GeneMark

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

PROGRAM INFORMATION

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Sequence         :  Mycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1
Analysis Date    :  3/28/24 at 4:56:49
Pages            :  75
Sequence Length  :  155582 bp
GC Content       :   64.71%

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

PS-Version       :  1.2

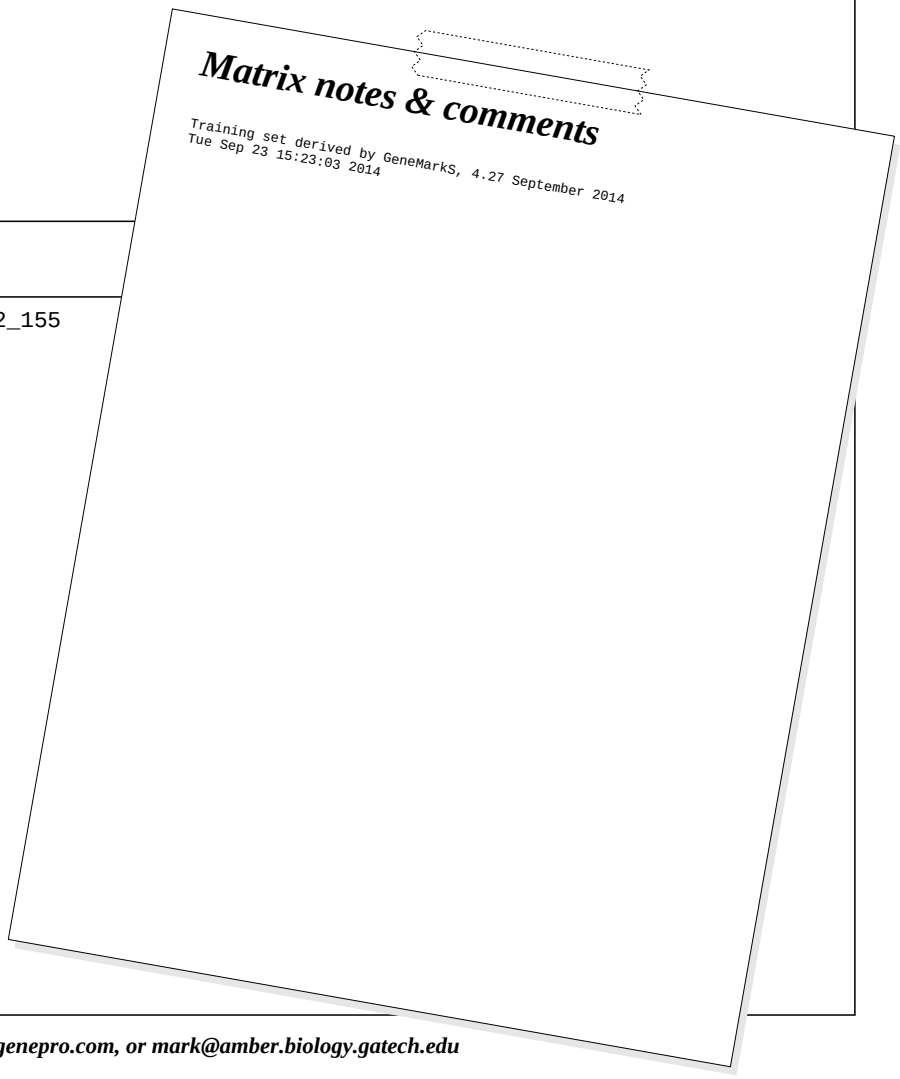
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MATRIX INFORMATION

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Author     :  -

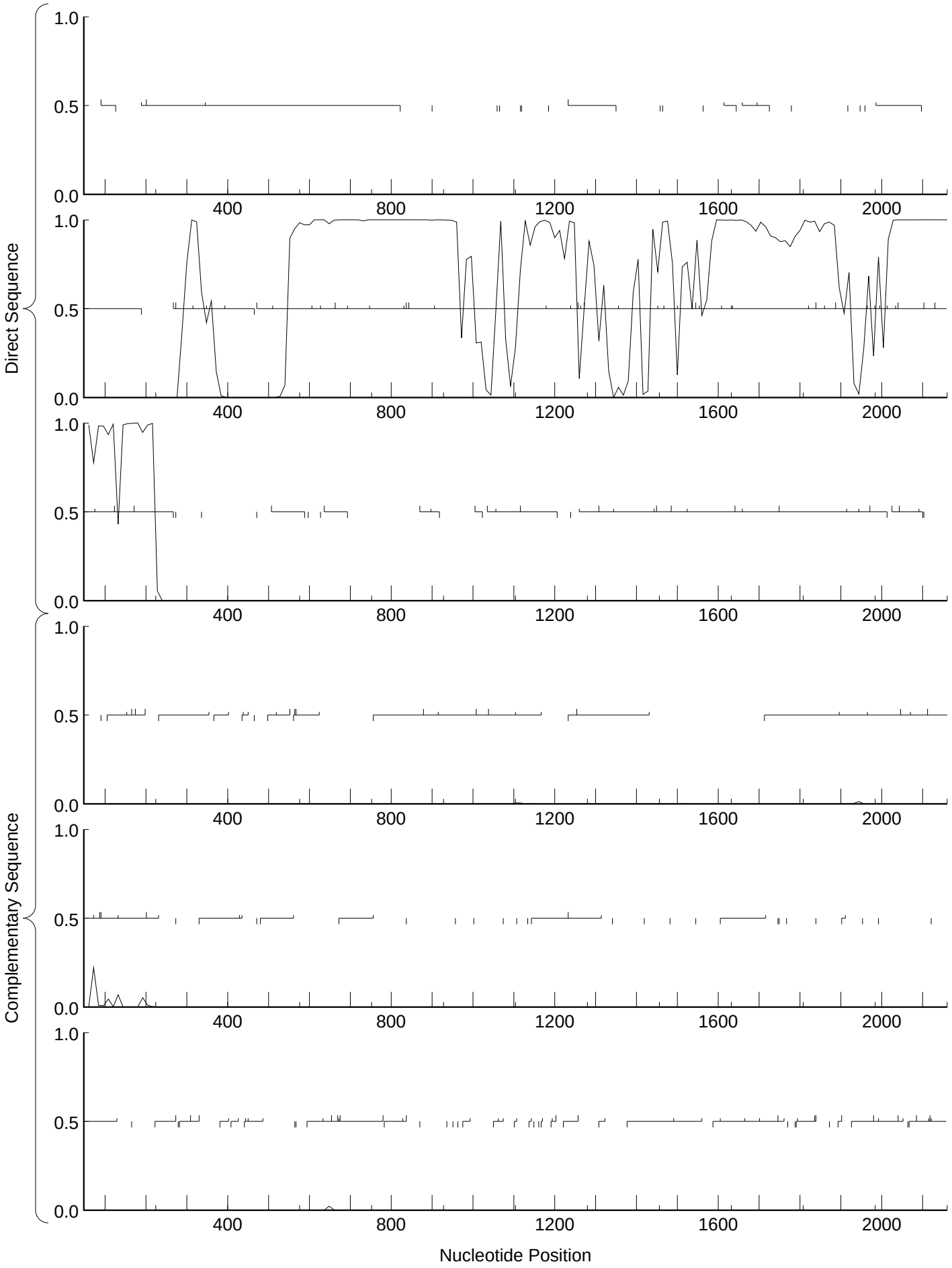
Order      :  4


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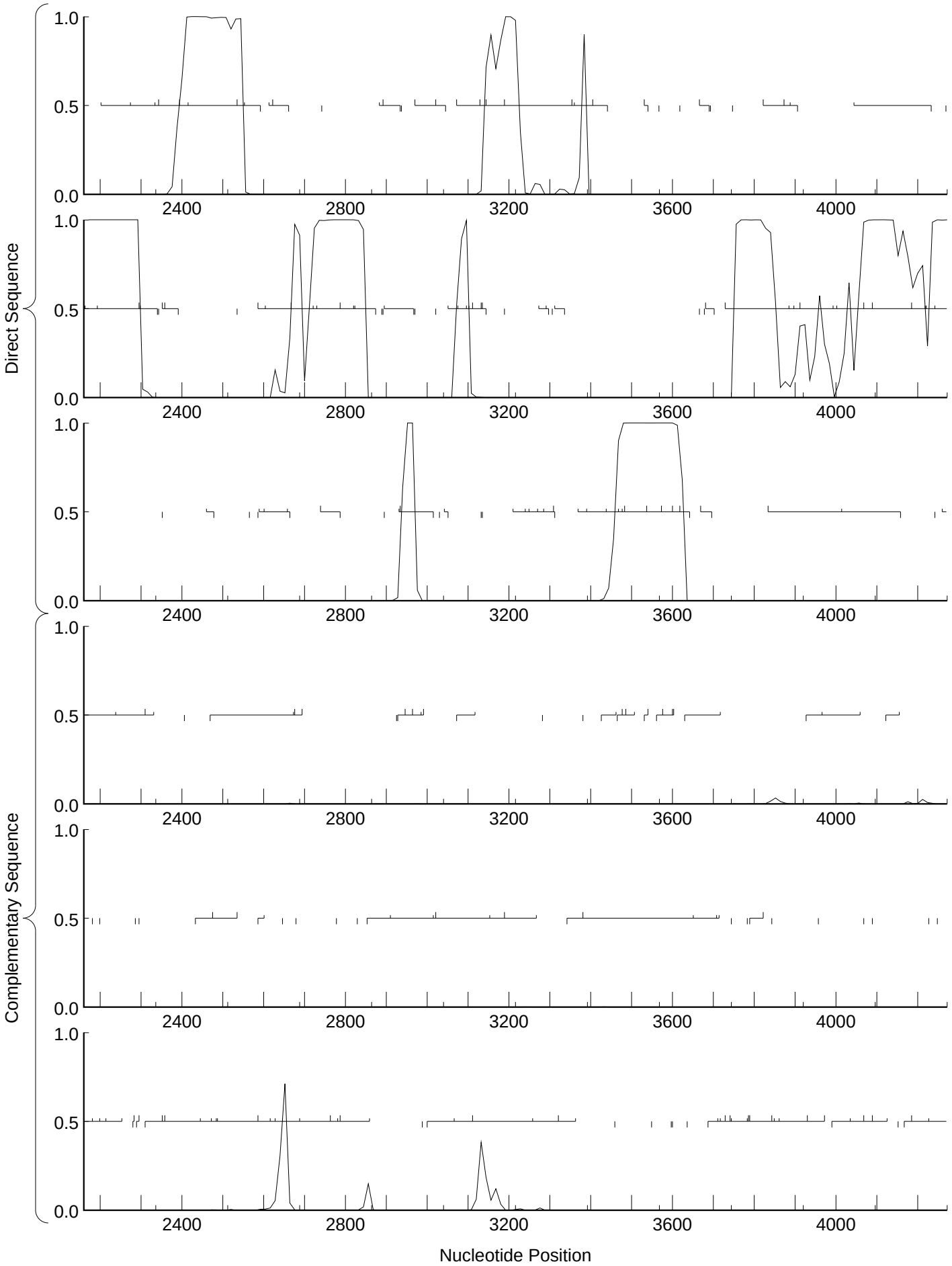


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

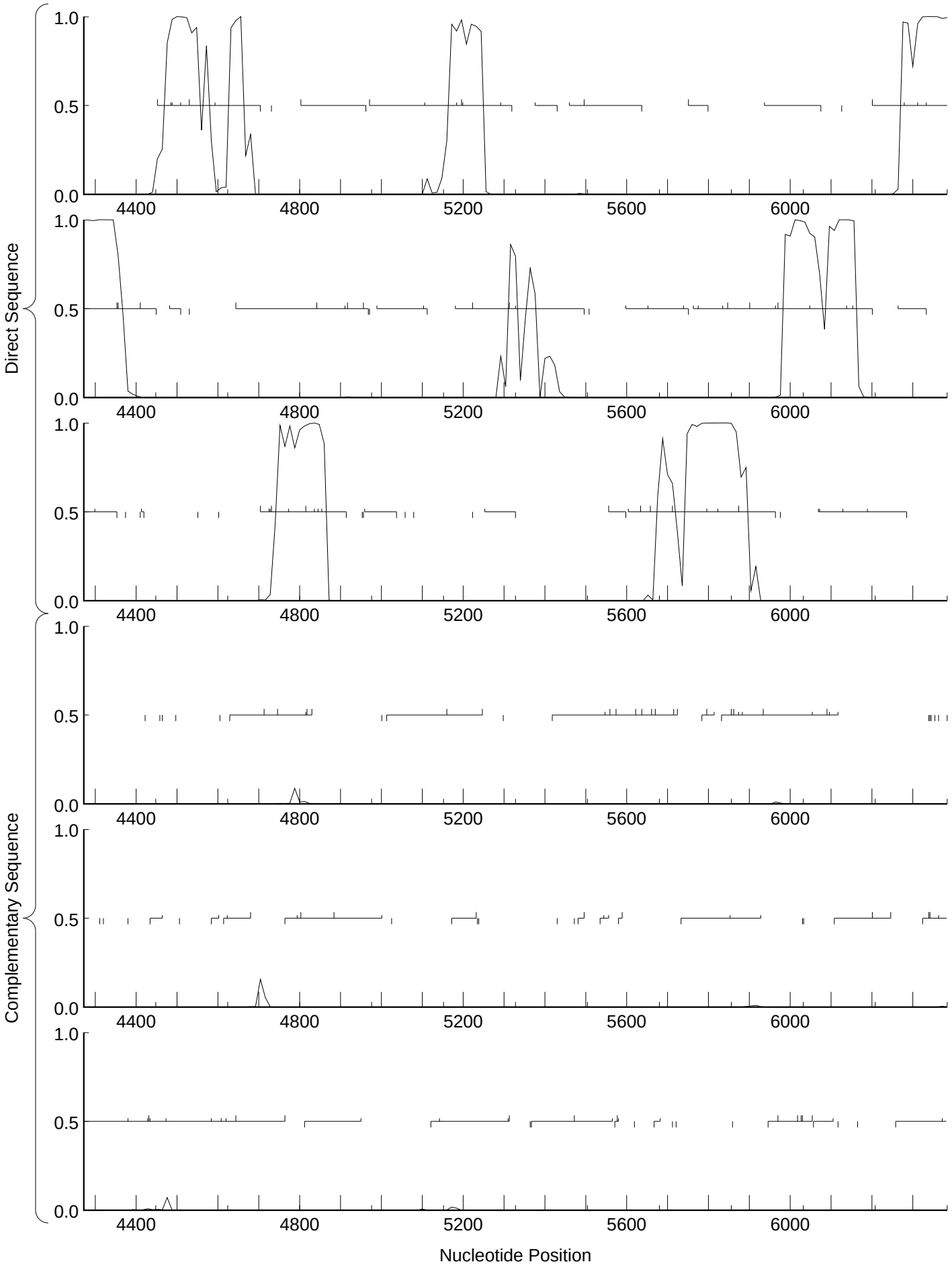
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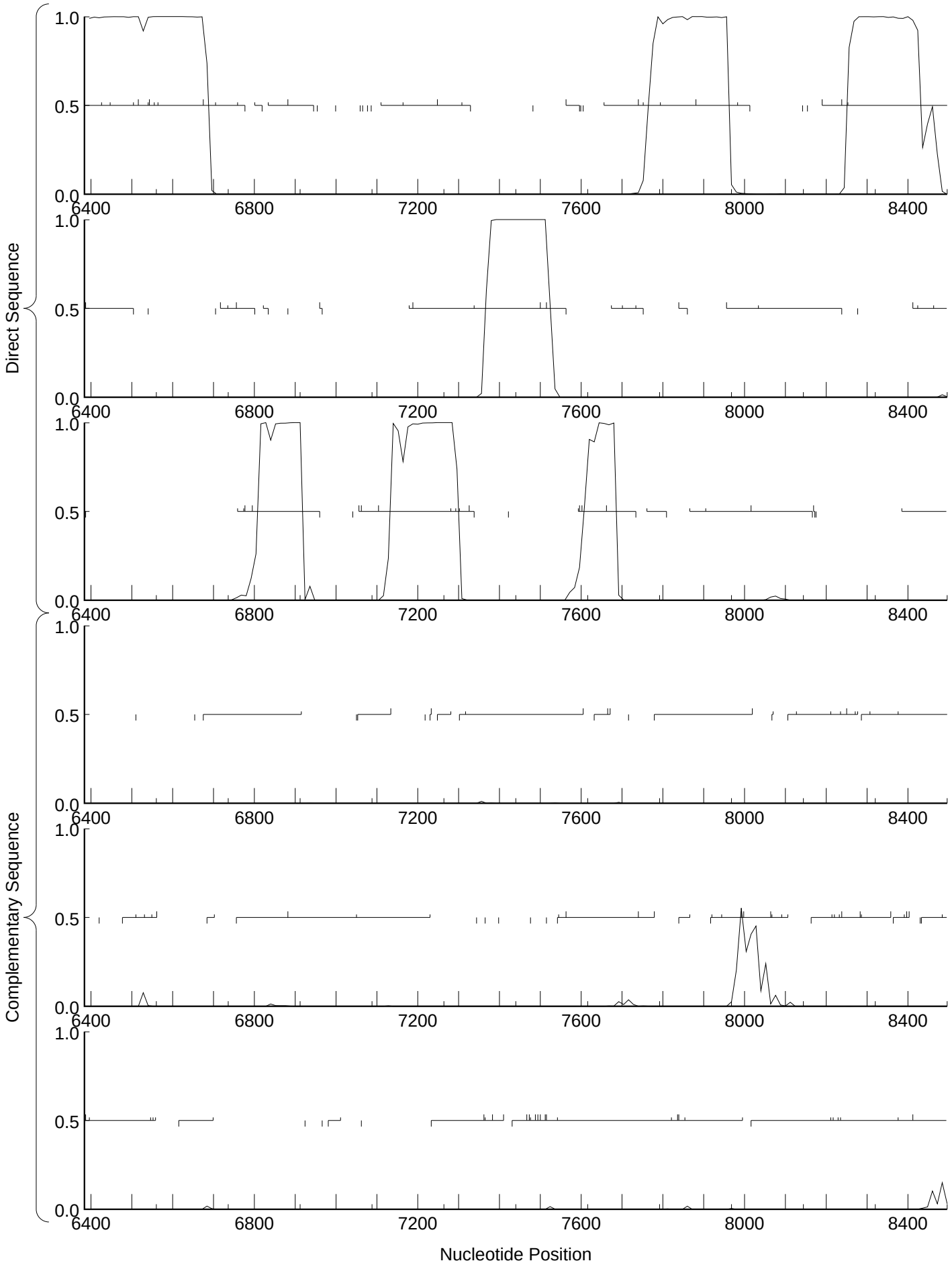
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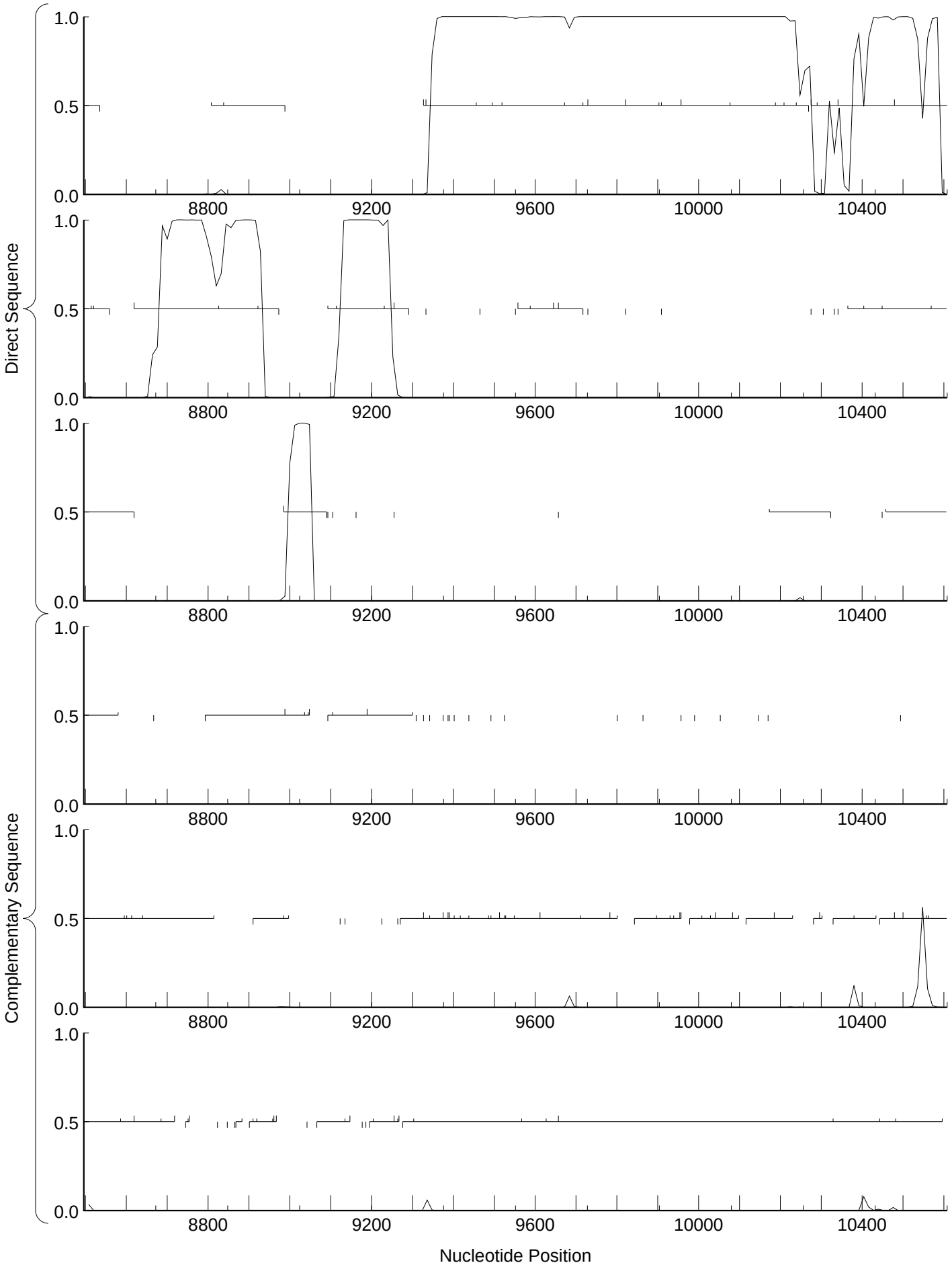
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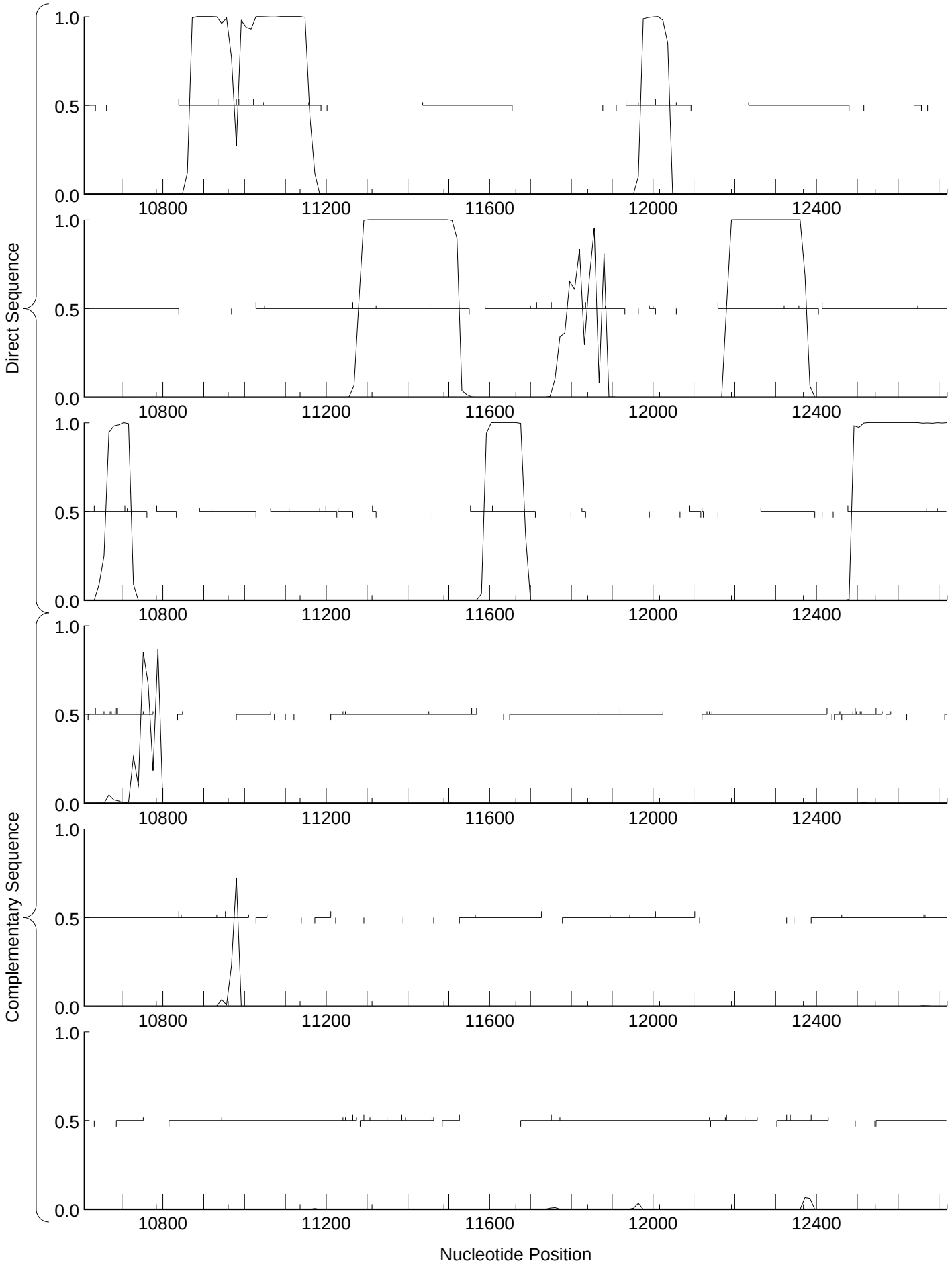
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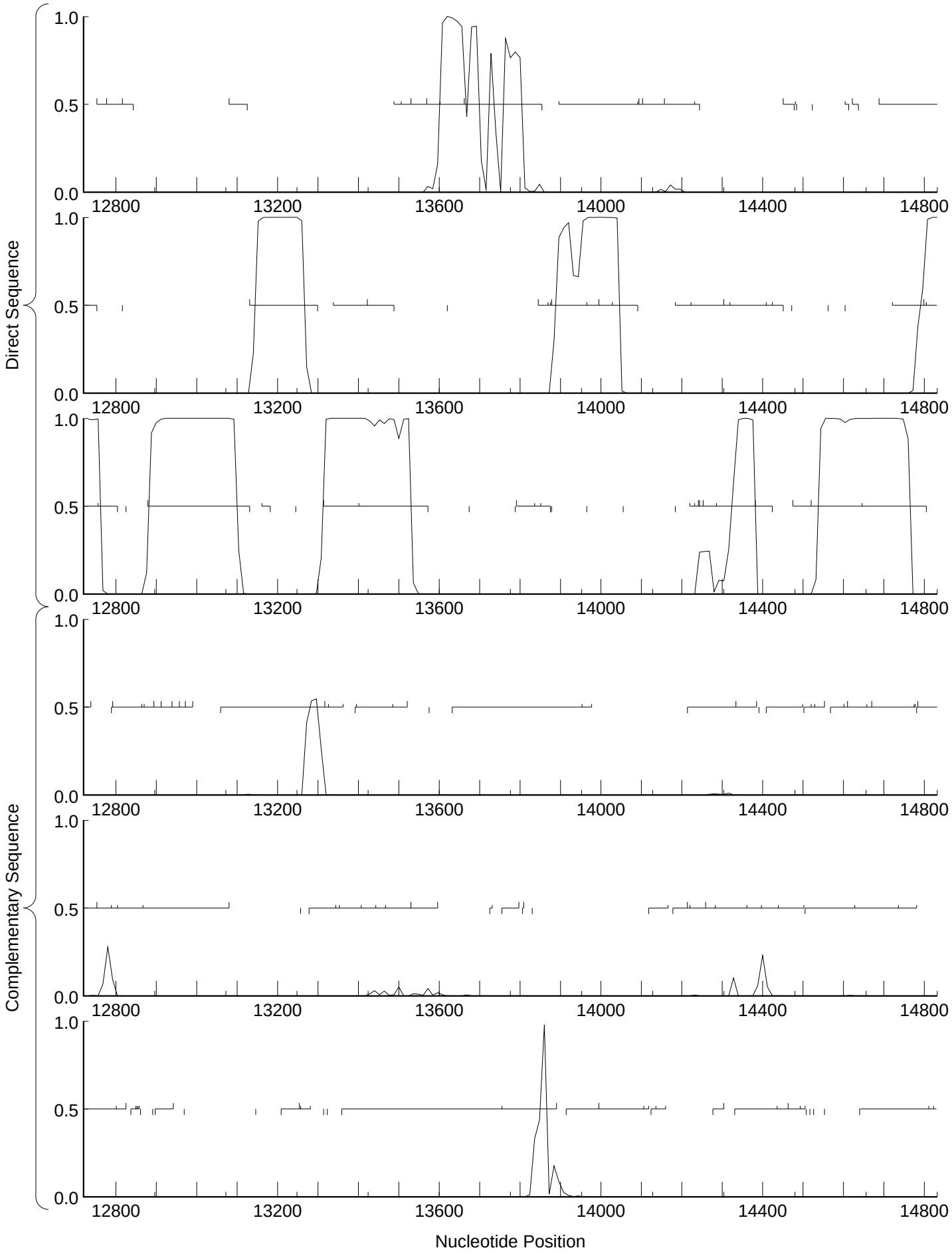
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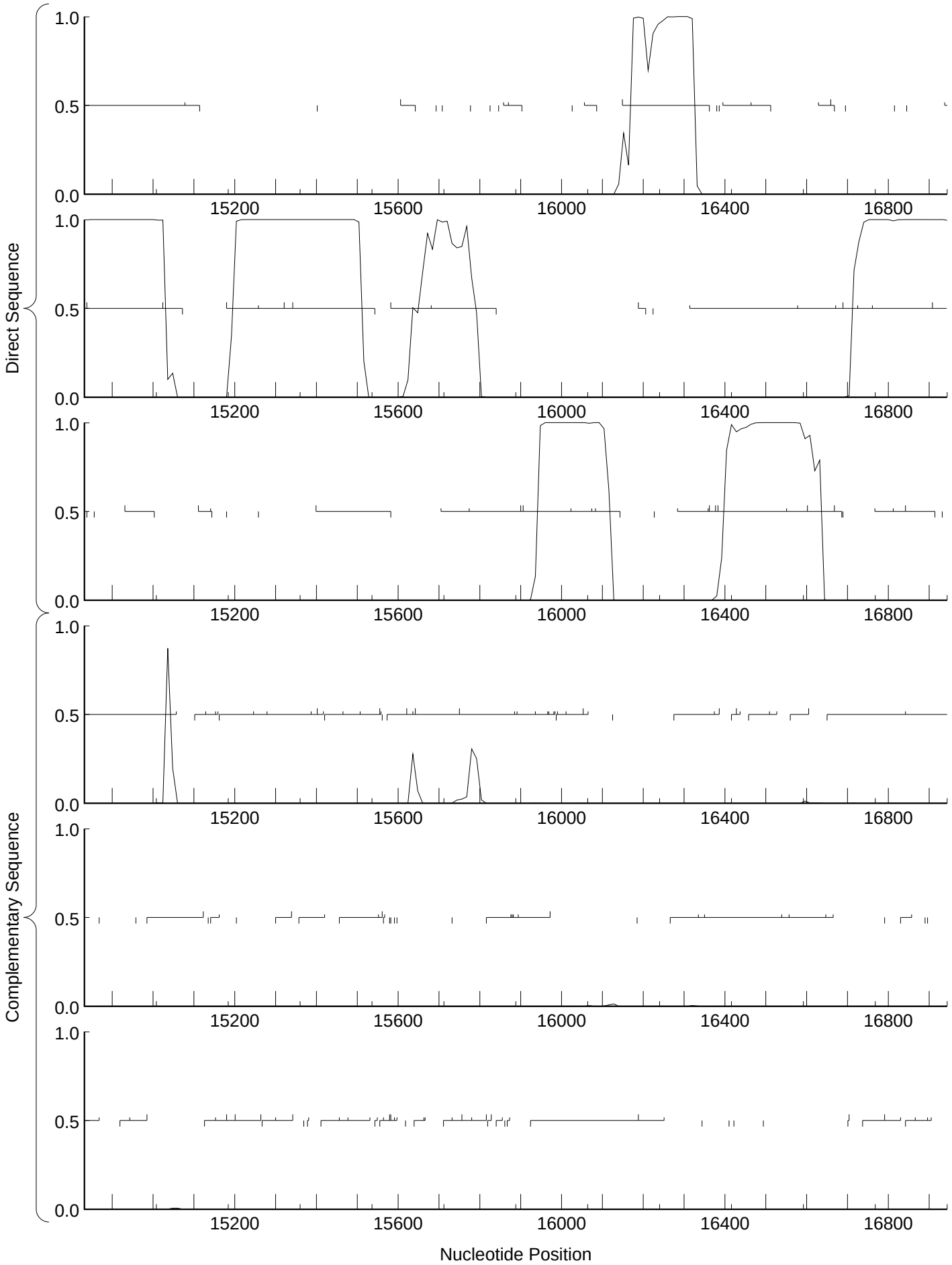
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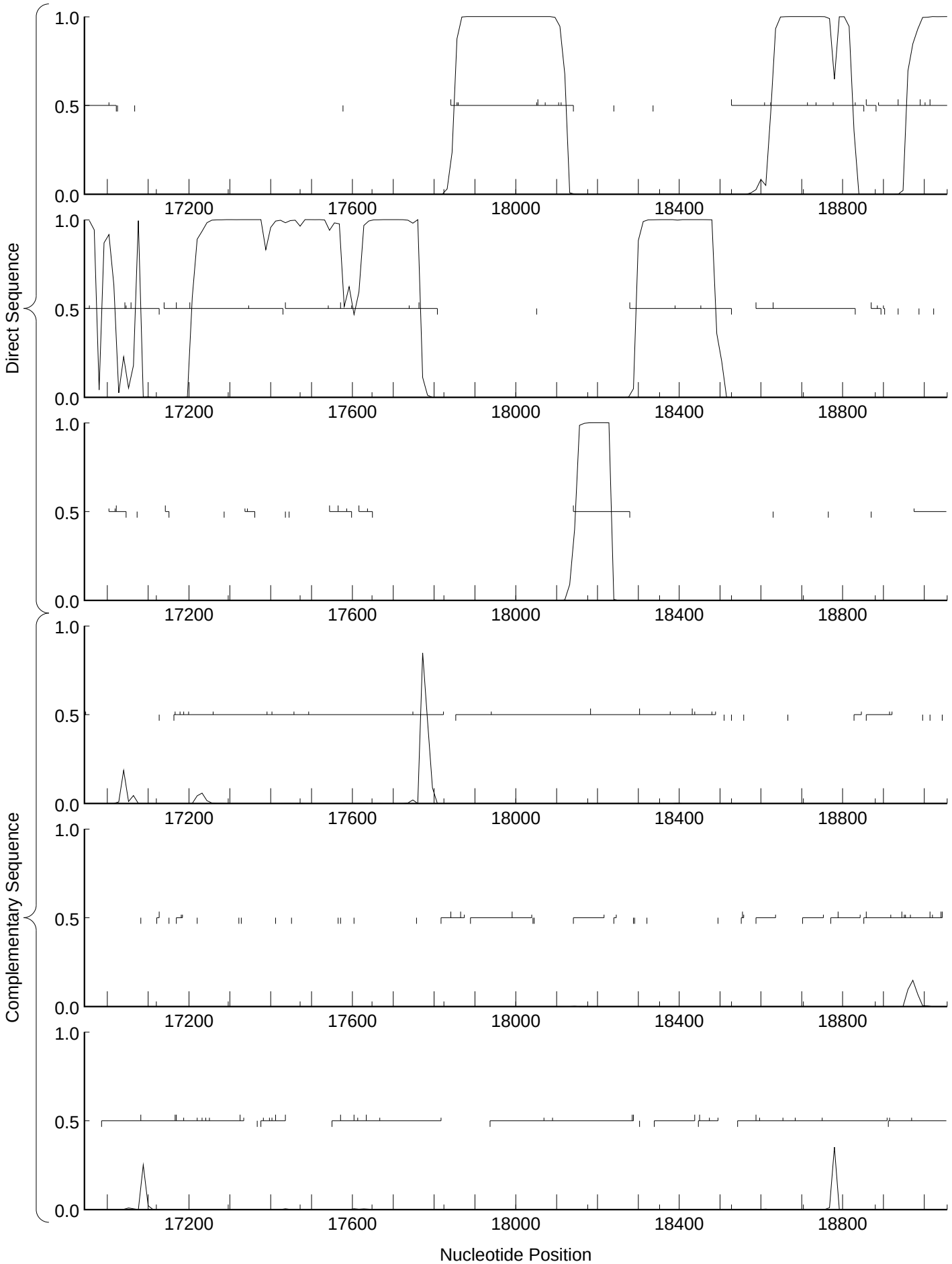
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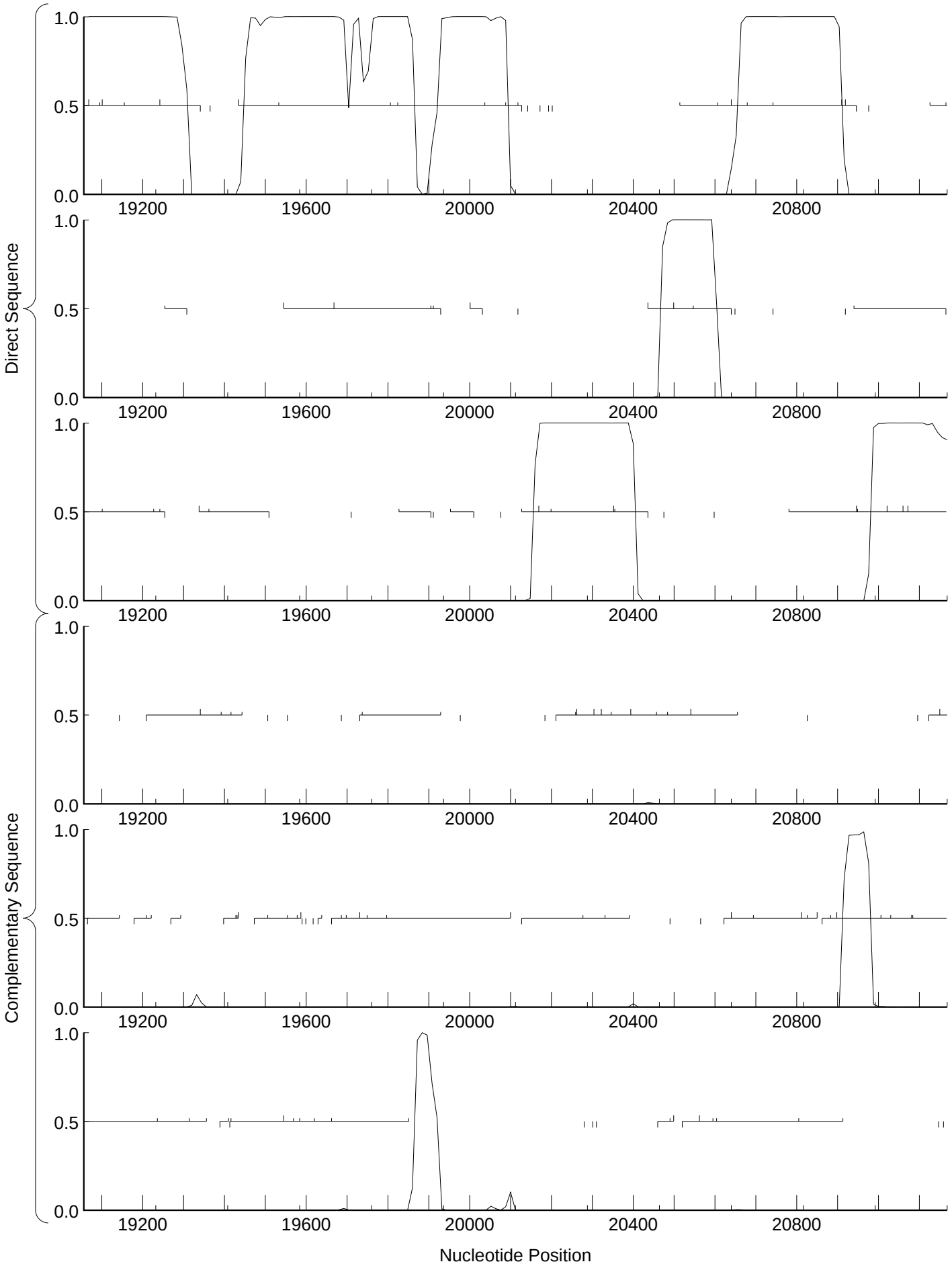
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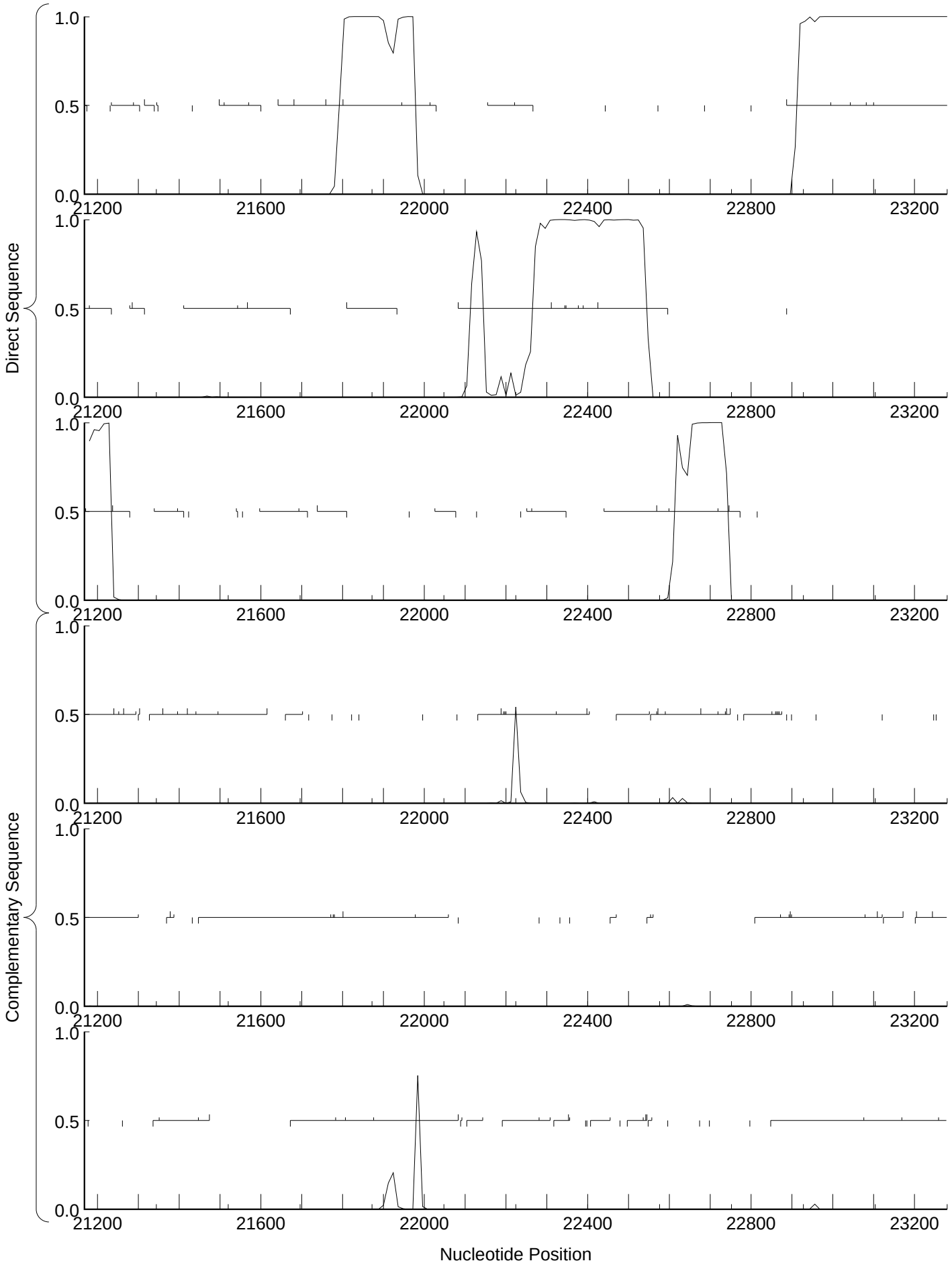
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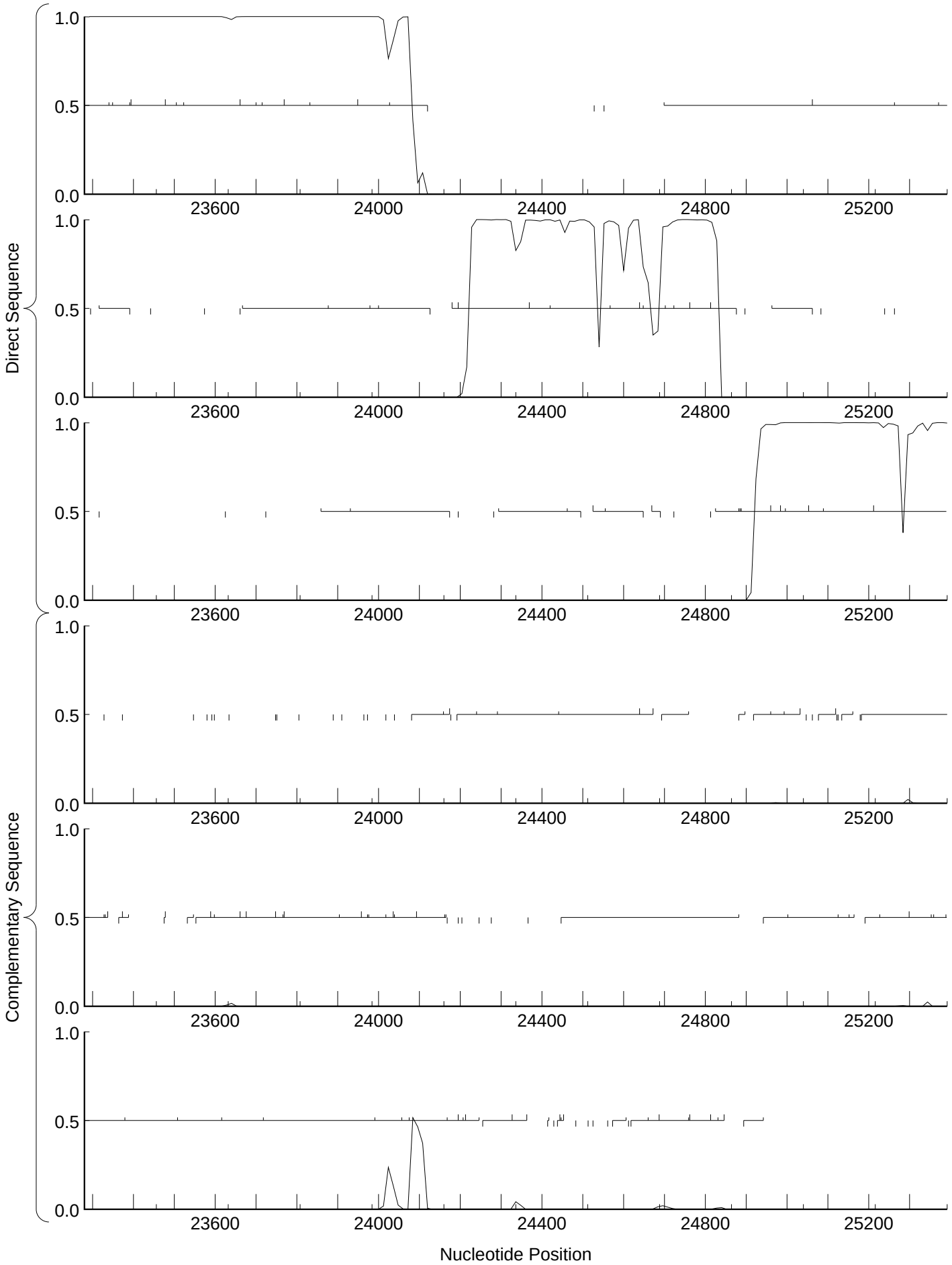
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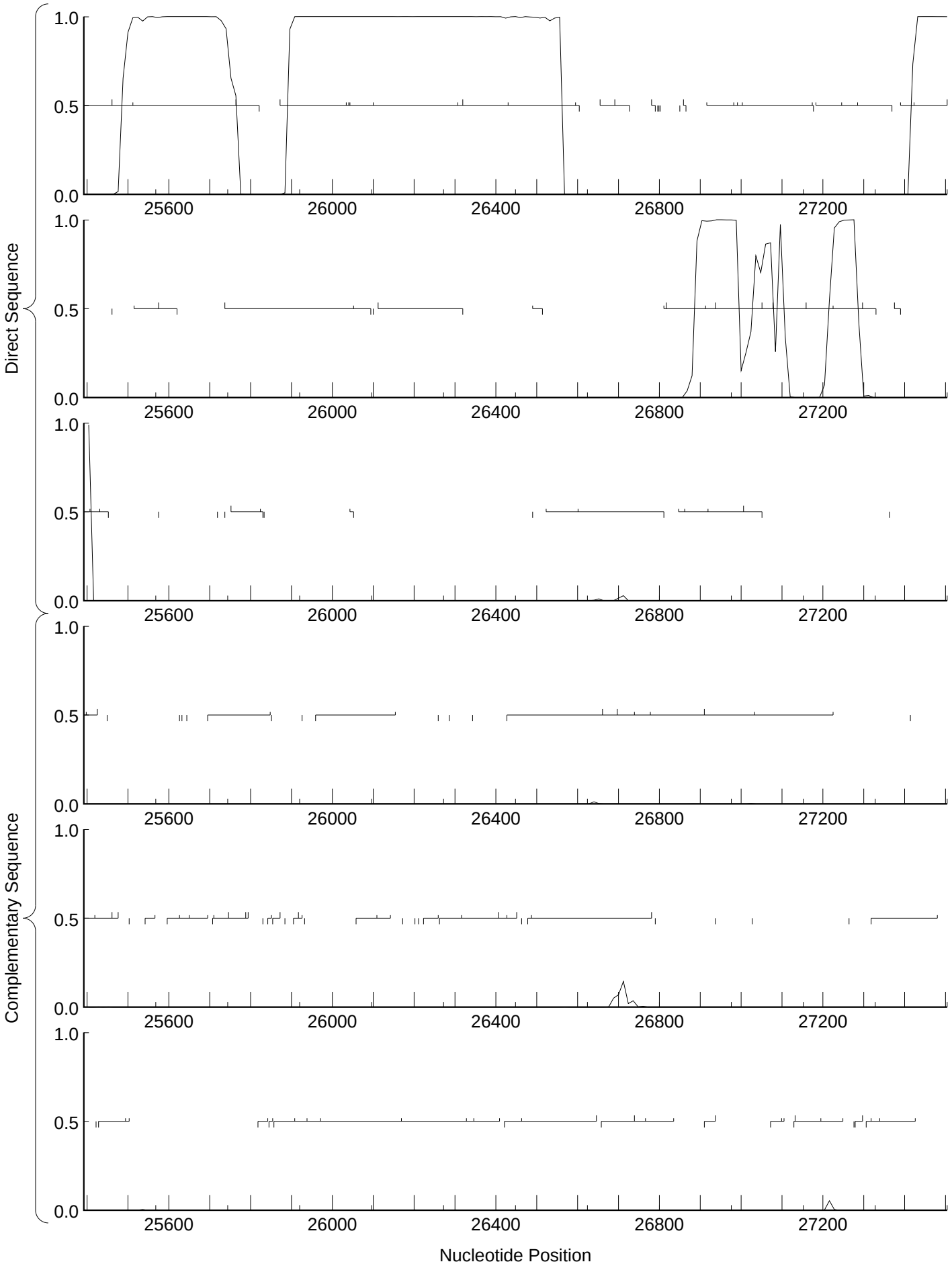
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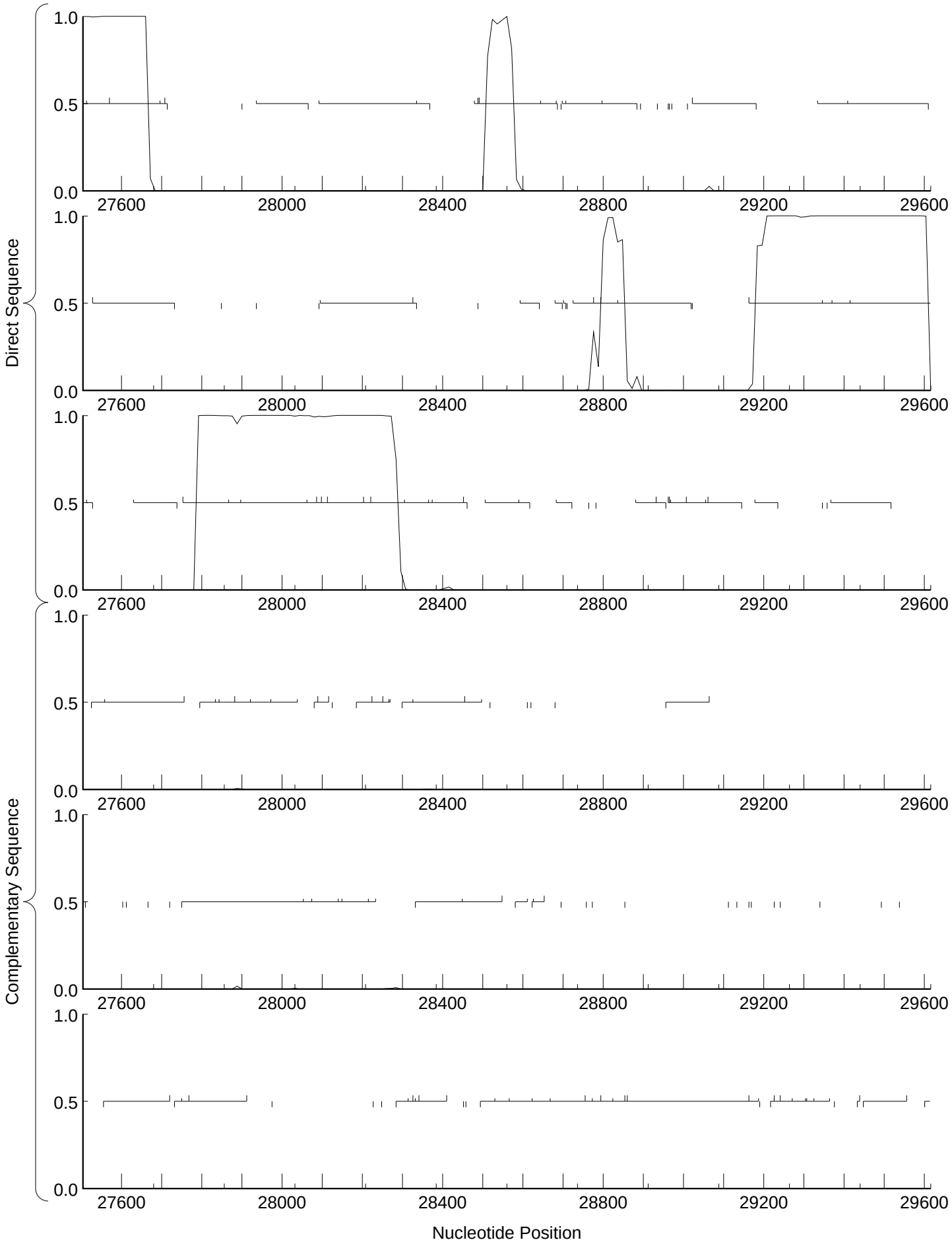
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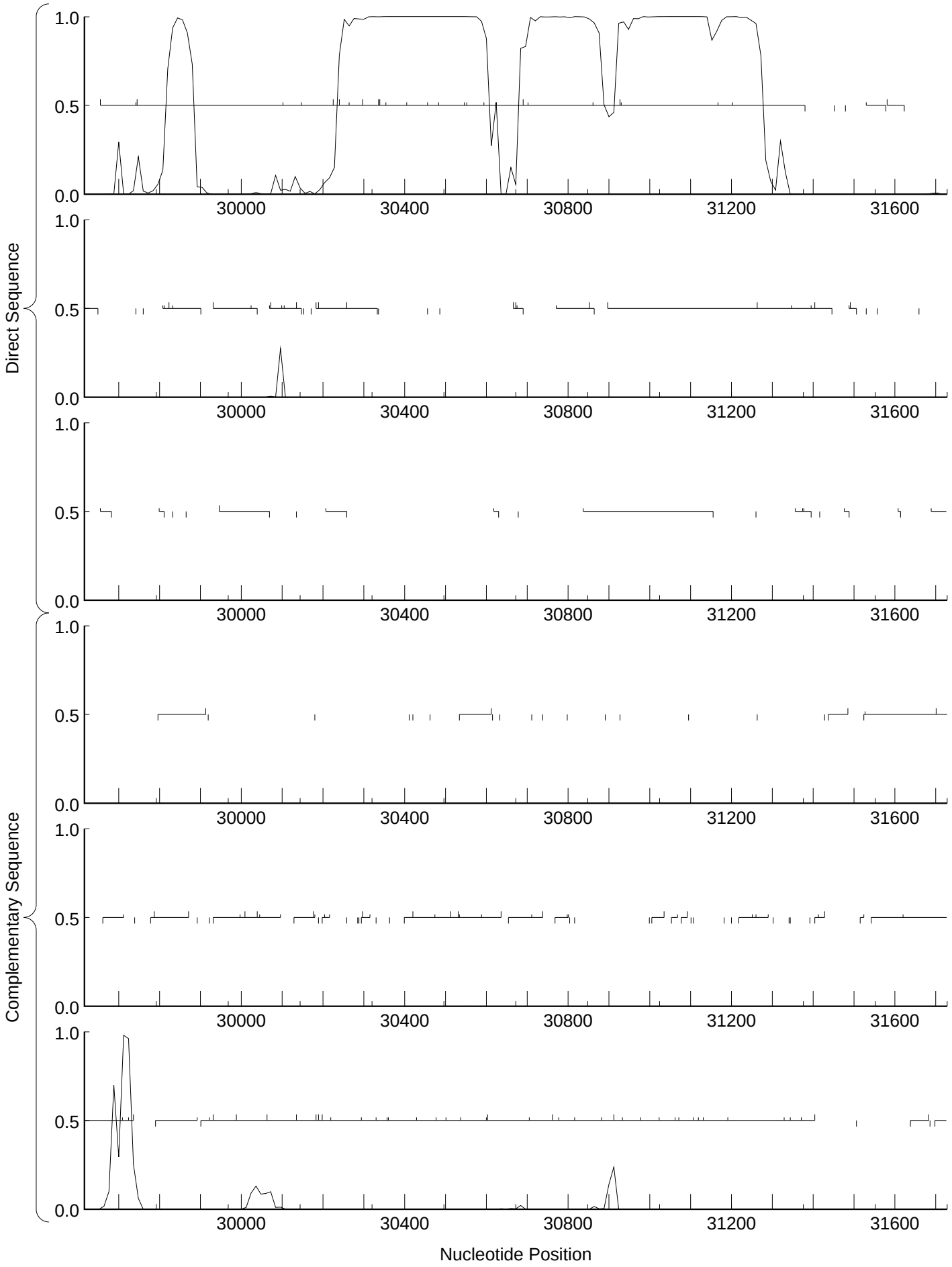
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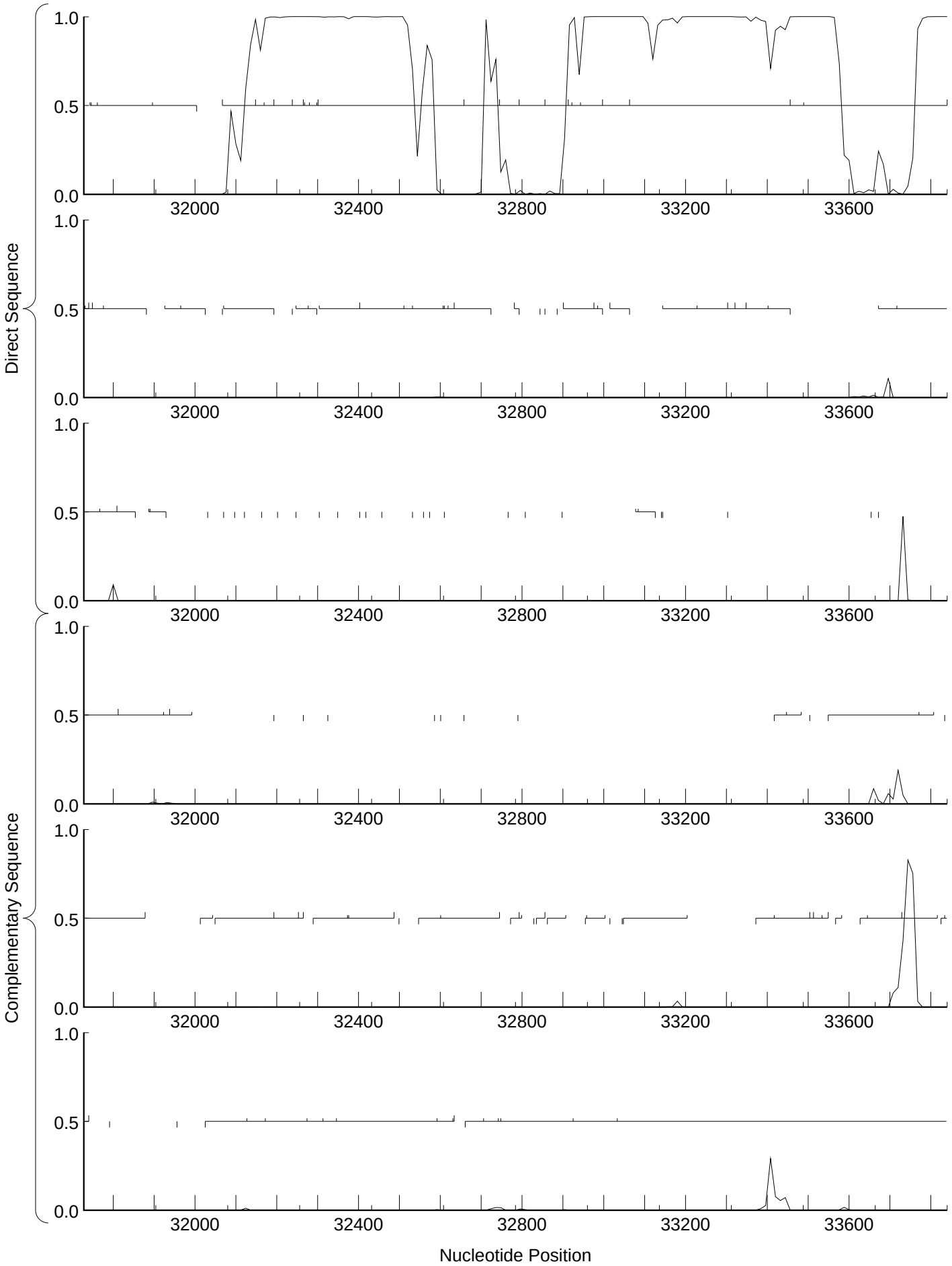
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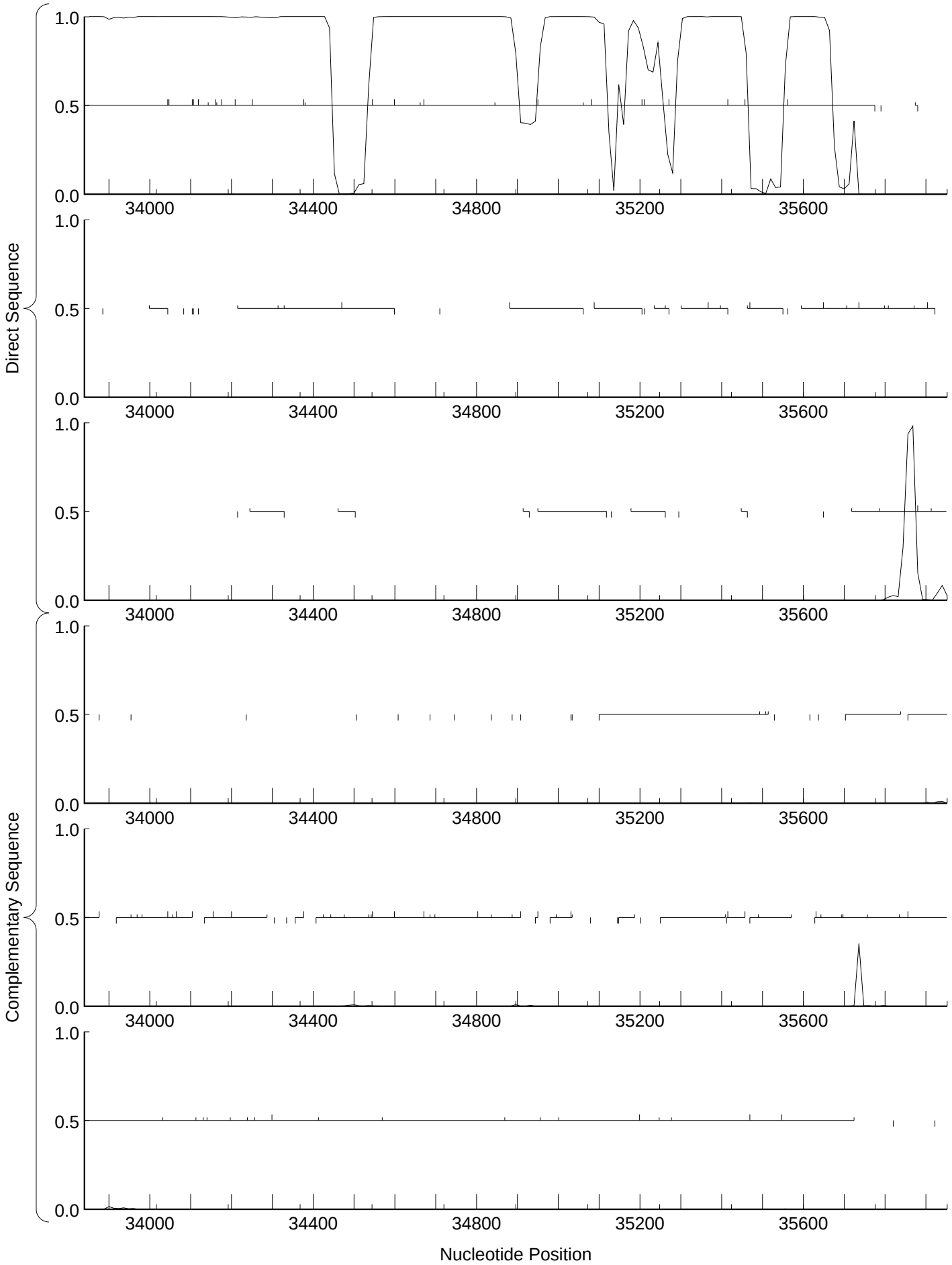
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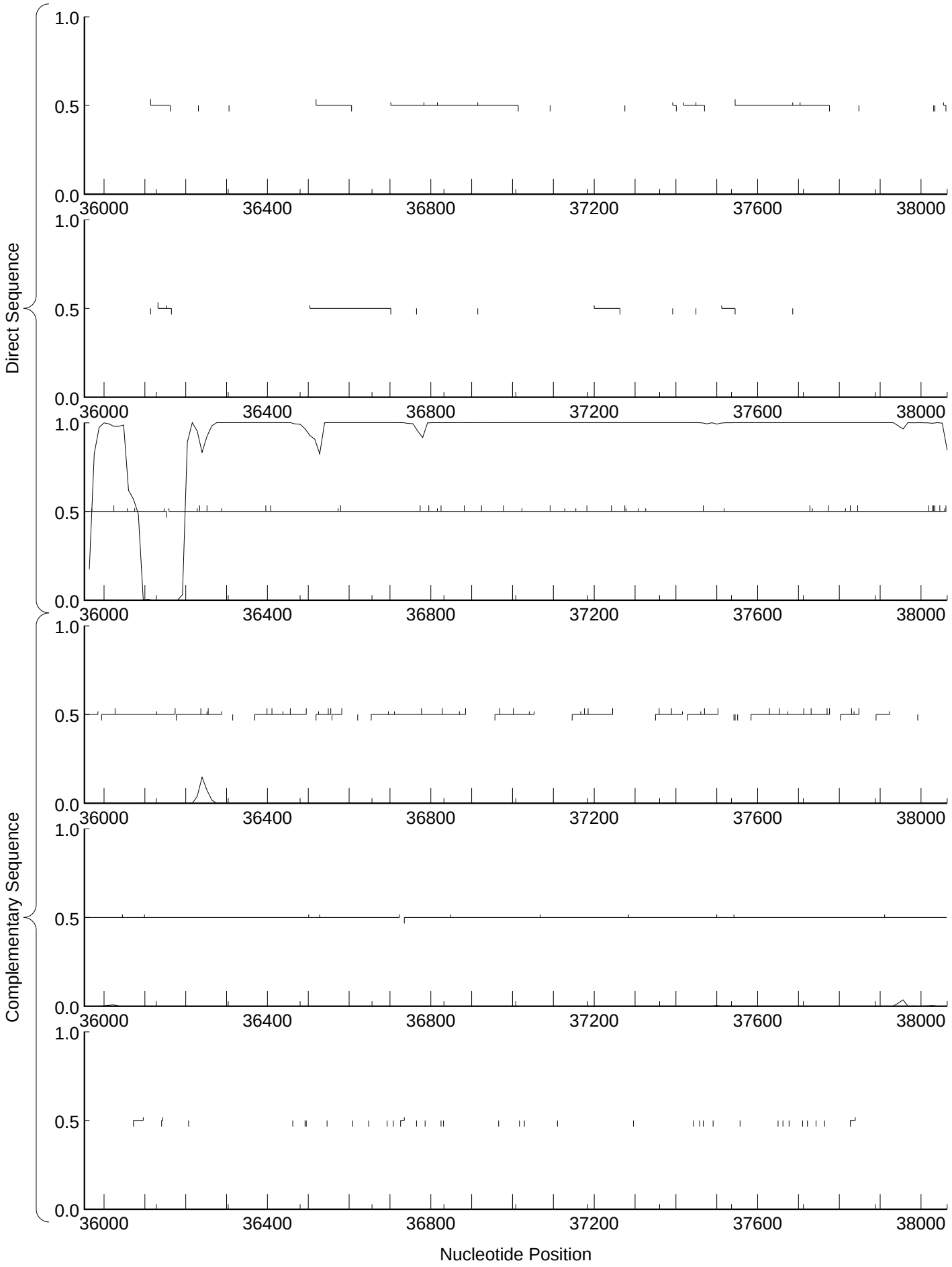
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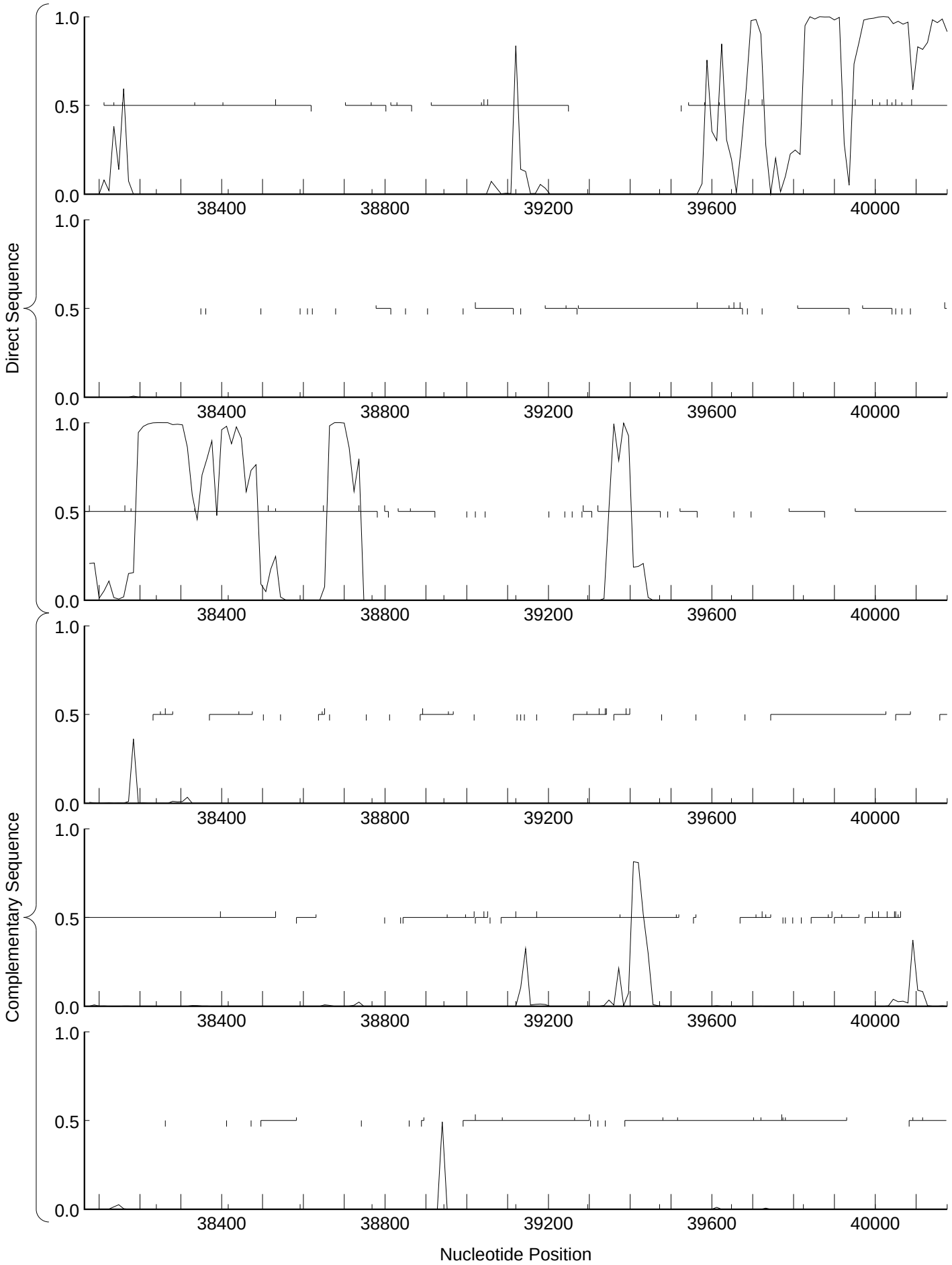
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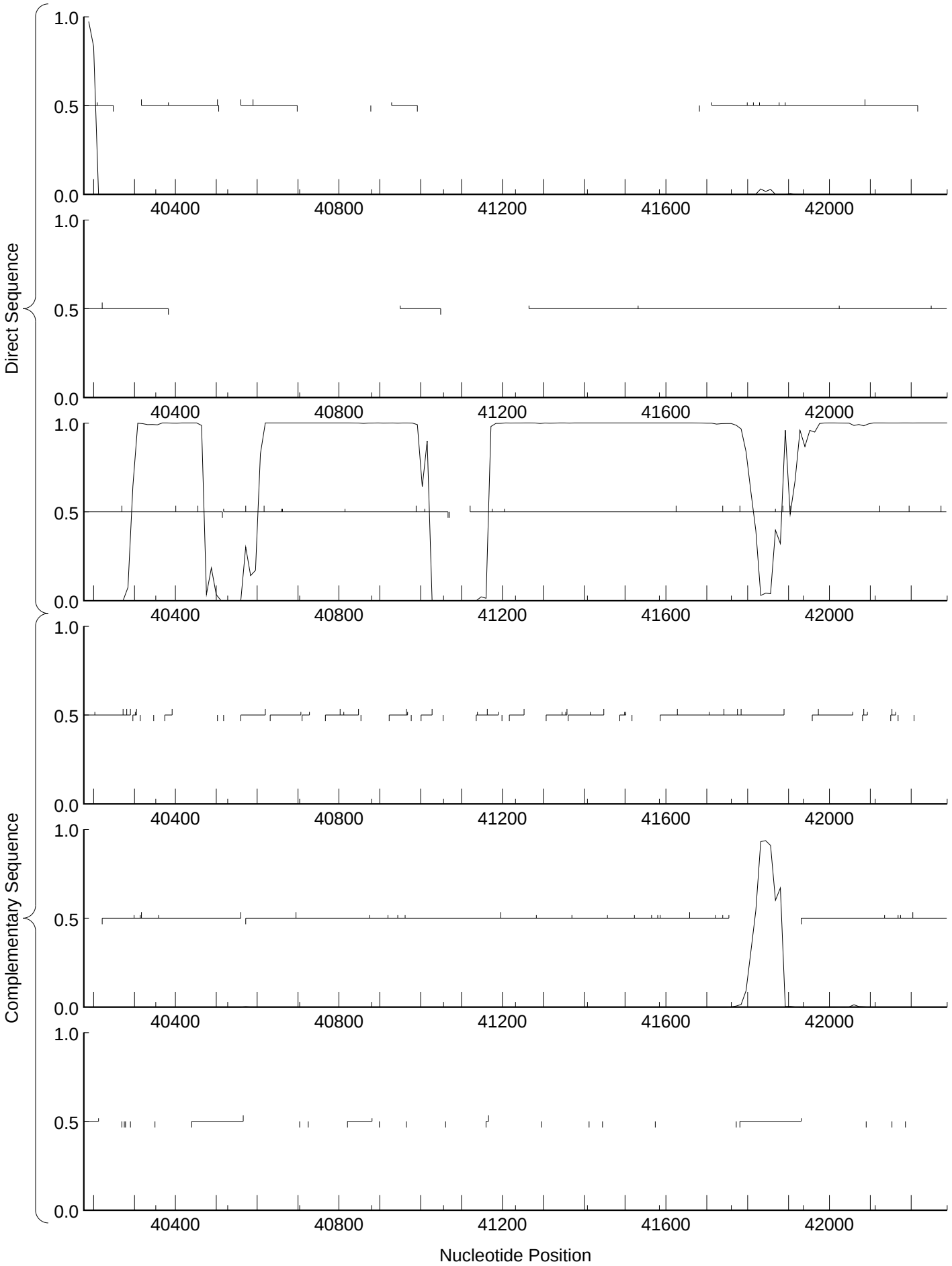
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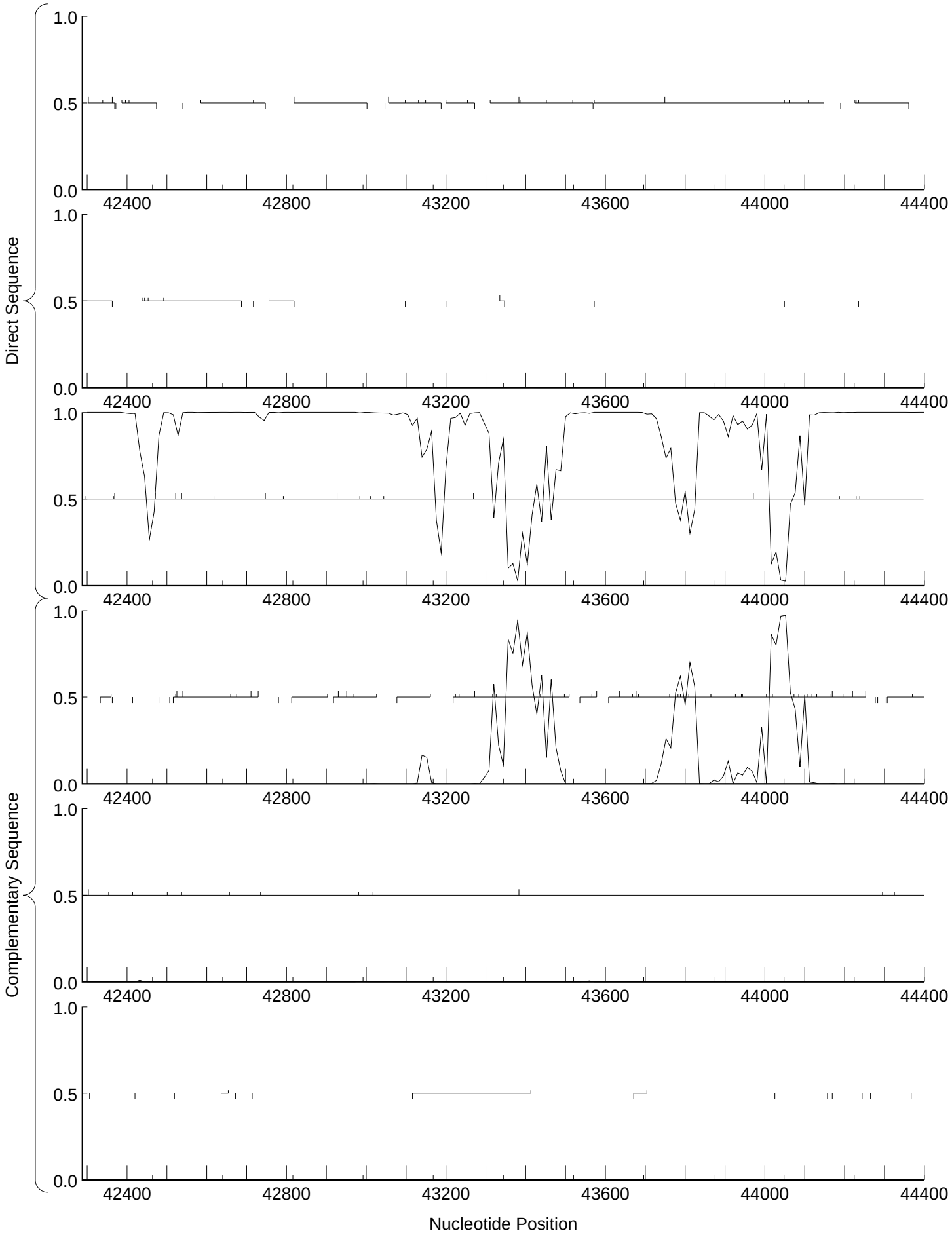
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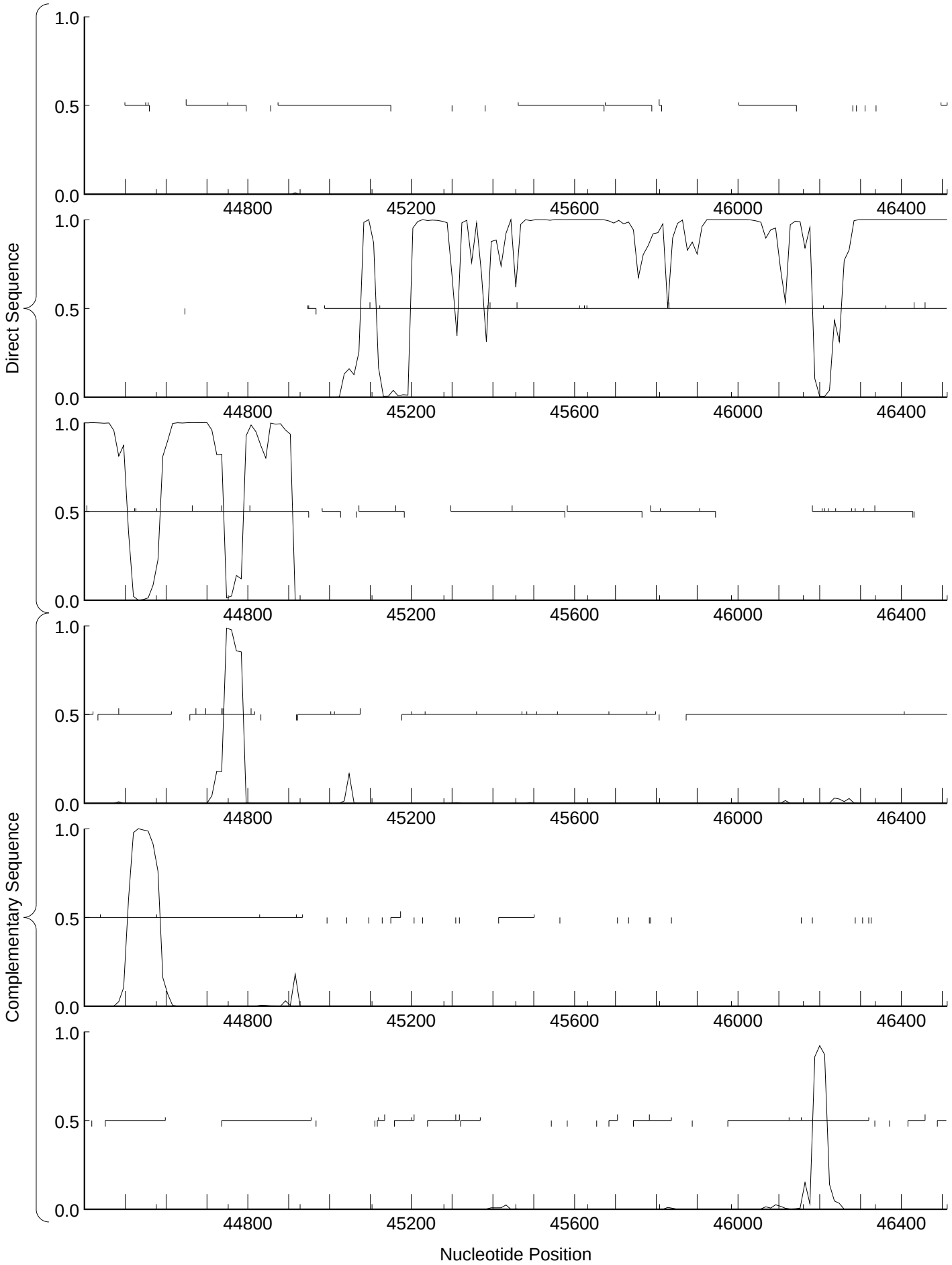
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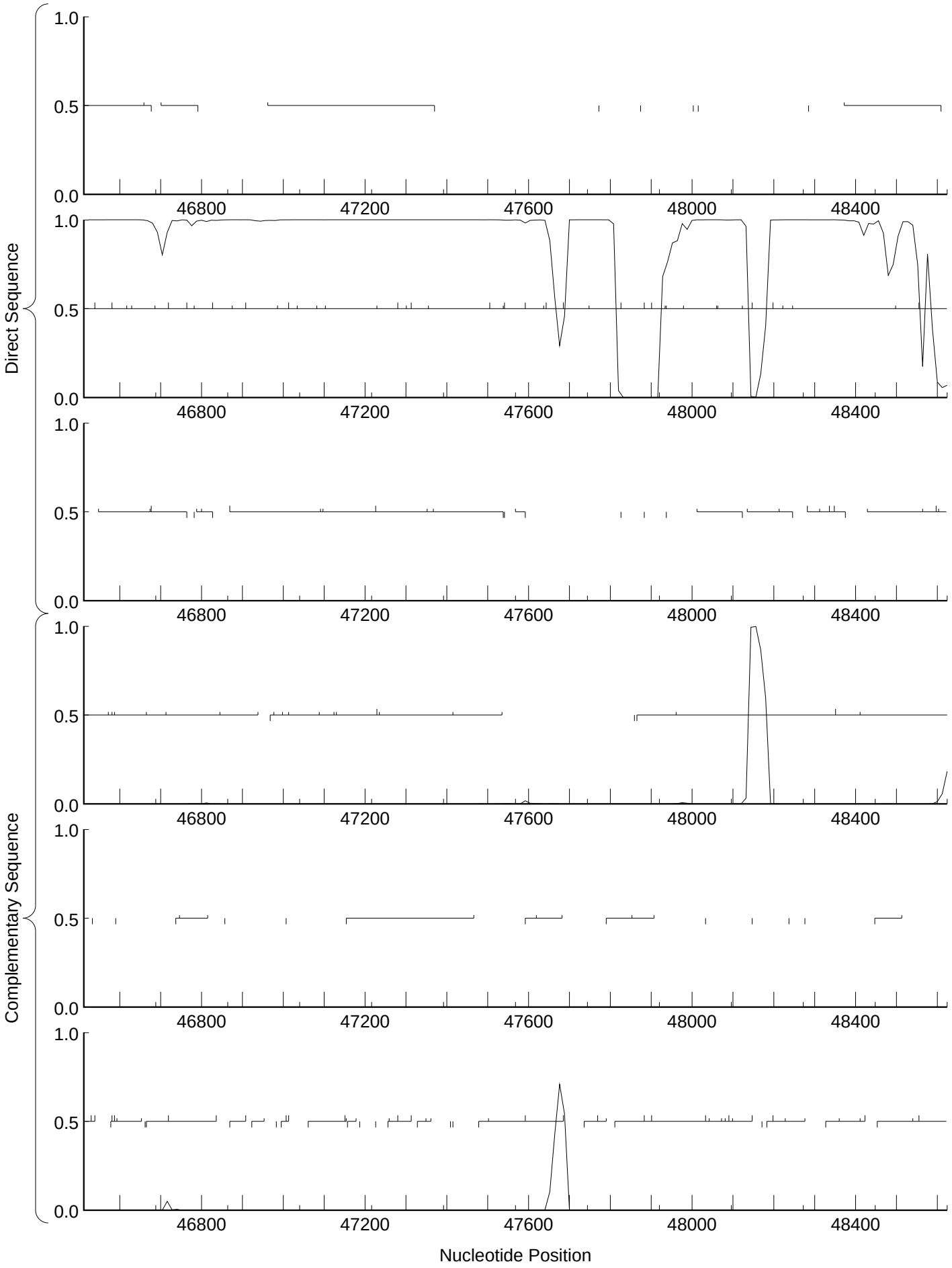
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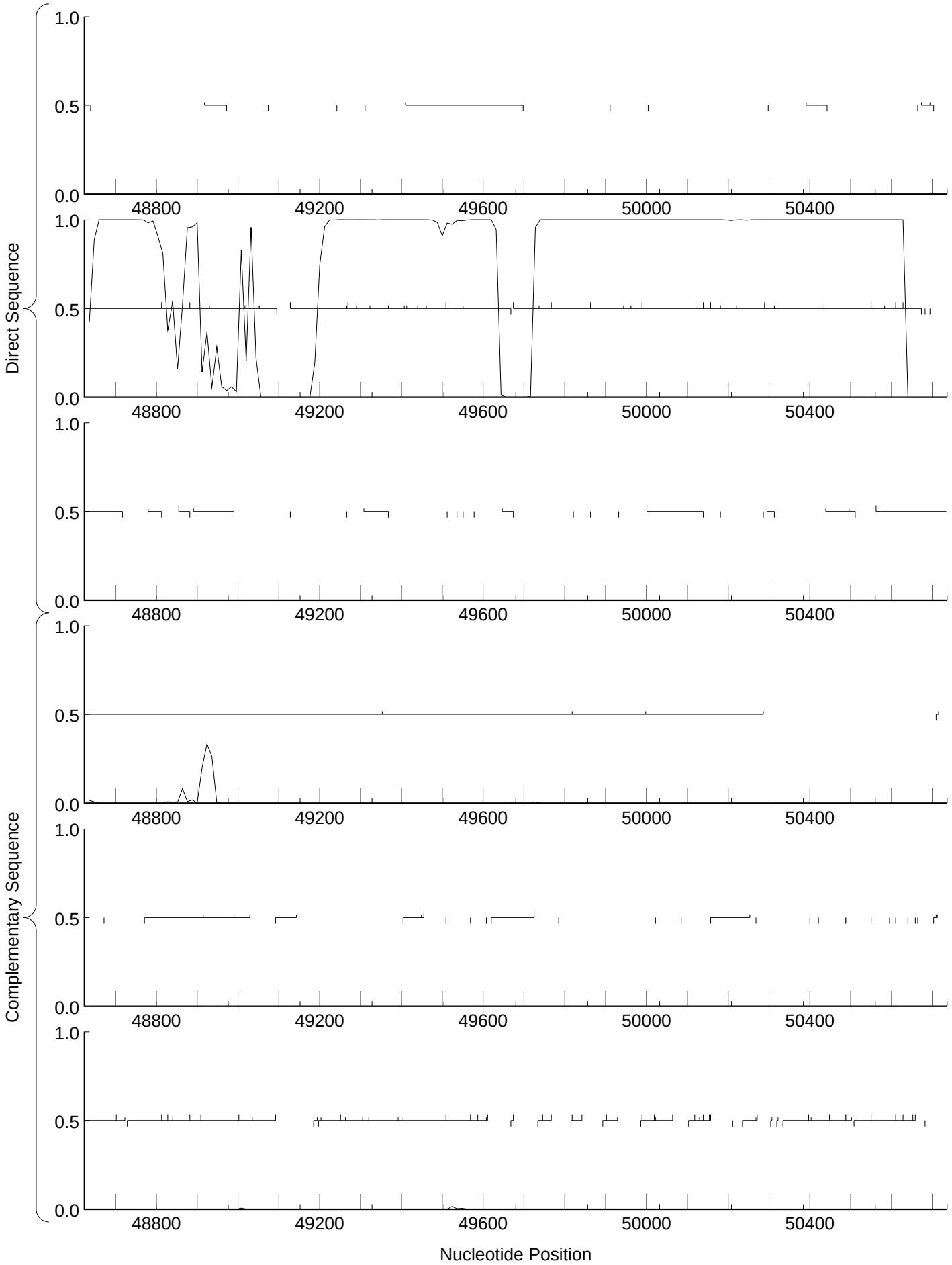
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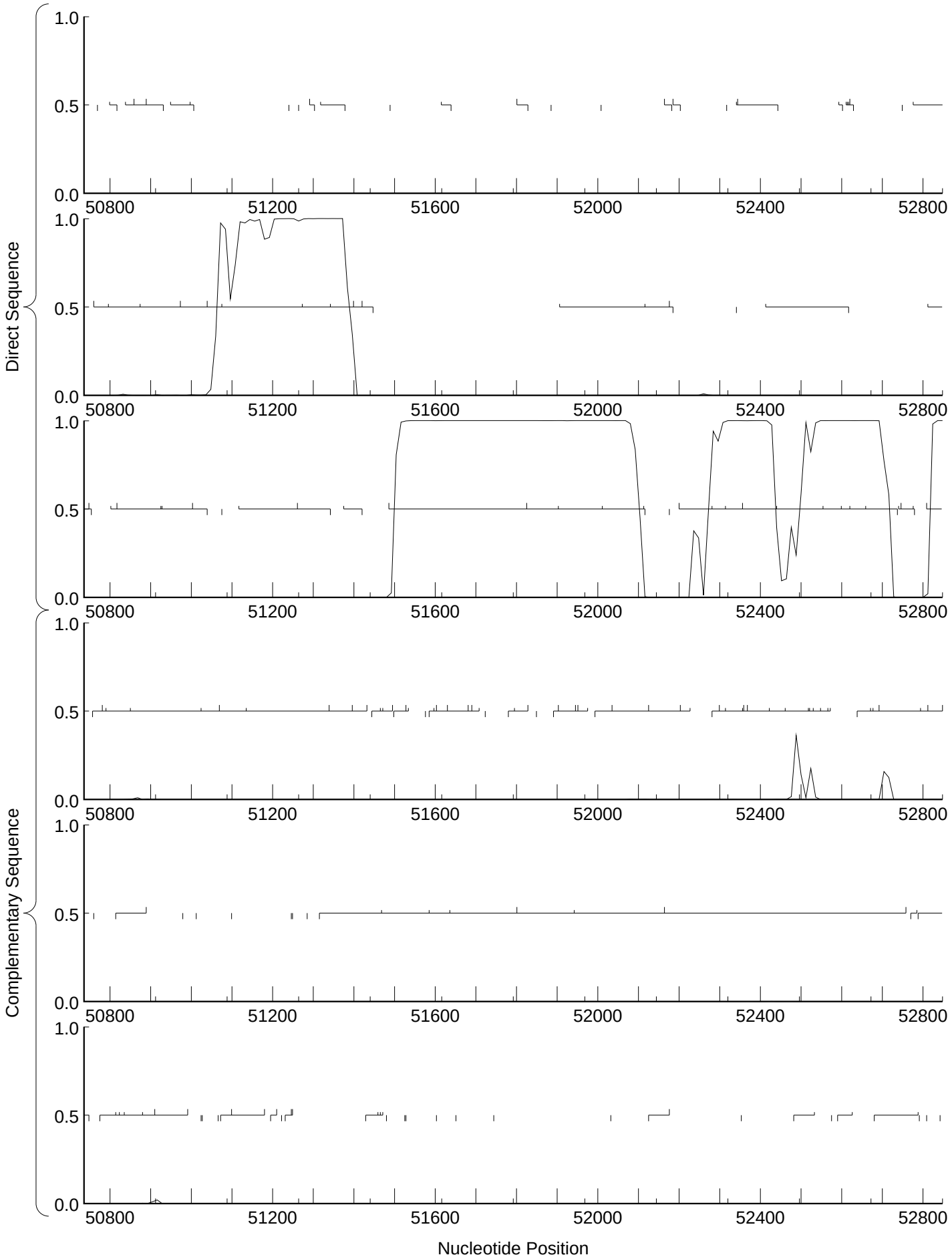
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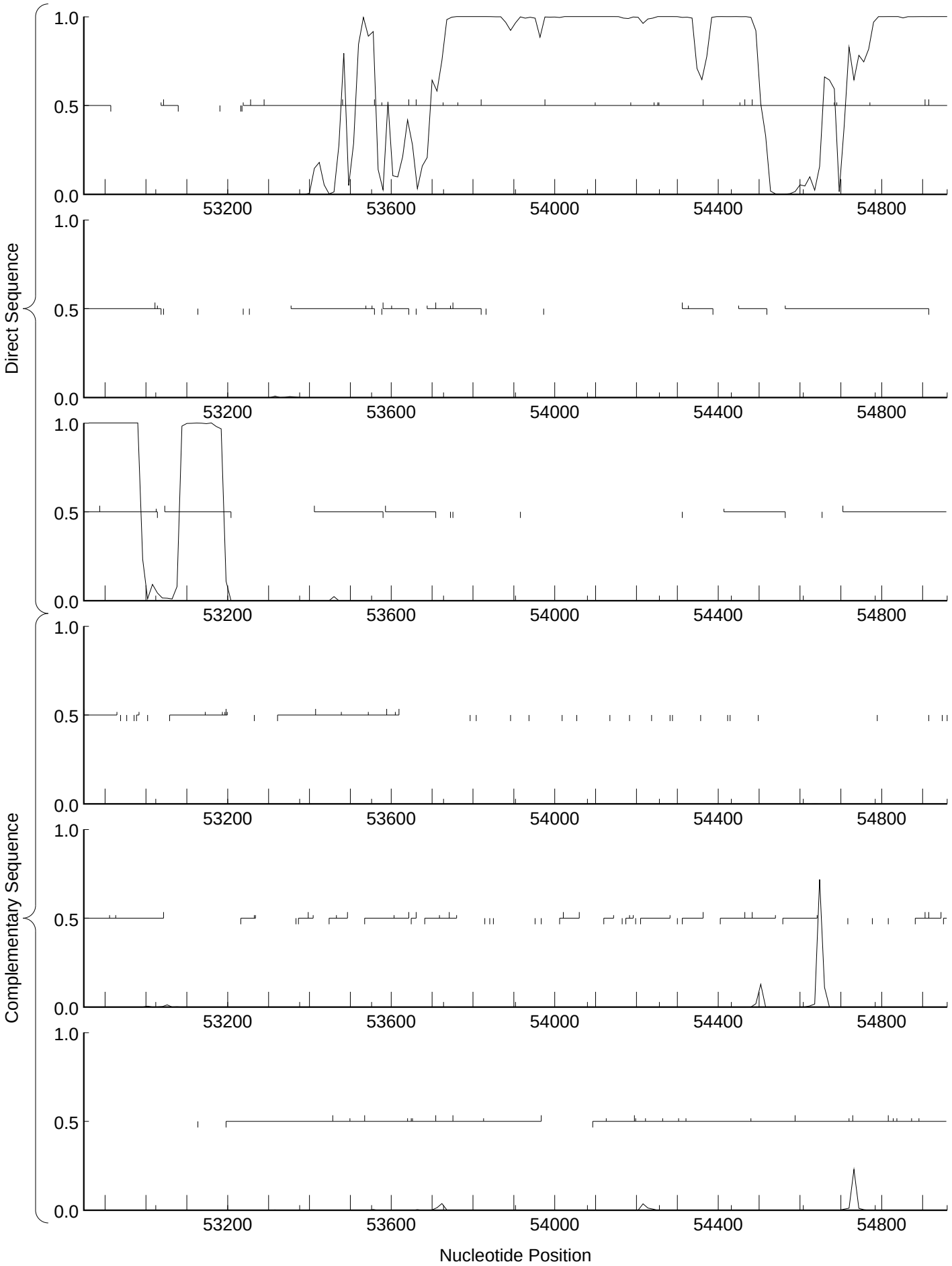
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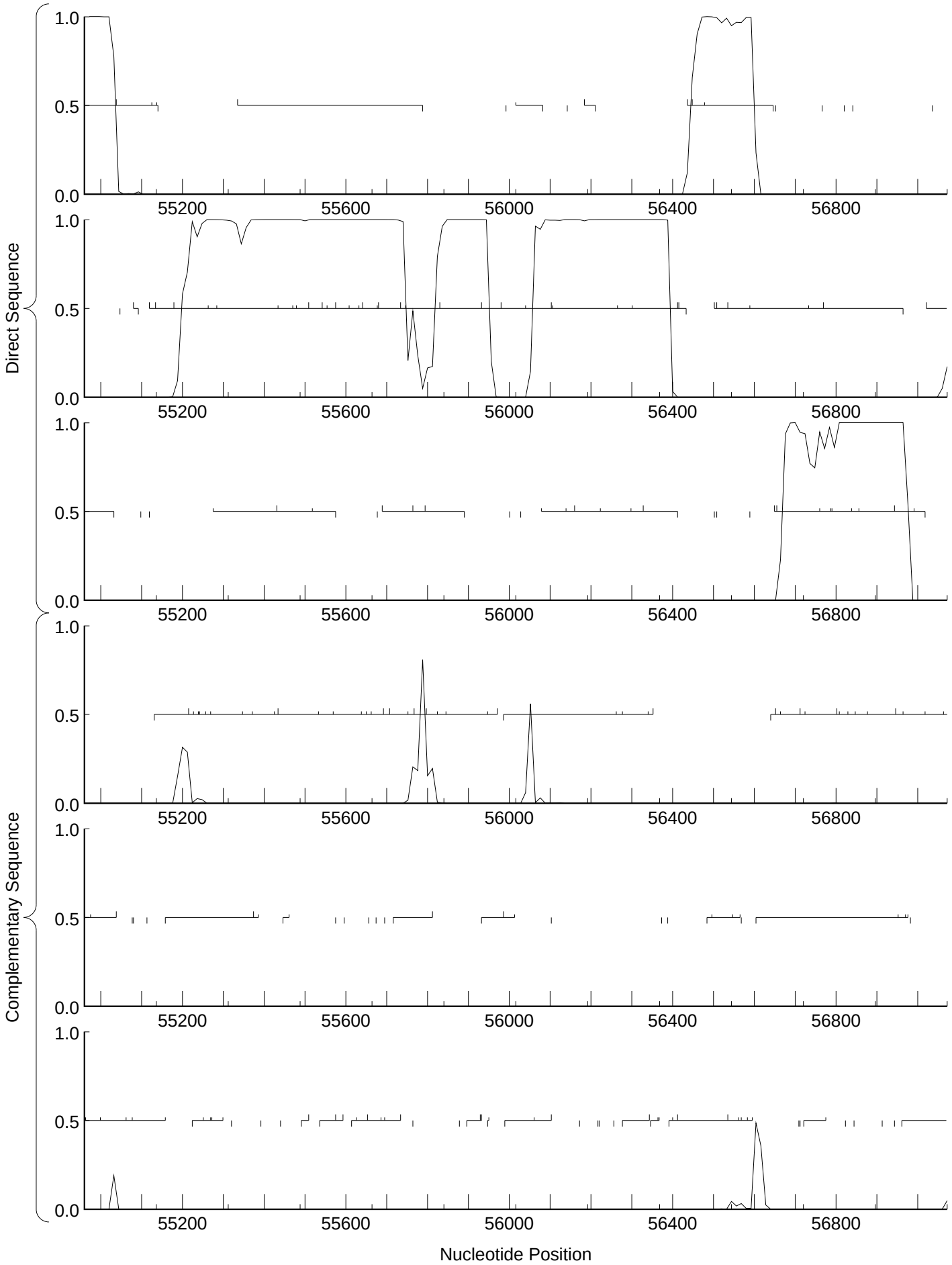
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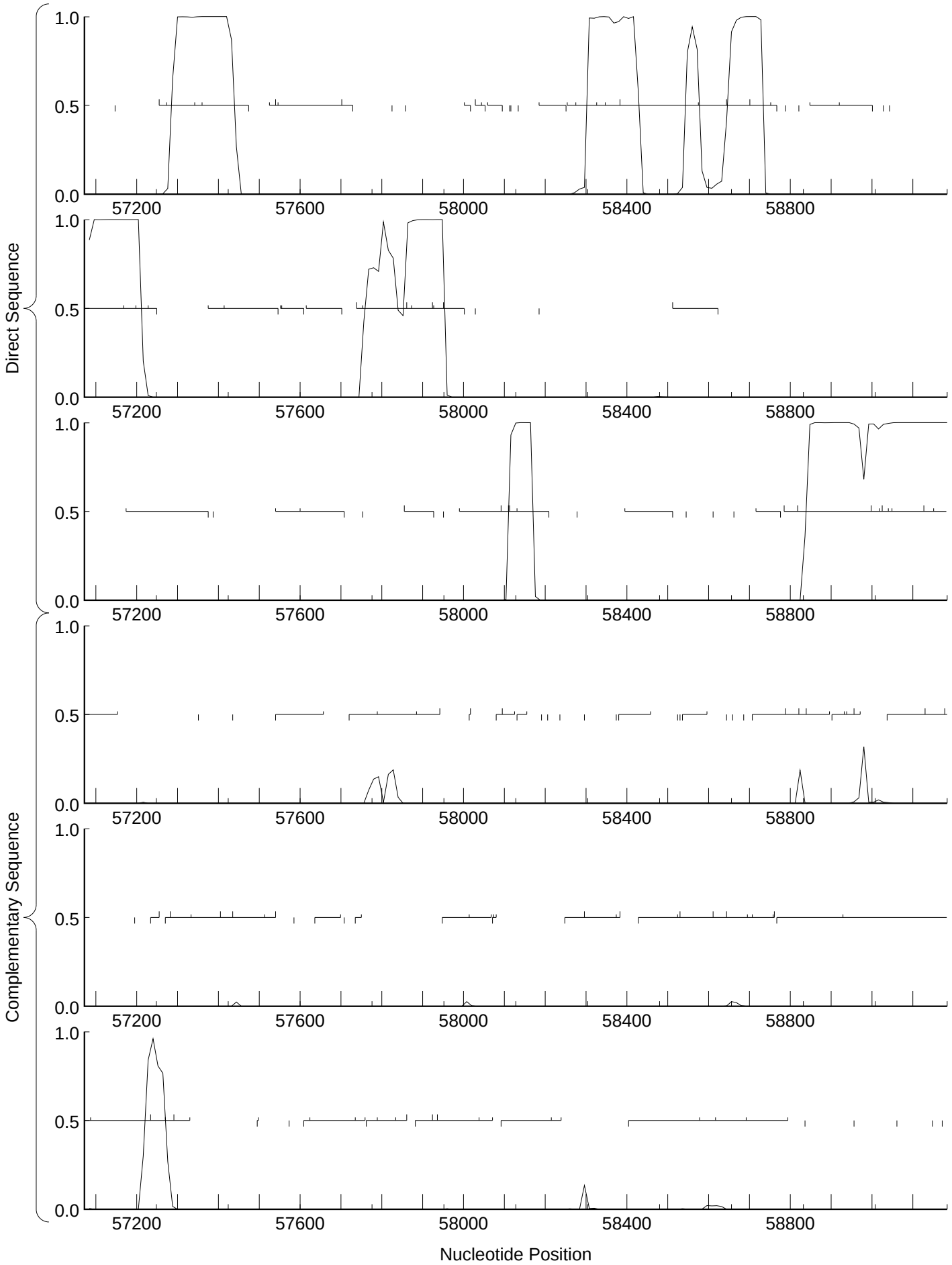
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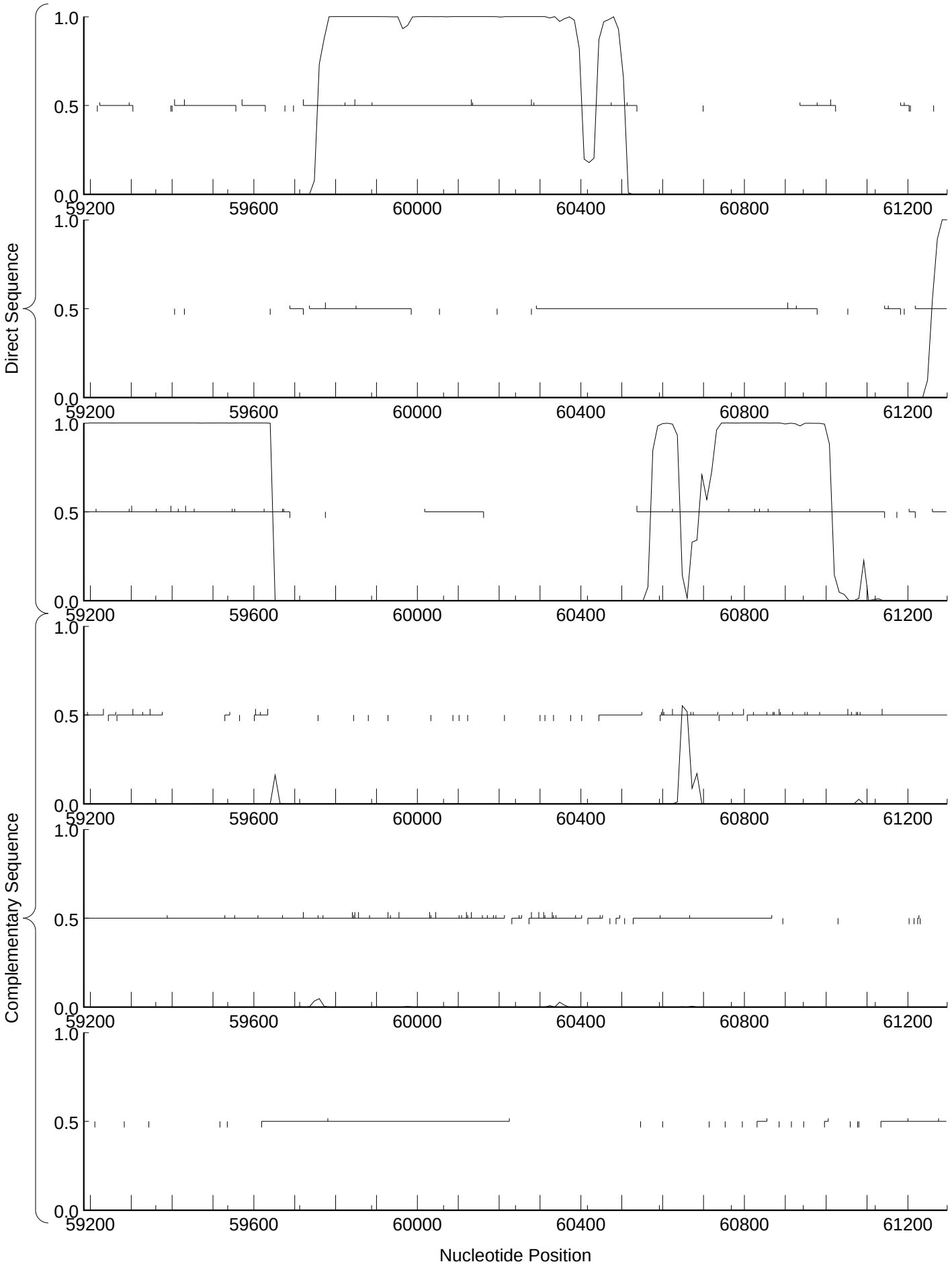
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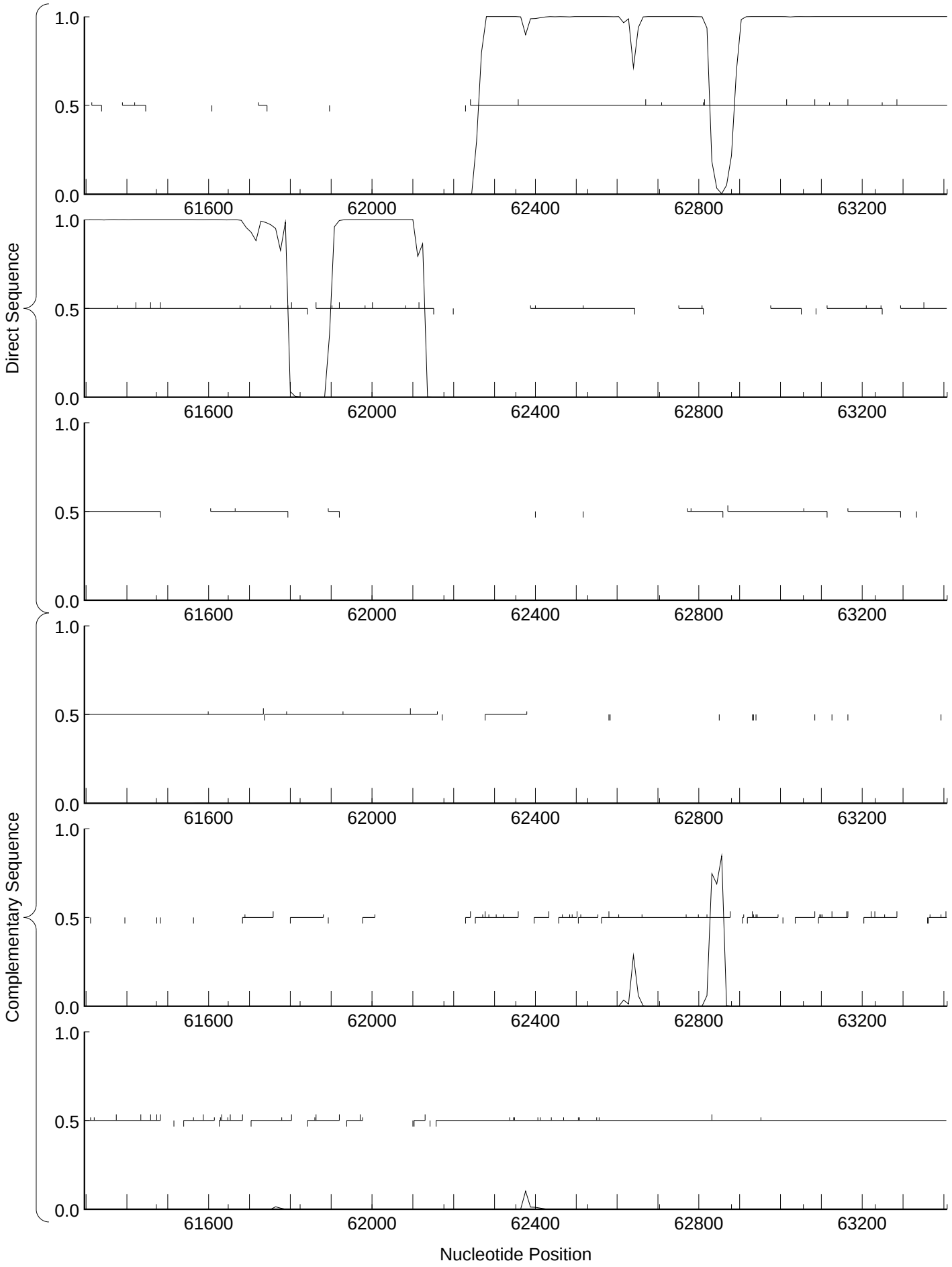
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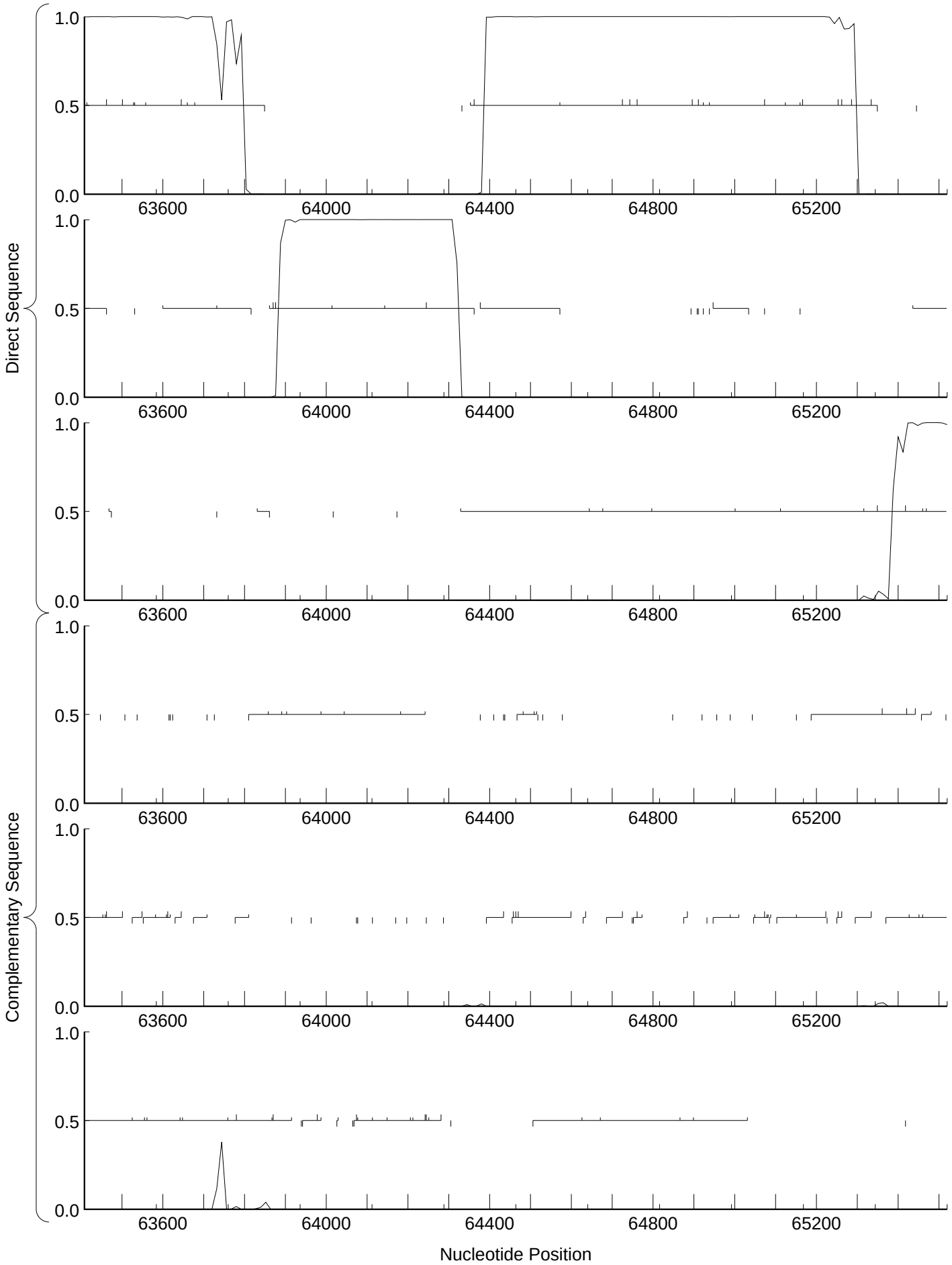
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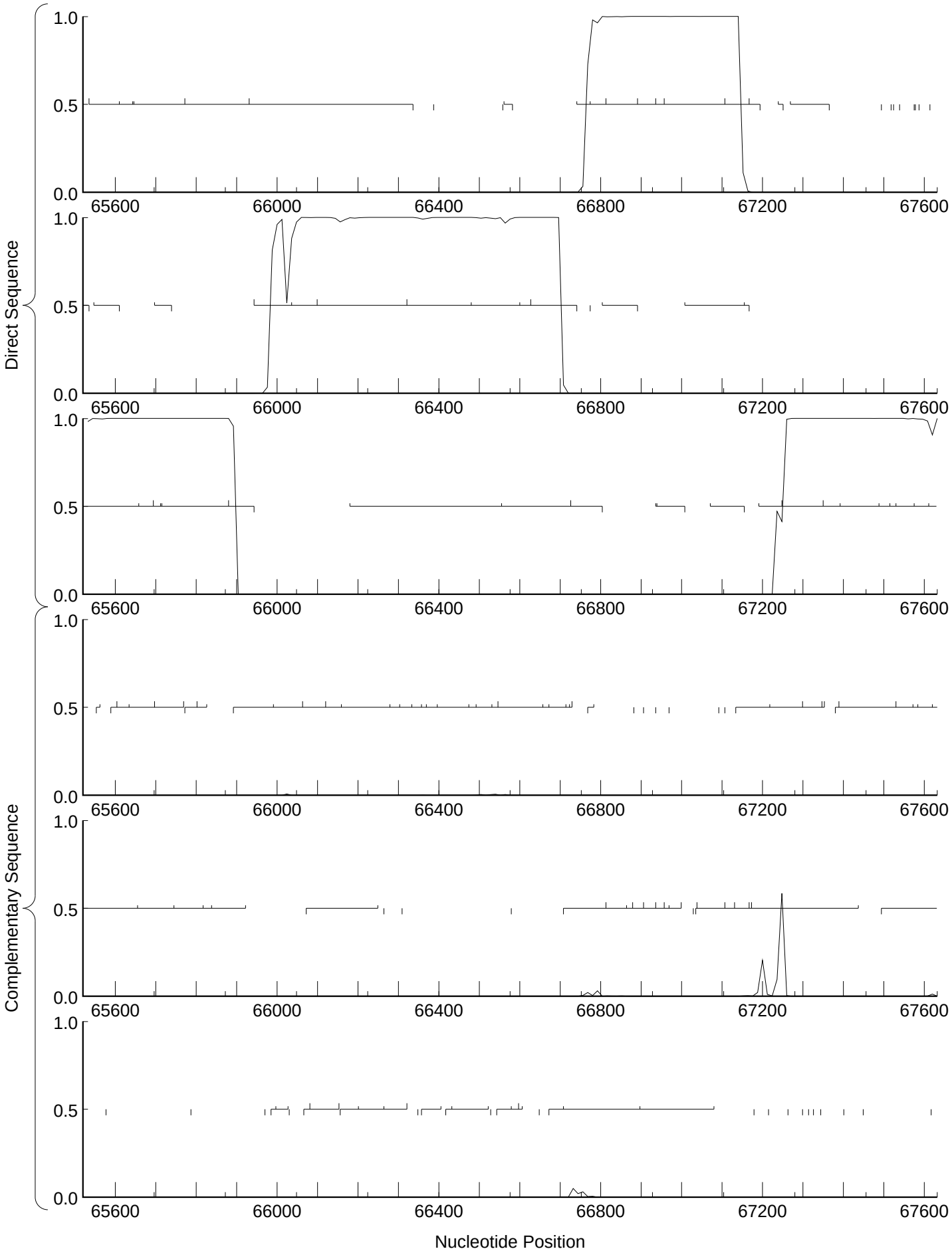
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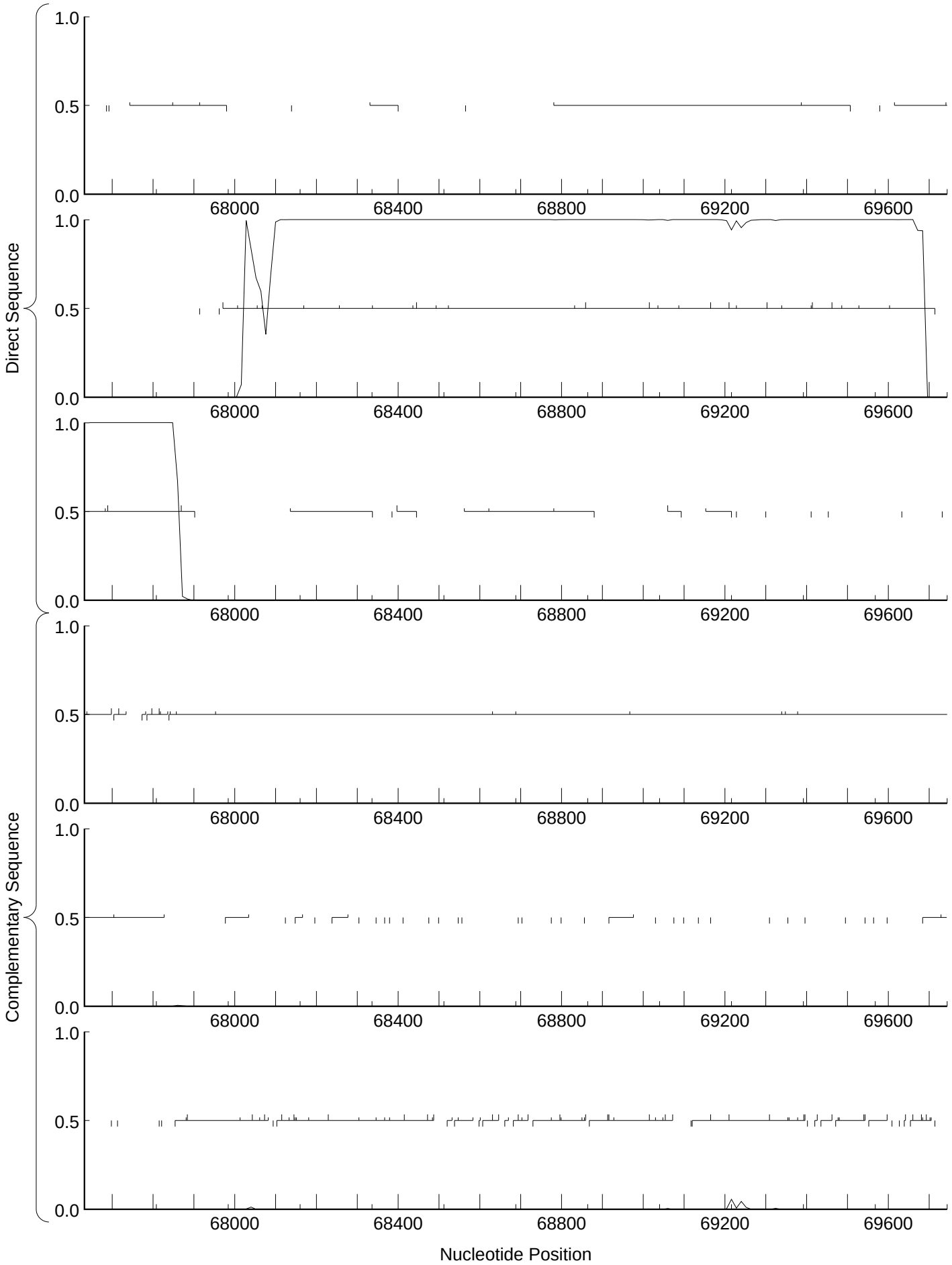
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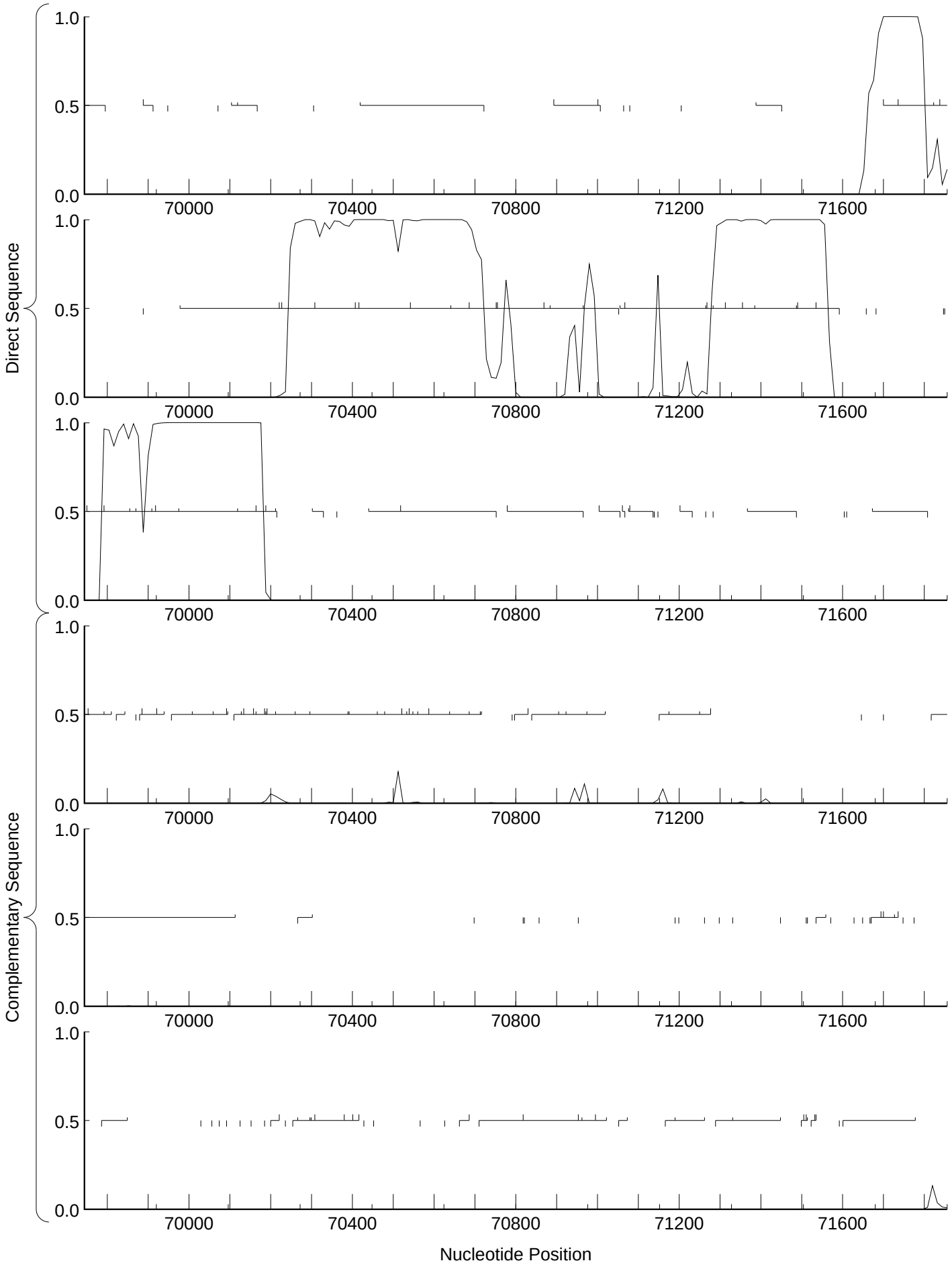
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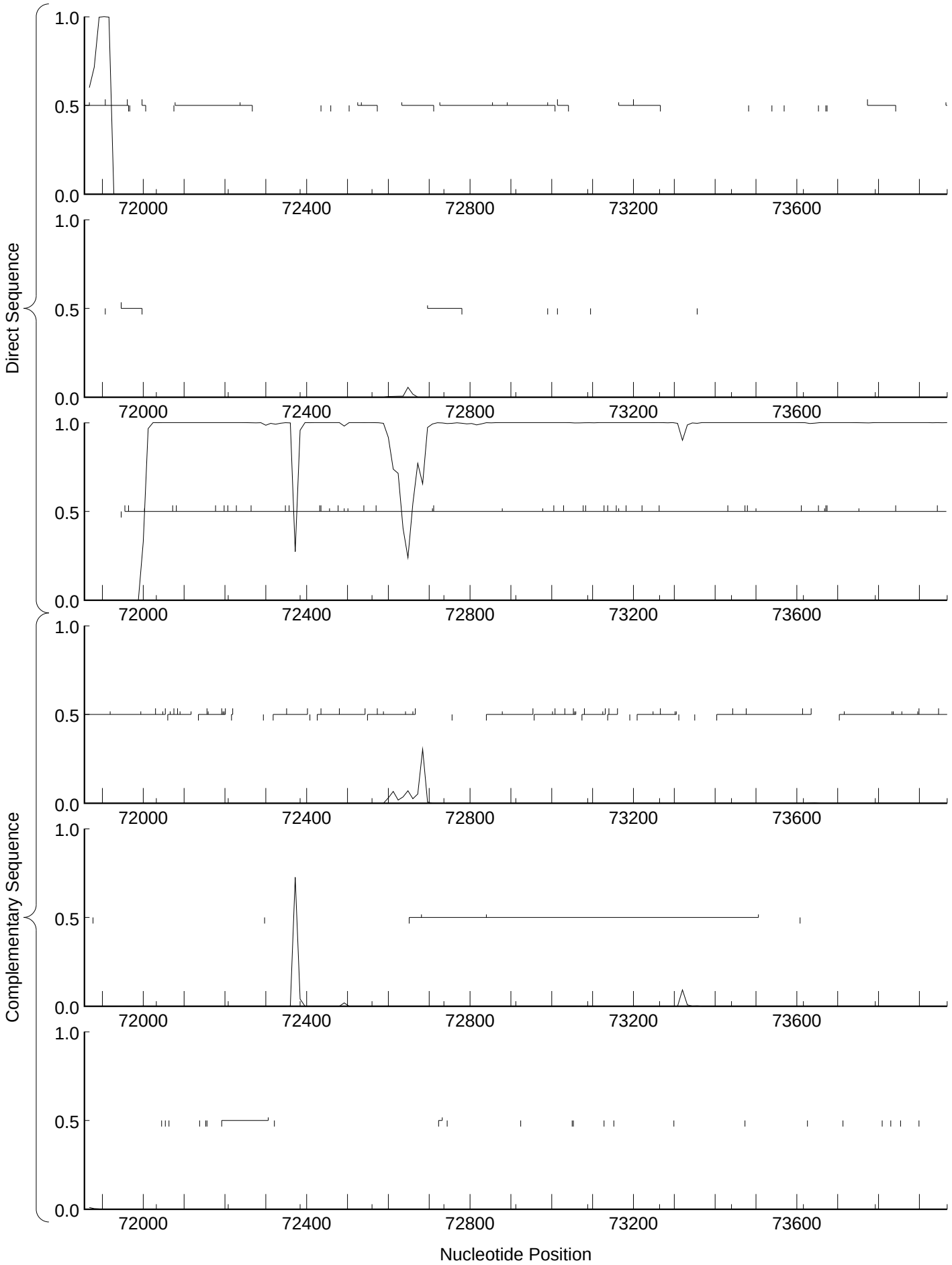
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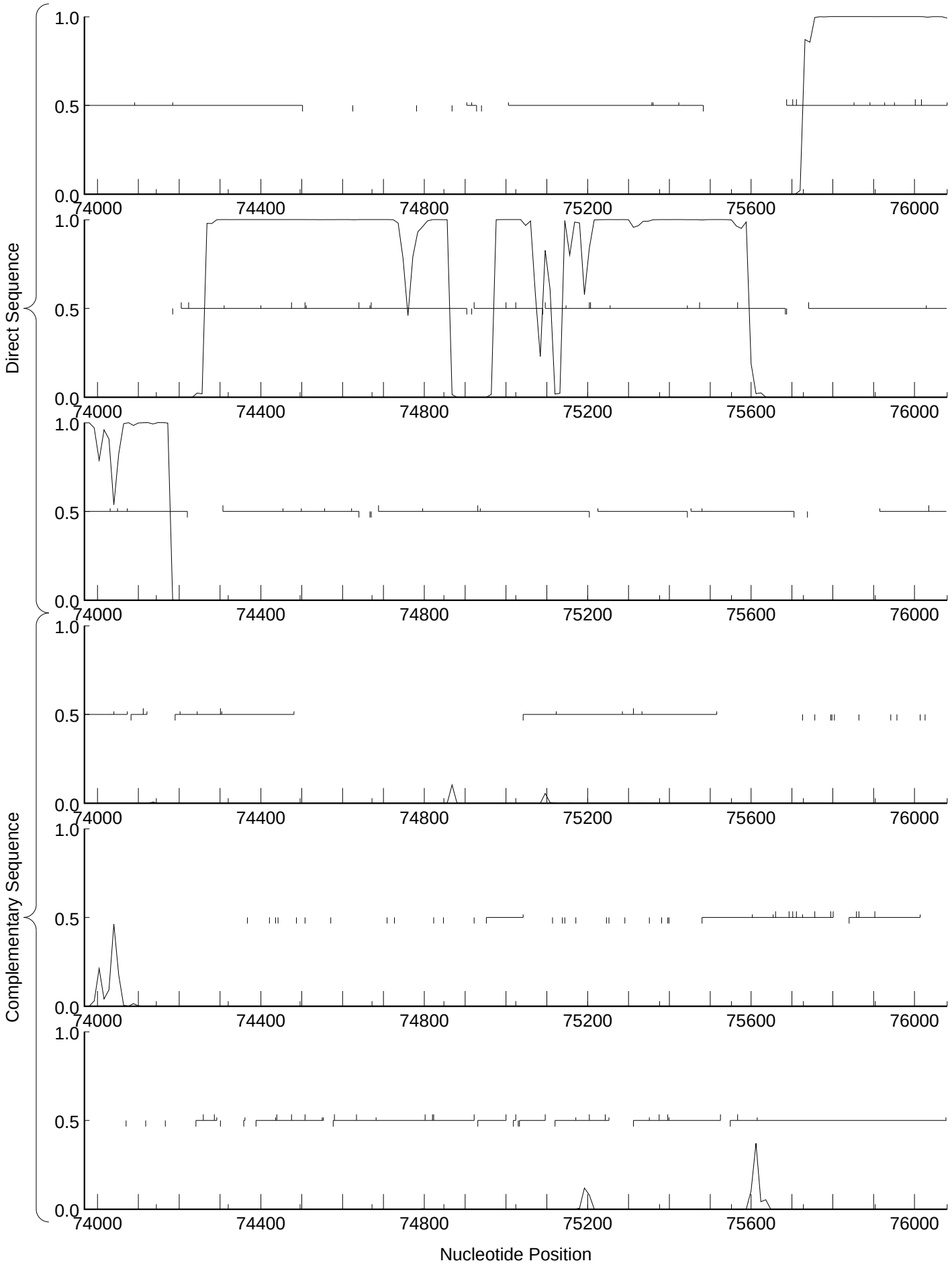
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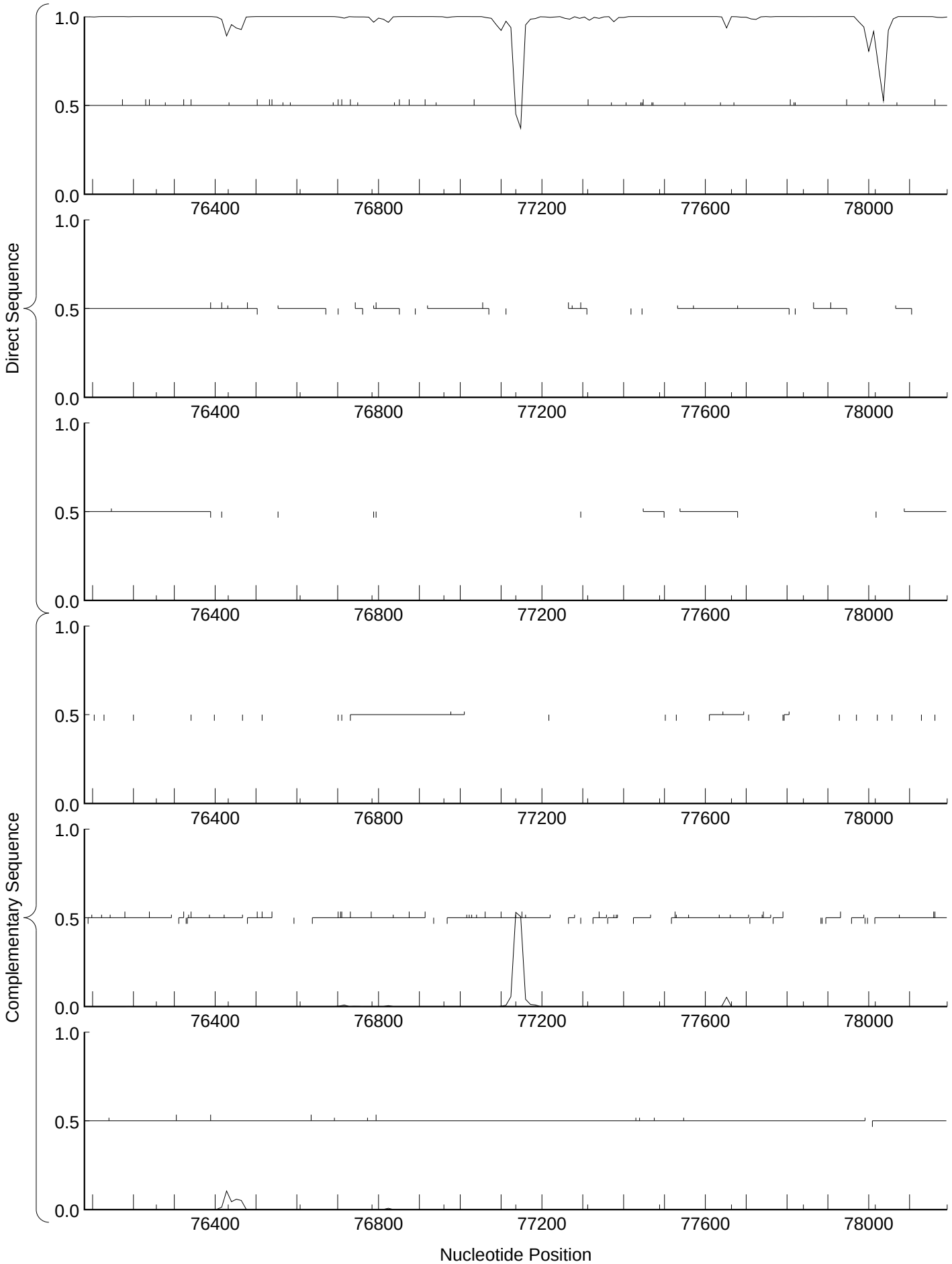
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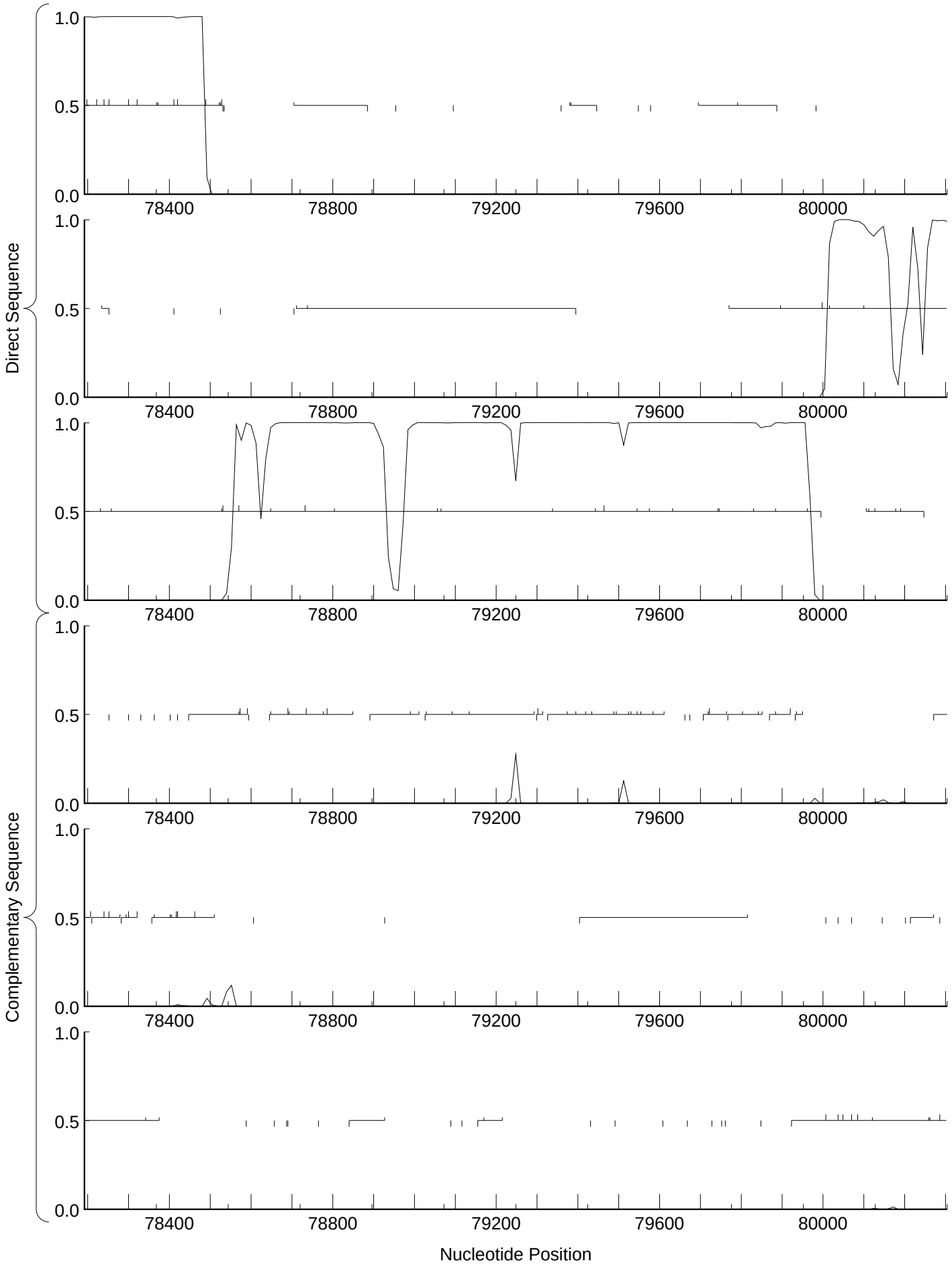
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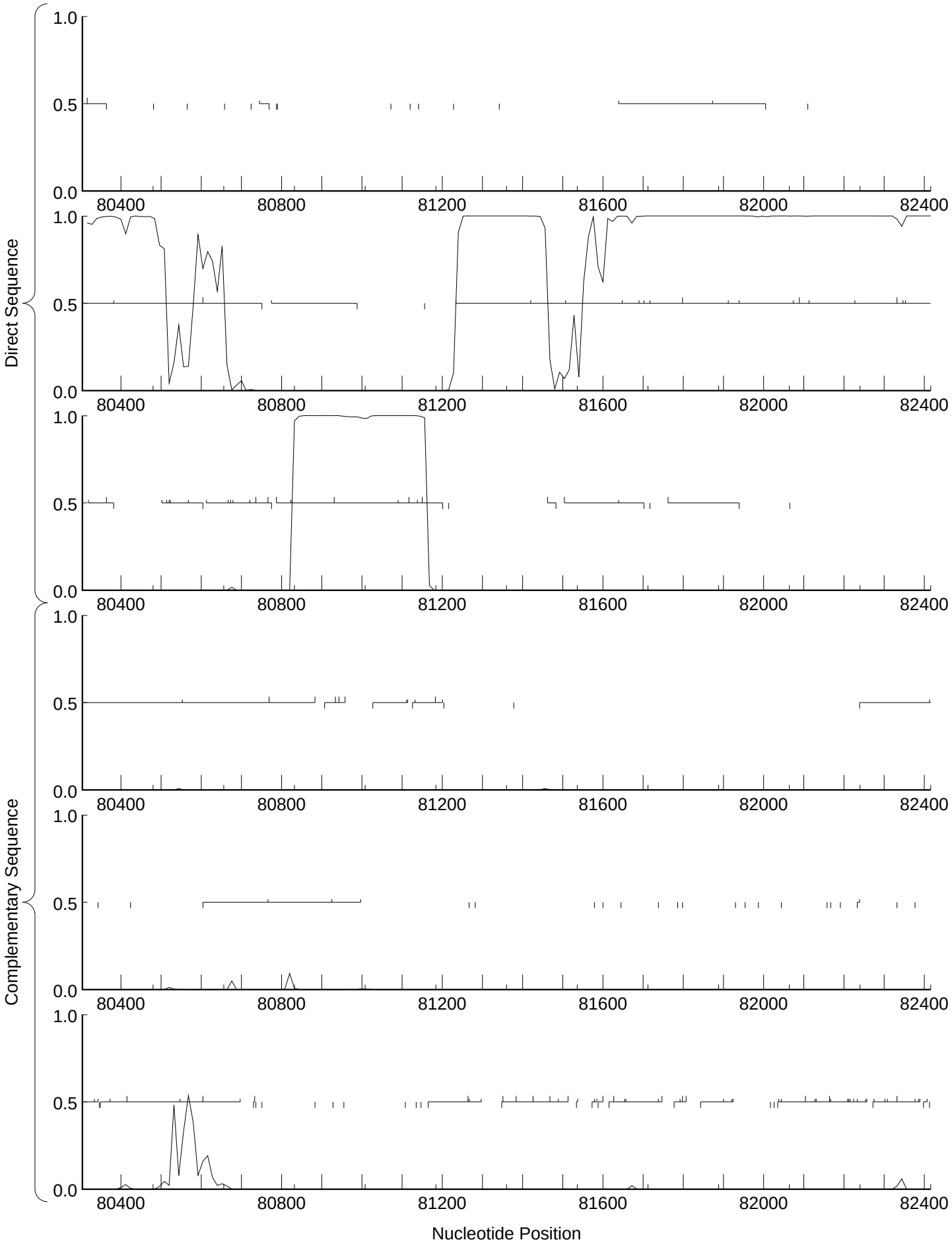
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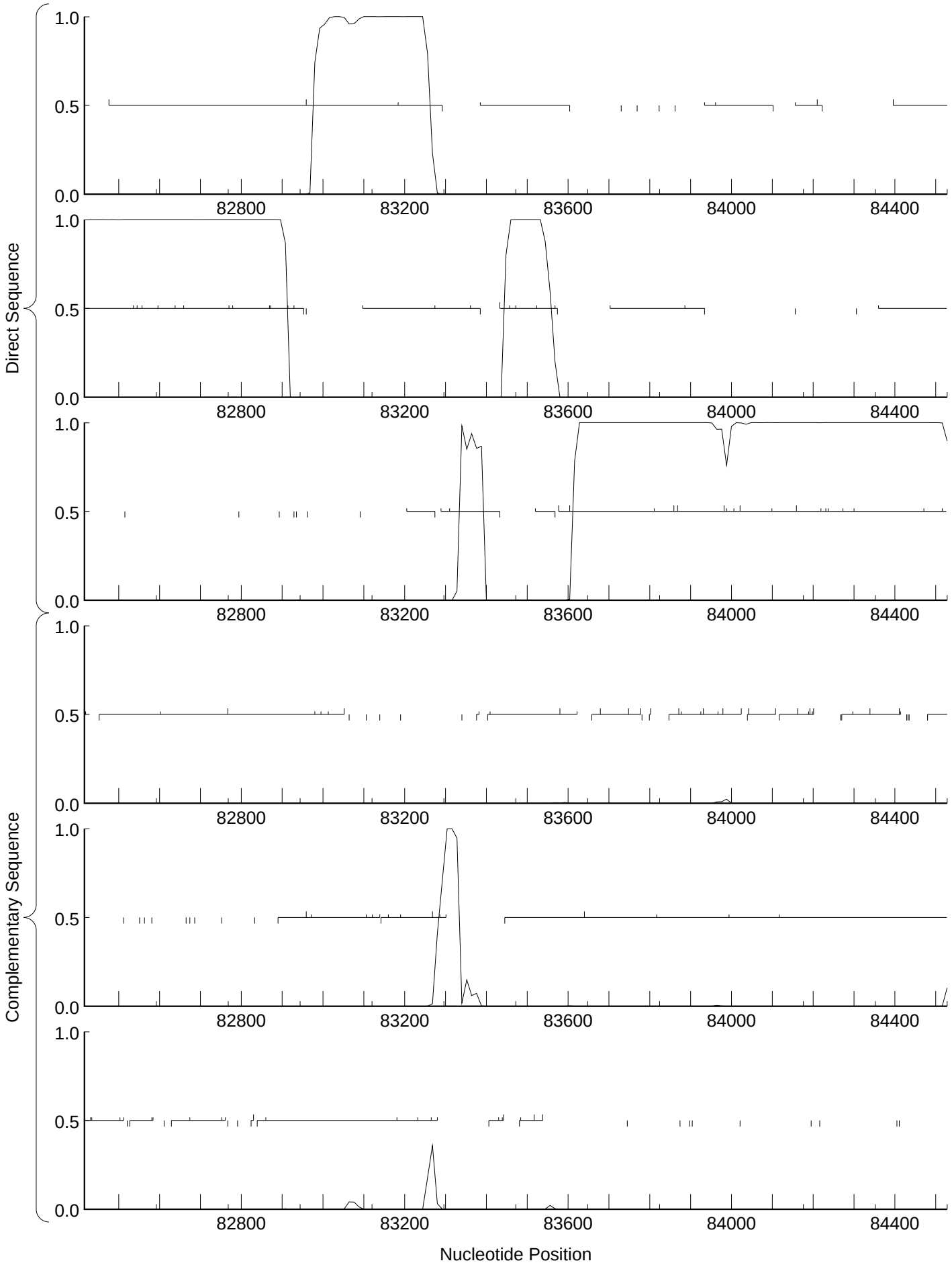
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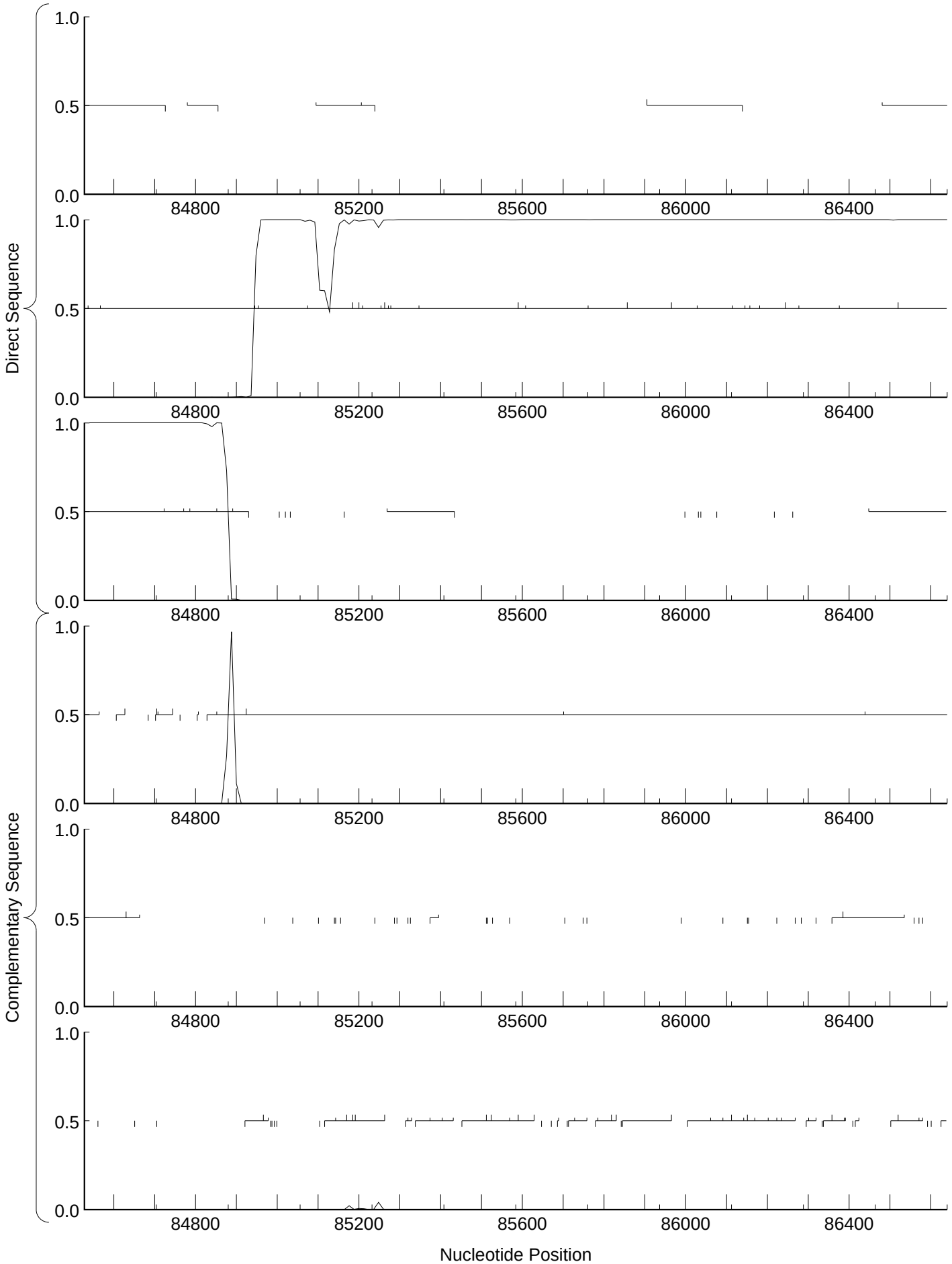
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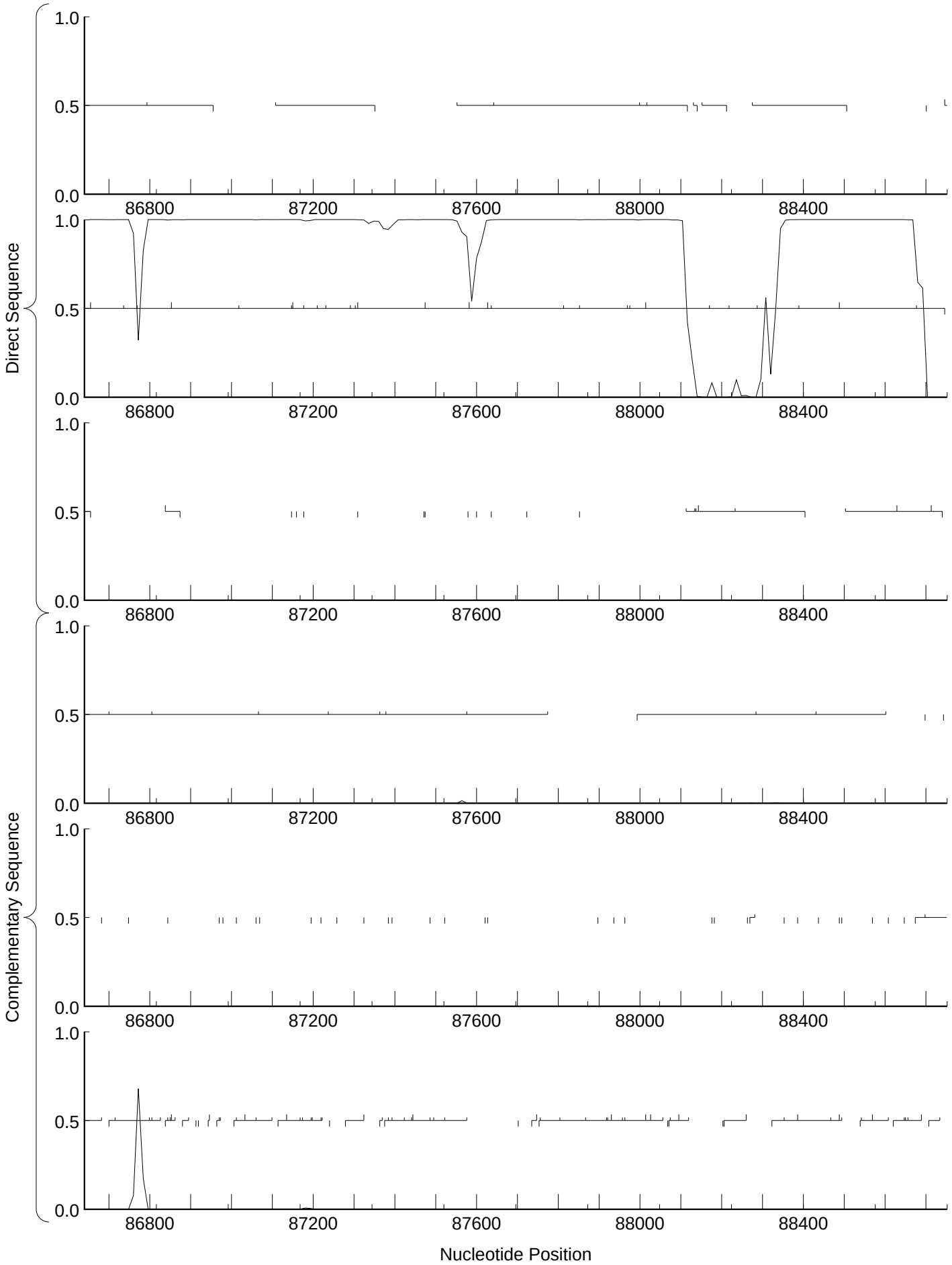
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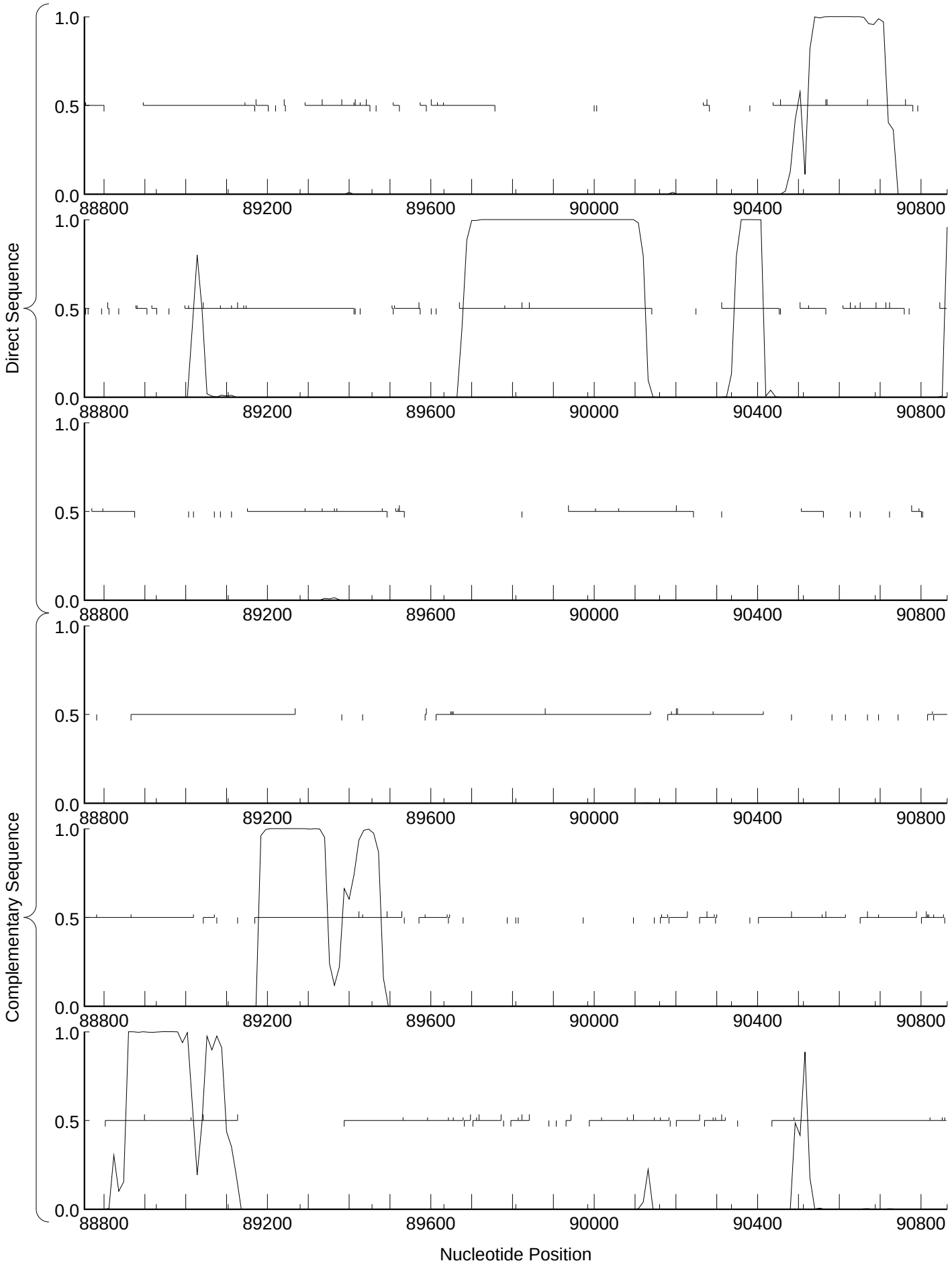
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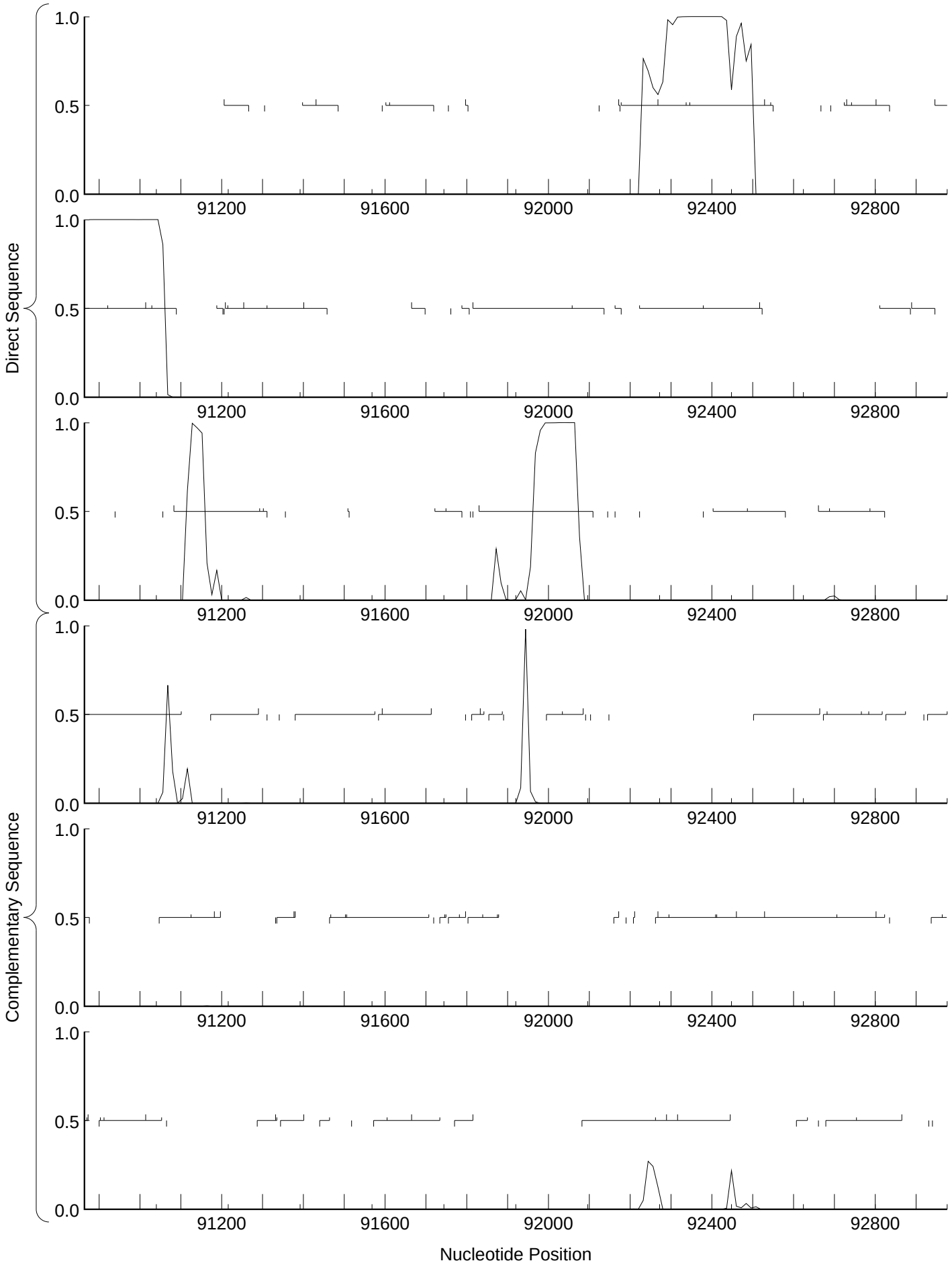
ycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 43/75



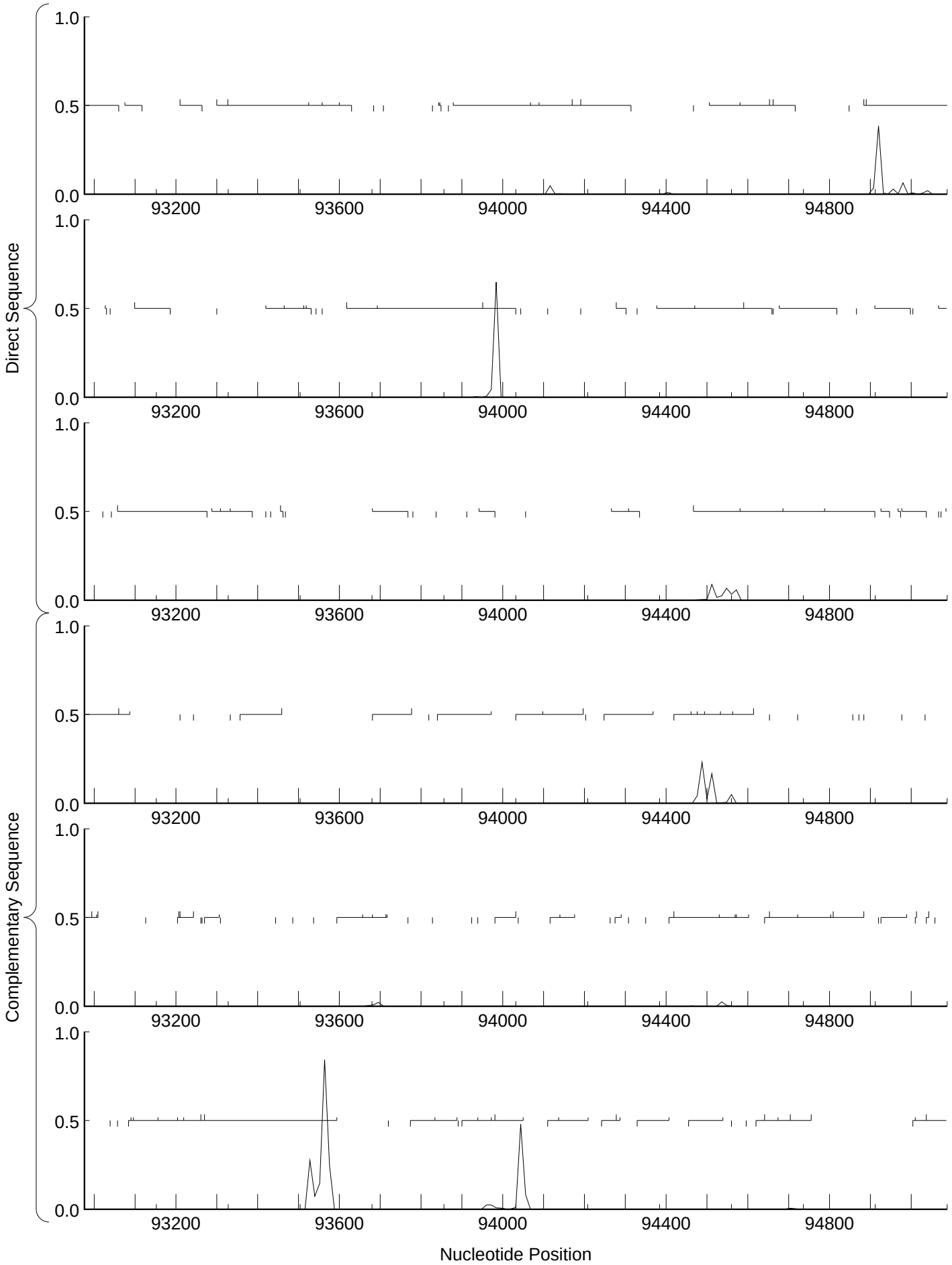
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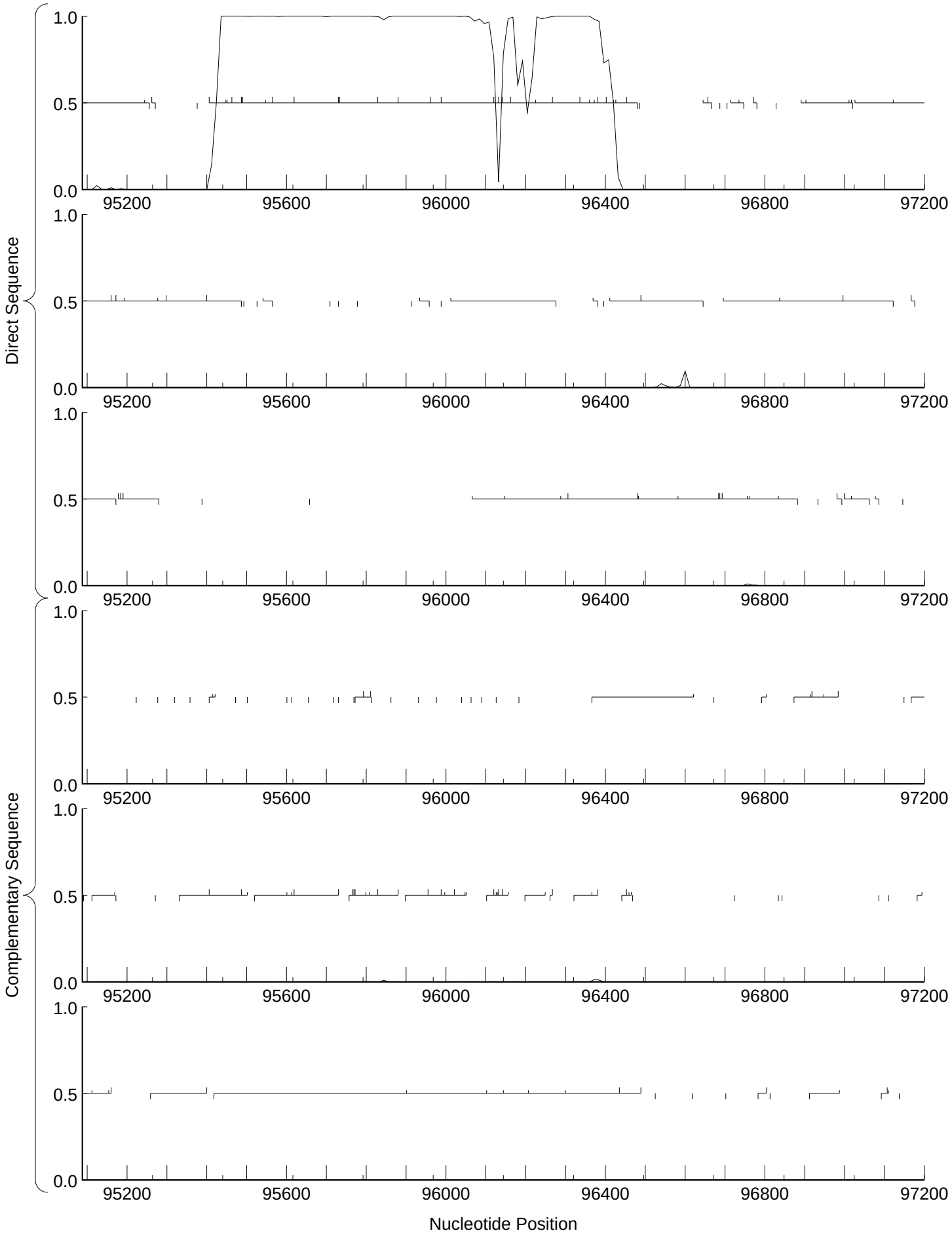
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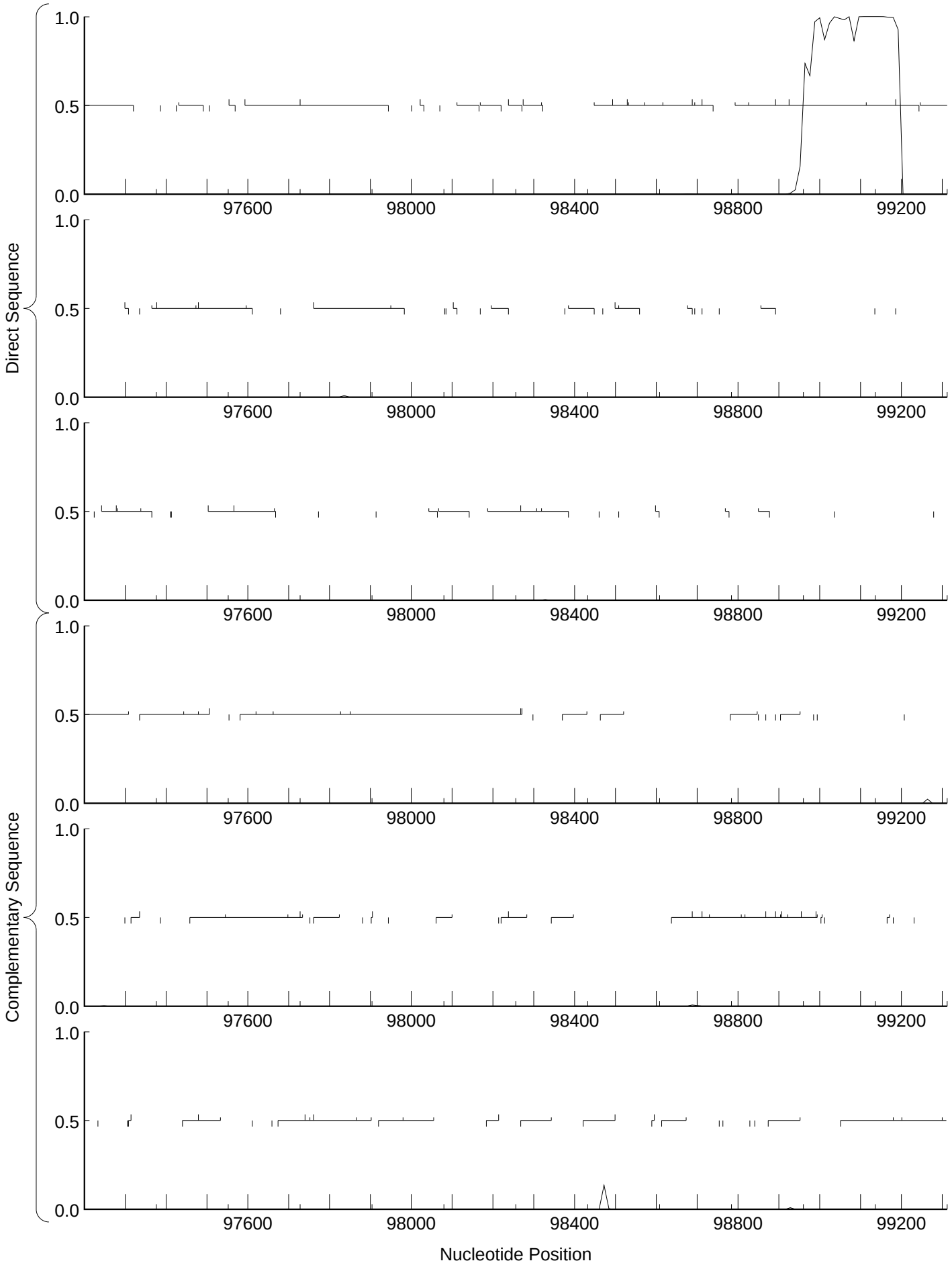
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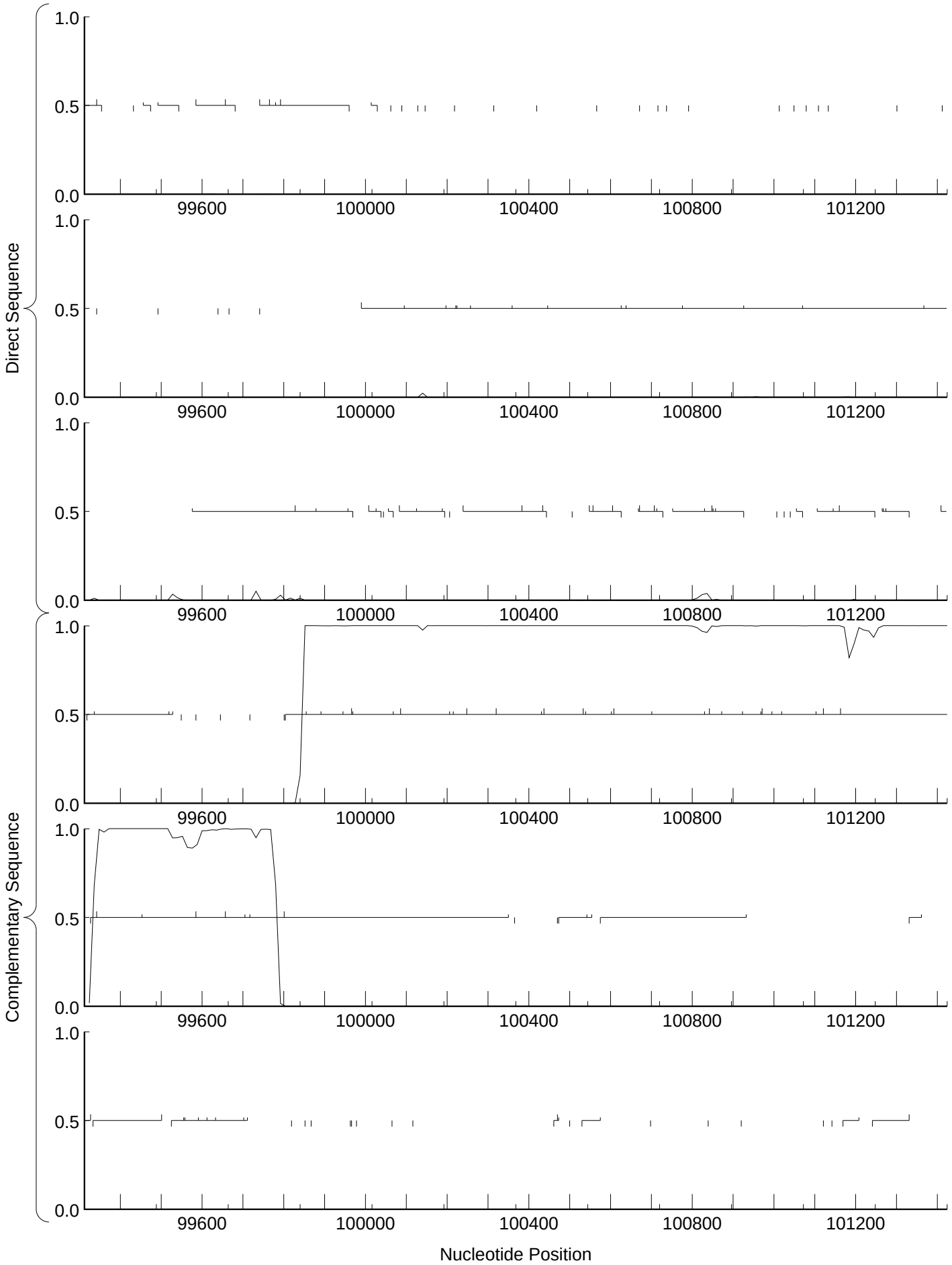
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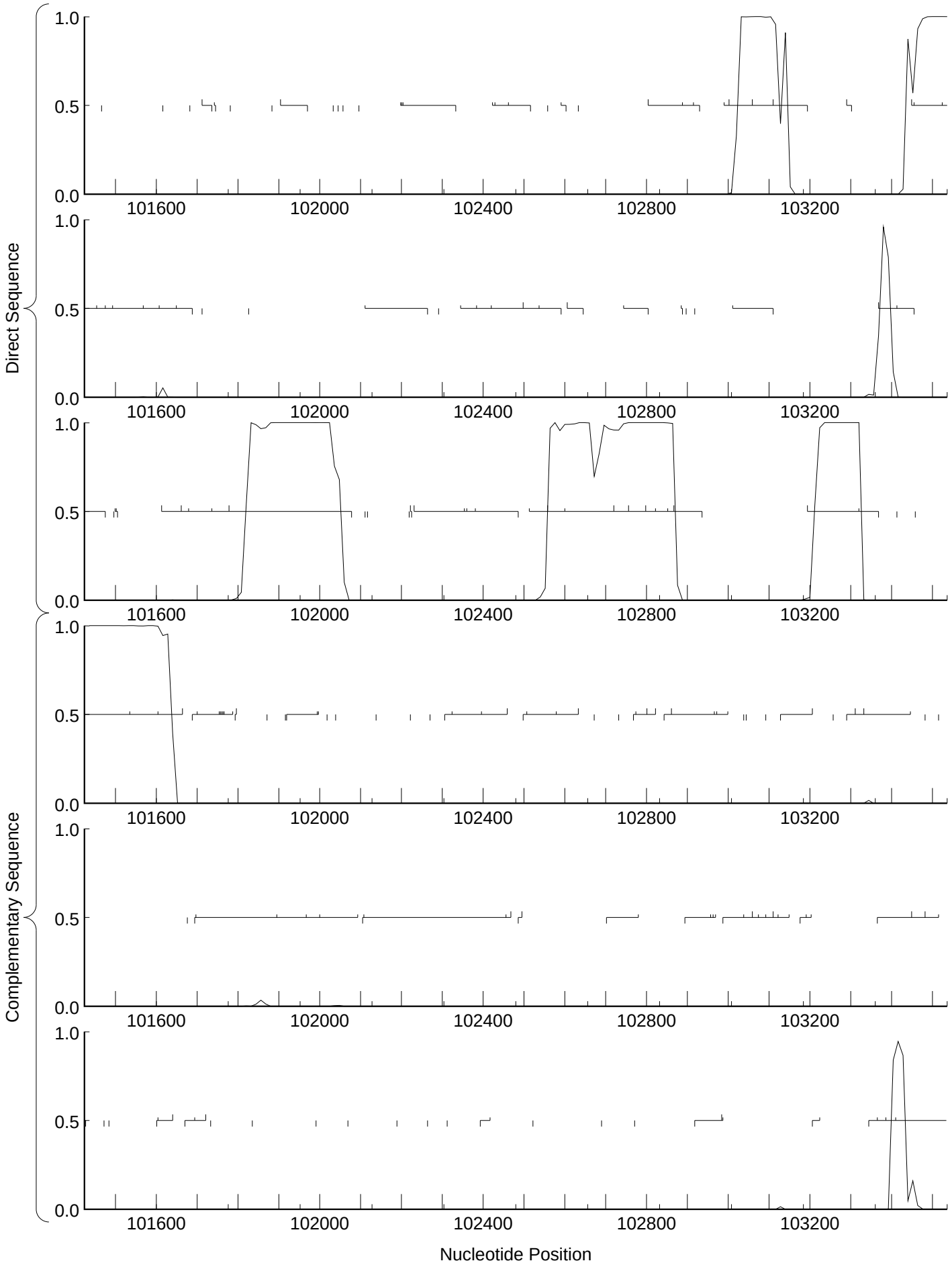
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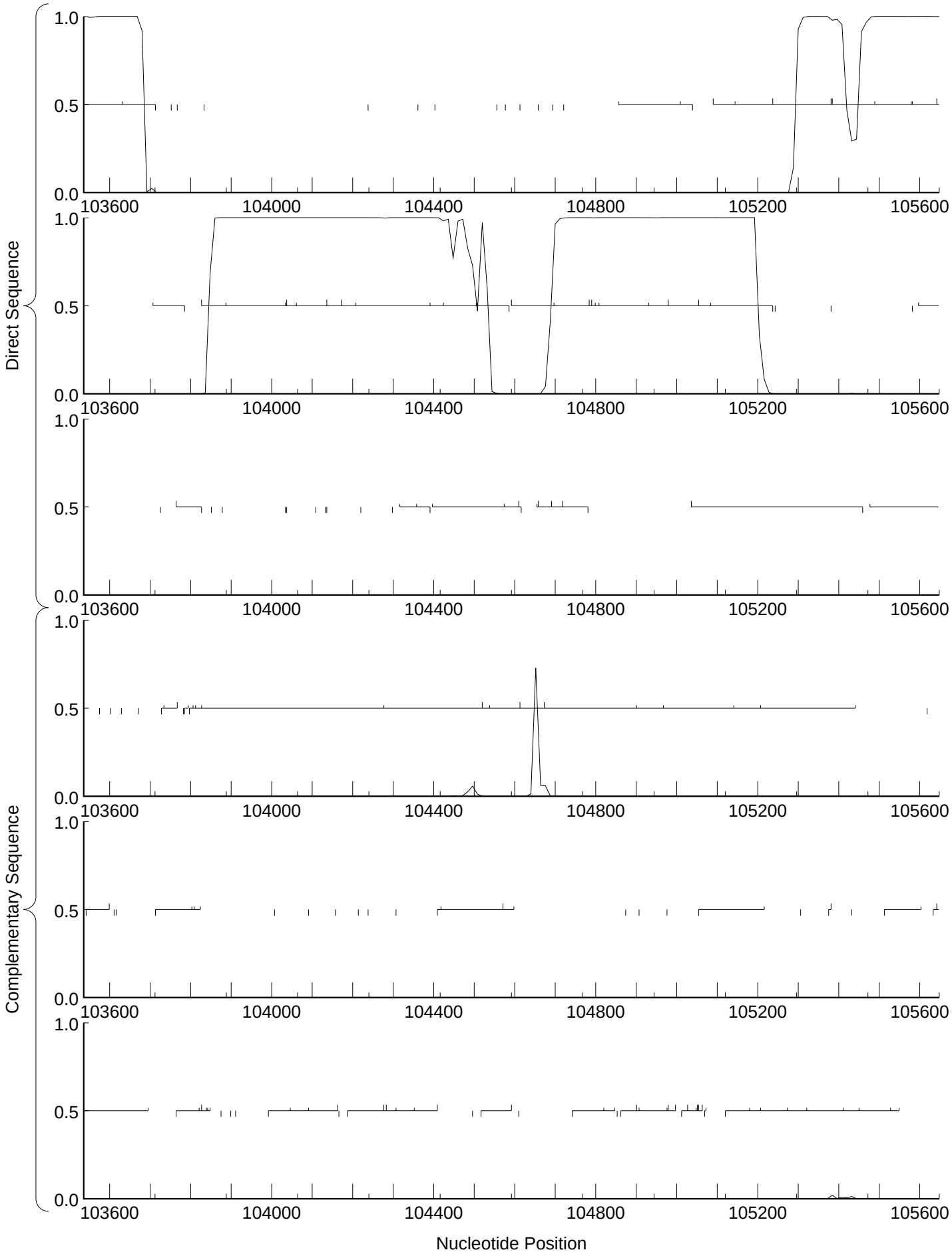
ycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 49/75



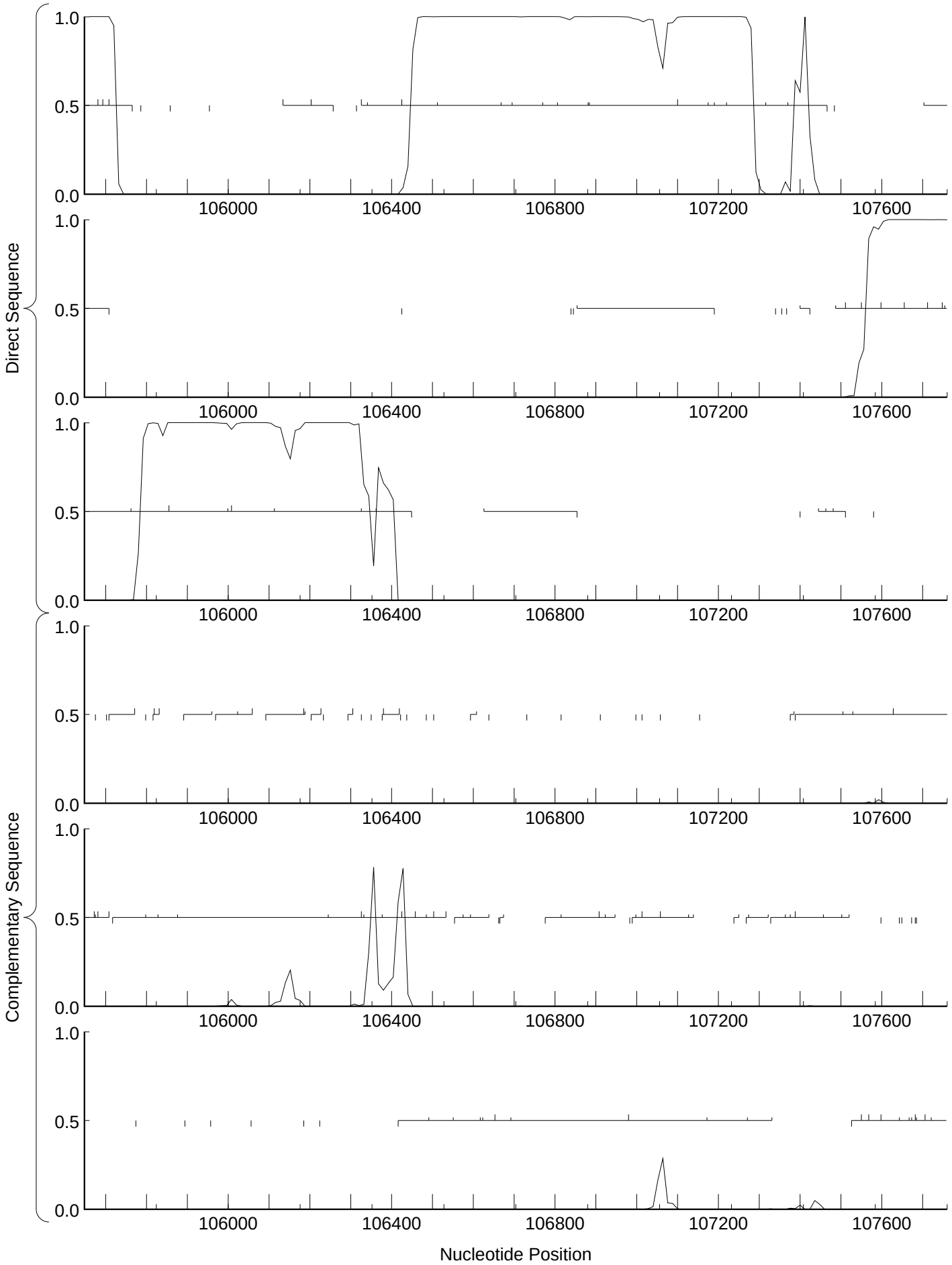
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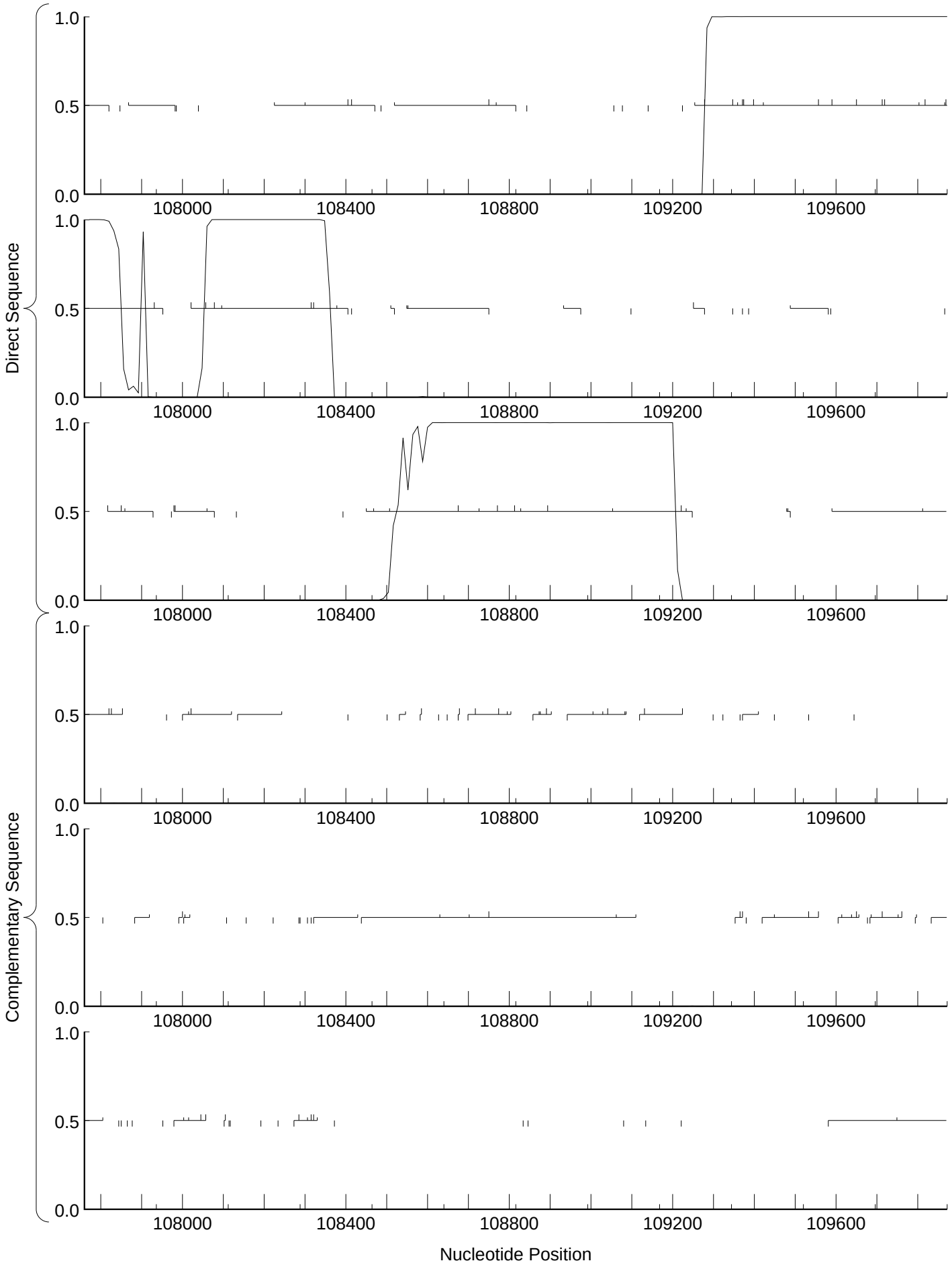
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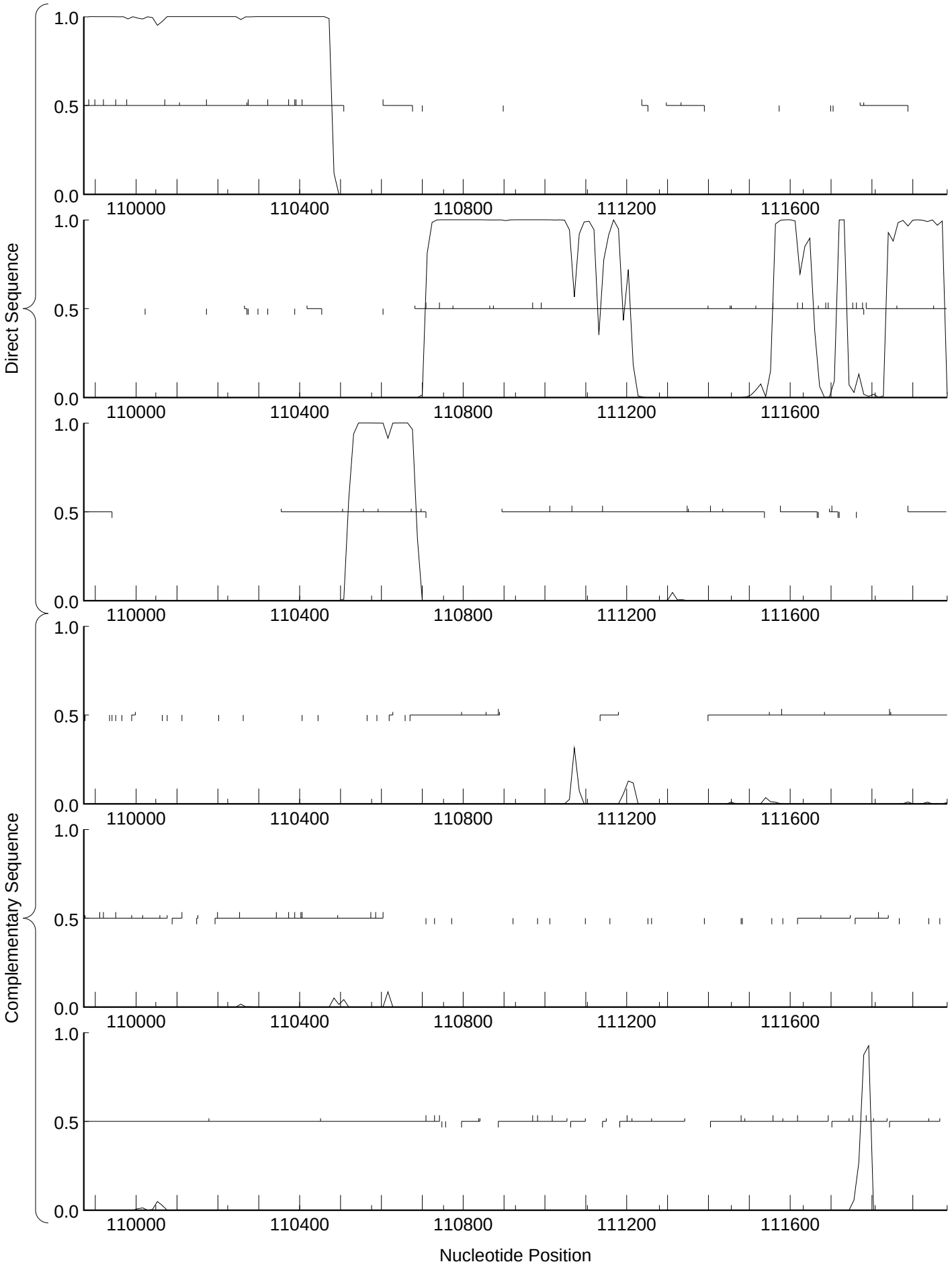
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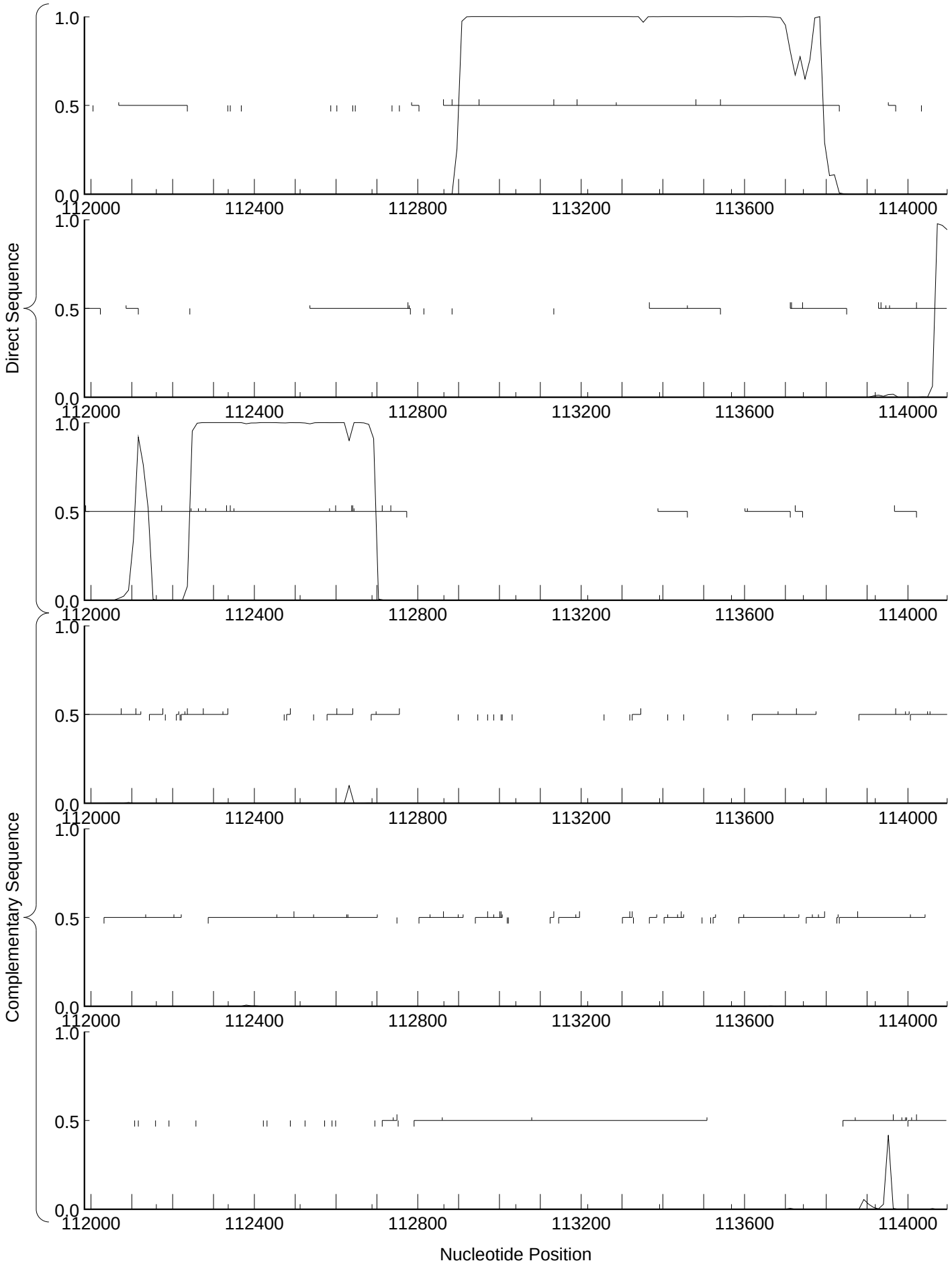
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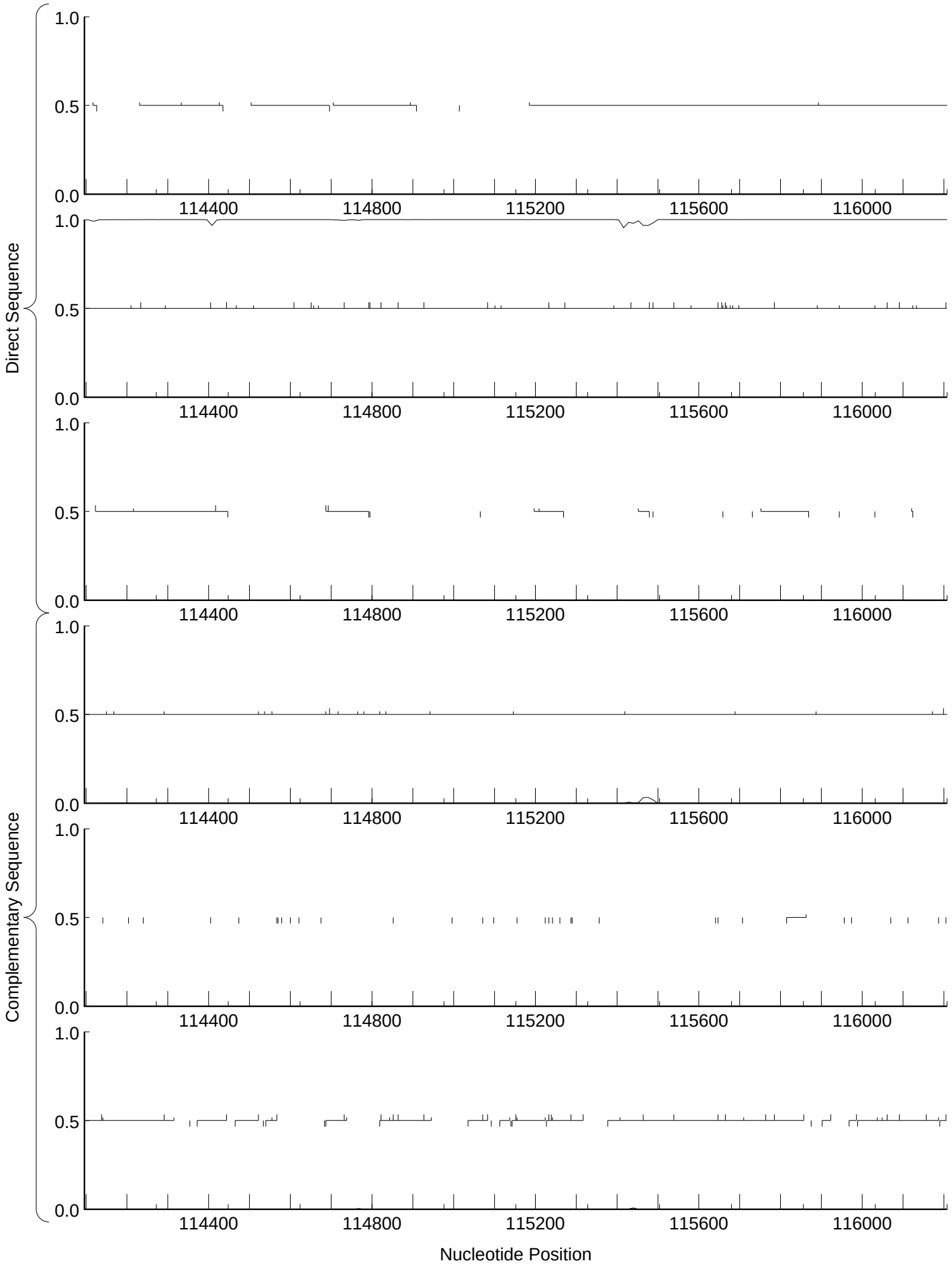
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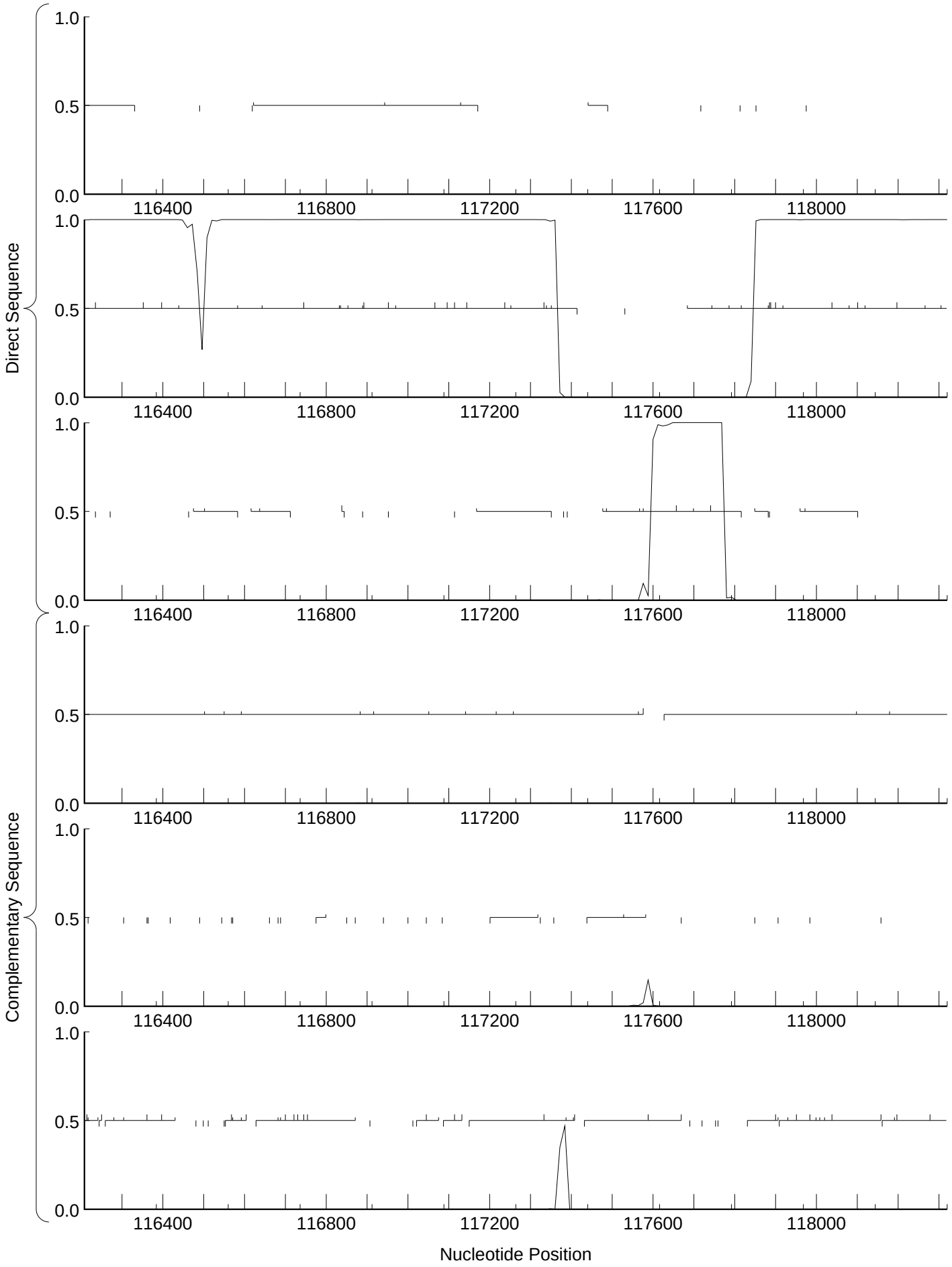
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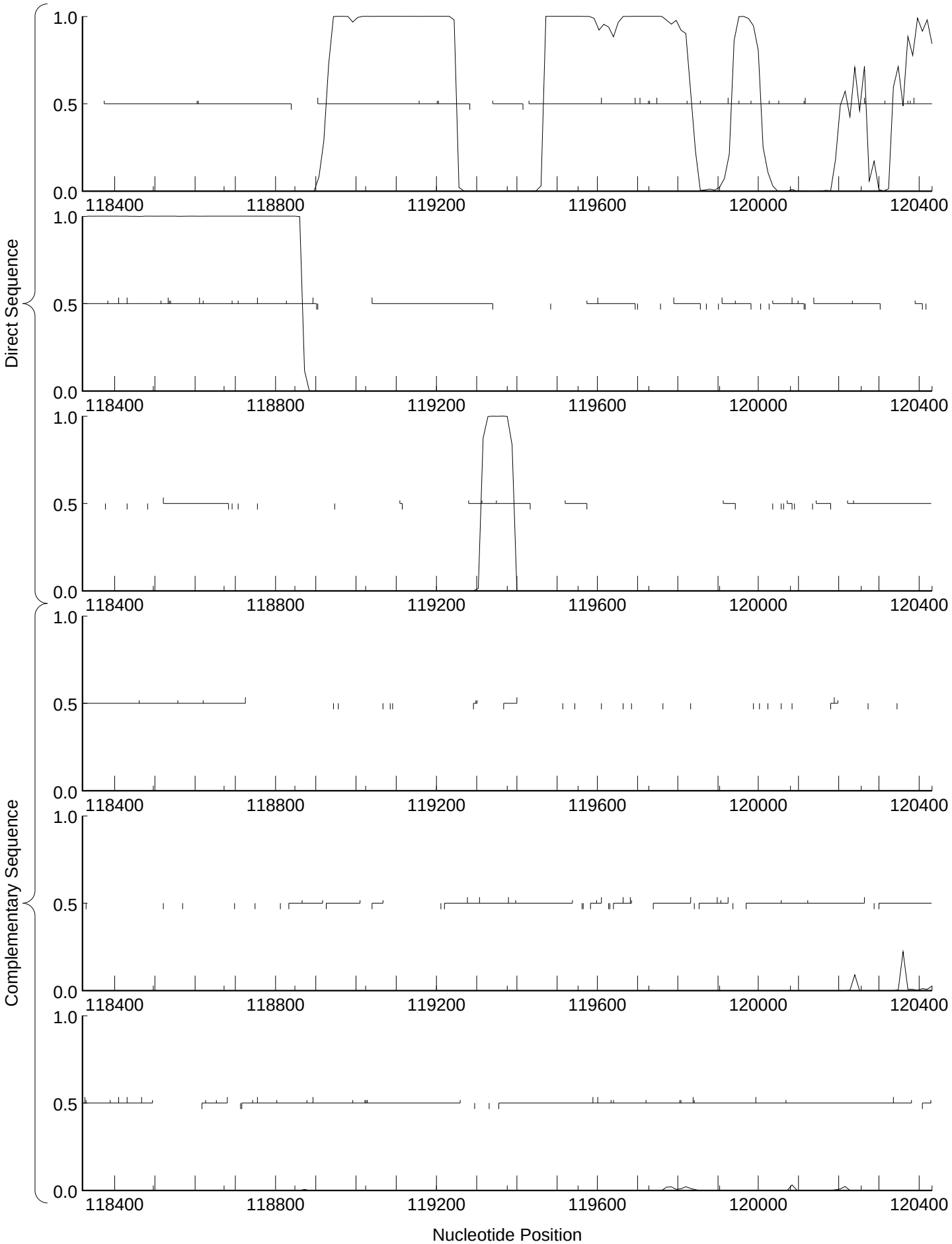
ycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 56/75



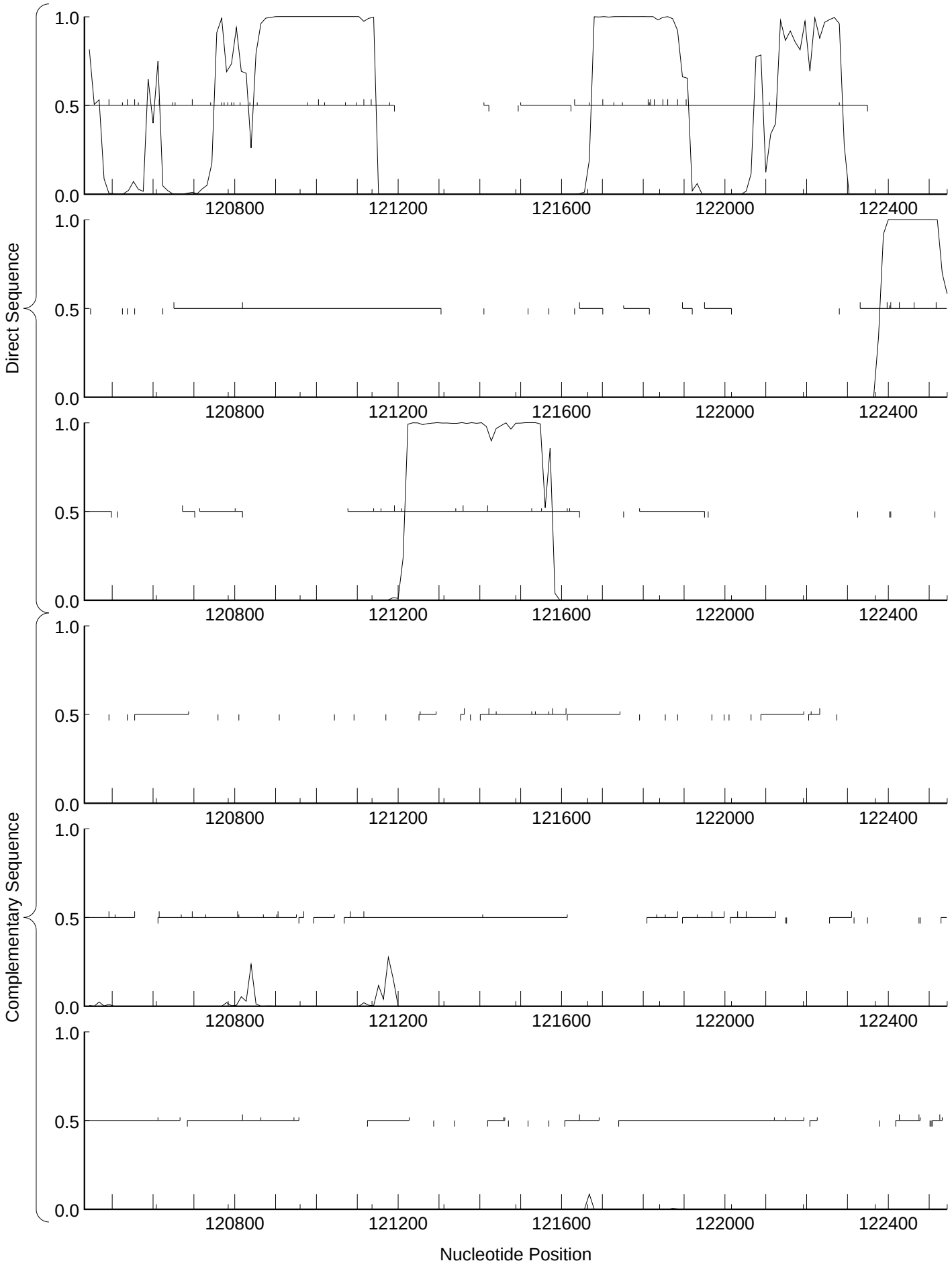
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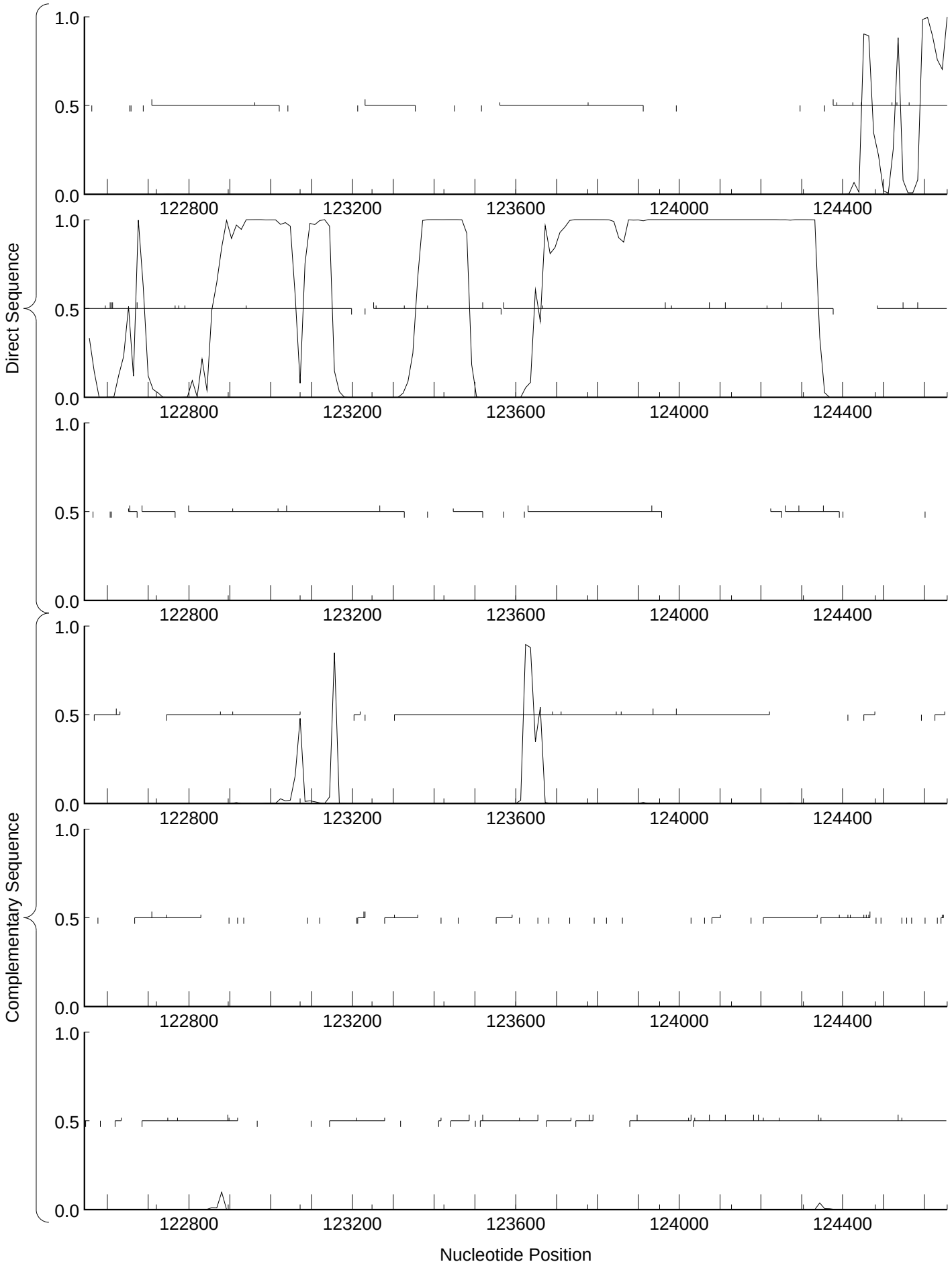
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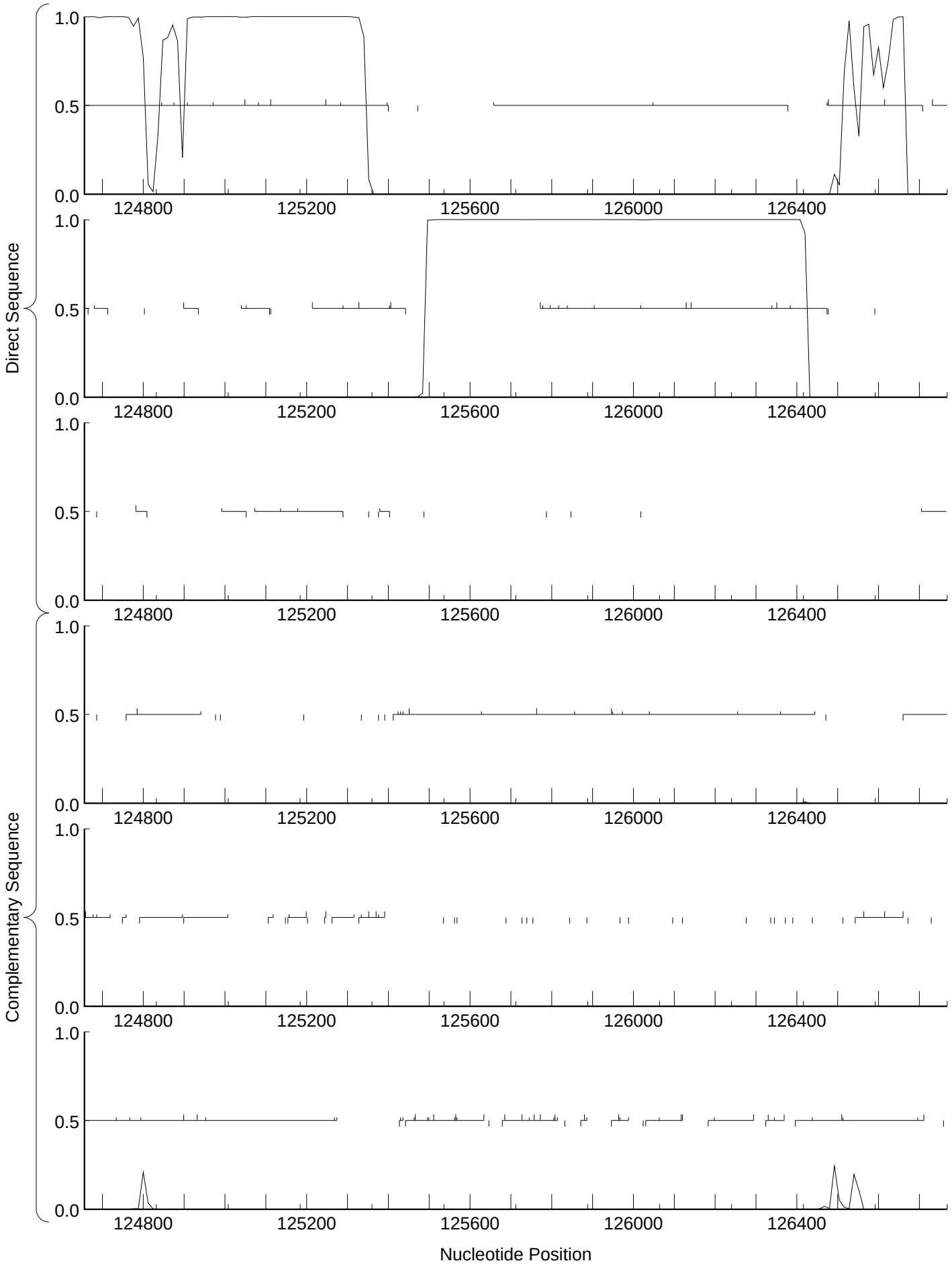
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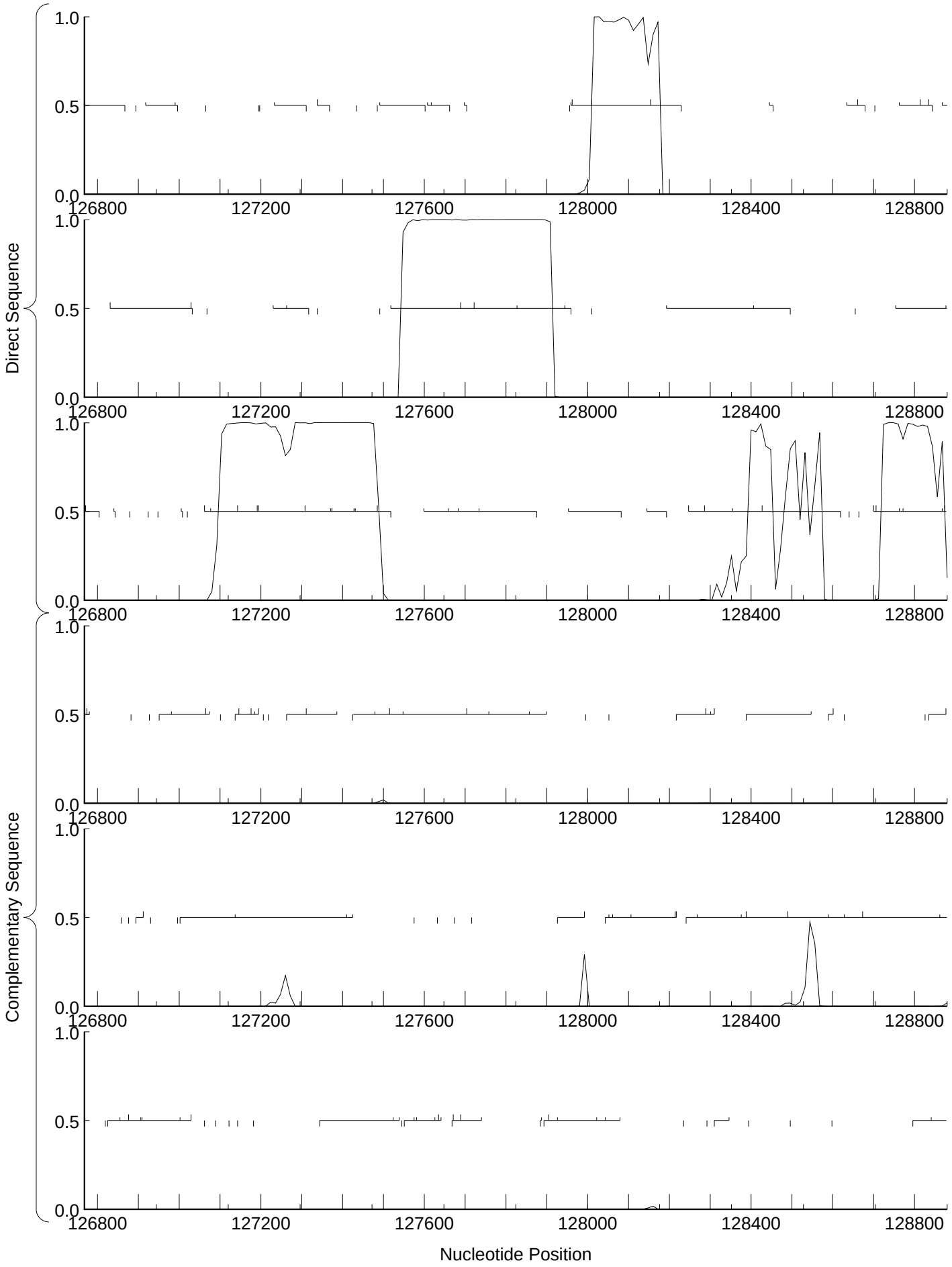
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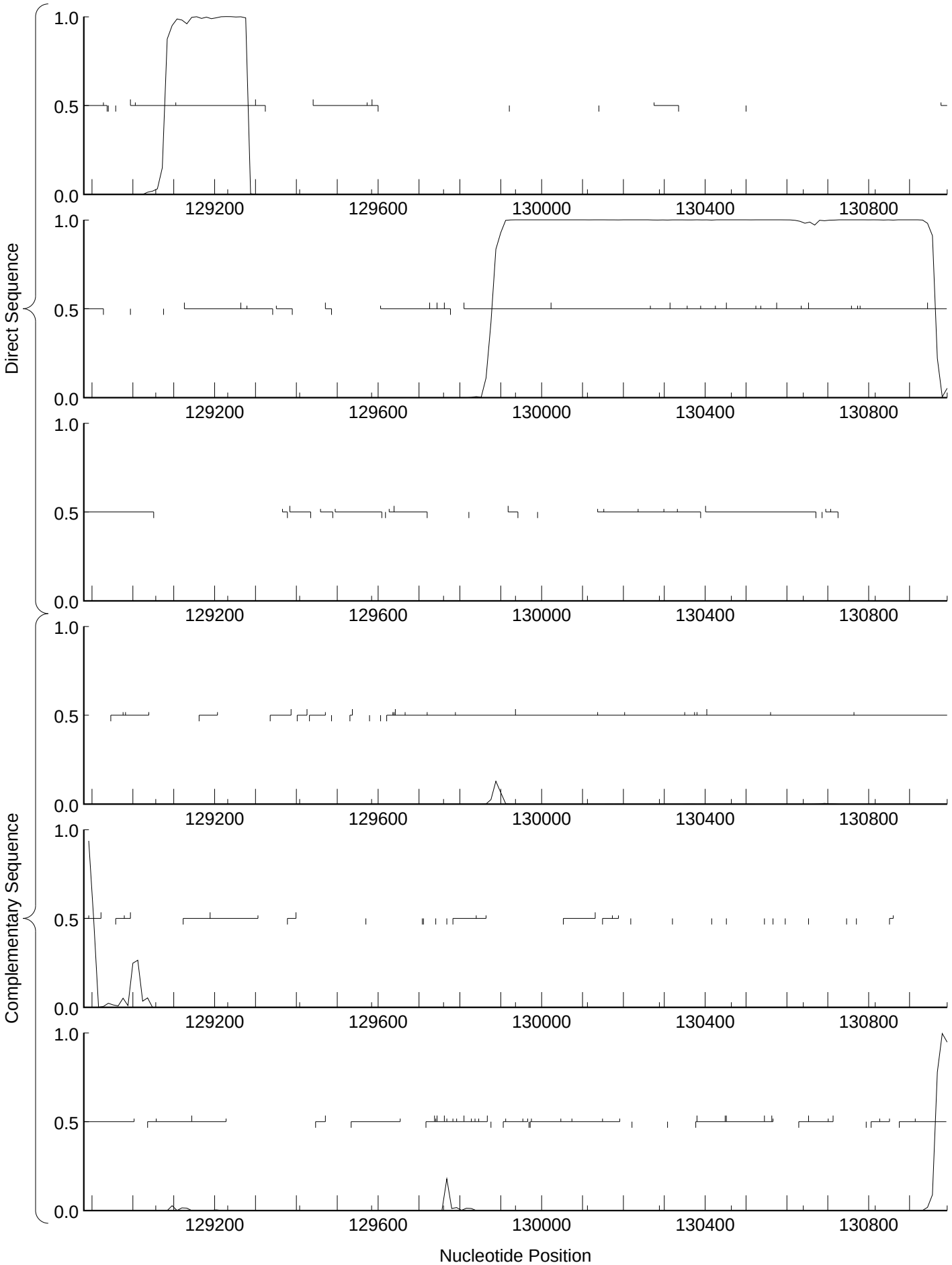
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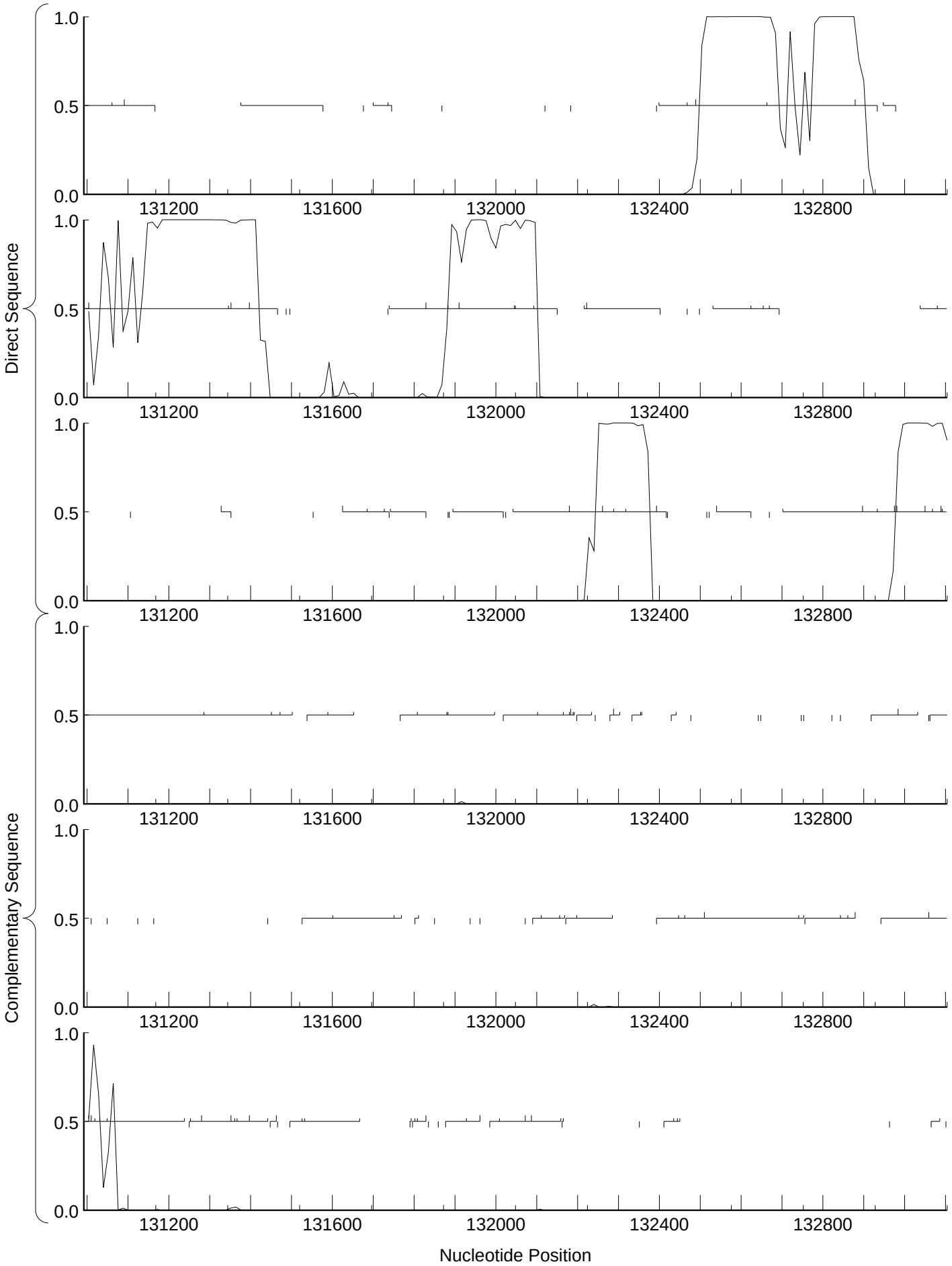
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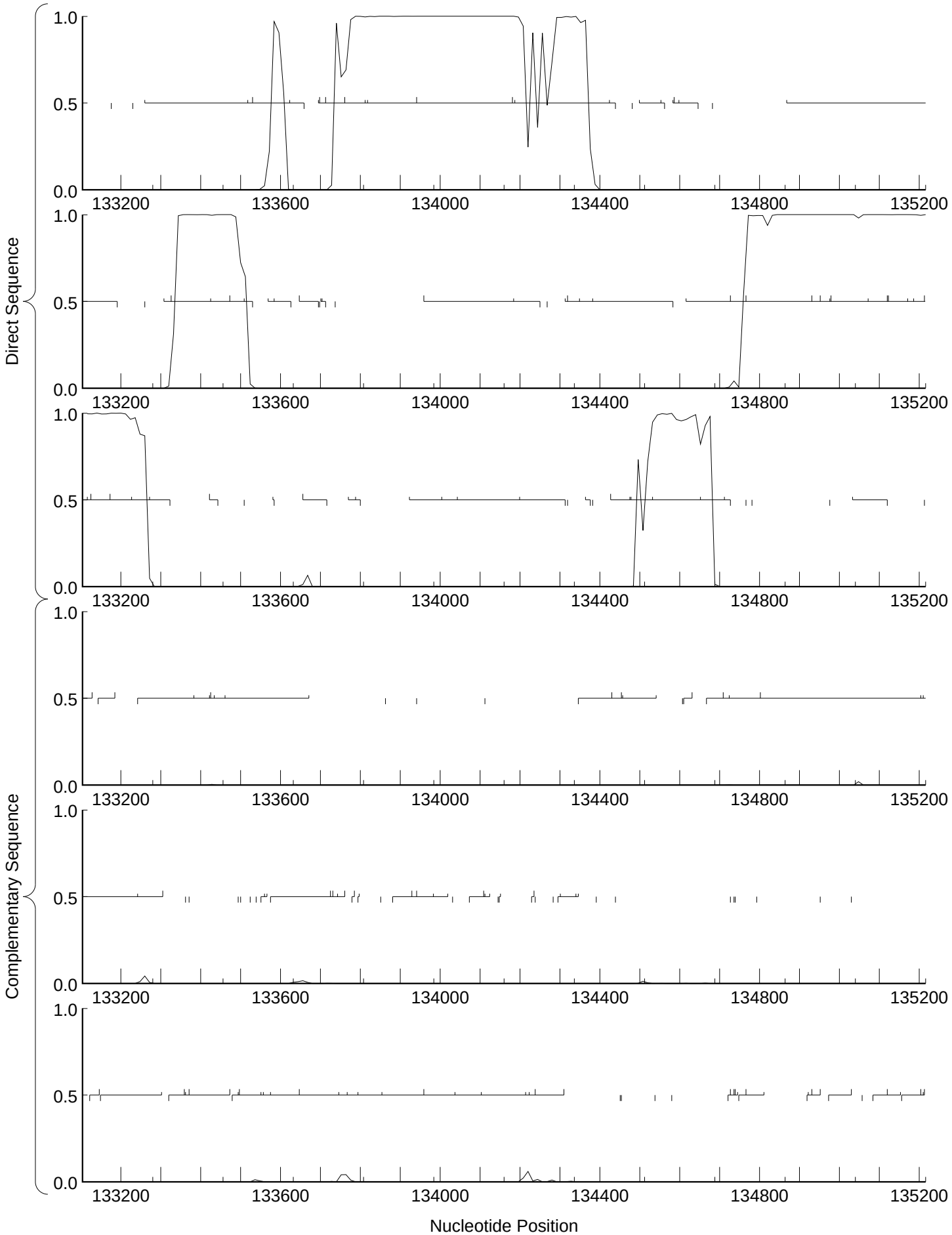
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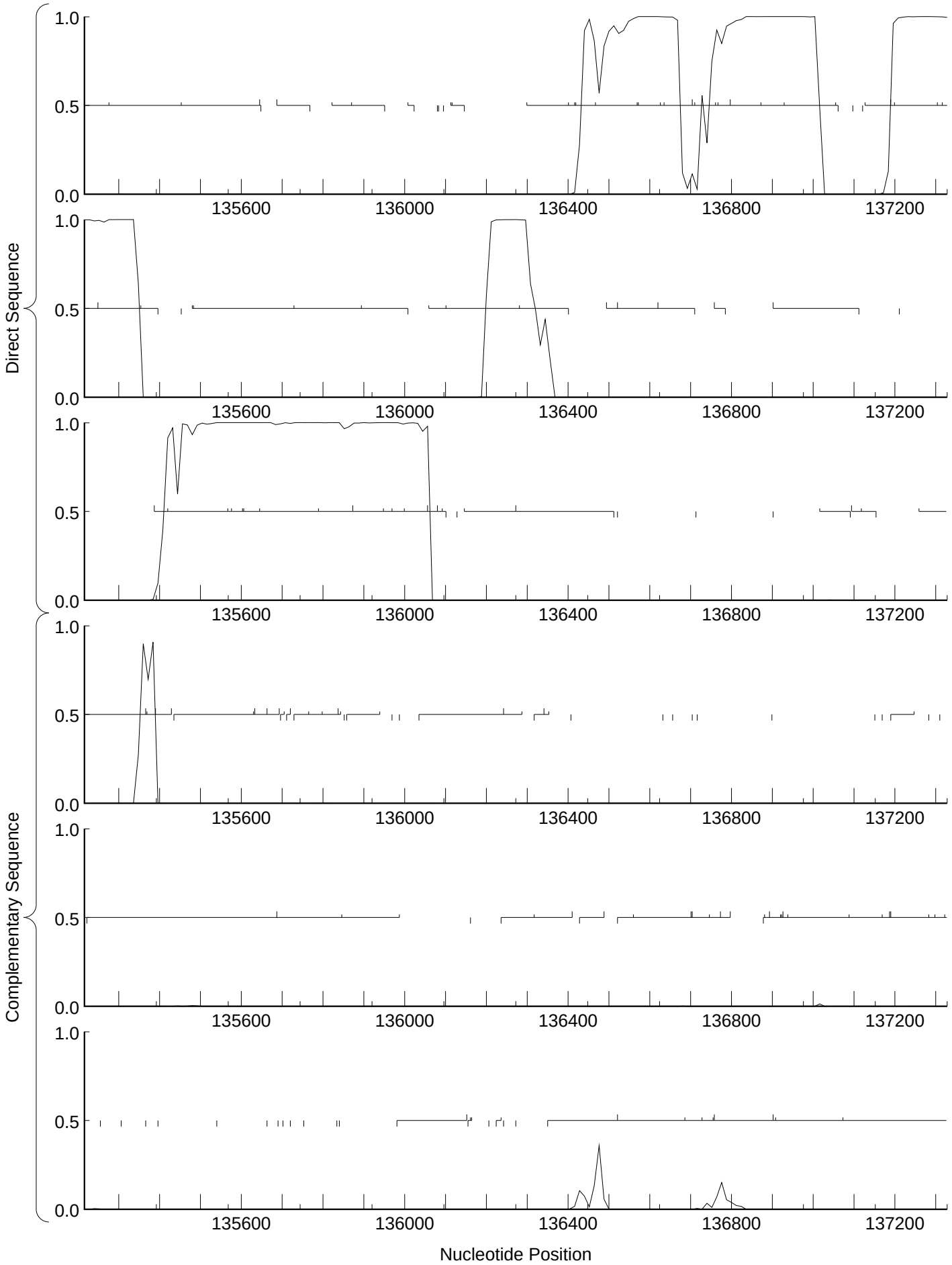
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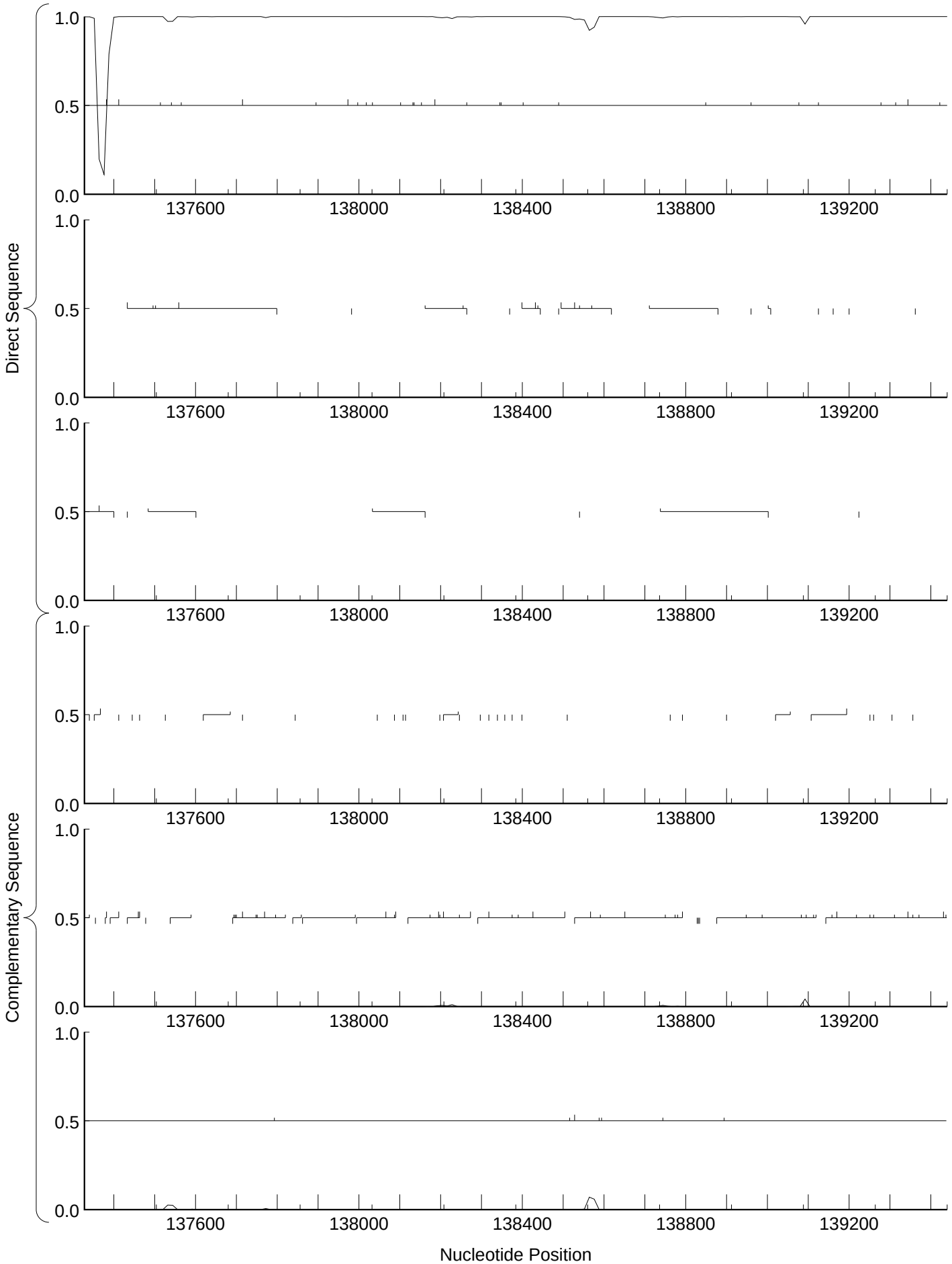
ycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 65/75



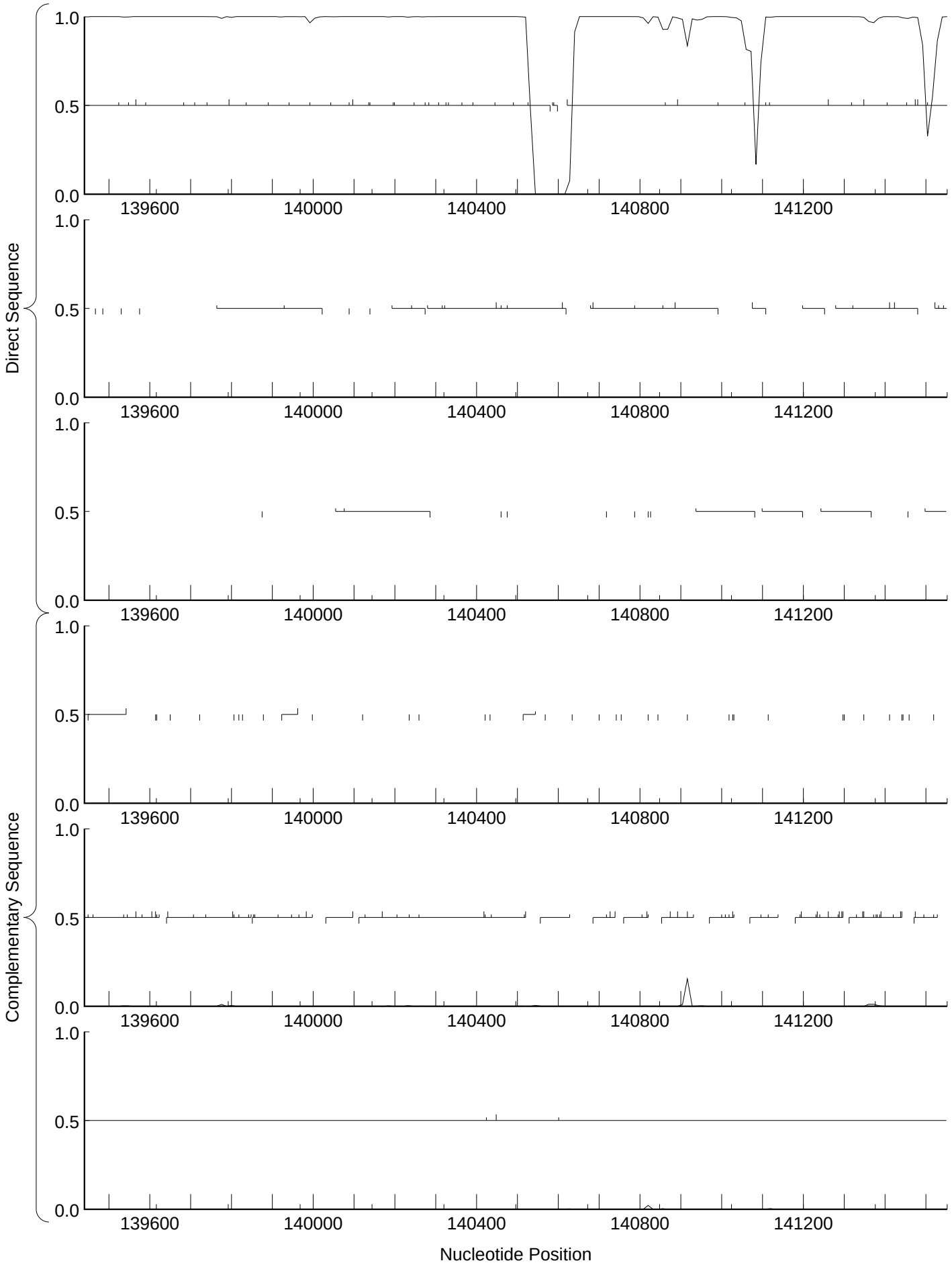
ycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 66/75



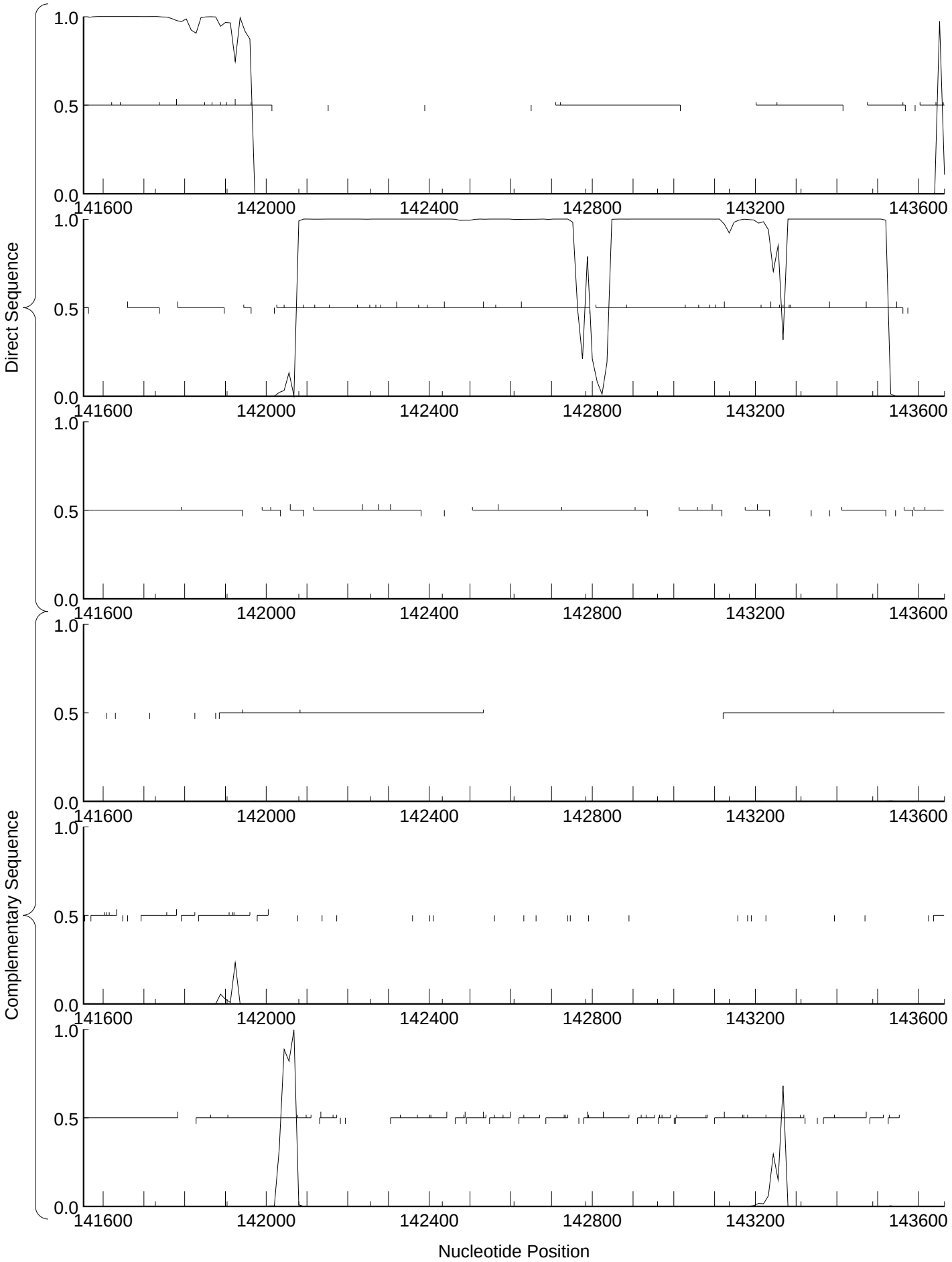
ycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 67/75



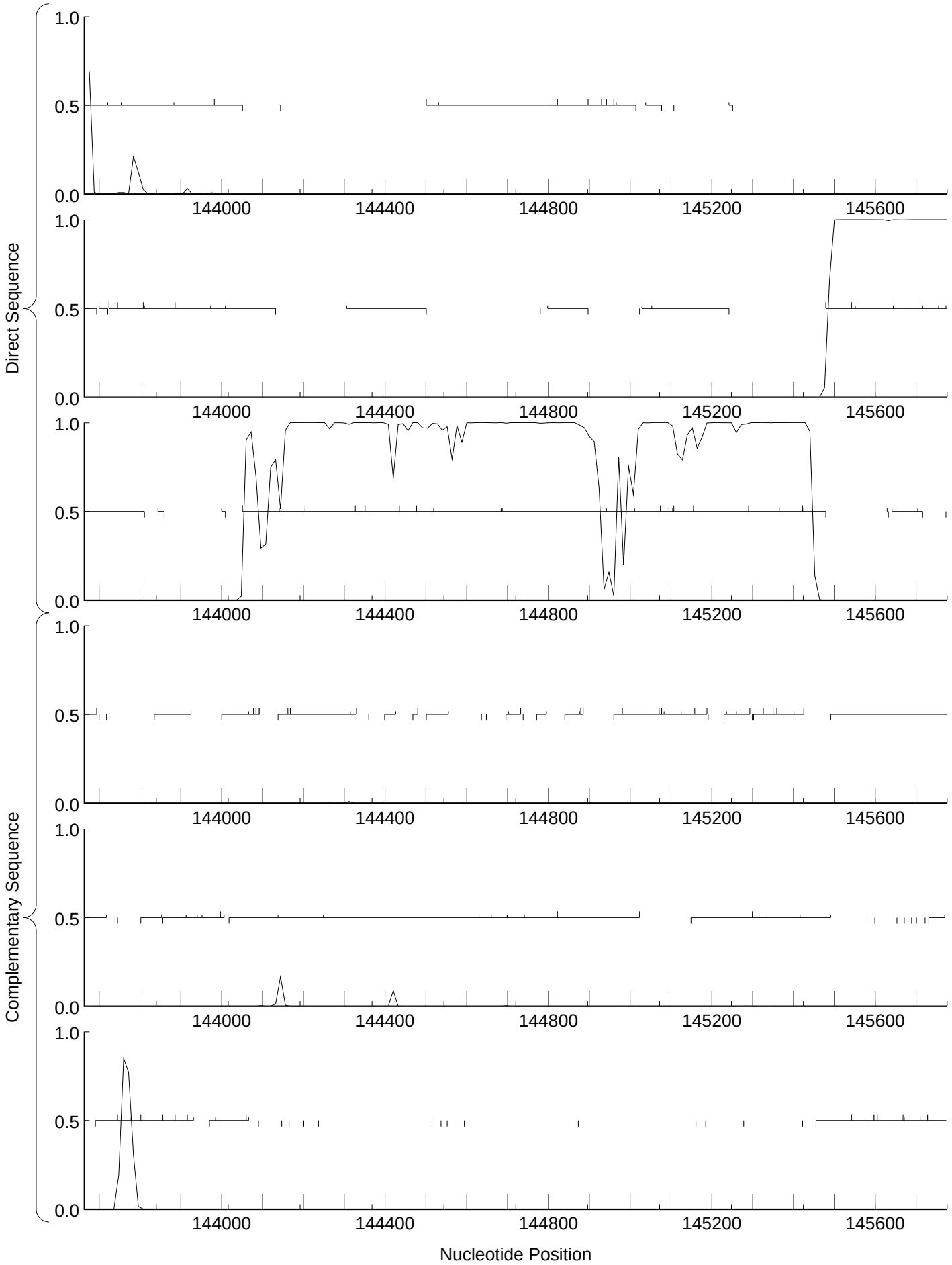
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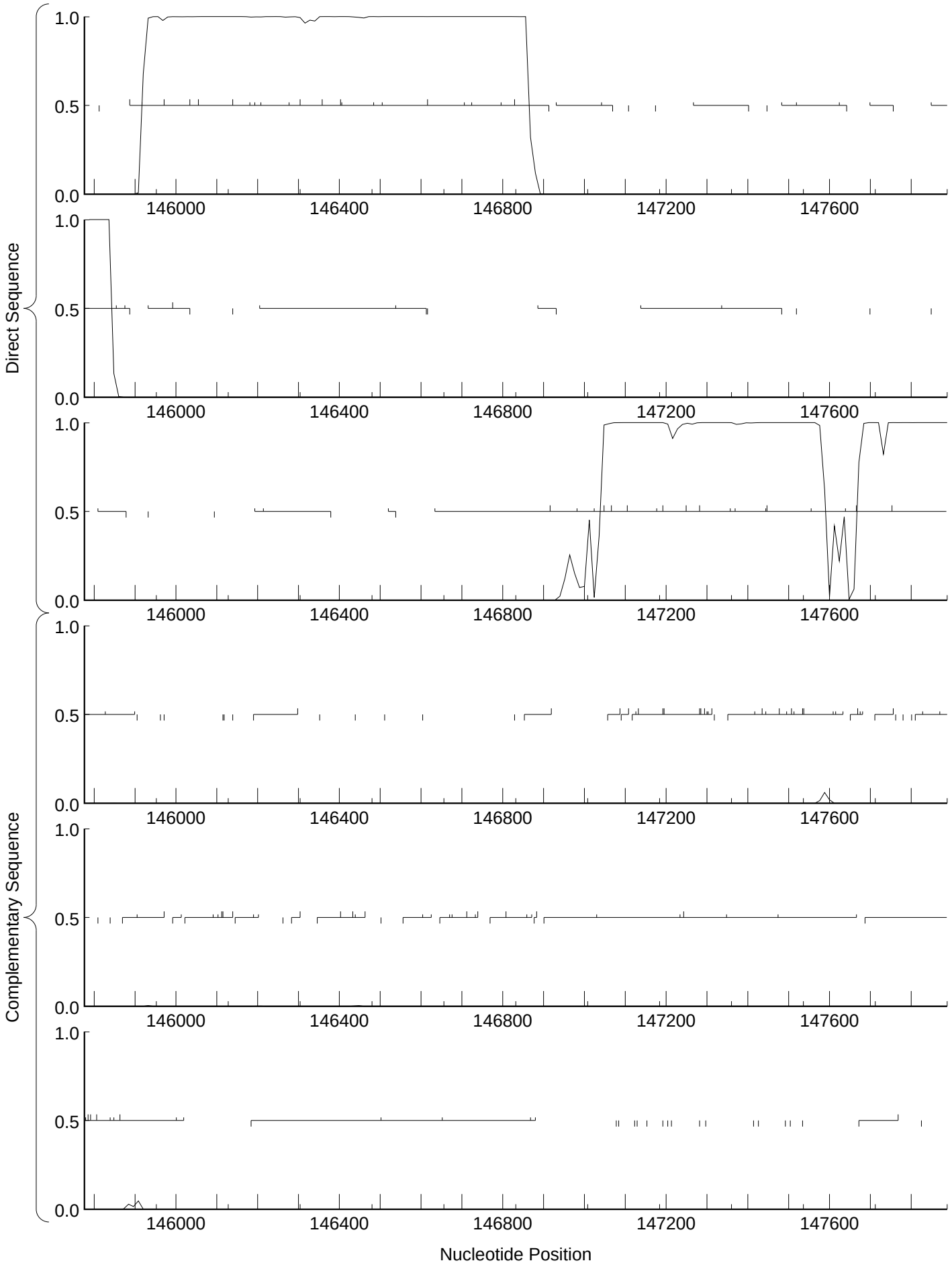
ycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 69/75



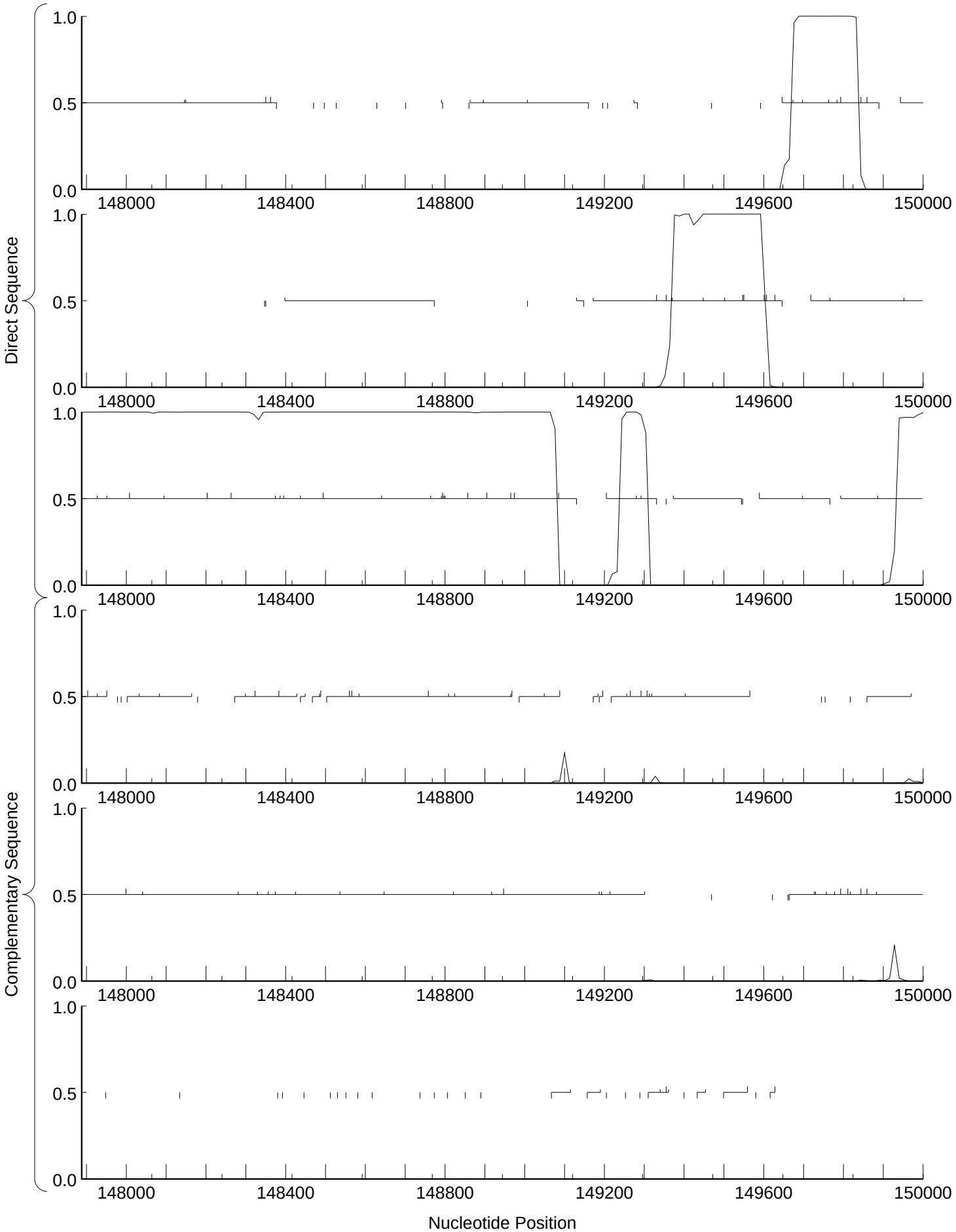
ycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 70/75



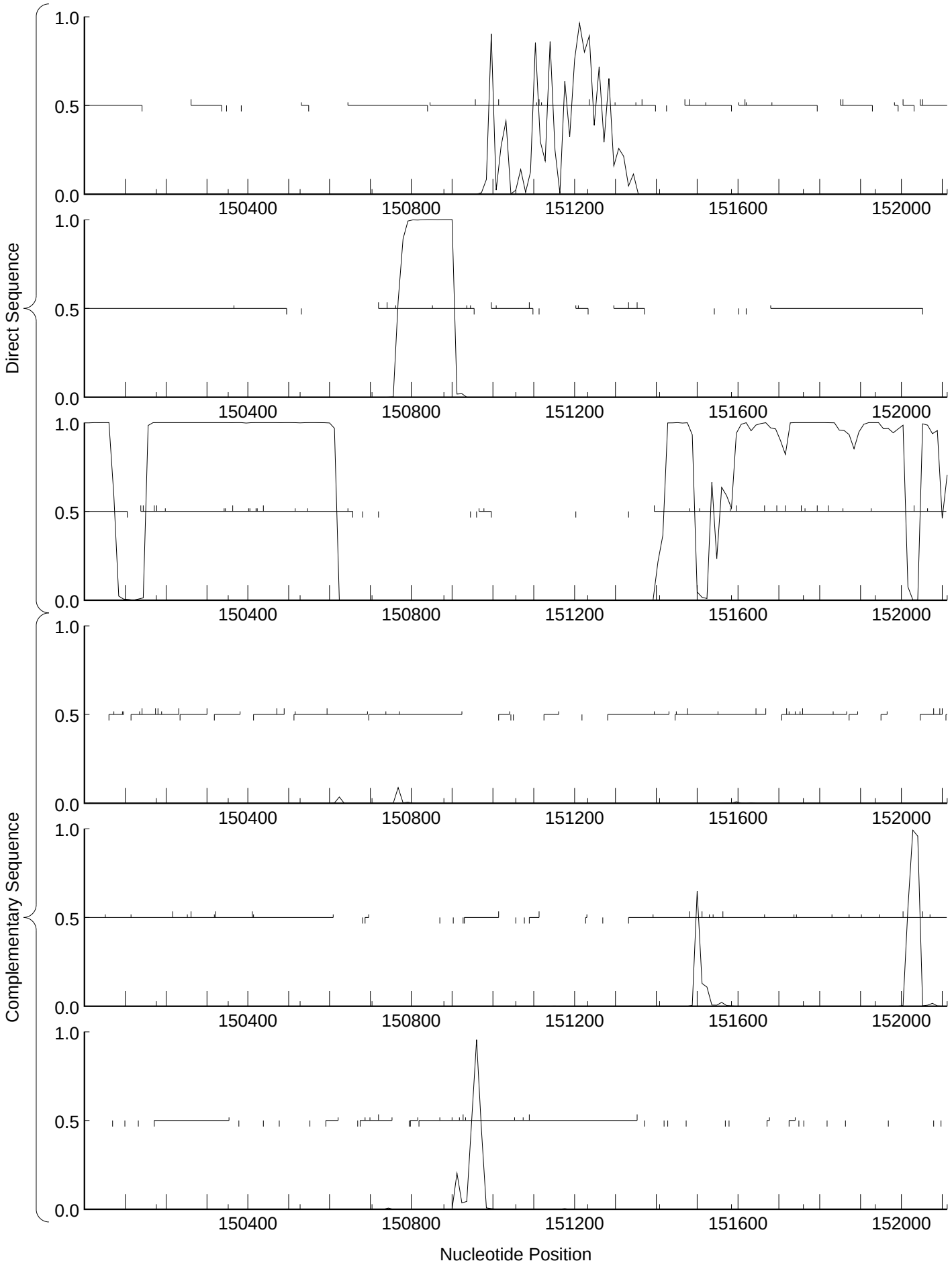
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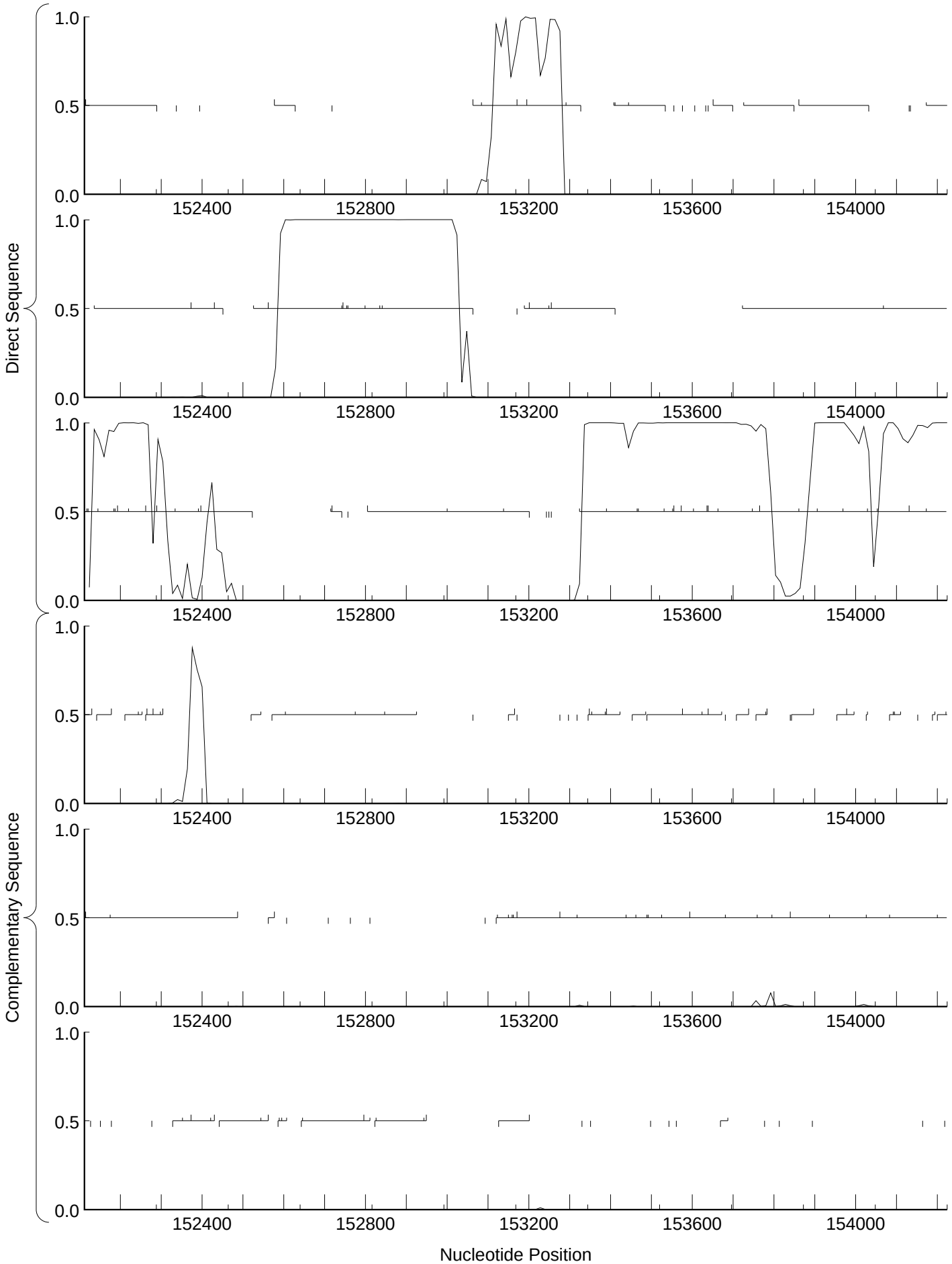
ycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 72/75



ycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 73/75



ycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 74/75



ycobacterium phage Yucca complete sequence, 155582 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 75/75

