GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Herbertwm complete sequence, 51546 bp including 10-base 3’ overhang (CG
Analysis Date    :  3/29/24 at 6:08:58
Pages            :  26
Sequence Length  :  51546 bp
GC Content       :   62.60%

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

PS-Version       :  1.2

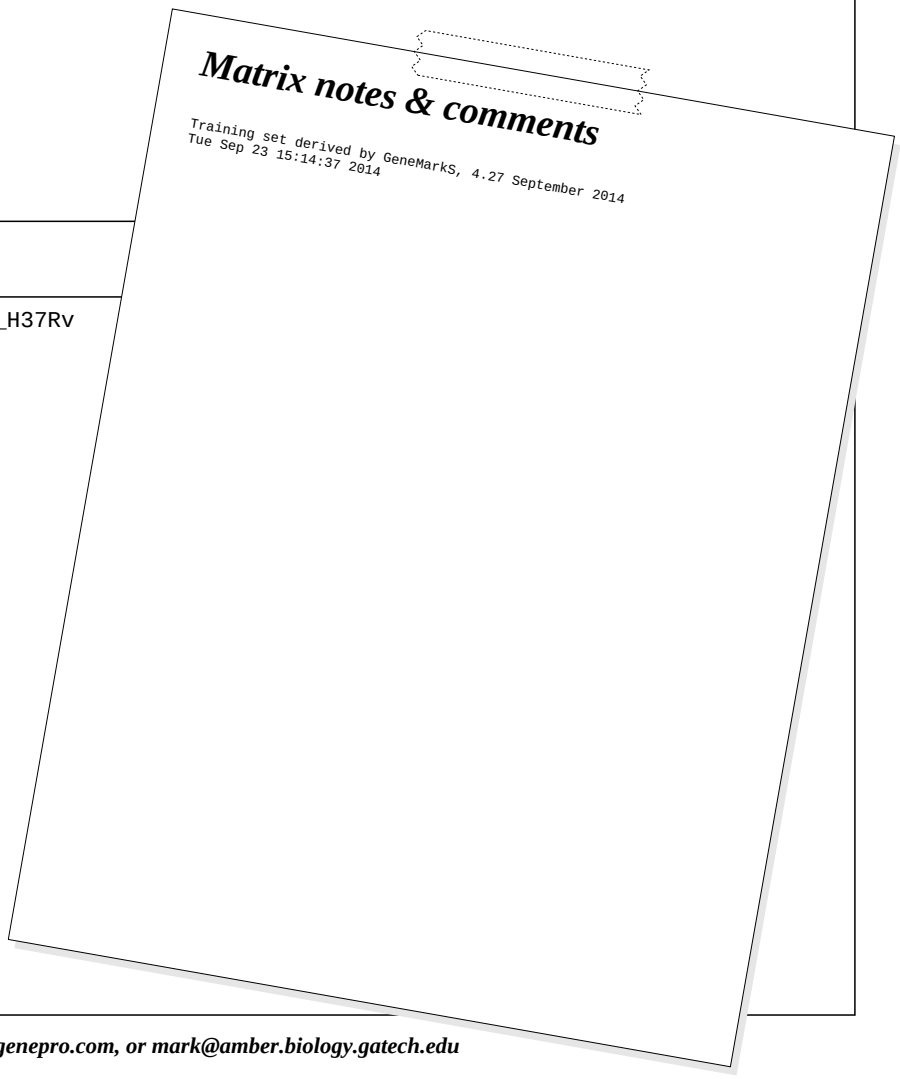
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Author     :  -

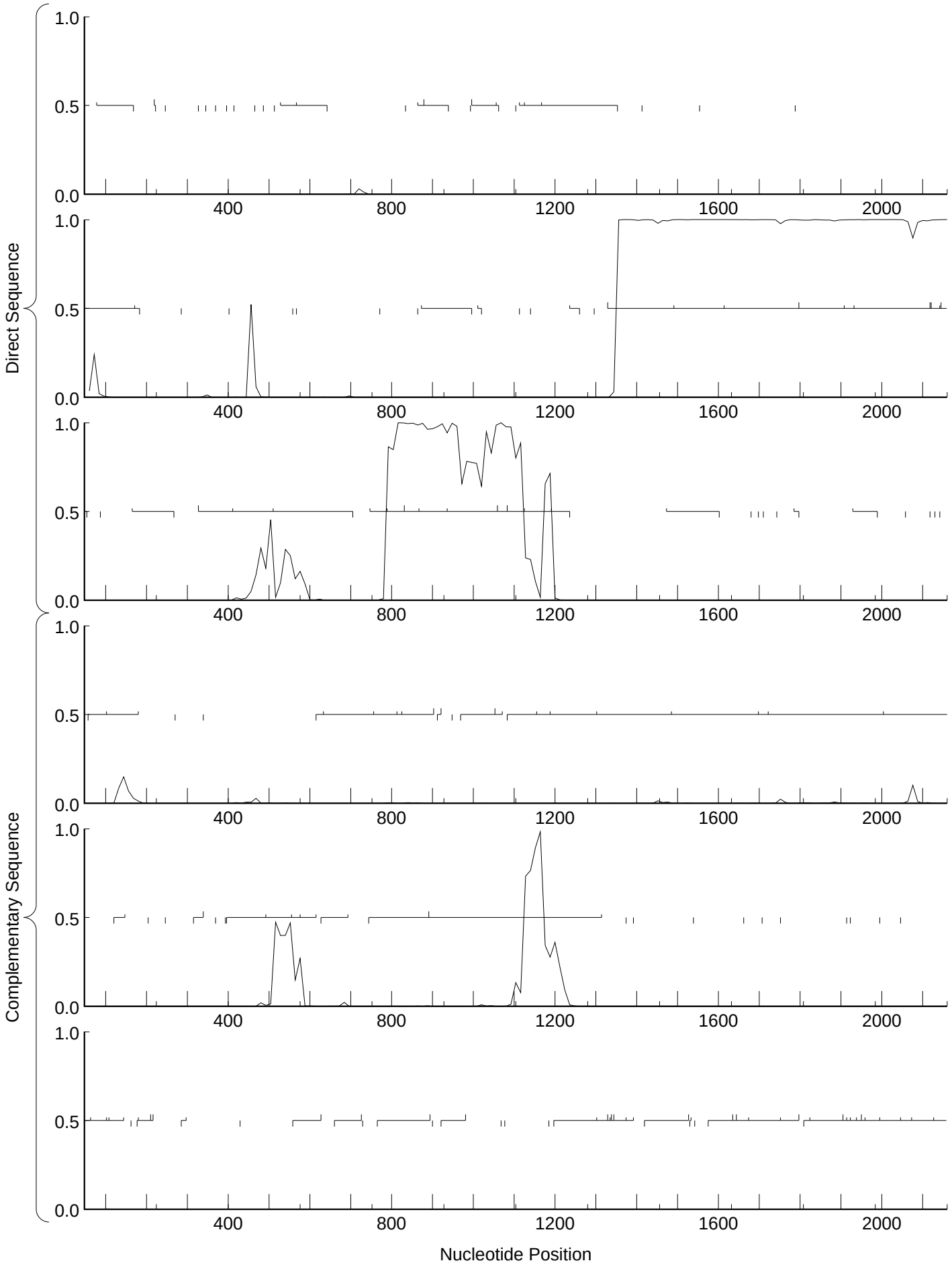
Order      :  4


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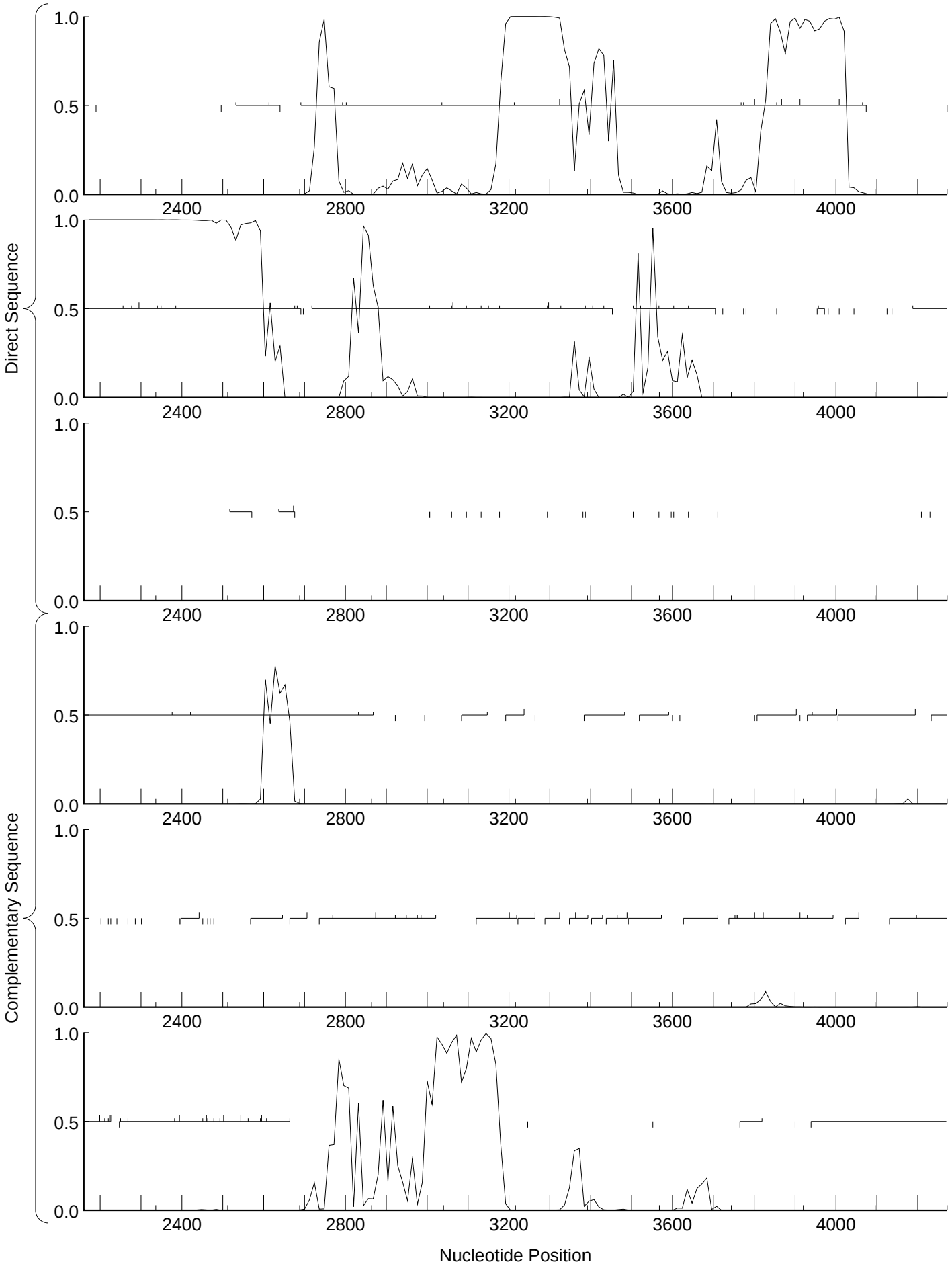


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

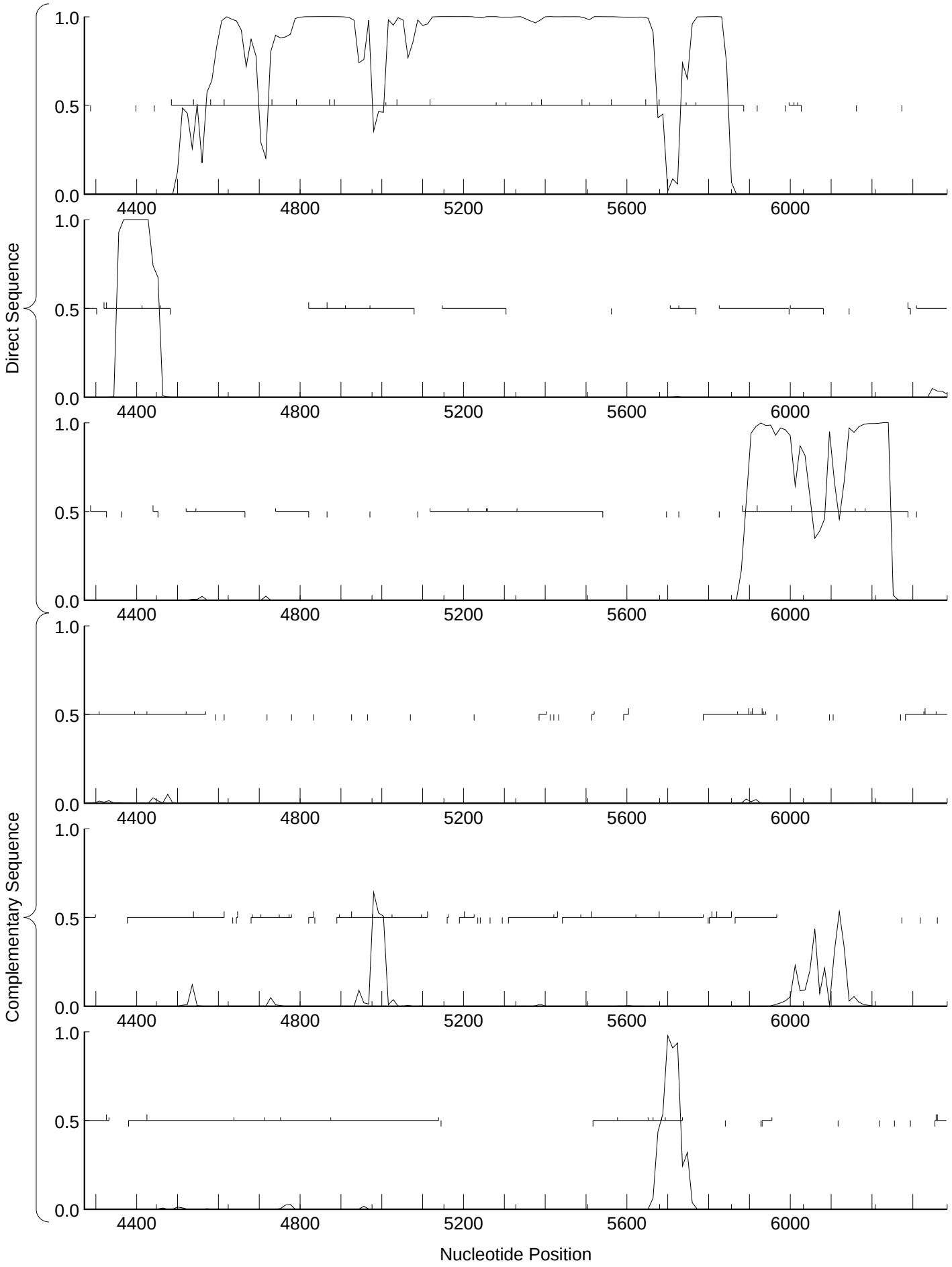
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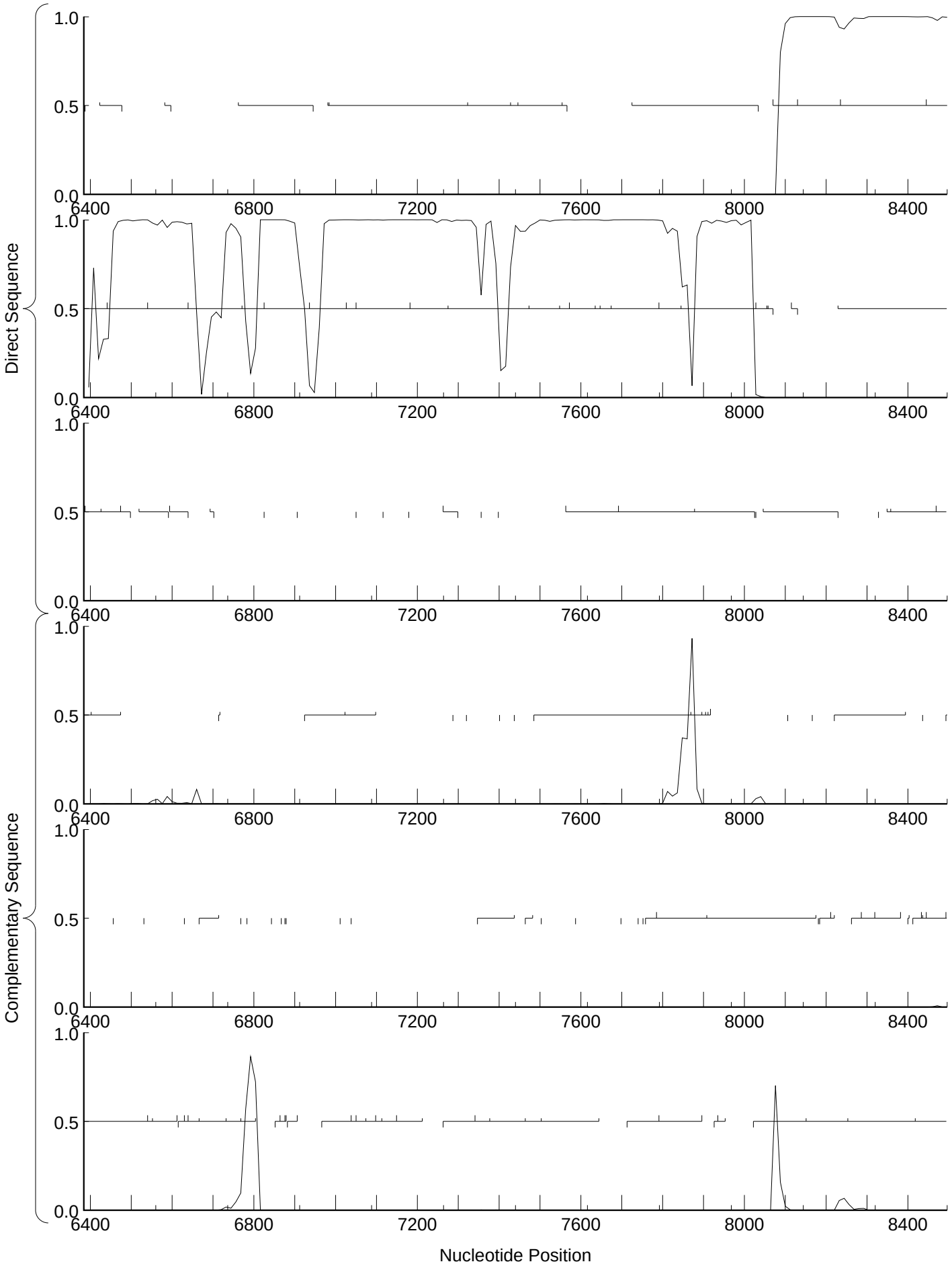
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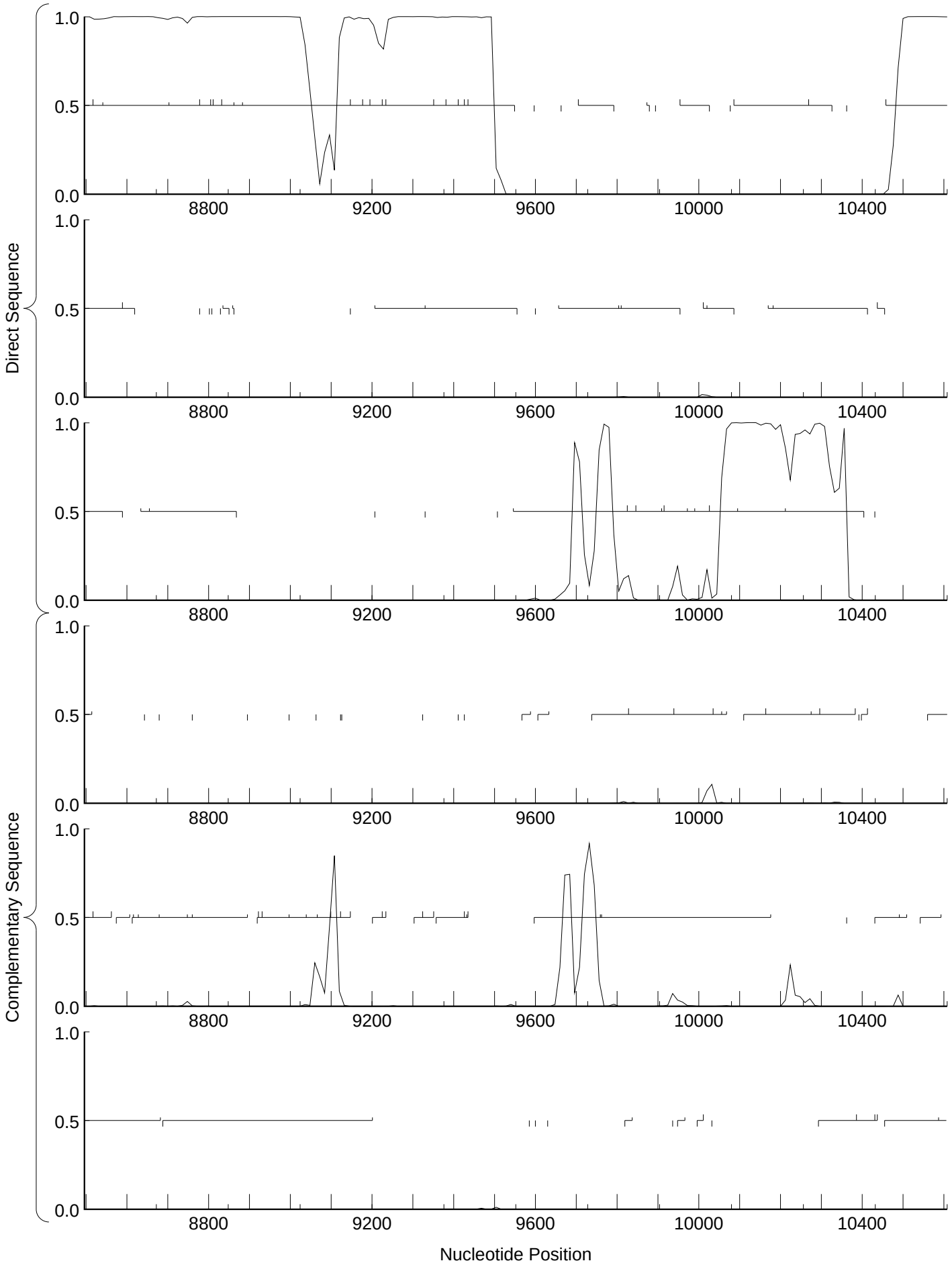
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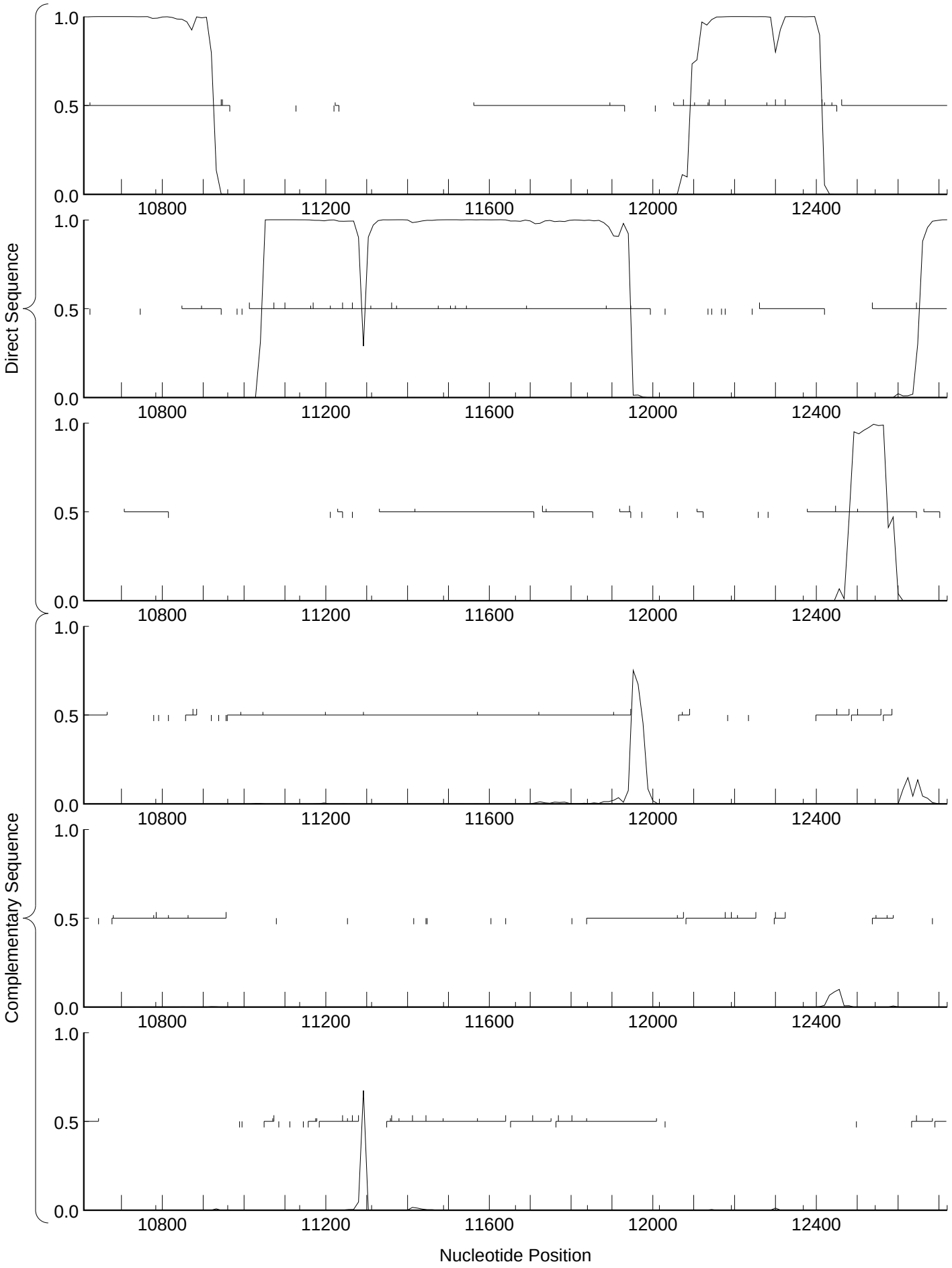
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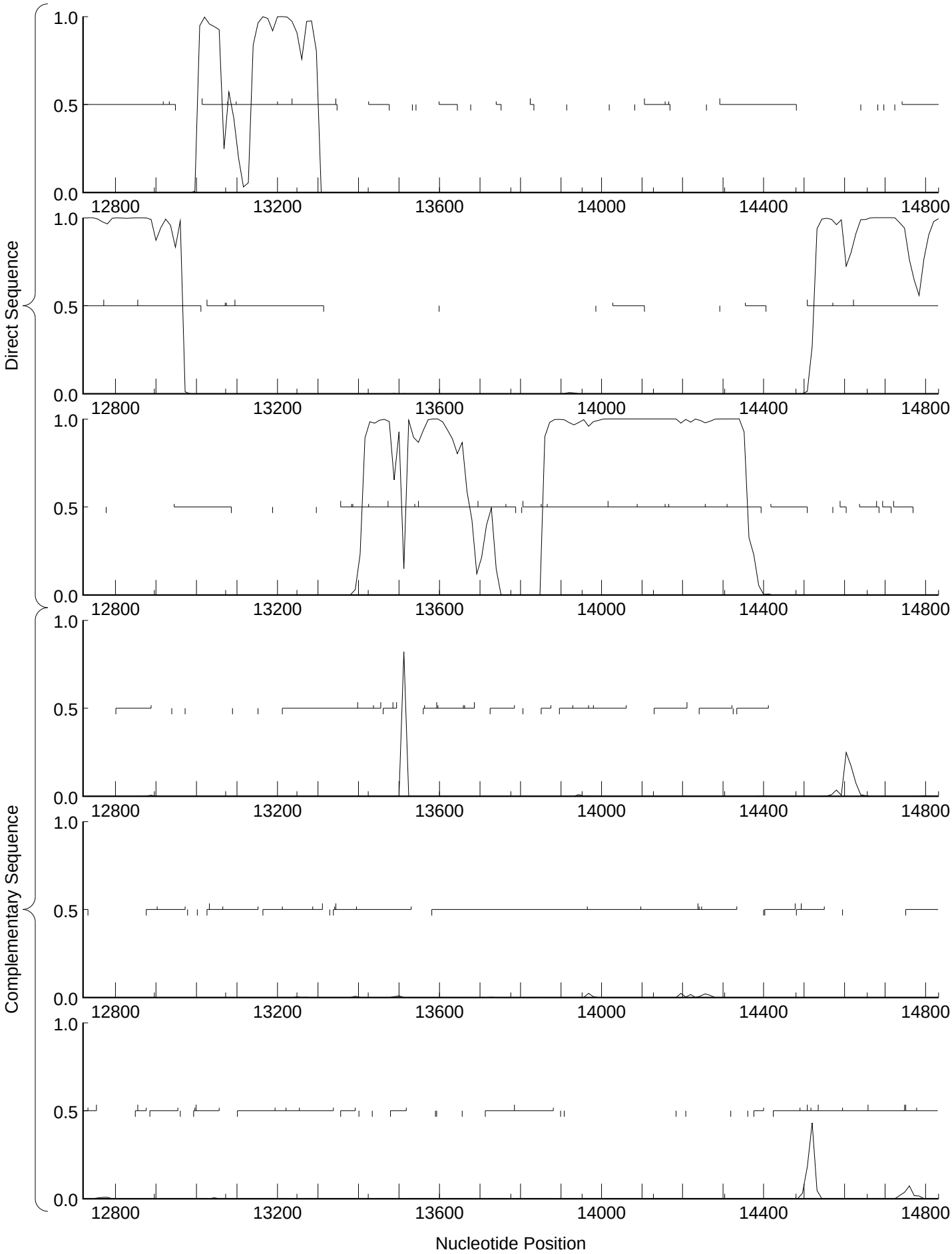
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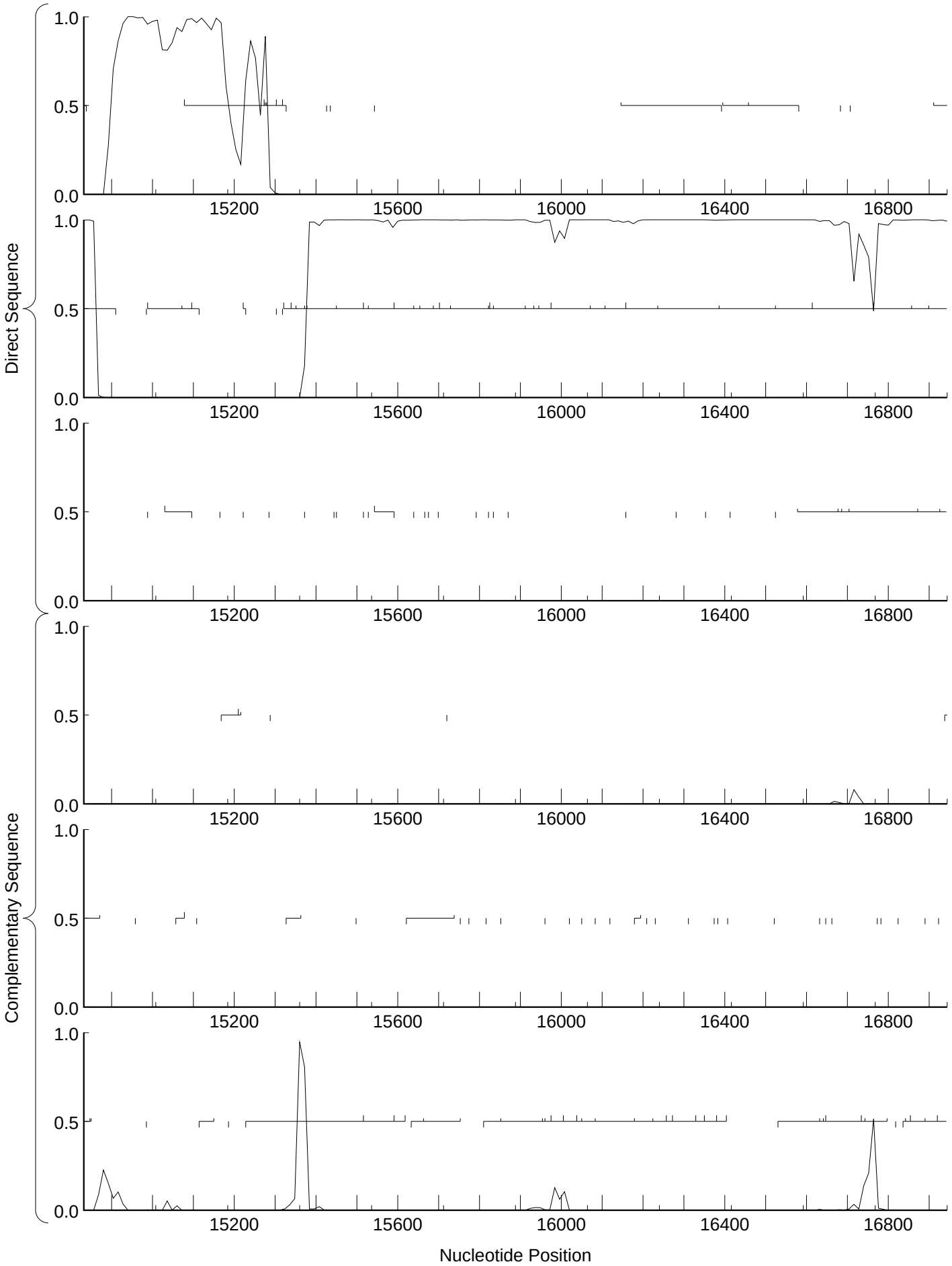
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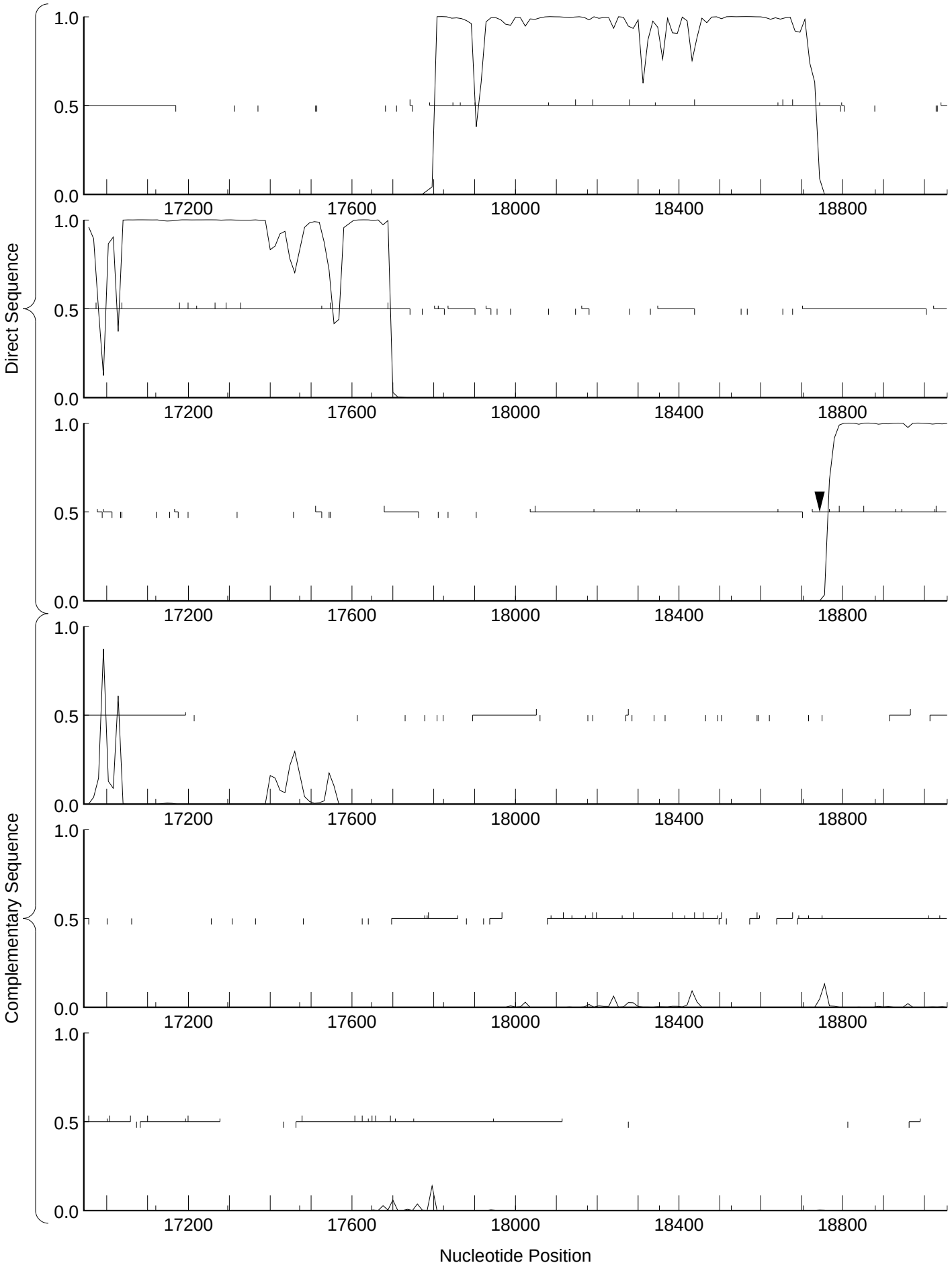
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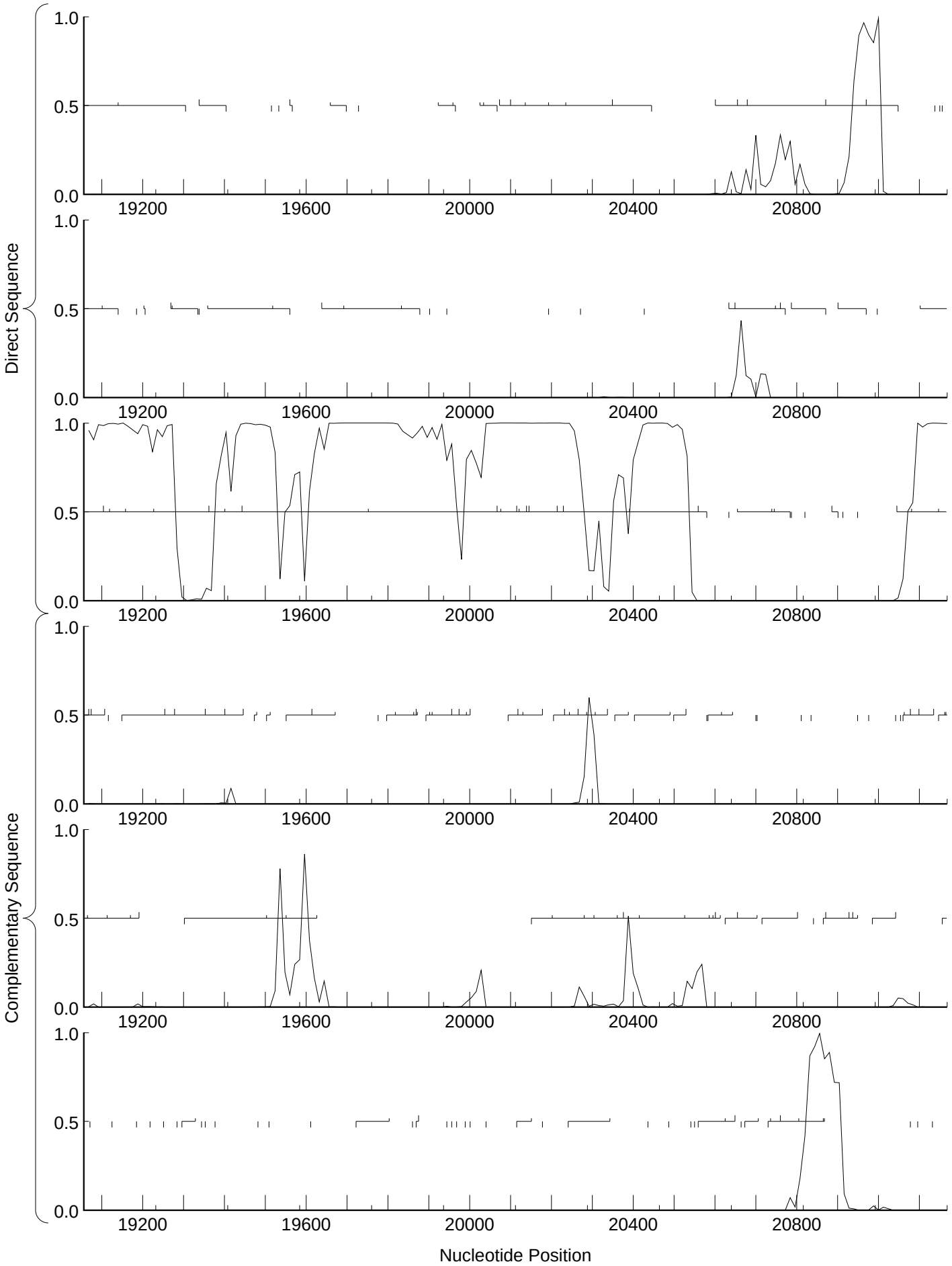
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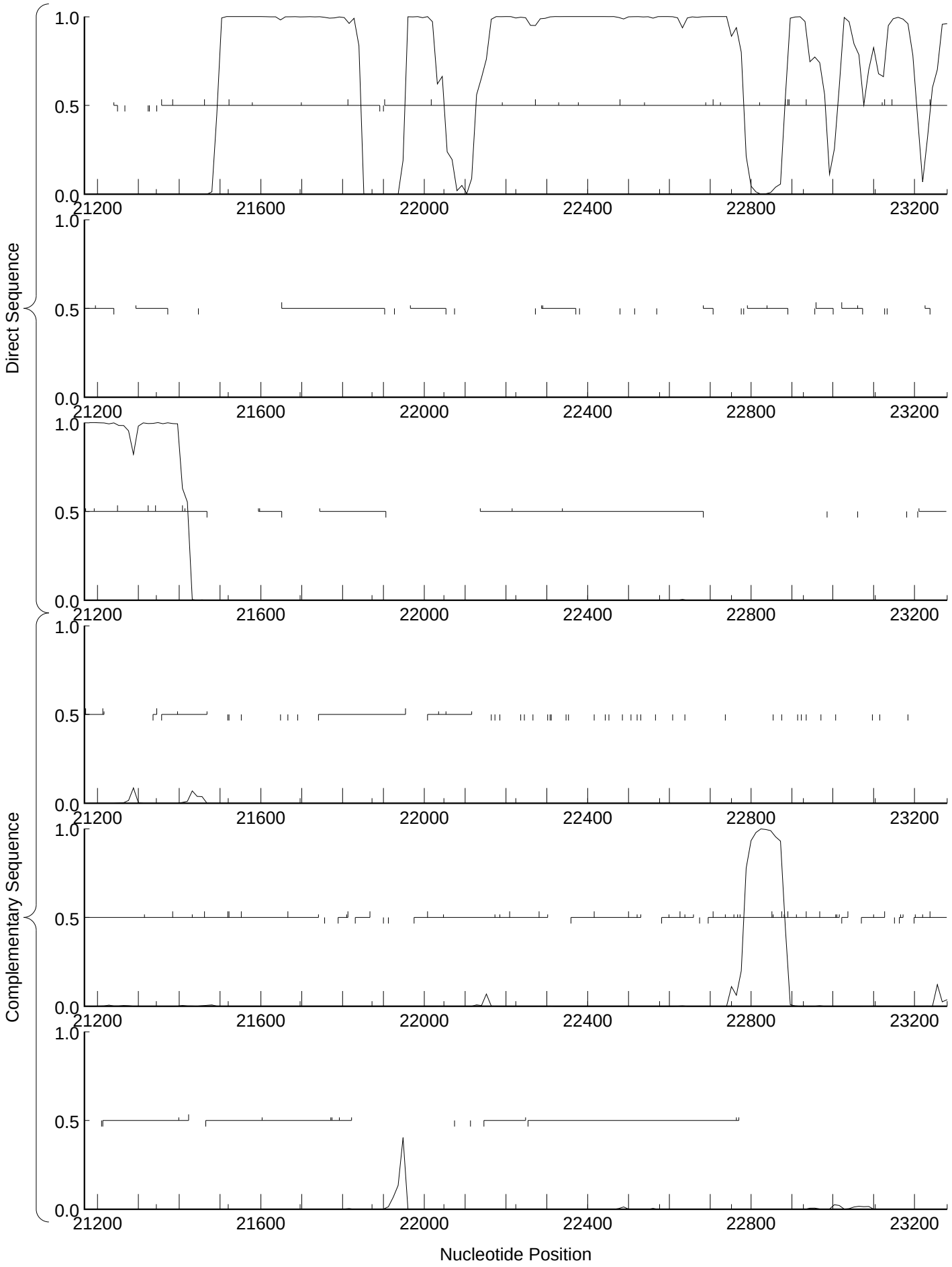
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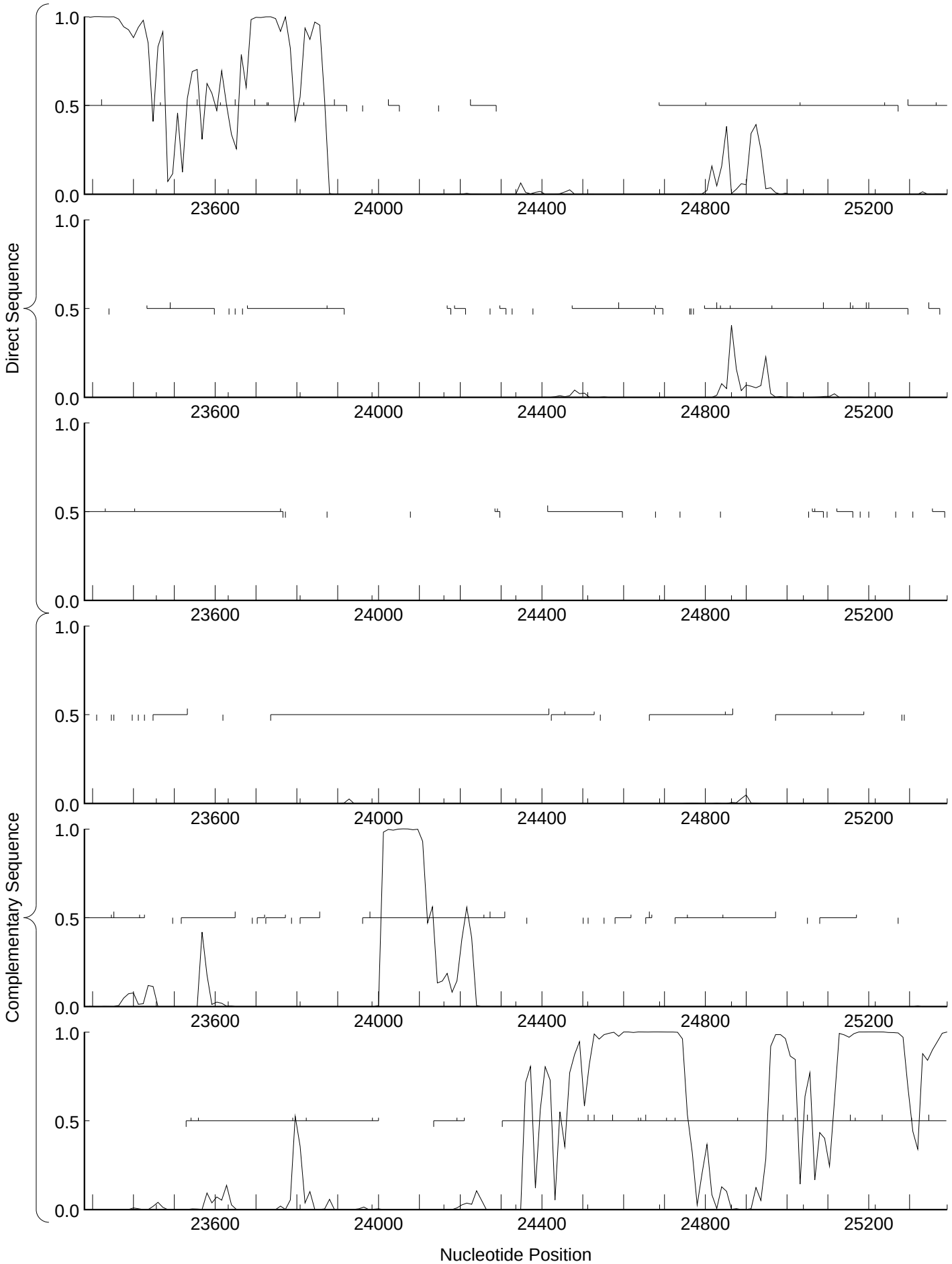
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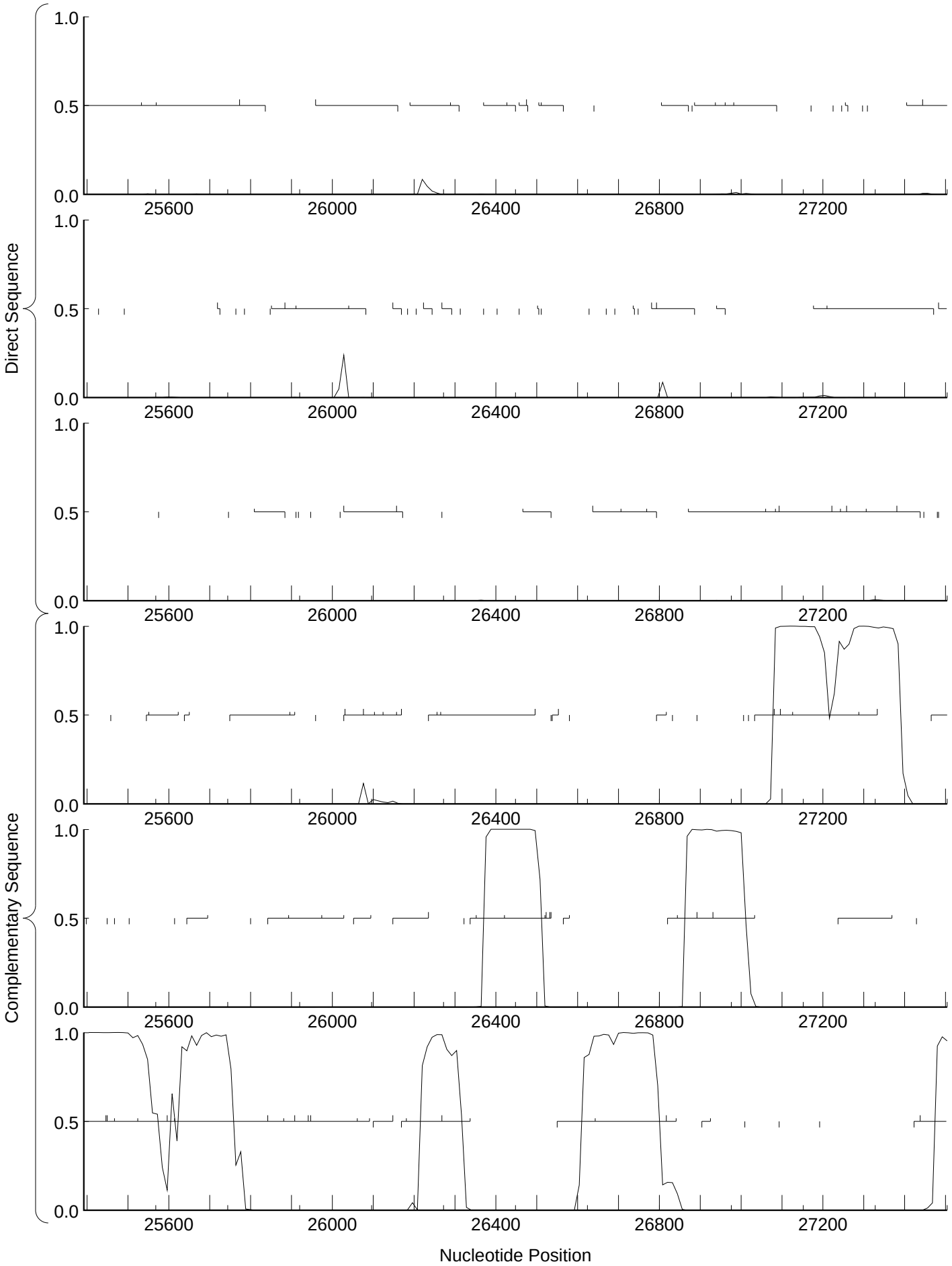


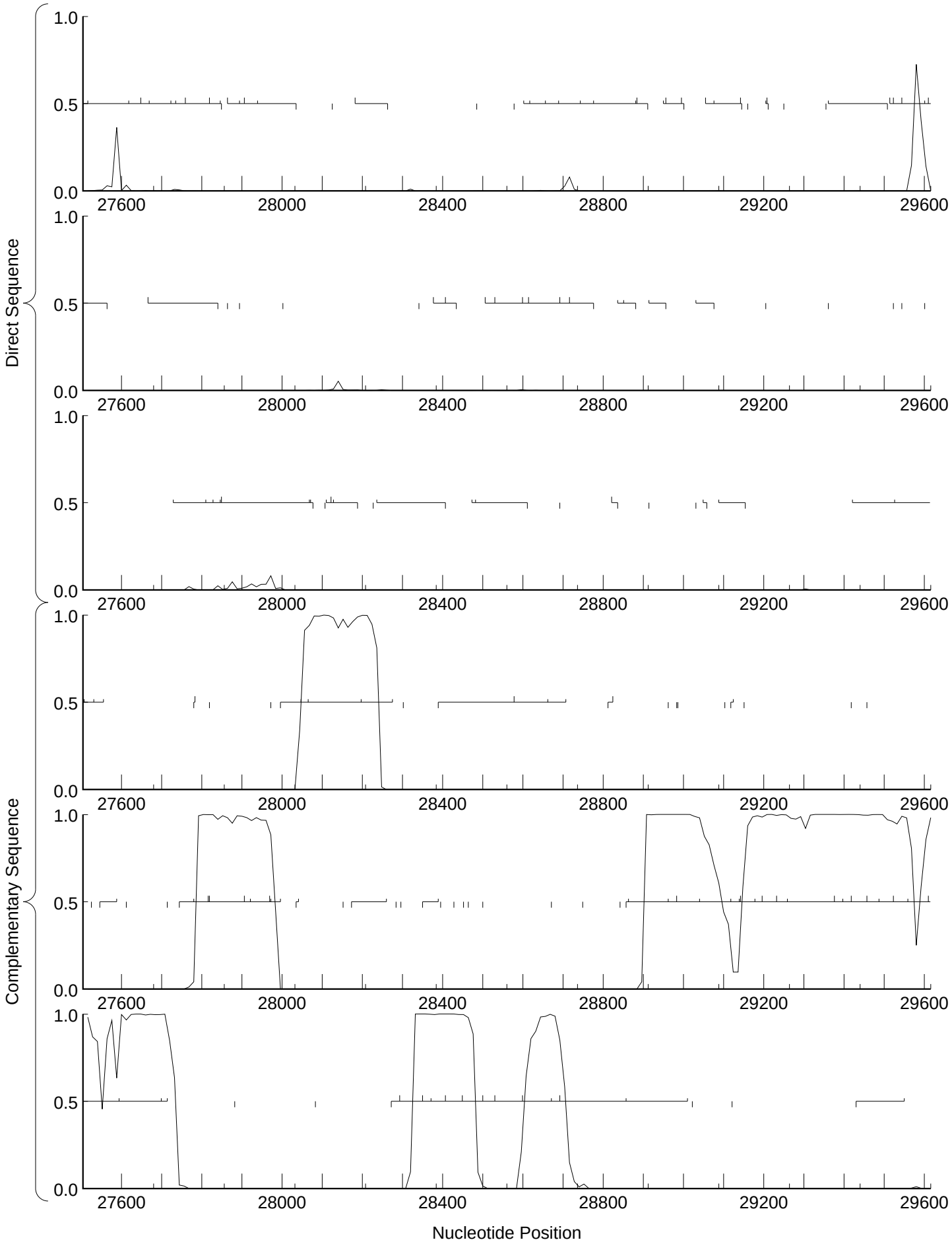
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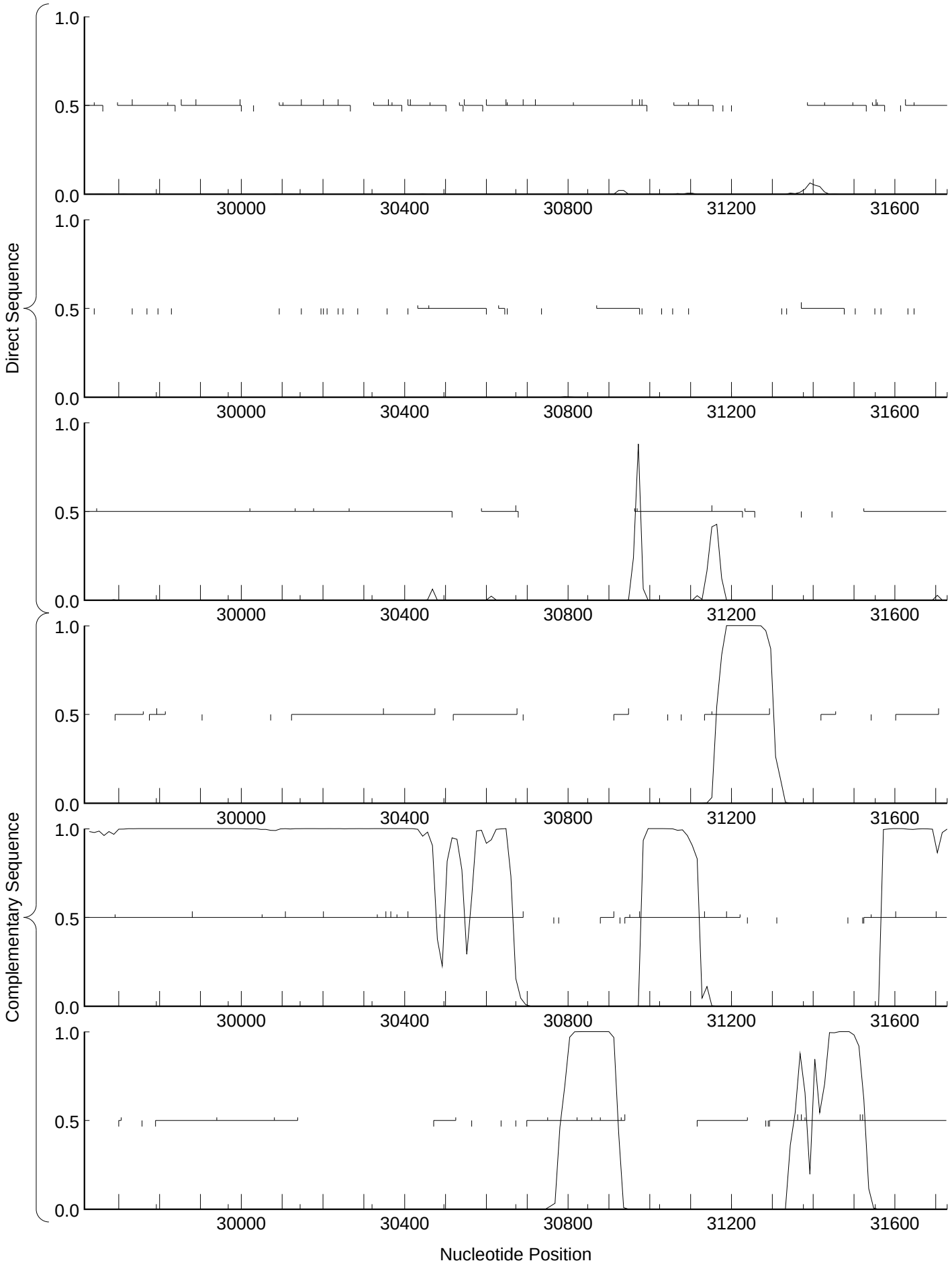


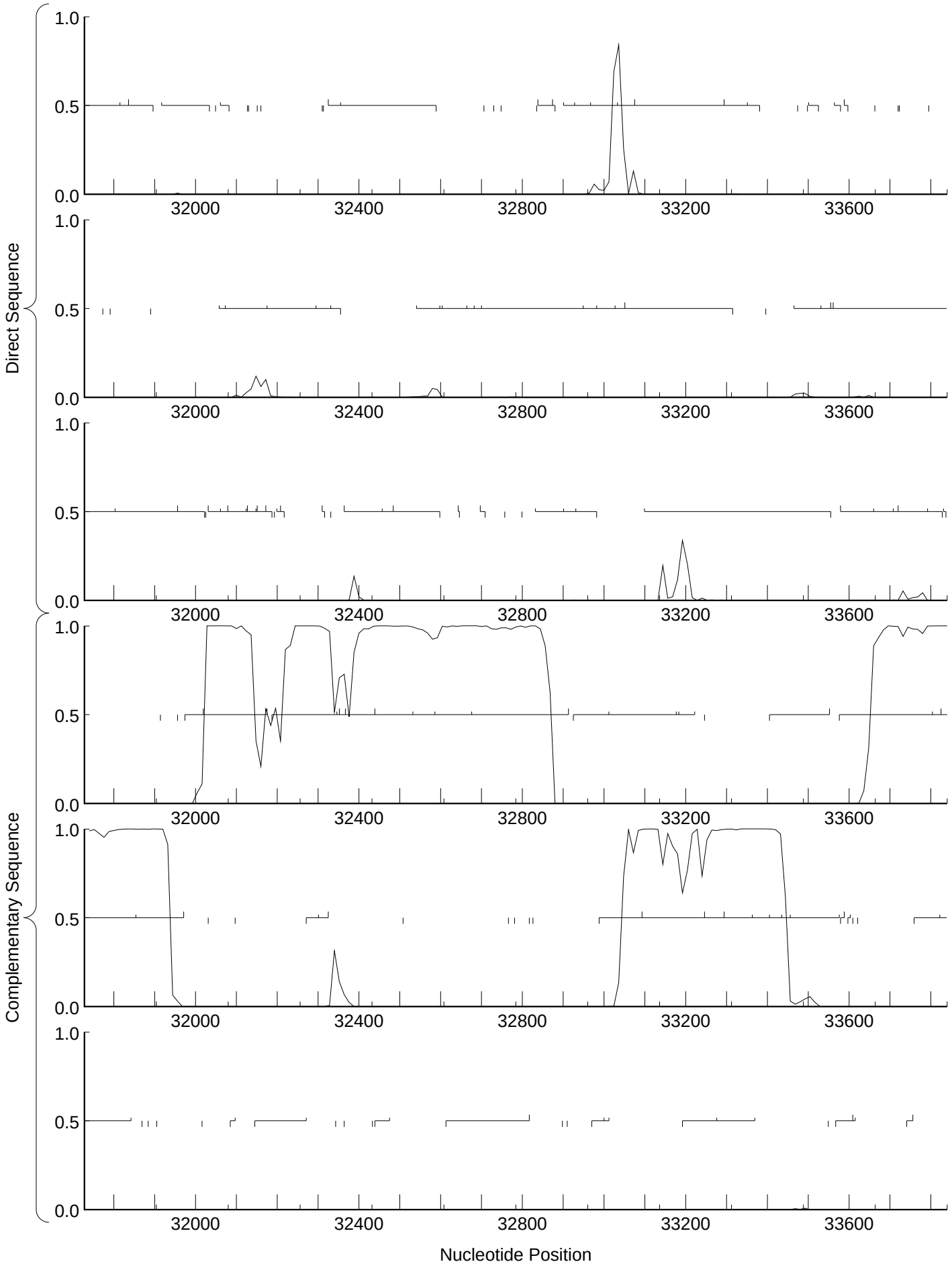
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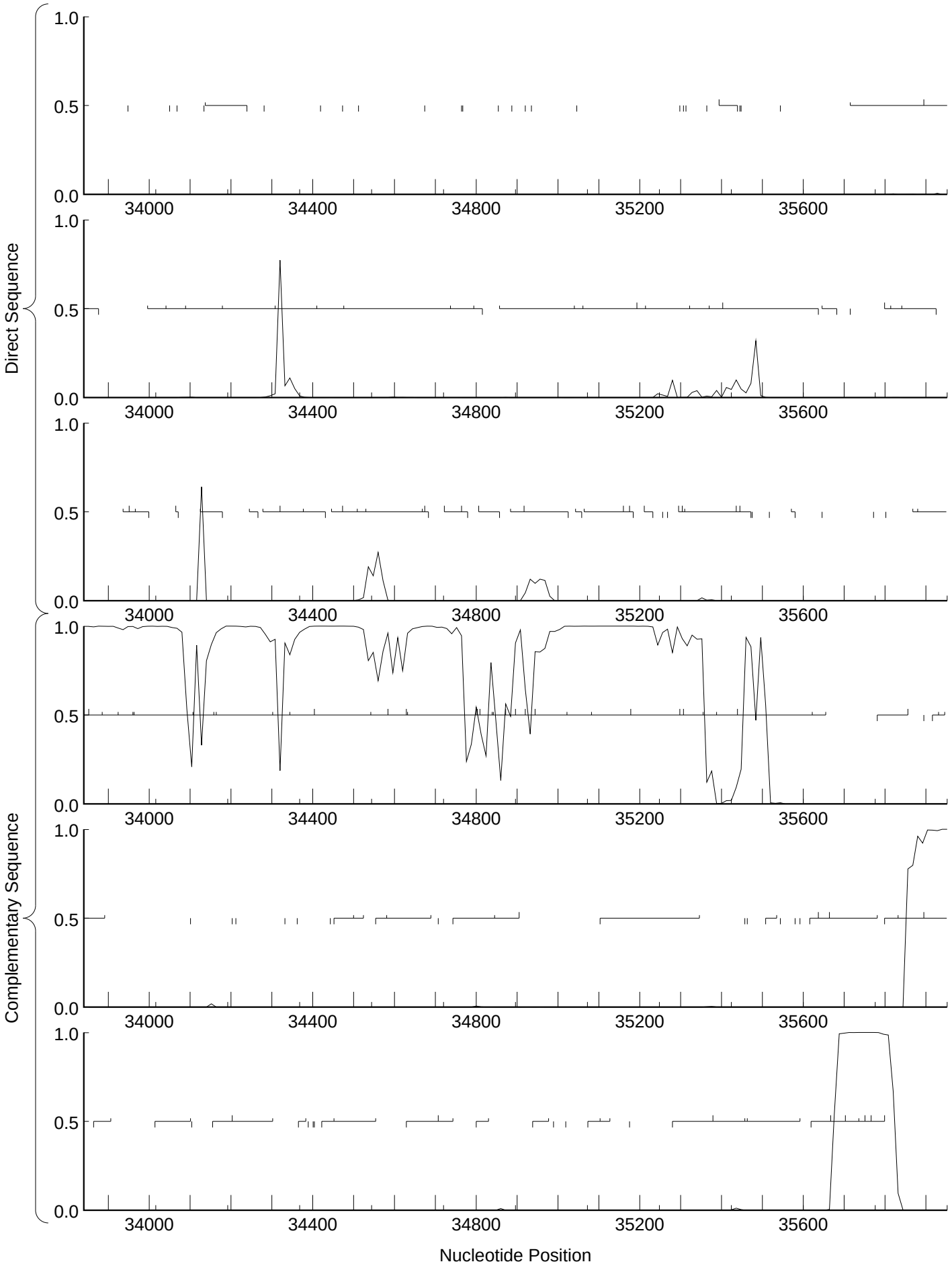


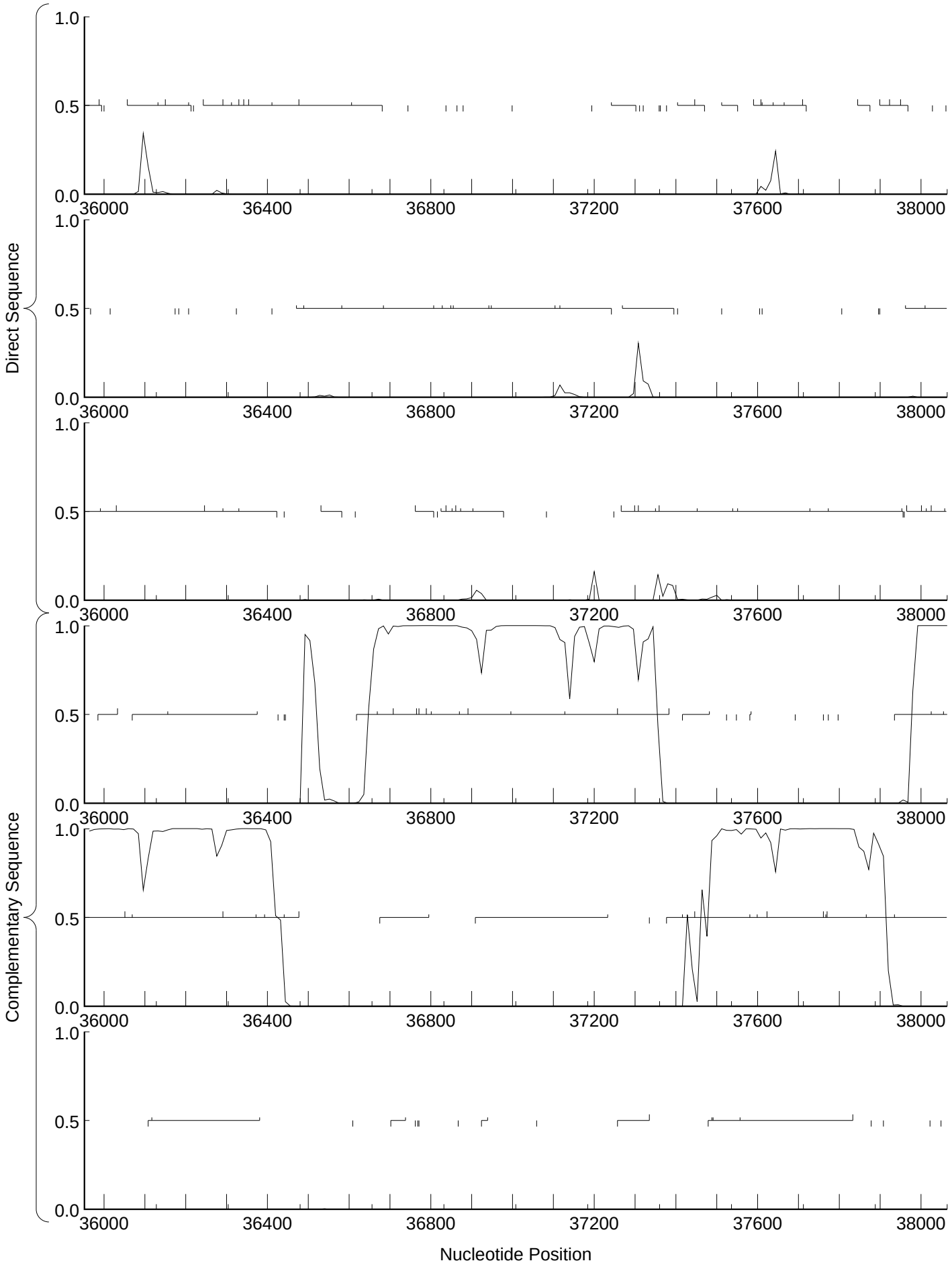

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Complementary Sequence
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26000
26400
26800
27200
Nucleotide Position

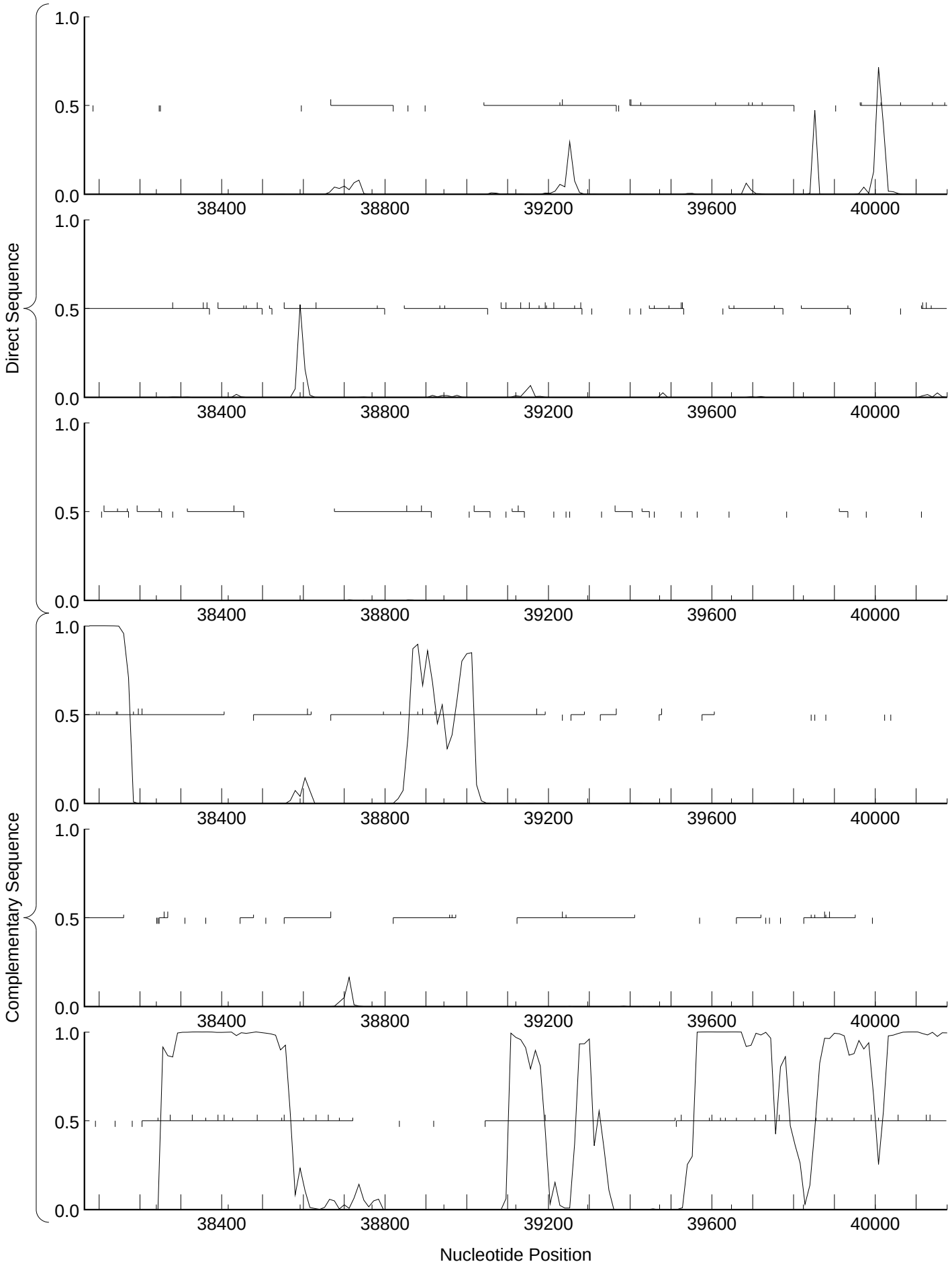

Direct Sequence
Complementary Sequence
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0.0
27600
28000
28400
28800
29200
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Nucleotide Position


Direct Sequence
Complementary Sequence
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Nucleotide Position

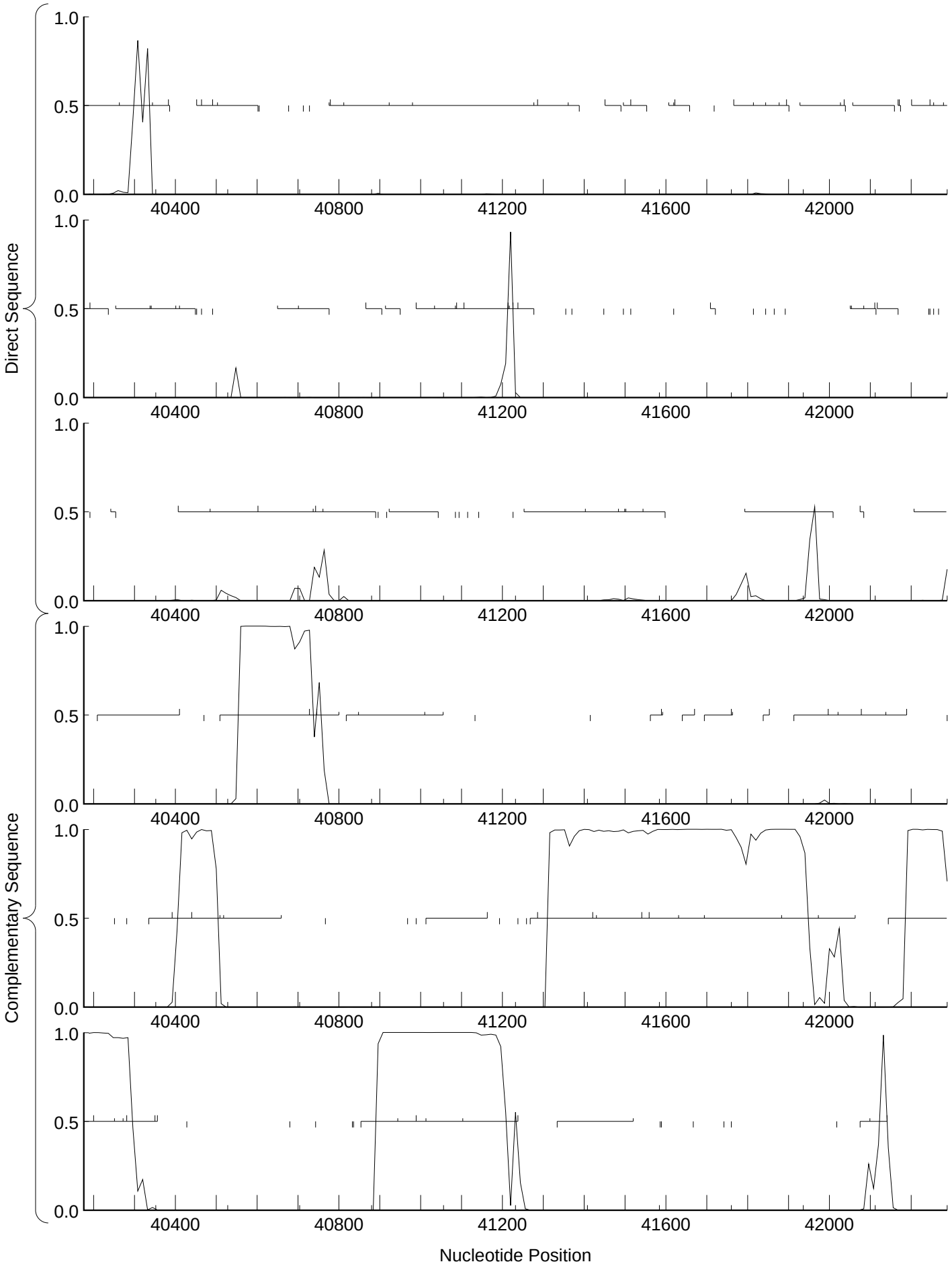

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33200
33600
Nucleotide Position

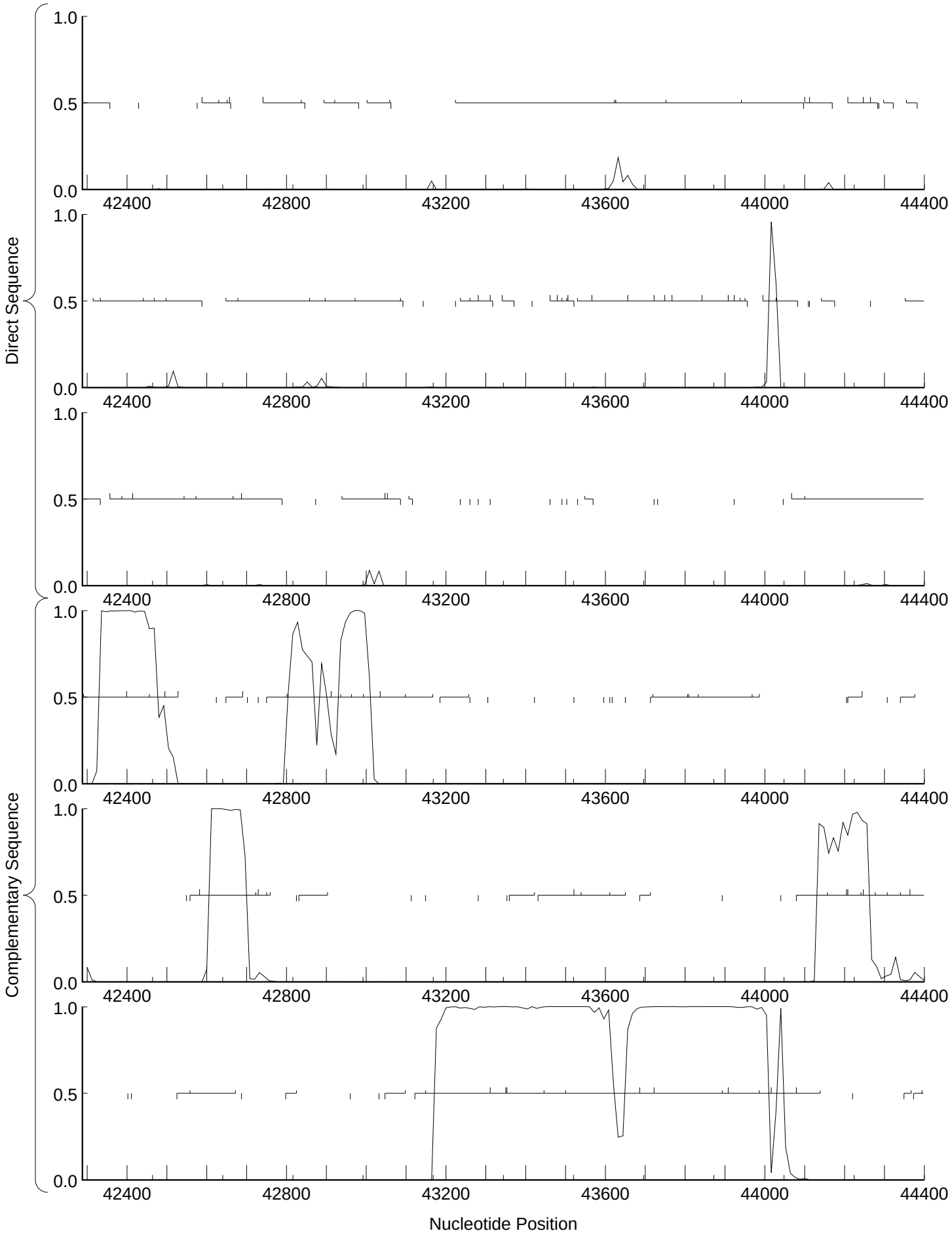

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Complementary Sequence
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35200
35600
Nucleotide Position

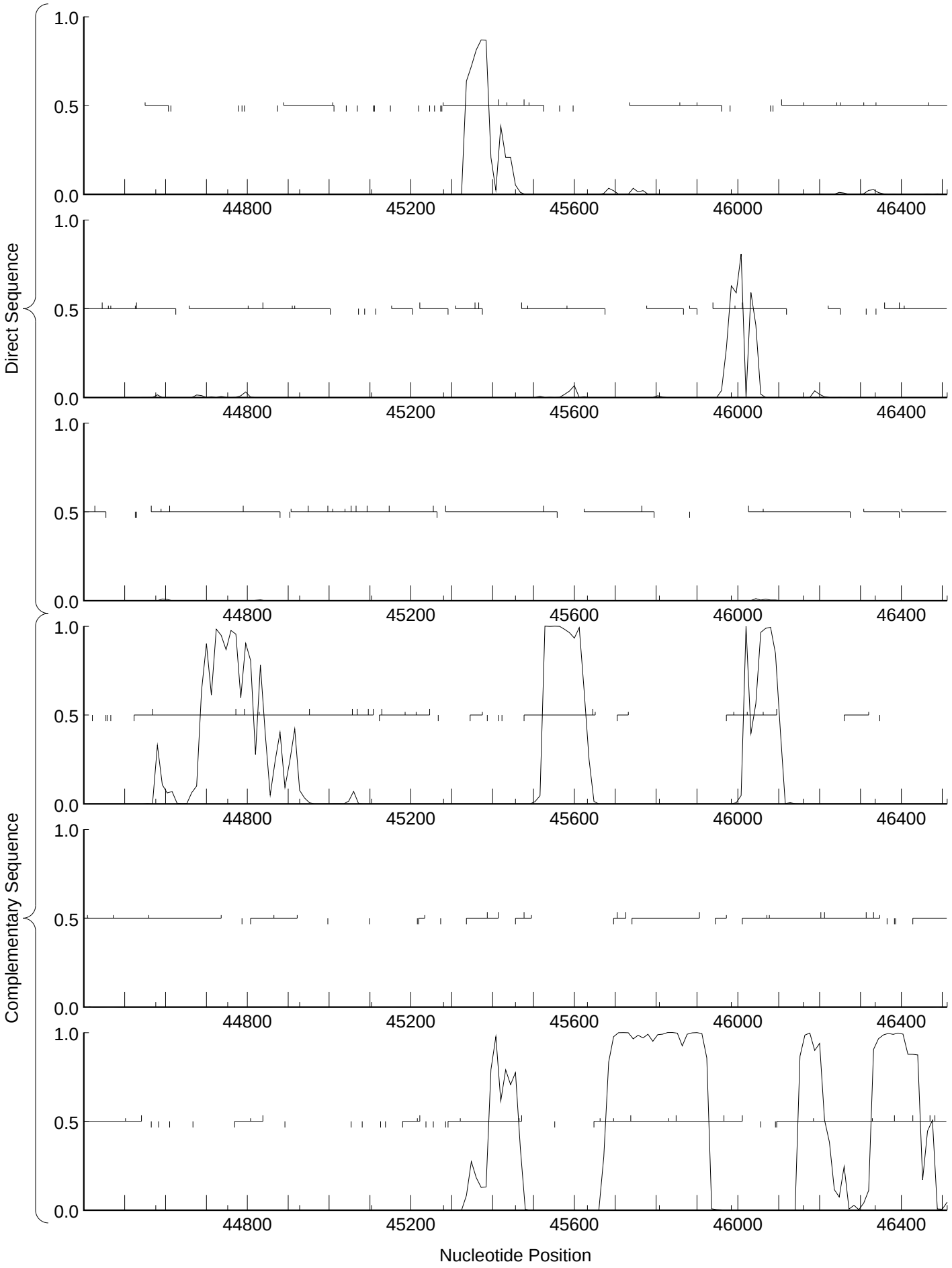

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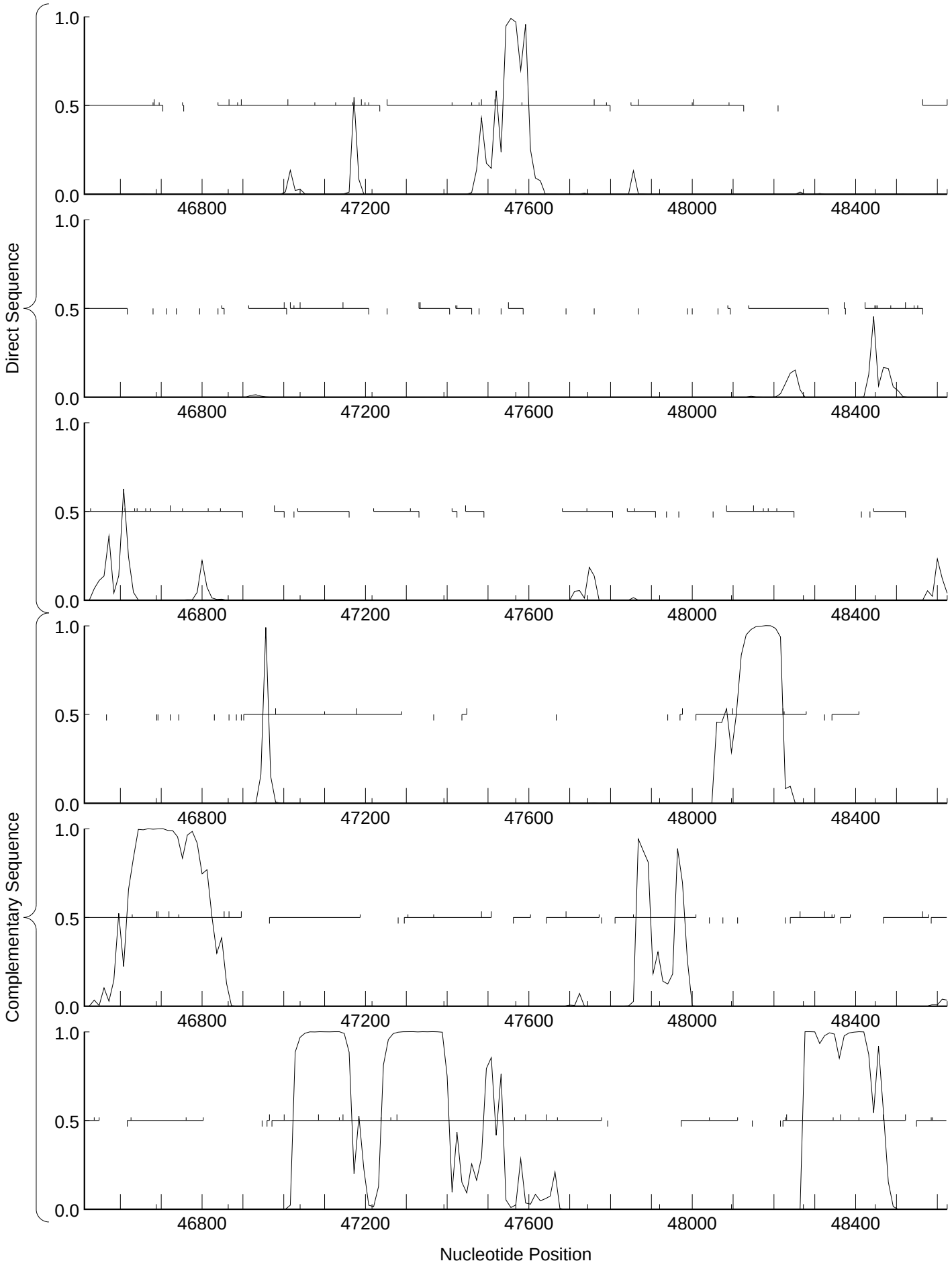

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Nucleotide Position

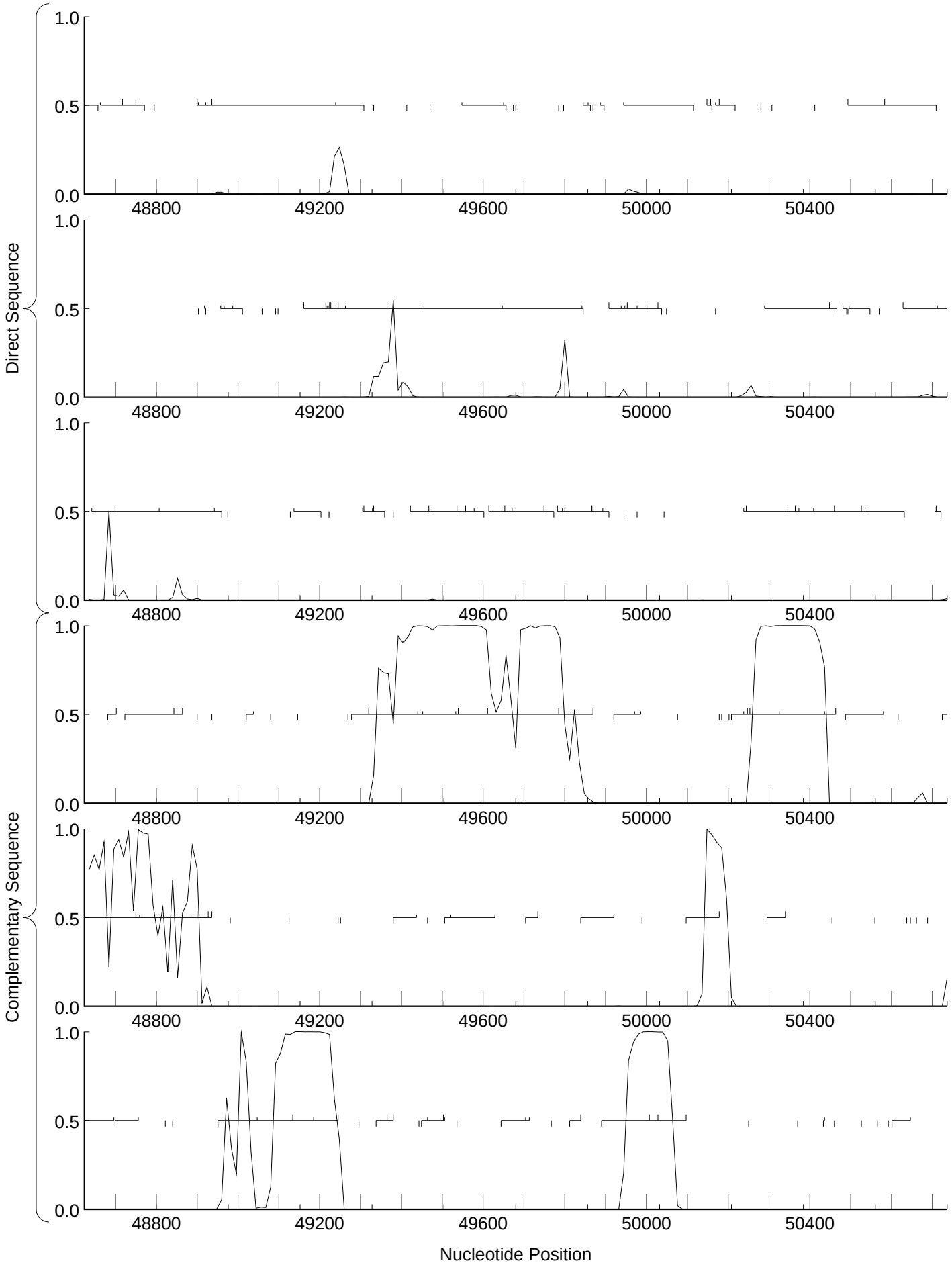
n complete sequence, 51546 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 21/26




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


Direct Sequence
Complementary Sequence
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0.0
44800
45200
45600
46000
46400
Nucleotide Position


Direct Sequence
Complementary Sequence
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46800
47200
47600
48000
48400
Nucleotide Position


Direct Sequence
Complementary Sequence
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0.0
48800
49200
49600
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Nucleotide Position

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