GeneMark

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

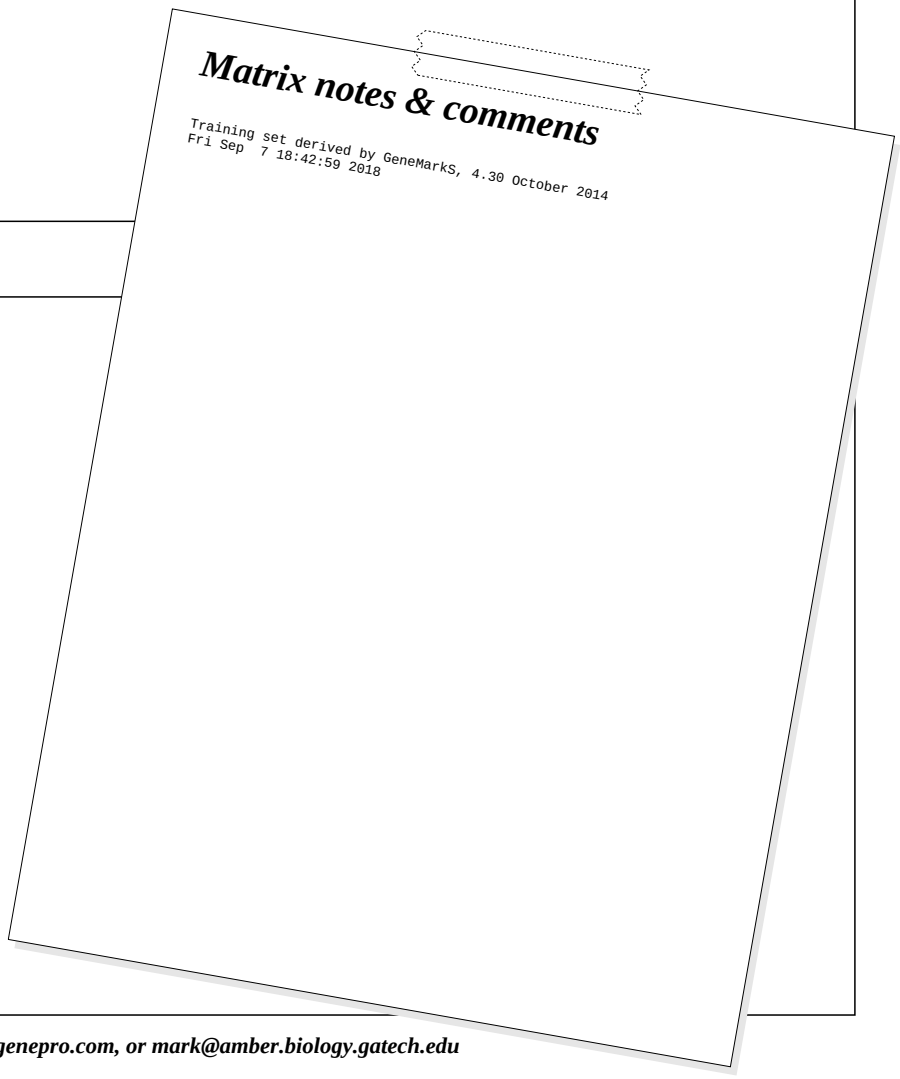
PROGRAM INFORMATION

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Sequence         :  Microbacterium phage Espinosa complete sequence, 41553 bp, circularly permuted, Cluster EA1
Analysis Date    :  3/29/24 at 2:36:38
Pages            :  21
Sequence Length  :  41553 bp
GC Content       :   63.41%

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

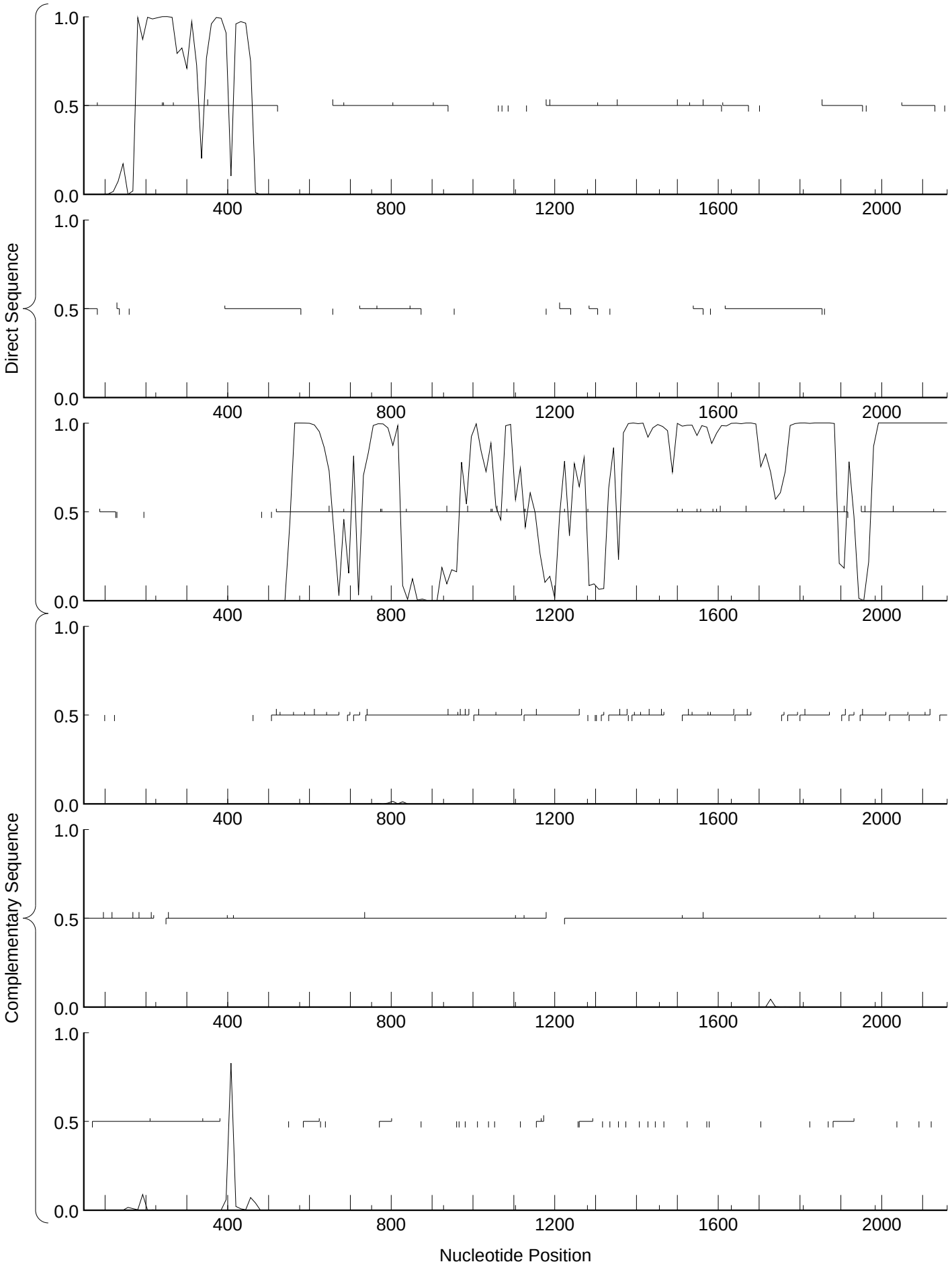
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Author     :  -

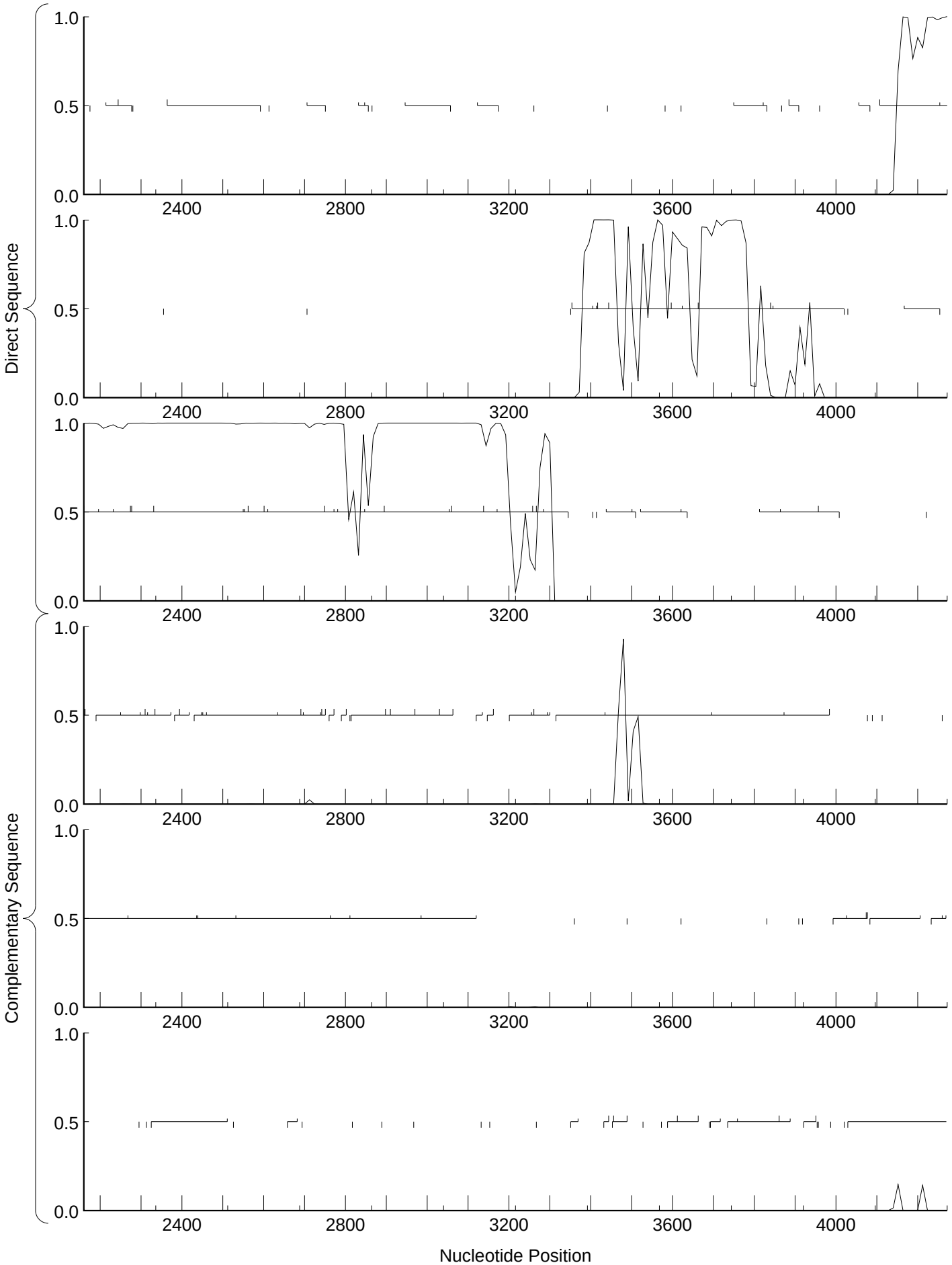
Order      :  4
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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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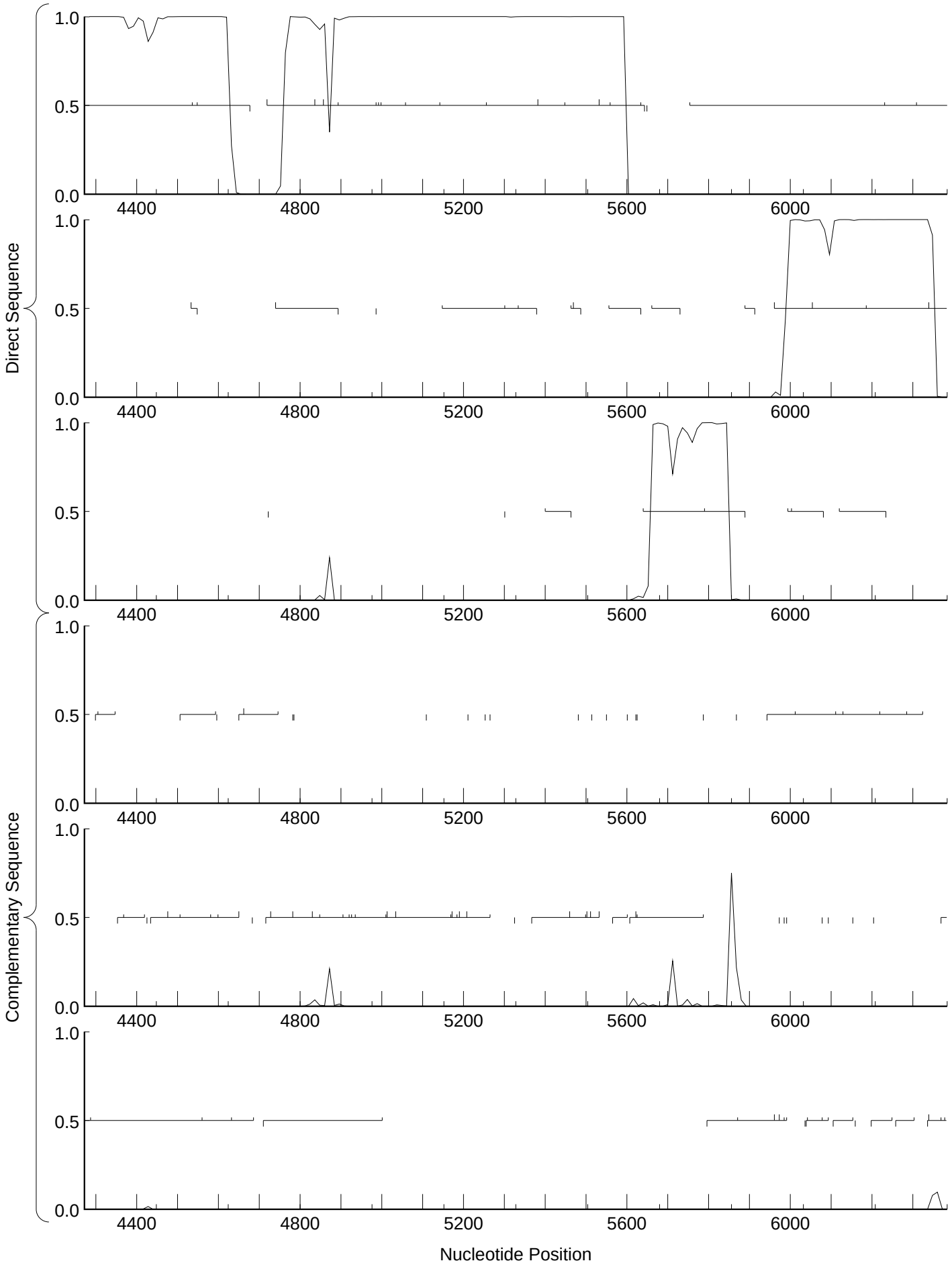

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

robacterium phage Espinosa complete sequence, 41553 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 2/21

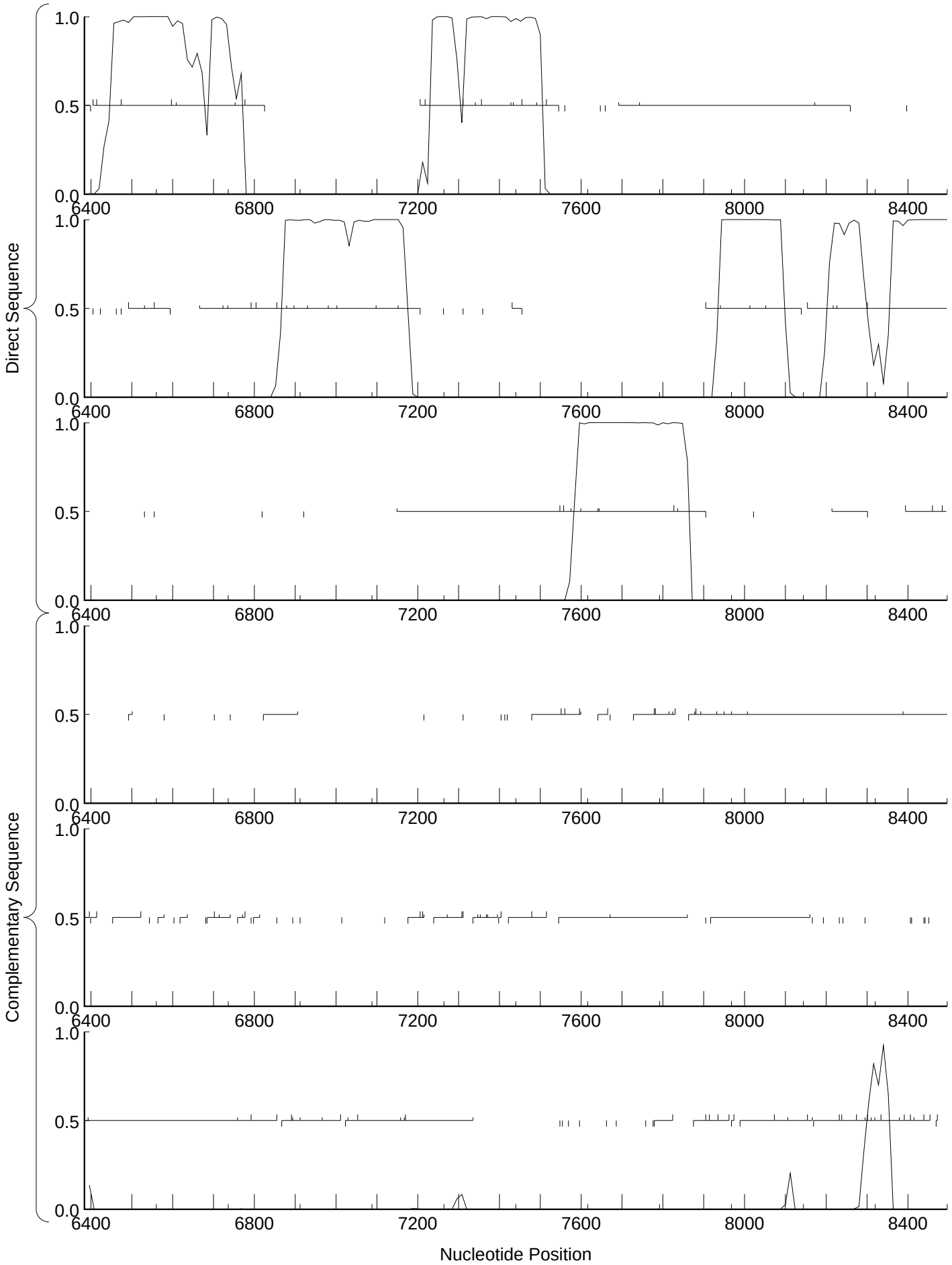



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
2400
2800
3200
3600
4000
Nucleotide Position

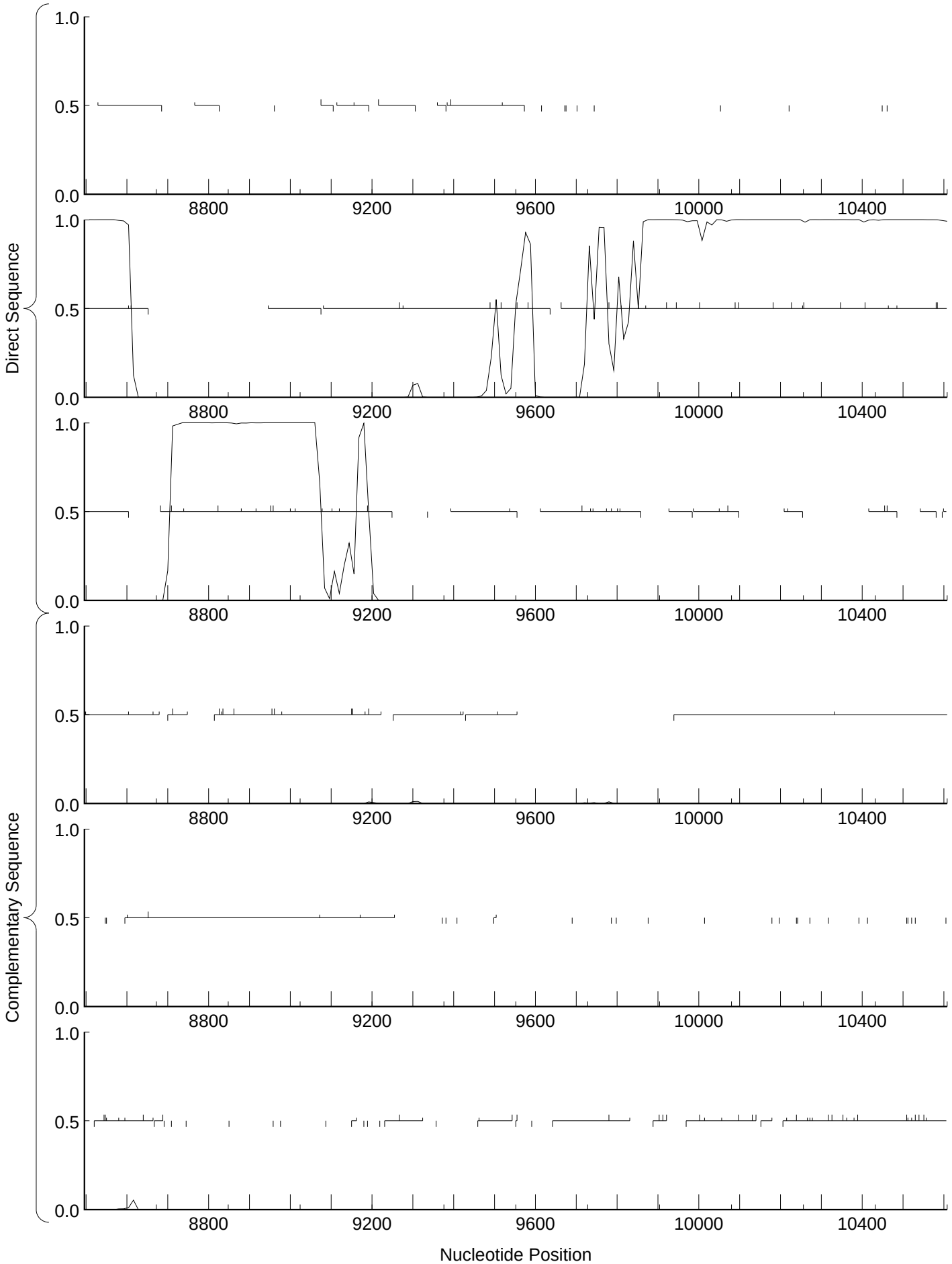
robacterium phage Espinosa complete sequence, 41553 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 4/21



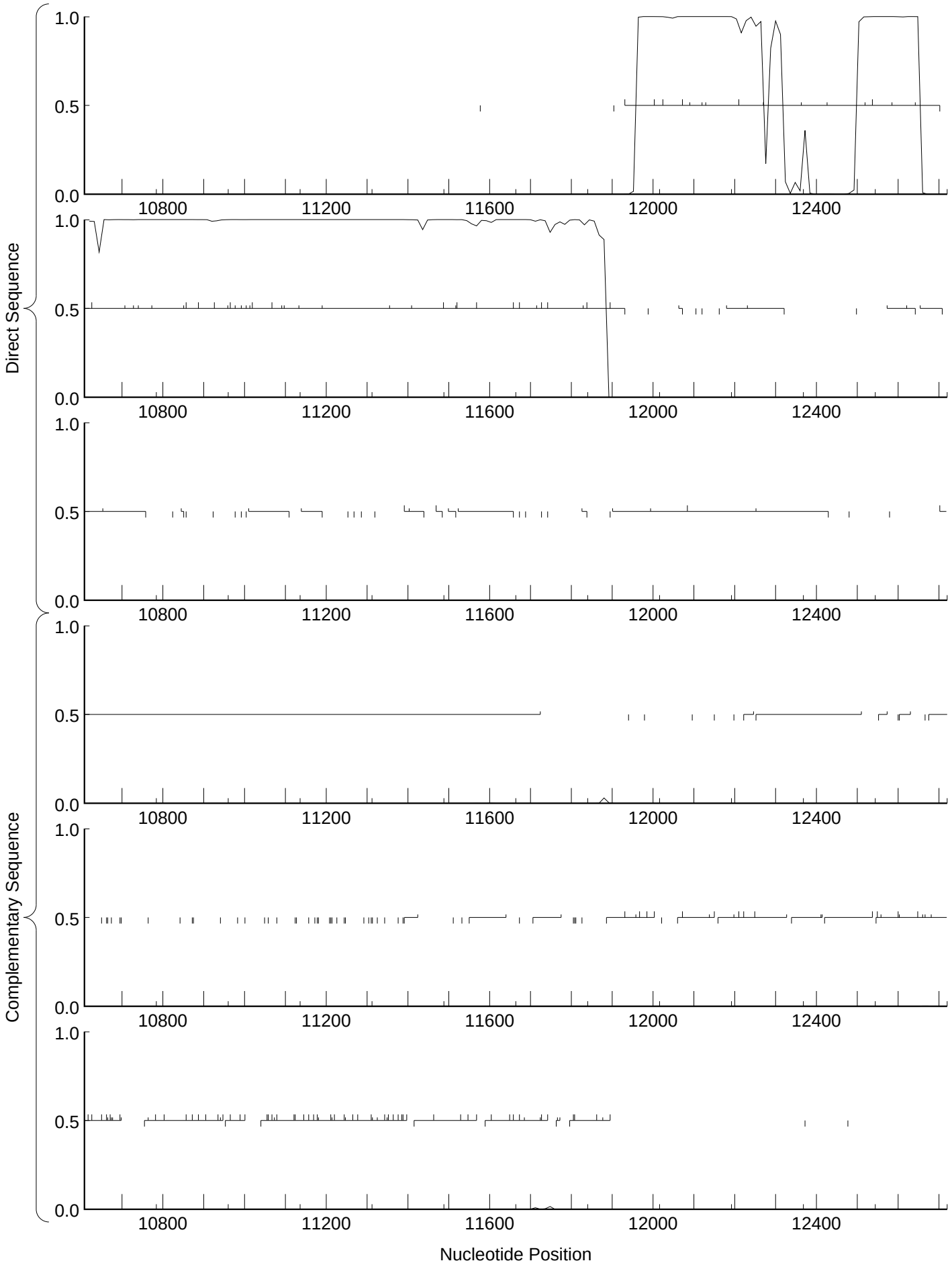
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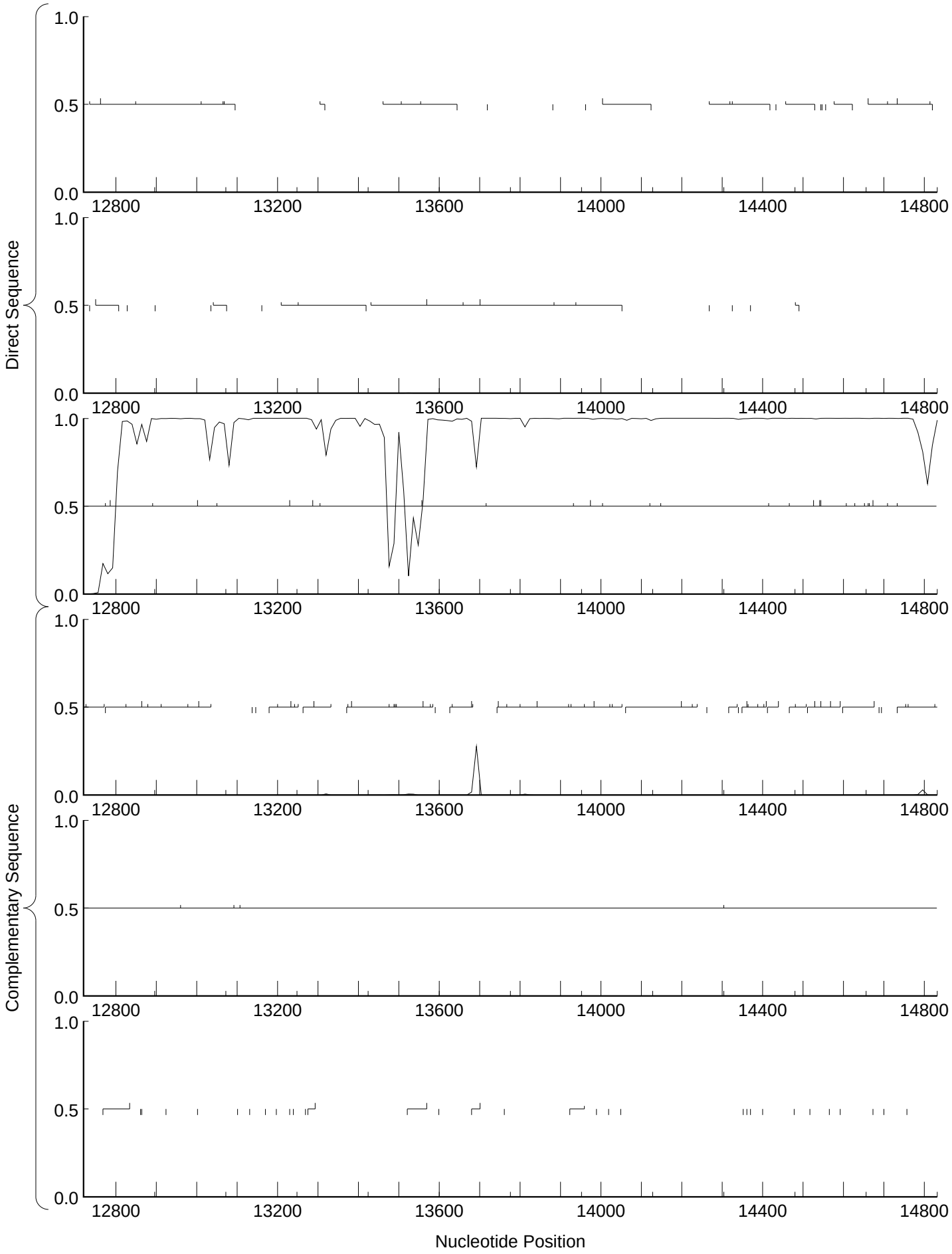
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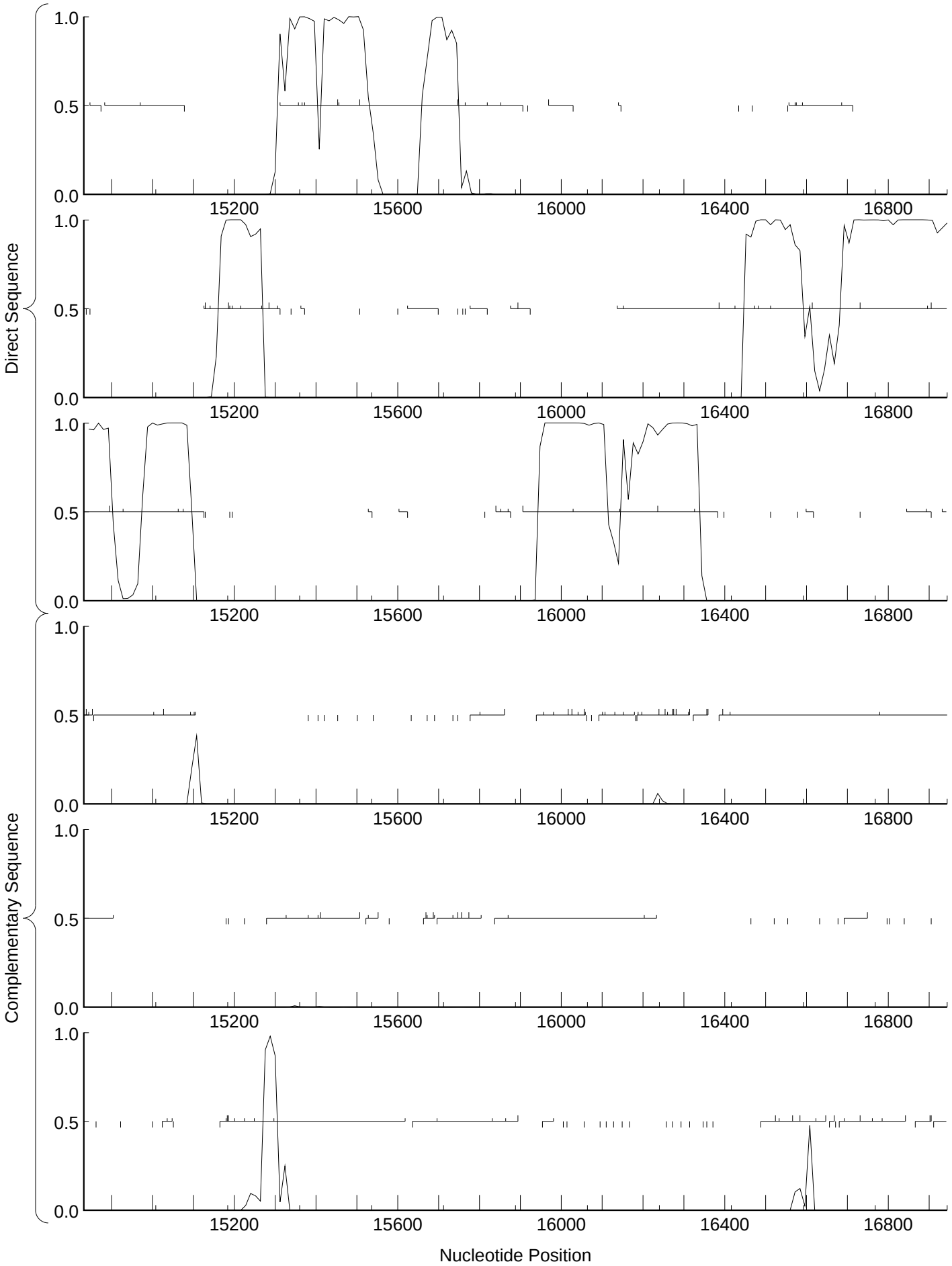
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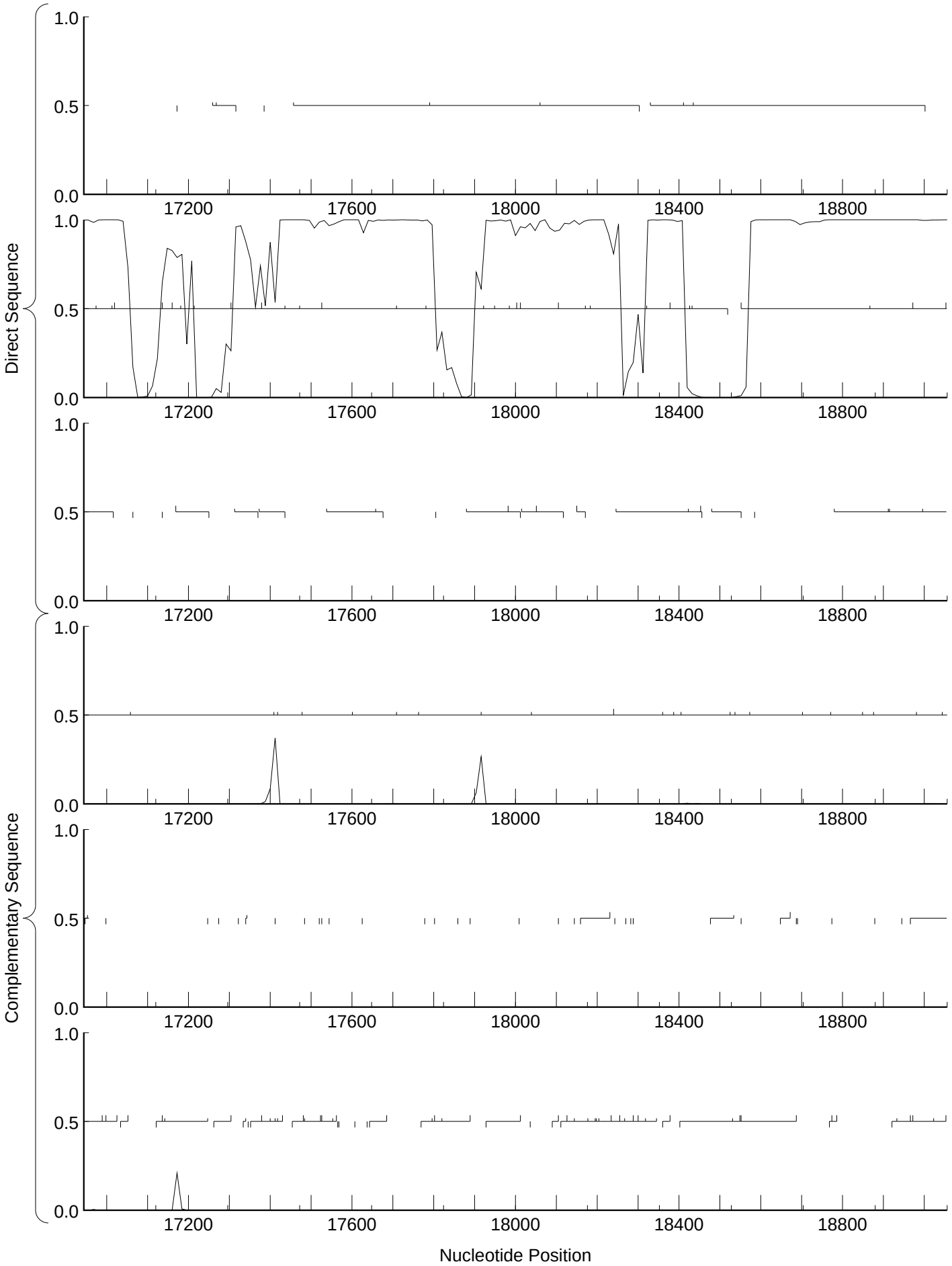
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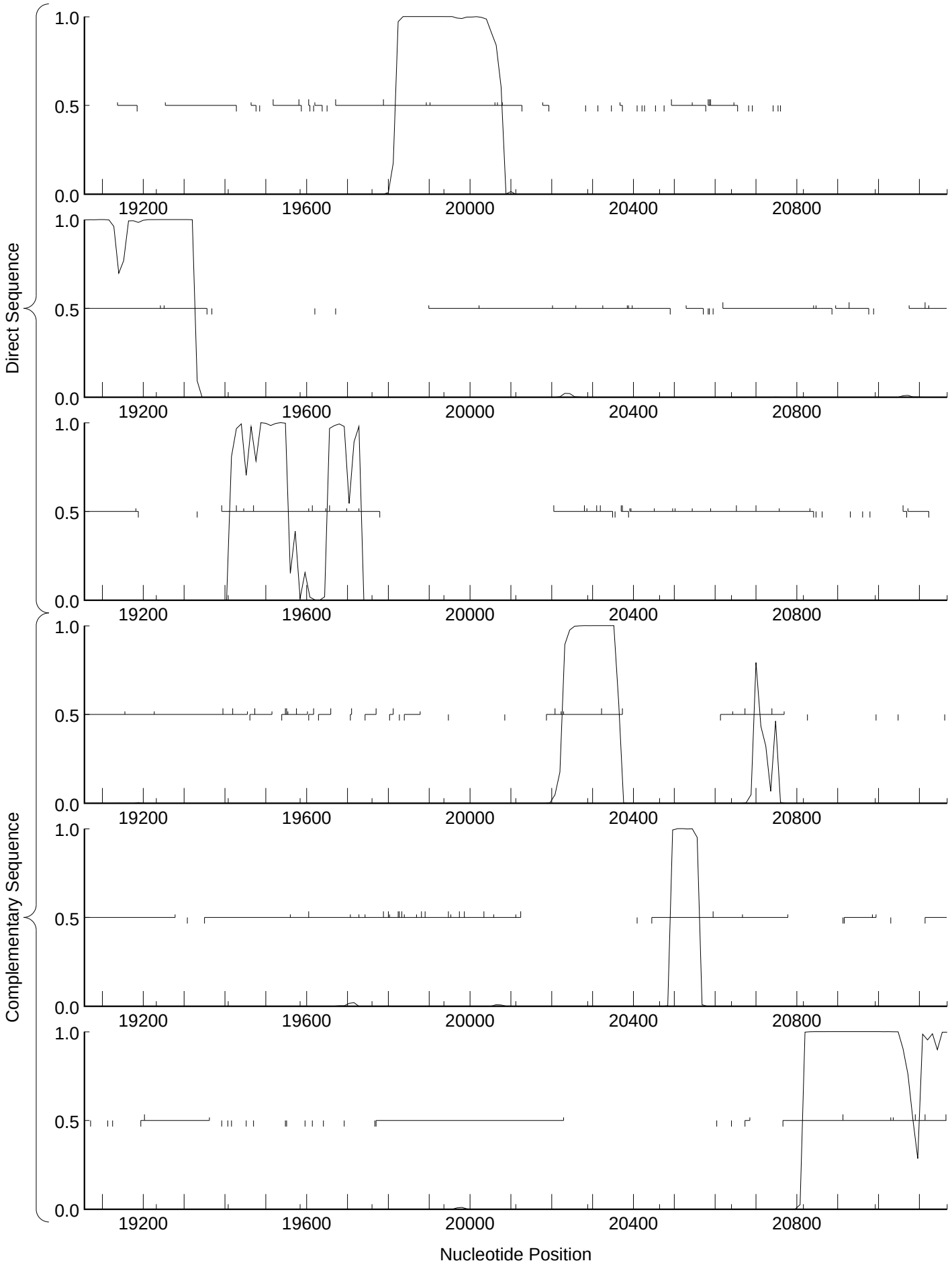
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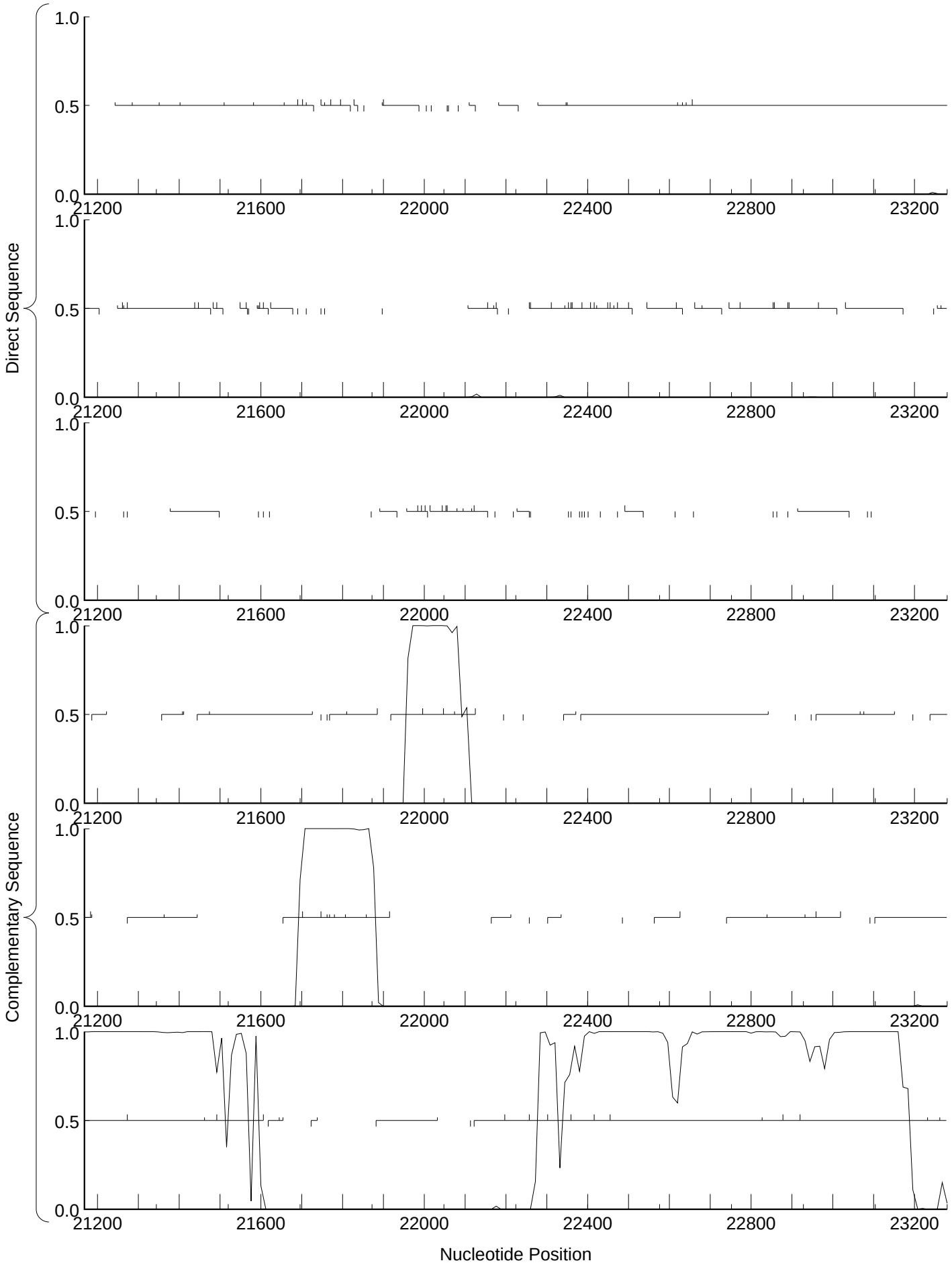
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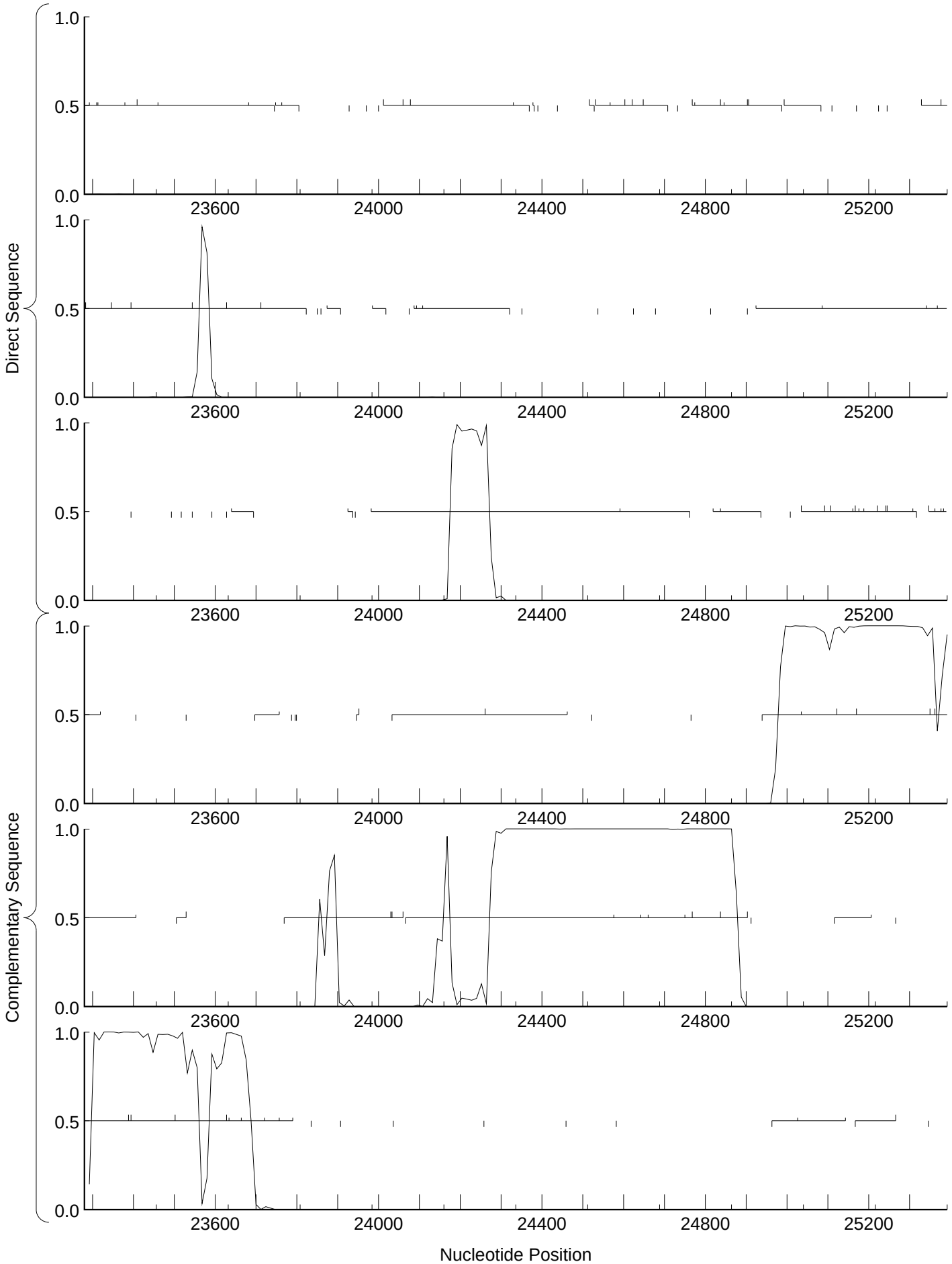
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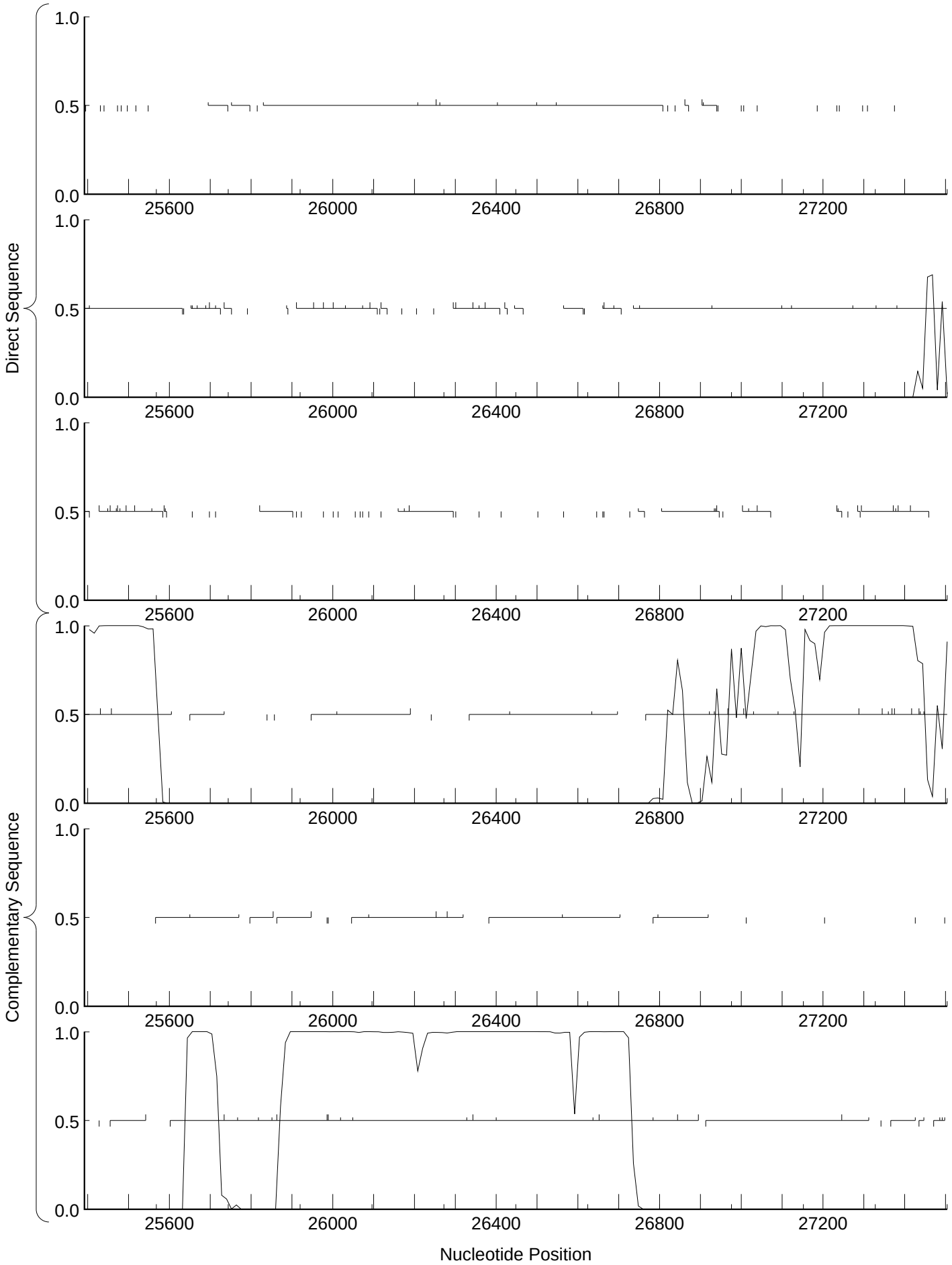
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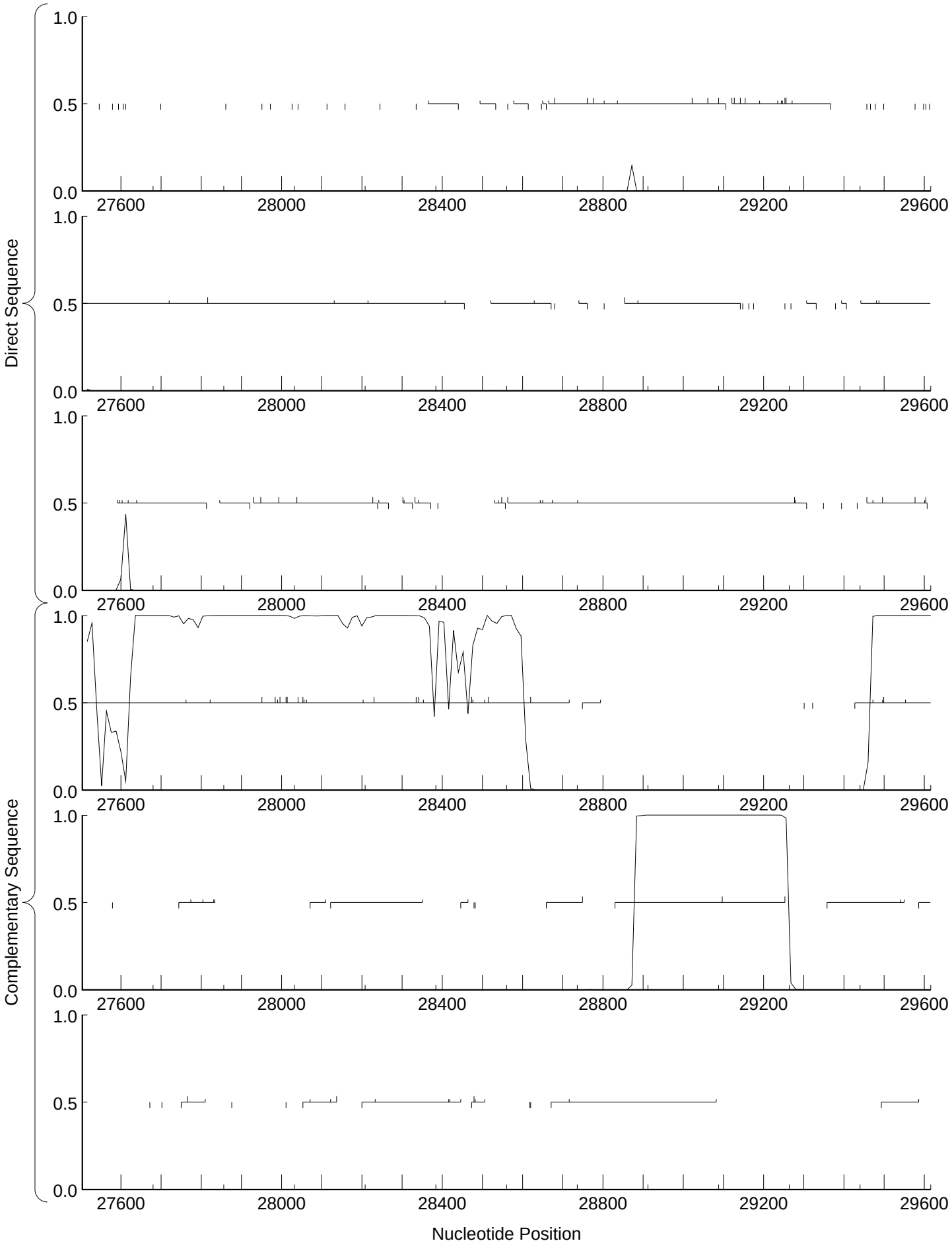
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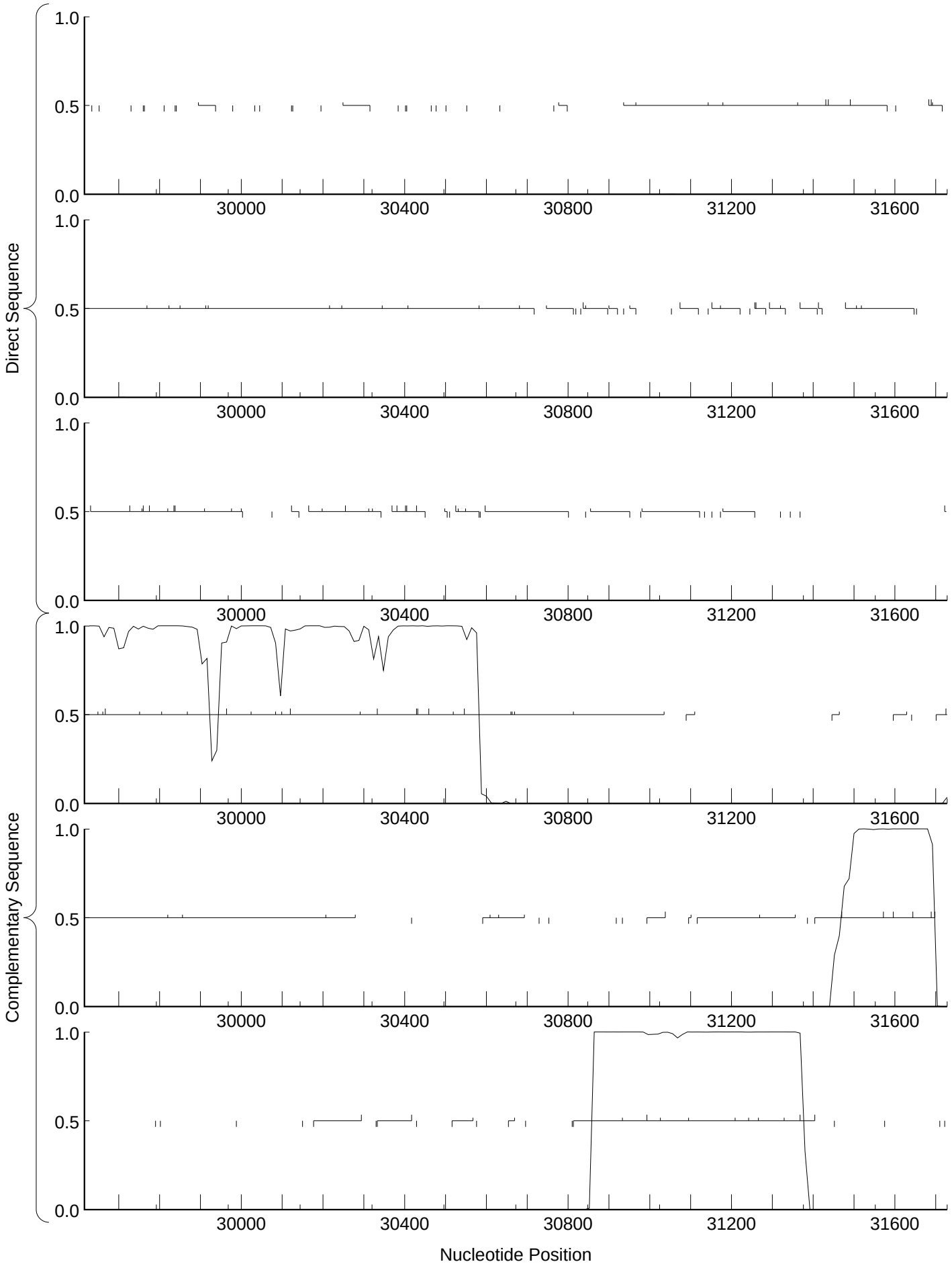
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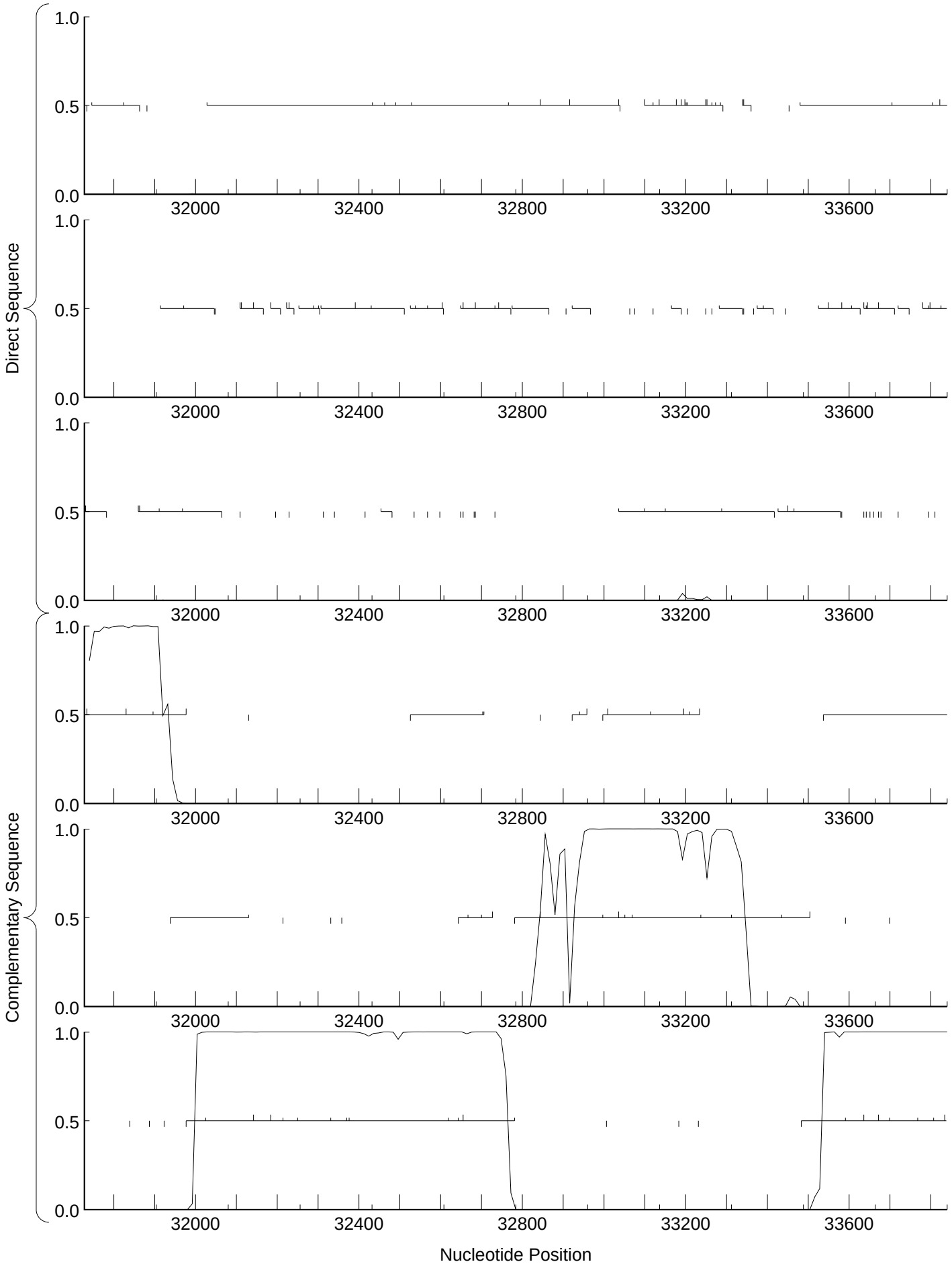
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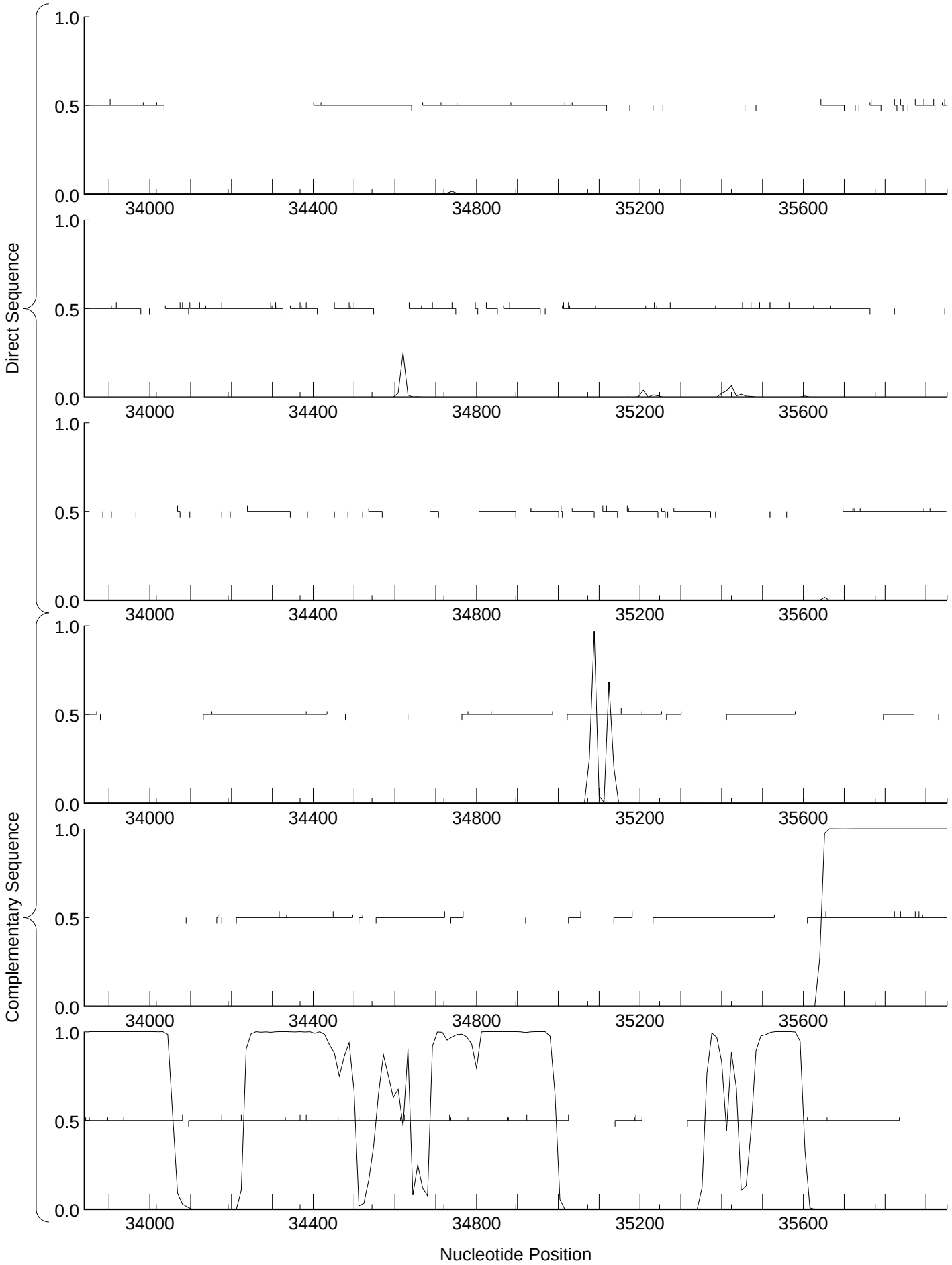
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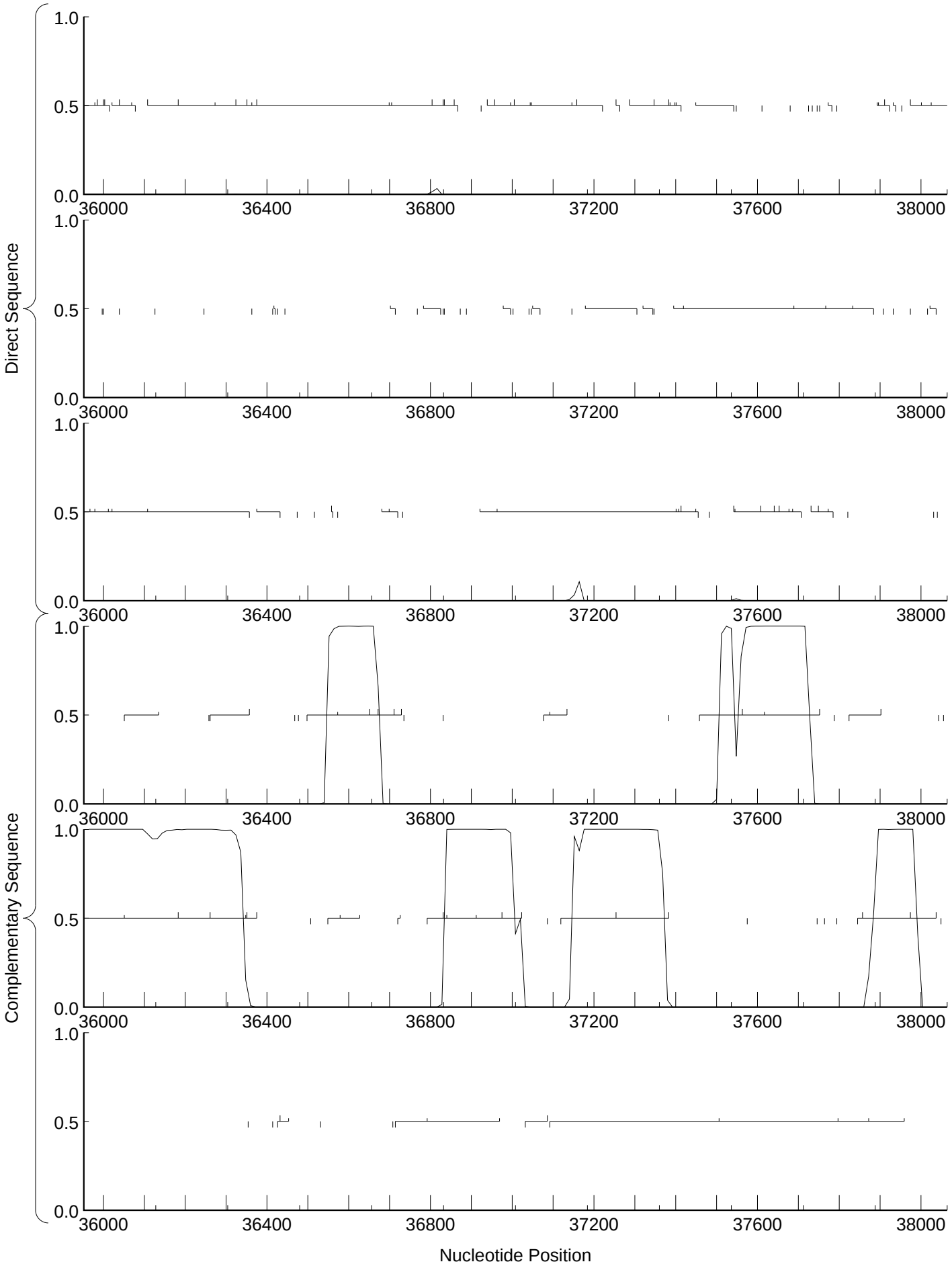
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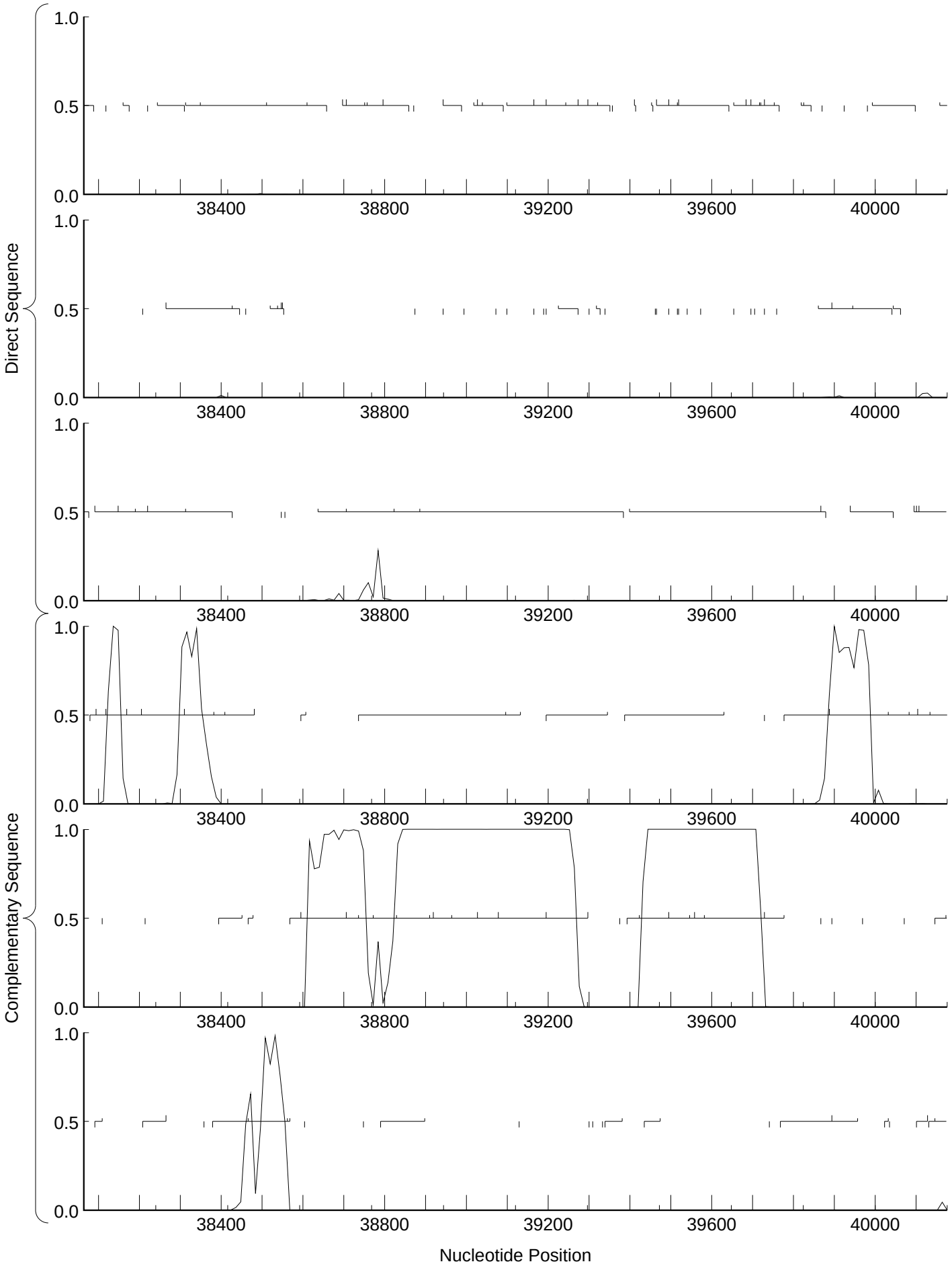
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obacterium phage Espinosa complete sequence, 41553 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 19/21



obacterium phage Espinosa complete sequence, 41553 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 20/21



obacterium phage Espinosa complete sequence, 41553 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 21/21

