GeneMark

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

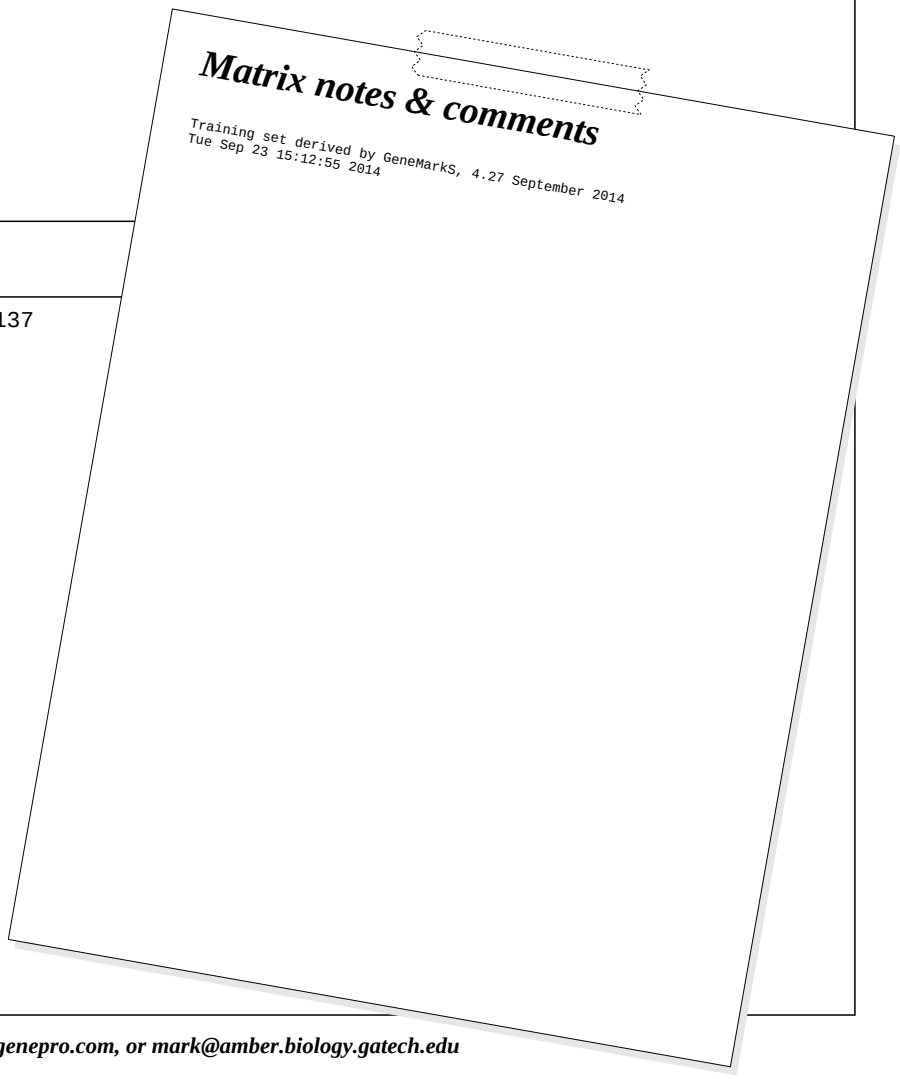
PROGRAM INFORMATION

```
Sequence         :  Propionibacterium phage P1.1 complete genome, 29348 bp including 11-base 3' overhang (TCGTA
Analysis Date    :  3/28/24 at 22:20:01
Pages            :  15
Sequence Length  :  29348 bp
GC Content       :   54.36%

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

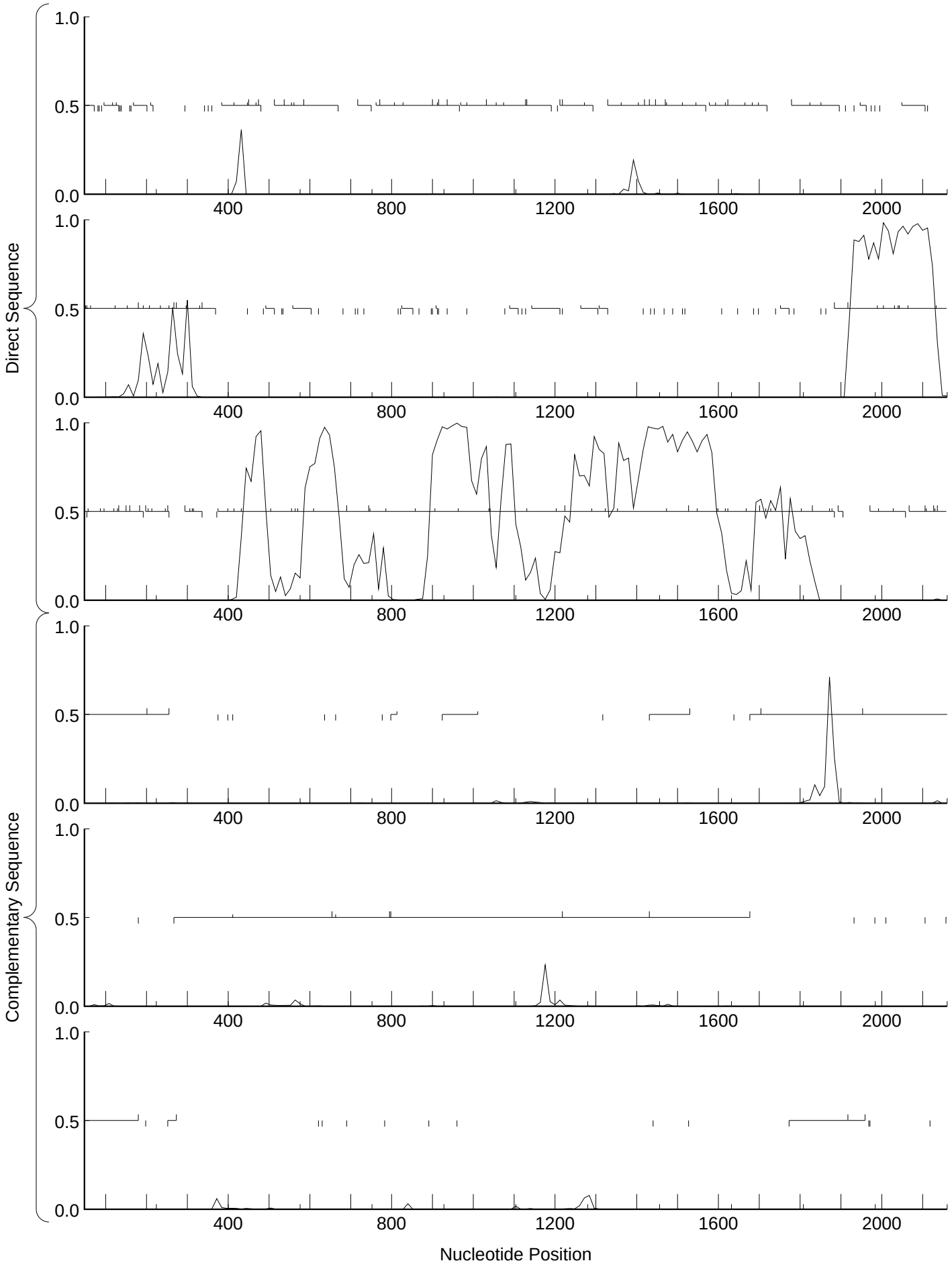
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Matrix     :  Propionibacterium_acnes_SK137
Author     :  -

Order      :  4
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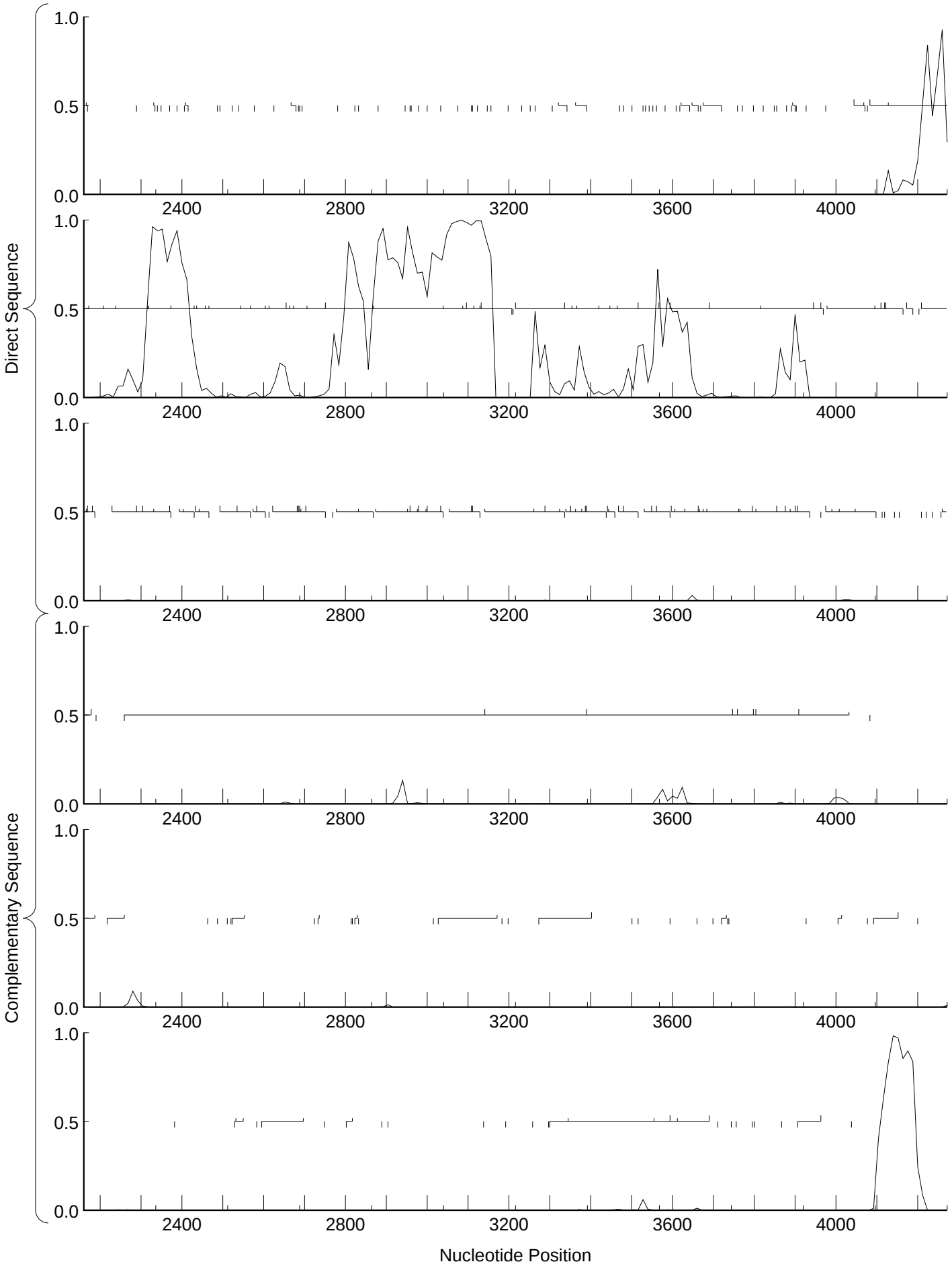

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

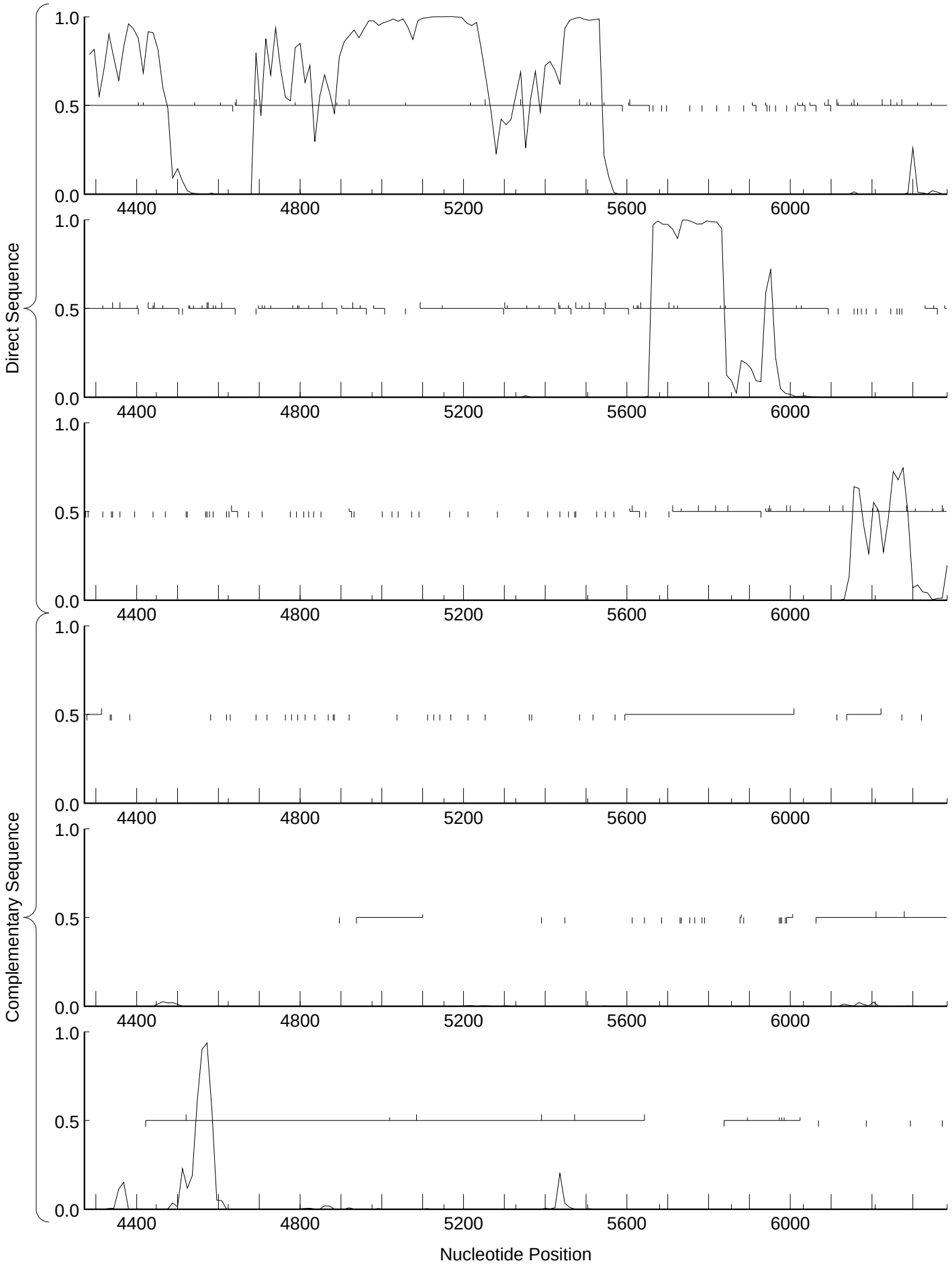
1 complete genome, 29348 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 2/15



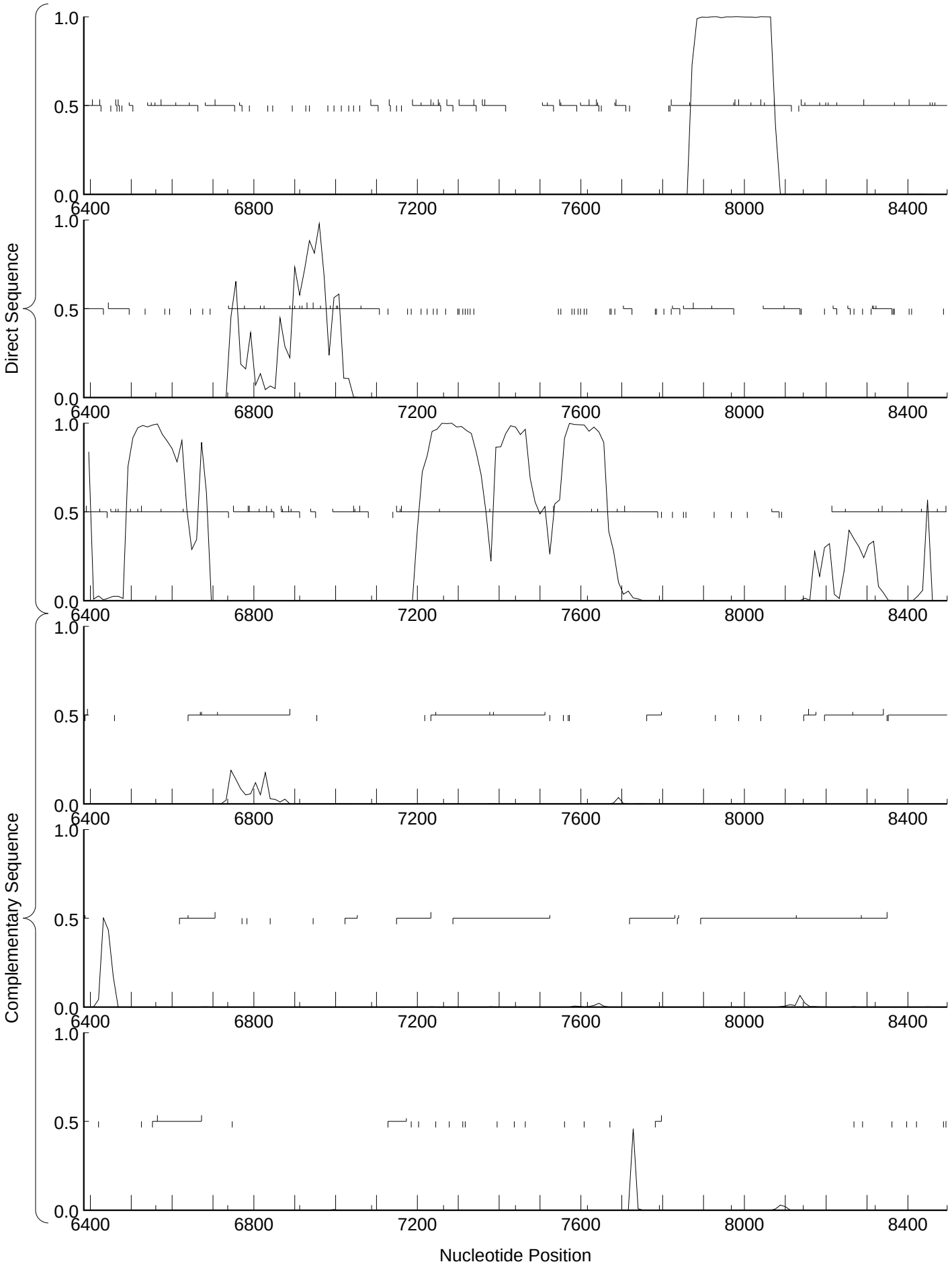
1 complete genome, 29348 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 3/15



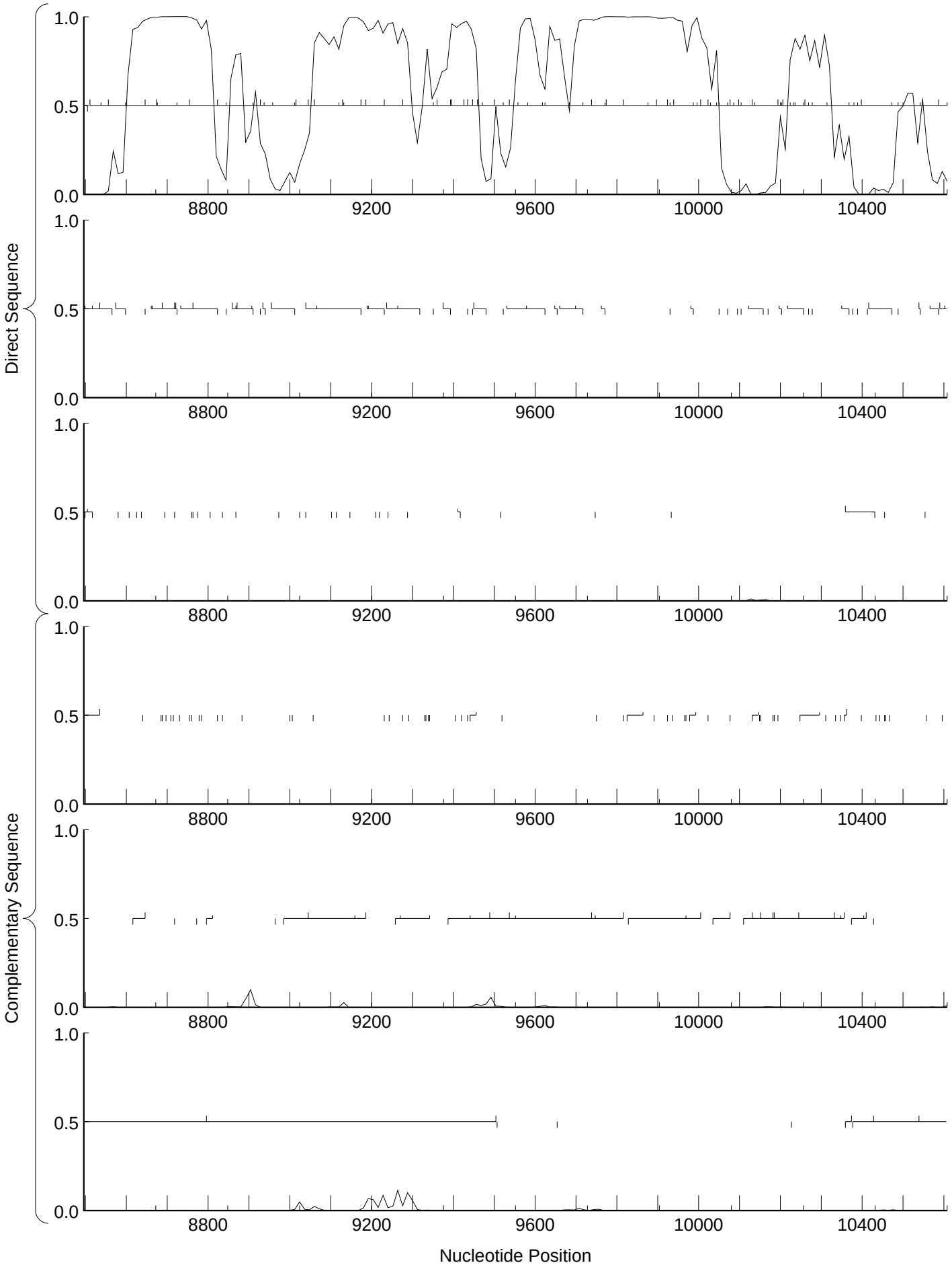
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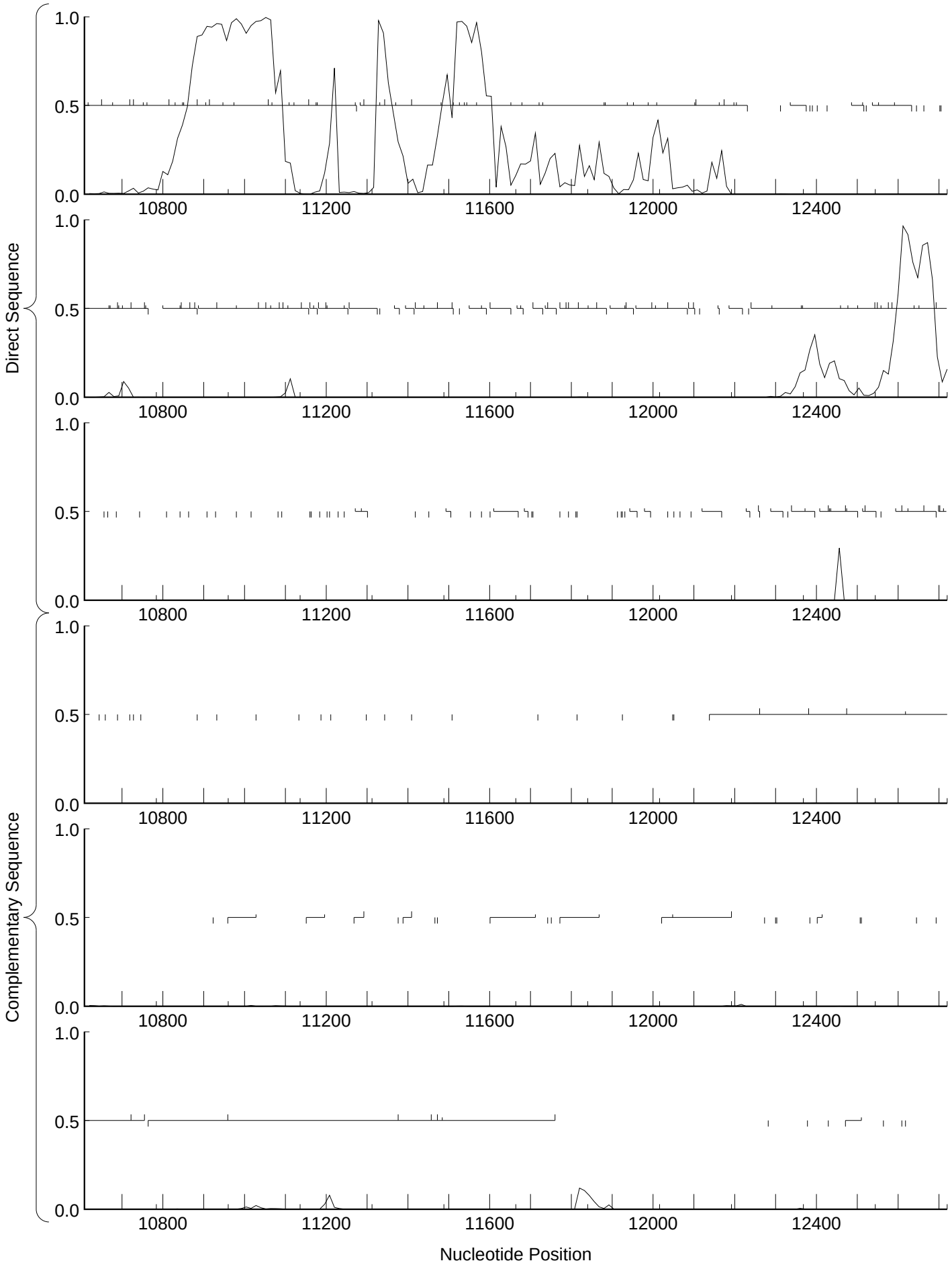
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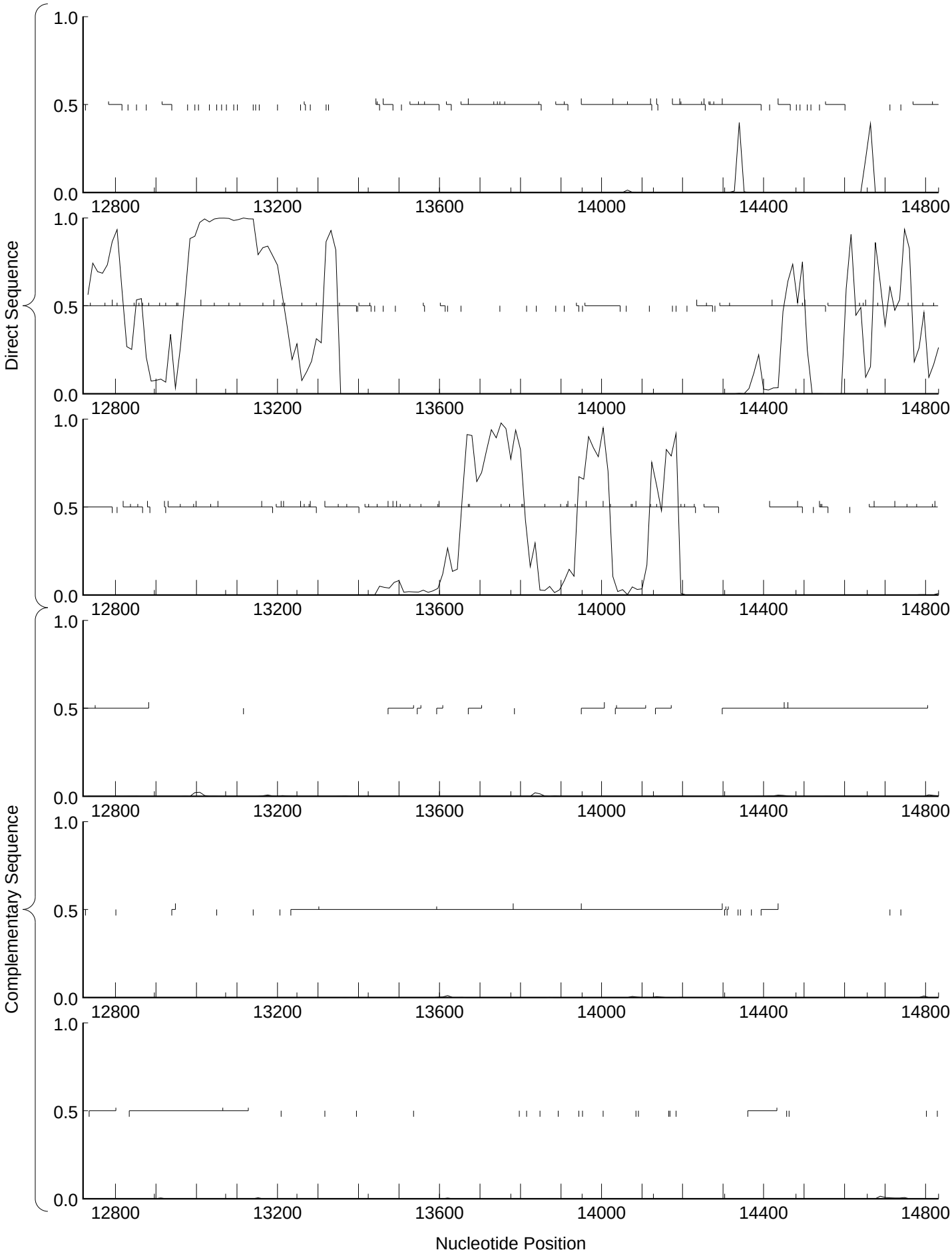
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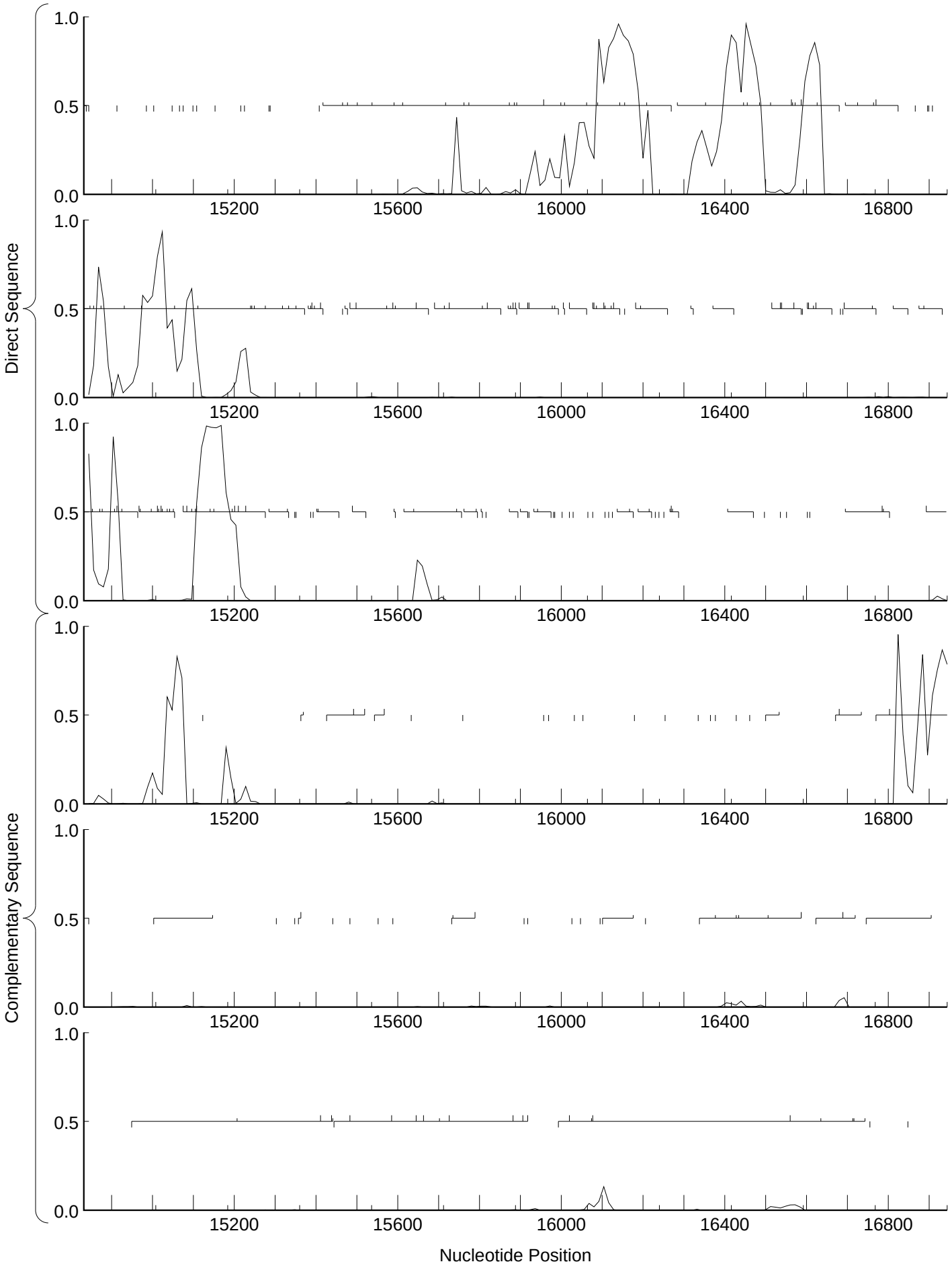
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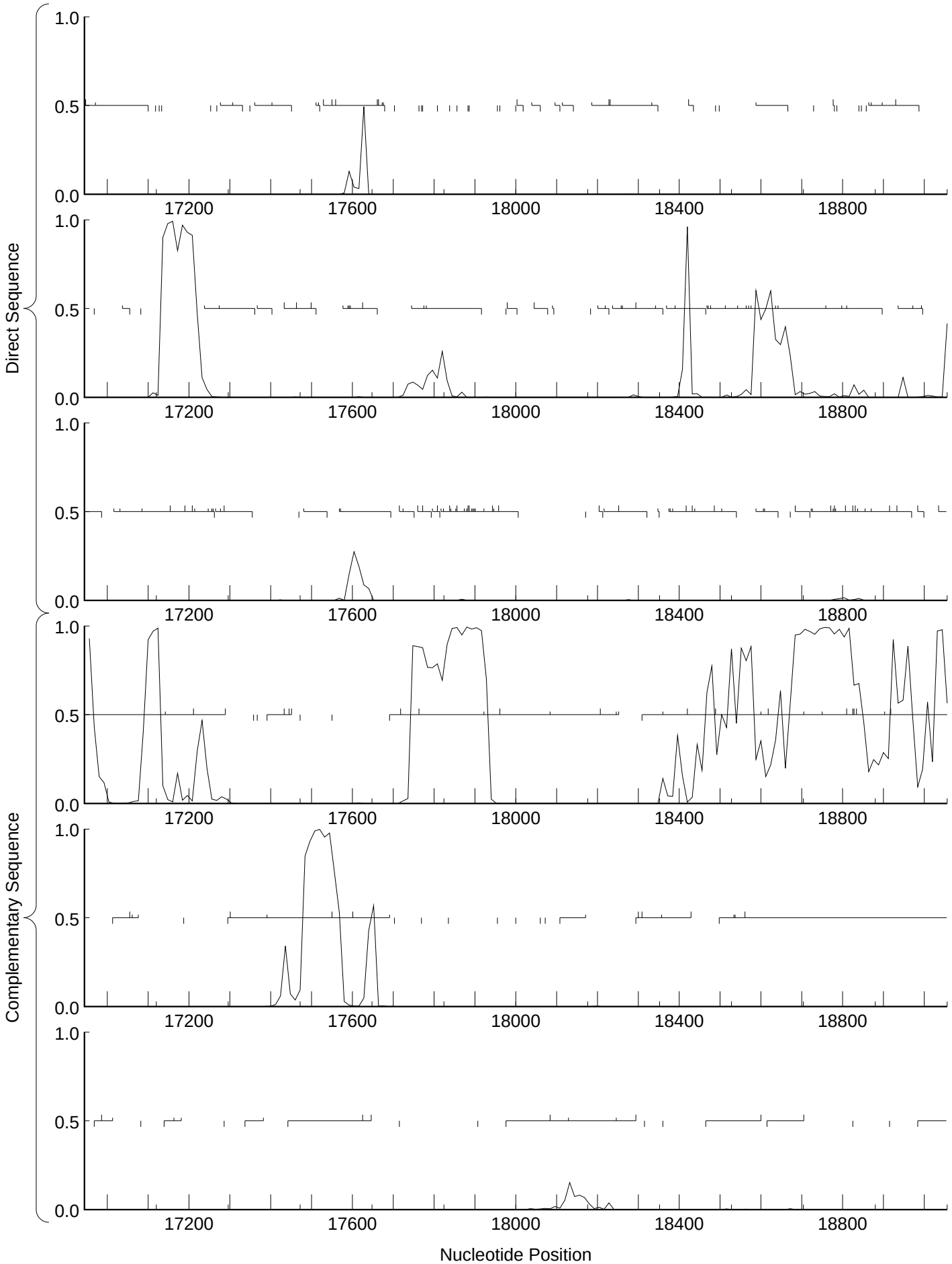


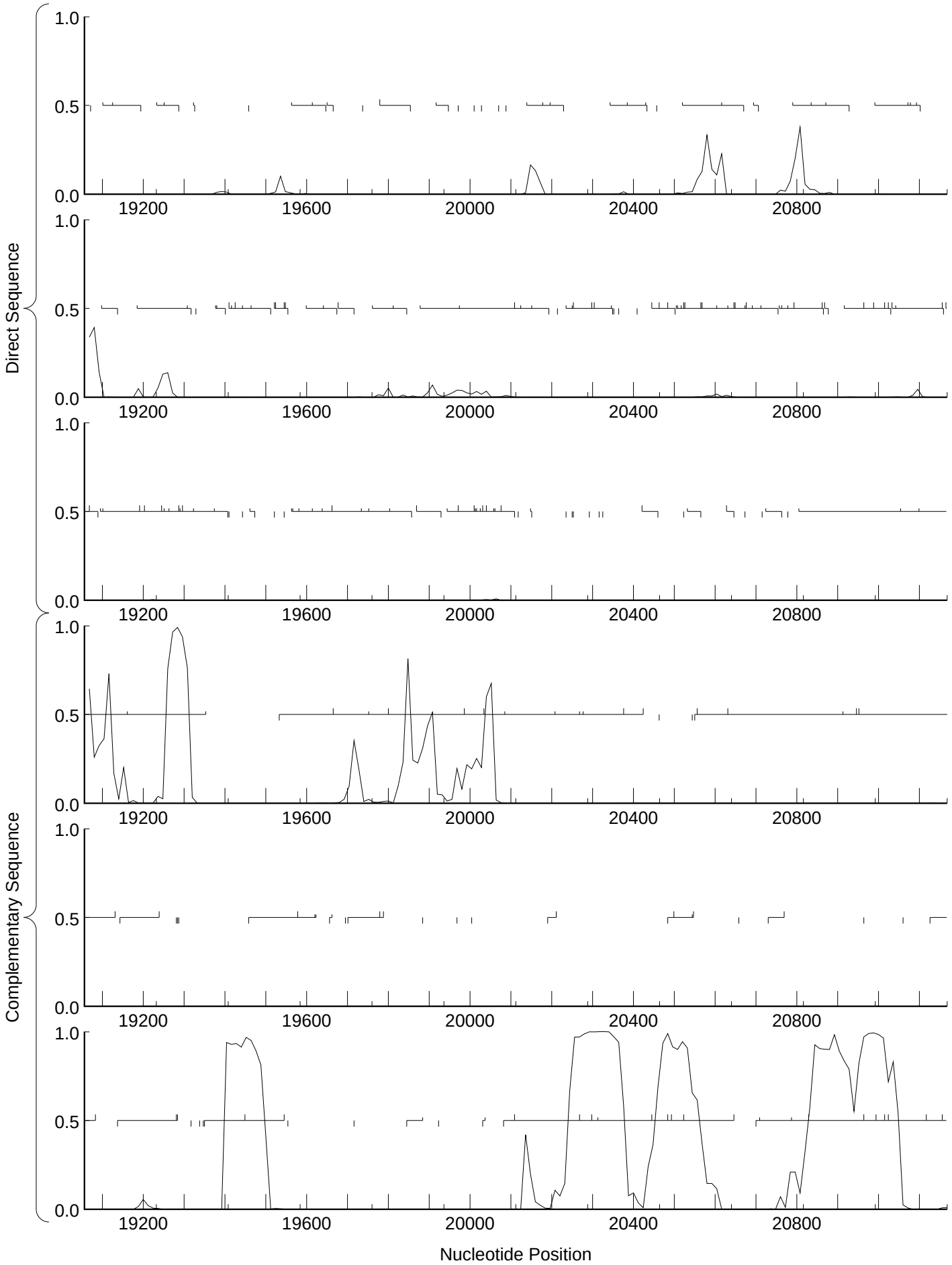
1 complete genome, 29348 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 8/15

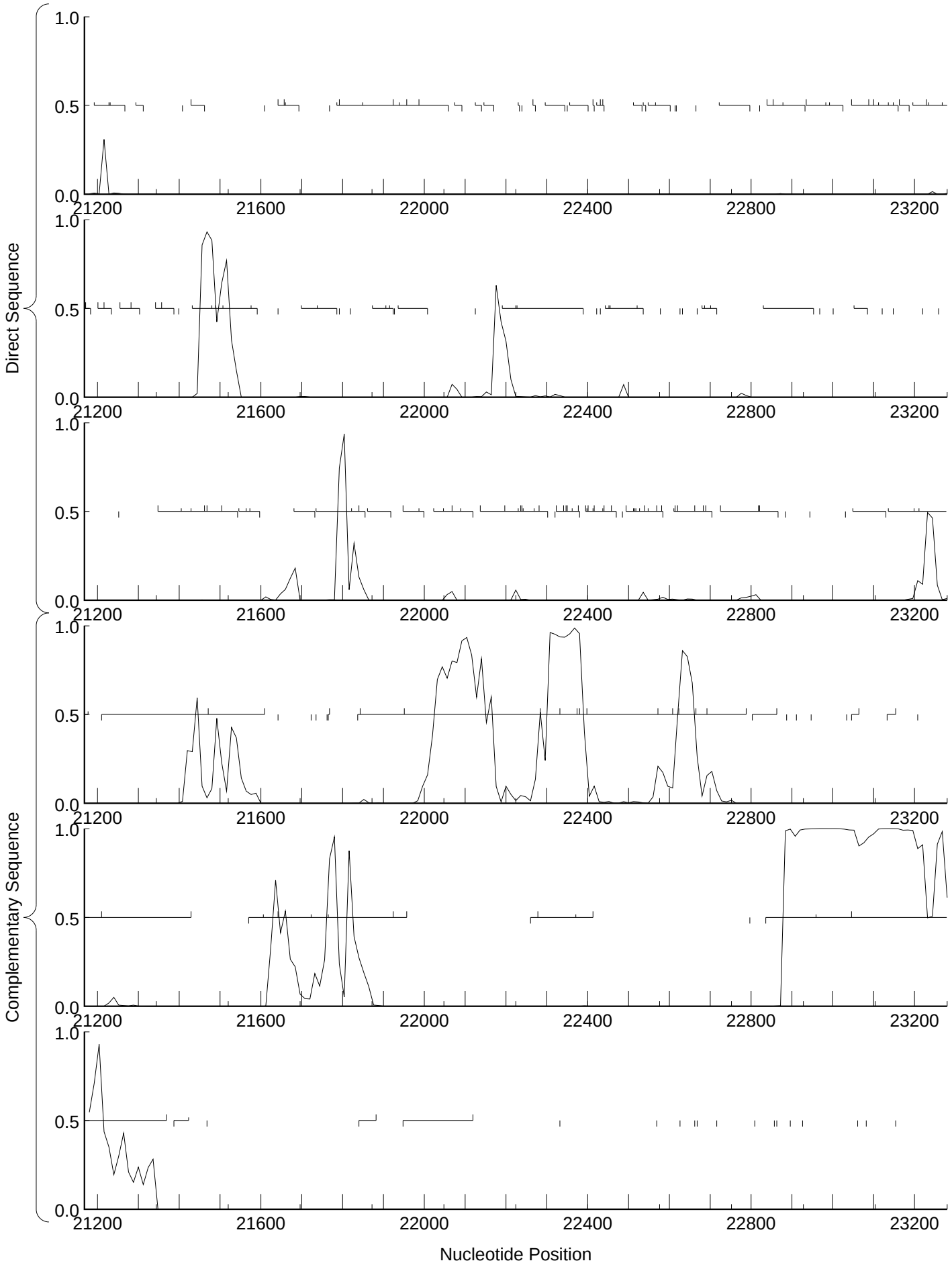


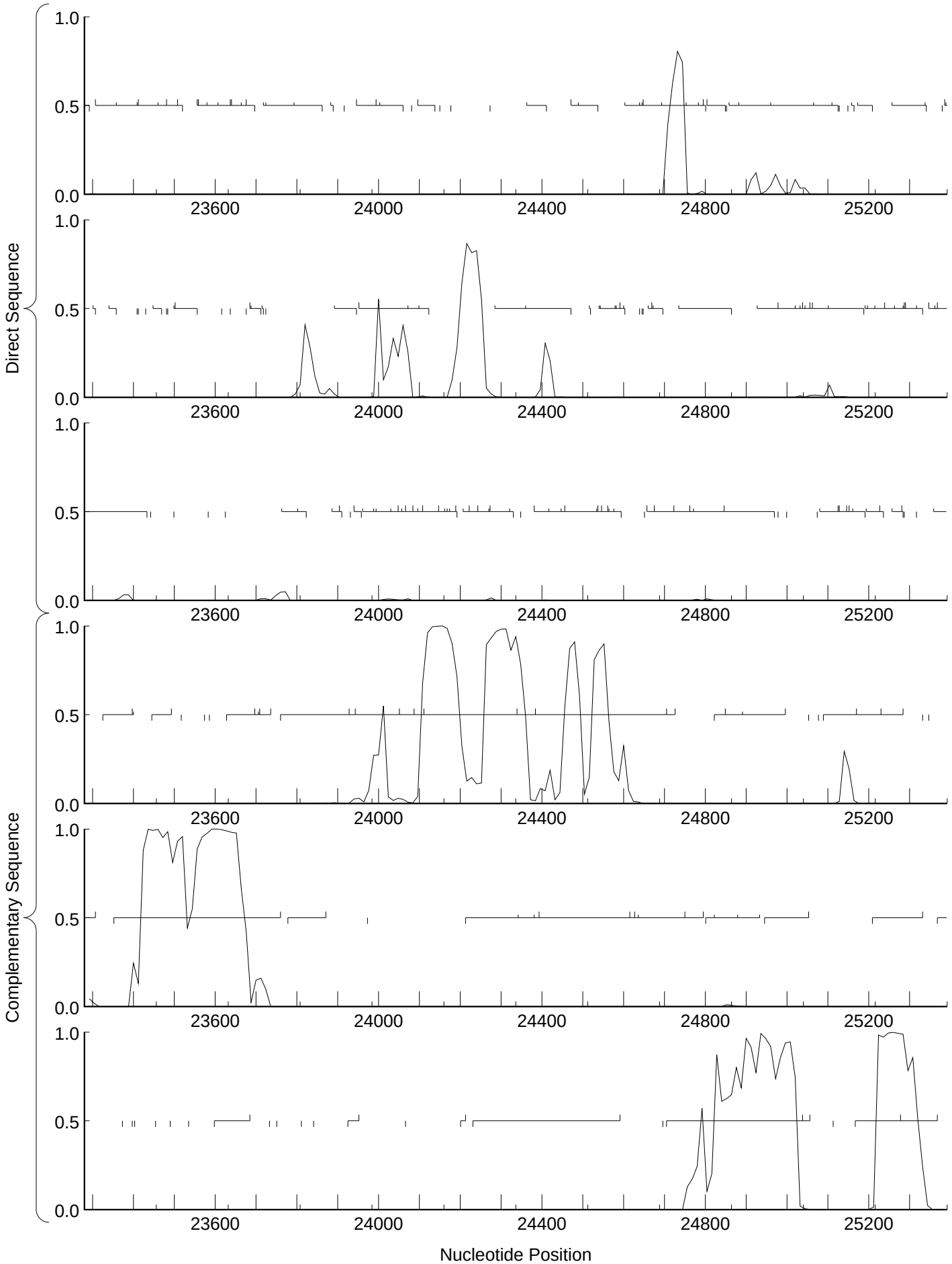
1 complete genome, 29348 bp including 11-base 3’ overhang (TCGTACGGCTT), Cluster BU, Order 4, Window 96, Step 12, 9/15

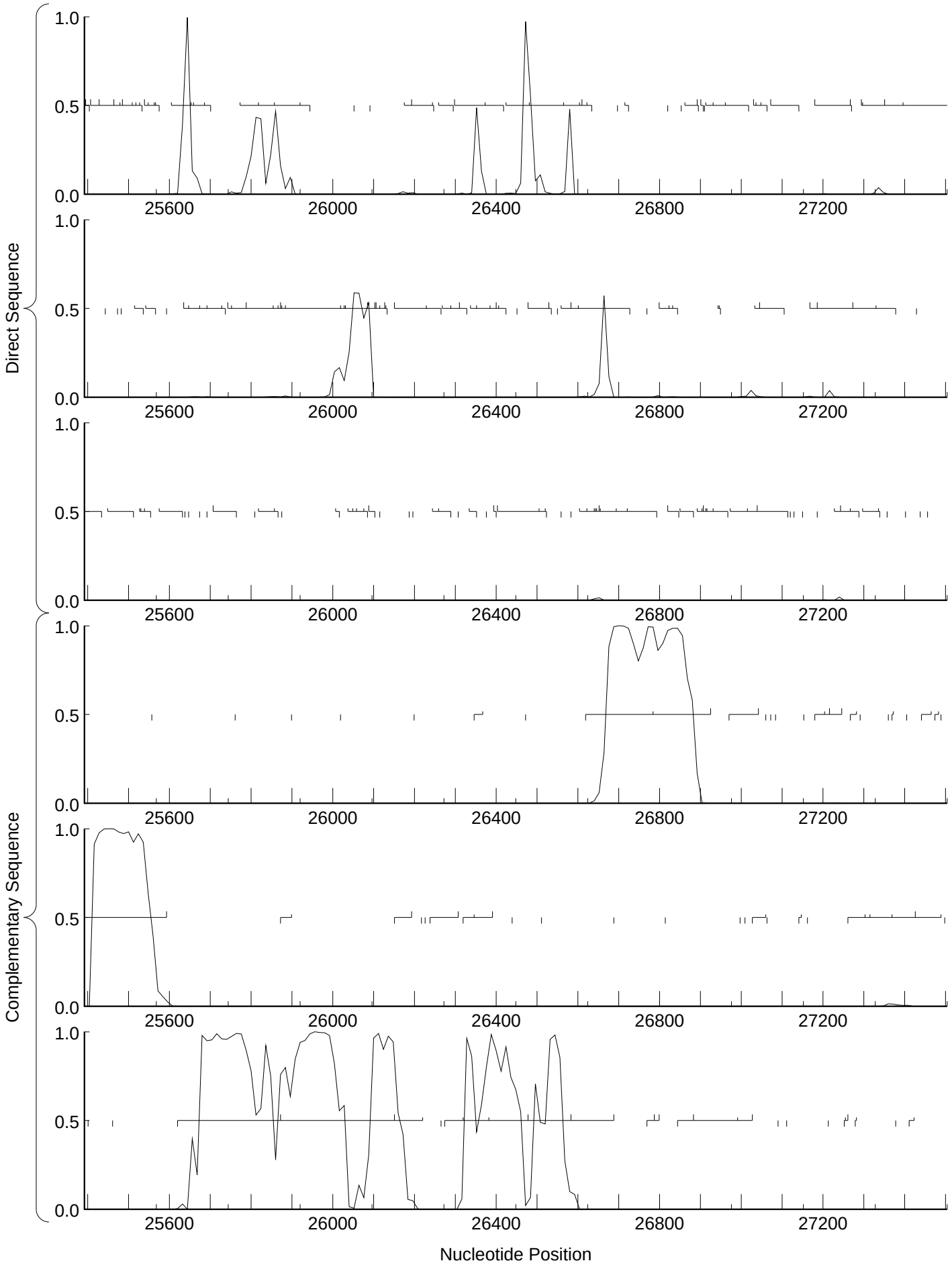


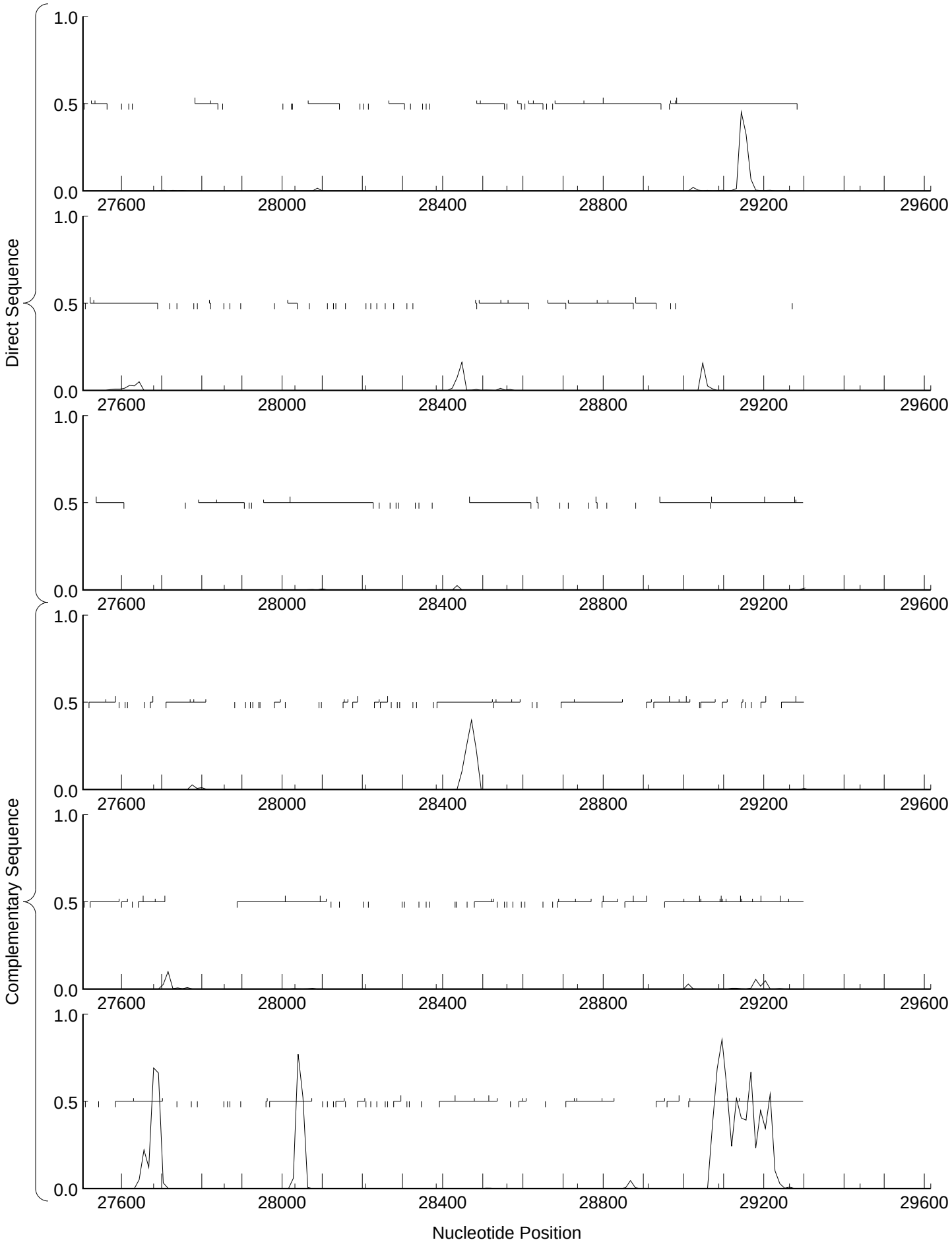

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
17200
17600
18000
18400
18800
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
19200
19600
20000
20400
20800
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
21200
21600
22000
22400
22800
23200
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
23600
24000
24400
24800
25200
Nucleotide Position


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


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
27600
28000
28400
28800
29200
29600
Nucleotide Position