GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage ShiVal Complete Sequence, 68355 bp, Circularly permuted, Cluster B1
Analysis Date    :  3/28/24 at 18:16:02
Pages            :  34
Sequence Length  :  68355 bp
GC Content       :   66.50%

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

PS-Version       :  1.2

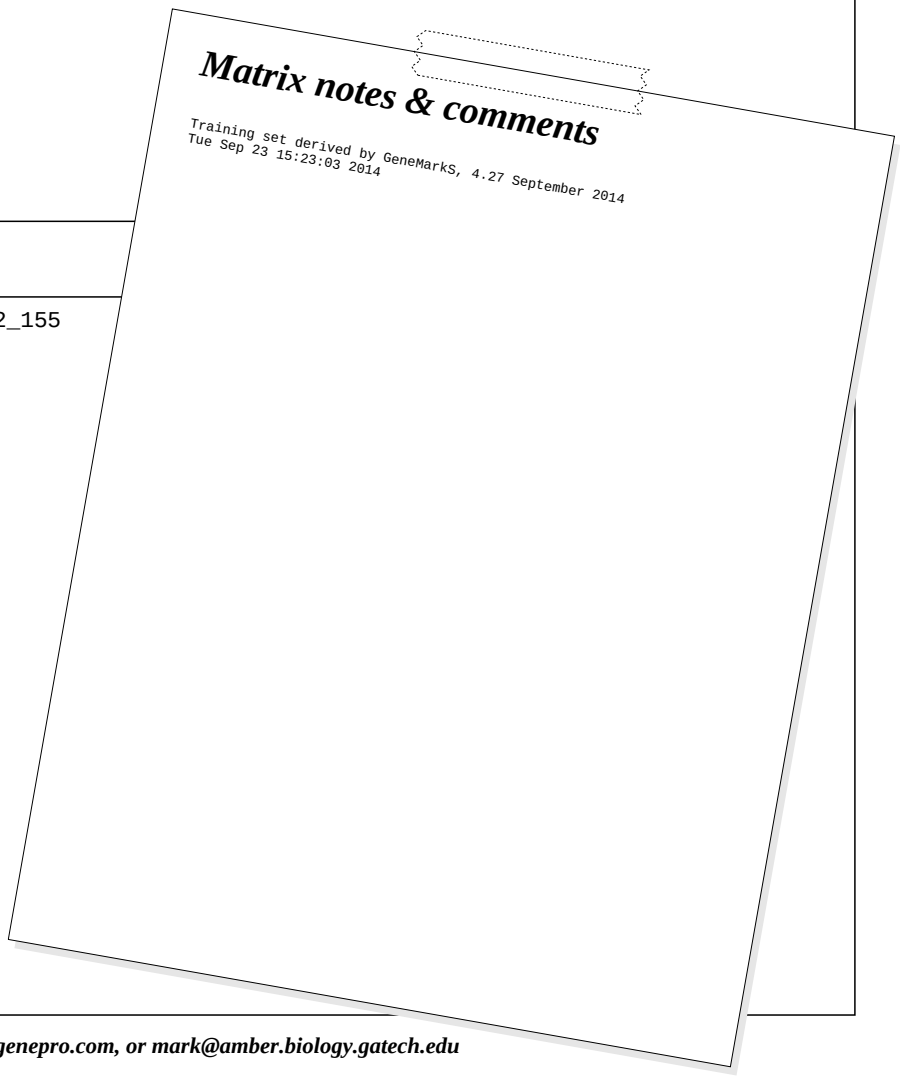
GeneMark Options :  PostScript graph,
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MATRIX INFORMATION

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Author     :  -

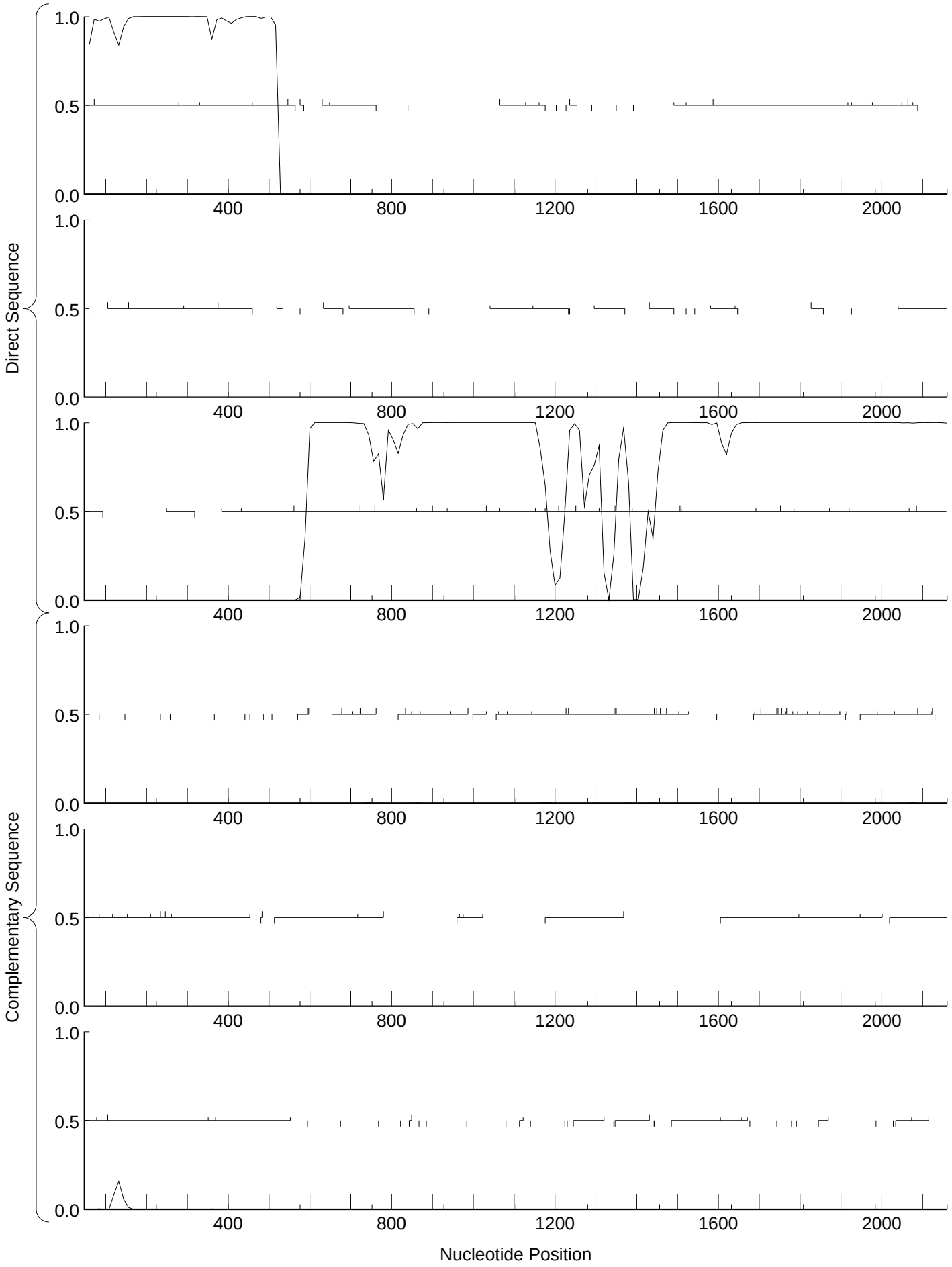
Order      :  4


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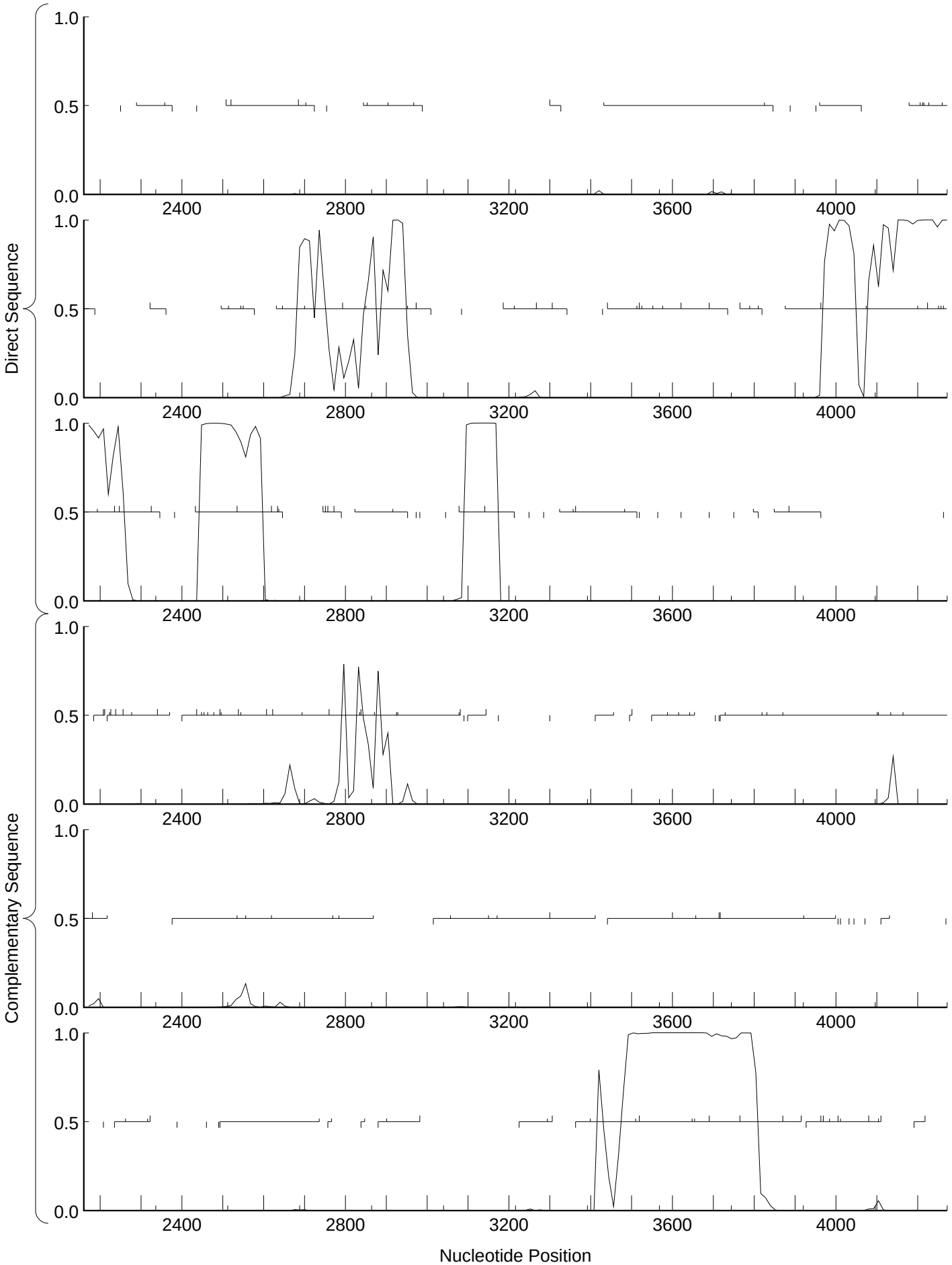


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

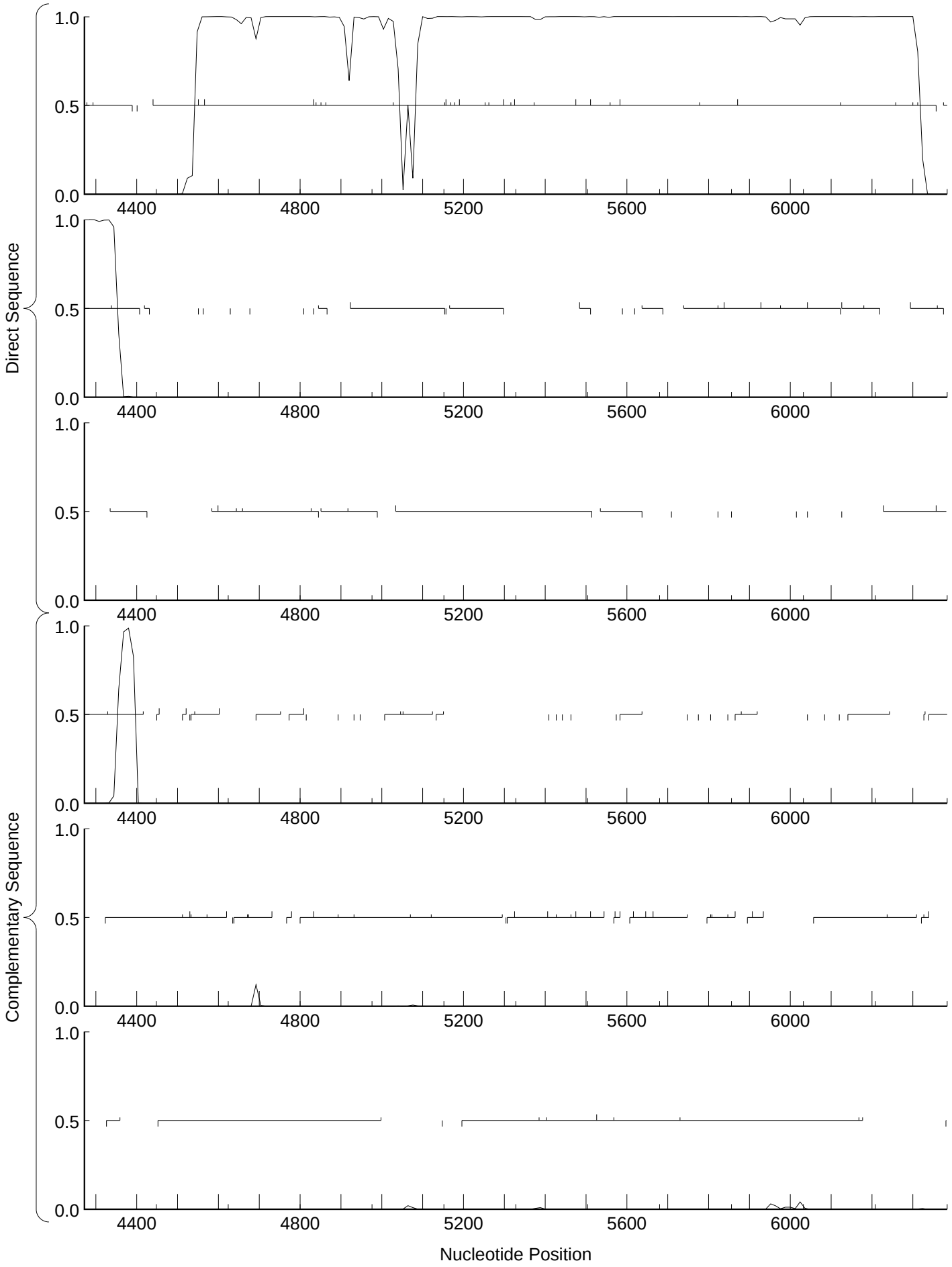
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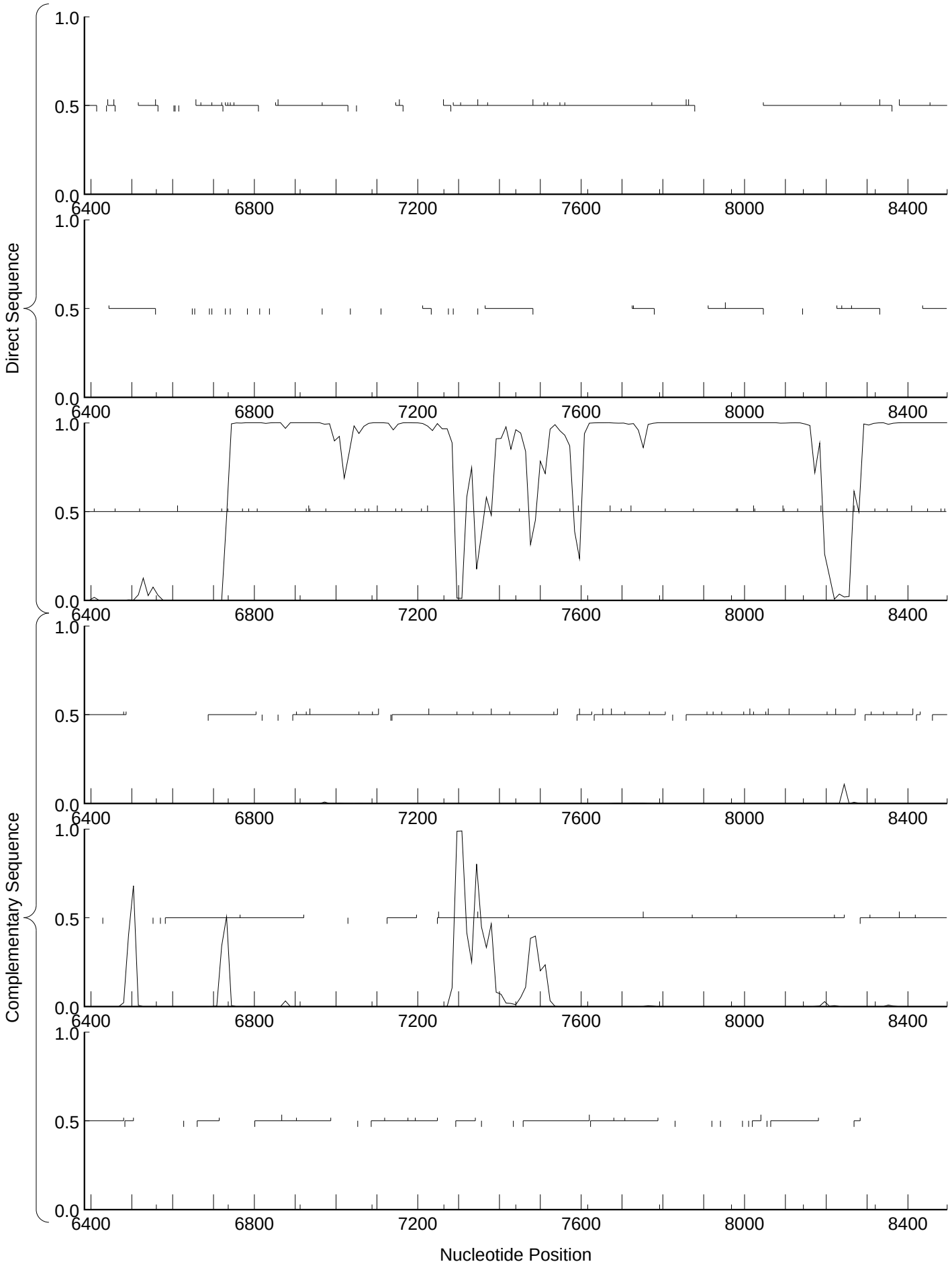
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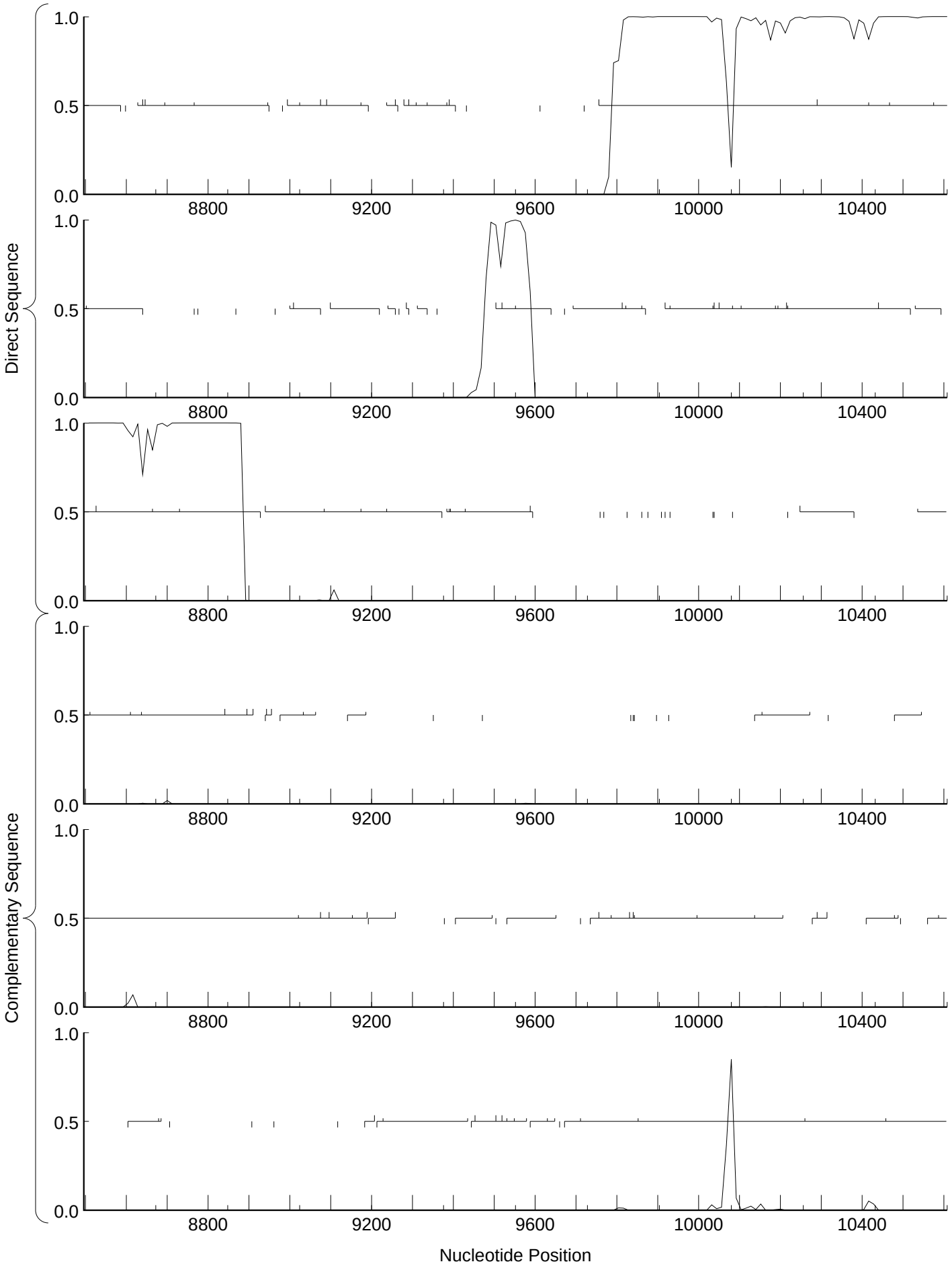
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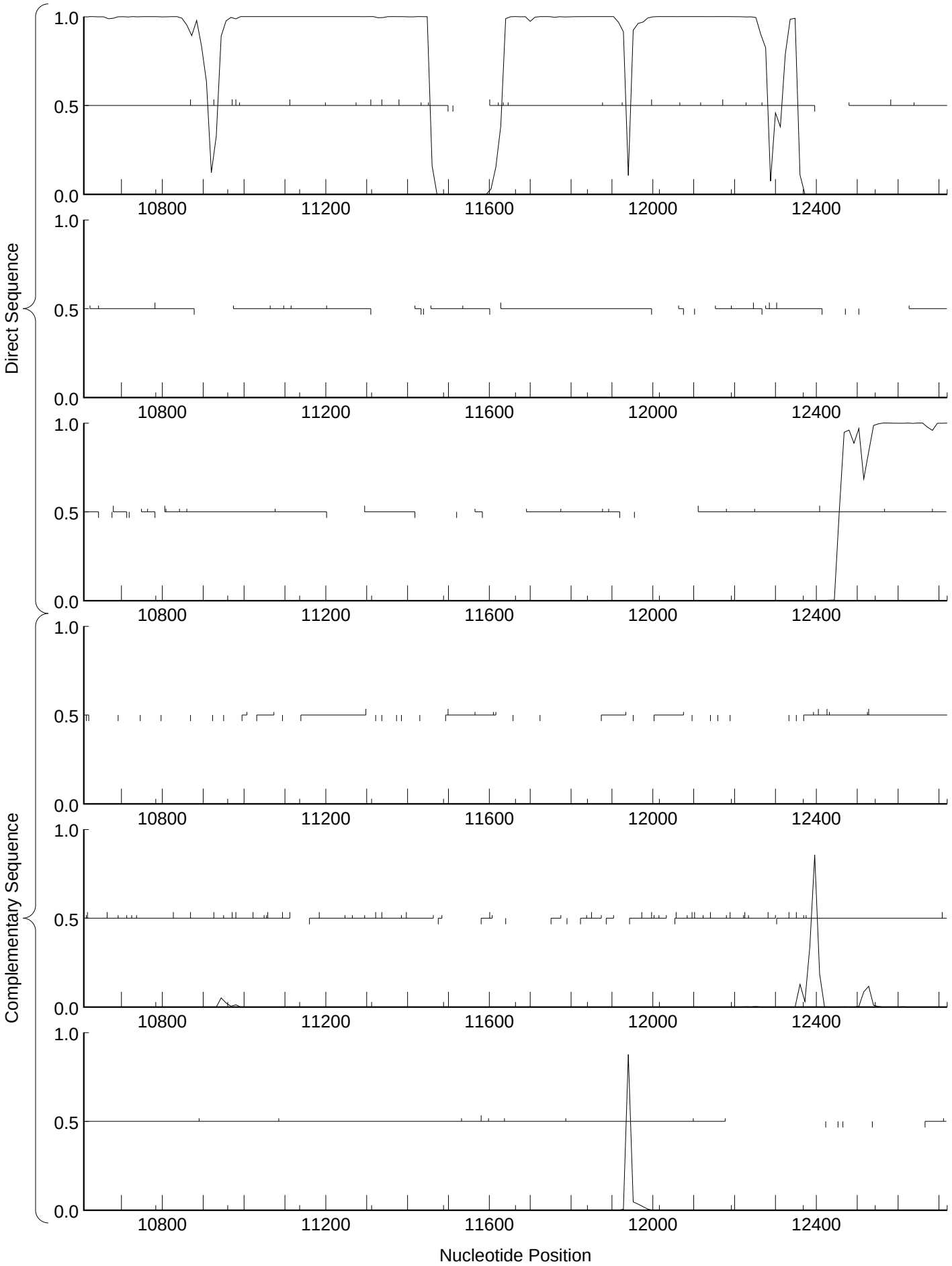
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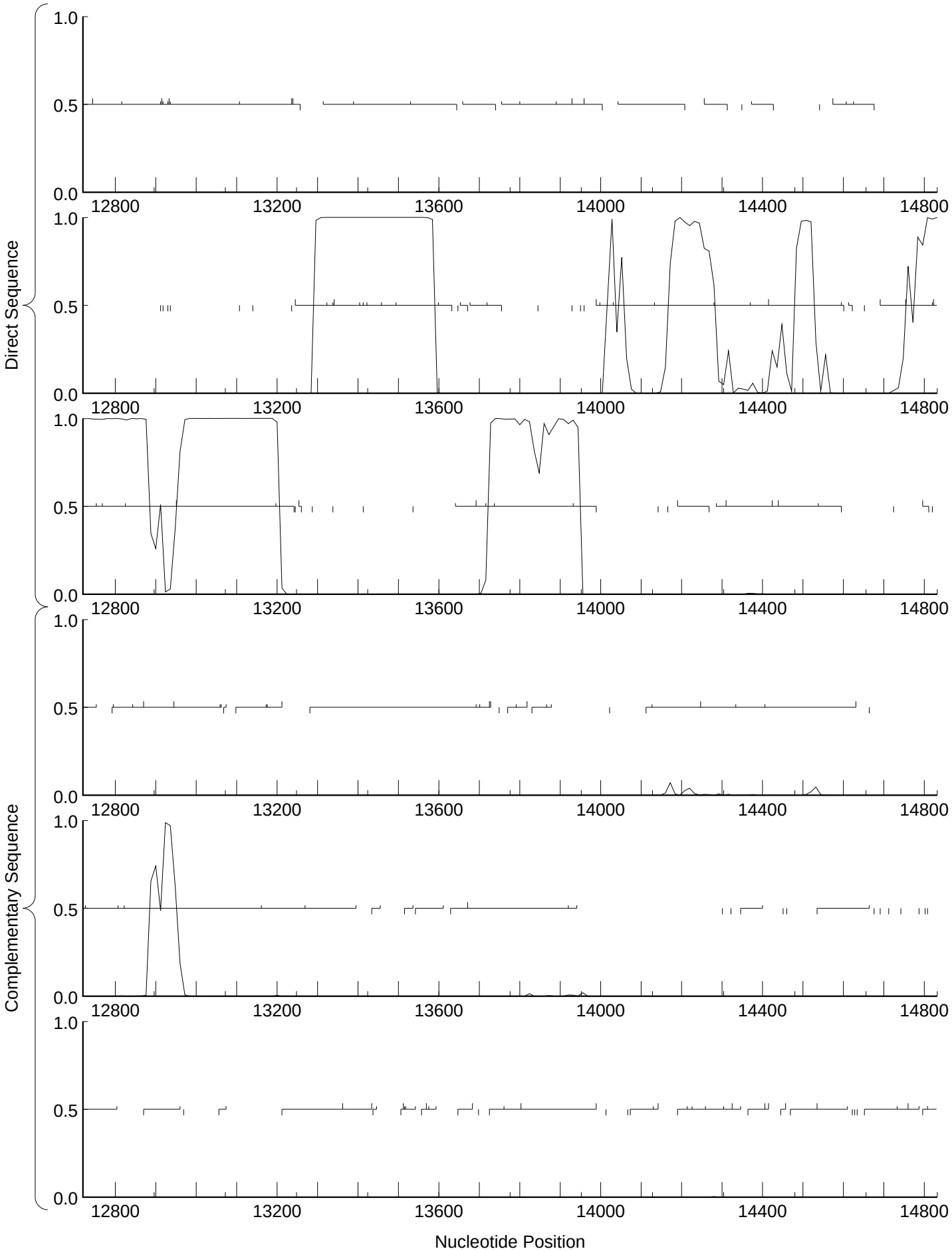
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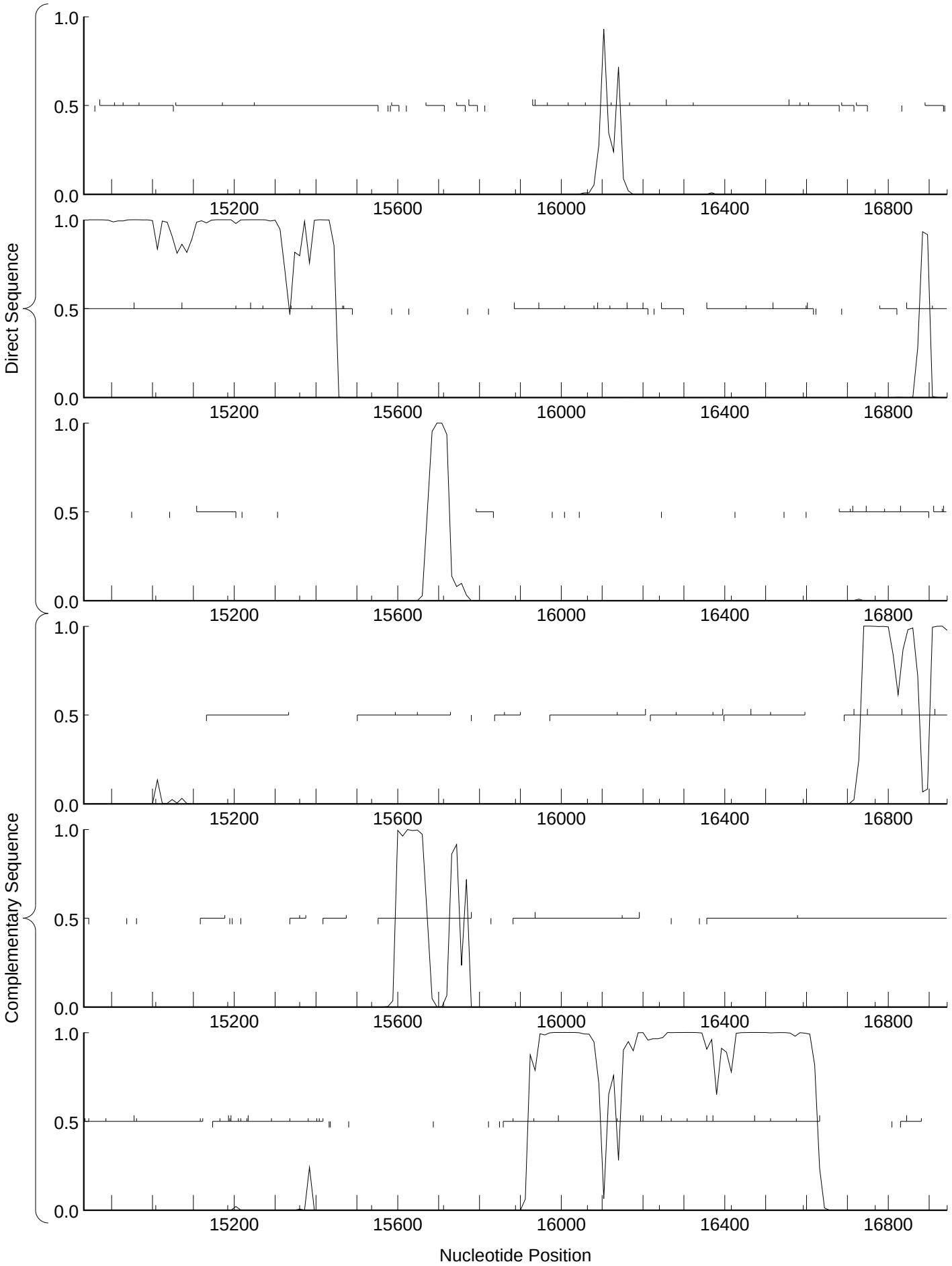
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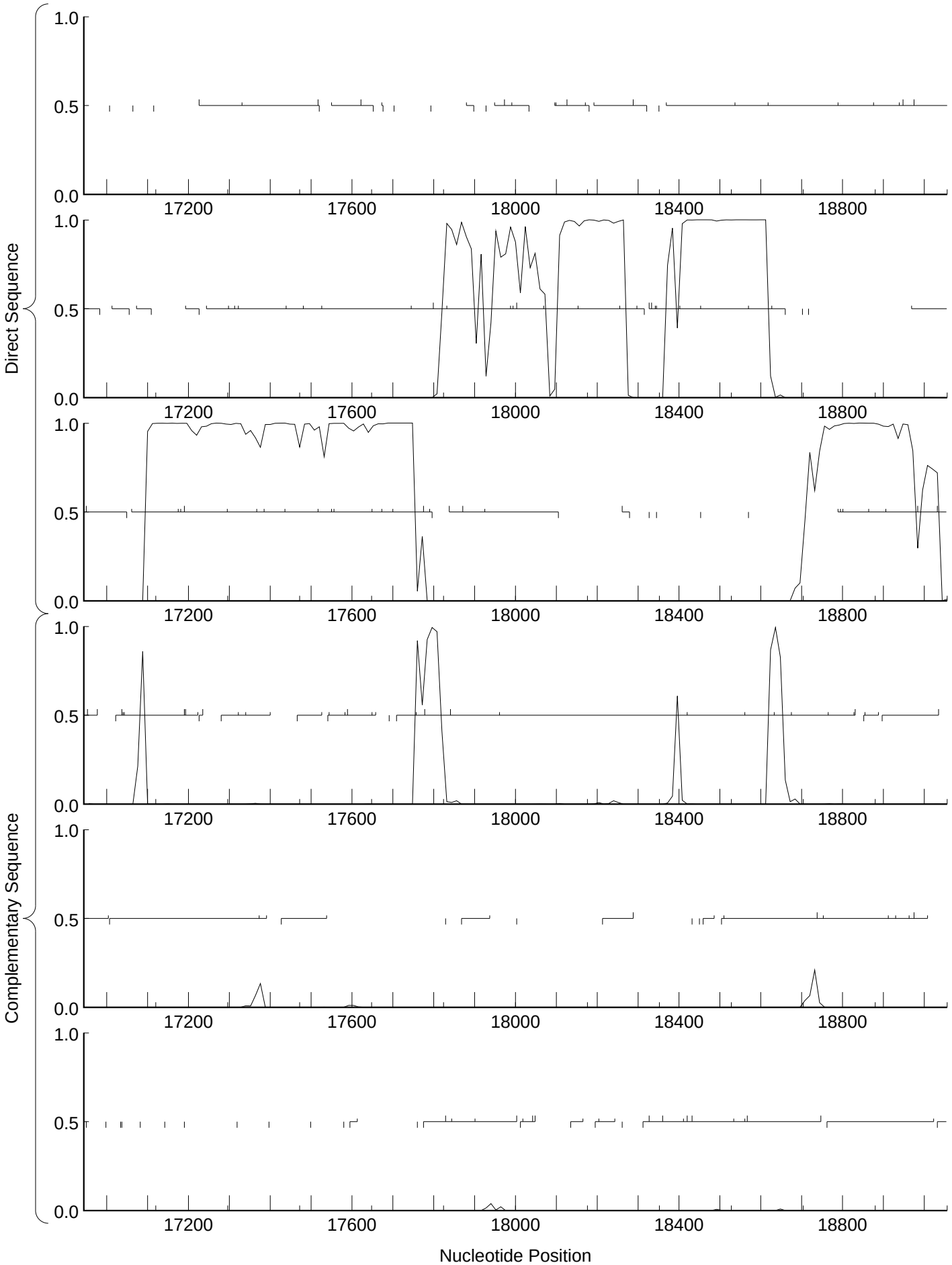
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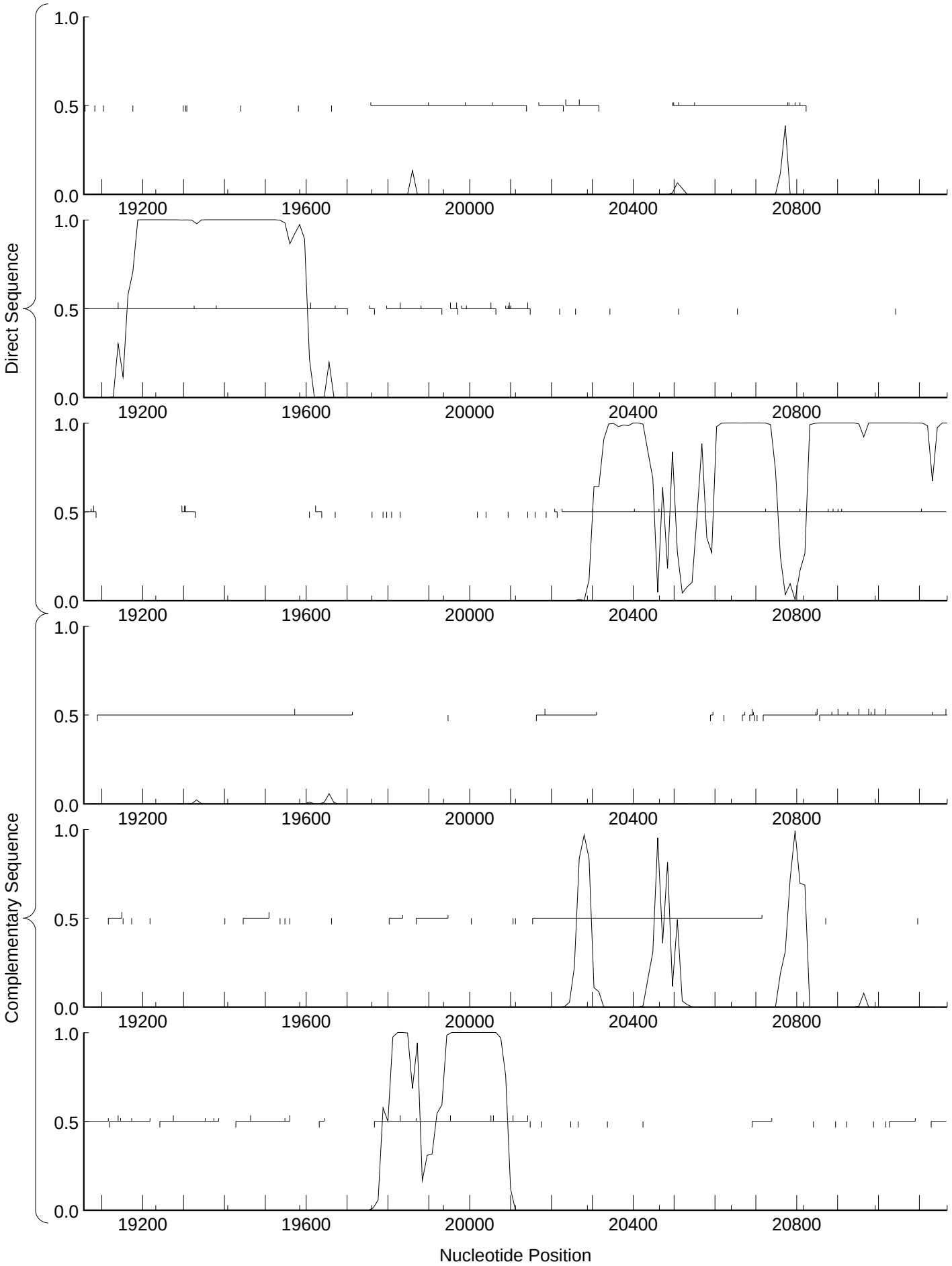
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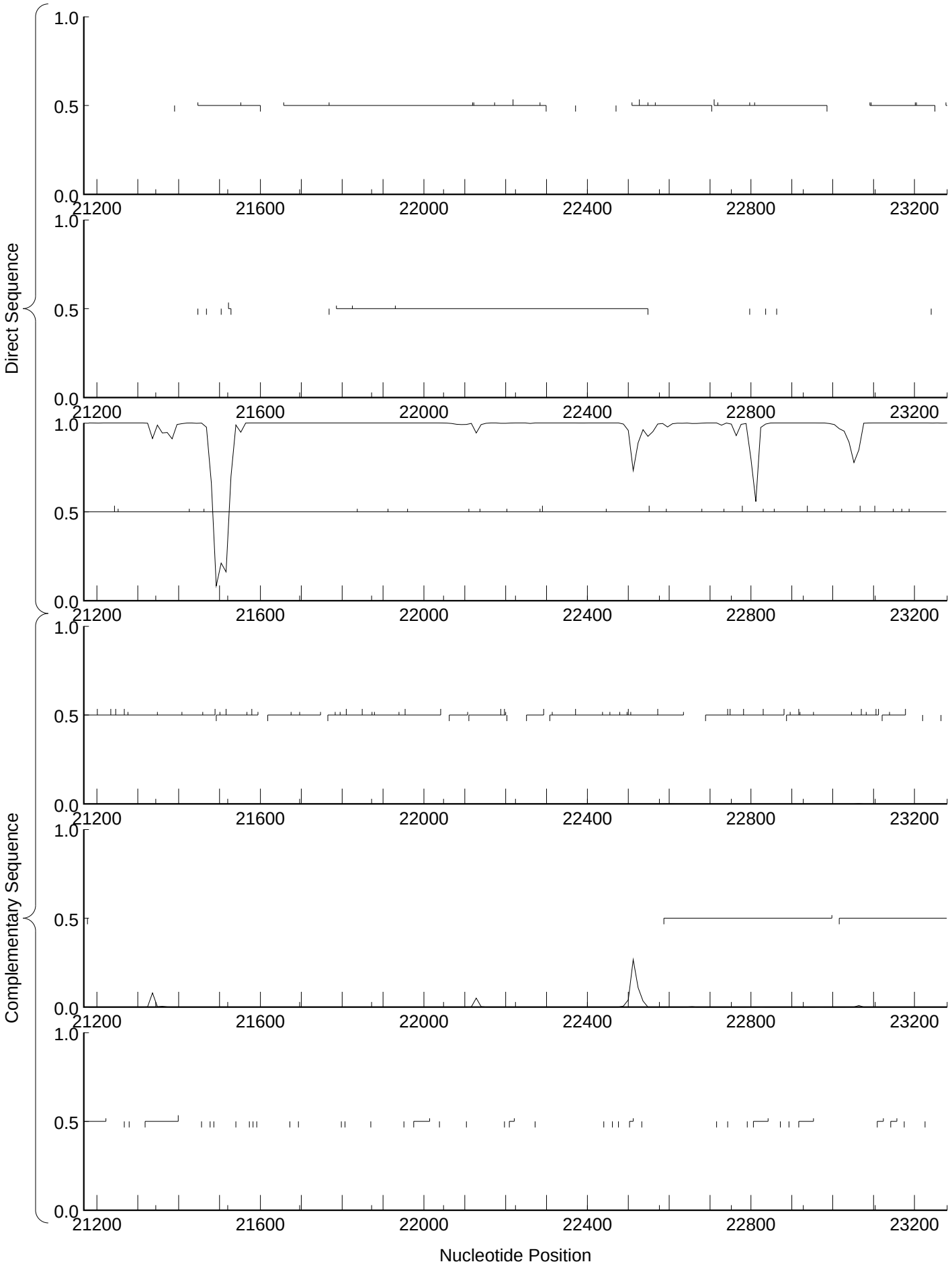
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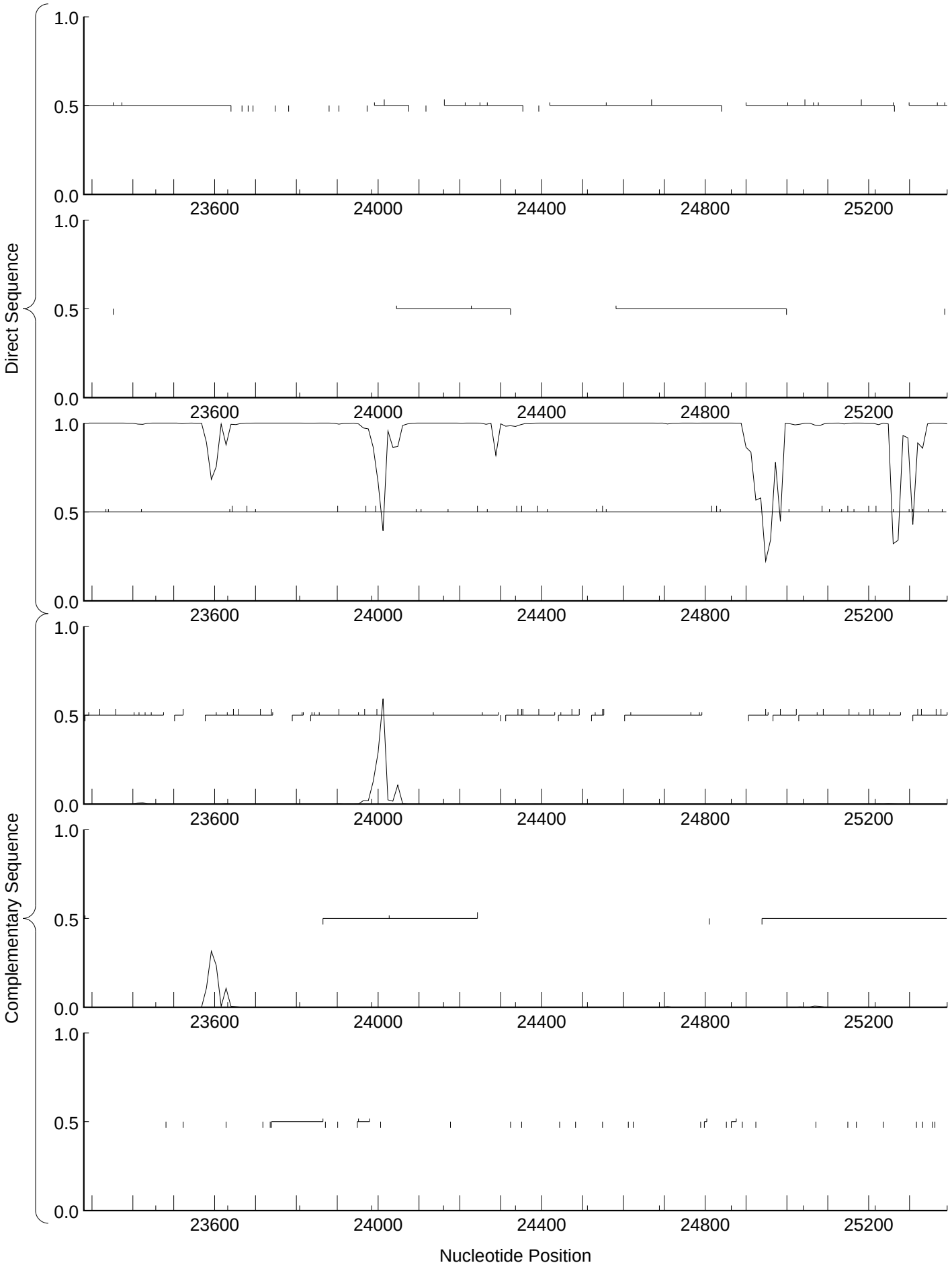
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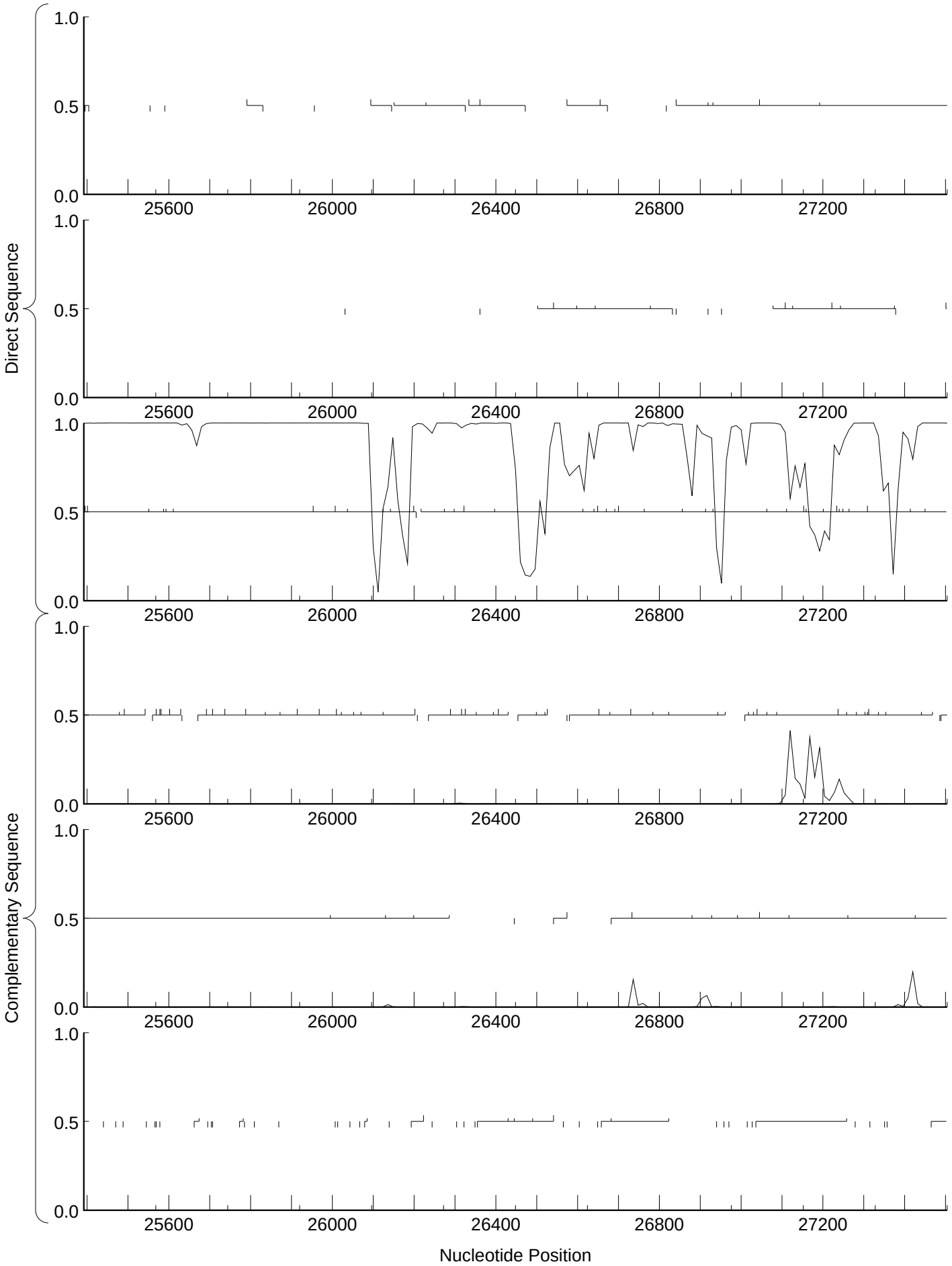
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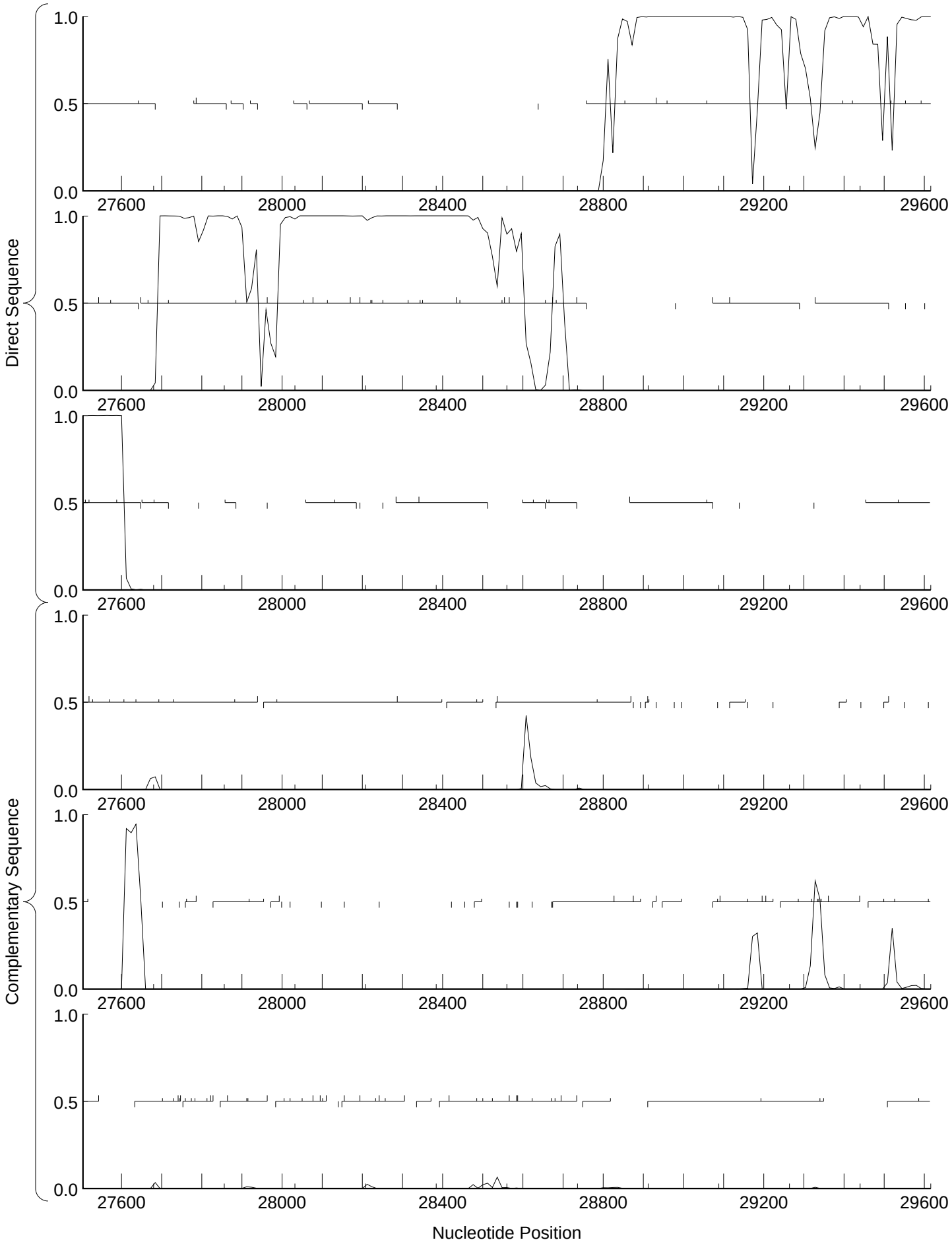
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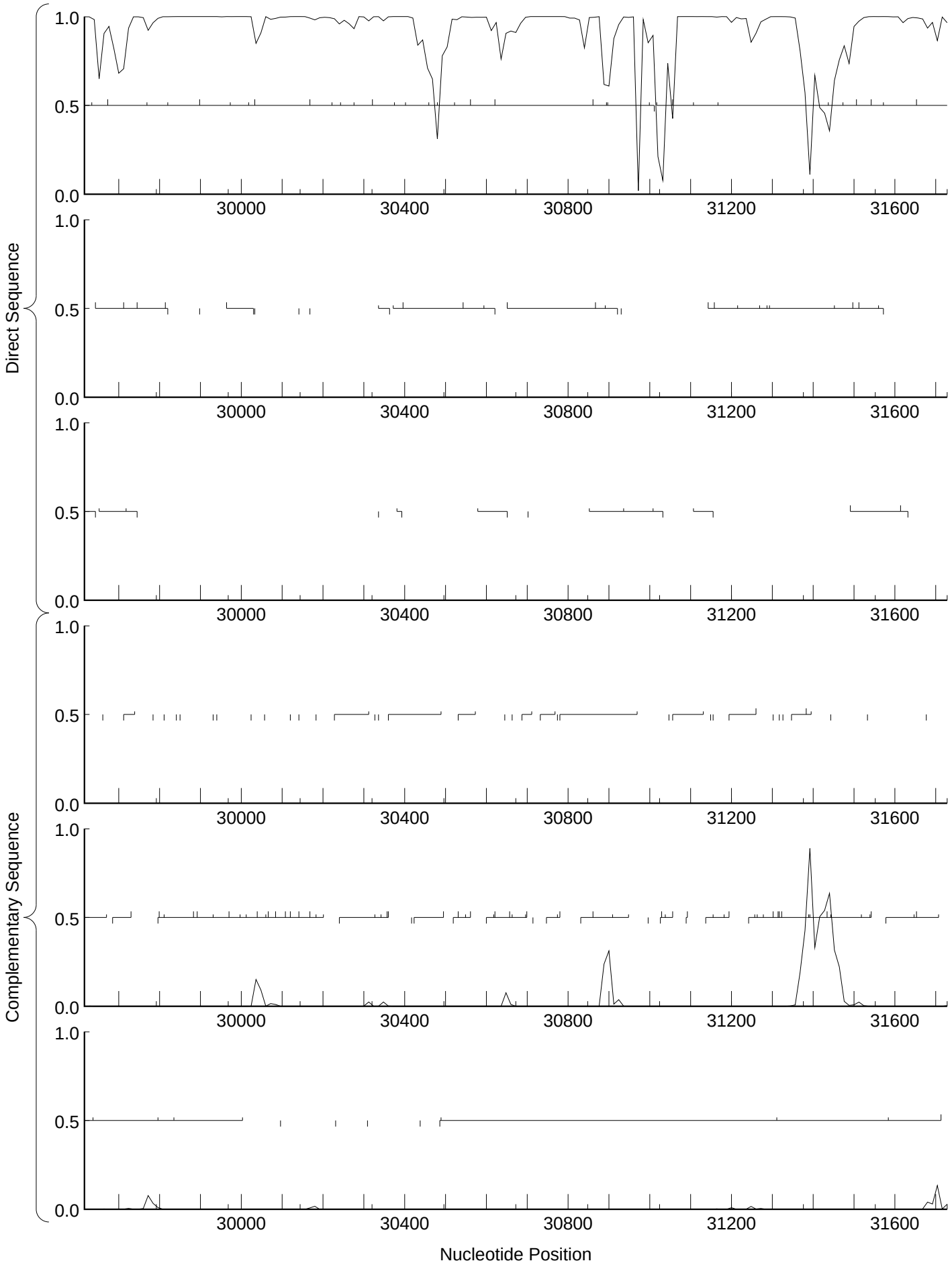
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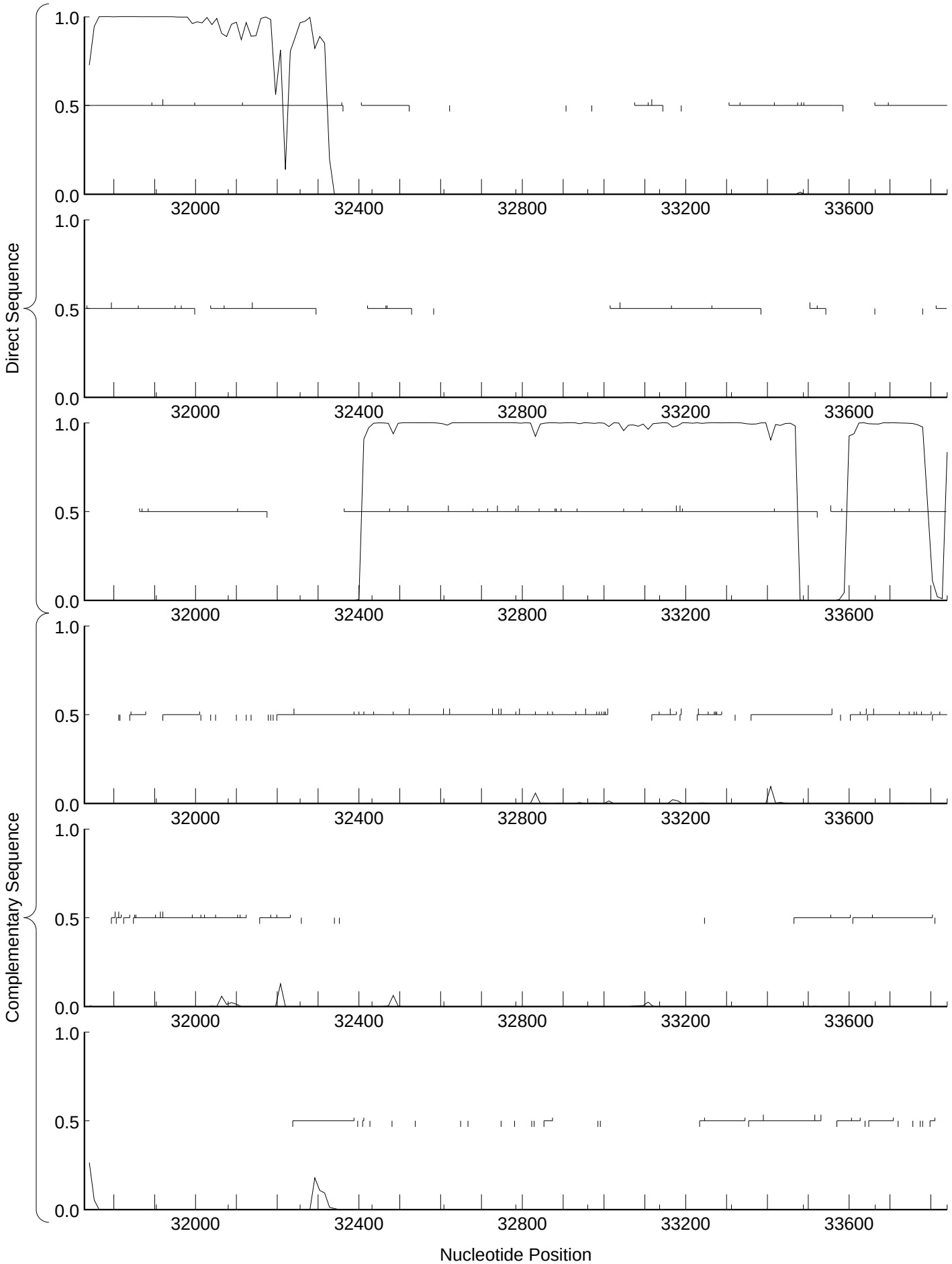
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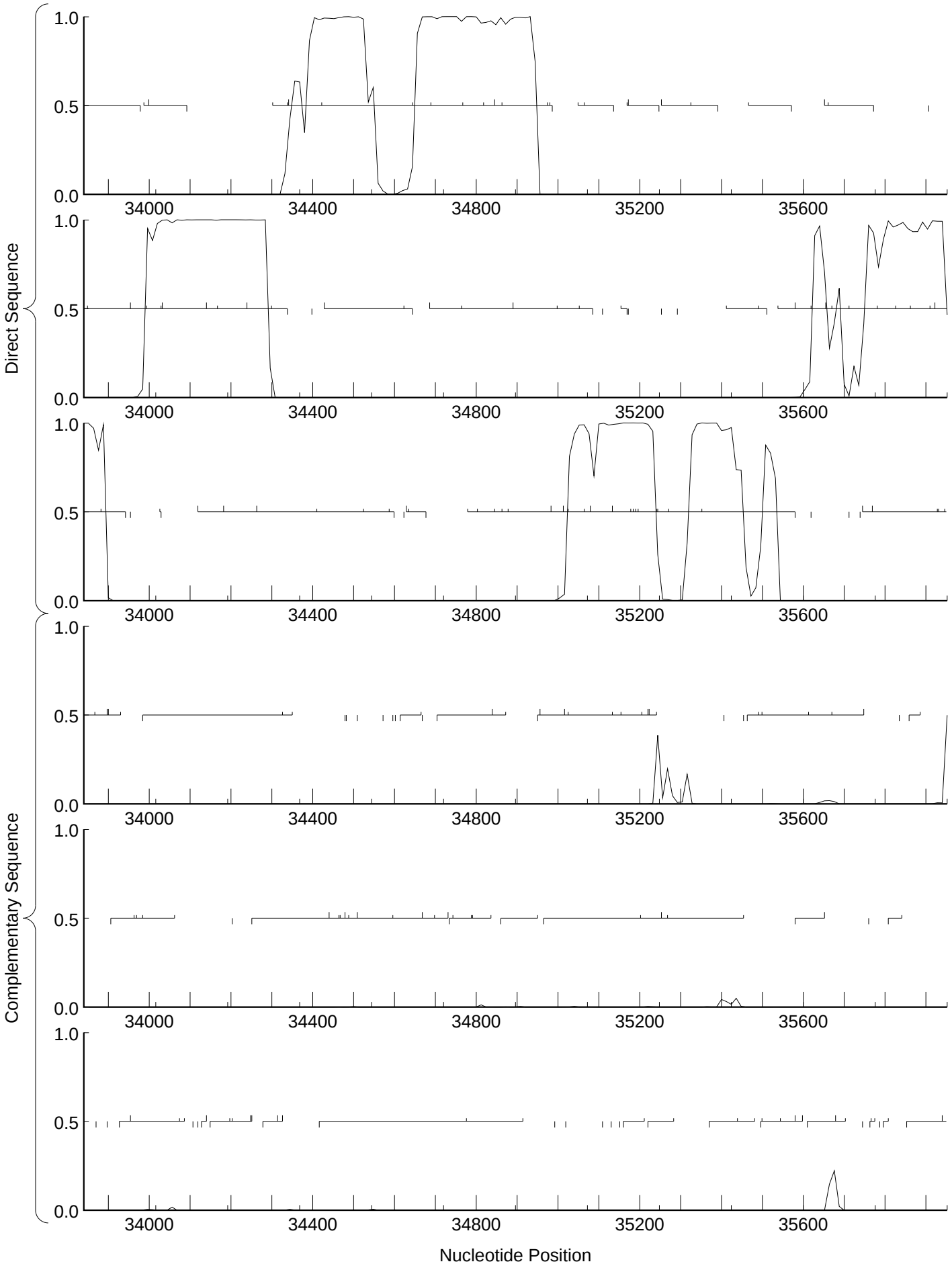
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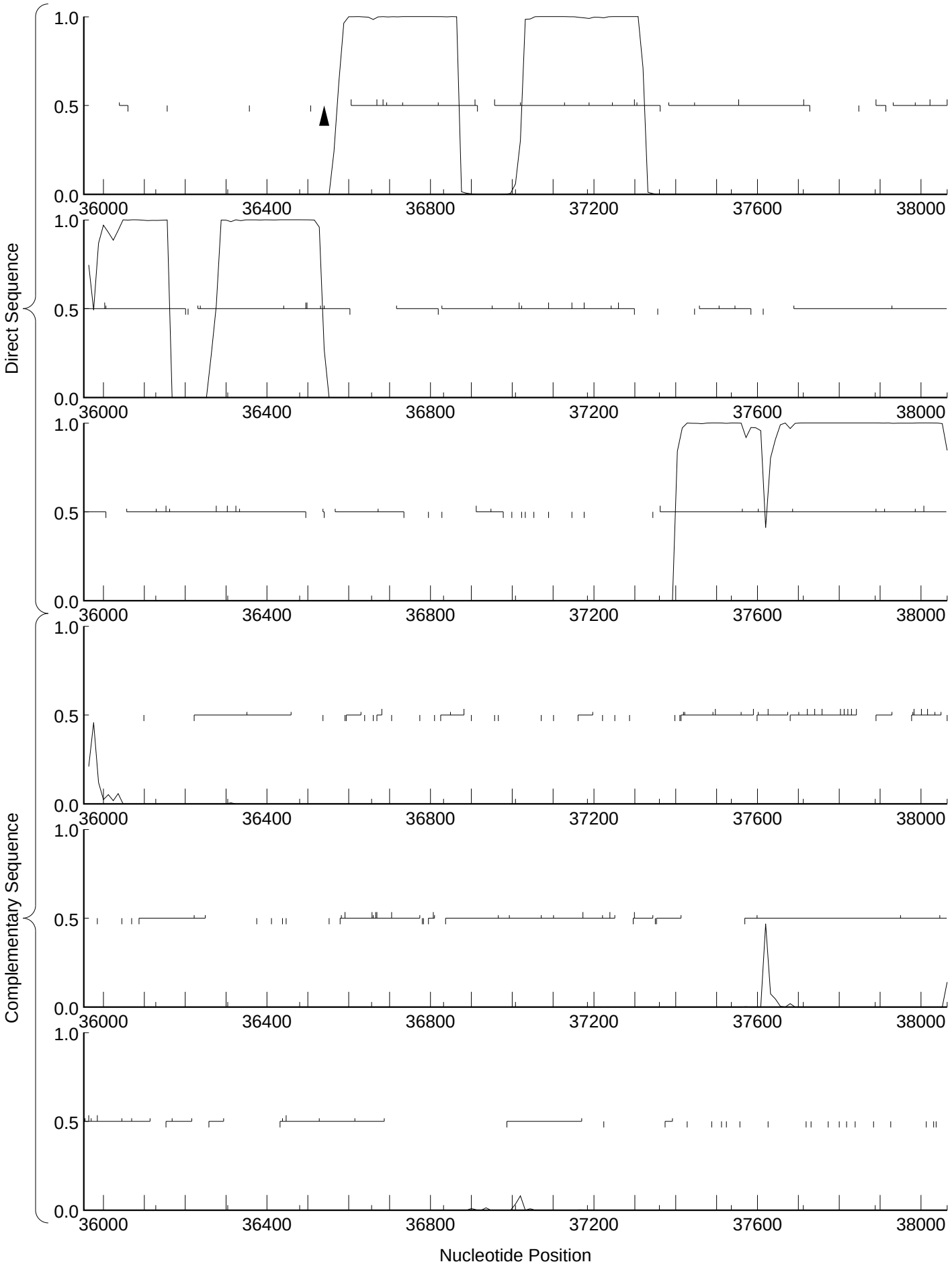
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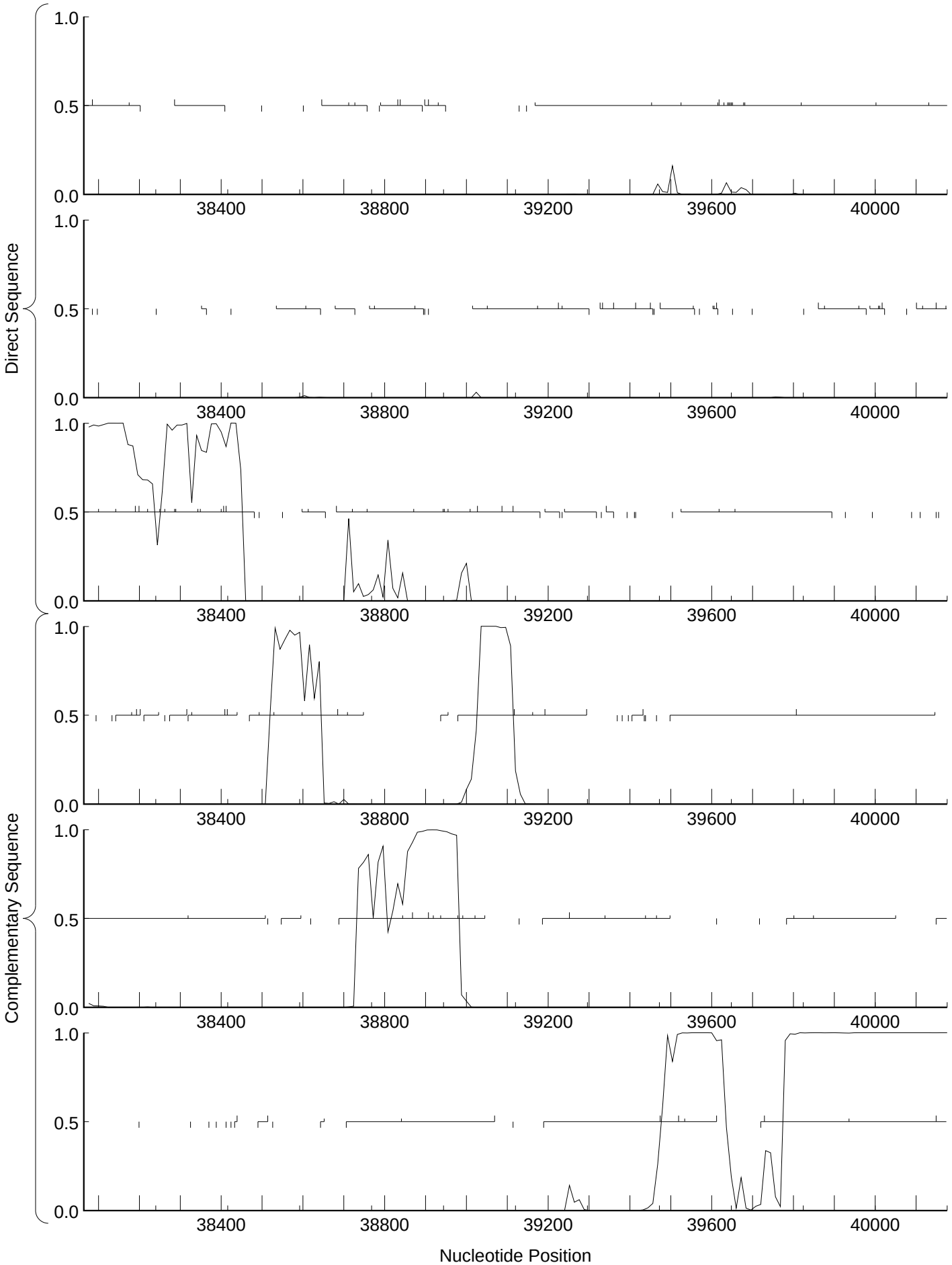
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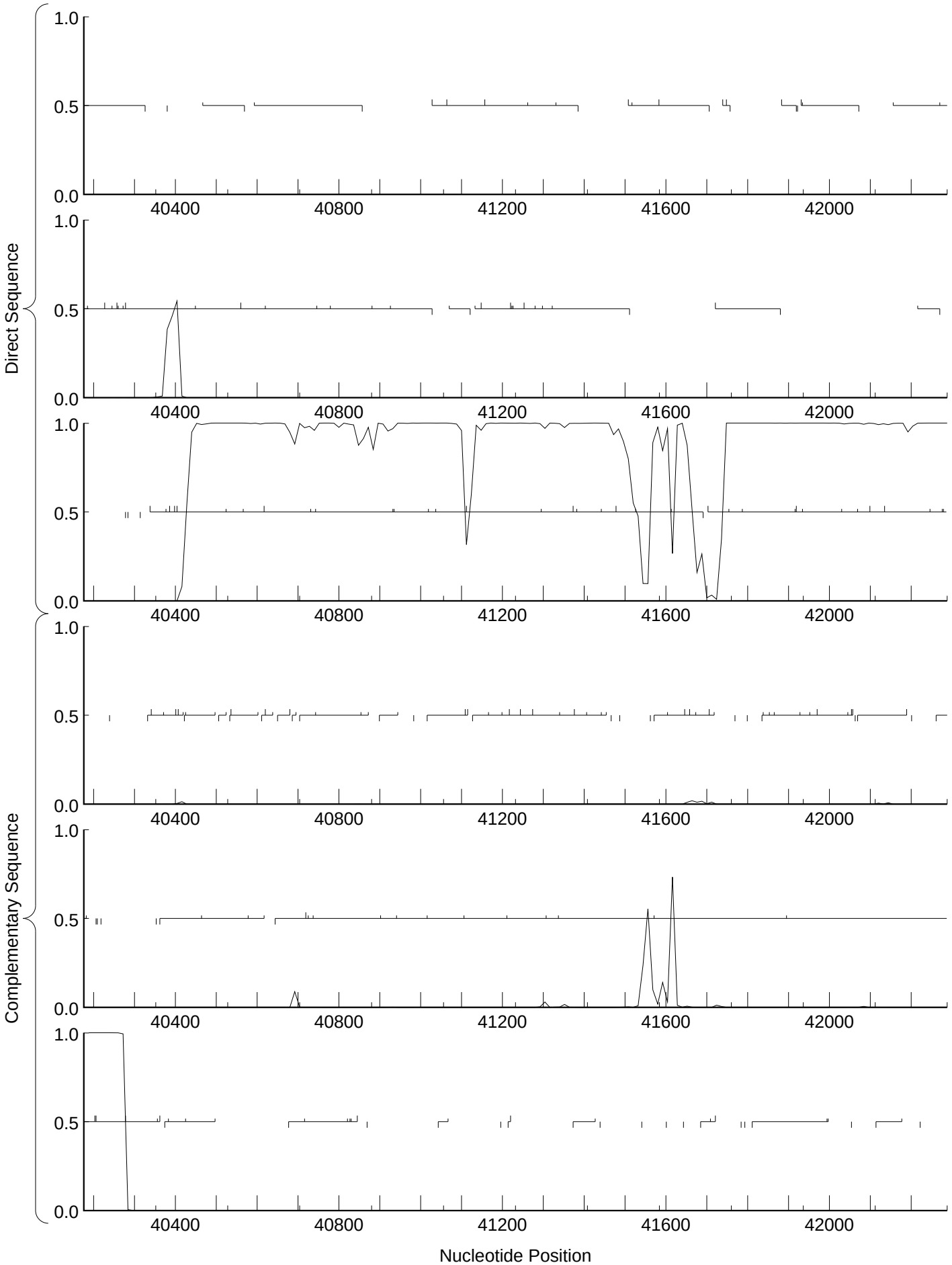
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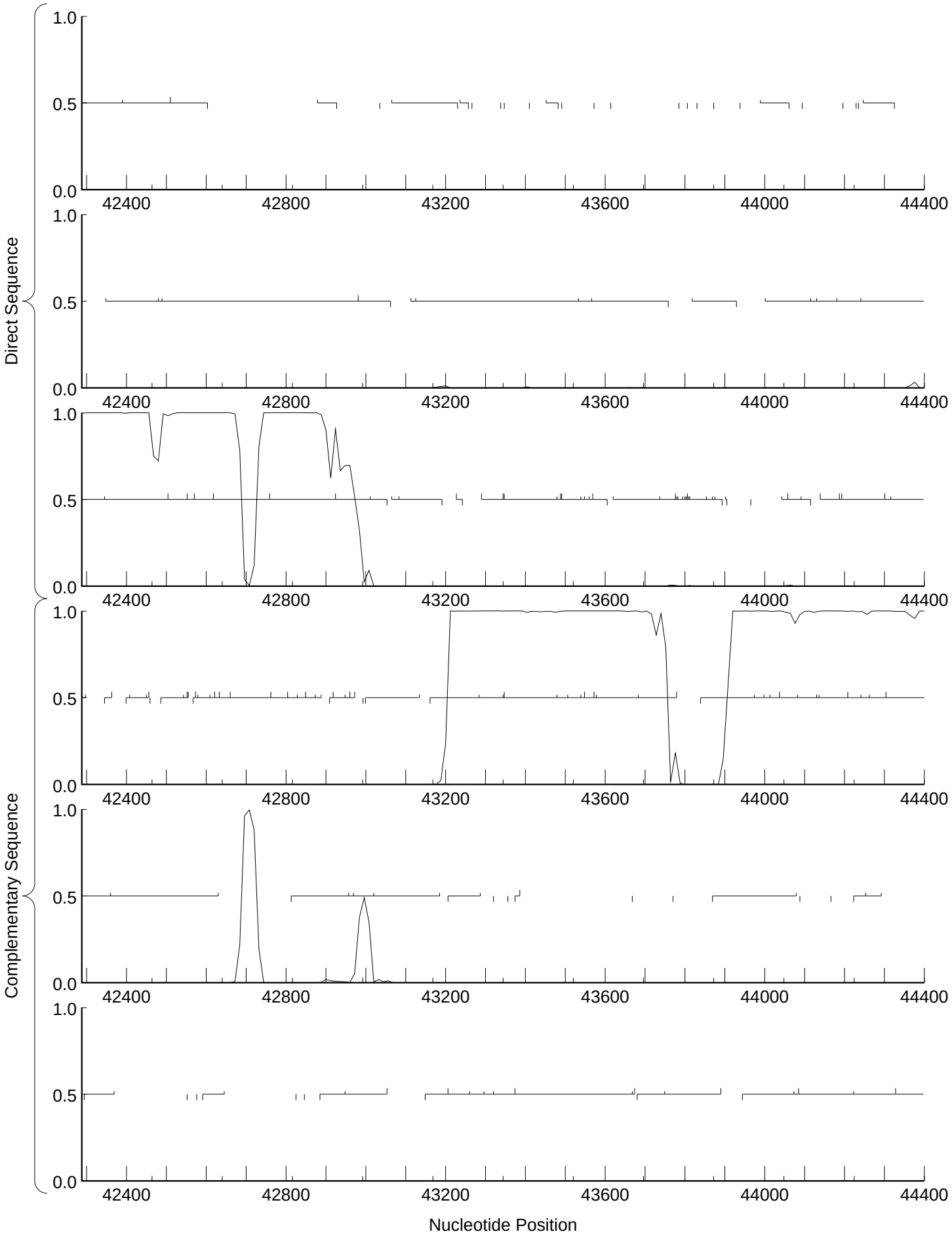
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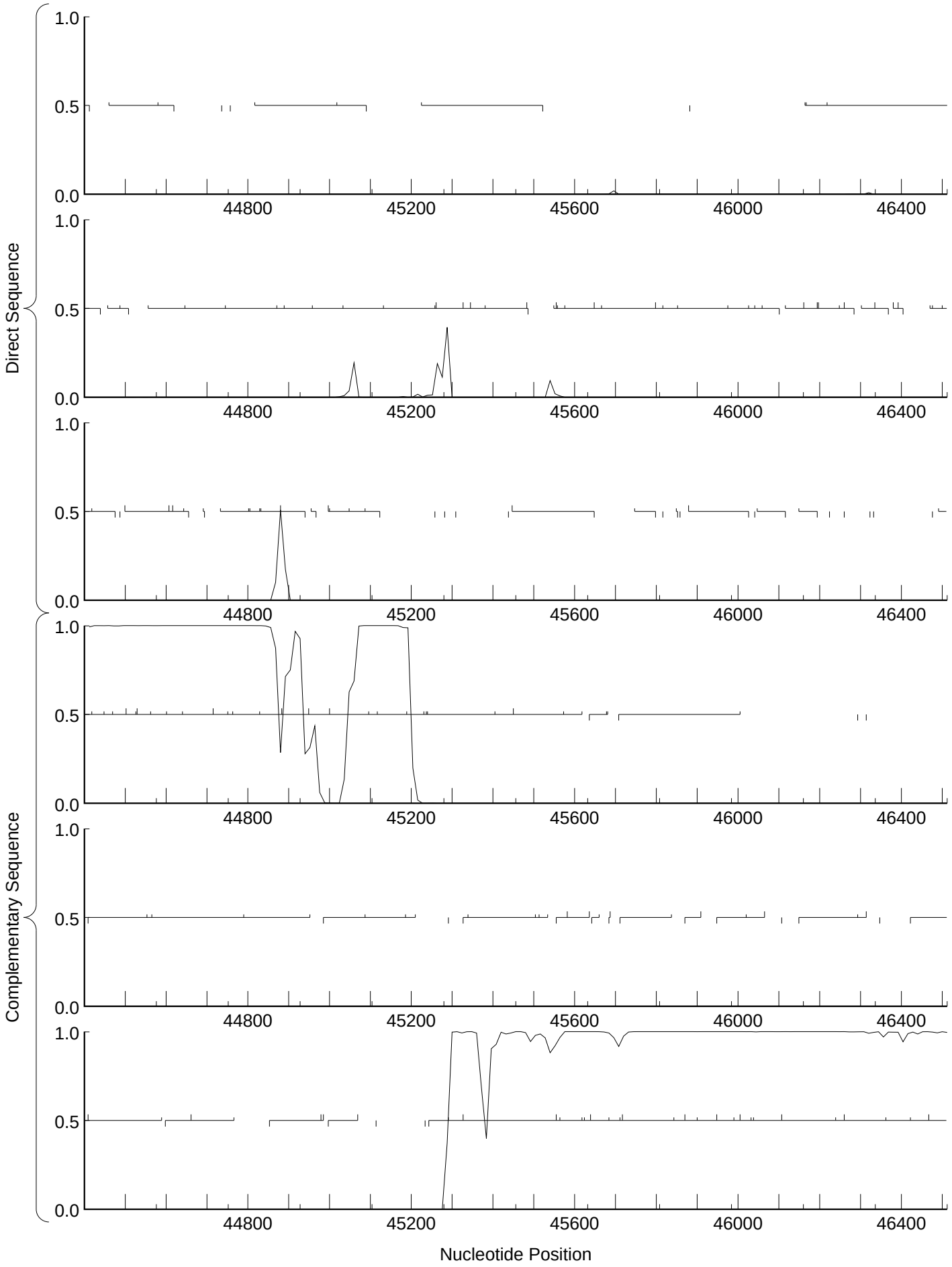
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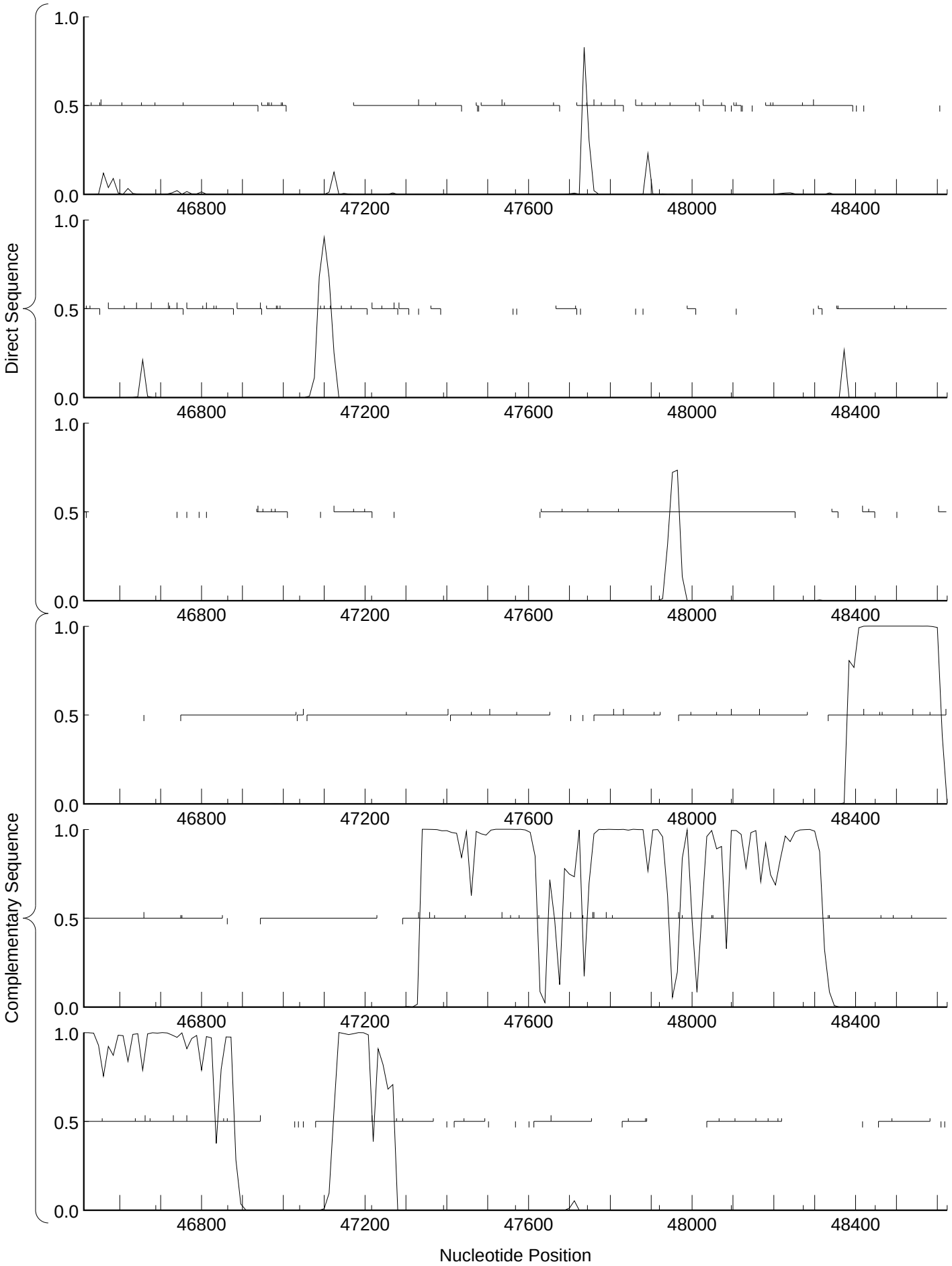
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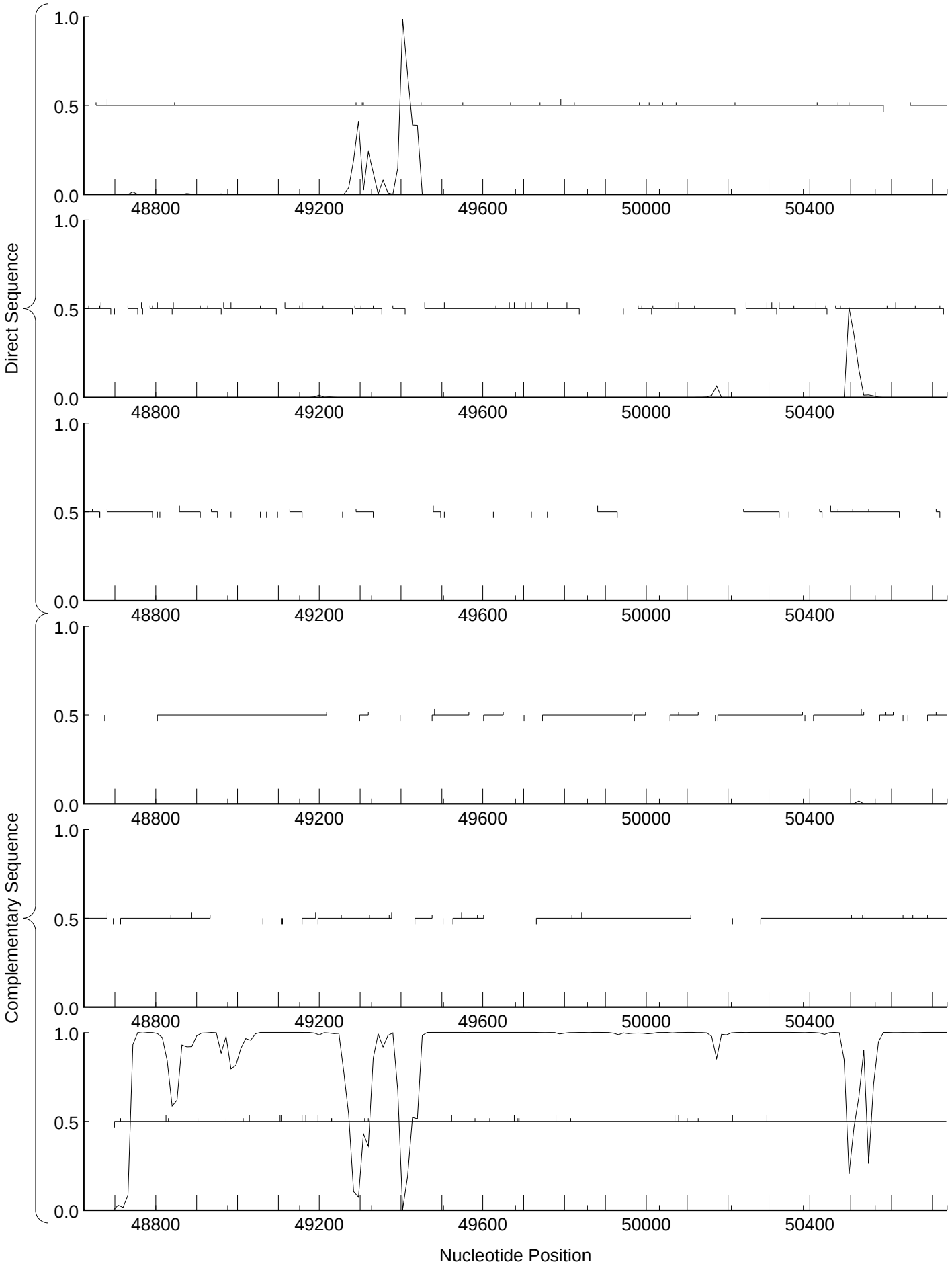
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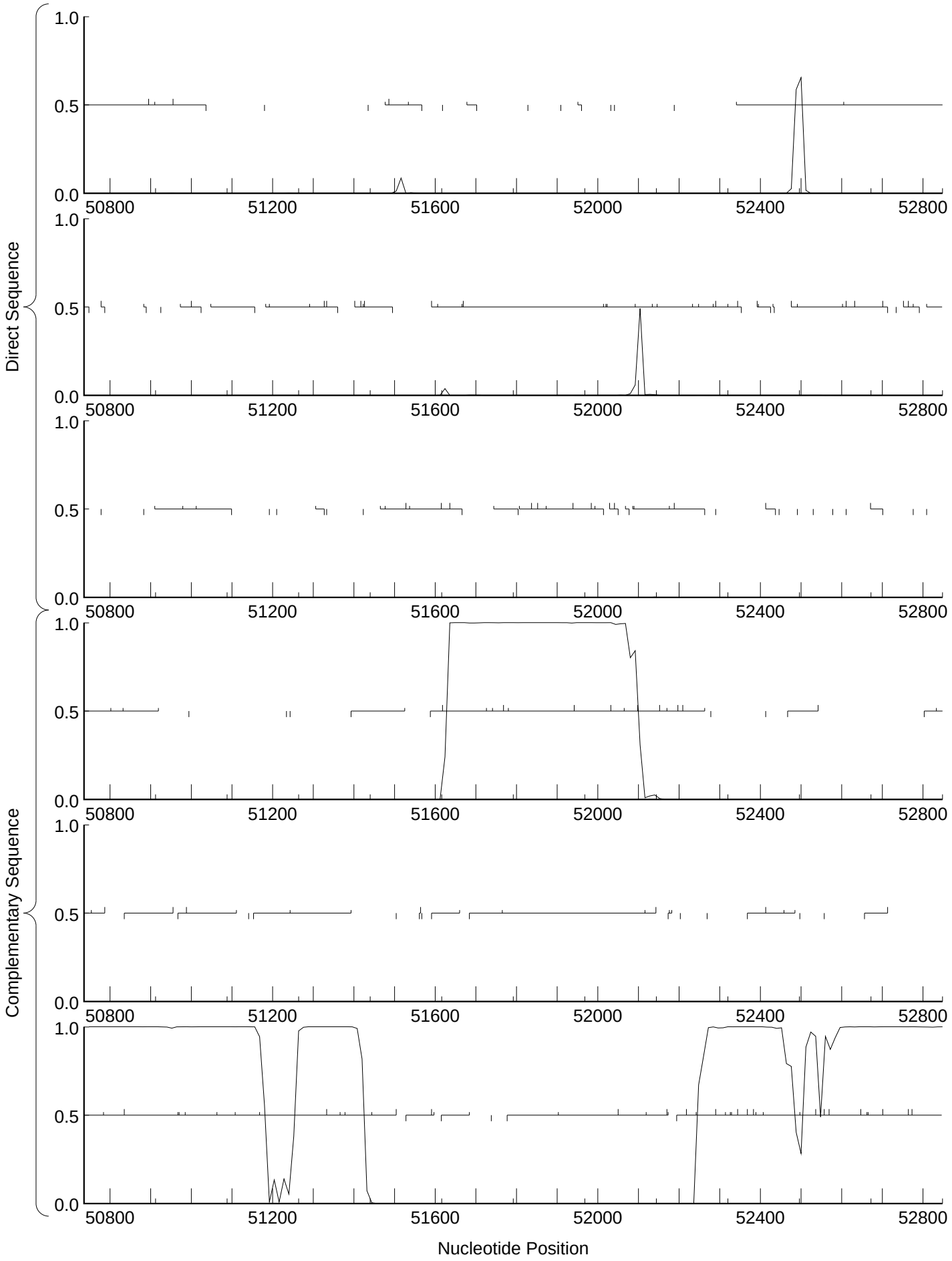
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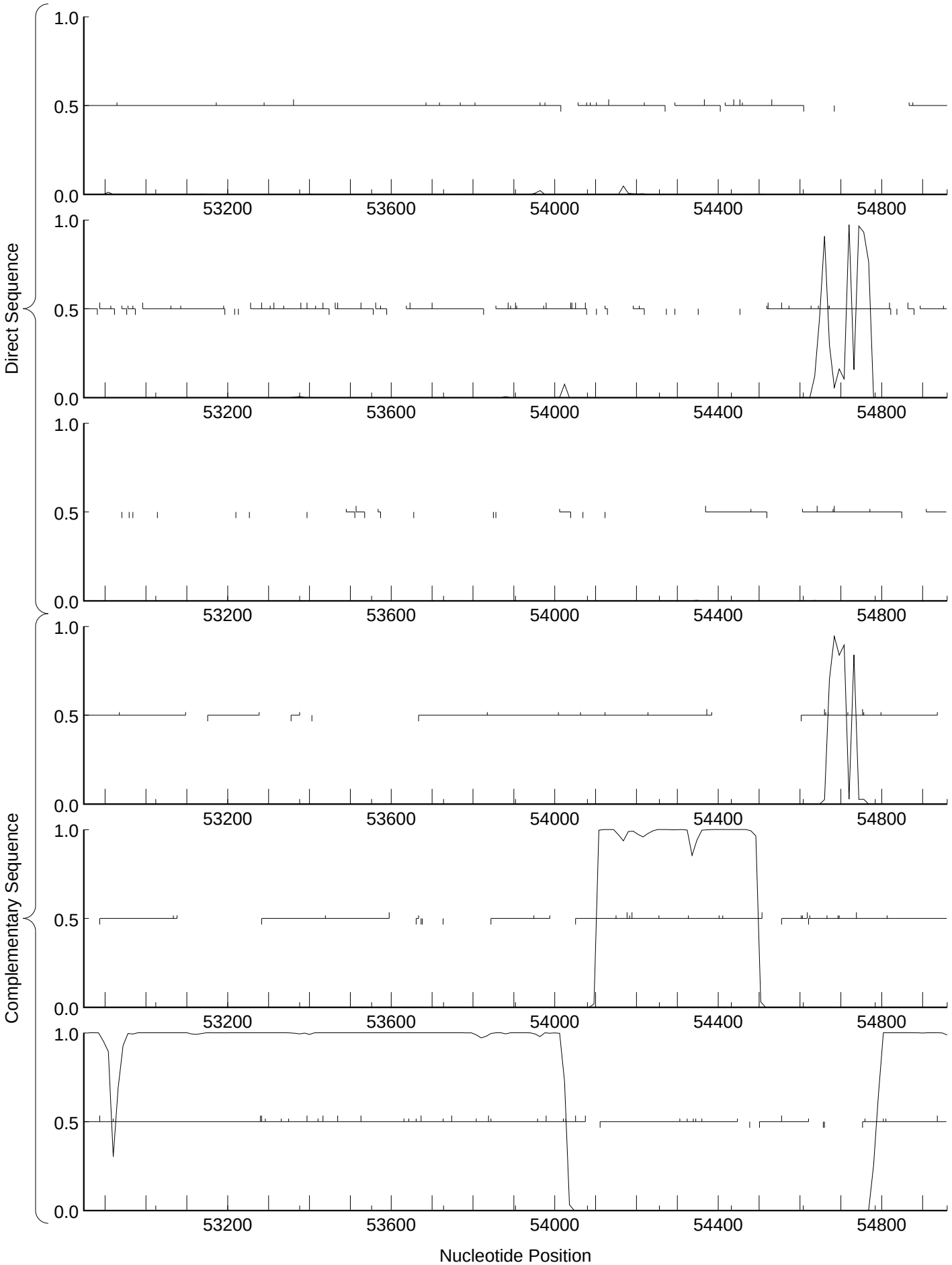
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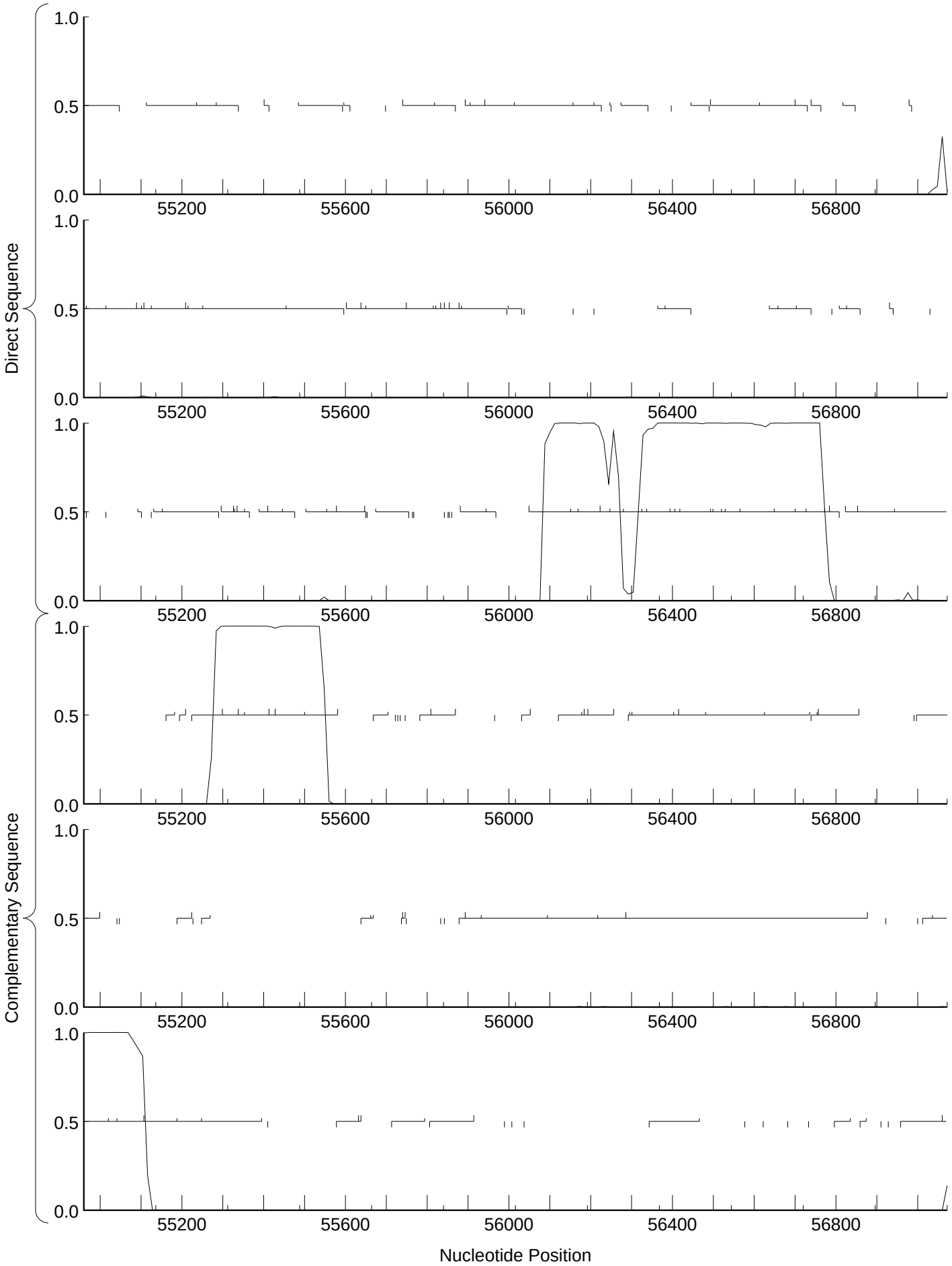
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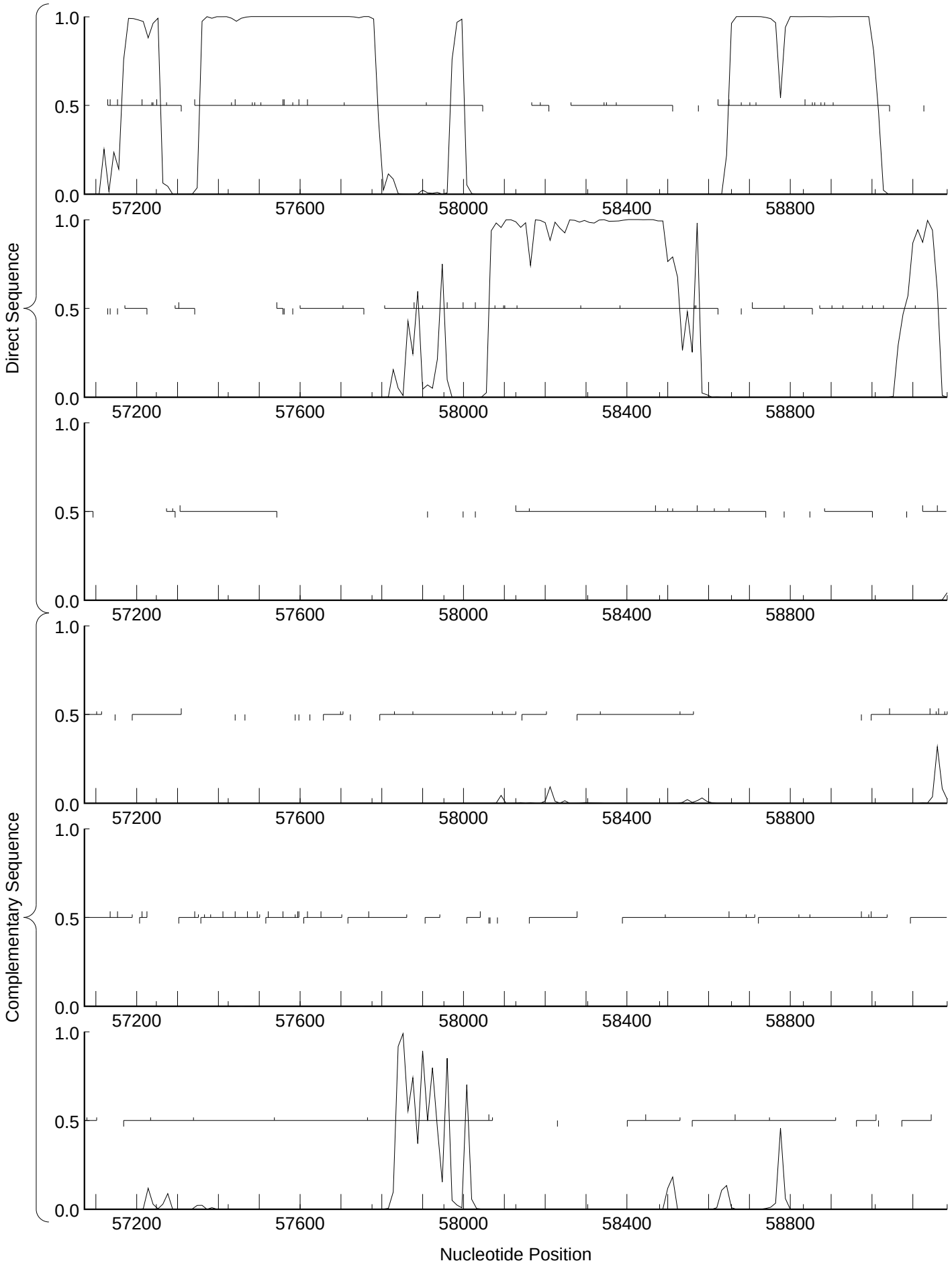
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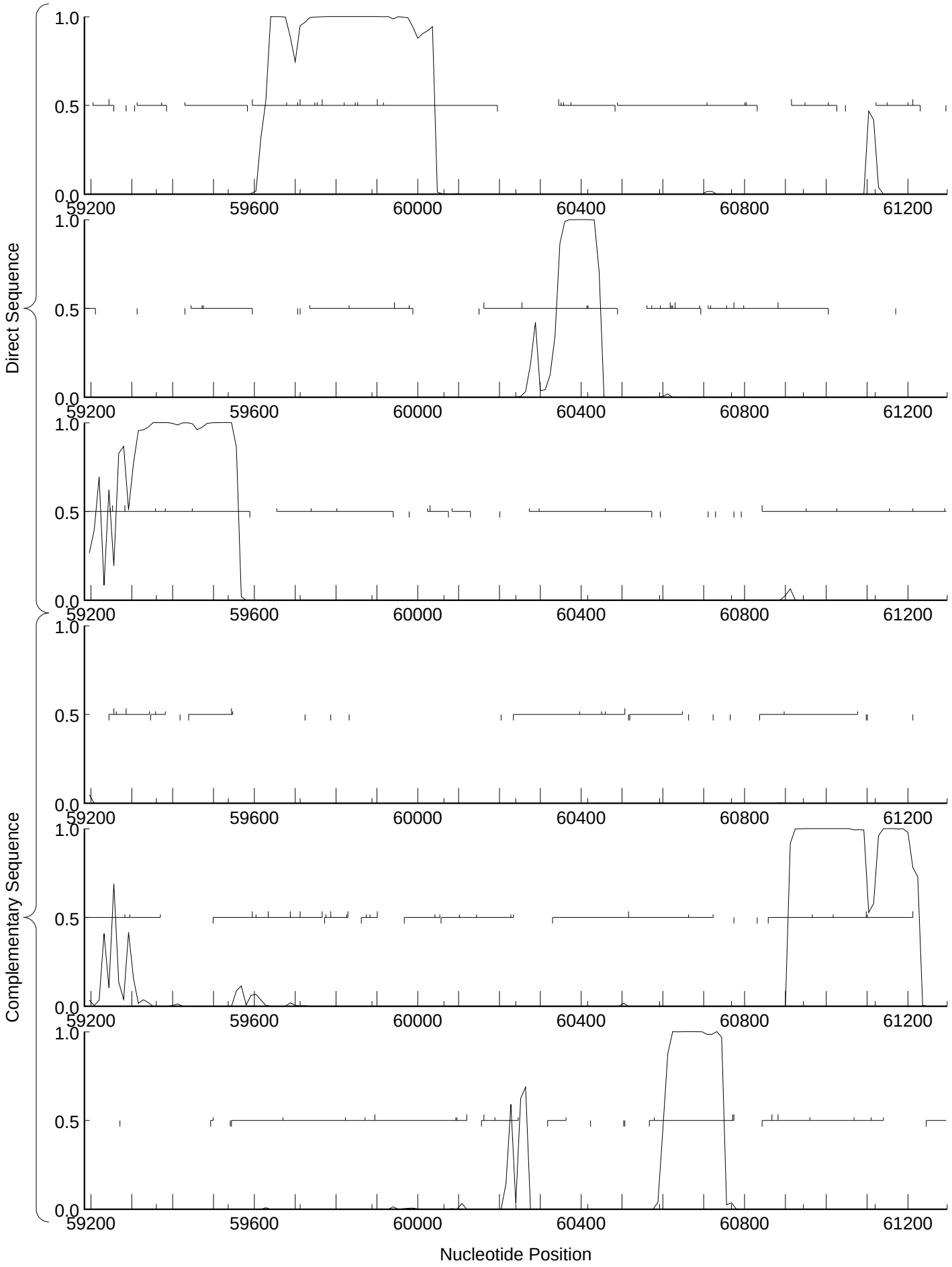
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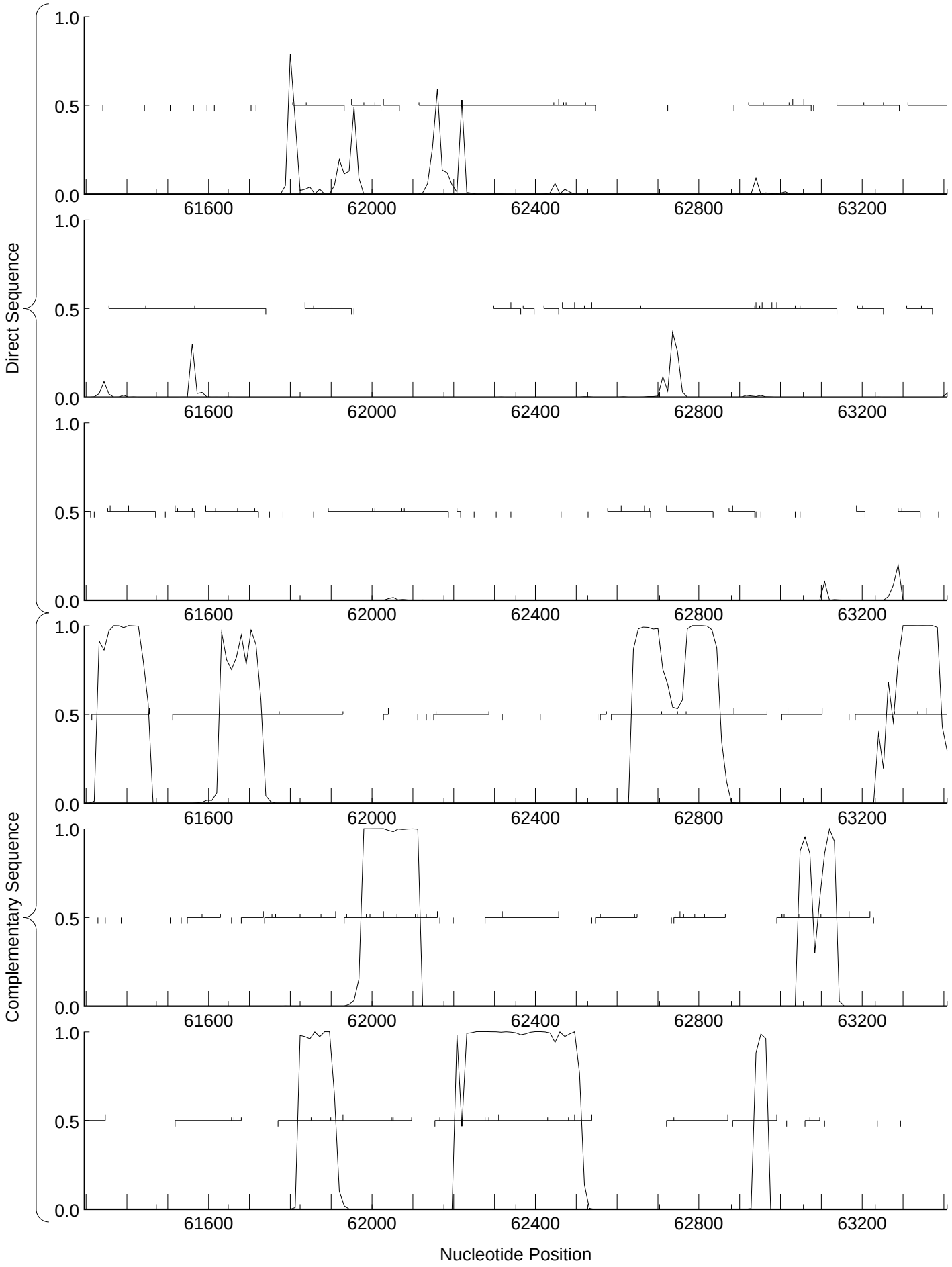


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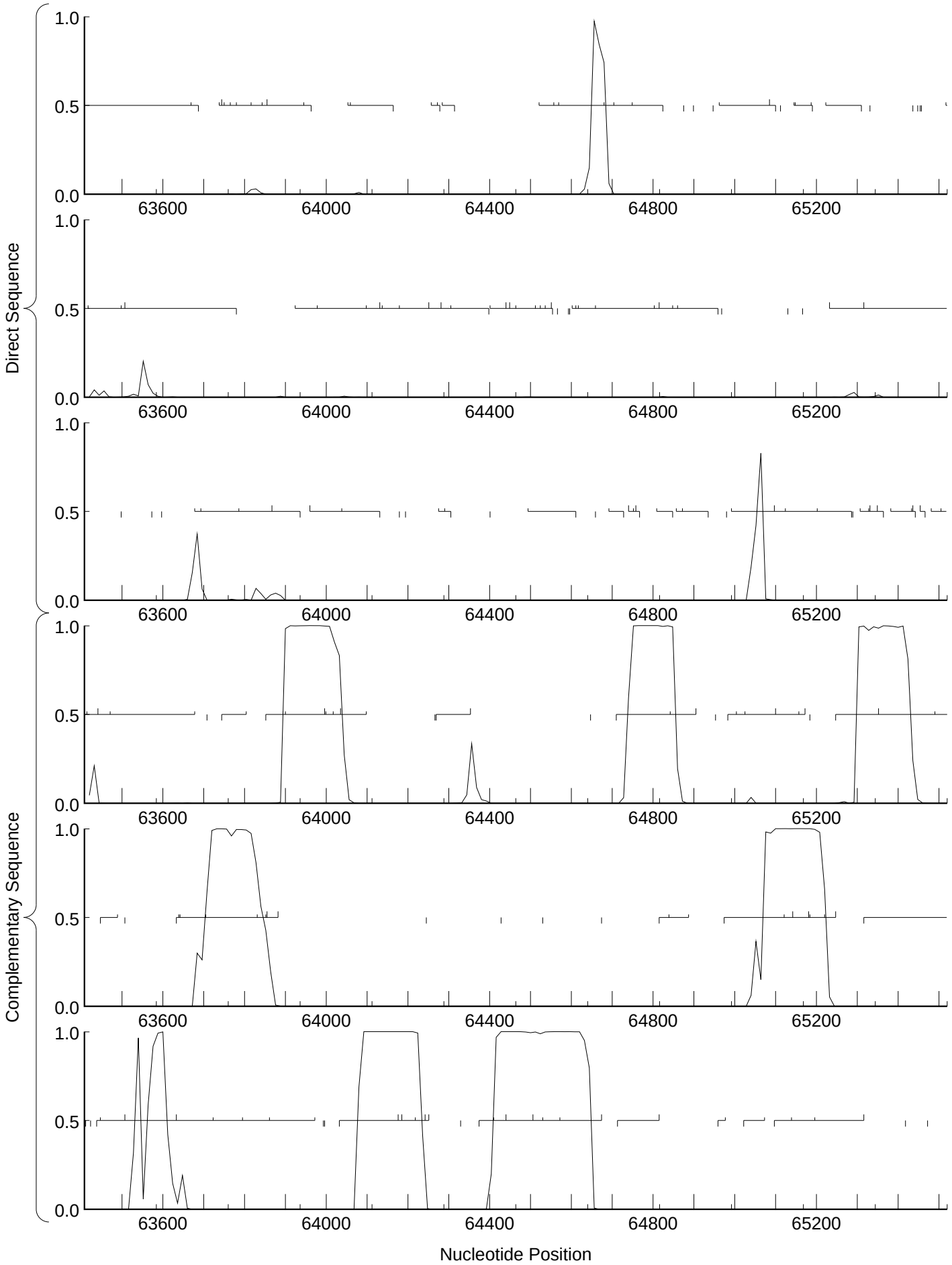


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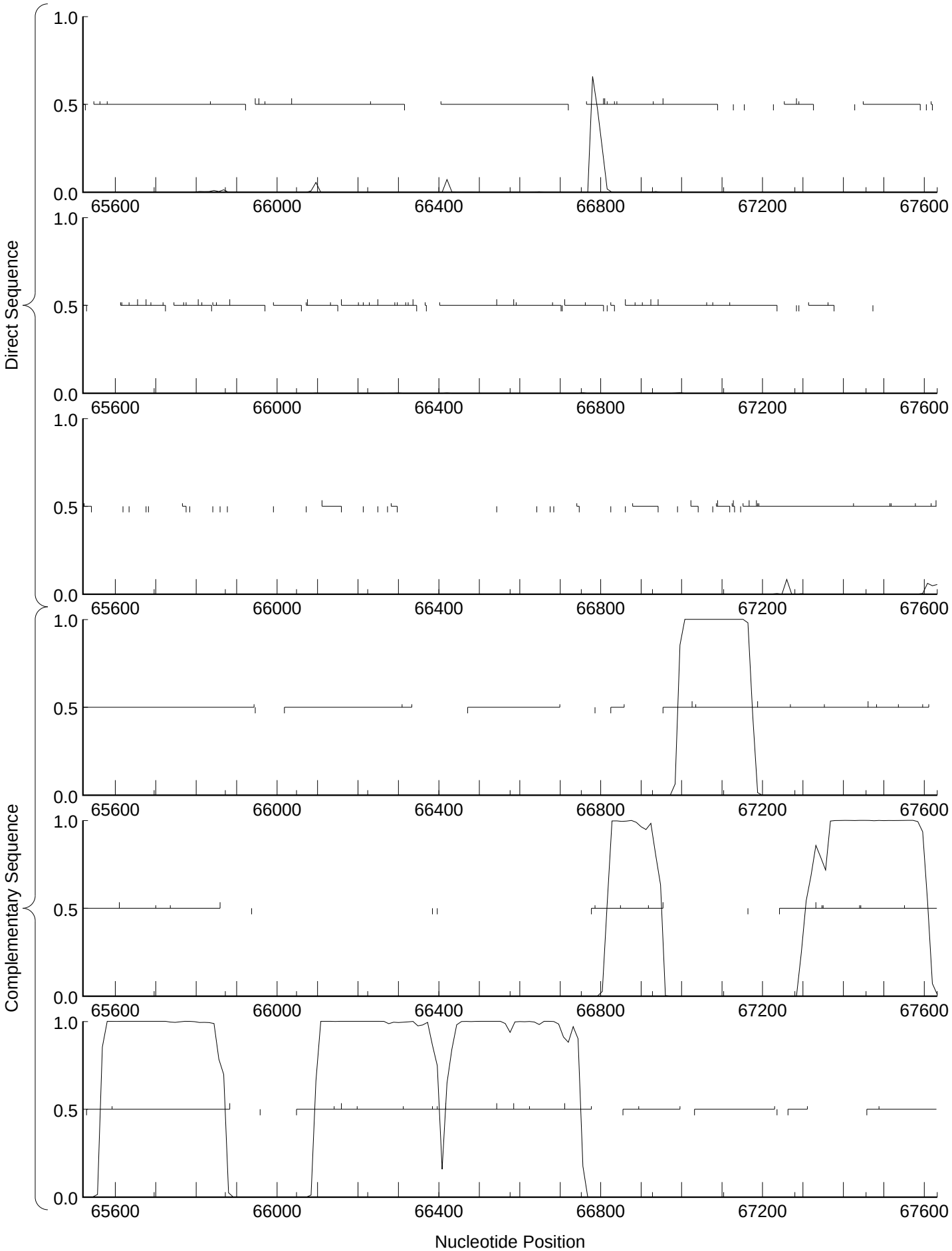



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
61600
62000
62400
62800
63200
Nucleotide Position

ycobacterium phage ShiVal Complete Sequence, 68355 bp, Circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 32/34



ycobacterium phage ShiVal Complete Sequence, 68355 bp, Circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 33/34



ycobacterium phage ShiVal Complete Sequence, 68355 bp, Circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 34/34

