GeneMark

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

PROGRAM INFORMATION

```
Sequence        :  Mycobacterium phage NaSiaTalie complete sequence, 52920 bp including 10-base 3’ overhang (C
Analysis Date   :  3/28/24 at 16:11:31
Pages           :  27
Sequence Length :  52920 bp
GC Content      :   63.40%

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 notes & comments

Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:23:03 2014

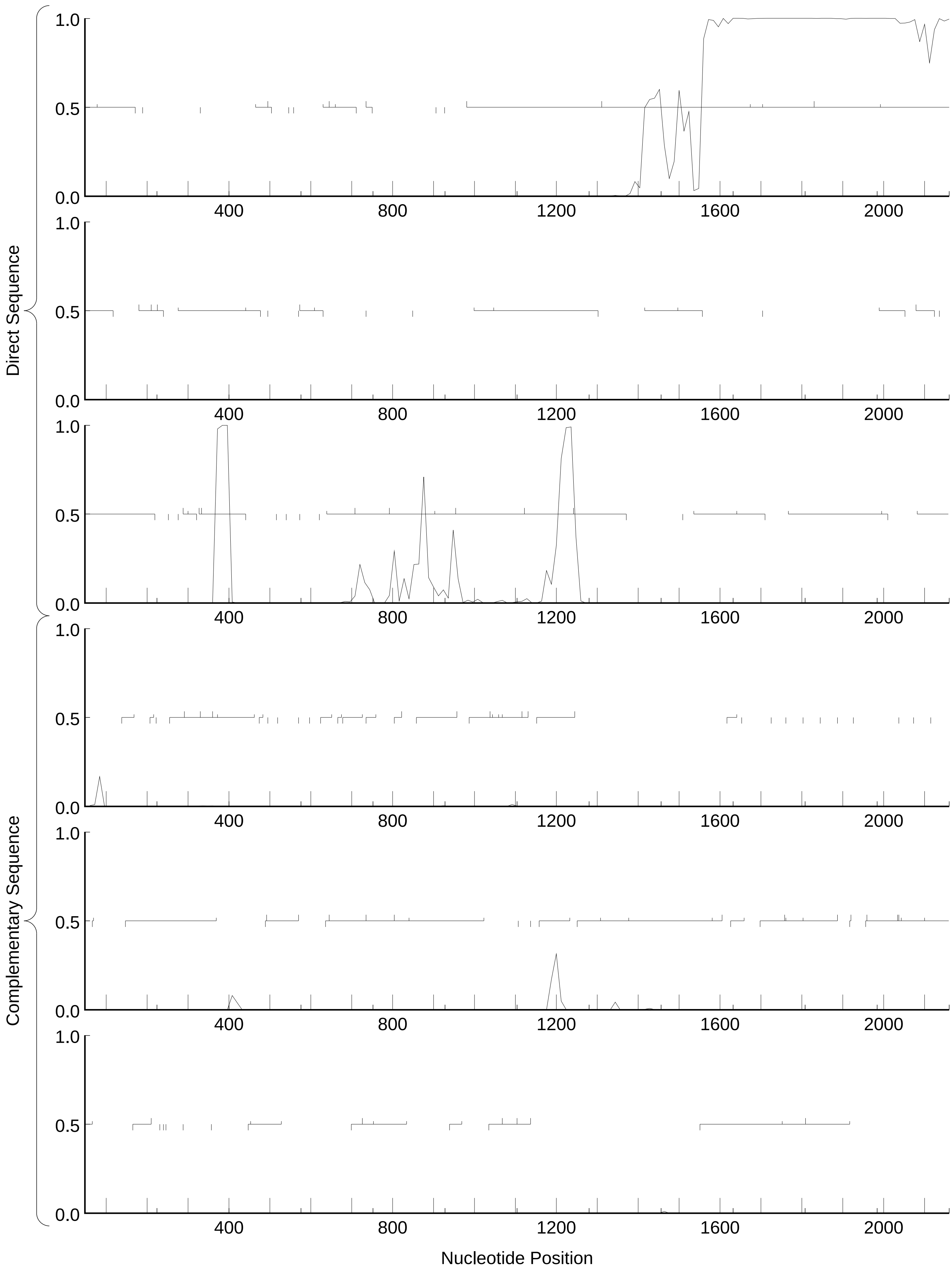
MATRIX INFORMATION

```
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

ie complete sequence, 52920 bp including 10-base 3' overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 2/27



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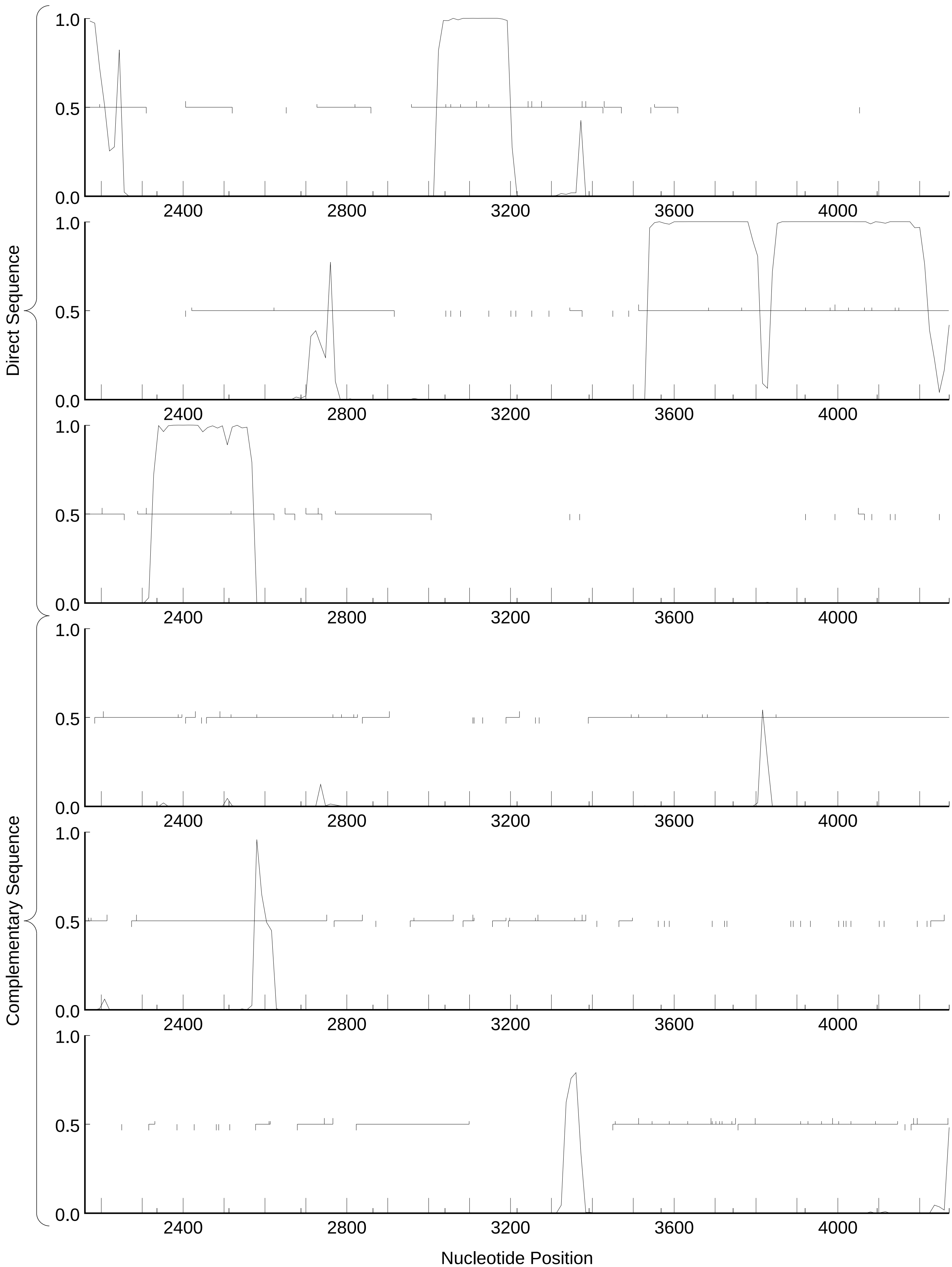

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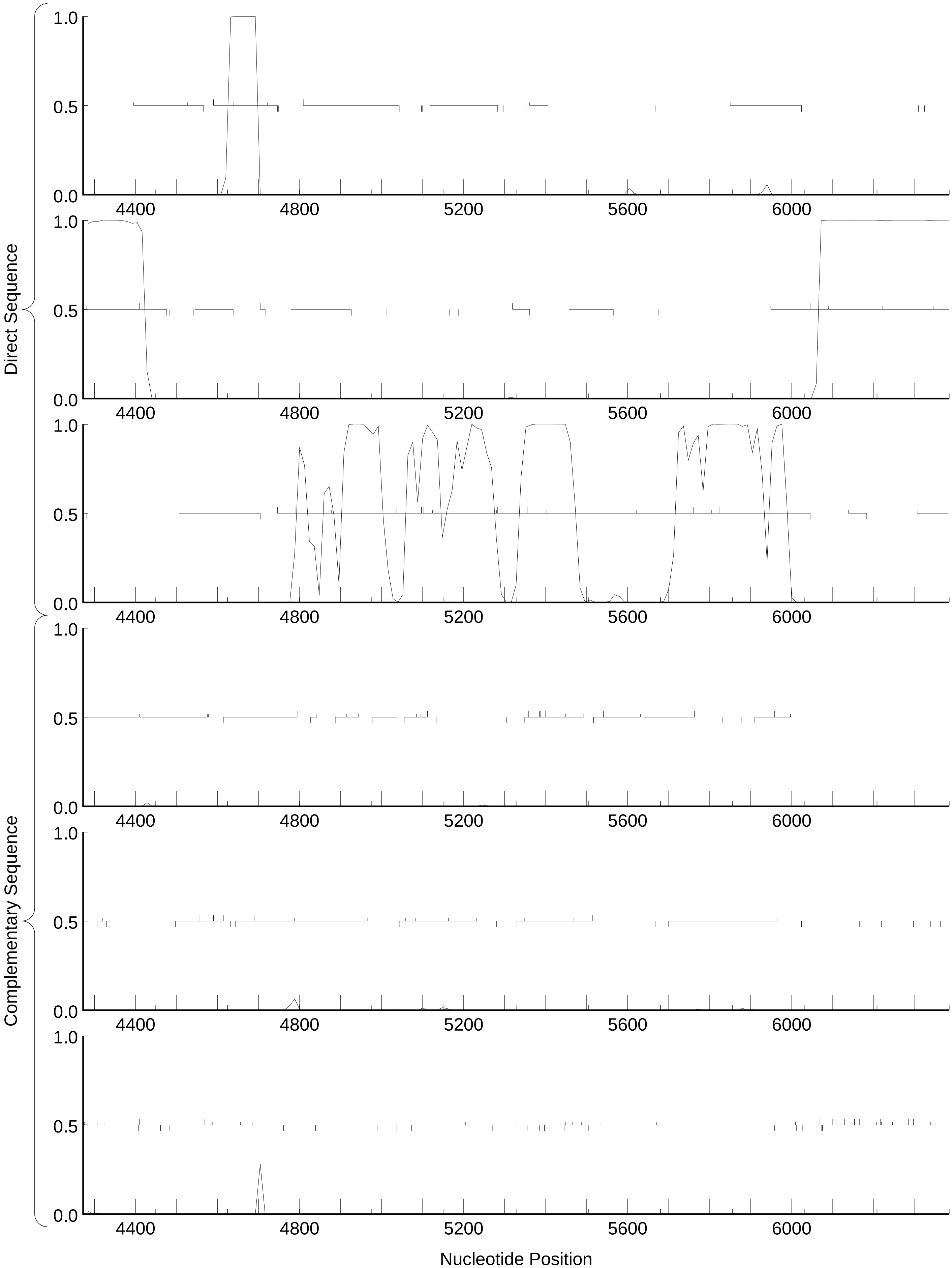

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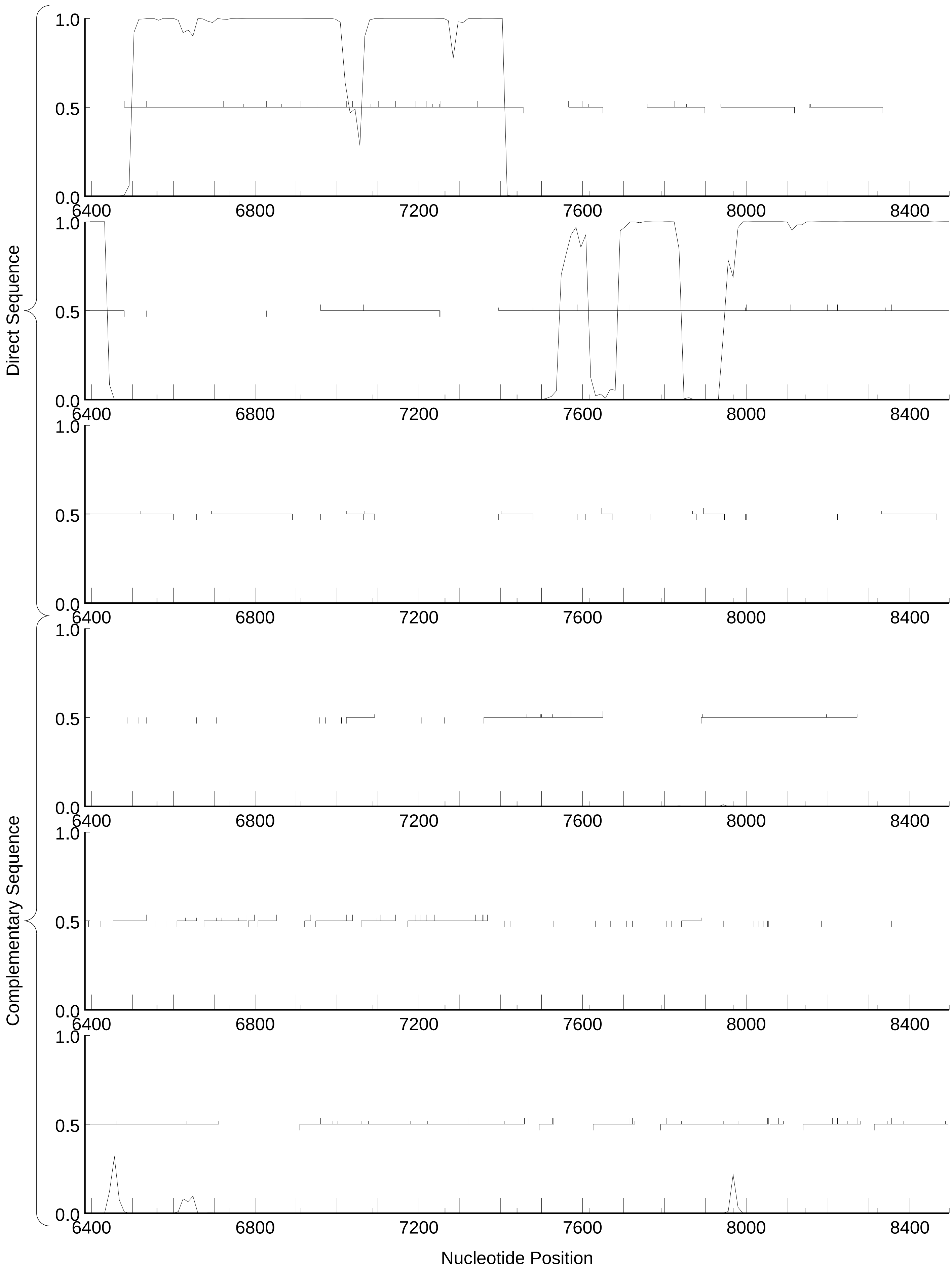

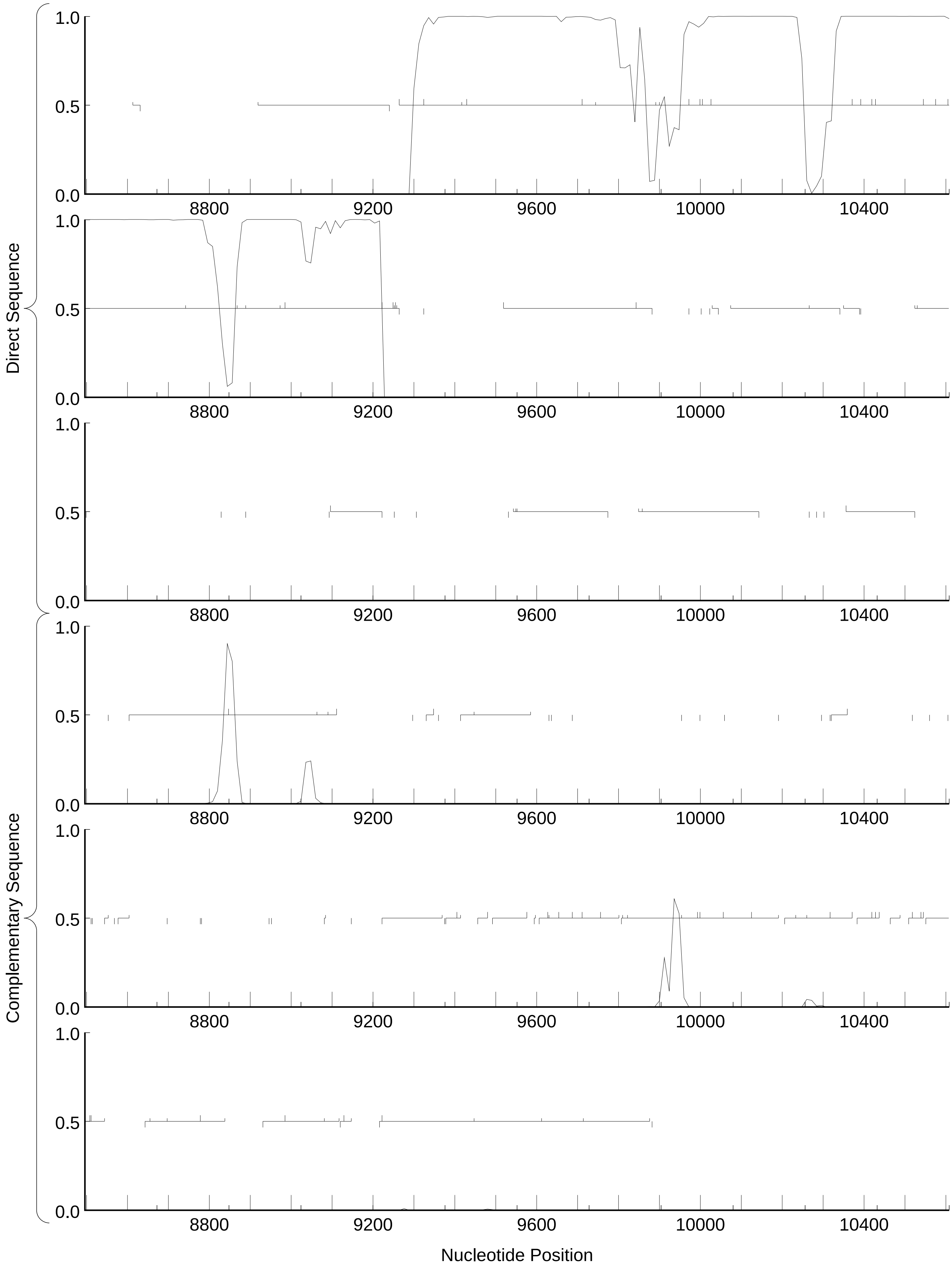

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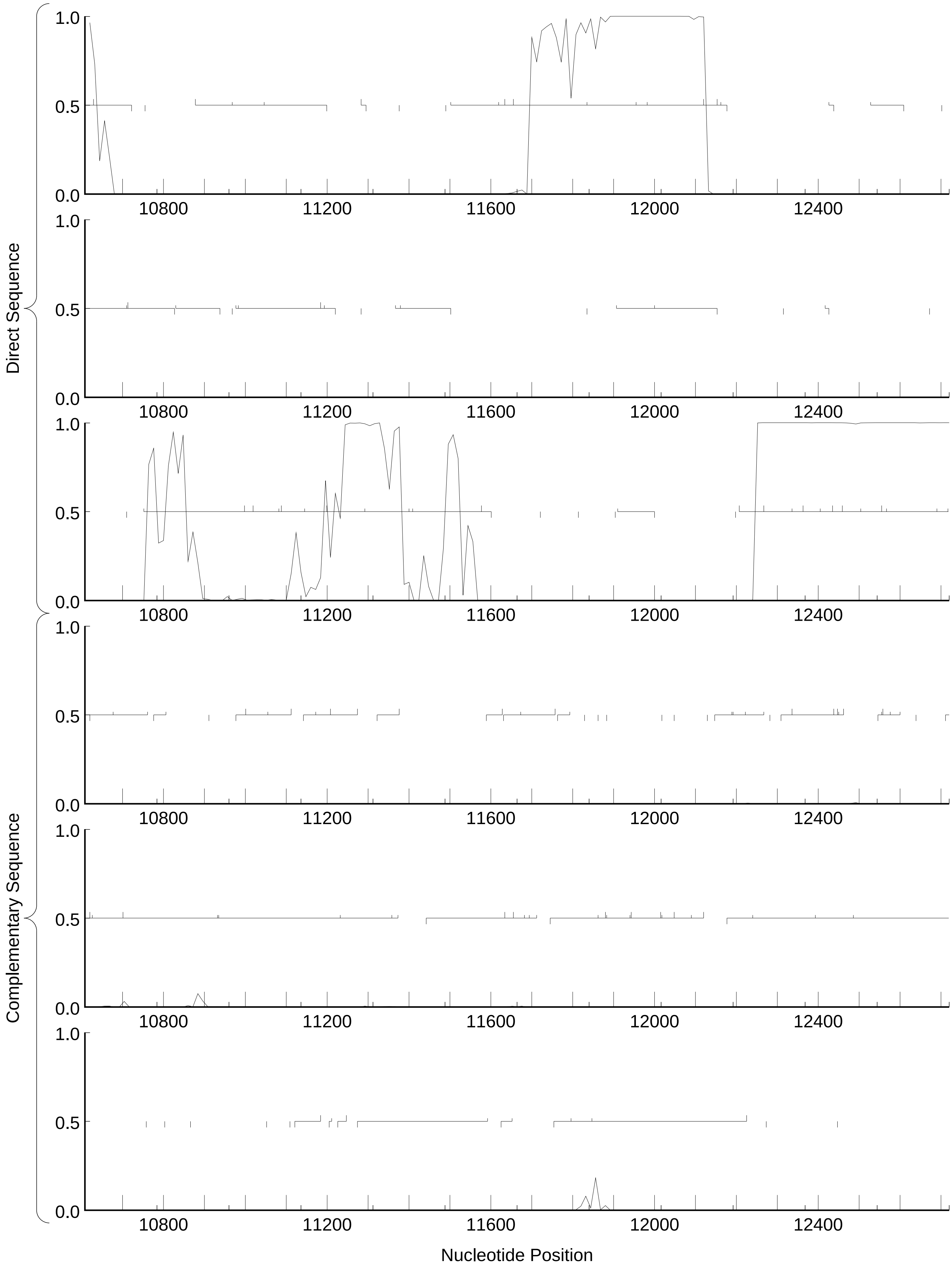
Nucleotide Position



Direct Sequence
Complementary Sequence
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10800
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11600
12000
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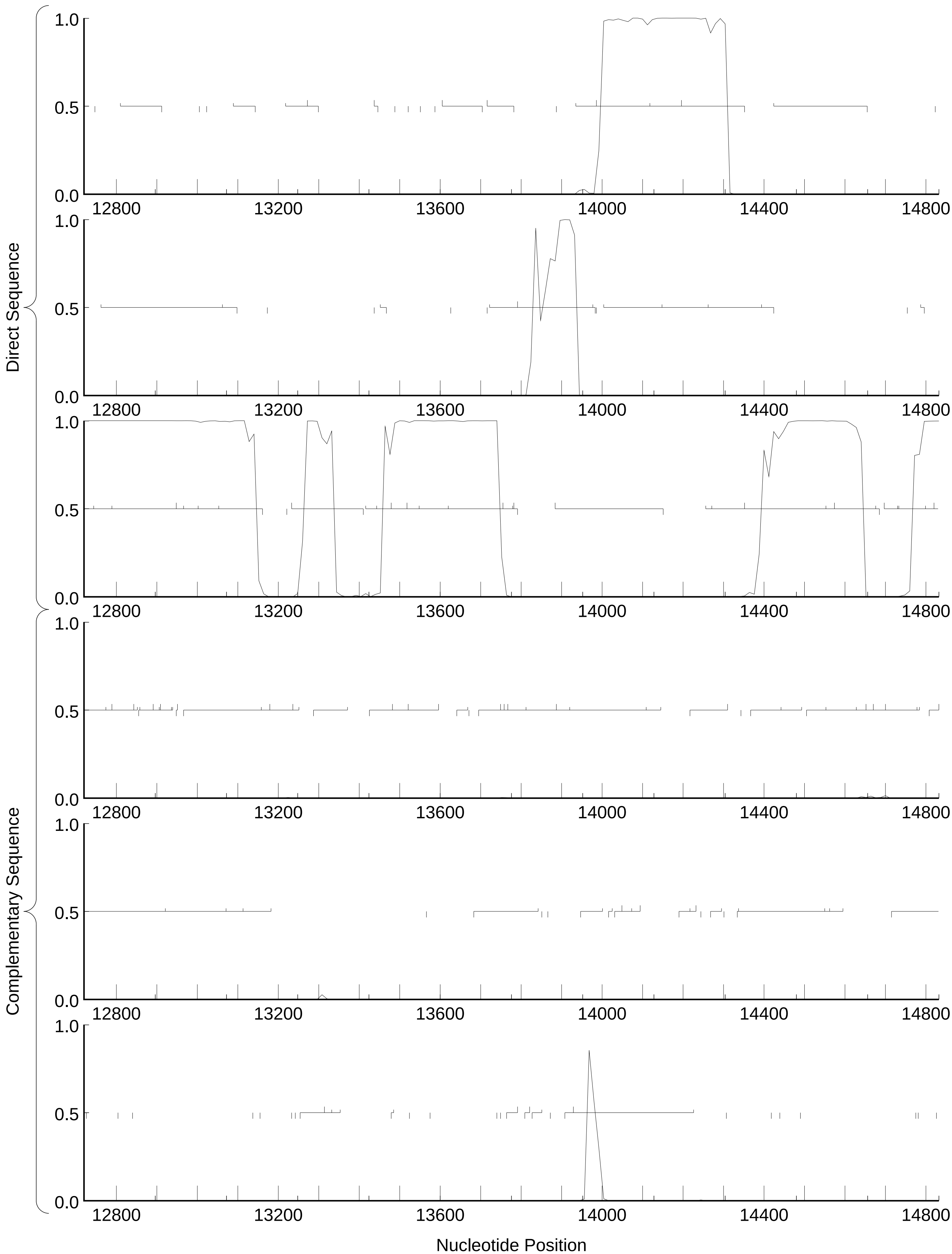
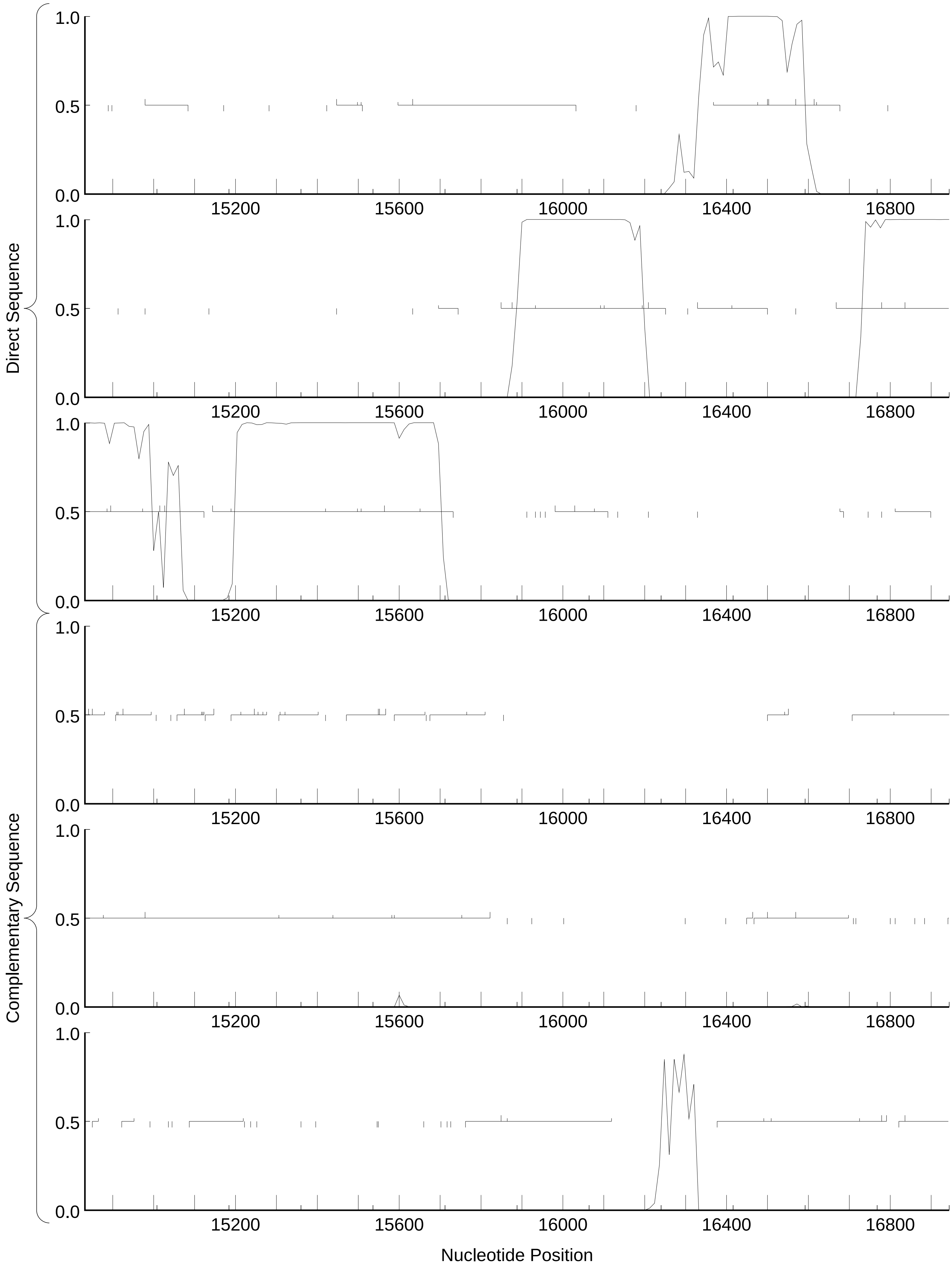
Nucleotide Position

ie complete sequence, 52920 bp including 10-base 3' overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 8/27

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Nucleotide Position

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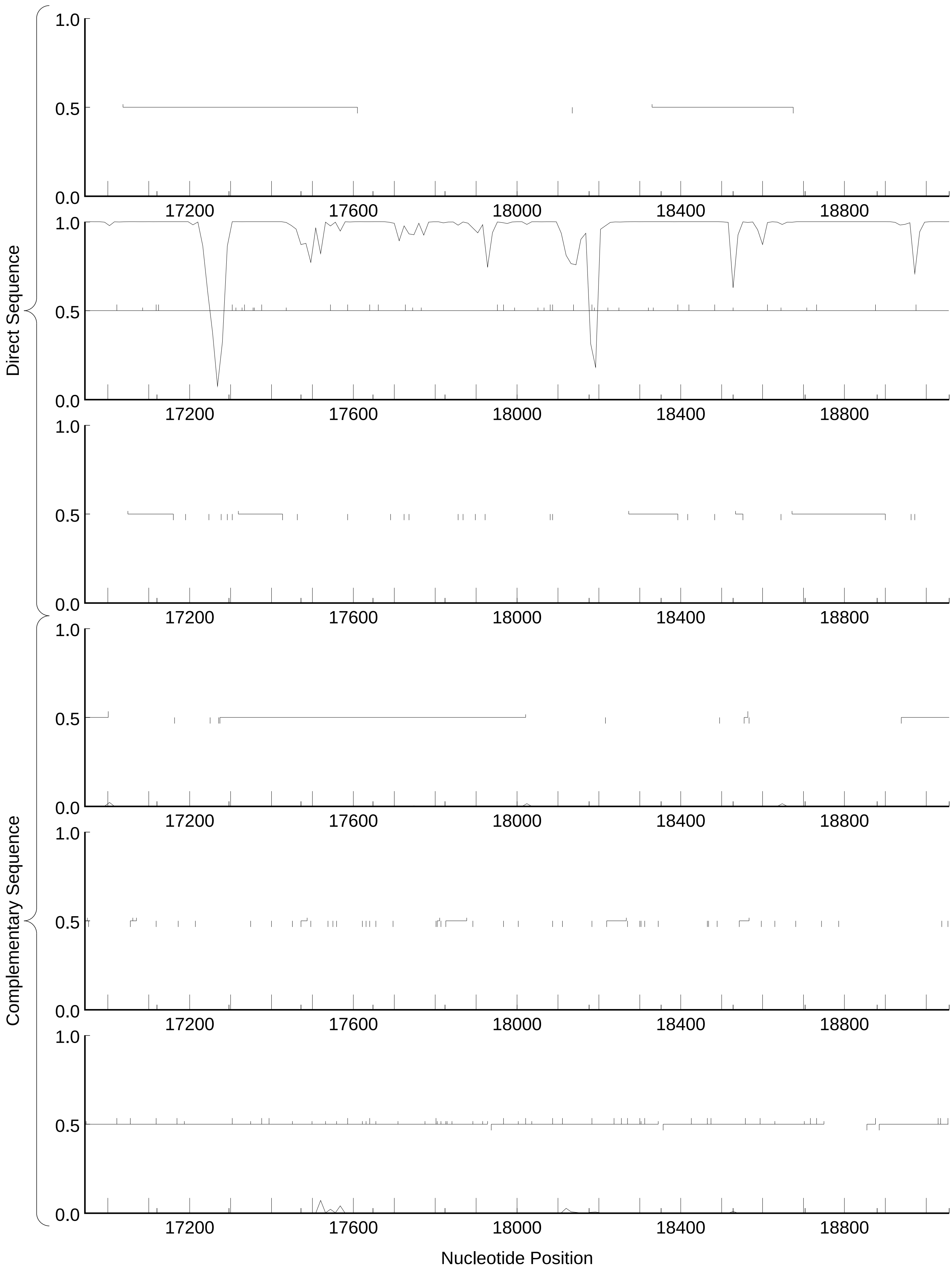

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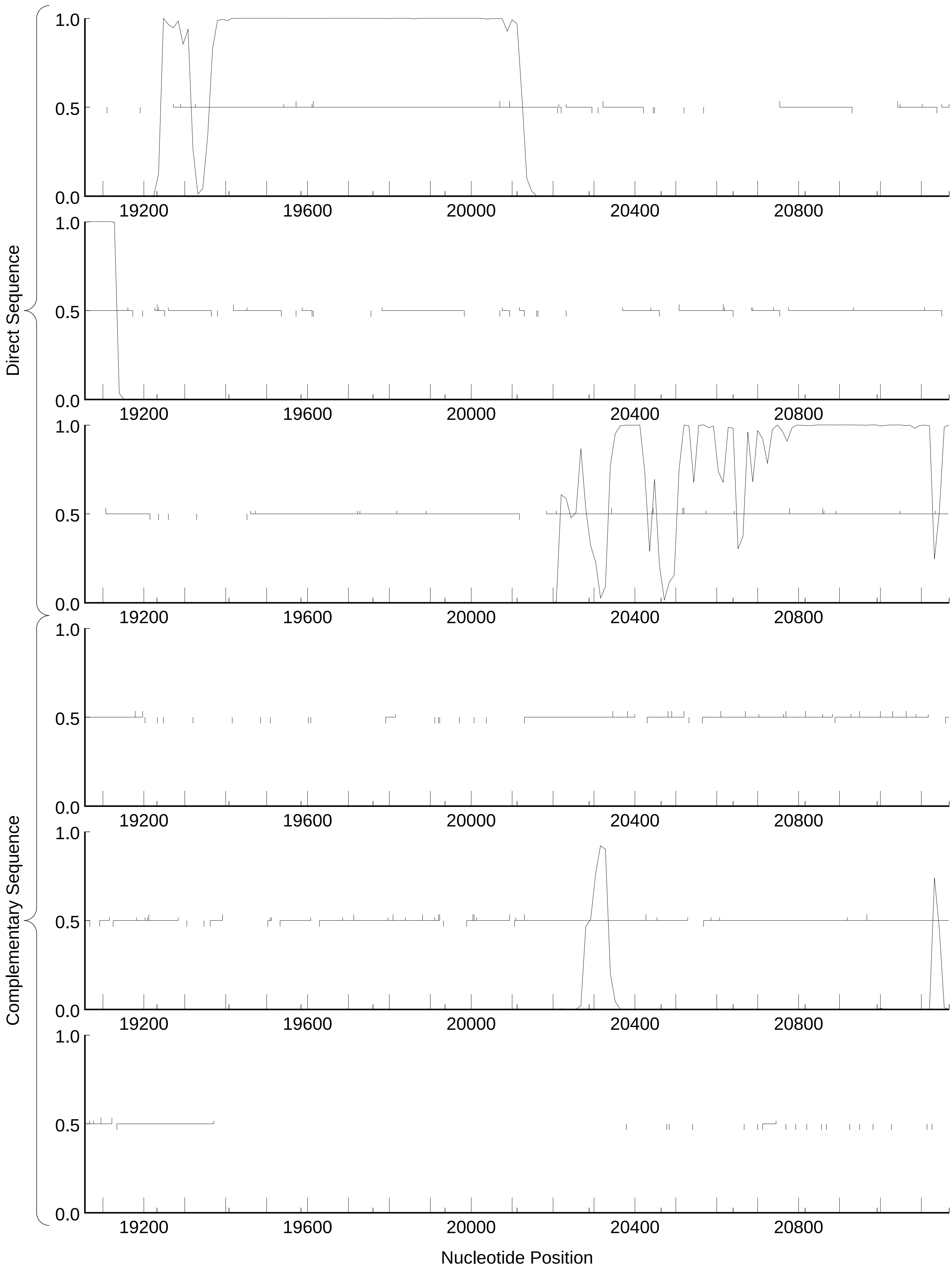

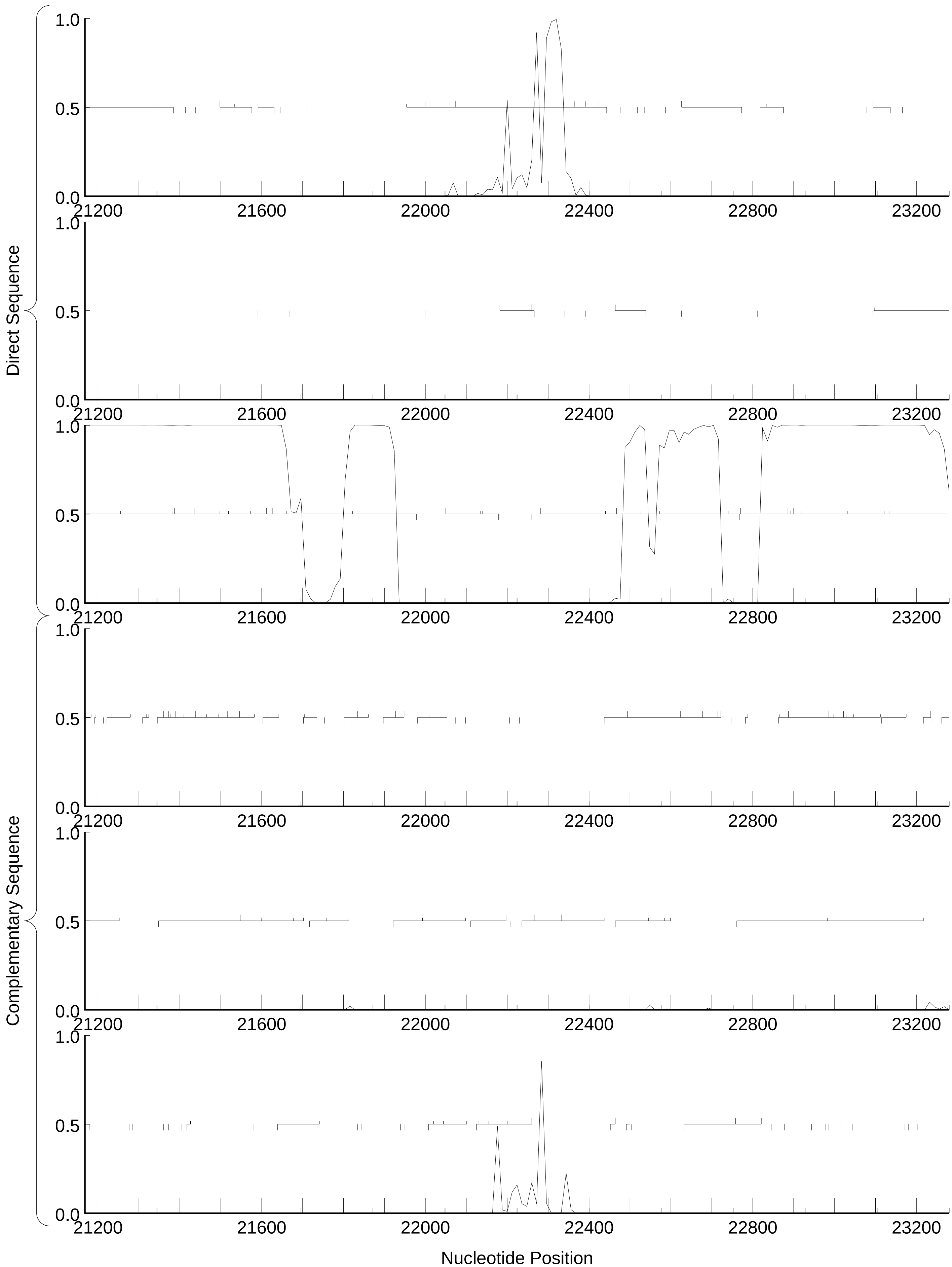
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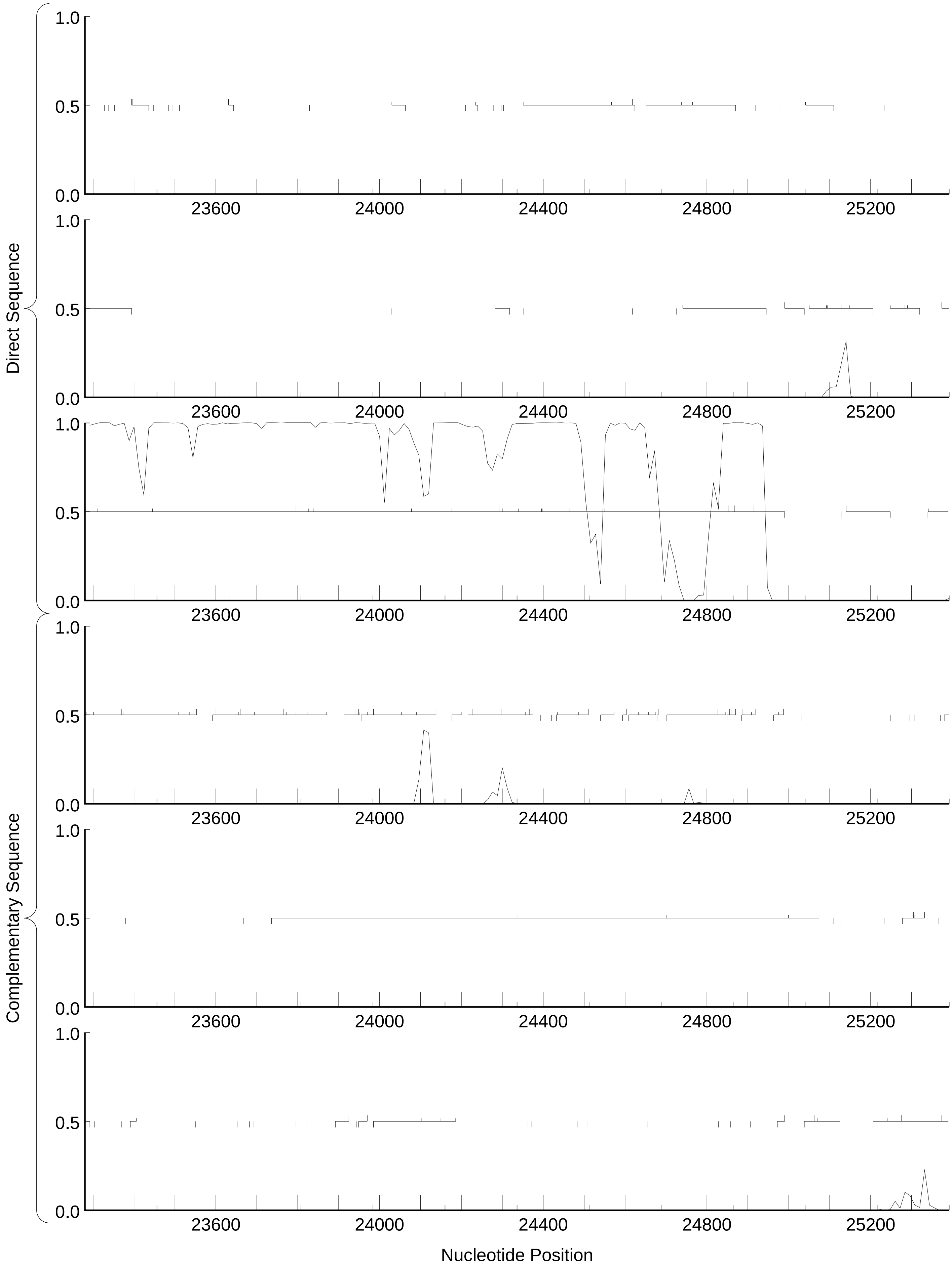
Nucleotide Position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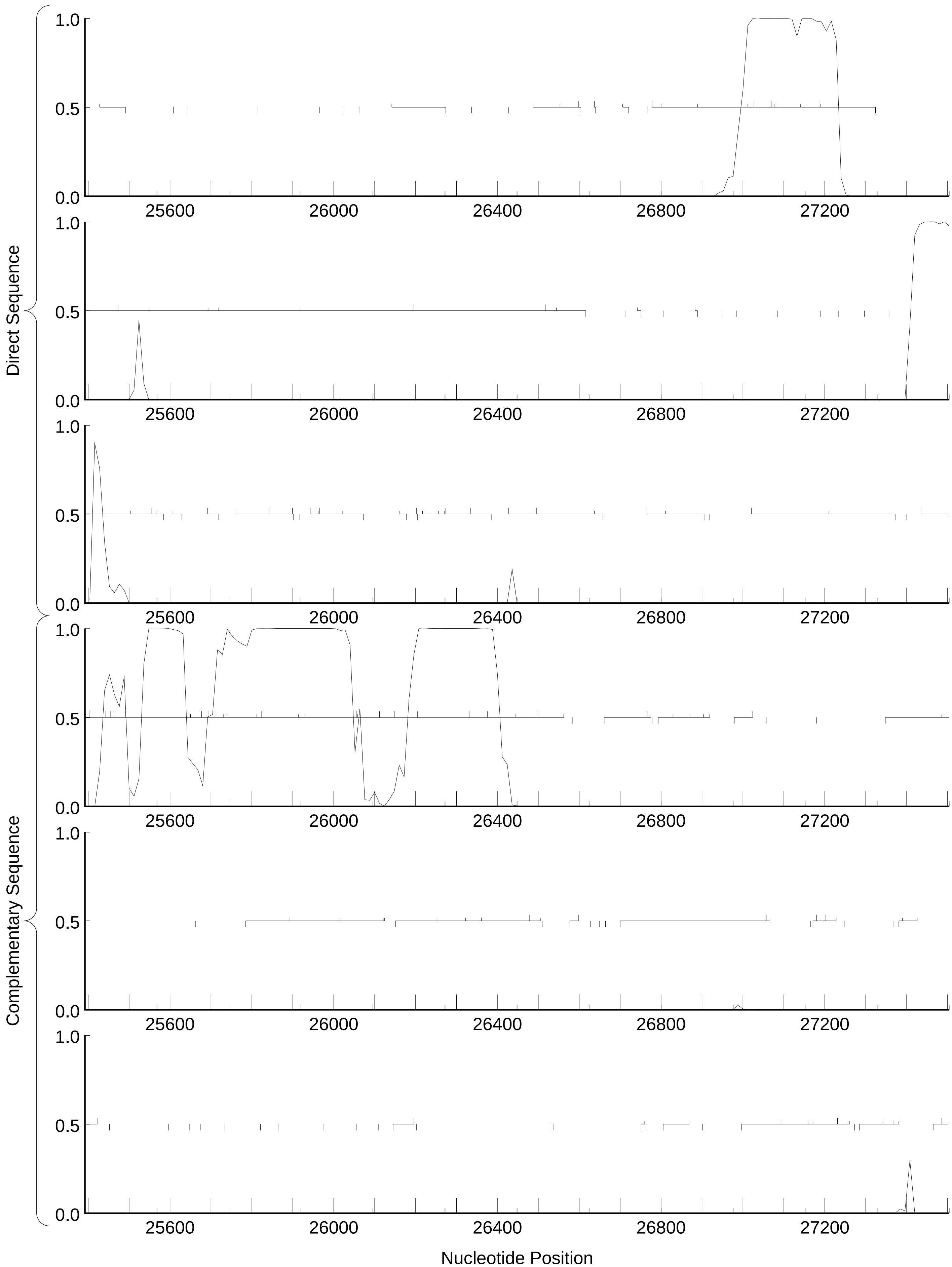
Nucleotide Position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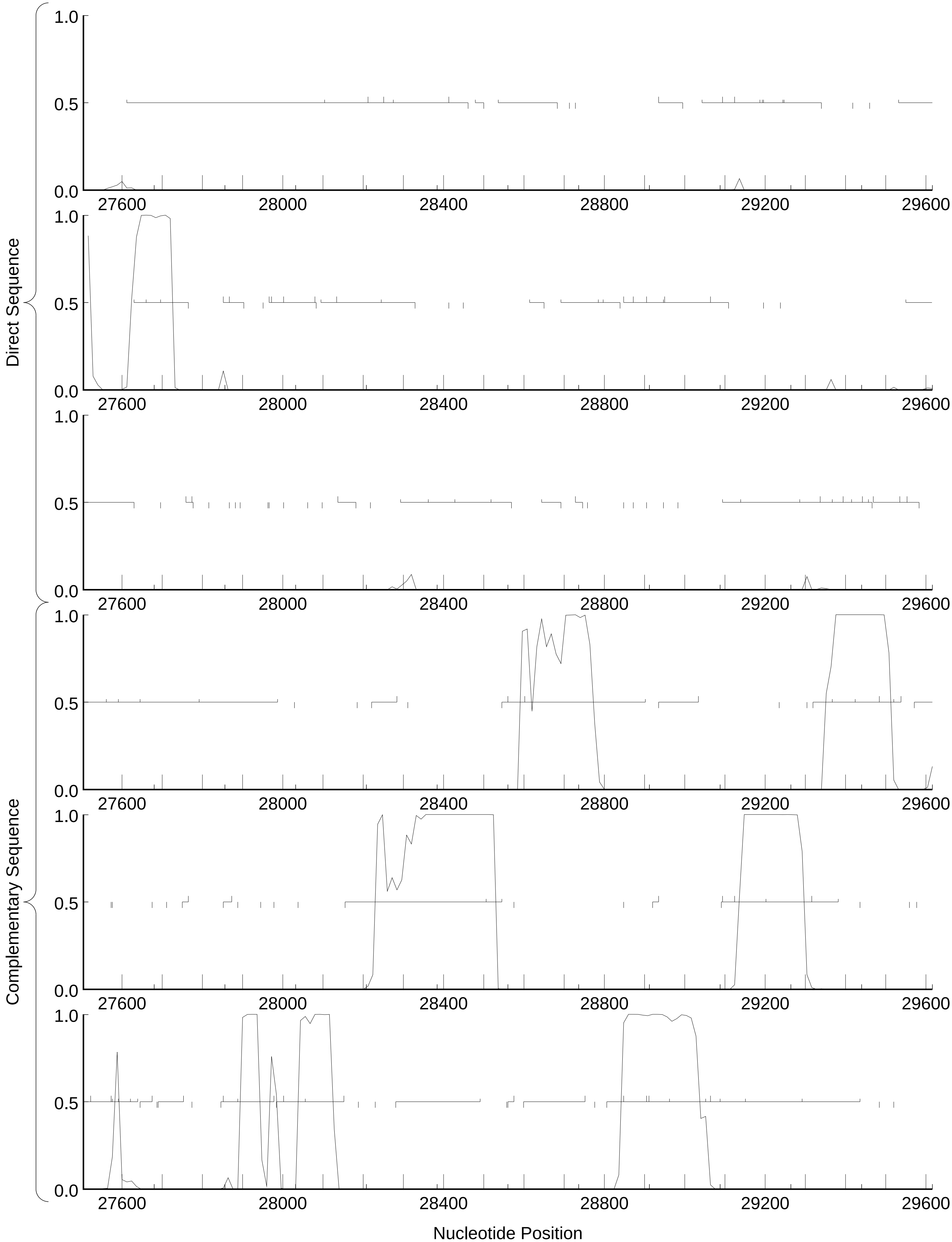
Nucleotide Position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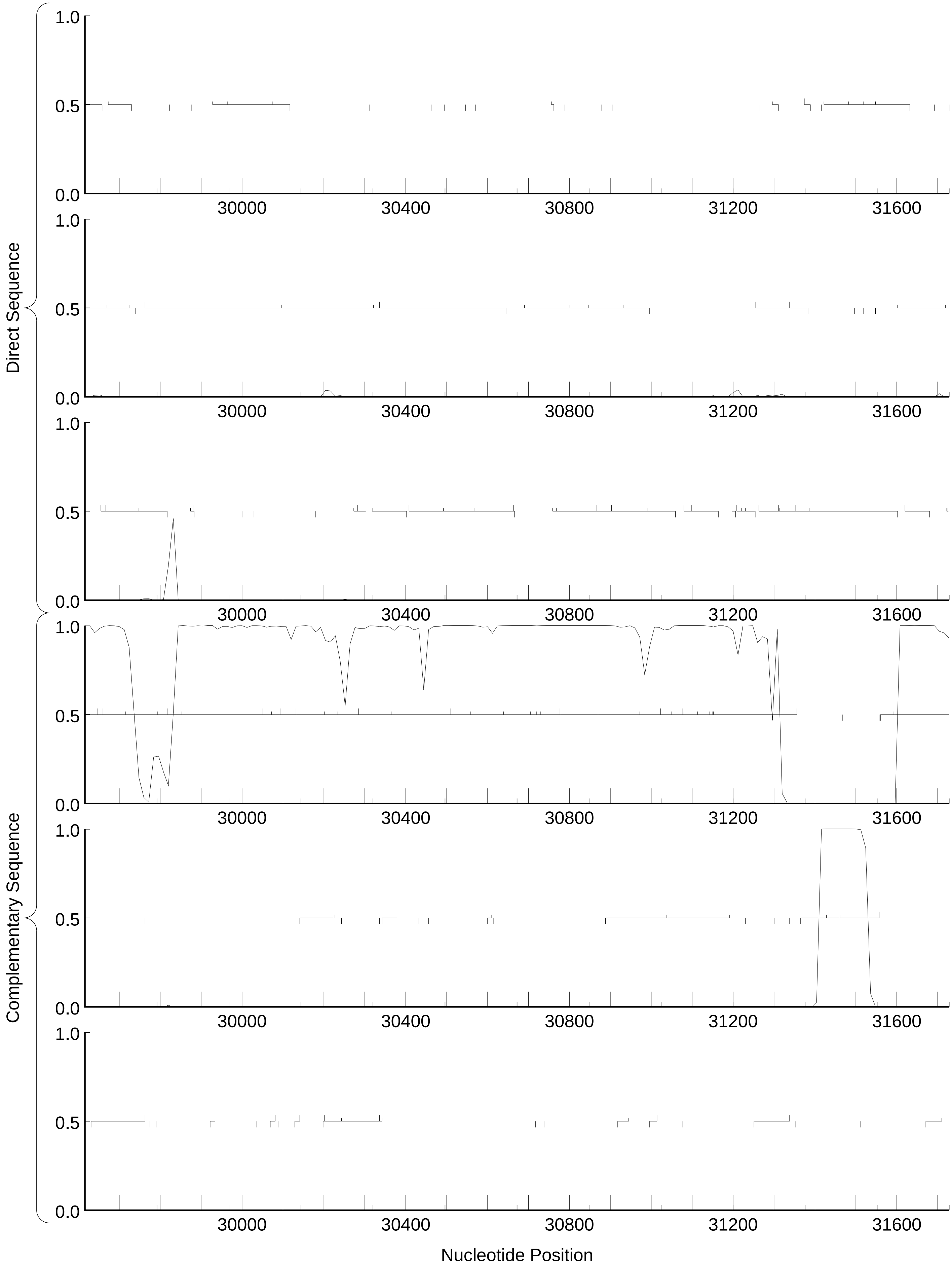
Nucleotide Position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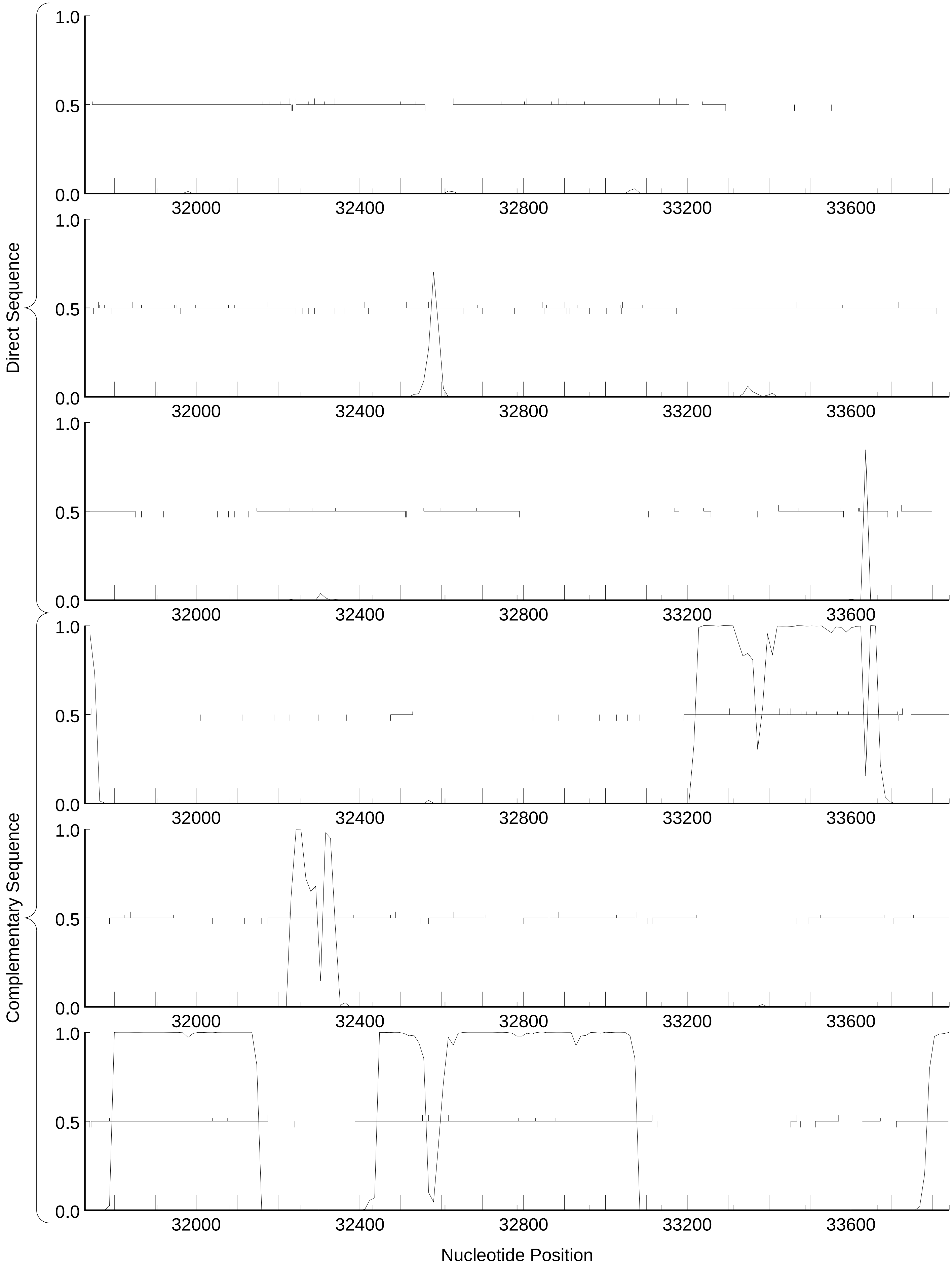
Nucleotide Position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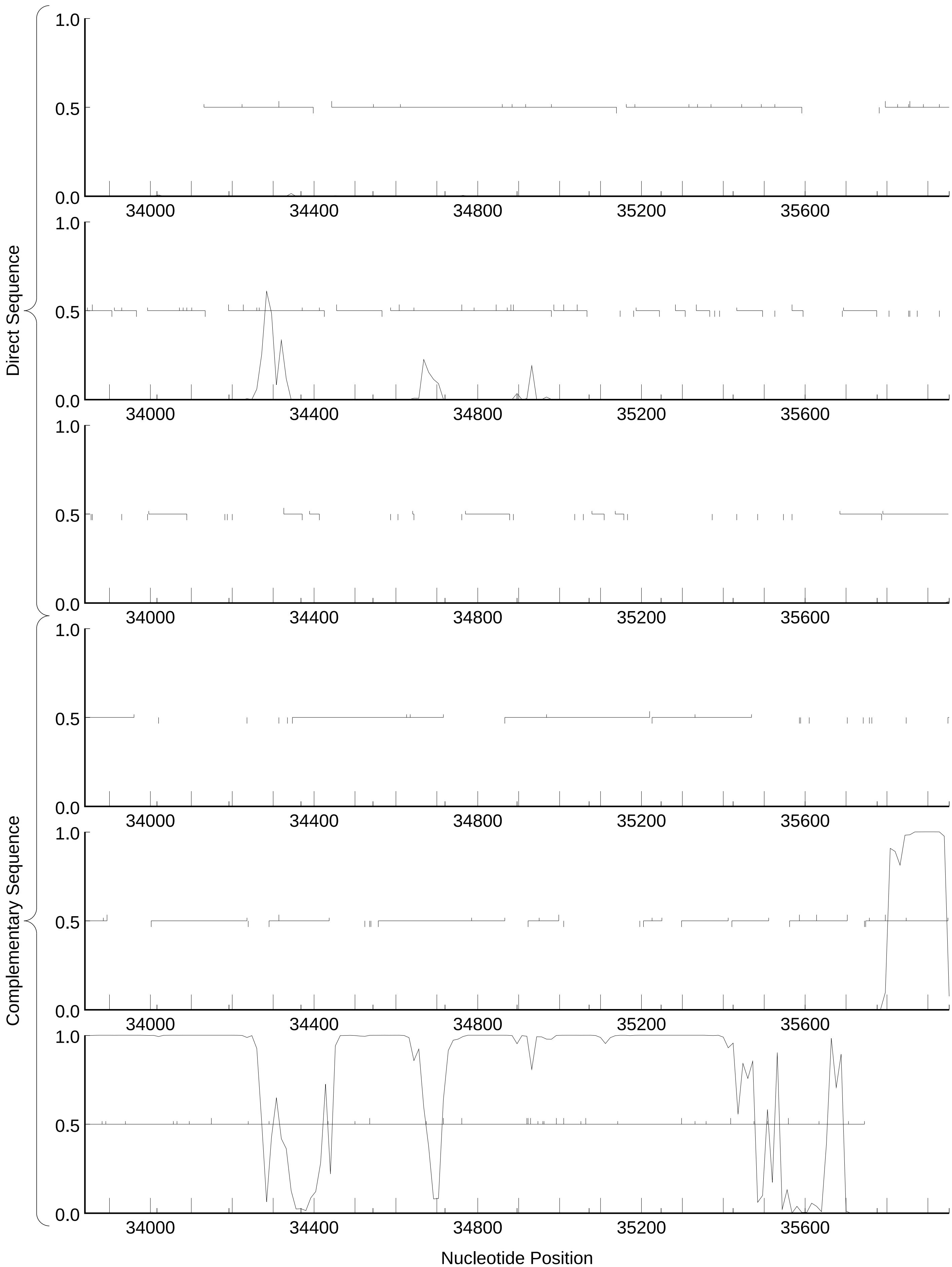
Nucleotide Position



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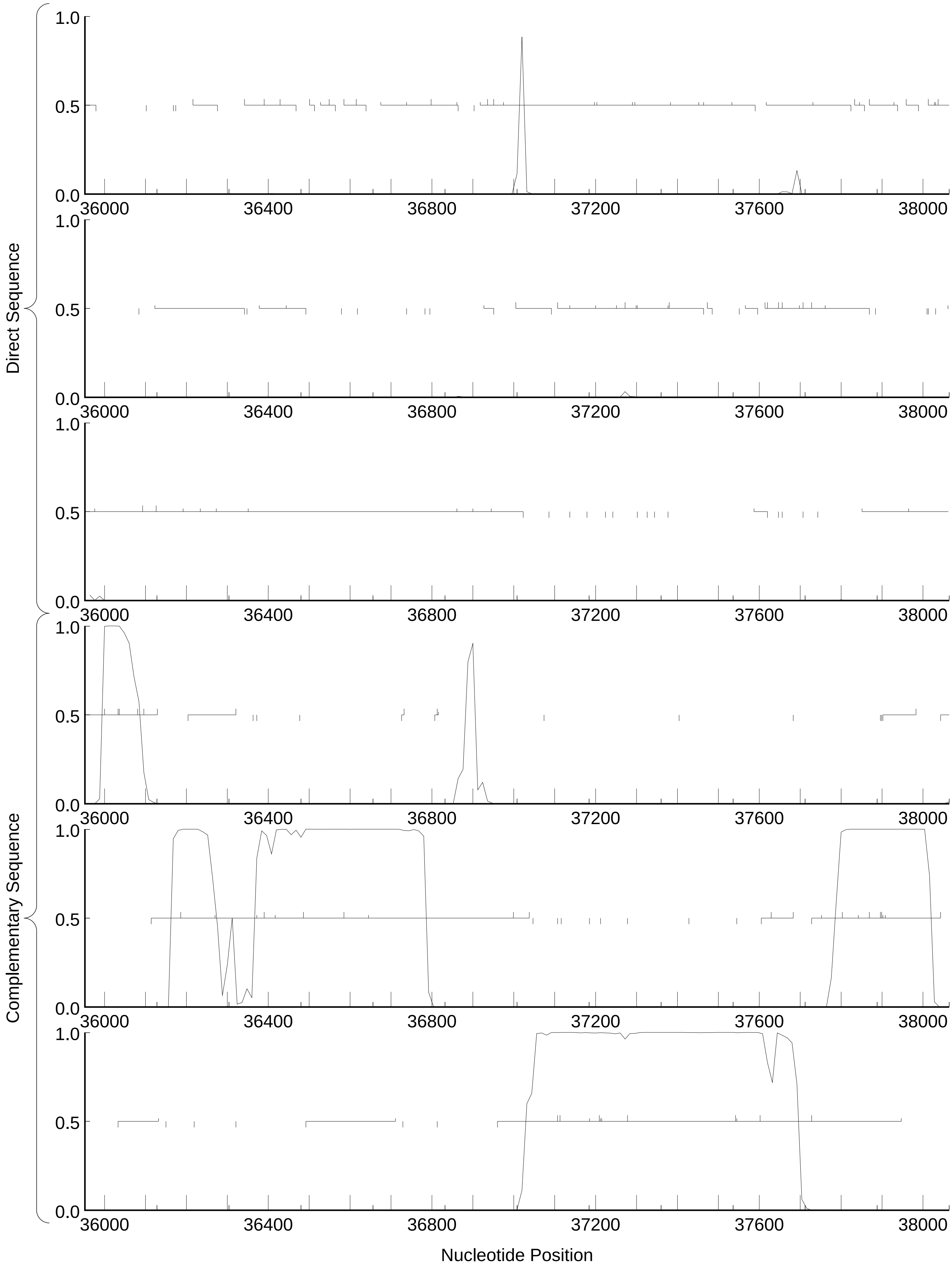
Nucleotide Position



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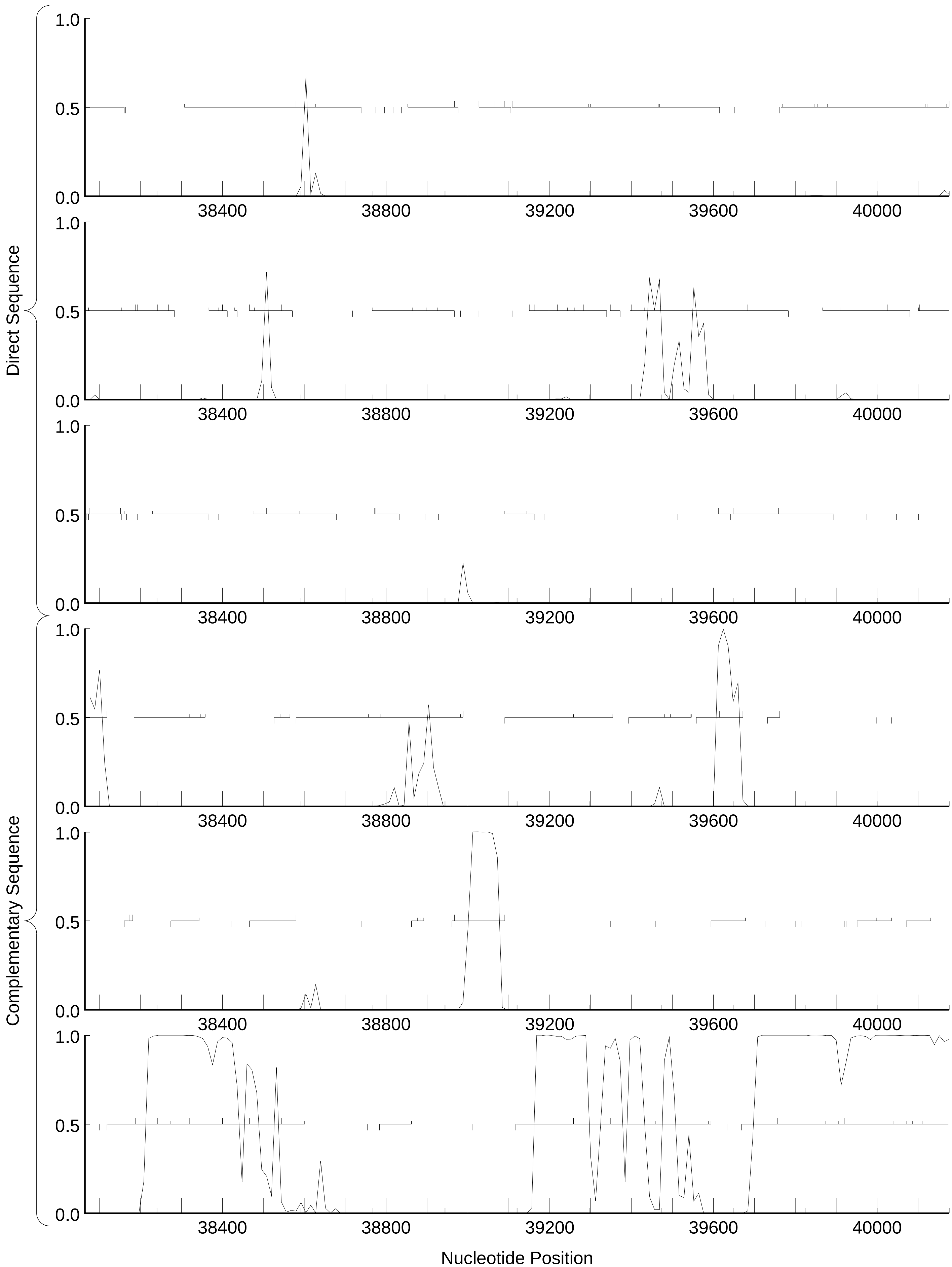
Nucleotide Position



Direct Sequence
Complementary Sequence
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0.0
38400
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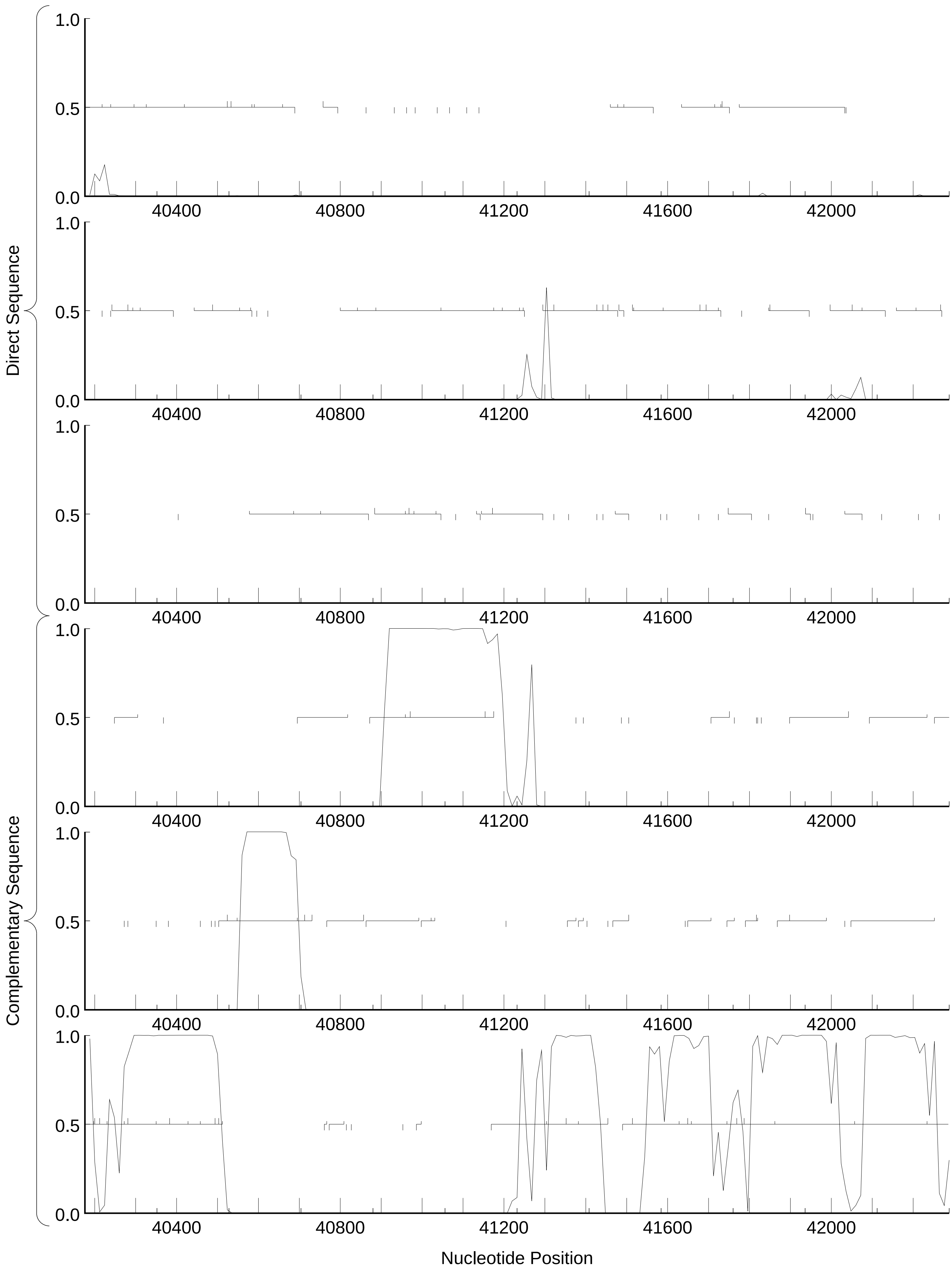
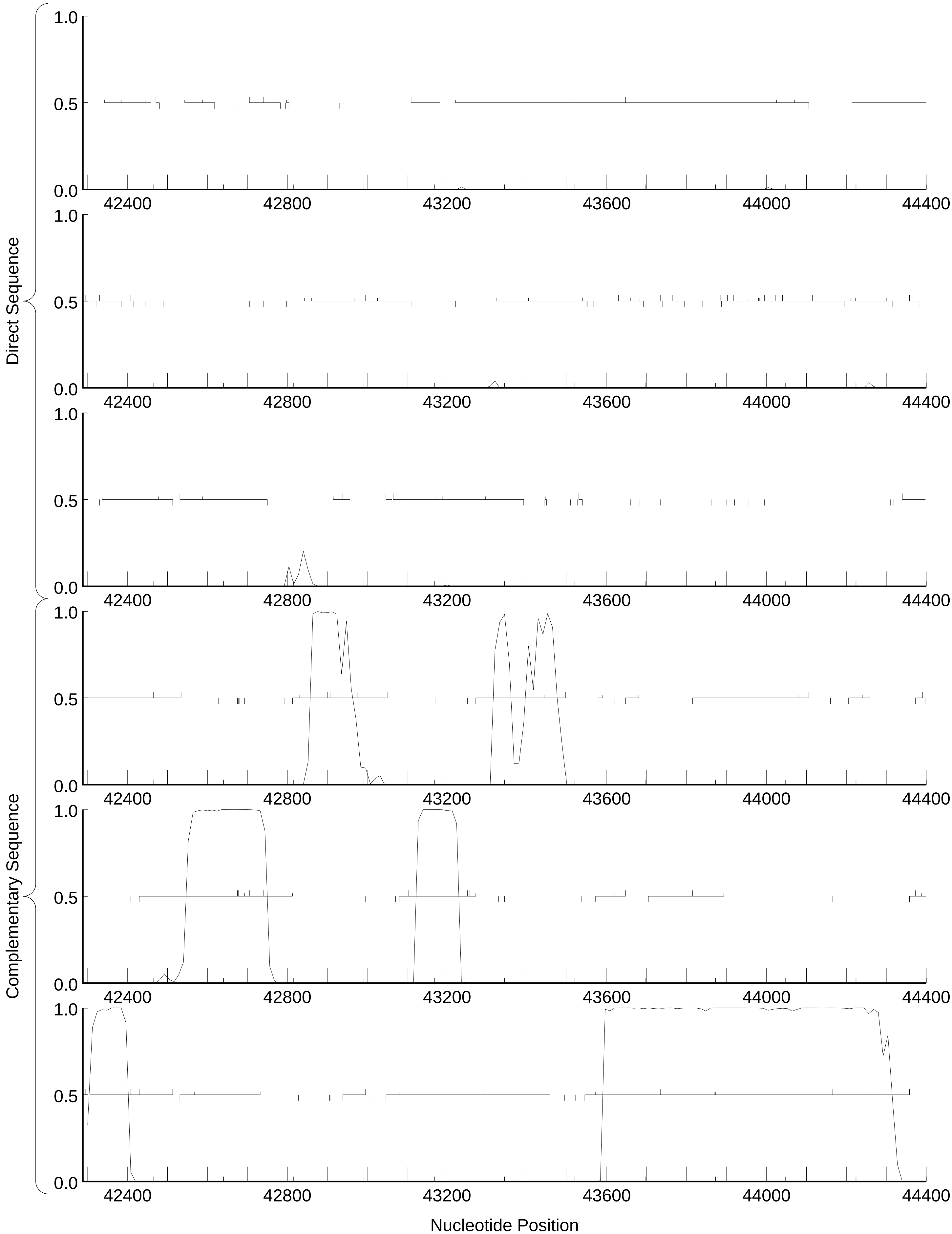
Nucleotide Position

e complete sequence, 52920 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 21/27

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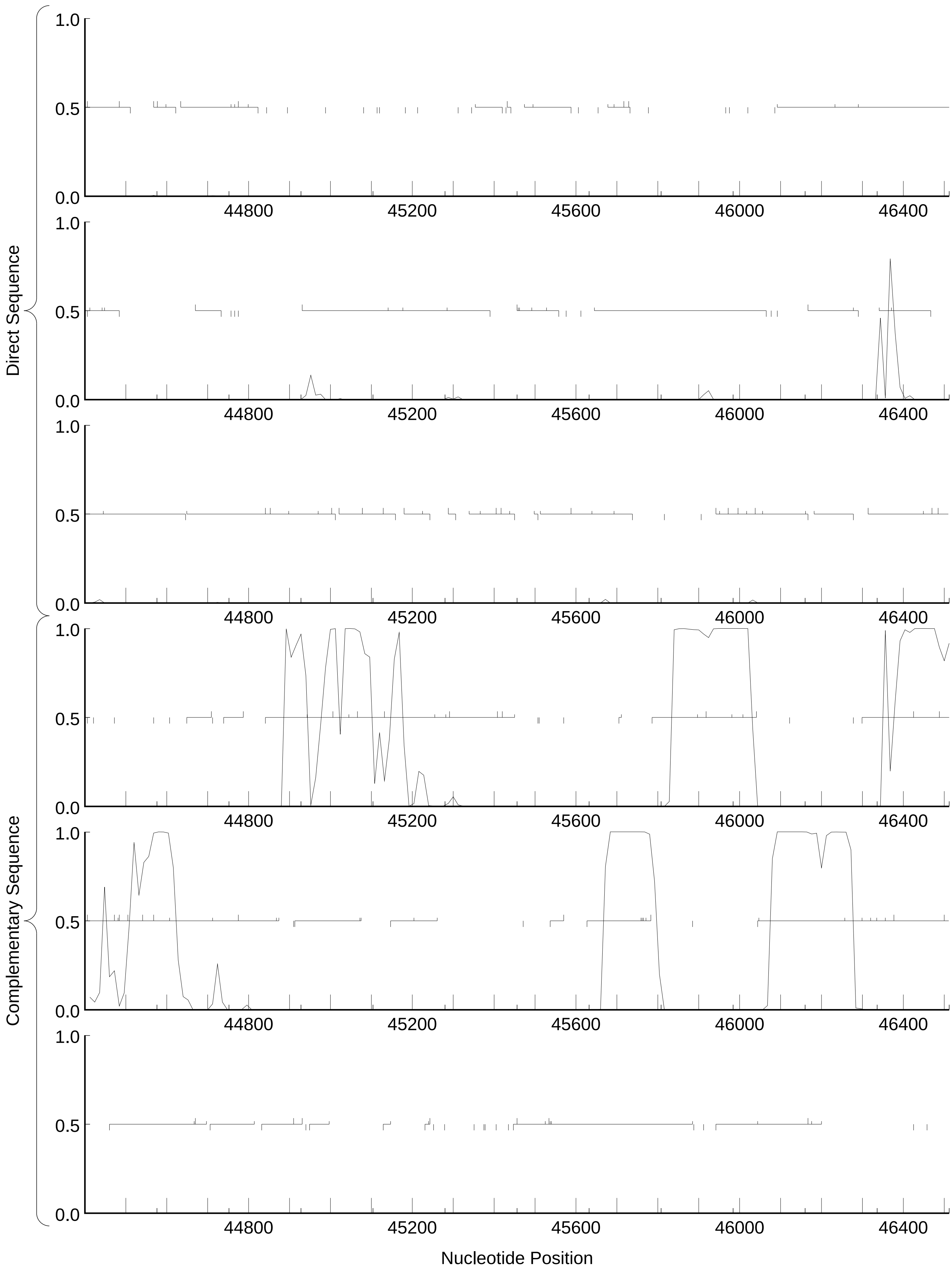
Nucleotide Position



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Complementary Sequence
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Nucleotide Position

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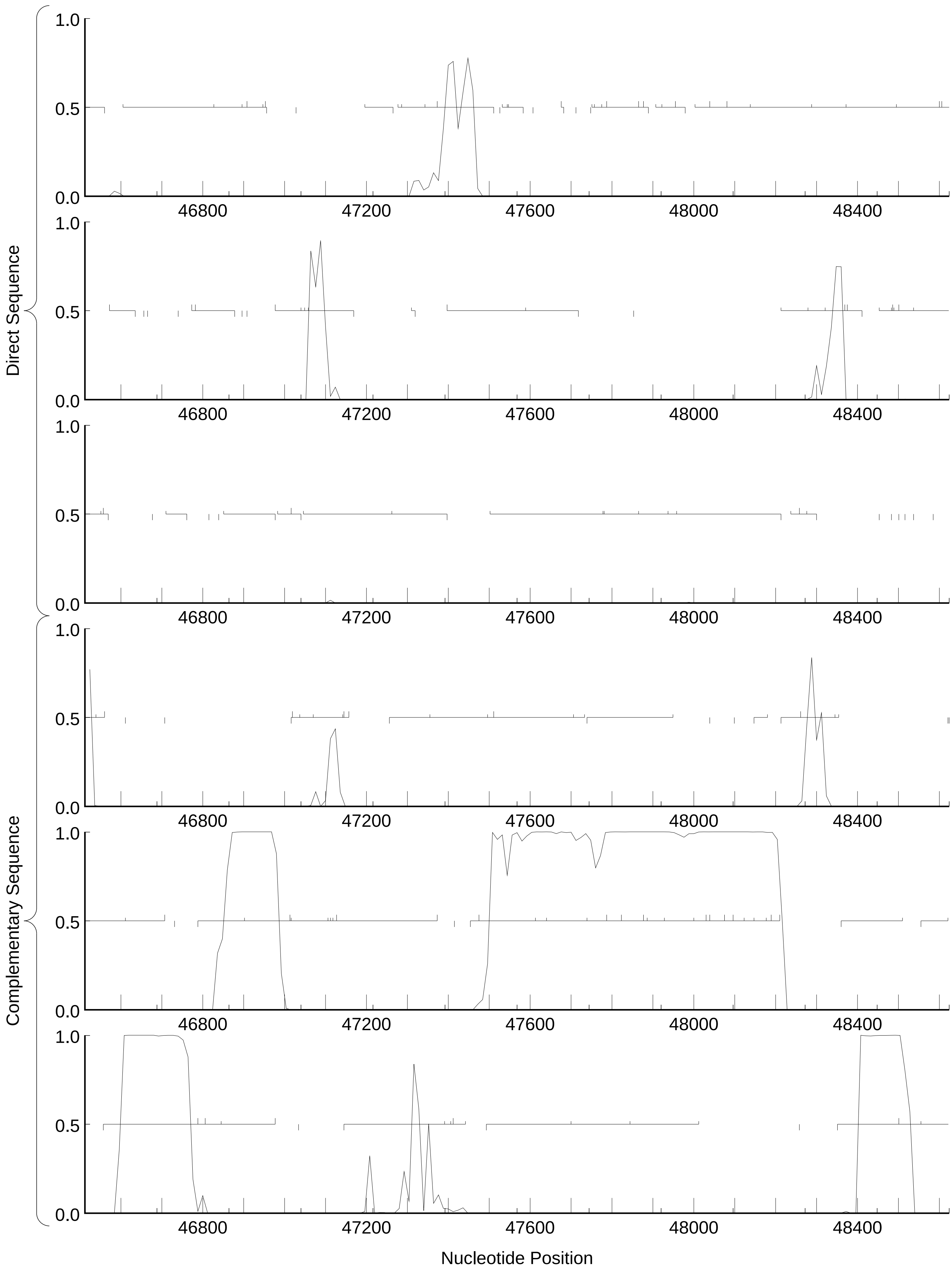

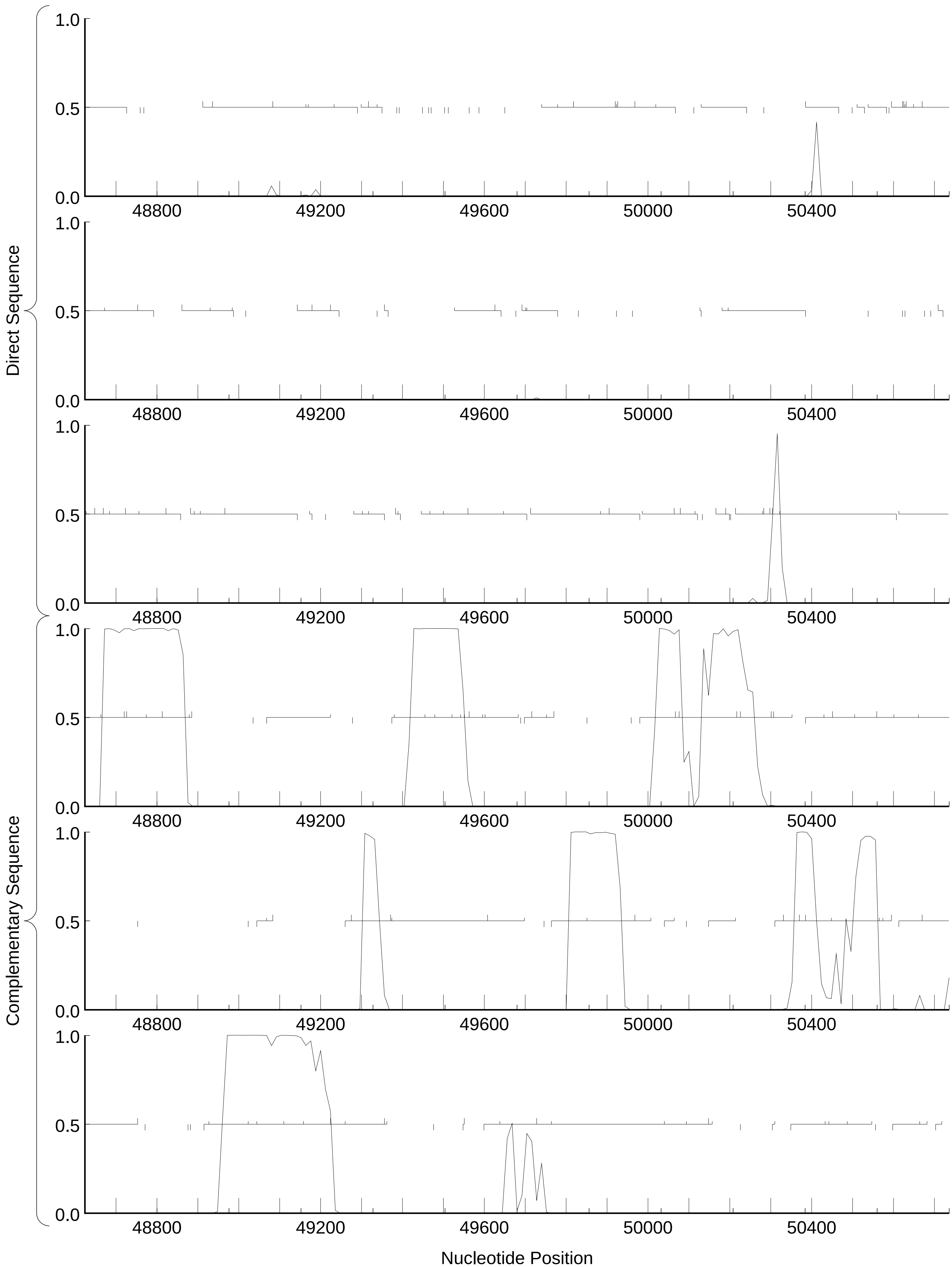
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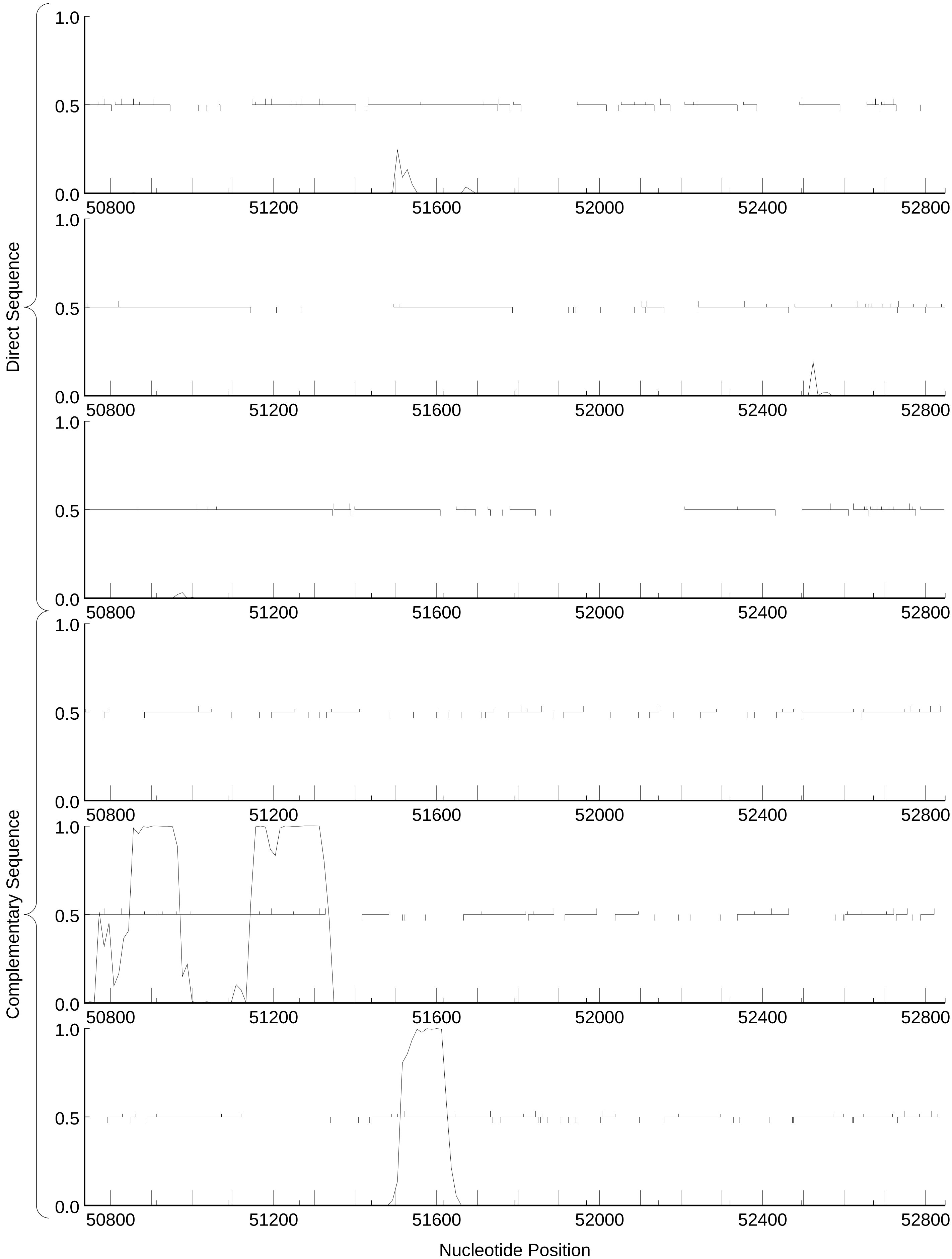
Complementary Sequence
Nucleotide Position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Complementary Sequence
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52000
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Nucleotide Position

e complete sequence, 52920 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A2, Order 4, Window 96, Step 12, 27/27