GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Gordonia phage Francois complete sequence, 50499 bp including 10-base 3’ overhang (TGCCGCGG
Analysis Date    :  3/28/24 at 7:42:27
Pages            :  25
Sequence Length  :  50499 bp
GC Content       :   67.36%

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,
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MATRIX INFORMATION

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Matrix     :  -
Author     :  -

Order      :  4

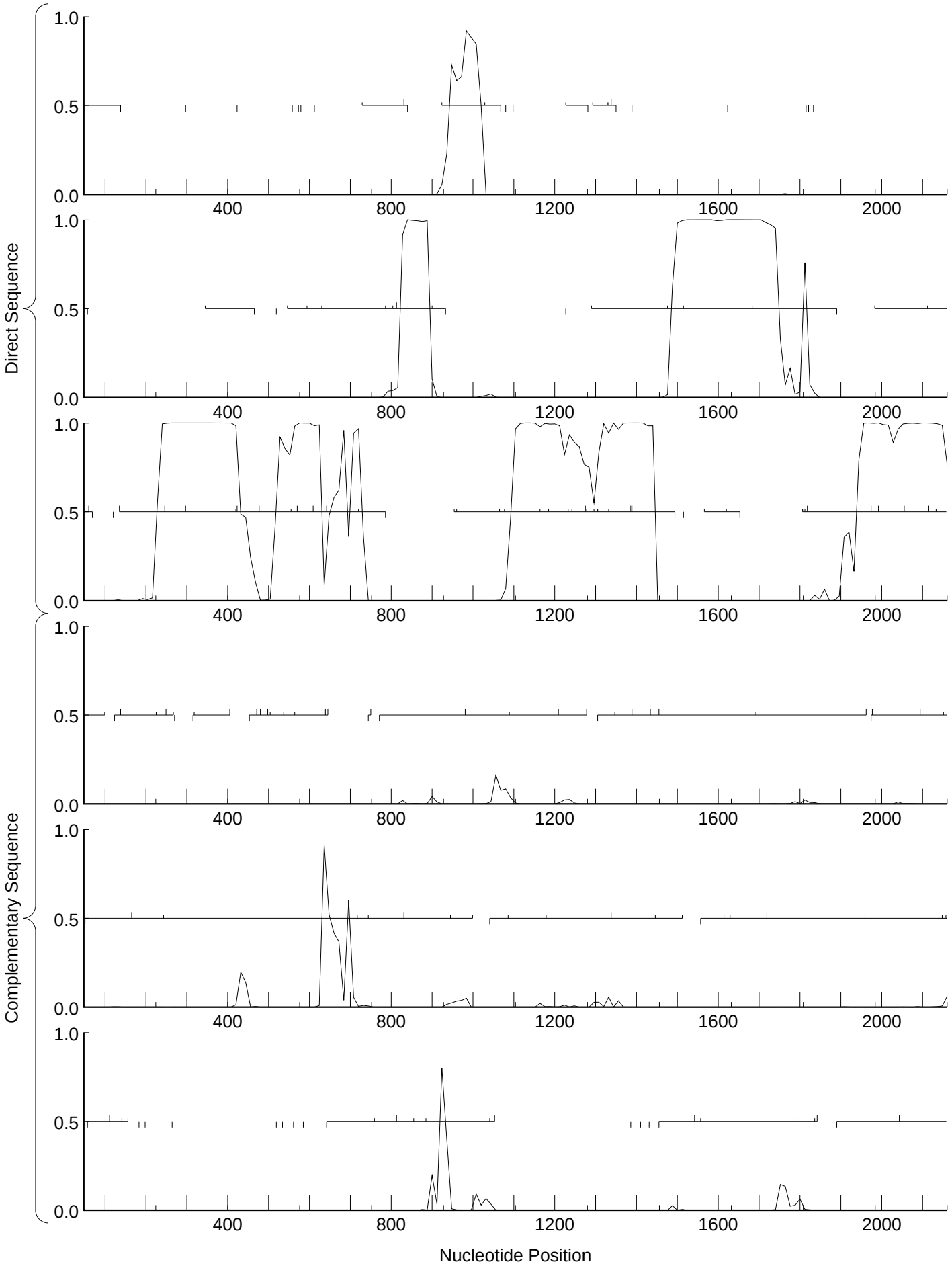

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

Matrix notes & comments

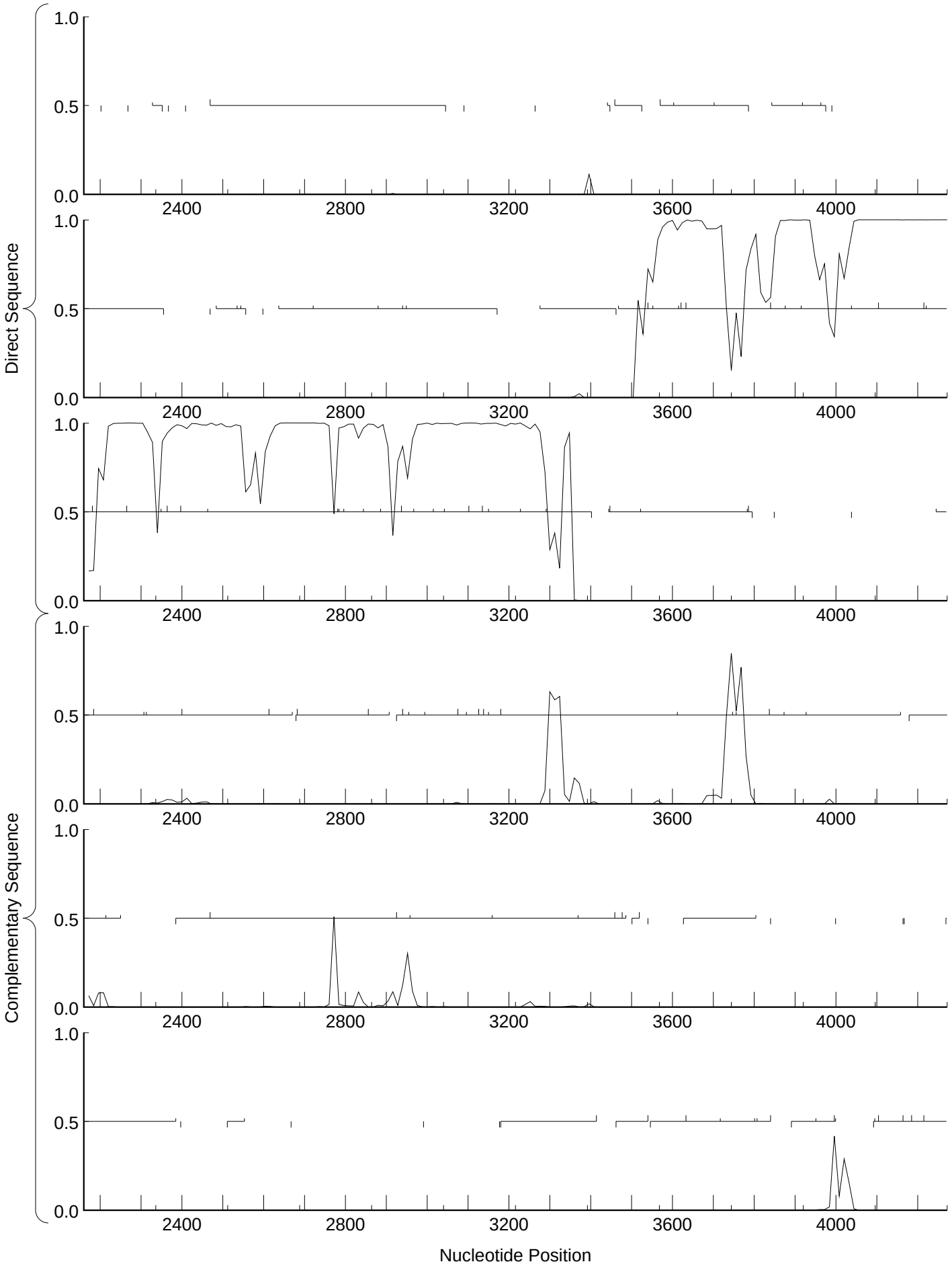
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Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep  8 00:36:14 2018
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For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

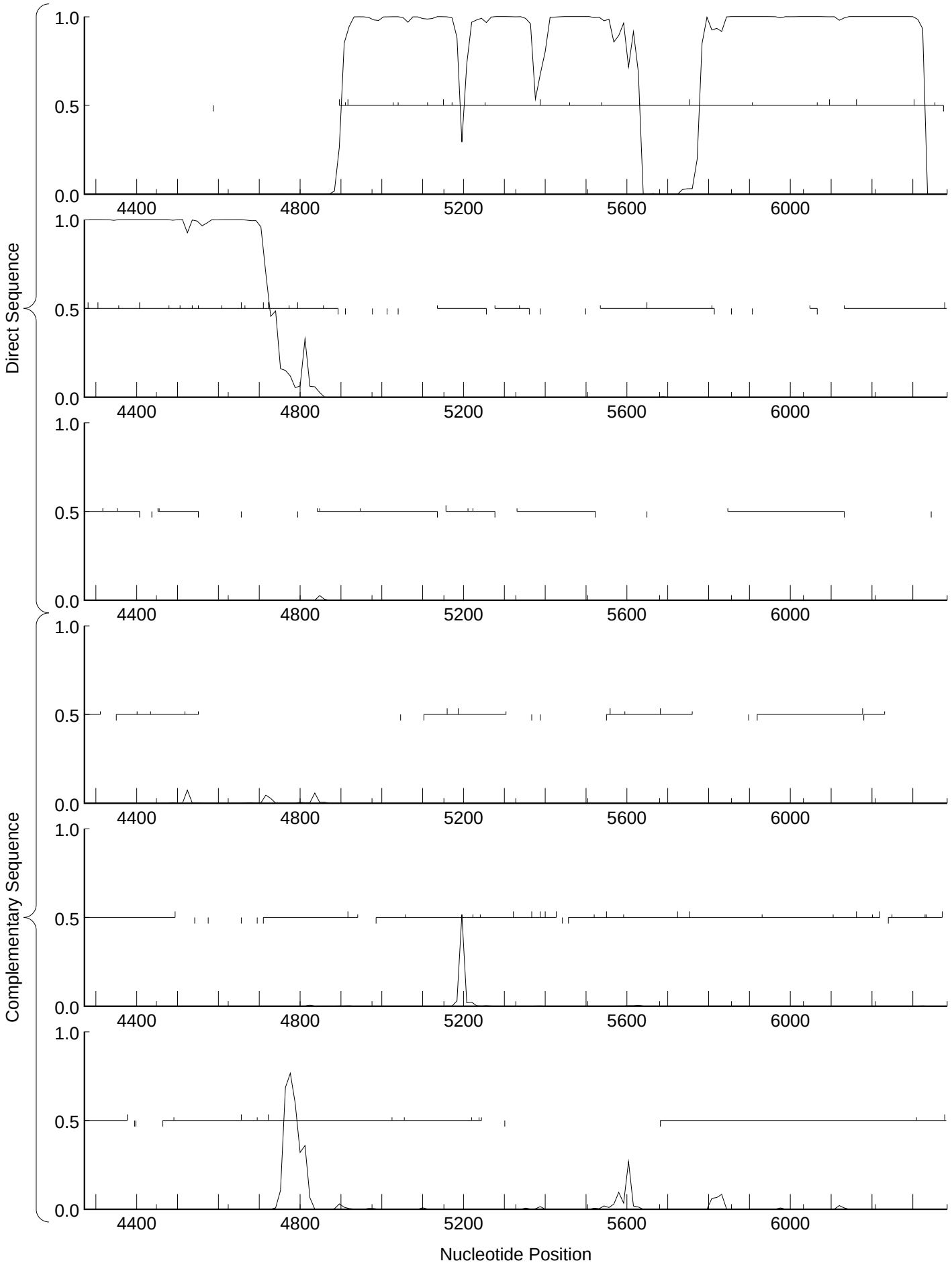
ois complete sequence, 50499 bp including 10-base 3’ overhang (TGCCGCGGTA), Cluster DI, Order 4, Window 96, Step 12, 2/25

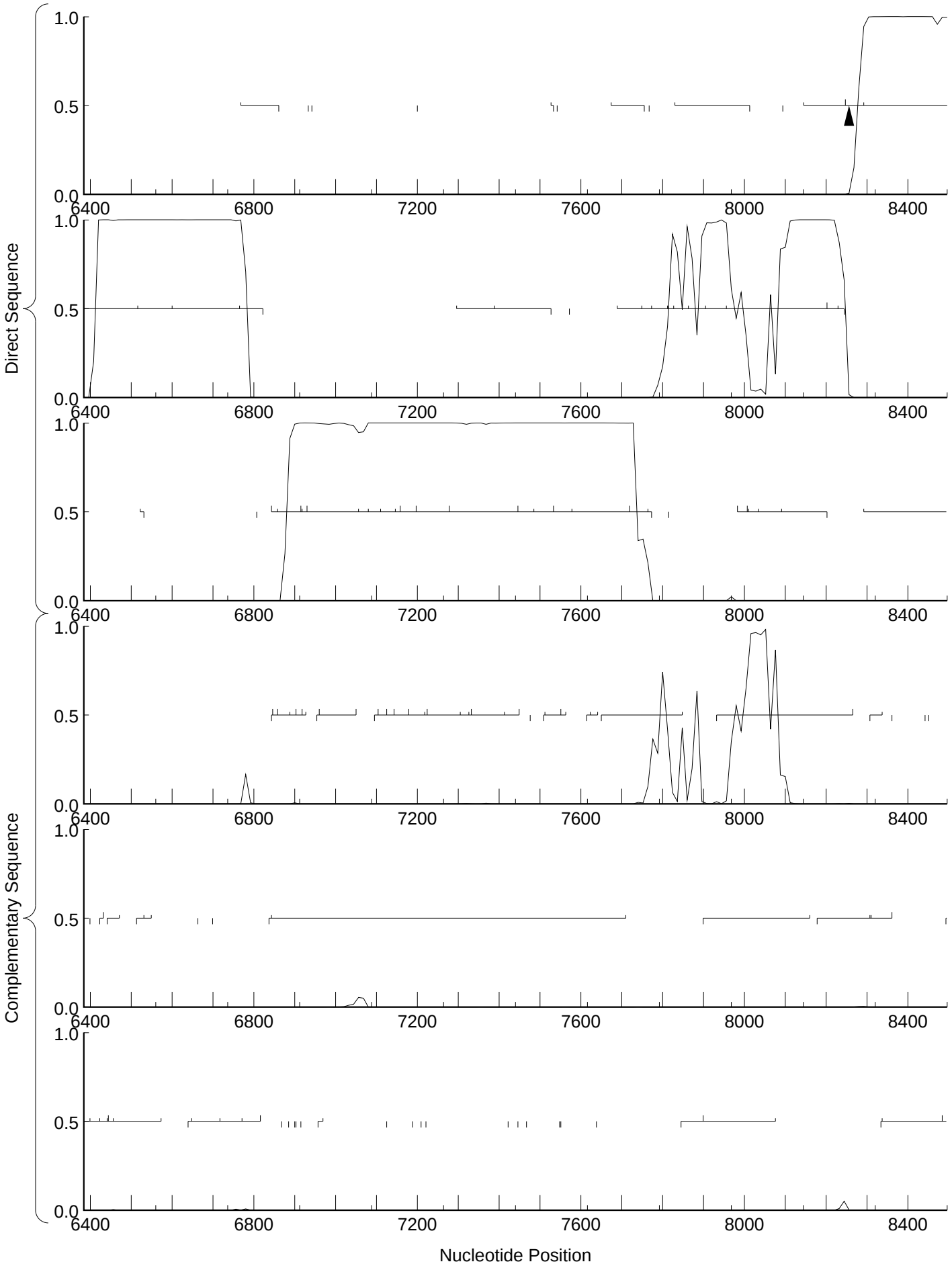


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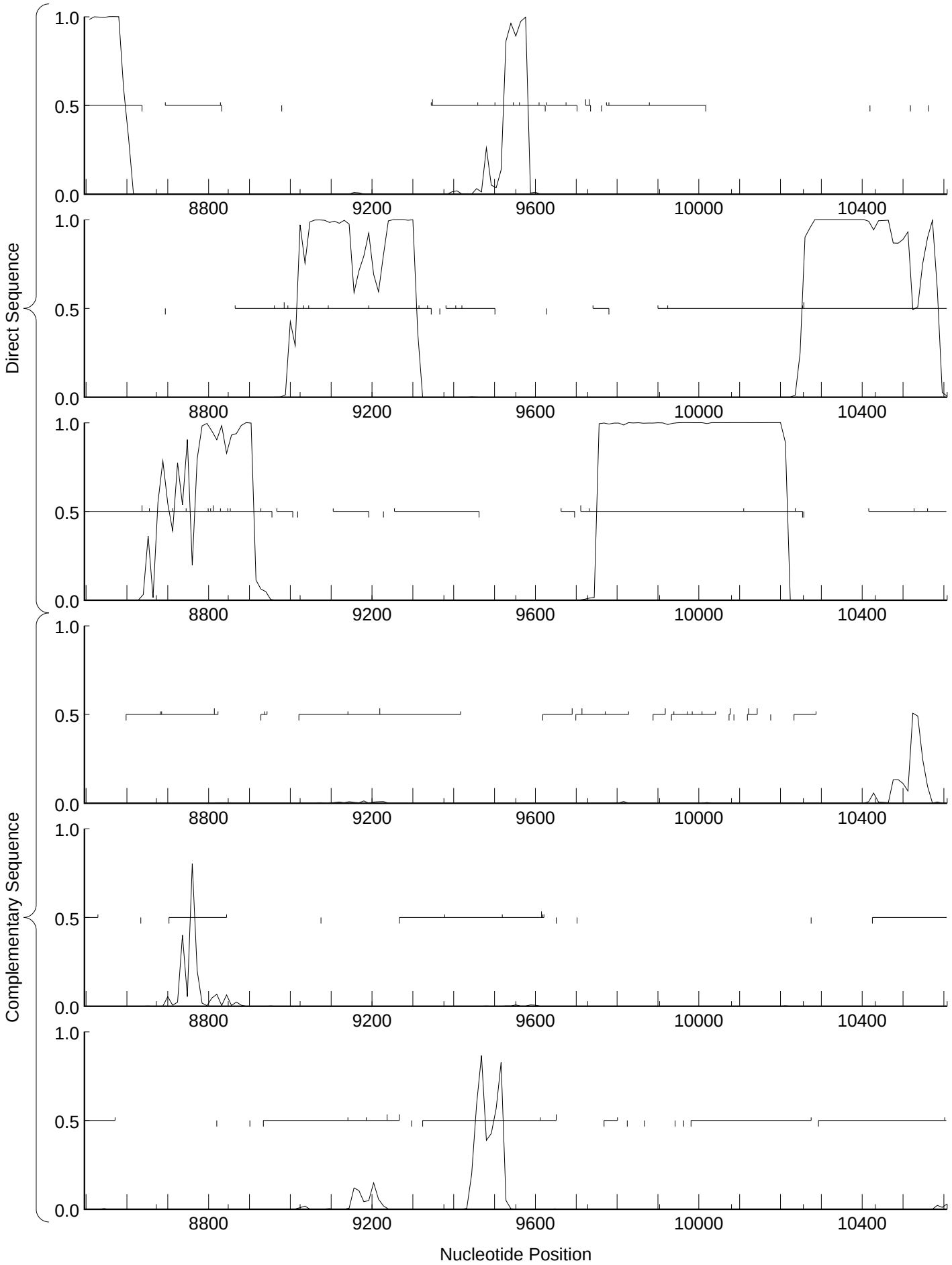


ois complete sequence, 50499 bp including 10-base 3’ overhang (TGCCGCGGTA), Cluster DI, Order 4, Window 96, Step 12, 4/25

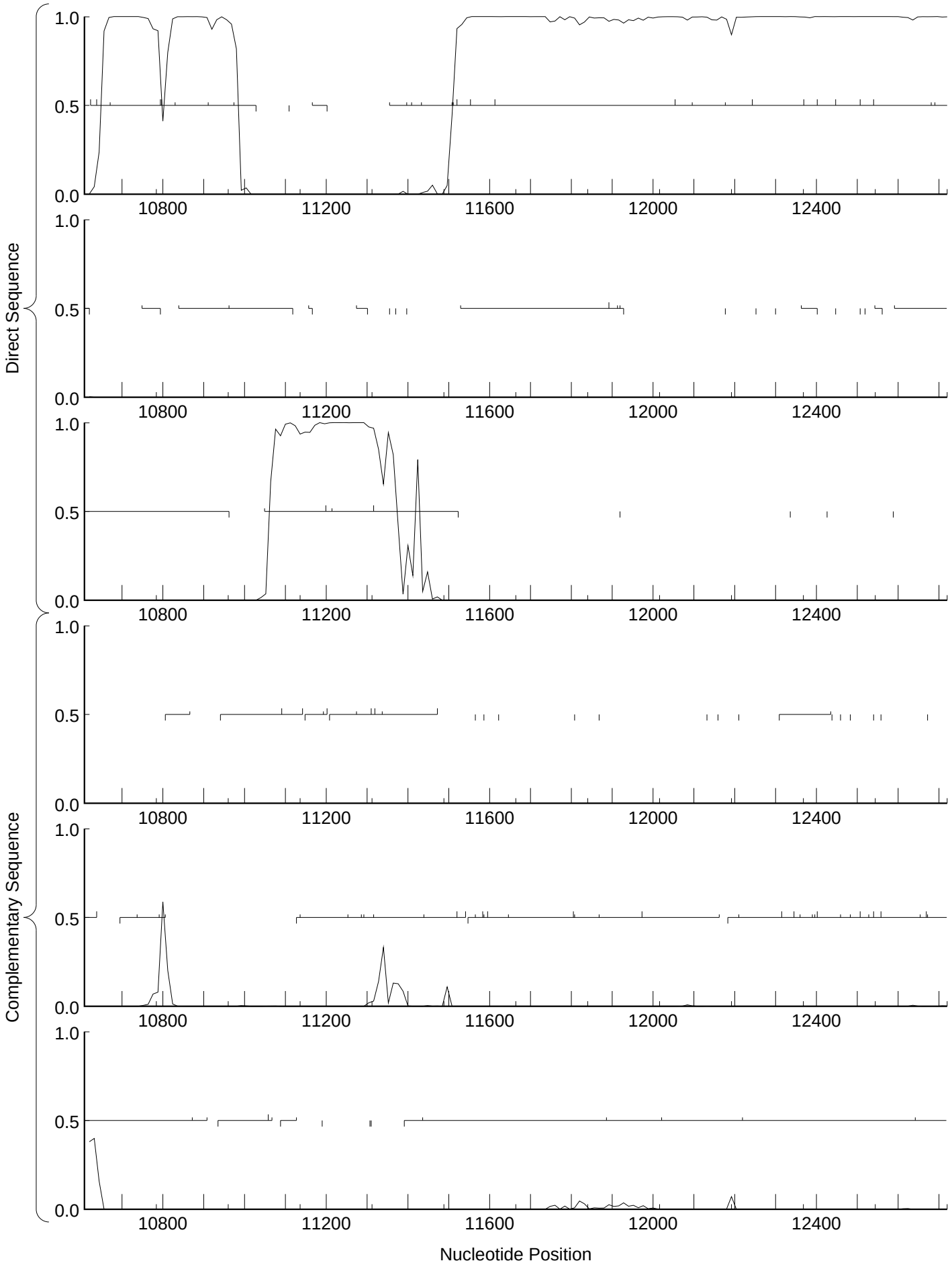



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
6400
6800
7200
7600
8000
8400
Nucleotide Position

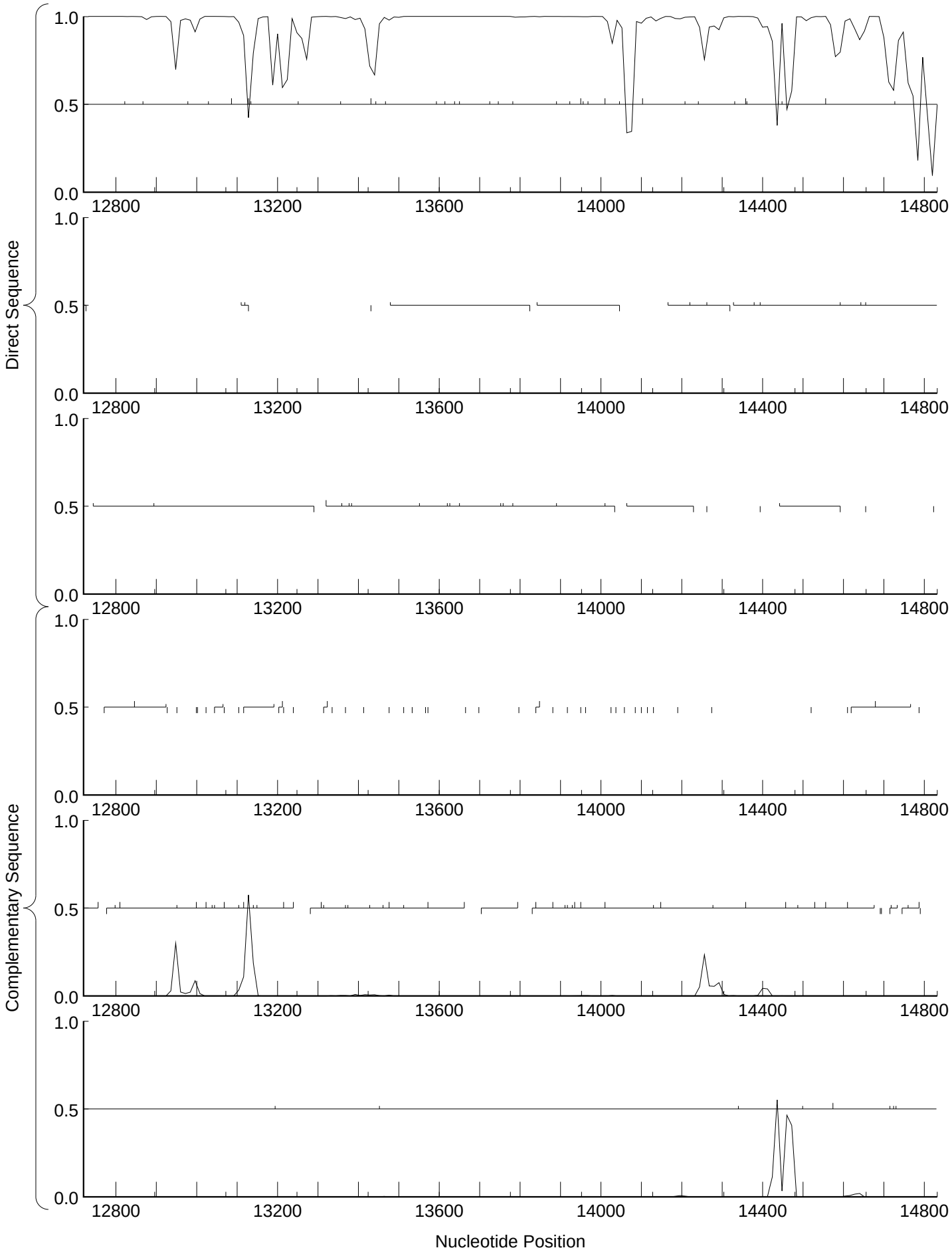
ois complete sequence, 50499 bp including 10-base 3’ overhang (TGCCGCGGTA), Cluster DI, Order 4, Window 96, Step 12, 6/25



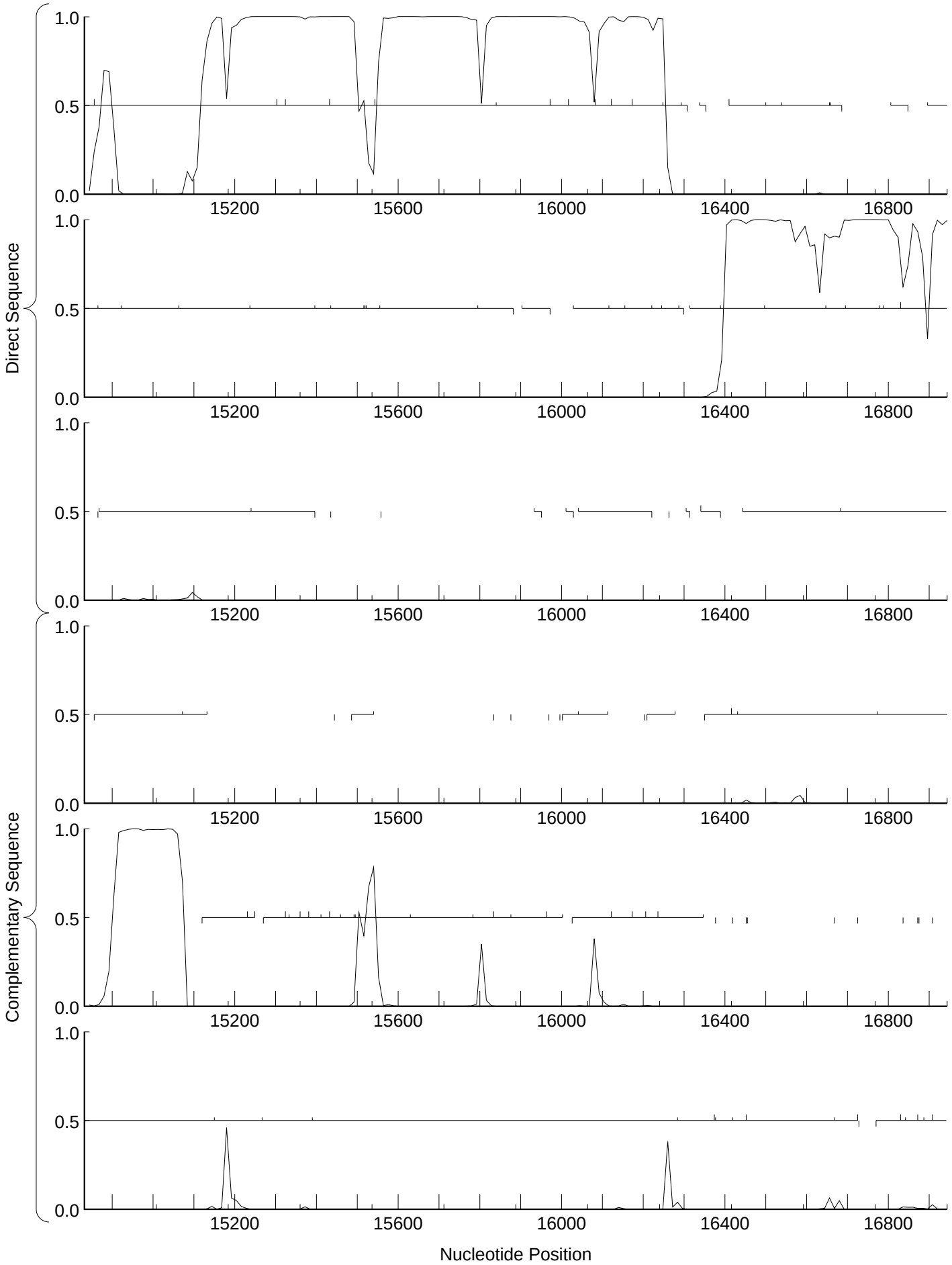
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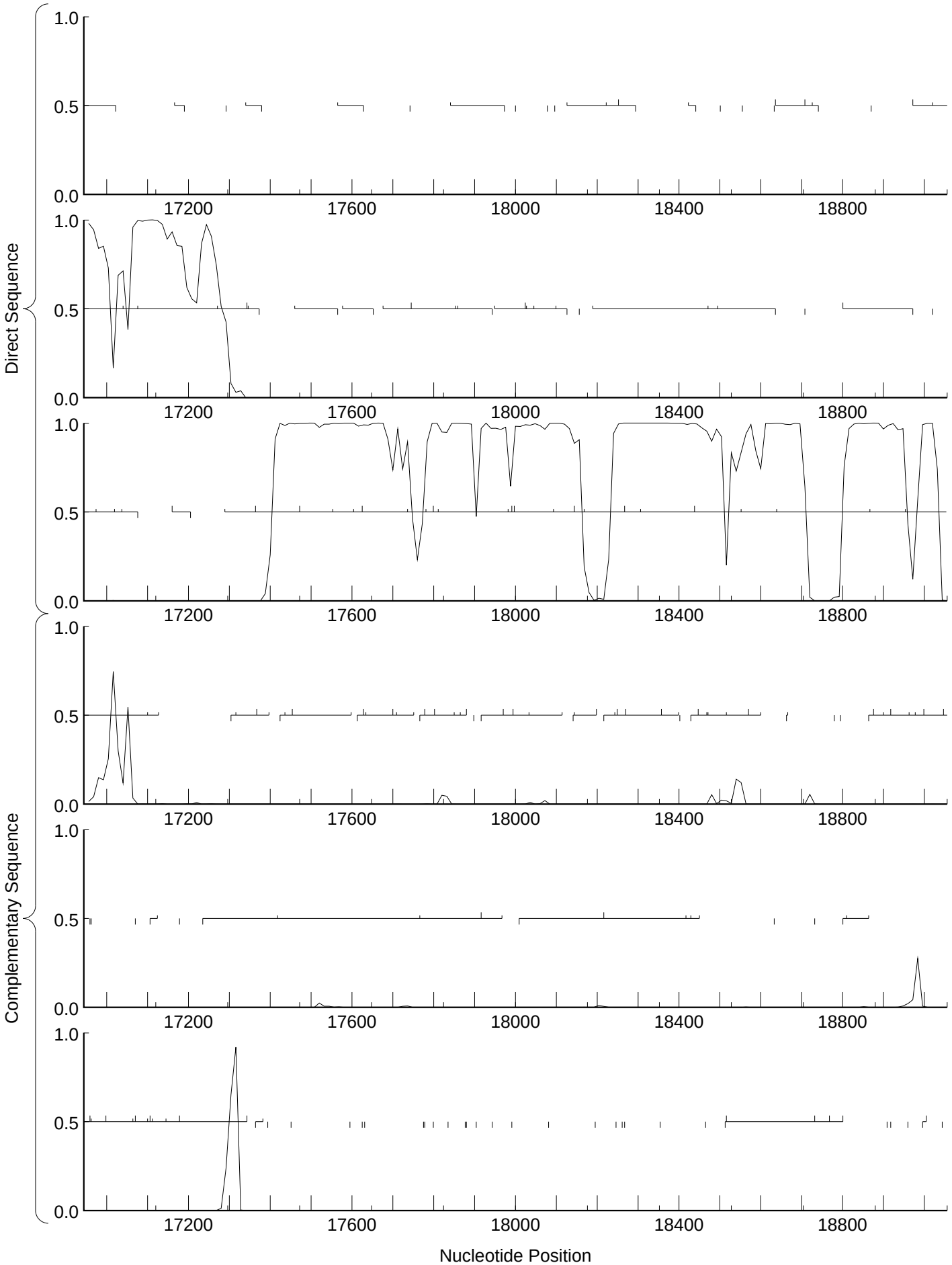
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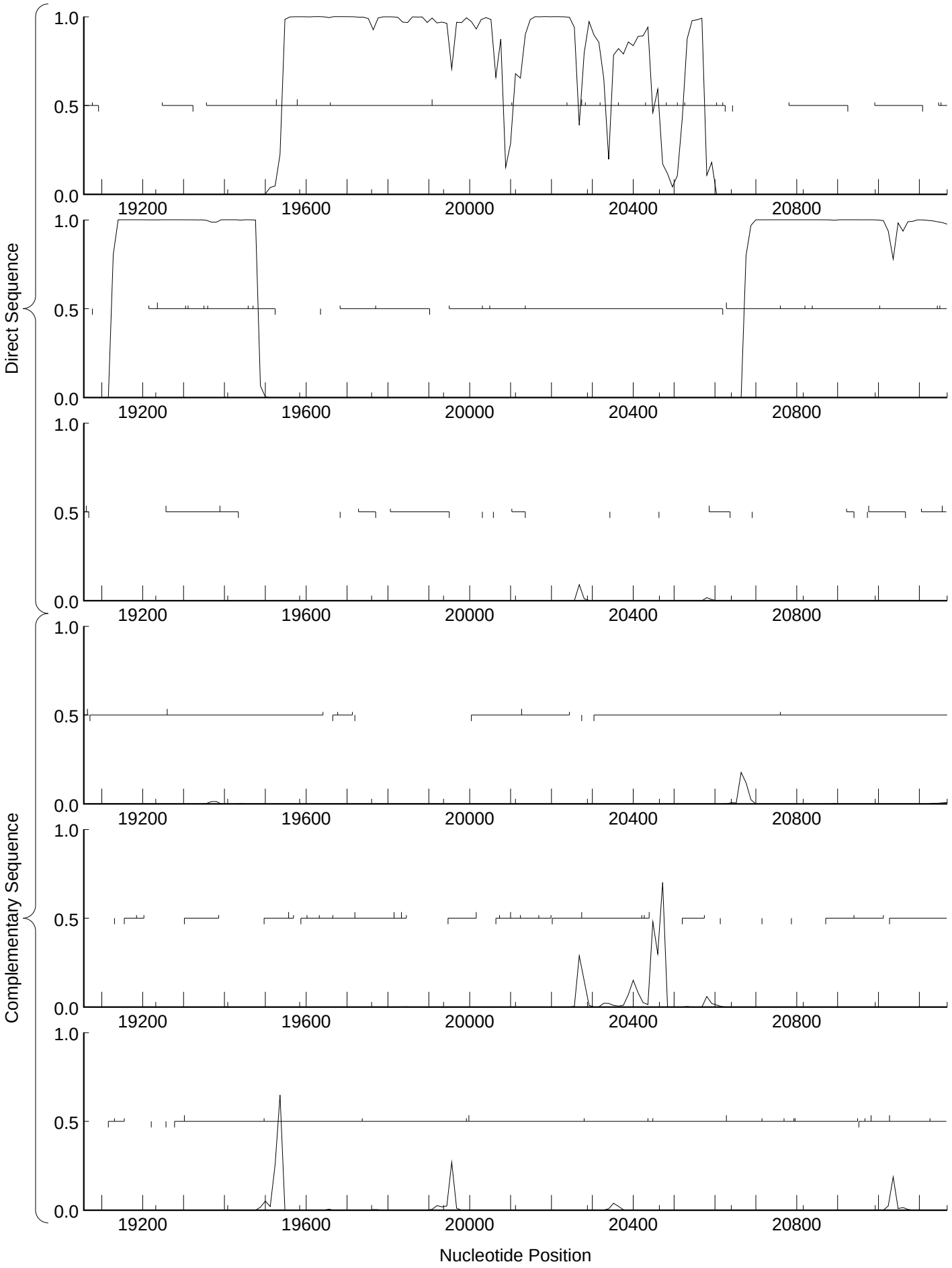
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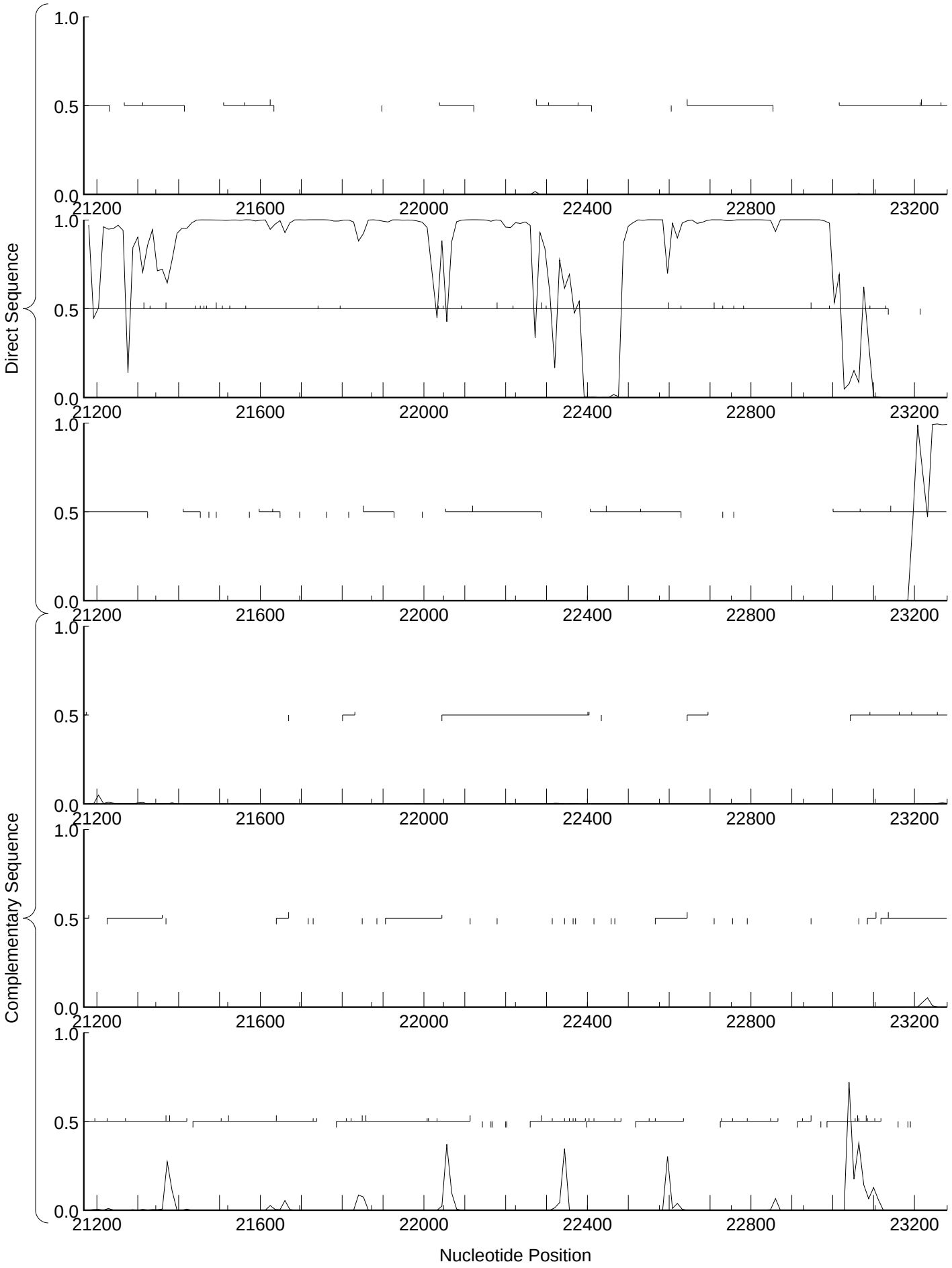
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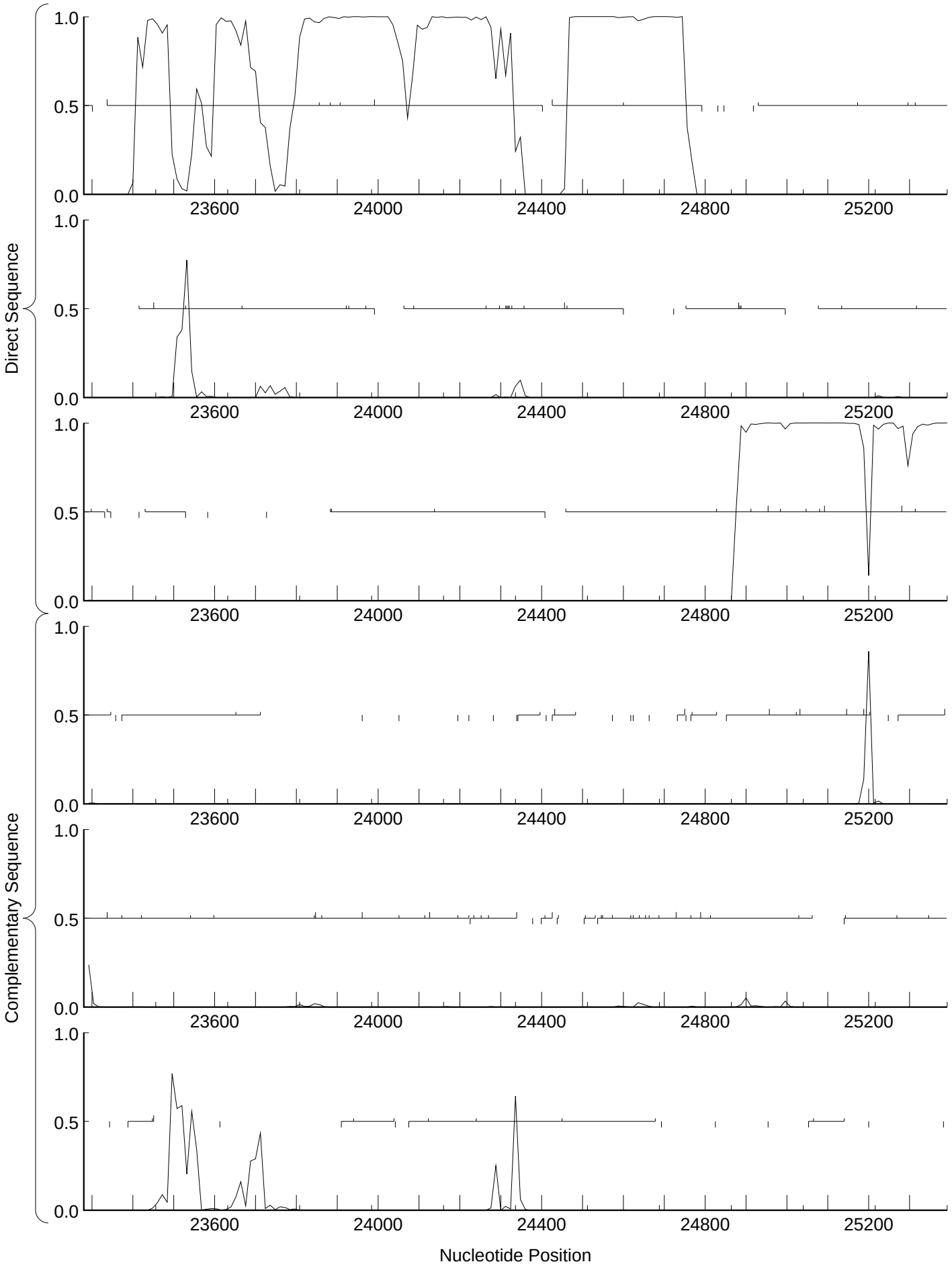
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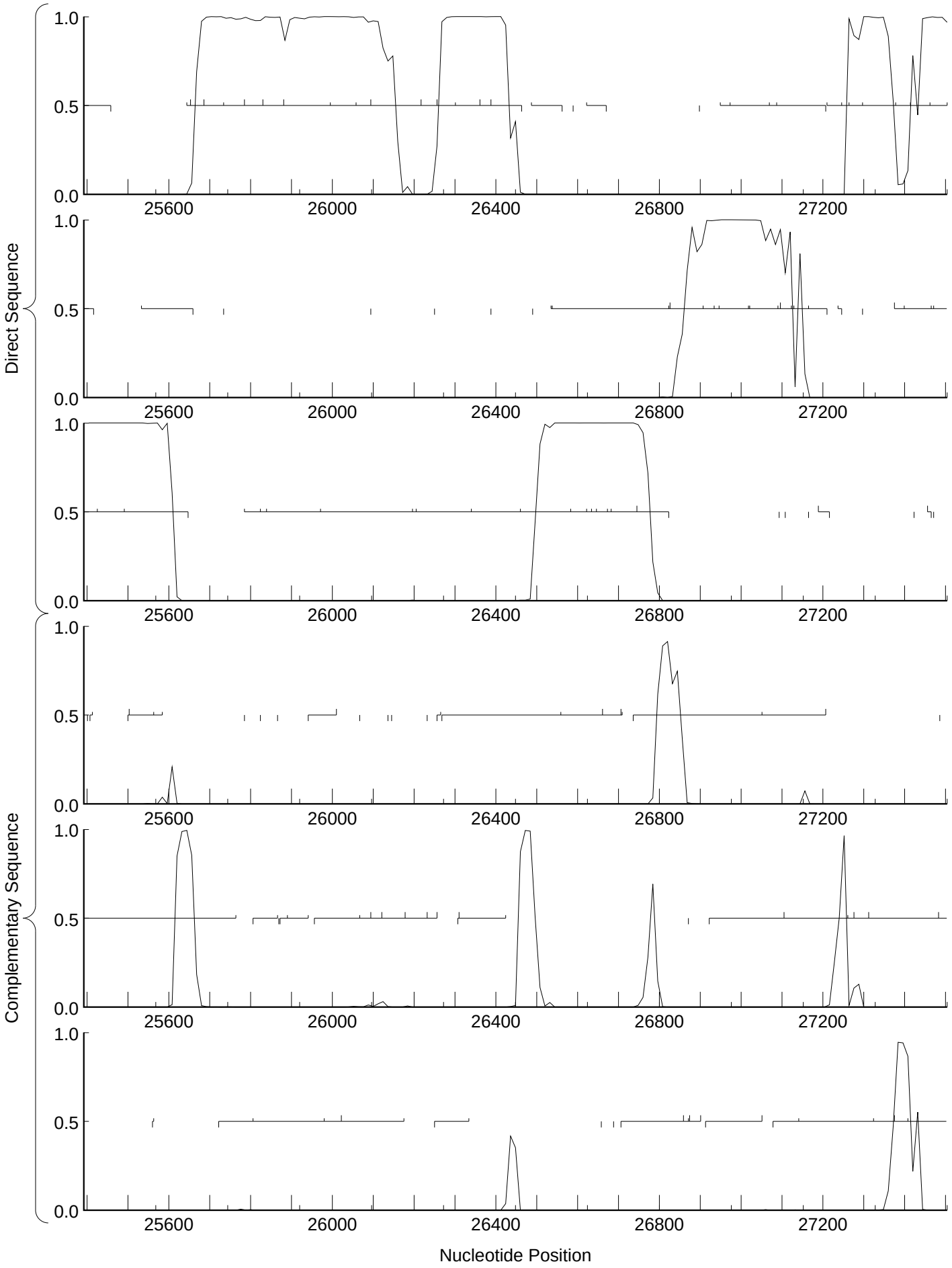
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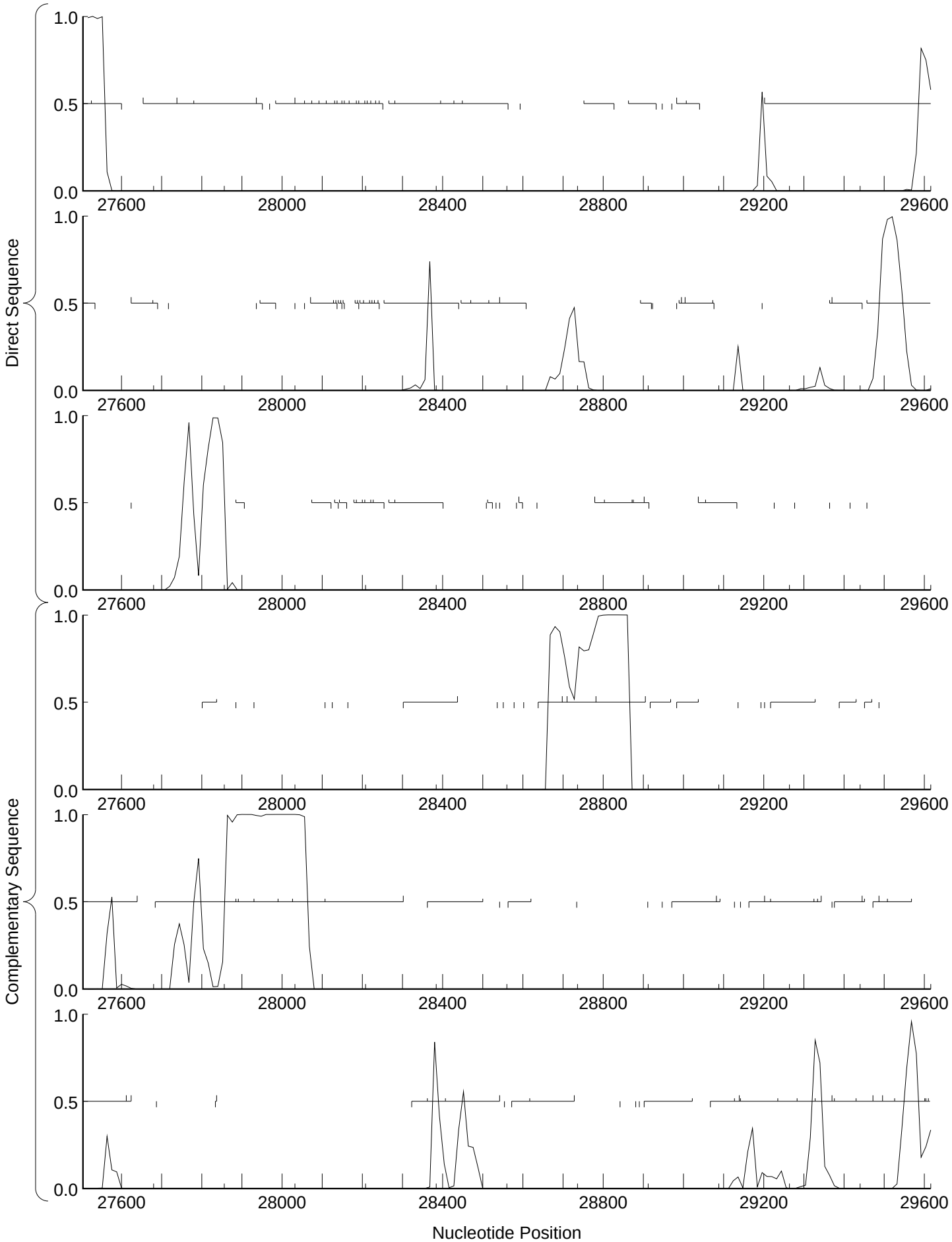
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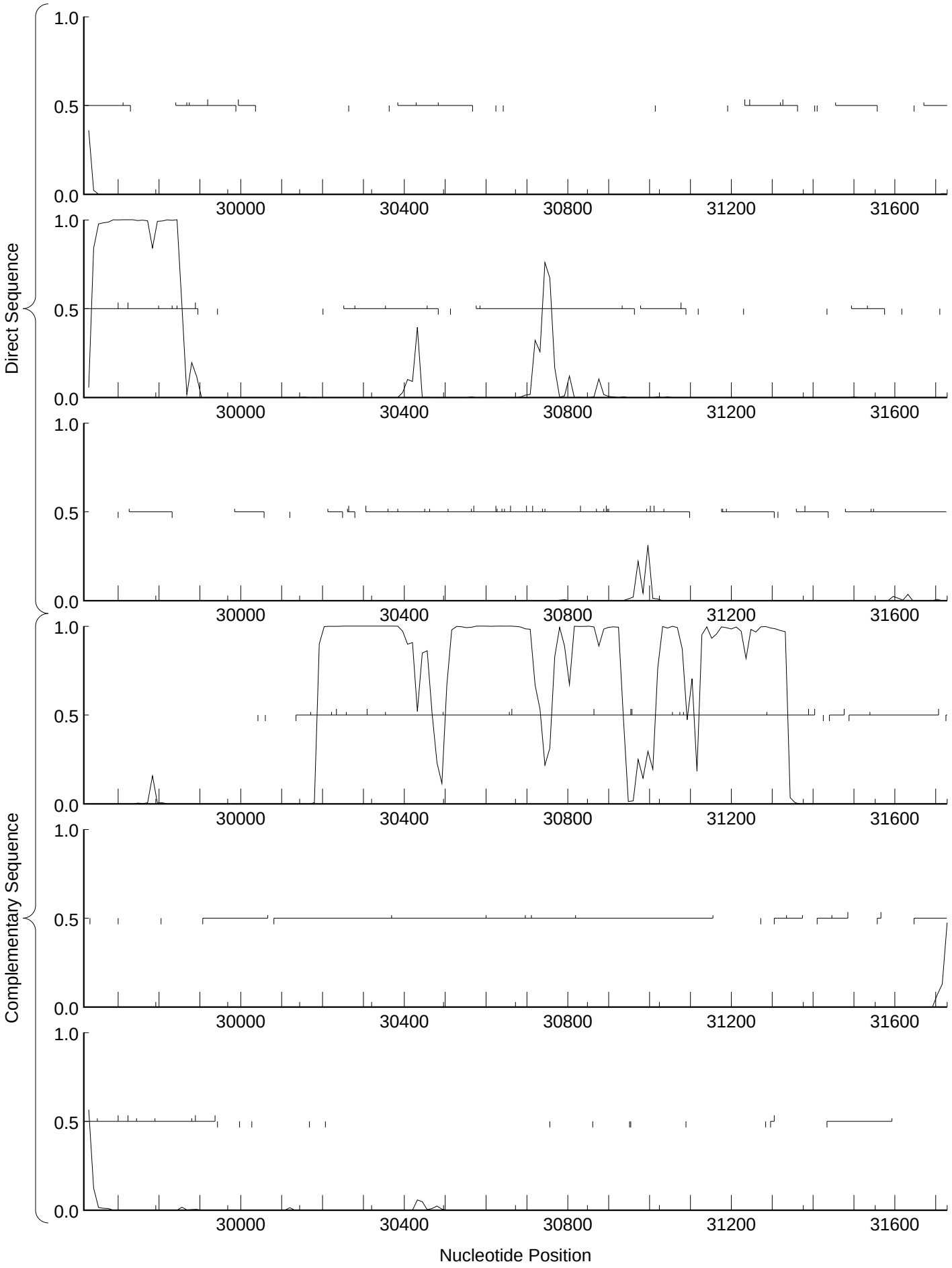
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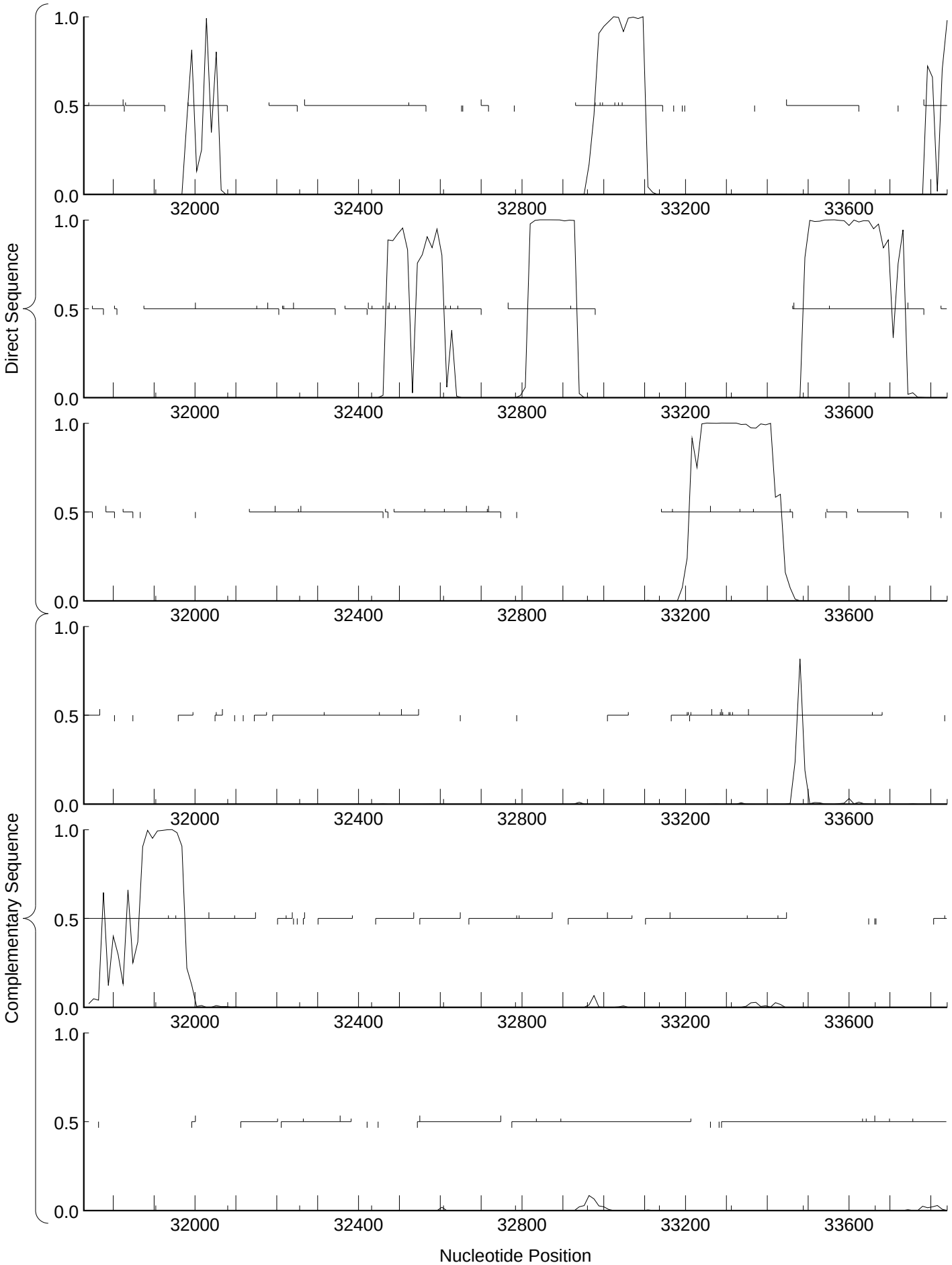
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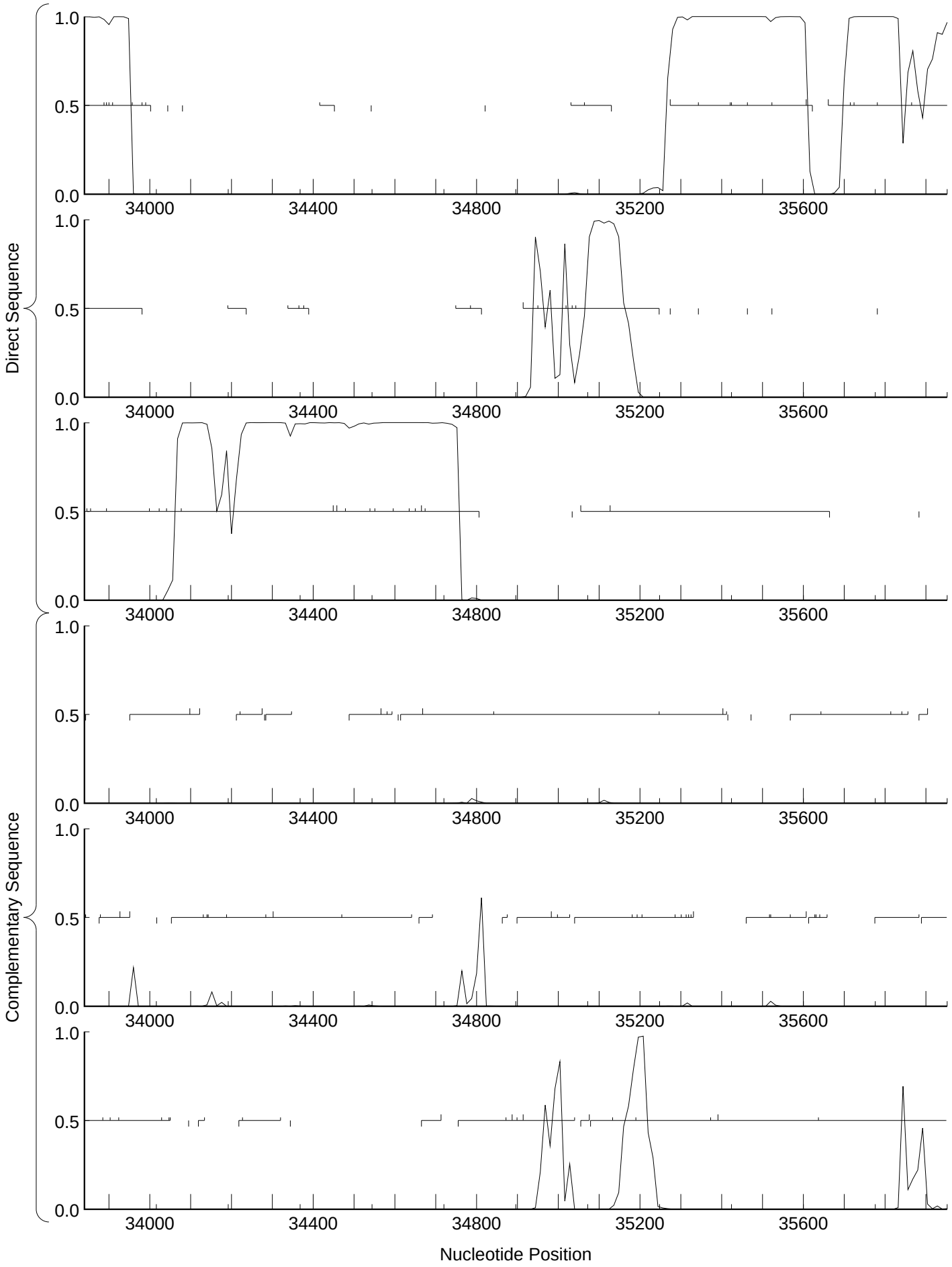
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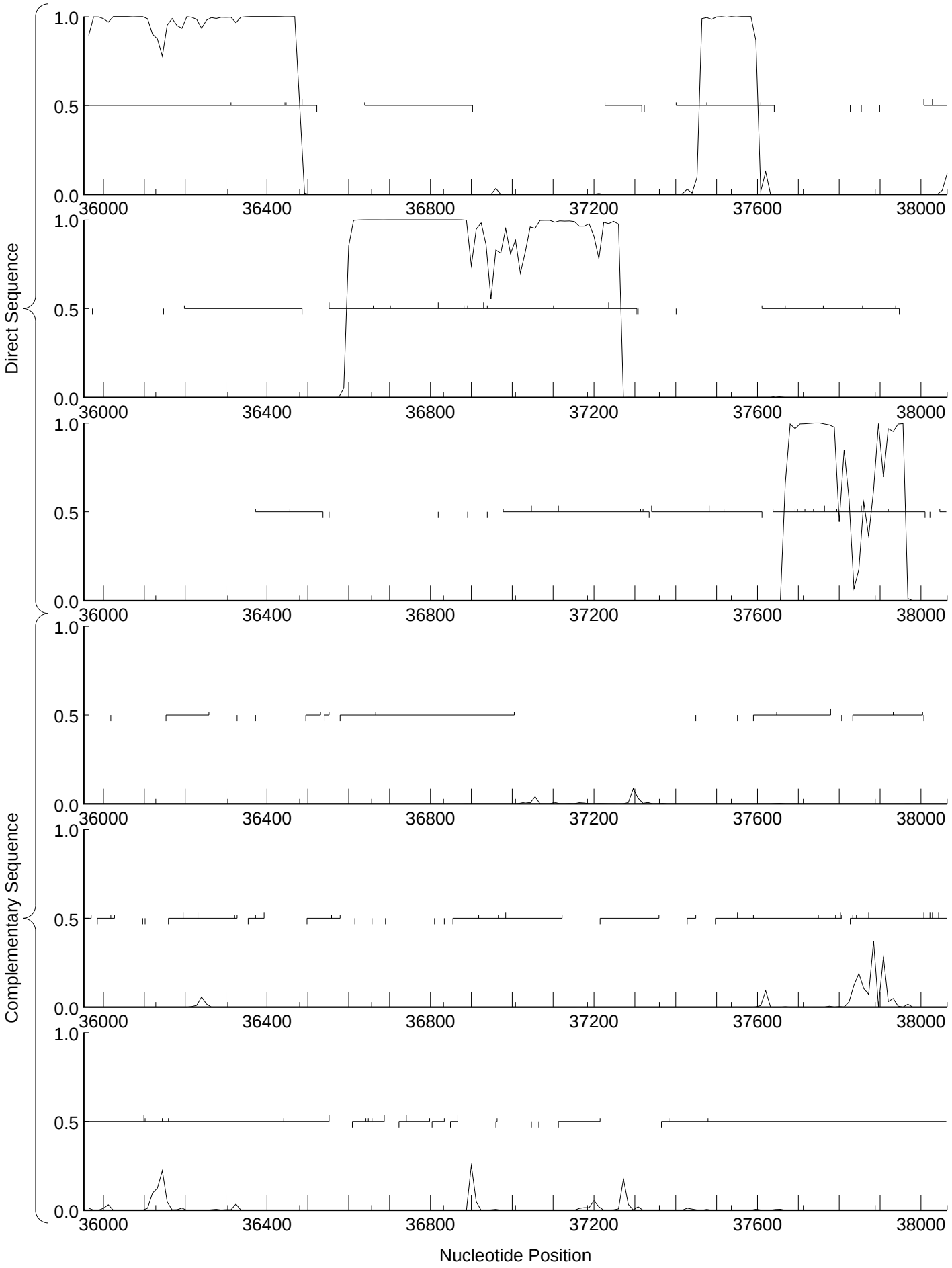
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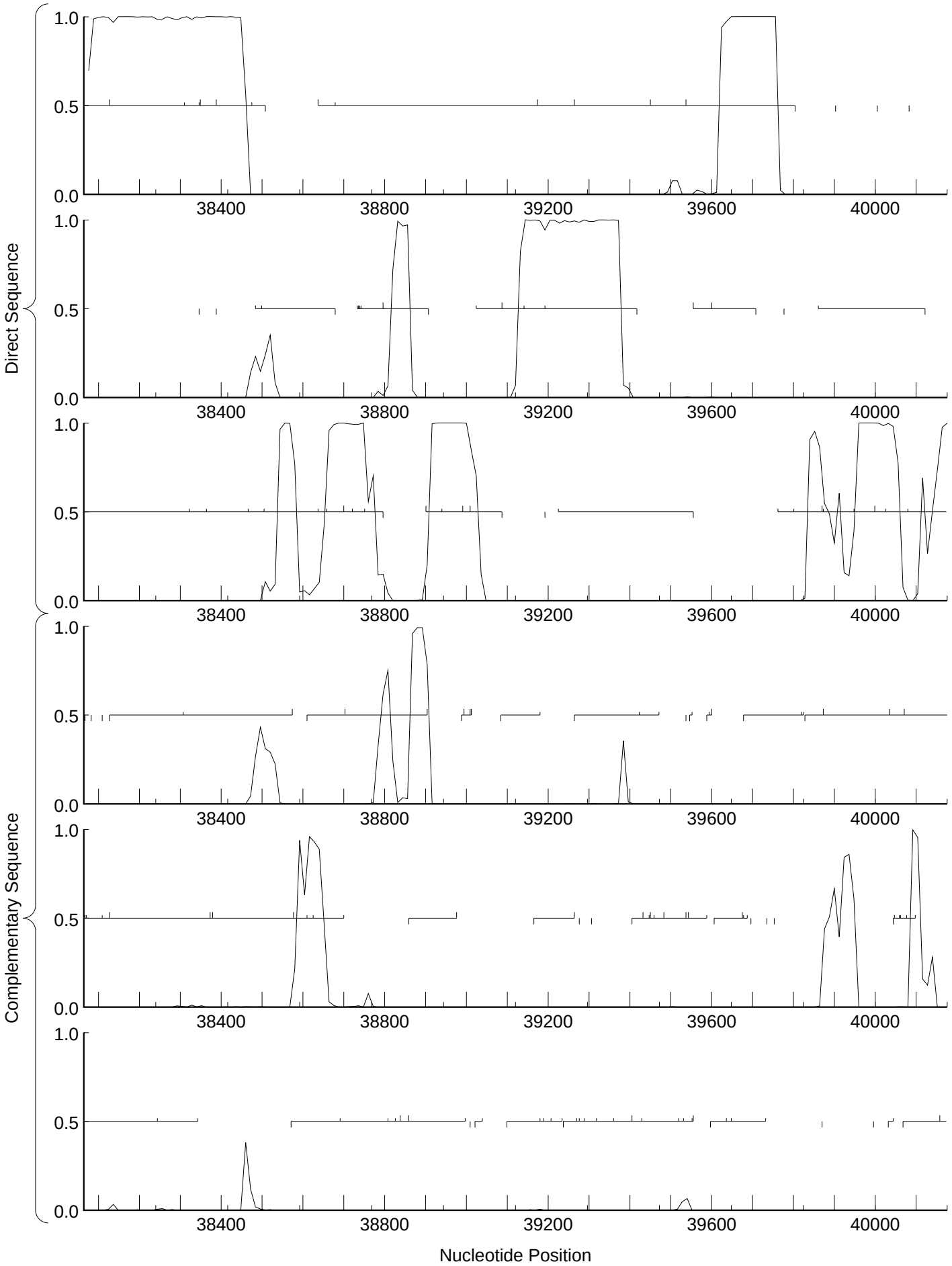
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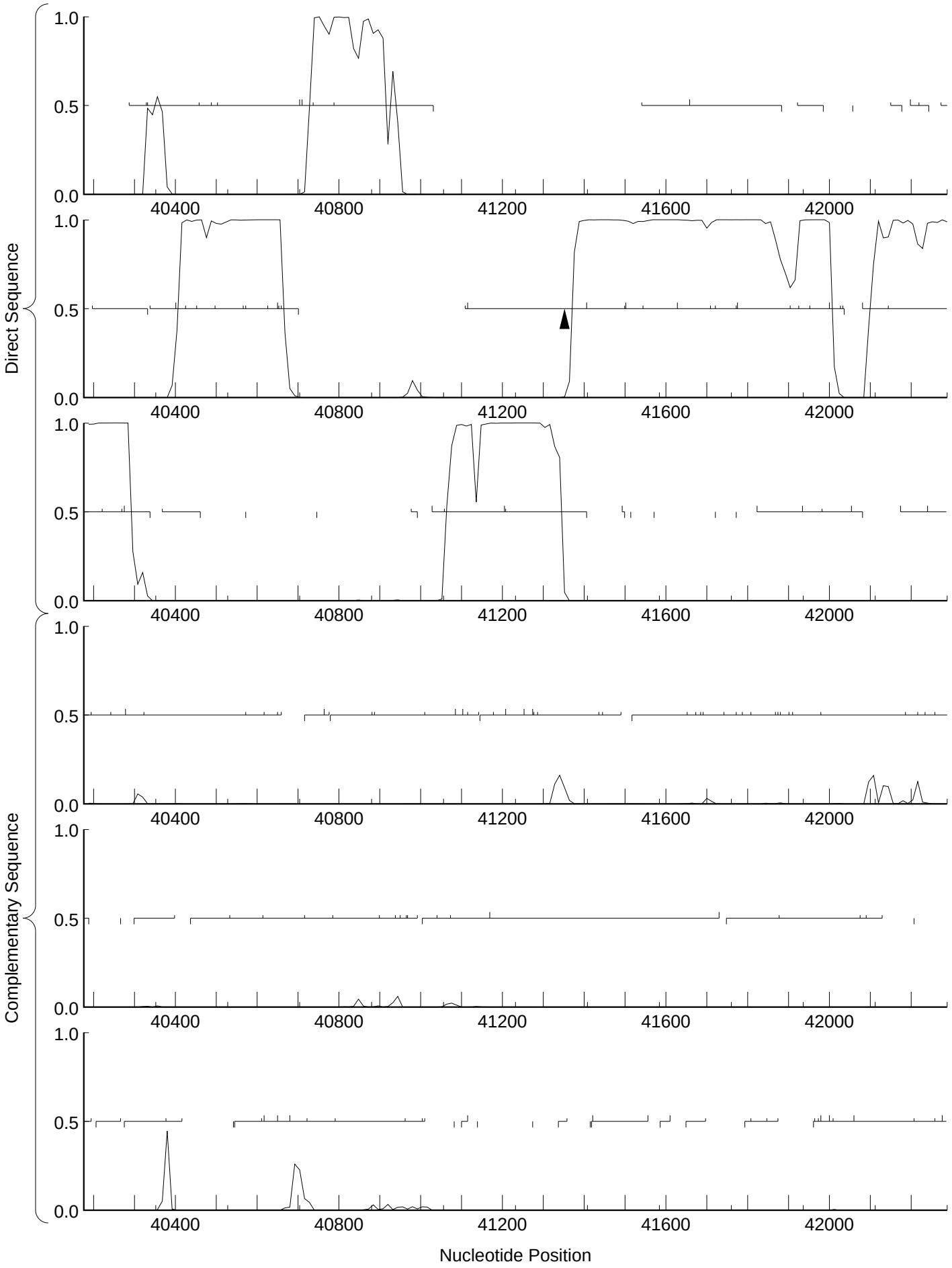
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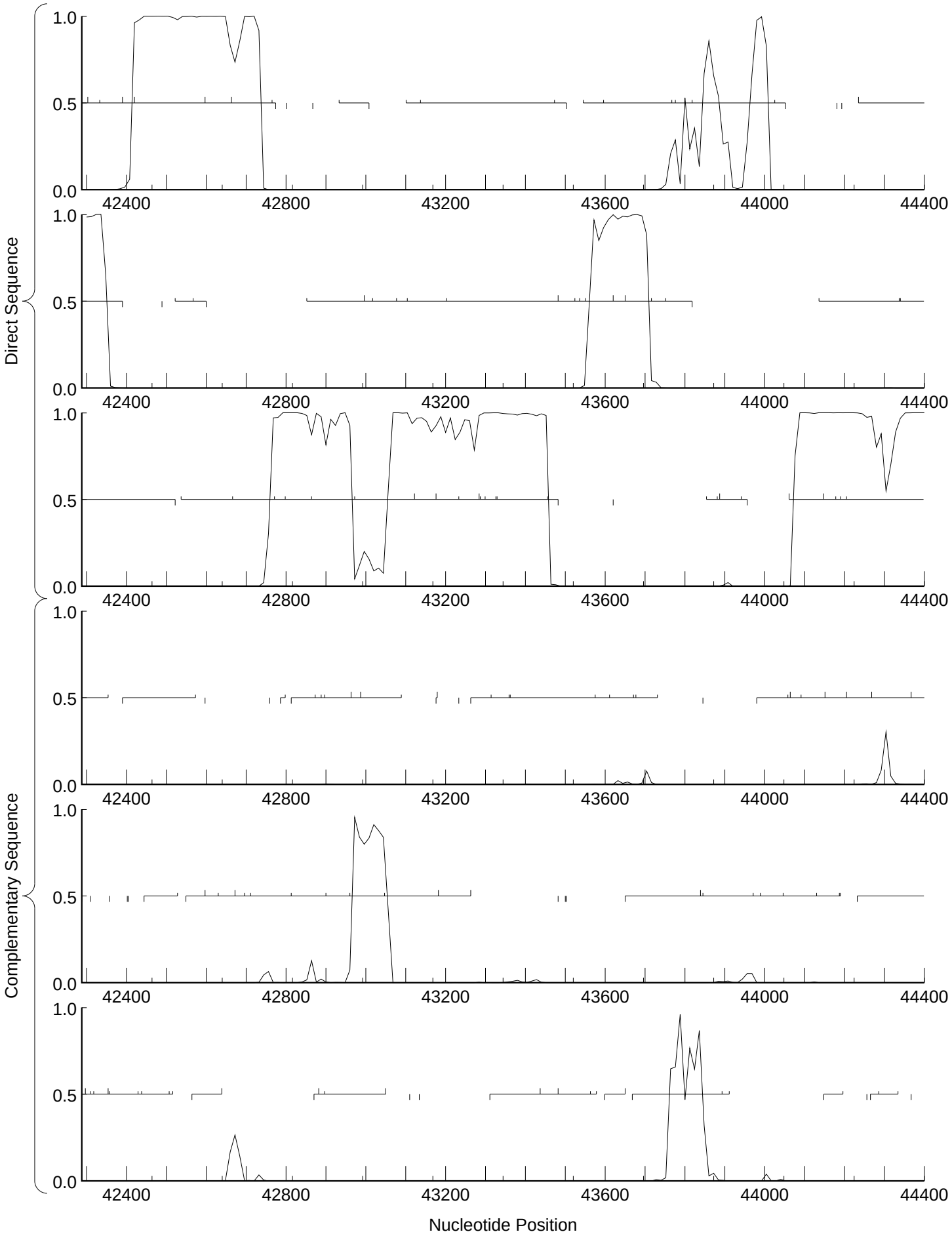
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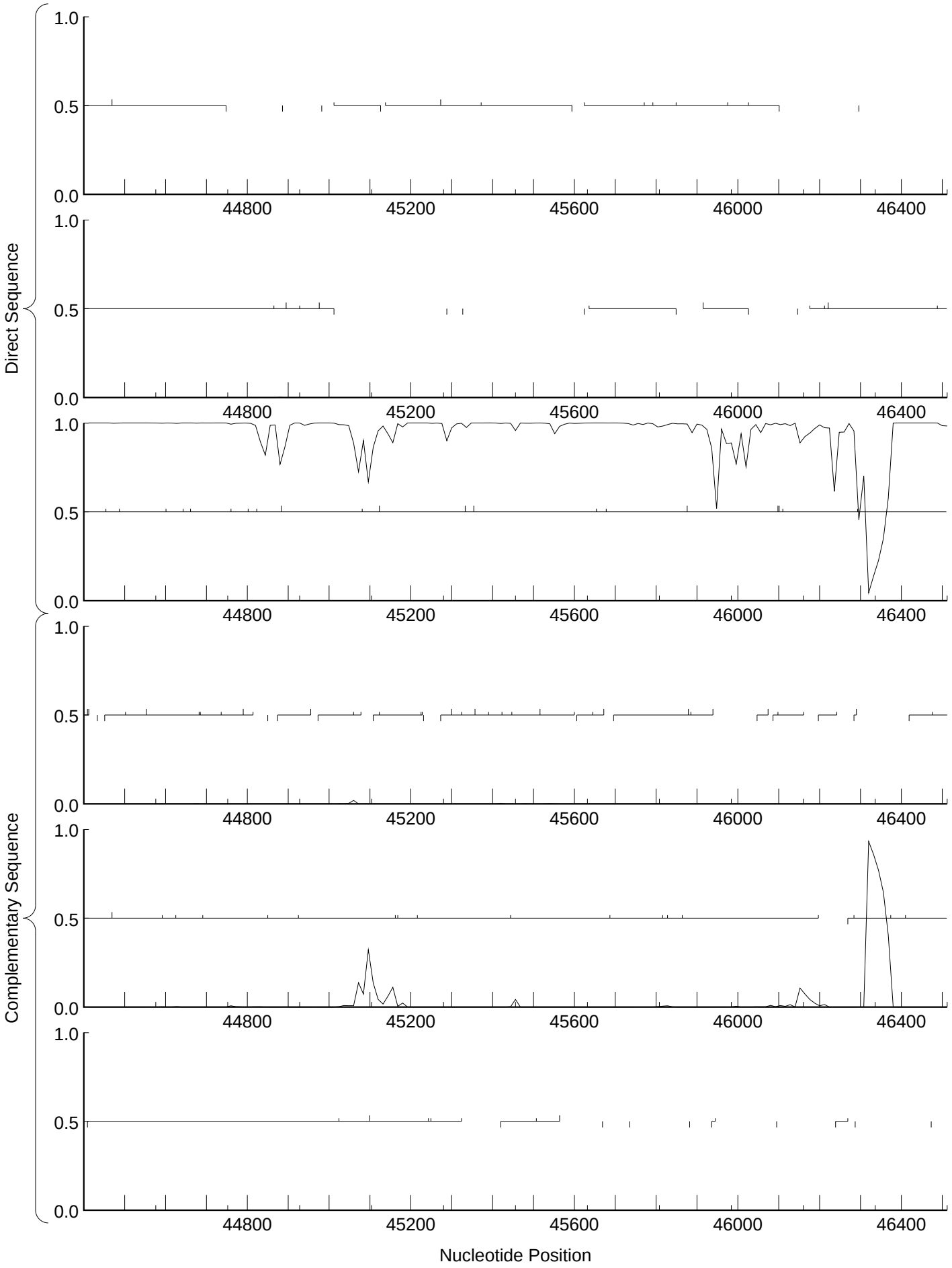
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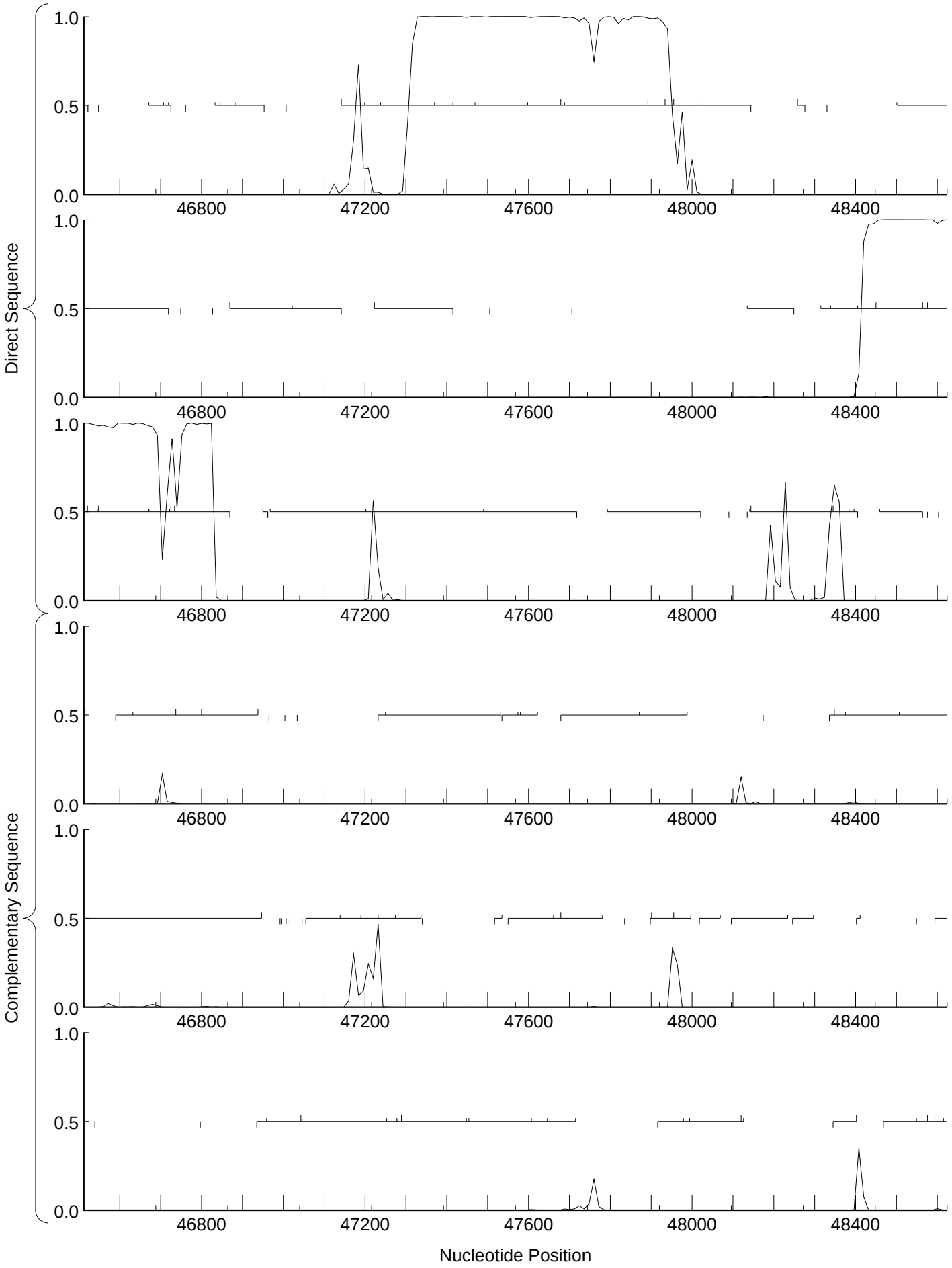
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s complete sequence, 50499 bp including 10-base 3’ overhang (TGCCGCGGTA), Cluster DI, Order 4, Window 96, Step 12, 23/25



s complete sequence, 50499 bp including 10-base 3' overhang (TGCCGCGGTA), Cluster DI, Order 4, Window 96, Step 12, 24/25



s complete sequence, 50499 bp including 10-base 3’ overhang (TGCCGCGGTA), Cluster DI, Order 4, Window 96, Step 12, 25/25

