GeneMark

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

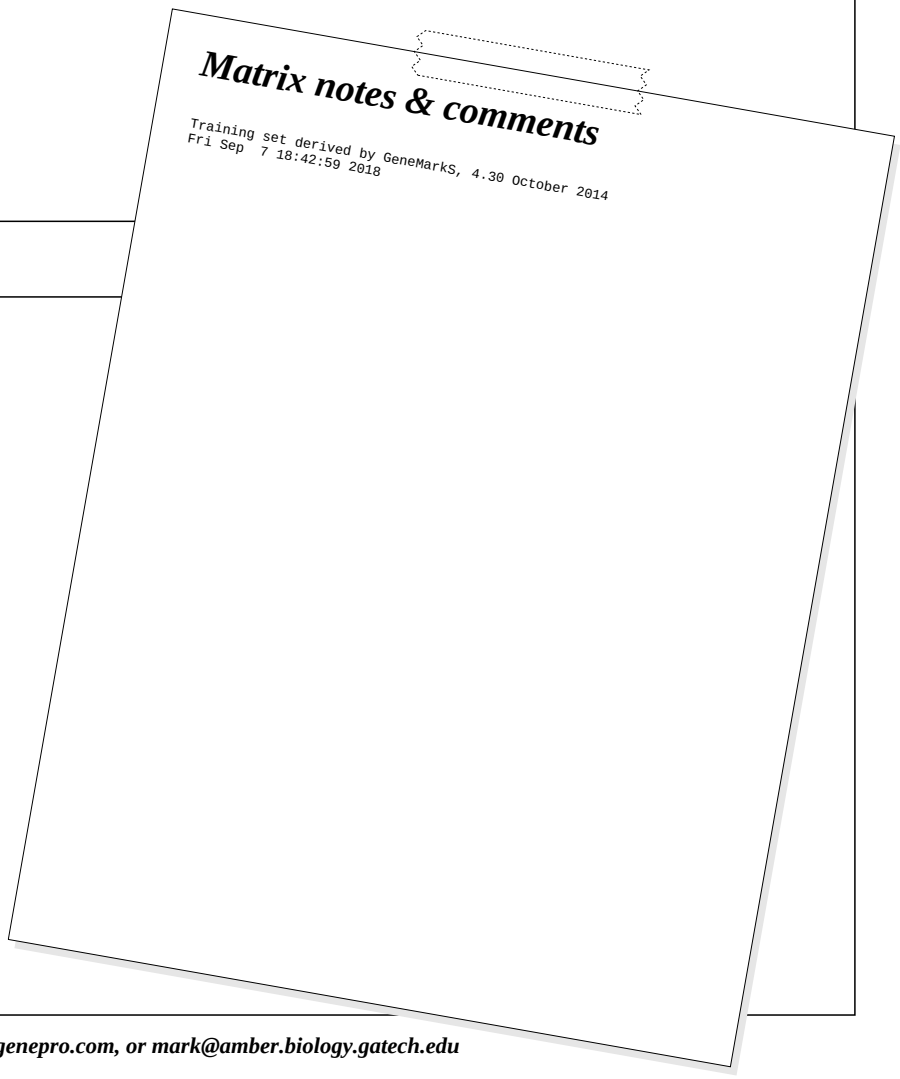
PROGRAM INFORMATION

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Sequence         :  Microbacterium phage KaiHaiDragon complete sequence, 52992 bp, circularly permuted, Cluster
Analysis Date    :  3/28/24 at 11:56:02
Pages            :  27
Sequence Length  :  52992 bp
GC Content       :   68.90%

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

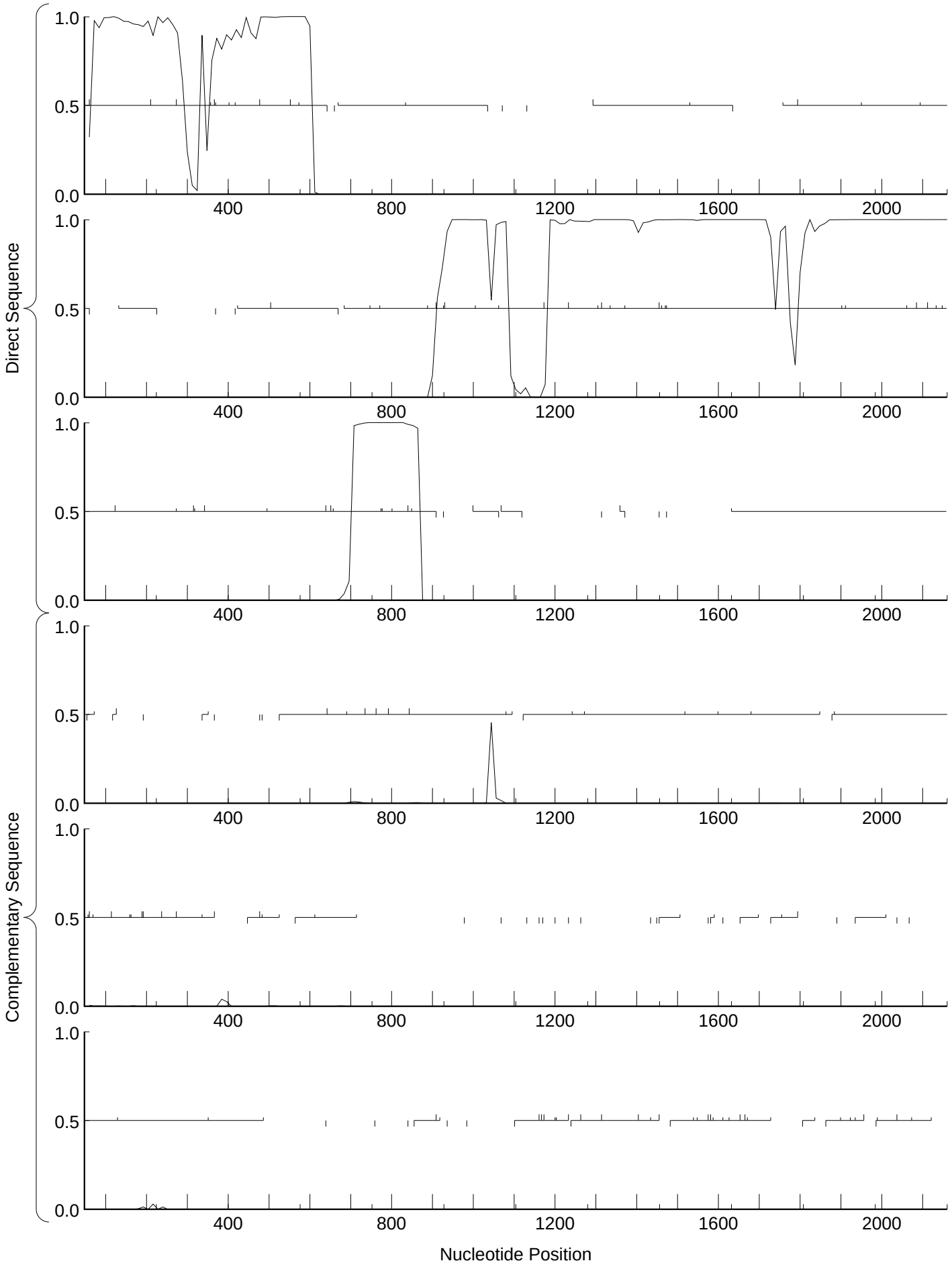
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Matrix     :  -
Author     :  -

Order      :  4
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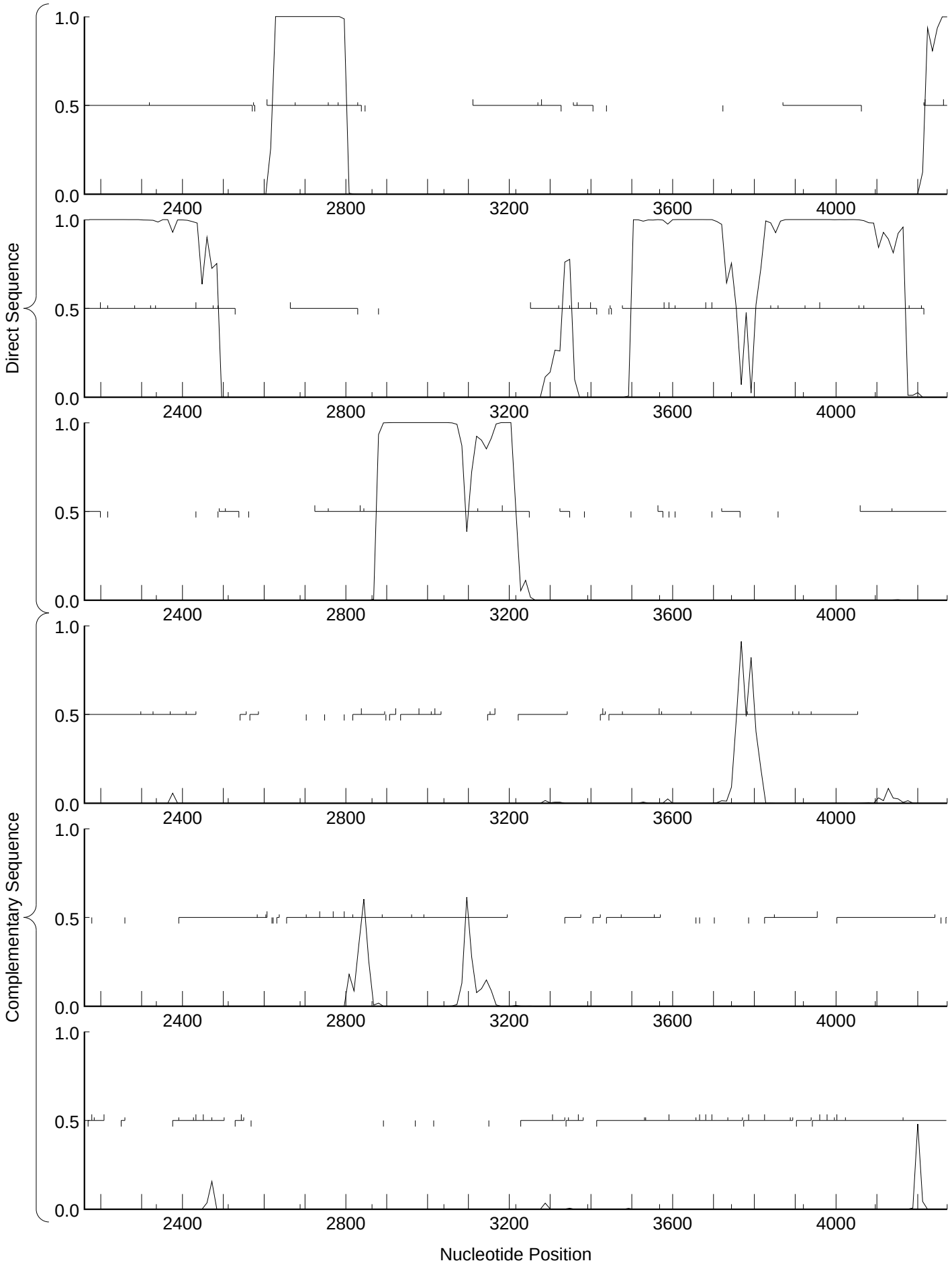

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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

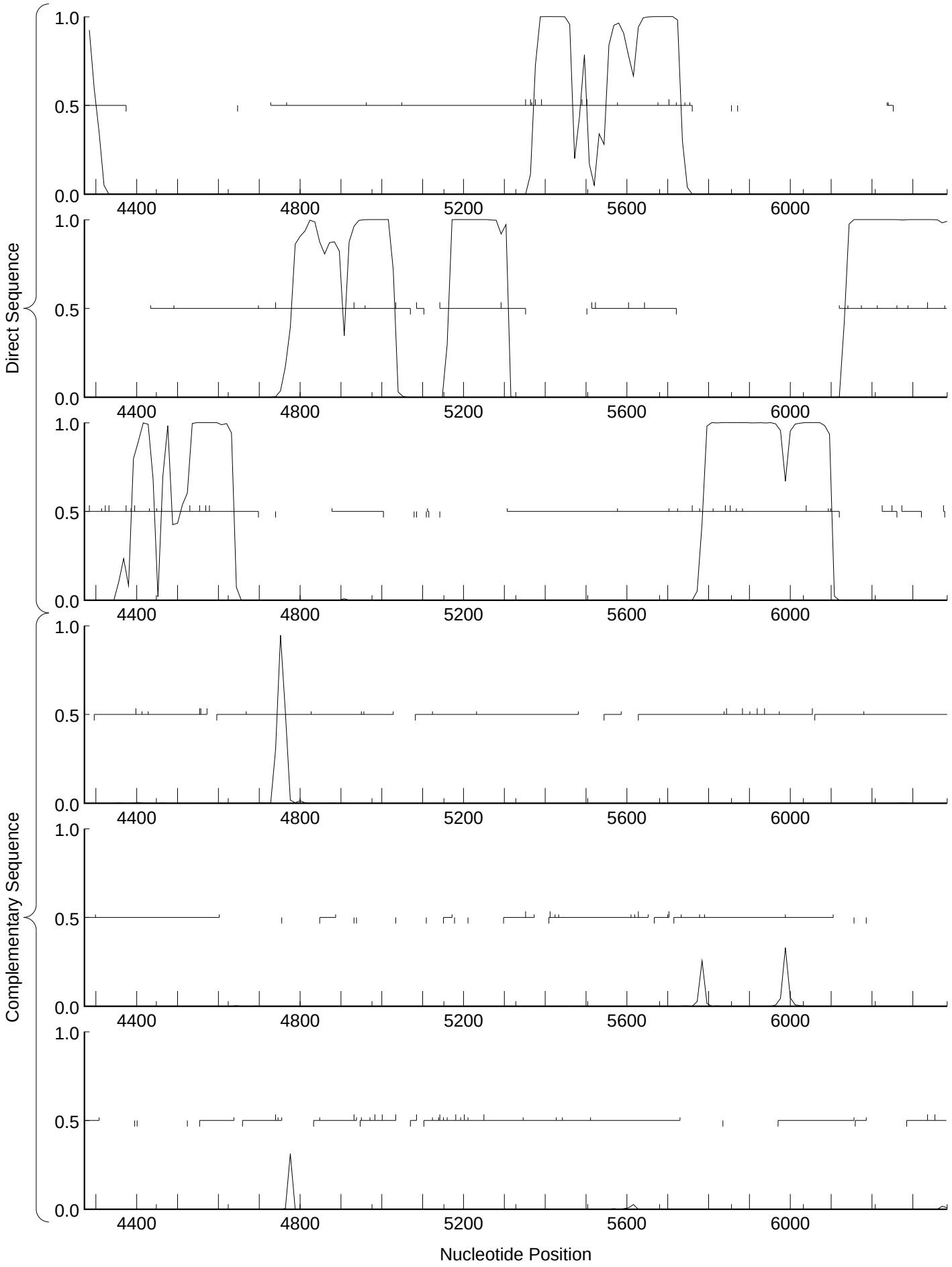

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

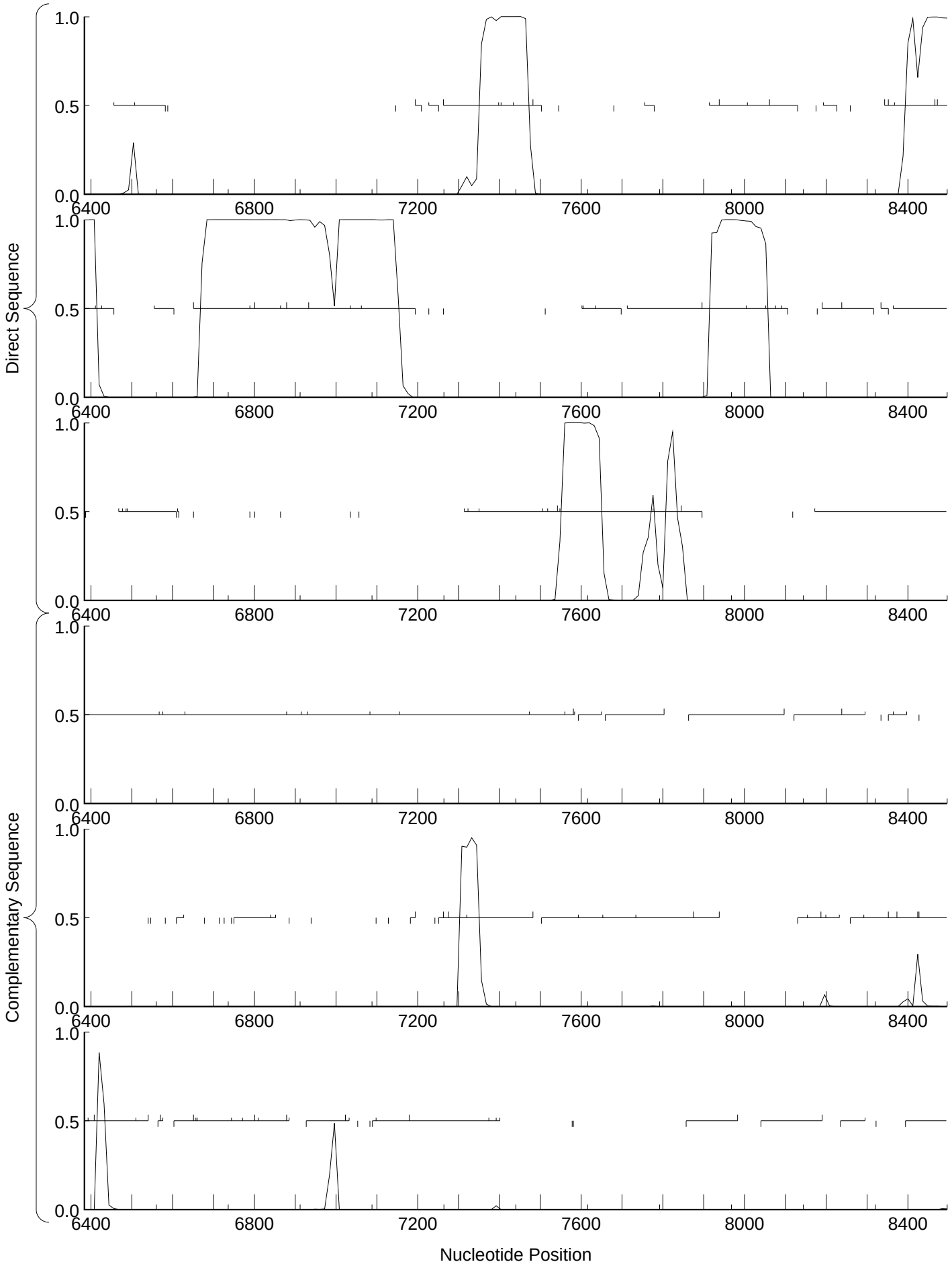
acterium phage KaiHaiDragon complete sequence, 52992 bp, circularly permuted, Cluster EC, Order 4, Window 96, Step 12, 2/27



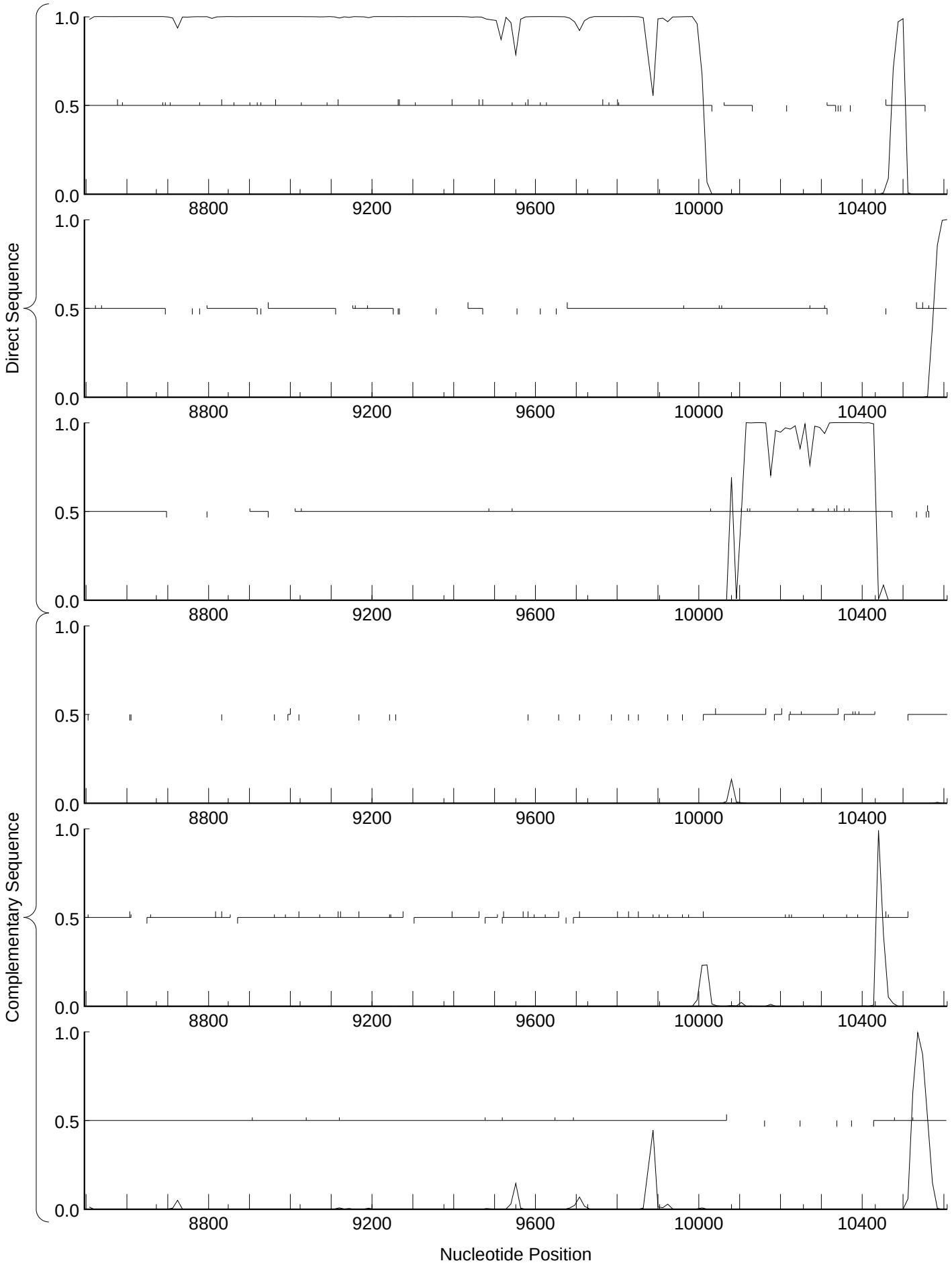
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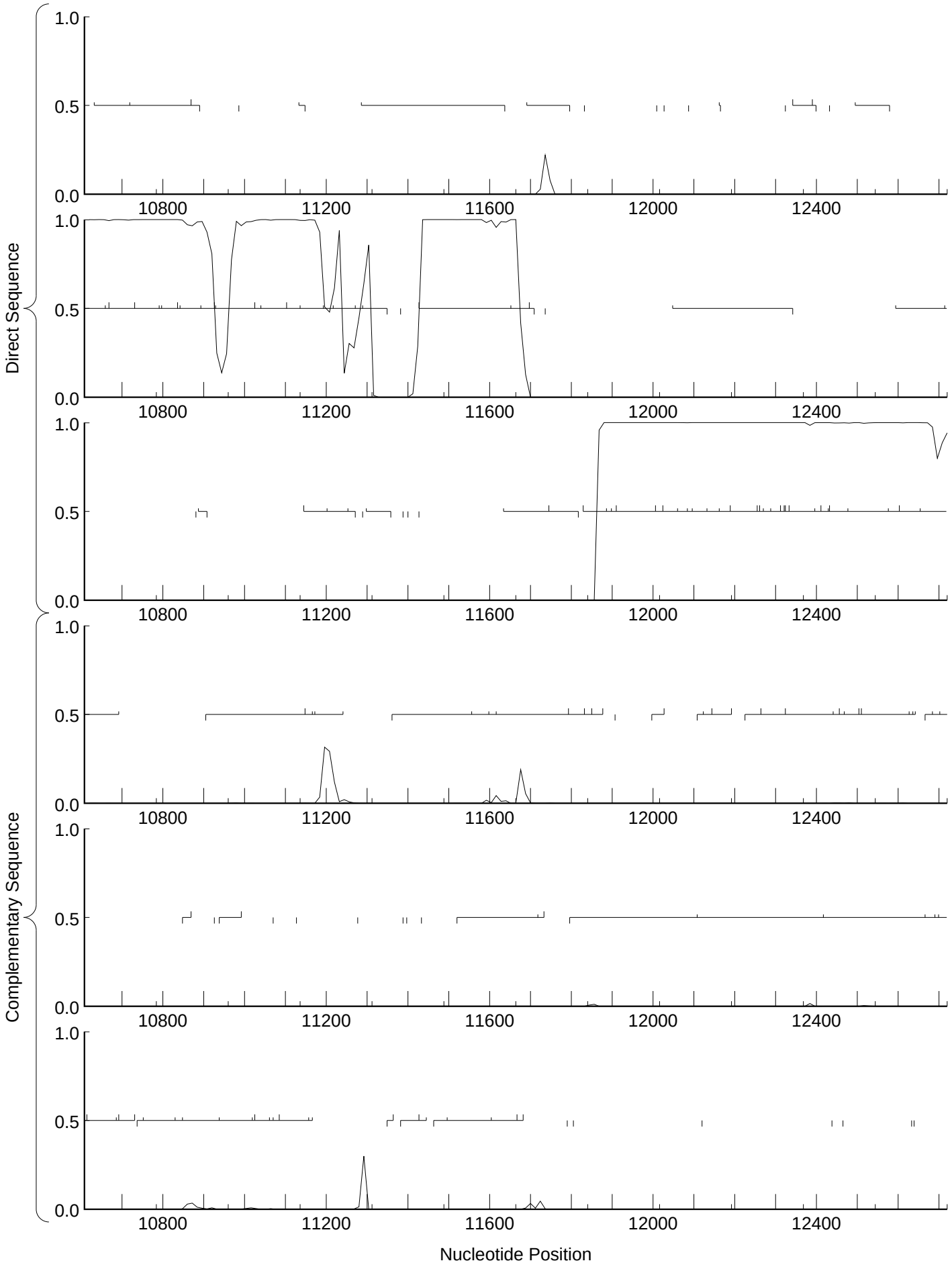

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
4400
4800
5200
5600
6000
Nucleotide Position


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

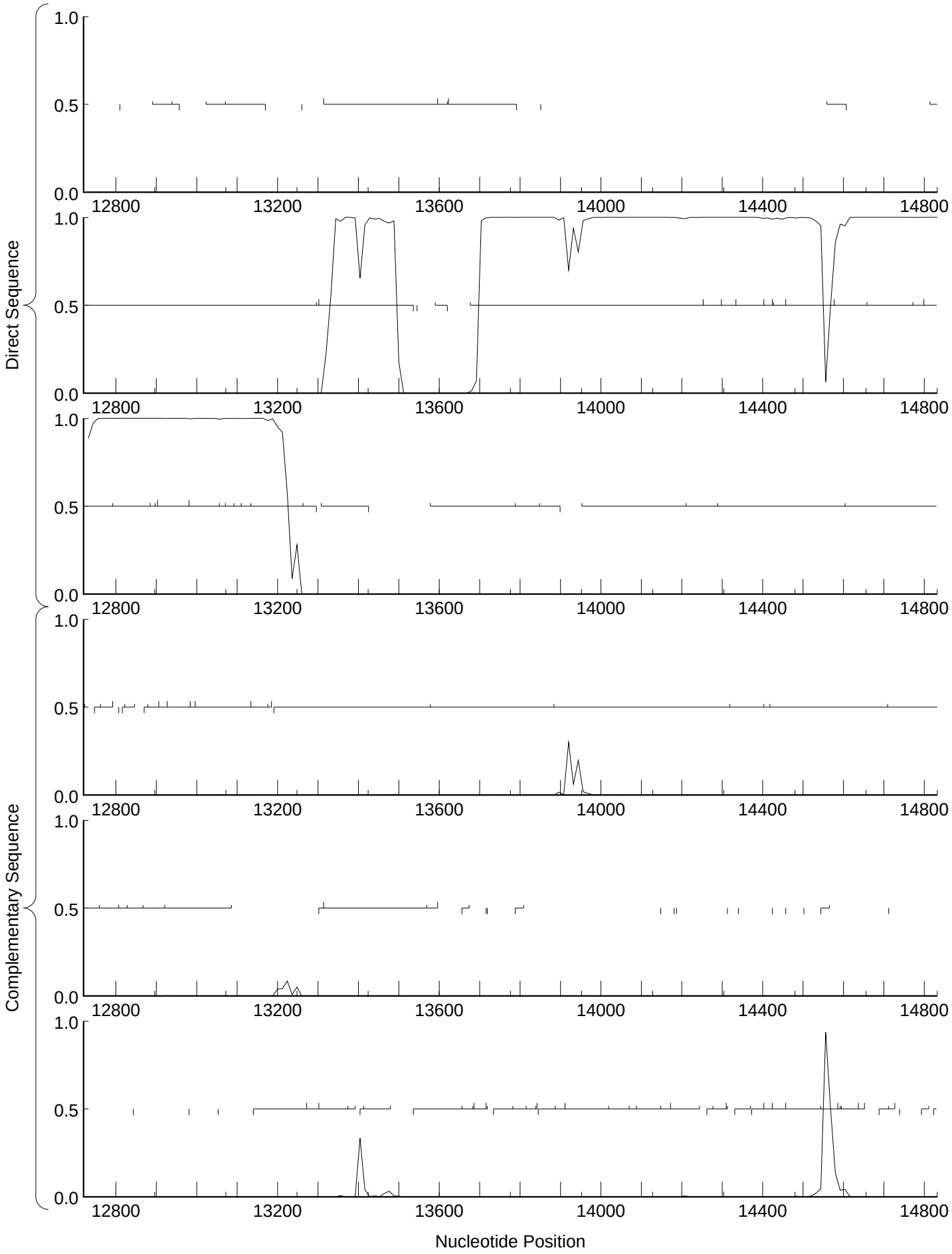
acterium phage KaiHaiDragon complete sequence, 52992 bp, circularly permuted, Cluster EC, Order 4, Window 96, Step 12, 6/27



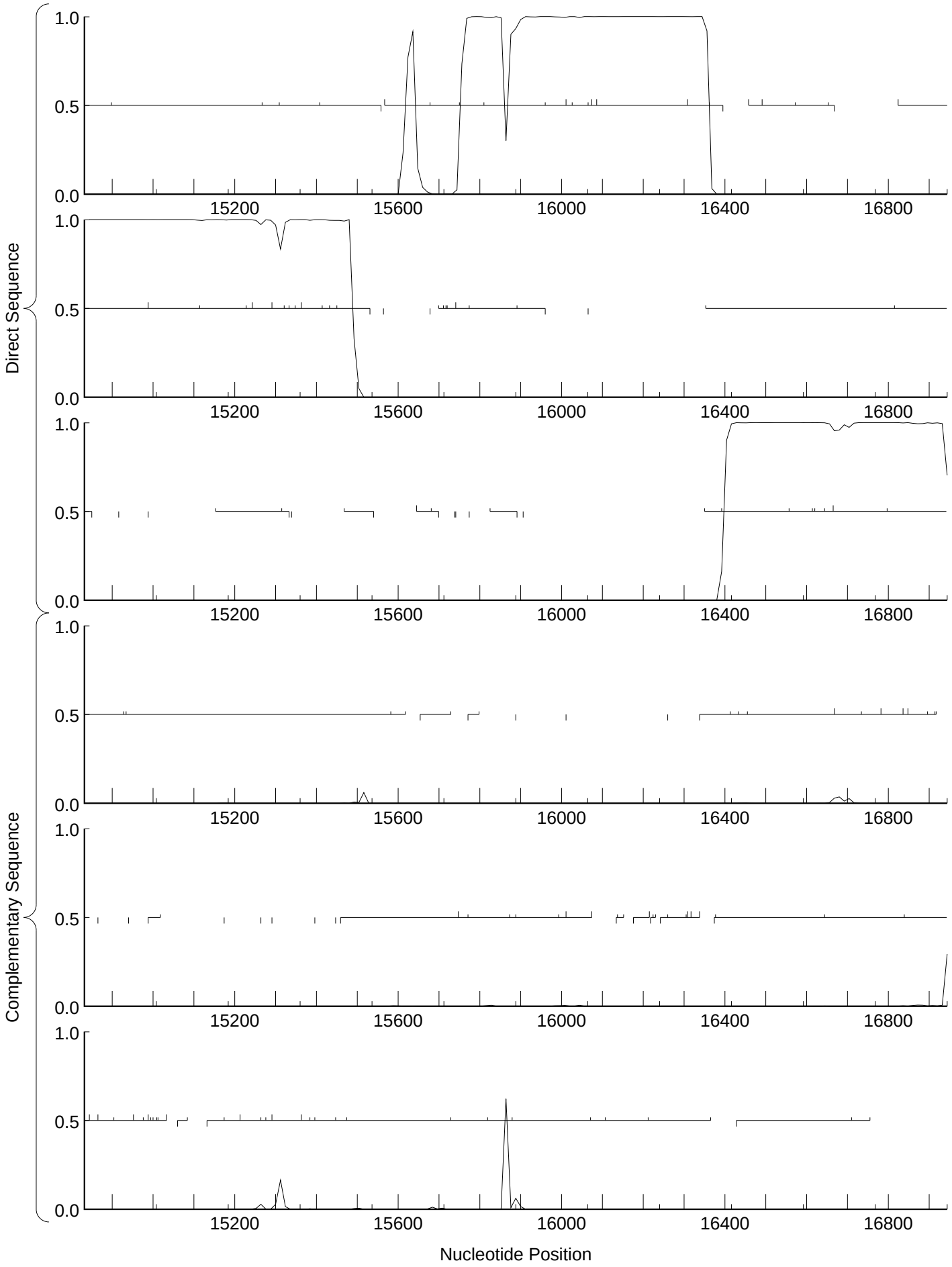
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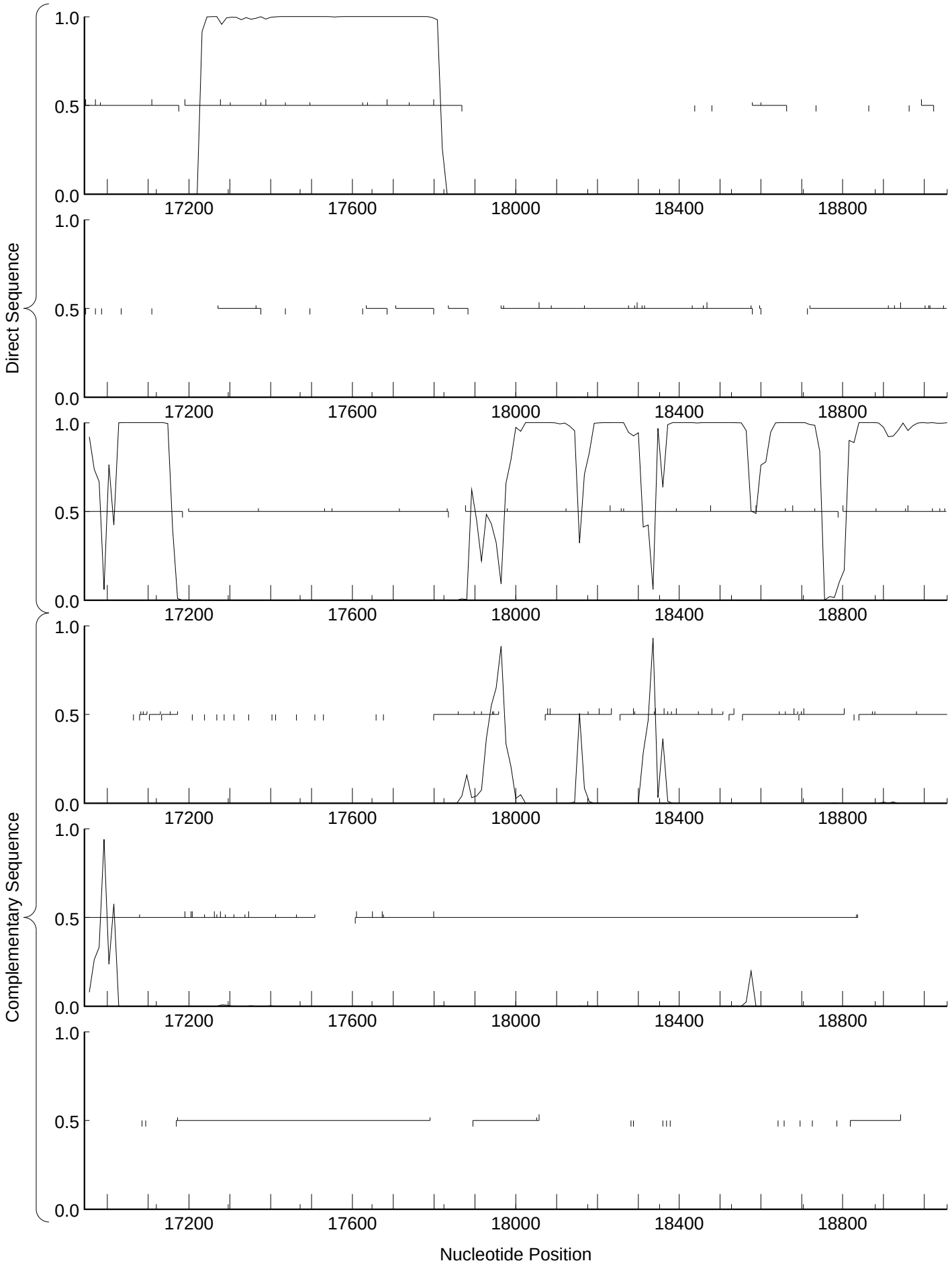
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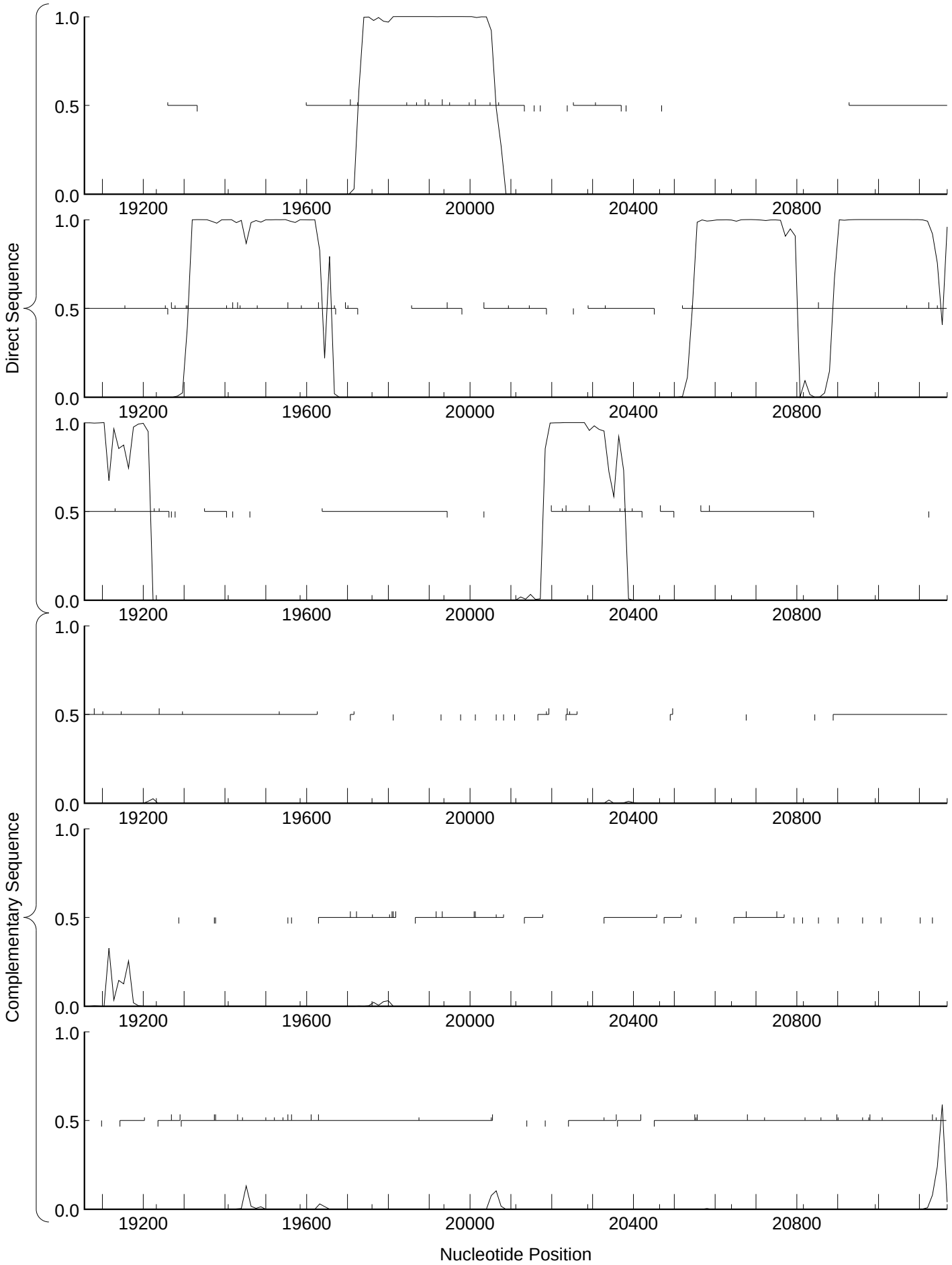
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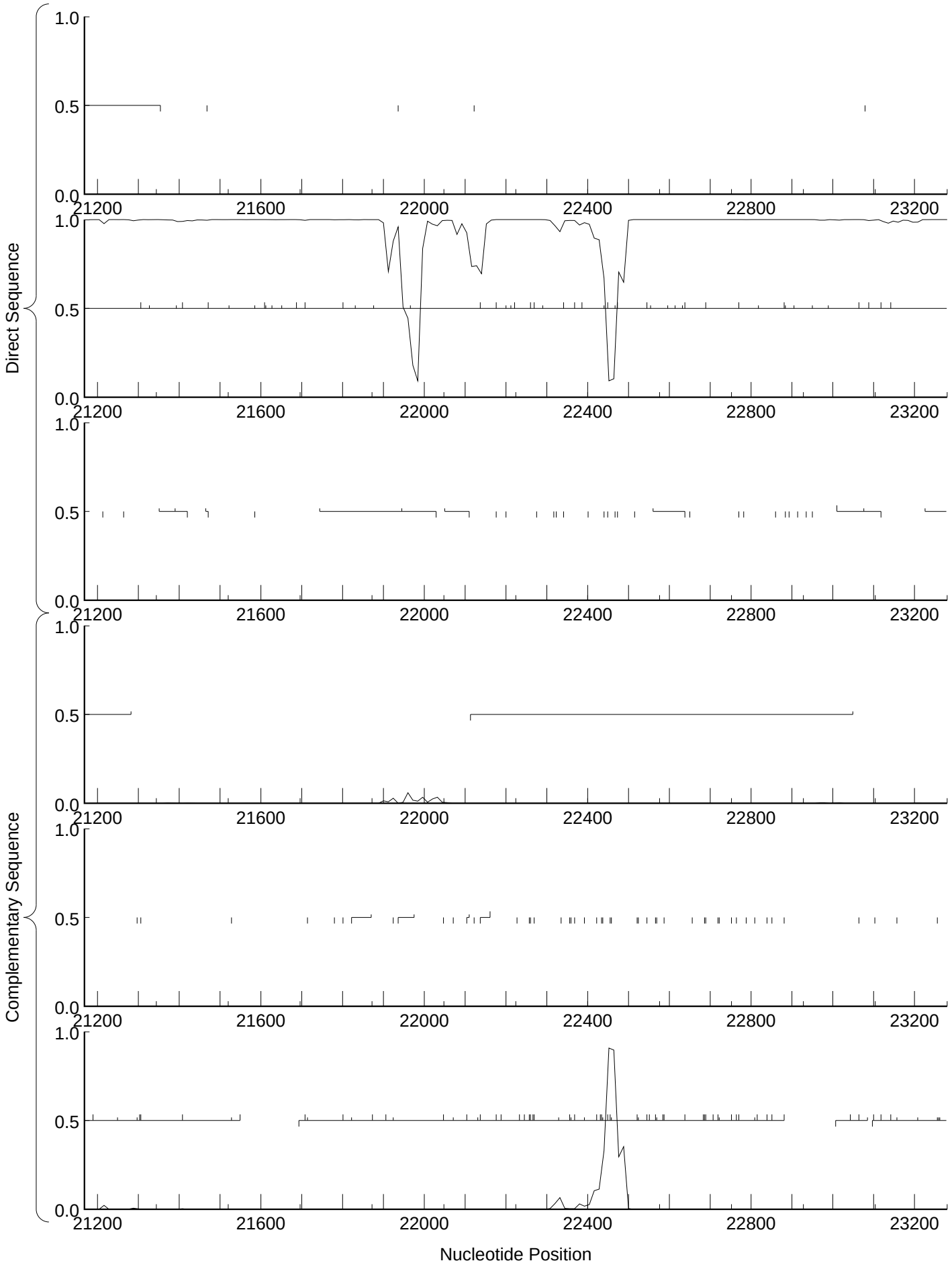
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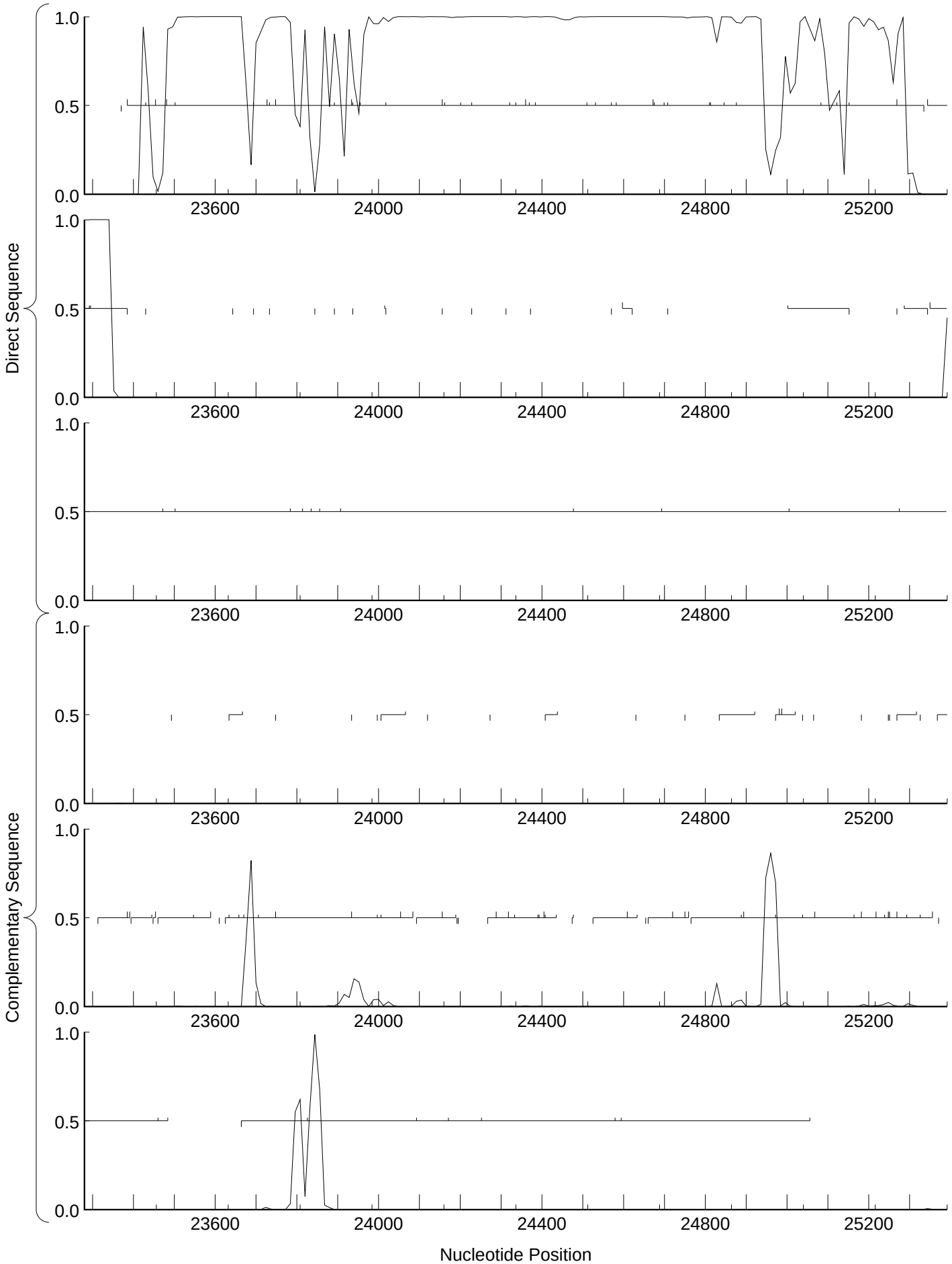
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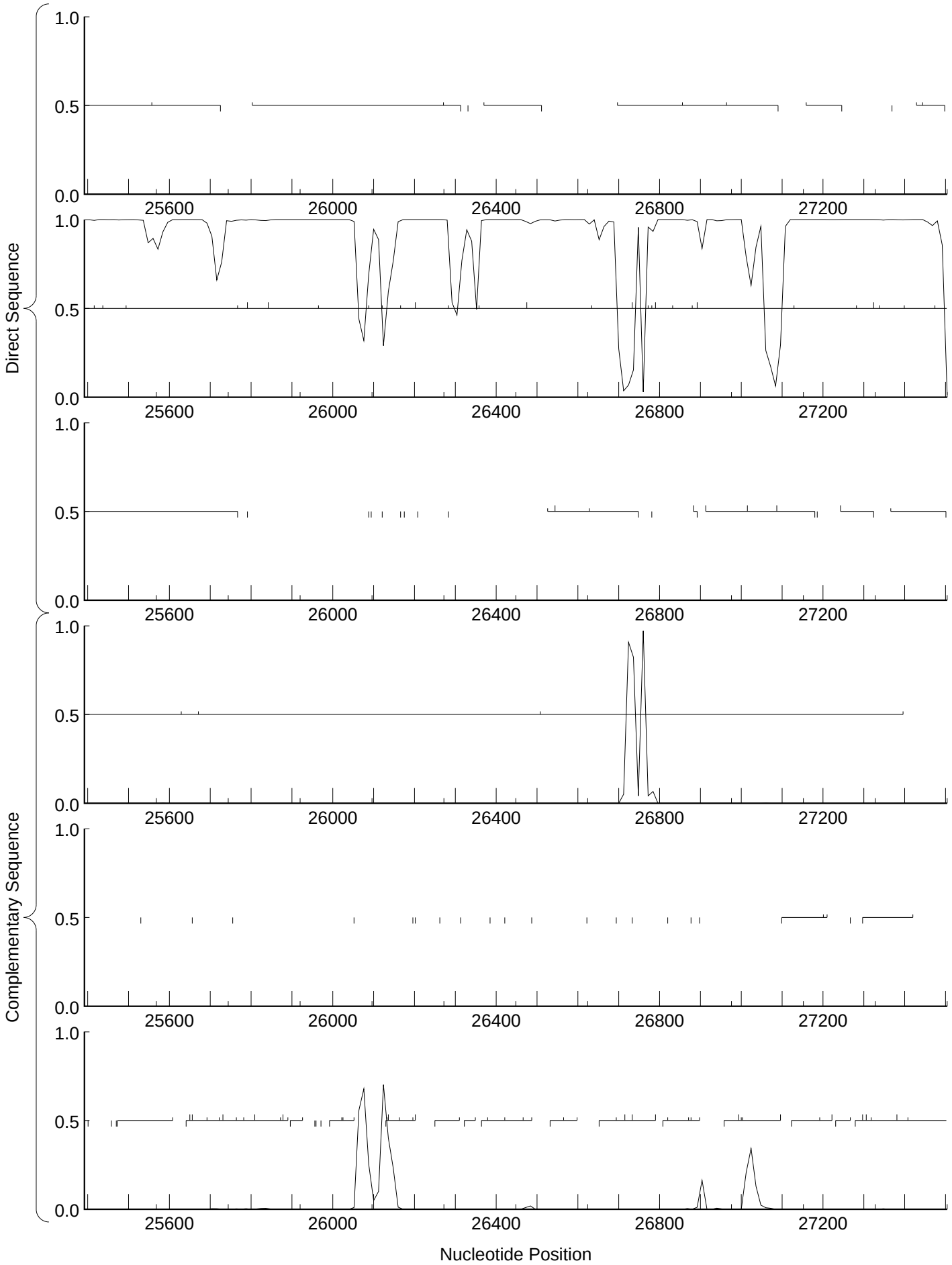
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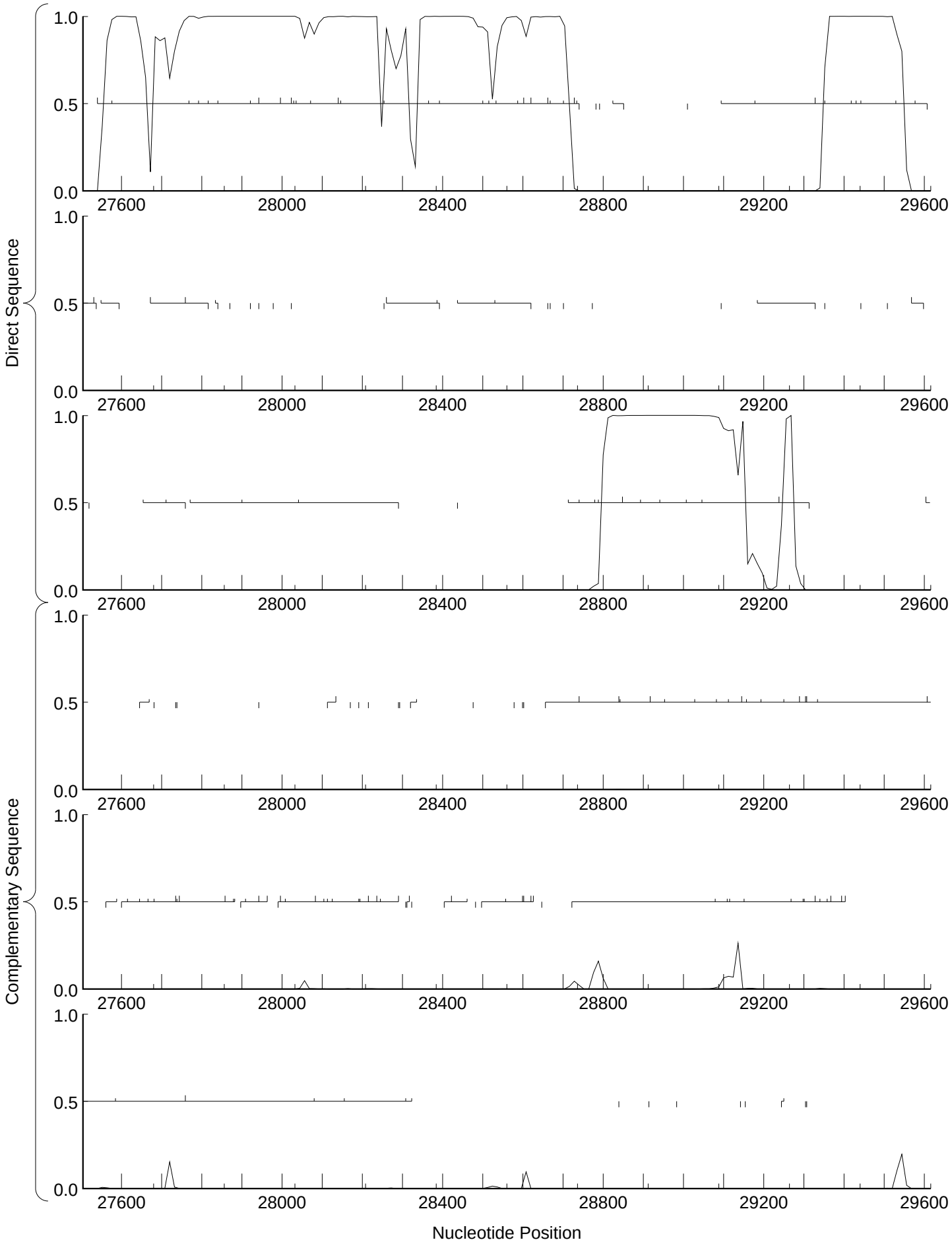
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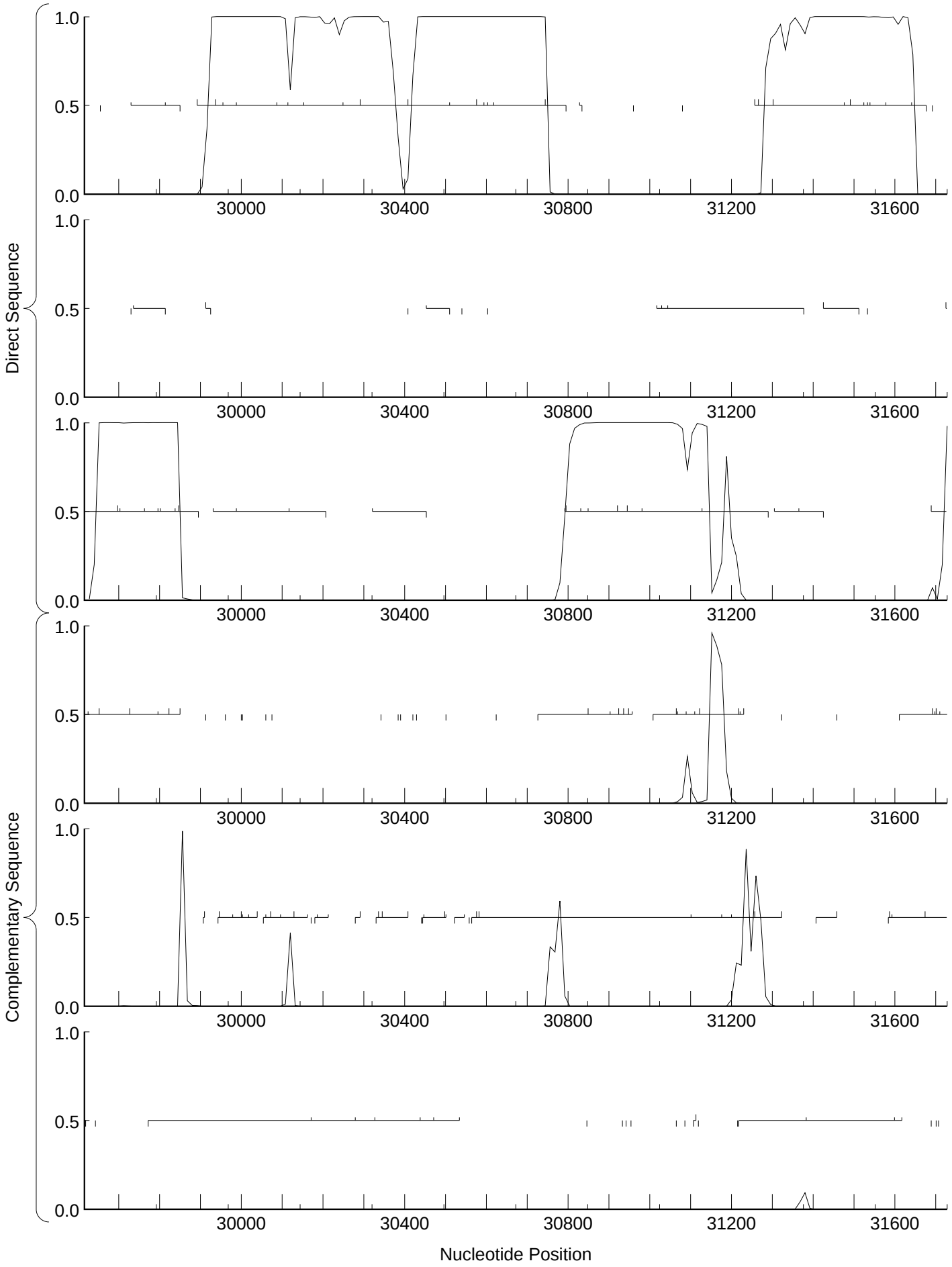
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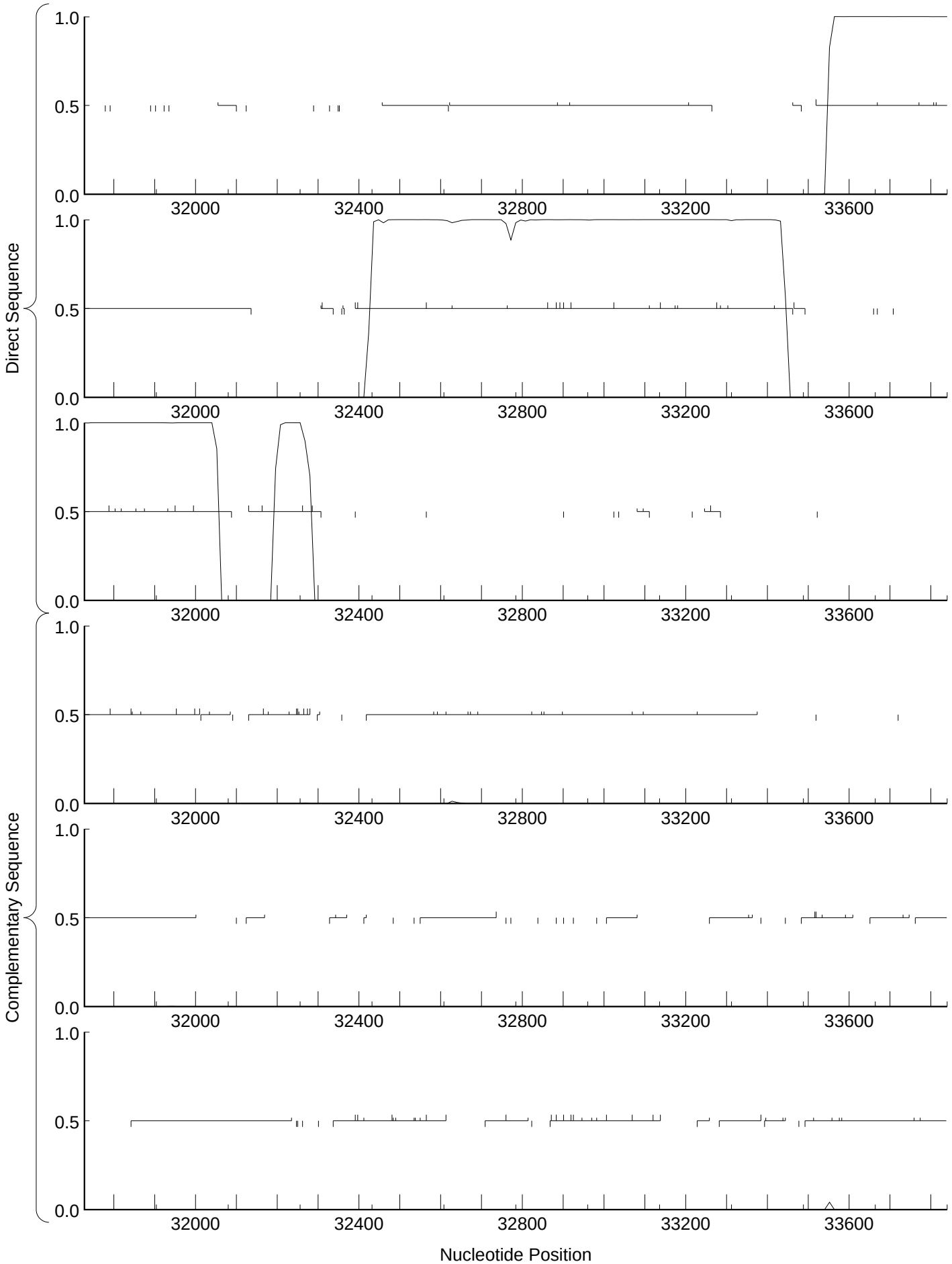
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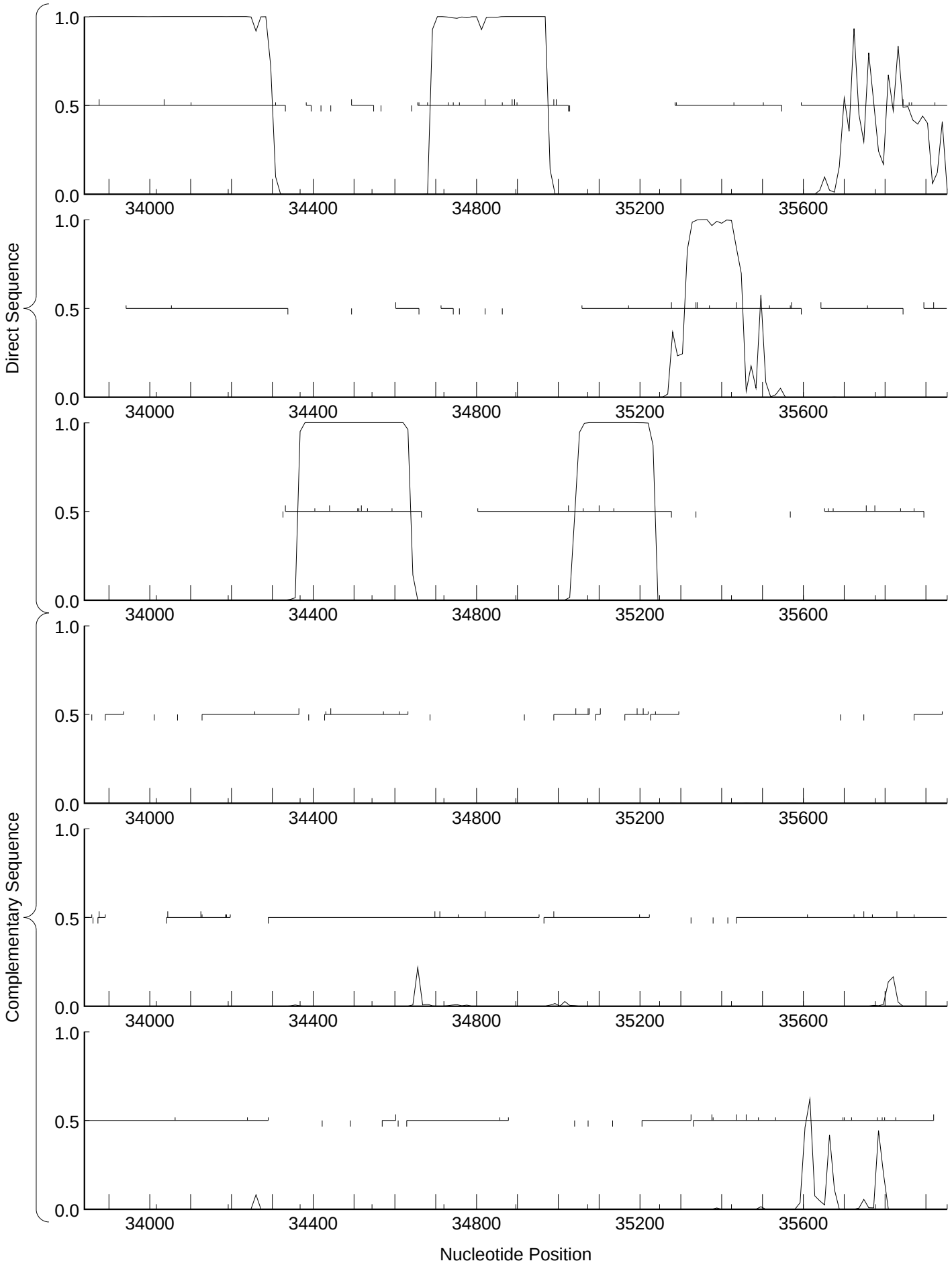
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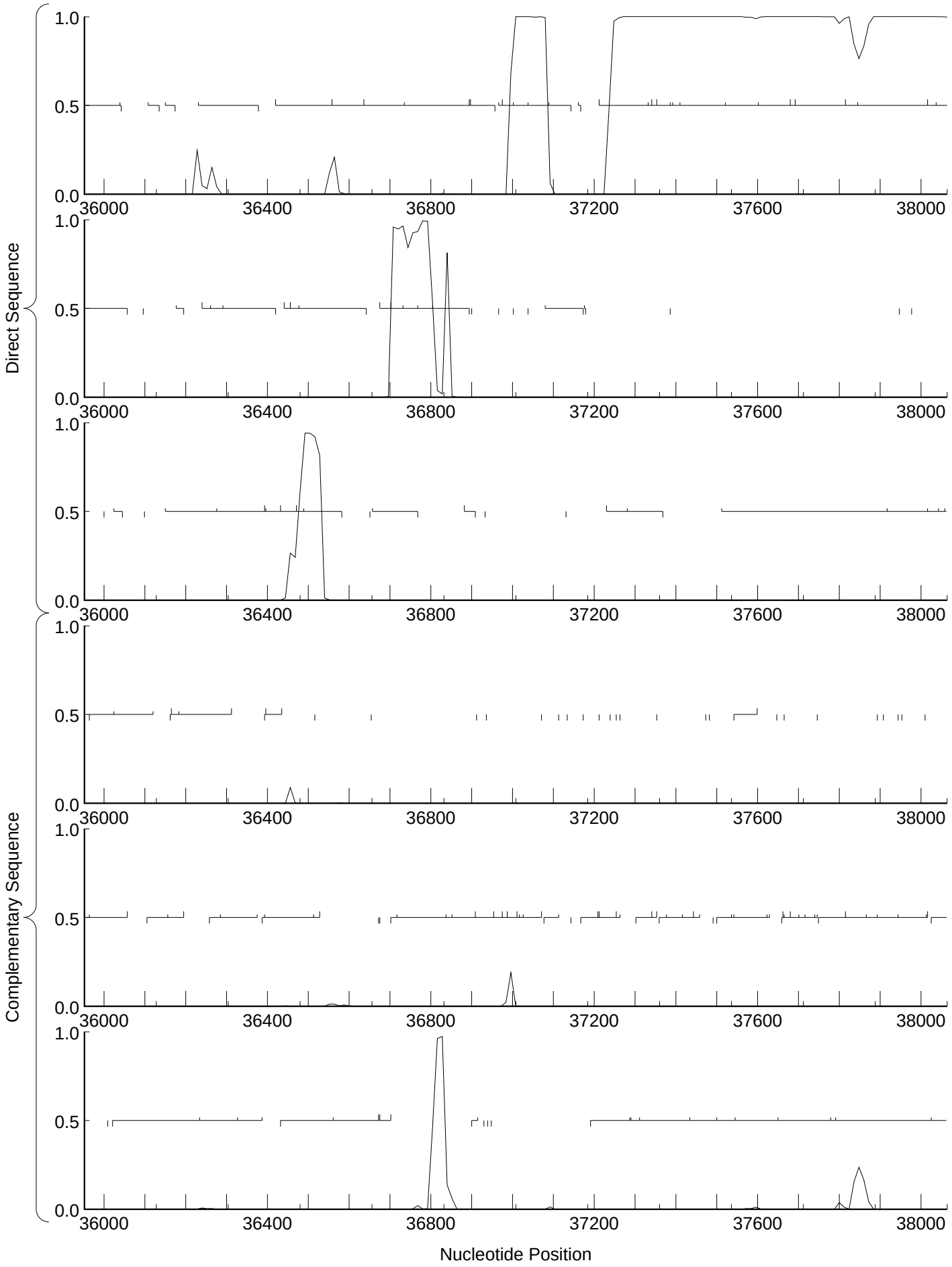
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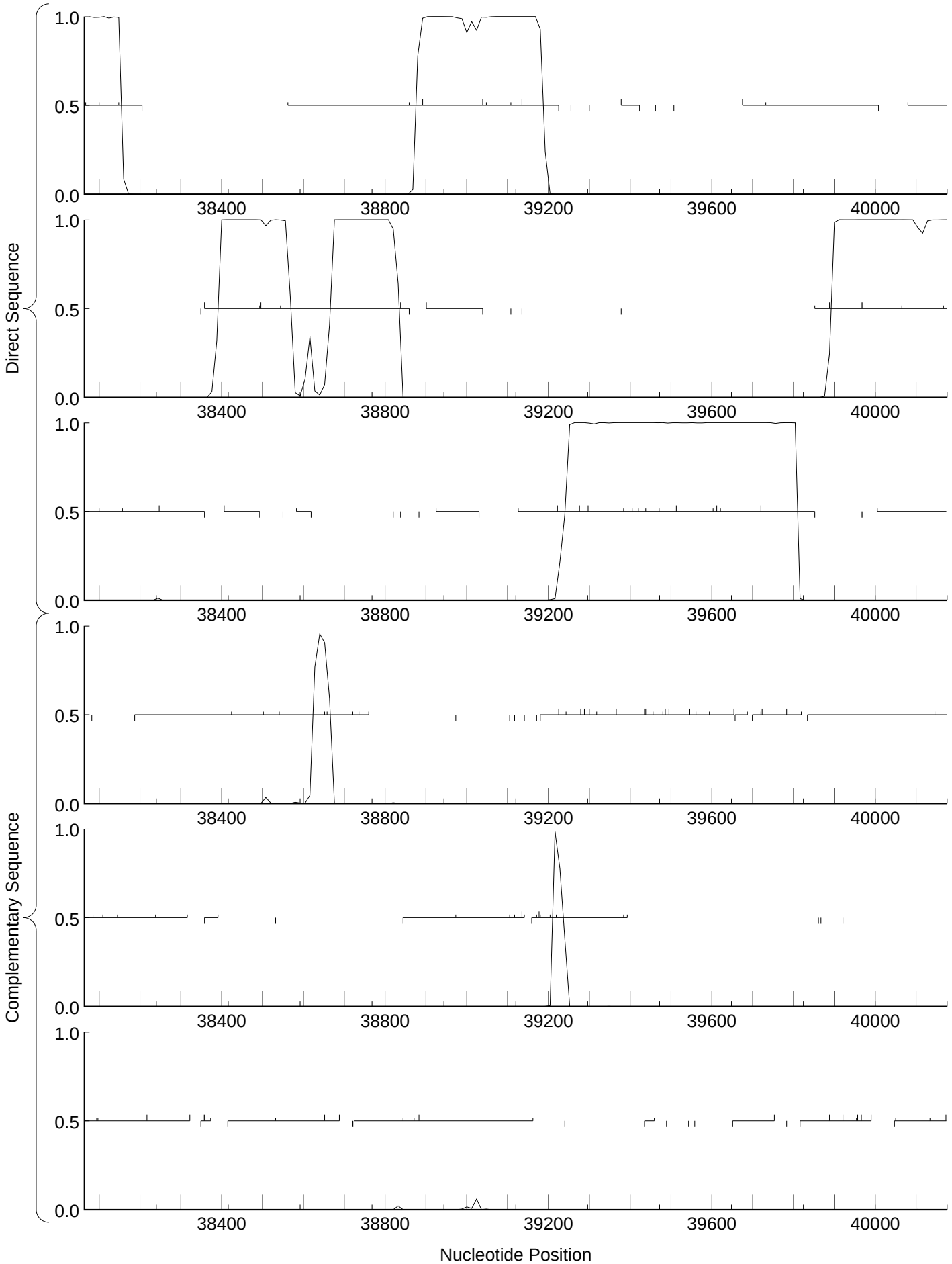
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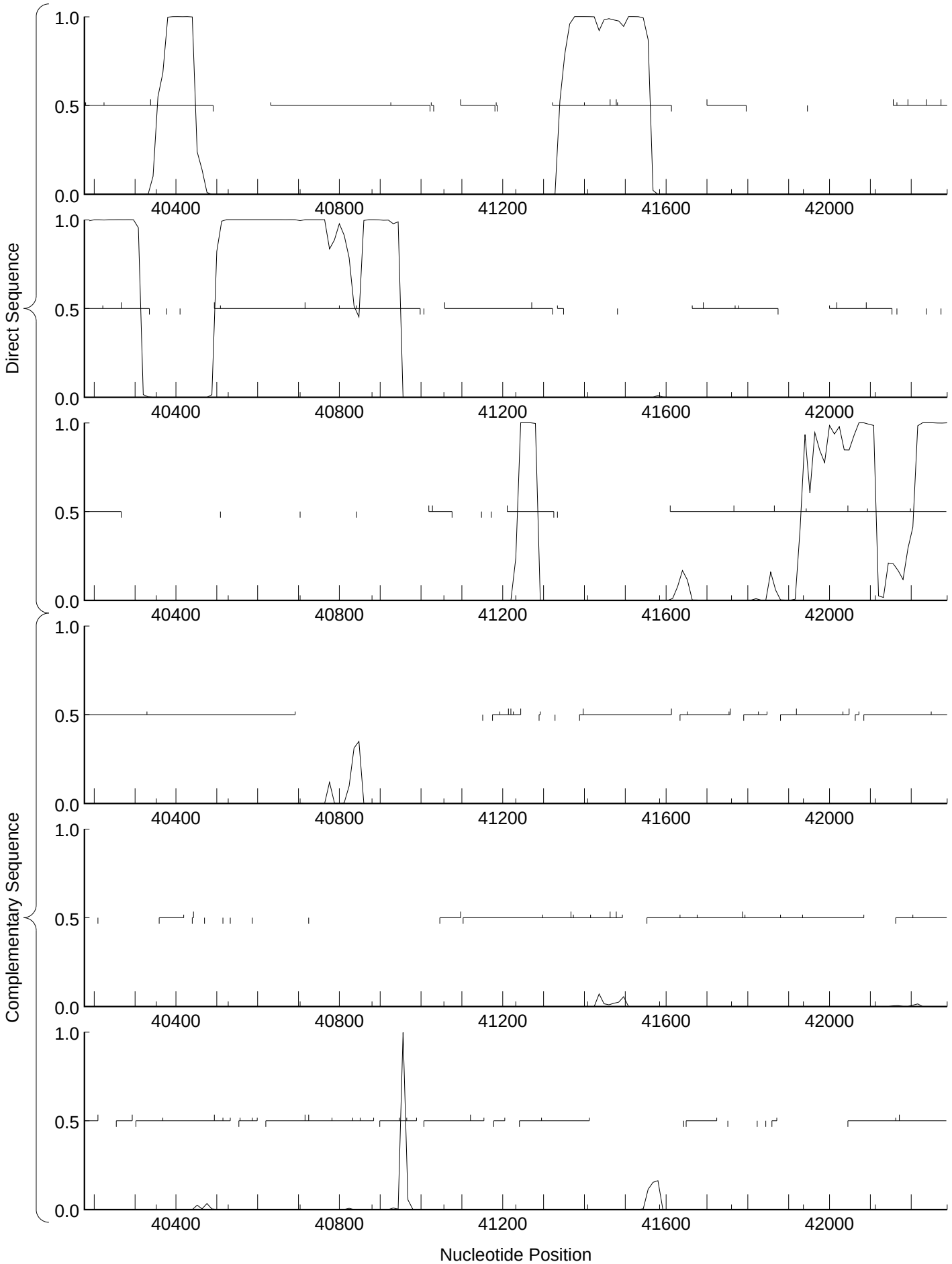
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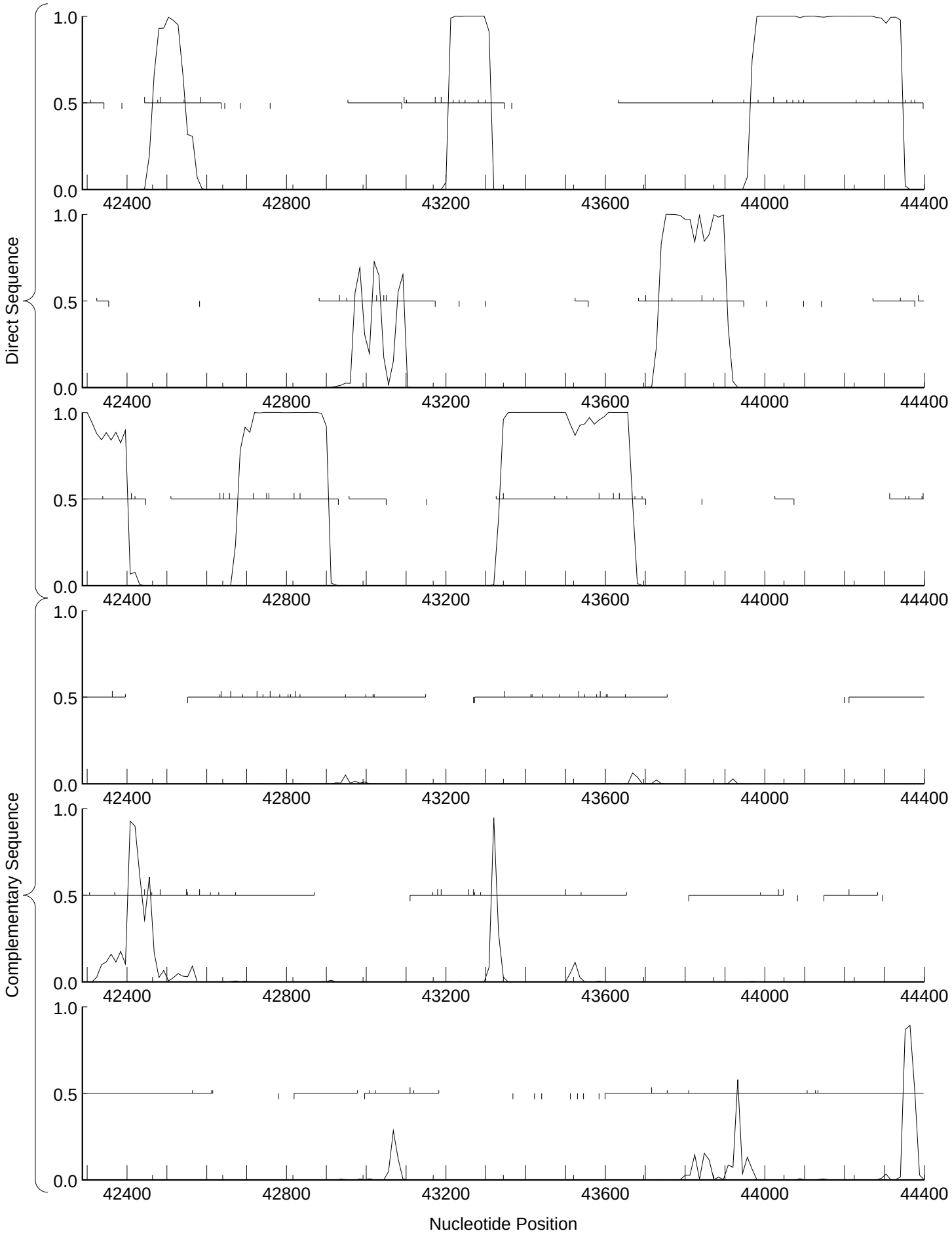
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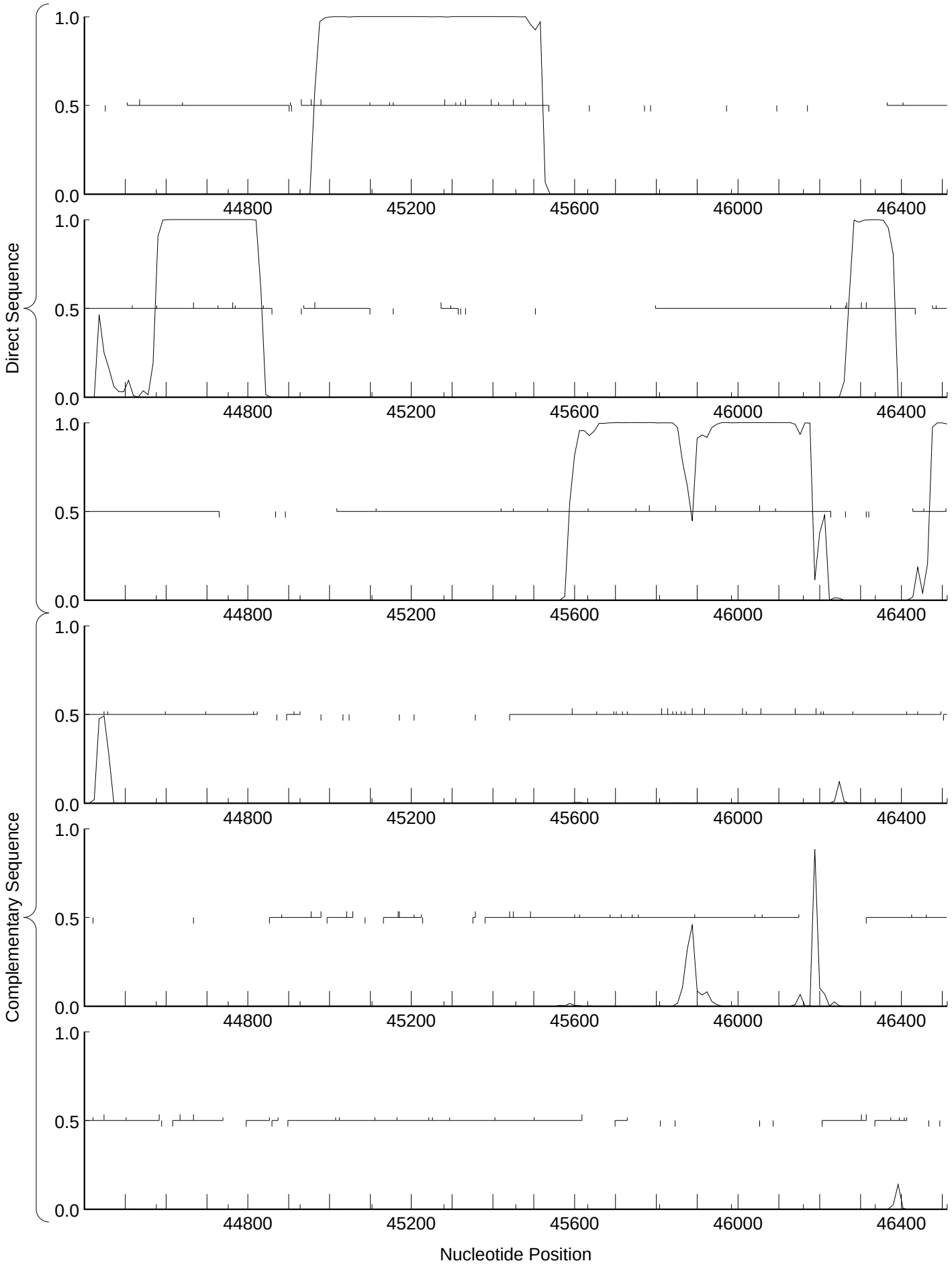
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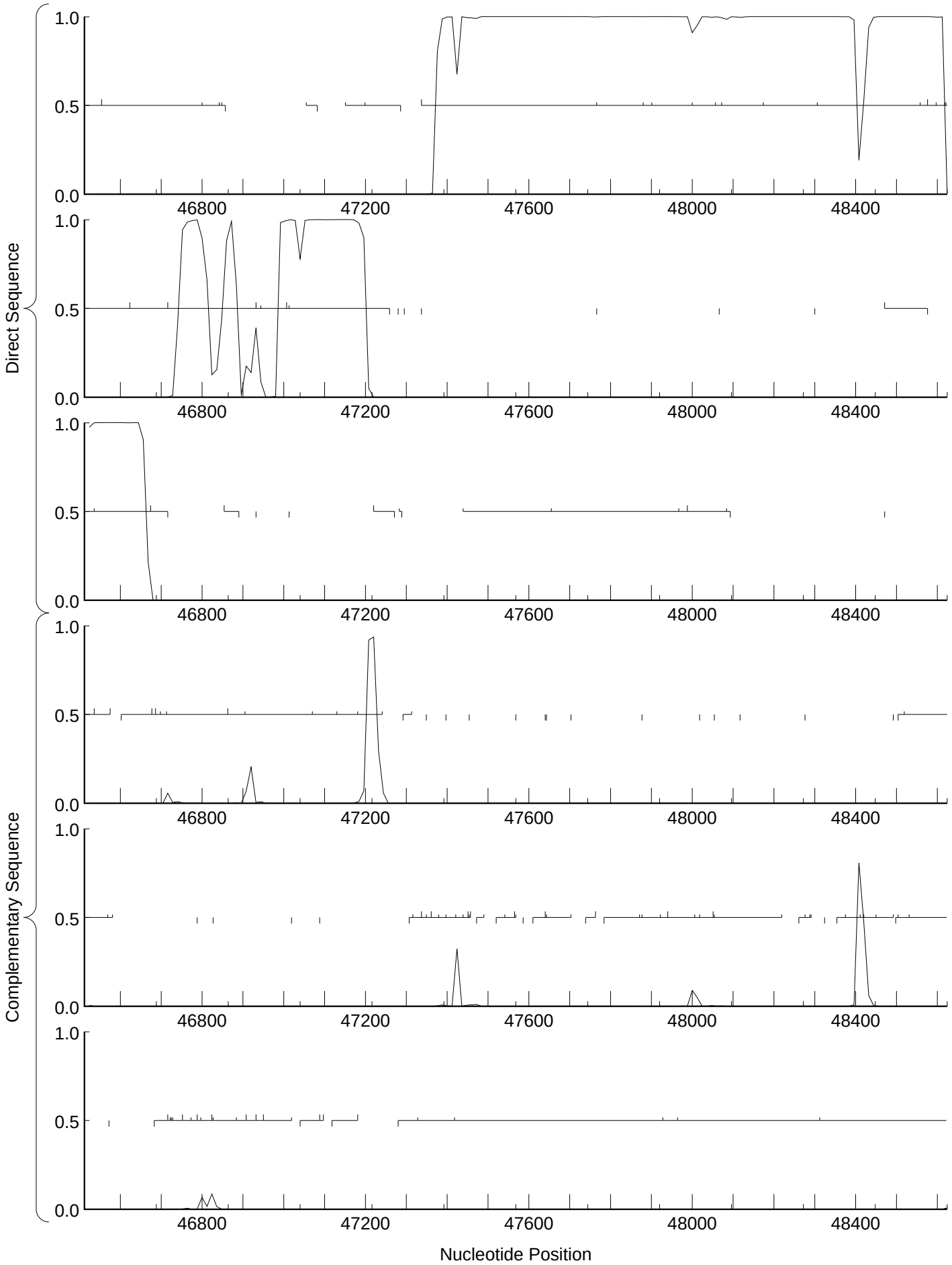
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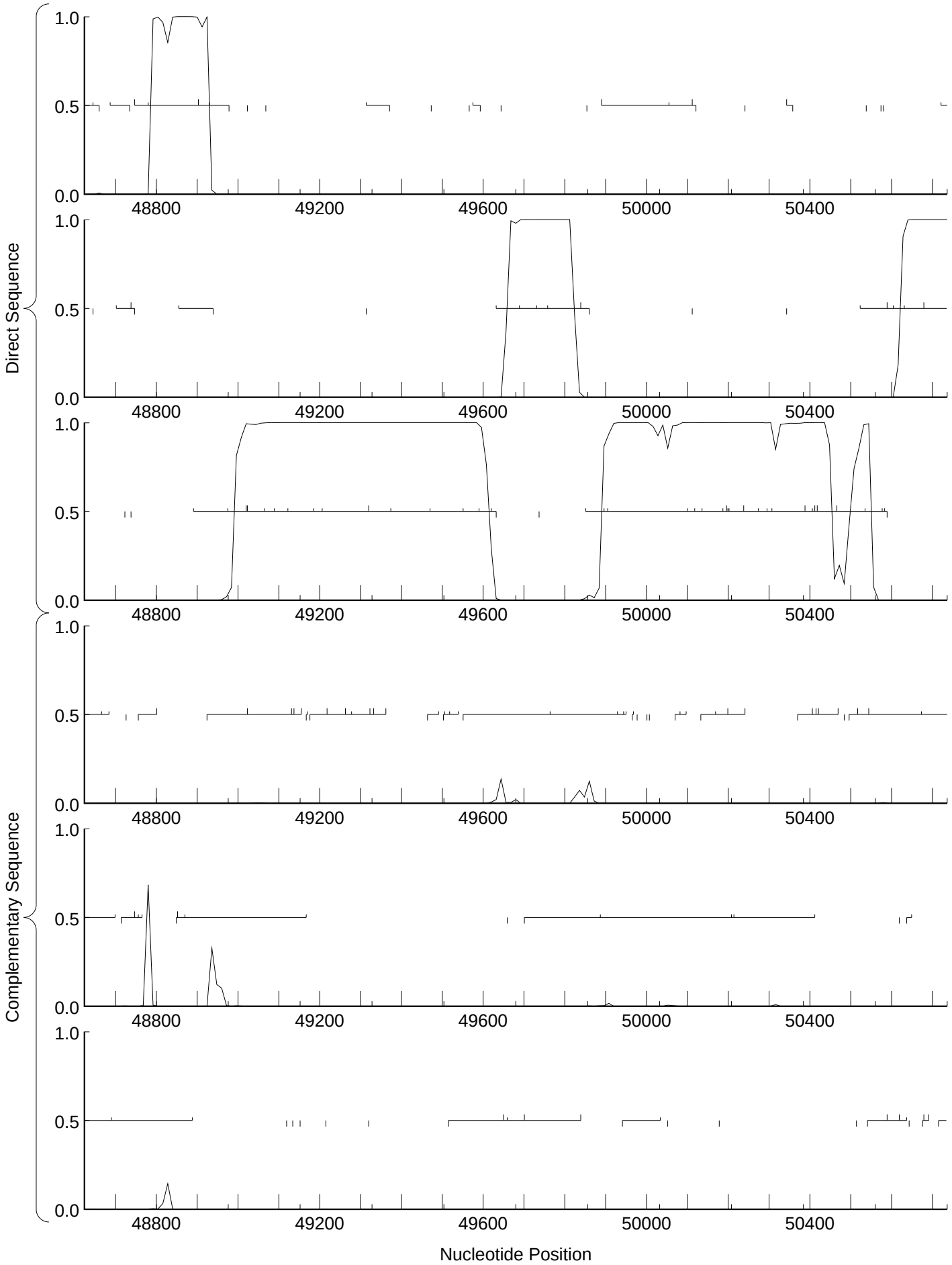
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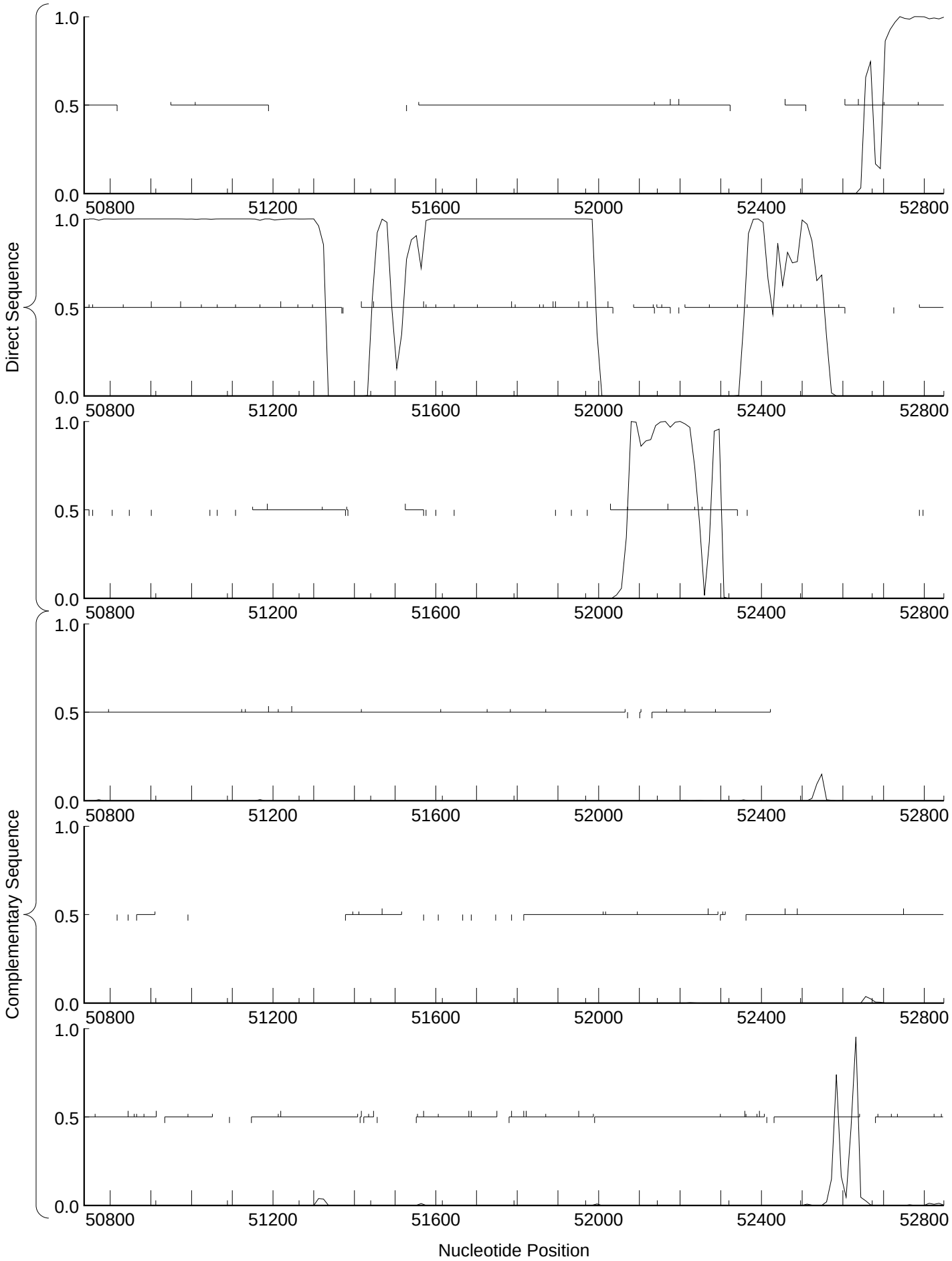
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cterium phage KaiHaiDragon complete sequence, 52992 bp, circularly permuted, Cluster EC, Order 4, Window 96, Step 12, 25/27



cterium phage KaiHaiDragon complete sequence, 52992 bp, circularly permuted, Cluster EC, Order 4, Window 96, Step 12, 26/27



cterium phage KaiHaiDragon complete sequence, 52992 bp, circularly permuted, Cluster EC, Order 4, Window 96, Step 12, 27/27

