GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Kalpine complete sequence, 53330 bp including 10-base 3’ overhang (CGGT
Analysis Date    :  3/19/24 at 2:46:01
Pages            :  27
Sequence Length  :  53330 bp
GC Content       :   64.37%

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

PS-Version       :  1.2

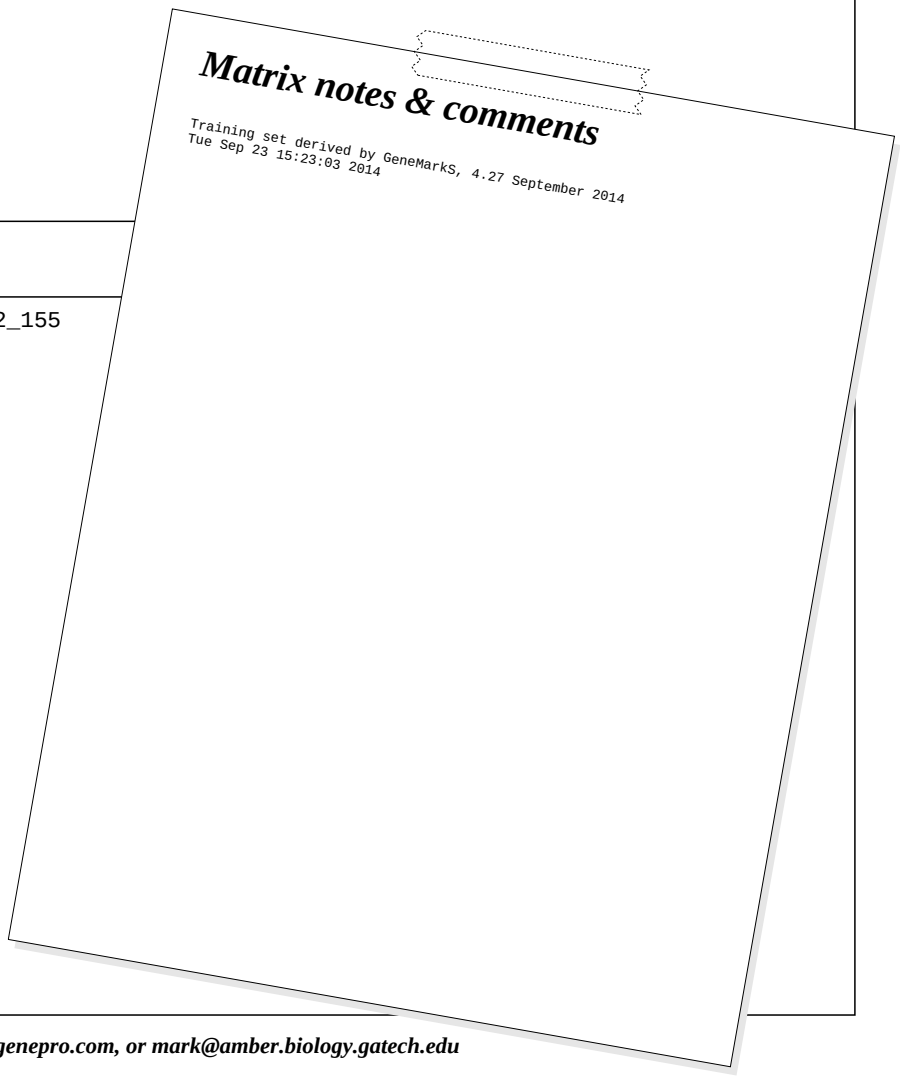
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
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MATRIX INFORMATION

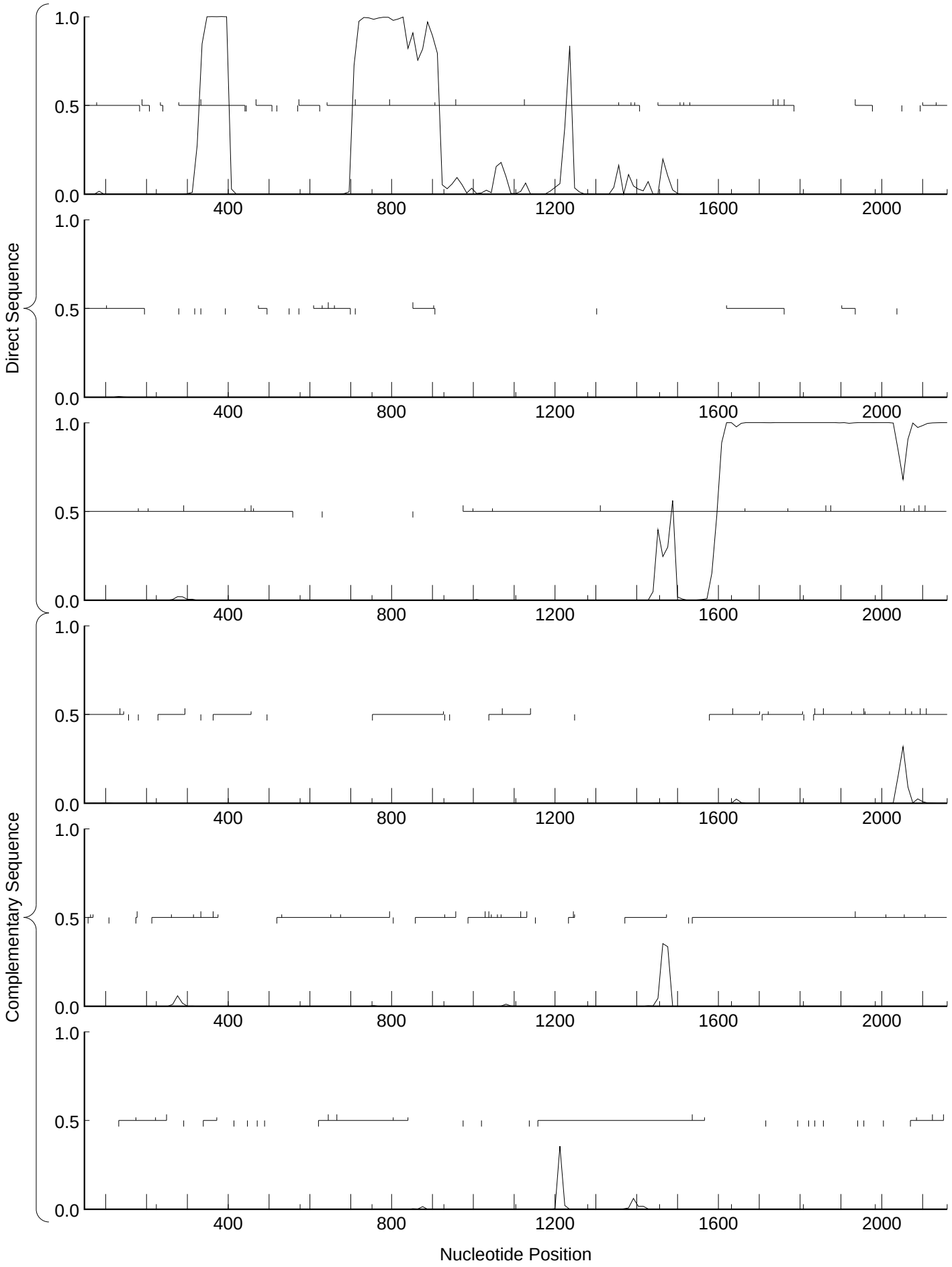
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Author     :  -

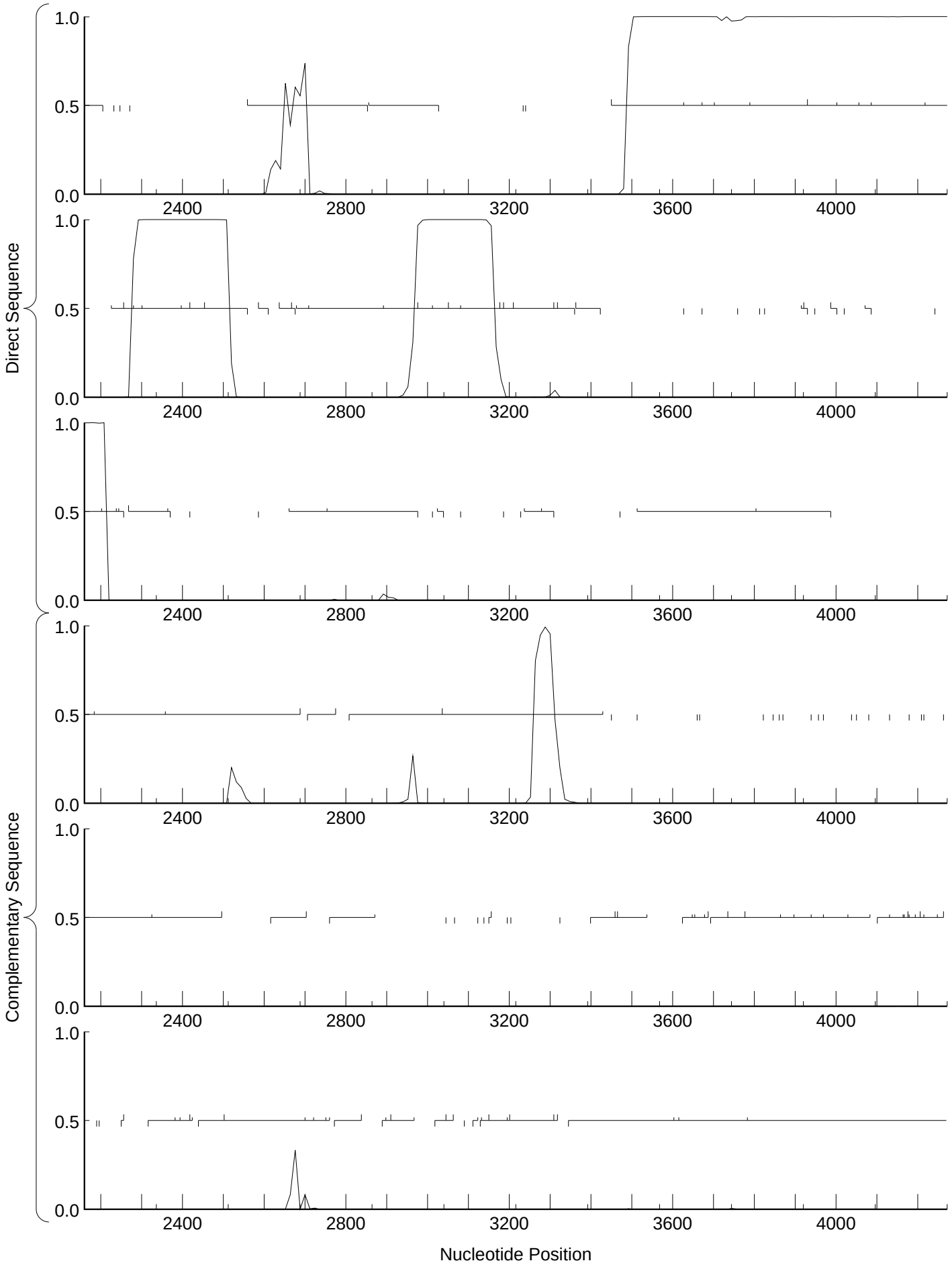
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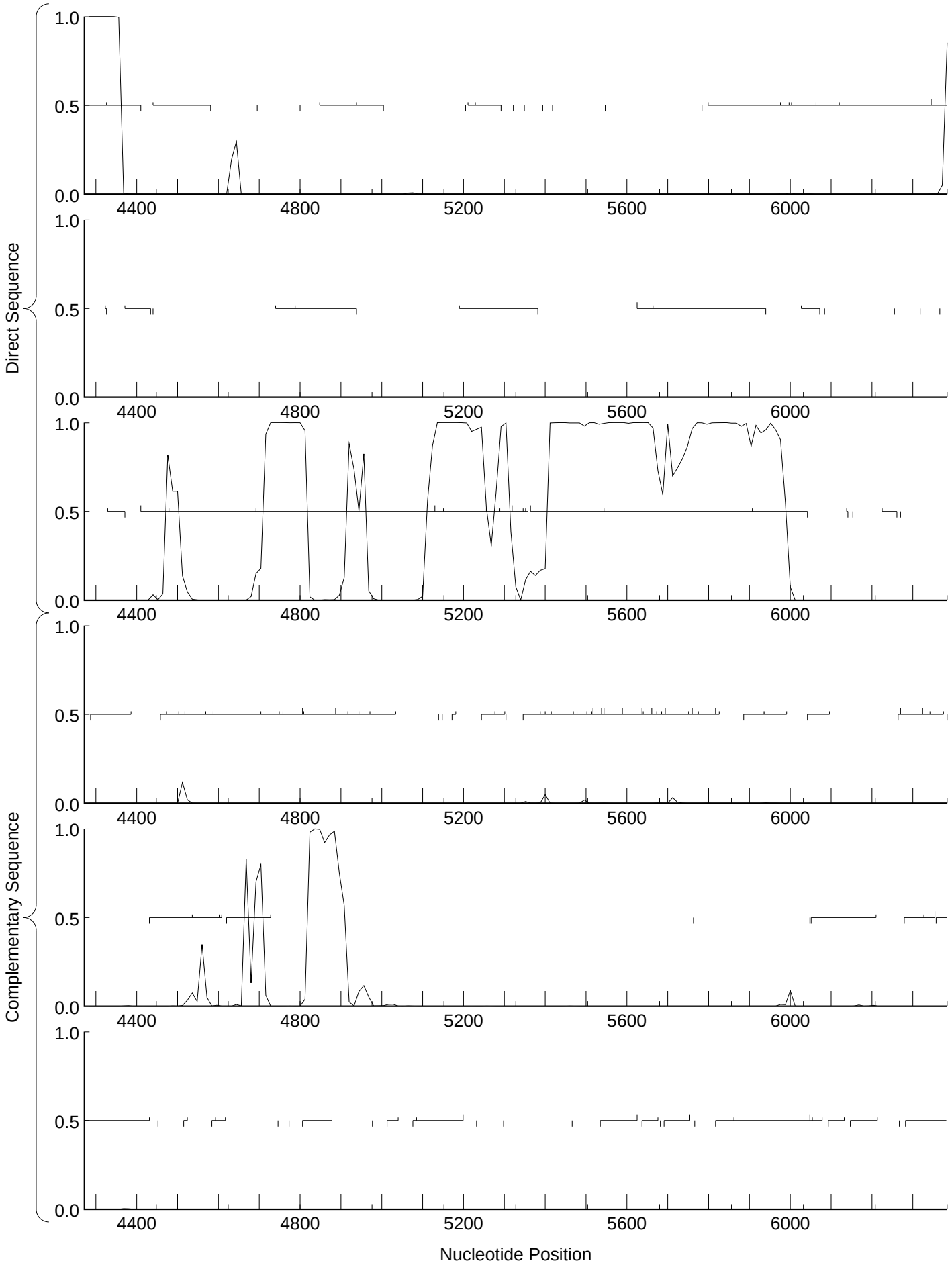

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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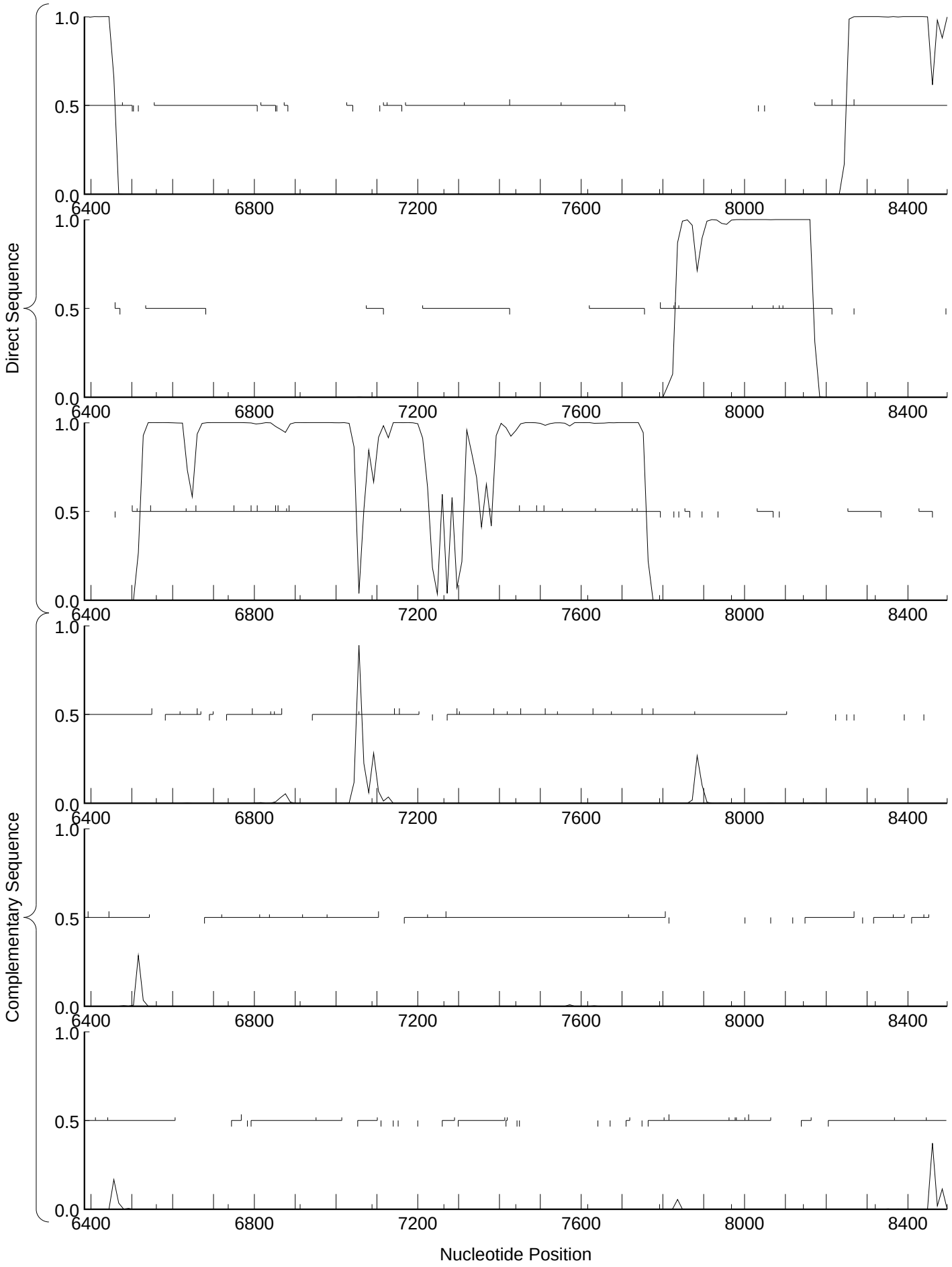


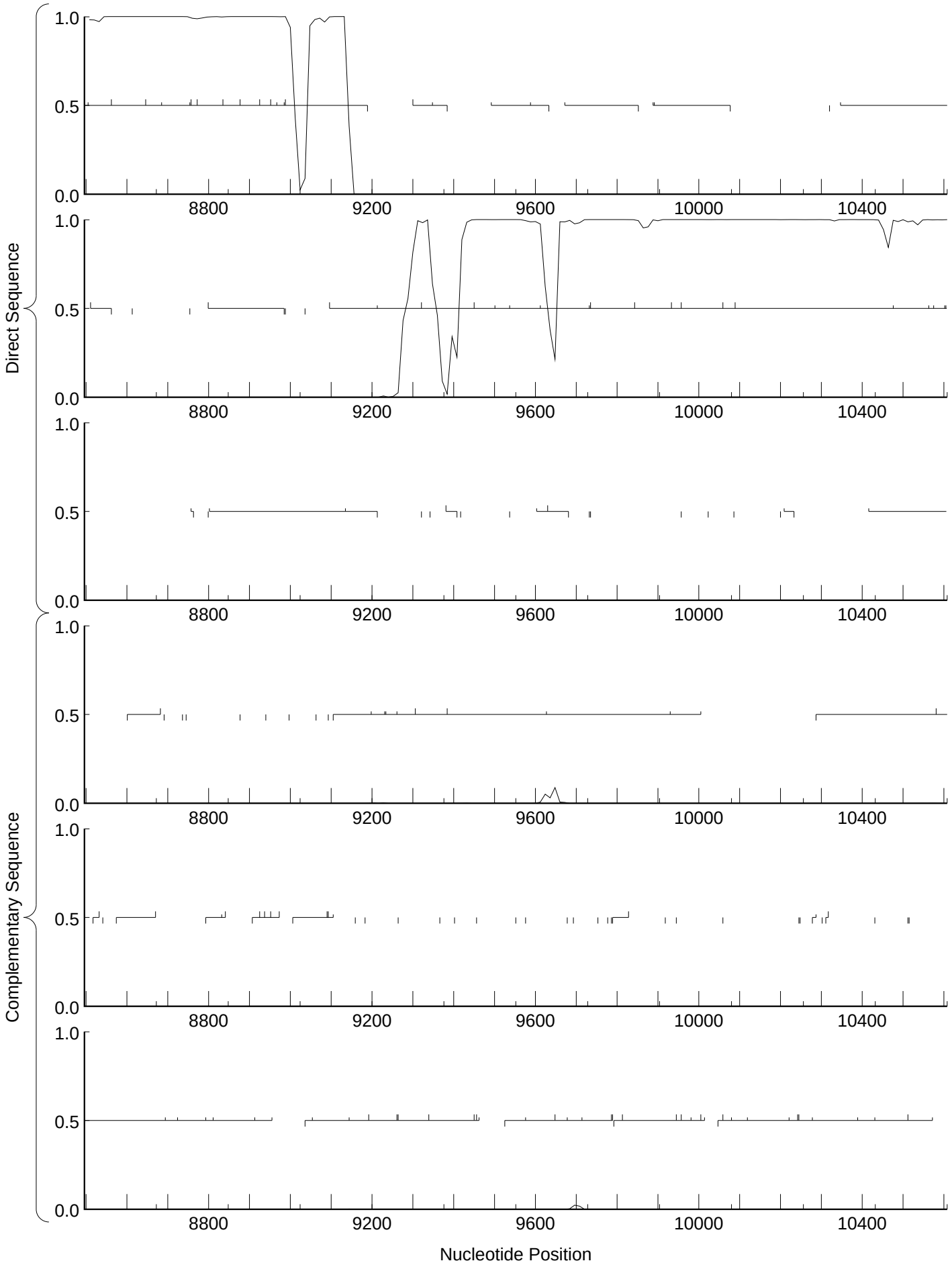
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

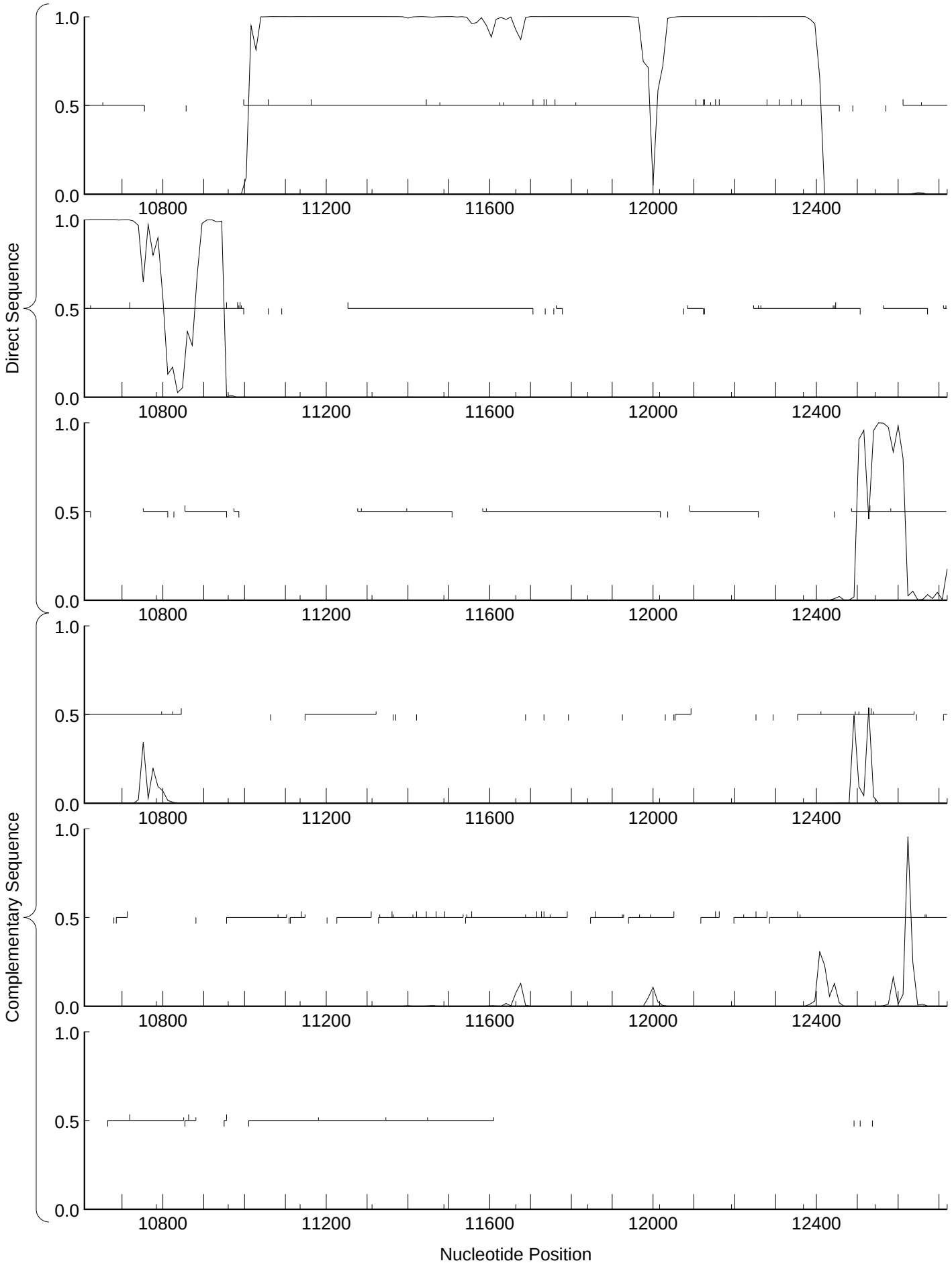

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800
1200
1600
2000
Nucleotide Position

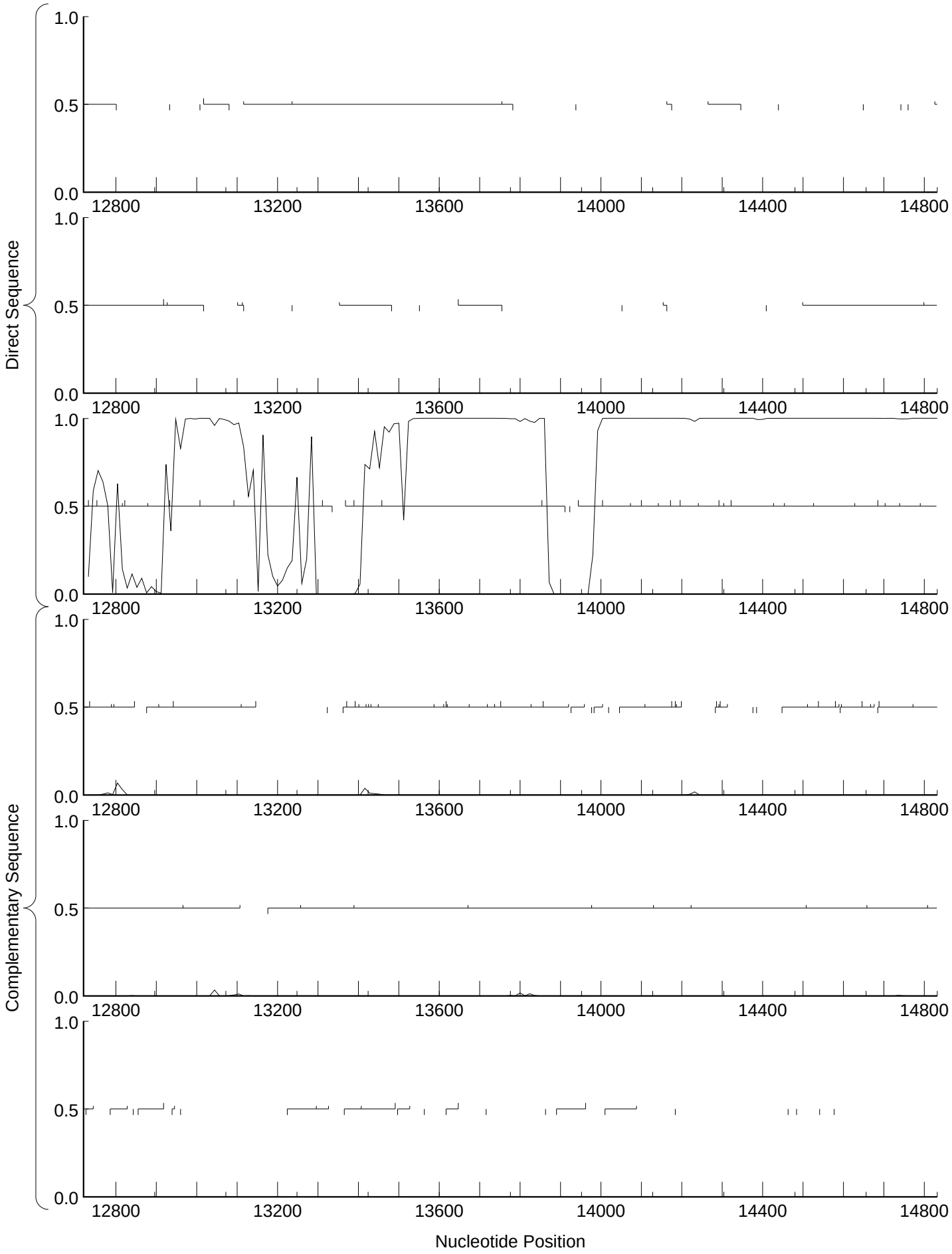

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0.0
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3200
3600
4000
Nucleotide Position

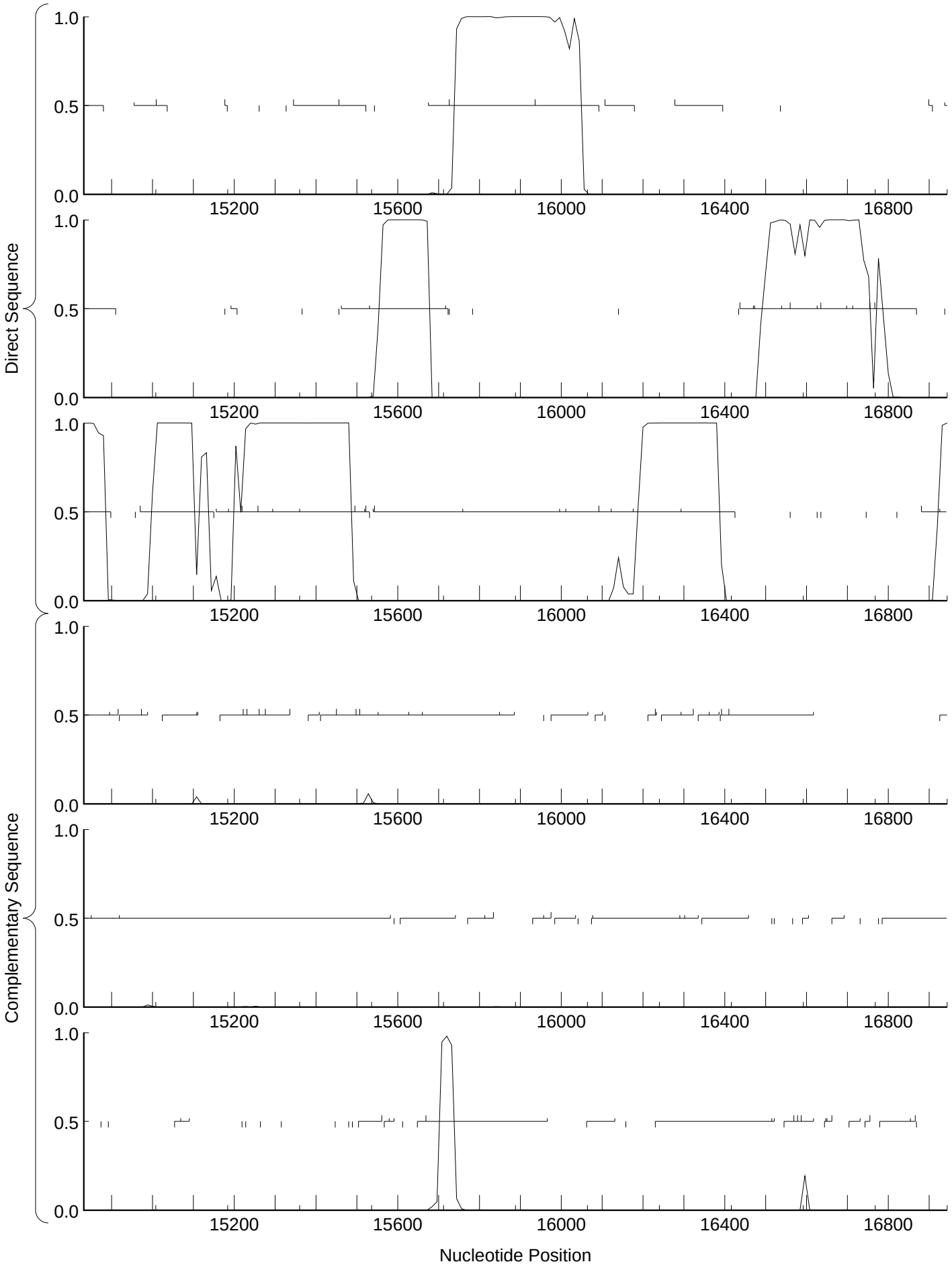

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4800
5200
5600
6000
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7600
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Nucleotide Position

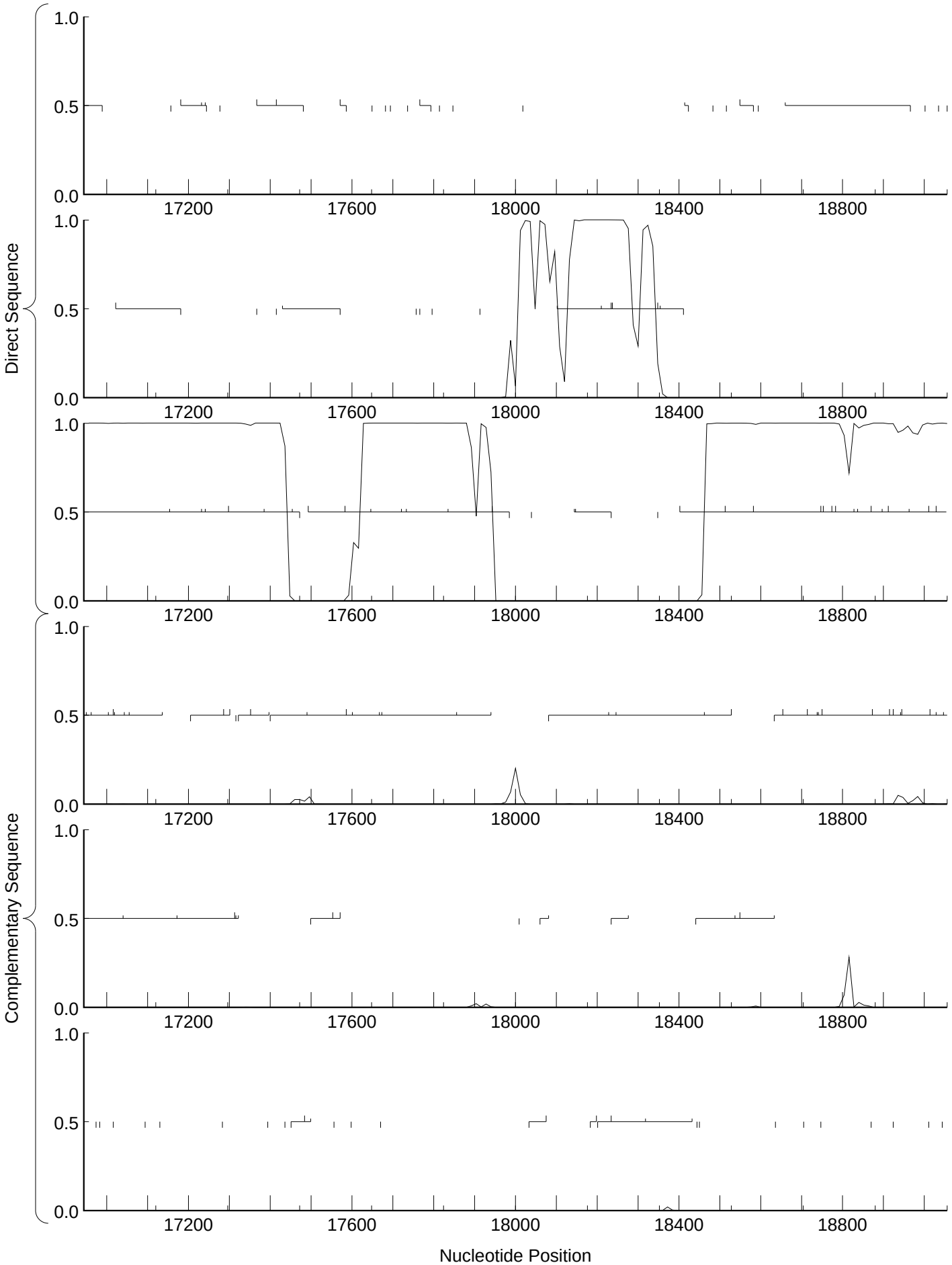

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8800
9200
9600
10000
10400
Nucleotide Position


Direct Sequence
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11600
12000
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Nucleotide Position

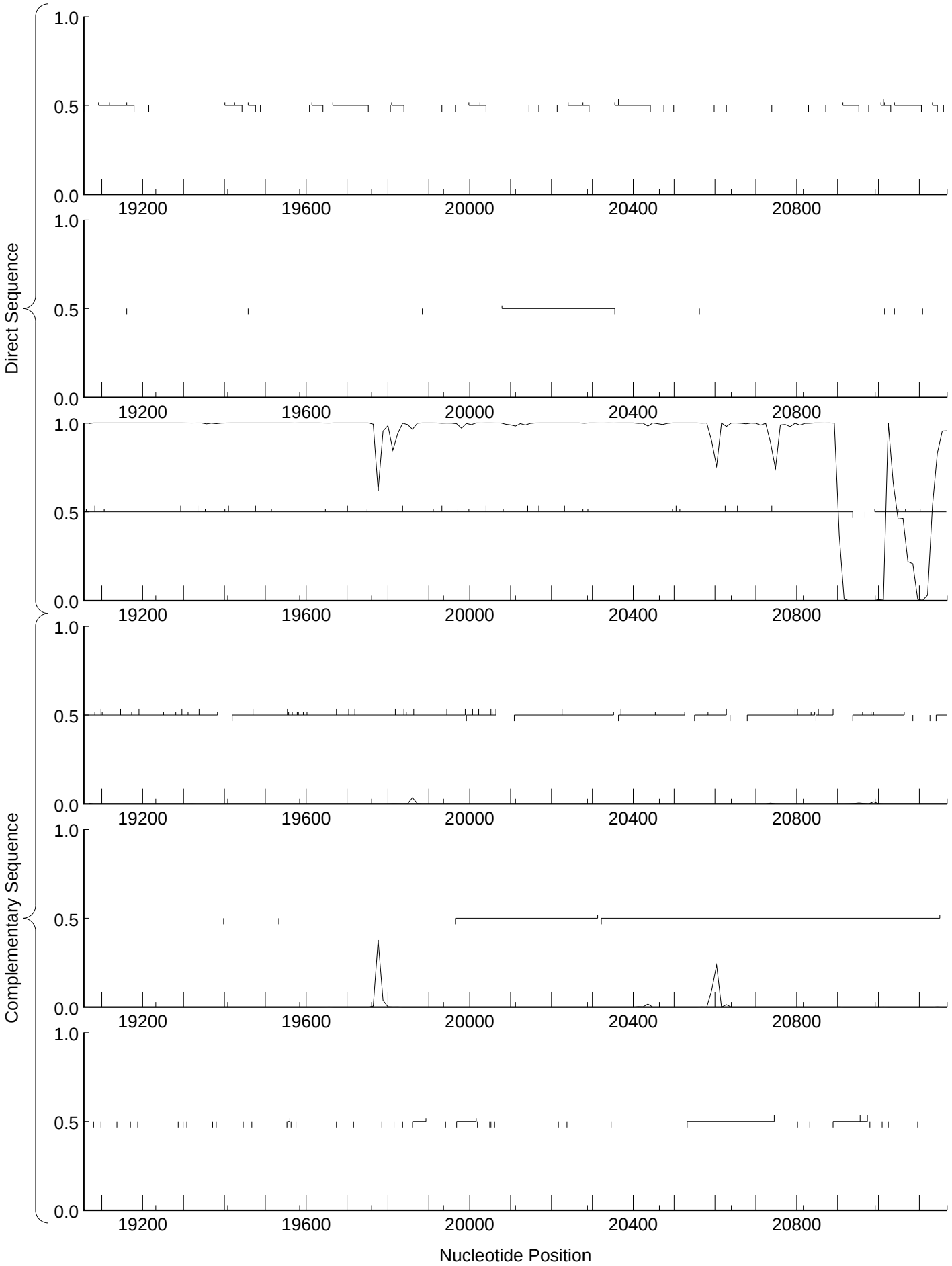

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13200
13600
14000
14400
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Nucleotide Position


Direct Sequence
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16000
16400
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Nucleotide Position

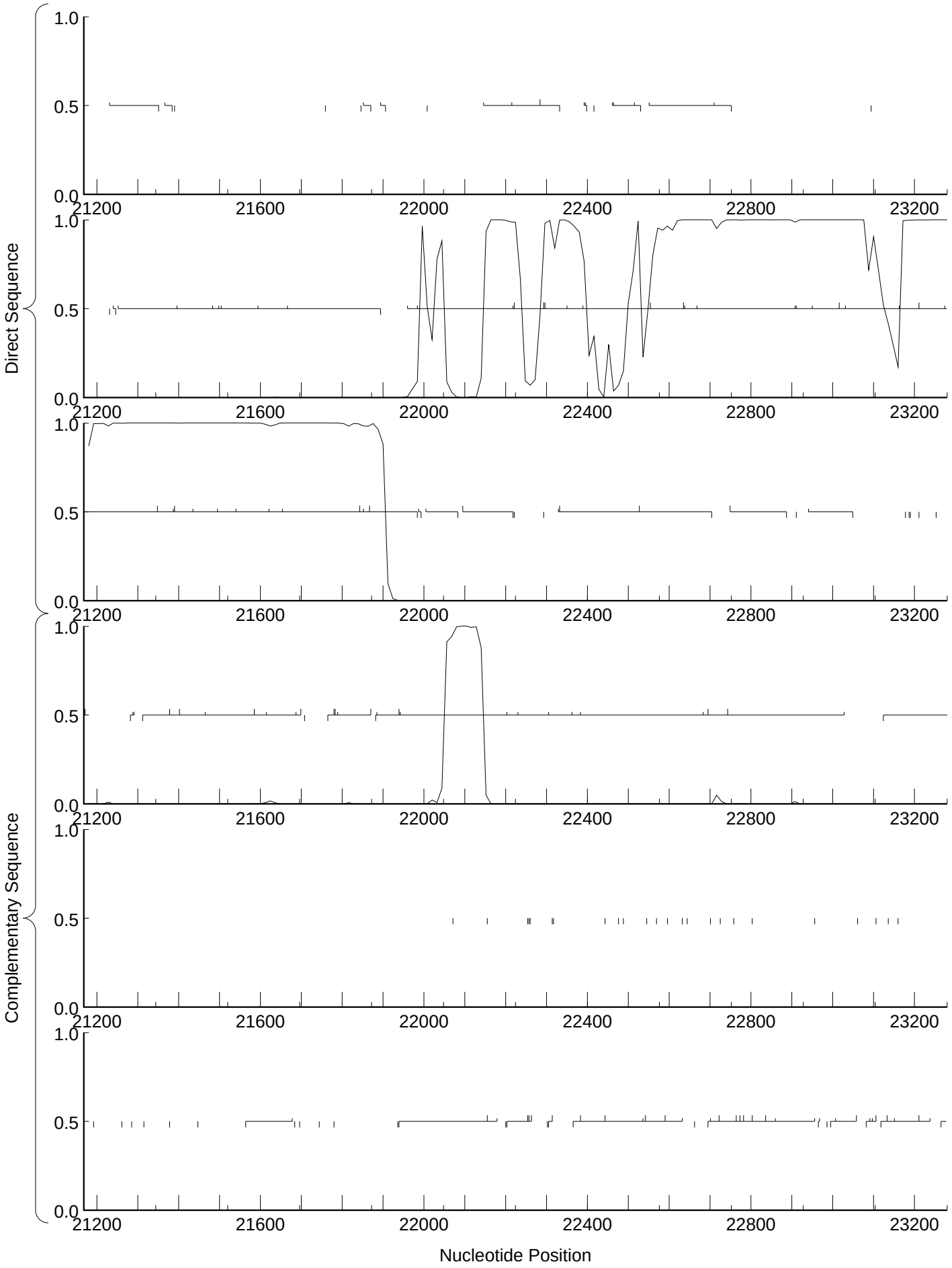
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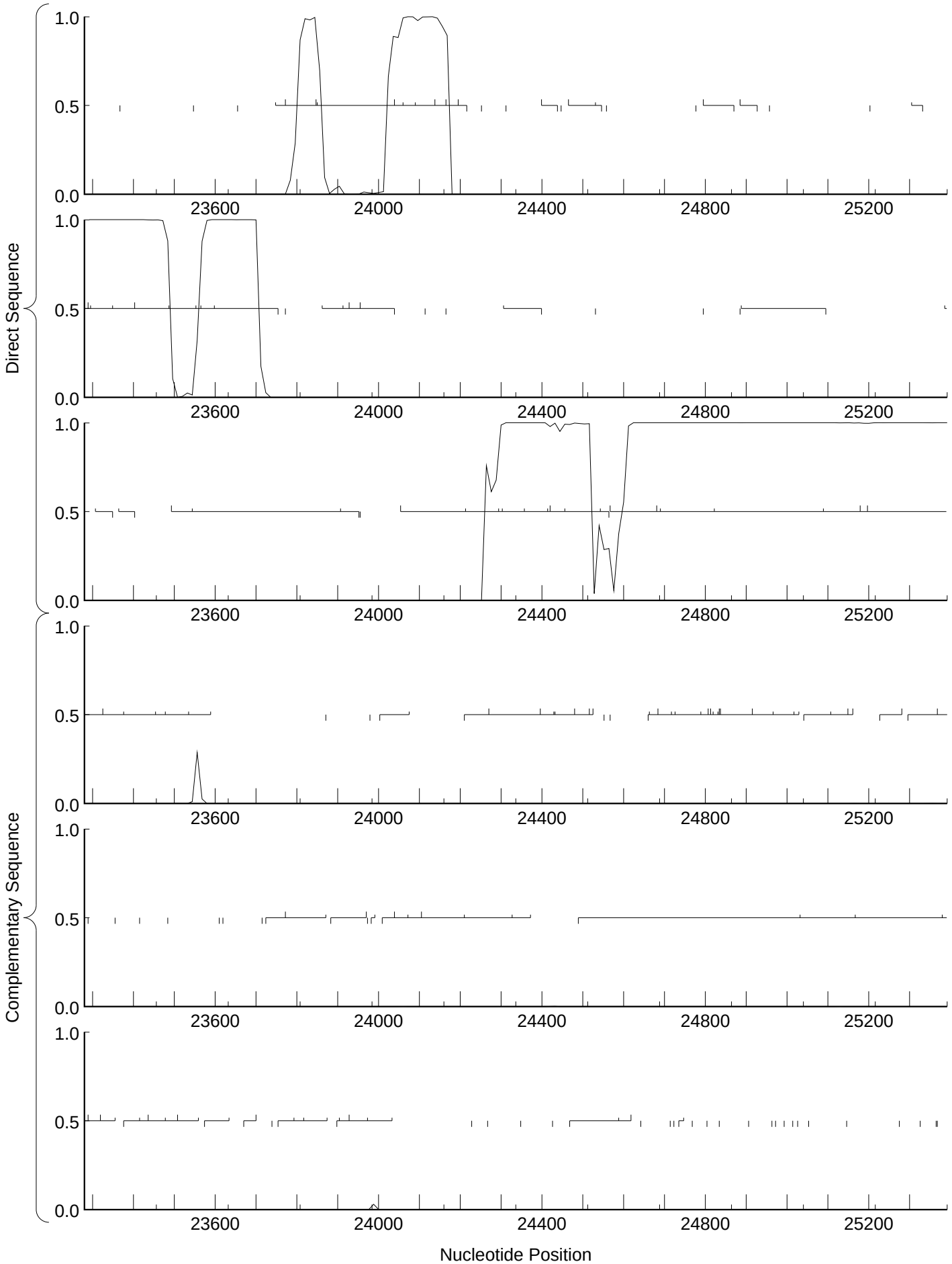
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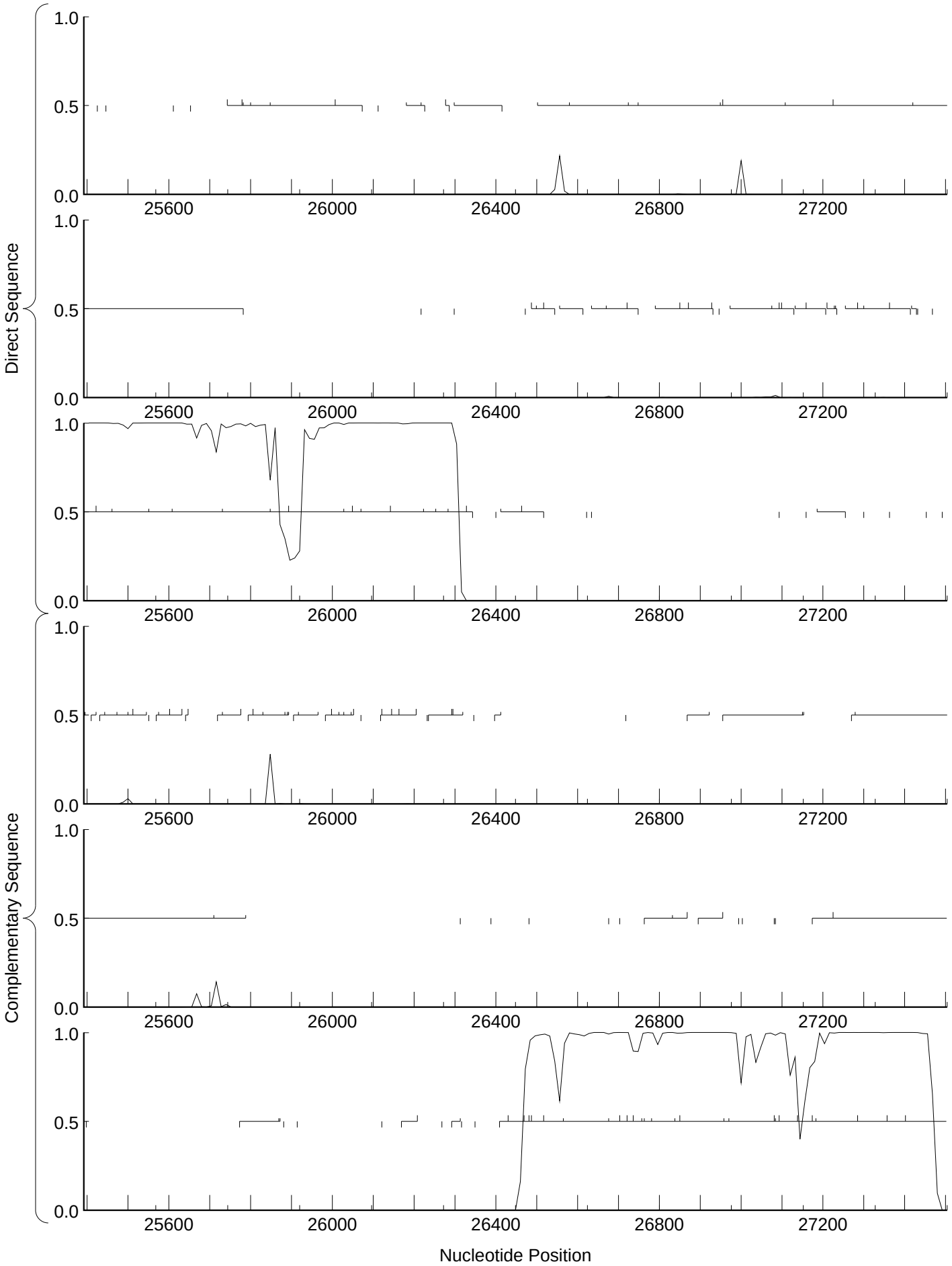
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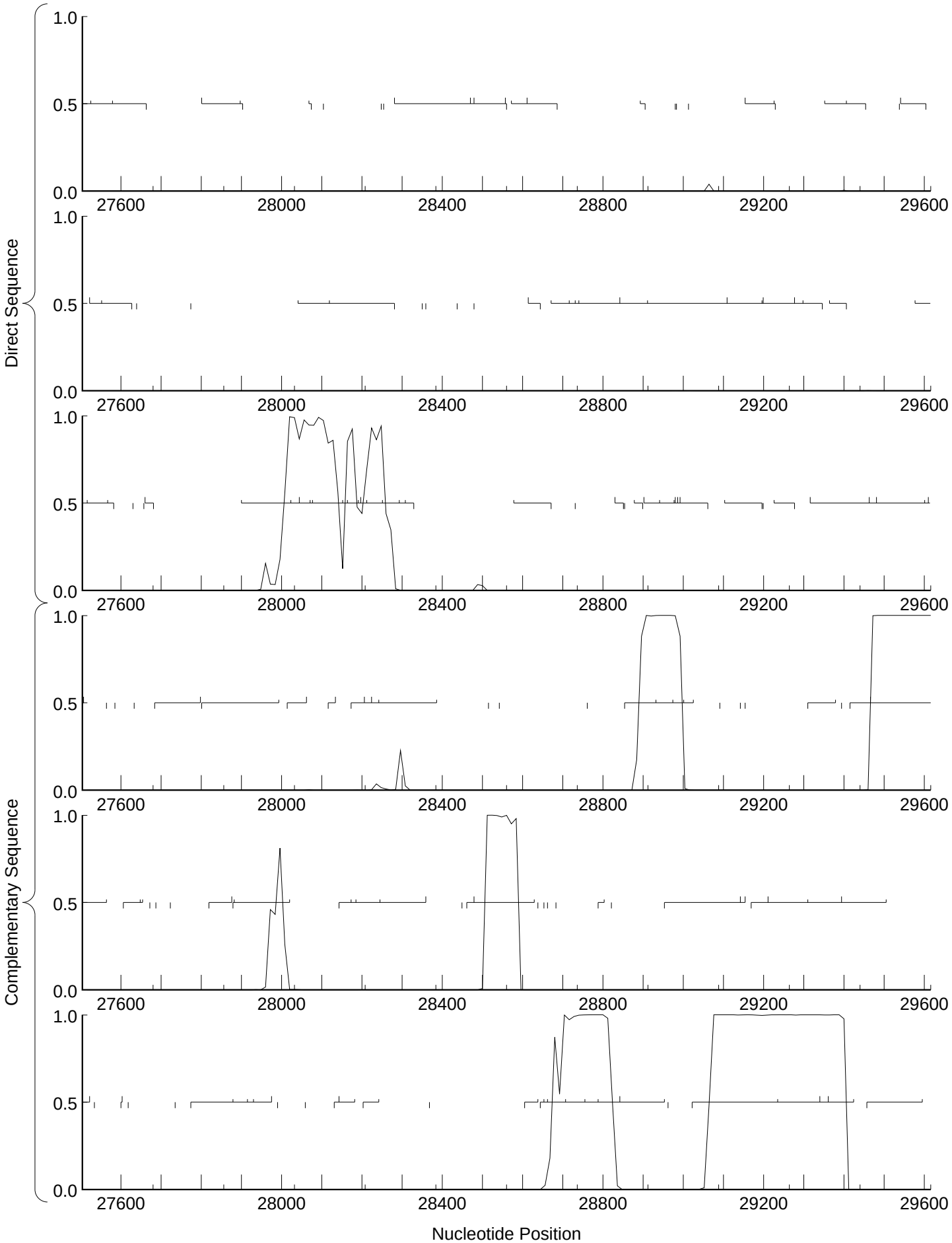
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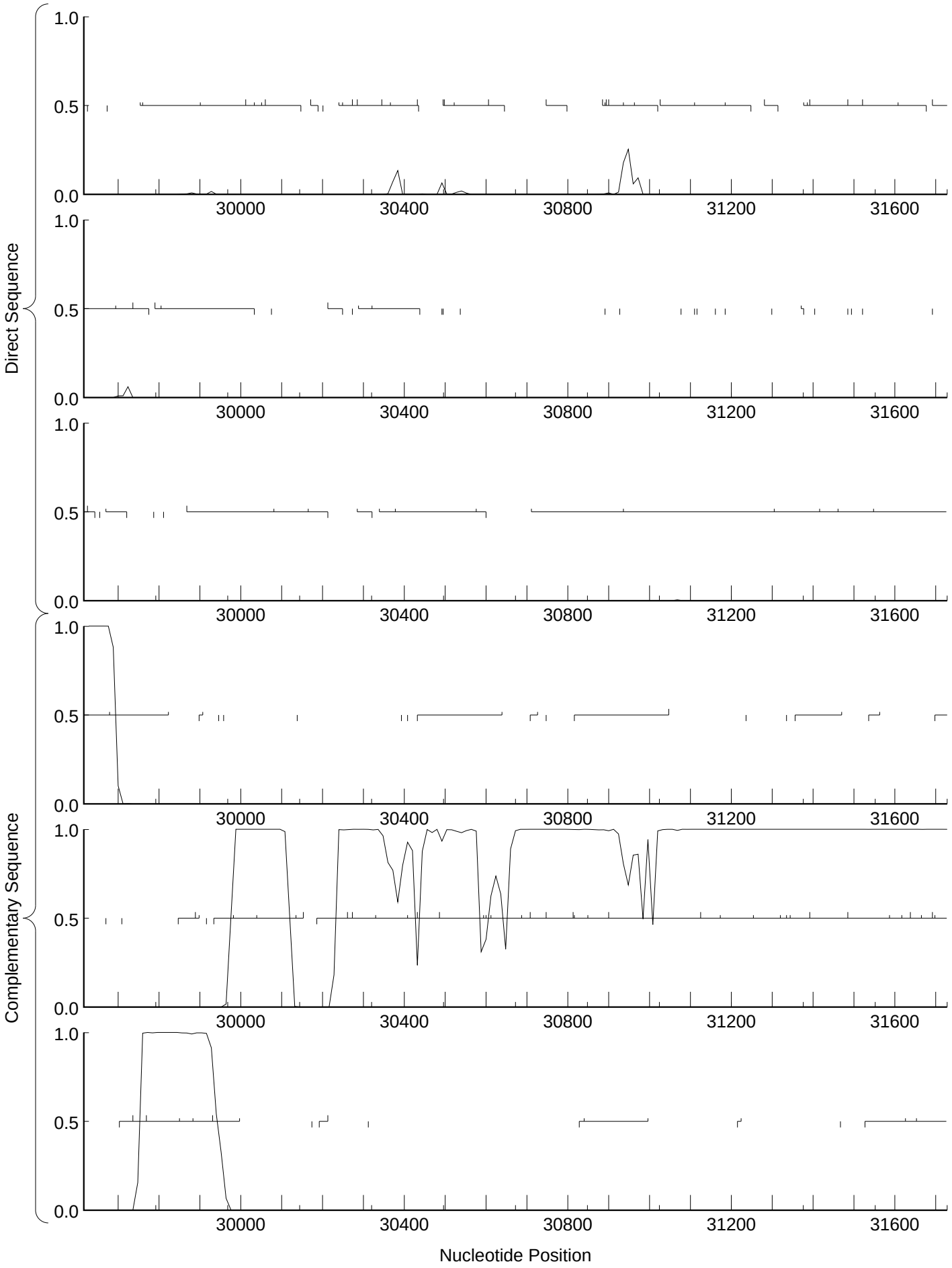
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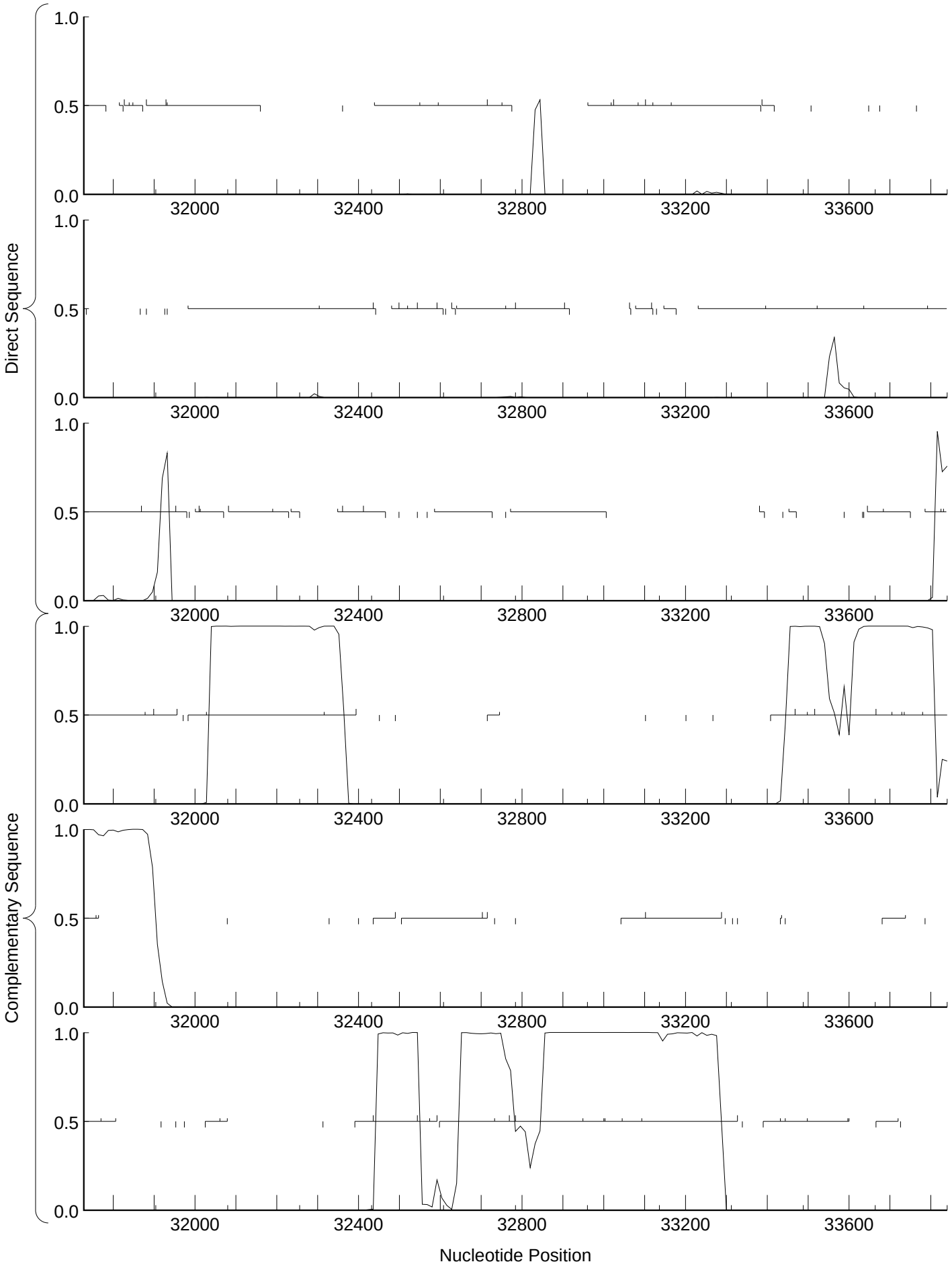
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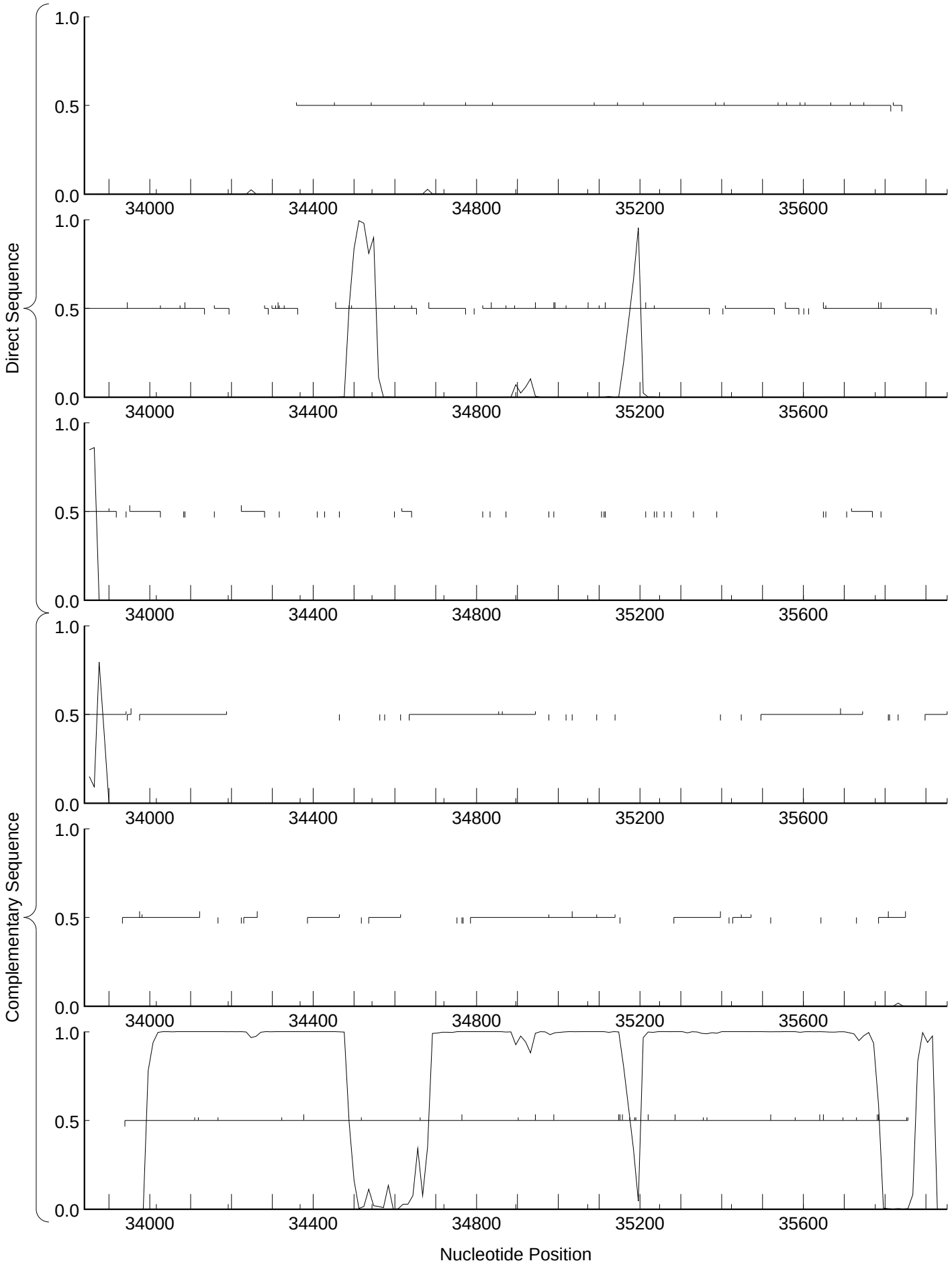
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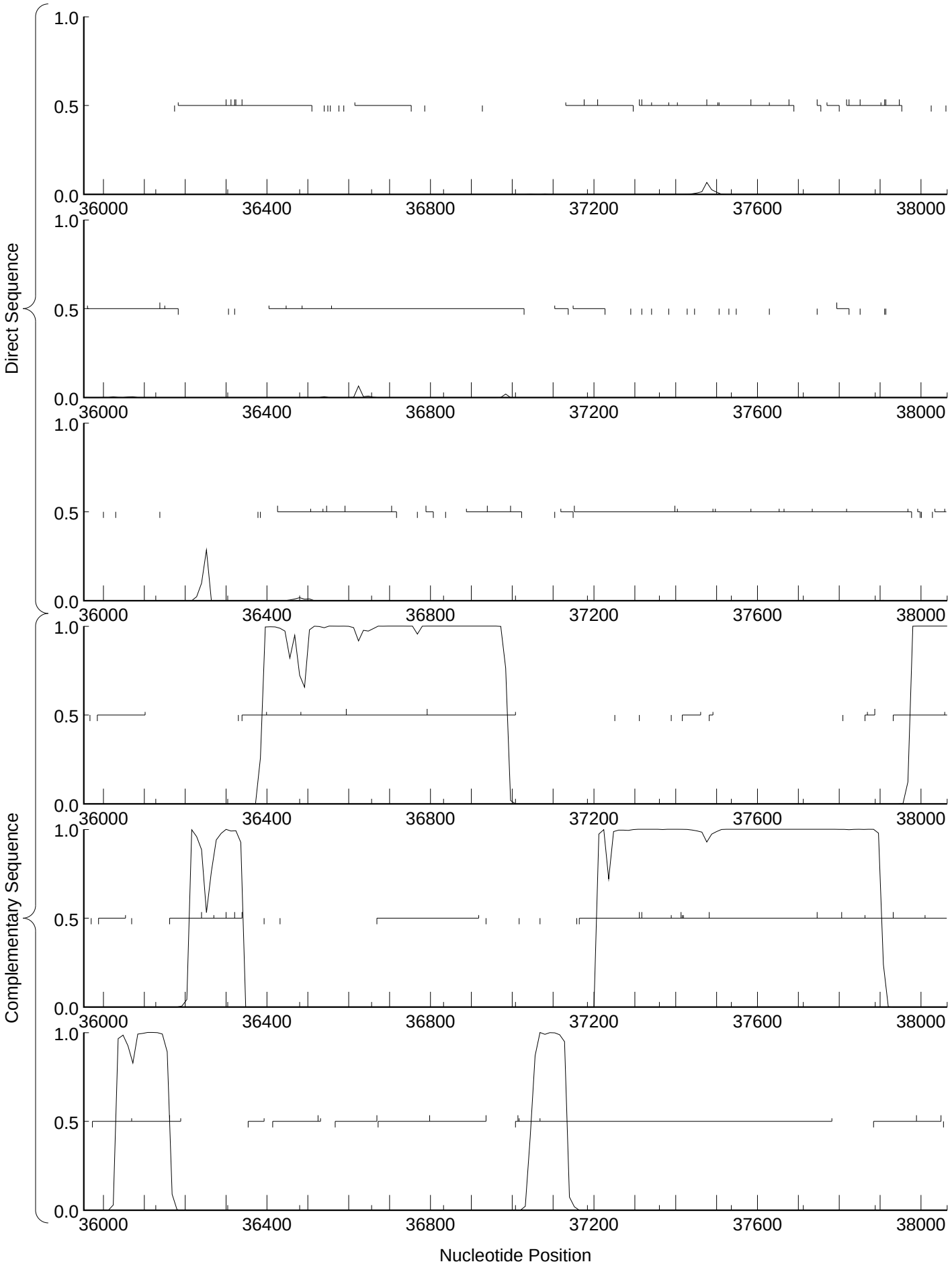
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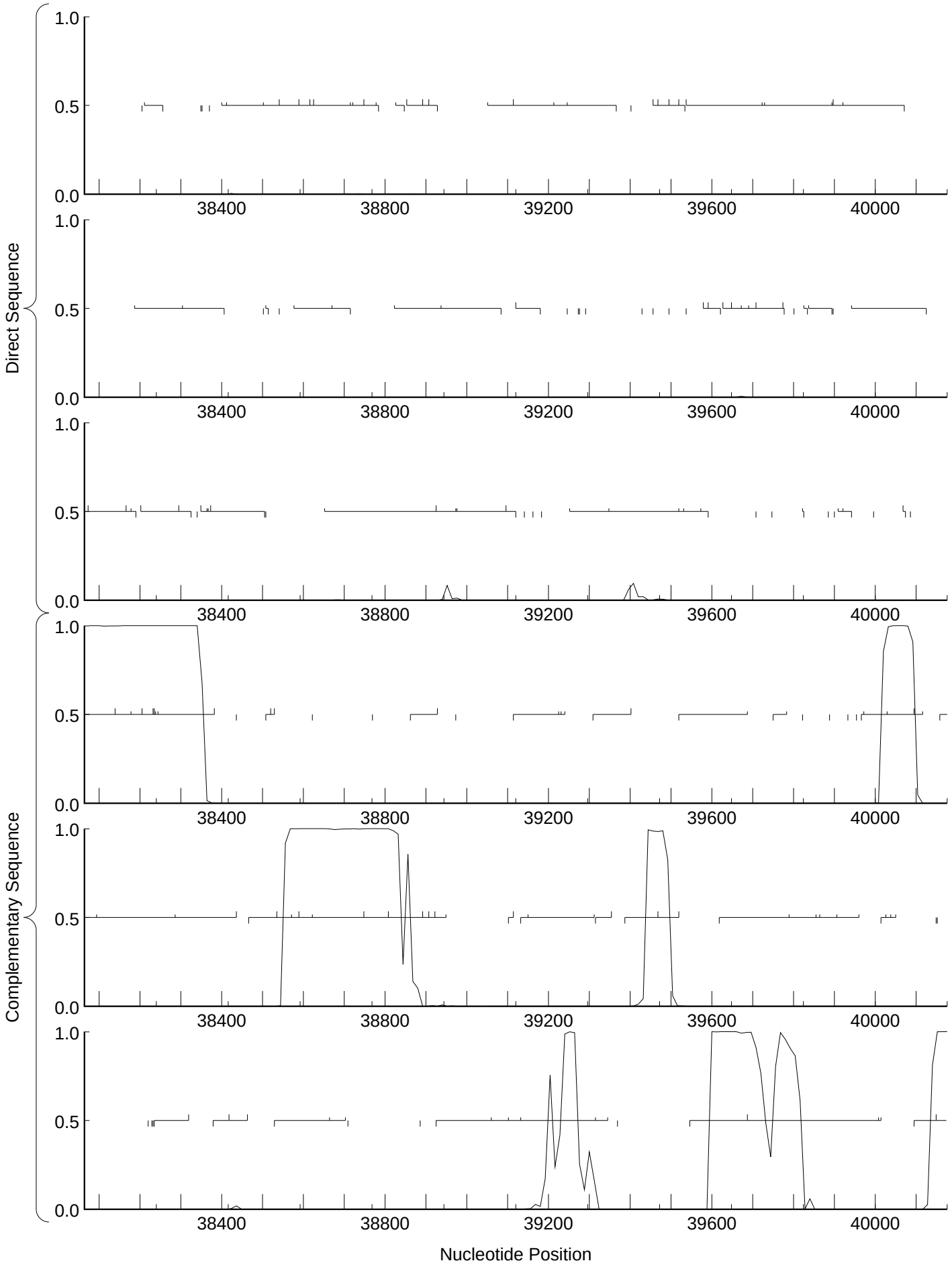
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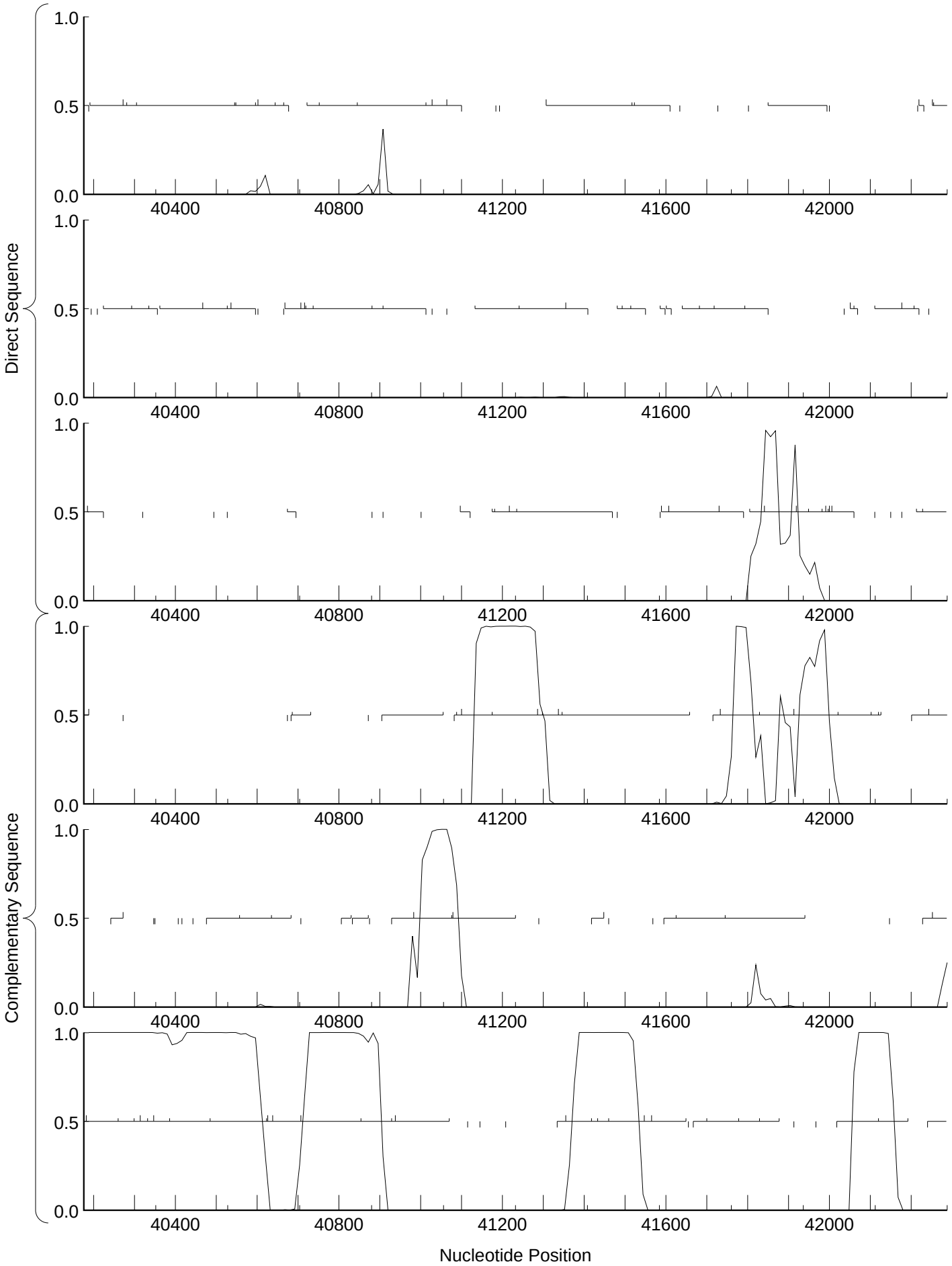
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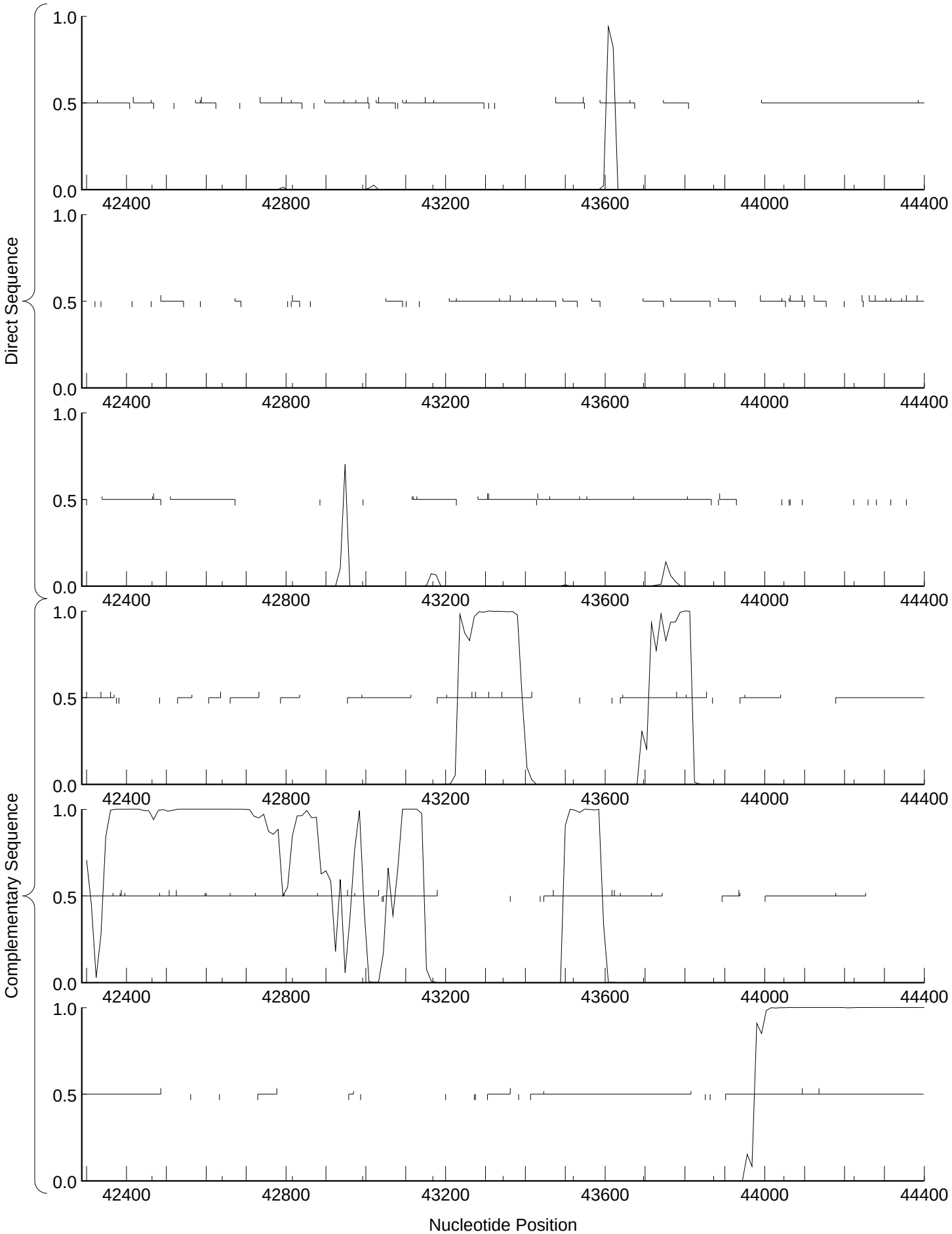
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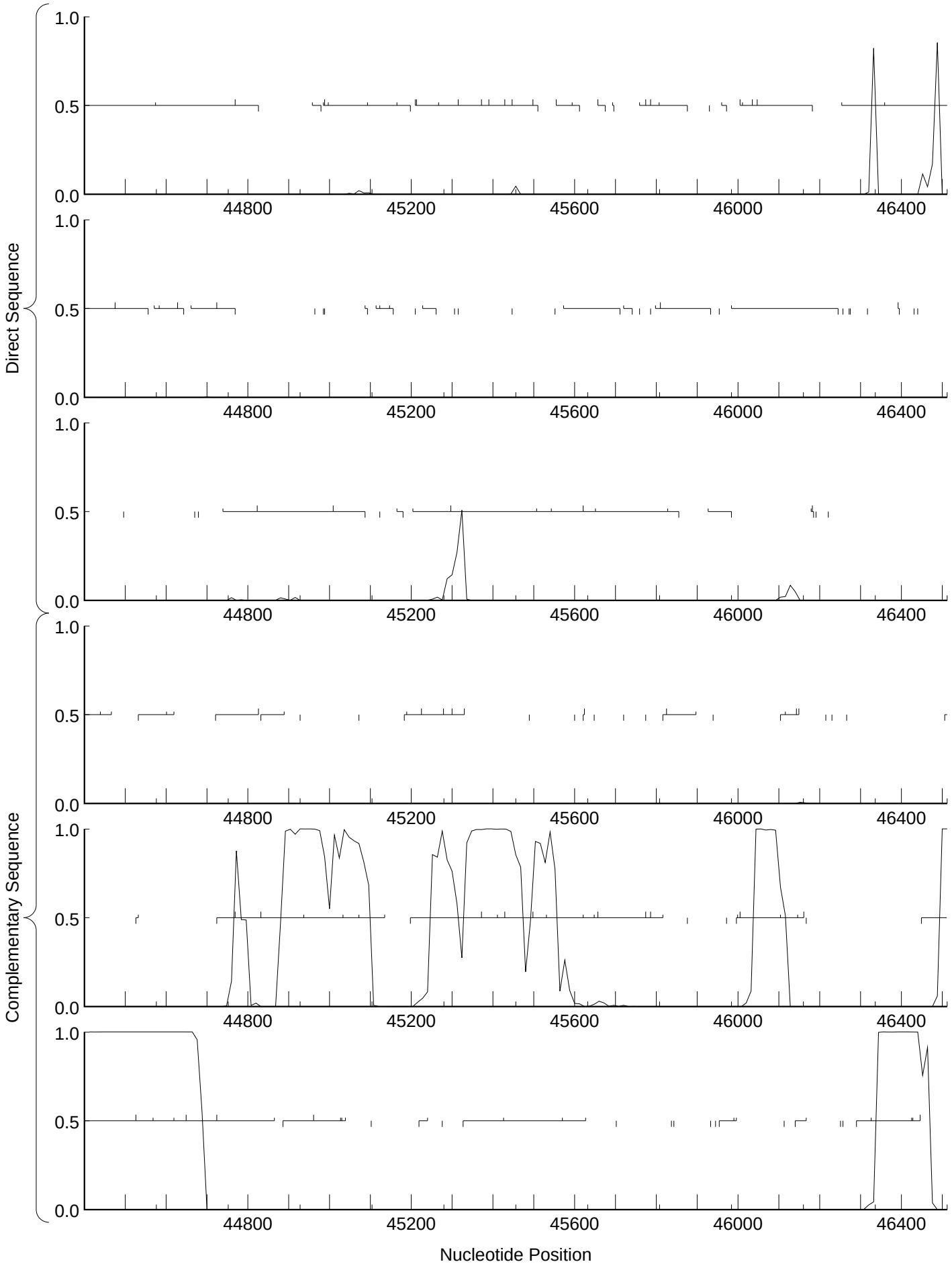
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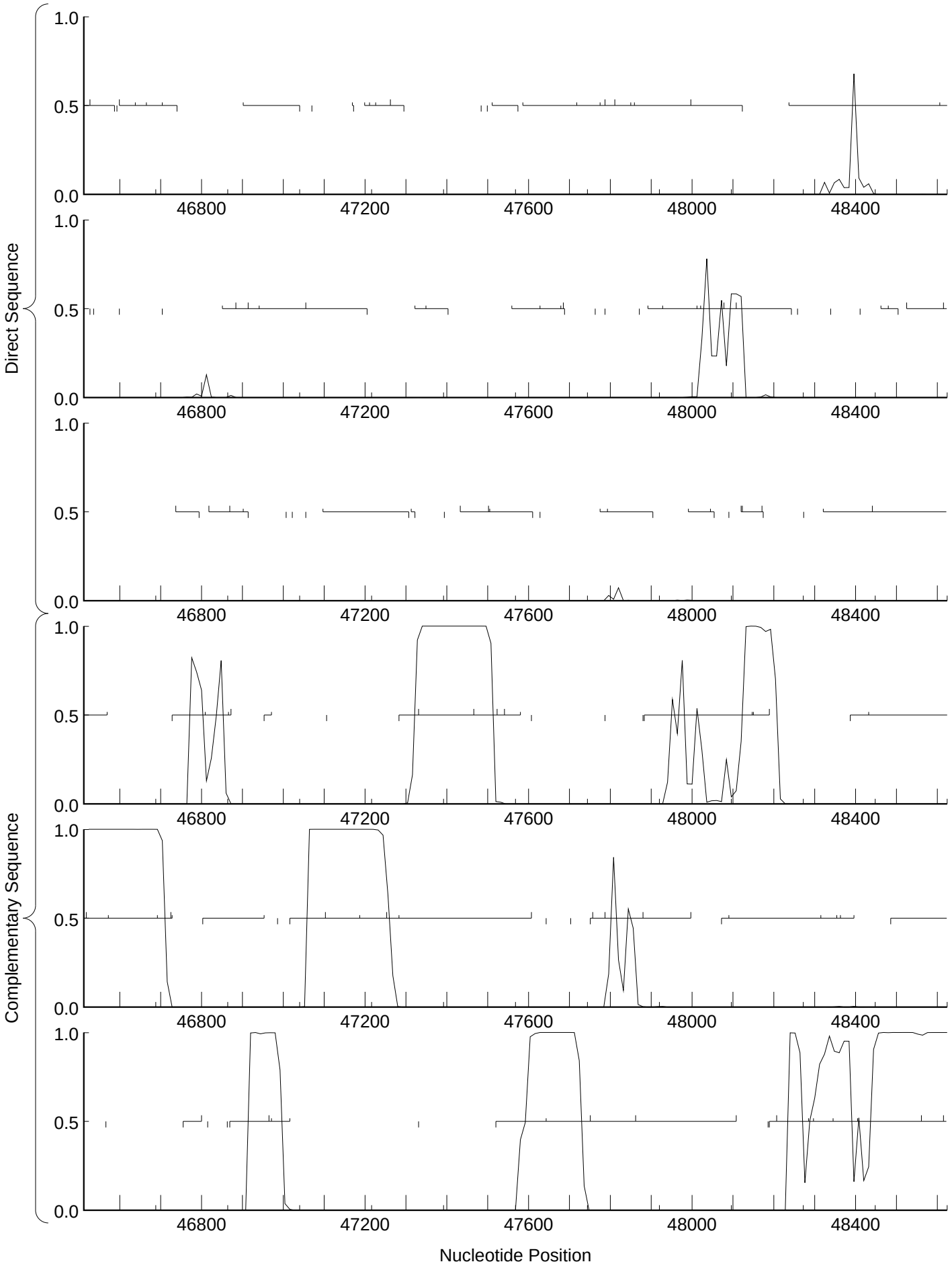
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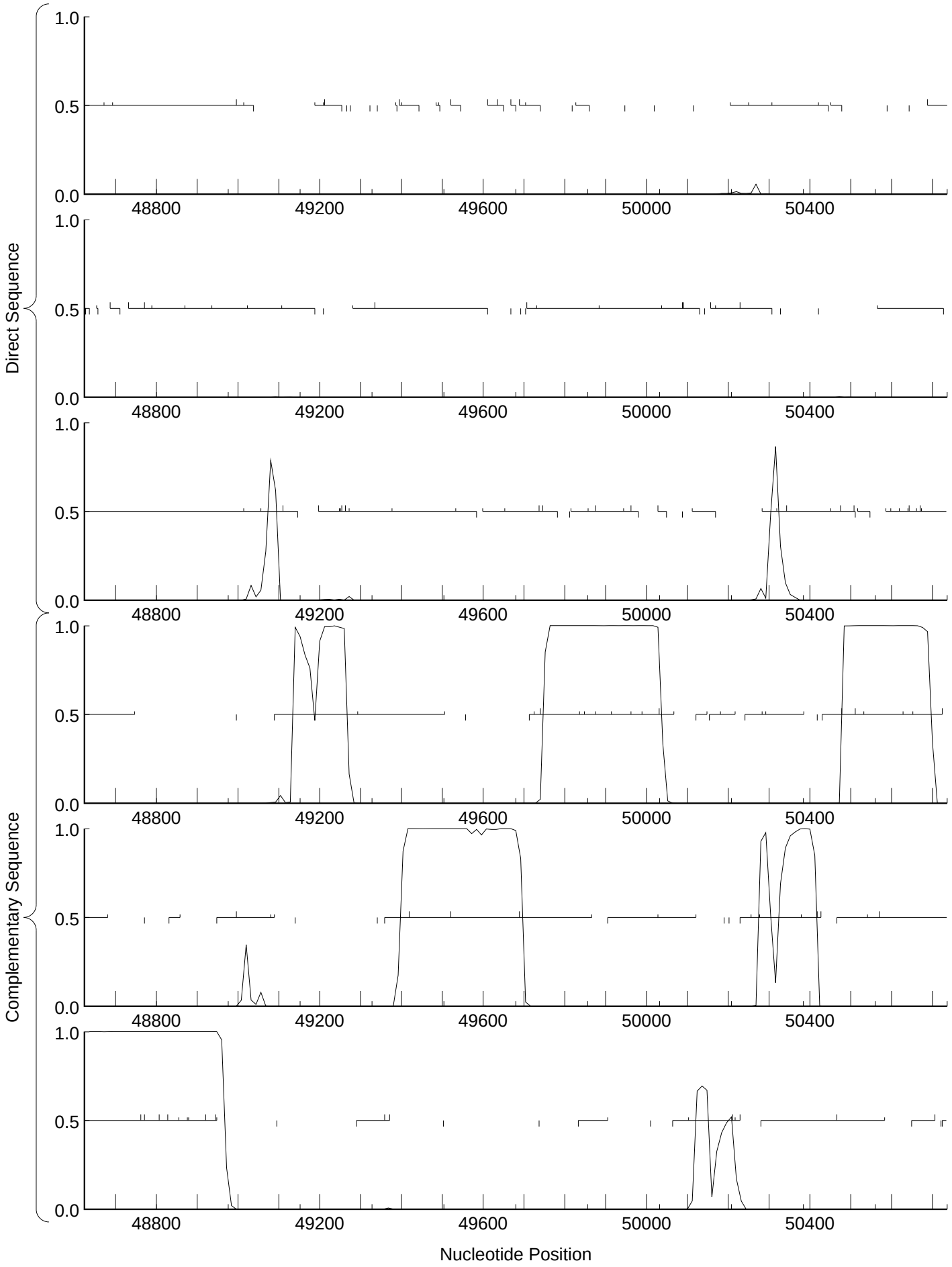
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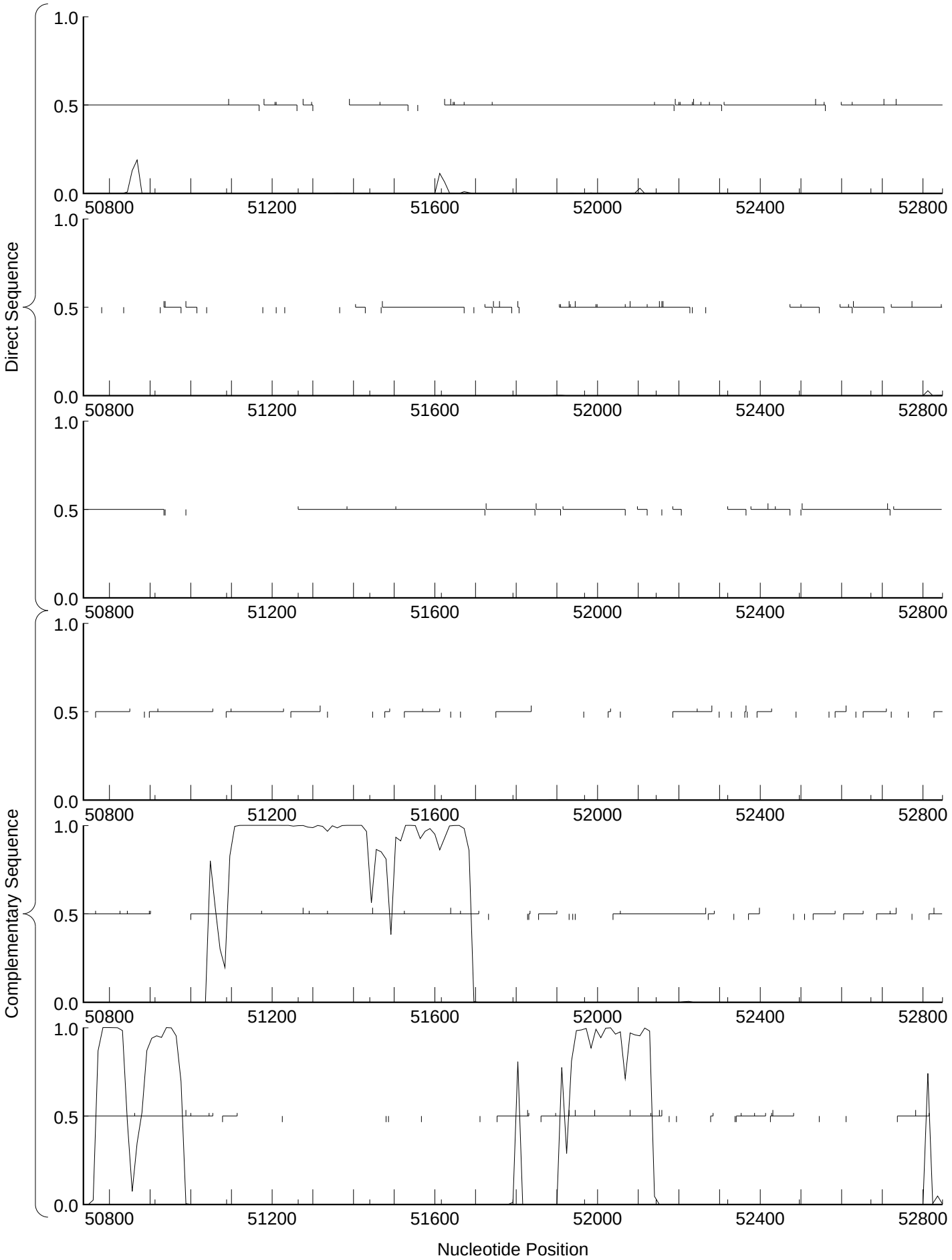
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e complete sequence, 53330 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 25/27



e complete sequence, 53330 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 26/27



e complete sequence, 53330 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 27/27

