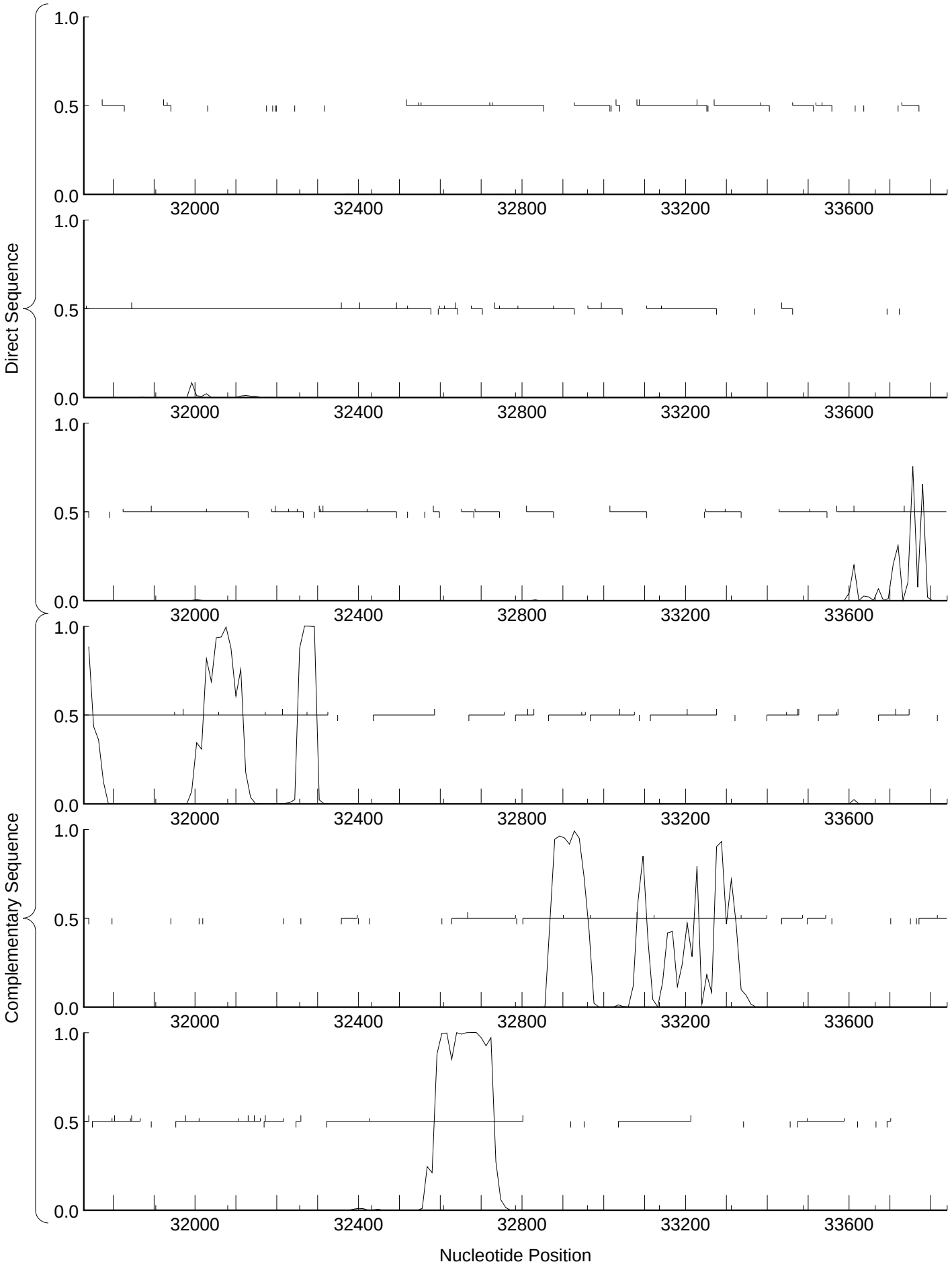
omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 15/25



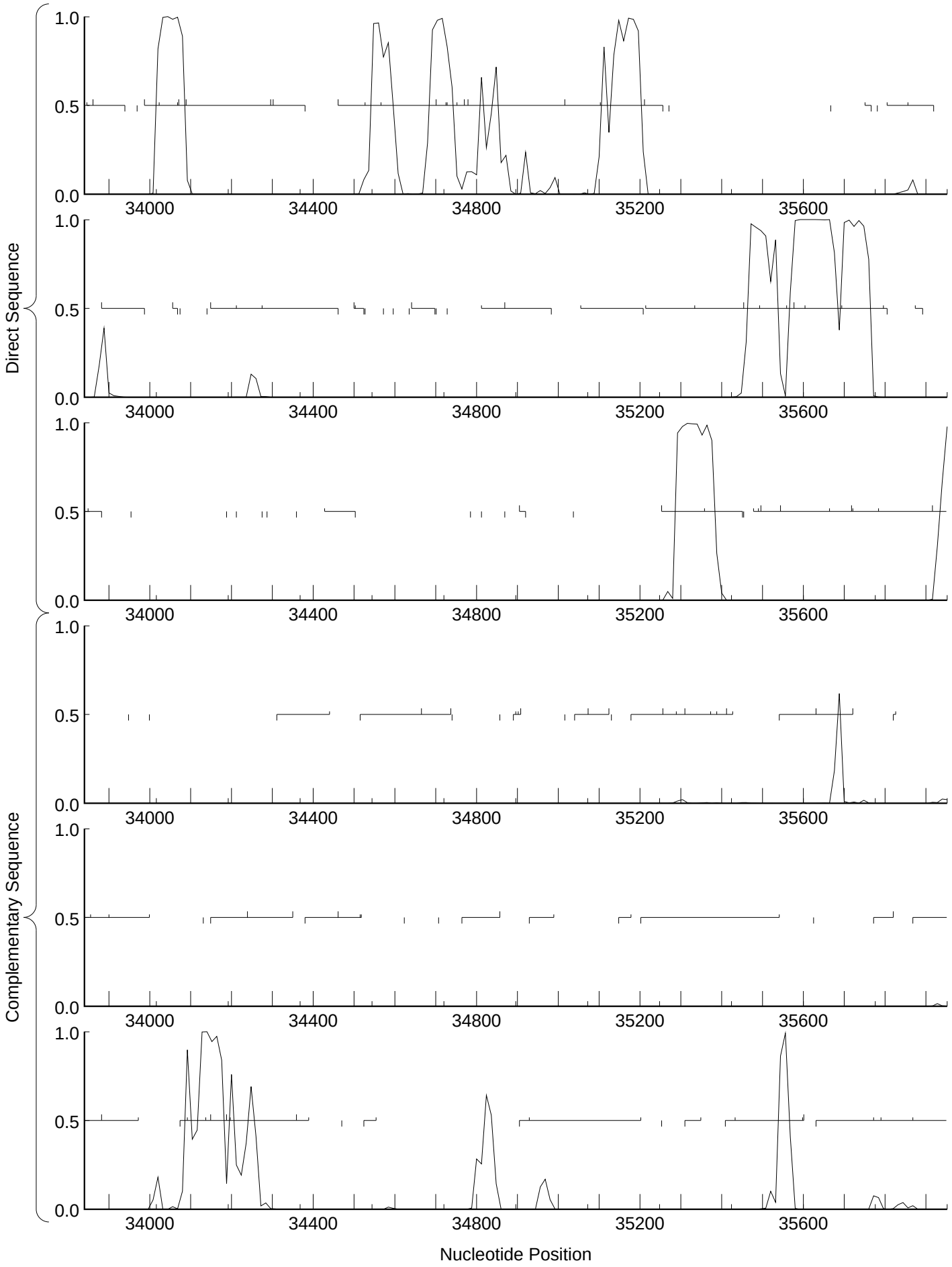
omplete sequence, 49658 bp including 12-base 3' overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 16/25



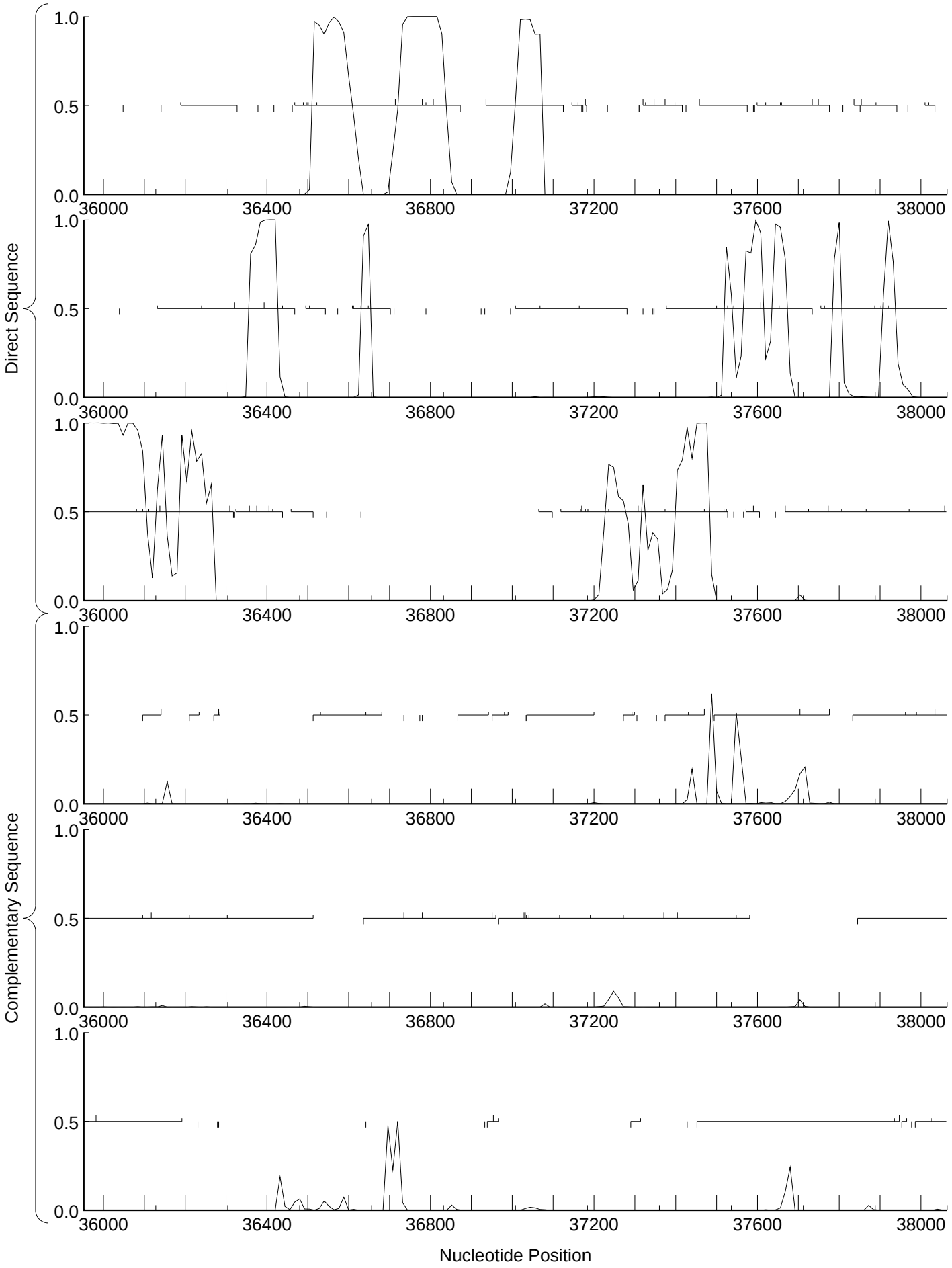
omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 17/25



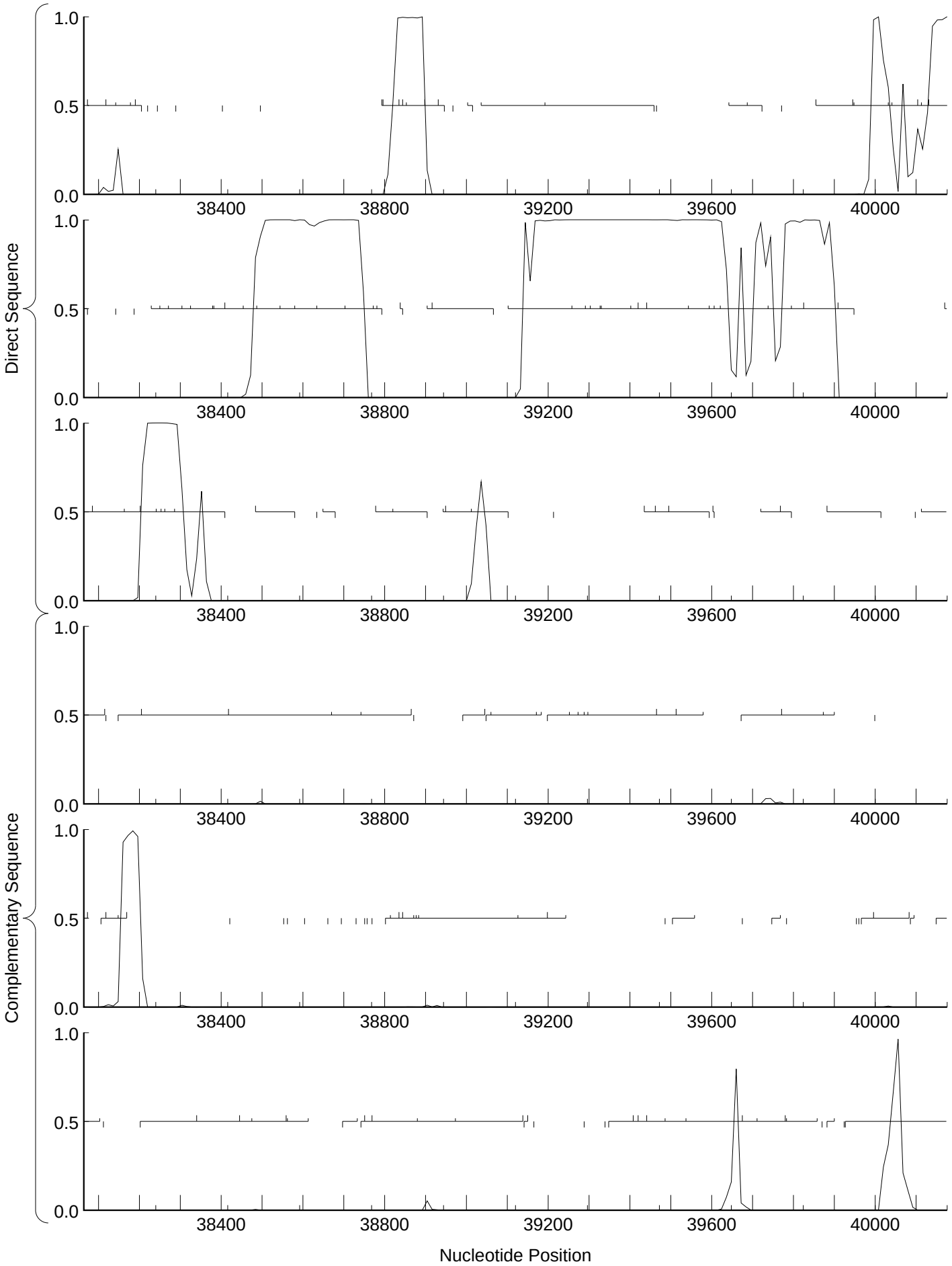
omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 18/25



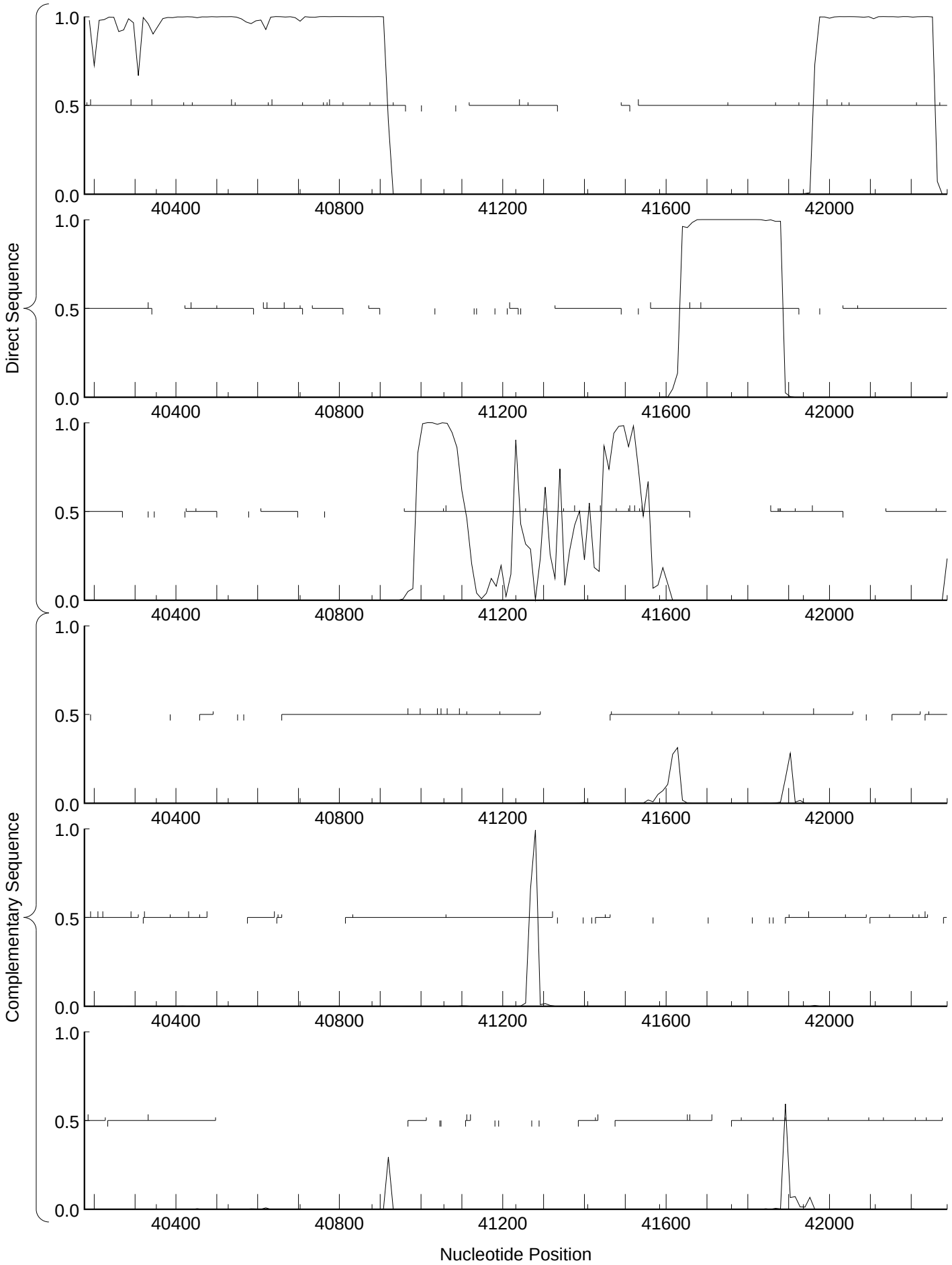
omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 19/25



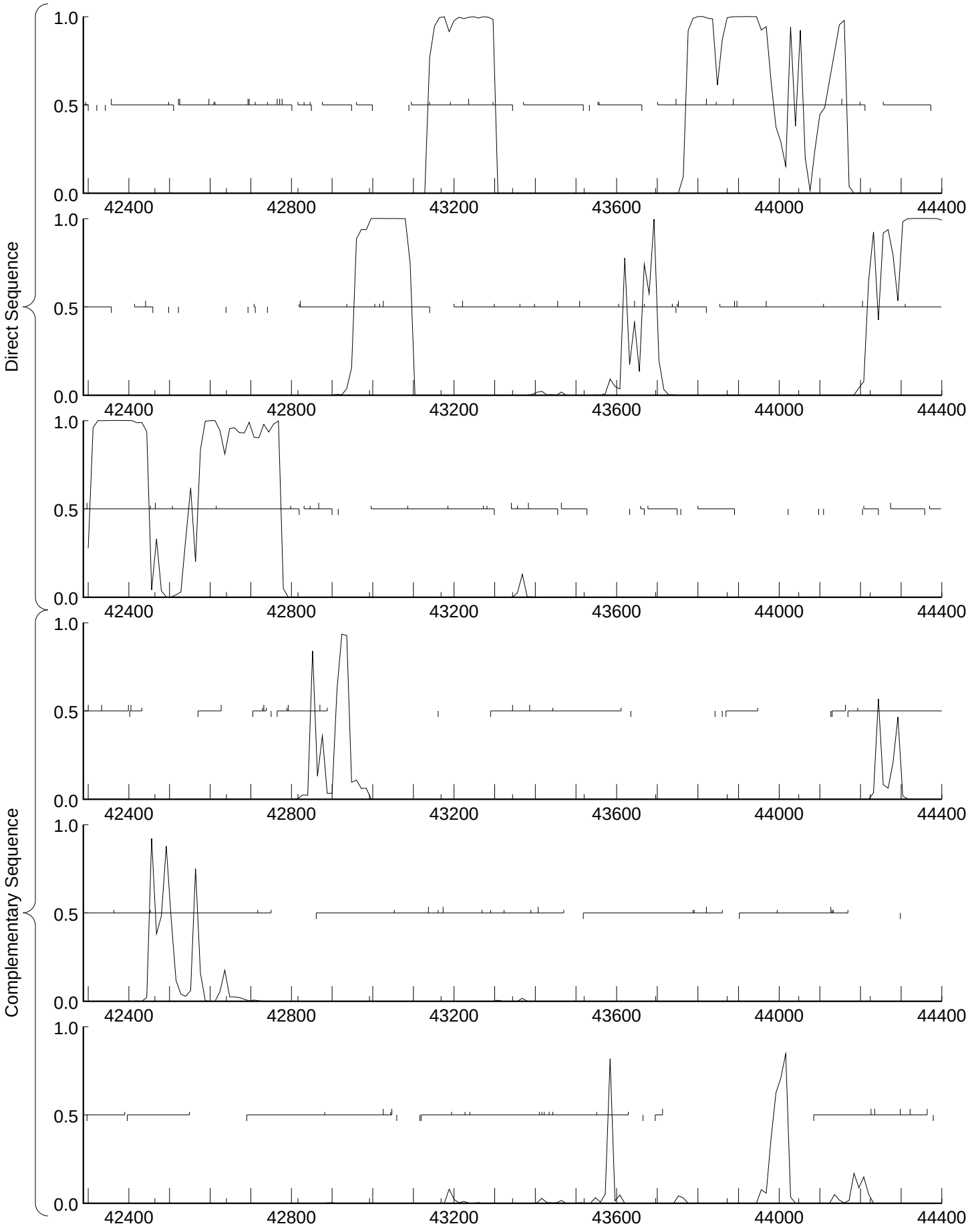
omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 20/25



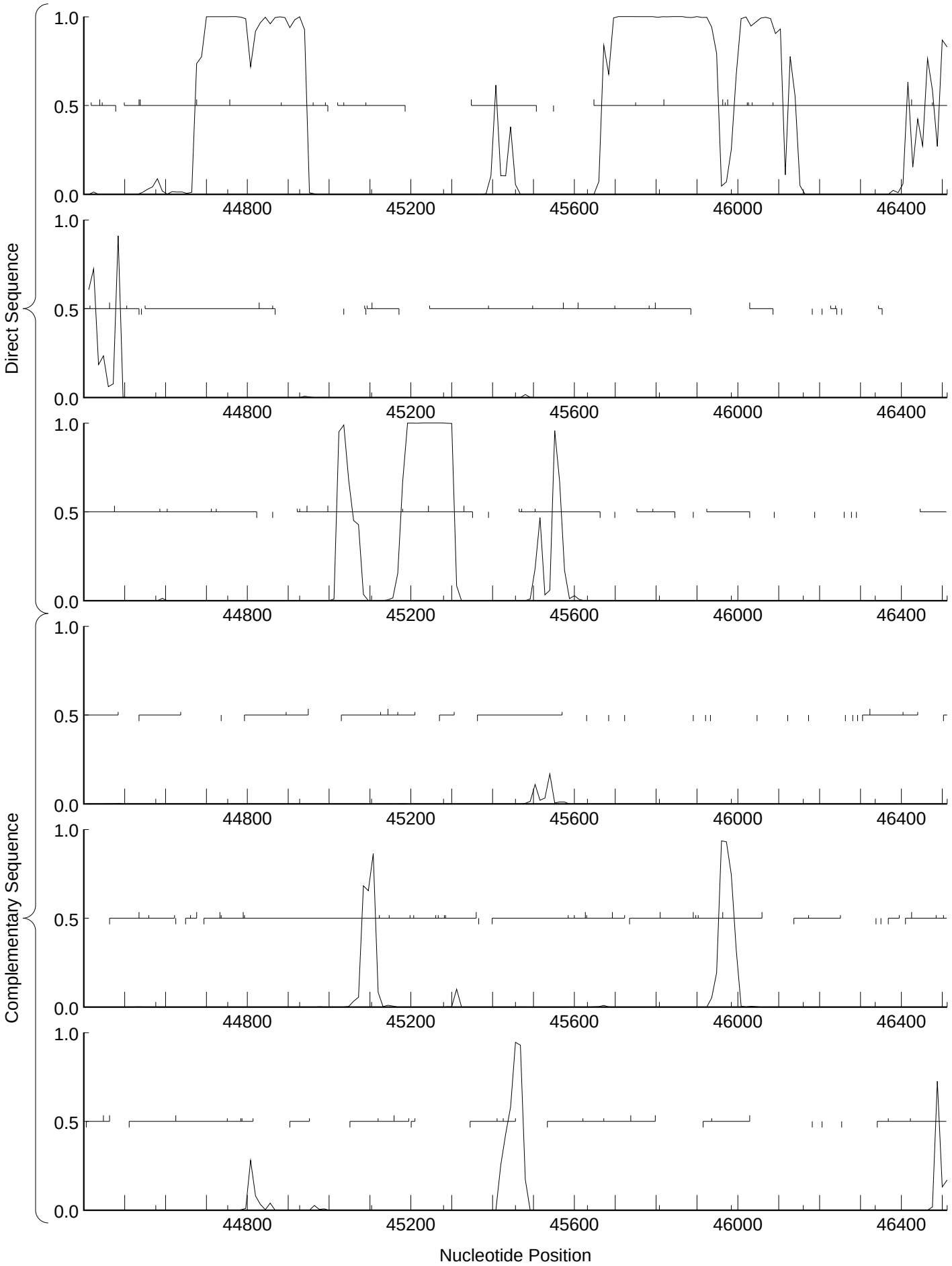
omplete sequence, 49658 bp including 12-base 3' overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 21/25



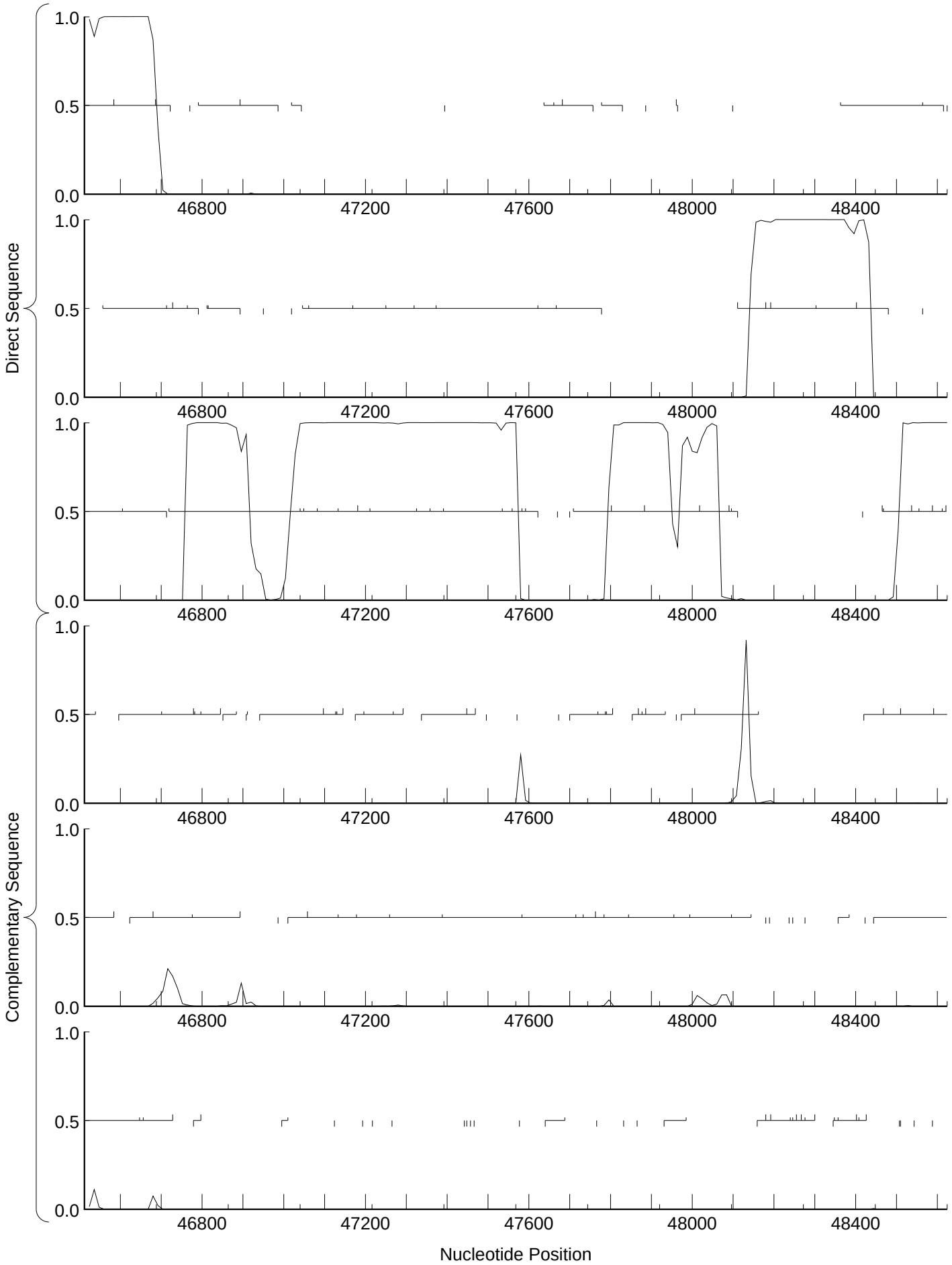
omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 22/25



omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 23/25



omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 24/25



omplete sequence, 49658 bp including 12-base 3’ overhang (CCTGCCGCCCGC), Cluster P, Order 4, Window 96, Step 12, 25/25

