GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Amymech complete sequence, 53746 bp including 14-base 3’ overhang (TAAG
Analysis Date    :  3/29/24 at 2:48:22
Pages            :  27
Sequence Length  :  53746 bp
GC Content       :   67.45%

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,
```

MATRIX INFORMATION

```
Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

Order      :  4
```

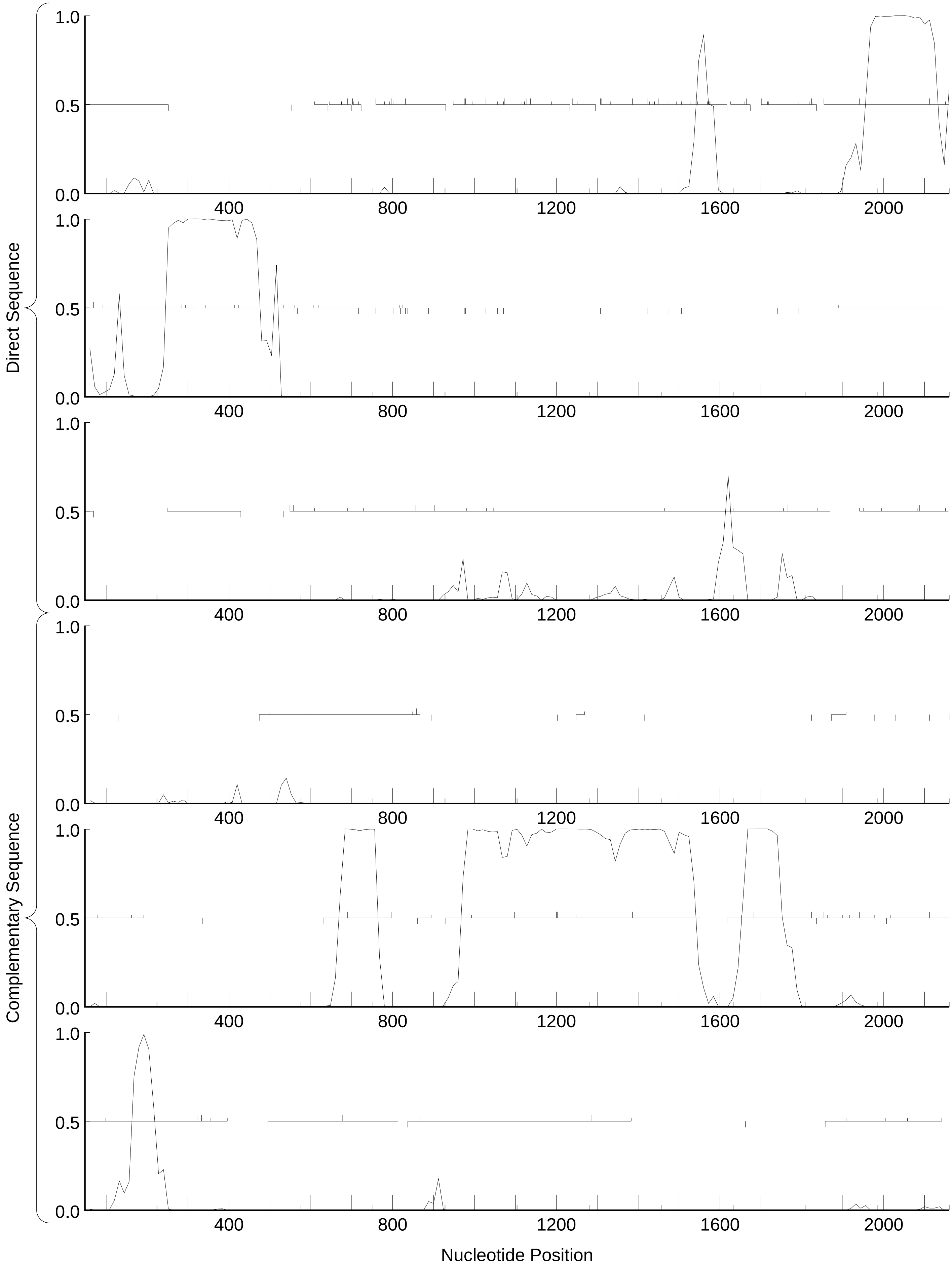
Send questions / comments to:

Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

Matrix notes & comments

Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu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0.0
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Nucleotide Position

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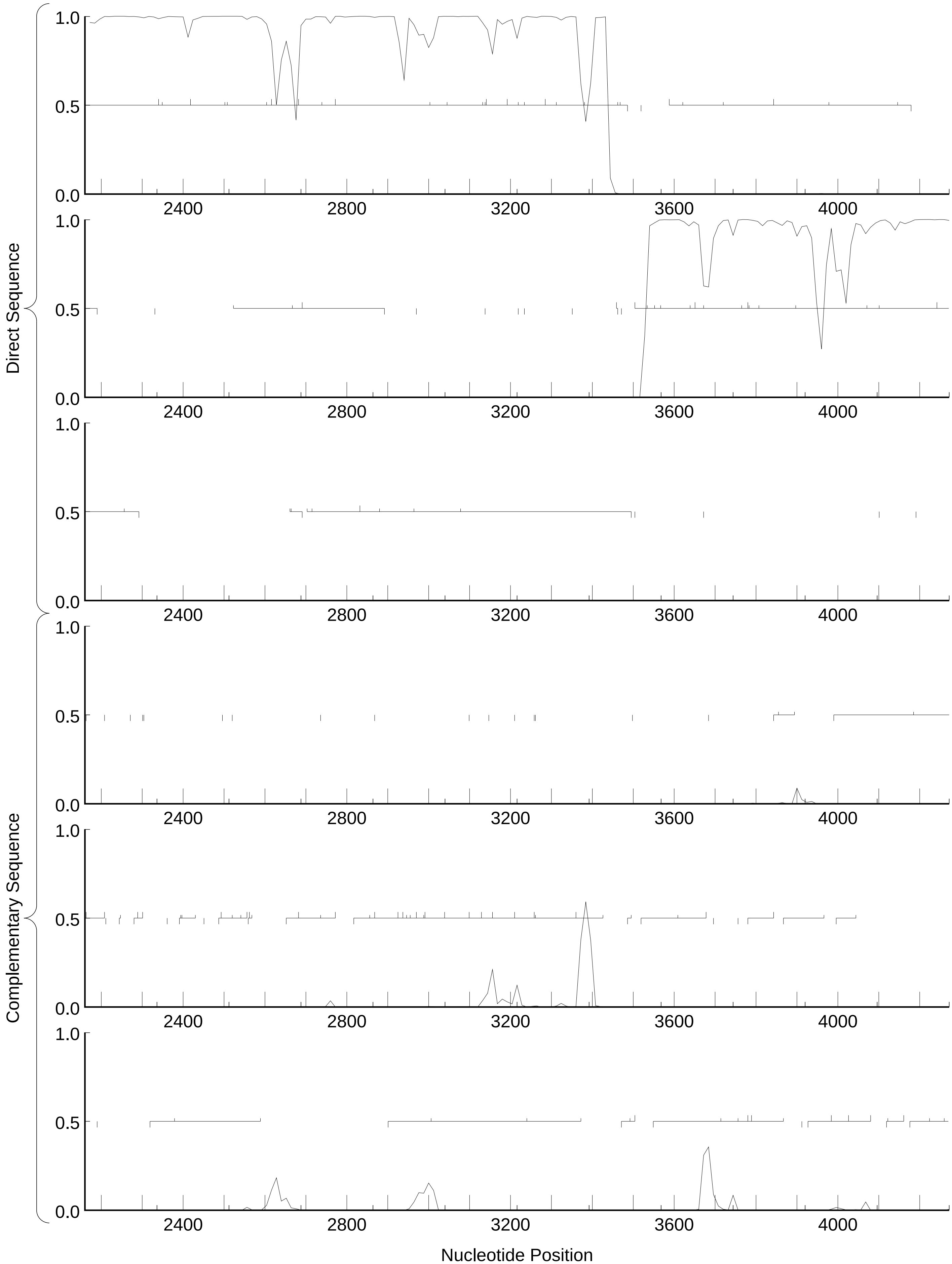

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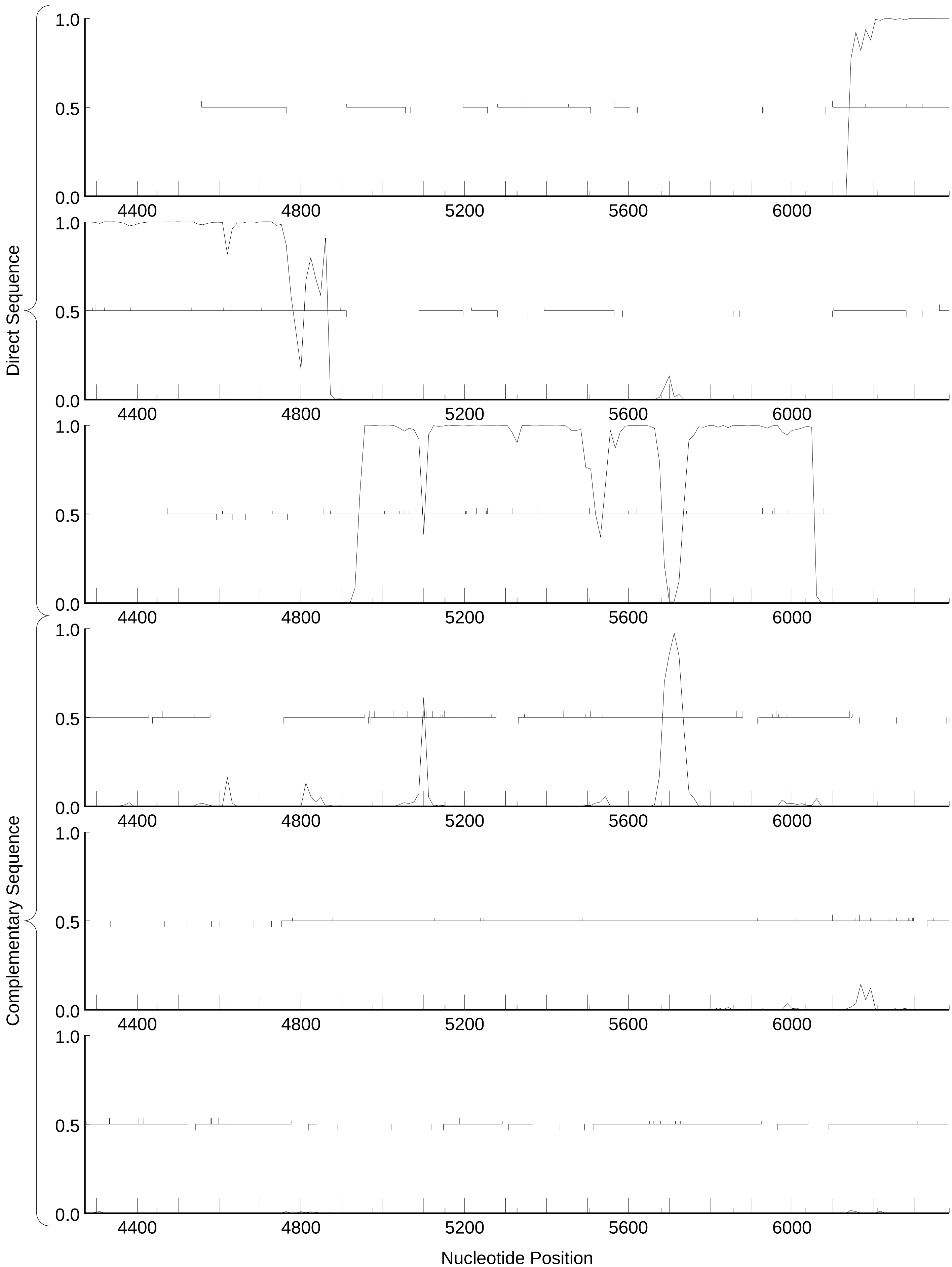

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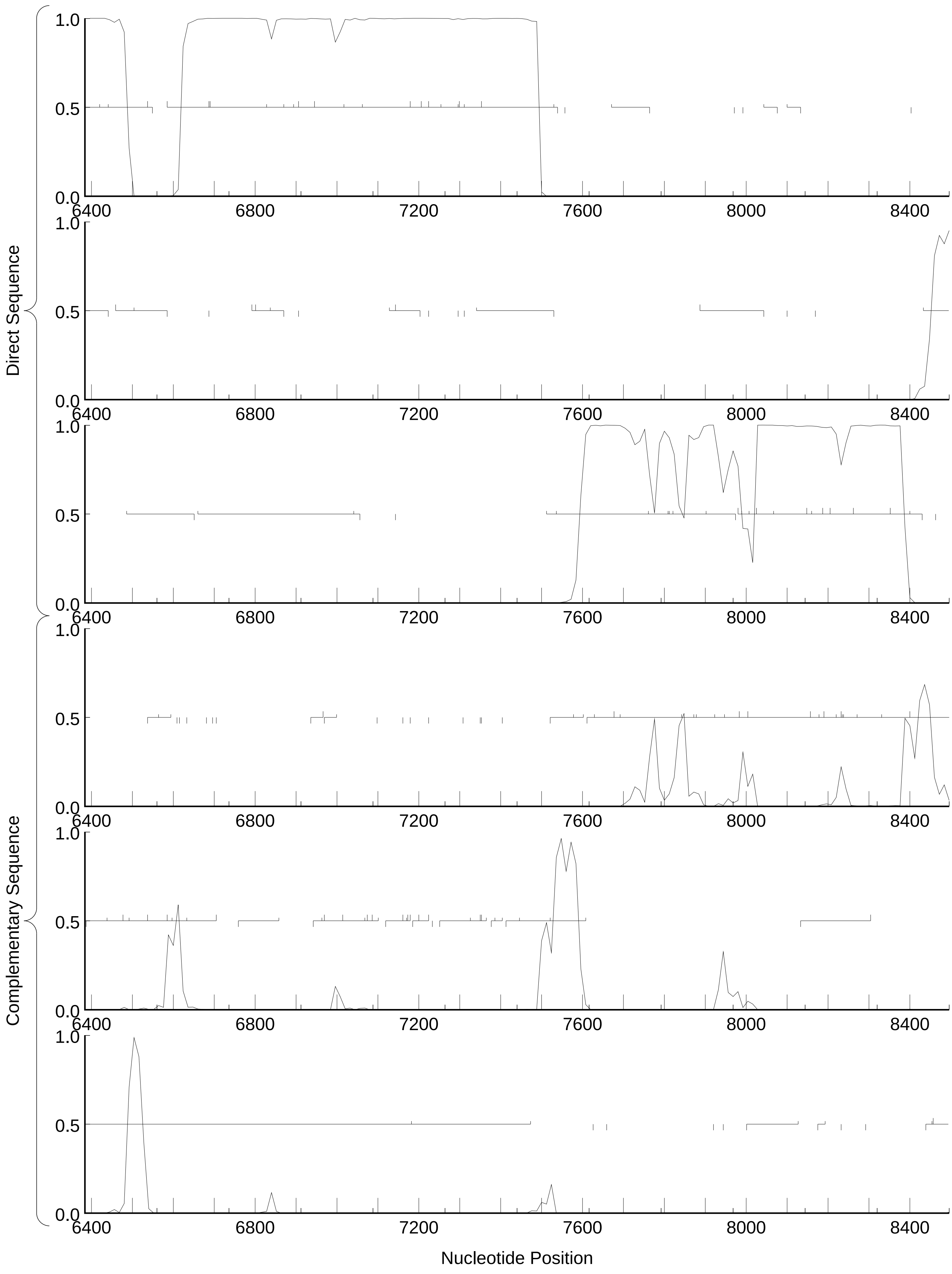

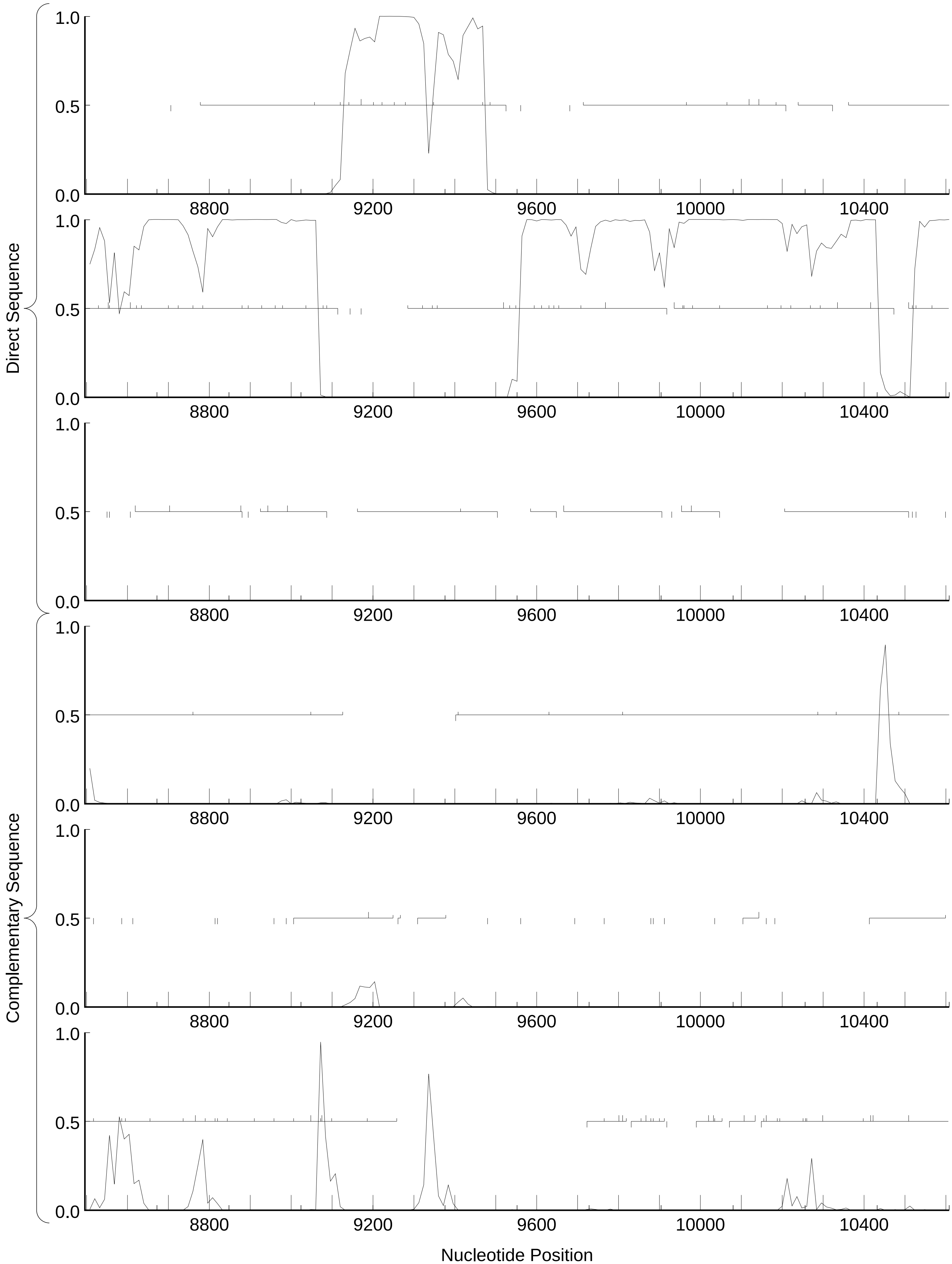
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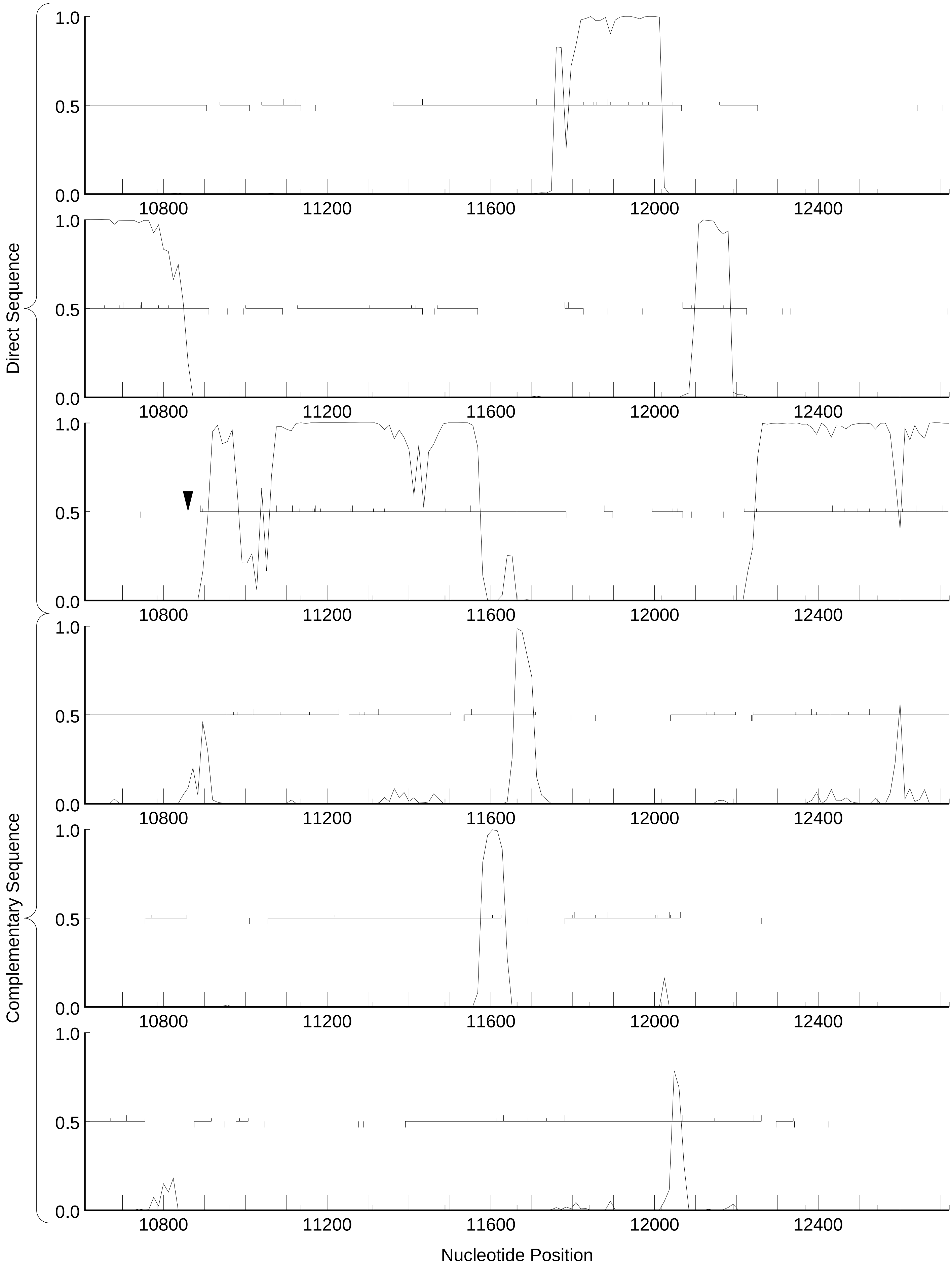
Nucleotide Position



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0.0
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11600
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Nucleotide Position

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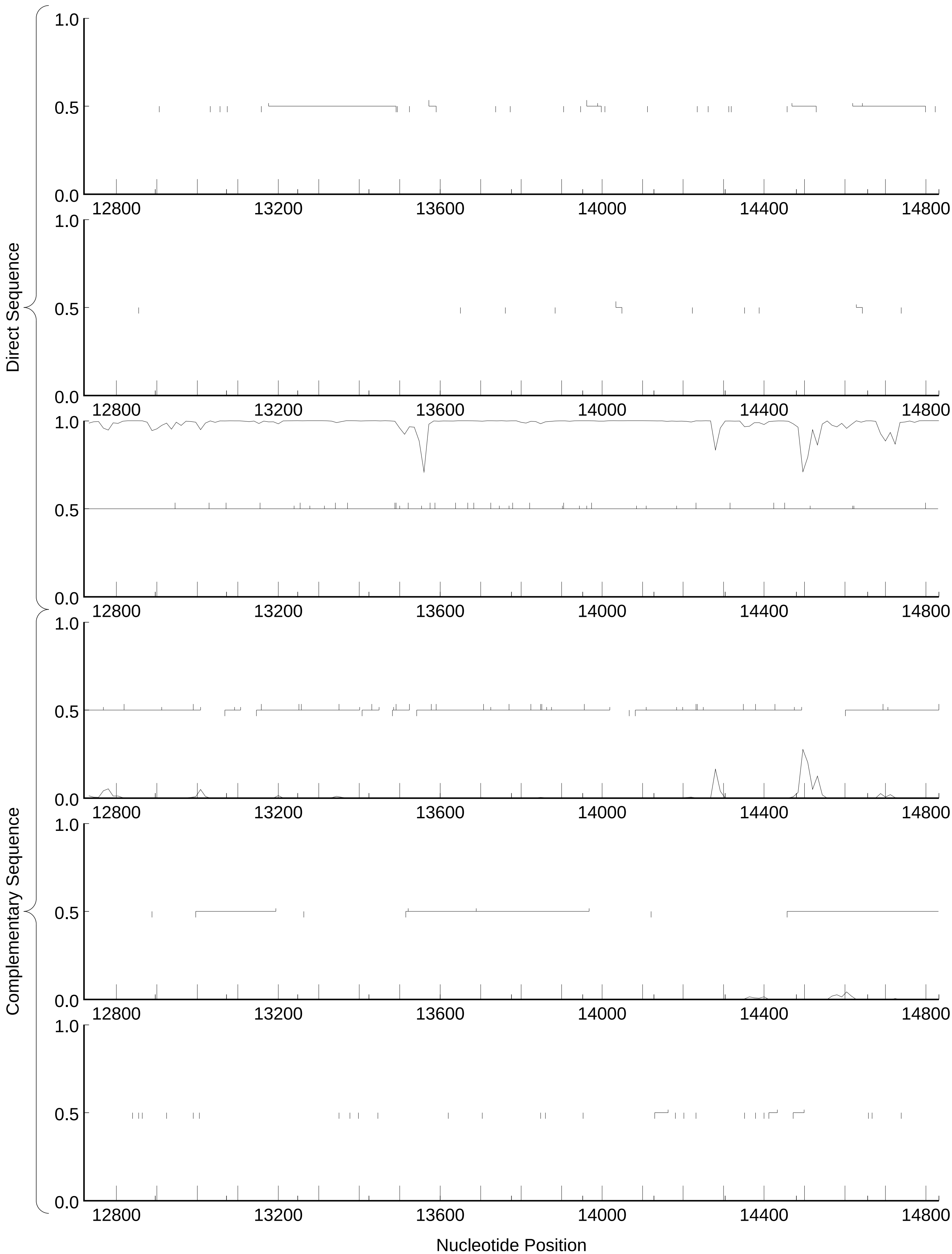

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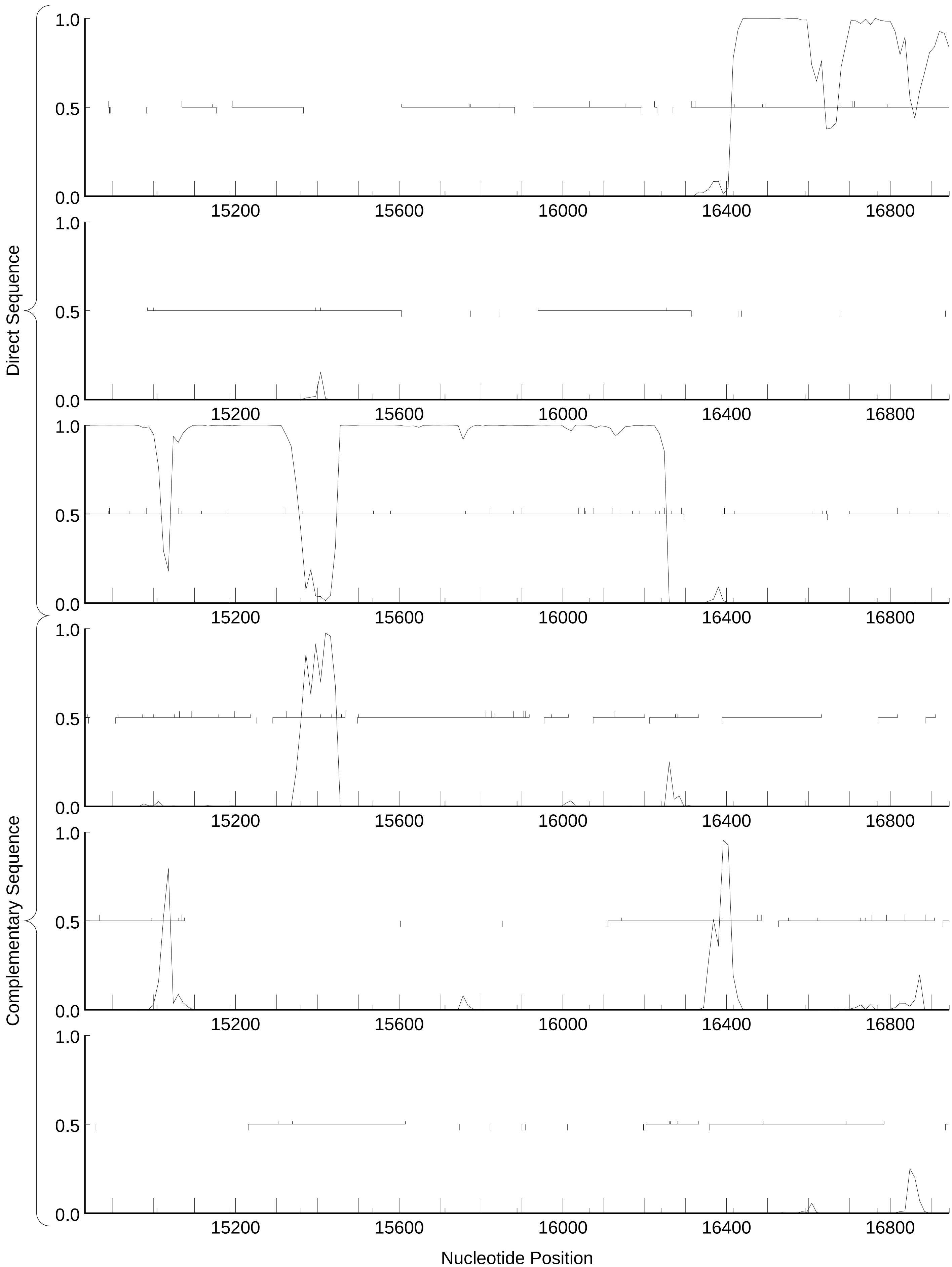

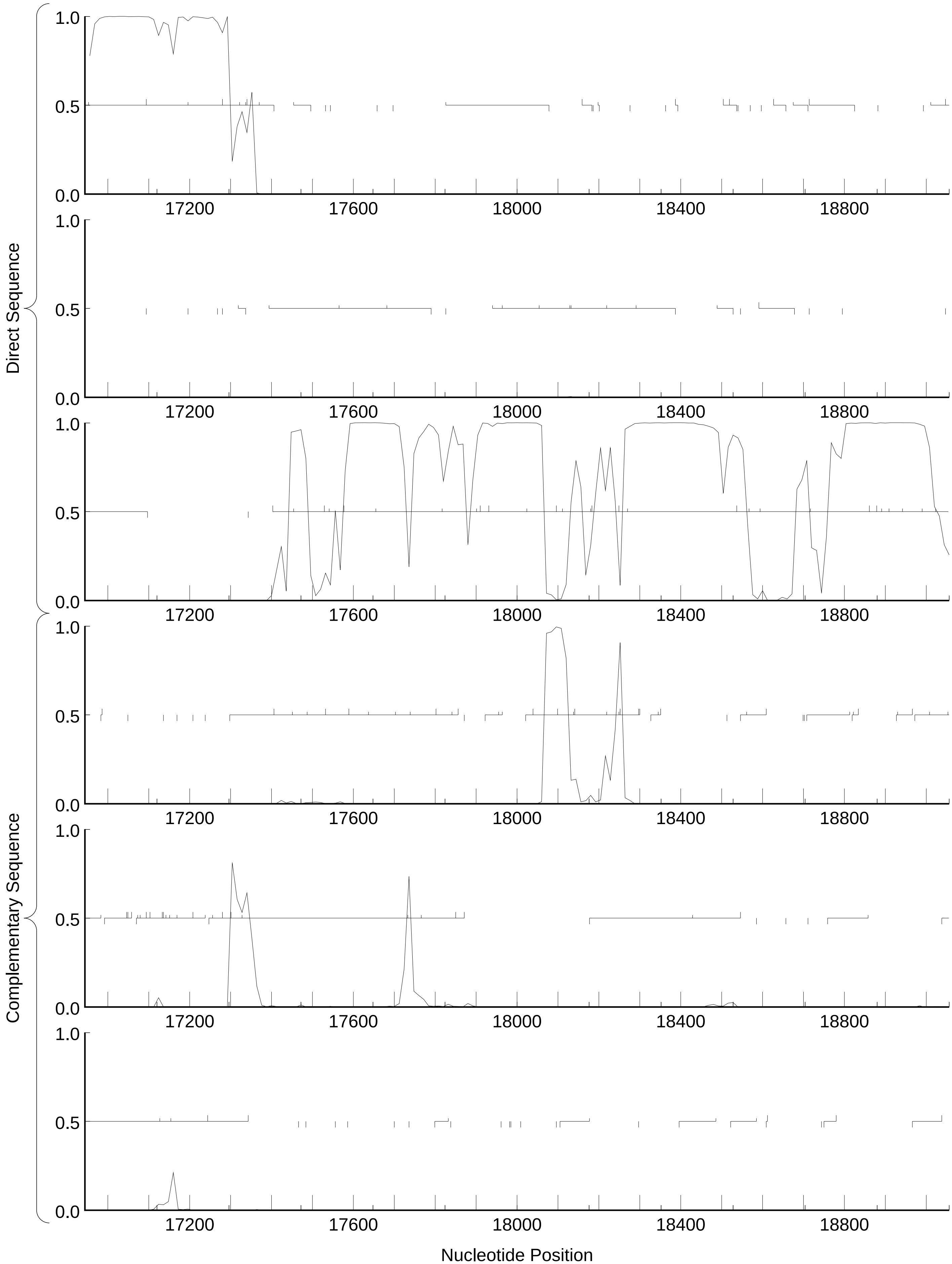

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18000
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Nucleotide Position

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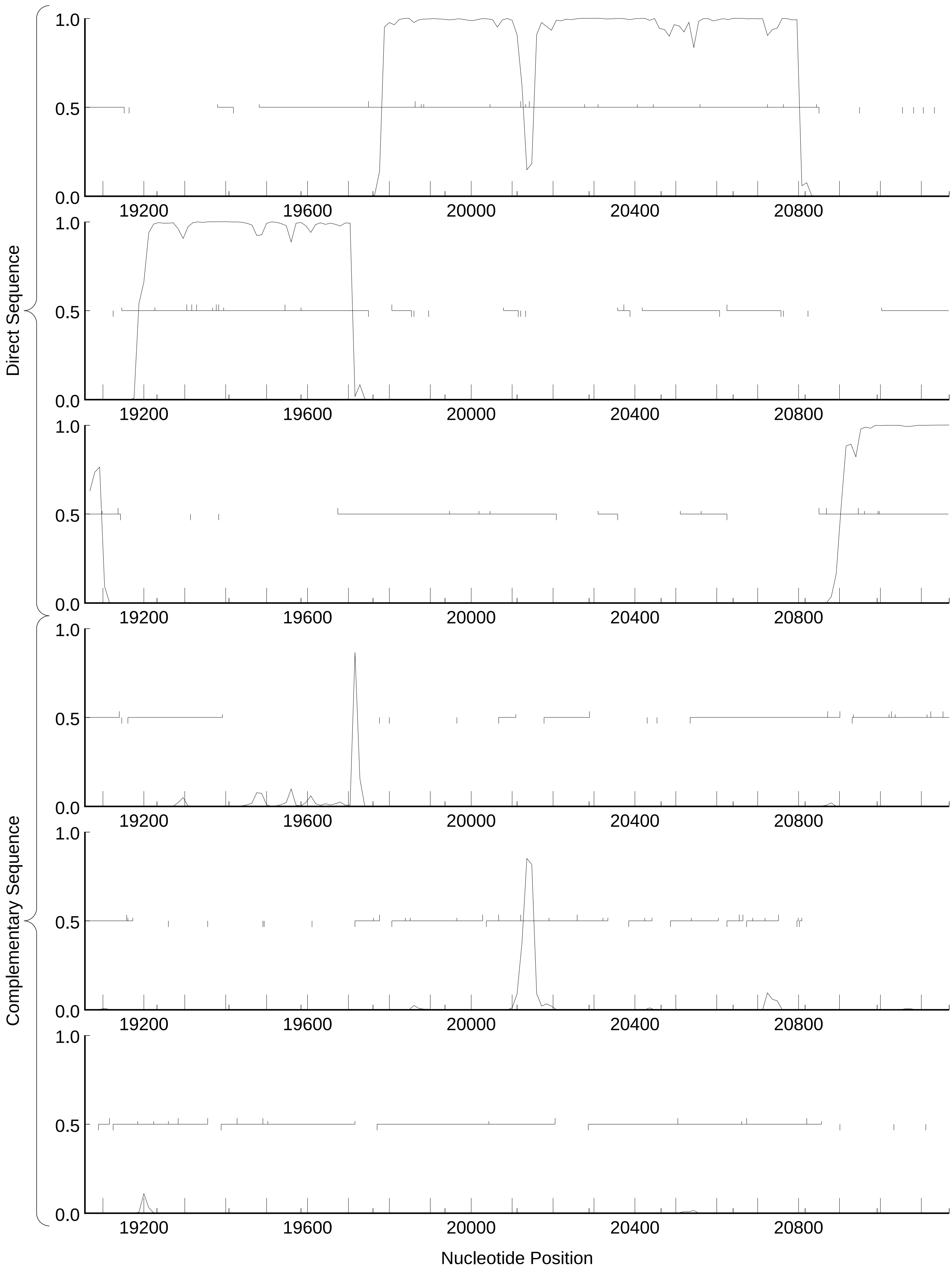

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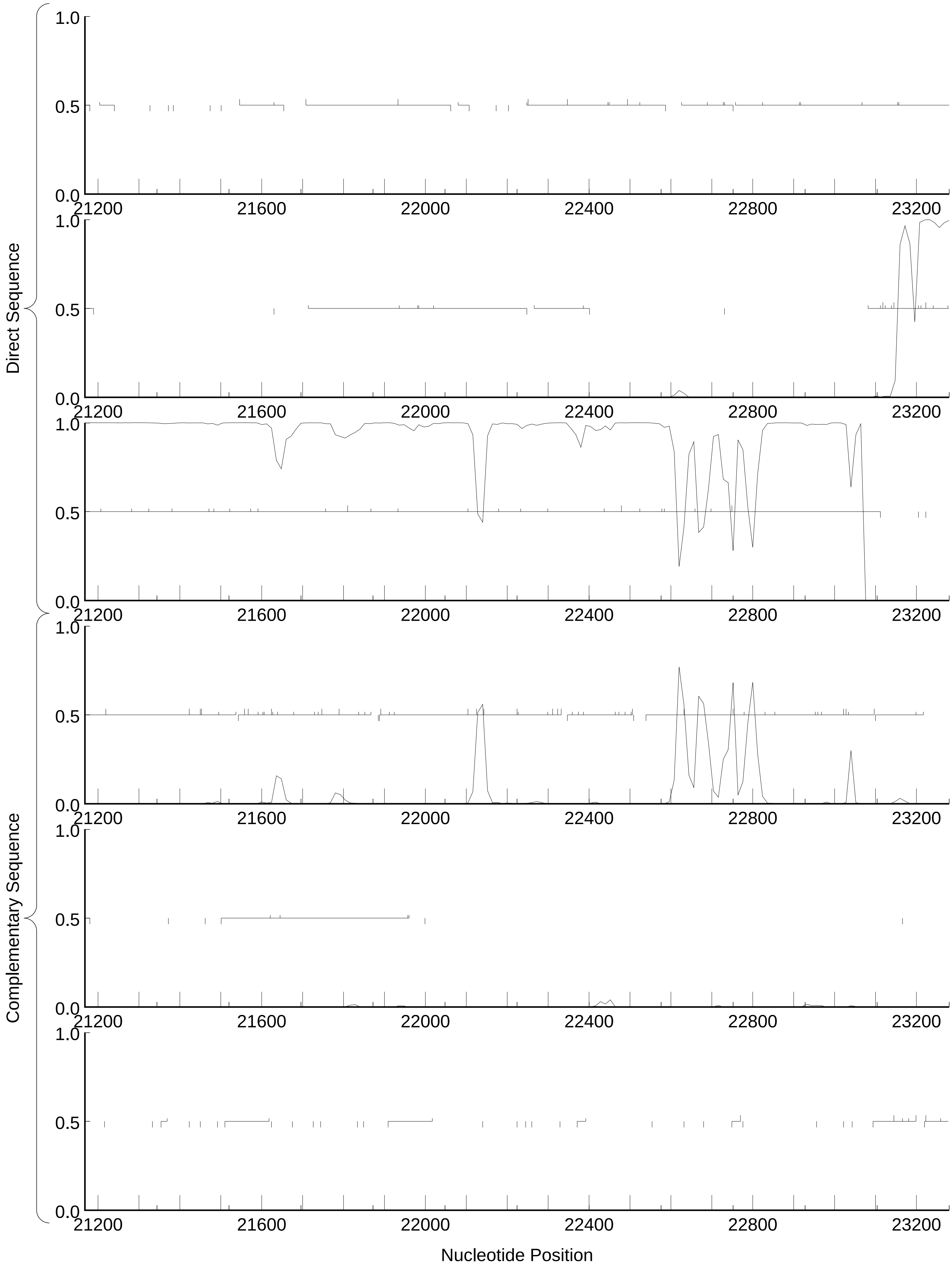

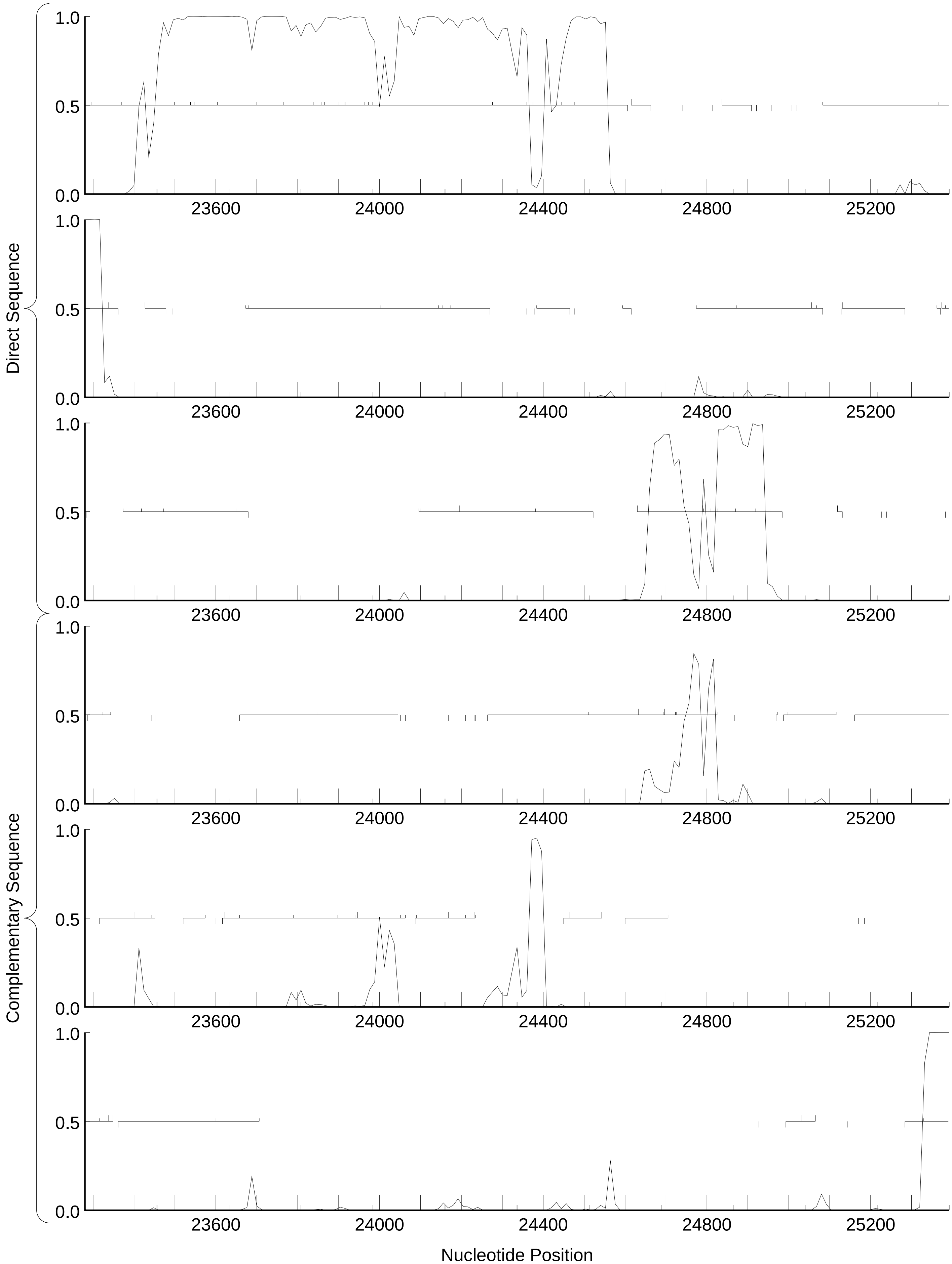
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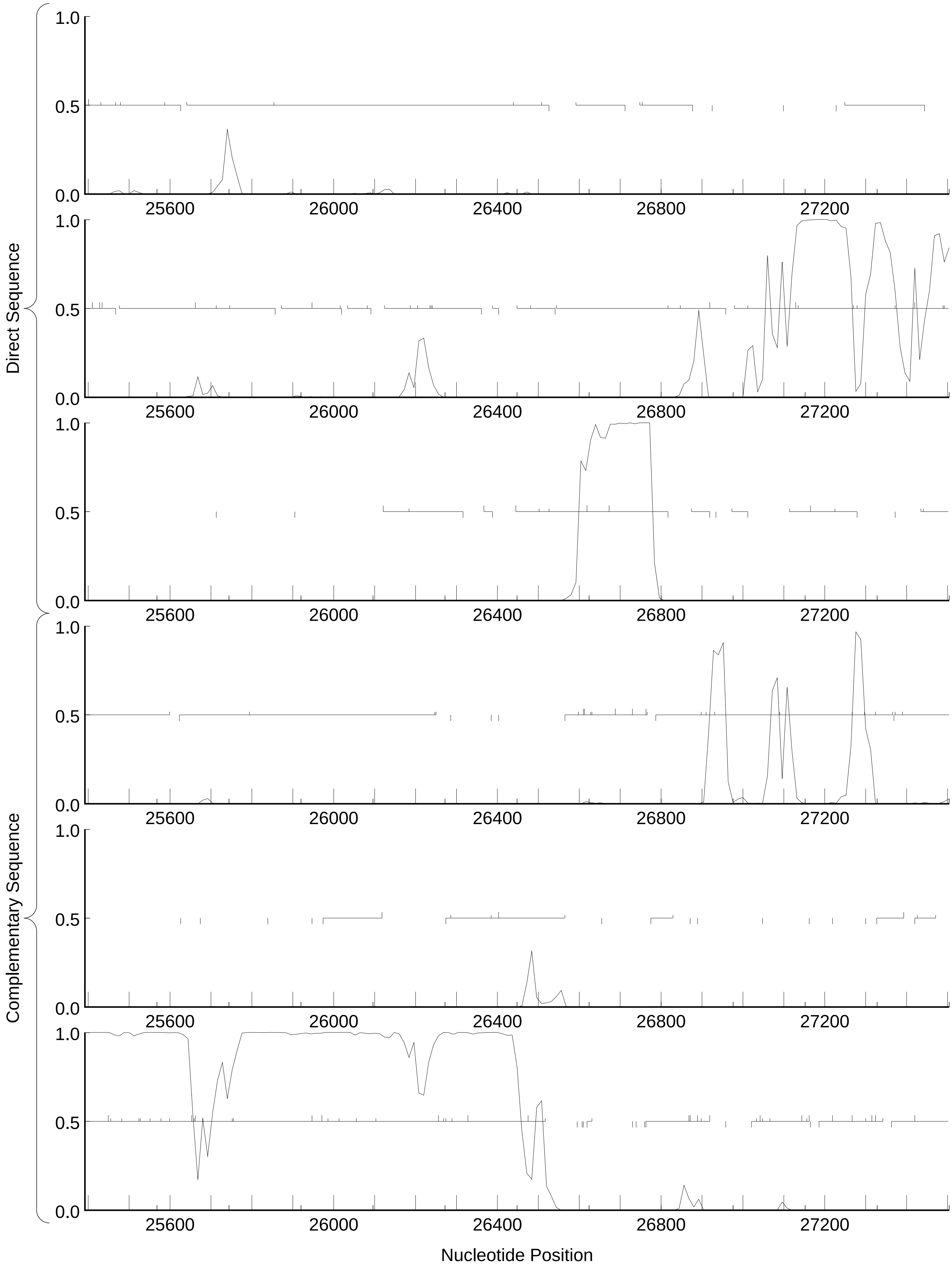

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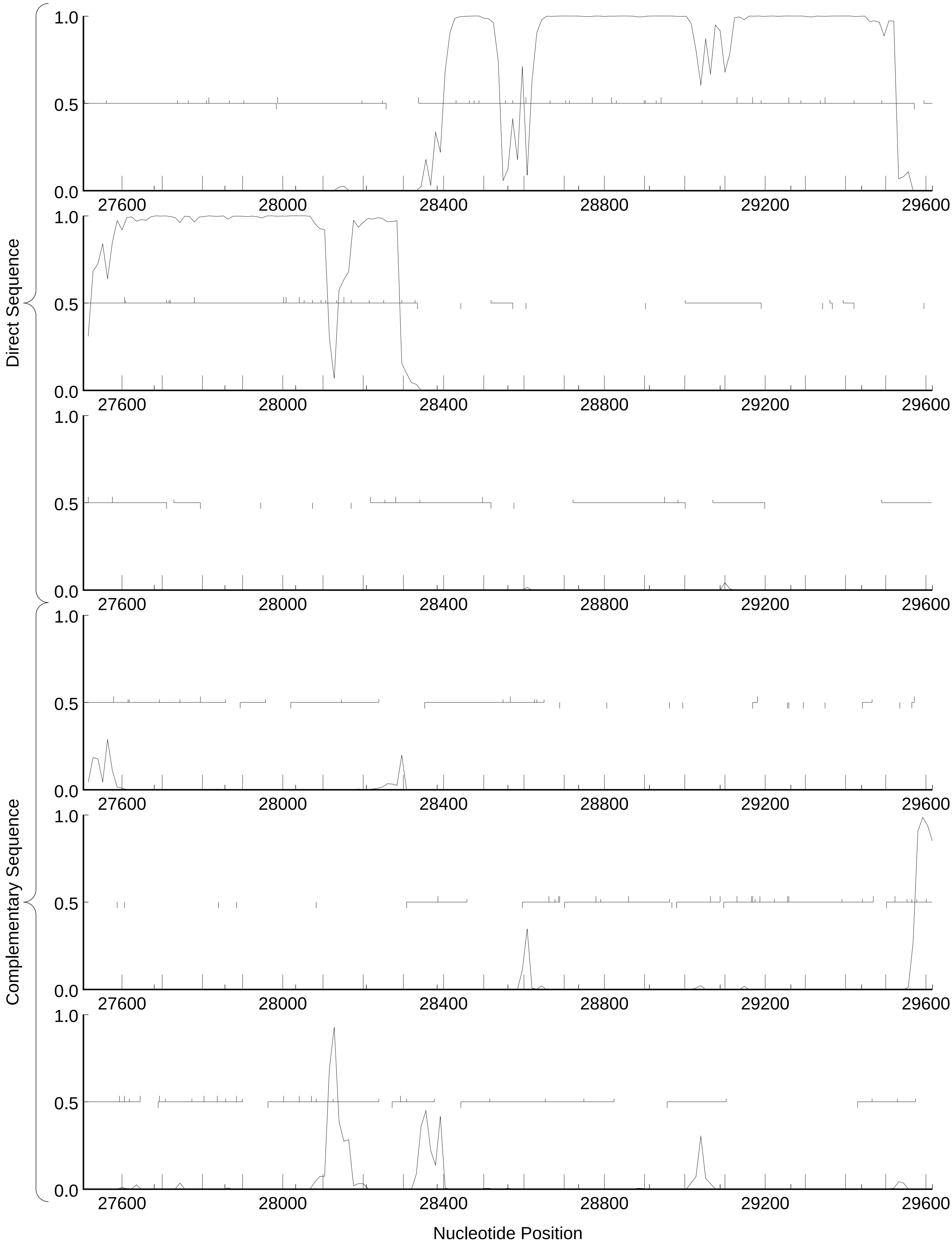
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27600
28000
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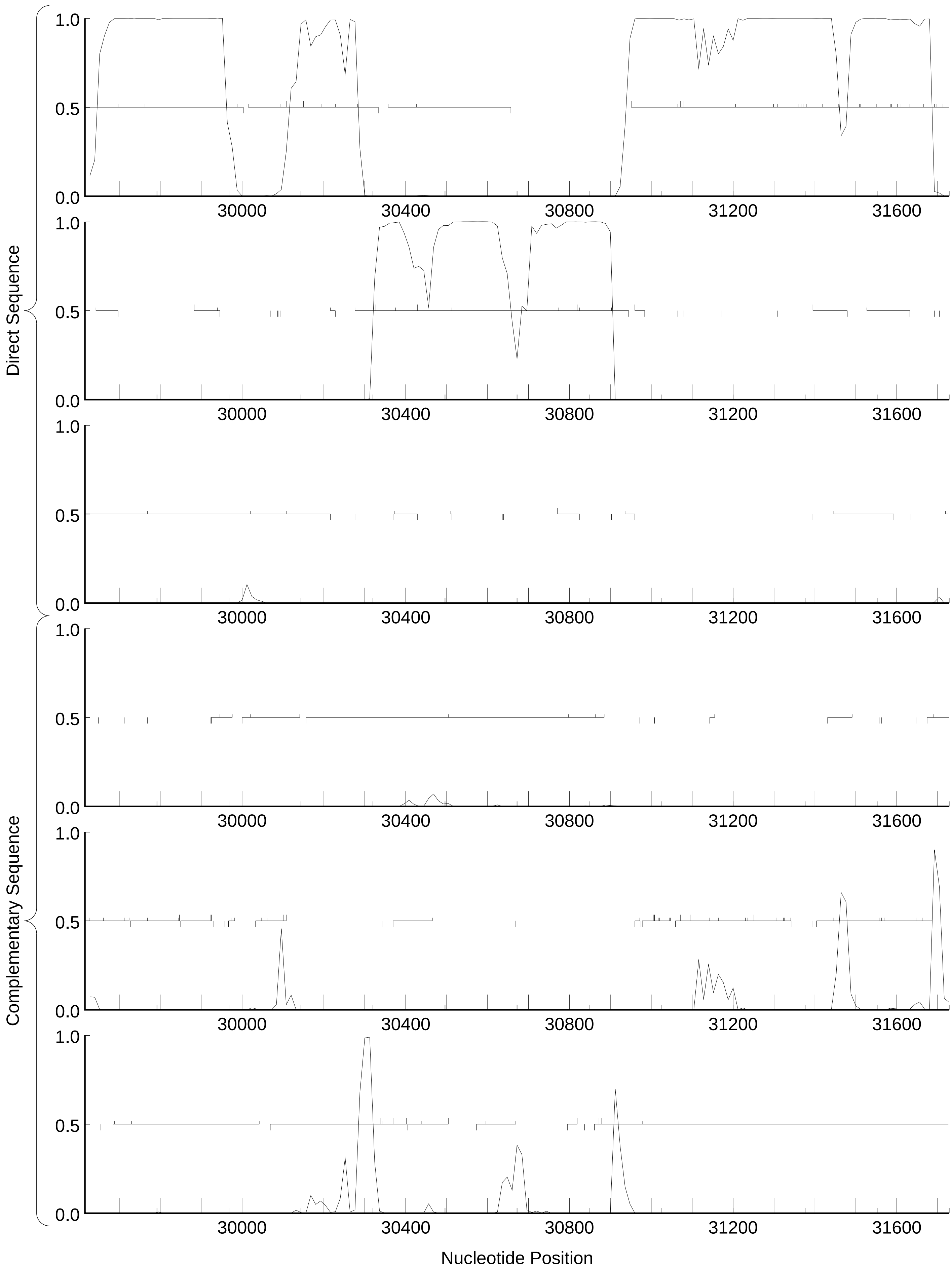
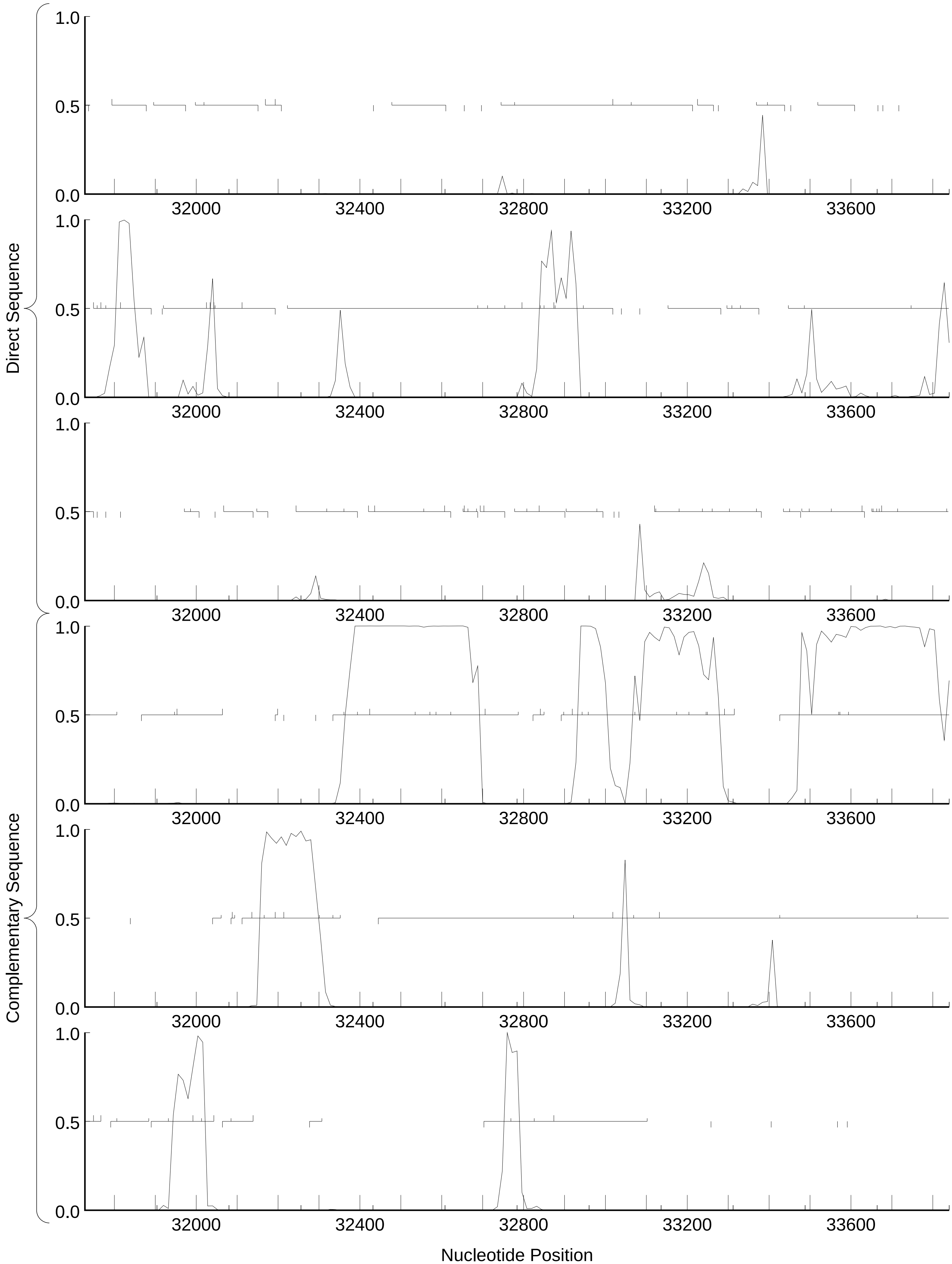
Nucleotide Position

blete sequence, 53746 bp including 14-base 3’ overhang (TAAGCCGCGCGGTA), Cluster Q, Order 4, Window 96, Step 12, 16/27

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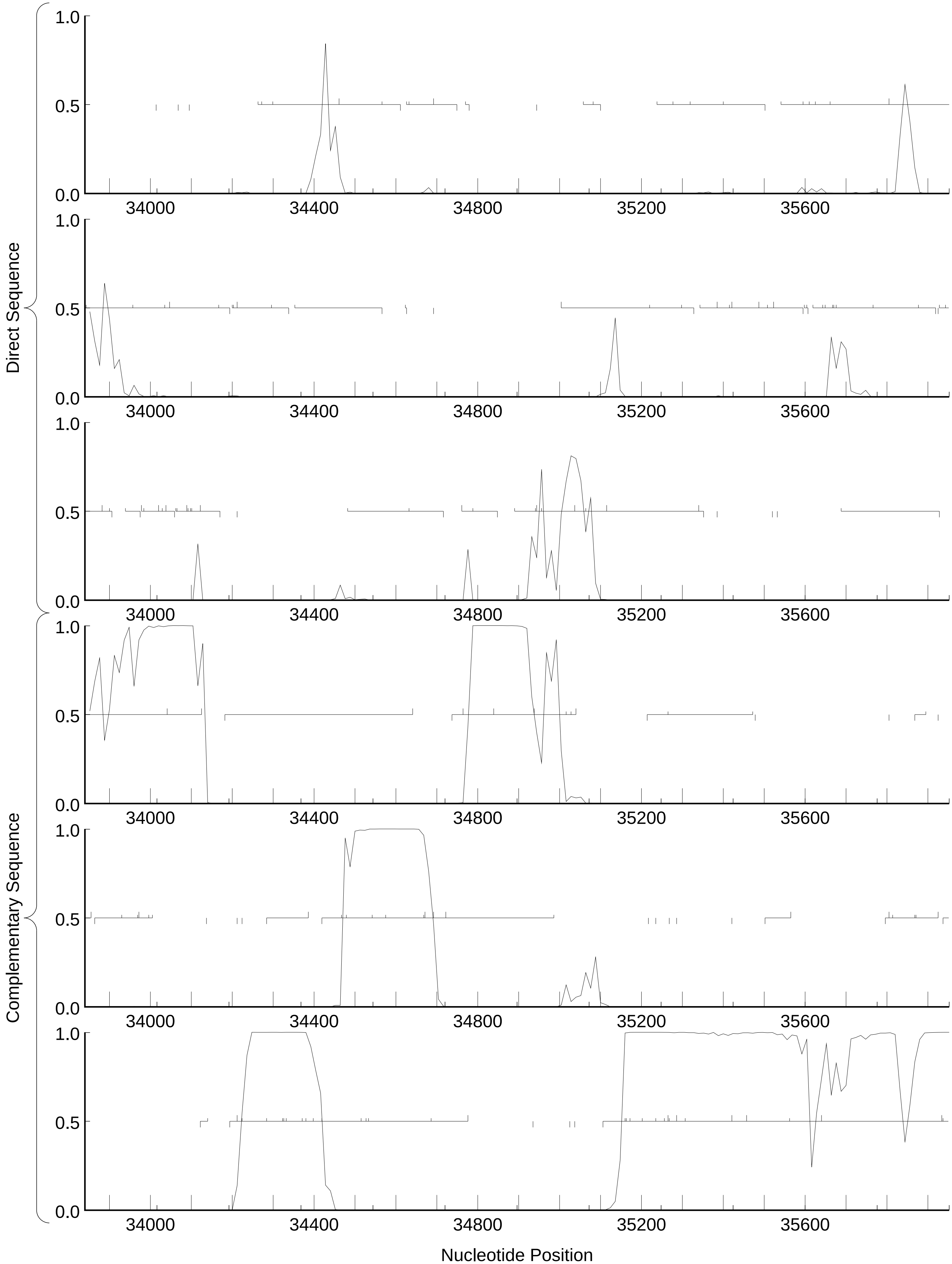
Nucleotide Position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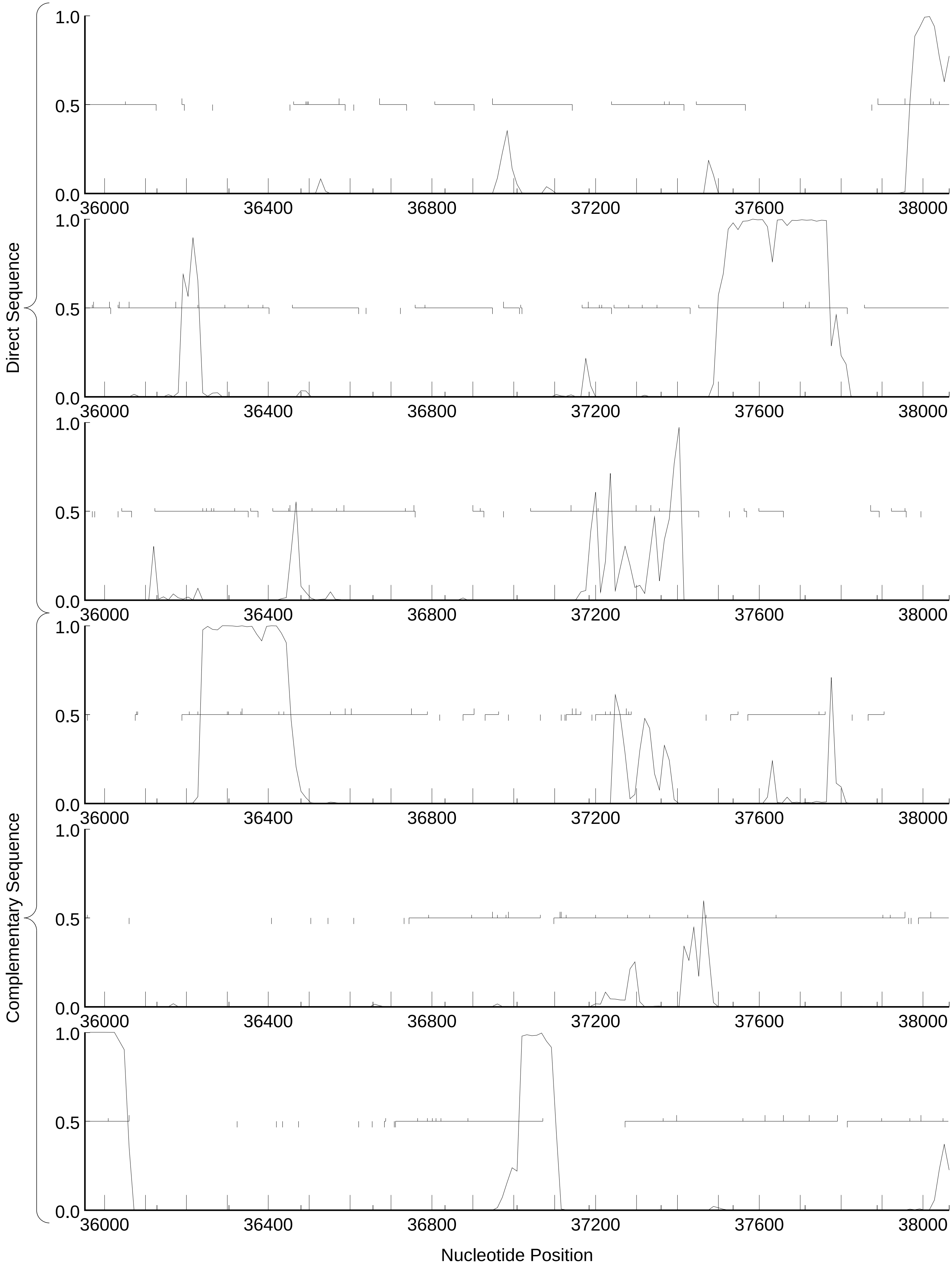
Nucleotide Position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Nucleotide Position

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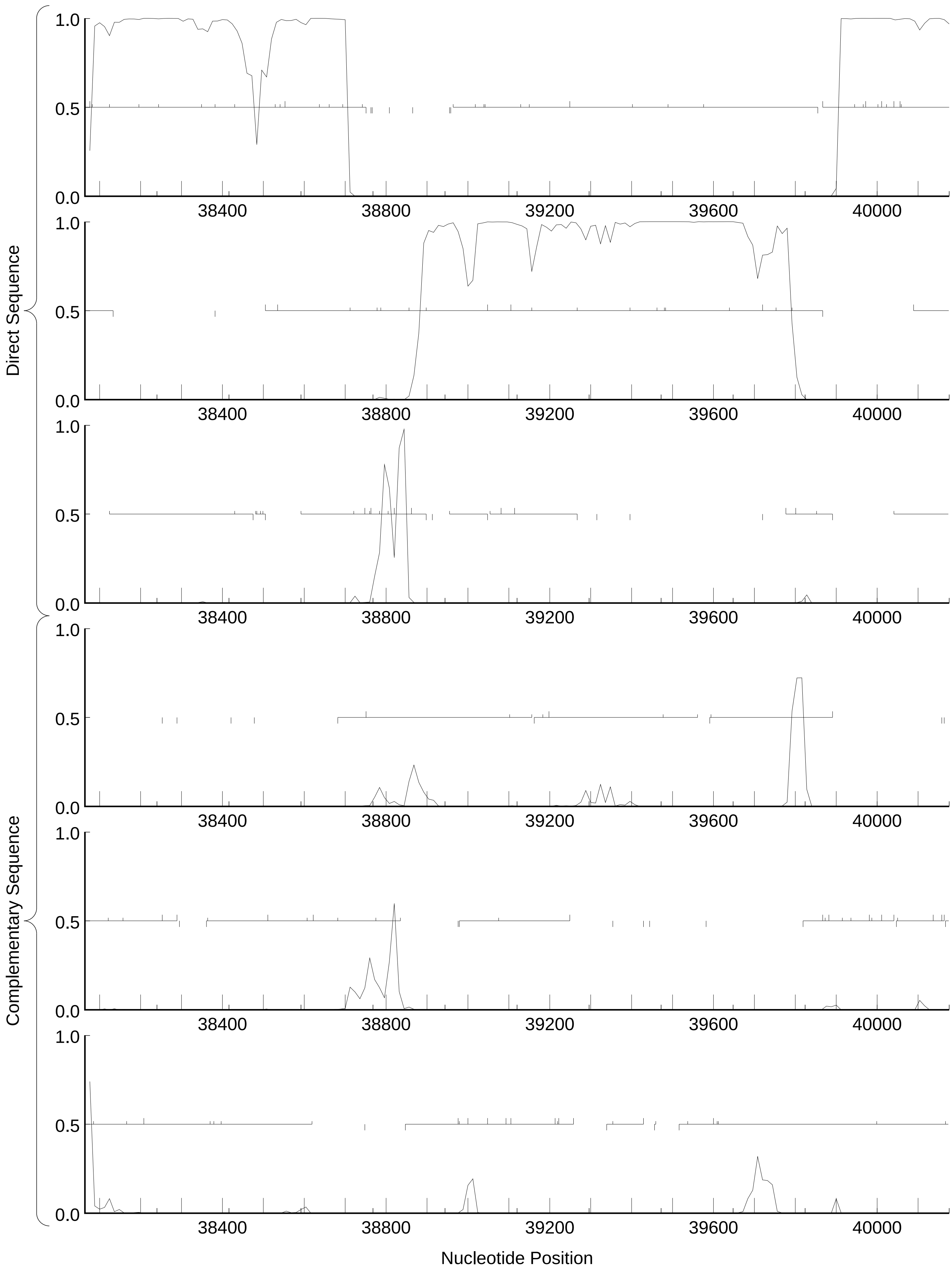

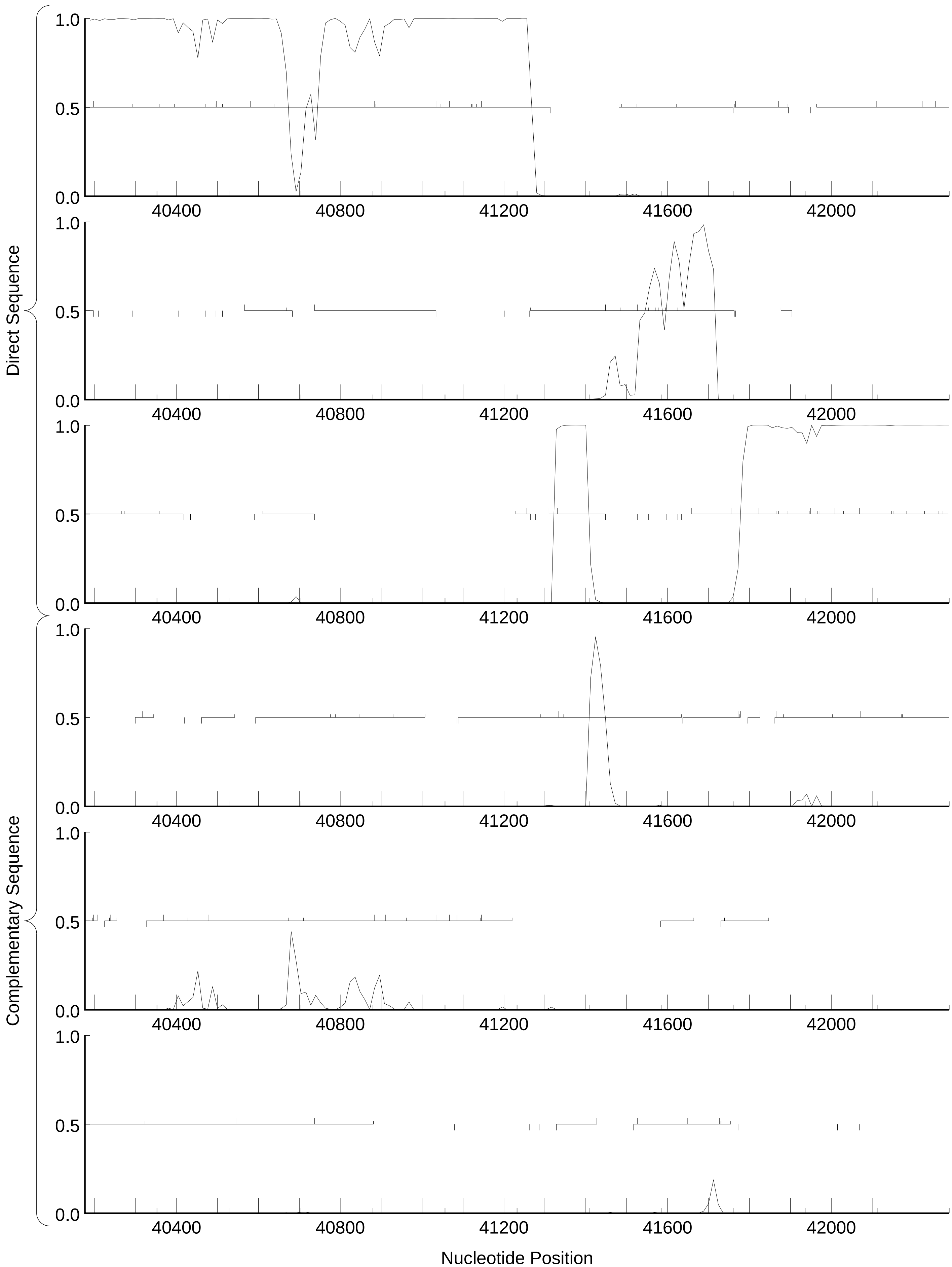
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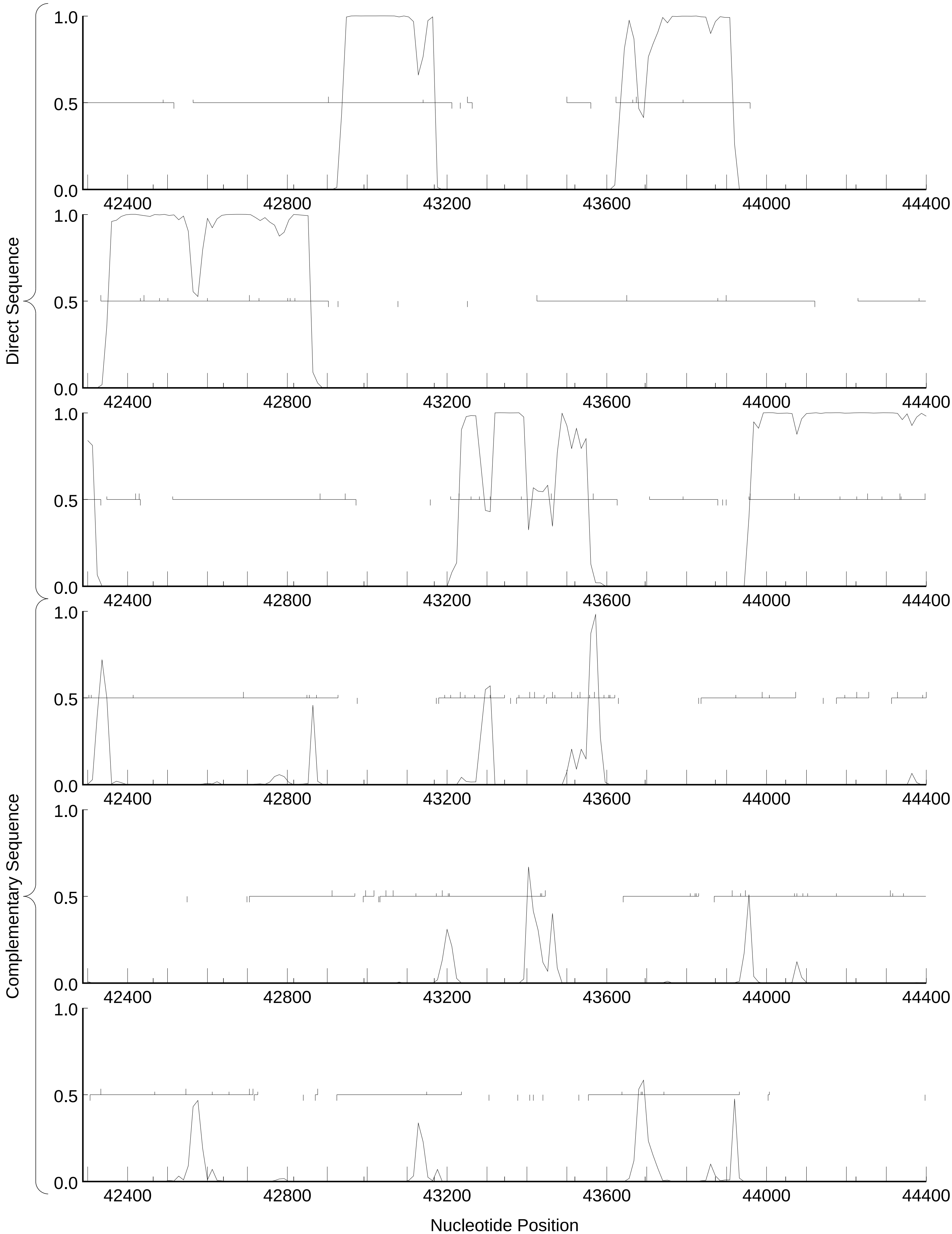

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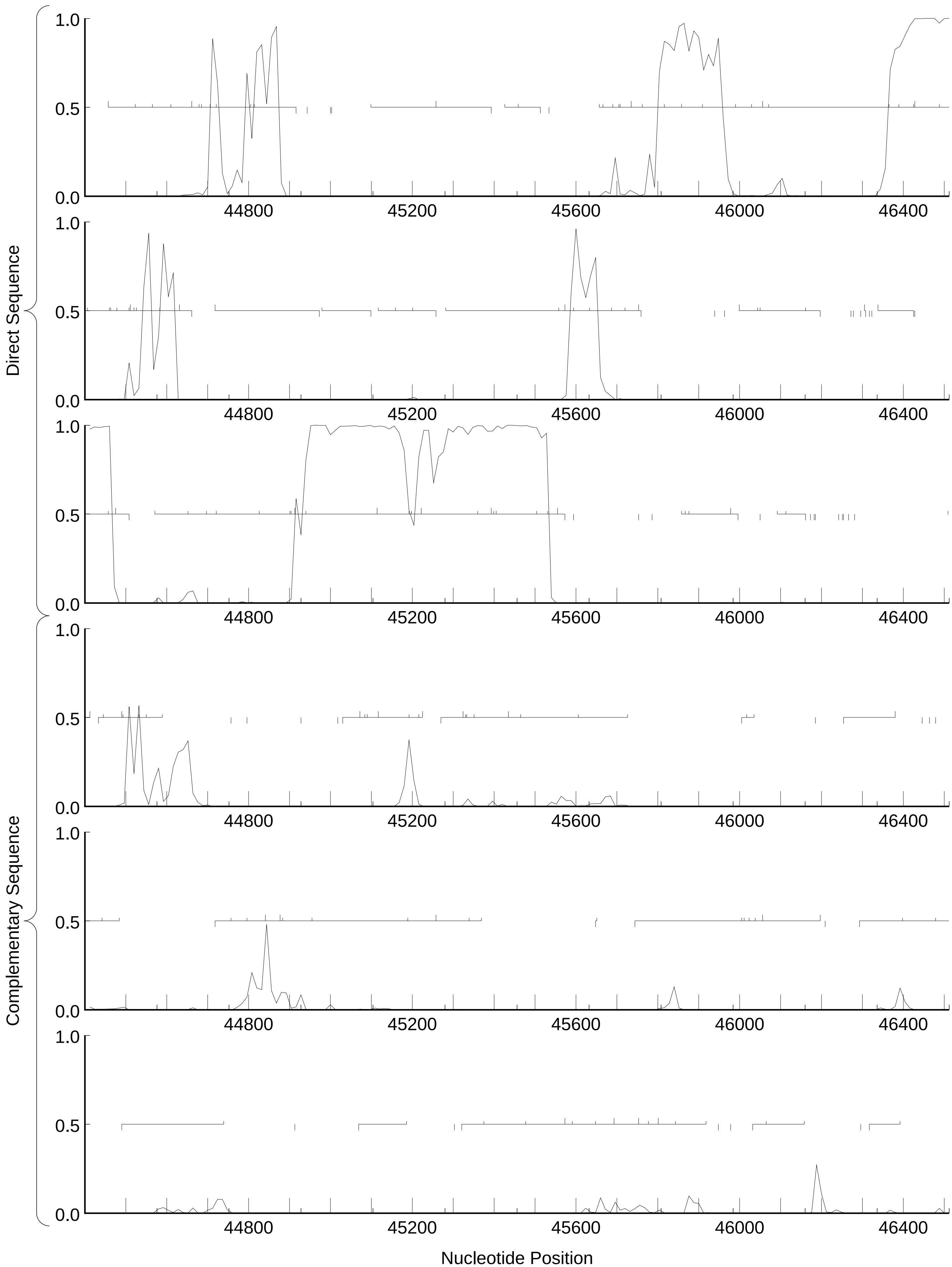

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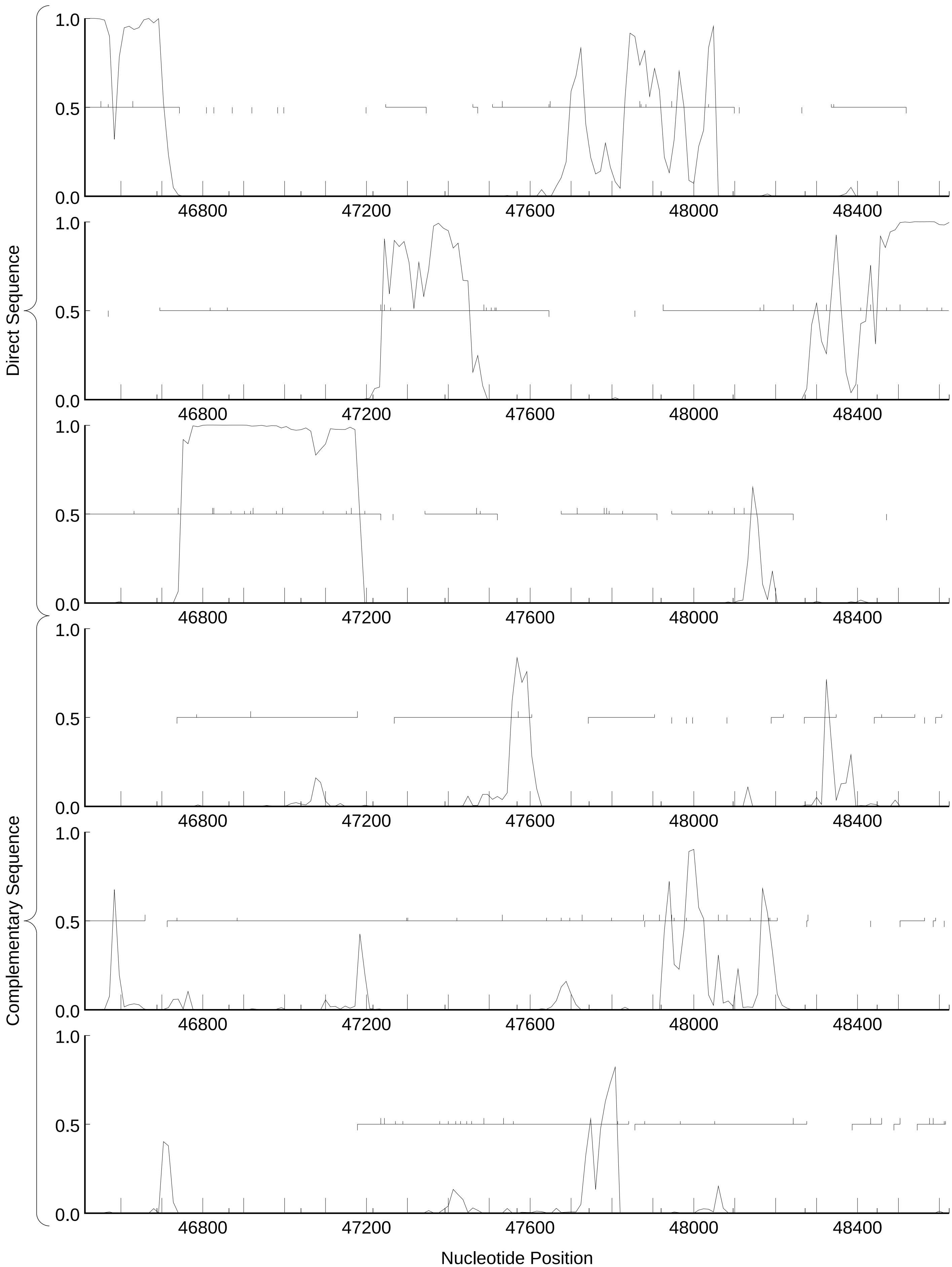

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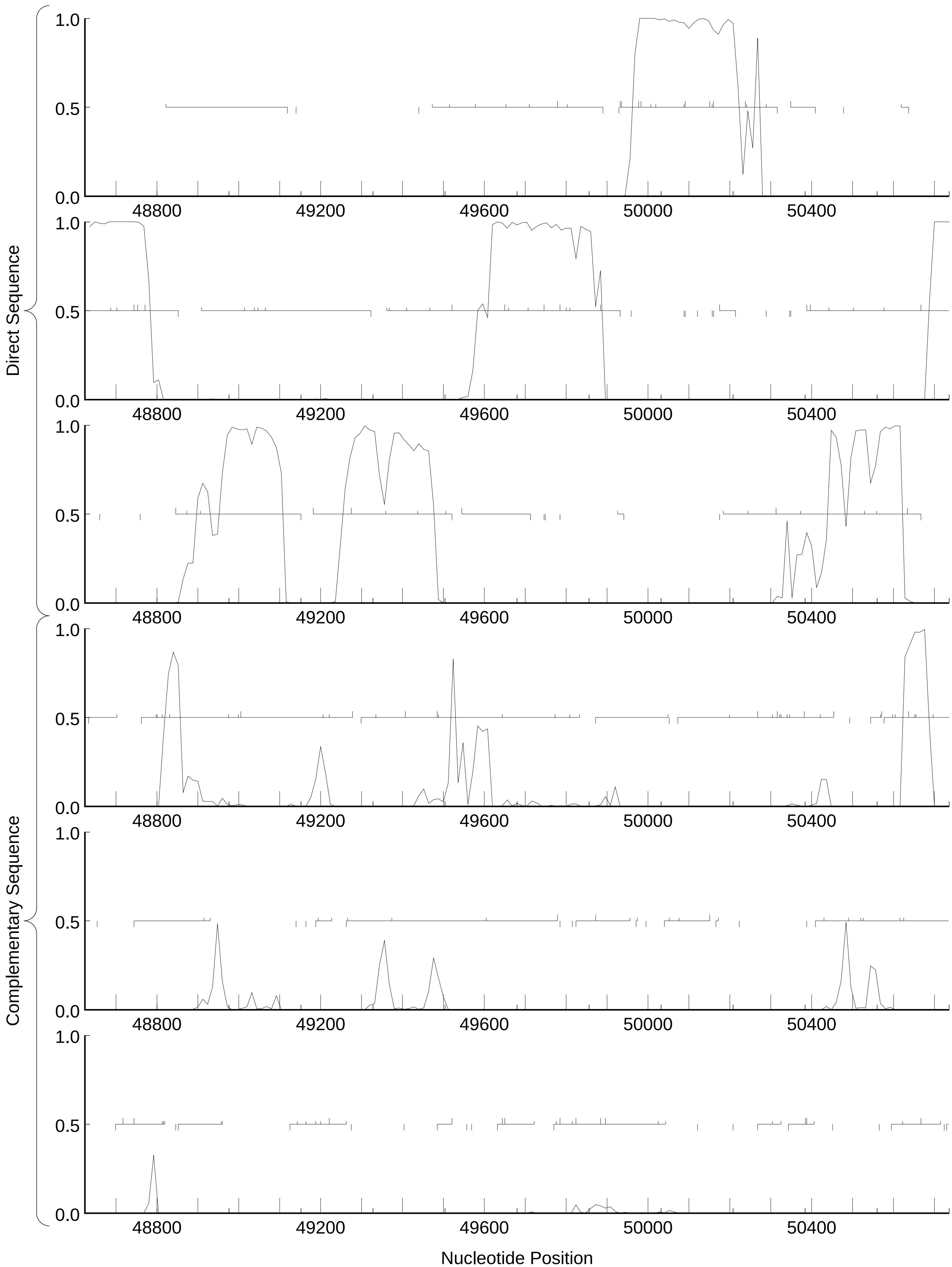

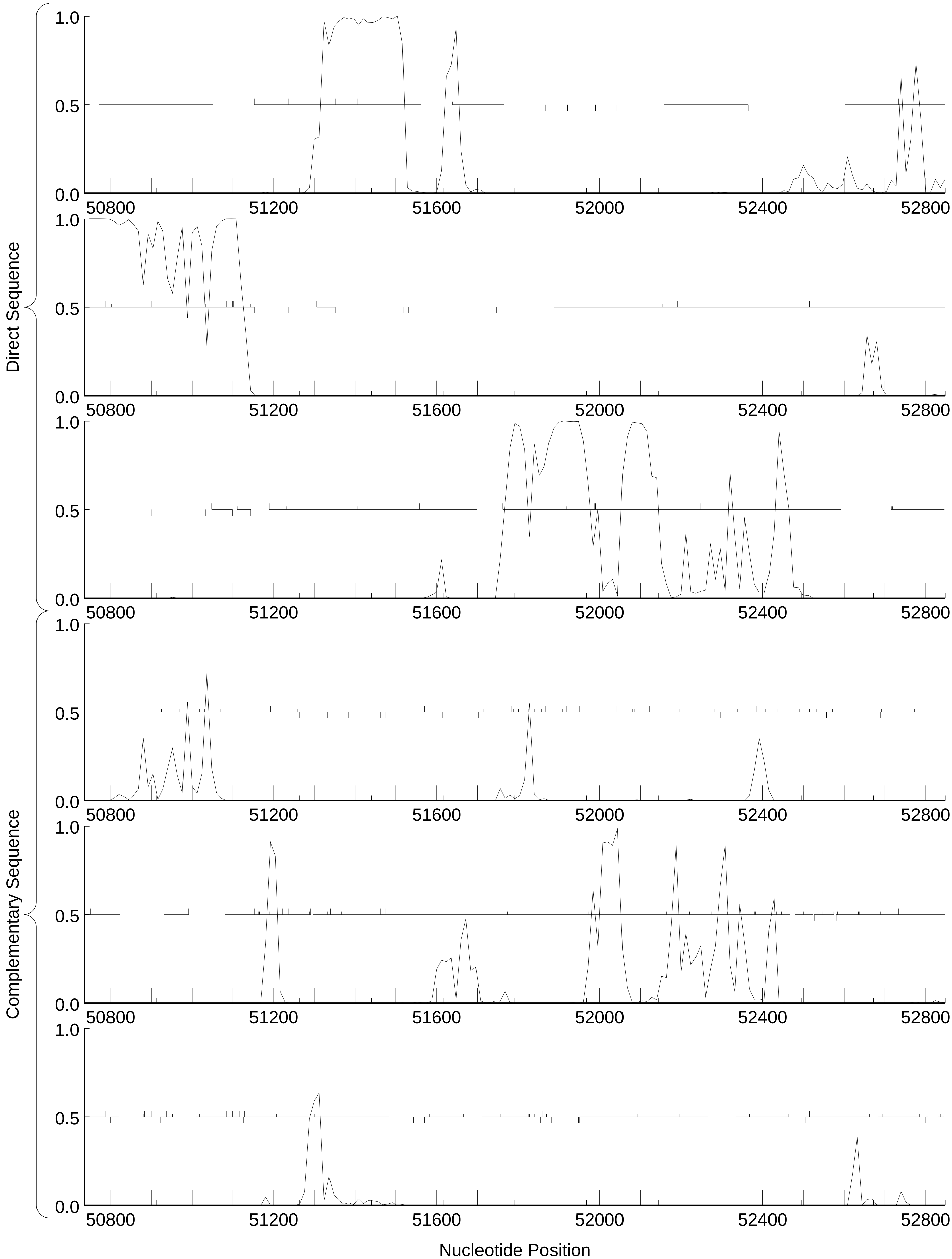
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Complementary Sequence
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Nucleotide Position

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