GeneMark

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

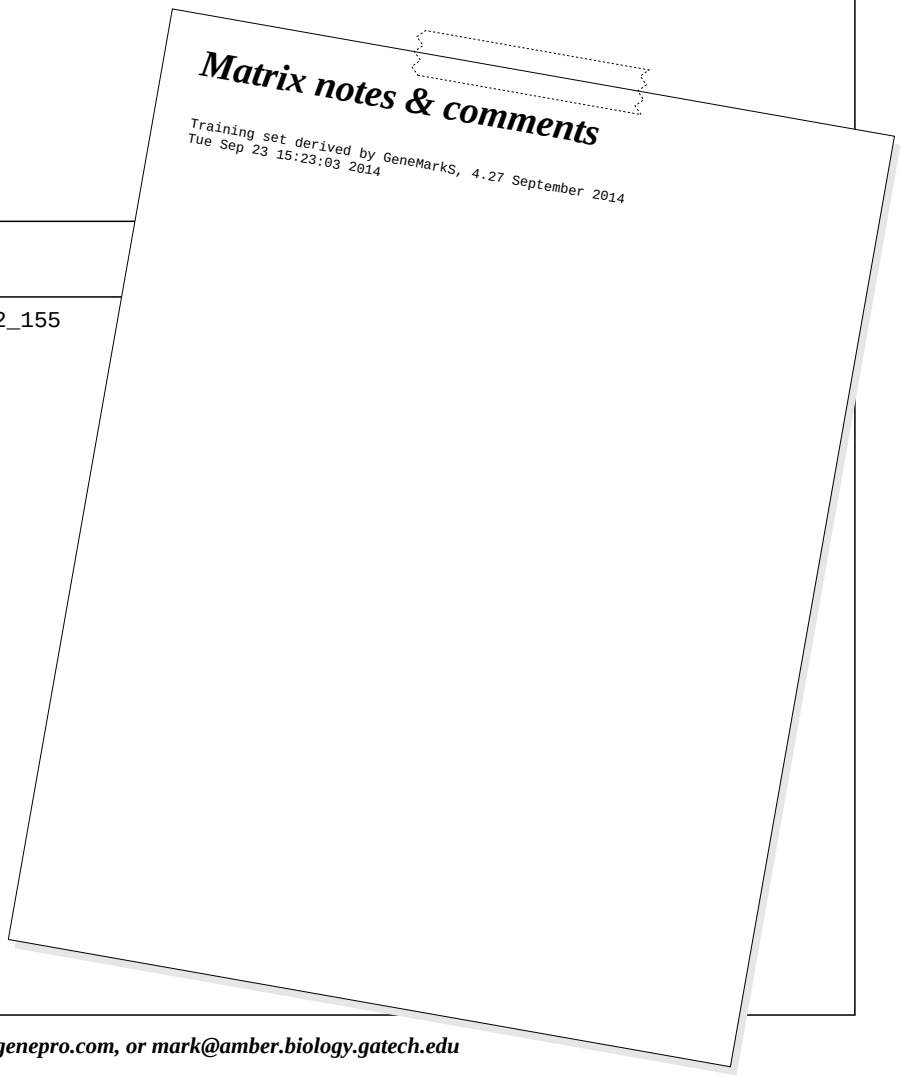
PROGRAM INFORMATION

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Sequence         :  Mycobacterium phage Pinkman complete sequence, 68938 bp, circularly permuted, Cluster B1
Analysis Date    :  3/28/24 at 7:49:54
Pages            :  34
Sequence Length  :  68938 bp
GC Content       :   66.37%

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

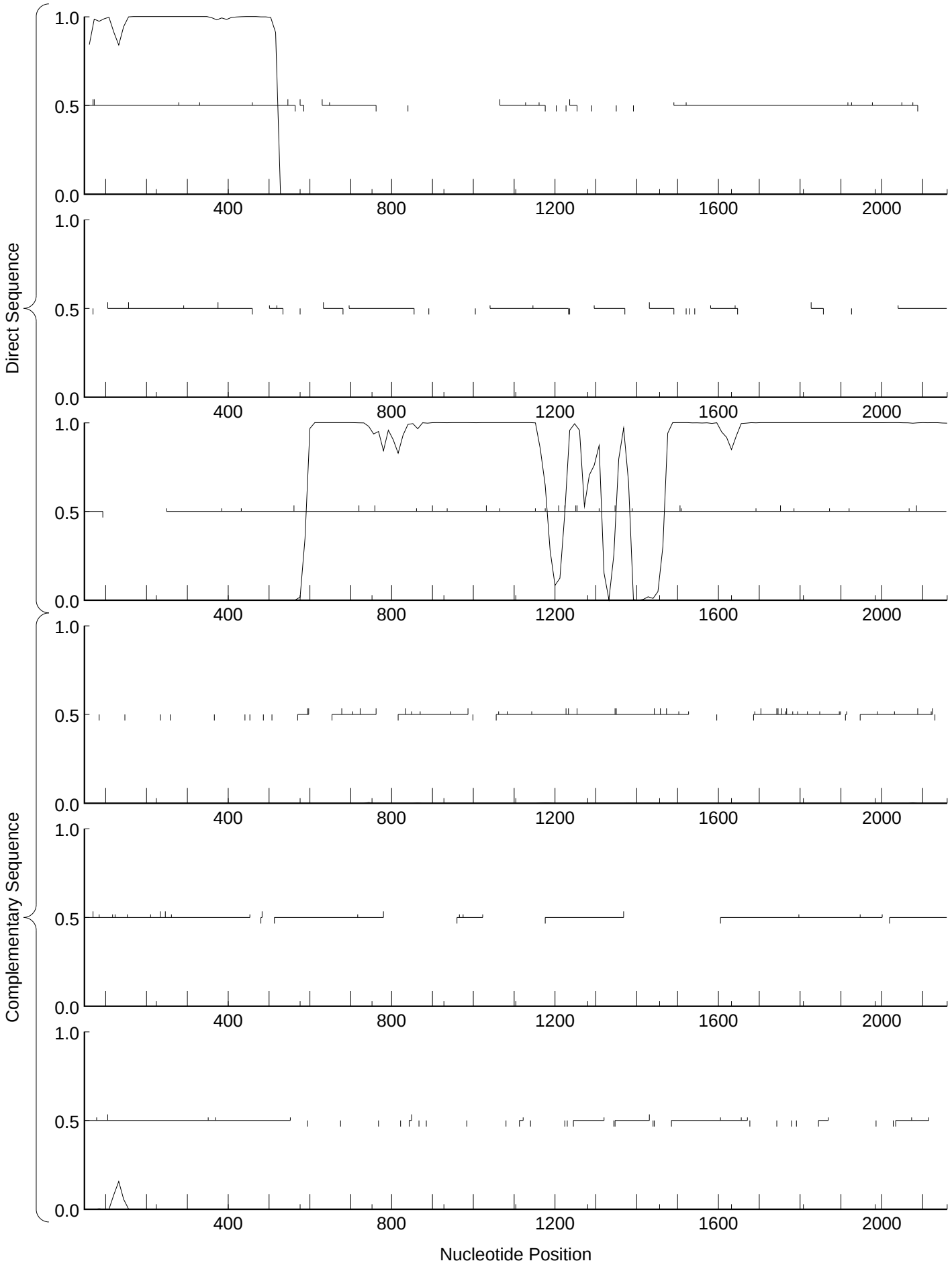
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Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

Order      :  4
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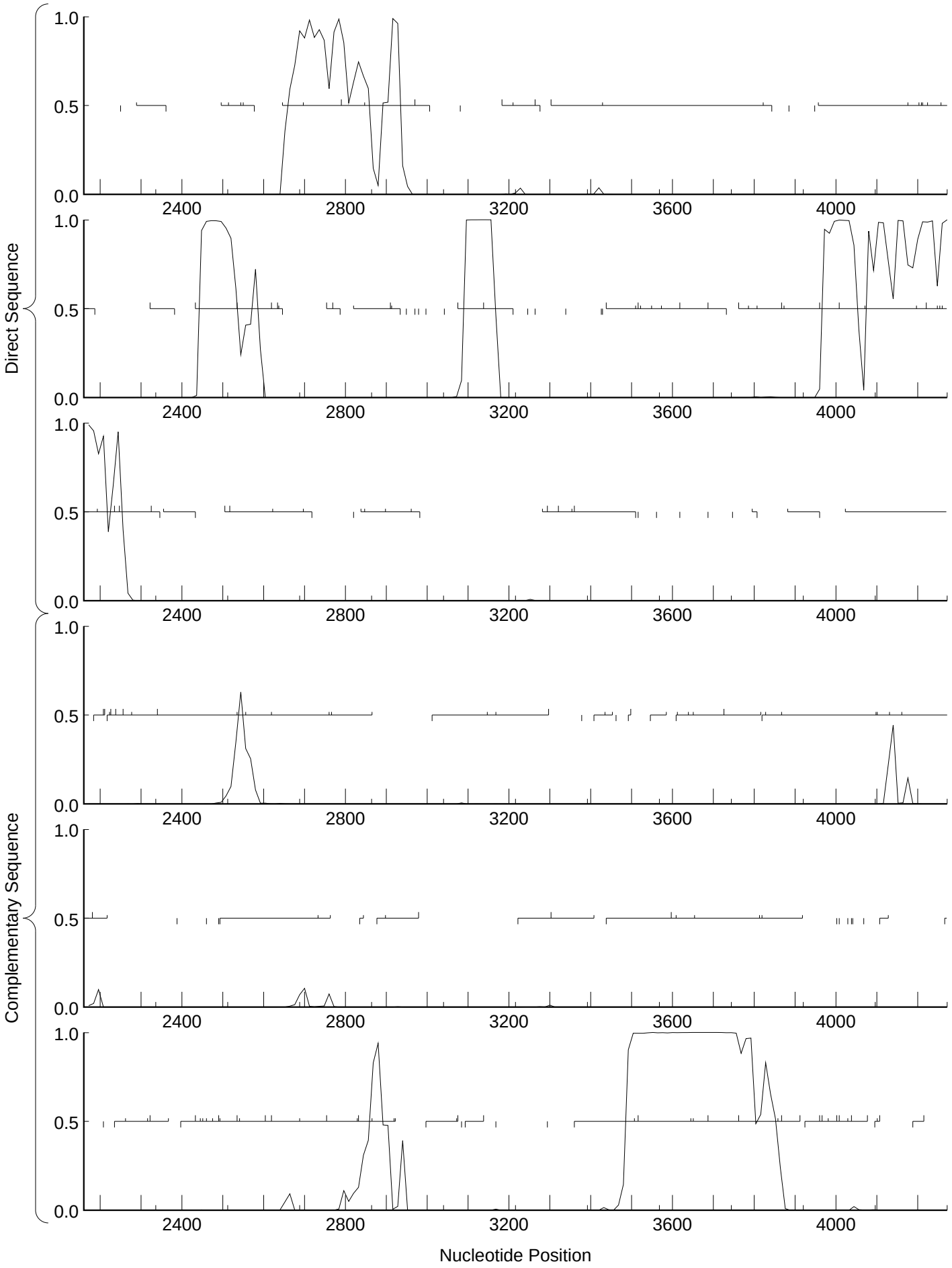

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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

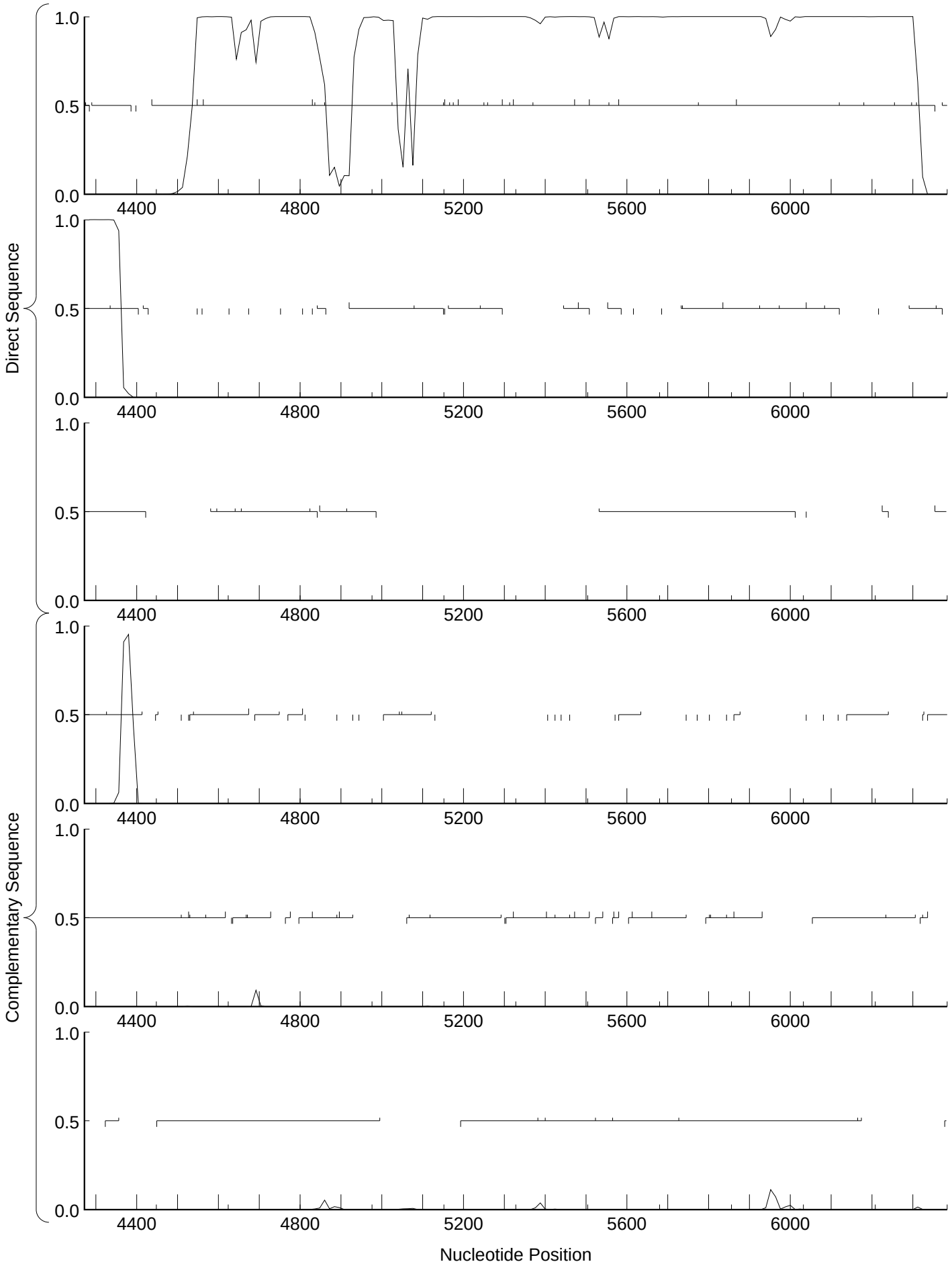

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

Mycobacterium phage Pinkman complete sequence, 68938 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 2/34

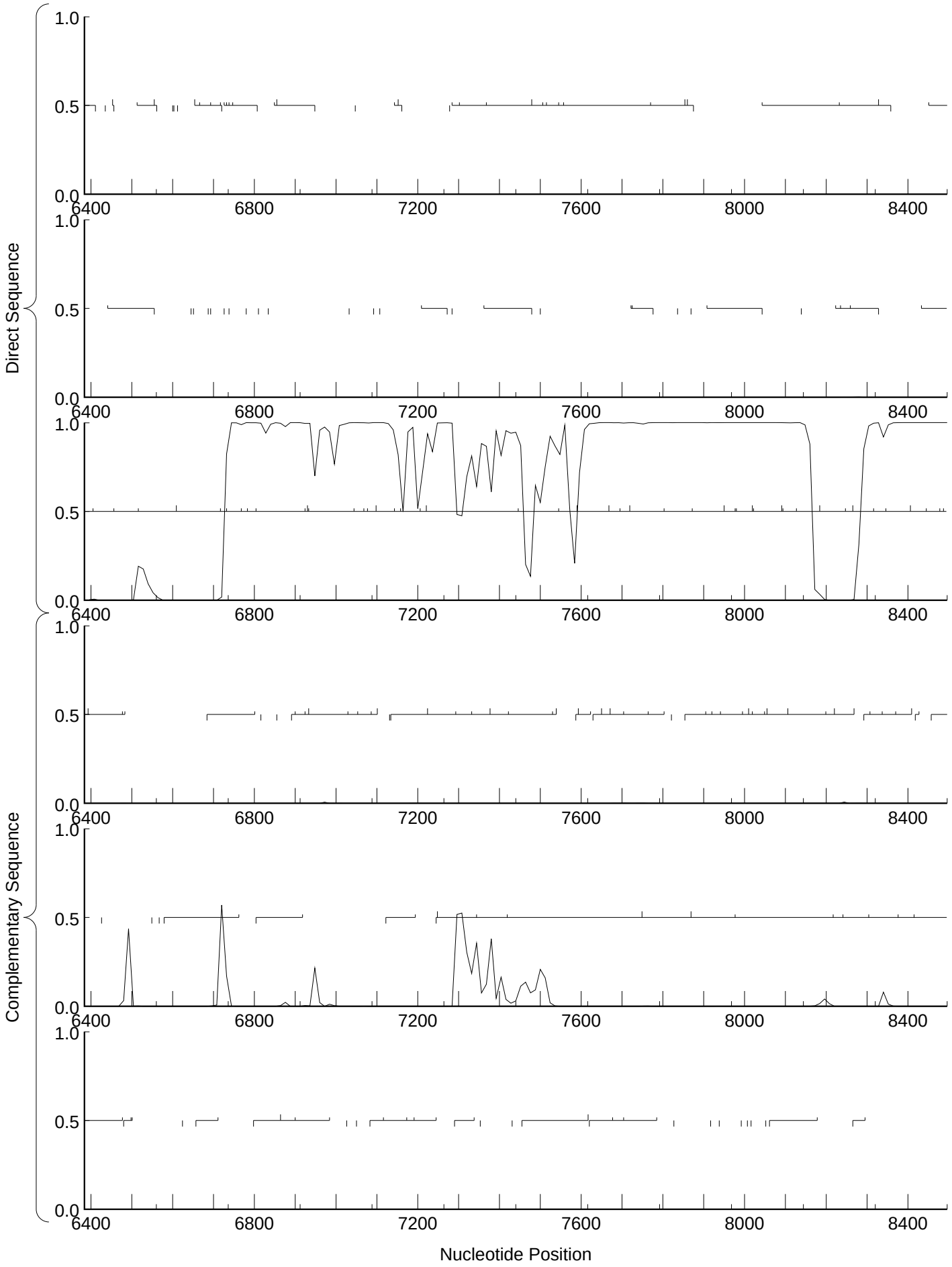


Mycobacterium phage Pinkman complete sequence, 68938 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 3/34

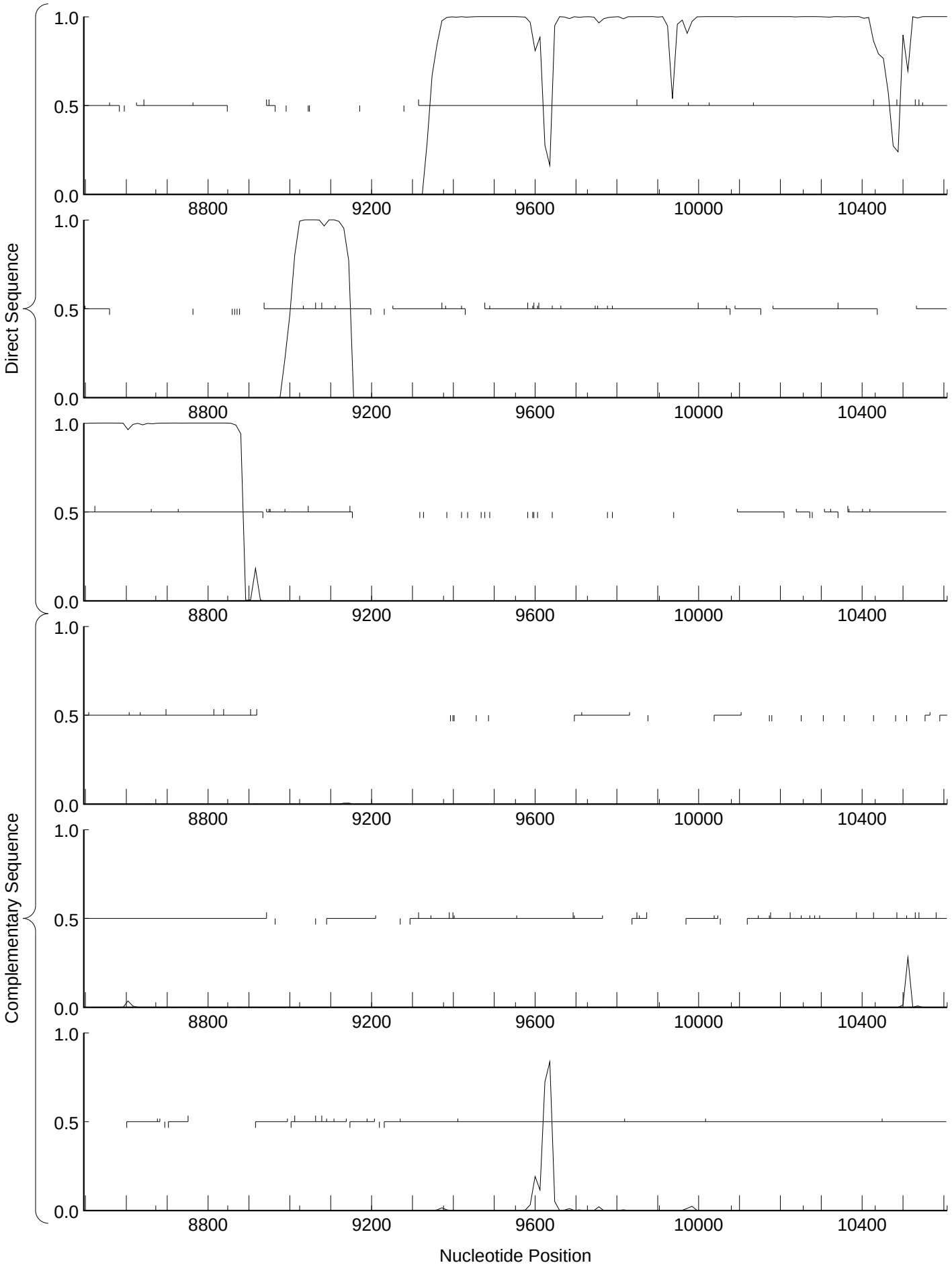



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
4400
4800
5200
5600
6000
Nucleotide Position

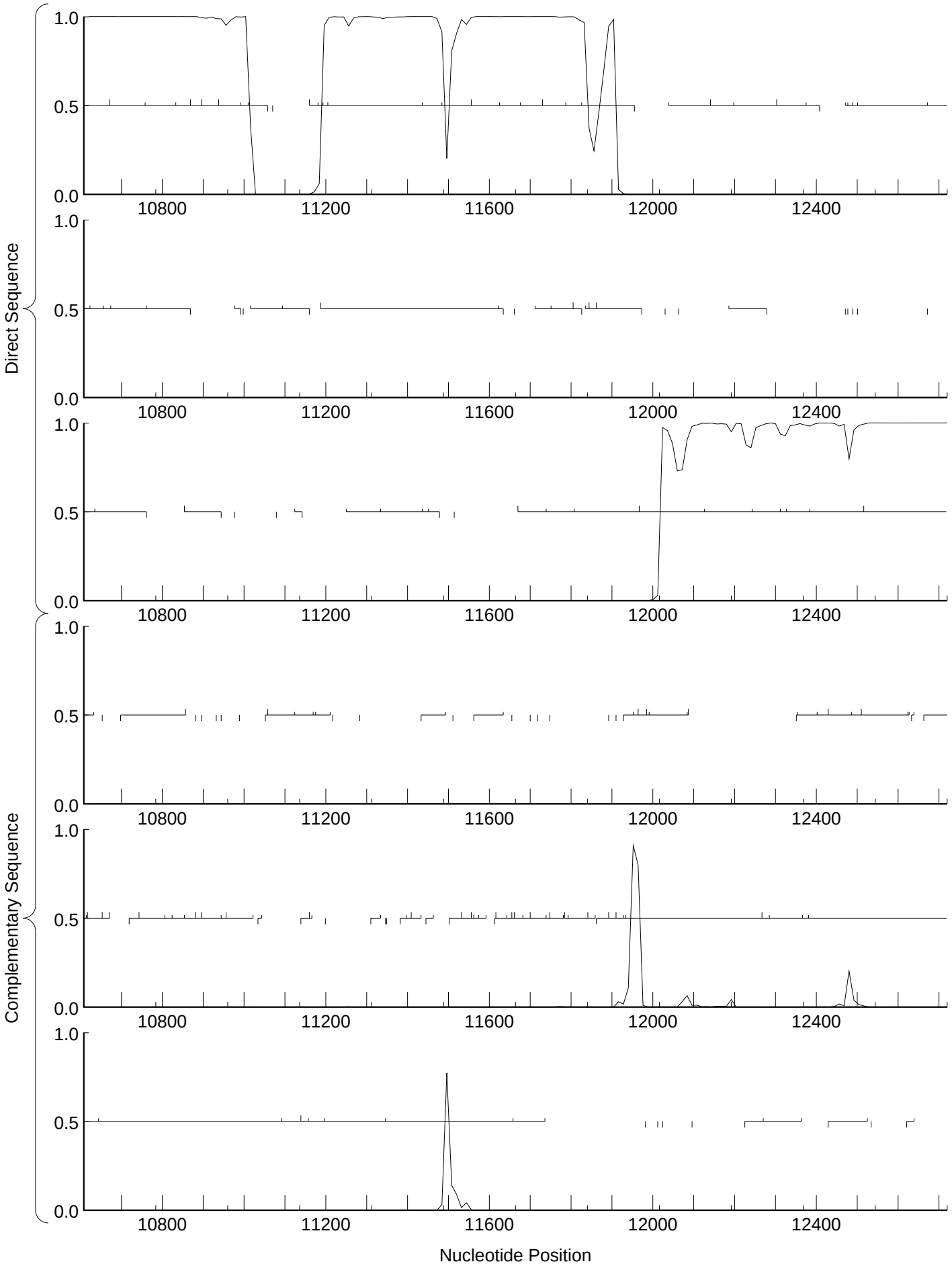
Mycobacterium phage Pinkman complete sequence, 68938 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 5/34



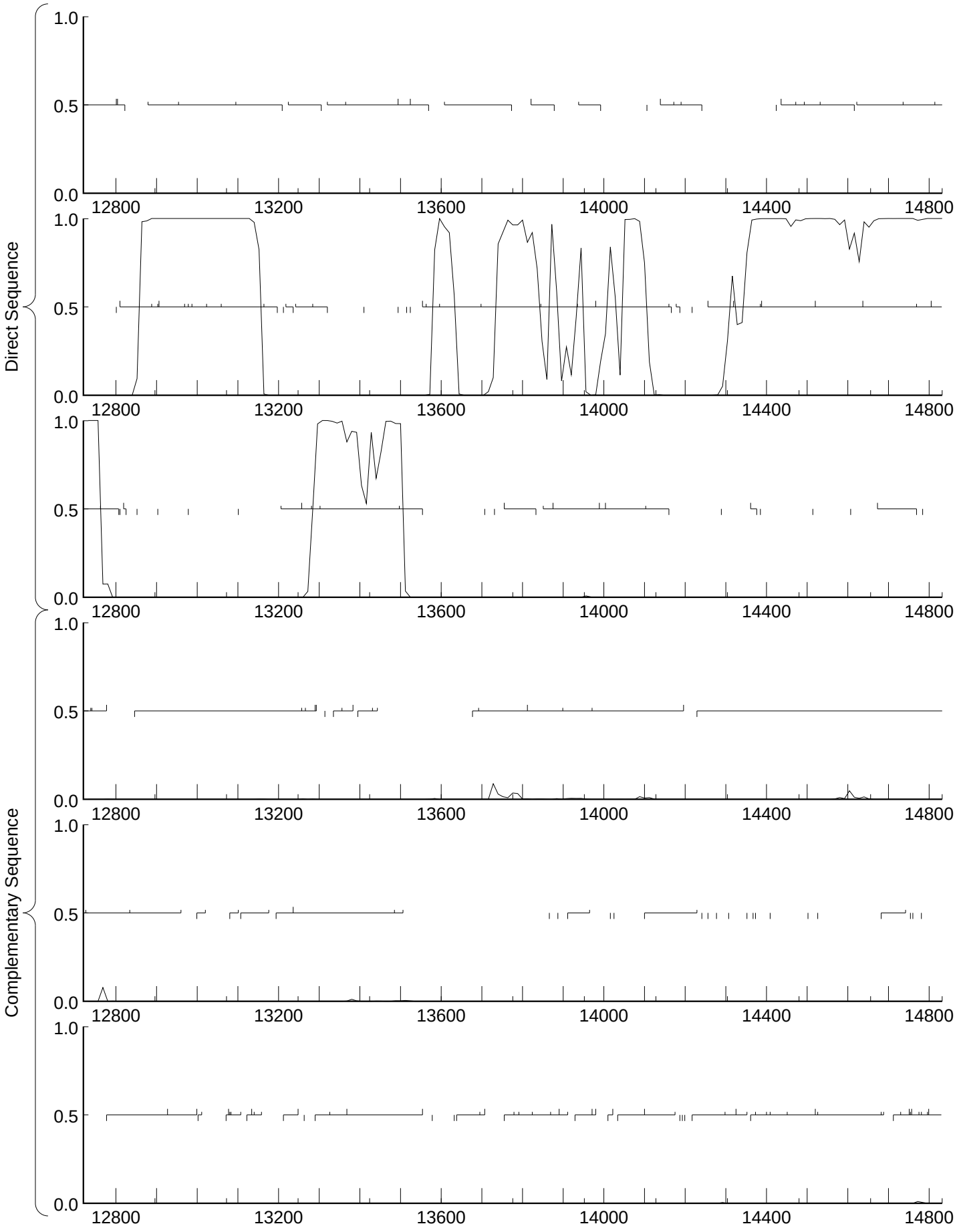
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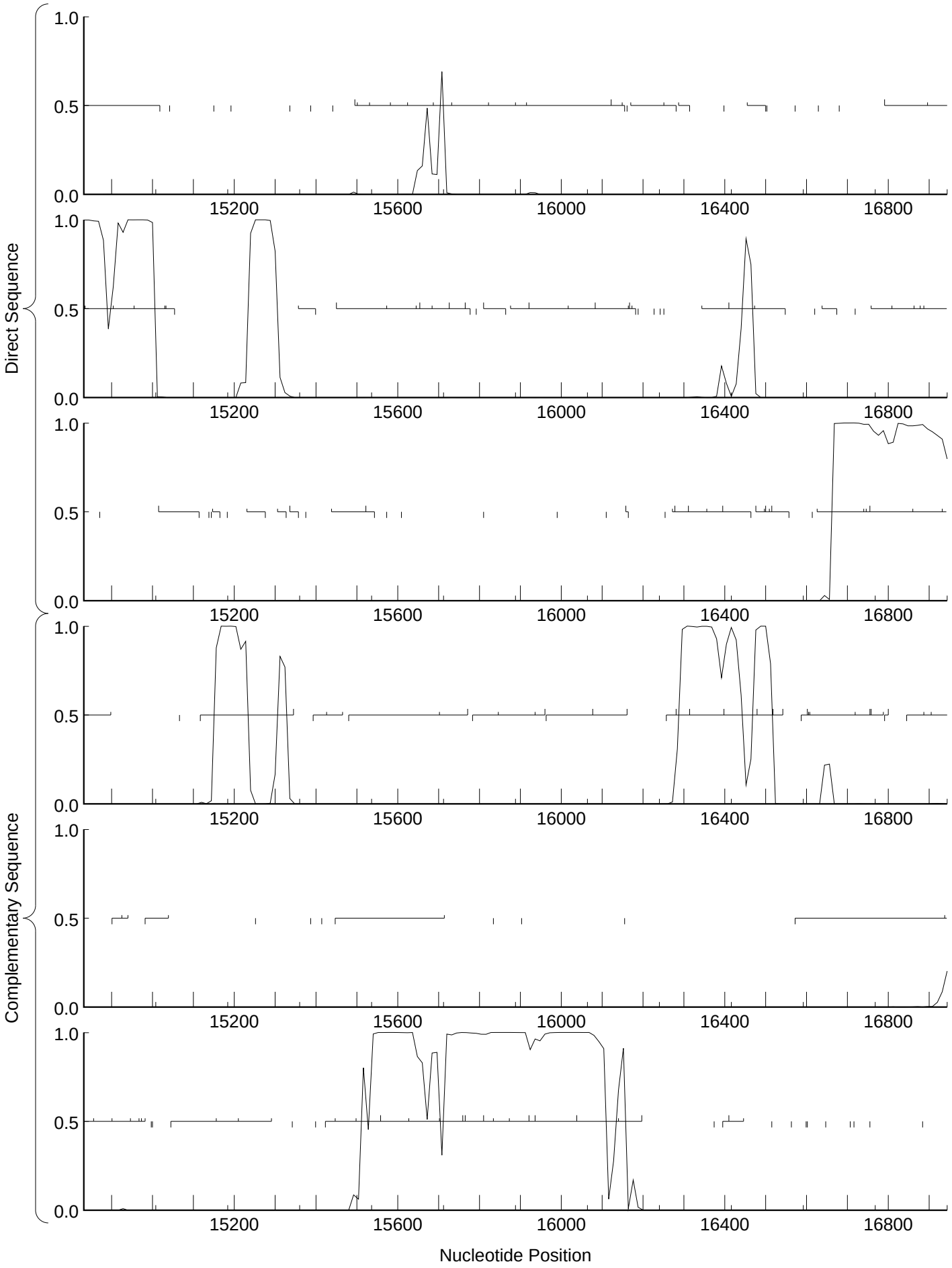
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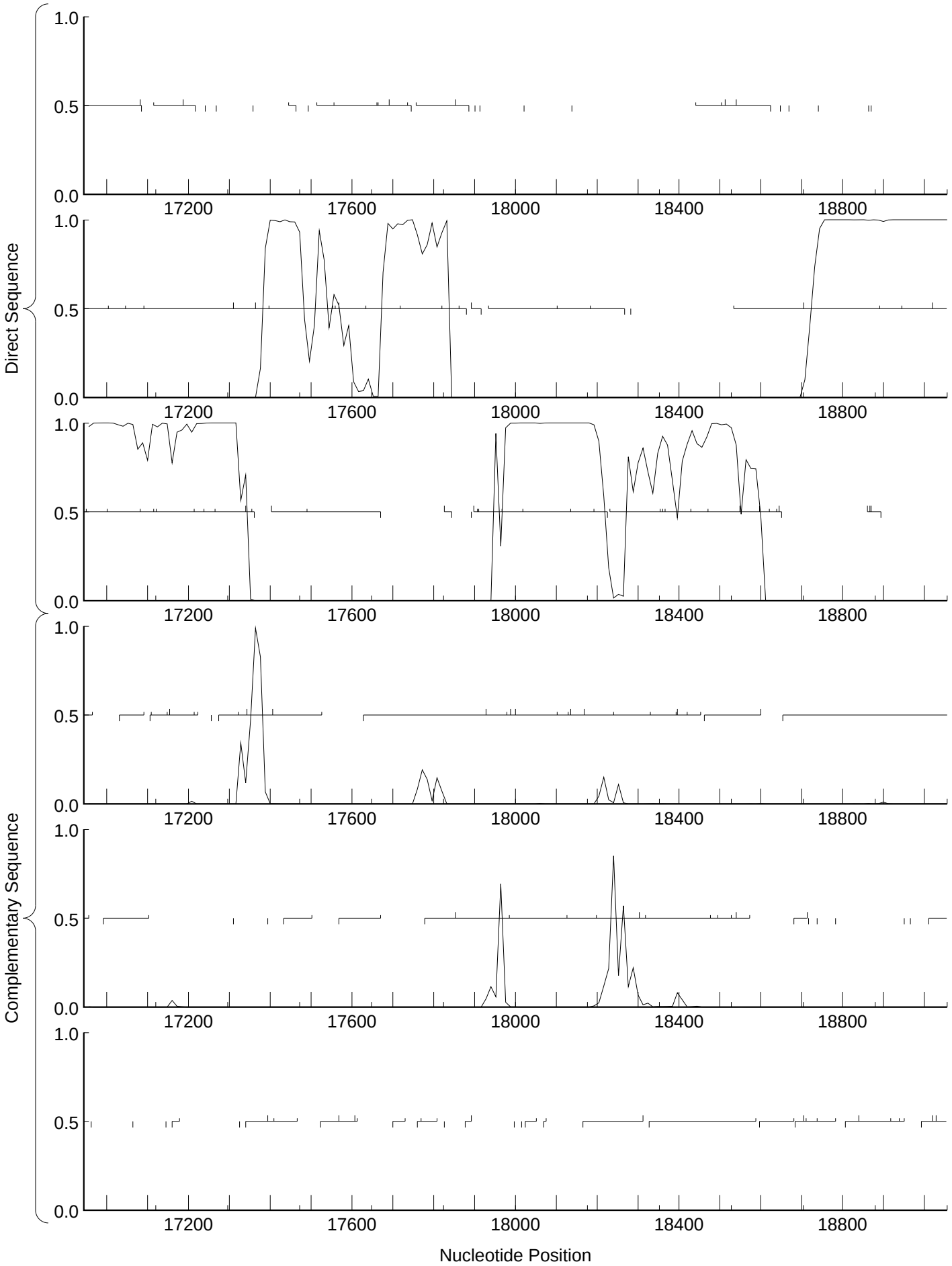
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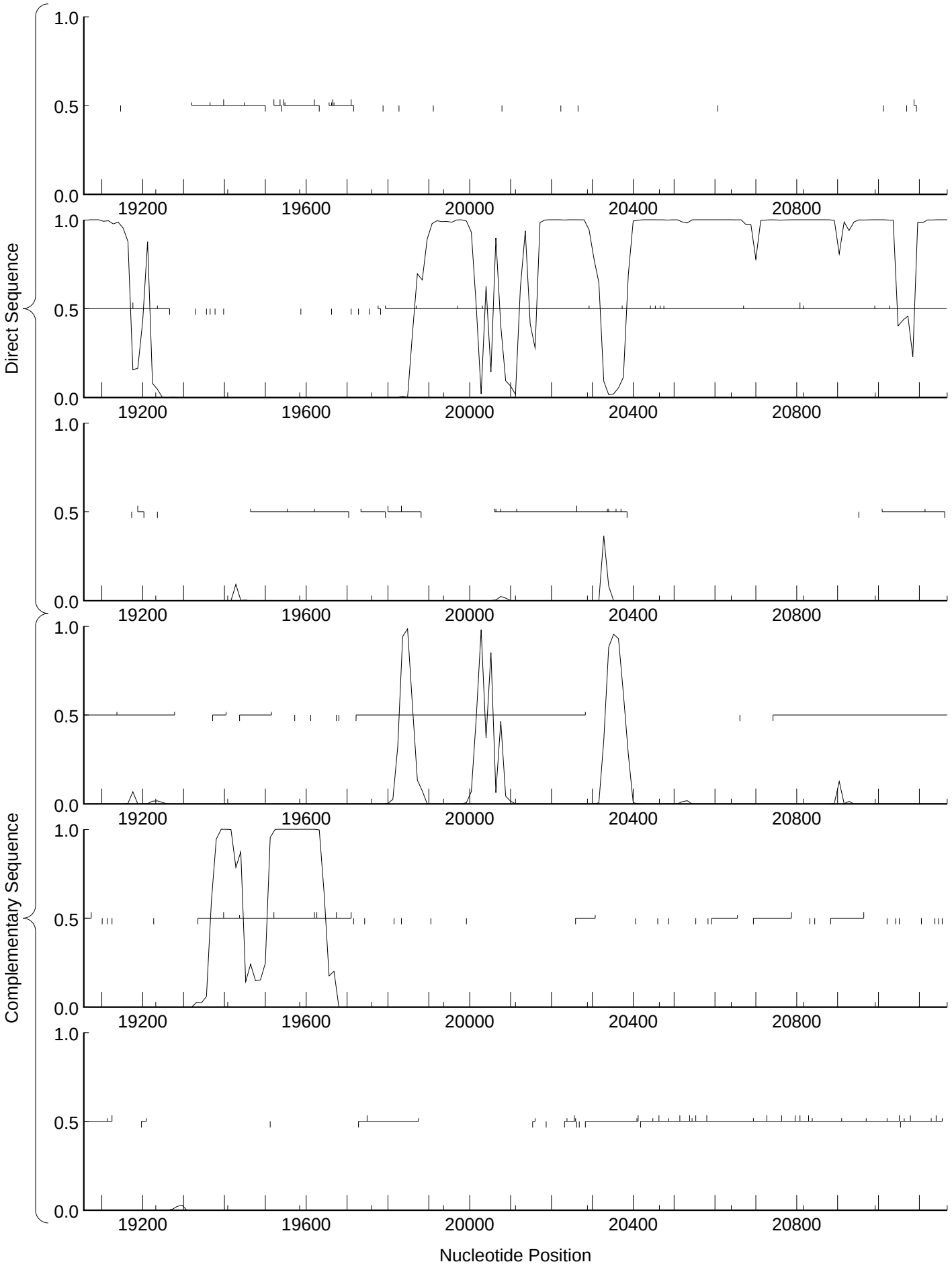
Mycobacterium phage Pinkman complete sequence, 68938 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 9/34



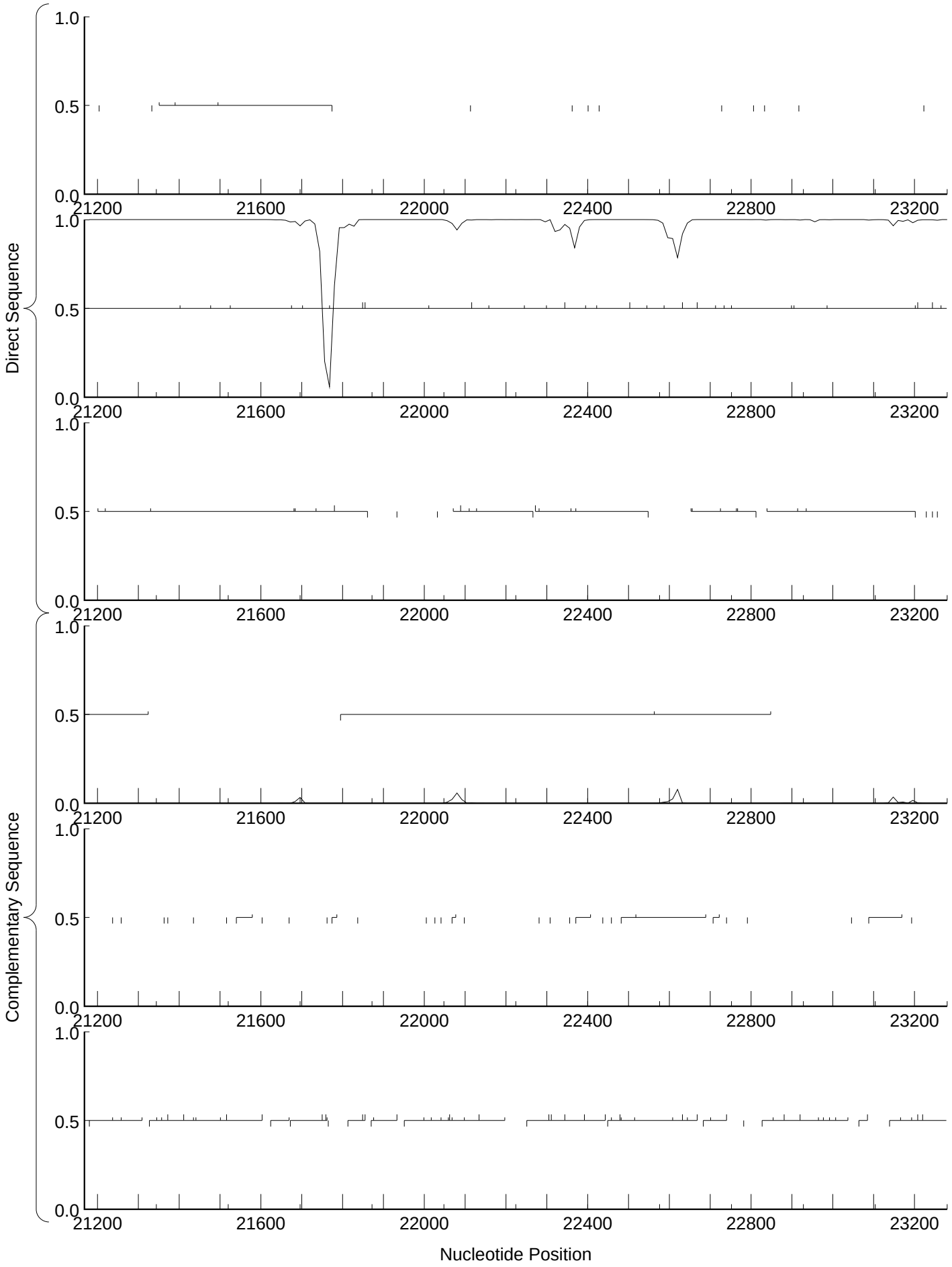
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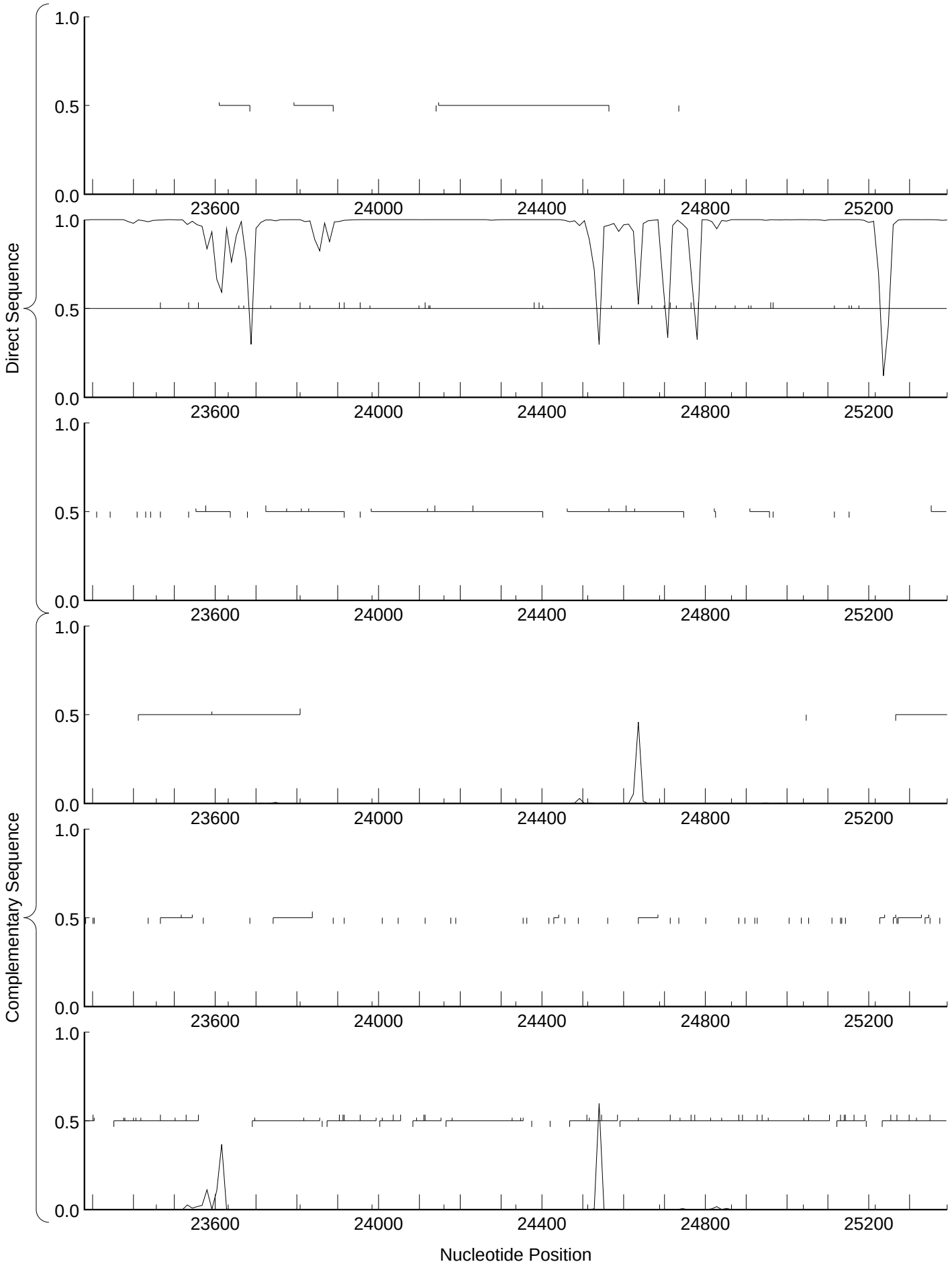
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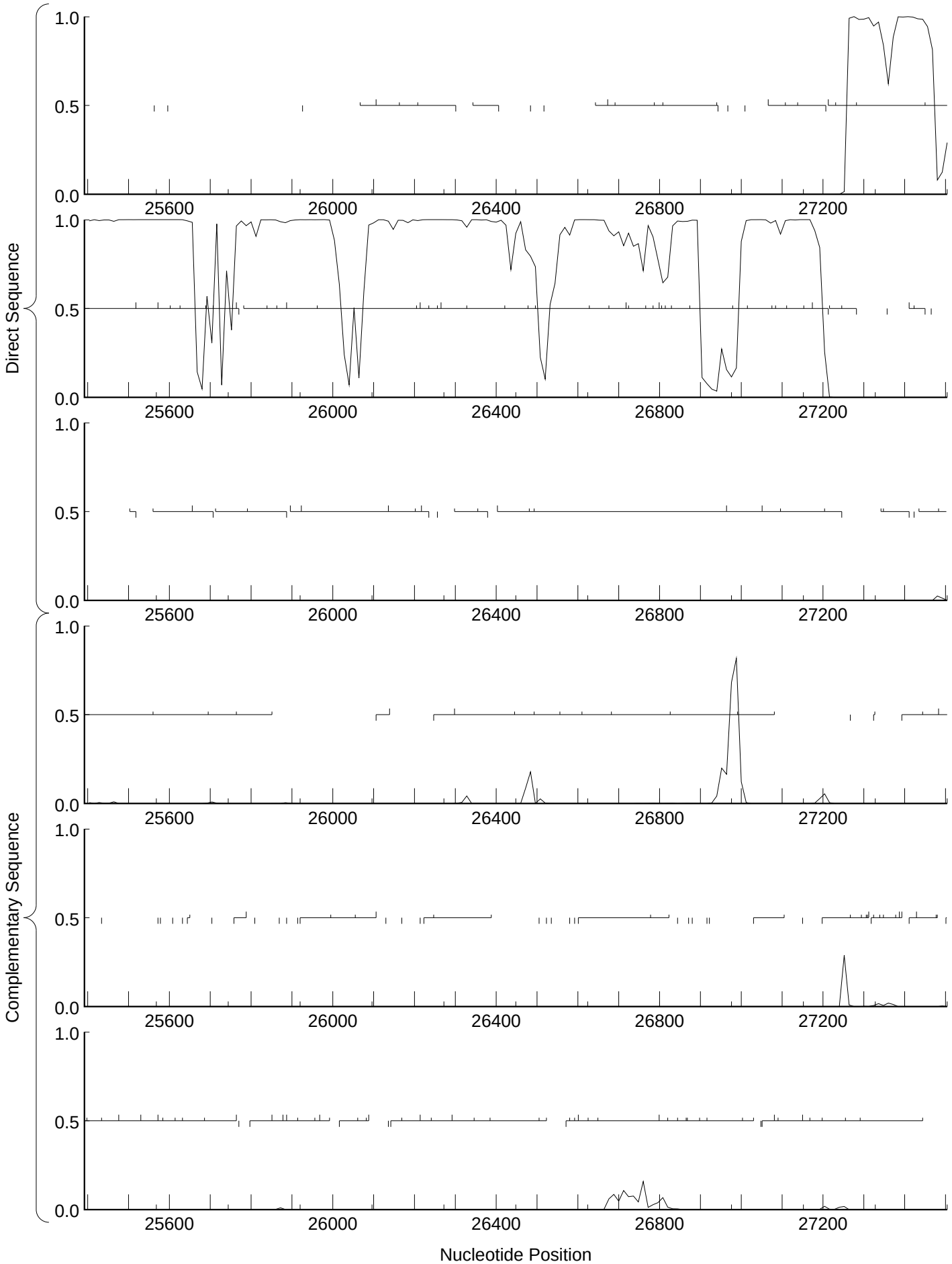
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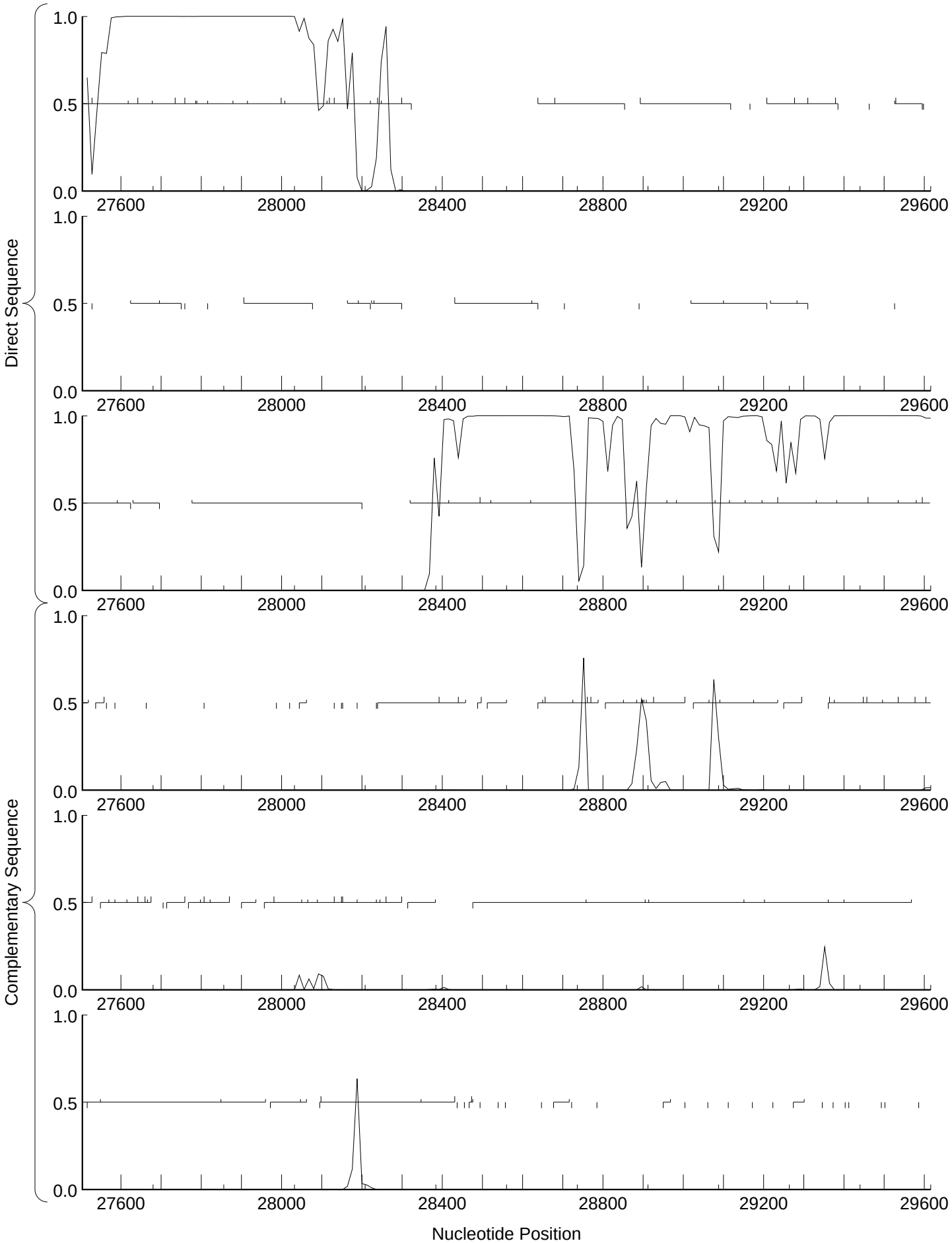
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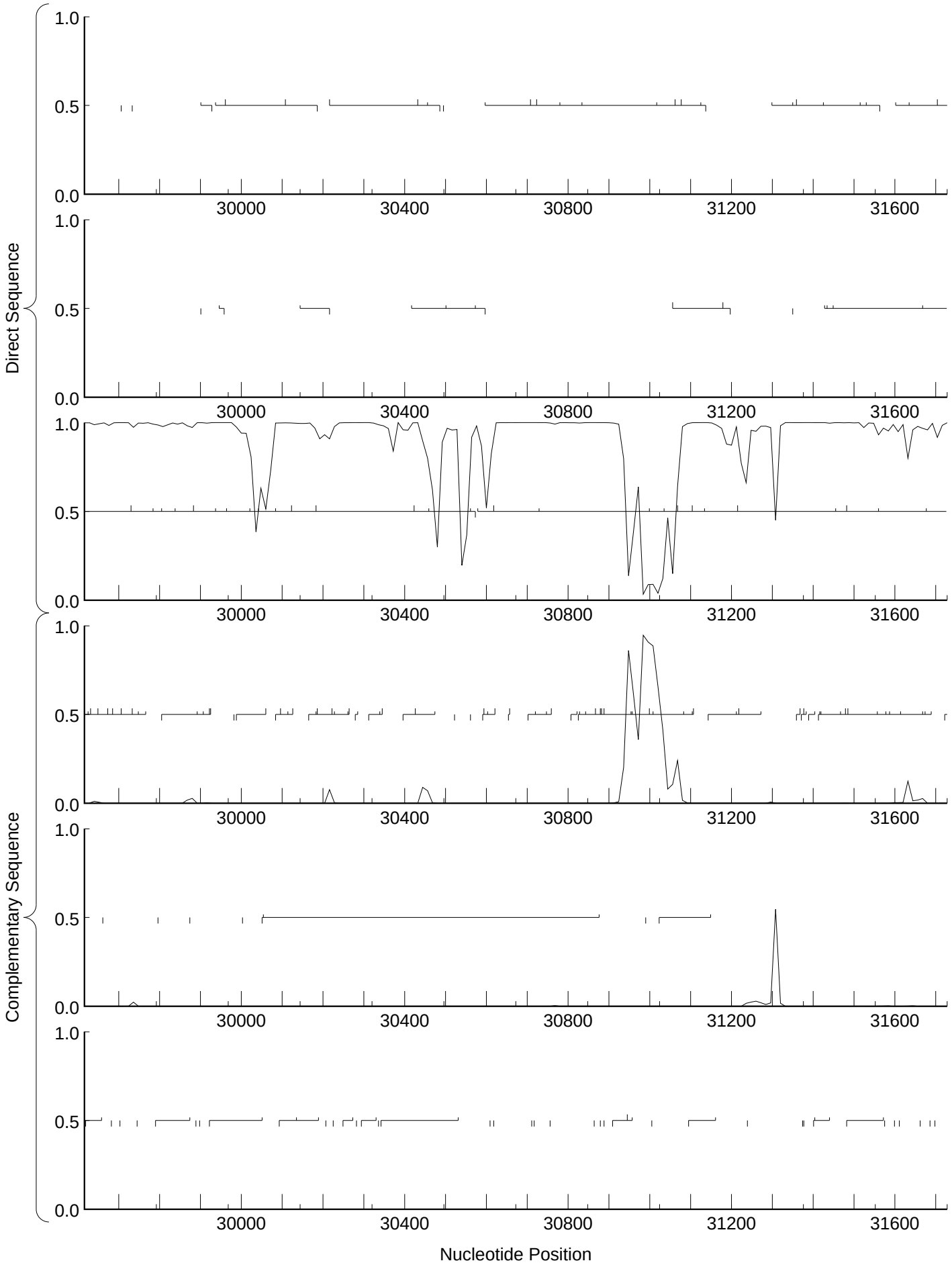
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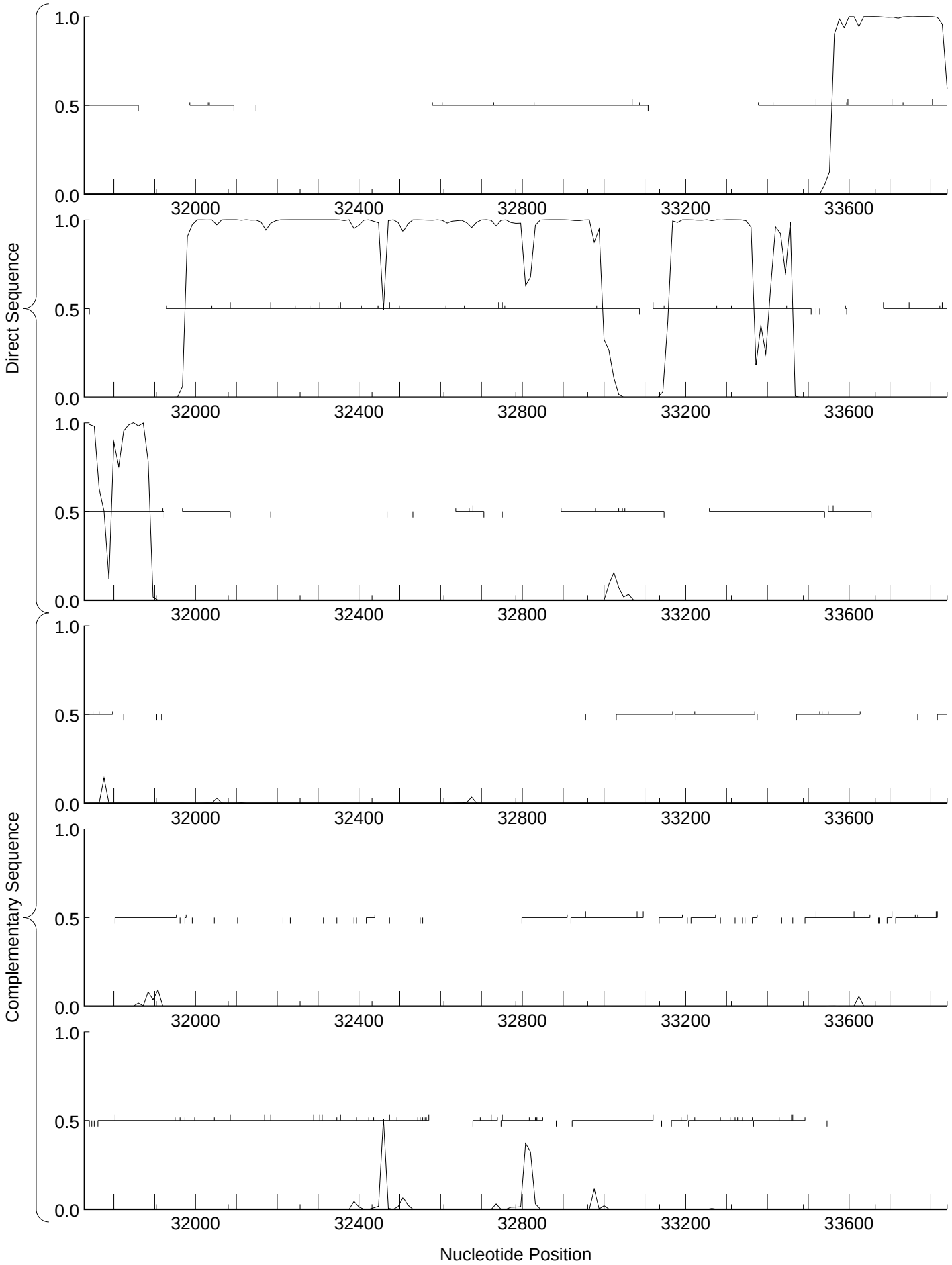
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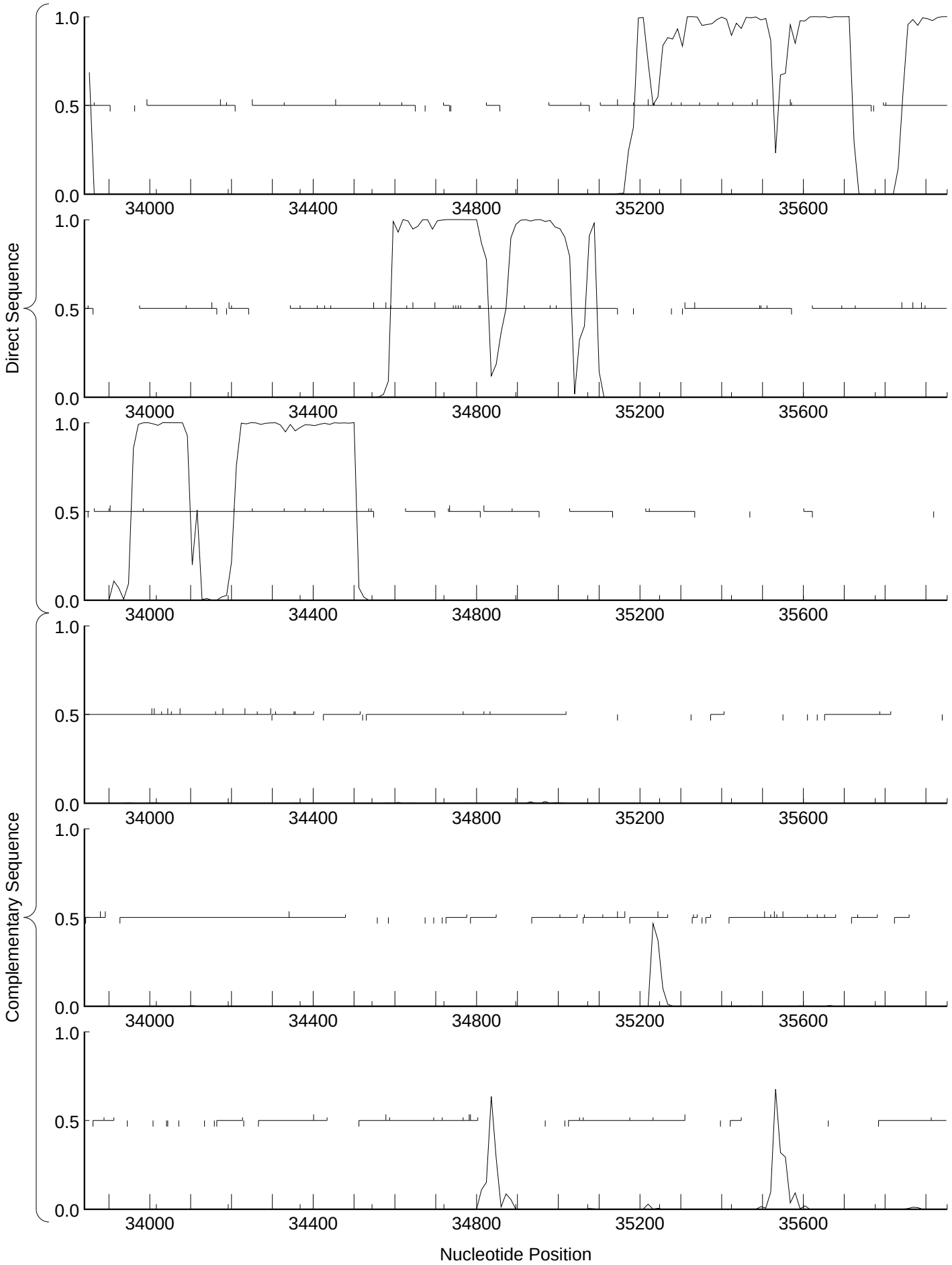
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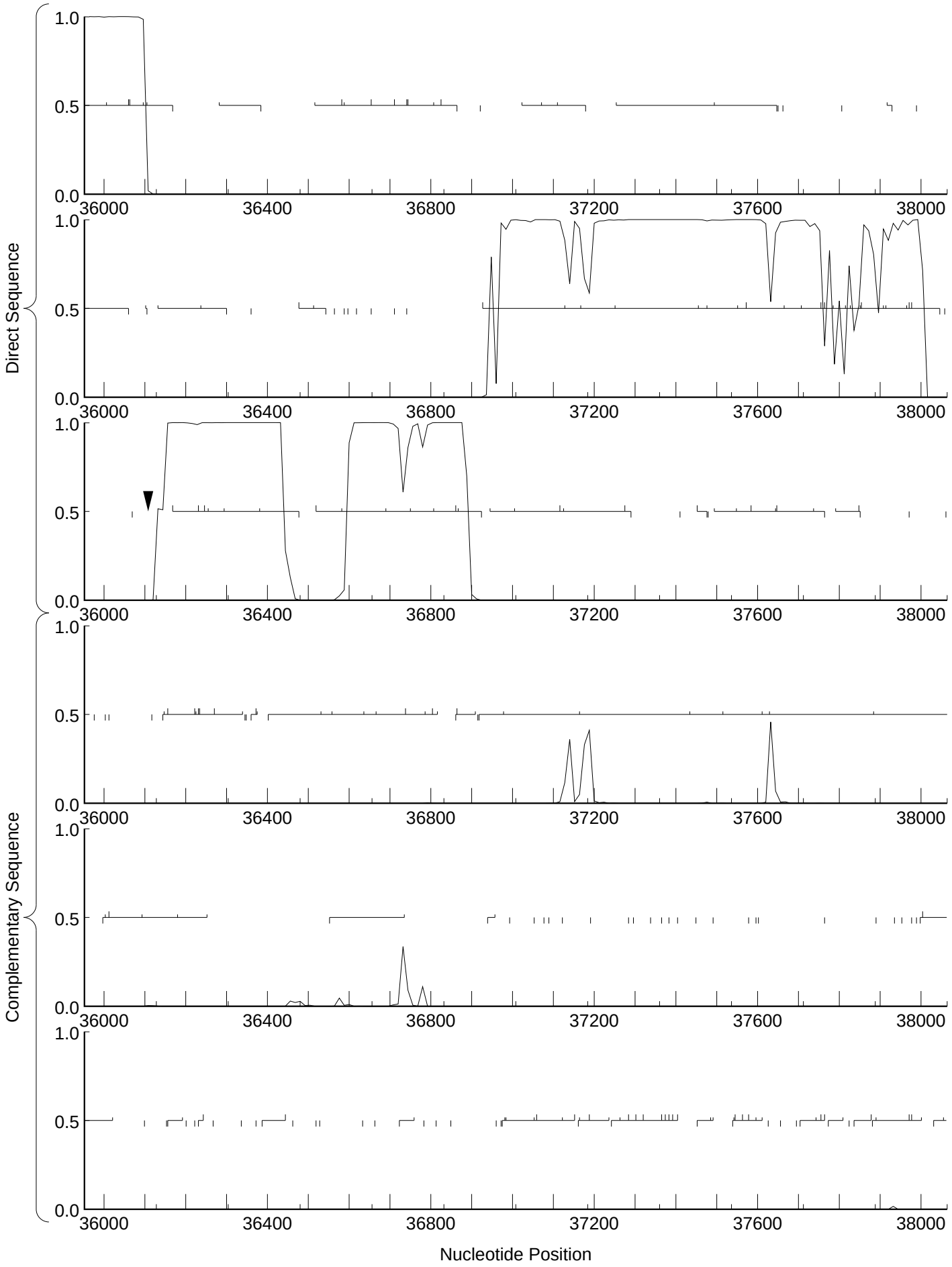
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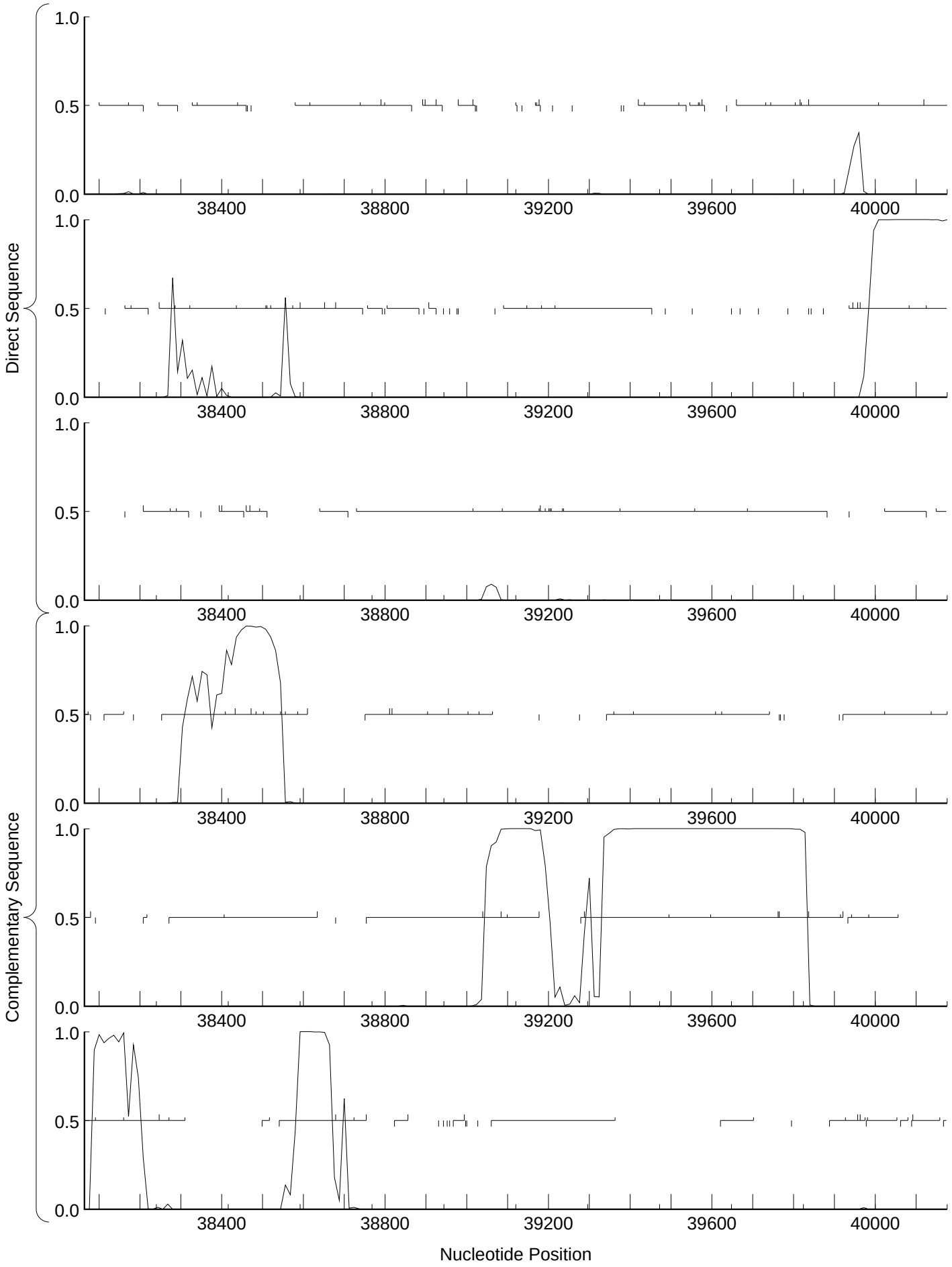
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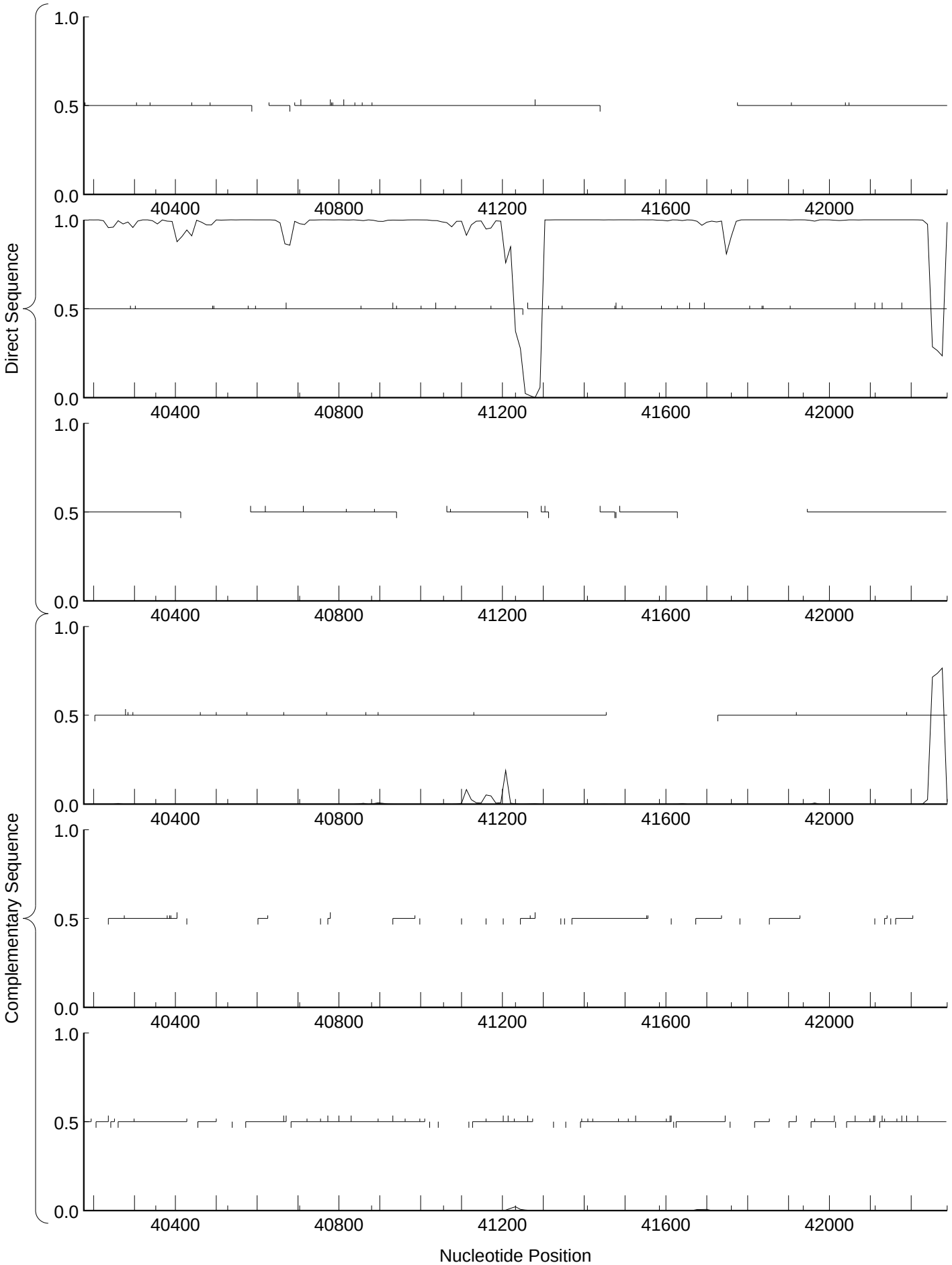
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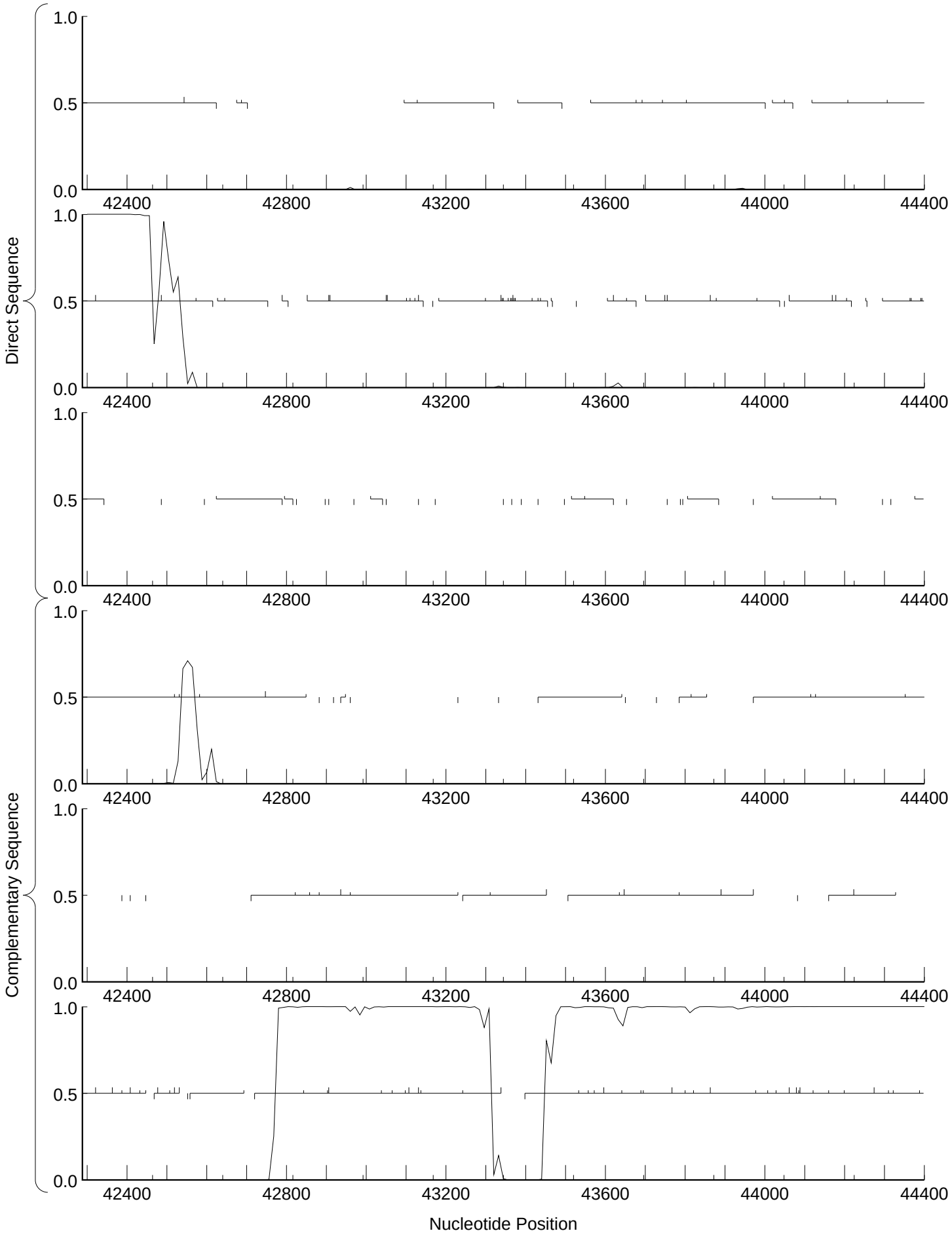
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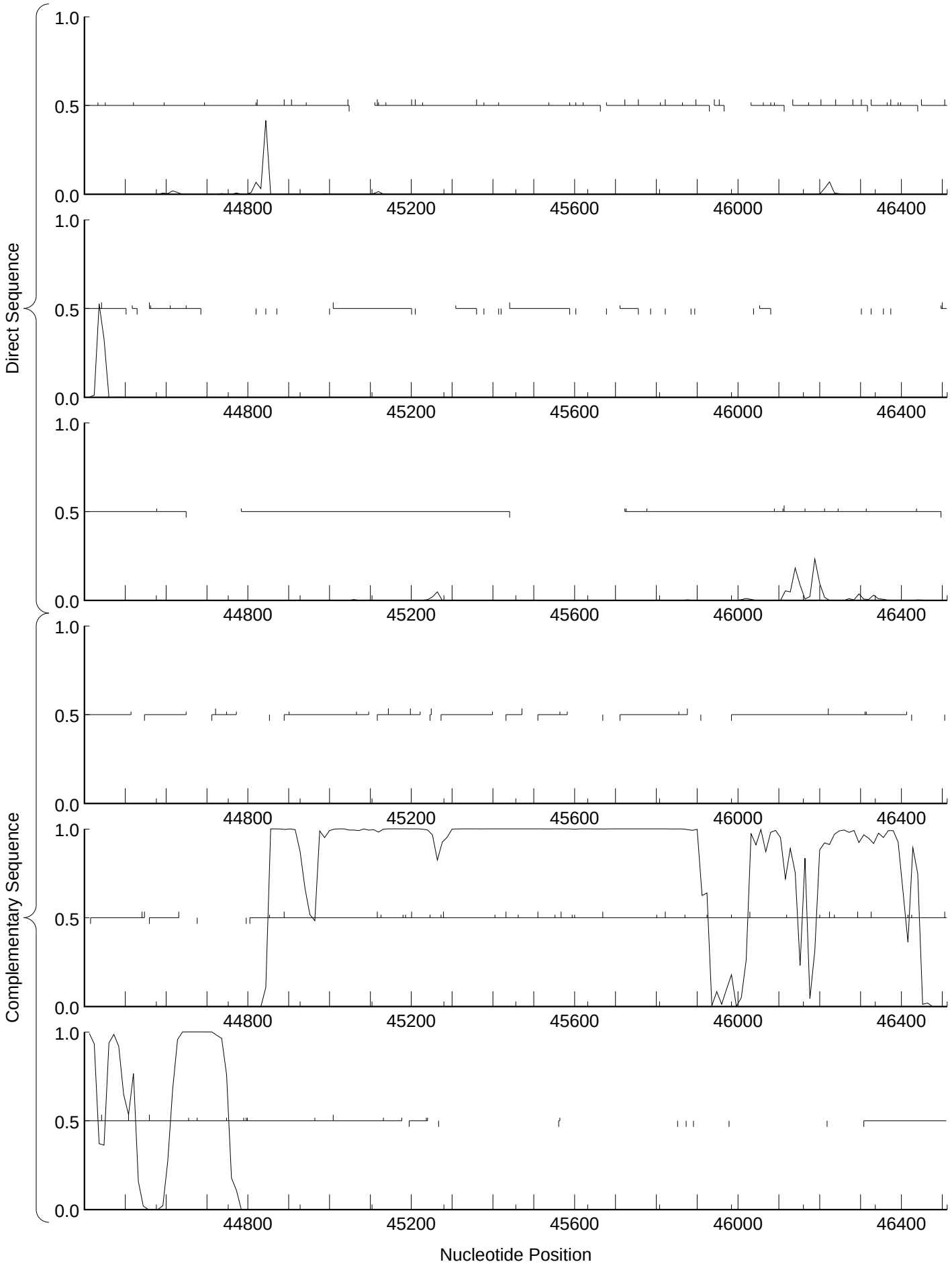
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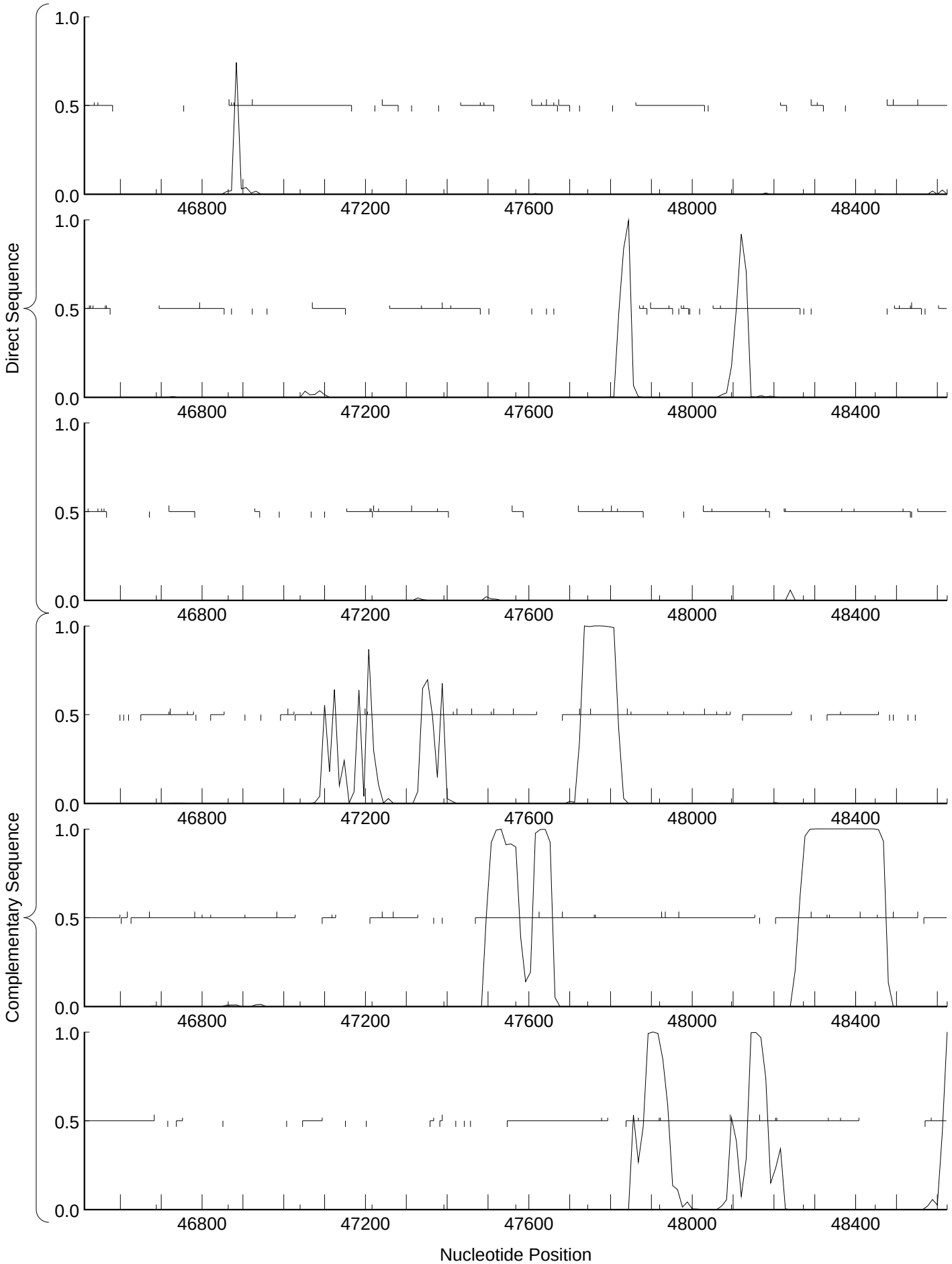
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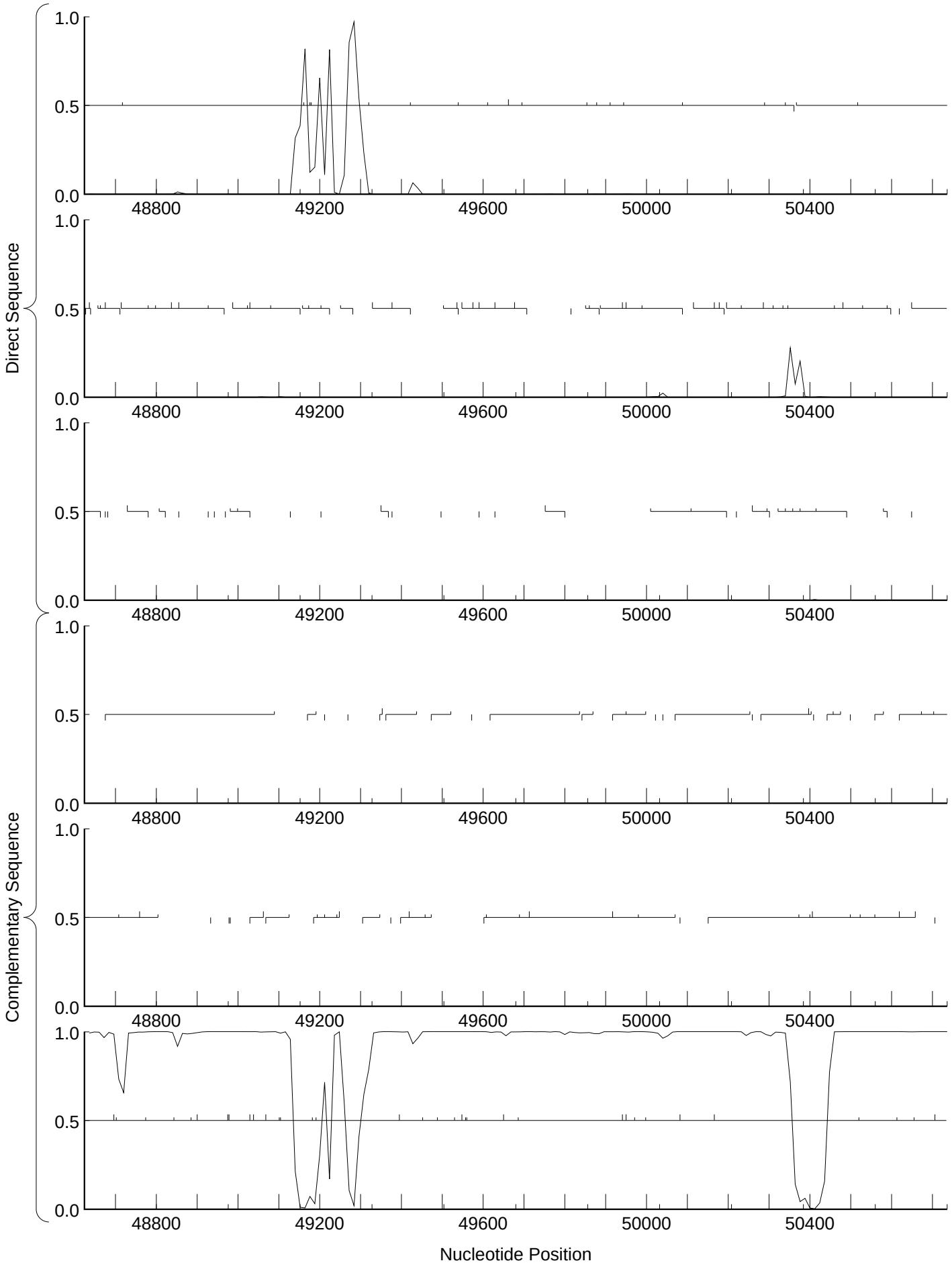
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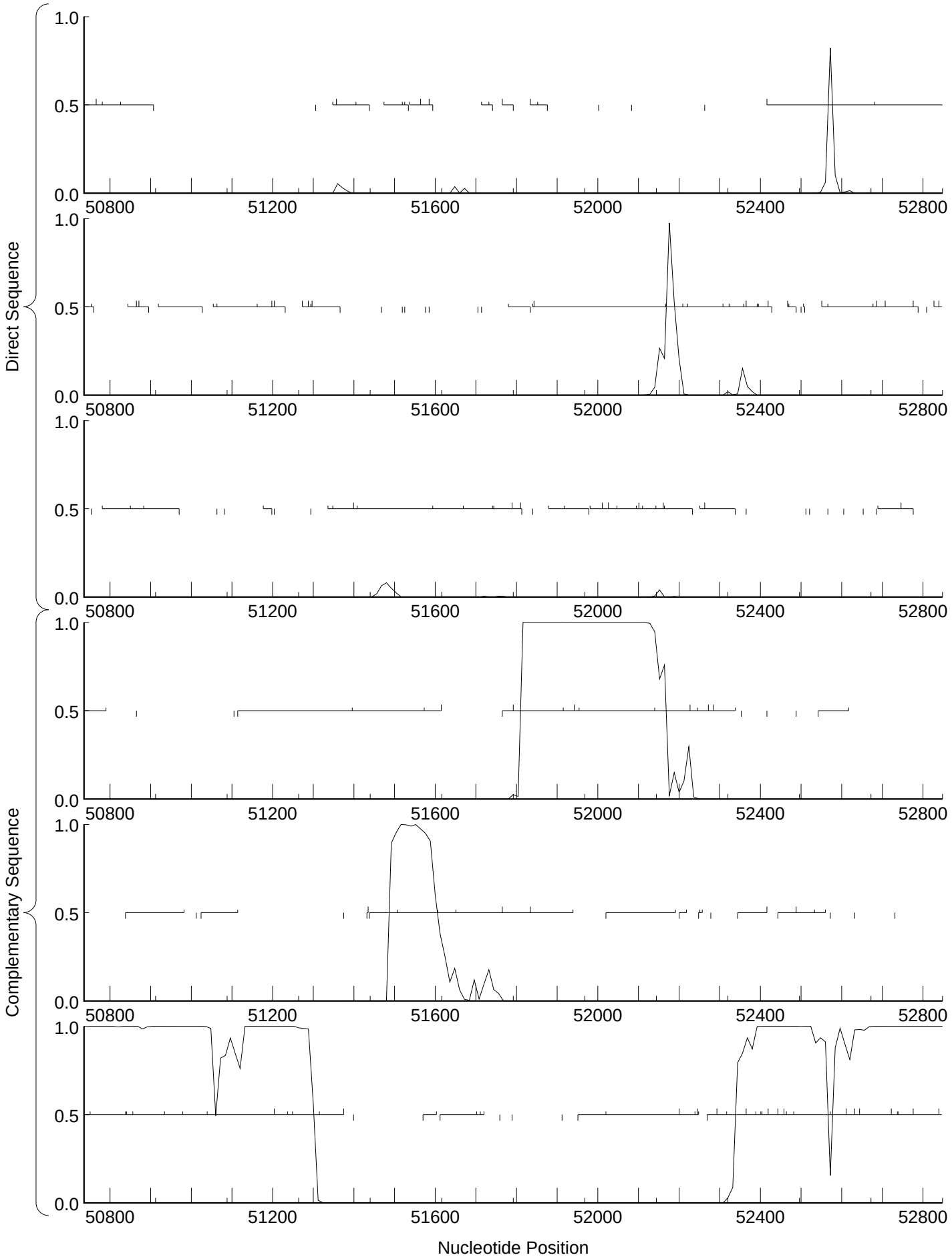
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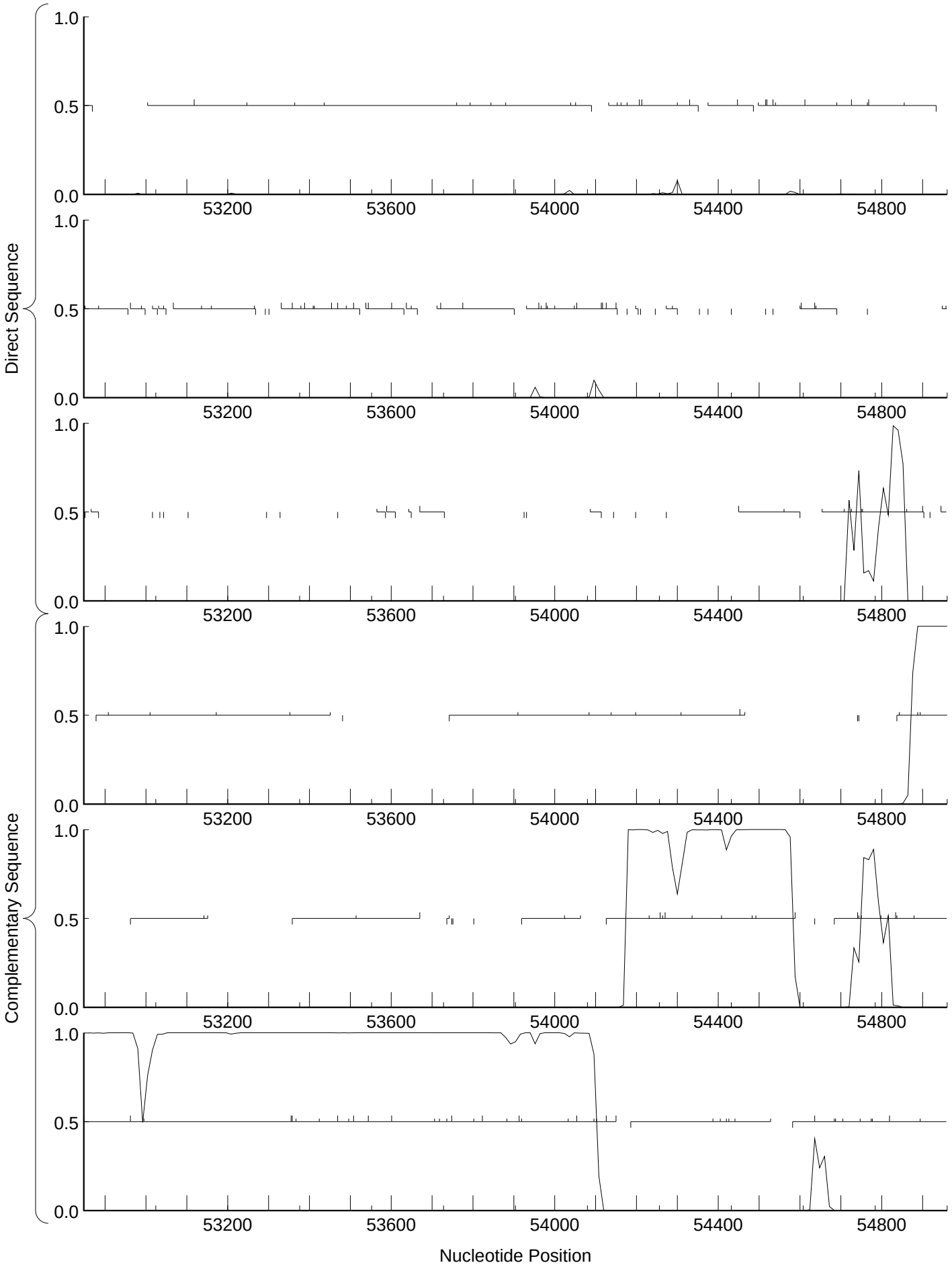
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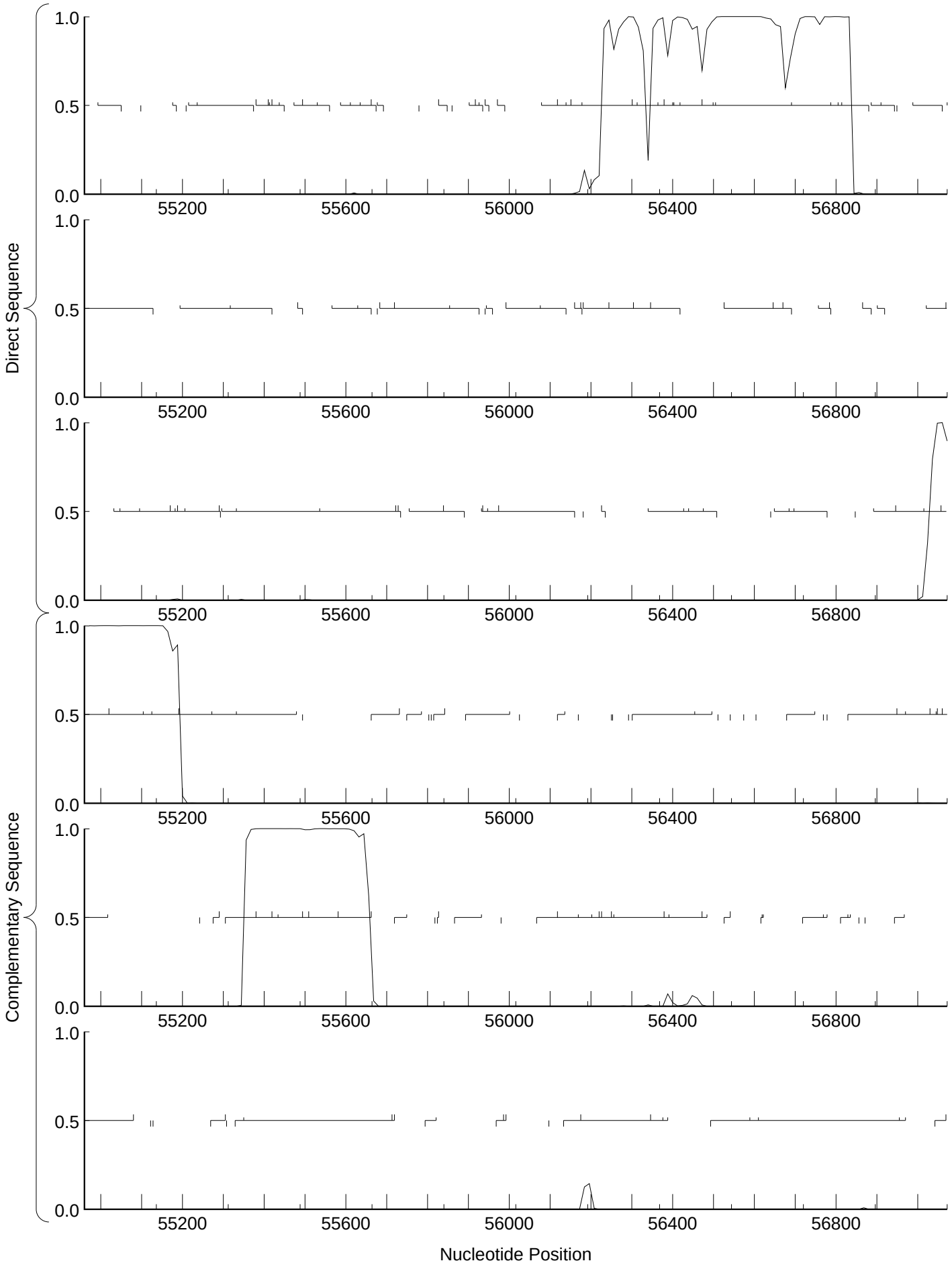
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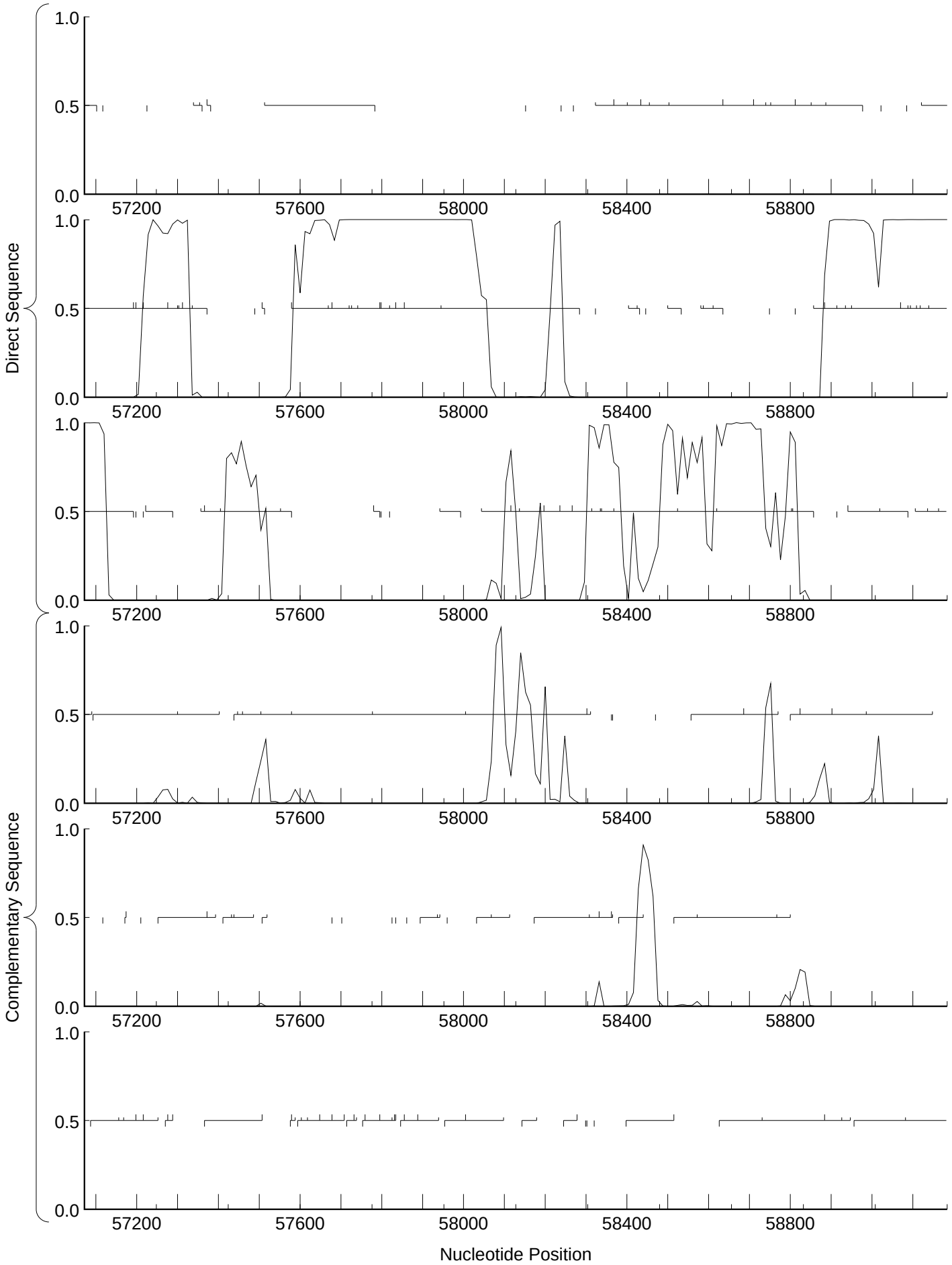
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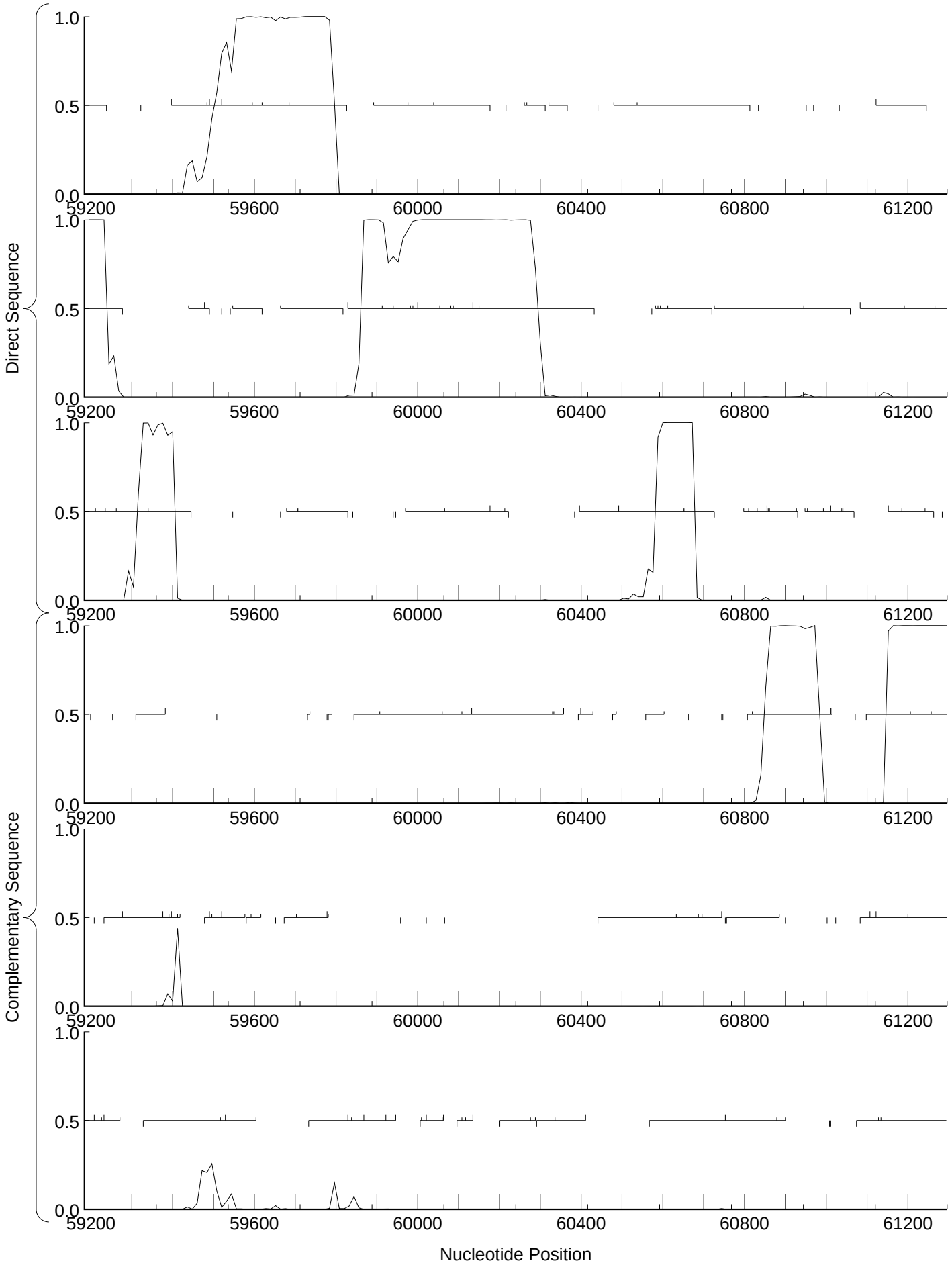
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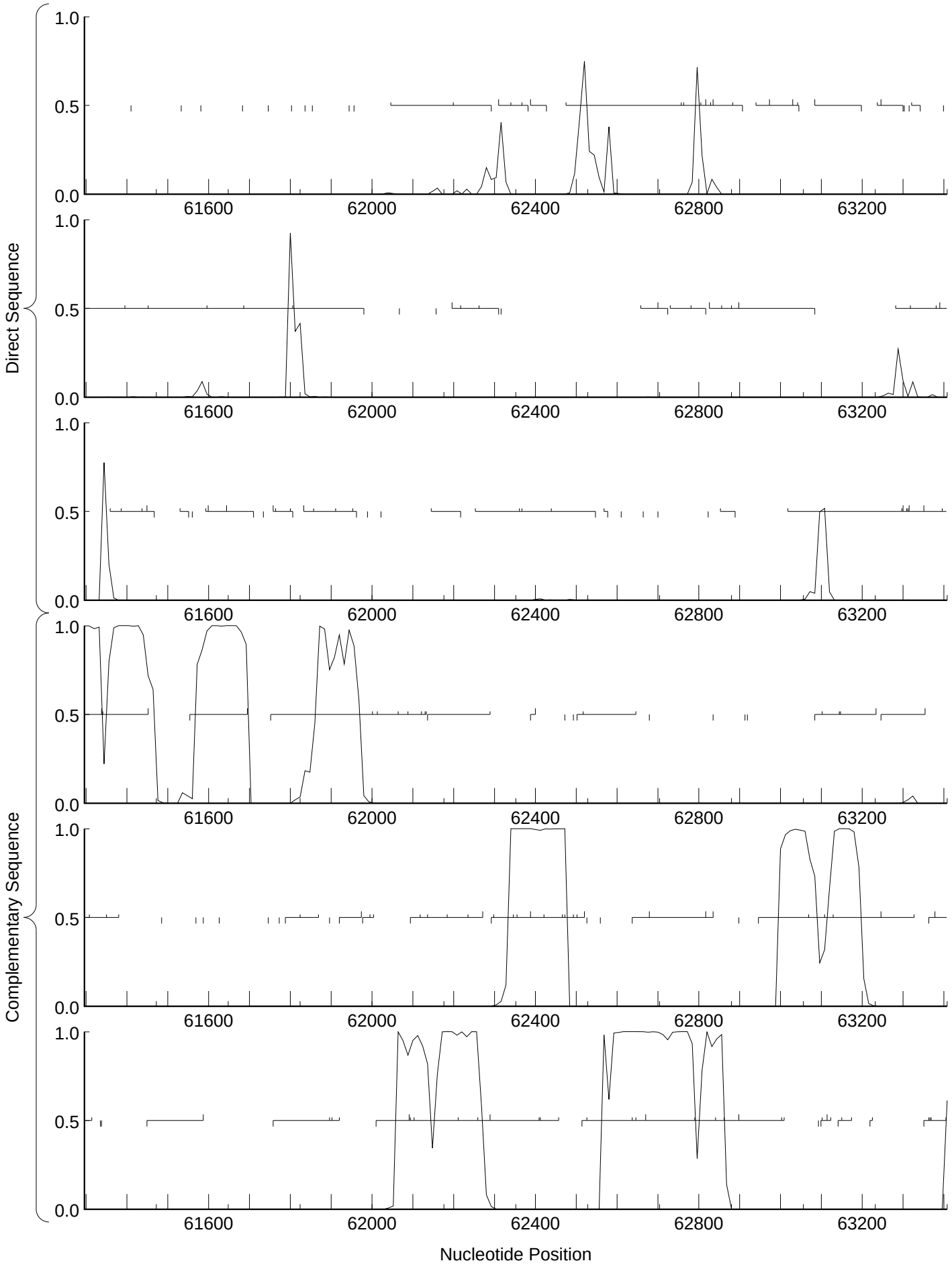
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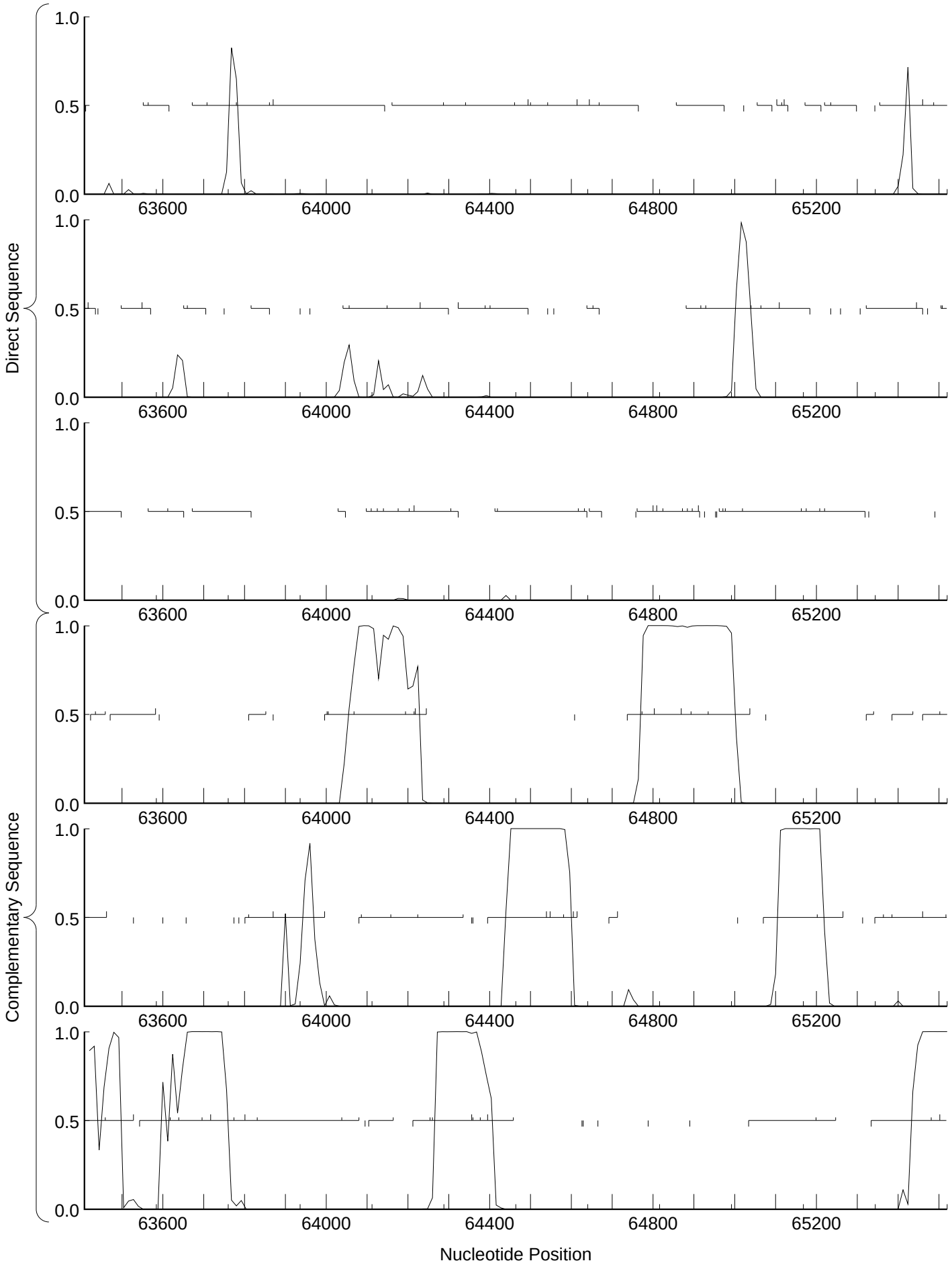
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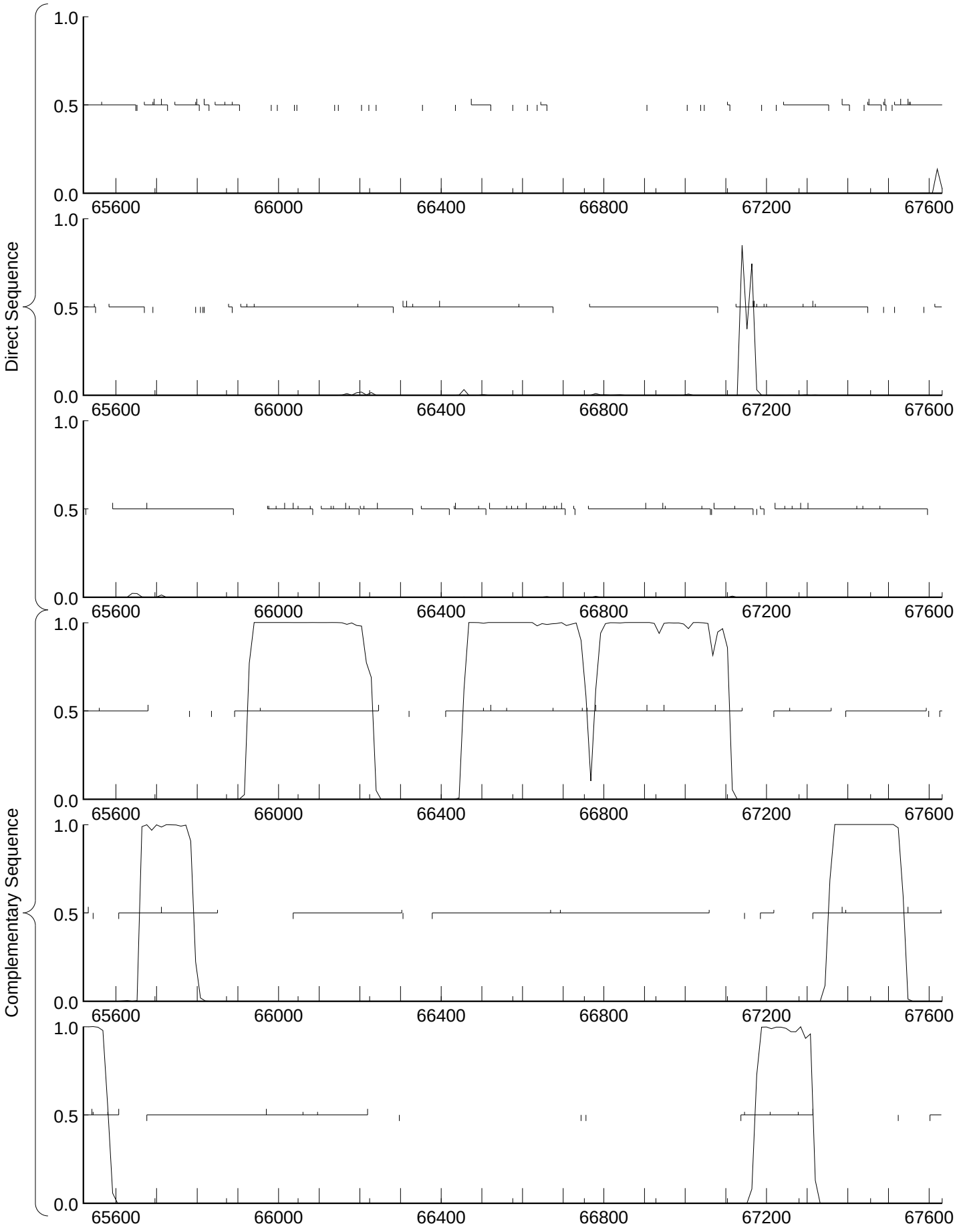
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rcobacterium phage Pinkman complete sequence, 68938 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 32/34



rcobacterium phage Pinkman complete sequence, 68938 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 33/34



rcobacterium phage Pinkman complete sequence, 68938 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 34/34

