GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Microbacterium phage Gargoyle complete sequence, 41803 bp, circularly permuted, Cluster EA1
Analysis Date    :  3/28/24 at 14:52:44
Pages            :  21
Sequence Length  :  41803 bp
GC Content       :   63.51%

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

PS-Version       :  1.2

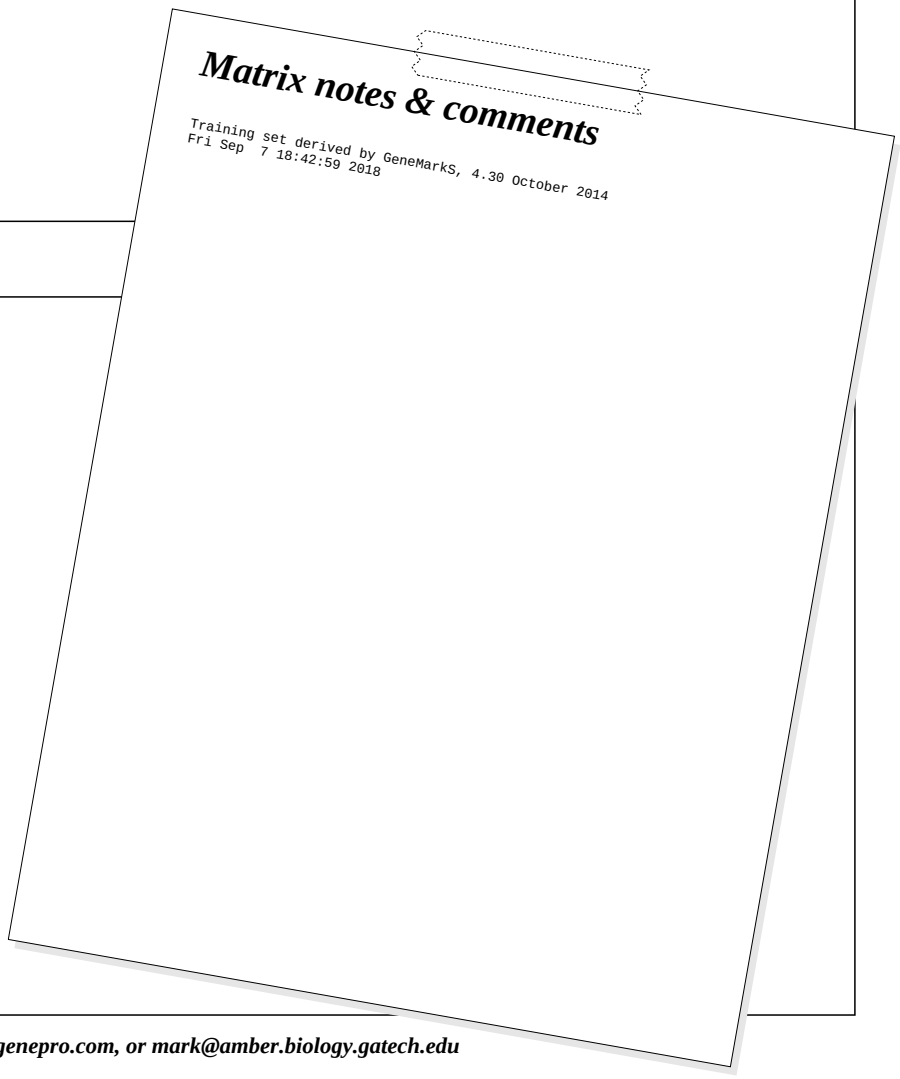
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Author     :  -

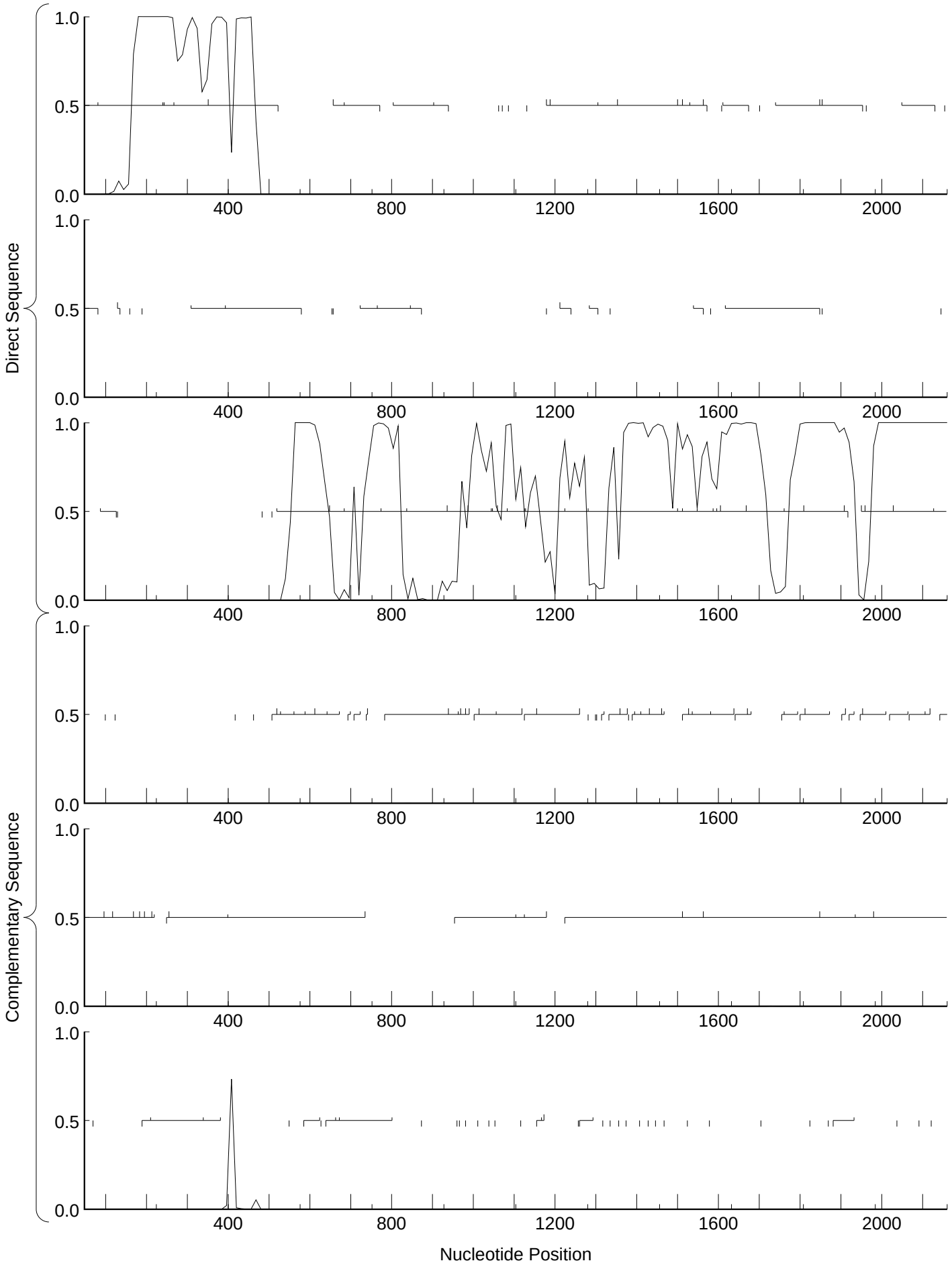
Order      :  4


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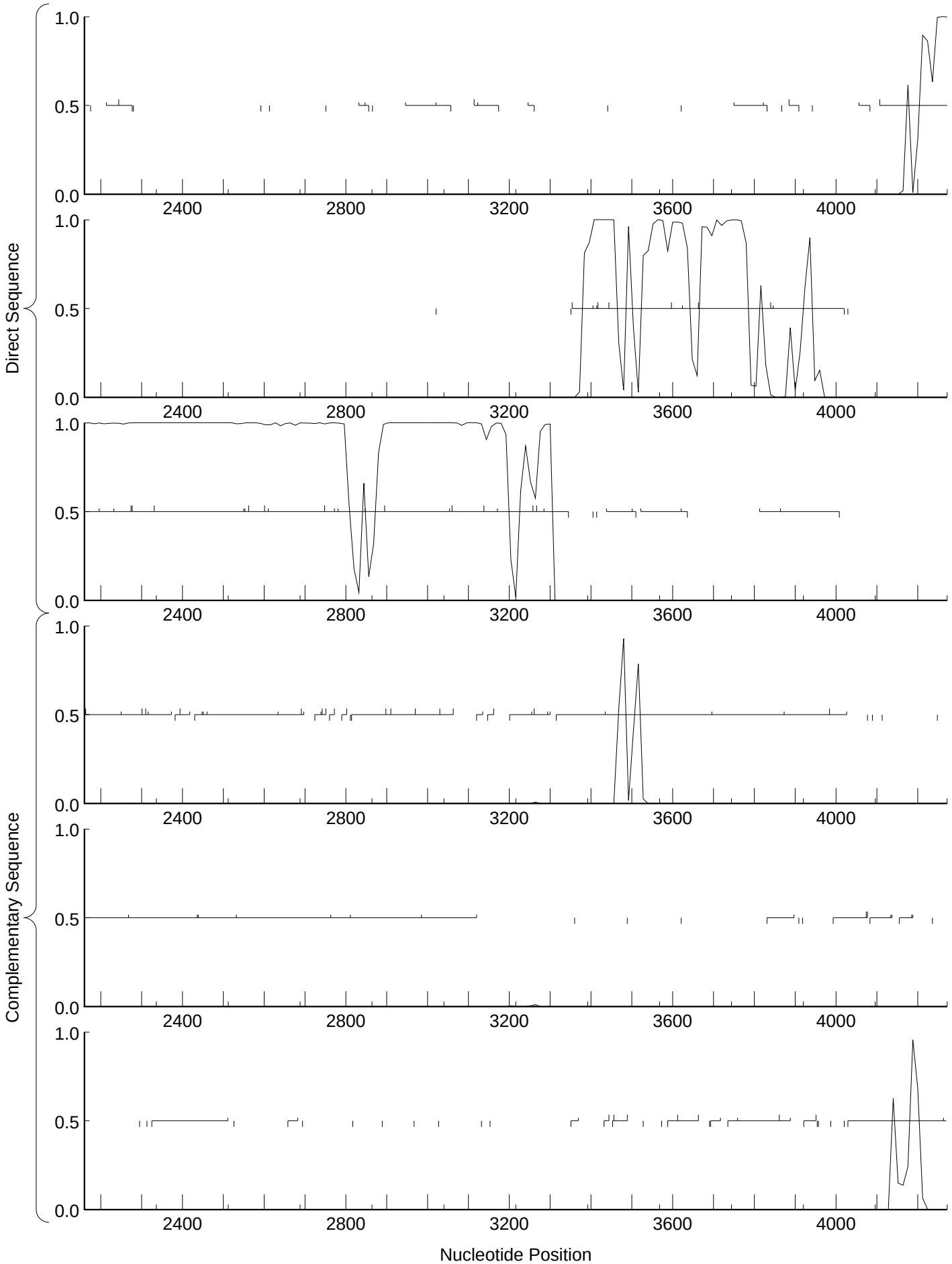


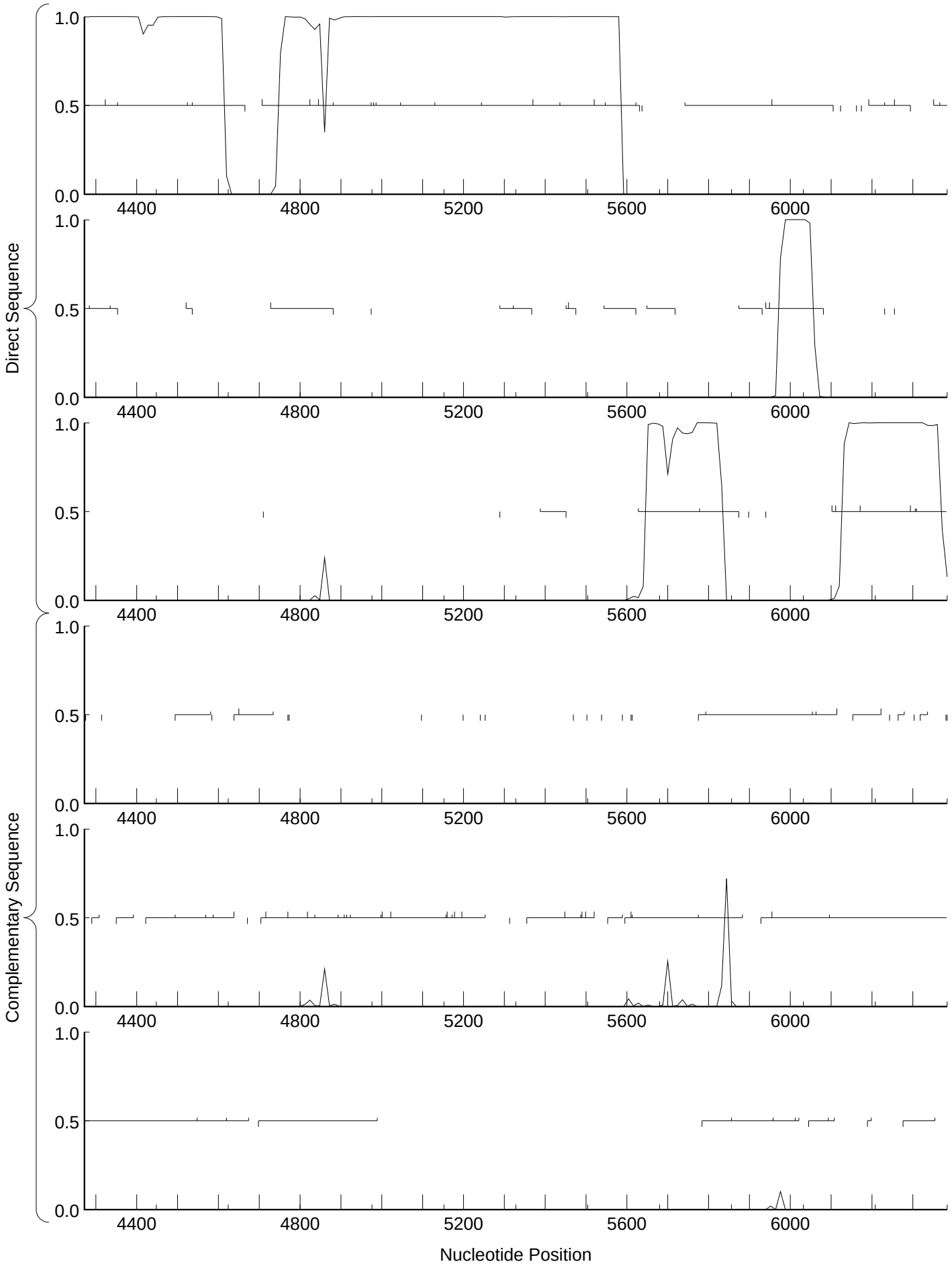
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

robacterium phage Gargoyle complete sequence, 41803 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 2/21

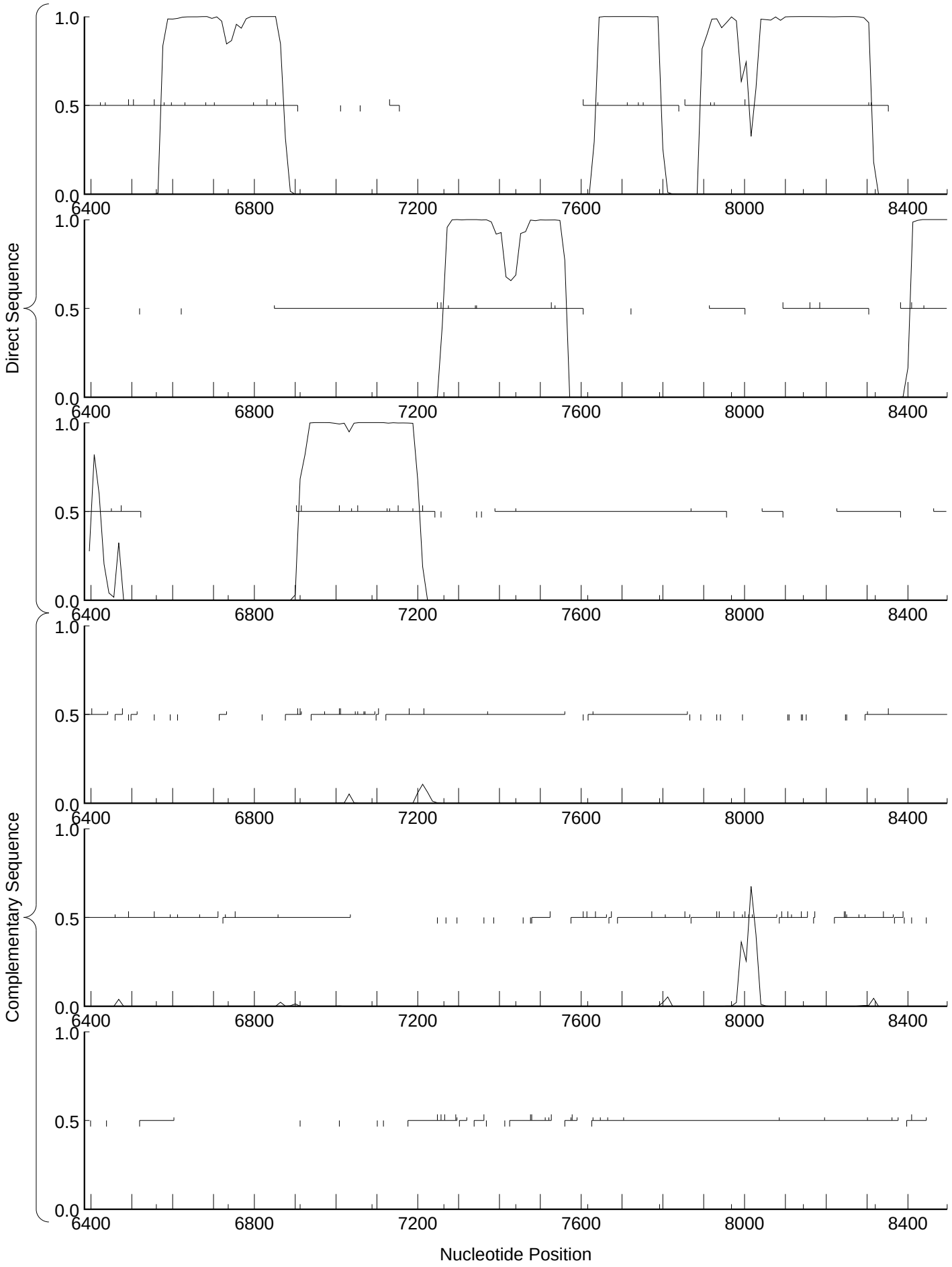


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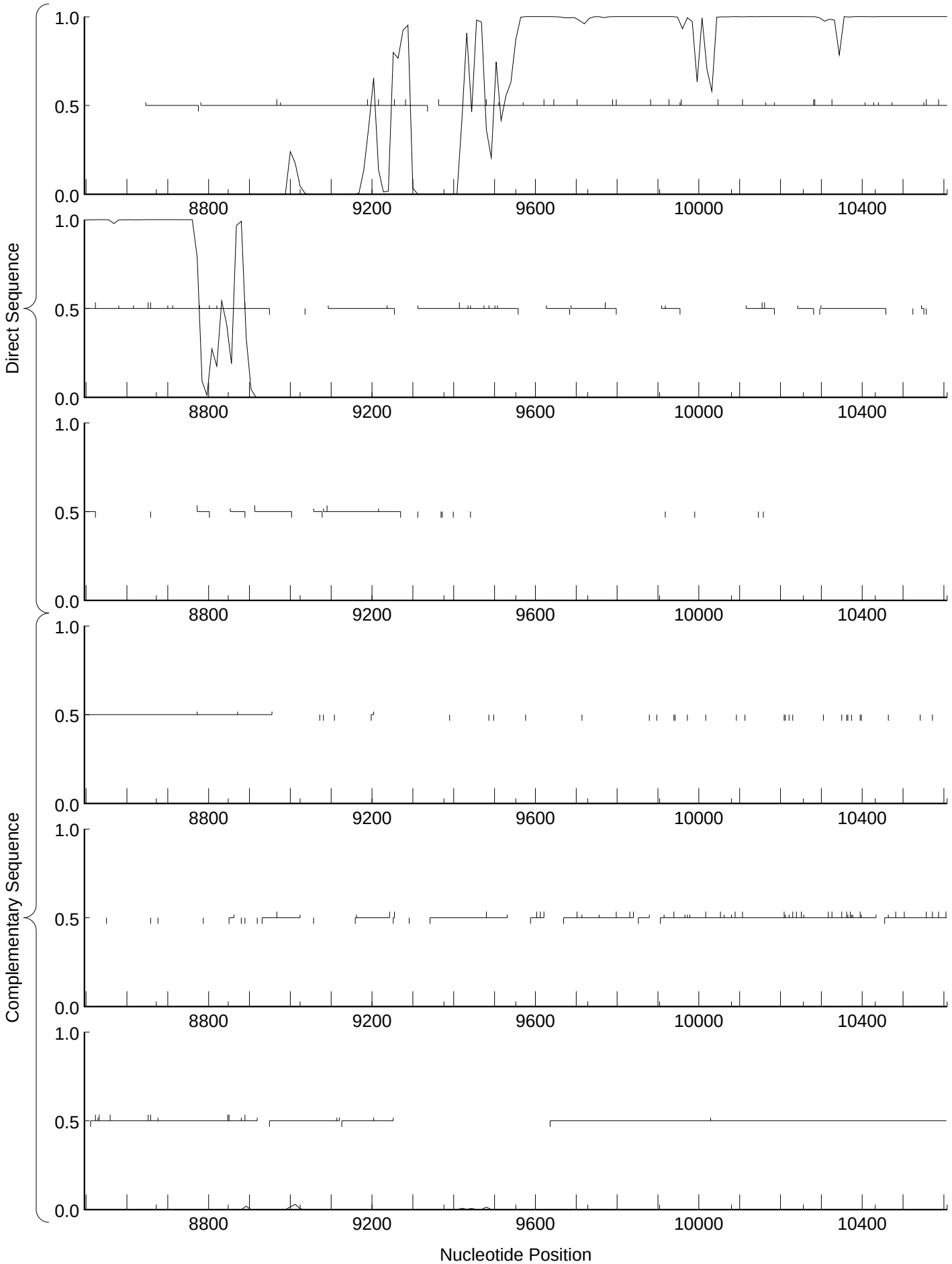



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
4400
4800
5200
5600
6000
Nucleotide Position

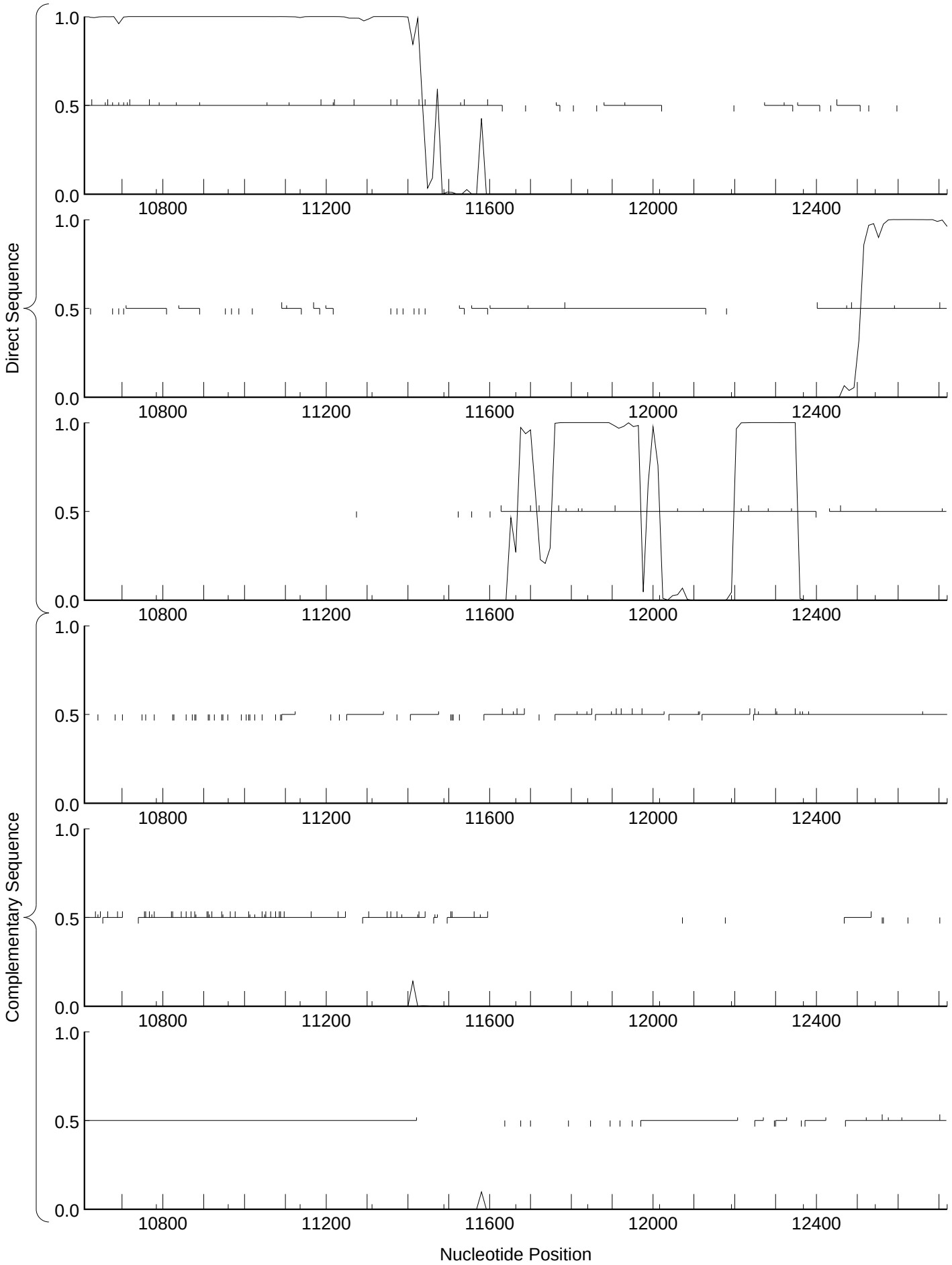
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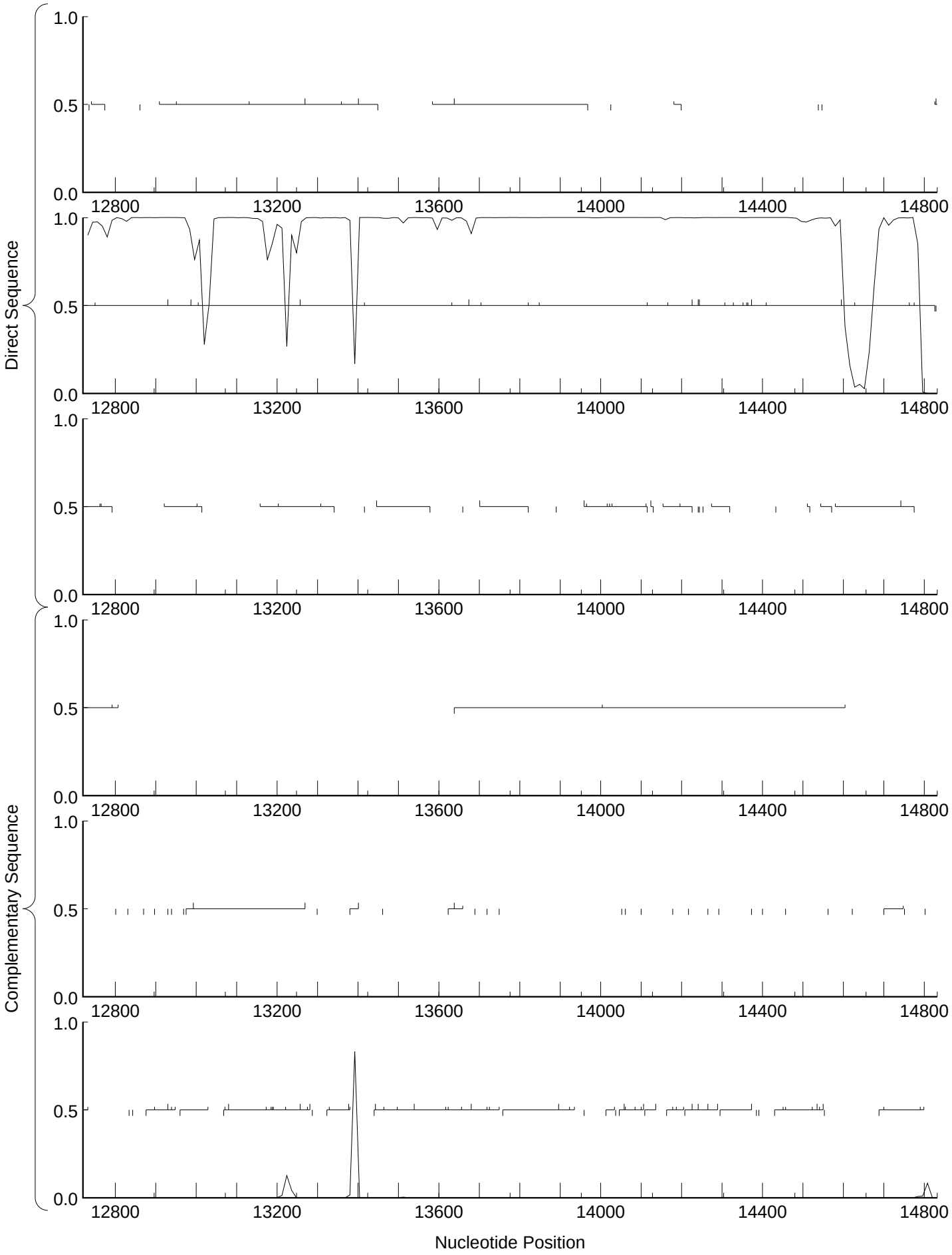
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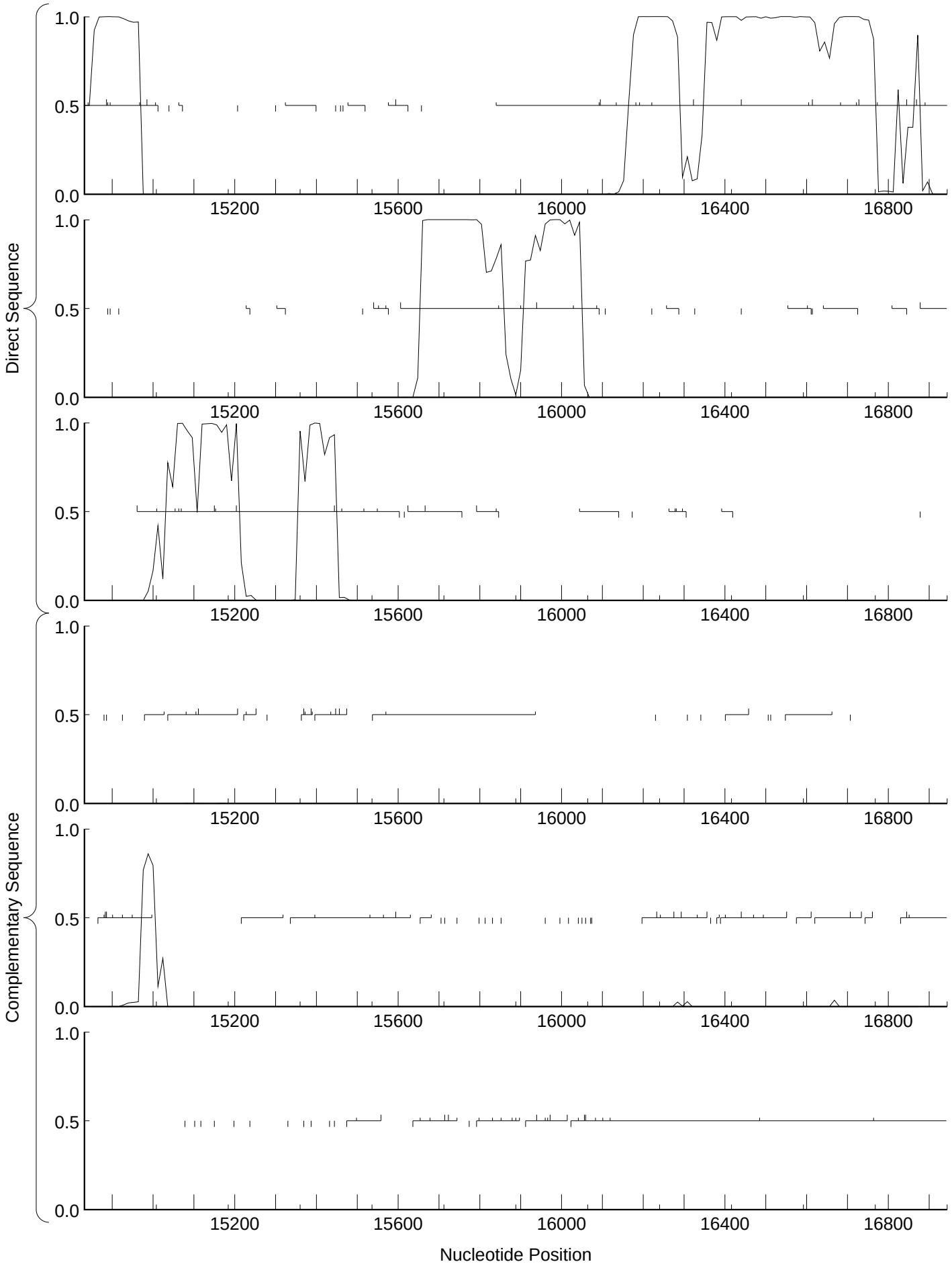
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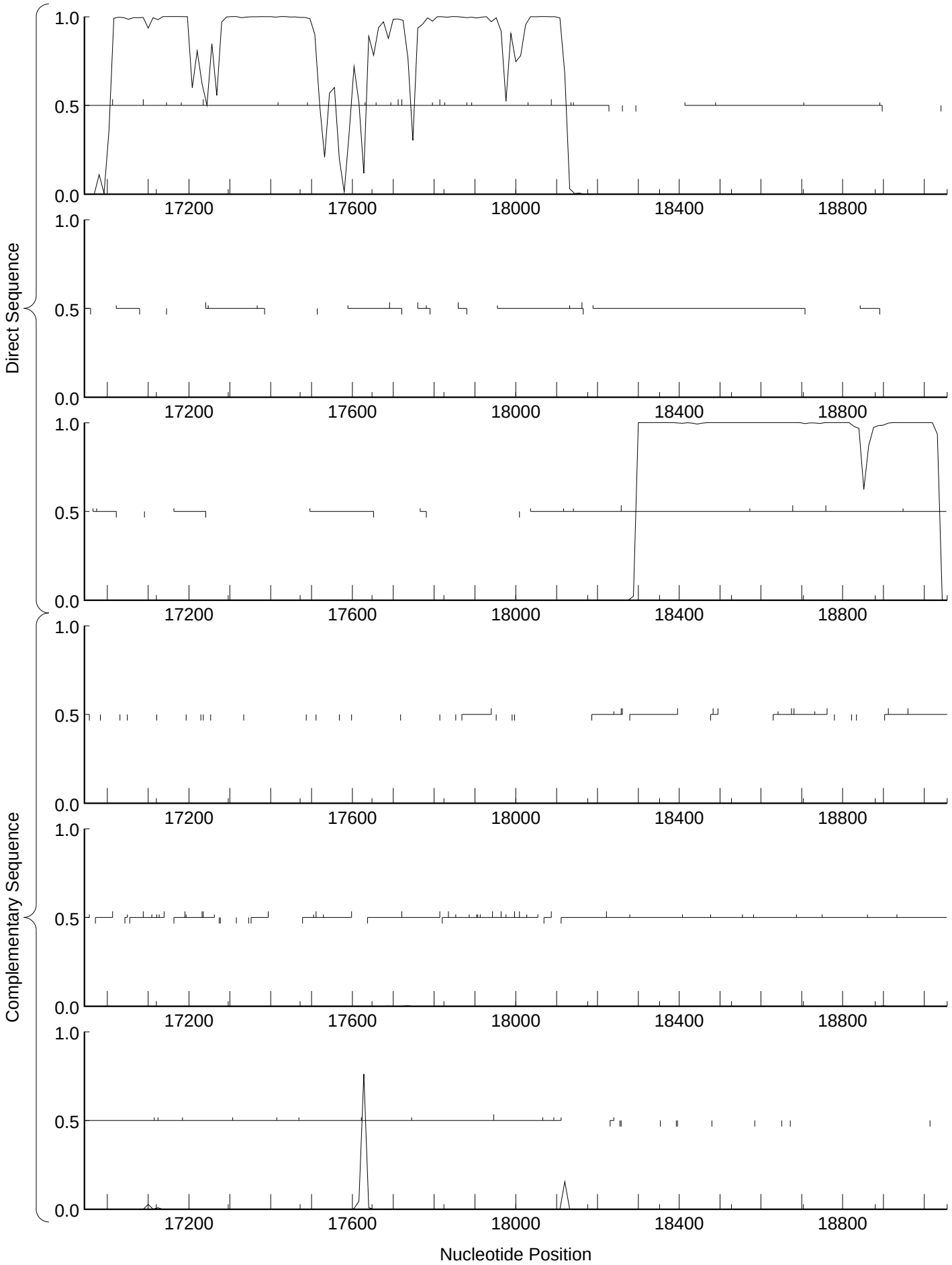
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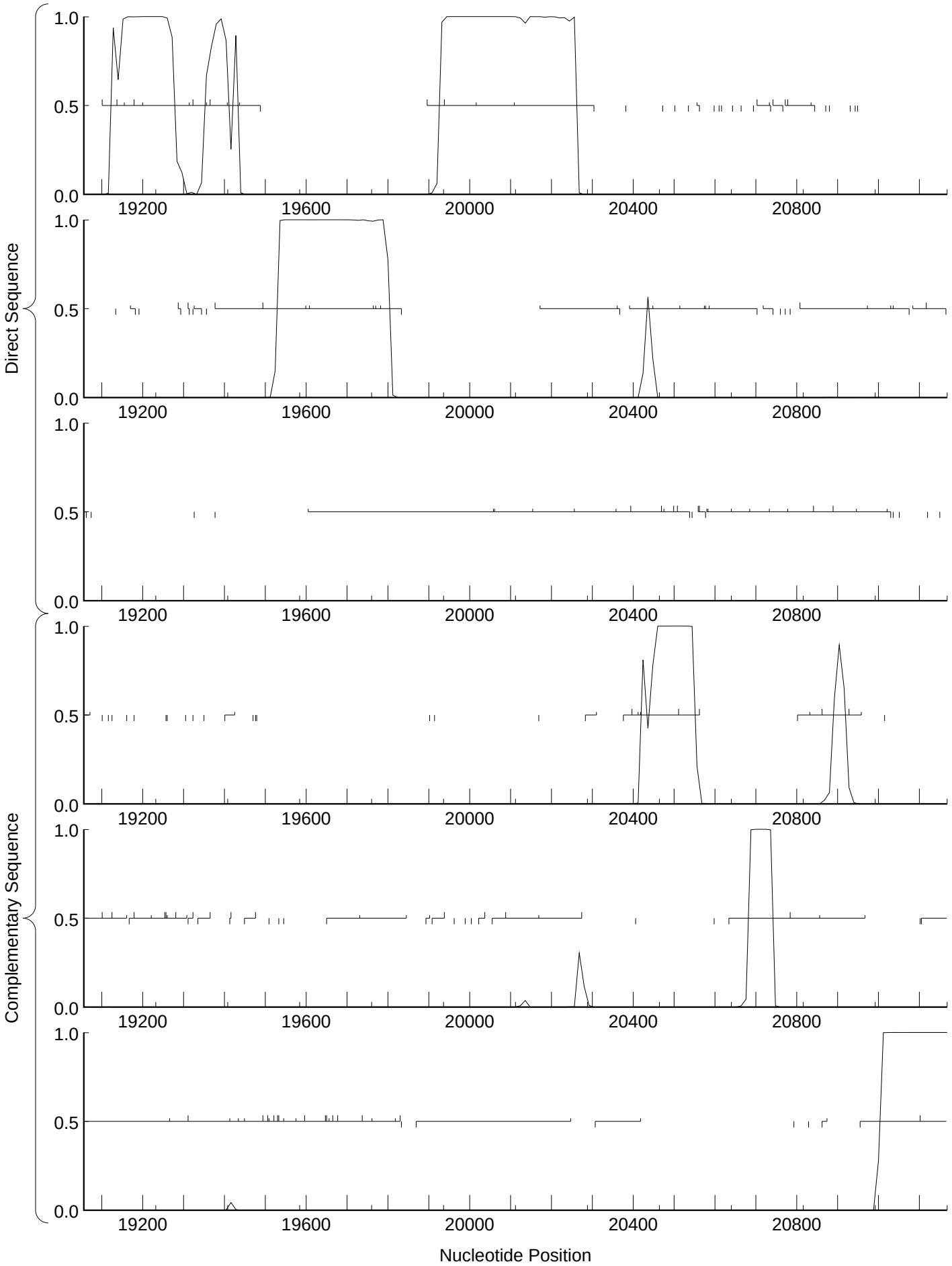
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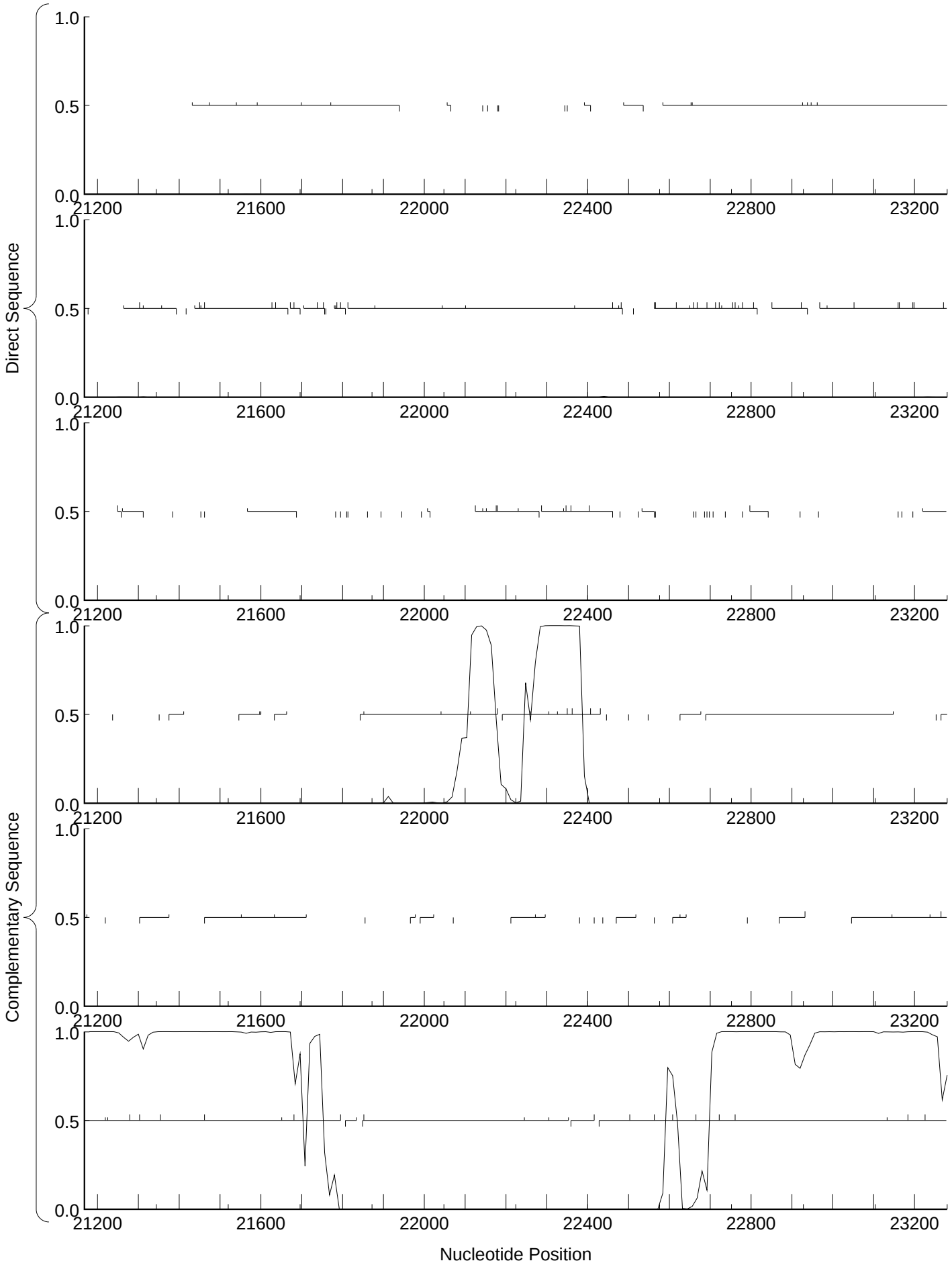
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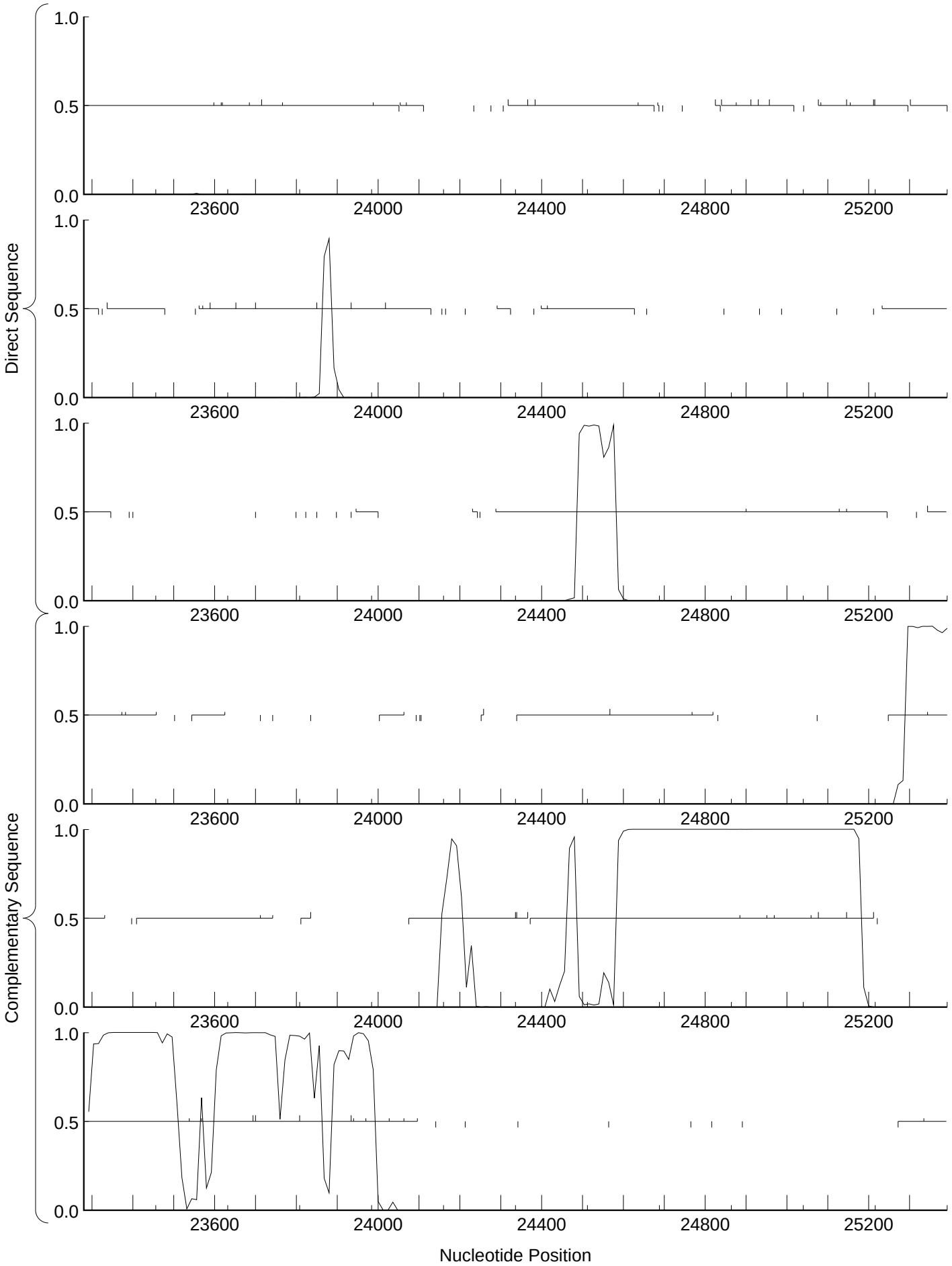
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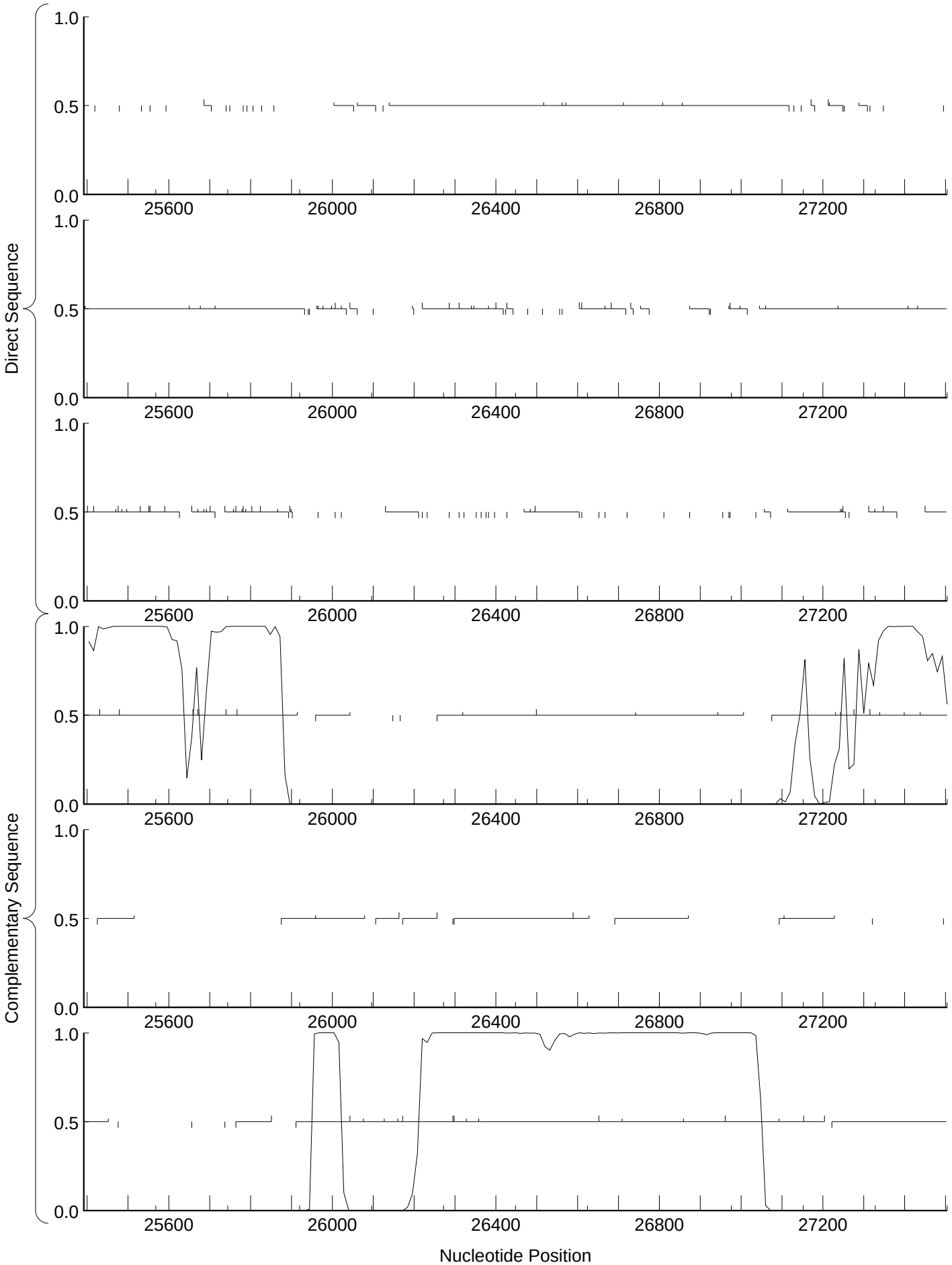
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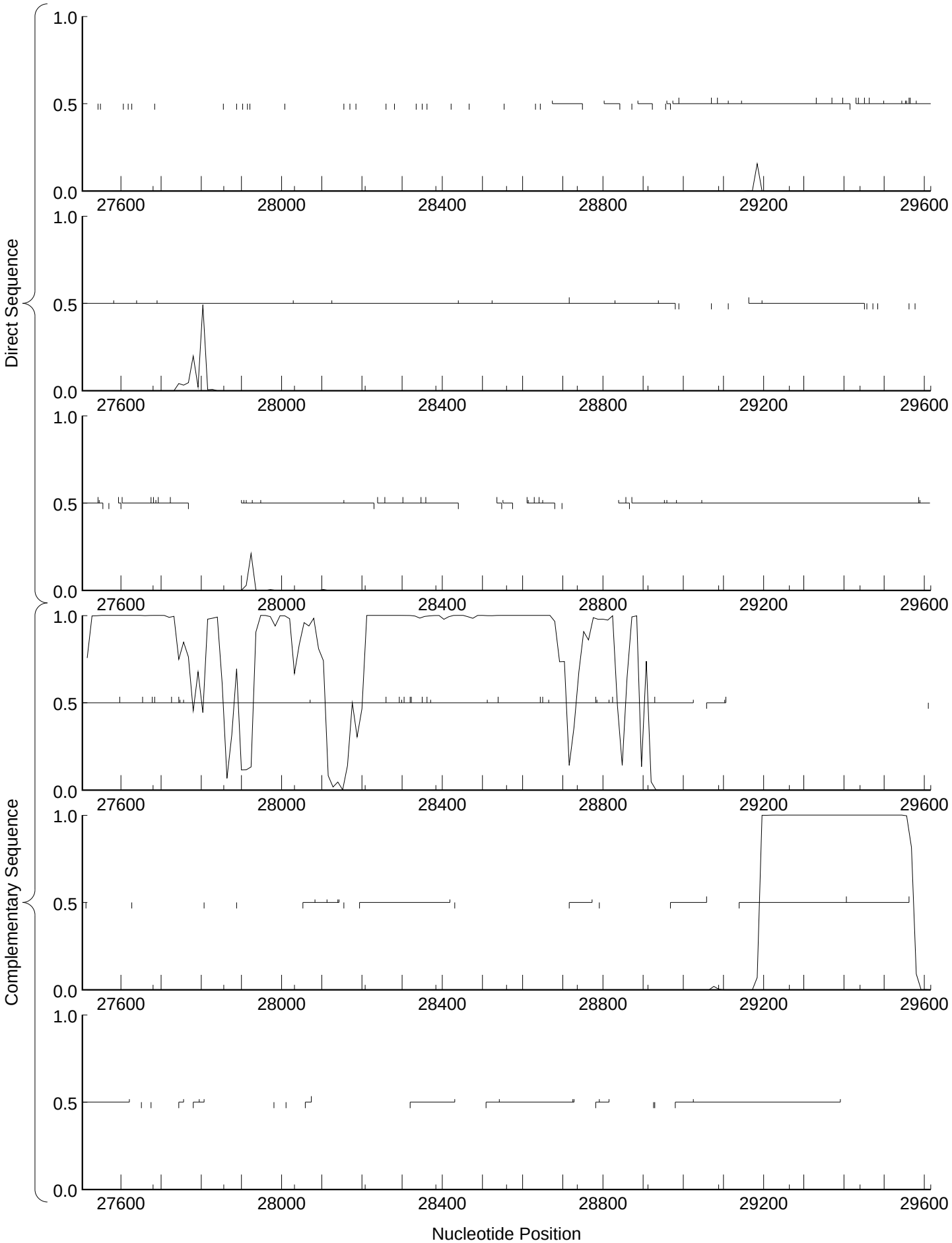
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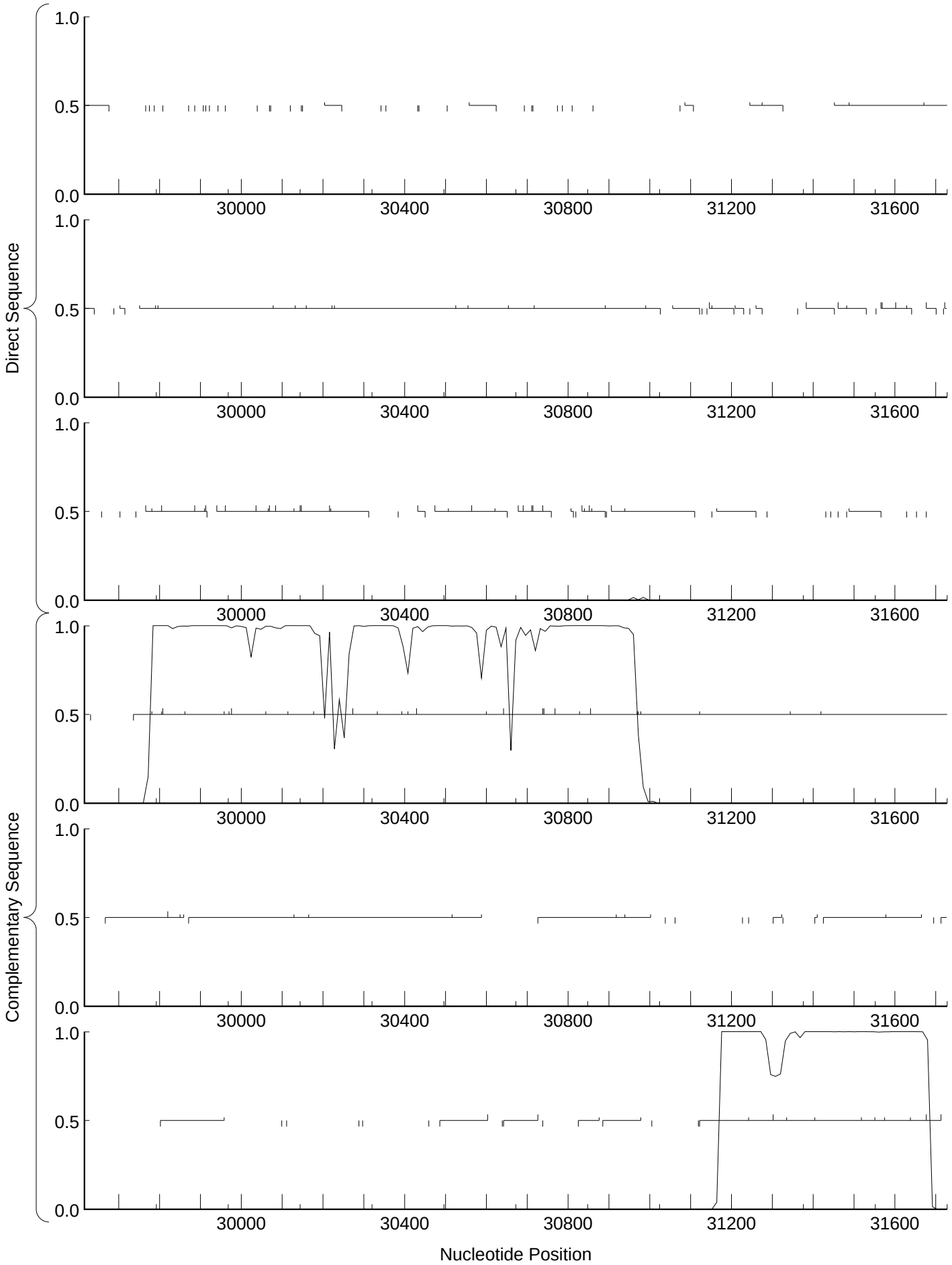
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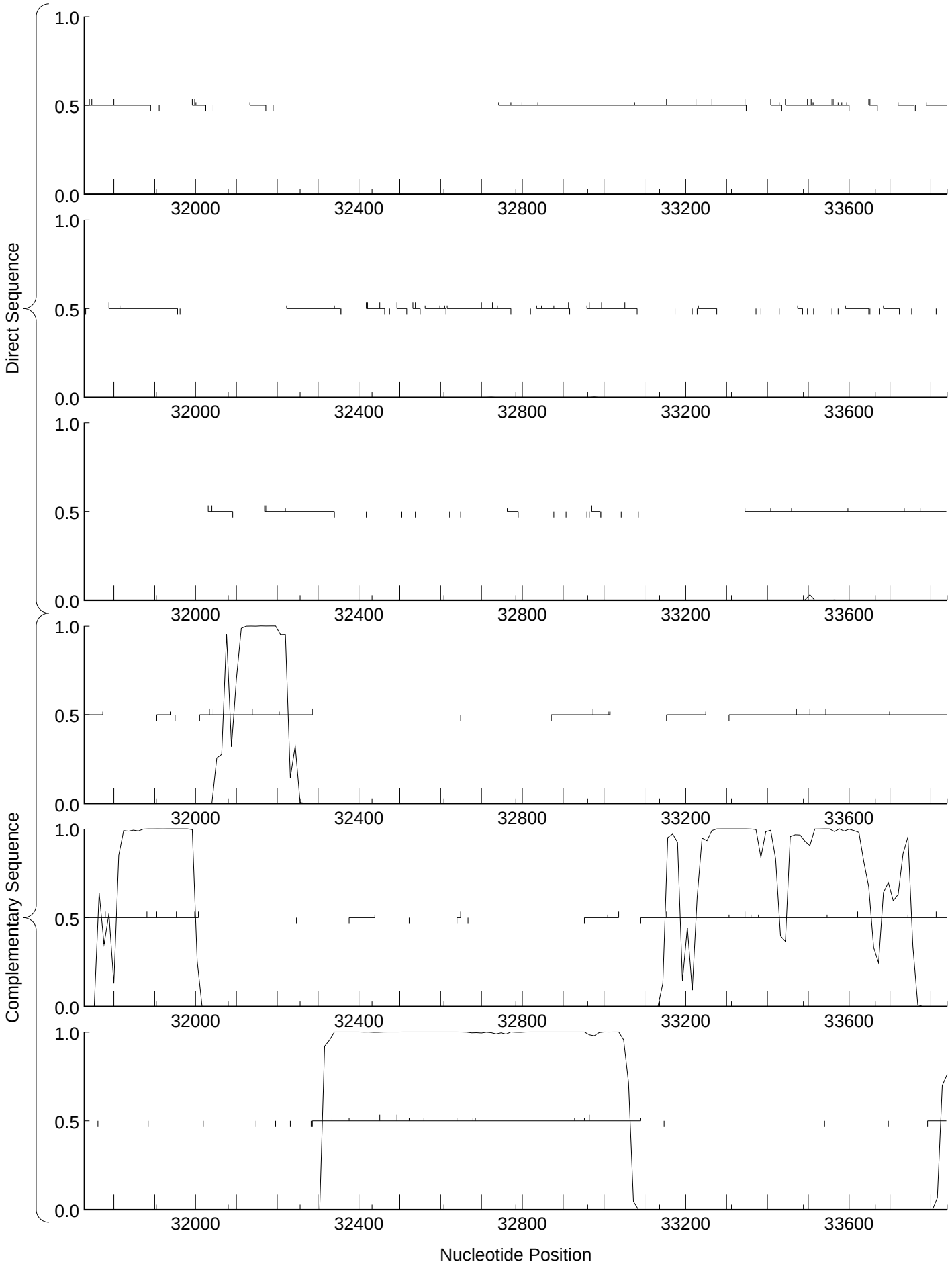
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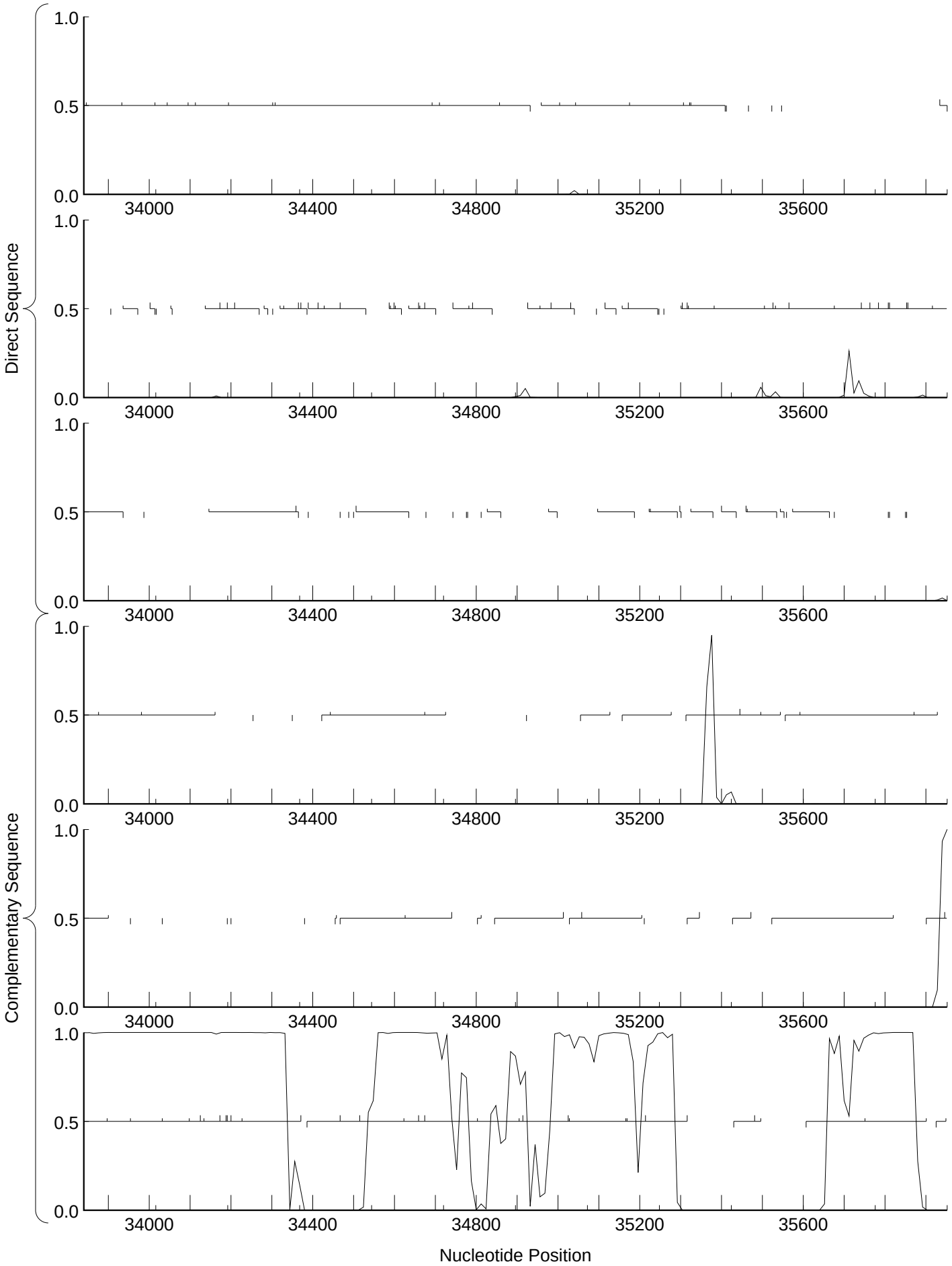
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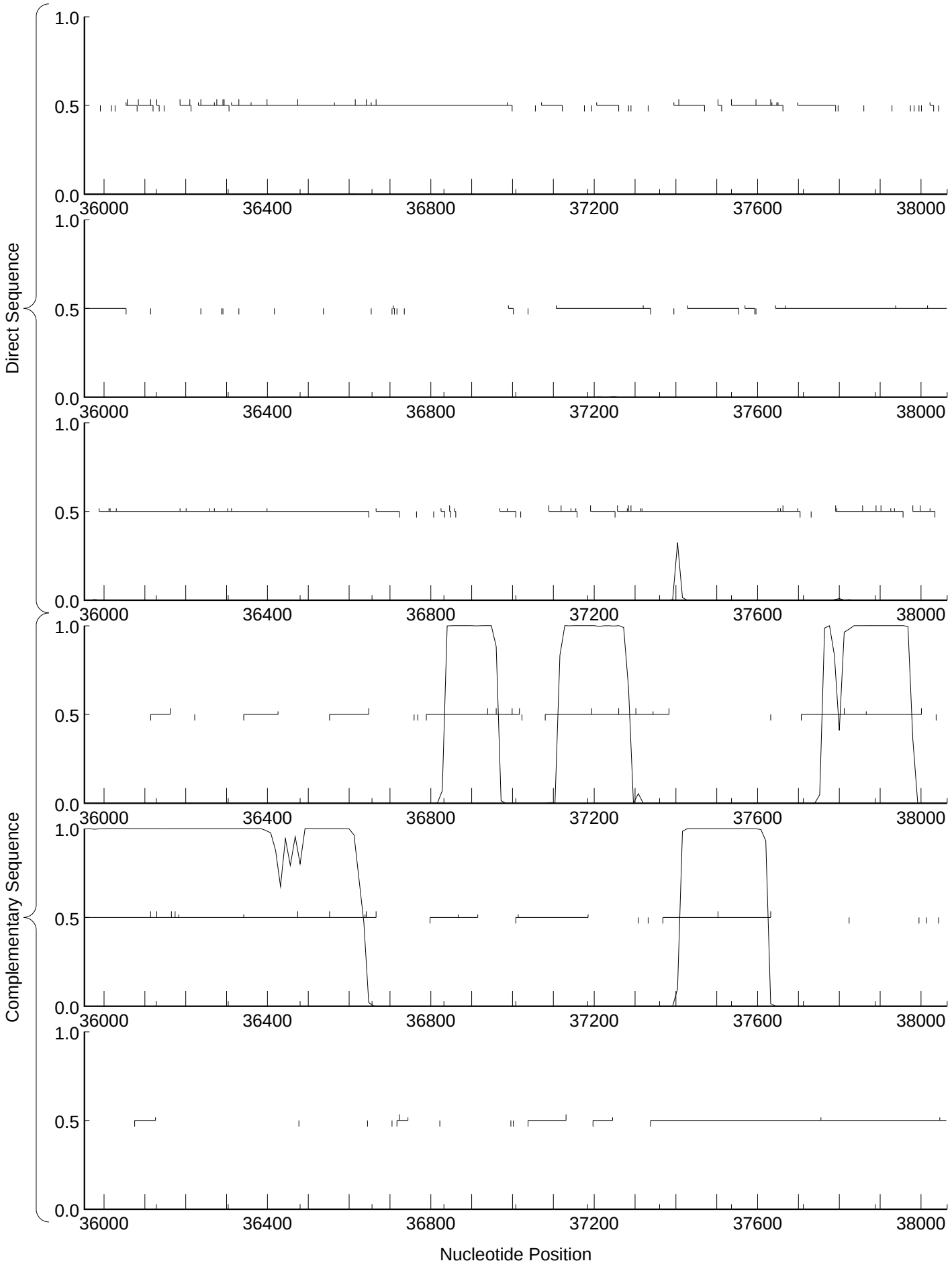
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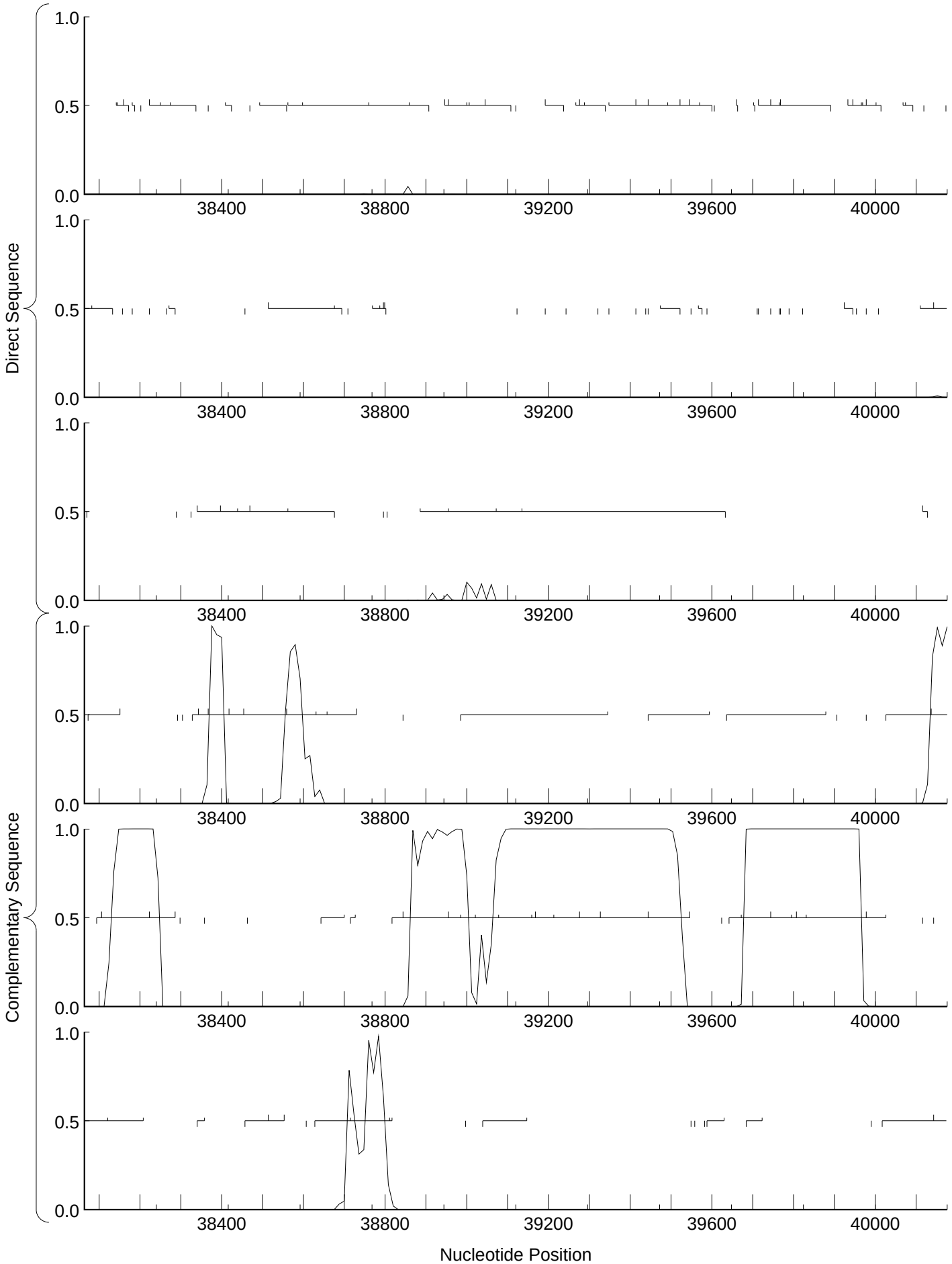
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obacterium phage Gargoyle complete sequence, 41803 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 19/21



obacterium phage Gargoyle complete sequence, 41803 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 20/21



obacterium phage Gargoyle complete sequence, 41803 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 21/21

