GeneMark

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

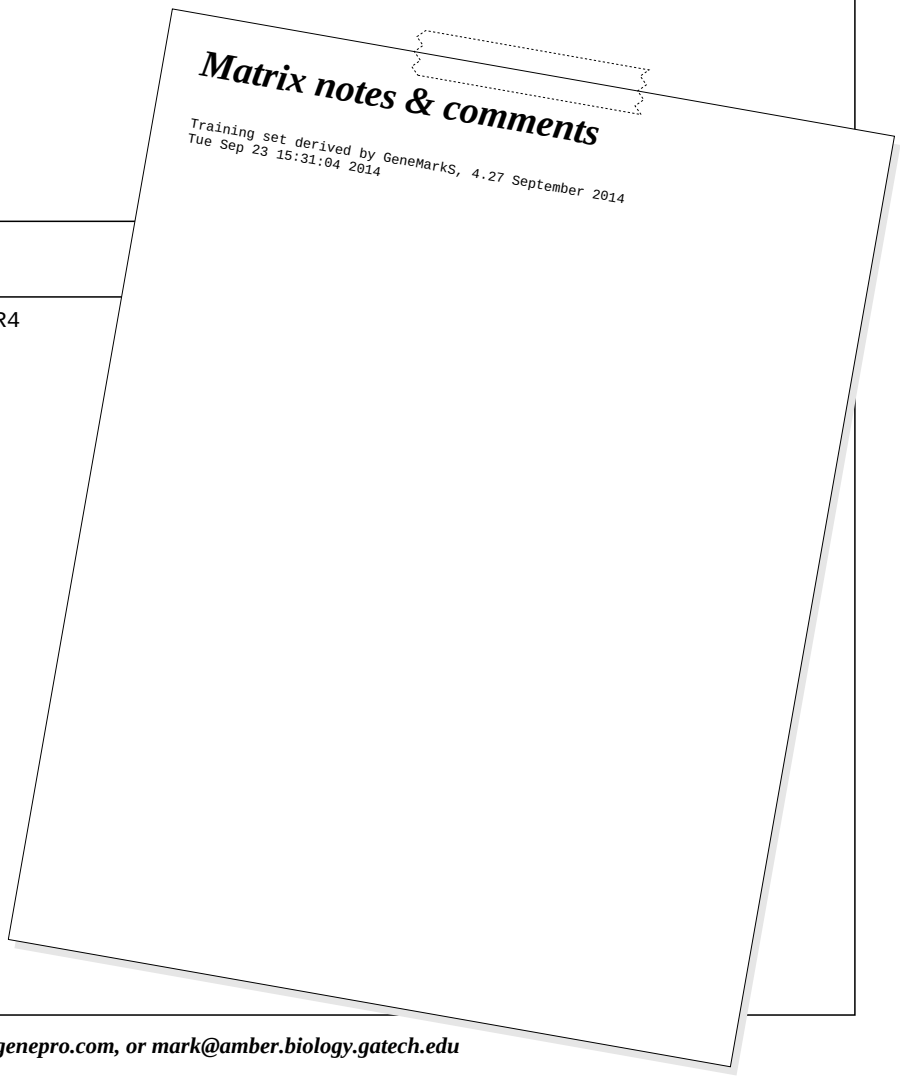
PROGRAM INFORMATION

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Sequence         :  Rhodococcus phage RER2 complete genome, 46596 bp including 10-base 3' overhang (CGGCCGTGAT)
Analysis Date    :  3/28/24 at 20:31:21
Pages            :  24
Sequence Length  :  46596 bp
GC Content       :   58.57%

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

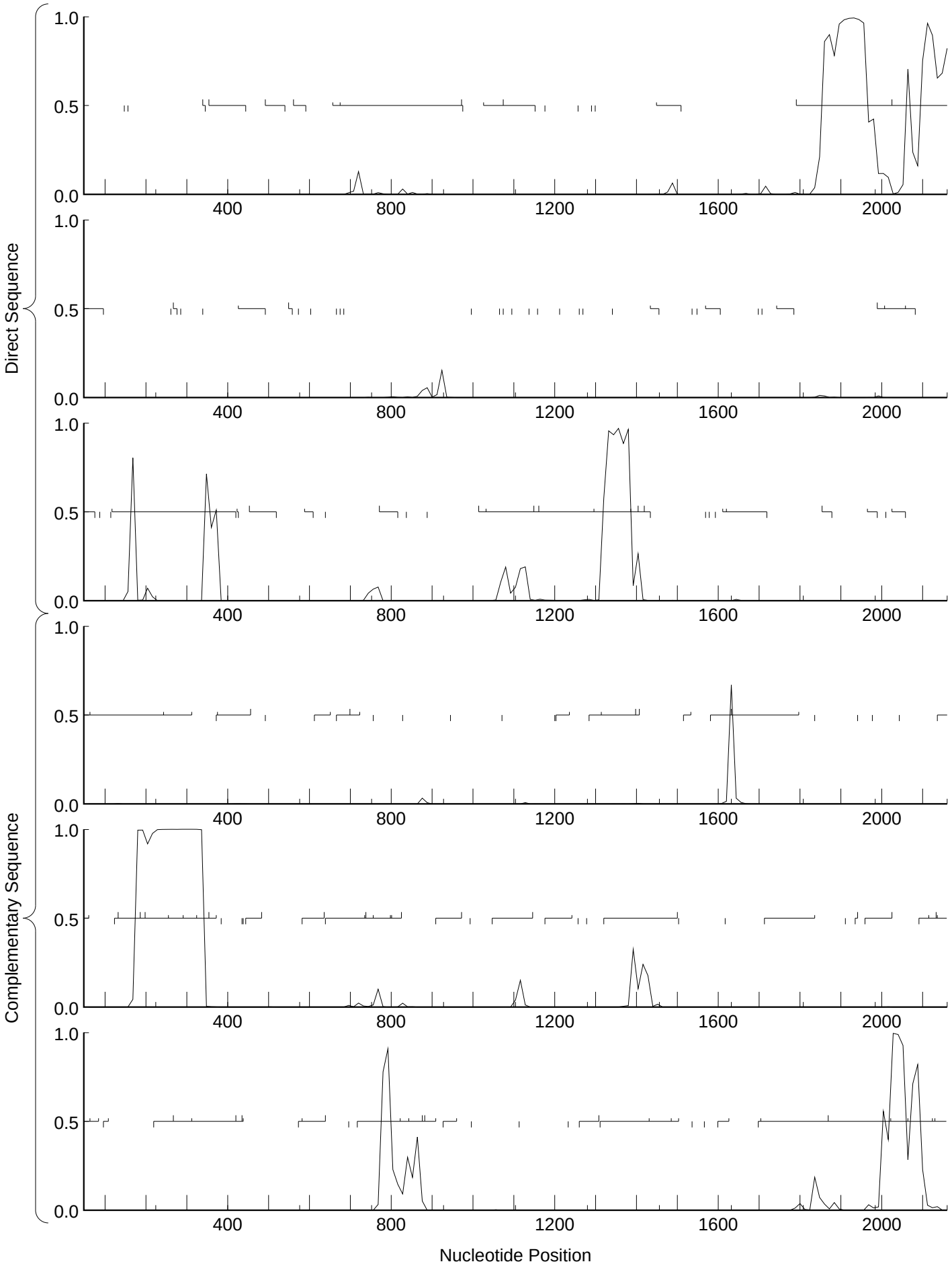
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Matrix     :  Rhodococcus_erythropolis_PR4
Author     :  -

Order      :  4
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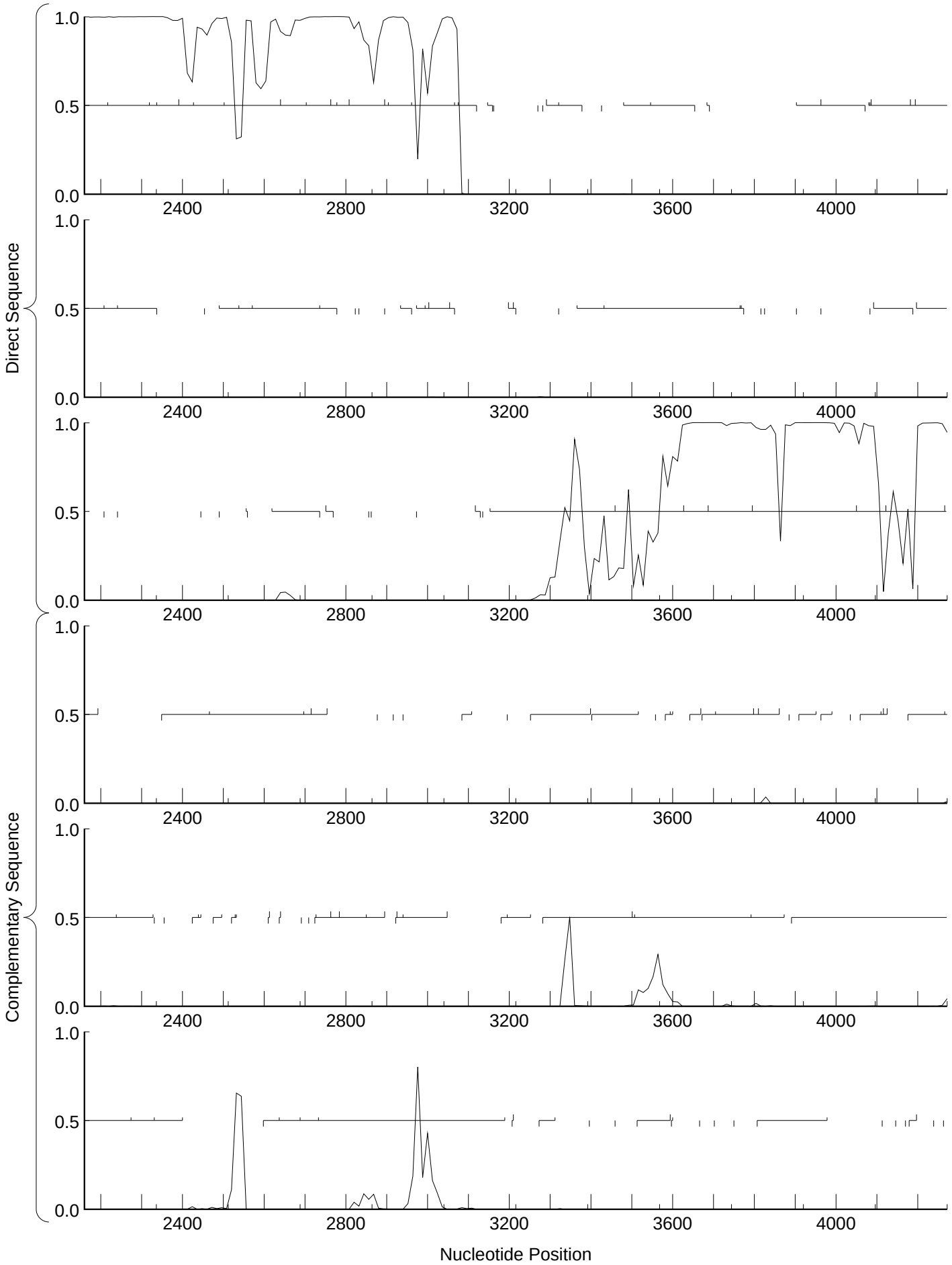

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

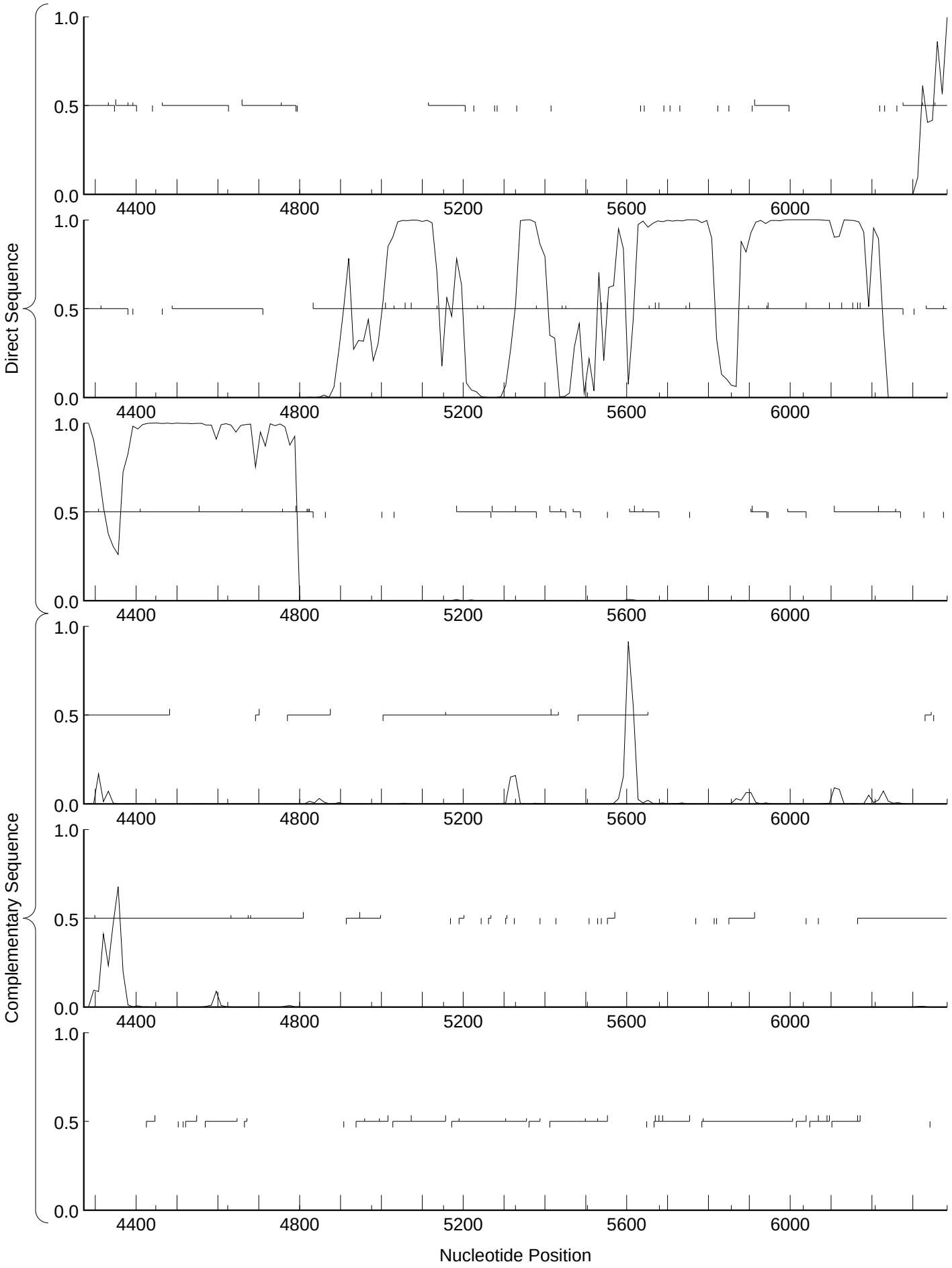
R2 complete genome, 46596 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 2/24



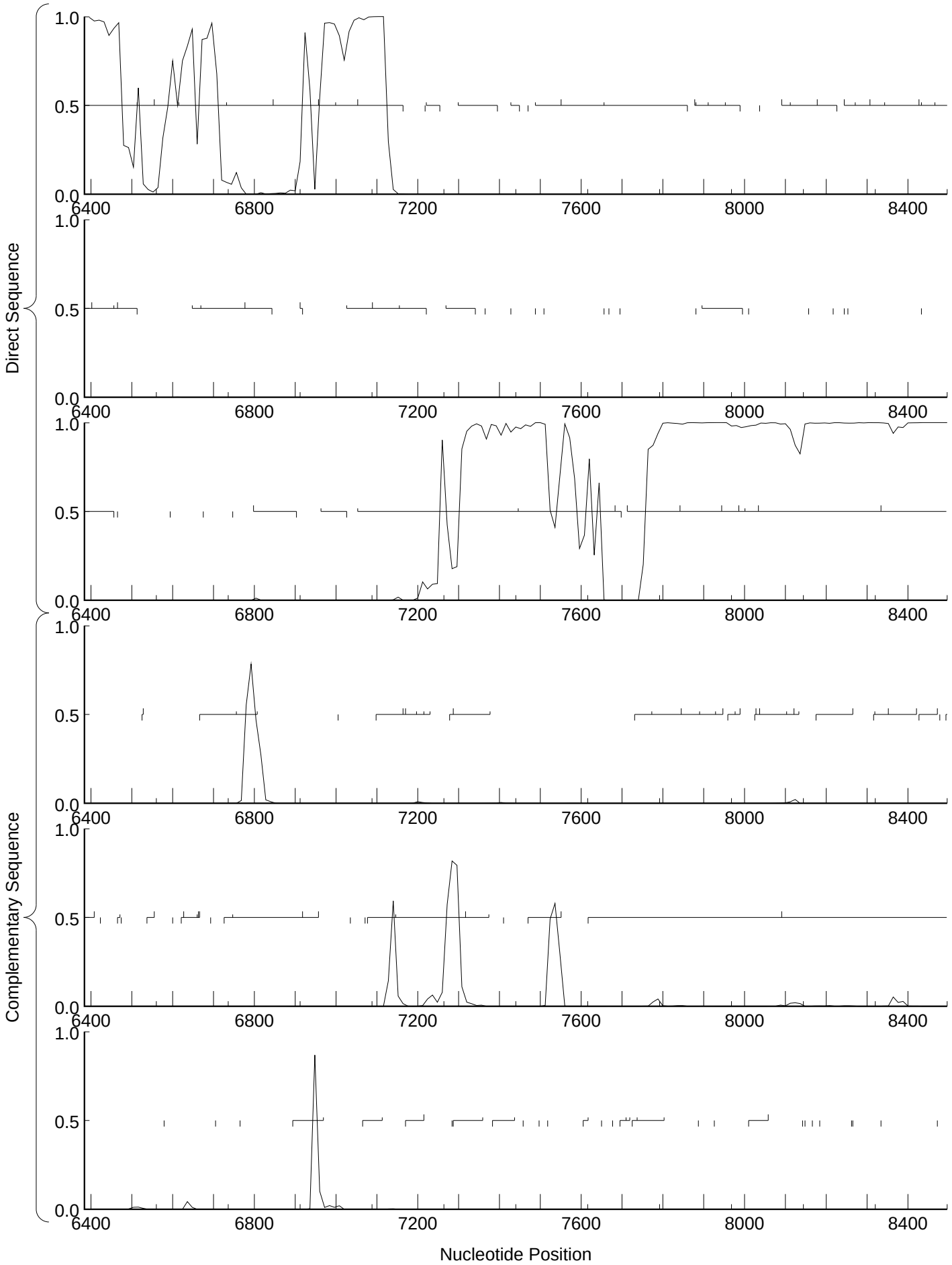
R2 complete genome, 46596 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 3/24



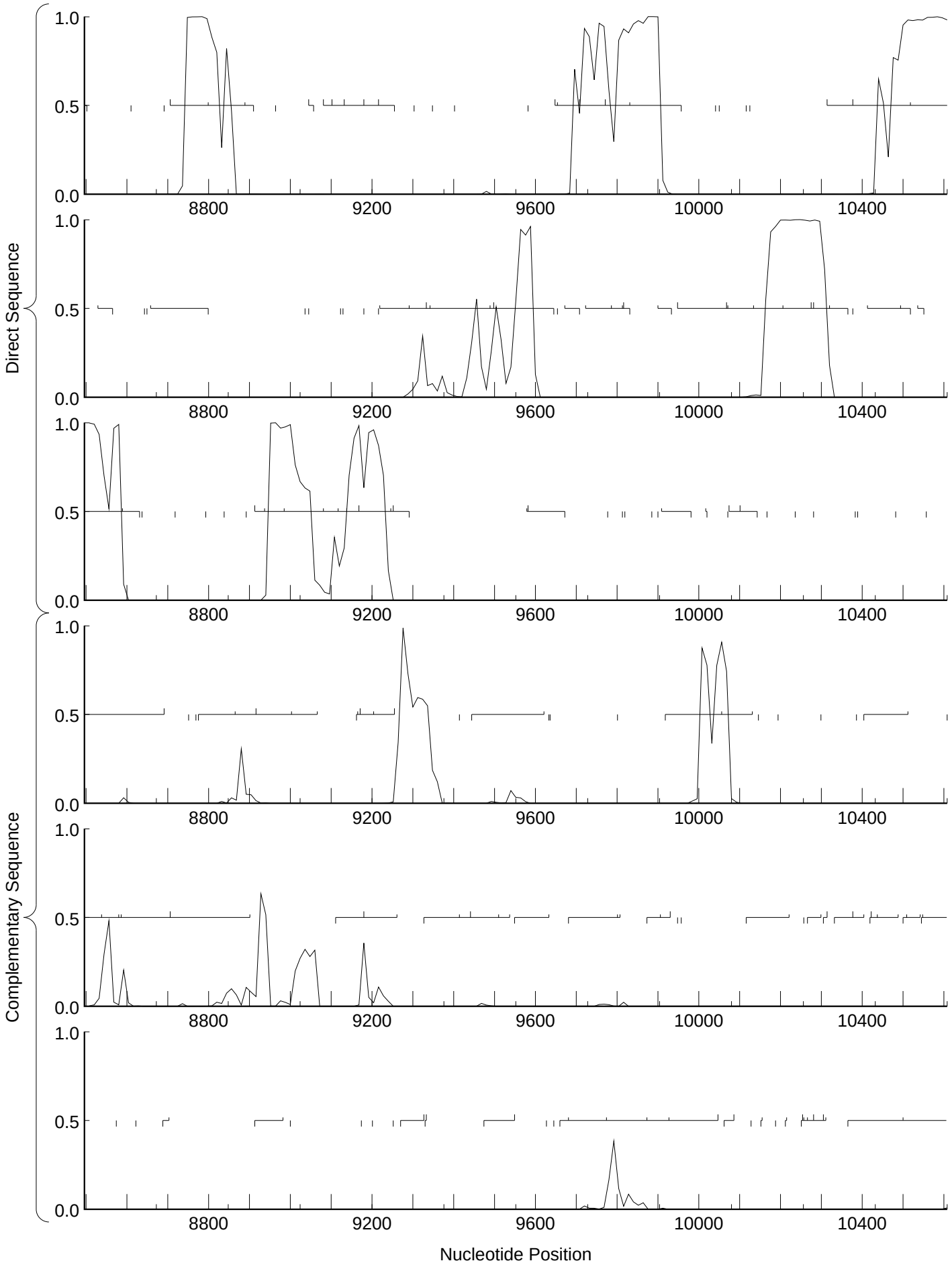
R2 complete genome, 46596 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 4/24



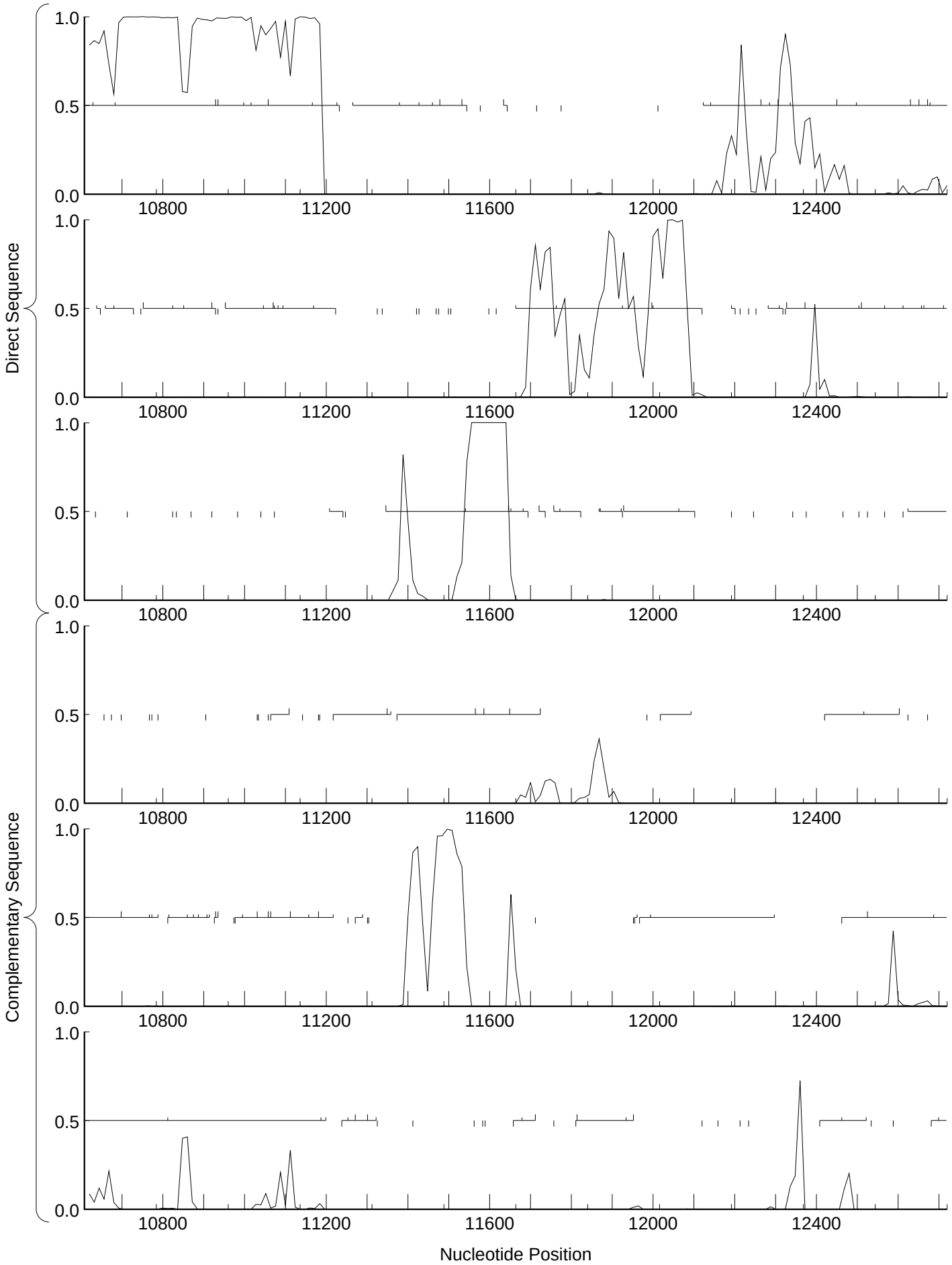
R2 complete genome, 46596 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 5/24



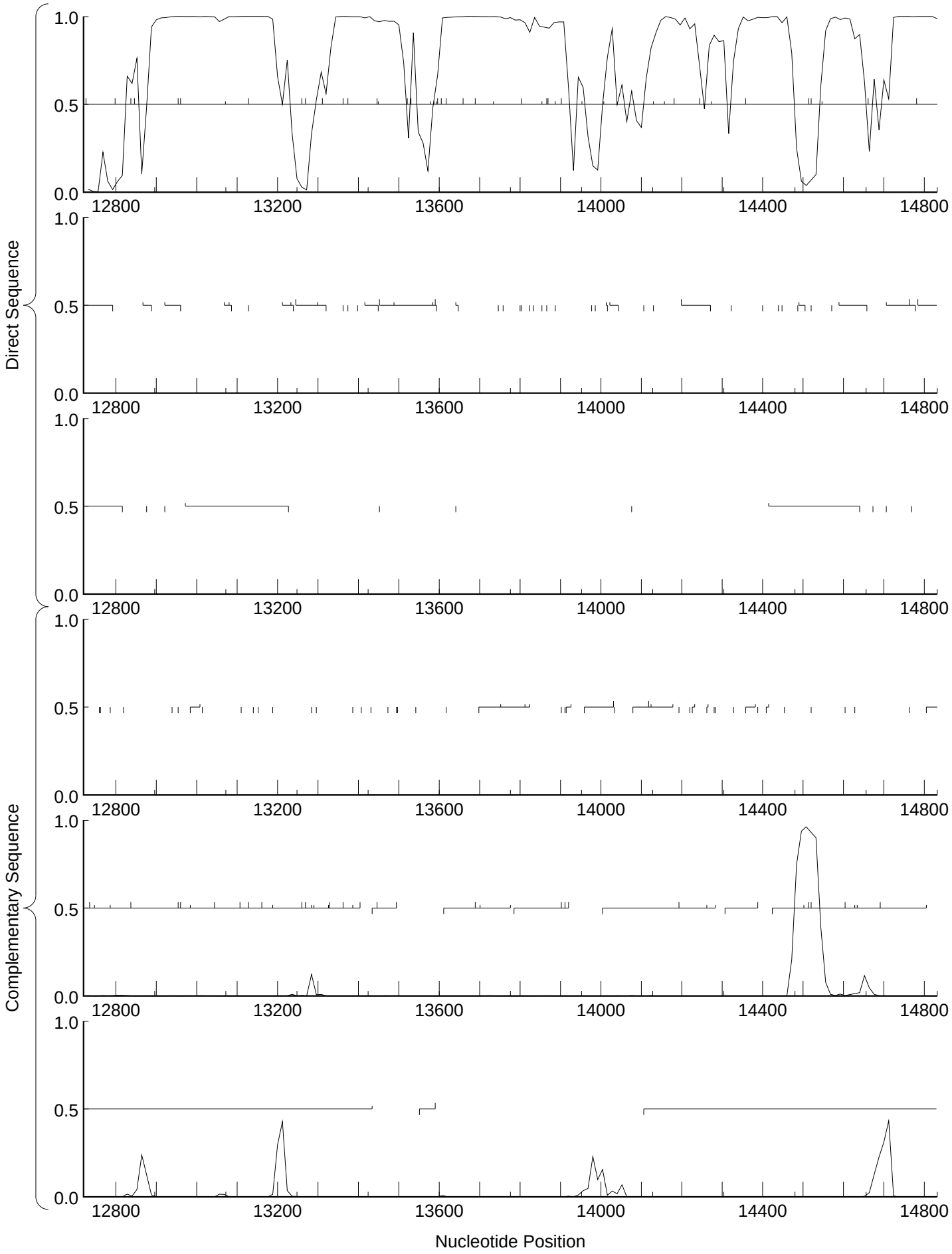
R2 complete genome, 46596 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 6/24



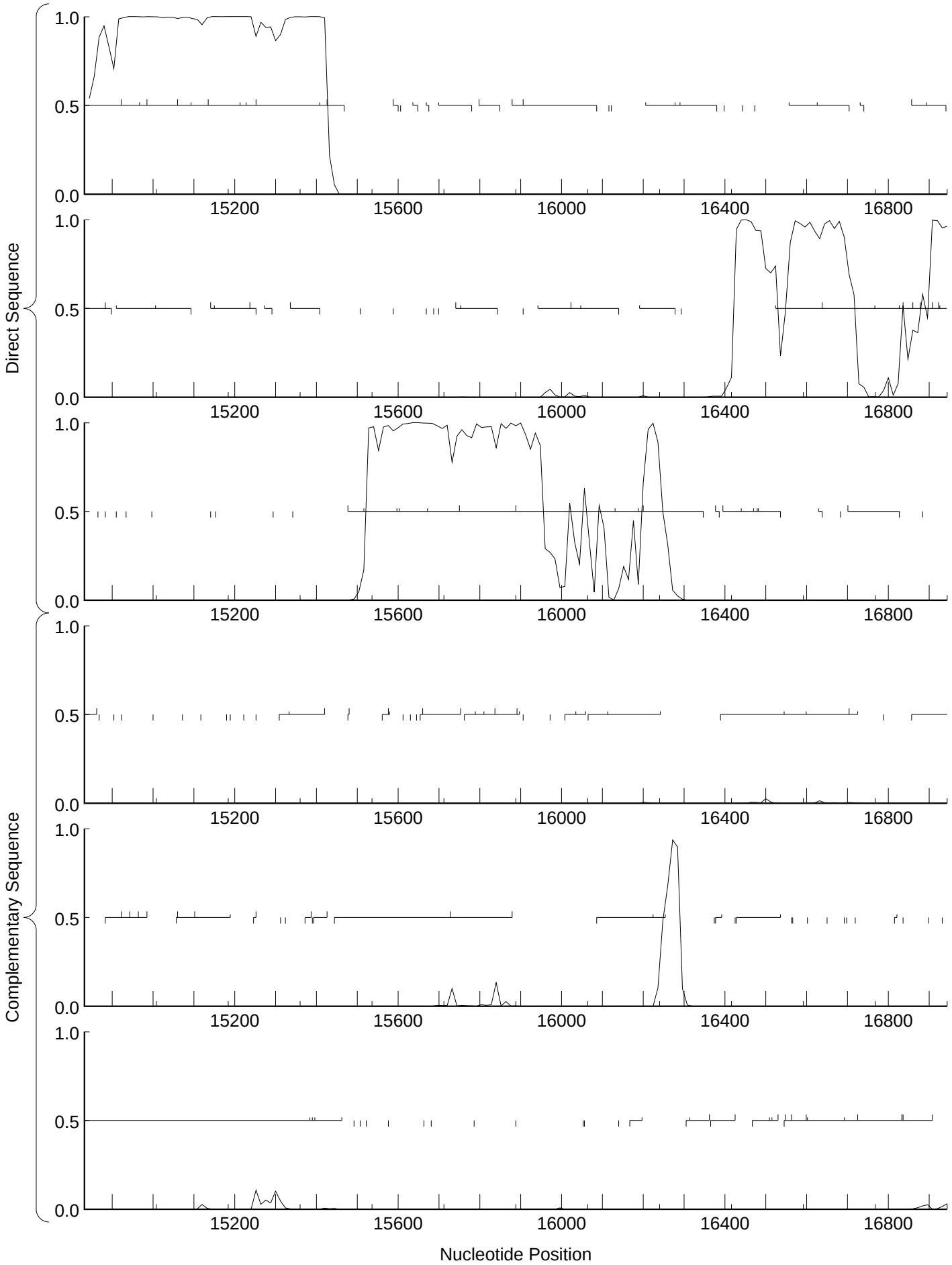
R2 complete genome, 46596 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 7/24



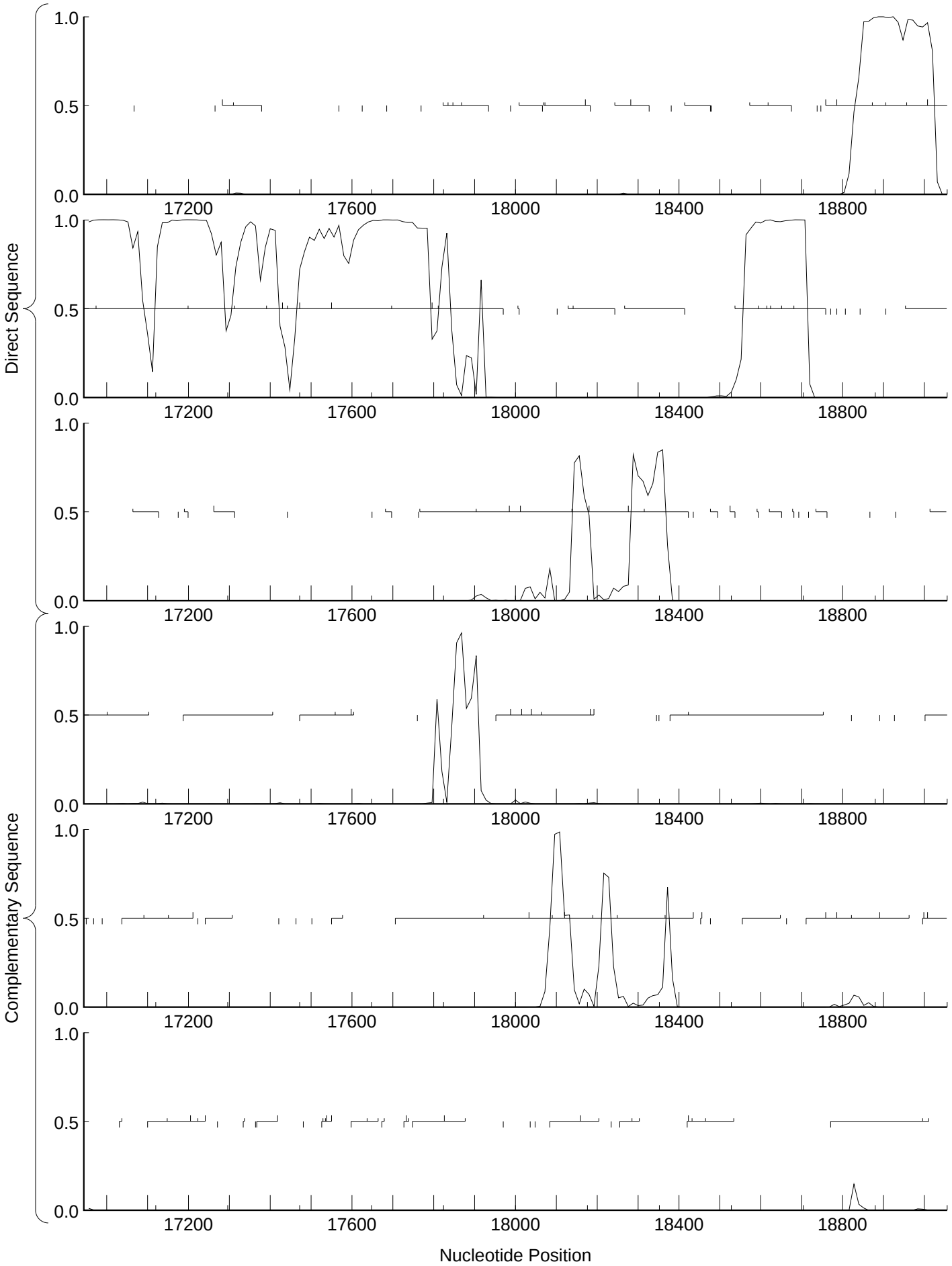
R2 complete genome, 46596 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 8/24



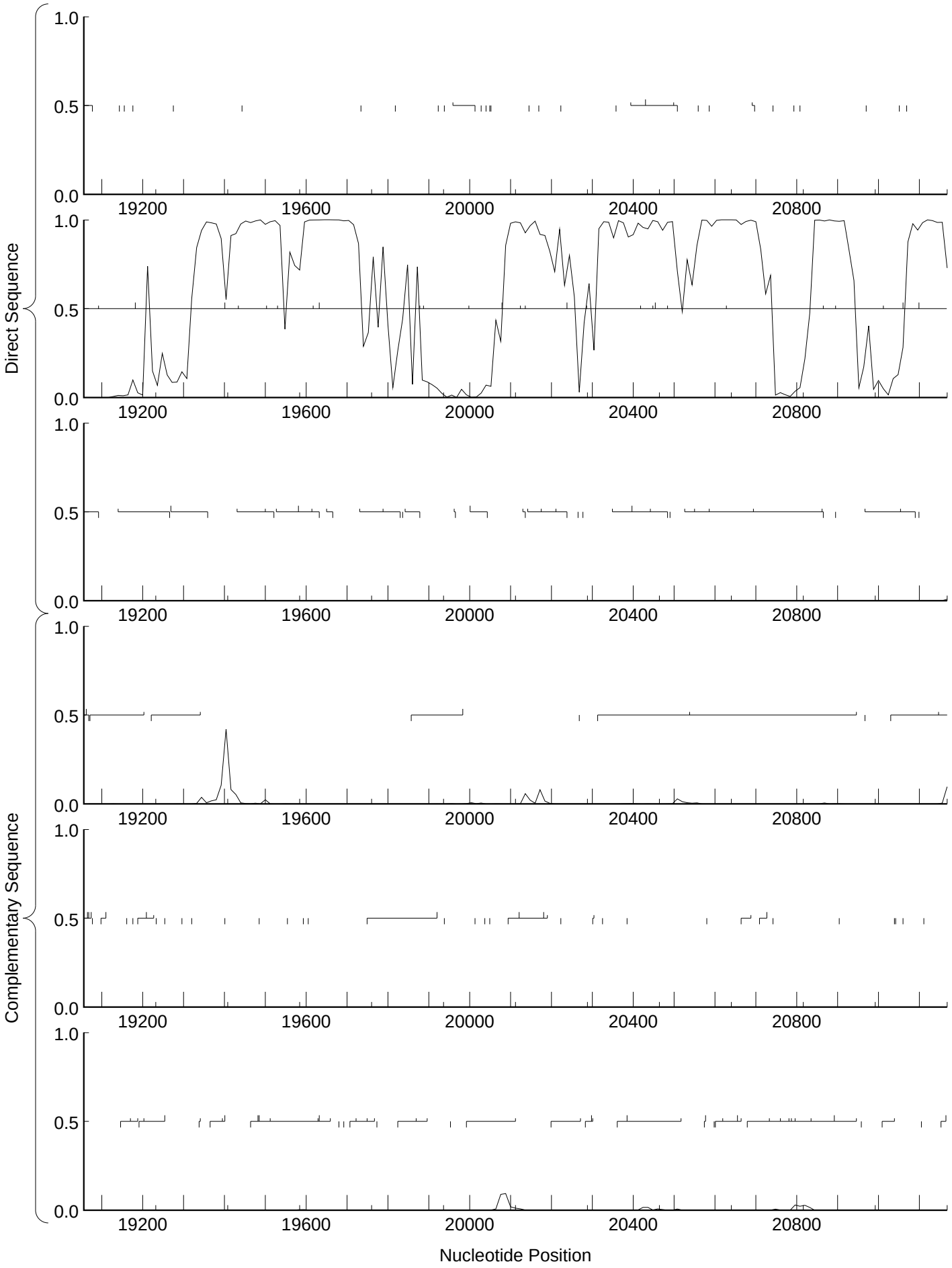
R2 complete genome, 46596 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 9/24



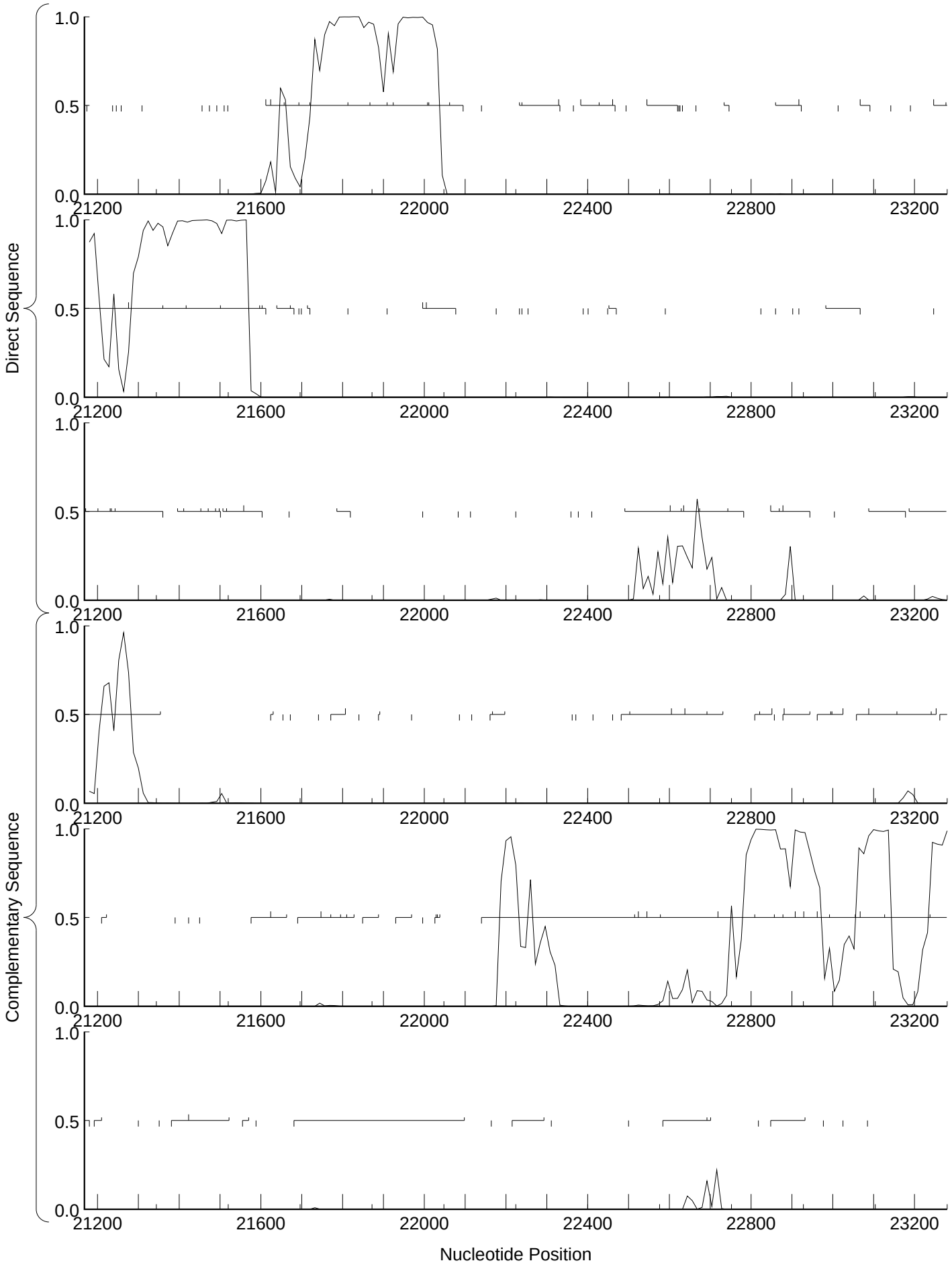
2 complete genome, 46596 bp including 10-base 3’ overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 10/24



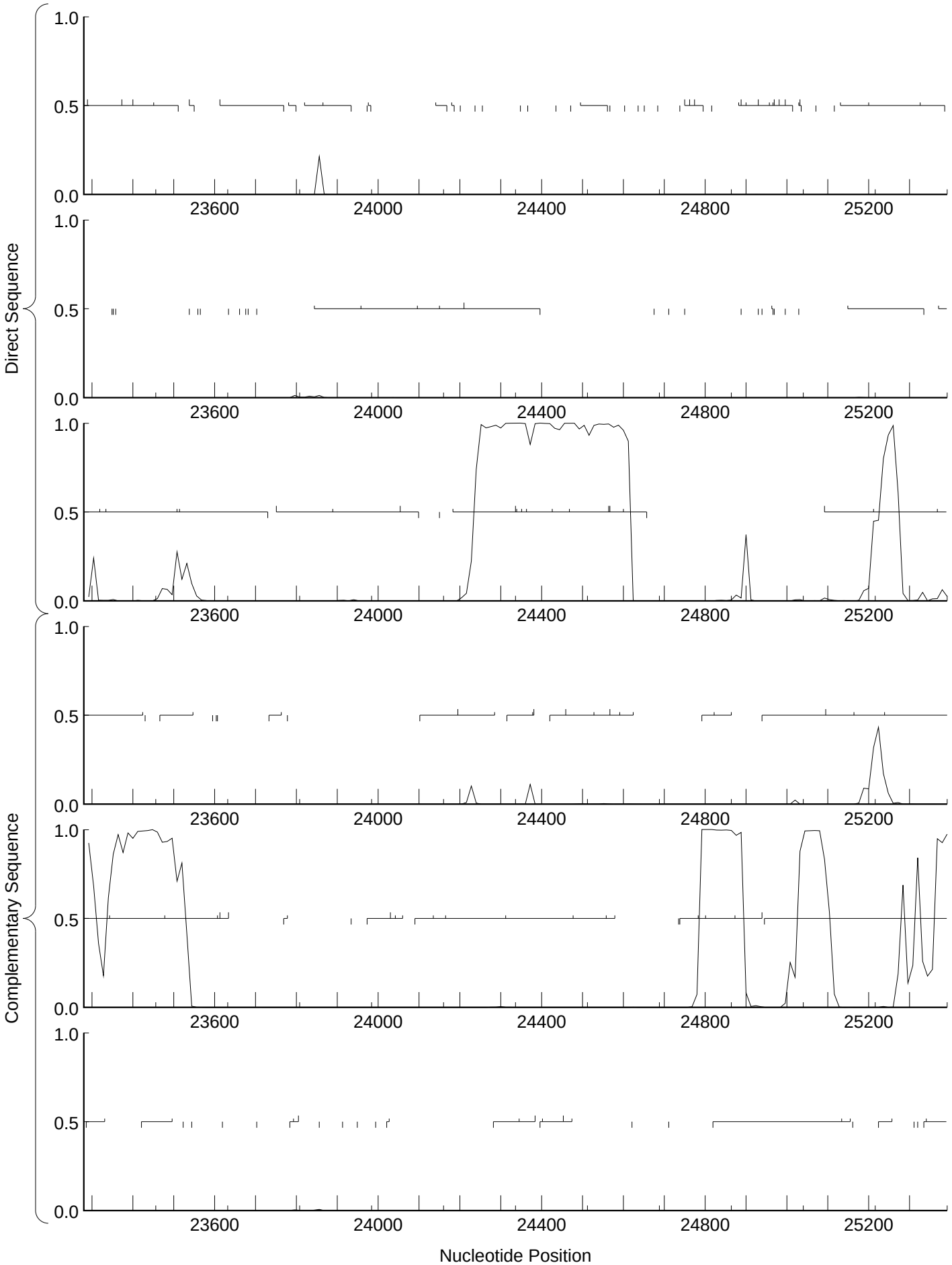
2 complete genome, 46596 bp including 10-base 3’ overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 11/24



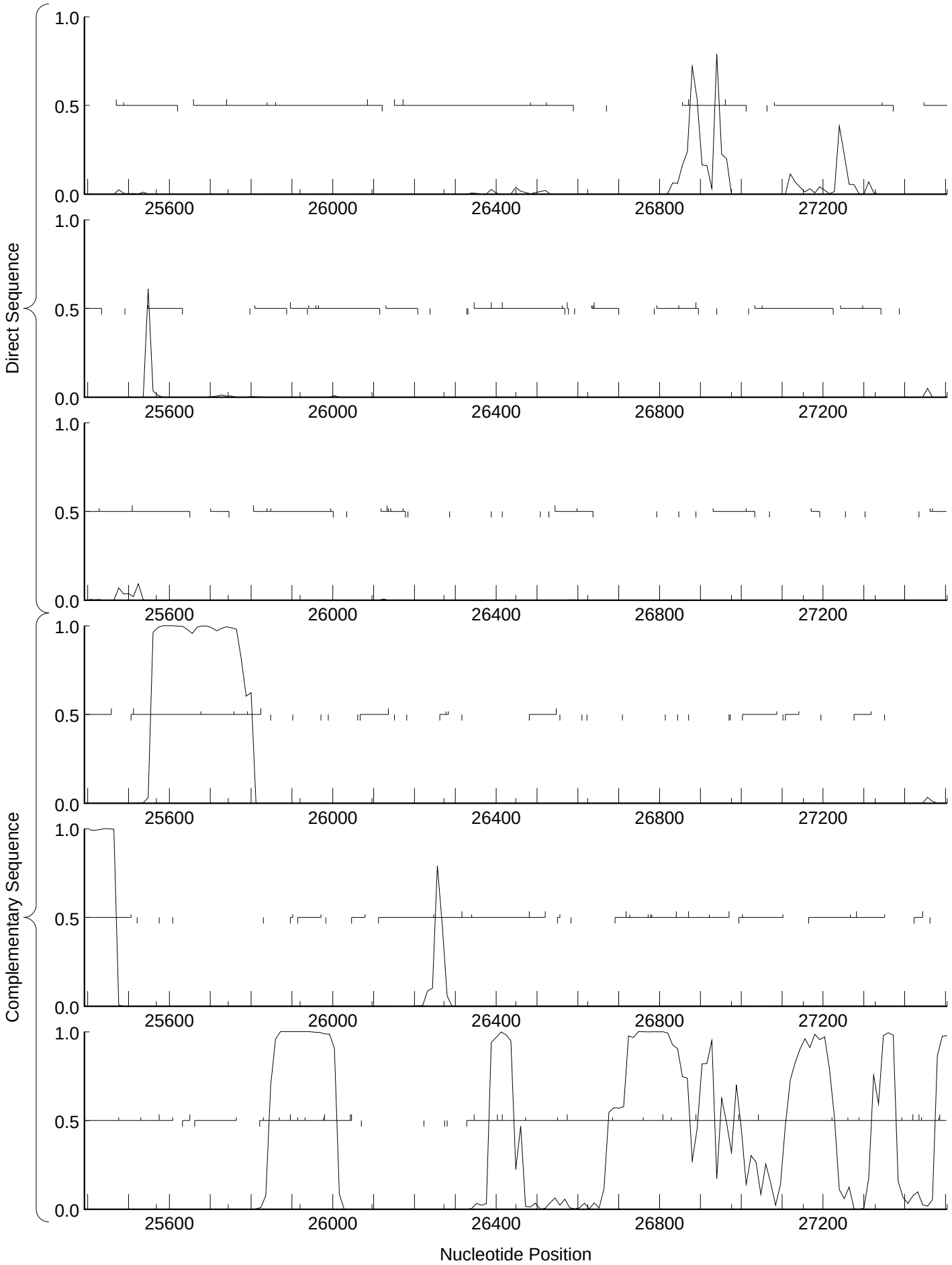
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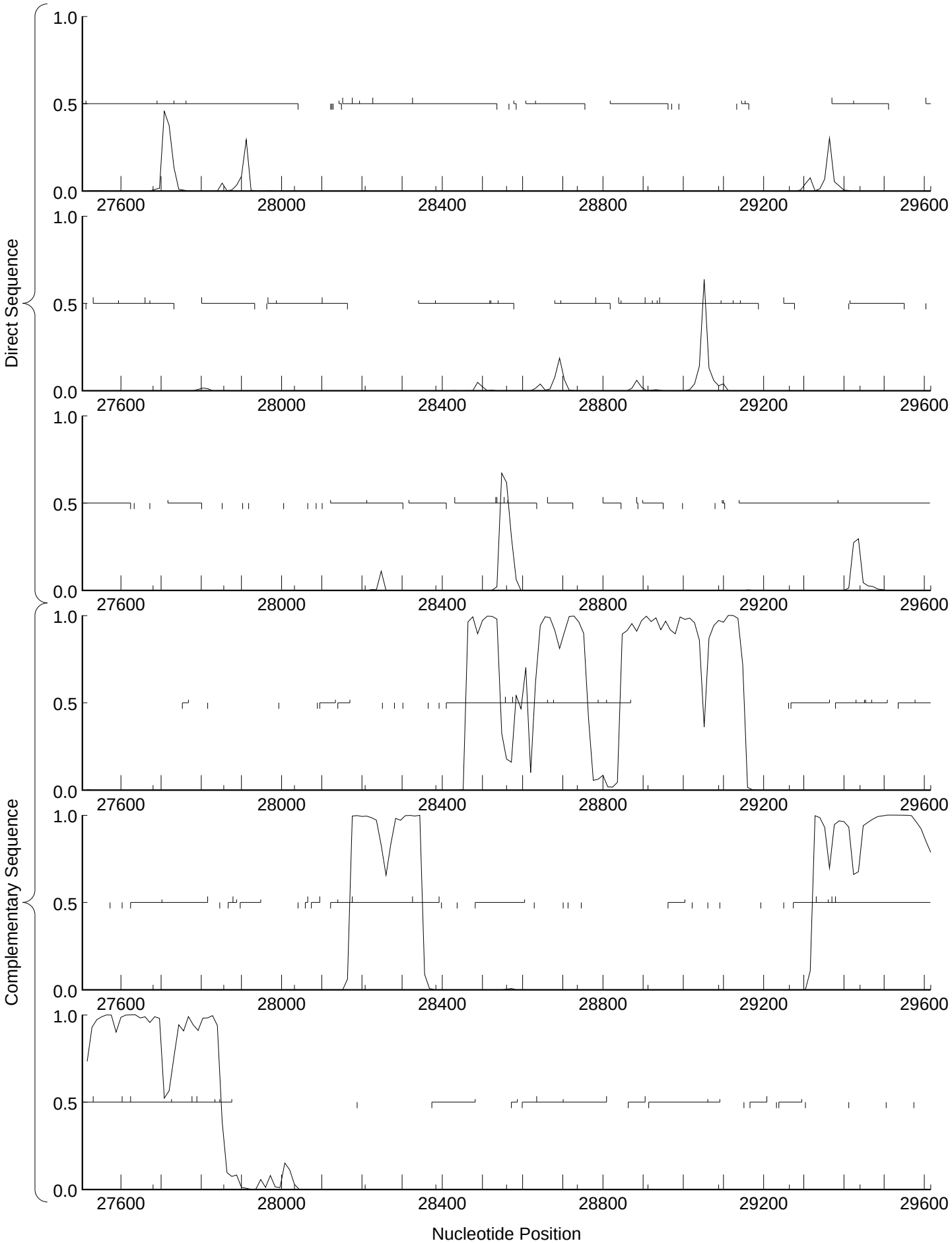
2 complete genome, 46596 bp including 10-base 3’ overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 13/24



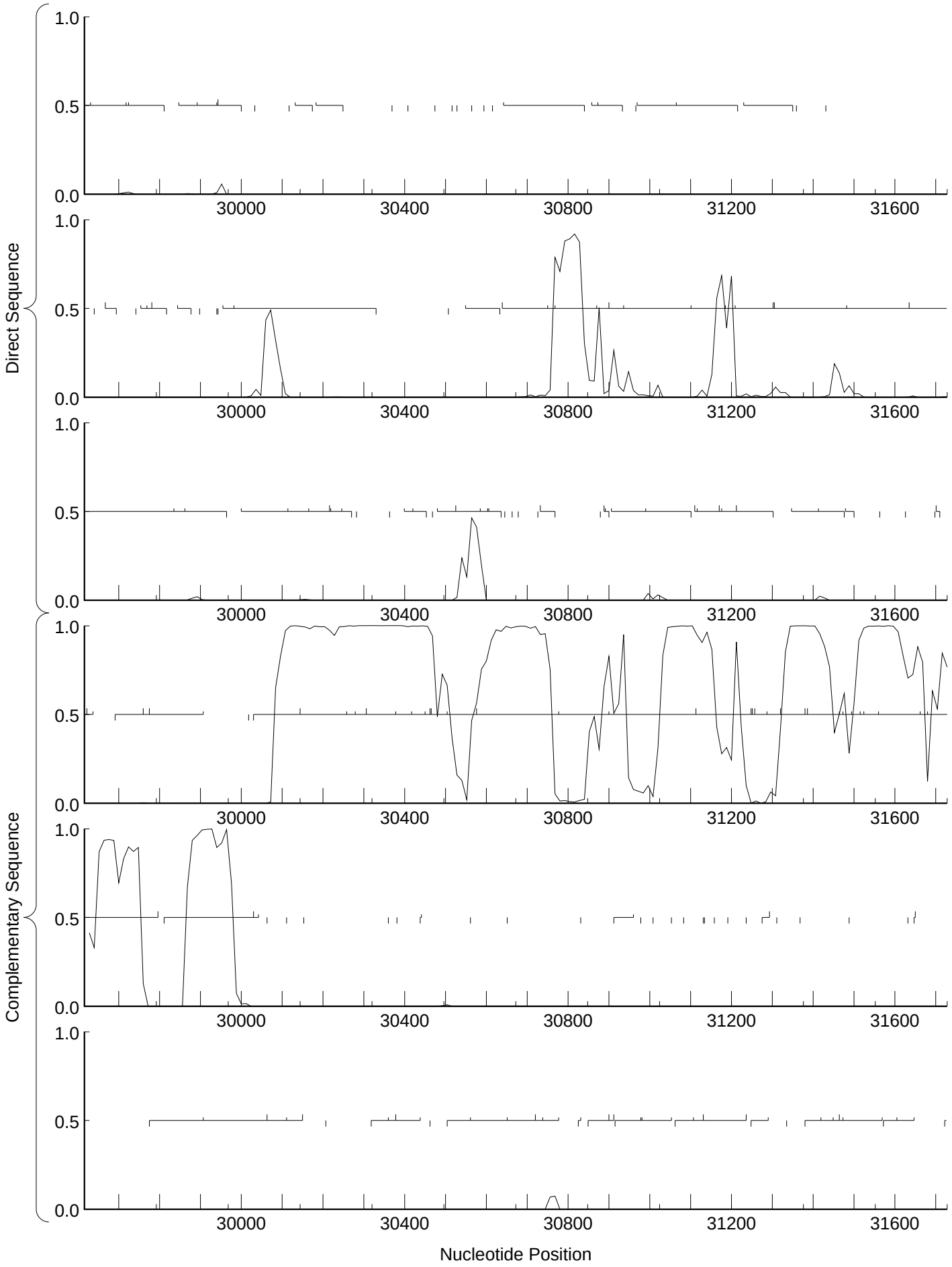
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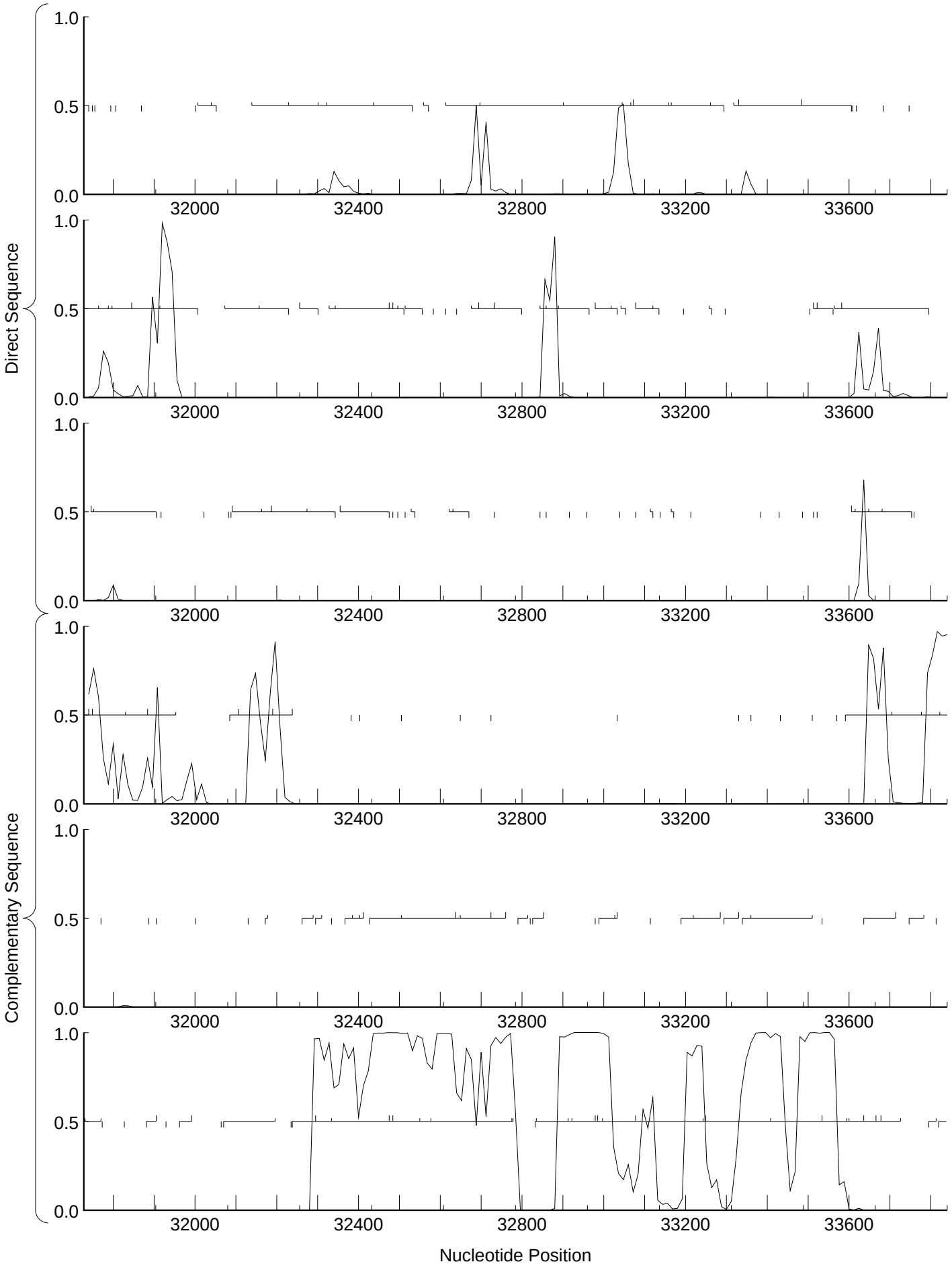
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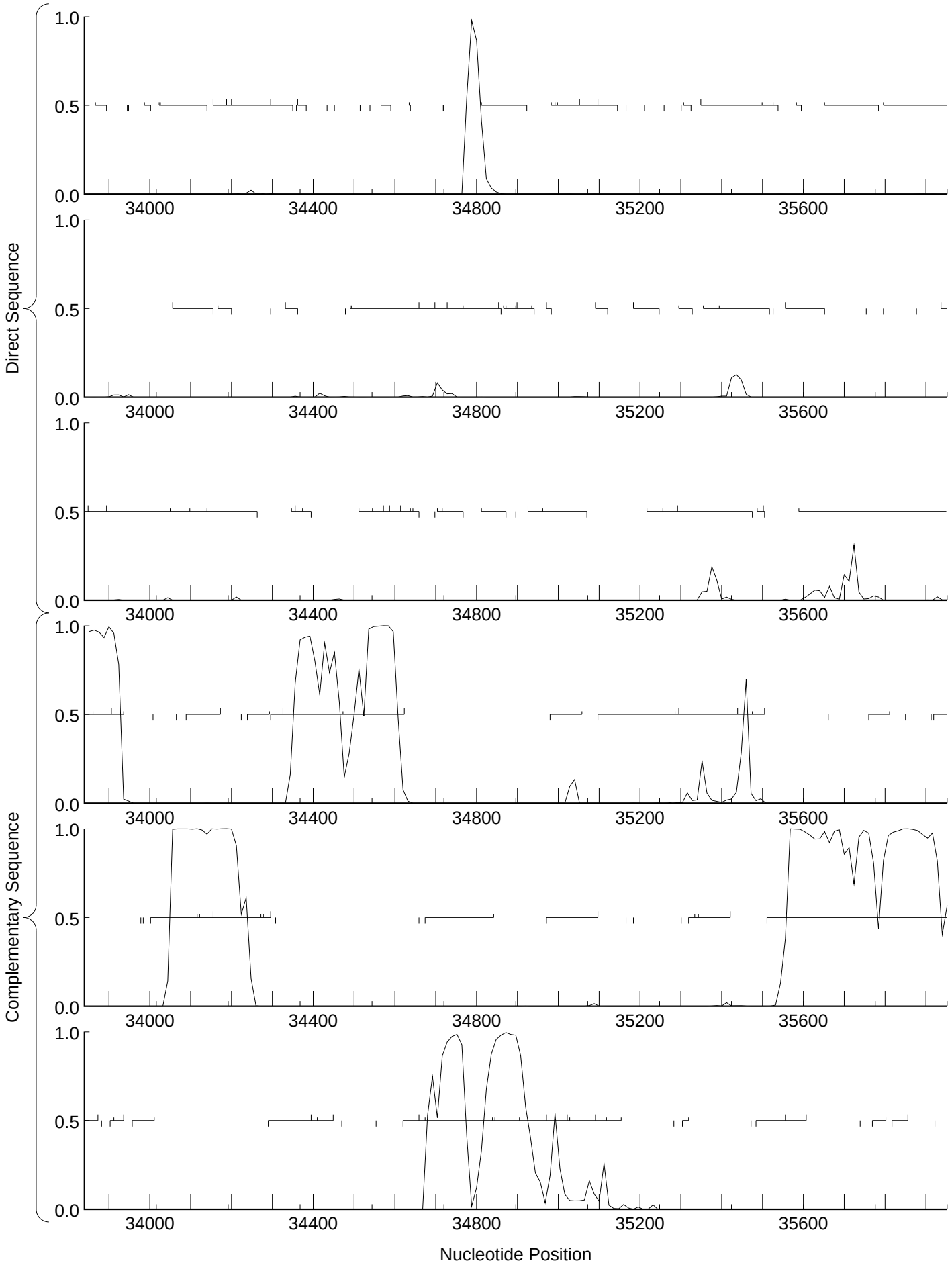
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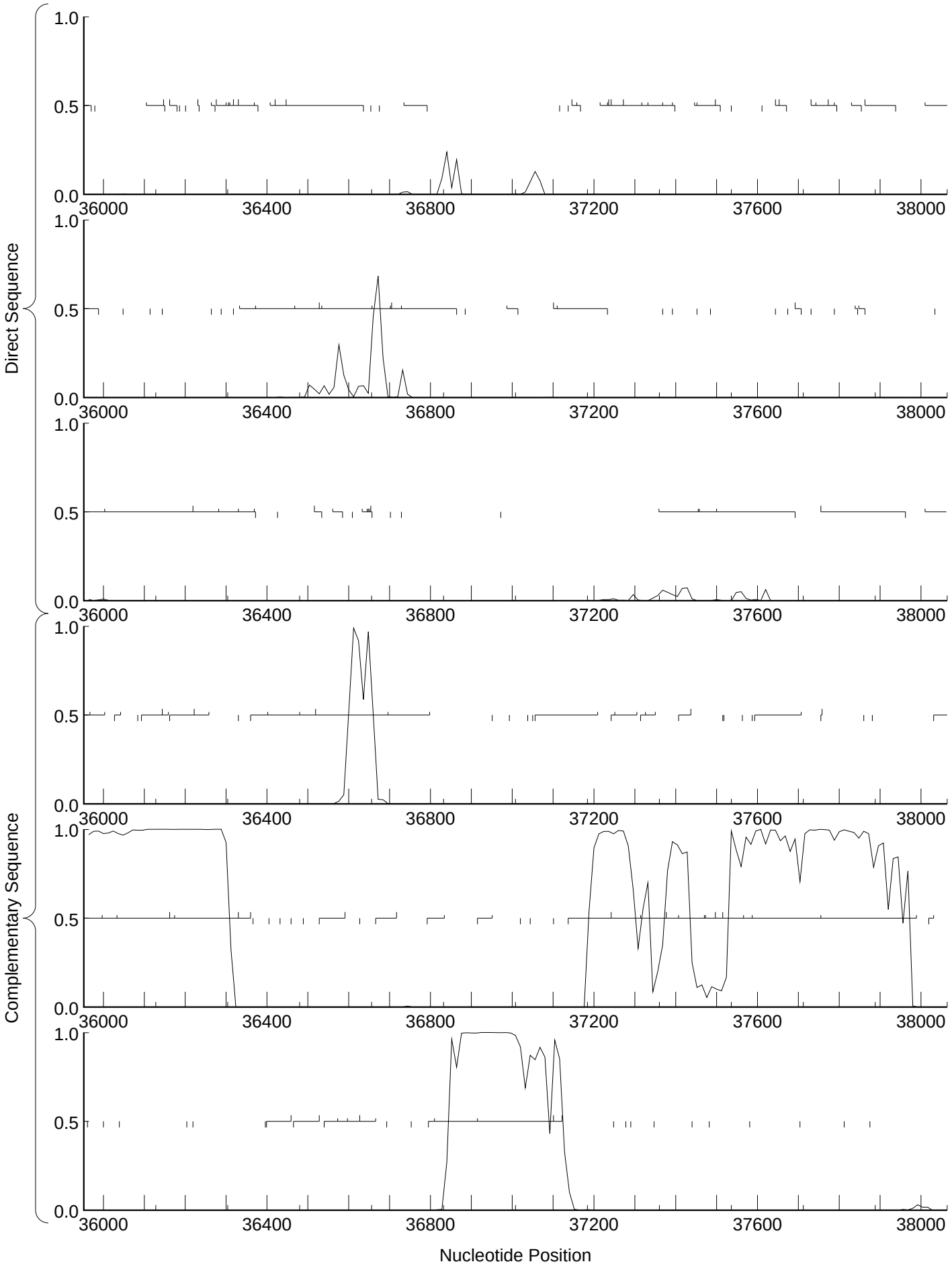
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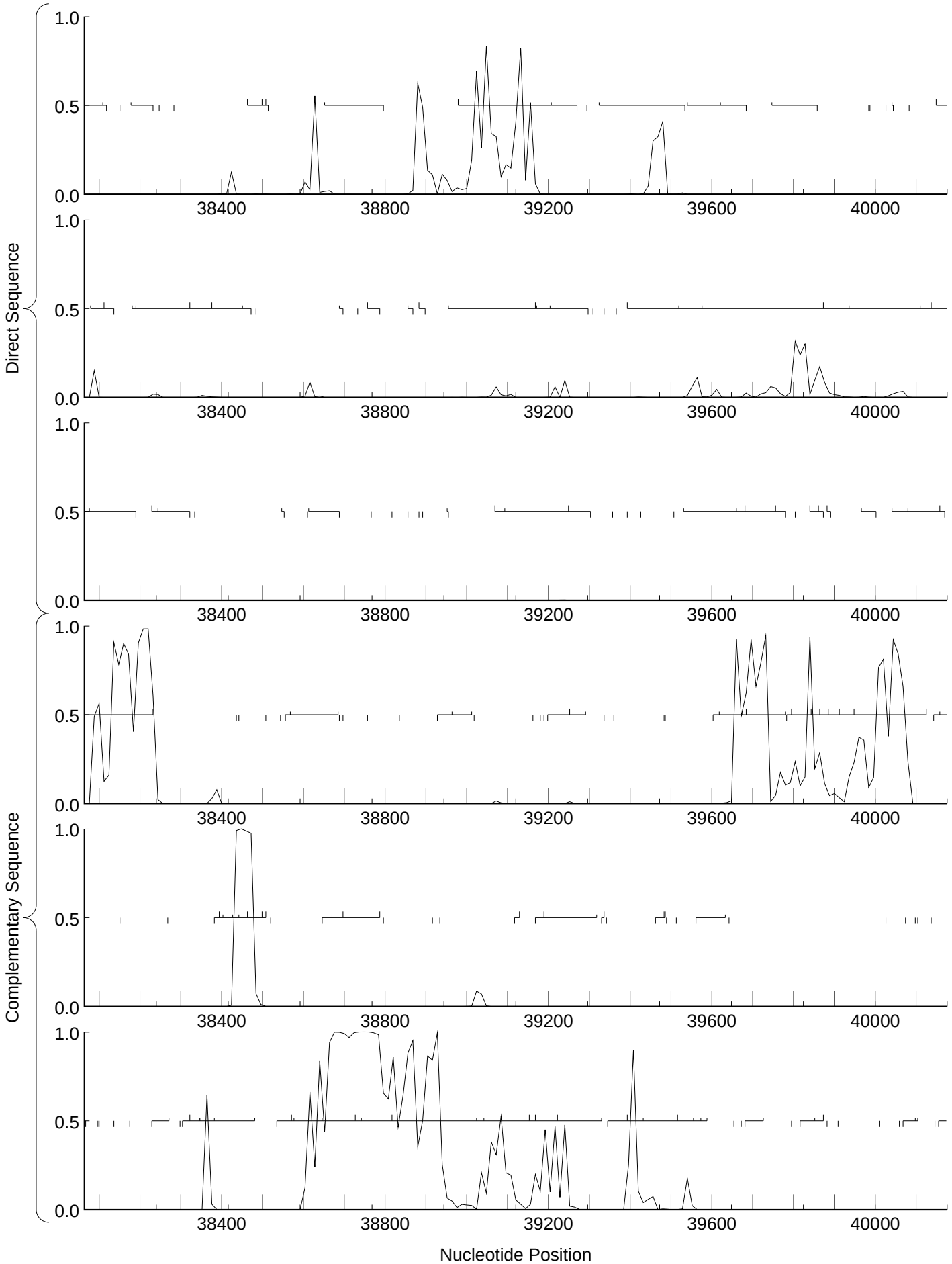
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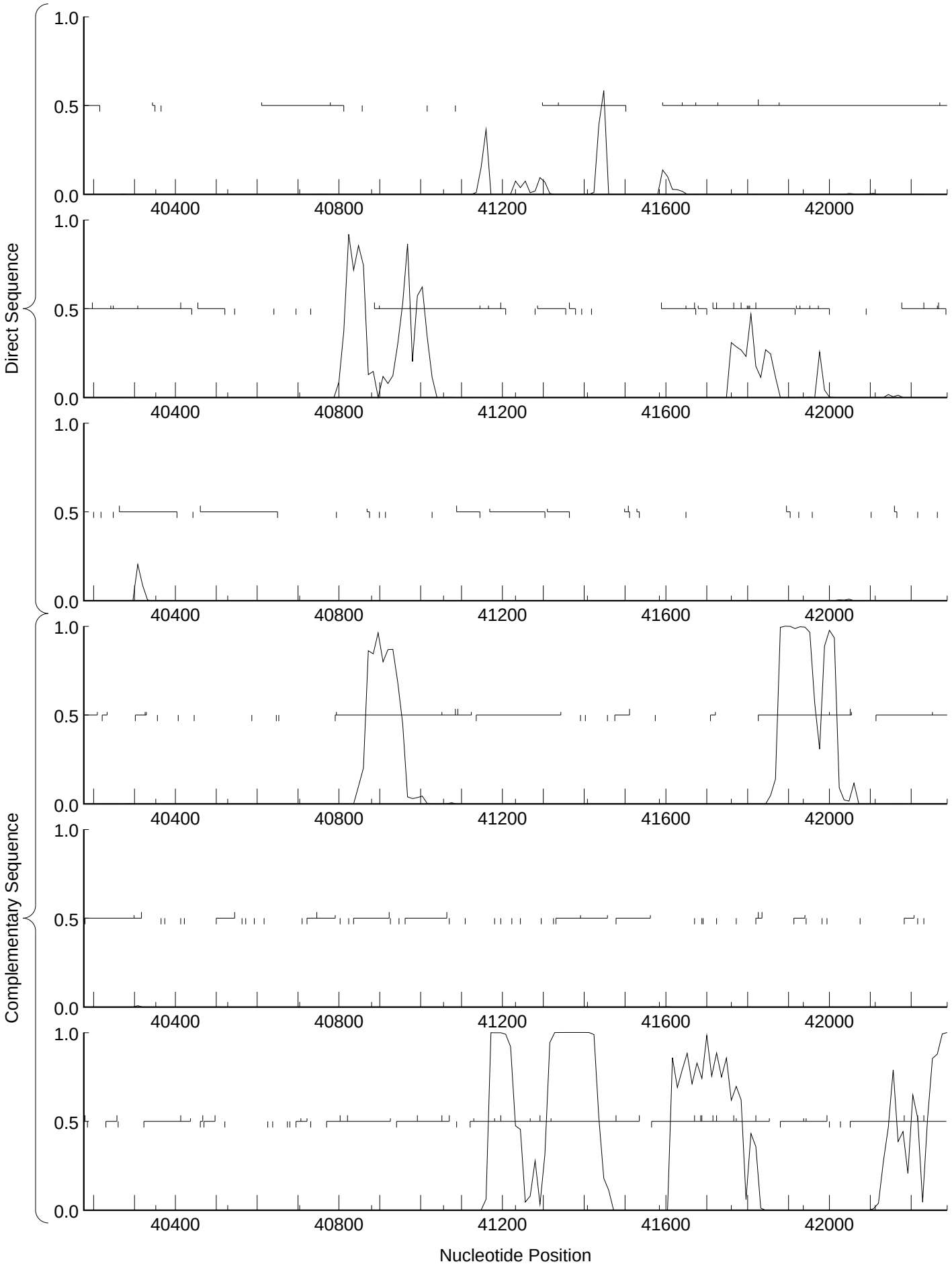
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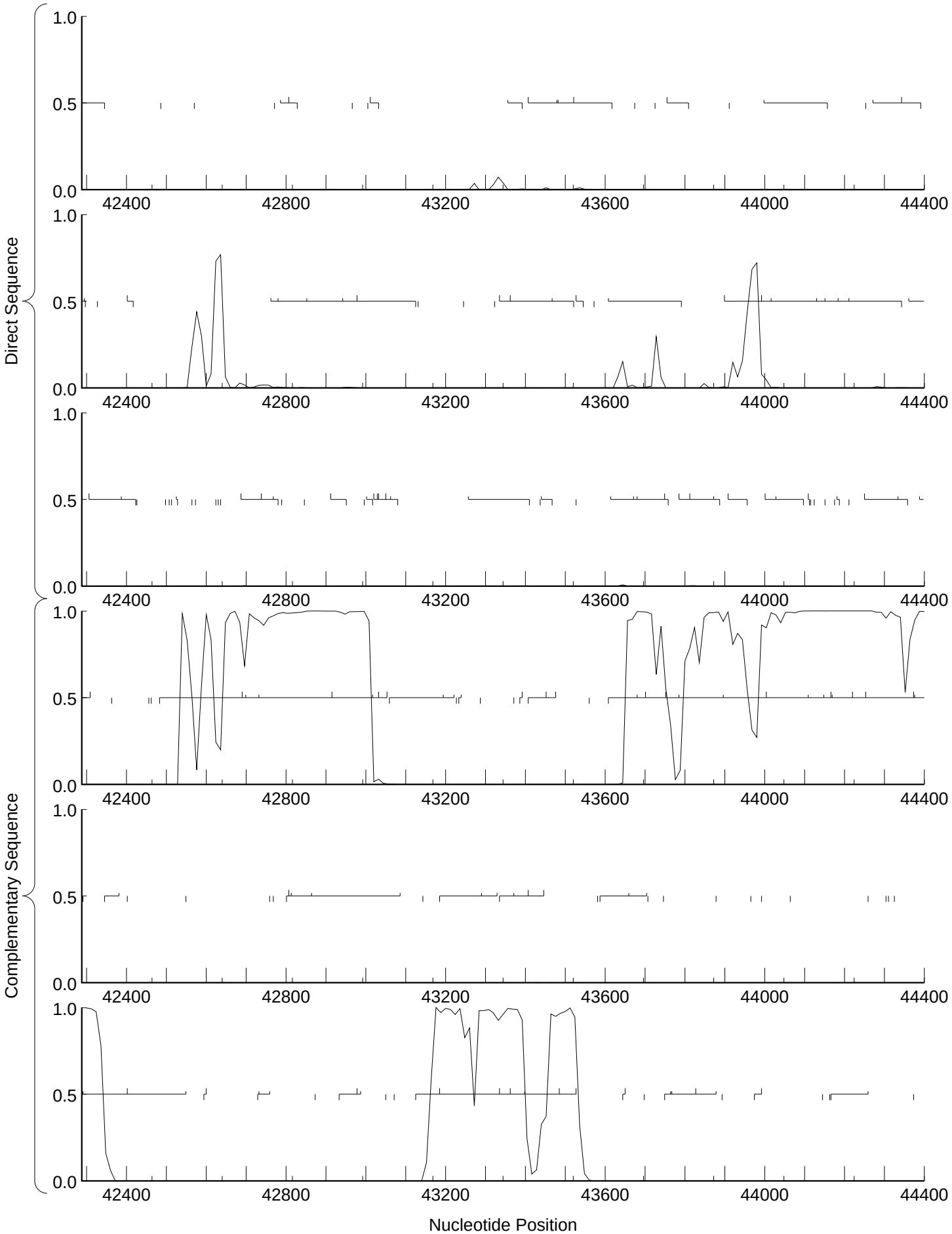
2 complete genome, 46596 bp including 10-base 3’ overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 20/24



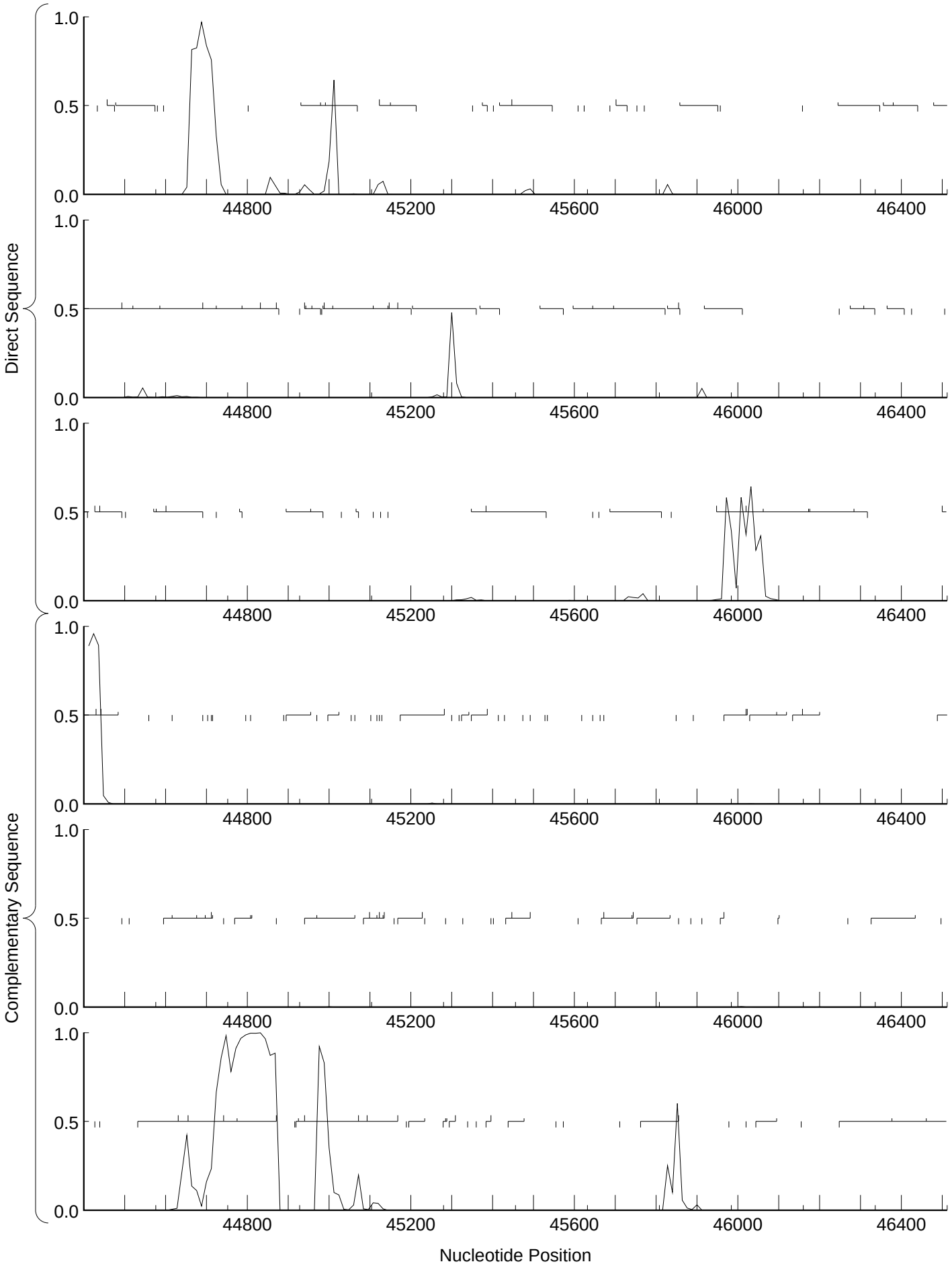
2 complete genome, 46596 bp including 10-base 3’ overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 21/24



2 complete genome, 46596 bp including 10-base 3’ overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 22/24



2 complete genome, 46596 bp including 10-base 3’ overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 23/24



2 complete genome, 46596 bp including 10-base 3’ overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 24/24

