GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage EagleEye complete sequence, 52974 bp including 10-base 3' overhang (CGG
Analysis Date    :  3/28/24 at 17:48:48
Pages            :  27
Sequence Length  :  52974 bp
GC Content       :   61.42%

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

PS-Version       :  1.2

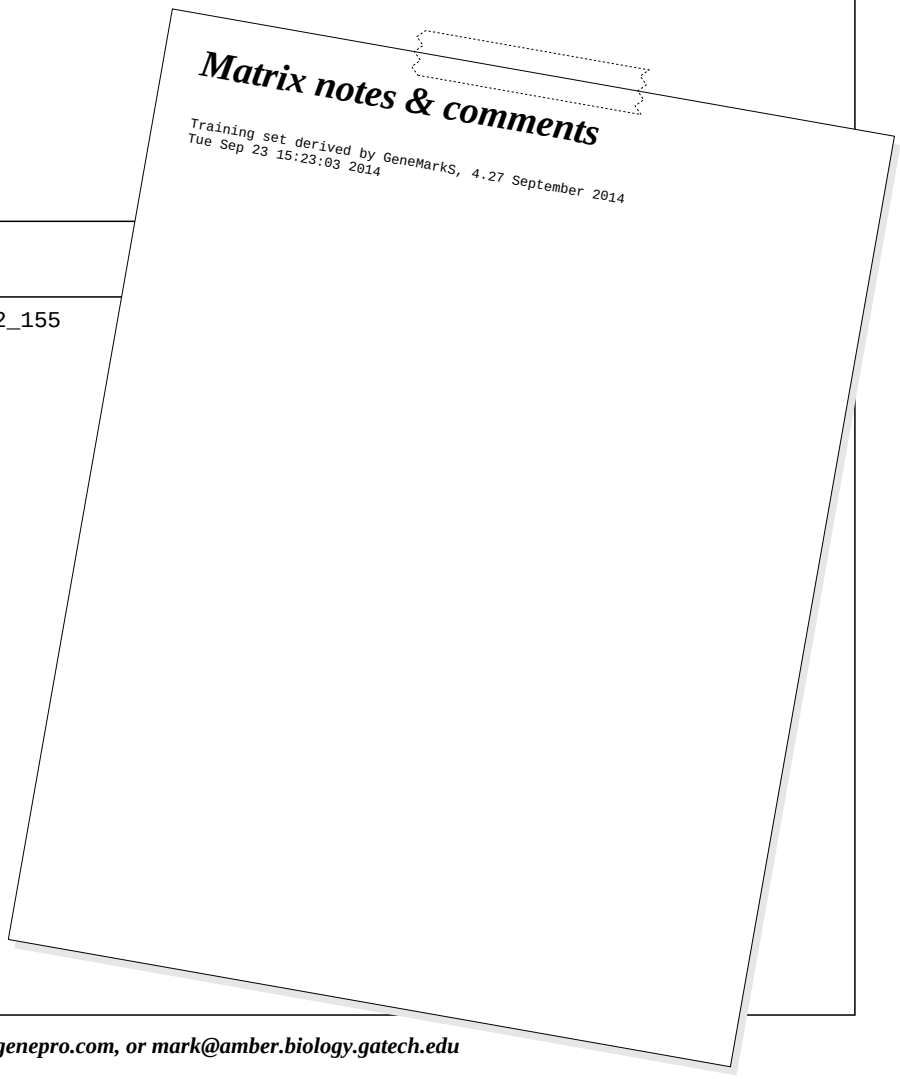
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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                    List regions and/or splice sites,
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MATRIX INFORMATION

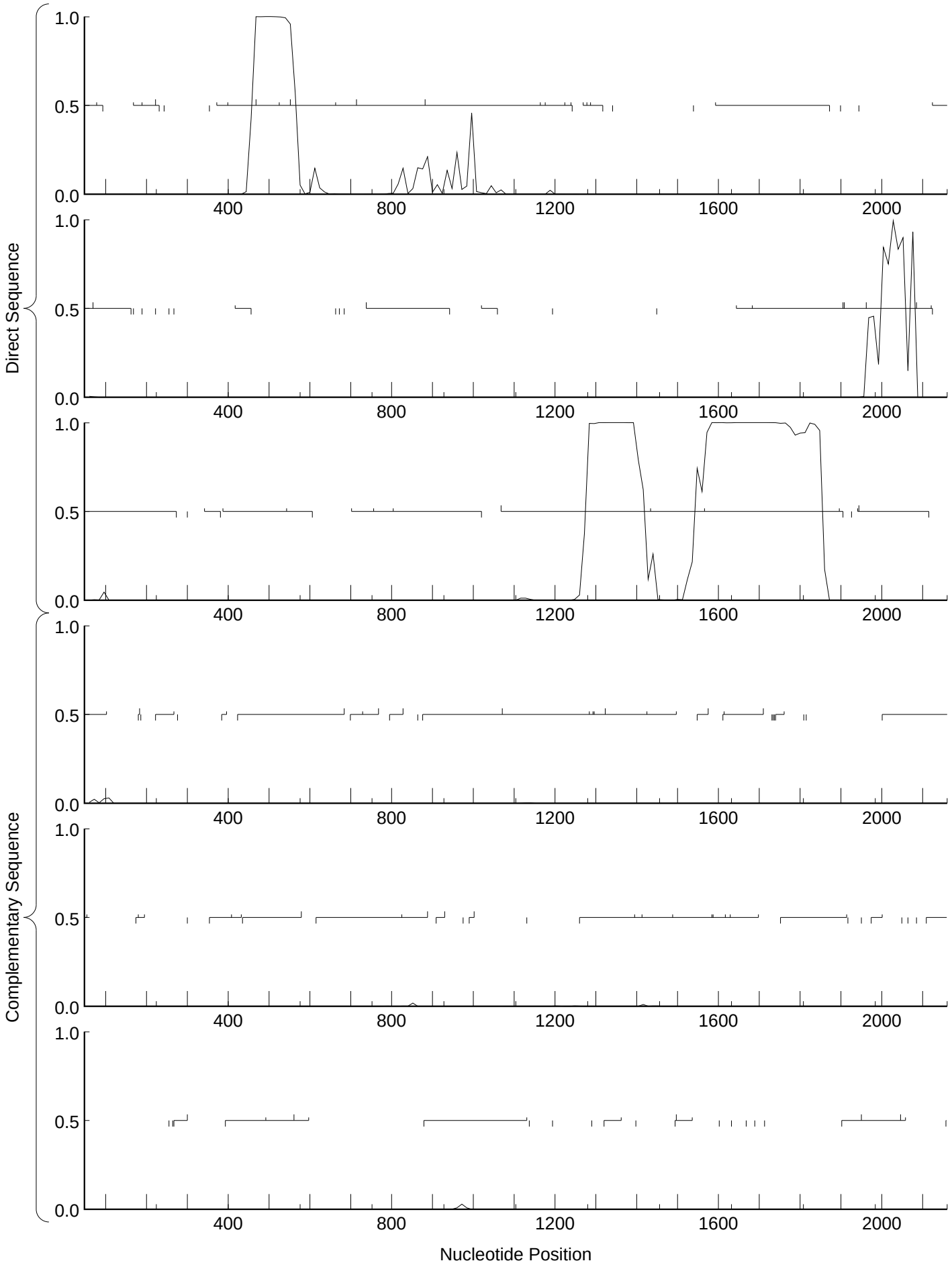
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Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

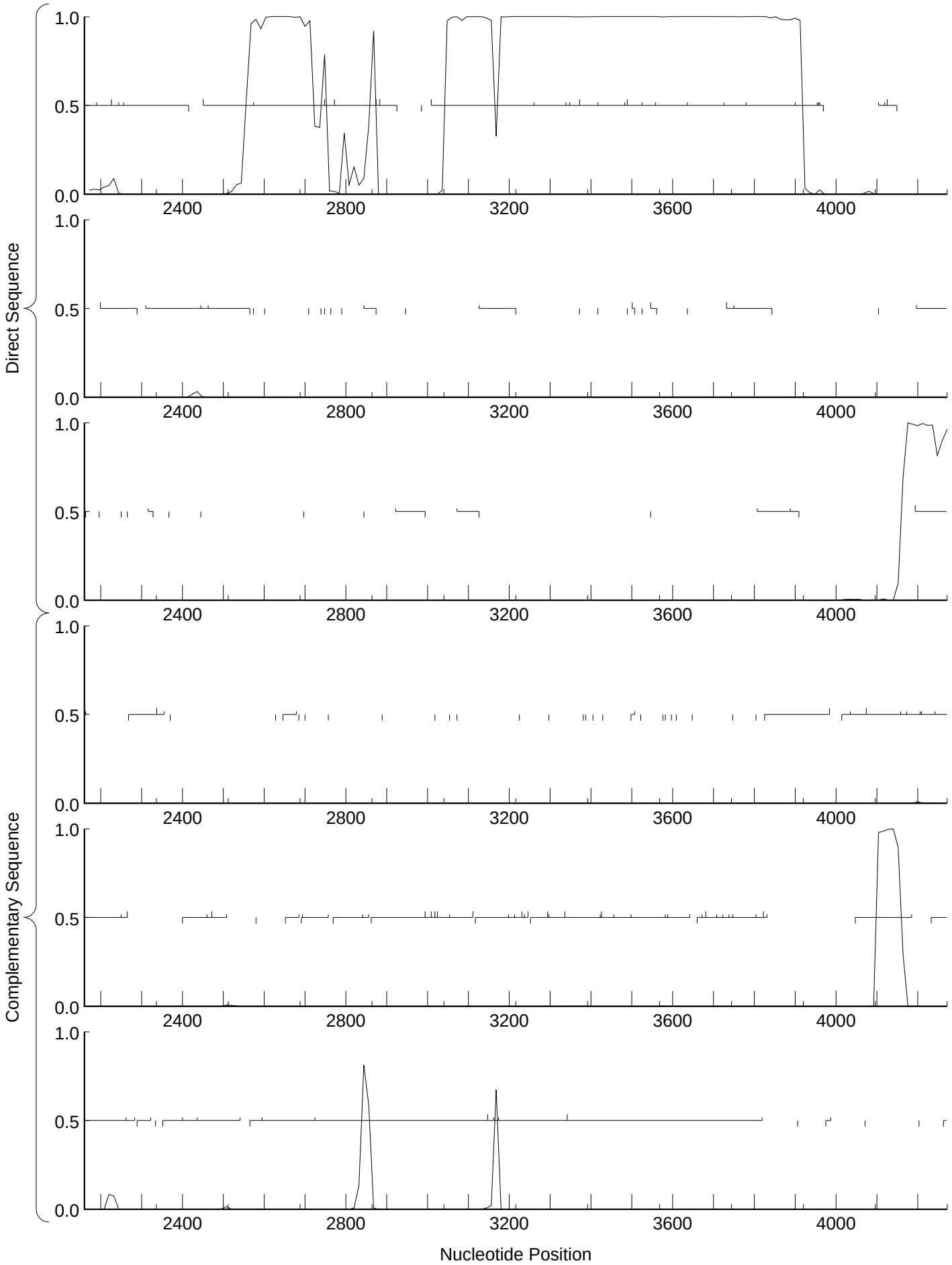
Order      :  4

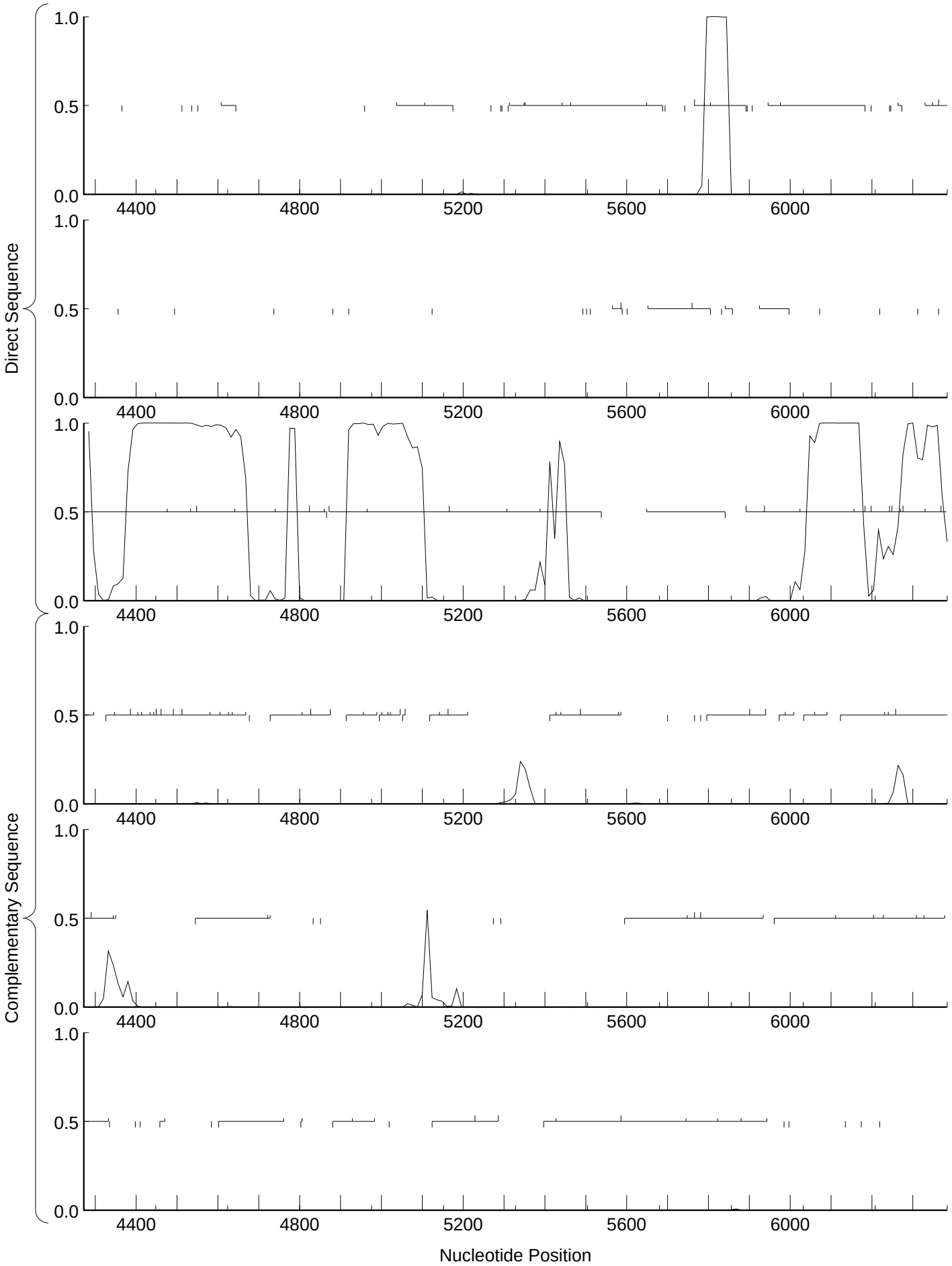

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

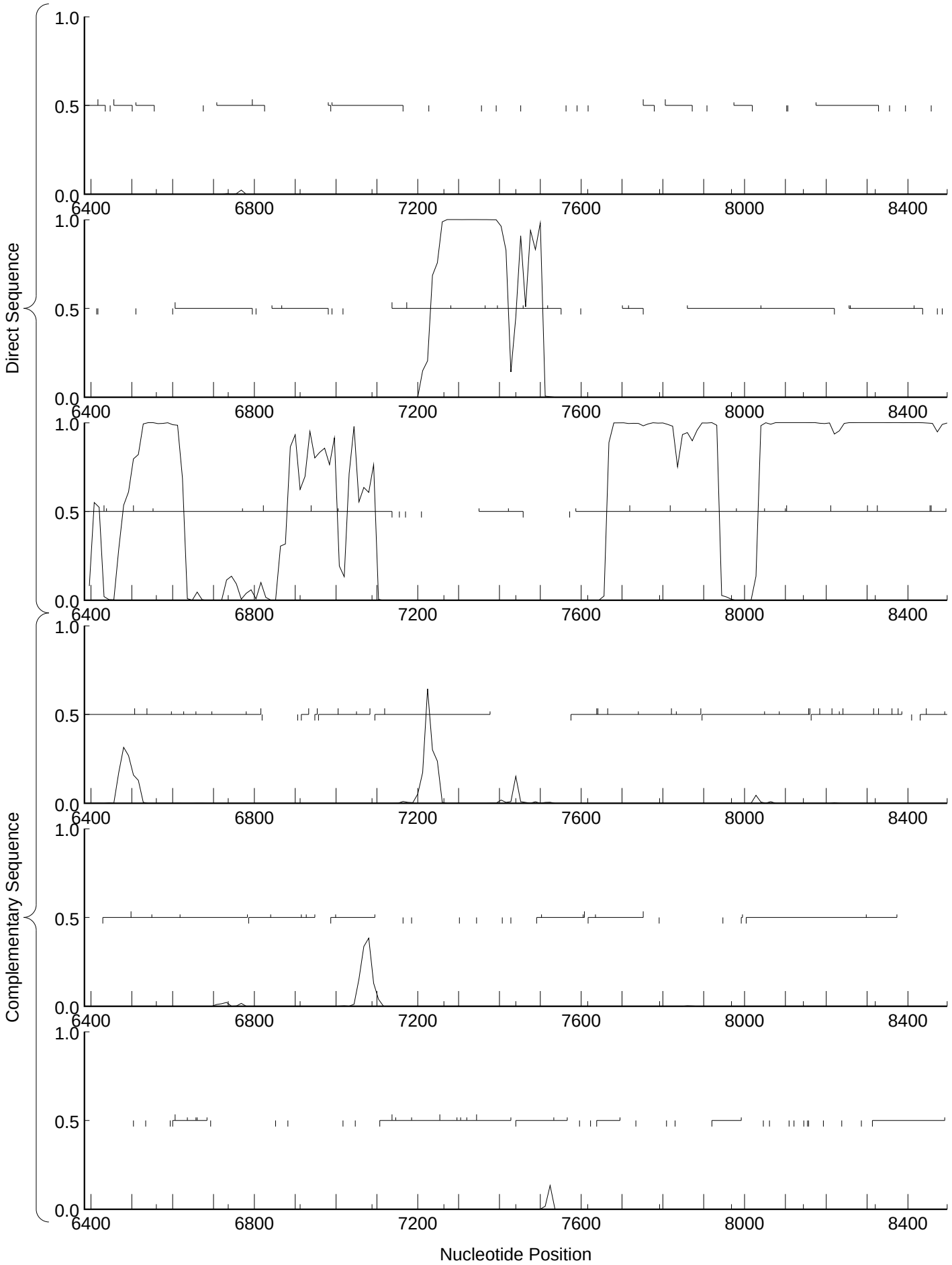


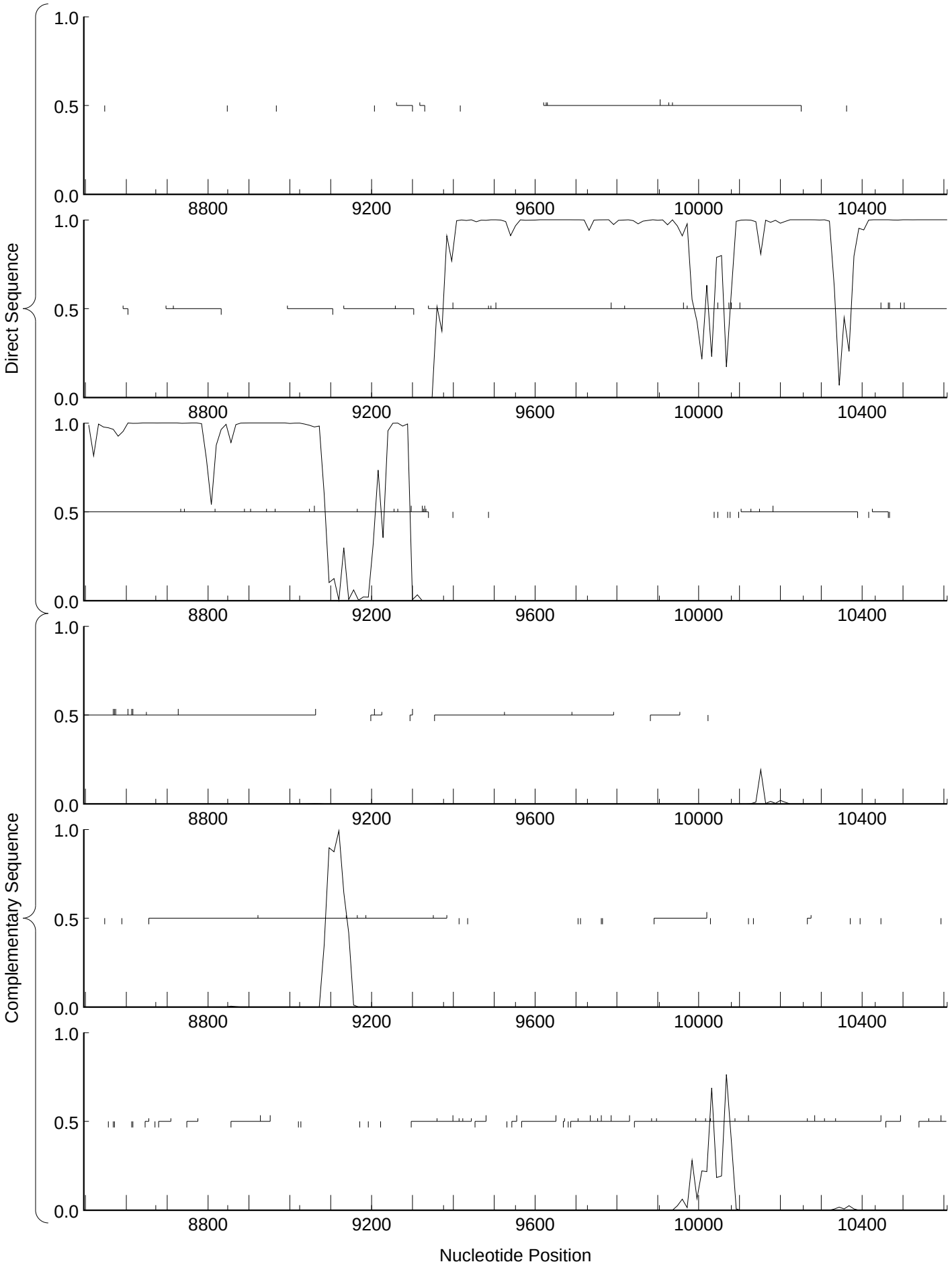
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu


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Complementary Sequence
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0.0
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1600
2000
Nucleotide Position

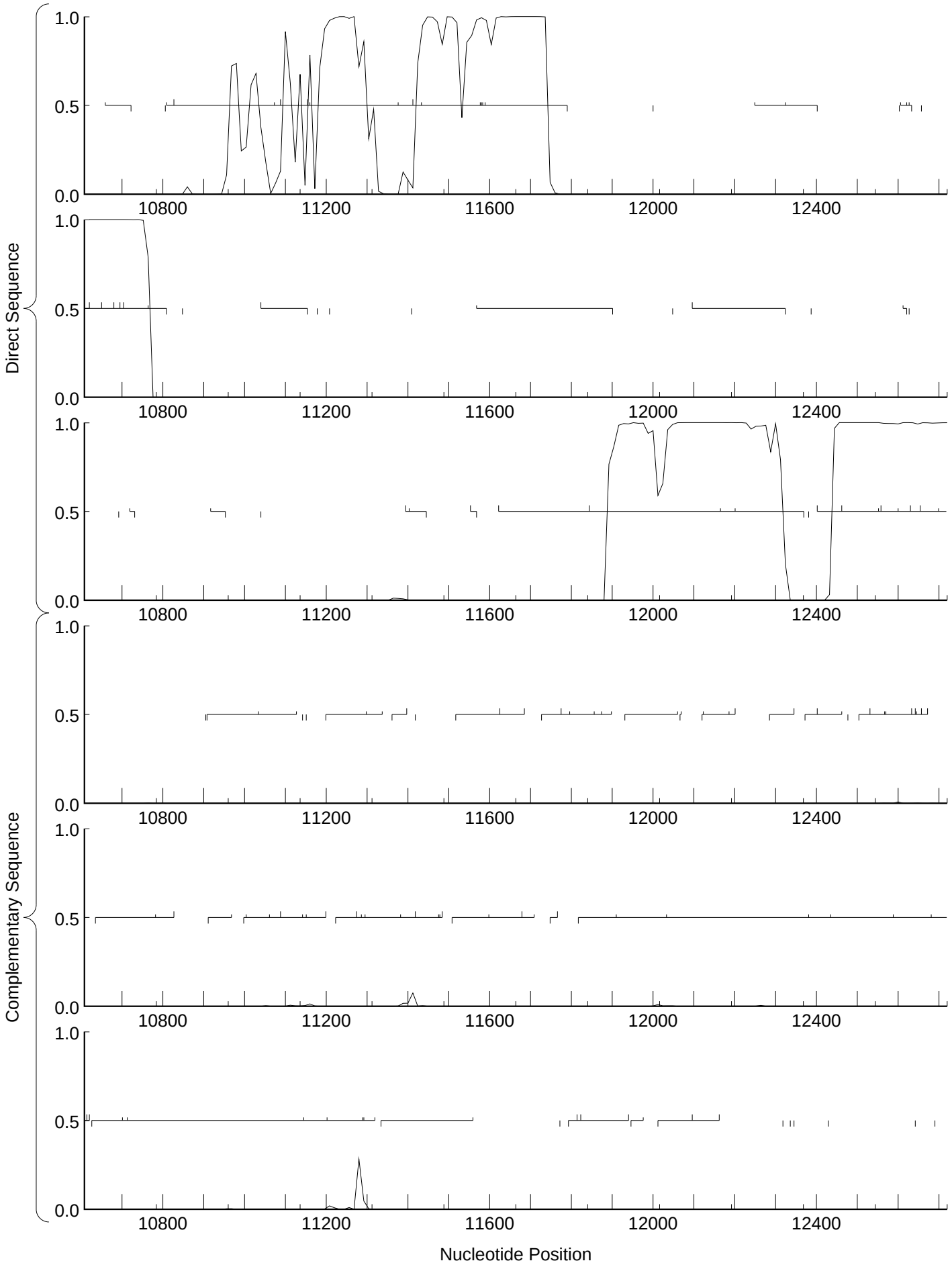

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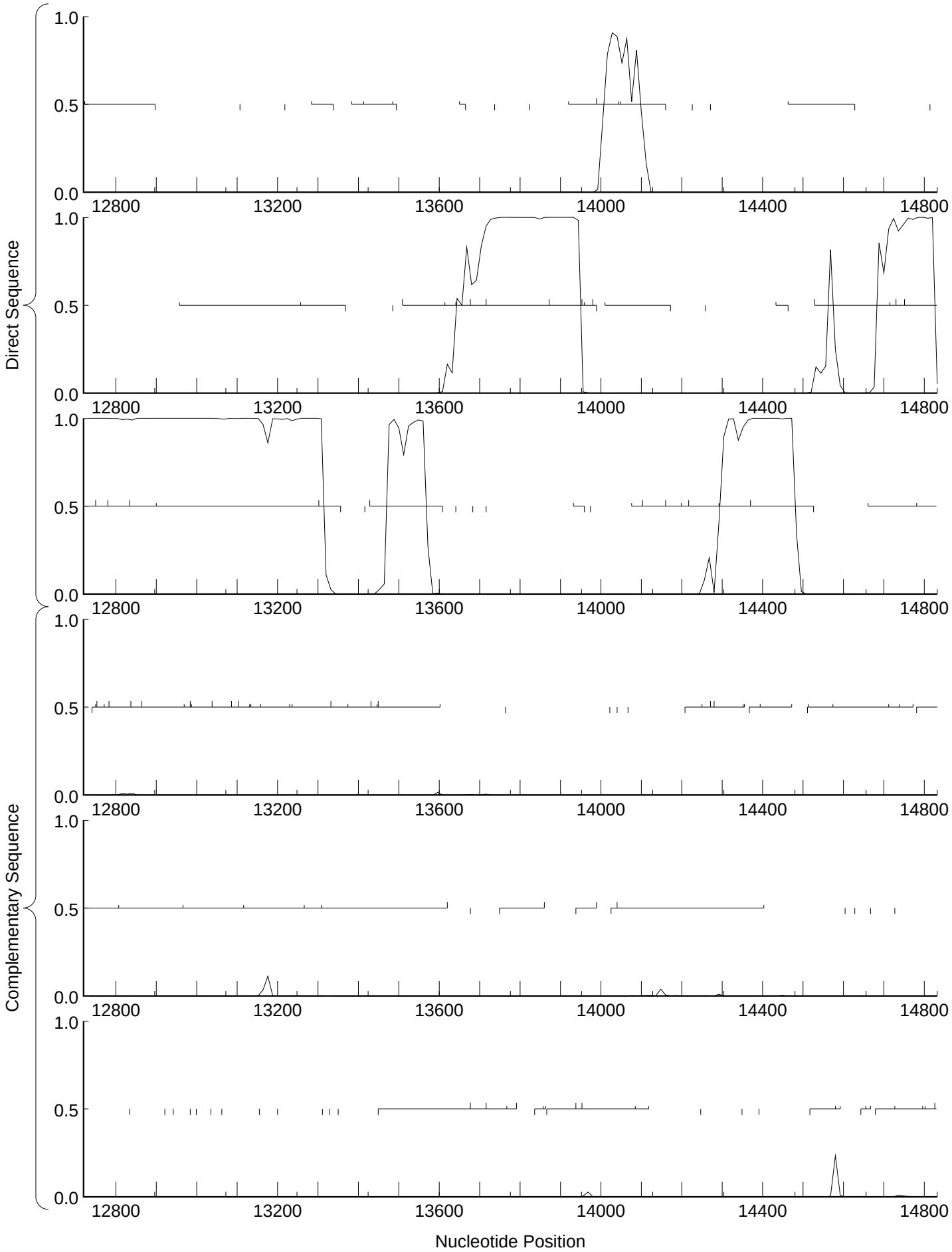

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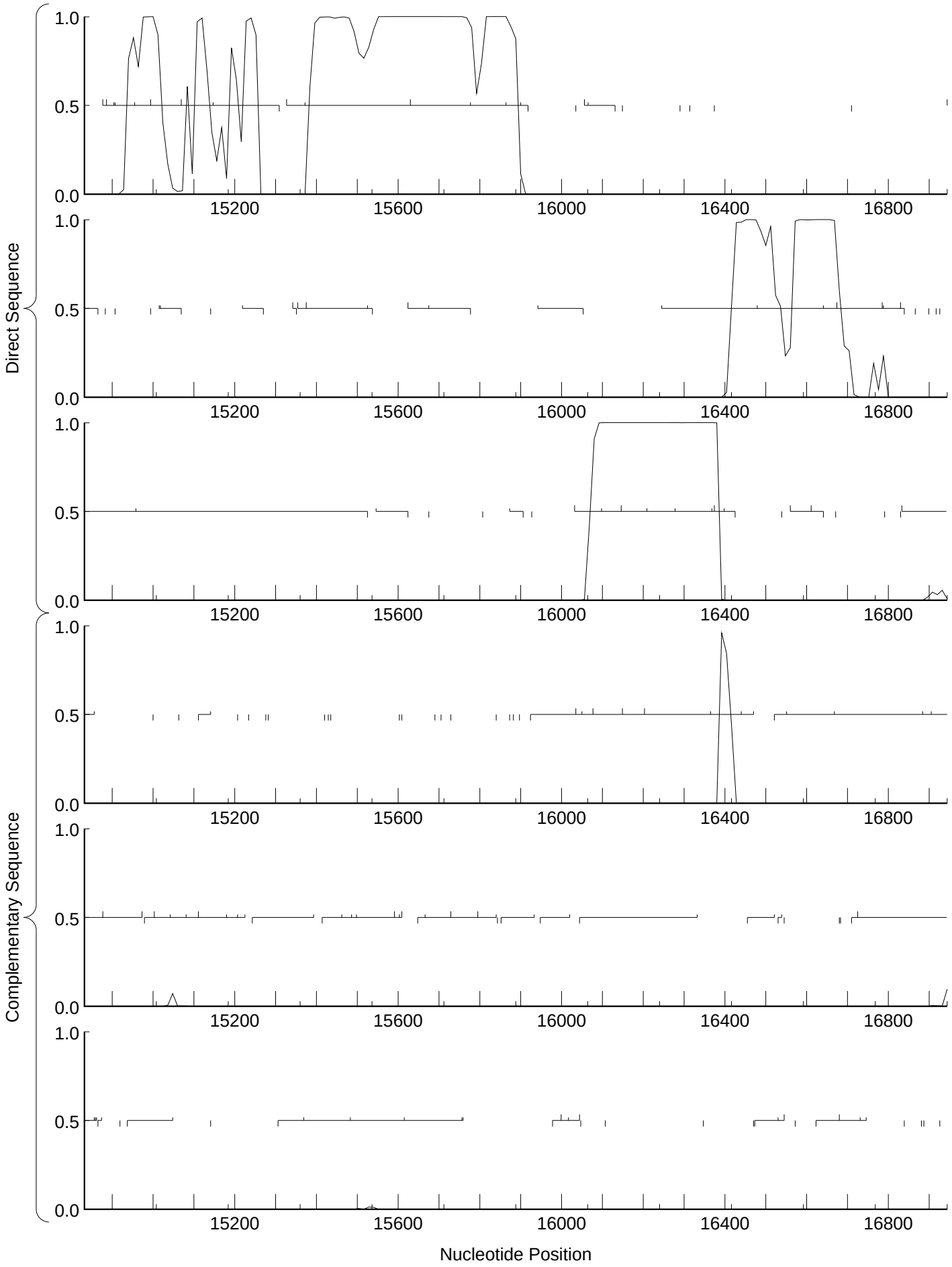

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Nucleotide Position

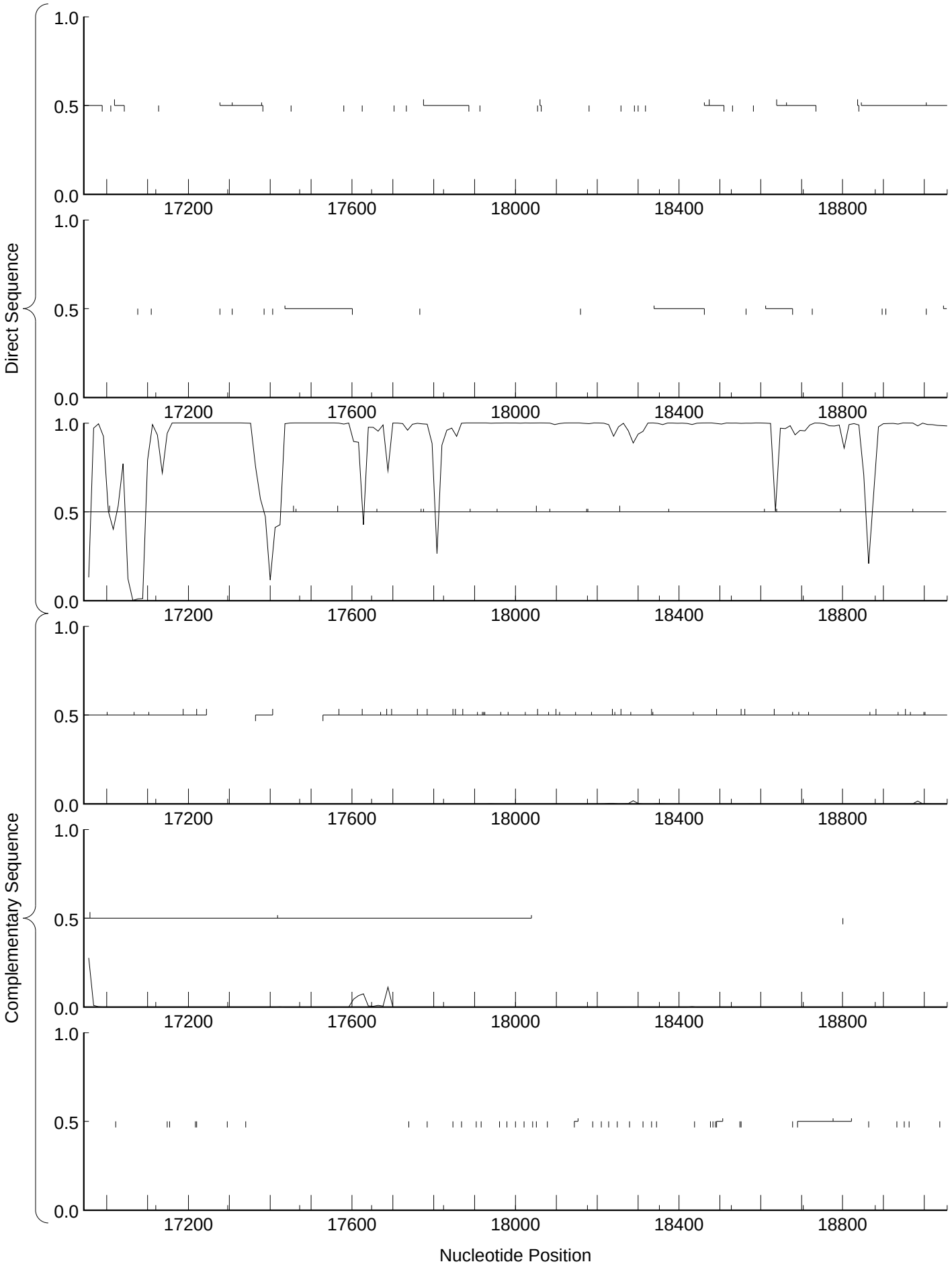
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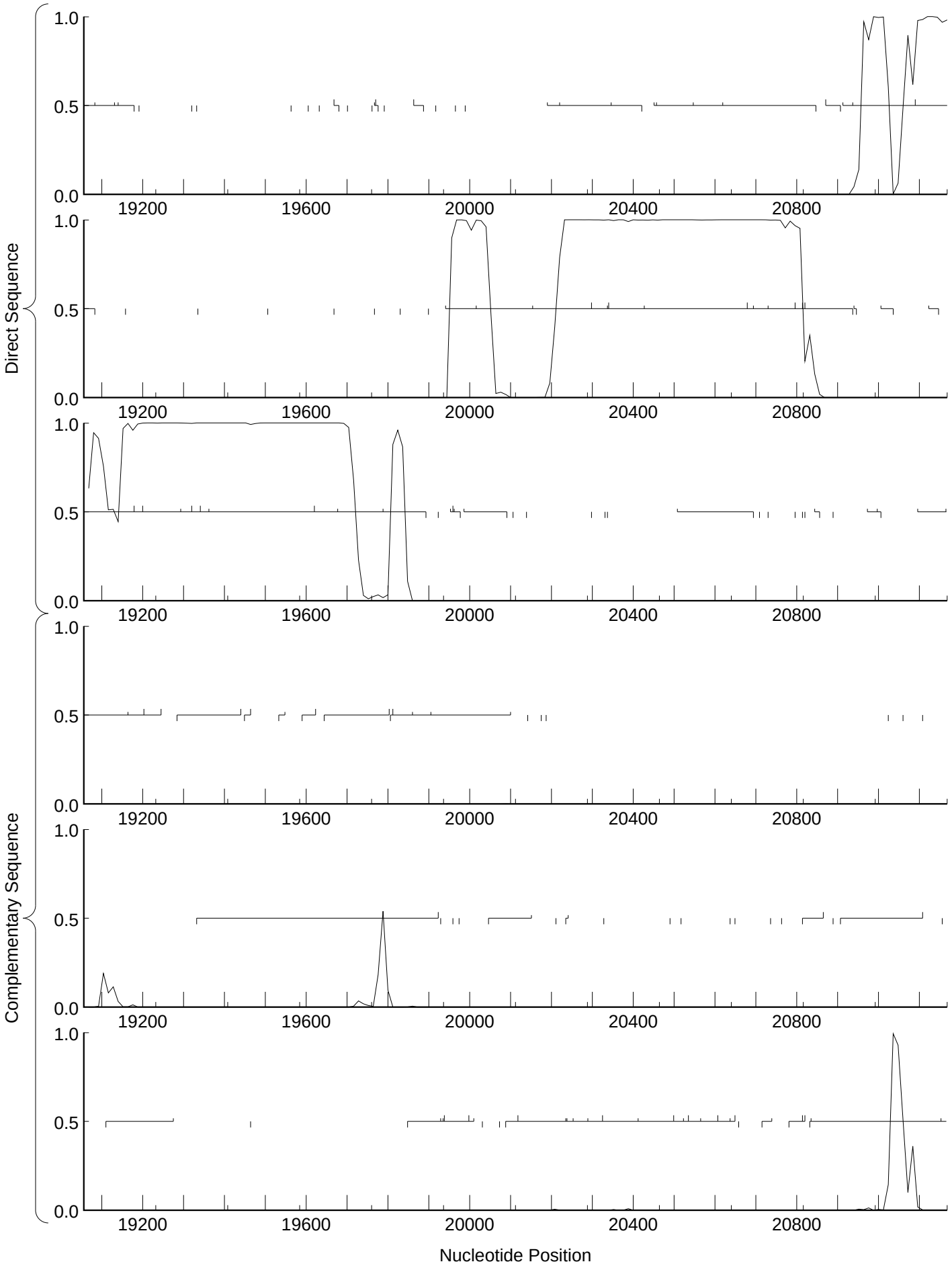

Direct Sequence
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13600
14000
14400
14800
Nucleotide Position


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Nucleotide Position

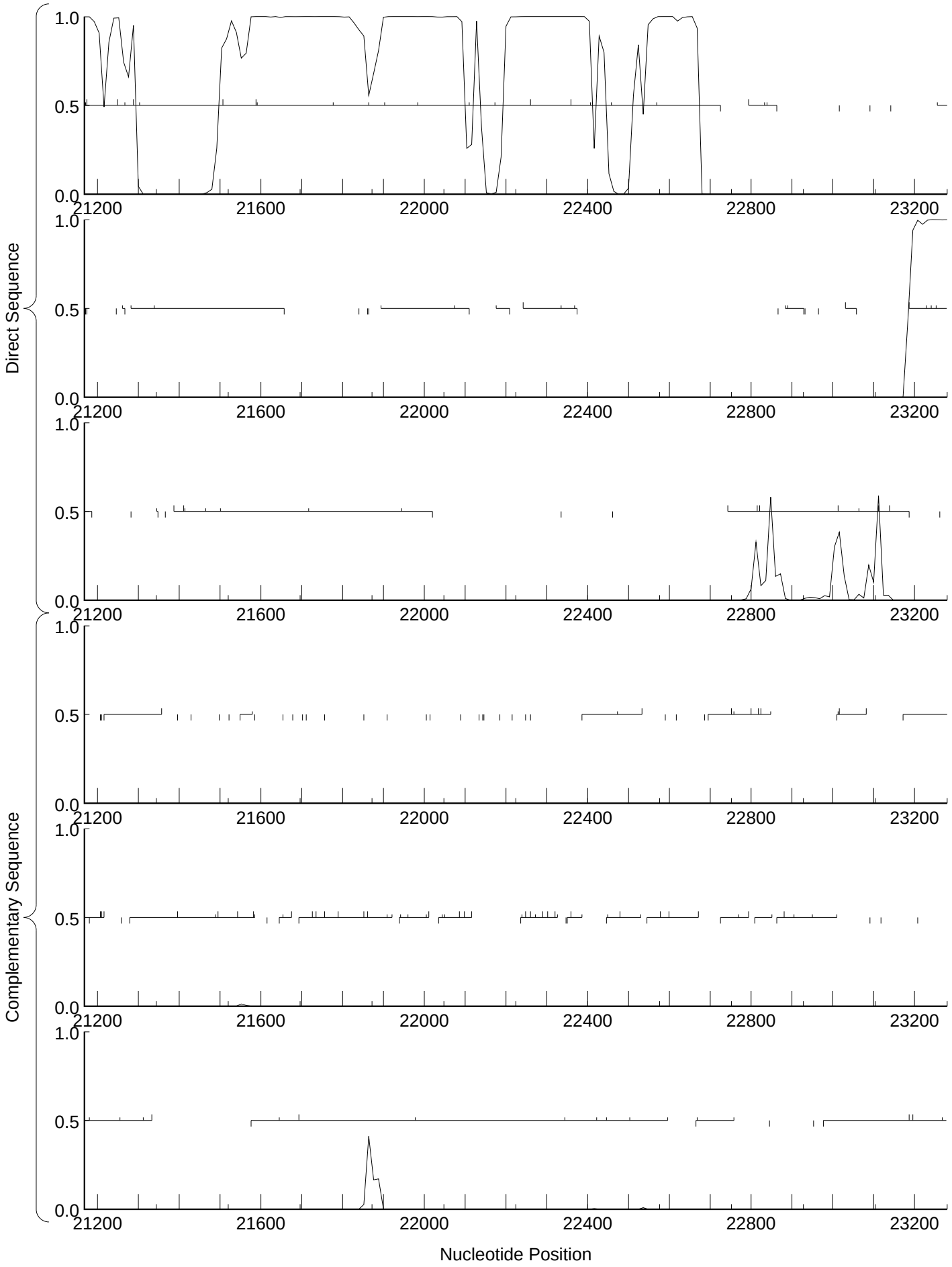
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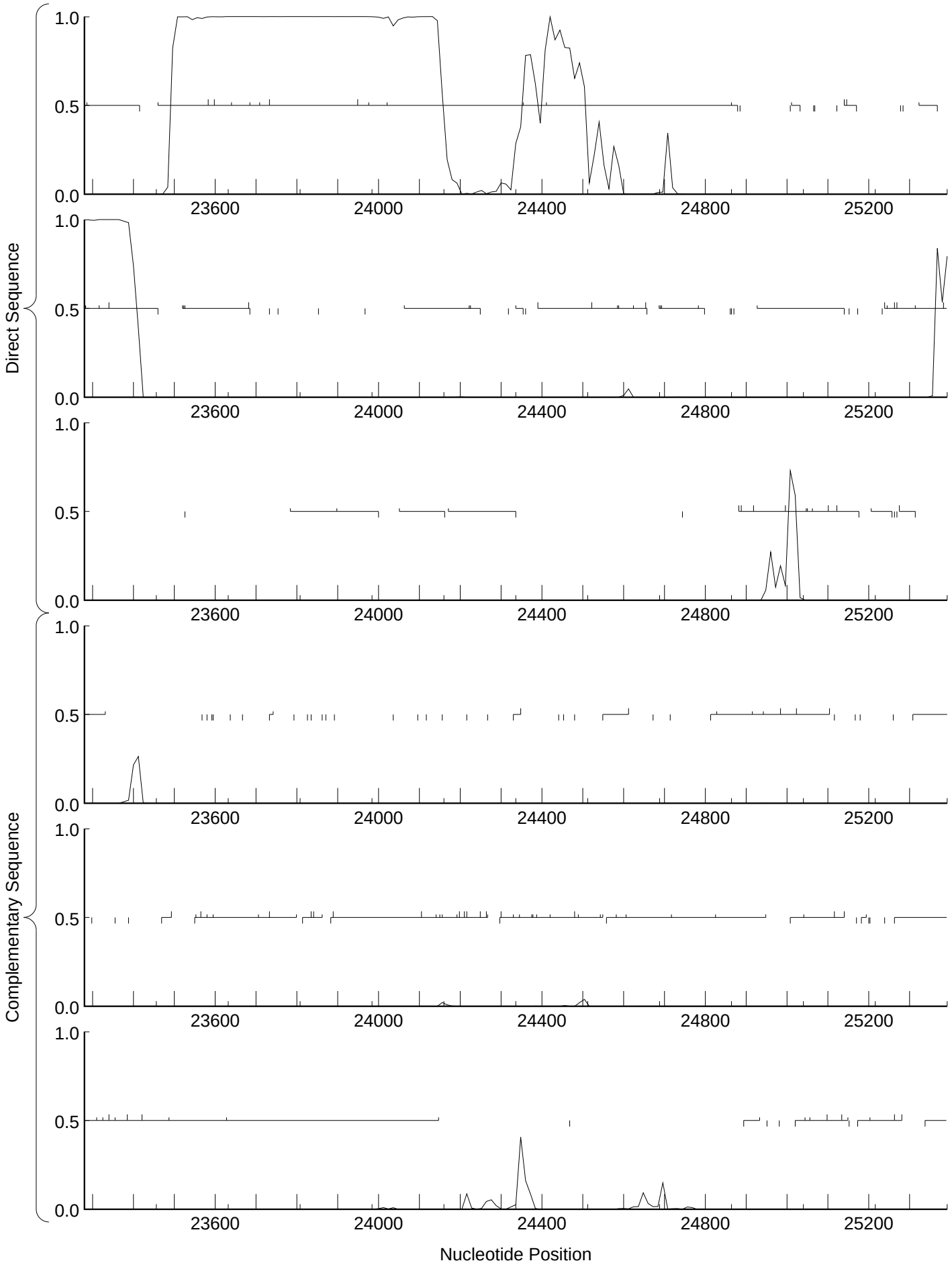
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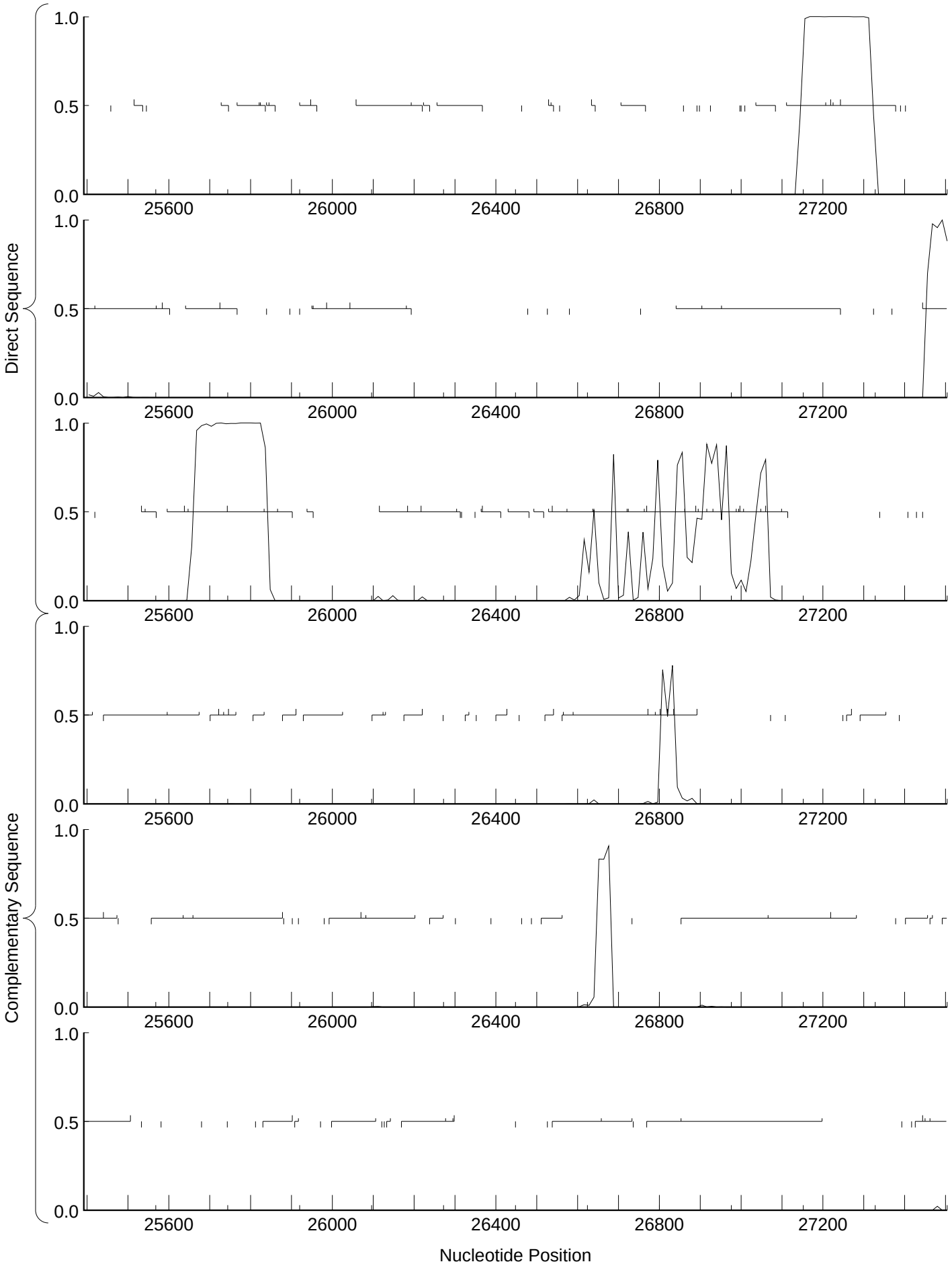


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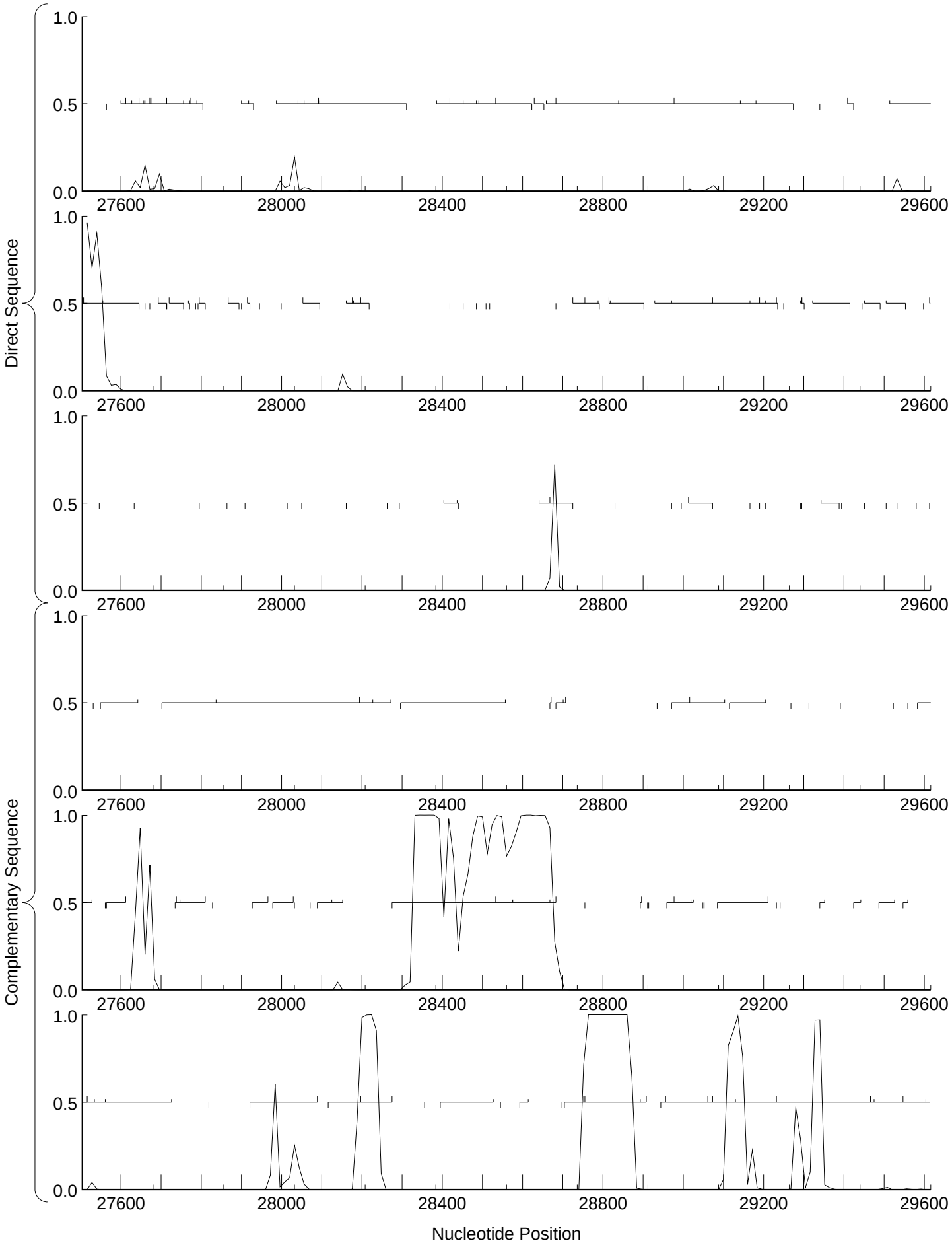


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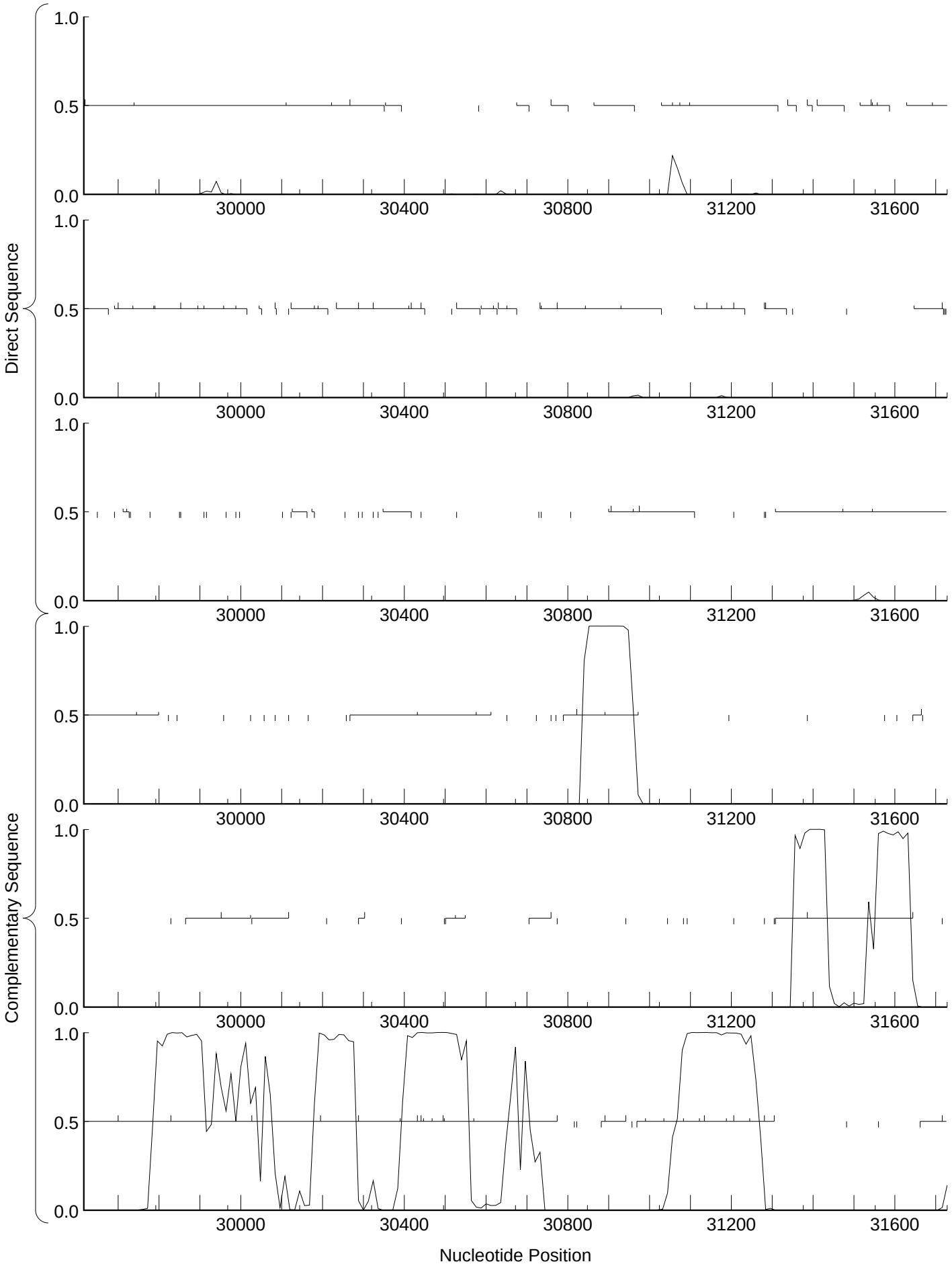



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Nucleotide Position

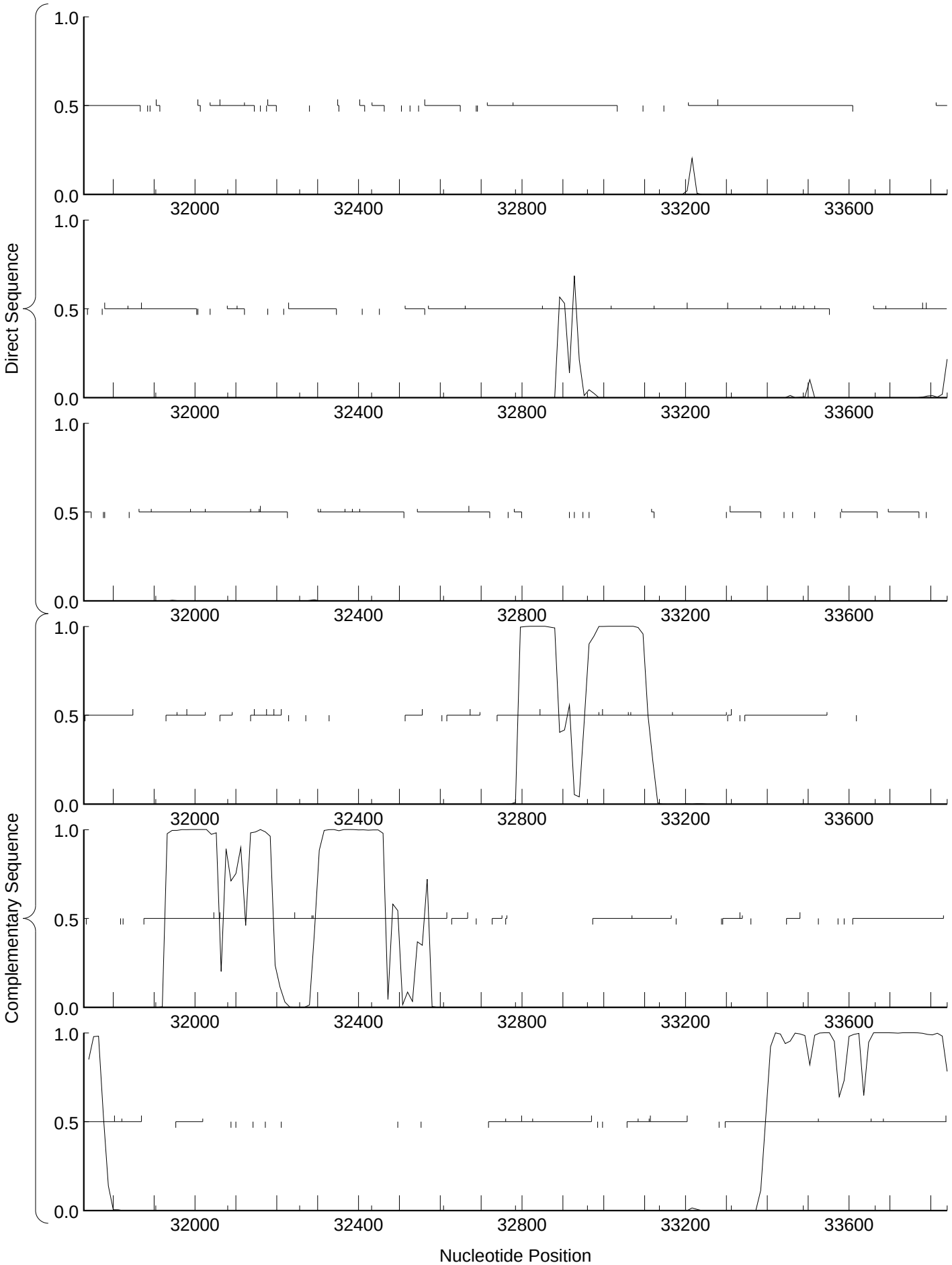
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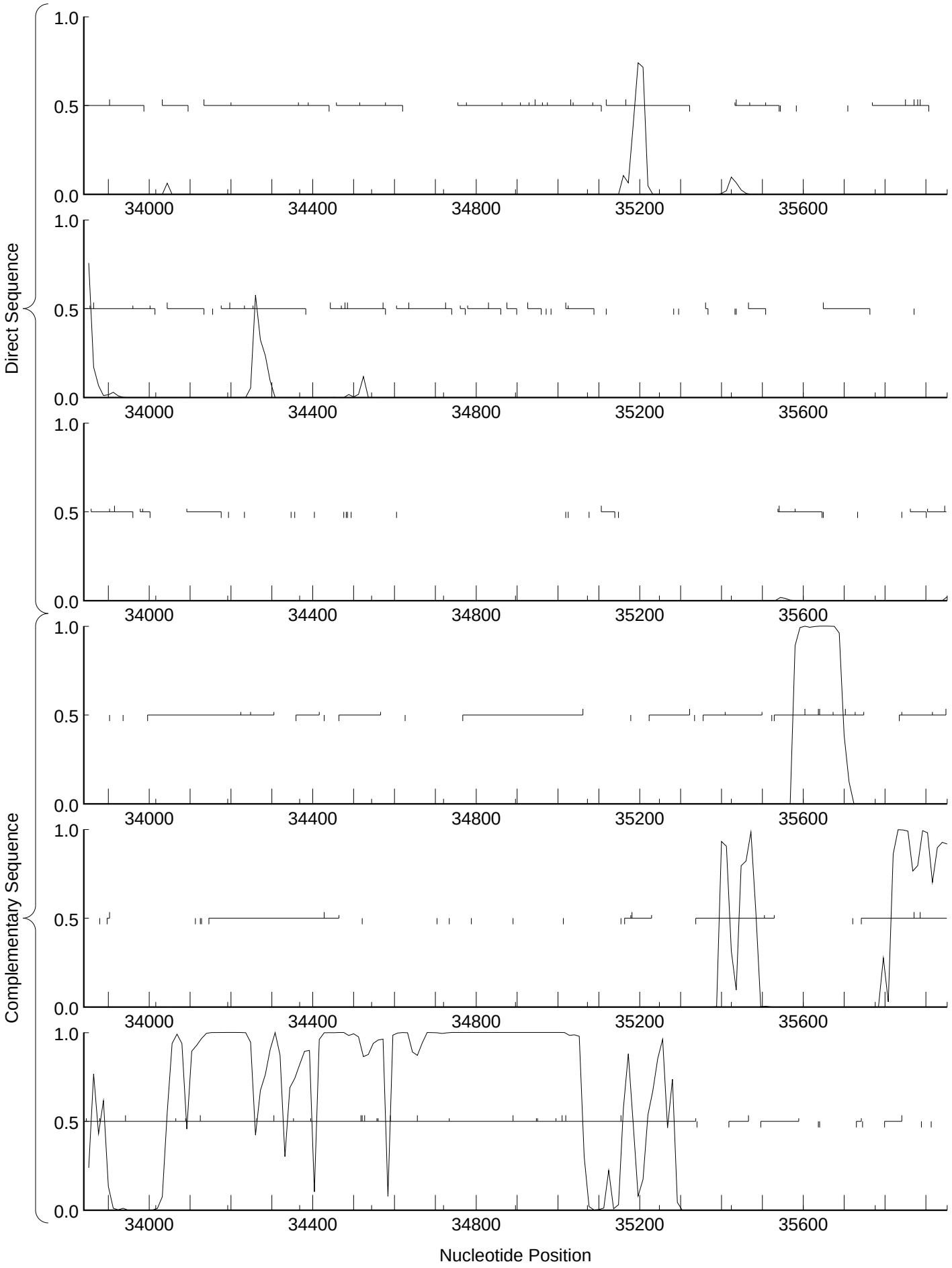
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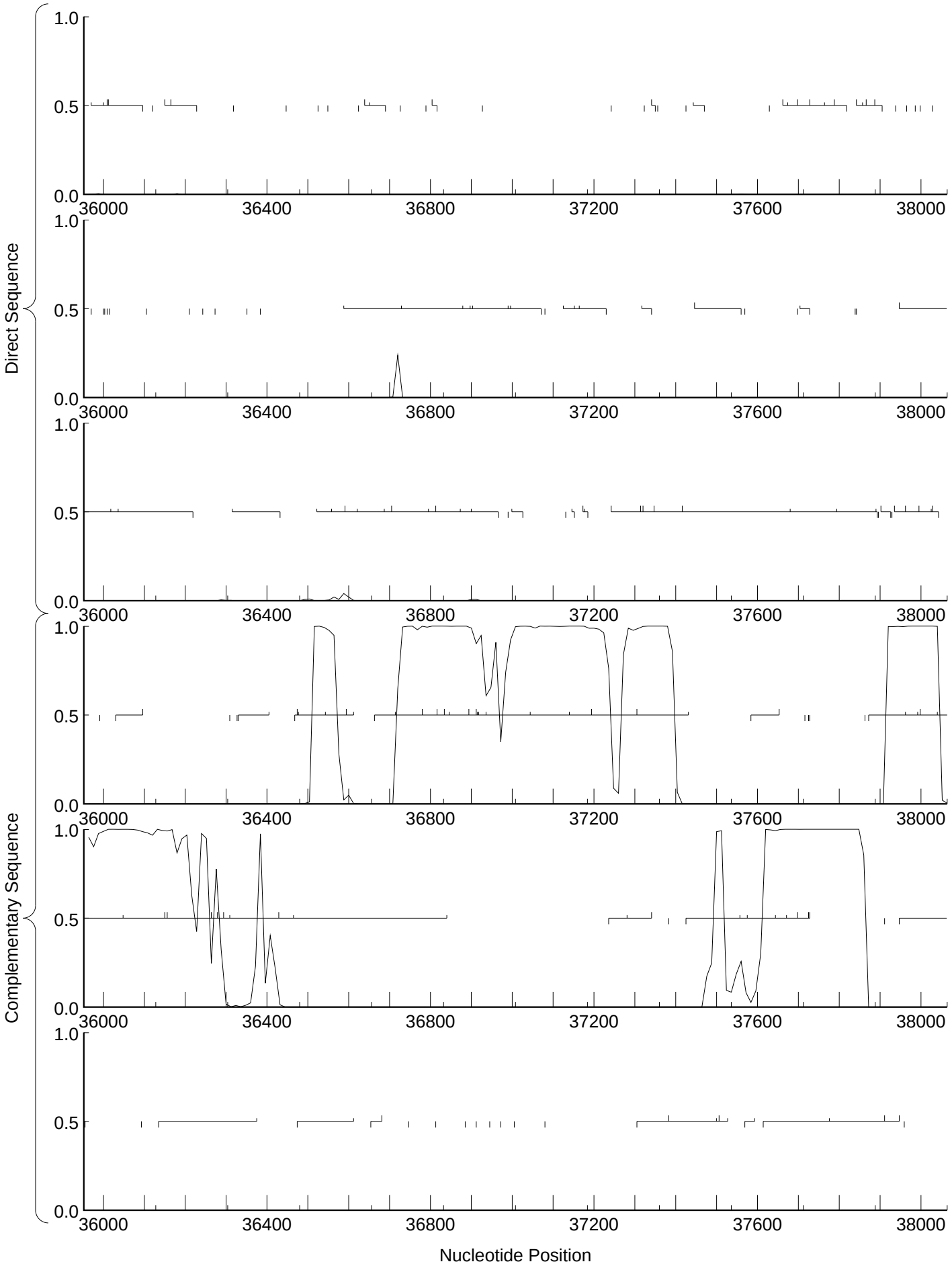
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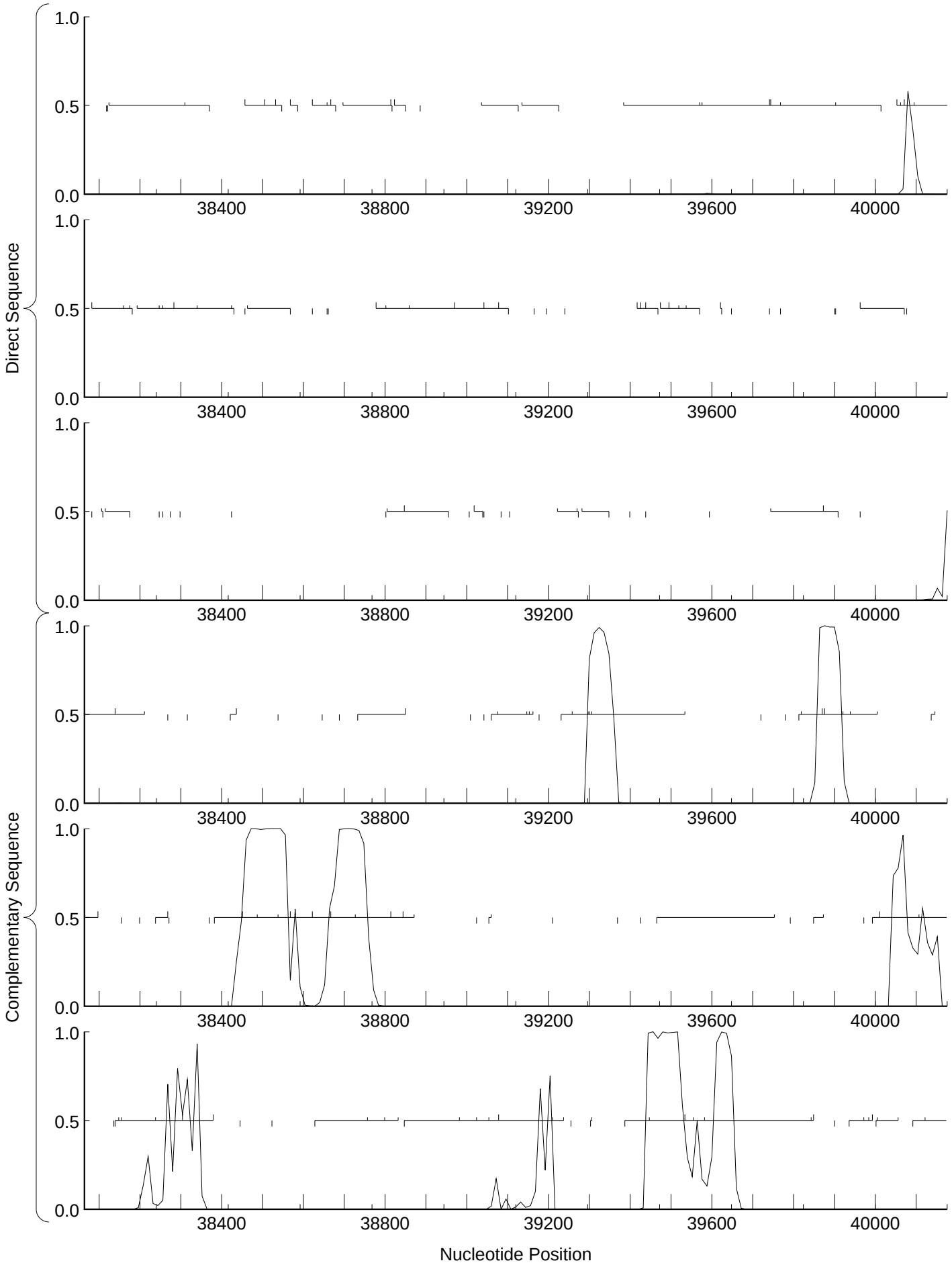
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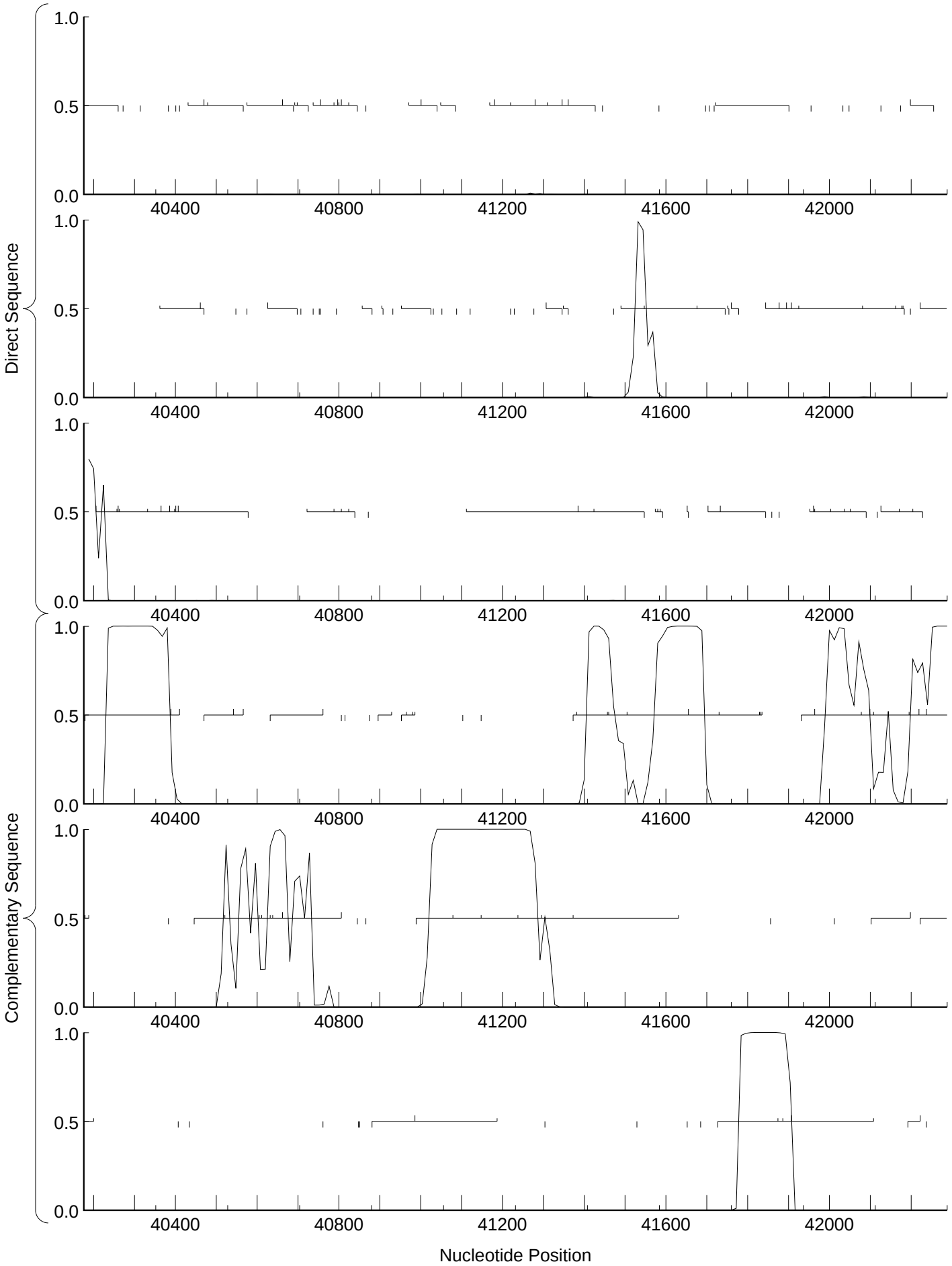
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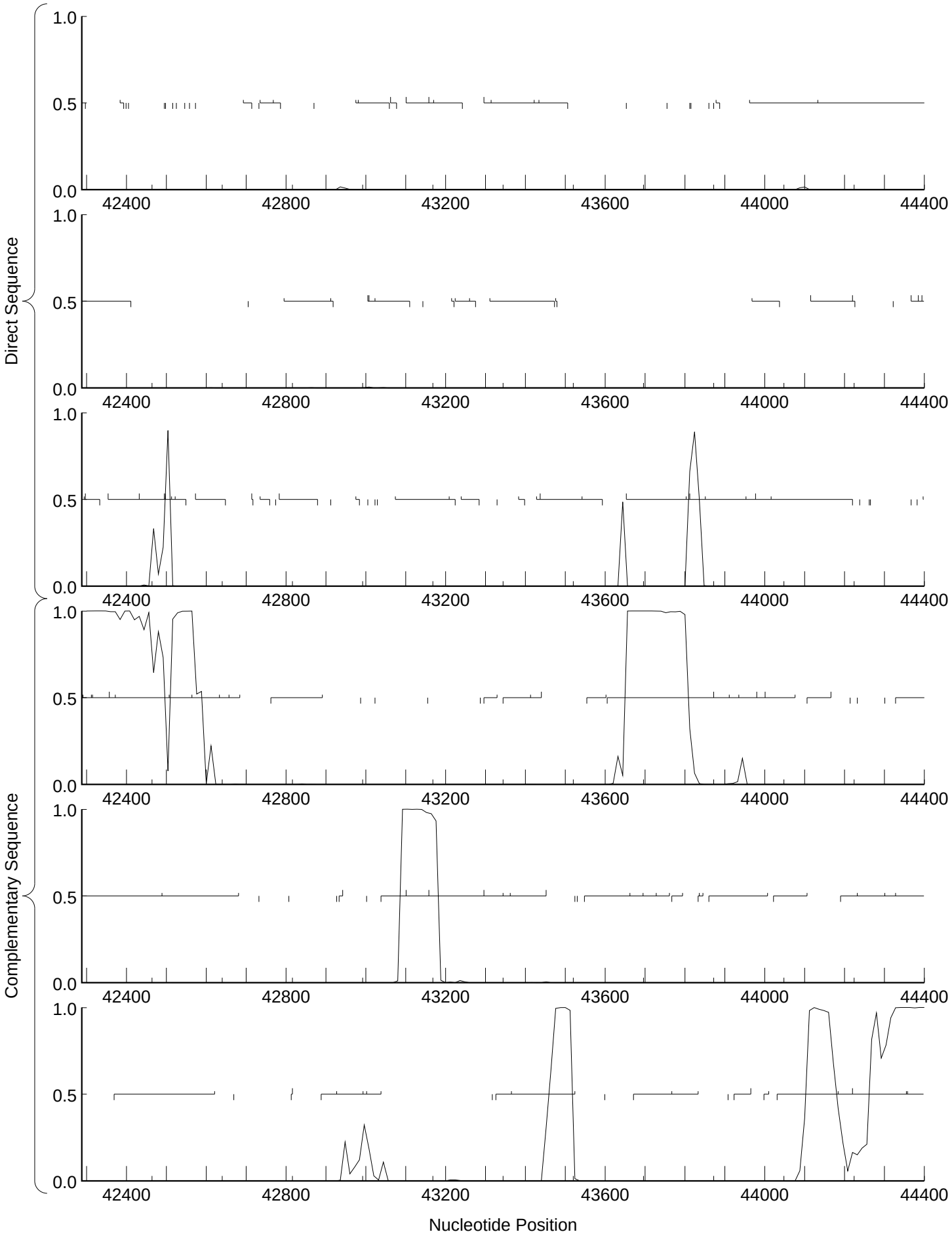
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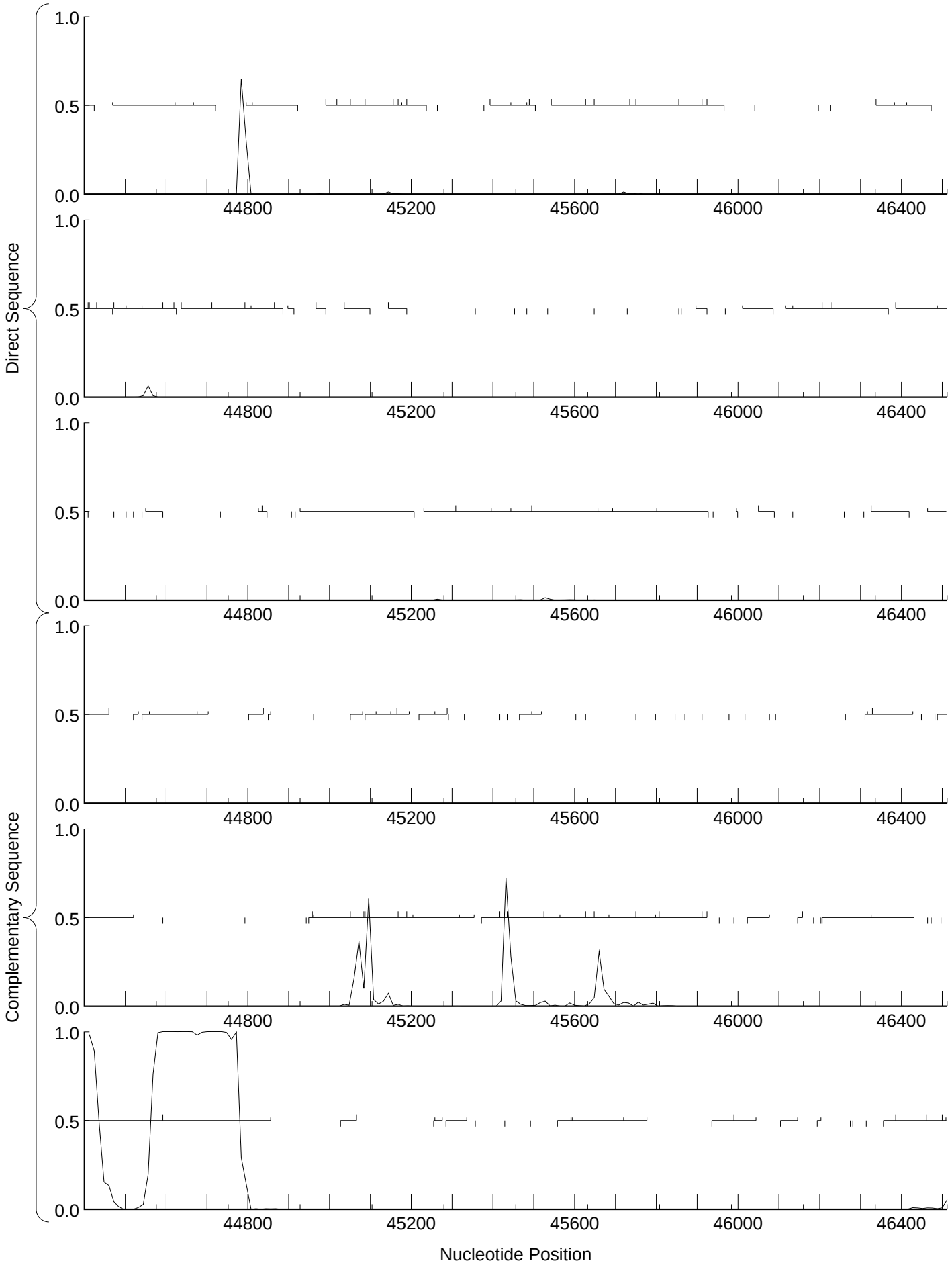
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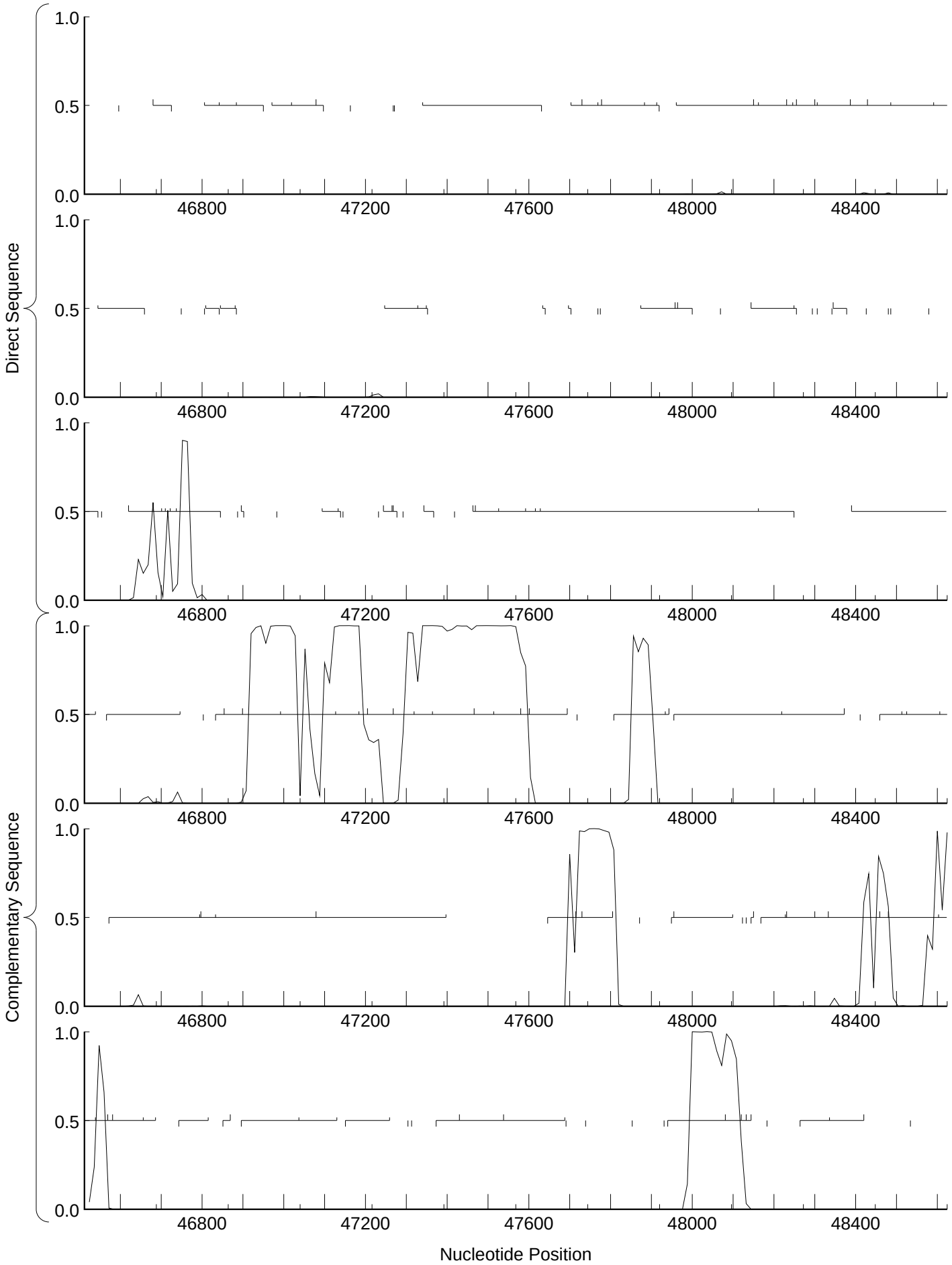
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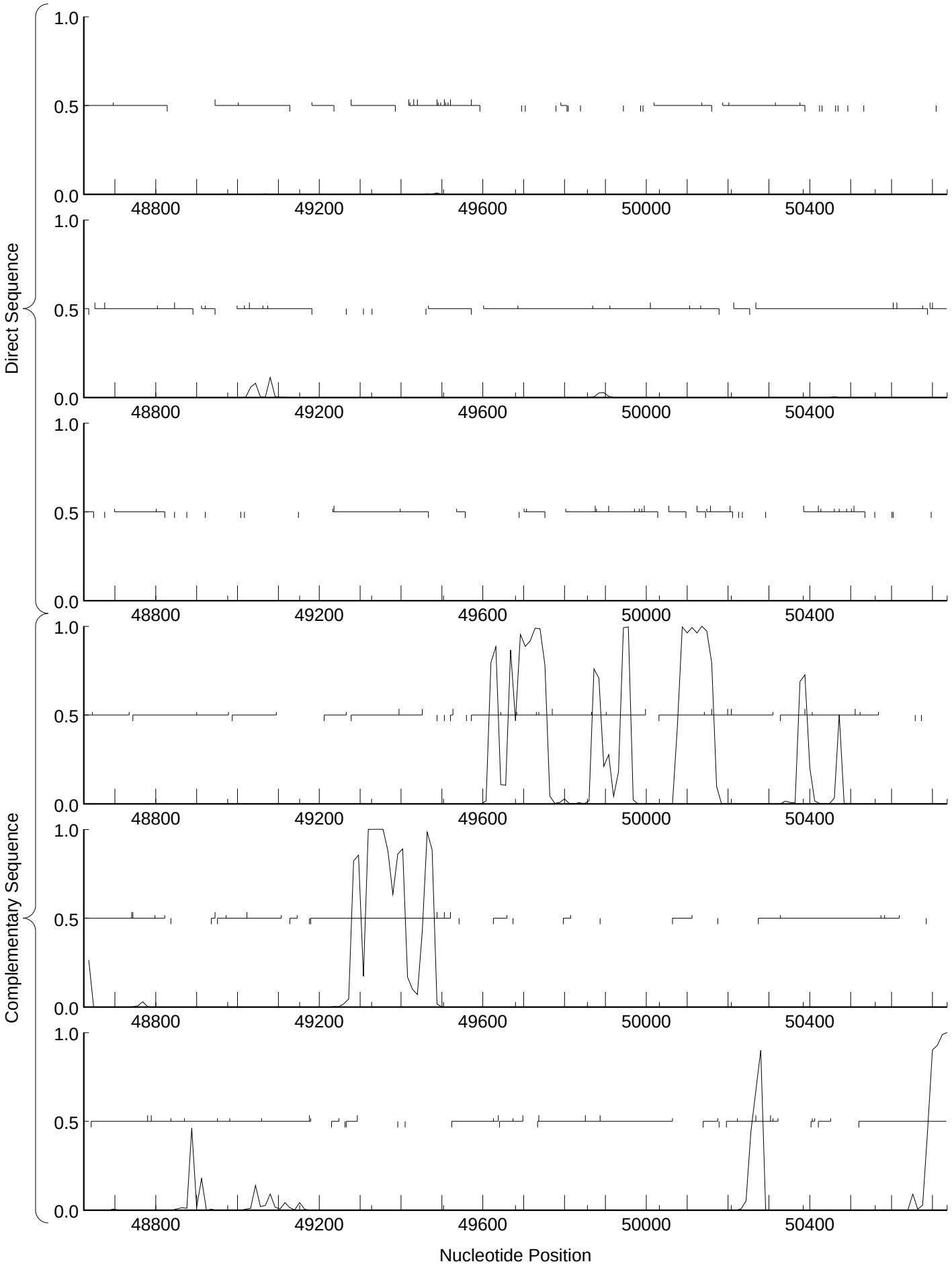
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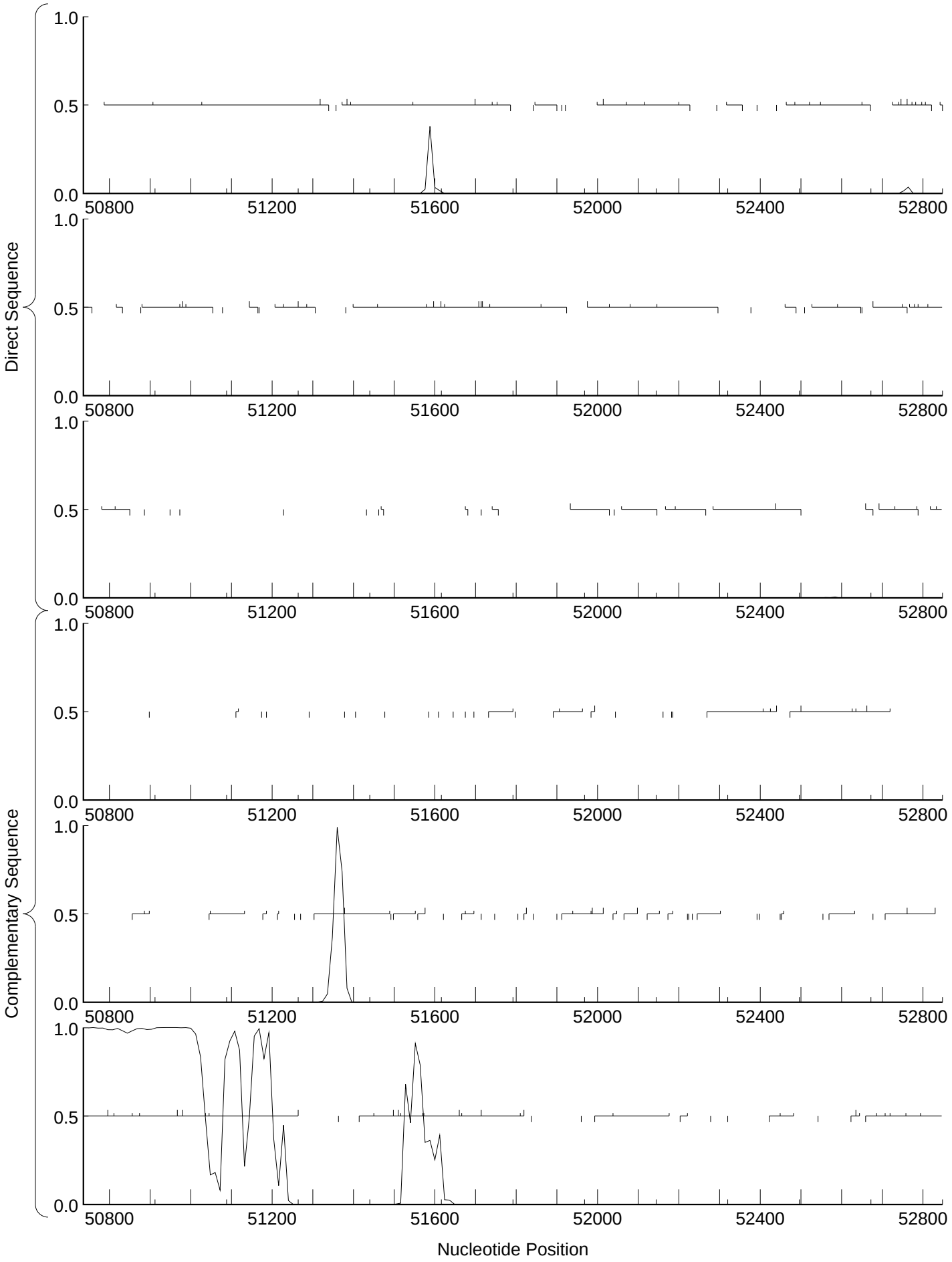
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e complete sequence, 52974 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 25/27



e complete sequence, 52974 bp including 10-base 3' overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 26/27



e complete sequence, 52974 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 27/27

