GeneMark

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

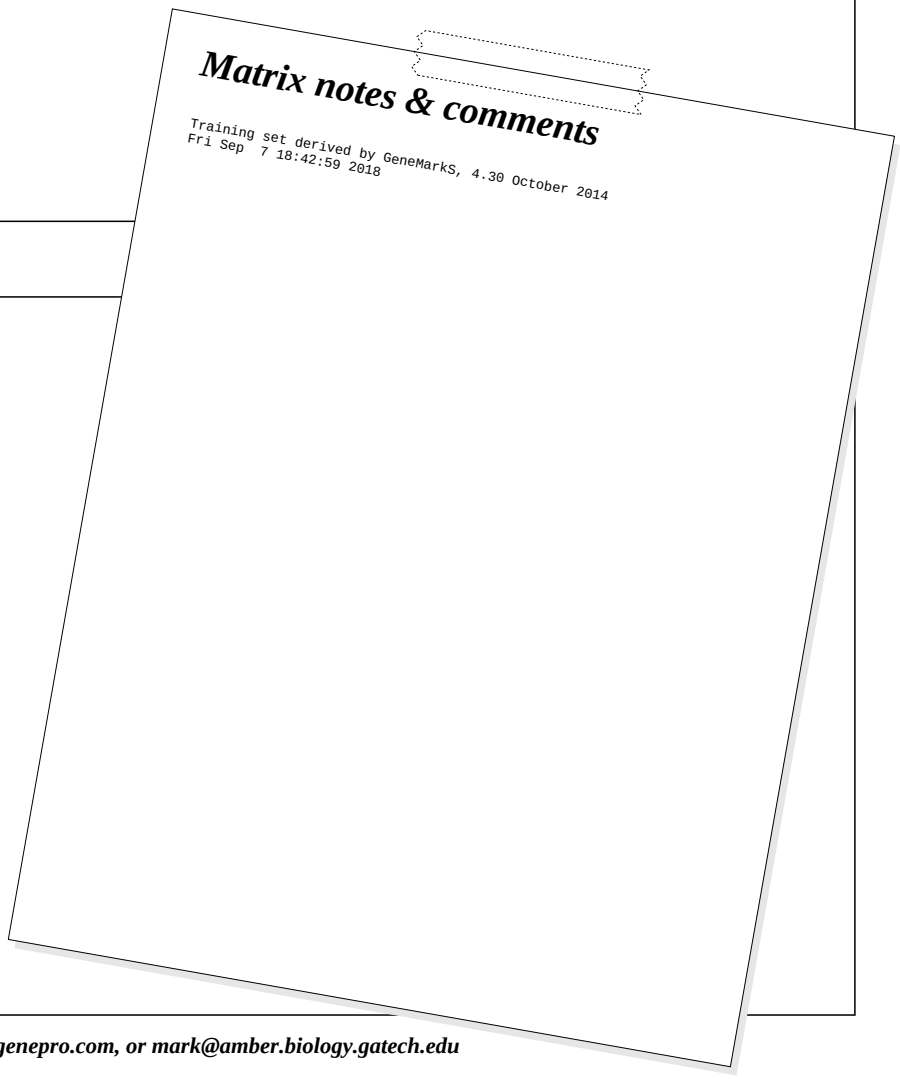
PROGRAM INFORMATION

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Sequence         :  Microbacterium phage LittleFortune complete sequence, 53161 bp, circularly permuted, Cluste
Analysis Date    :  3/29/24 at 5:21:21
Pages            :  27
Sequence Length  :  53161 bp
GC Content       :   68.96%

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

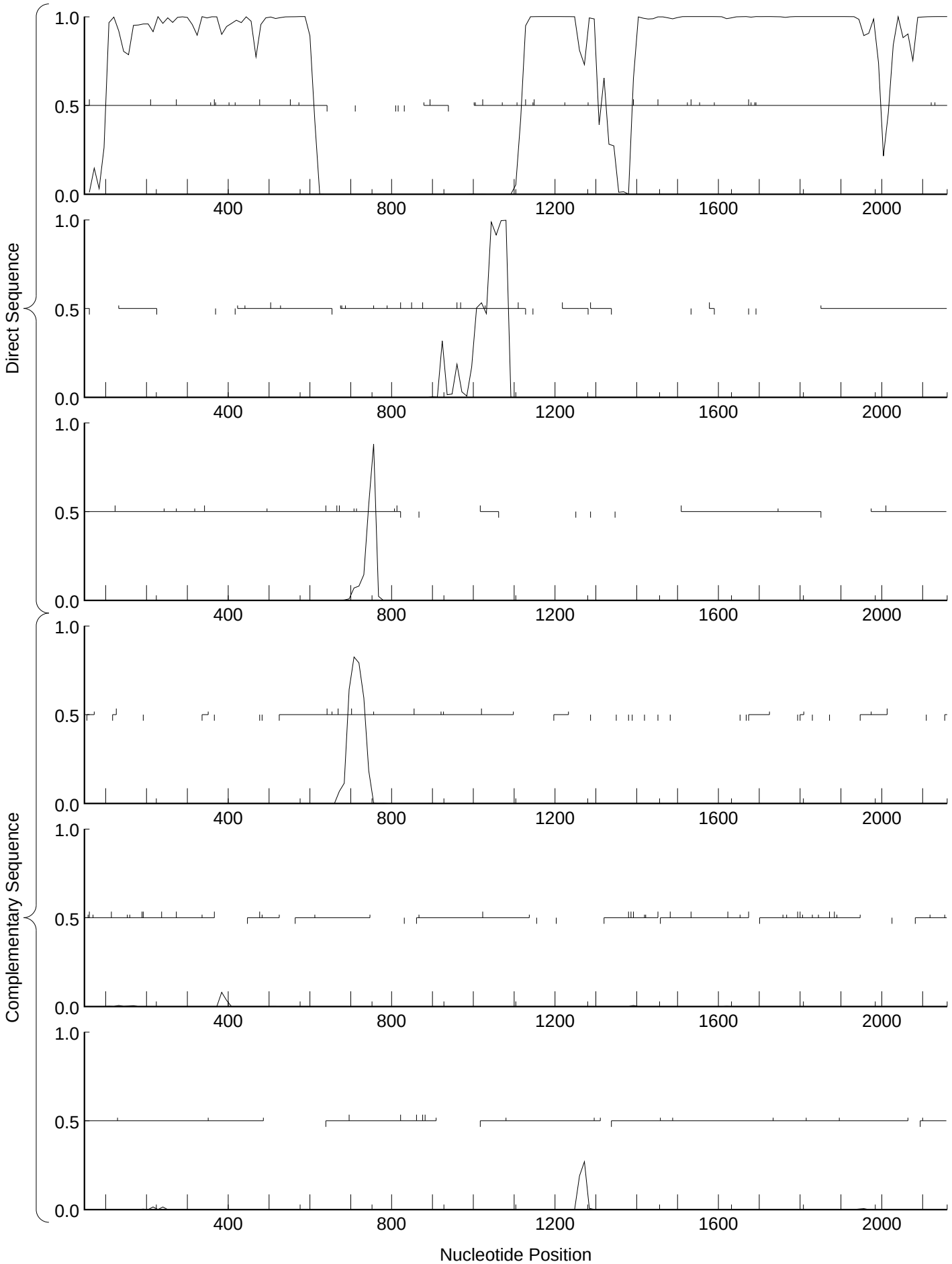
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Matrix     :  -
Author     :  -

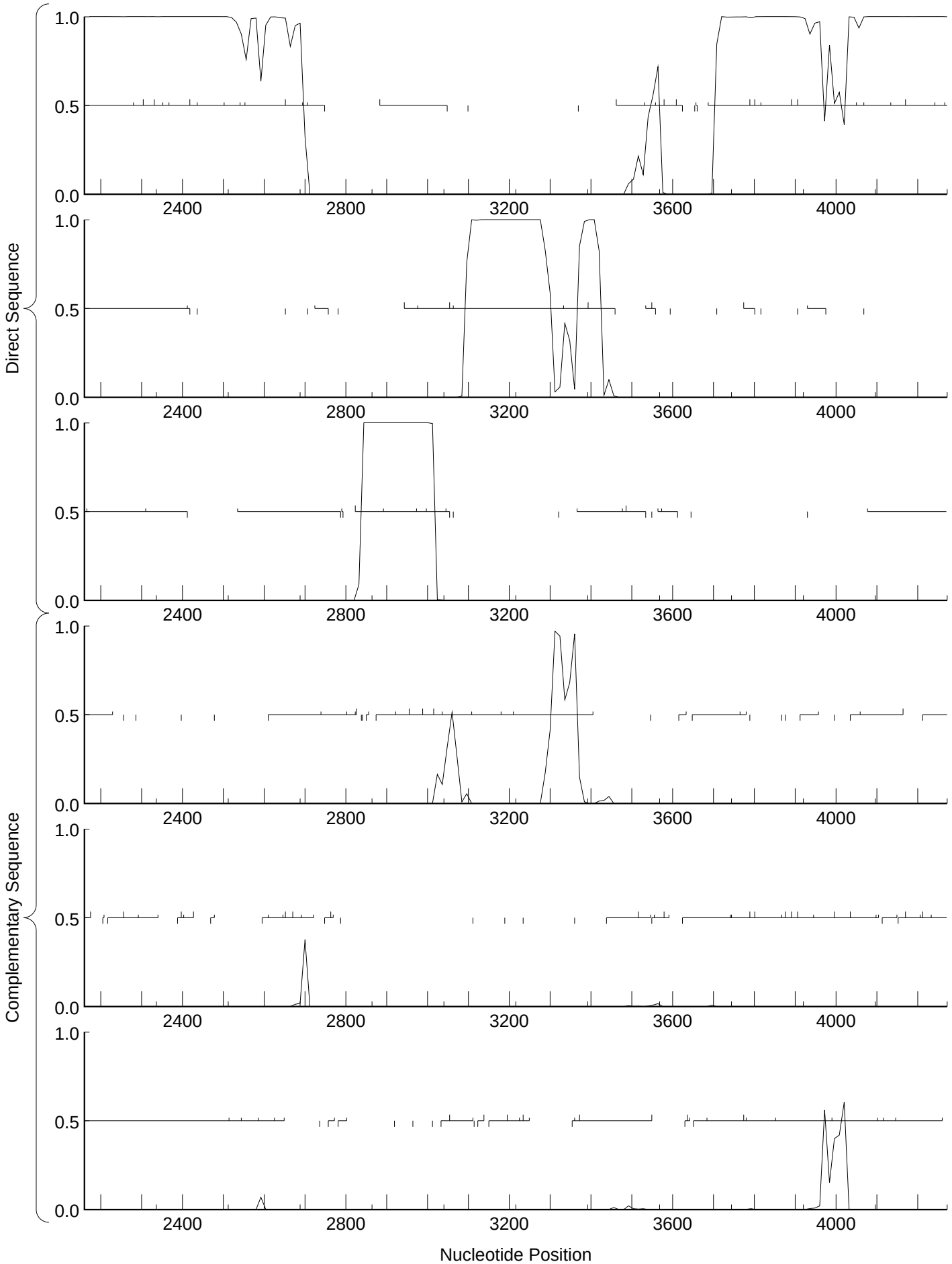
Order      :  4
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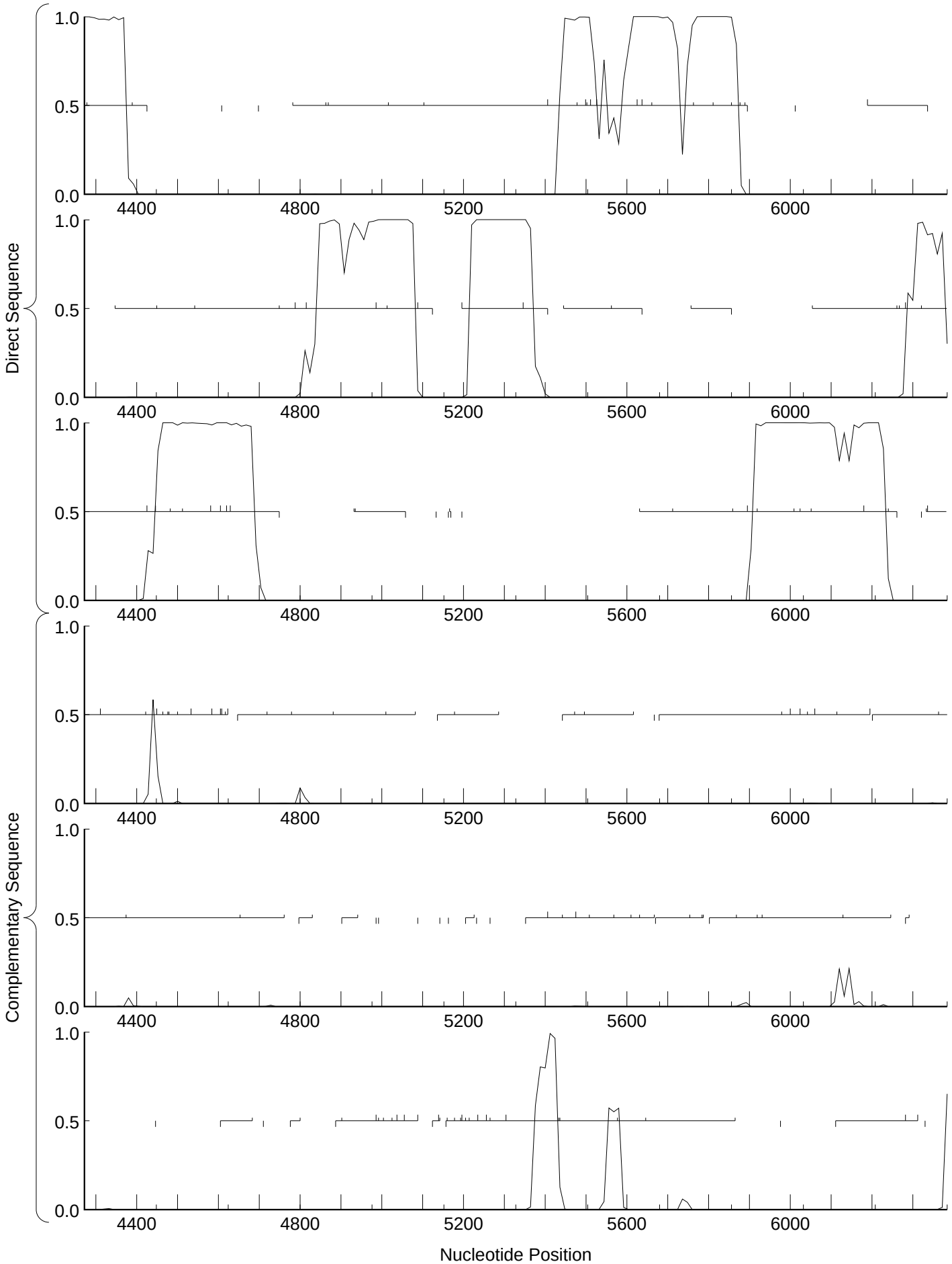

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

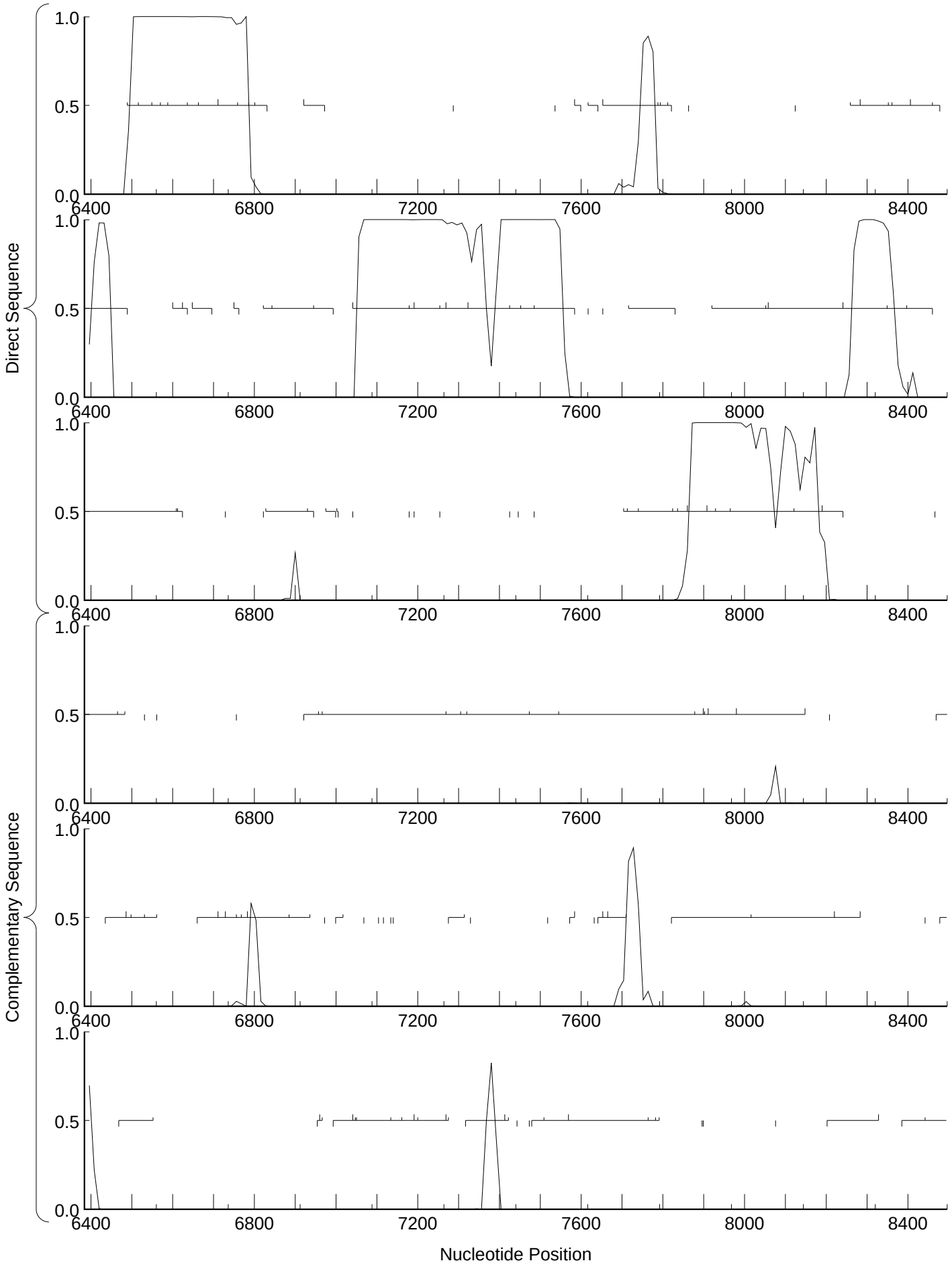
bacterium phage LittleFortune complete sequence, 53161 bp, circularly permuted, Cluster EC, Order 4, Window 96, Step 12, 2/27

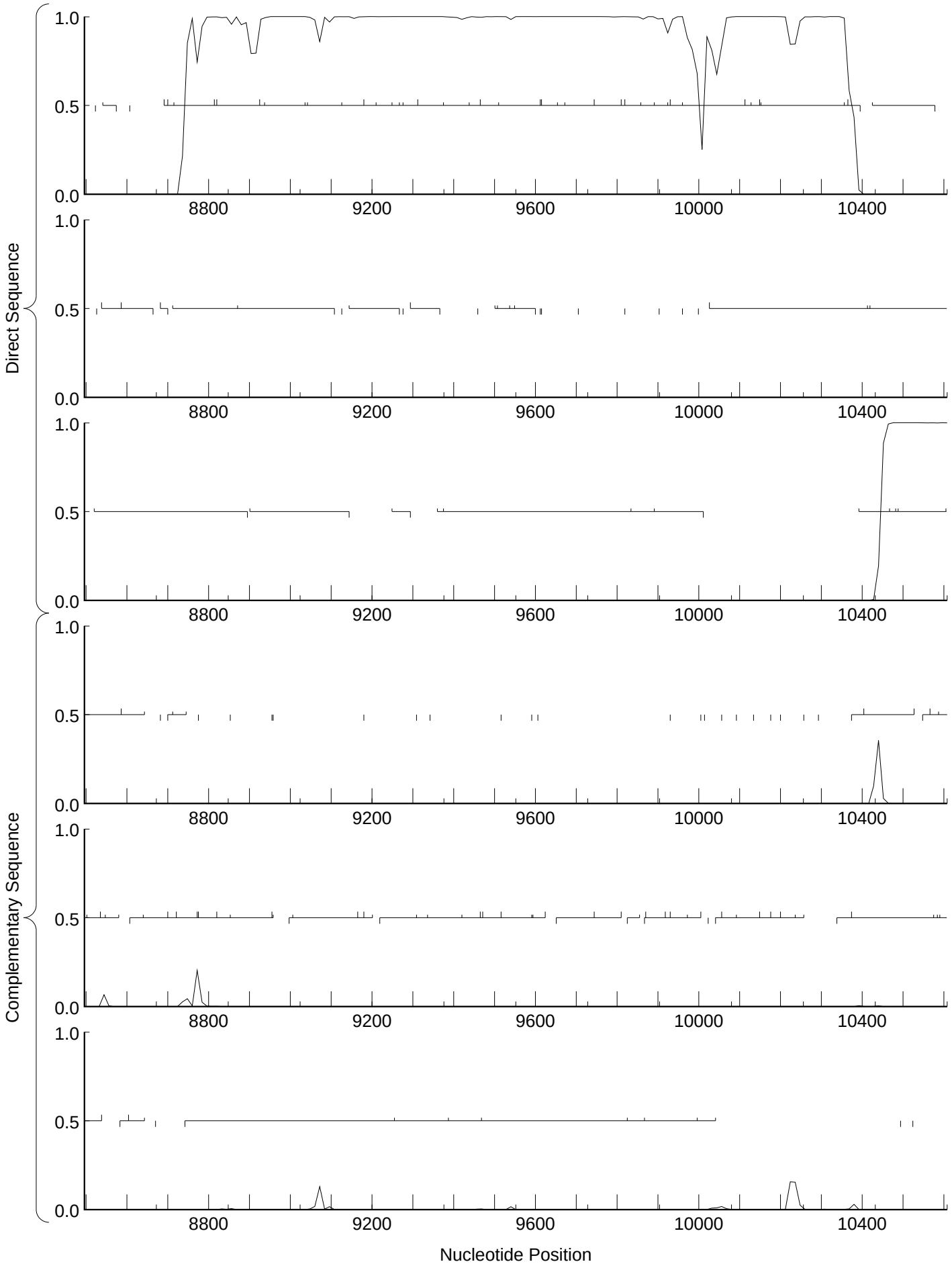



Direct Sequence
Complementary Sequence
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0.5
0.0
2400
2800
3200
3600
4000
Nucleotide Position

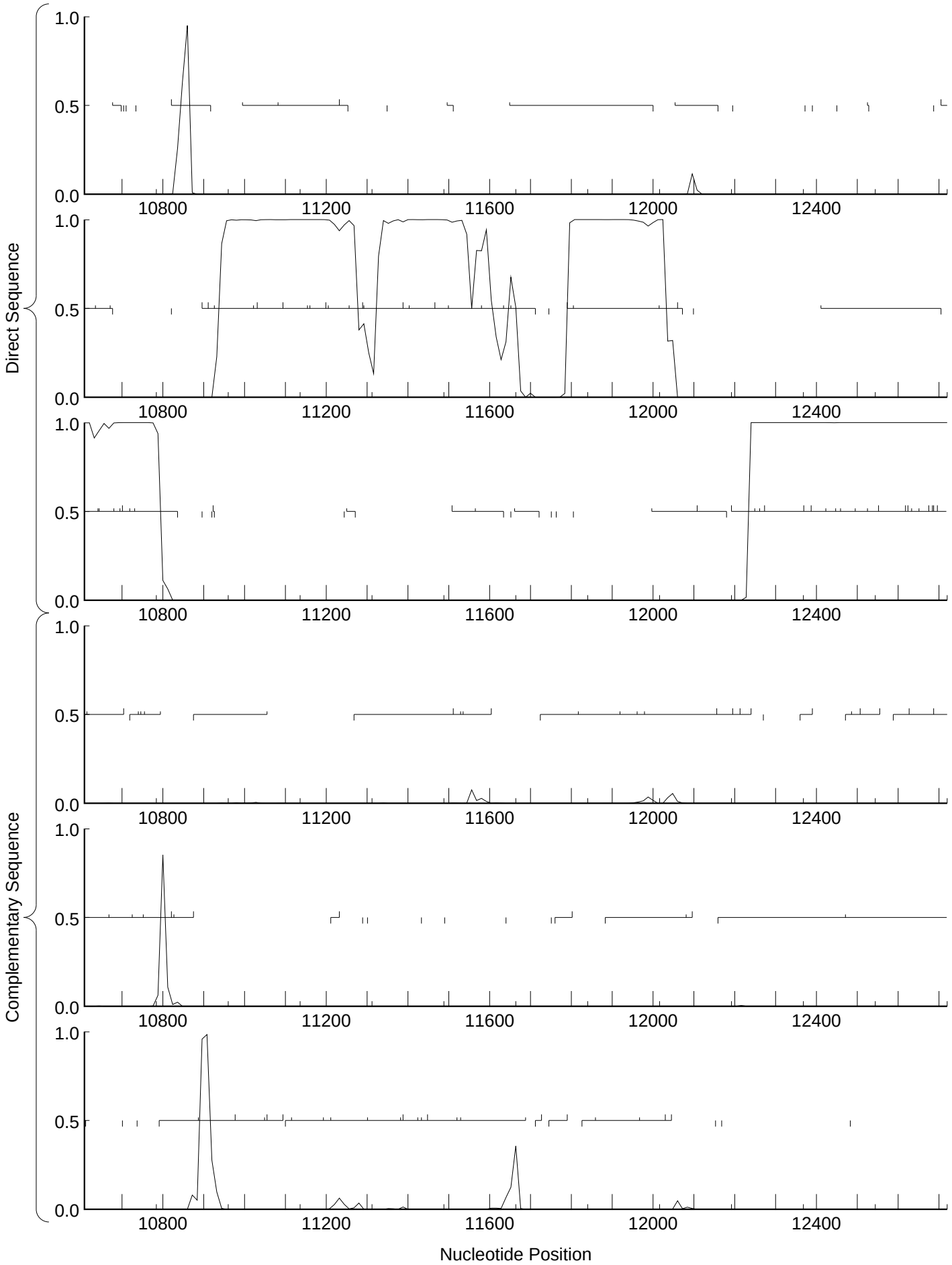

Direct Sequence
Complementary Sequence
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0.0
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5200
5600
6000
Nucleotide Position

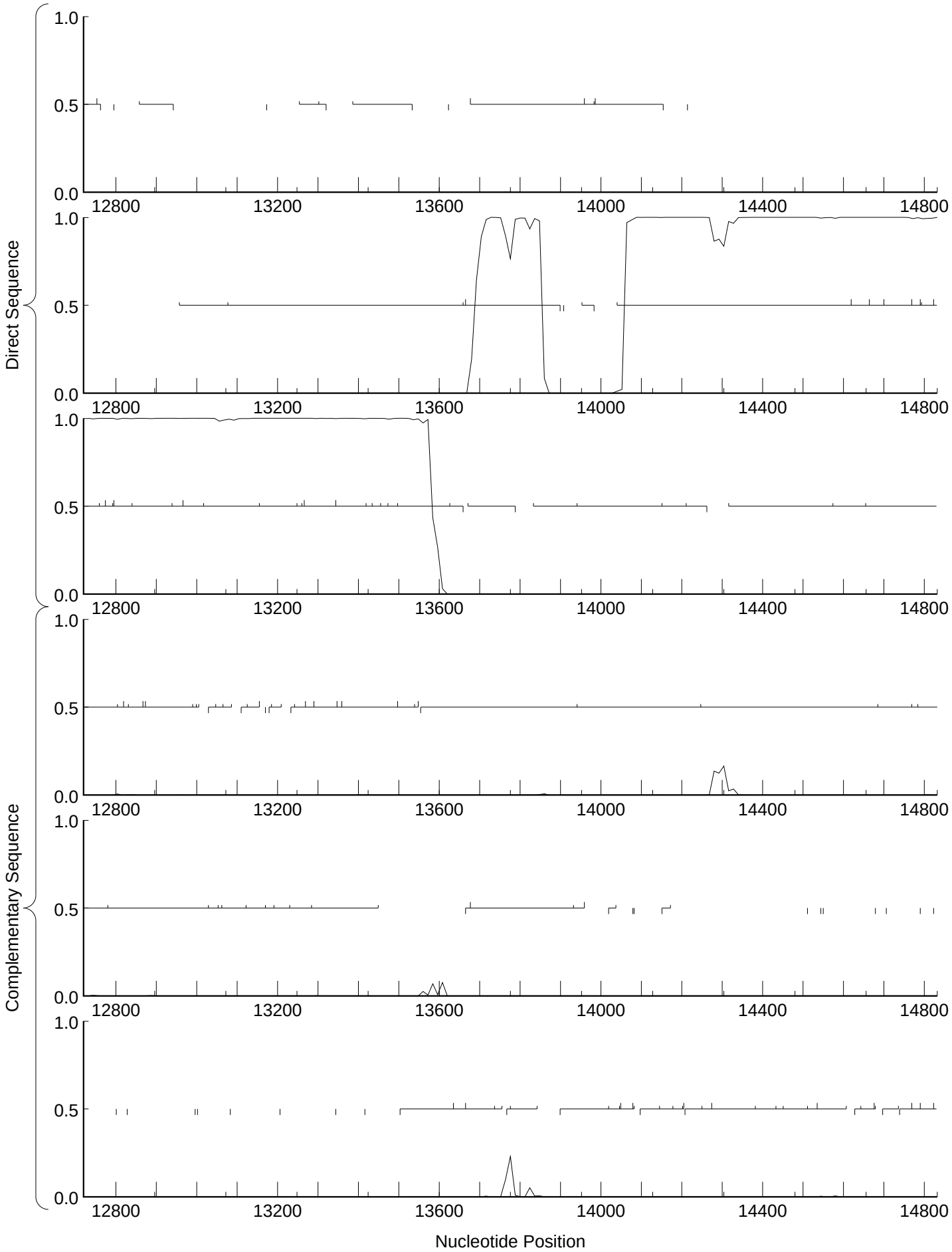
bacterium phage LittleFortune complete sequence, 53161 bp, circularly permuted, Cluster EC, Order 4, Window 96, Step 12, 5/27



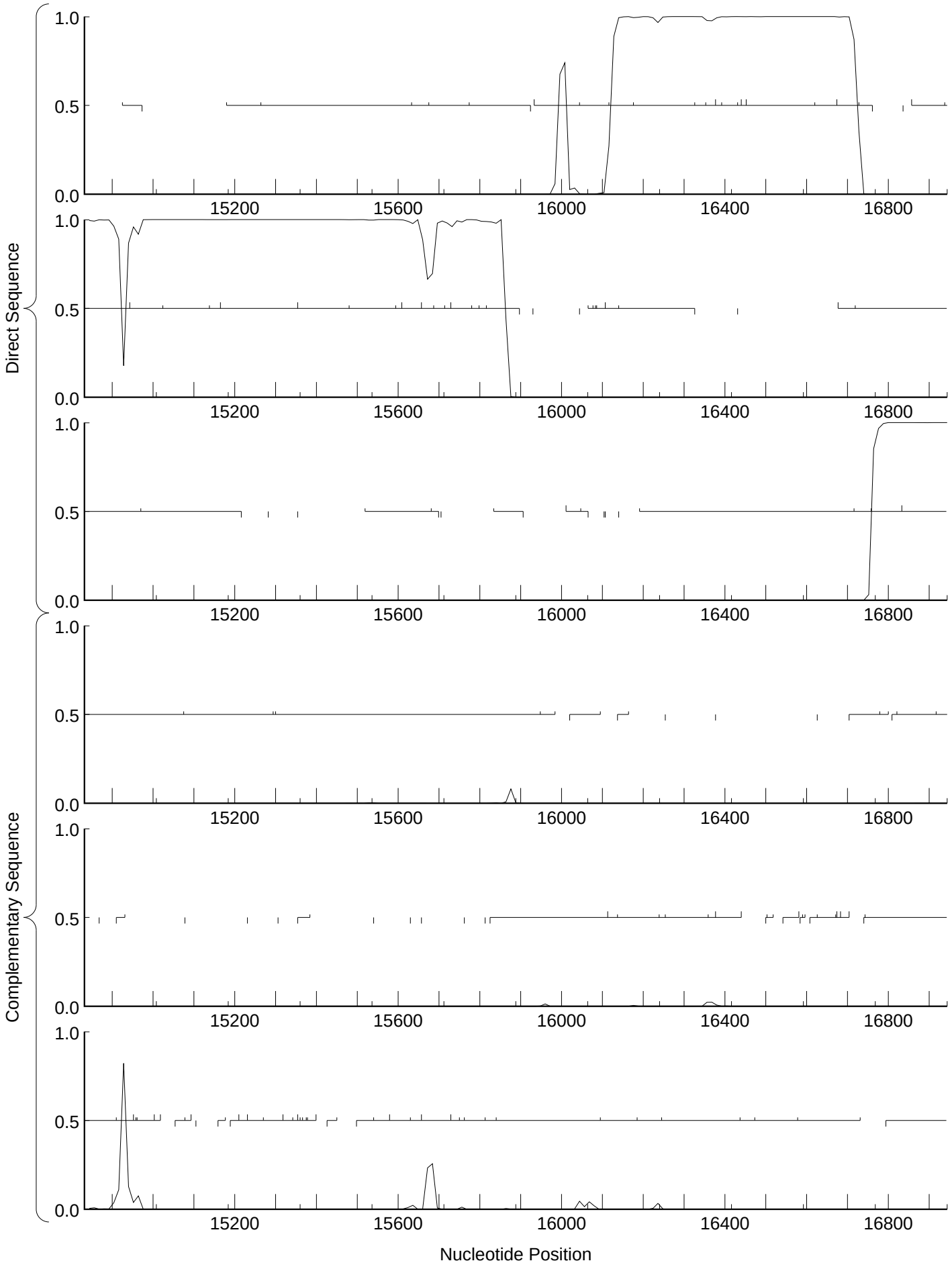

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
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9200
9600
10000
10400
Nucleotide Position

bacterium phage LittleFortune complete sequence, 53161 bp, circularly permuted, Cluster EC, Order 4, Window 96, Step 12, 7/27

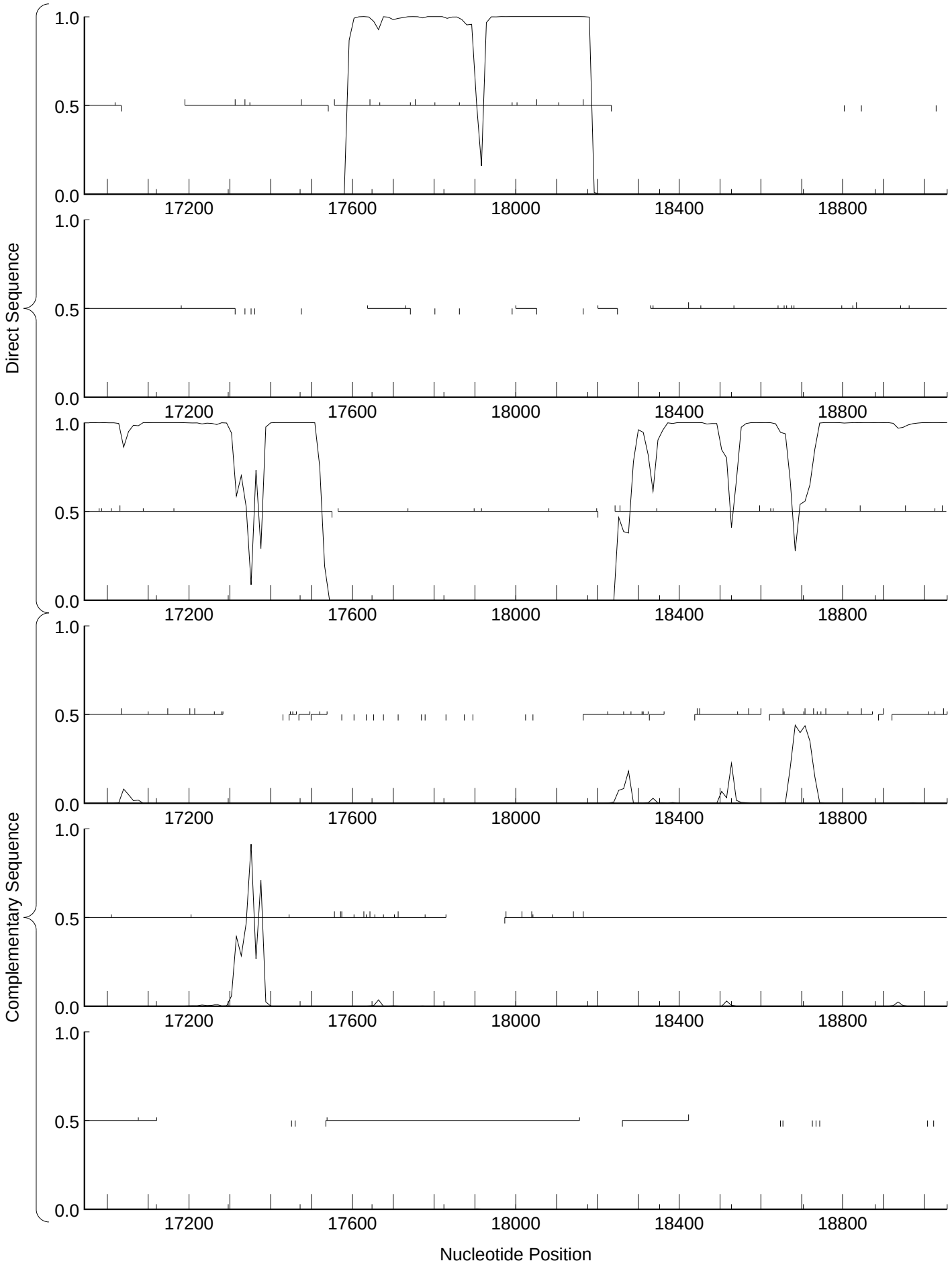



Direct Sequence
Complementary Sequence
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0.5
0.0
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13200
13600
14000
14400
14800
Nucleotide Position

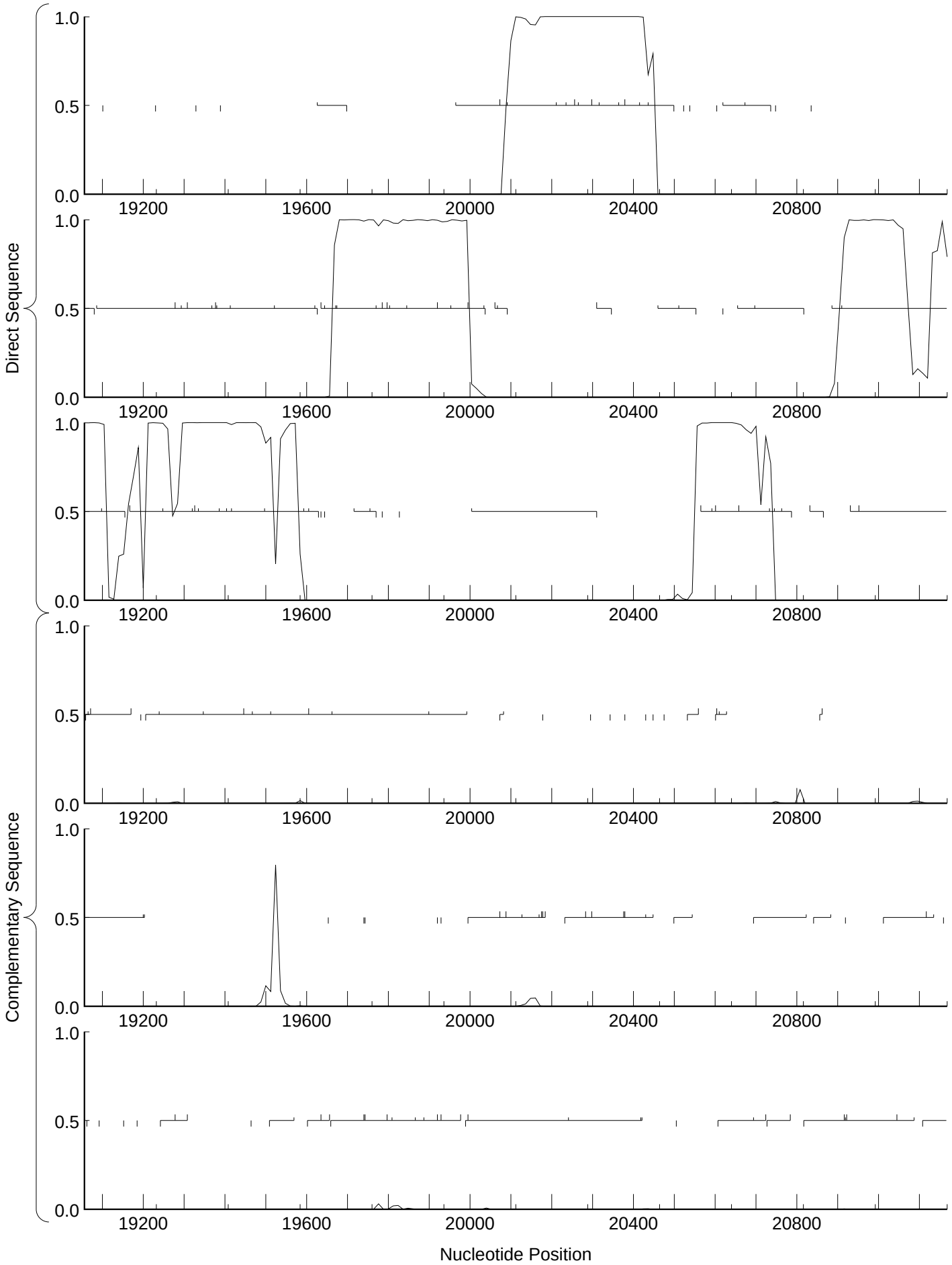
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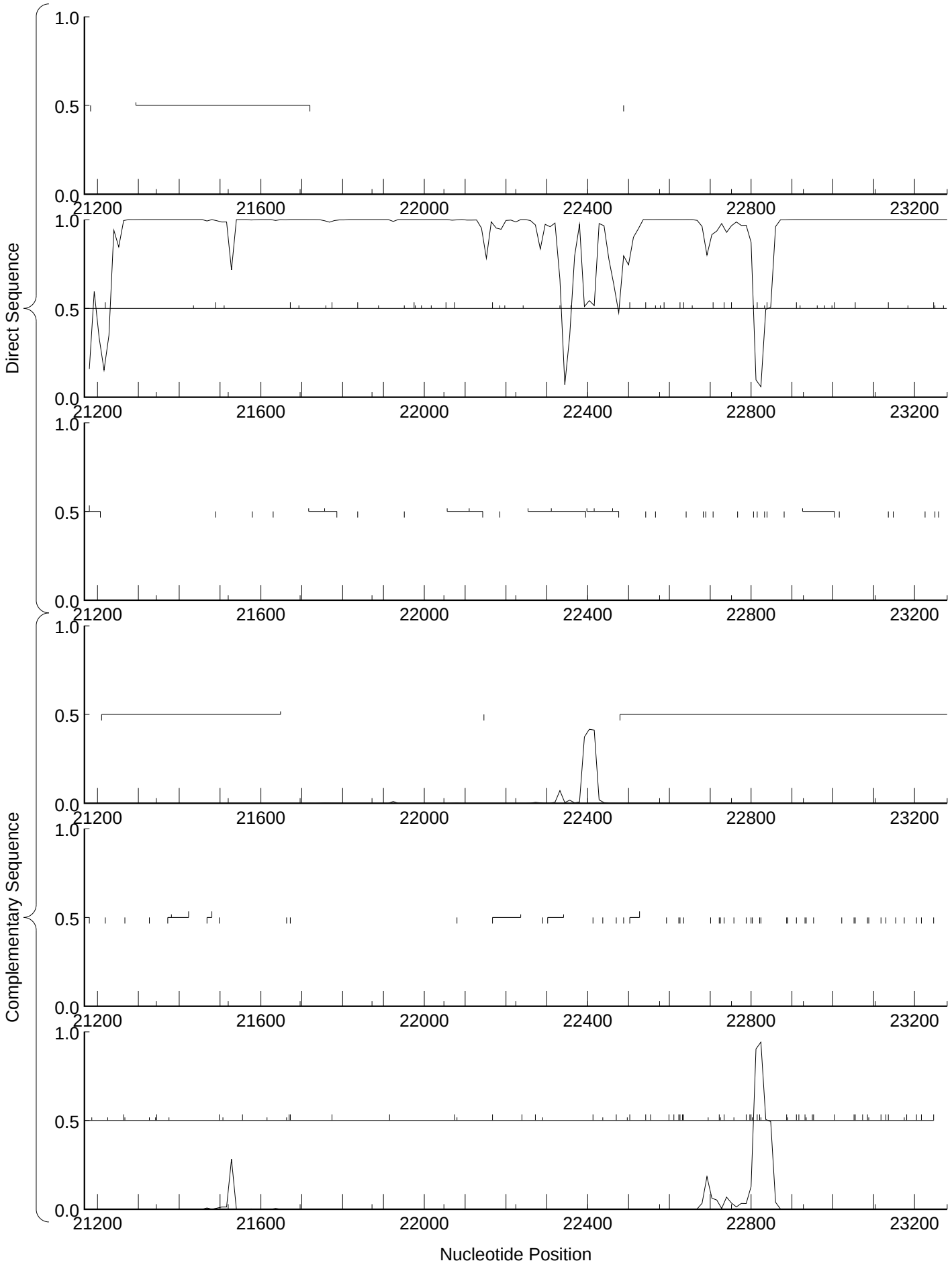
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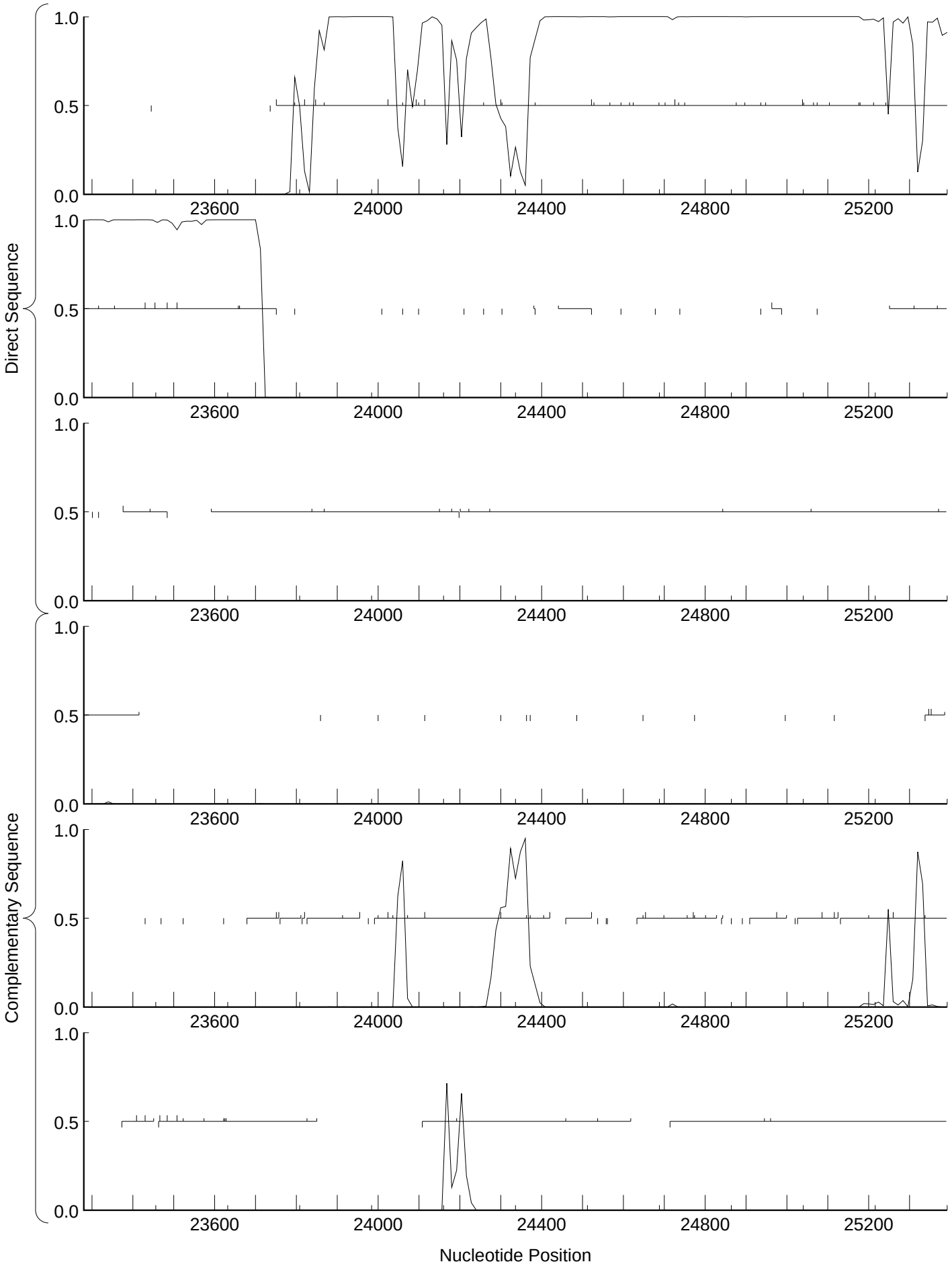
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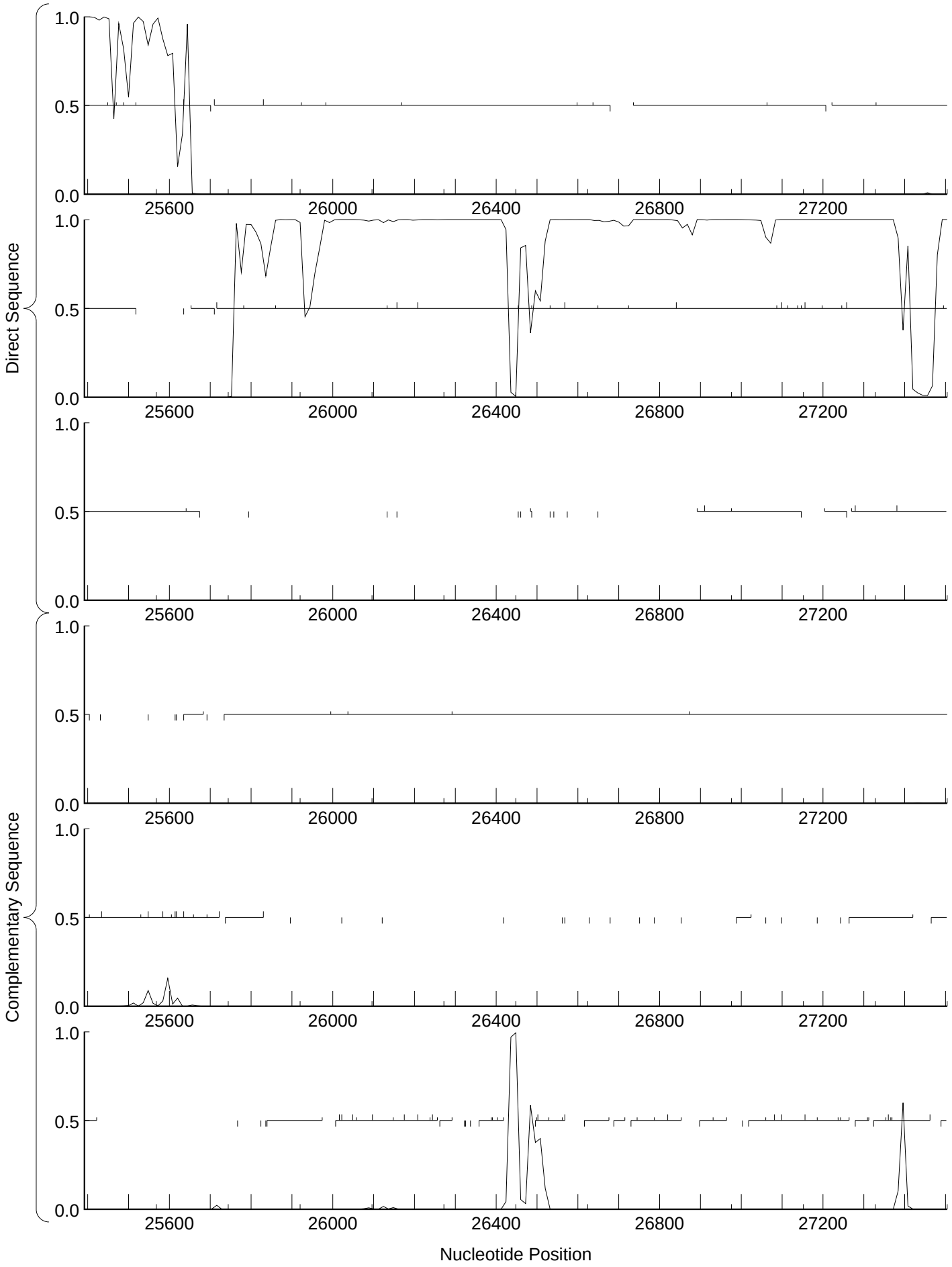
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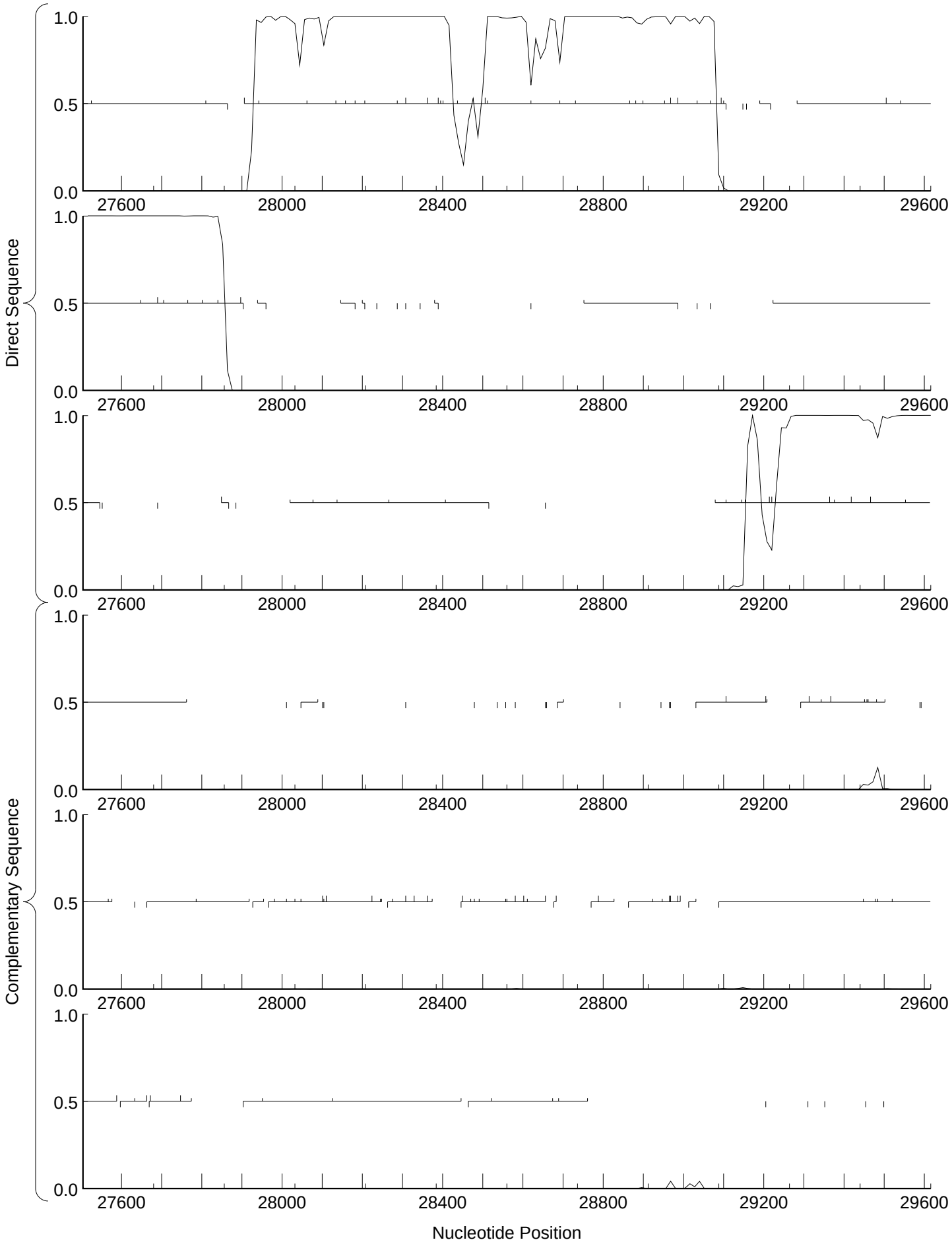
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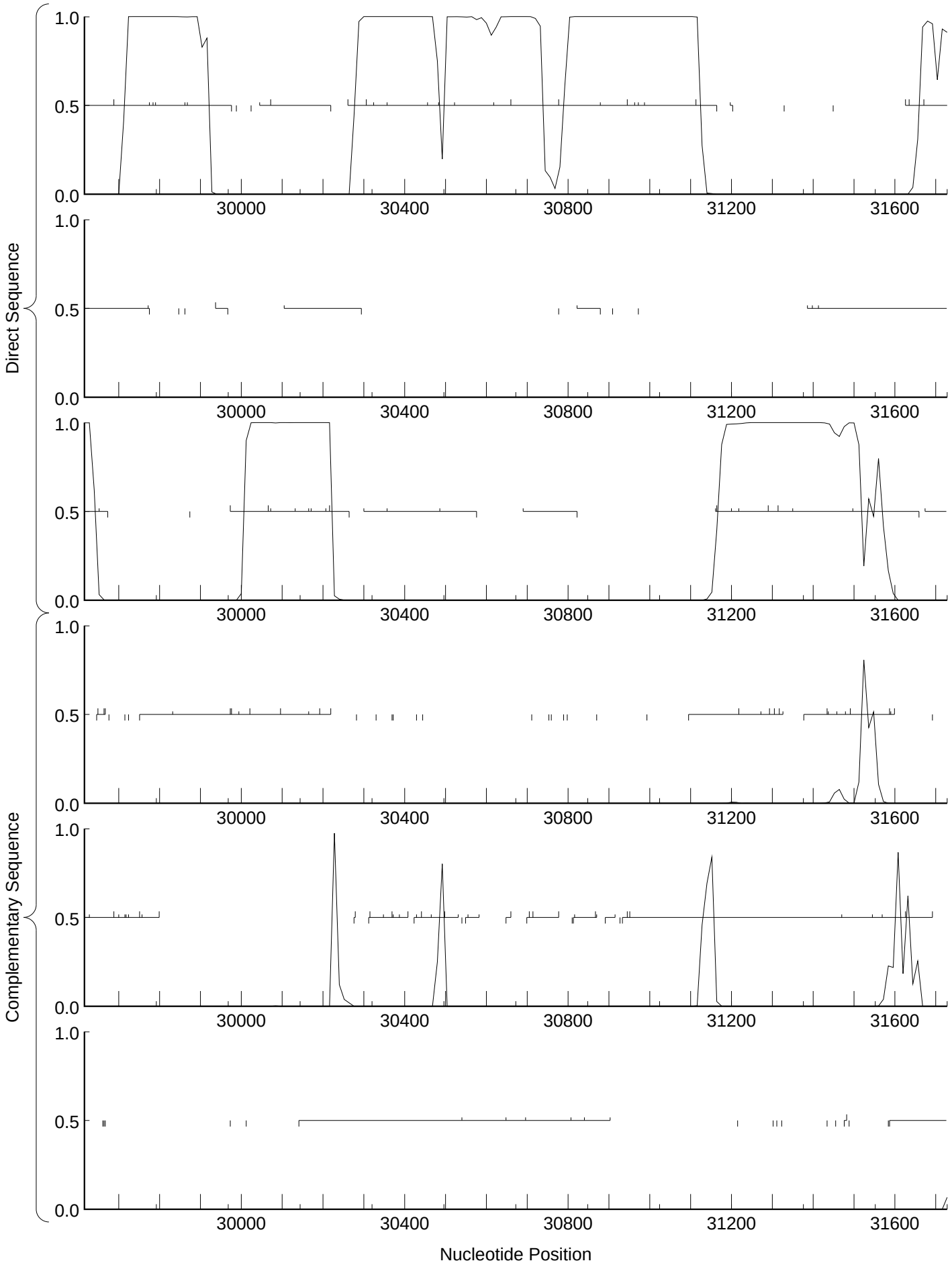
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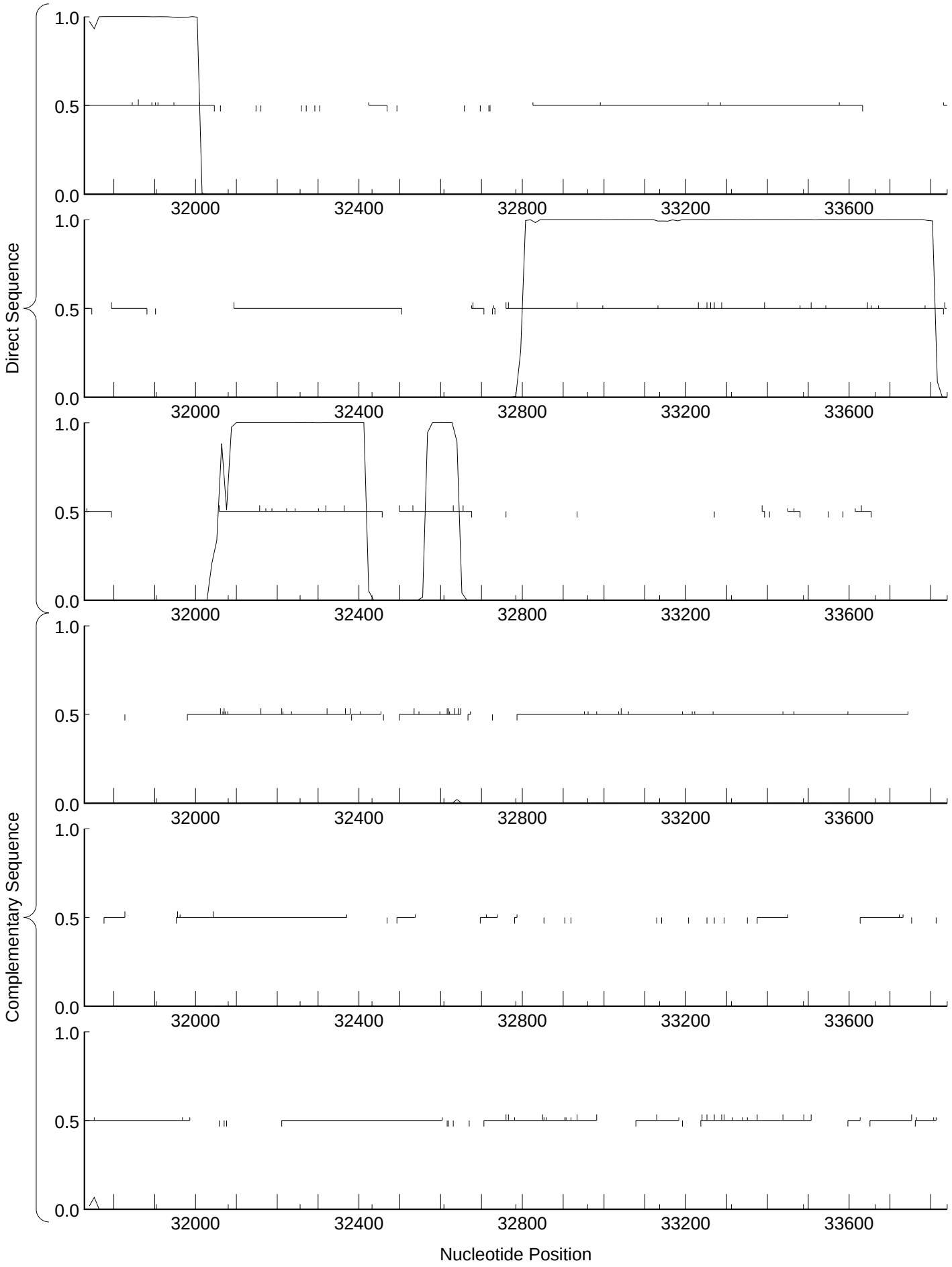
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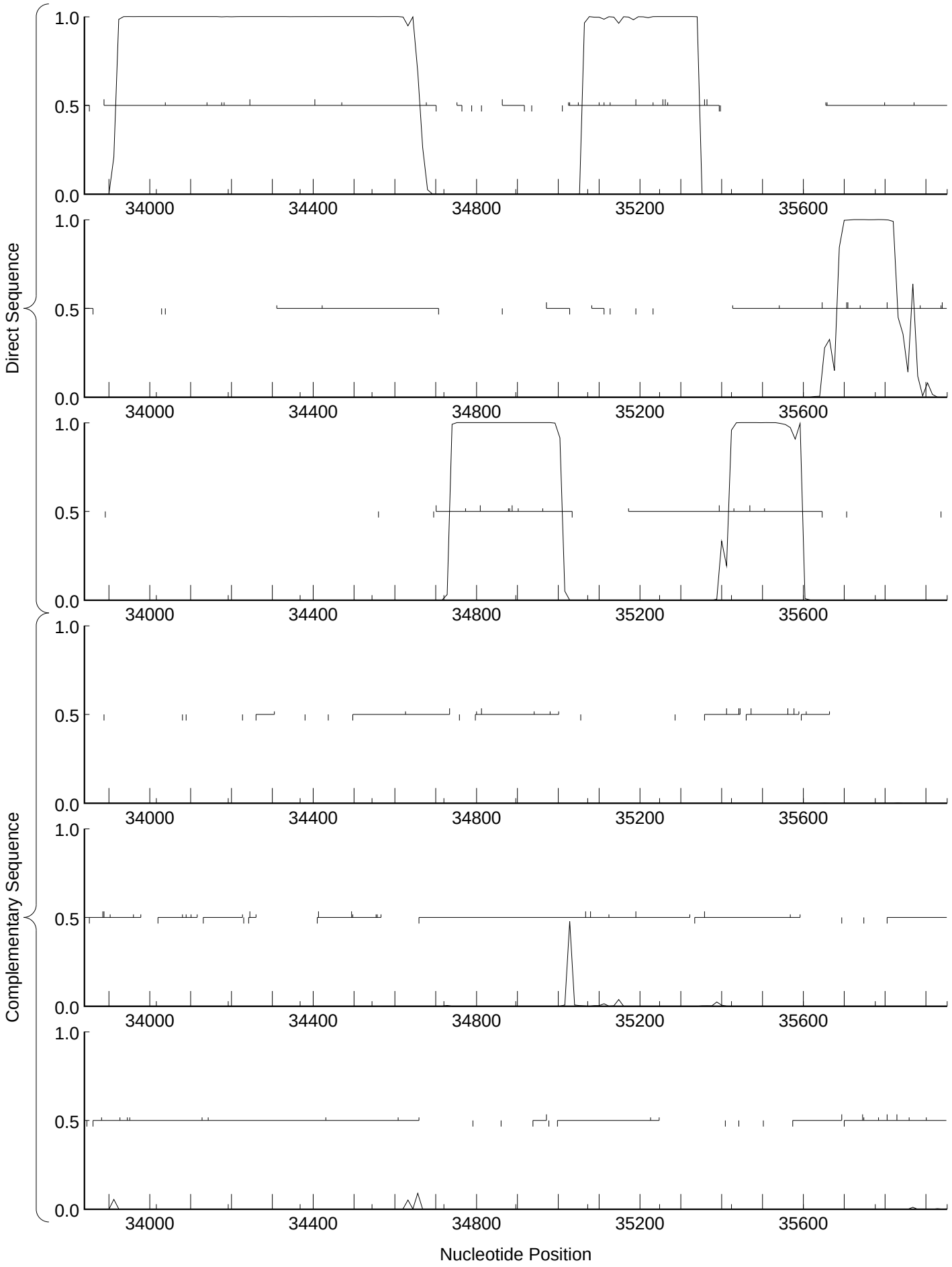
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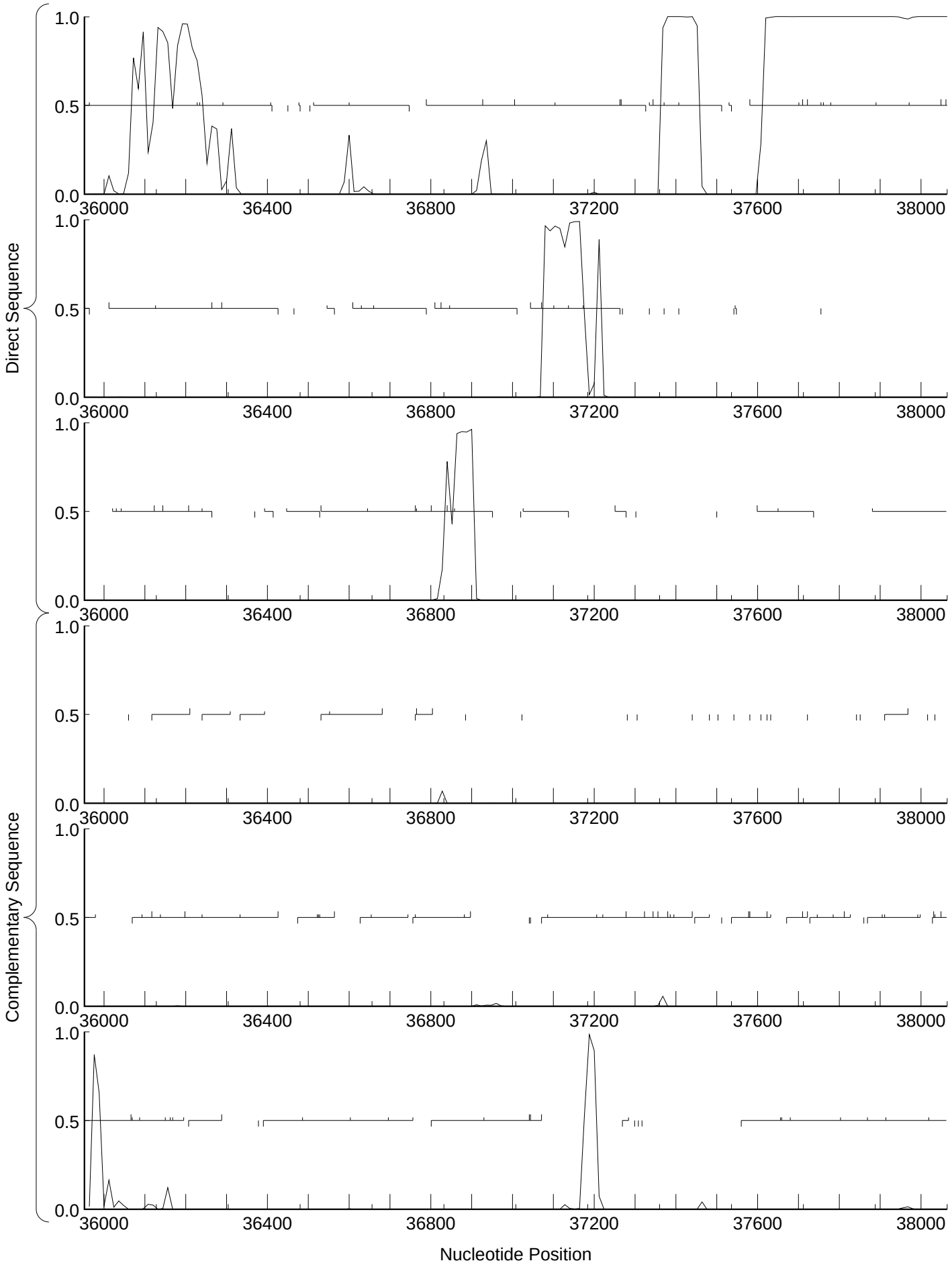
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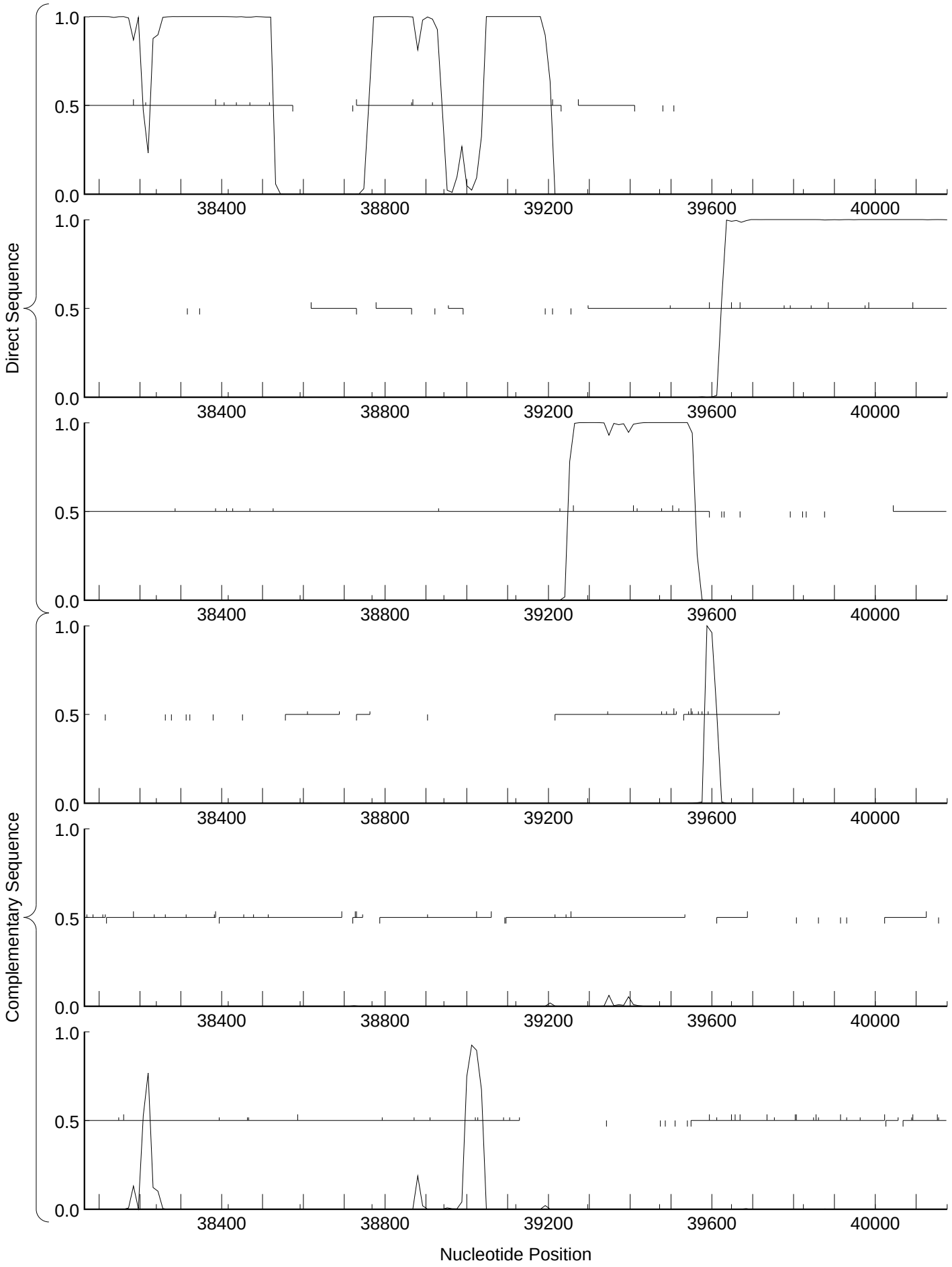
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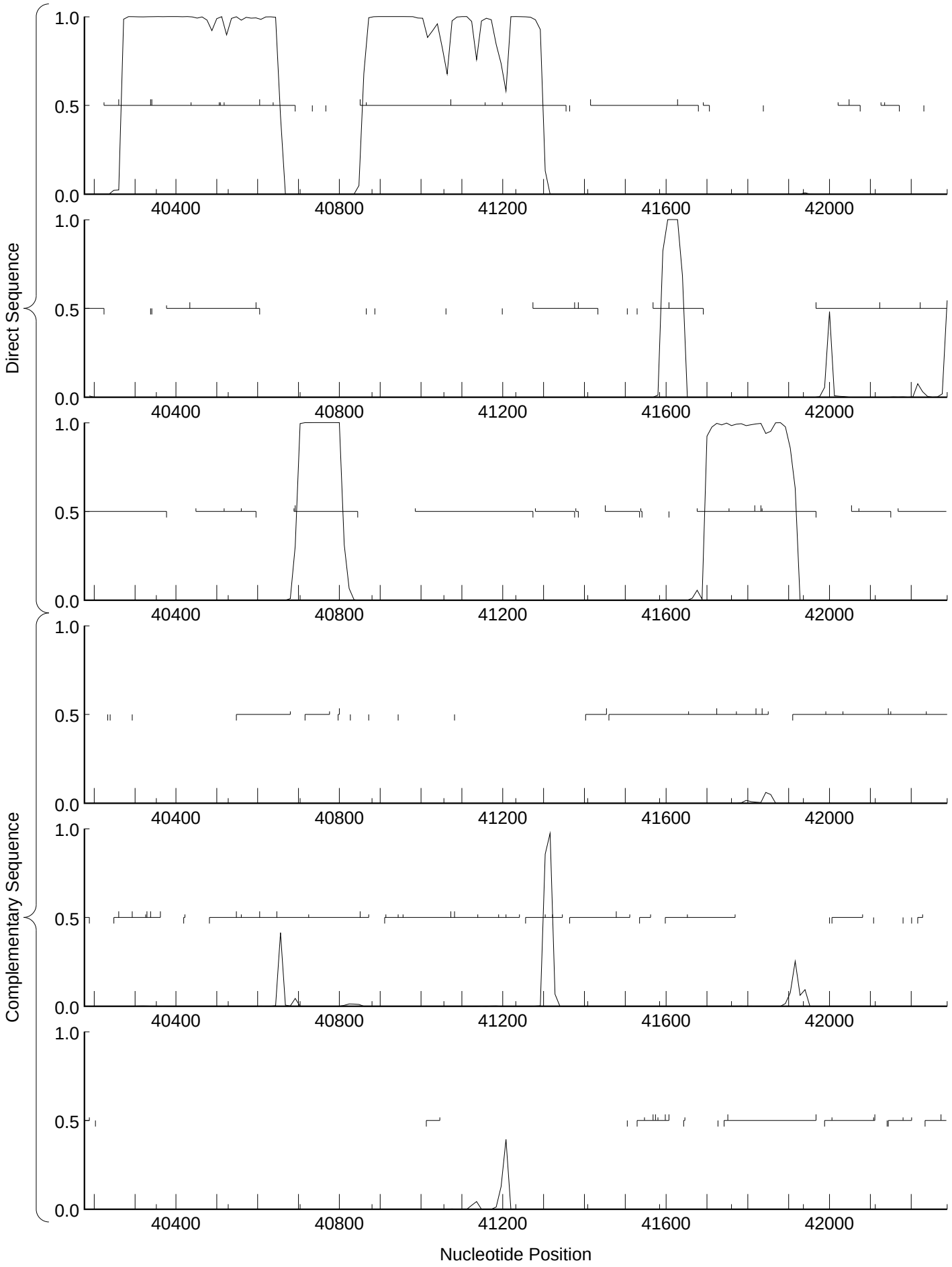
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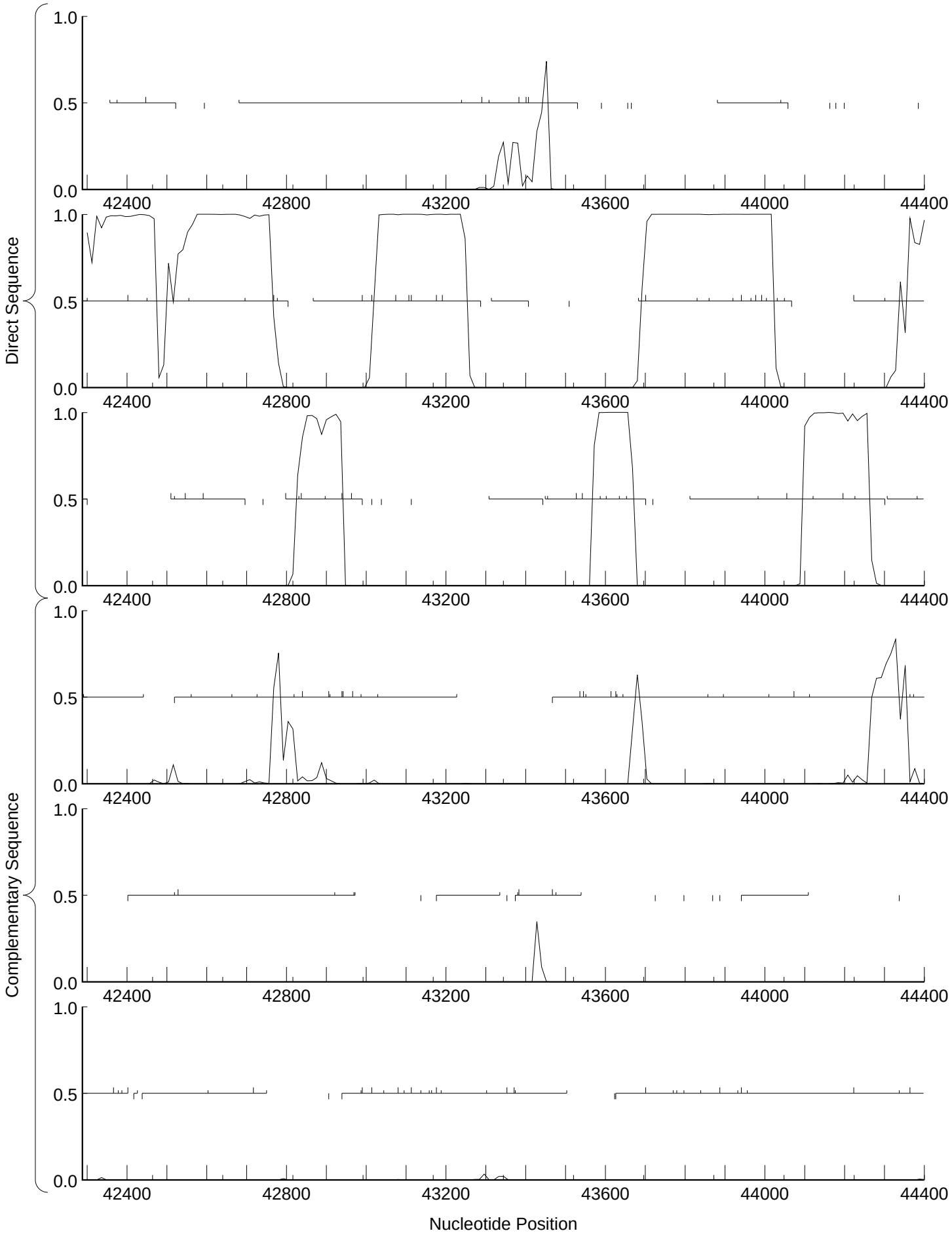
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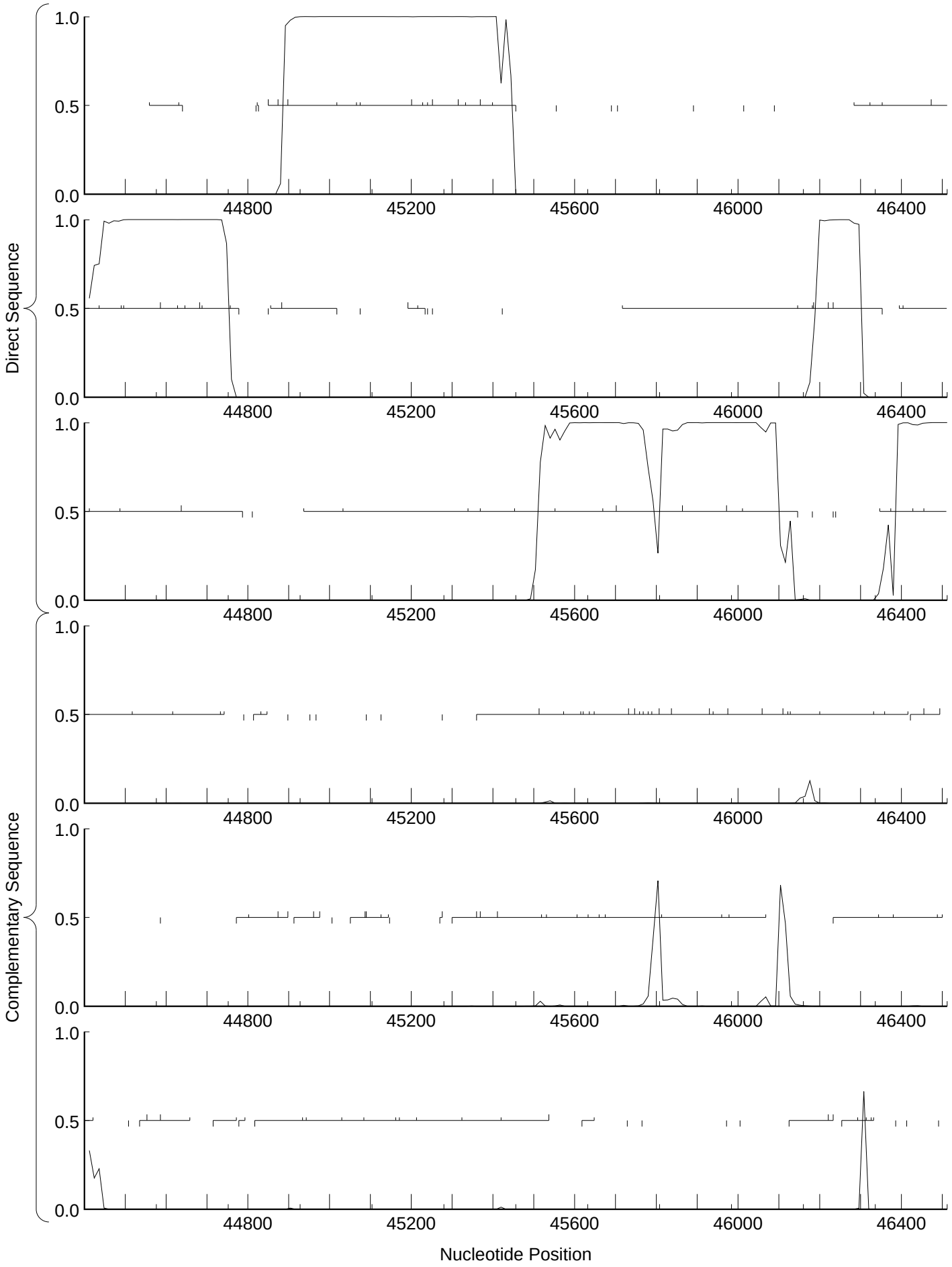
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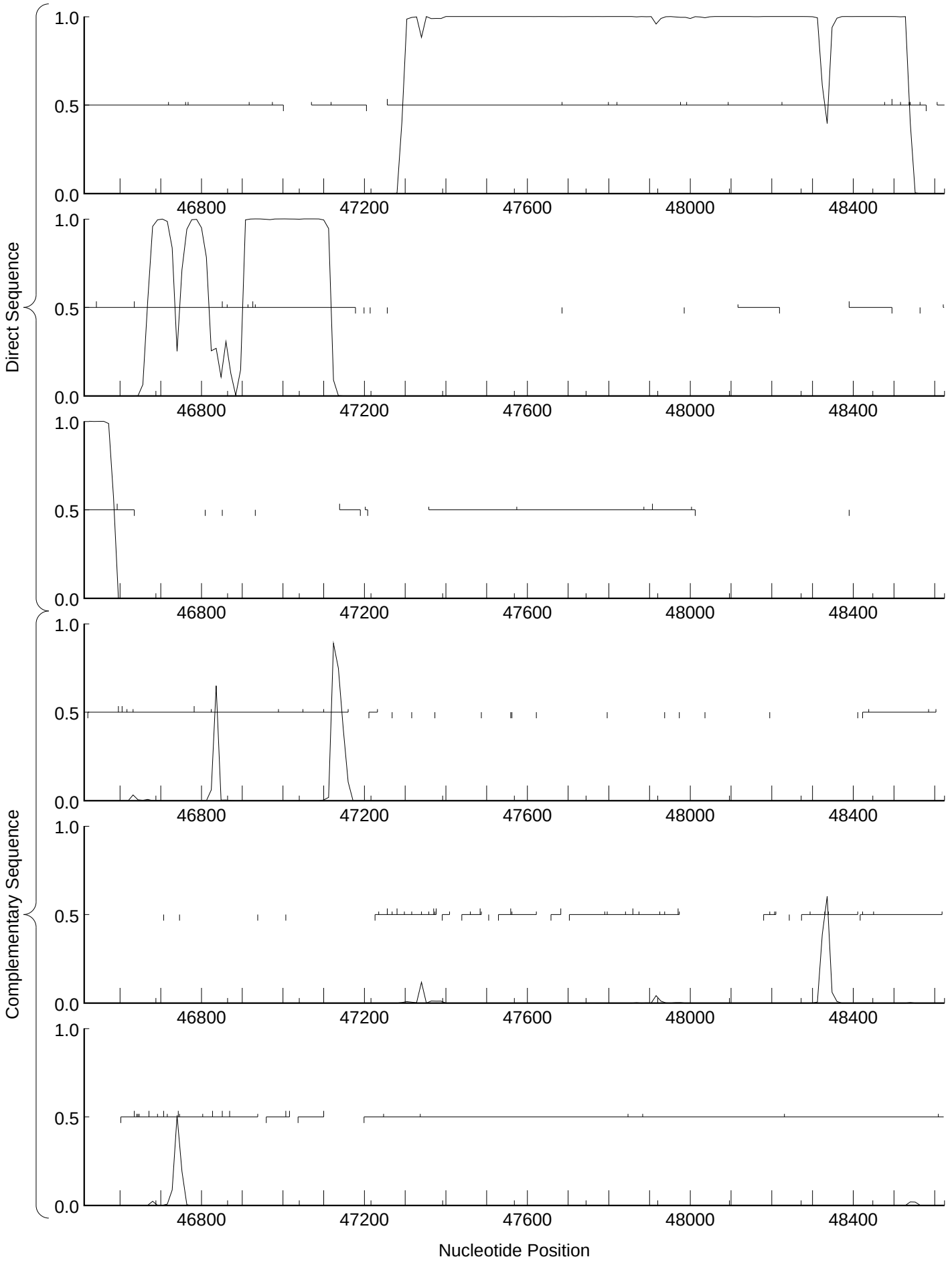
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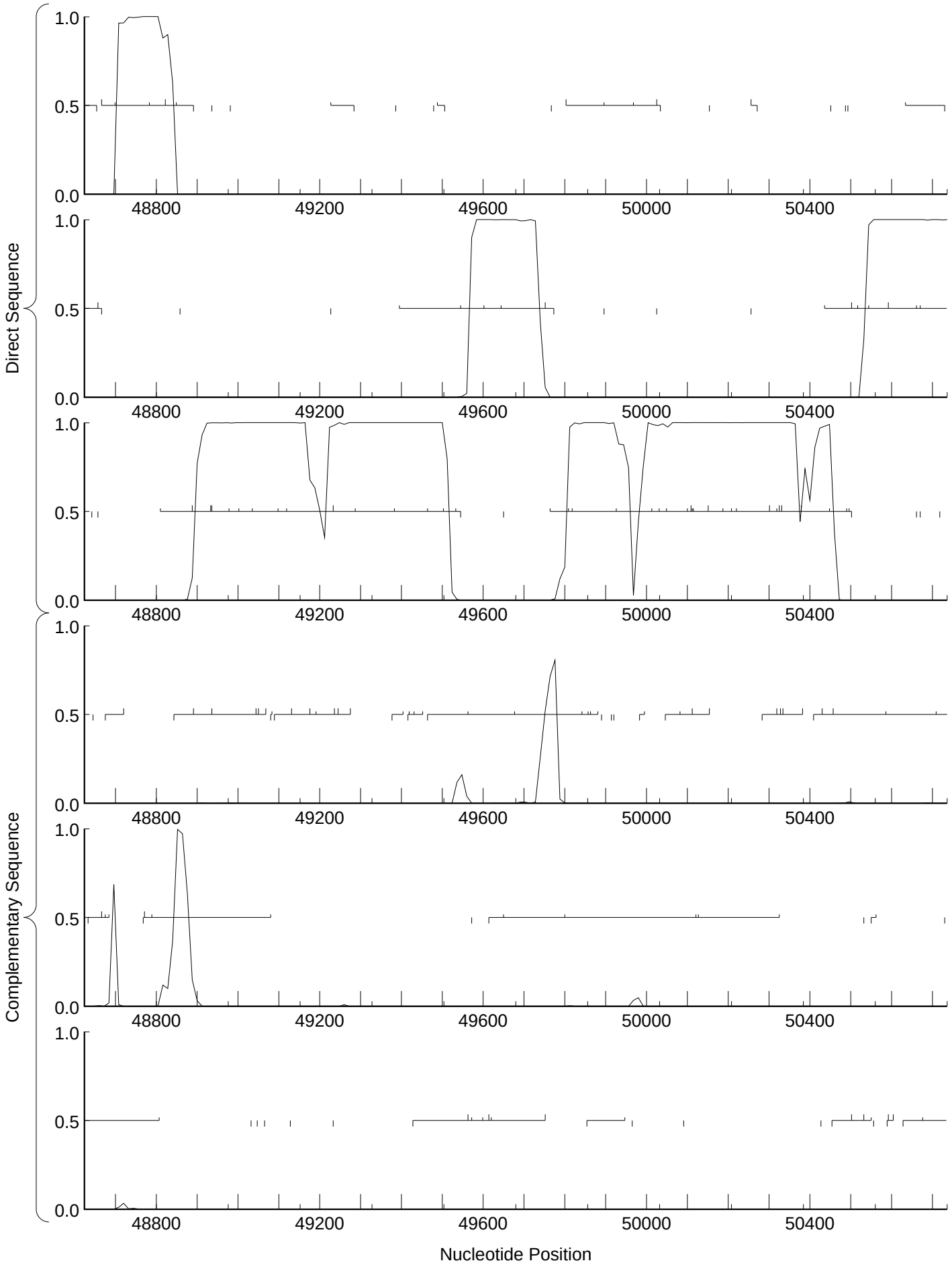
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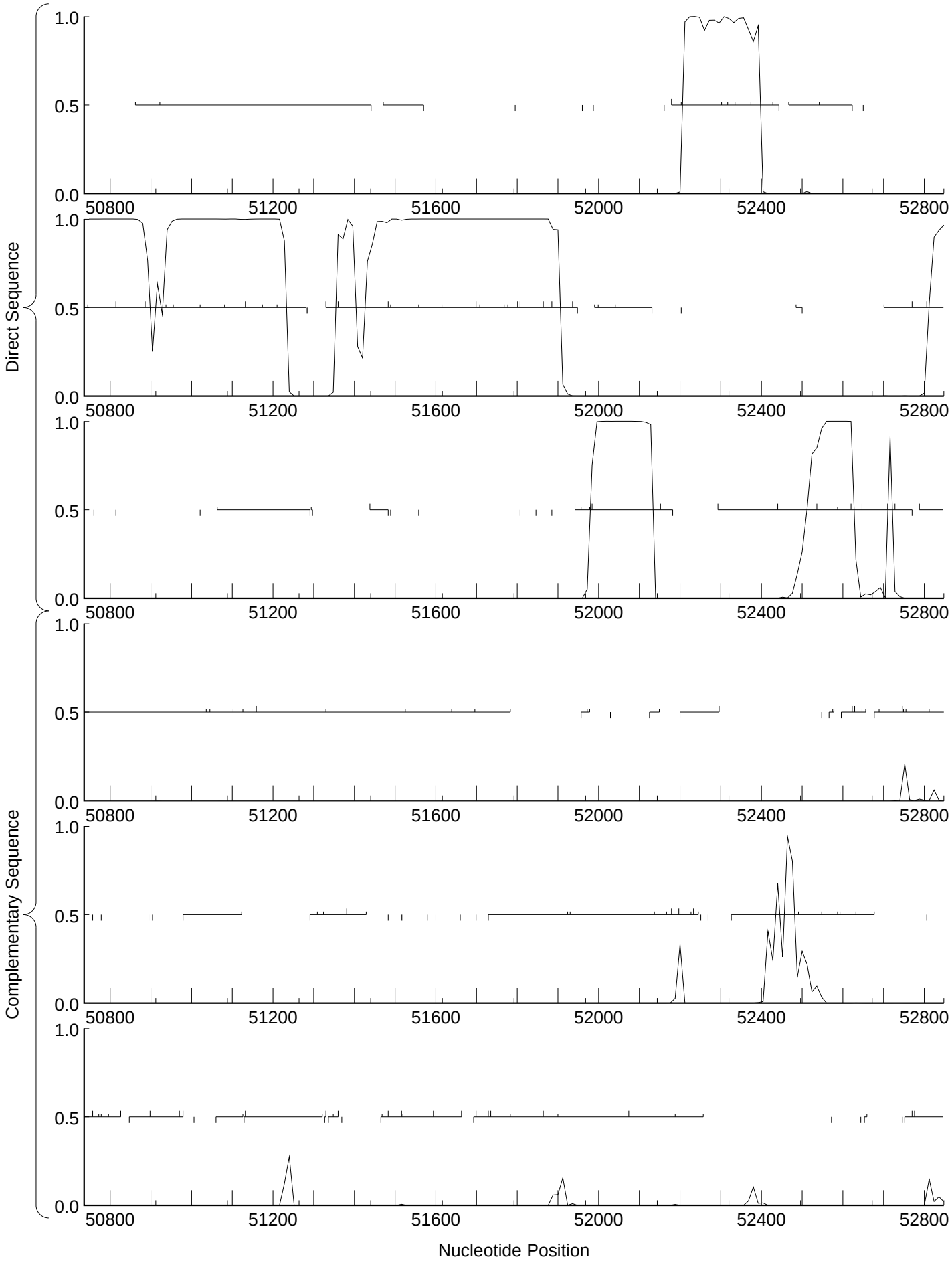
acterium phage LittleFortune complete sequence, 53161 bp, circularly permuted, Cluster EC, Order 4, Window 96, Step 12, 24/27



acterium phage LittleFortune complete sequence, 53161 bp, circularly permuted, Cluster EC, Order 4, Window 96, Step 12, 25/27



acterium phage LittleFortune complete sequence, 53161 bp, circularly permuted, Cluster EC, Order 4, Window 96, Step 12, 26/27



acterium phage LittleFortune complete sequence, 53161 bp, circularly permuted, Cluster EC, Order 4, Window 96, Step 12, 27/27

