GeneMark

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

PROGRAM INFORMATION

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Sequence         :  Mycobacterium phage Silverleaf complete sequence, 73210 bp including 10-base 3' overhang (T
Analysis Date    :  3/29/24 at 10:34:19
Pages            :  36
Sequence Length  :  73210 bp
GC Content       :   58.78%

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

PS-Version       :  1.2

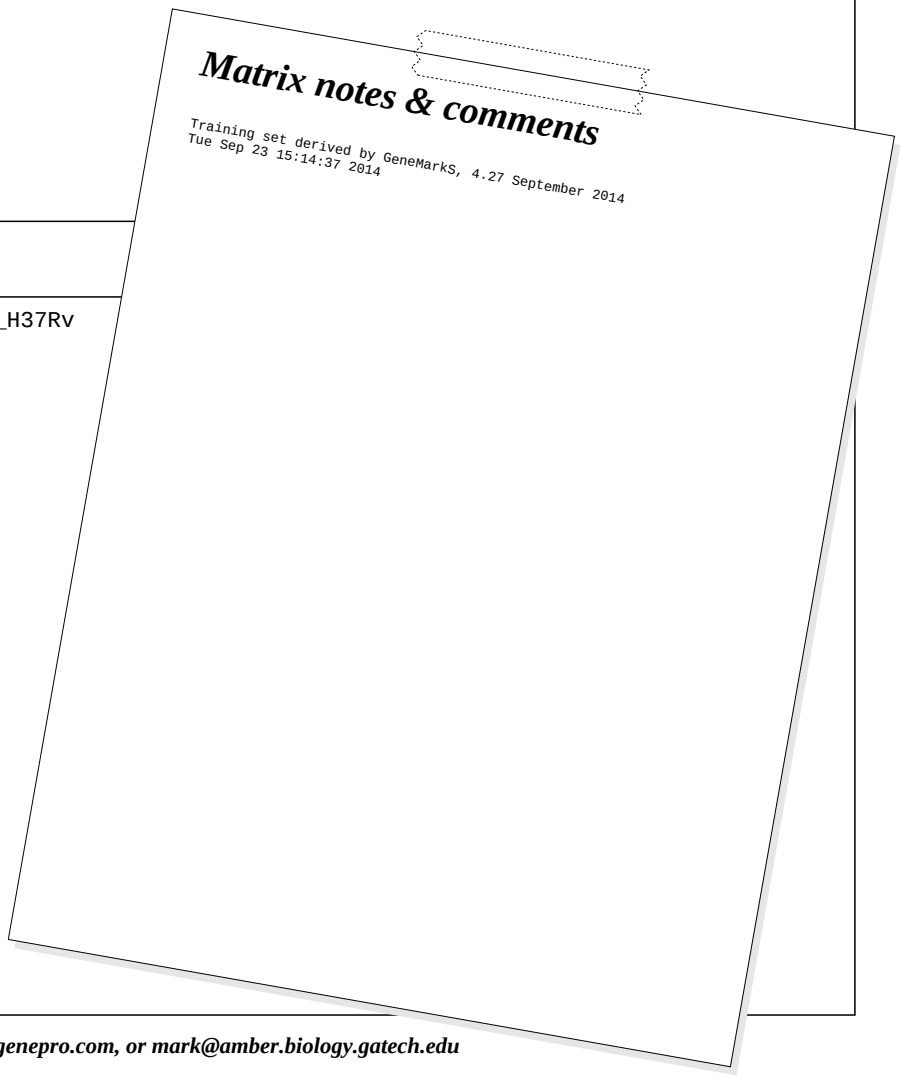
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Author     :  -

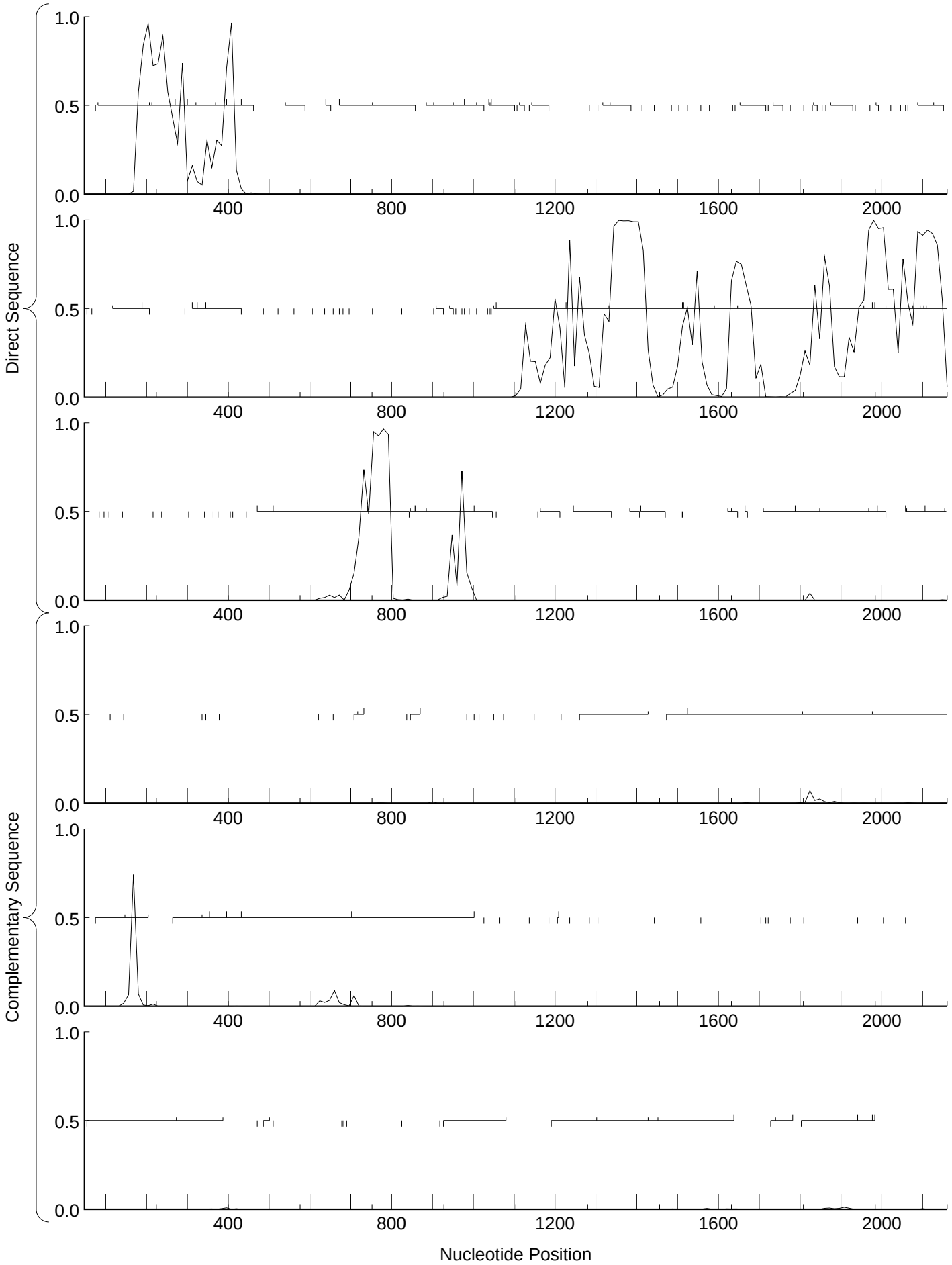
Order      :  4


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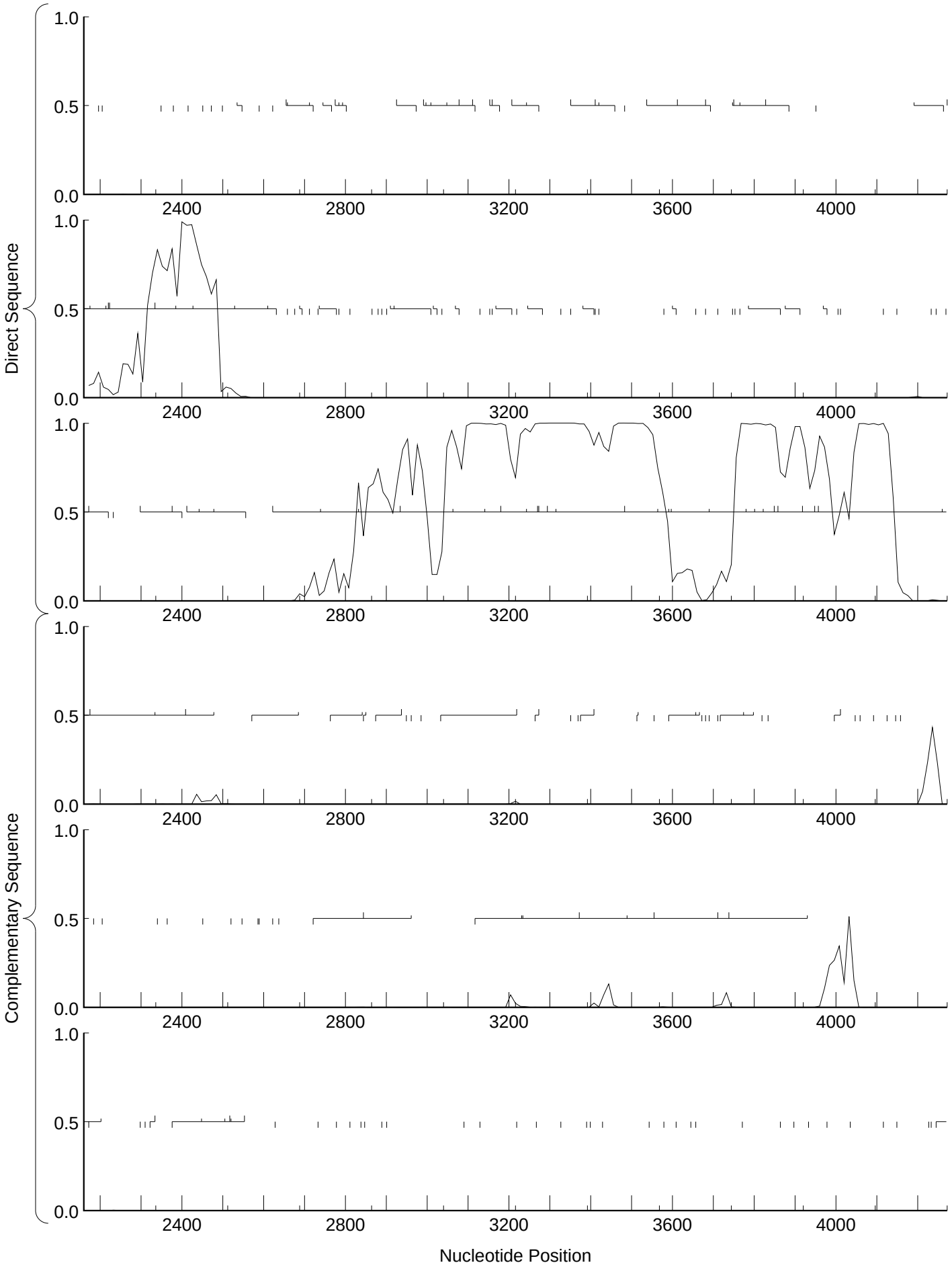


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

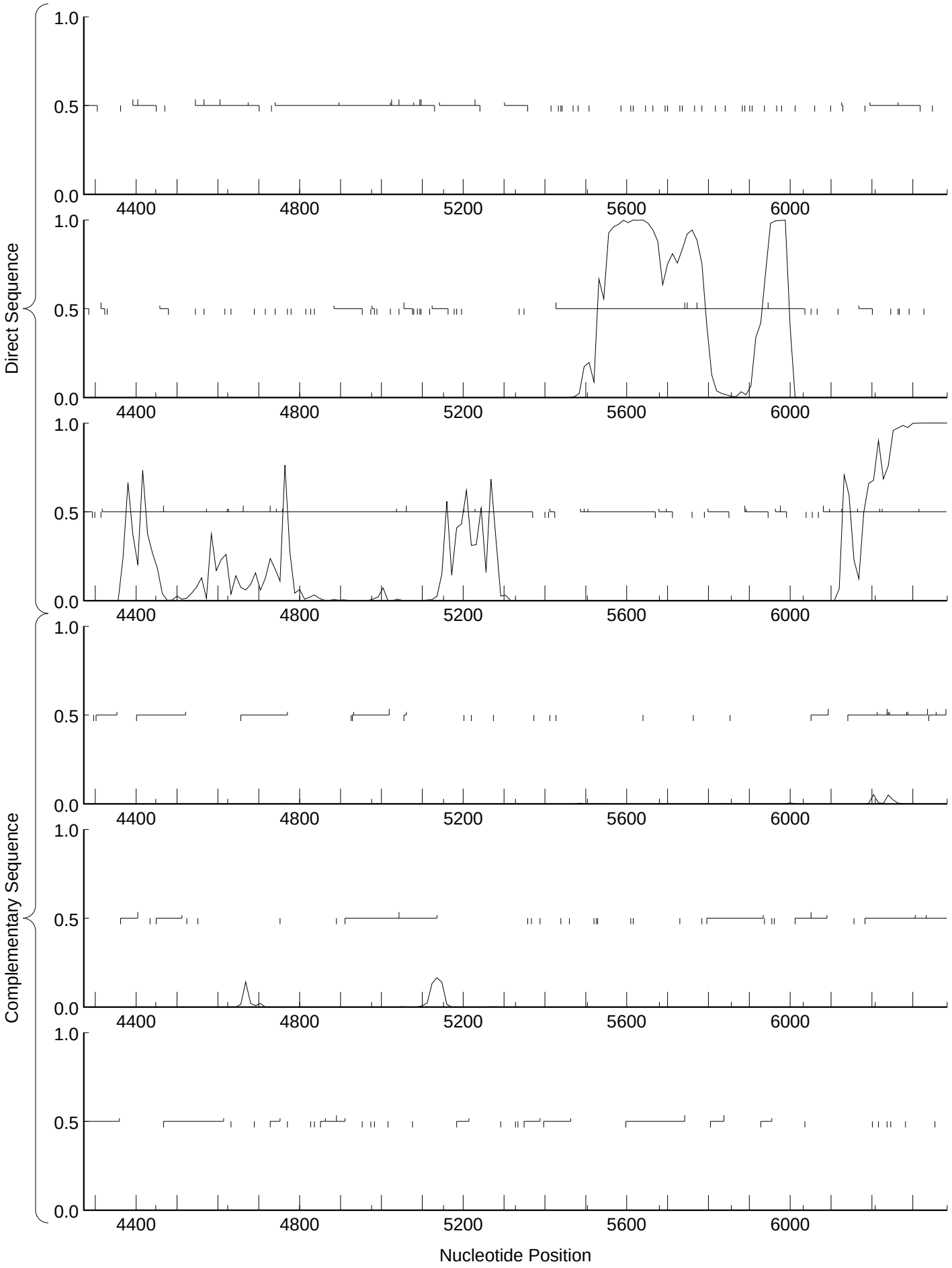
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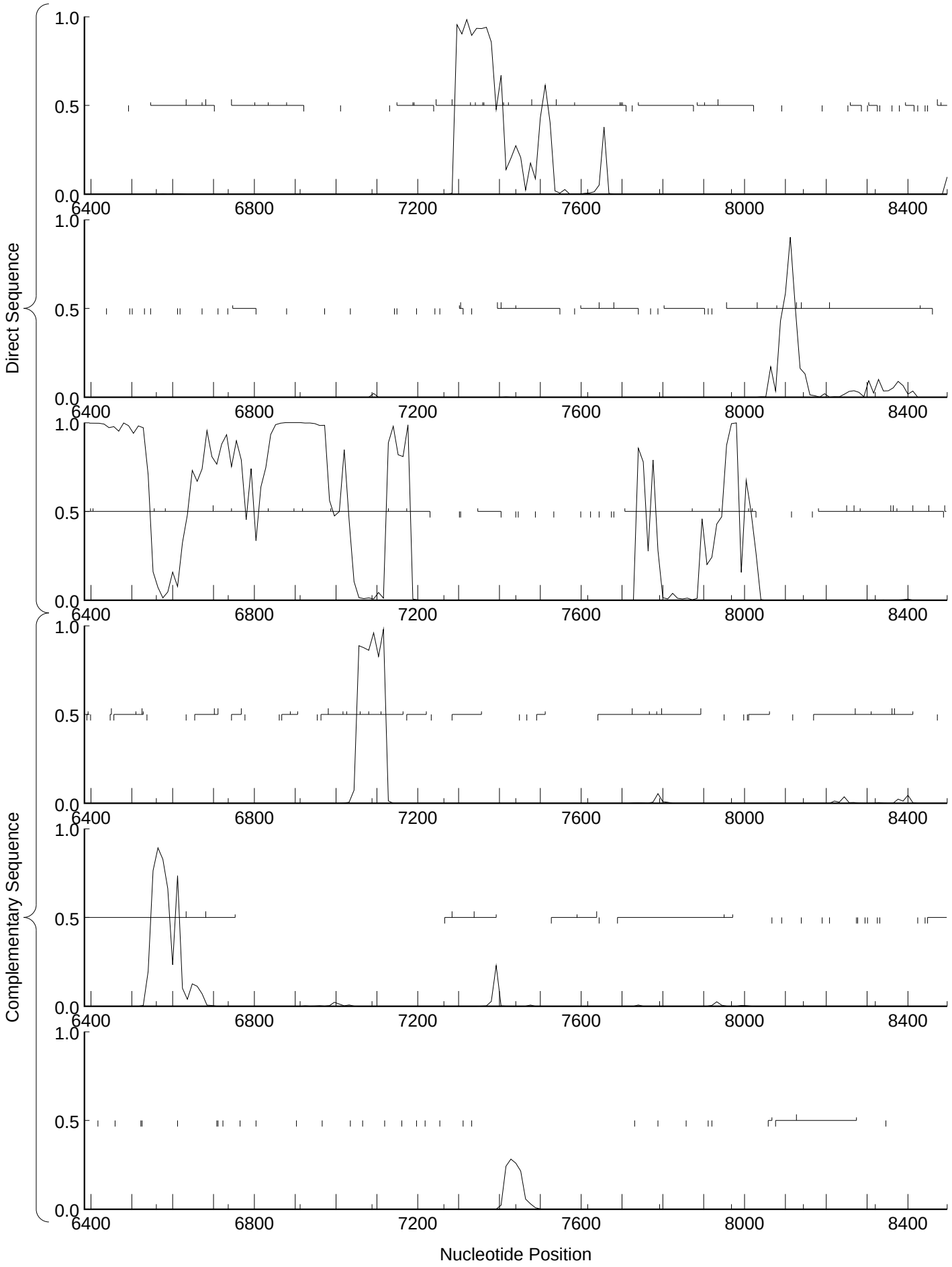
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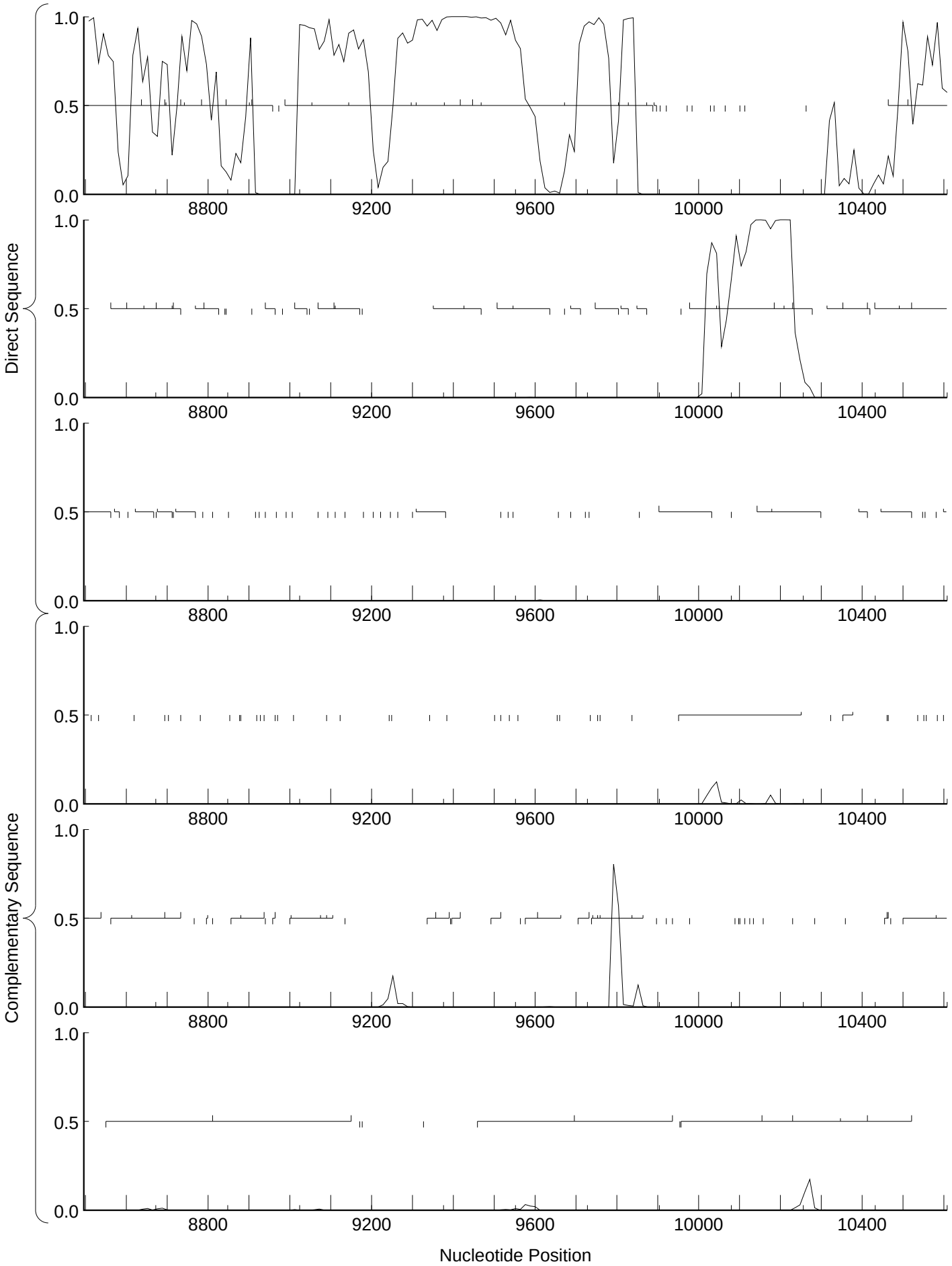
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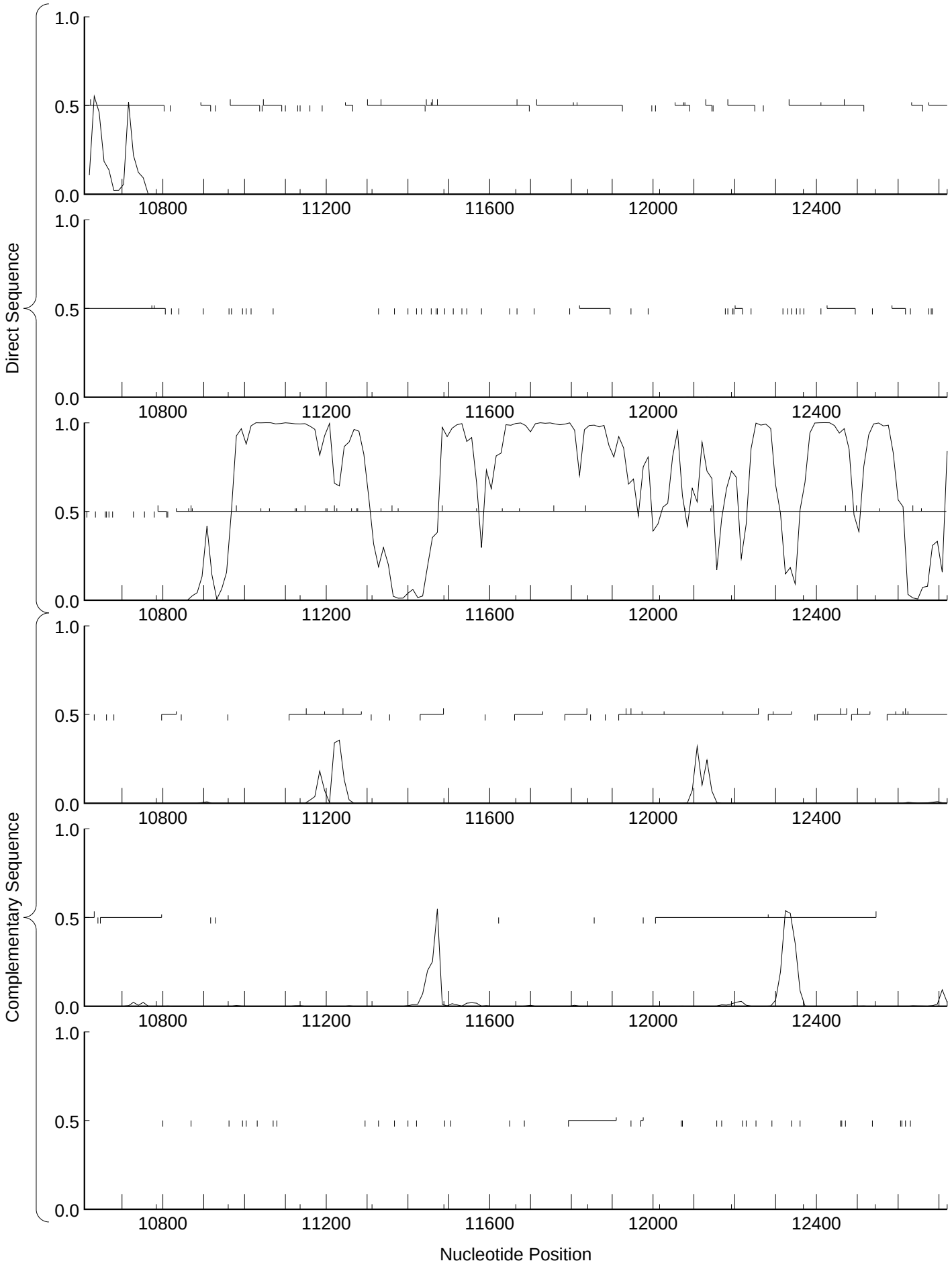
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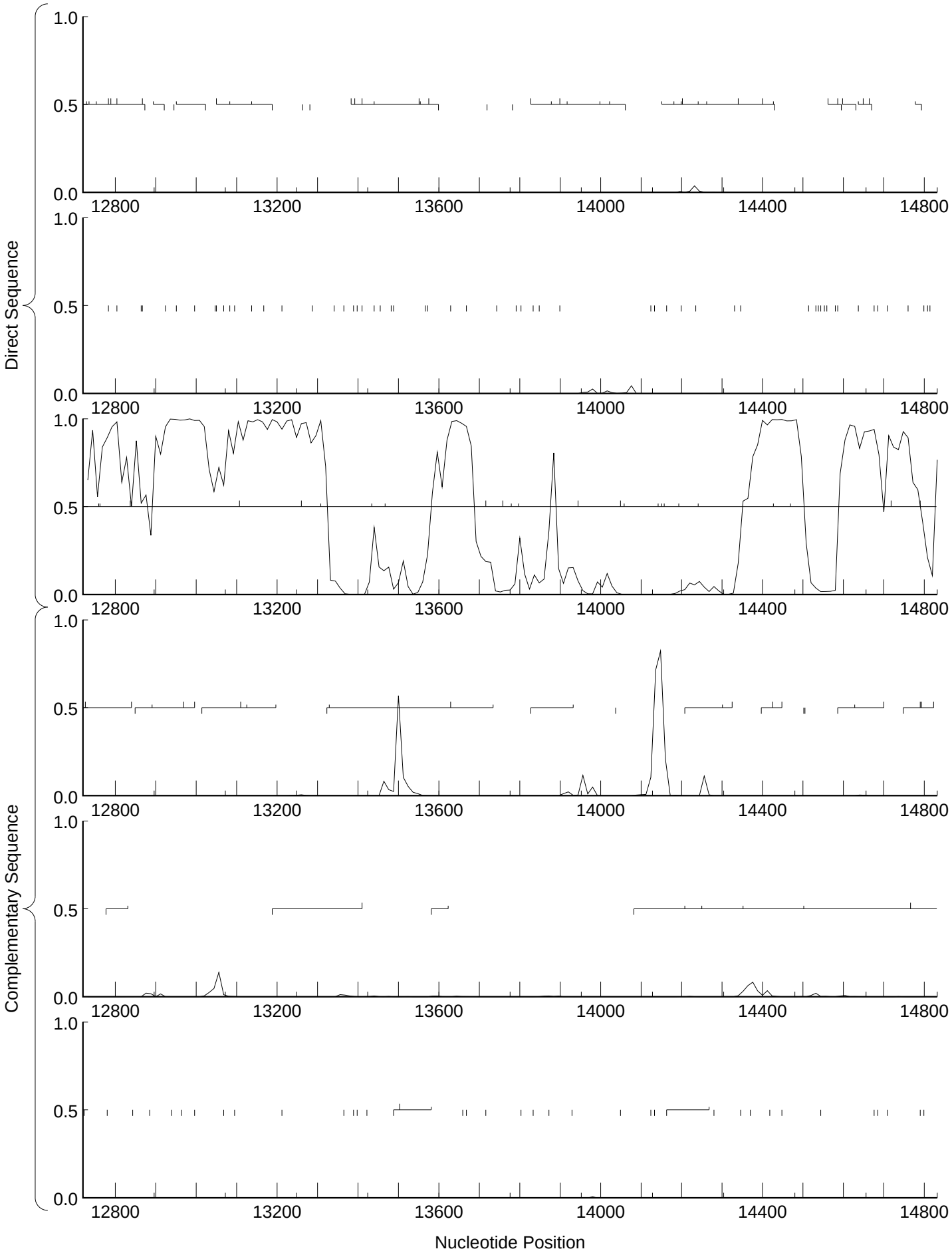
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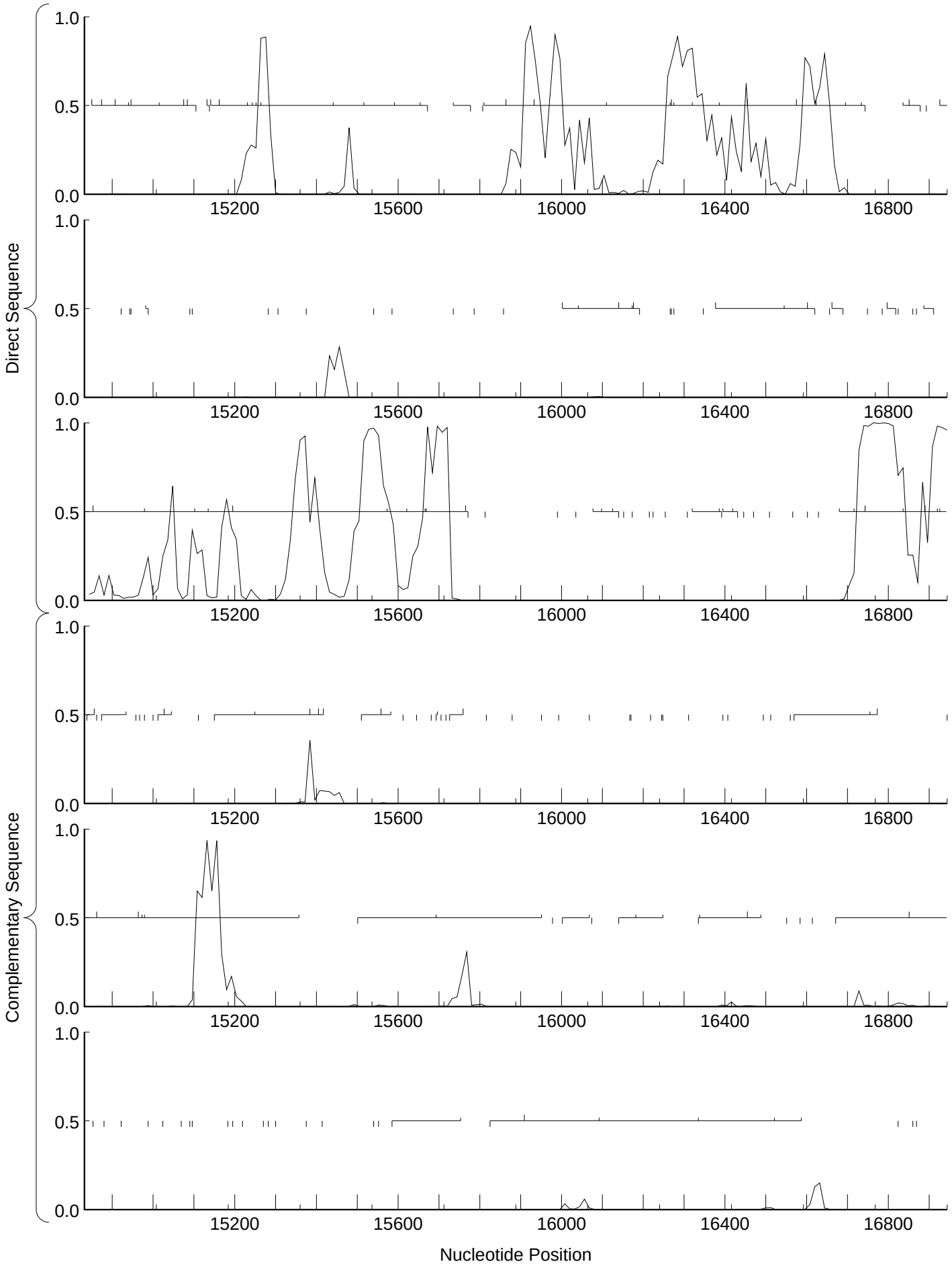
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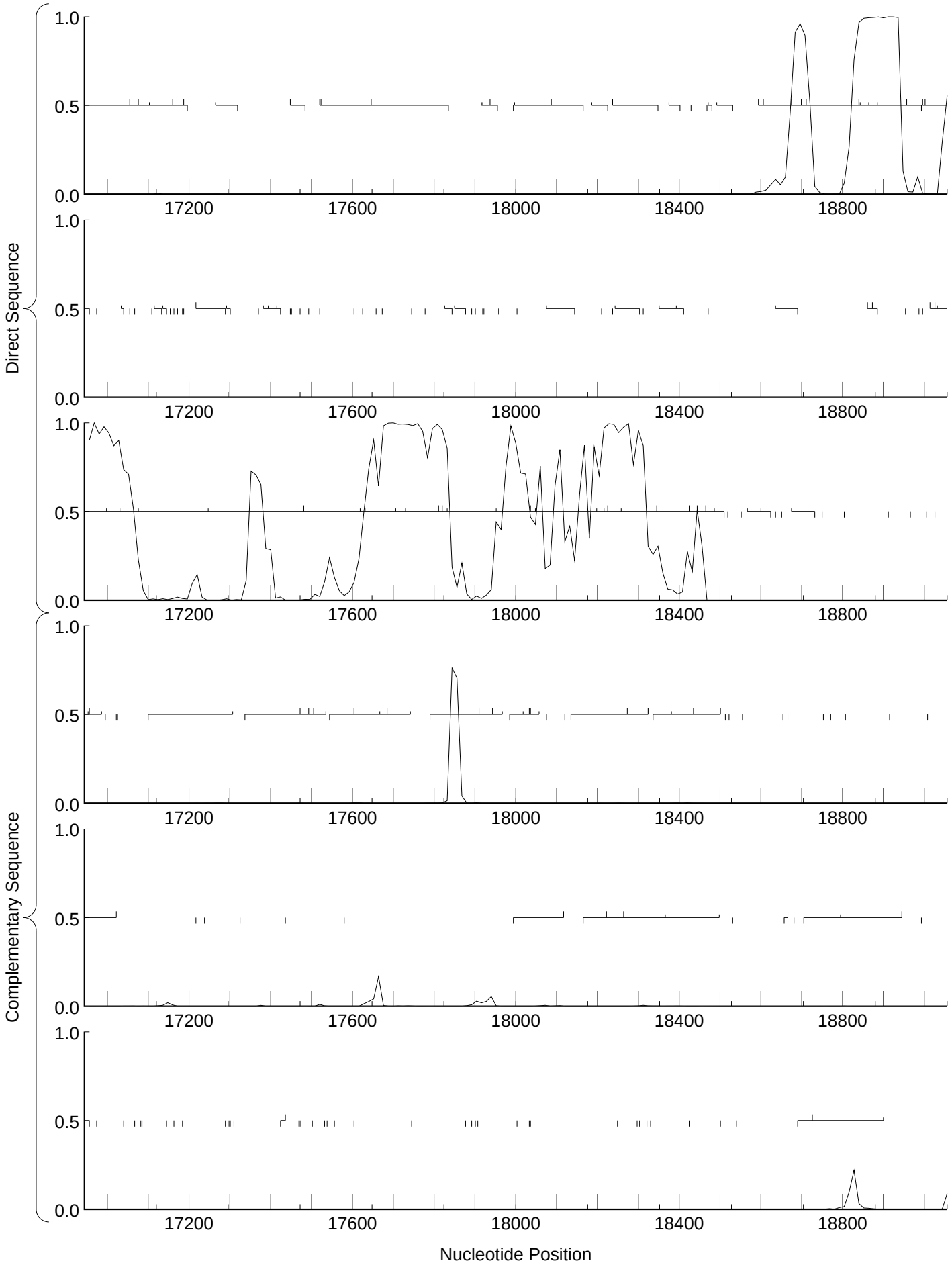
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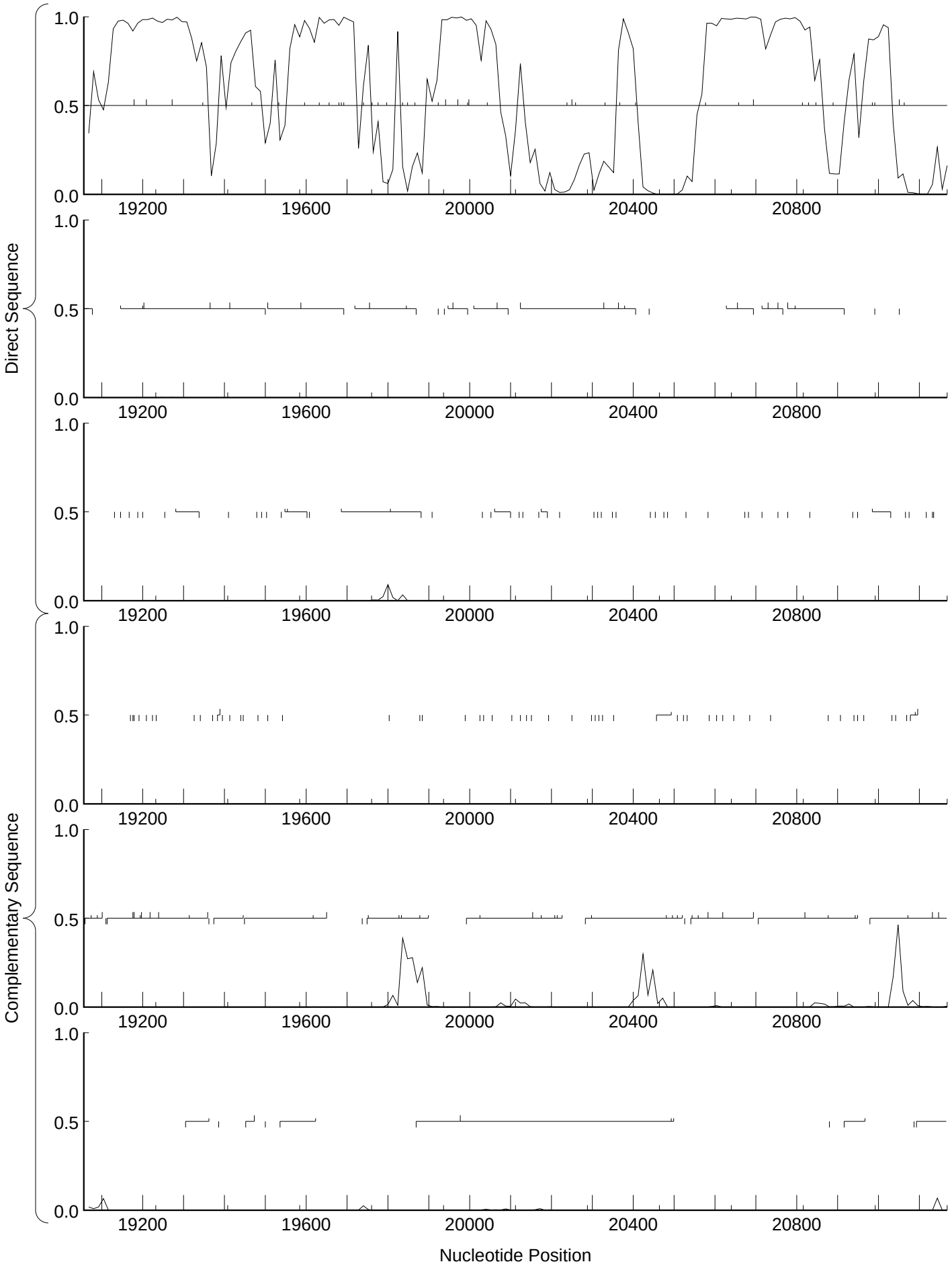
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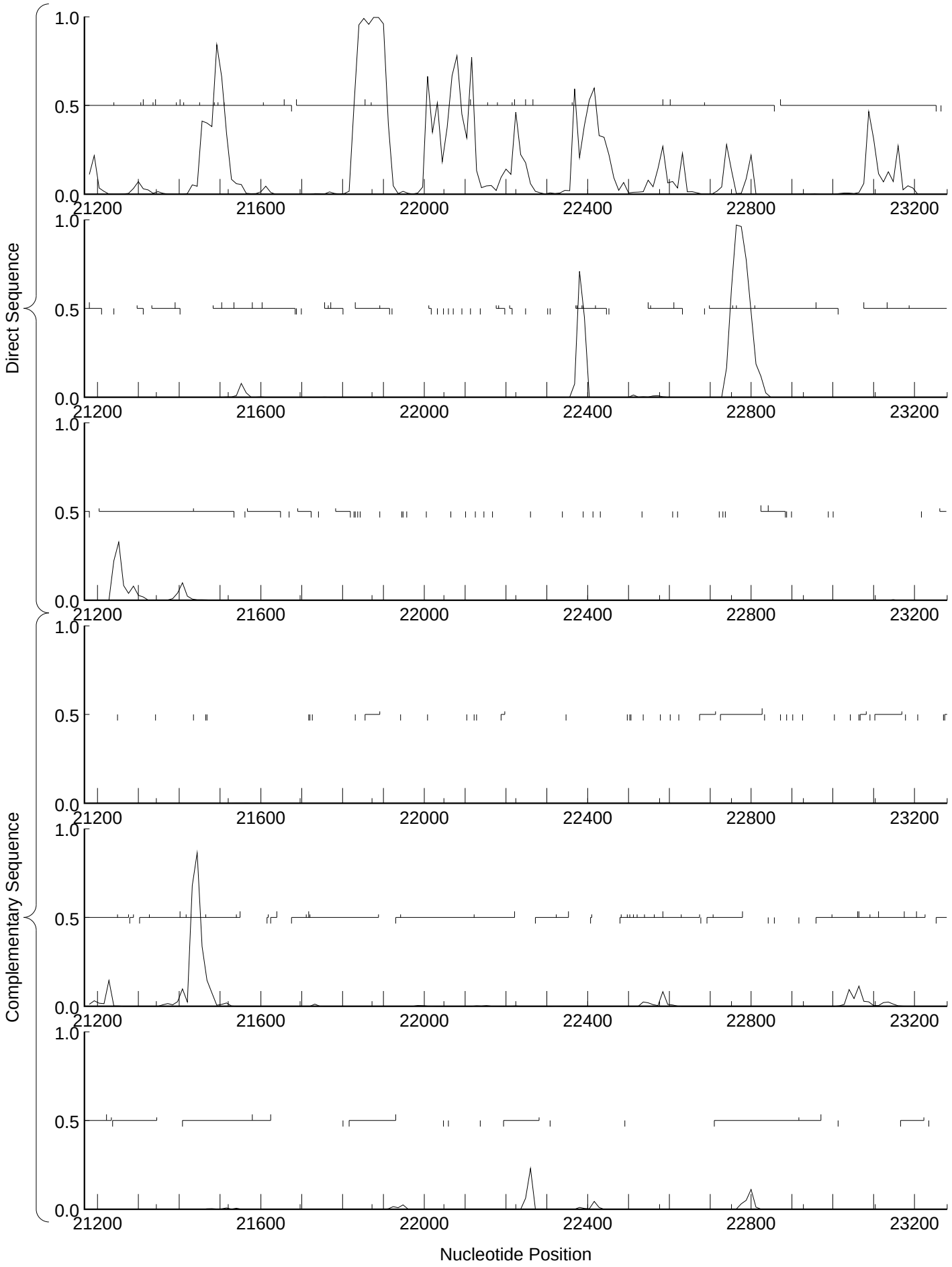
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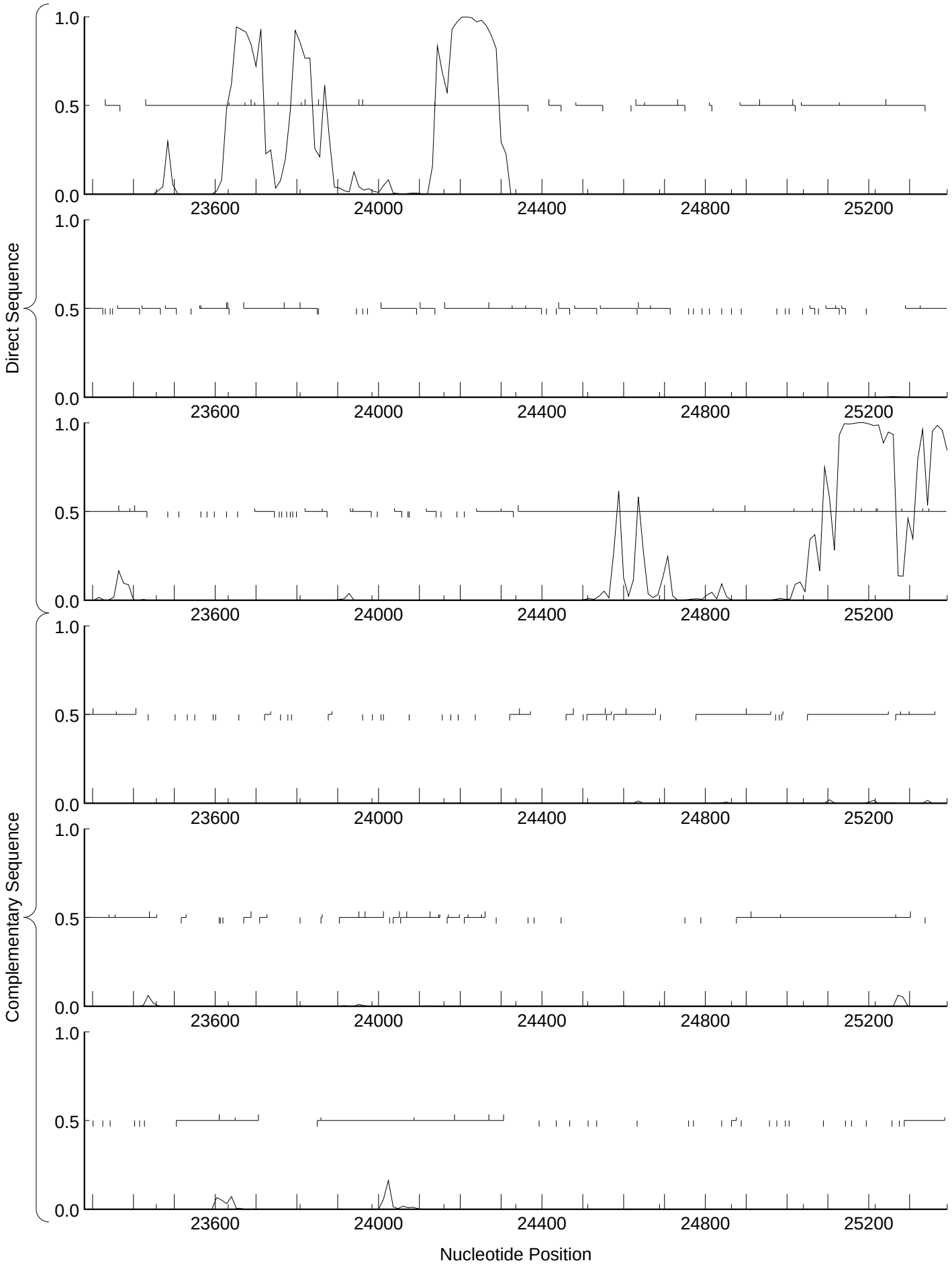
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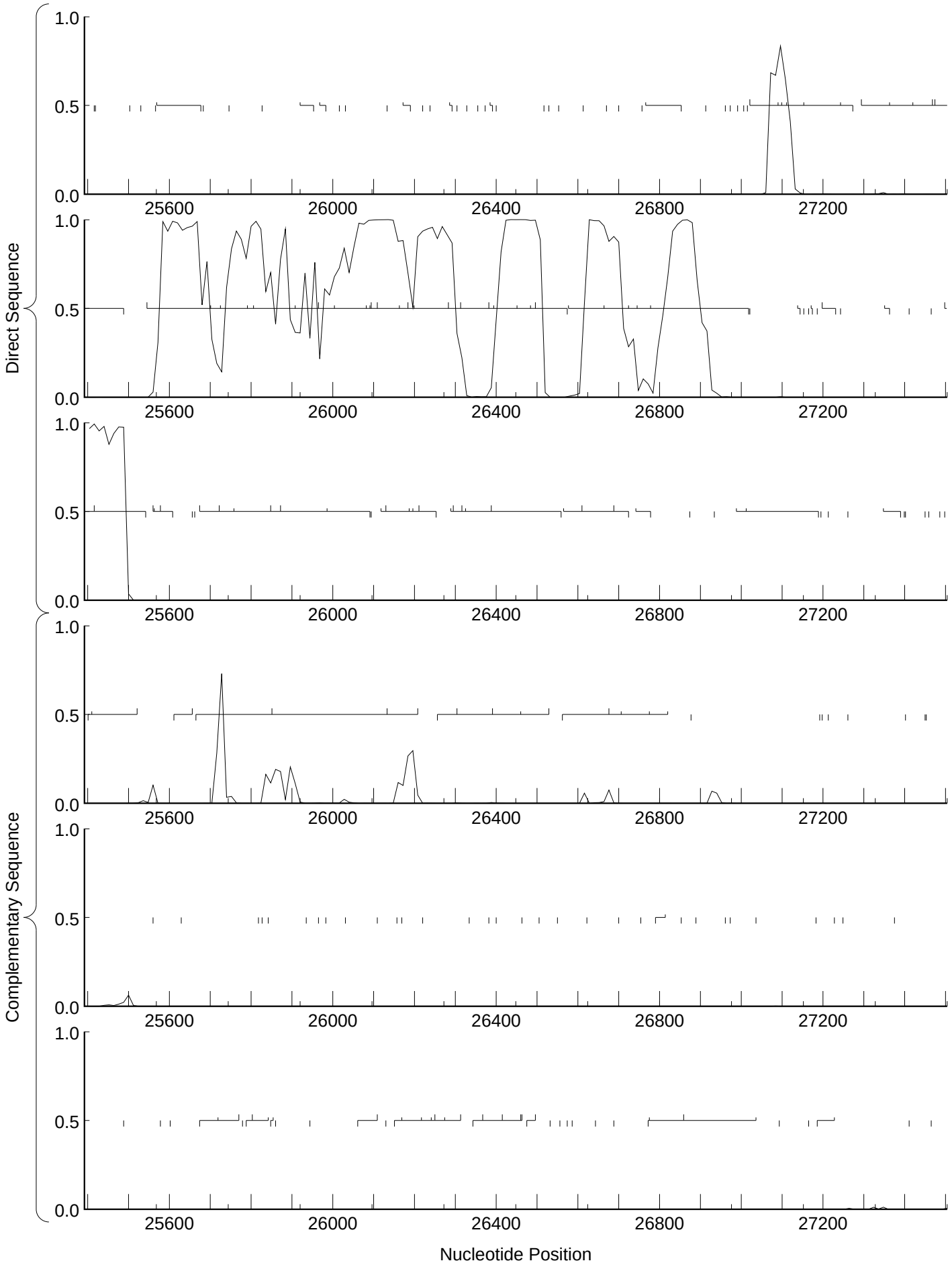
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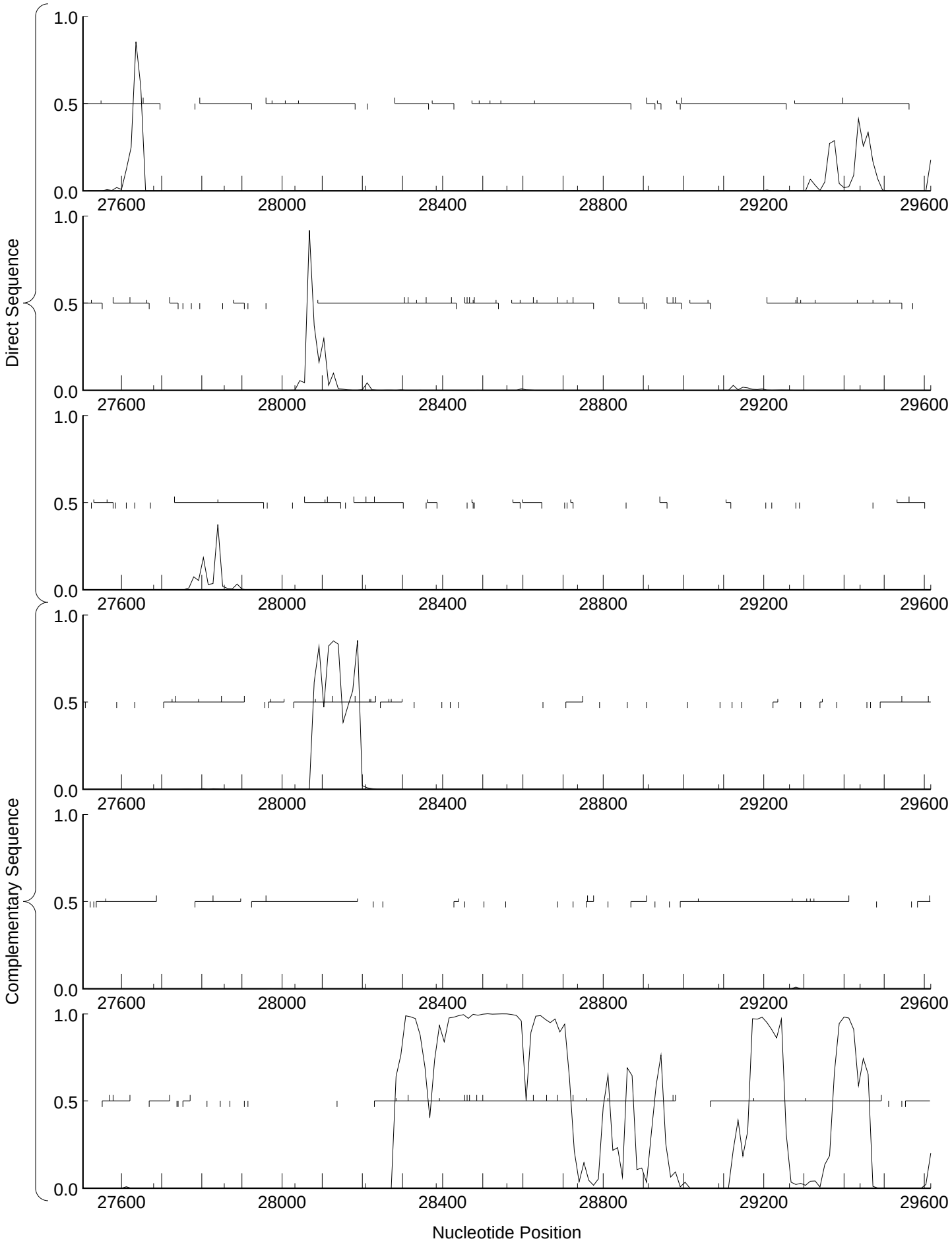


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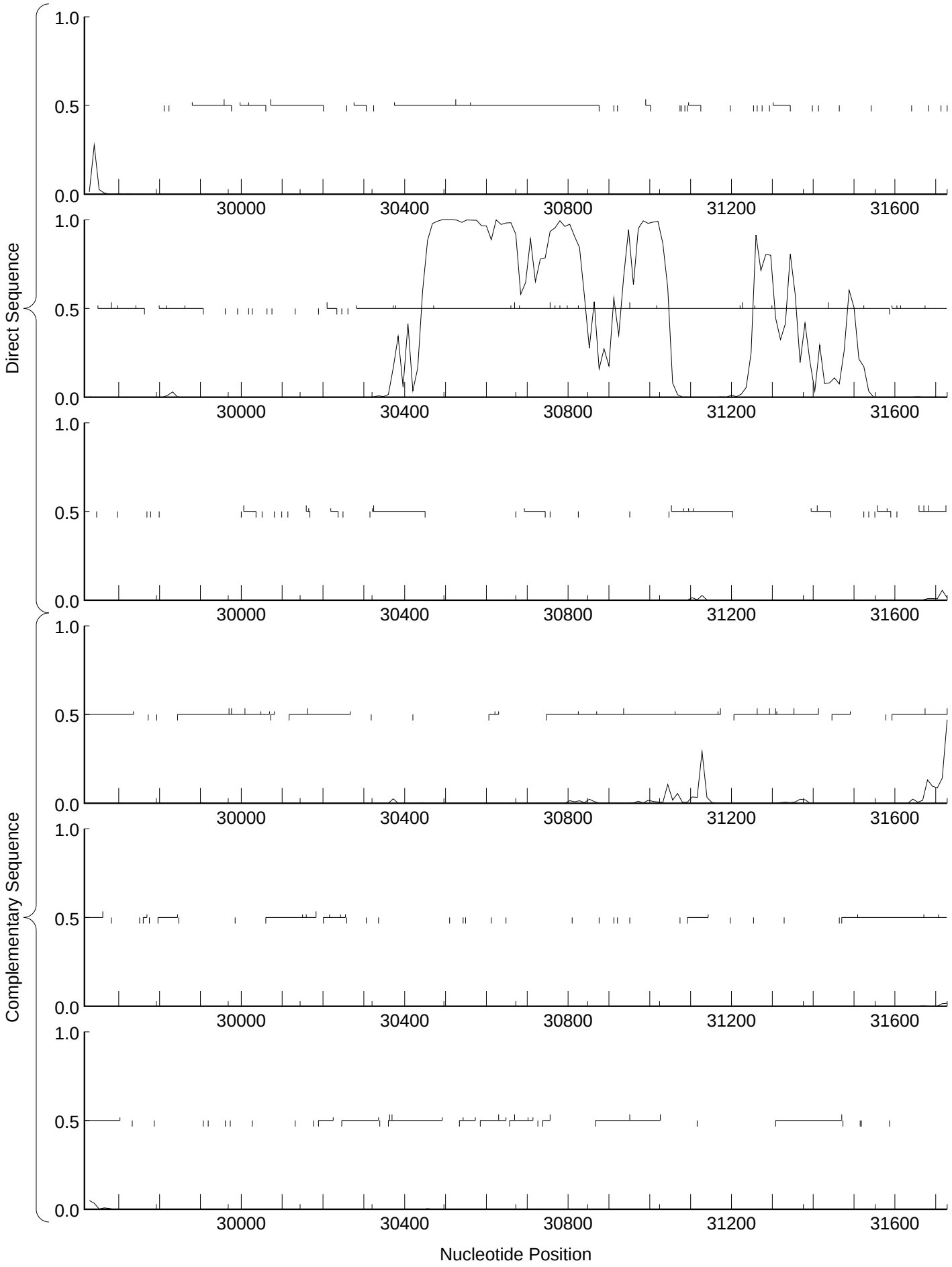


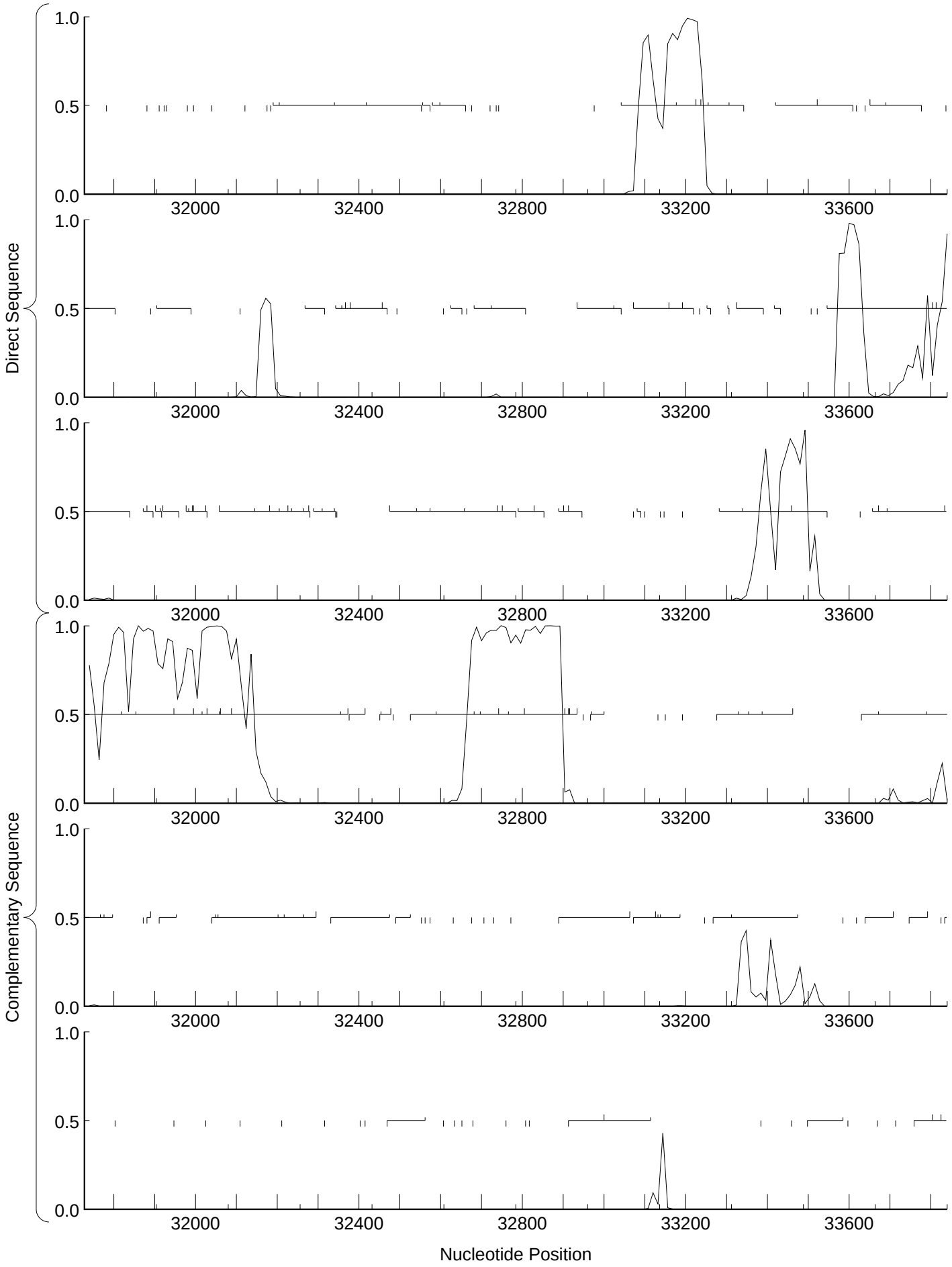
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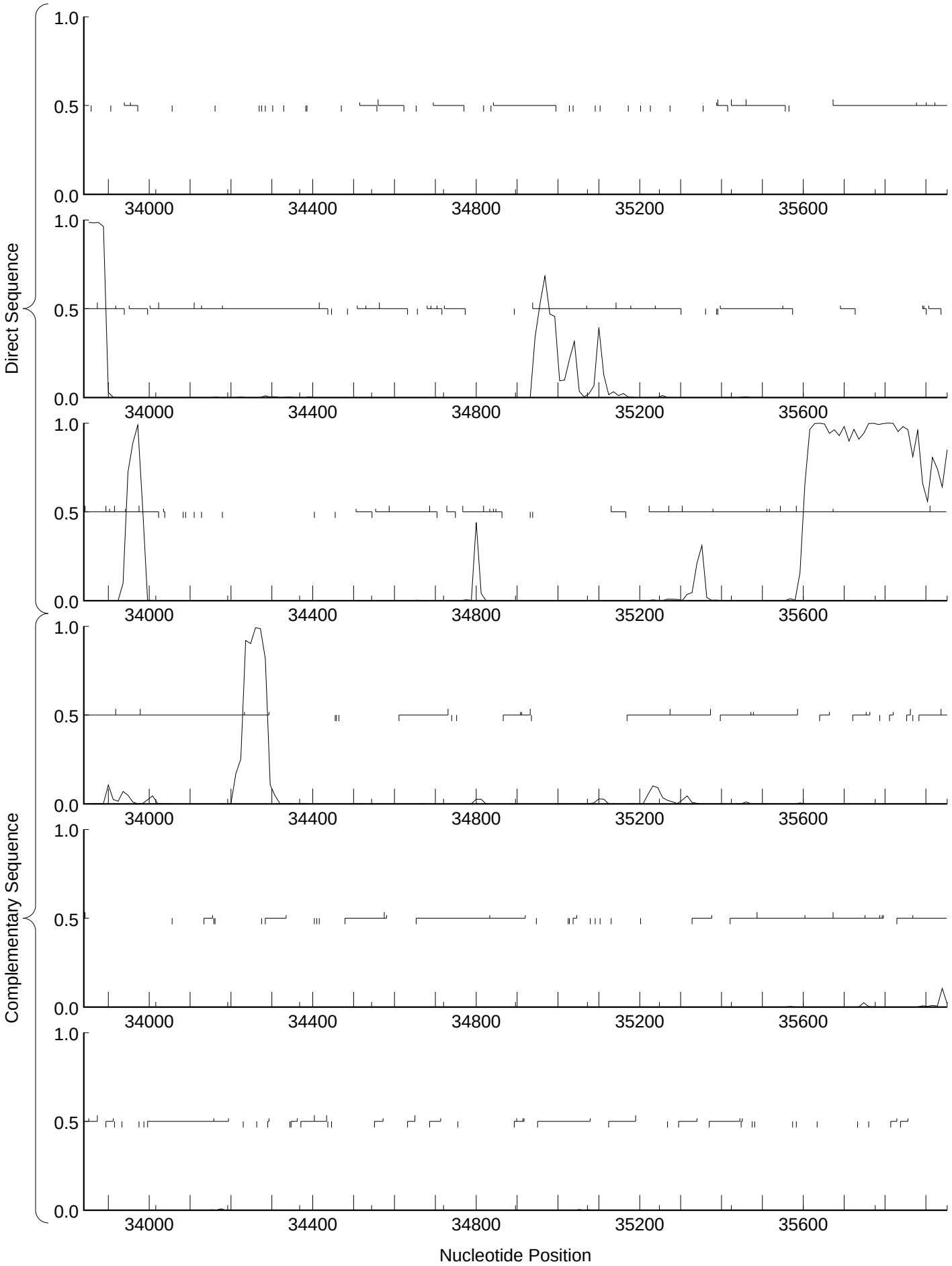

Direct Sequence
Complementary Sequence
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0.5
0.0
27600
28000
28400
28800
29200
29600
Nucleotide Position

f complete sequence, 73210 bp including 10-base 3' overhang (TCGAACGGCC), Cluster L1, Order 4, Window 96, Step 12, 16/36

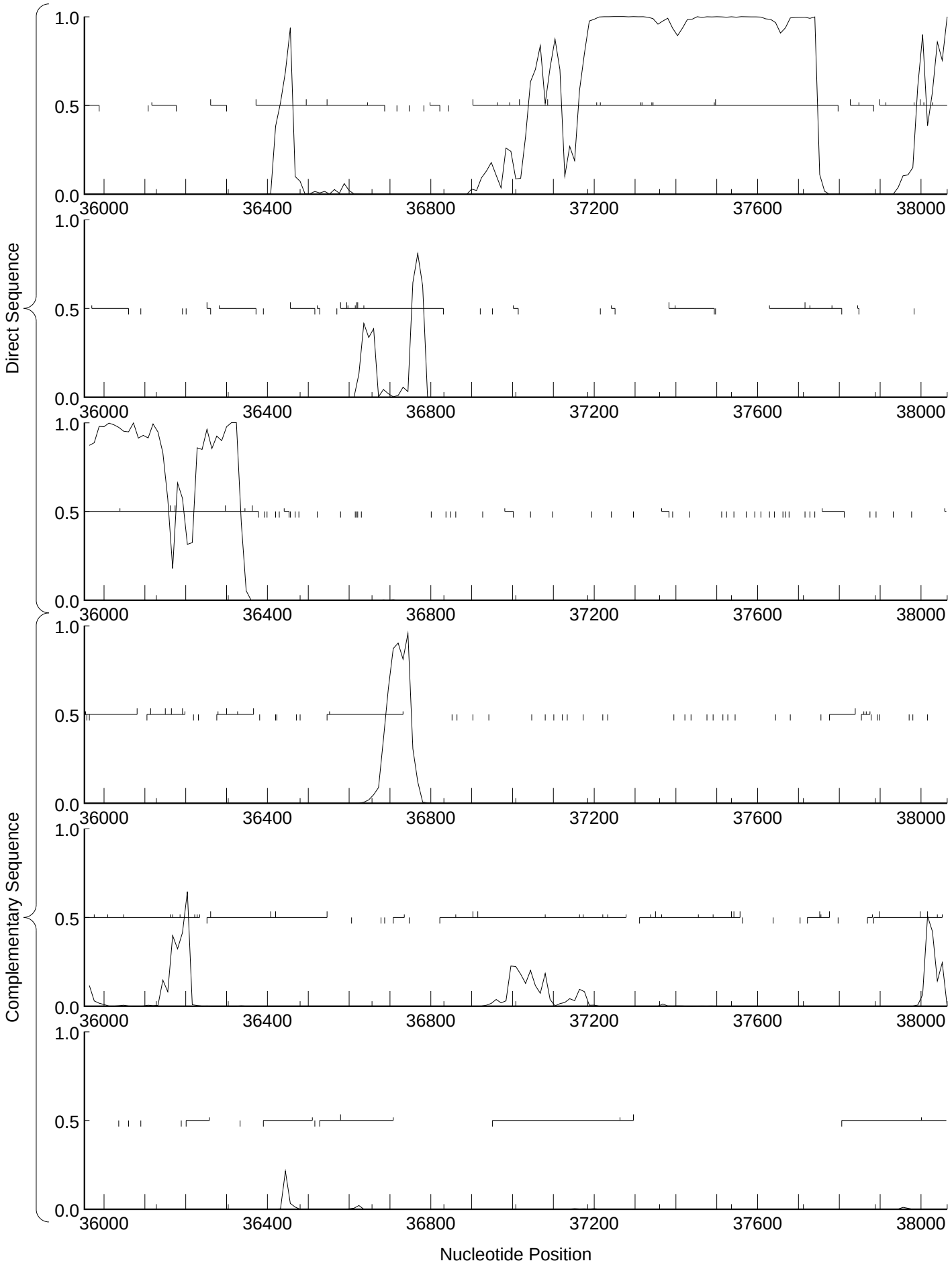



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
32000
32400
32800
33200
33600
Nucleotide Position

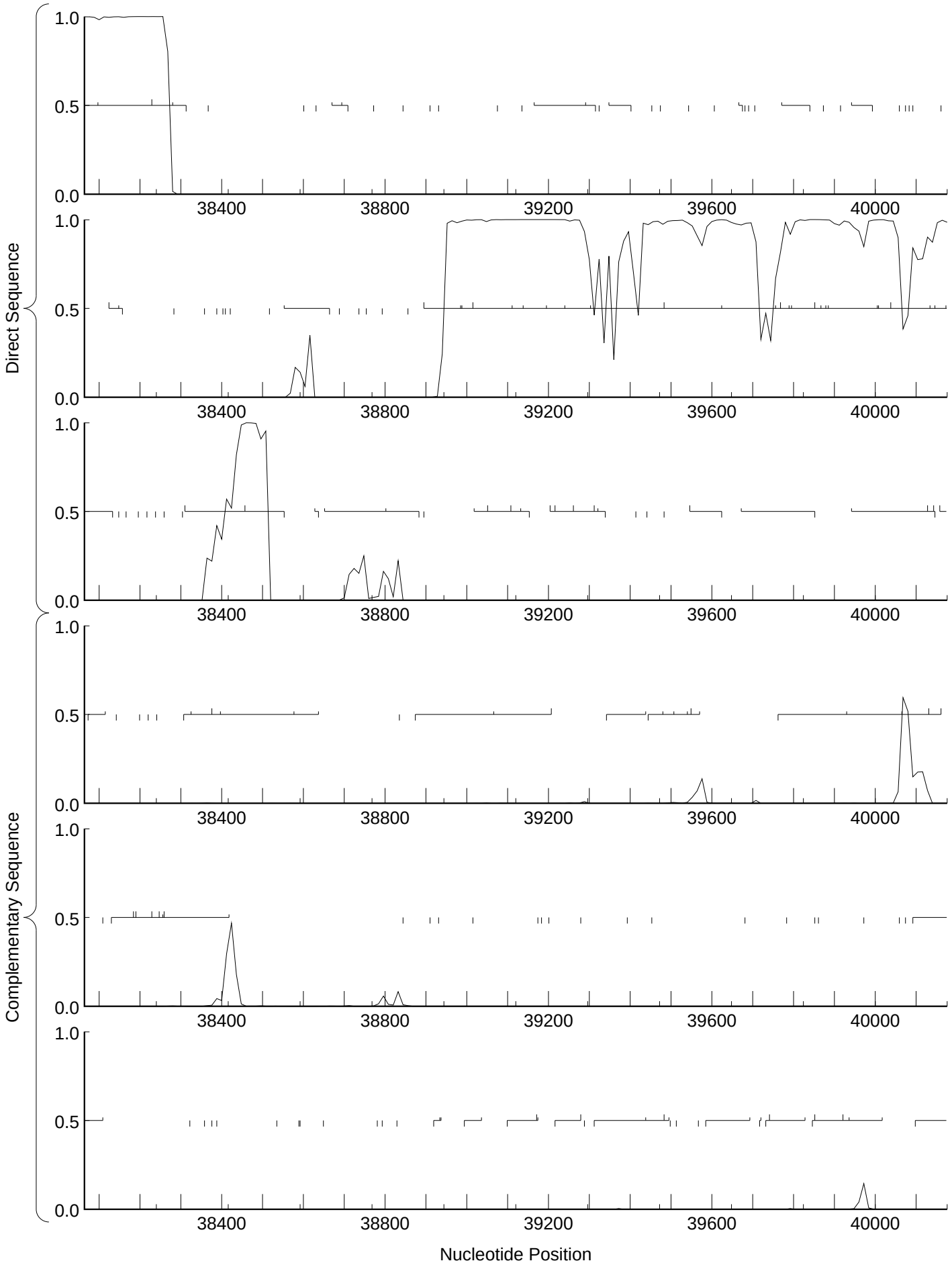
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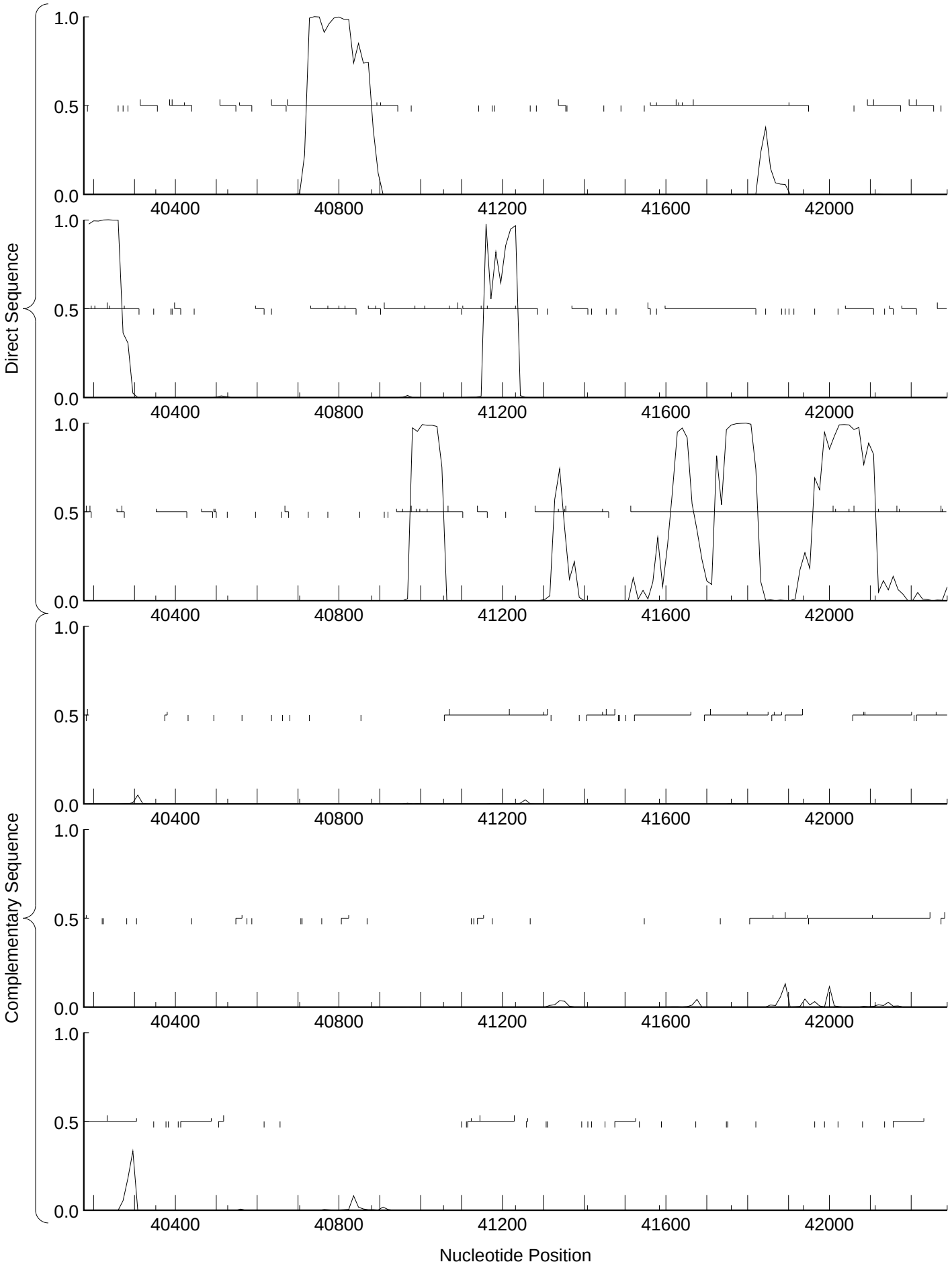
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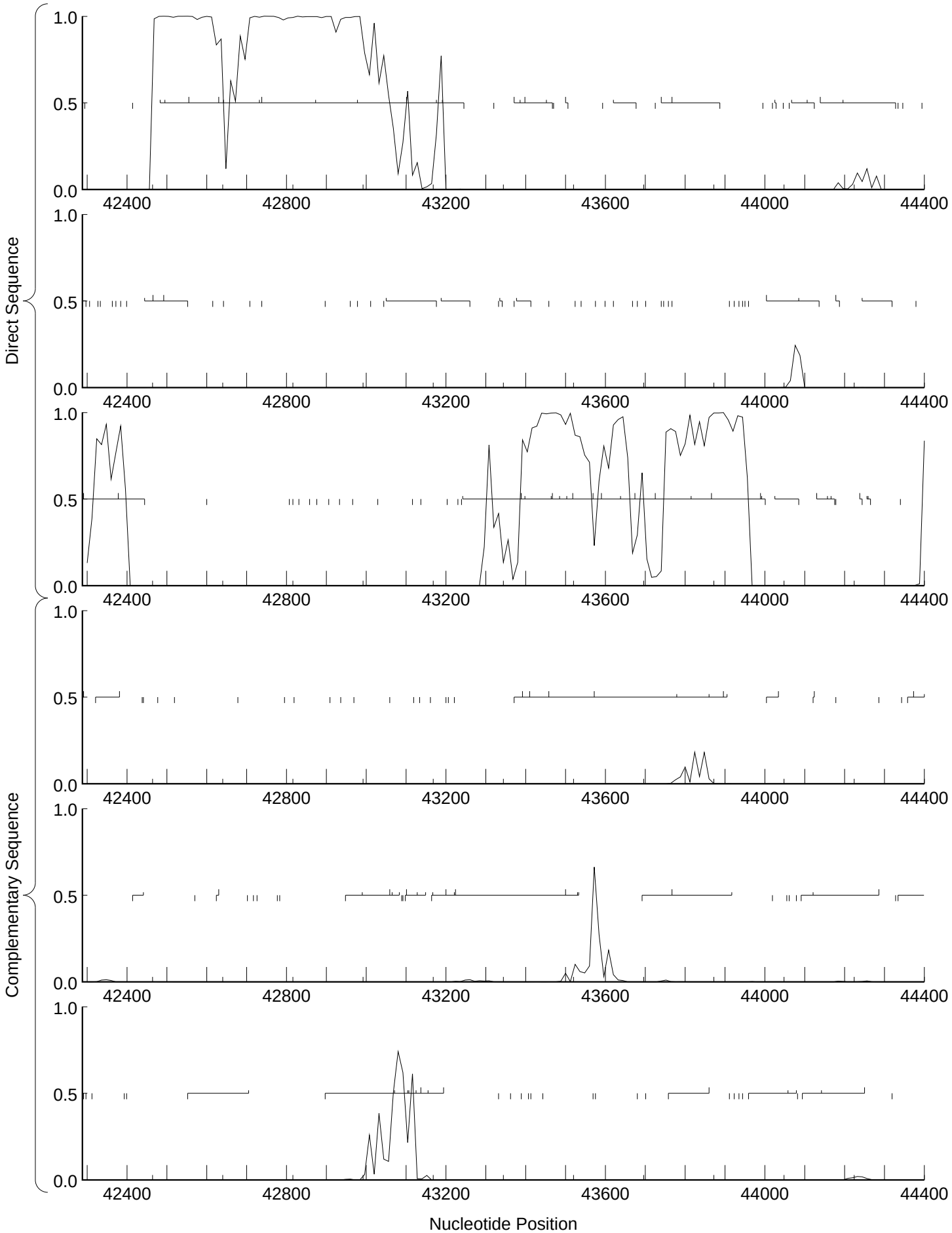


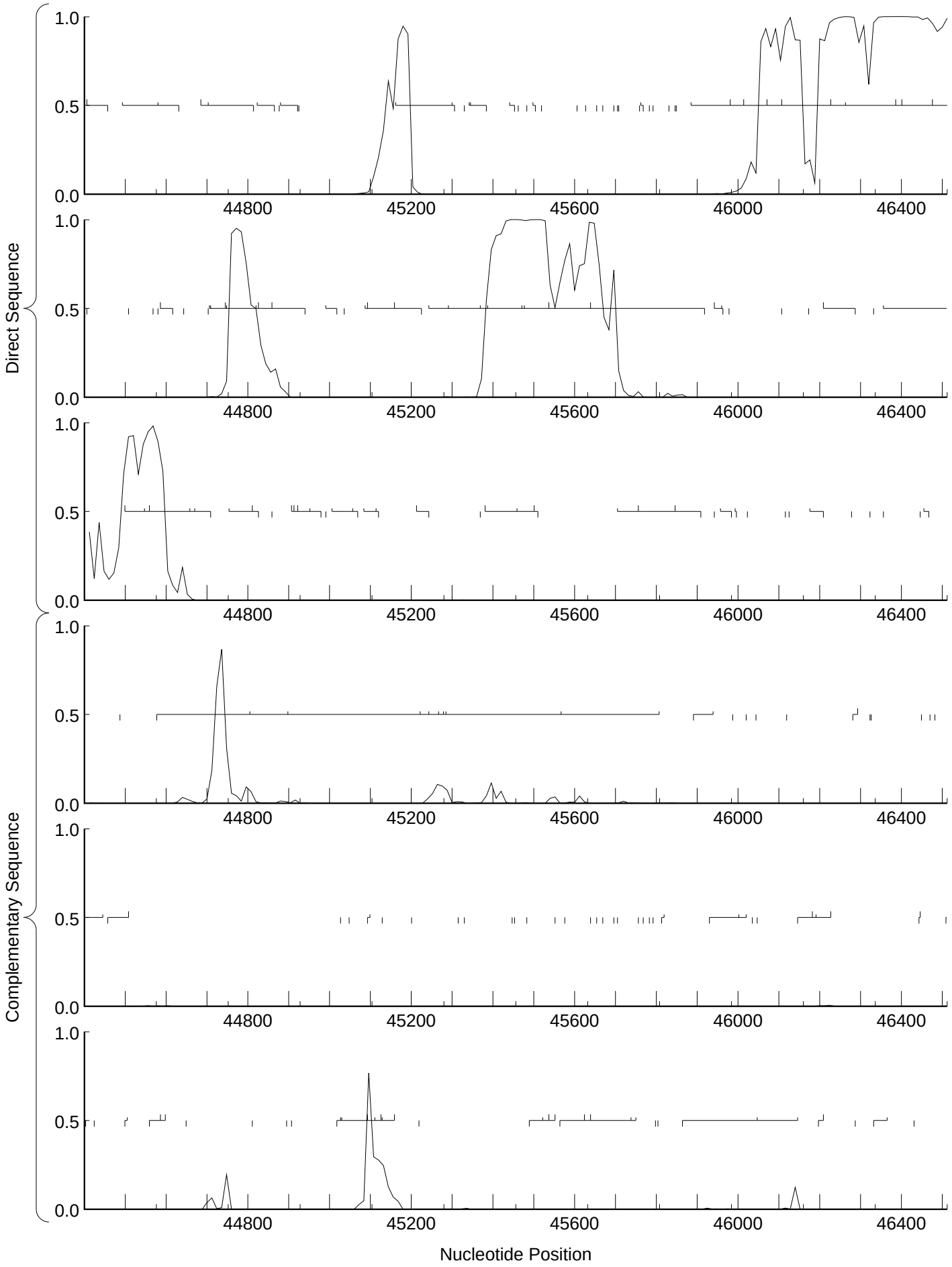
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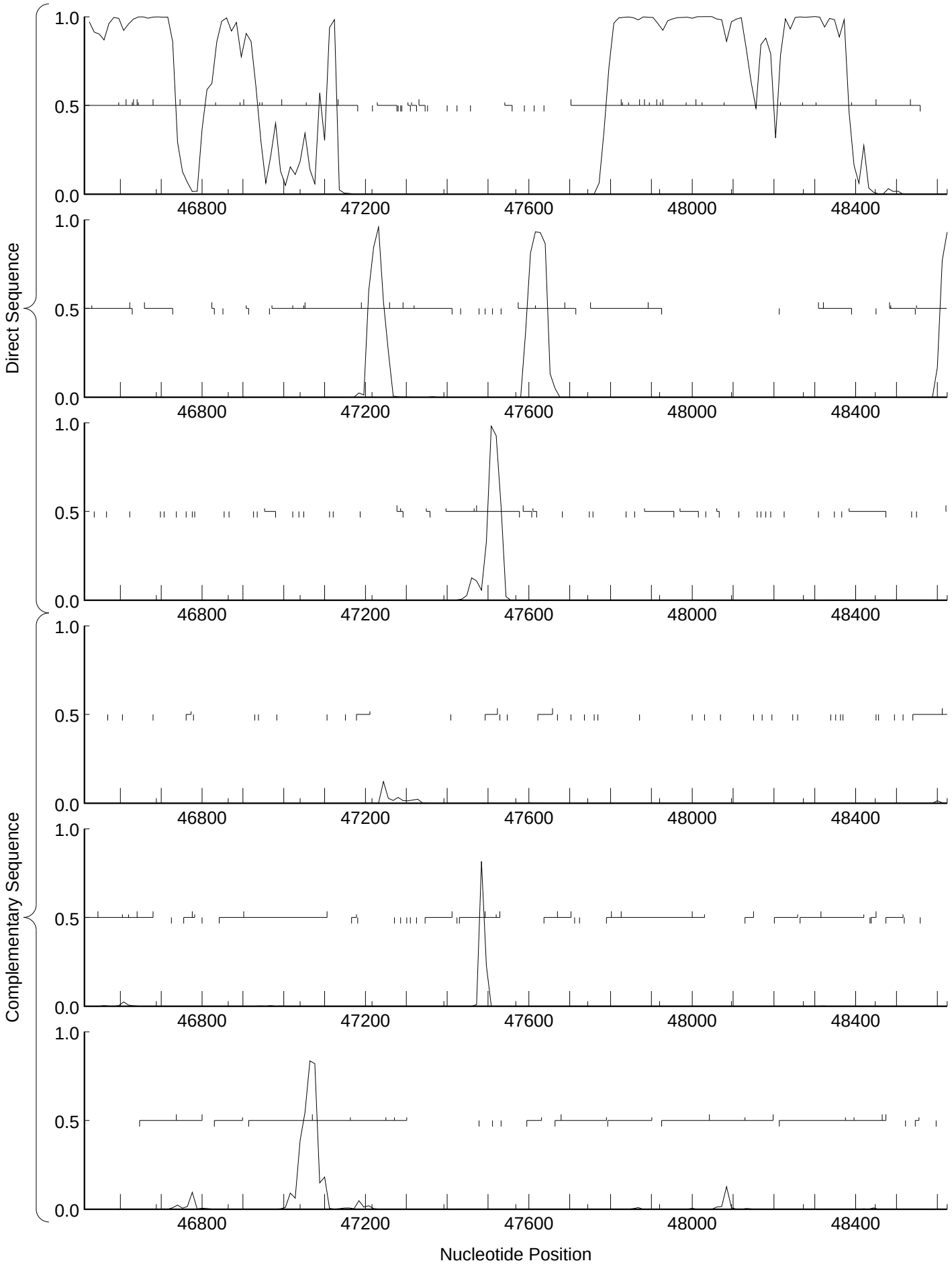
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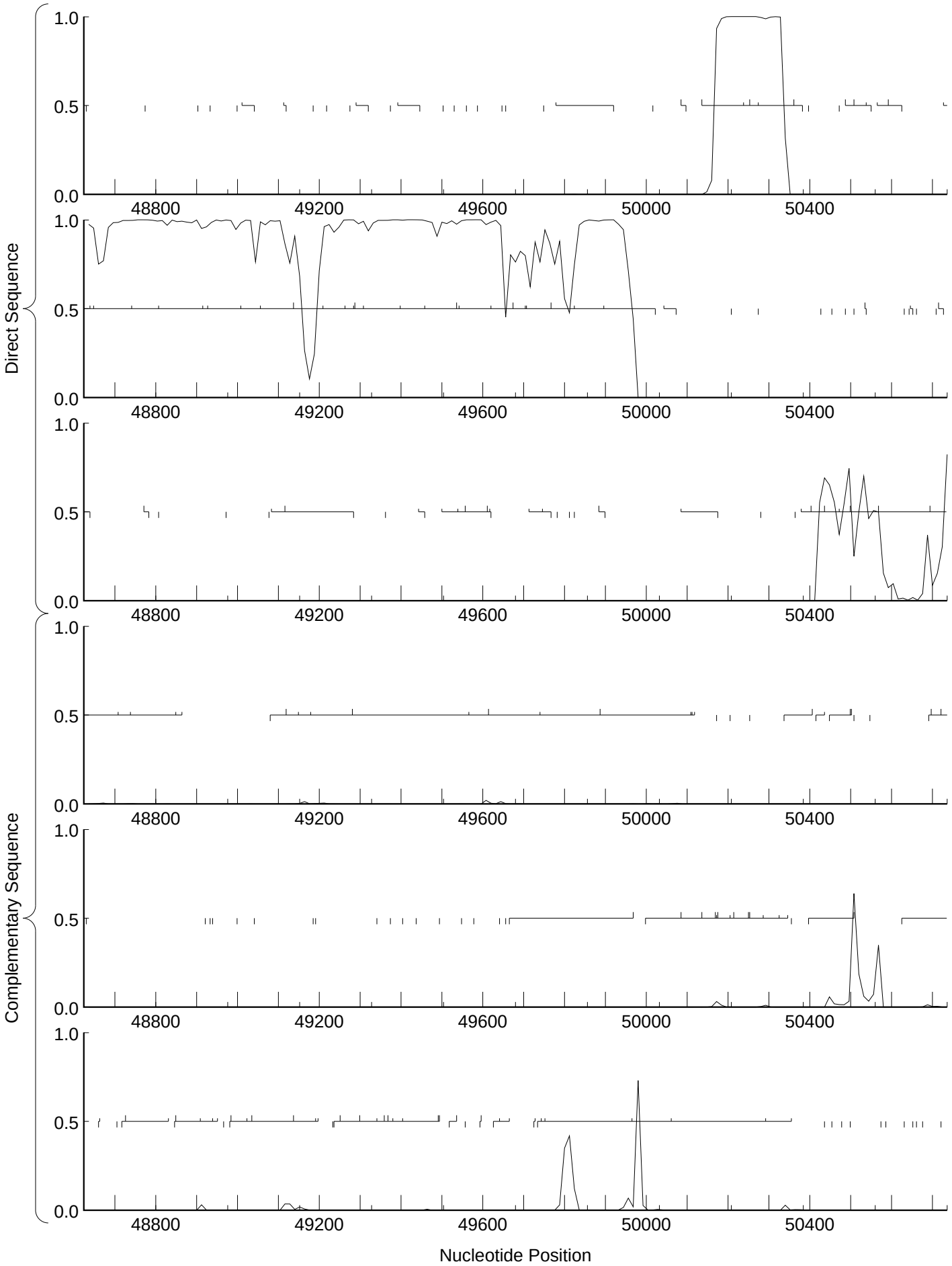

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
42400
42800
43200
43600
44000
44400
Nucleotide Position


Direct Sequence
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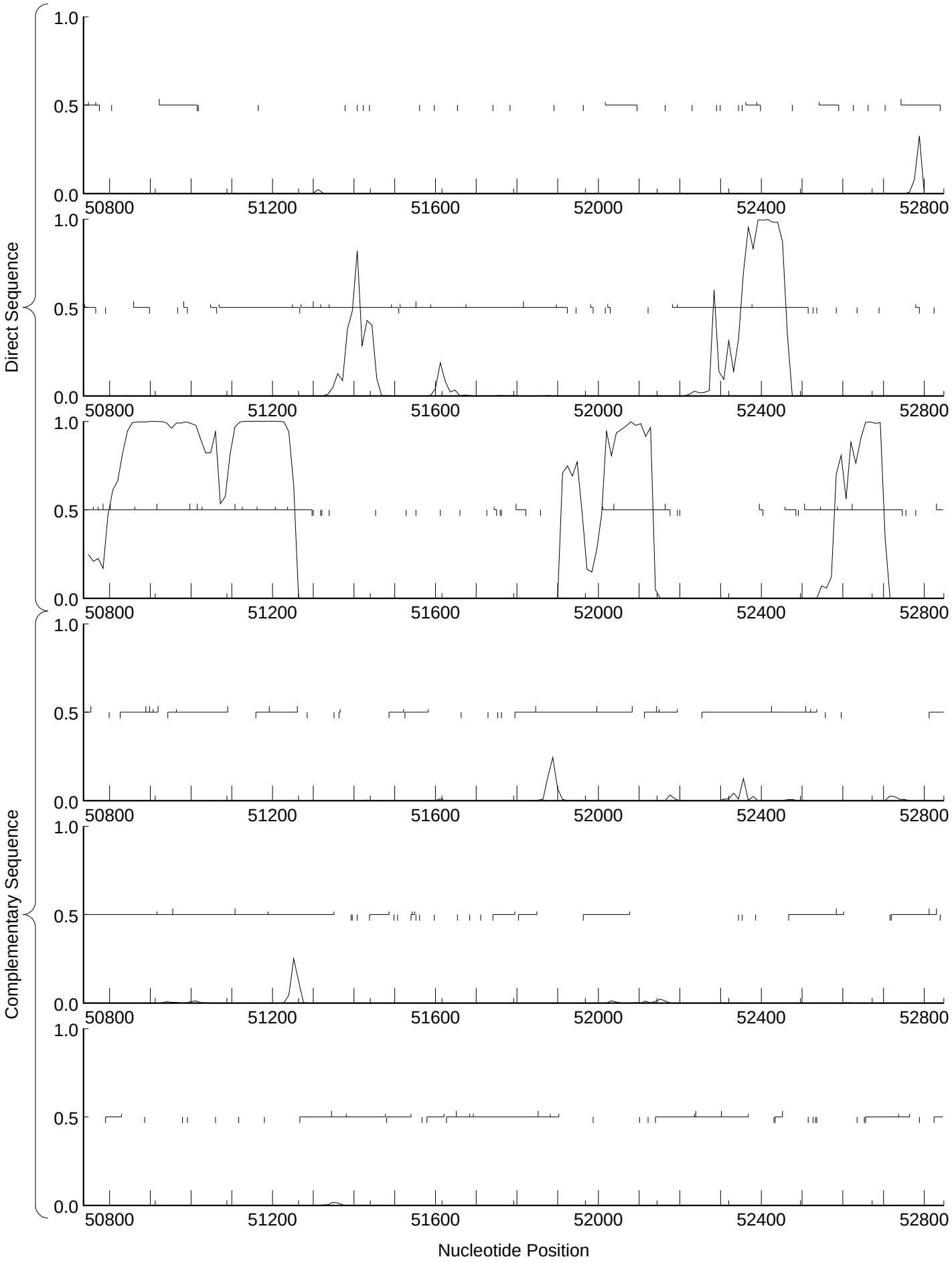
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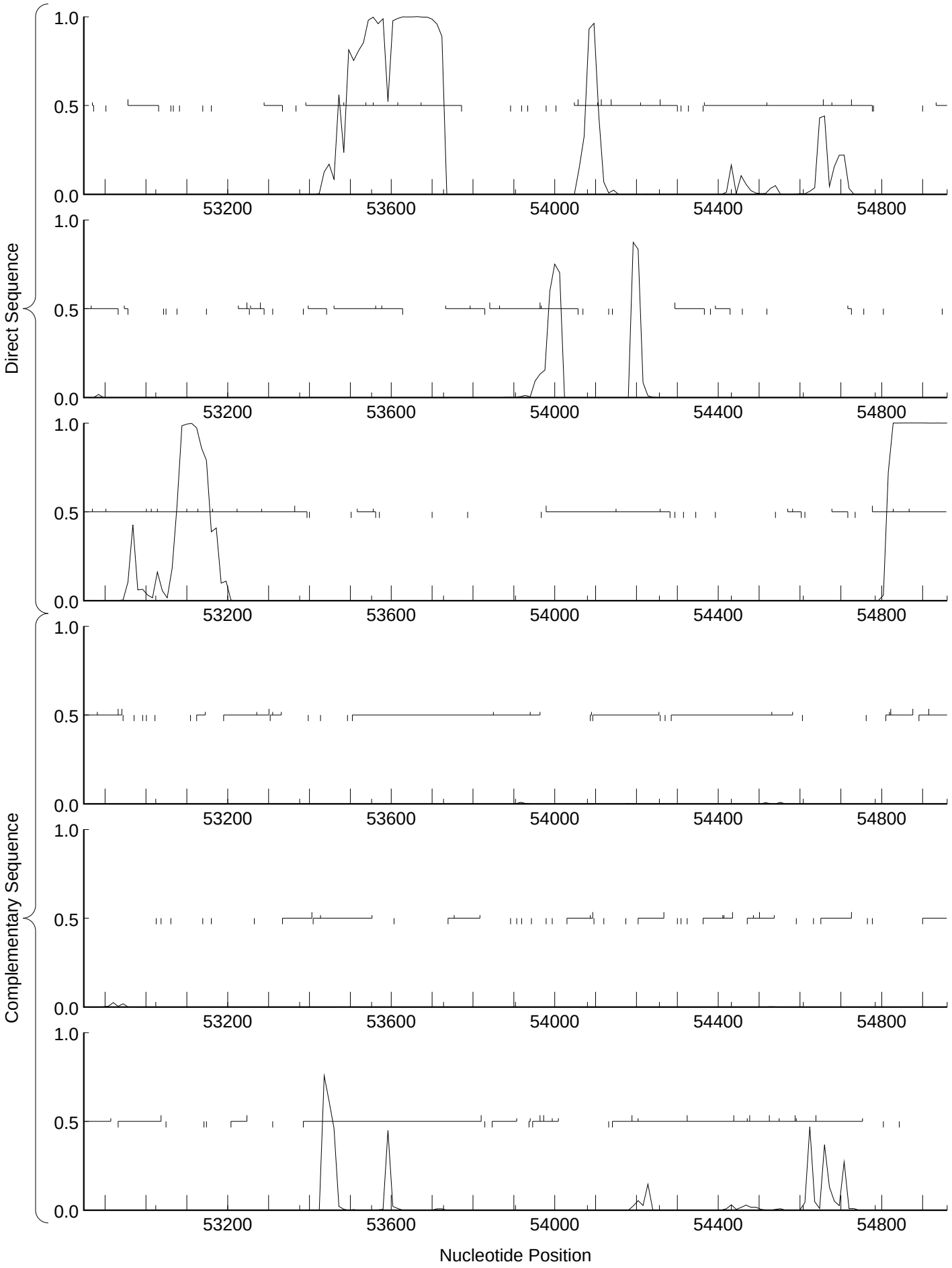
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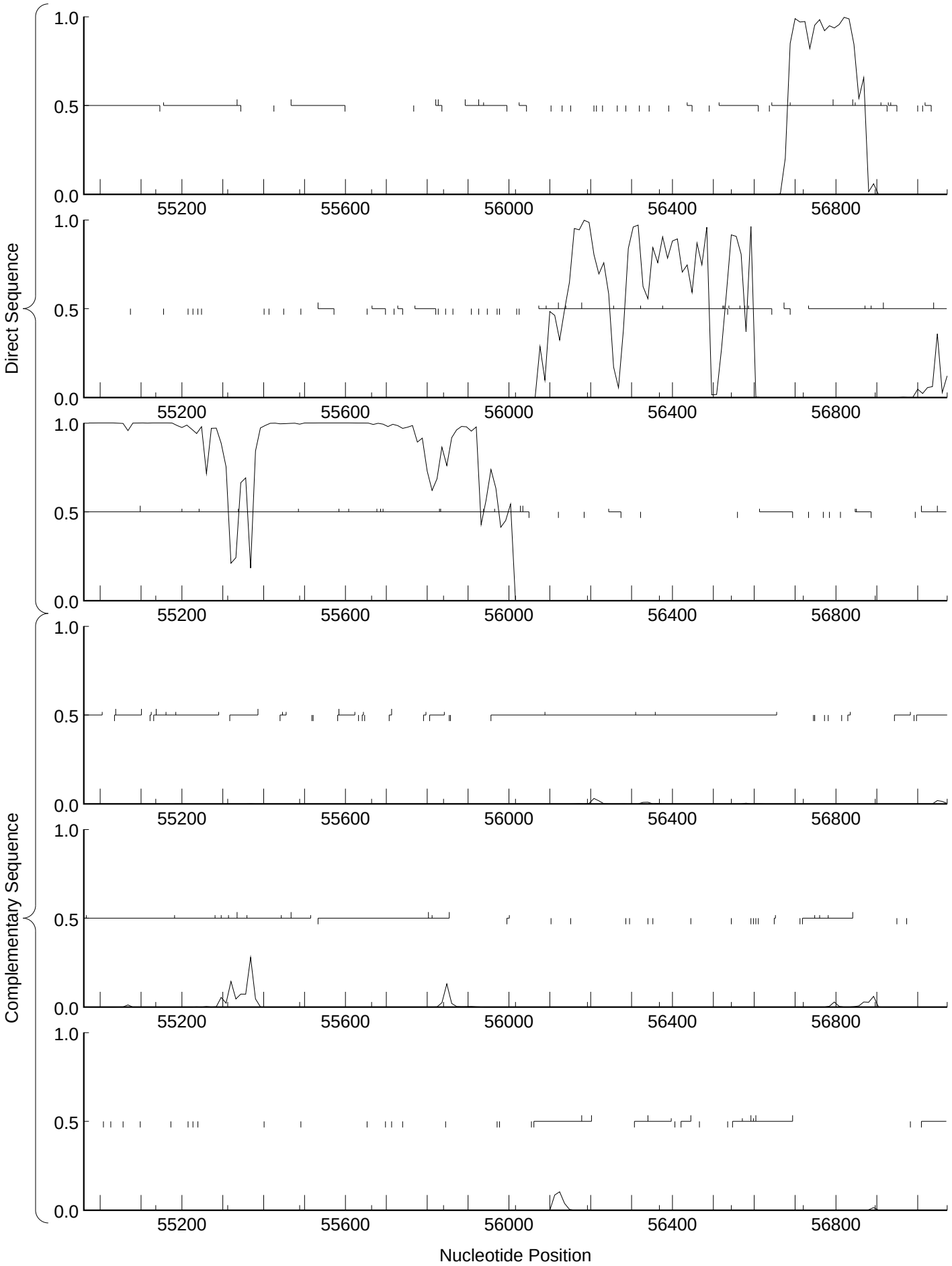
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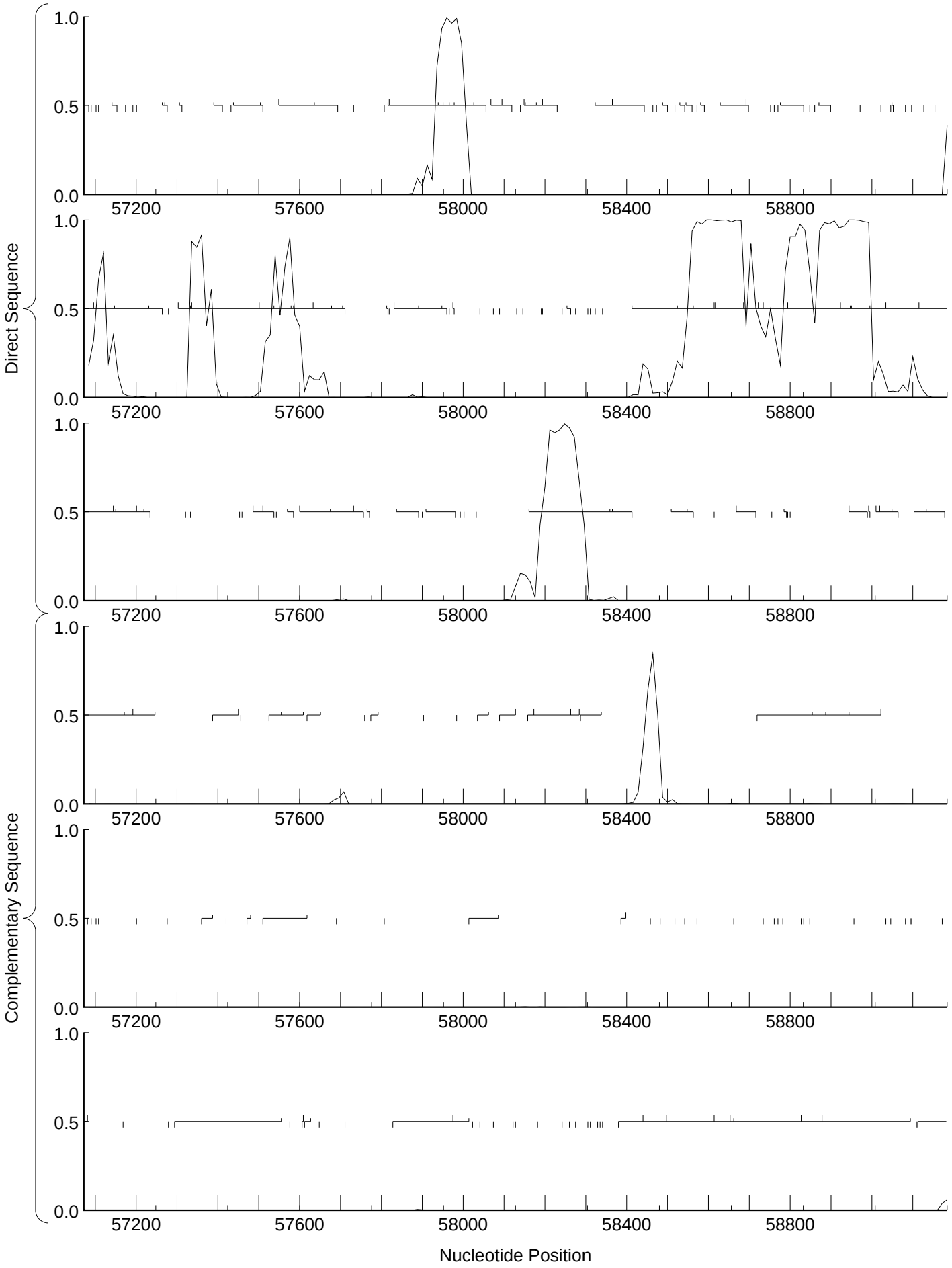
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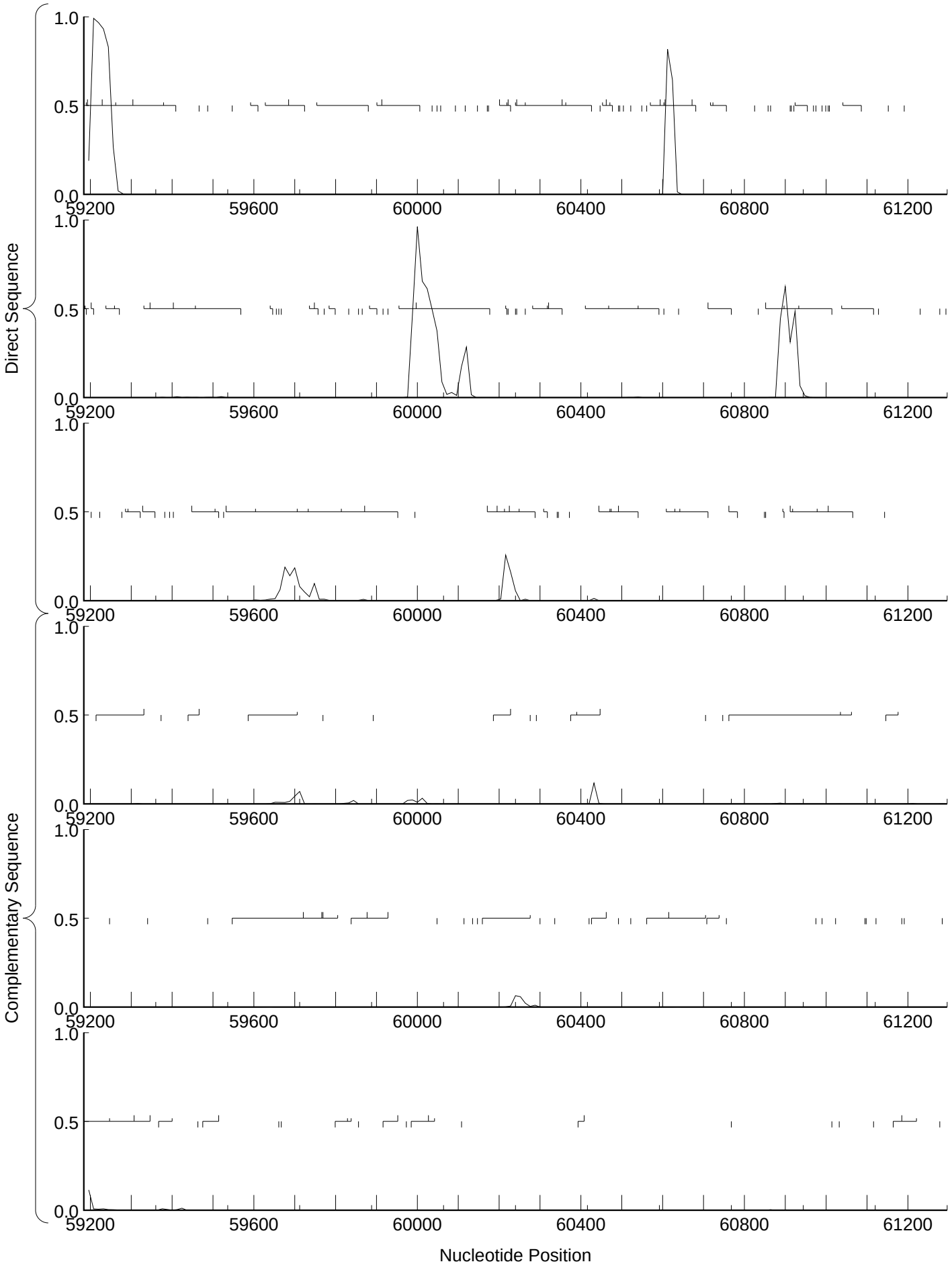
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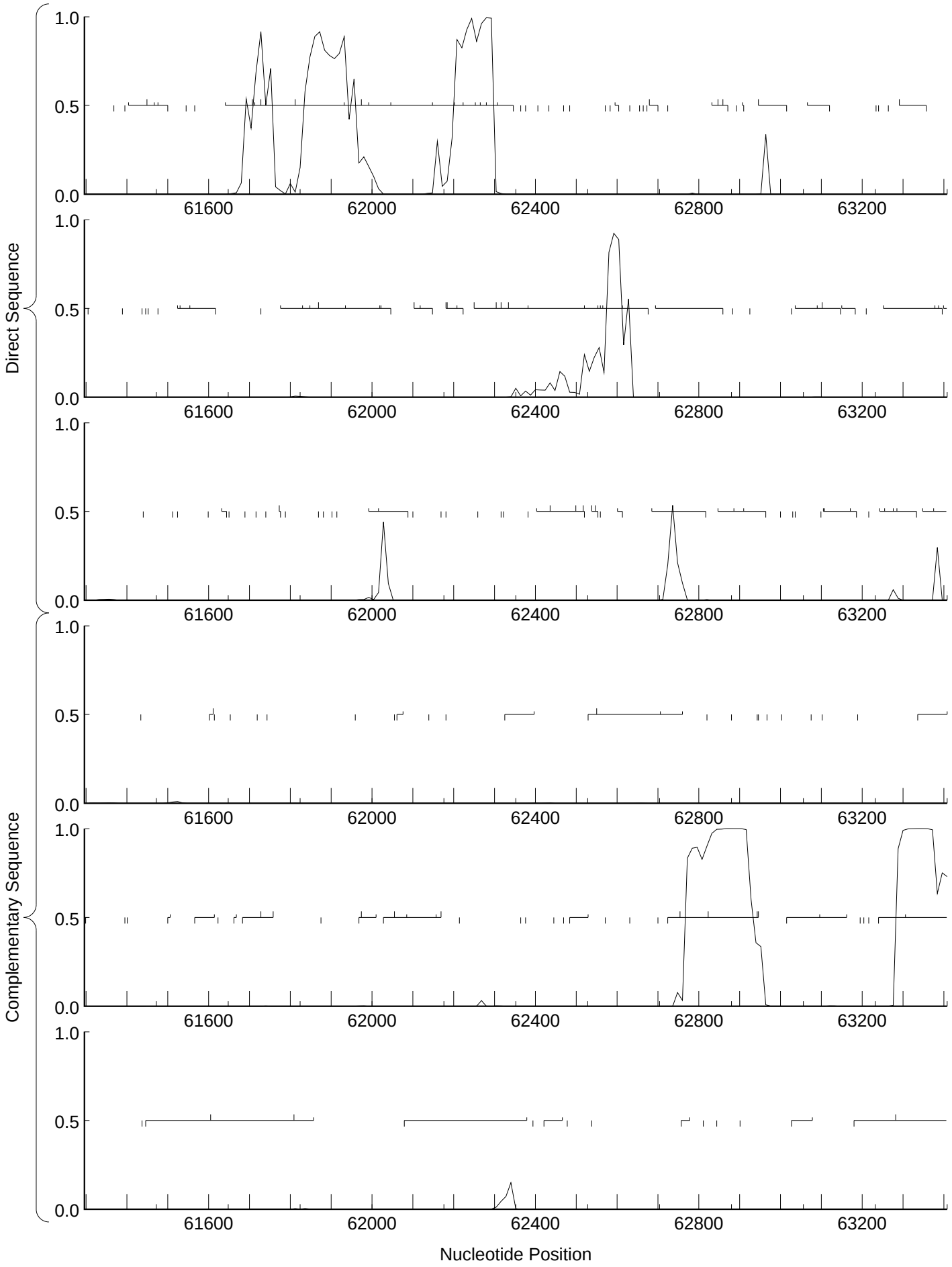
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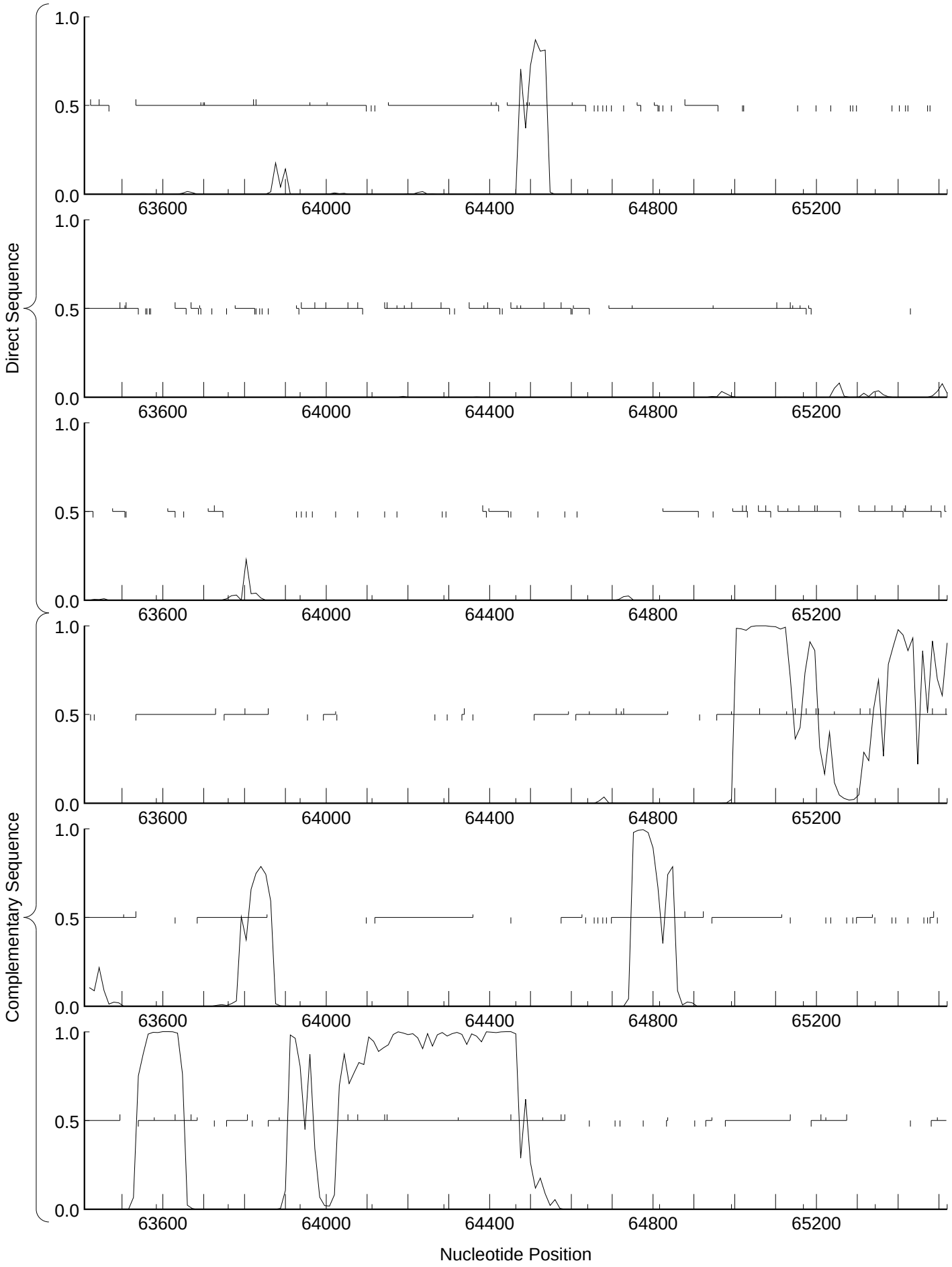
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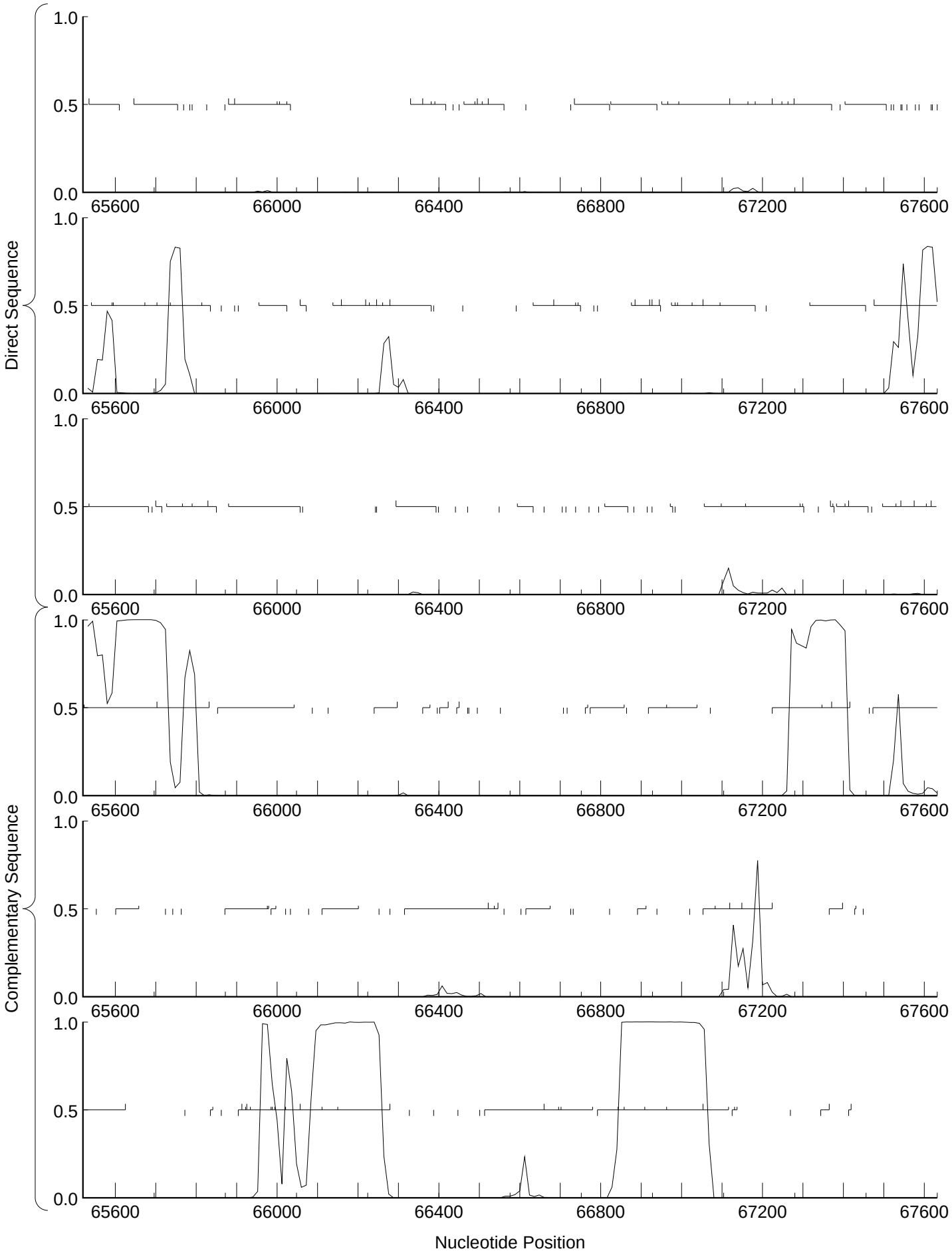
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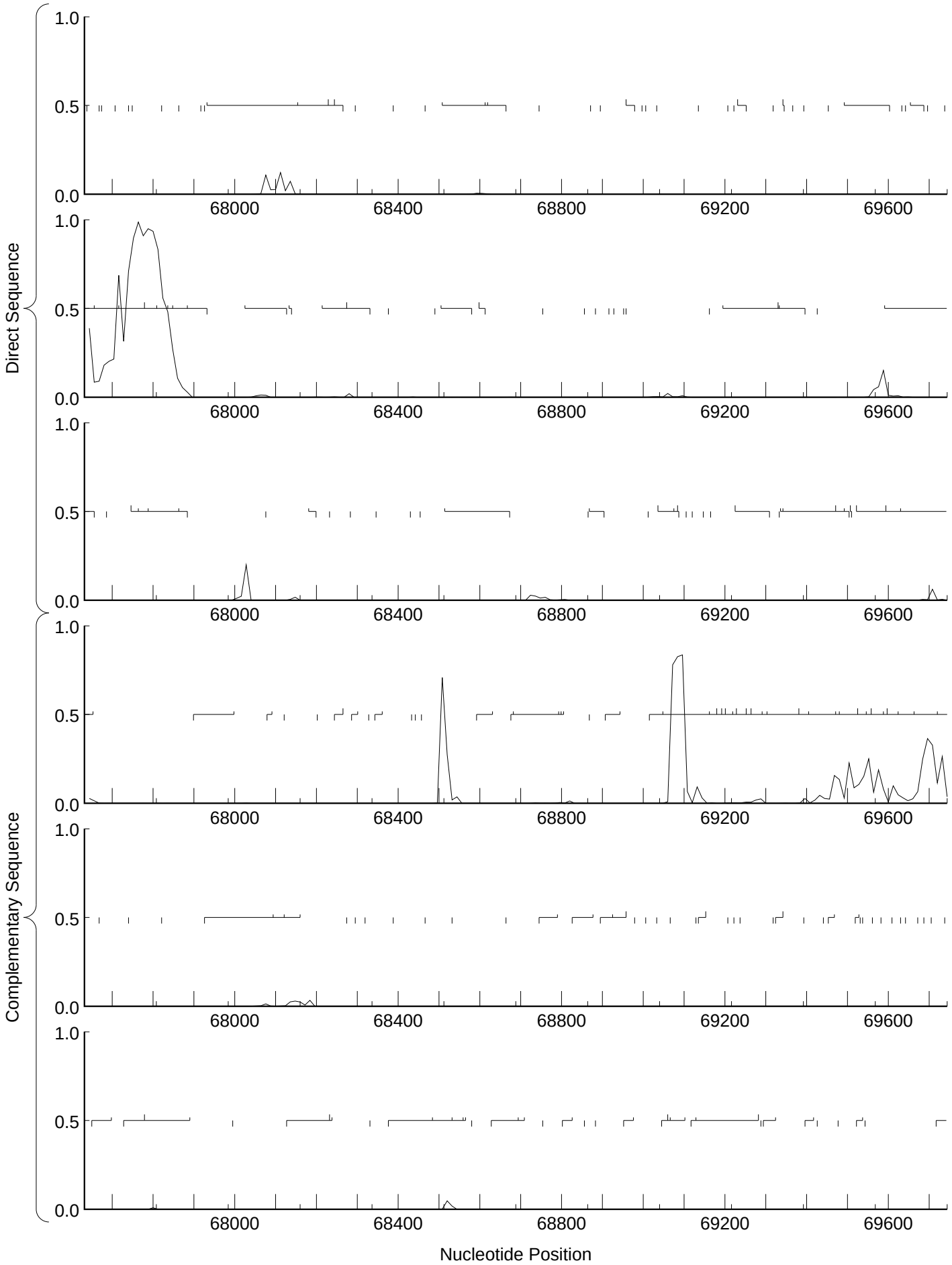
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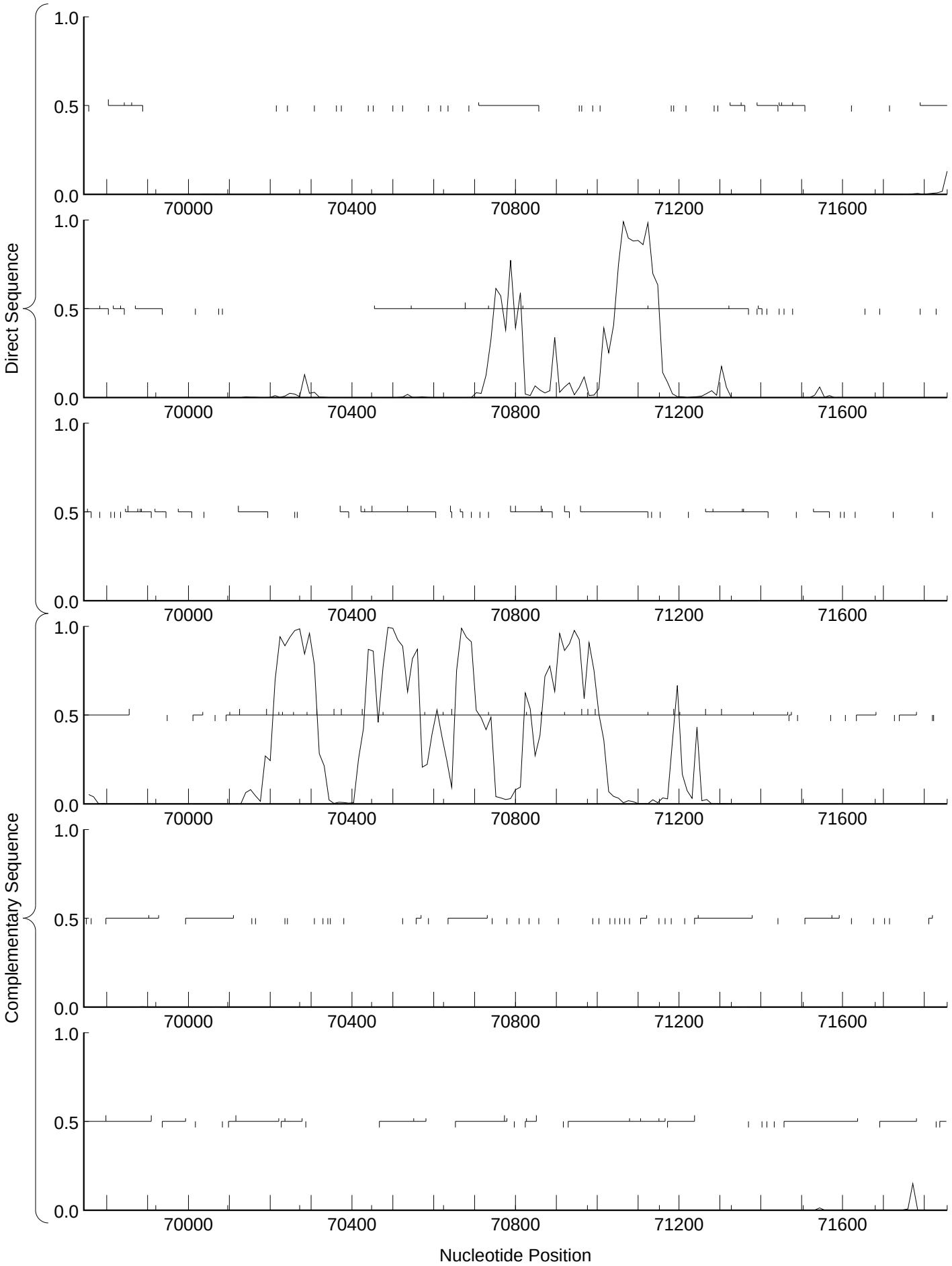
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f complete sequence, 73210 bp including 10-base 3' overhang (TCGAACGGCC), Cluster L1, Order 4, Window 96, Step 12, 35/36



f complete sequence, 73210 bp including 10-base 3' overhang (TCGAACGGCC), Cluster L1, Order 4, Window 96, Step 12, 36/36

