GeneMark

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

PROGRAM INFORMATION

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Sequence         :  Mycobacterium phage Tonenili complete sequence, 160985 bp, circularly permuted, Cluster C1
Analysis Date    :  3/29/24 at 4:58:53
Pages            :  78
Sequence Length  :  160985 bp
GC Content       :   64.09%

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

PS-Version       :  1.2

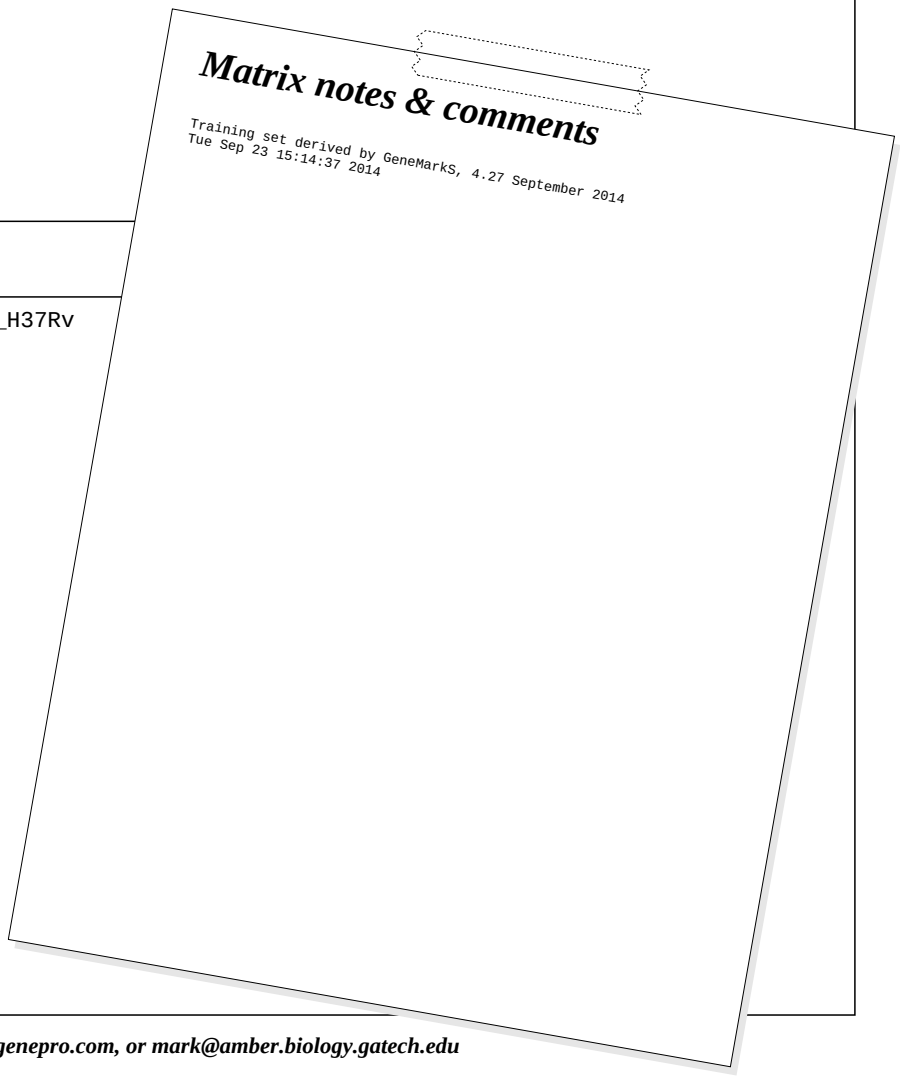
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MATRIX INFORMATION

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Matrix     :  Mycobacterium_tuberculosis_H37Rv
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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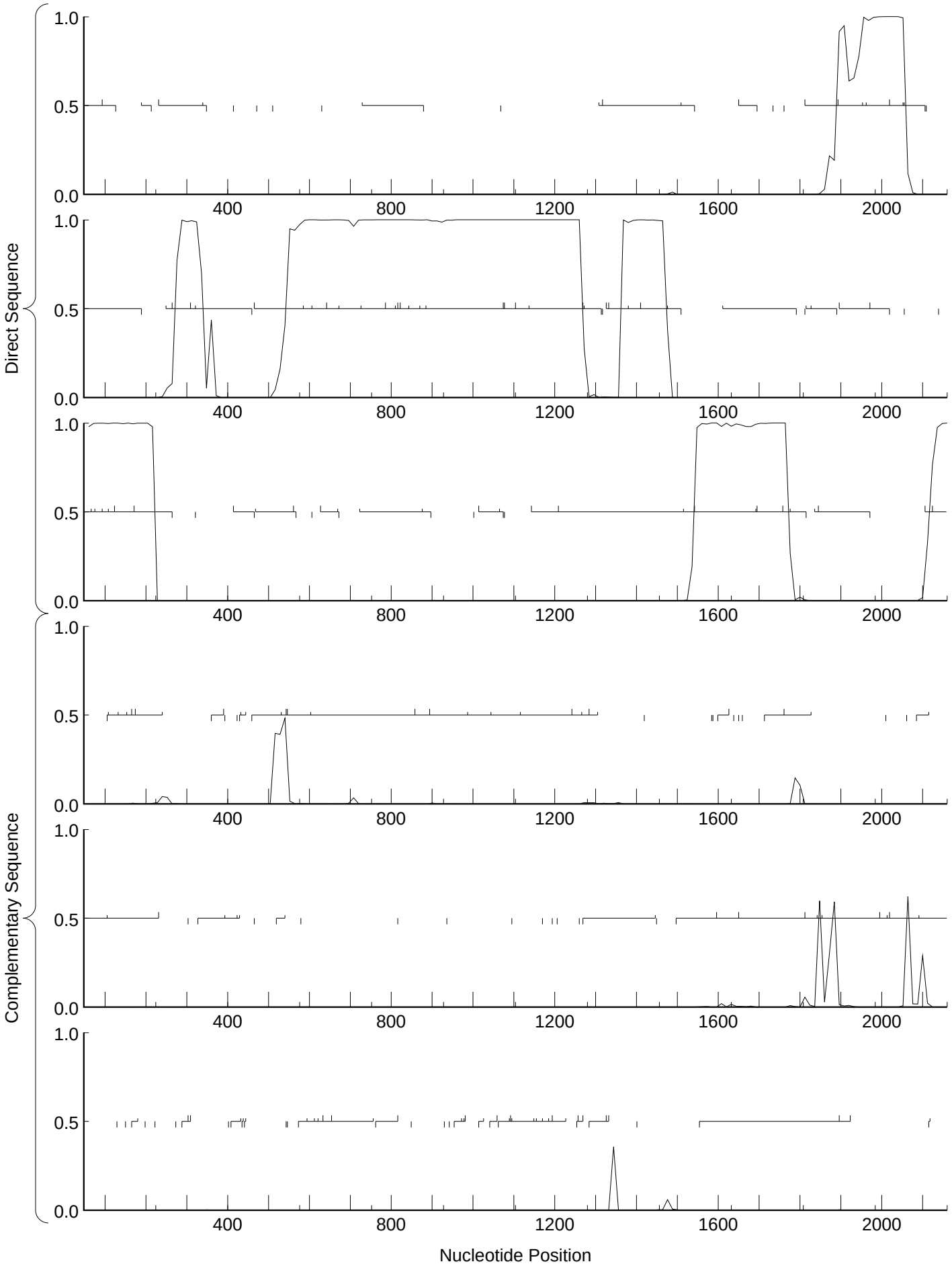


Matrix notes & comments

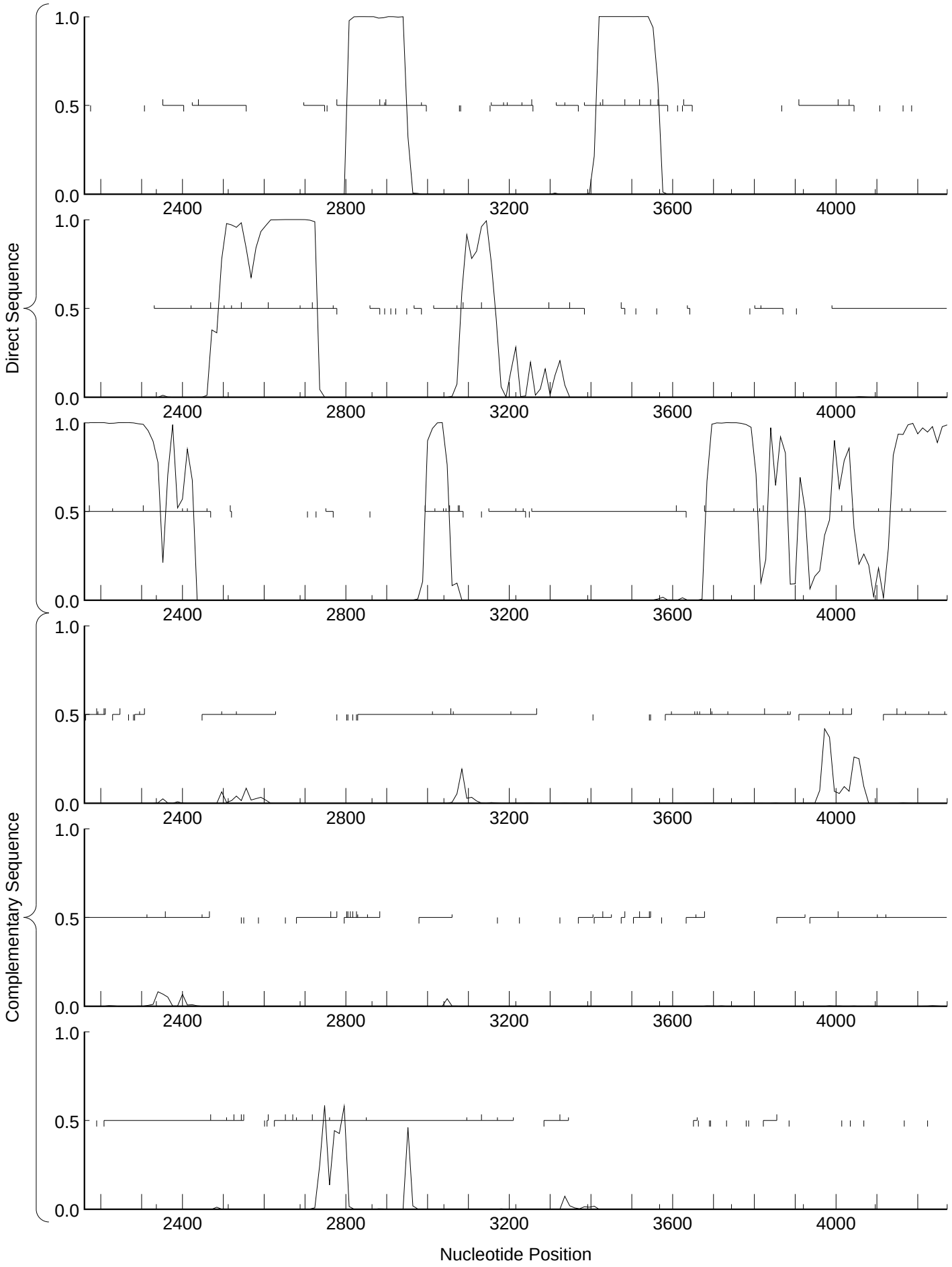
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

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

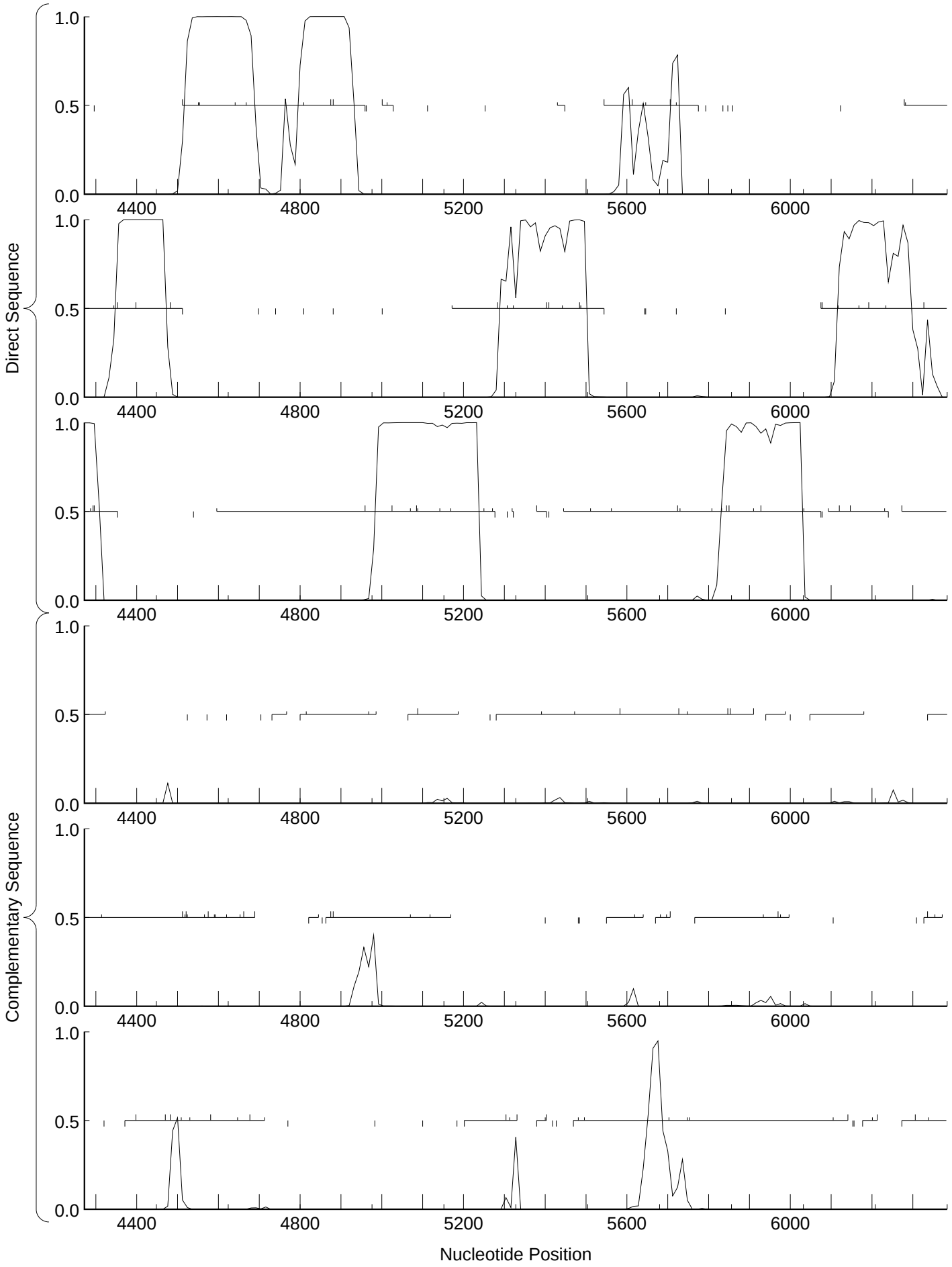
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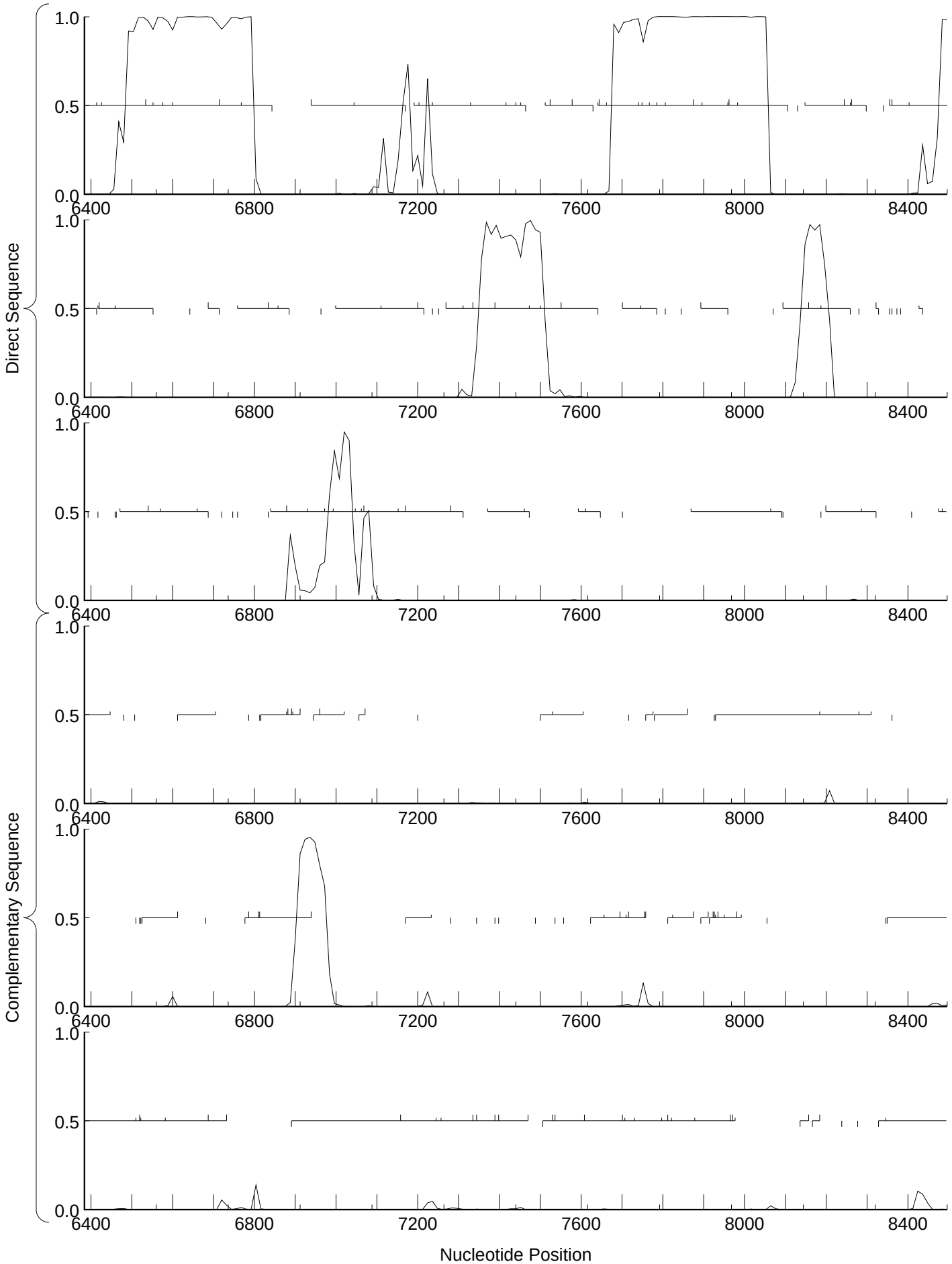
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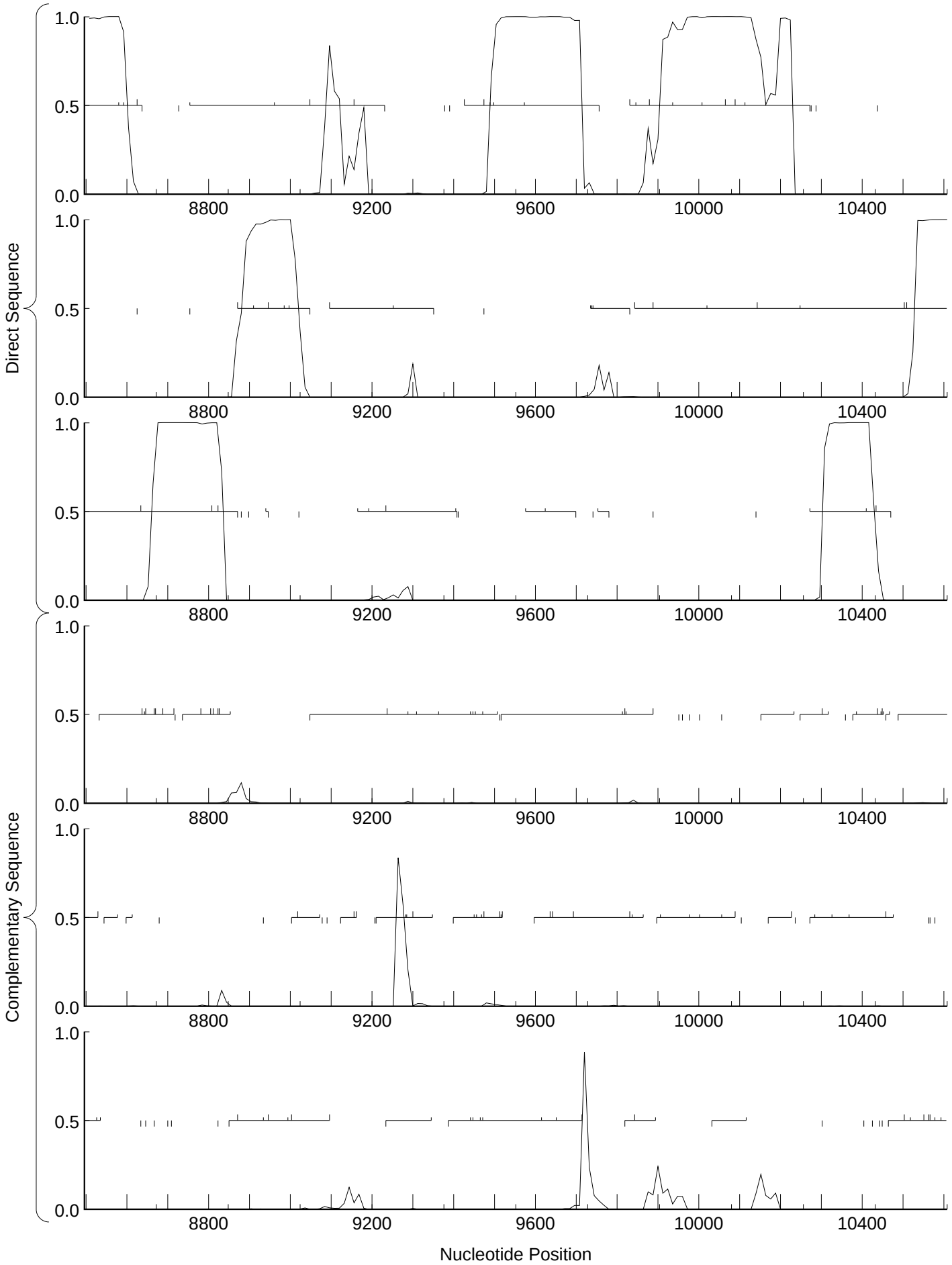
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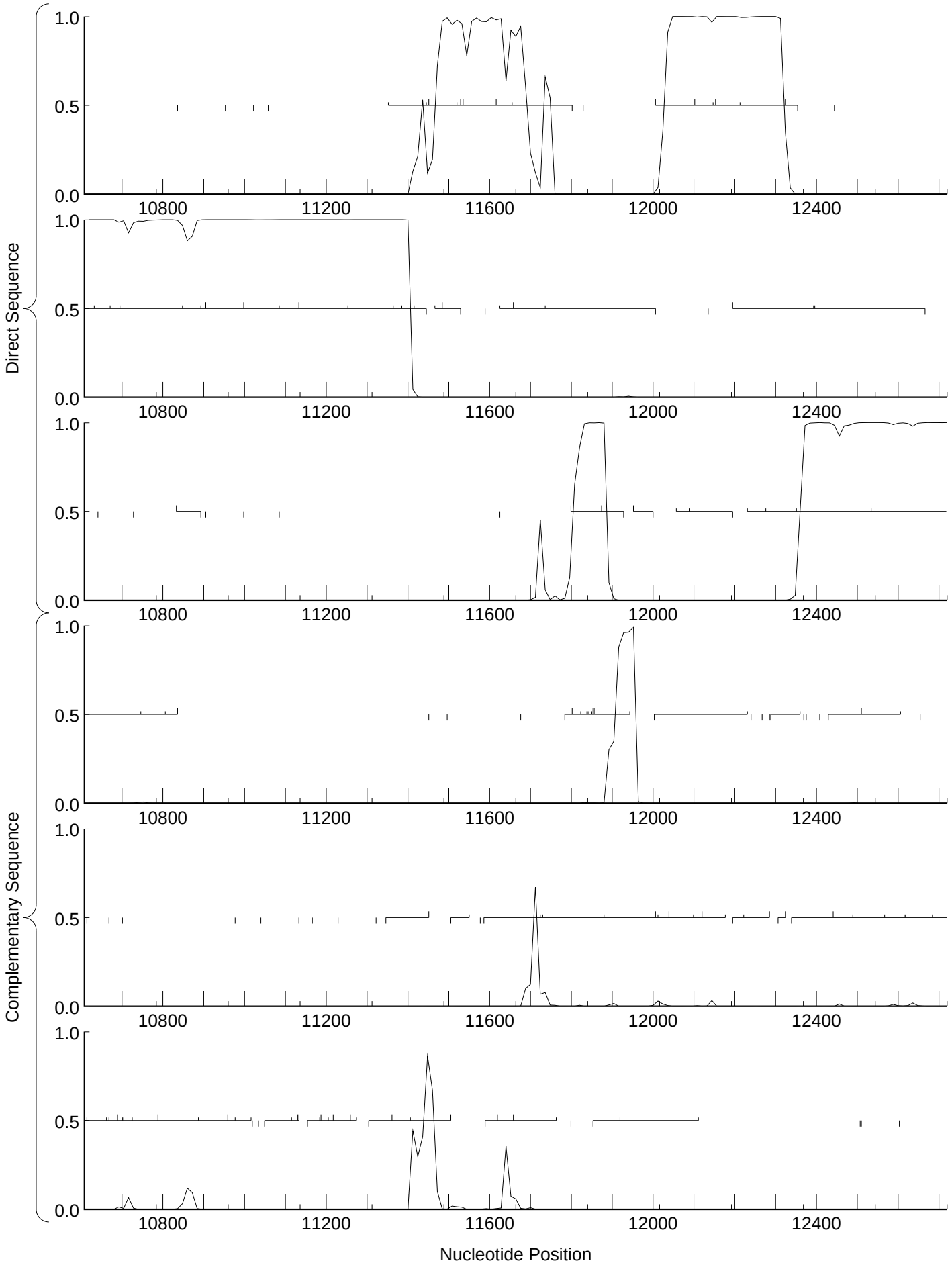
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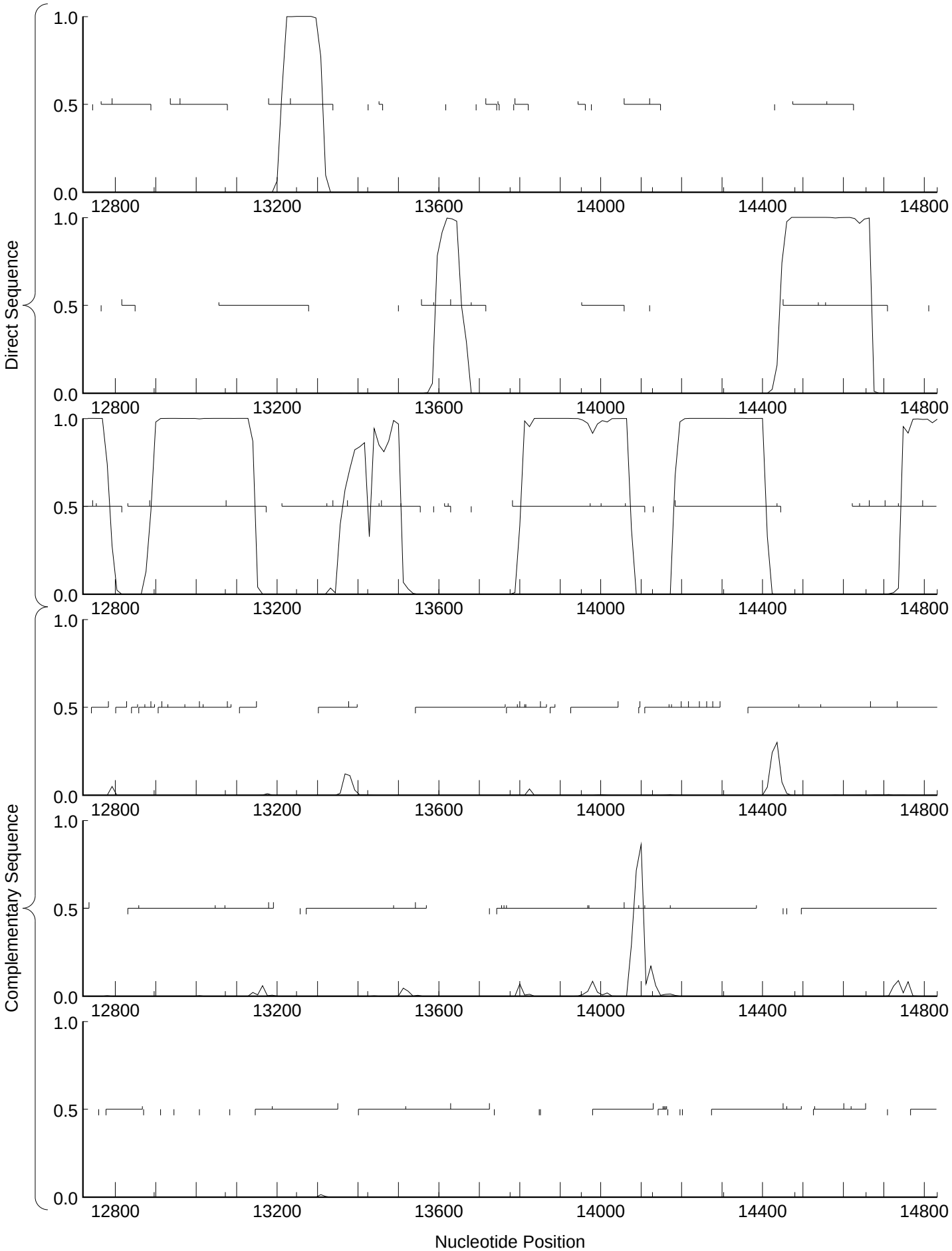
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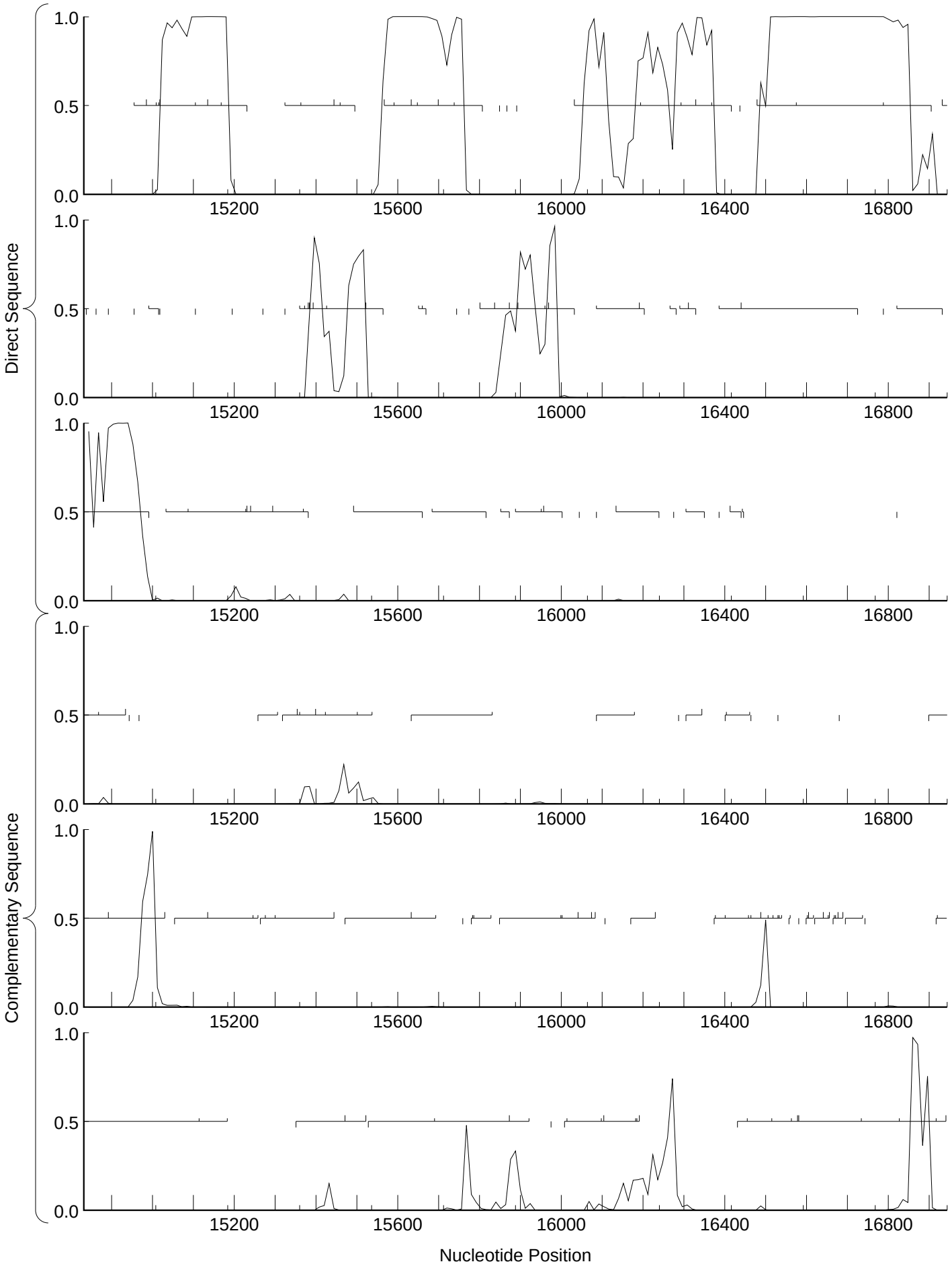
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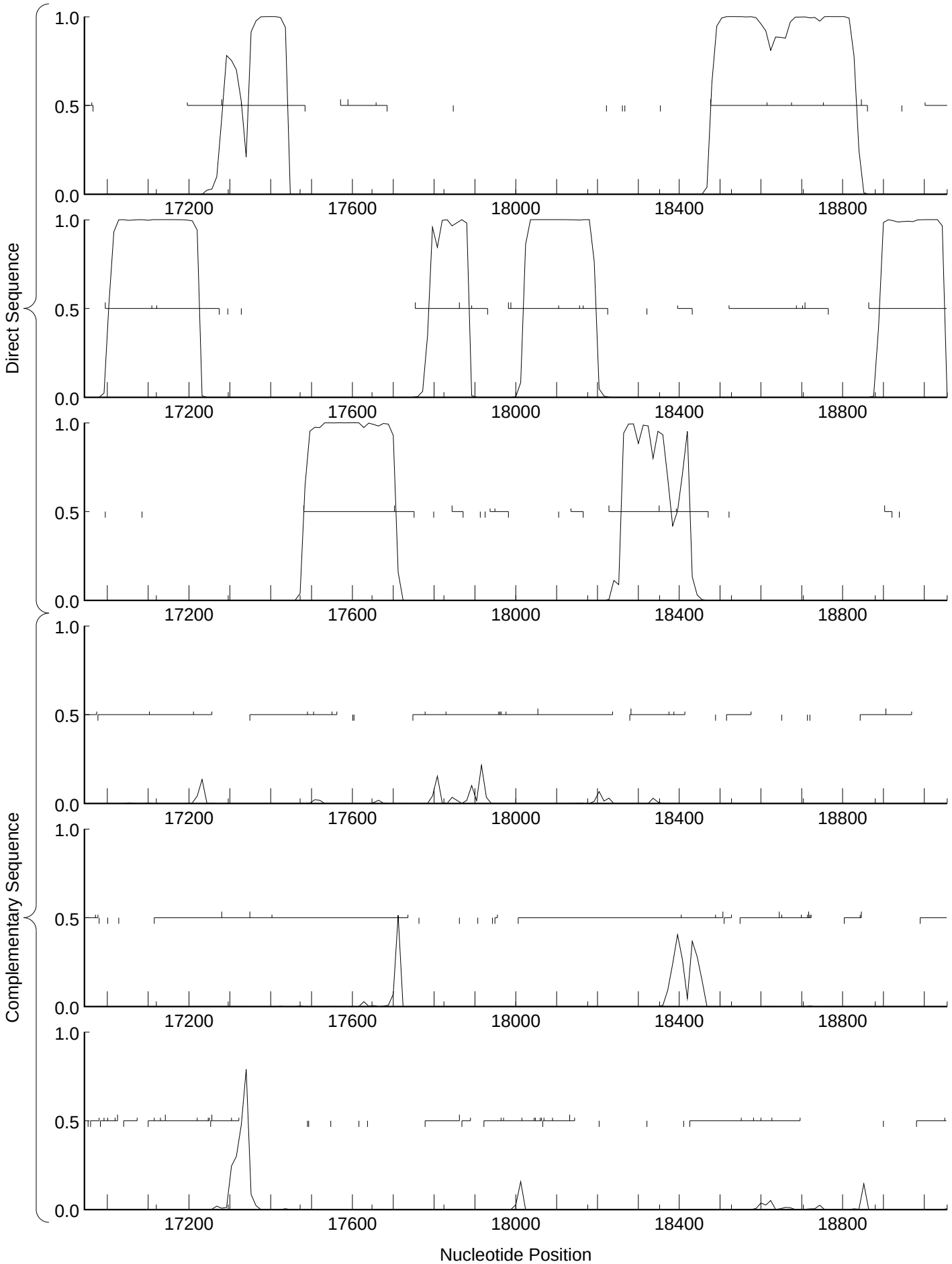
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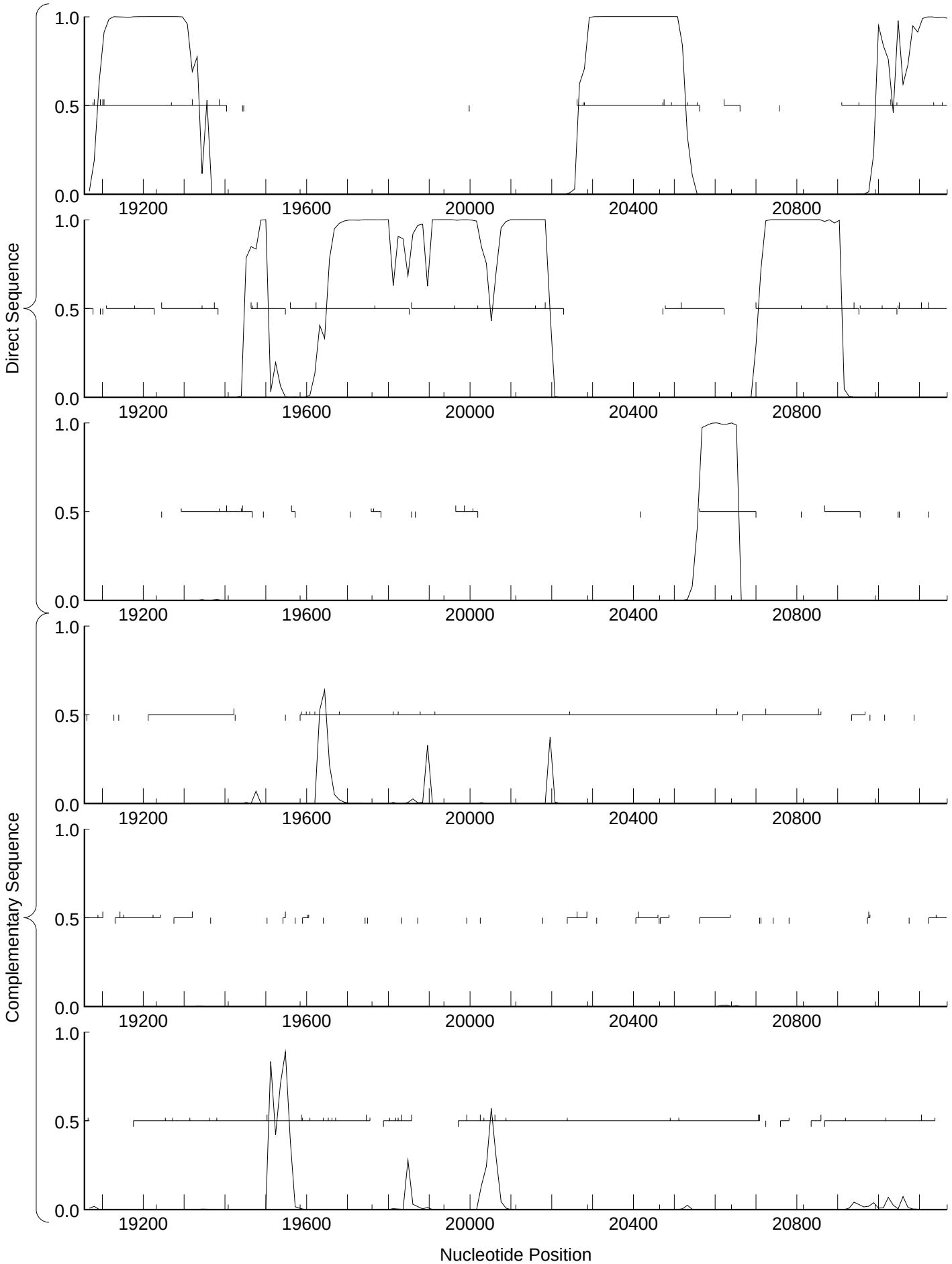
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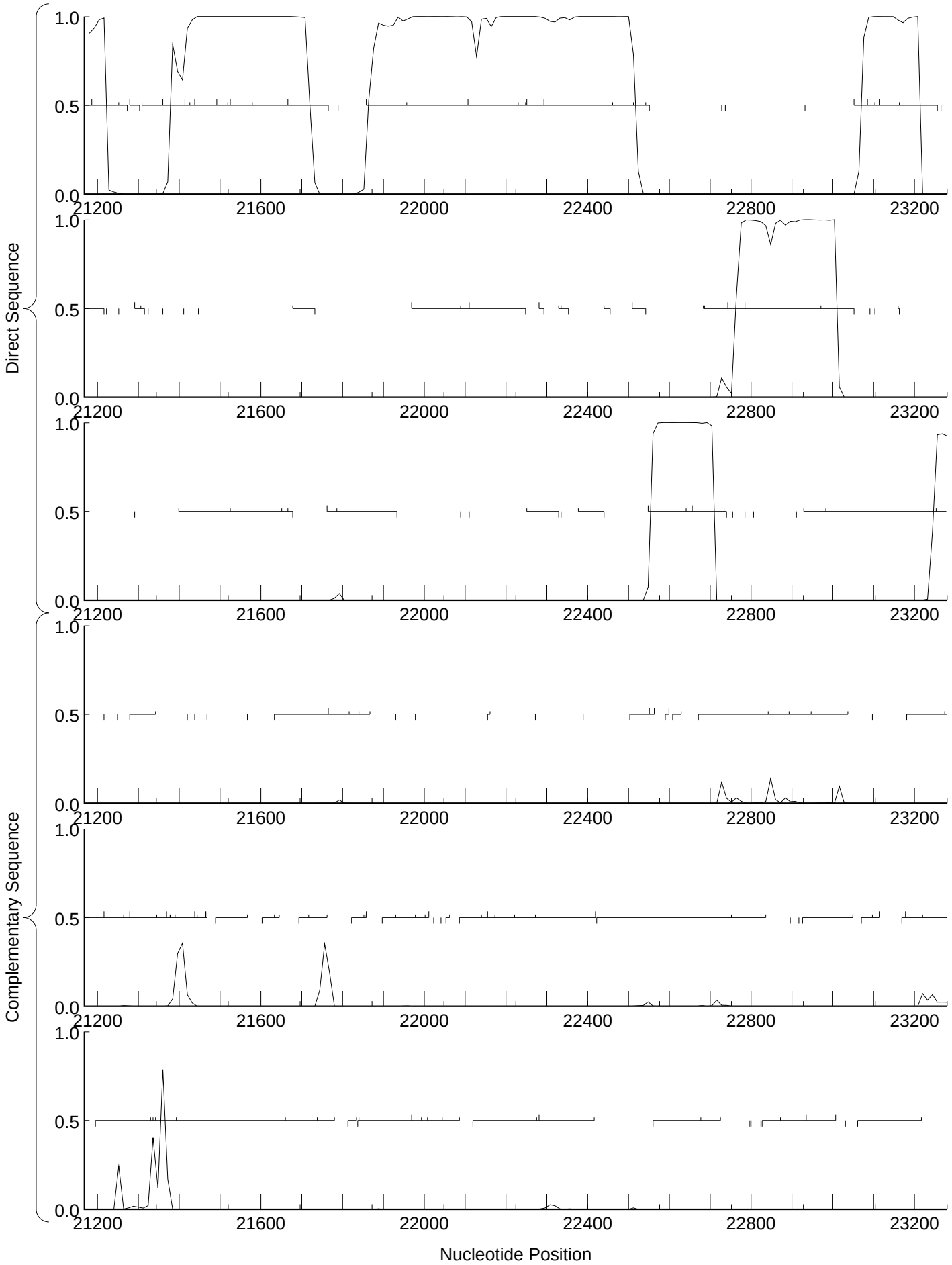
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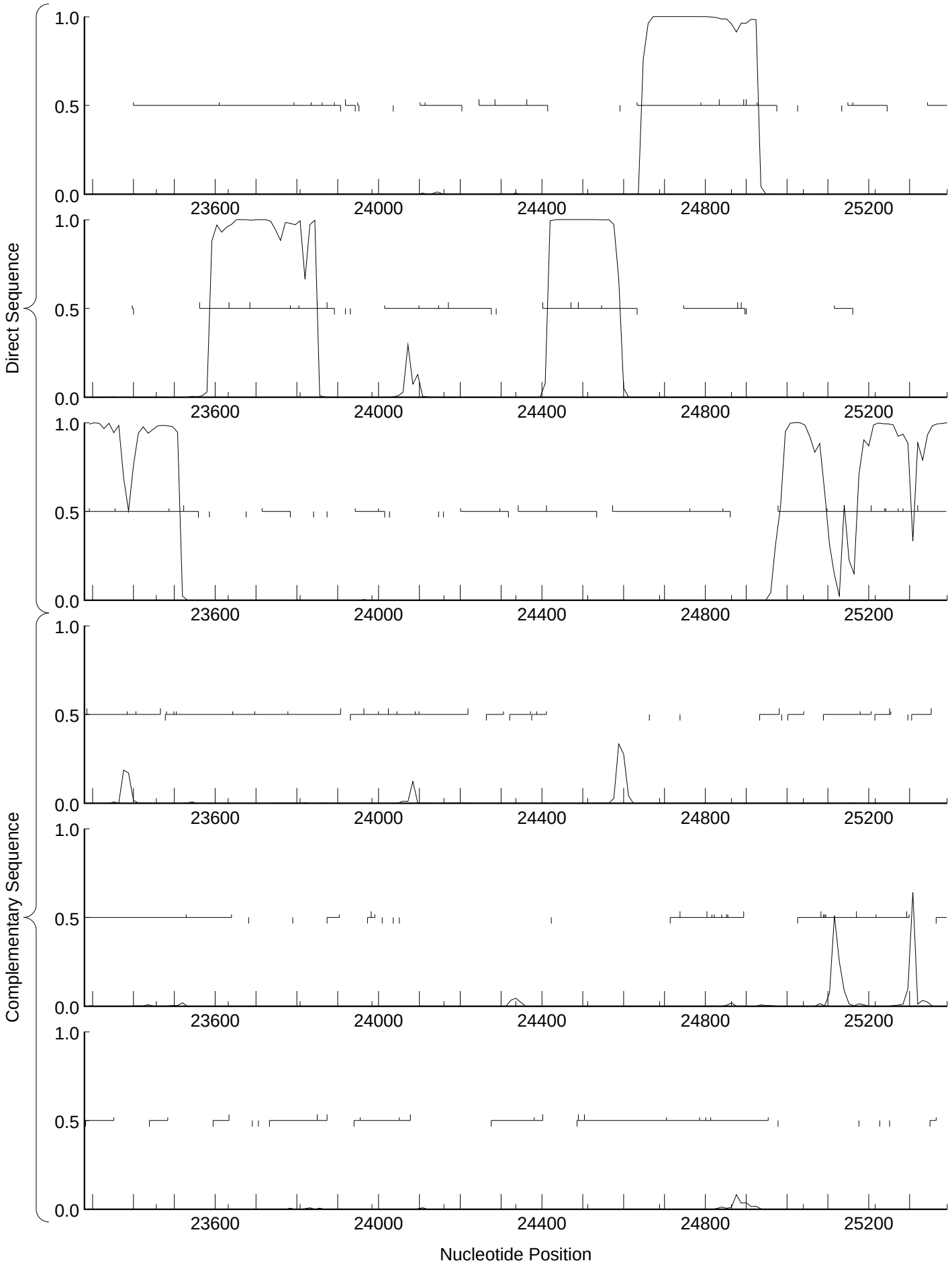
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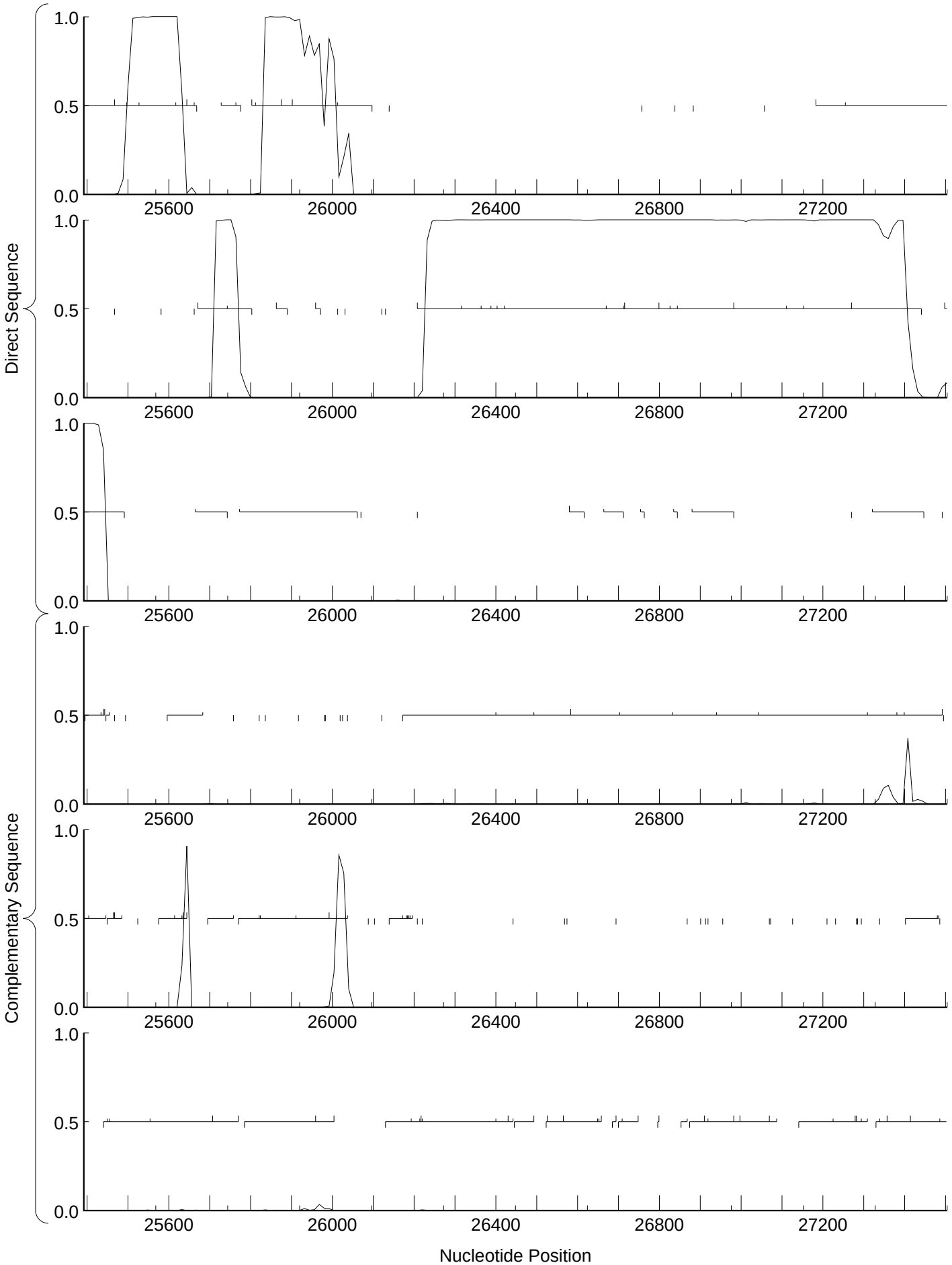
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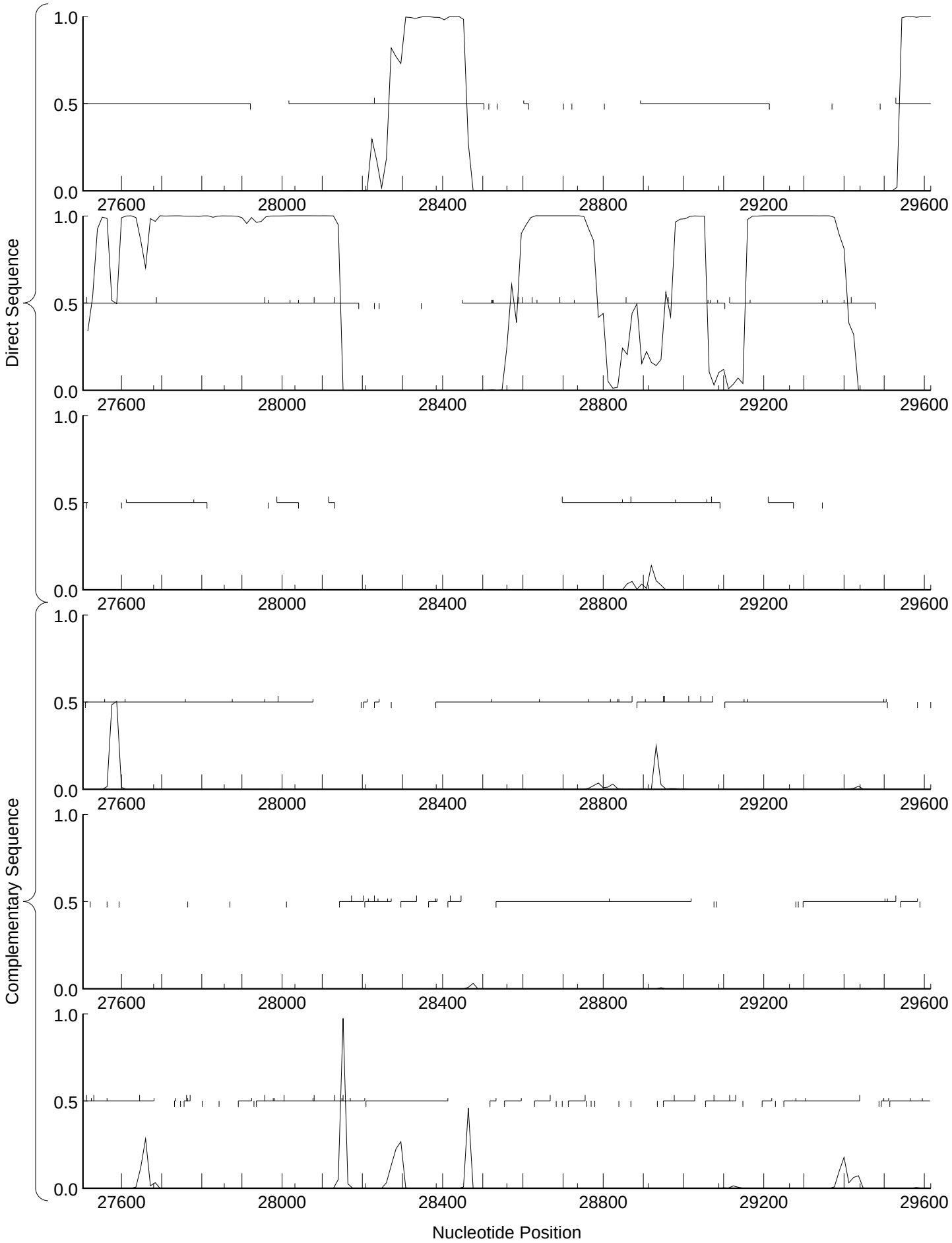
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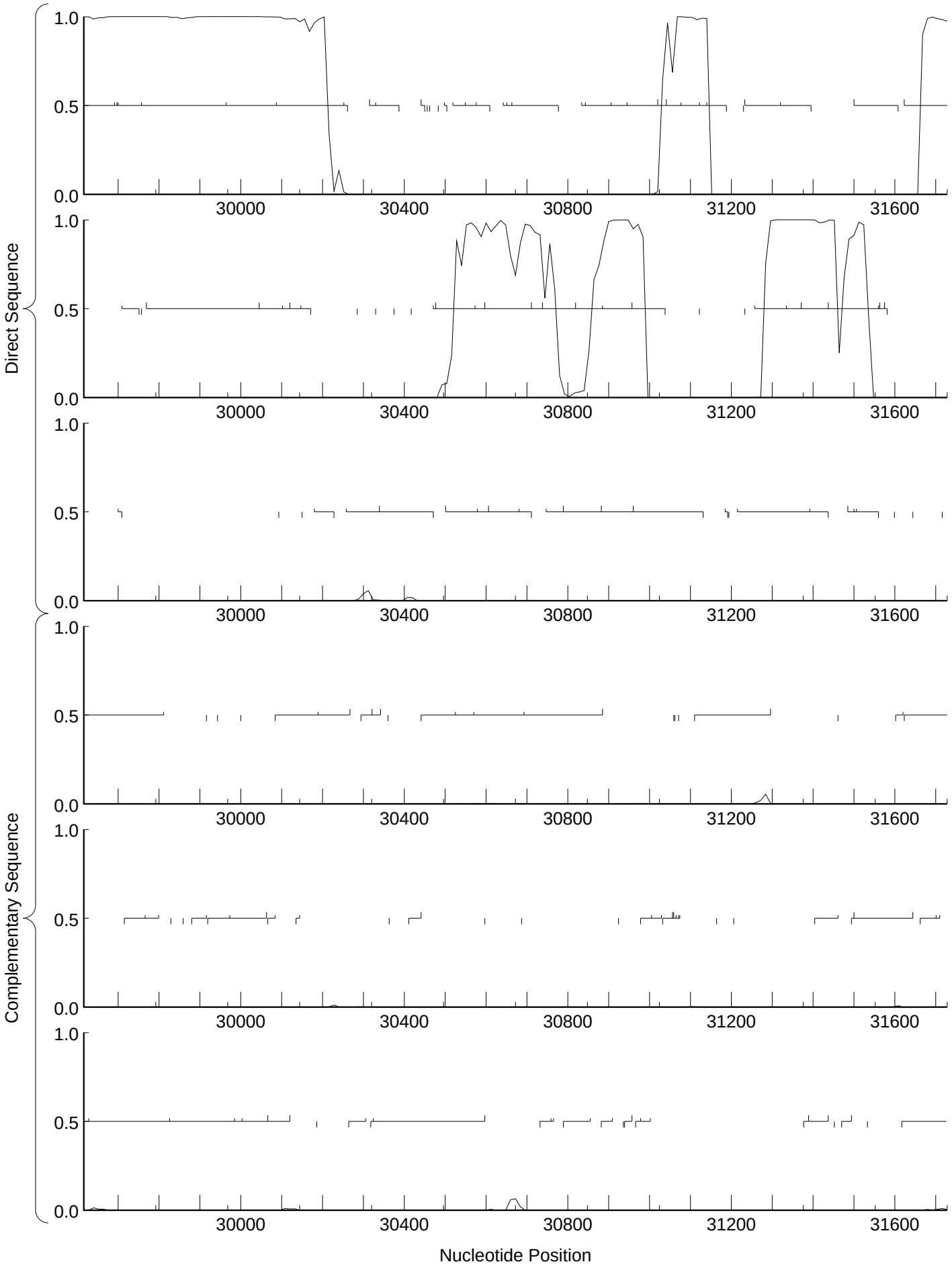
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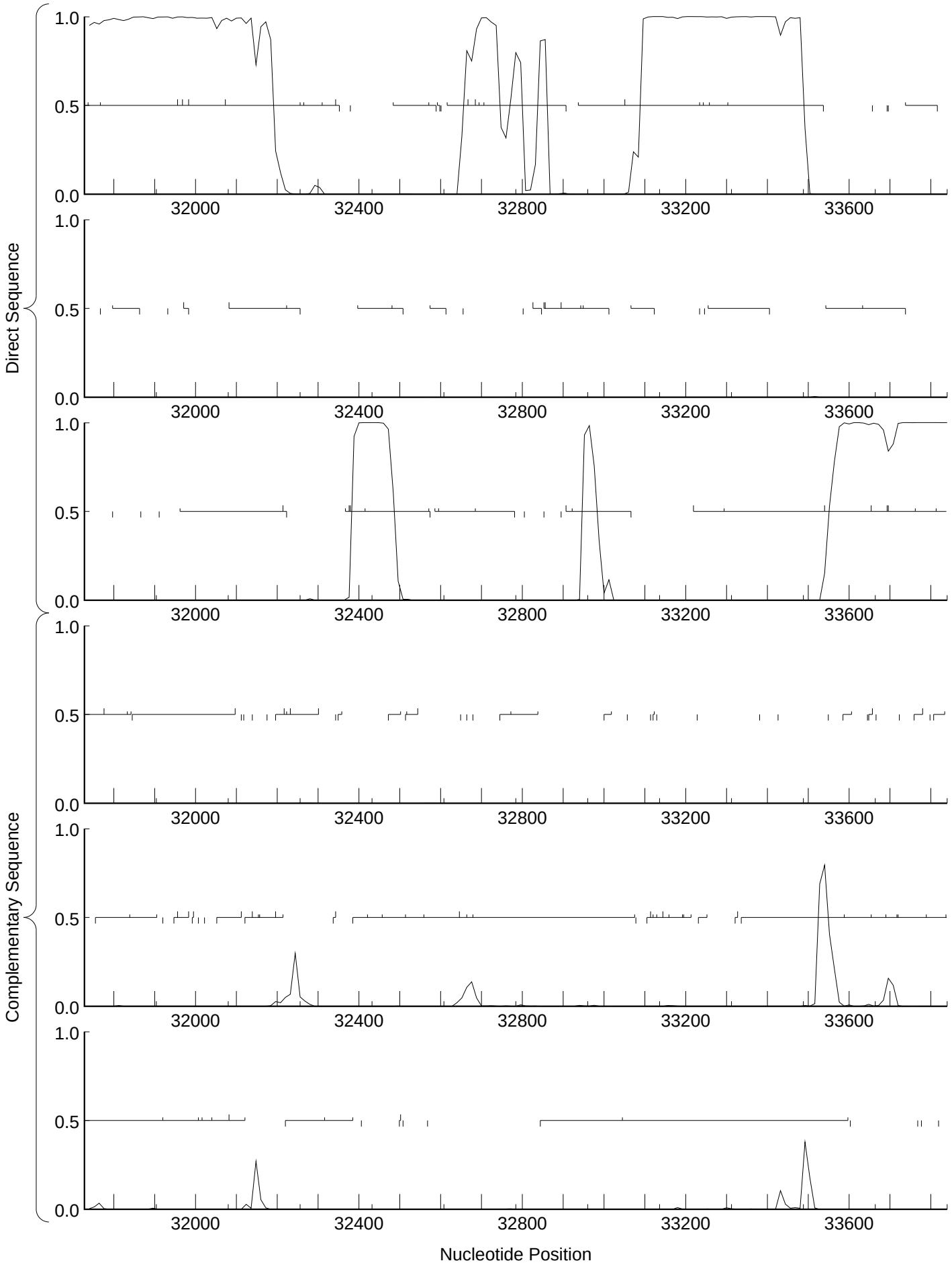
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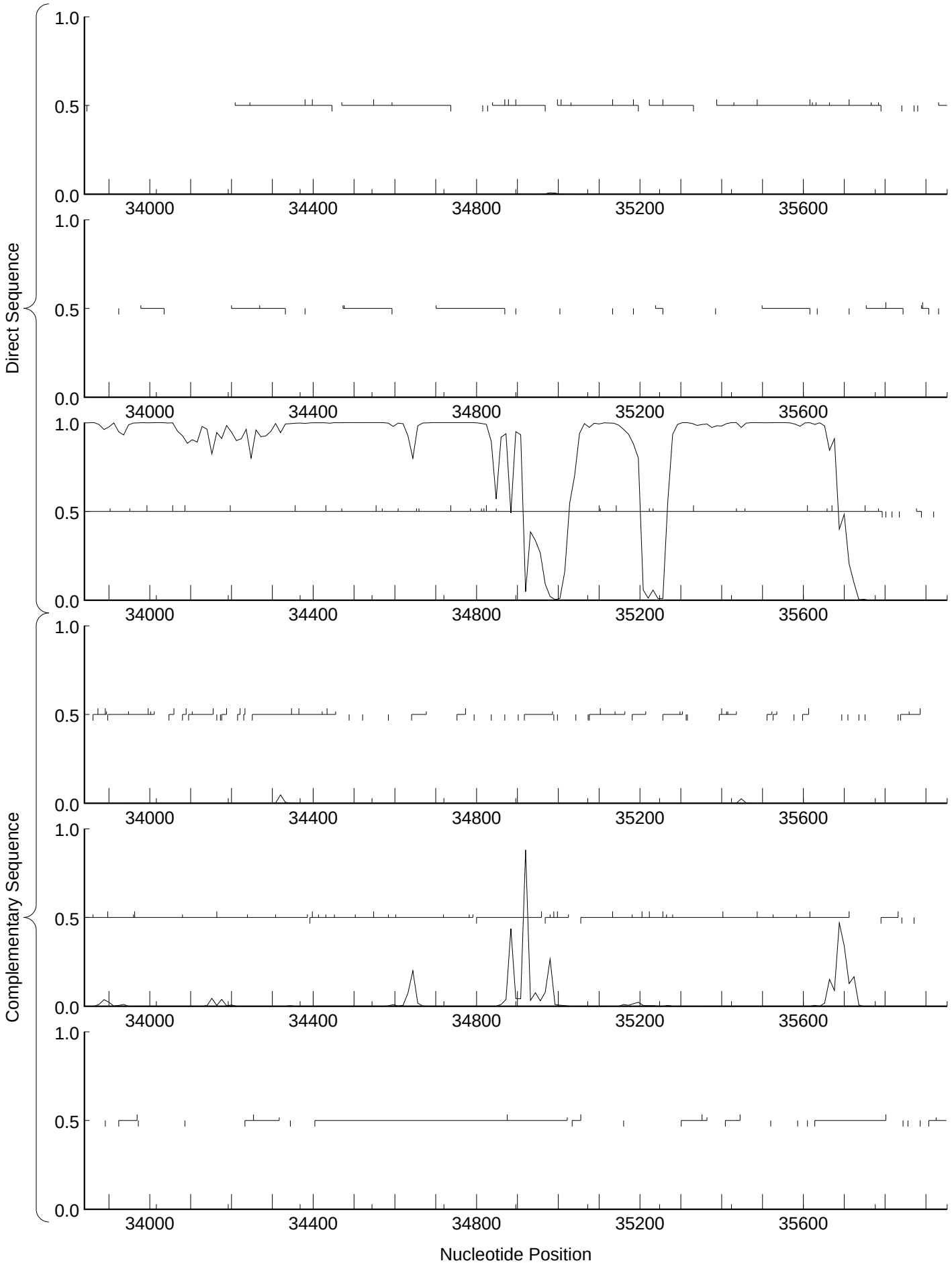
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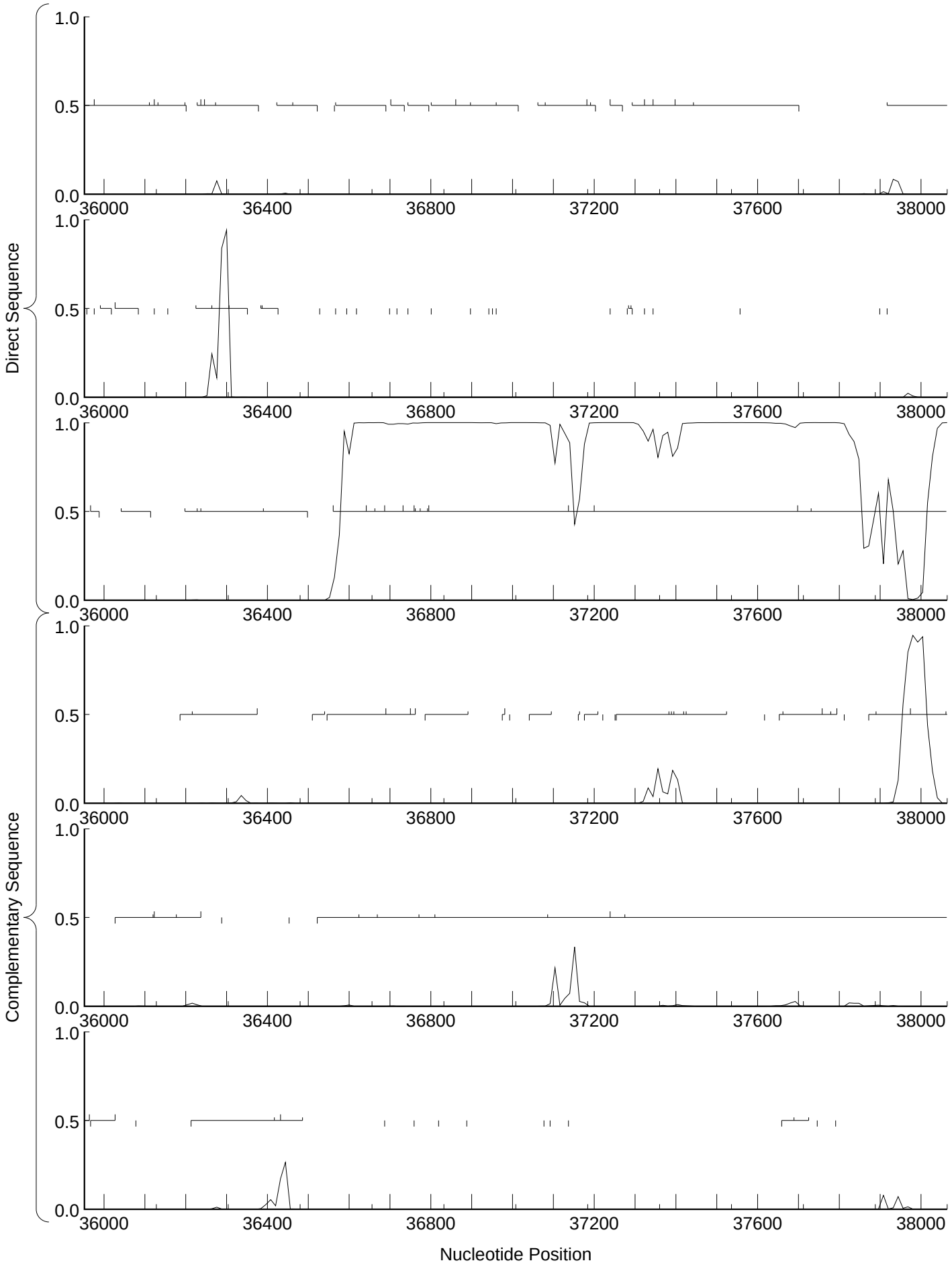
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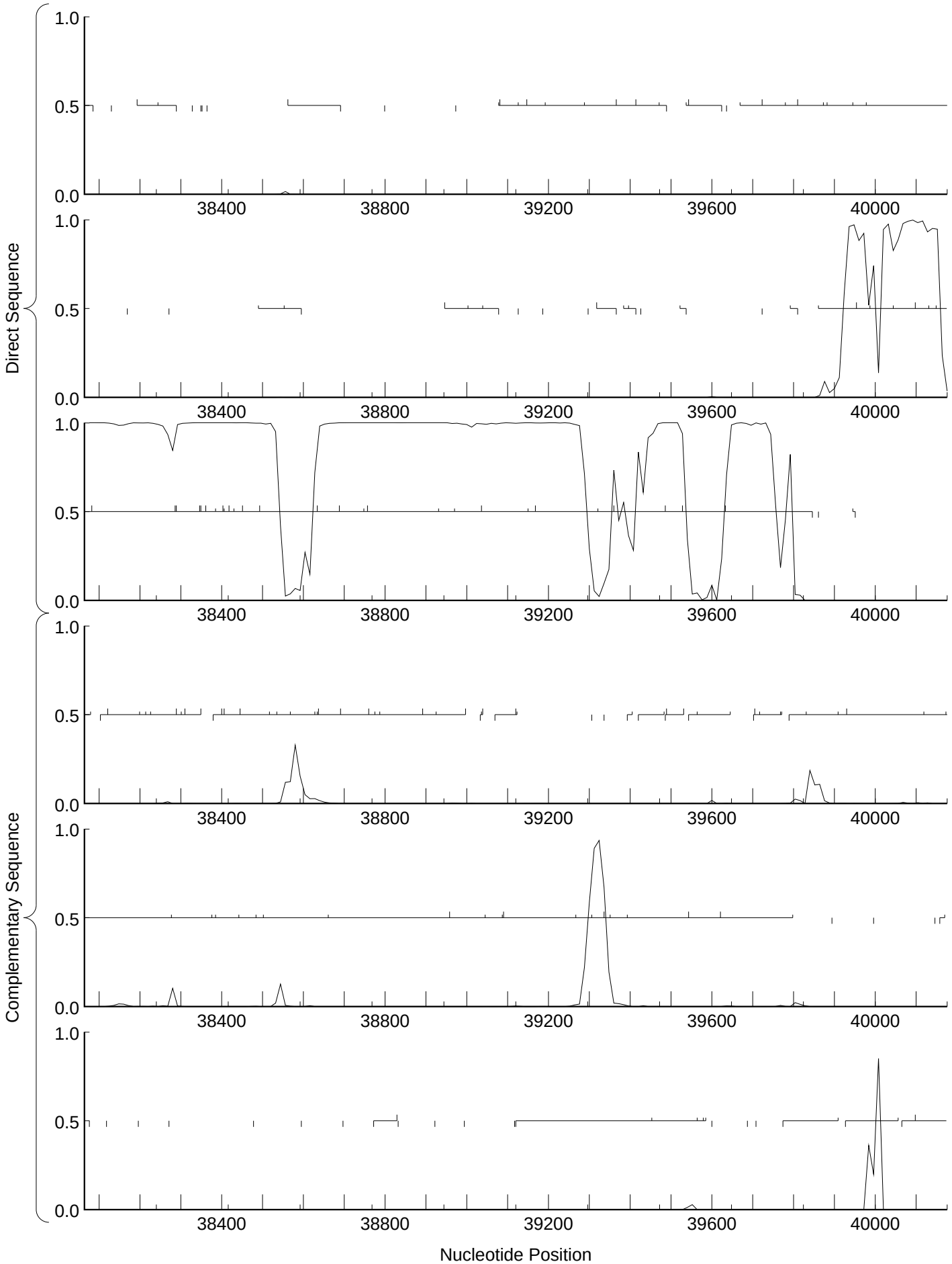
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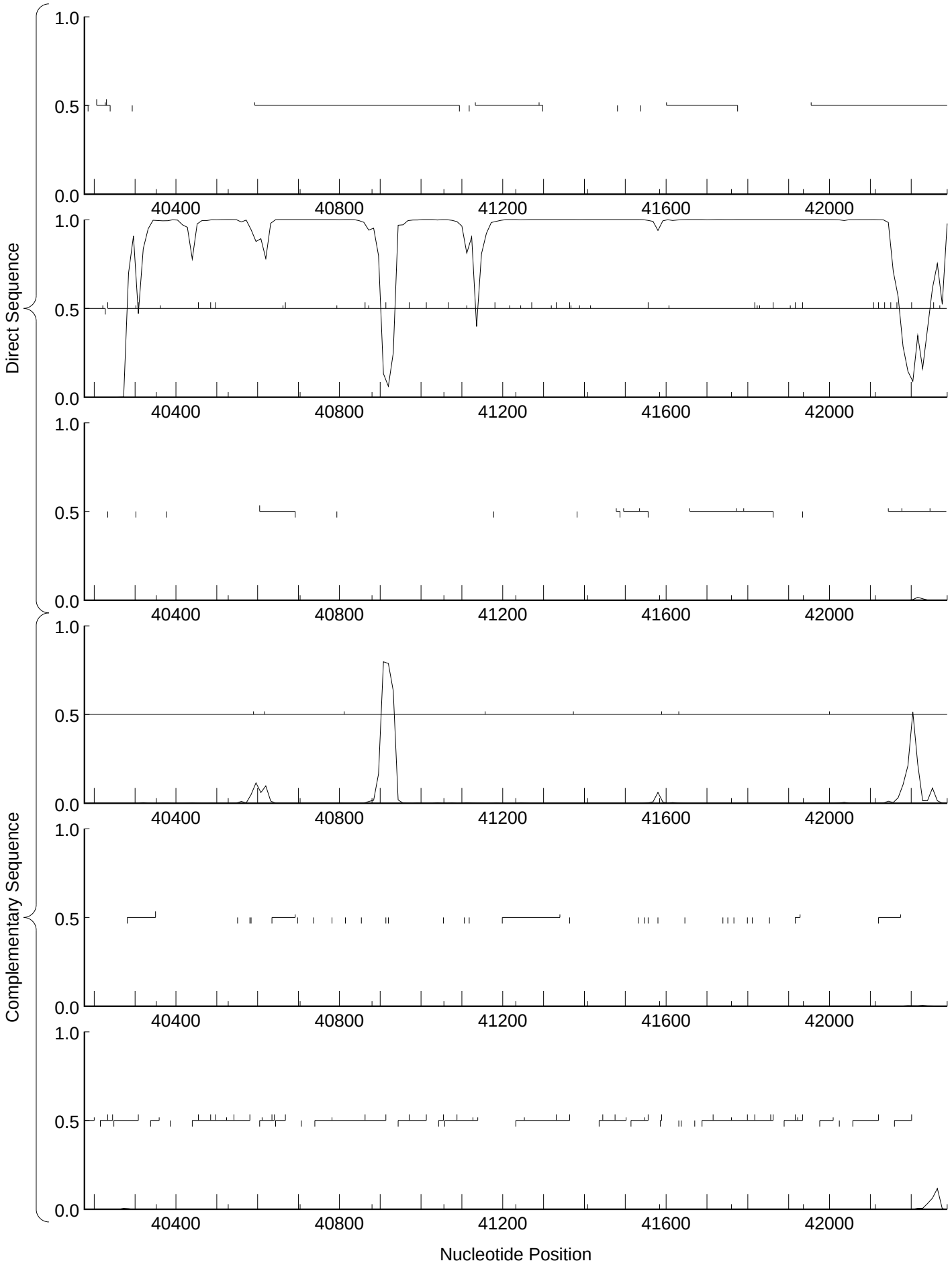
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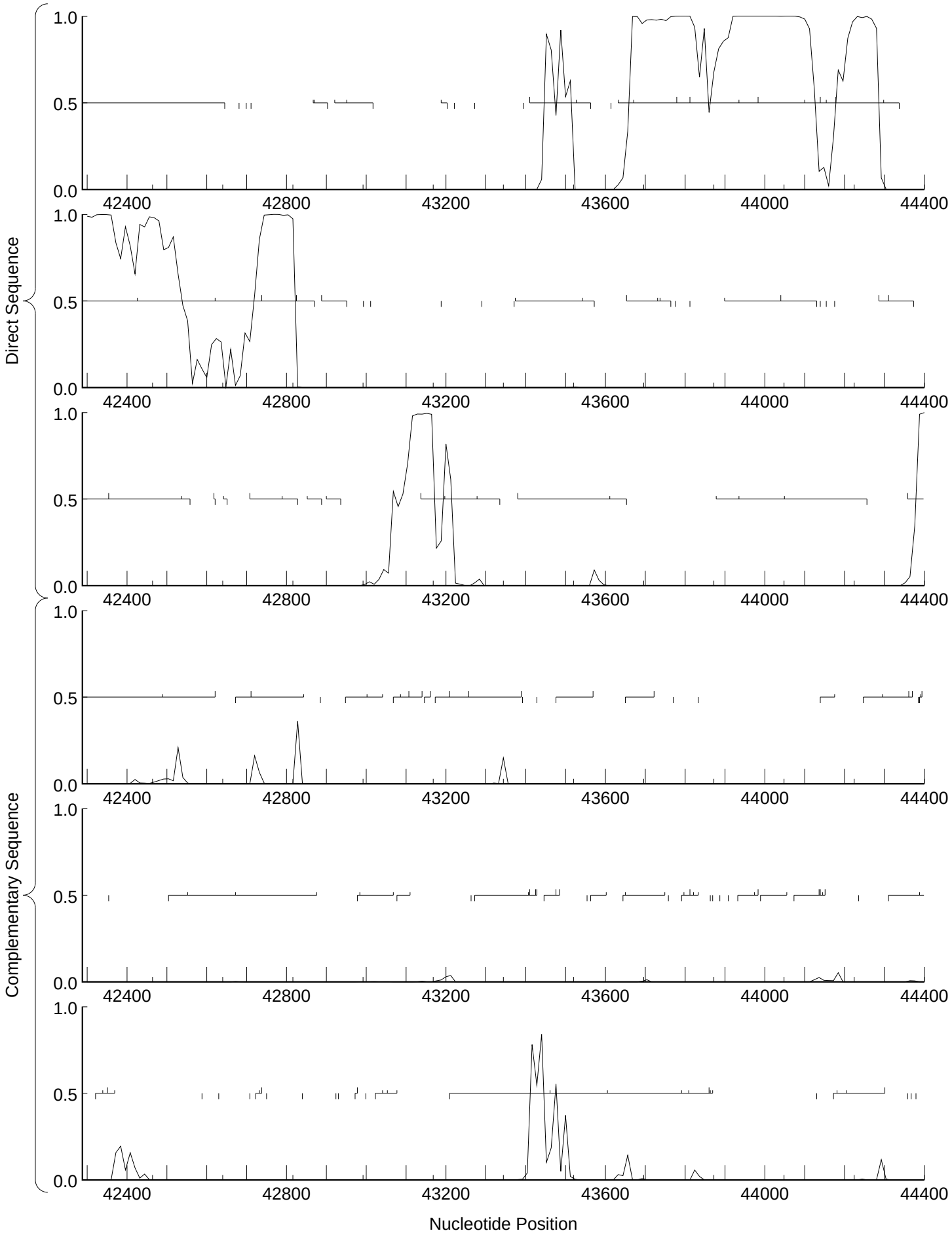
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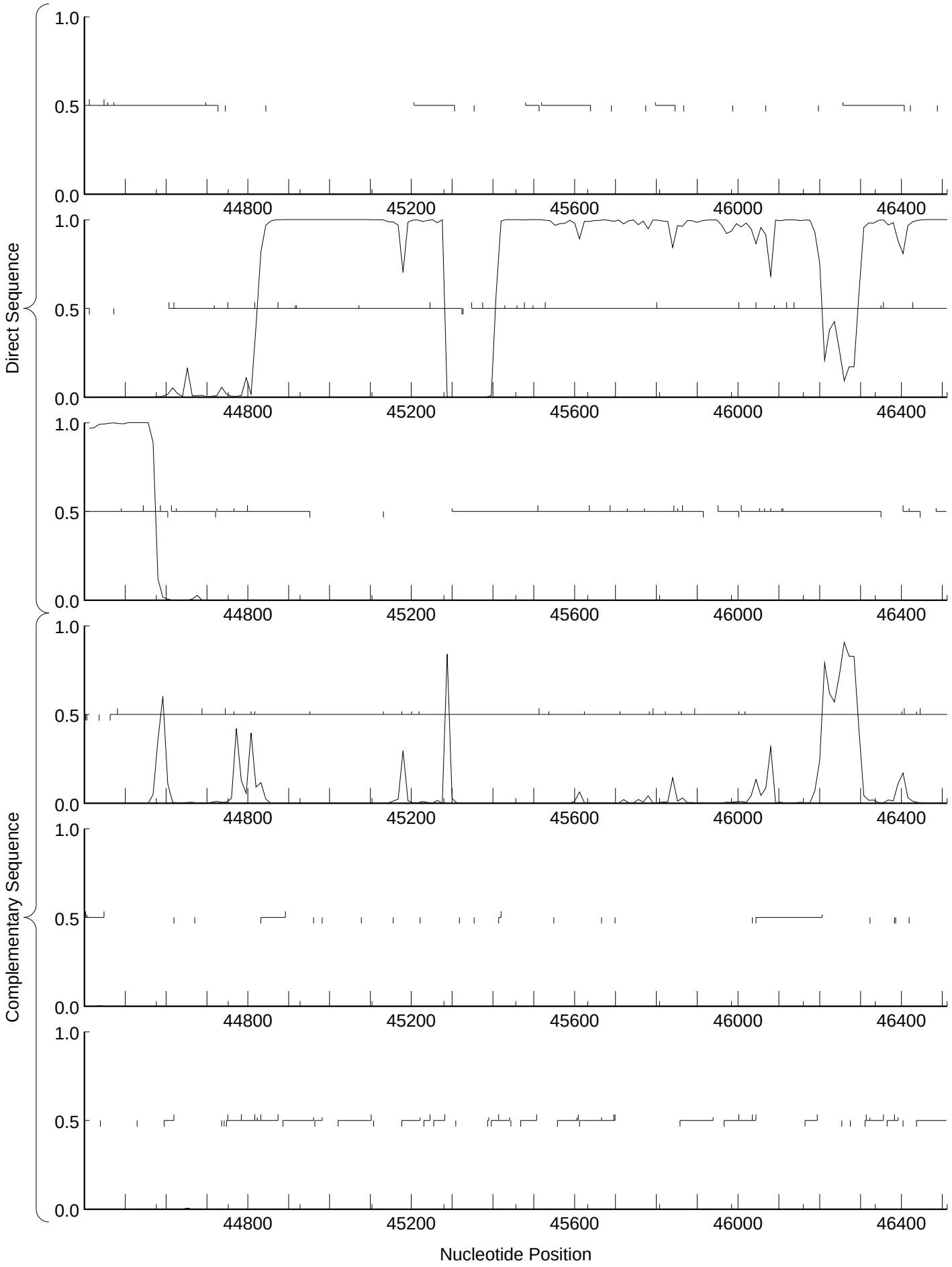
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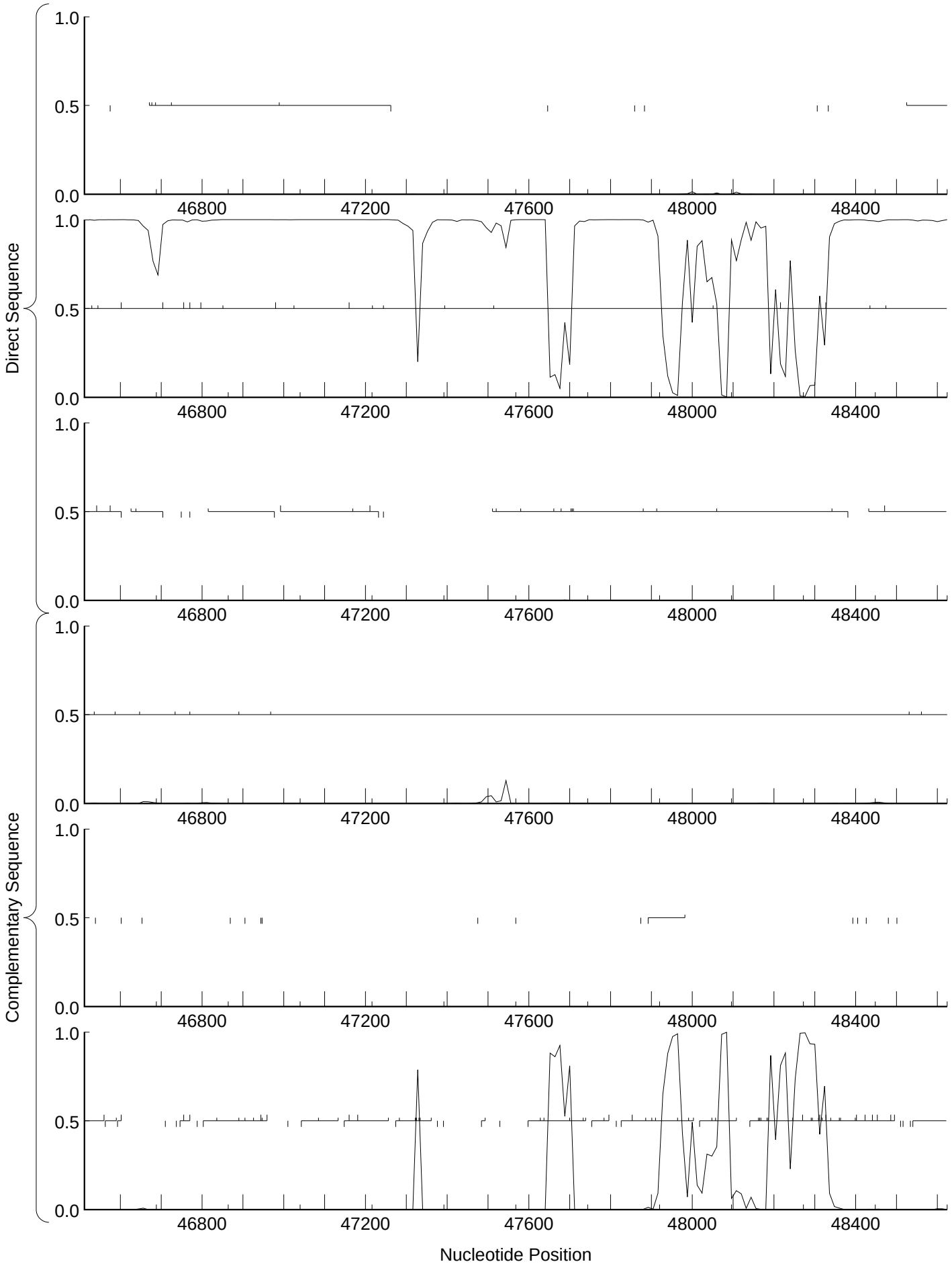
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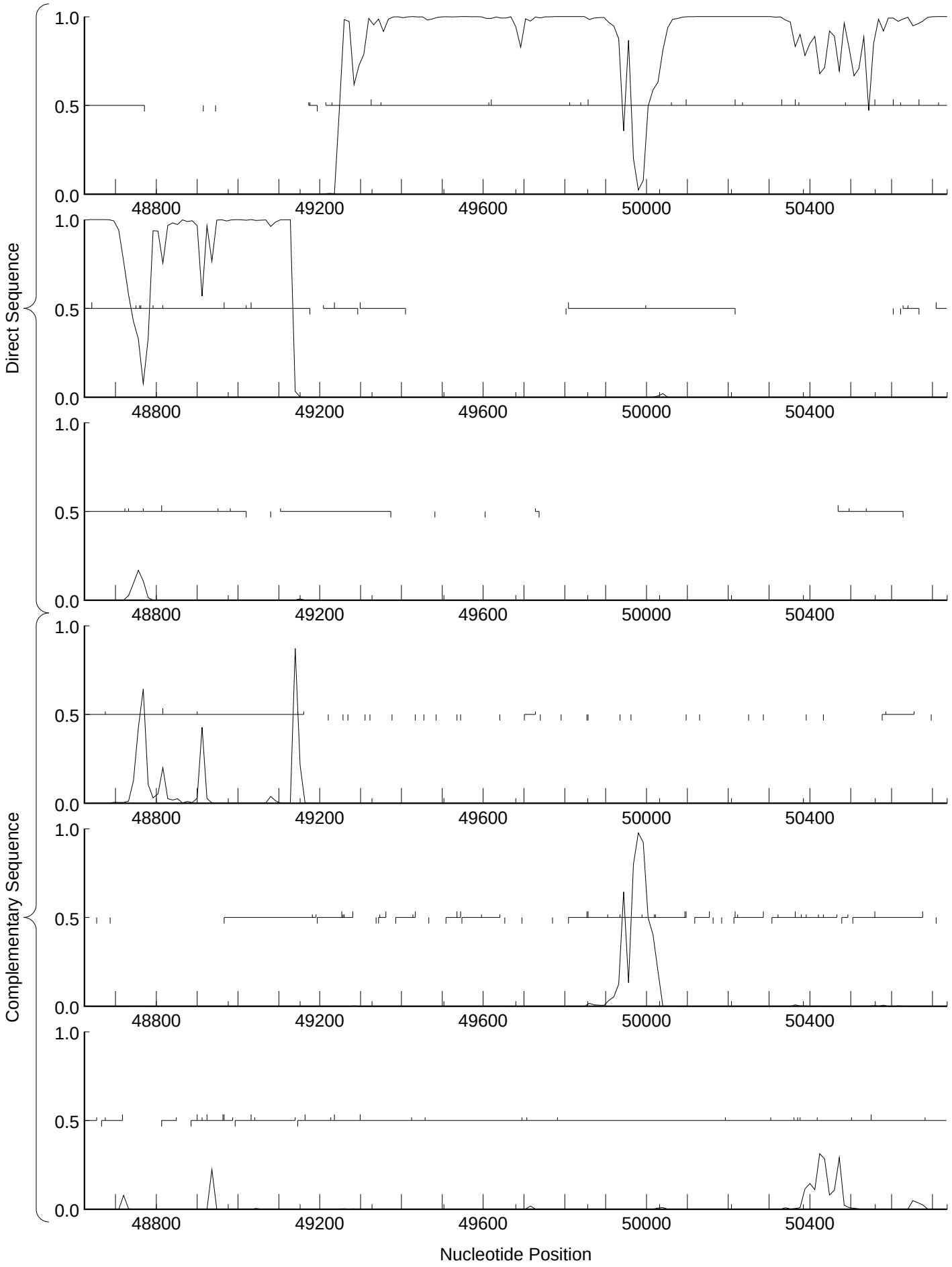
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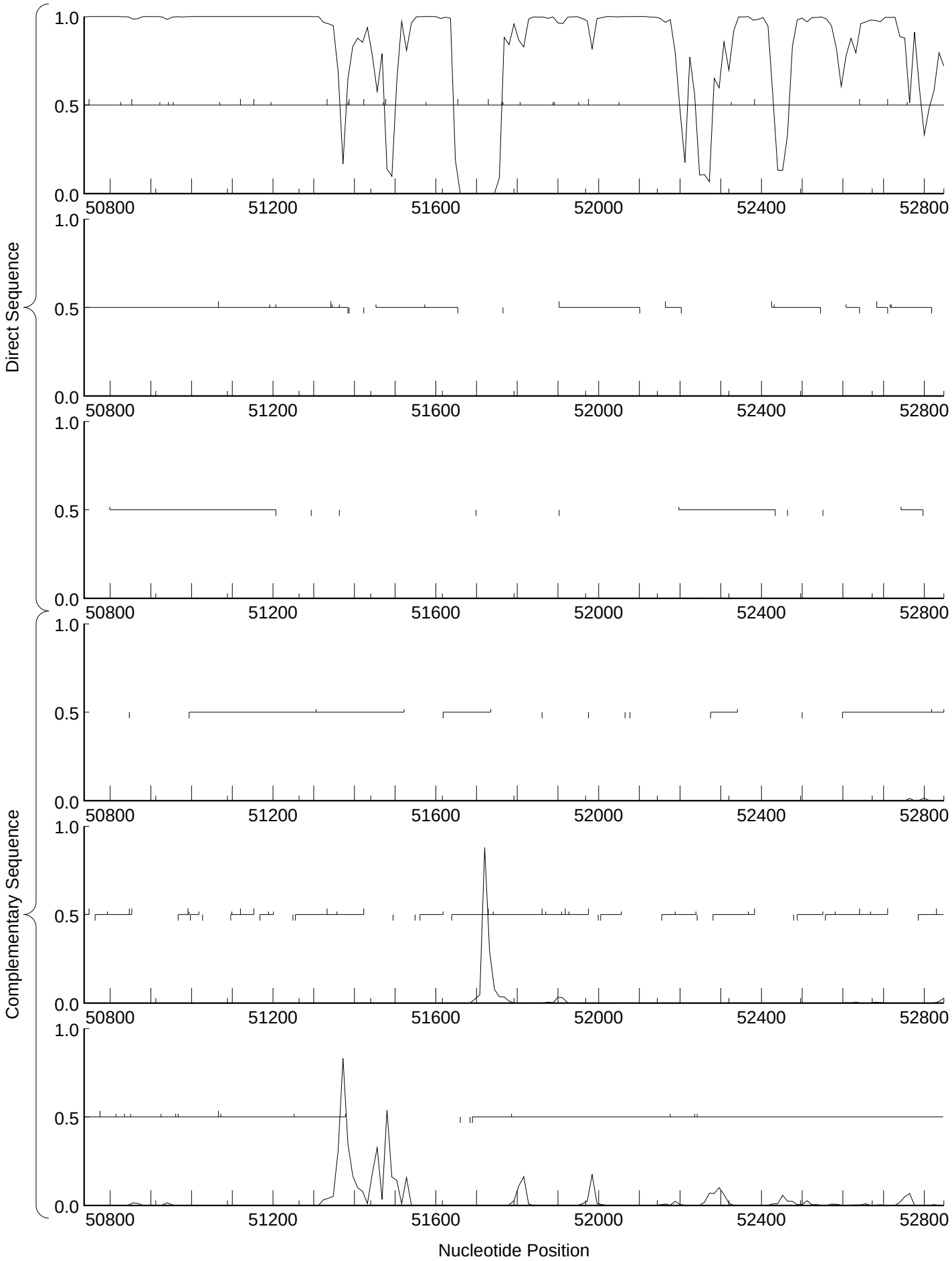
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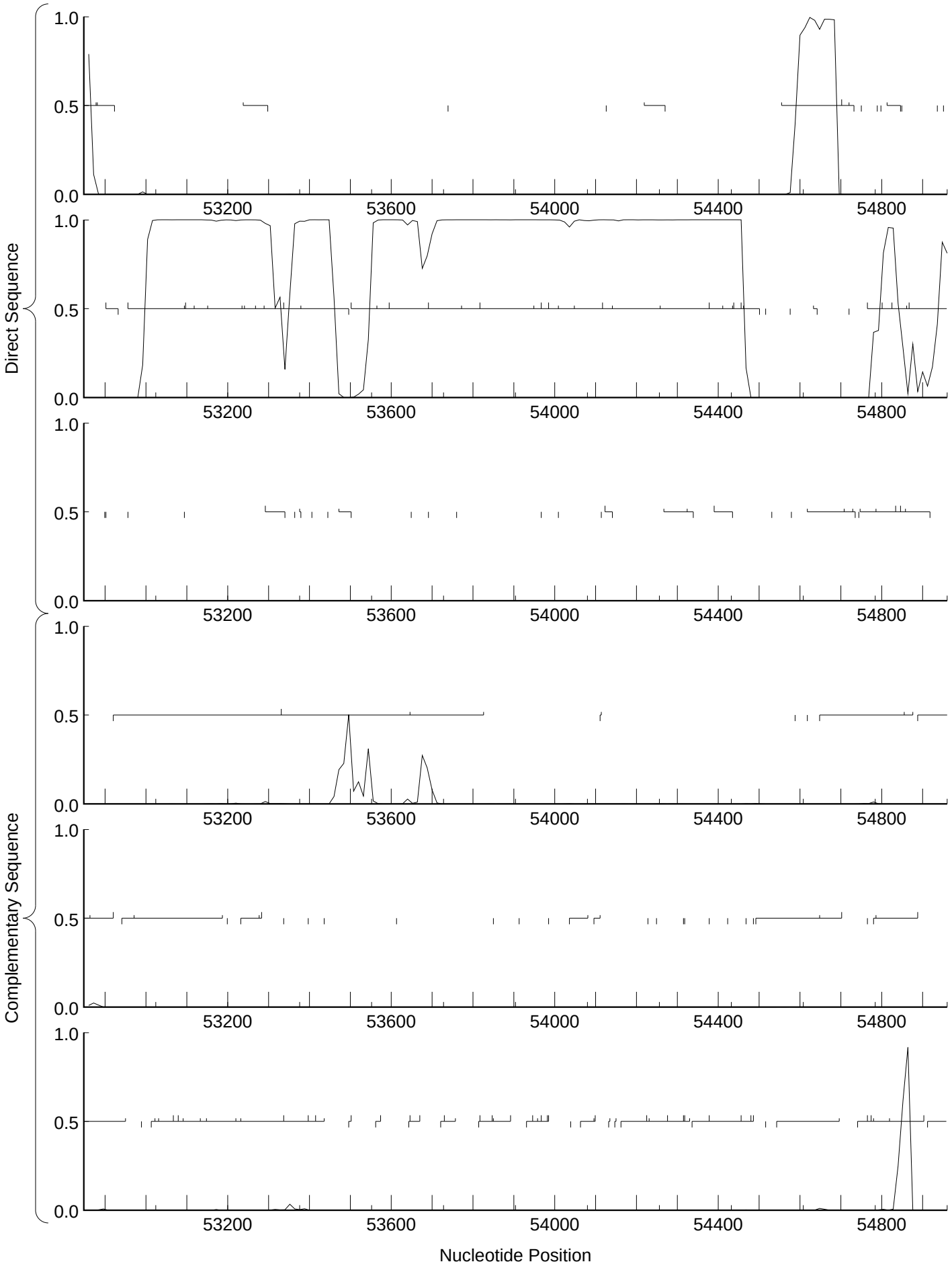
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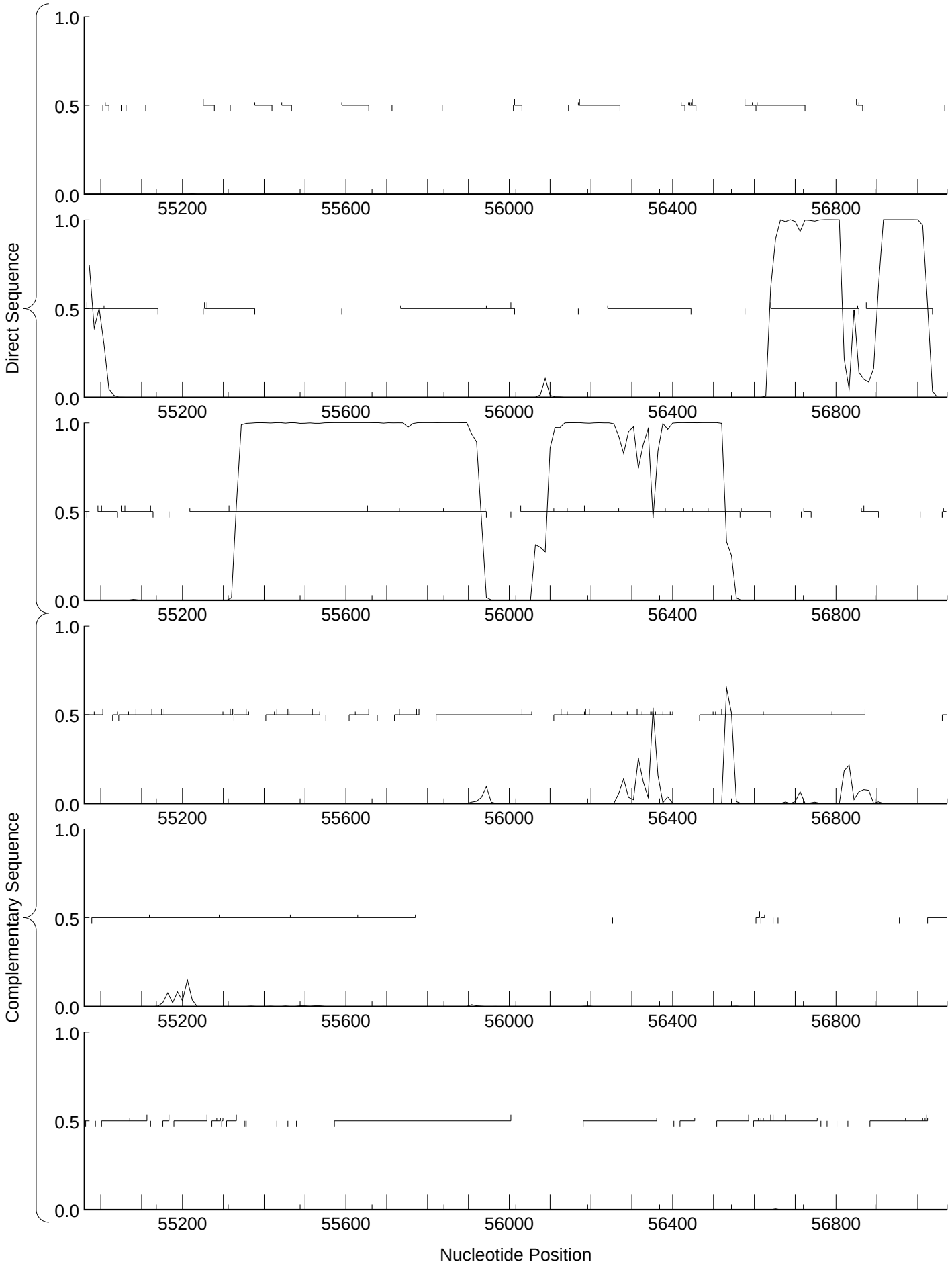
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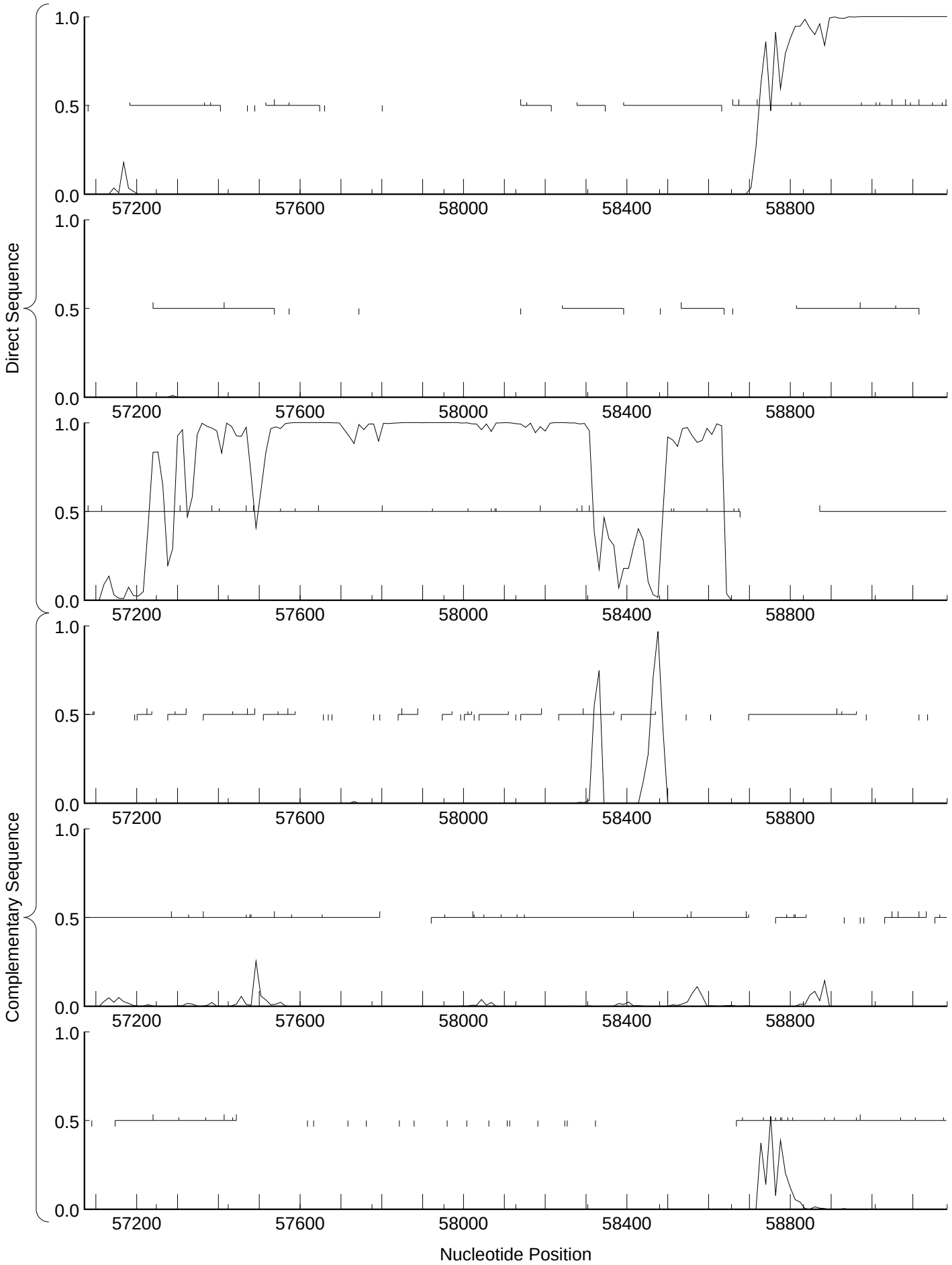
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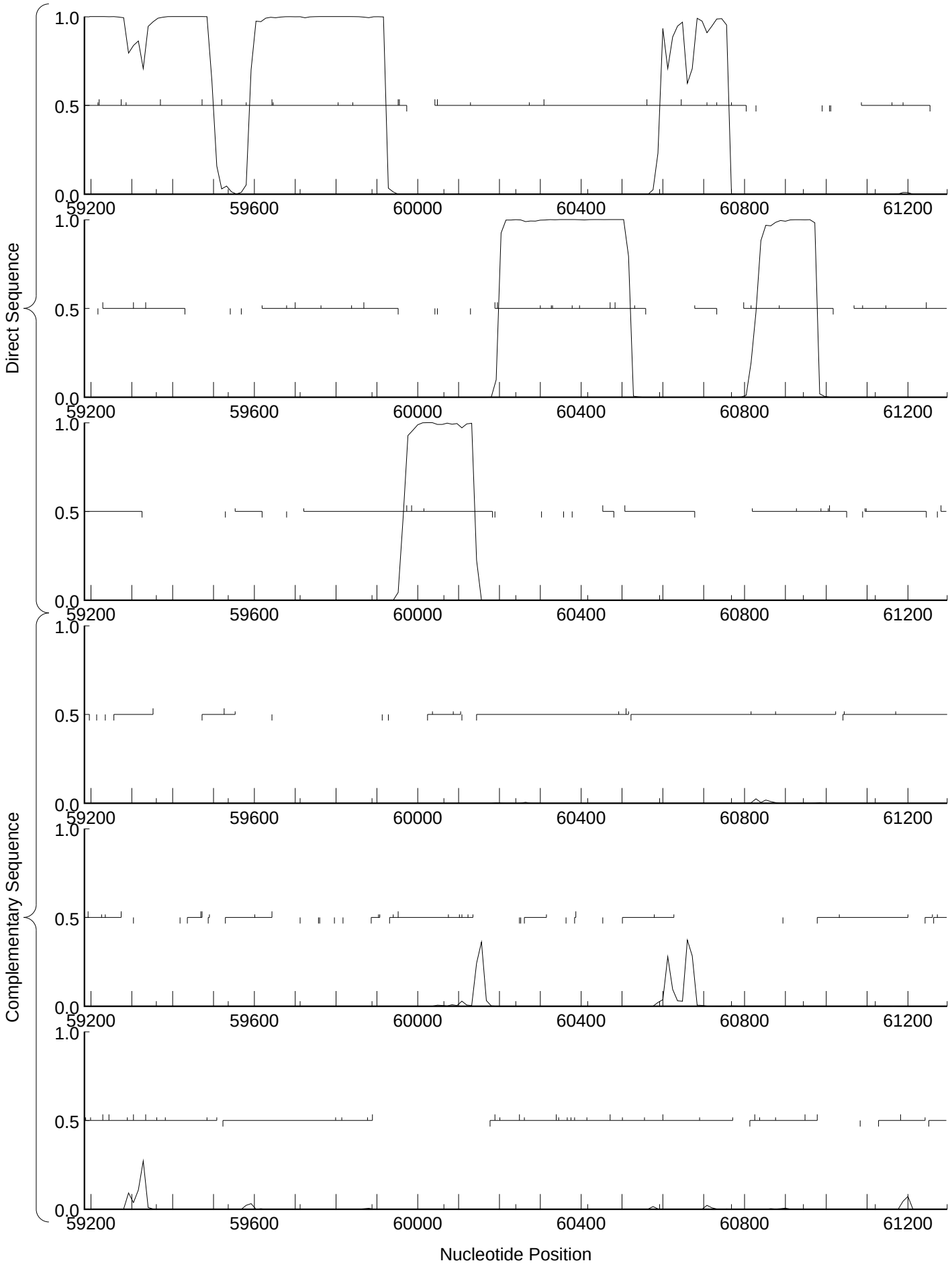
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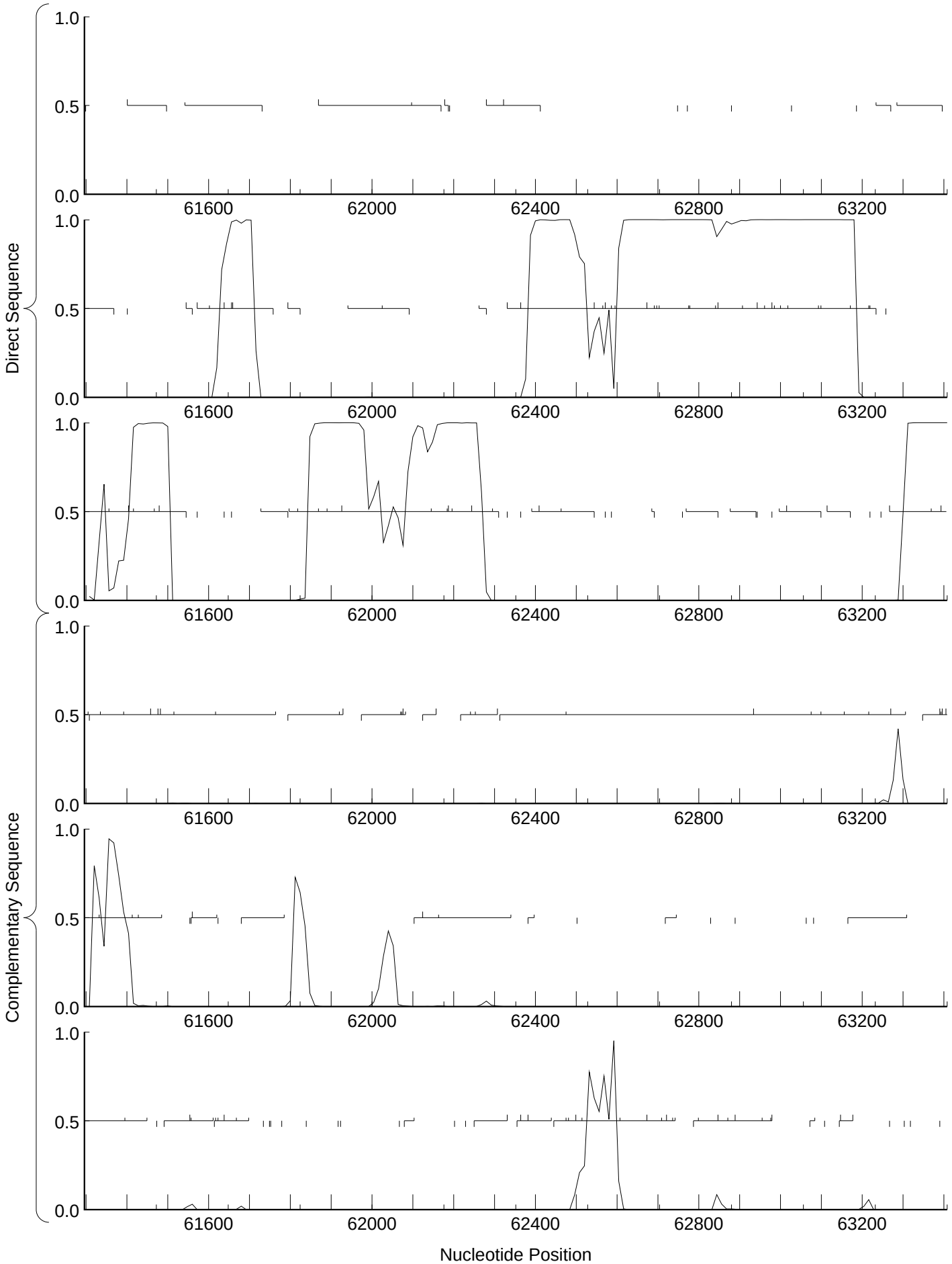
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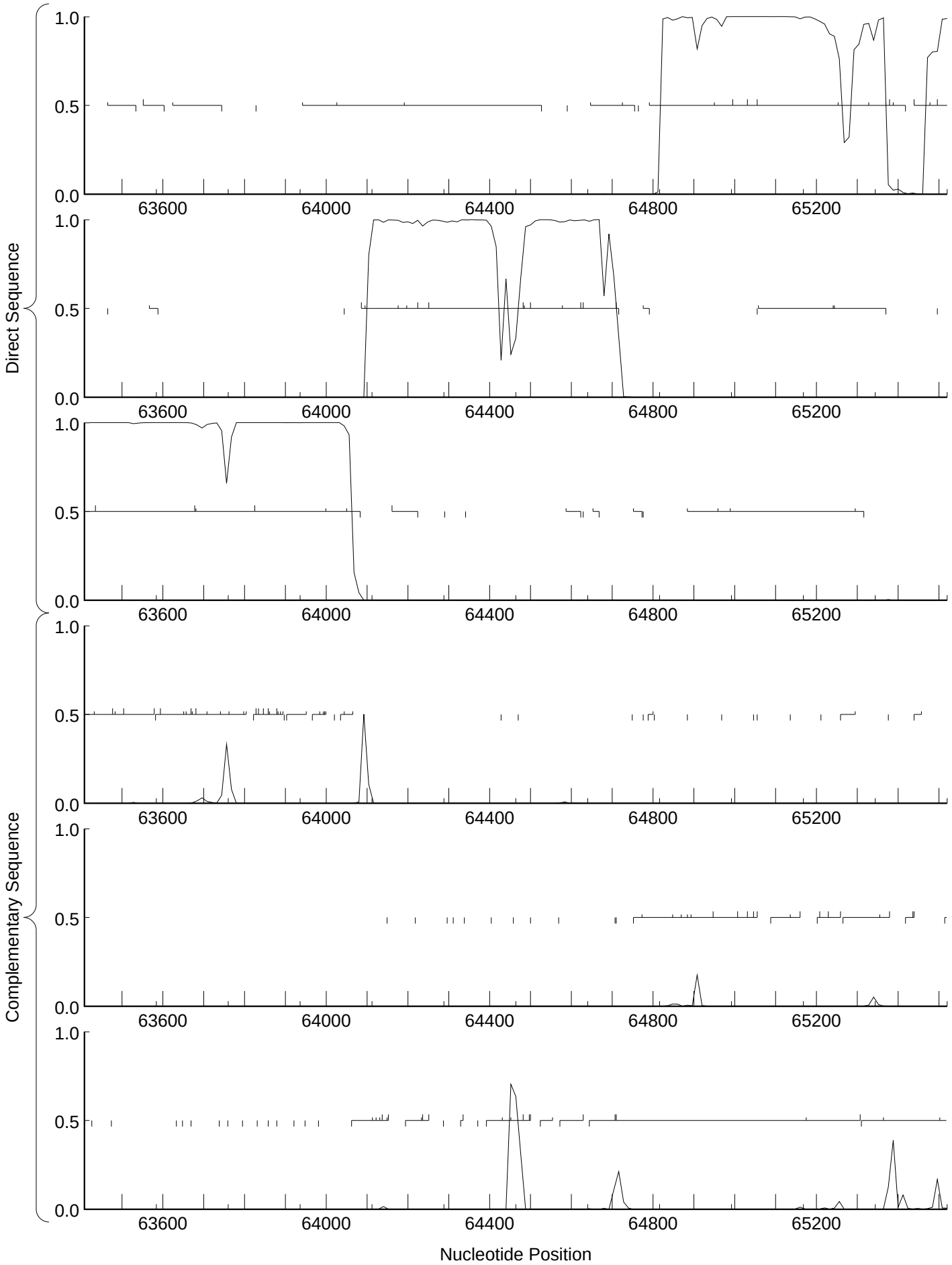
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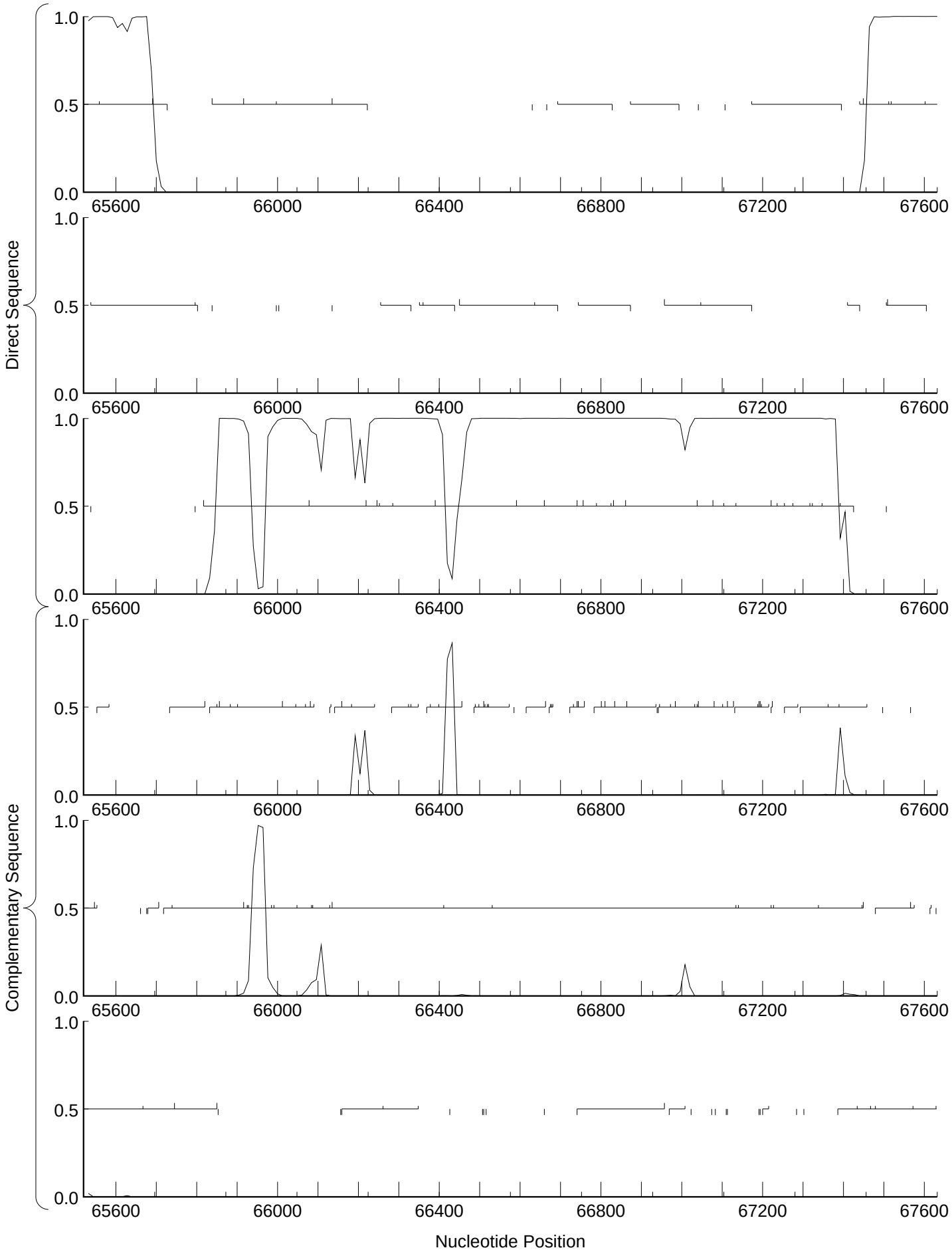
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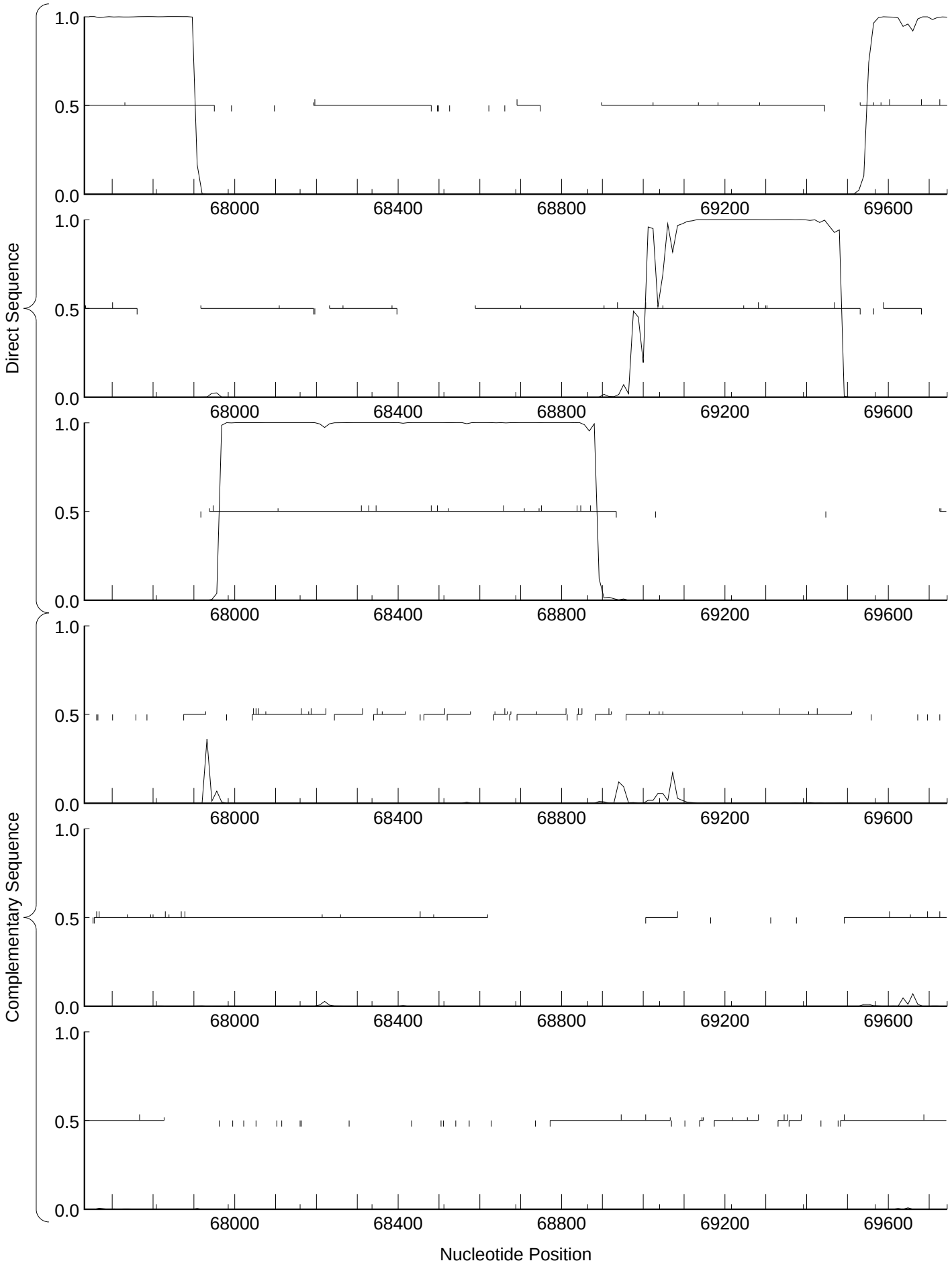
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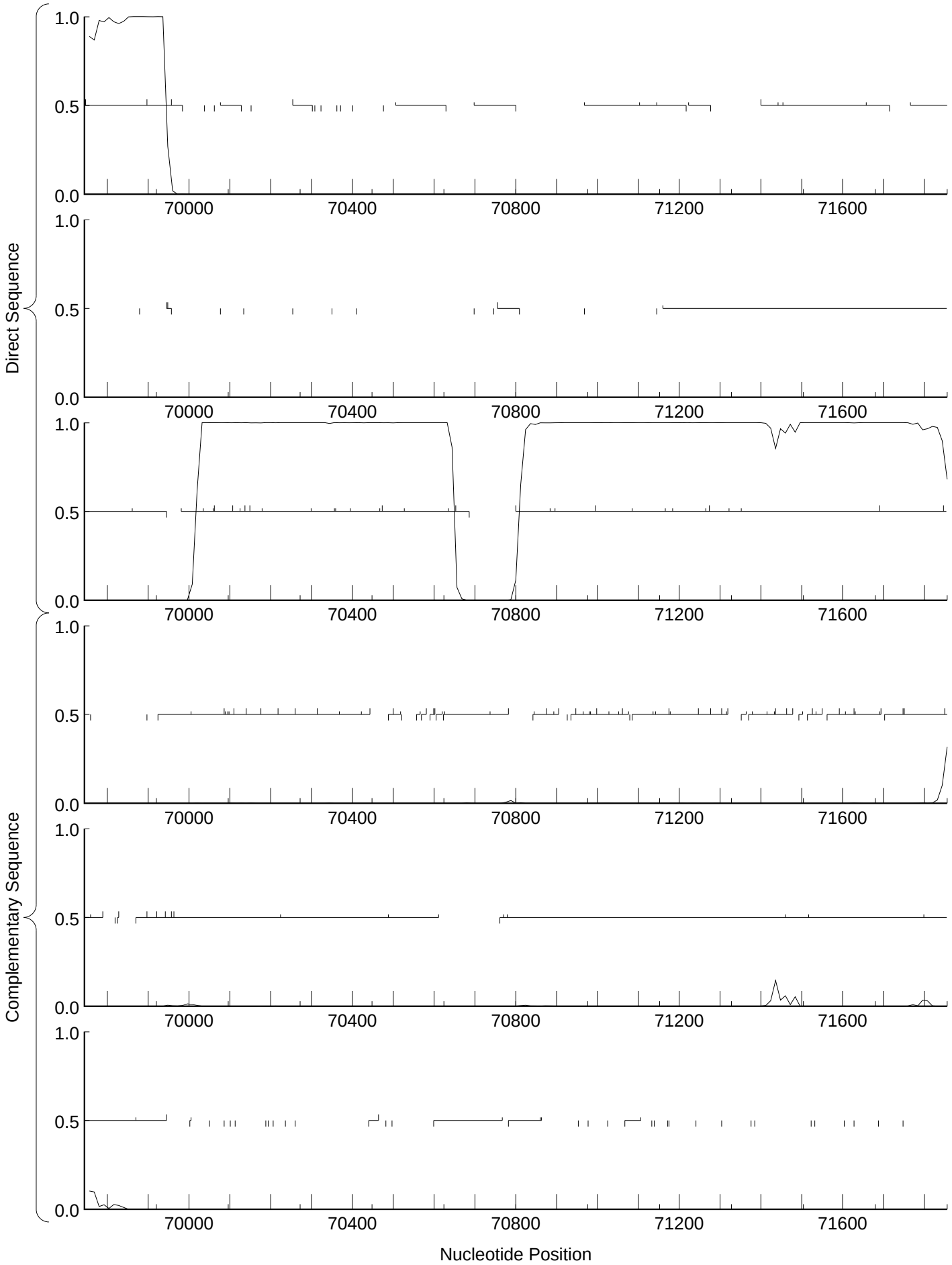
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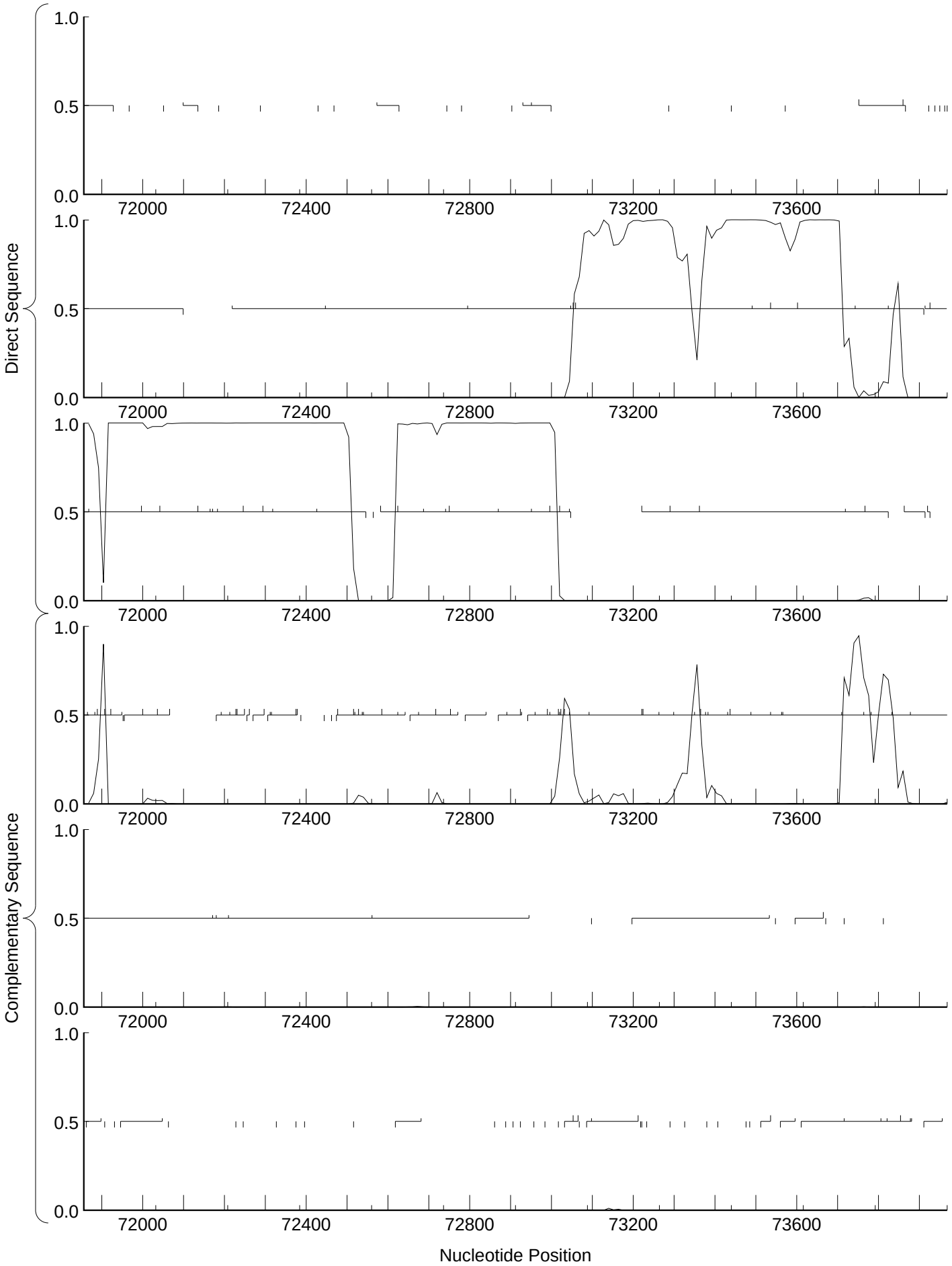
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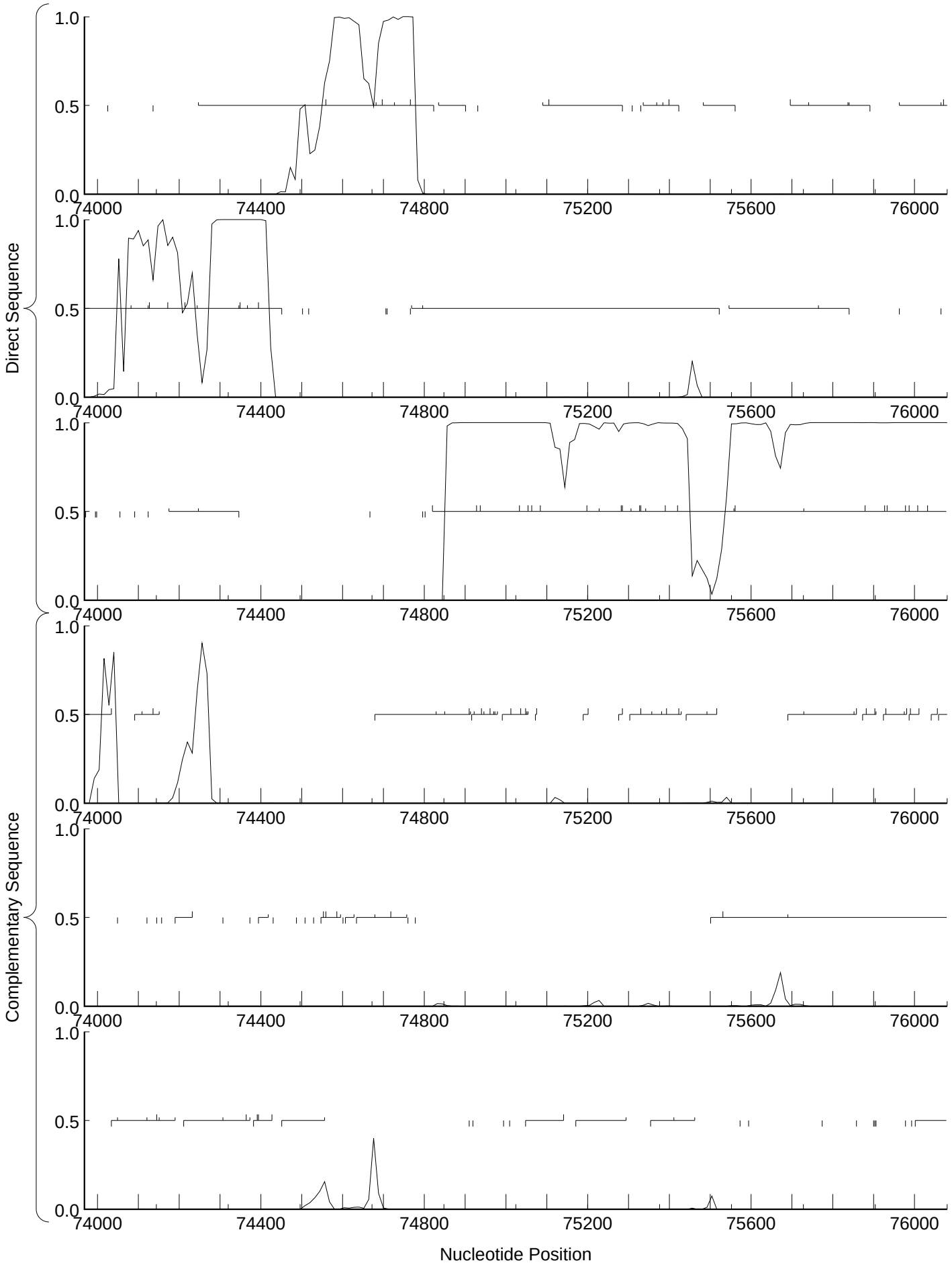
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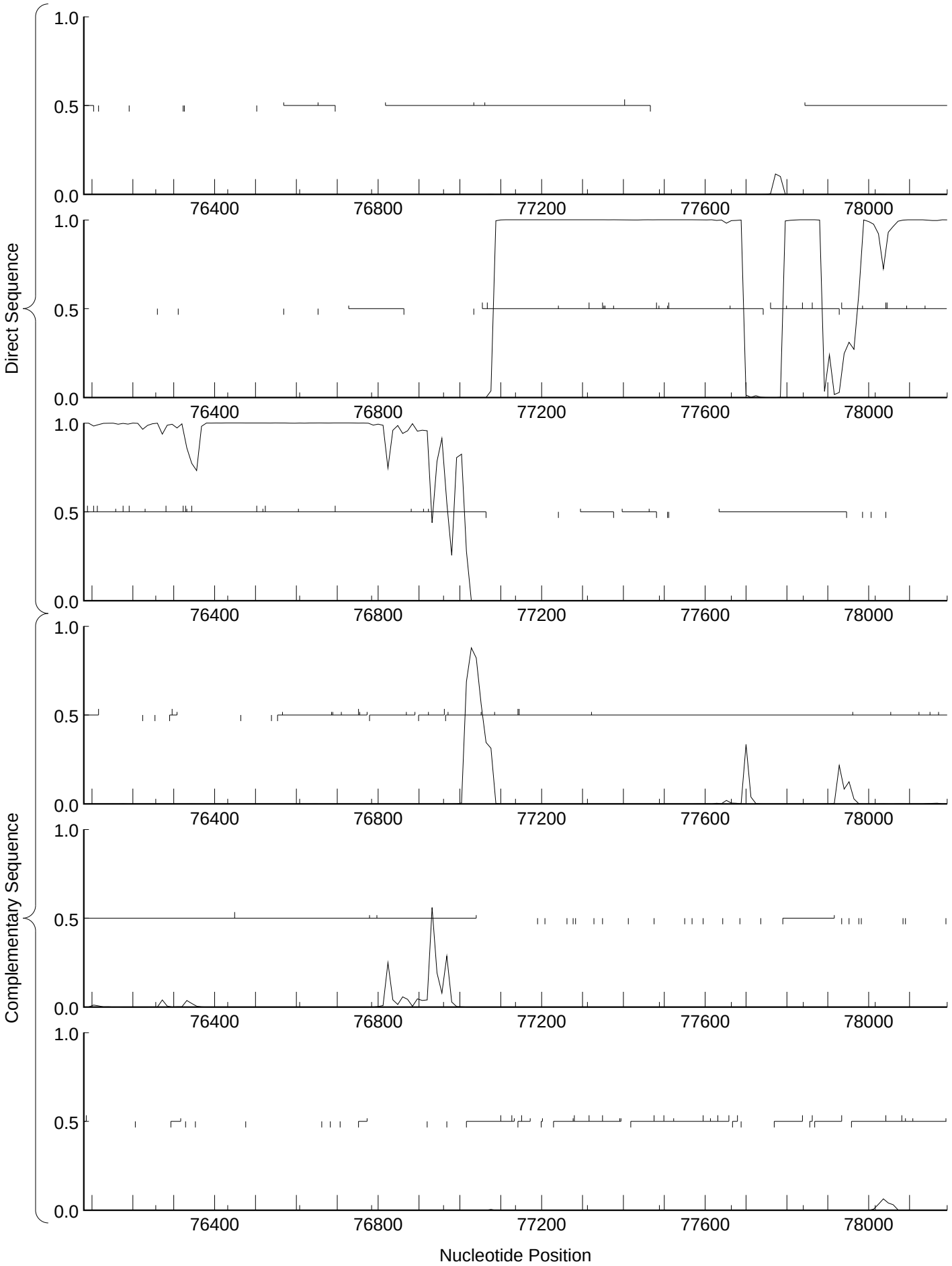
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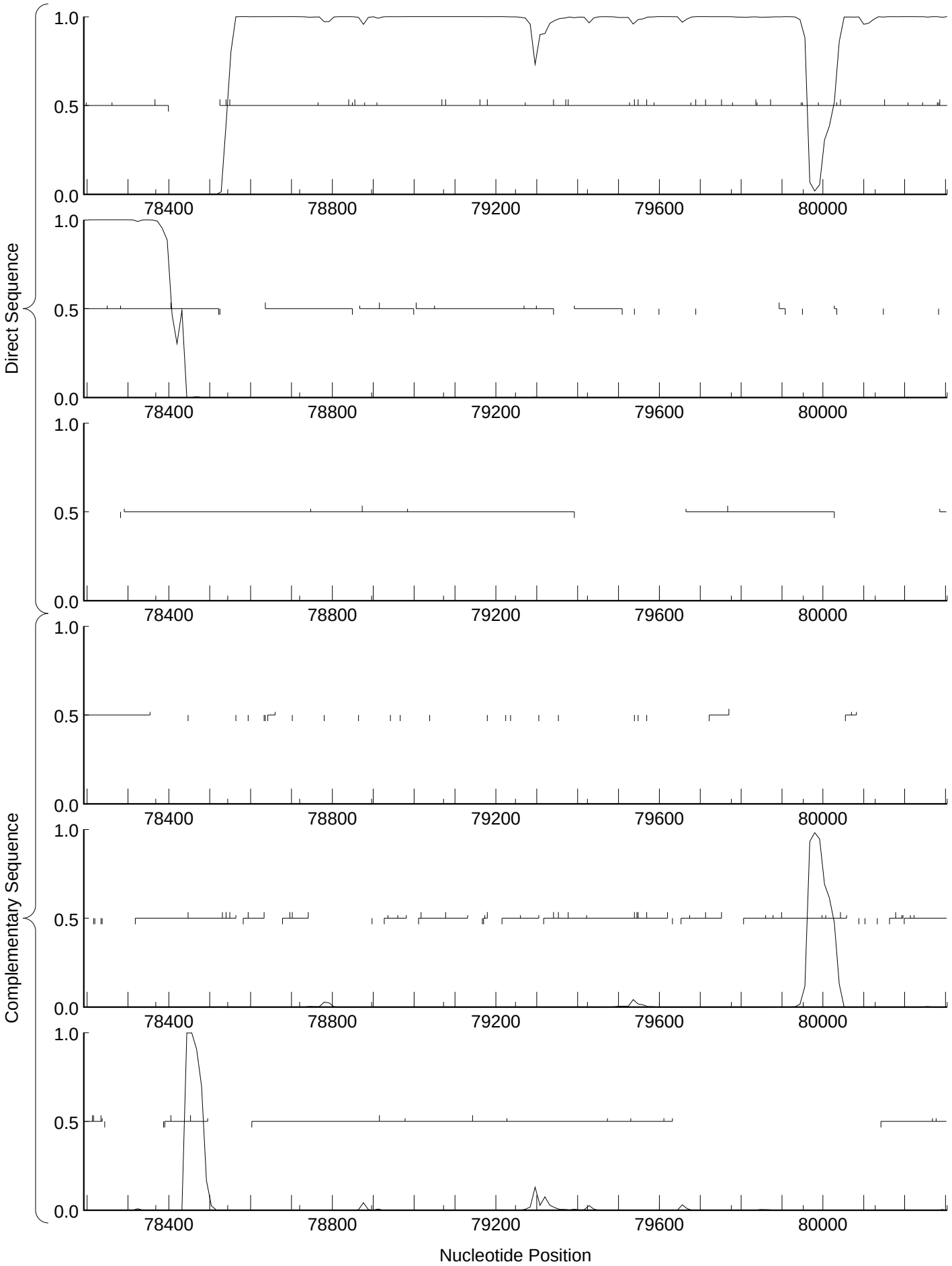
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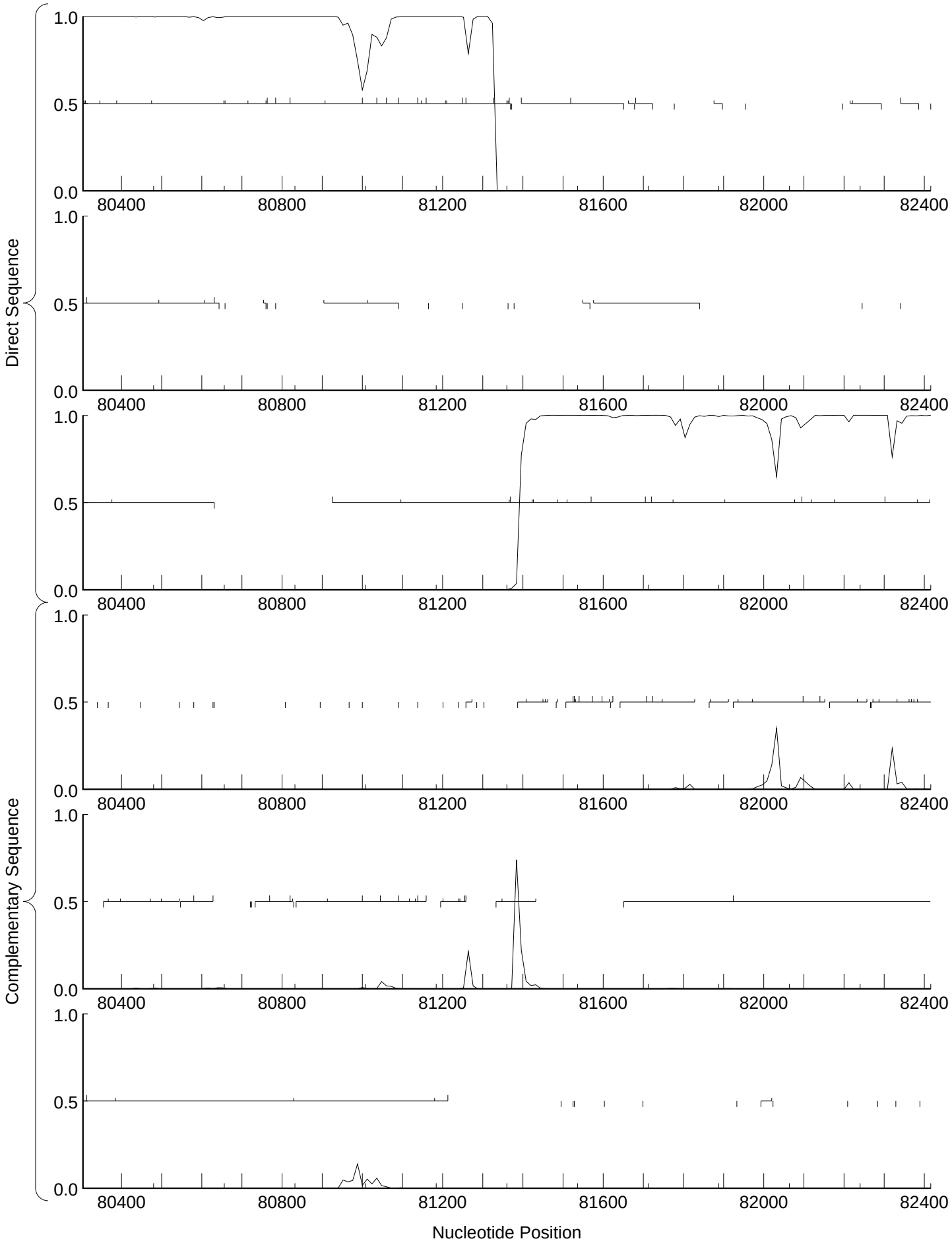
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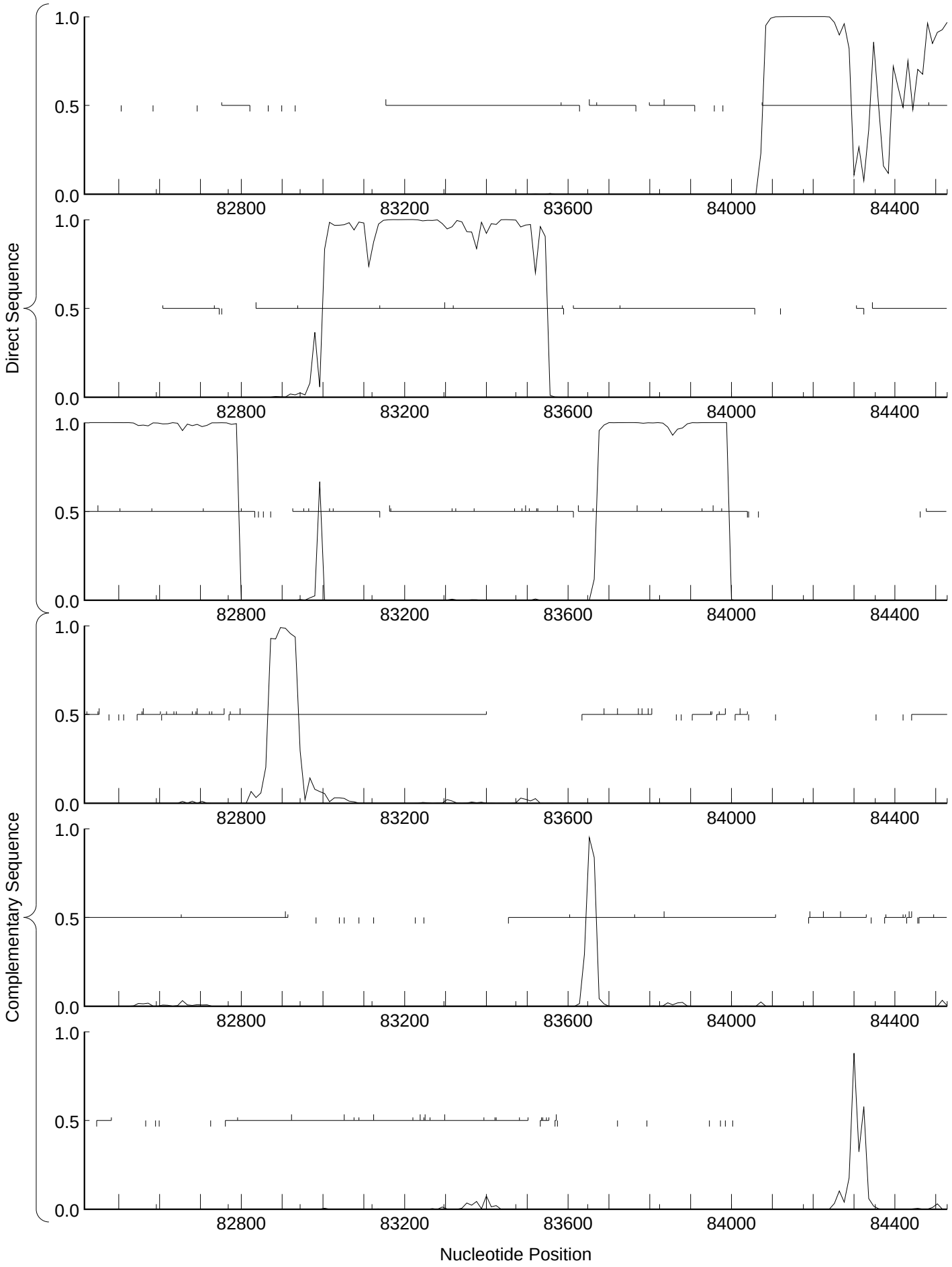
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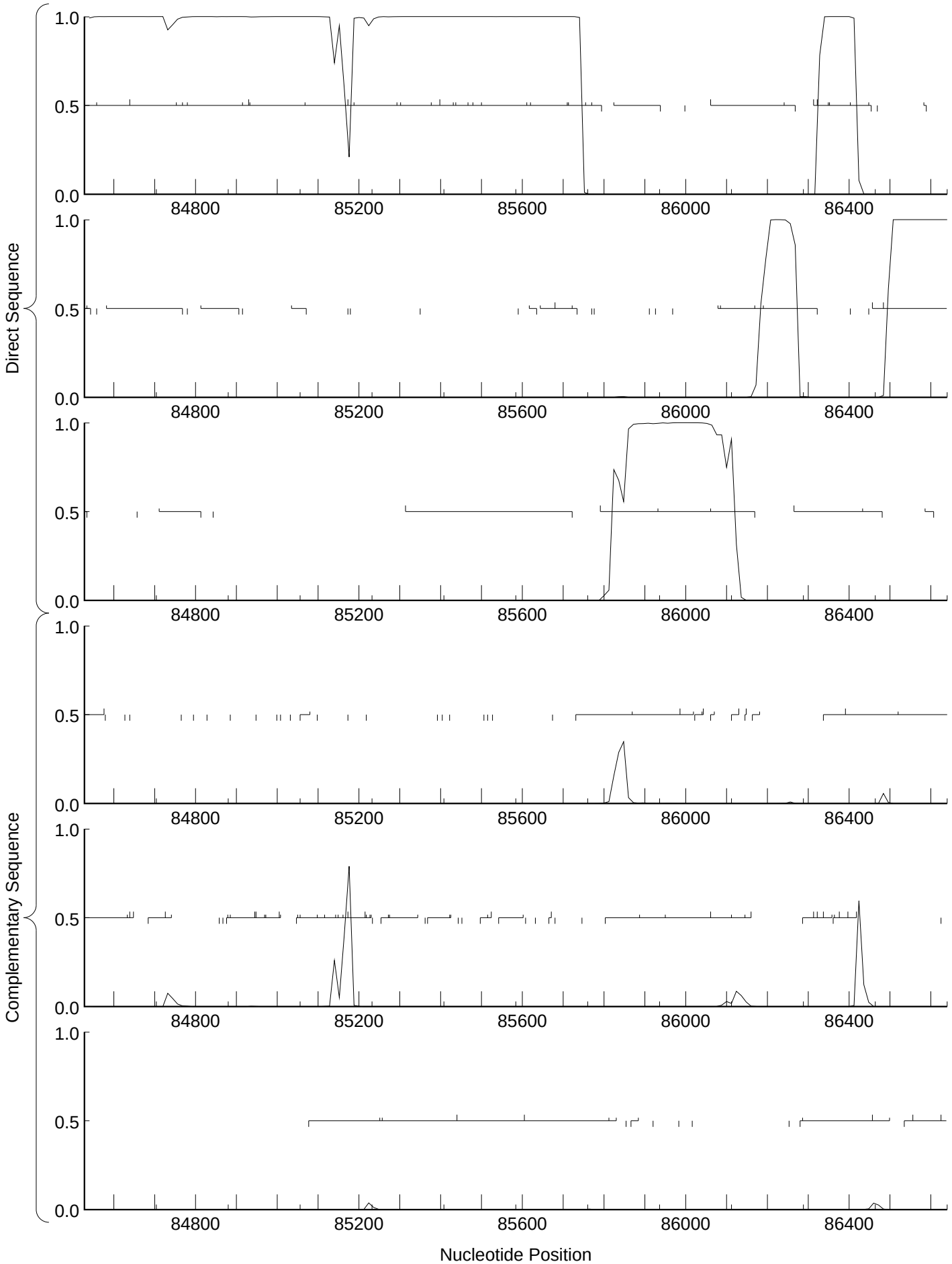
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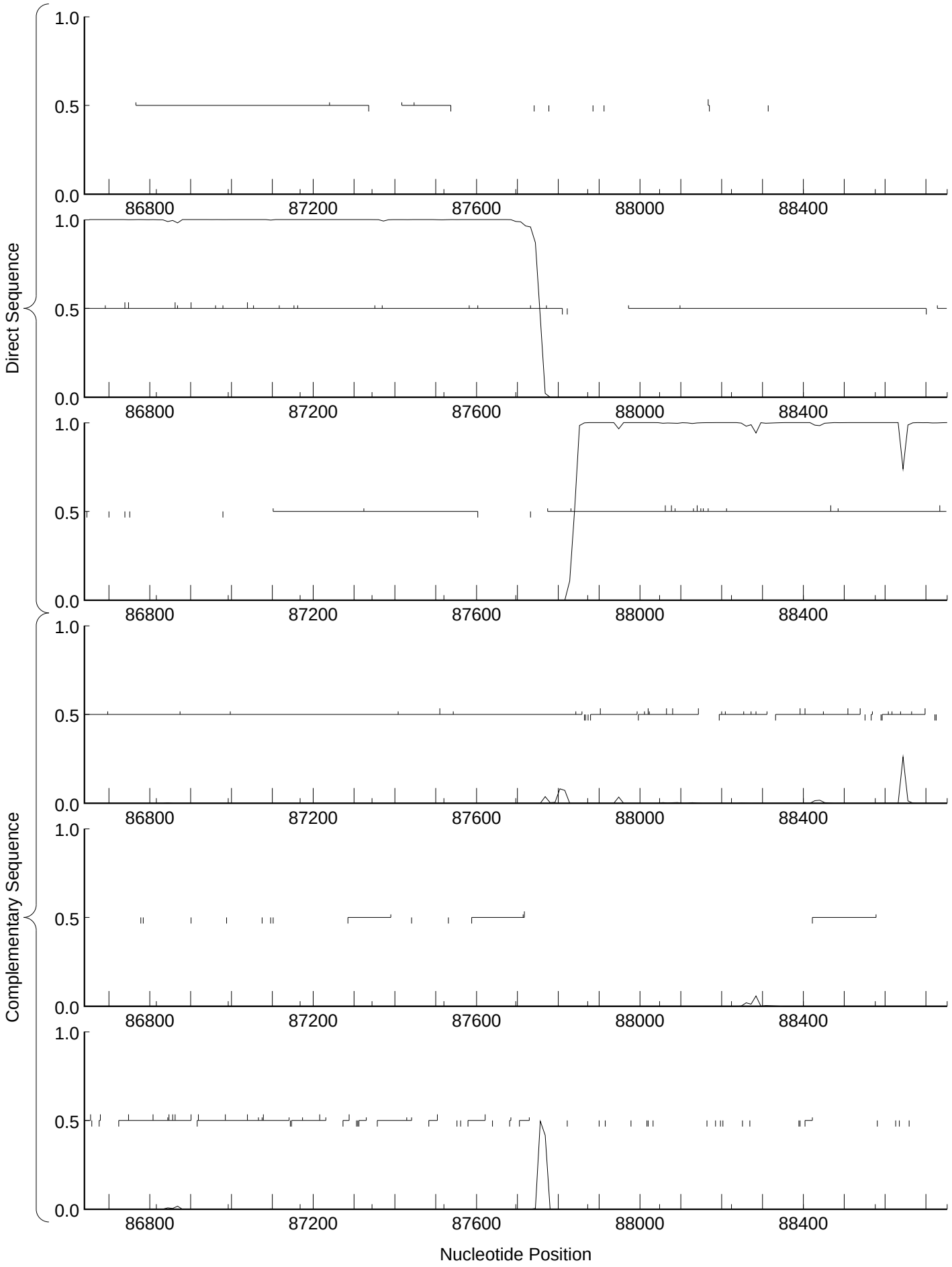
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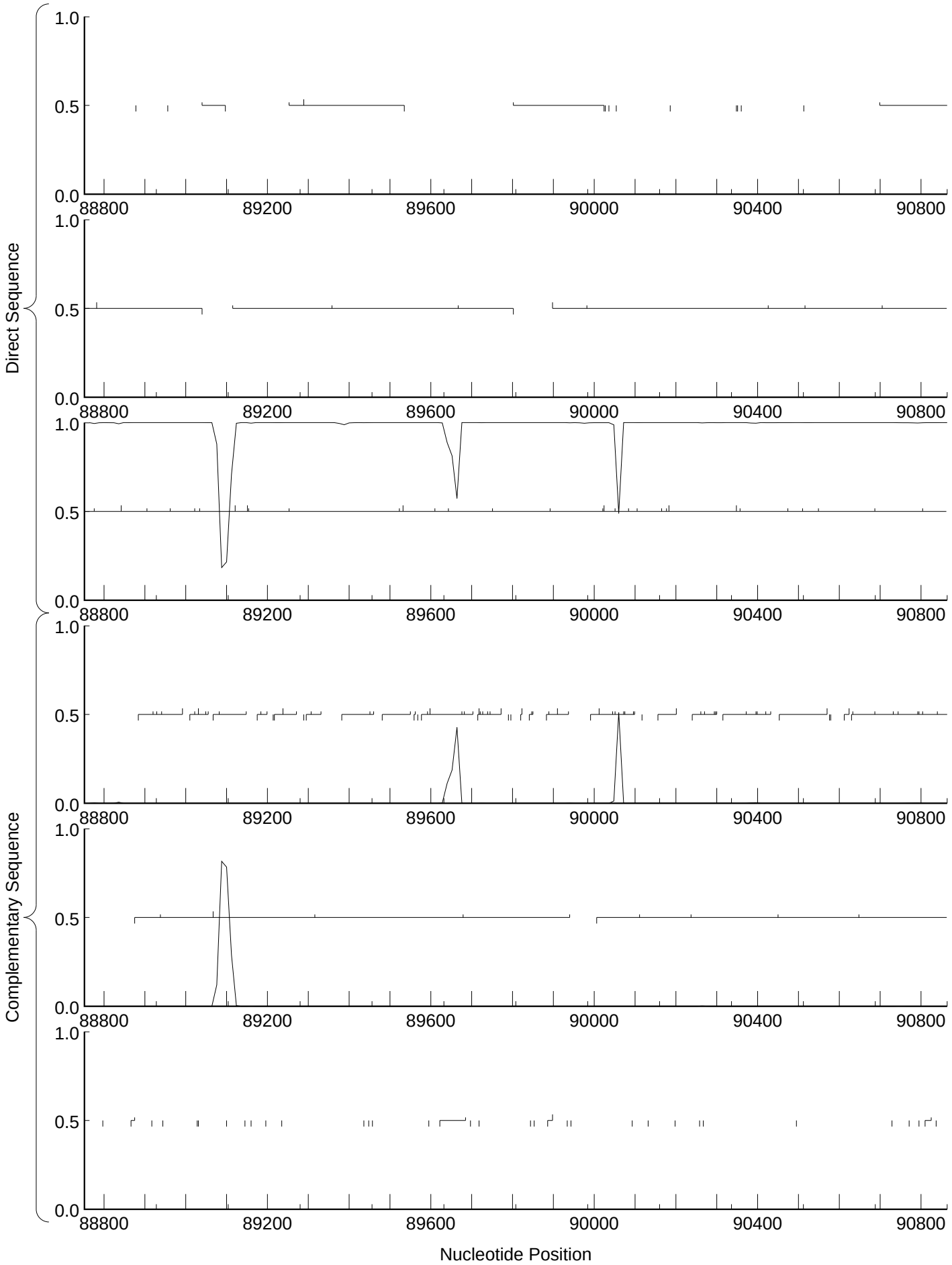
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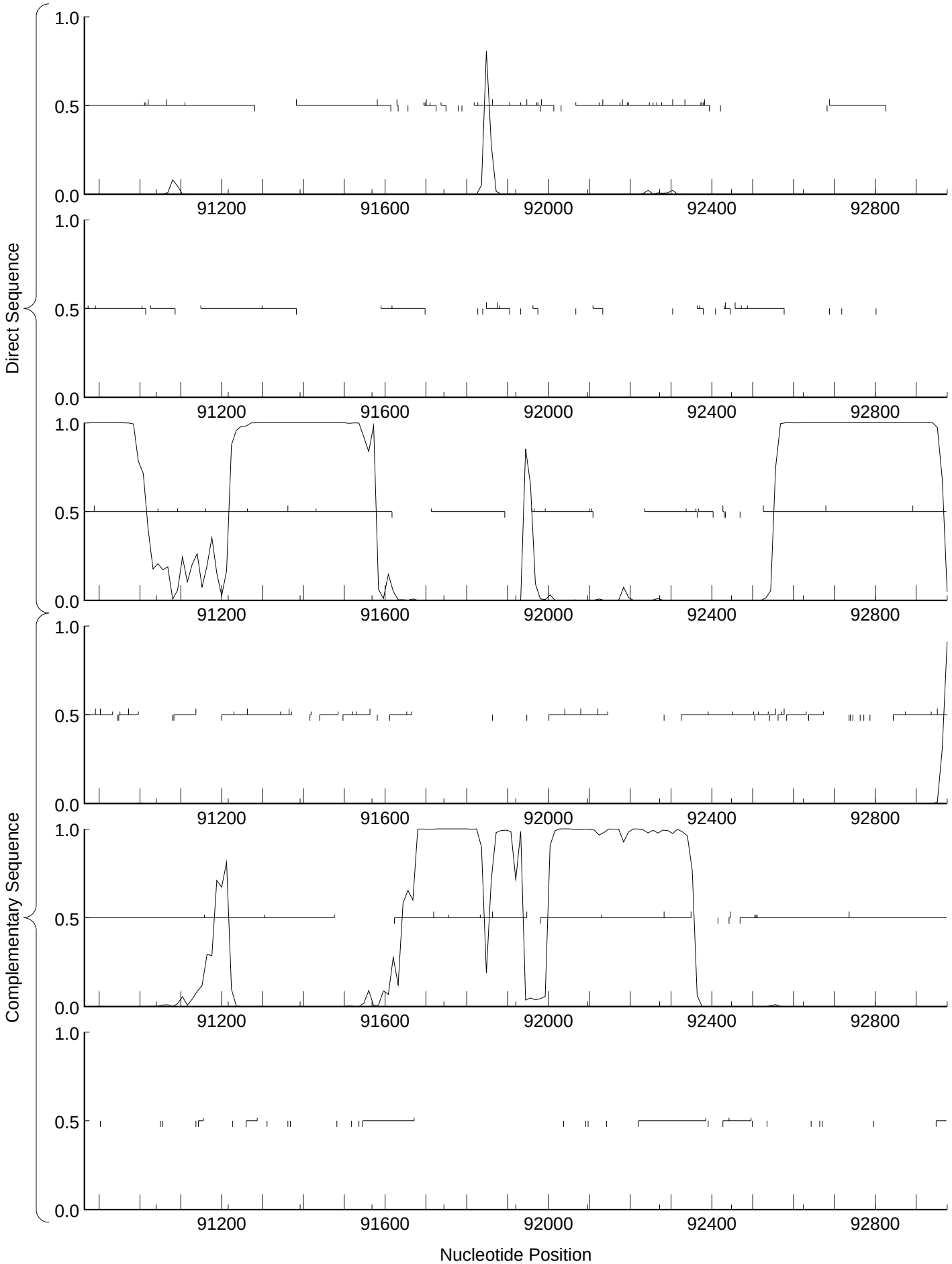
cobacterium phage Tonenili complete sequence, 160985 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 43/78



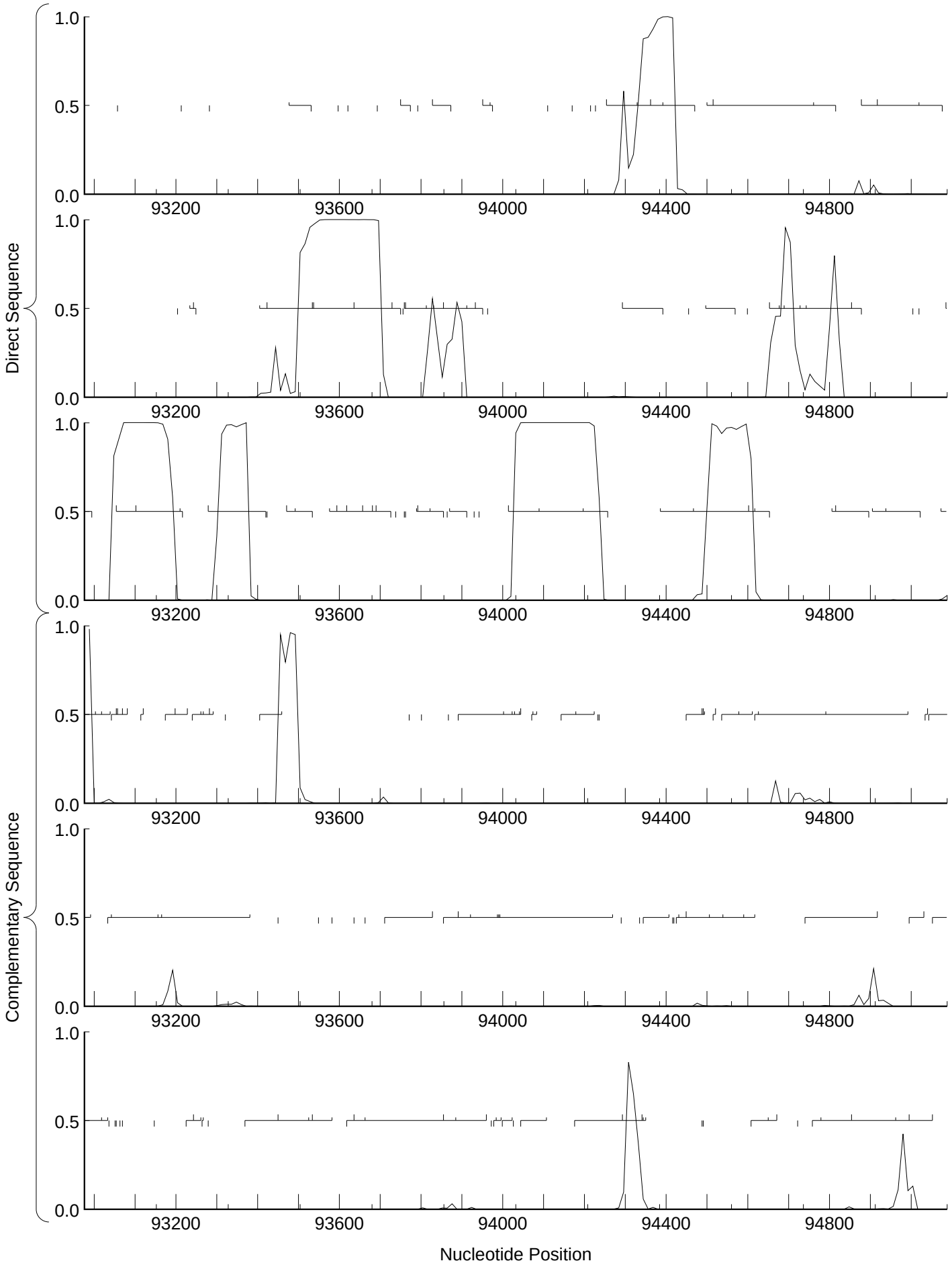
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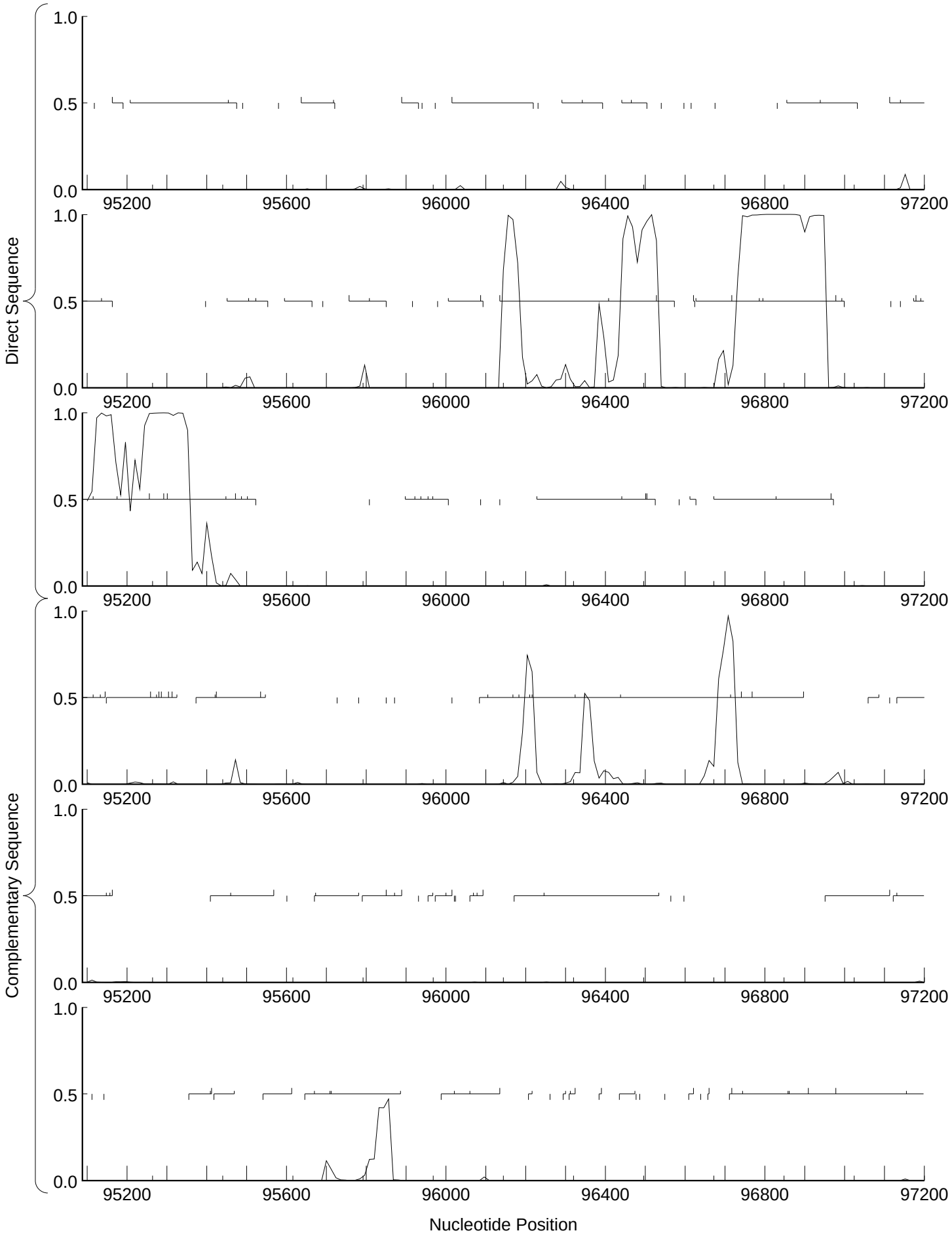
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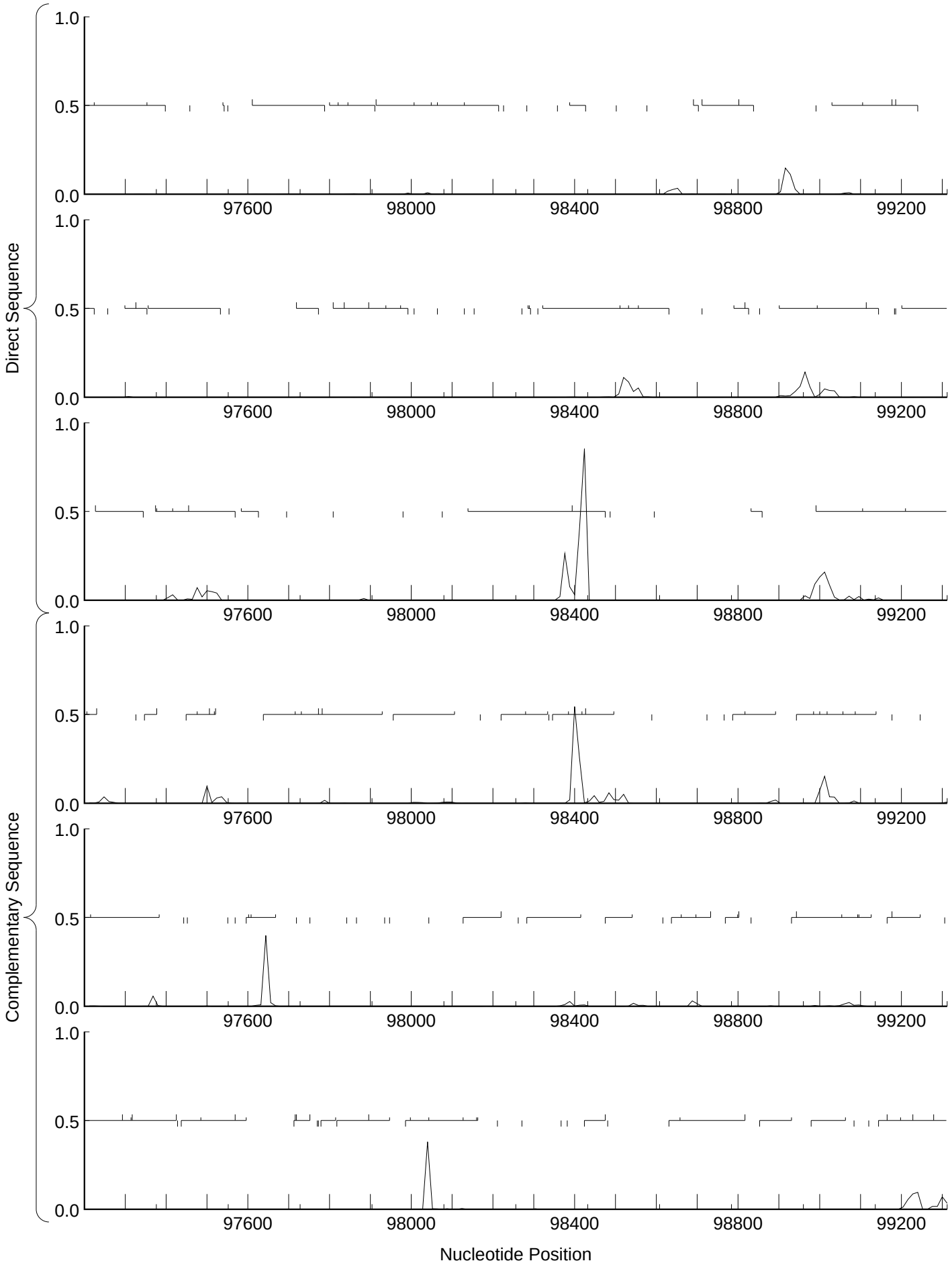
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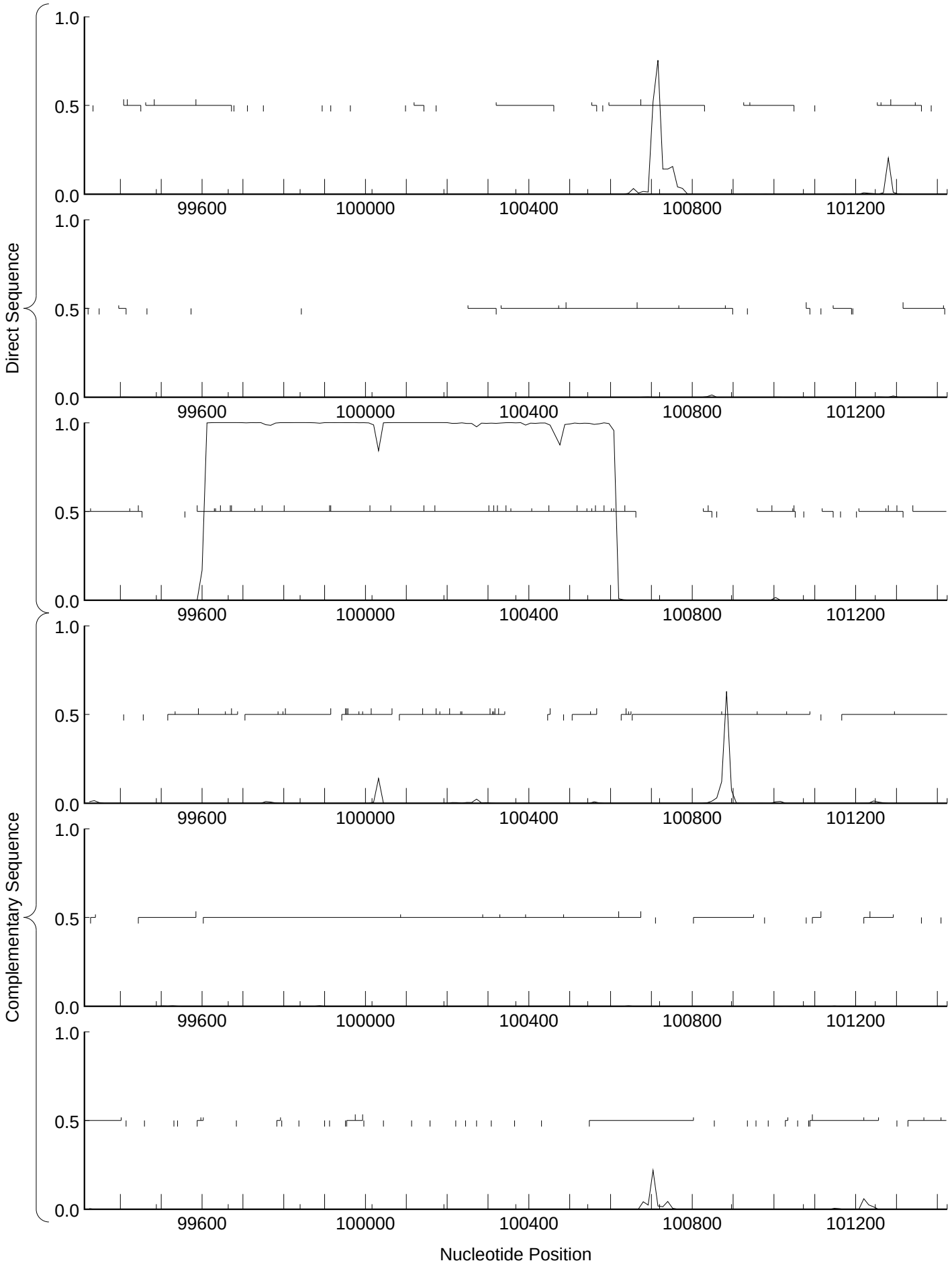
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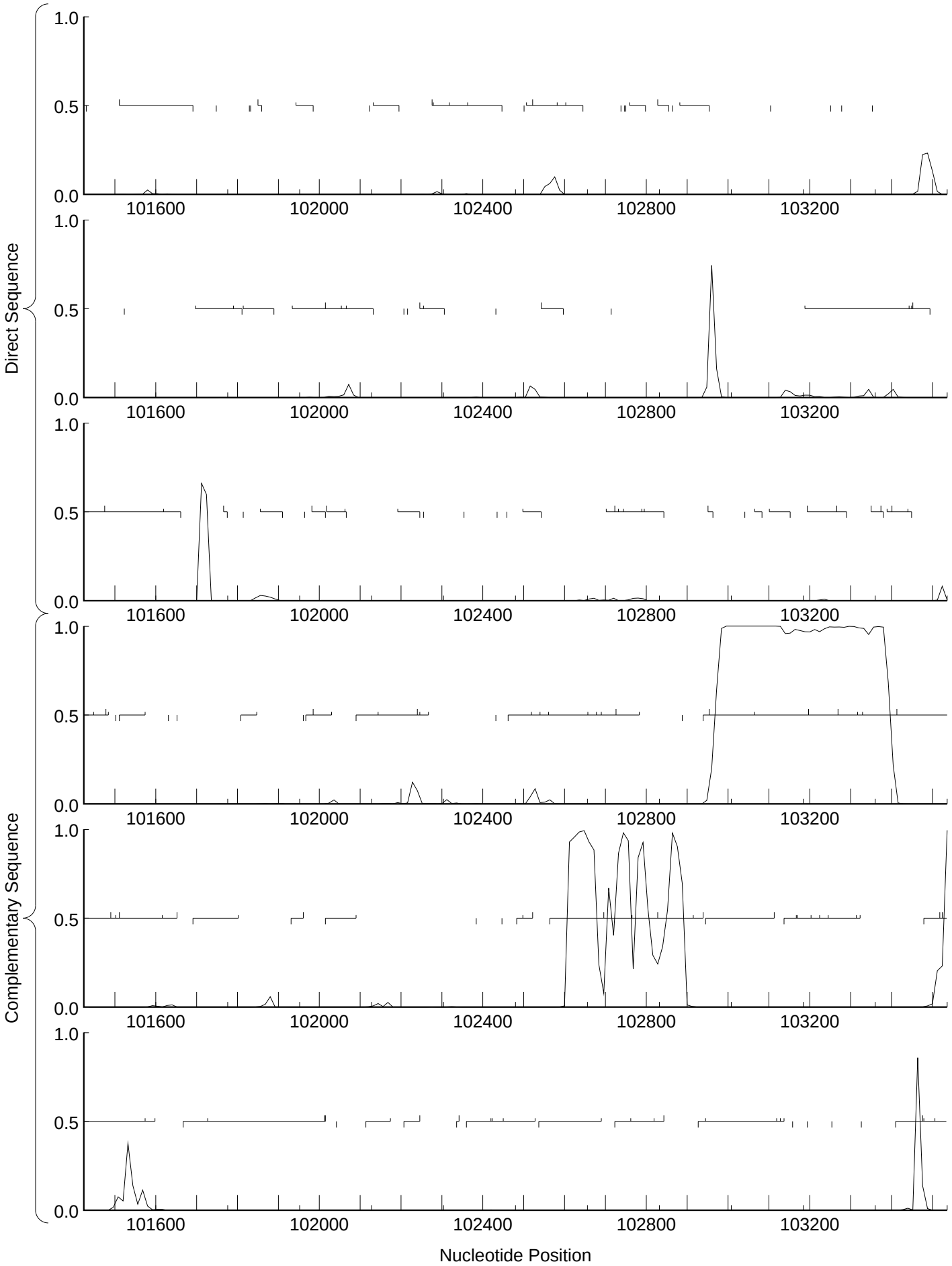
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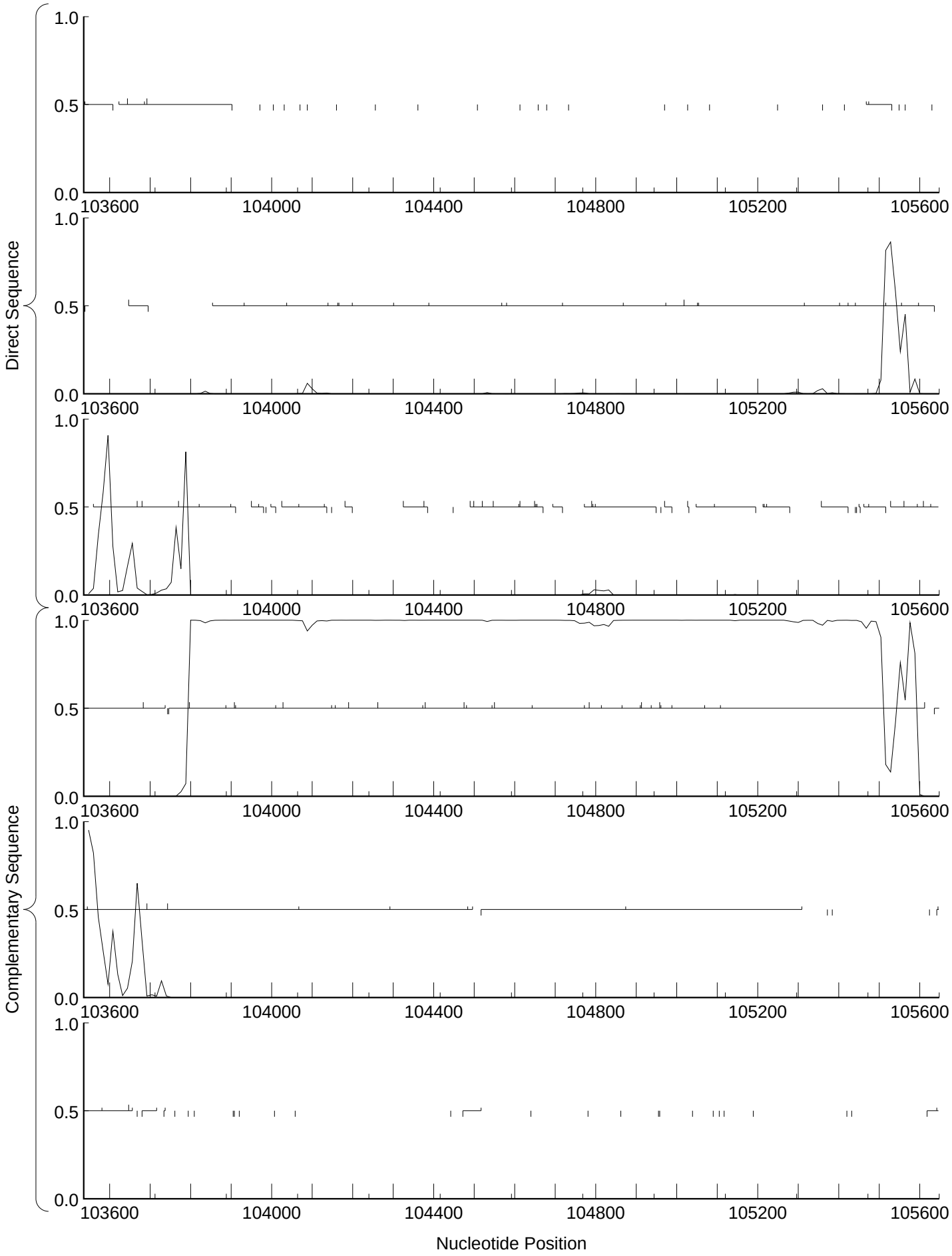
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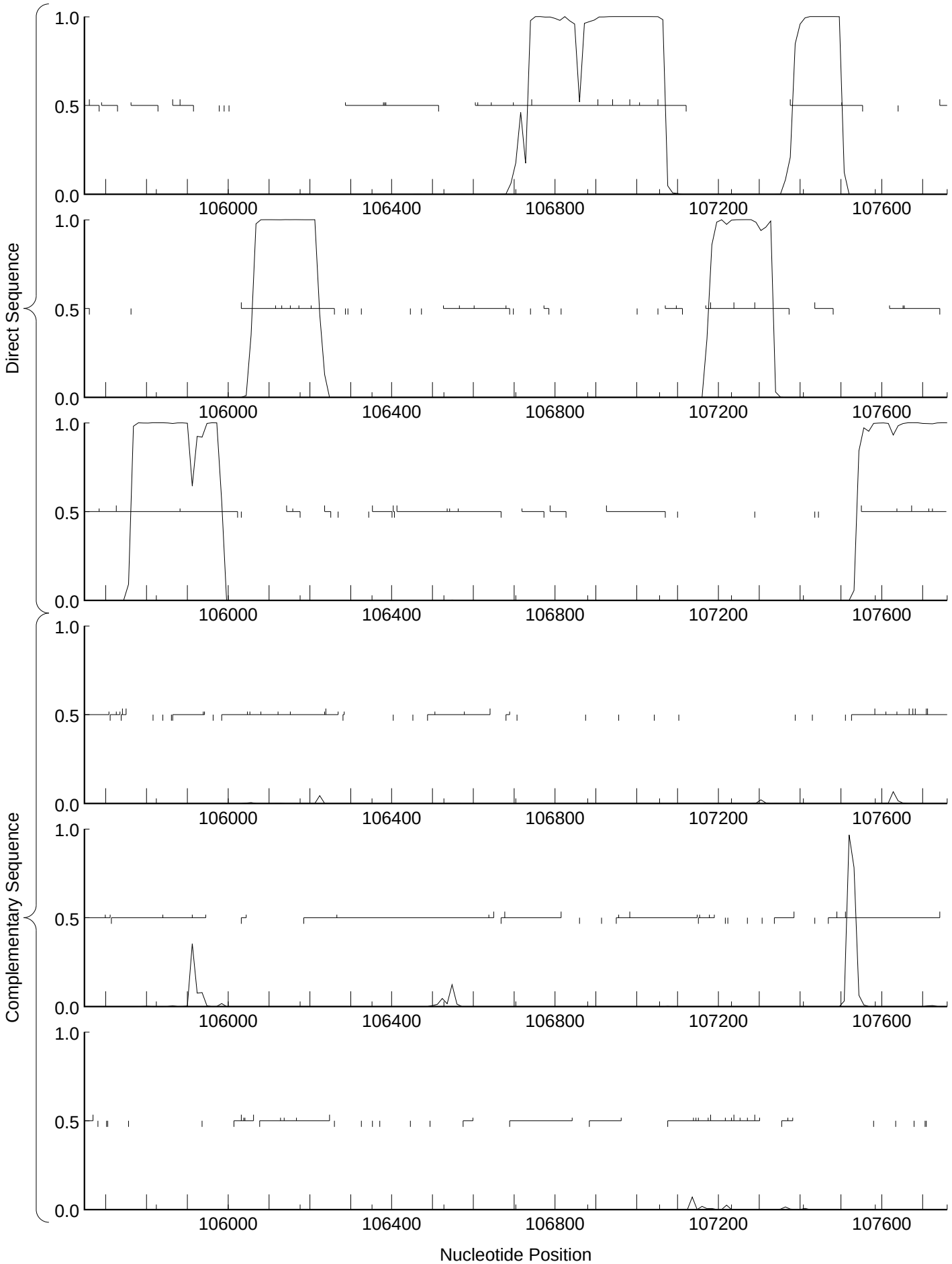
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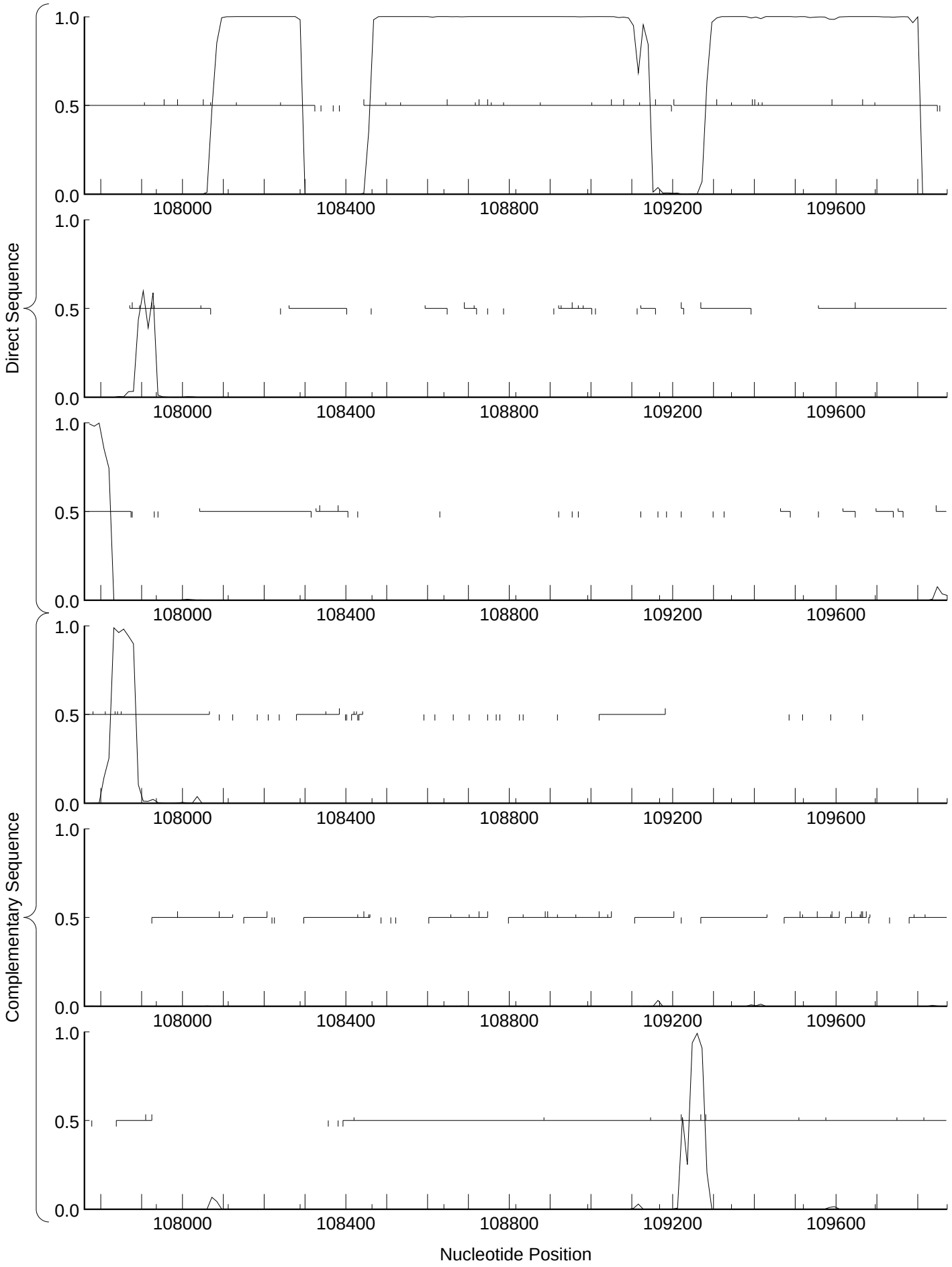
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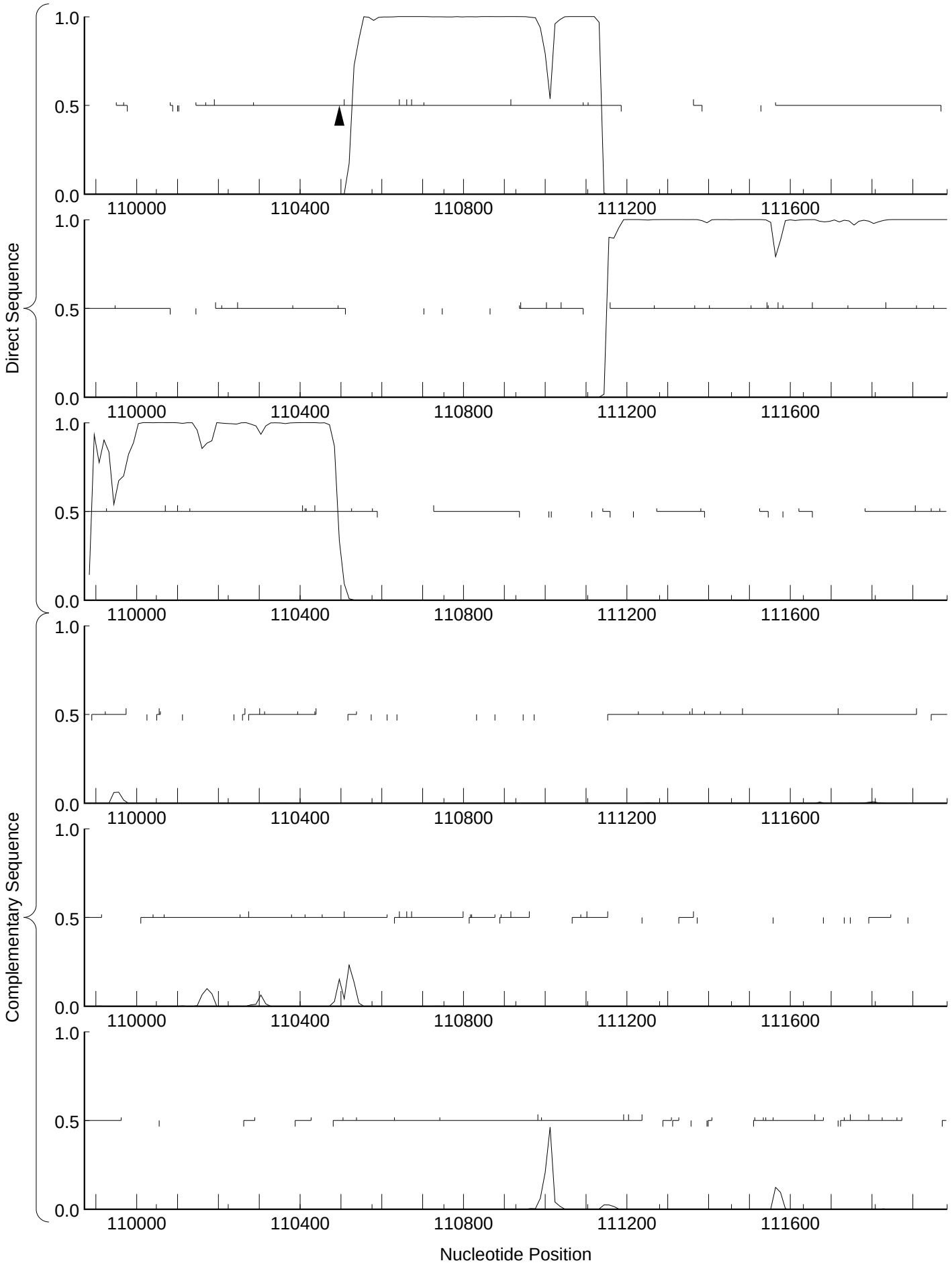
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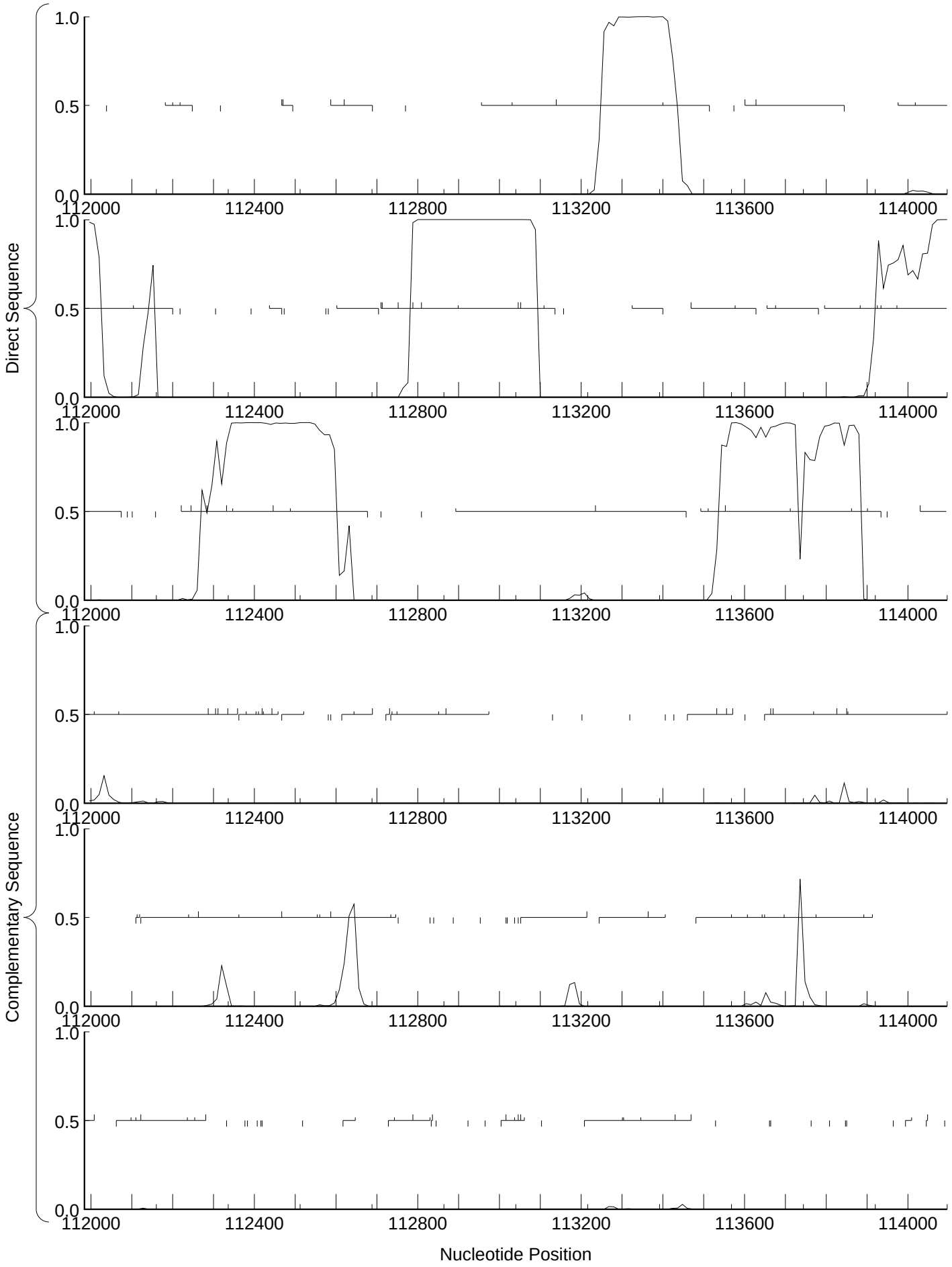
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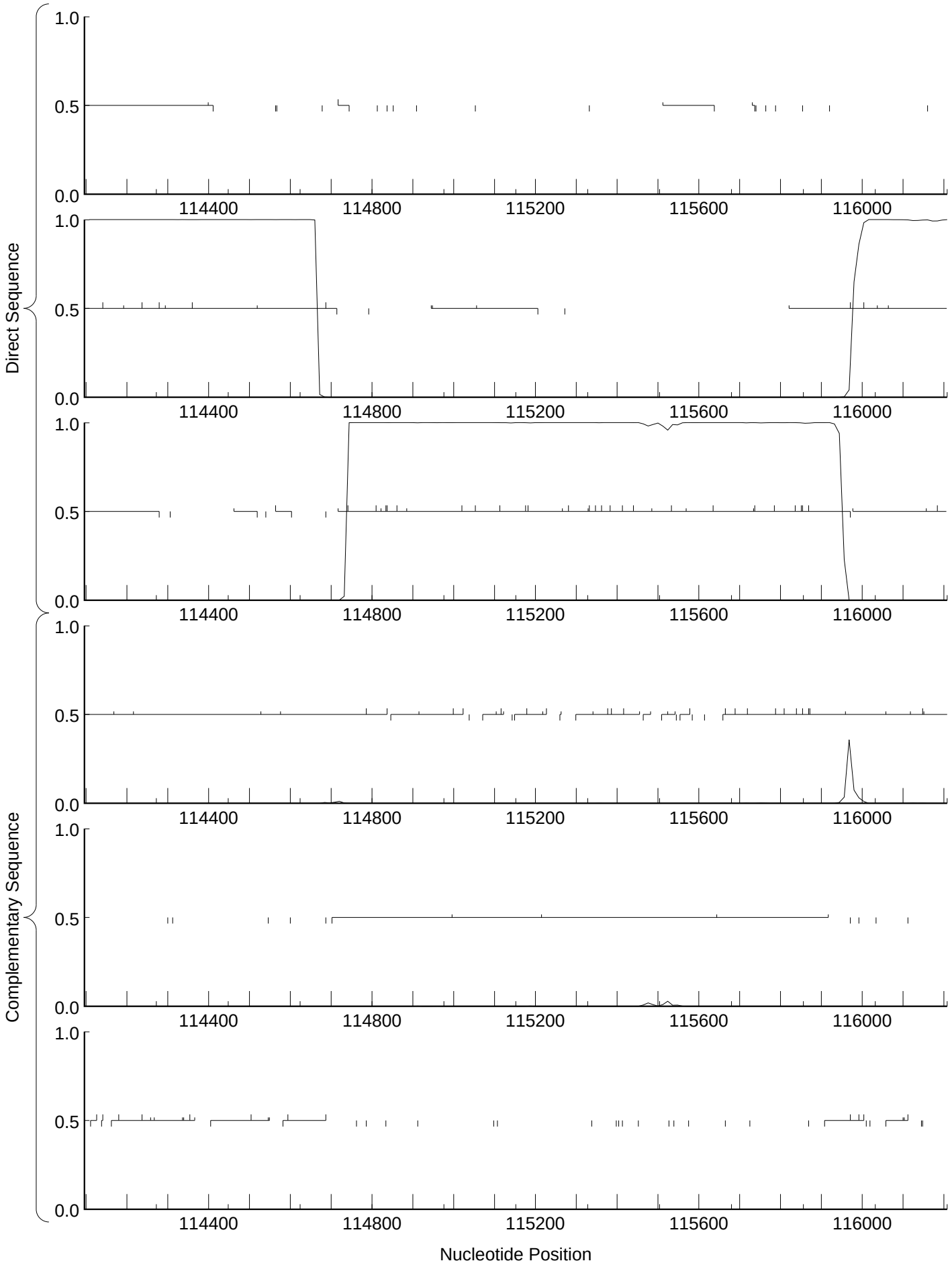
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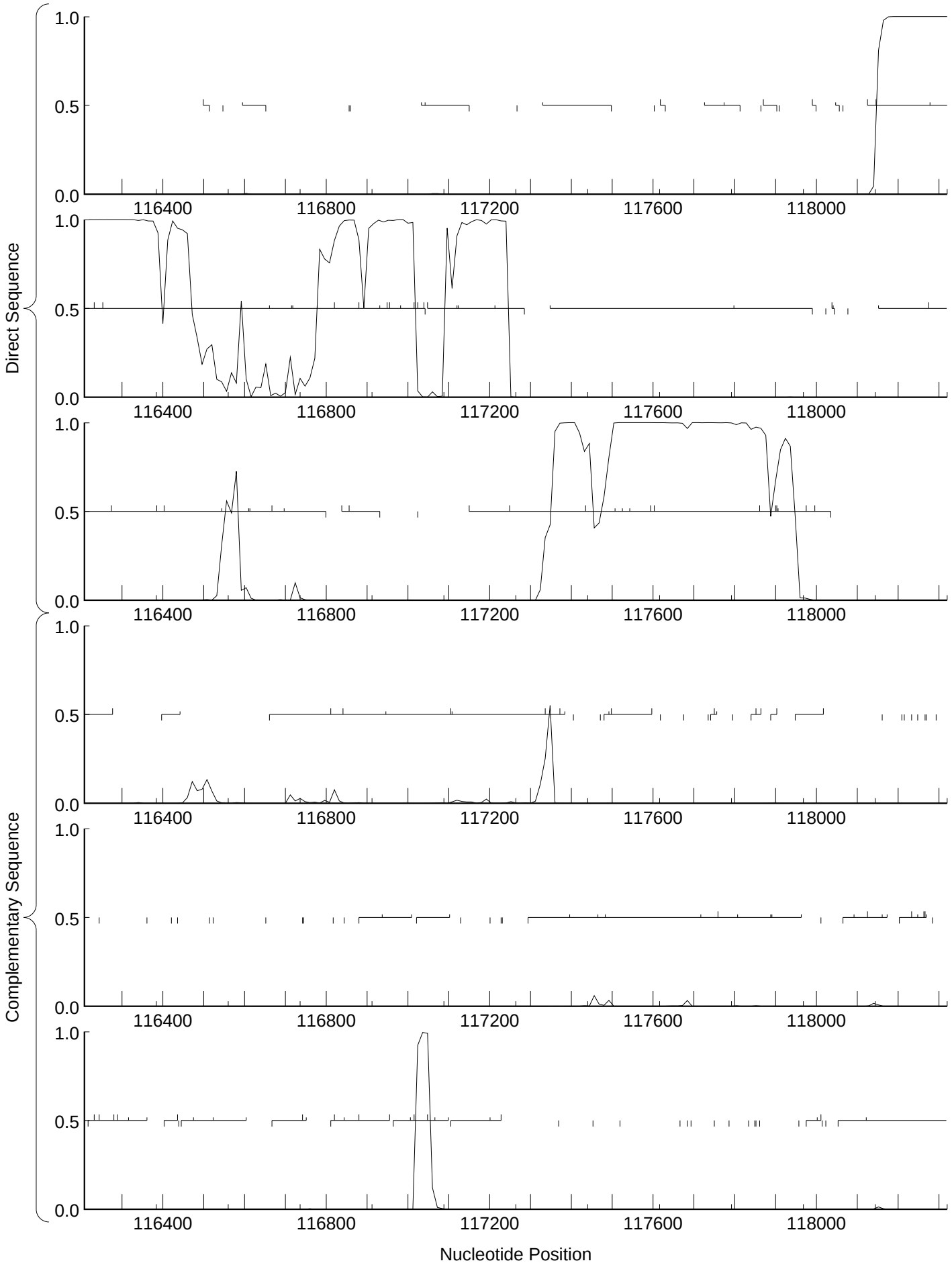
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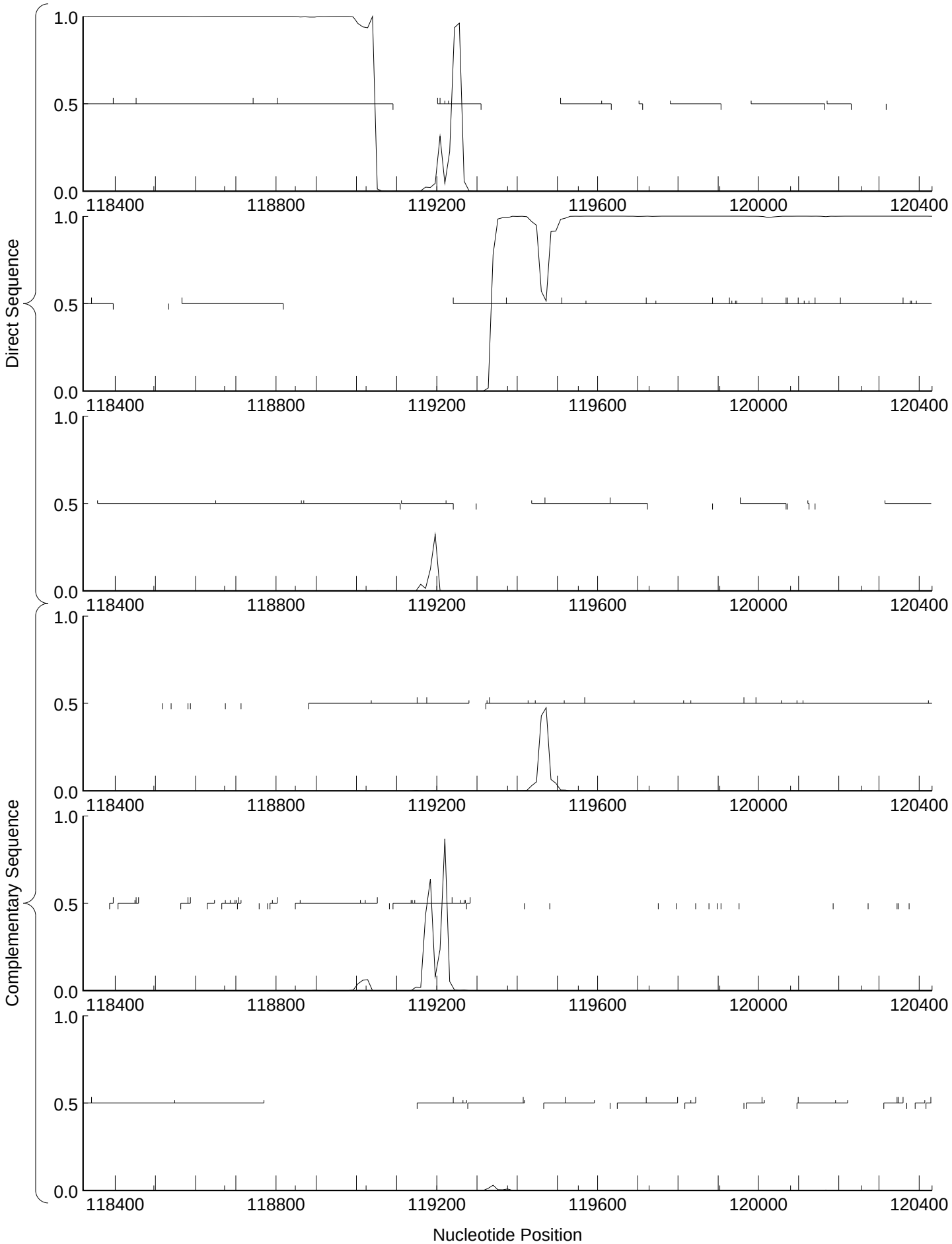
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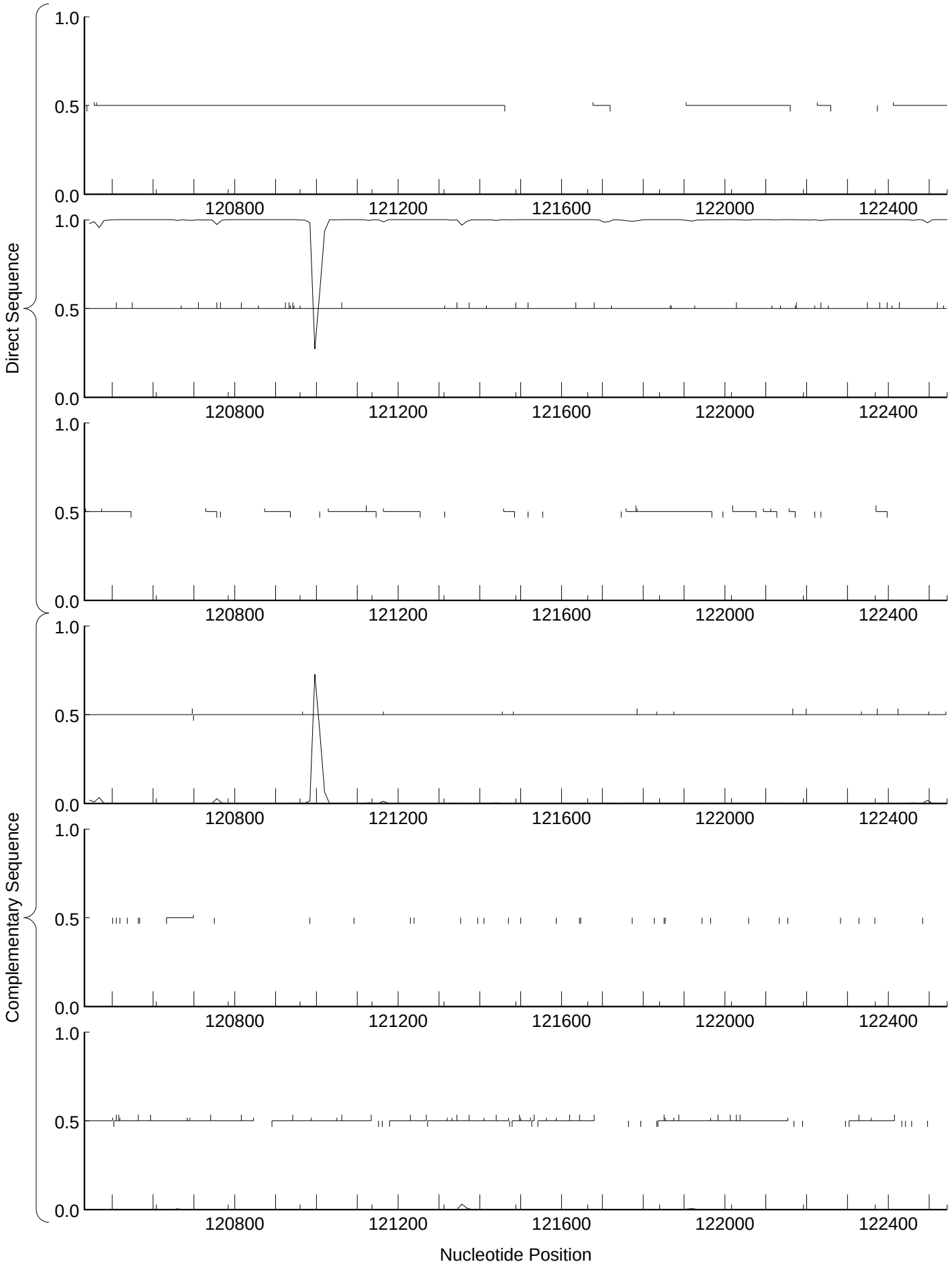
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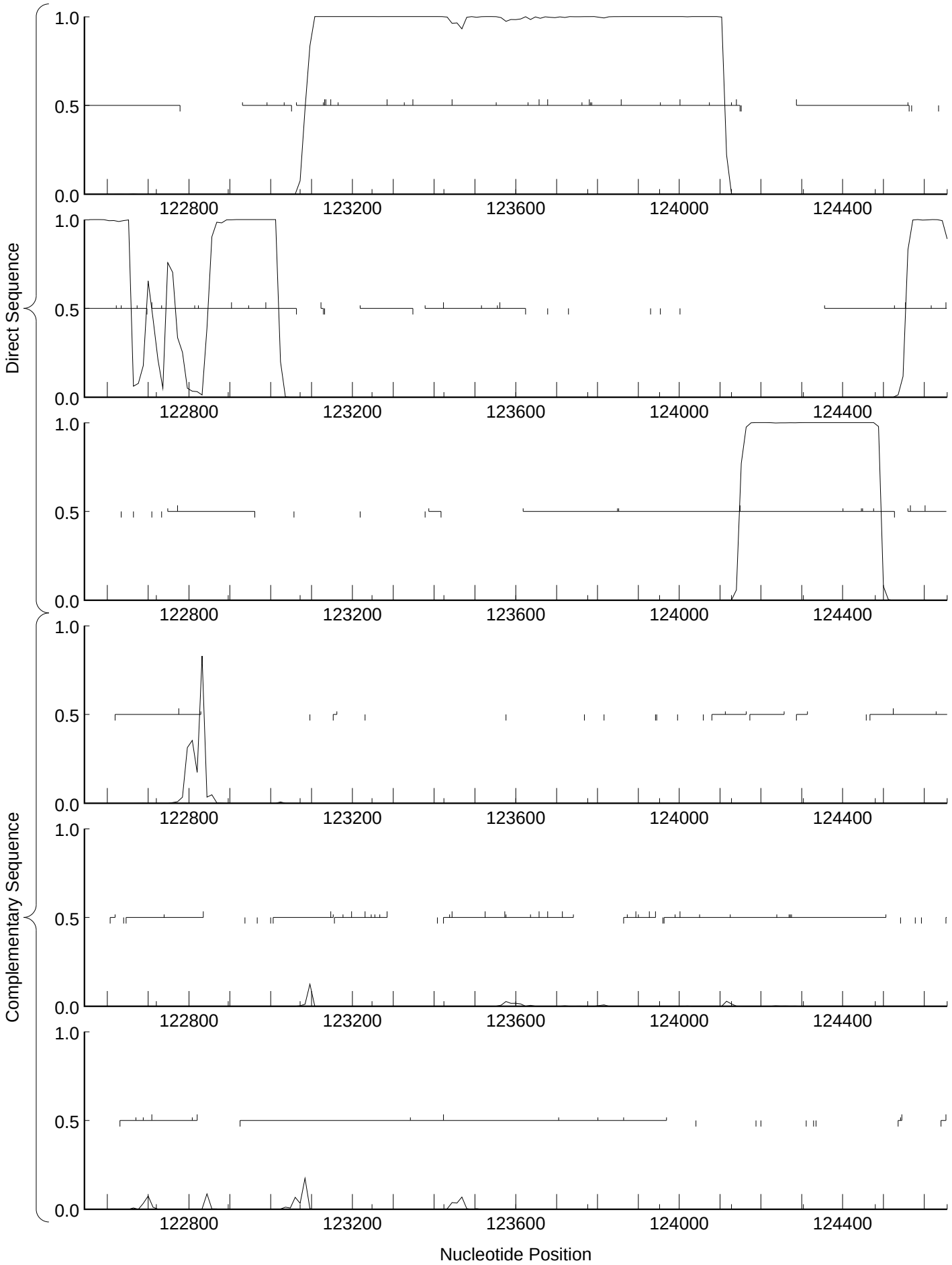
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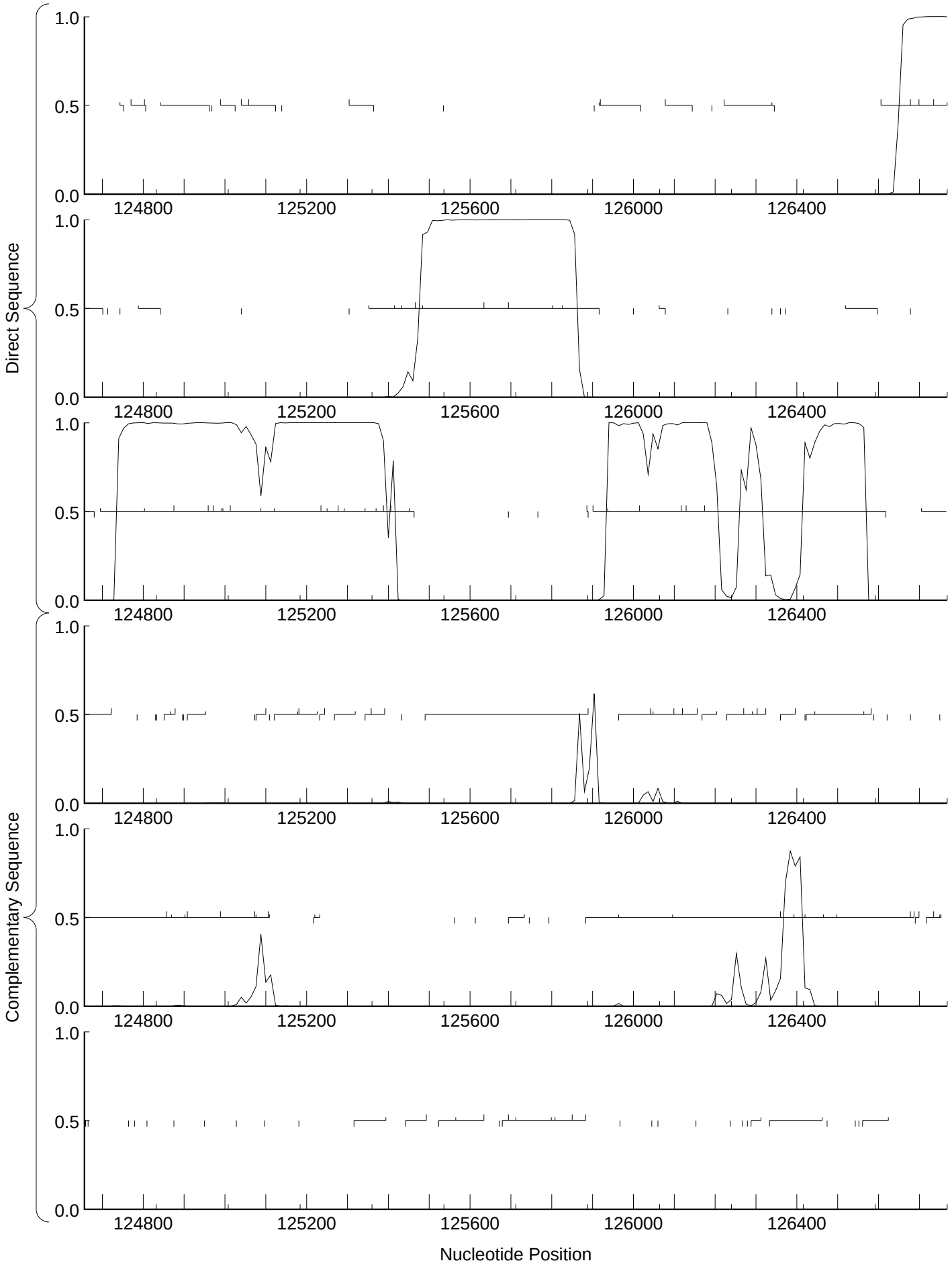
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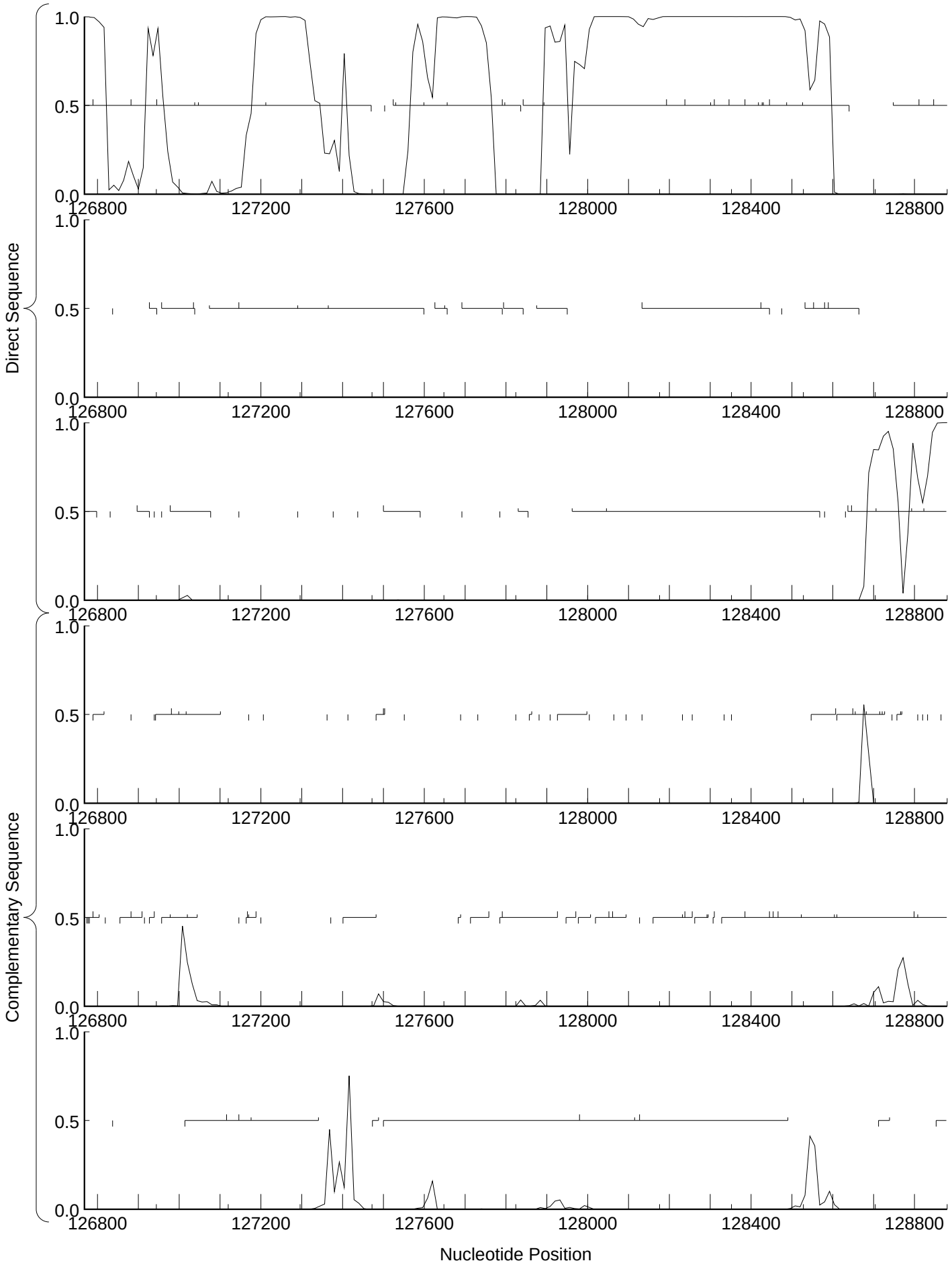
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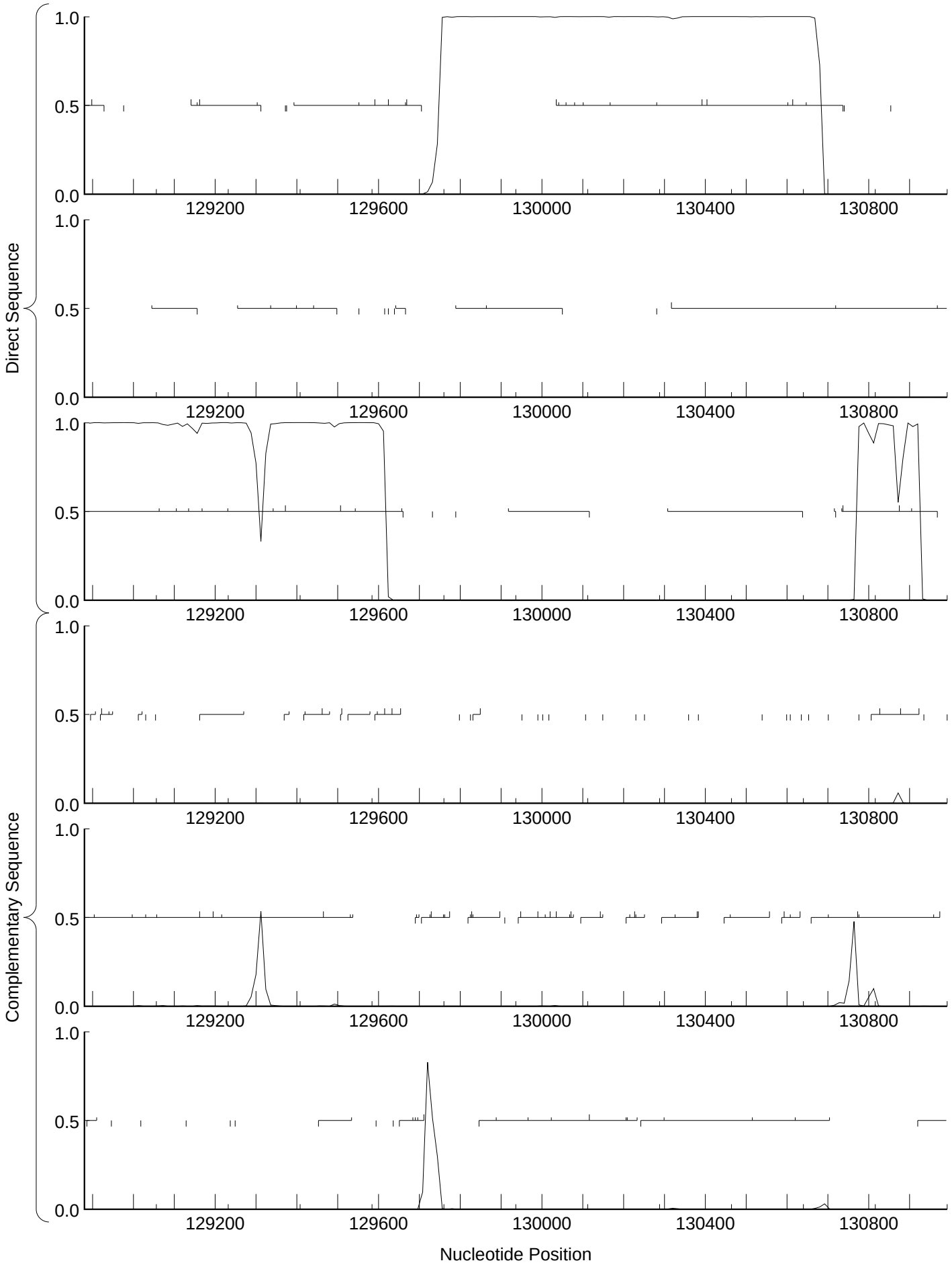
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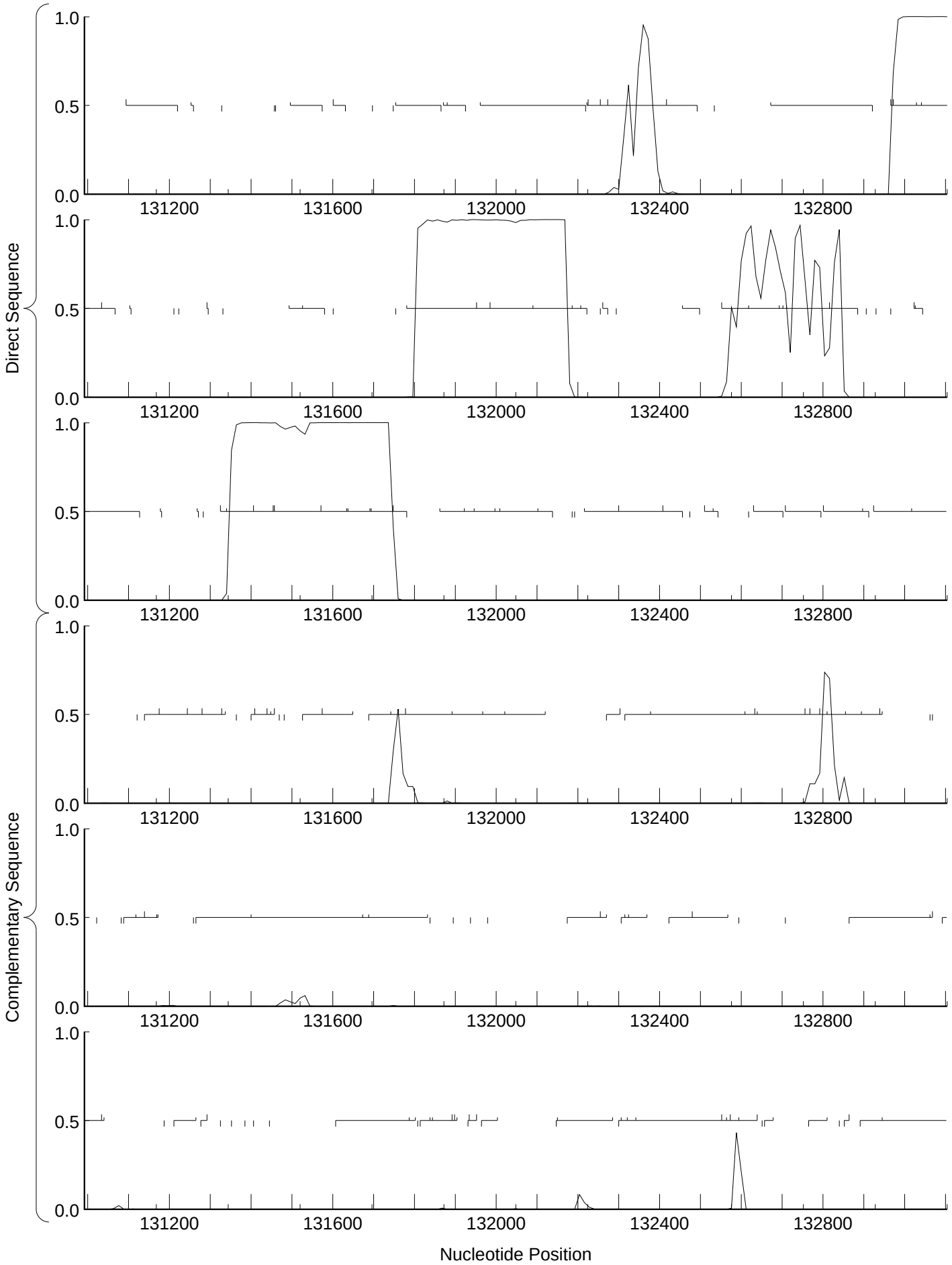
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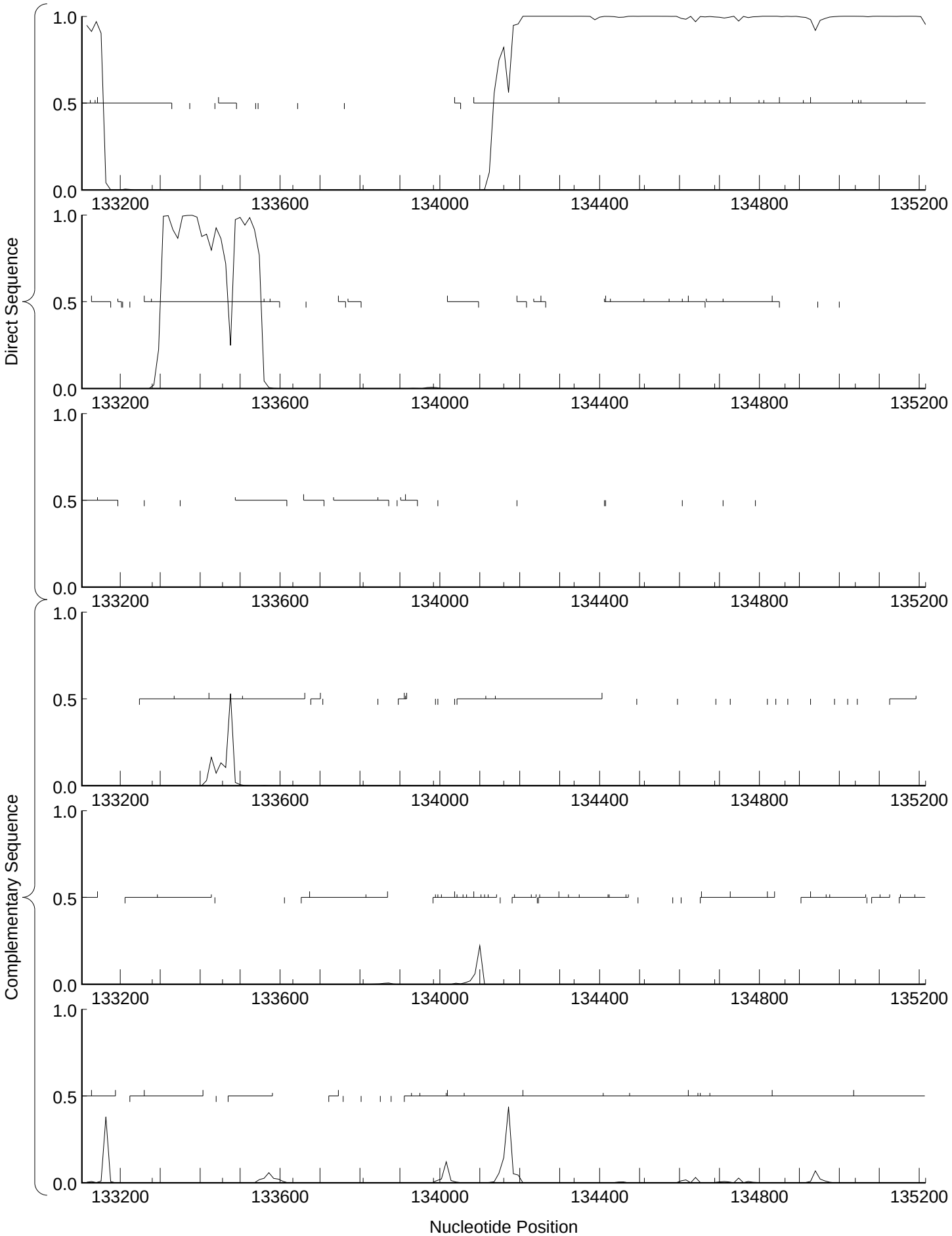
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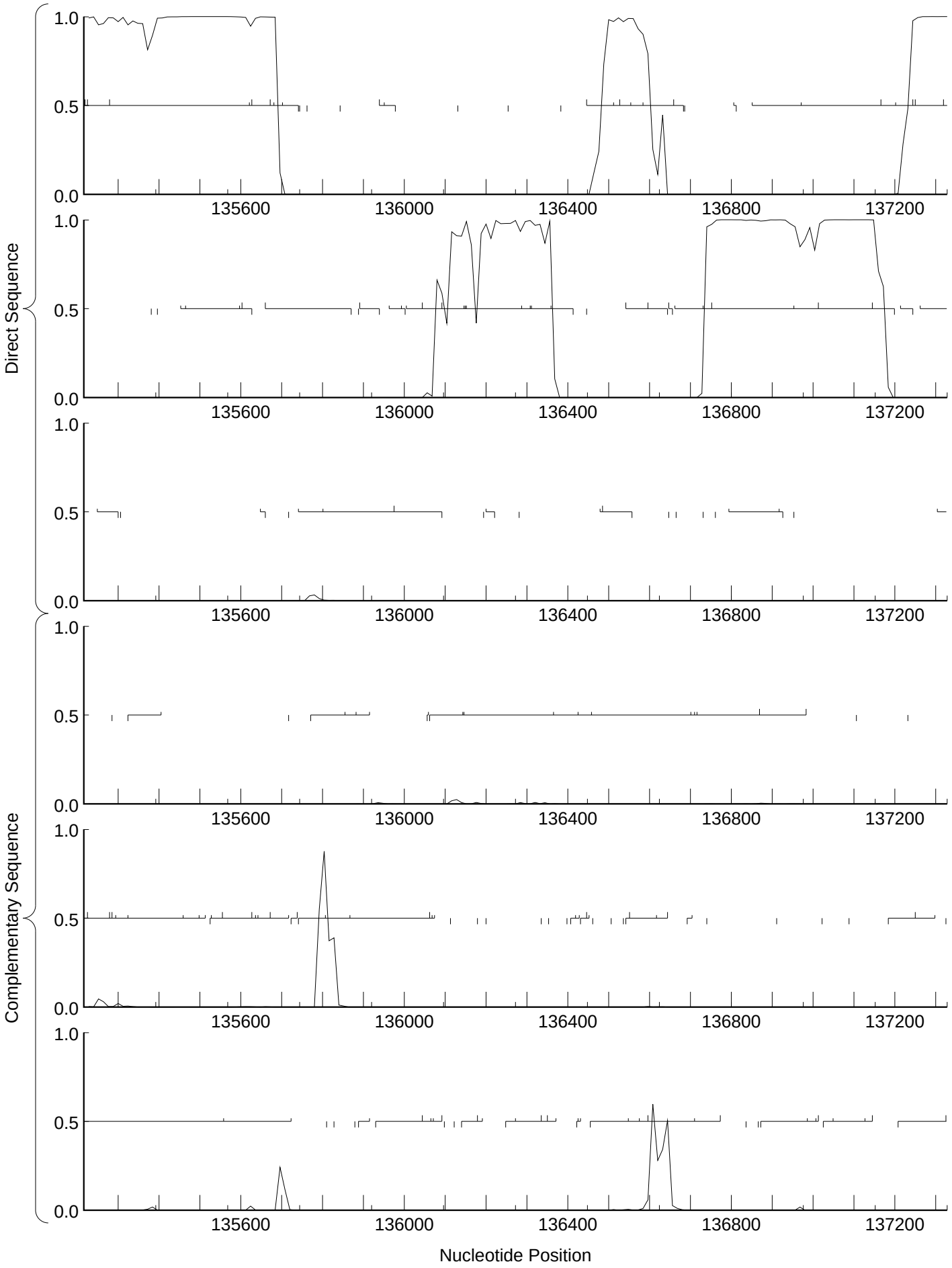
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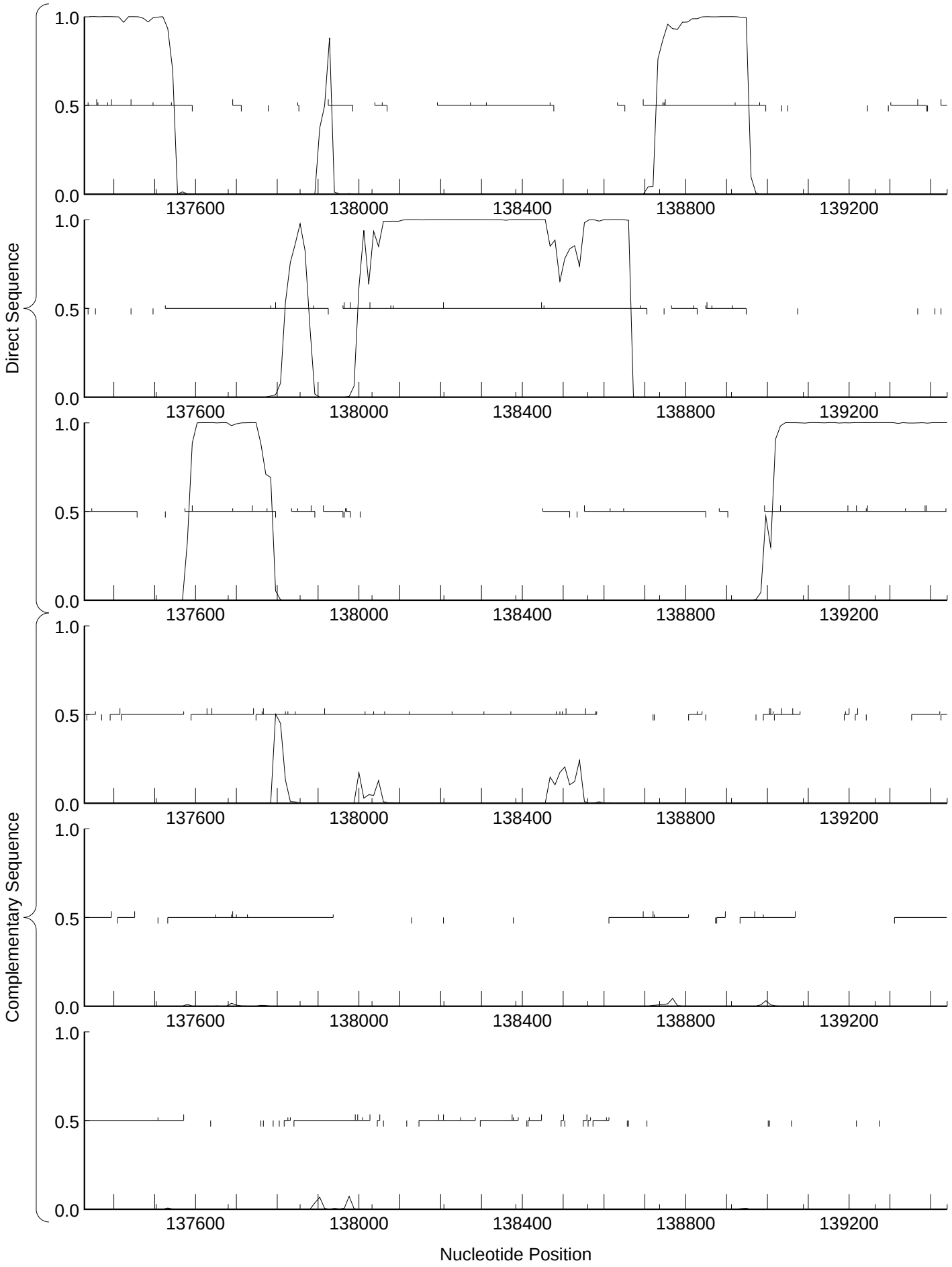
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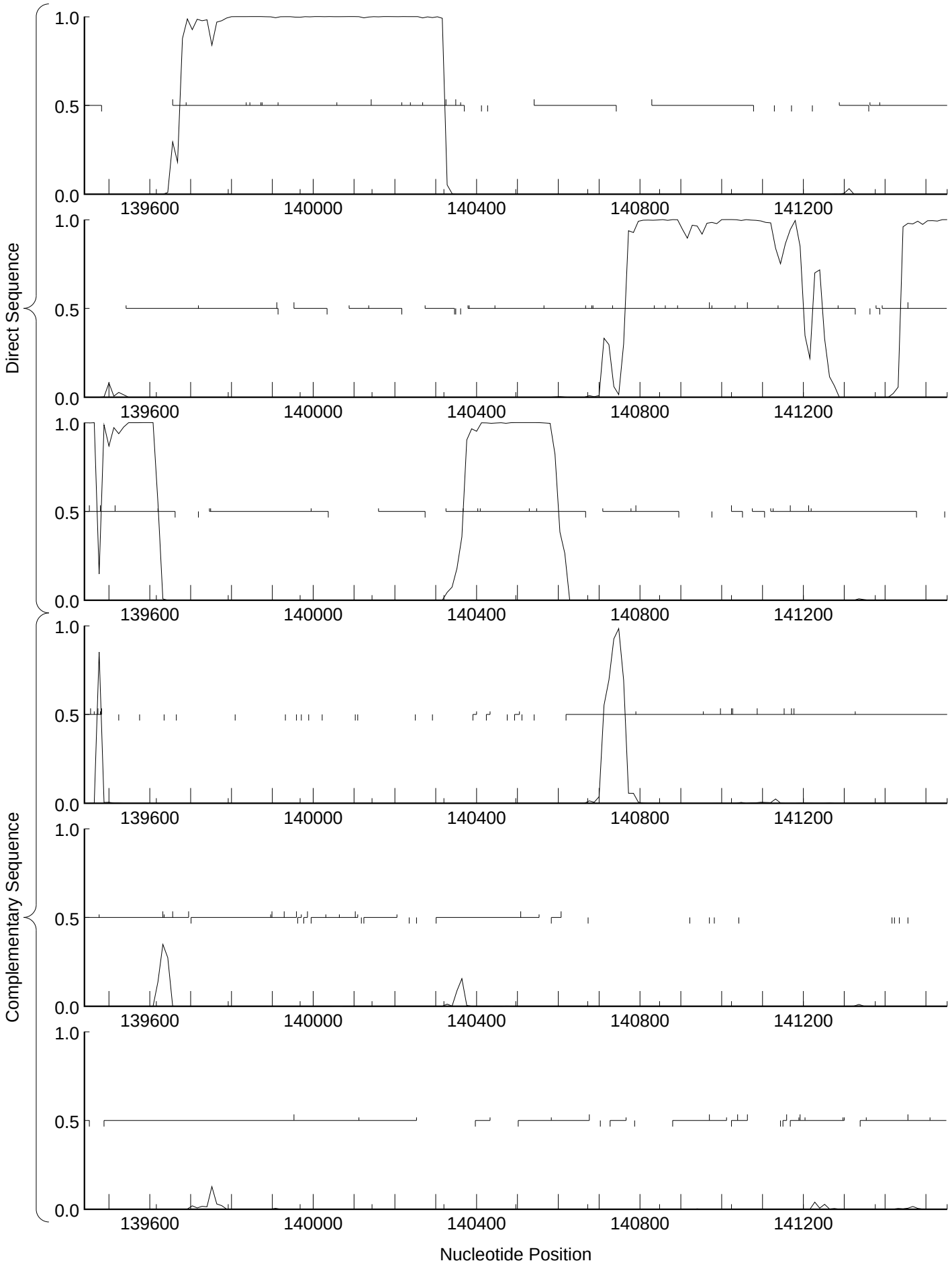
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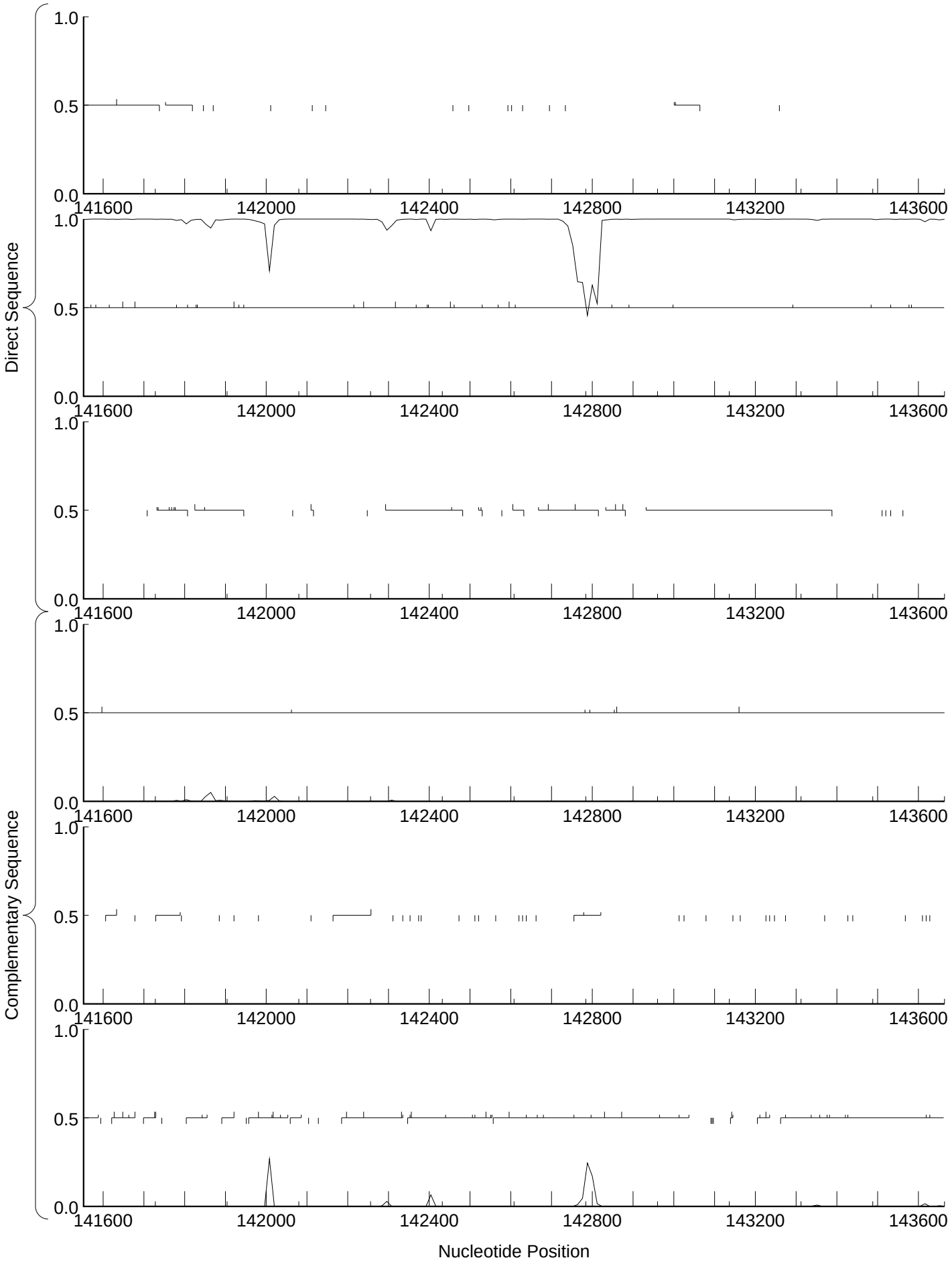
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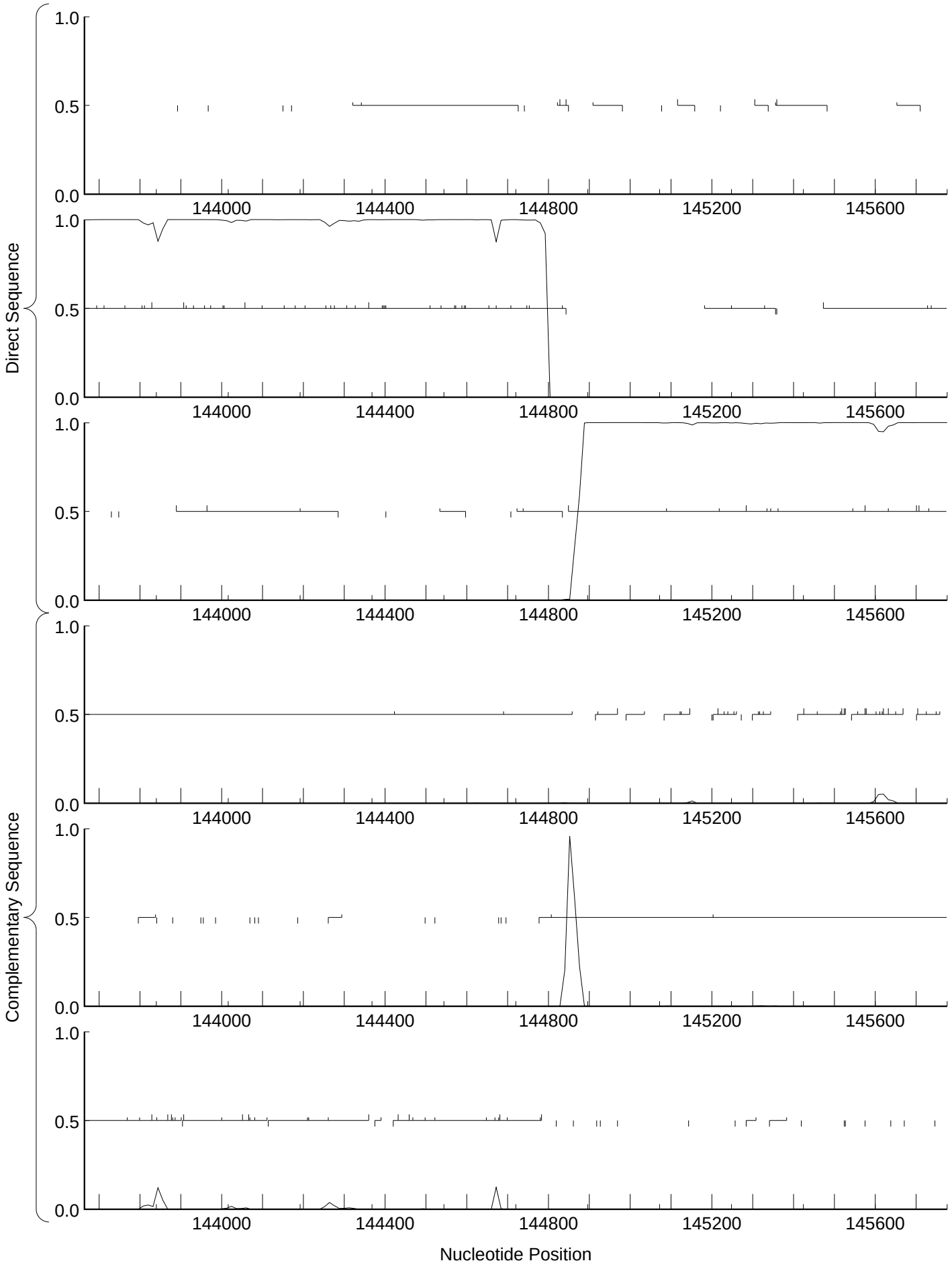
cobacterium phage Tonenili complete sequence, 160985 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 68/78



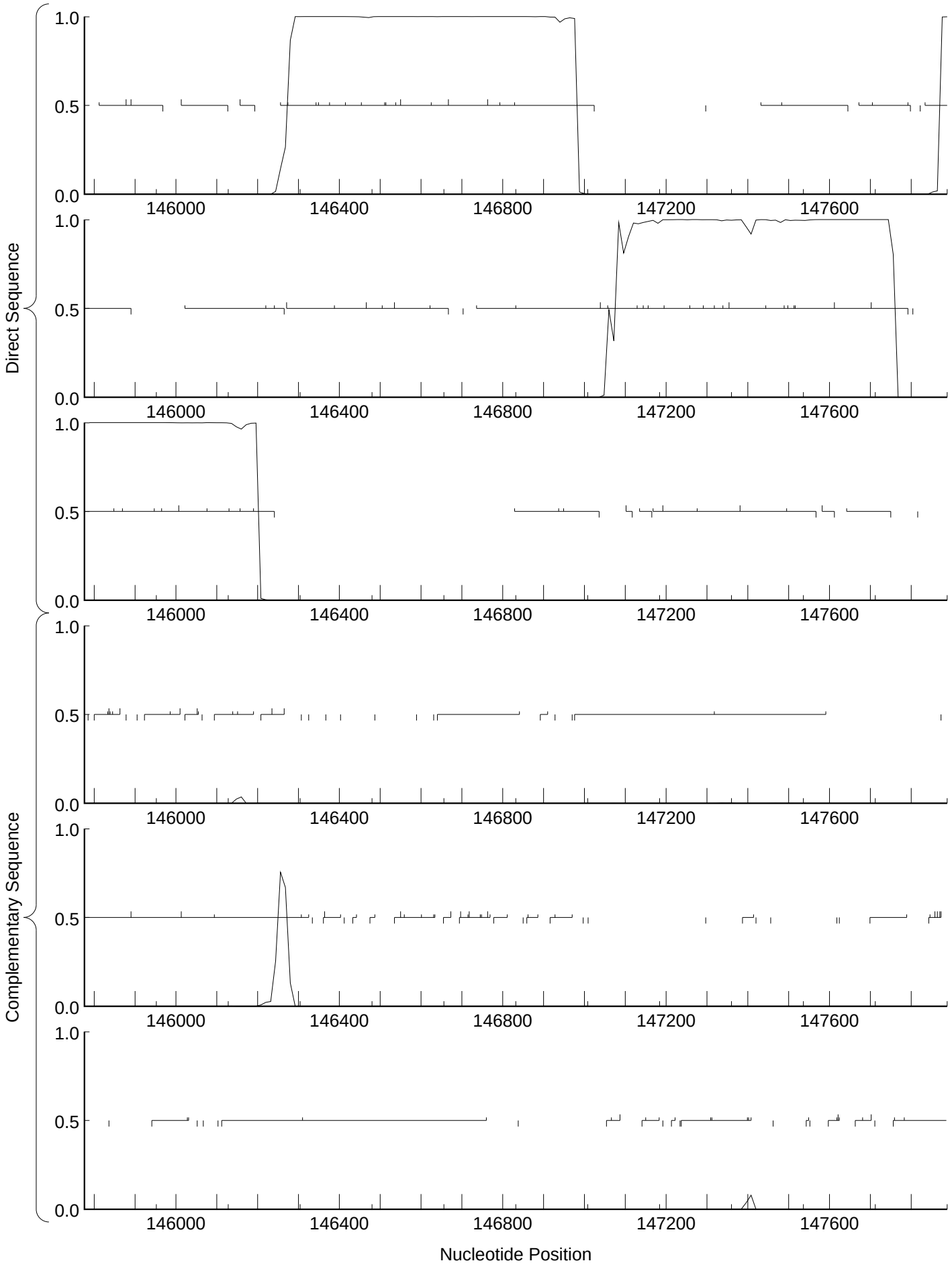
cobacterium phage Tonenili complete sequence, 160985 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 69/78



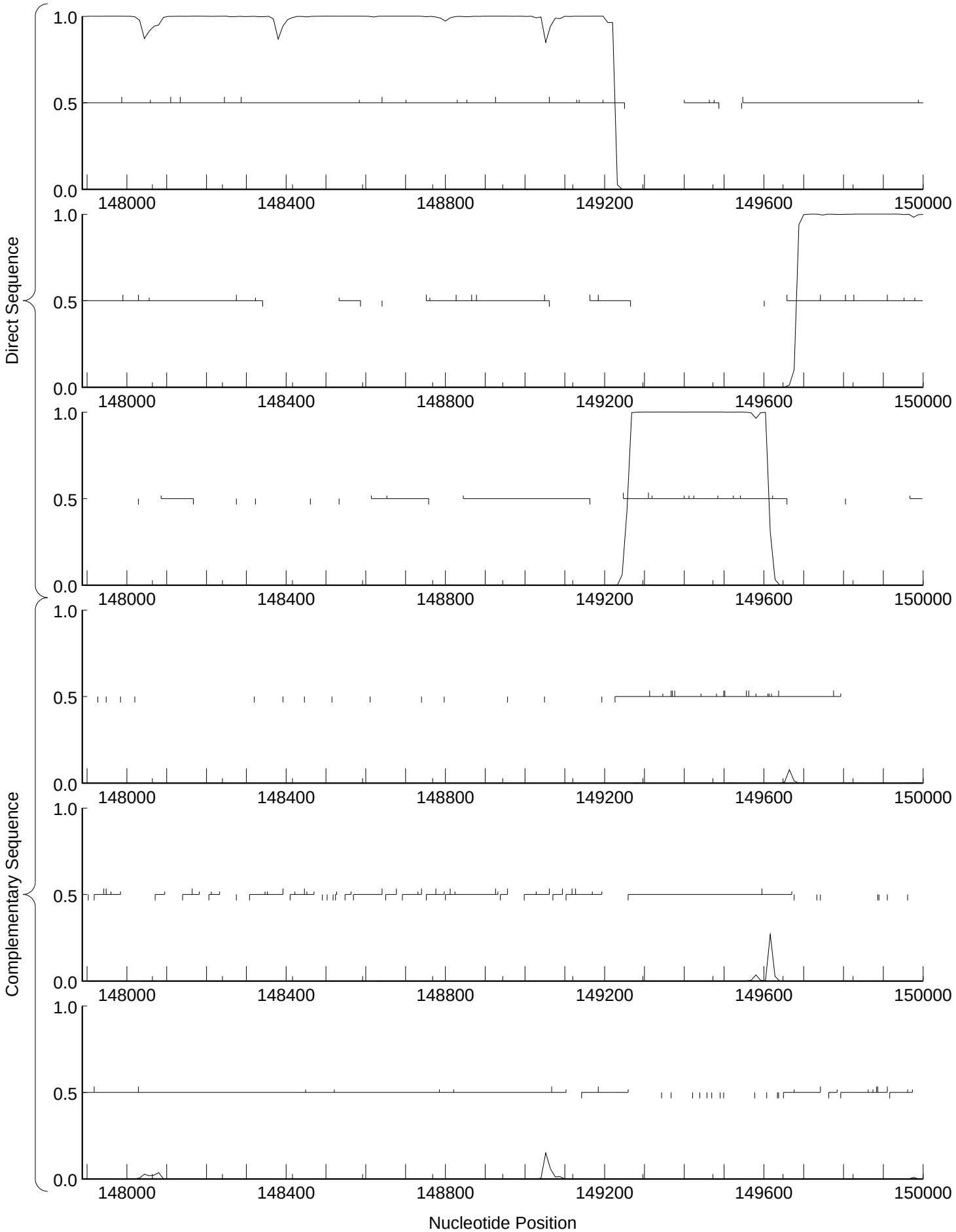
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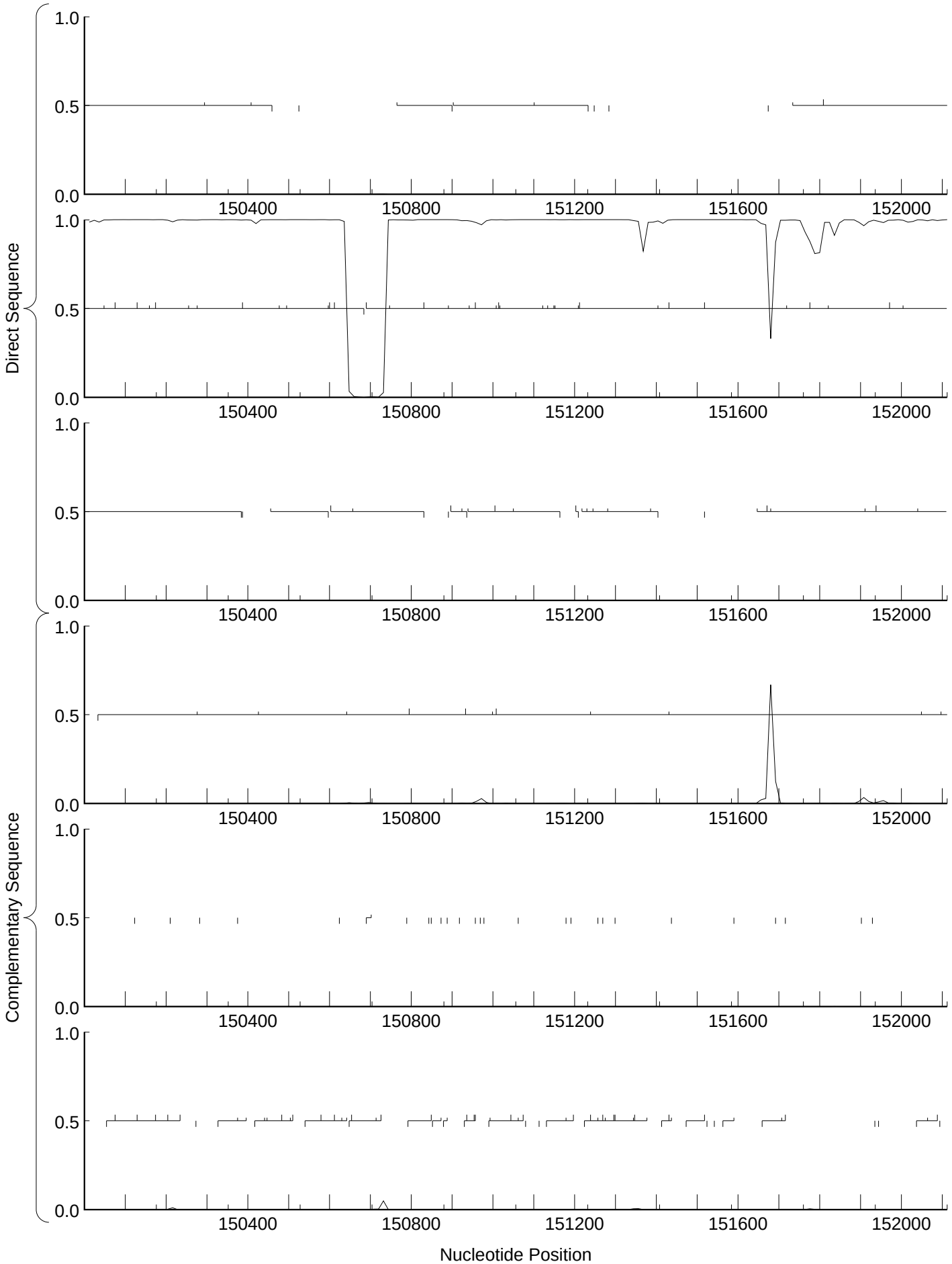
cobacterium phage Tonenili complete sequence, 160985 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 71/78



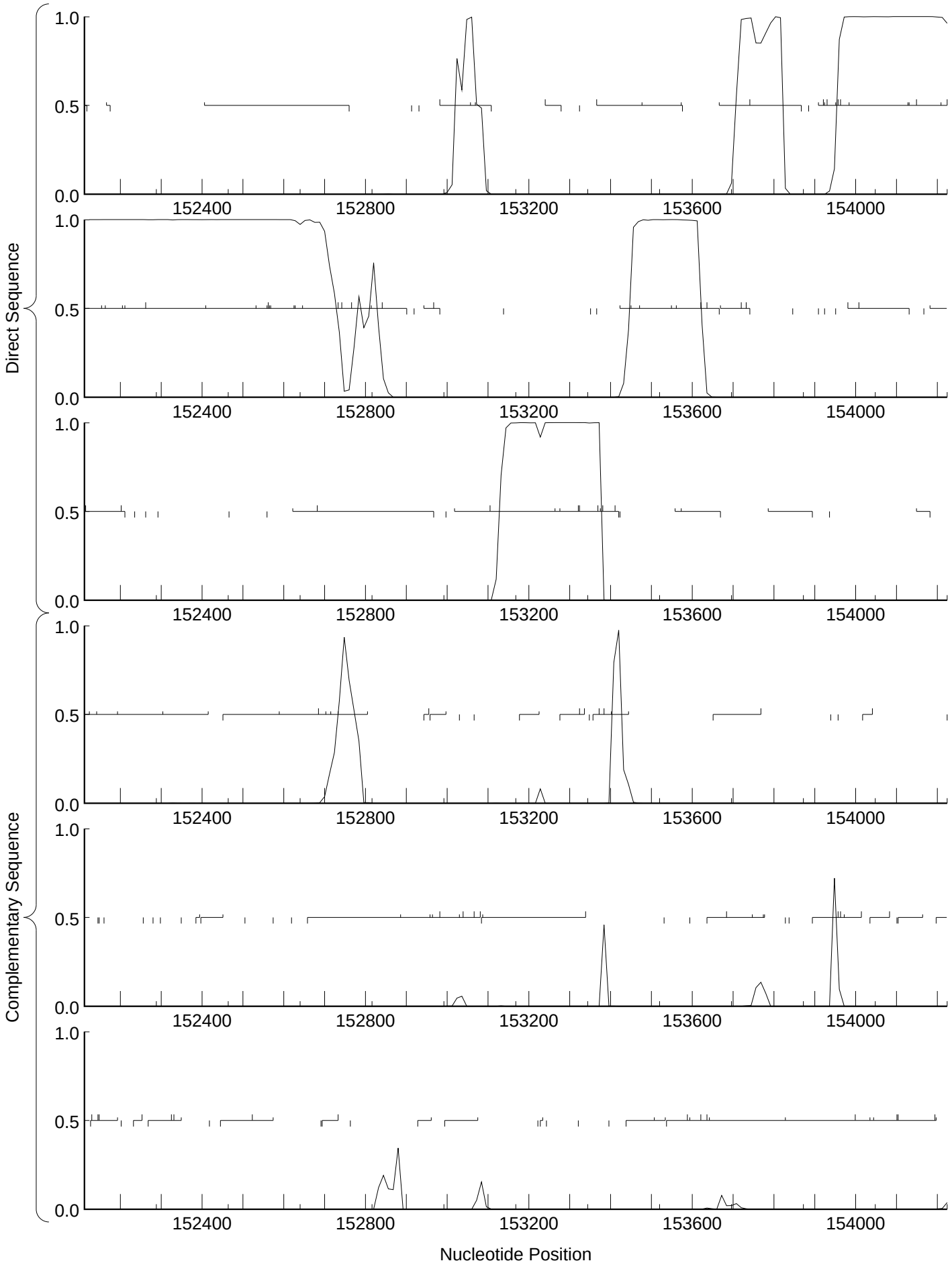
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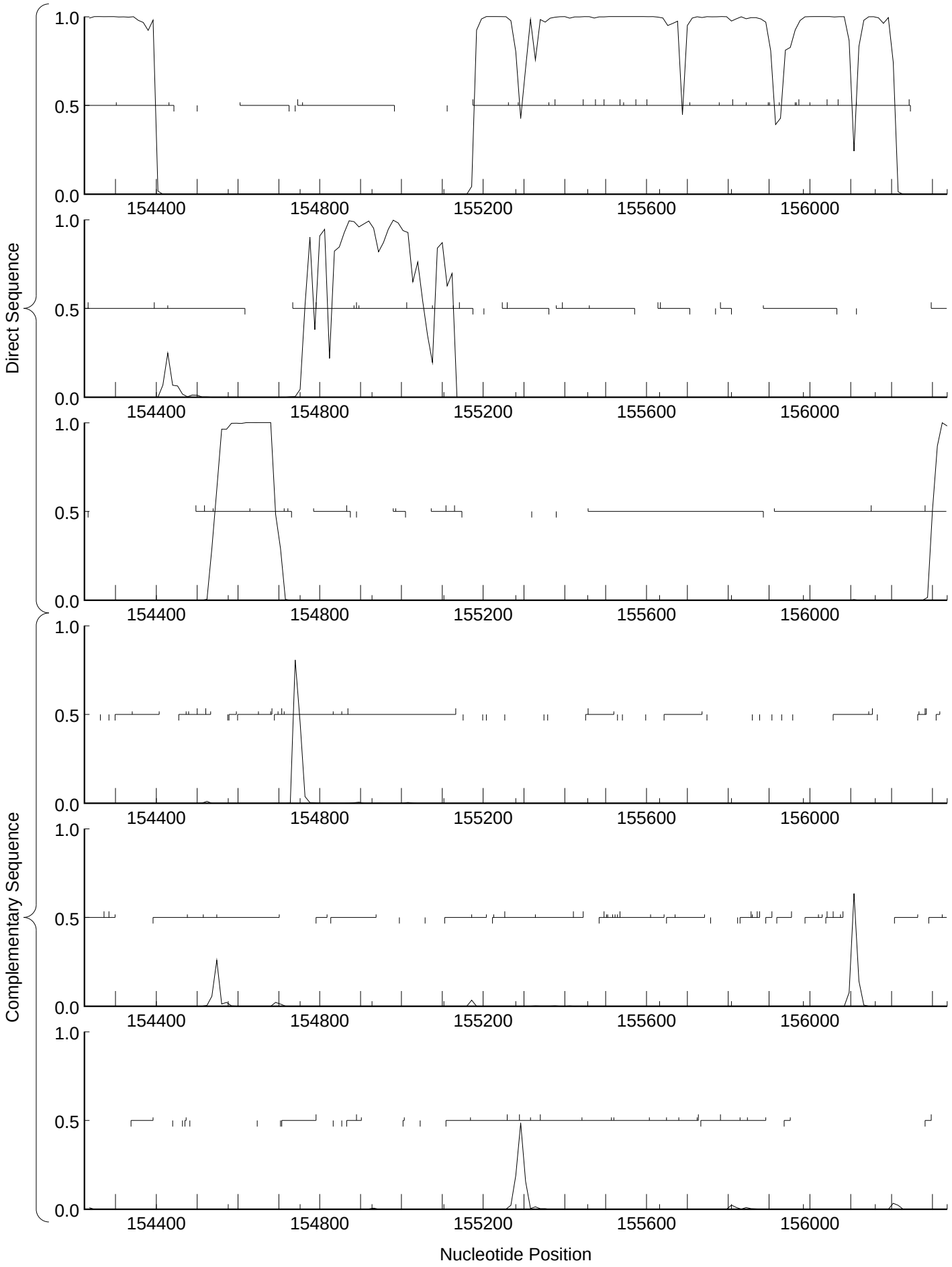
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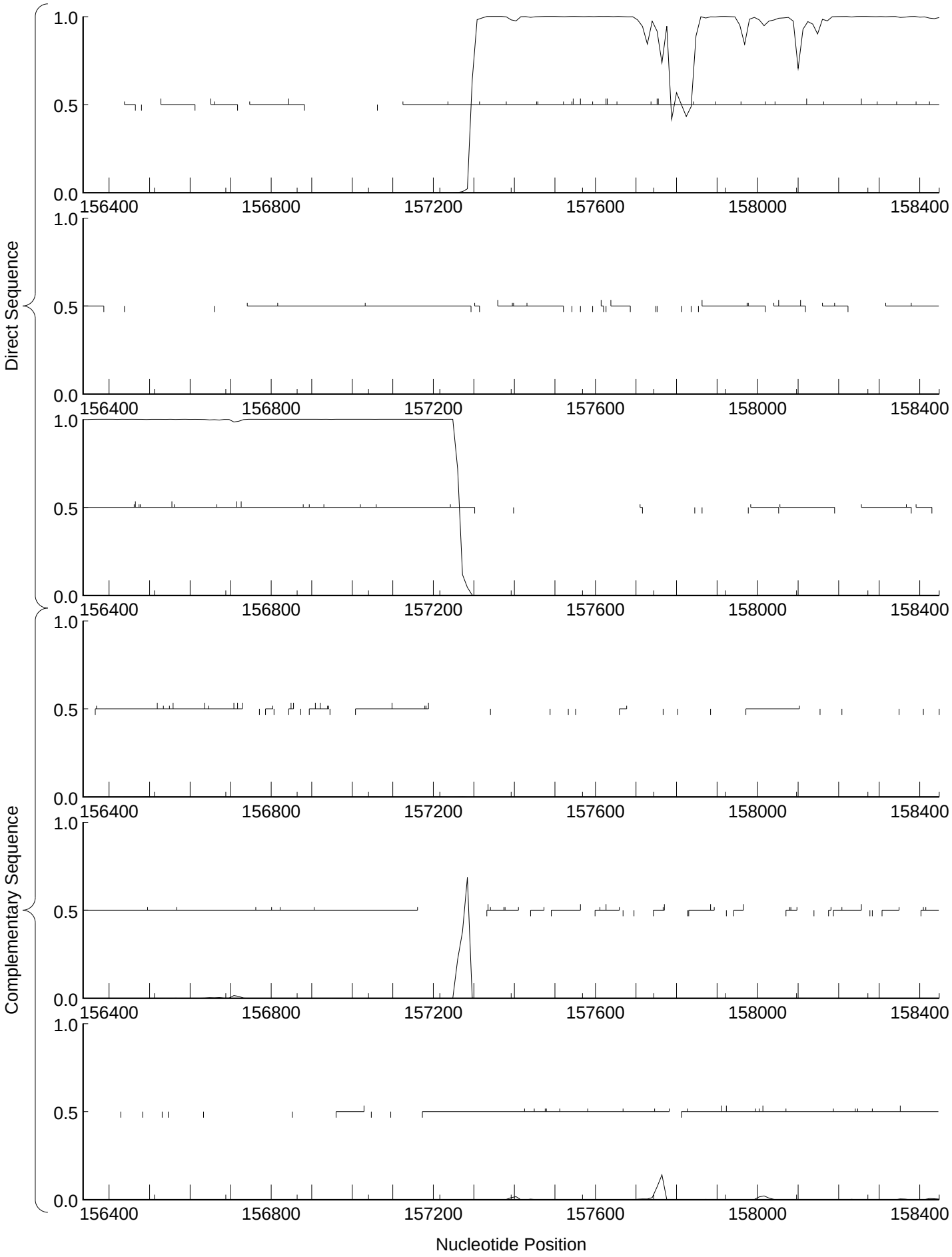
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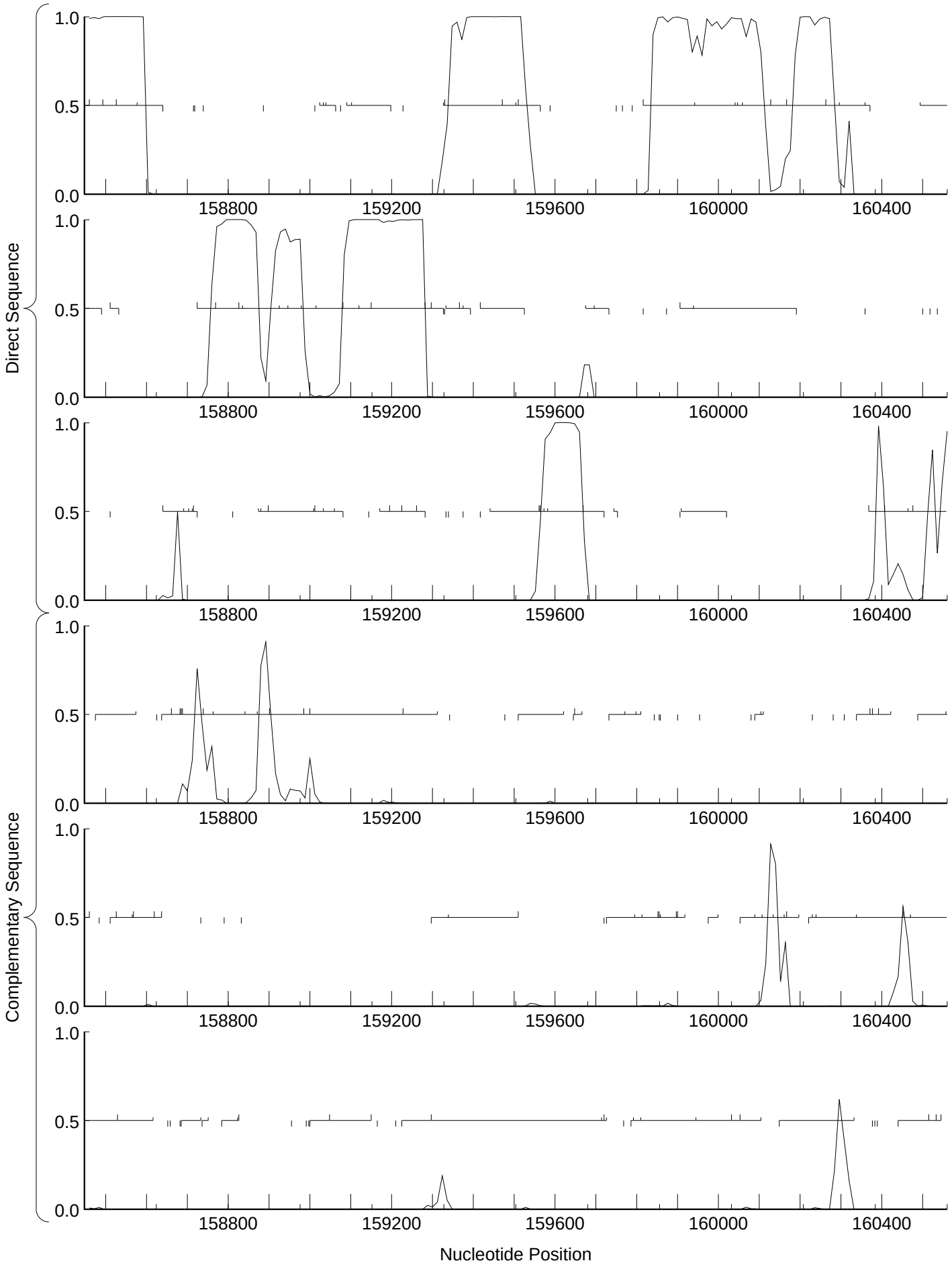
cobacterium phage Tonenili complete sequence, 160985 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 75/78



cobacterium phage Tonenili complete sequence, 160985 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 76/78



cobacterium phage Tonenili complete sequence, 160985 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 77/78



cobacterium phage Tonenili complete sequence, 160985 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 78/78

