GeneMark

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

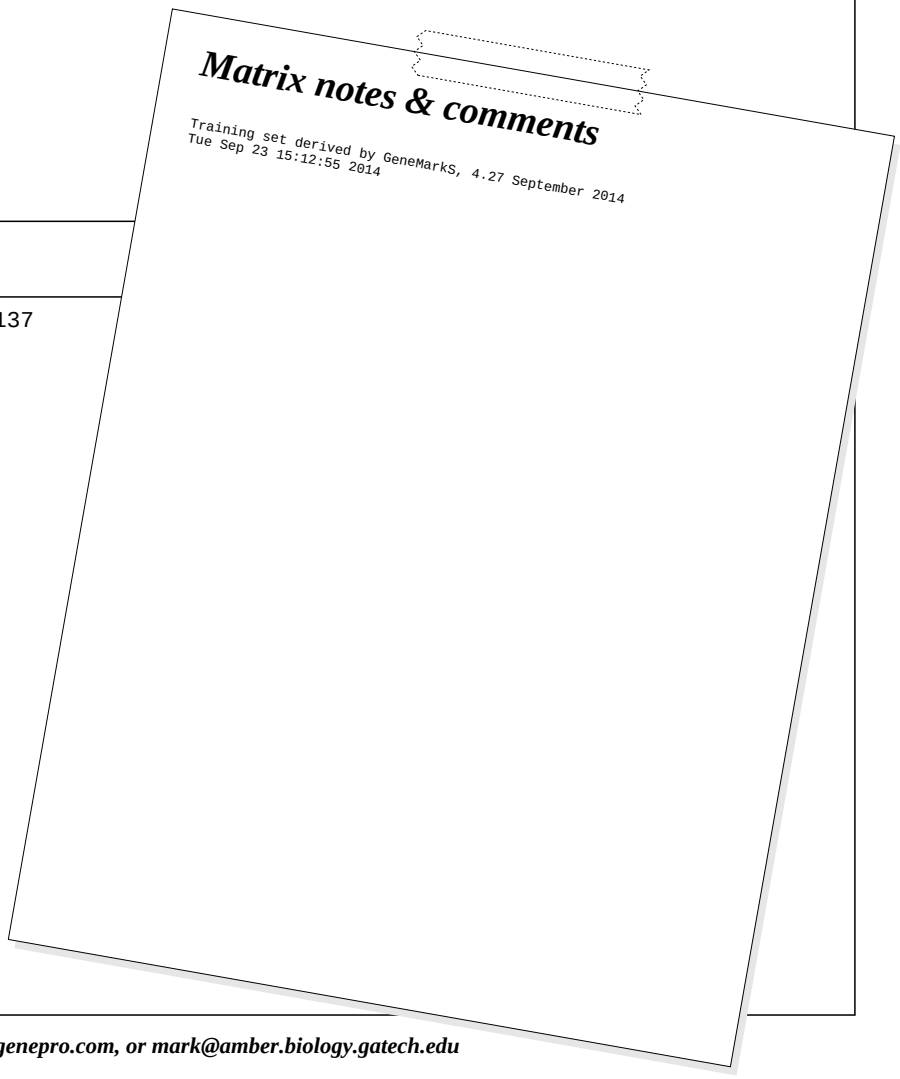
PROGRAM INFORMATION

```
Sequence         :  Propionibacterium phage Keiki complete sequence, 29339 bp including 11-base 3’ overhang (TC
Analysis Date    :  3/28/24 at 20:34:55
Pages            :  15
Sequence Length  :  29339 bp
GC Content       :   54.03%

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

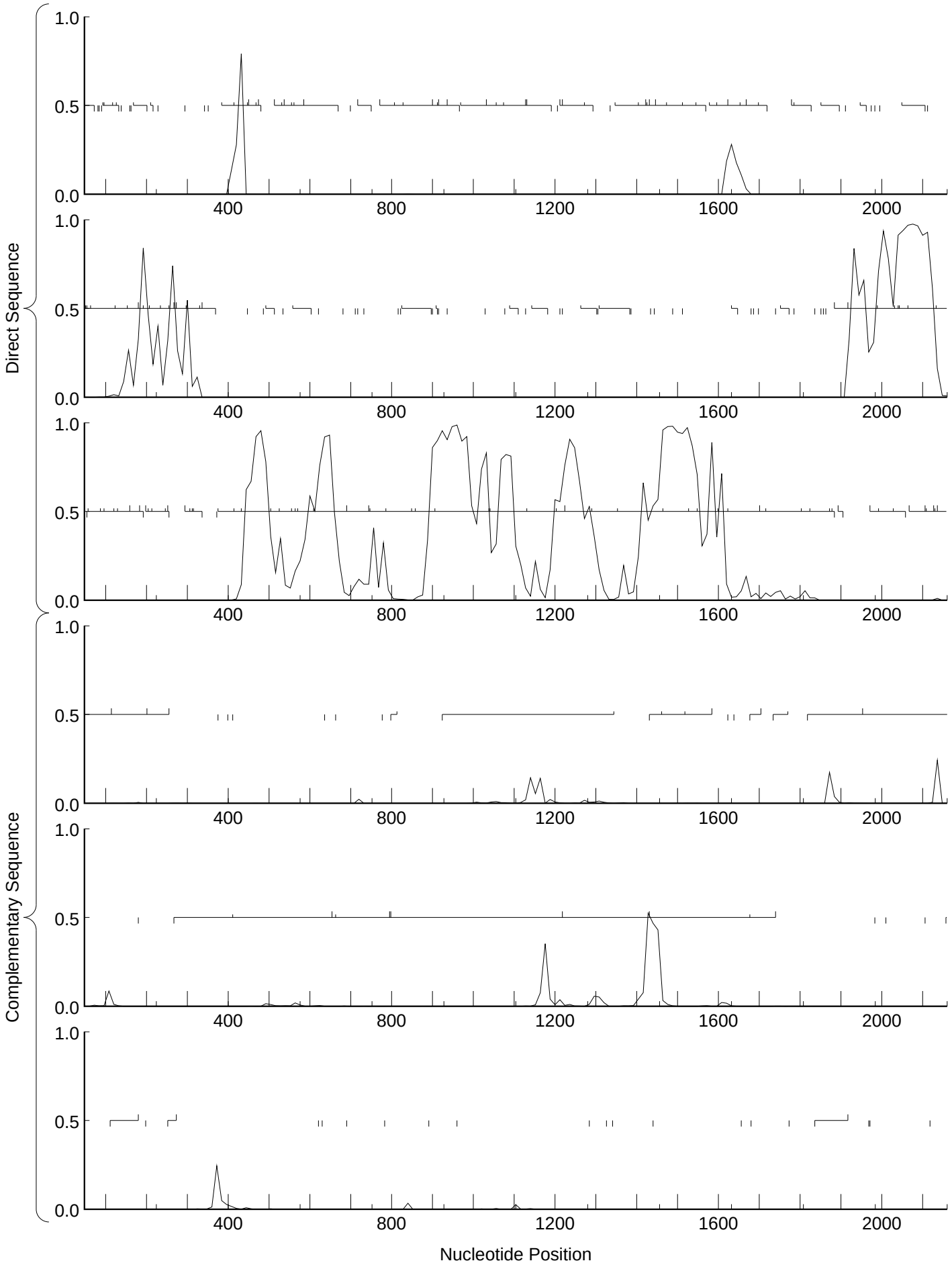
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Matrix     :  Propionibacterium_acnes_SK137
Author     :  -

Order      :  4
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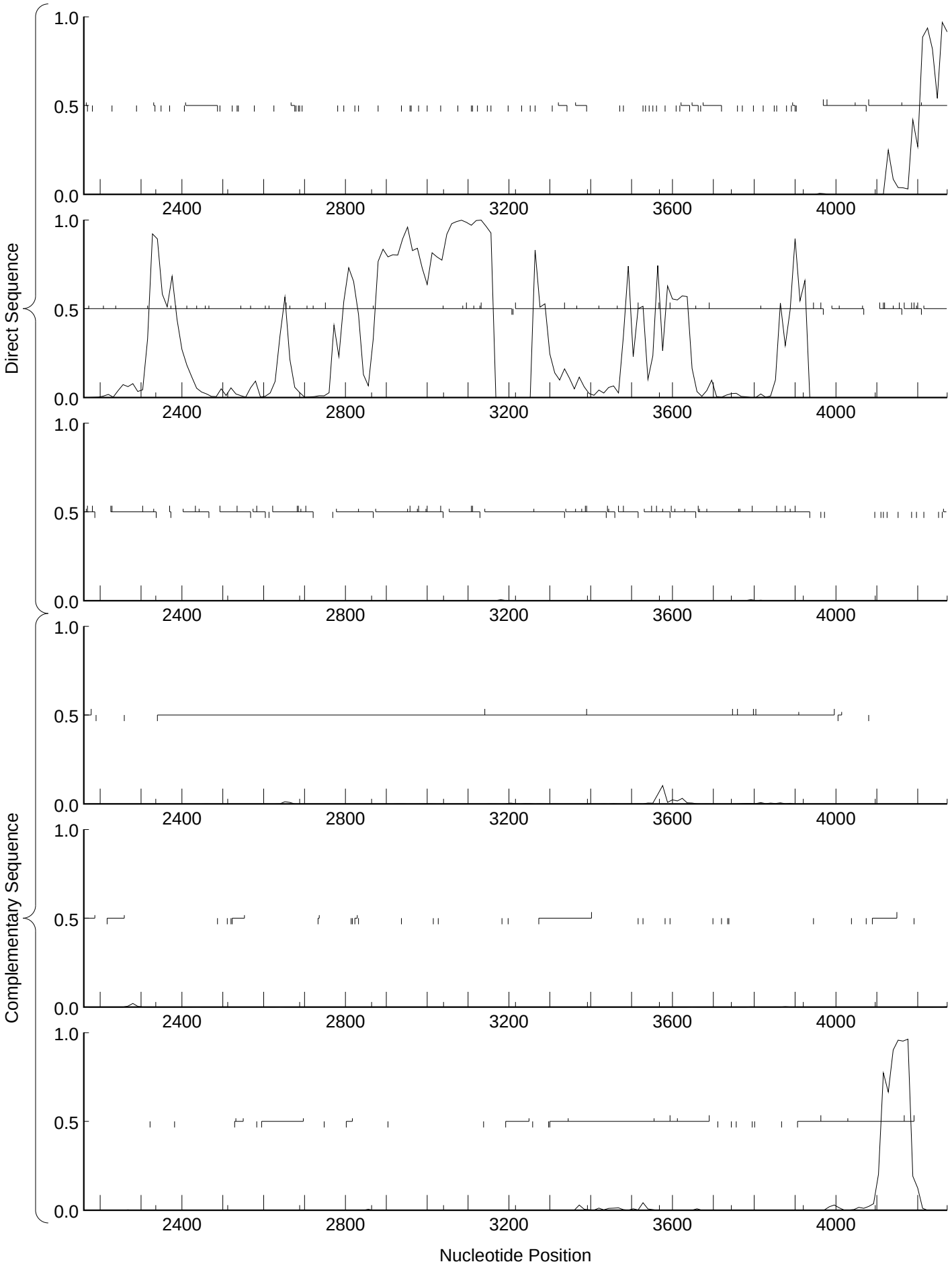

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

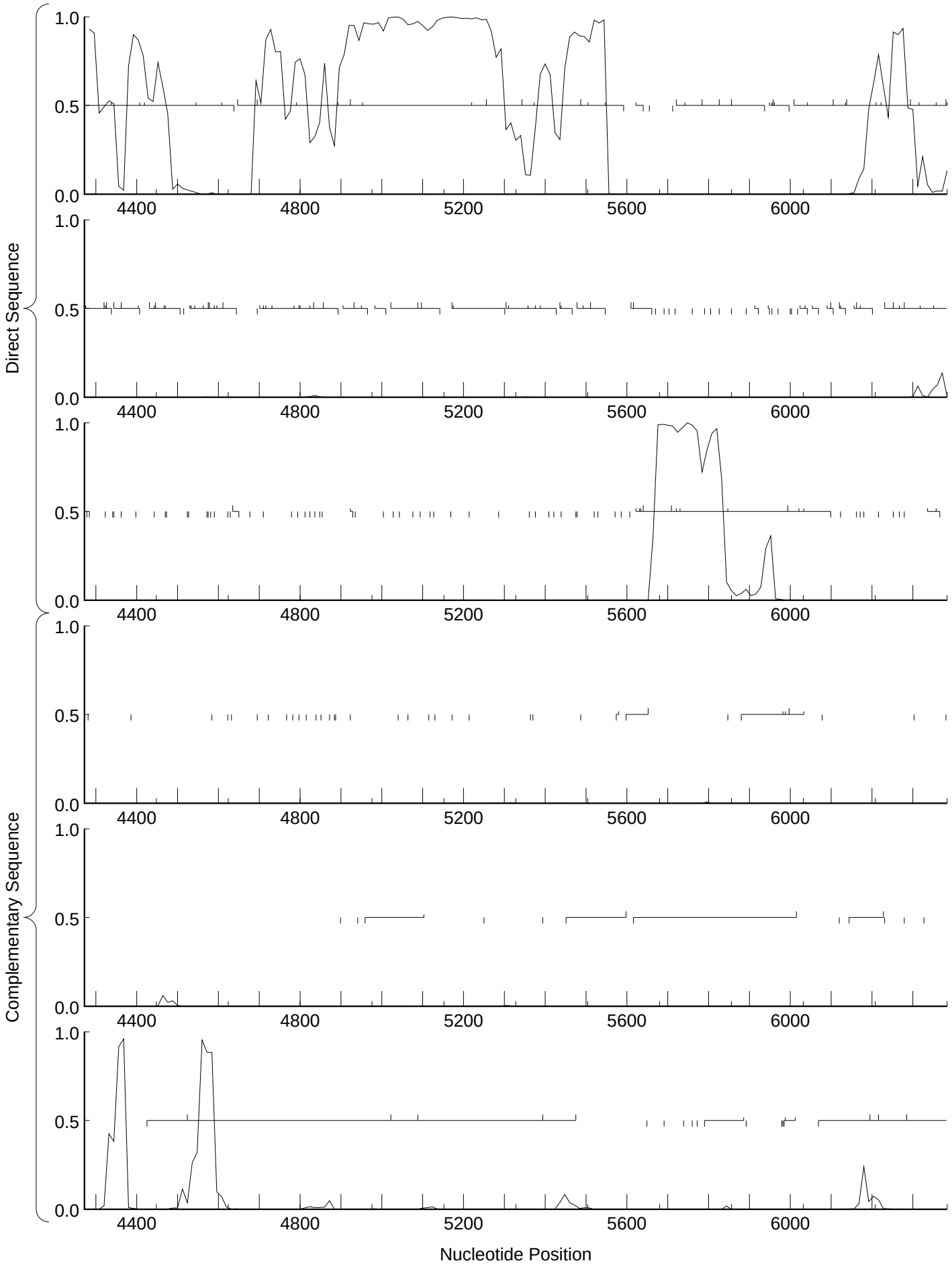
complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 2/15



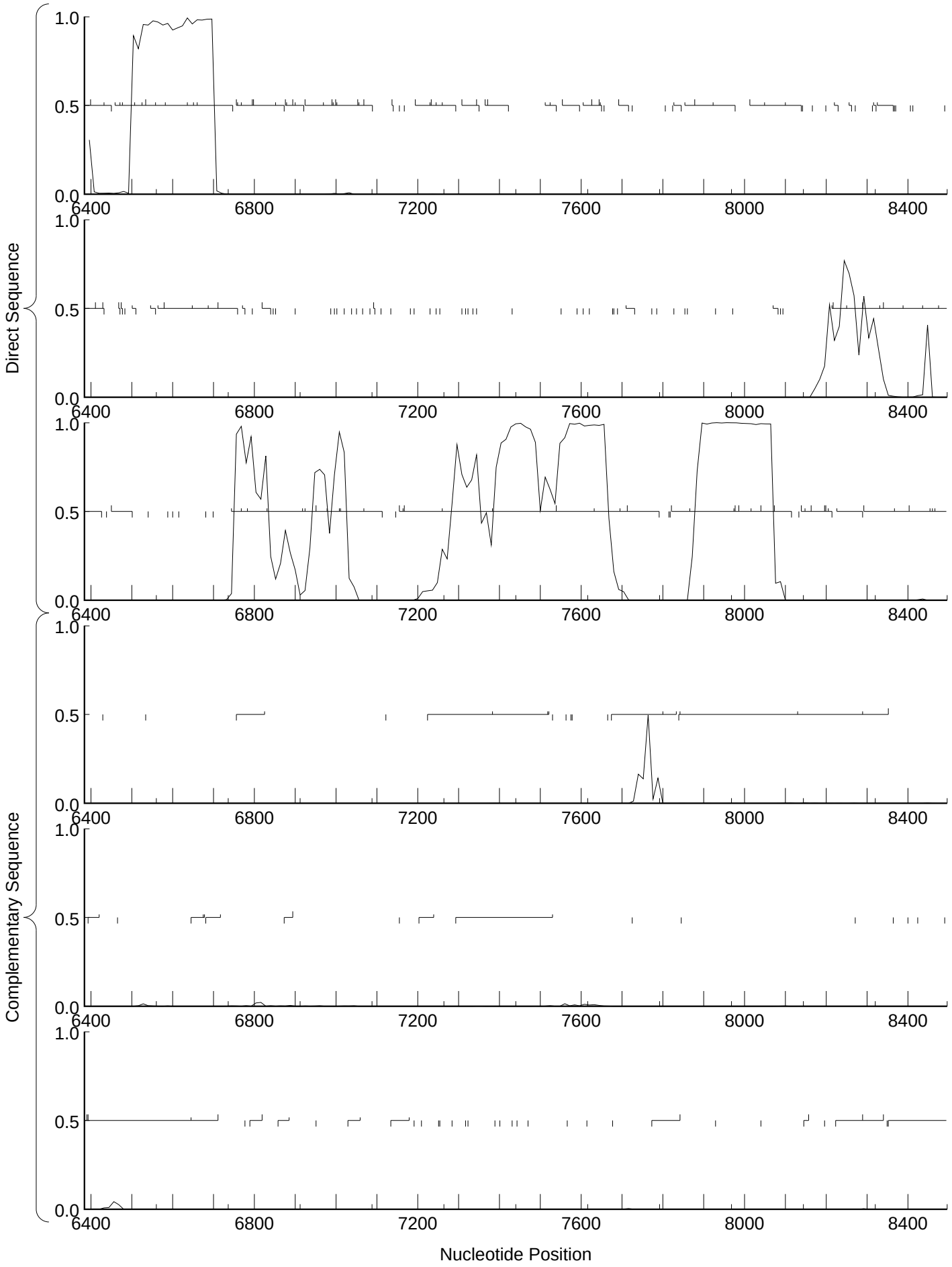
complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 3/15



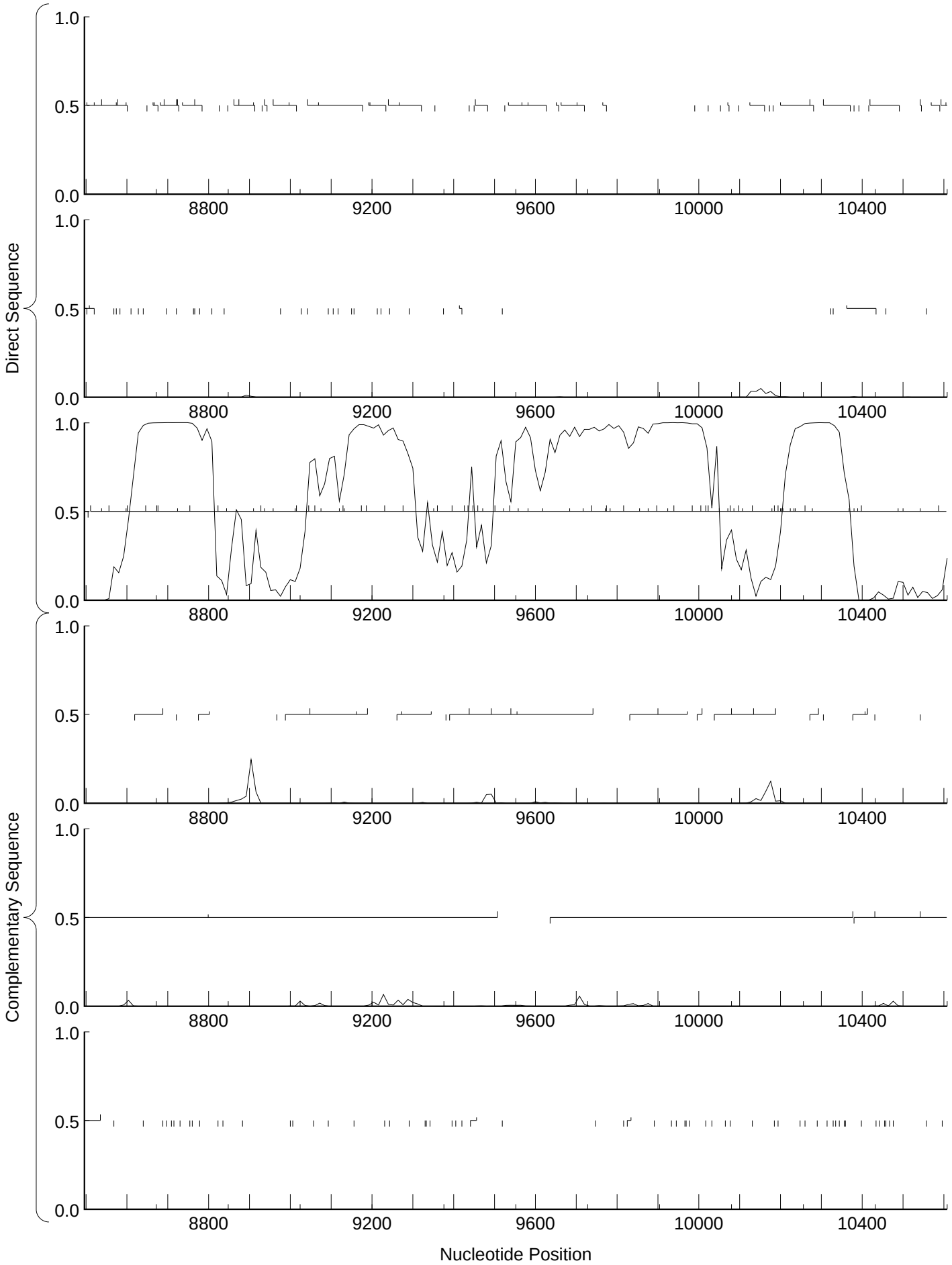
complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 4/15



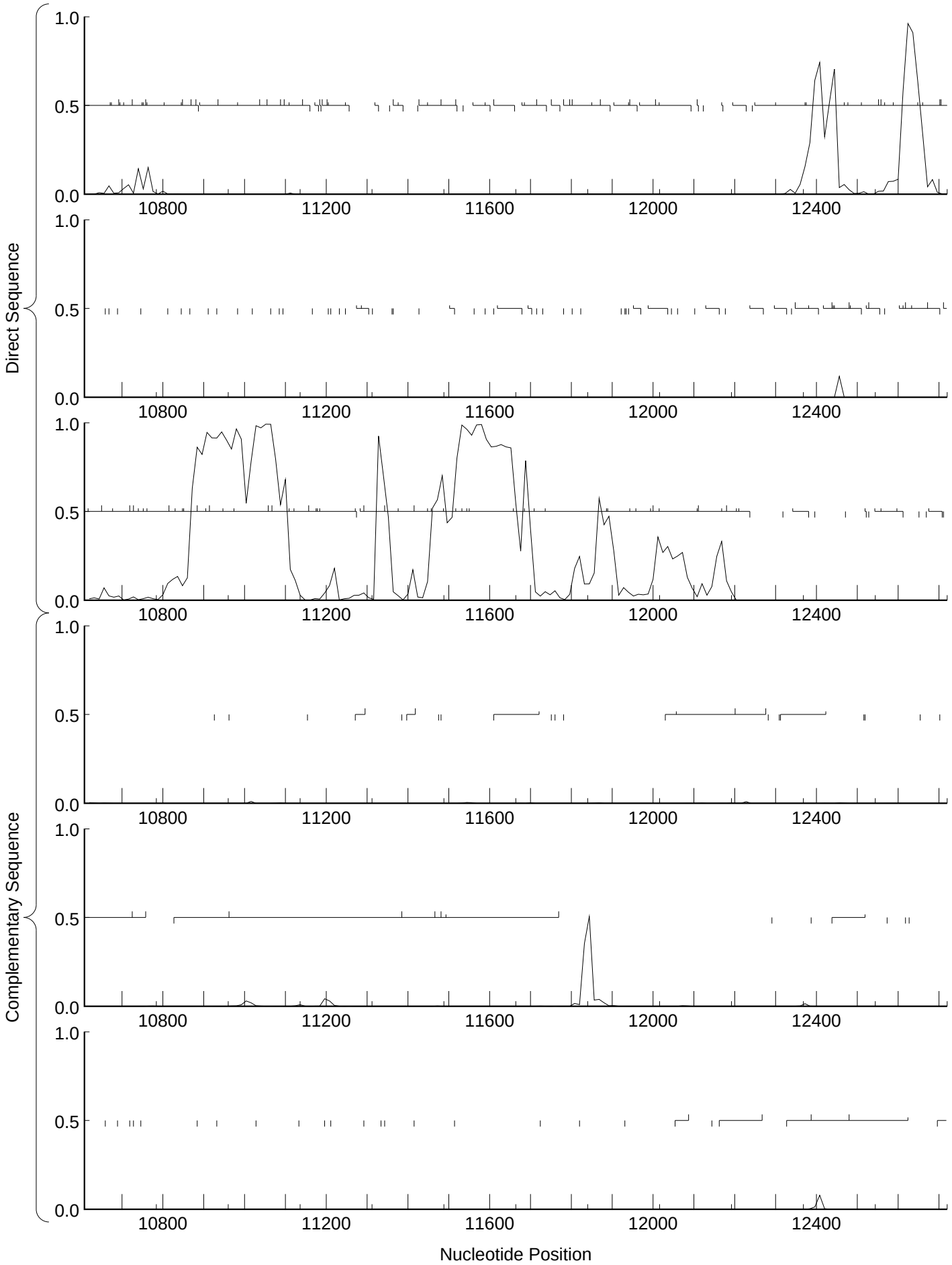
complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 5/15



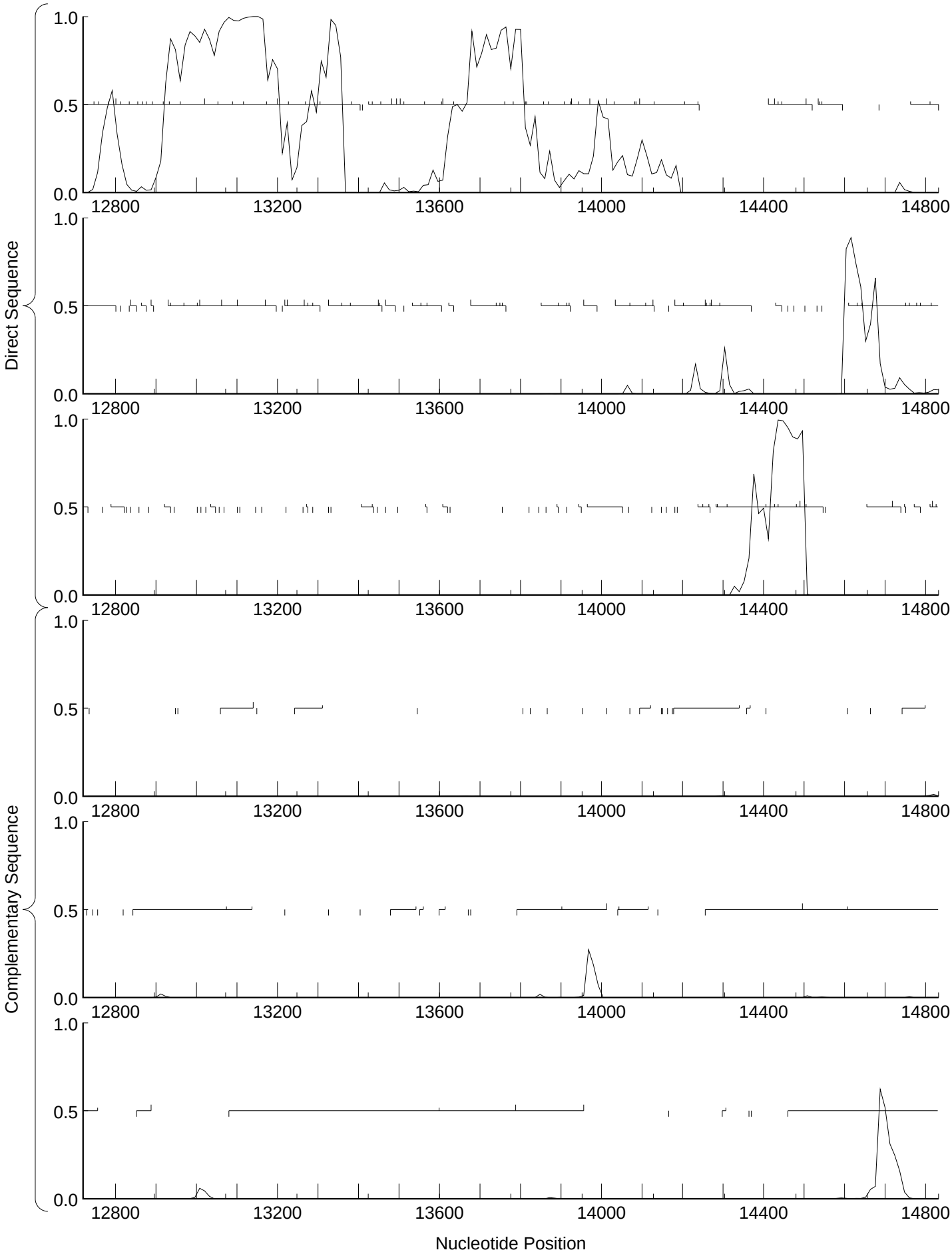
complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 6/15



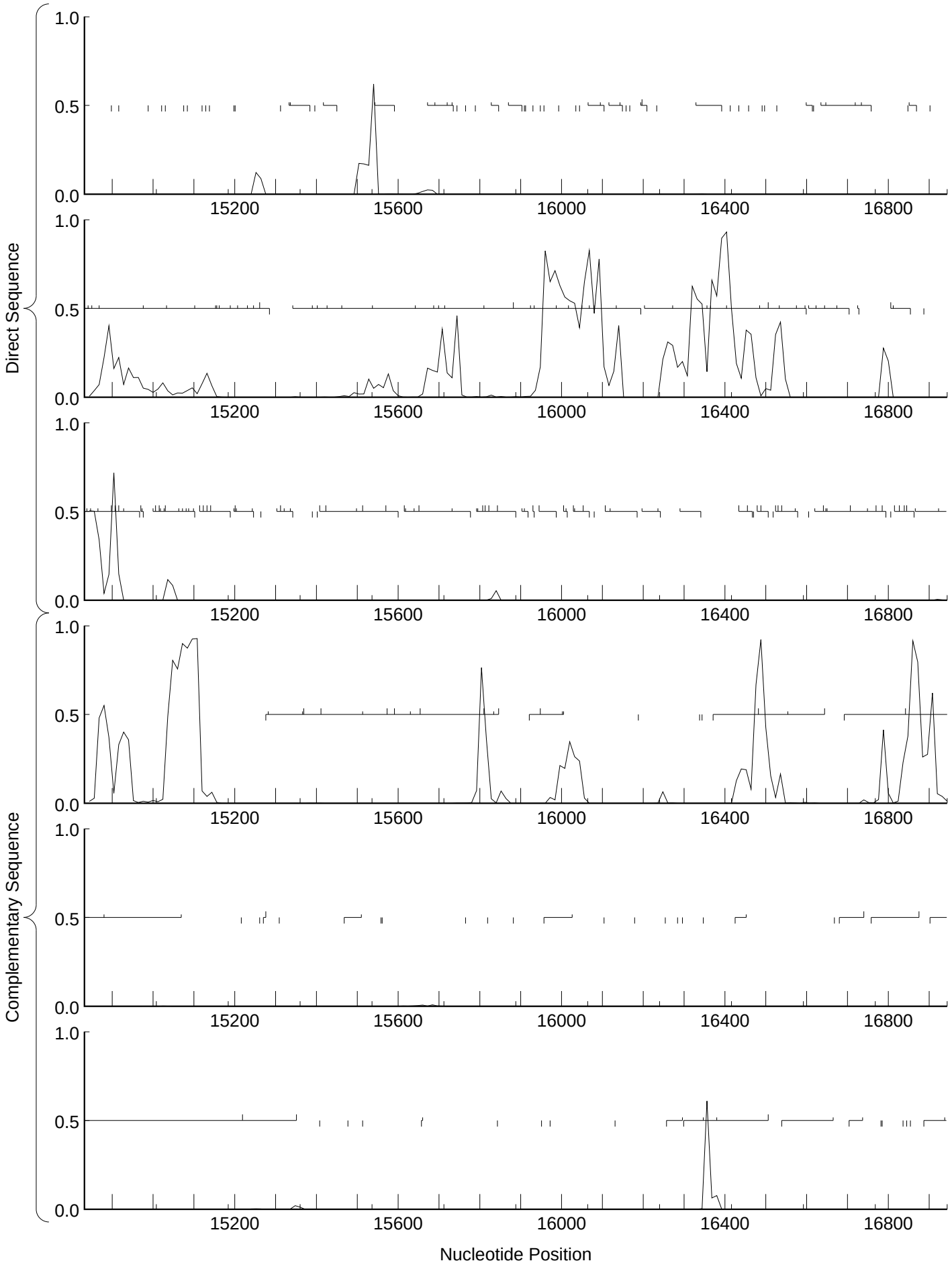
complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 7/15



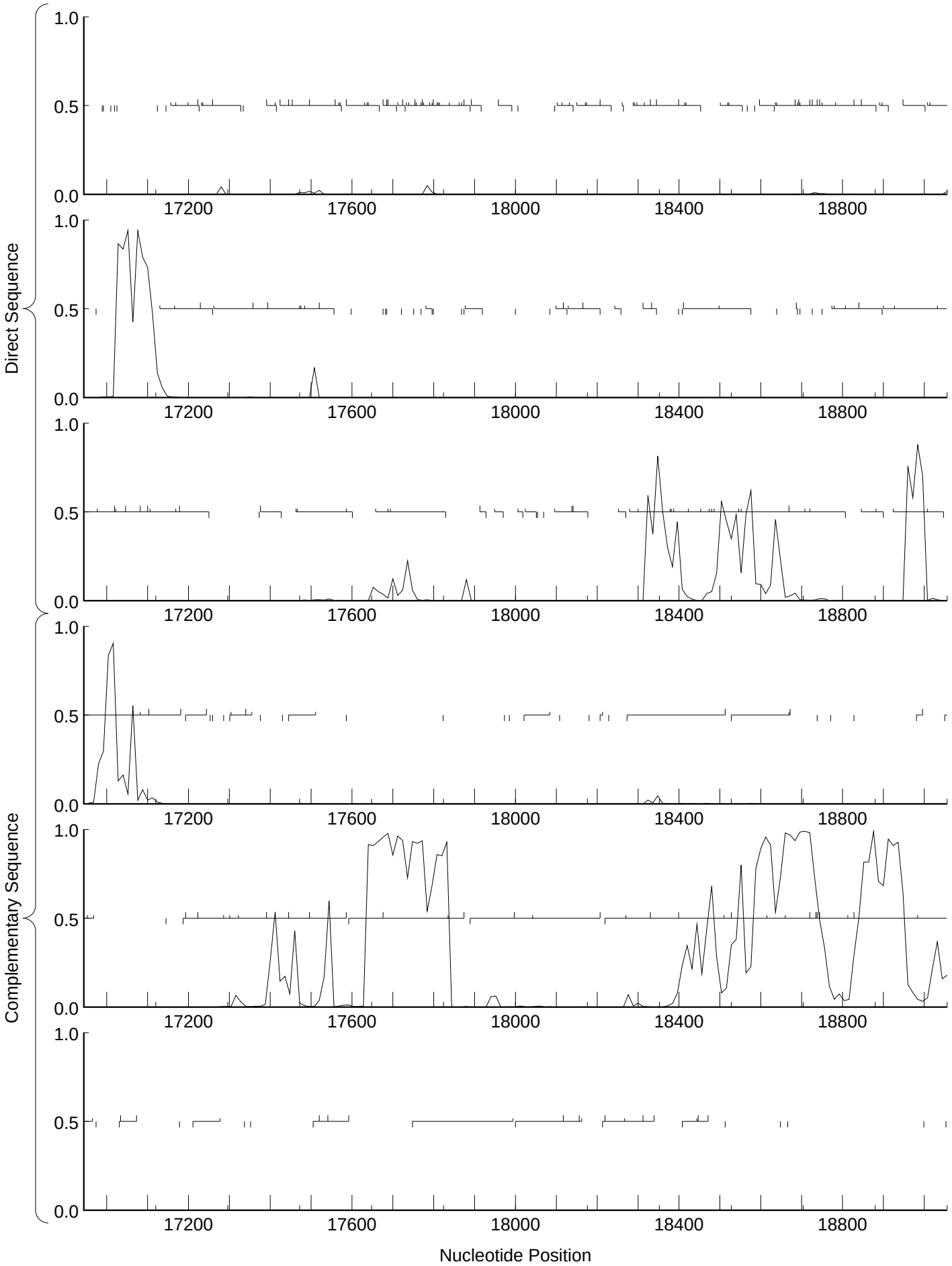
complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 8/15



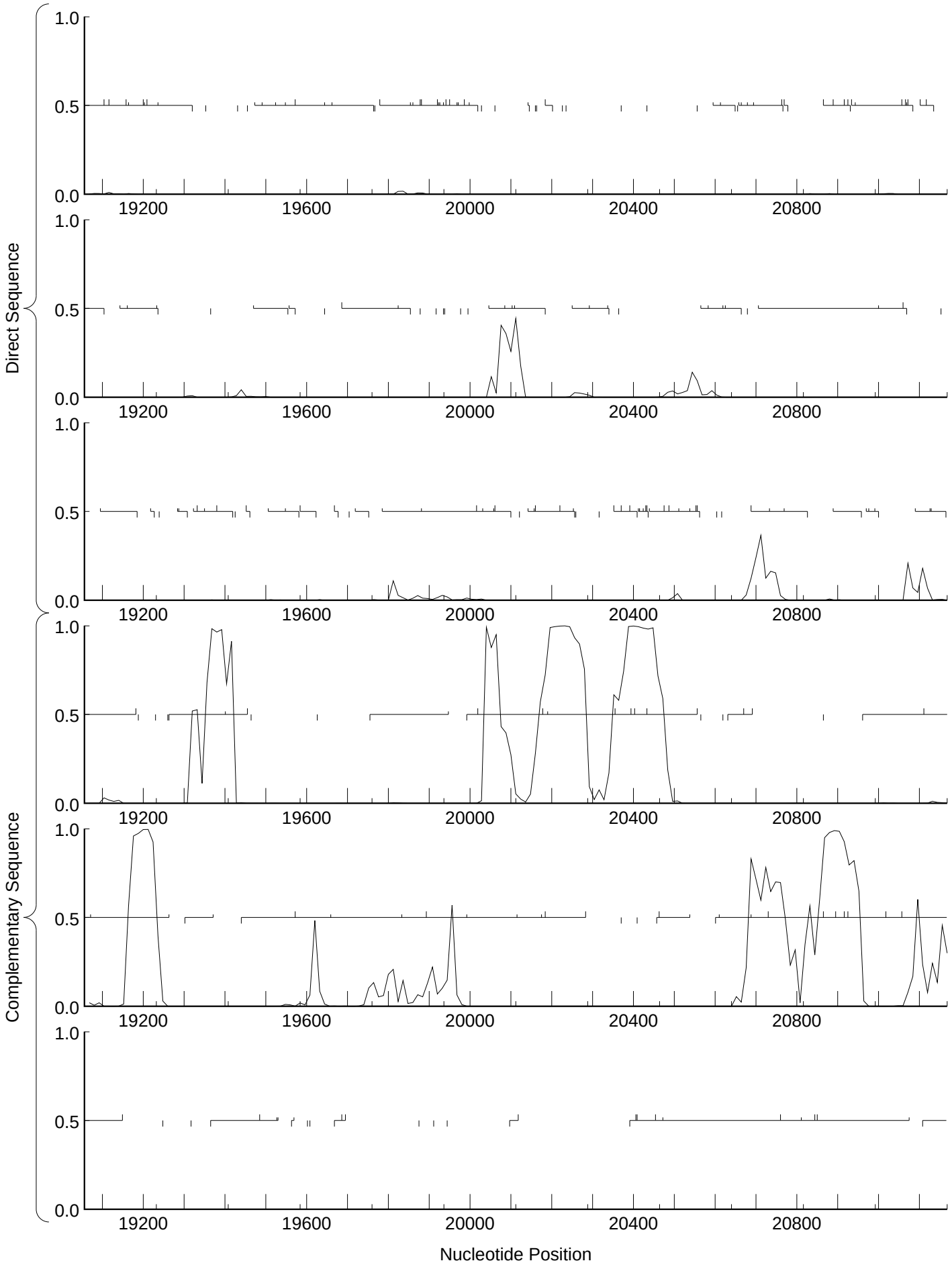
complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 9/15



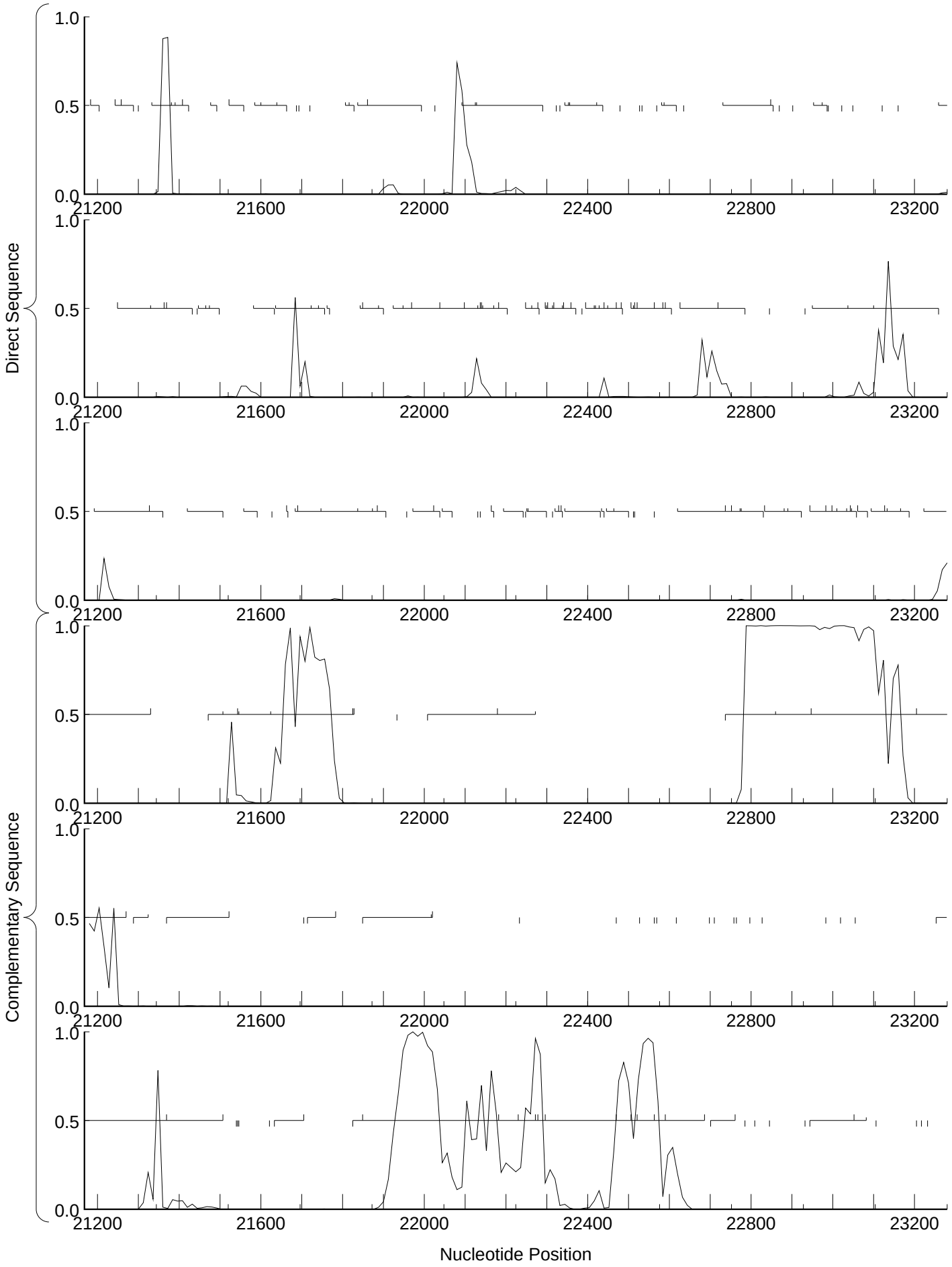
complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 10/15



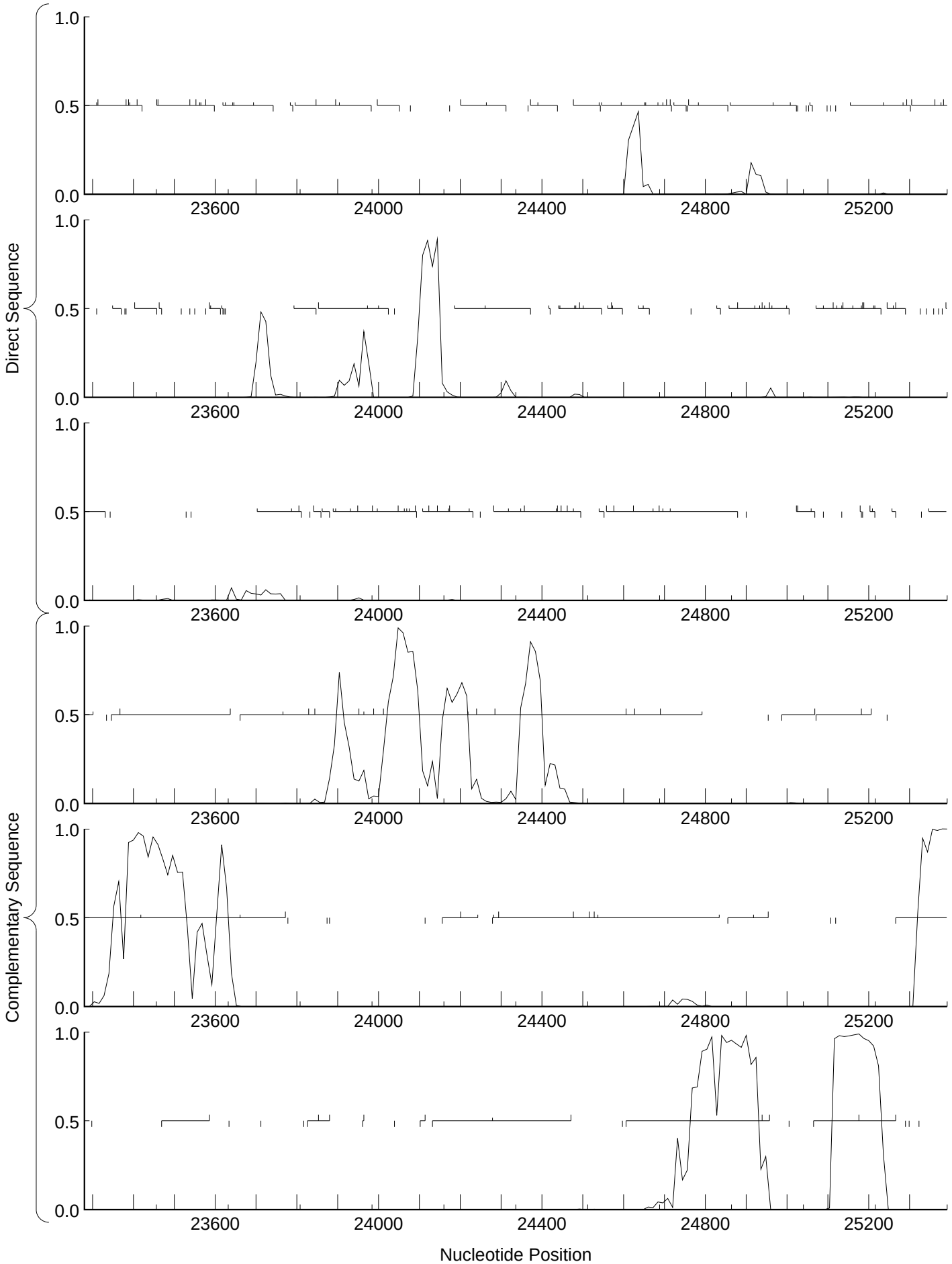
complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 11/15

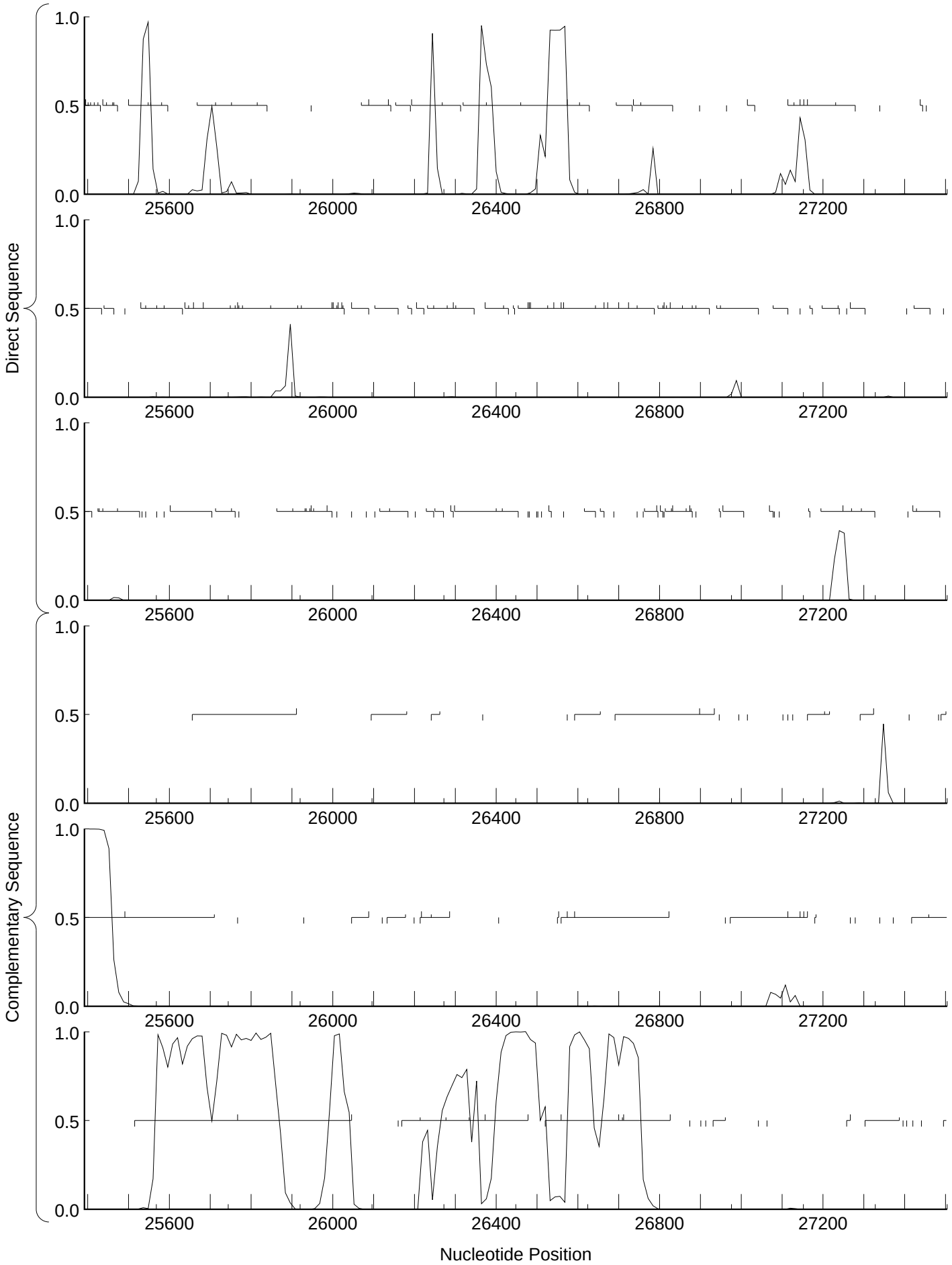


complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 12/15



complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 13/15




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
25600
26000
26400
26800
27200
Nucleotide Position

complete sequence, 29339 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 15/15

