GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Arthrobacter phage Timinator complete sequence, 51138 bp, circularly permuted, Cluster AO2
Analysis Date    :  3/29/24 at 4:47:04
Pages            :  26
Sequence Length  :  51138 bp
GC Content       :   60.92%

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

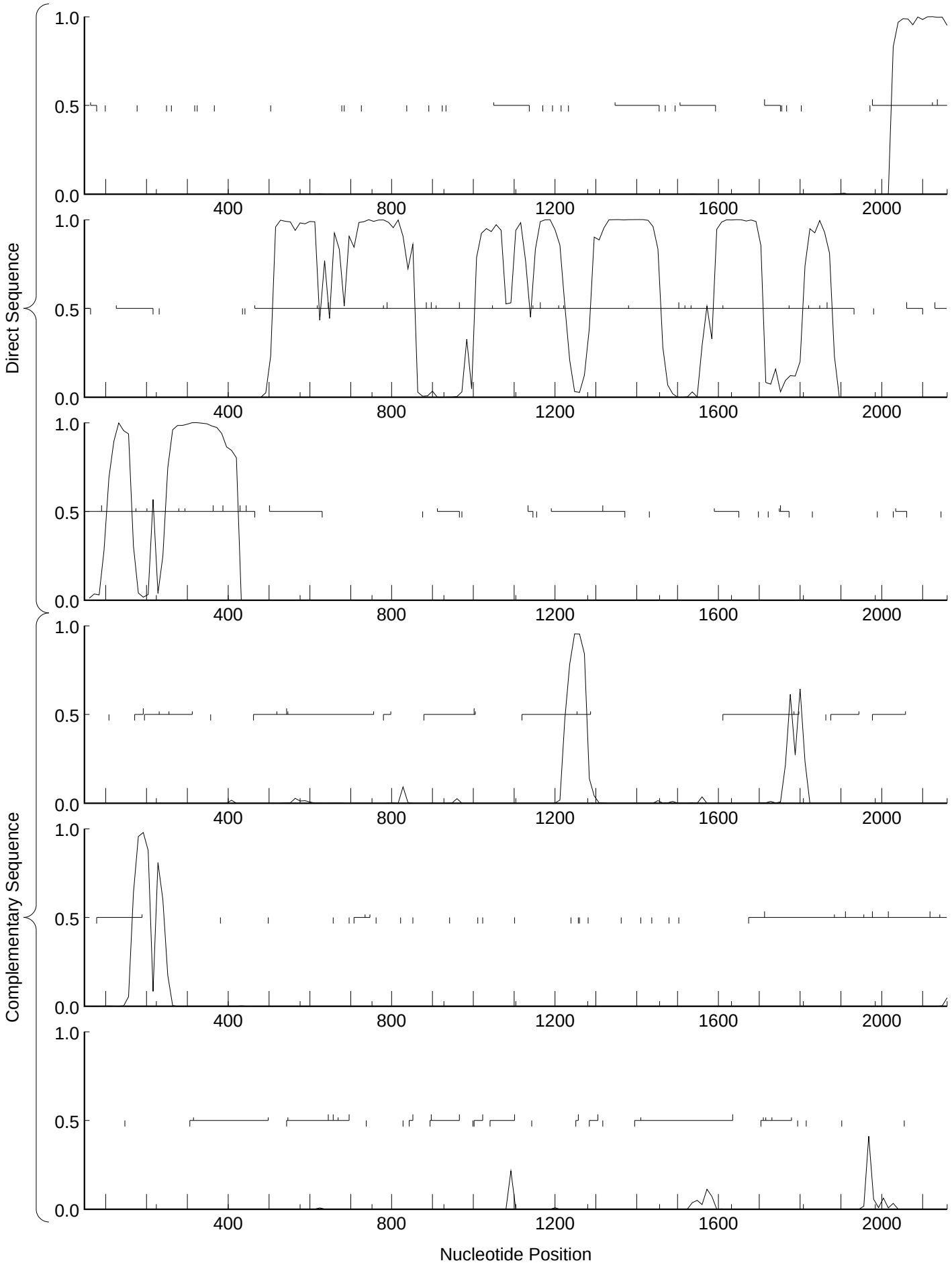
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Order      :  4
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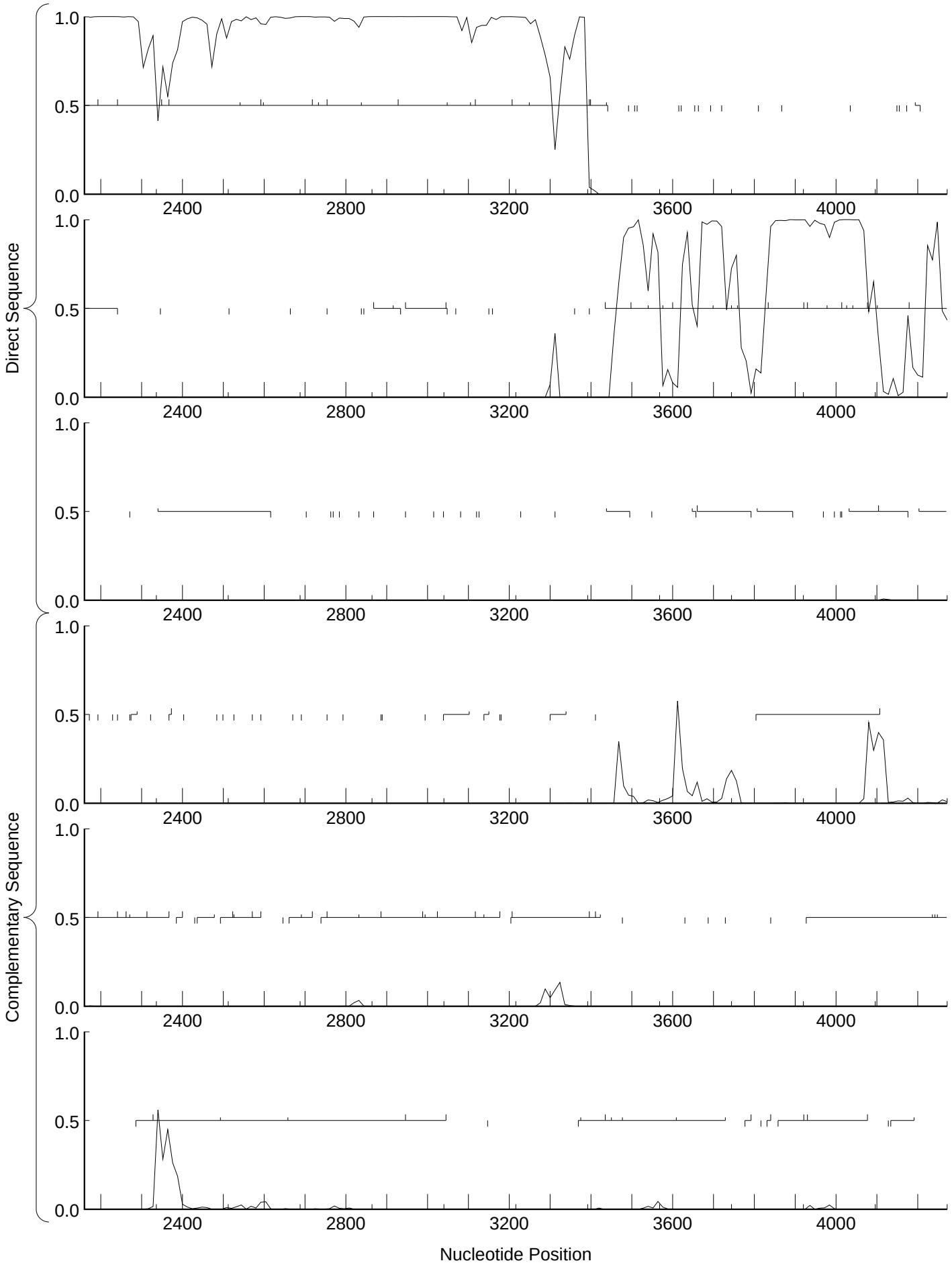

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

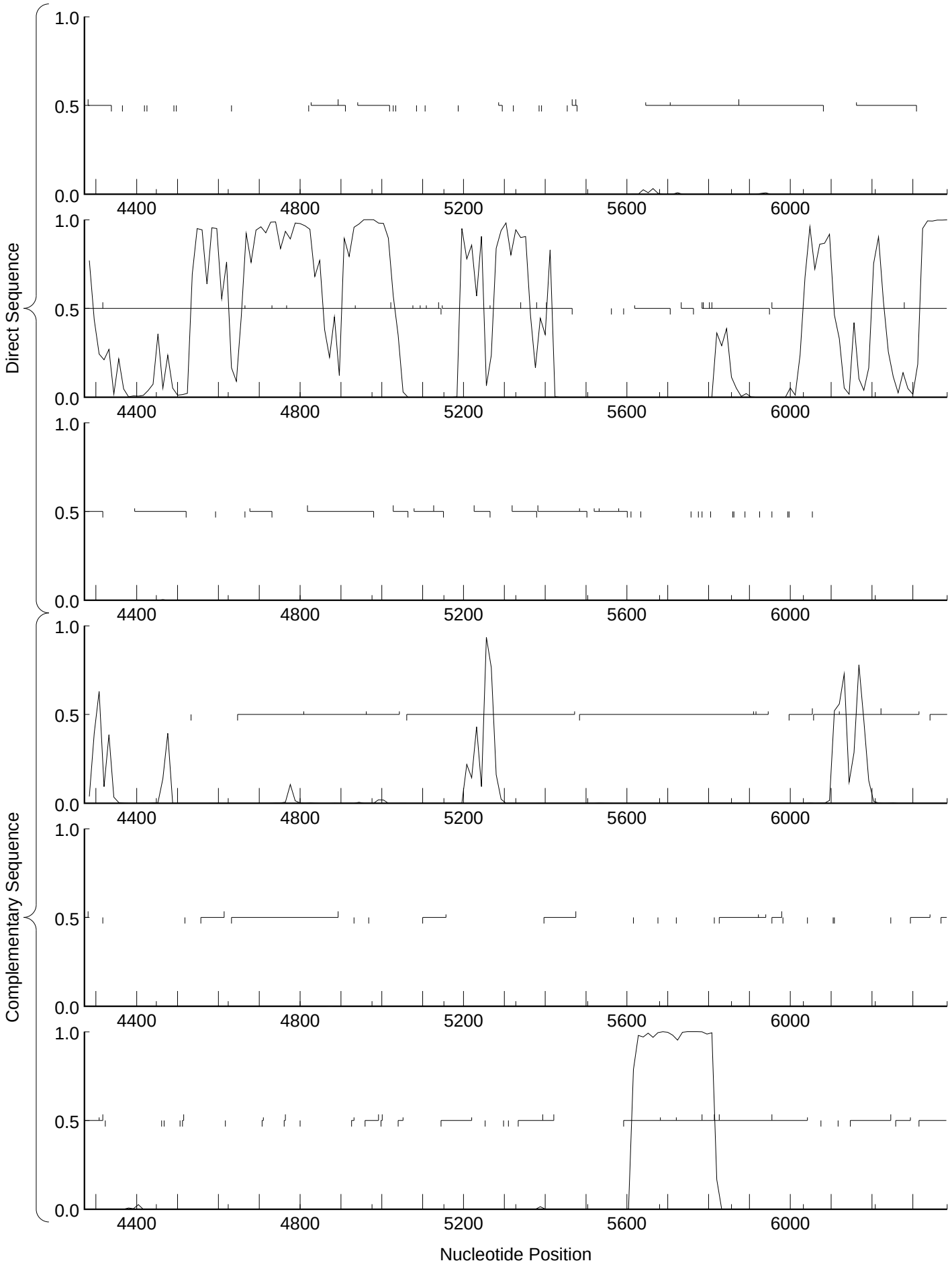
Arthrobacter phage Timinator complete sequence, 51138 bp, circularly permuted, Cluster AO2, Order 4, Window 96, Step 12, 2/26



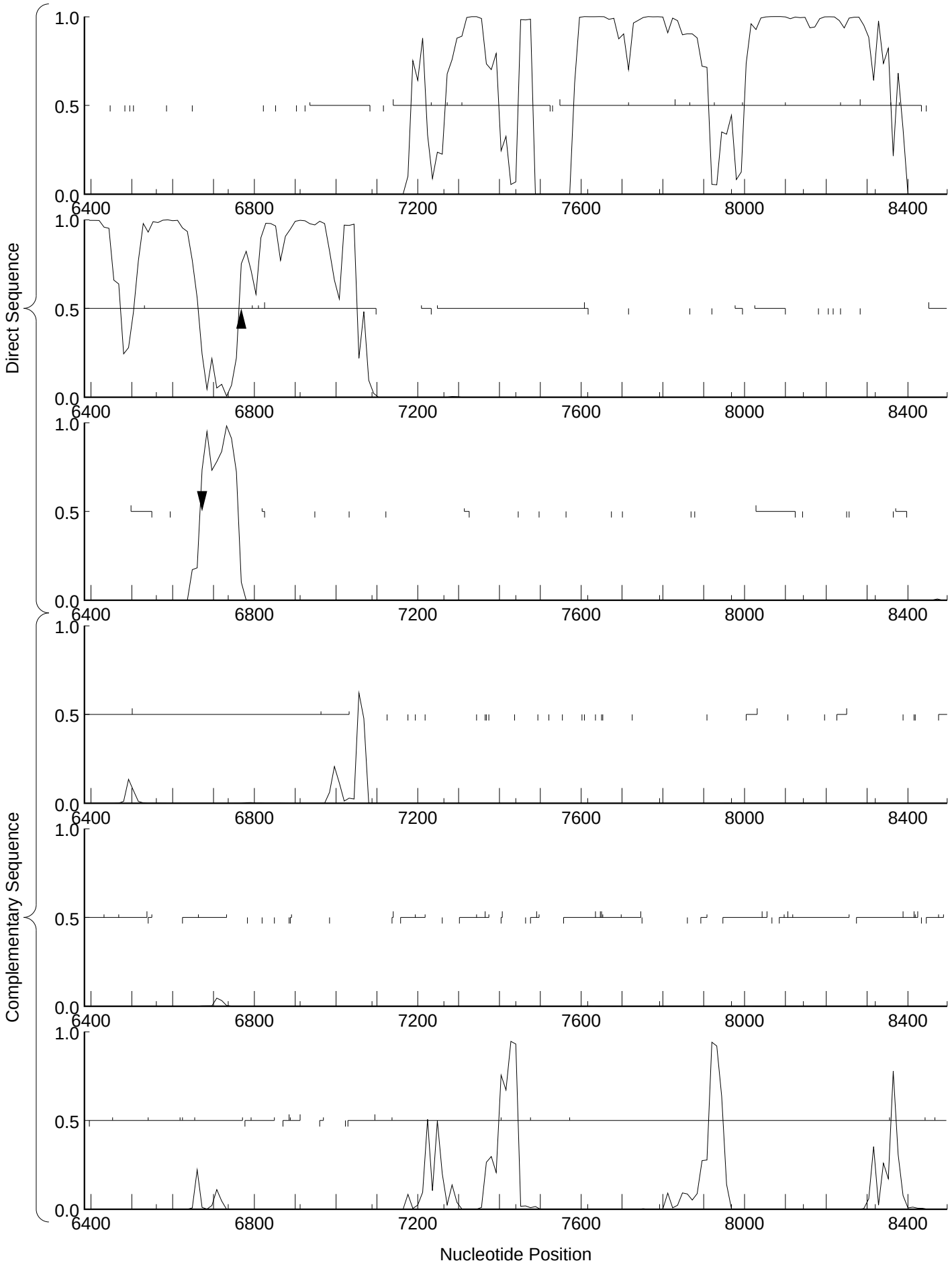
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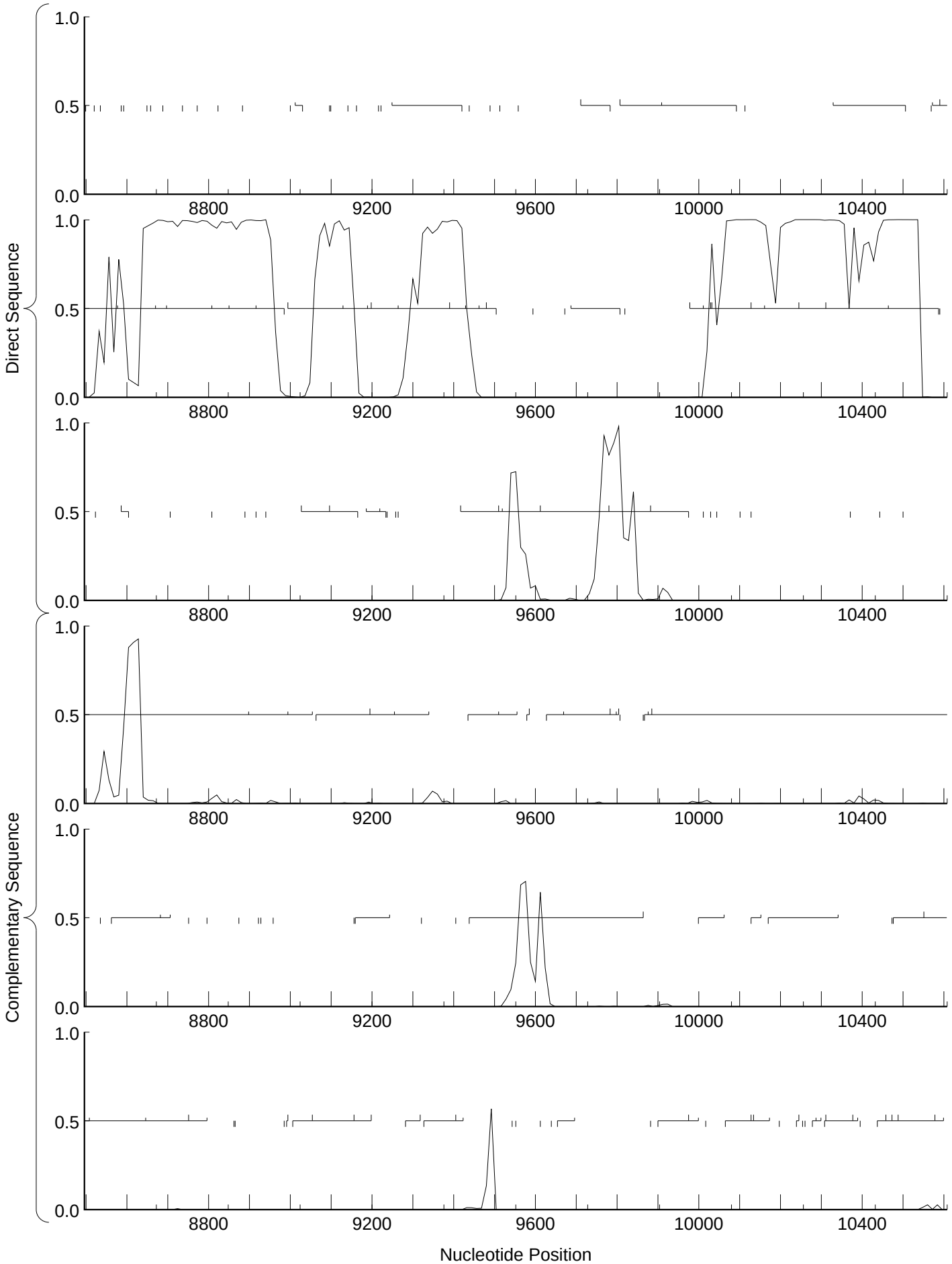
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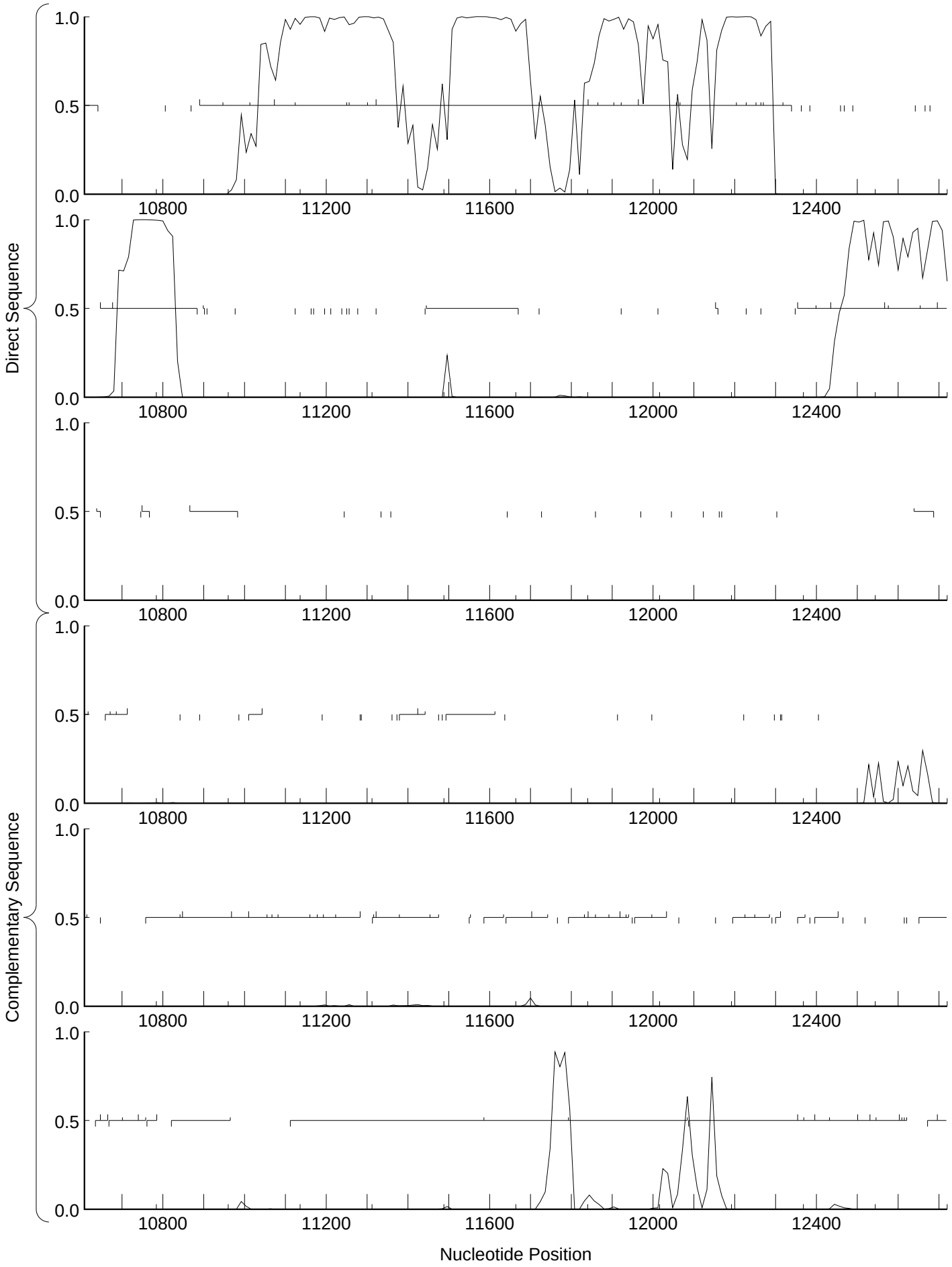
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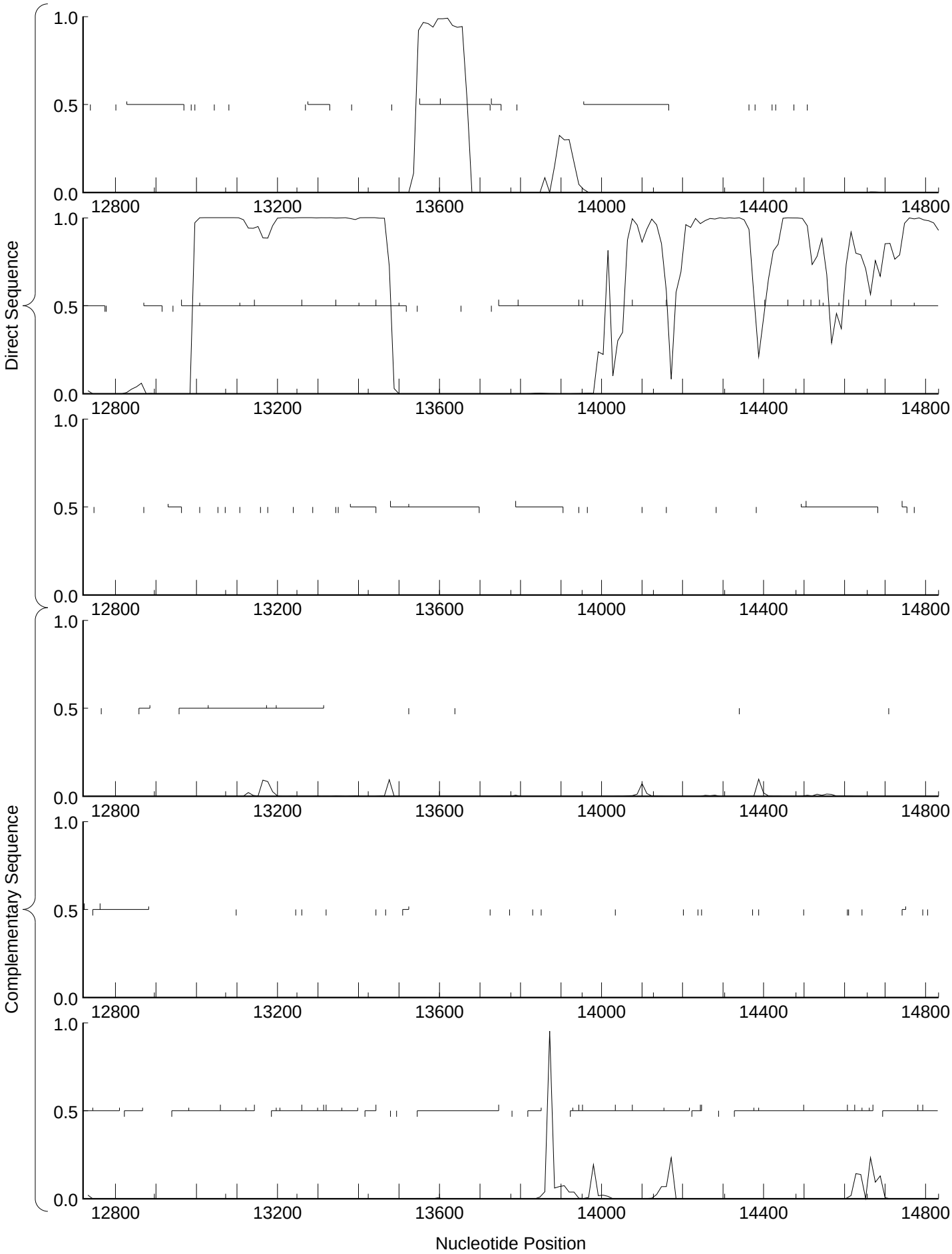
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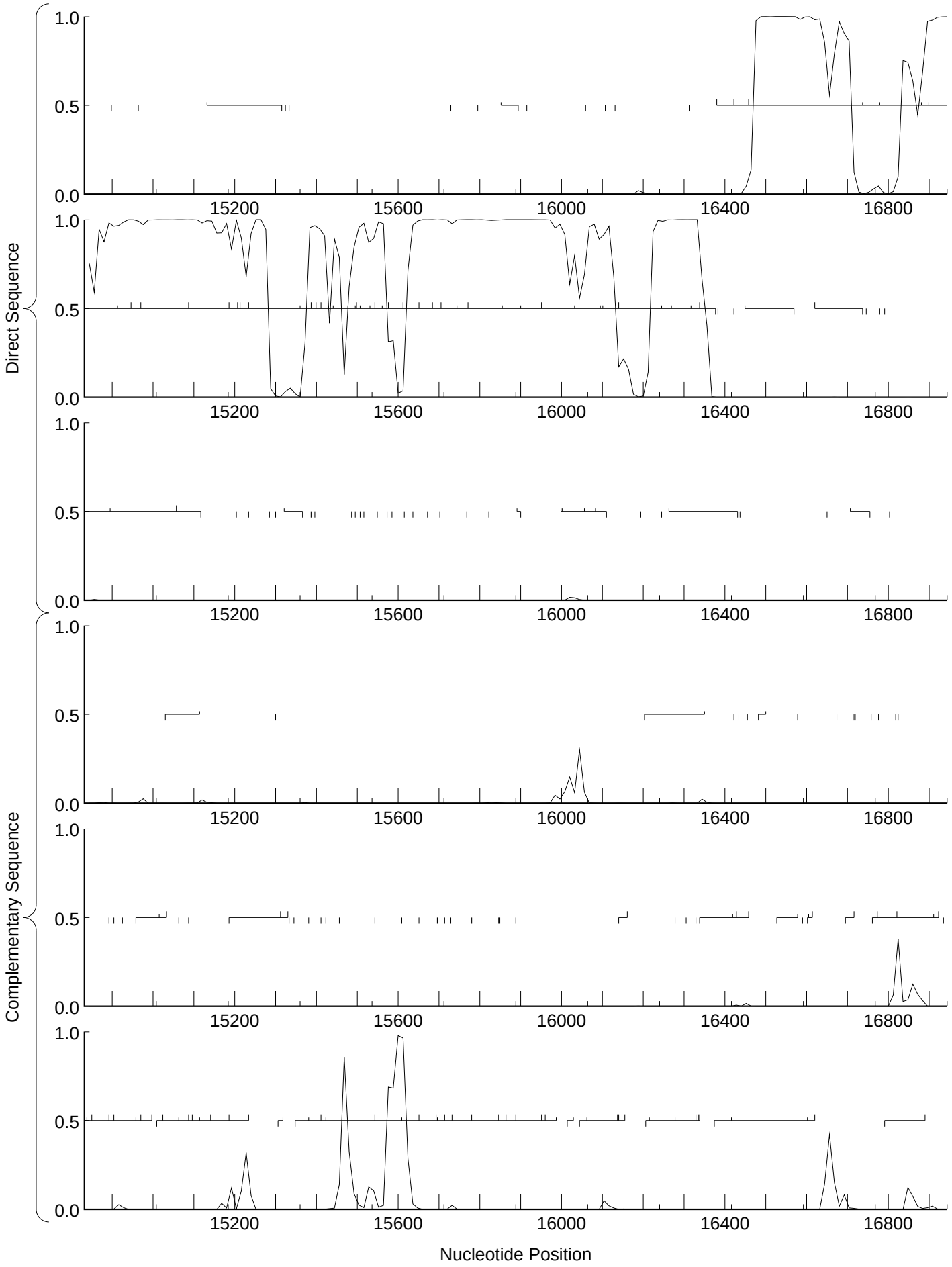
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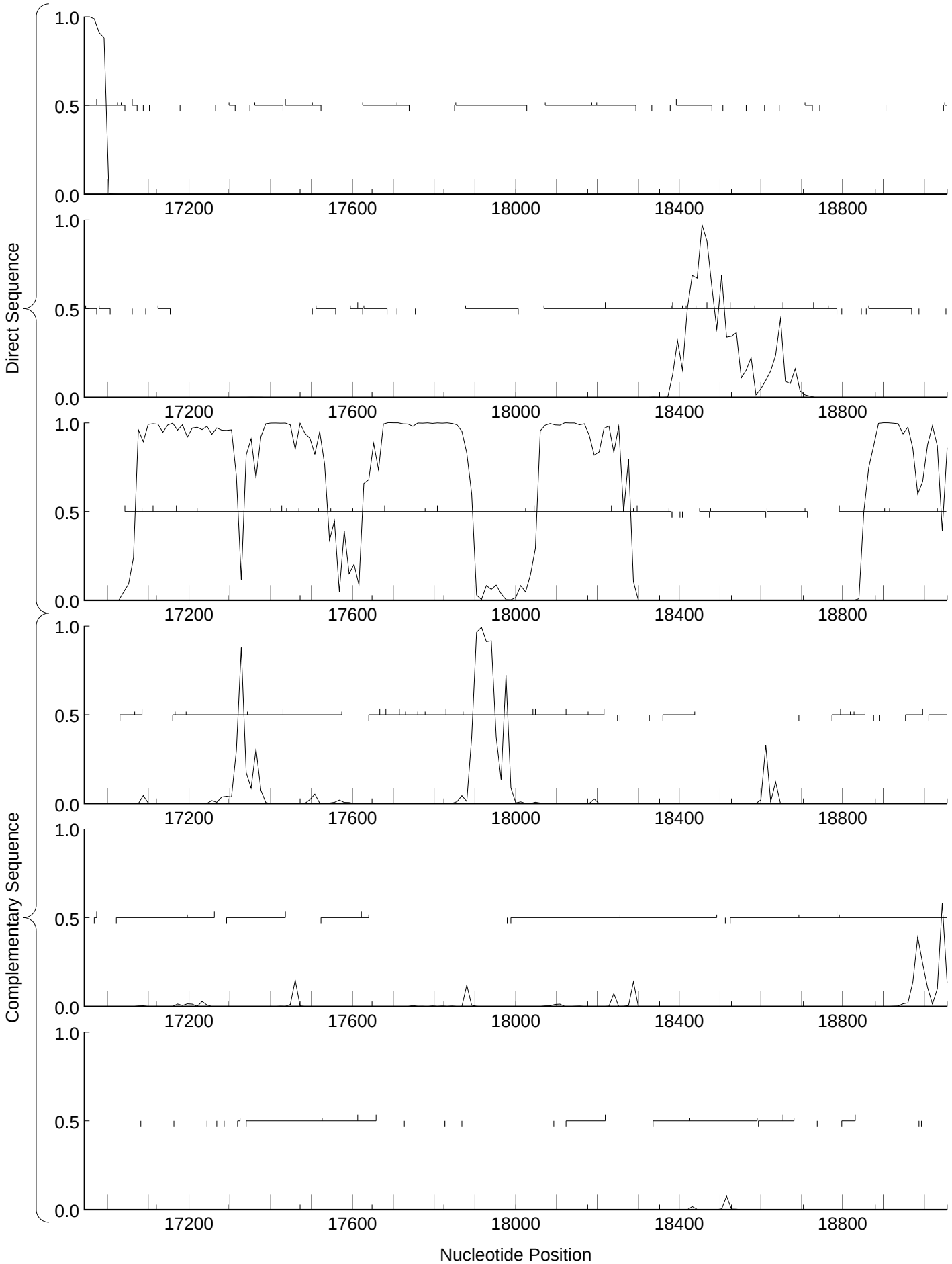
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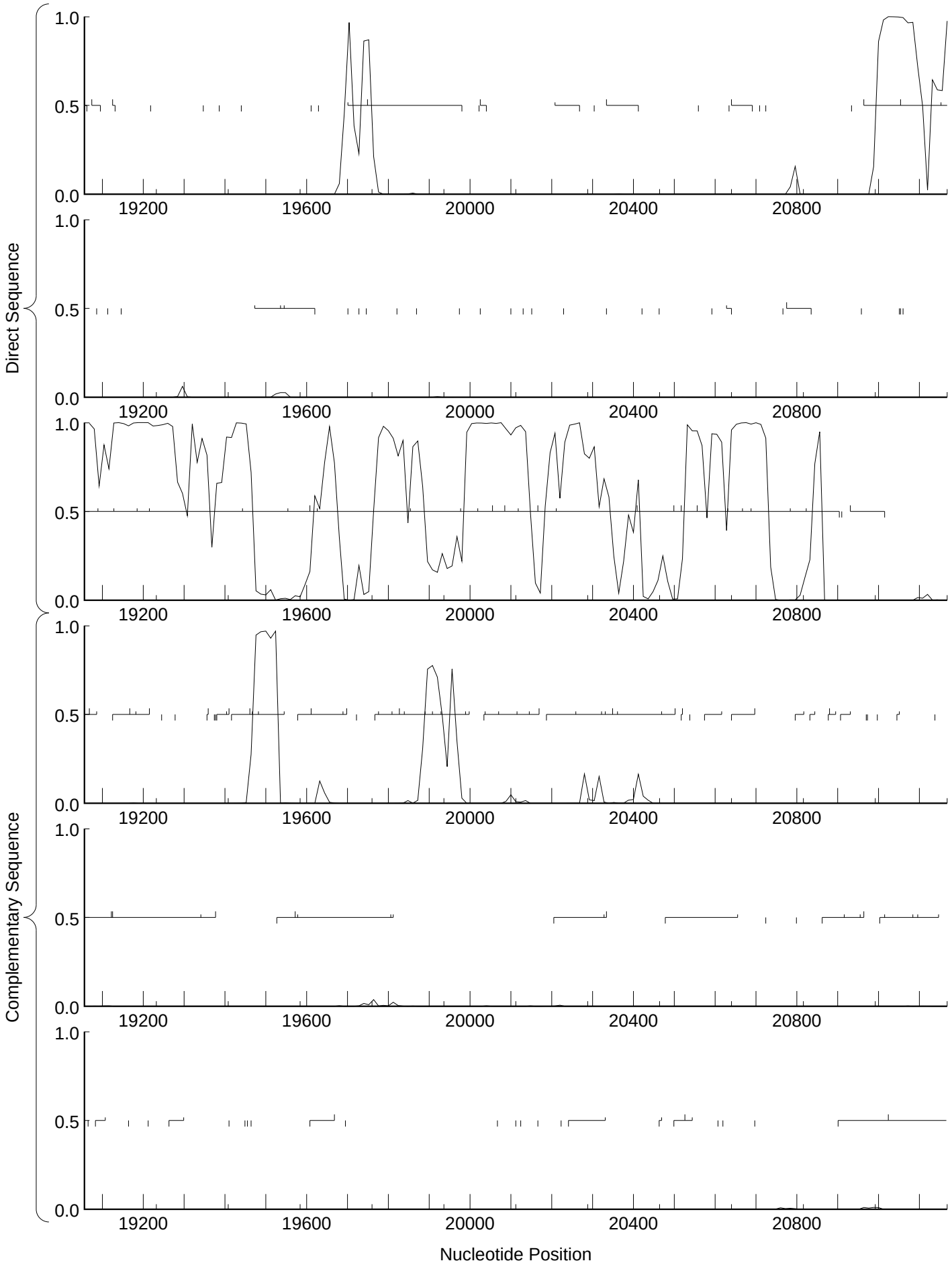
Arthrobacter phage Timinator complete sequence, 51138 bp, circularly permuted, Cluster AO2, Order 4, Window 96, Step 12, 9/26



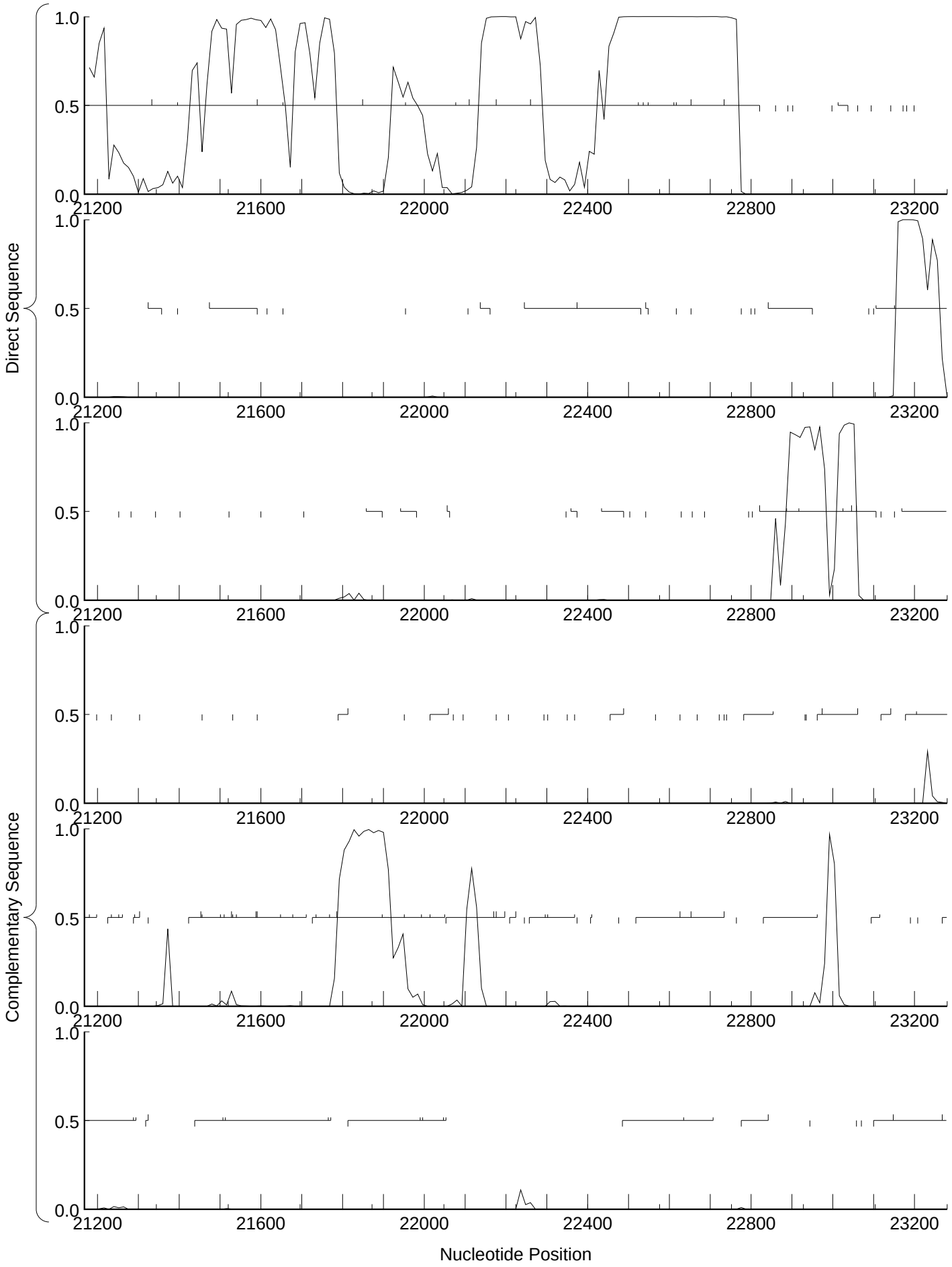
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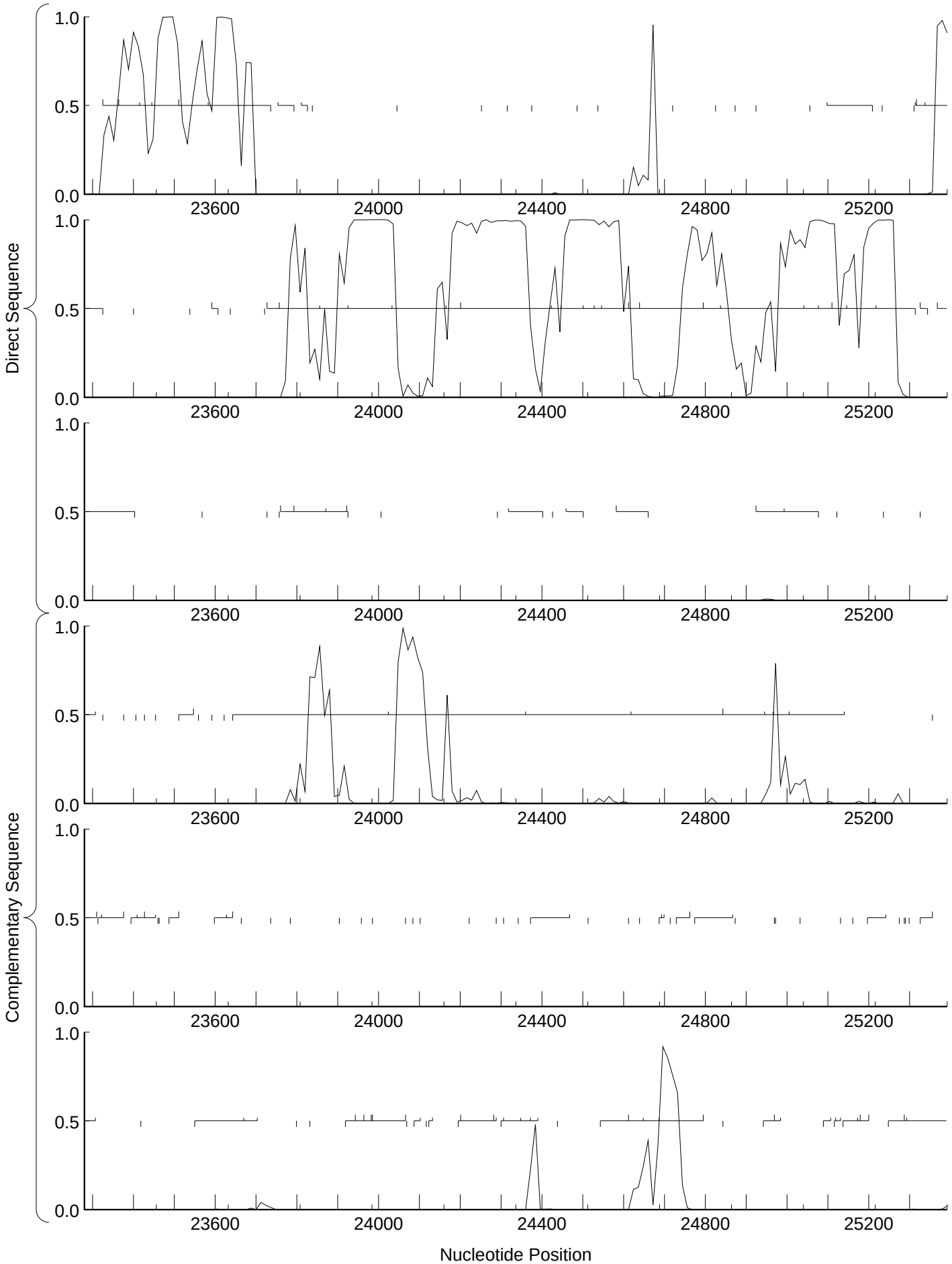
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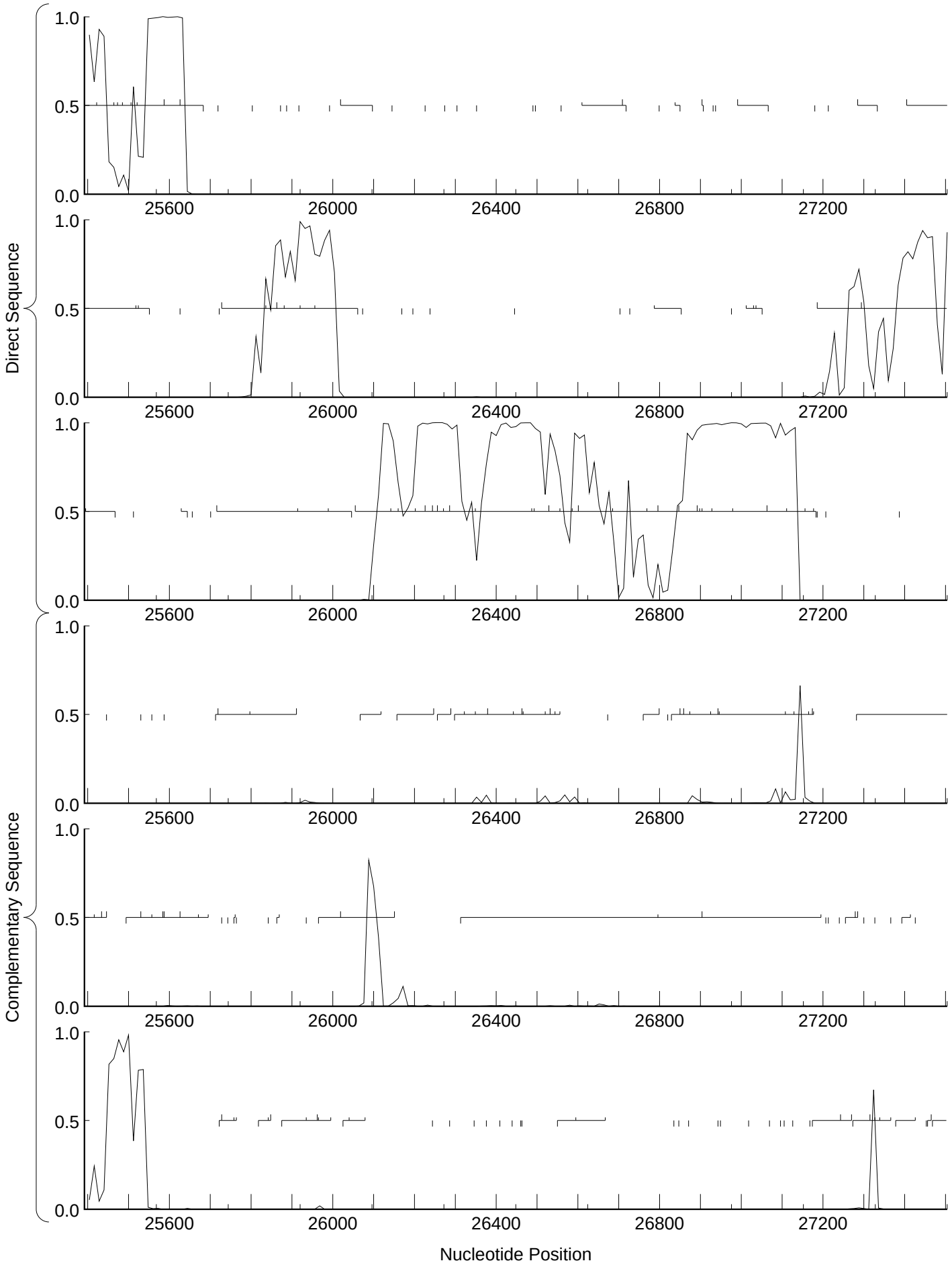
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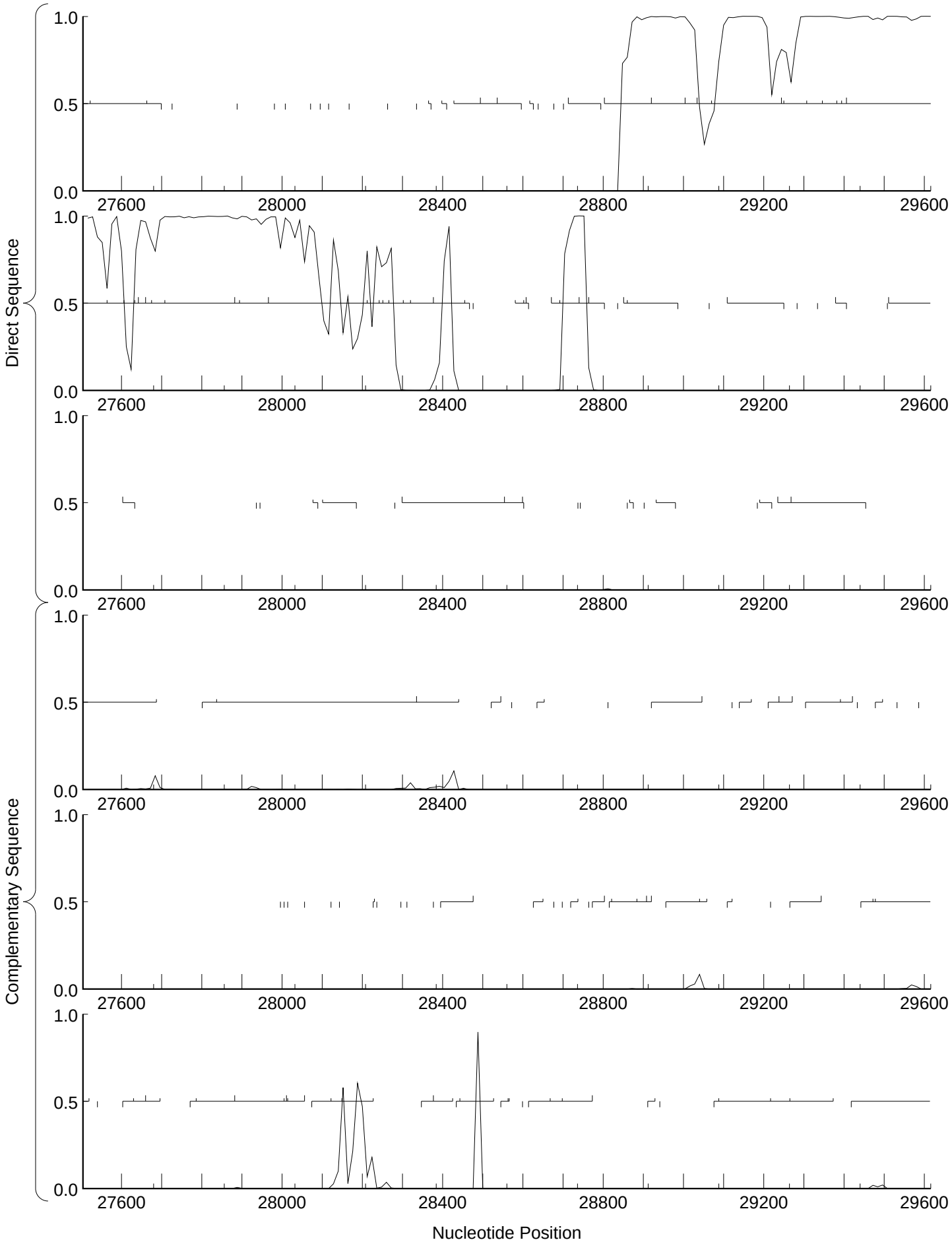
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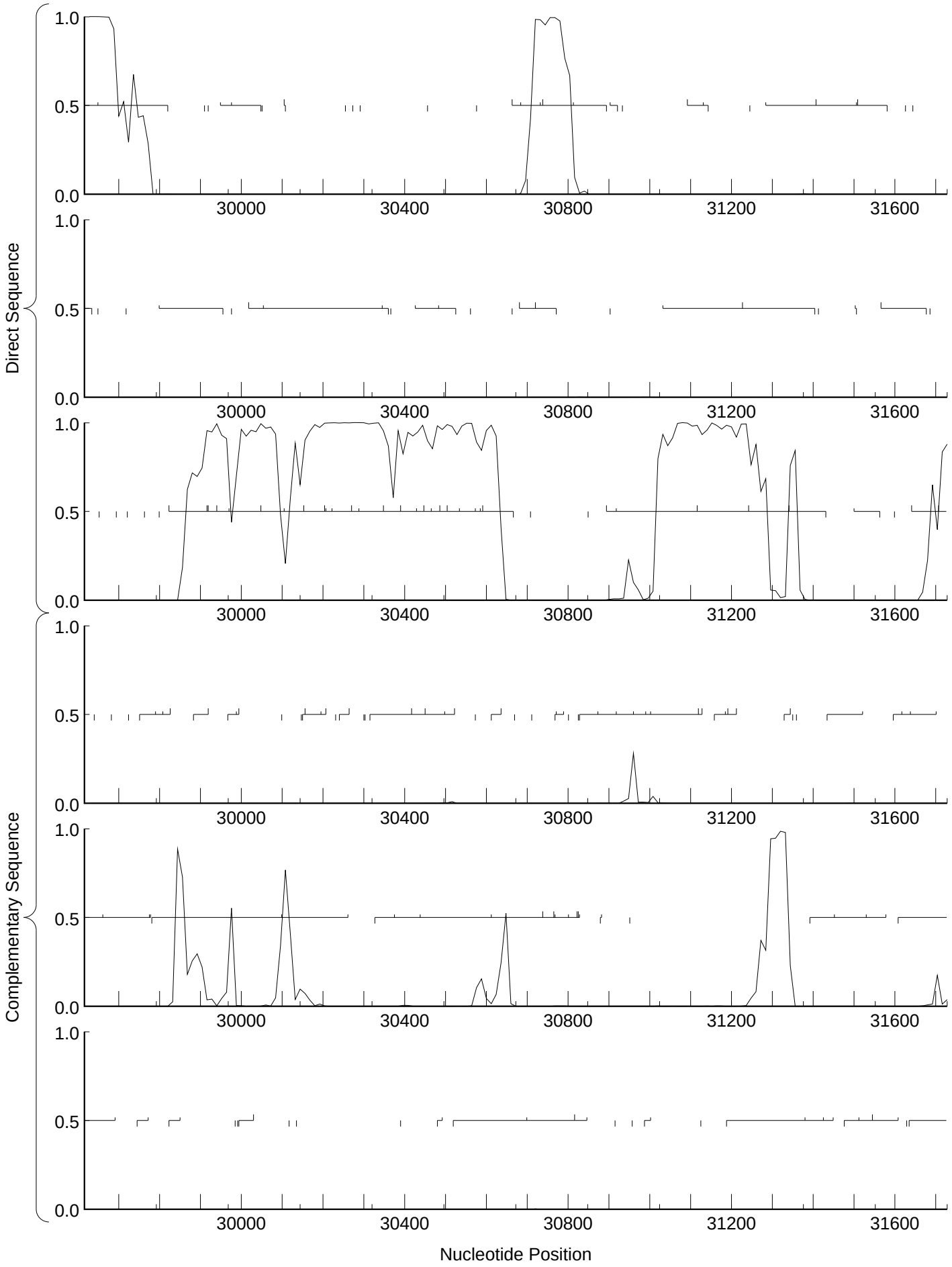
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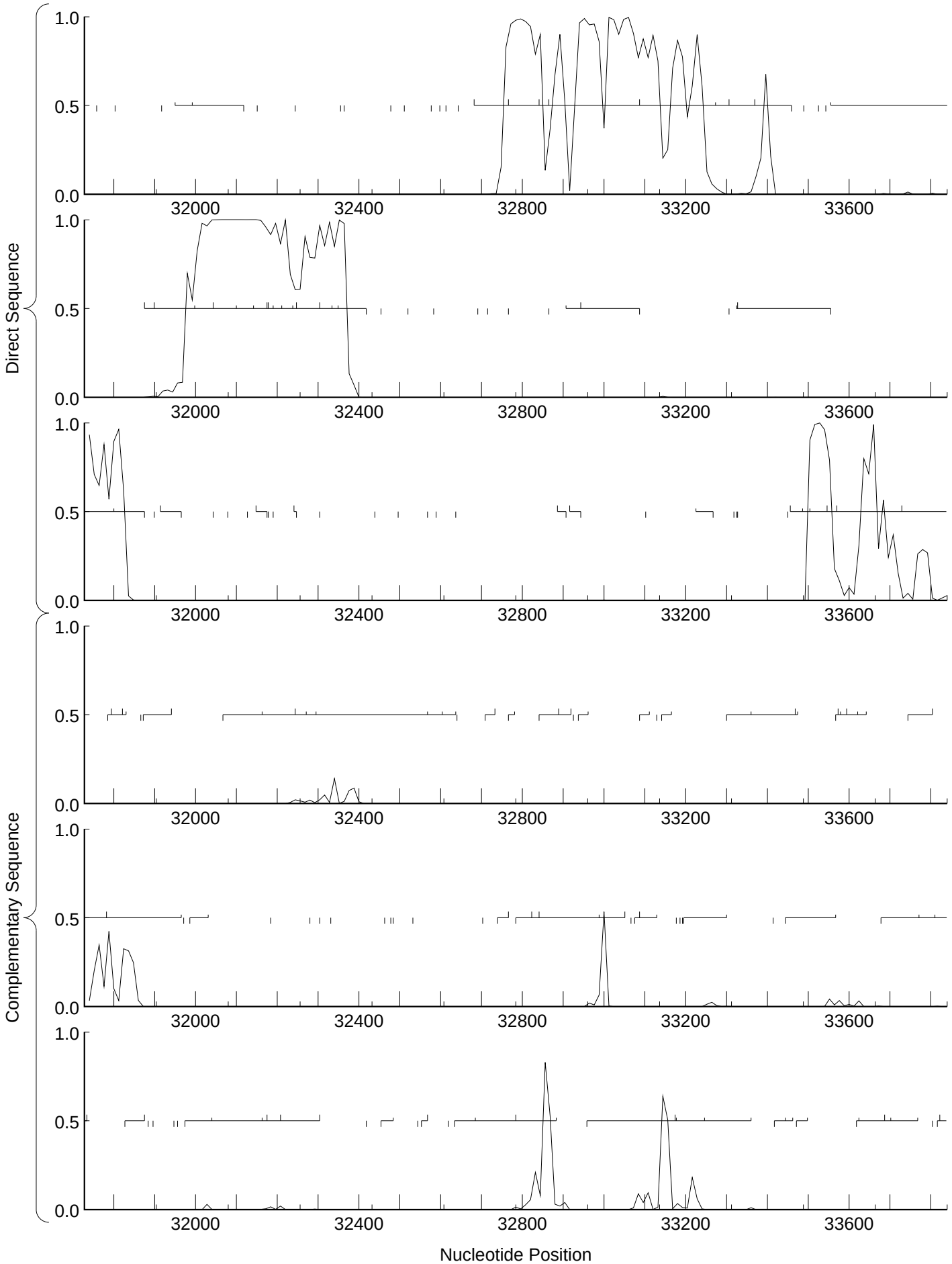
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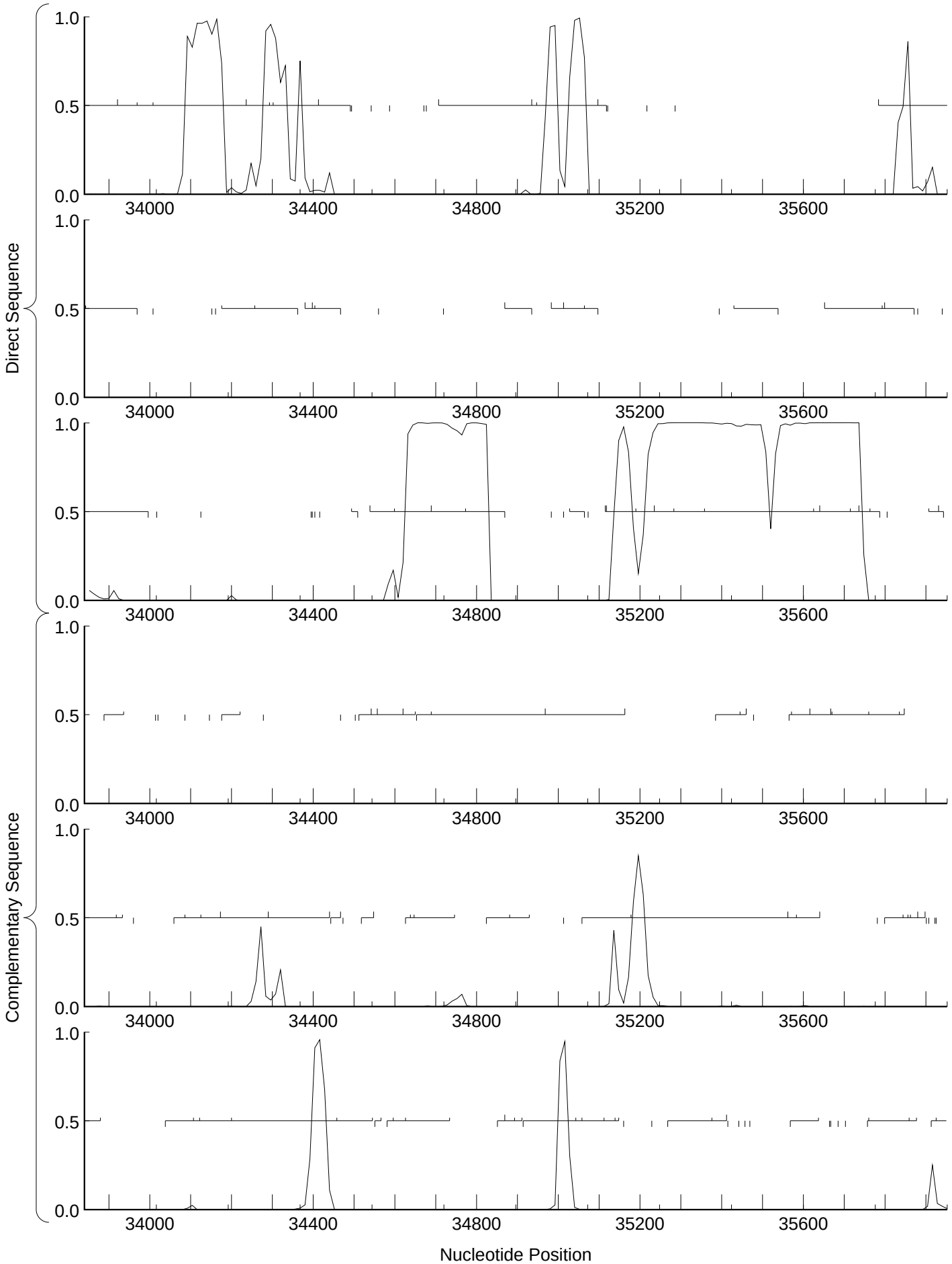


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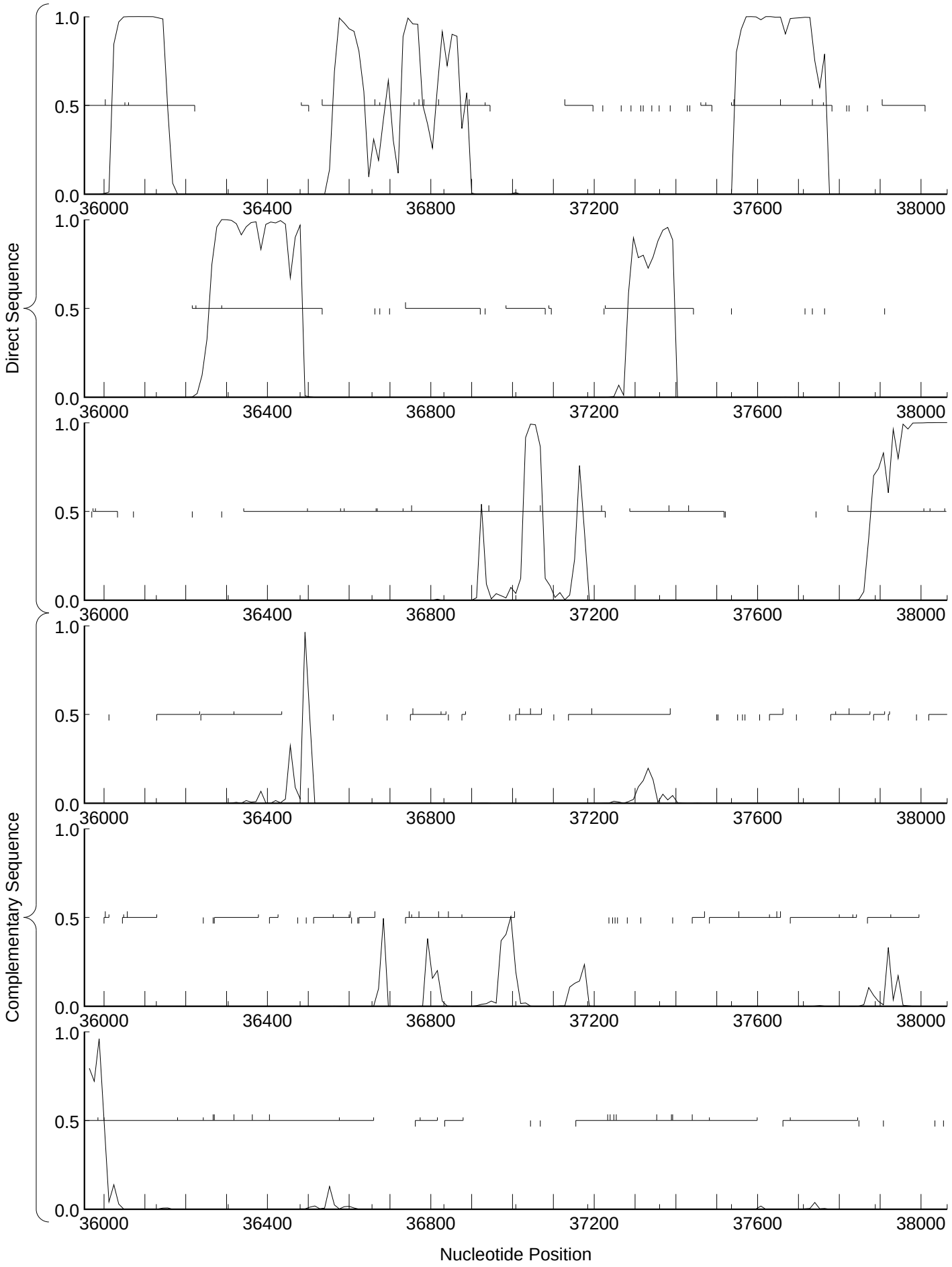


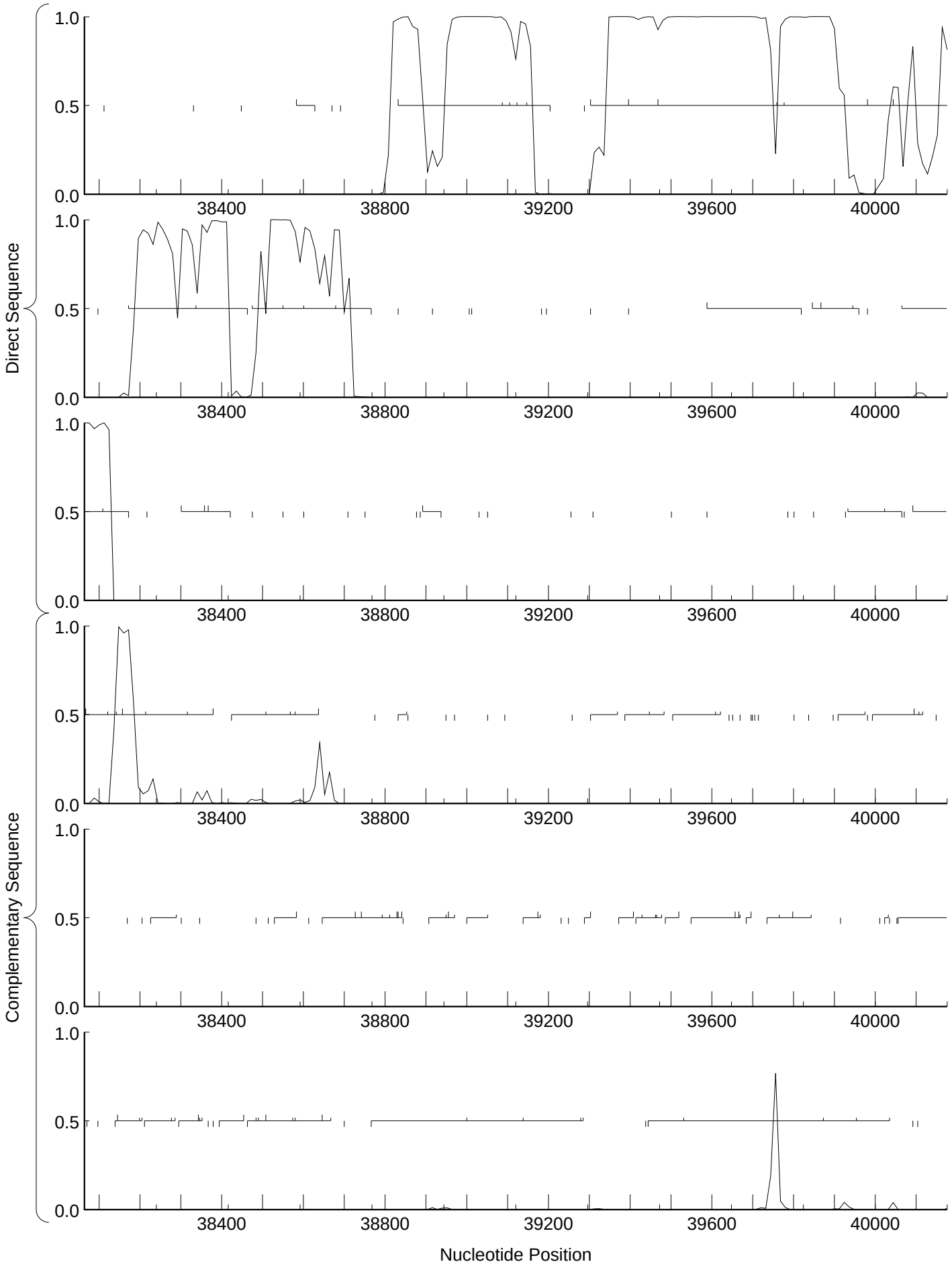
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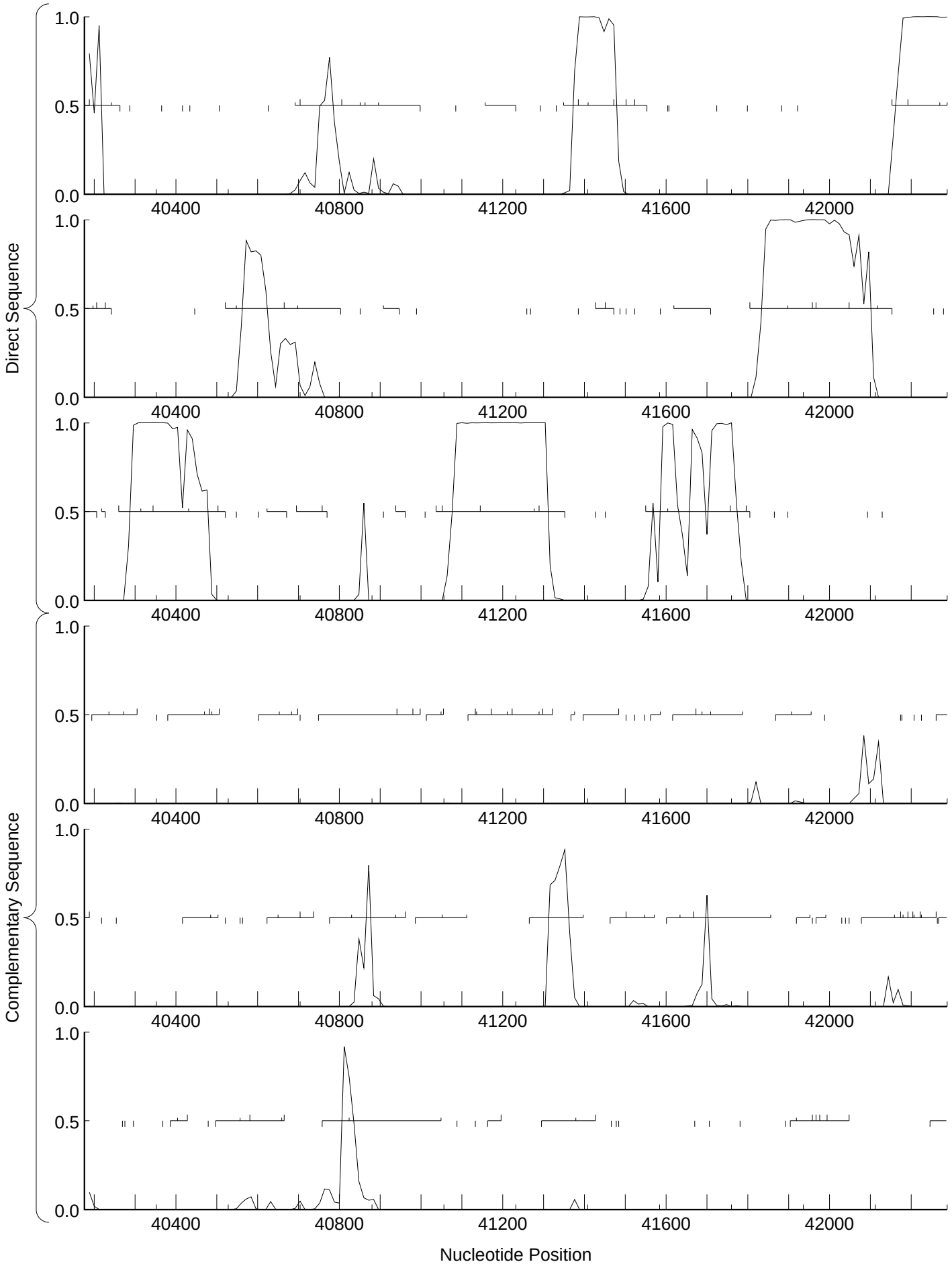

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

throbacter phage Timinator complete sequence, 51138 bp, circularly permuted, Cluster AO2, Order 4, Window 96, Step 12, 19/26

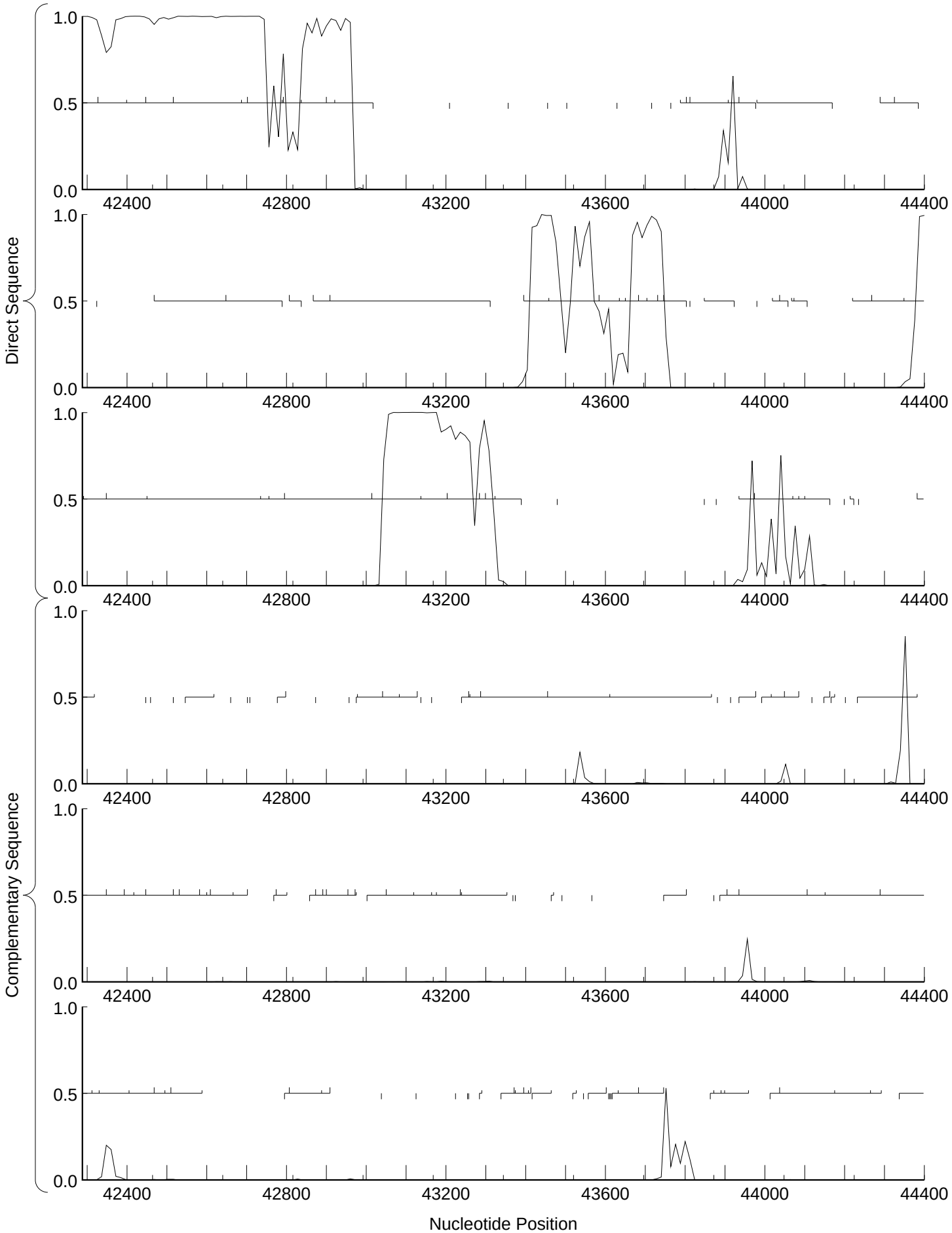



Direct Sequence
Complementary Sequence
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0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

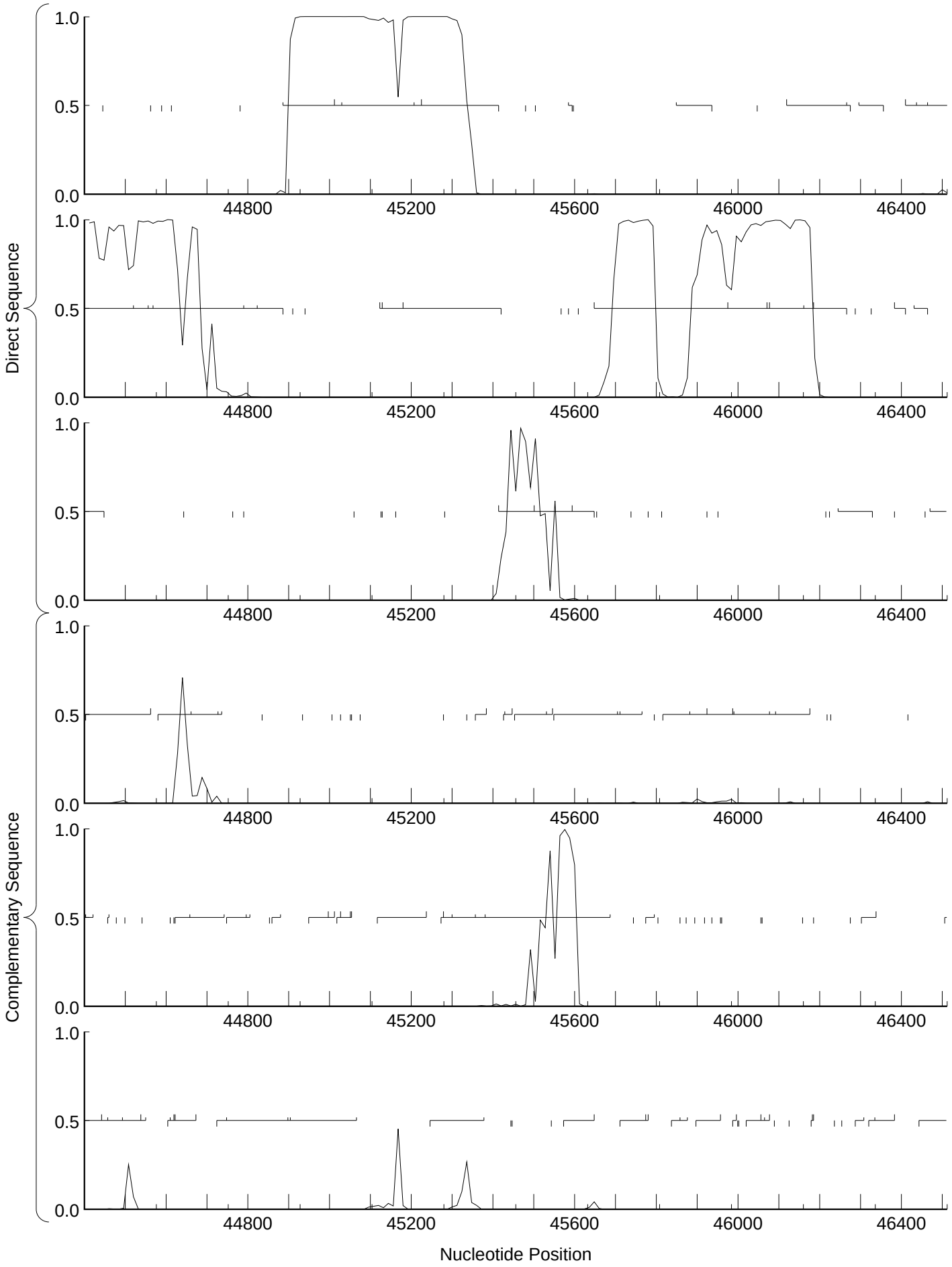
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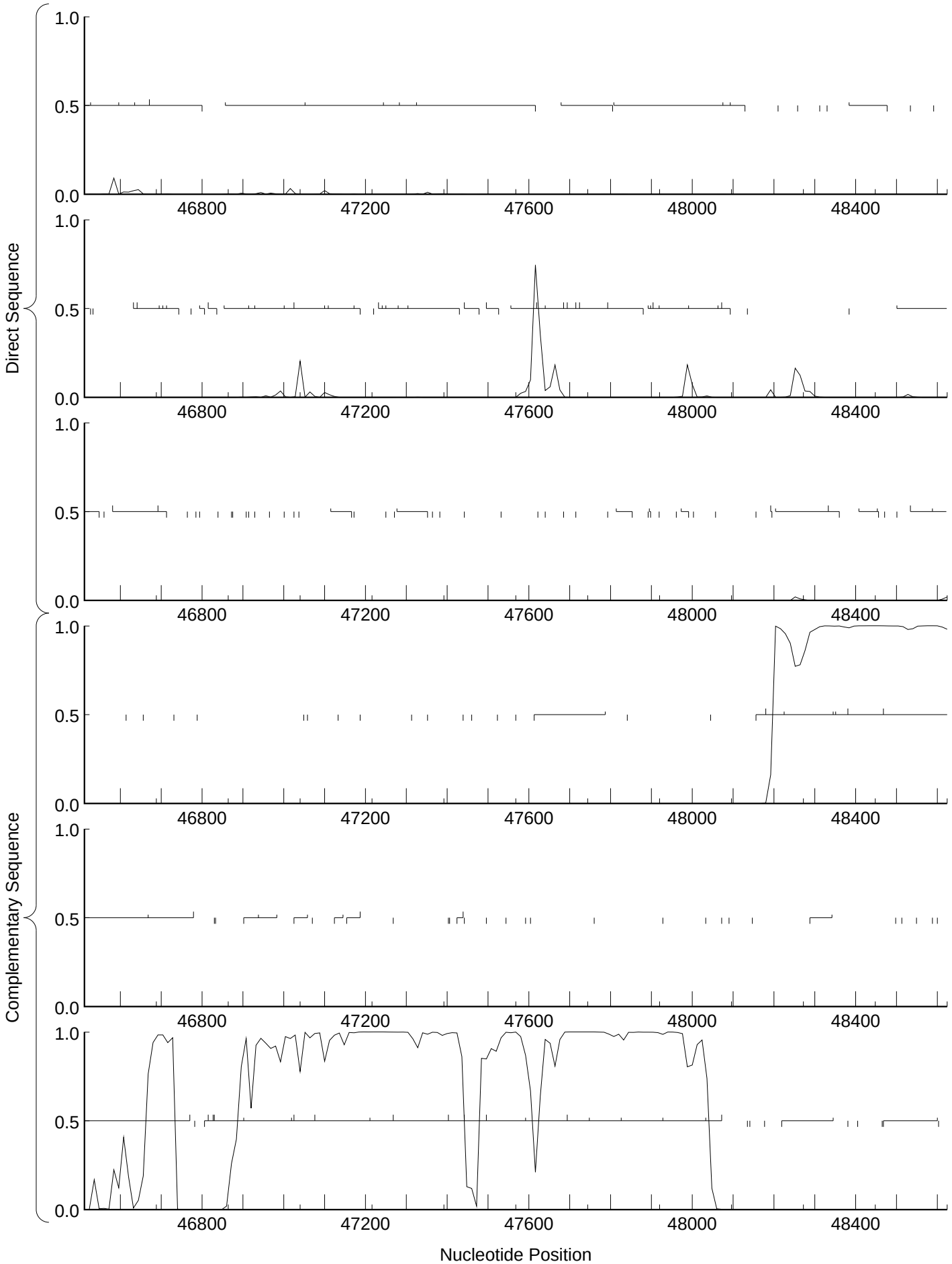


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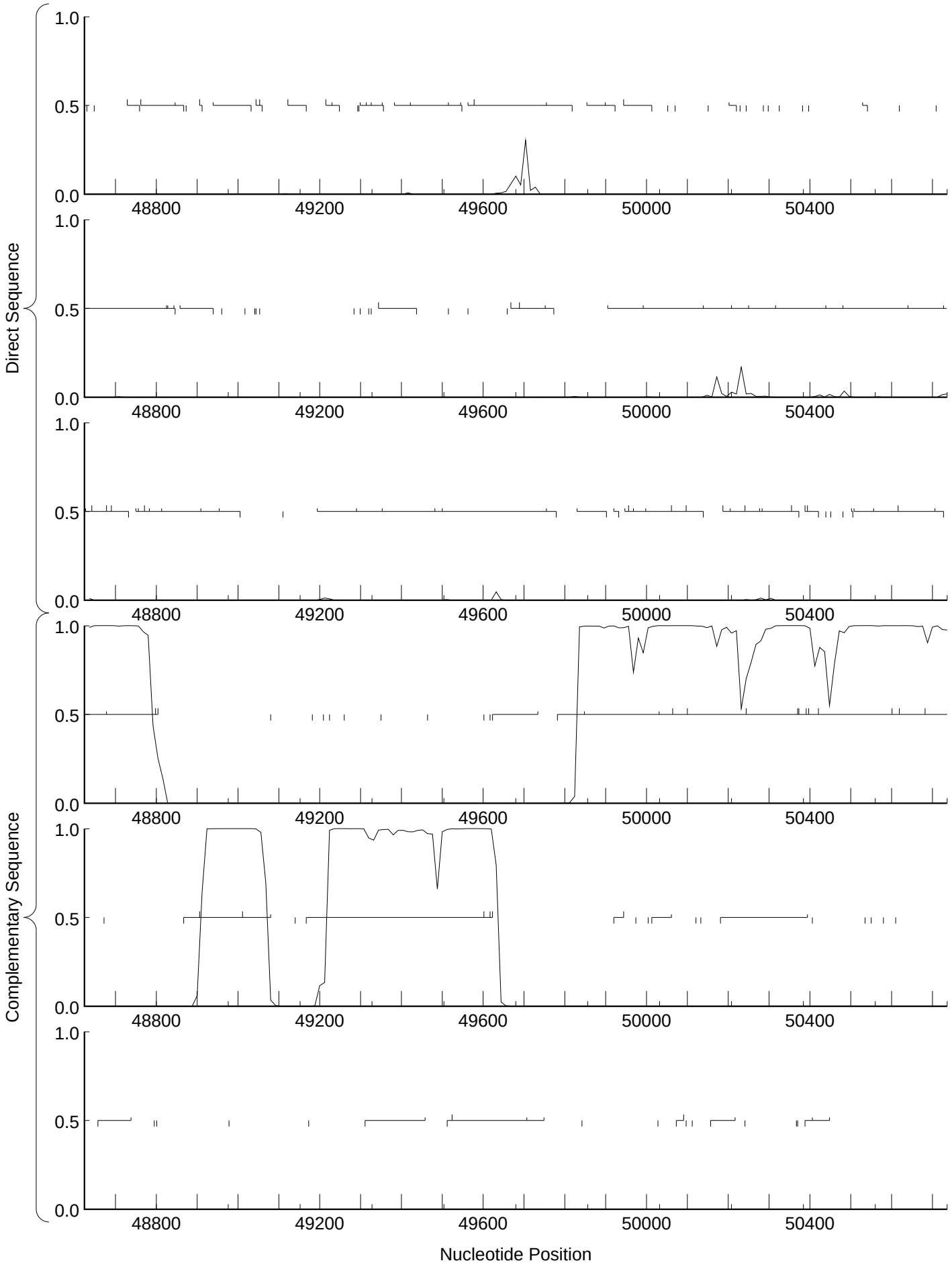


throbacter phage Timinator complete sequence, 51138 bp, circularly permuted, Cluster AO2, Order 4, Window 96, Step 12, 23/26




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
46800
47200
47600
48000
48400
Nucleotide Position

throbacter phage Timinator complete sequence, 51138 bp, circularly permuted, Cluster AO2, Order 4, Window 96, Step 12, 25/26



throbacter phage Timinator complete sequence, 51138 bp, circularly permuted, Cluster AO2, Order 4, Window 96, Step 12, 26/26

