GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage BlackStallion complete sequence, 68714 bp, circularly permuted, Cluster
Analysis Date    :  3/28/24 at 15:38:13
Pages            :  34
Sequence Length  :  68714 bp
GC Content       :   66.50%

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

PS-Version       :  1.2

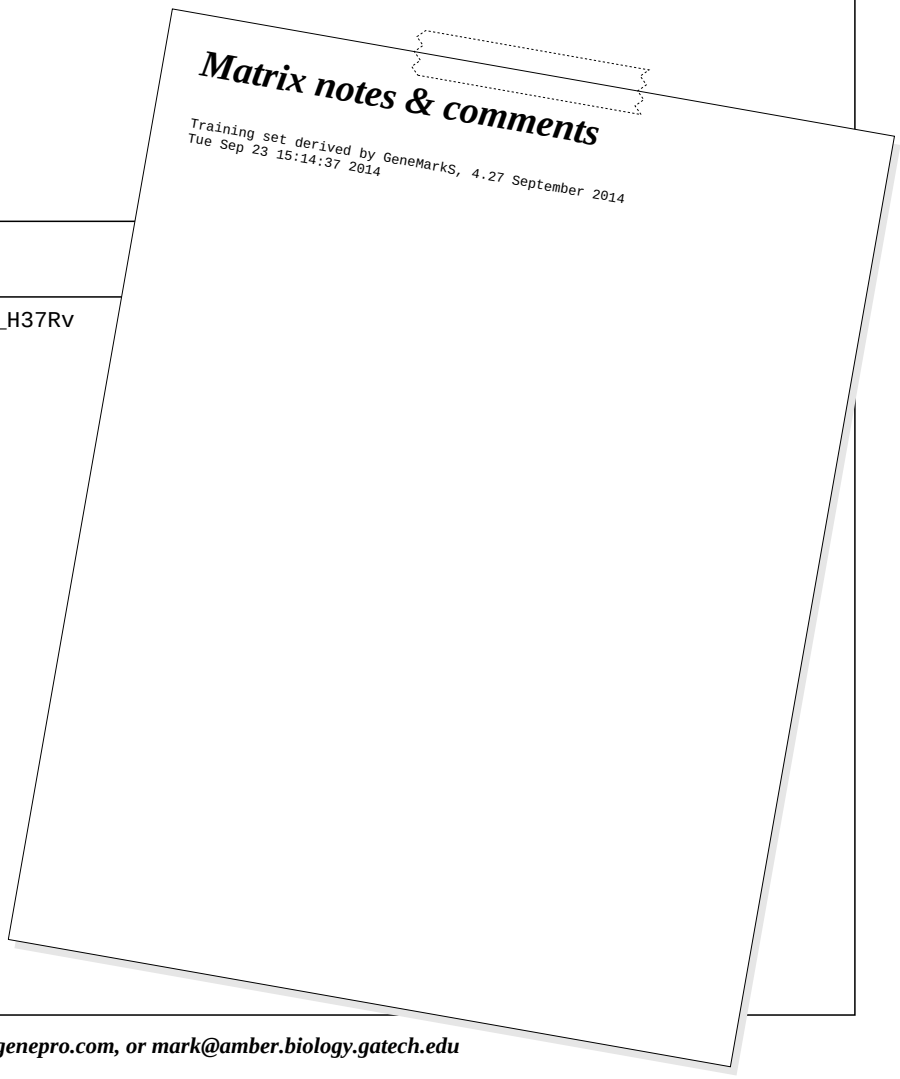
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                    Mark ORFs / splice sites,
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MATRIX INFORMATION

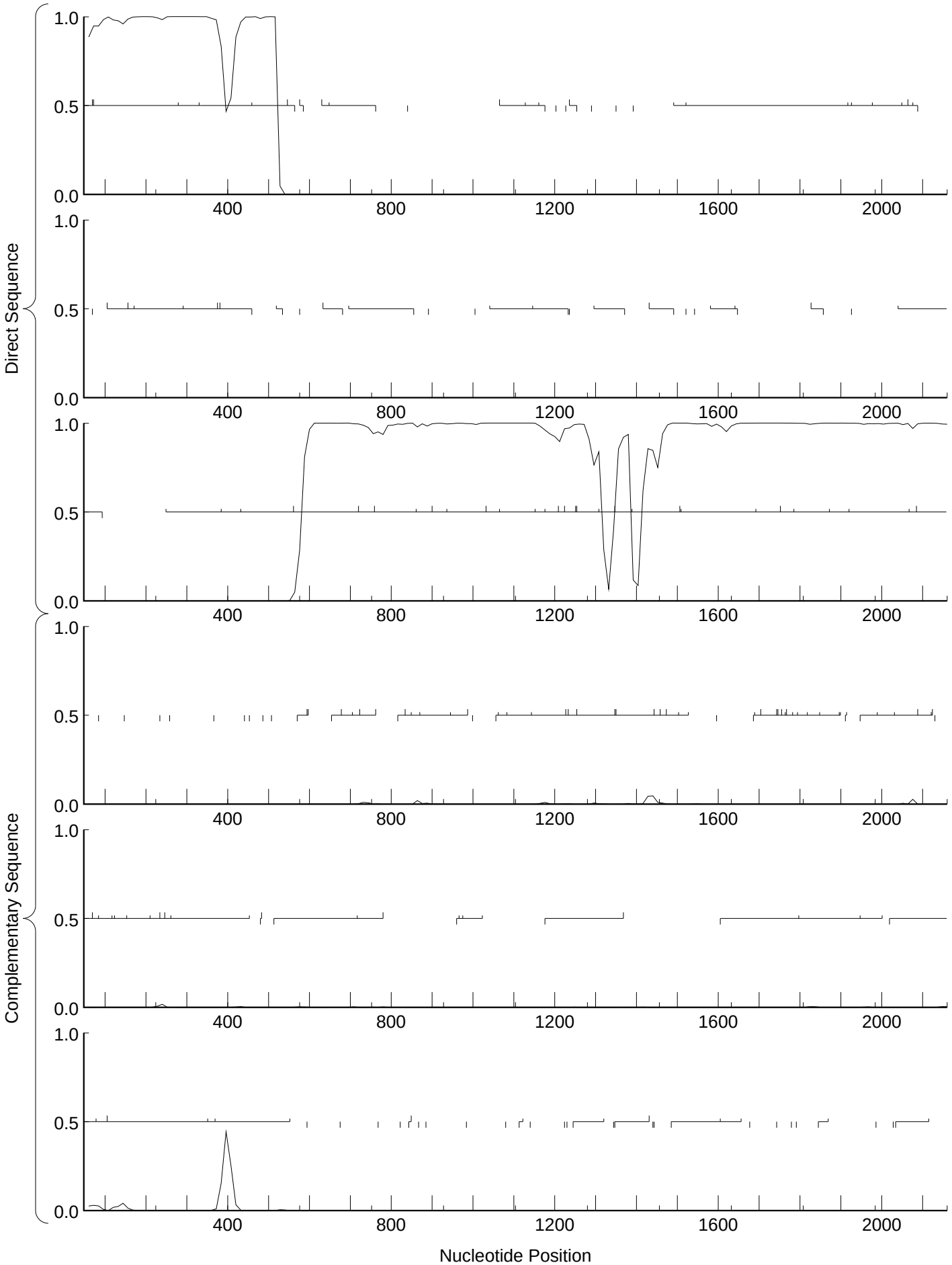
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Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

Order      :  4
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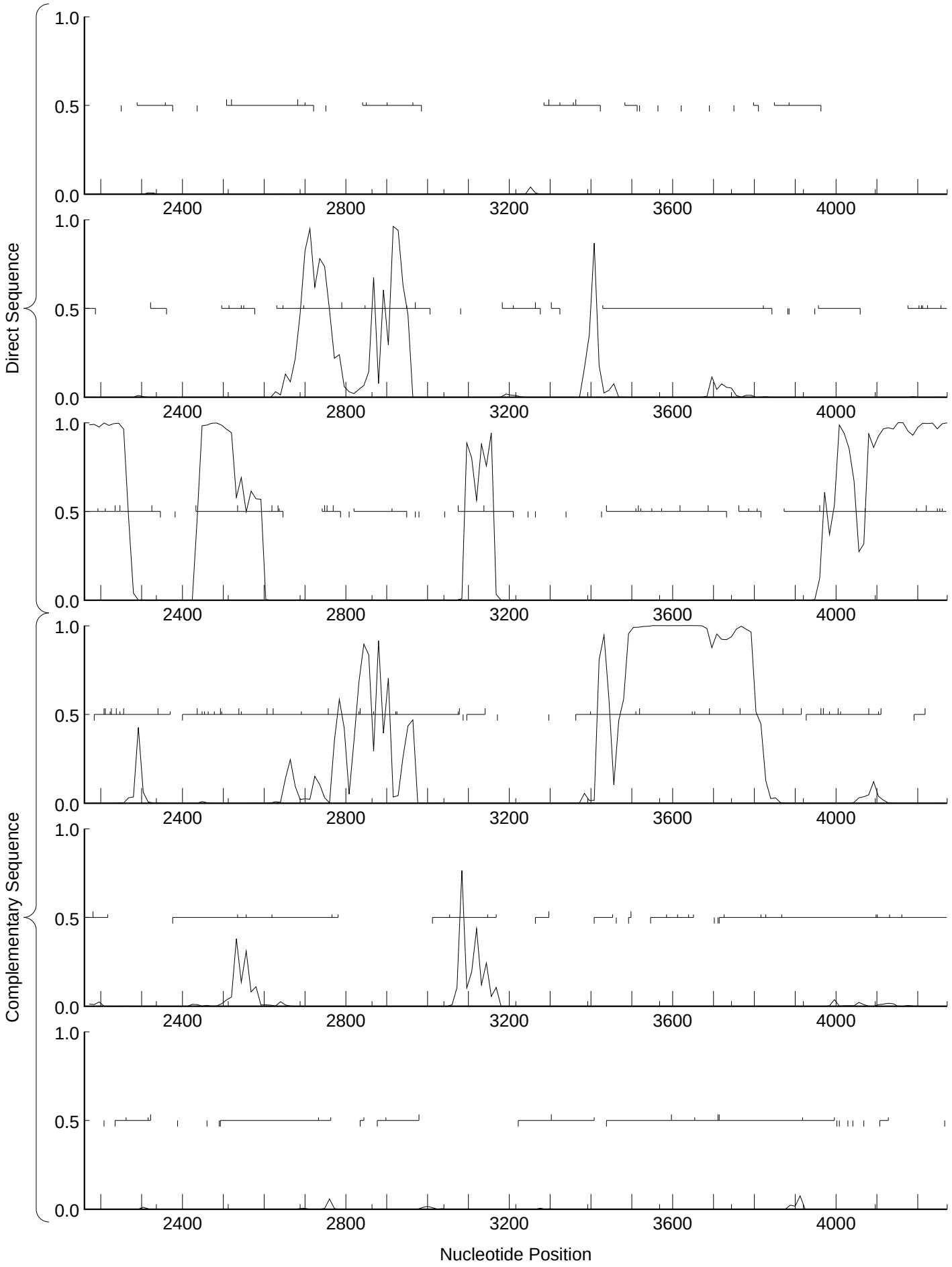

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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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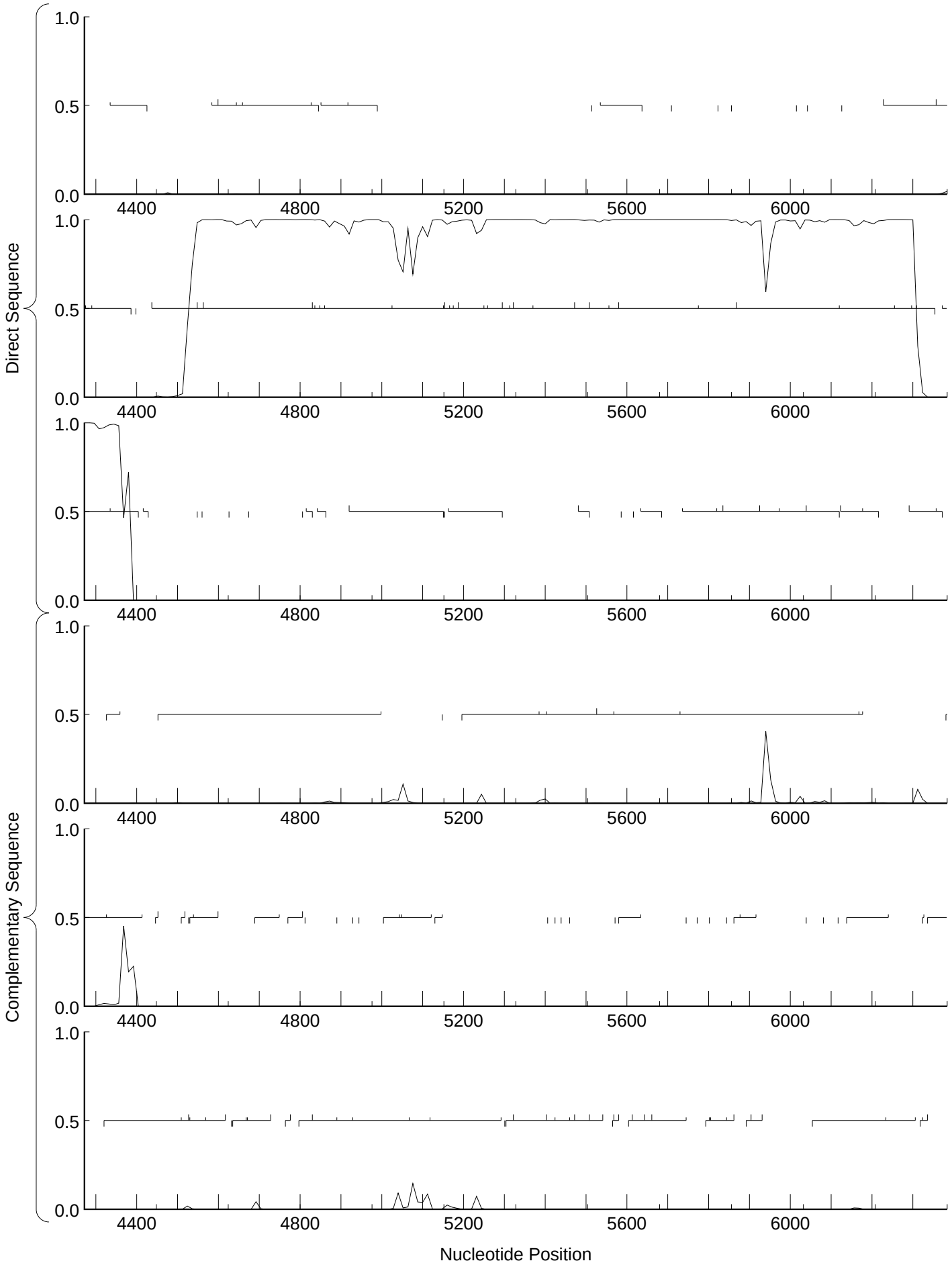



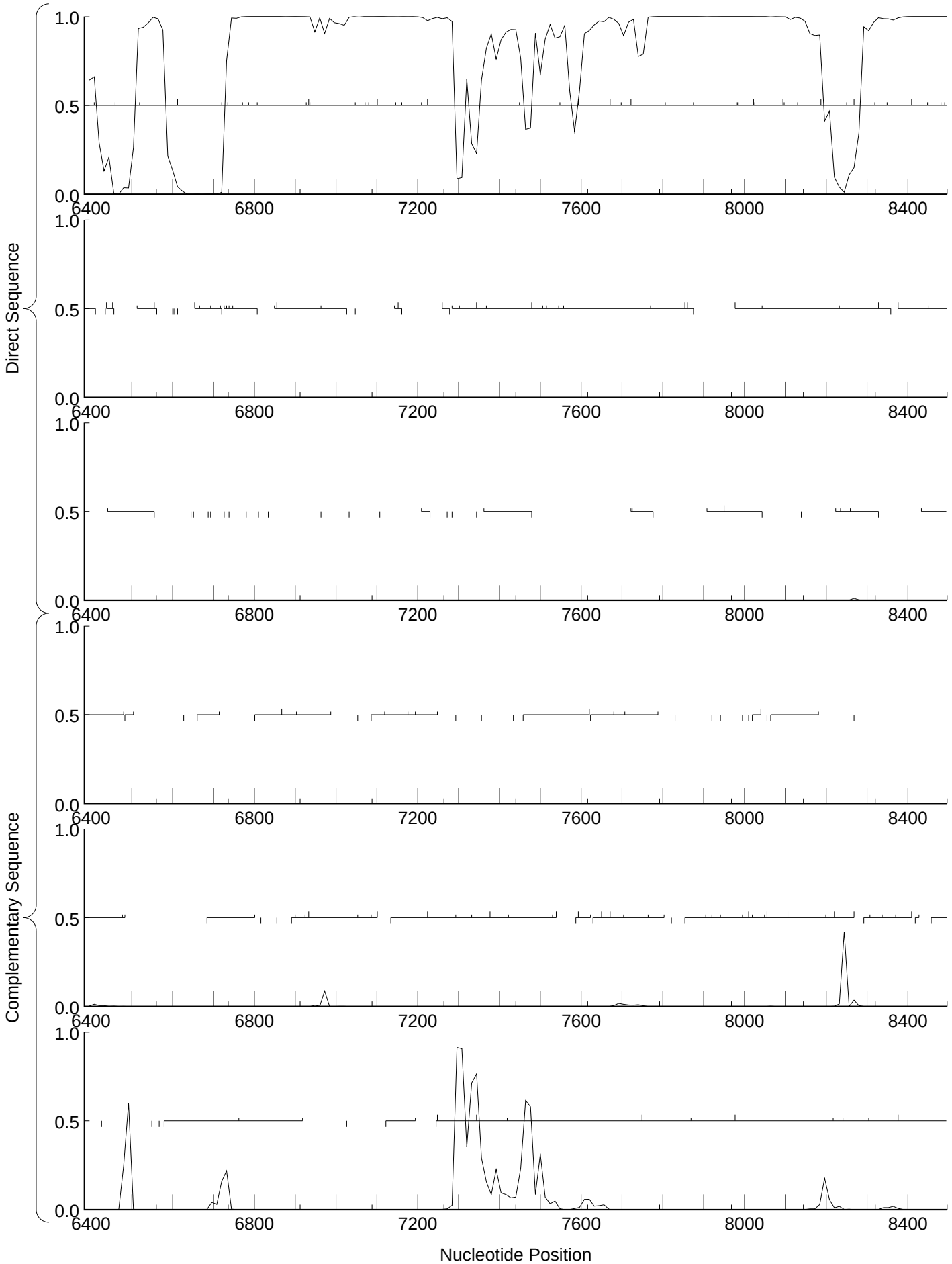
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu


Direct Sequence
Complementary Sequence
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0.0
400
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1200
1600
2000
Nucleotide Position

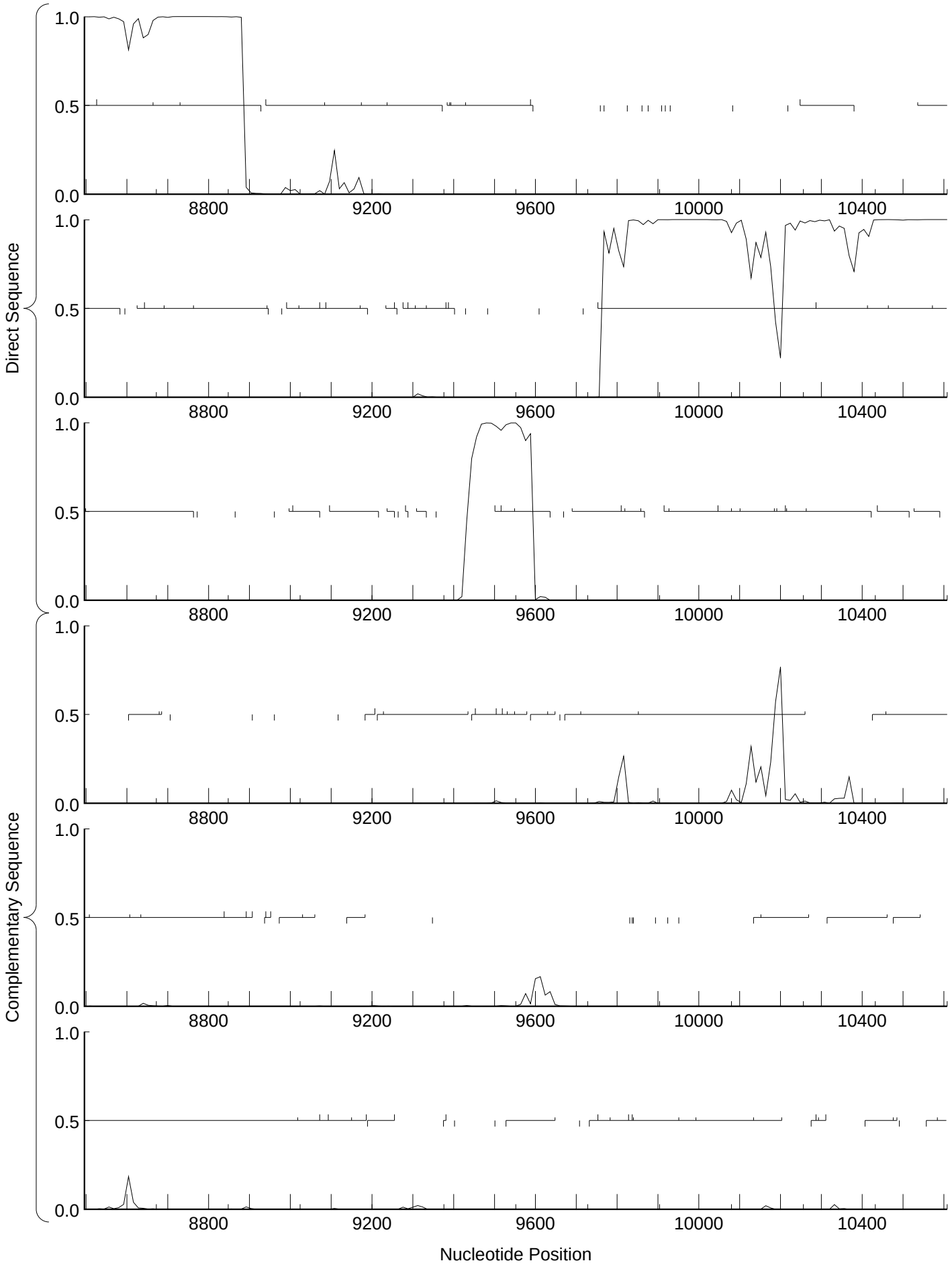
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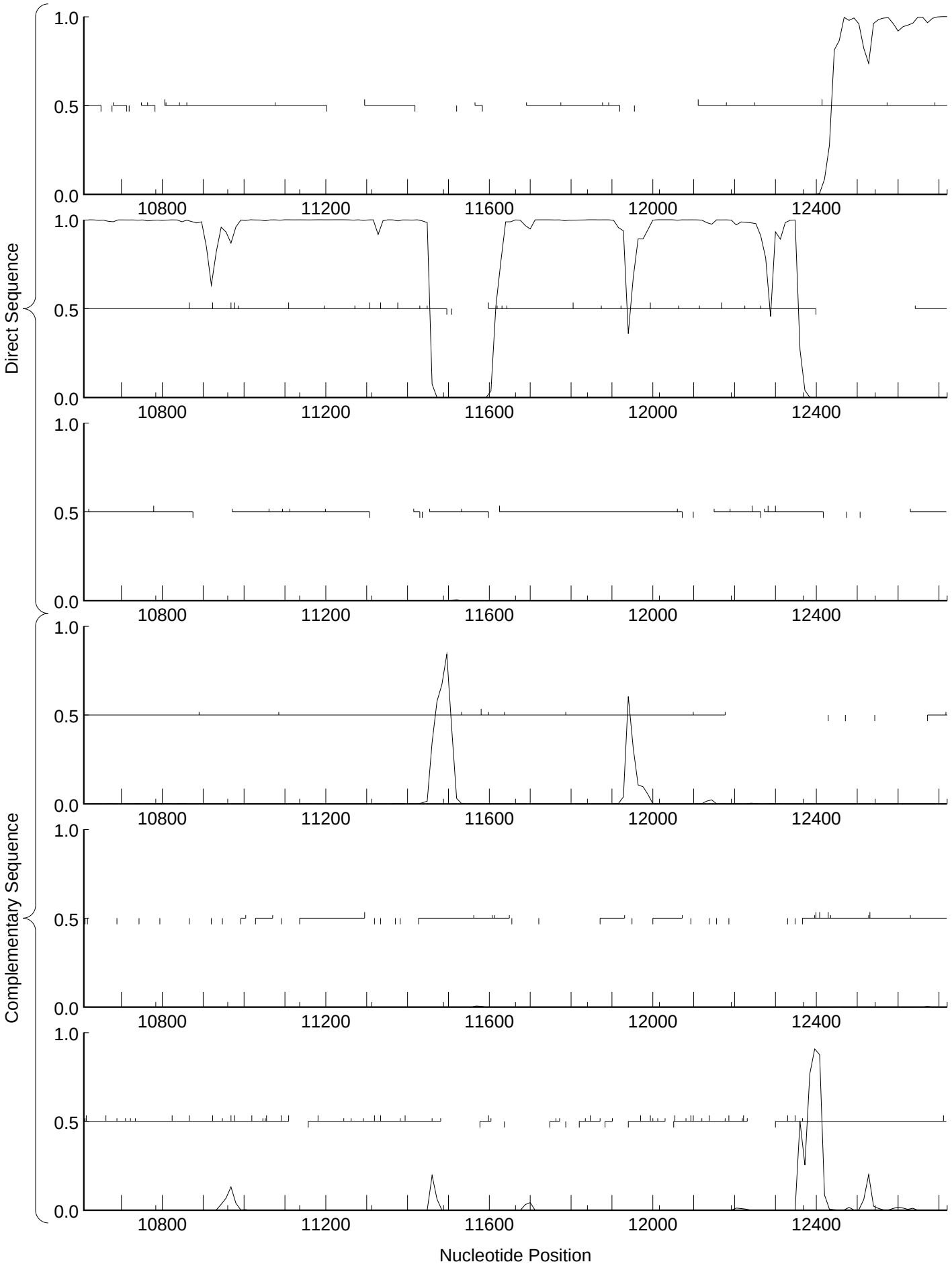

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Nucleotide Position

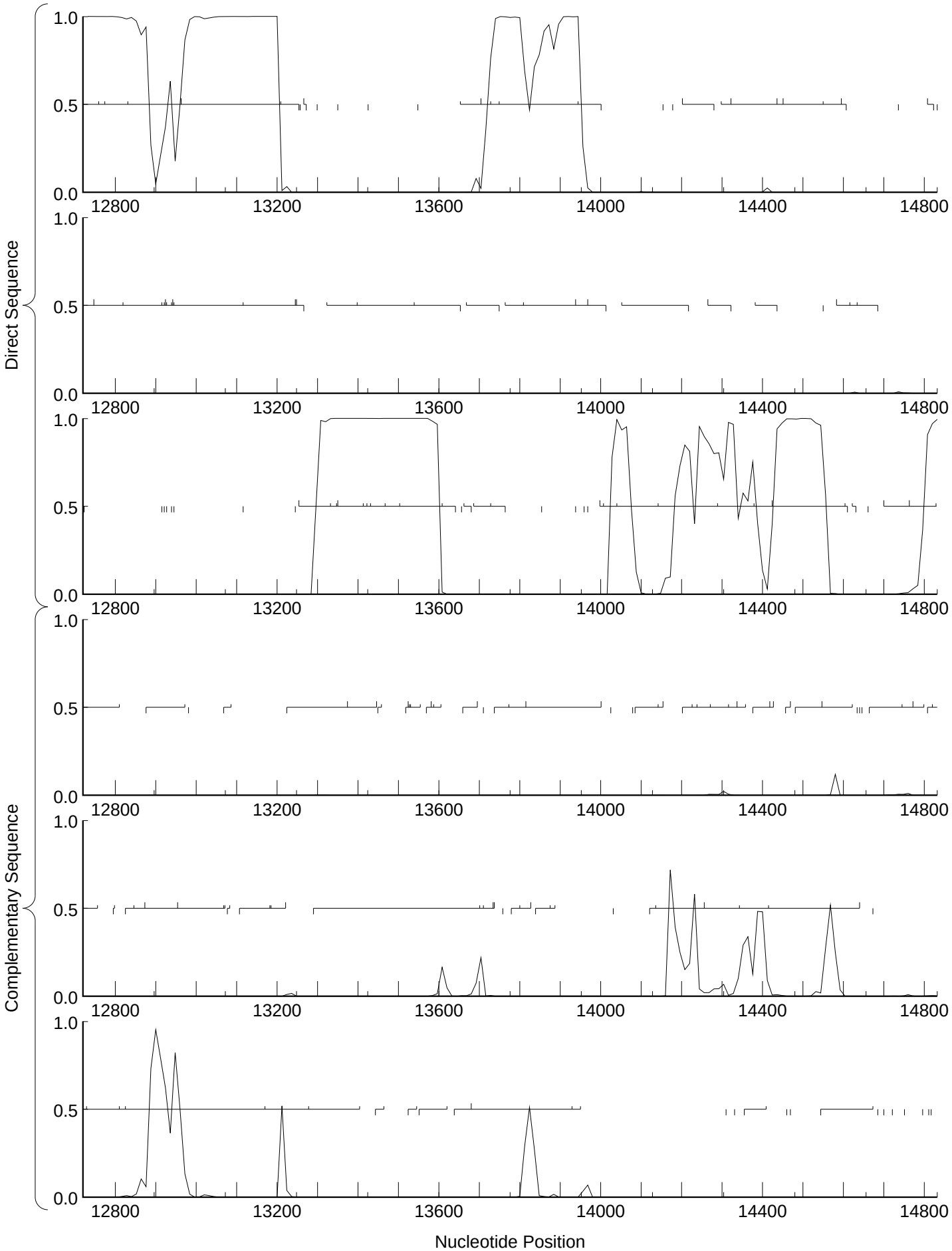

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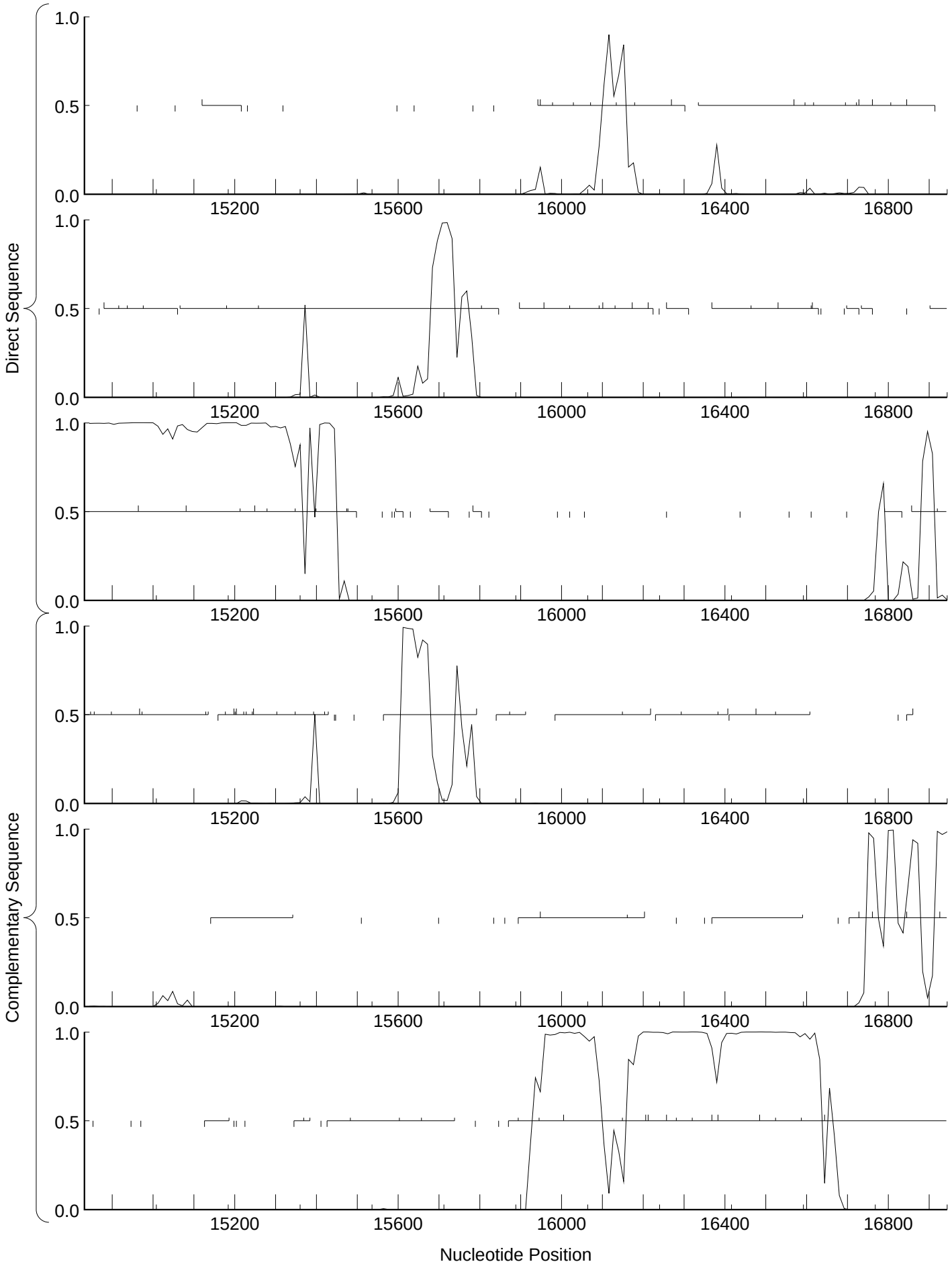
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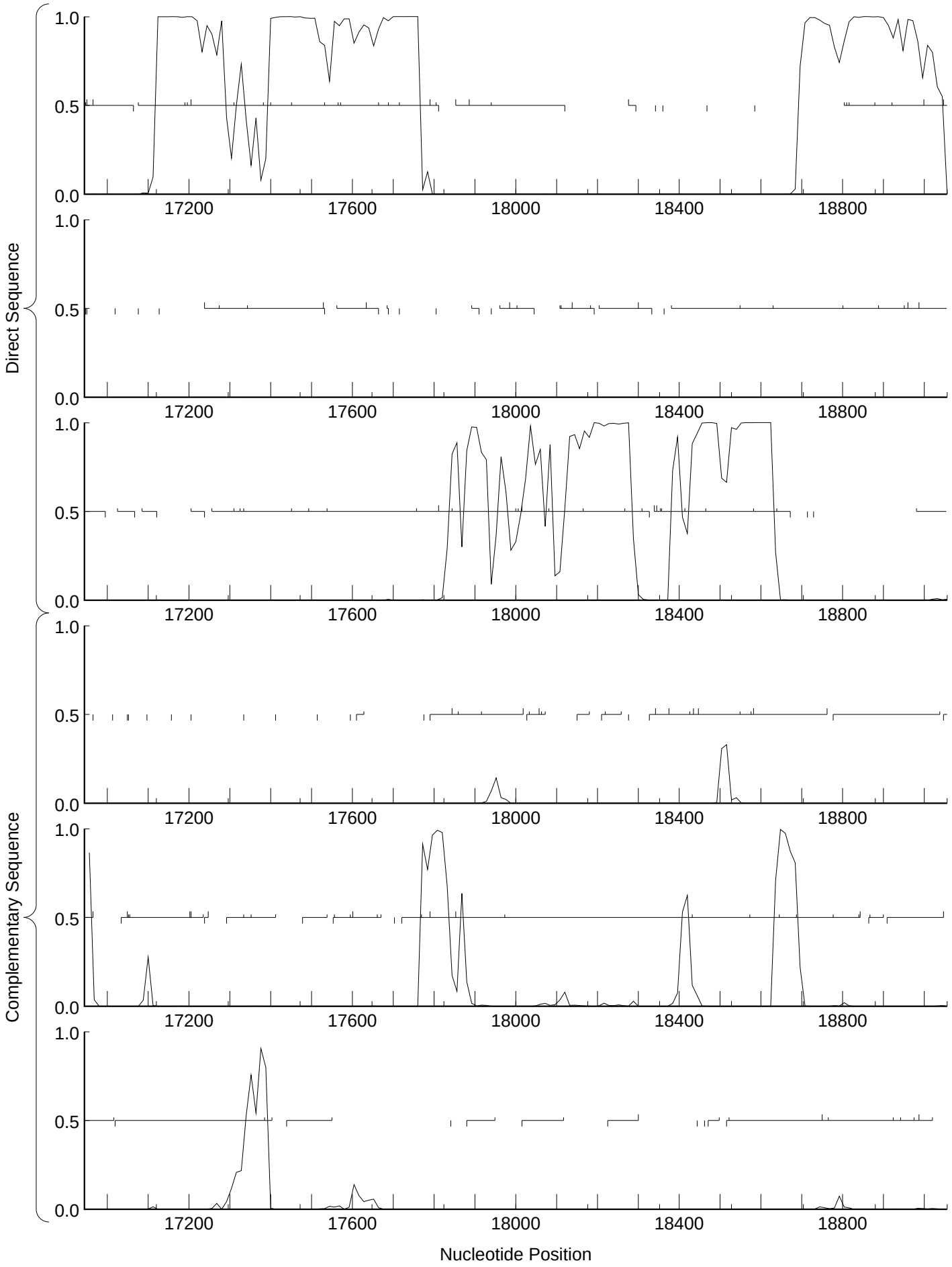
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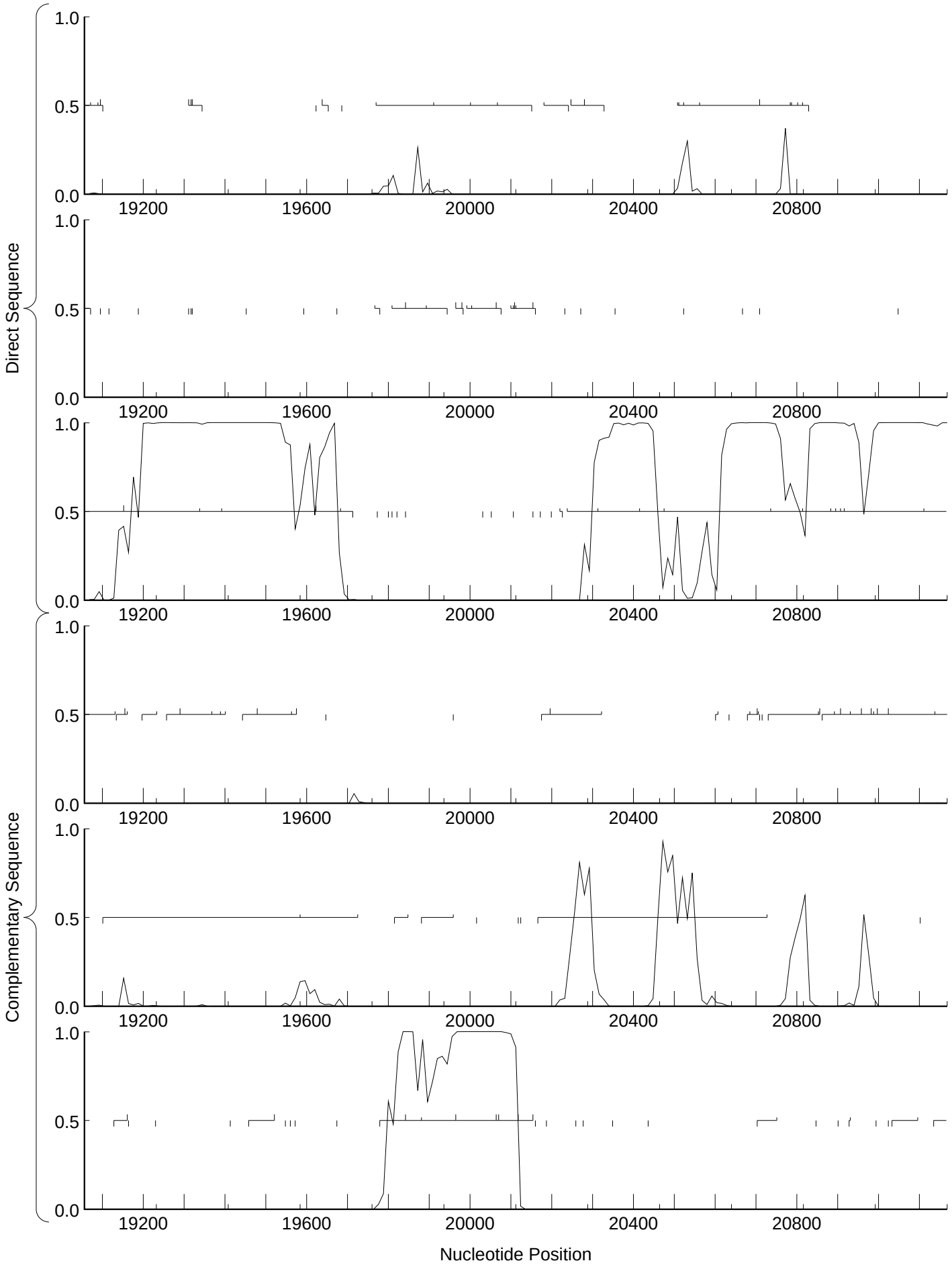

Direct Sequence
Complementary Sequence
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13200
13600
14000
14400
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Nucleotide Position


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Nucleotide Position

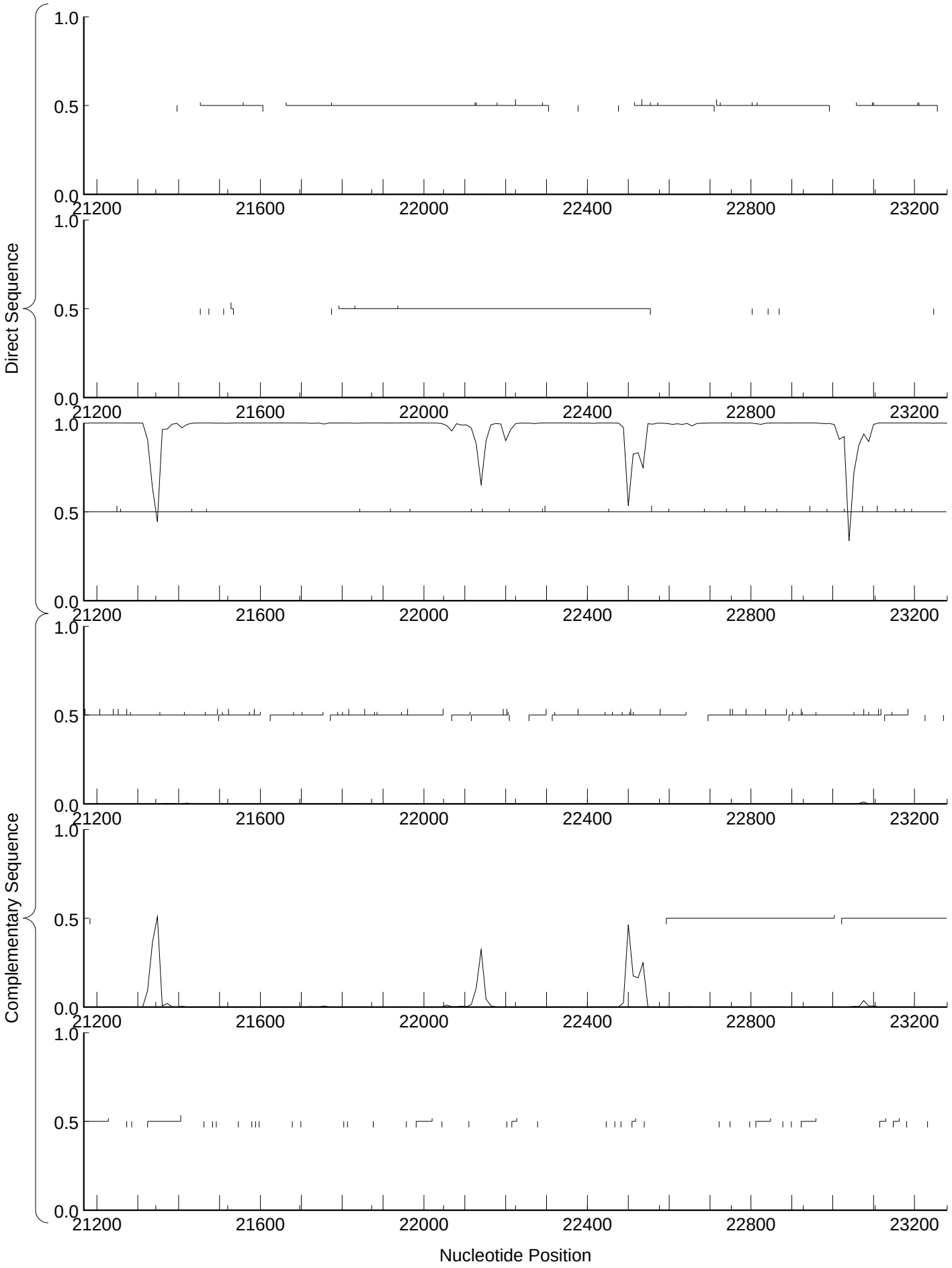
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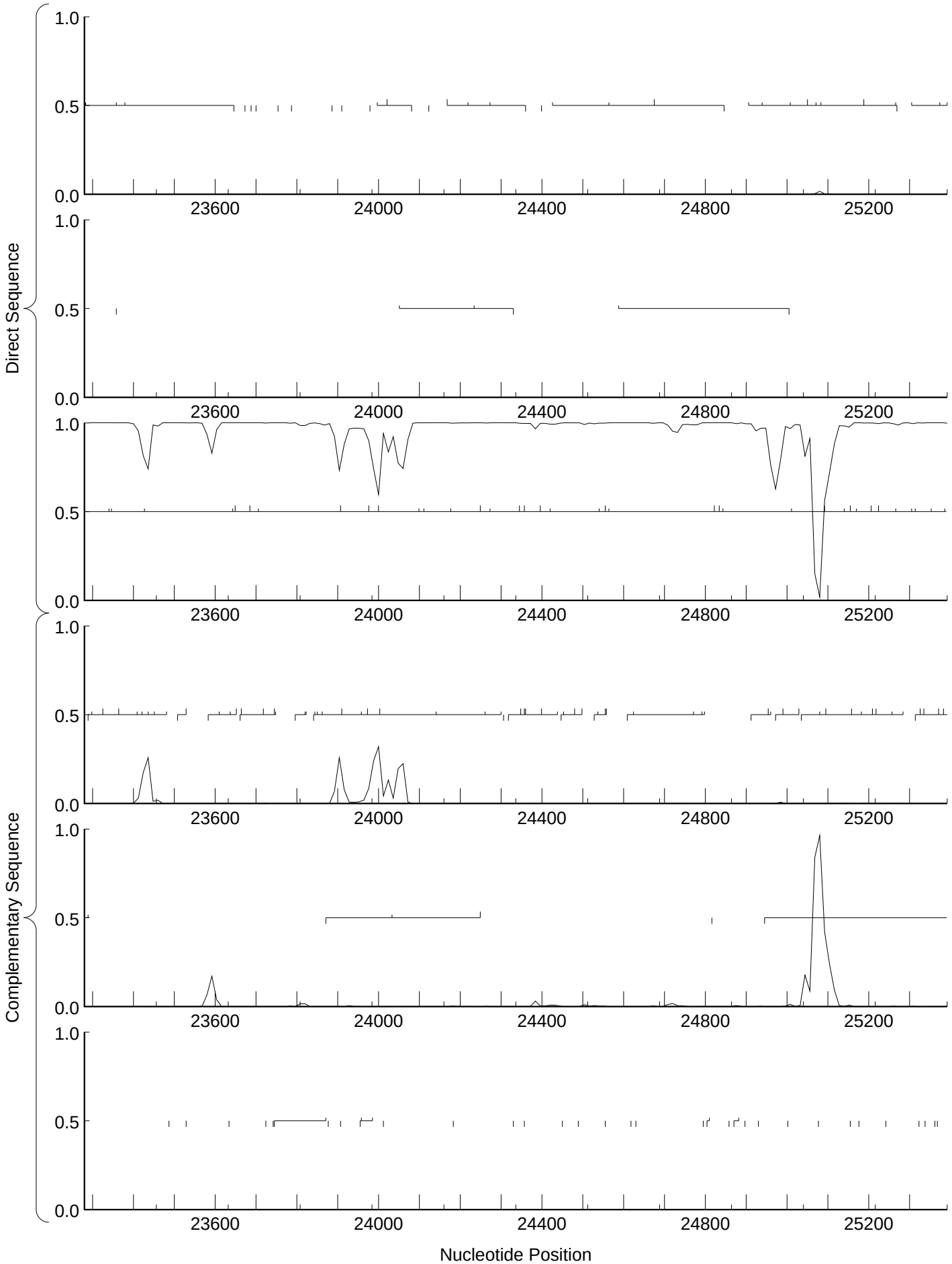
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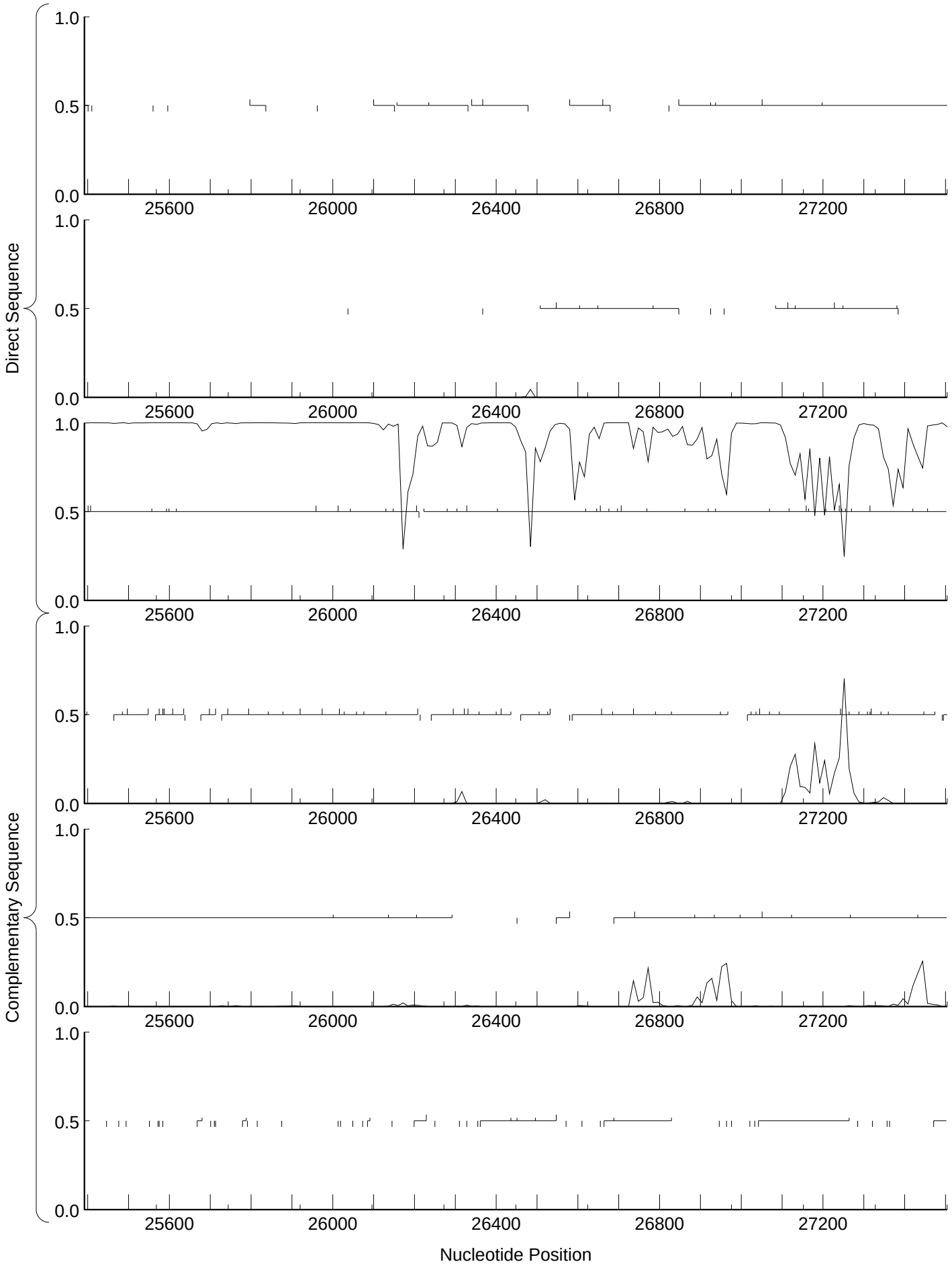
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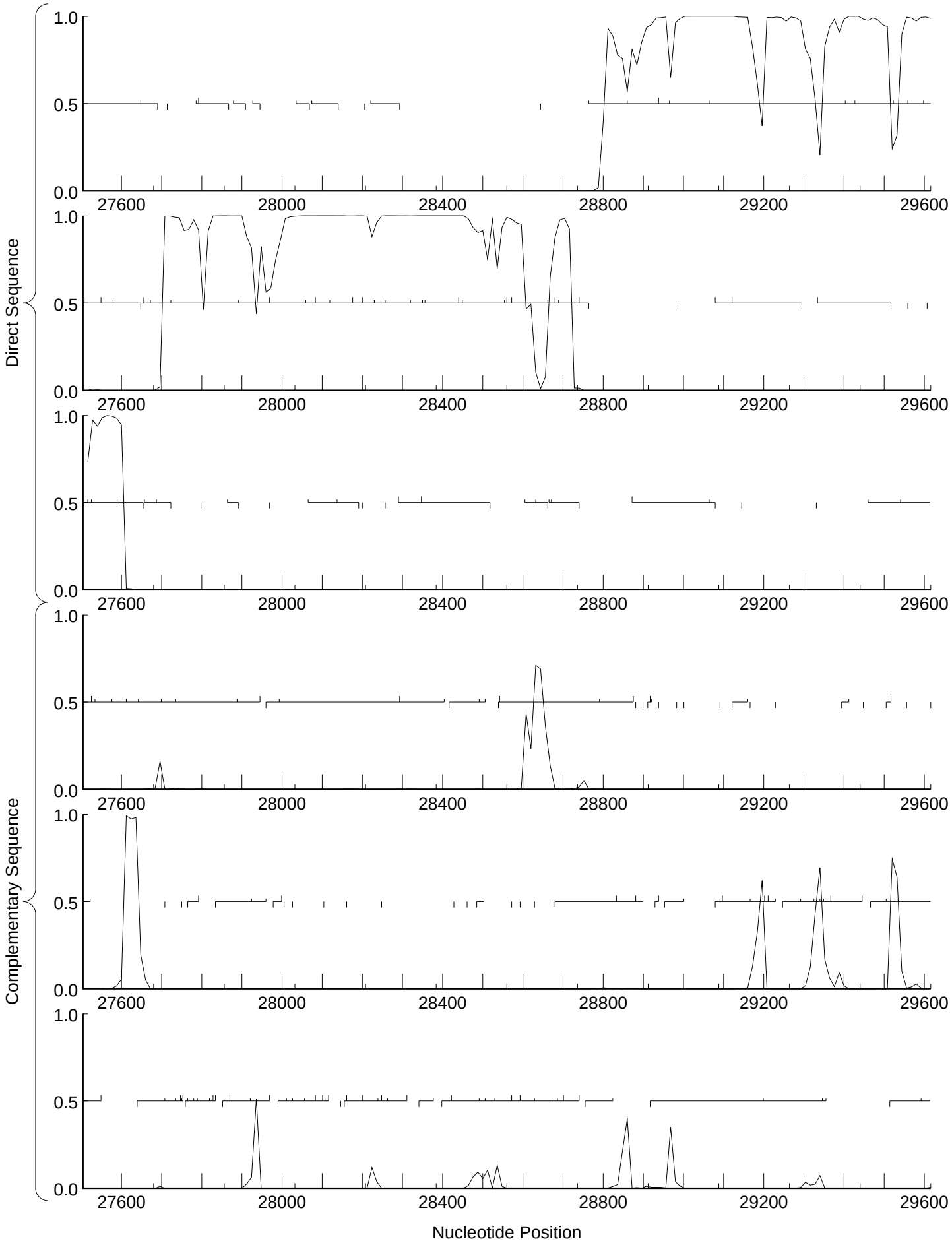
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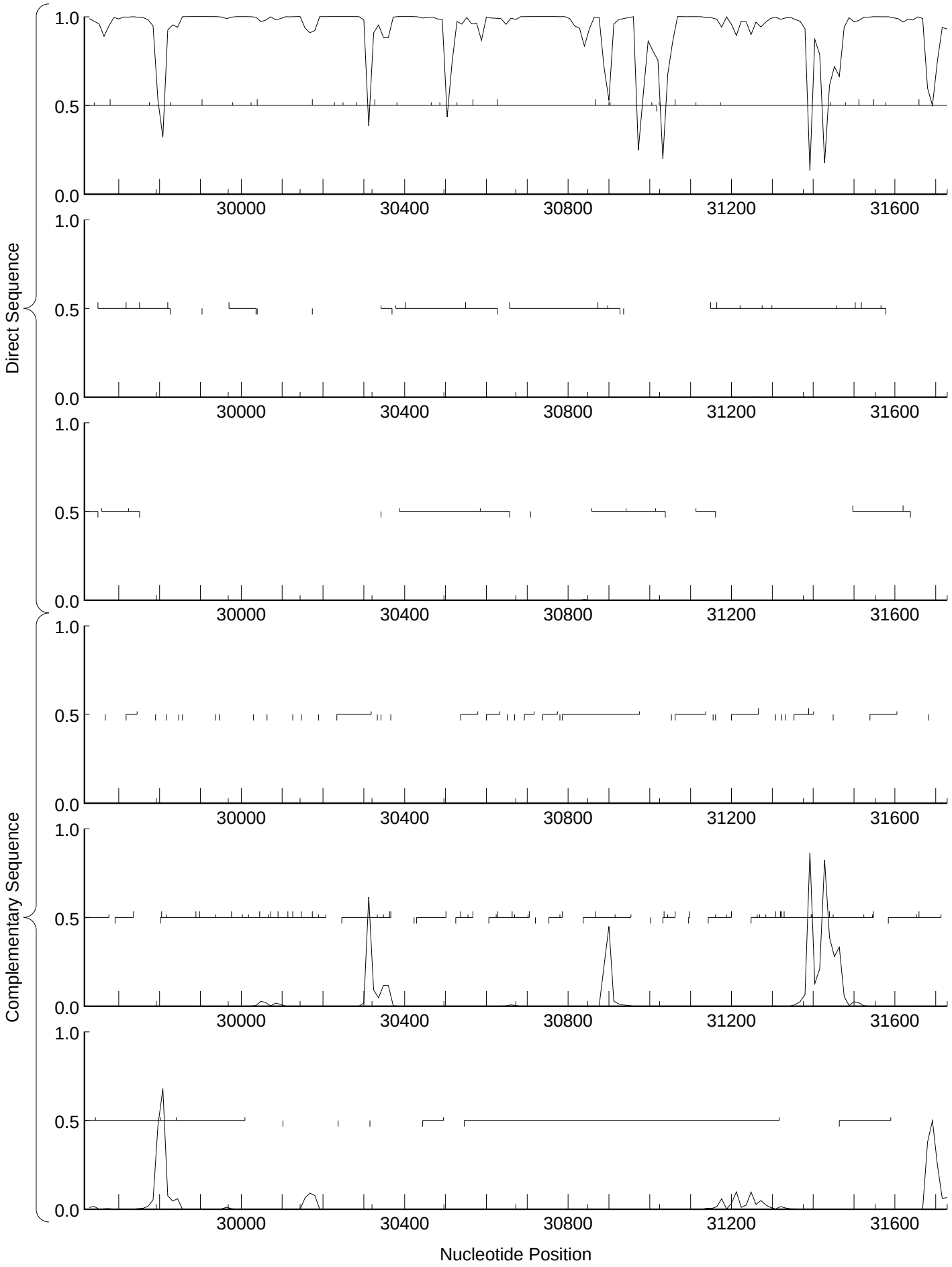
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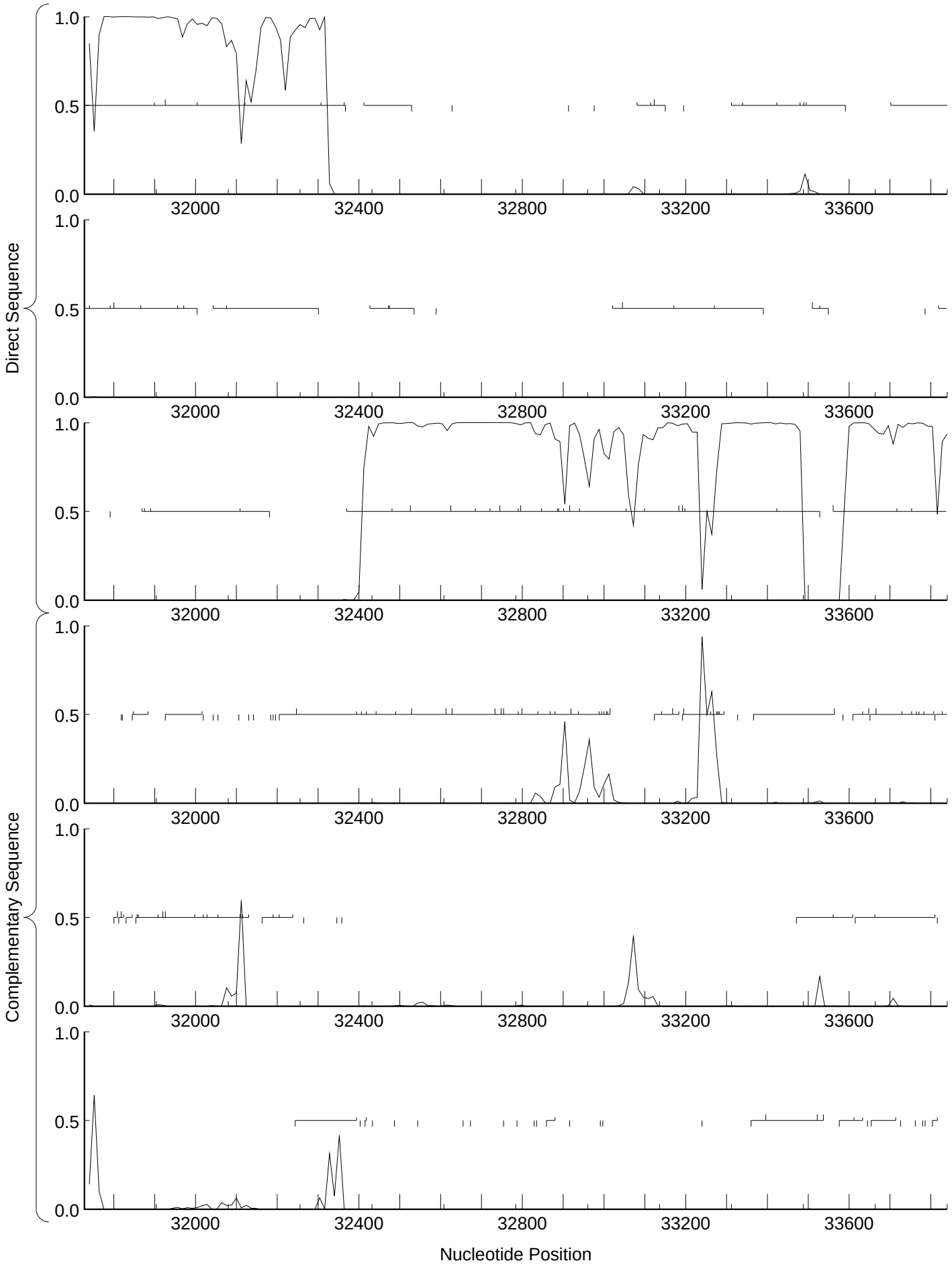
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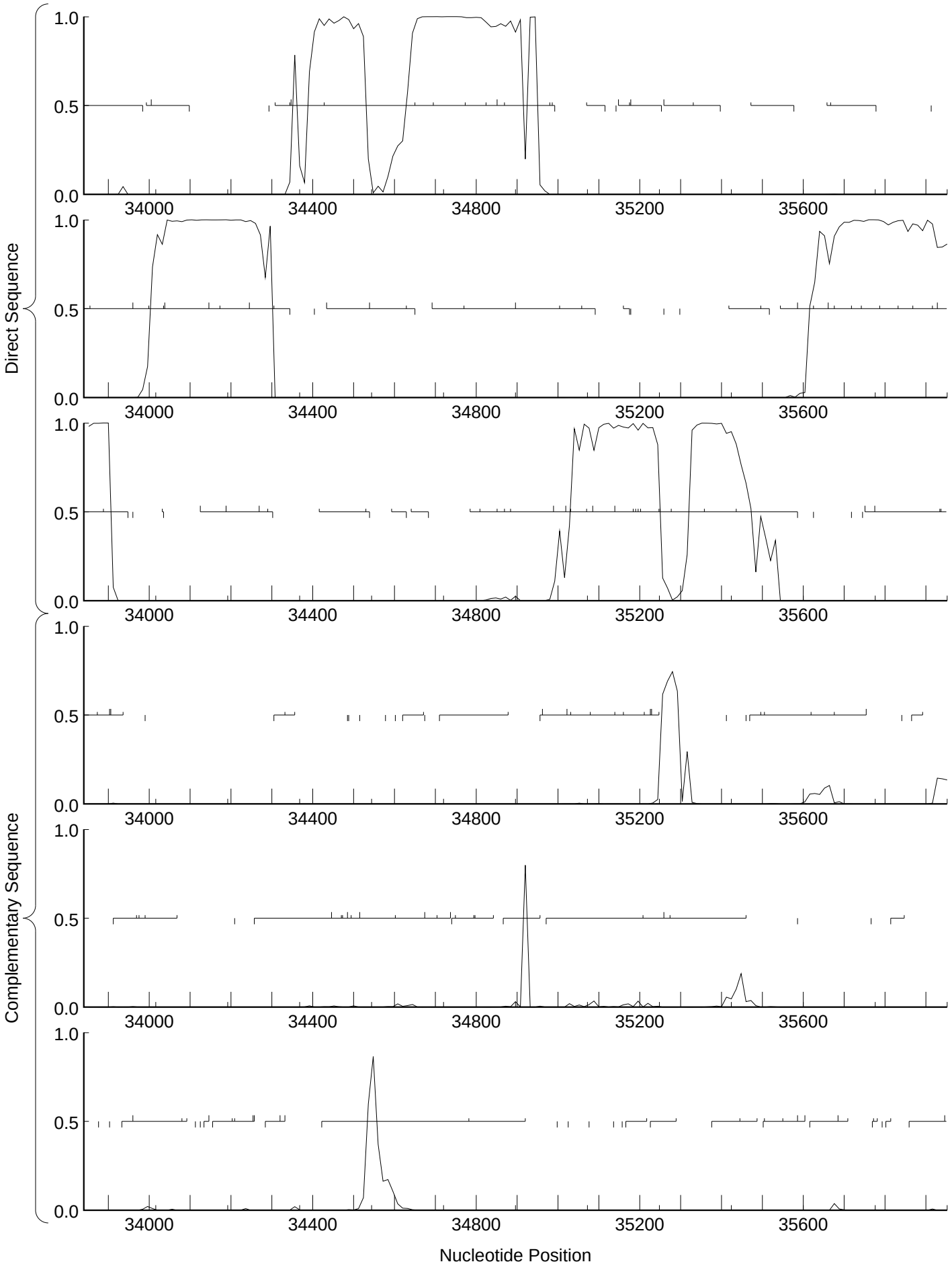
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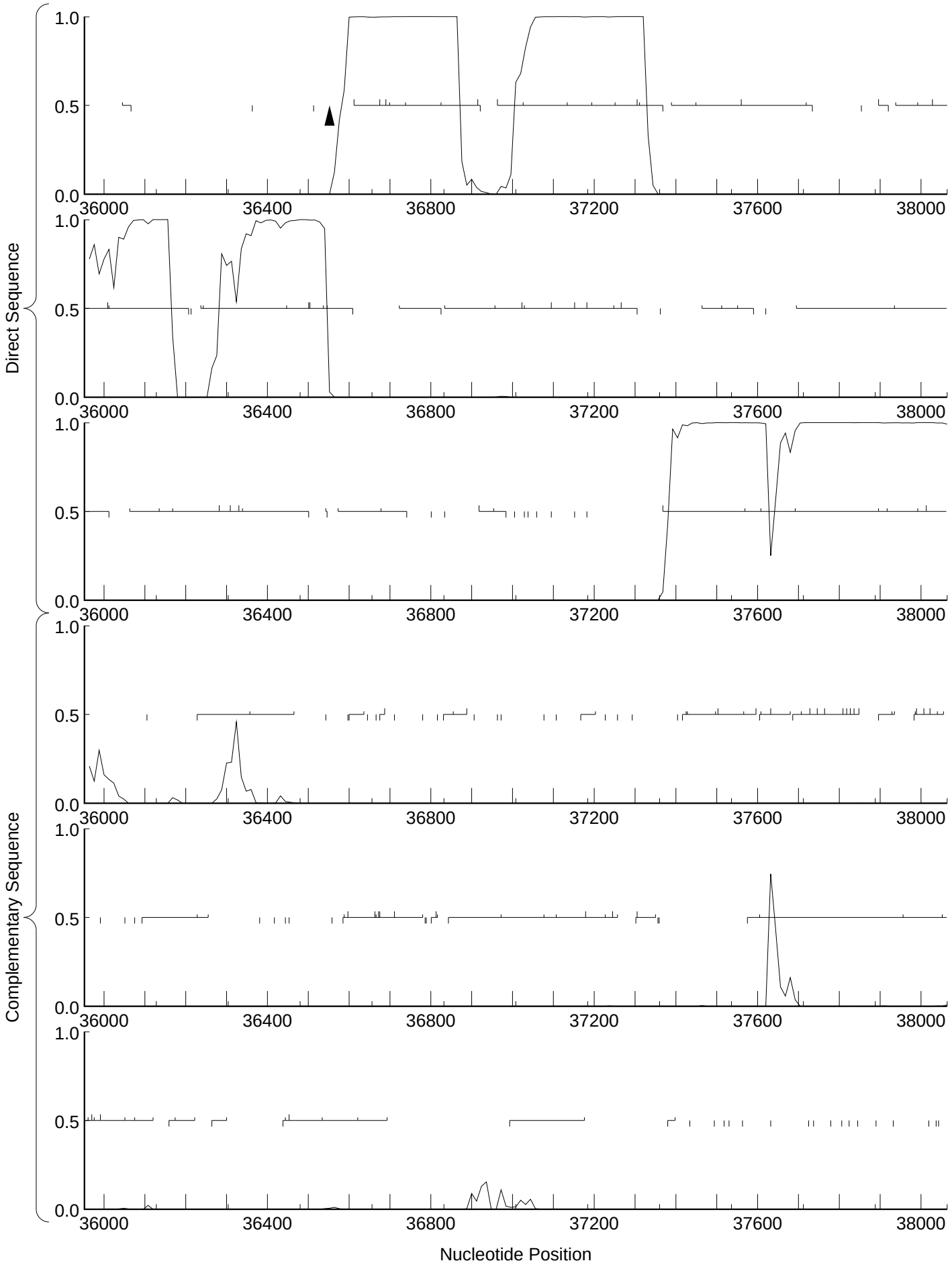
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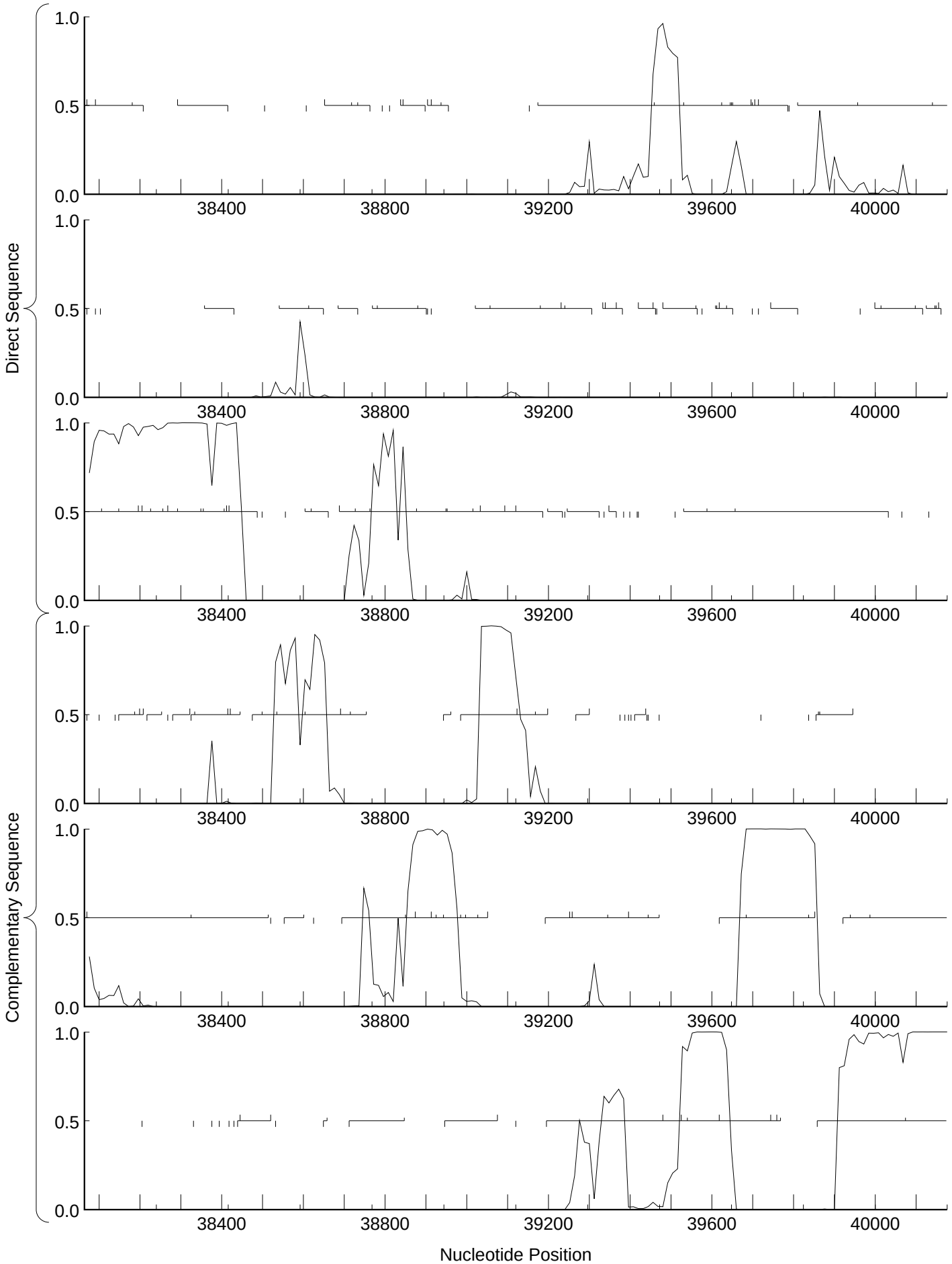


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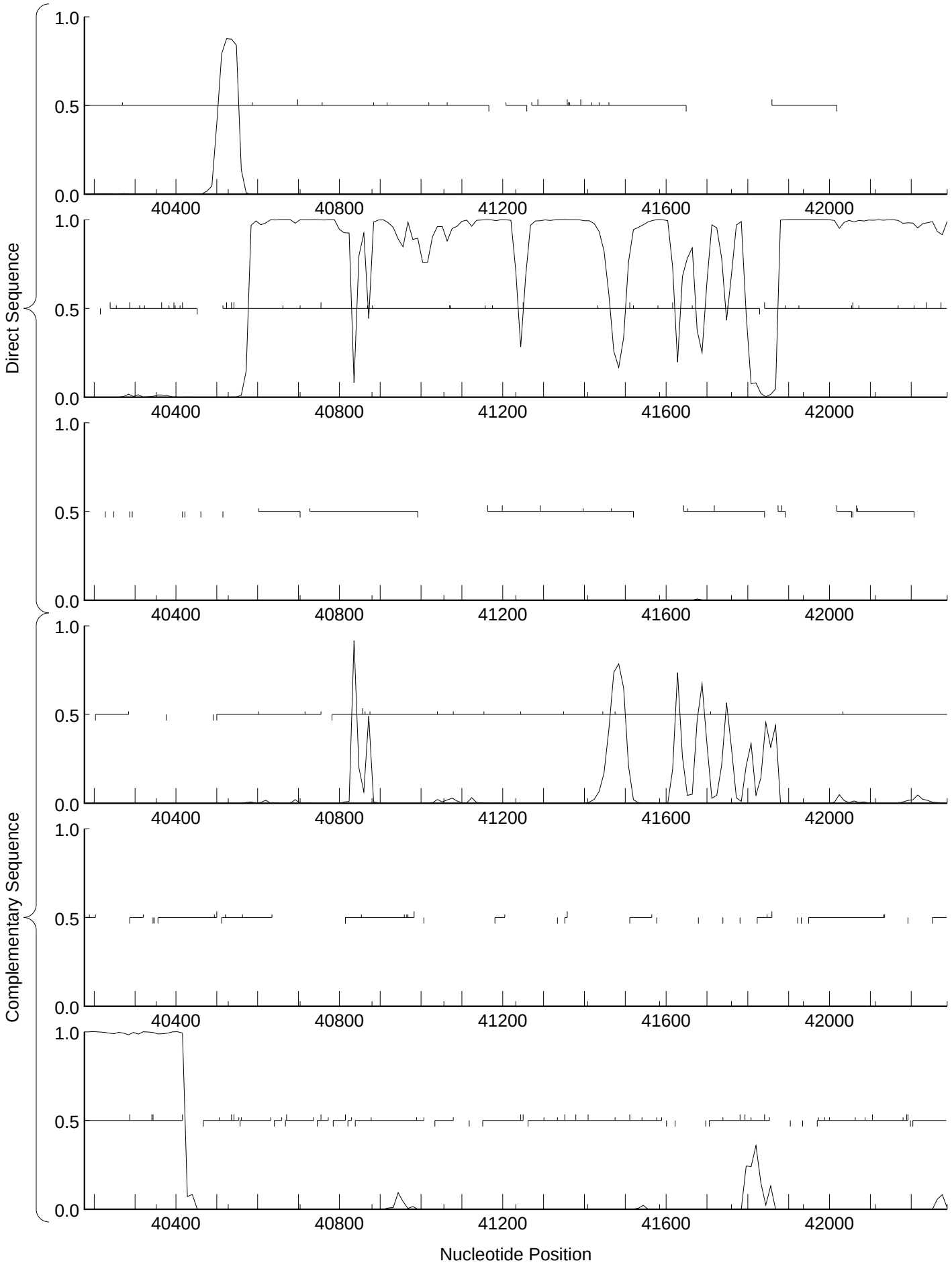


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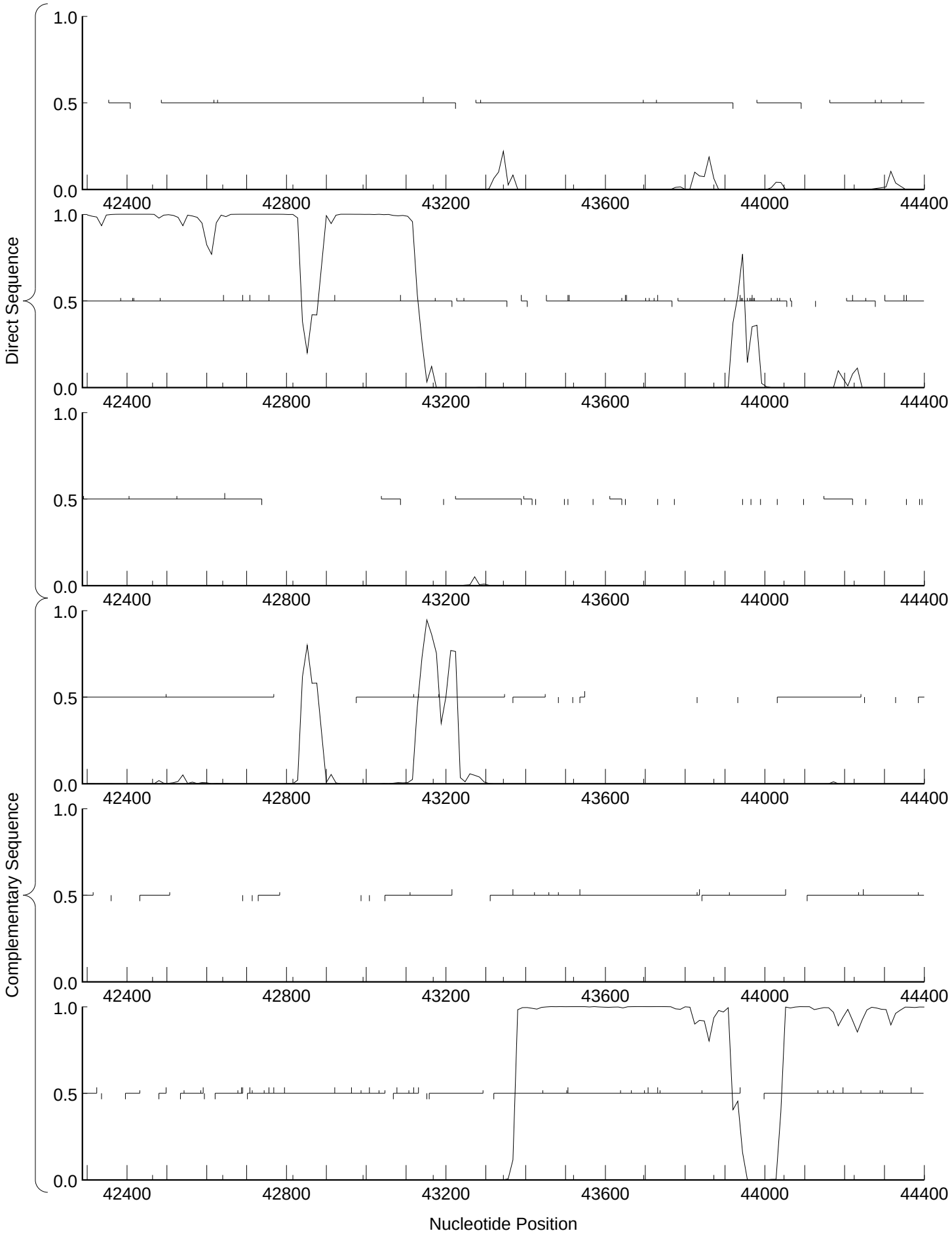



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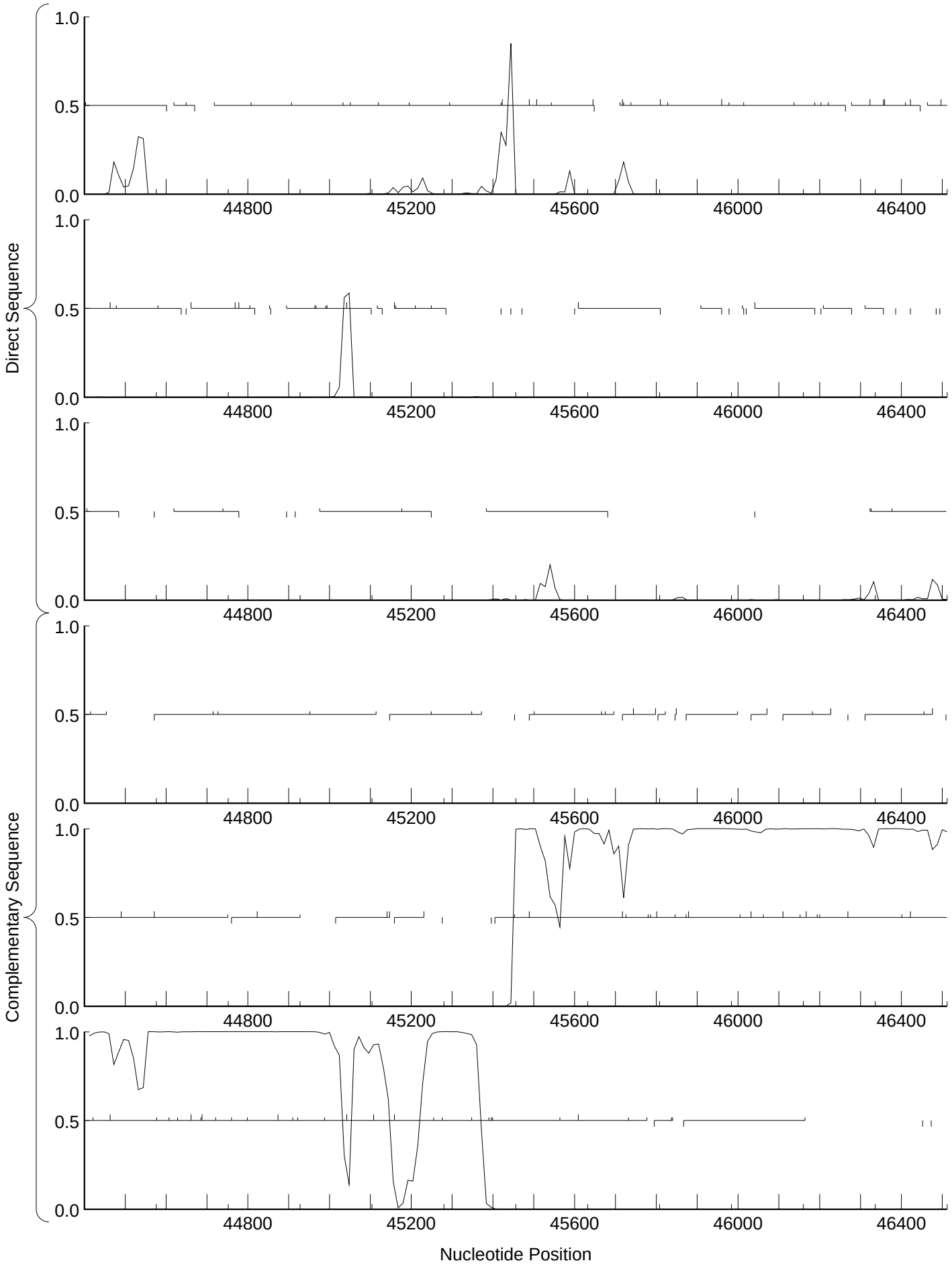
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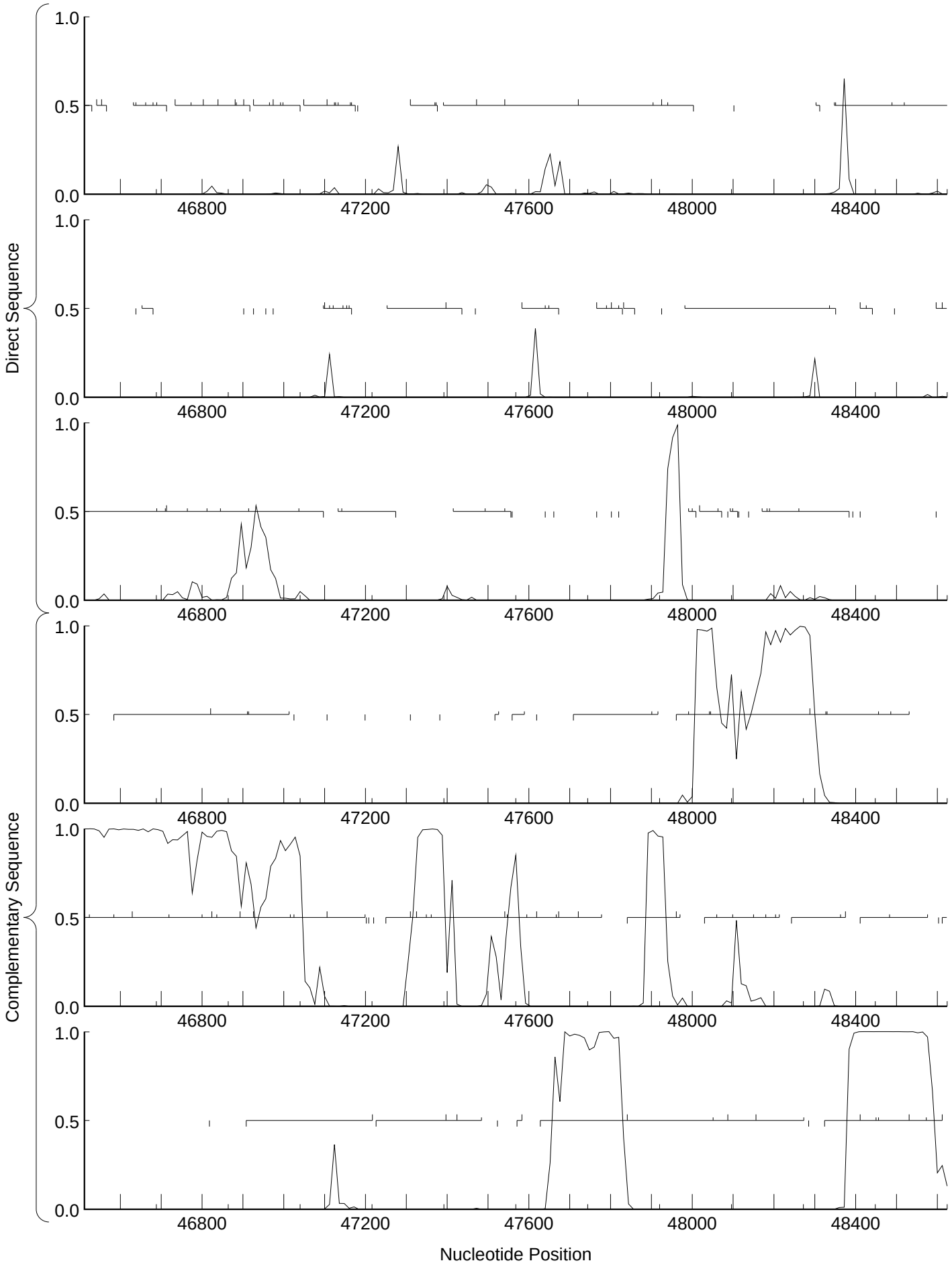
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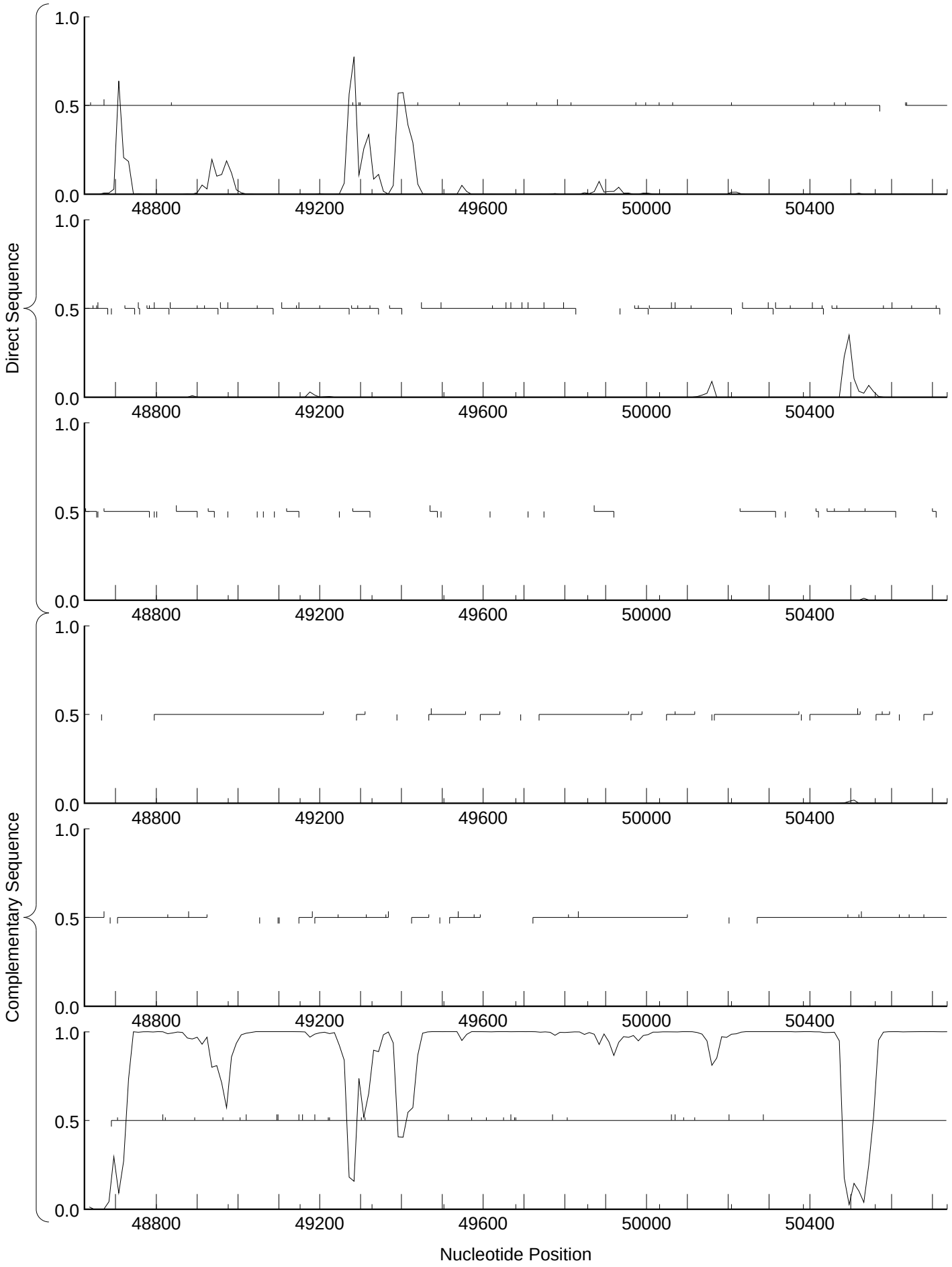
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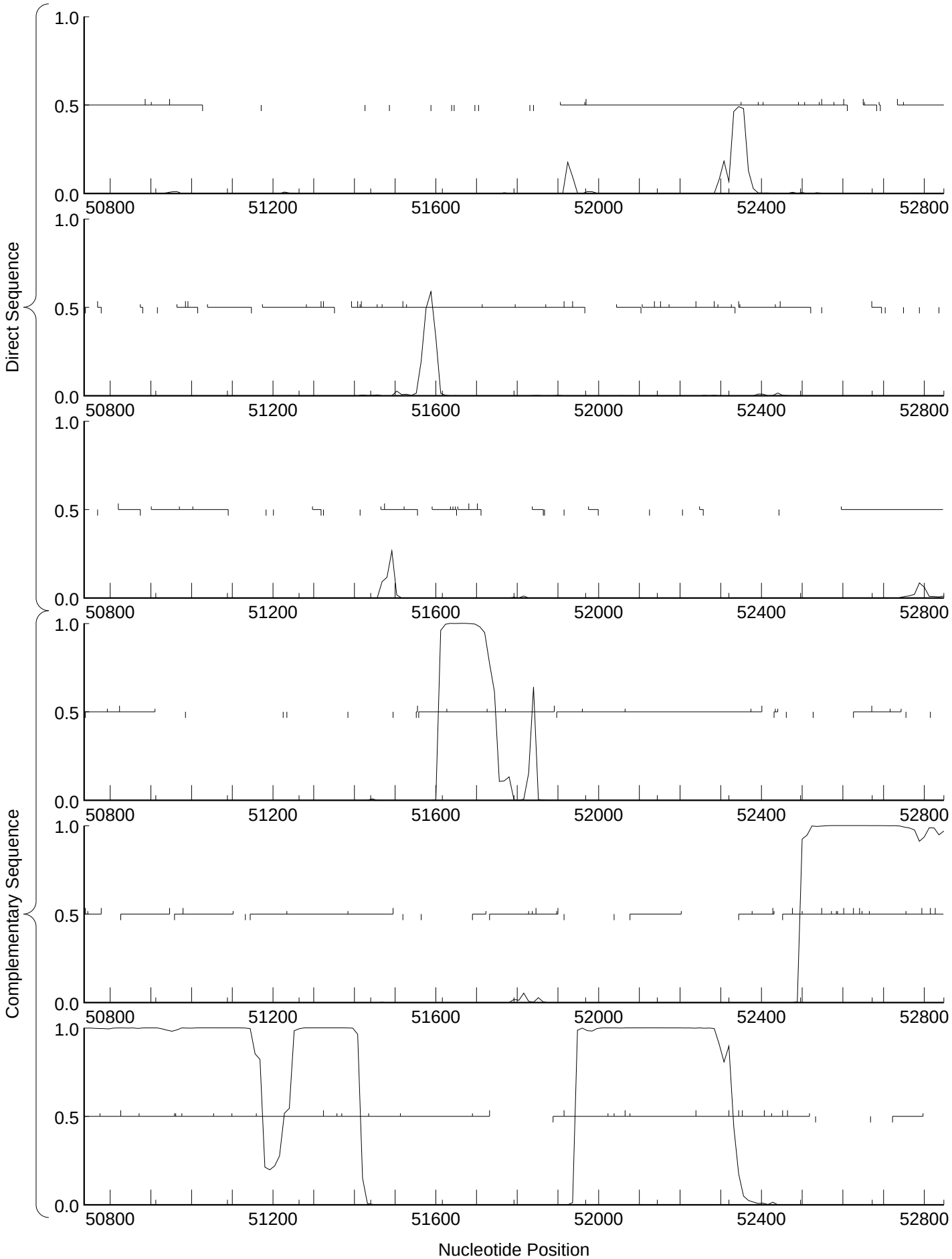
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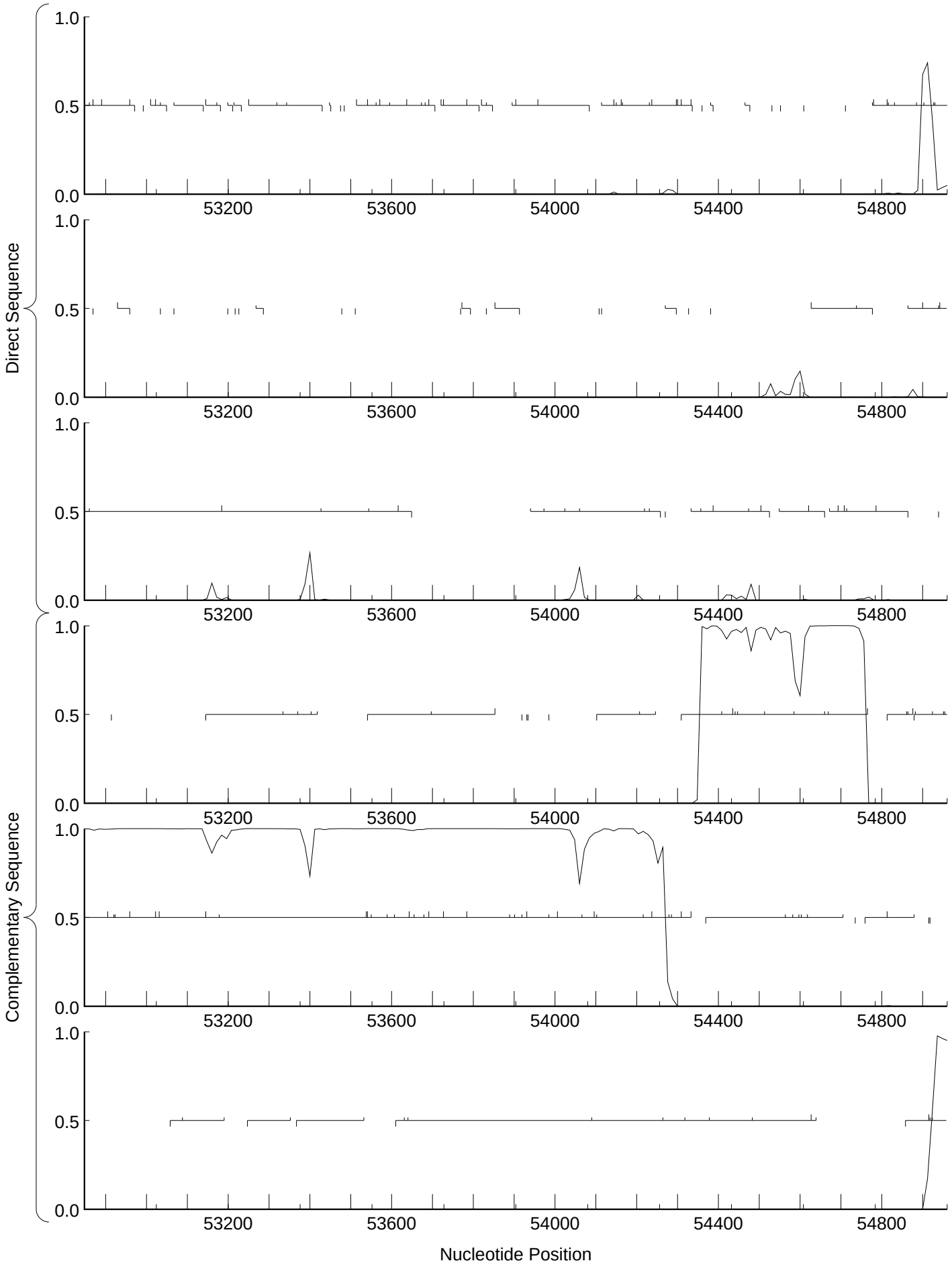
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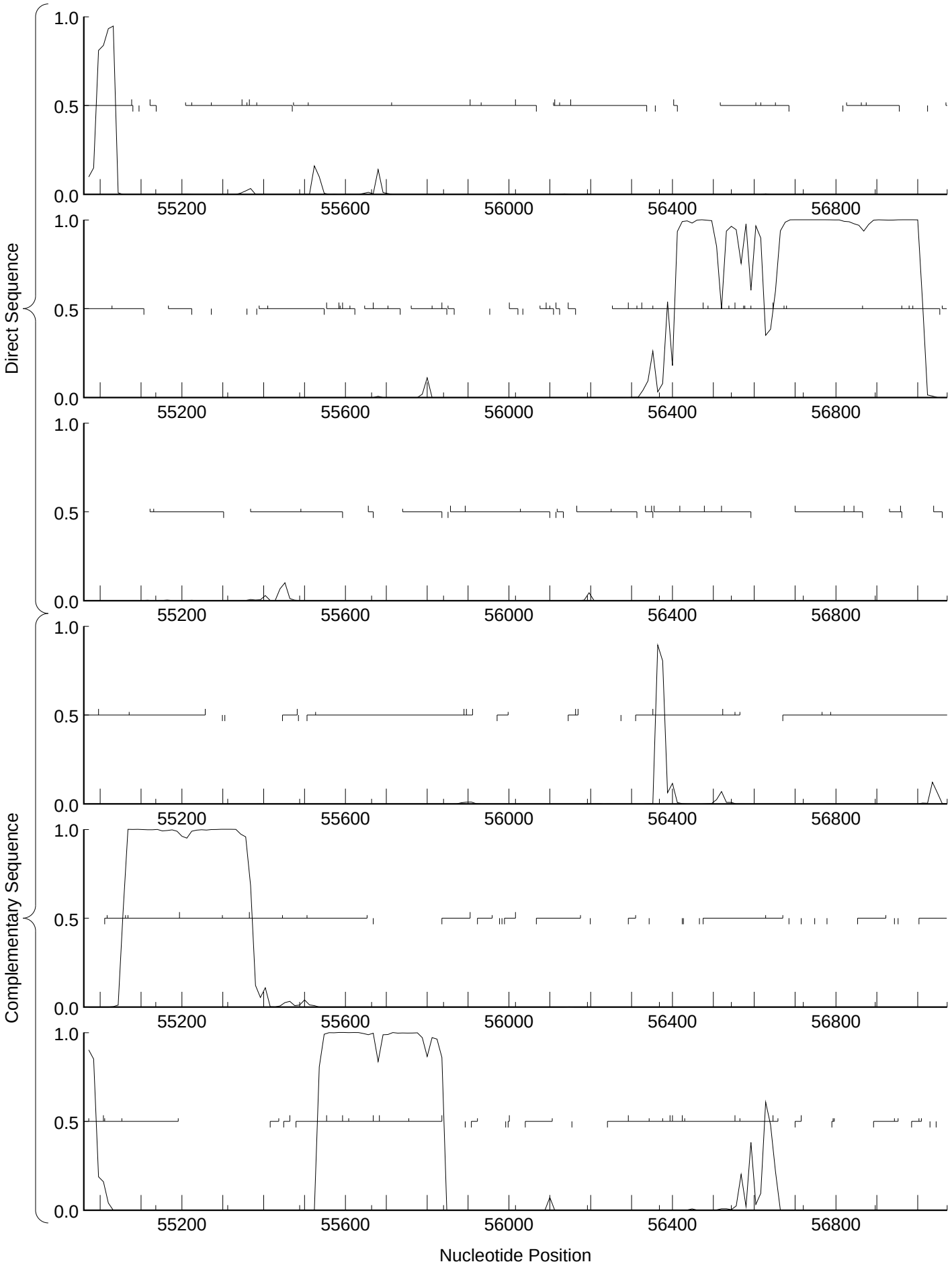
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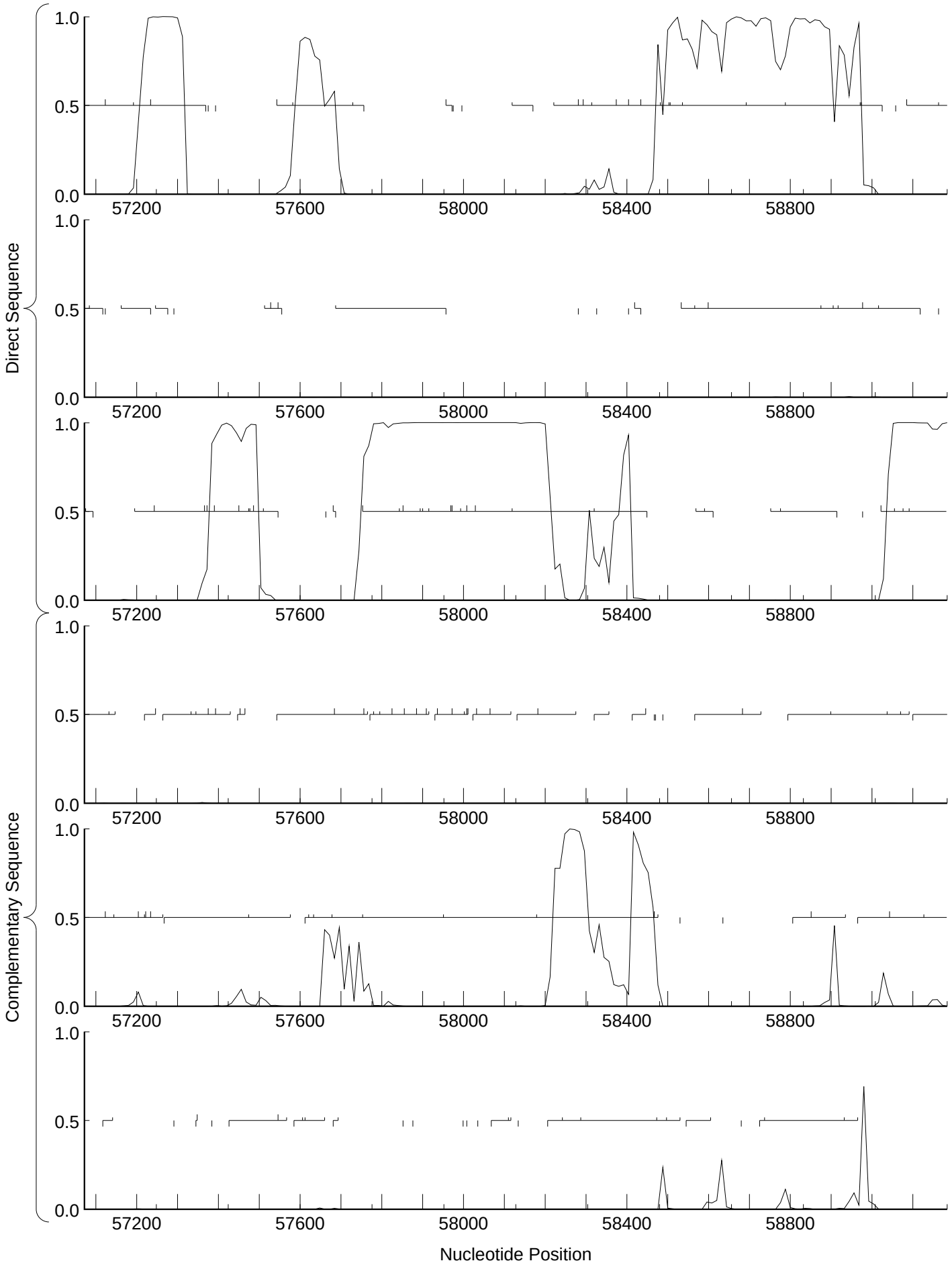
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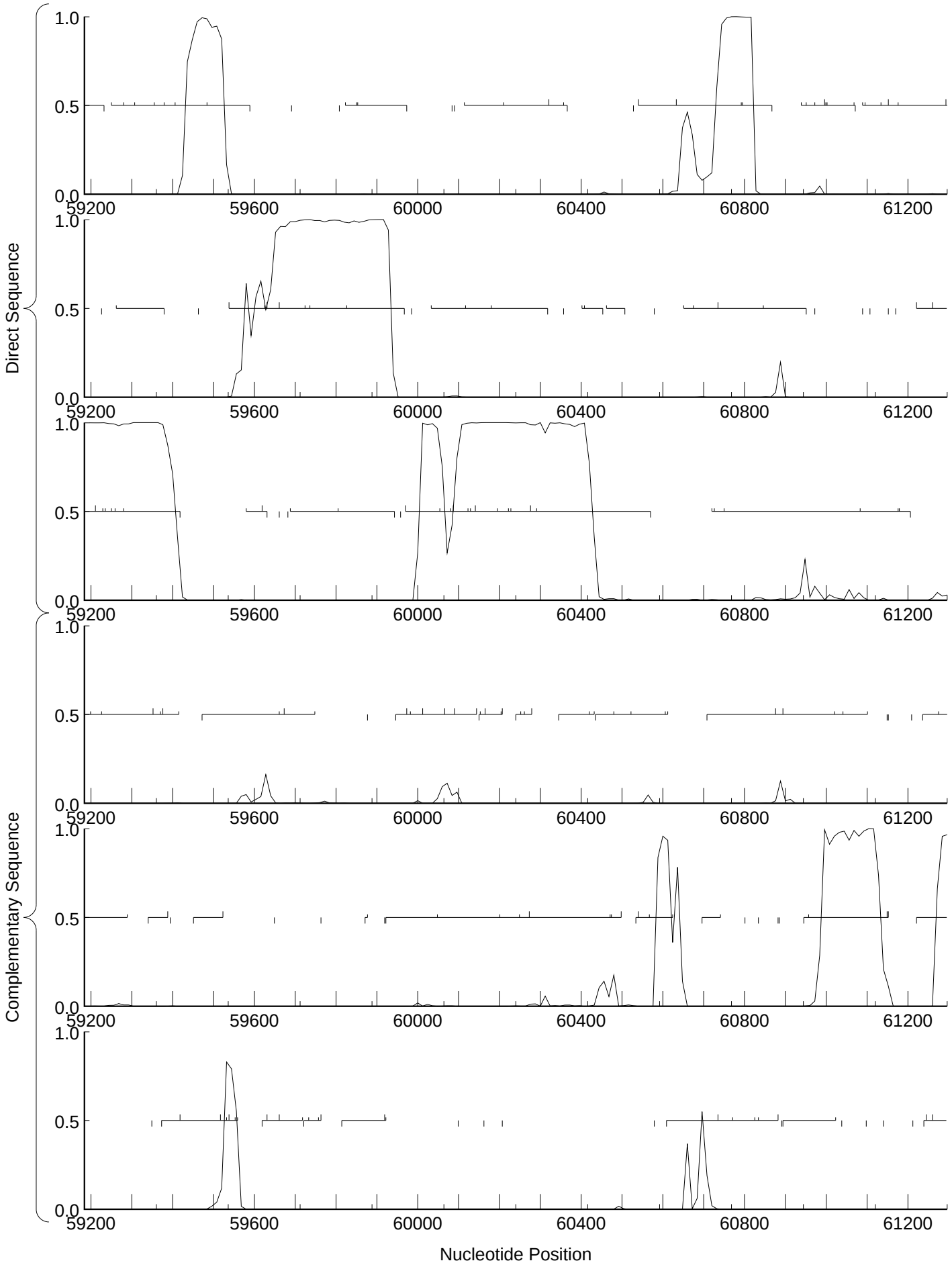
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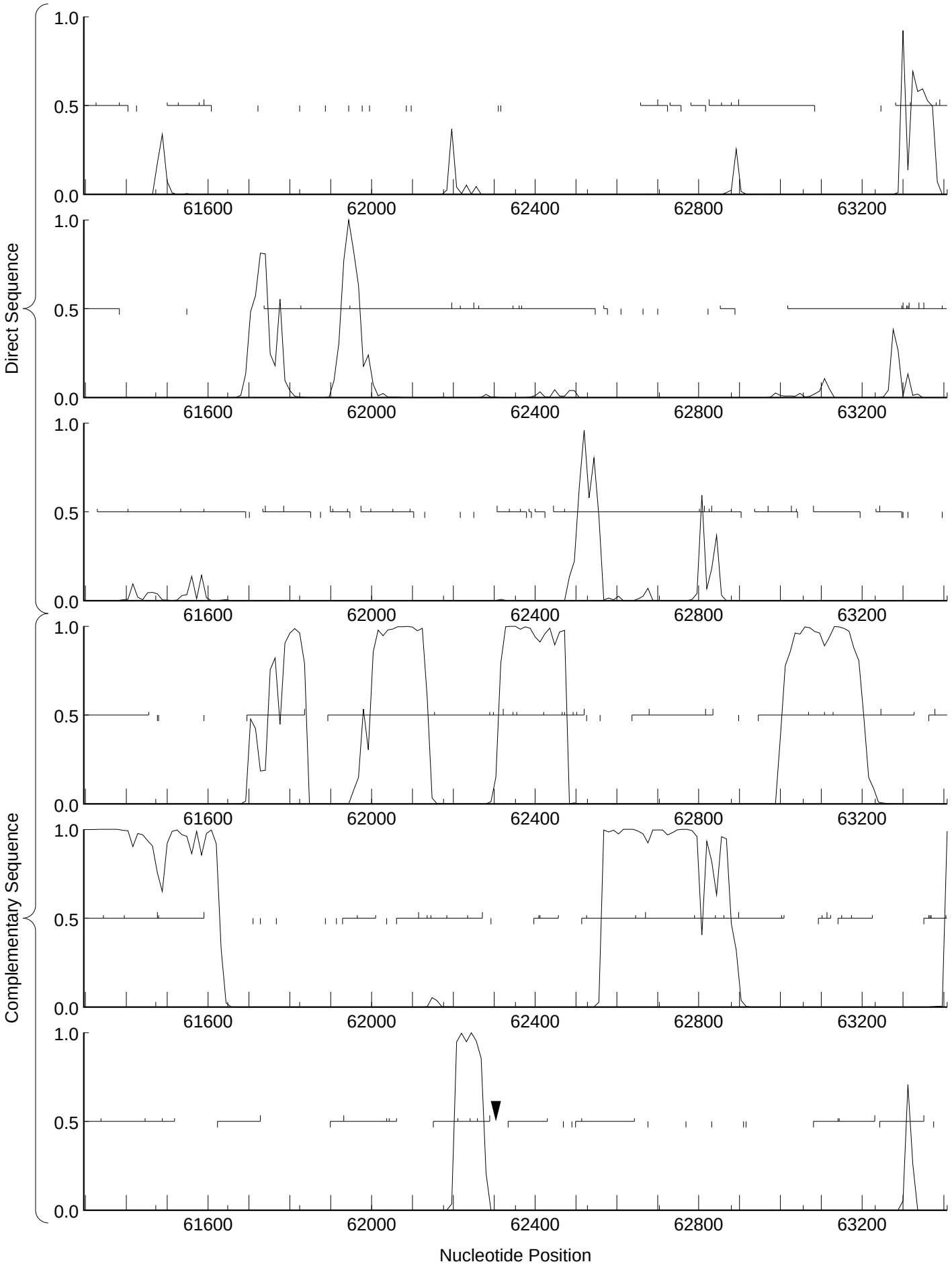
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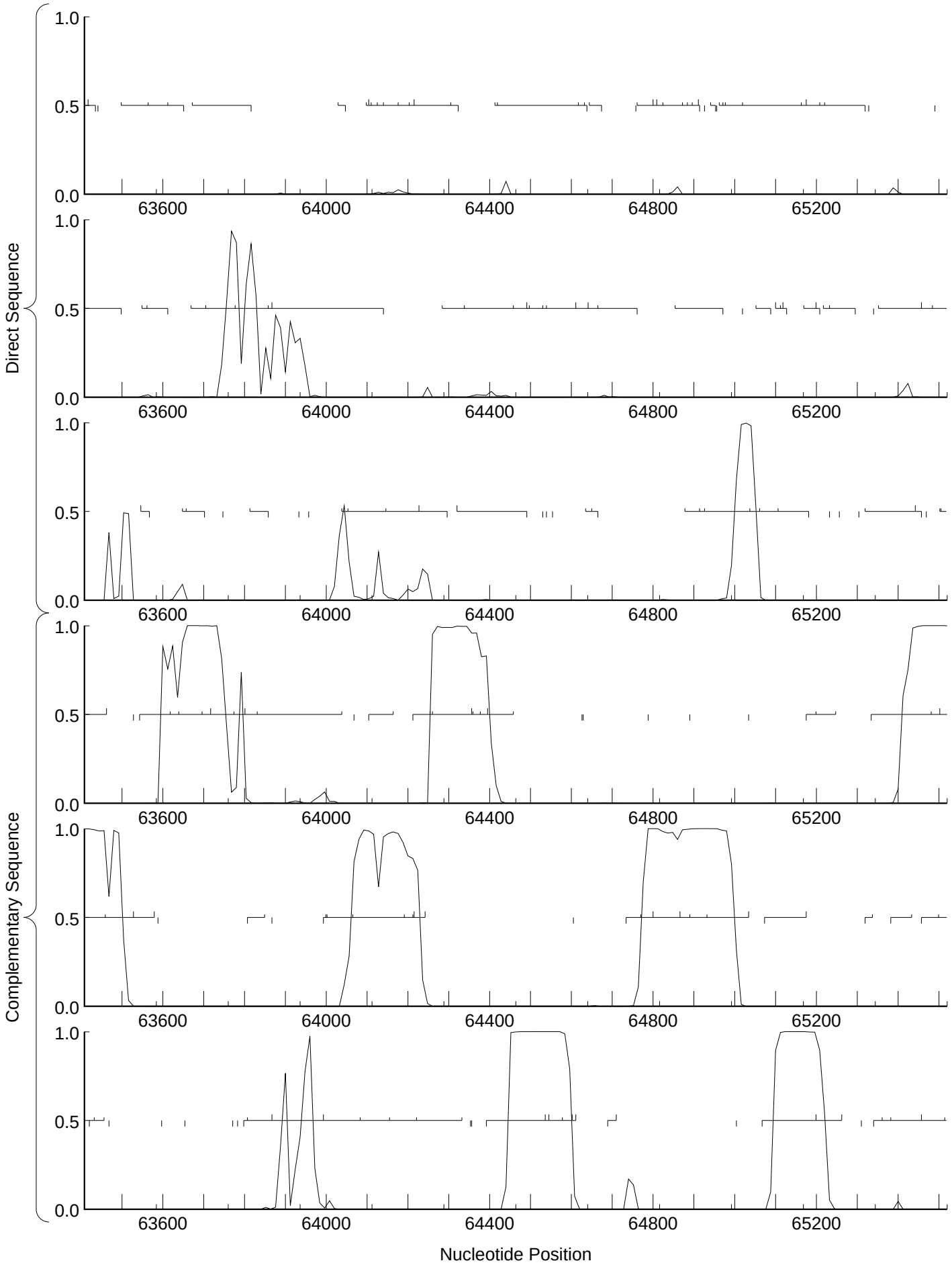
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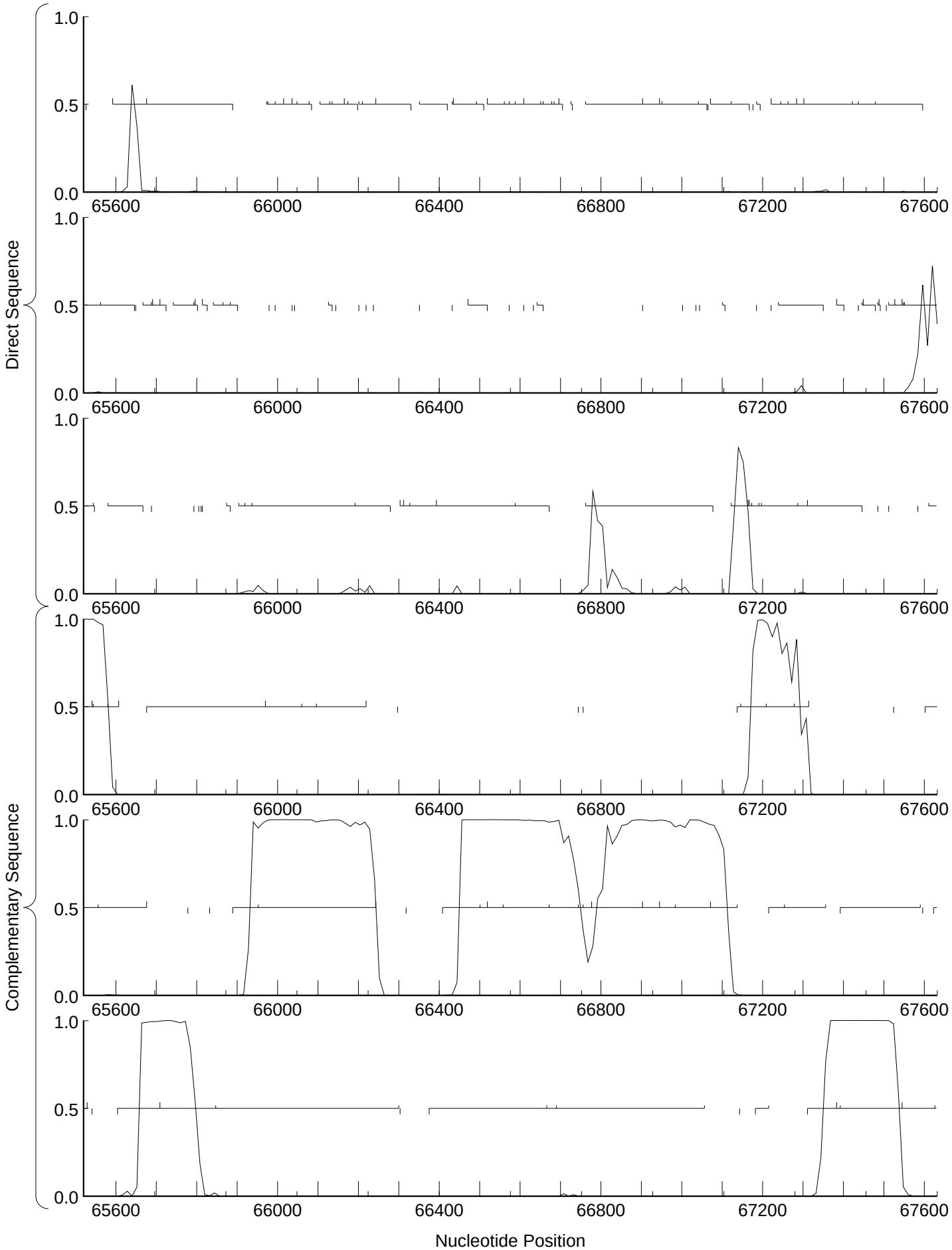
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acterium phage BlackStallion complete sequence, 68714 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 32/34



acterium phage BlackStallion complete sequence, 68714 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 33/34



acterium phage BlackStallion complete sequence, 68714 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 34/34

