# GeneMark

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

## PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Saguaro complete sequence, 69448 bp, circularly permuted, Cluster B7
Analysis Date    :  3/29/24 at 1:30:29
Pages            :  34
Sequence Length  :  69448 bp
GC Content       :   69.54%

Window Length    :  96 bp
Window Step      :  12 bp
Threshold Value  :  0.500

PS-Version       :  1.2

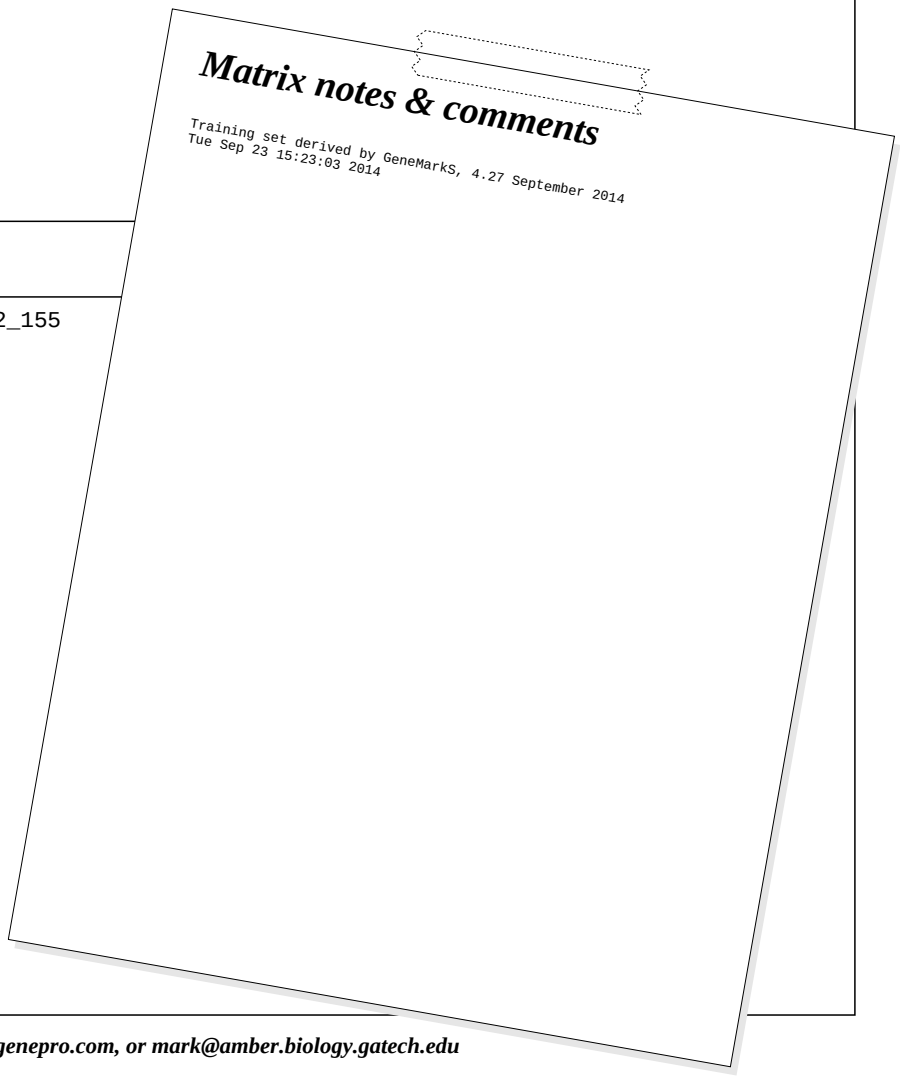
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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## MATRIX INFORMATION

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Author     :  -

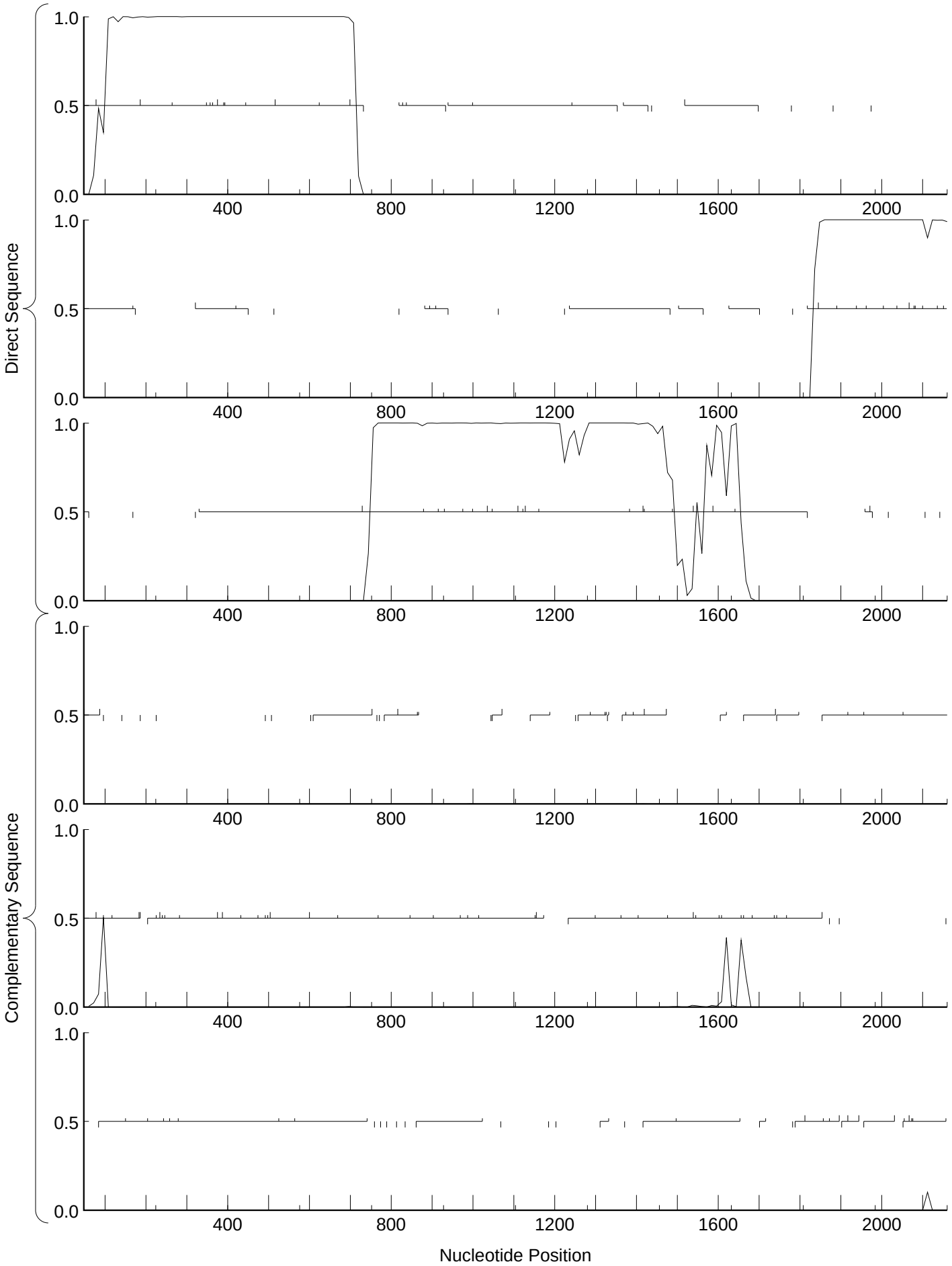
Order      :  4


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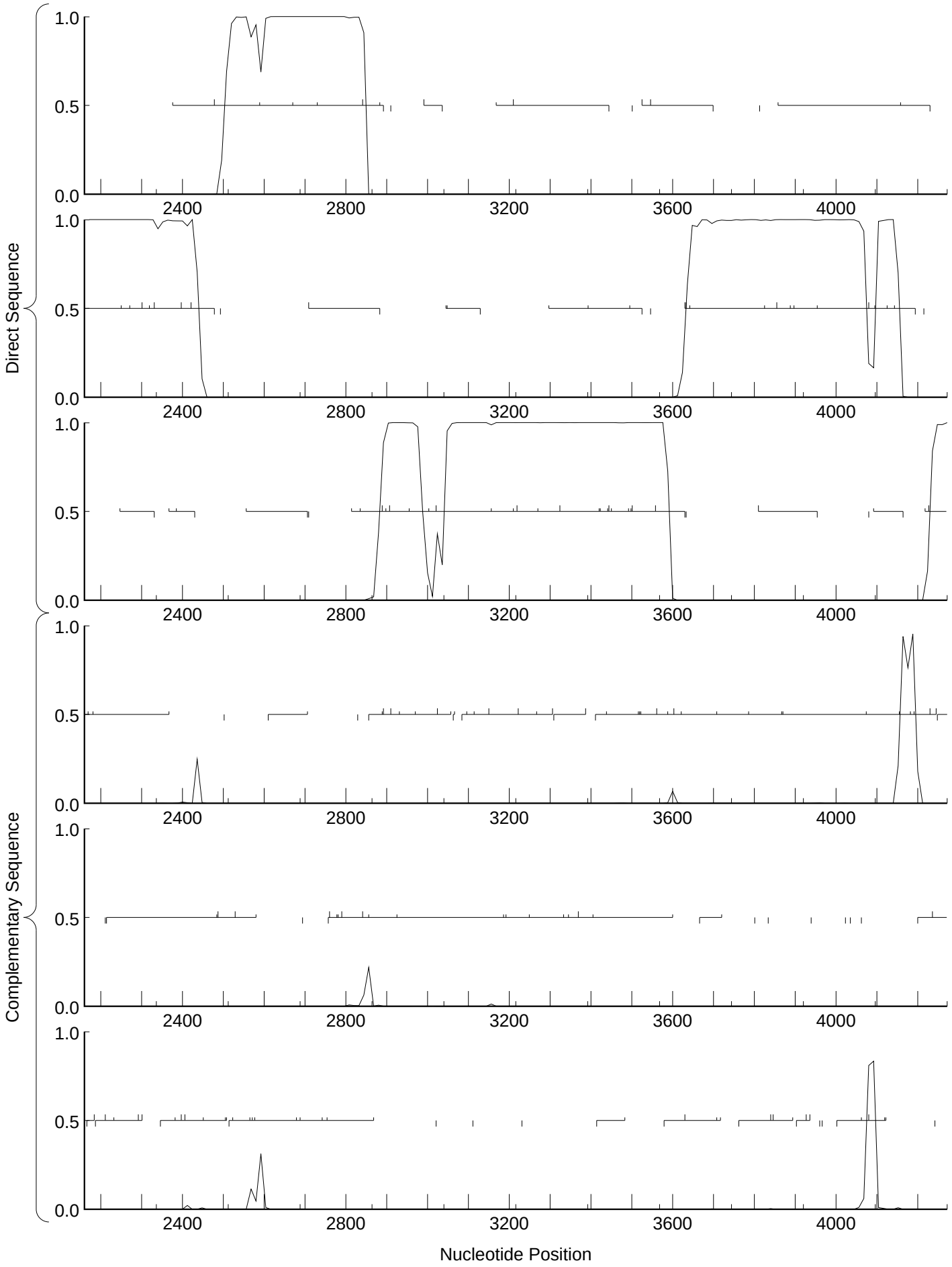



*For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu*

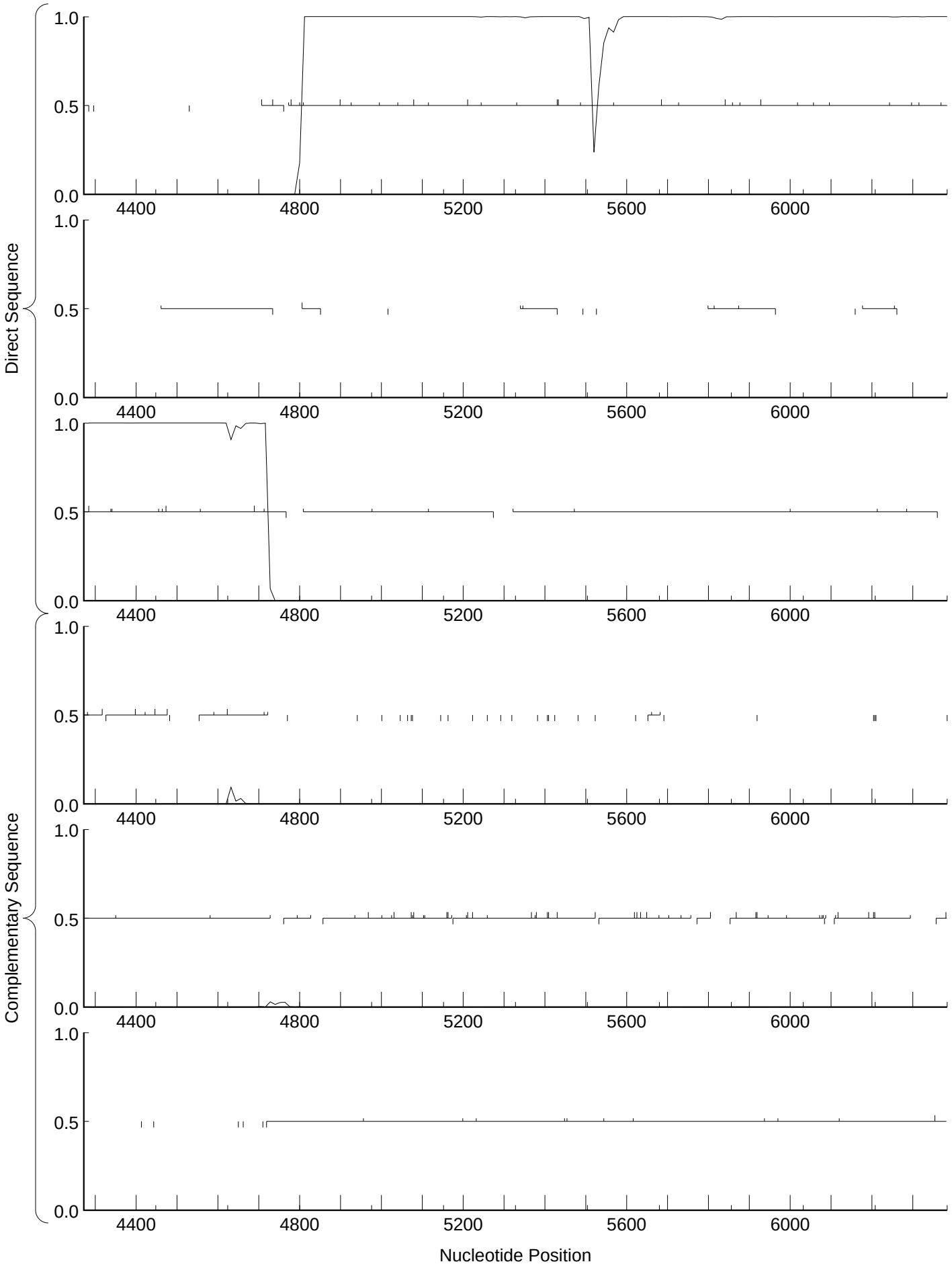
Mycobacterium phage Saguaro complete sequence, 69448 bp, circularly permuted, Cluster B7, Order 4, Window 96, Step 12, 2/34



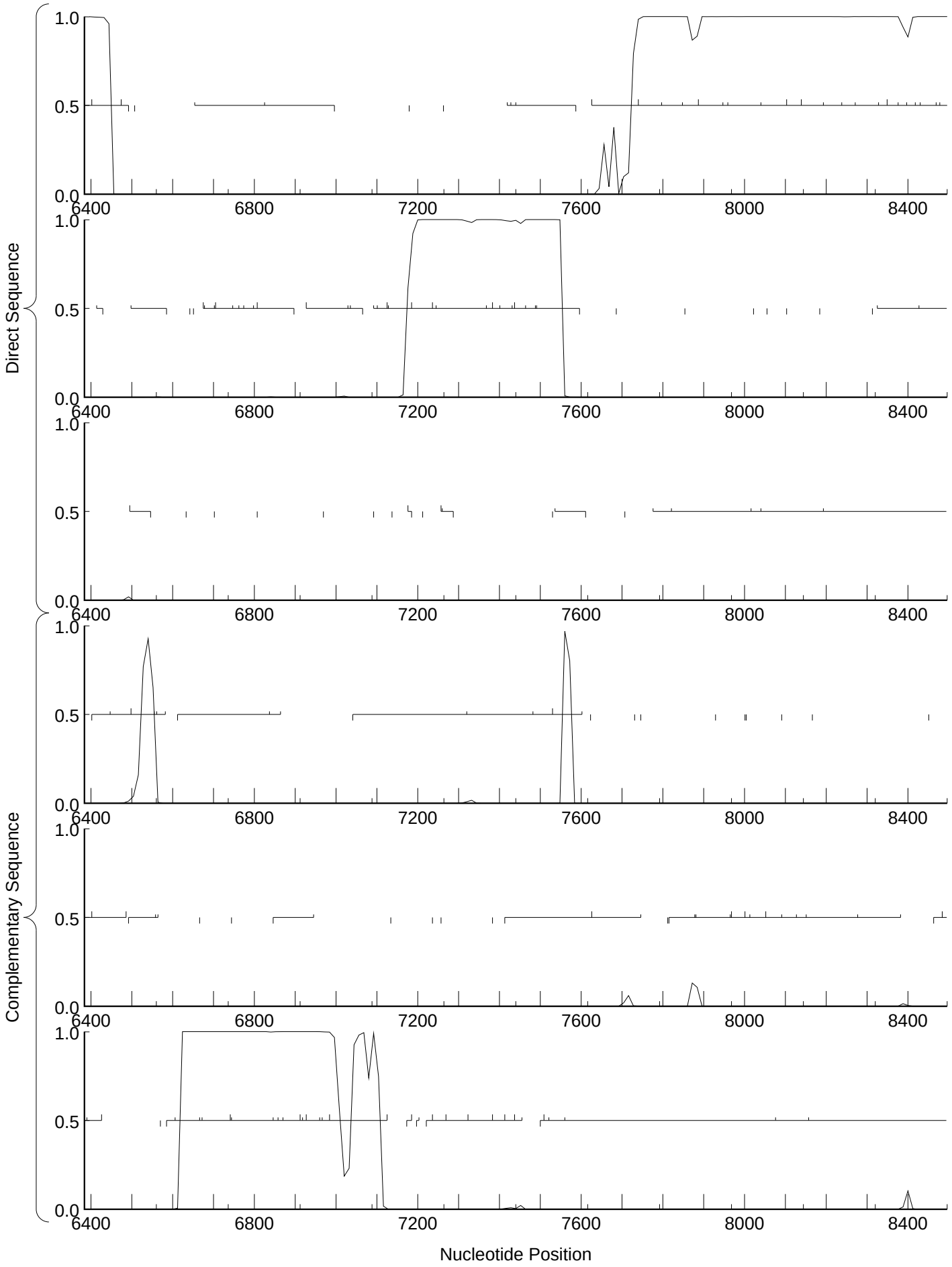
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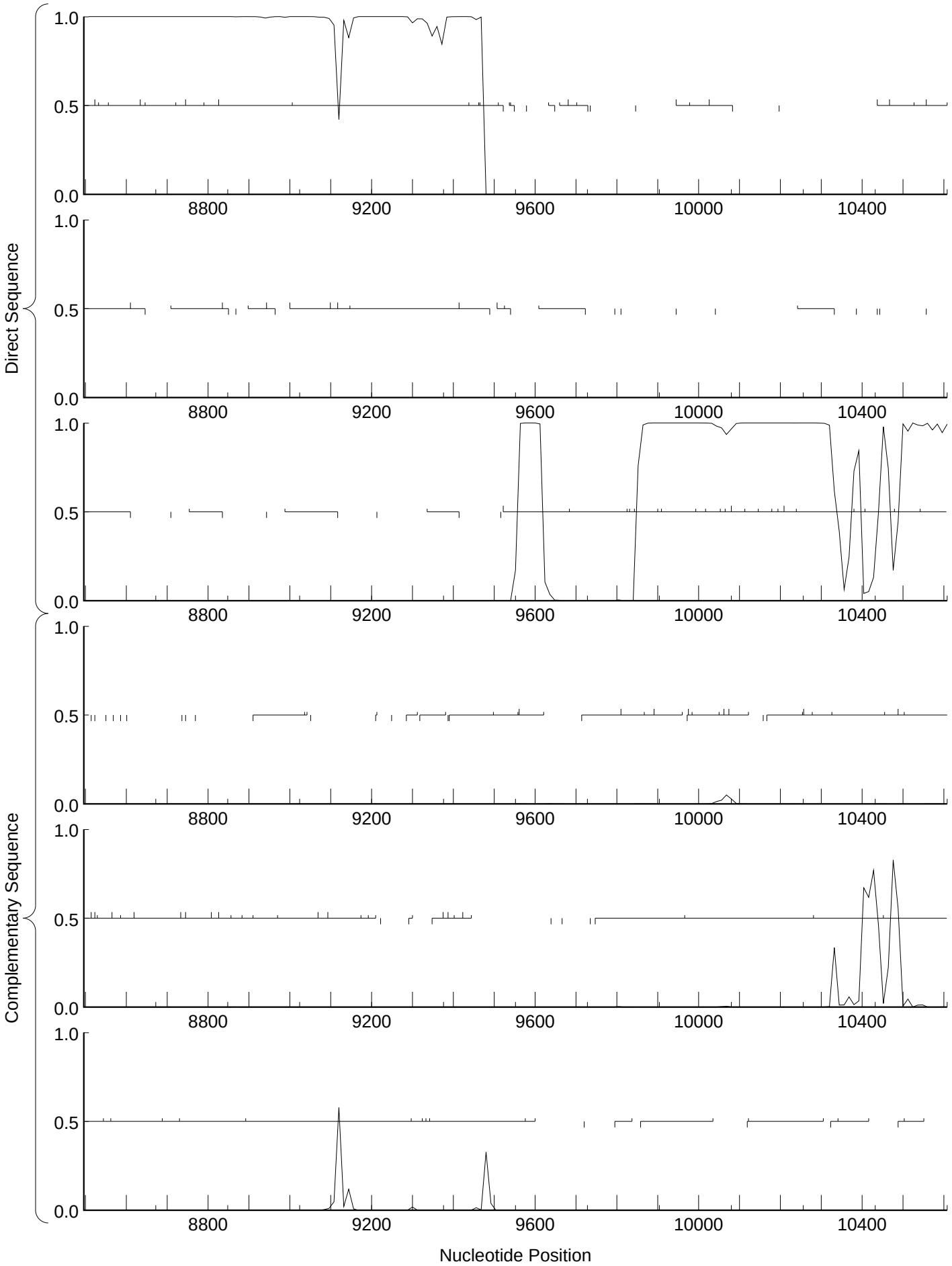
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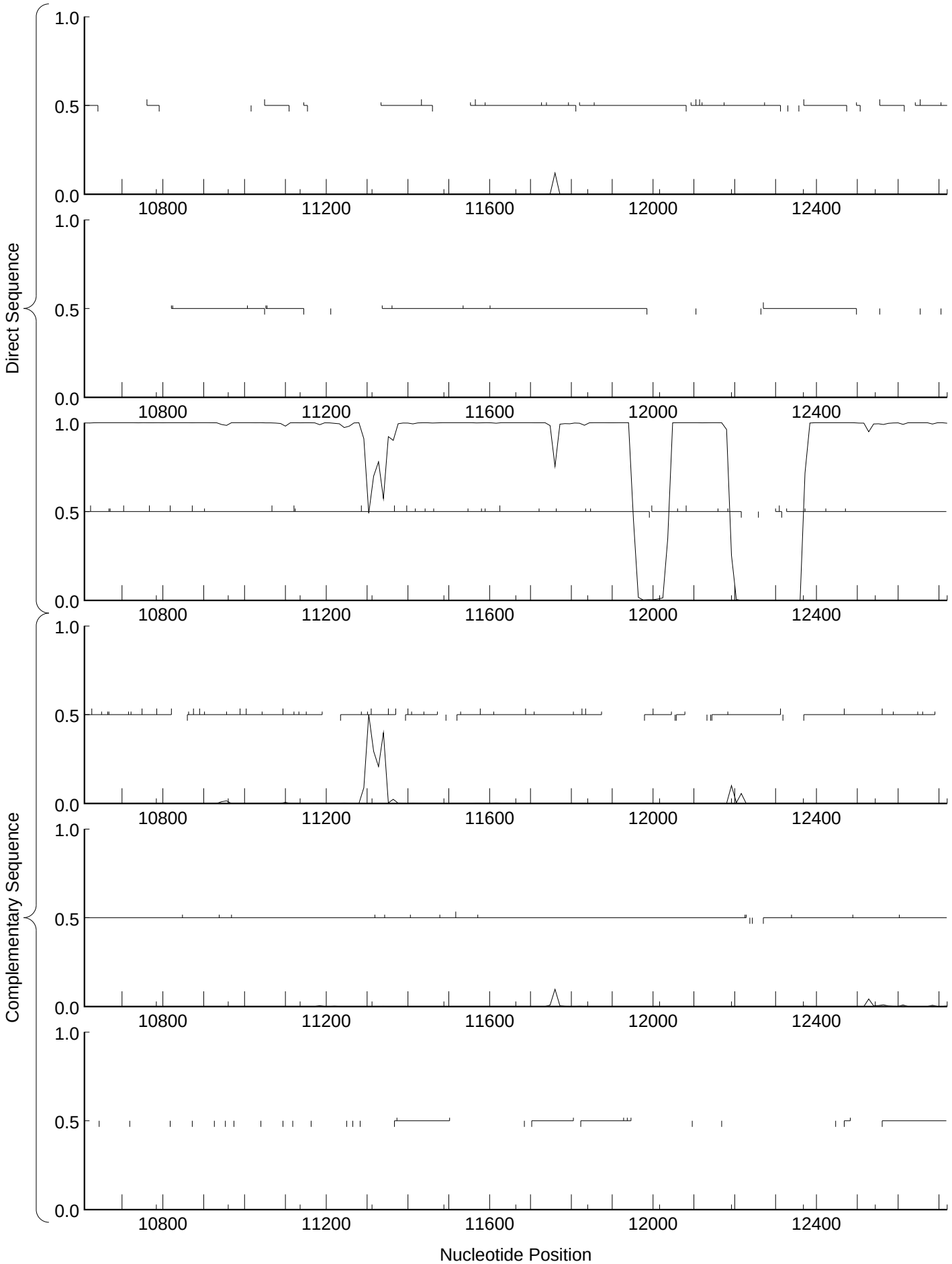
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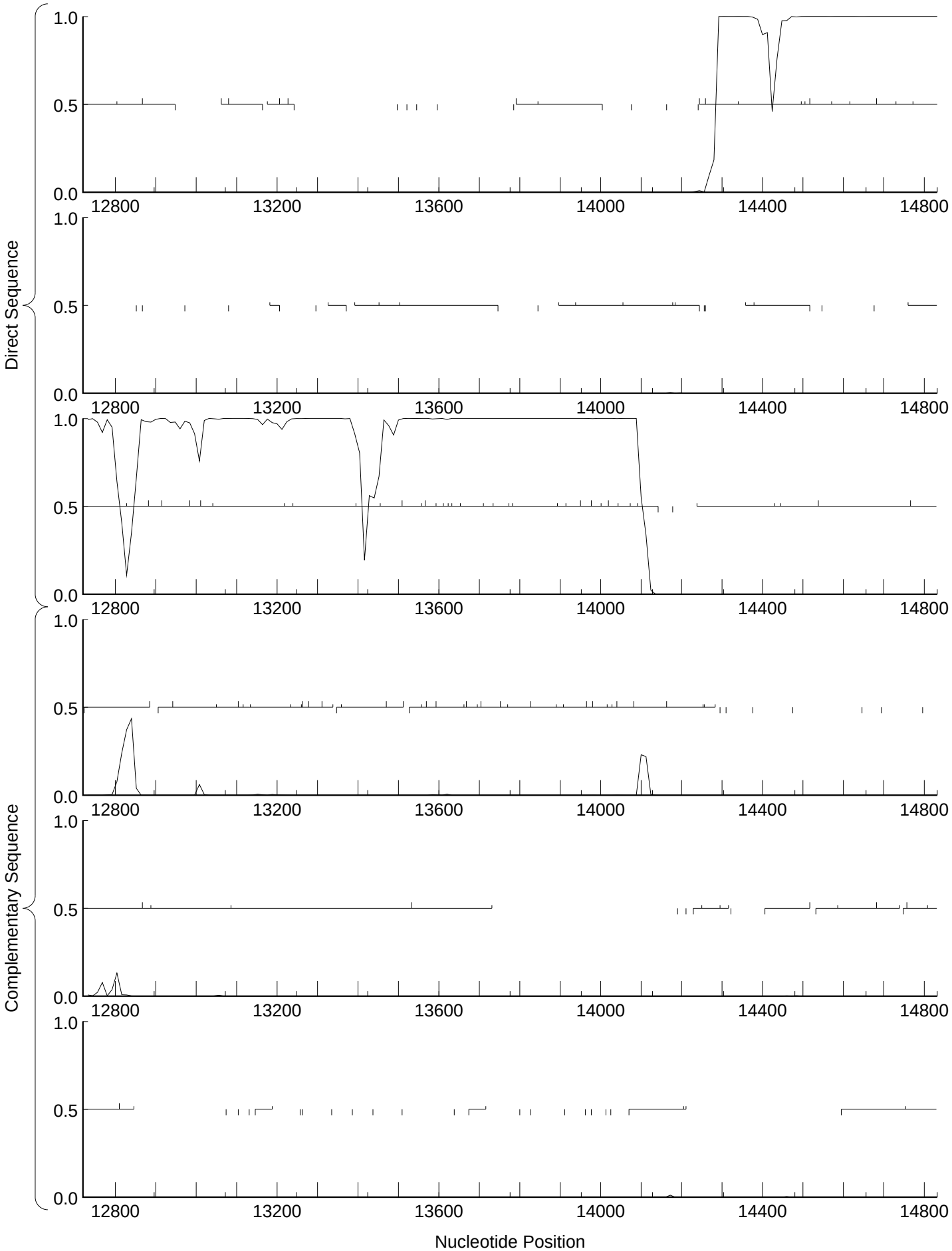
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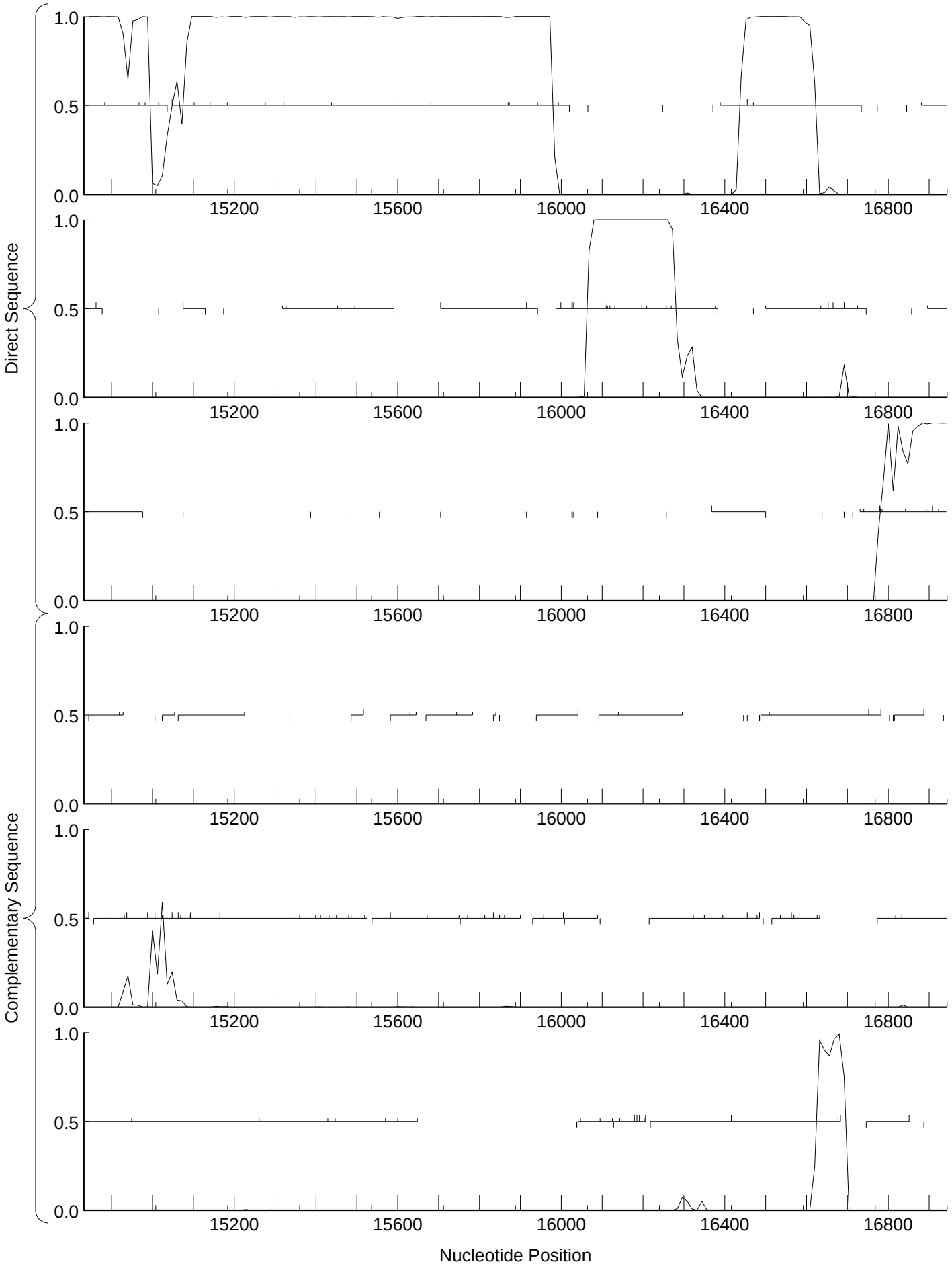
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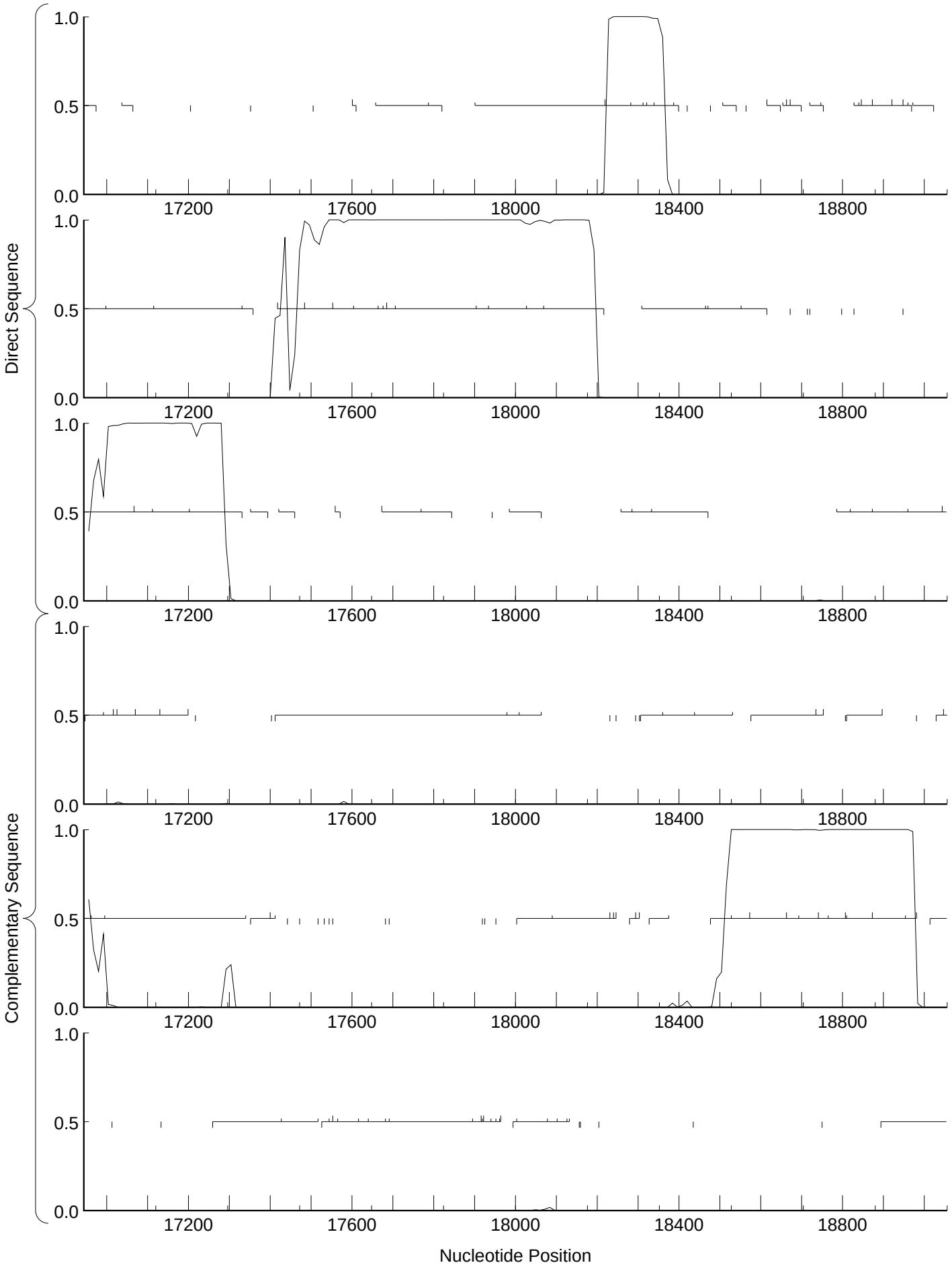
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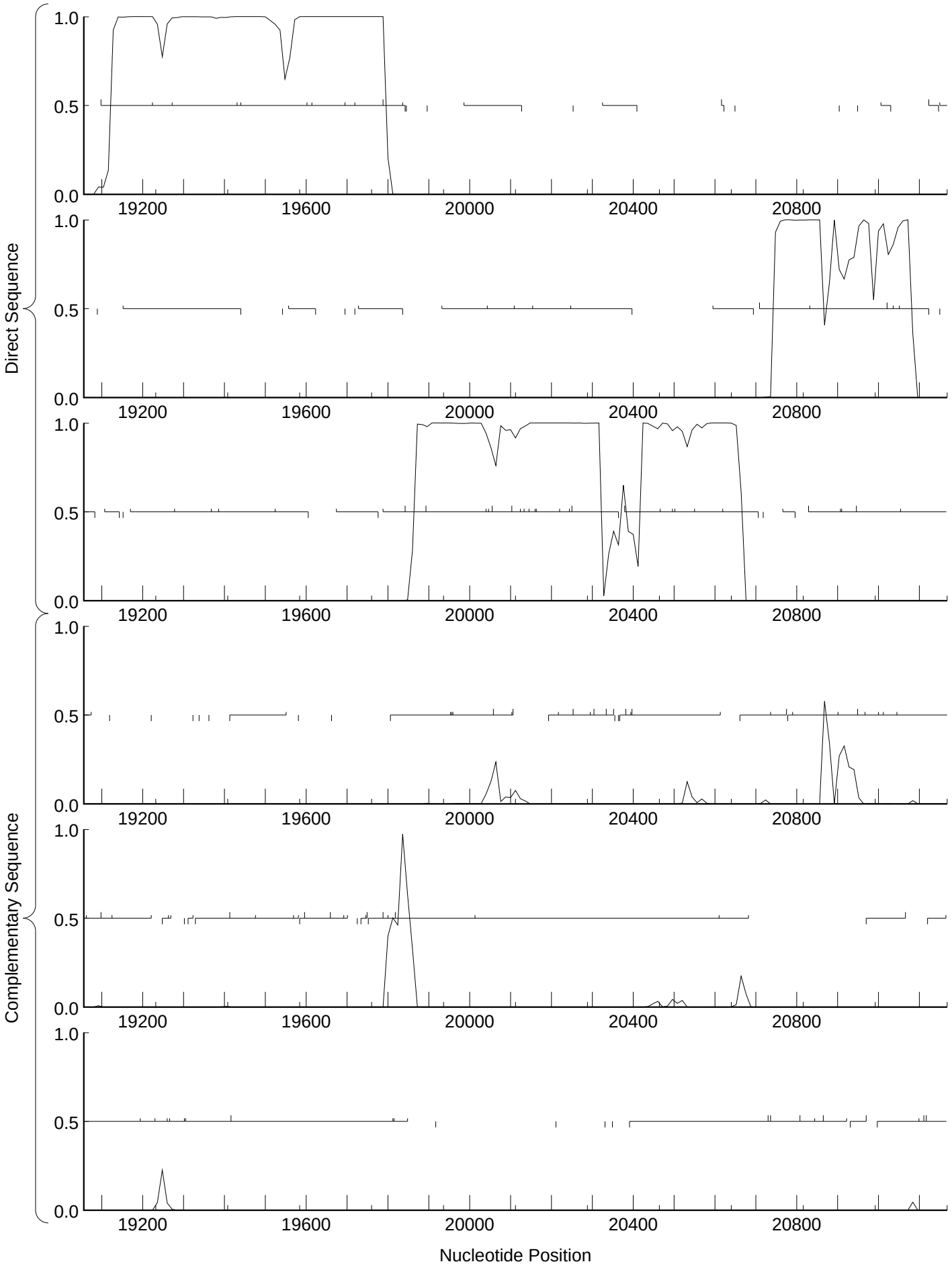
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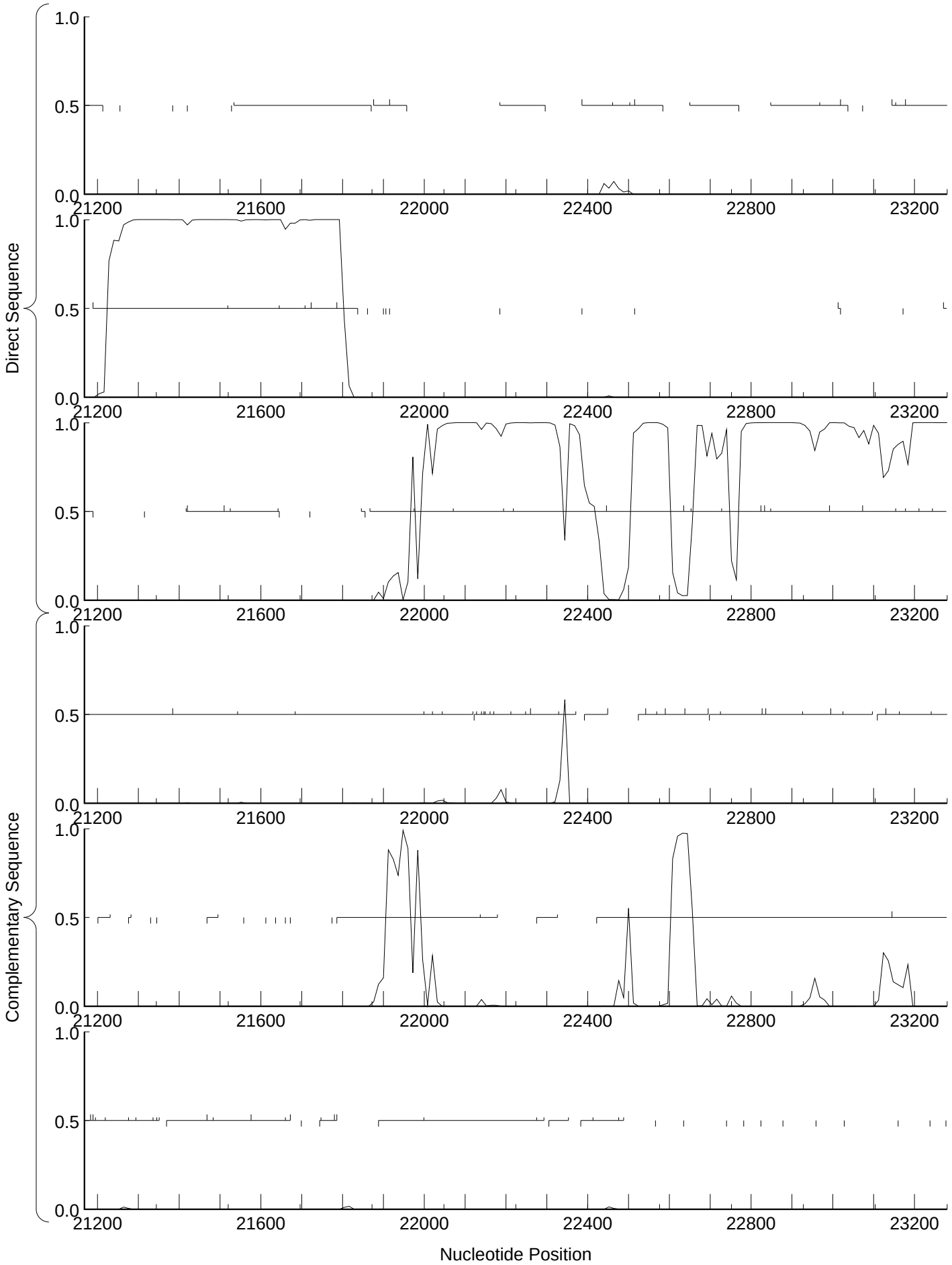
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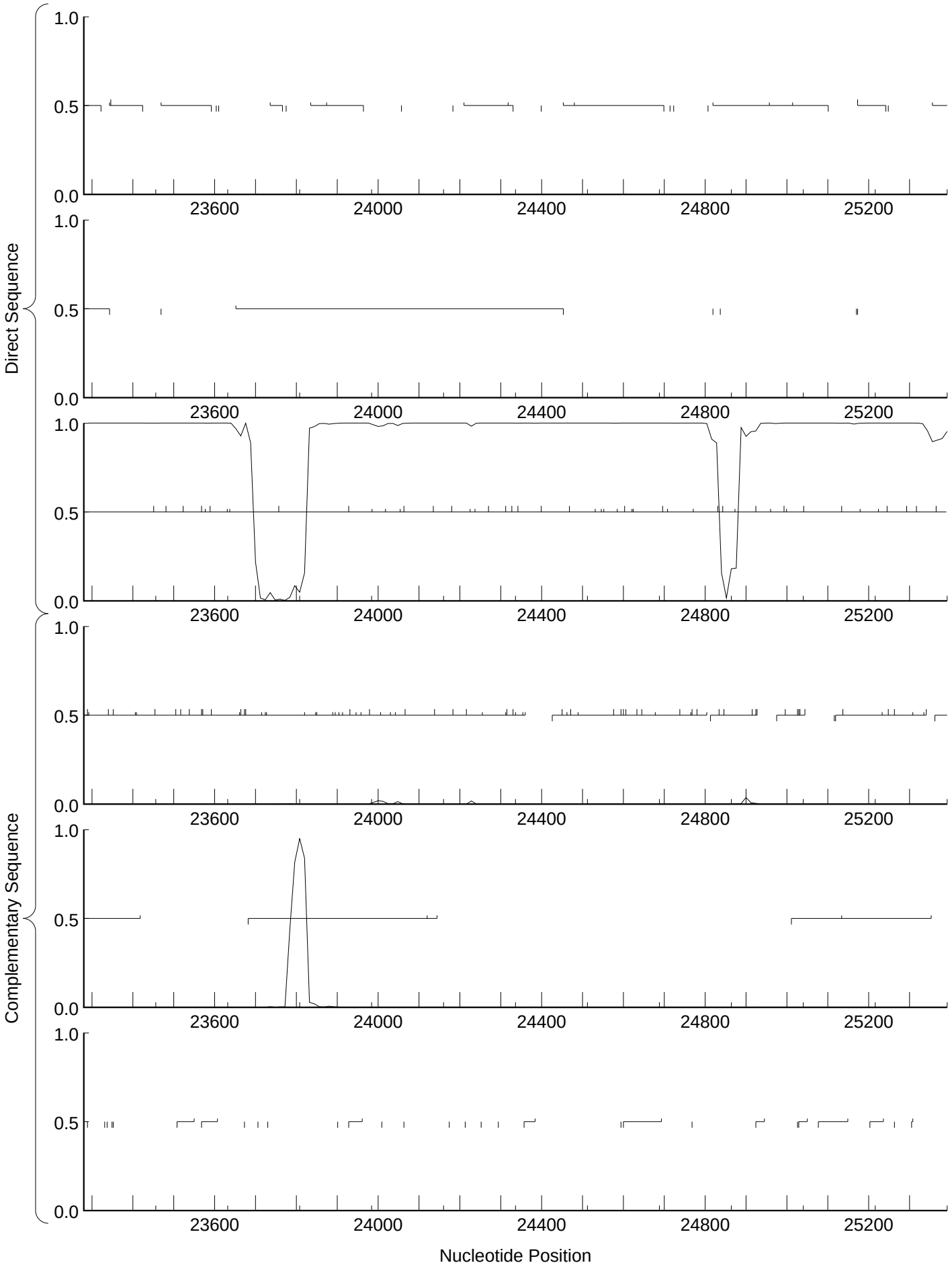
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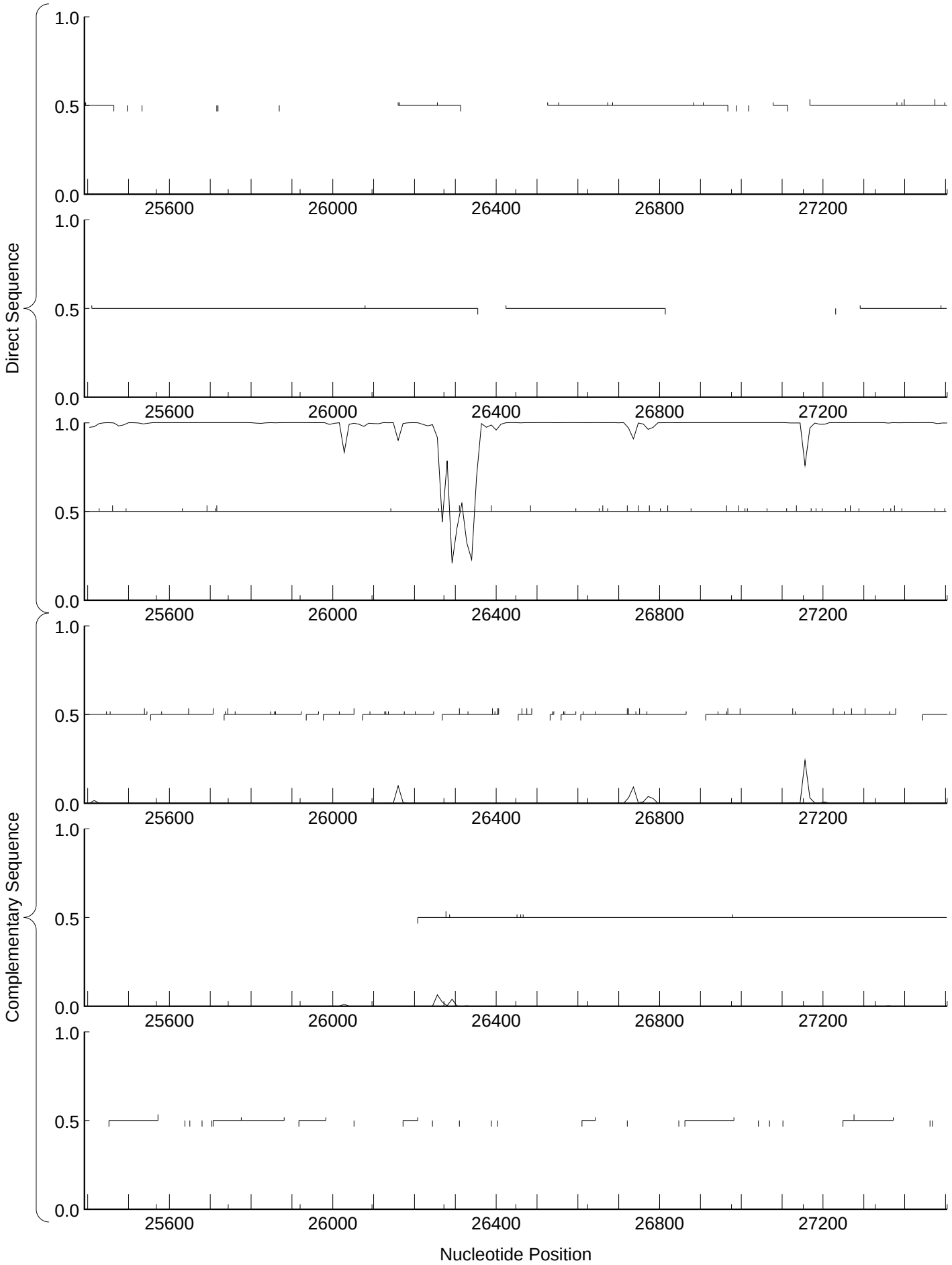
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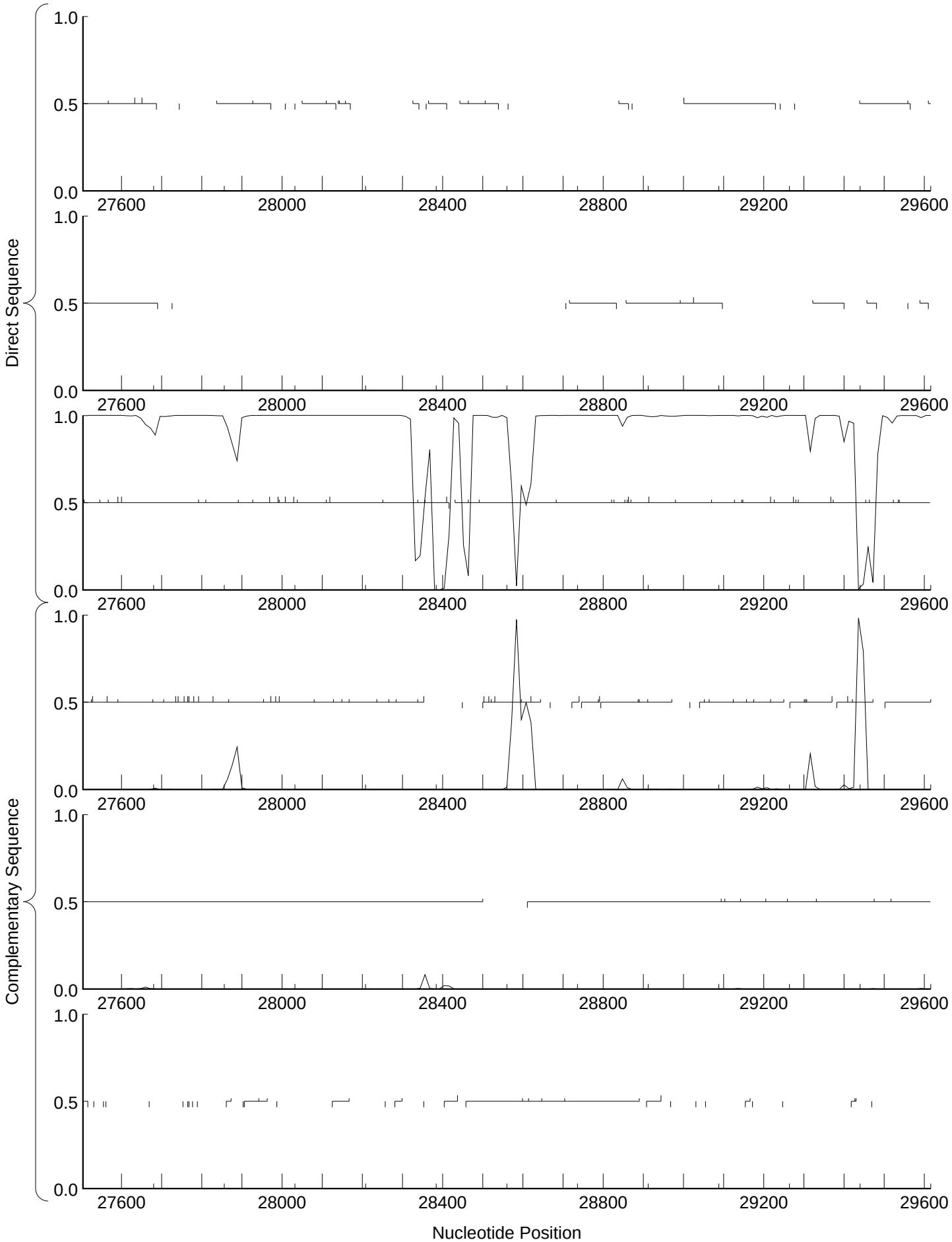
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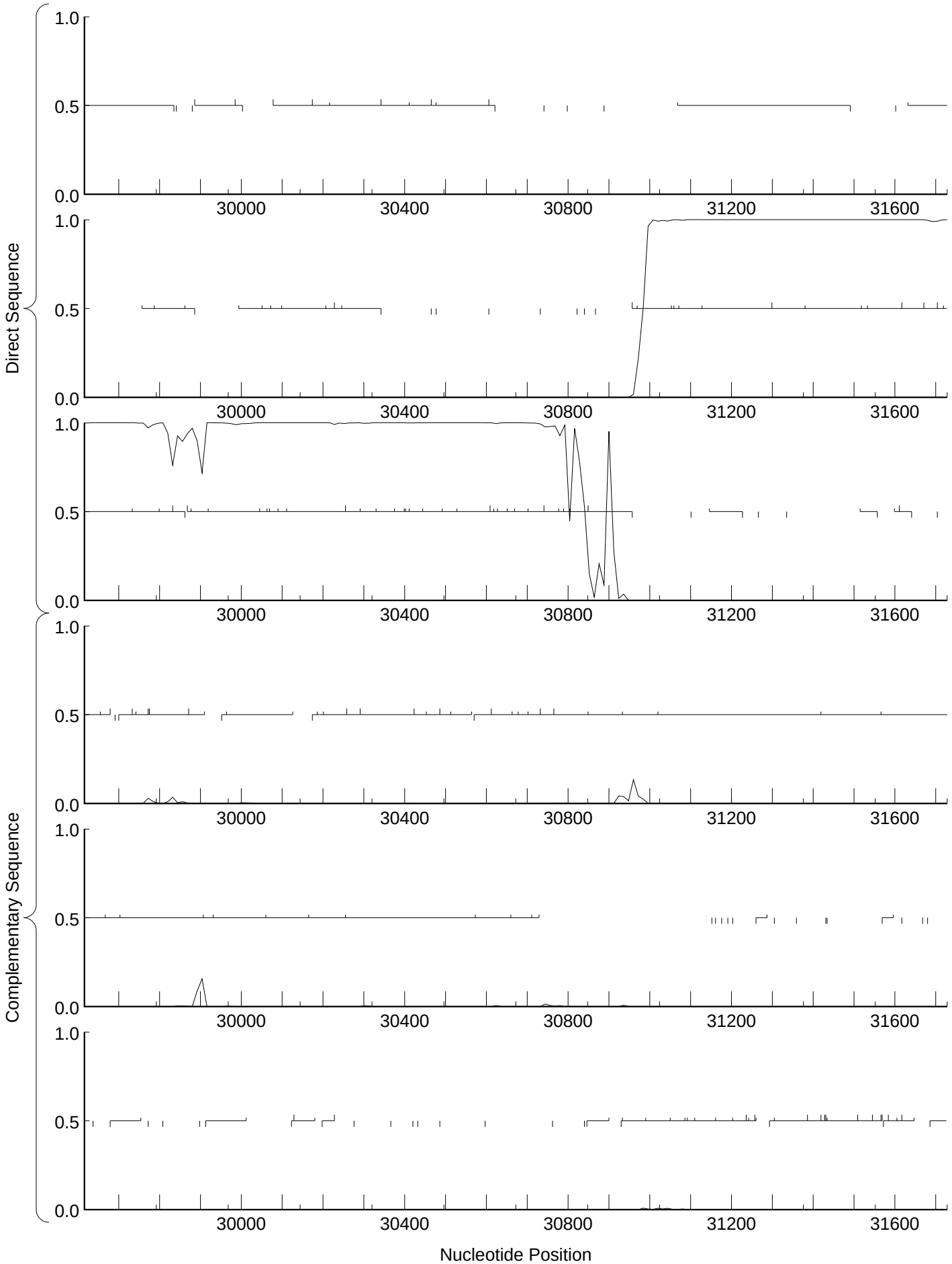
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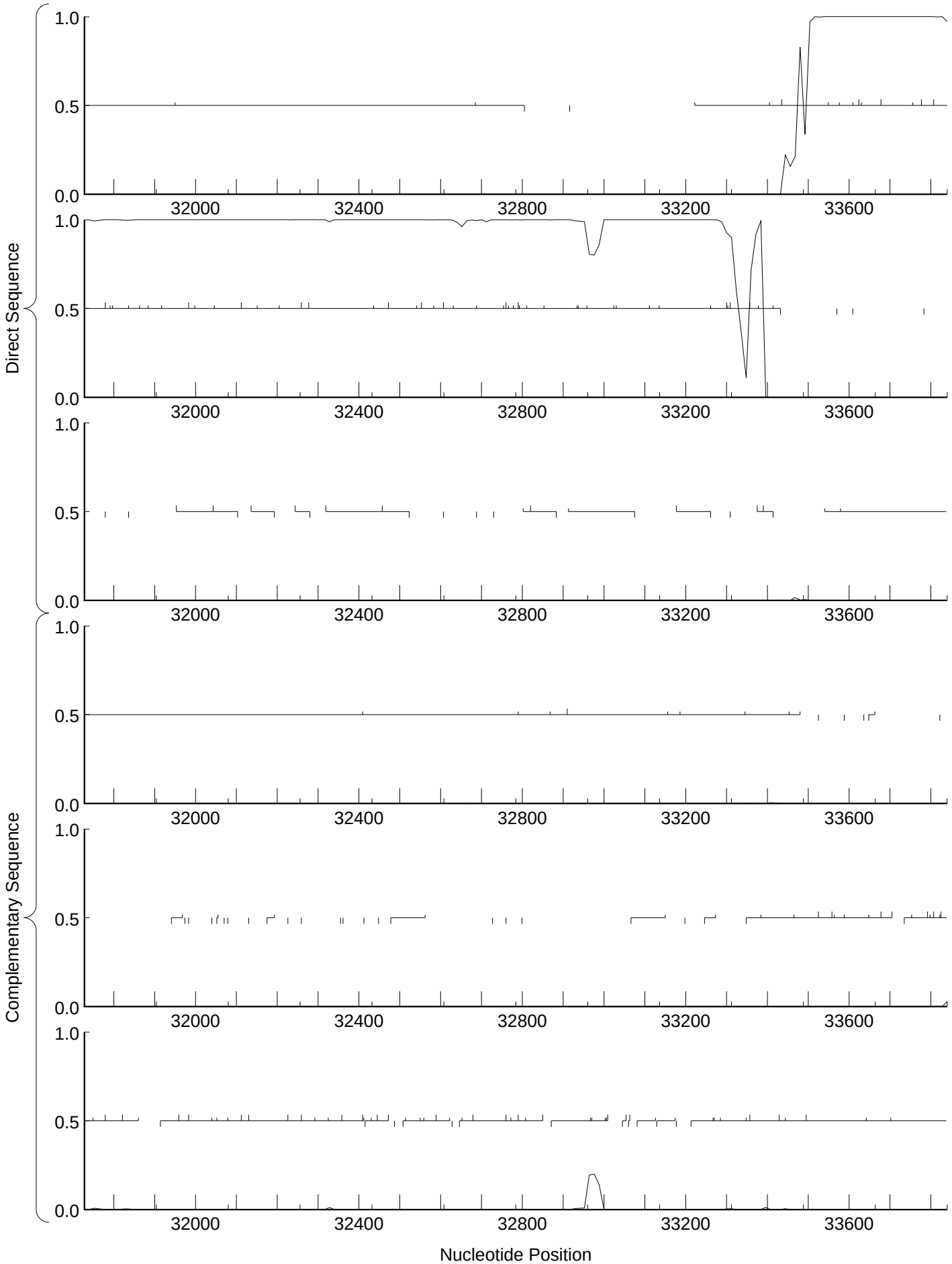
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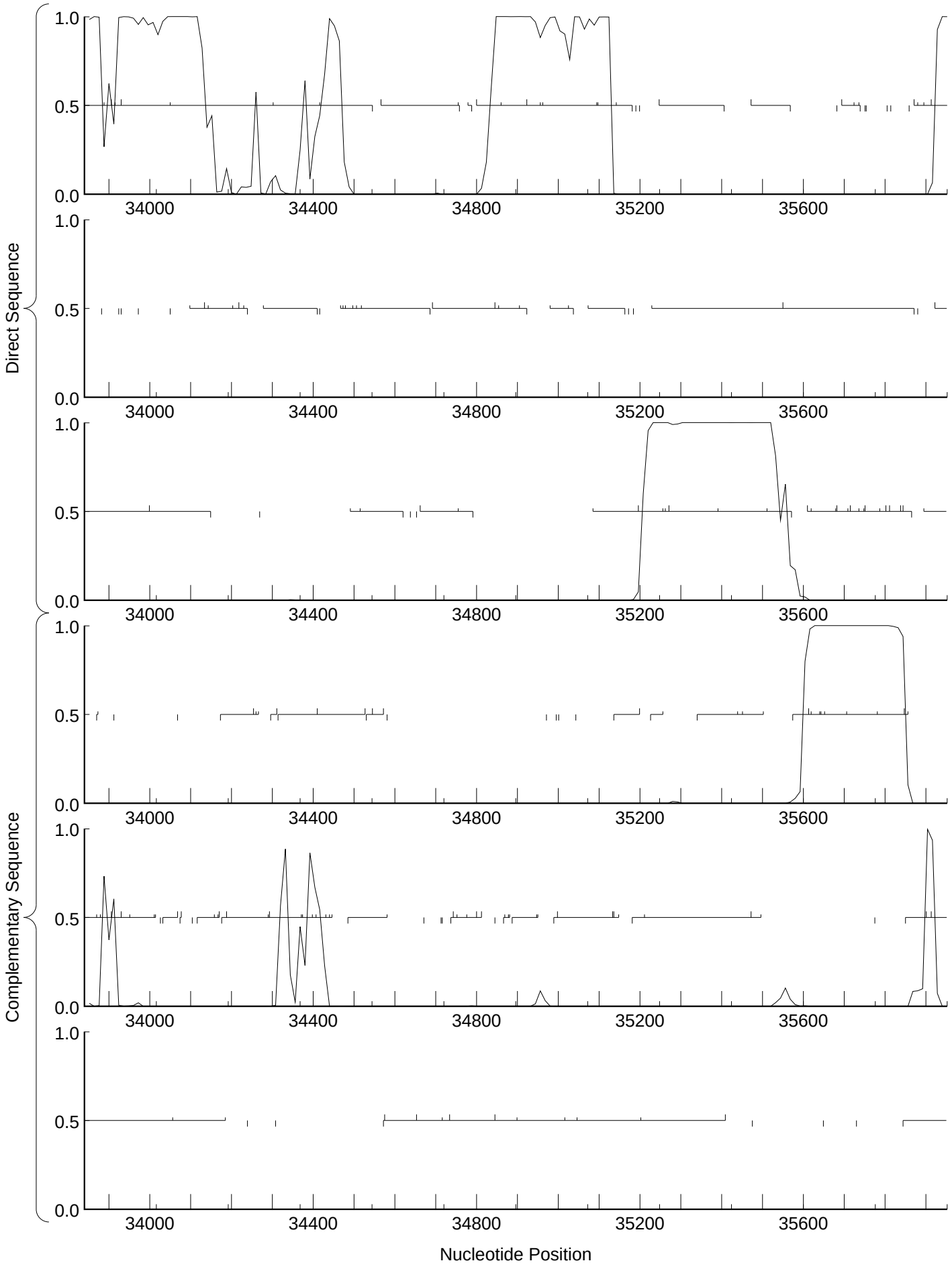
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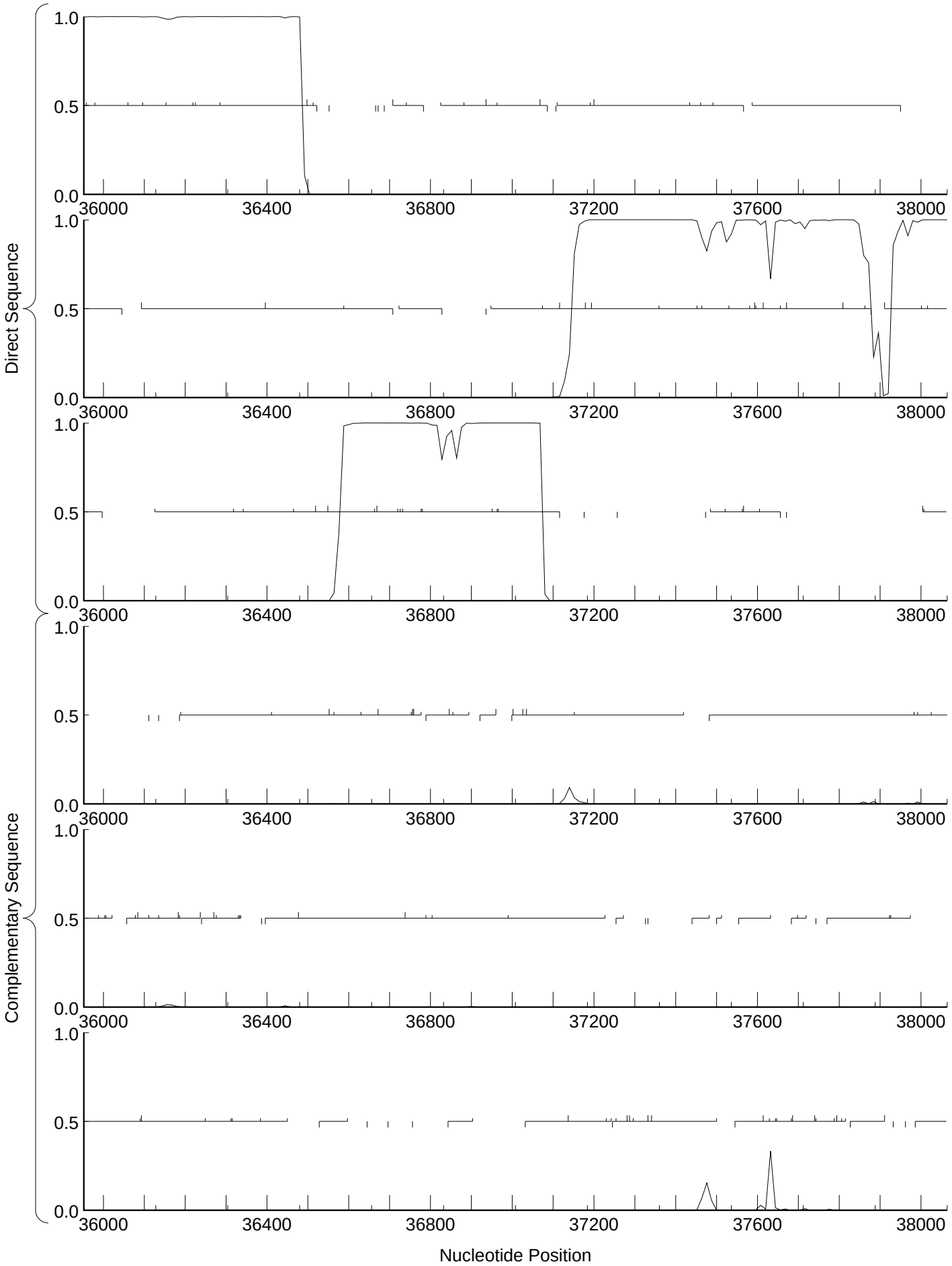
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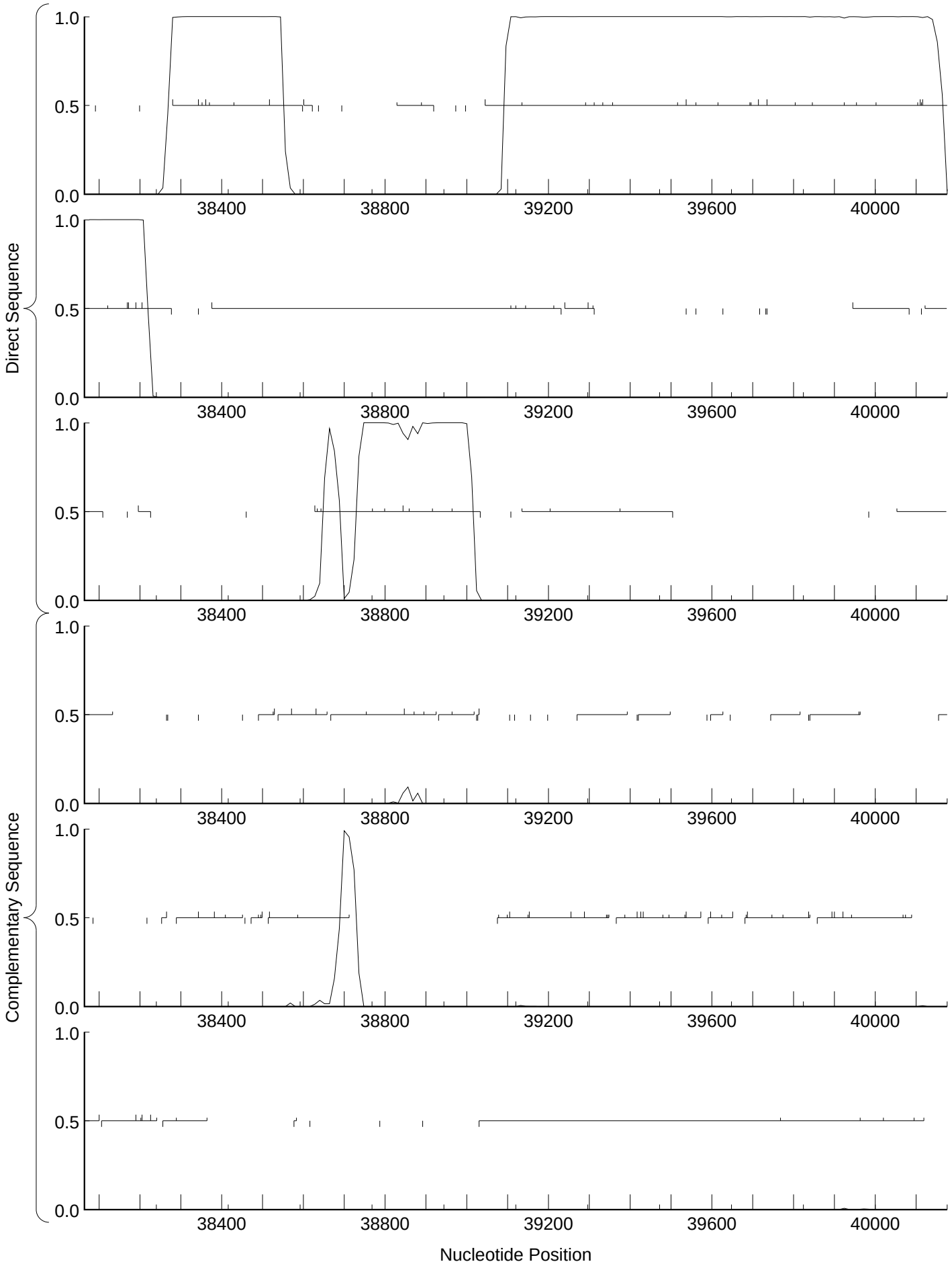


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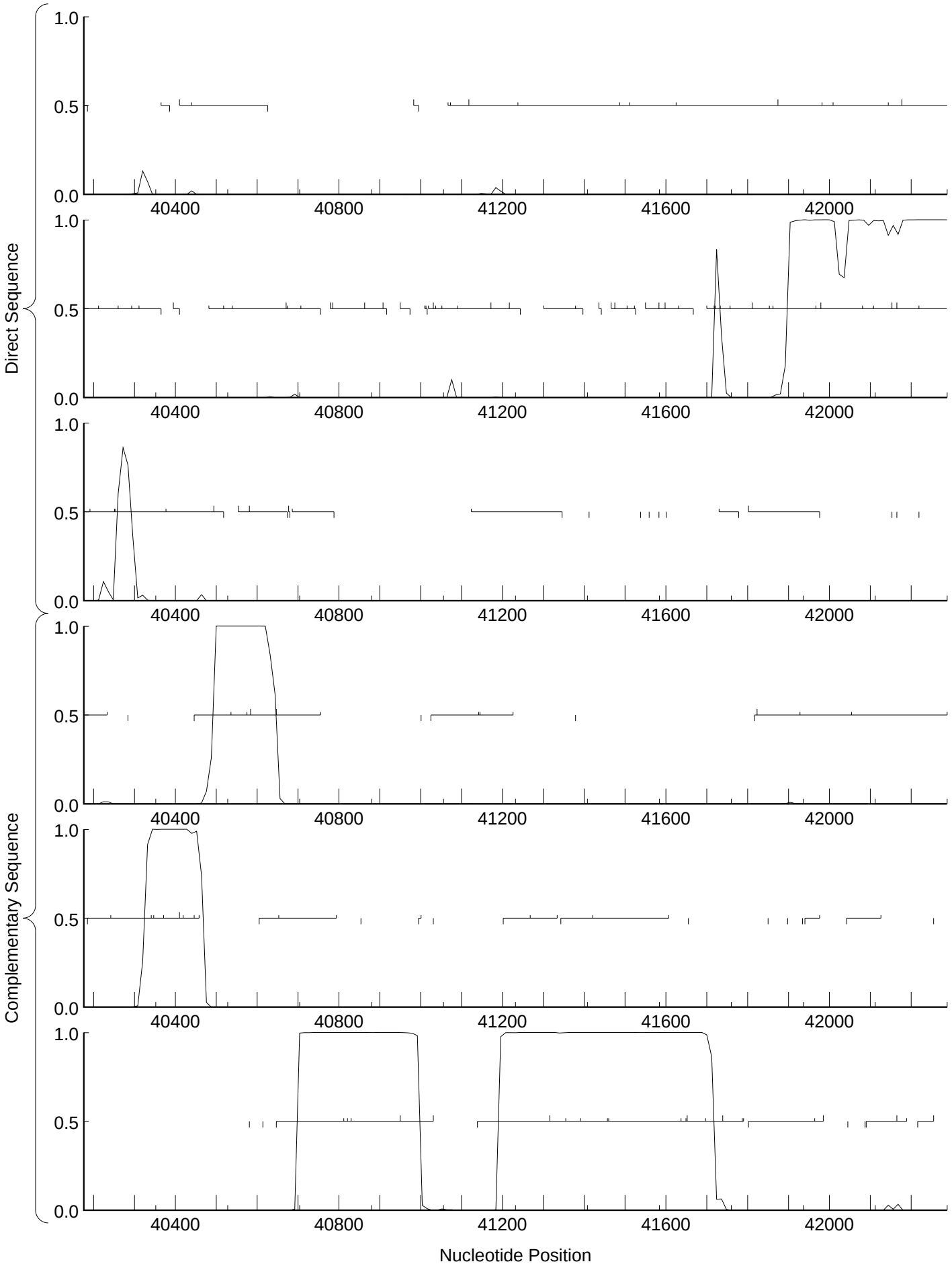


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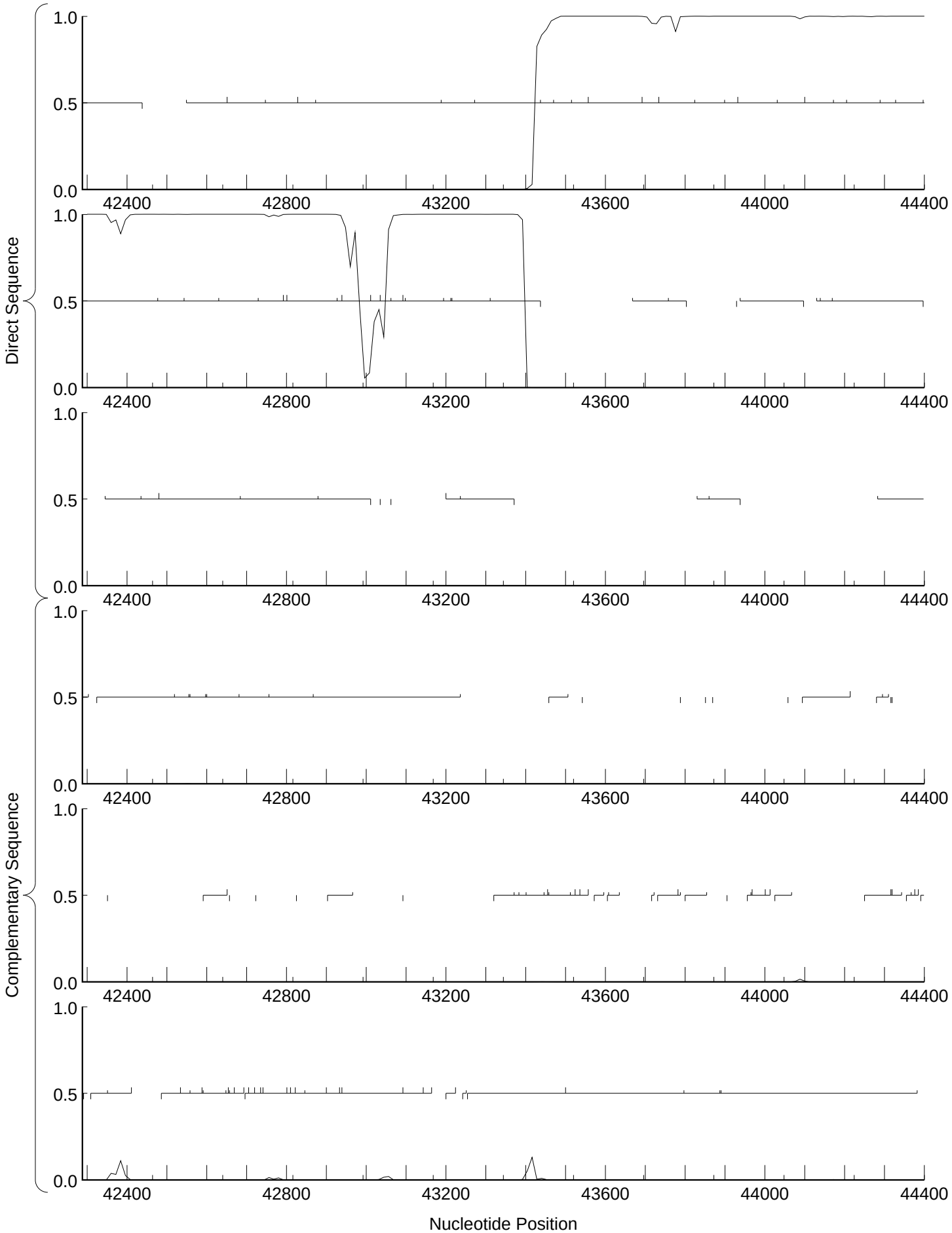



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

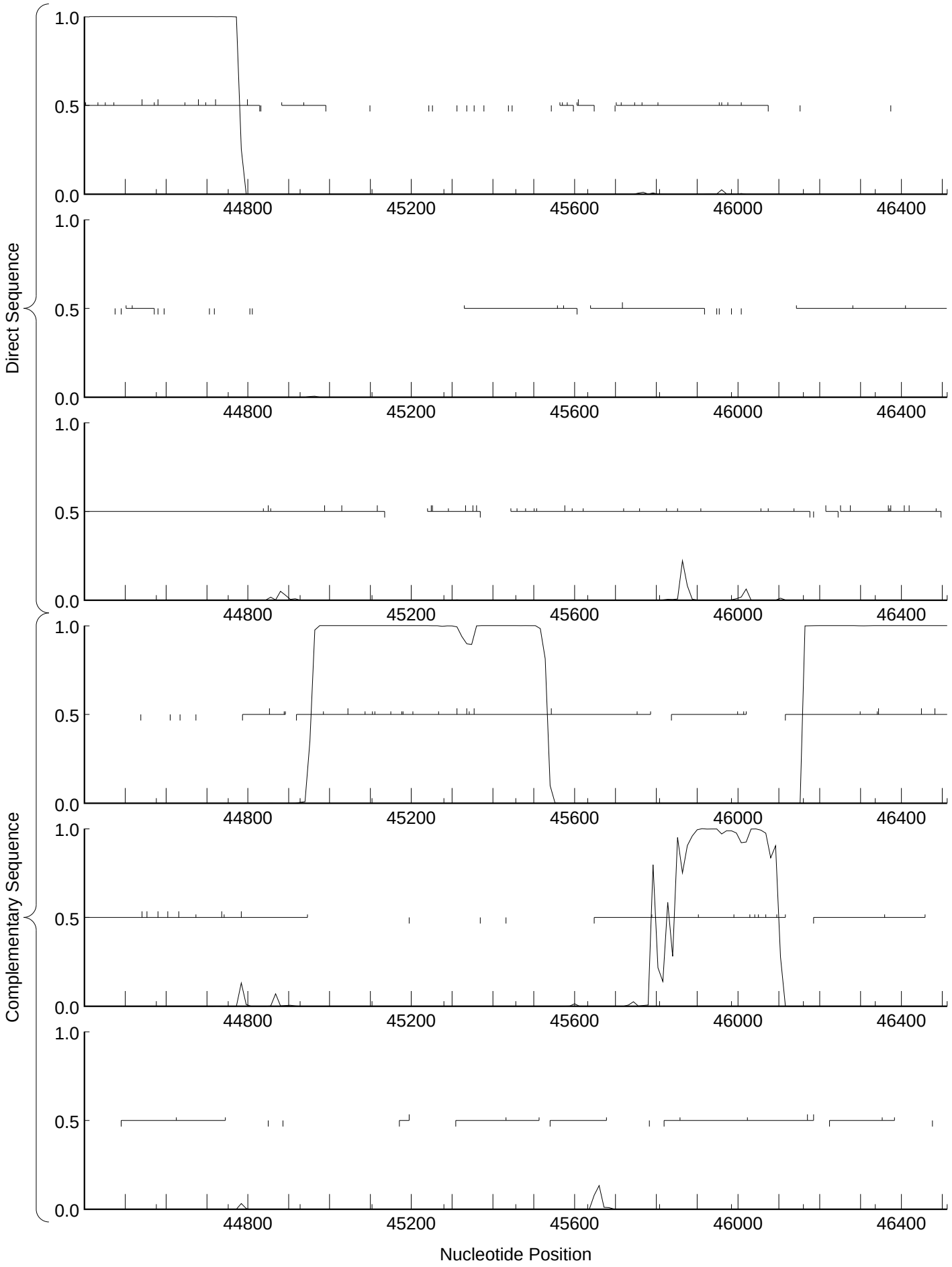
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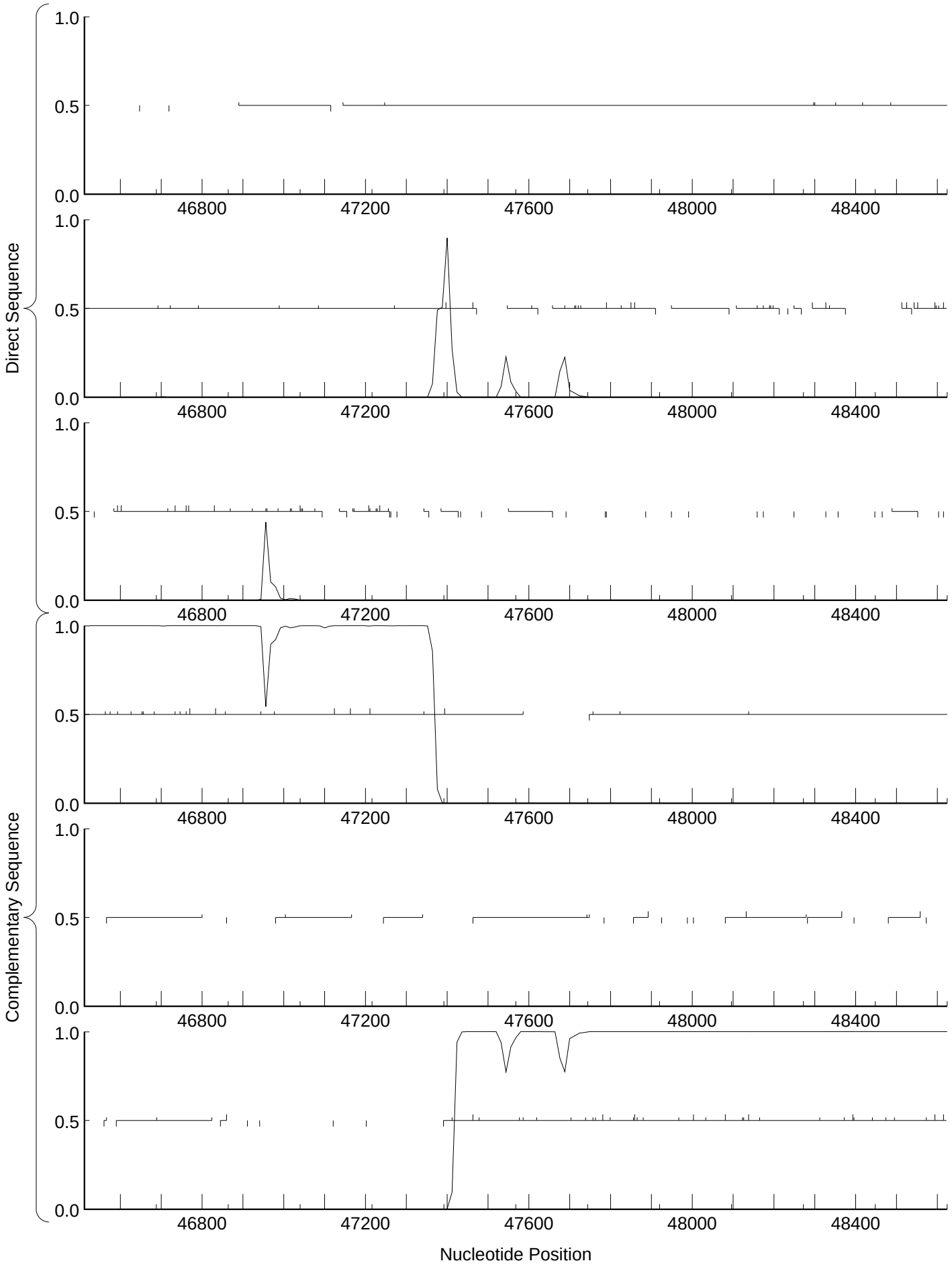
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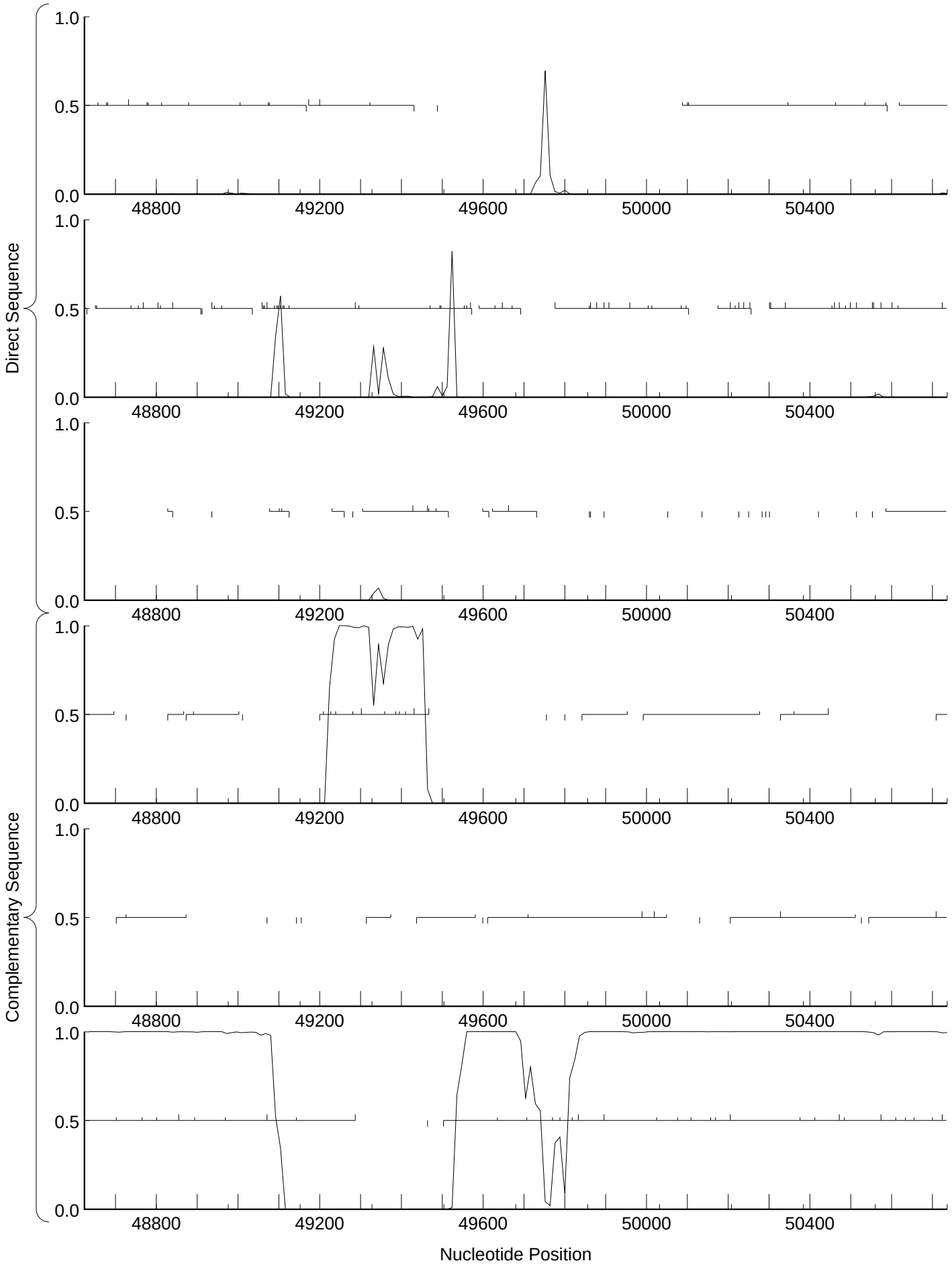
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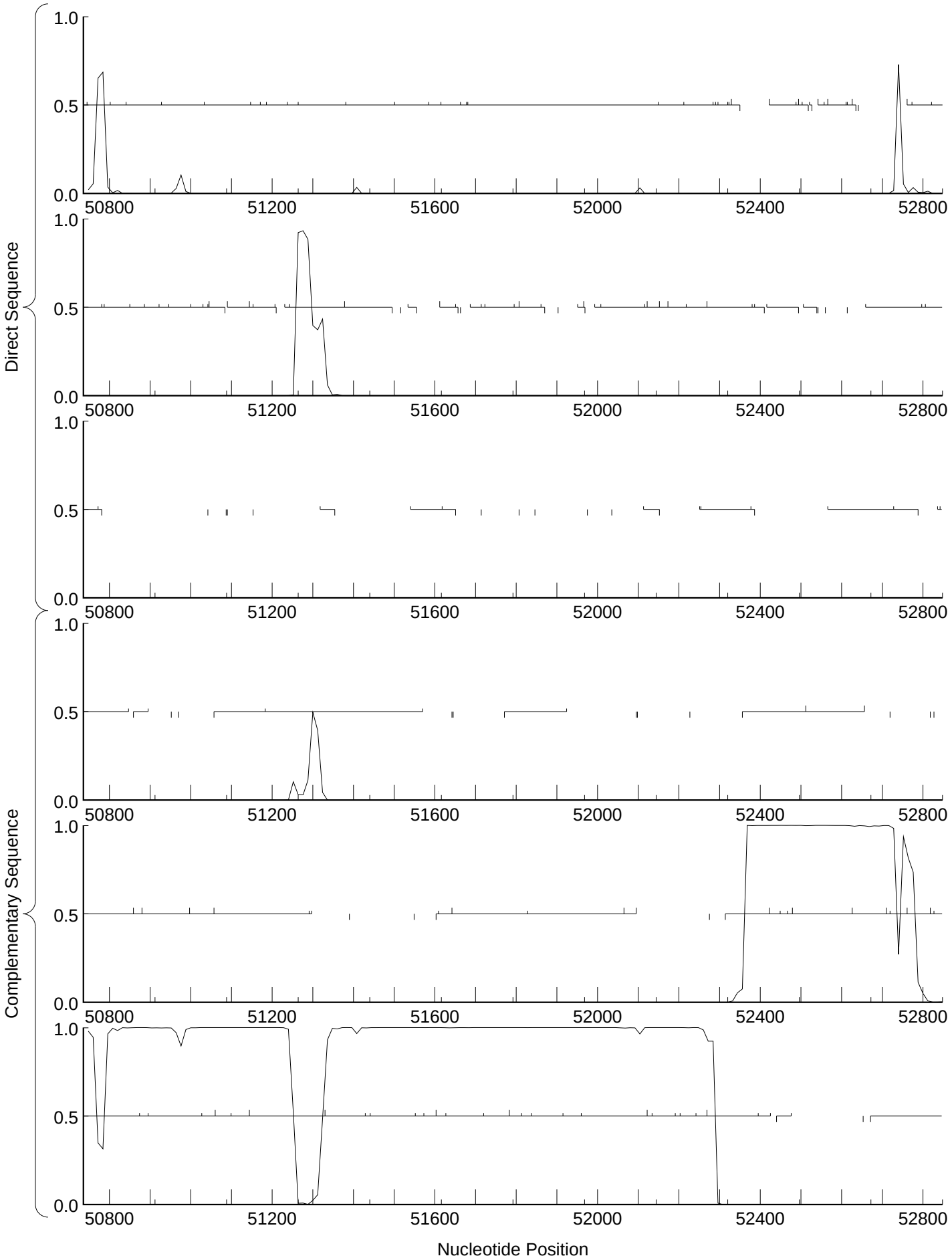
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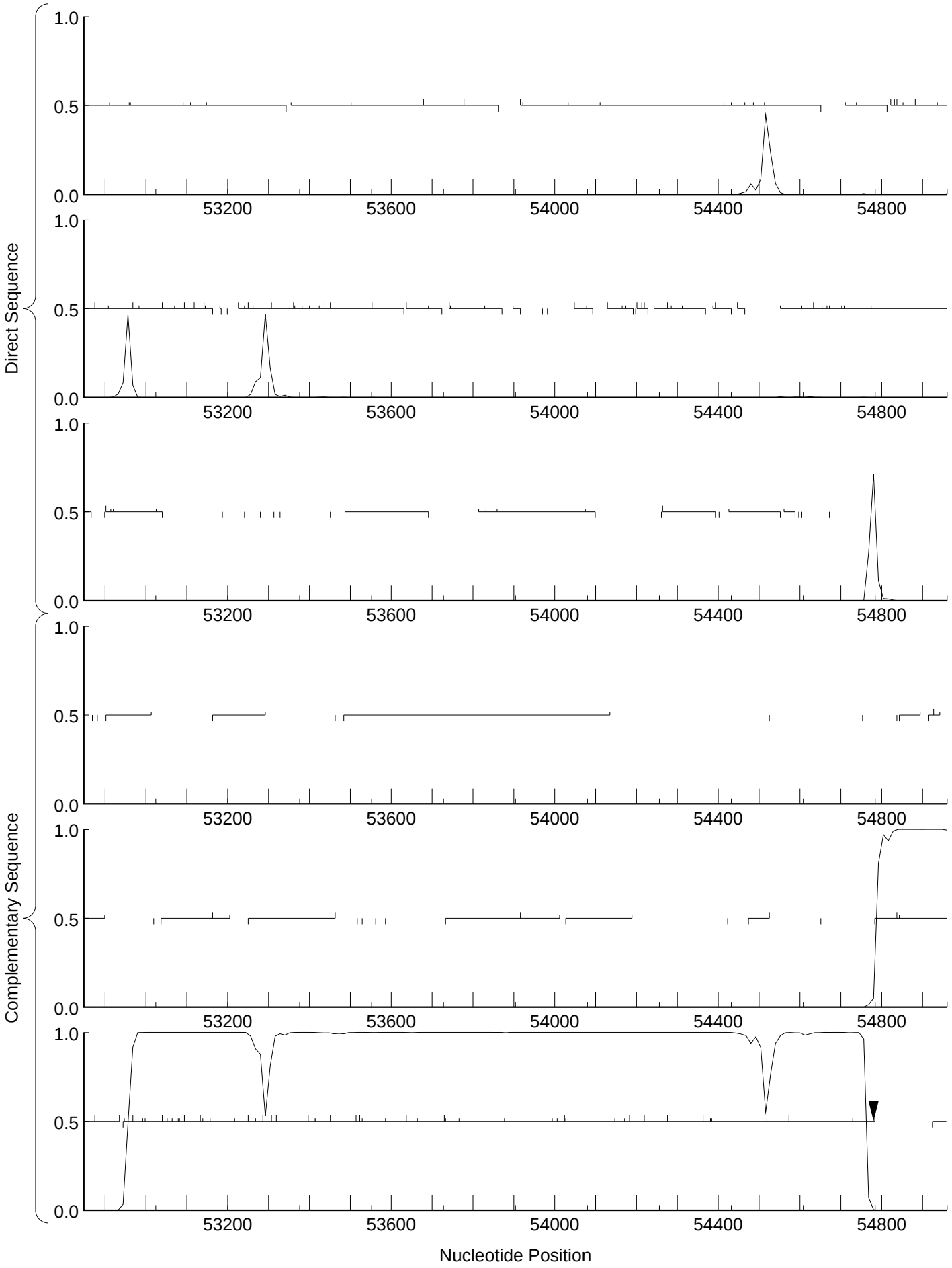
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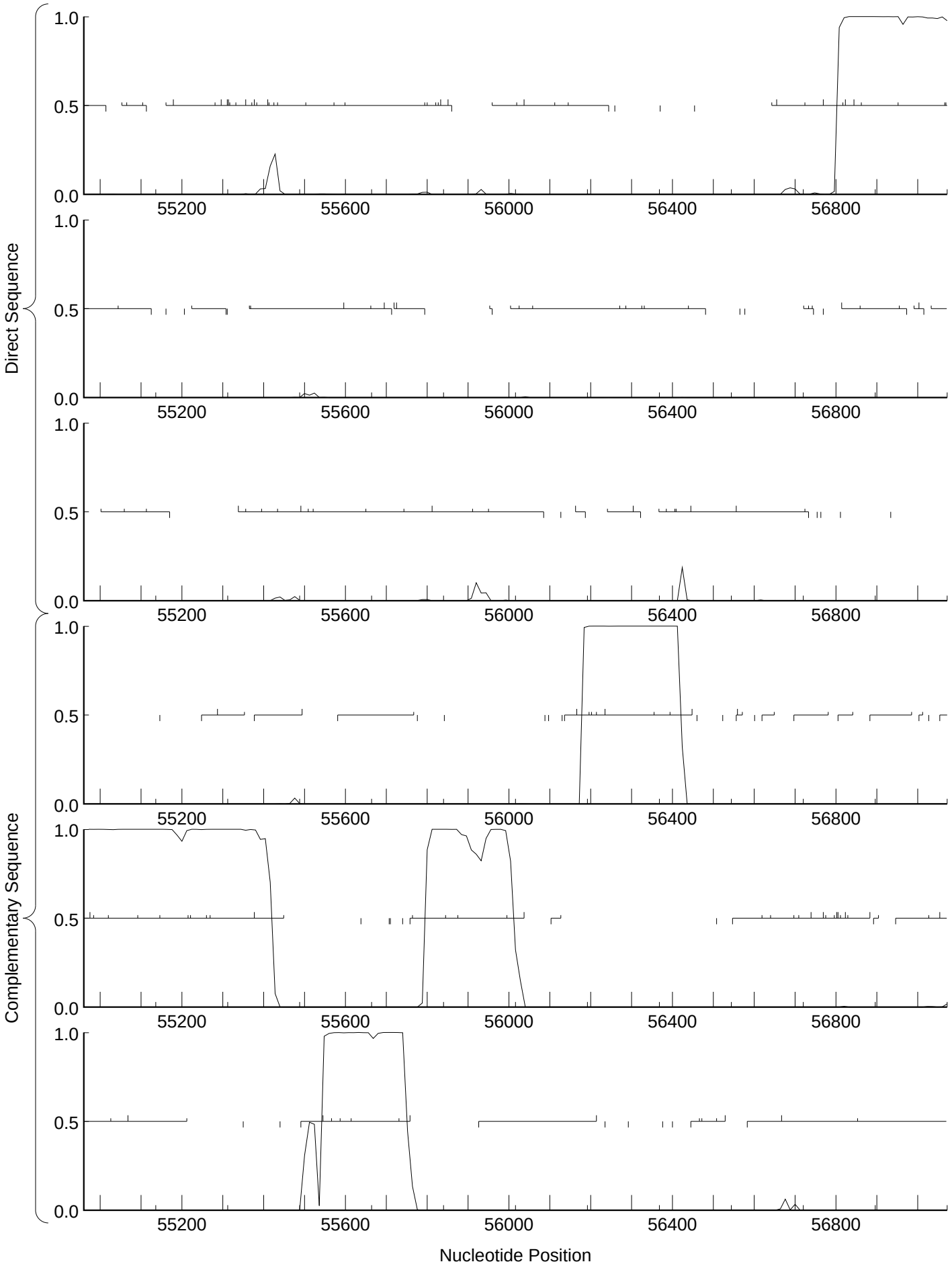
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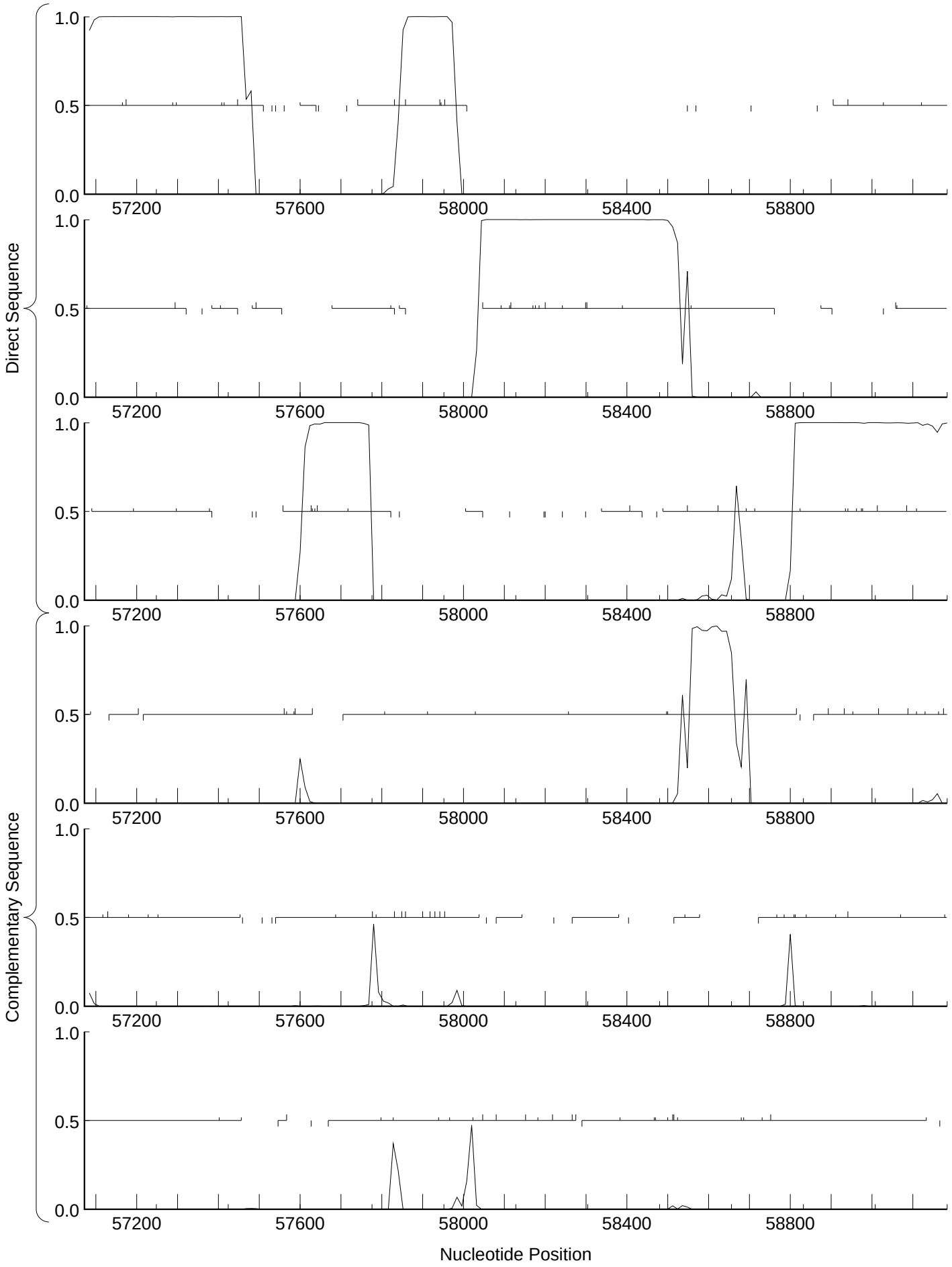
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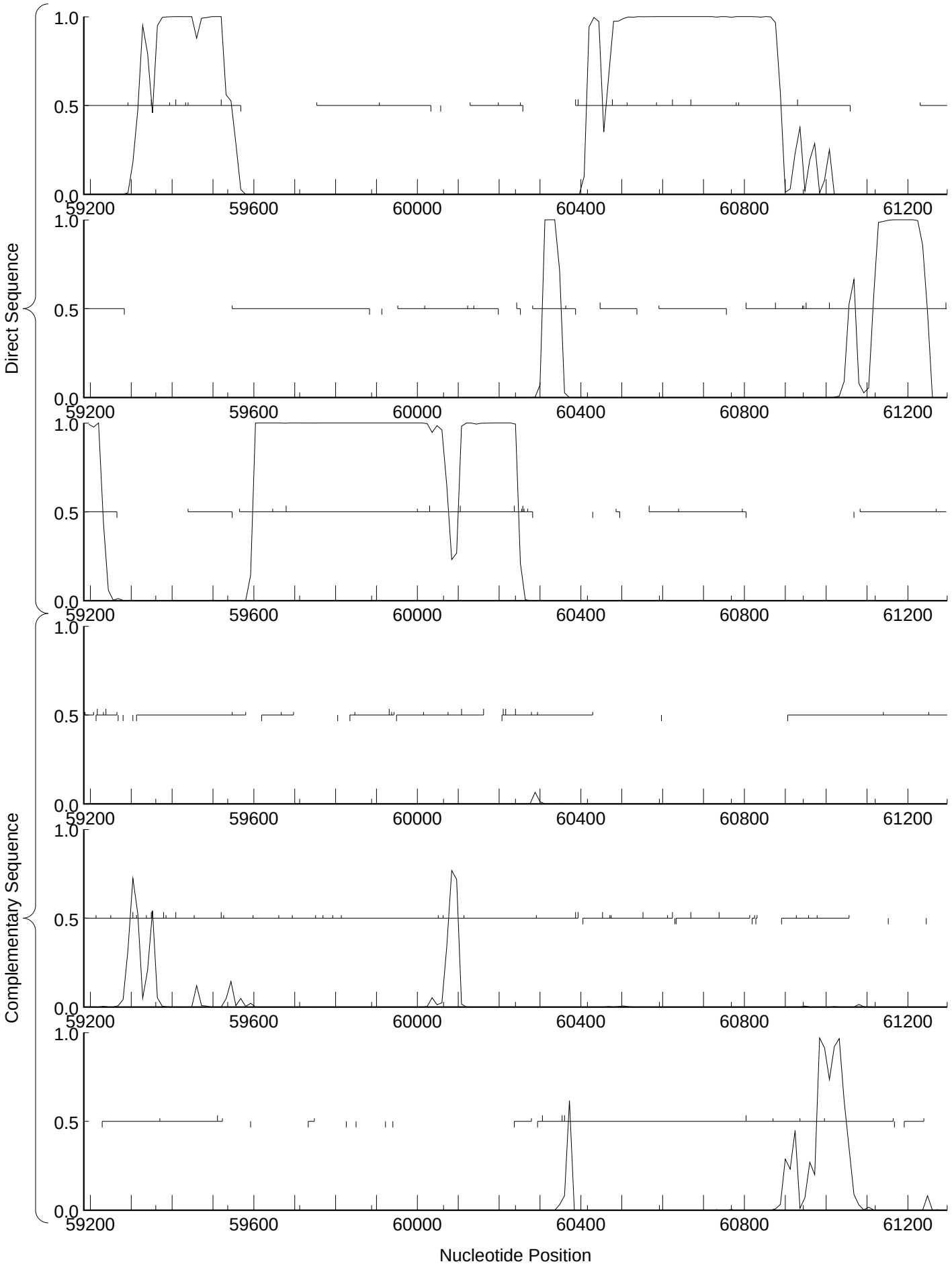
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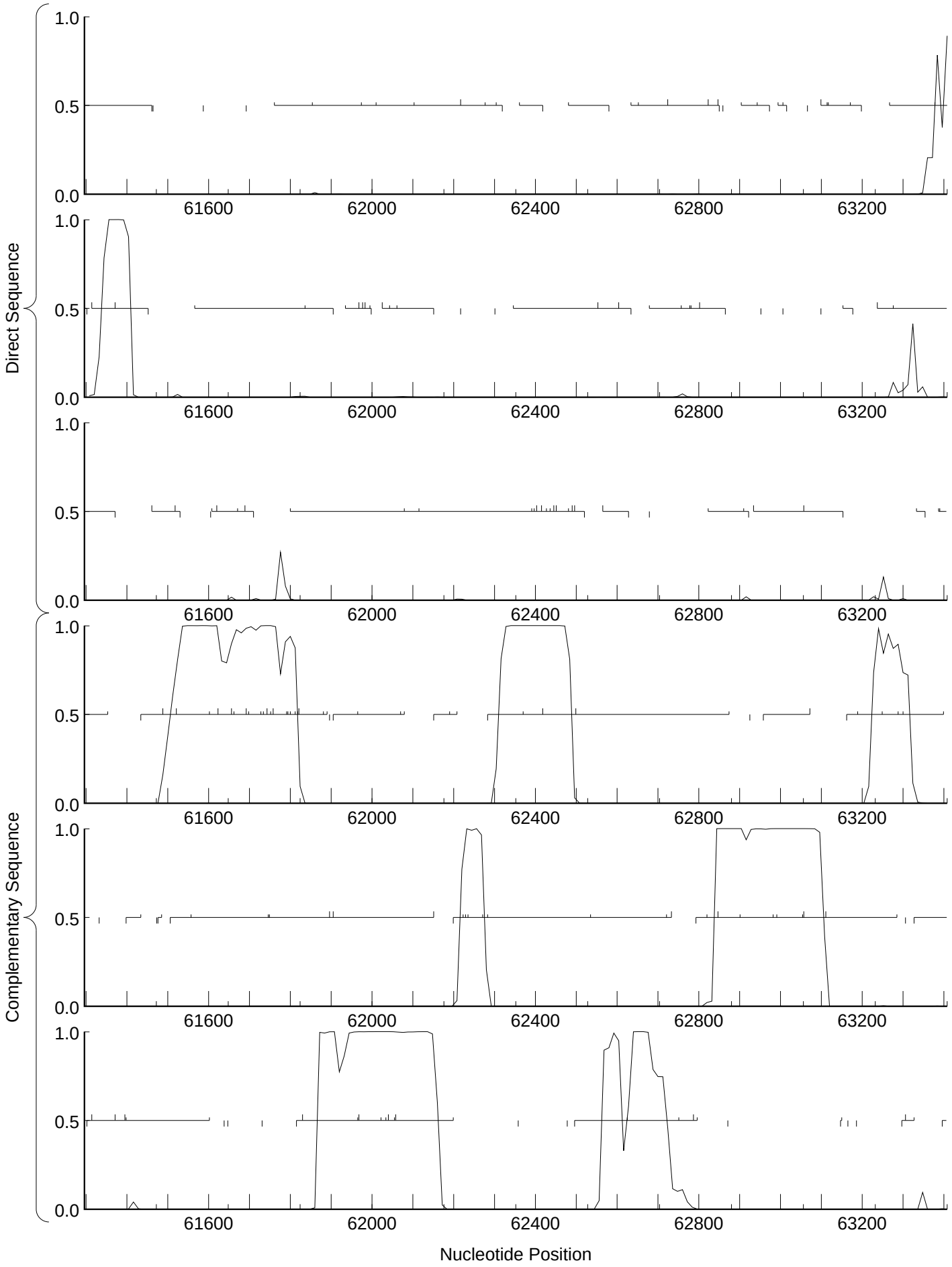
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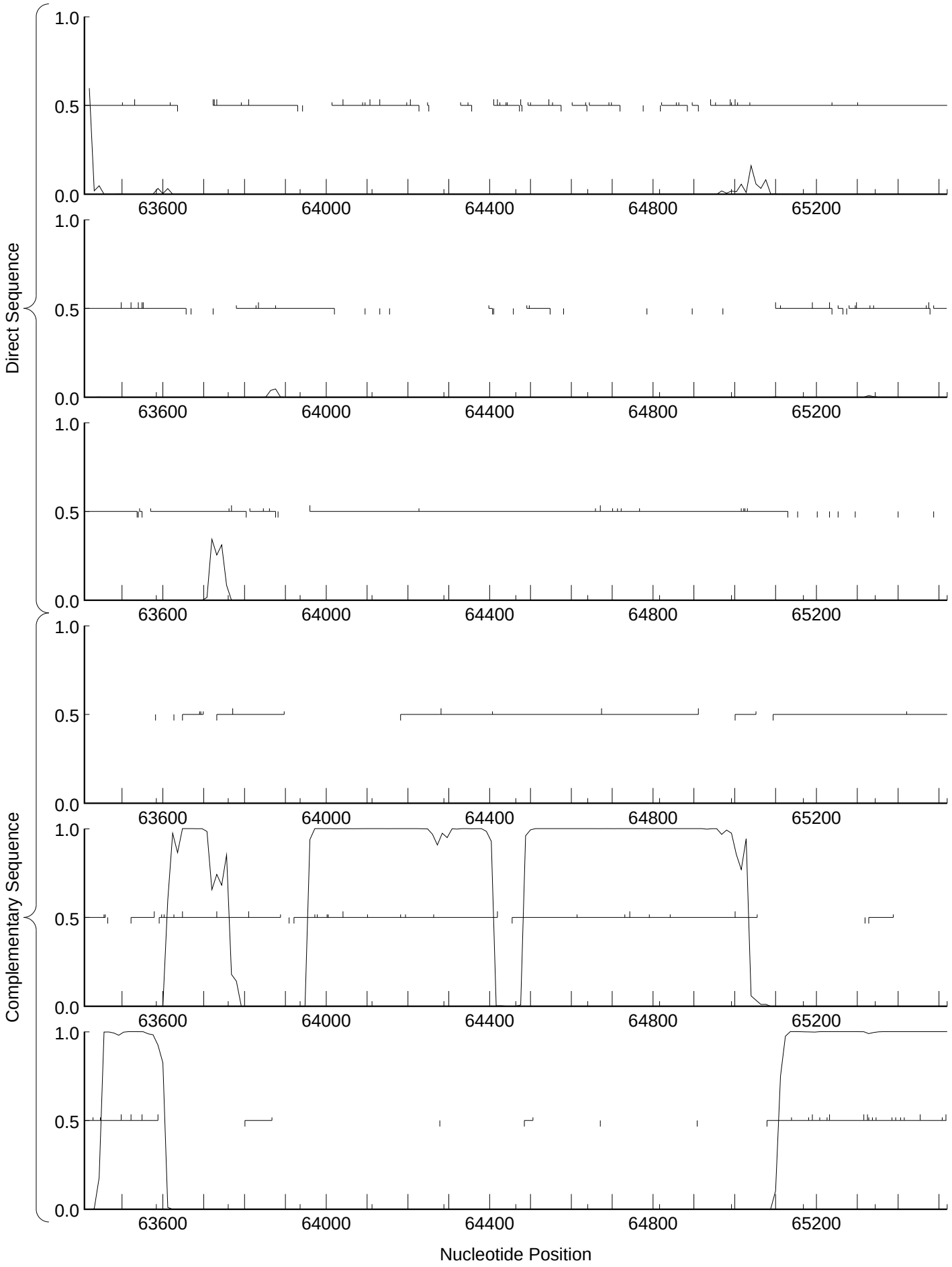
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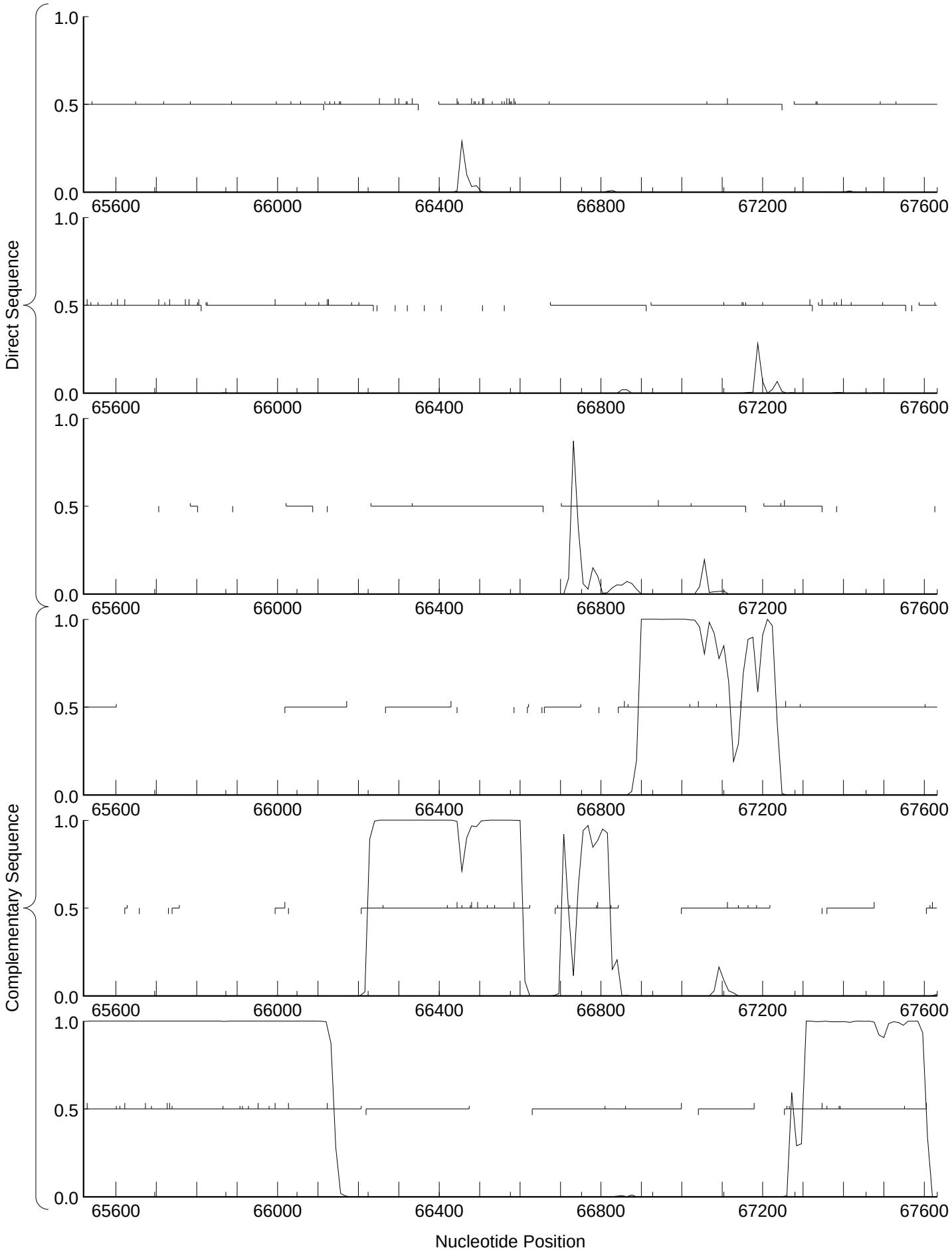
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ycobacterium phage Saguaro complete sequence, 69448 bp, circularly permuted, Cluster B7, Order 4, Window 96, Step 12, 32/34



ycobacterium phage Saguaro complete sequence, 69448 bp, circularly permuted, Cluster B7, Order 4, Window 96, Step 12, 33/34



ycobacterium phage Saguaro complete sequence, 69448 bp, circularly permuted, Cluster B7, Order 4, Window 96, Step 12, 34/34

