GeneMark

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

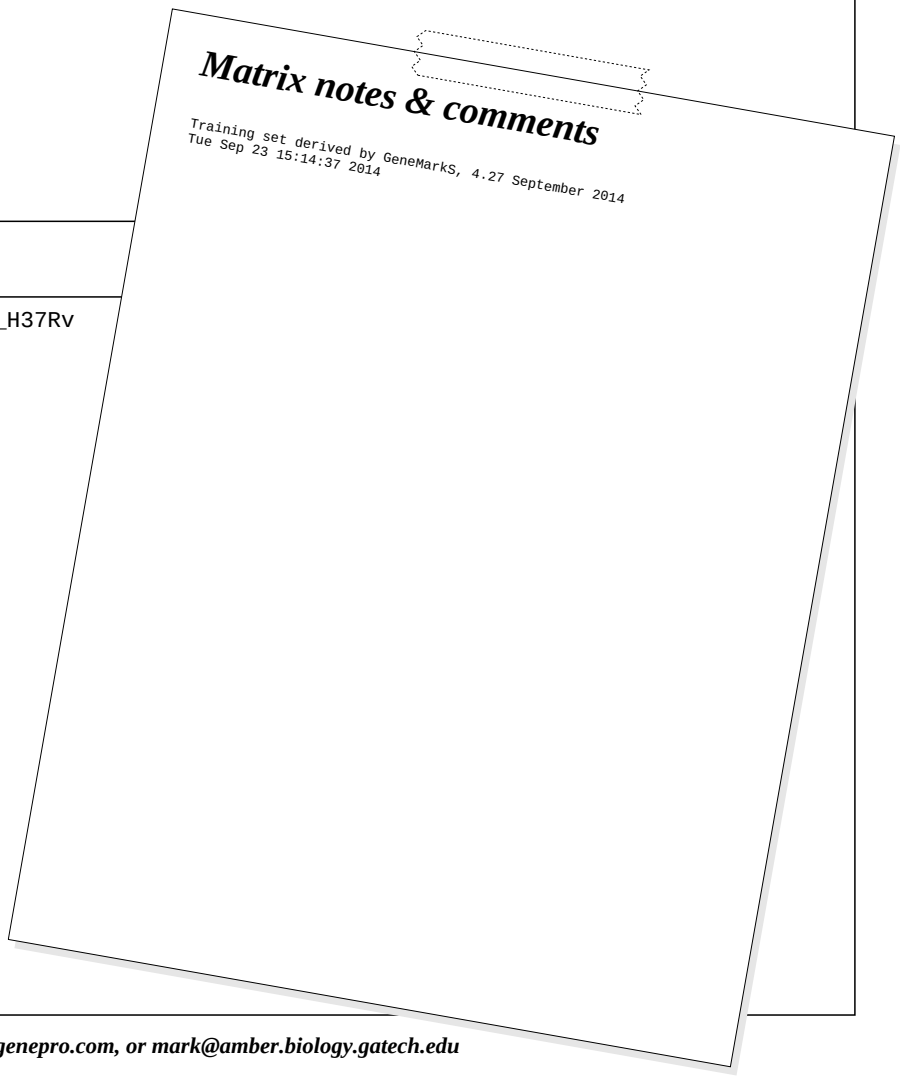
PROGRAM INFORMATION

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Analysis Date    :  3/29/24 at 8:06:44
Pages            :  76
Sequence Length  :  156319 bp
GC Content       :   64.71%

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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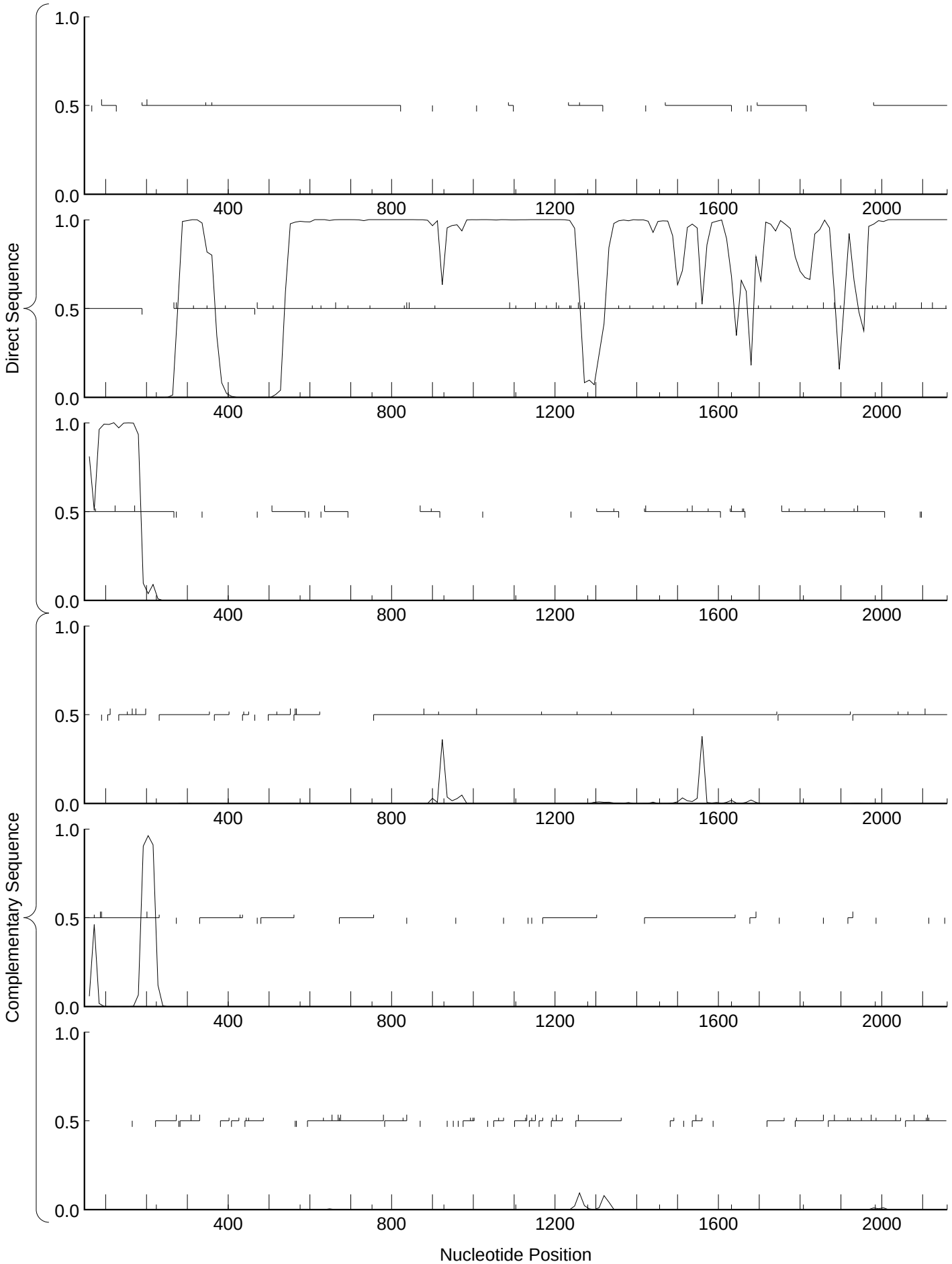
MATRIX INFORMATION

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Author     :  -

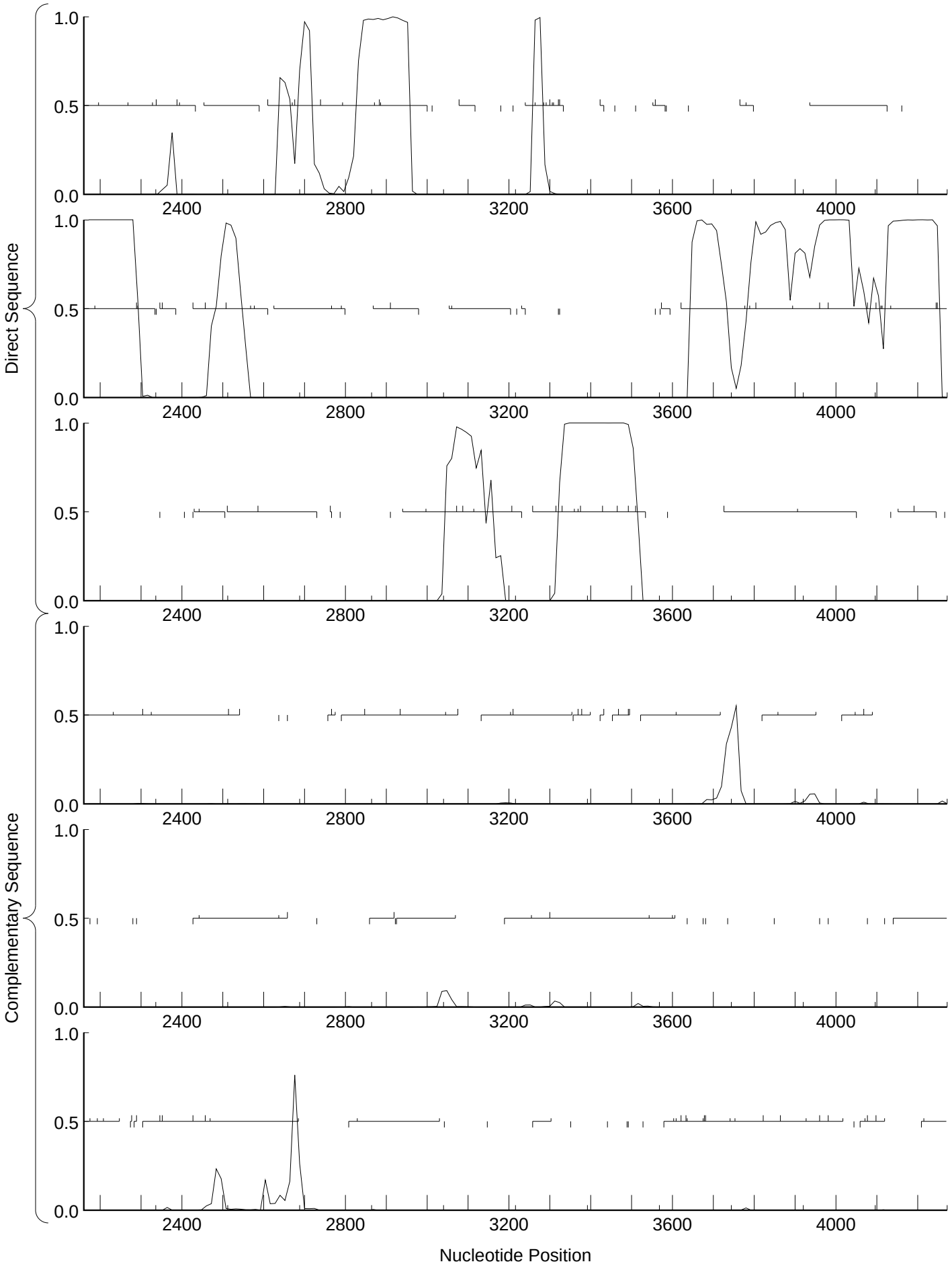
Order      :  4
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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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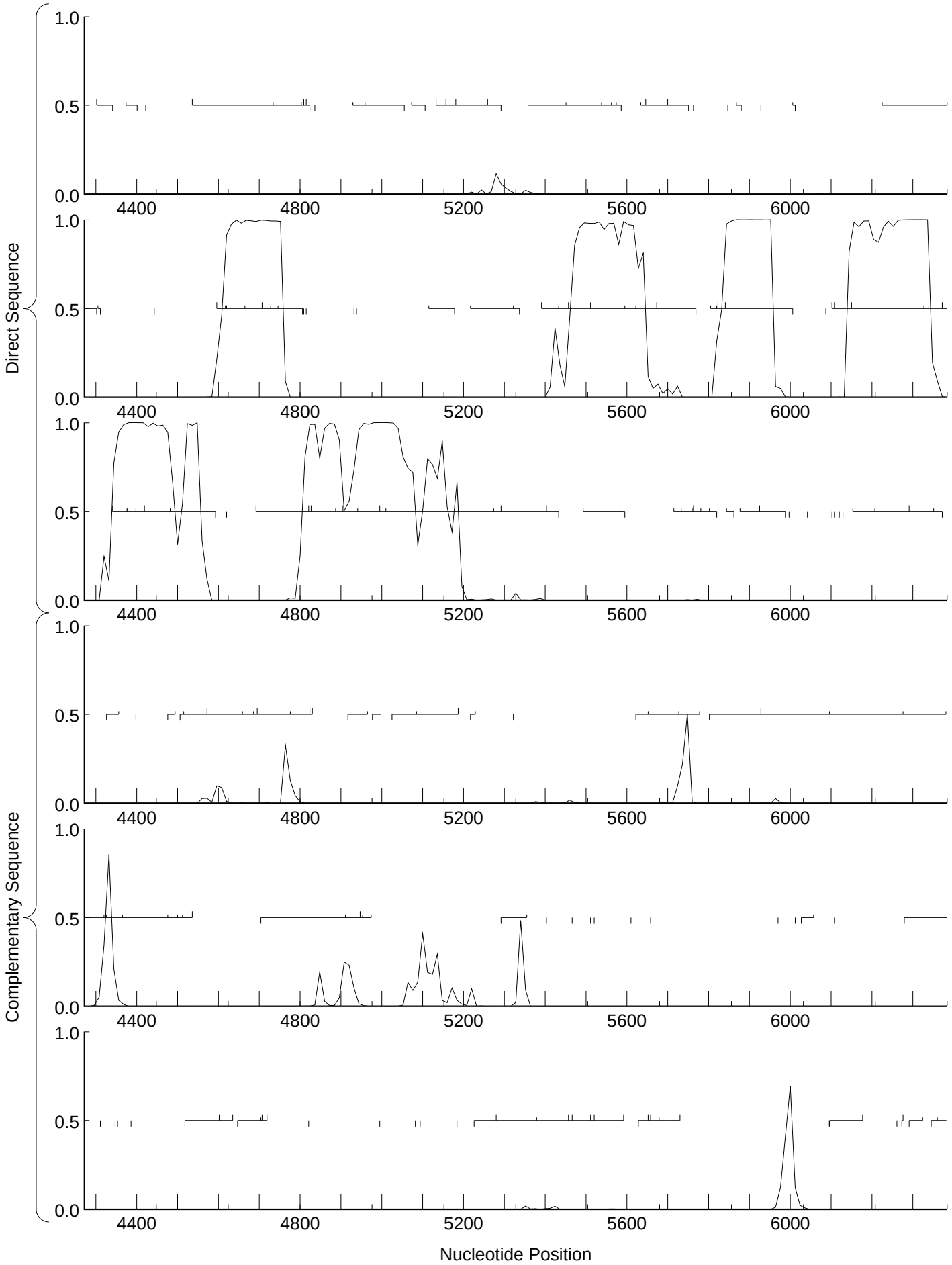

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
400
800
1200
1600
2000
Nucleotide Position

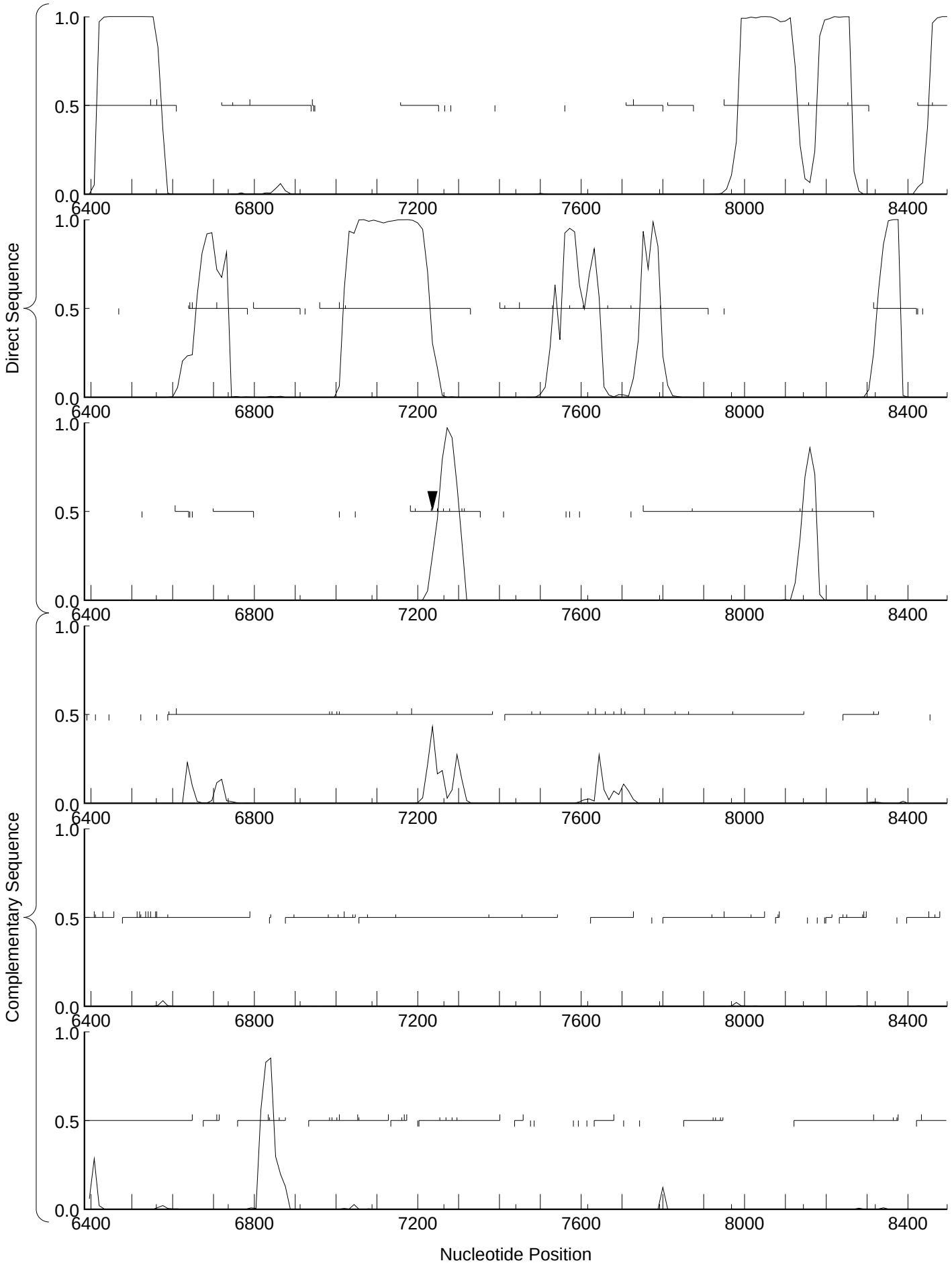
Mycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 3/76

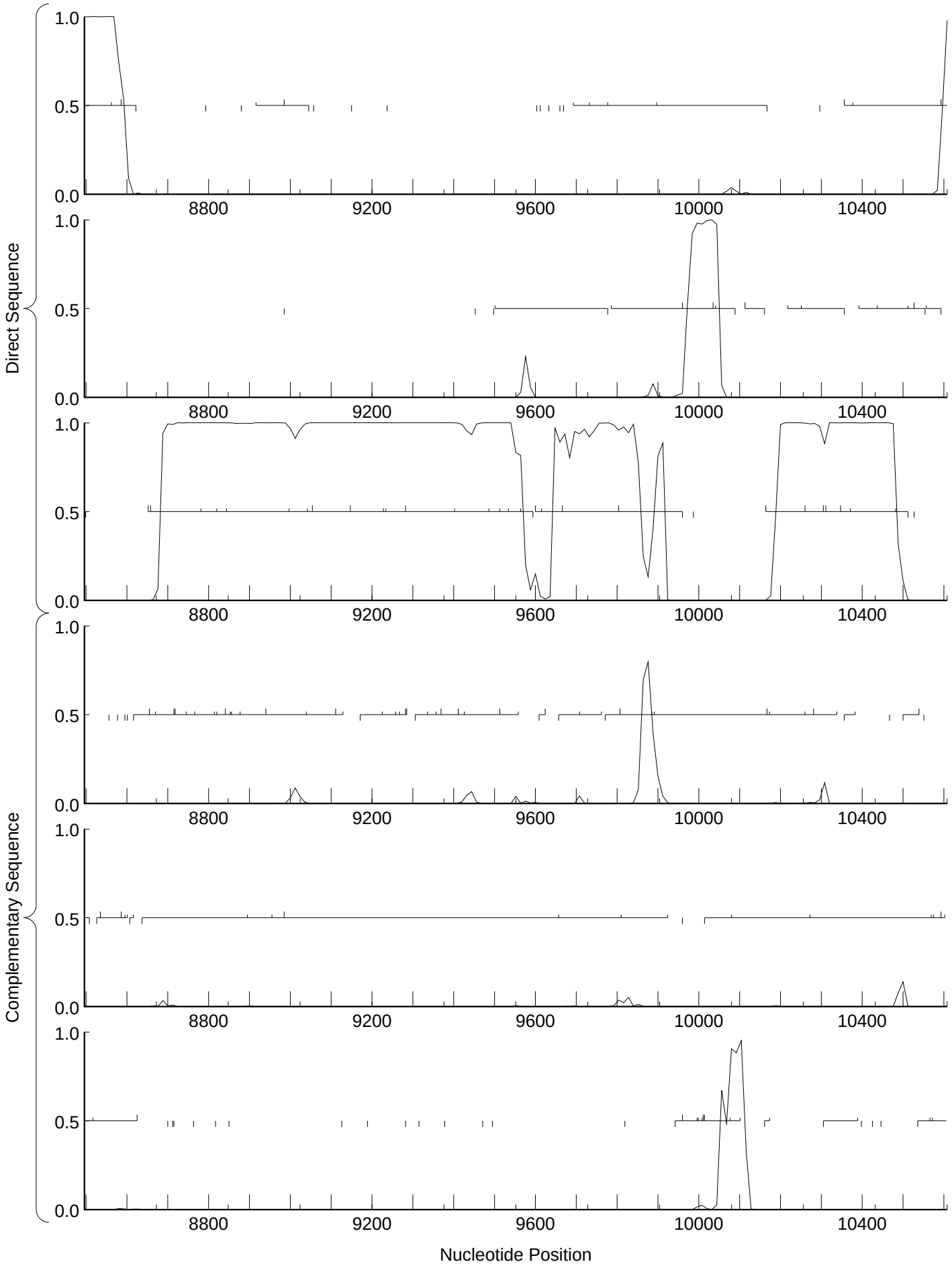


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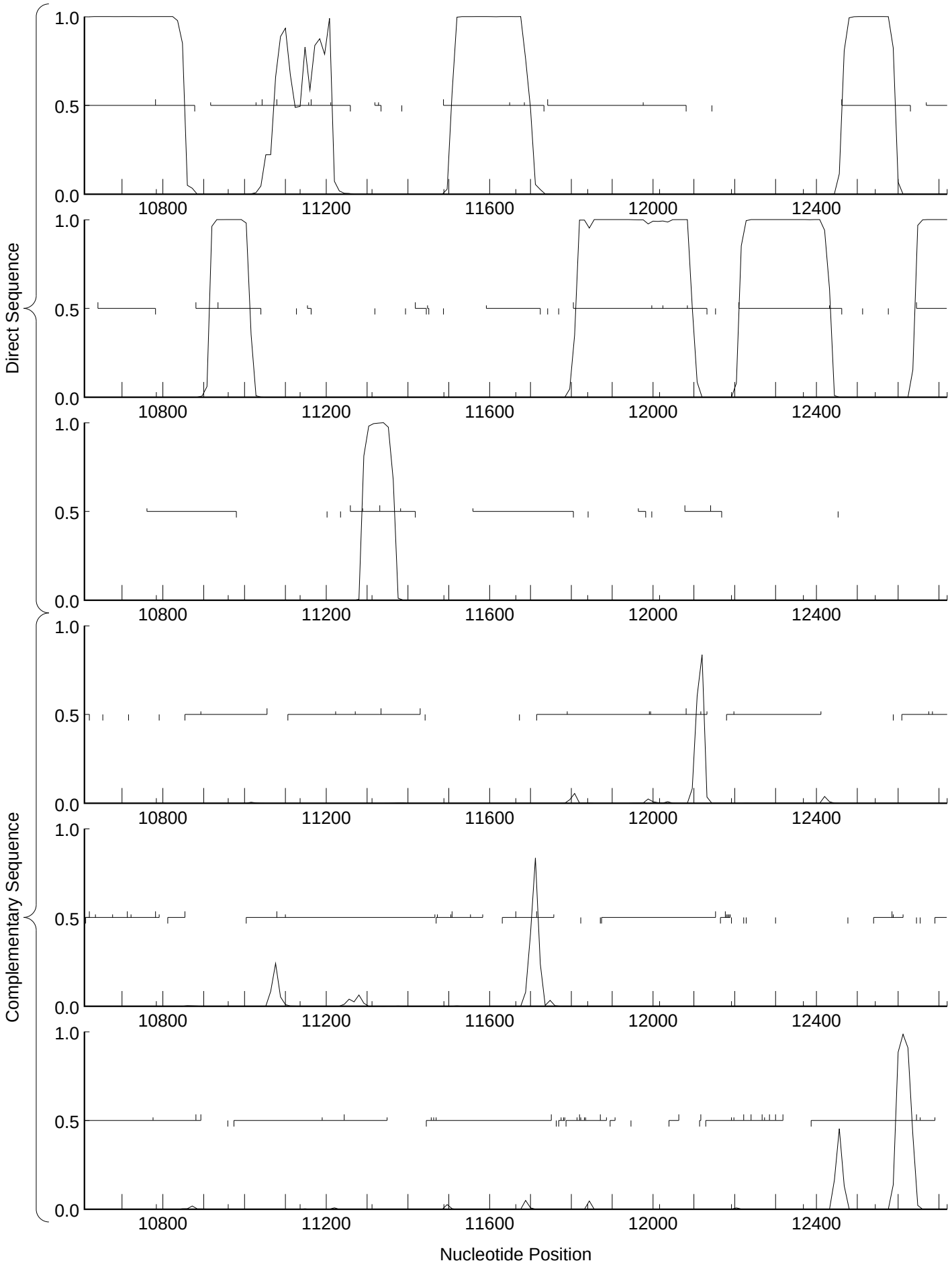


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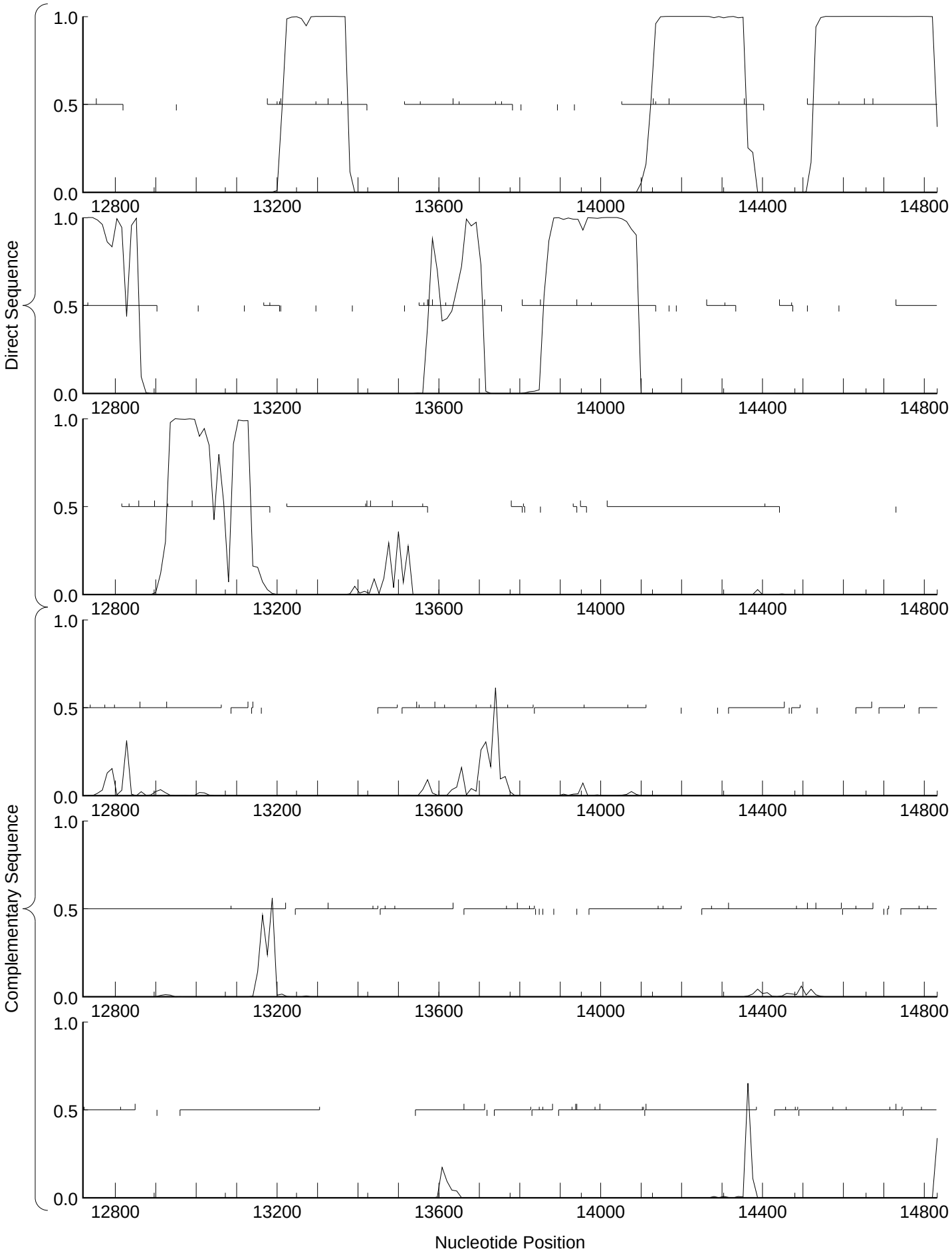



Direct Sequence
Complementary Sequence
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0.5
0.0
8800
9200
9600
10000
10400
Nucleotide Position

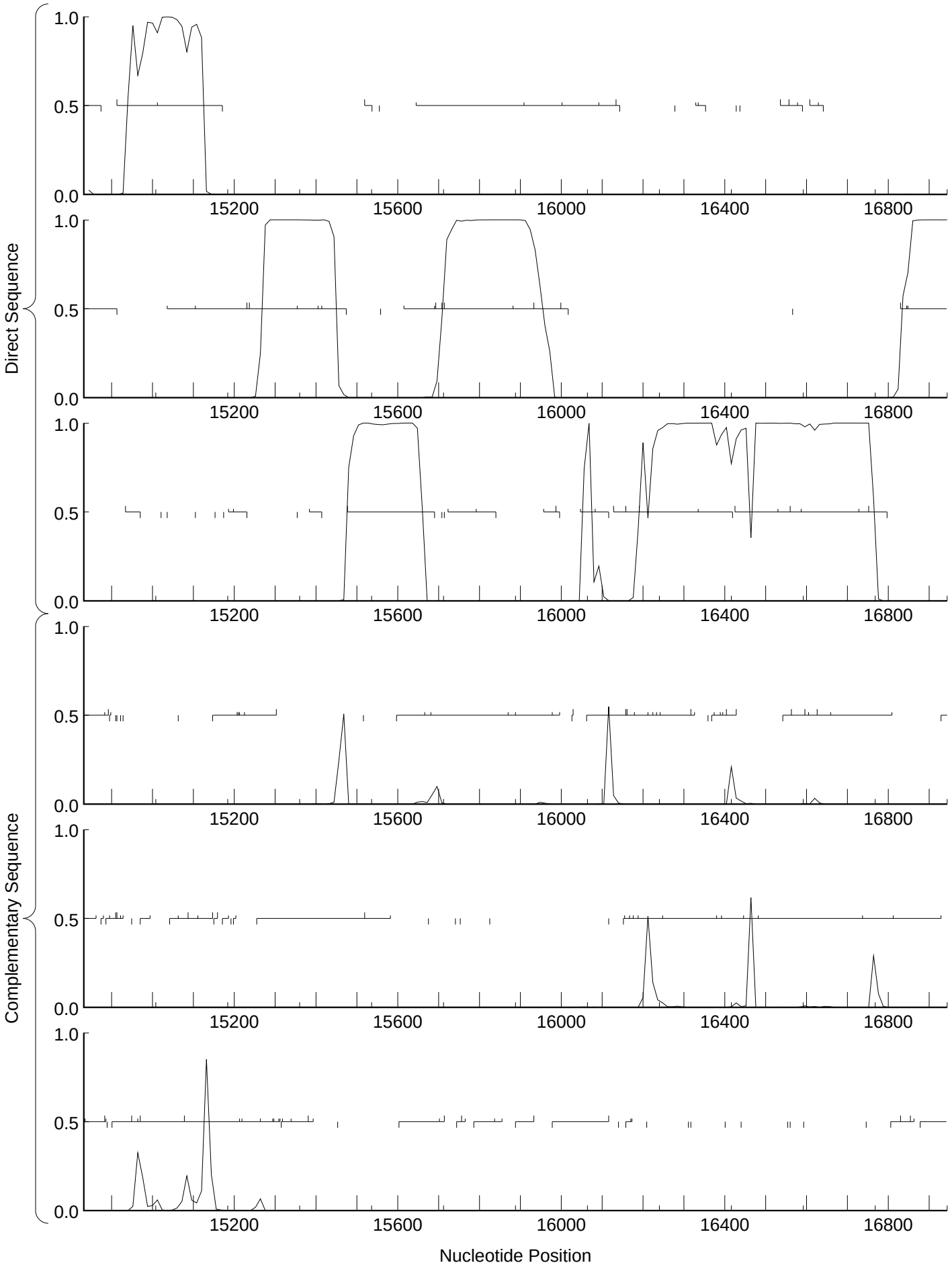
Mycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 7/76



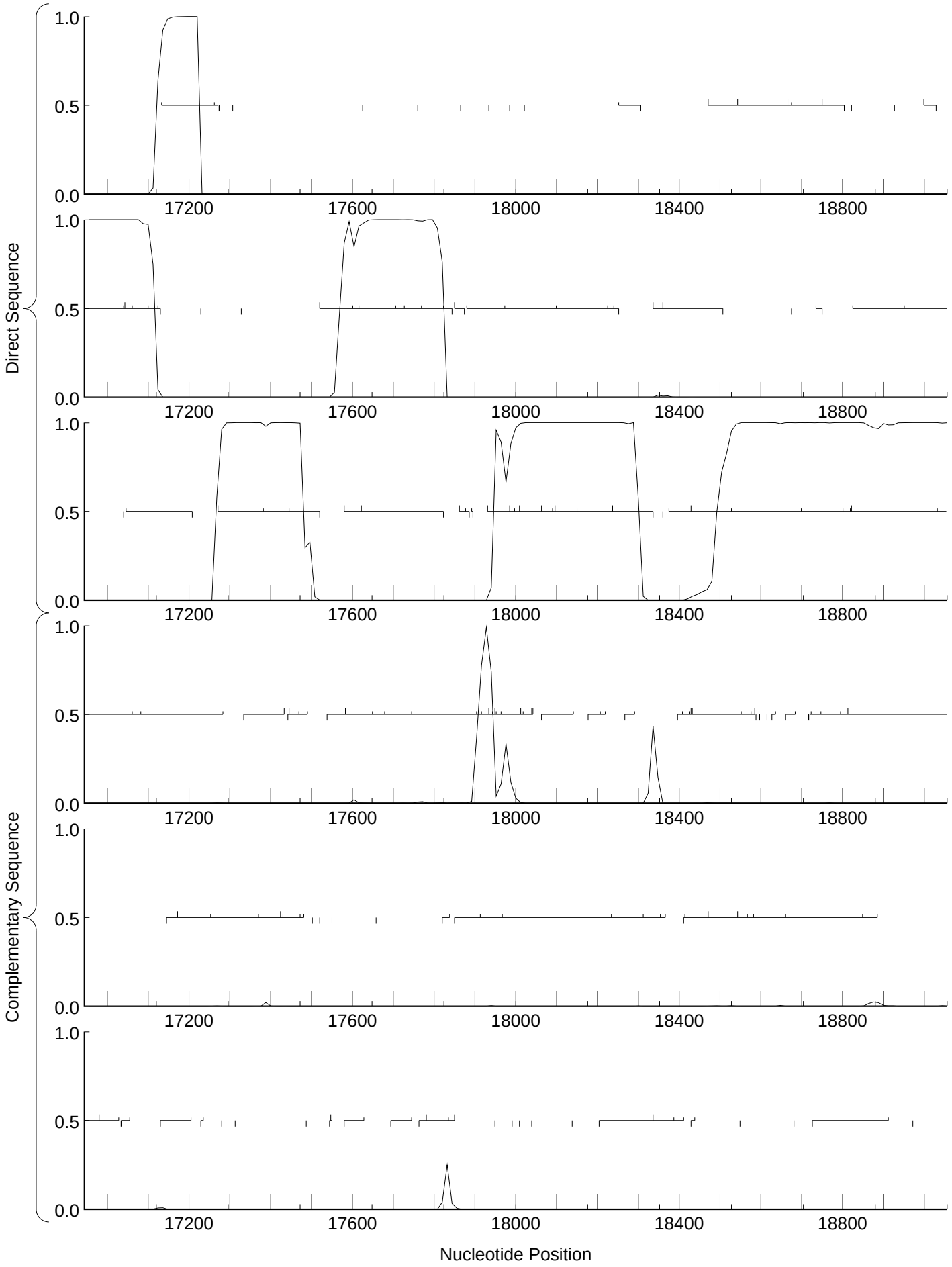
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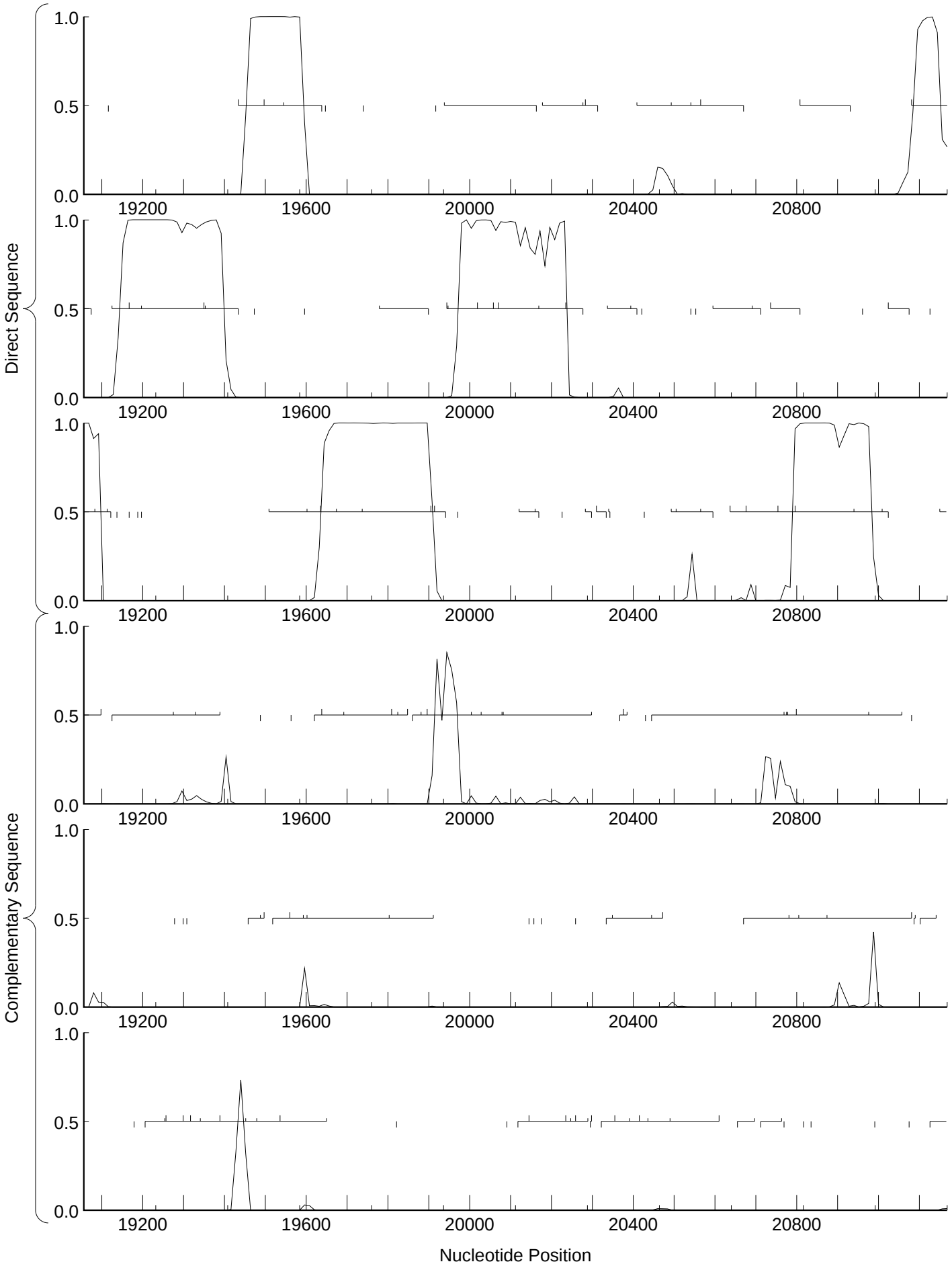
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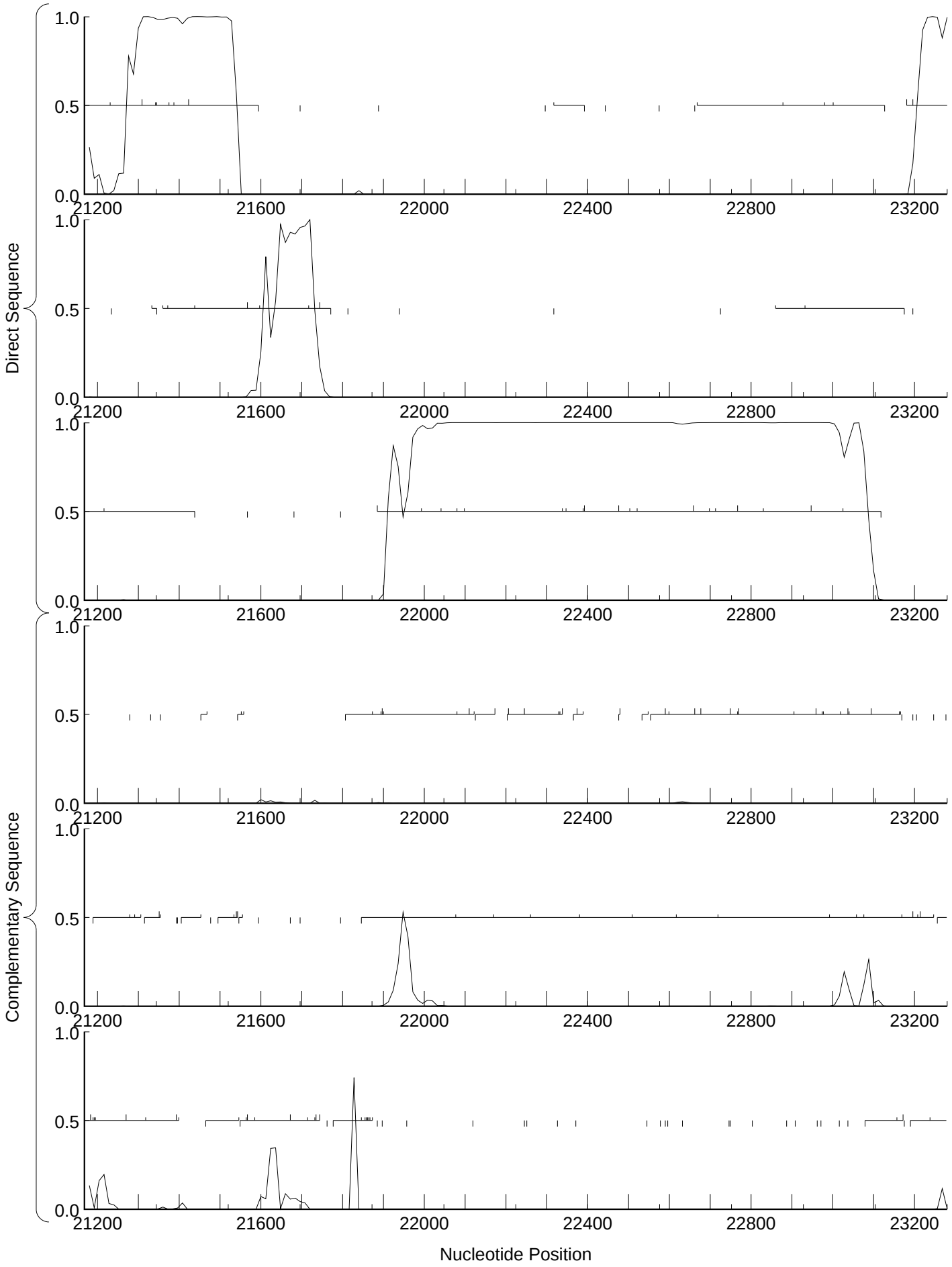
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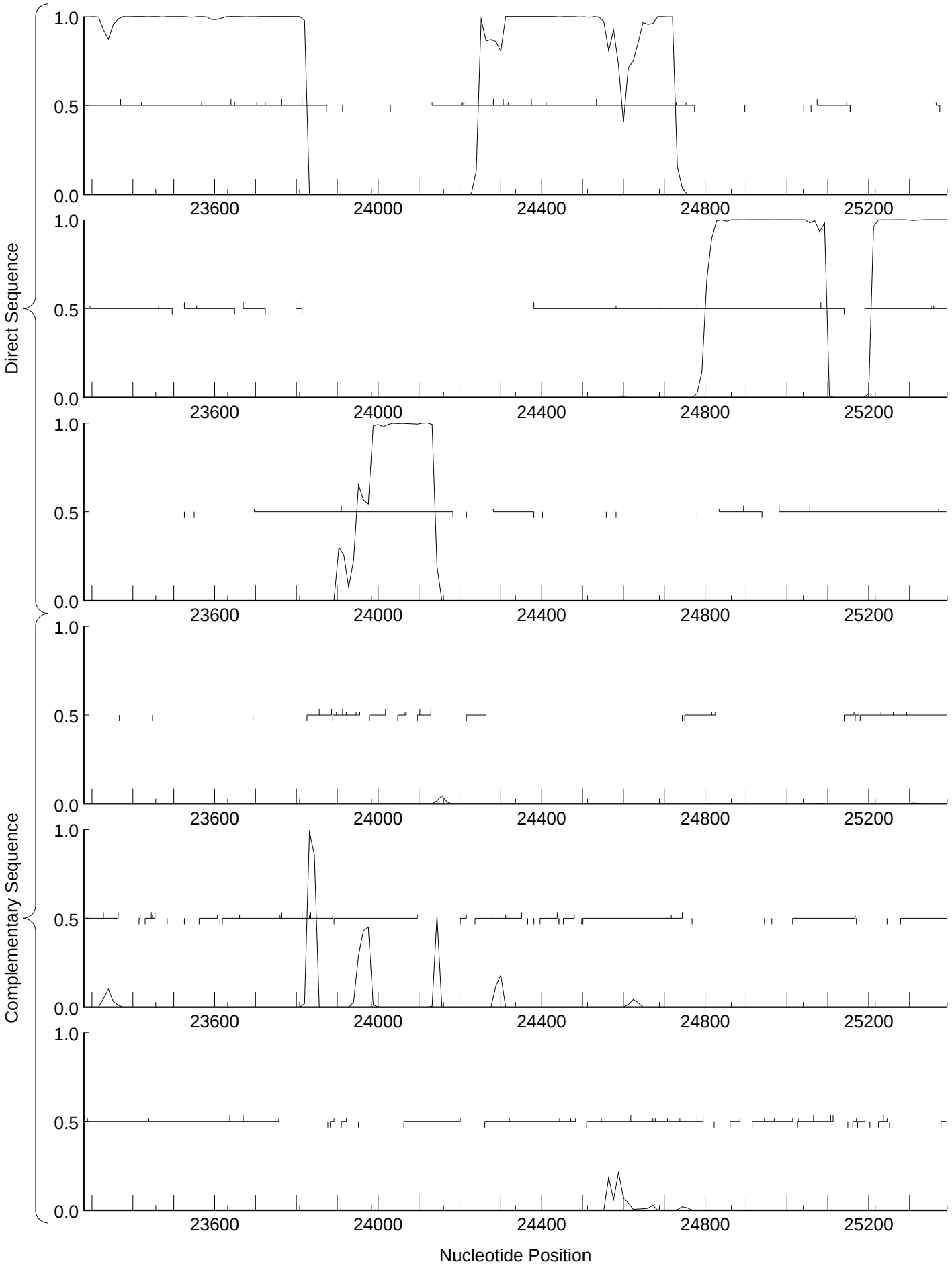
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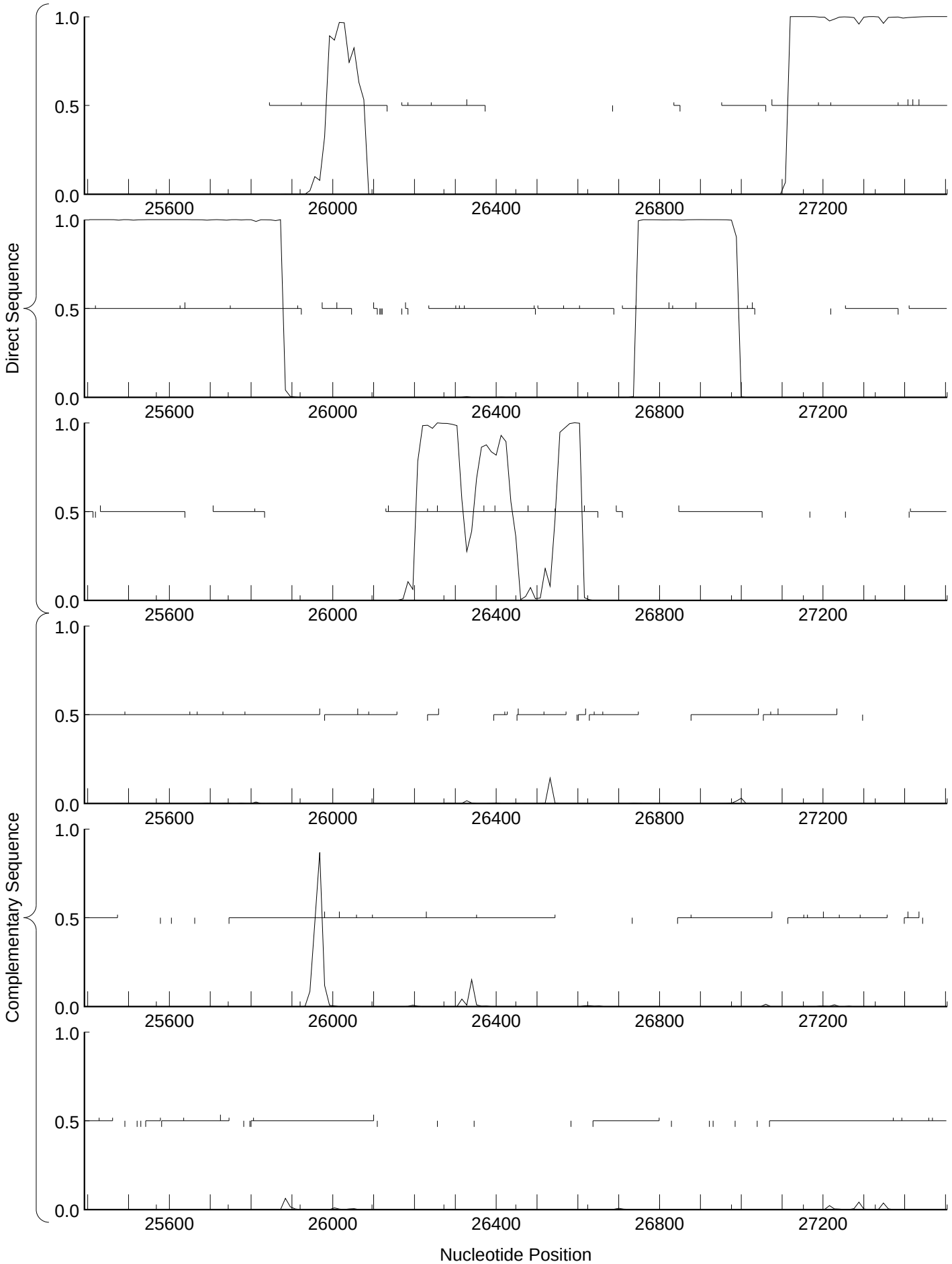
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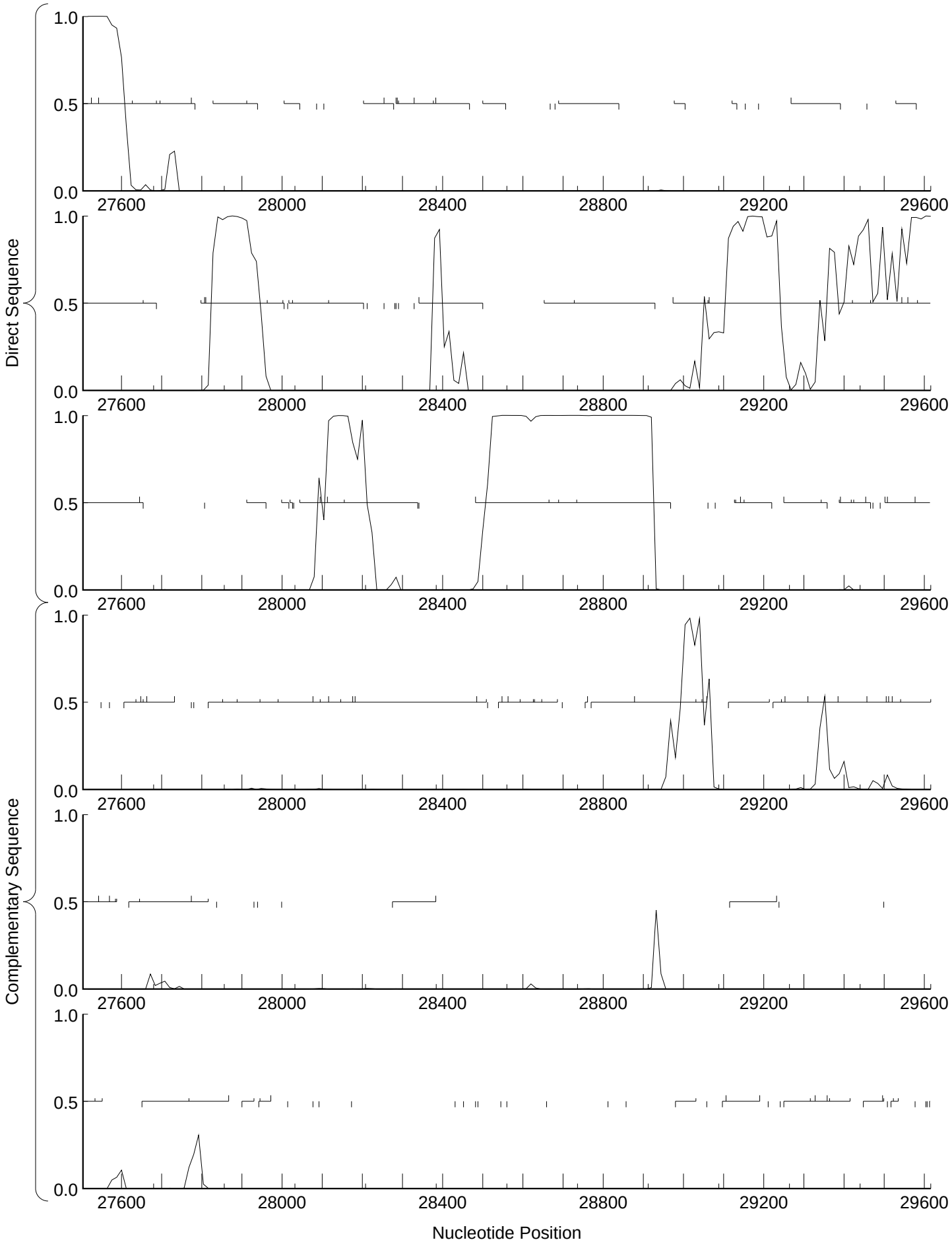
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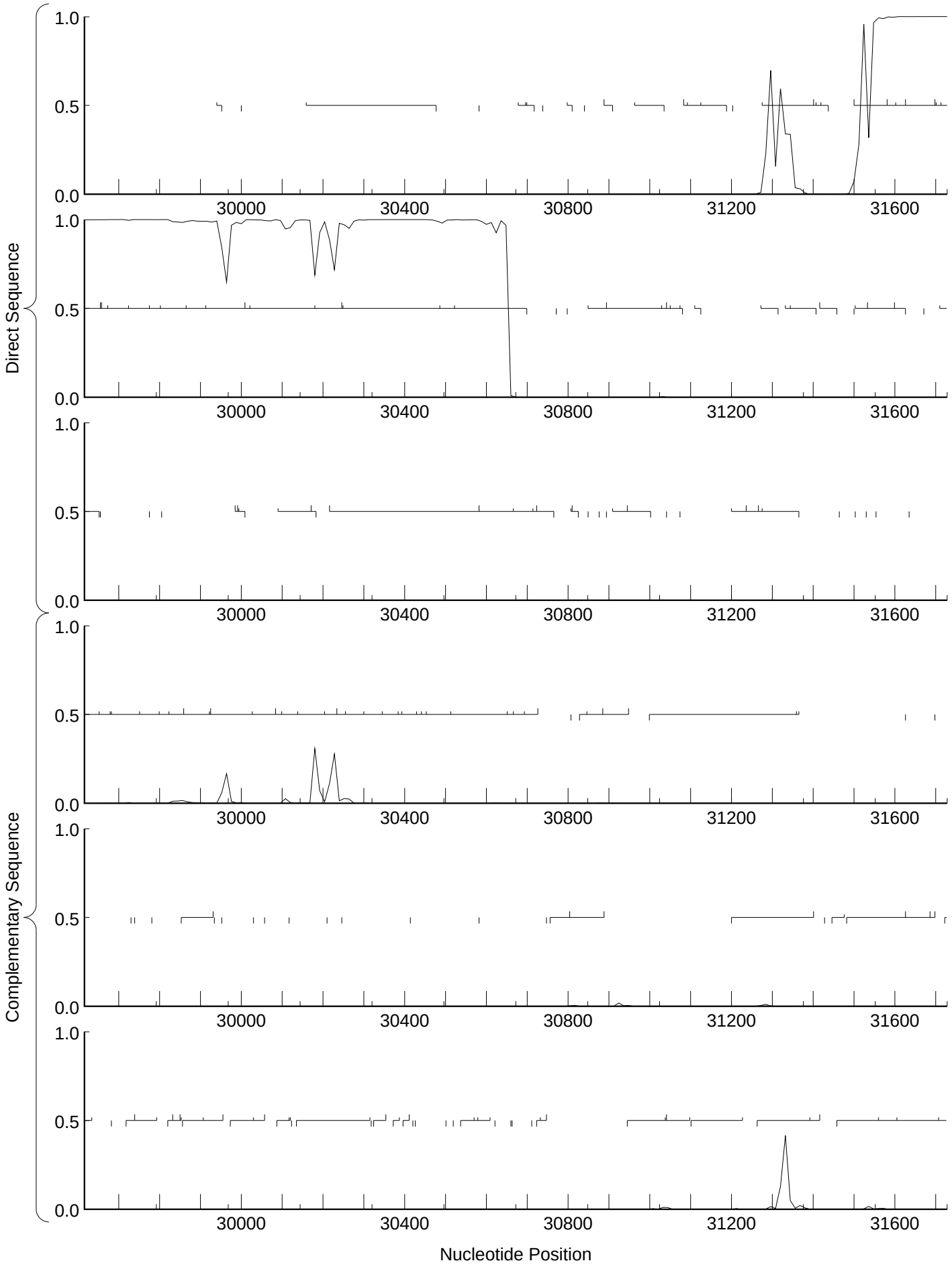
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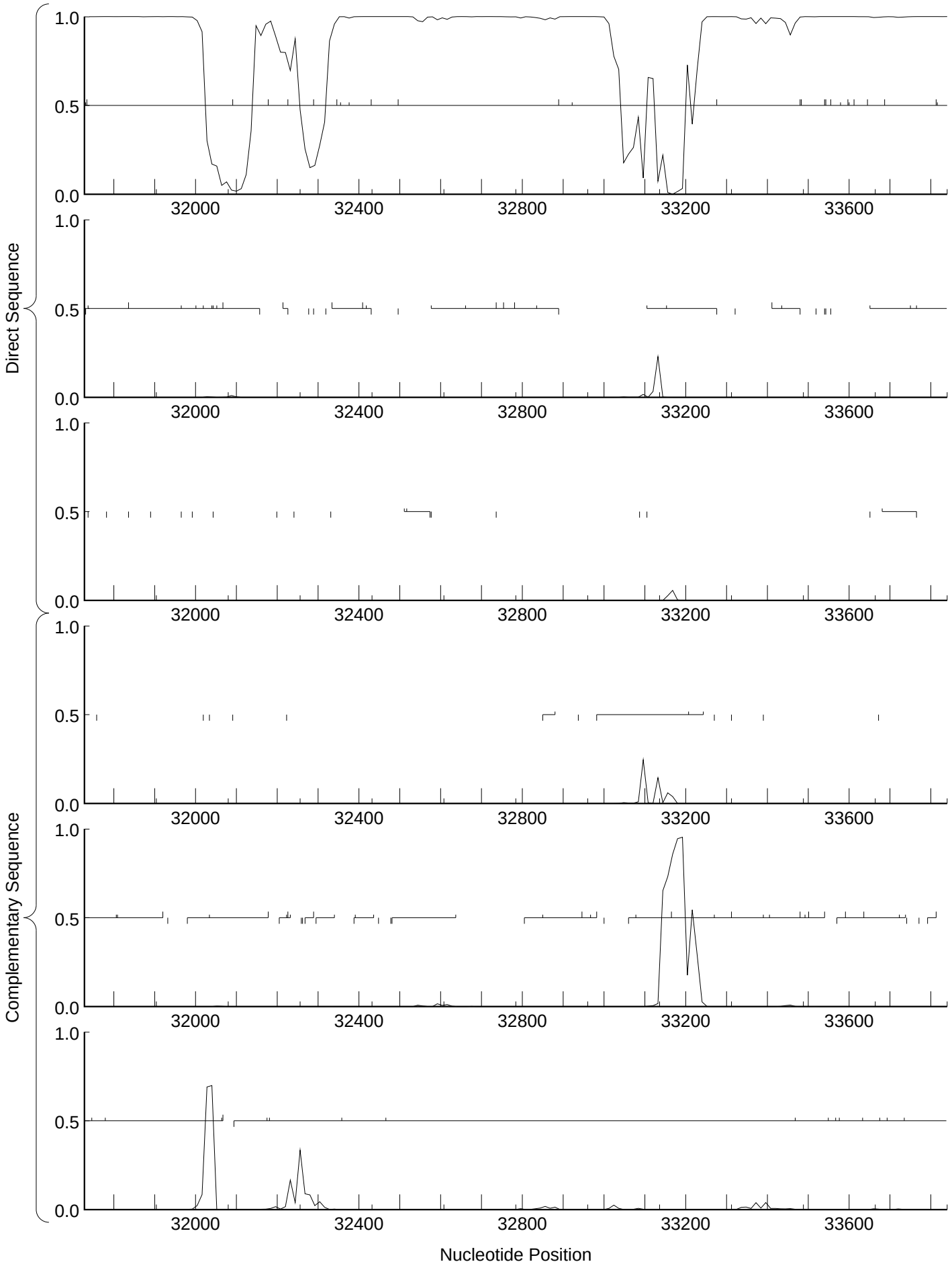
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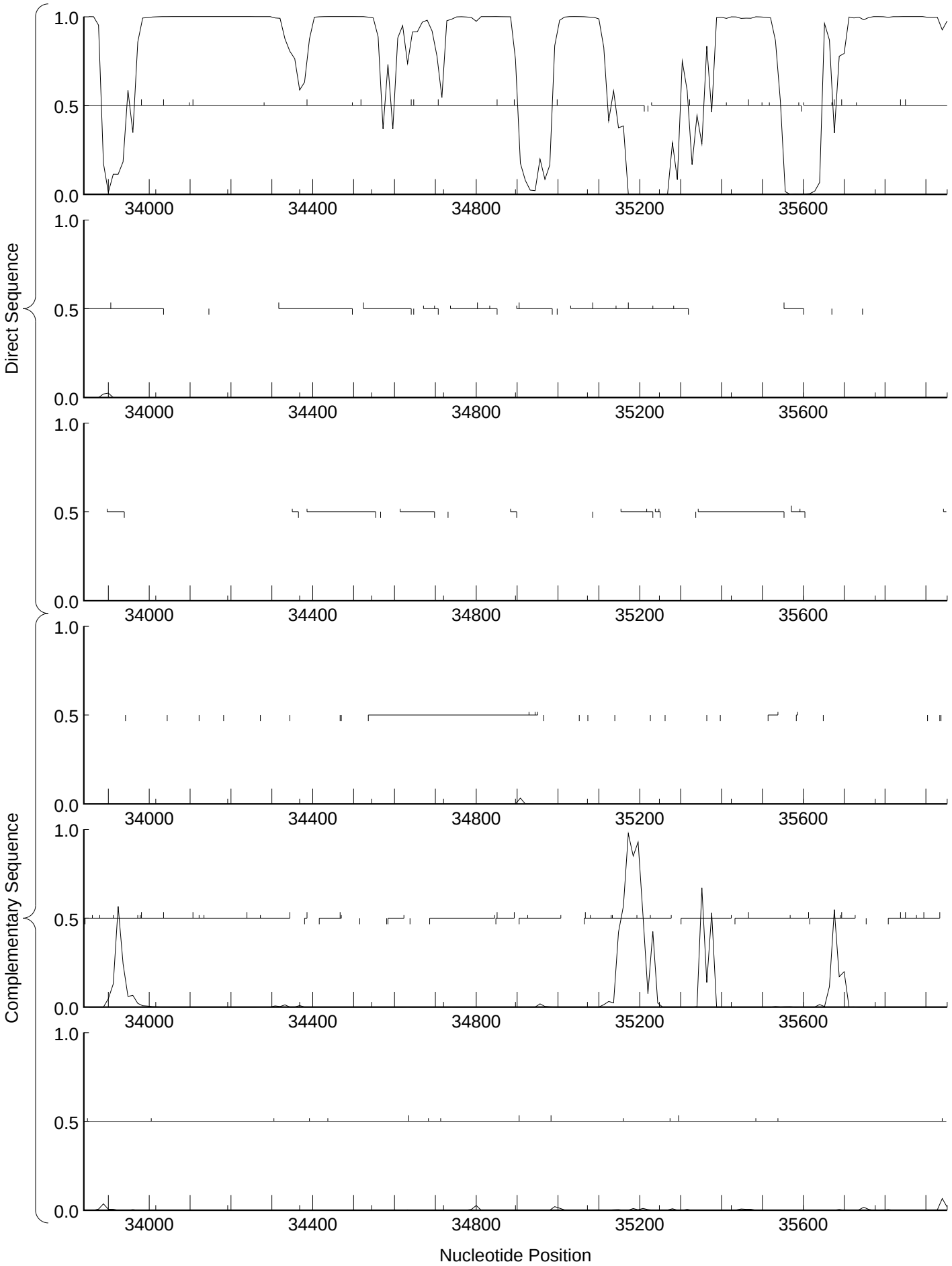
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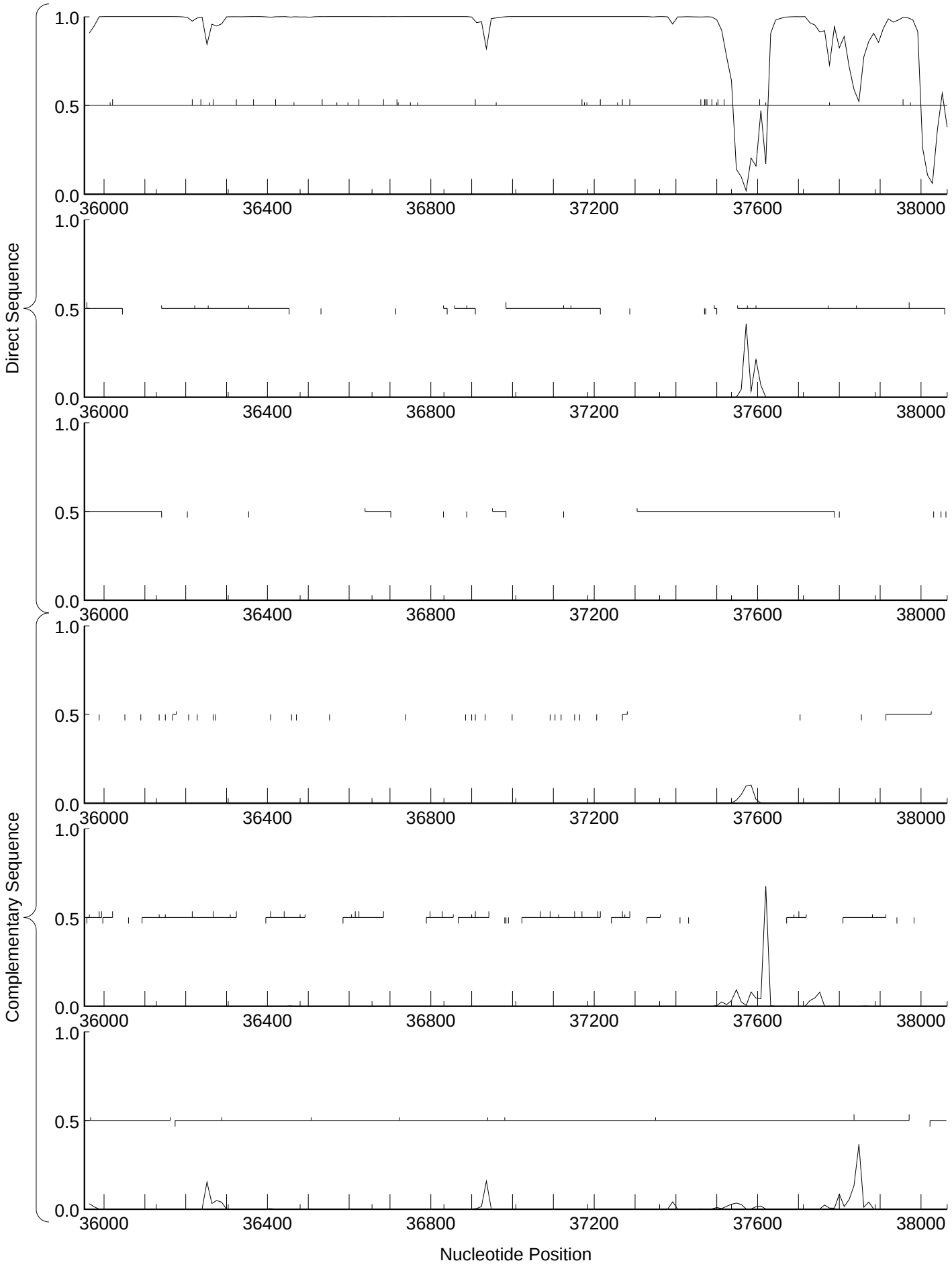
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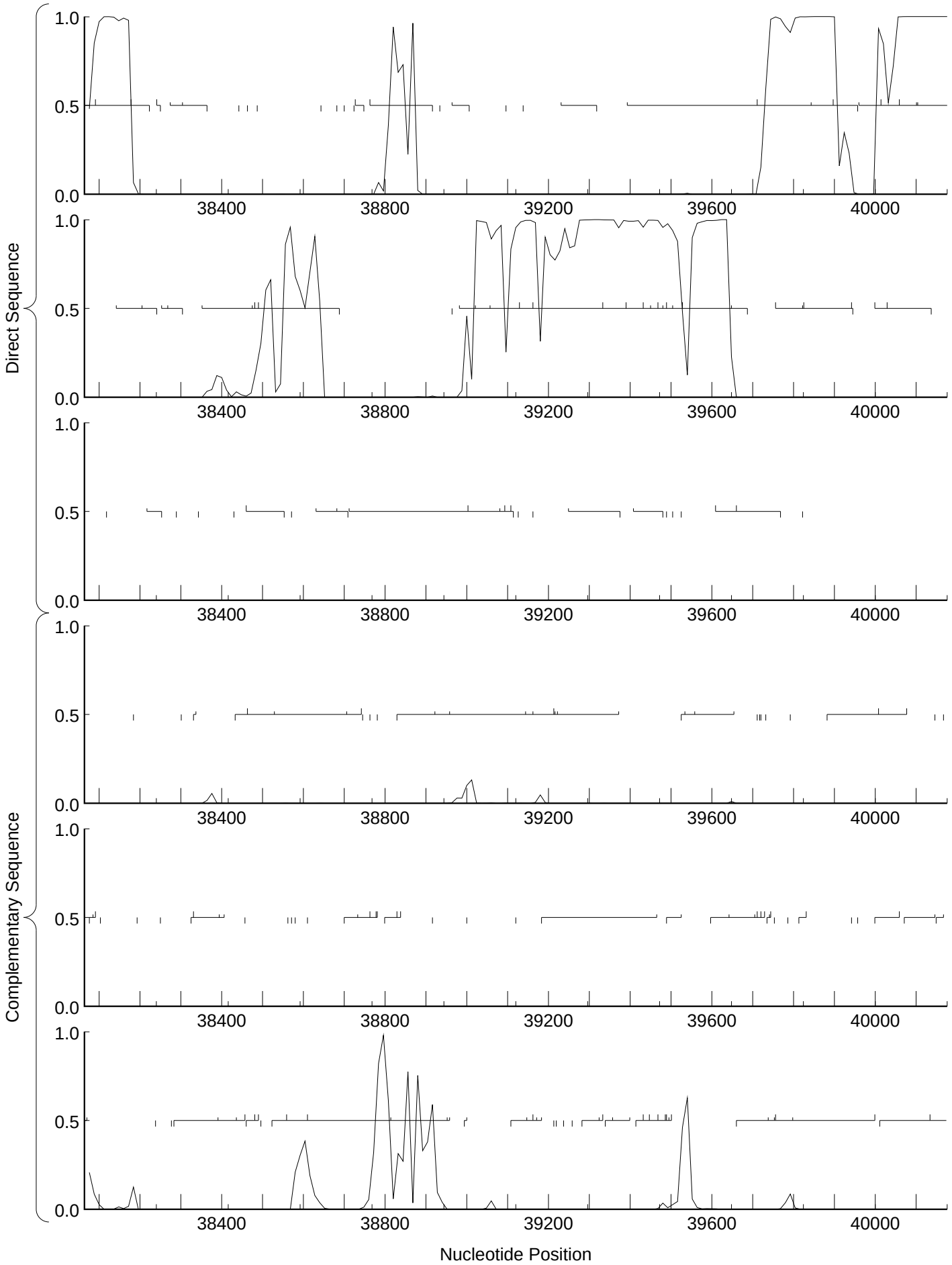
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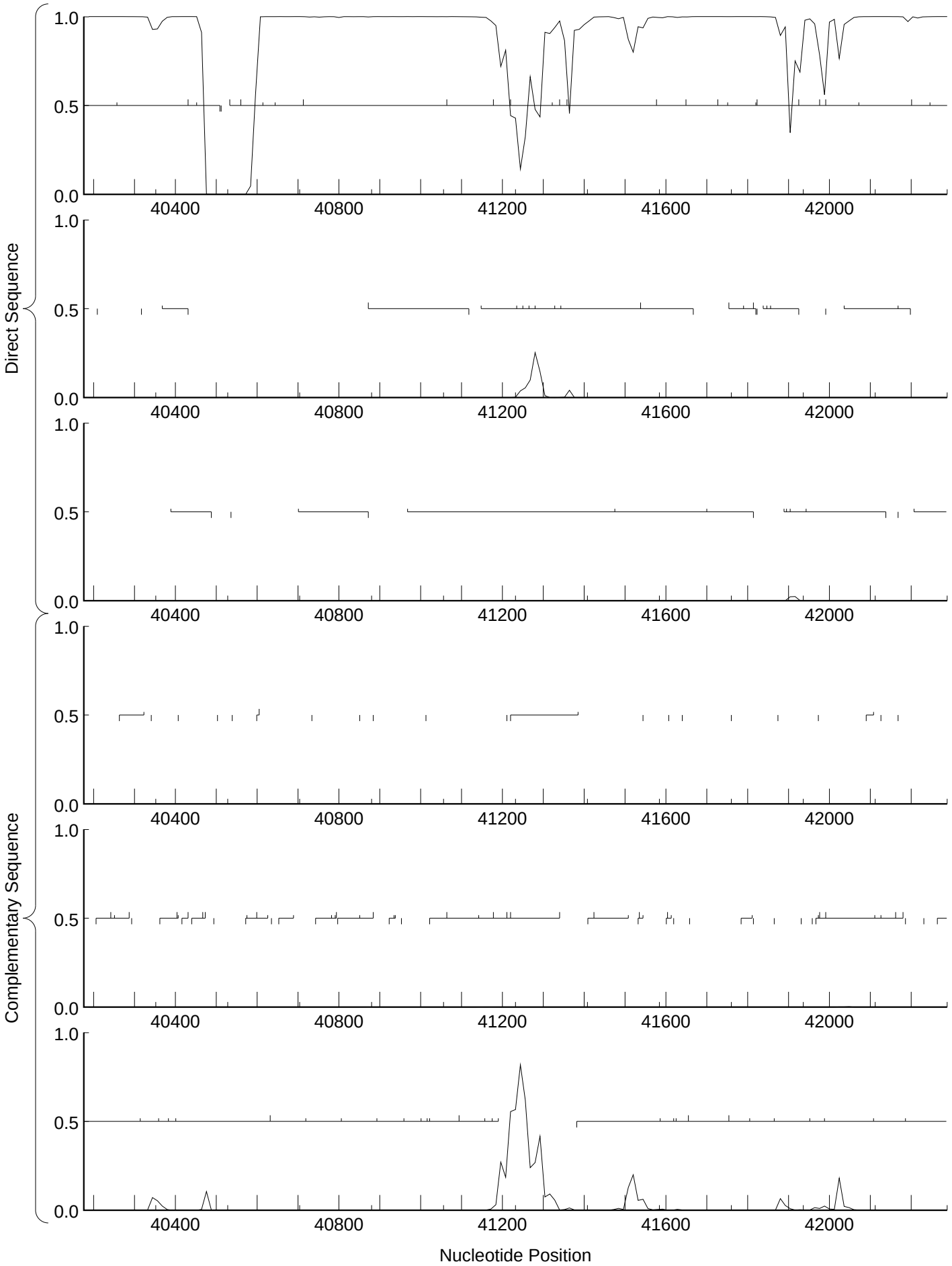
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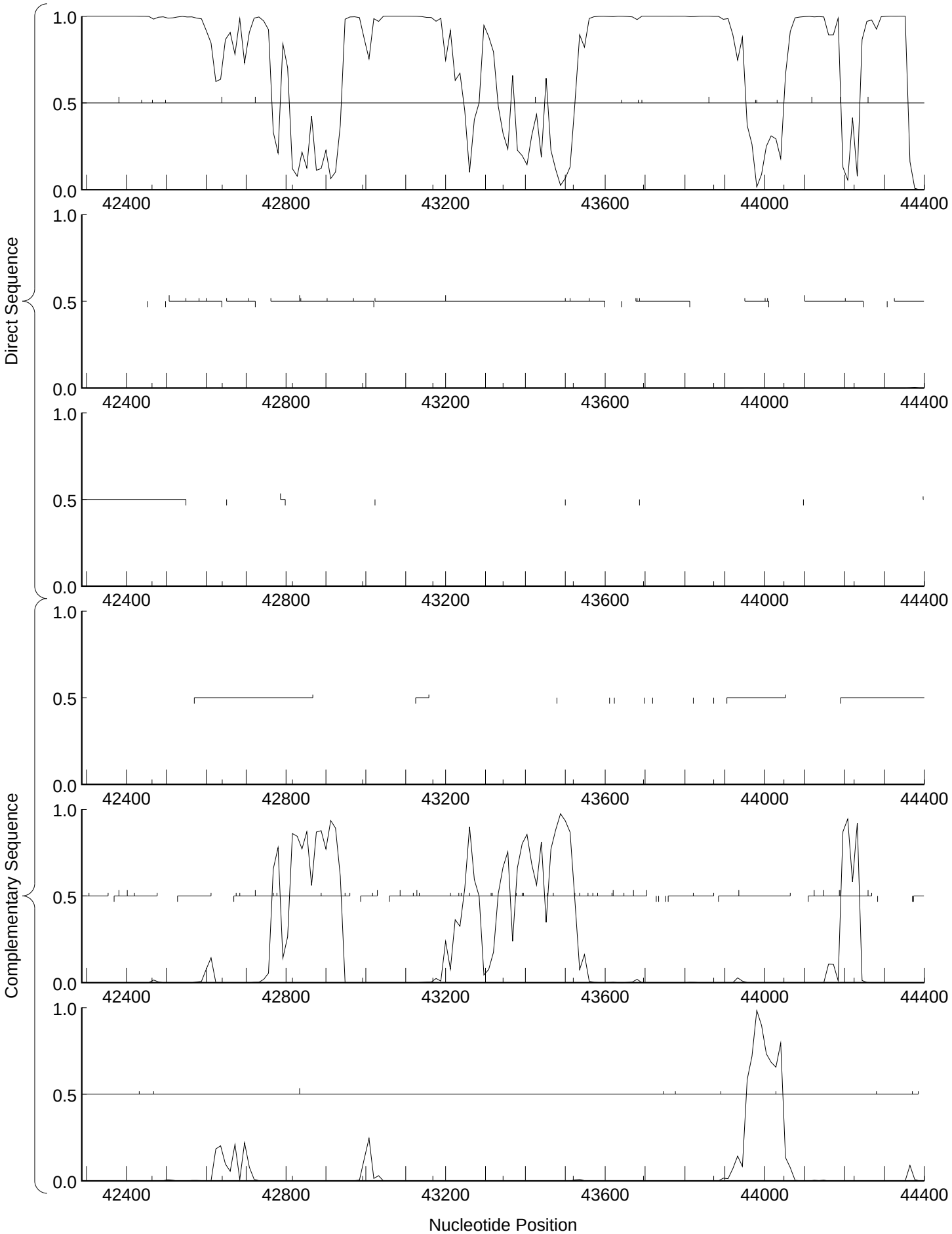
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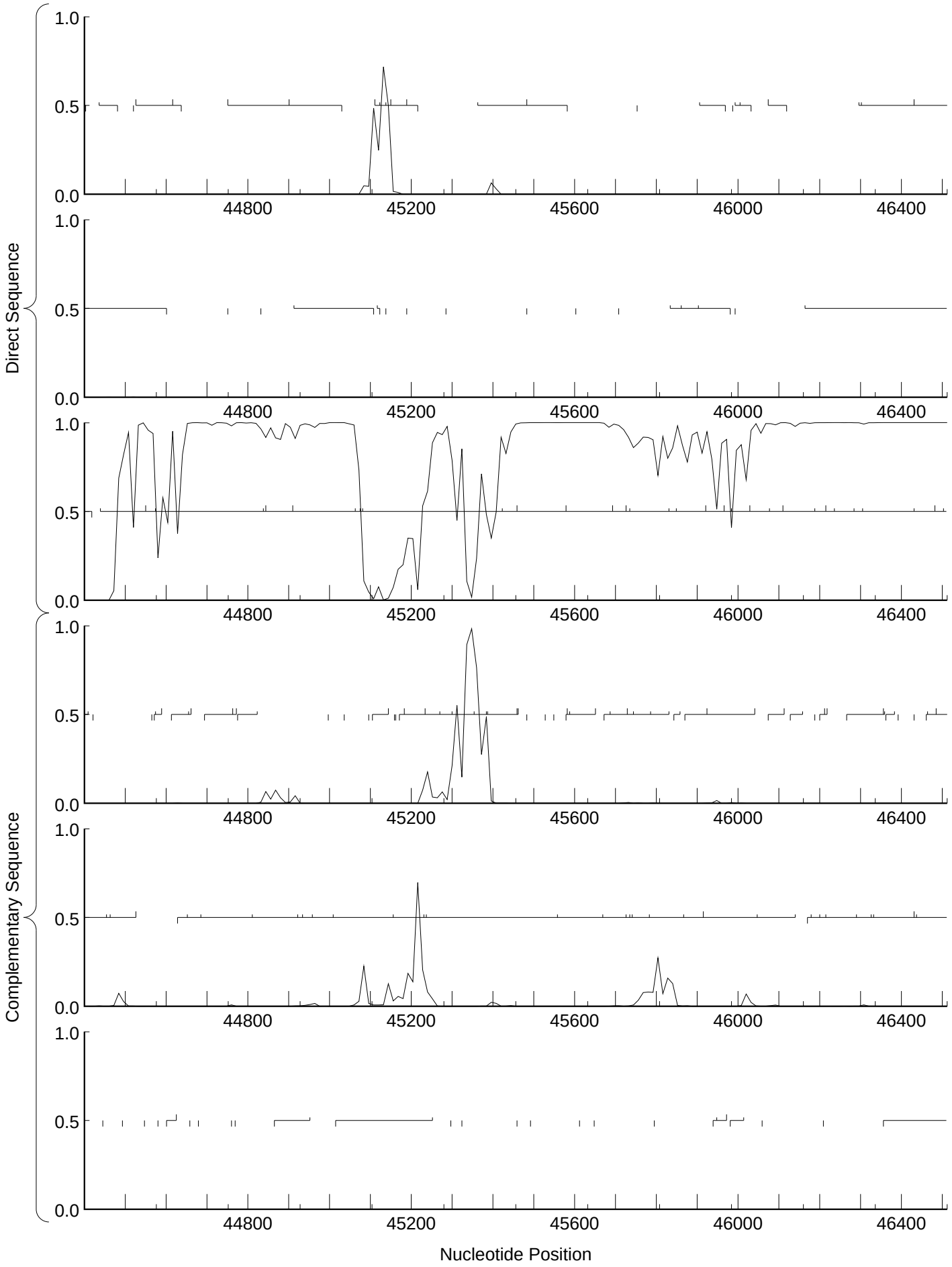
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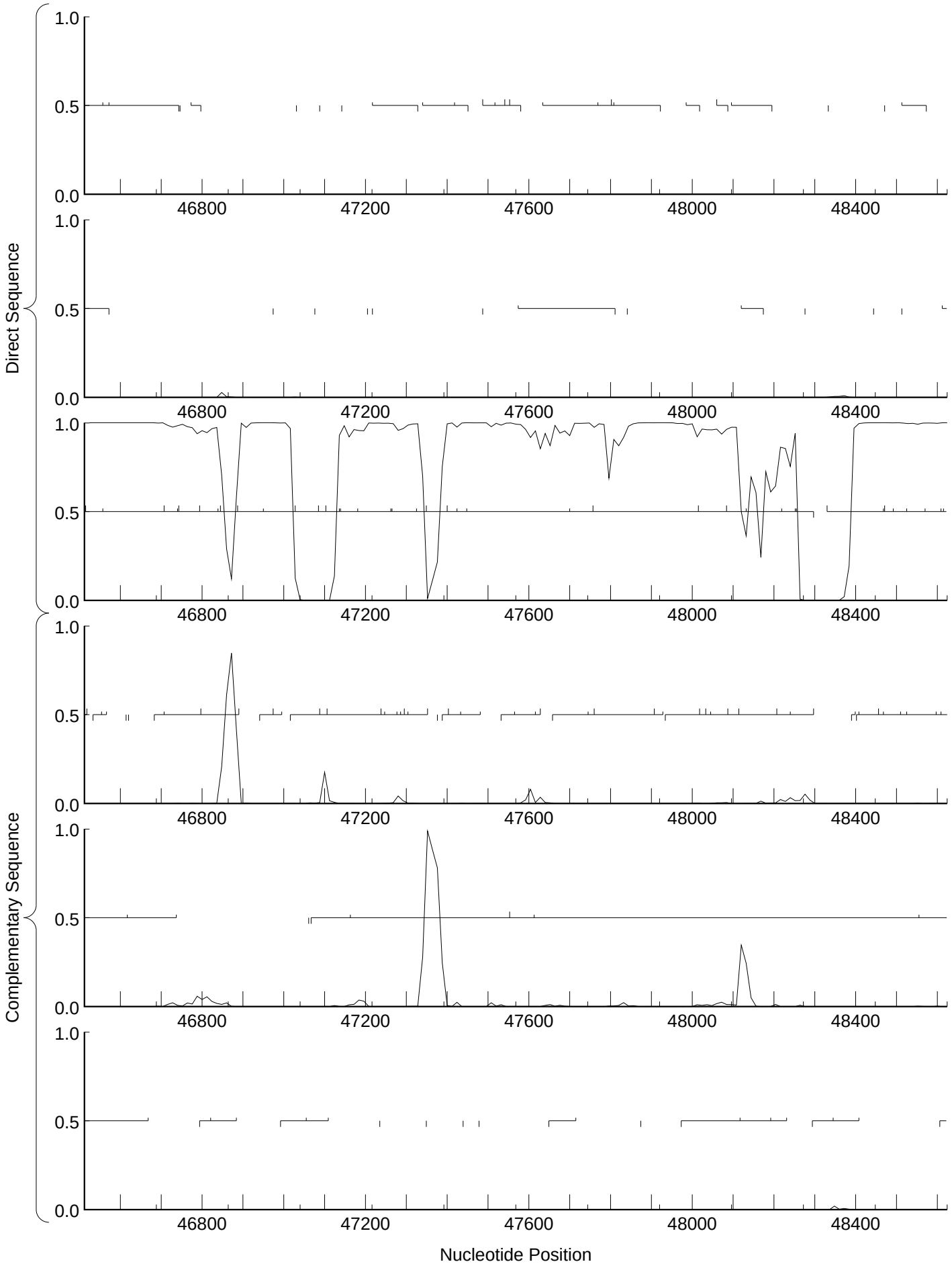
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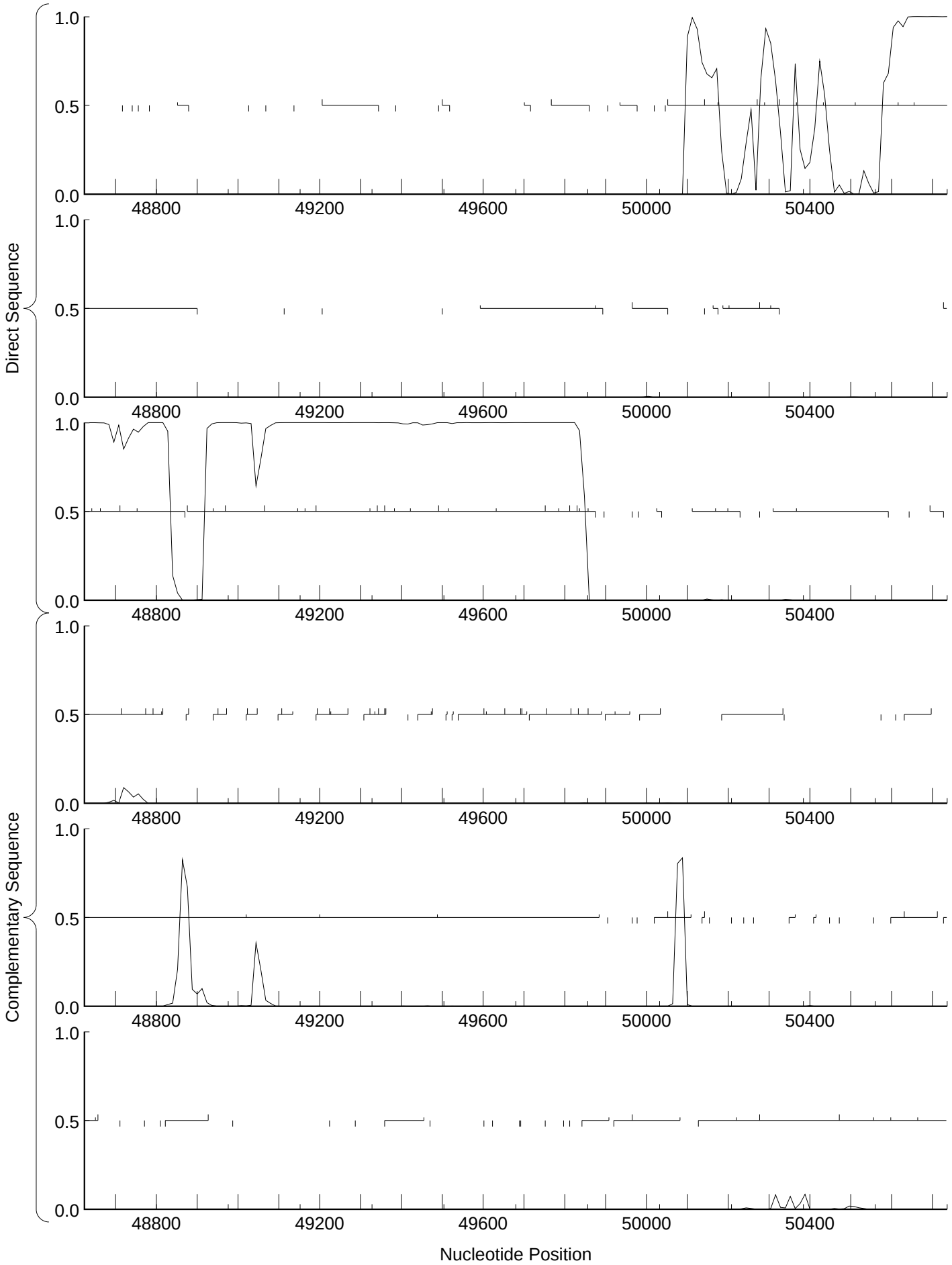
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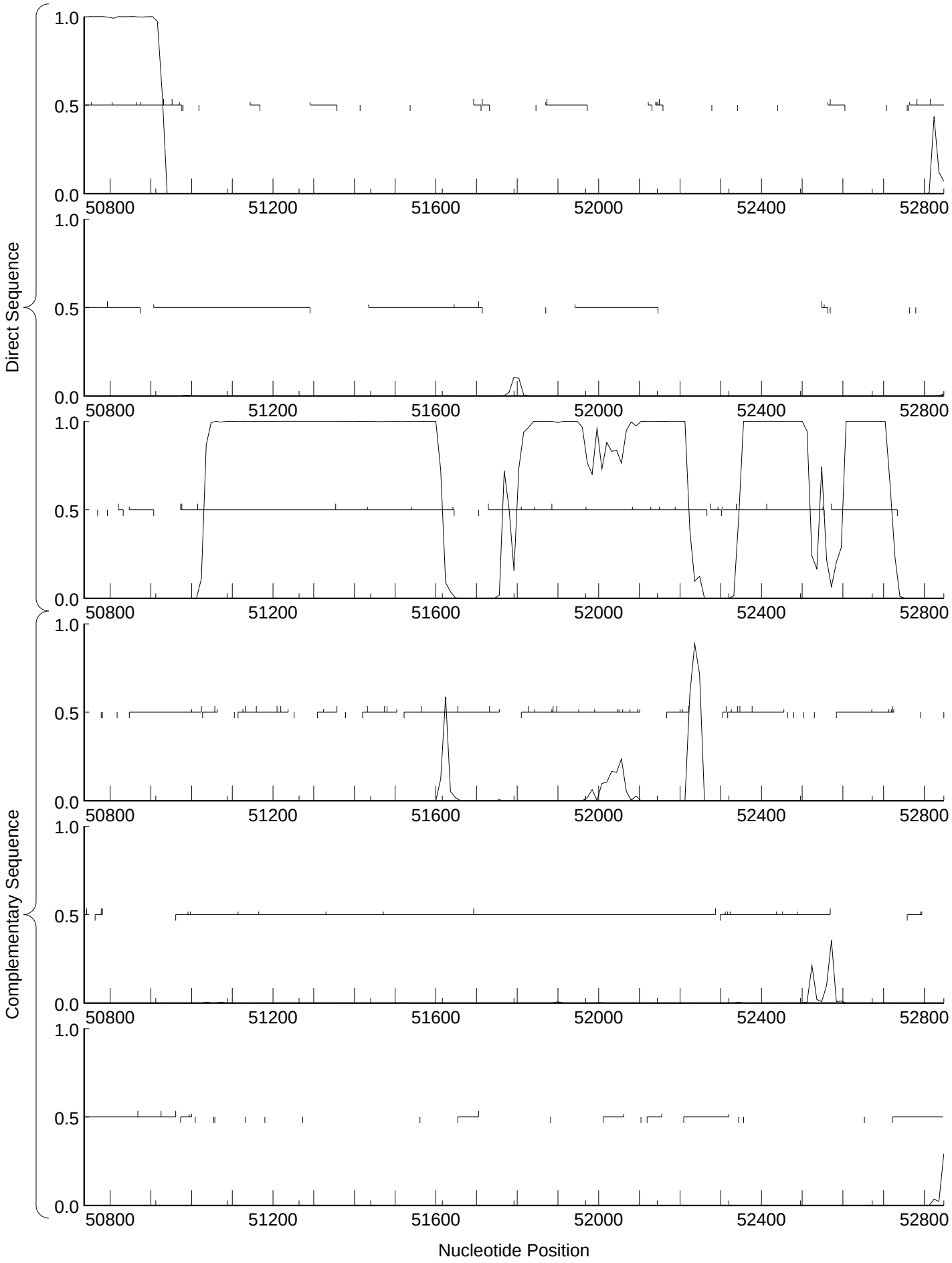
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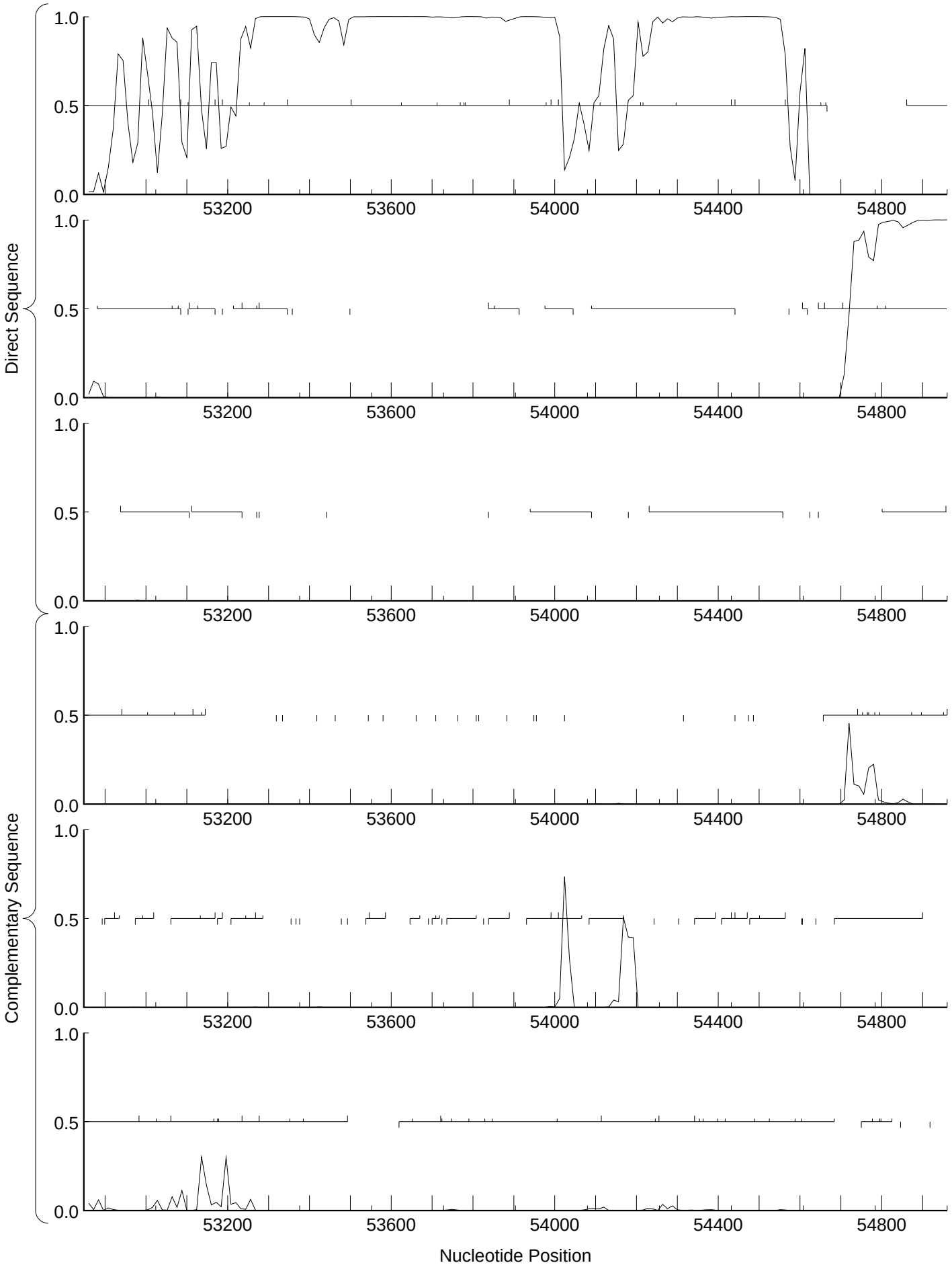
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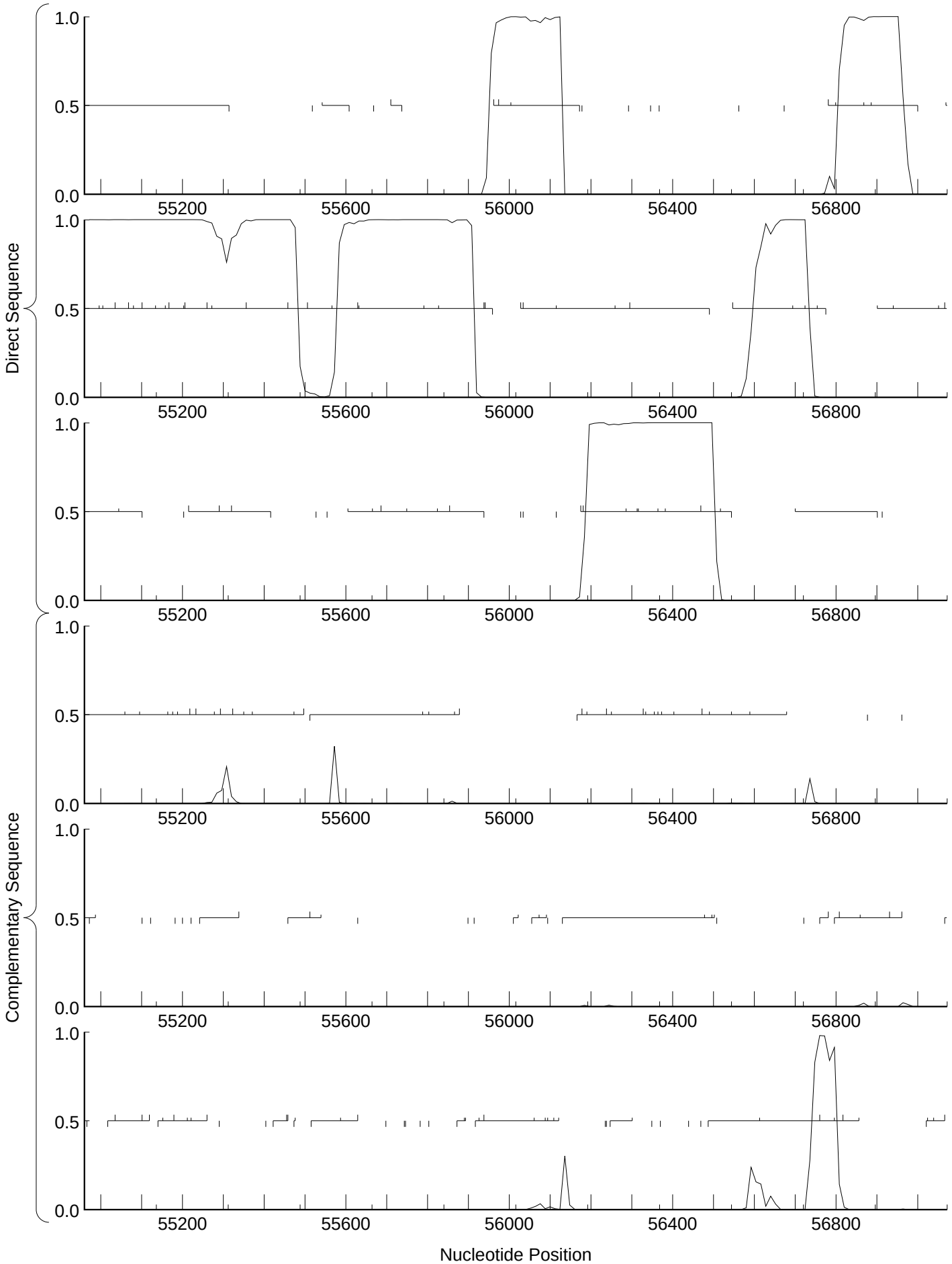
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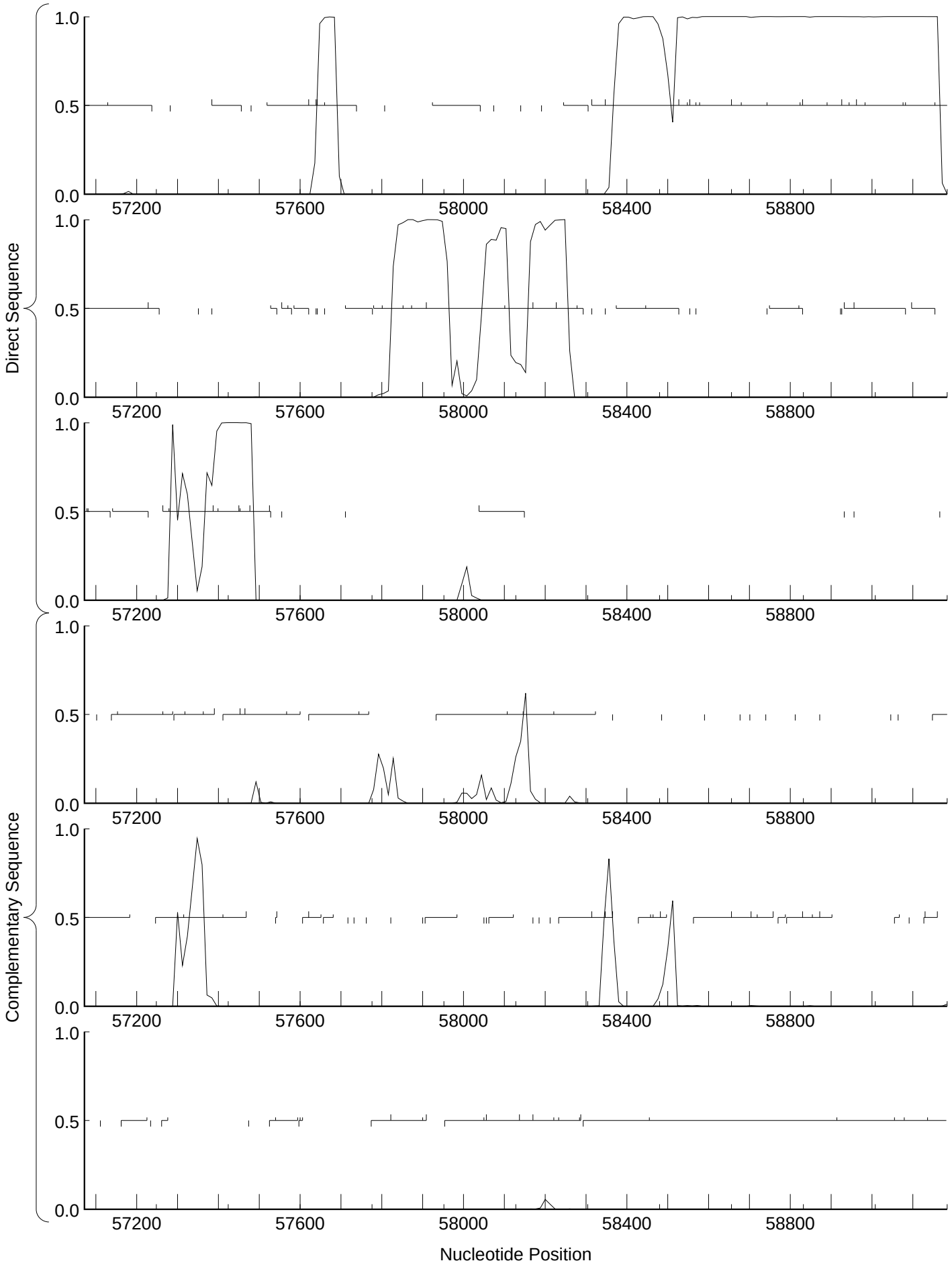
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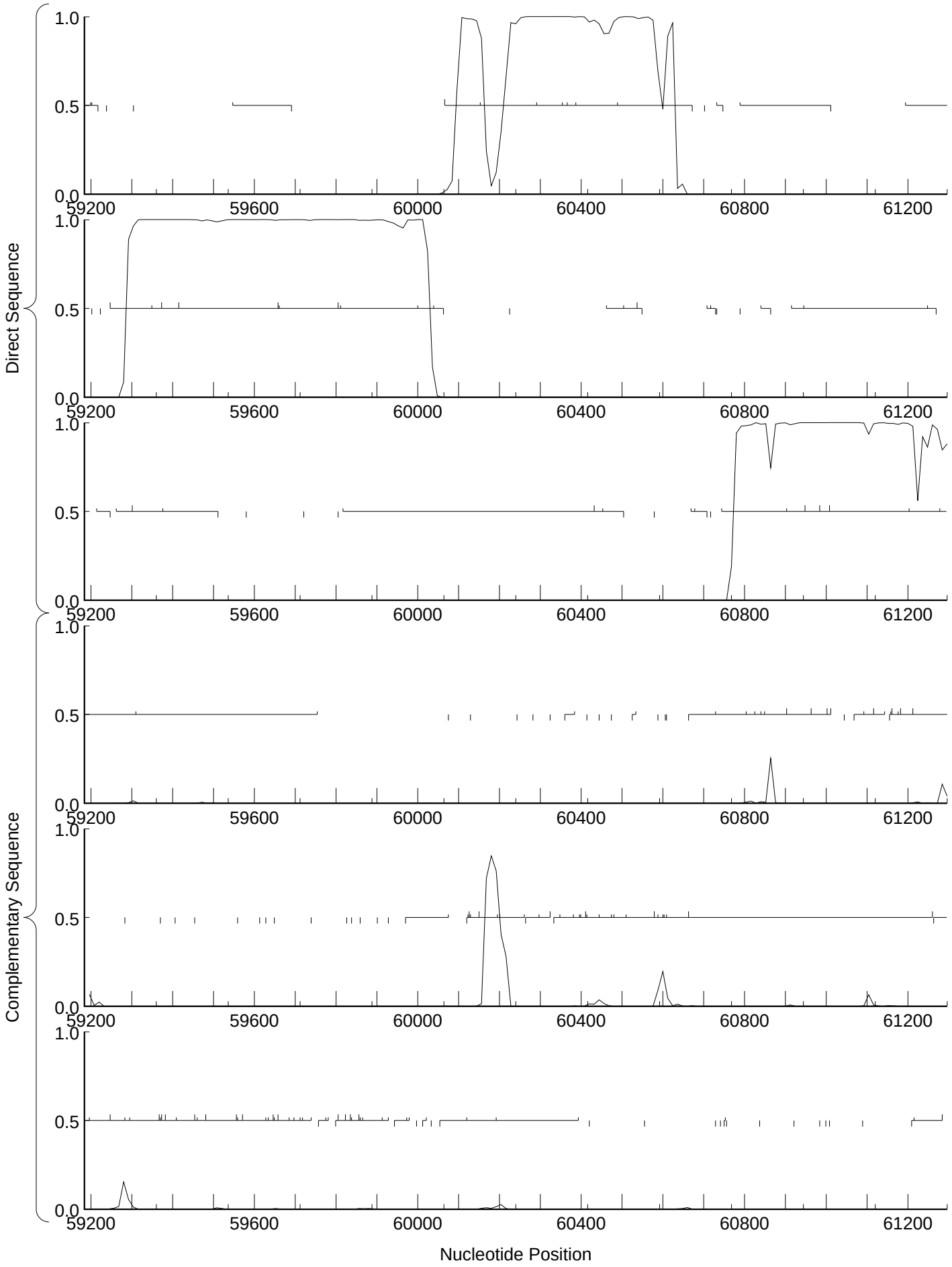
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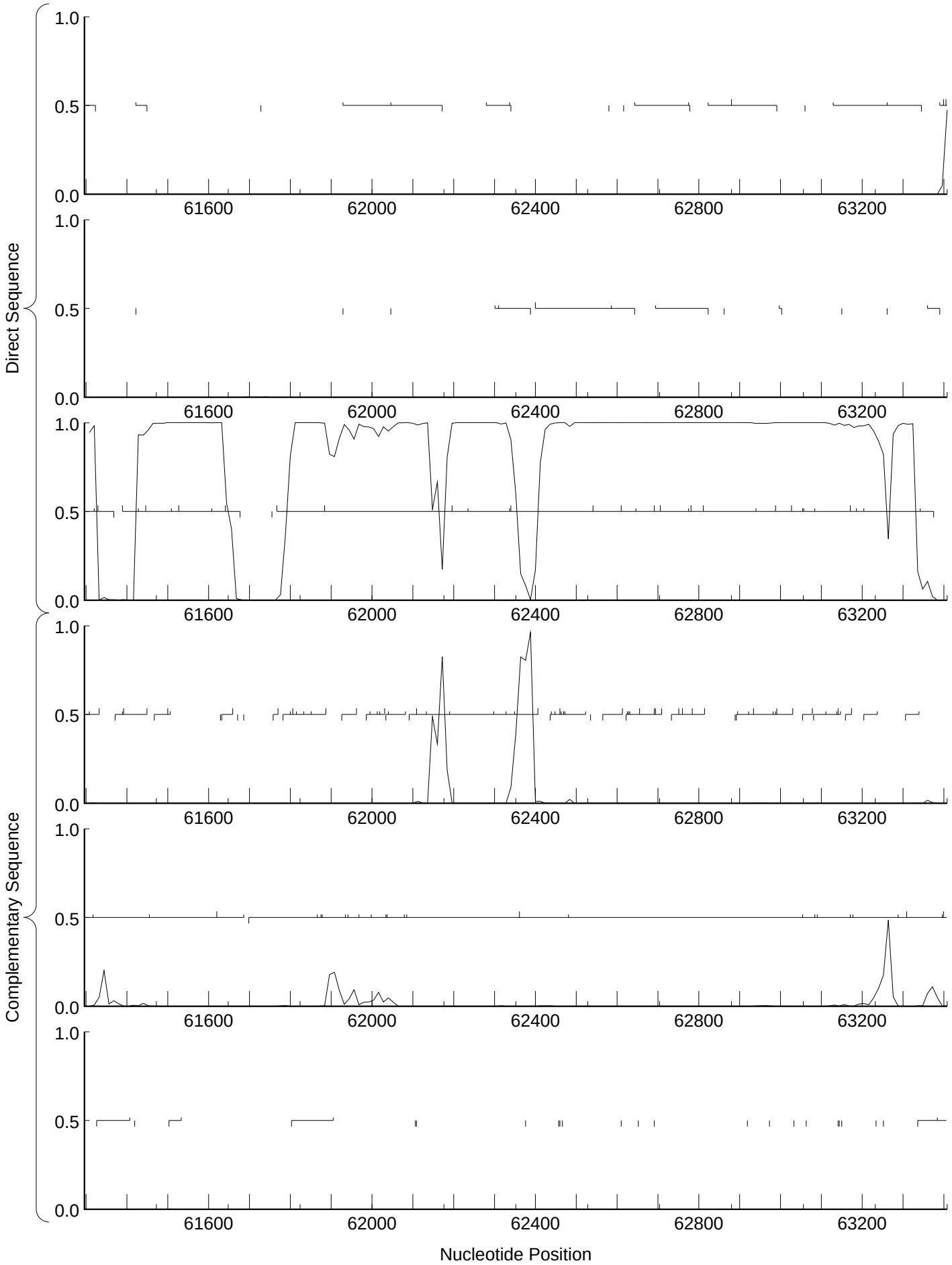
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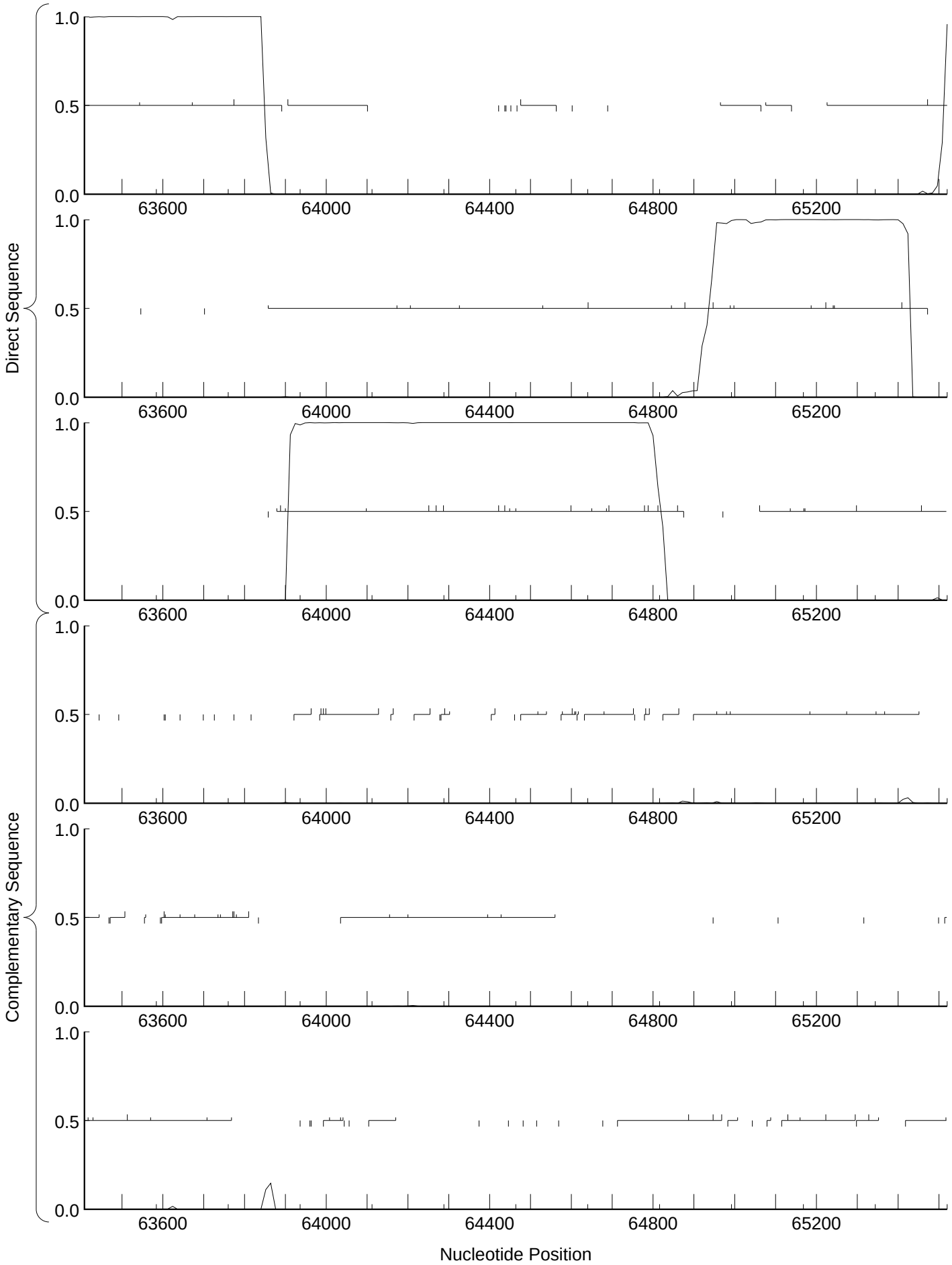
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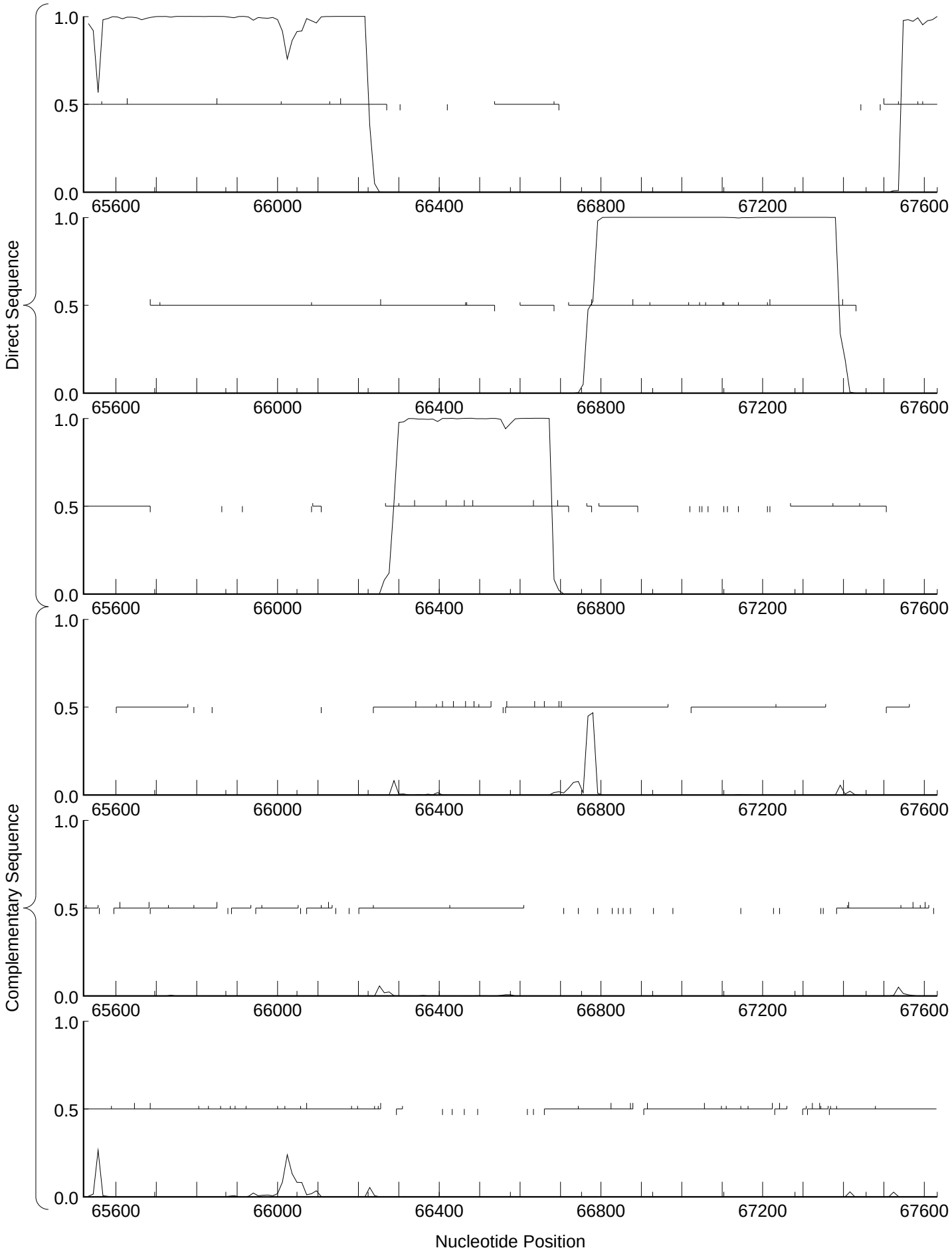
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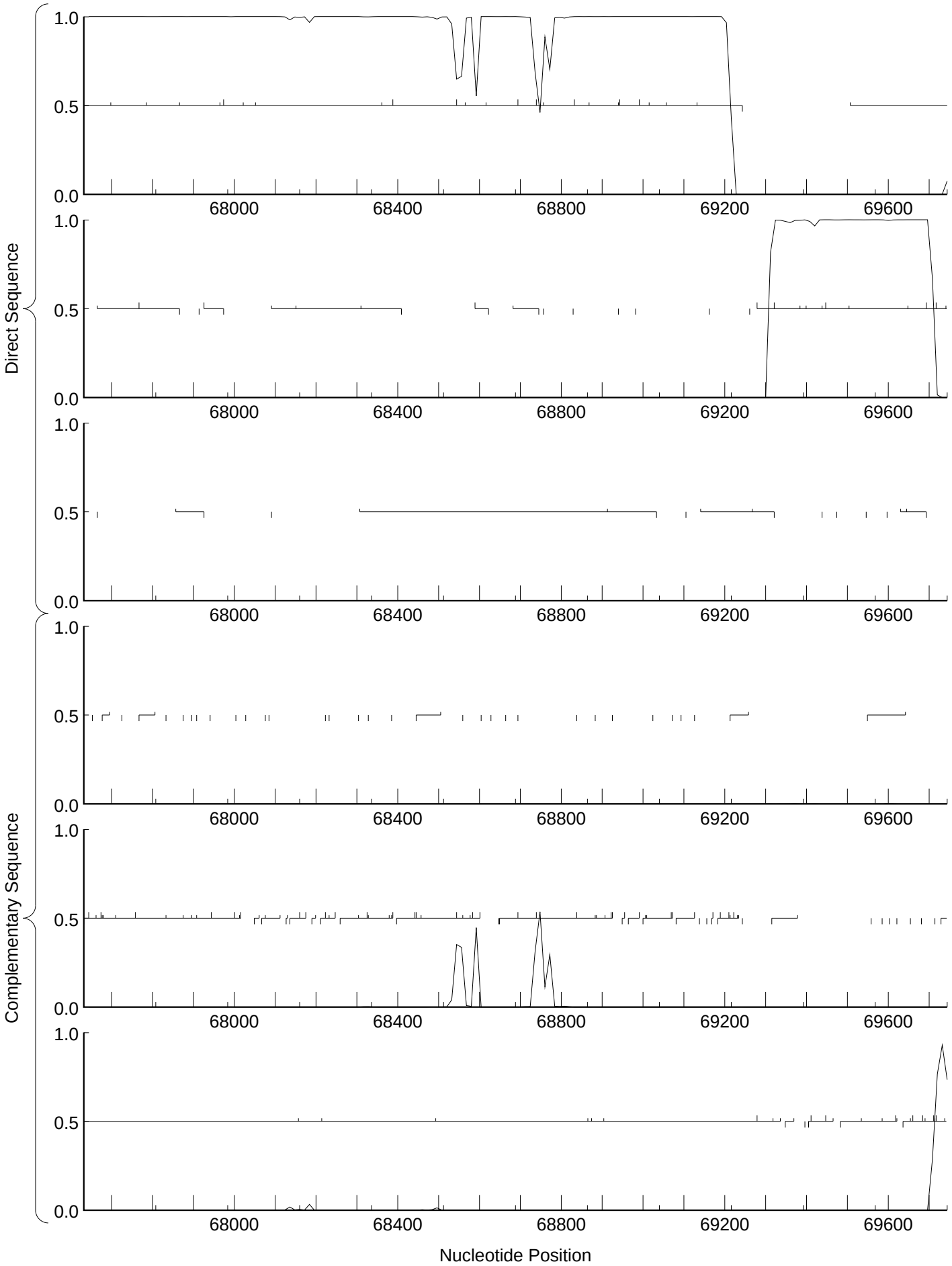
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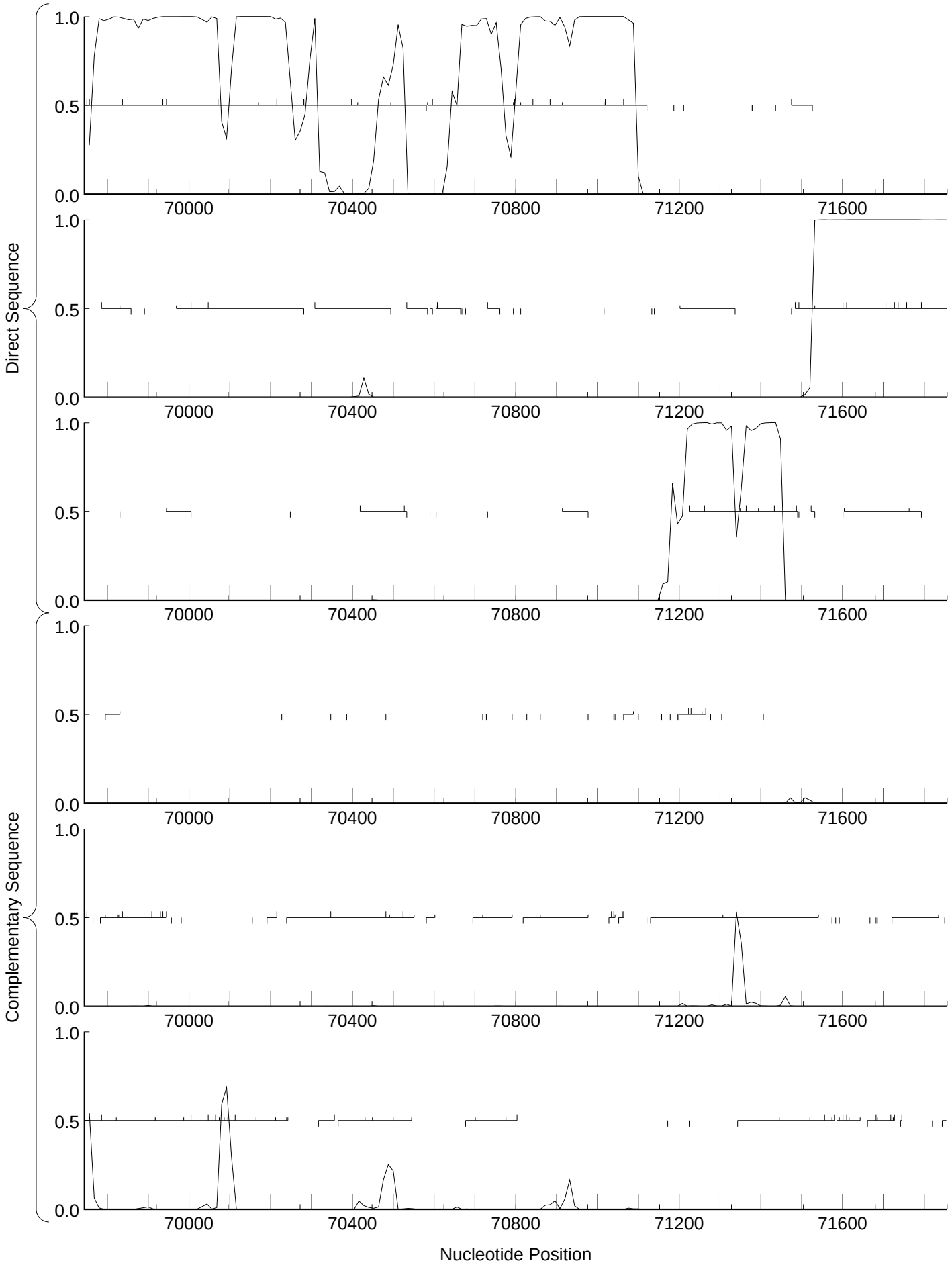
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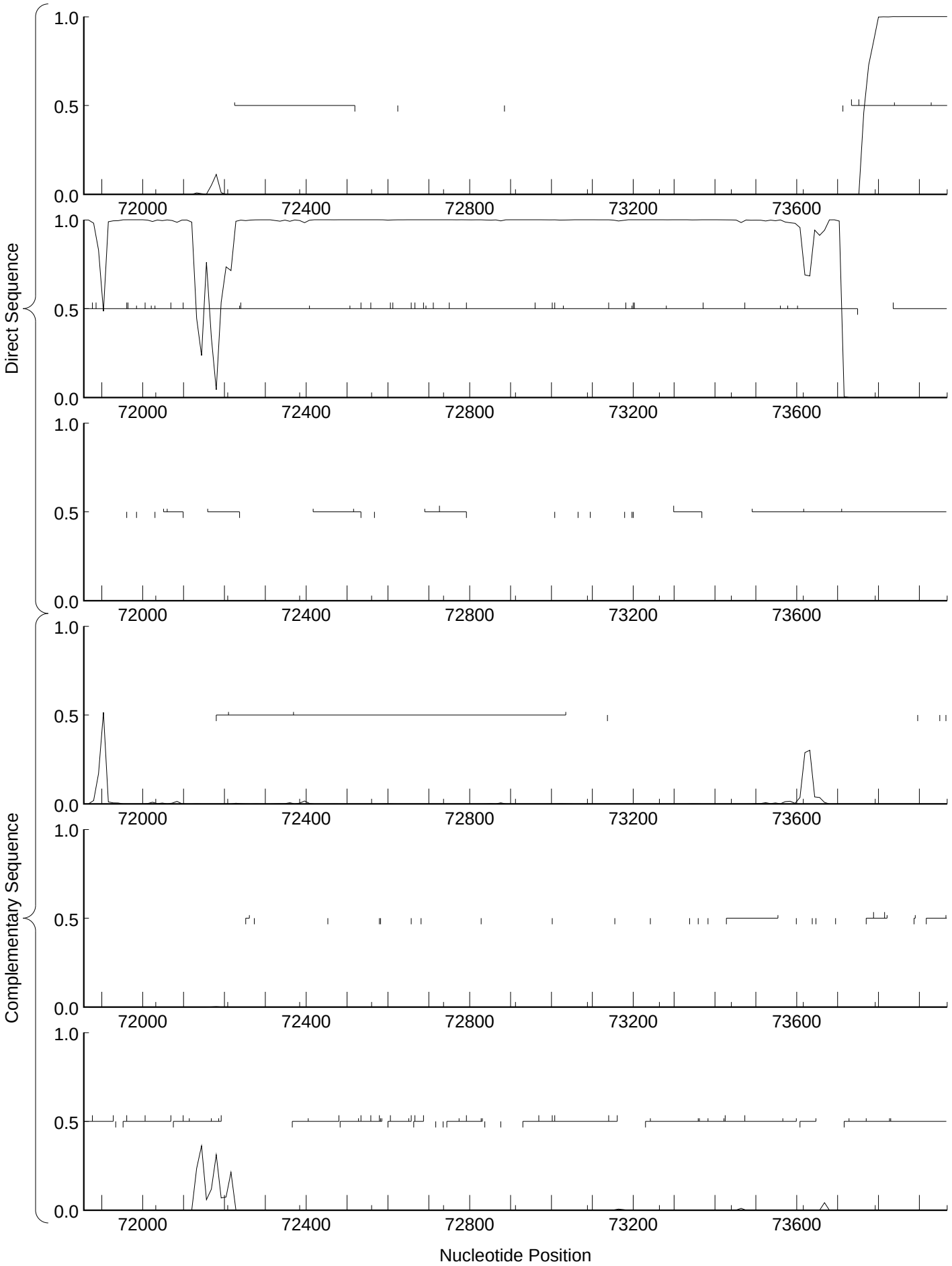
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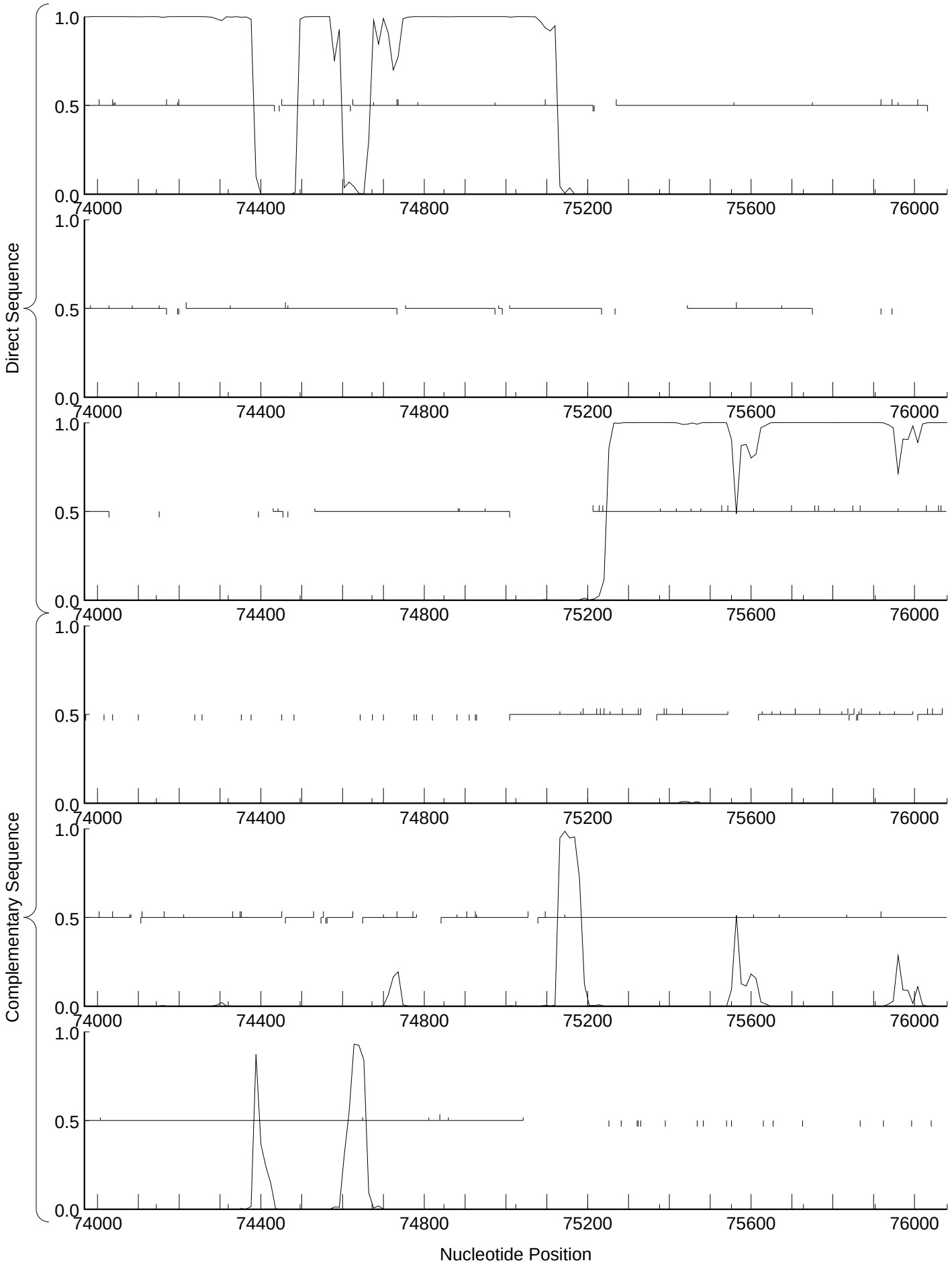
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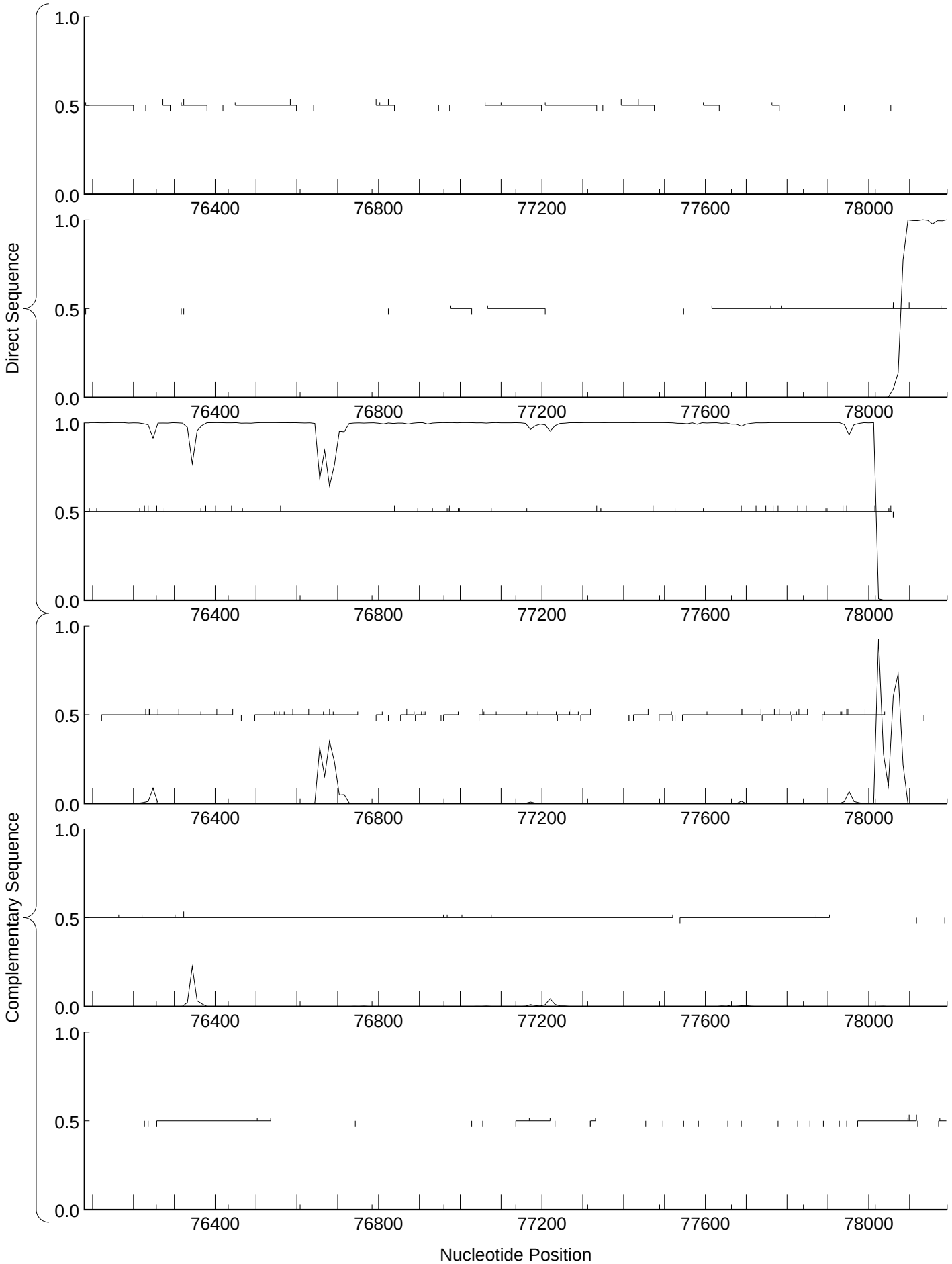
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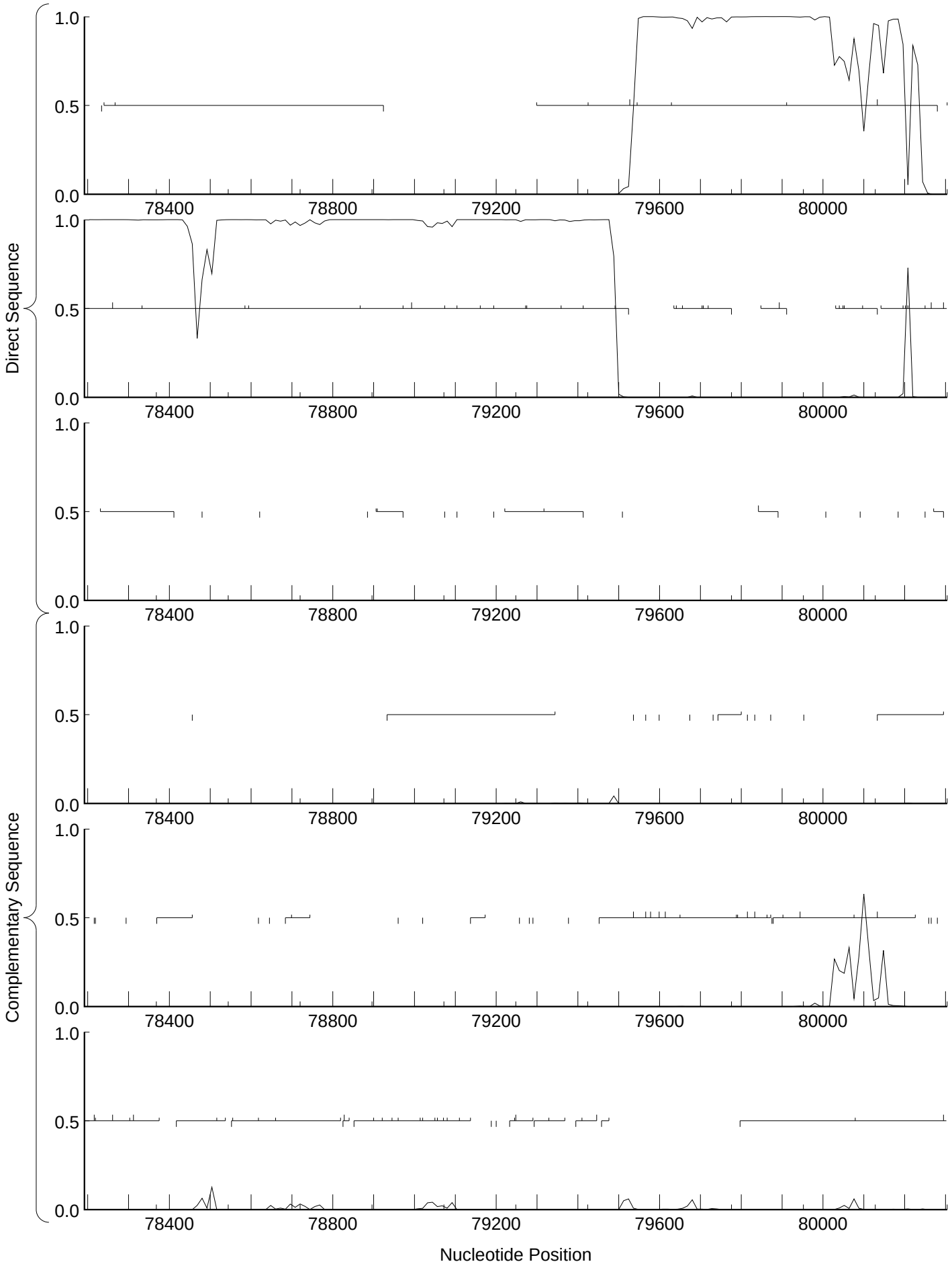
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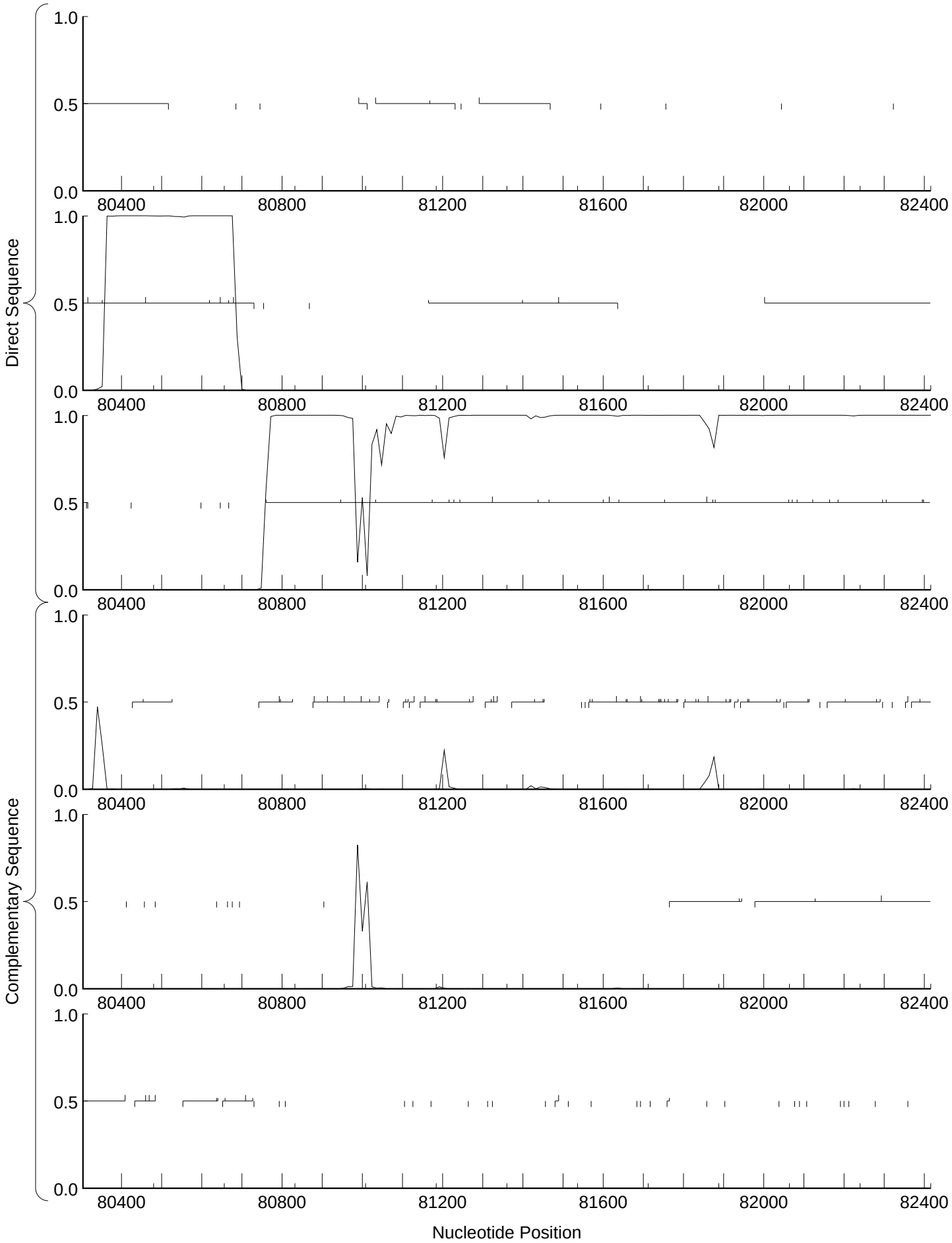
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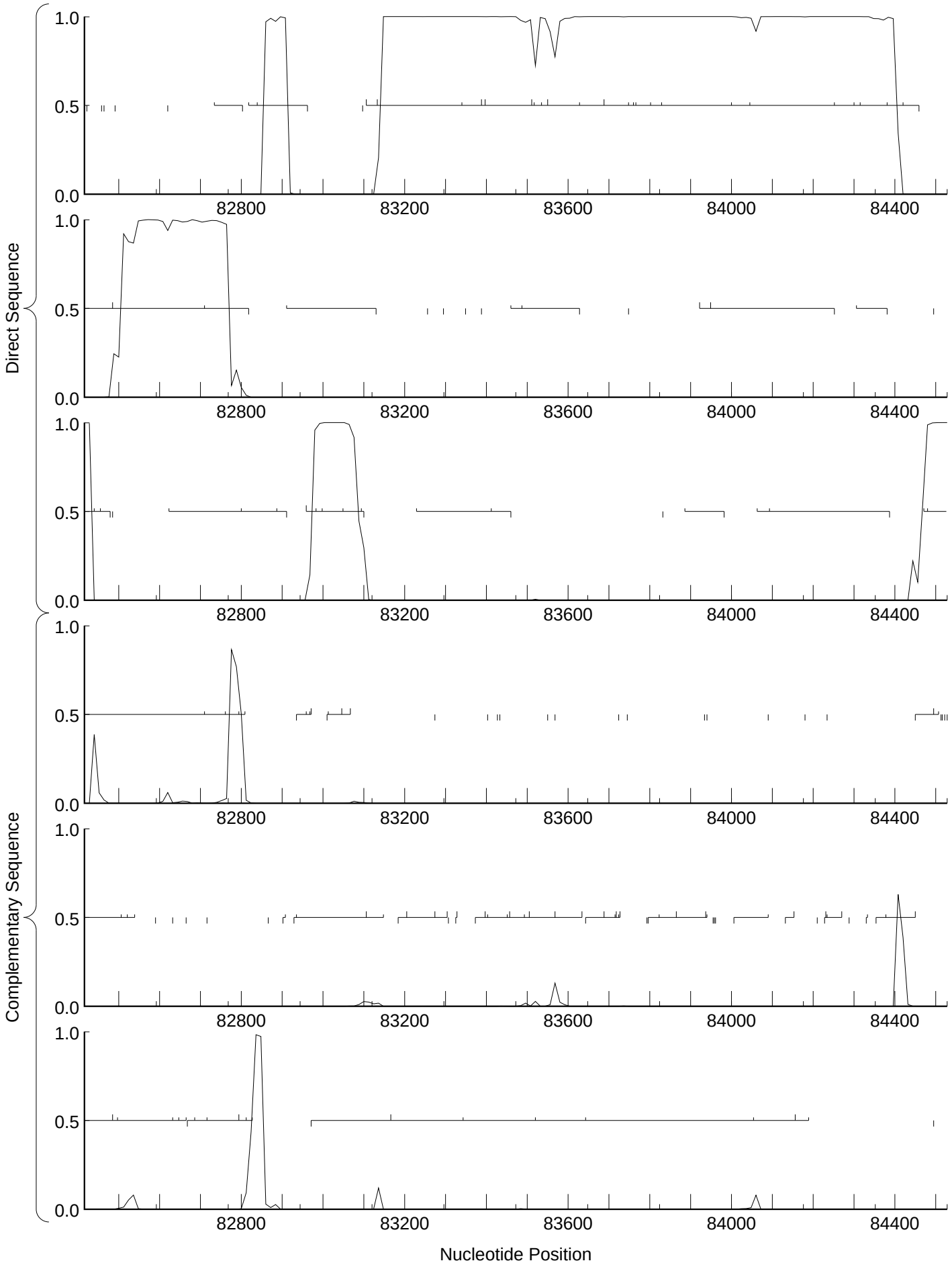
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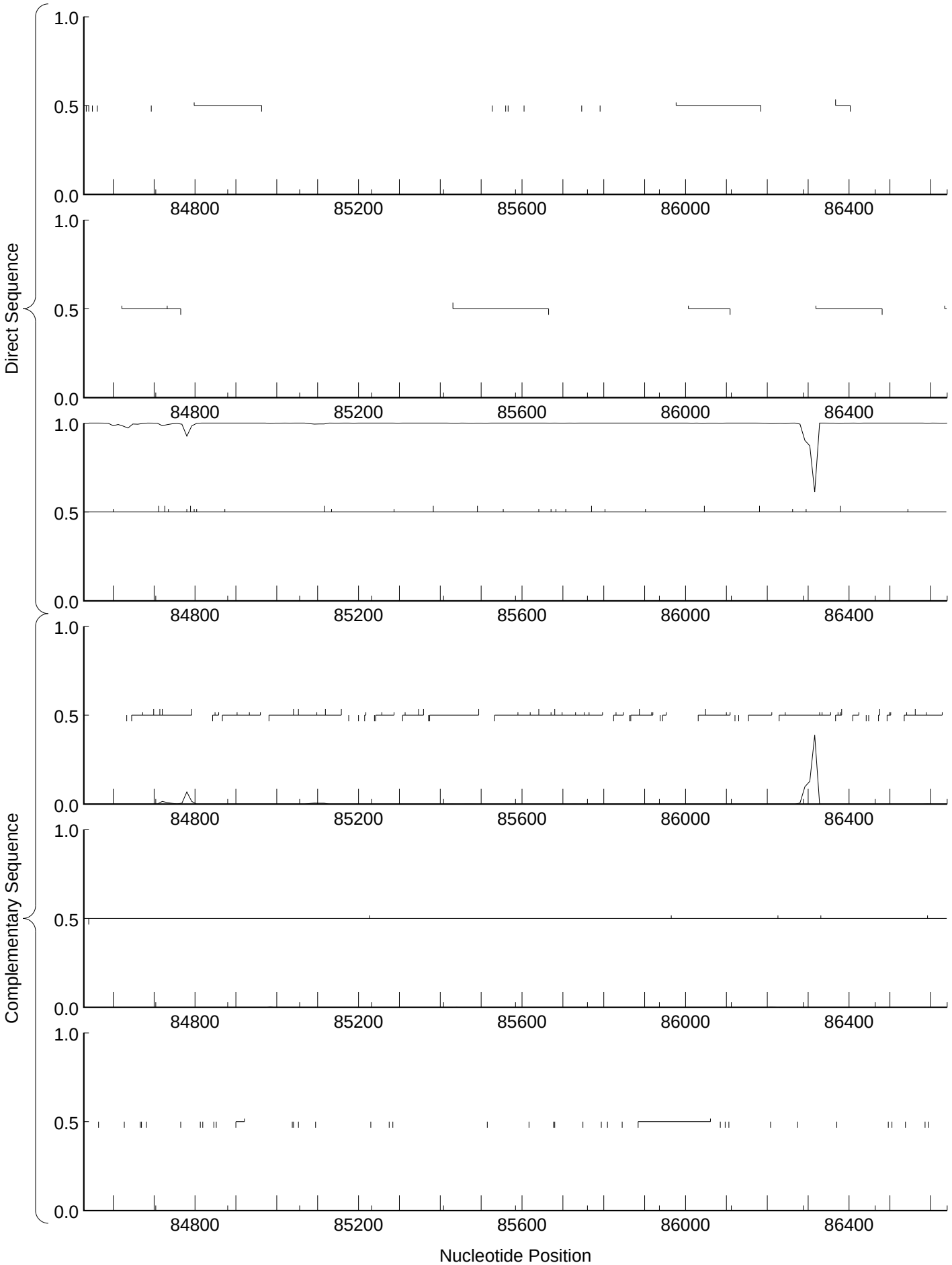
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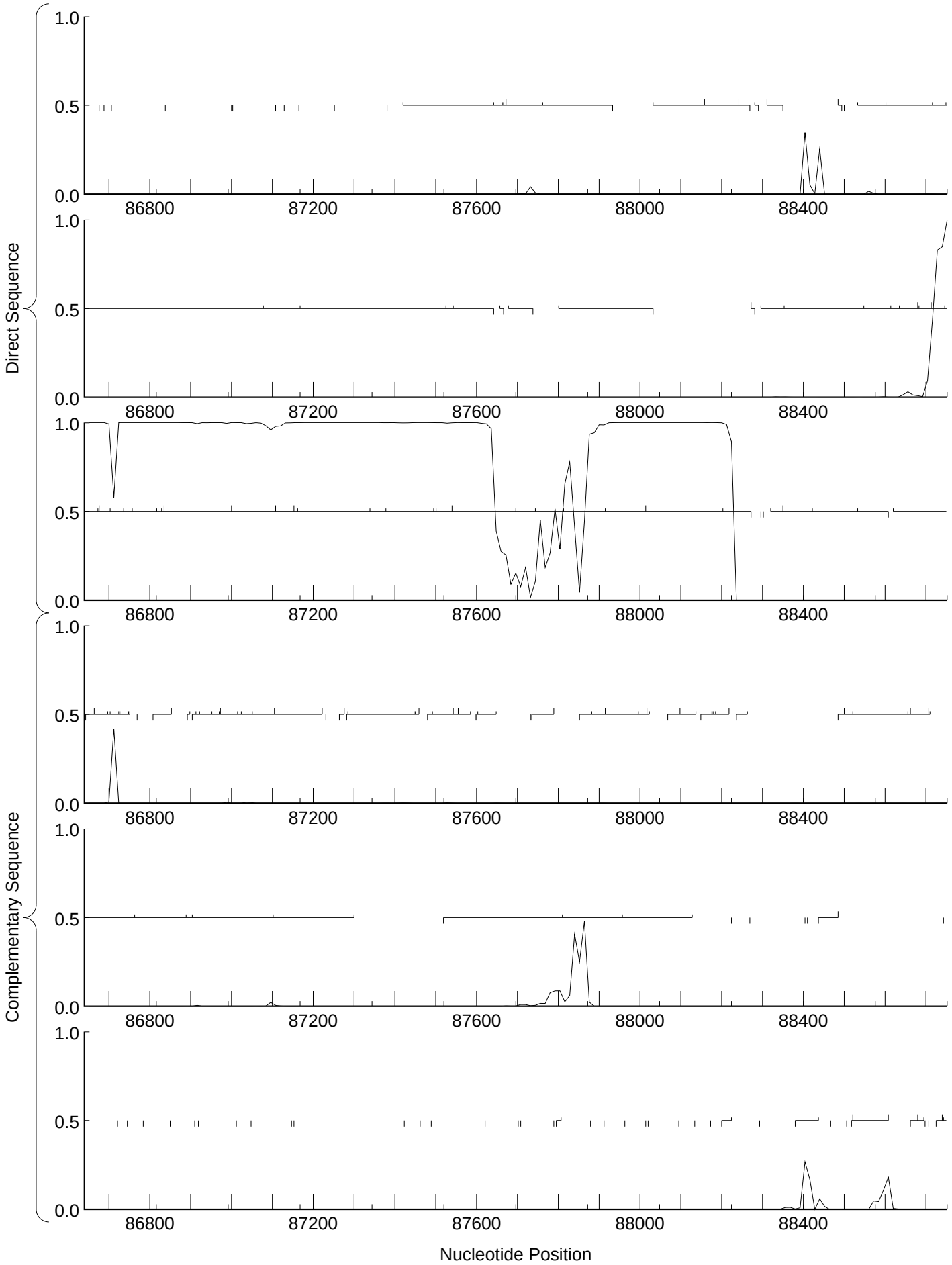
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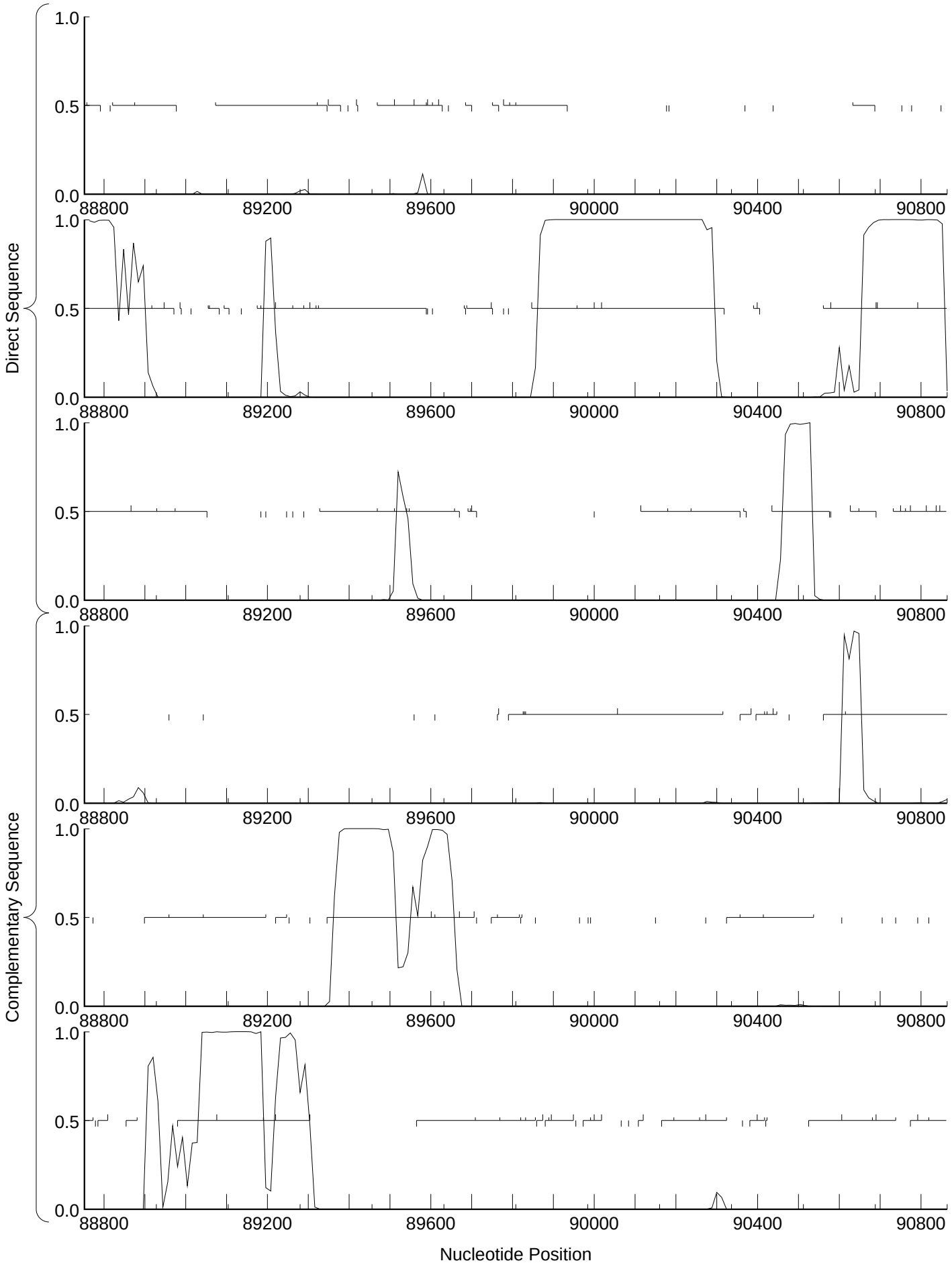
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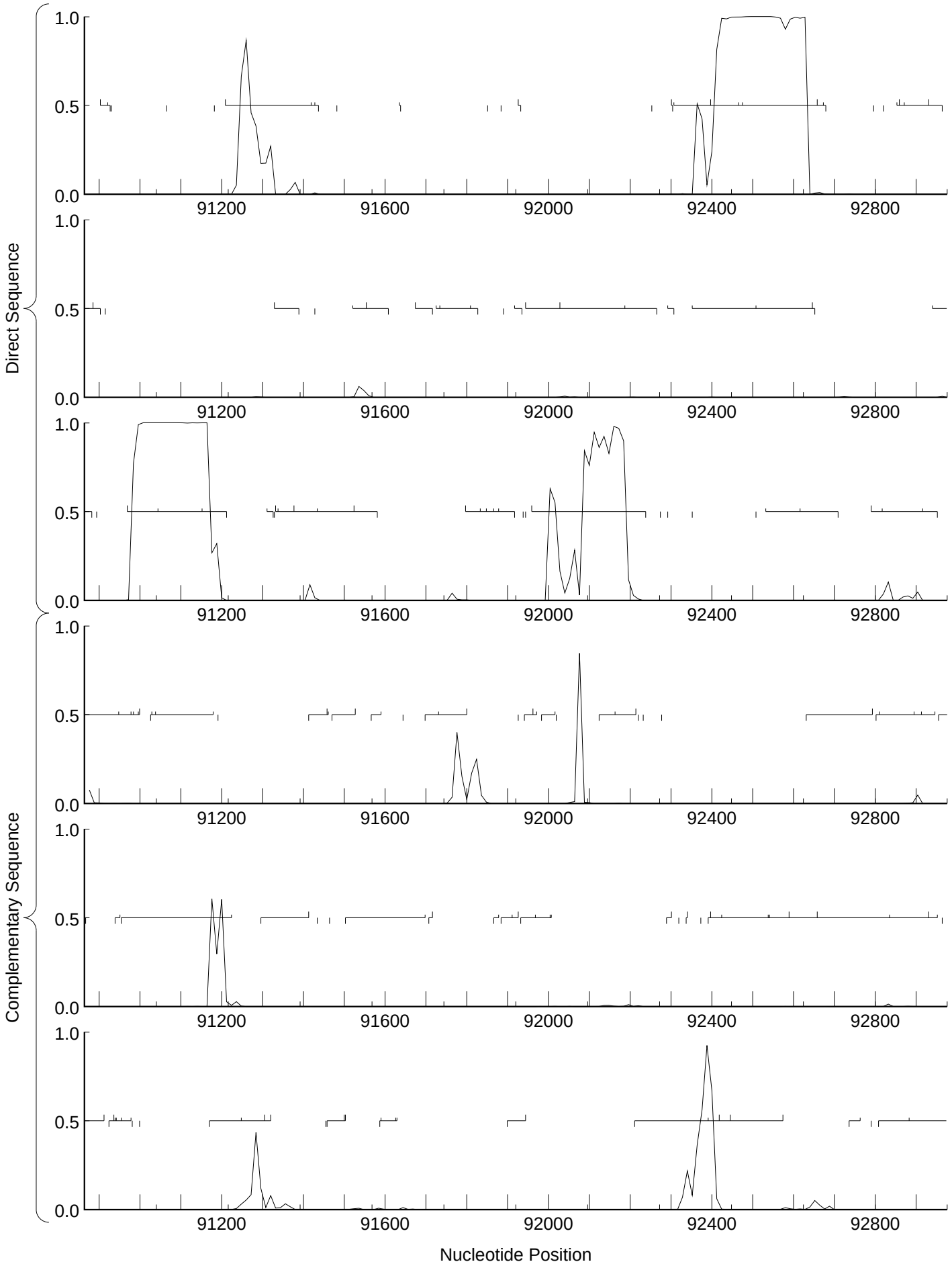
ycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 43/76



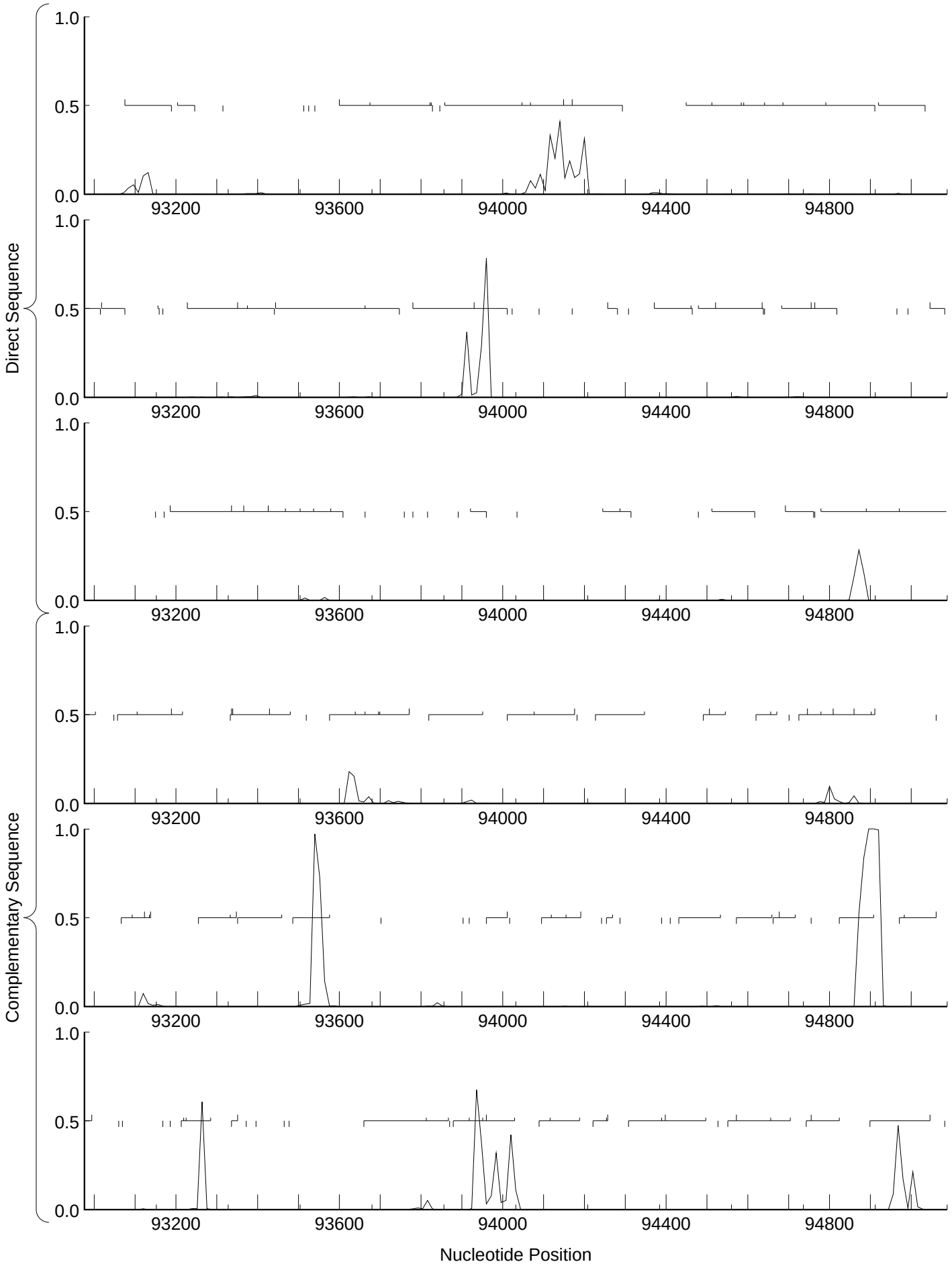
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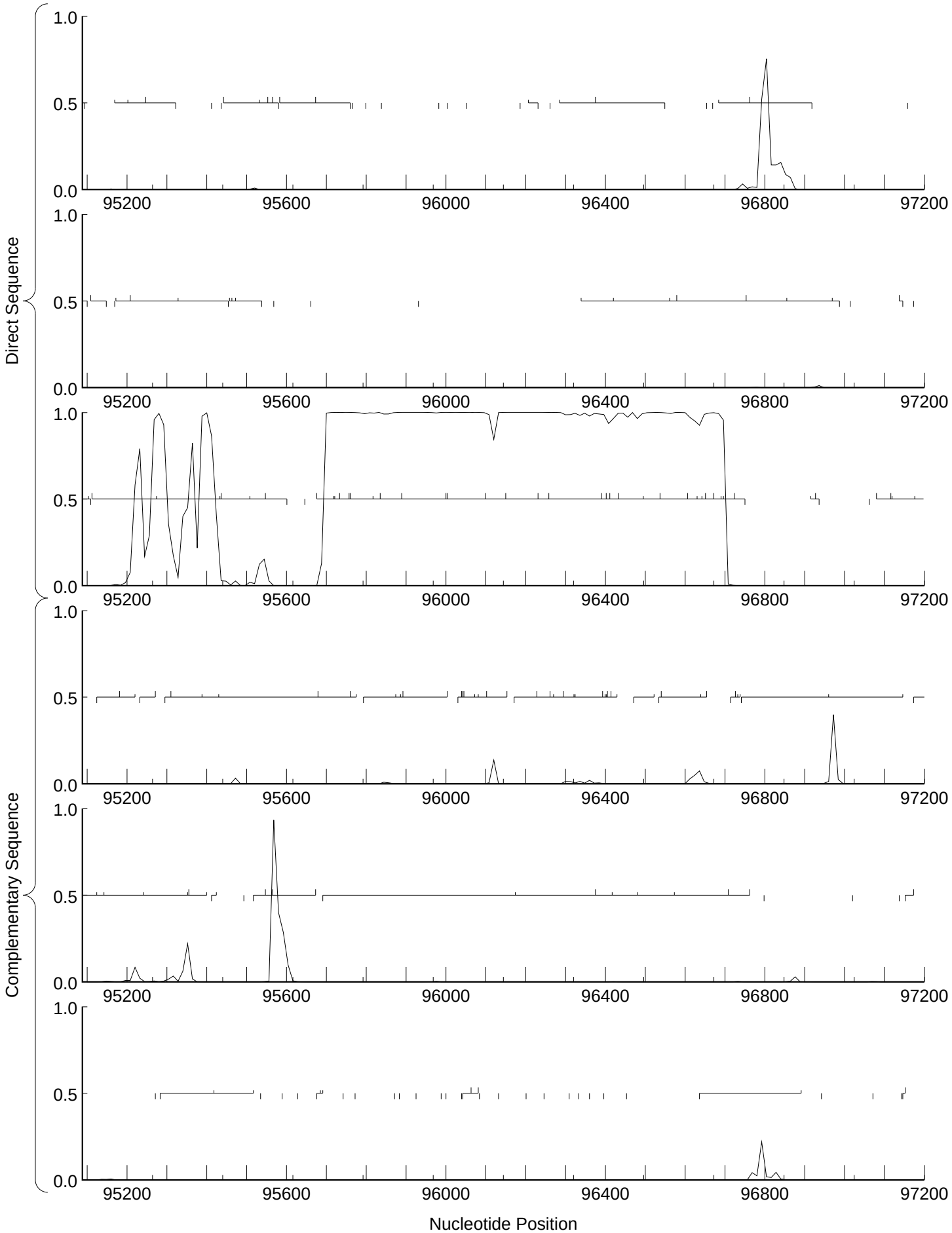
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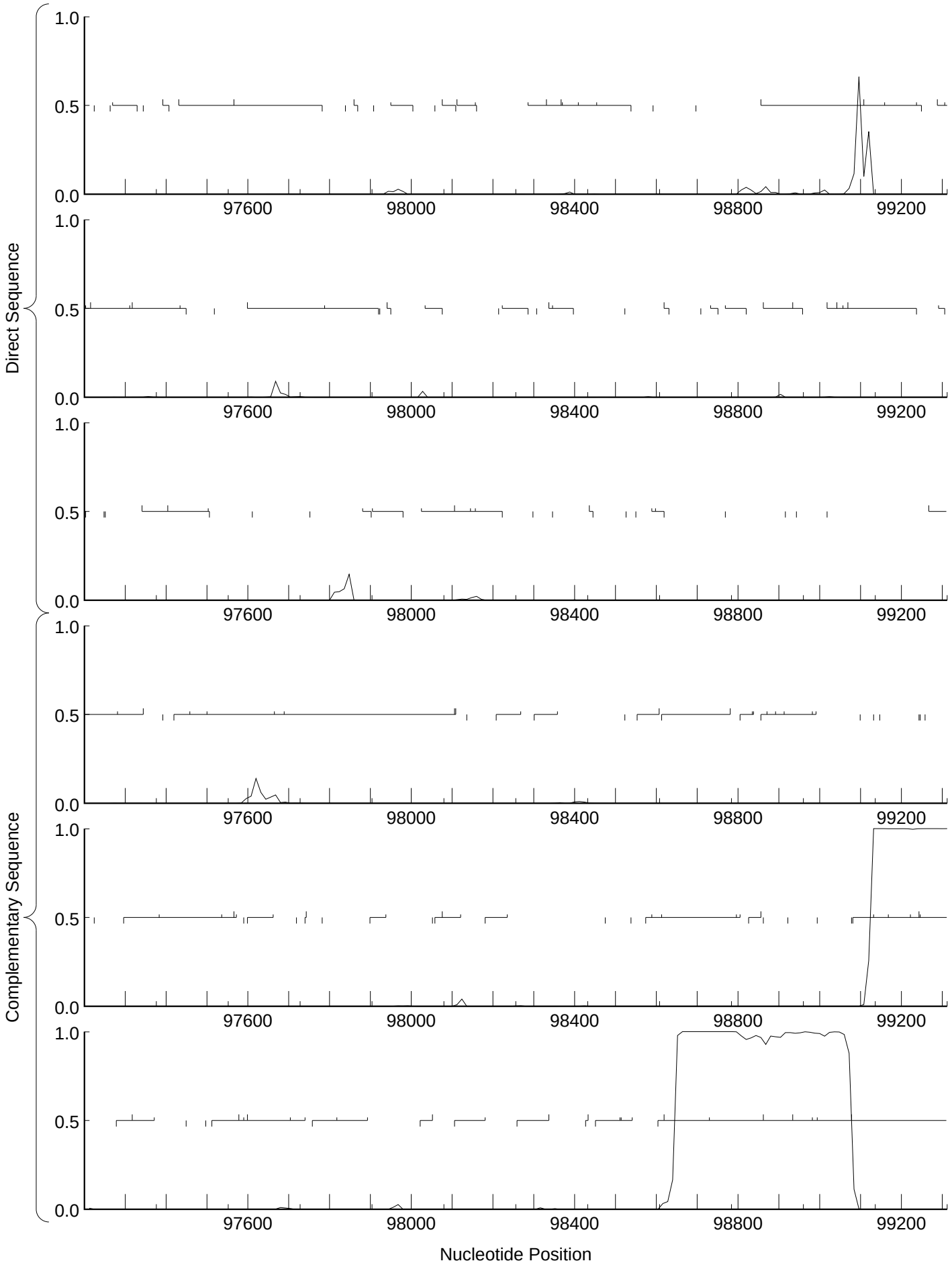
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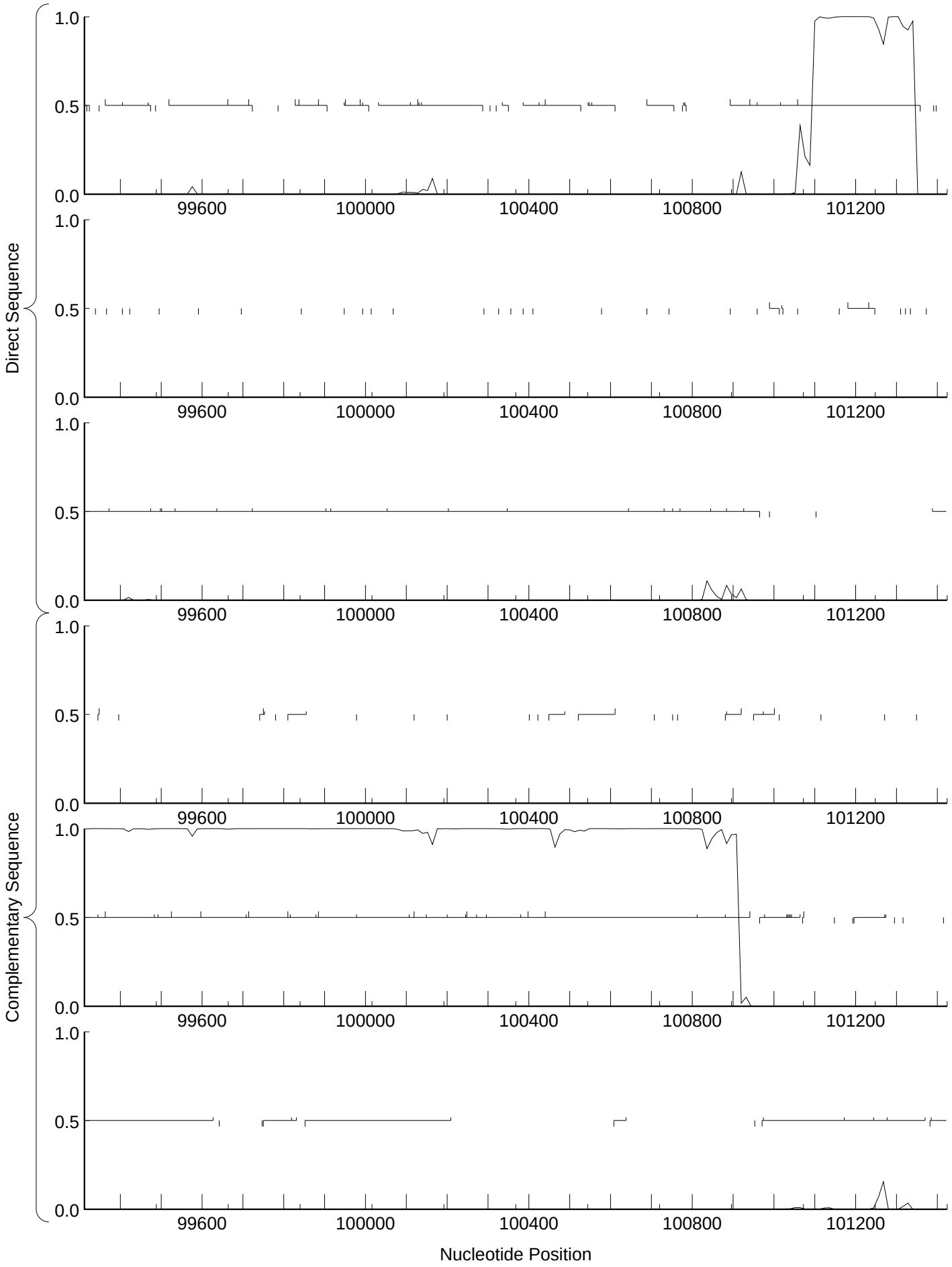
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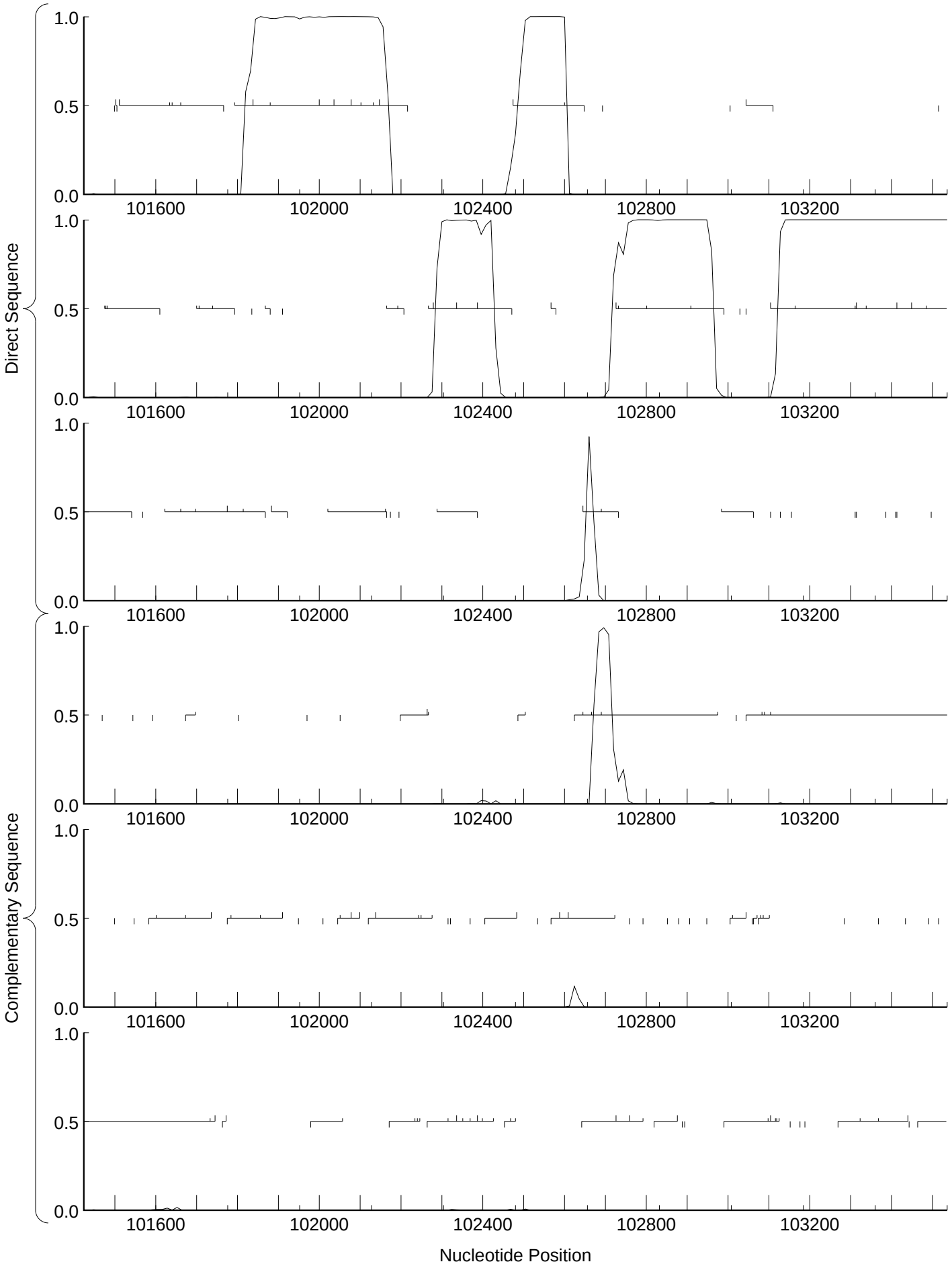
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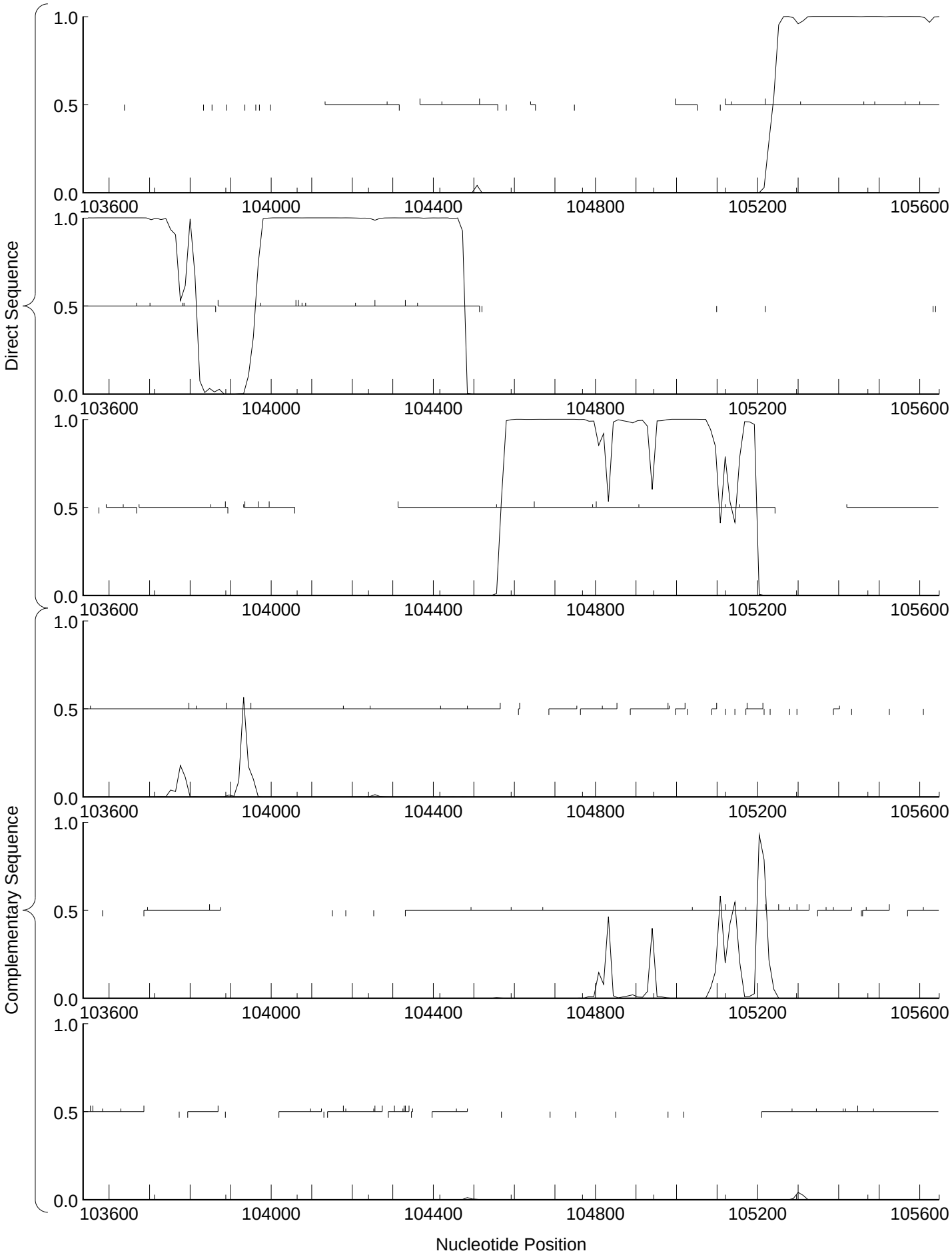
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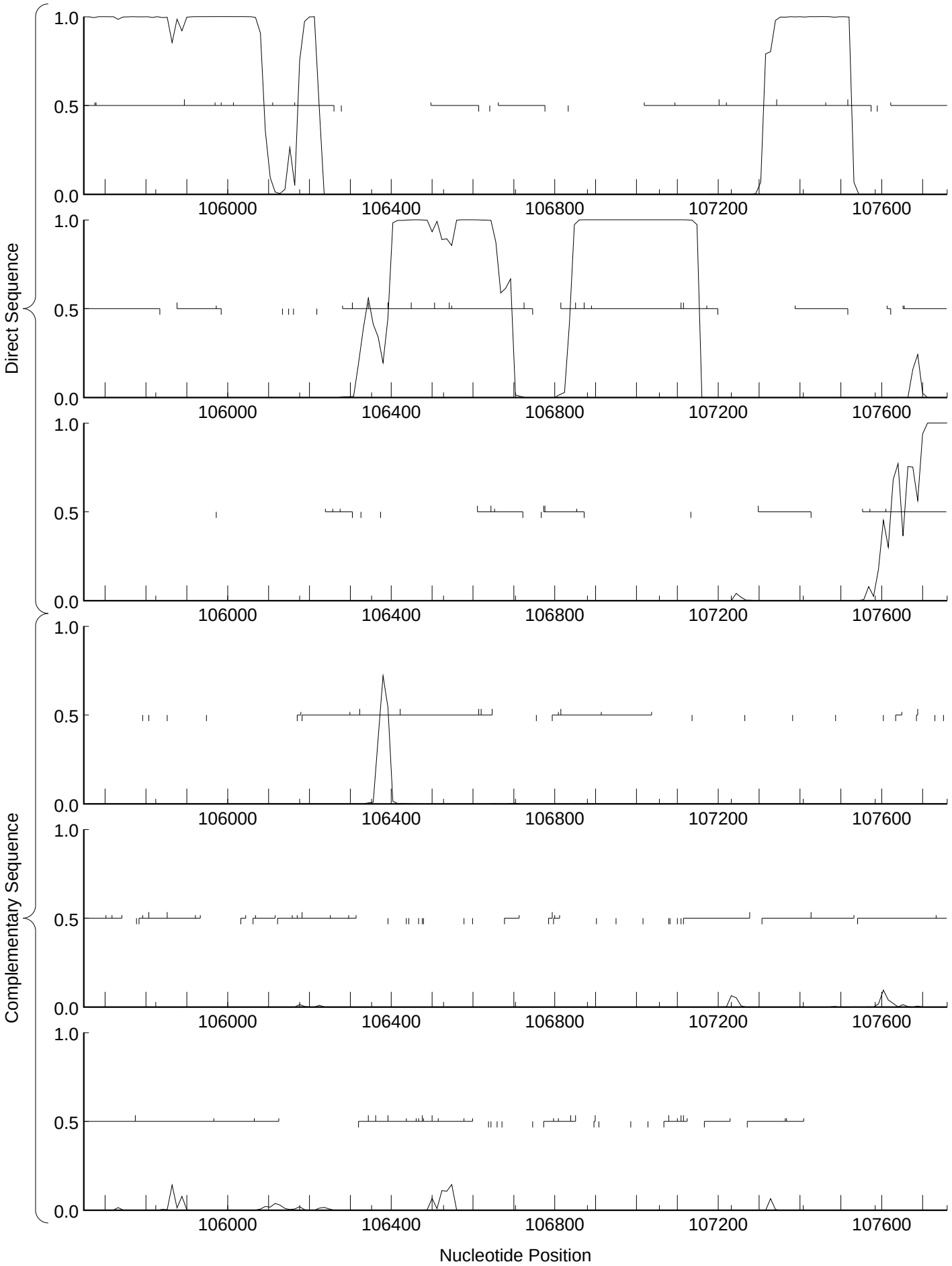
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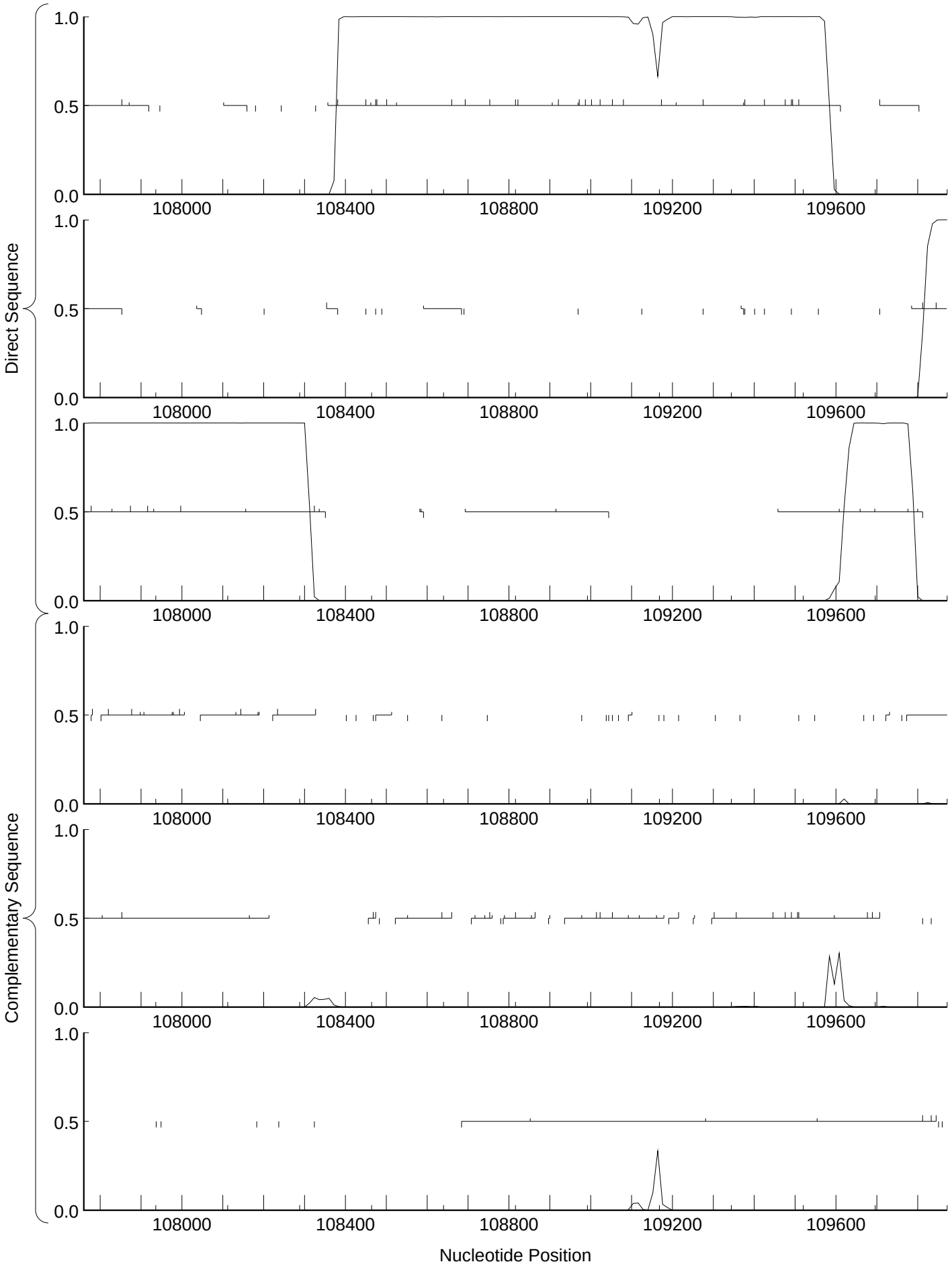
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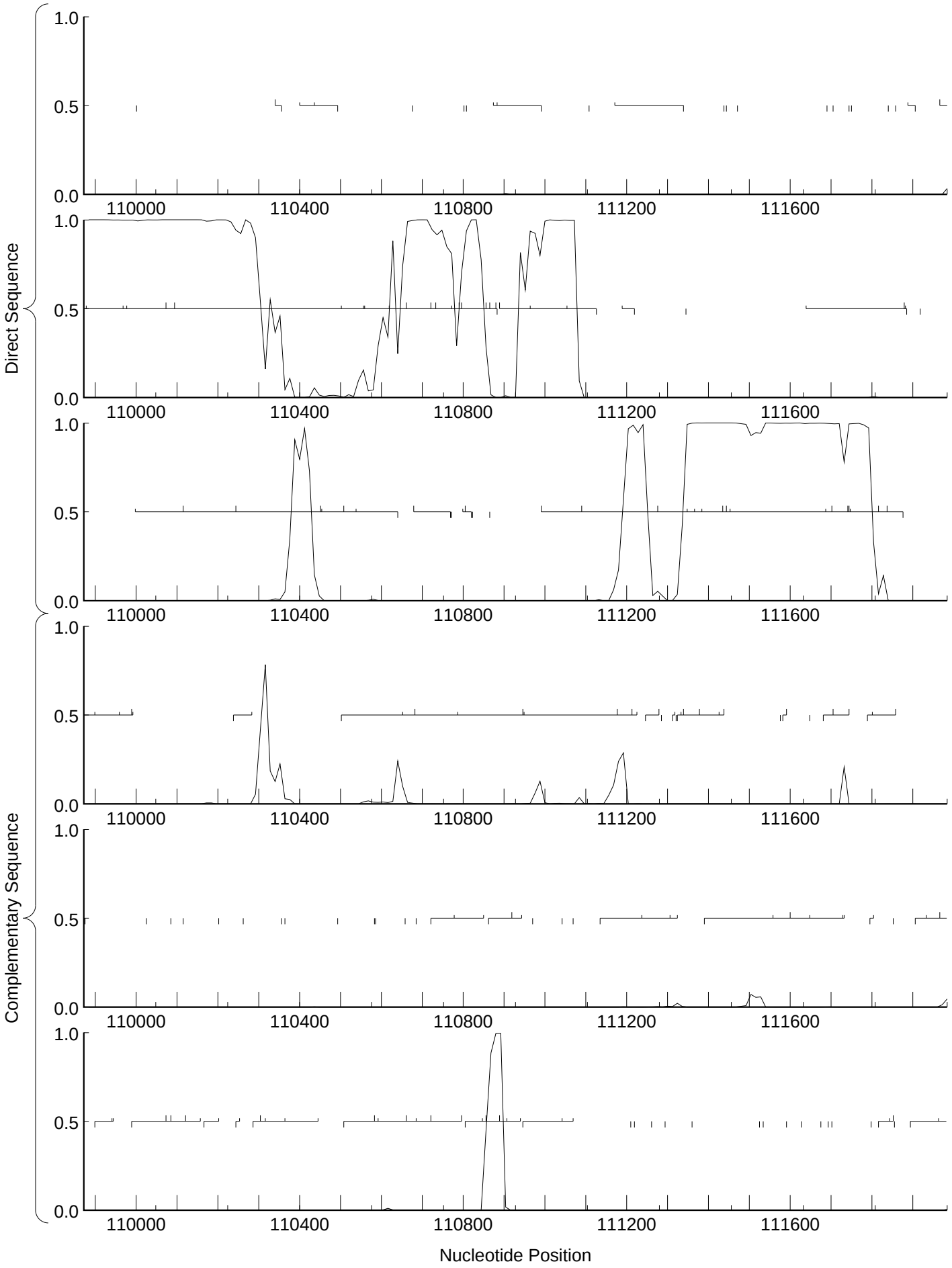
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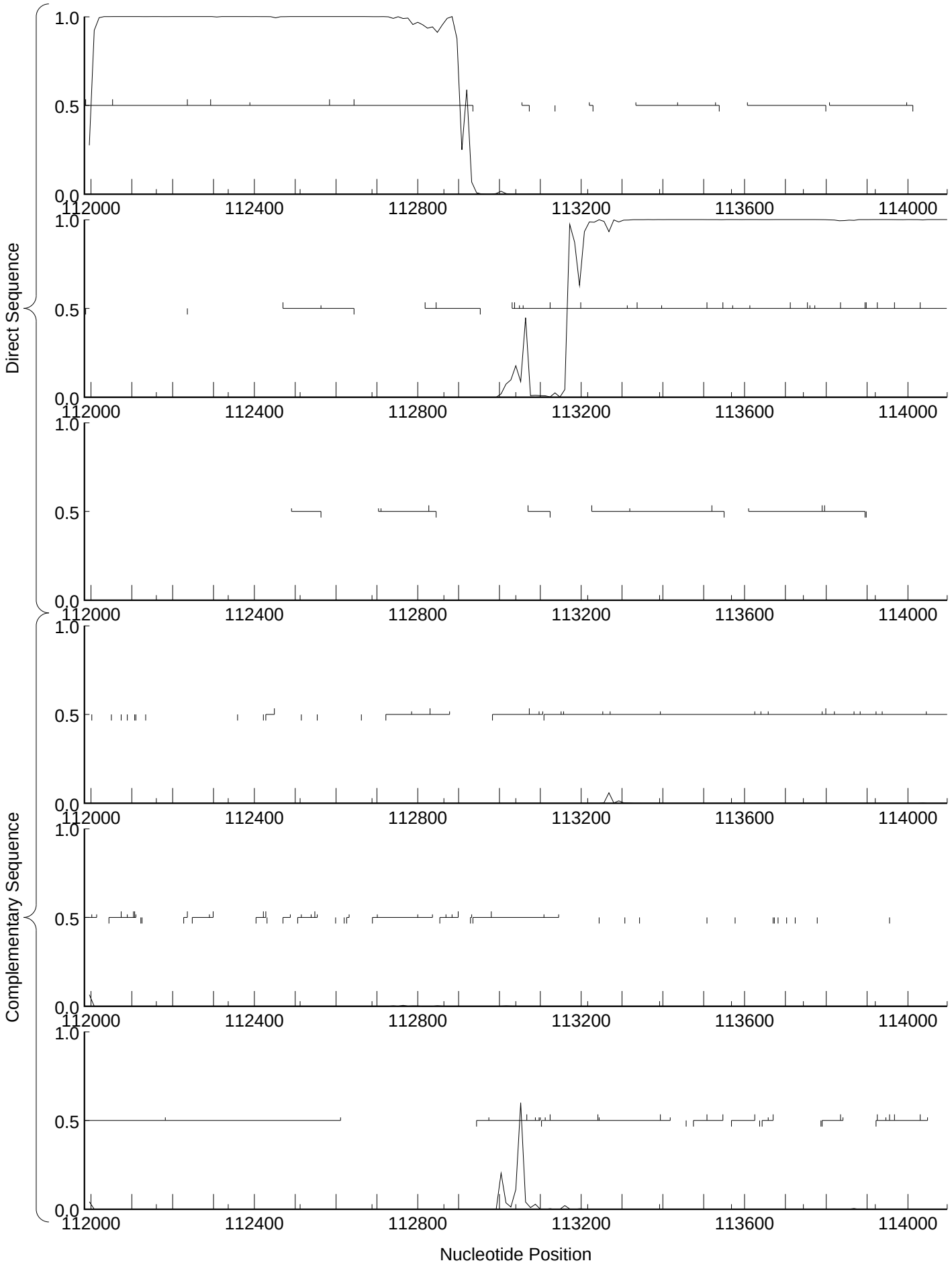
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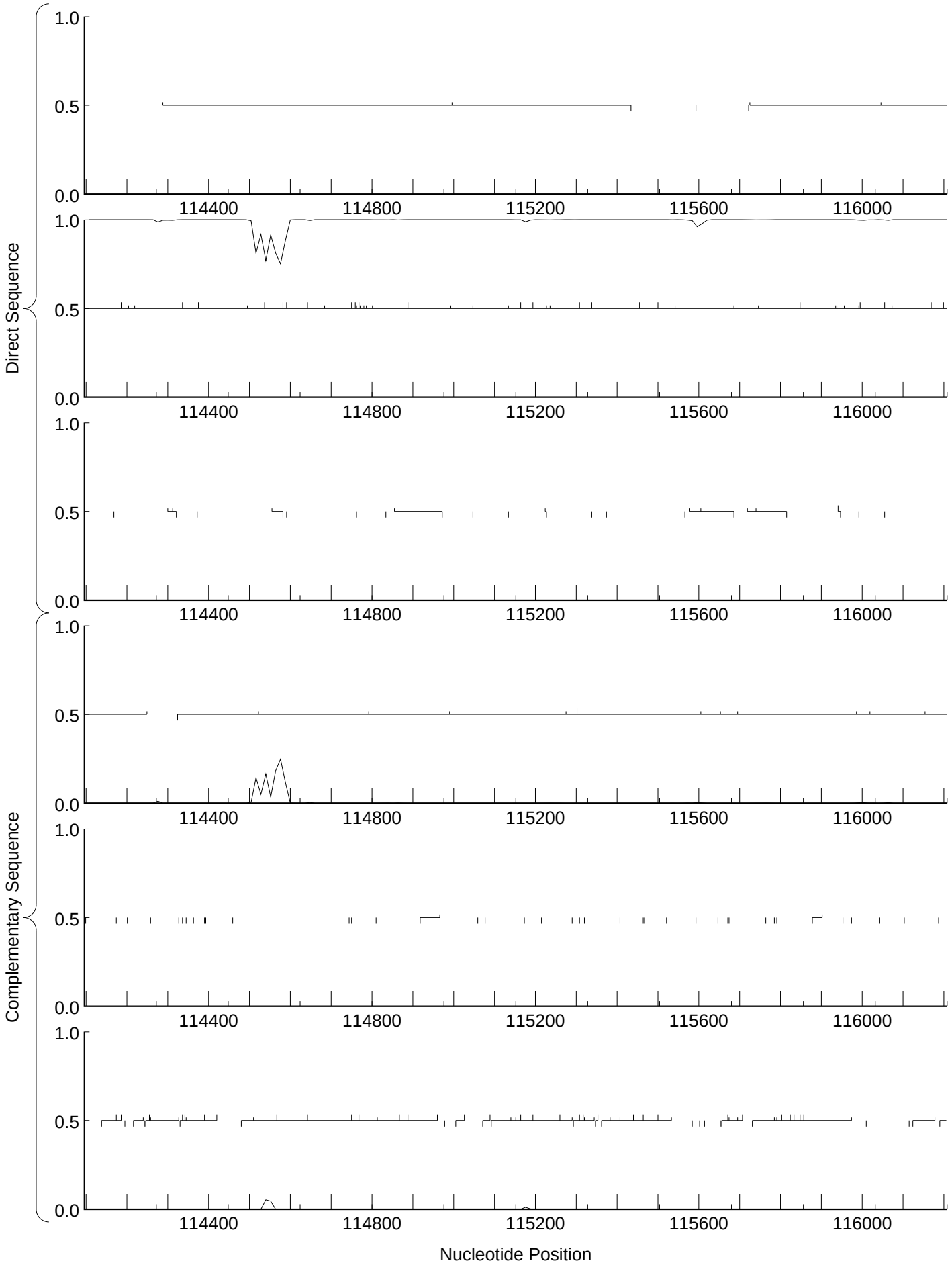
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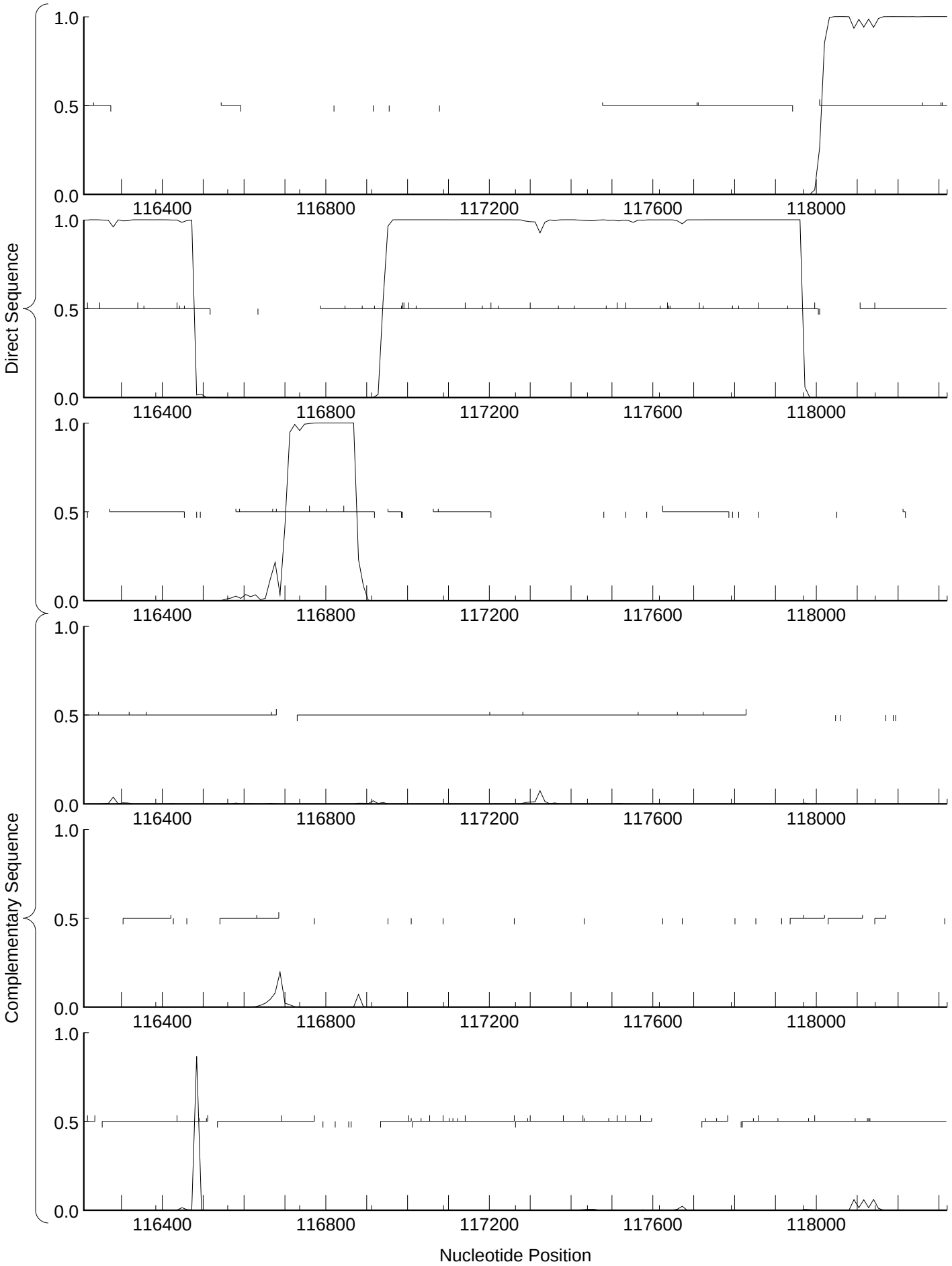
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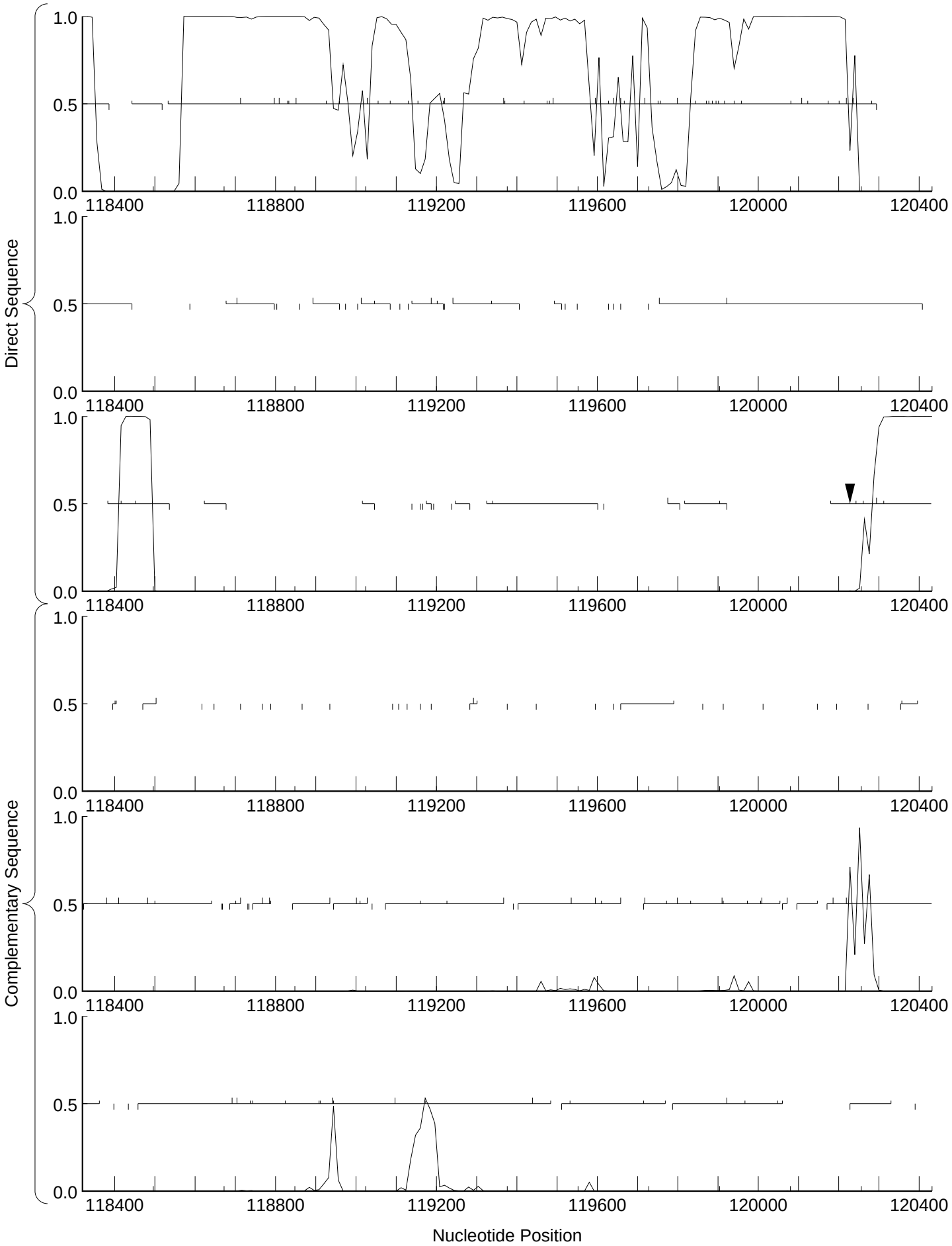
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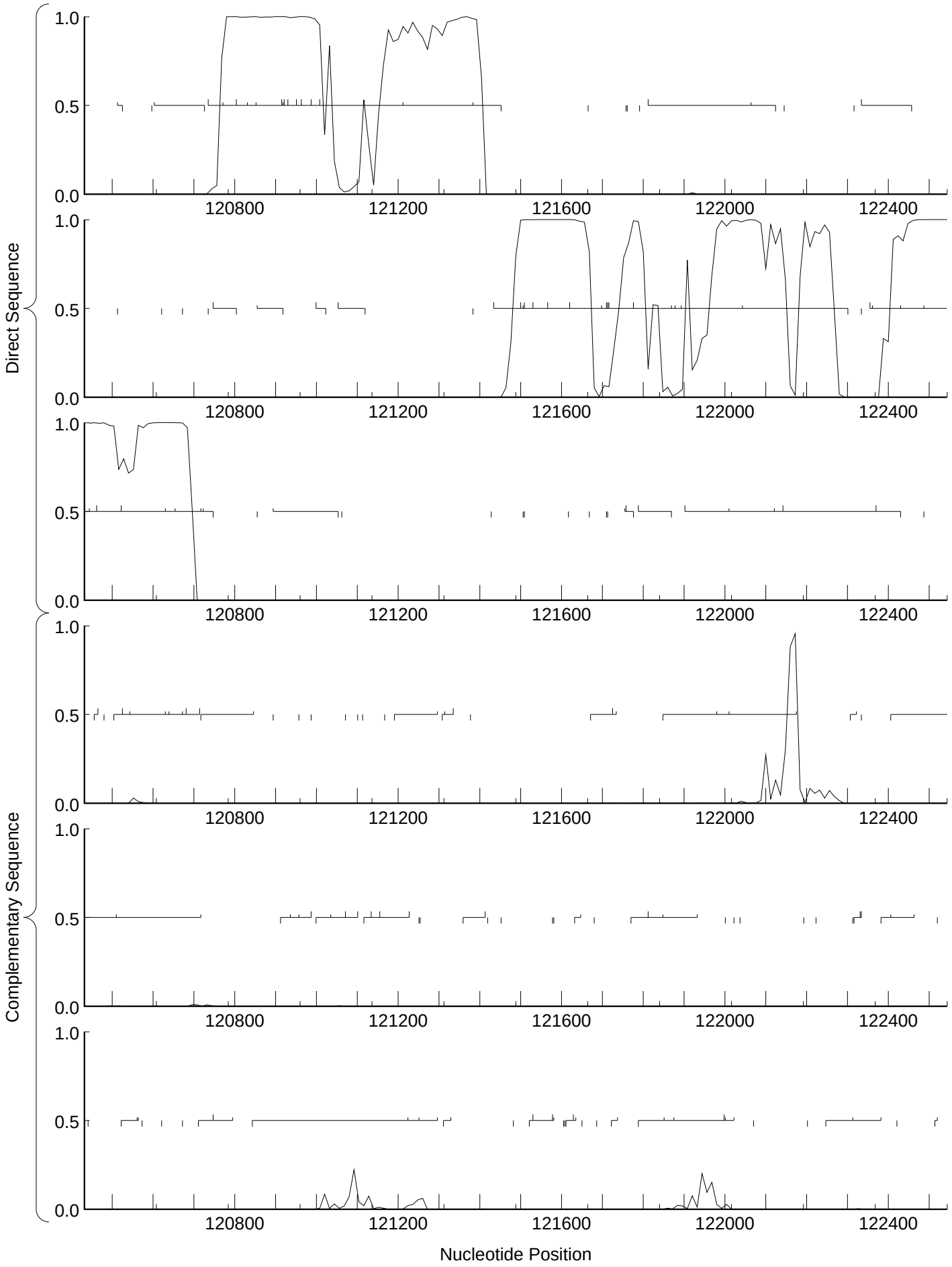
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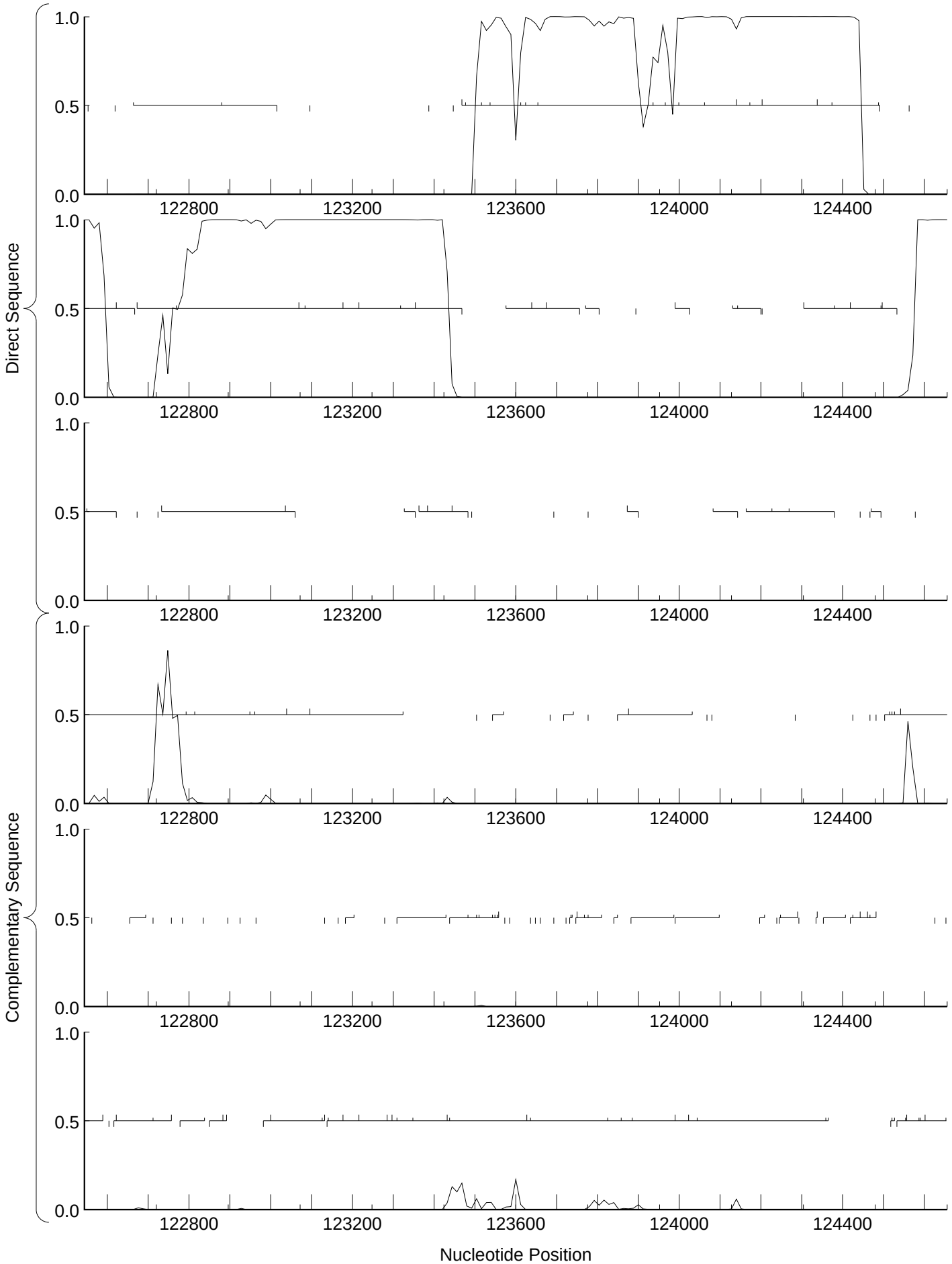
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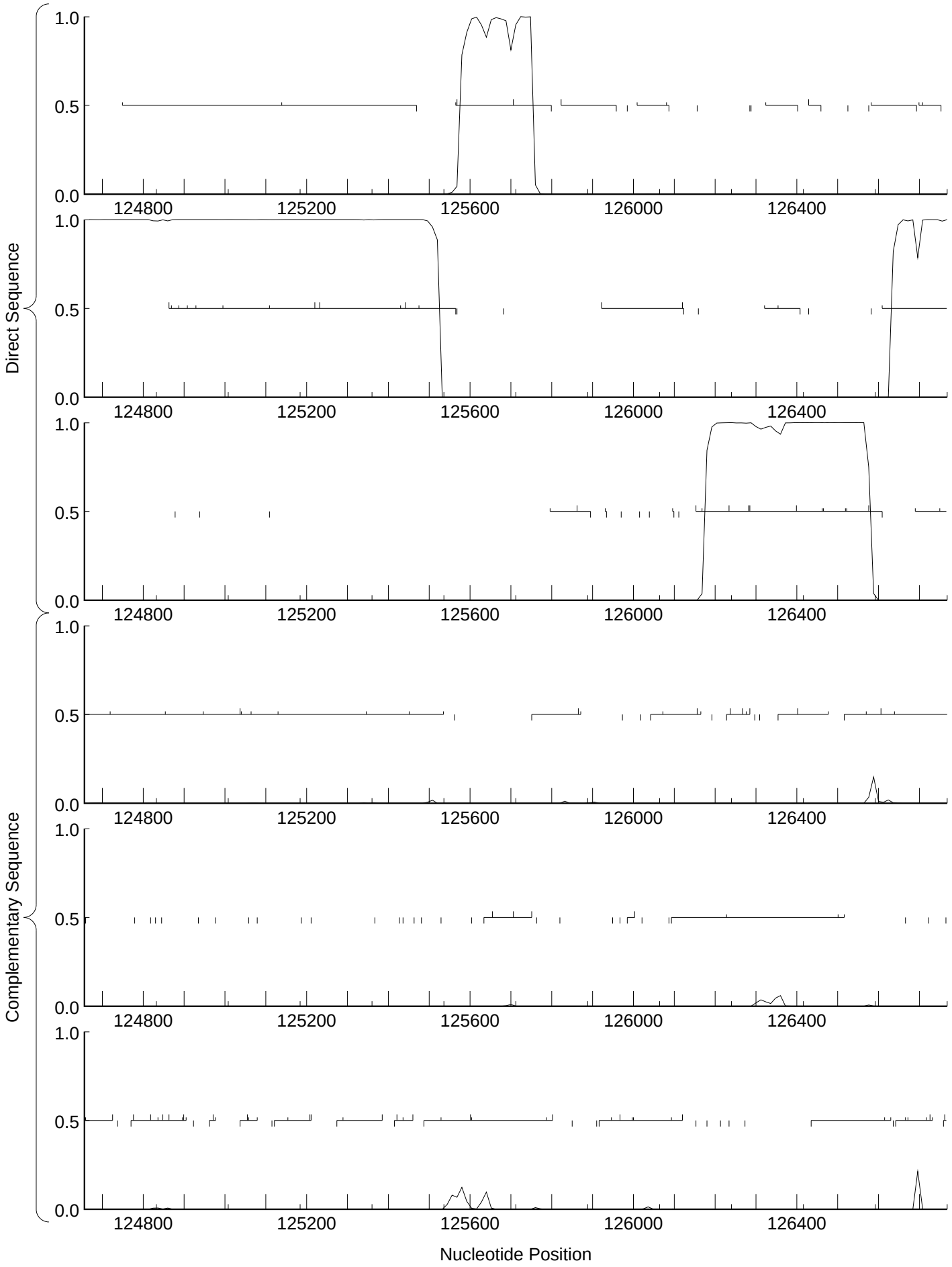
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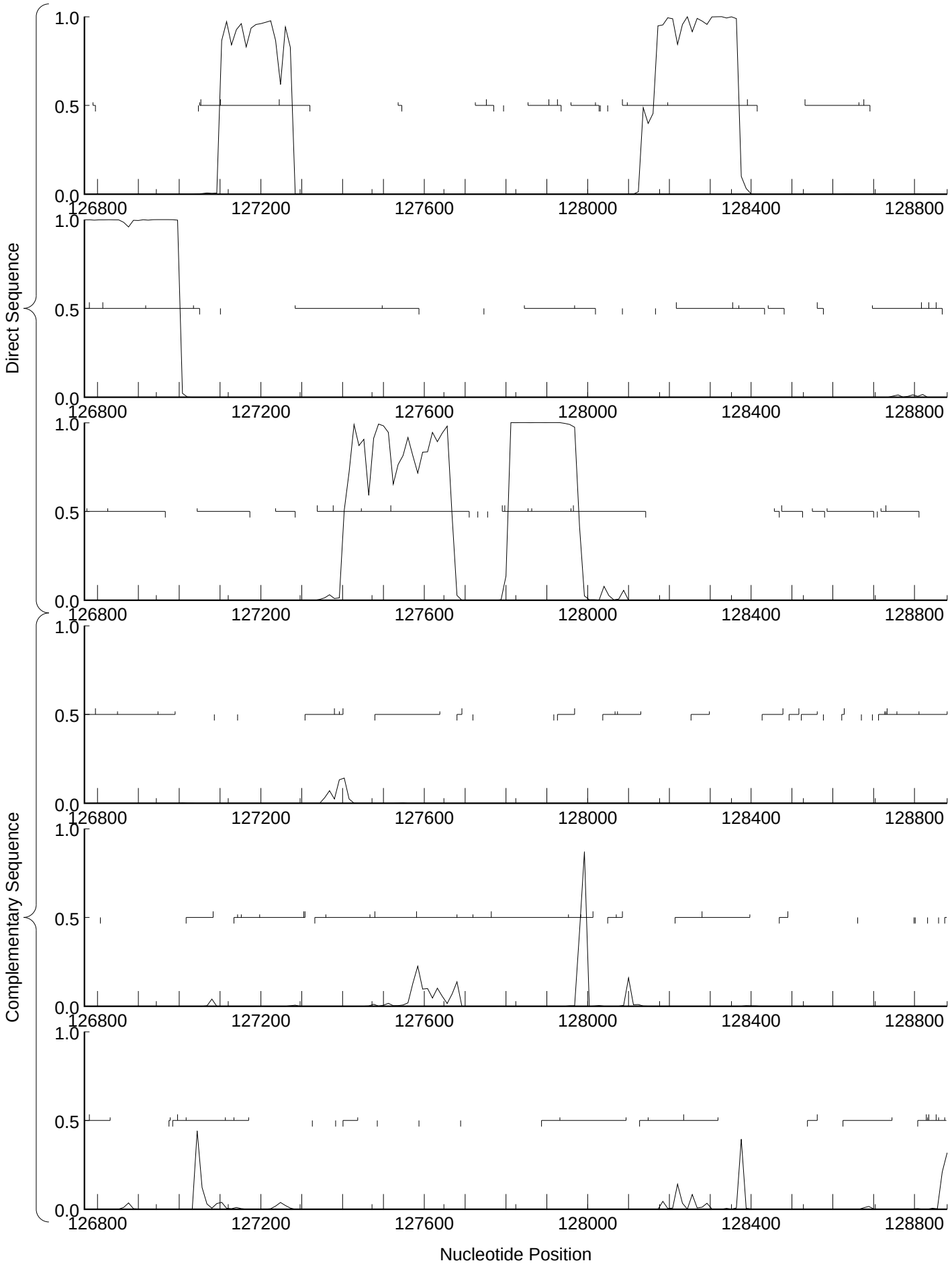
ycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 60/76



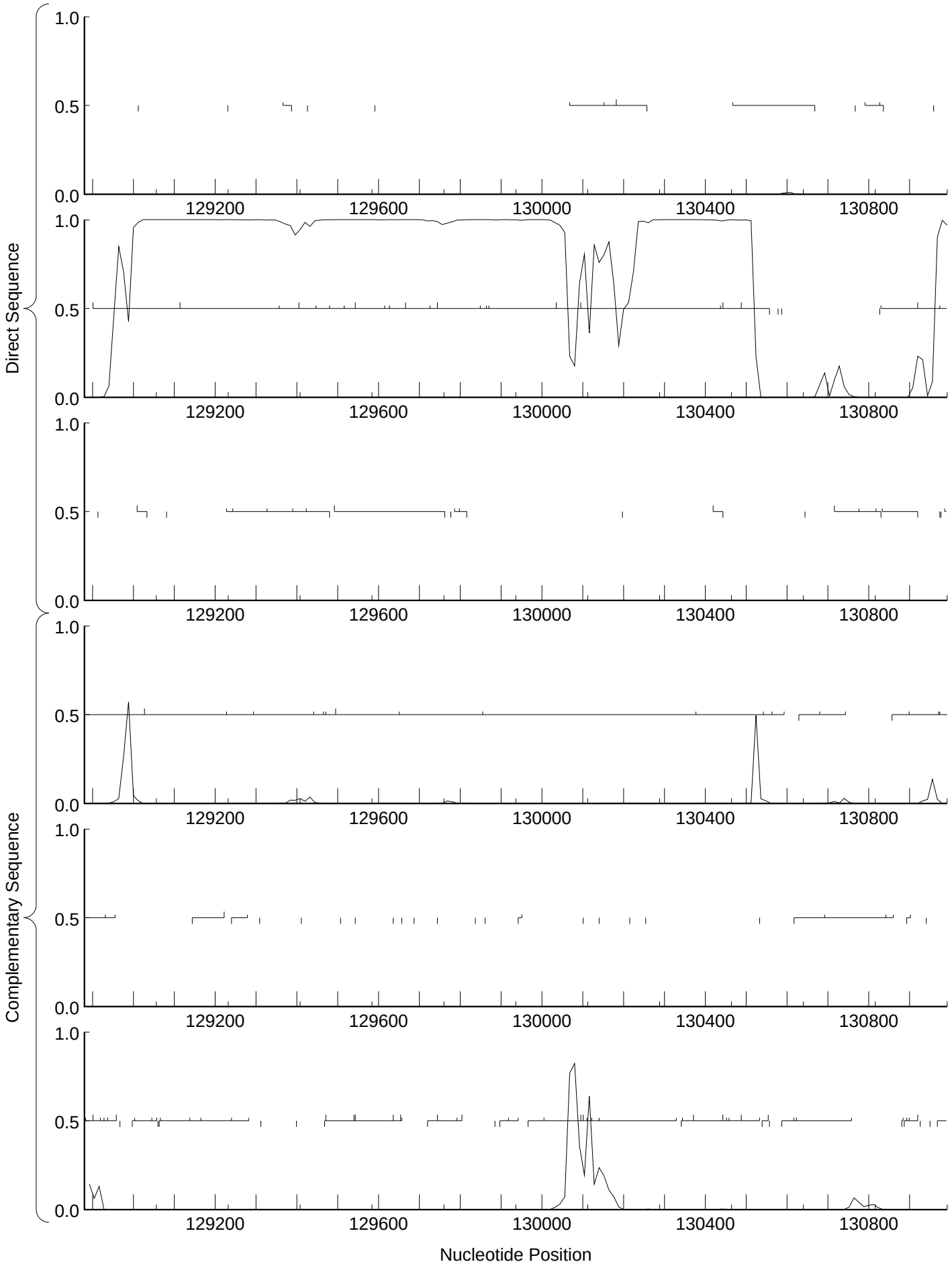
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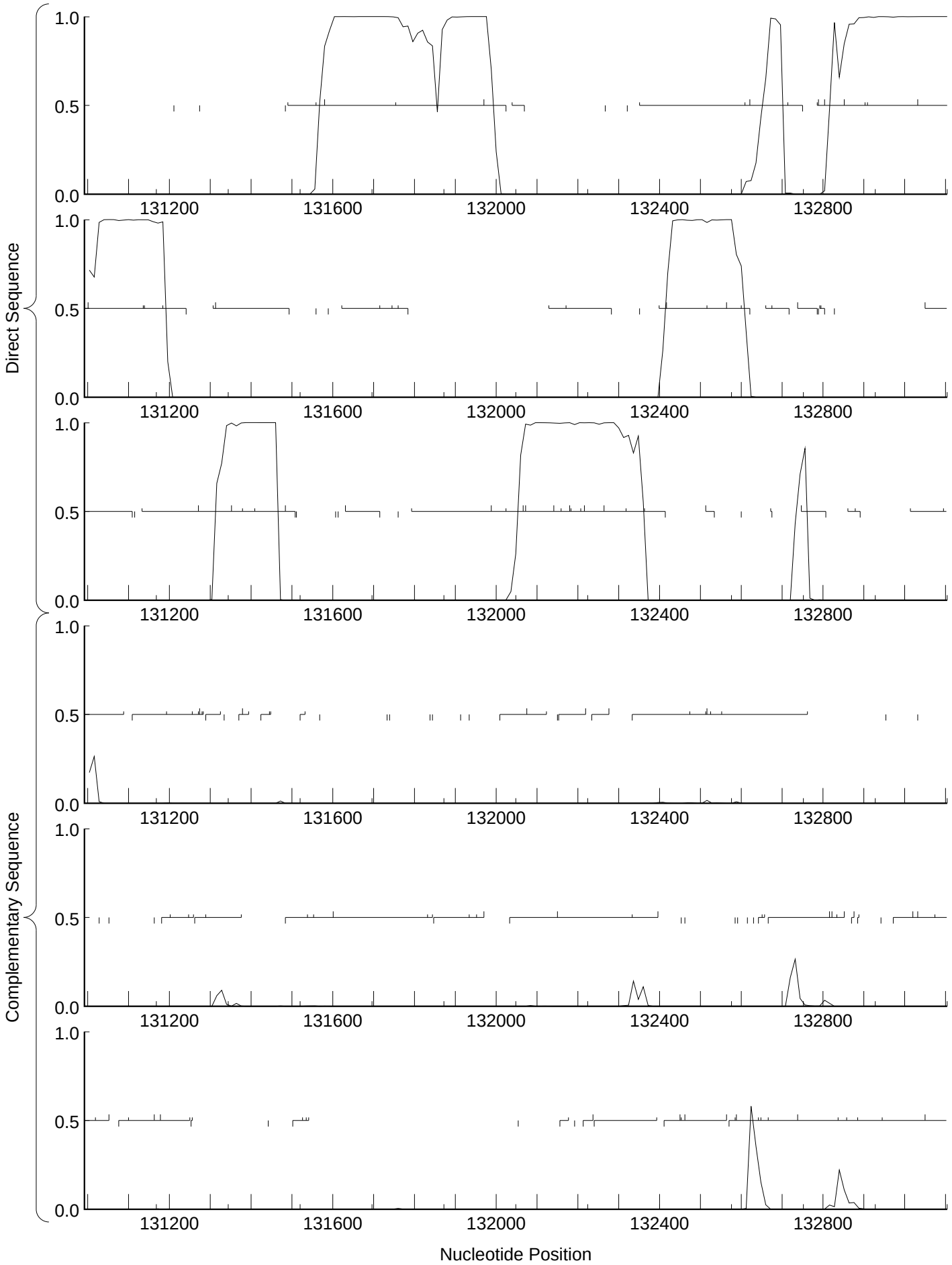
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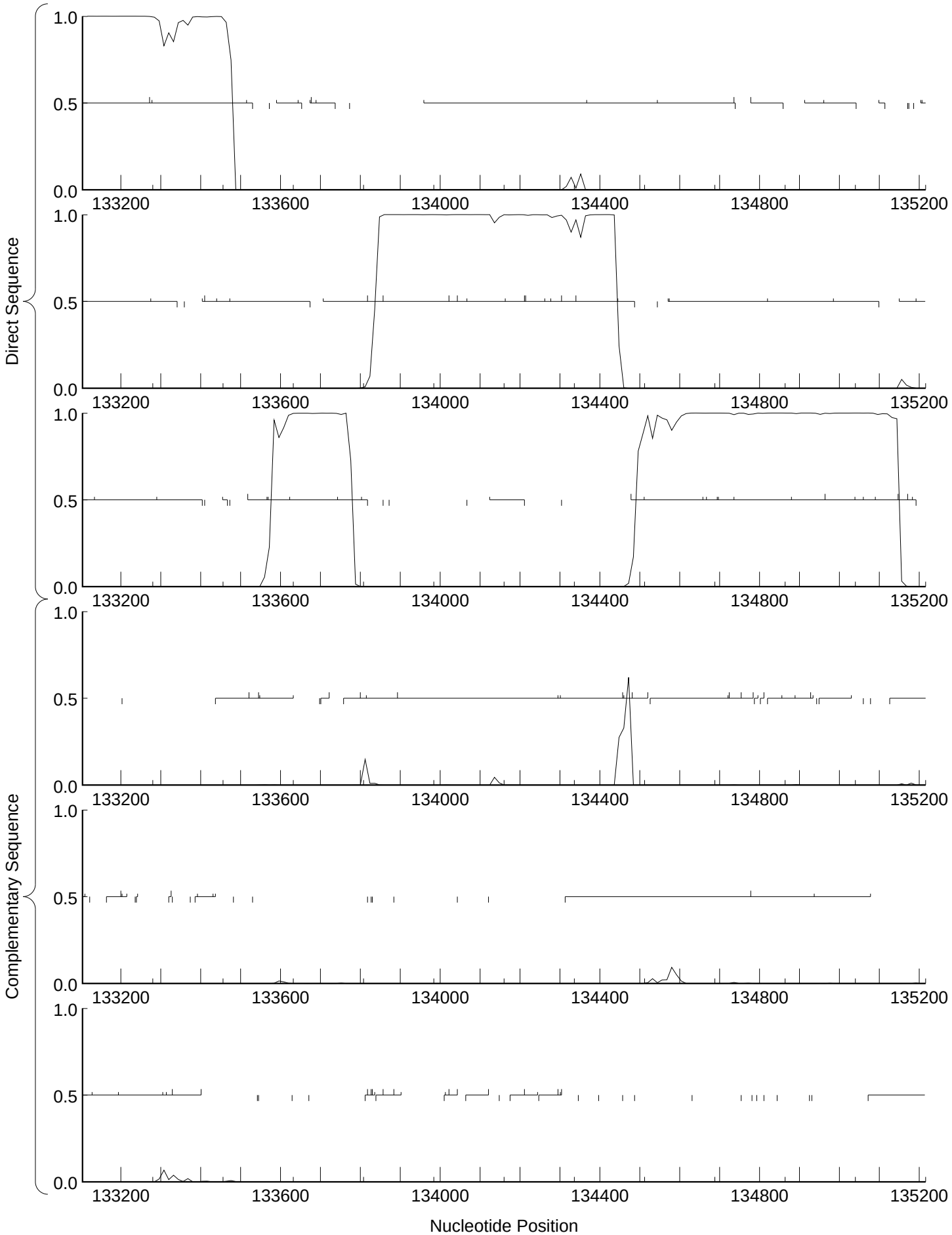
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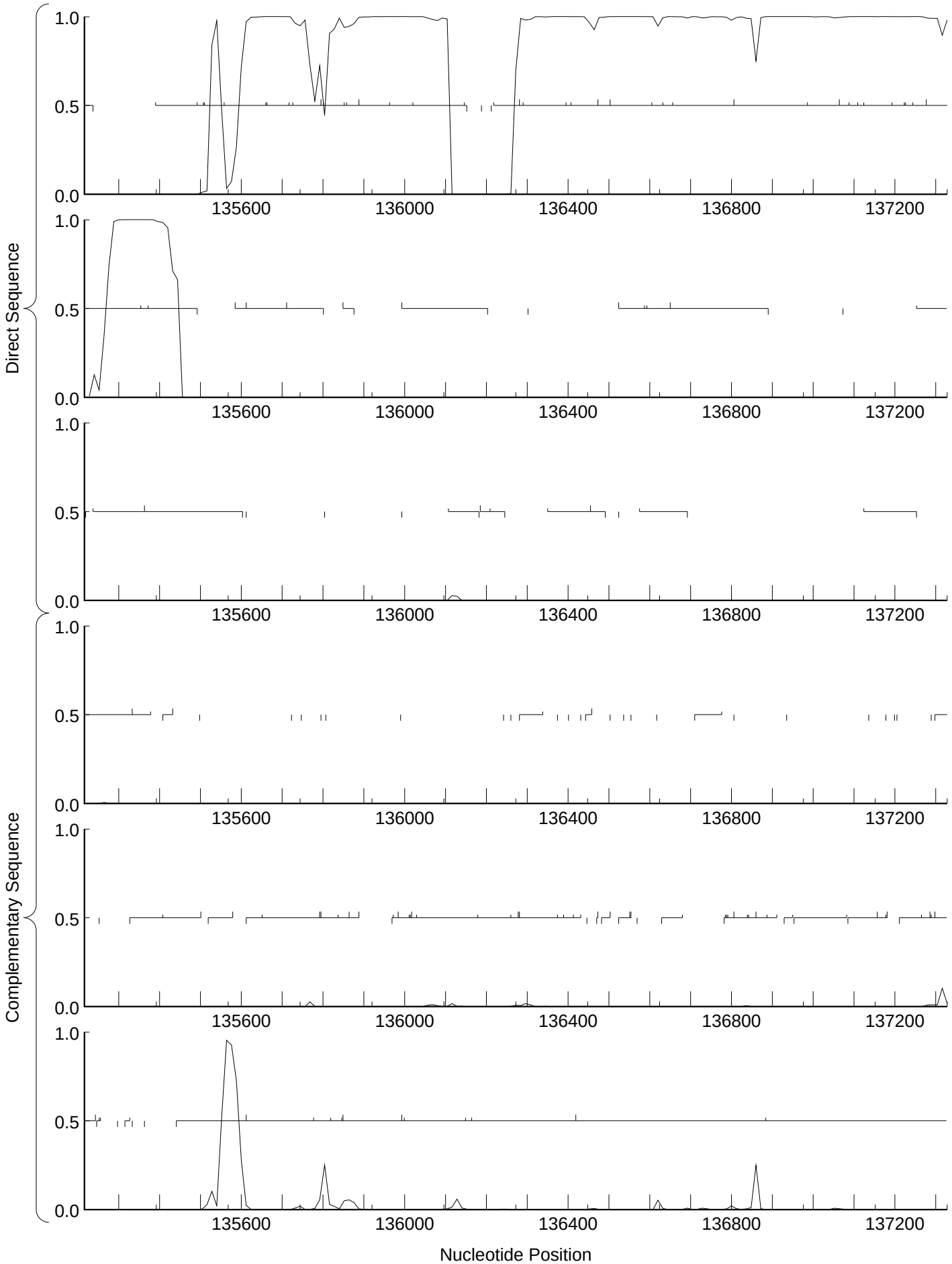
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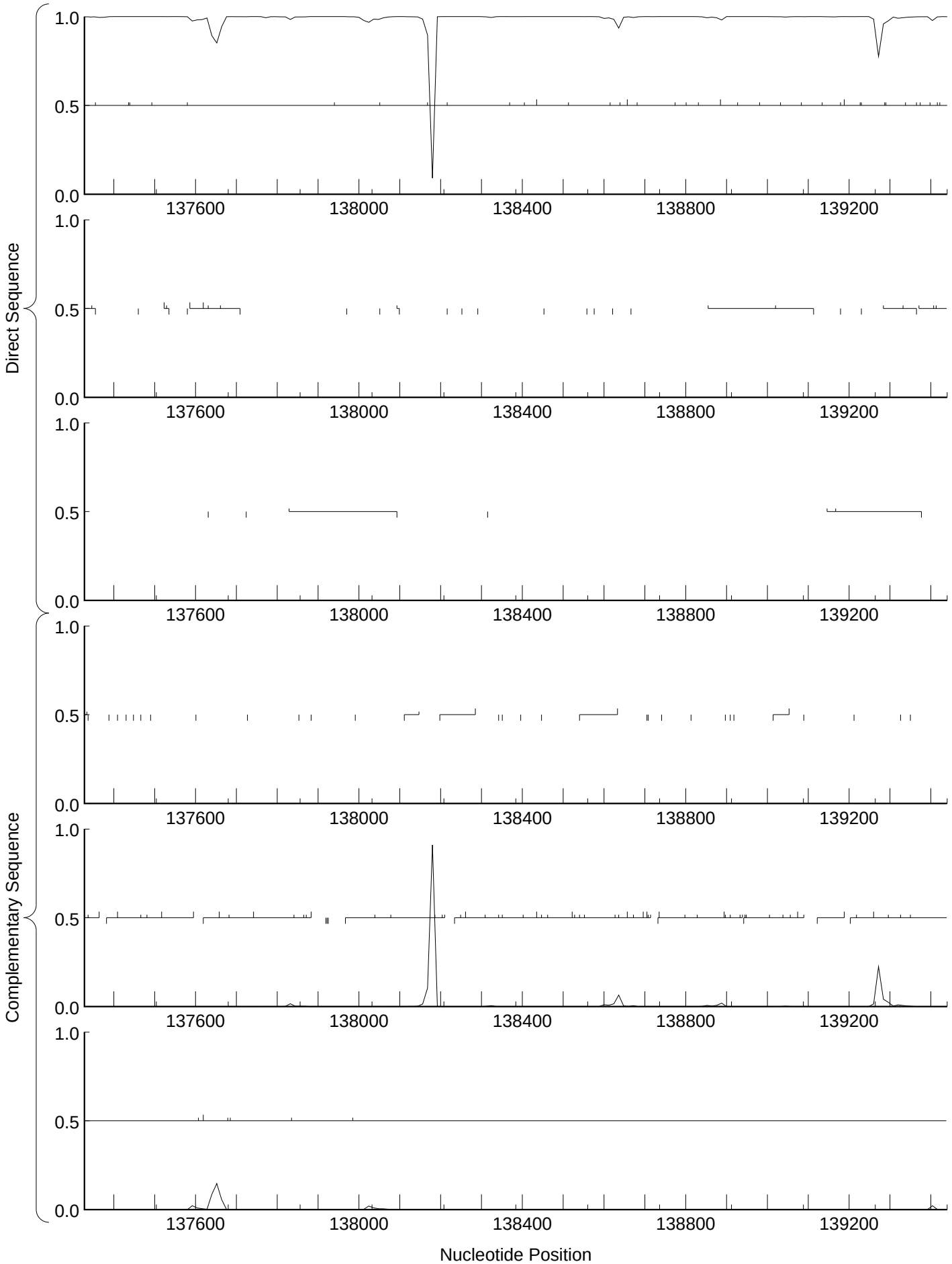
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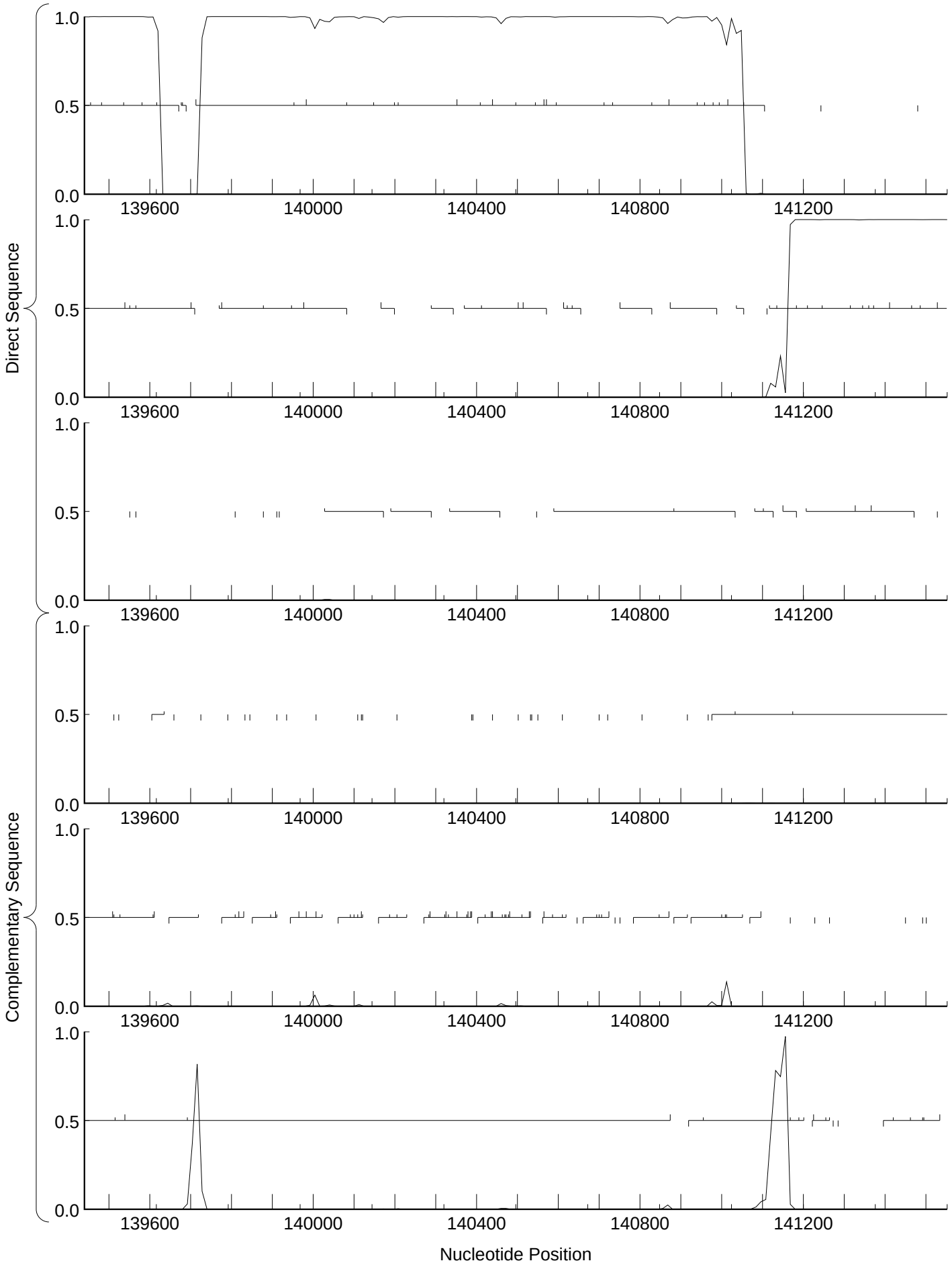
ycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 66/76



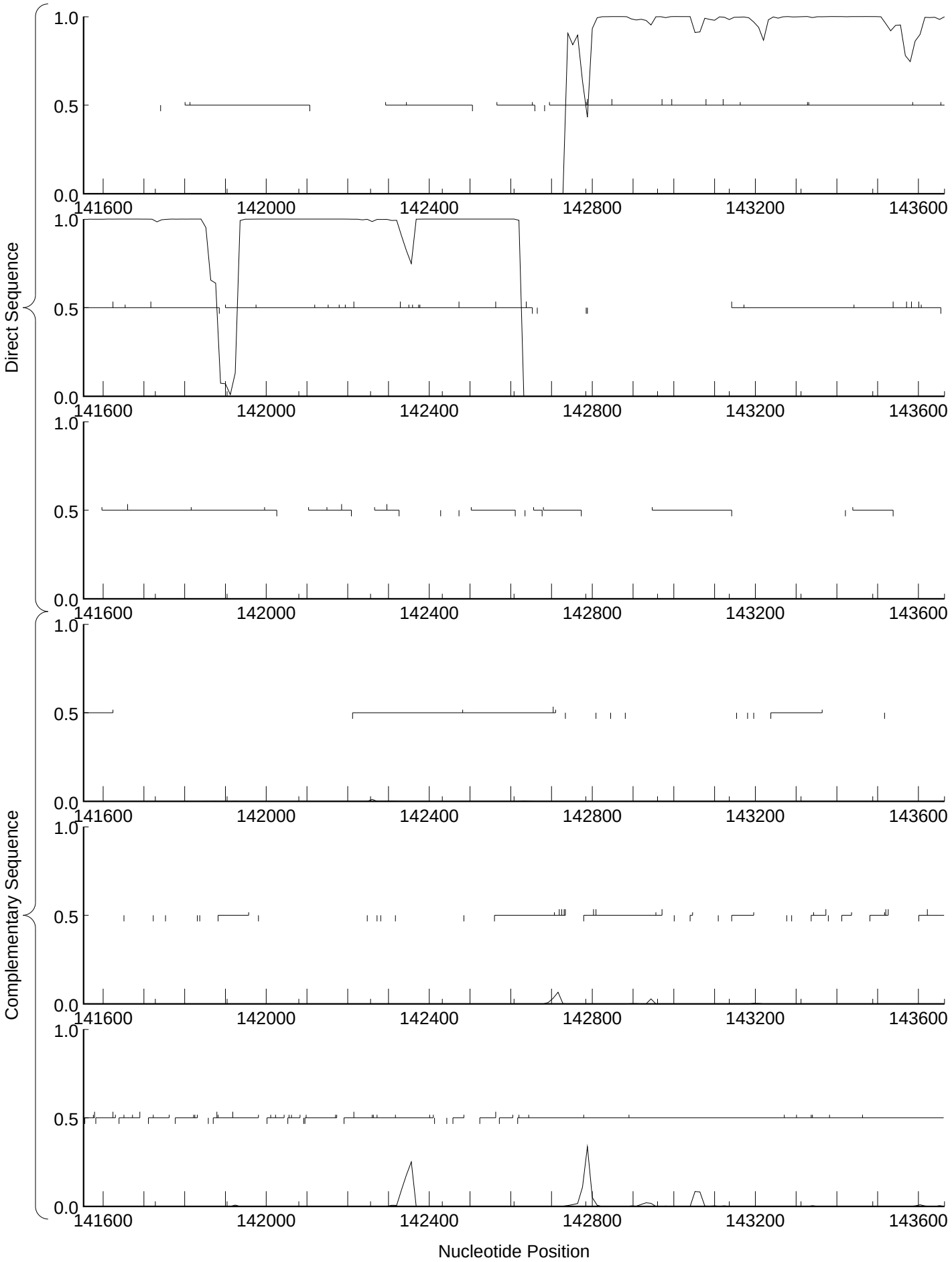
ycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 67/76



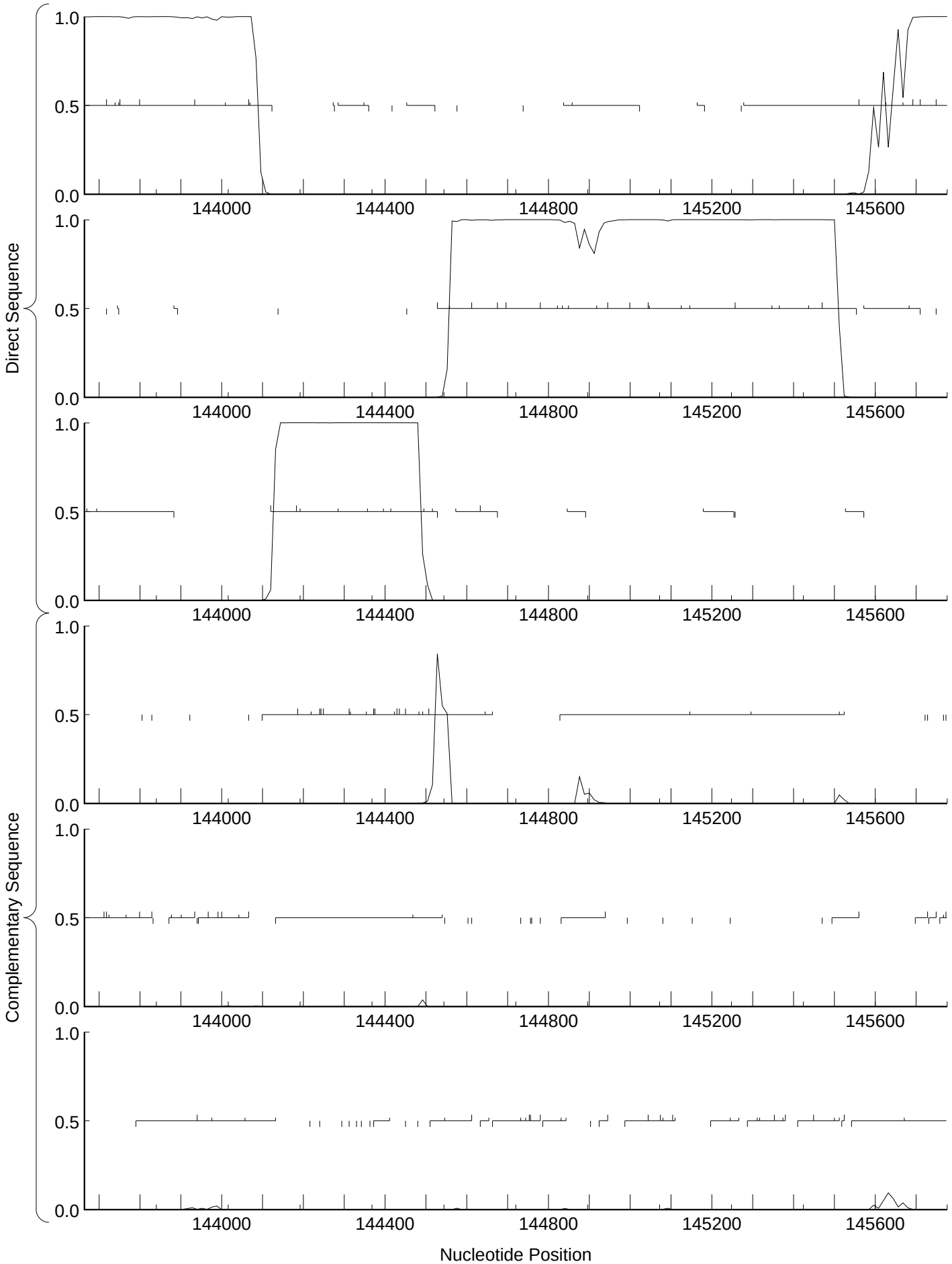
ycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 68/76



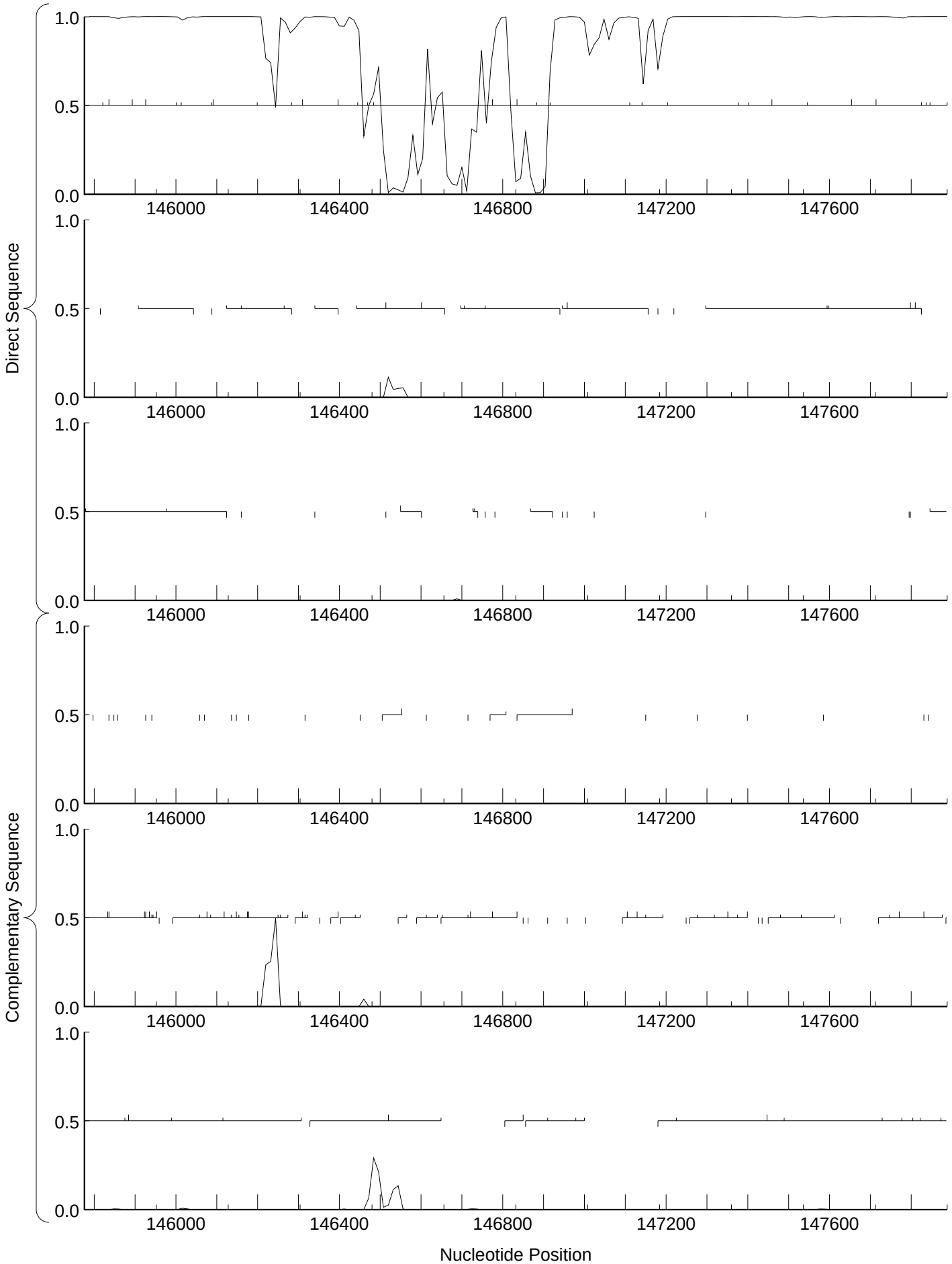
ycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 69/76



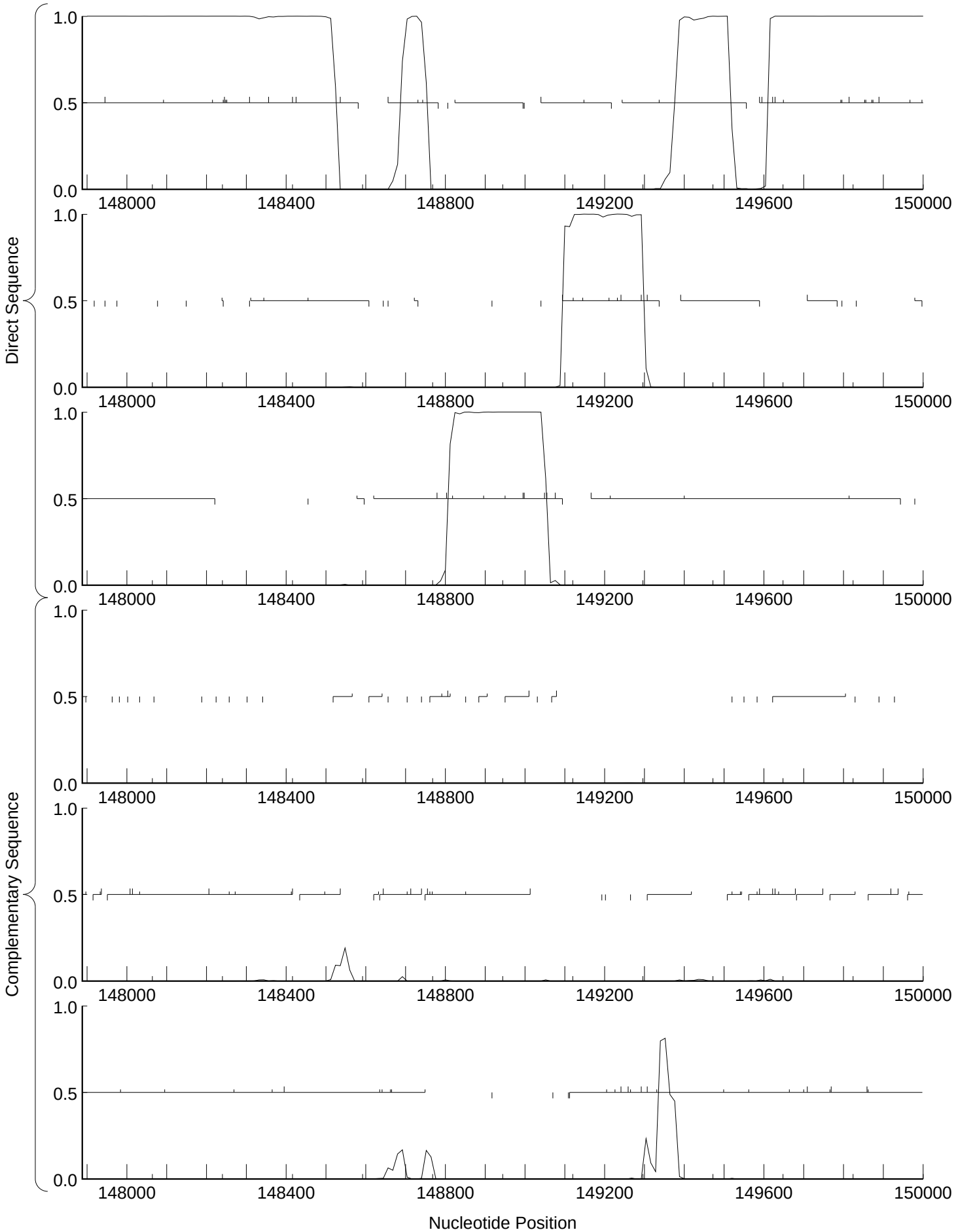
ycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 70/76



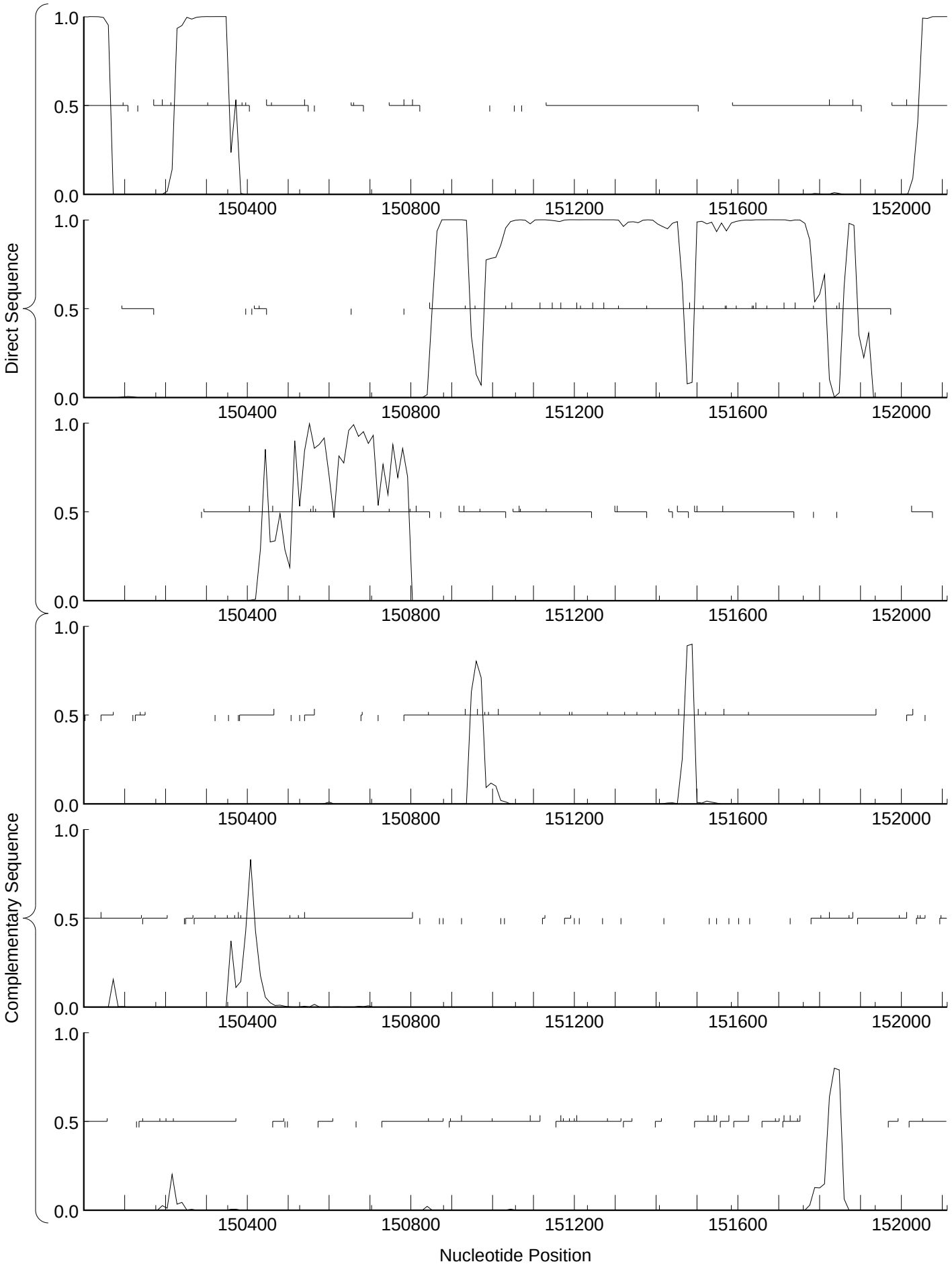
ycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 71/76



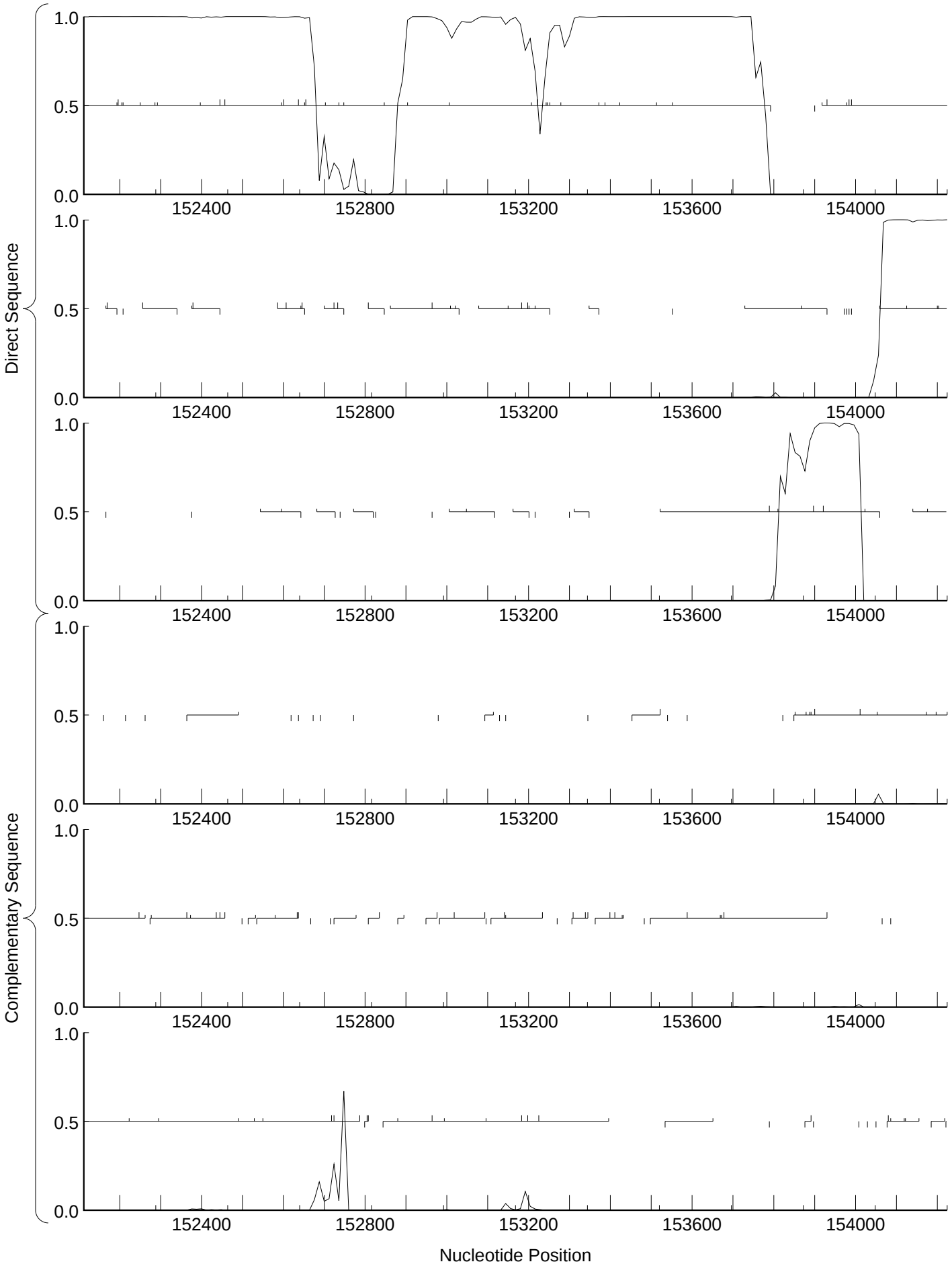
ycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 72/76



ycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 73/76



ycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 74/76



ycobacterium phage RoMag complete sequence, 156319 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 75/76

