GeneMark

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

PROGRAM INFORMATION

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Sequence         :  Mycobacterium Phage MrMagoo, complete sequence 84,303 bp including 11-base 3’ overhang (ACC
Analysis Date    :  3/28/24 at 14:36:50
Pages            :  41
Sequence Length  :  84303 bp
GC Content       :   60.80%

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

PS-Version       :  1.2

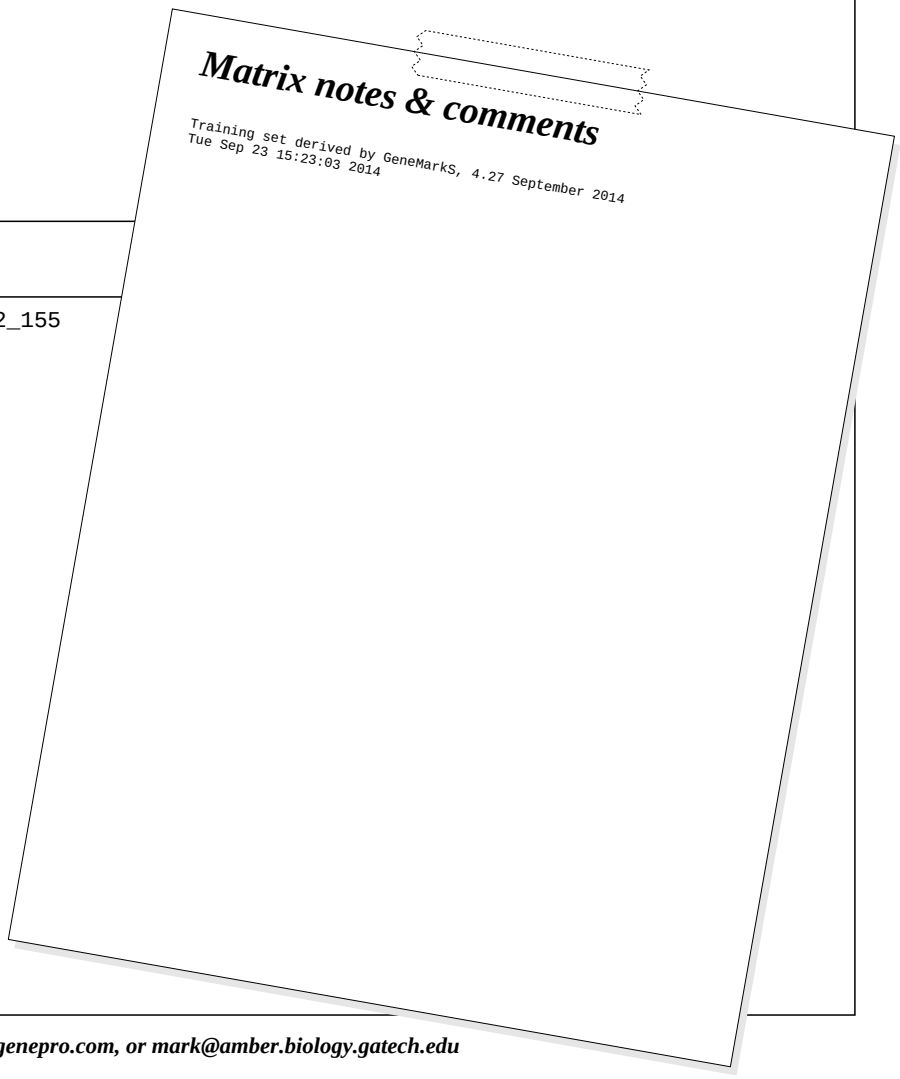
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MATRIX INFORMATION

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Author     :  -

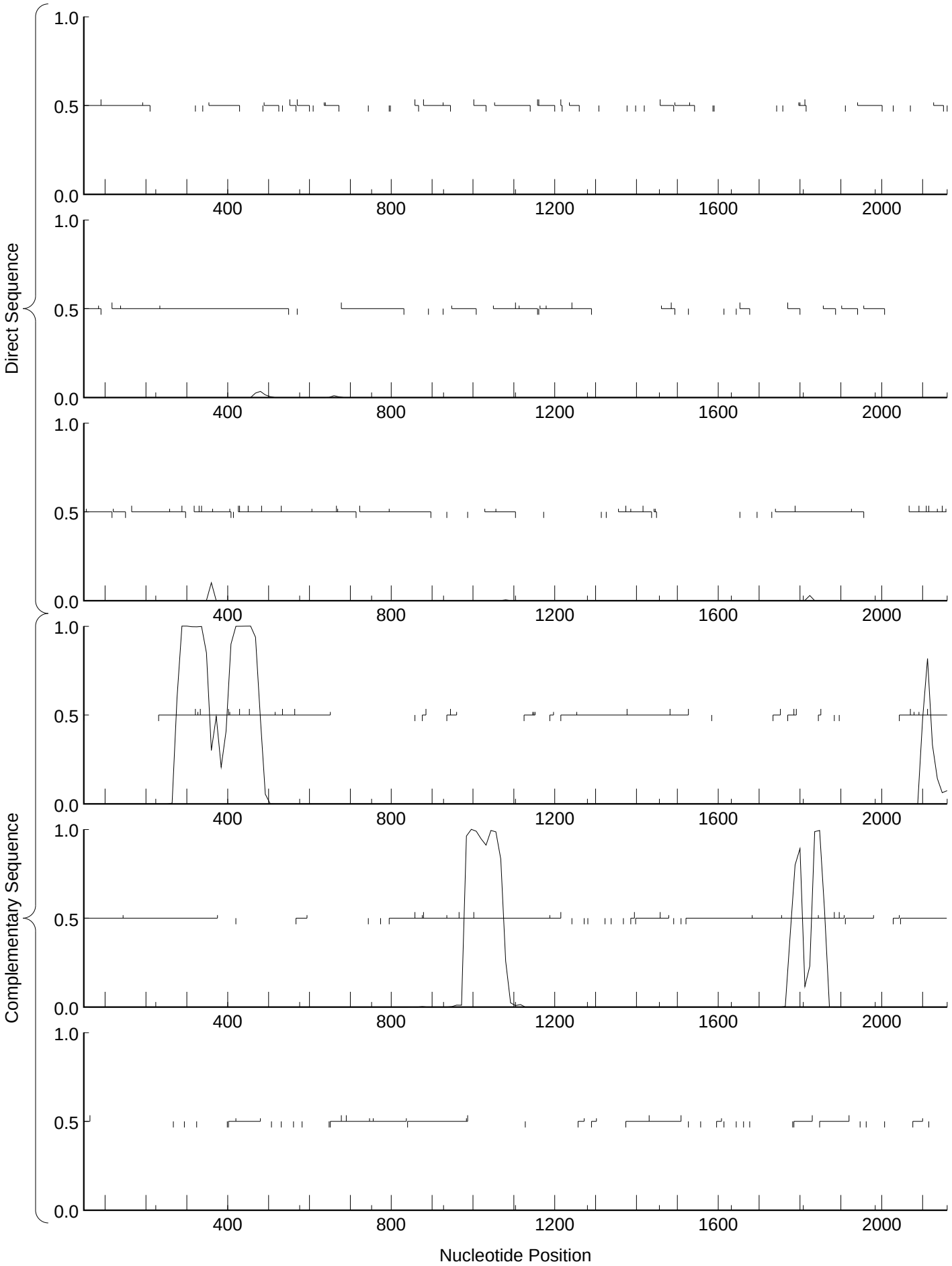
Order      :  4


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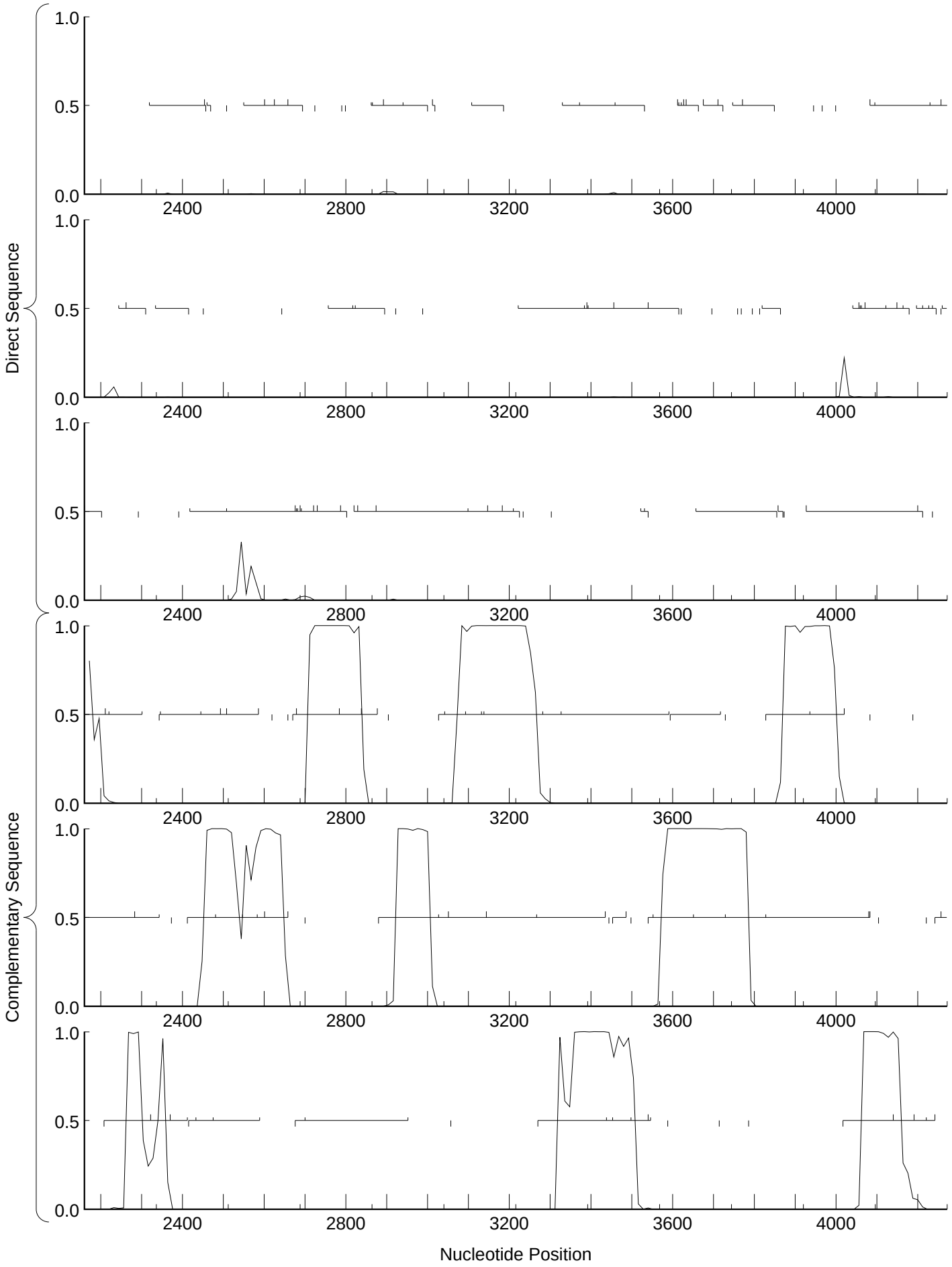


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

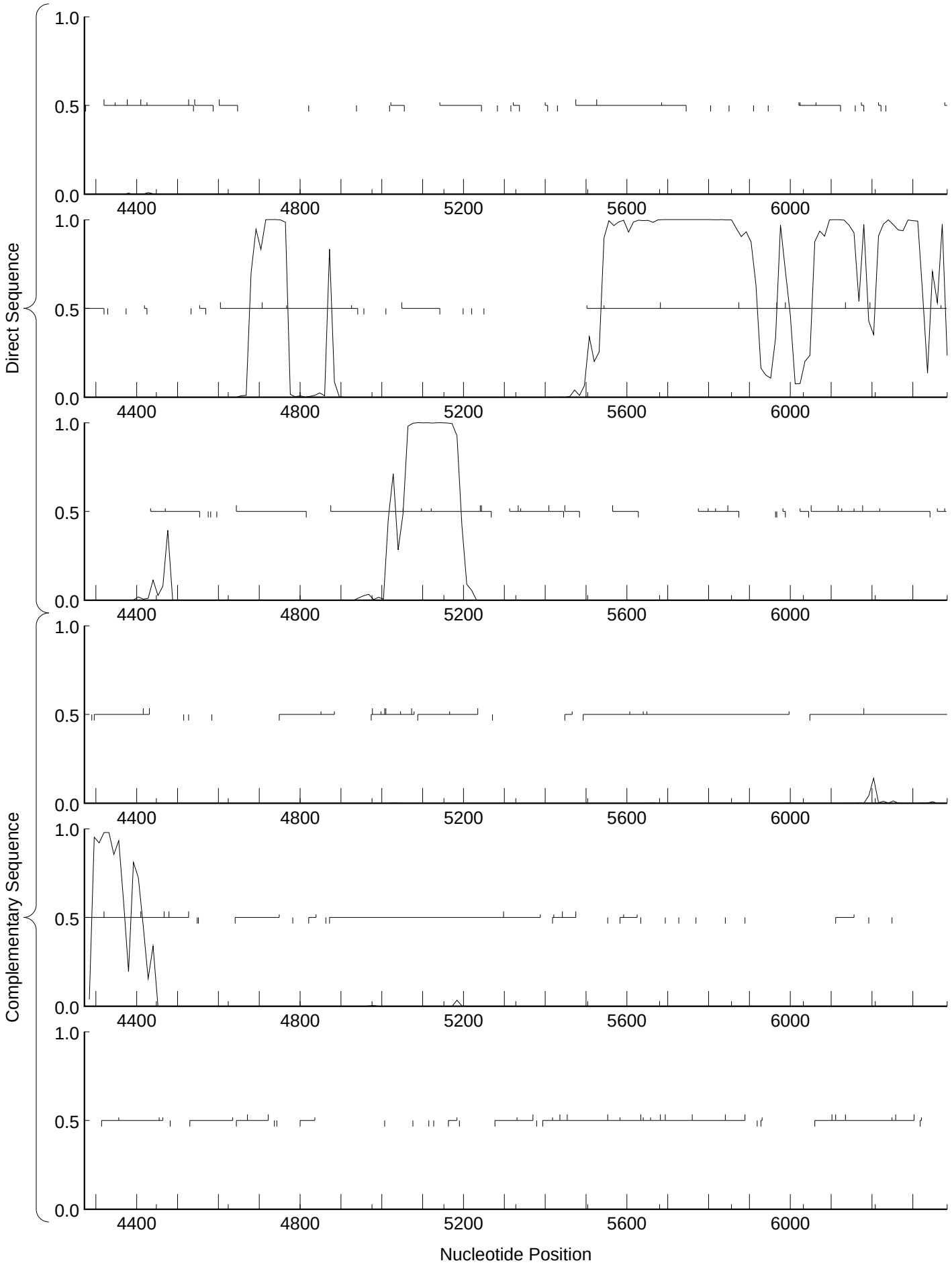
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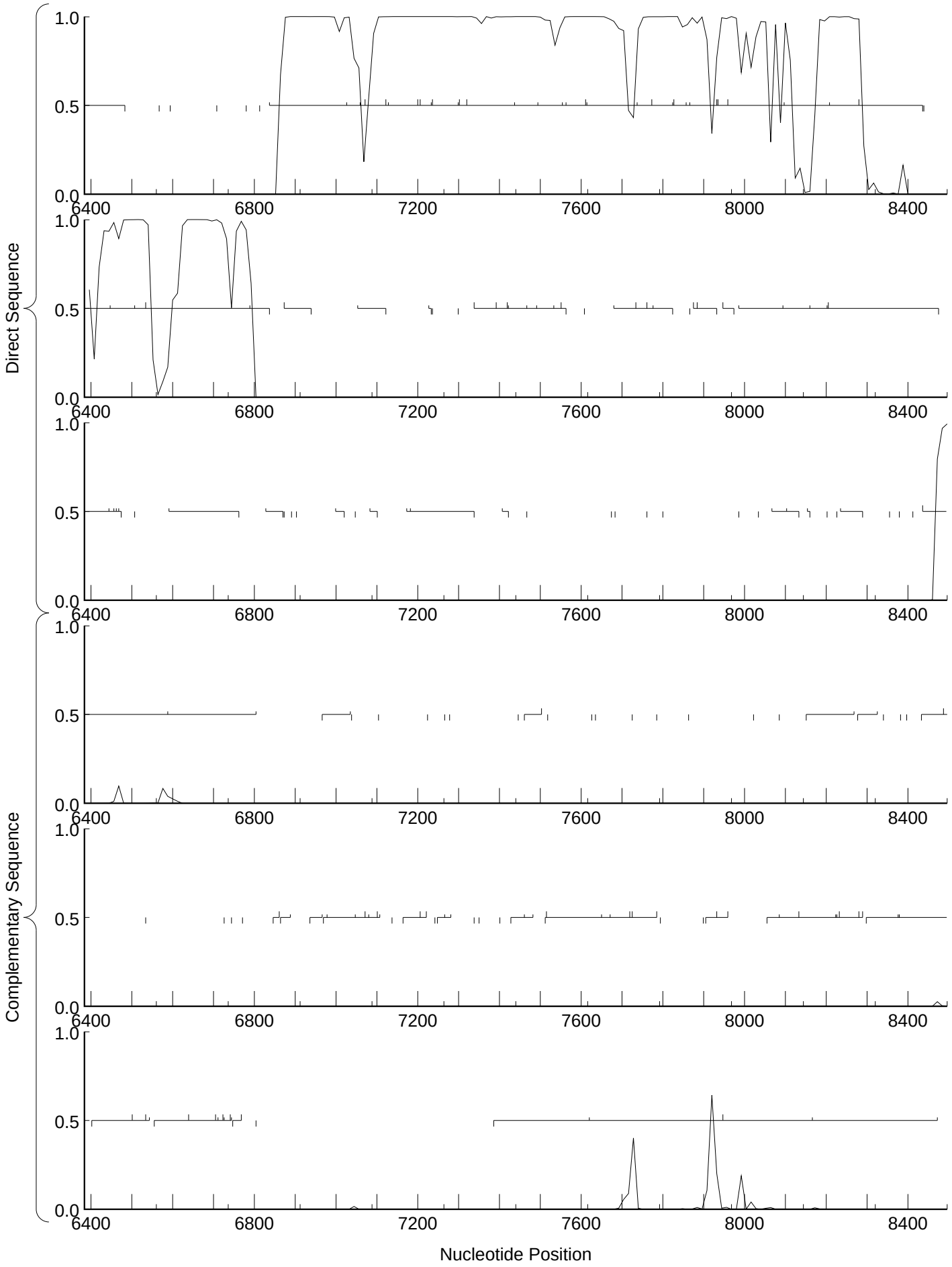
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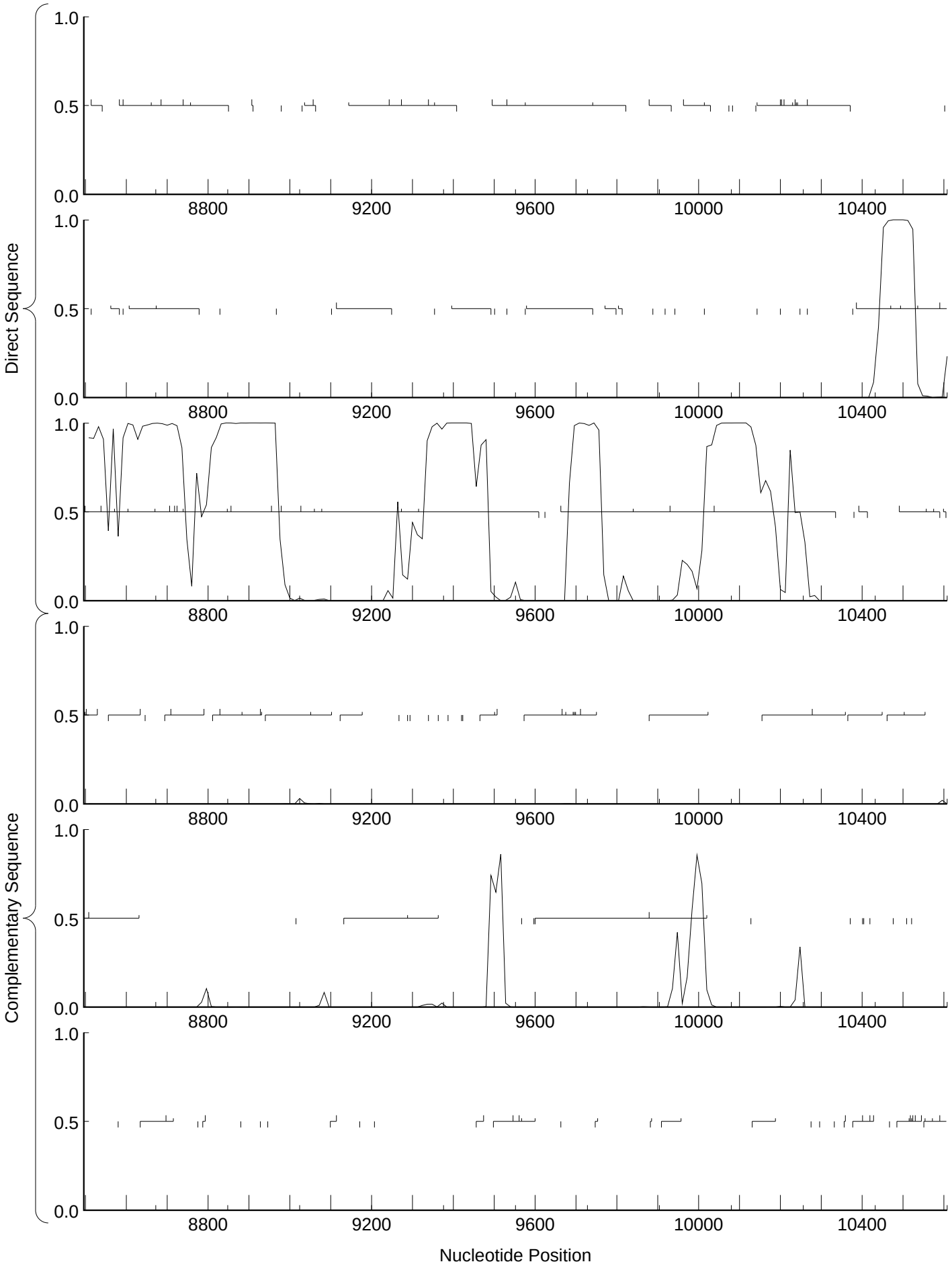
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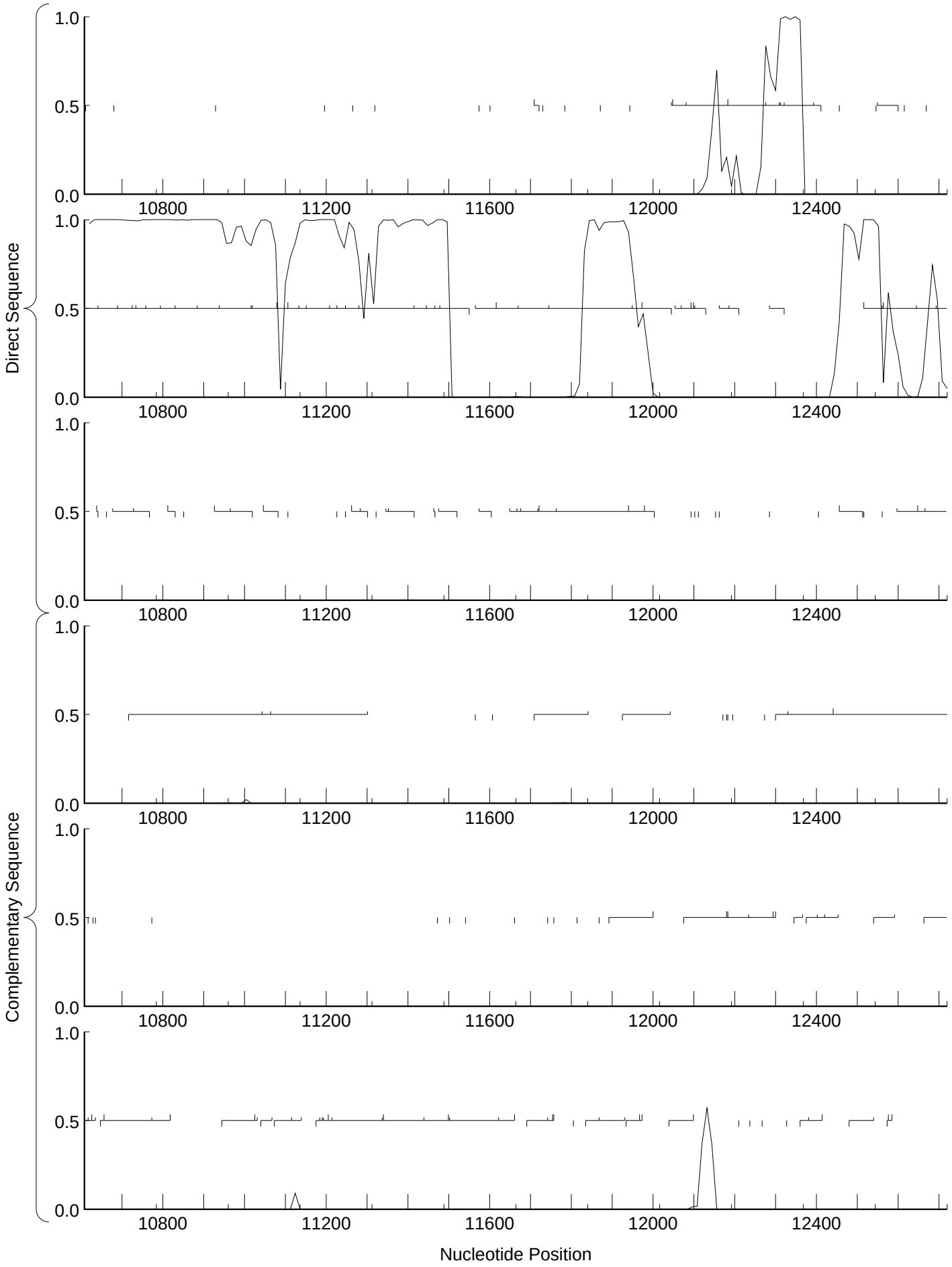
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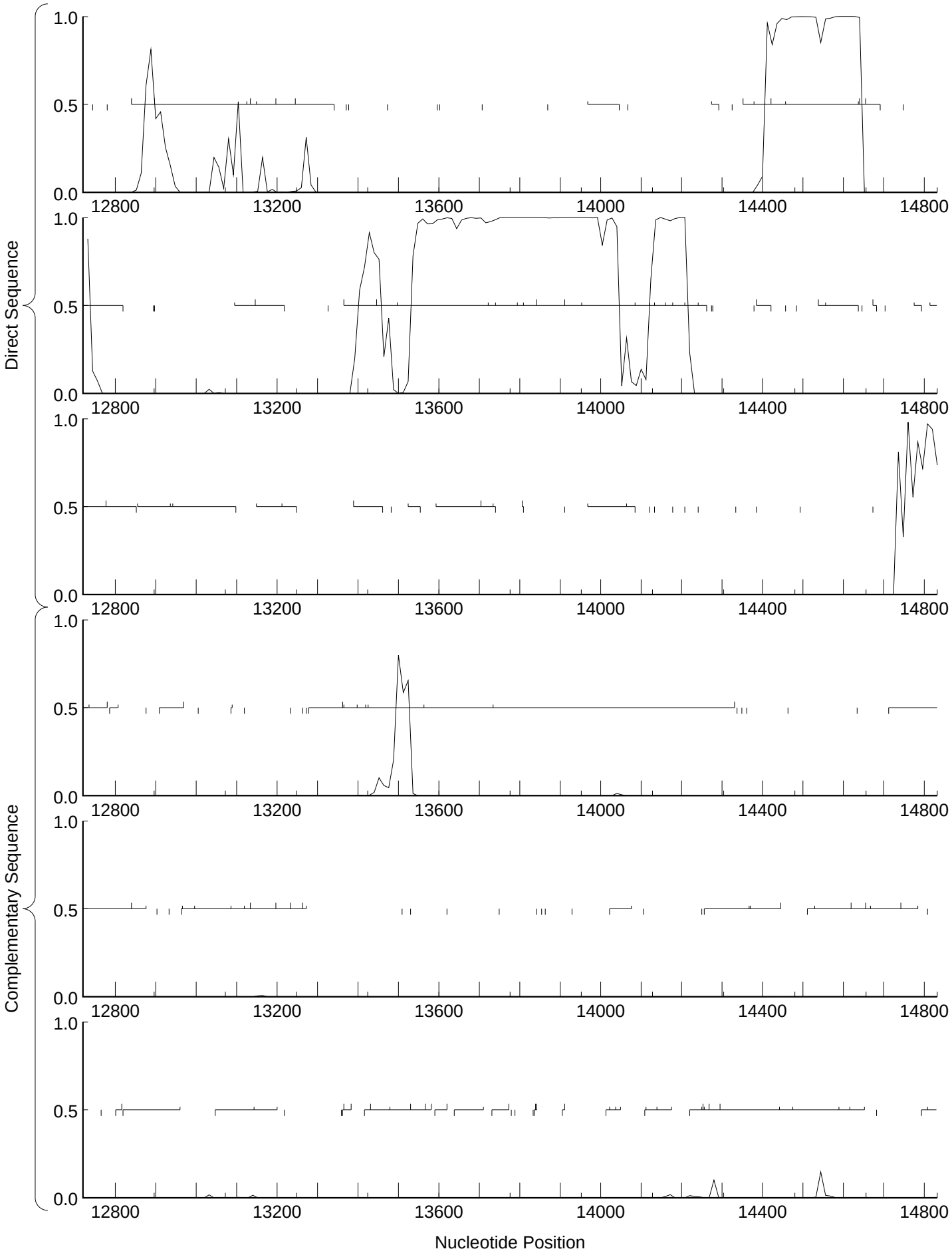
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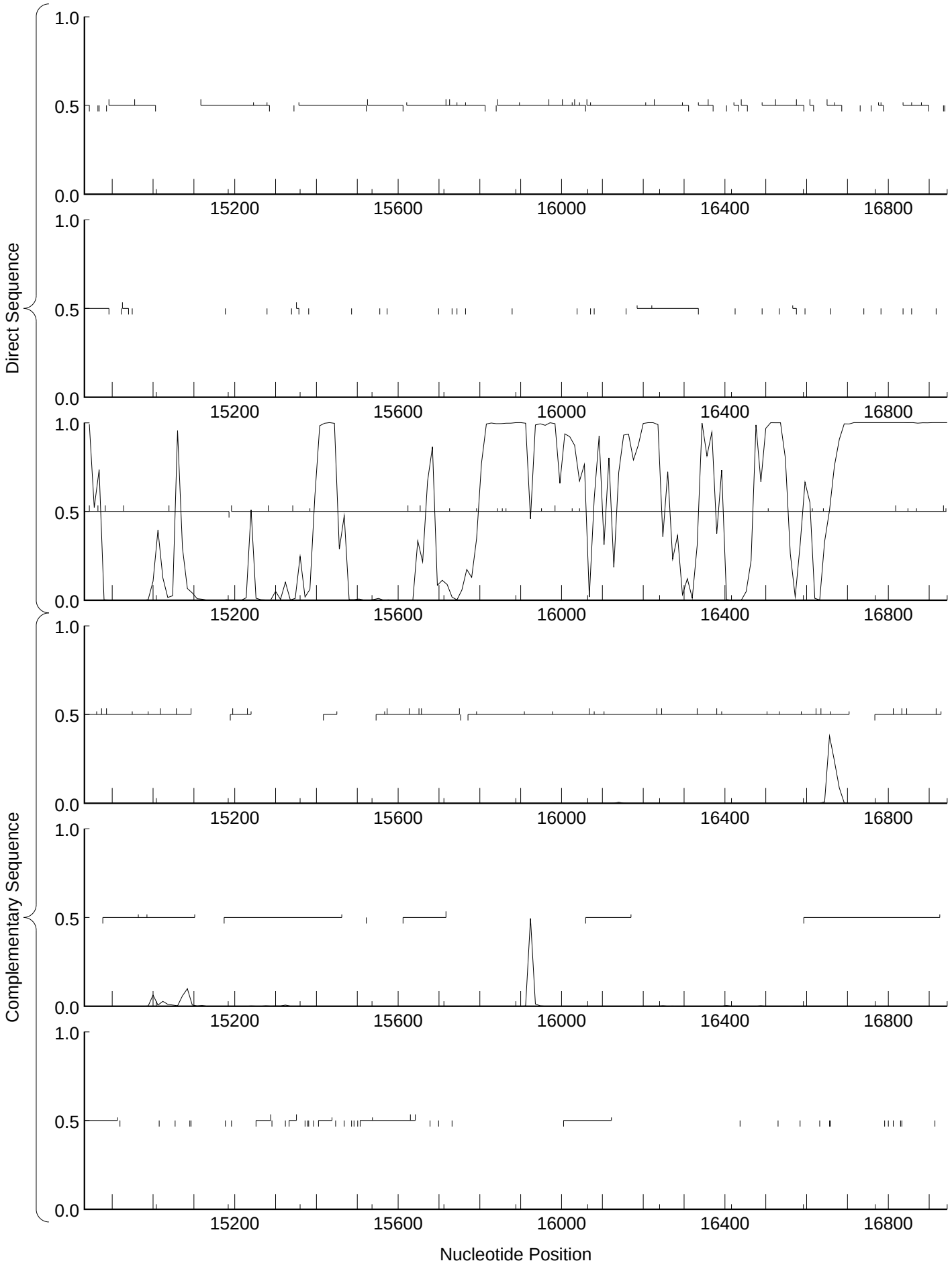
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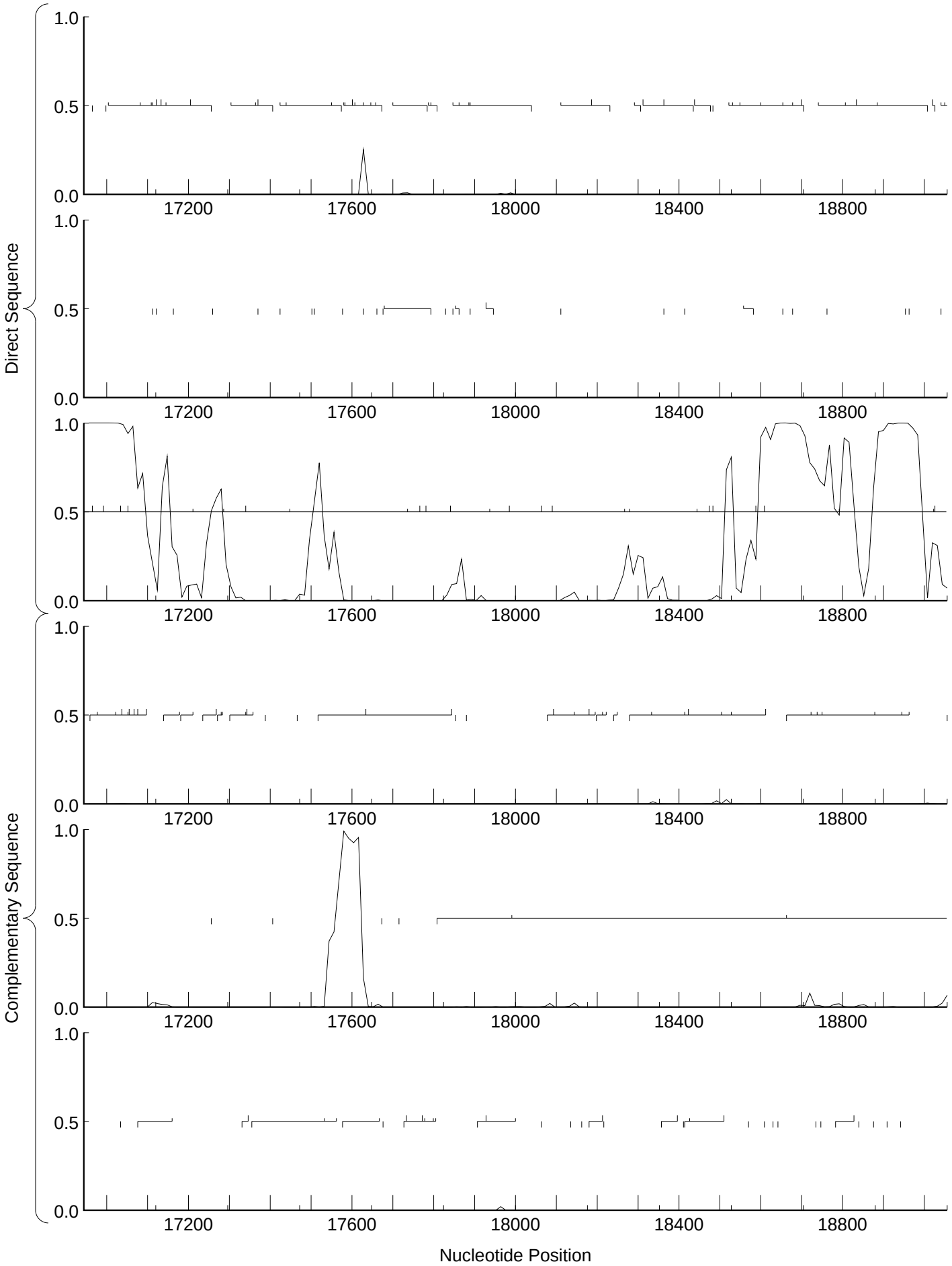
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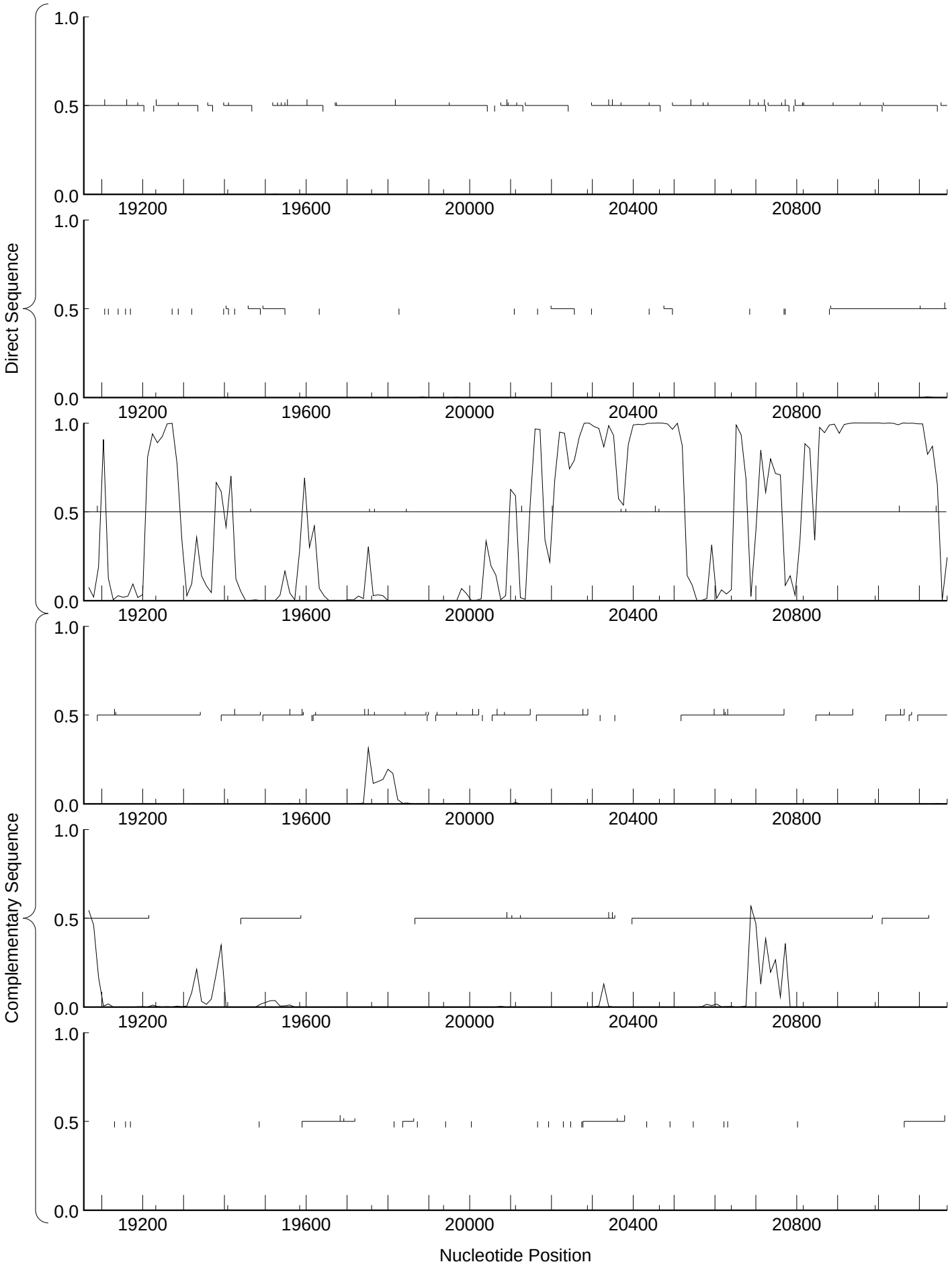
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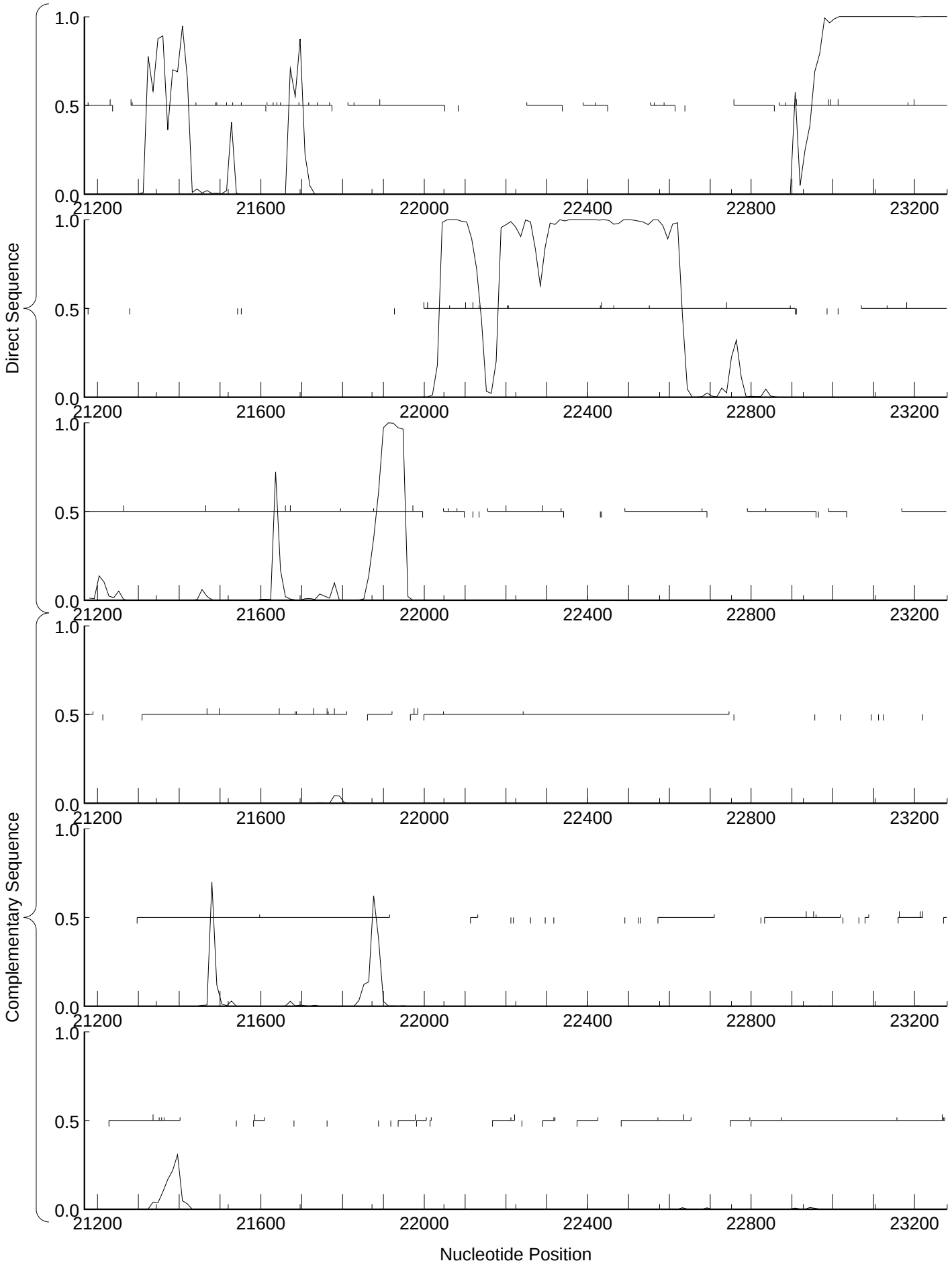
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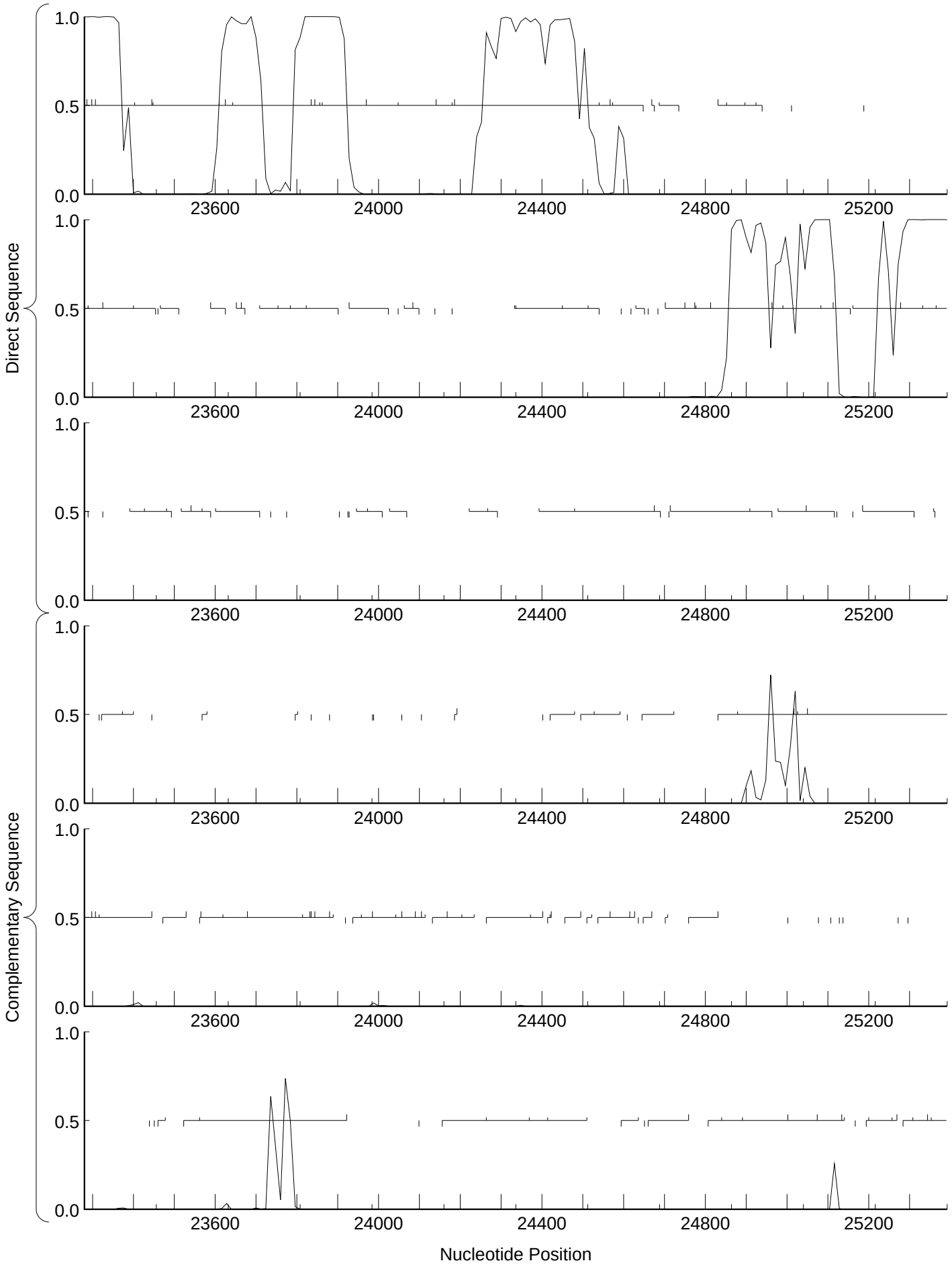
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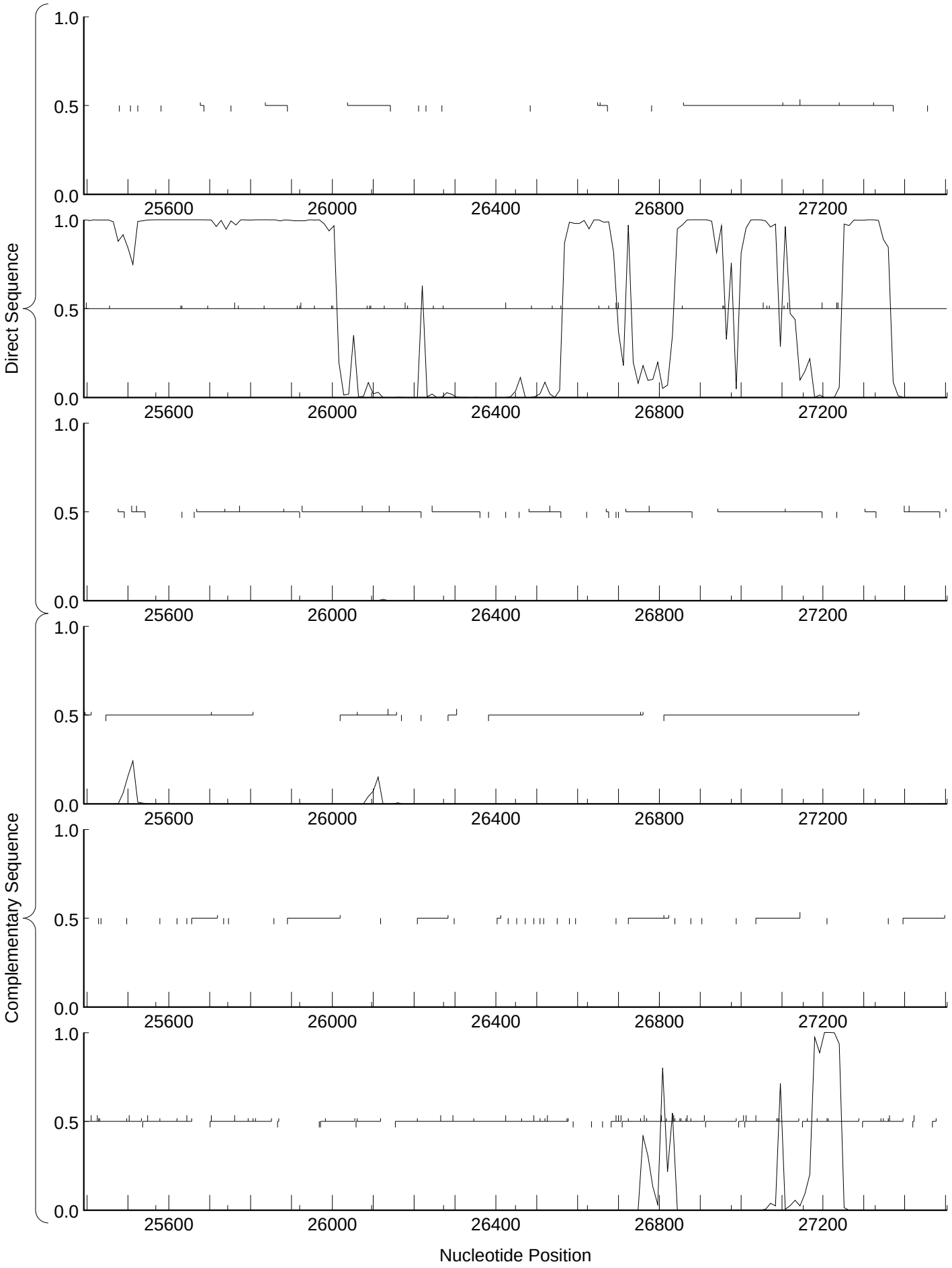
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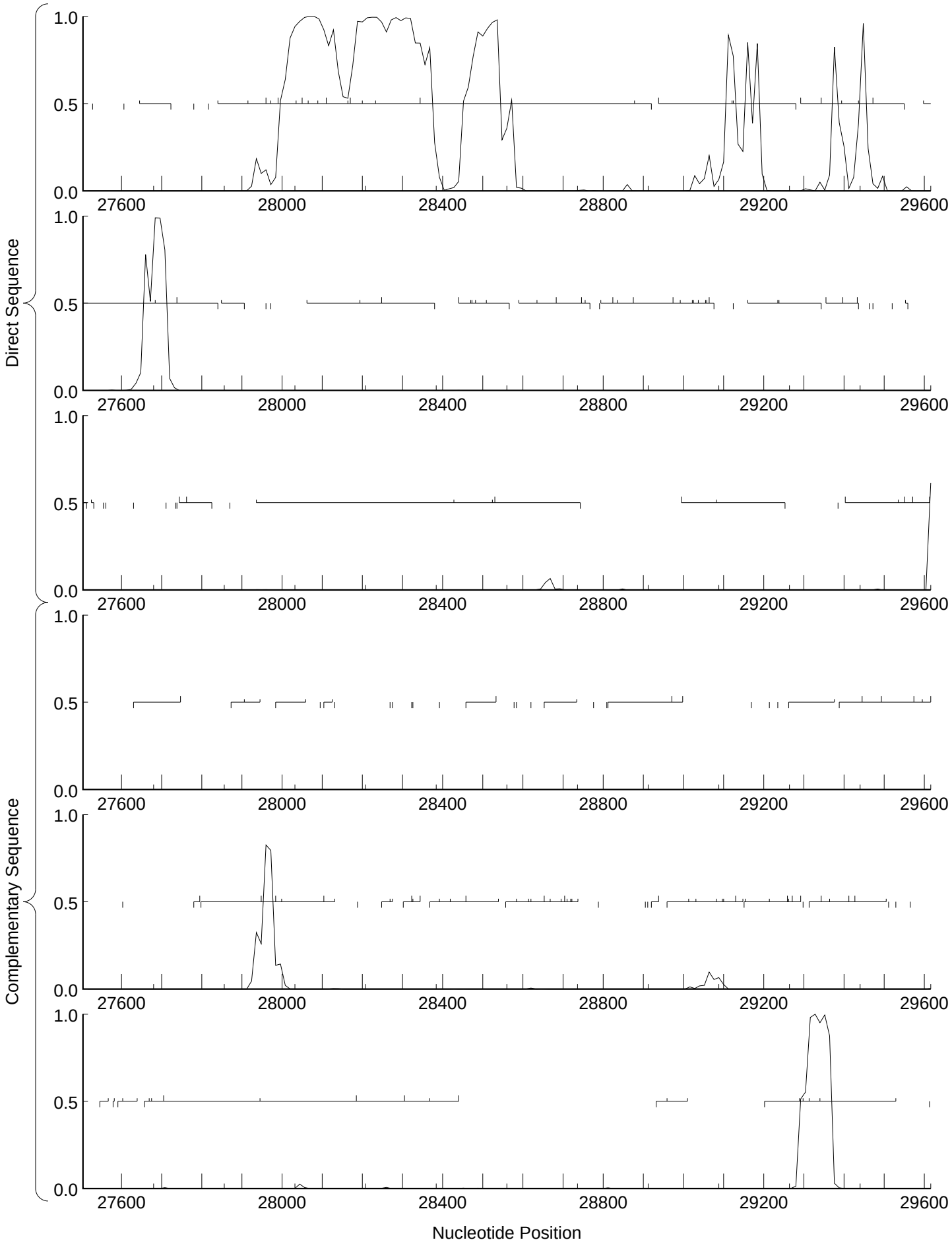
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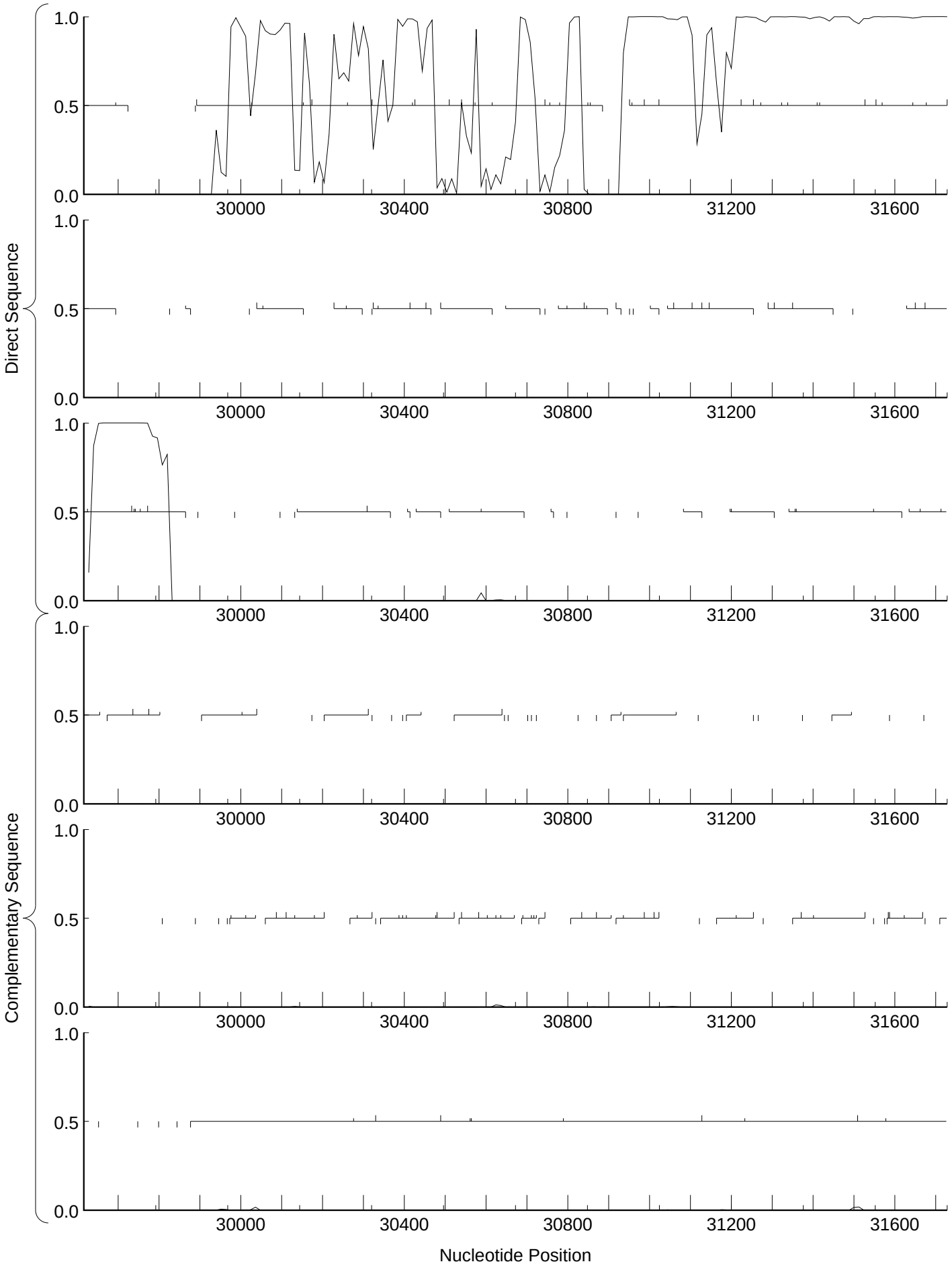
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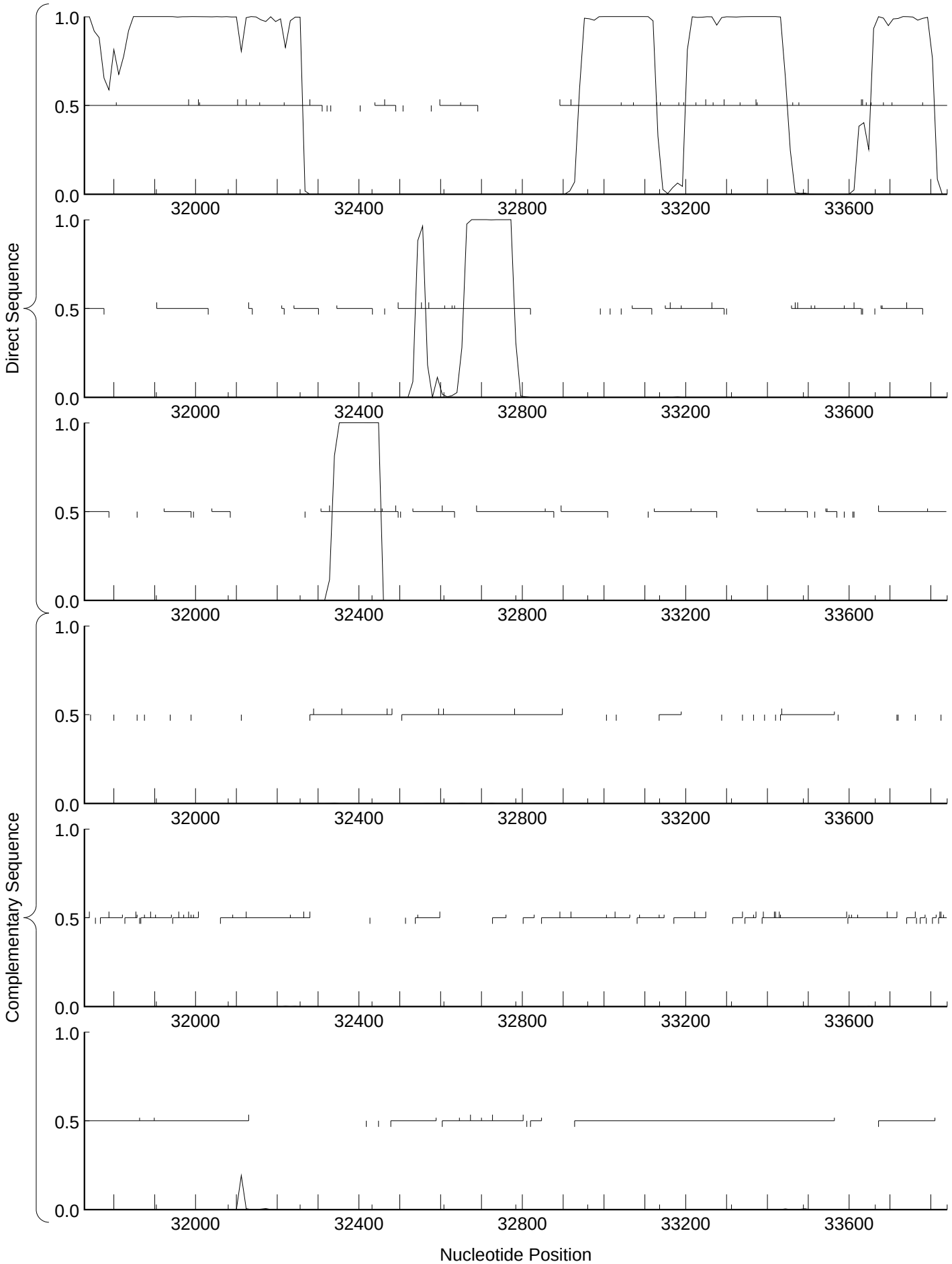


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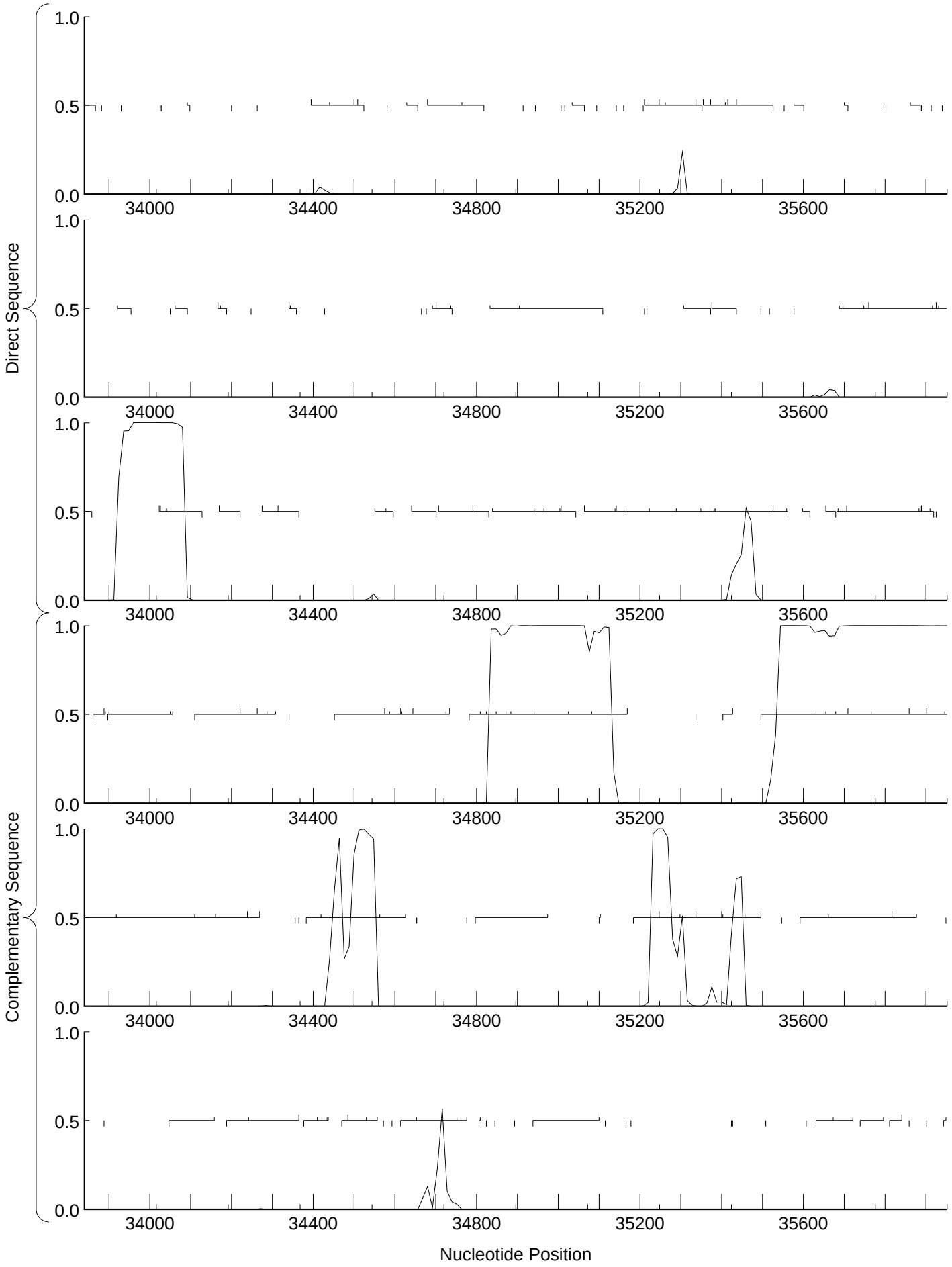


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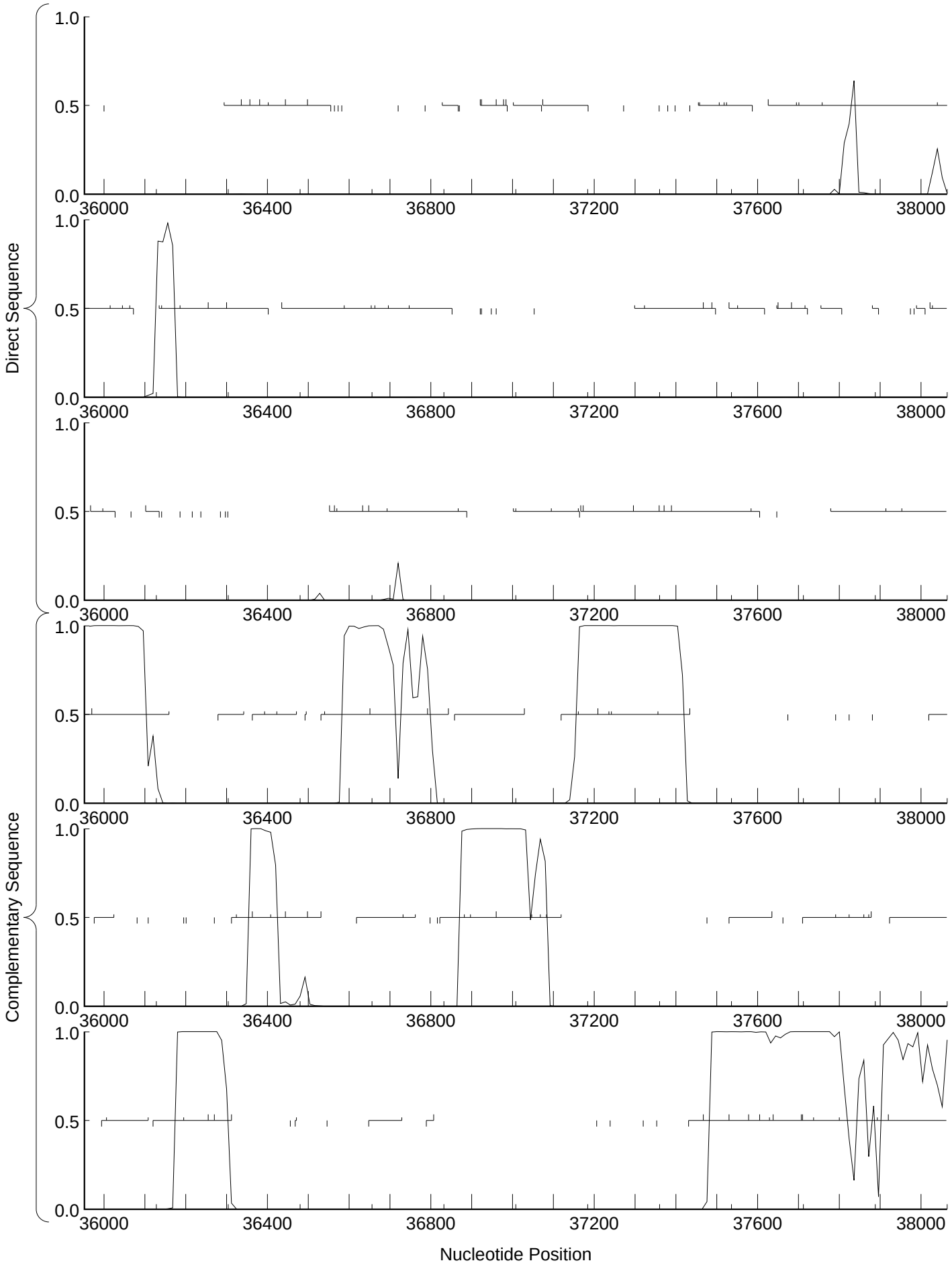



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Complementary Sequence
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0.5
0.0
32000
32400
32800
33200
33600
Nucleotide Position

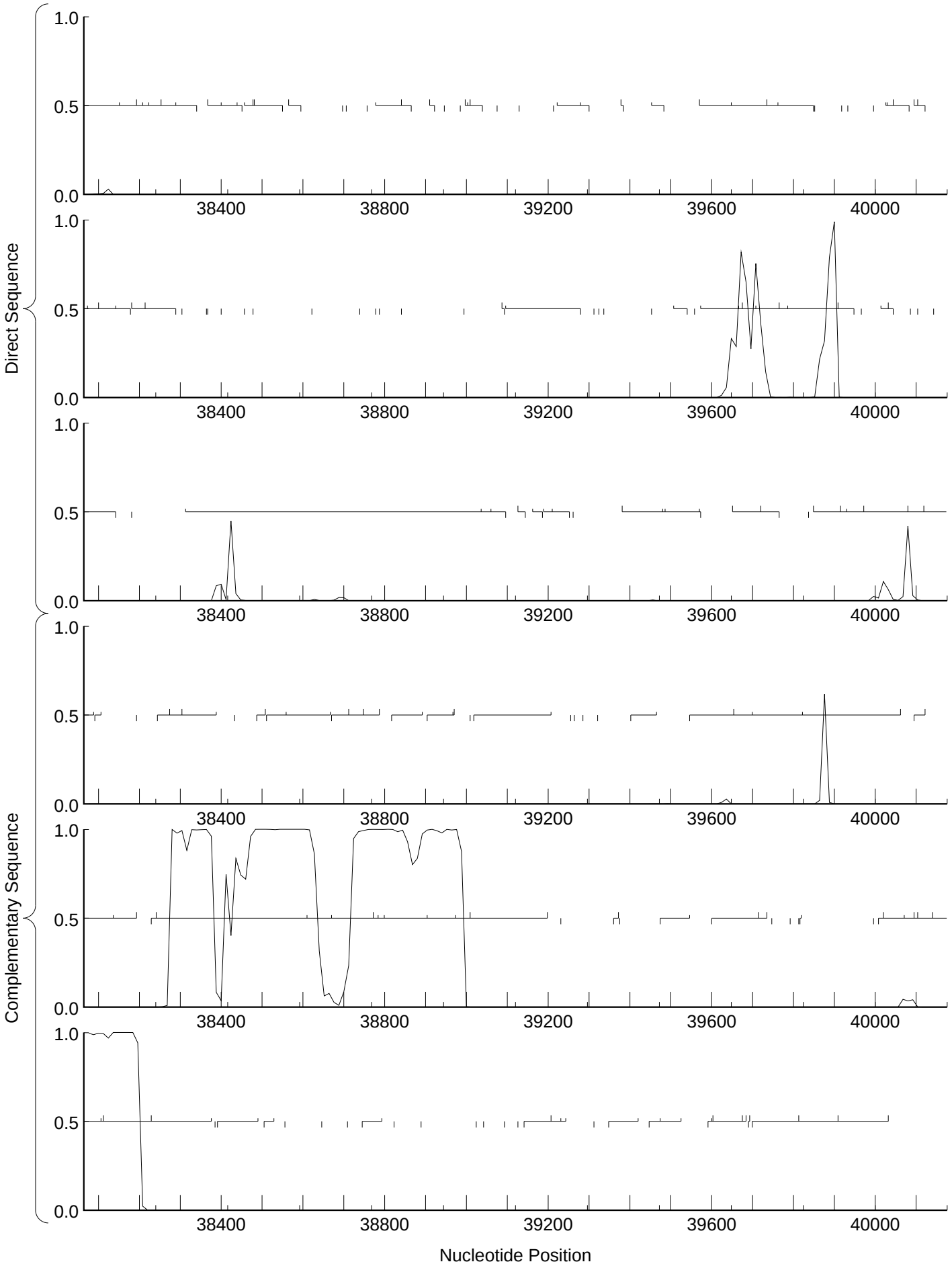
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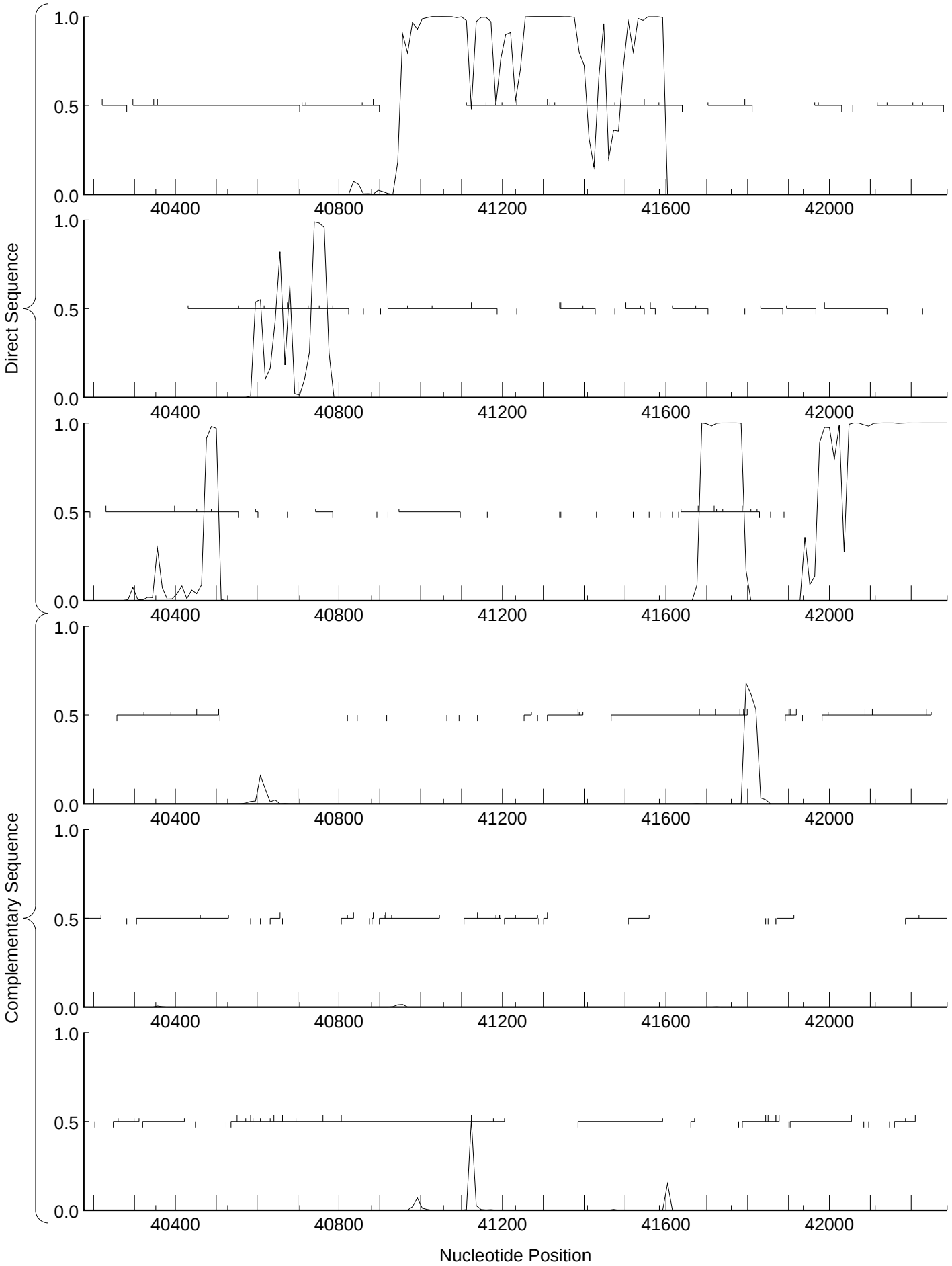
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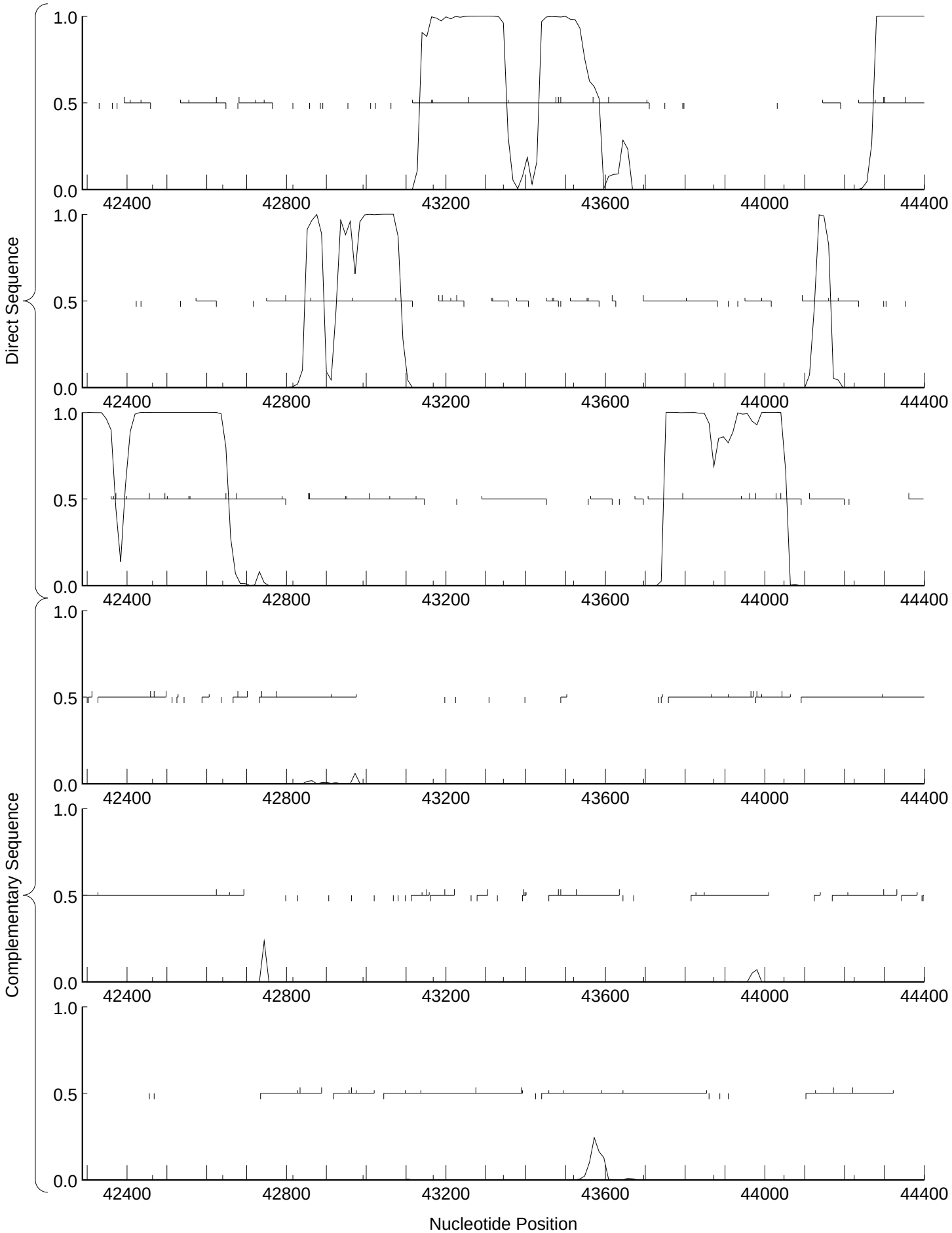
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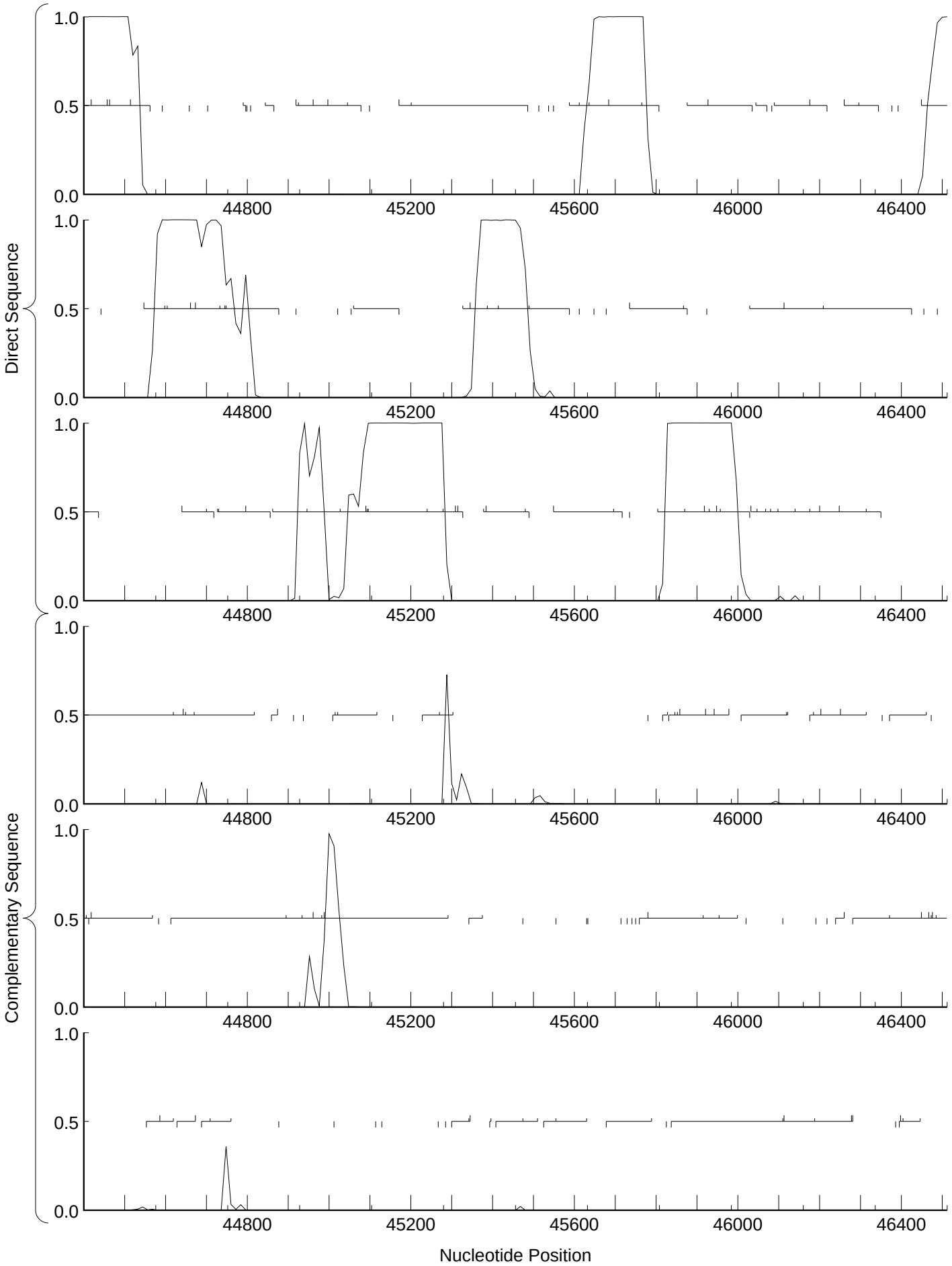
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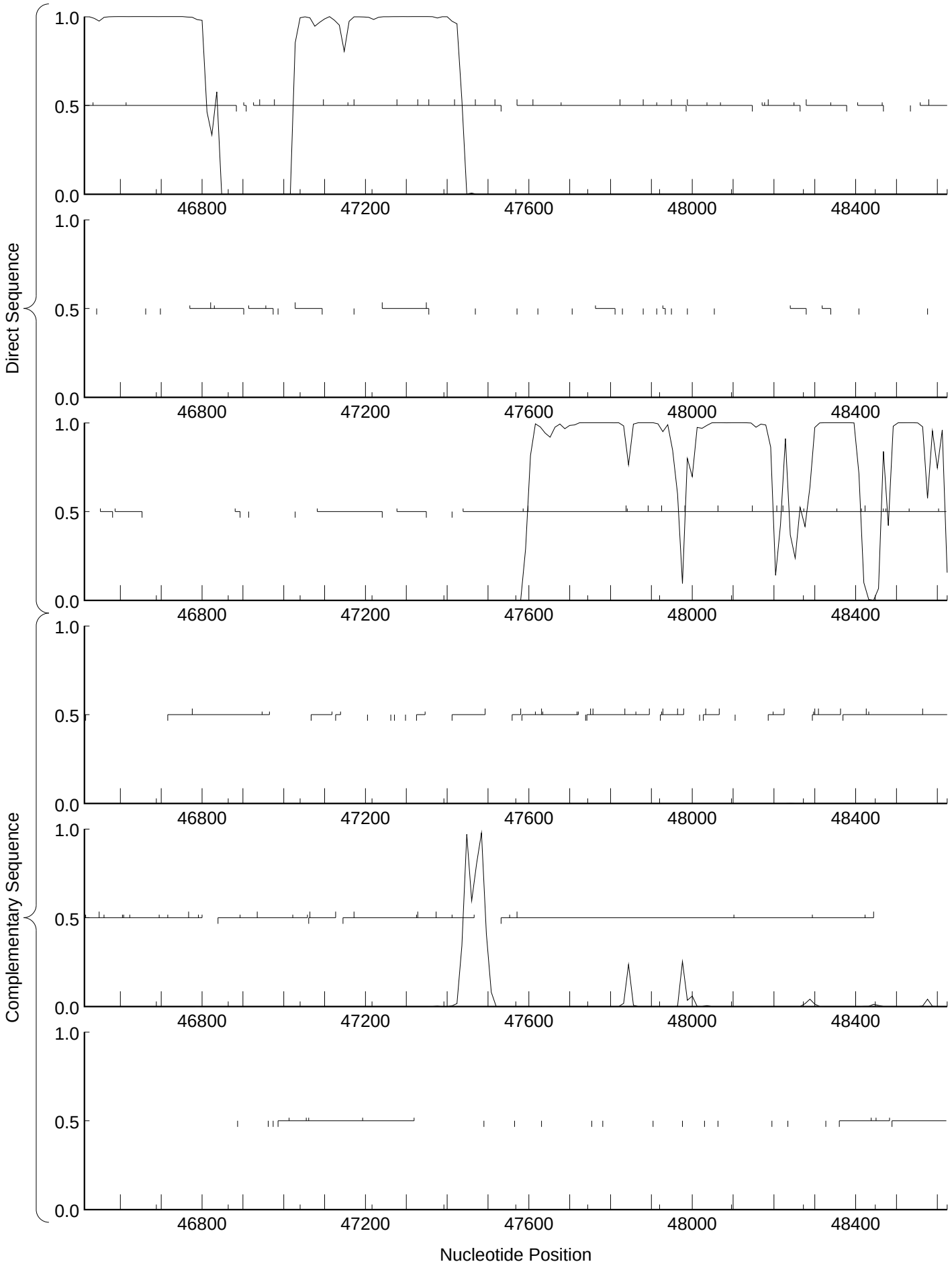
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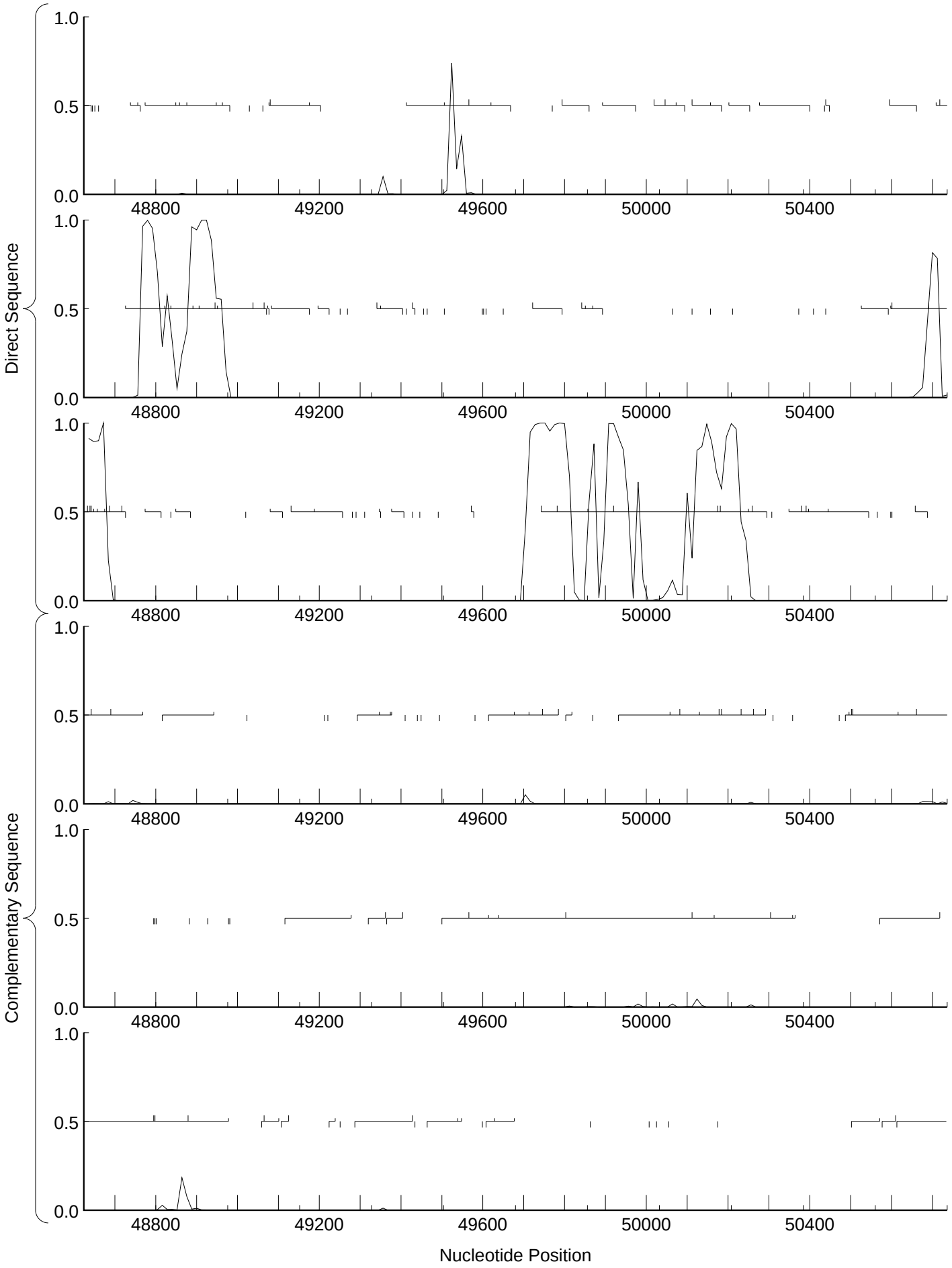
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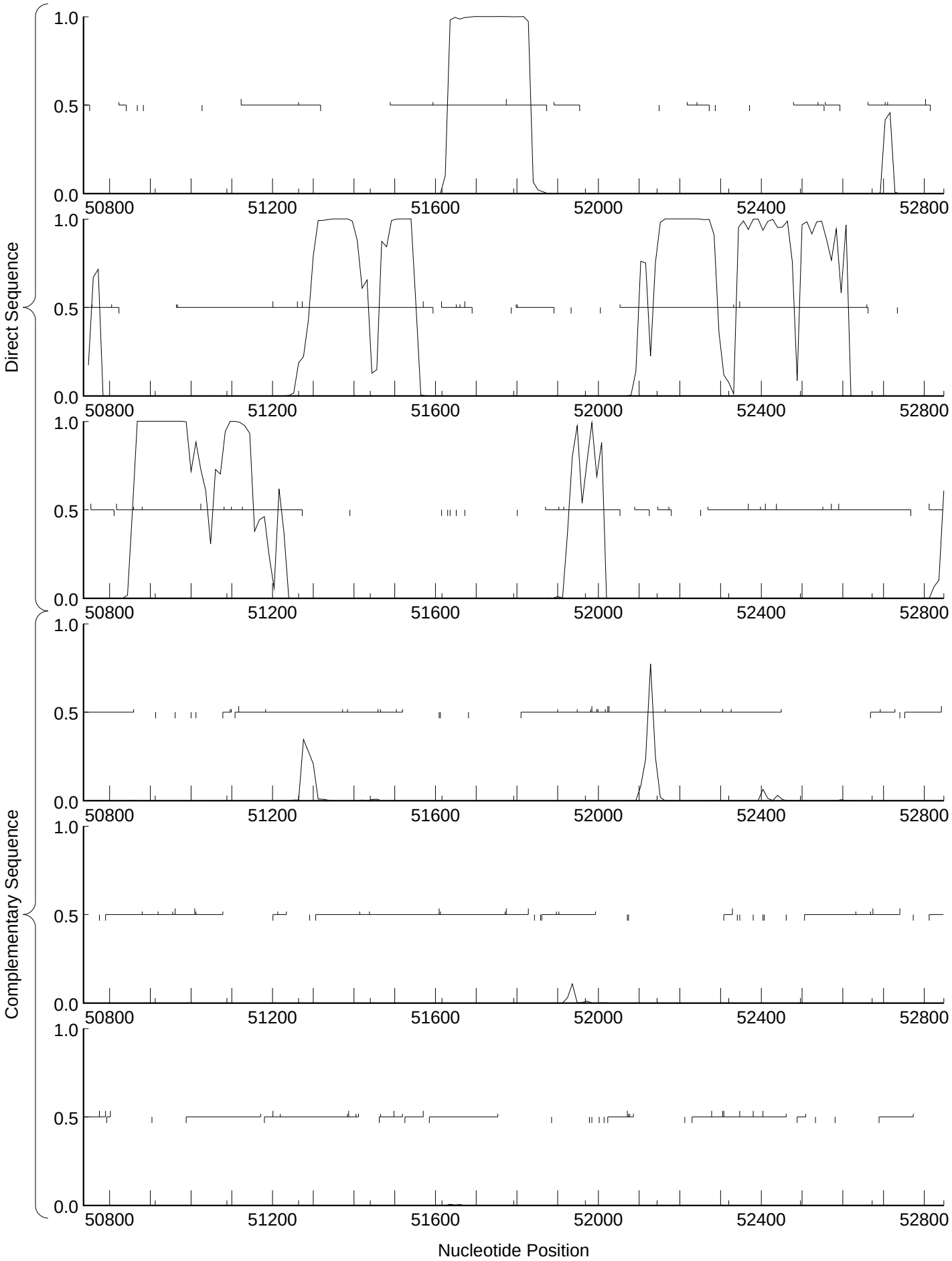
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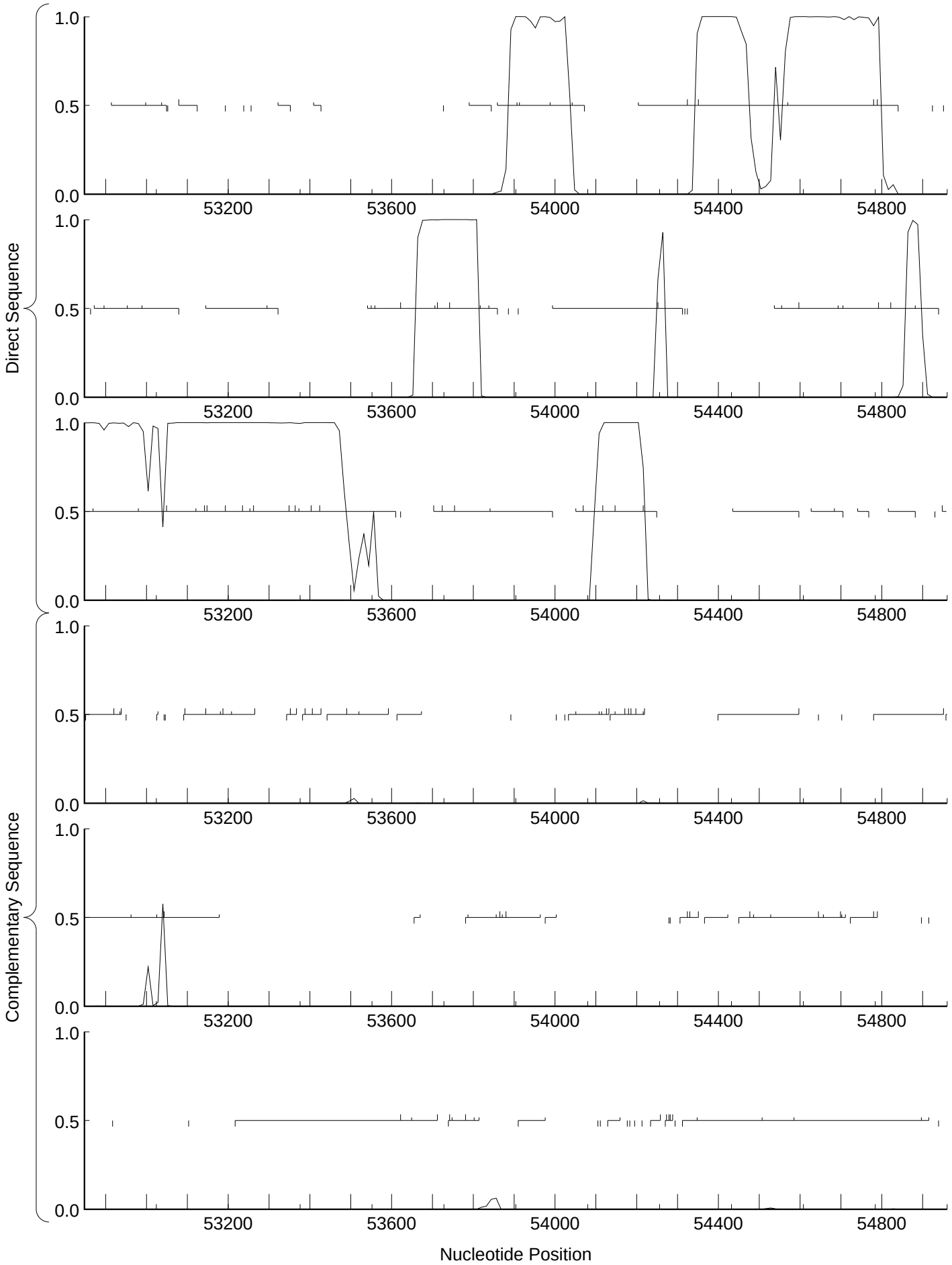
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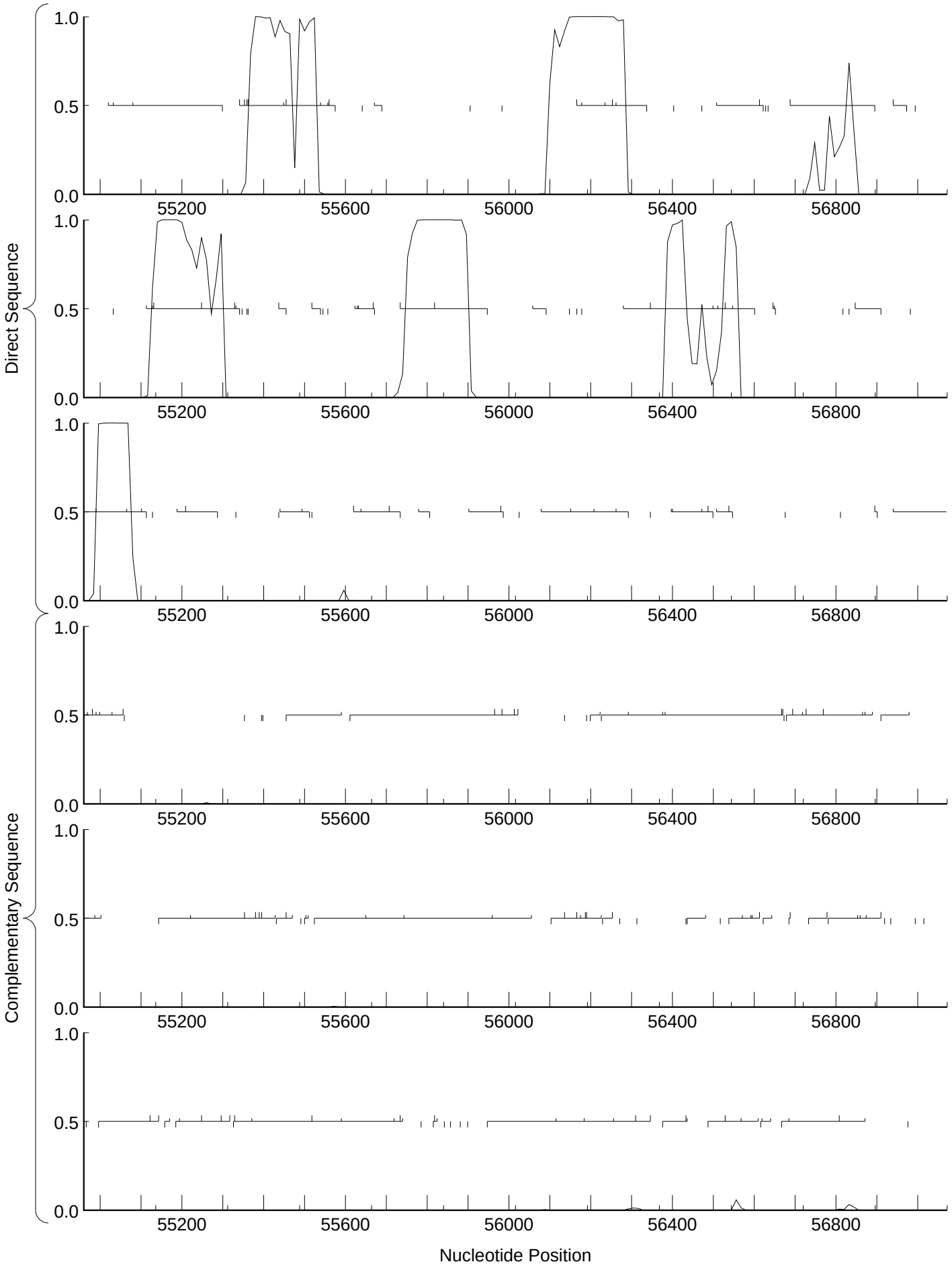
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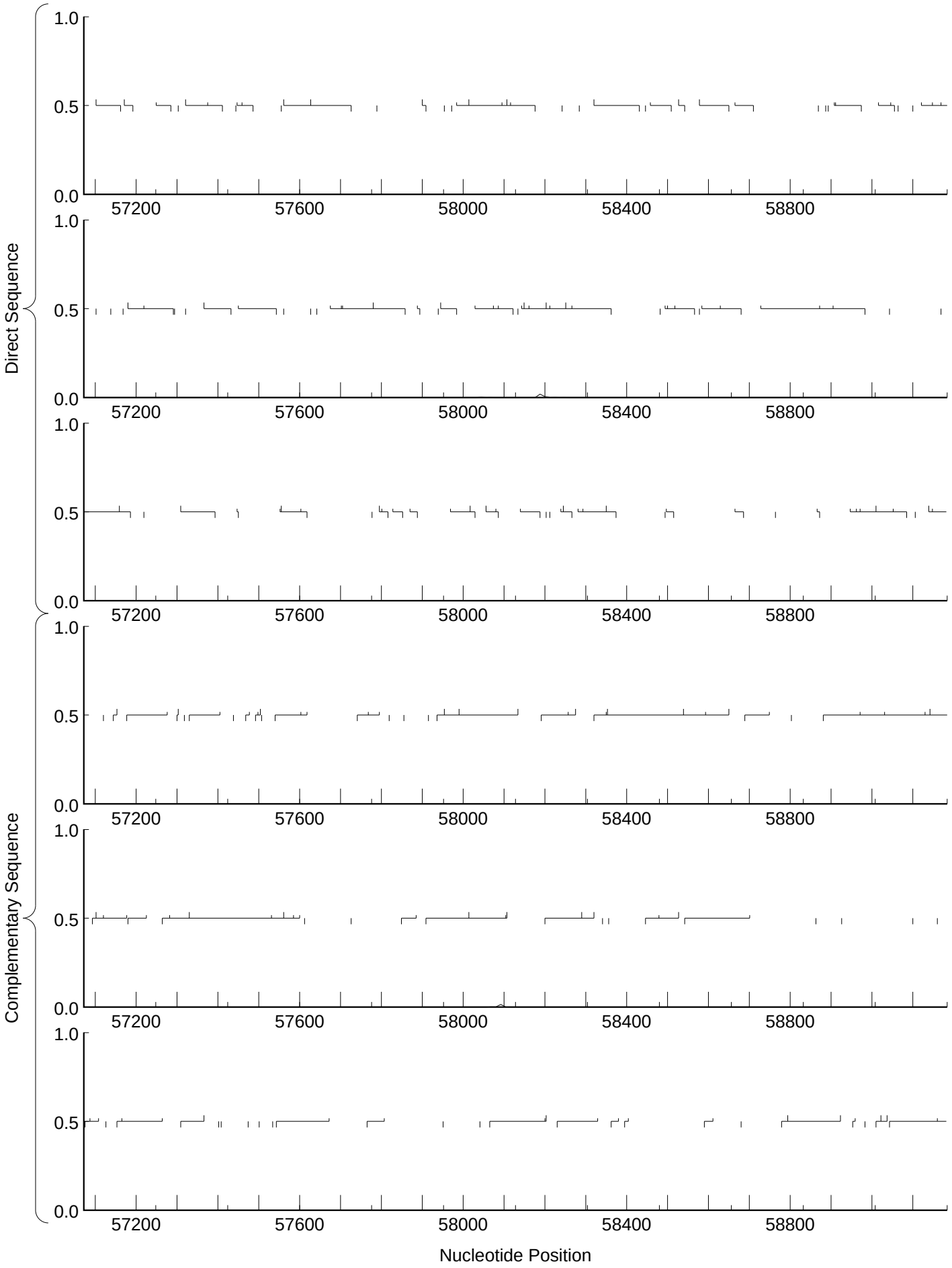
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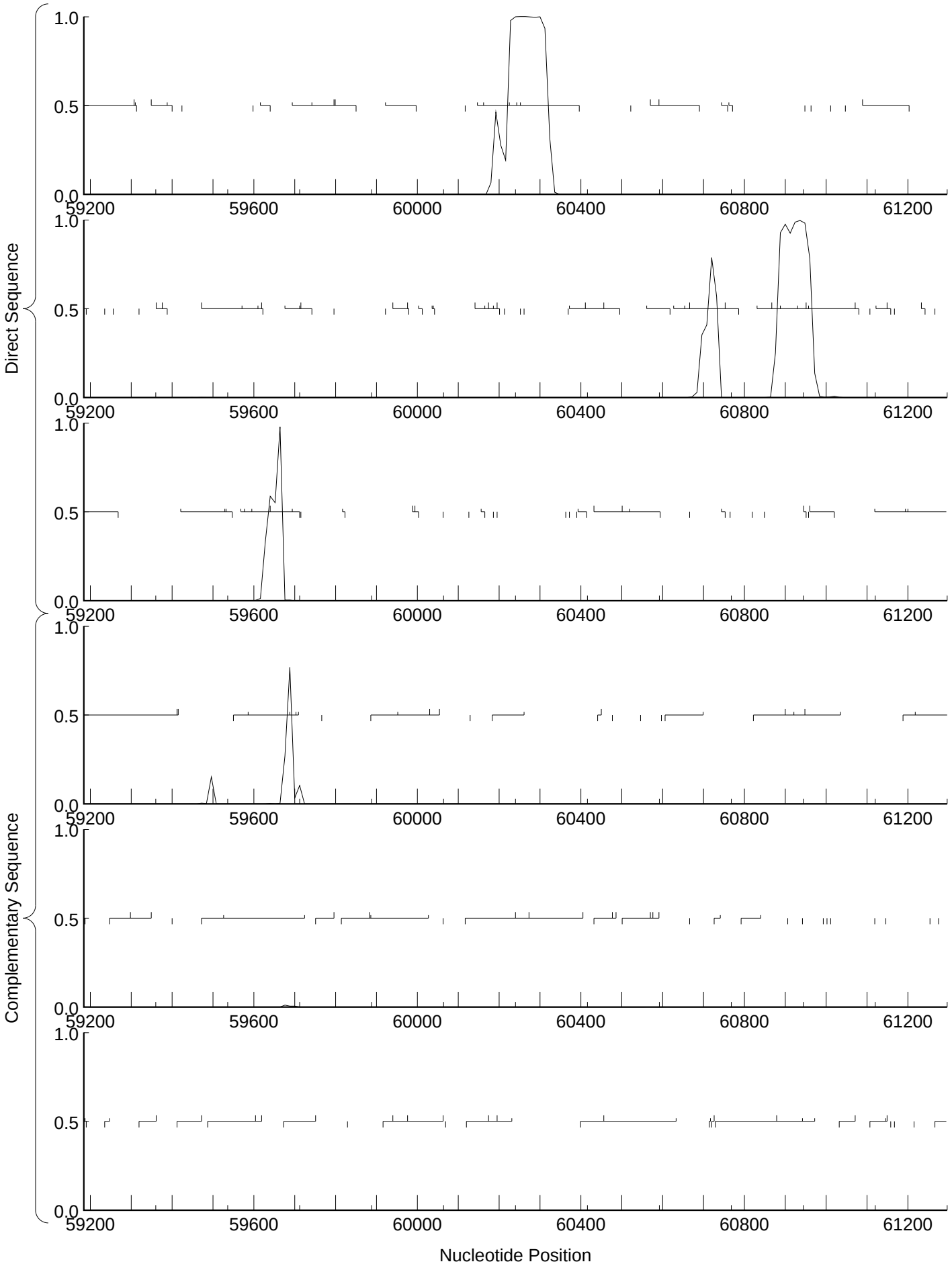
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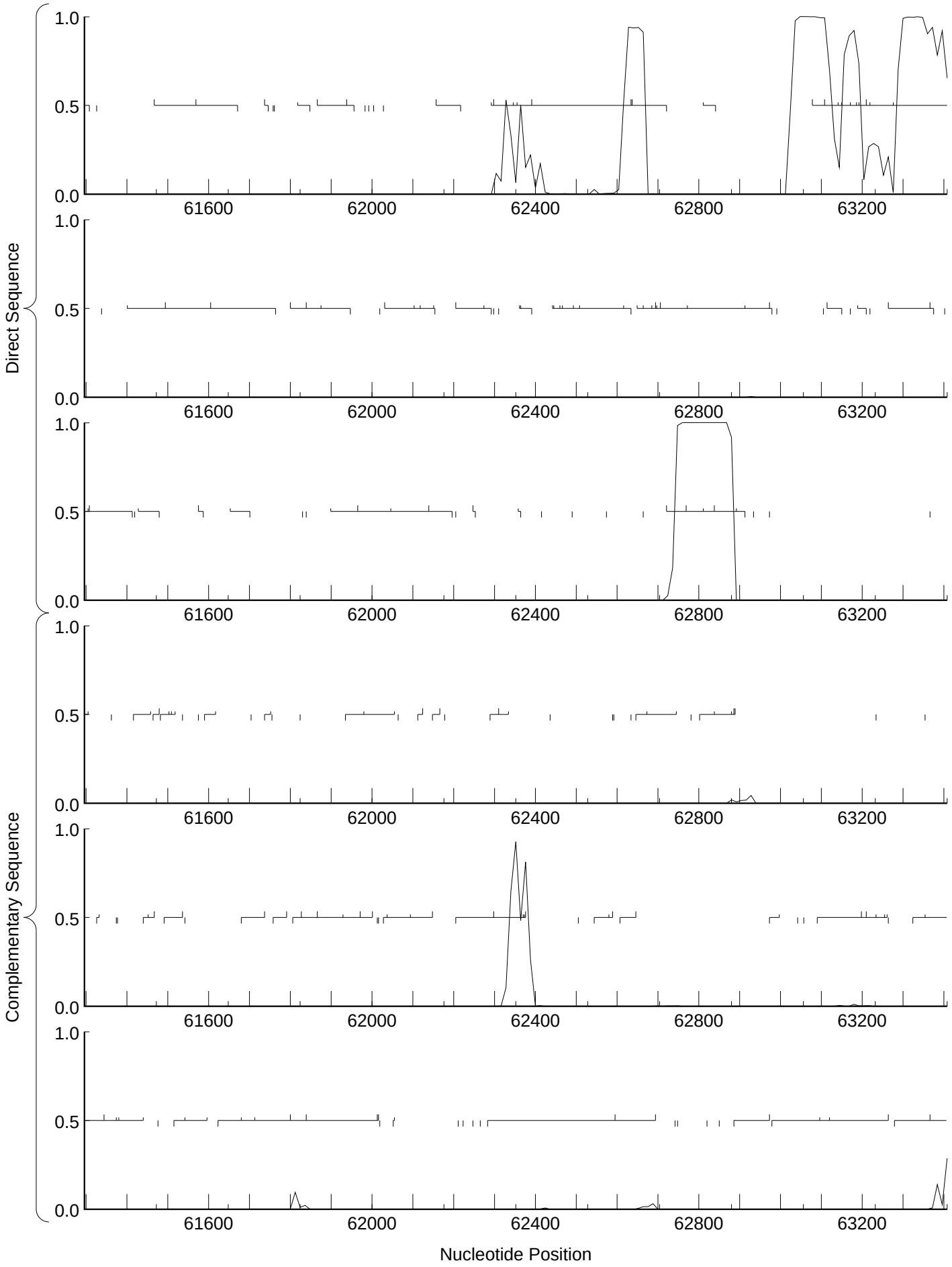
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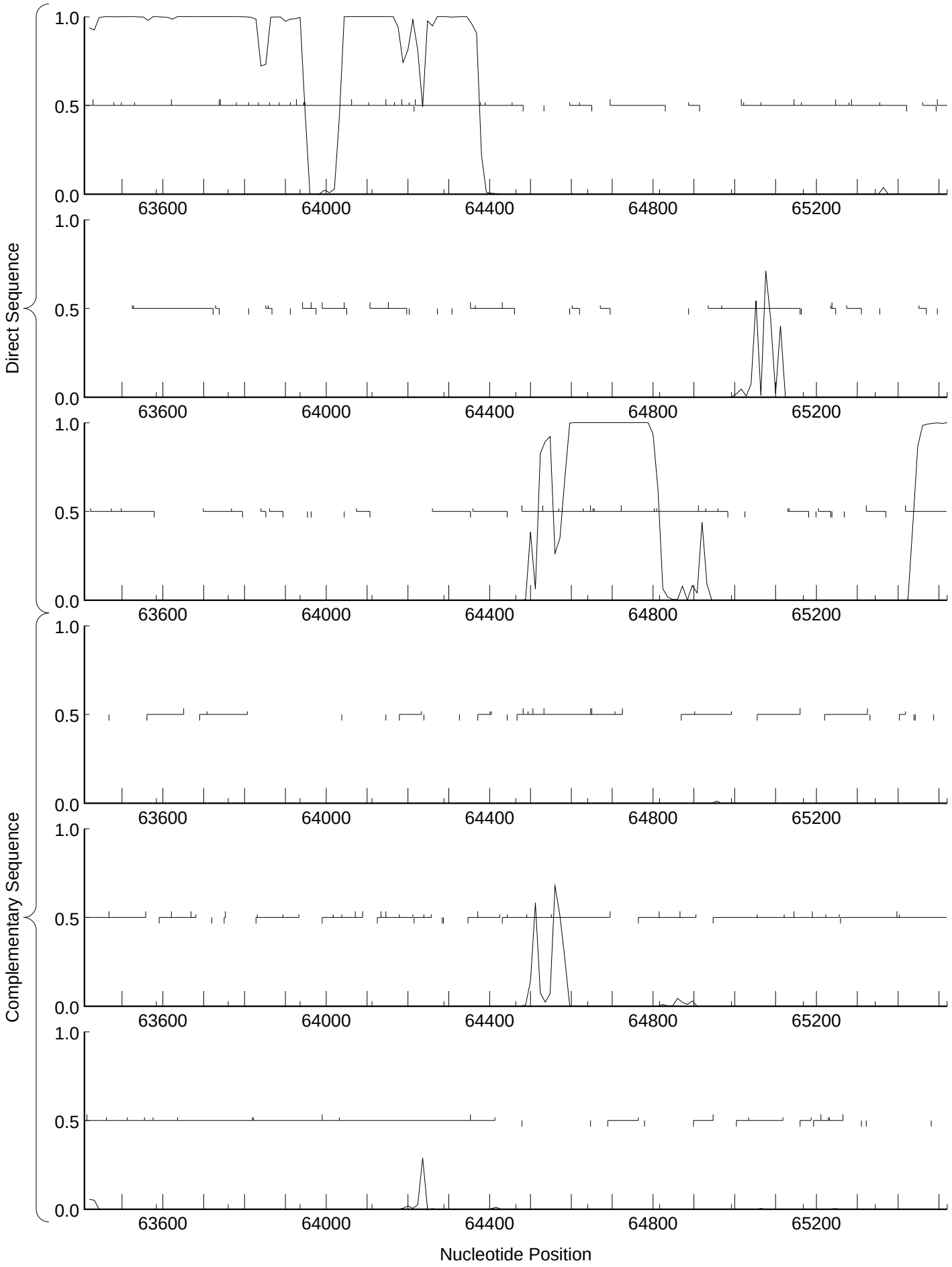
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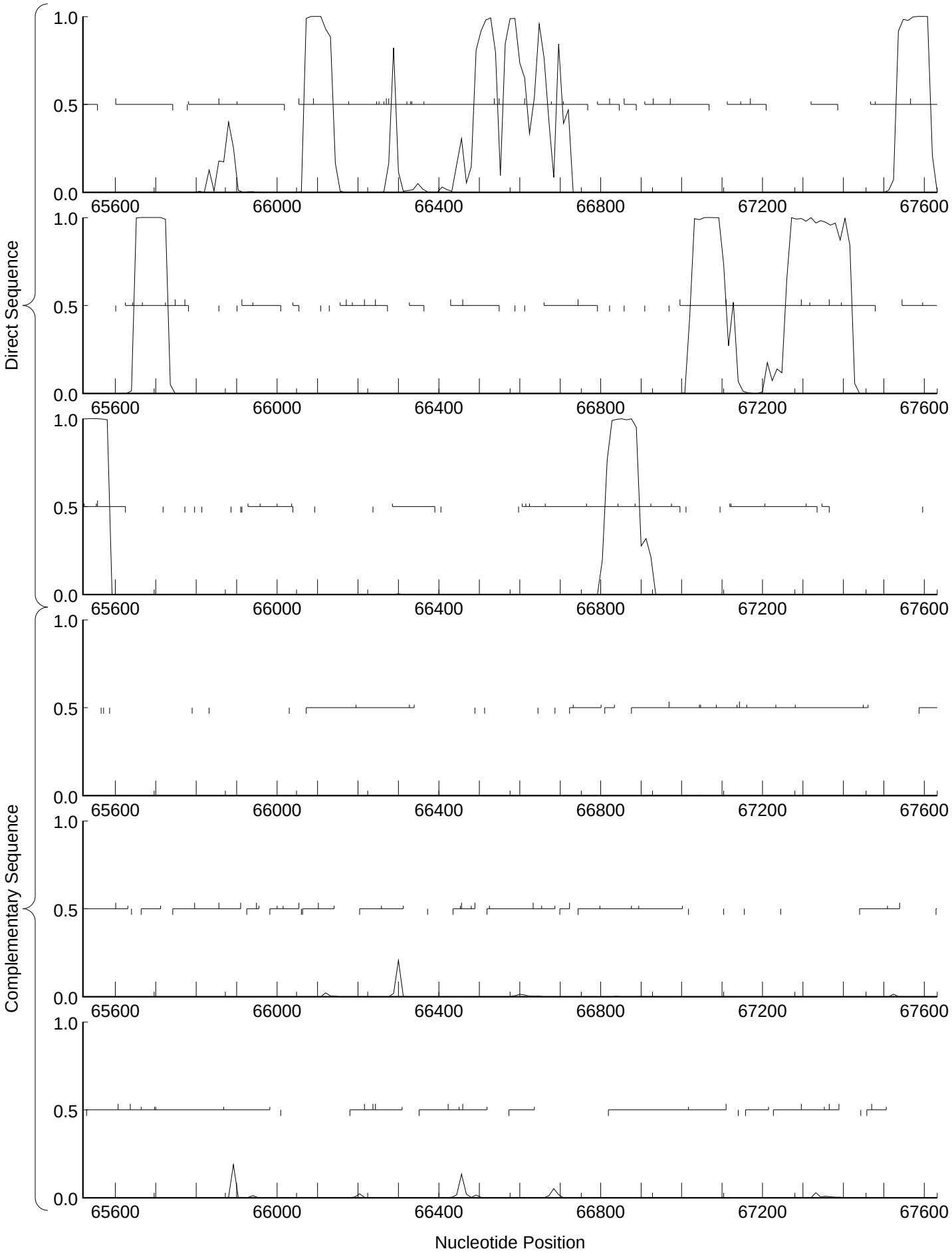
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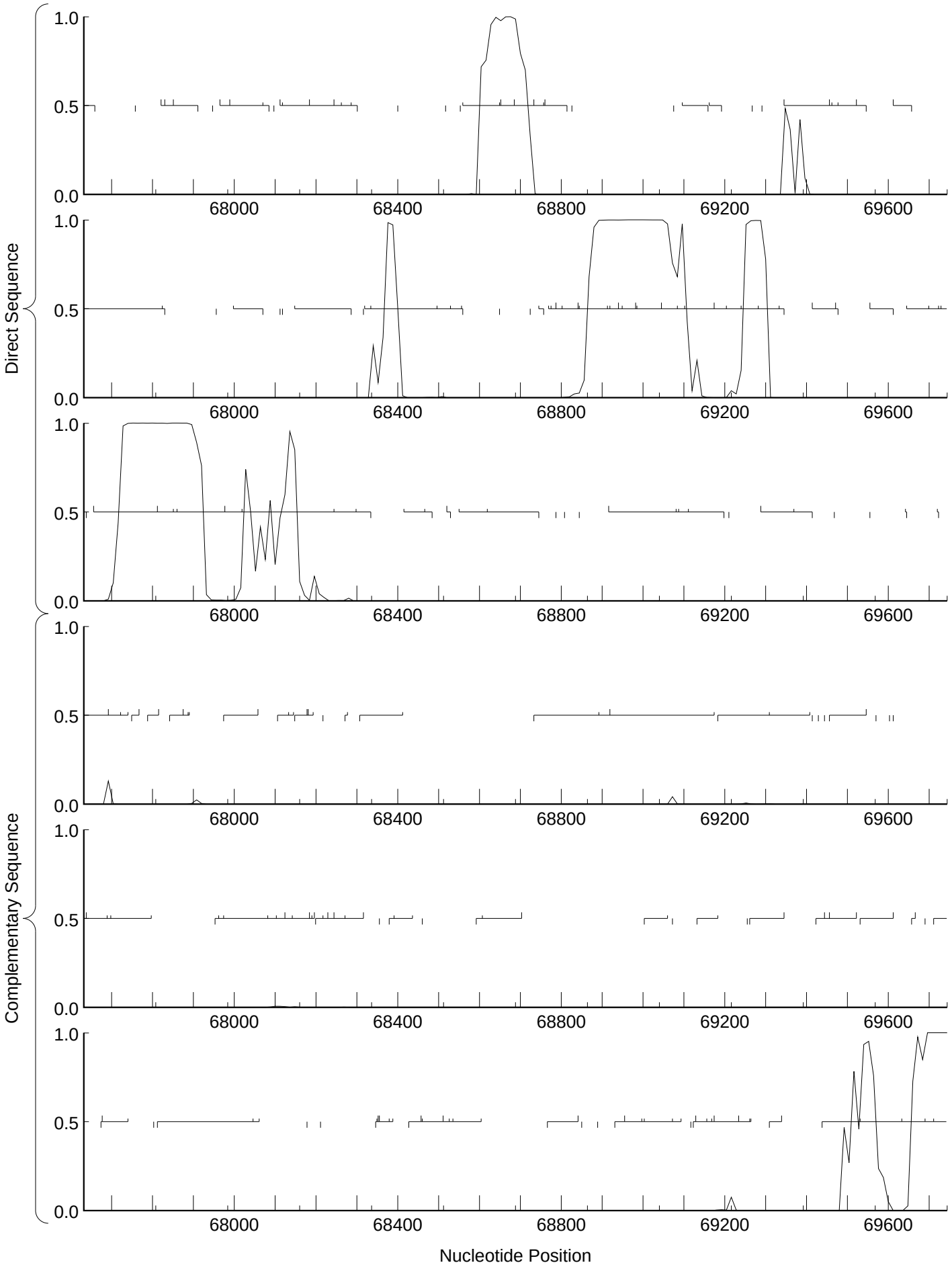
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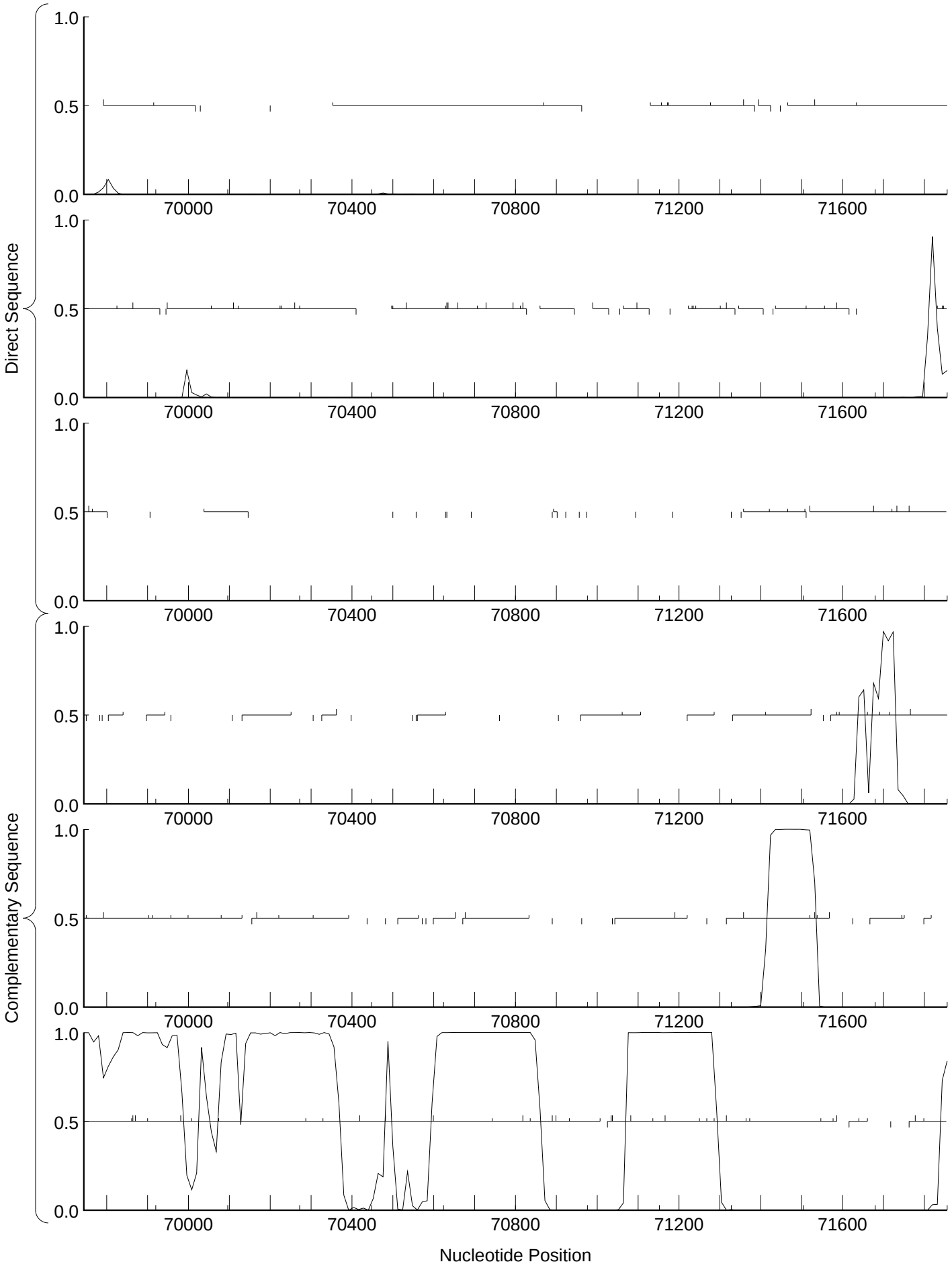
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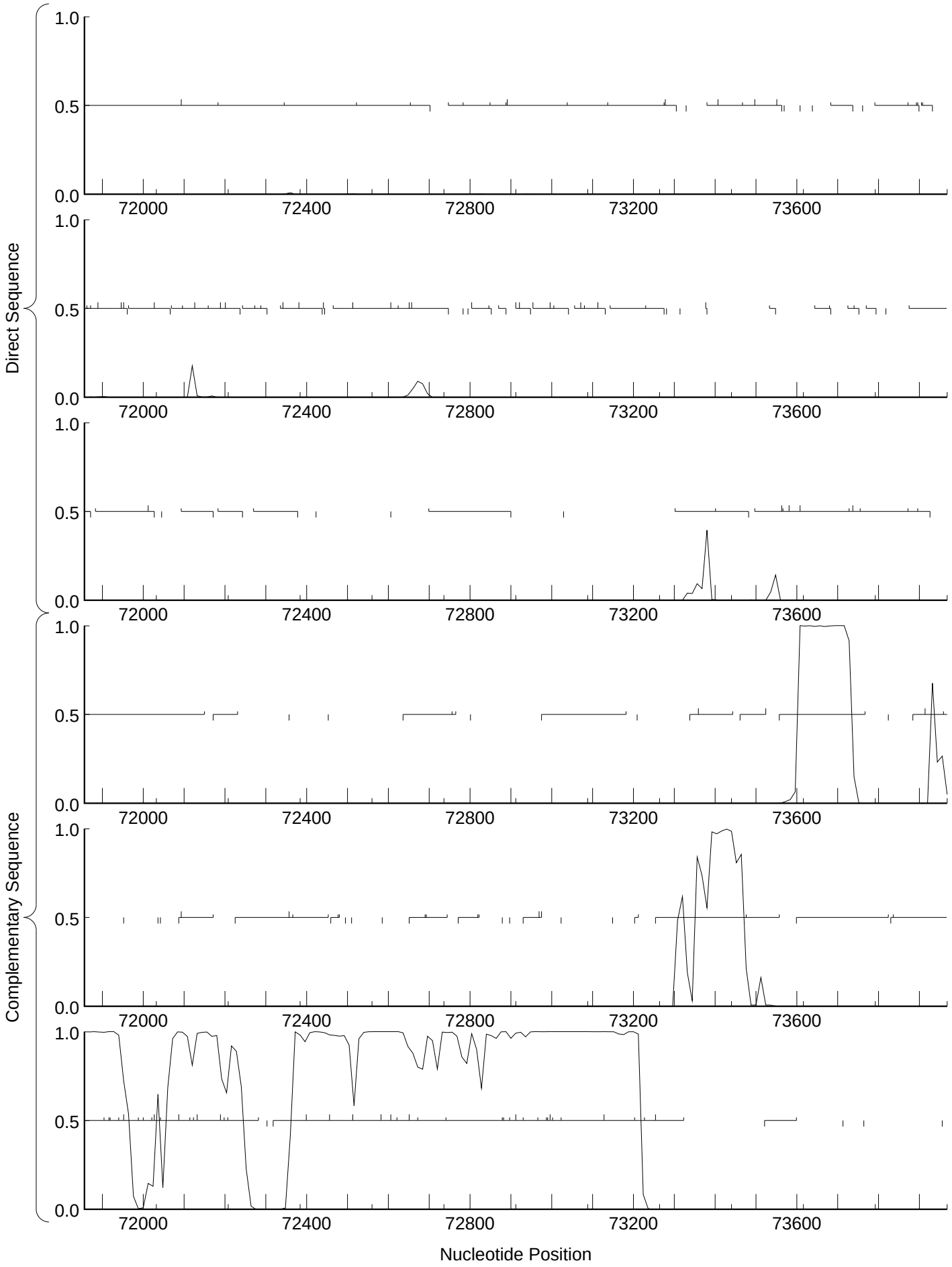


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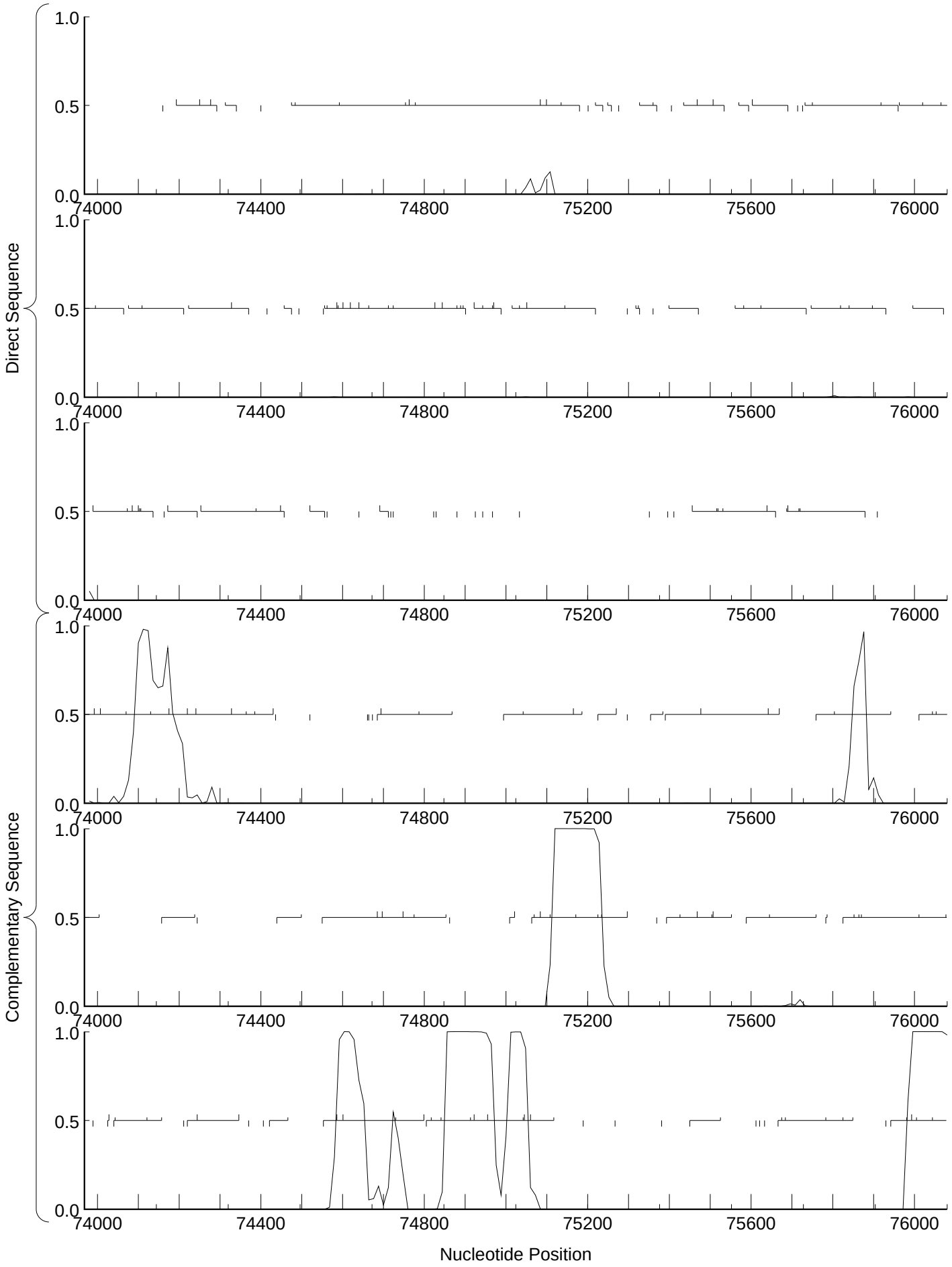


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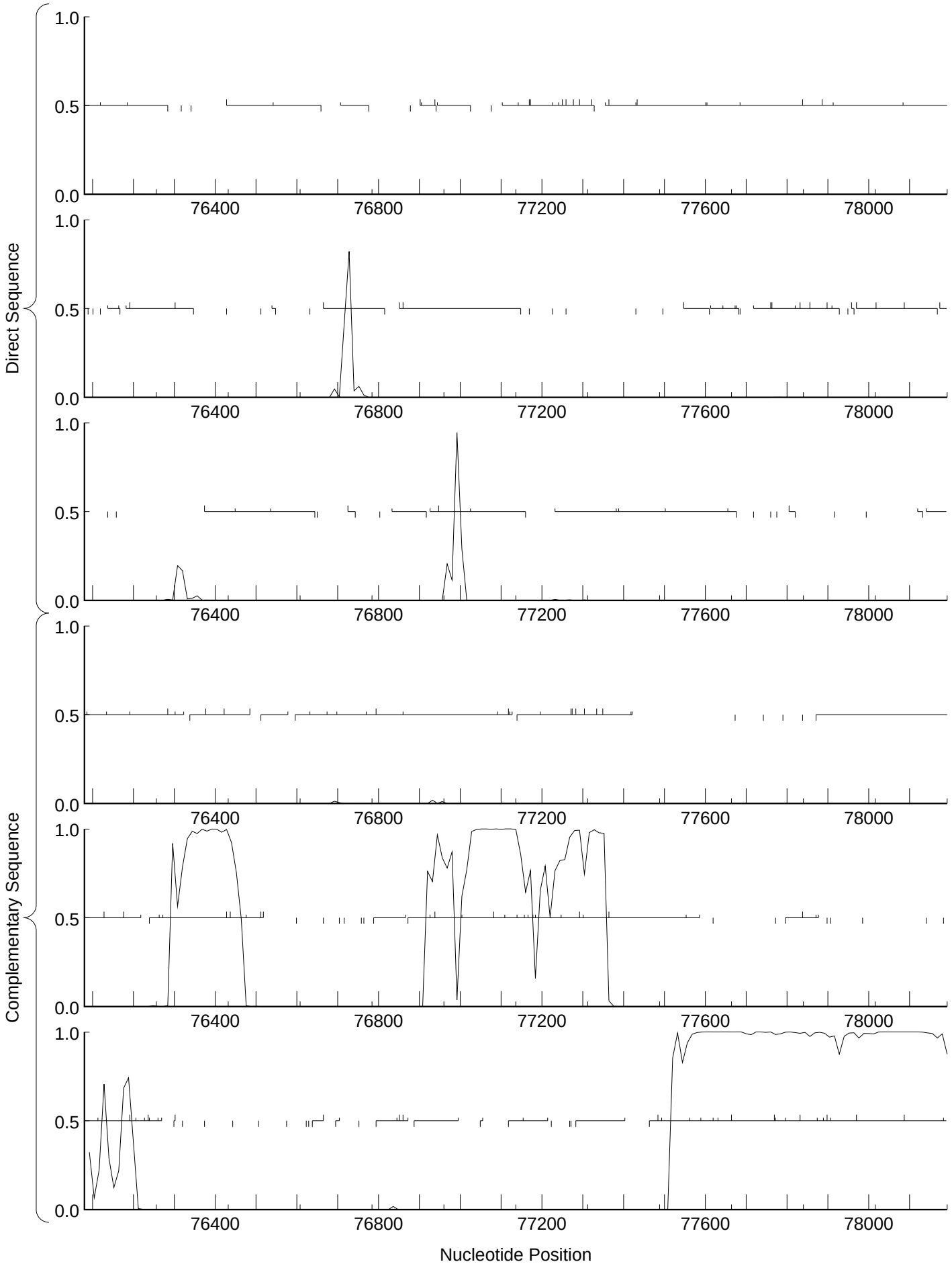



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Complementary Sequence
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0.5
0.0
72000
72400
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73200
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Nucleotide Position

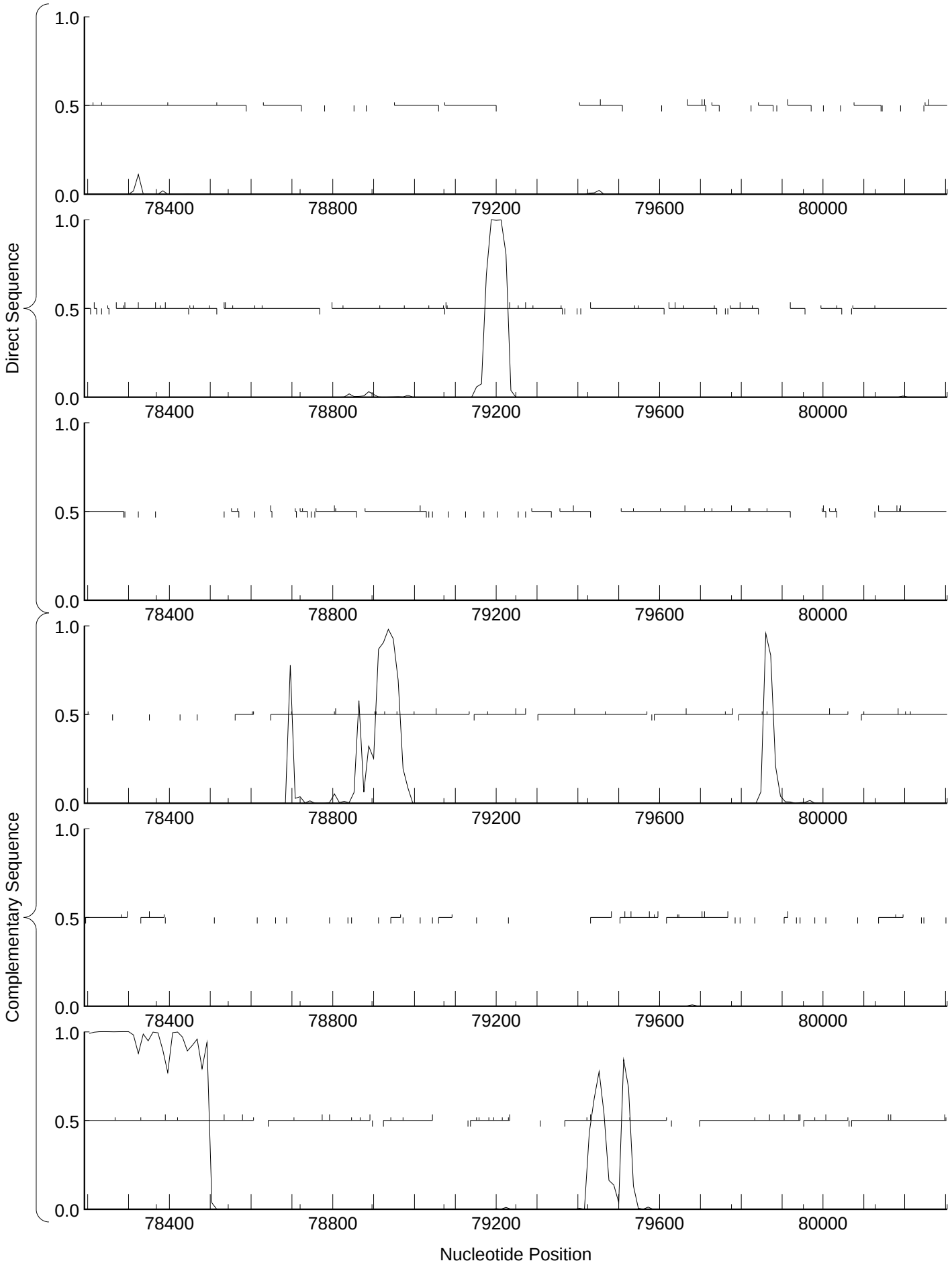
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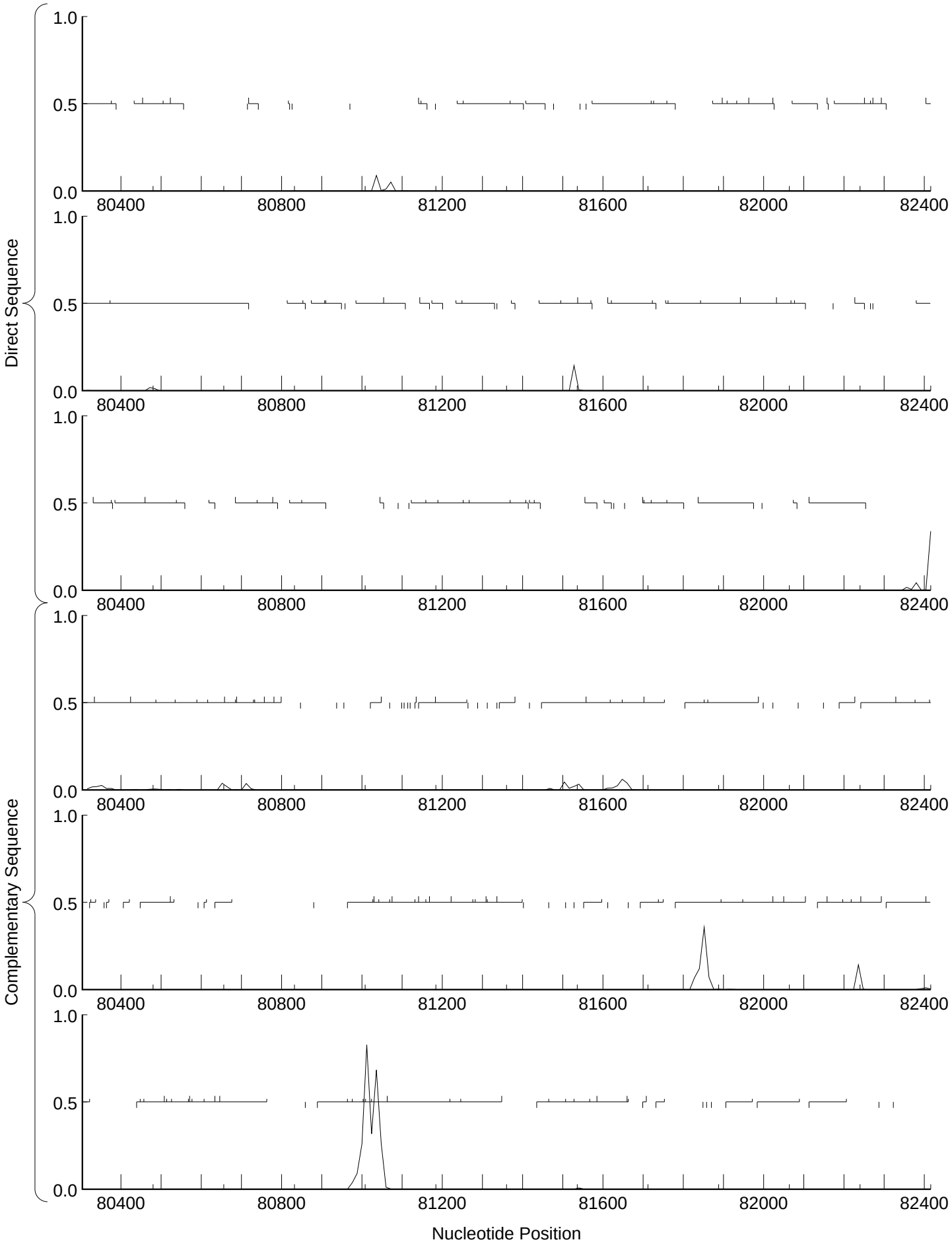
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complete sequence 84,303 bp including 11-base 3’ overhang (ACCCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 39/41



complete sequence 84,303 bp including 11-base 3' overhang (ACCCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 40/41



complete sequence 84,303 bp including 11-base 3’ overhang (ACCCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 41/41

