GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Streptomyces phage Animus complete sequence, 50793 bp including 11-base 3' overhang (CGCCGT
Analysis Date    :  3/29/24 at 8:52:58
Pages            :  26
Sequence Length  :  50793 bp
GC Content       :   67.02%

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

PS-Version       :  1.2

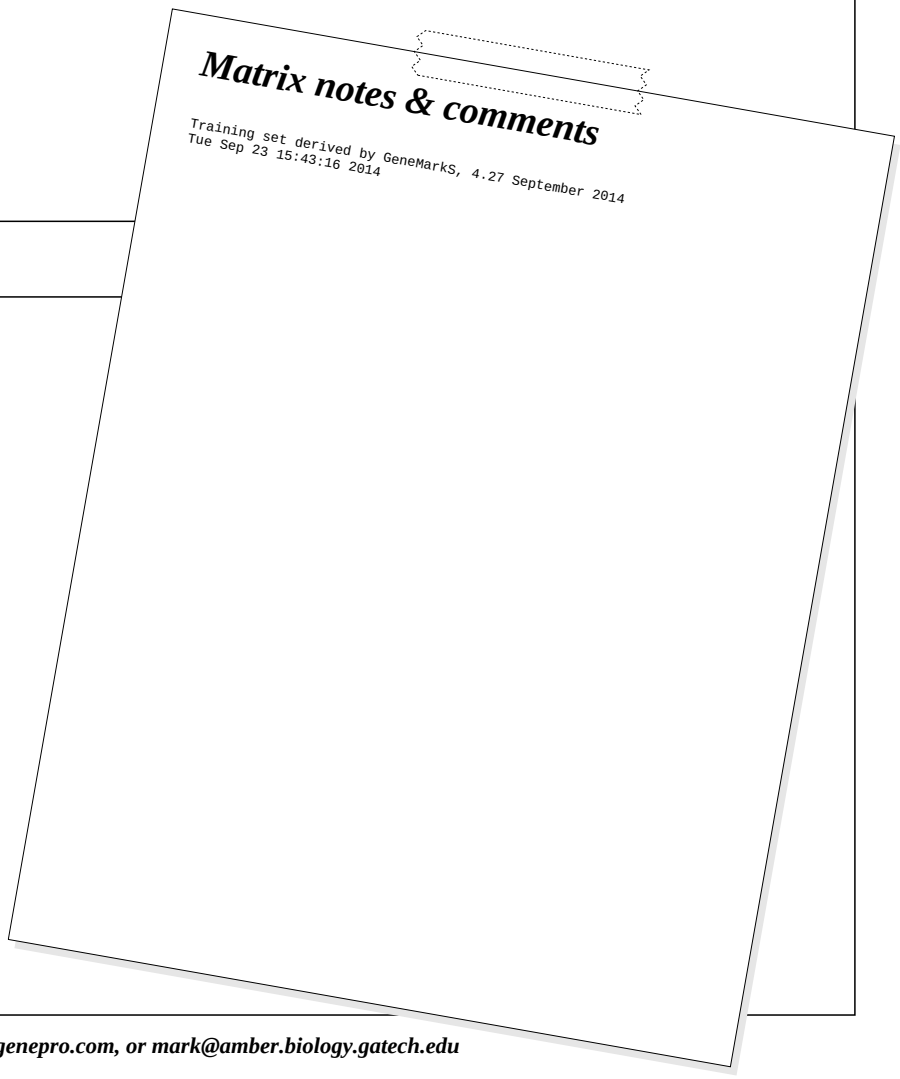
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Matrix     :  Streptomyces_scabiei_87_22
Author     :  -

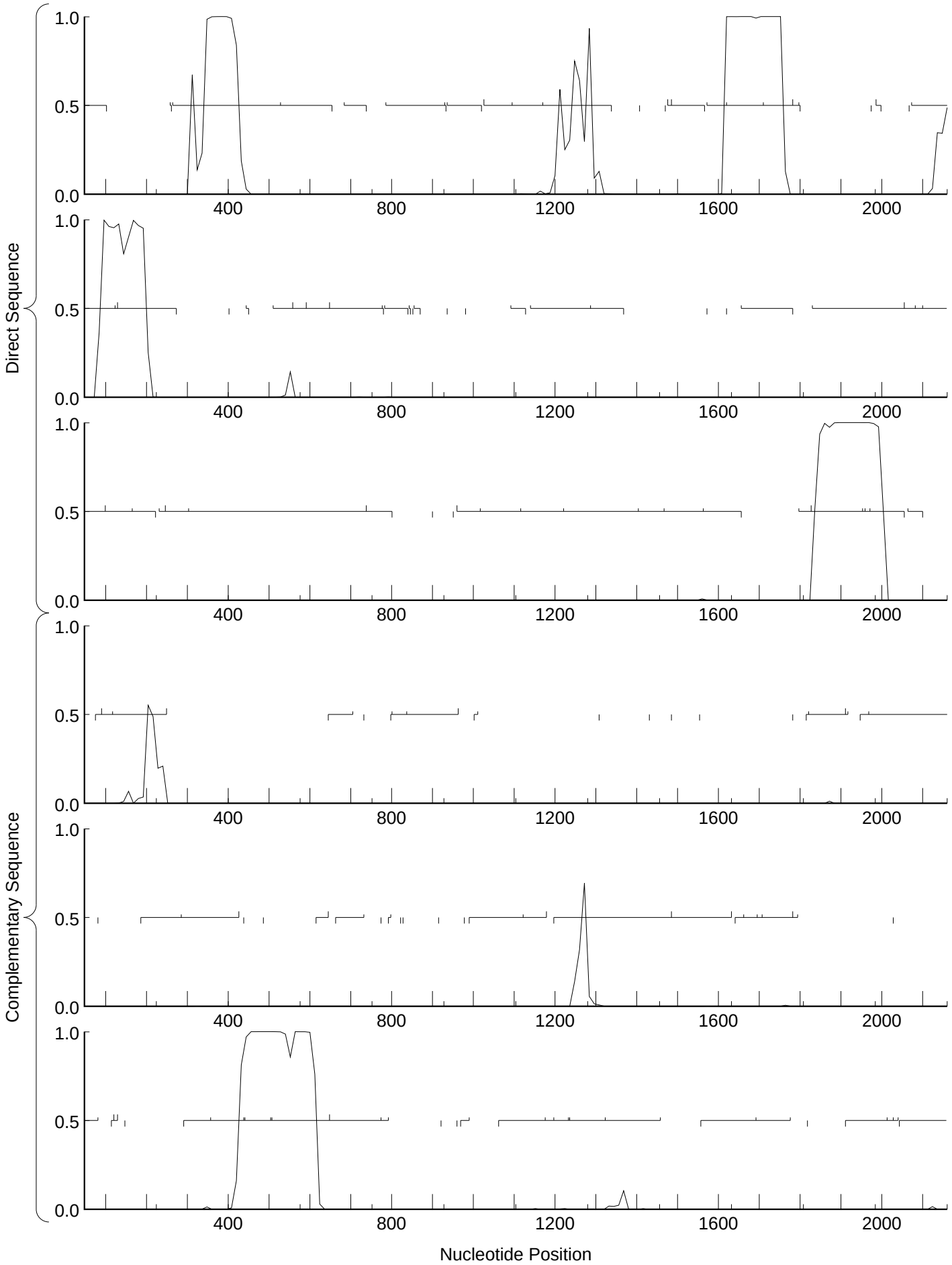
Order      :  4

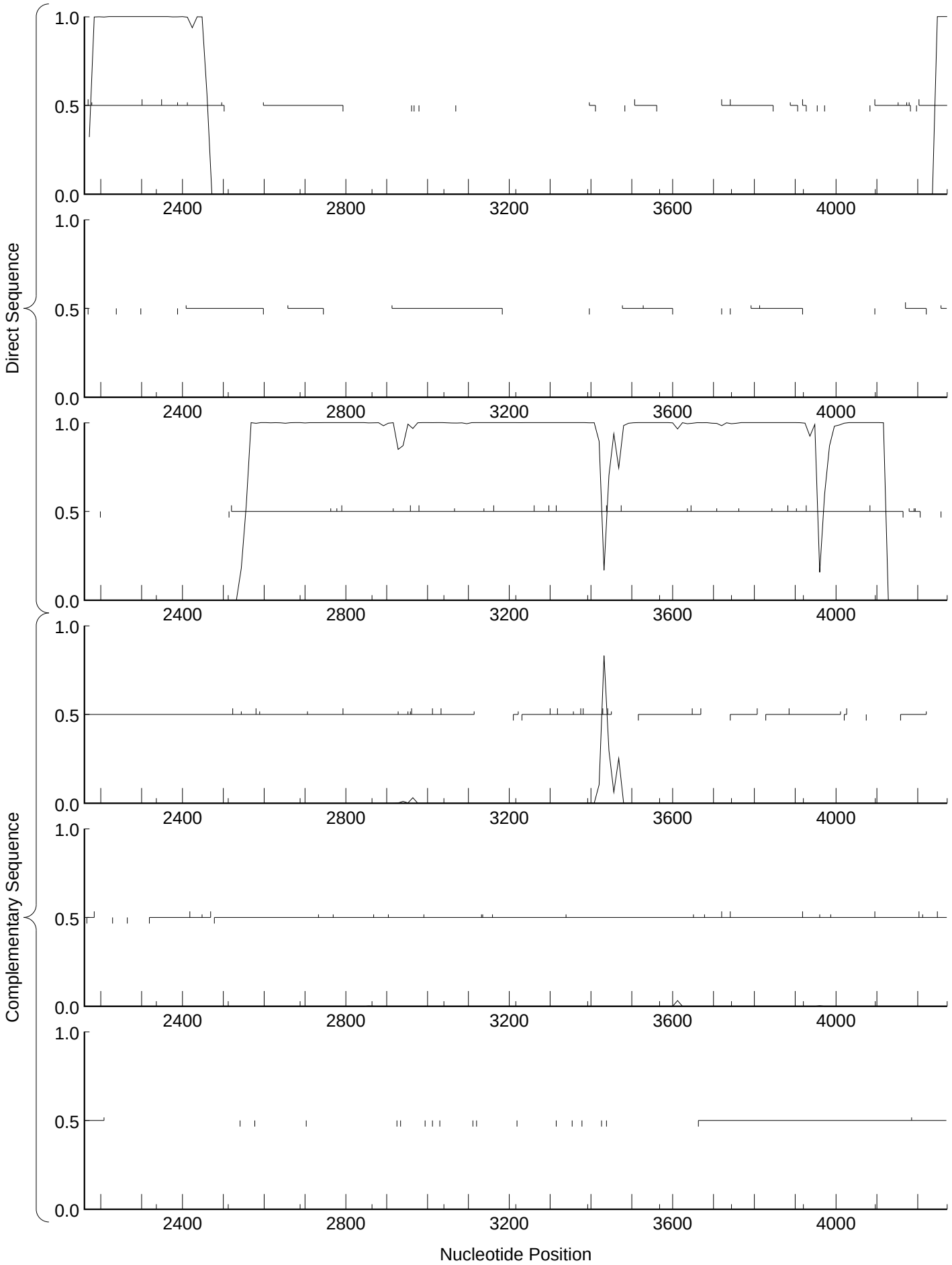

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```



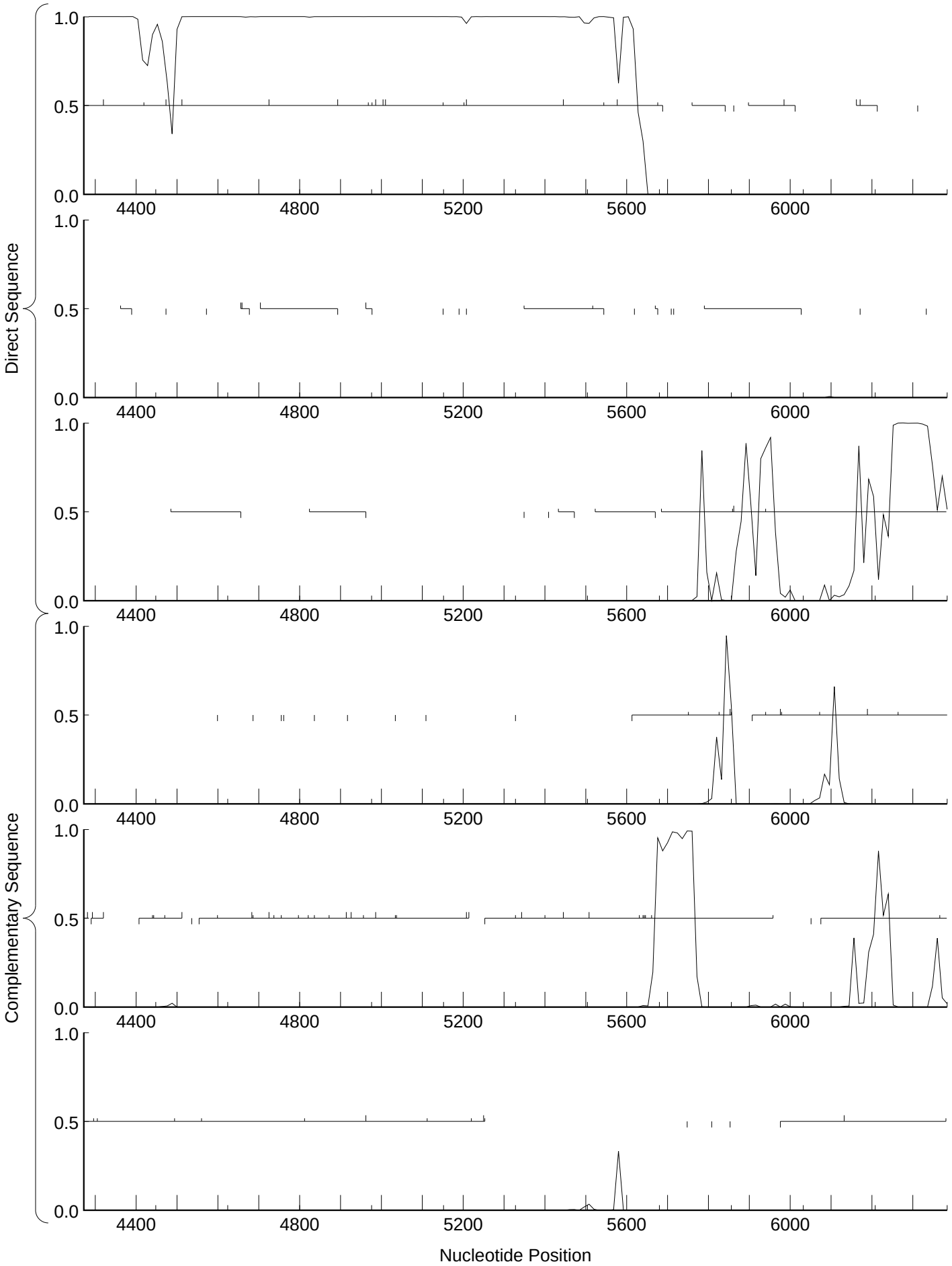
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

complete sequence, 50793 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 2/26

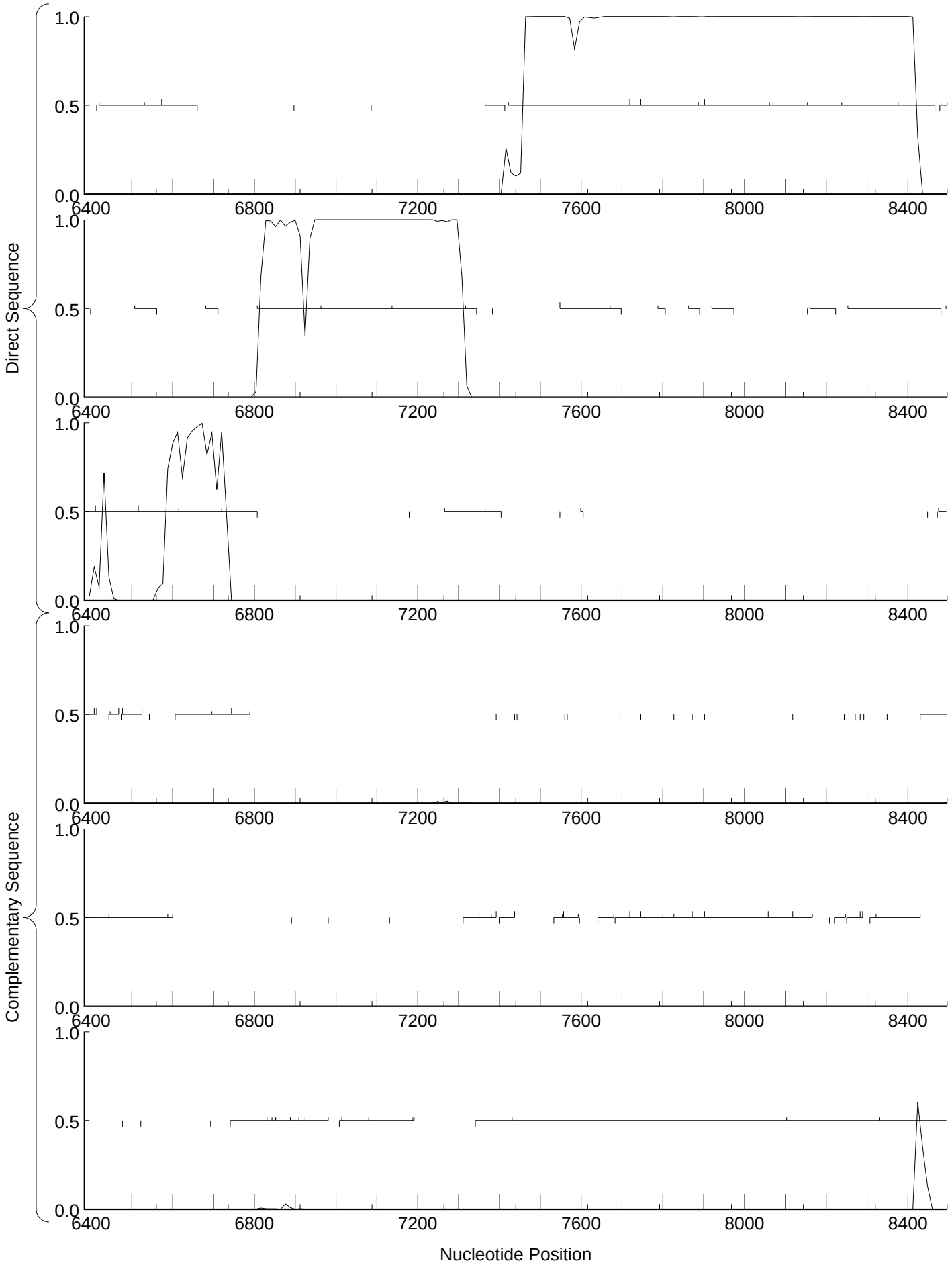



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Complementary Sequence
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Nucleotide Position

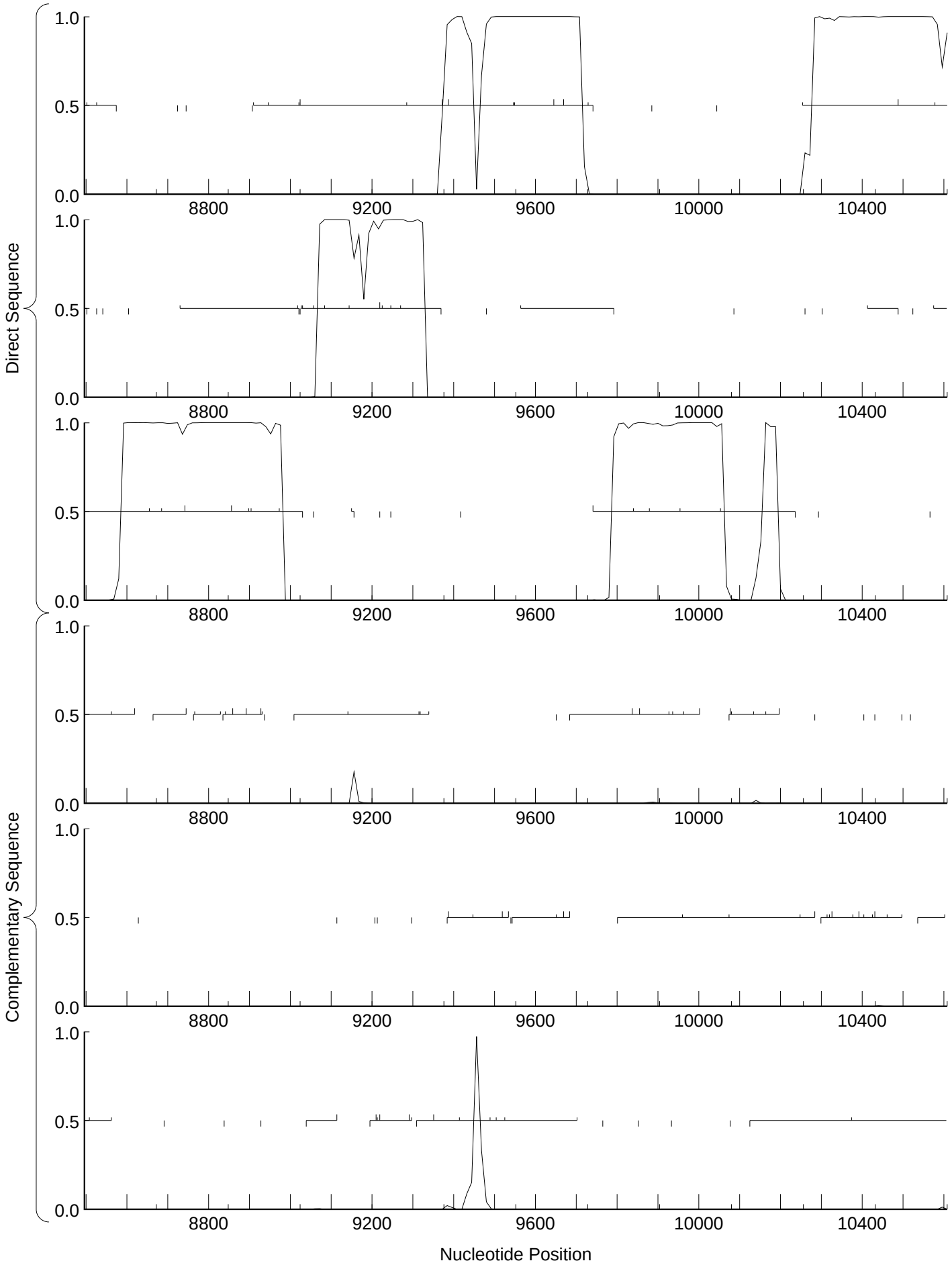
complete sequence, 50793 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 4/26



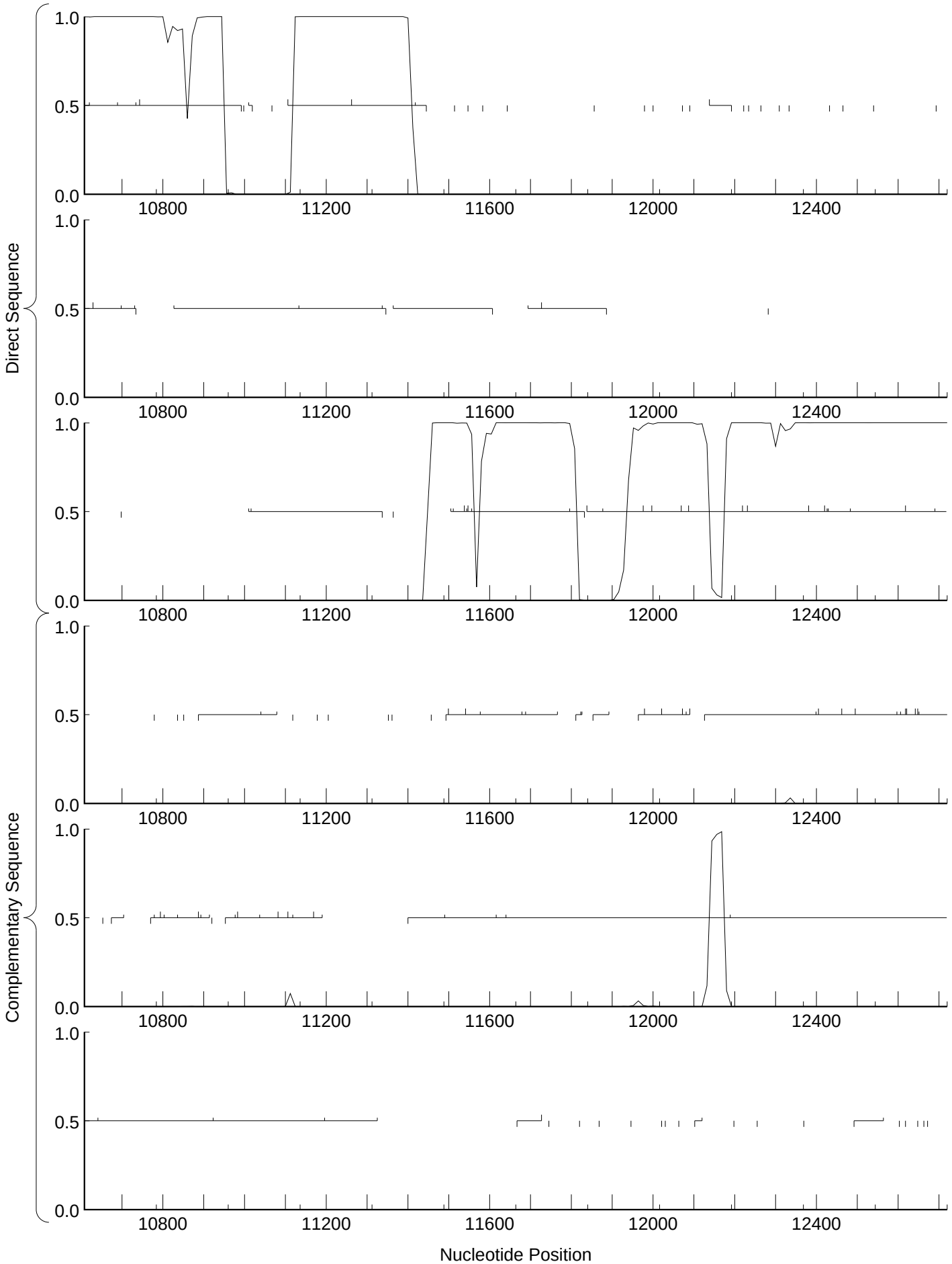
complete sequence, 50793 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 5/26



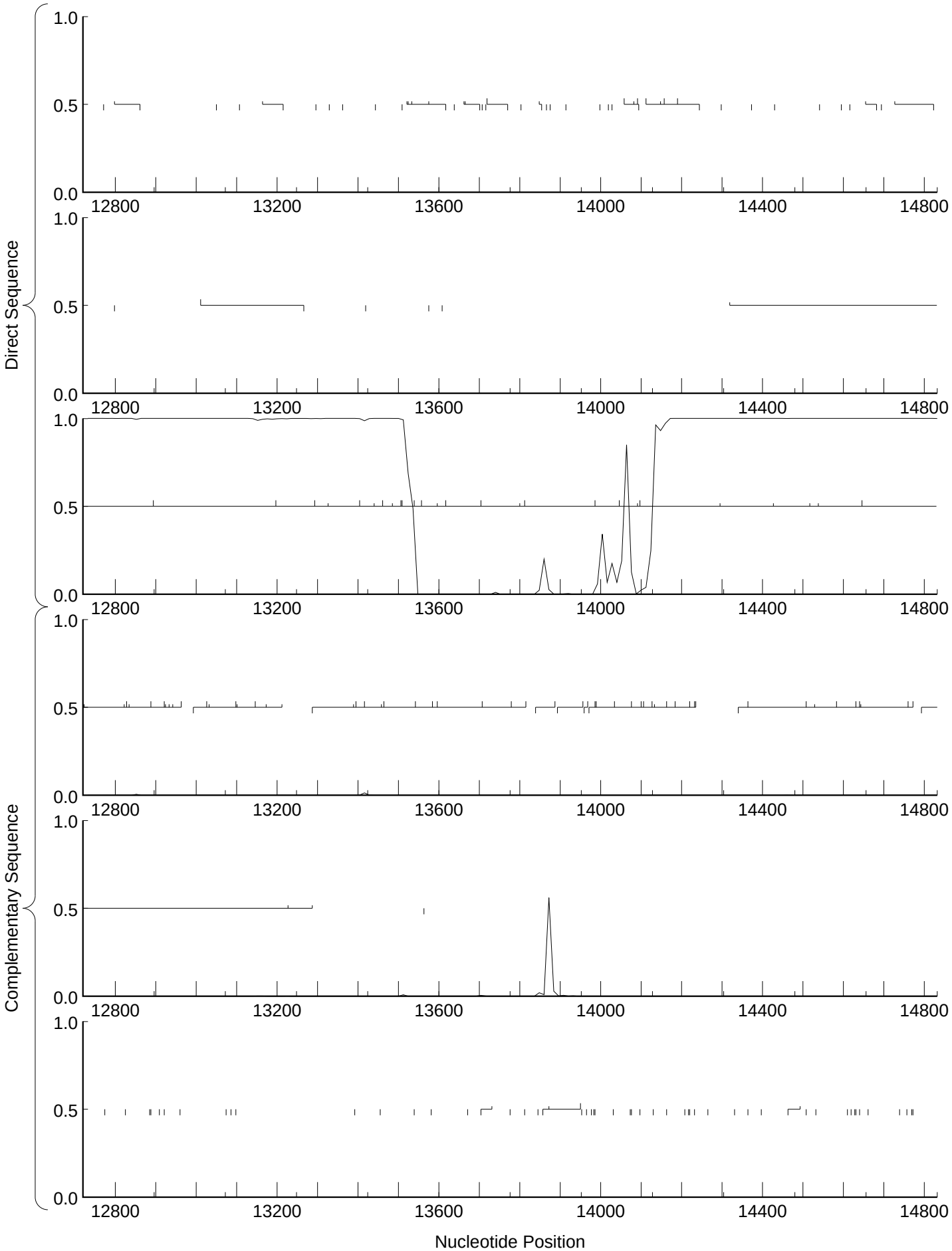
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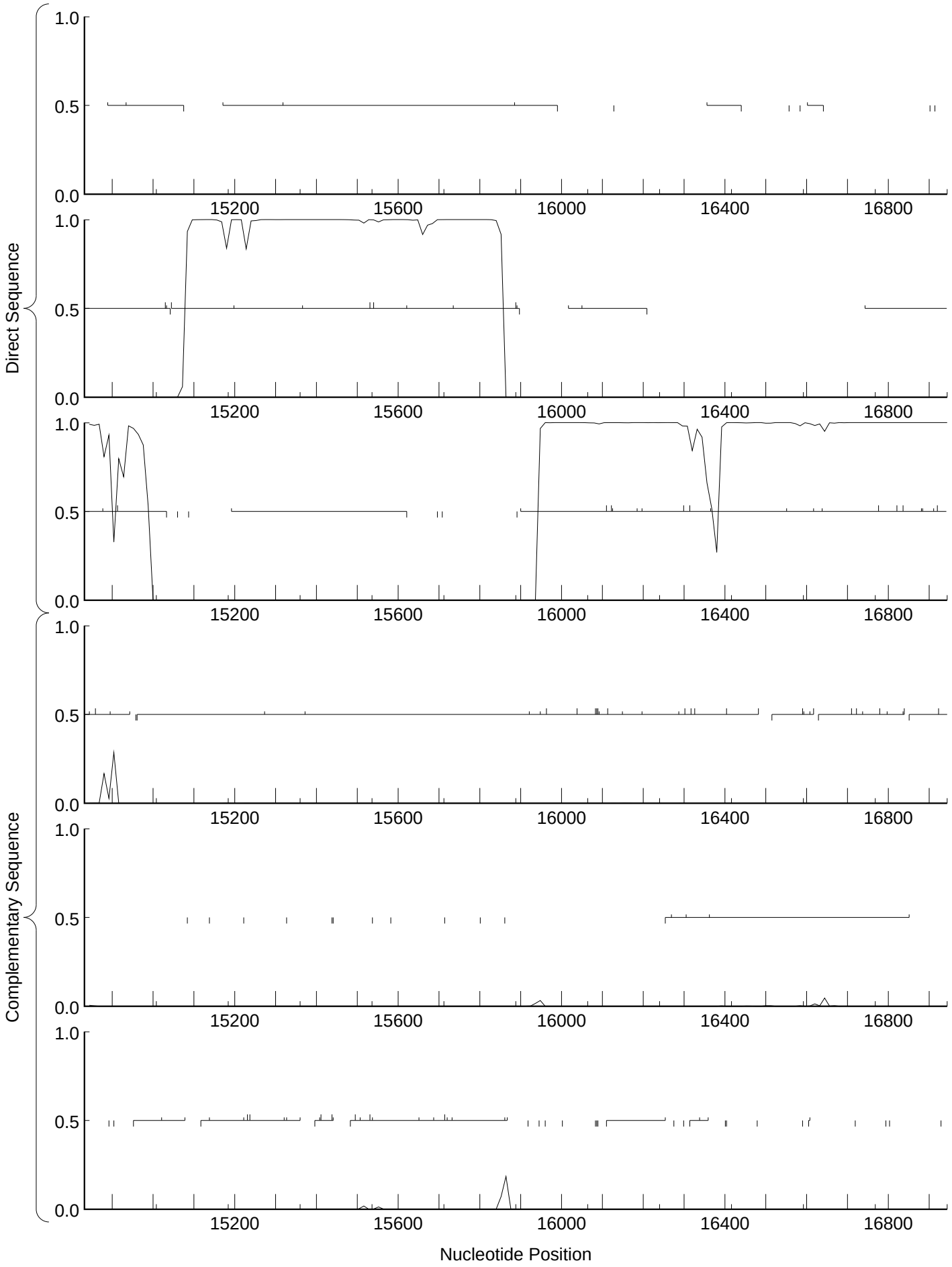
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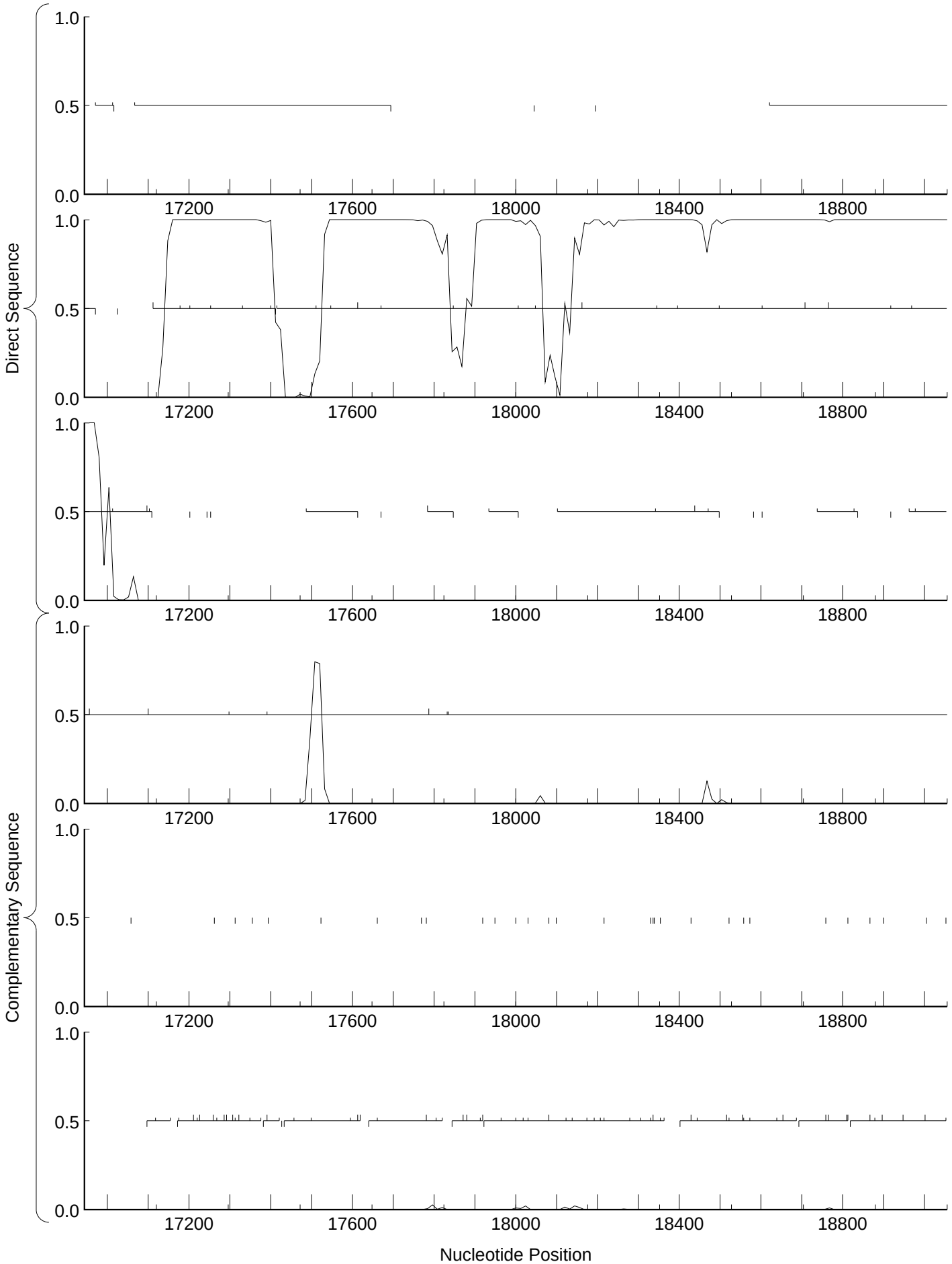
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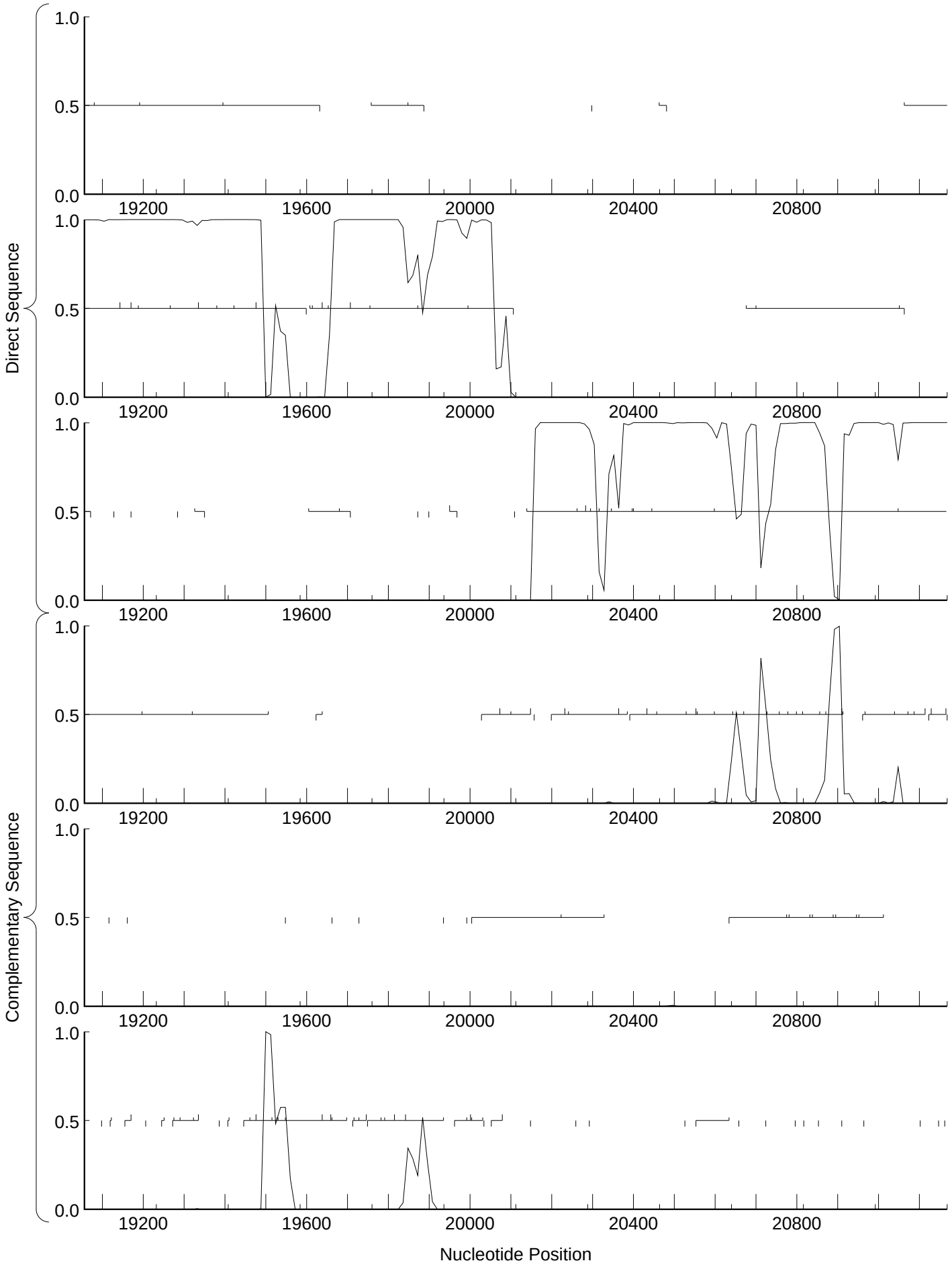
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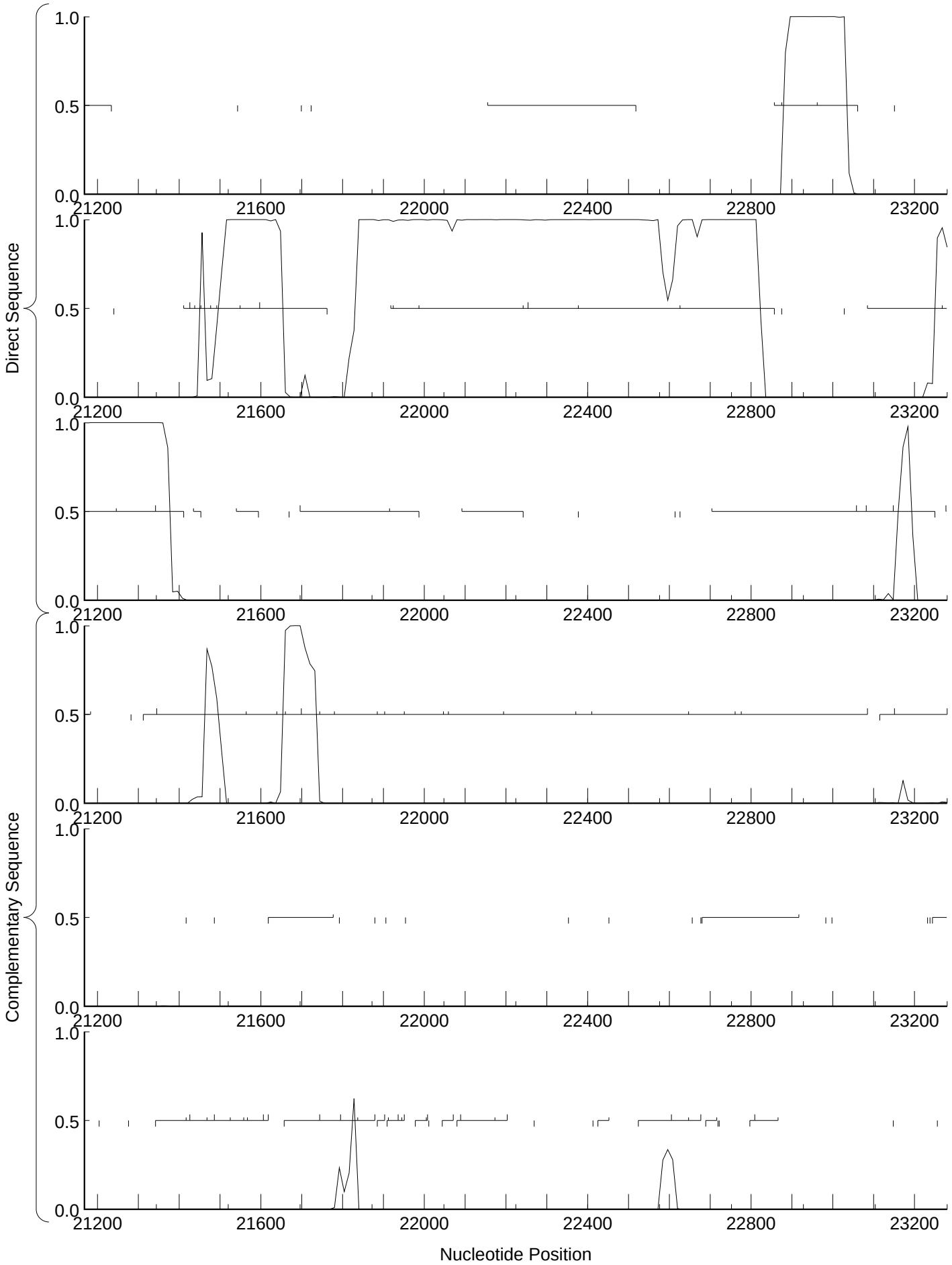
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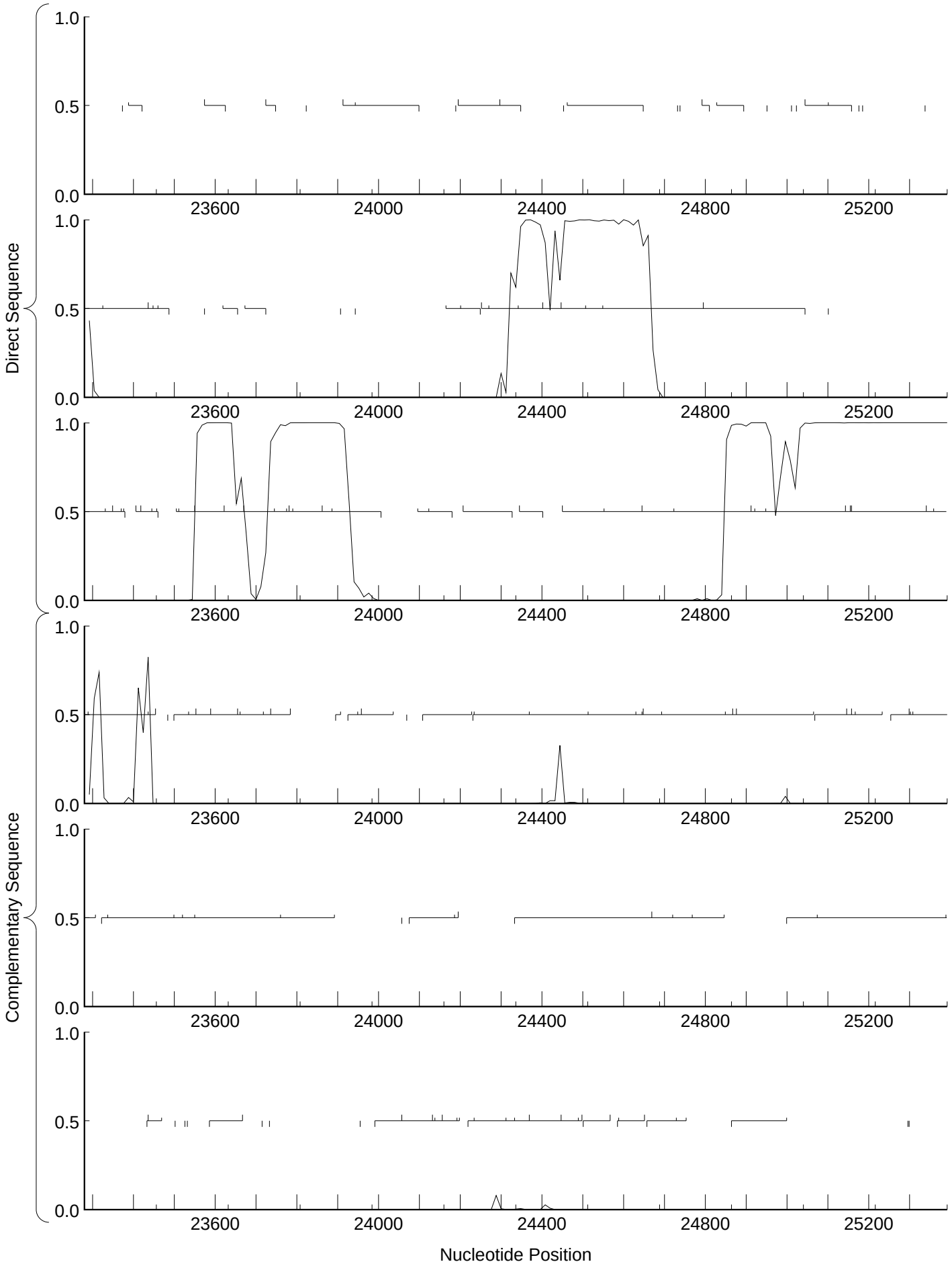
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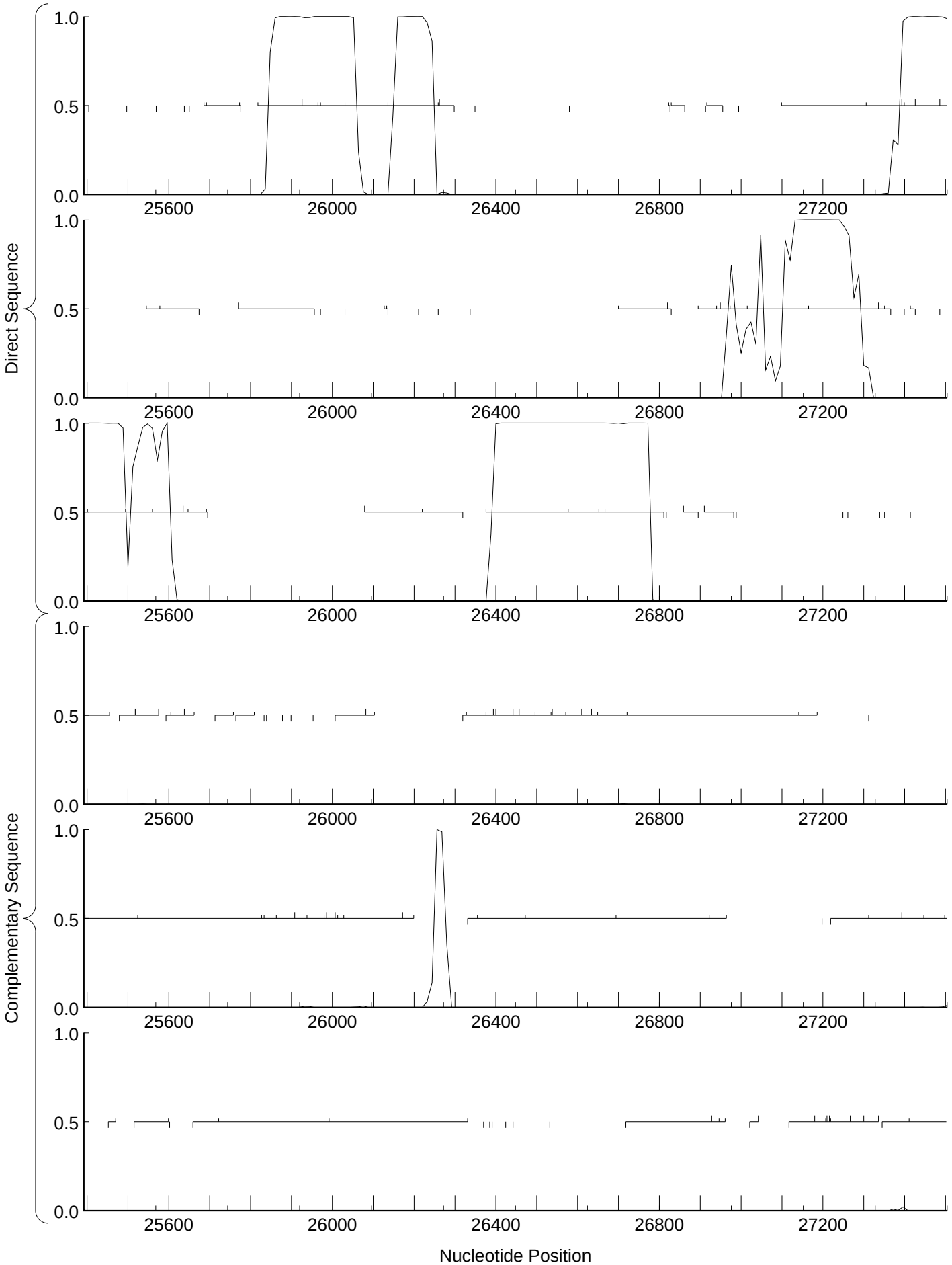
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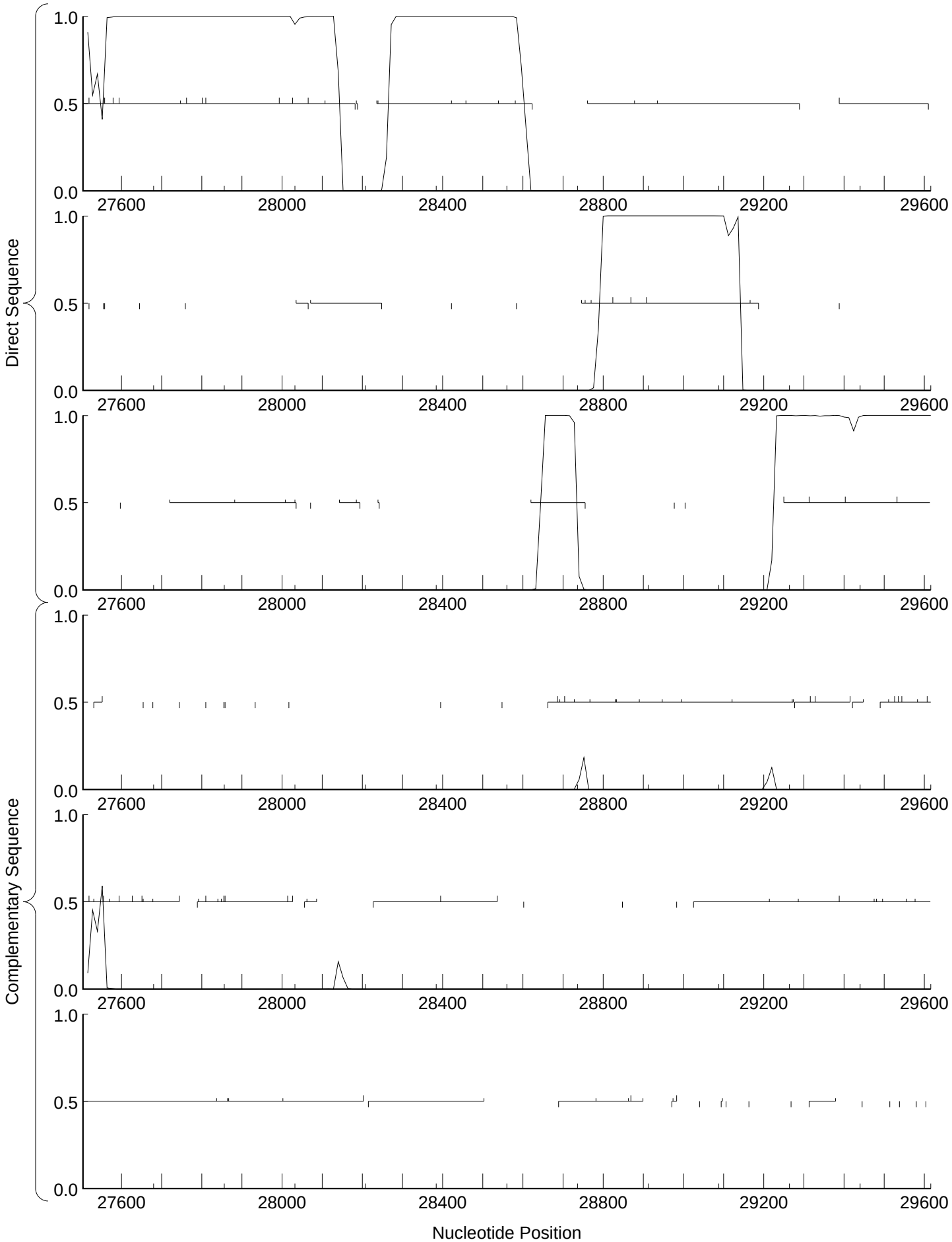
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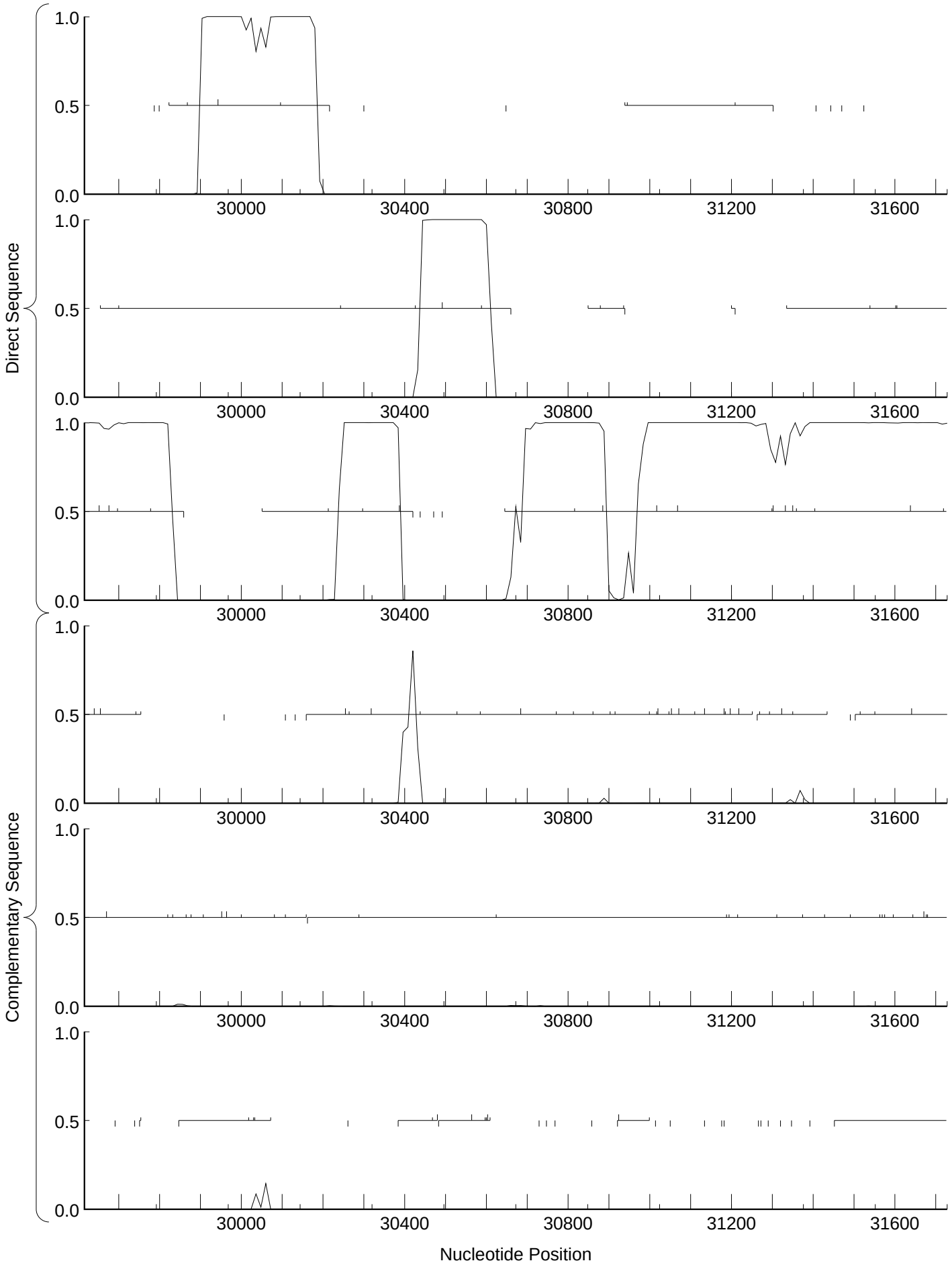


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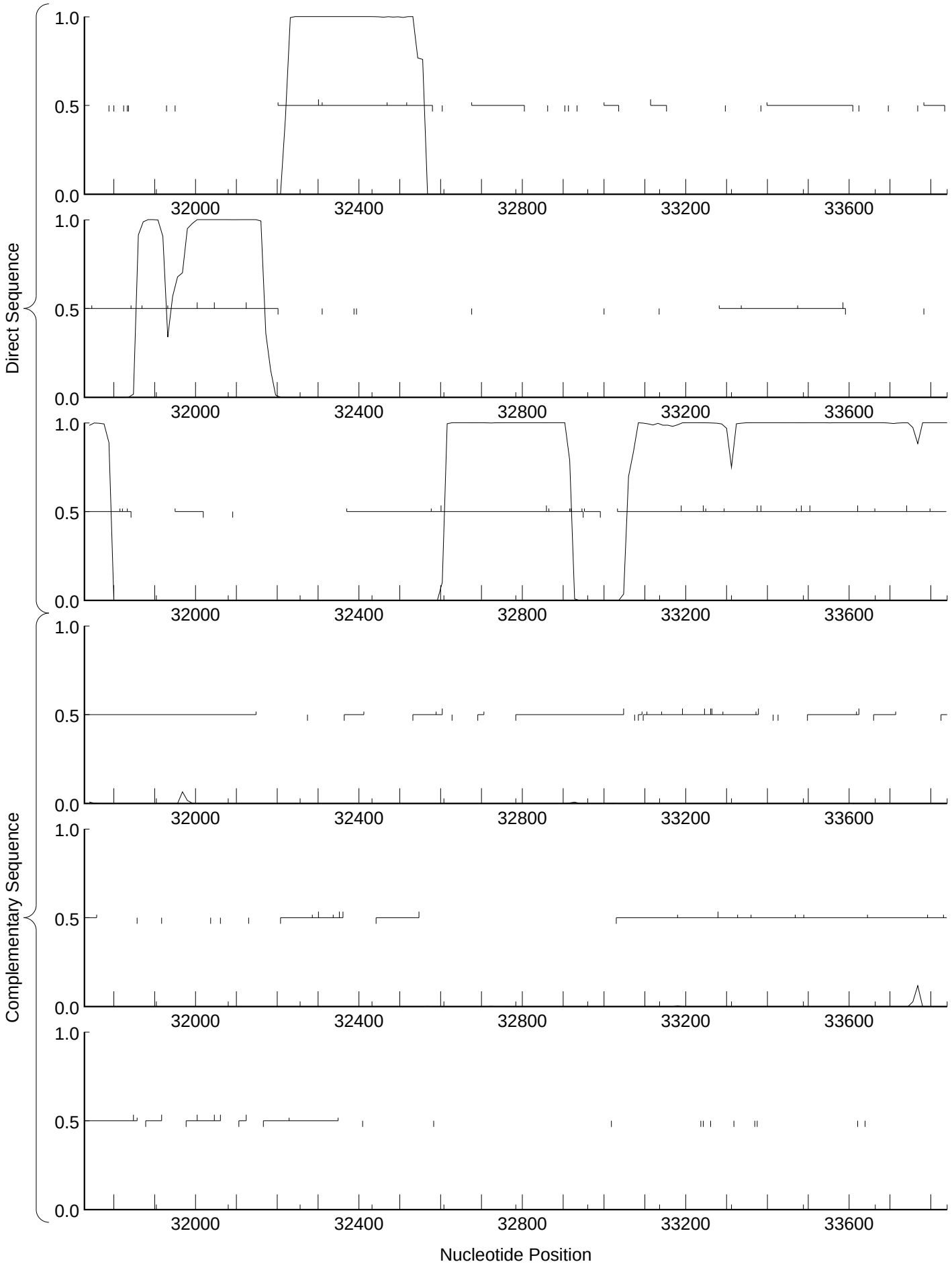


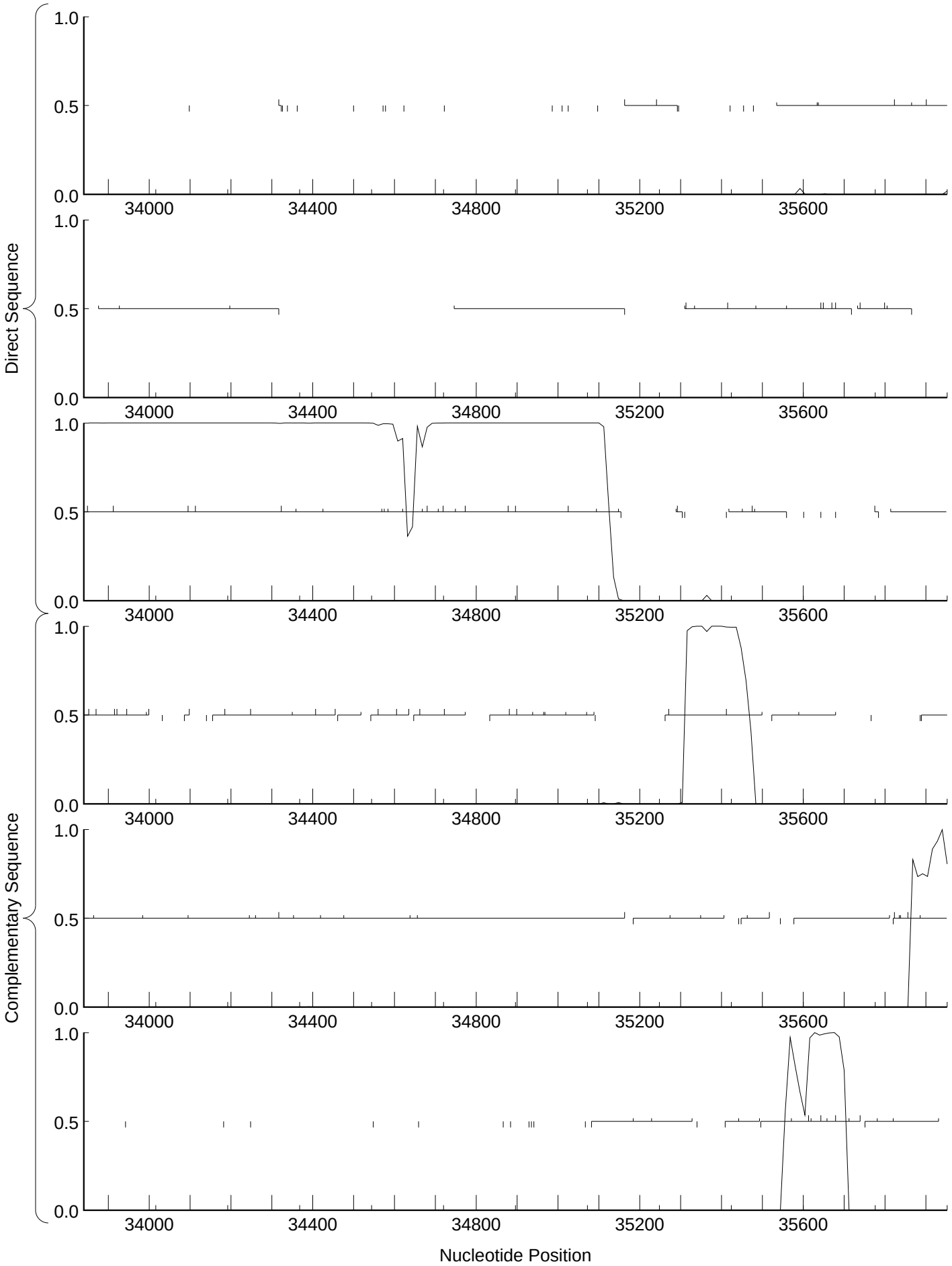
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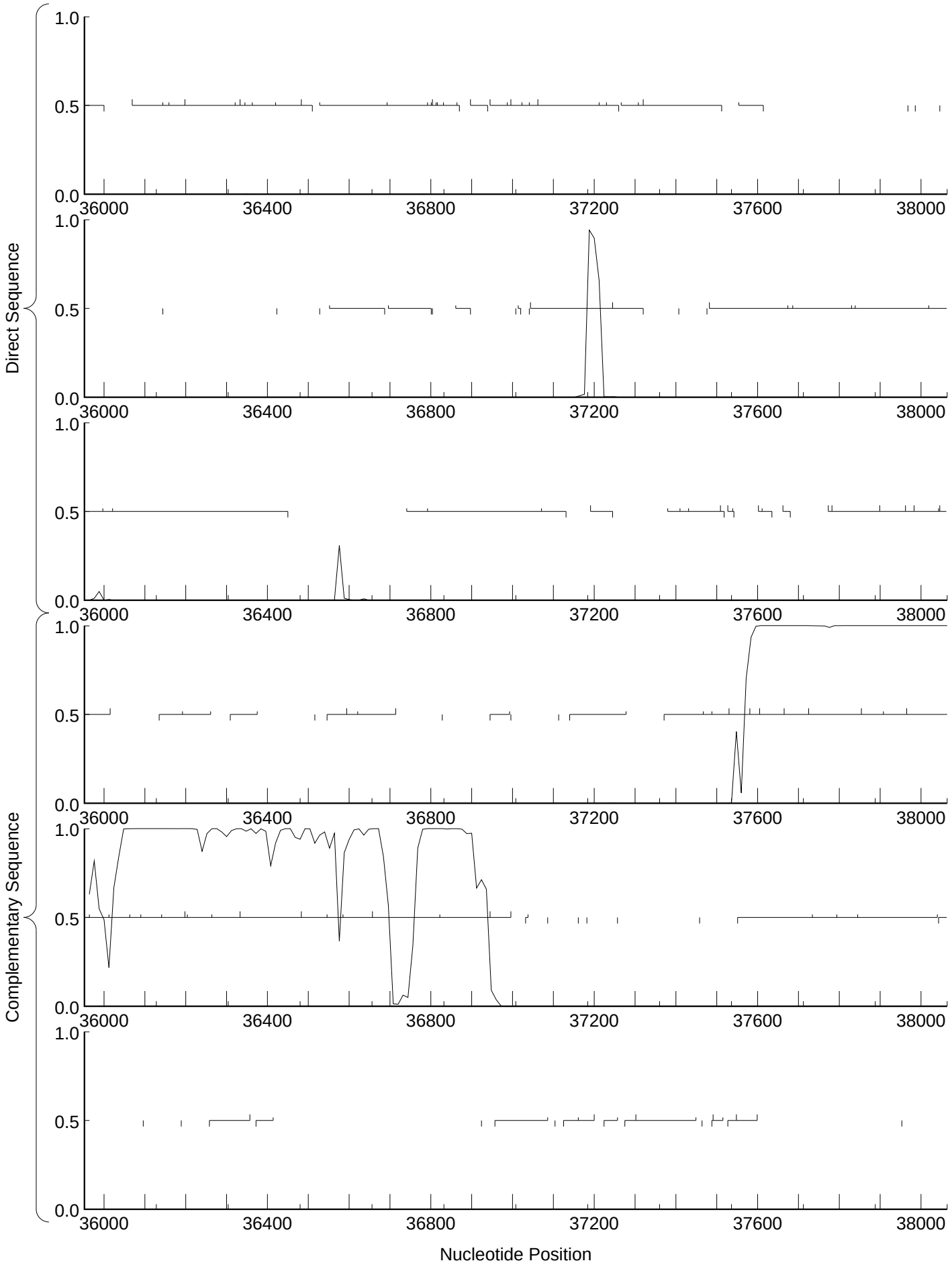


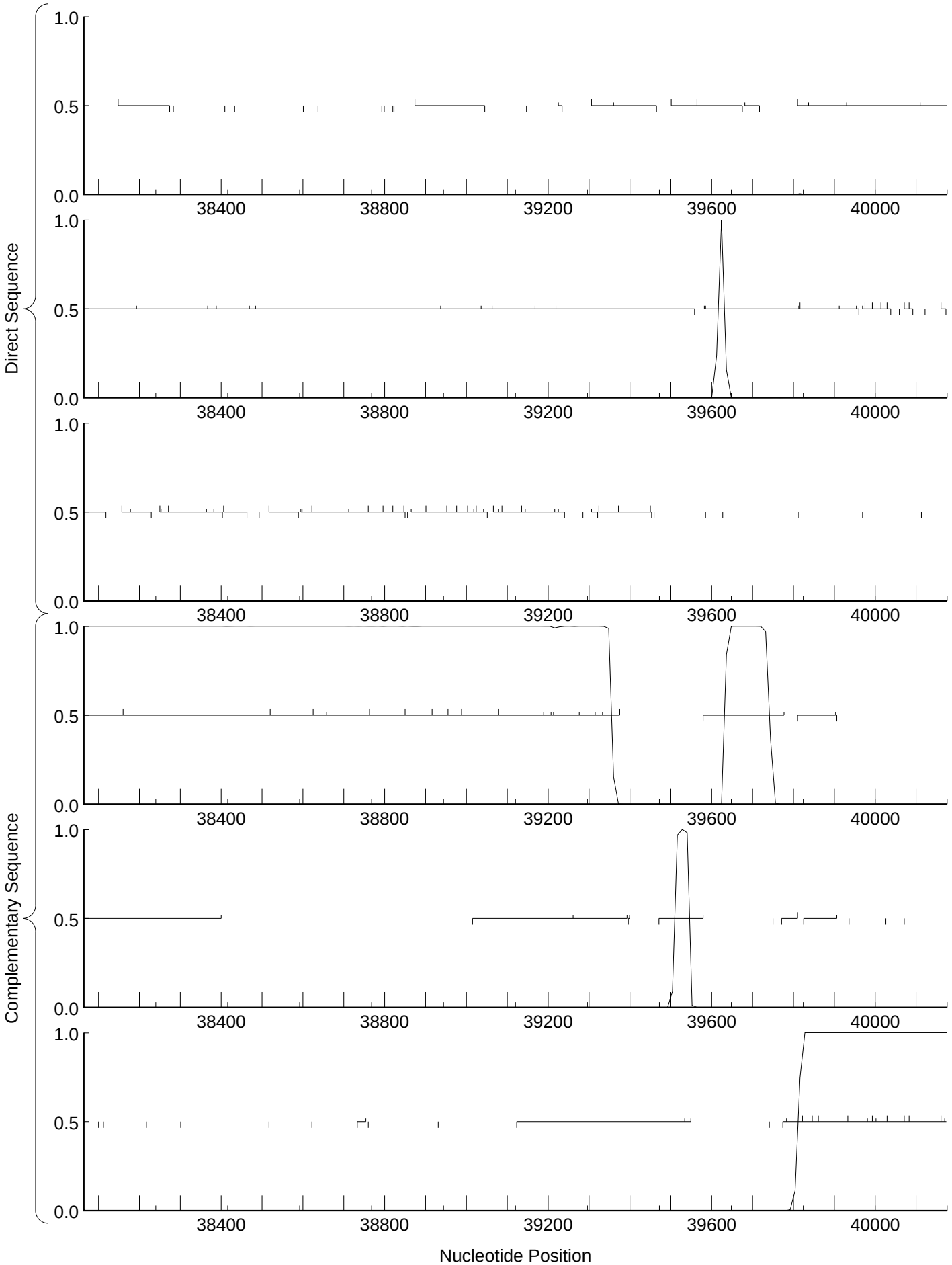

Direct Sequence
Complementary Sequence
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0.0
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Nucleotide Position

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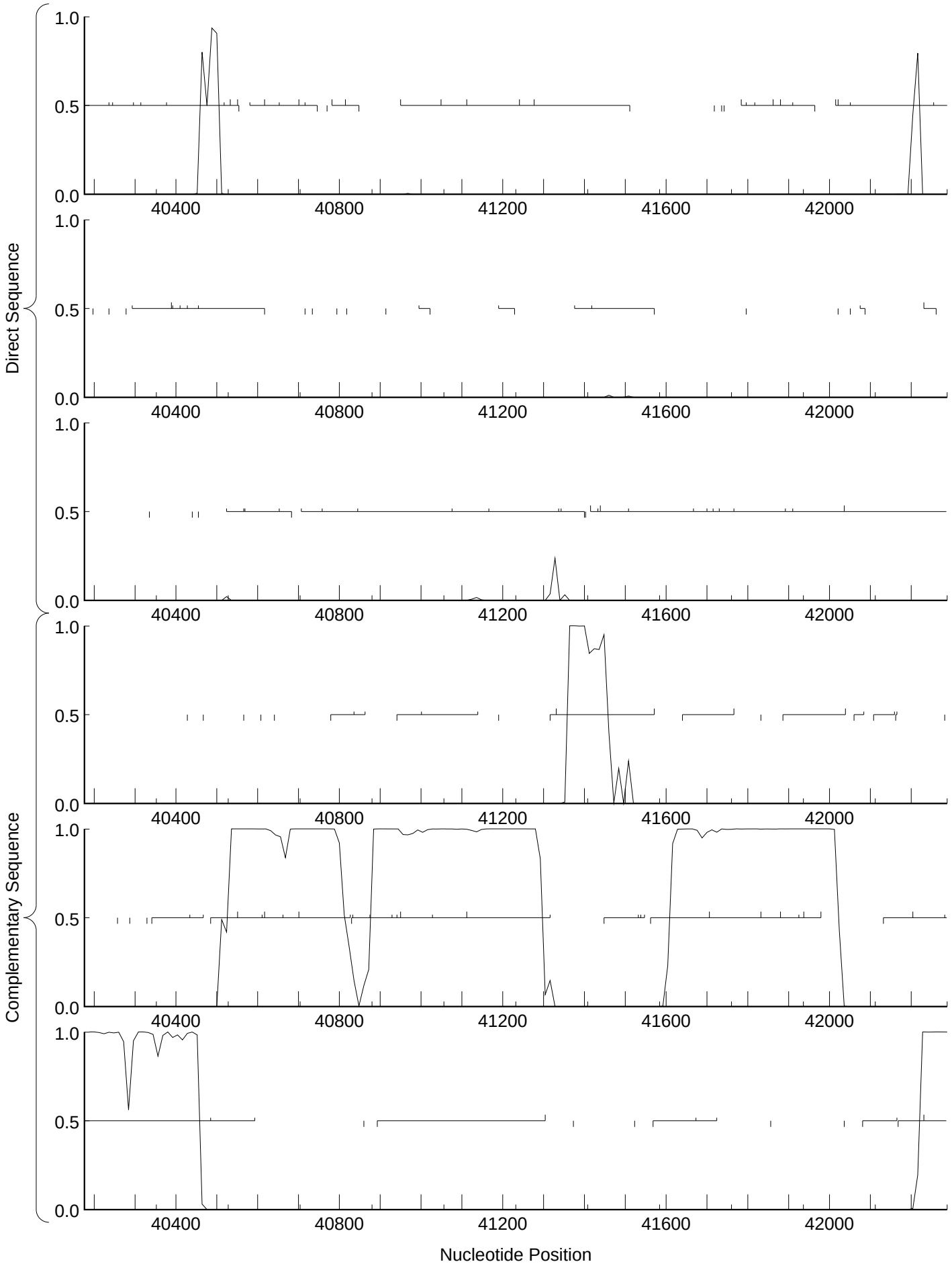



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Complementary Sequence
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Nucleotide Position

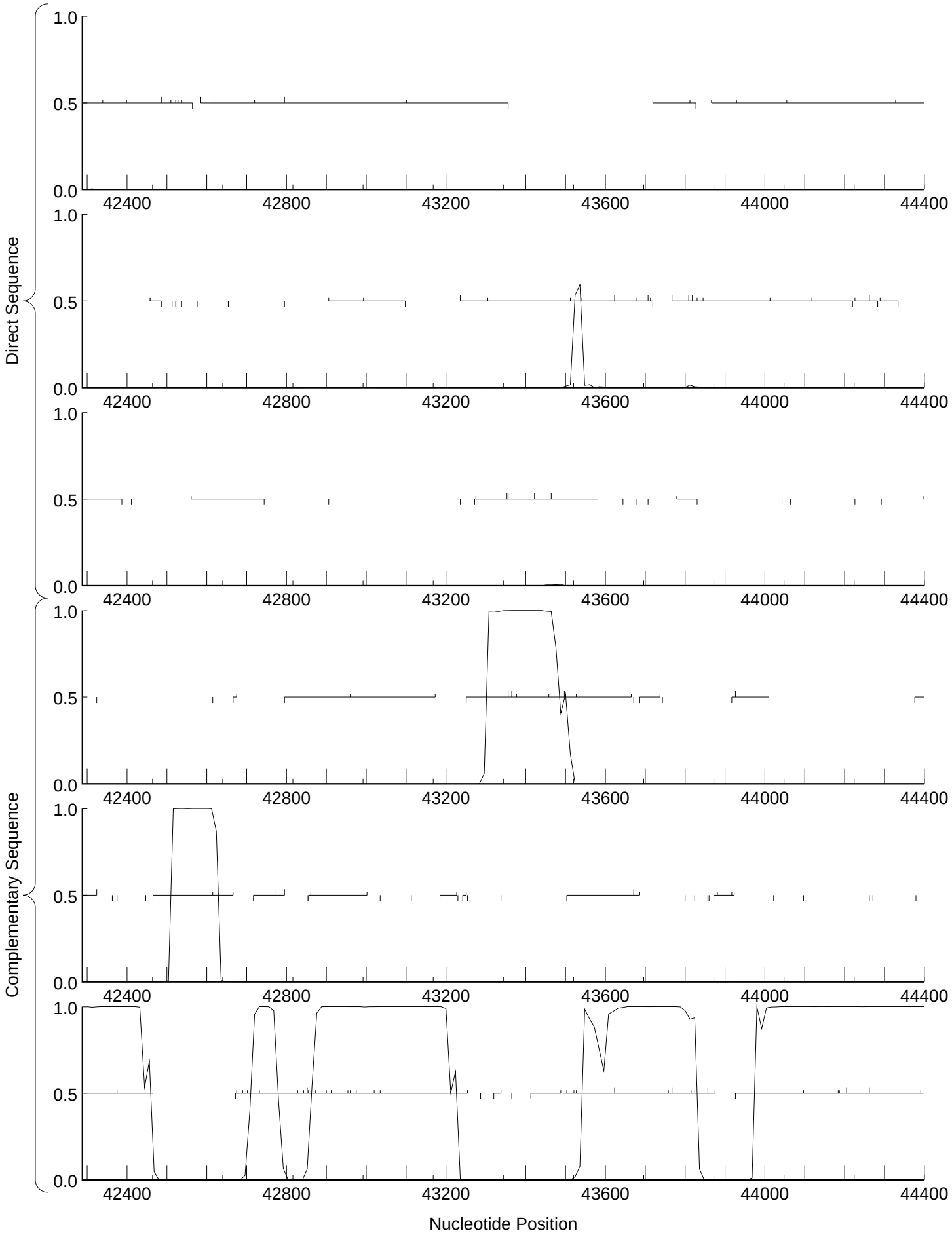

Direct Sequence
Complementary Sequence
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37200
37600
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Nucleotide Position


Direct Sequence
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Nucleotide Position

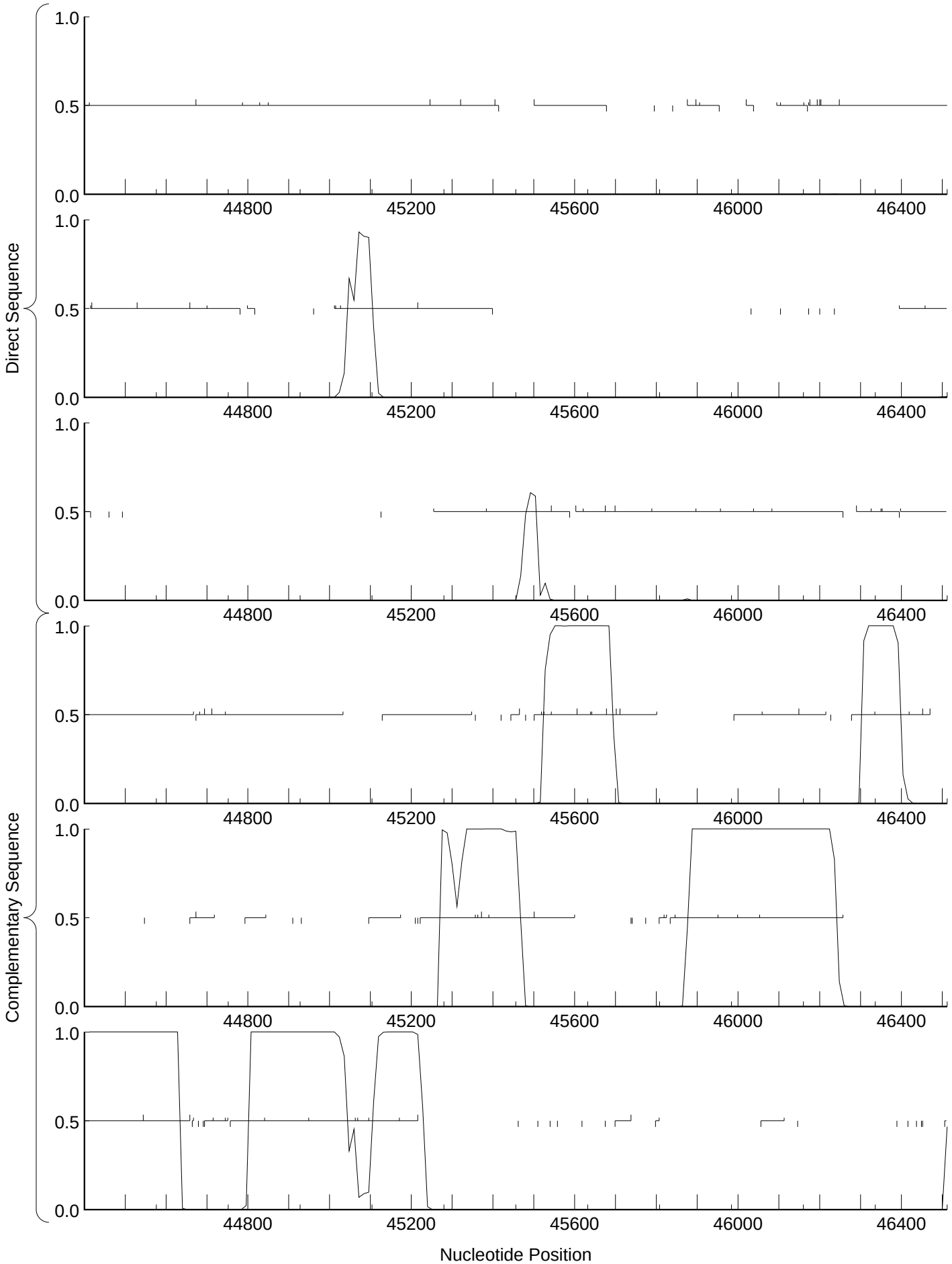
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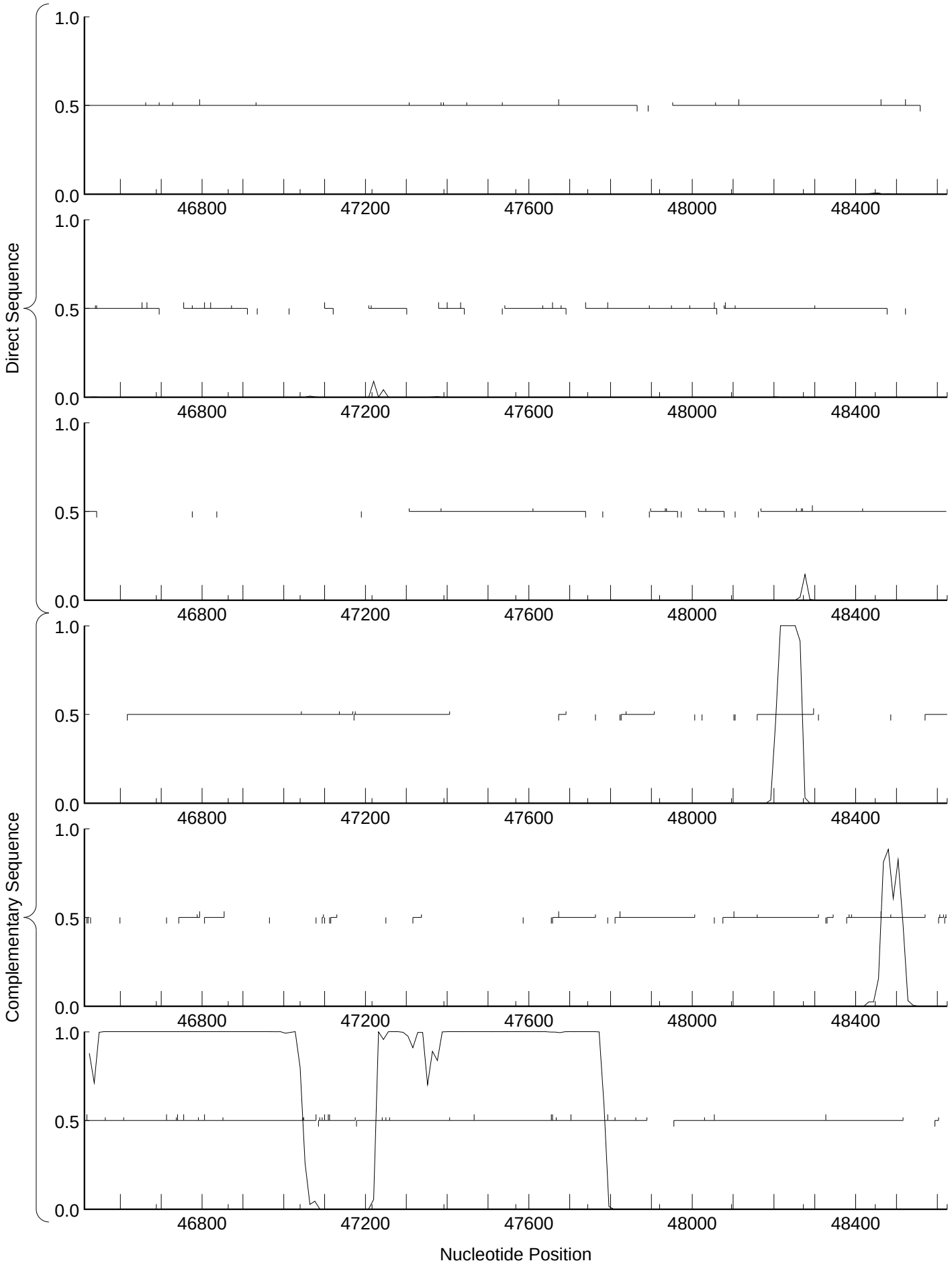
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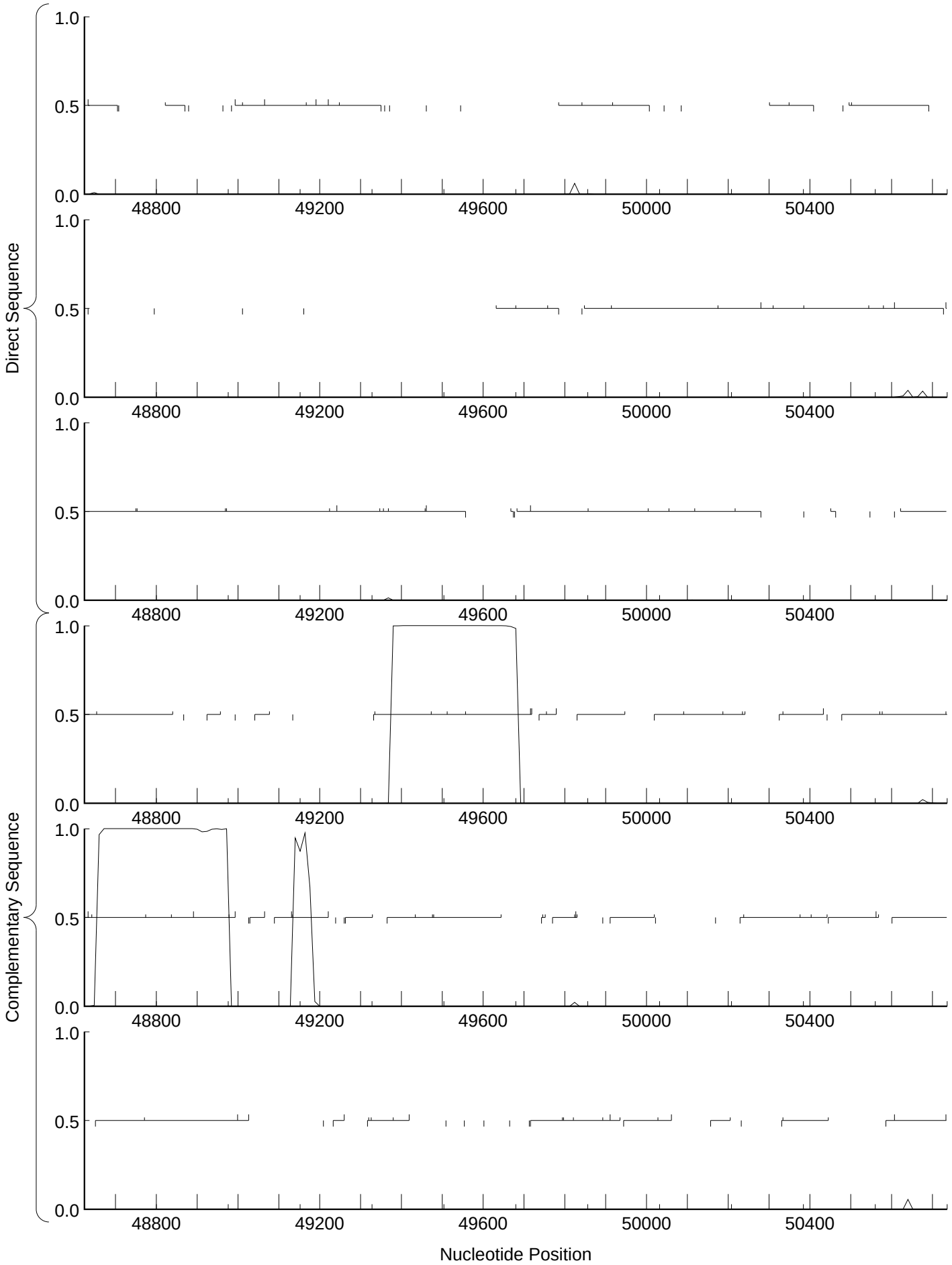
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omplete sequence, 50793 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 25/26



omplete sequence, 50793 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 26/26

