GeneMark

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

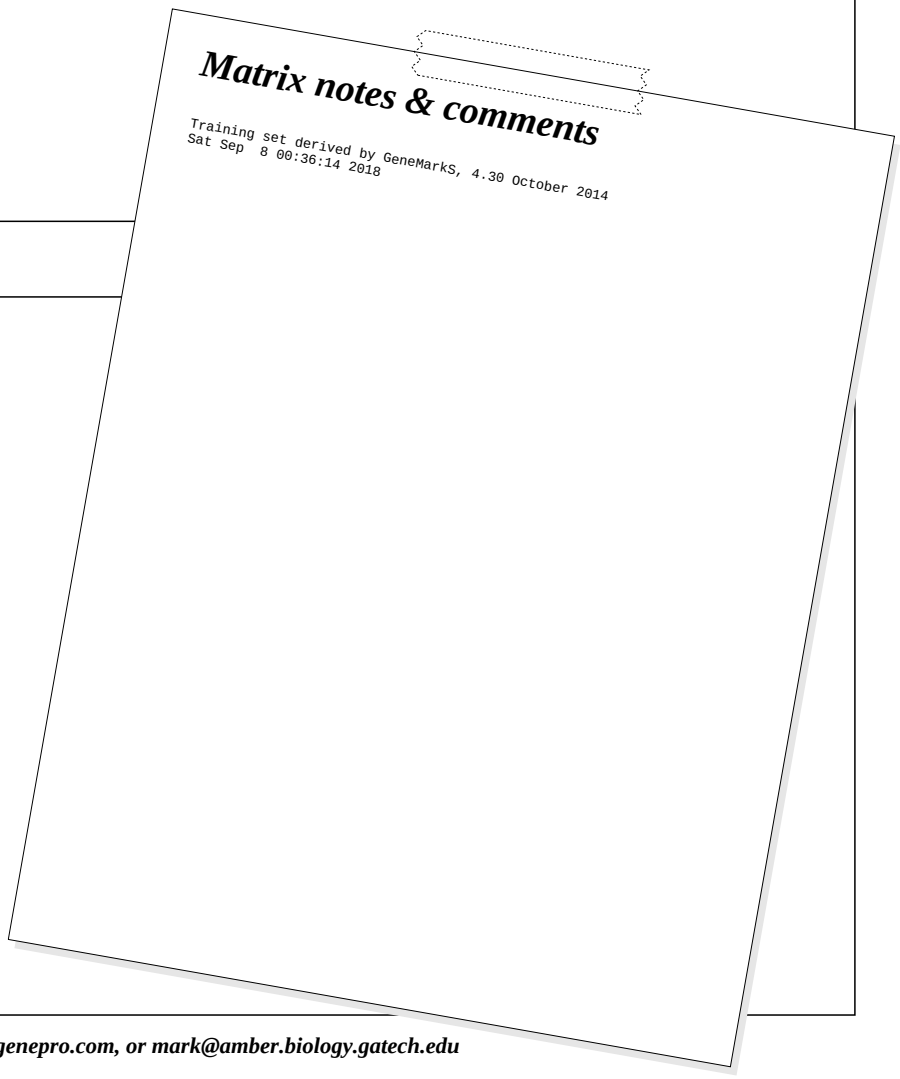
PROGRAM INFORMATION

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Sequence         :  Gordonia phage Switzerland complete sequence, 52206 bp including 10-base 3' overhang (CGGGT
Analysis Date    :  3/28/24 at 20:06:27
Pages            :  26
Sequence Length  :  52206 bp
GC Content       :   62.01%

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

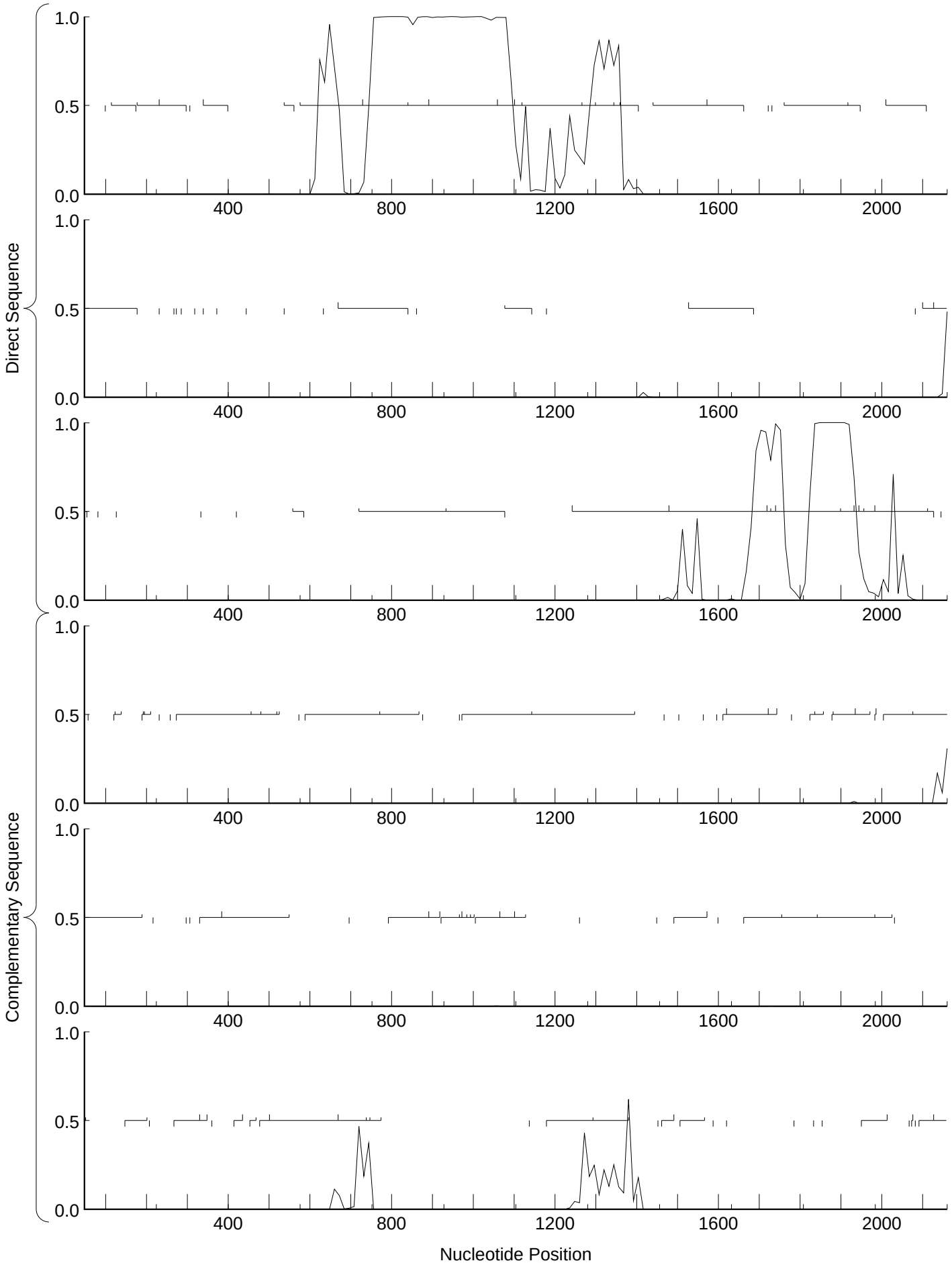
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Matrix     :  -
Author     :  -

Order      :  4
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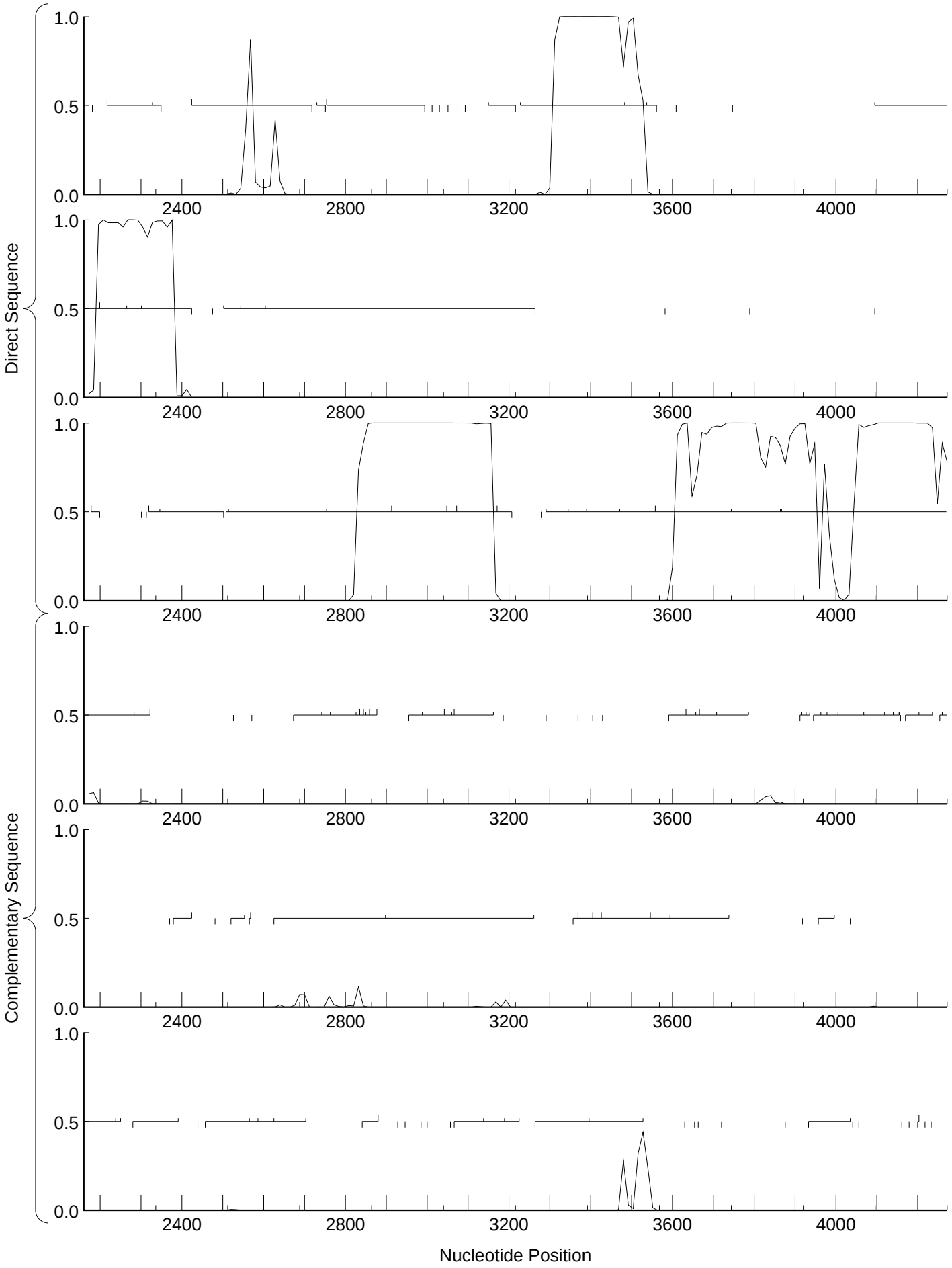

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

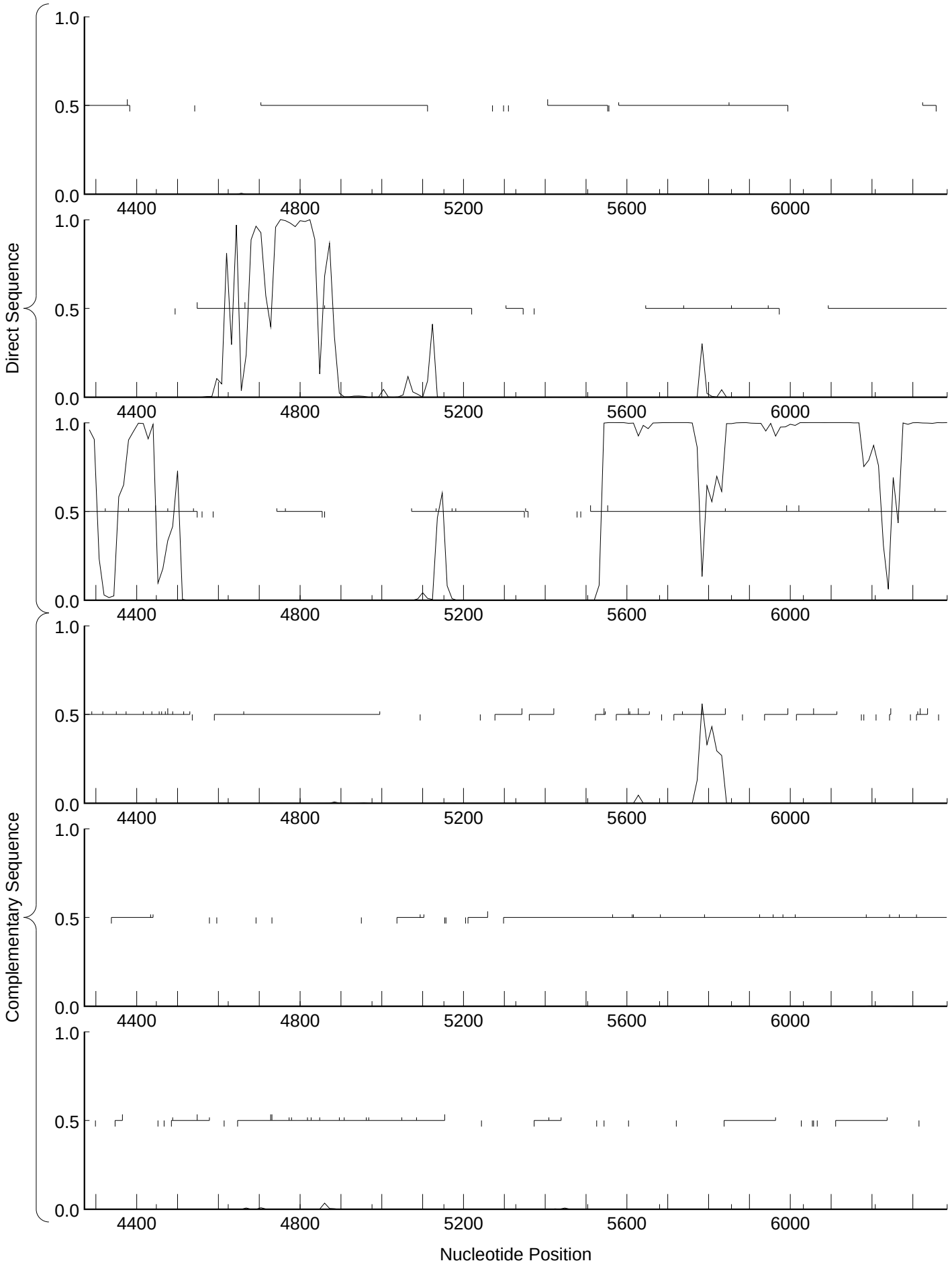
complete sequence, 52206 bp including 10-base 3’ overhang (CGGGTGGTTA), Cluster A15, Order 4, Window 96, Step 12, 2/26



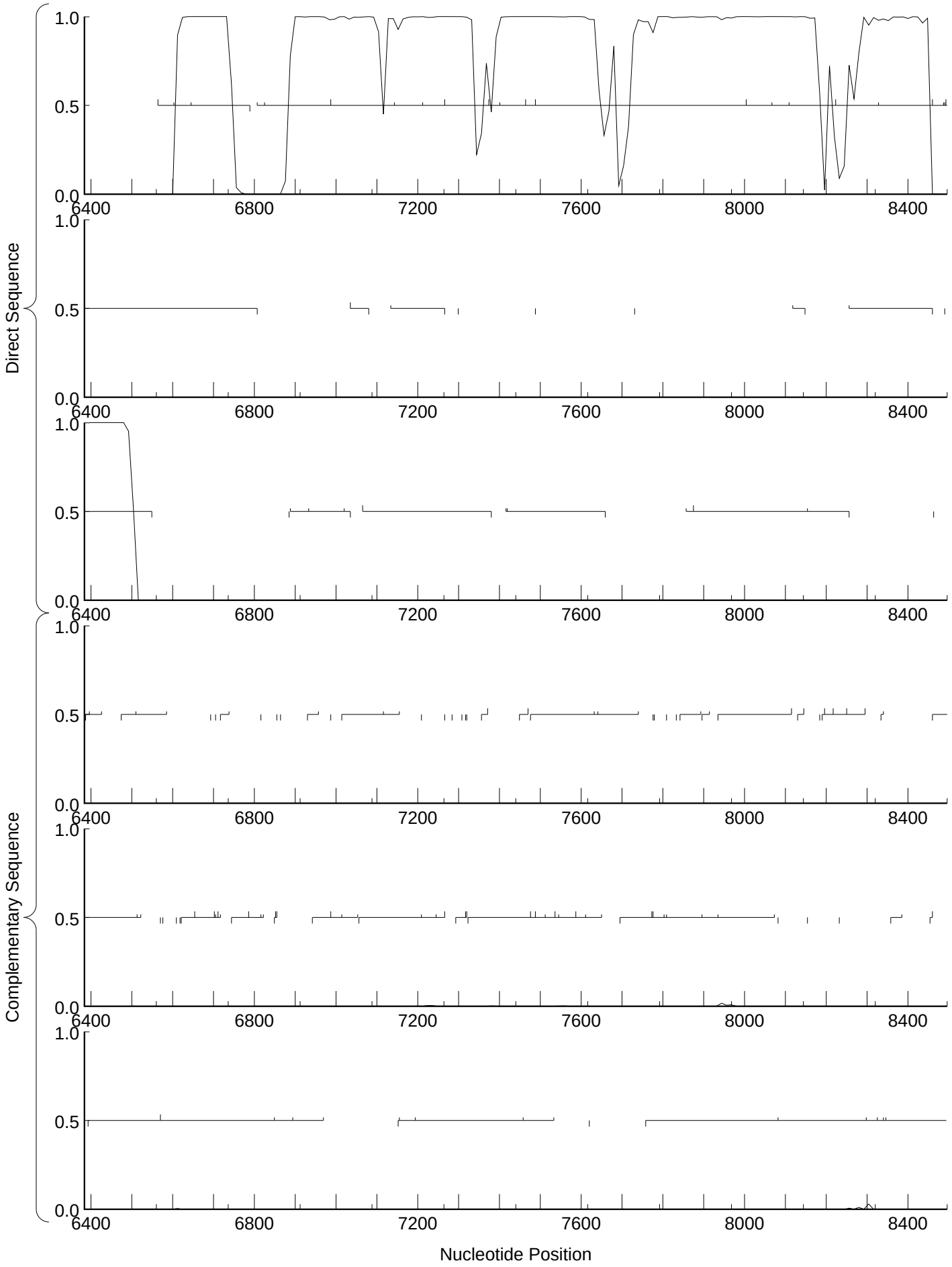
complete sequence, 52206 bp including 10-base 3’ overhang (CGGGTGGTTA), Cluster A15, Order 4, Window 96, Step 12, 3/26



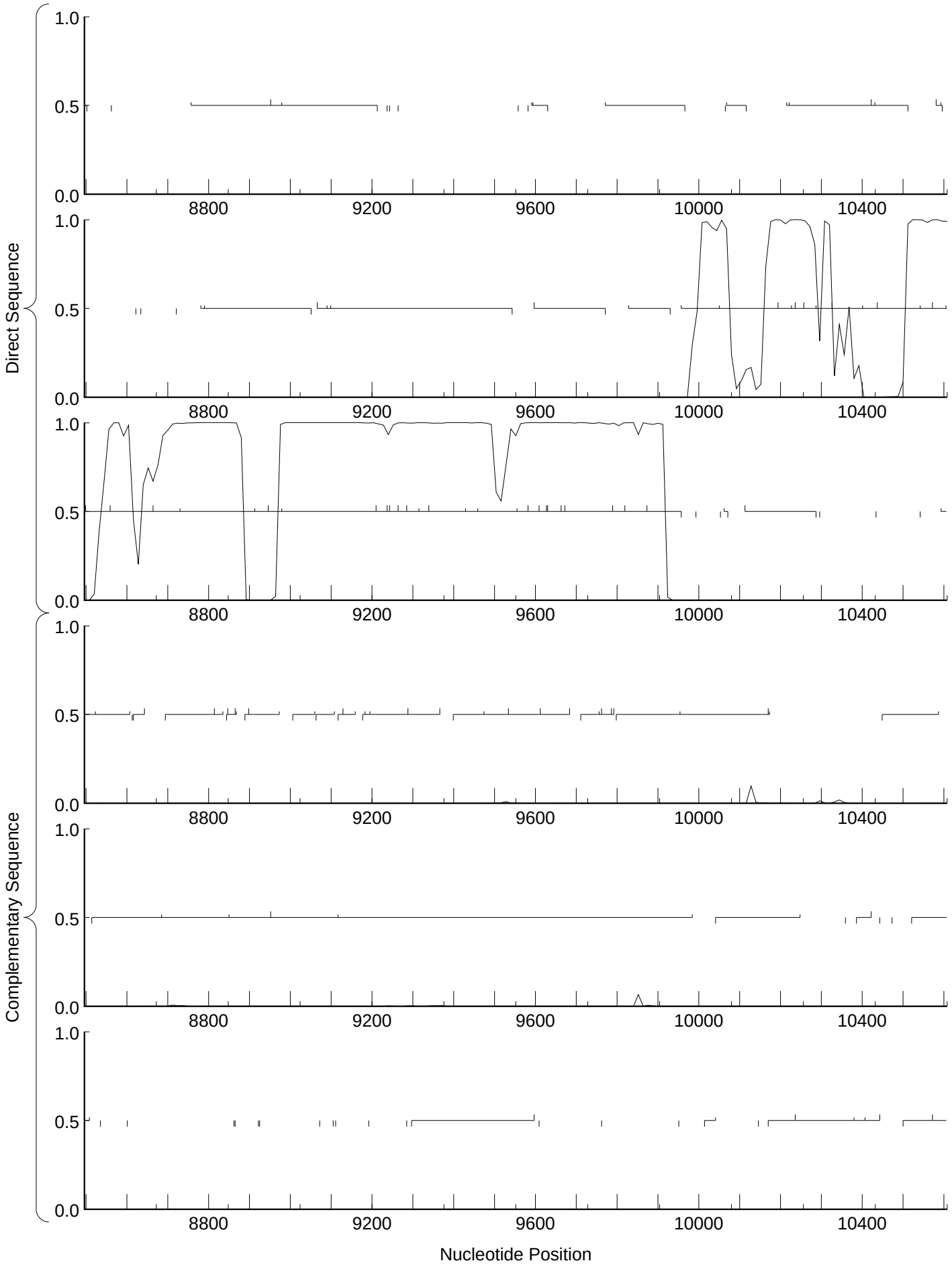
complete sequence, 52206 bp including 10-base 3’ overhang (CGGGTGGTTA), Cluster A15, Order 4, Window 96, Step 12, 4/26



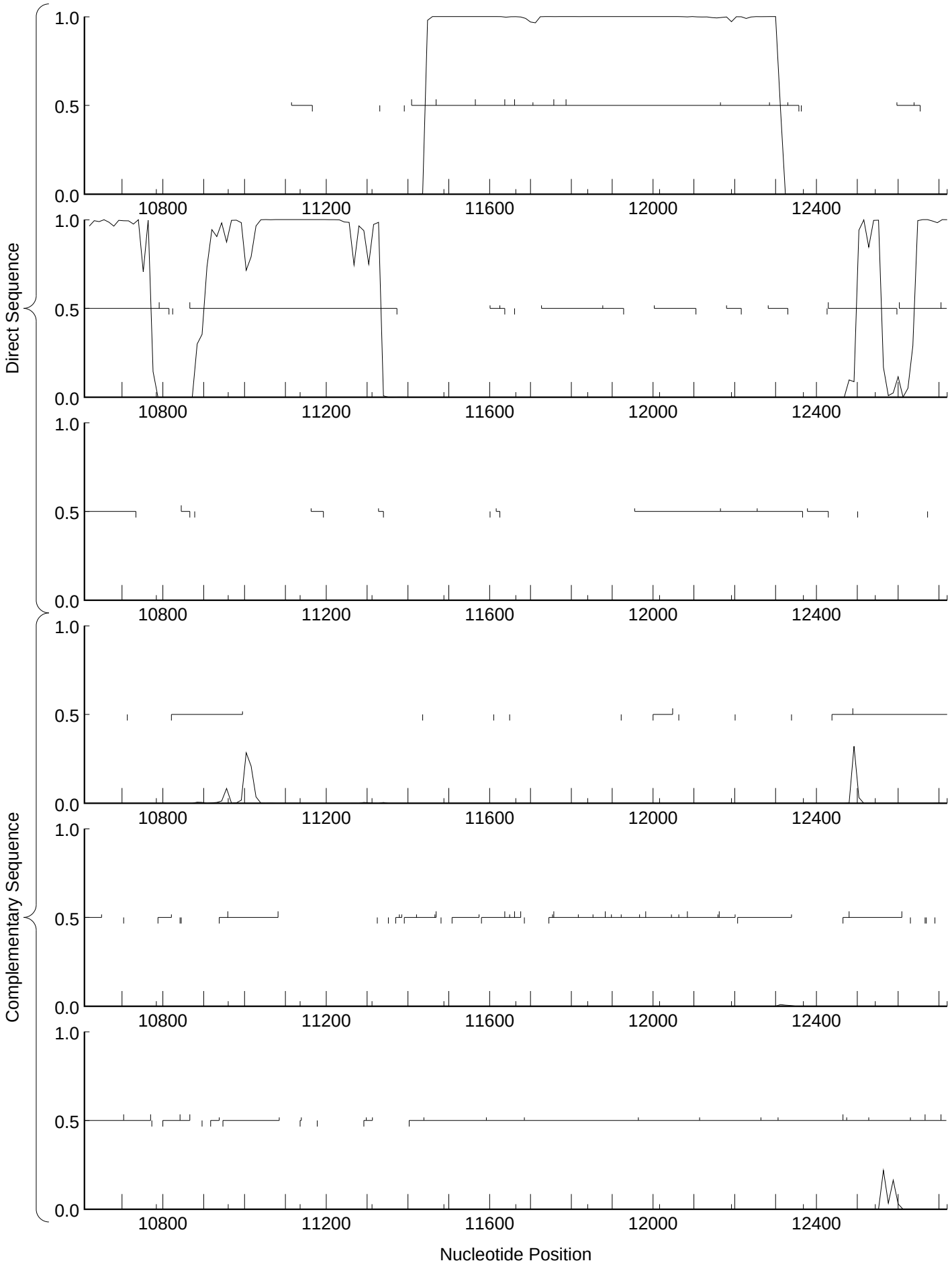
complete sequence, 52206 bp including 10-base 3’ overhang (CGGGTGGTTA), Cluster A15, Order 4, Window 96, Step 12, 5/26



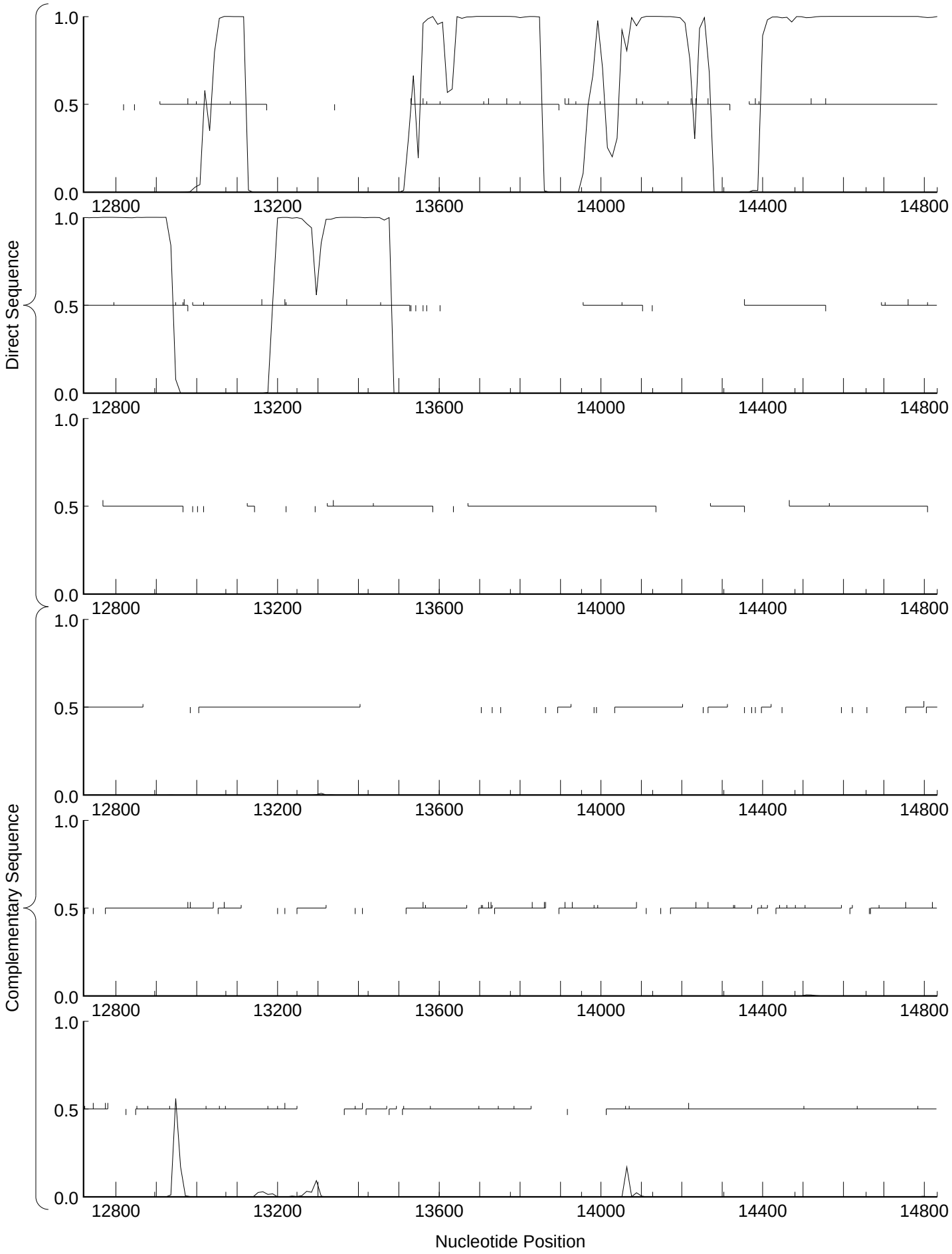
complete sequence, 52206 bp including 10-base 3’ overhang (CGGGTGGTTA), Cluster A15, Order 4, Window 96, Step 12, 6/26



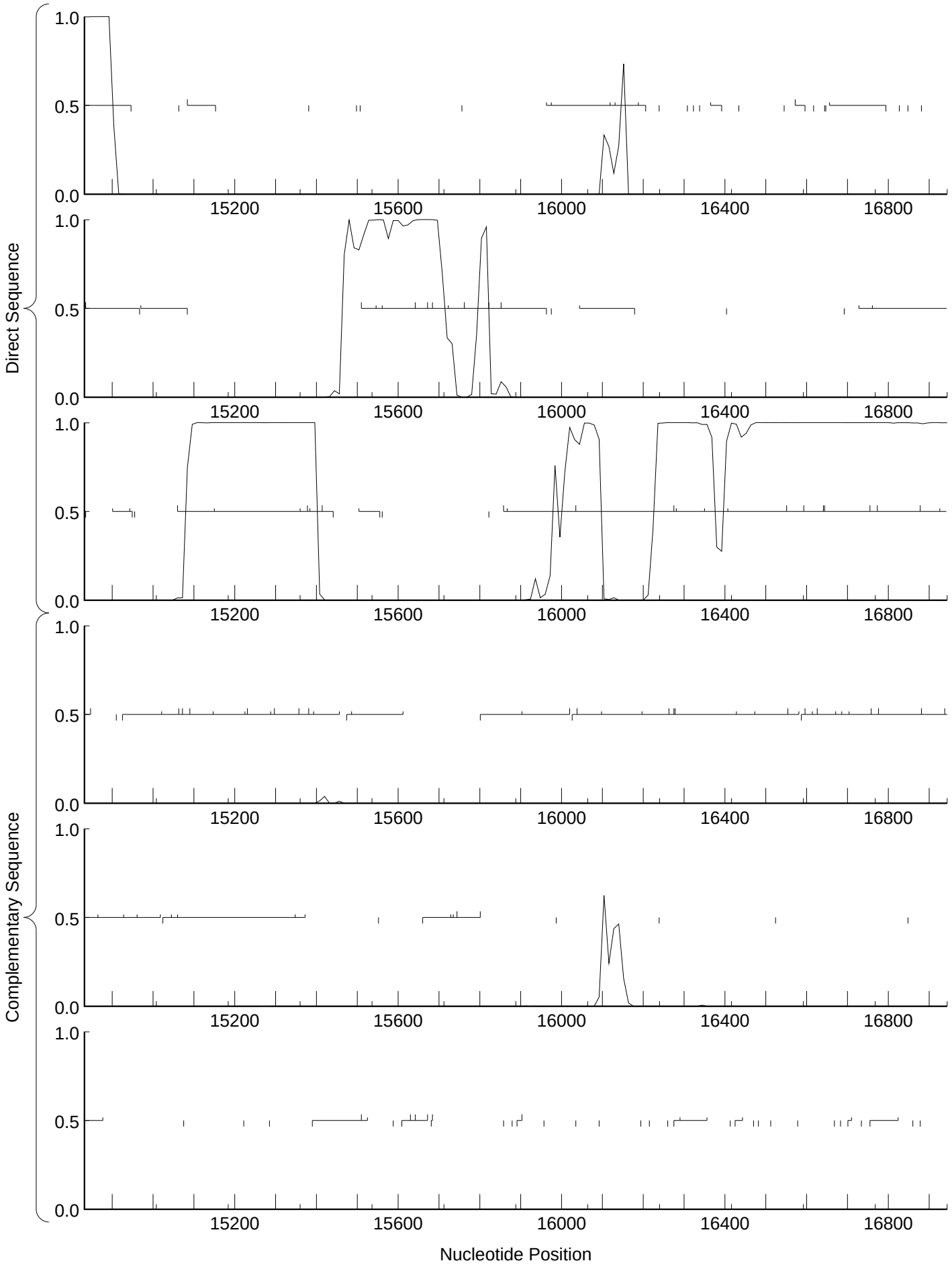
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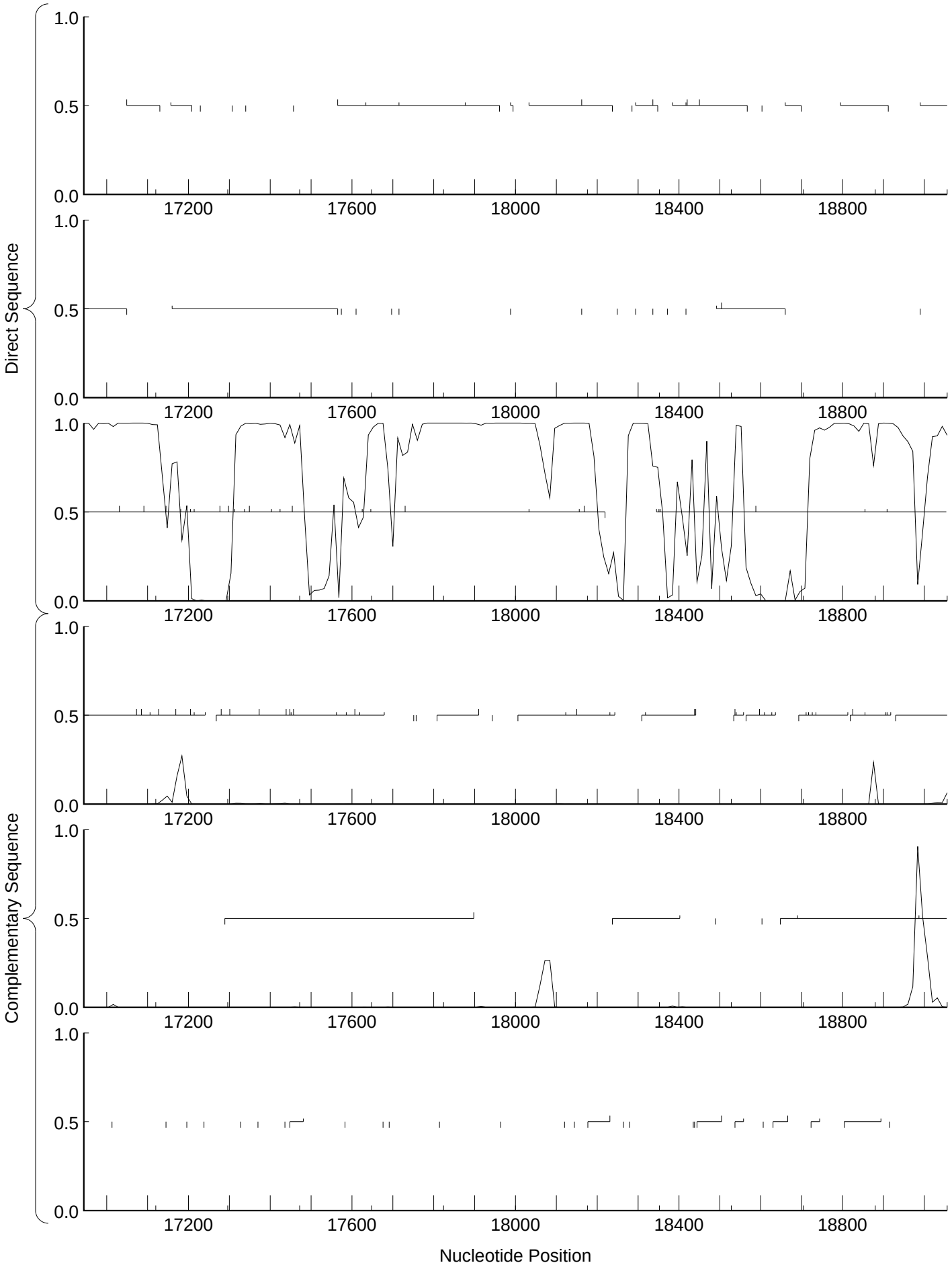
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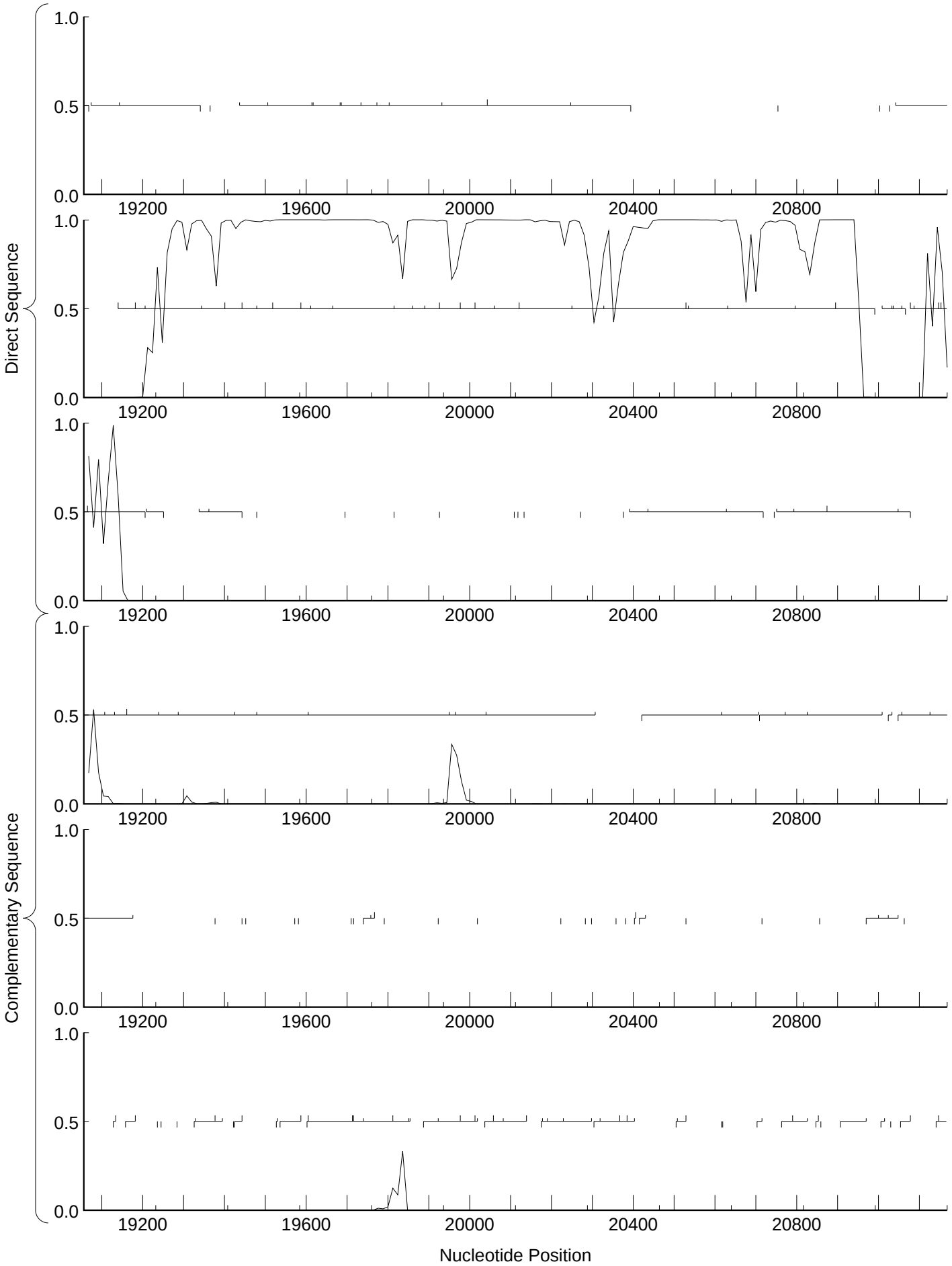
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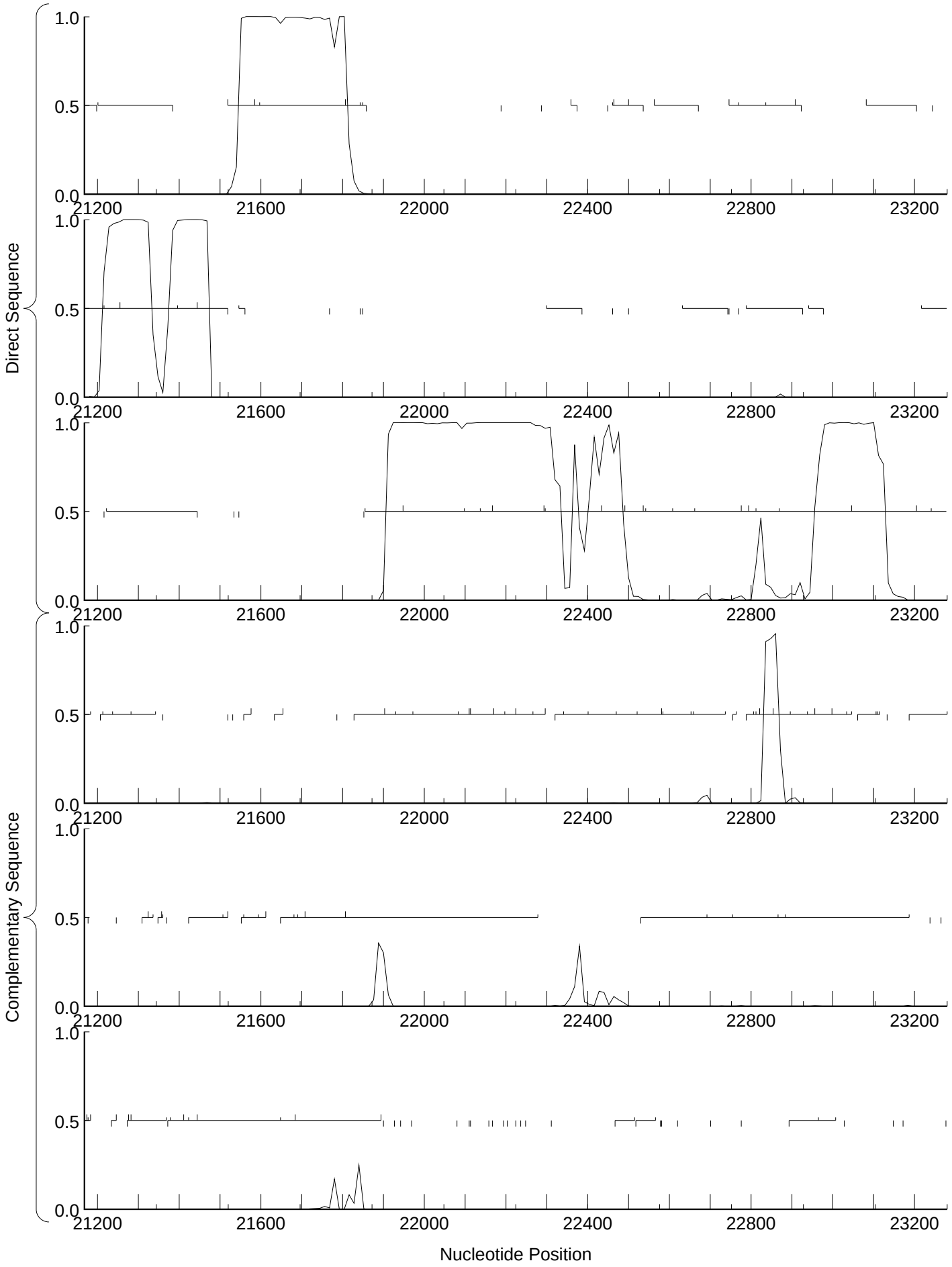
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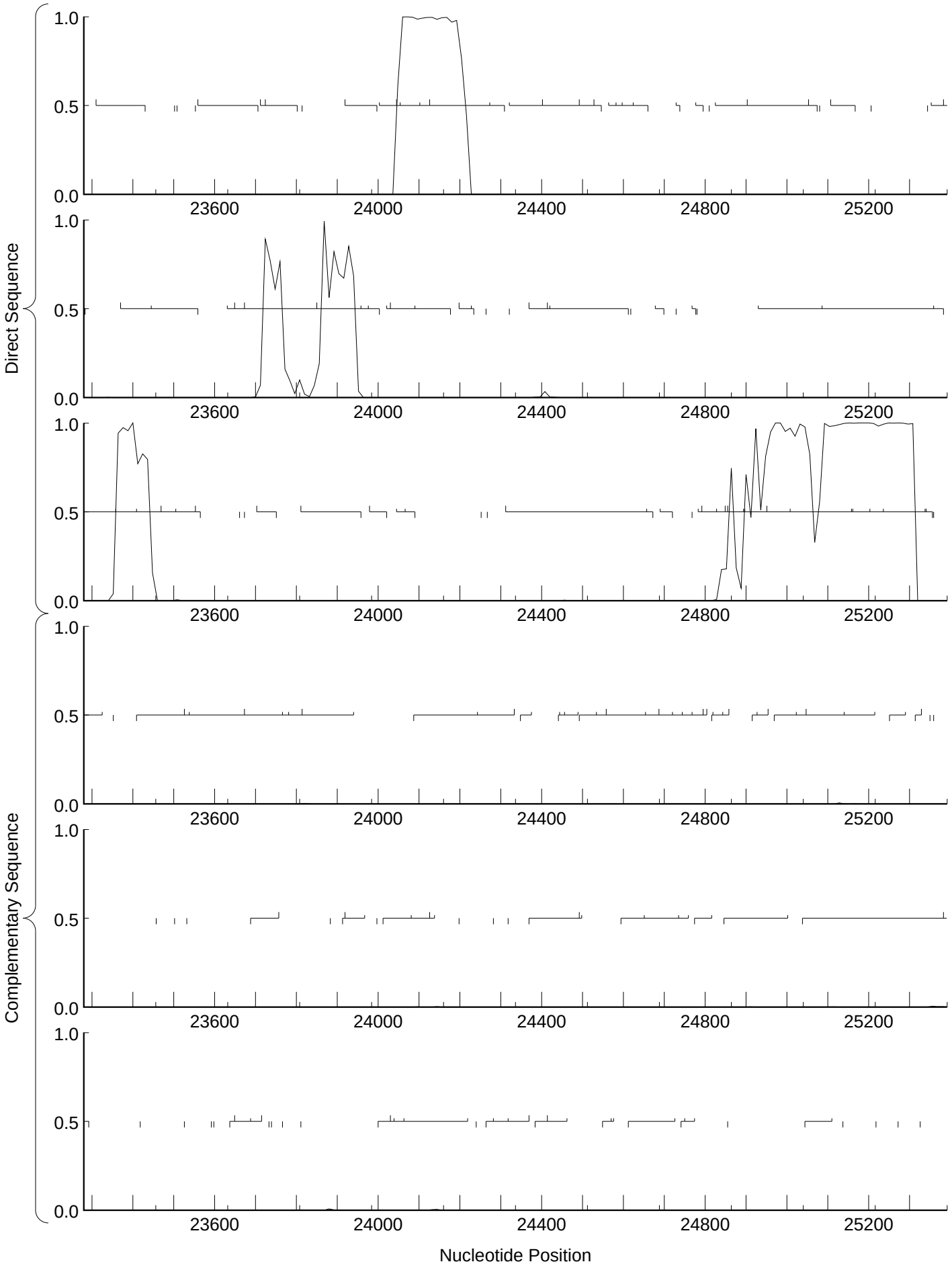
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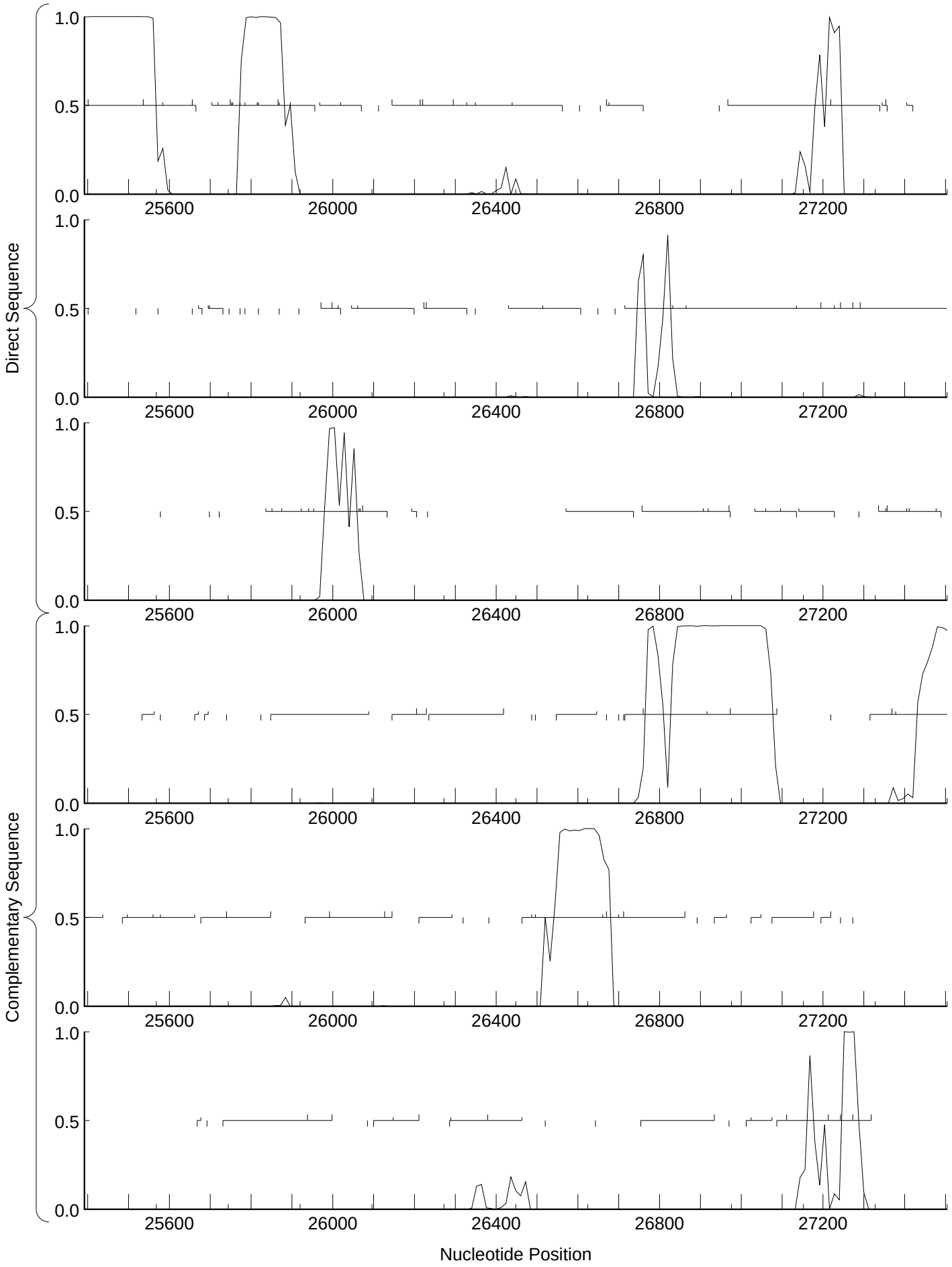
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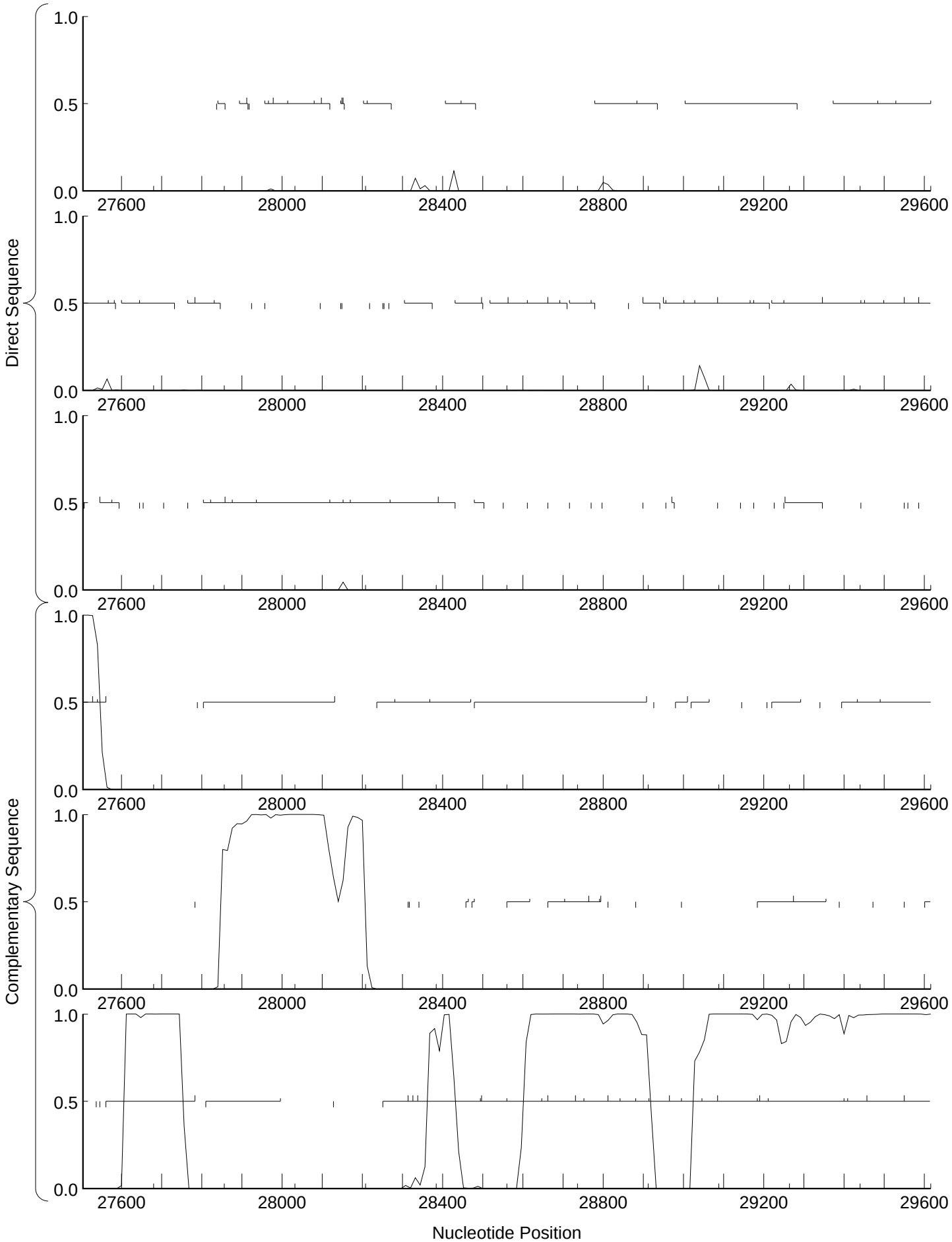
complete sequence, 52206 bp including 10-base 3’ overhang (CGGGTGGTTA), Cluster A15, Order 4, Window 96, Step 12, 13/26



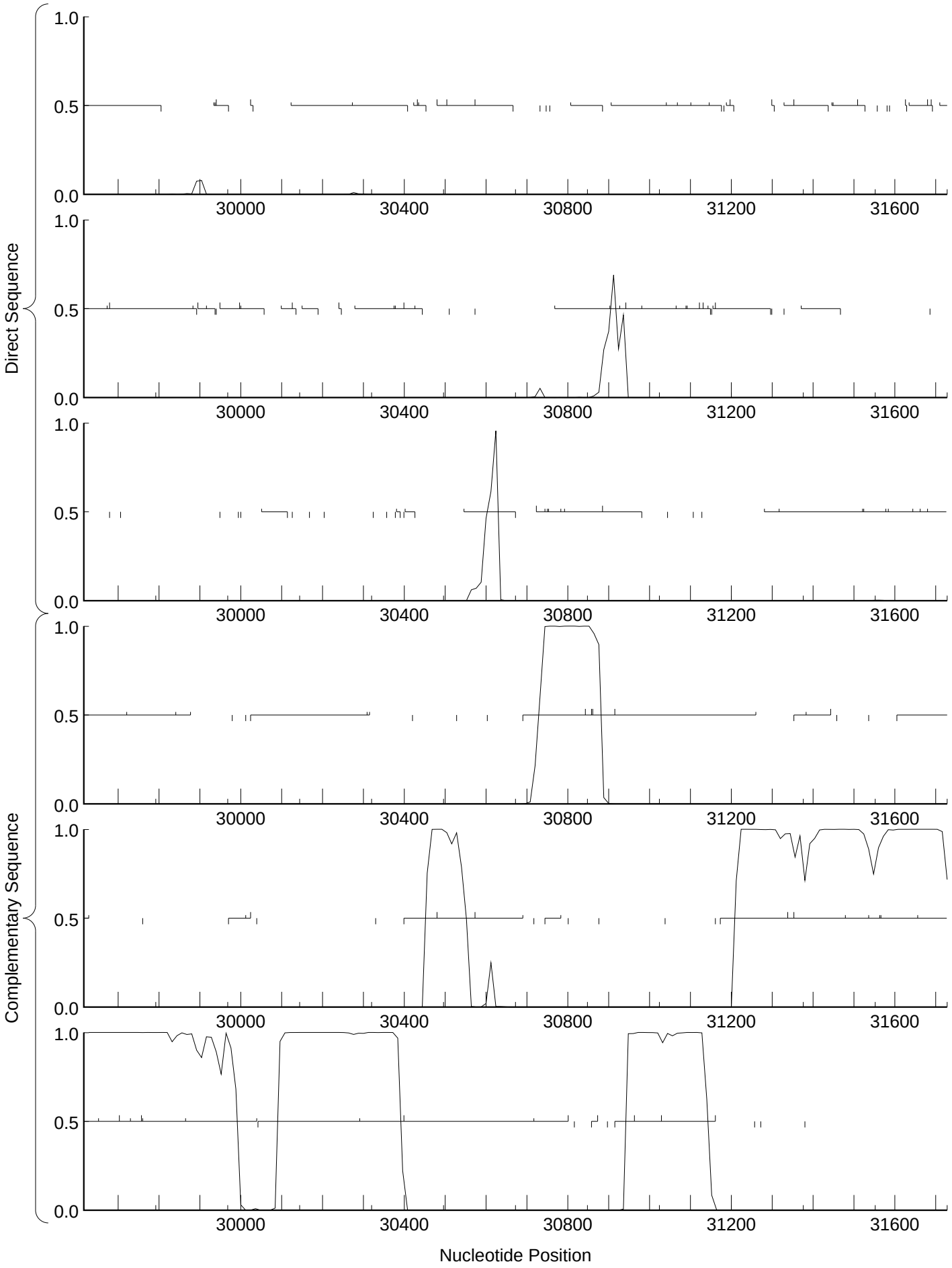
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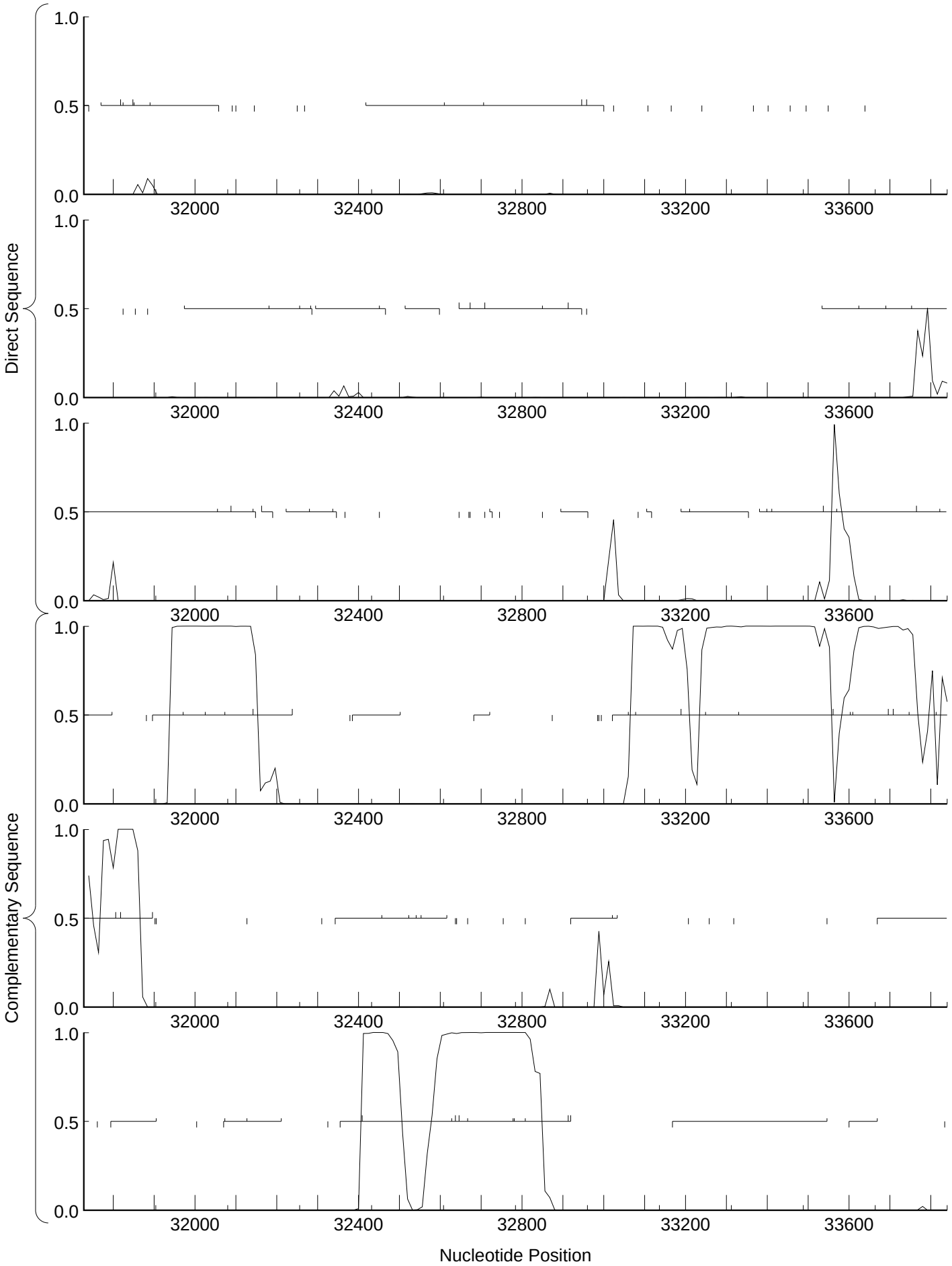
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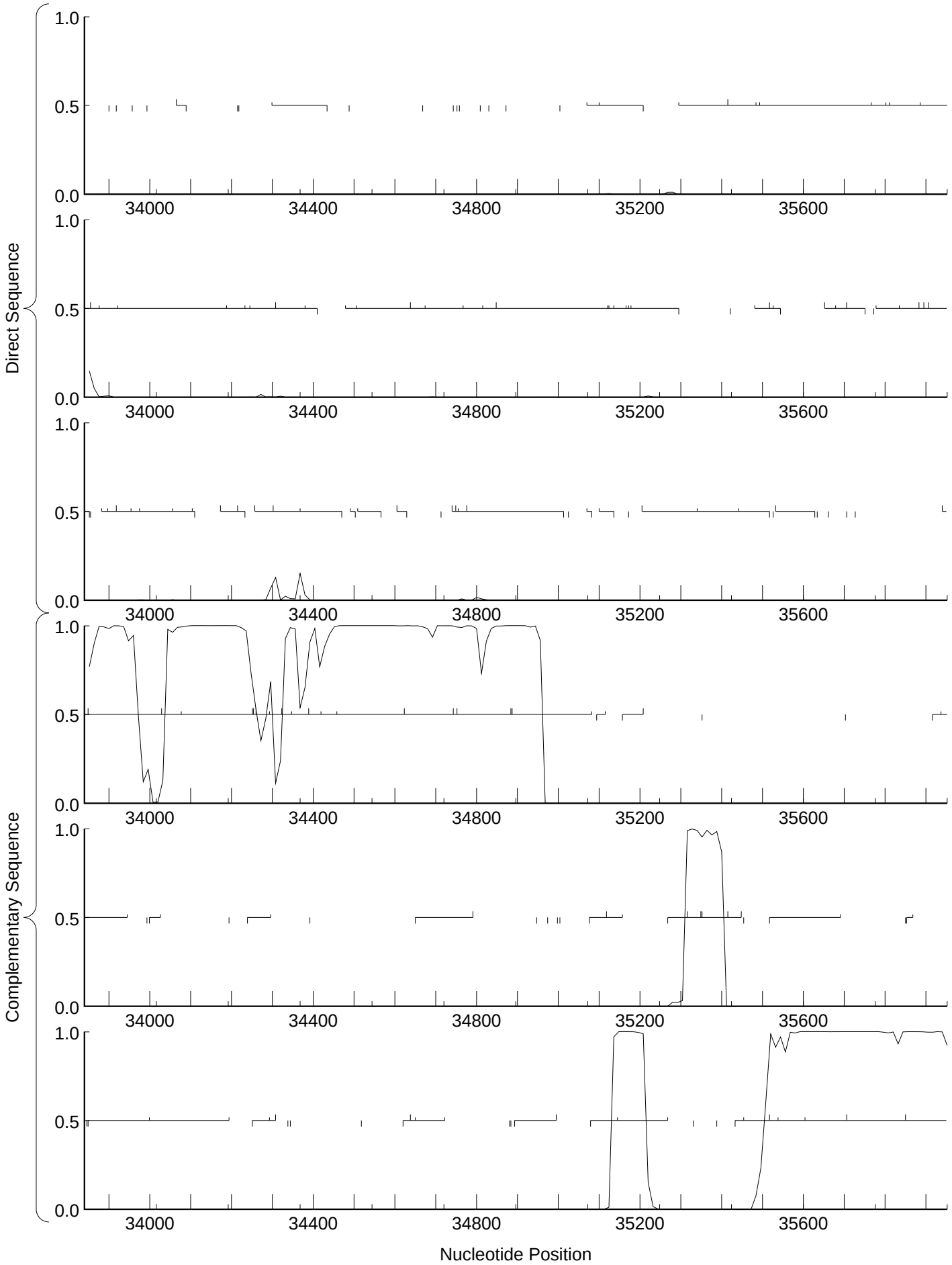


complete sequence, 52206 bp including 10-base 3’ overhang (CGGGTGGTTA), Cluster A15, Order 4, Window 96, Step 12, 16/26

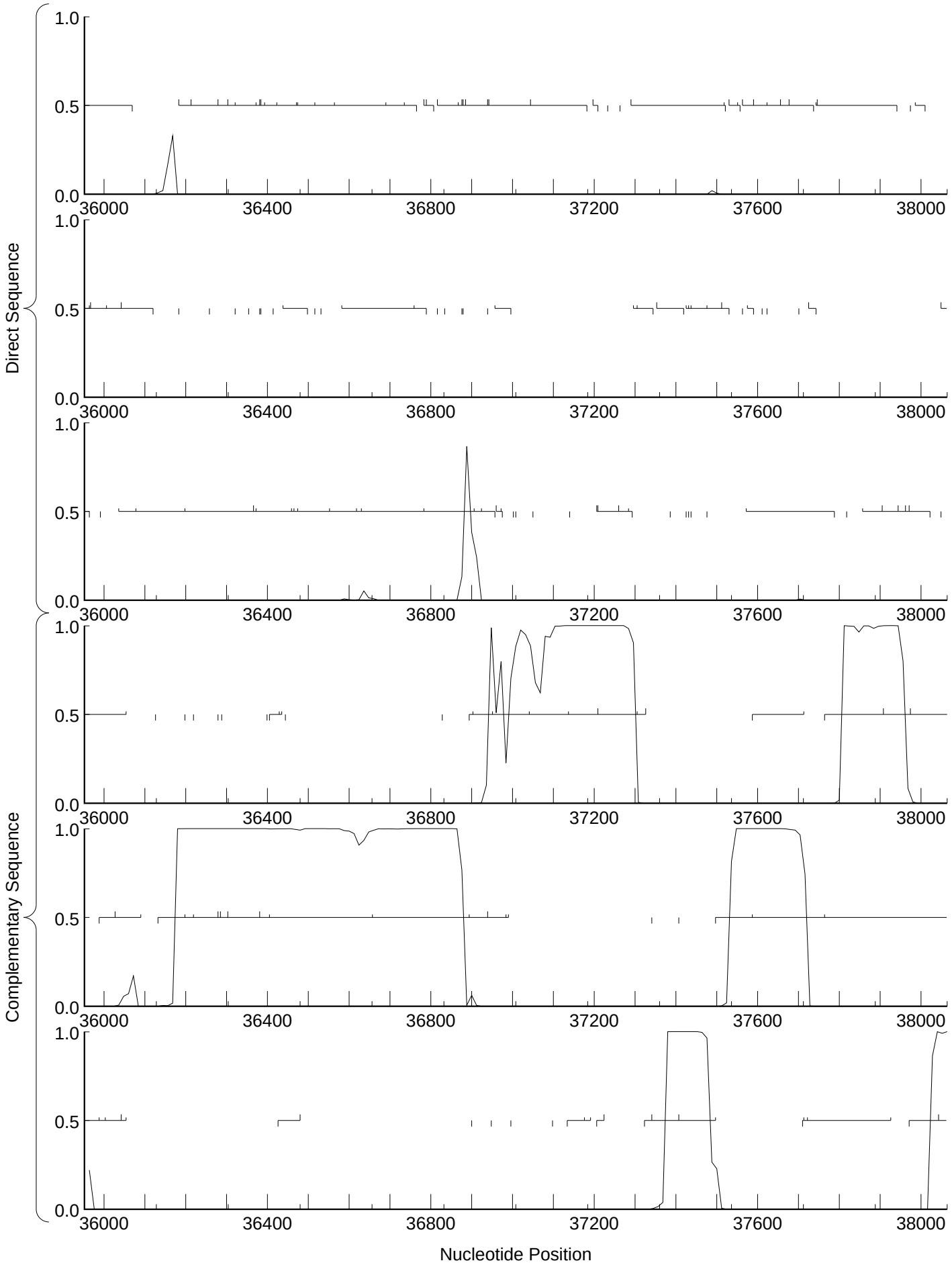


complete sequence, 52206 bp including 10-base 3’ overhang (CGGGTGGTTA), Cluster A15, Order 4, Window 96, Step 12, 17/26

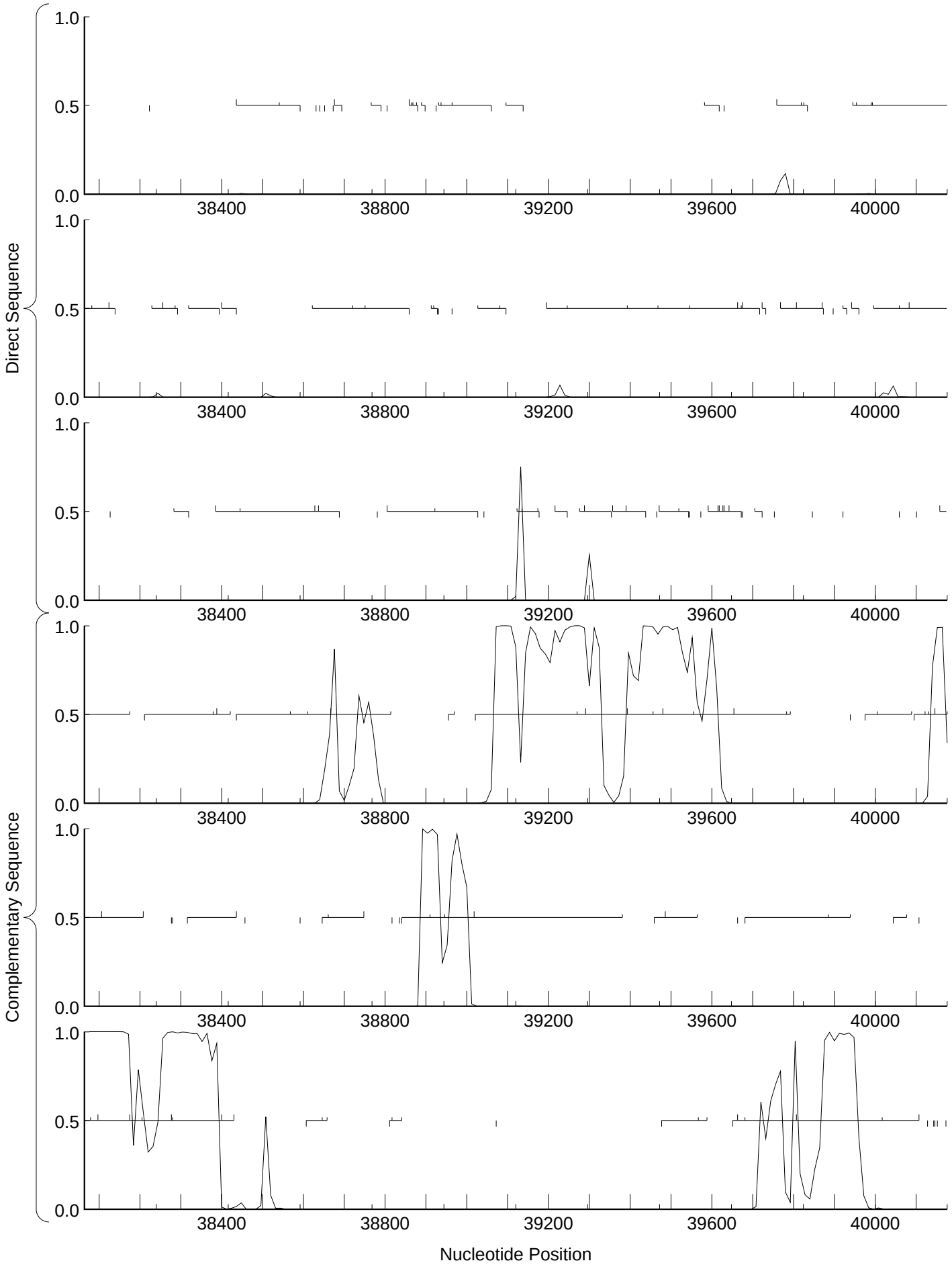



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

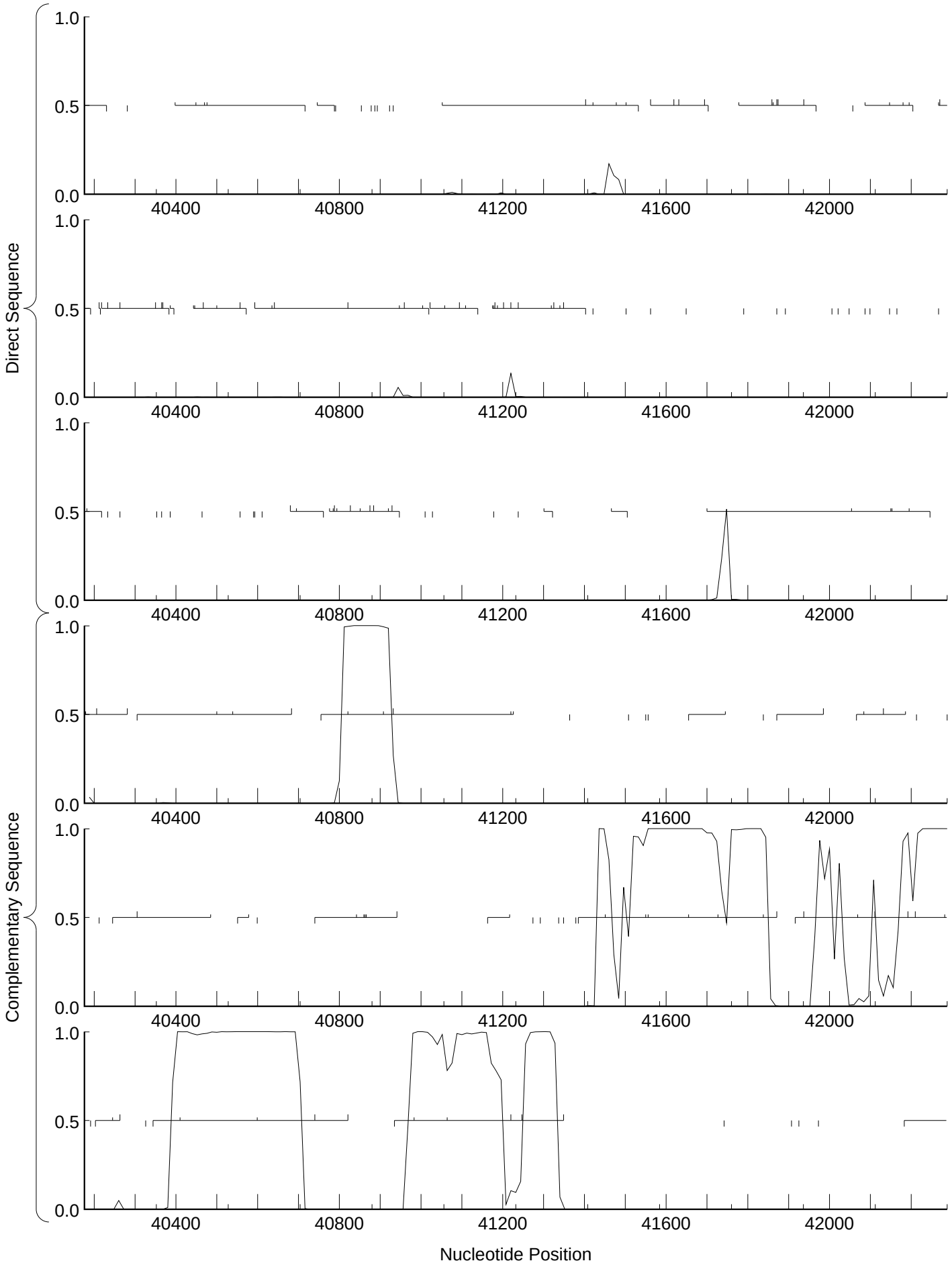
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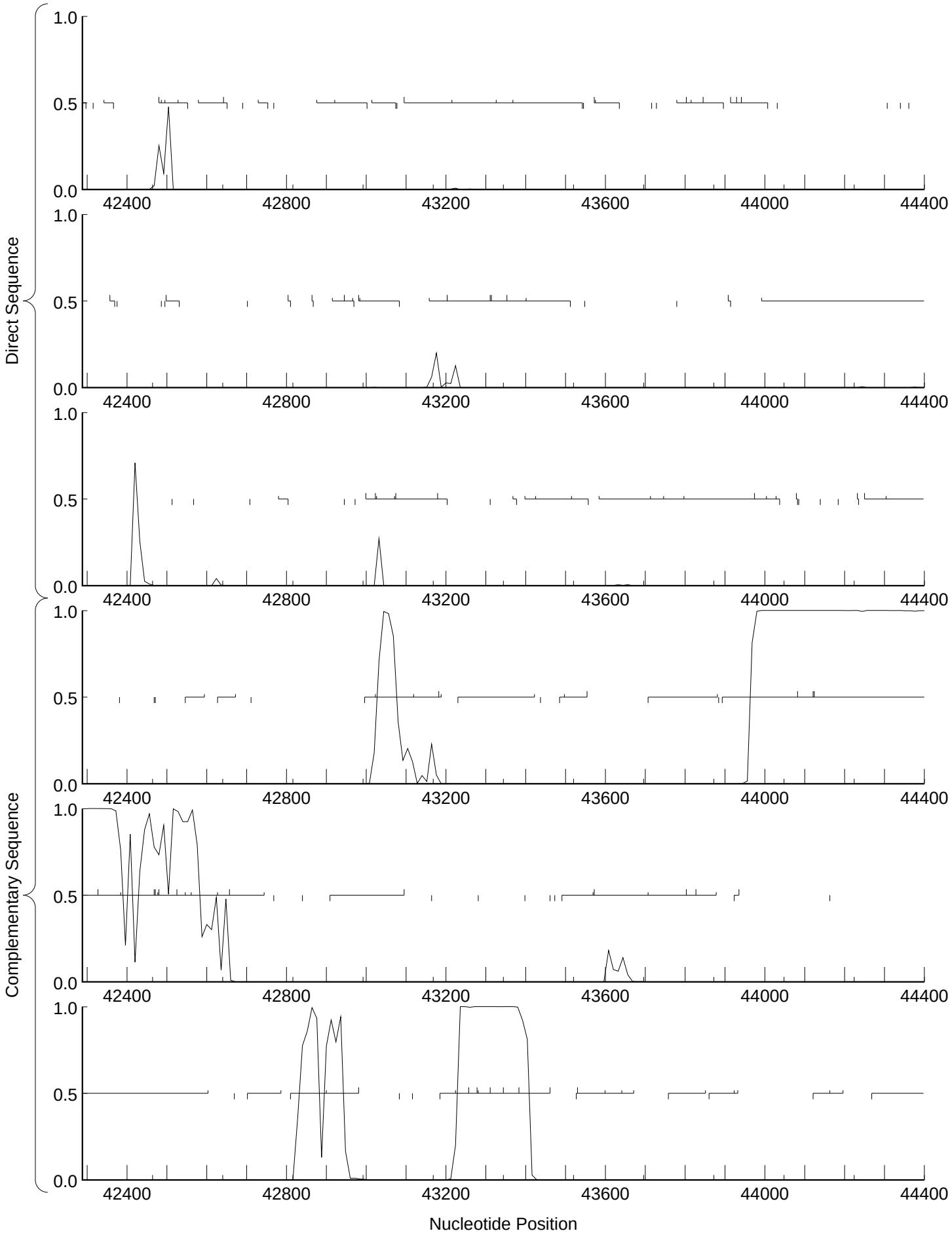
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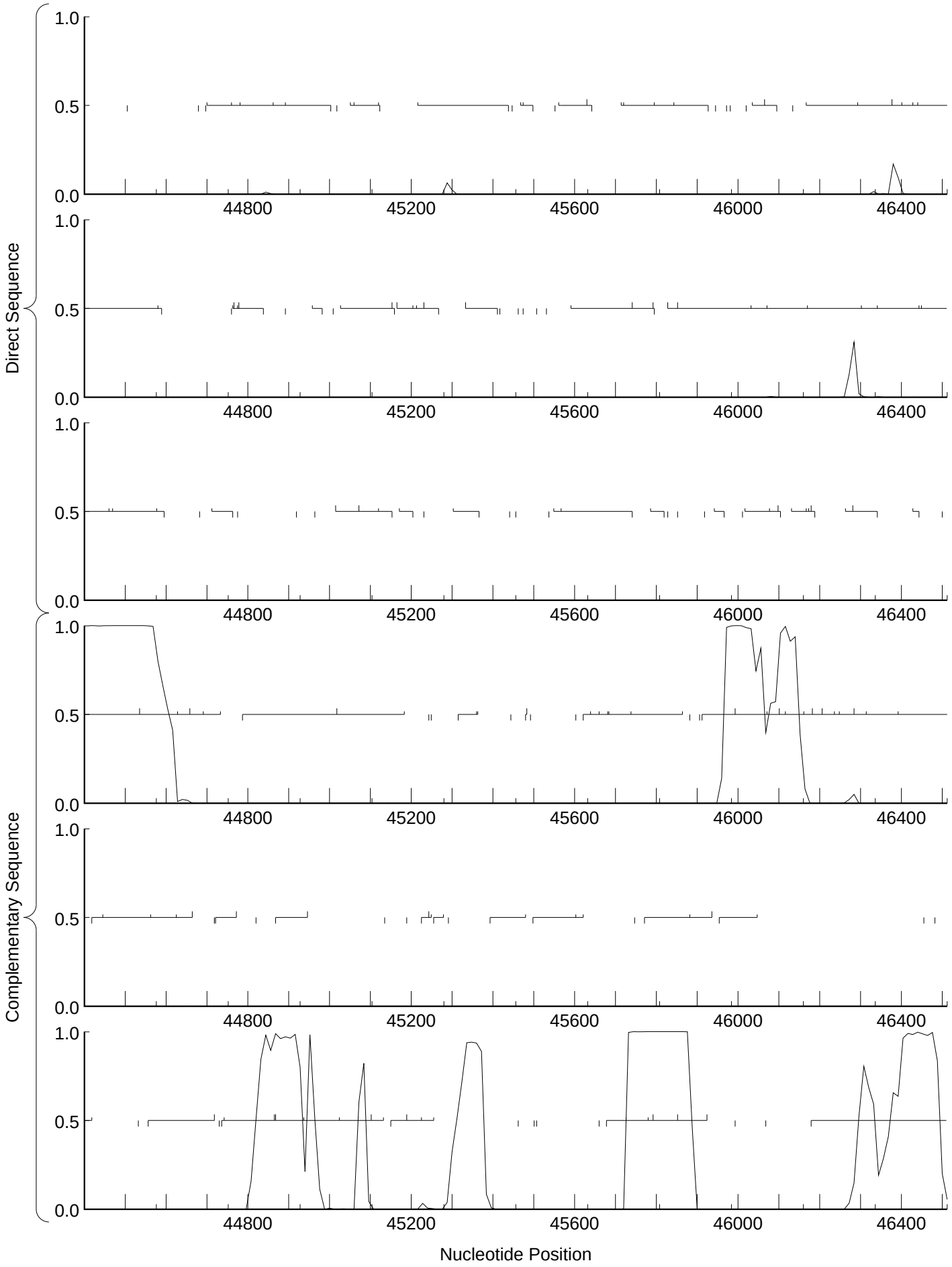
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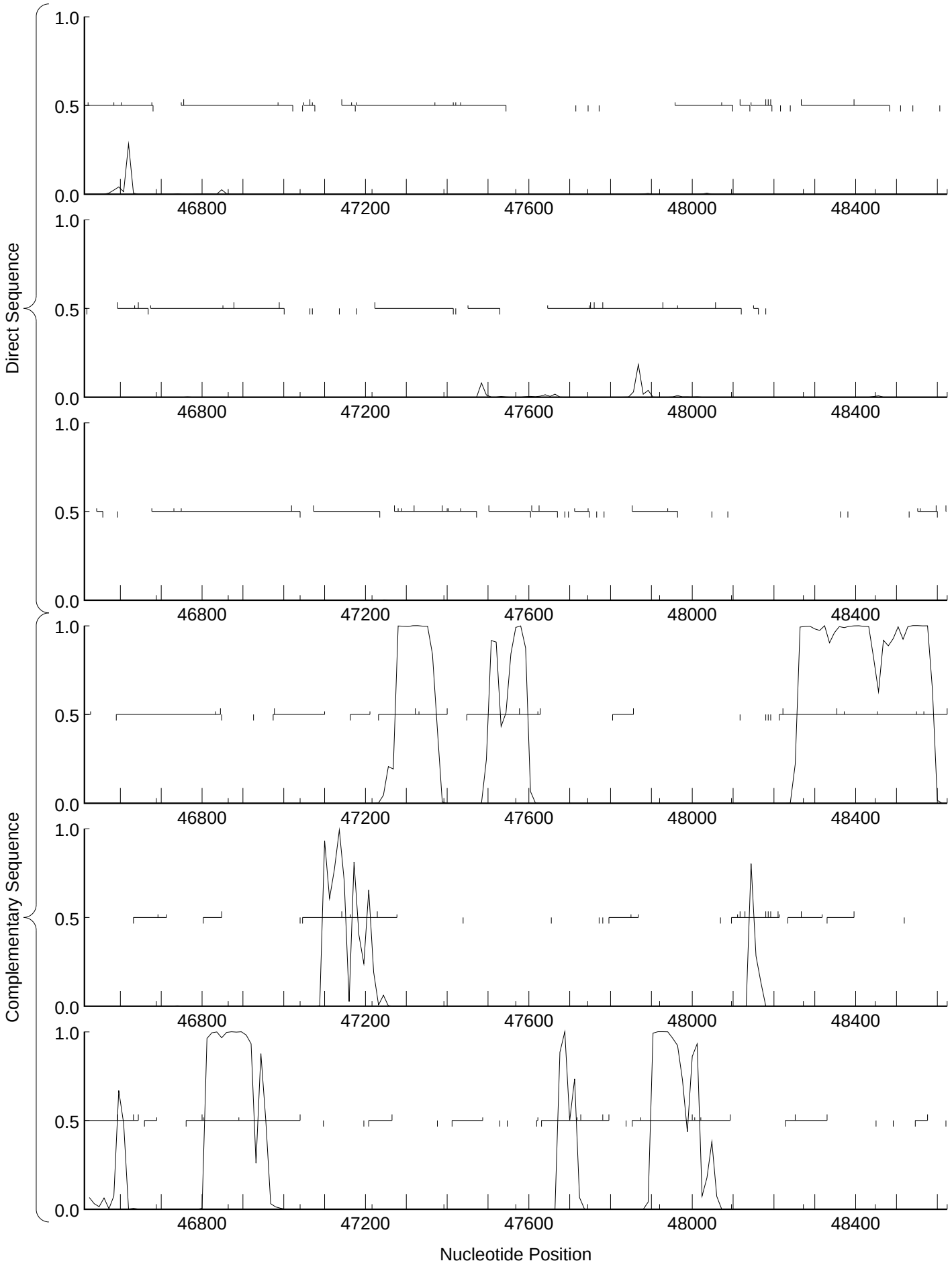
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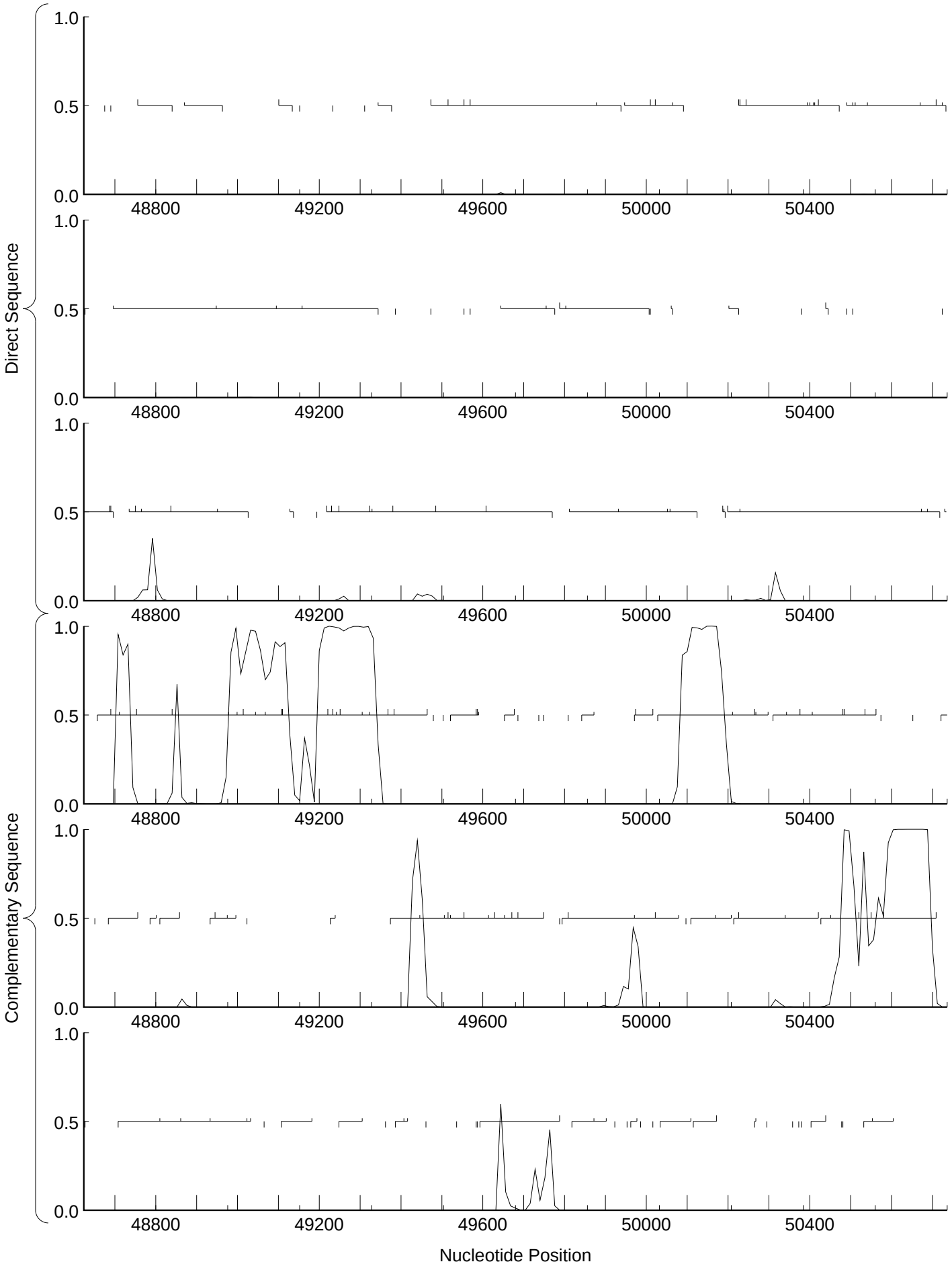
complete sequence, 52206 bp including 10-base 3’ overhang (CGGGTGGTTA), Cluster A15, Order 4, Window 96, Step 12, 23/26



complete sequence, 52206 bp including 10-base 3’ overhang (CGGGTGGTTA), Cluster A15, Order 4, Window 96, Step 12, 24/26



complete sequence, 52206 bp including 10-base 3’ overhang (CGGGTGGTTA), Cluster A15, Order 4, Window 96, Step 12, 25/26



complete sequence, 52206 bp including 10-base 3’ overhang (CGGGTGGTTA), Cluster A15, Order 4, Window 96, Step 12, 26/26

