# GeneMark

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

## PROGRAM INFORMATION

```
Sequence         :  Gordonia phage SoilAssassin complete sequence, 47880 bp including 11-base 3’ overhang (TACC
Analysis Date    :  3/28/24 at 8:27:22
Pages            :  24
Sequence Length  :  47880 bp
GC Content       :   66.82%

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 notes & comments**

Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep  8 00:36:14 2018

## MATRIX INFORMATION

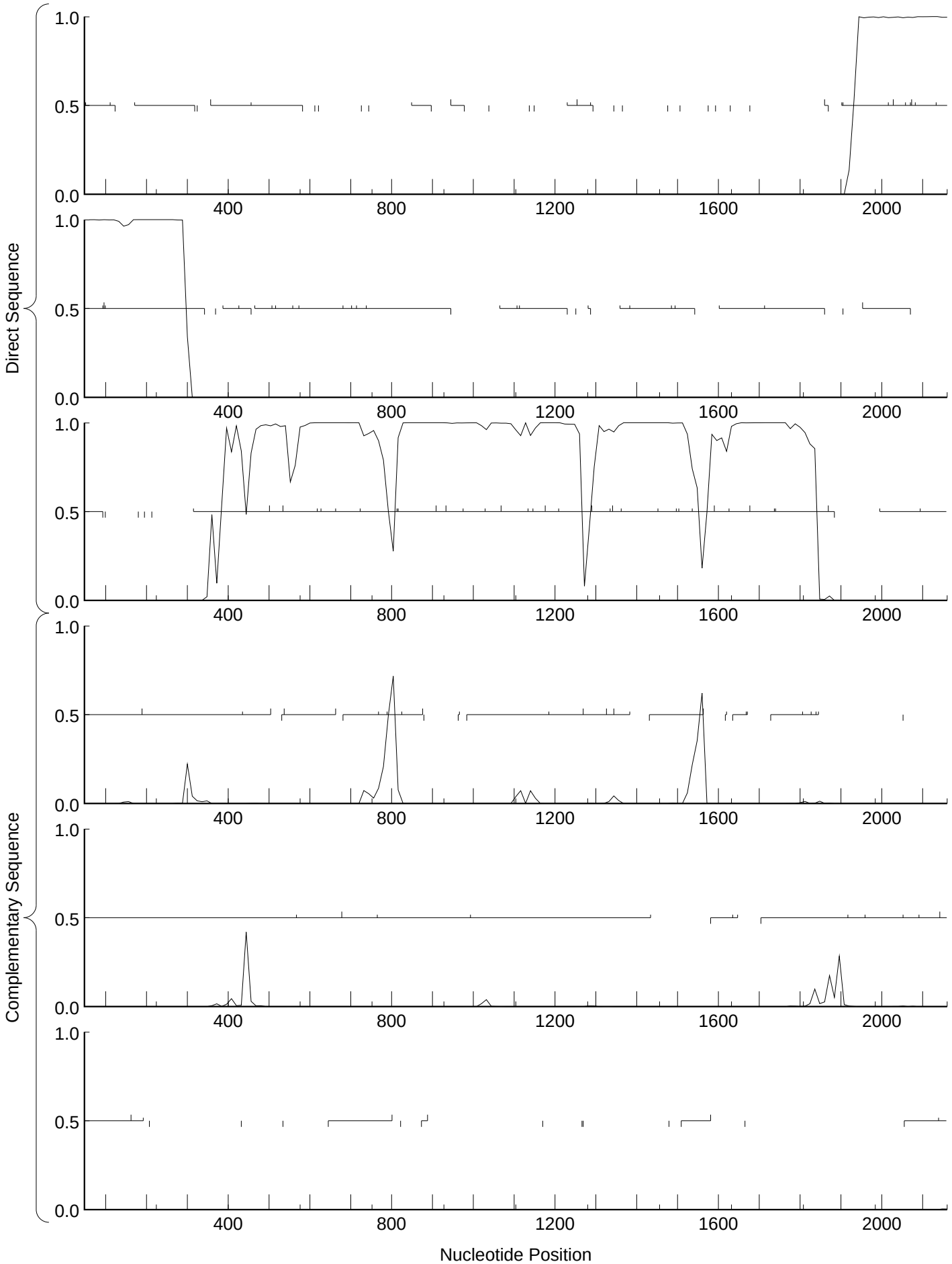
```
Matrix     :  -
Author     :  -

Order      :  4
```

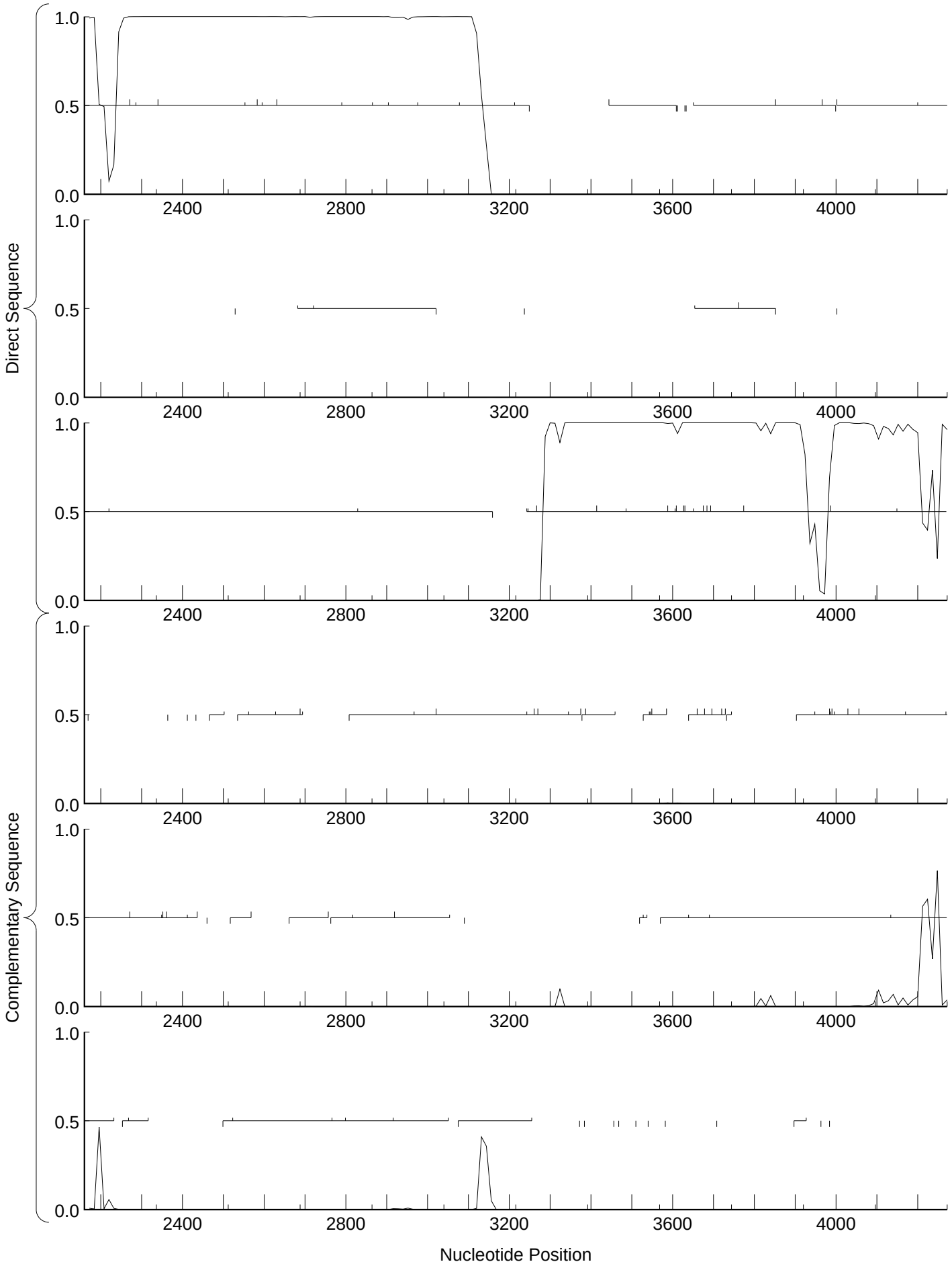

Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA  30332-0230


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

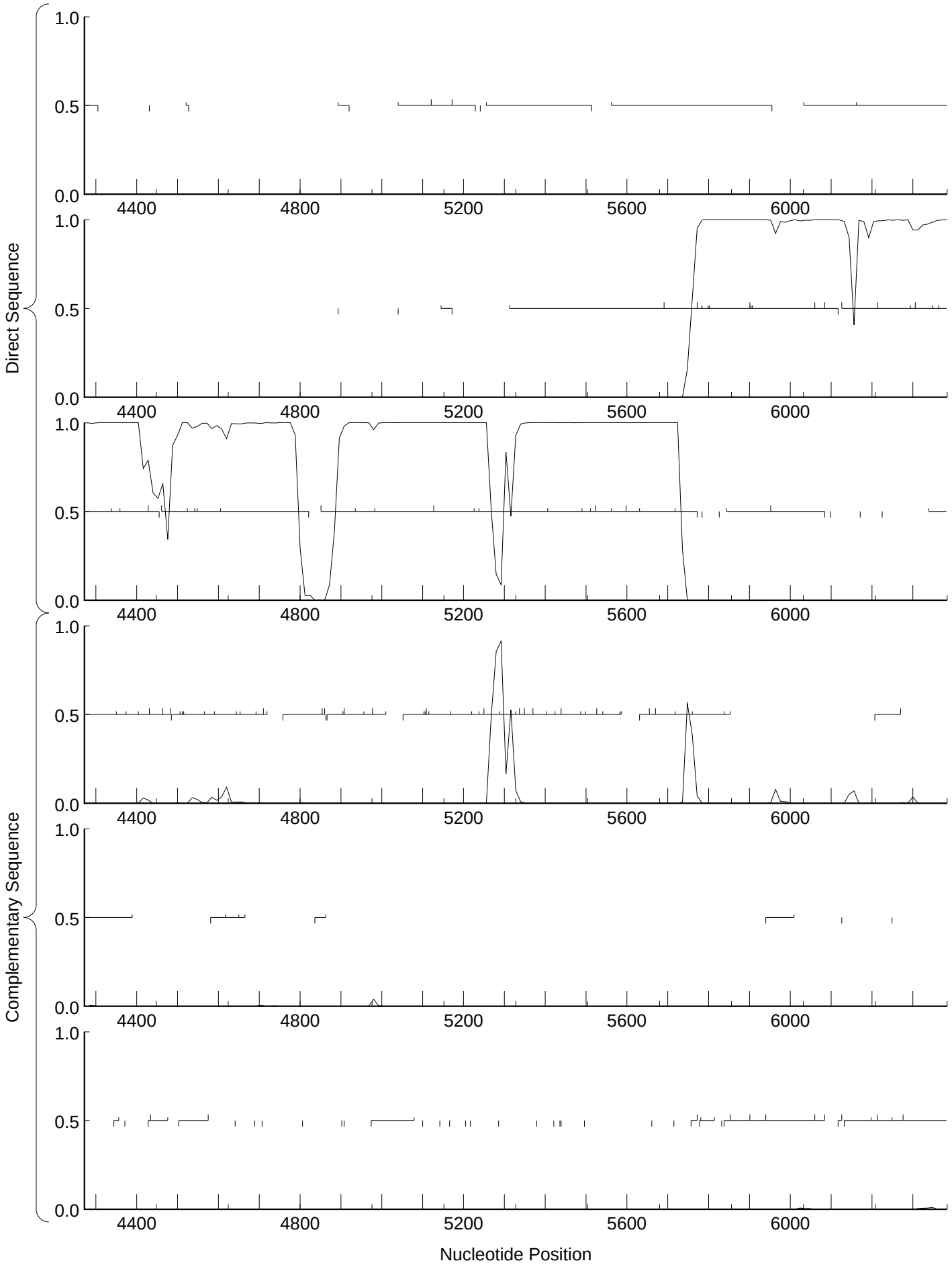
omplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 2/24



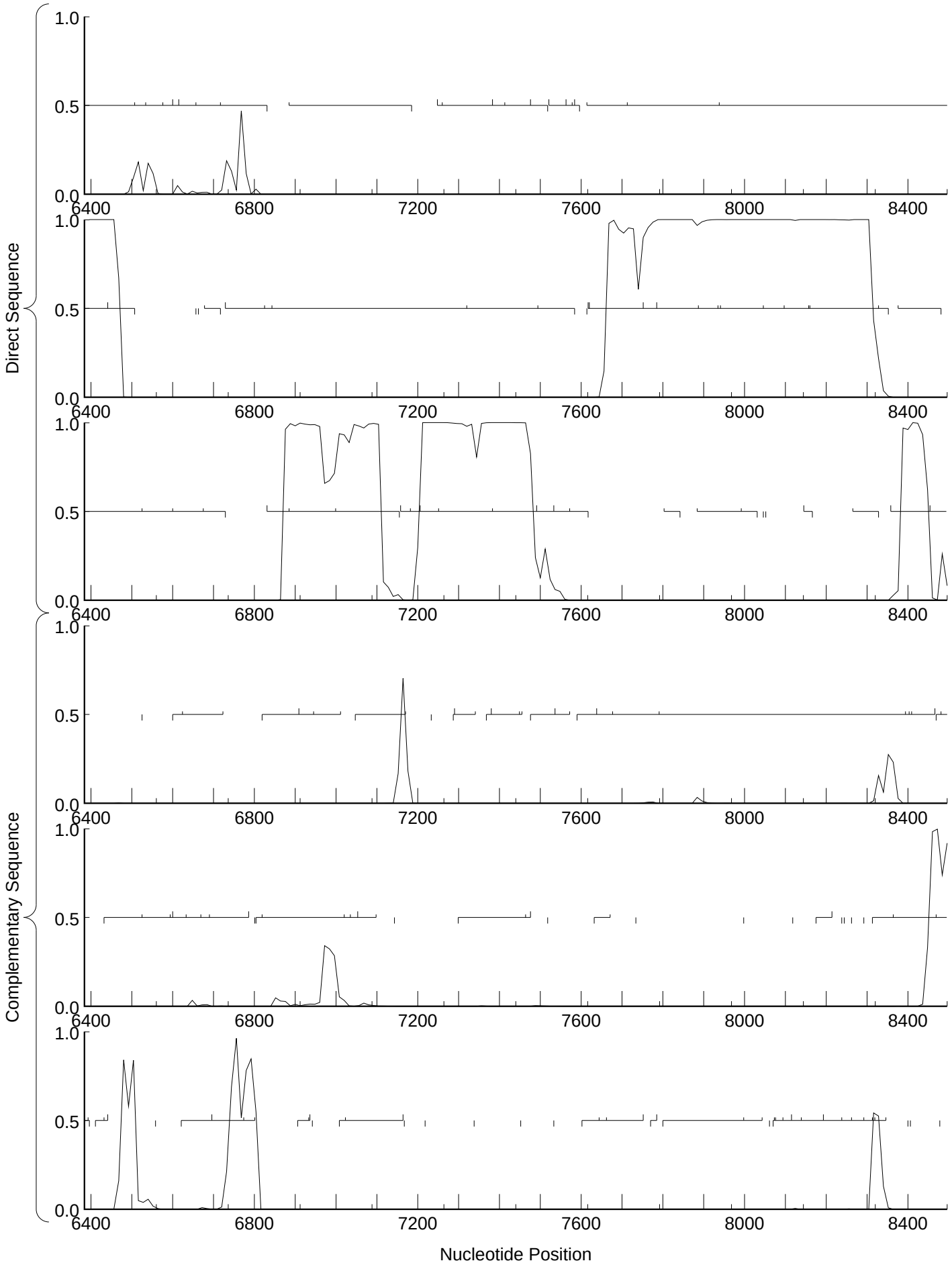
omplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 3/24



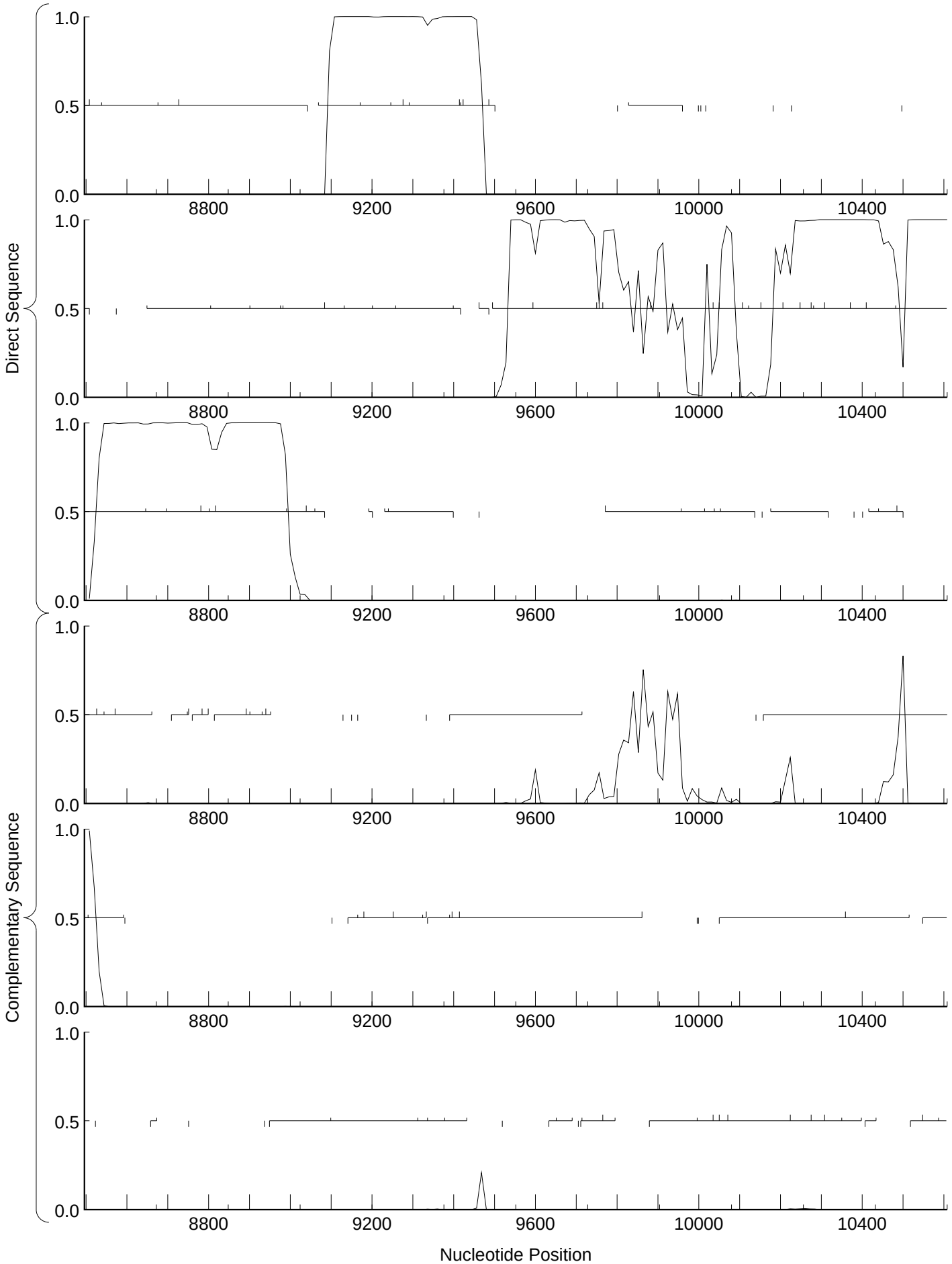
omplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 4/24



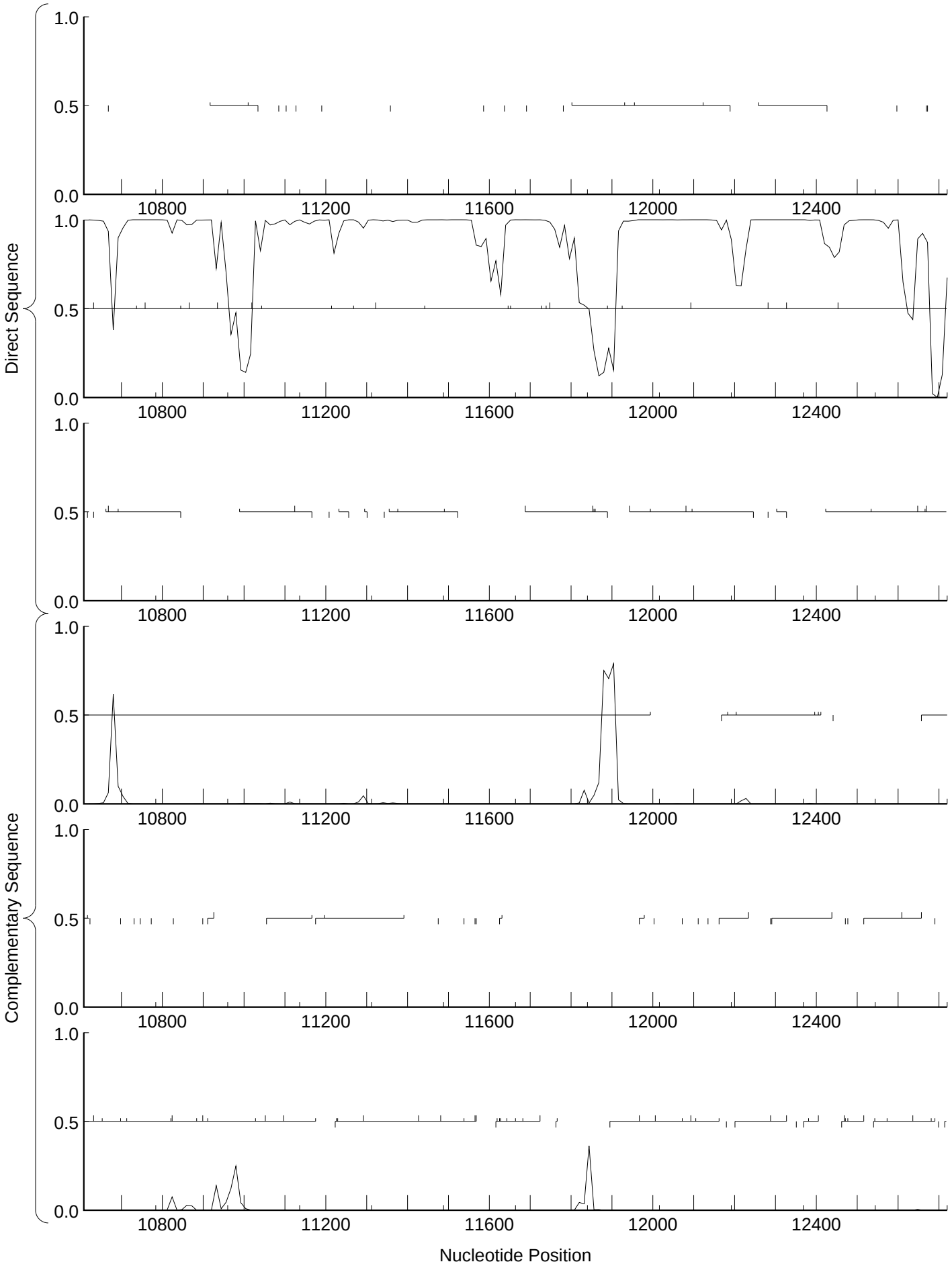
omplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 5/24



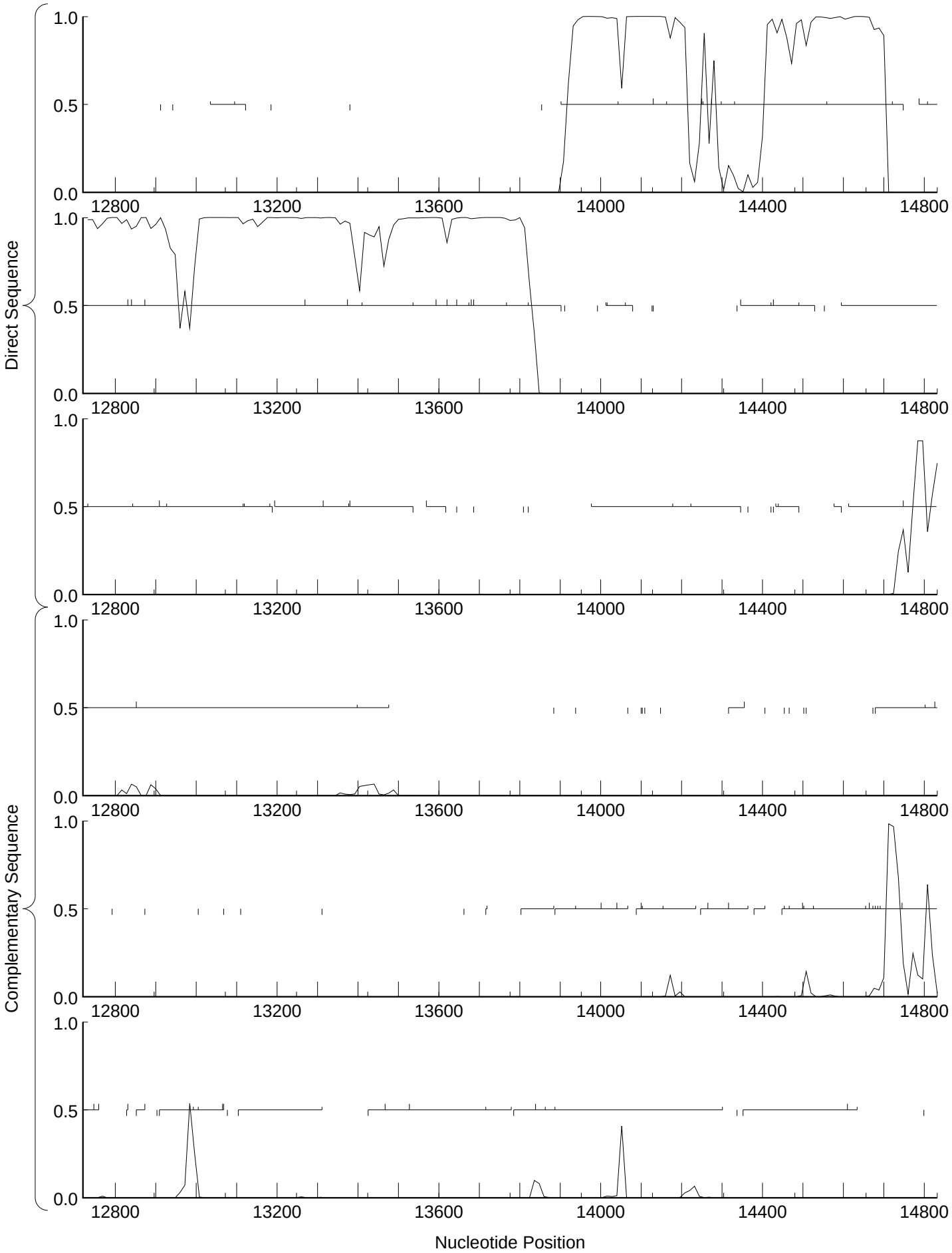
omplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 6/24



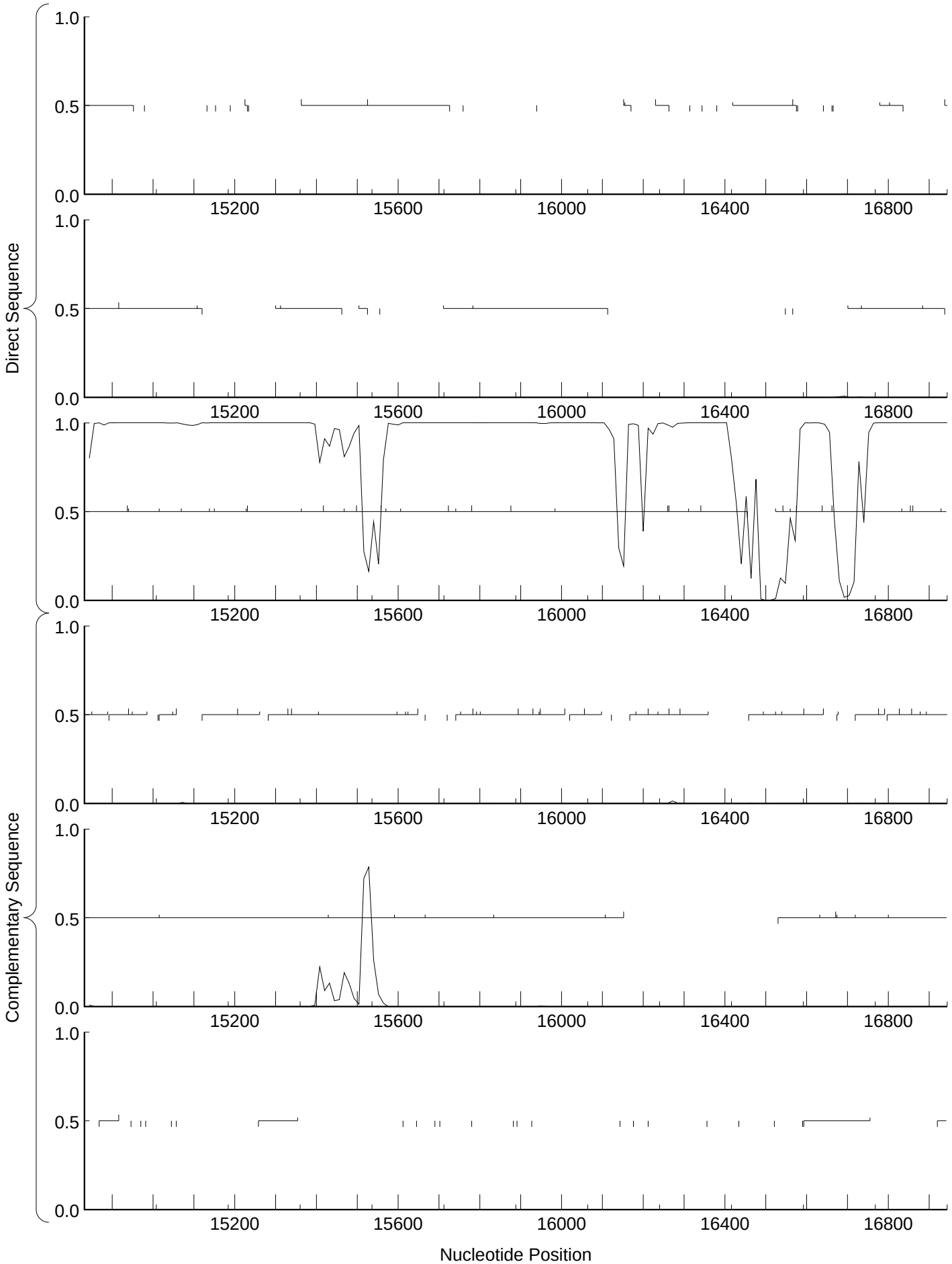
omplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 7/24



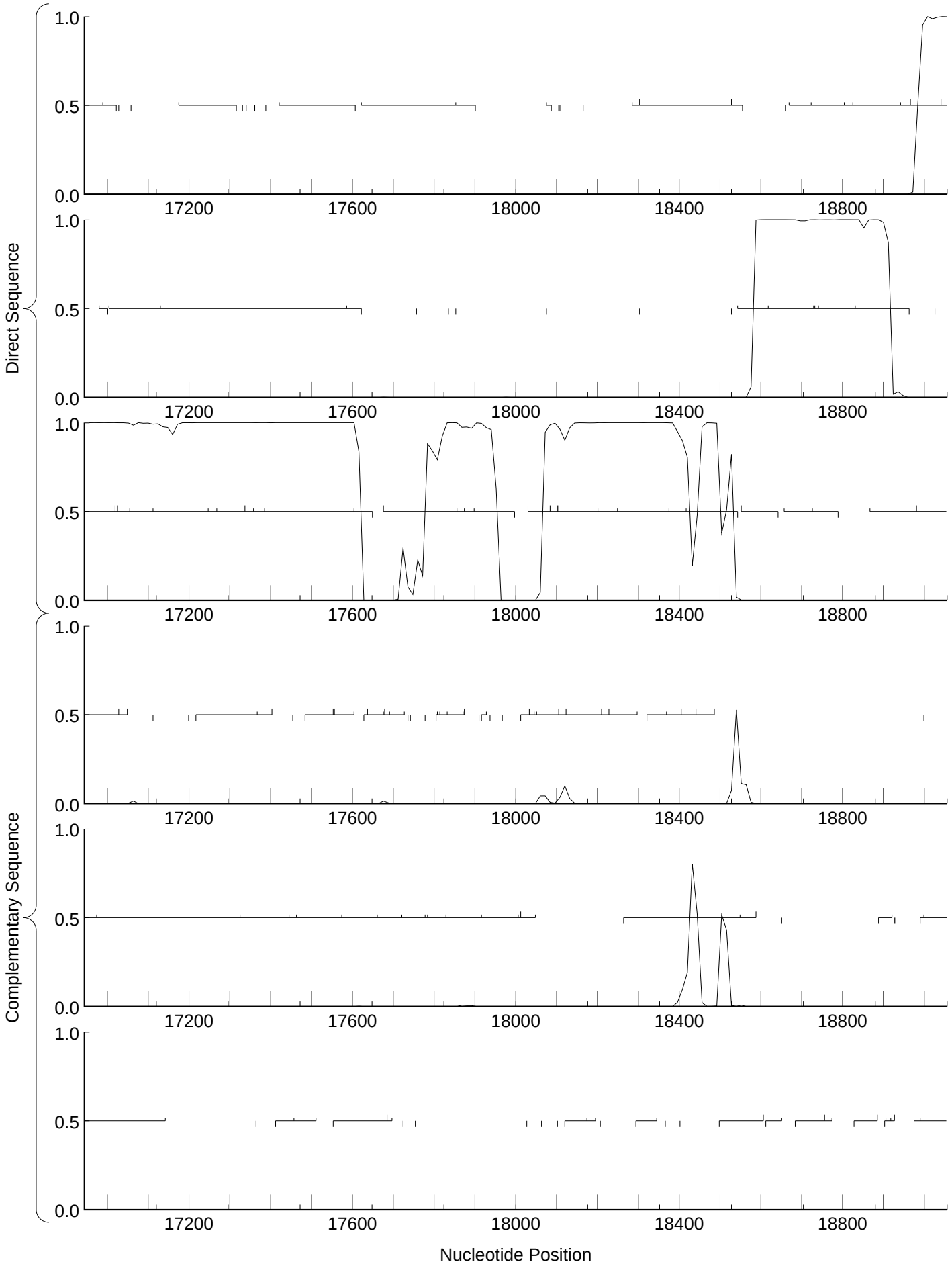
omplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 8/24



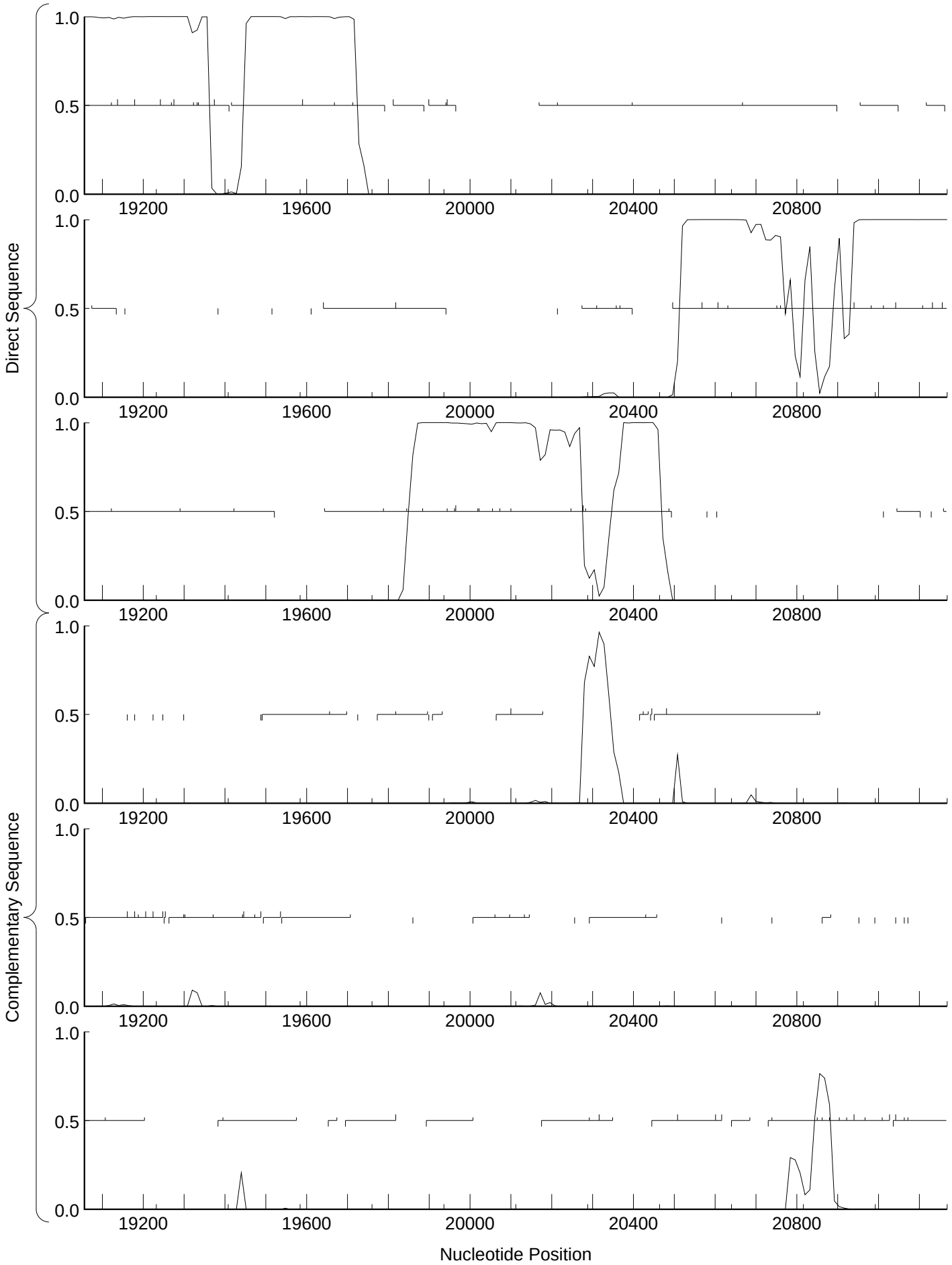
omplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 9/24



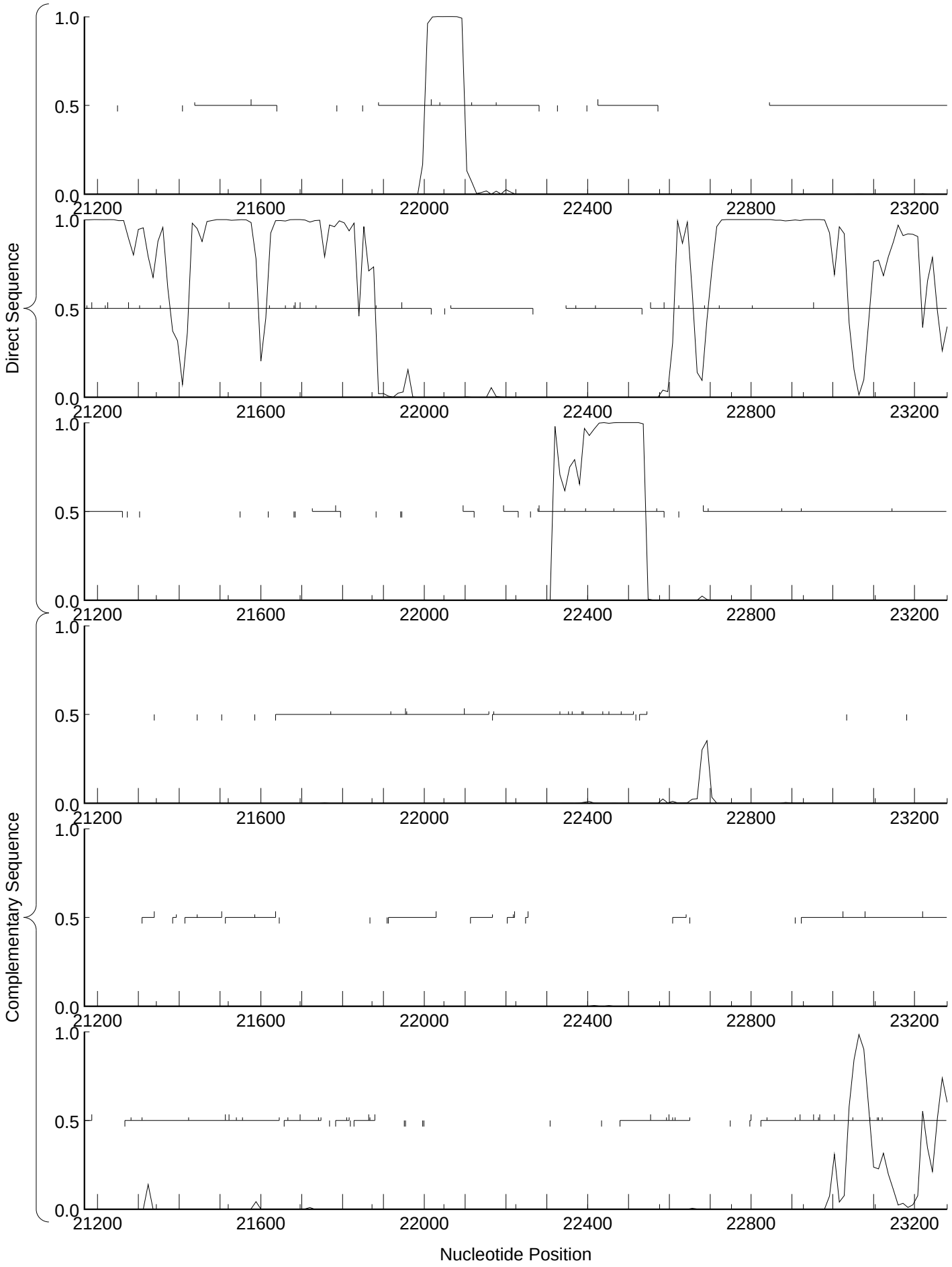
mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 10/24



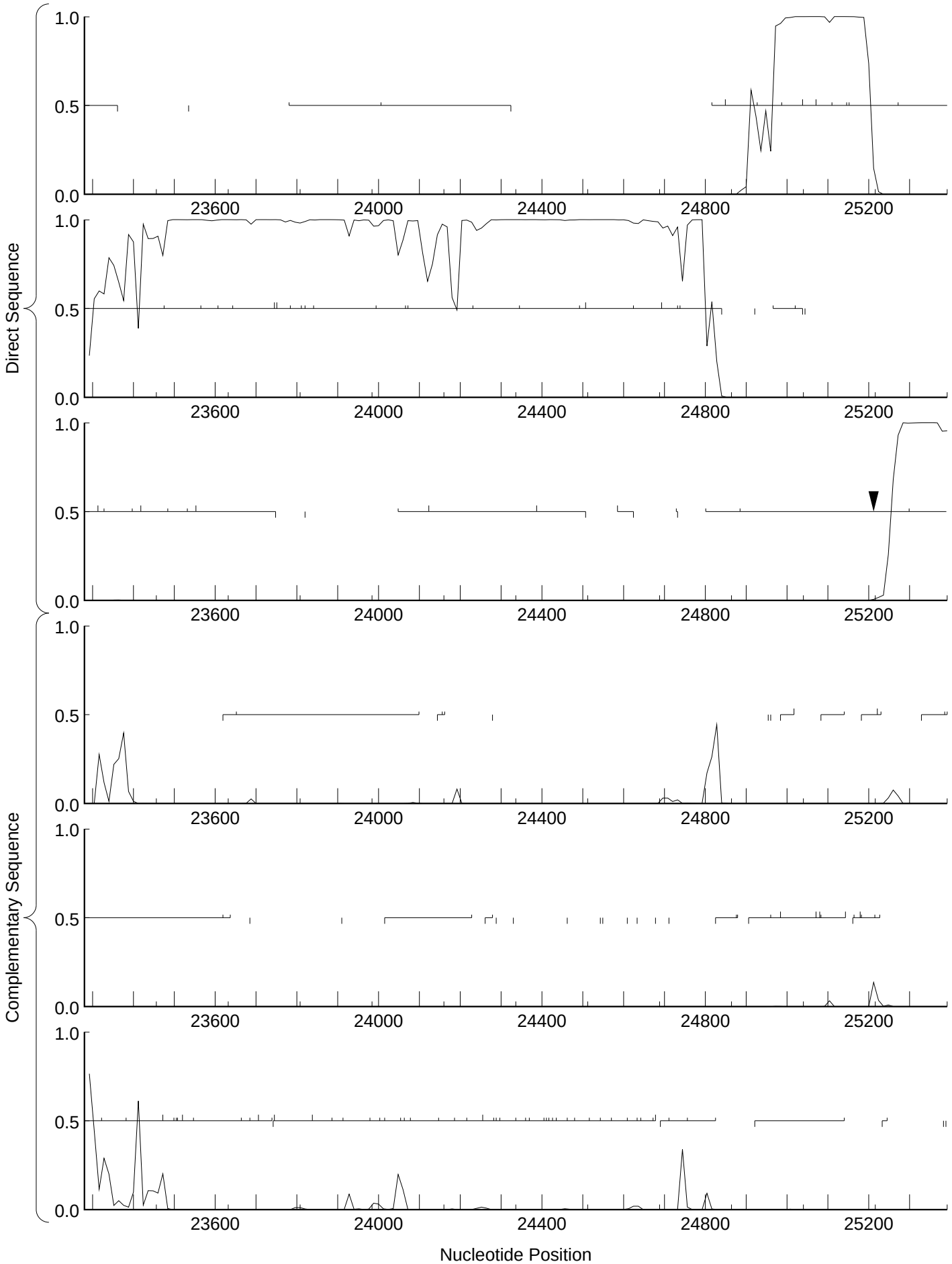
mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 11/24



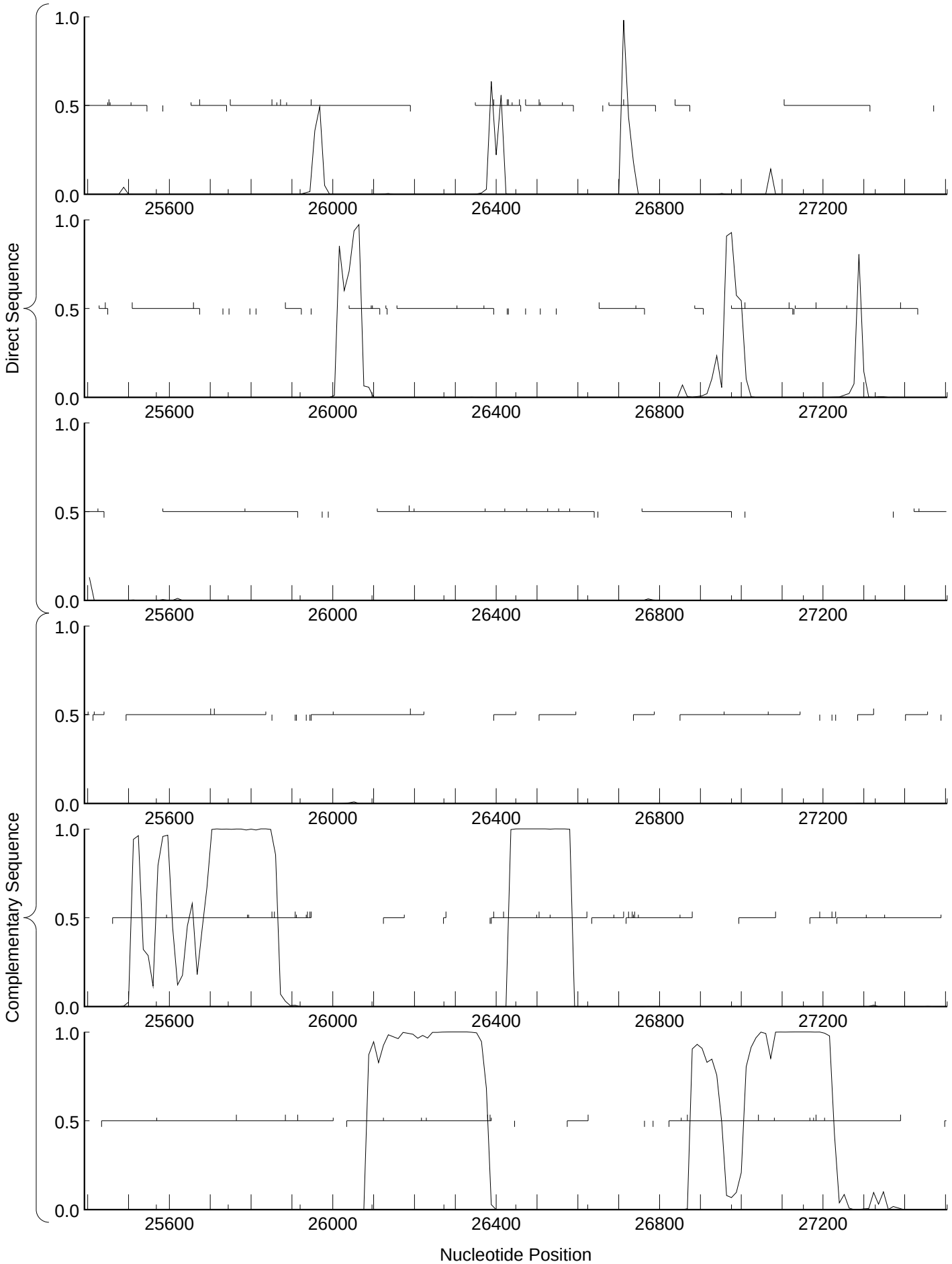
mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 12/24



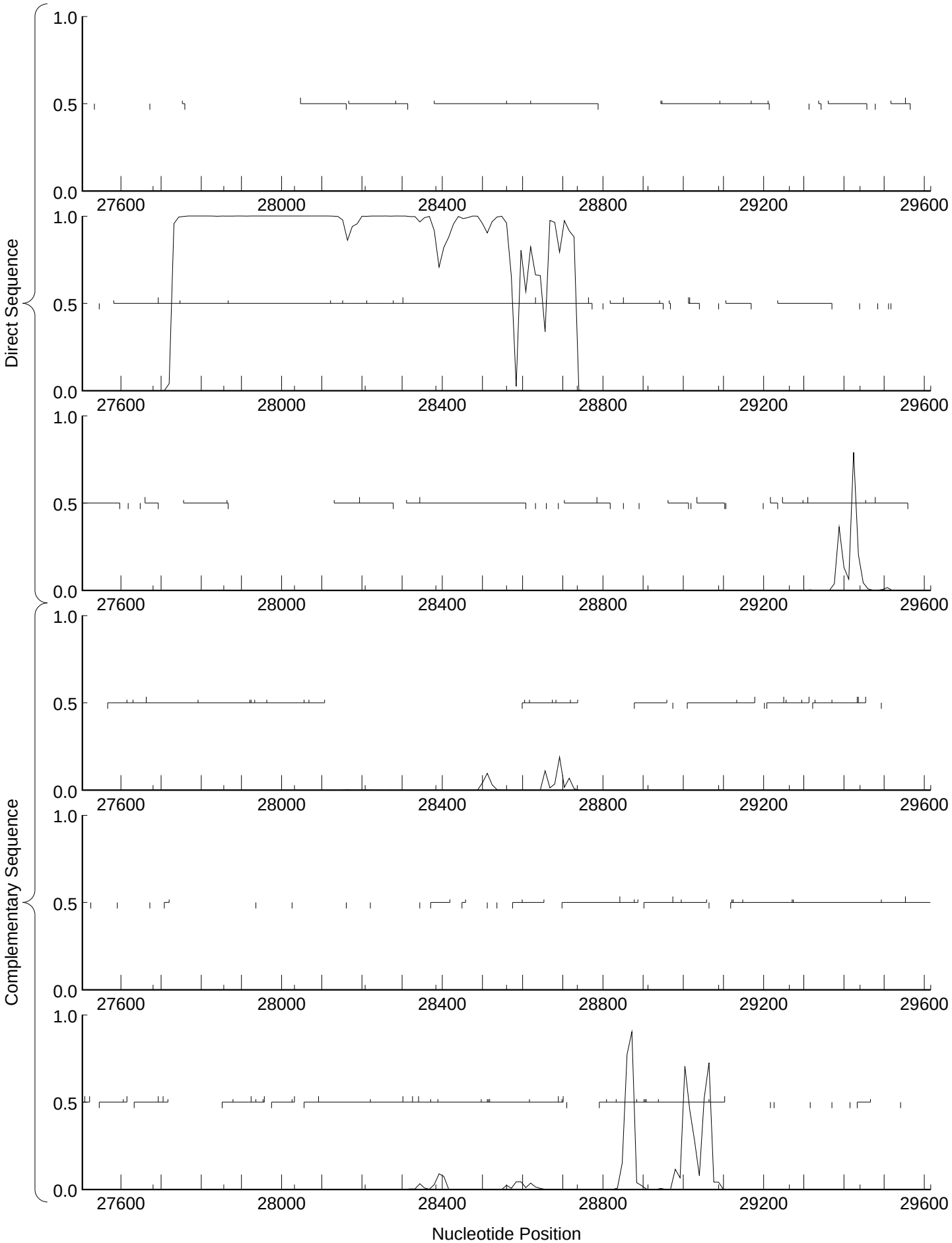
mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 13/24



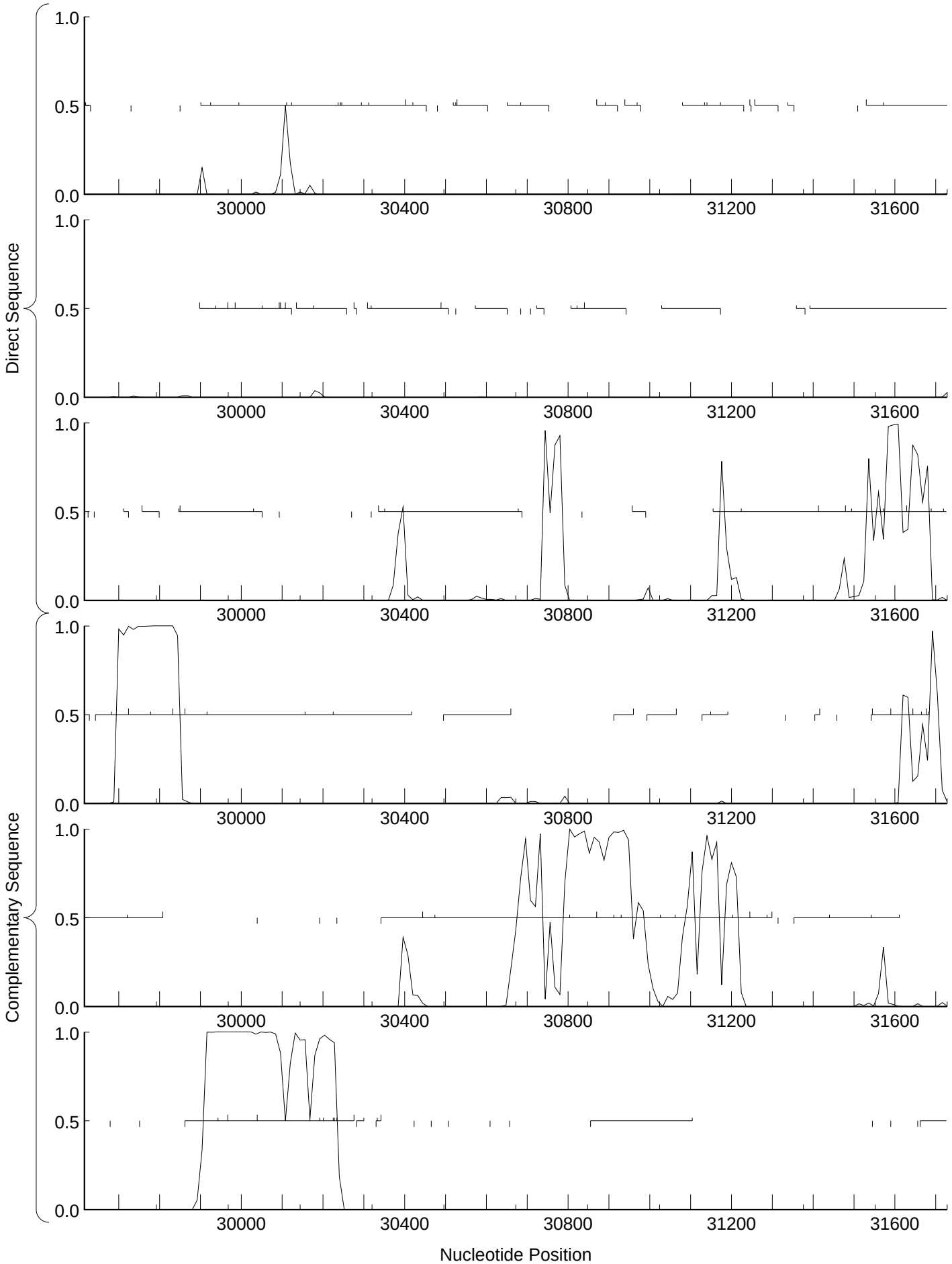
mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 14/24



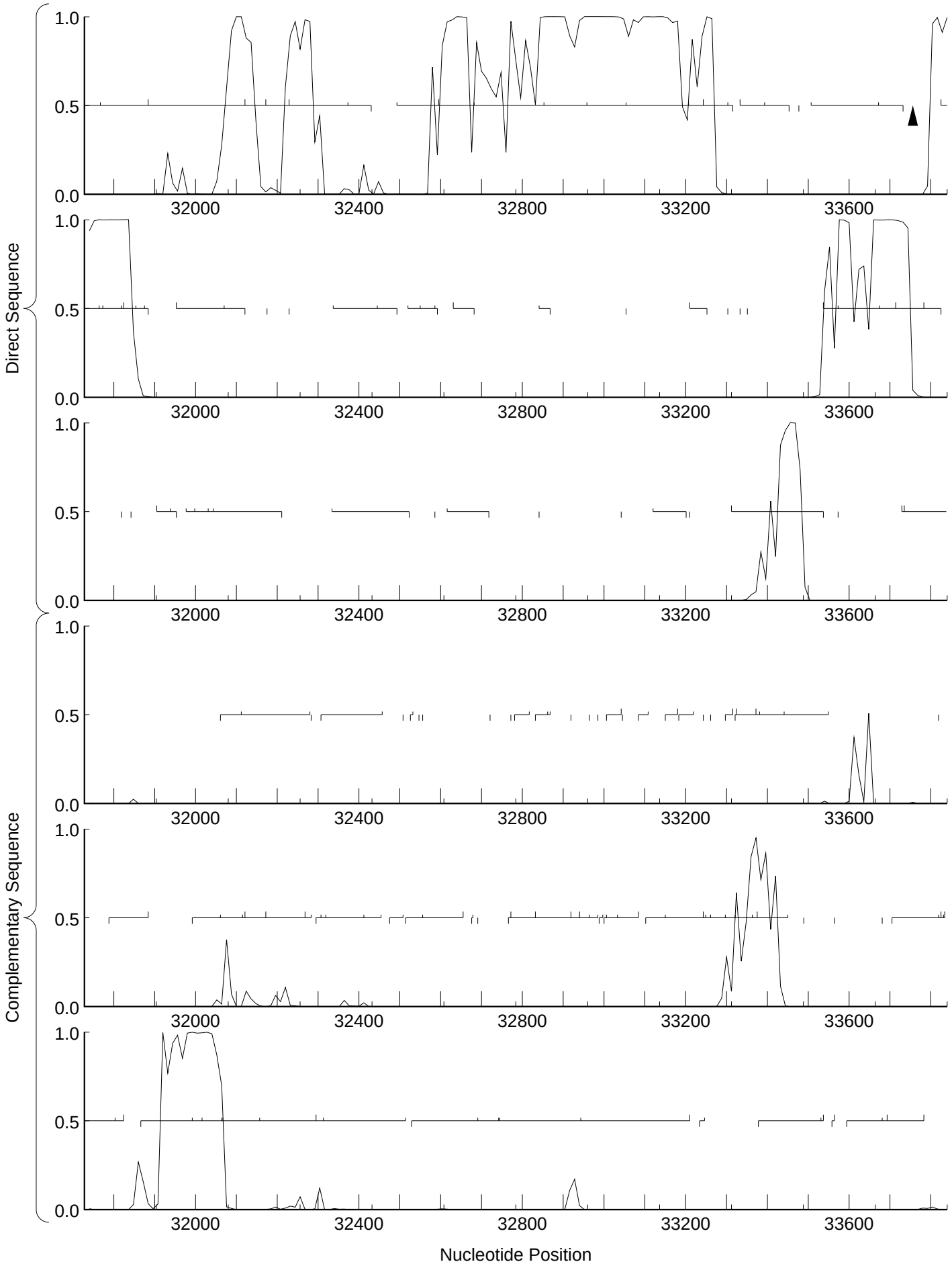
mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 15/24



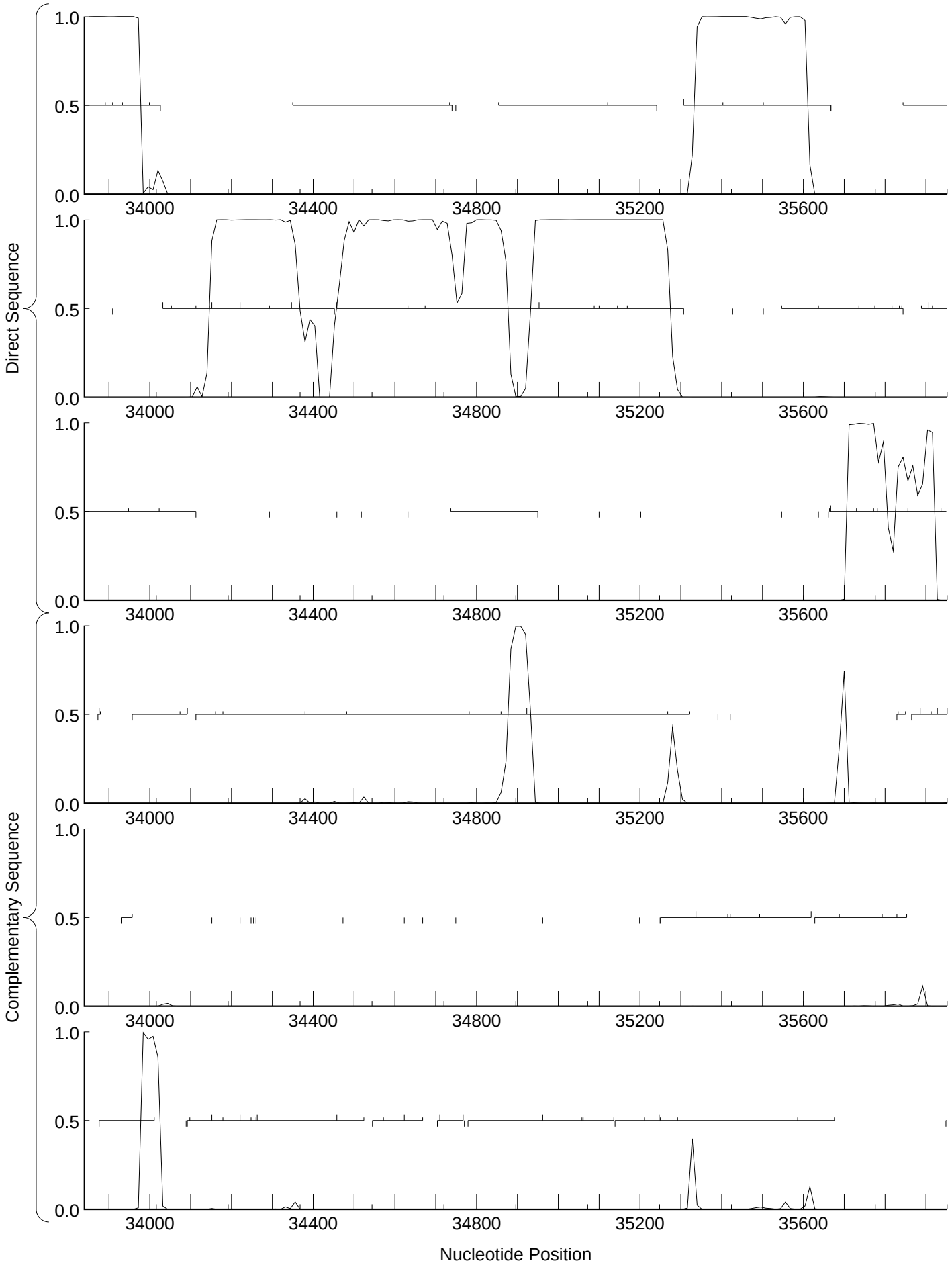
mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 16/24



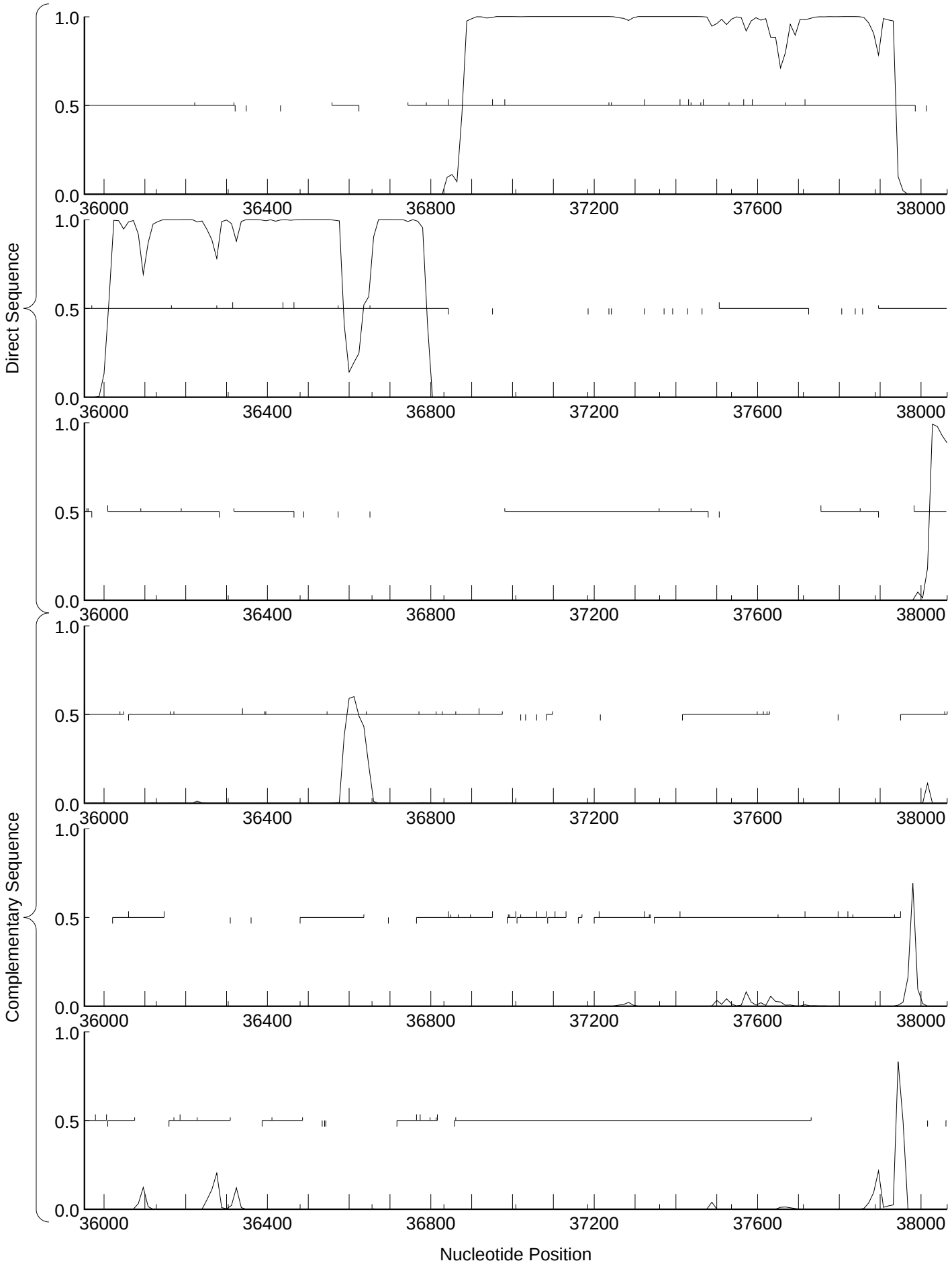
mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 17/24



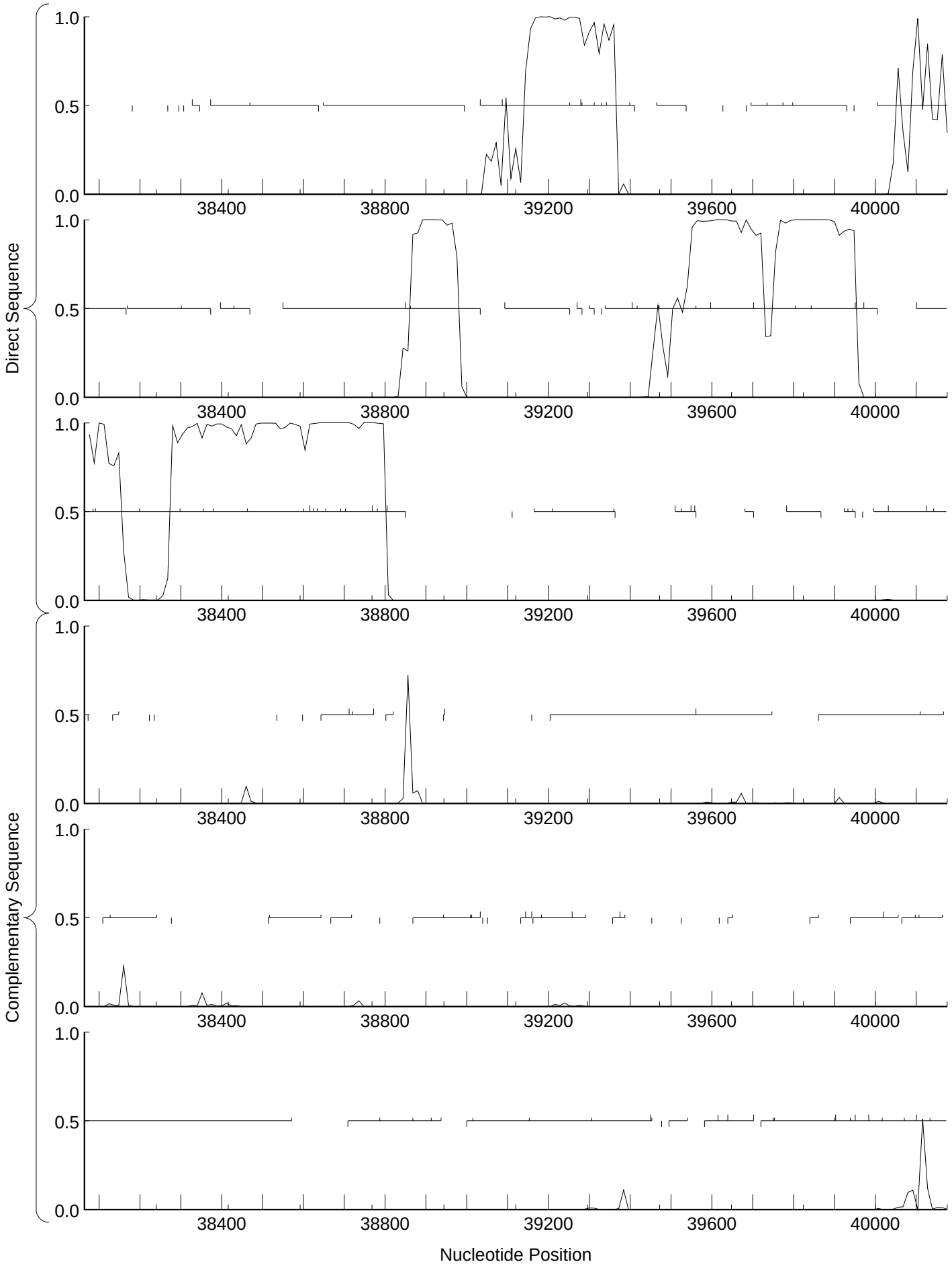
mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 18/24



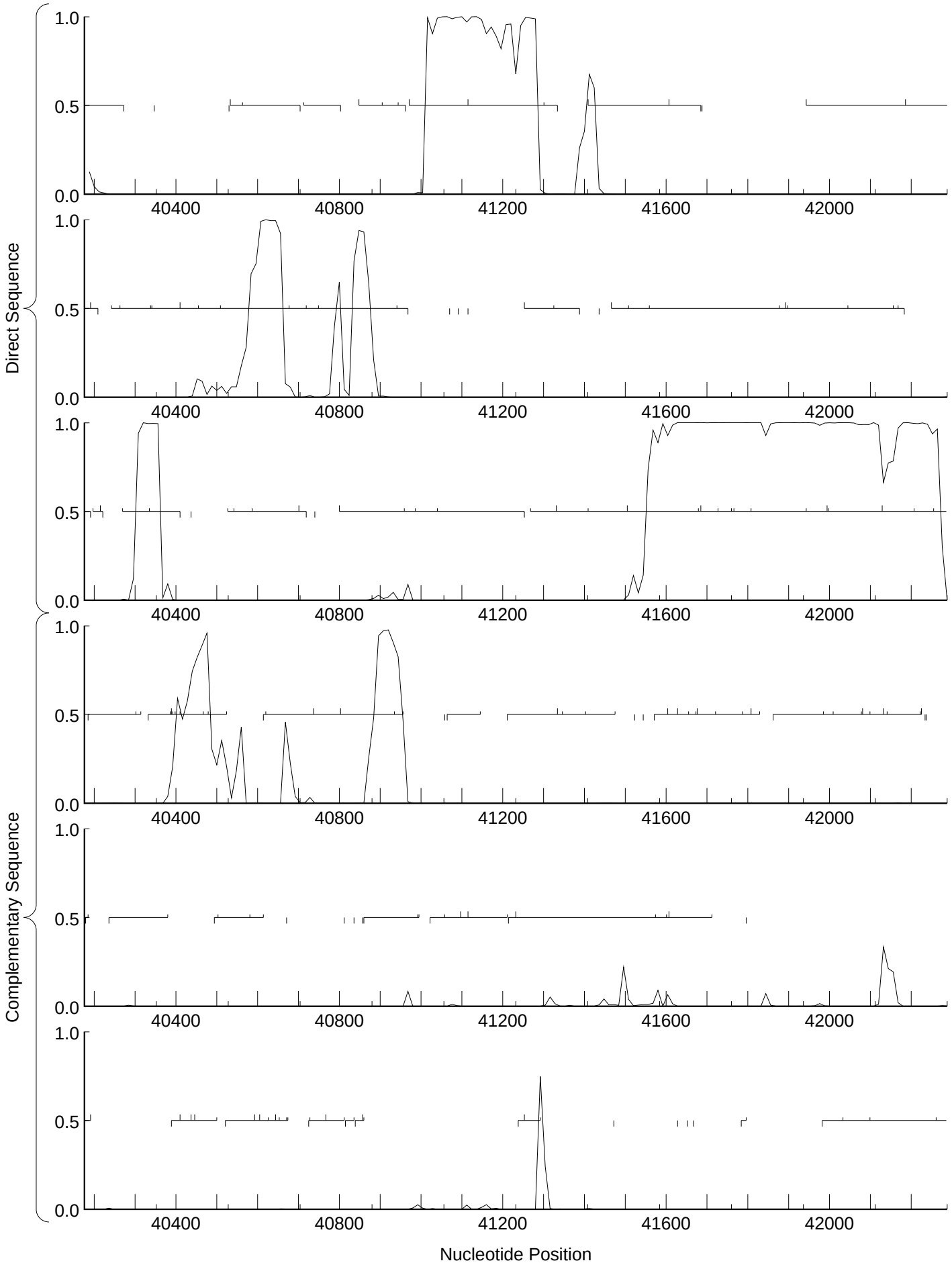
mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 19/24



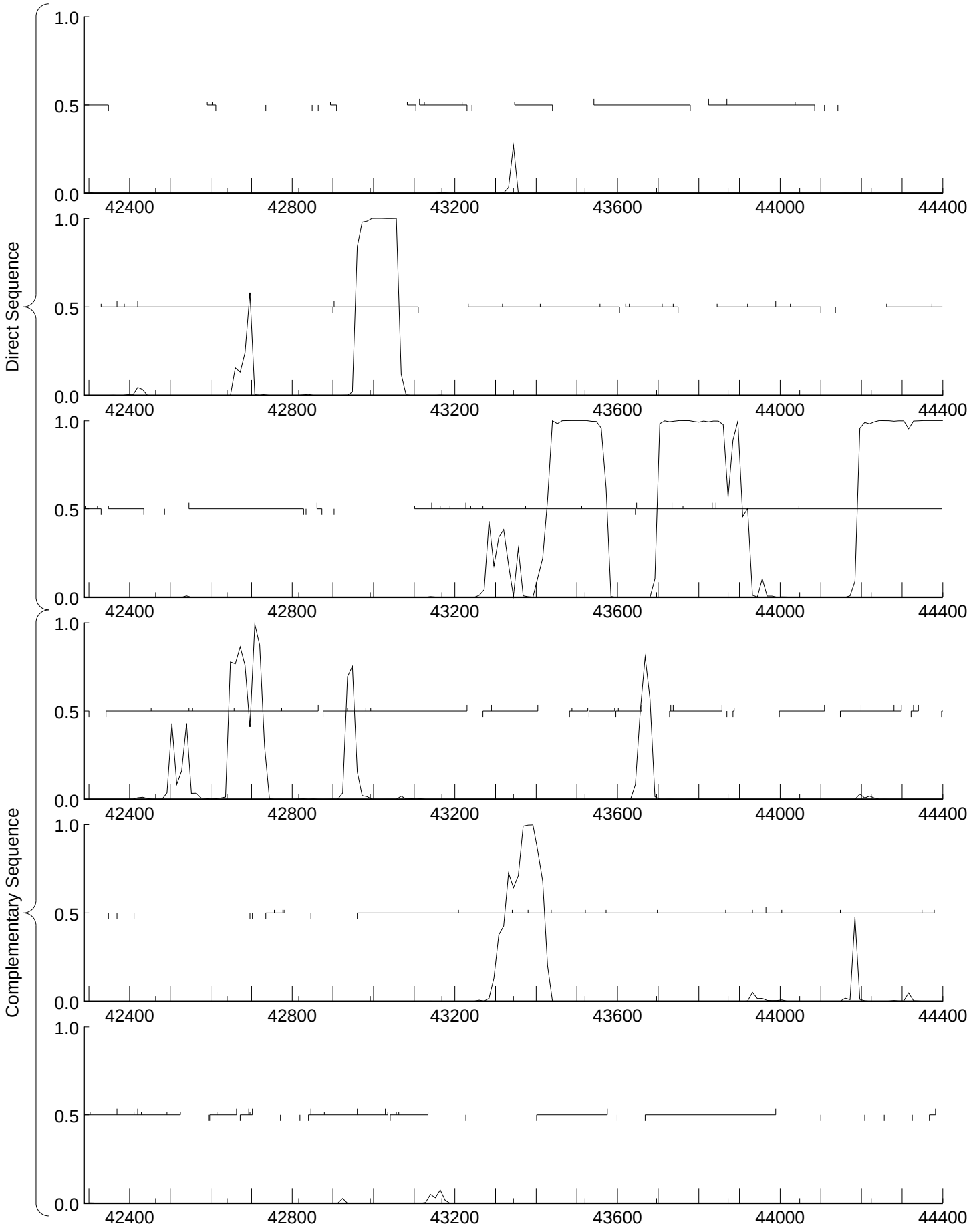
mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 20/24



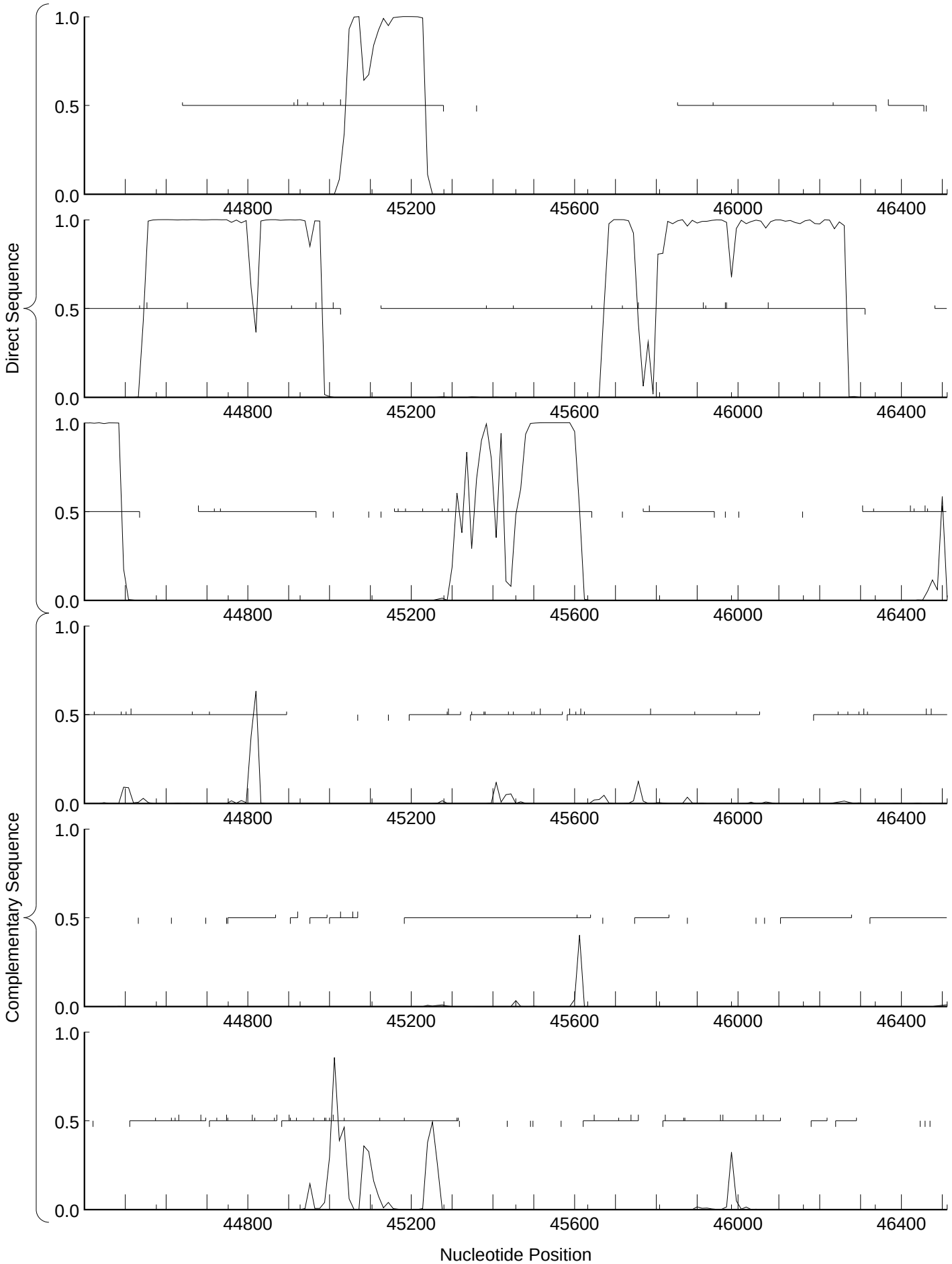
mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 21/24



mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 22/24



mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 23/24



mplete sequence, 47880 bp including 11-base 3’ overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 24/24

