GeneMark

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

PROGRAM INFORMATION

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Sequence         :  Mycobacterium phage Constella complete sequence, 110169 bp including 4-base 3’ overhang (AT
Analysis Date    :  3/29/24 at 11:21:07
Pages            :  54
Sequence Length  :  110169 bp
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Window Length    :  96 bp
Window Step      :  12 bp
Threshold Value  :  0.500

PS-Version       :  1.2

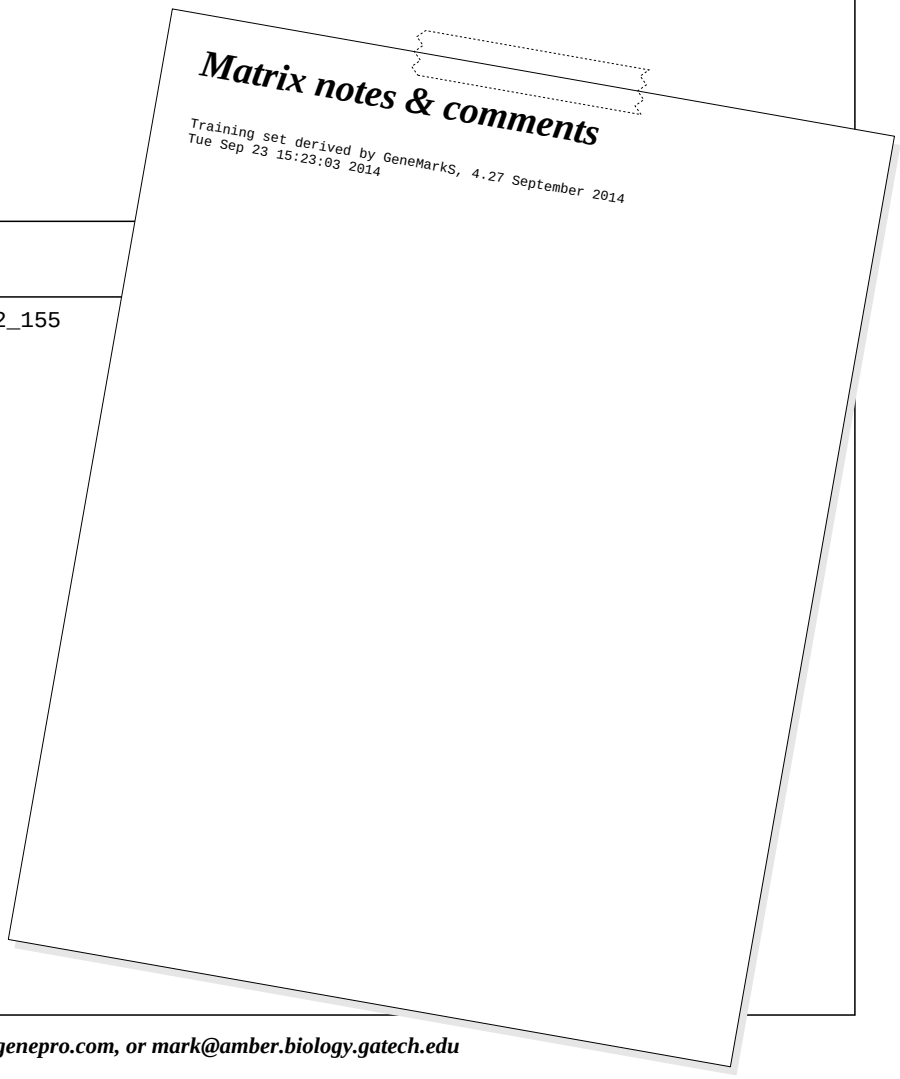
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MATRIX INFORMATION

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Author     :  -

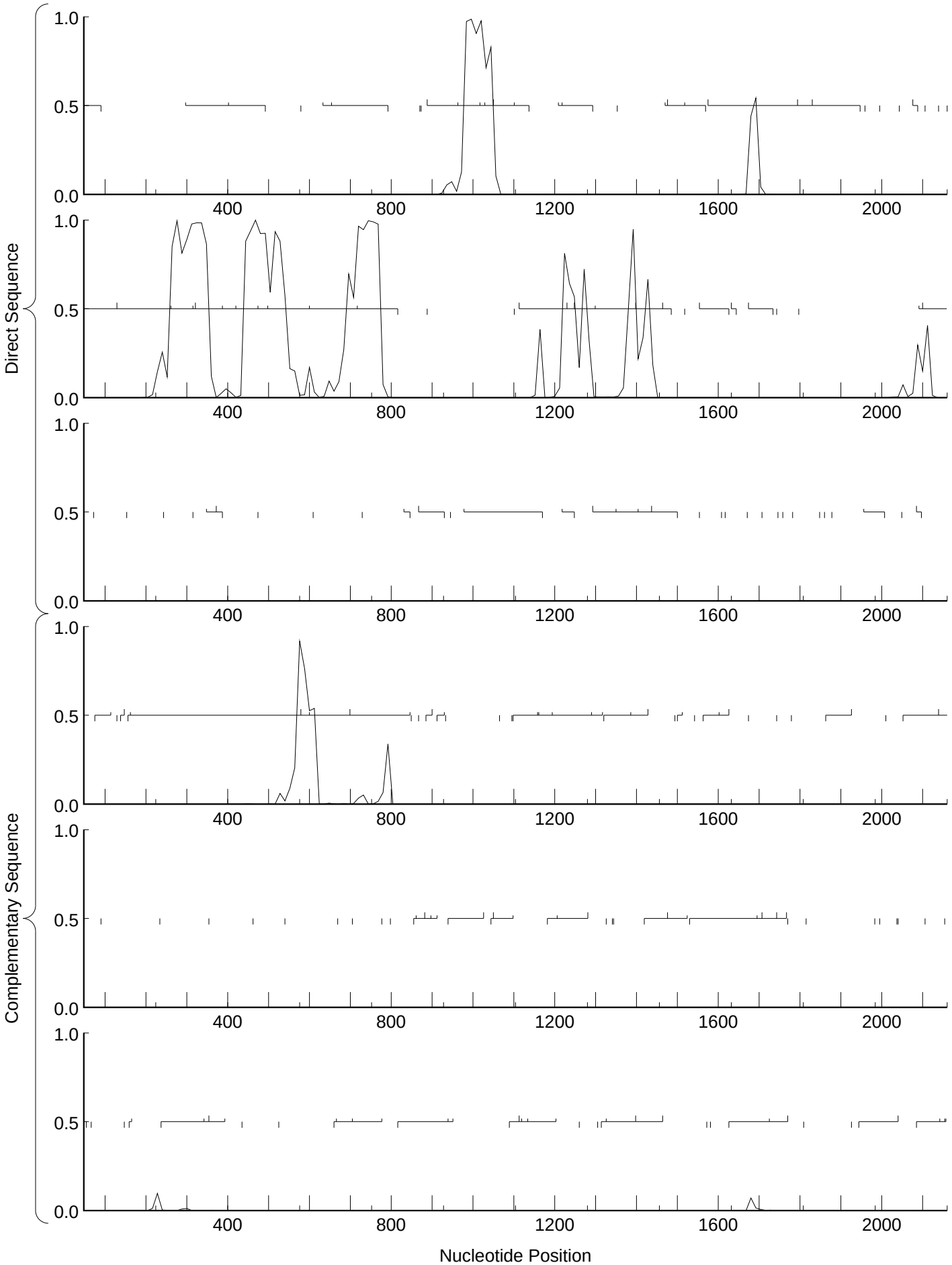
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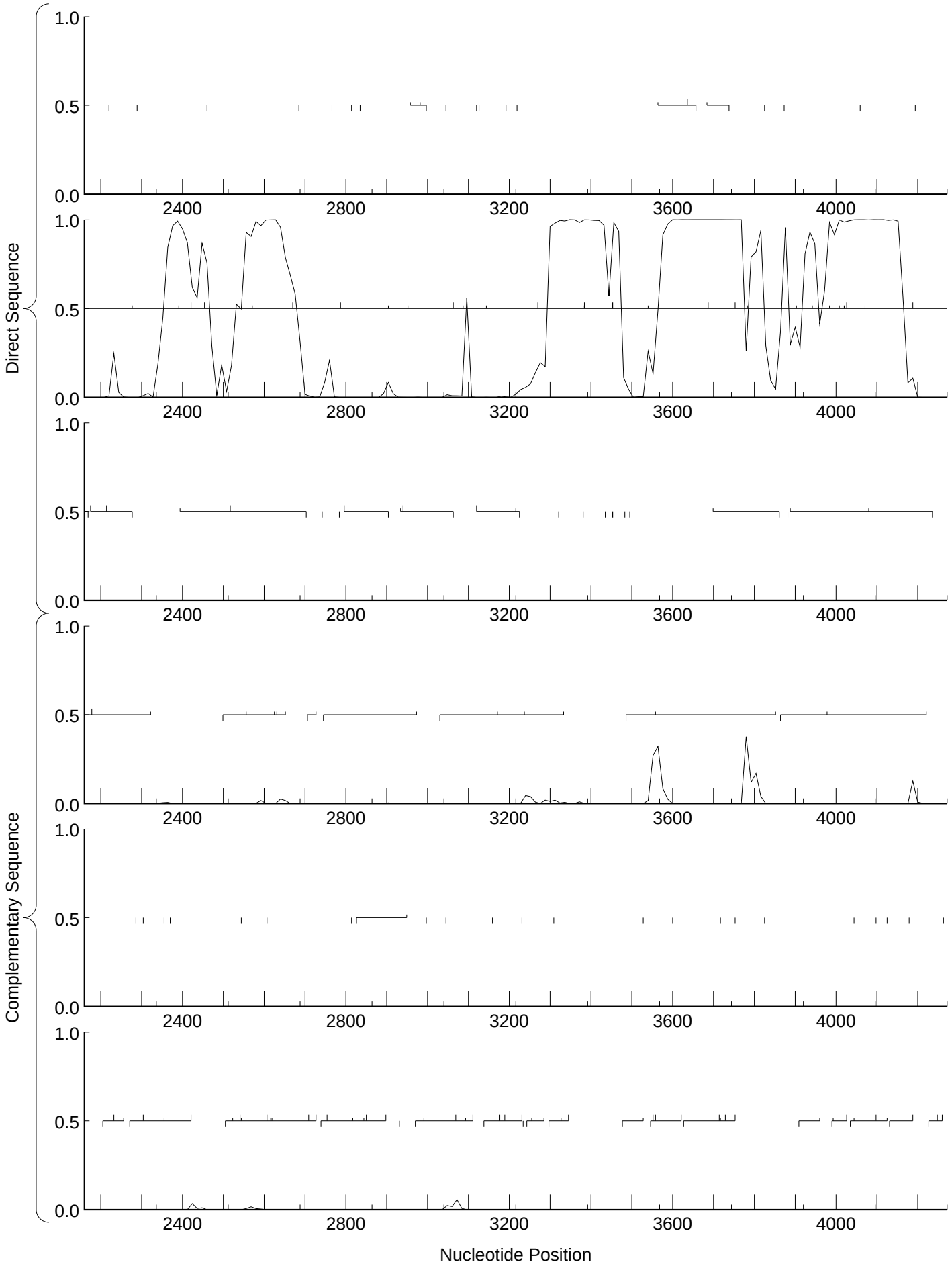

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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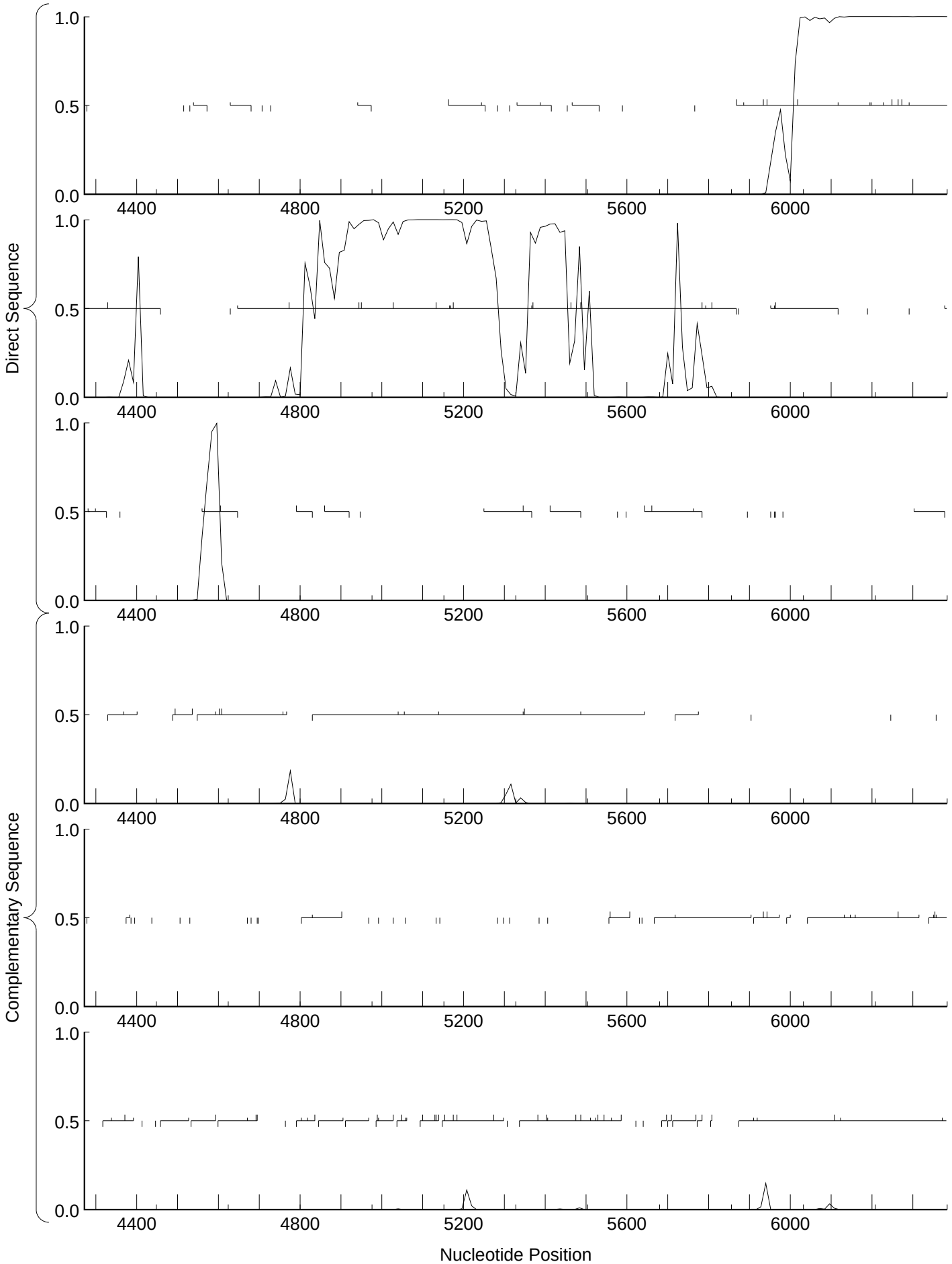


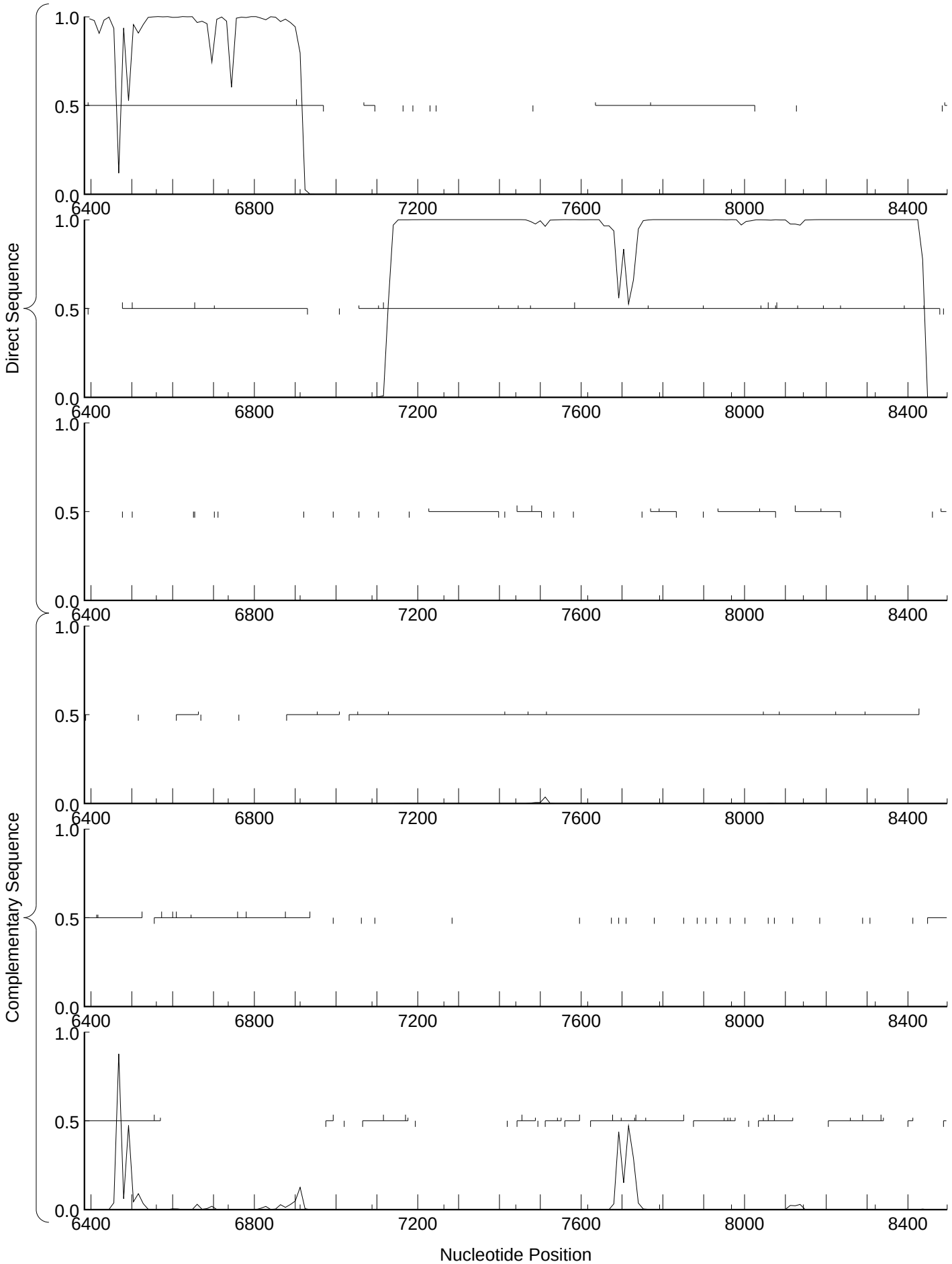
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

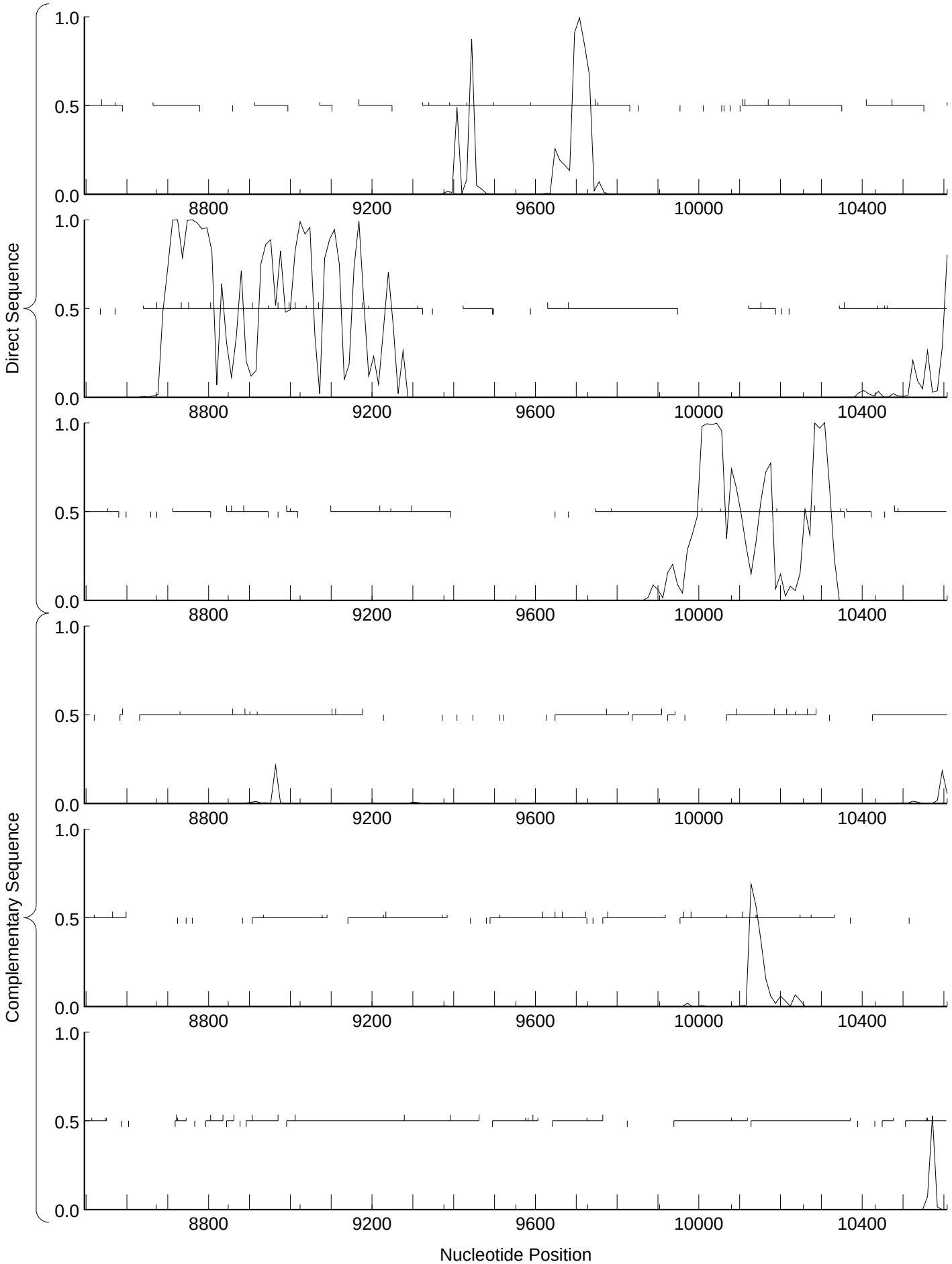
ge Constella complete sequence, 110169 bp including 4-base 3’ overhang (ATCC), Cluster J, Order 4, Window 96, Step 12, 2/54

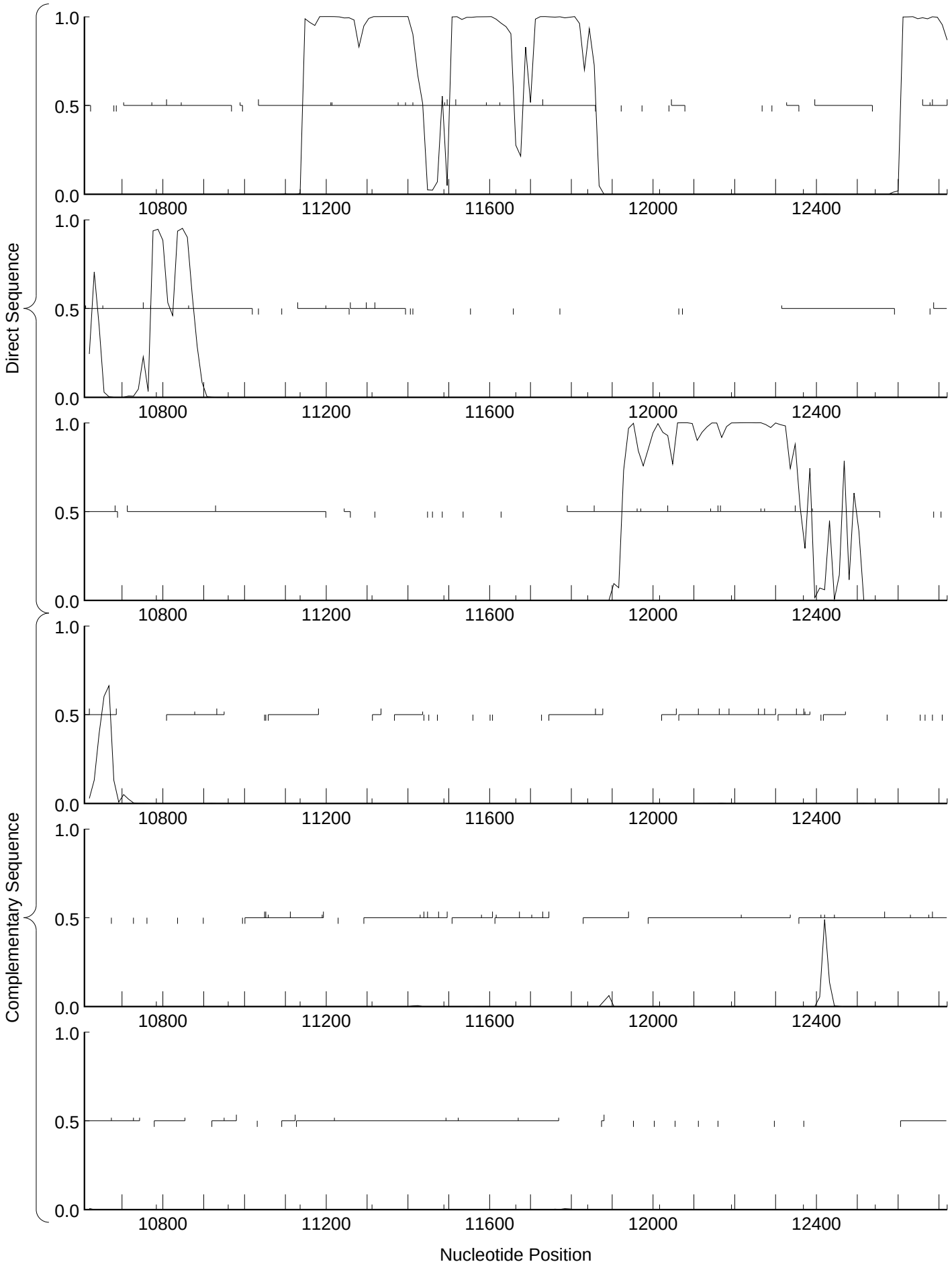


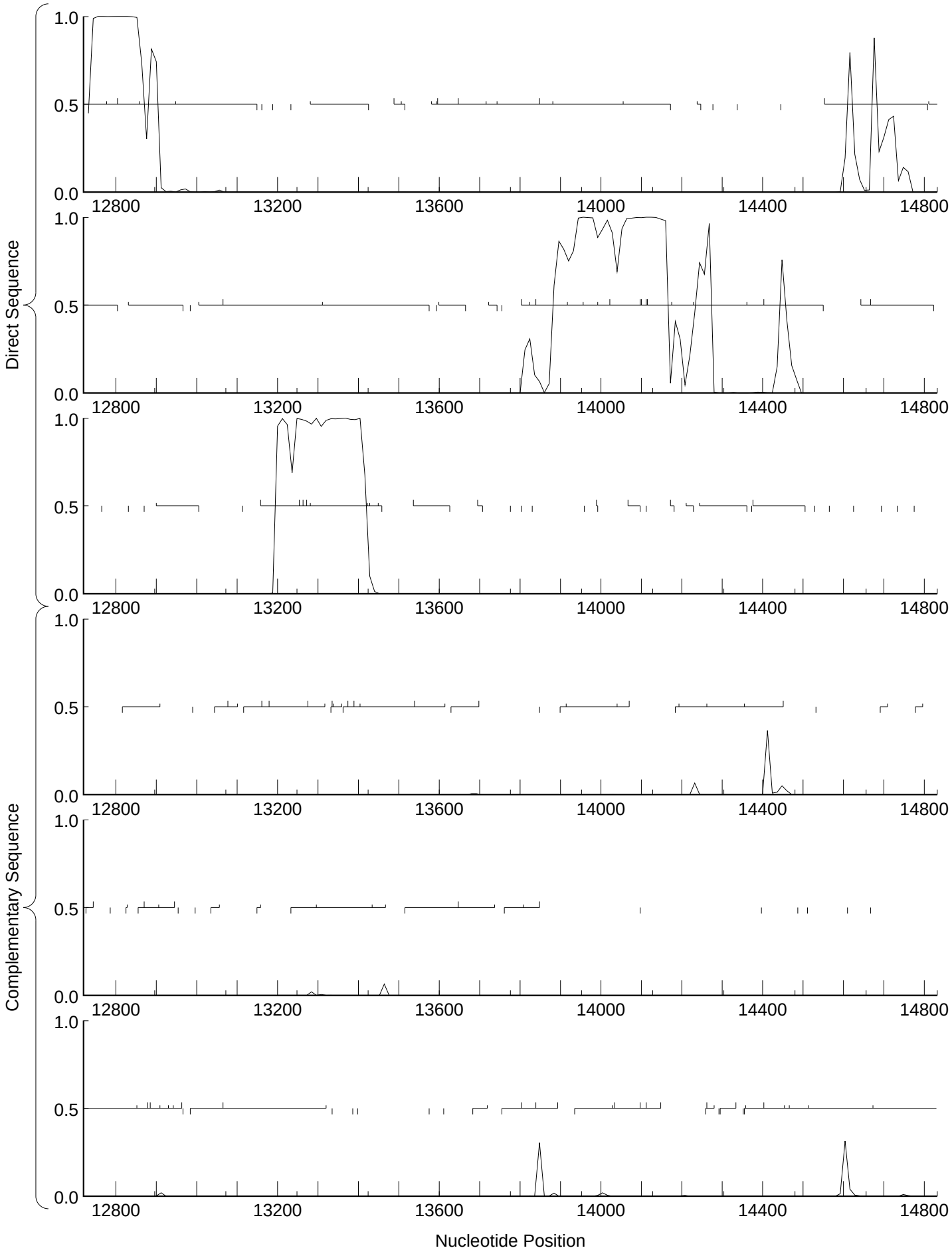

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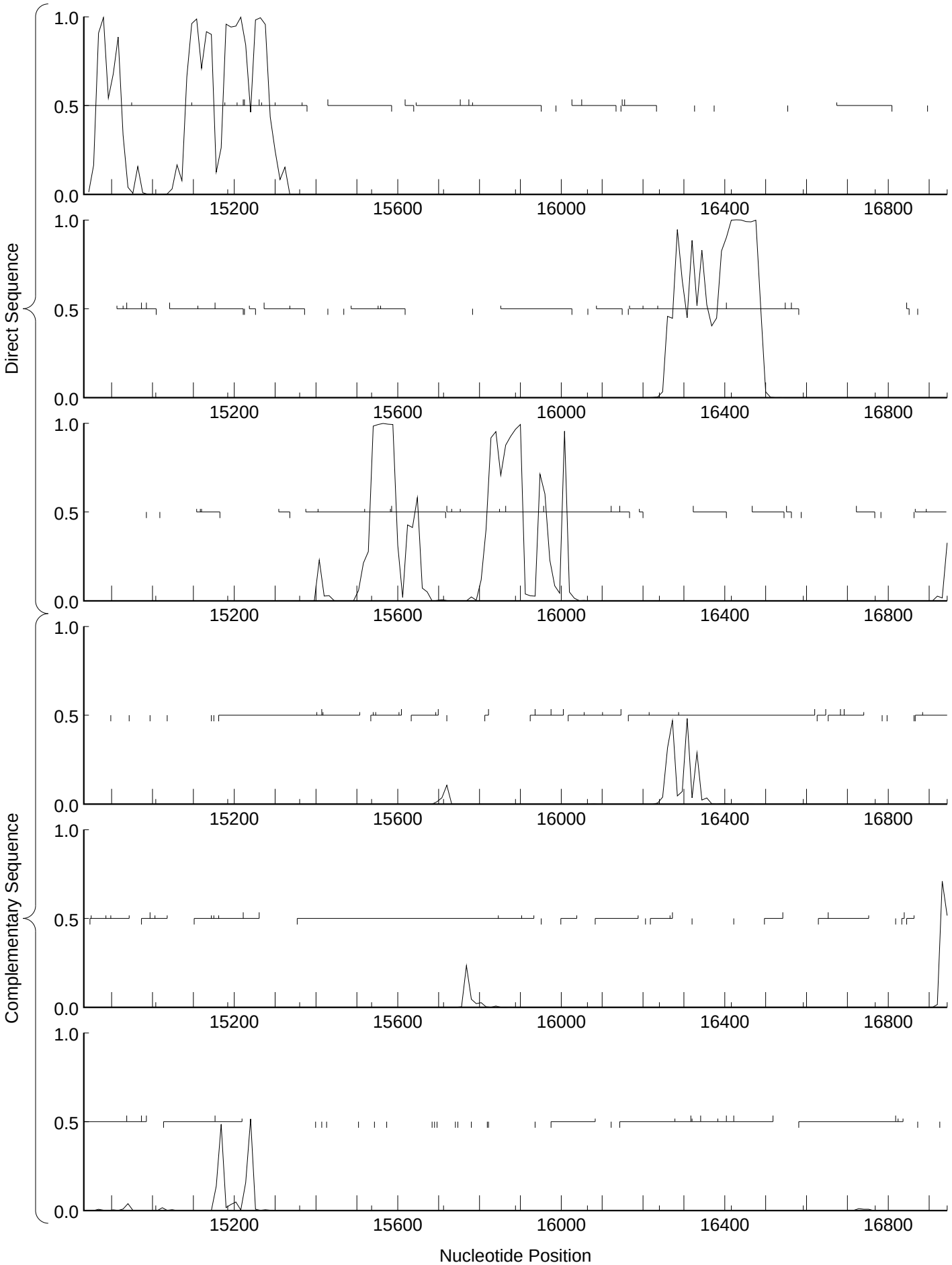

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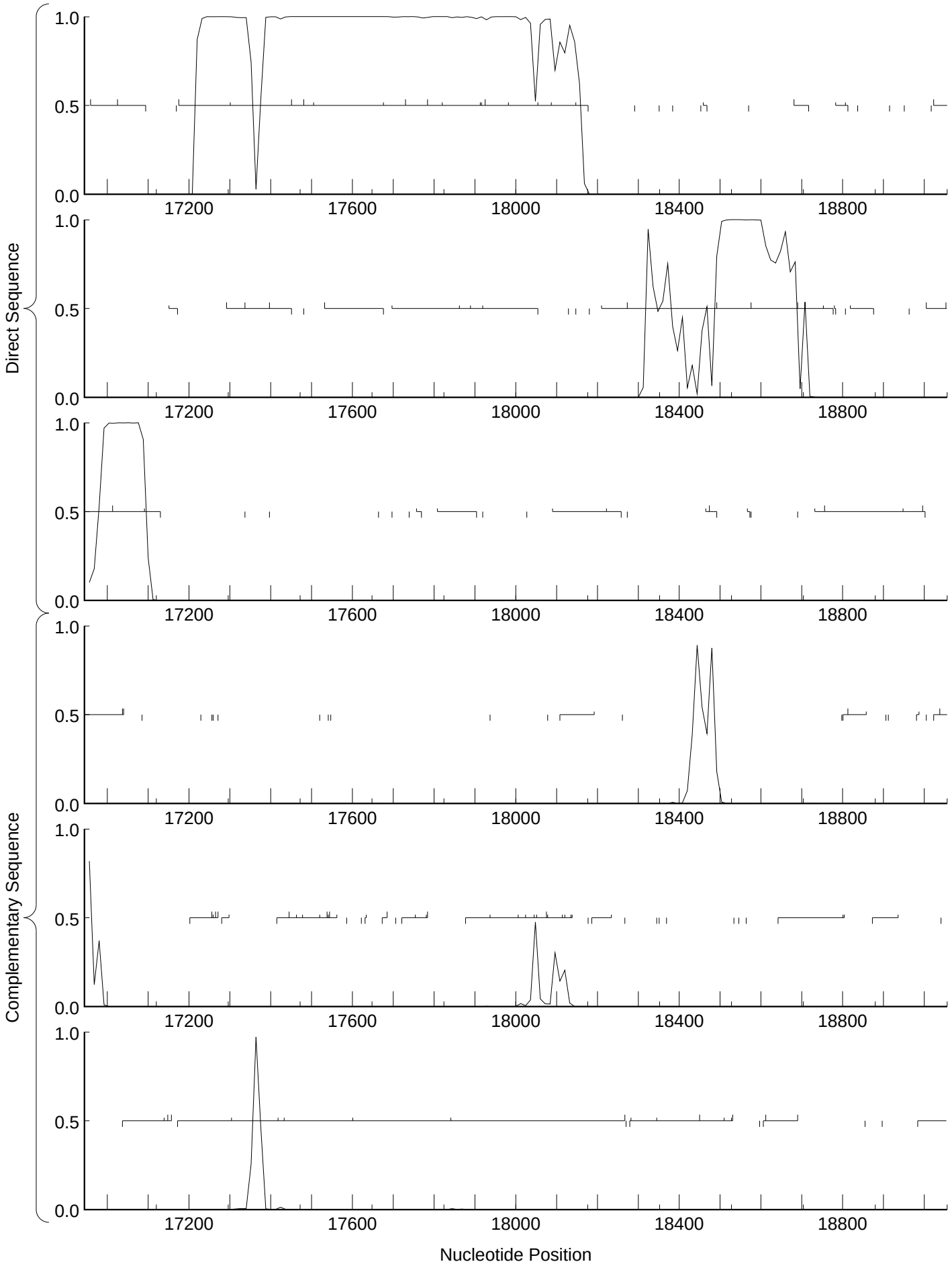

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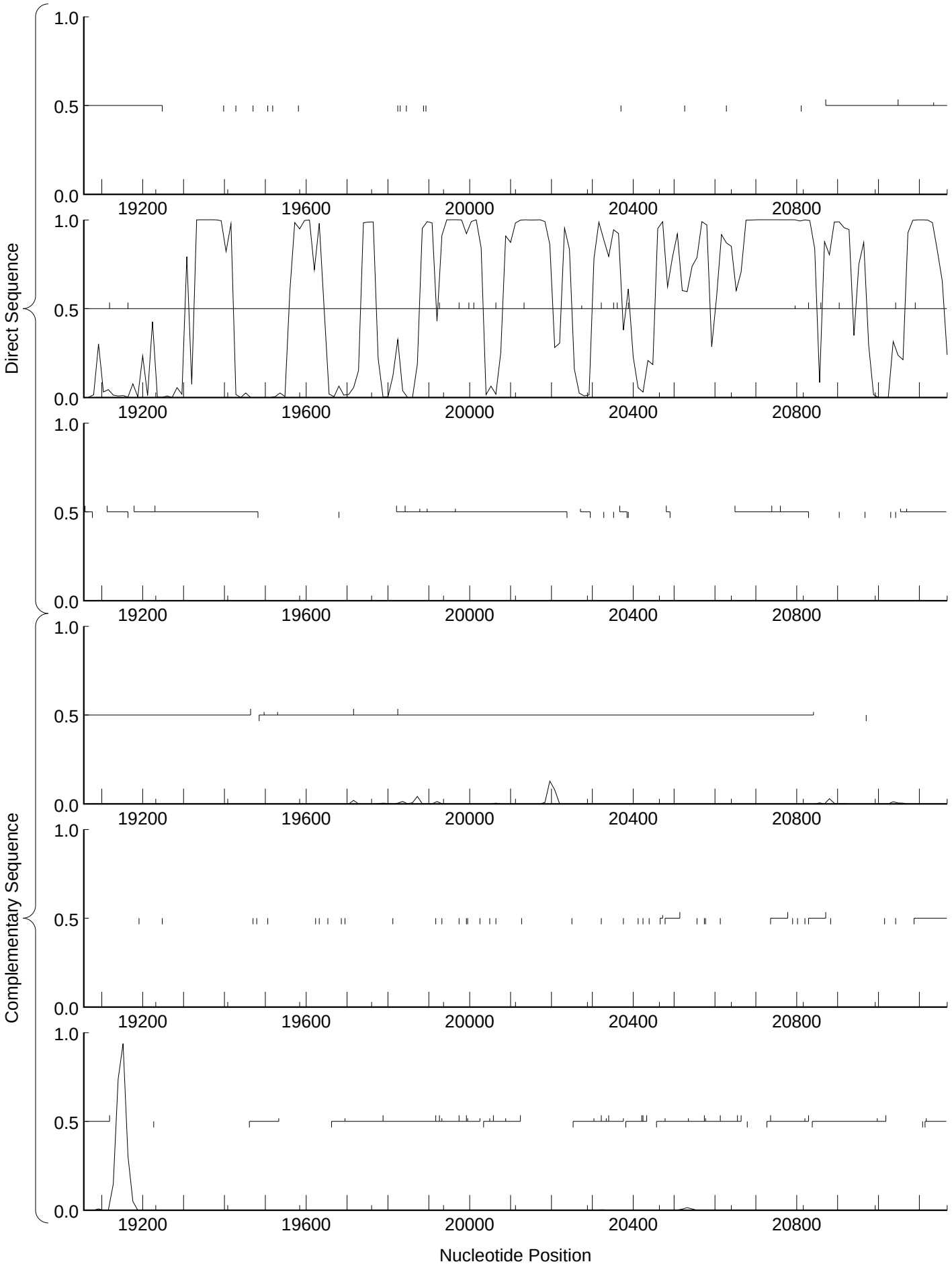

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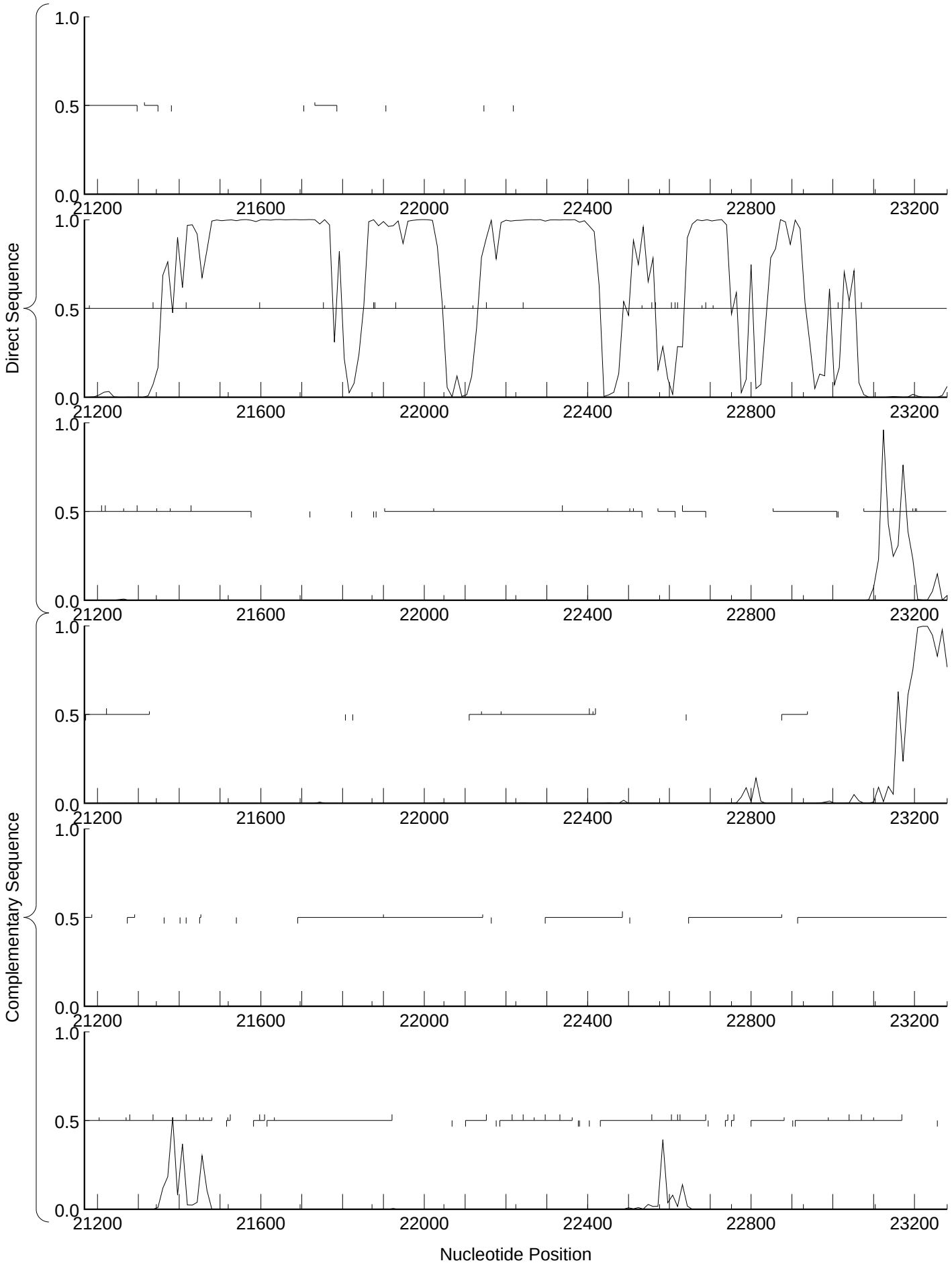
ge Constella complete sequence, 110169 bp including 4-base 3’ overhang (ATCC), Cluster J, Order 4, Window 96, Step 12, 10/54



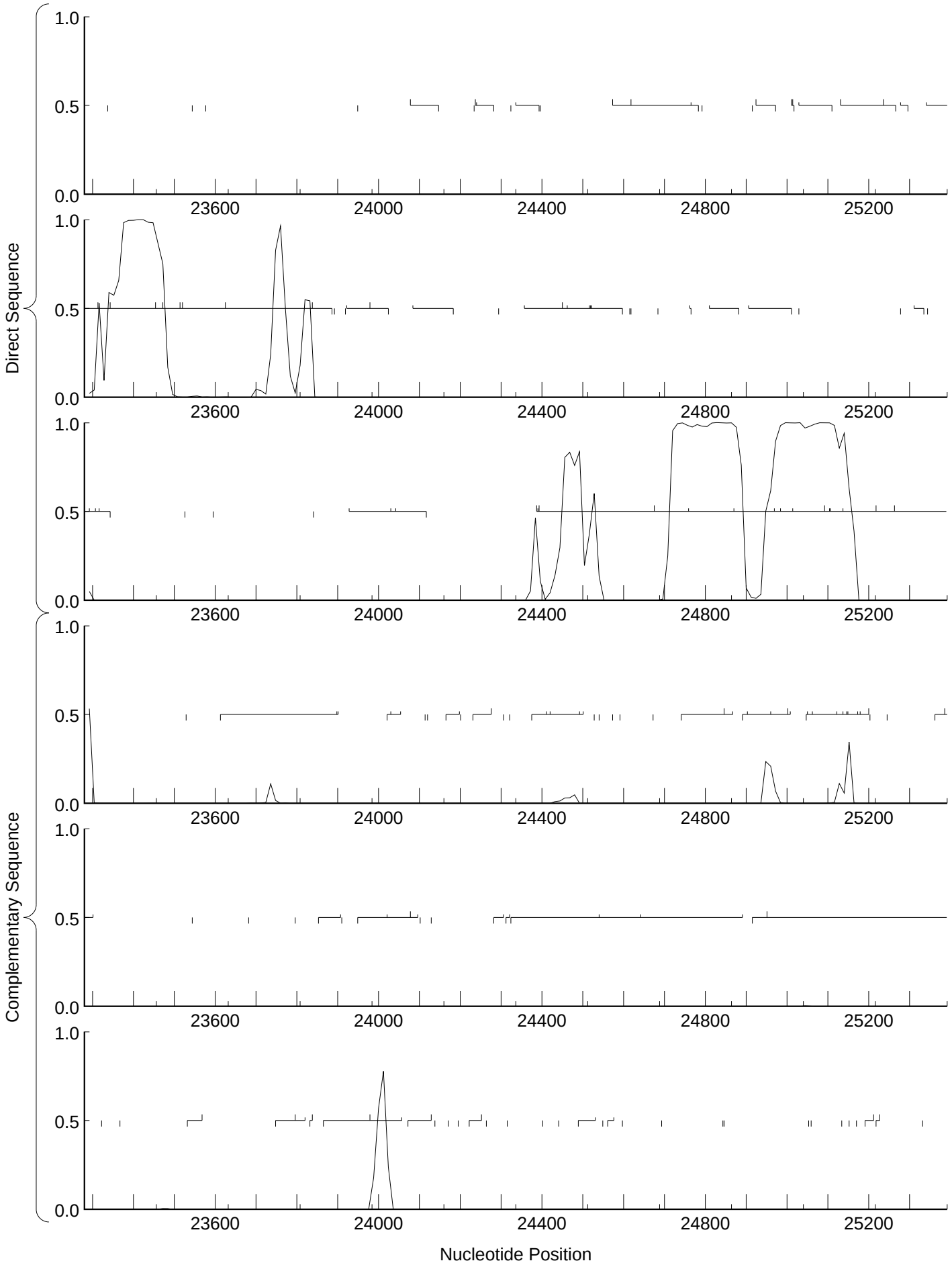
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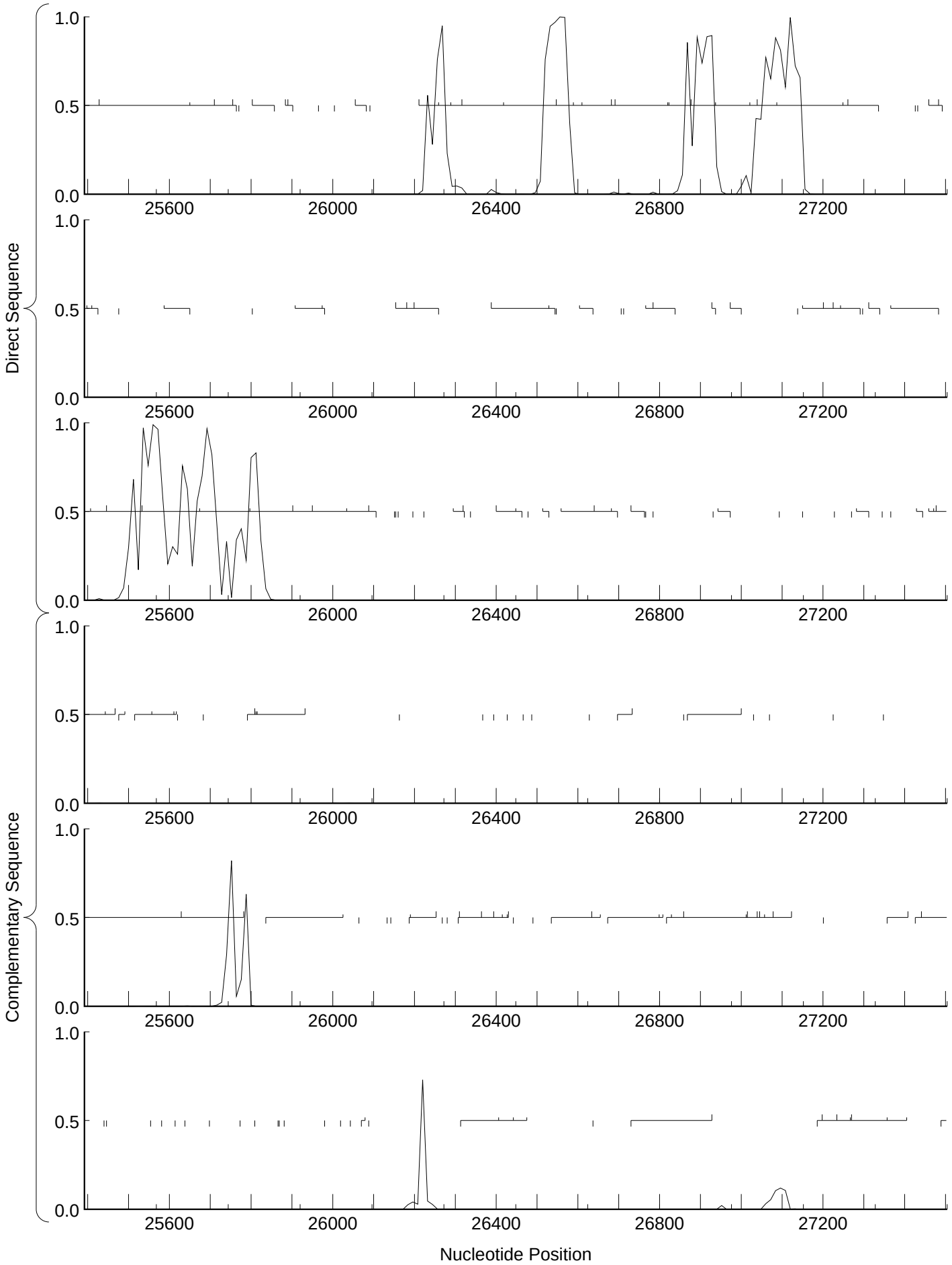
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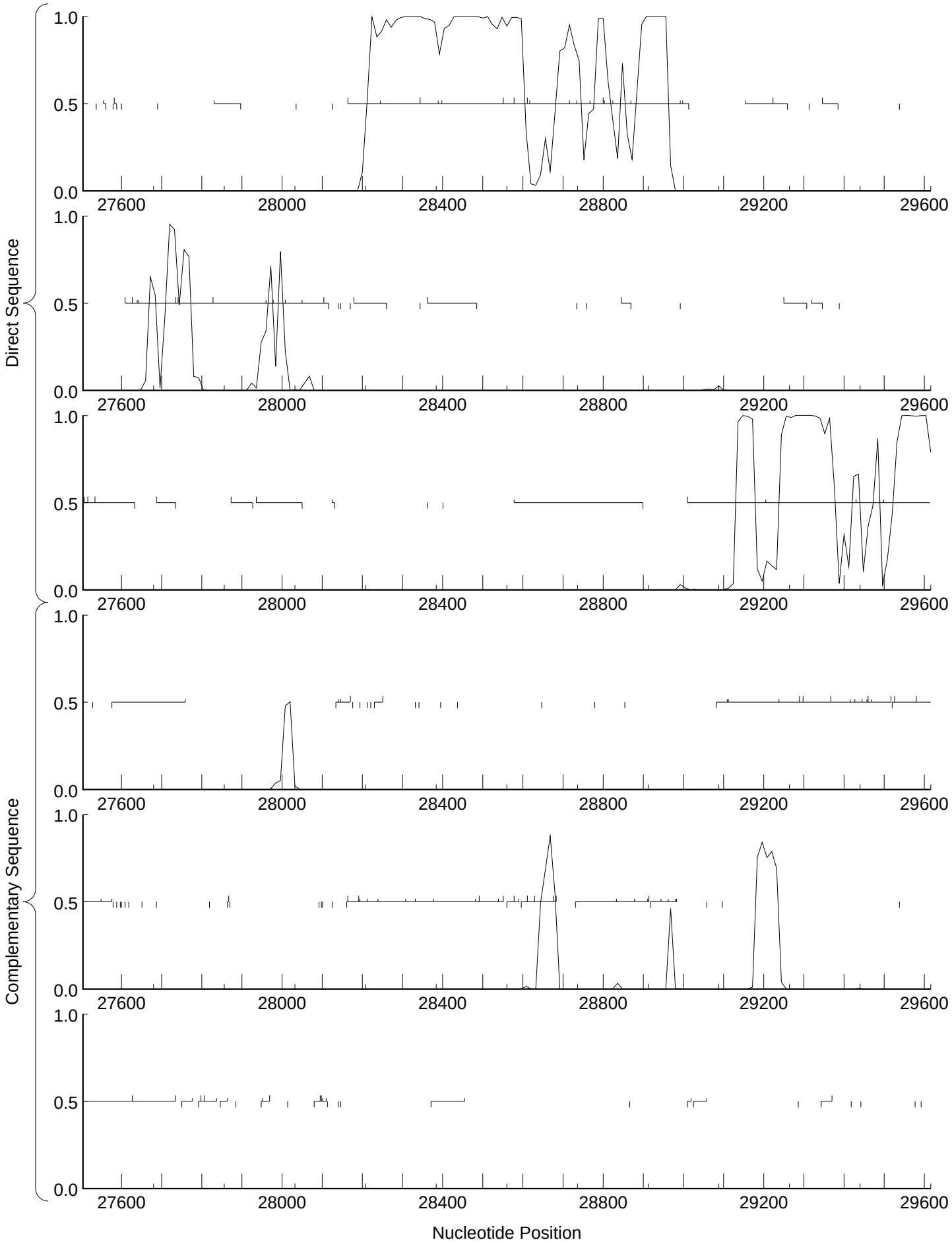
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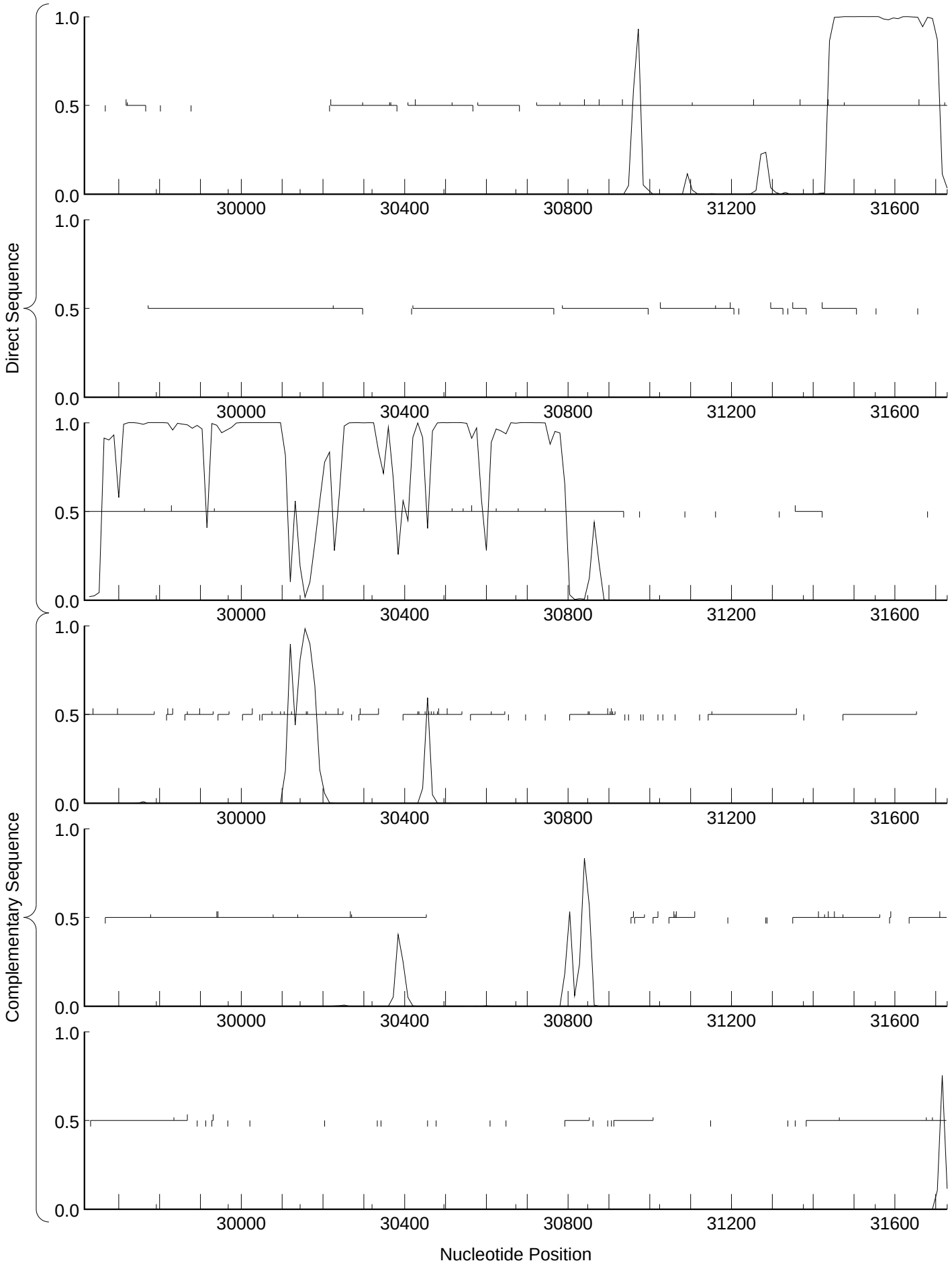
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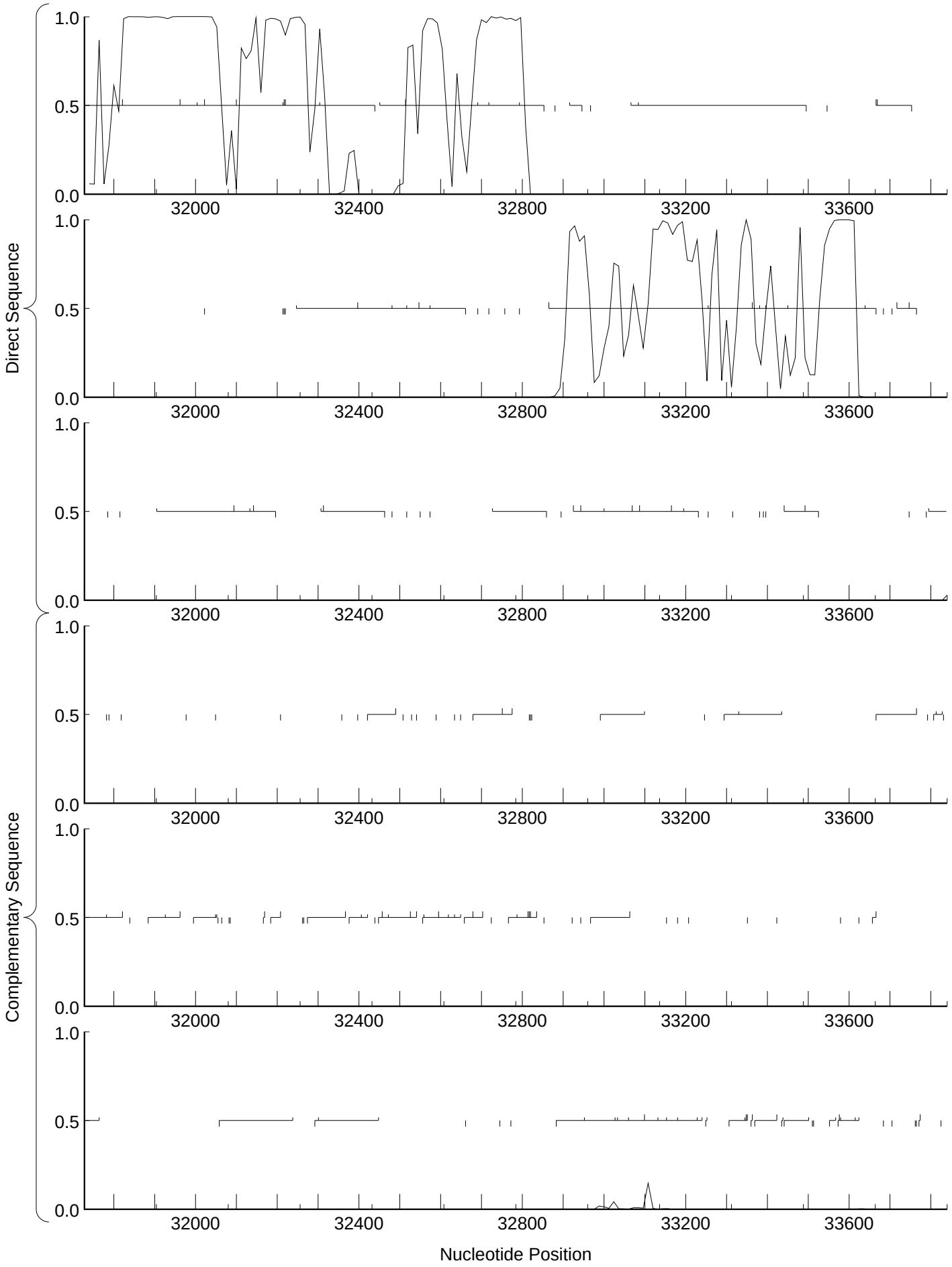
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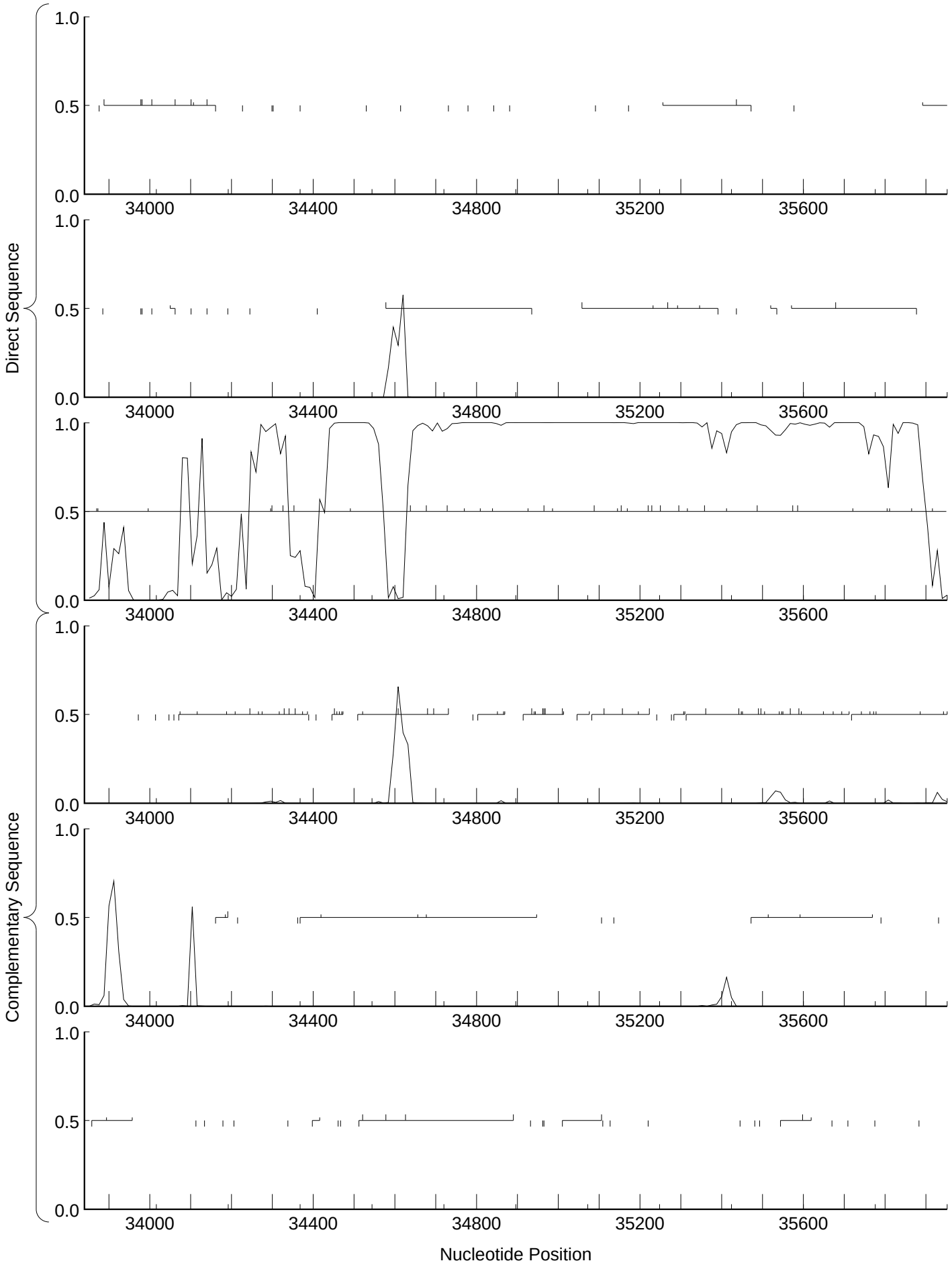
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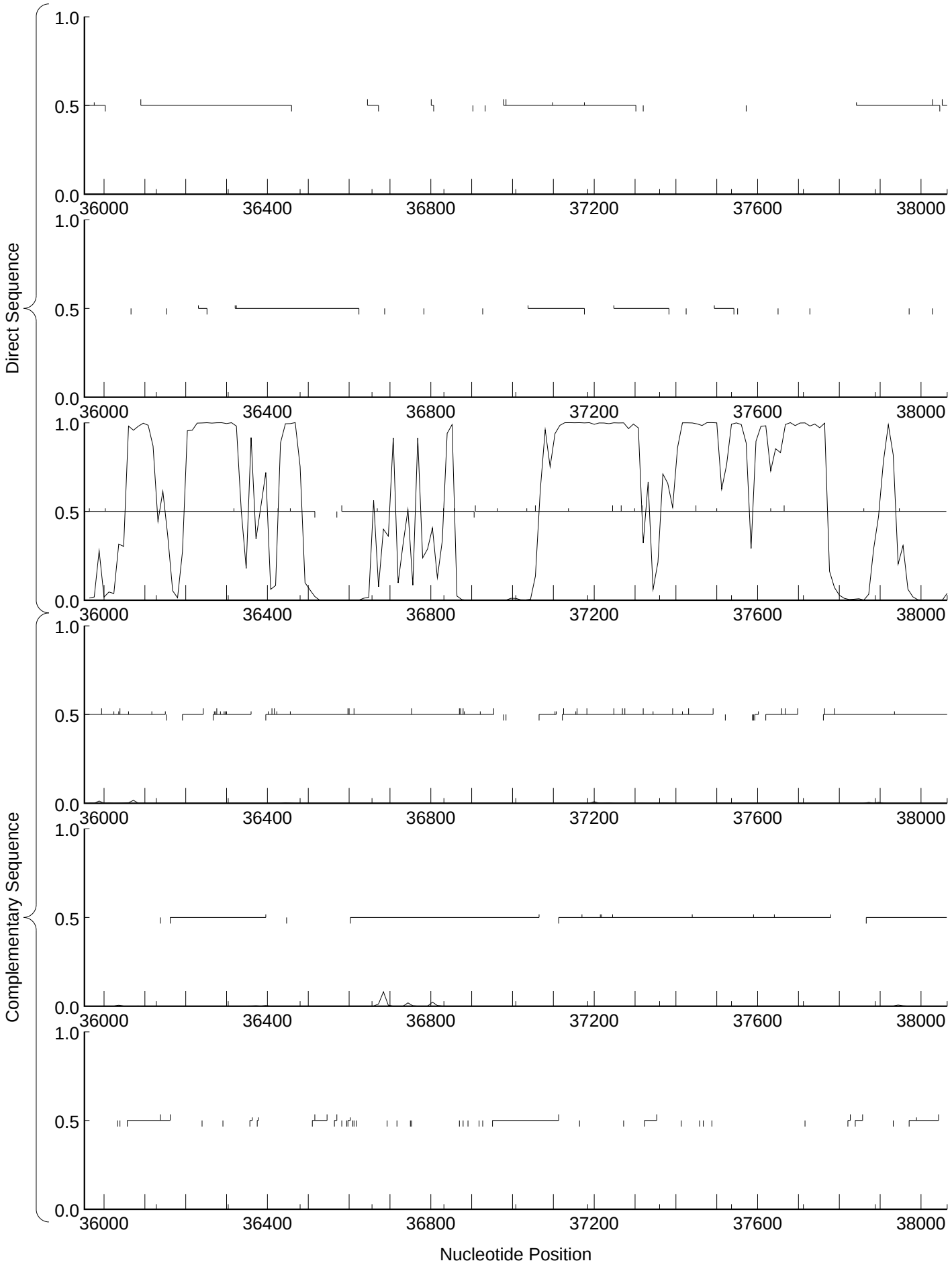


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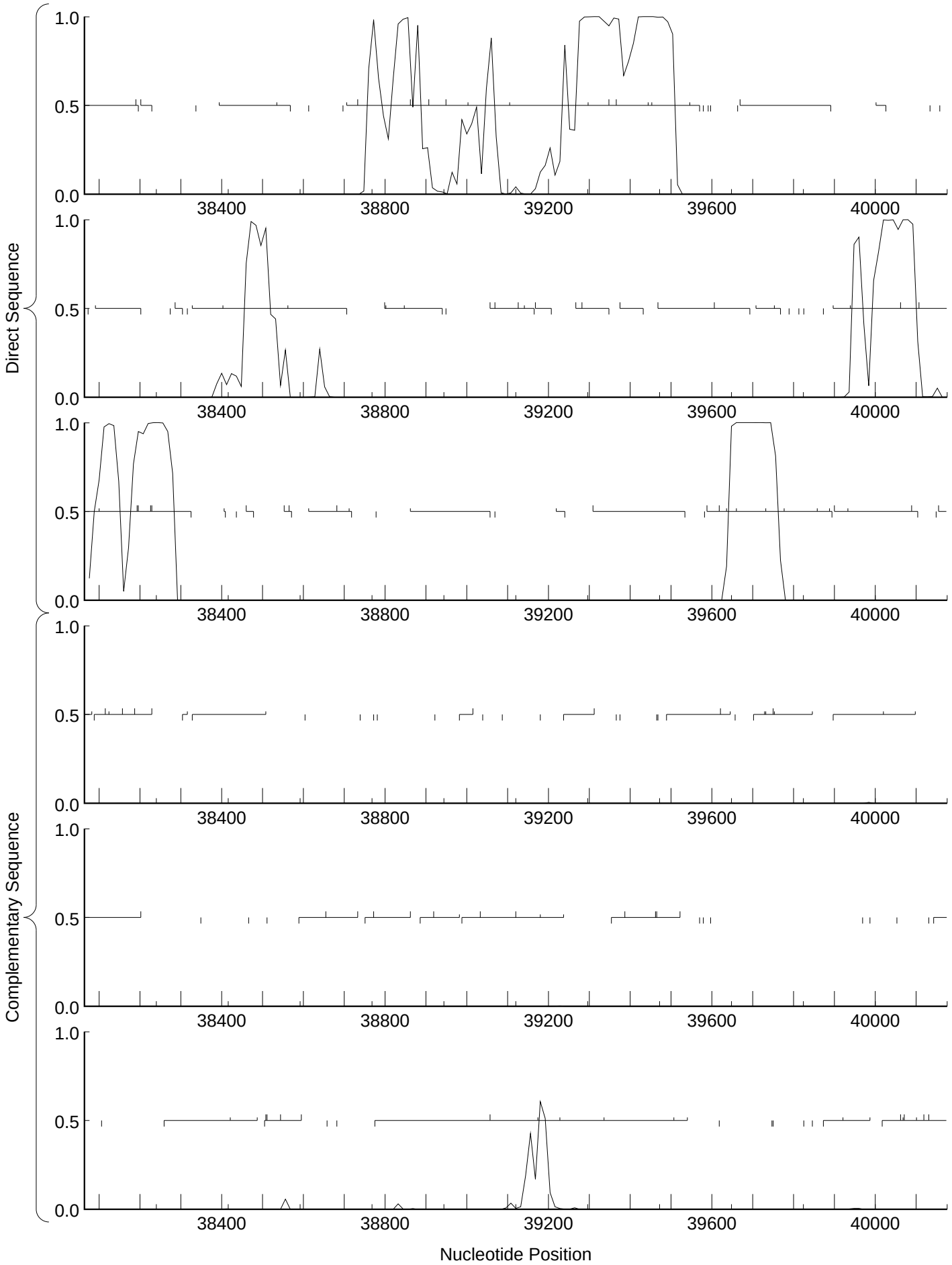


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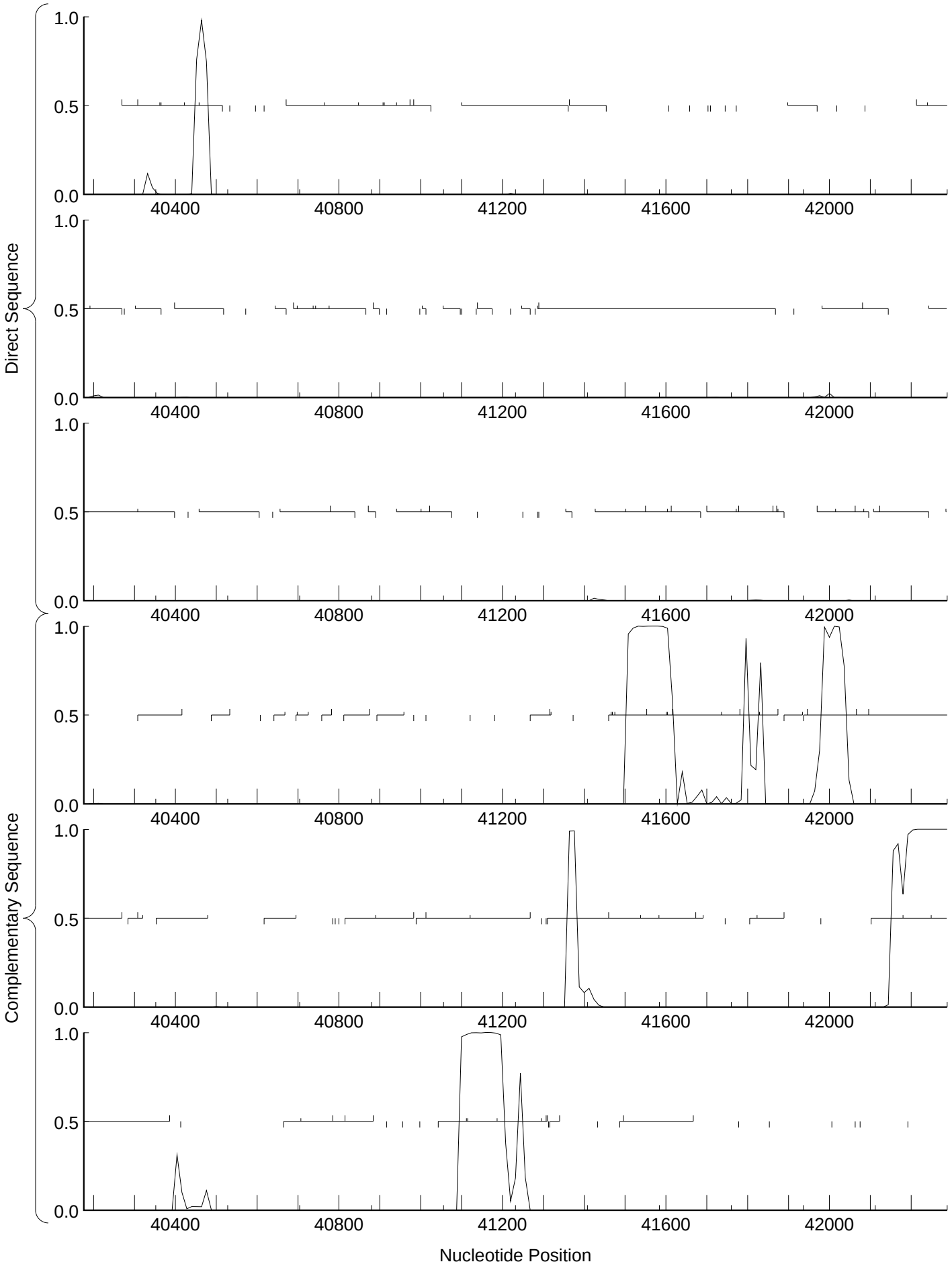


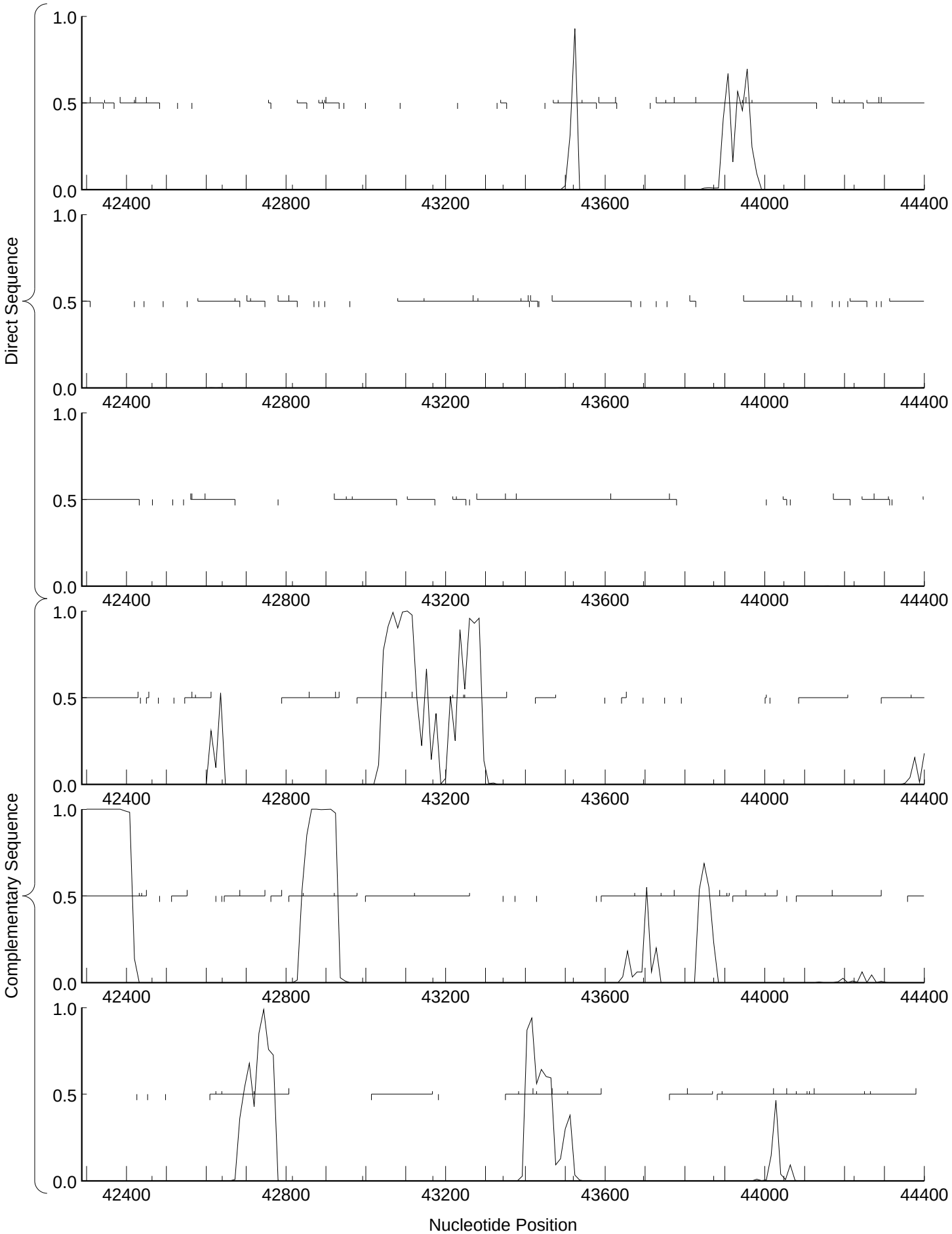

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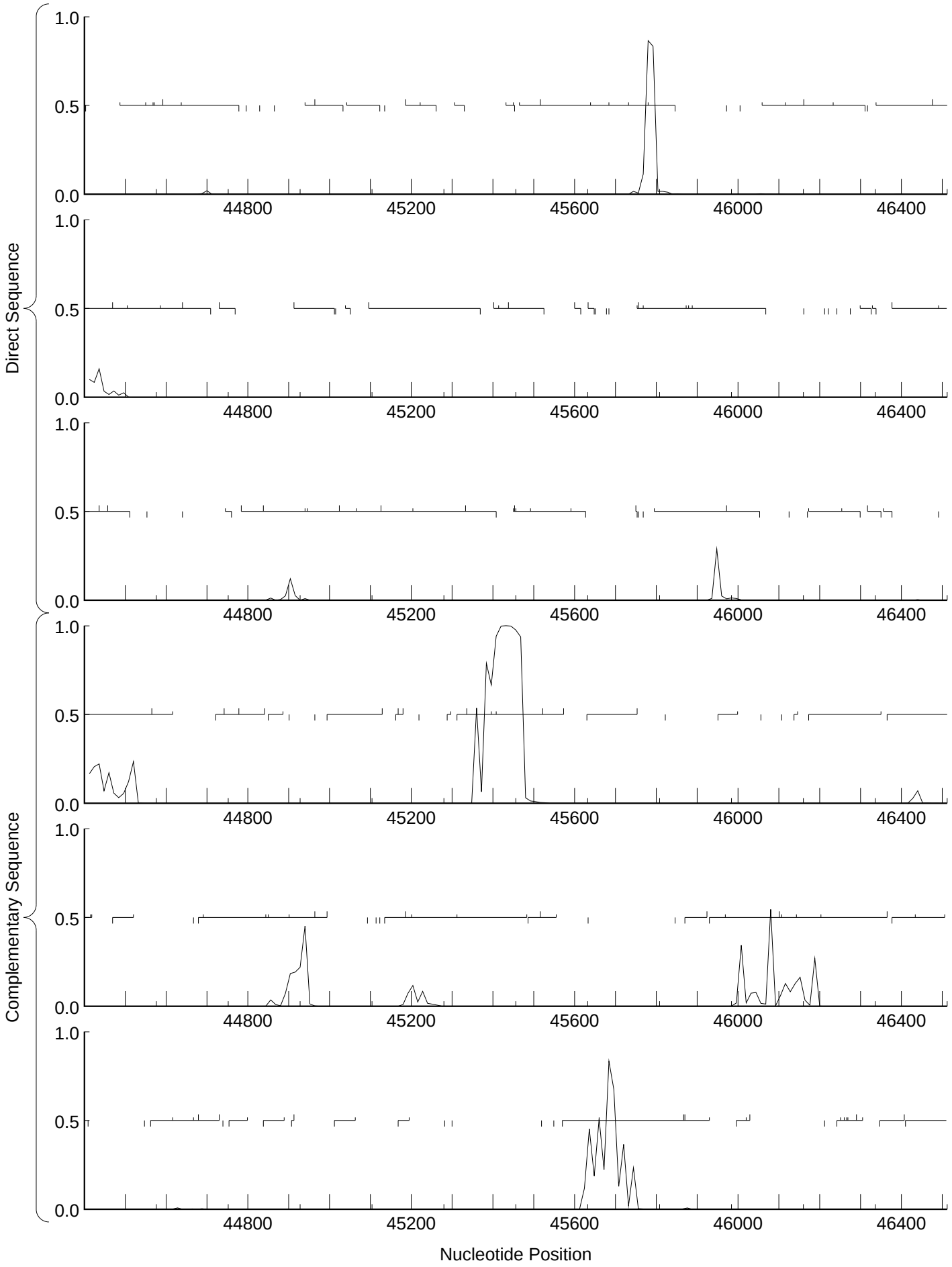


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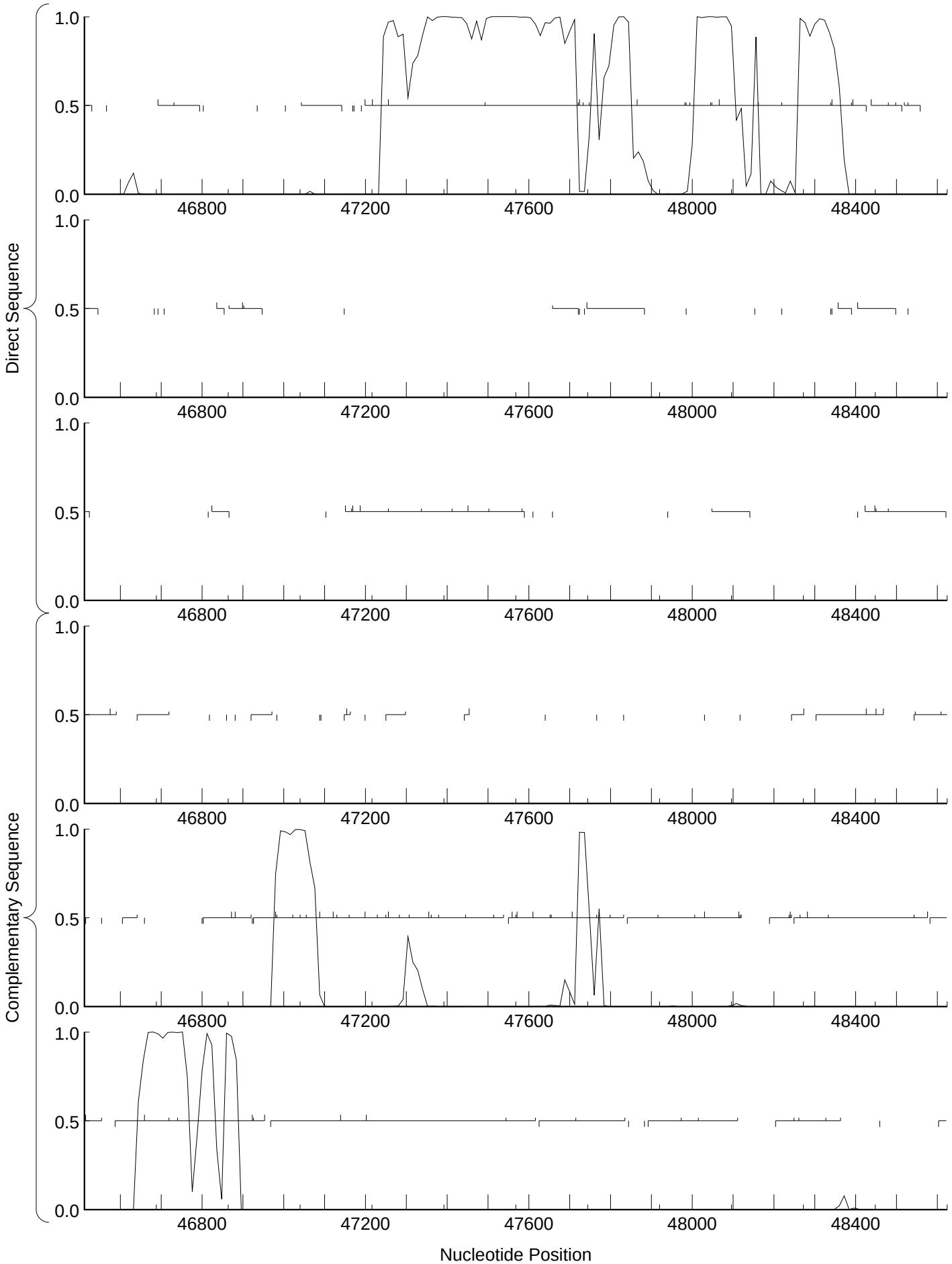



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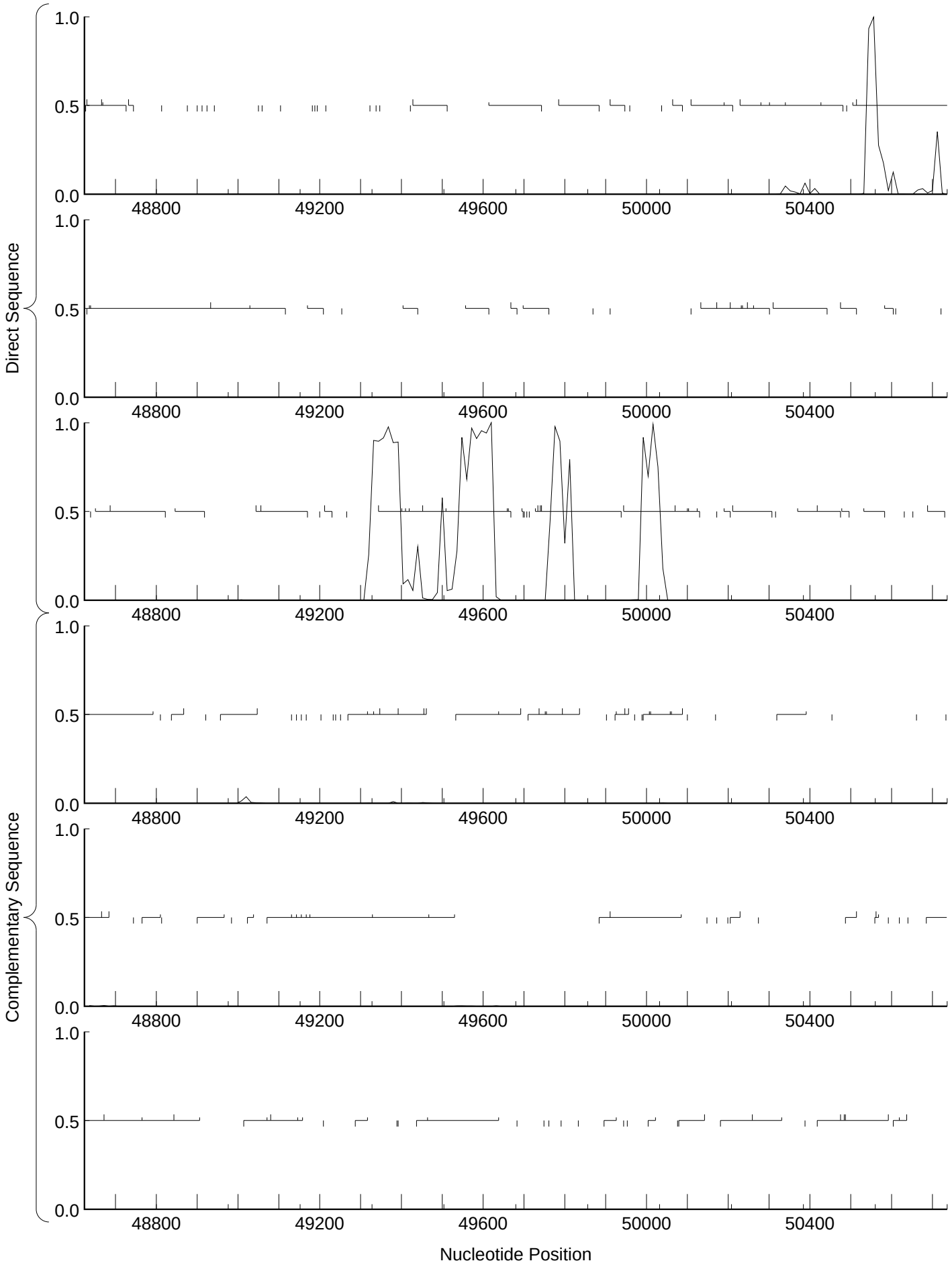
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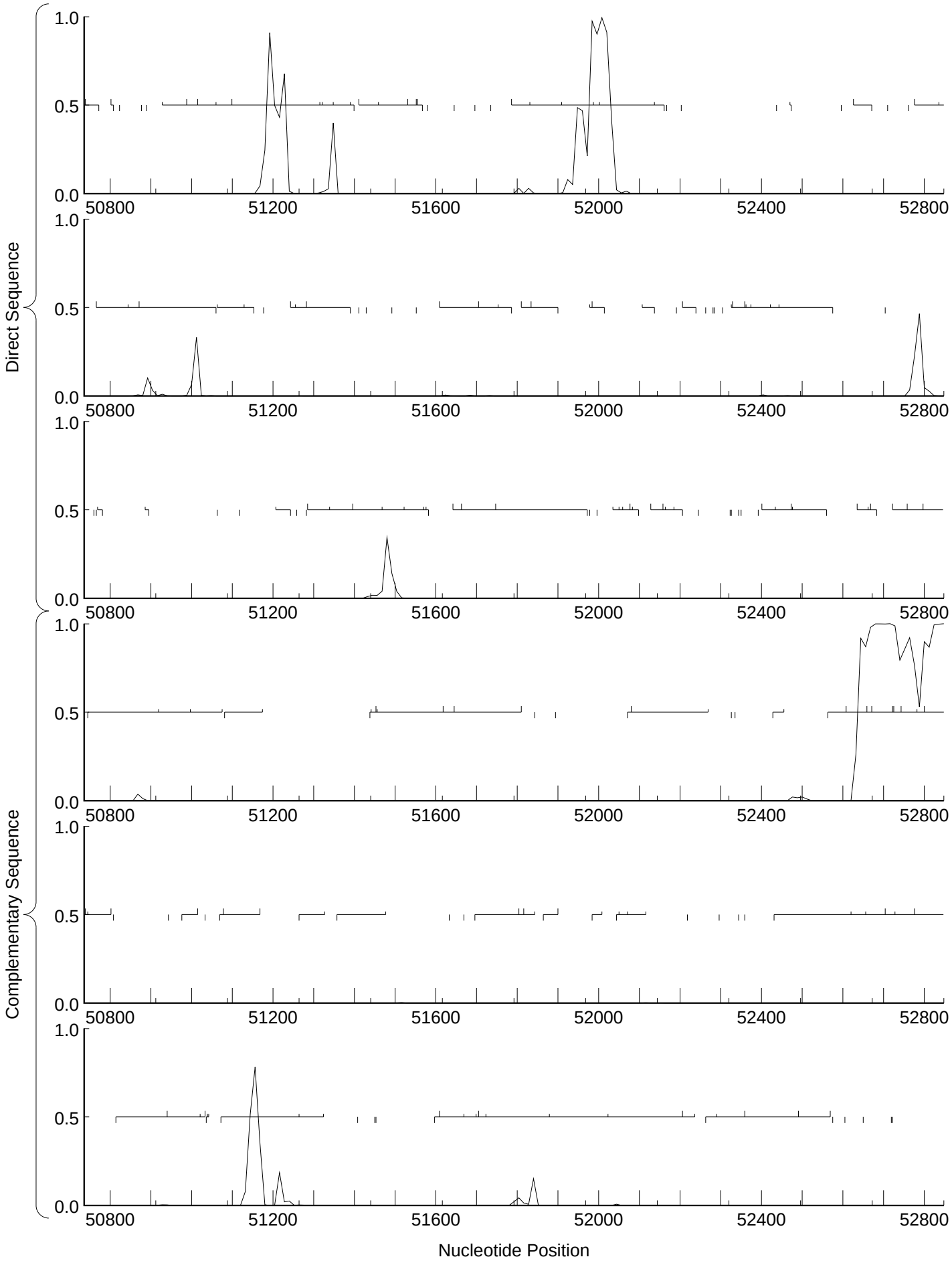
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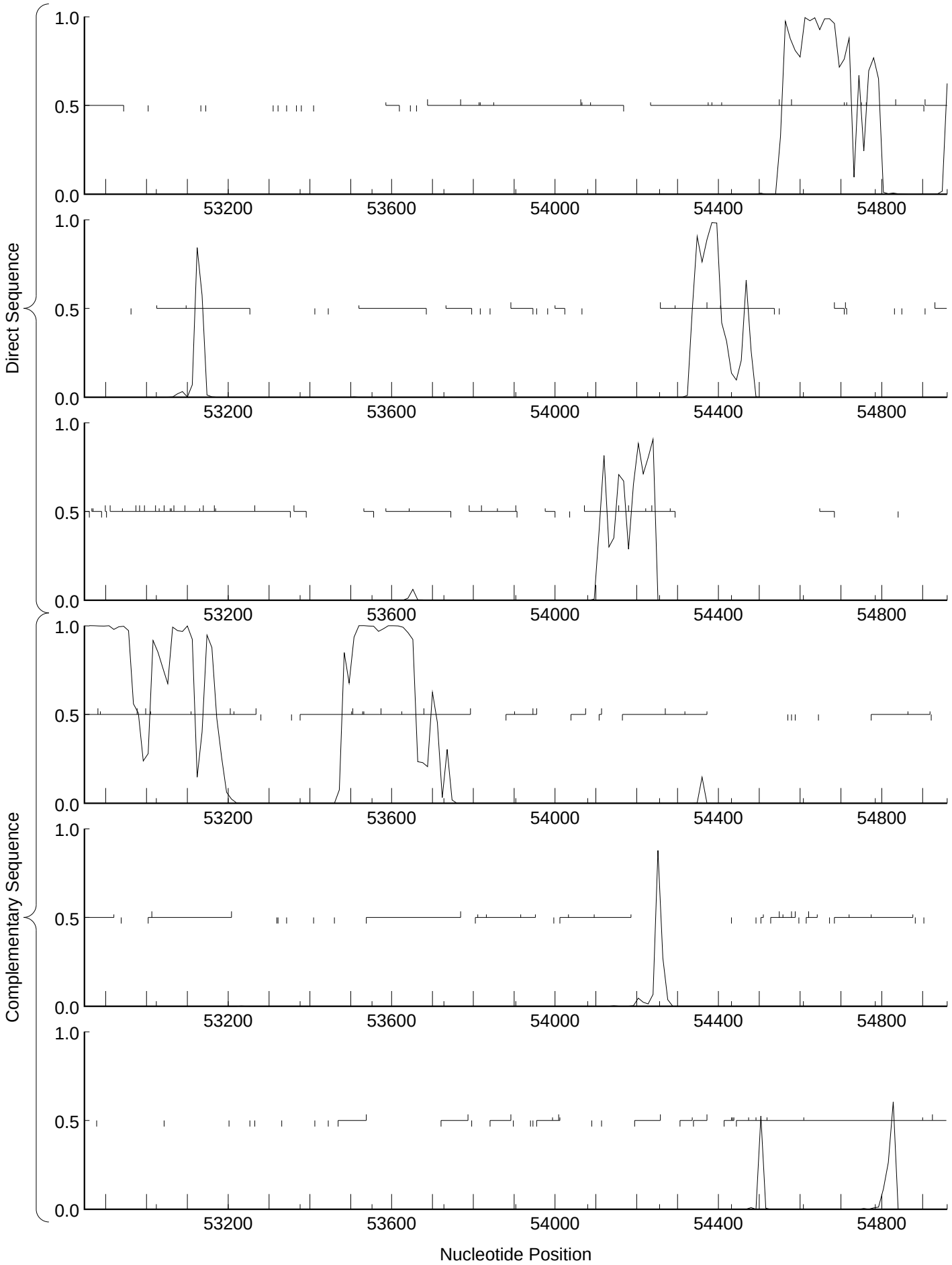
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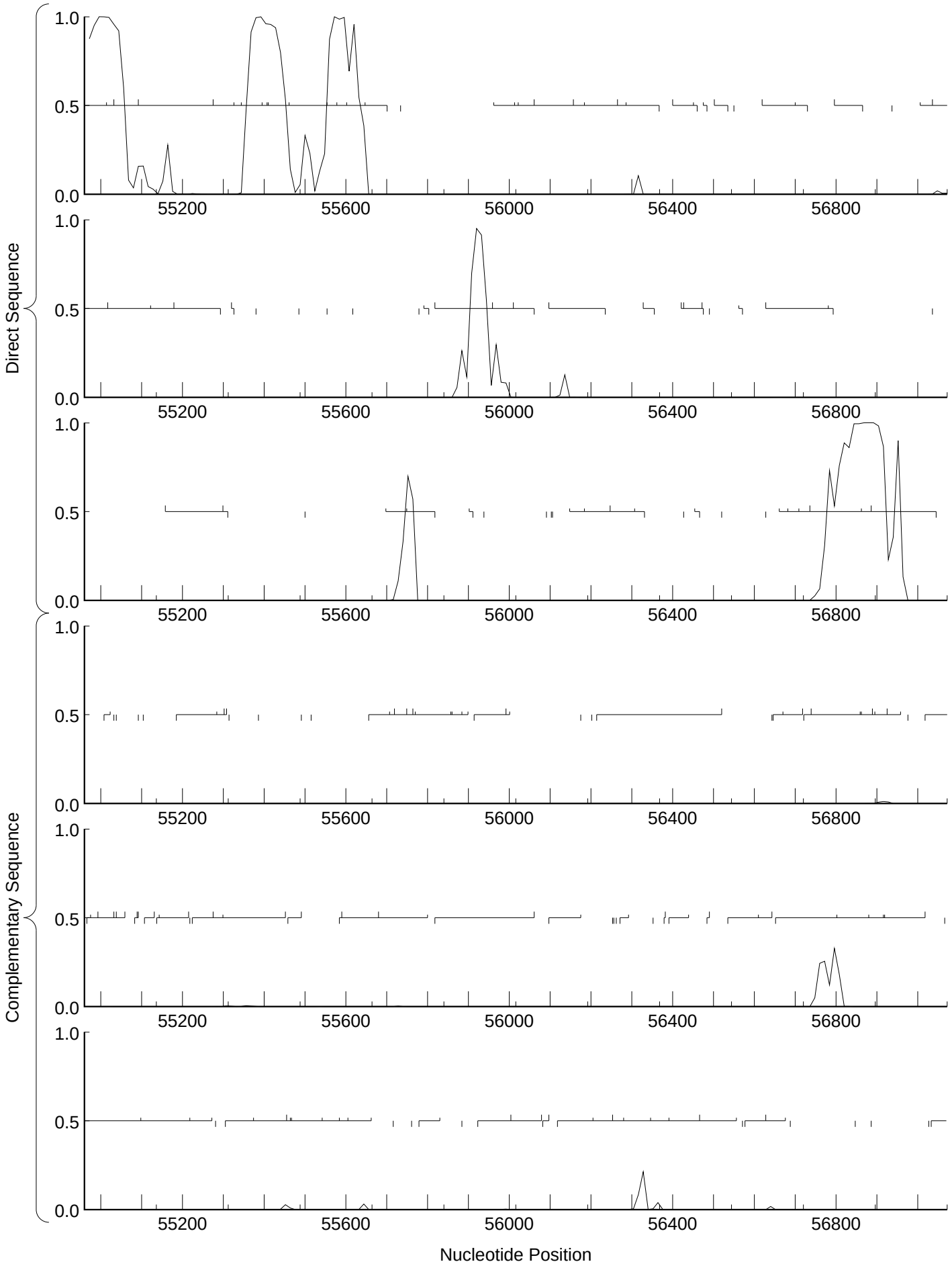


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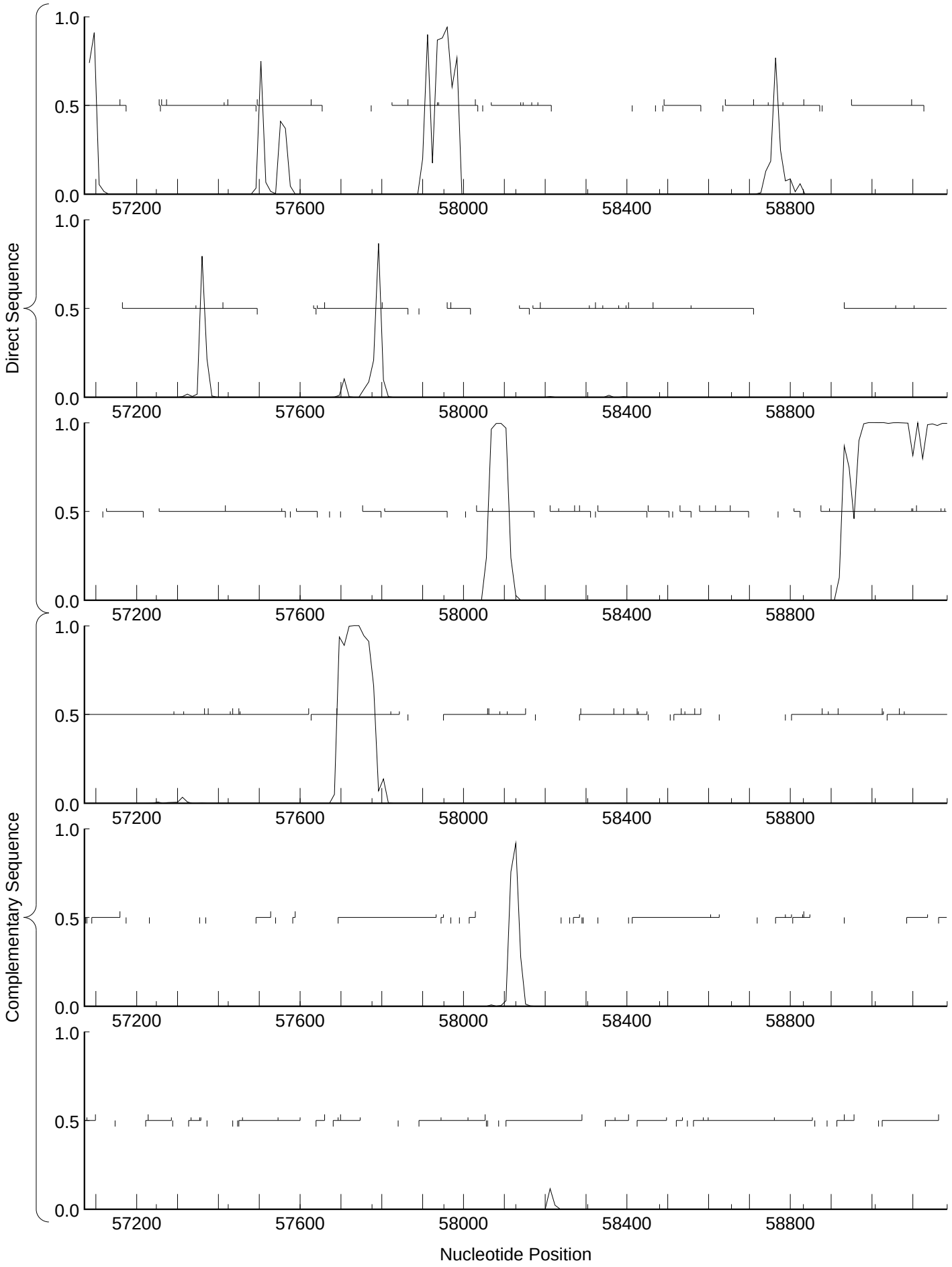


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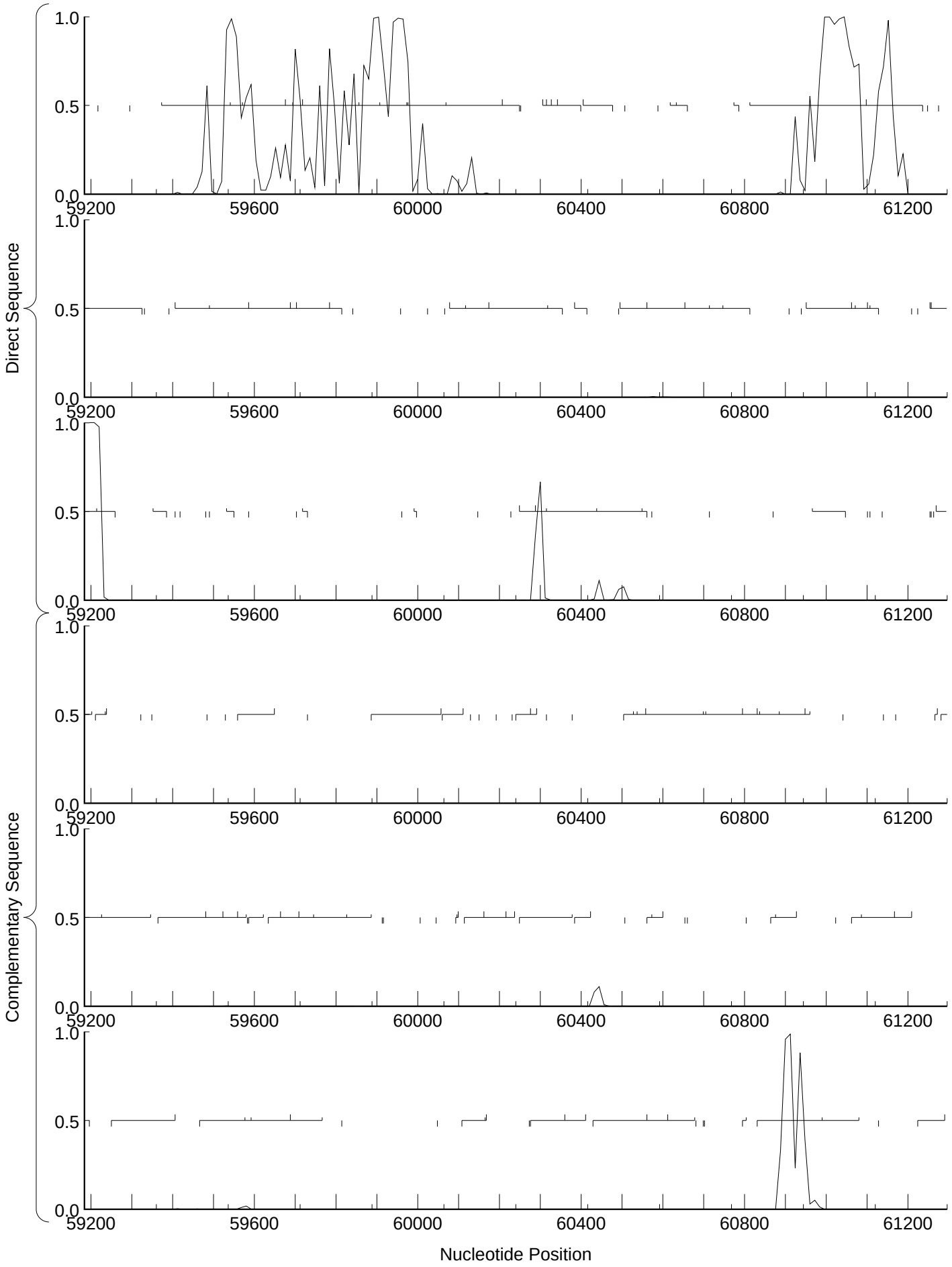



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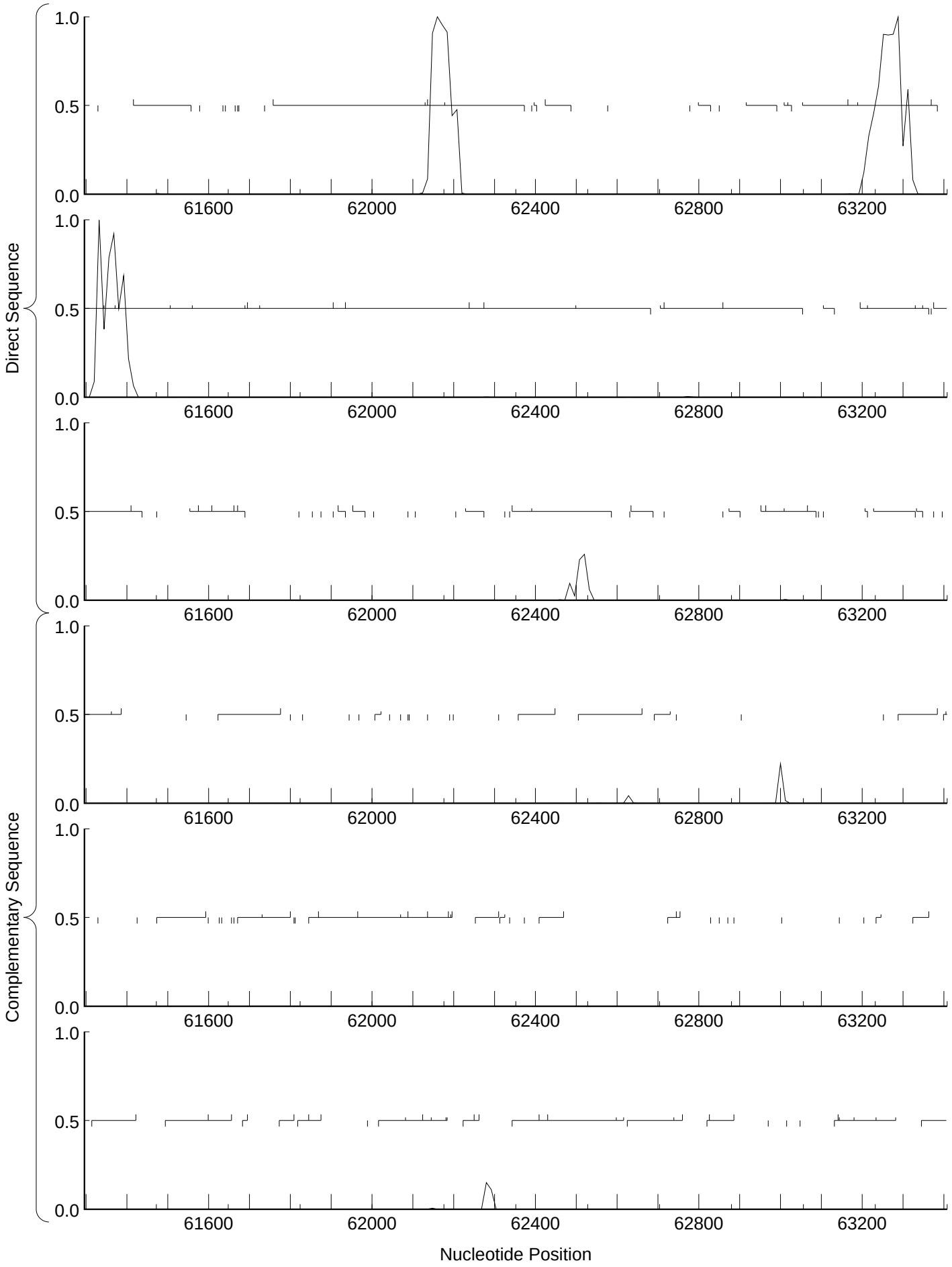
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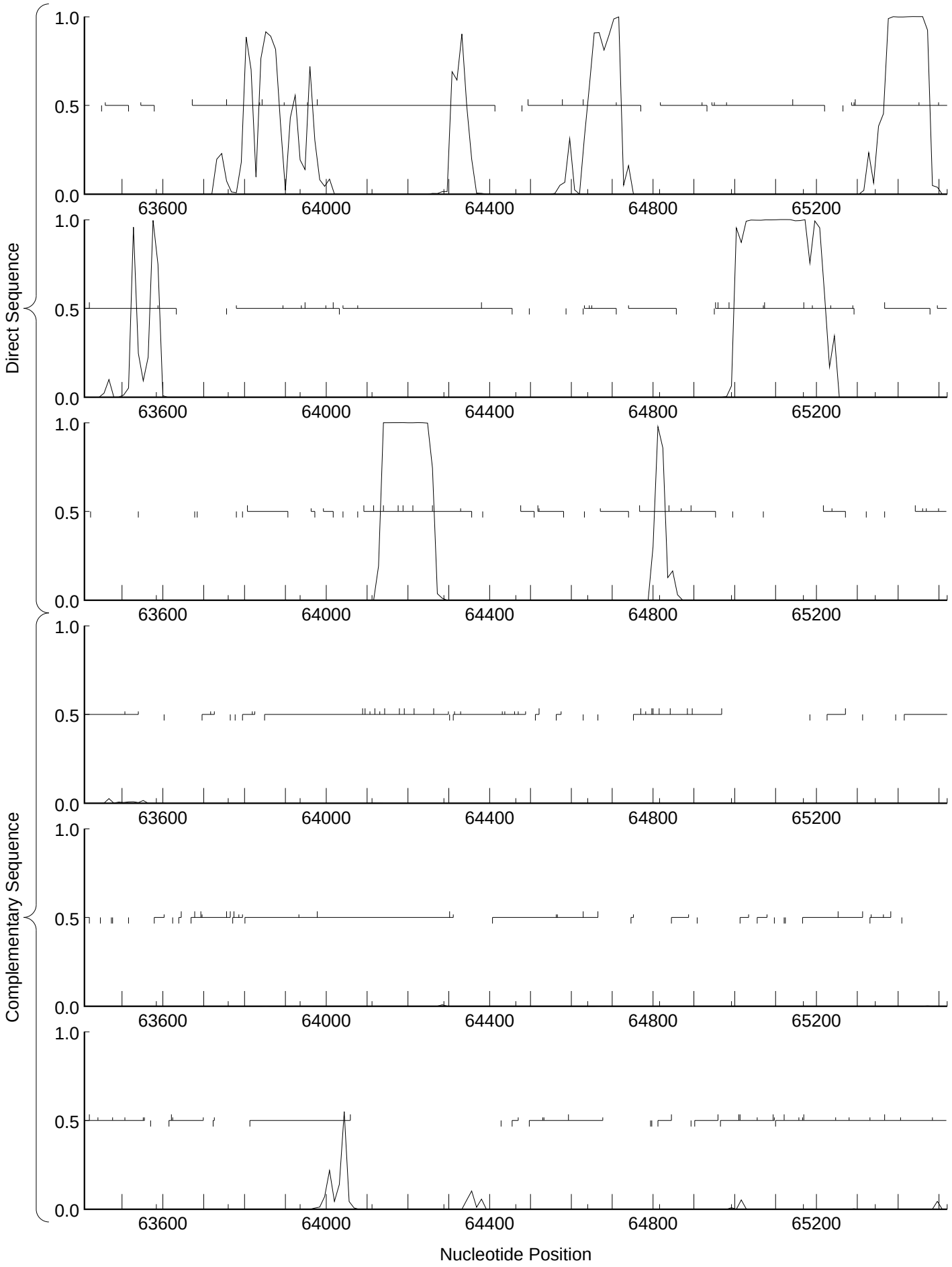
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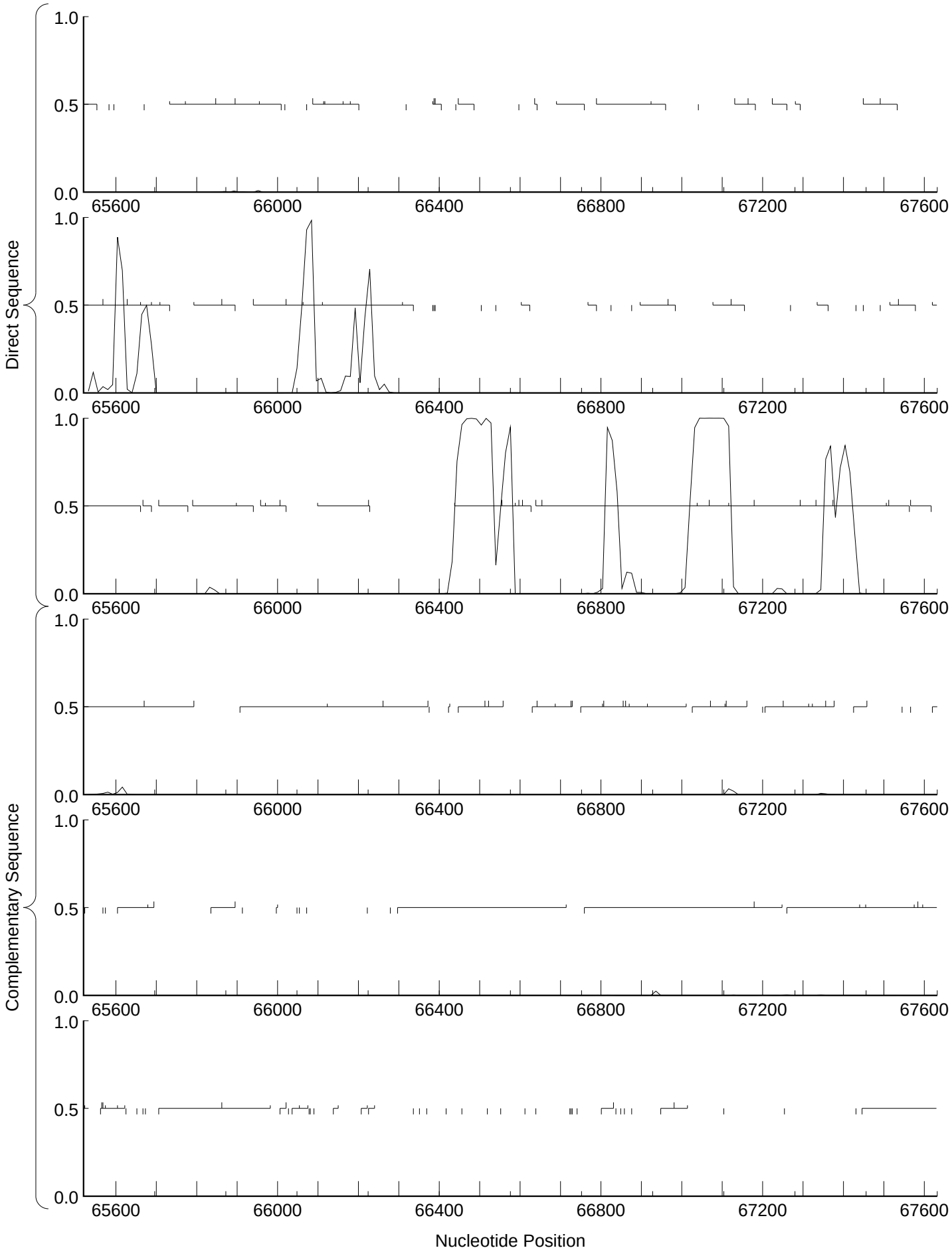
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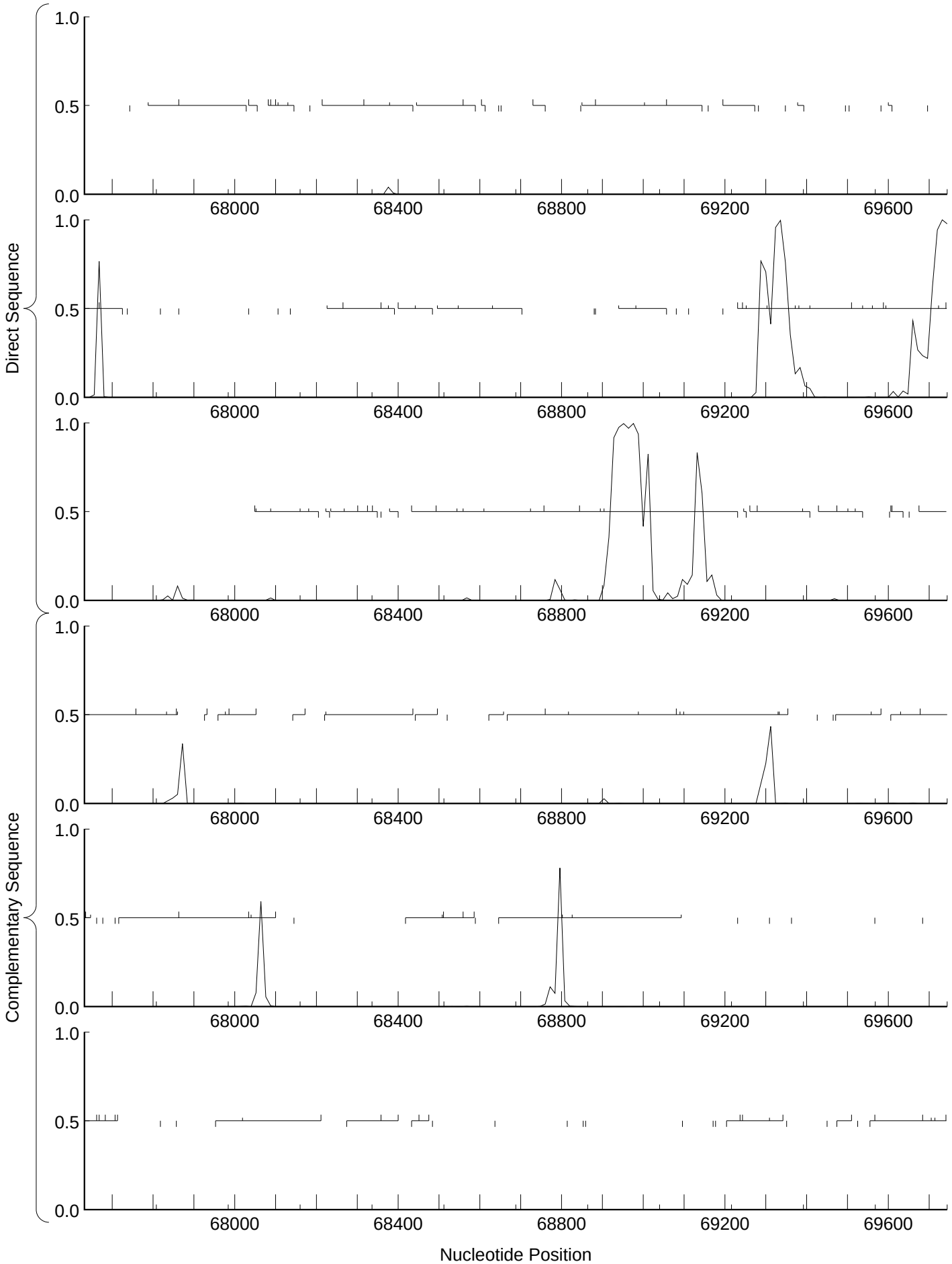
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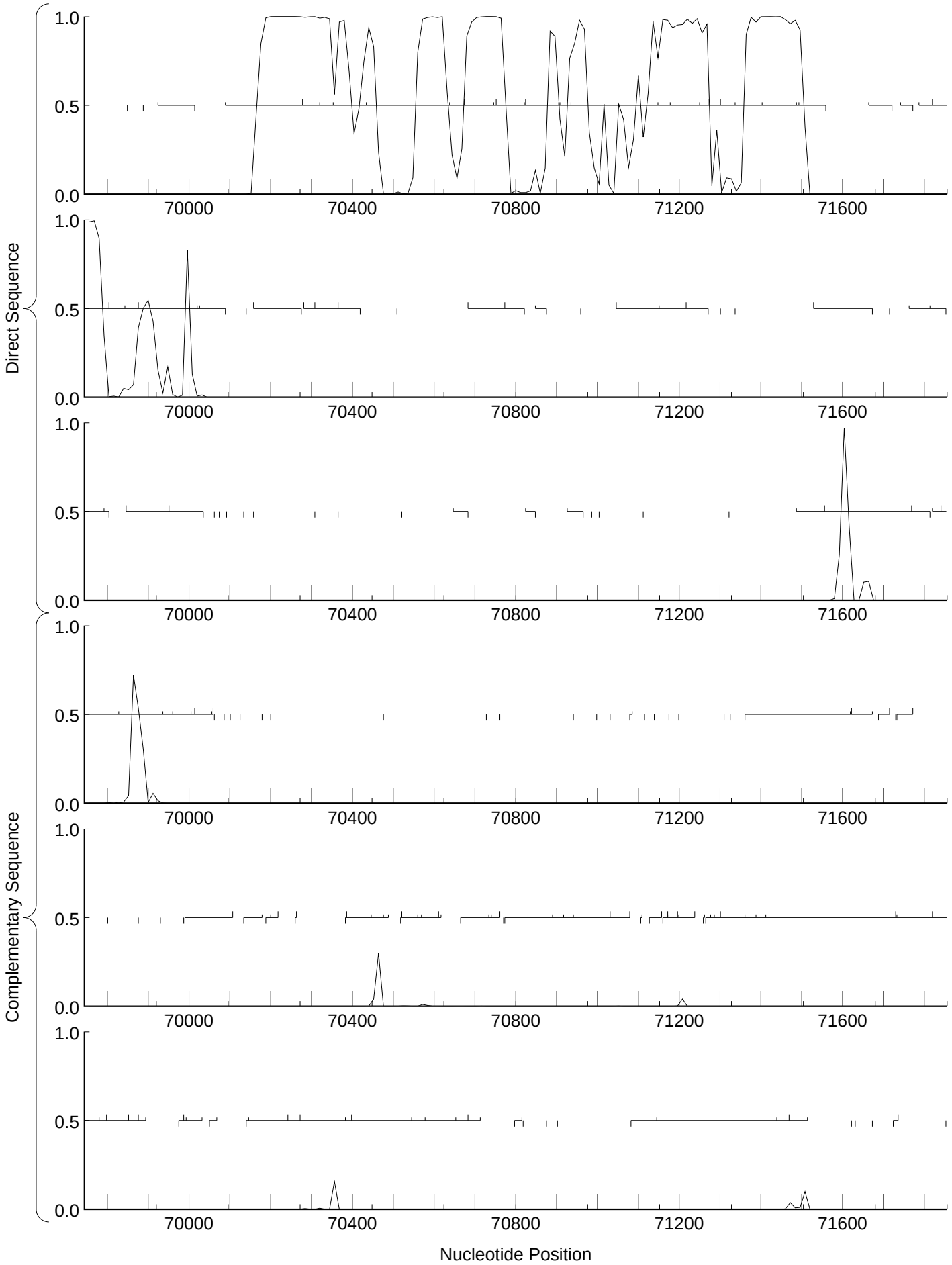
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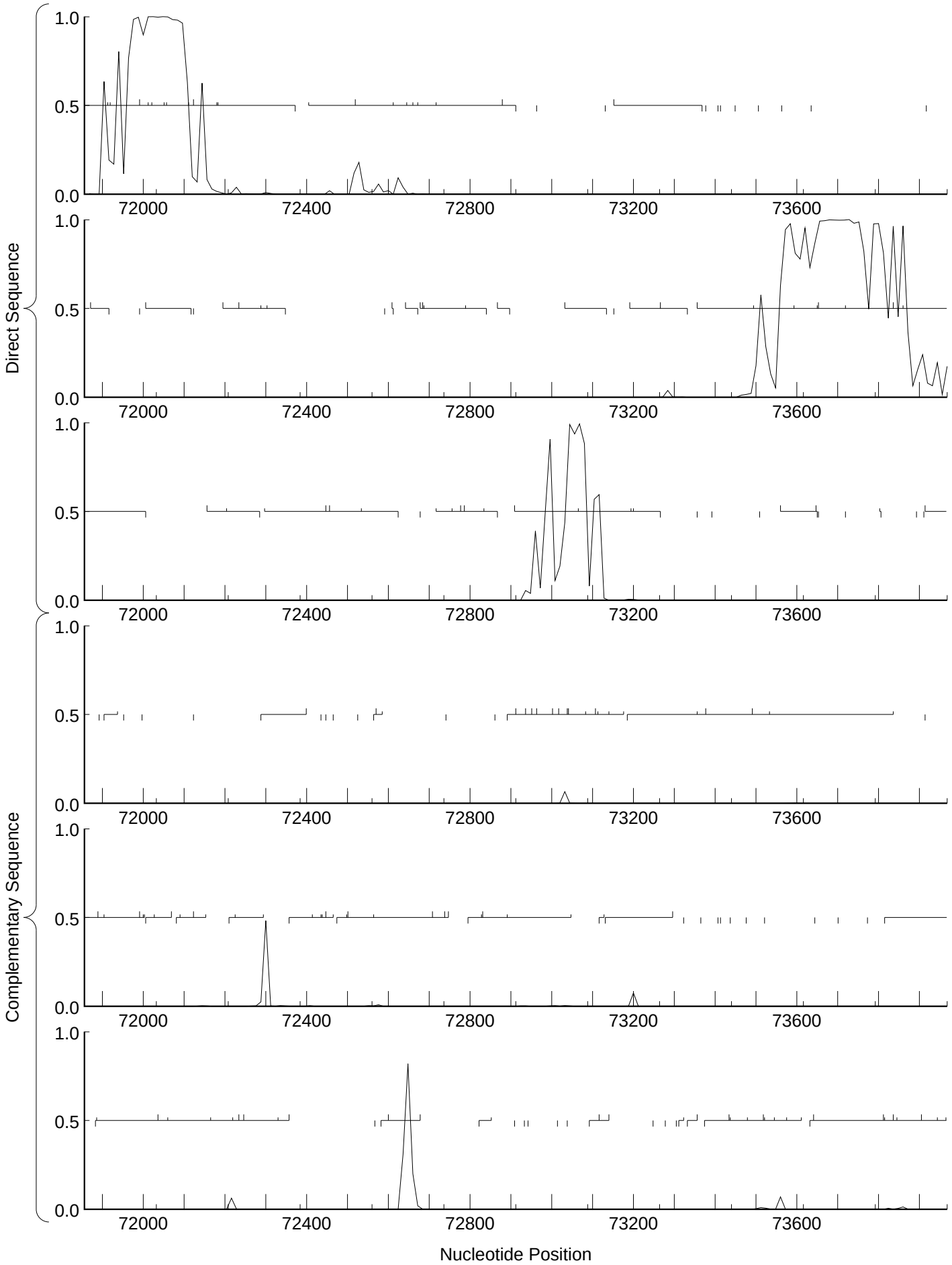
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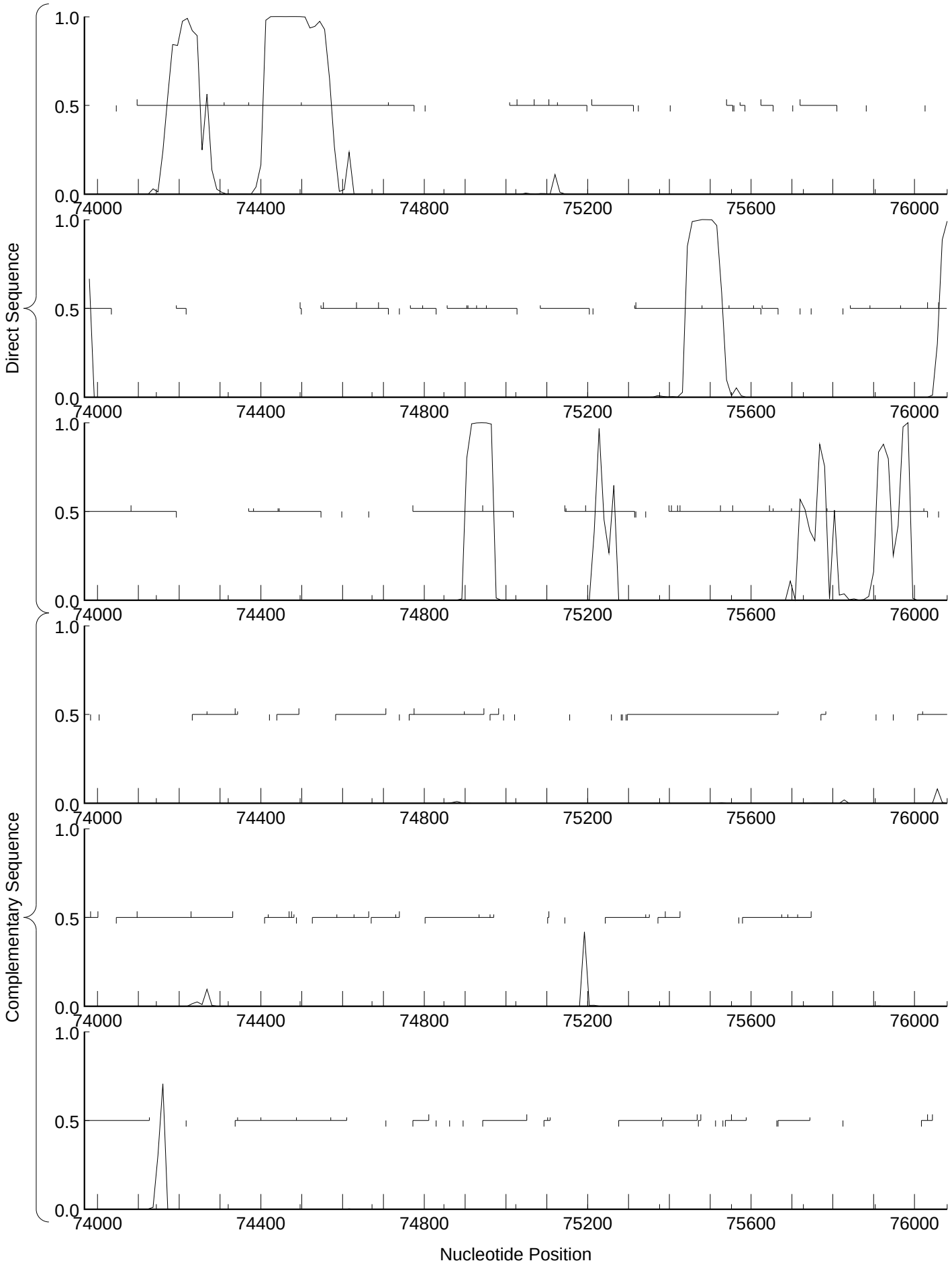
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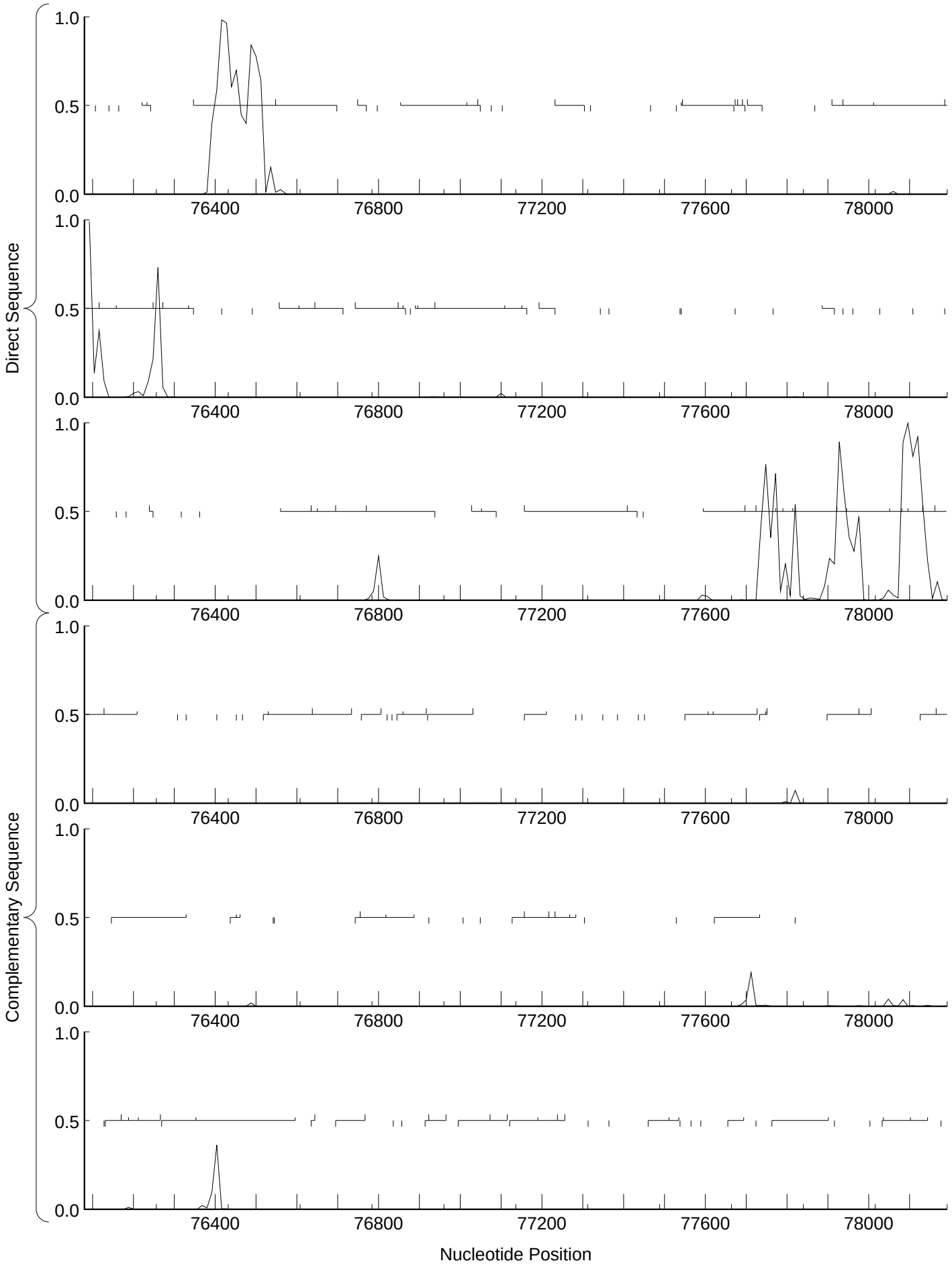
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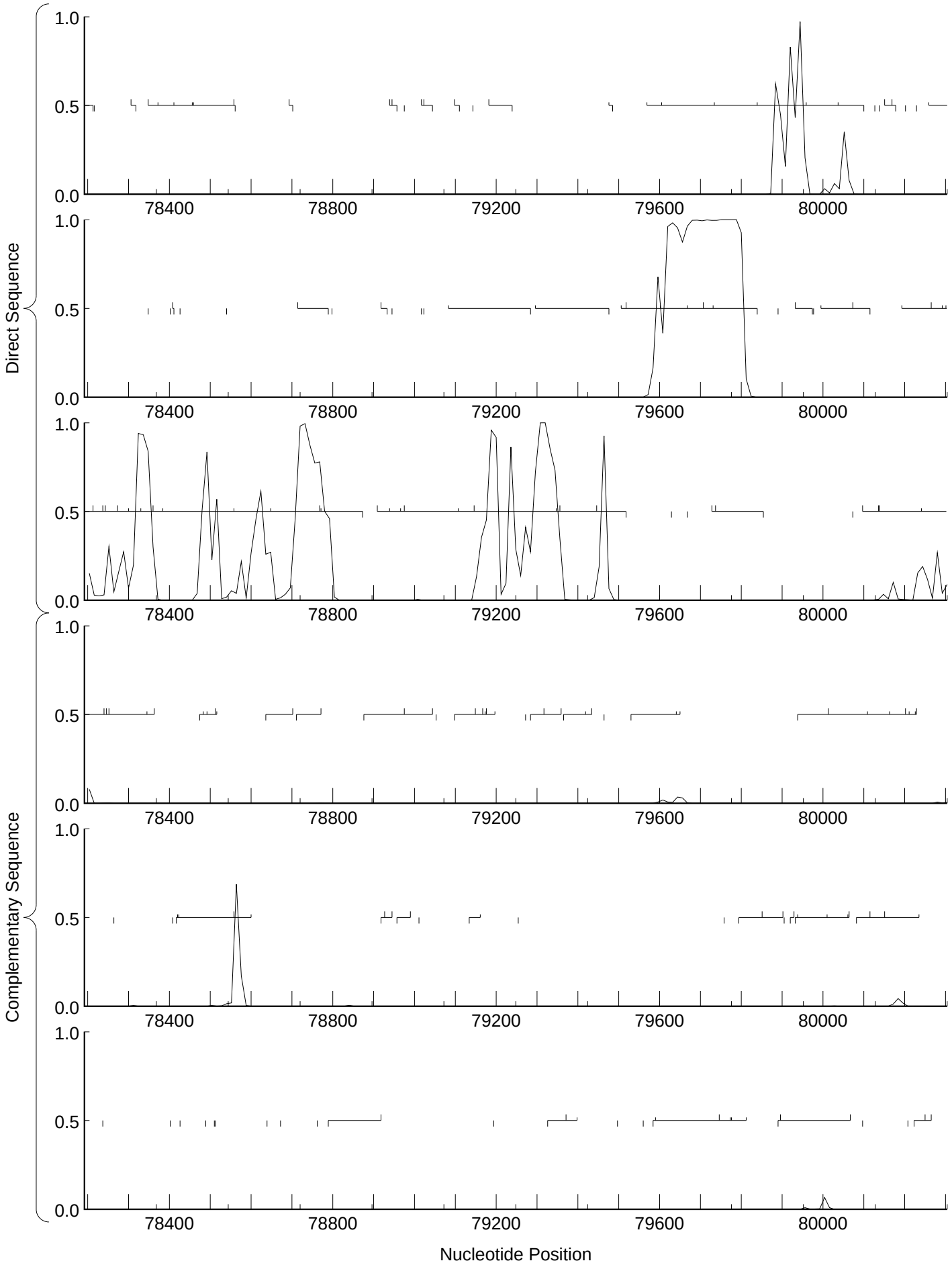
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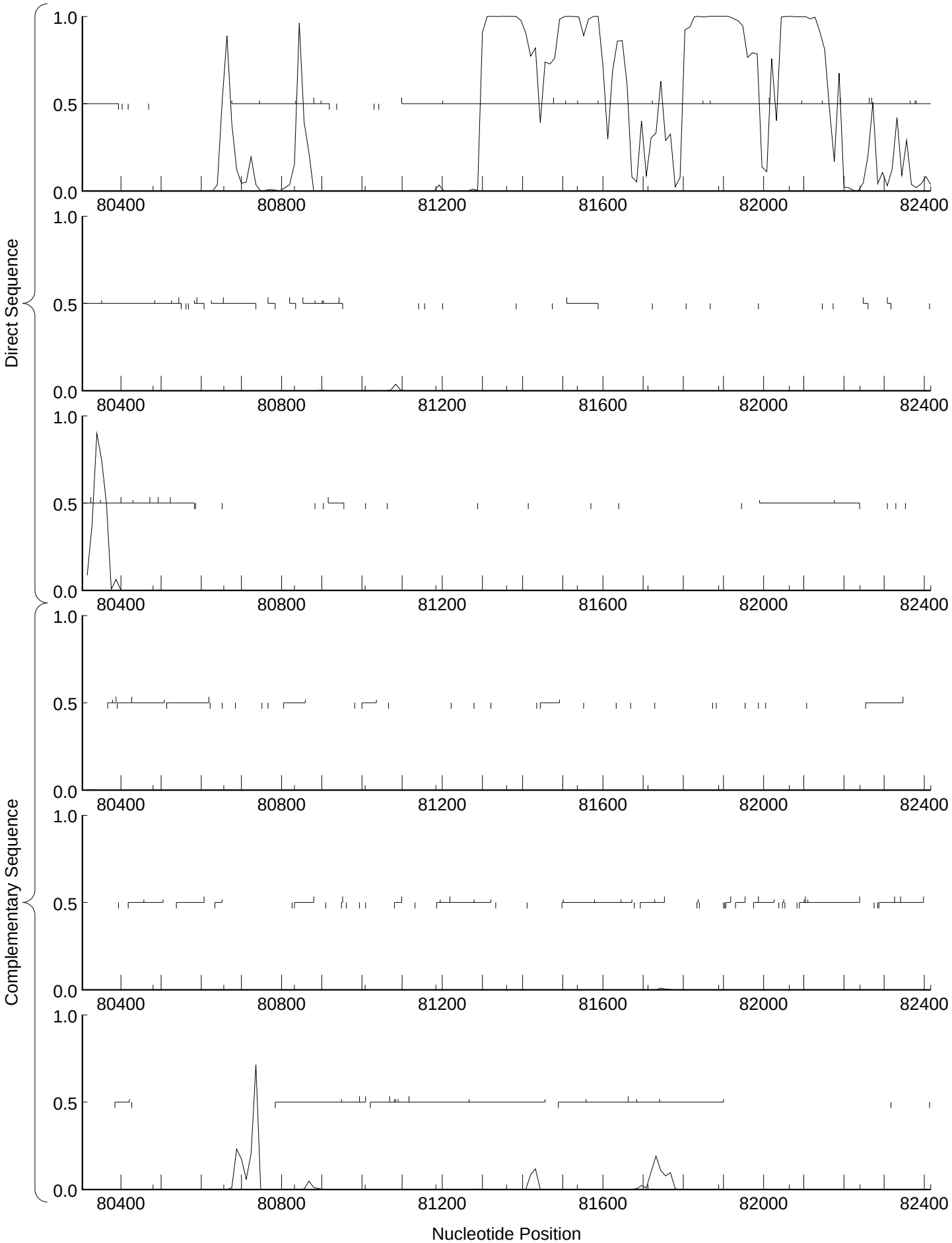
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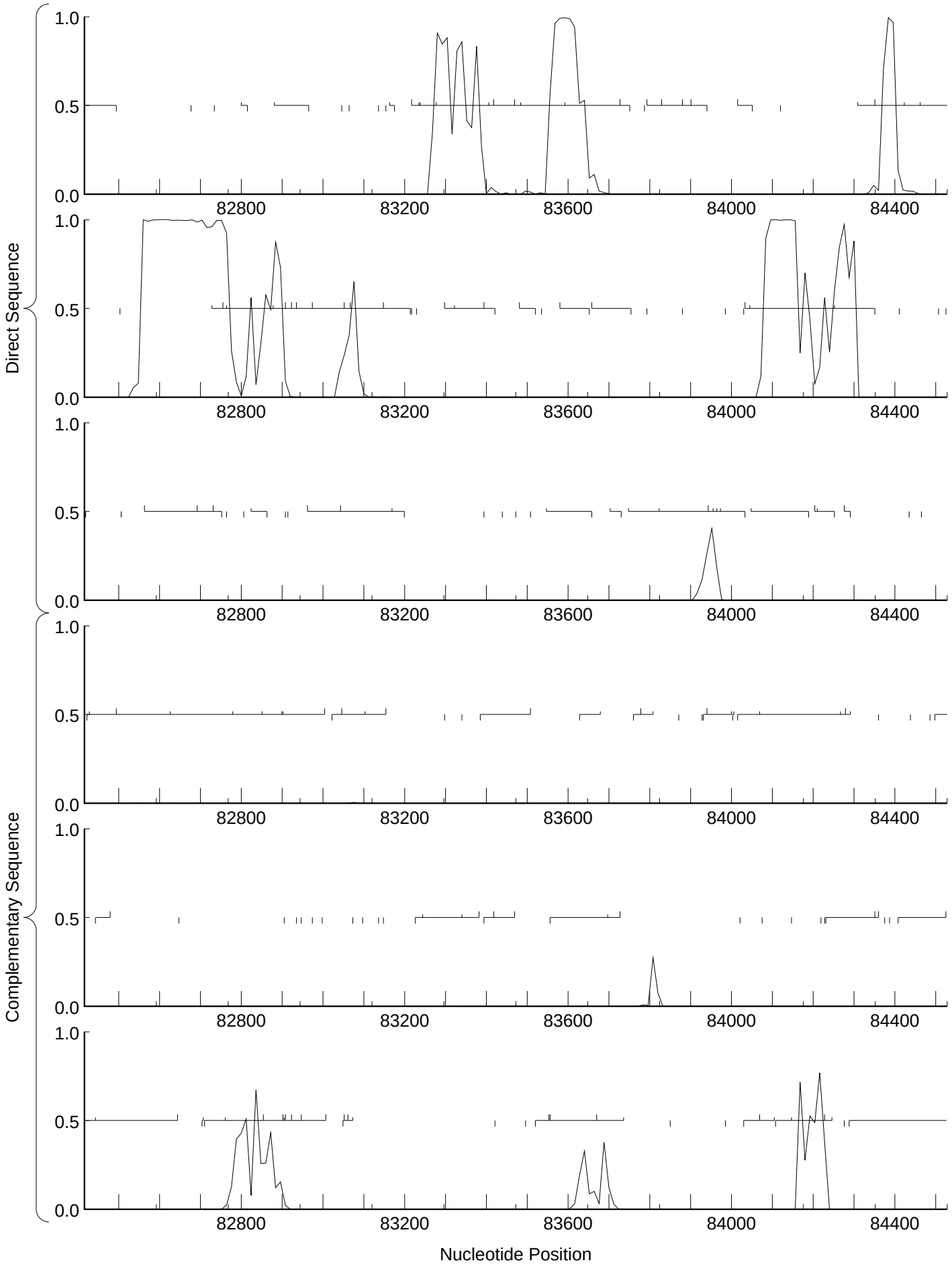


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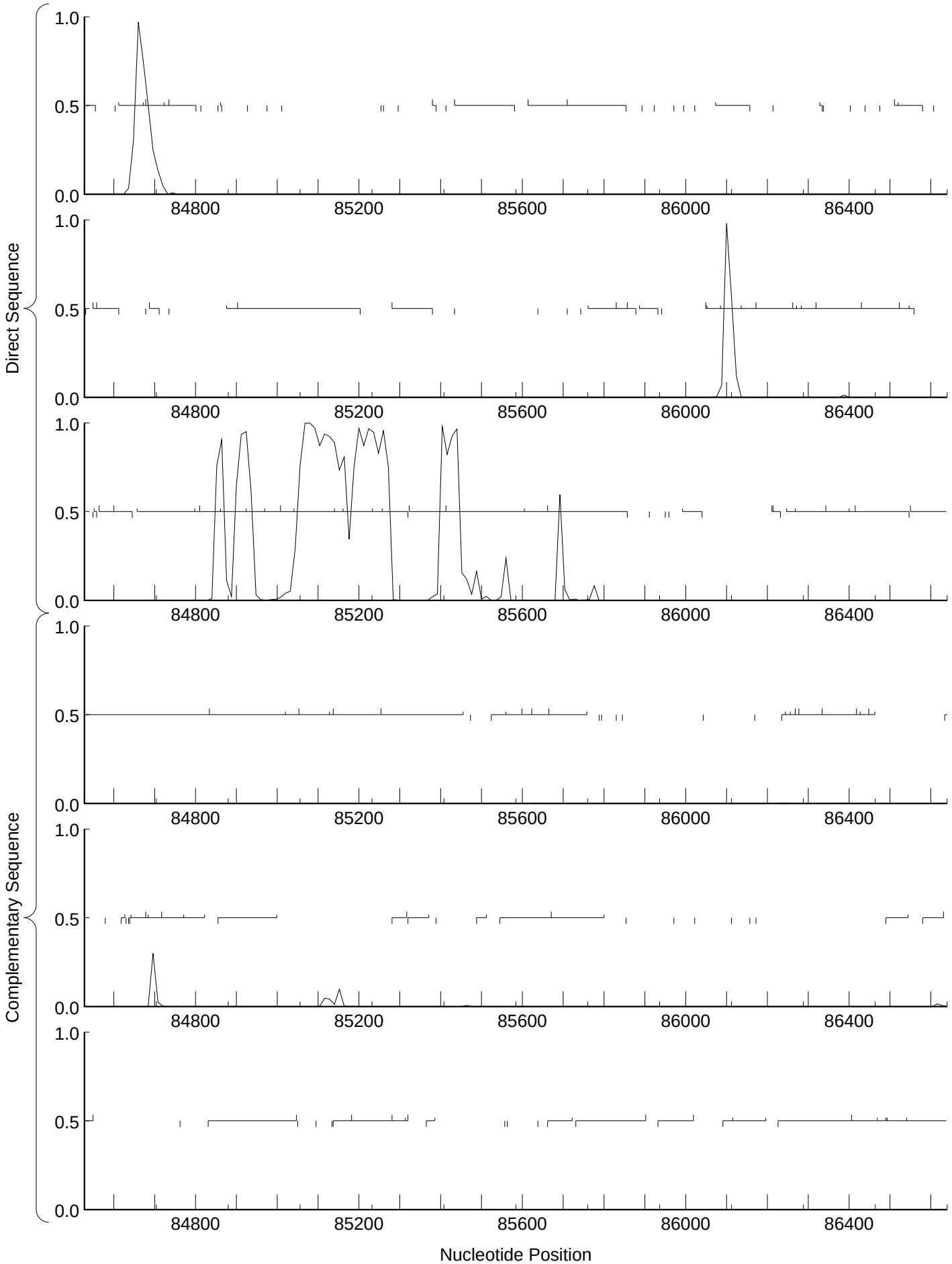


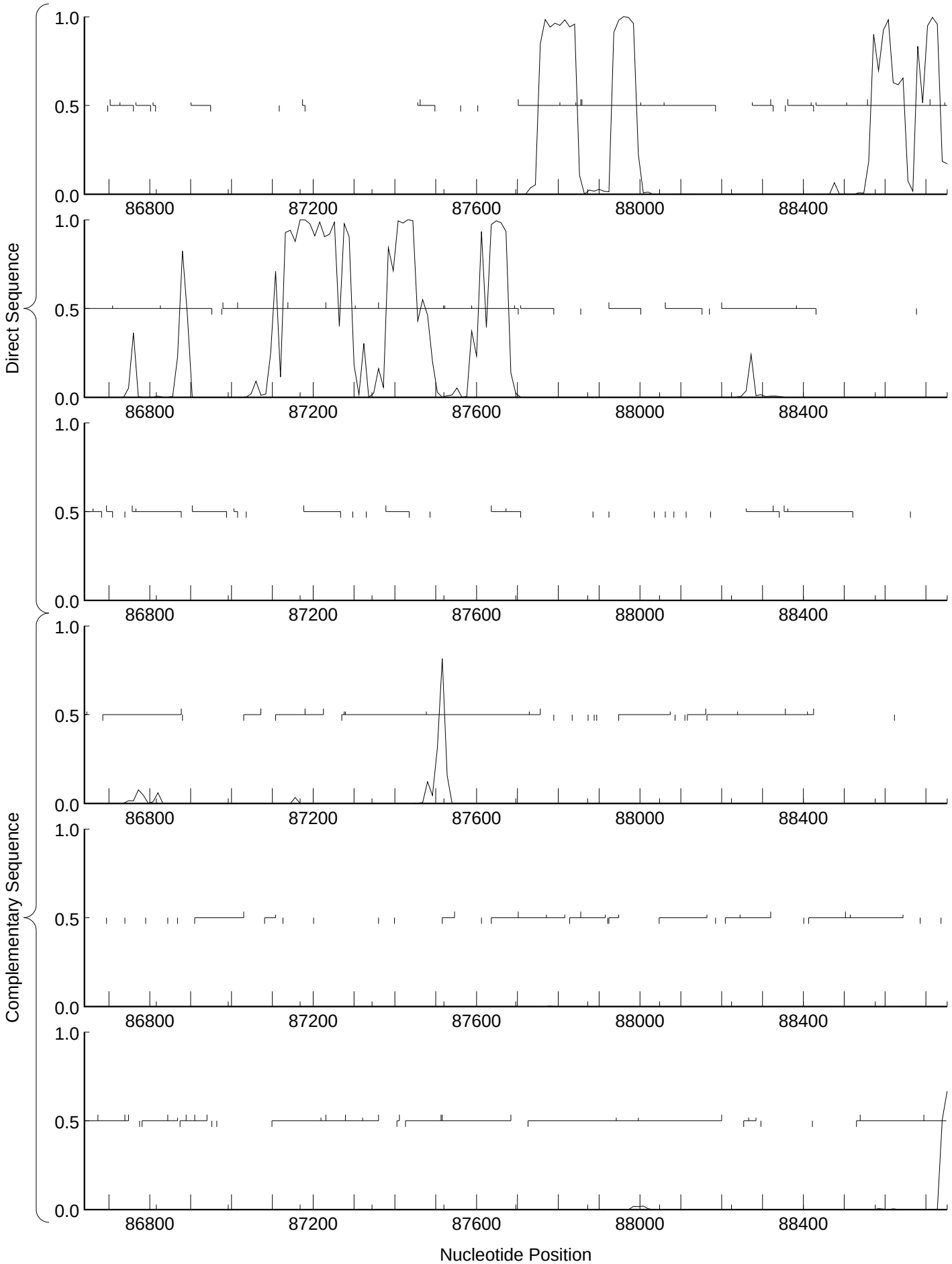
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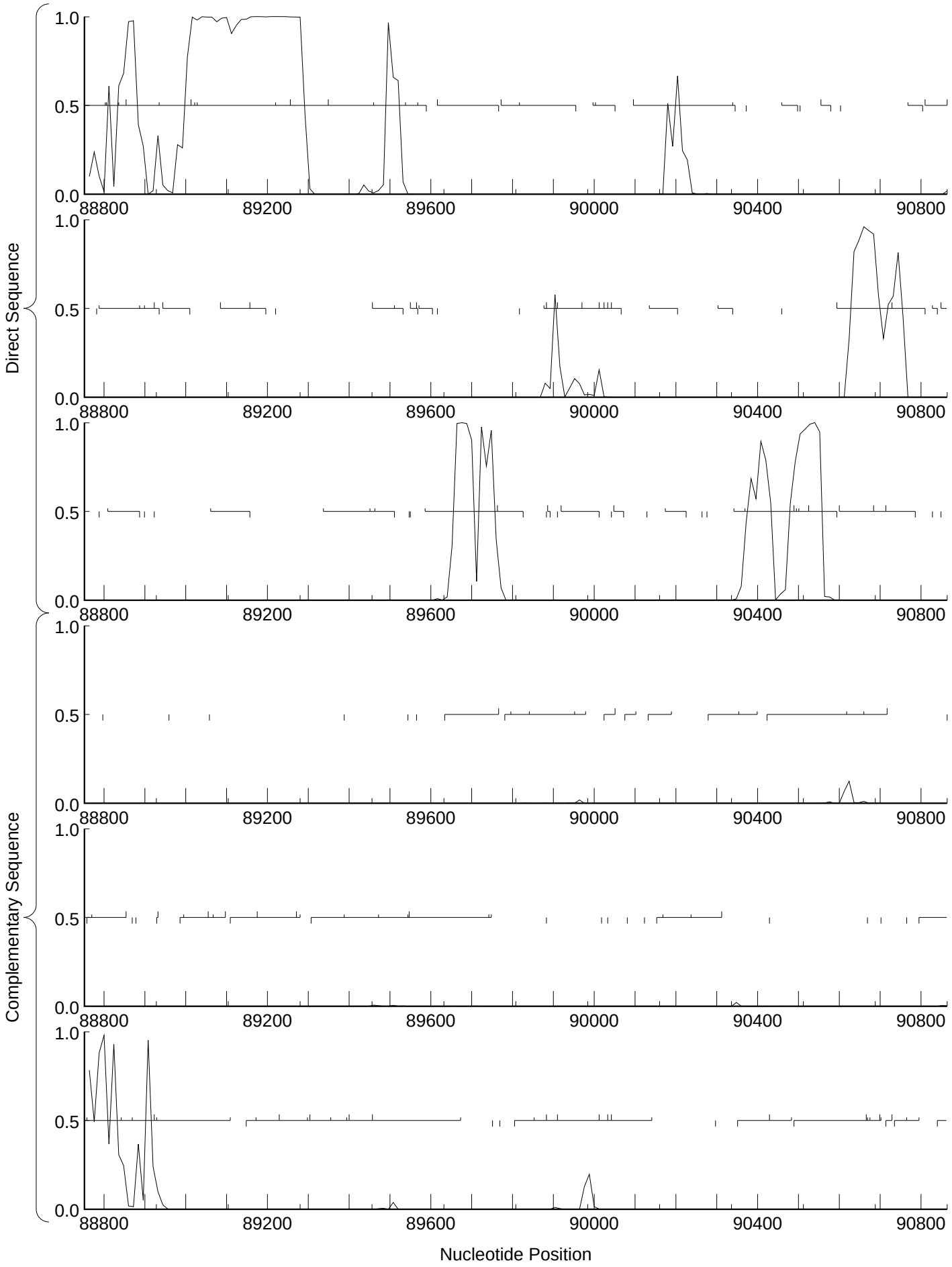


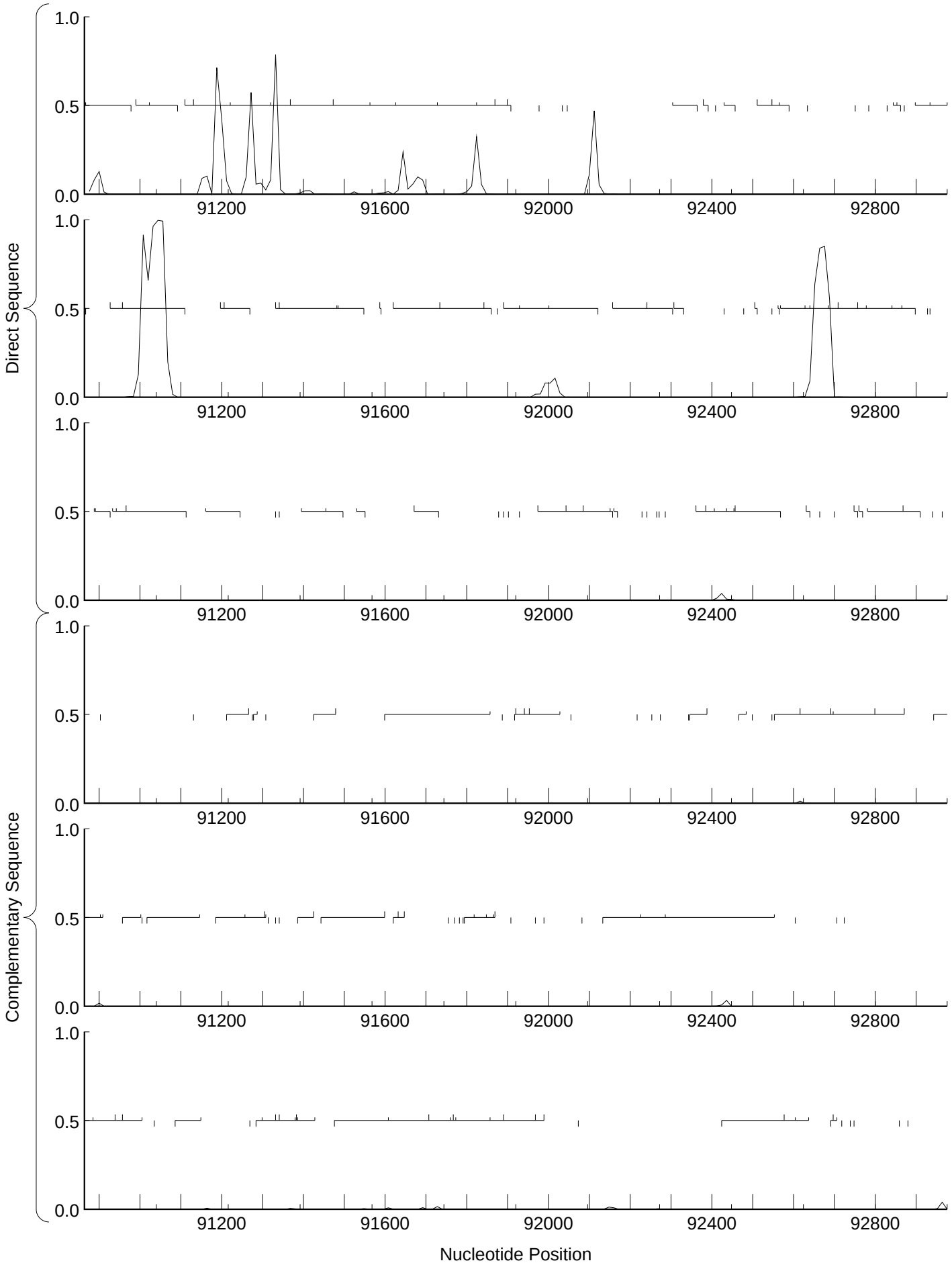

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84400
Nucleotide Position

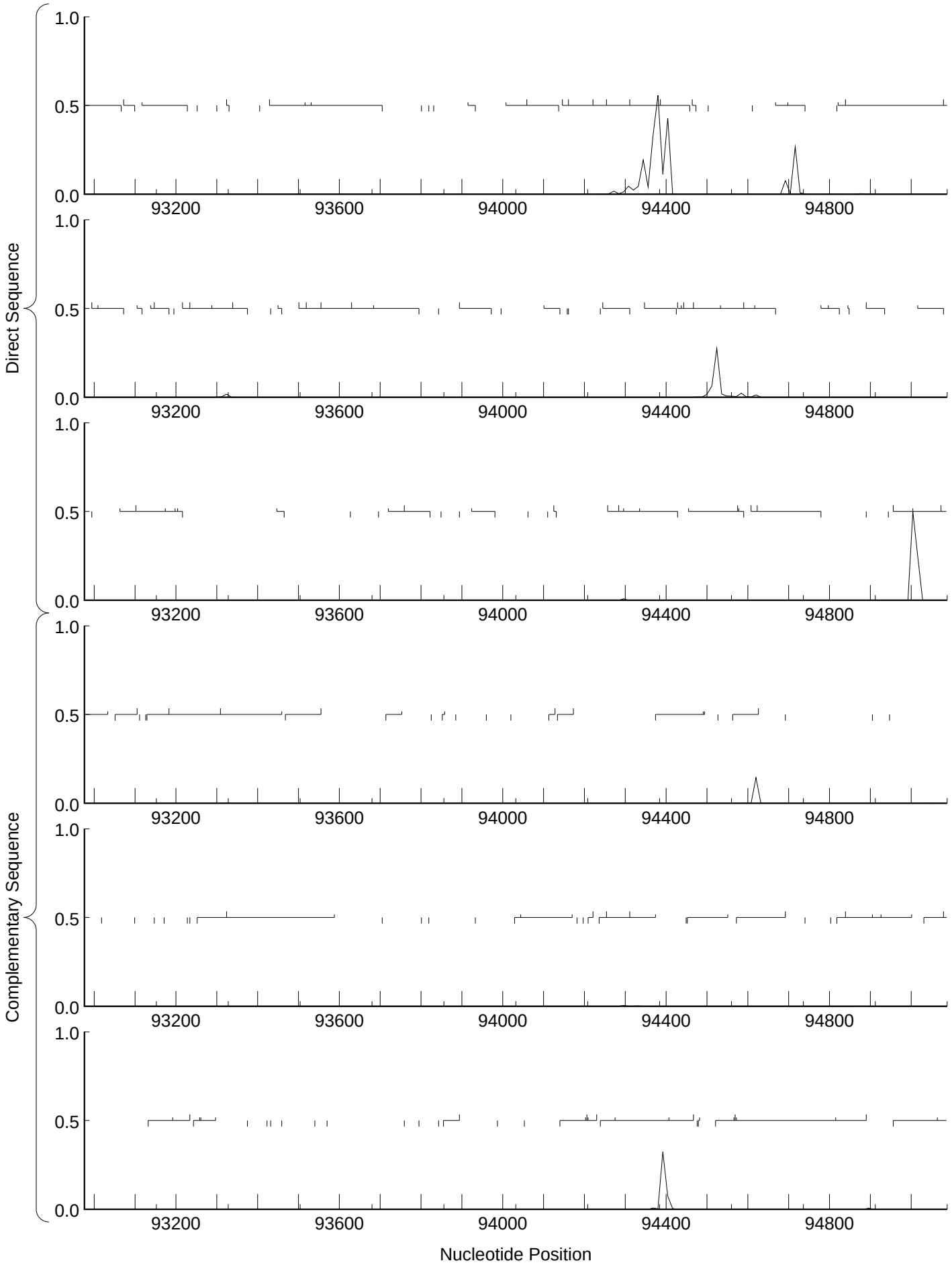
ge Constella complete sequence, 110169 bp including 4-base 3’ overhang (ATCC), Cluster J, Order 4, Window 96, Step 12, 42/54



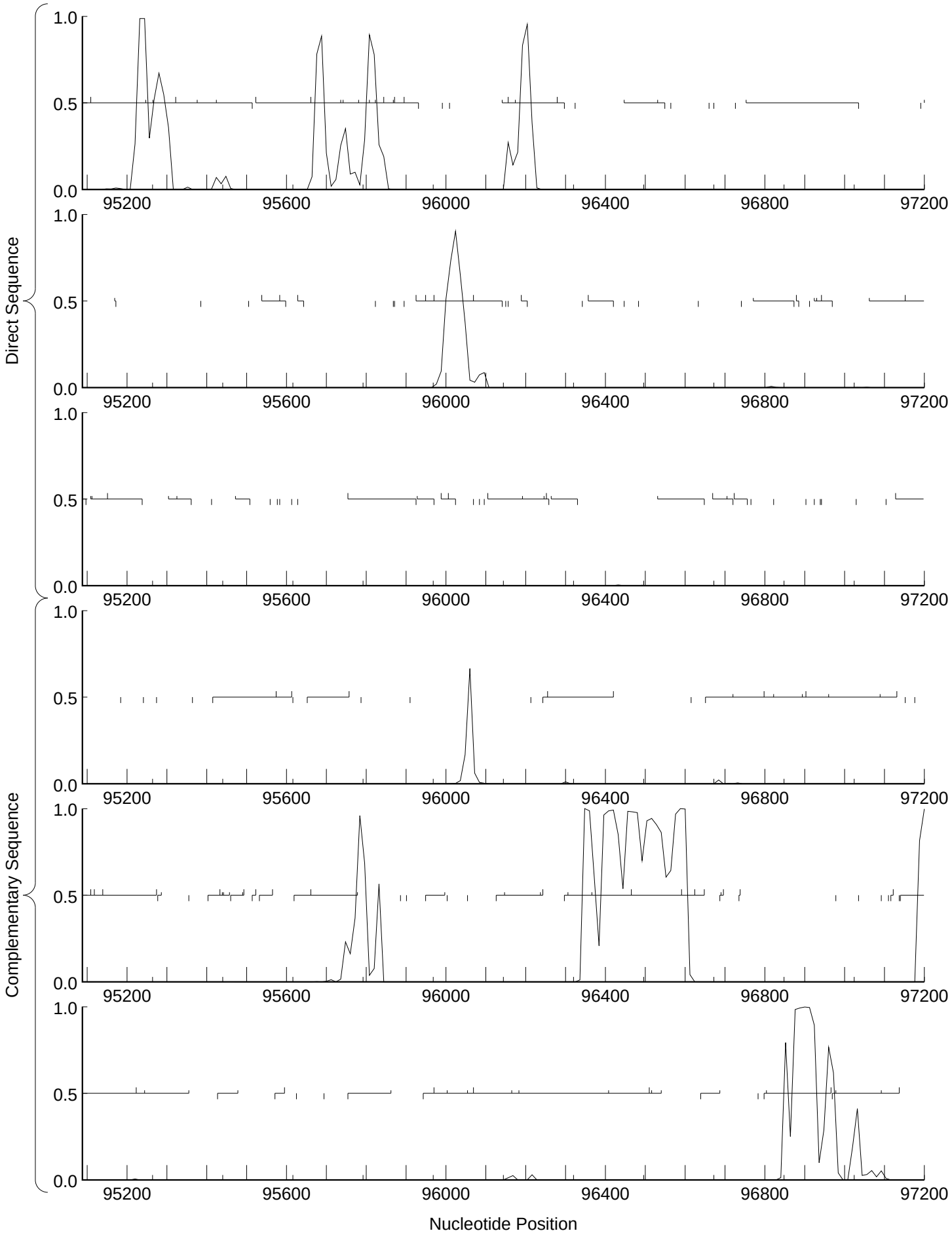

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
86800
87200
87600
88000
88400
Nucleotide Position

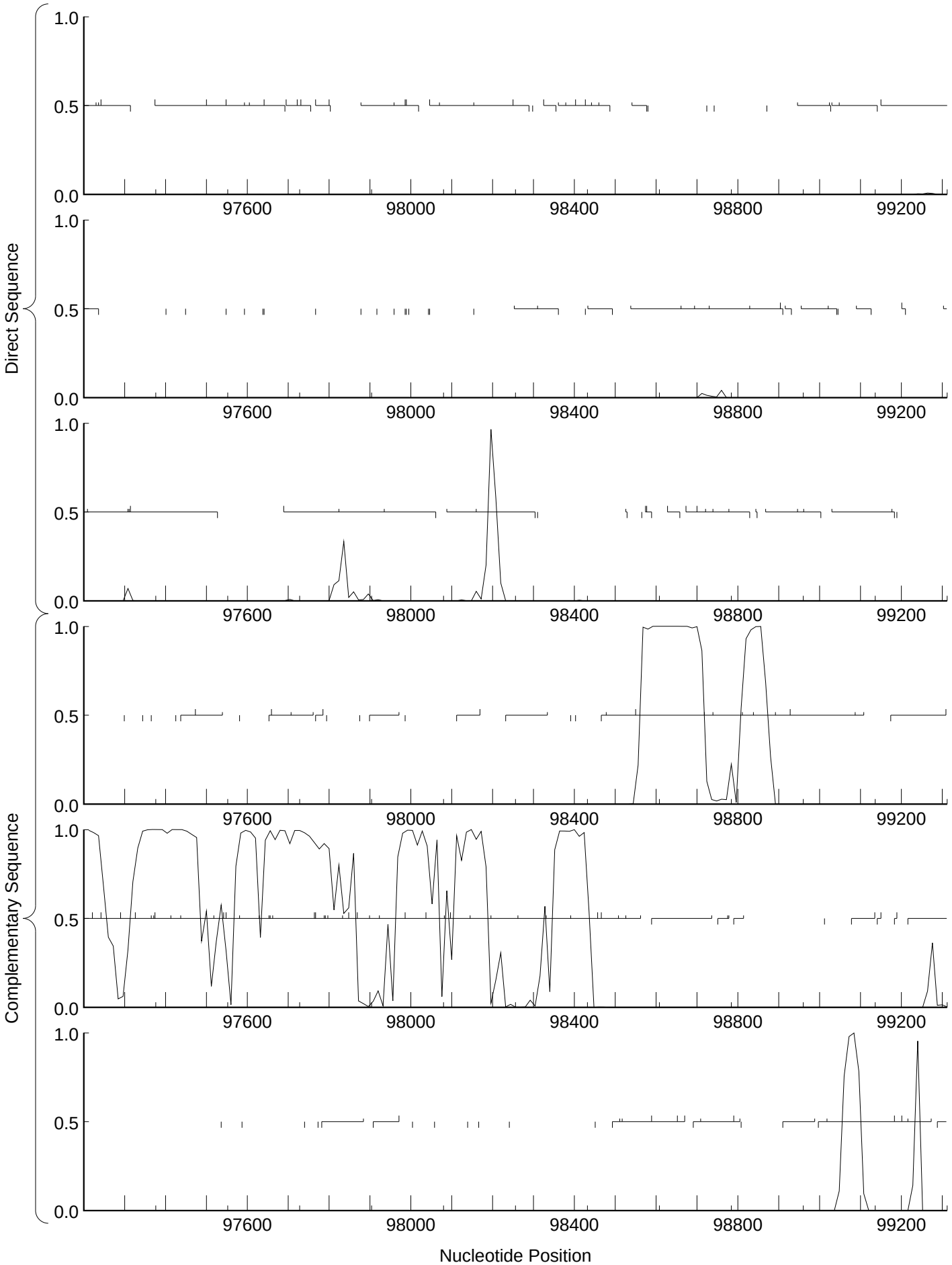

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
88800
89200
89600
90000
90400
90800
Nucleotide Position

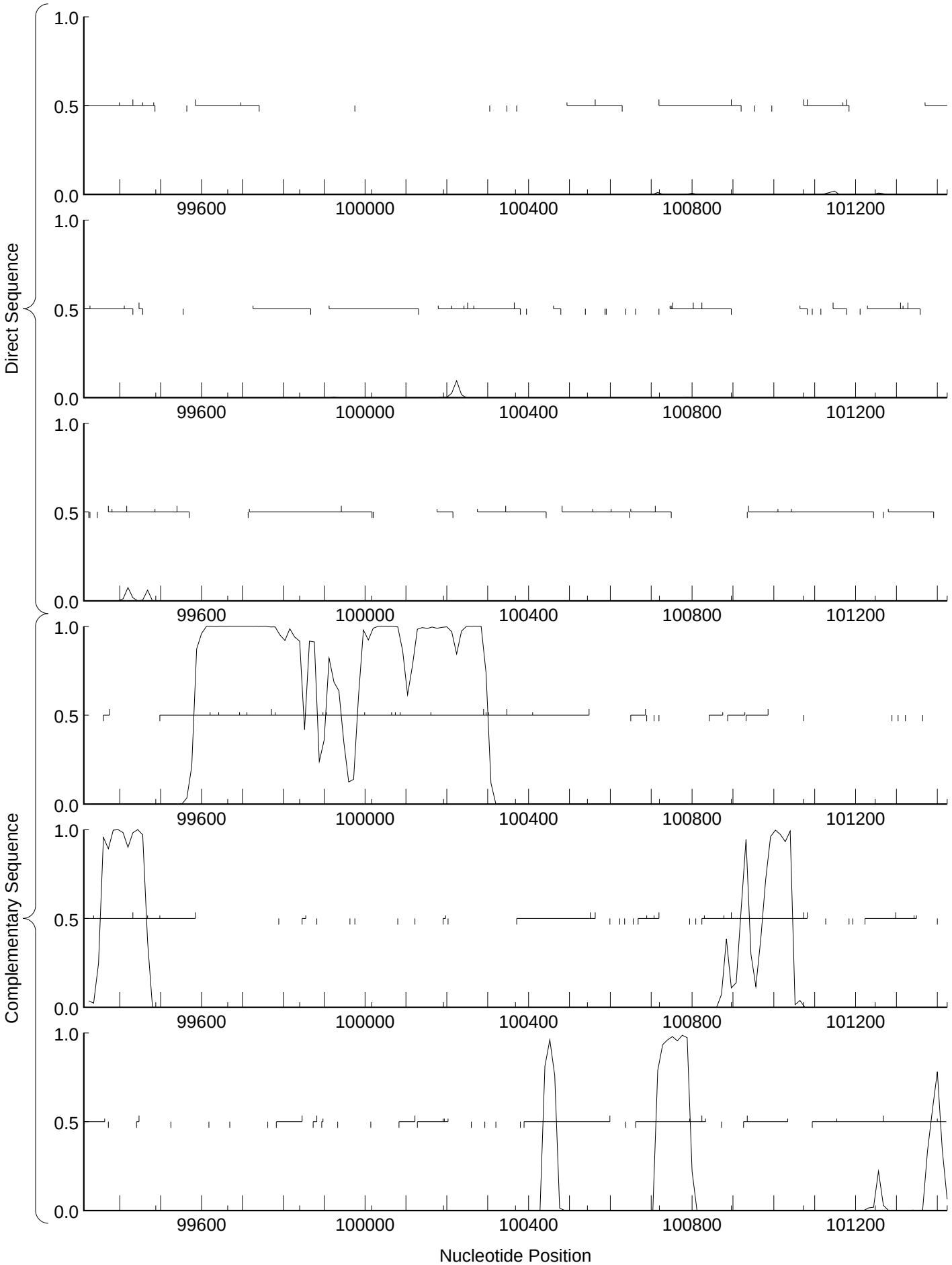

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
91200
91600
92000
92400
92800
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
93200
93600
94000
94400
94800
Nucleotide Position

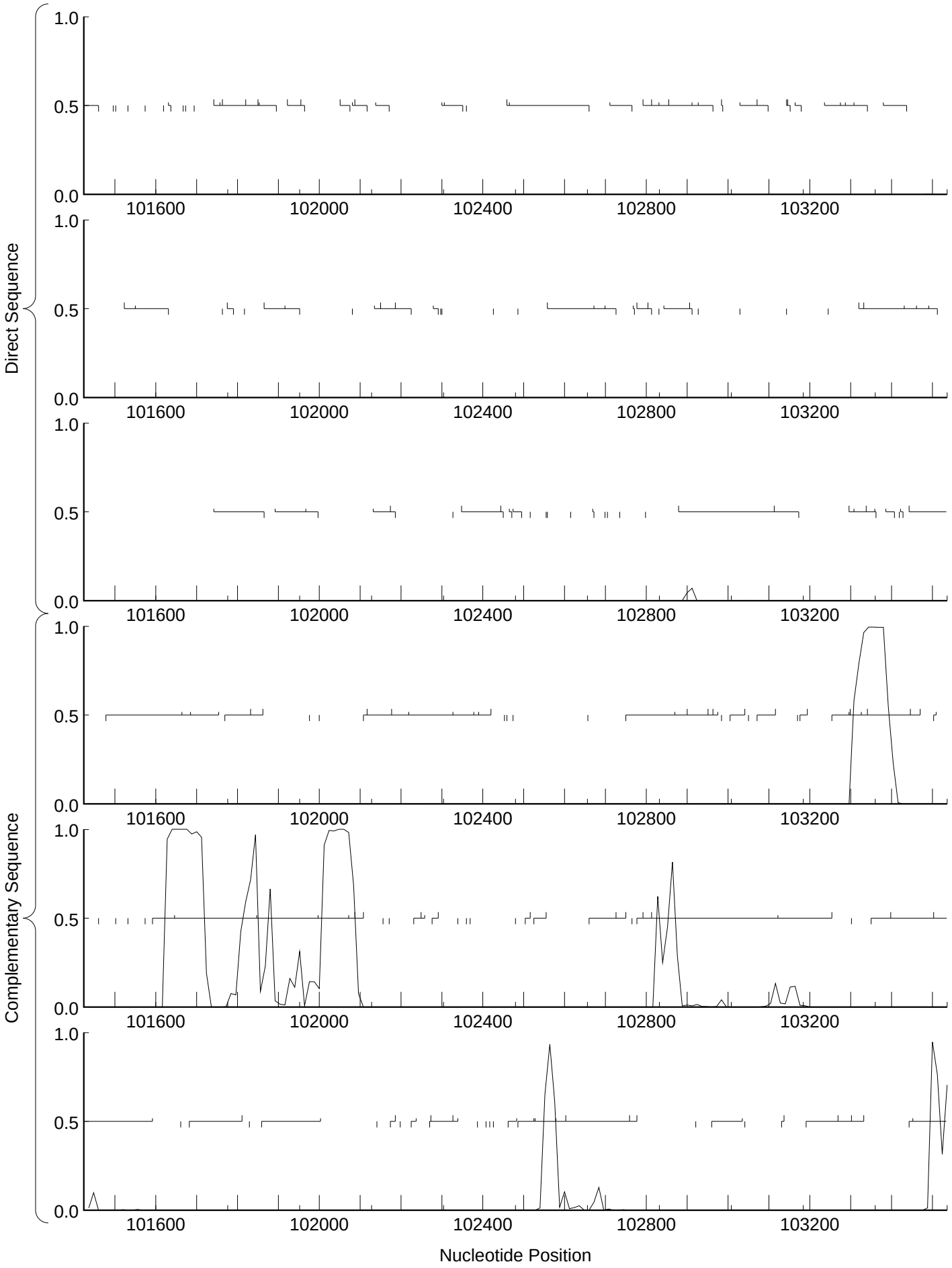
ge Constella complete sequence, 110169 bp including 4-base 3’ overhang (ATCC), Cluster J, Order 4, Window 96, Step 12, 47/54



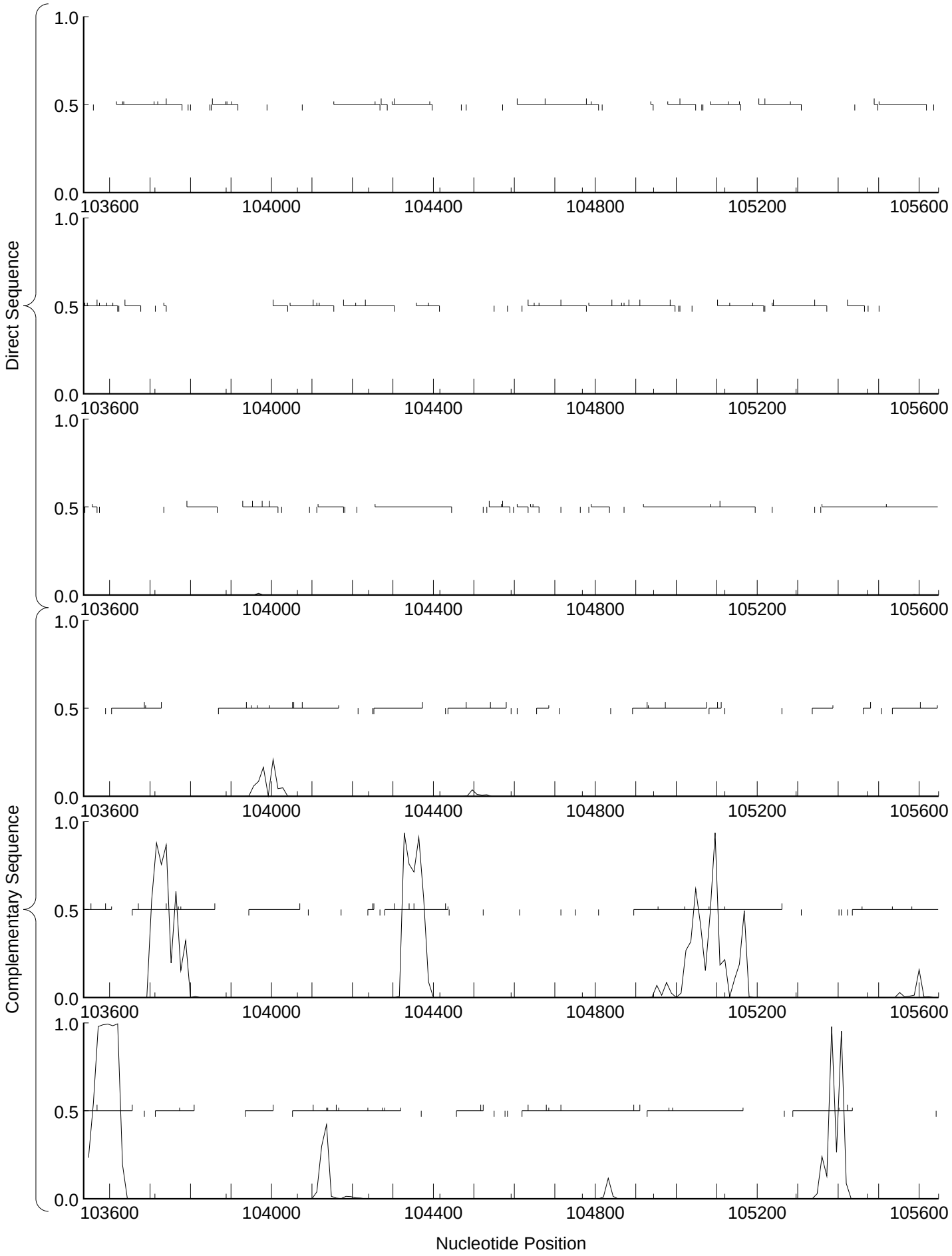

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
97600
98000
98400
98800
99200
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
99600
100000
100400
100800
101200
Nucleotide Position

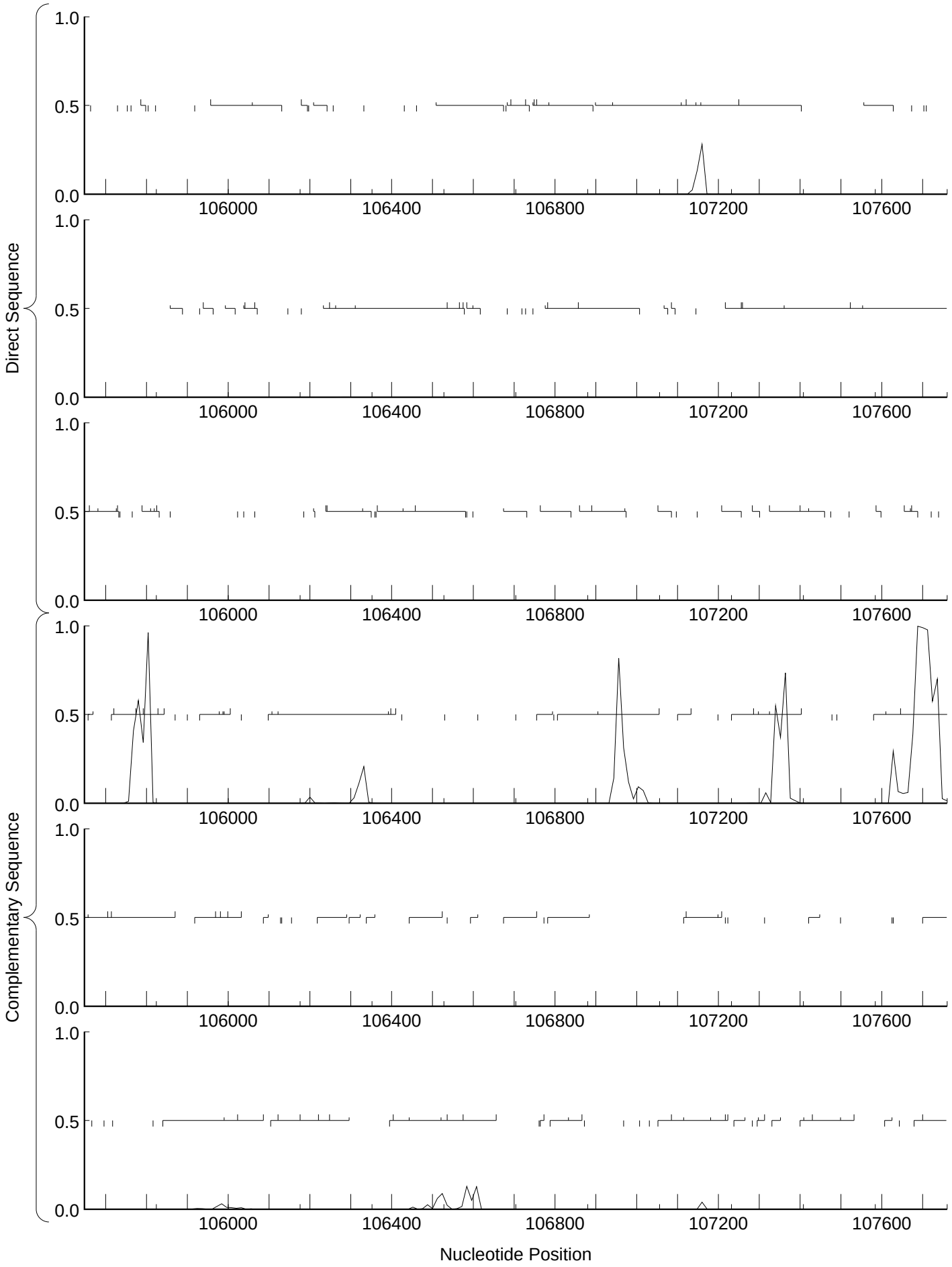
ge Constella complete sequence, 110169 bp including 4-base 3’ overhang (ATCC), Cluster J, Order 4, Window 96, Step 12, 50/54

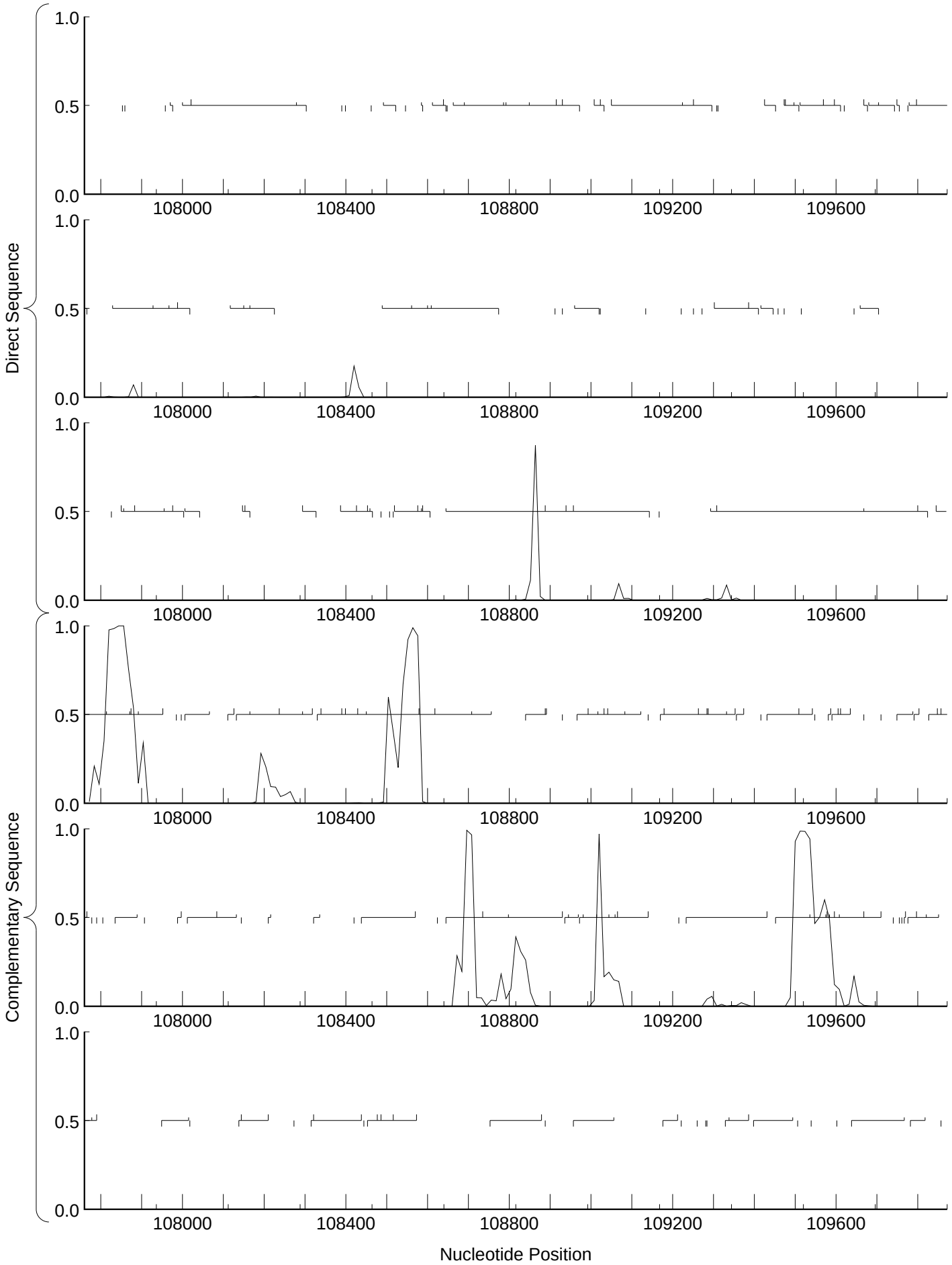


ge Constella complete sequence, 110169 bp including 4-base 3’ overhang (ATCC), Cluster J, Order 4, Window 96, Step 12, 51/54



ge Constella complete sequence, 110169 bp including 4-base 3’ overhang (ATCC), Cluster J, Order 4, Window 96, Step 12, 52/54




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
108000
108400
108800
109200
109600
Nucleotide Position

ge Constella complete sequence, 110169 bp including 4-base 3’ overhang (ATCC), Cluster J, Order 4, Window 96, Step 12, 54/54

