GeneMark

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

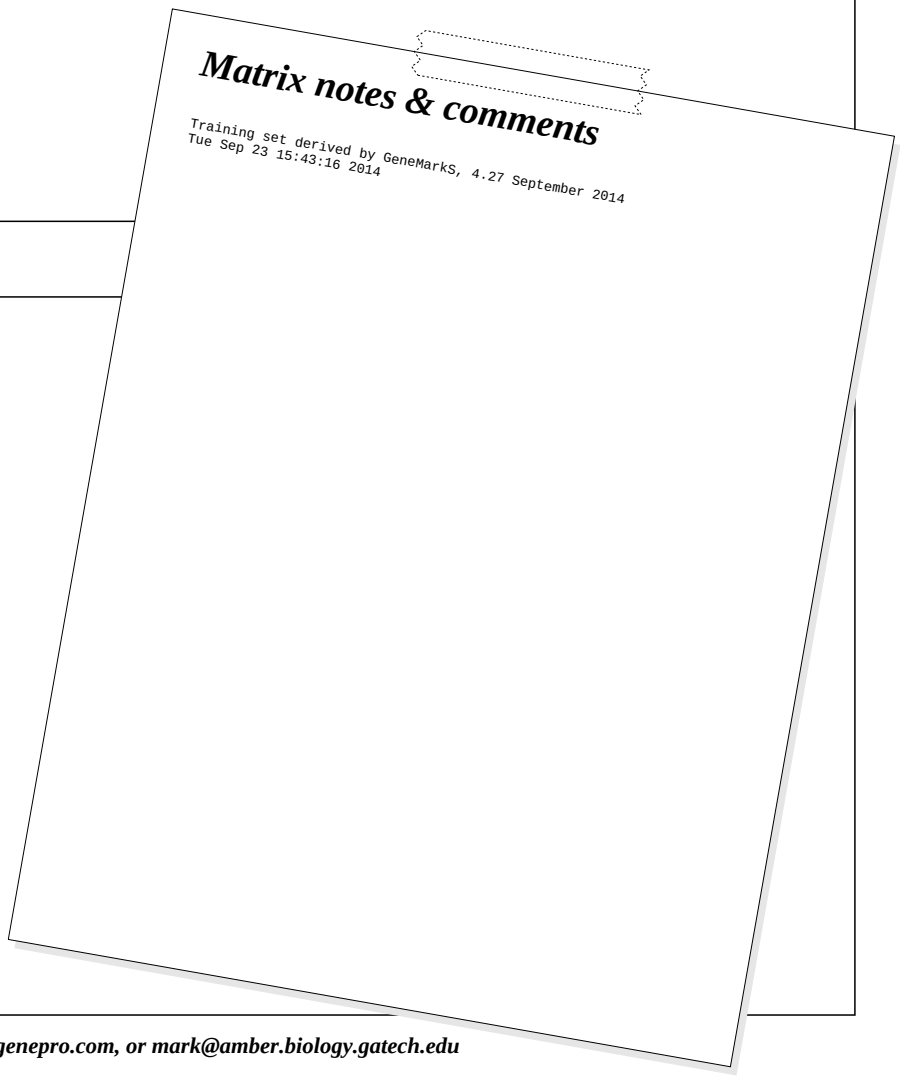
PROGRAM INFORMATION

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Sequence         :  Streptomyces phage GirlPower complete sequence, 55521 bp including 9-base 3’ overhang (CGCC
Analysis Date    :  3/29/24 at 11:32:40
Pages            :  28
Sequence Length  :  55521 bp
GC Content       :   59.10%

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

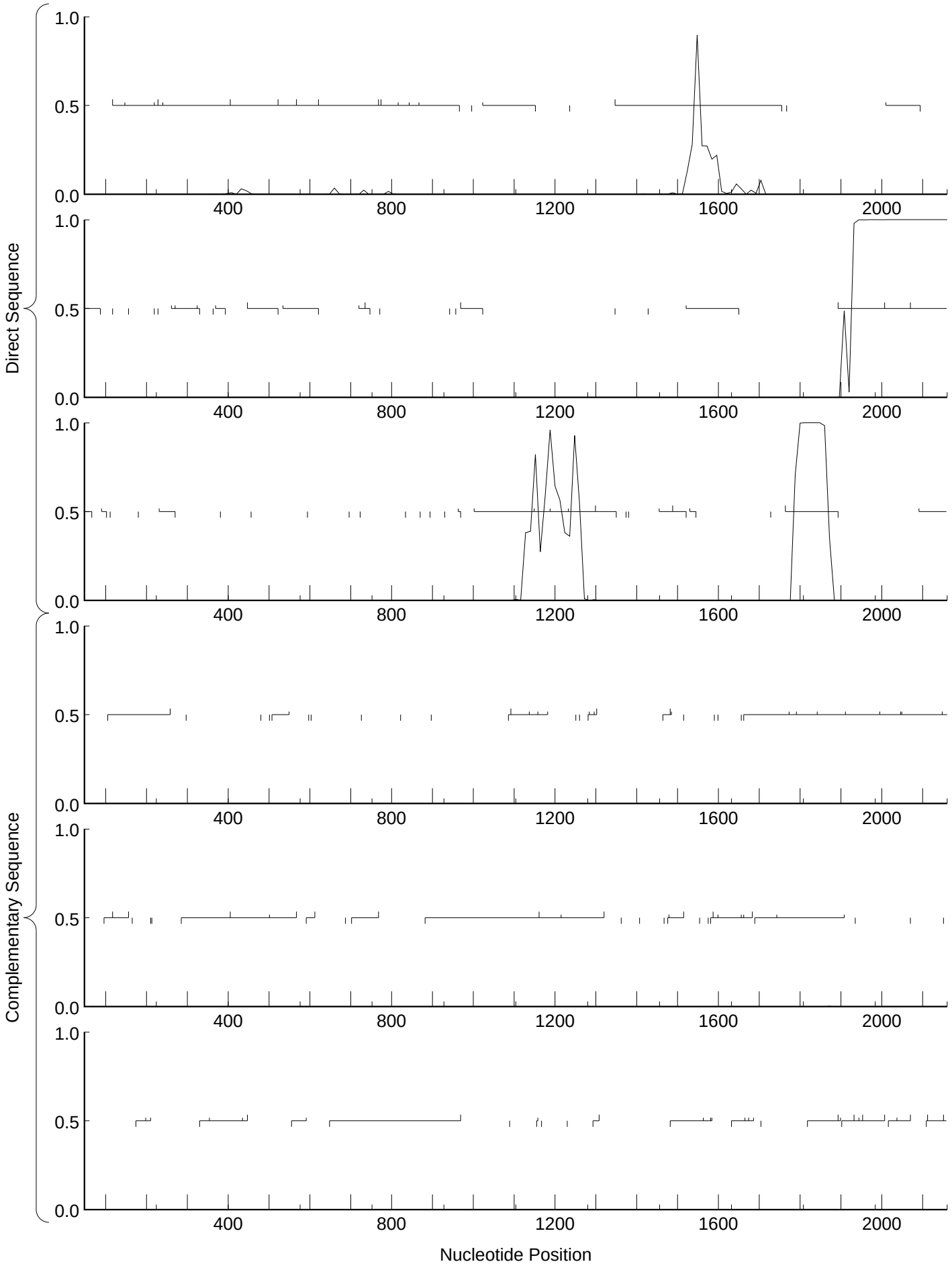
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Matrix     :  Streptomyces_scabiei_87_22
Author     :  -

Order      :  4
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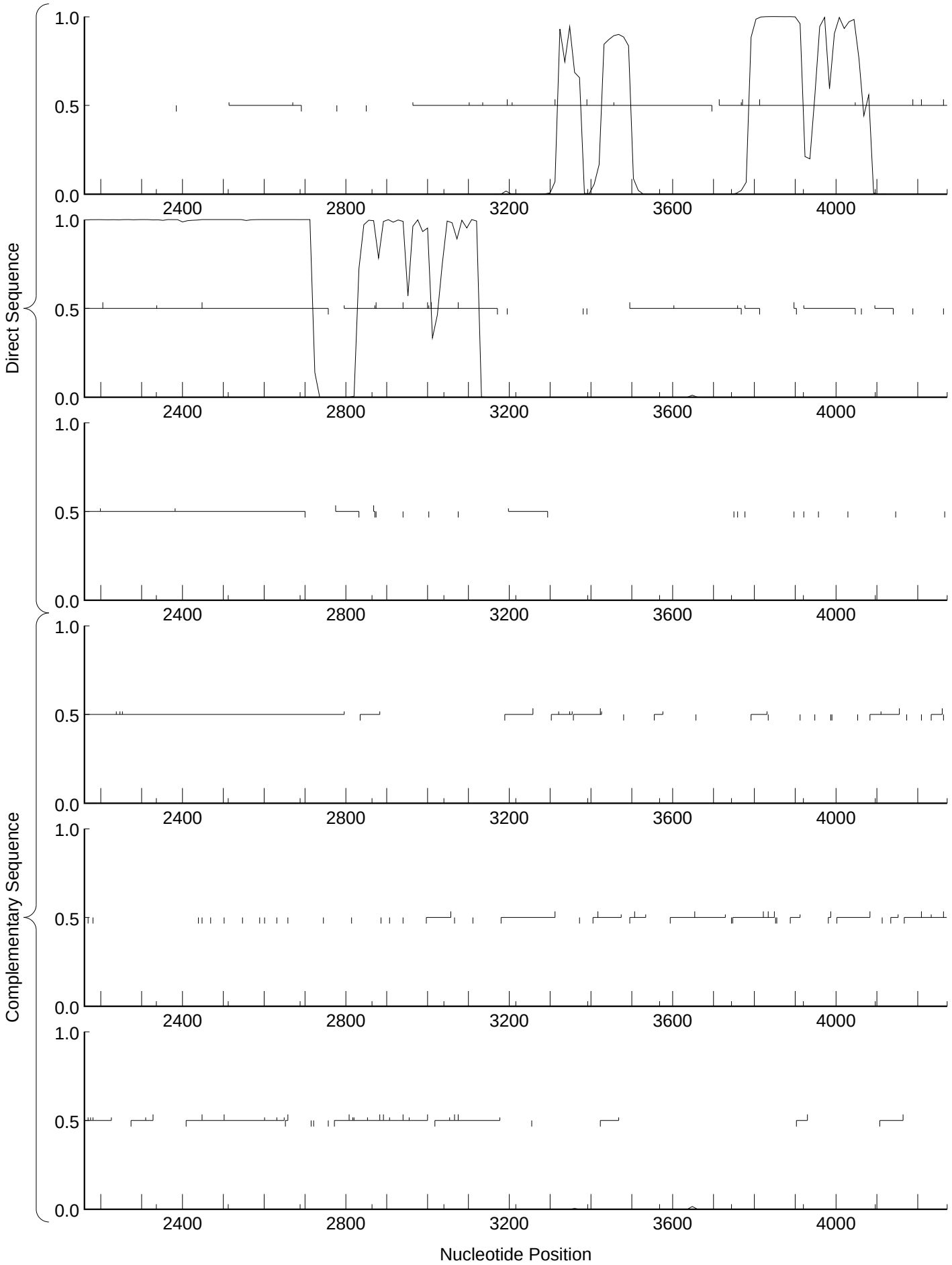

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

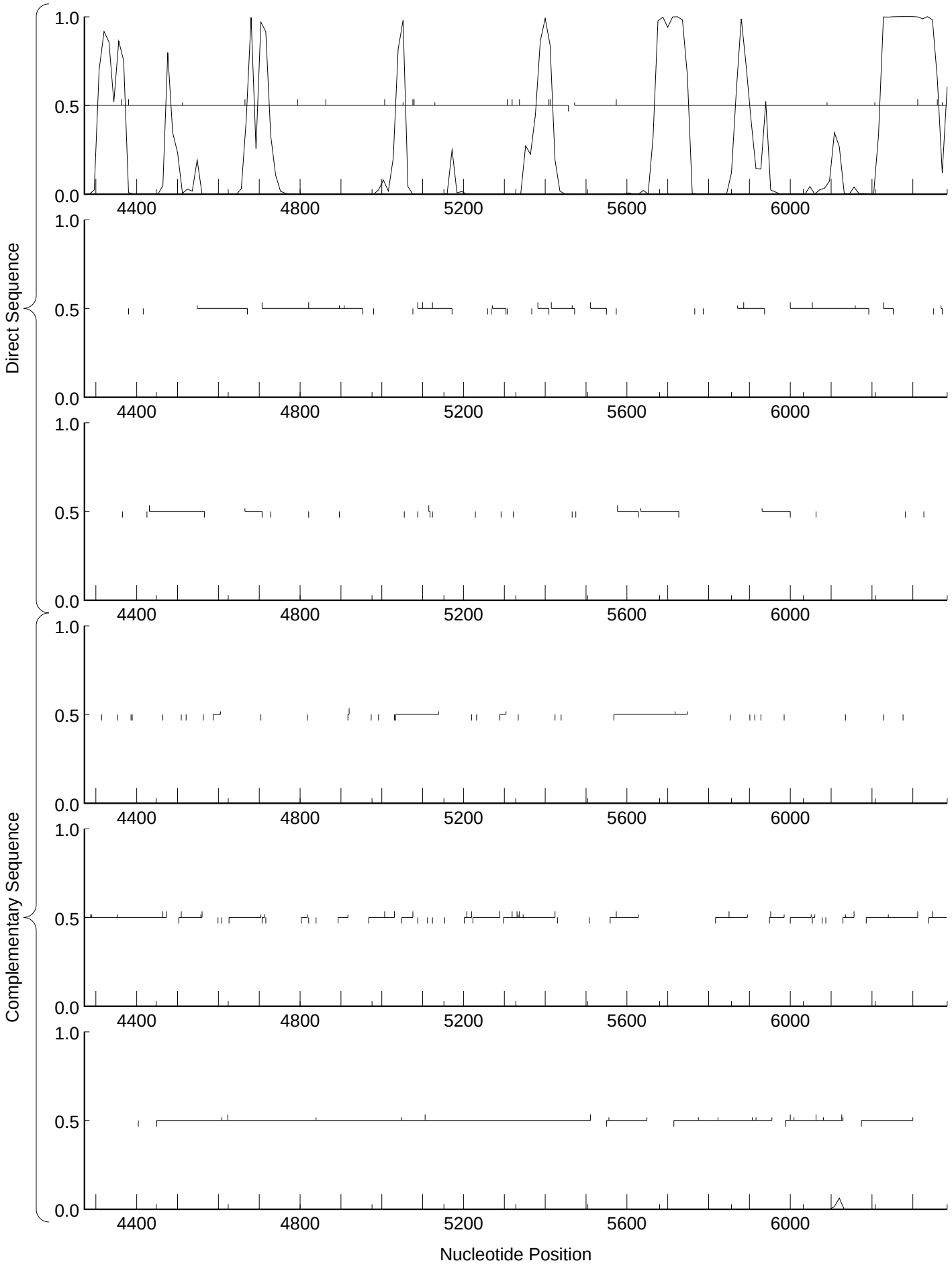
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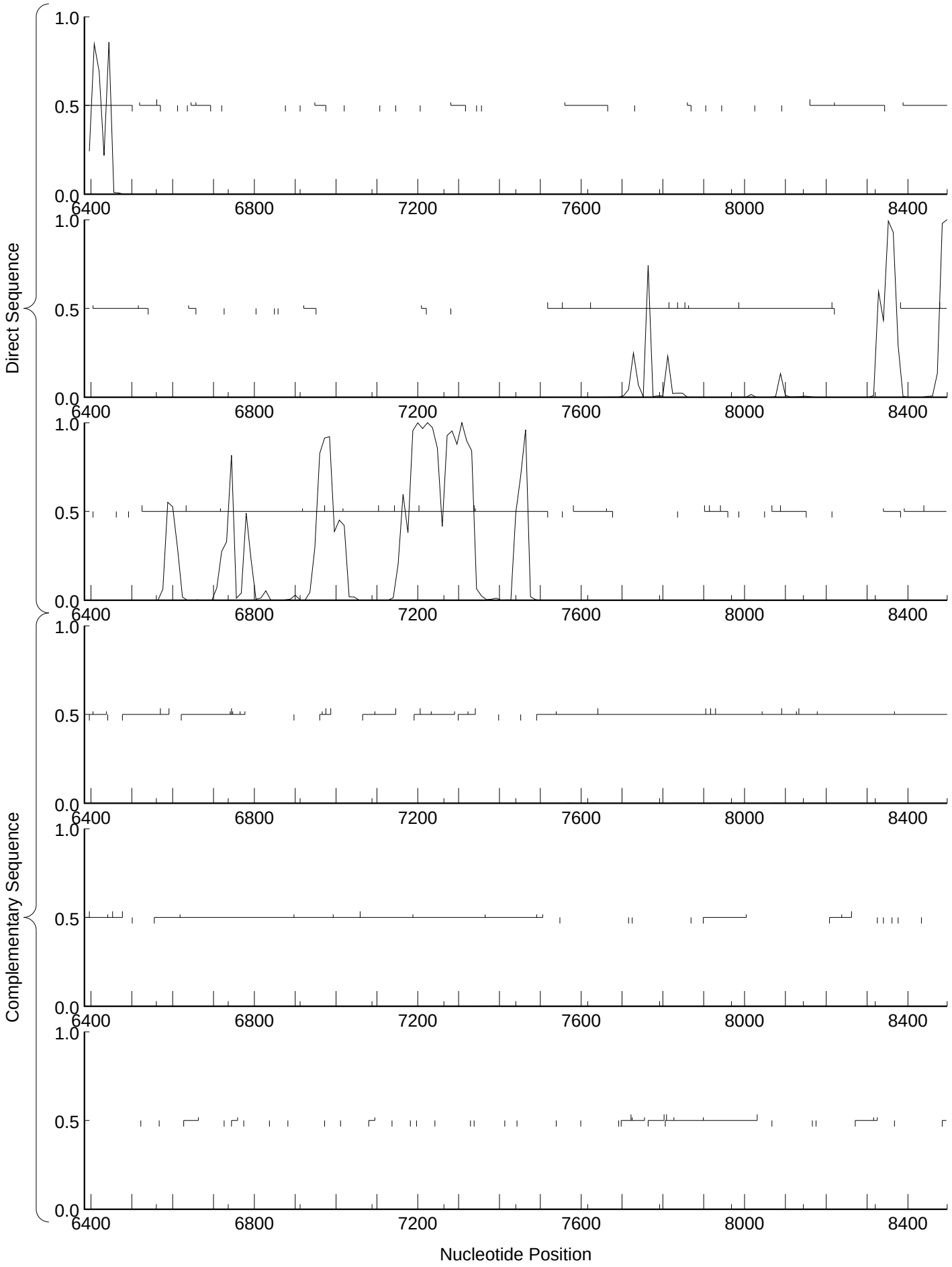
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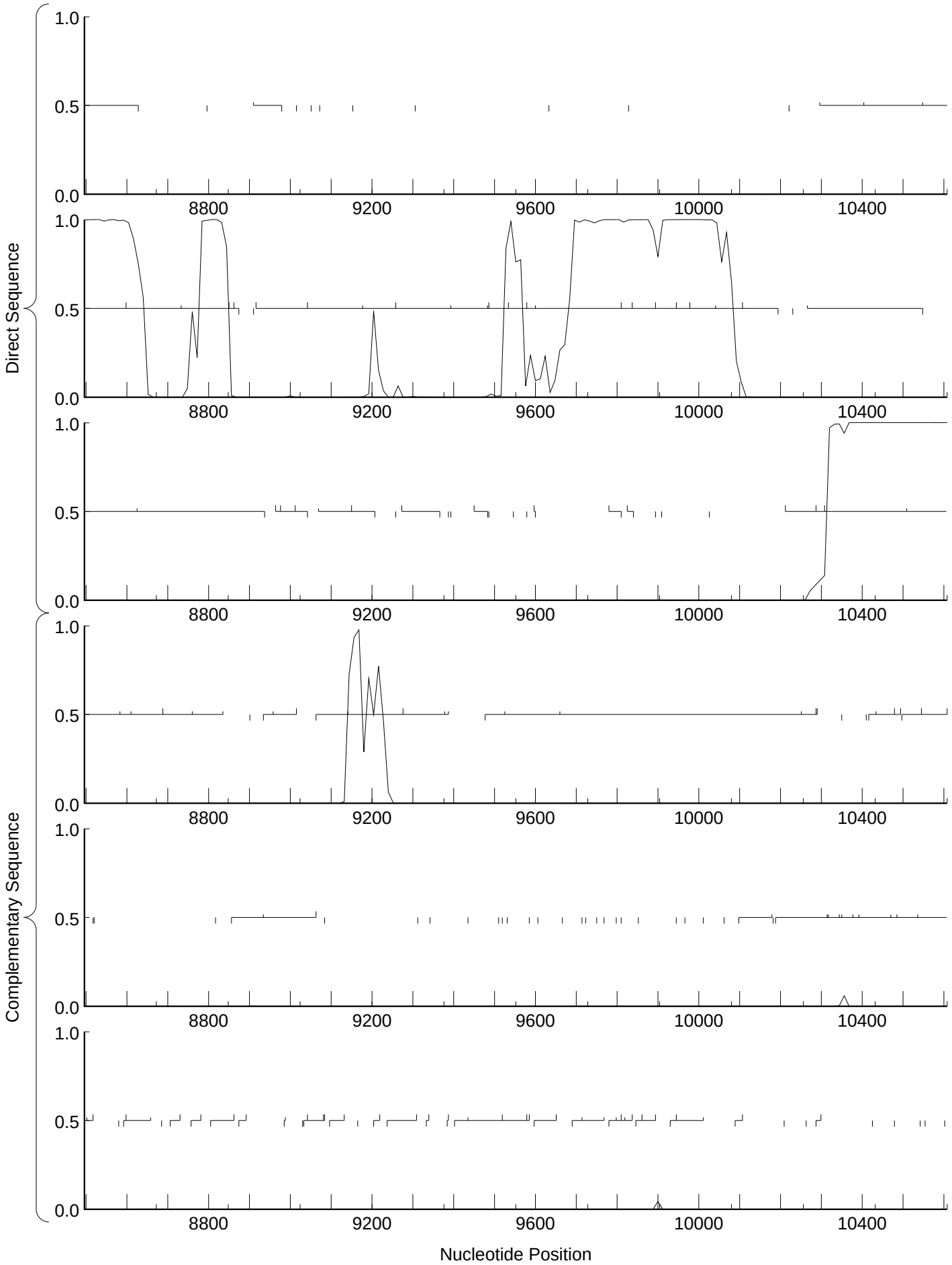
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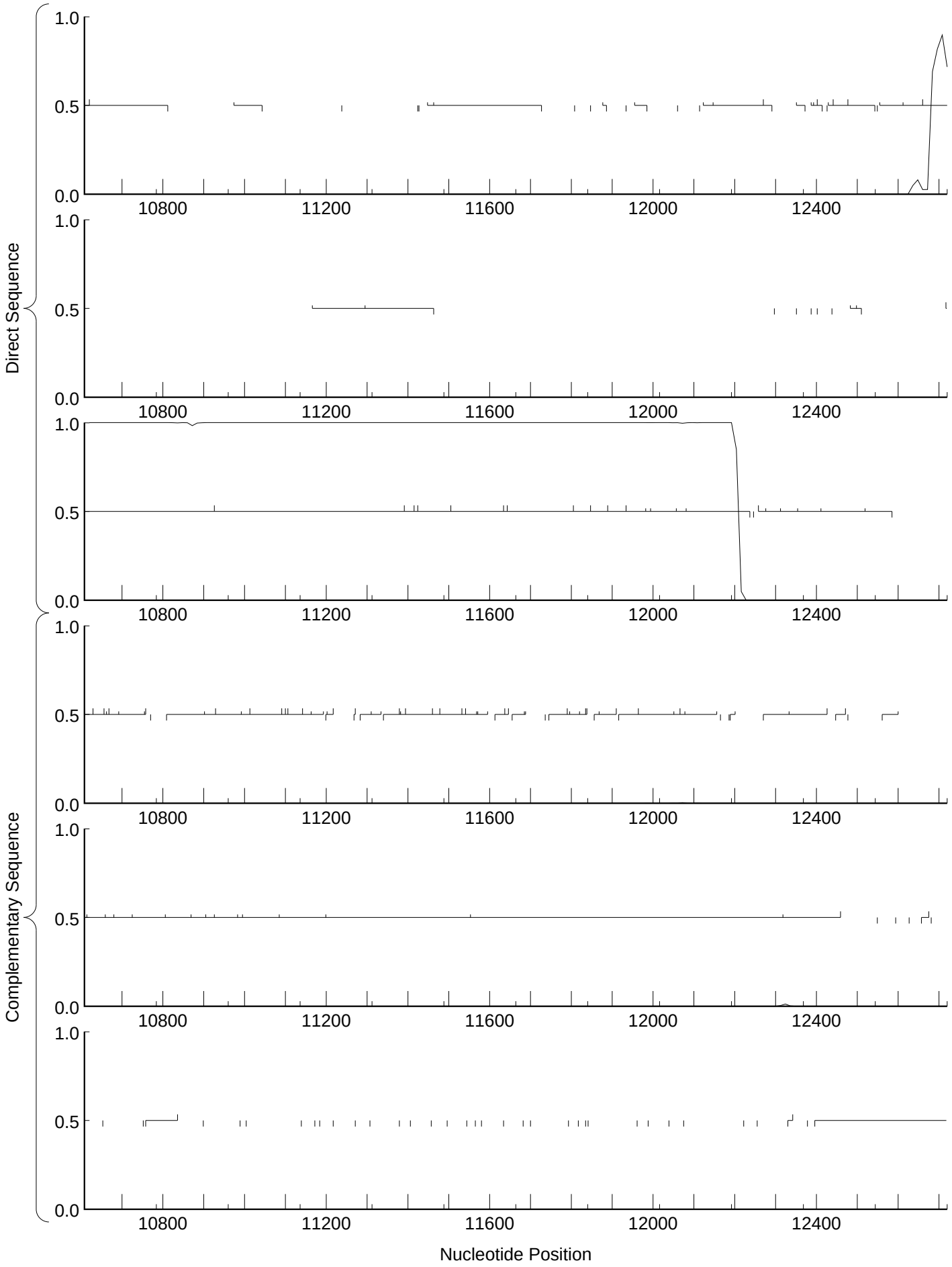
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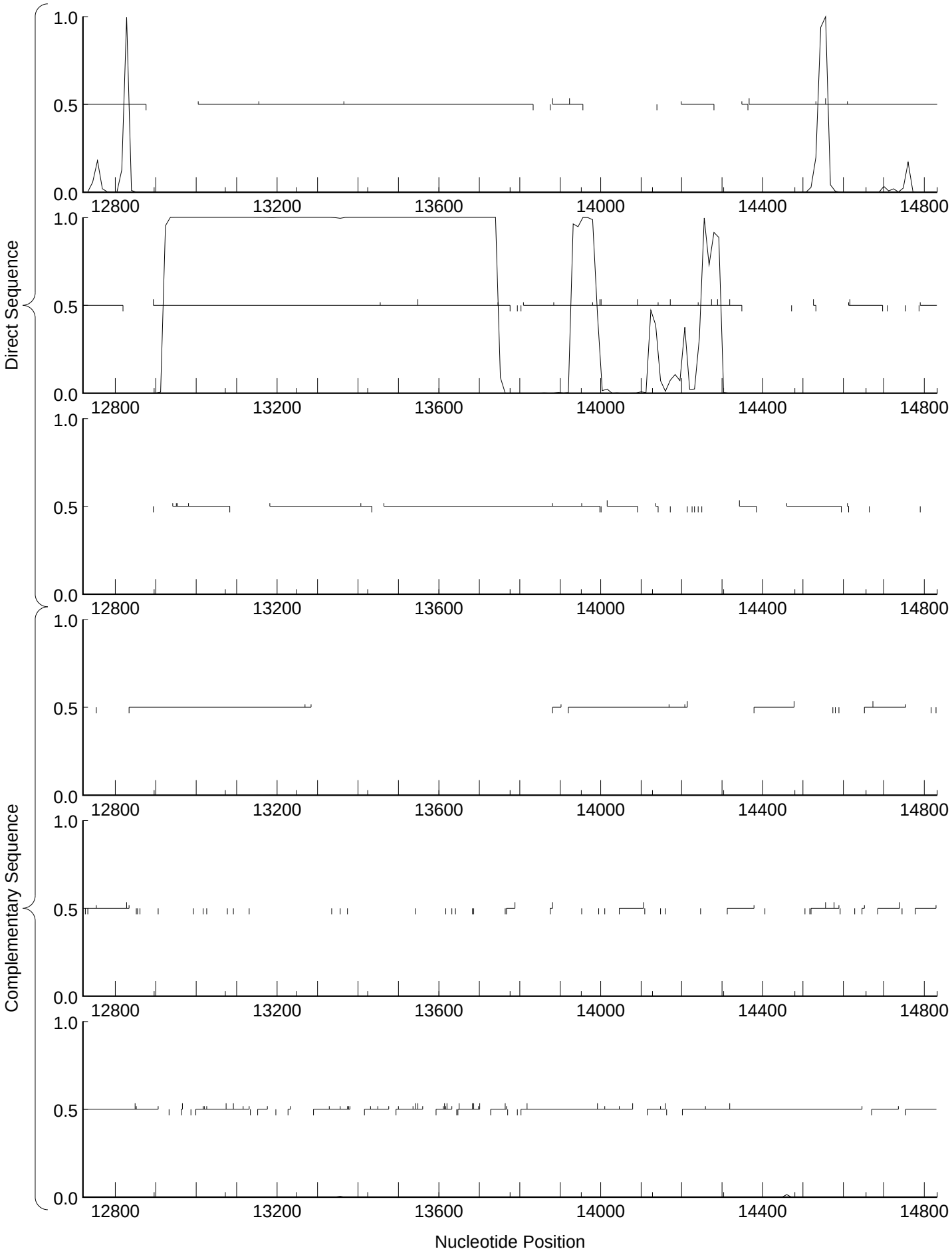
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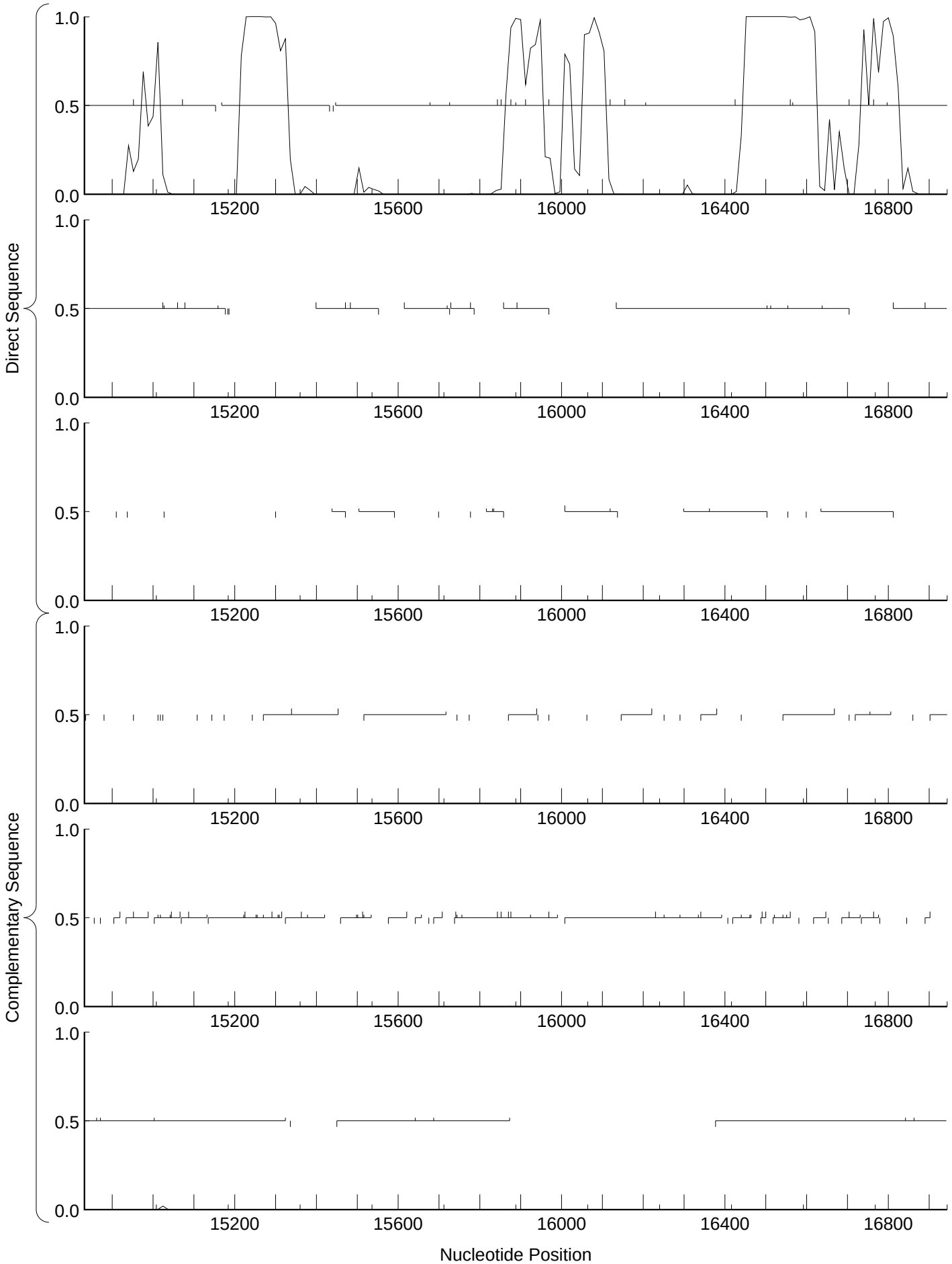
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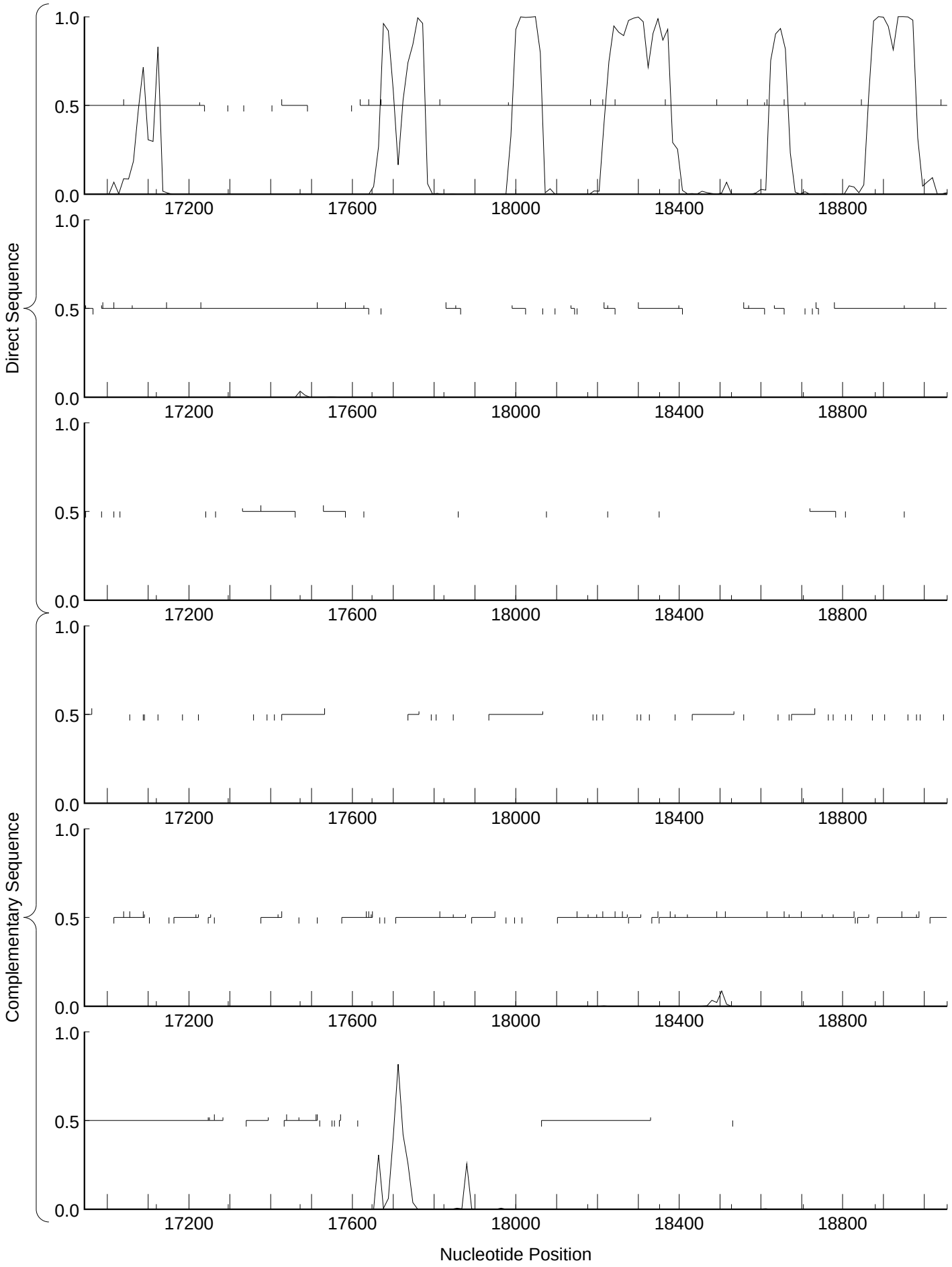
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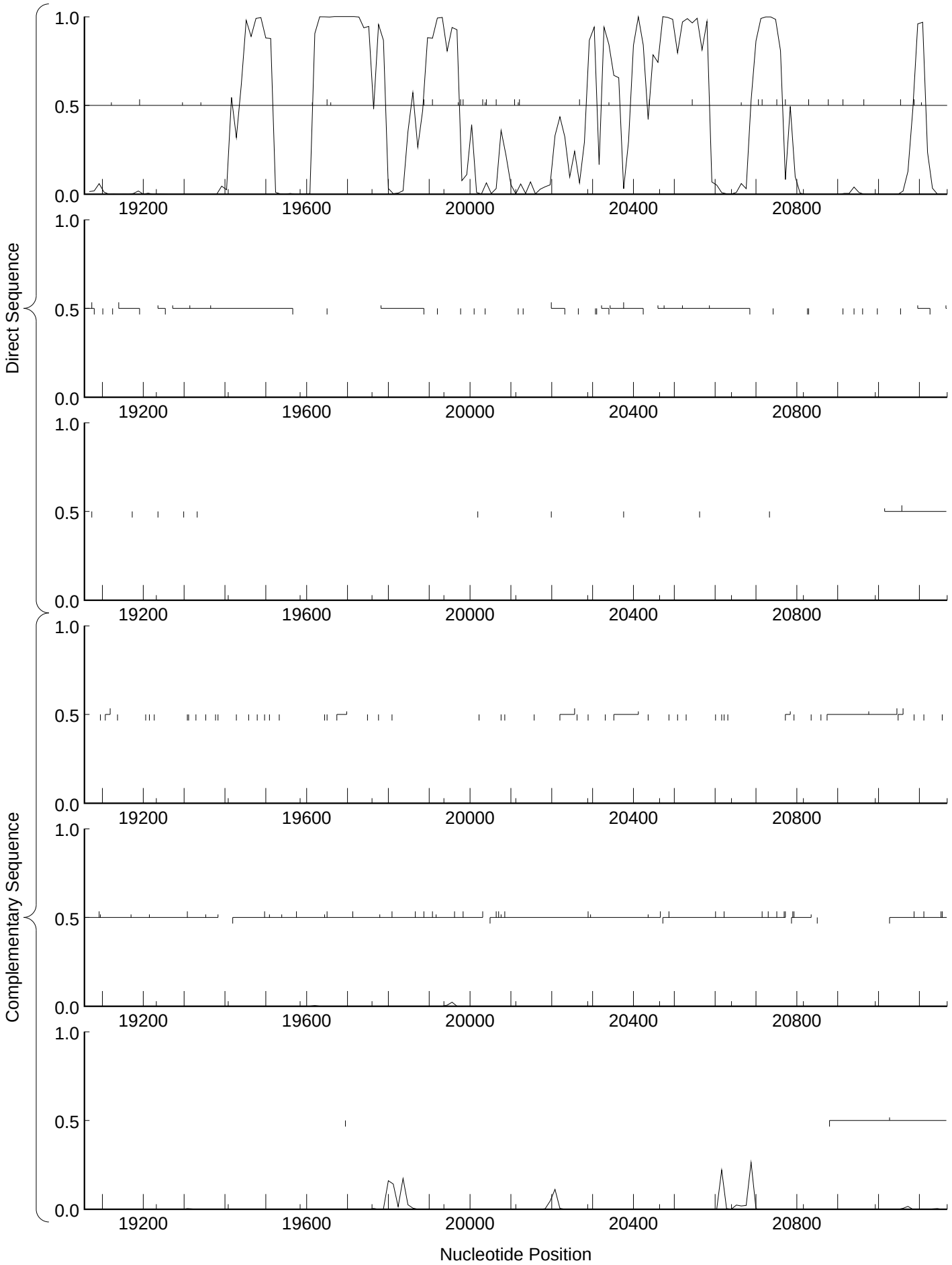
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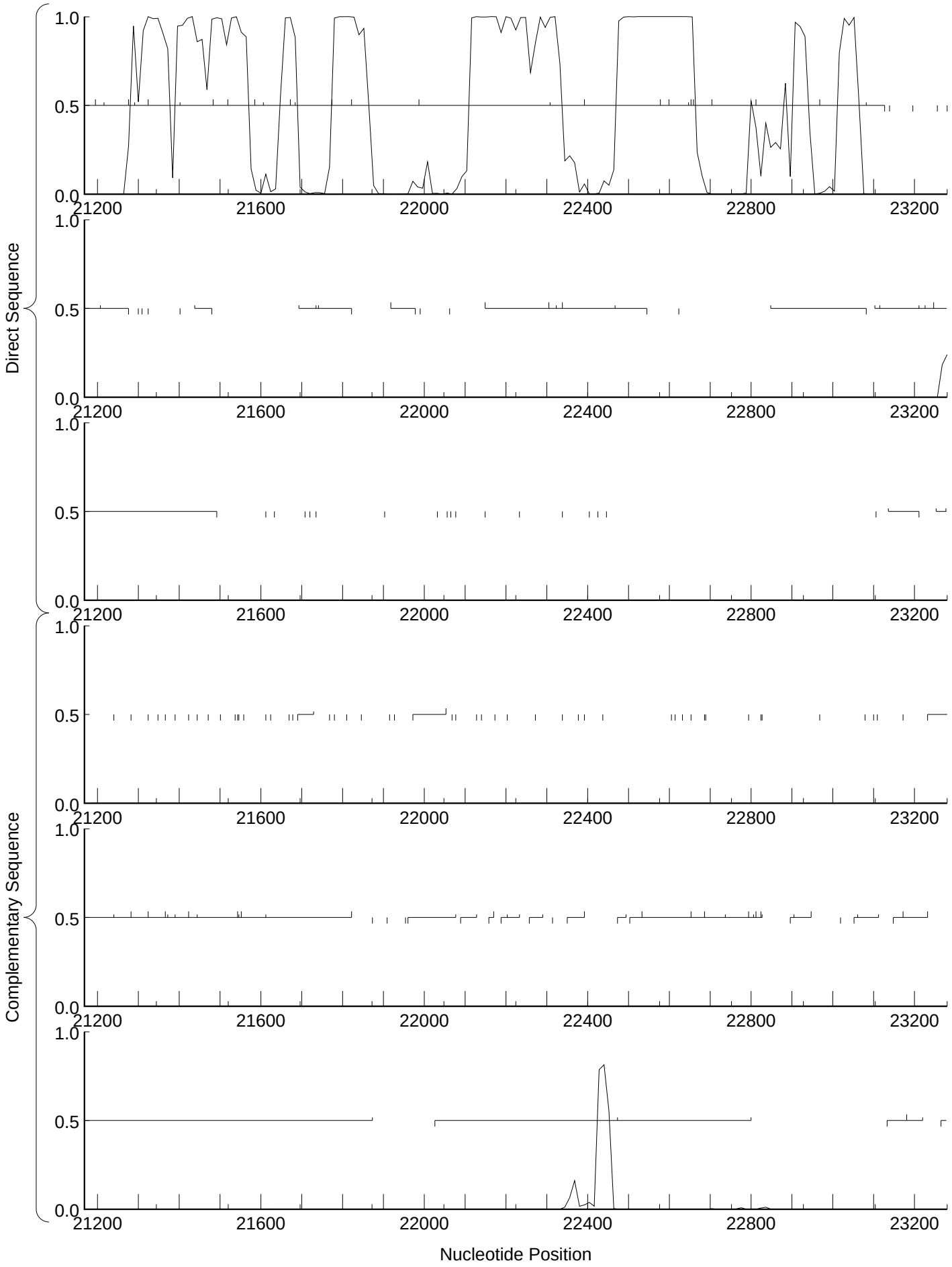
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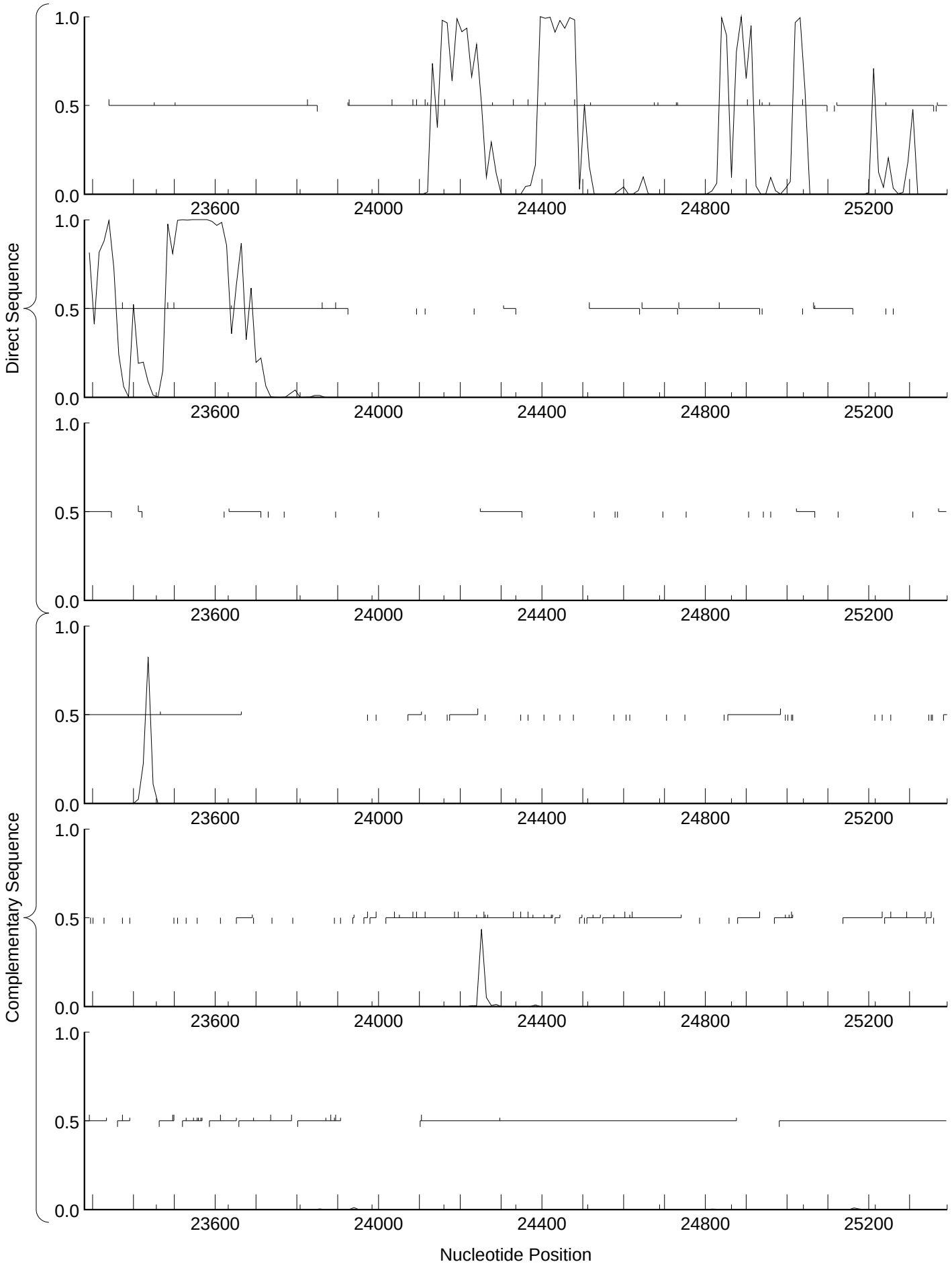
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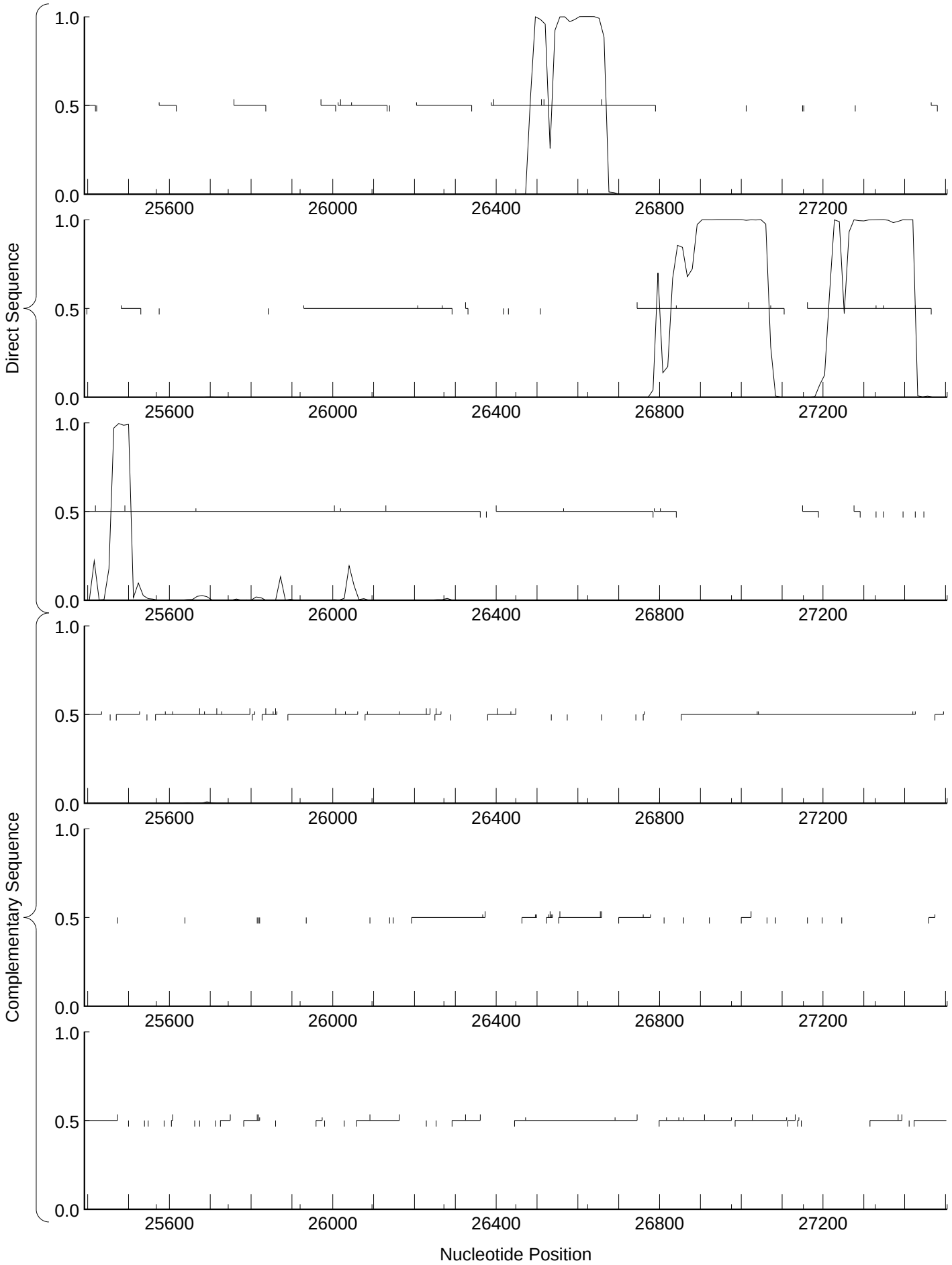
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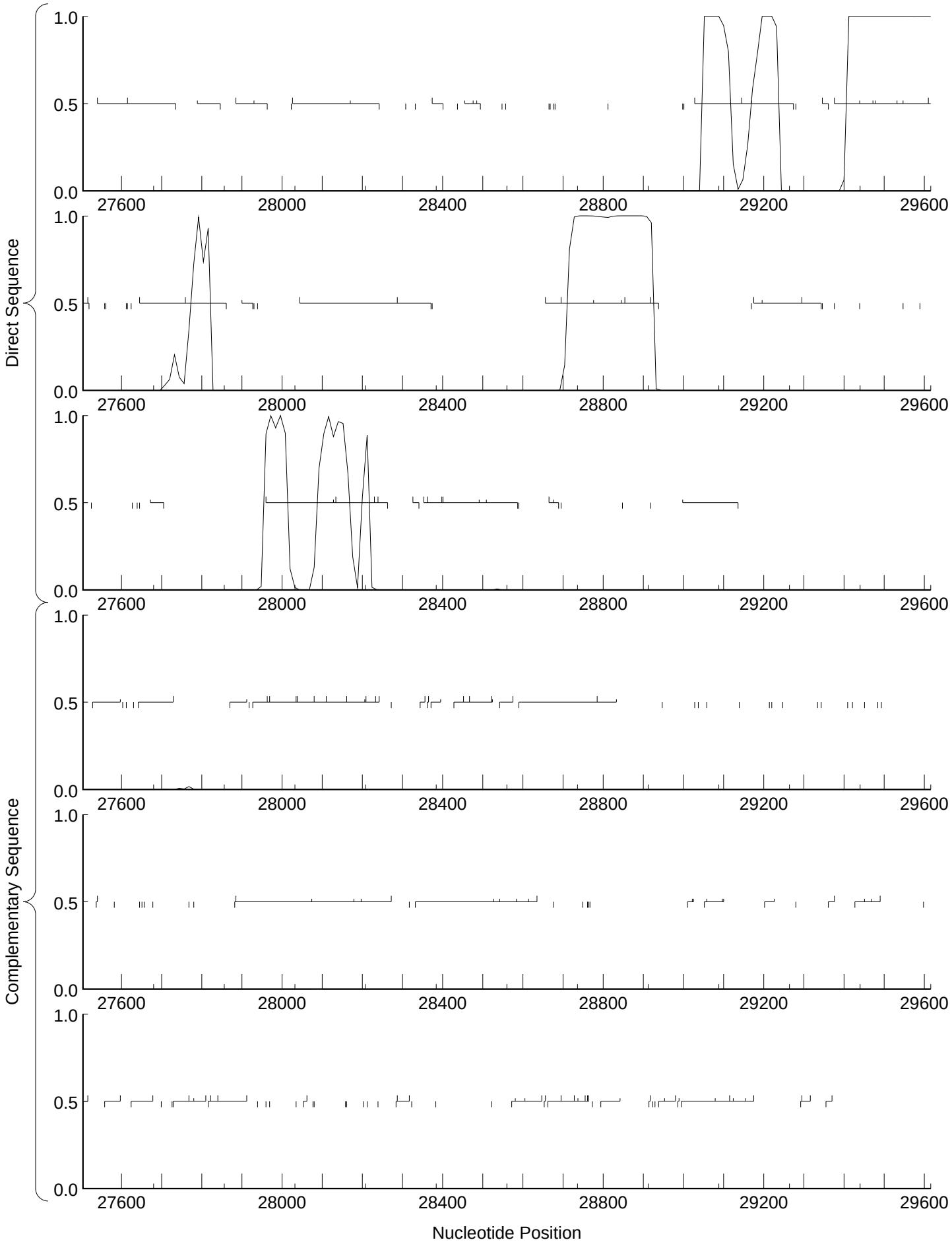
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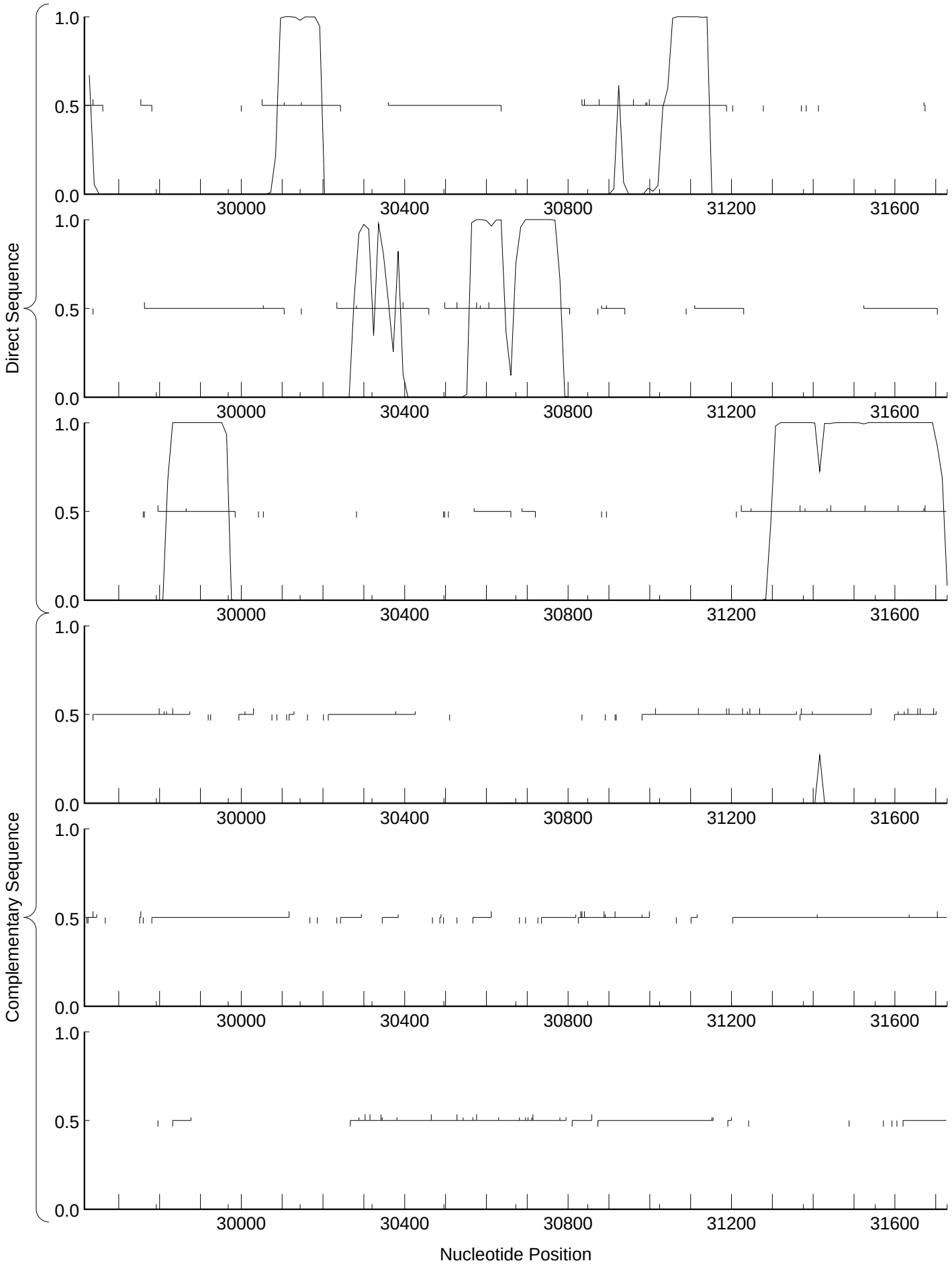
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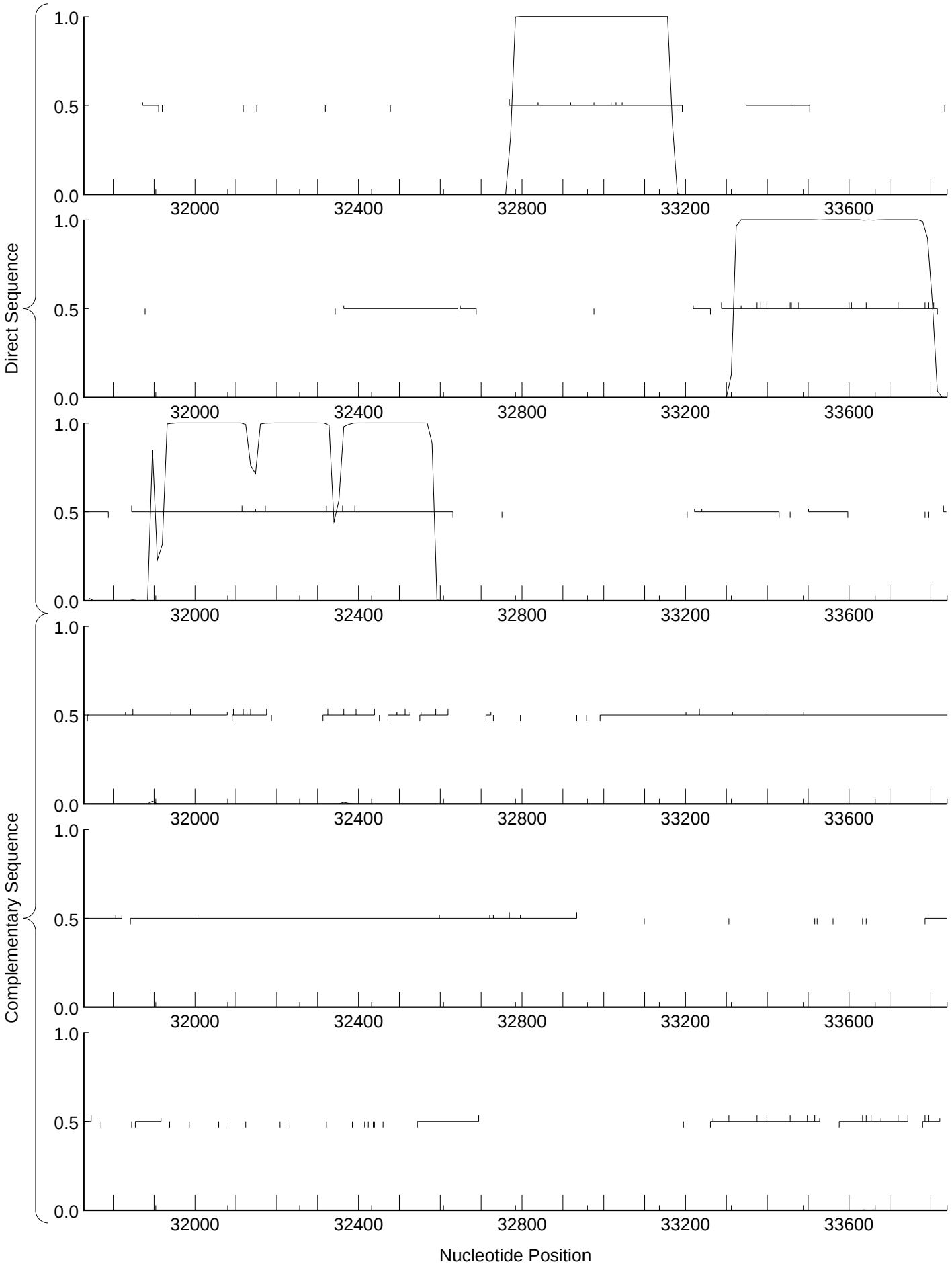
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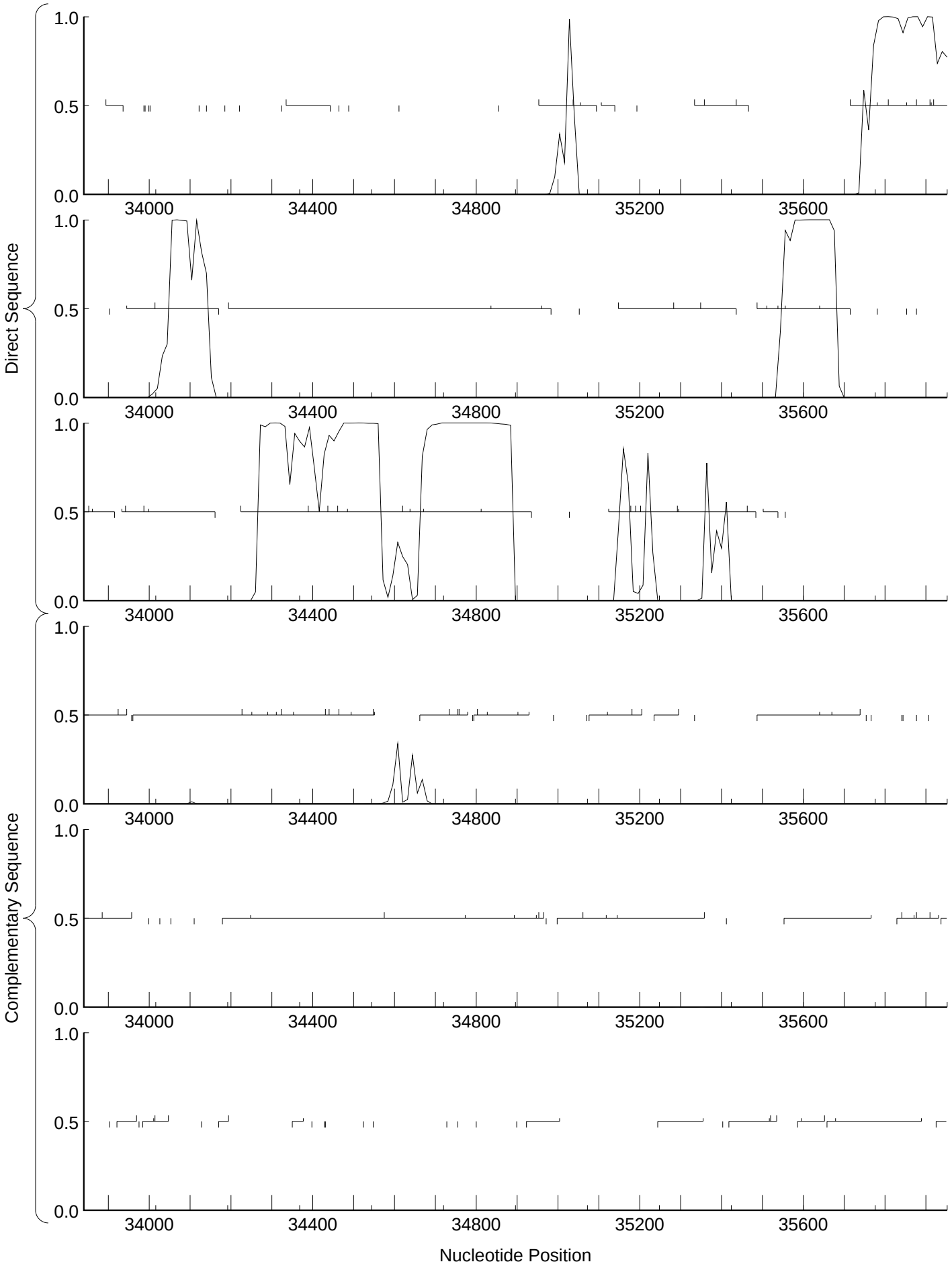
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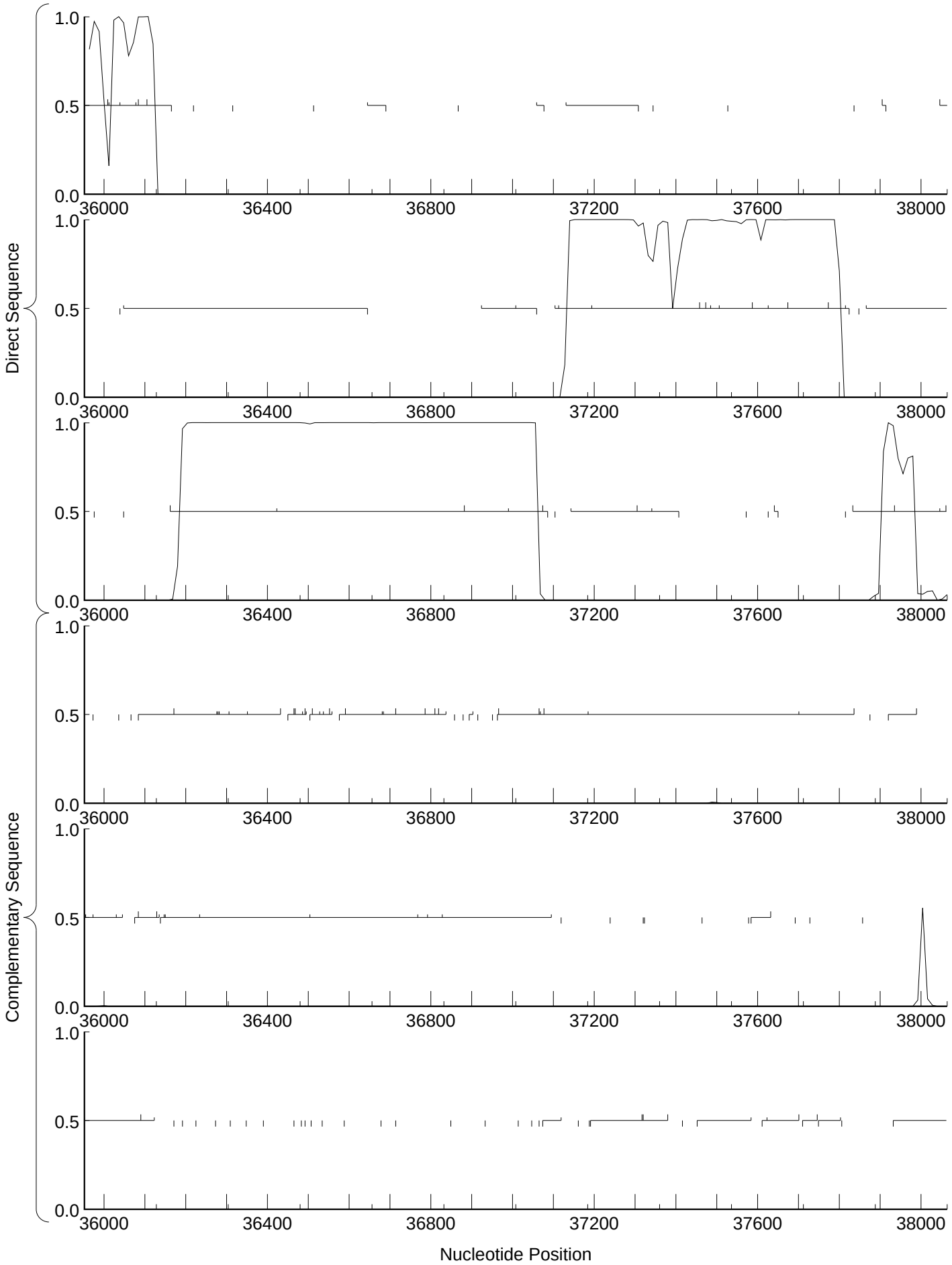
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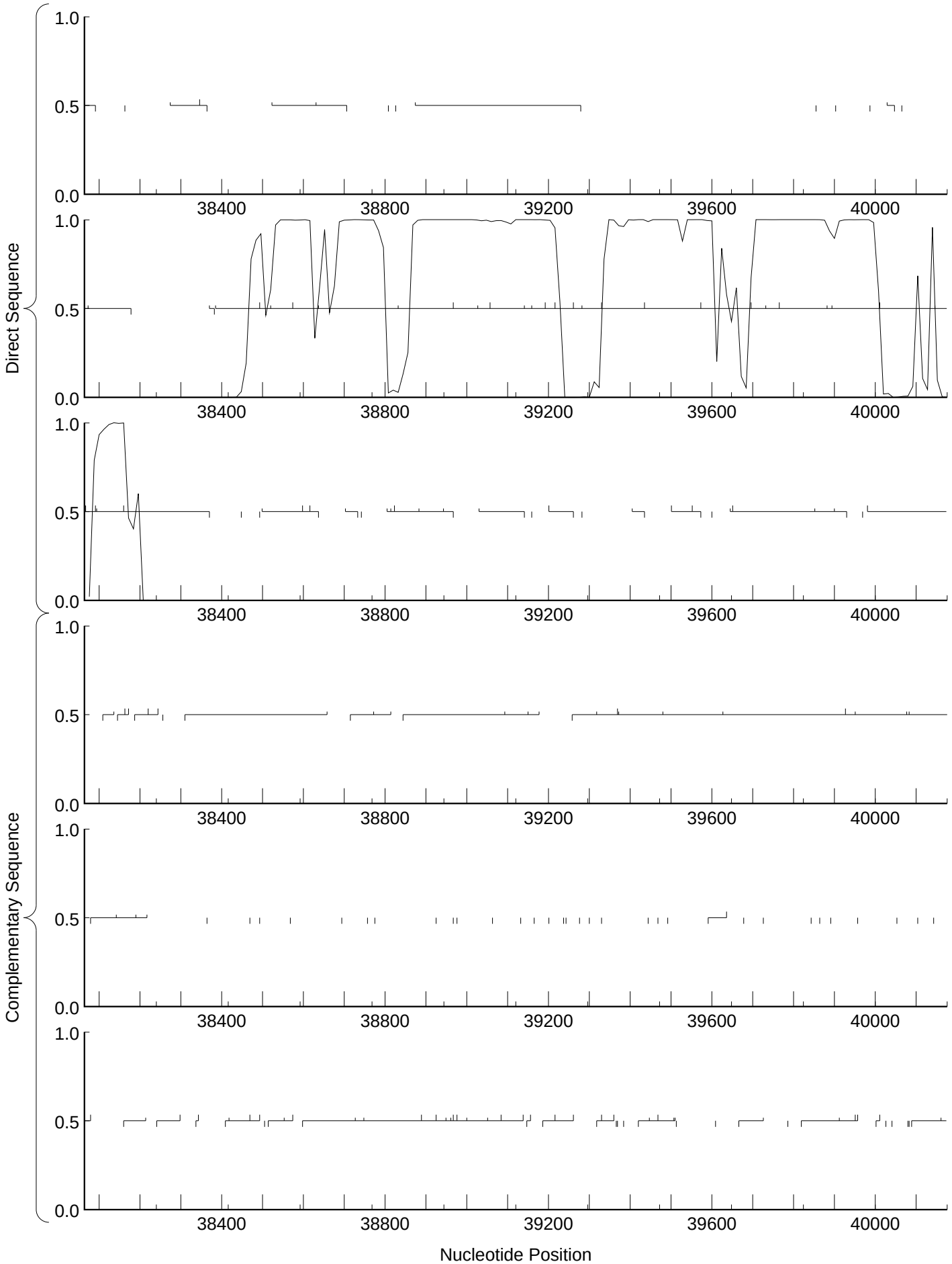
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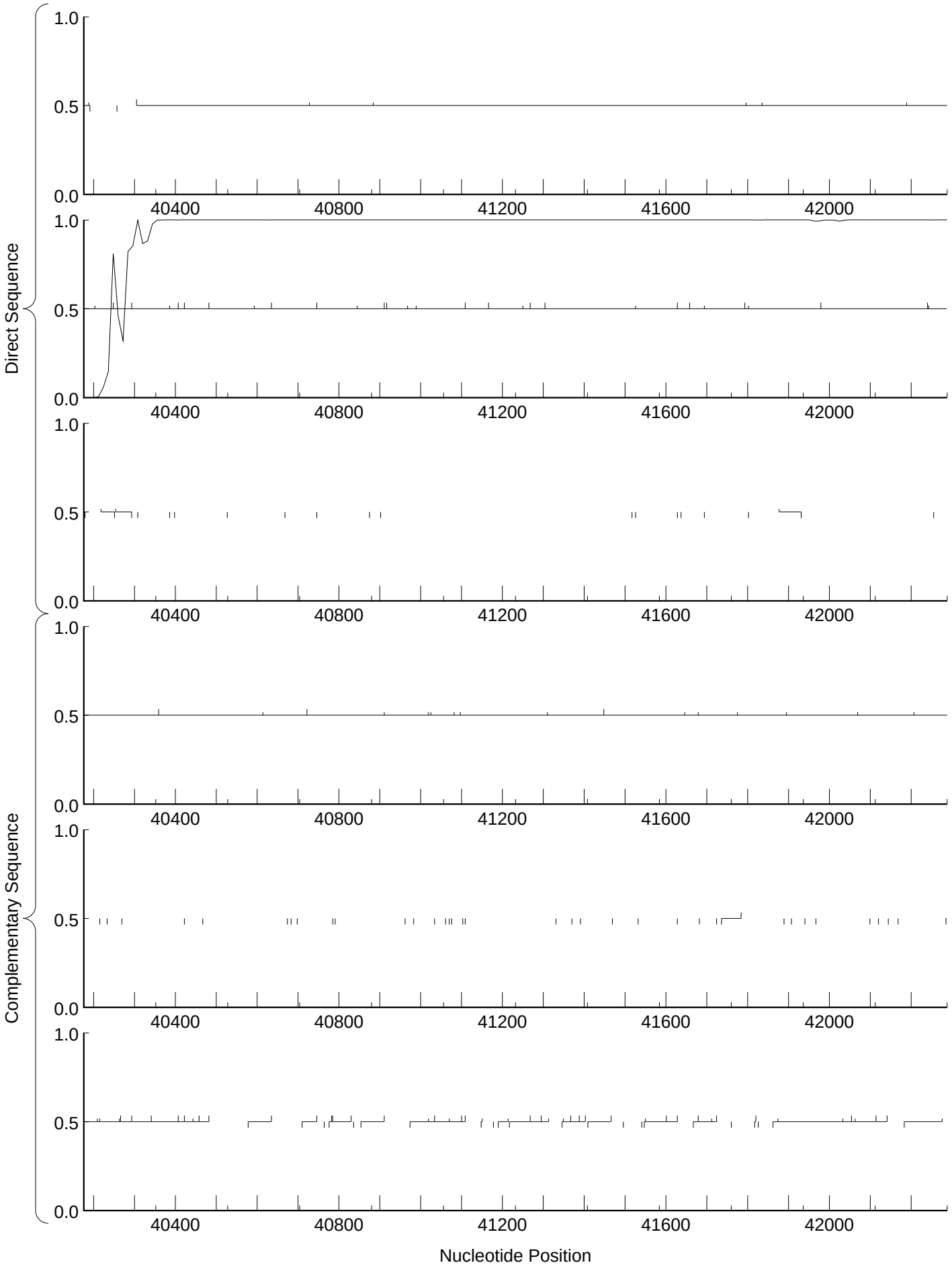
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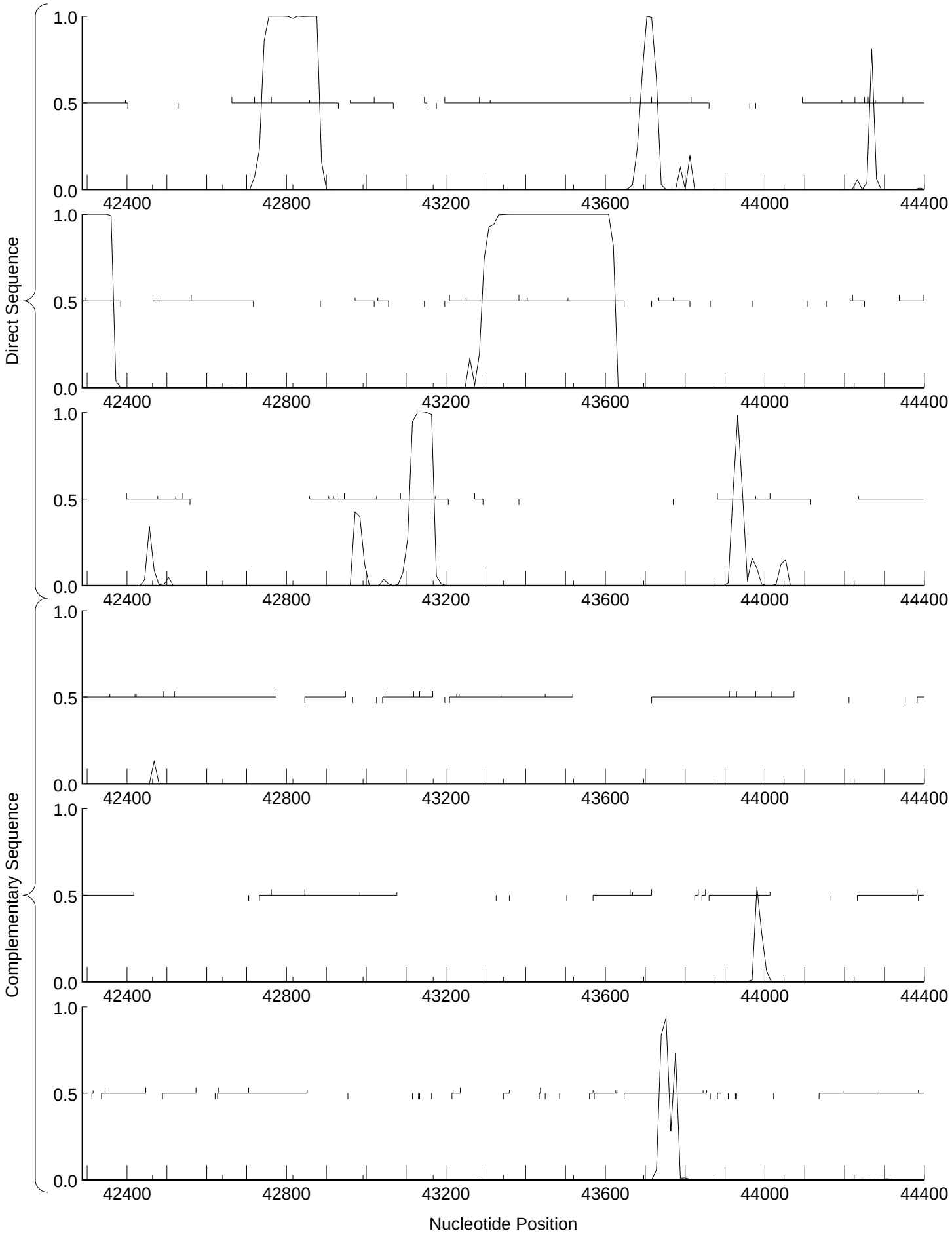
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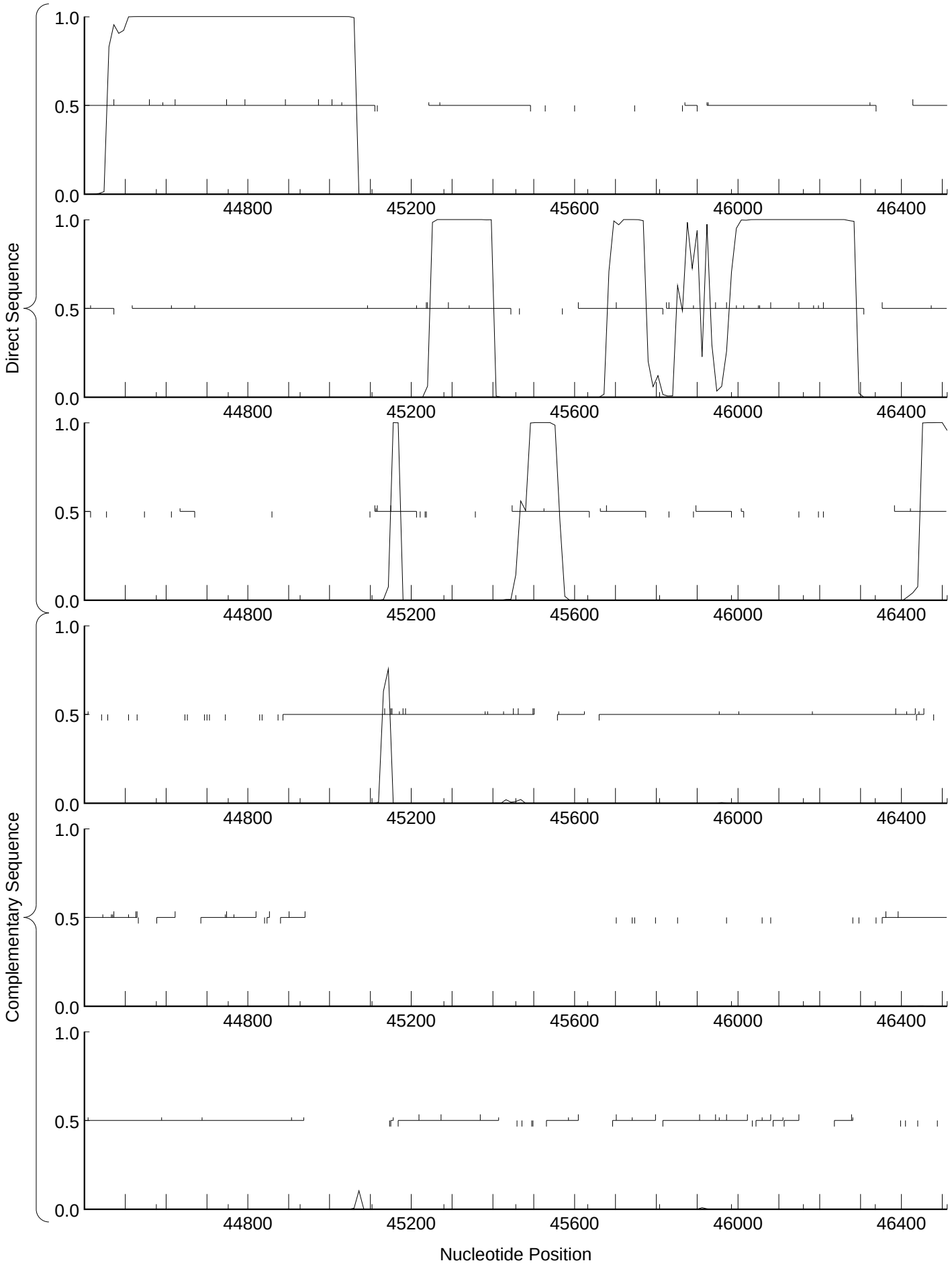


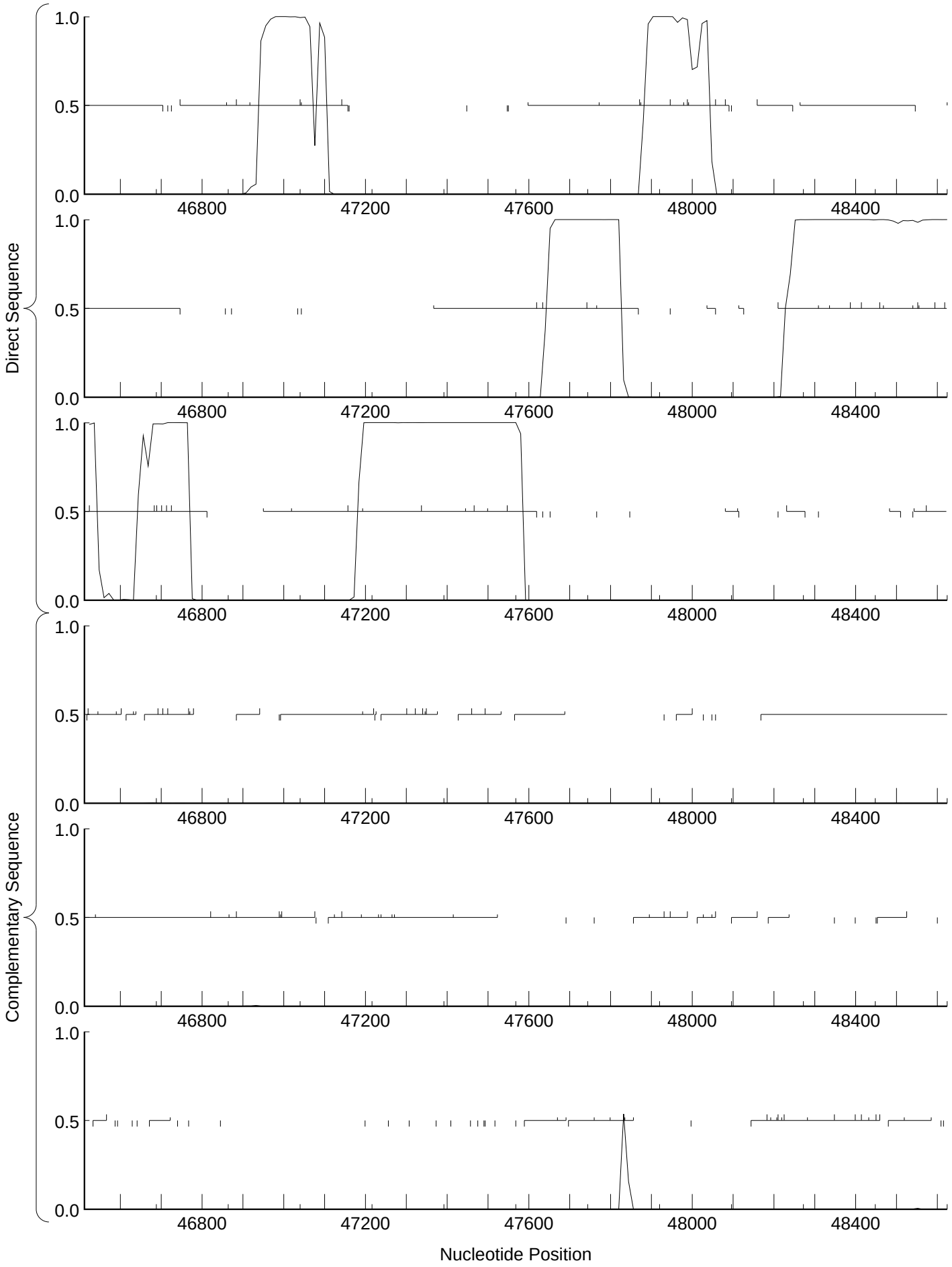
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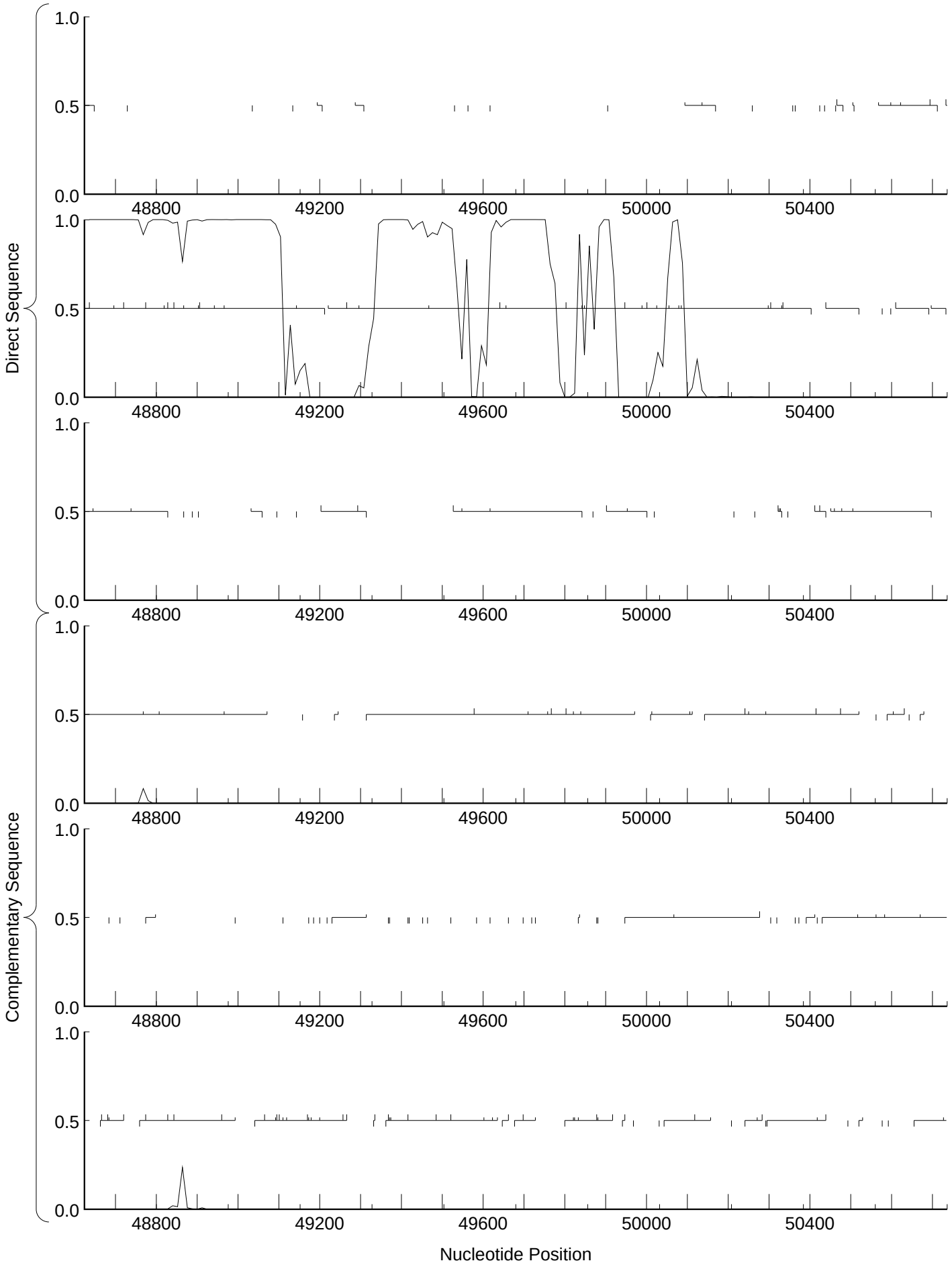


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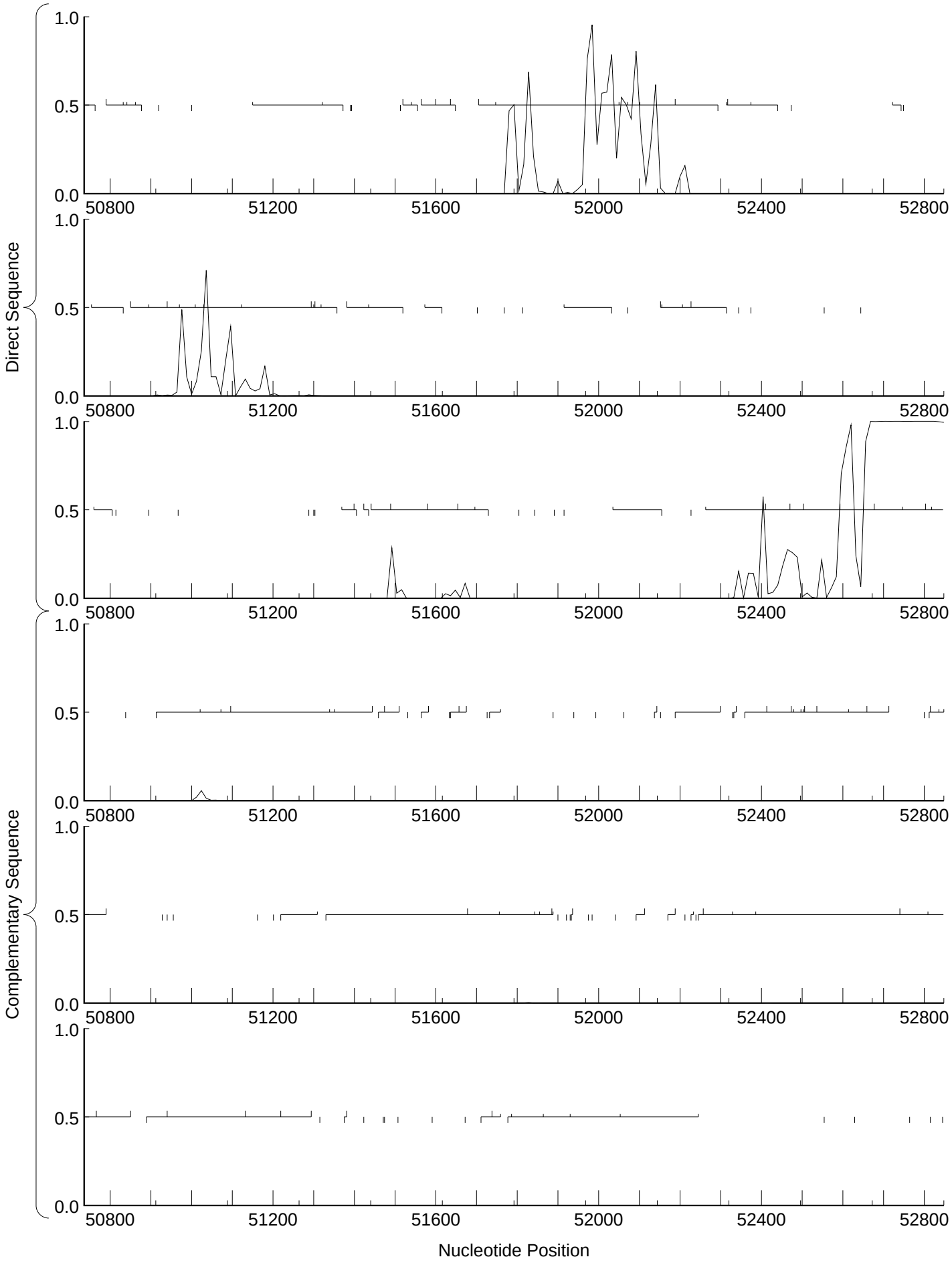



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
44800
45200
45600
46000
46400
Nucleotide Position

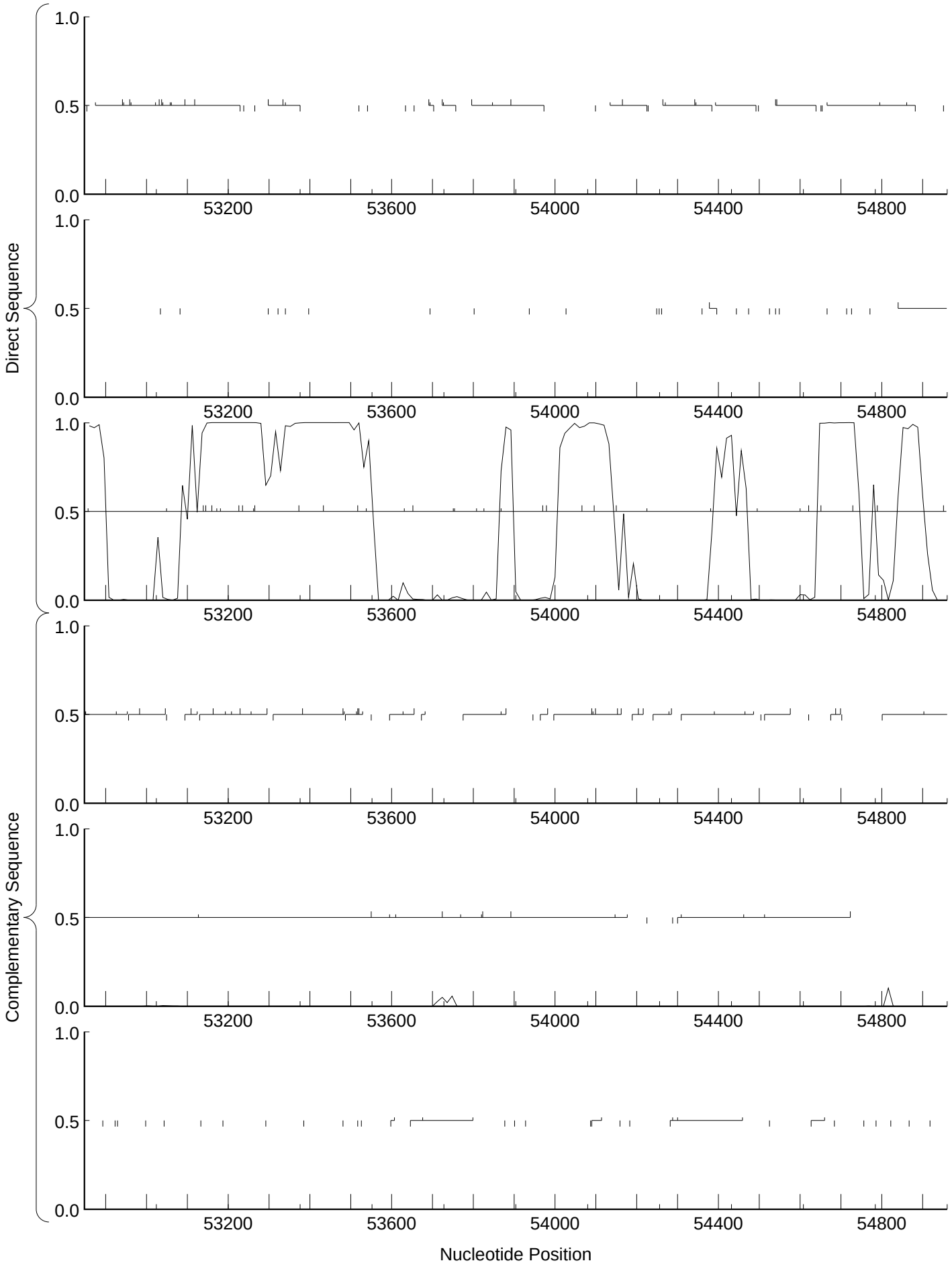

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
46800
47200
47600
48000
48400
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
48800
49200
49600
50000
50400
Nucleotide Position

ver complete sequence, 55521 bp including 9-base 3' overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 26/28



ver complete sequence, 55521 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 27/28



ver complete sequence, 55521 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 28/28

