GeneMark

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

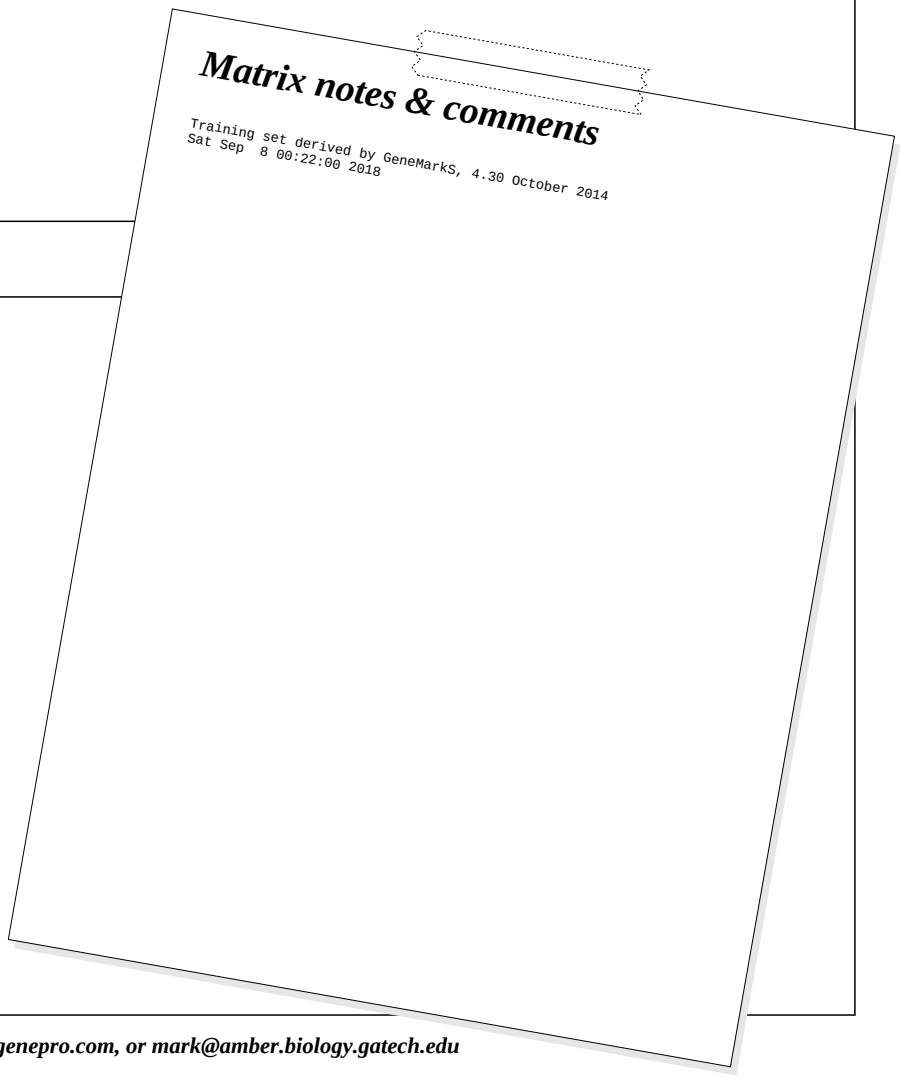
PROGRAM INFORMATION

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Sequence         :  Arthrobacter phage Inked complete sequence, 57862 bp including 9-base 3’ overhang (CGCCGGCC
Analysis Date    :  3/28/24 at 10:30:07
Pages            :  29
Sequence Length  :  57862 bp
GC Content       :   51.75%

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

PS-Version       :  1.2

GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
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MATRIX INFORMATION

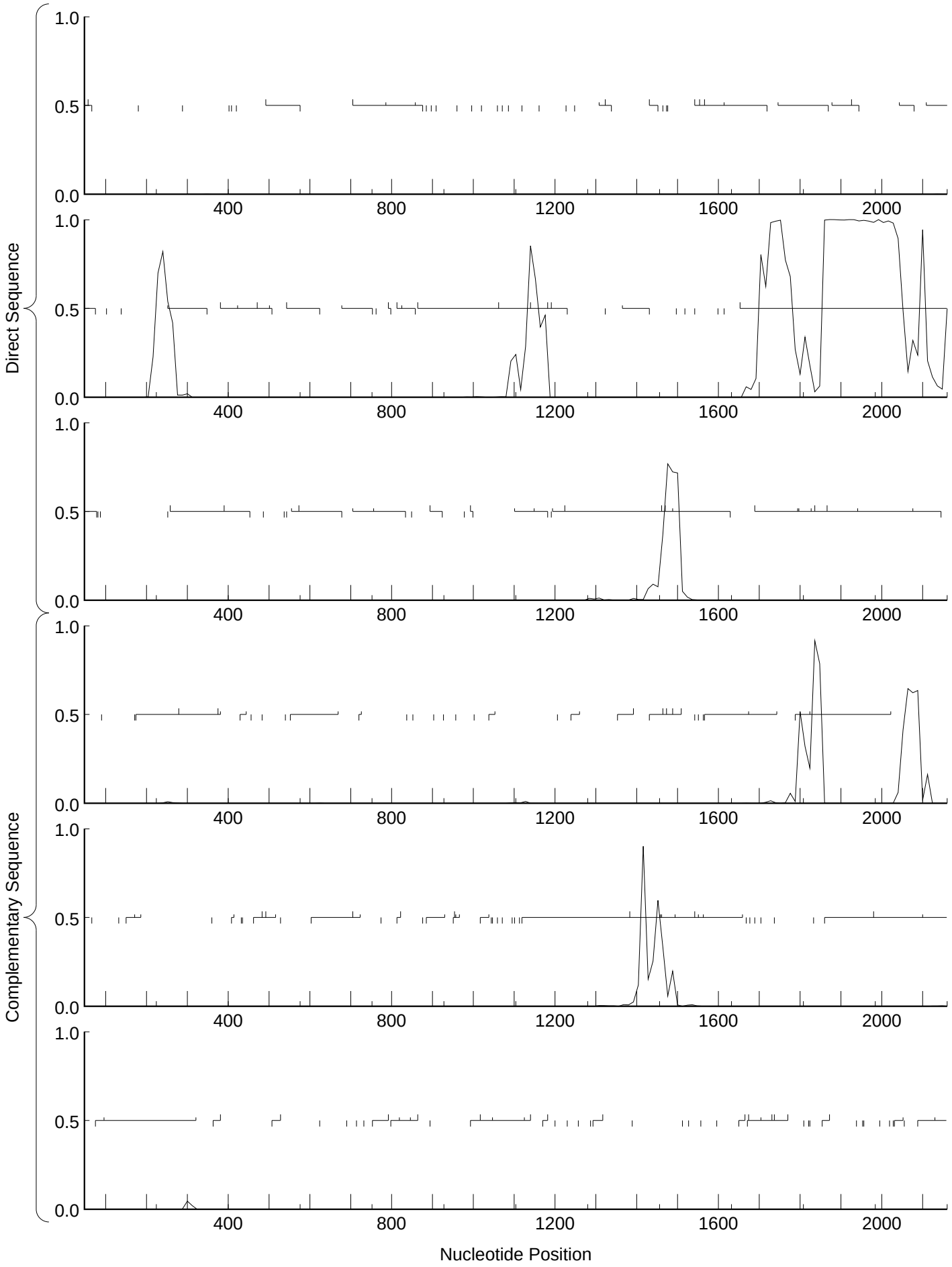
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Matrix     :  -
Author     :  -

Order      :  4
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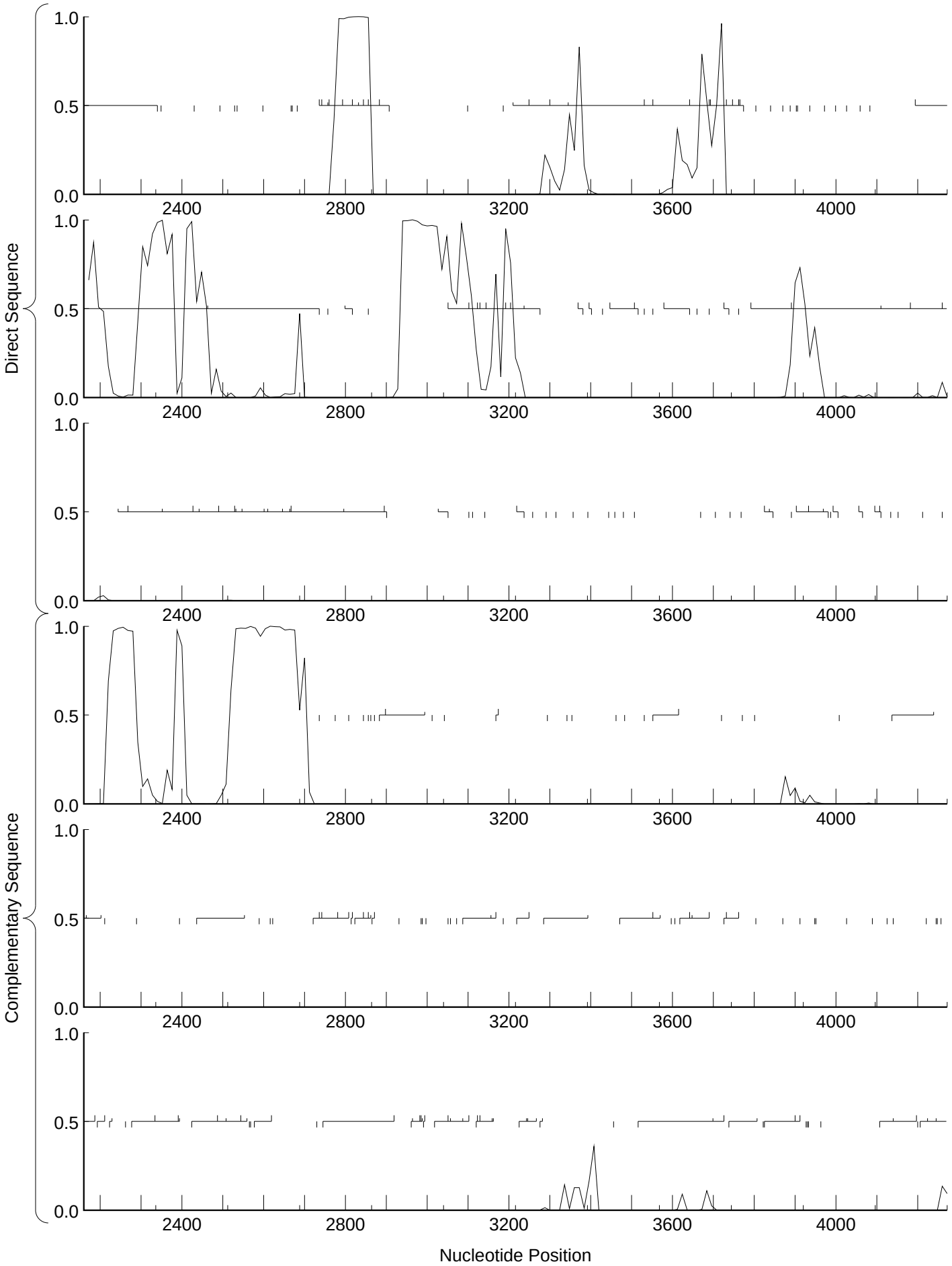

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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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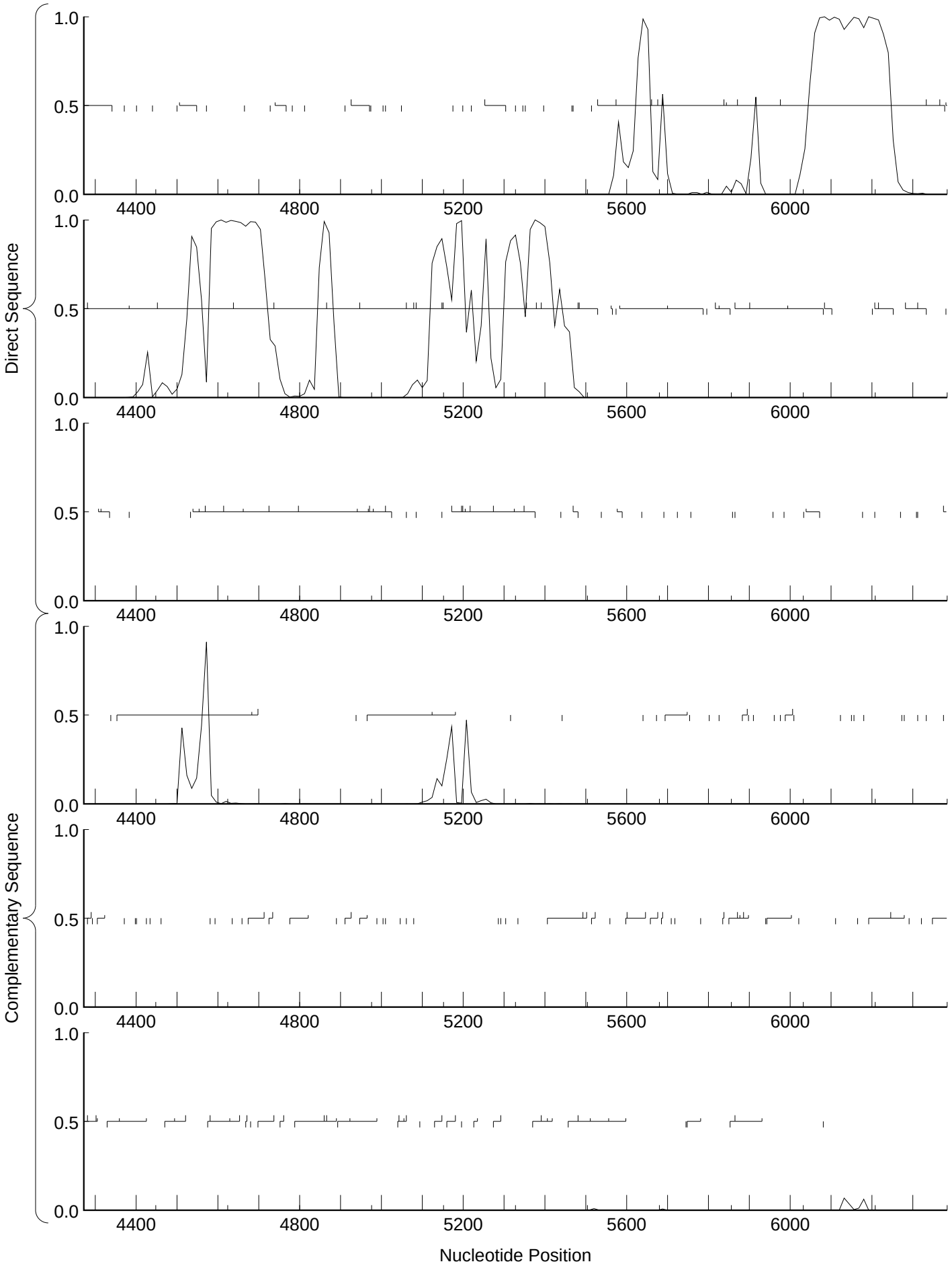

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

ked complete sequence, 57862 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AU, Order 4, Window 96, Step 12, 2/29

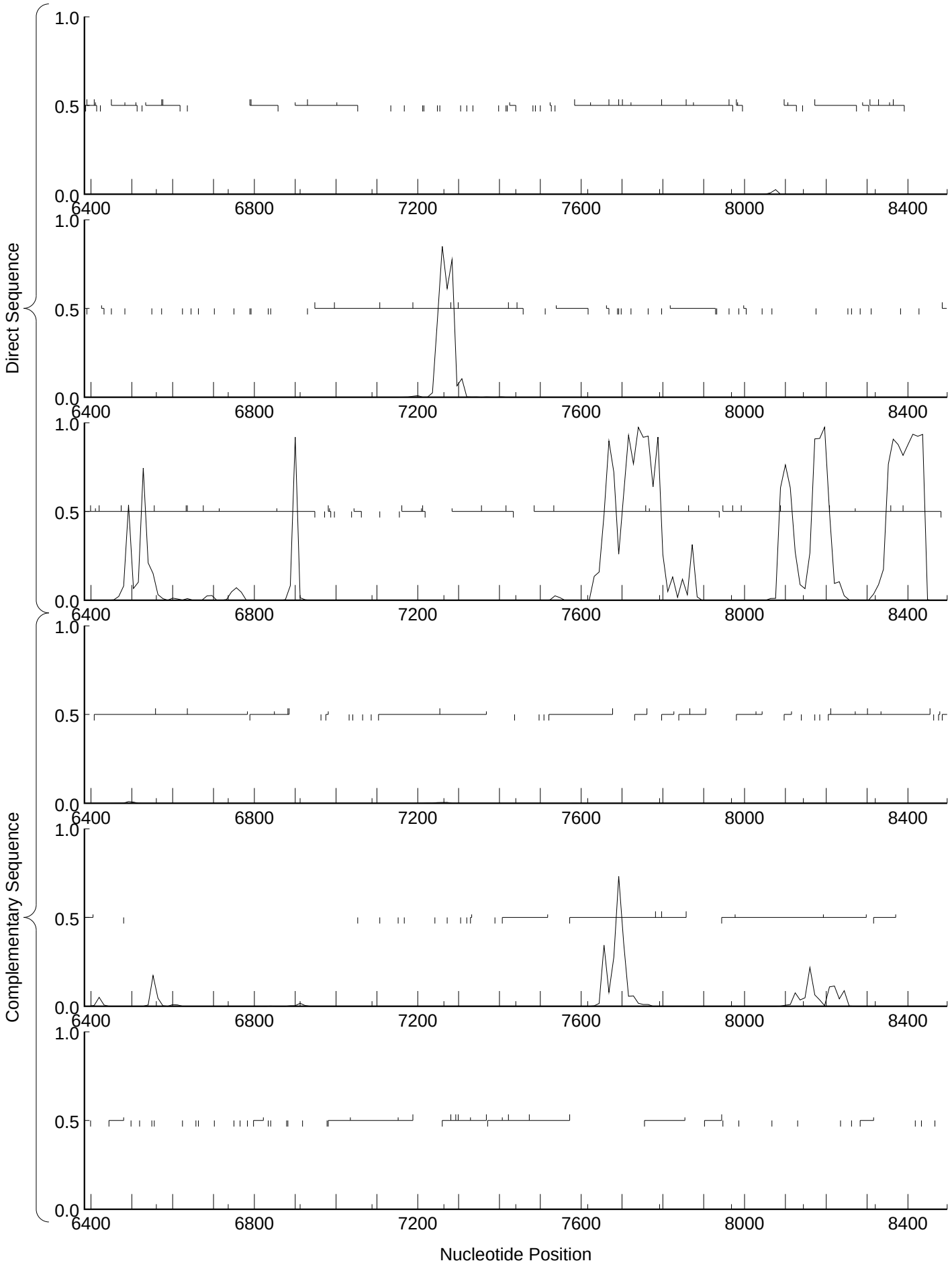


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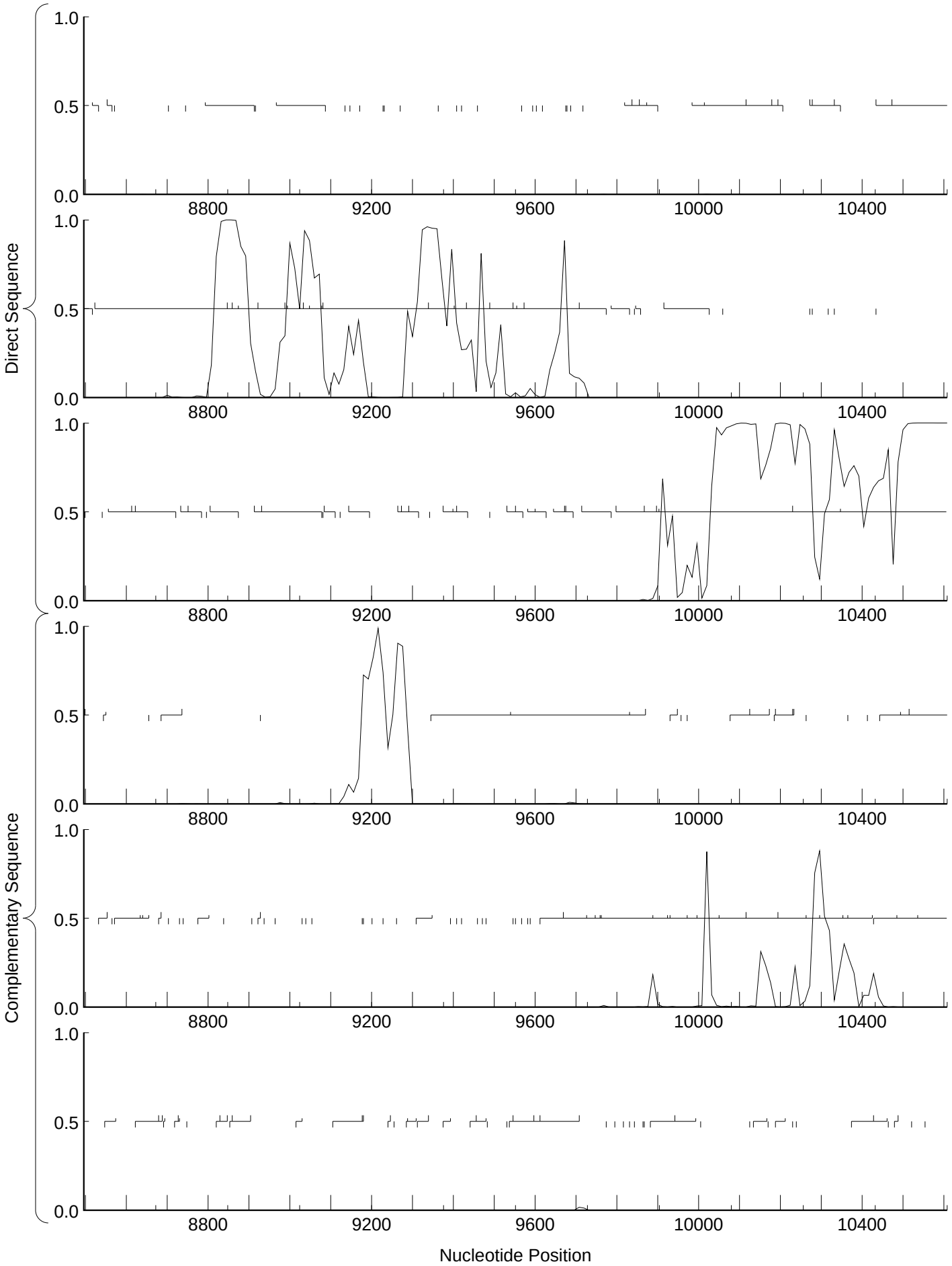



Direct Sequence
Complementary Sequence
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0.5
0.0
4400
4800
5200
5600
6000
Nucleotide Position

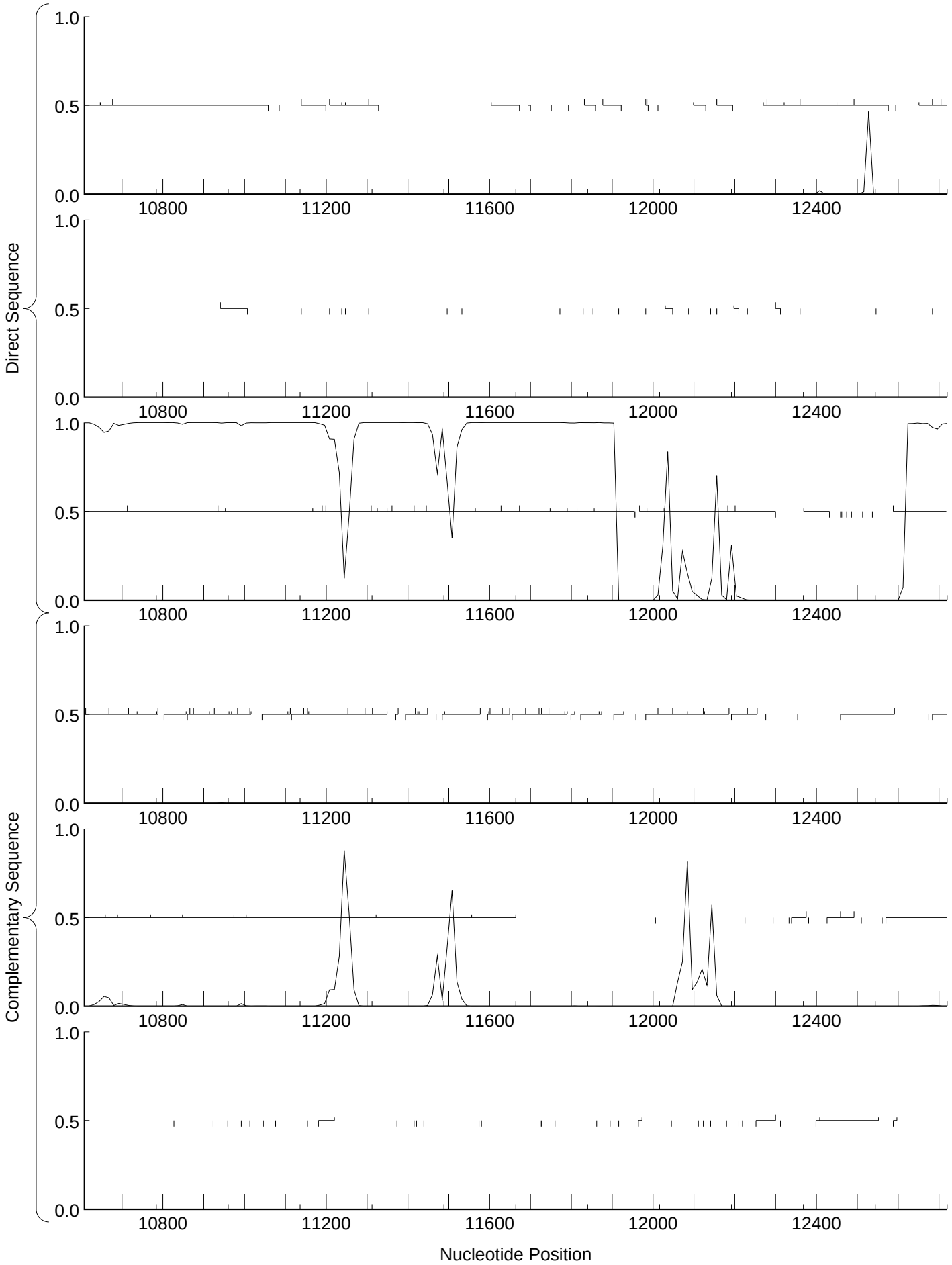
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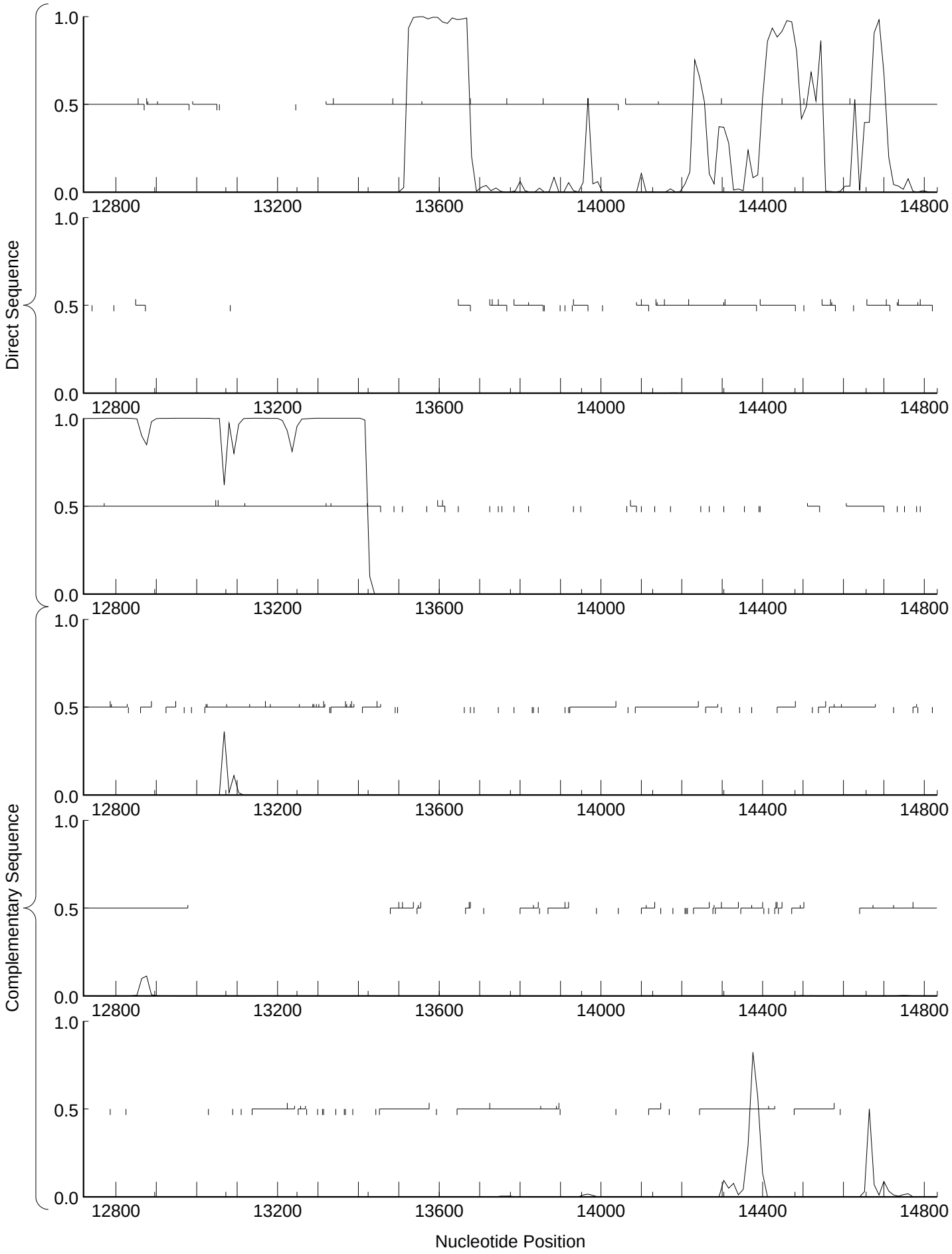
ked complete sequence, 57862 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AU, Order 4, Window 96, Step 12, 6/29



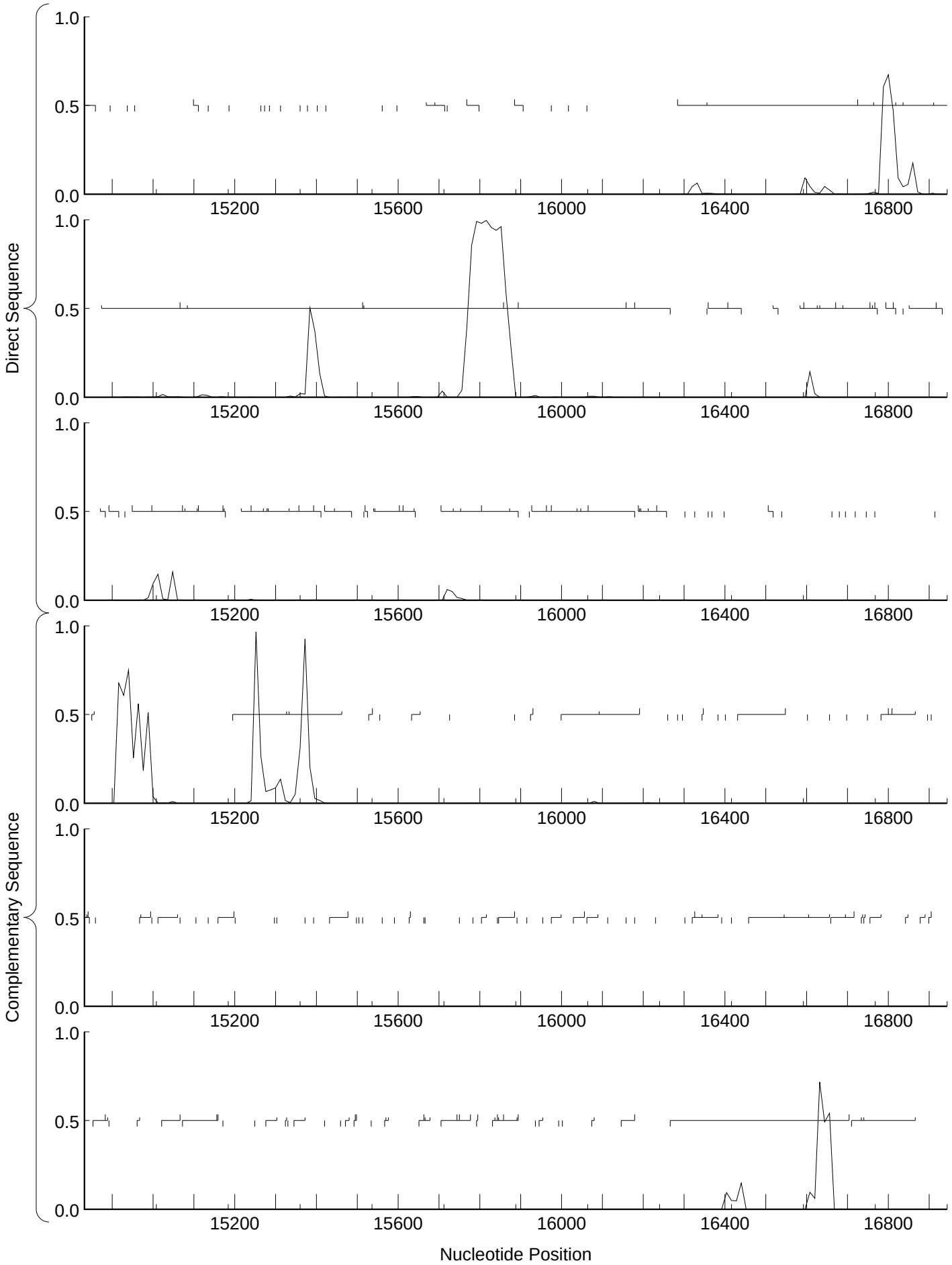
ked complete sequence, 57862 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AU, Order 4, Window 96, Step 12, 7/29



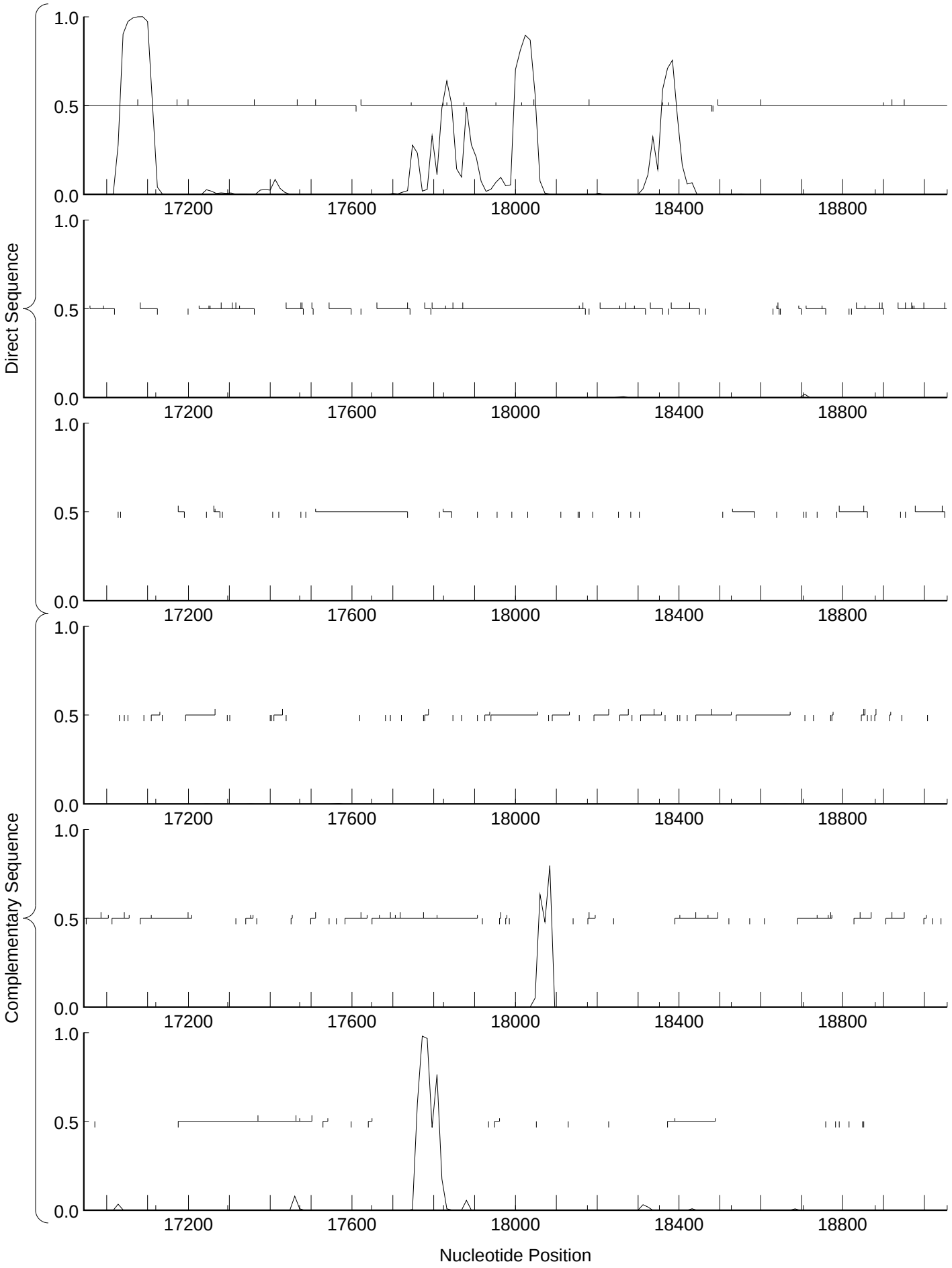
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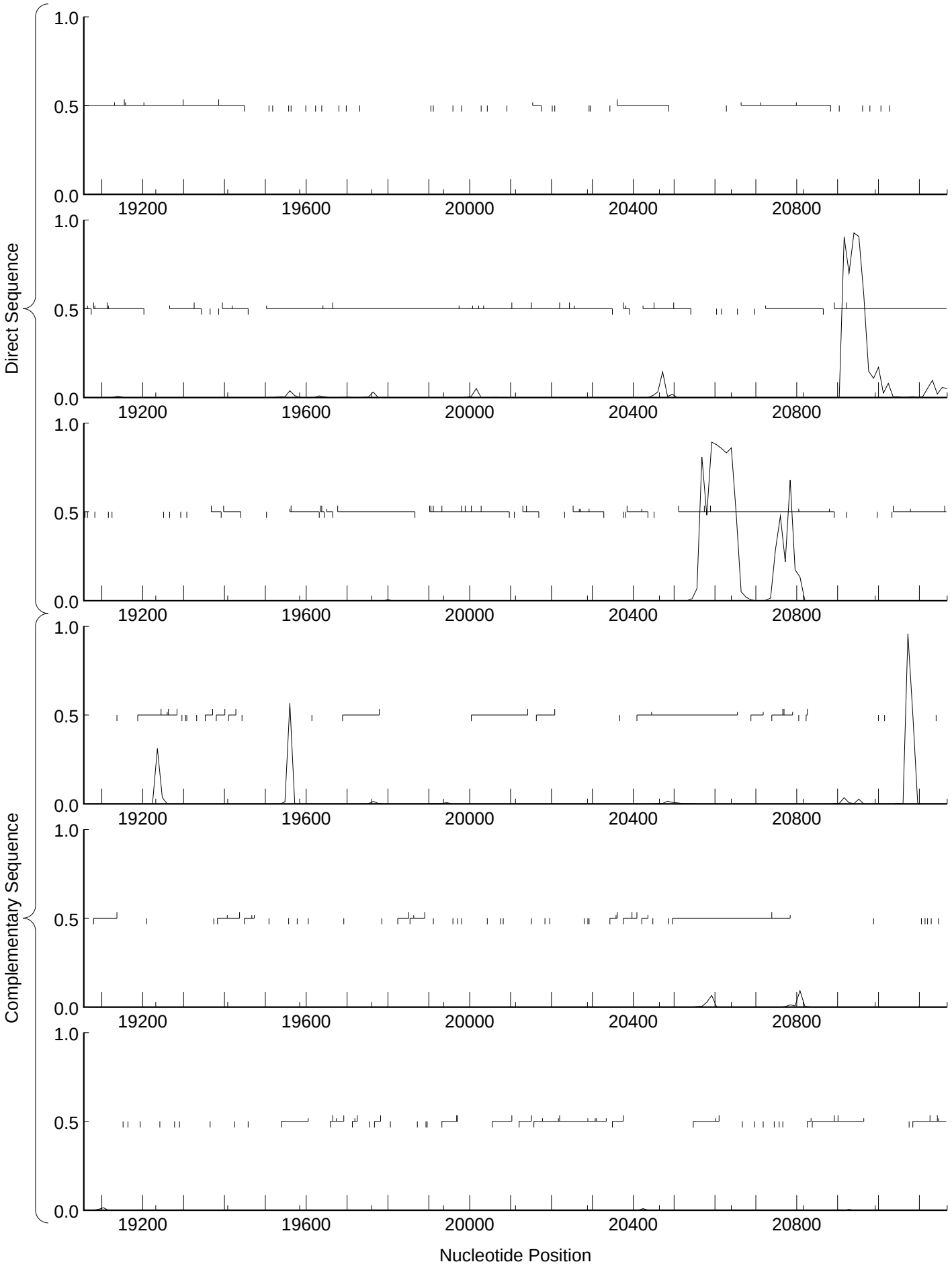
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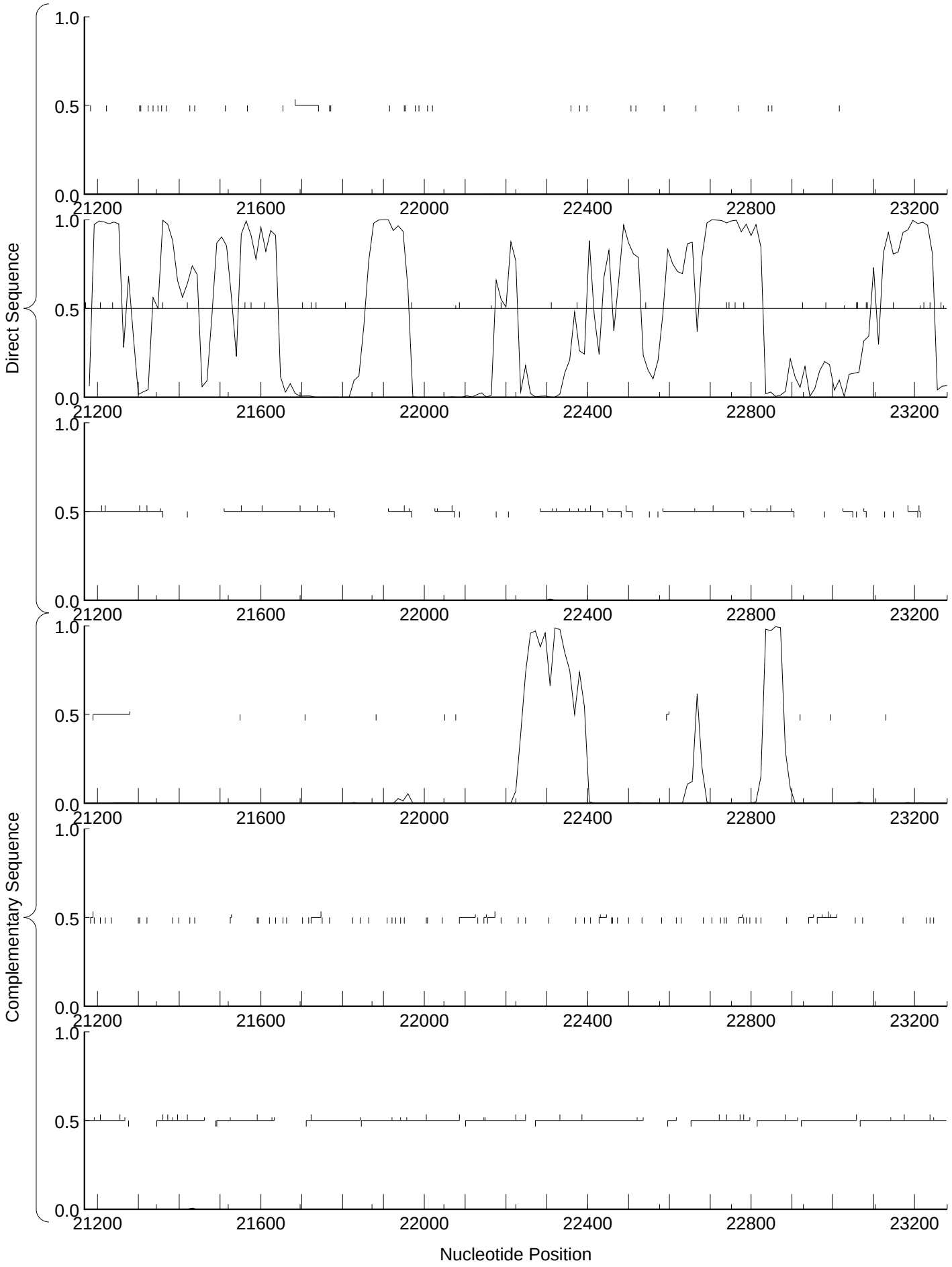
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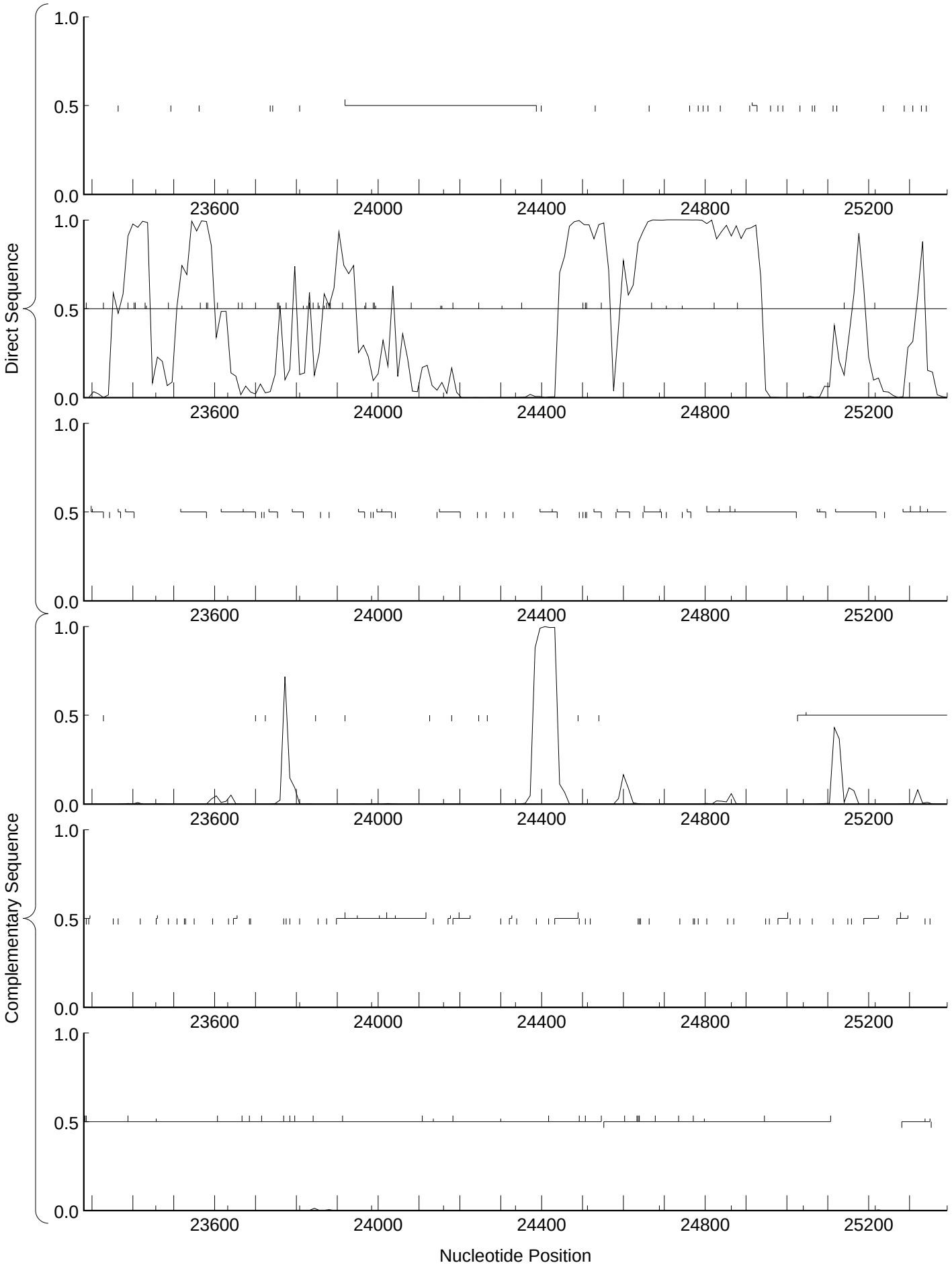
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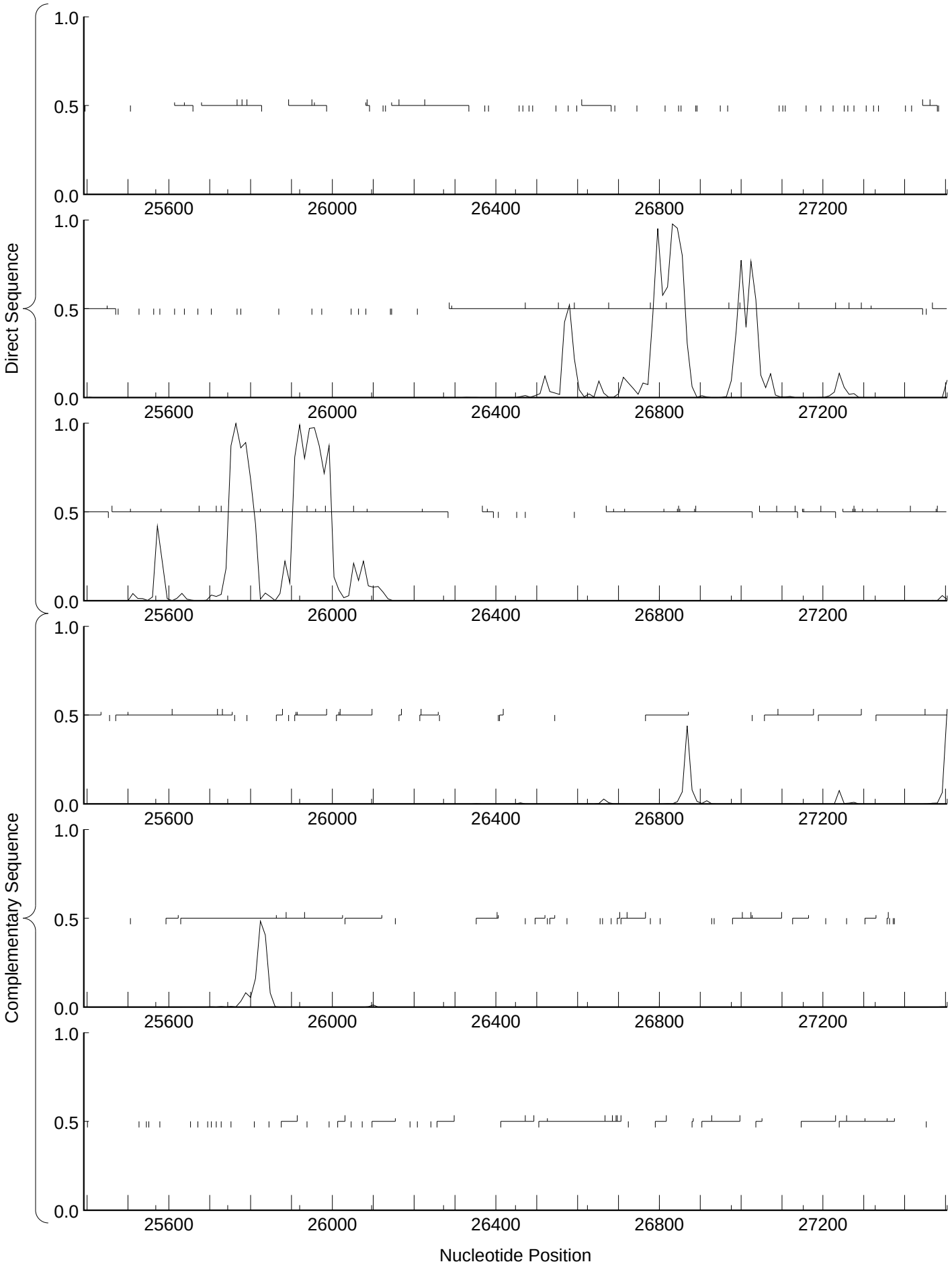
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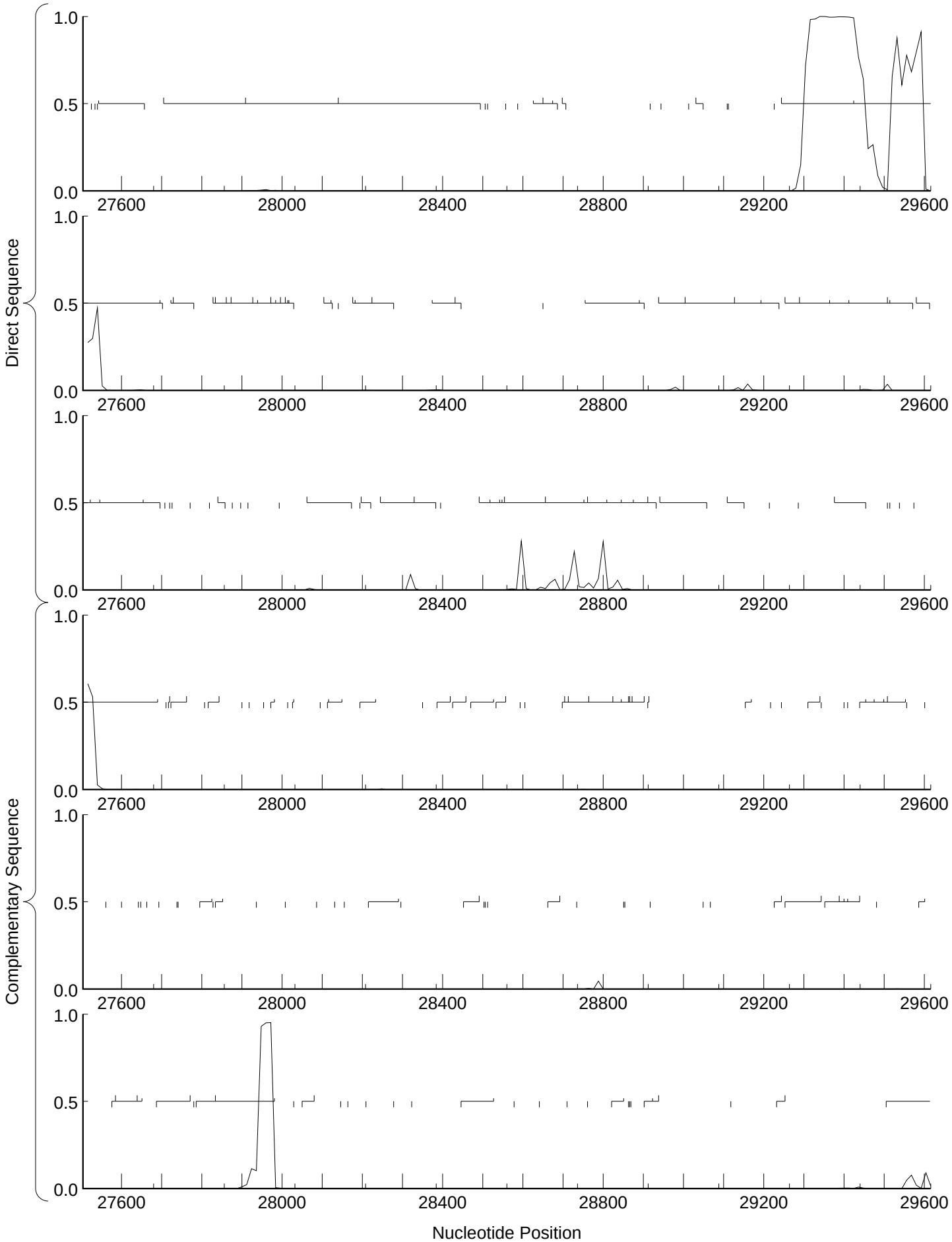
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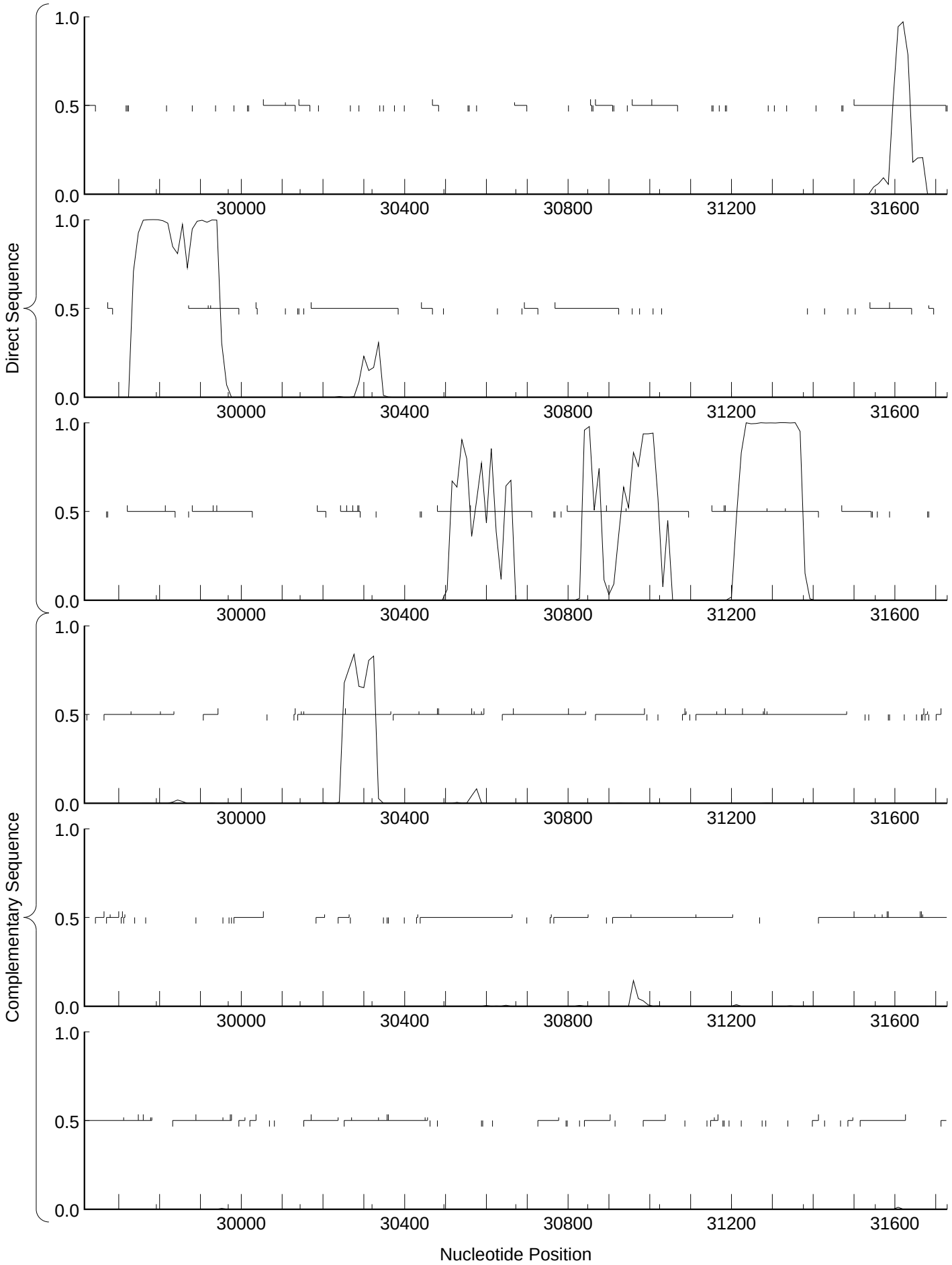
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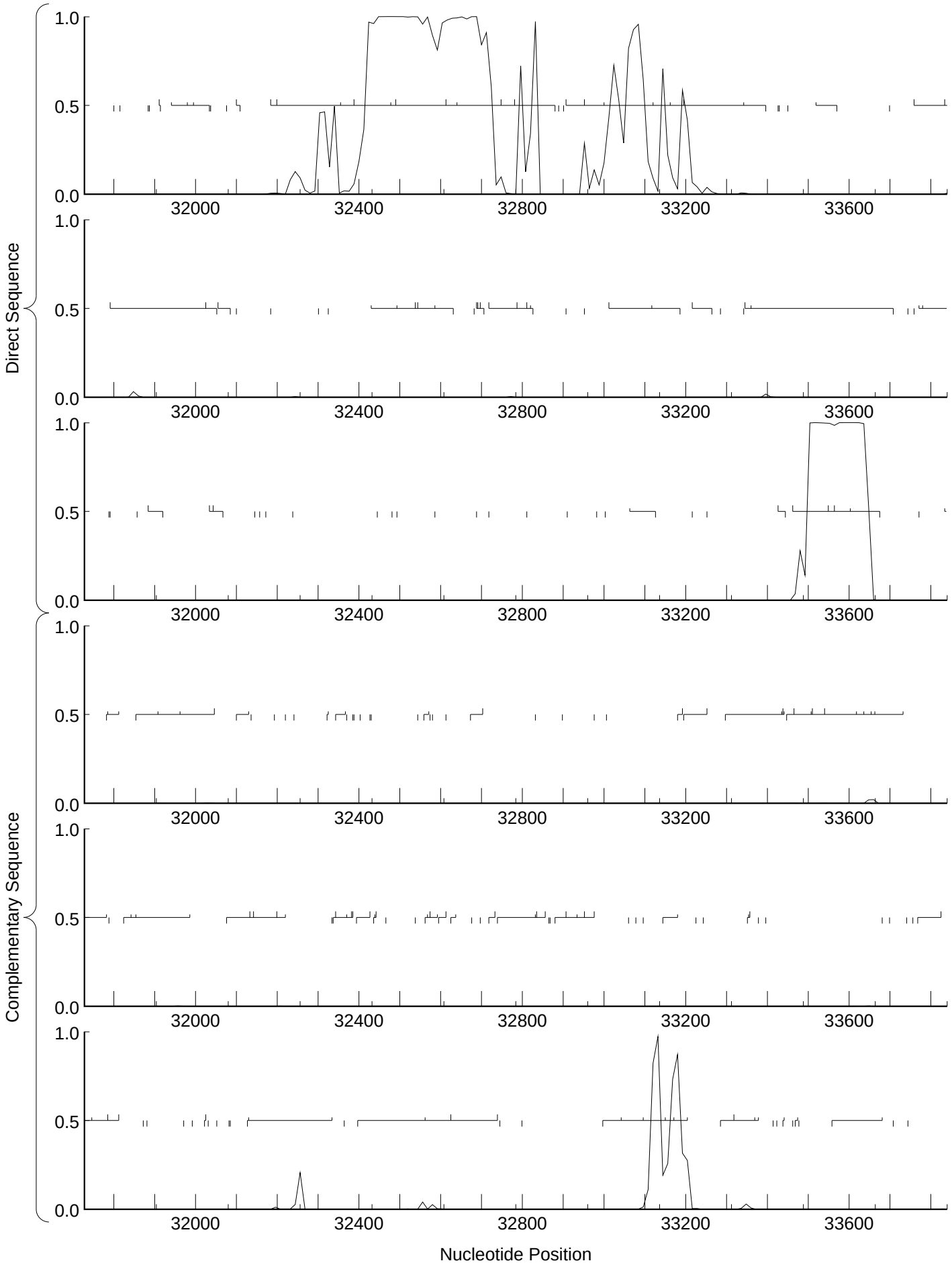
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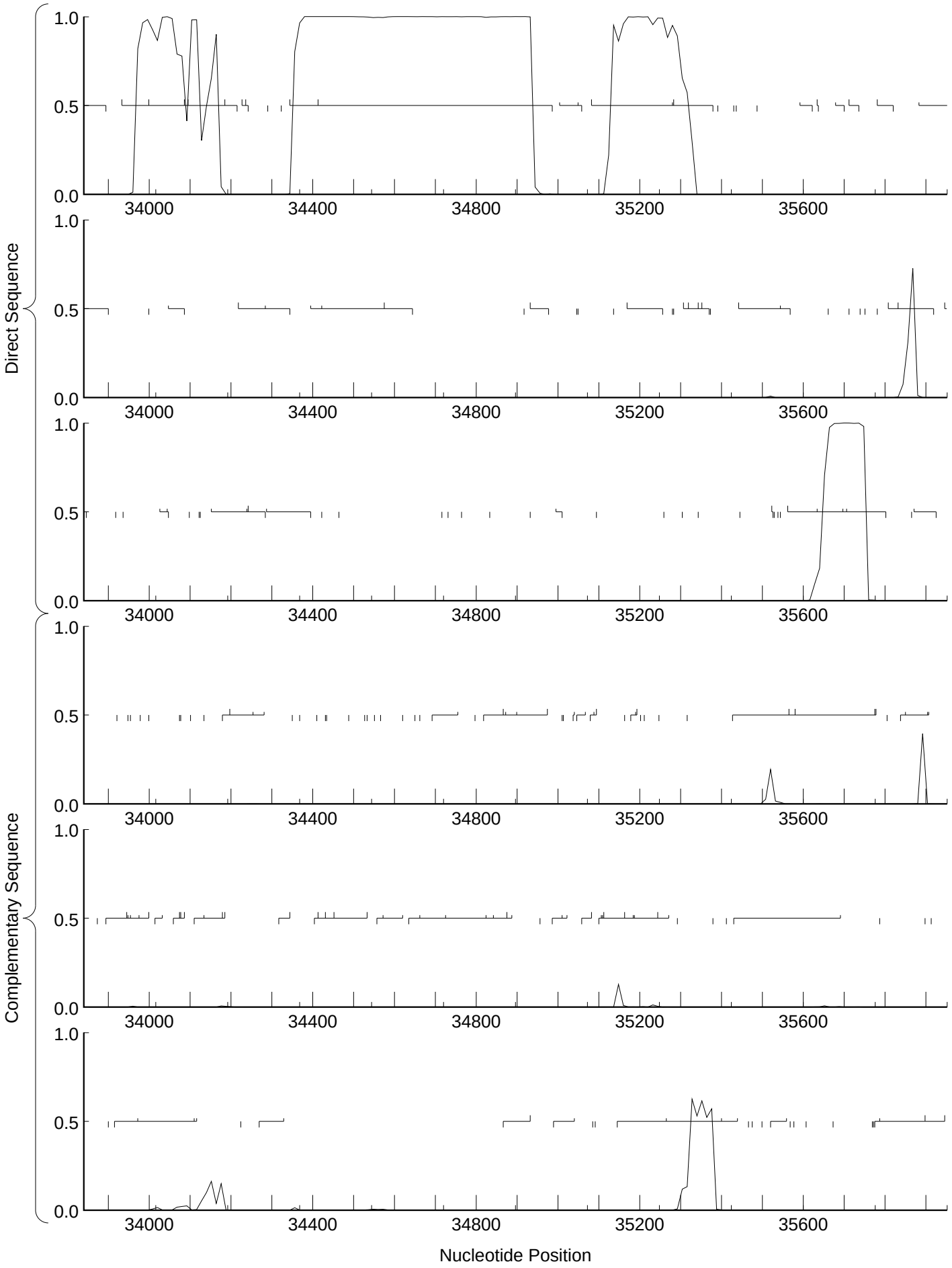
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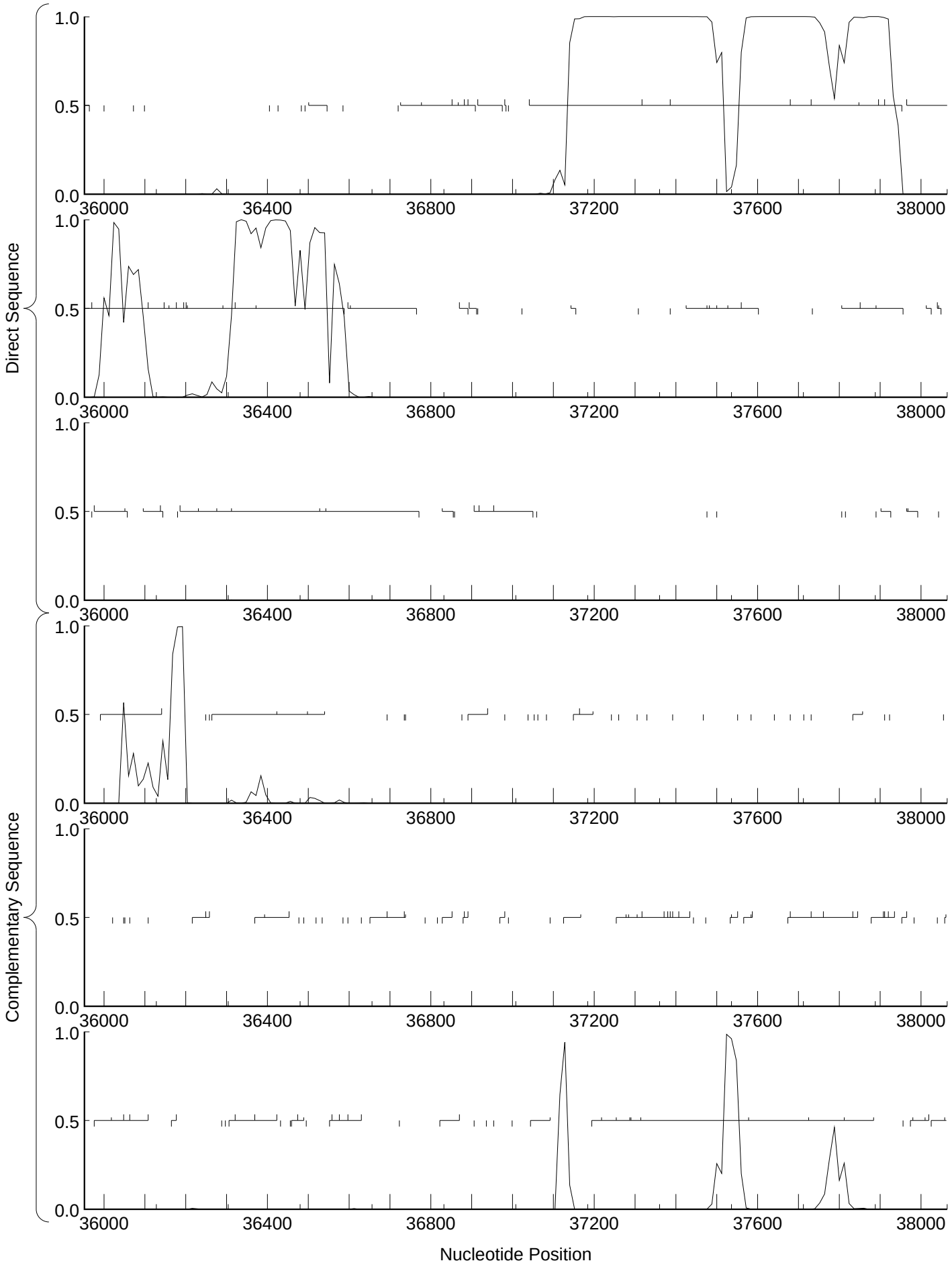
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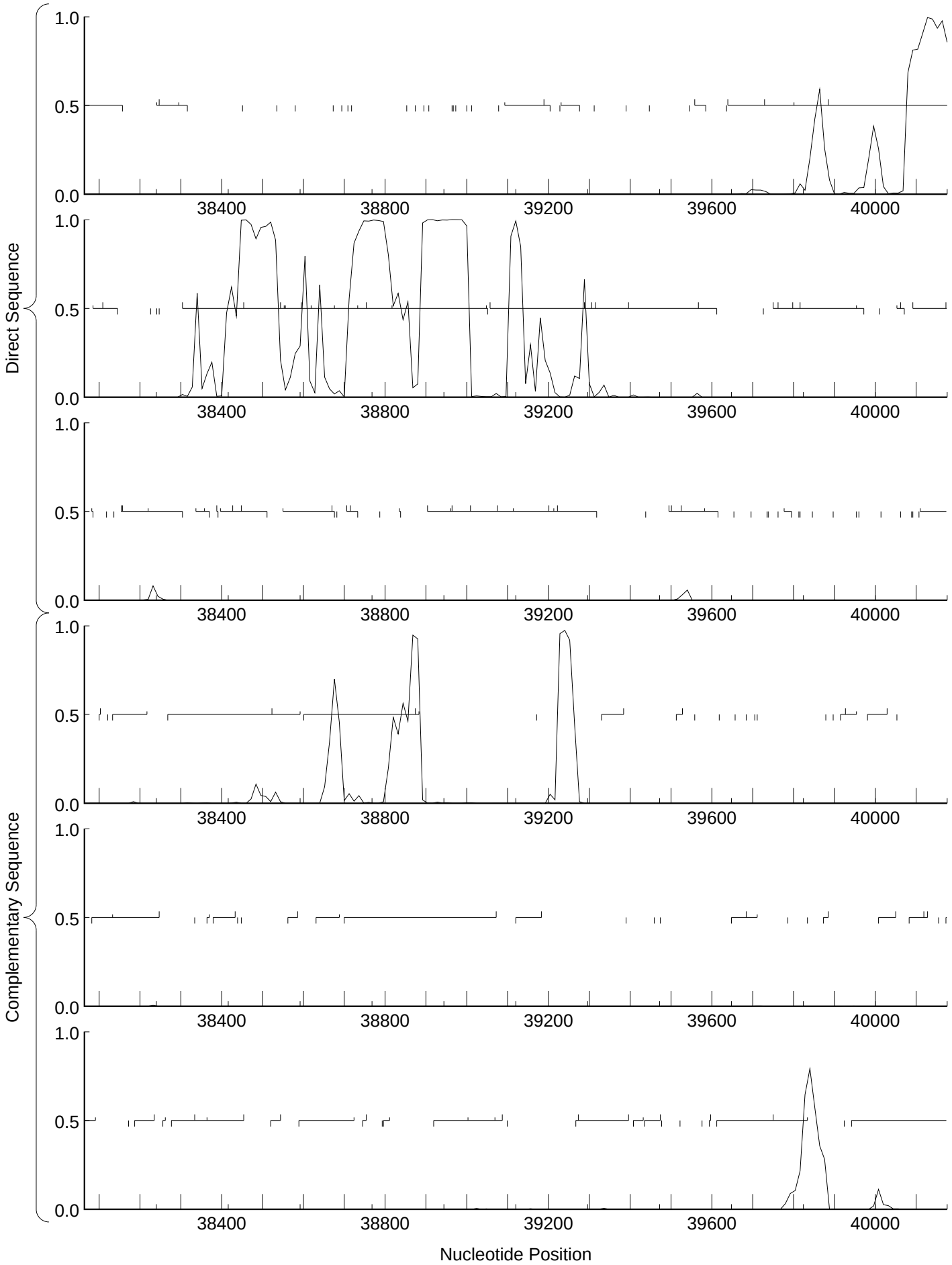


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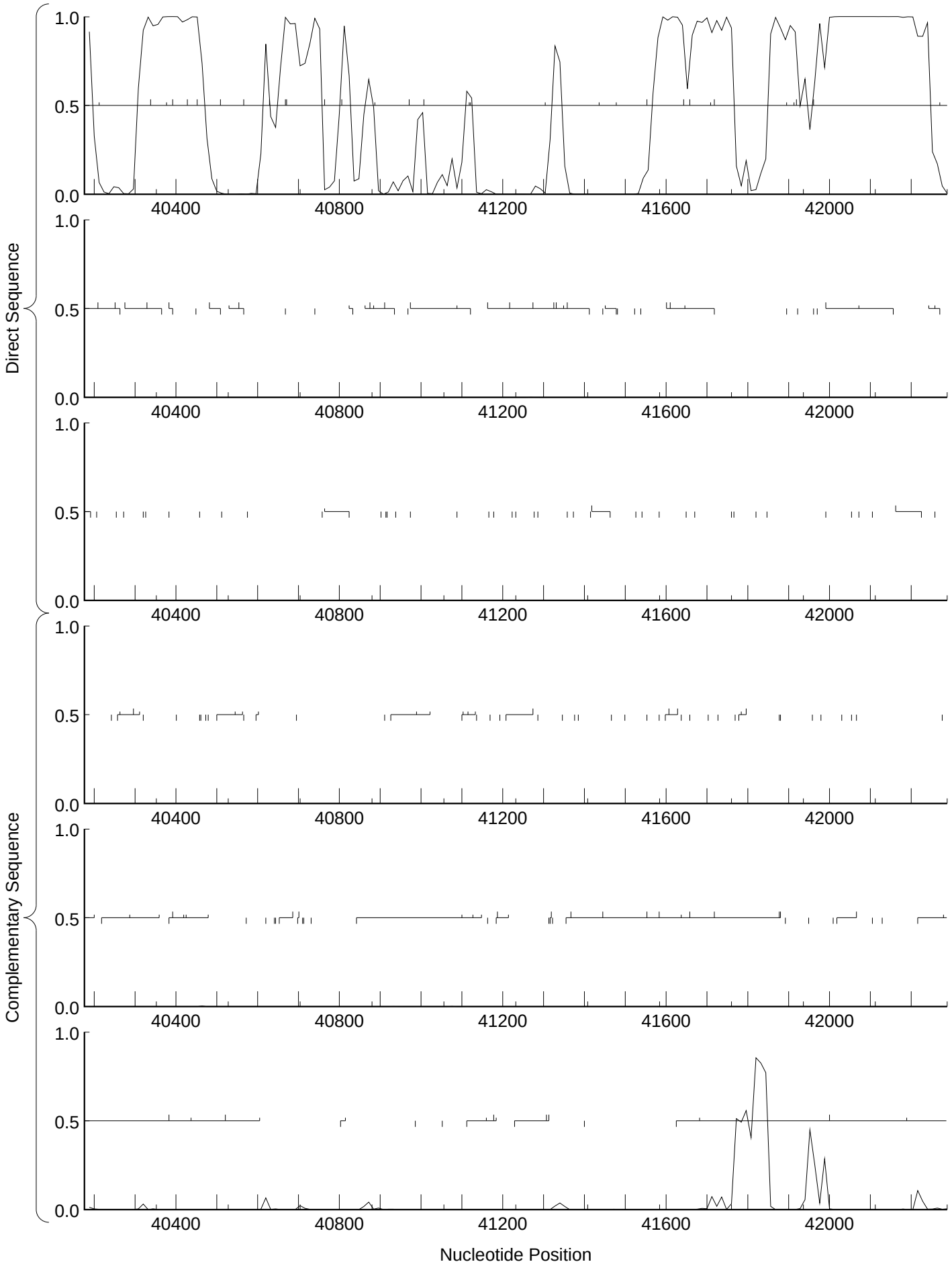


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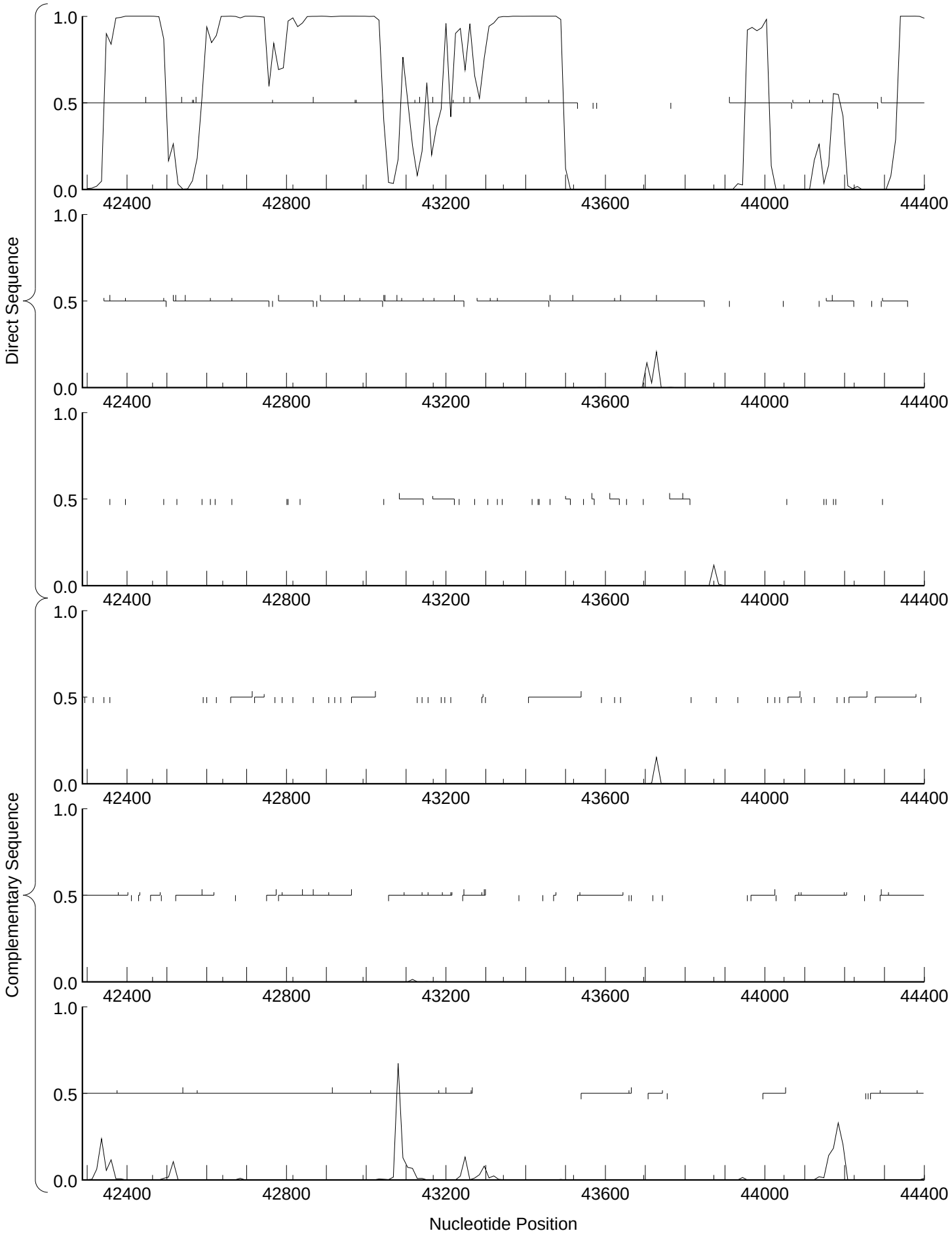



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

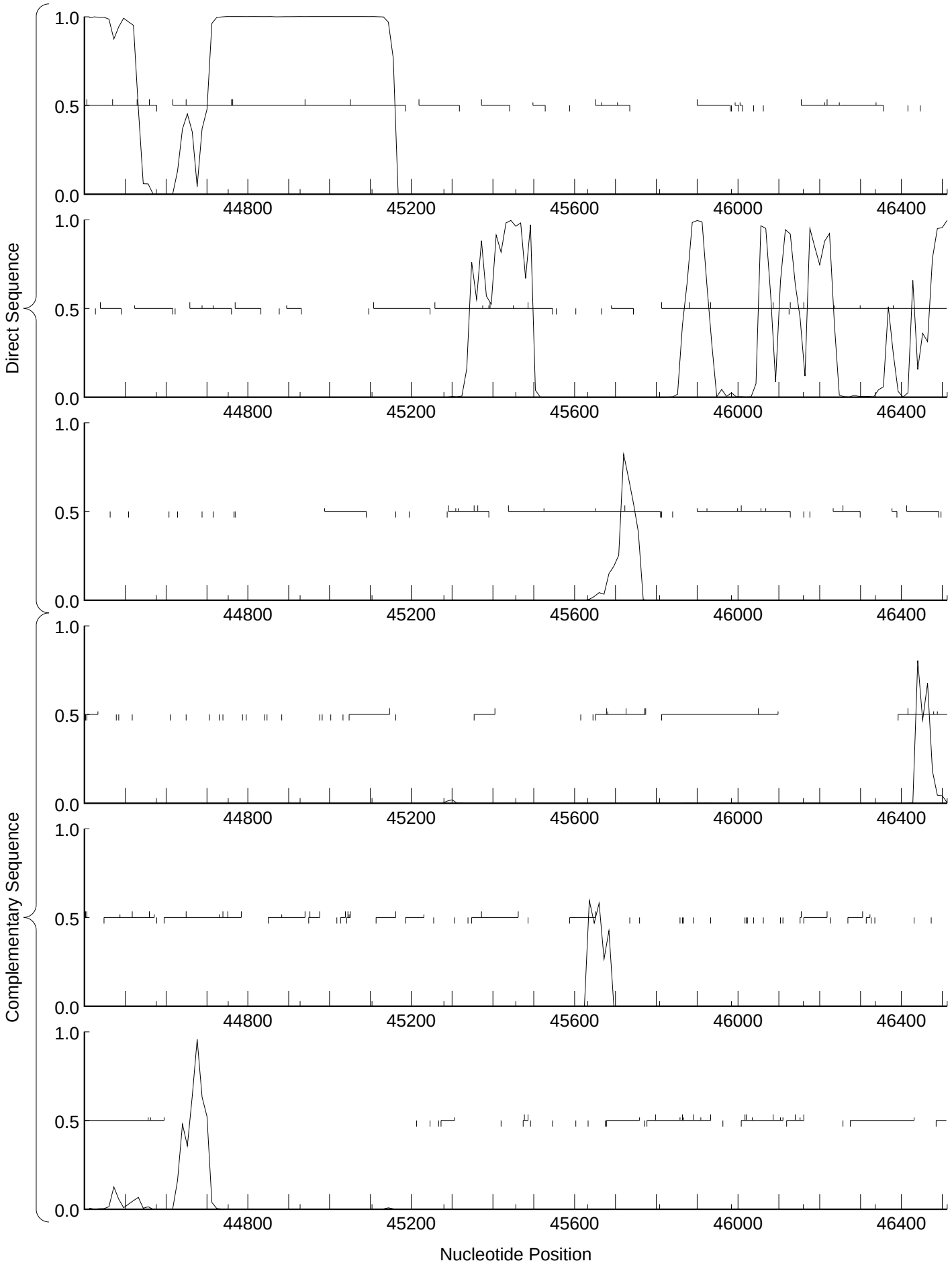
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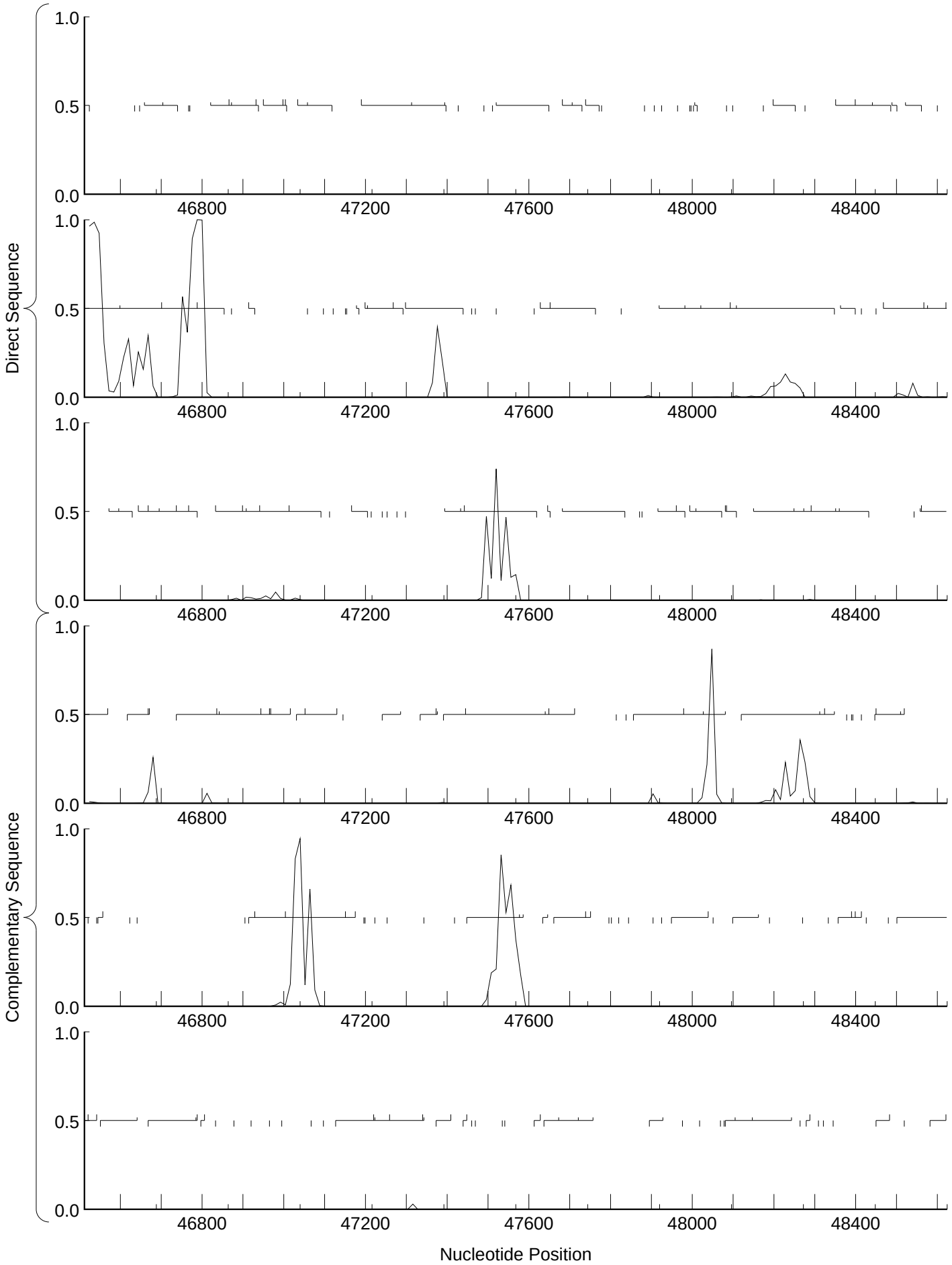
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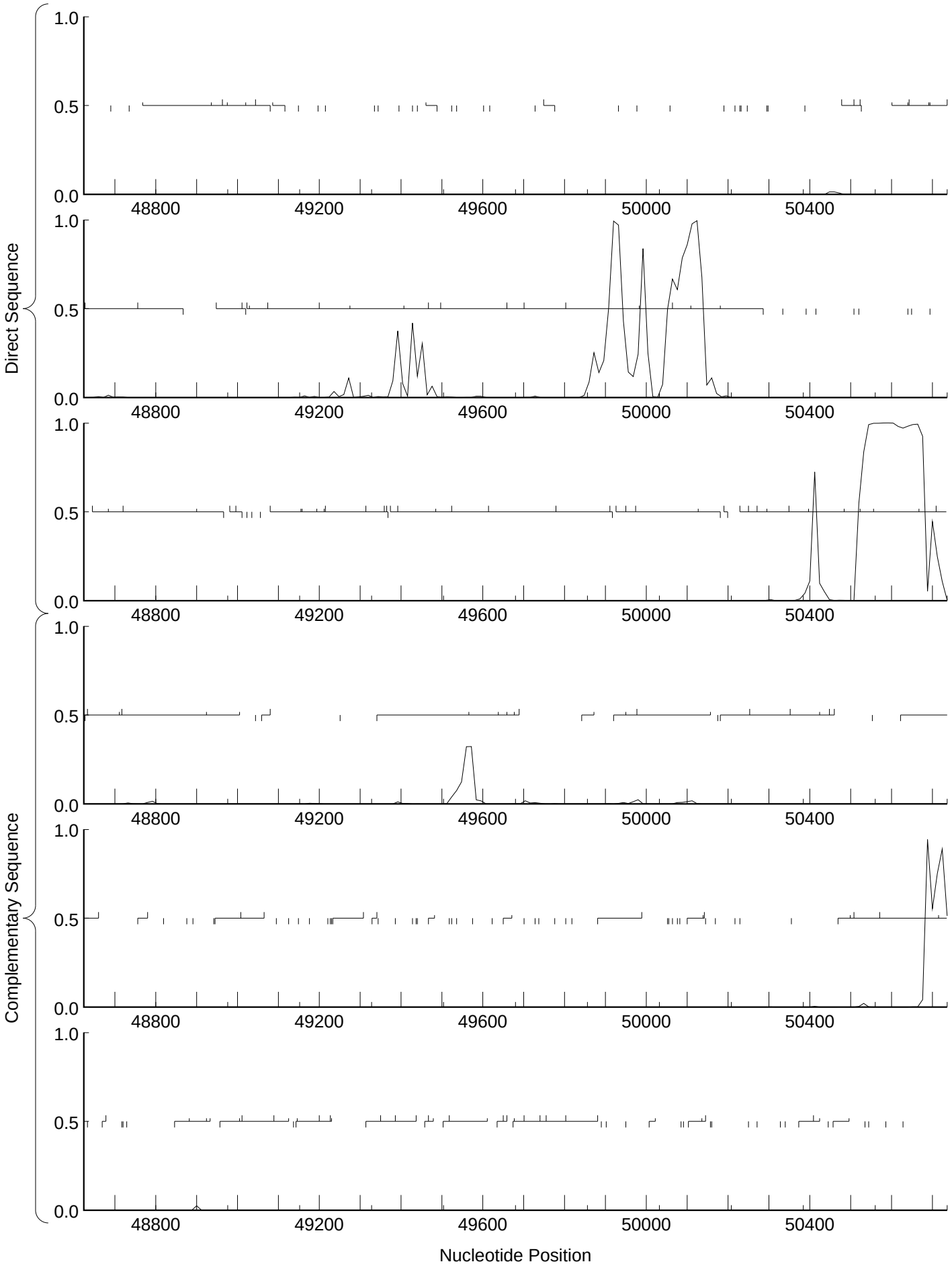
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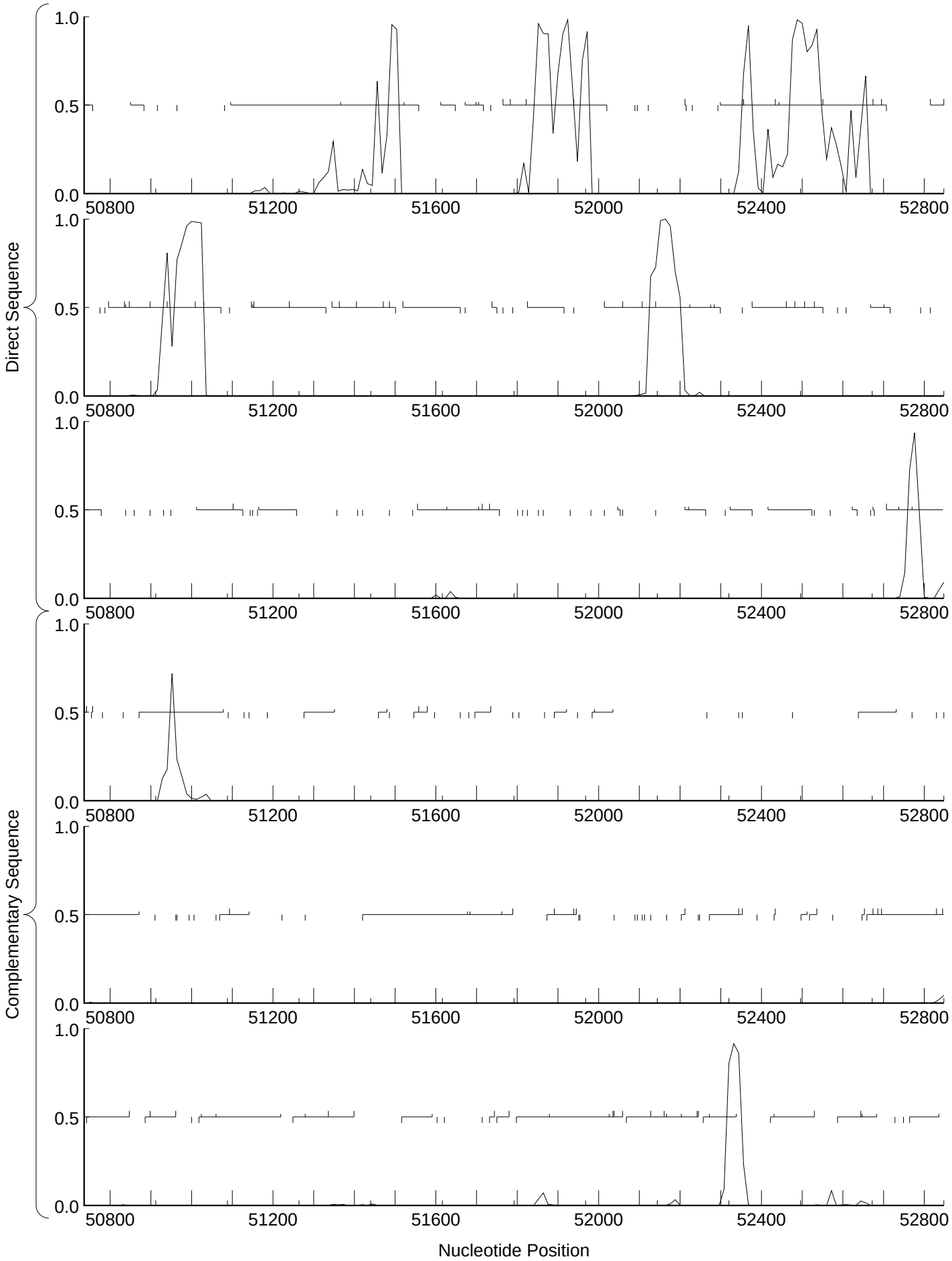
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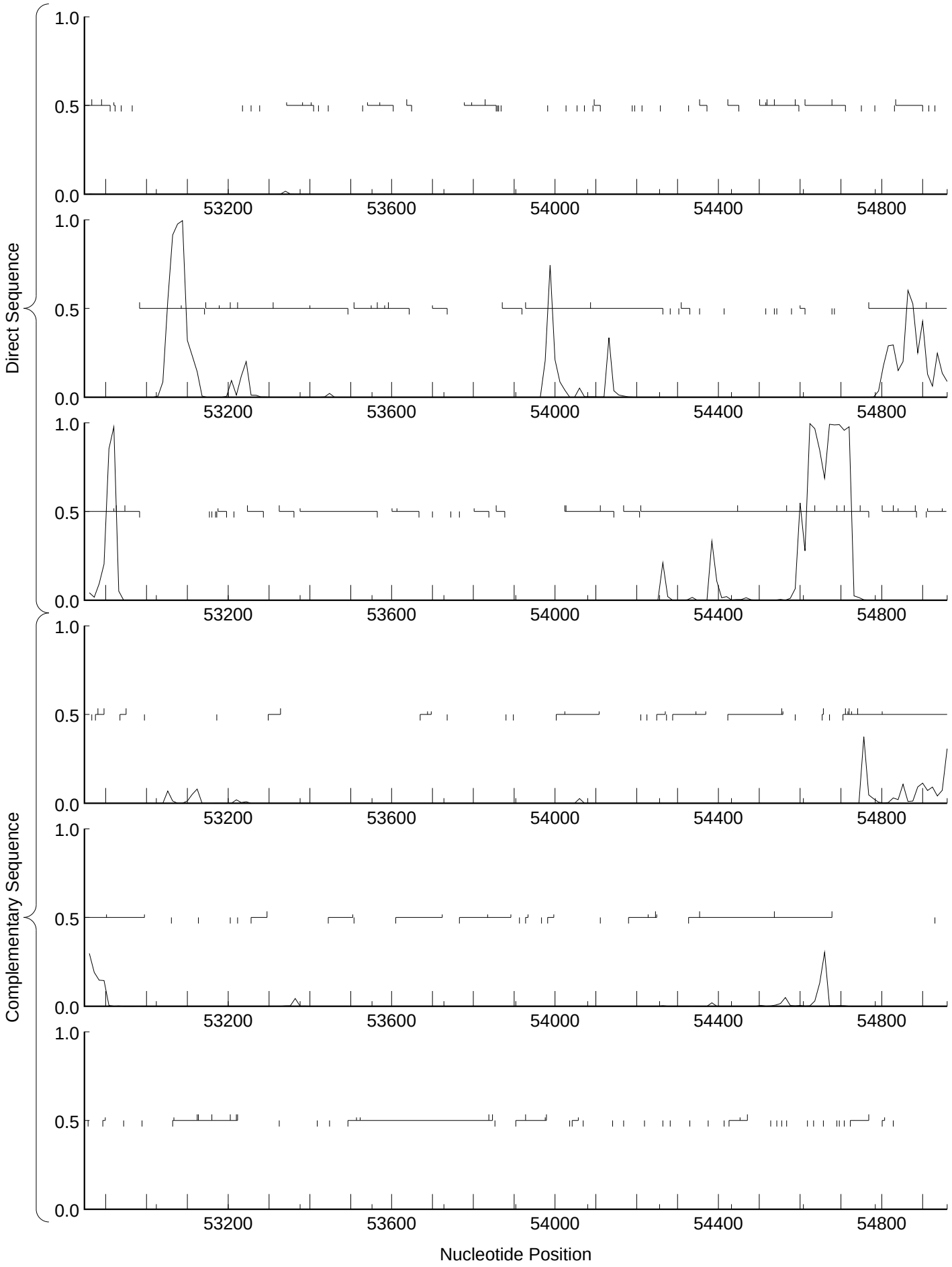
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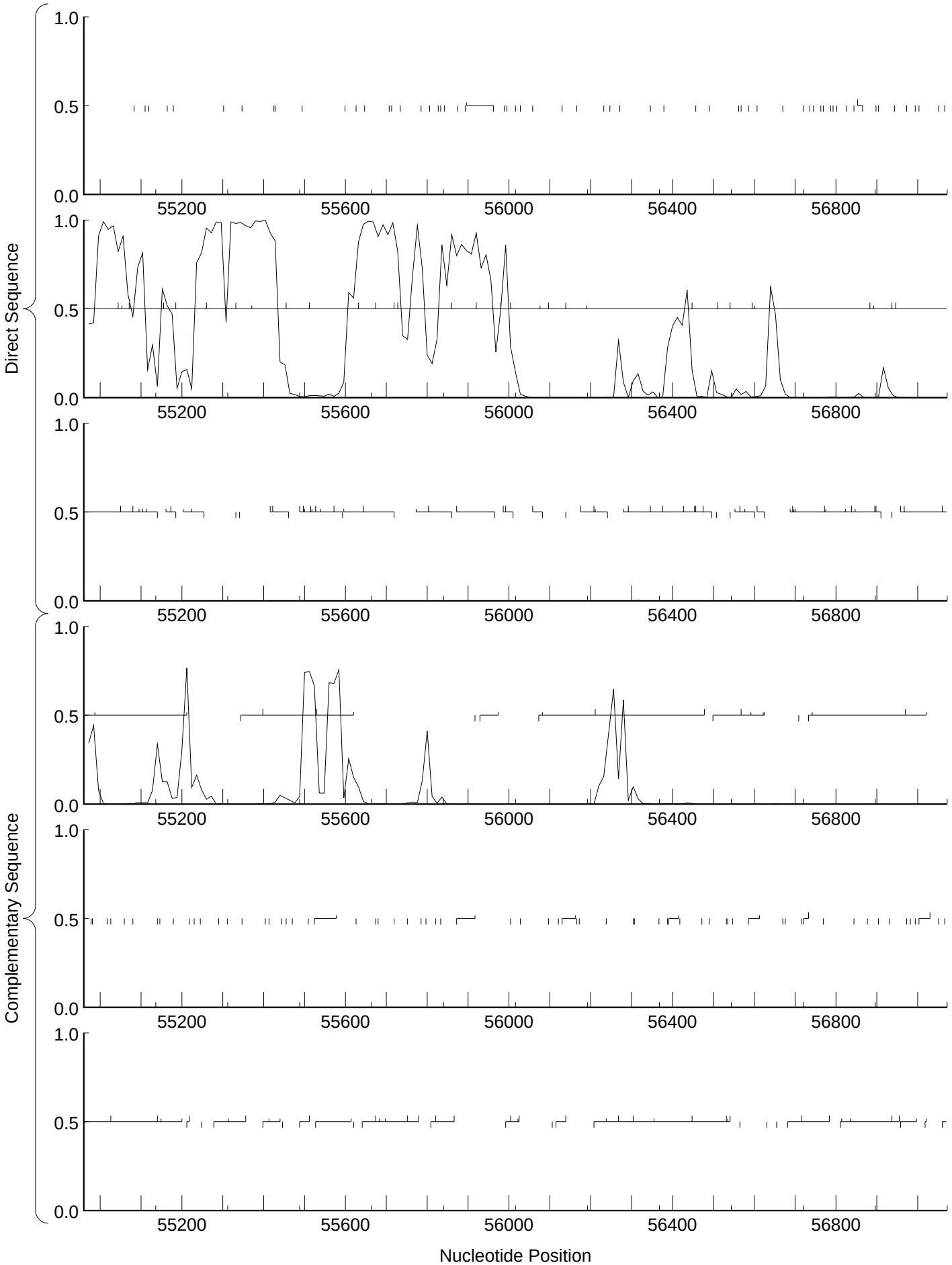
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ed complete sequence, 57862 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AU, Order 4, Window 96, Step 12, 27/29



ed complete sequence, 57862 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AU, Order 4, Window 96, Step 12, 28/29



ed complete sequence, 57862 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AU, Order 4, Window 96, Step 12, 29/29

