GeneMark

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

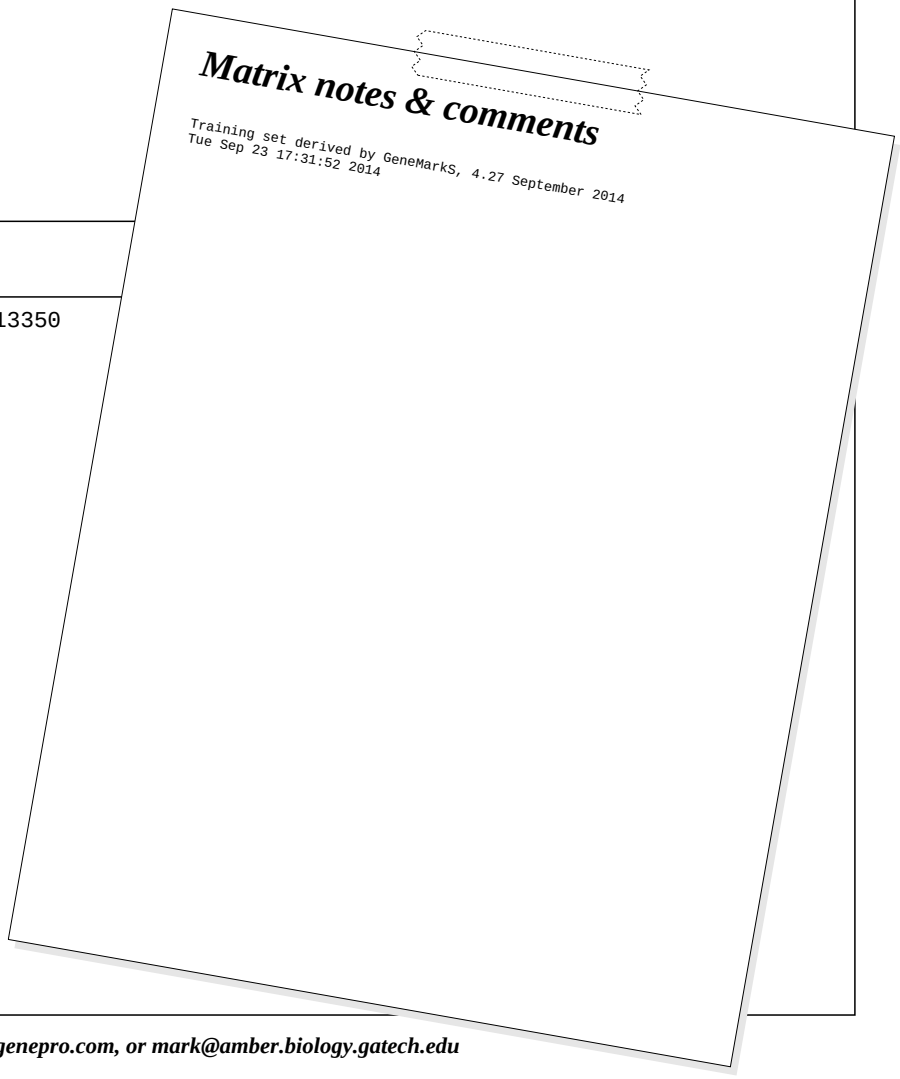
PROGRAM INFORMATION

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Sequence         :  Streptomyces phage Doxi13 complete sequence, 43696 bp including 9-base 3' overhang (CGCCGCC
Analysis Date    :  3/28/24 at 12:59:01
Pages            :  22
Sequence Length  :  43696 bp
GC Content       :   60.88%

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,
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MATRIX INFORMATION

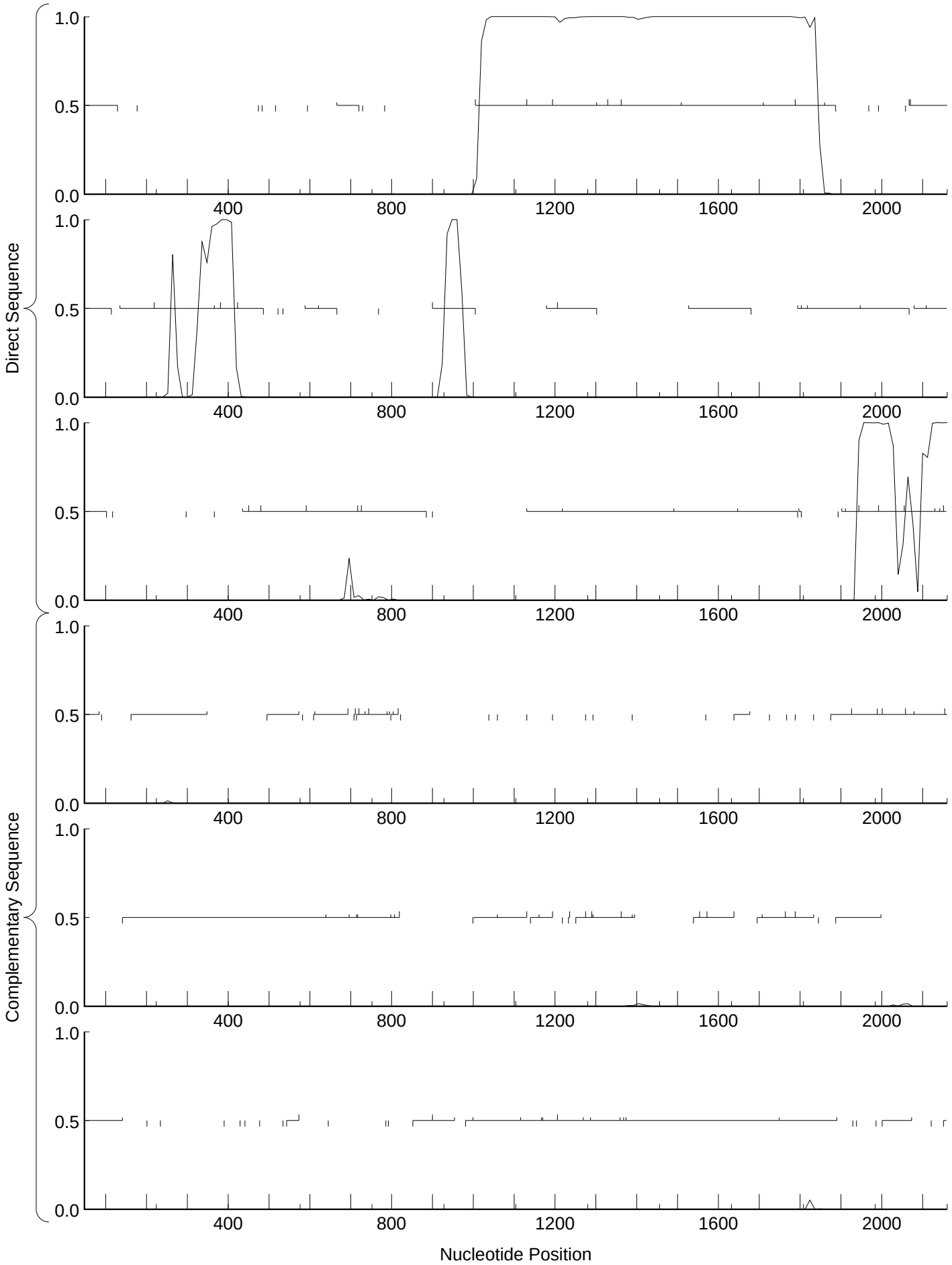
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Matrix     :  Streptomyces_griseus_NBRC_13350
Author     :  -

Order      :  4
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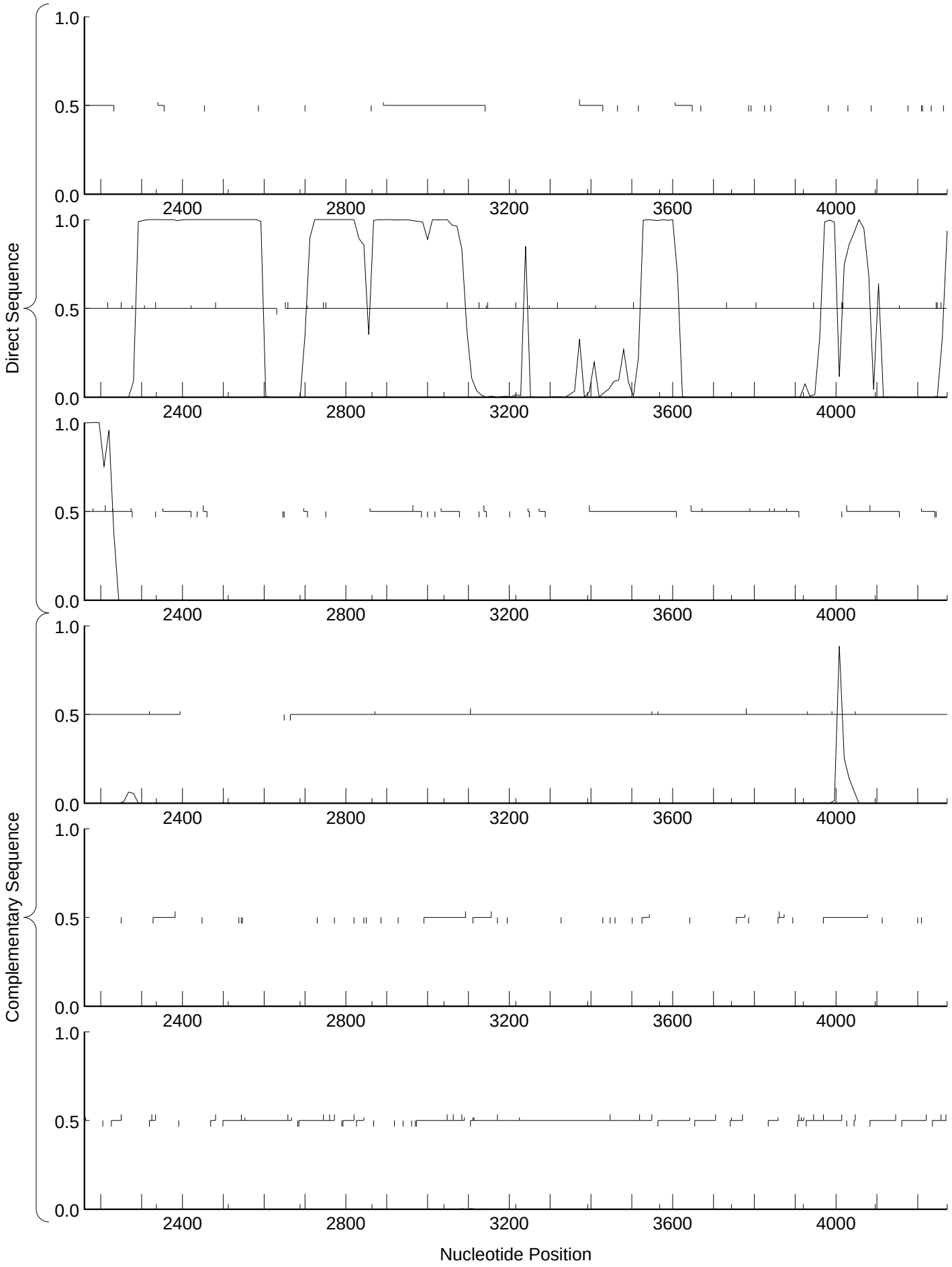

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

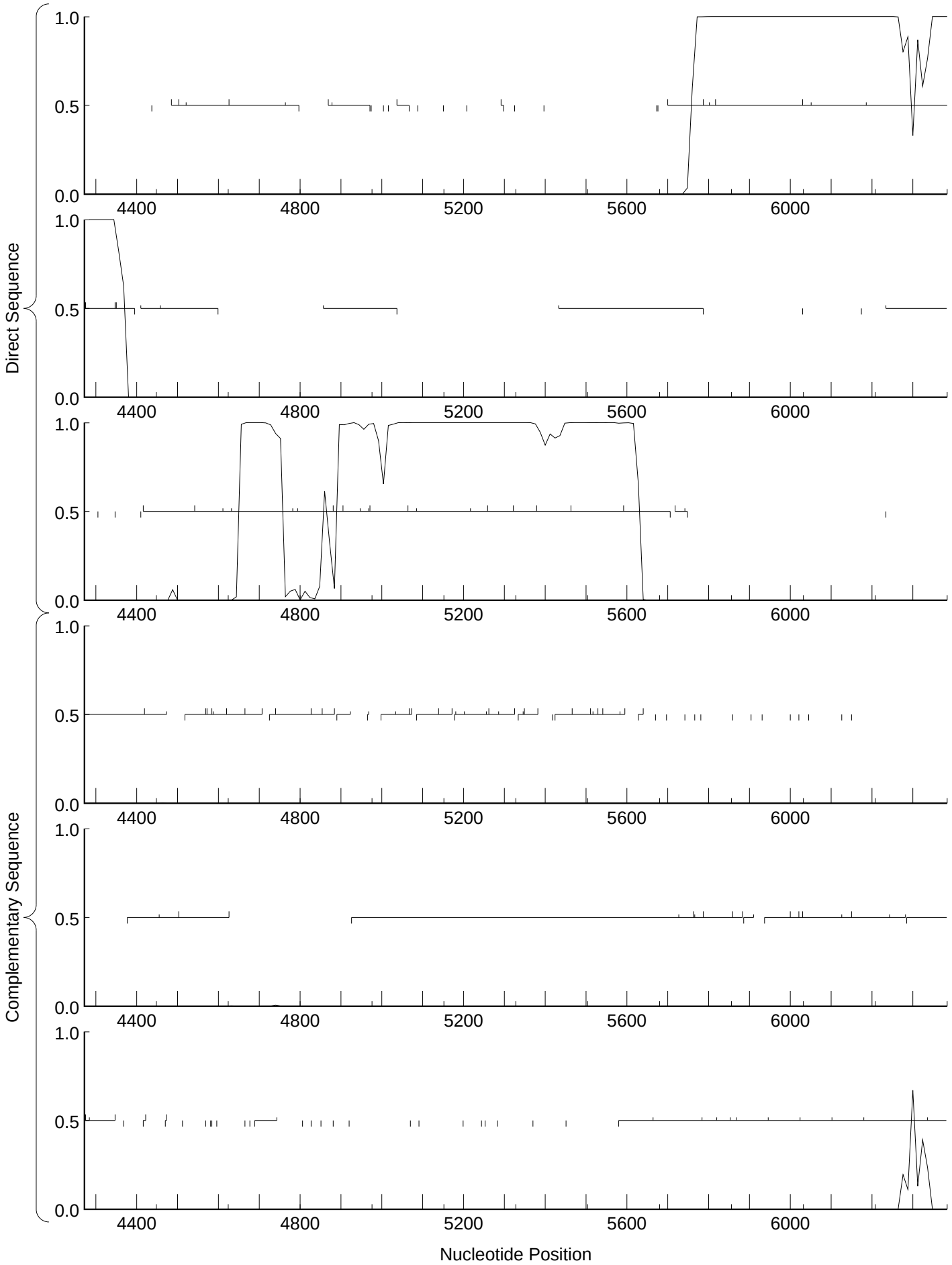
xi13 complete sequence, 43696 bp including 9-base 3’ overhang (CGCCGCCCT), Cluster BI2, Order 4, Window 96, Step 12, 2/22

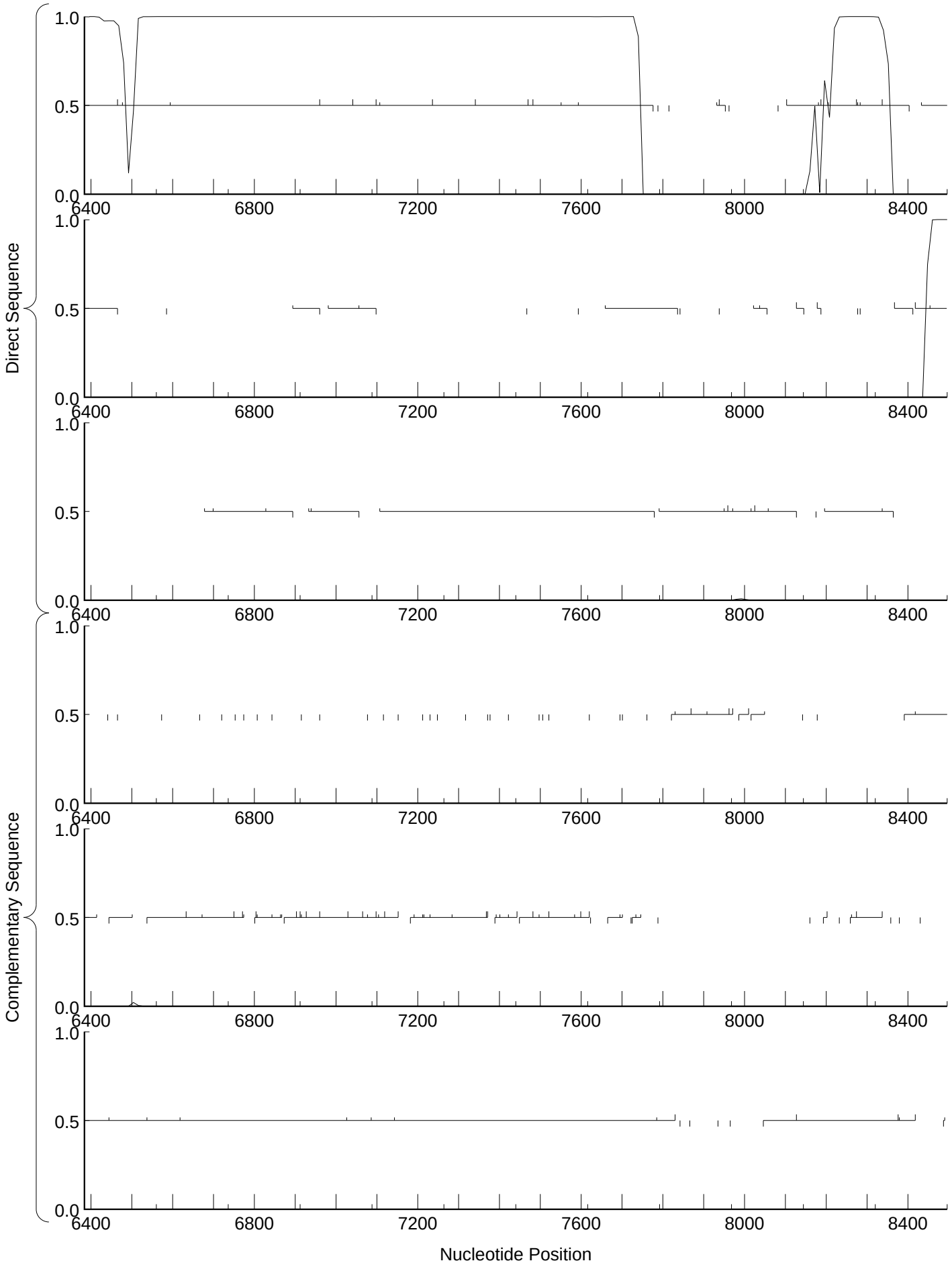


xi13 complete sequence, 43696 bp including 9-base 3' overhang (CGCCGCCCT), Cluster BI2, Order 4, Window 96, Step 12, 3/22

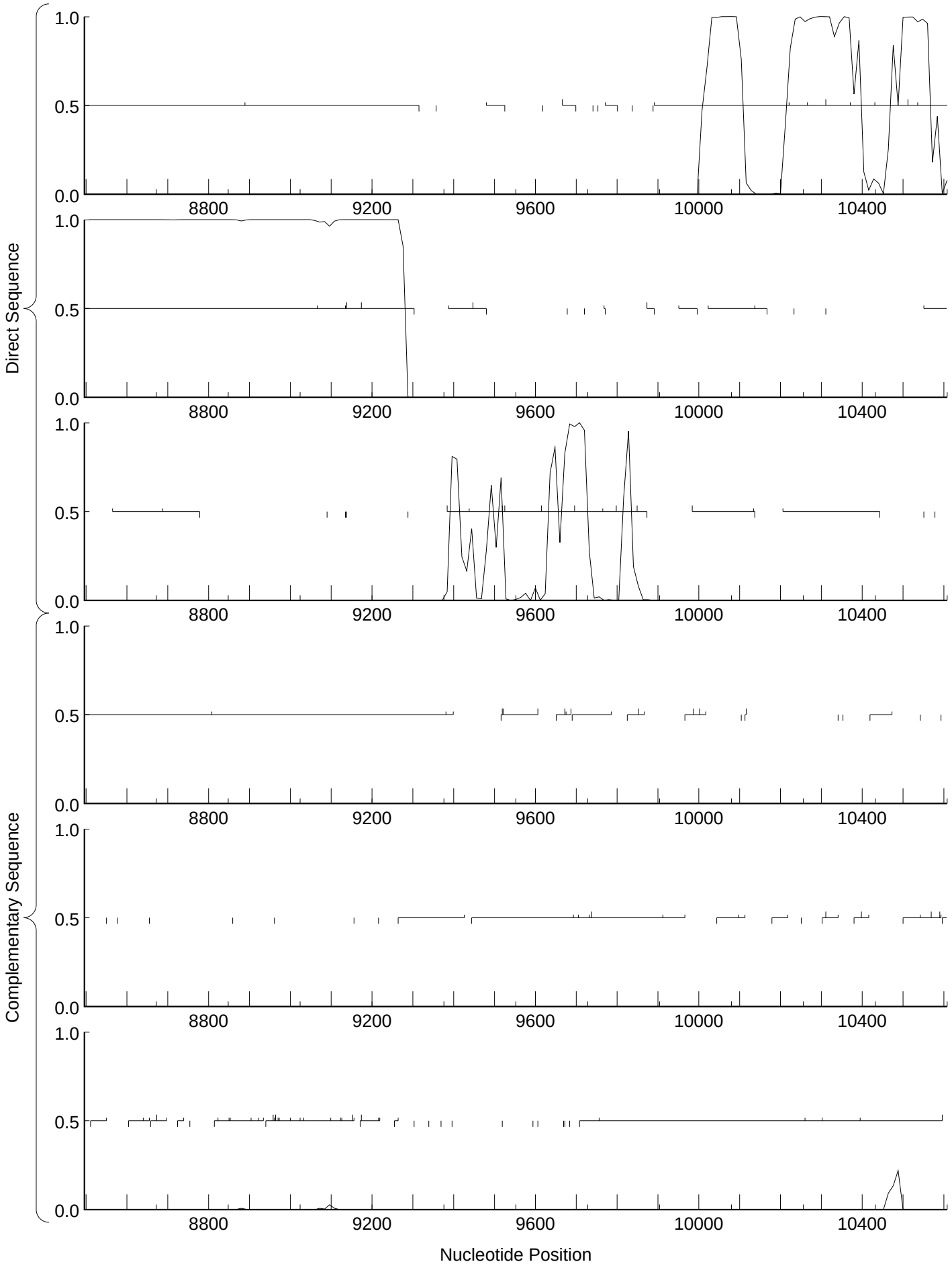


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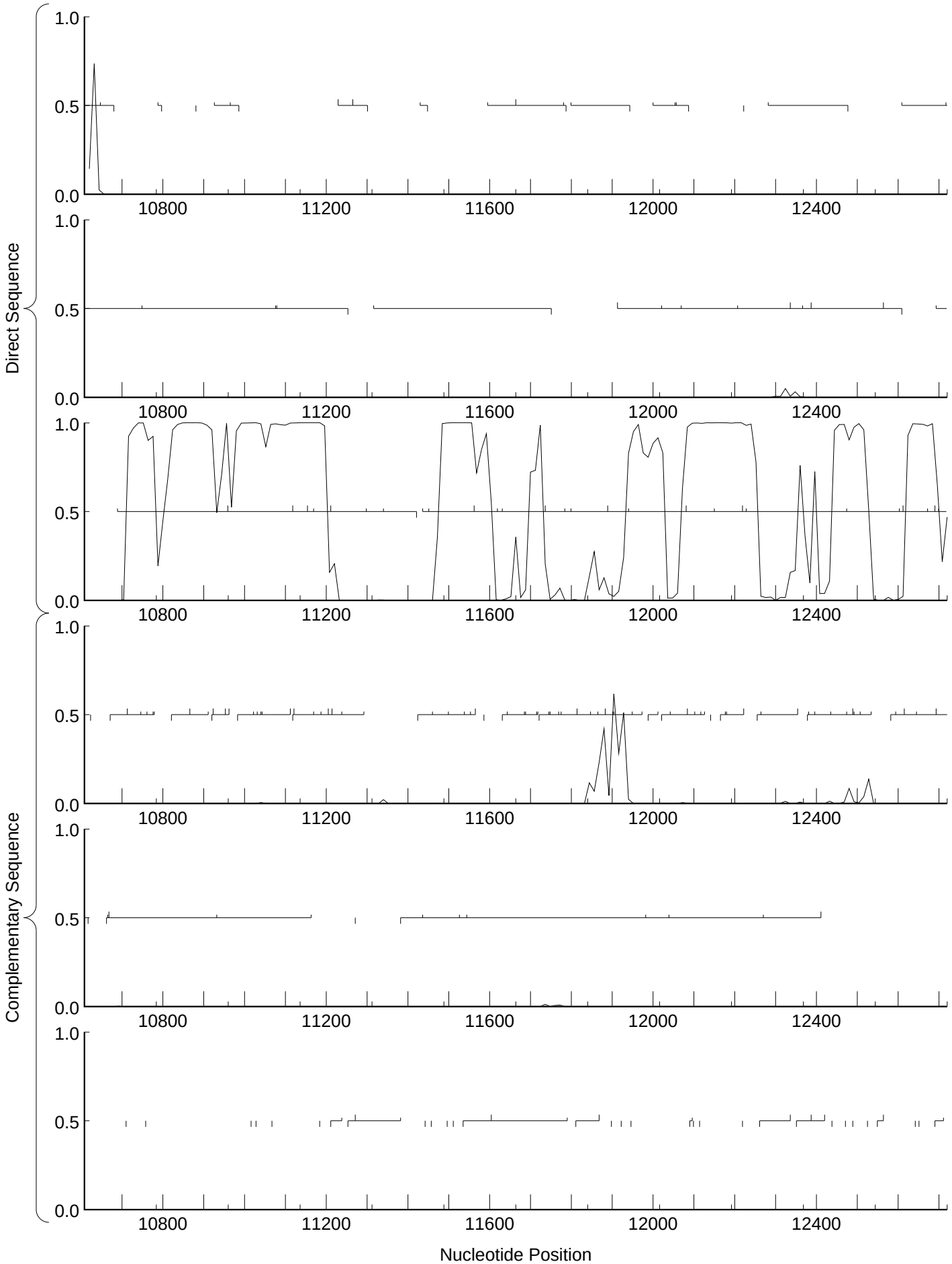



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
6400
6800
7200
7600
8000
8400
Nucleotide Position

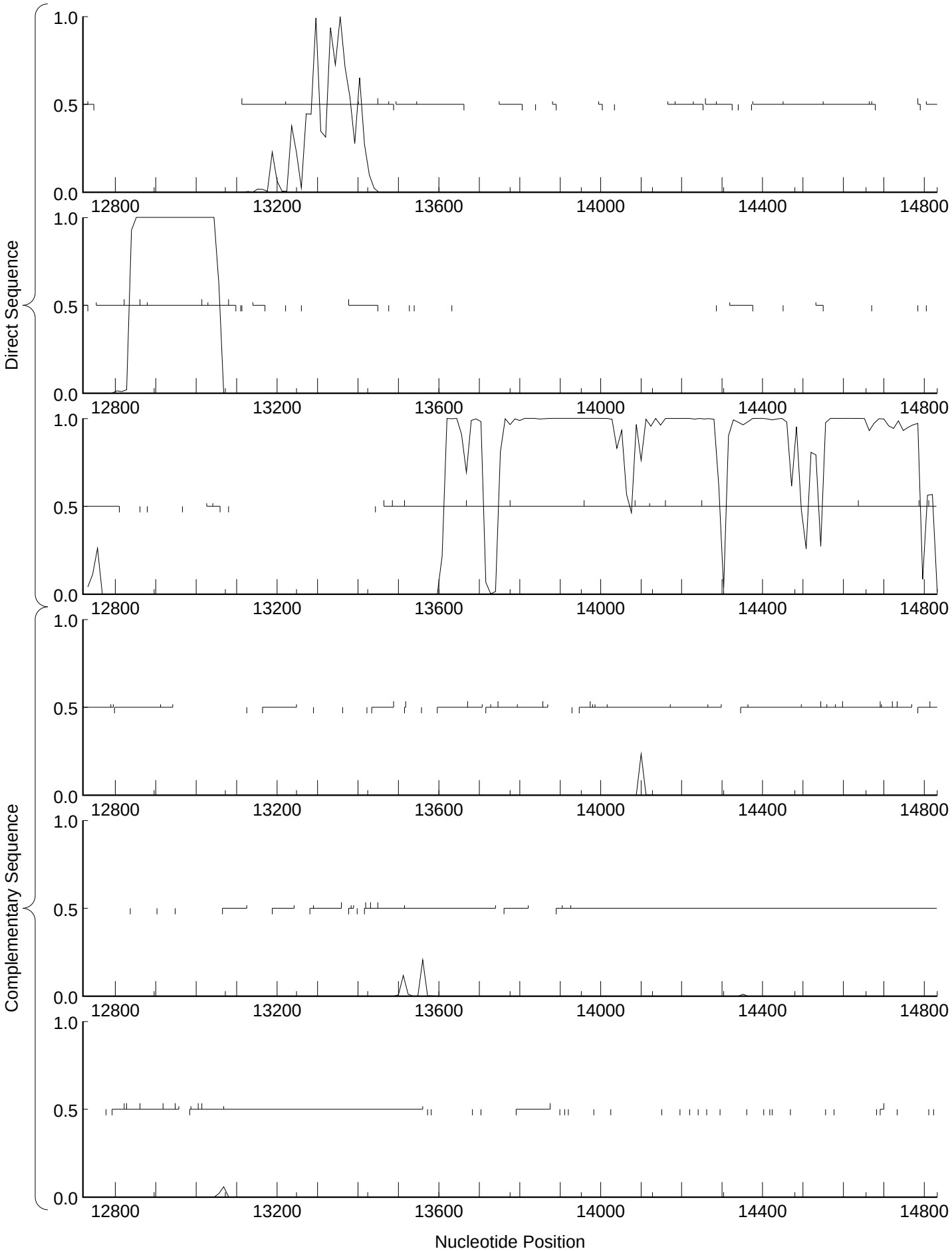
xi13 complete sequence, 43696 bp including 9-base 3’ overhang (CGCCGCCCT), Cluster BI2, Order 4, Window 96, Step 12, 6/22



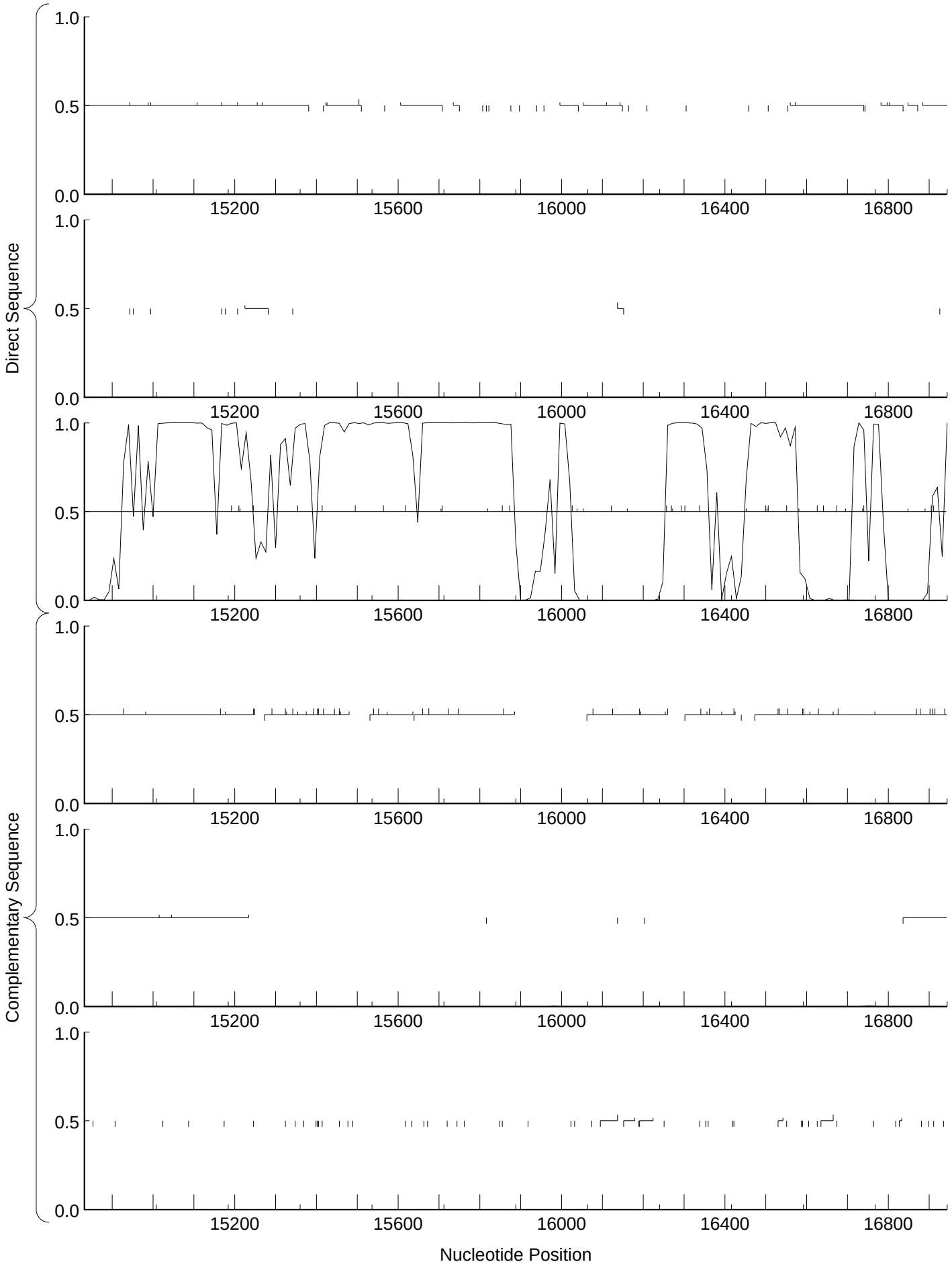
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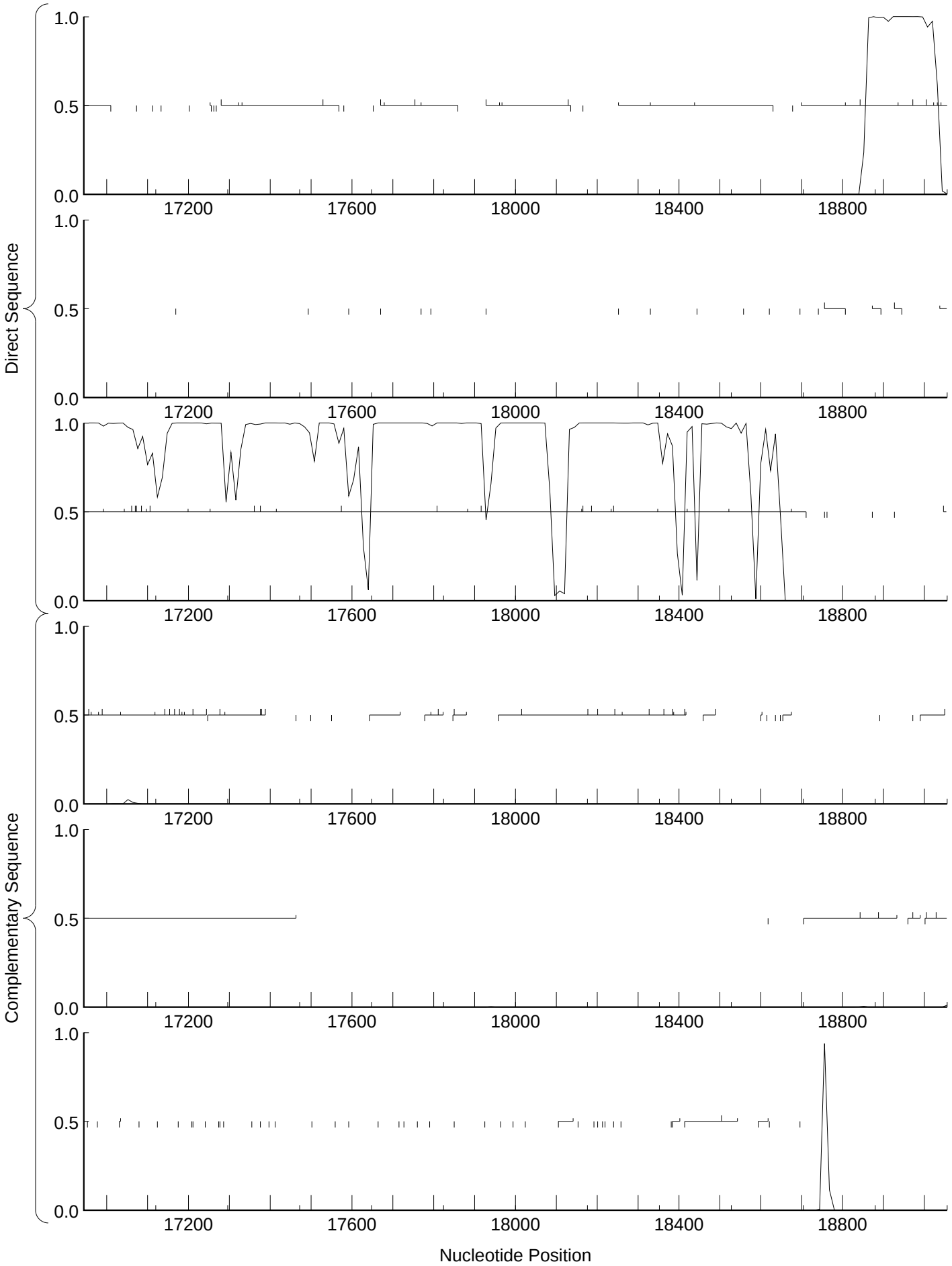
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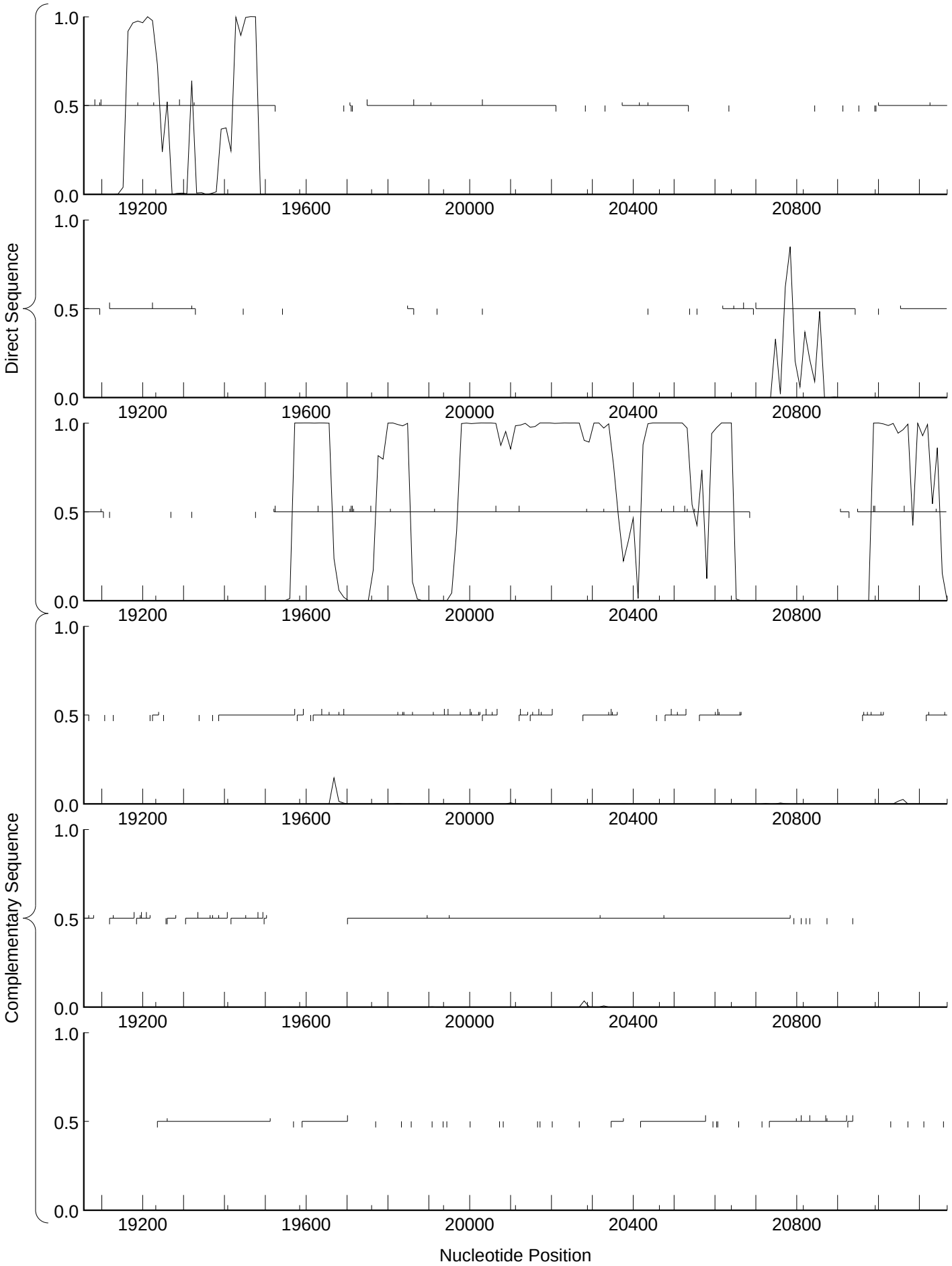
xi13 complete sequence, 43696 bp including 9-base 3’ overhang (CGCCGCCCT), Cluster BI2, Order 4, Window 96, Step 12, 9/22



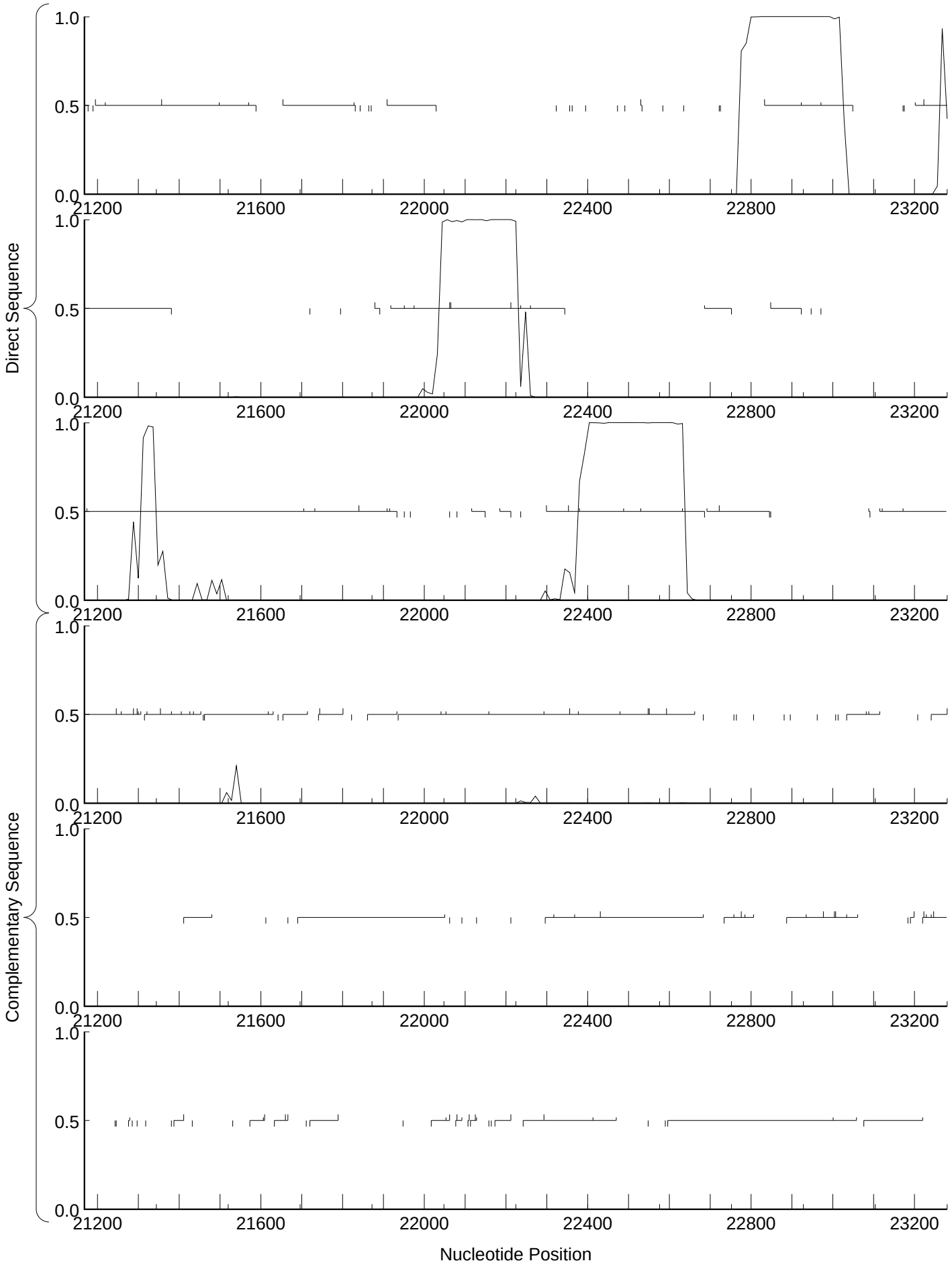
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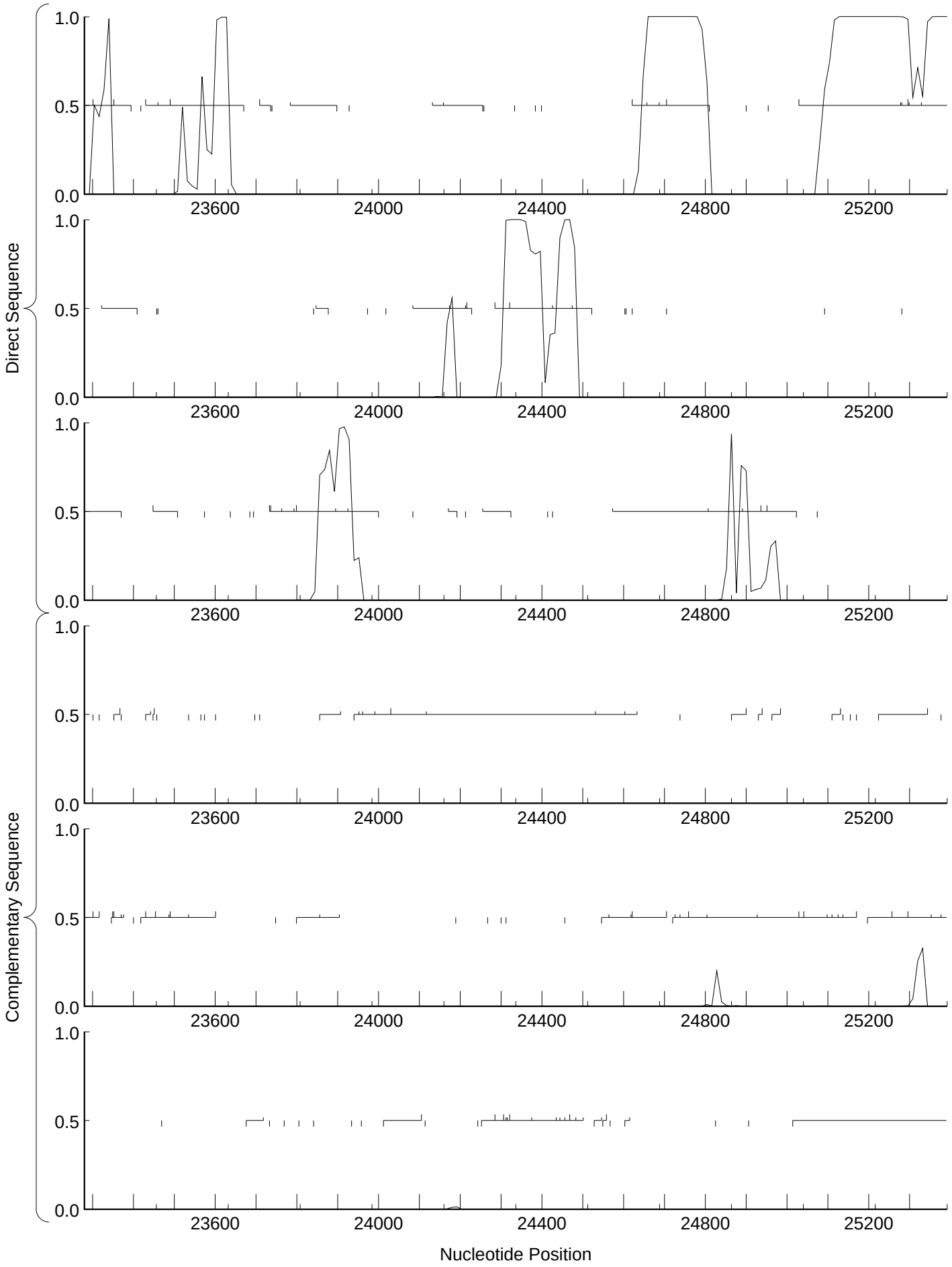
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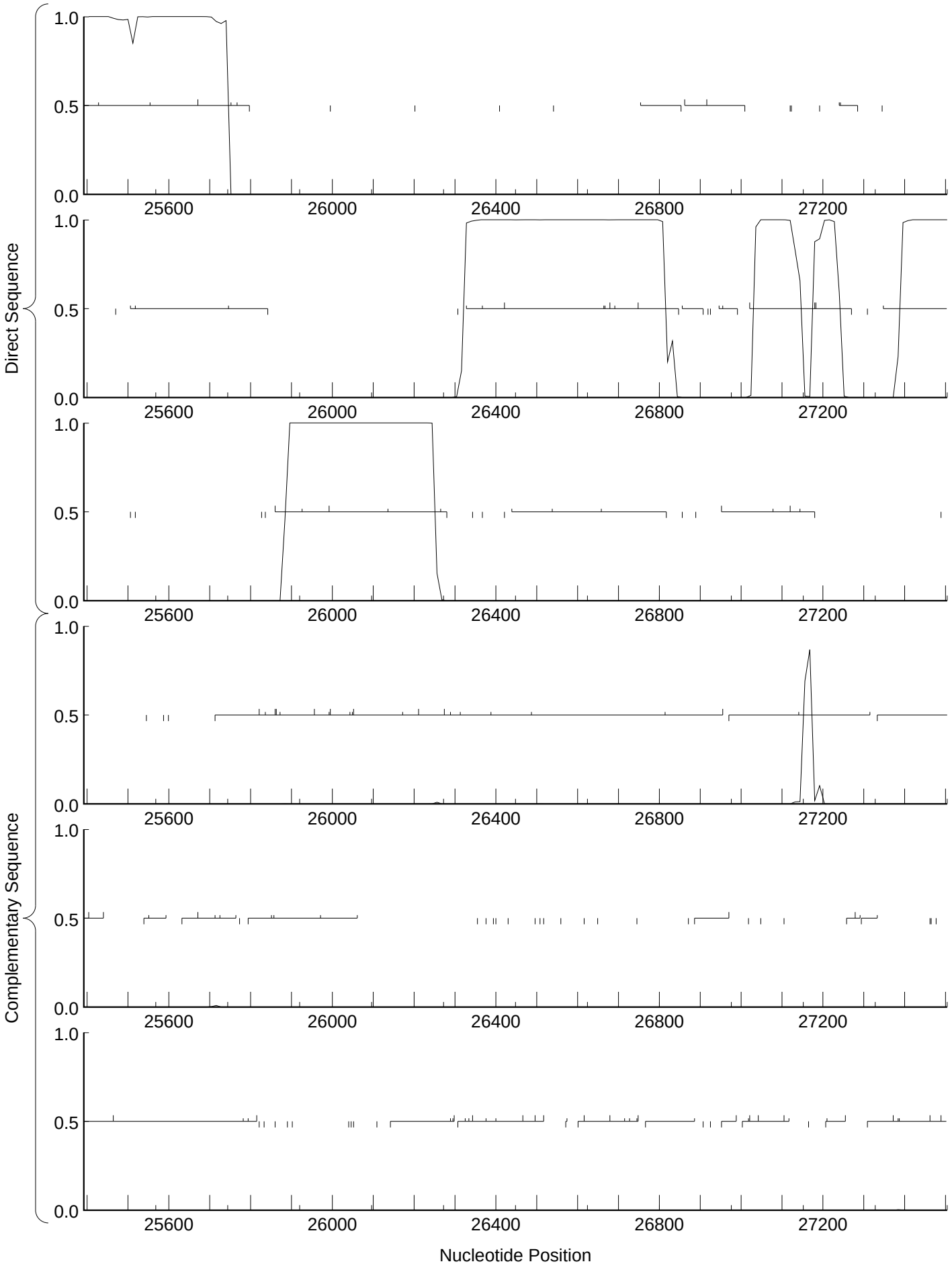
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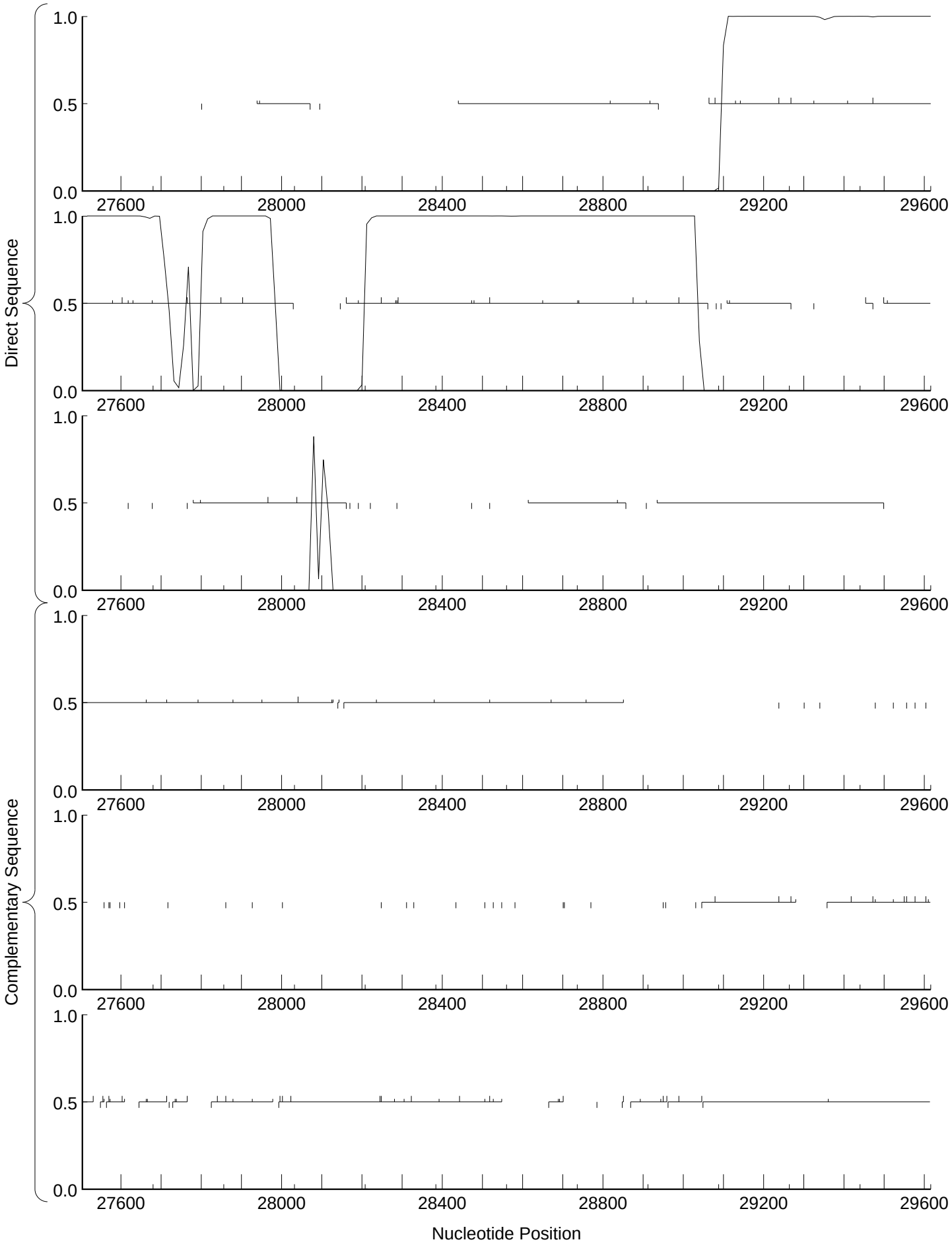
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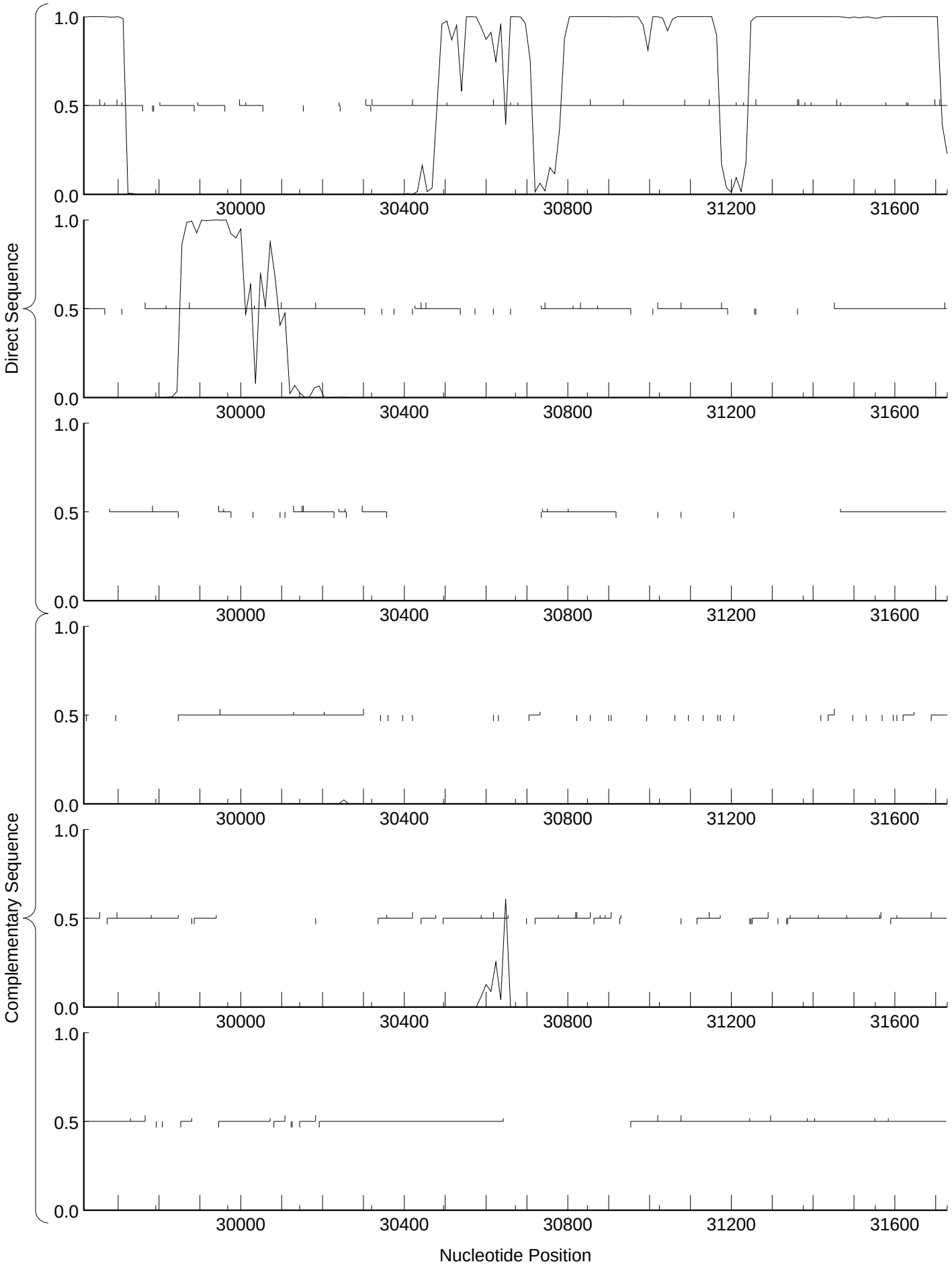
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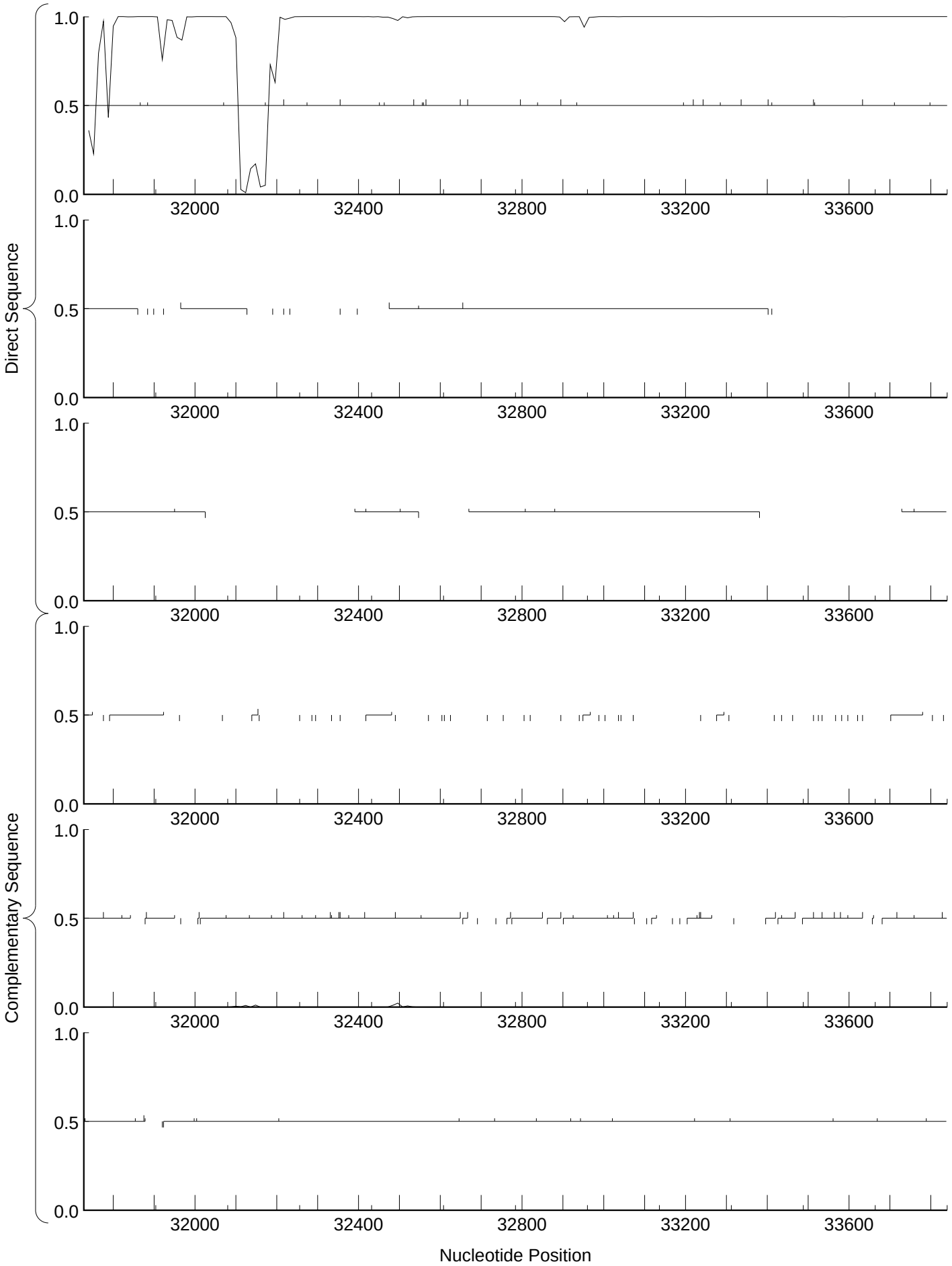
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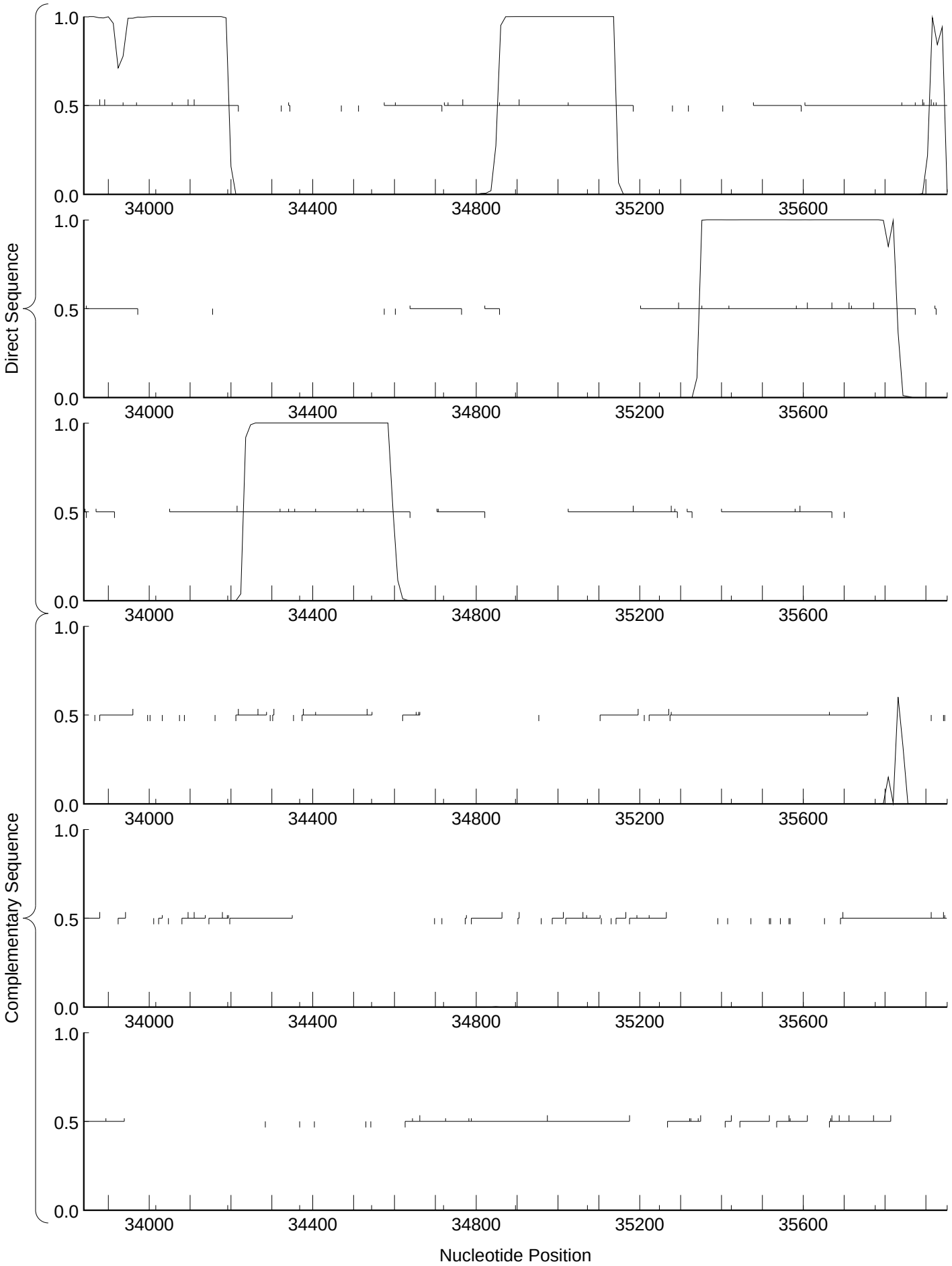
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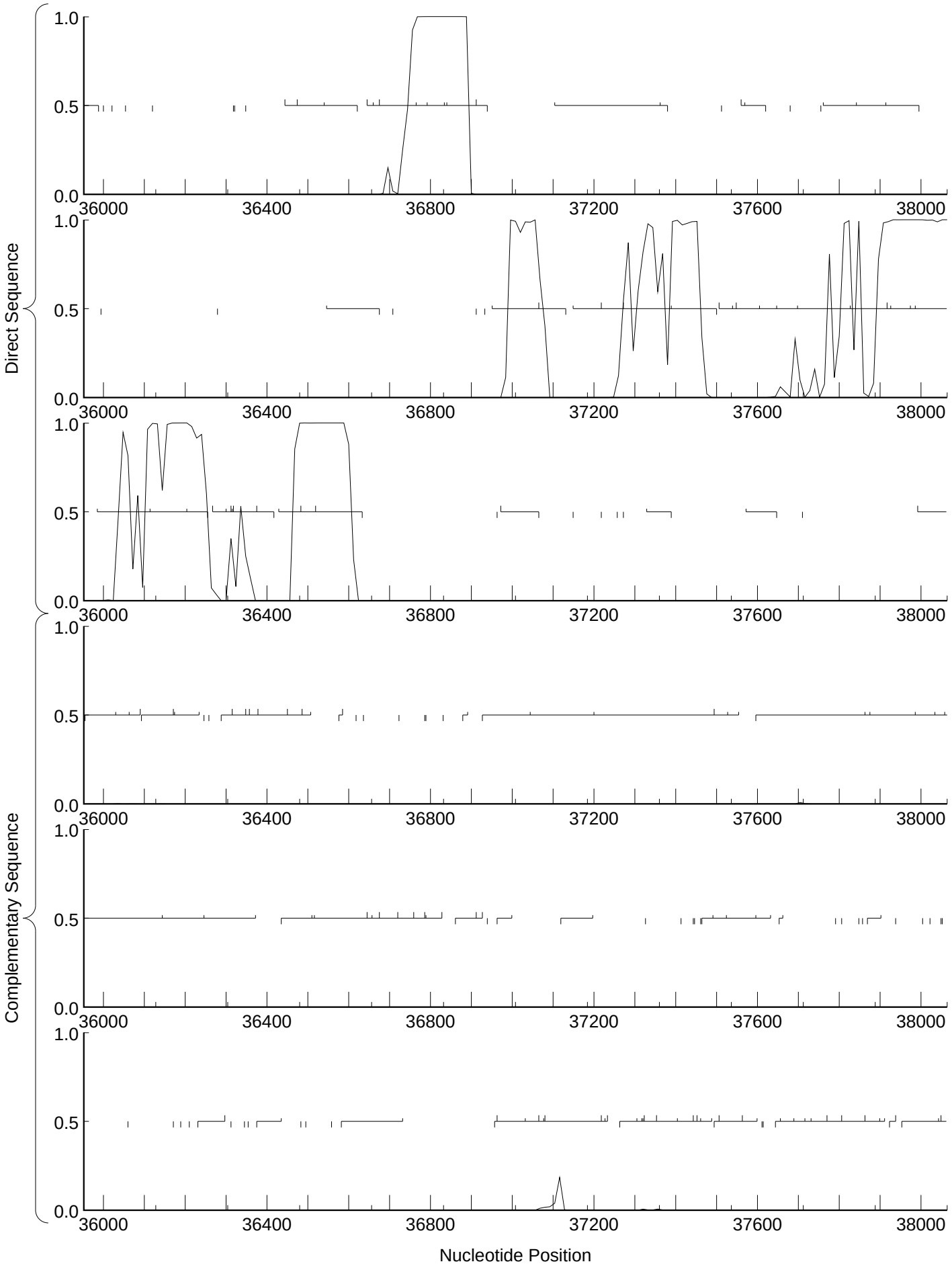
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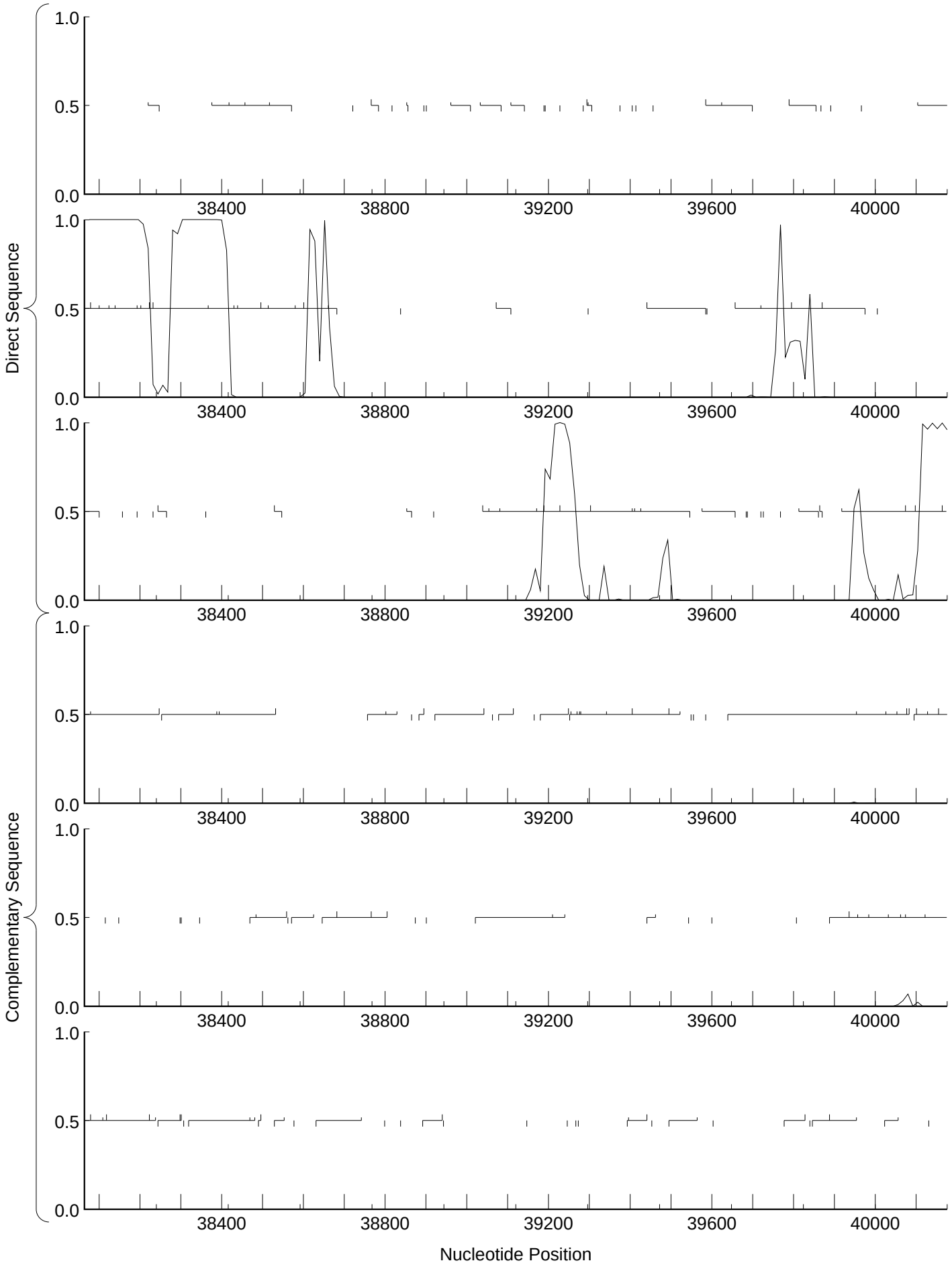
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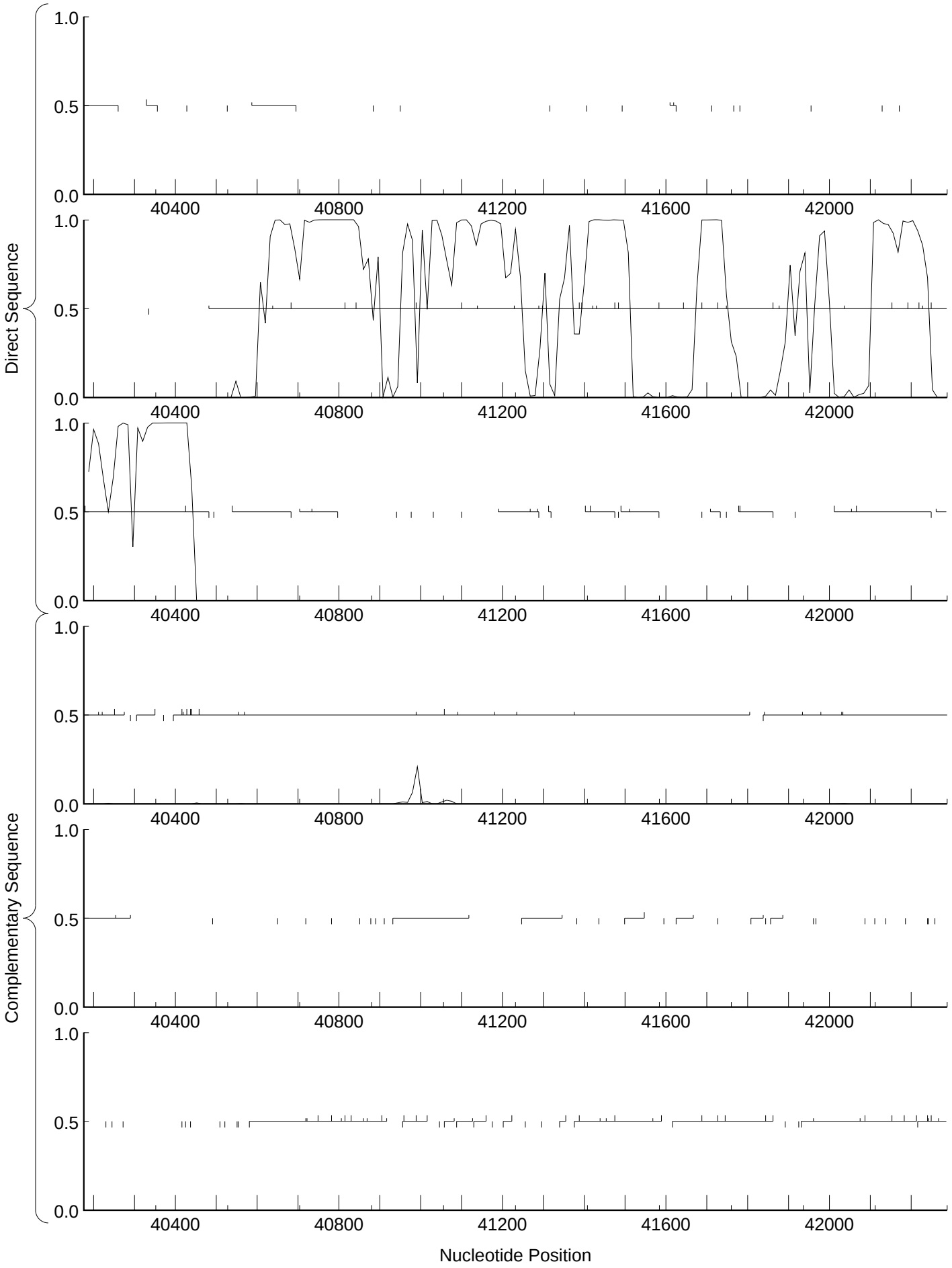
13 complete sequence, 43696 bp including 9-base 3' overhang (CGCCGCCCT), Cluster BI2, Order 4, Window 96, Step 12, 19/22



13 complete sequence, 43696 bp including 9-base 3' overhang (CGCCGCCCT), Cluster BI2, Order 4, Window 96, Step 12, 20/22



13 complete sequence, 43696 bp including 9-base 3' overhang (CGCCGCCCT), Cluster BI2, Order 4, Window 96, Step 12, 21/22



13 complete sequence, 43696 bp including 9-base 3’ overhang (CGCCGCCCT), Cluster BI2, Order 4, Window 96, Step 12, 22/22

