GeneMark

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

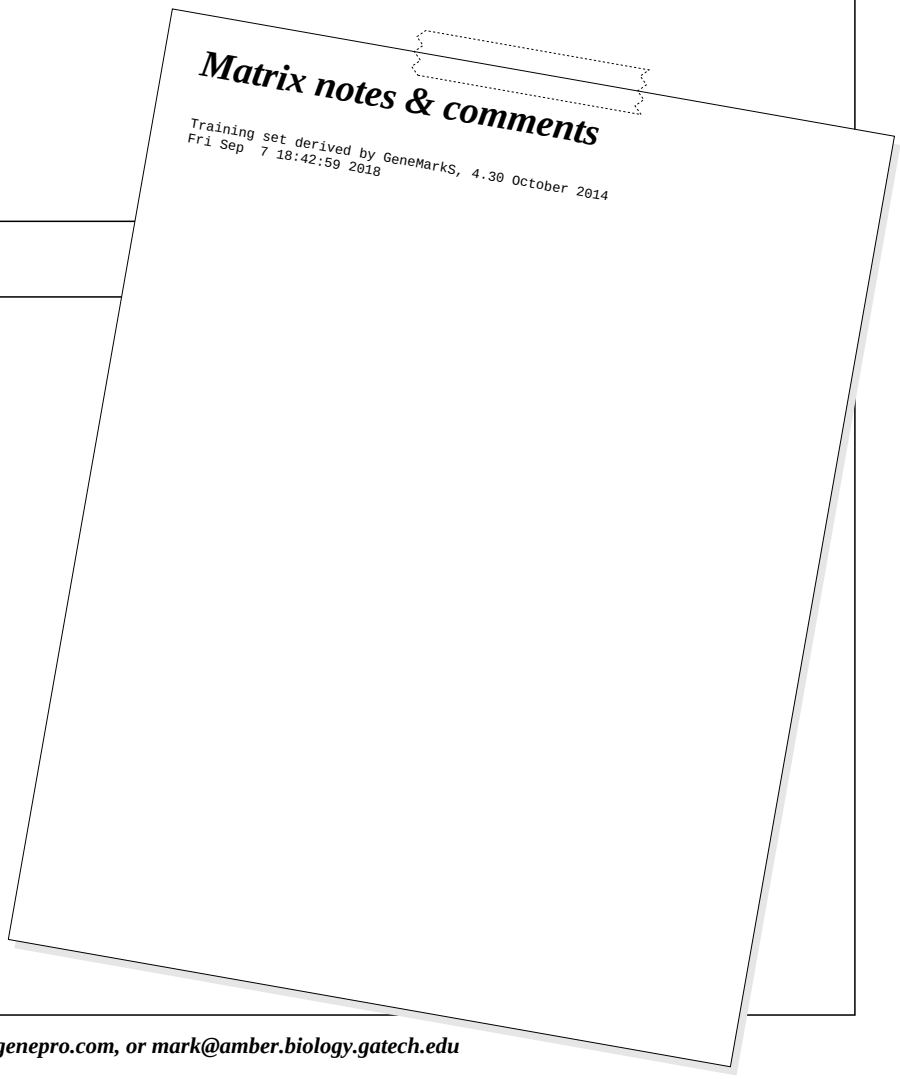
PROGRAM INFORMATION

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Sequence         :  Microbacterium phage BeeBee8 complete sequence, 42027 bp, circularly permuted, Cluster EA
Analysis Date    :  3/28/24 at 10:20:42
Pages            :  21
Sequence Length  :  42027 bp
GC Content       :   63.35%

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,
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MATRIX INFORMATION

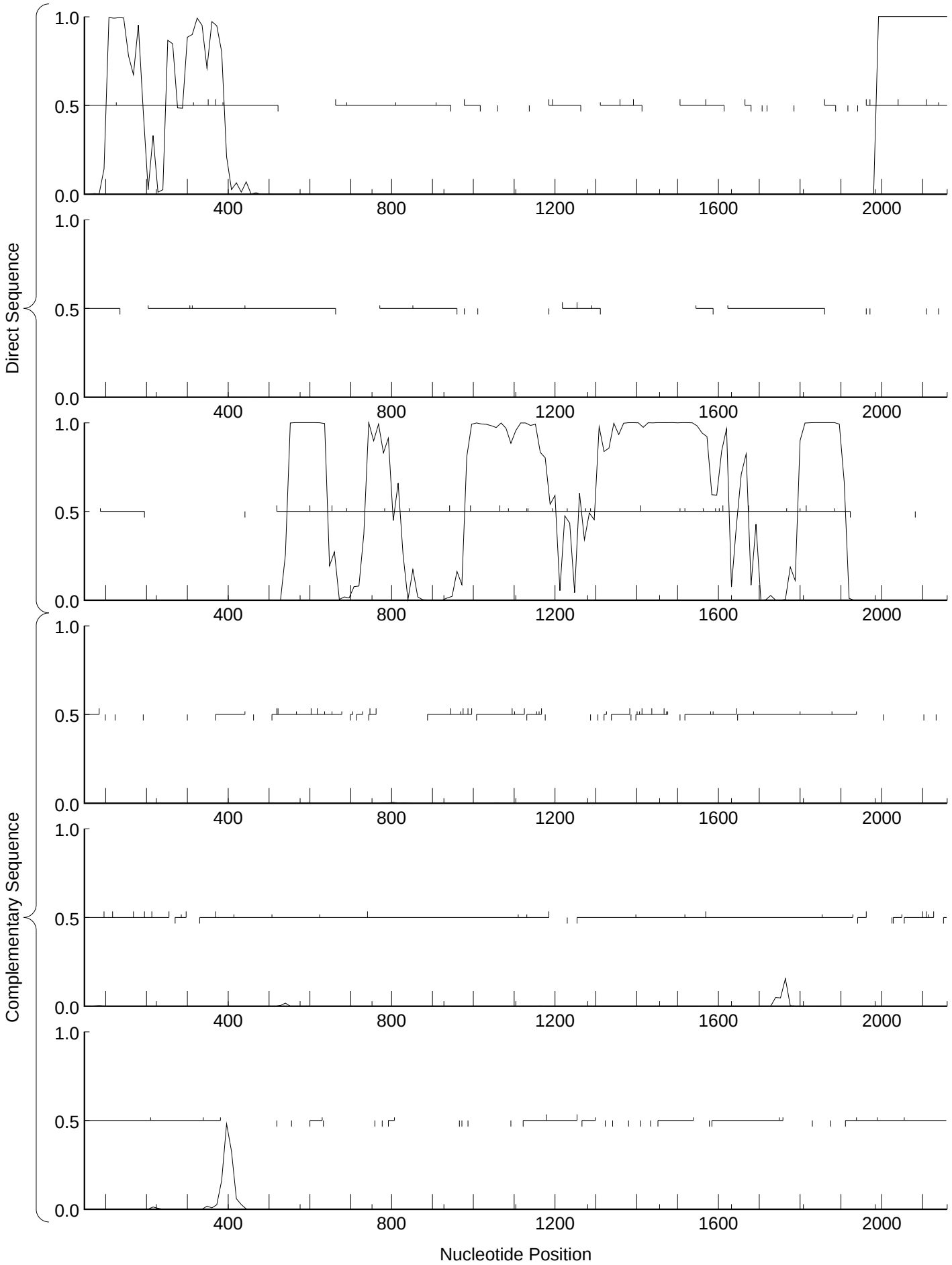
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Matrix     :  -
Author     :  -

Order      :  4
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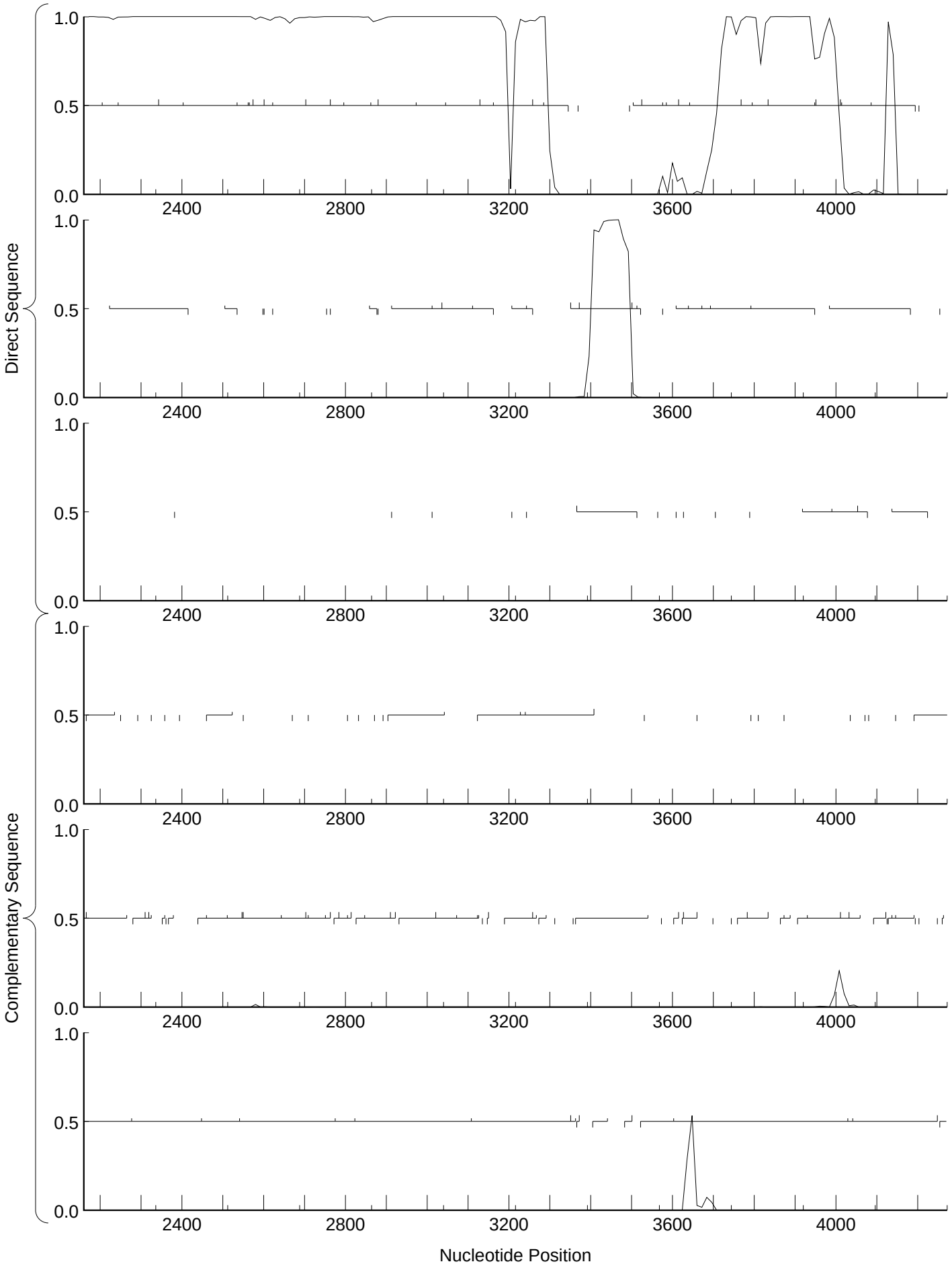

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

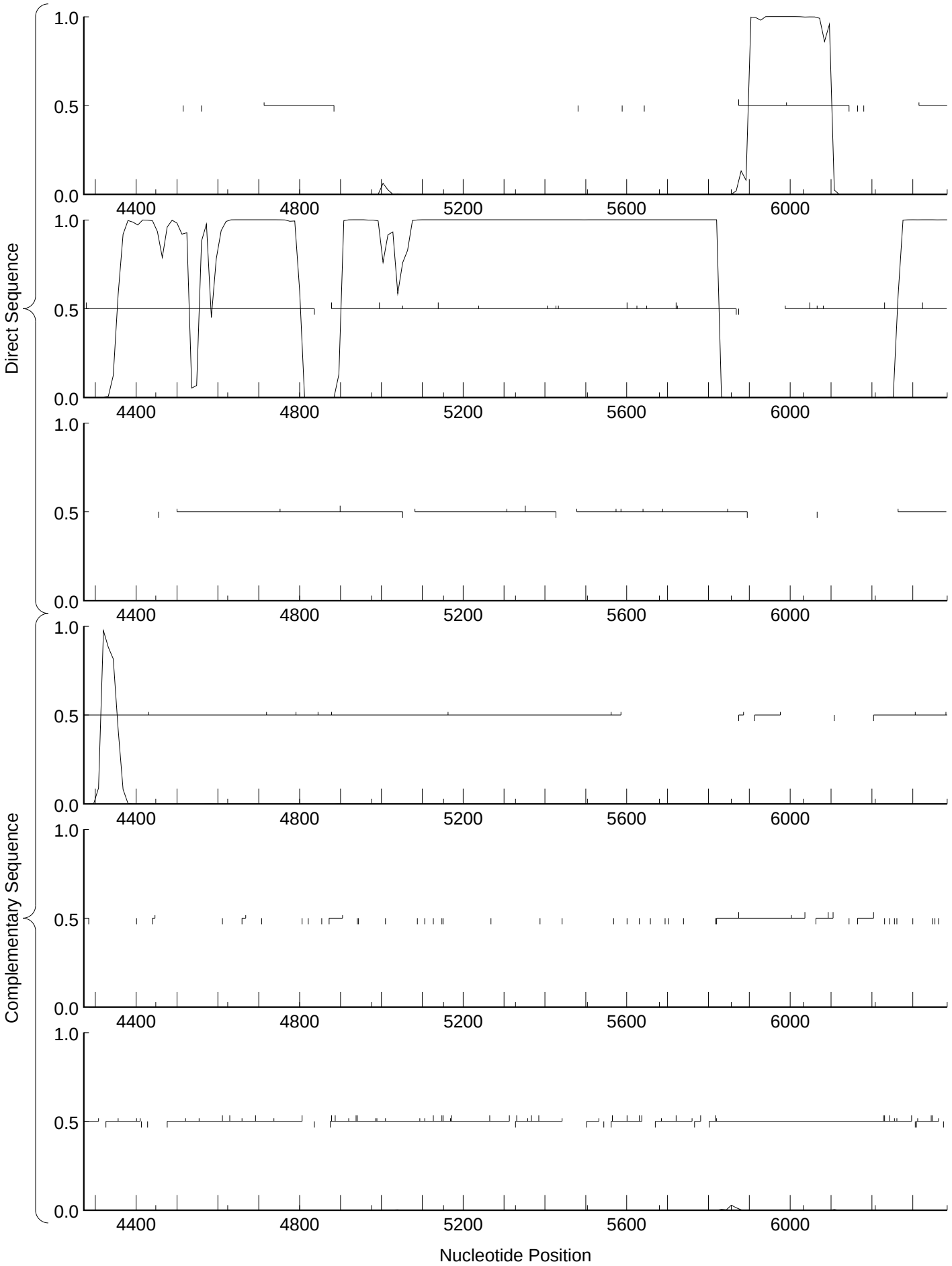
crobacterium phage BeeBee8 complete sequence, 42027 bp, circularly permuted, Cluster EA, Order 4, Window 96, Step 12, 2/21



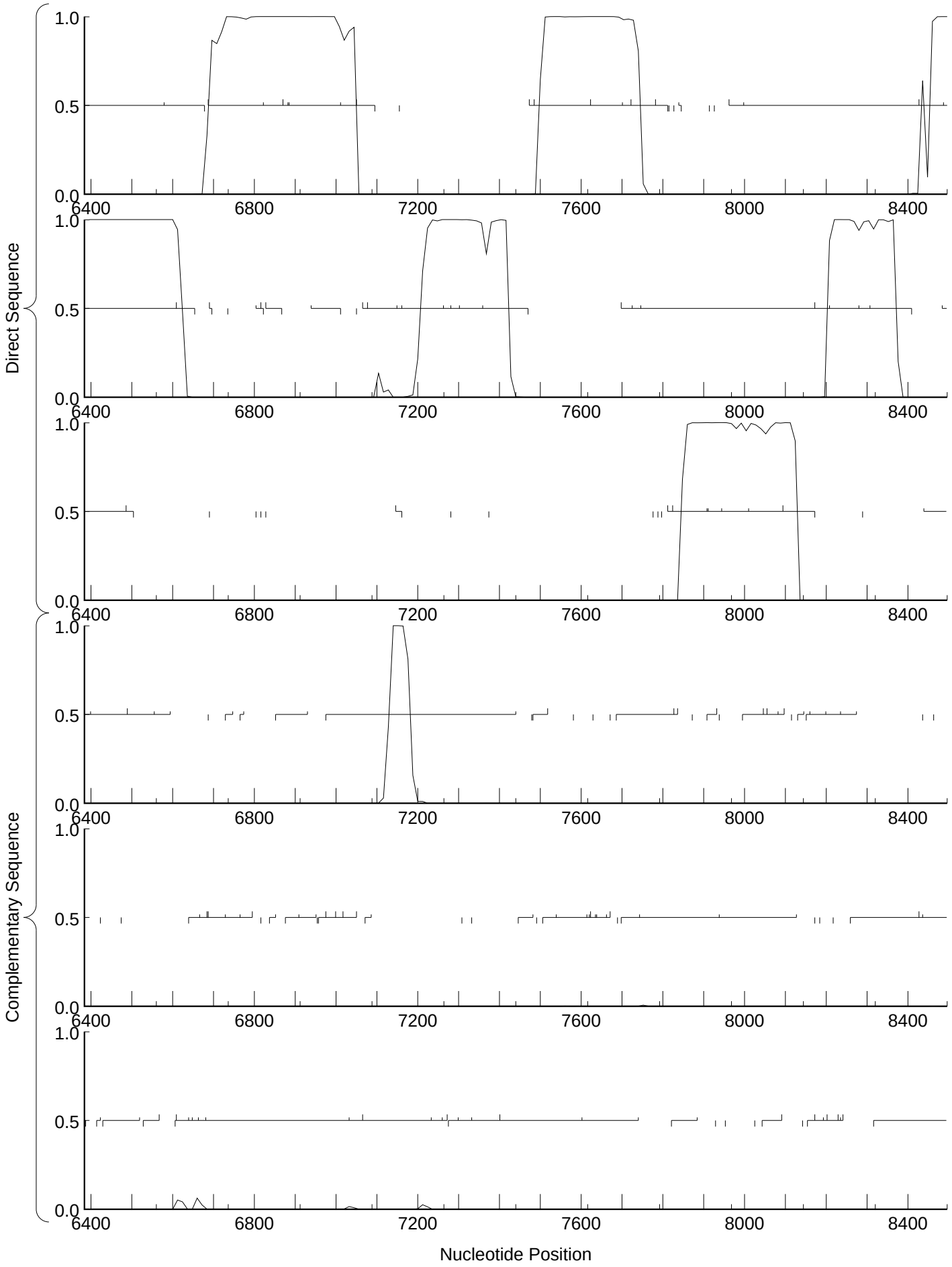
crobacterium phage BeeBee8 complete sequence, 42027 bp, circularly permuted, Cluster EA, Order 4, Window 96, Step 12, 3/21



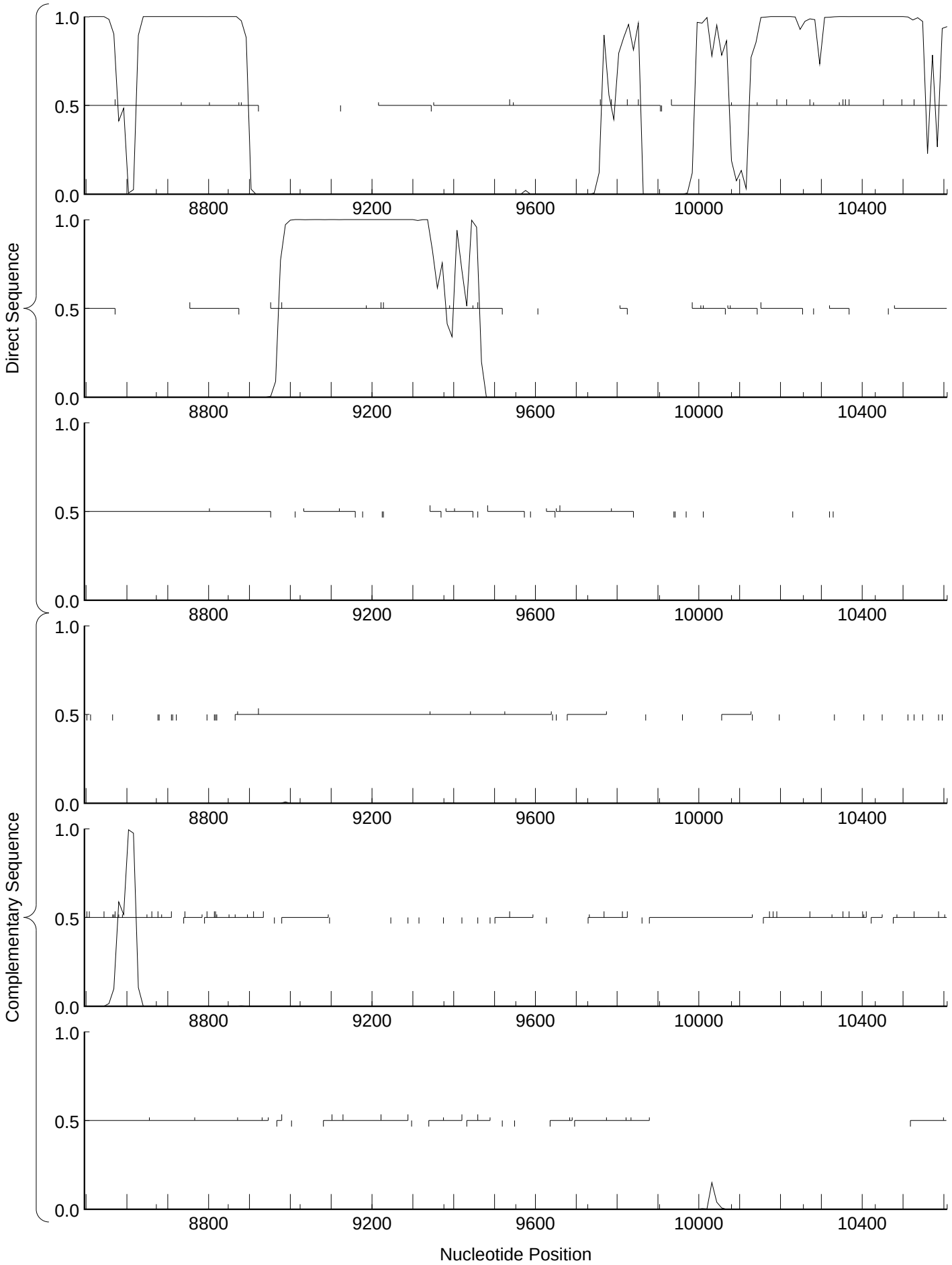
crobacterium phage BeeBee8 complete sequence, 42027 bp, circularly permuted, Cluster EA, Order 4, Window 96, Step 12, 4/21



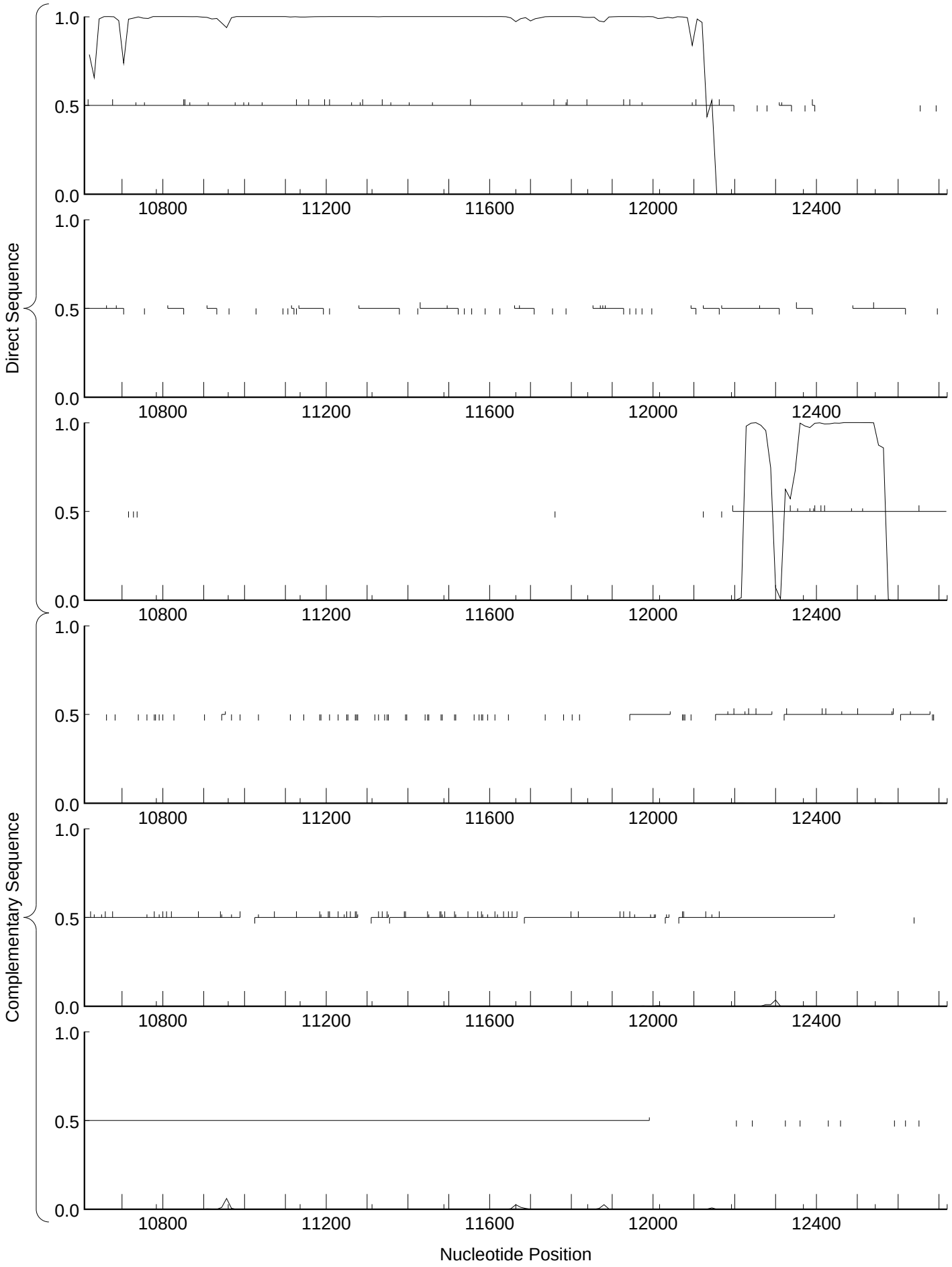
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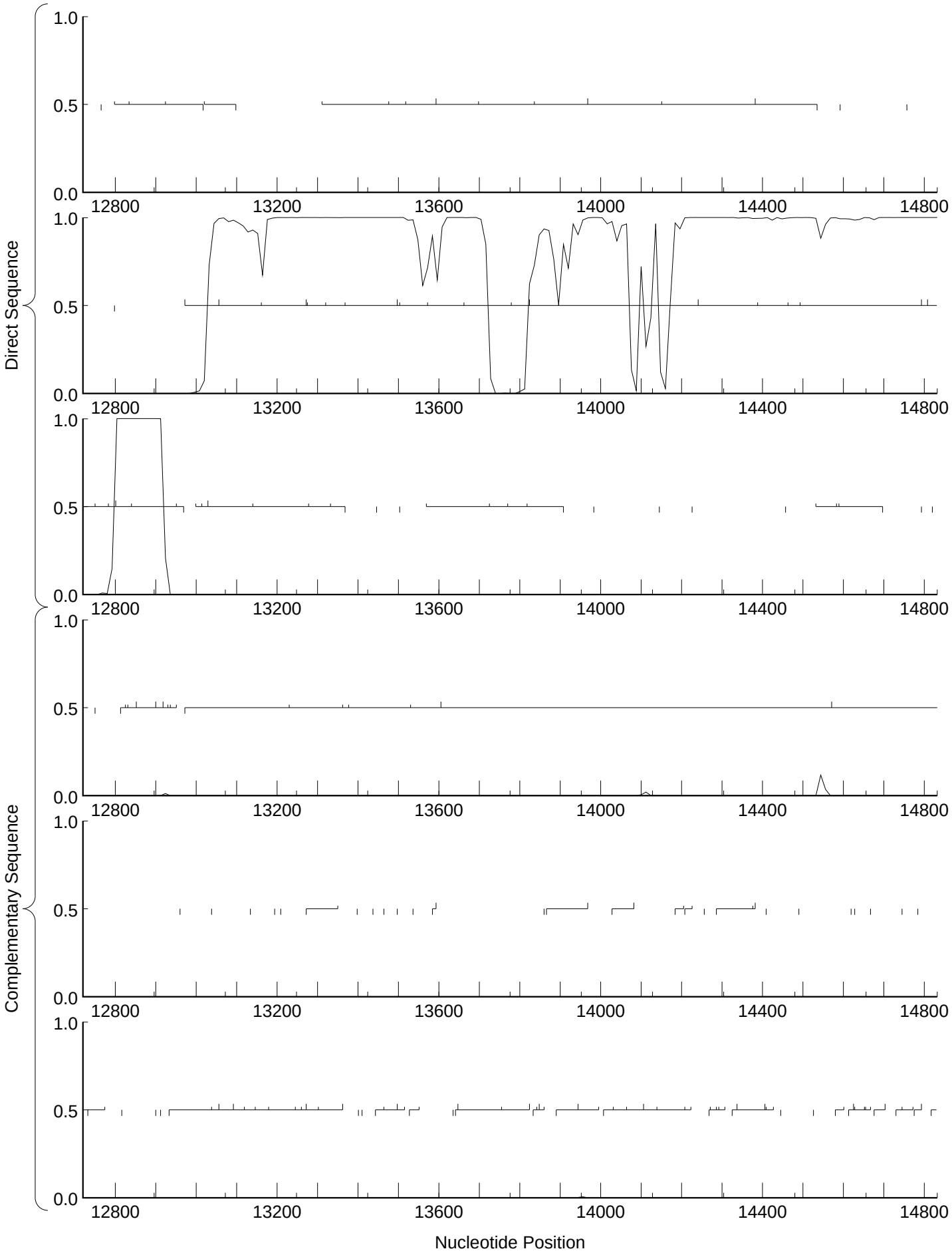
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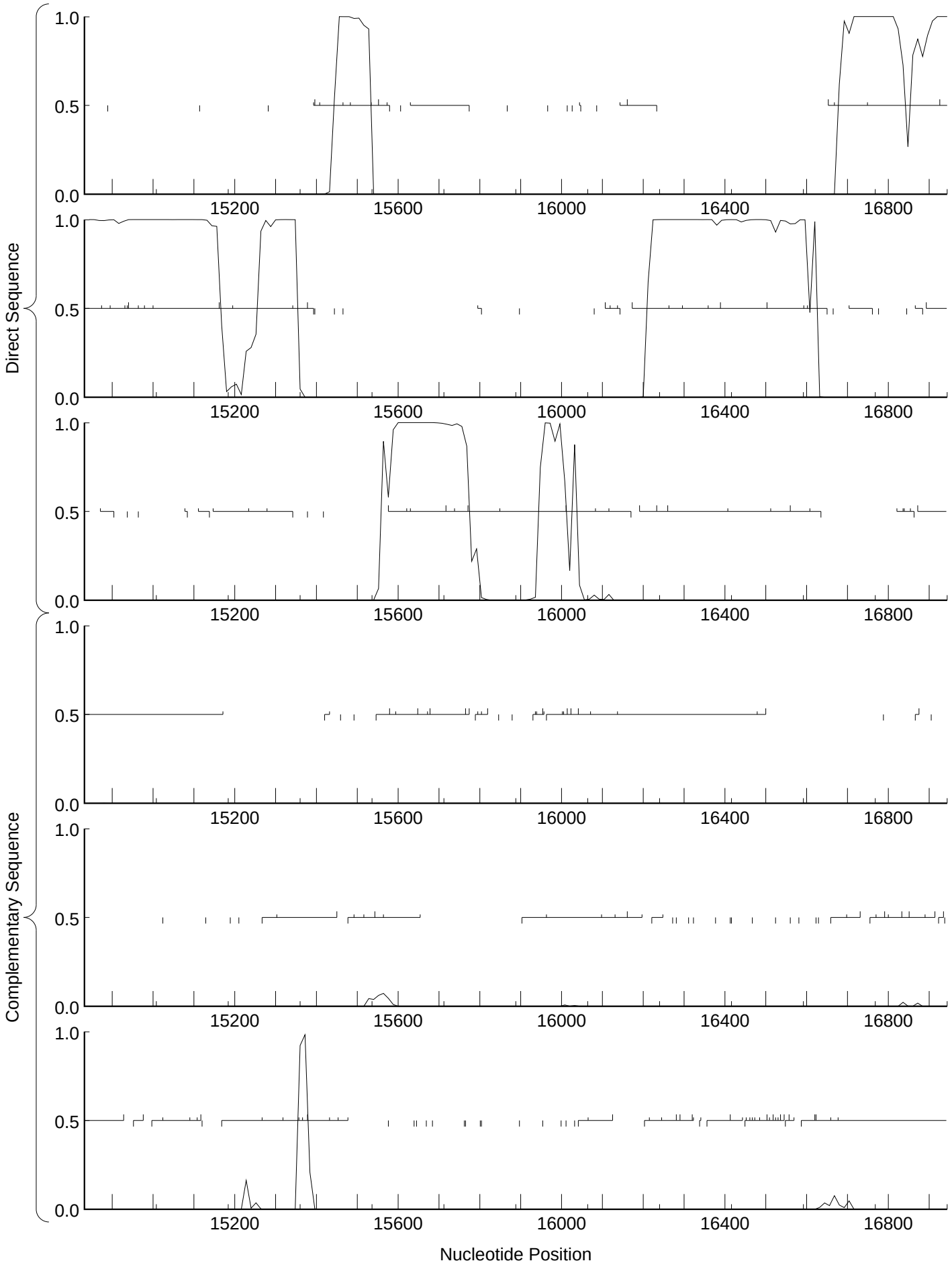
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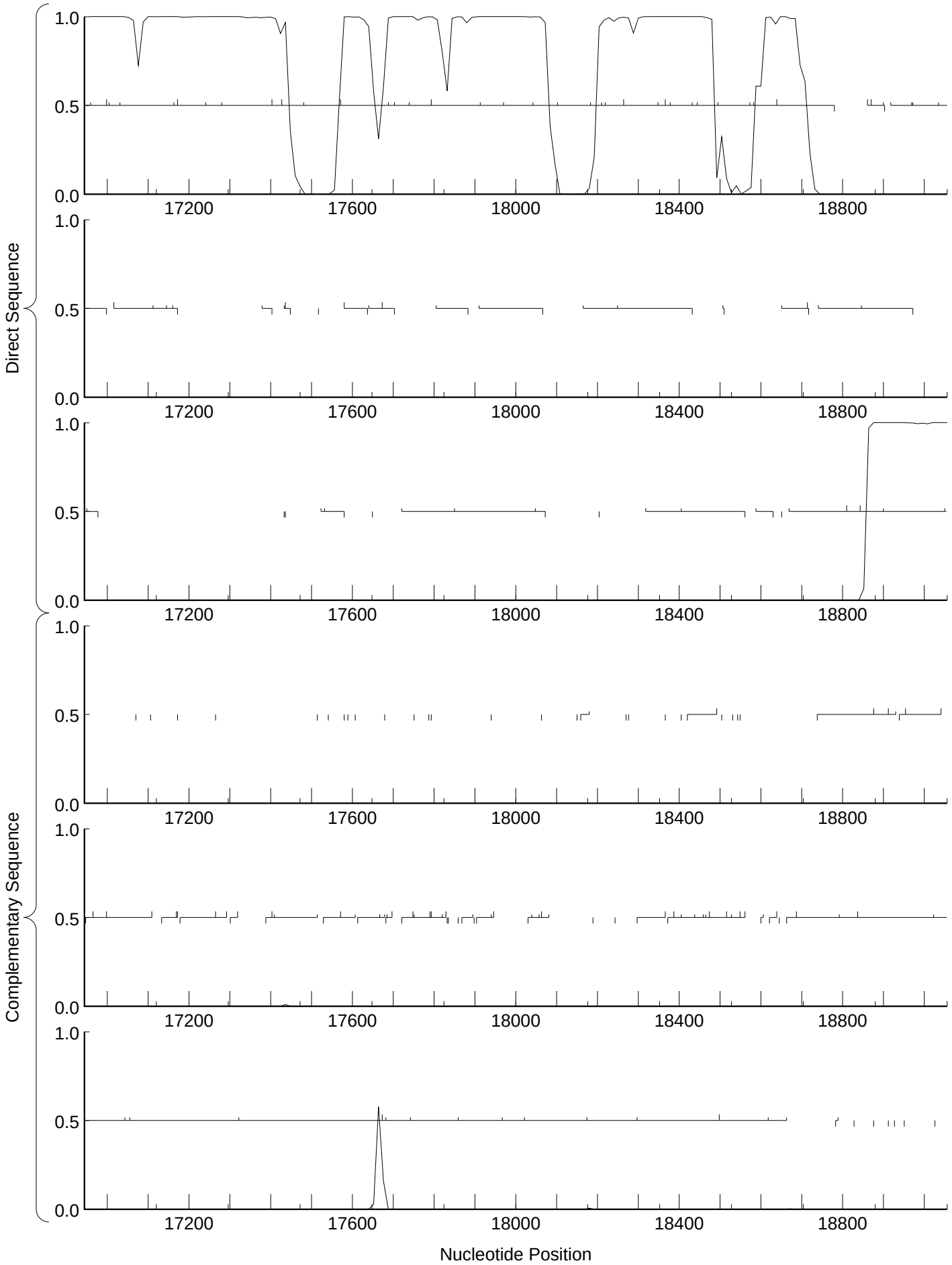
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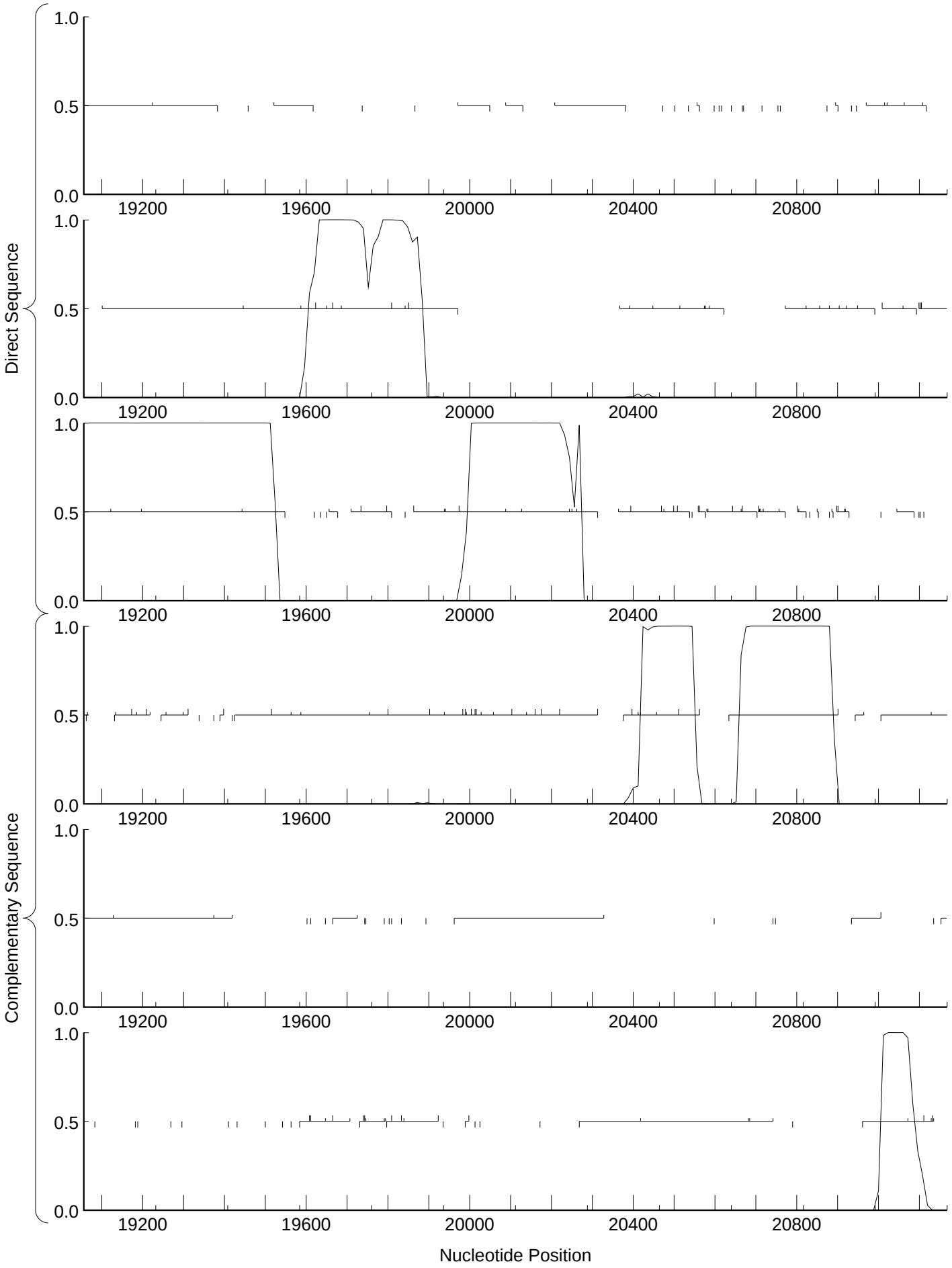
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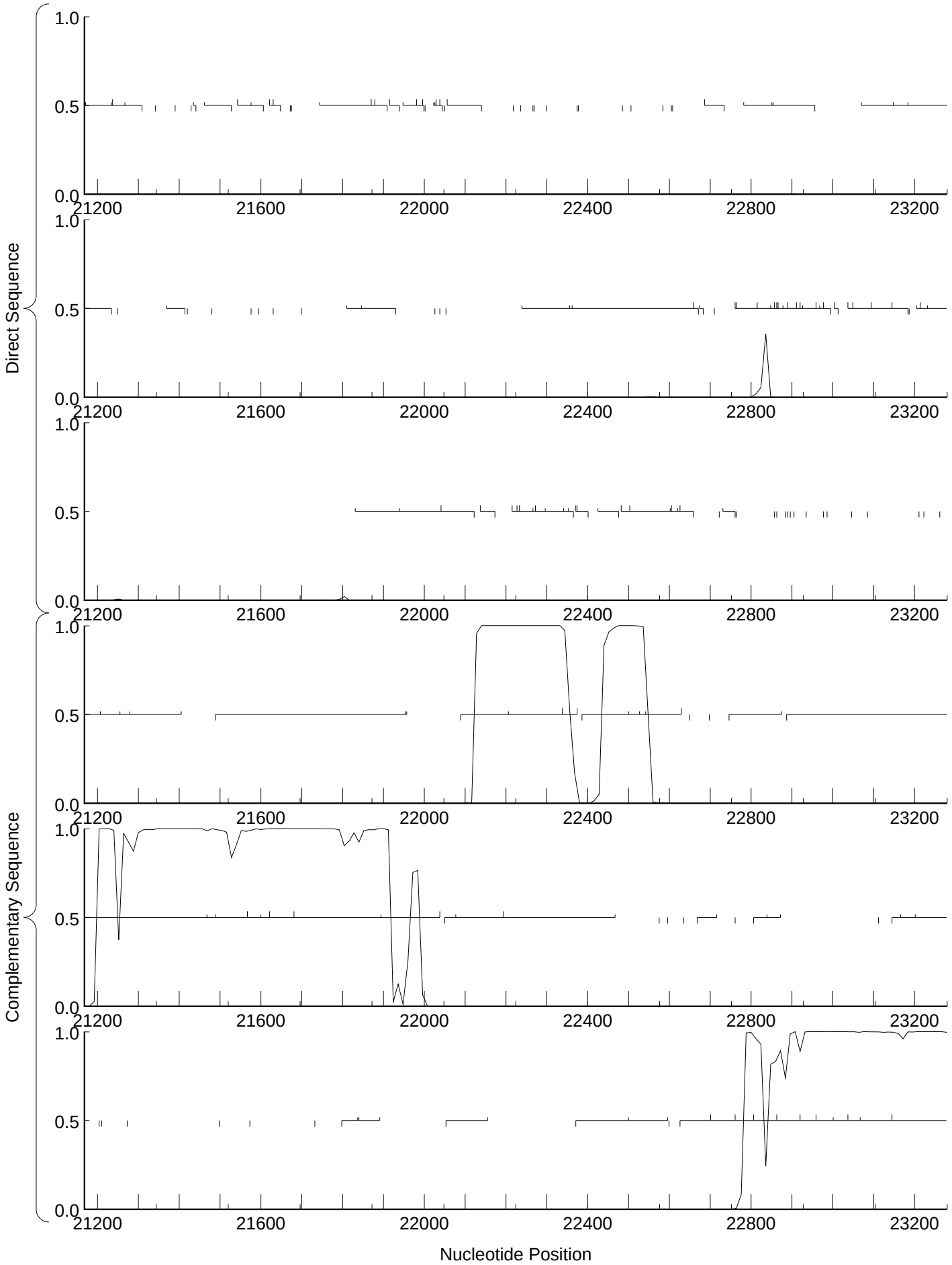
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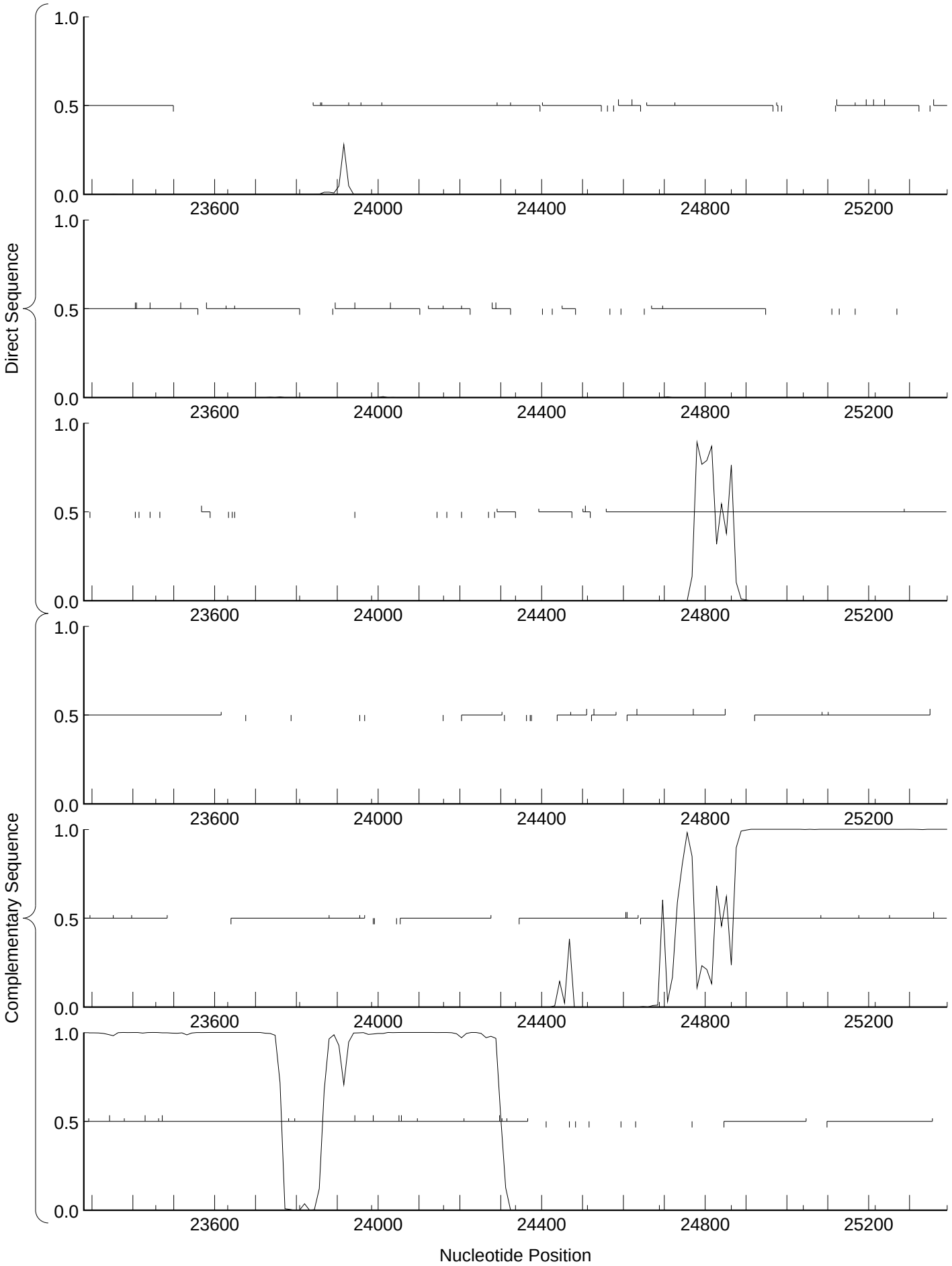
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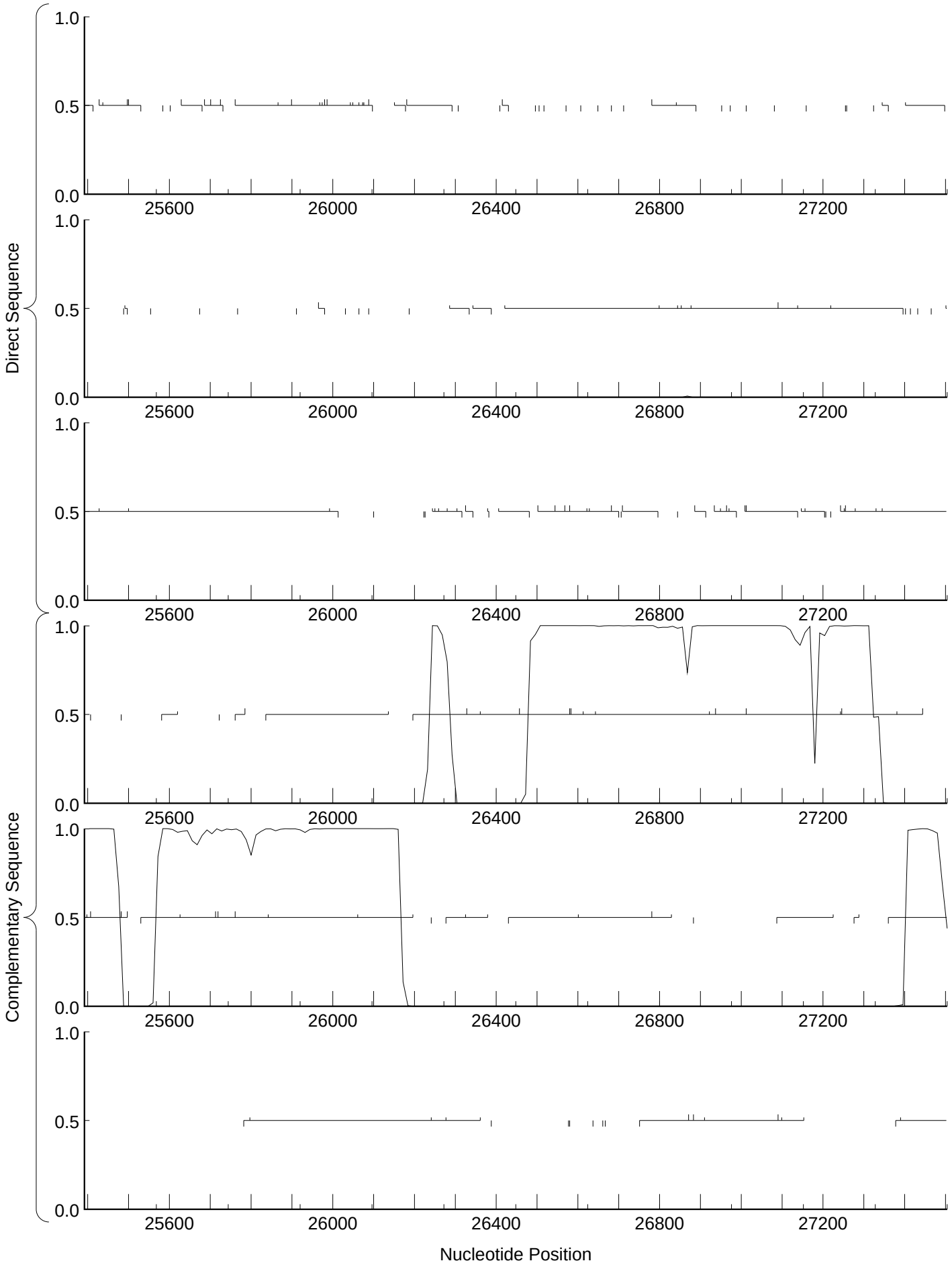
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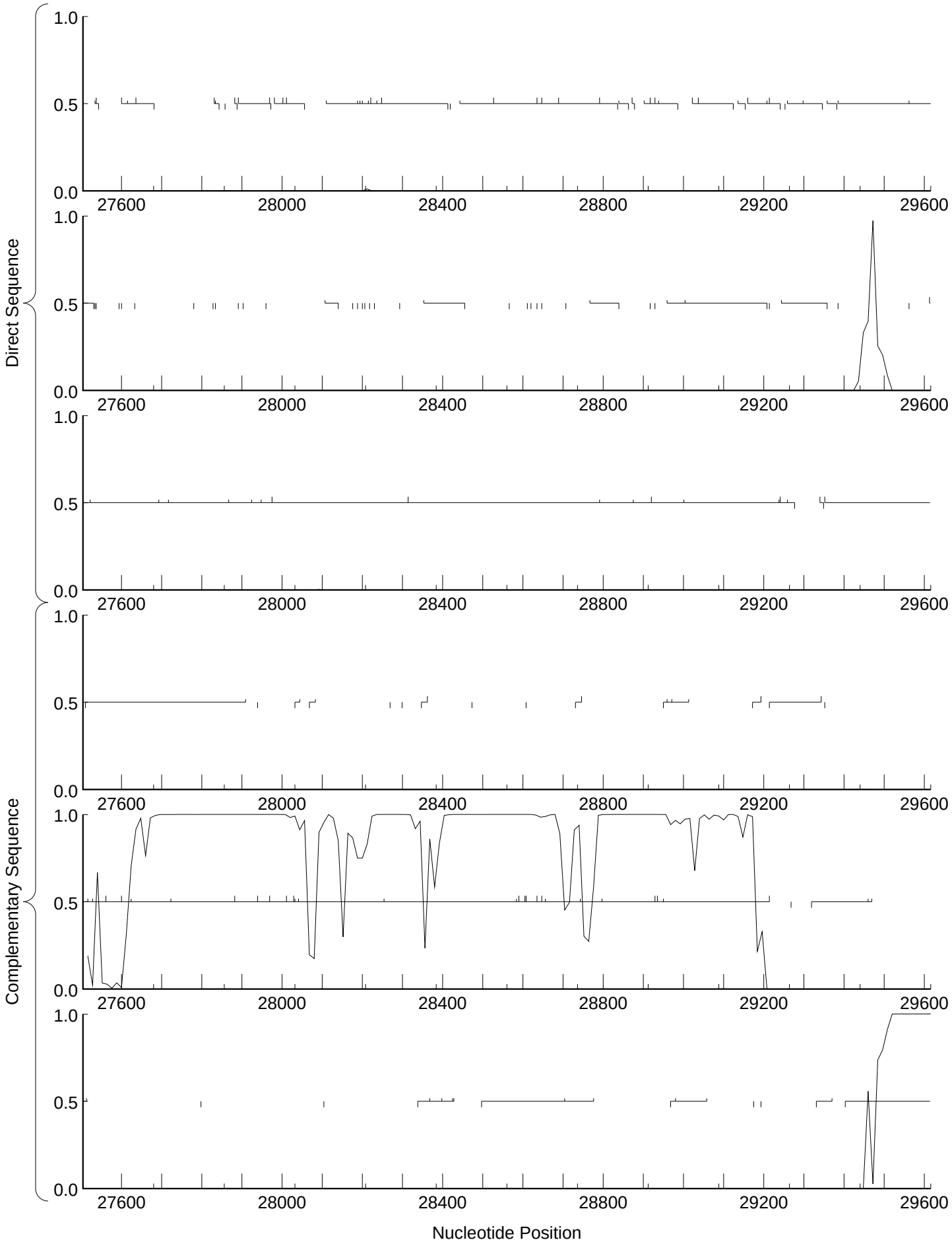
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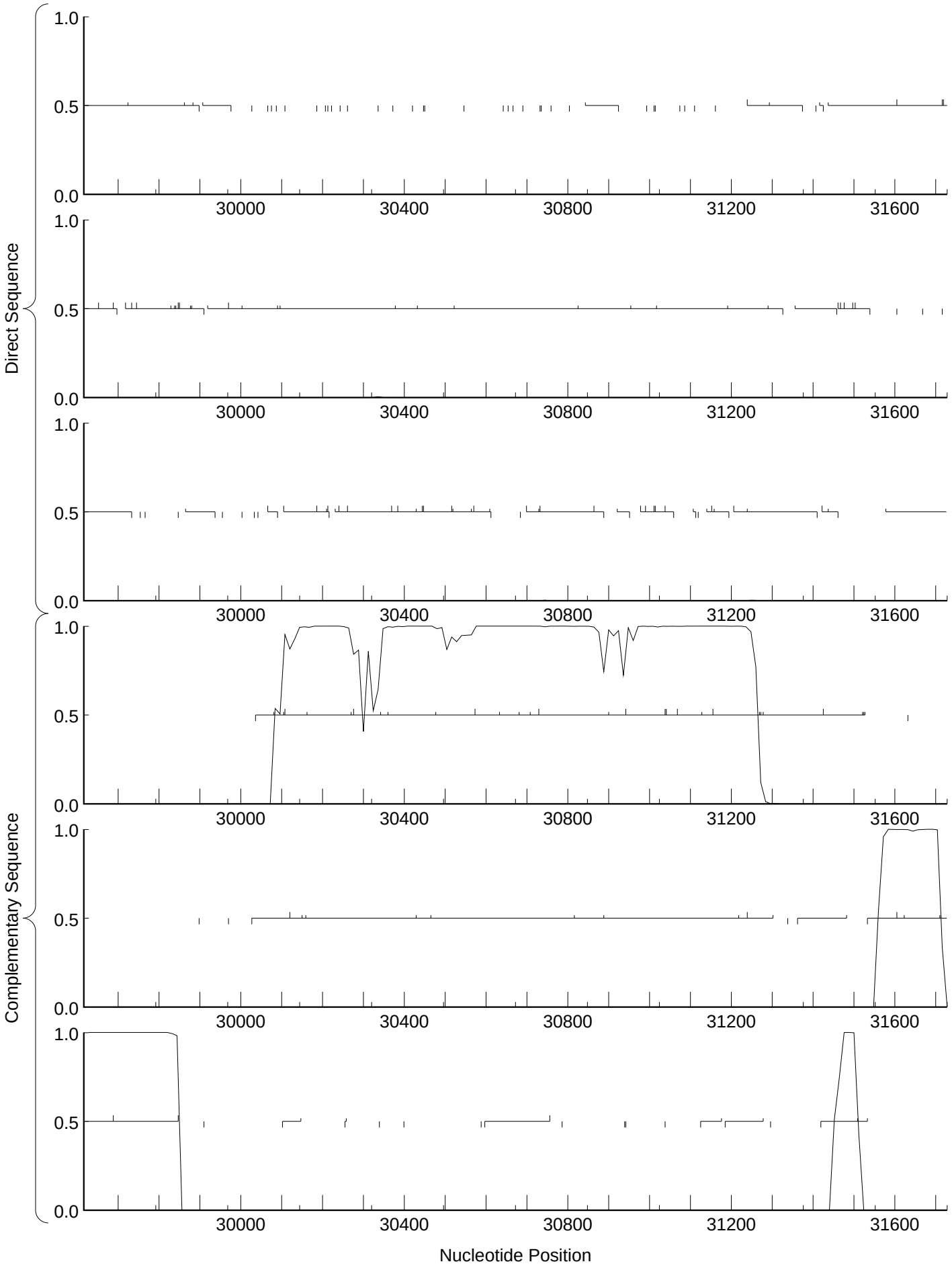
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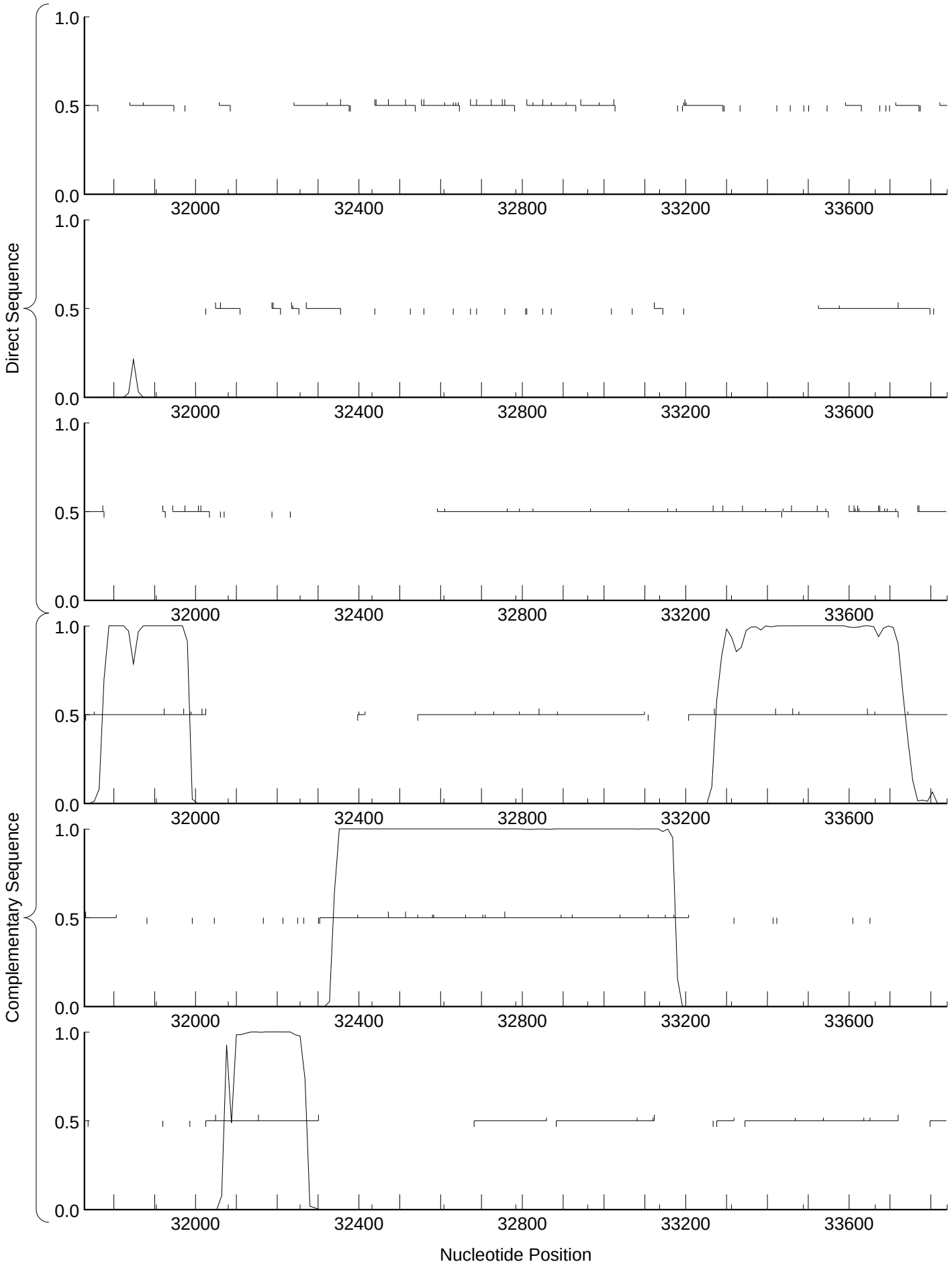


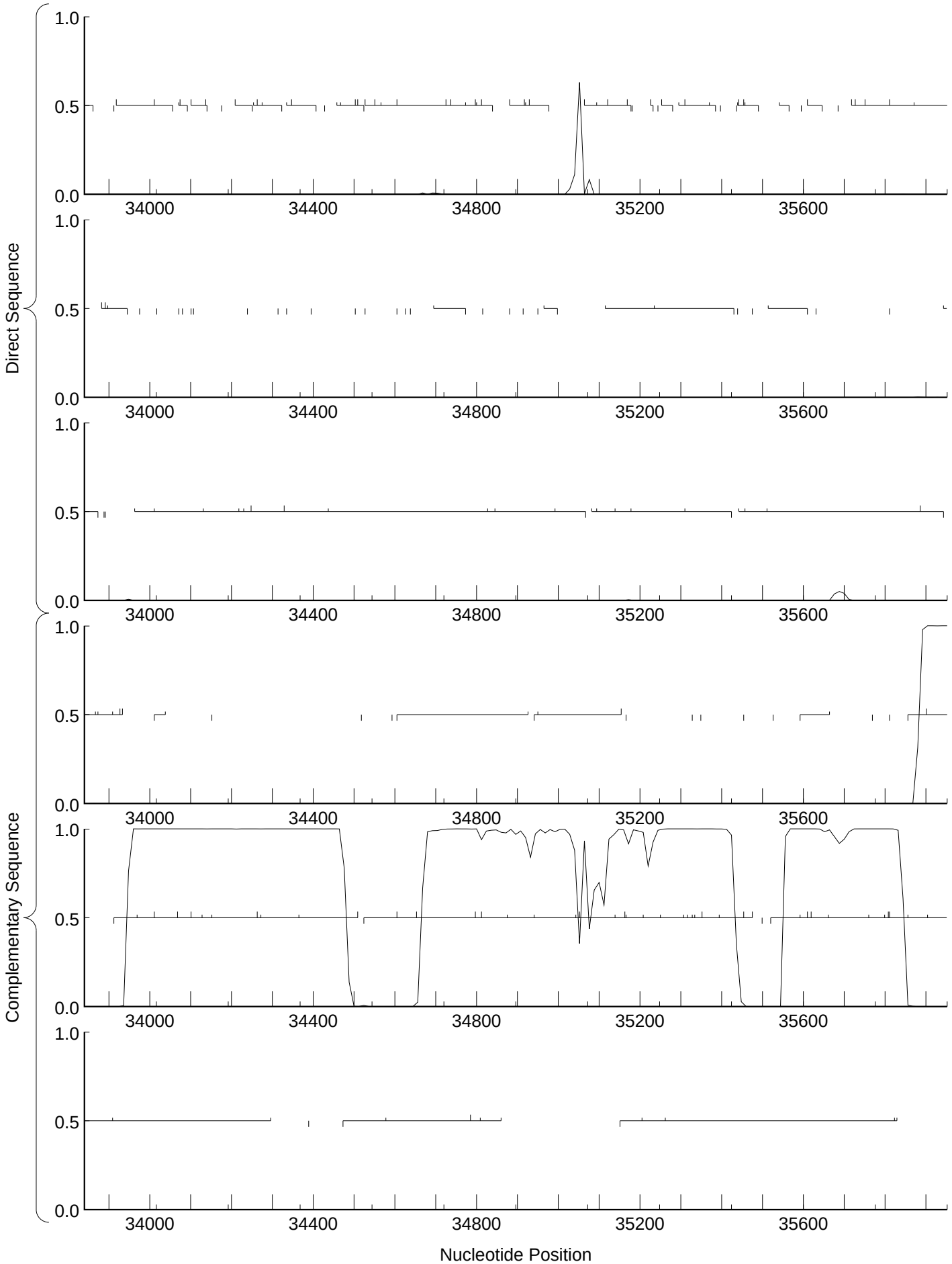
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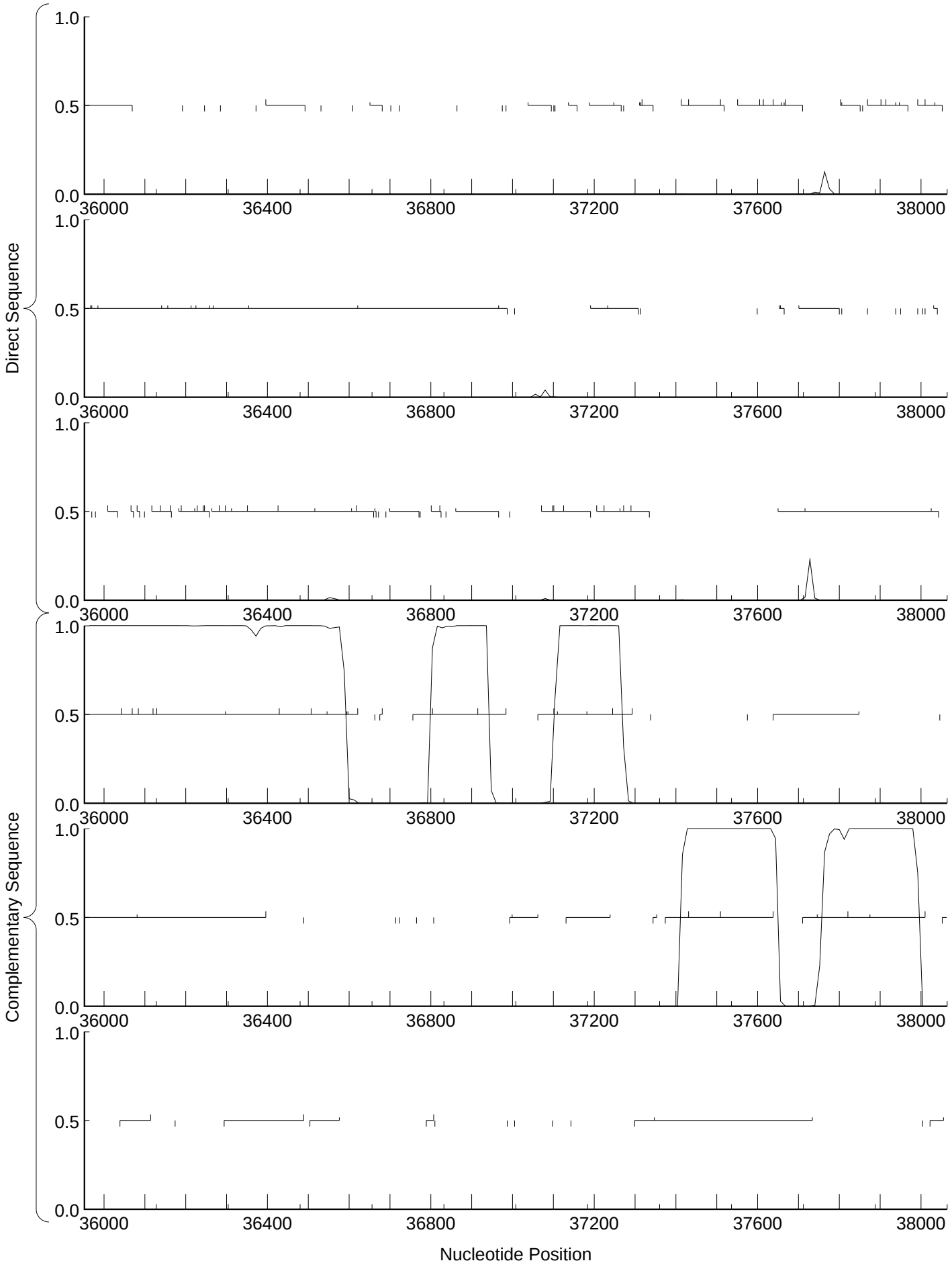
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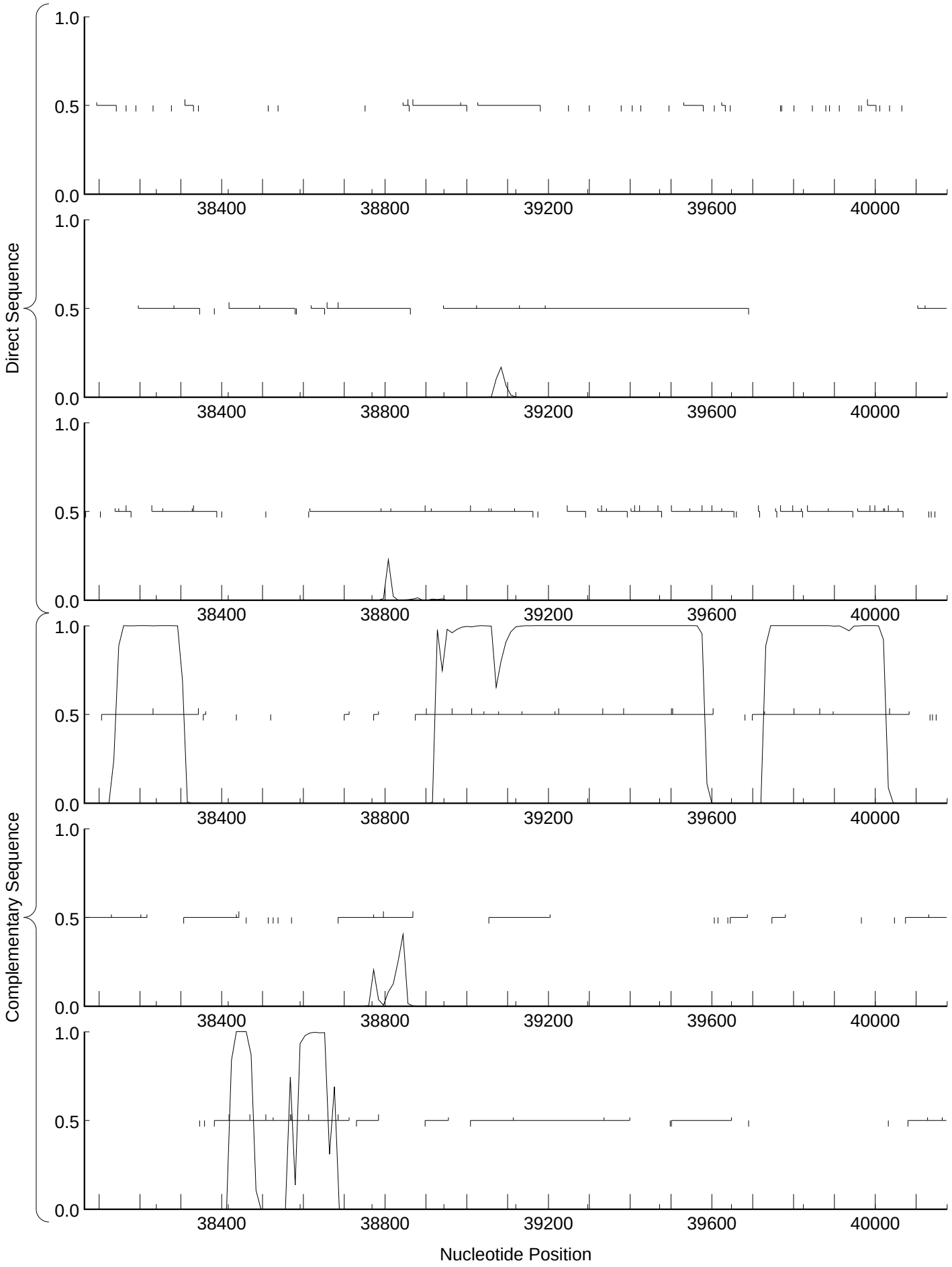

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0.0
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32400
32800
33200
33600
Nucleotide Position

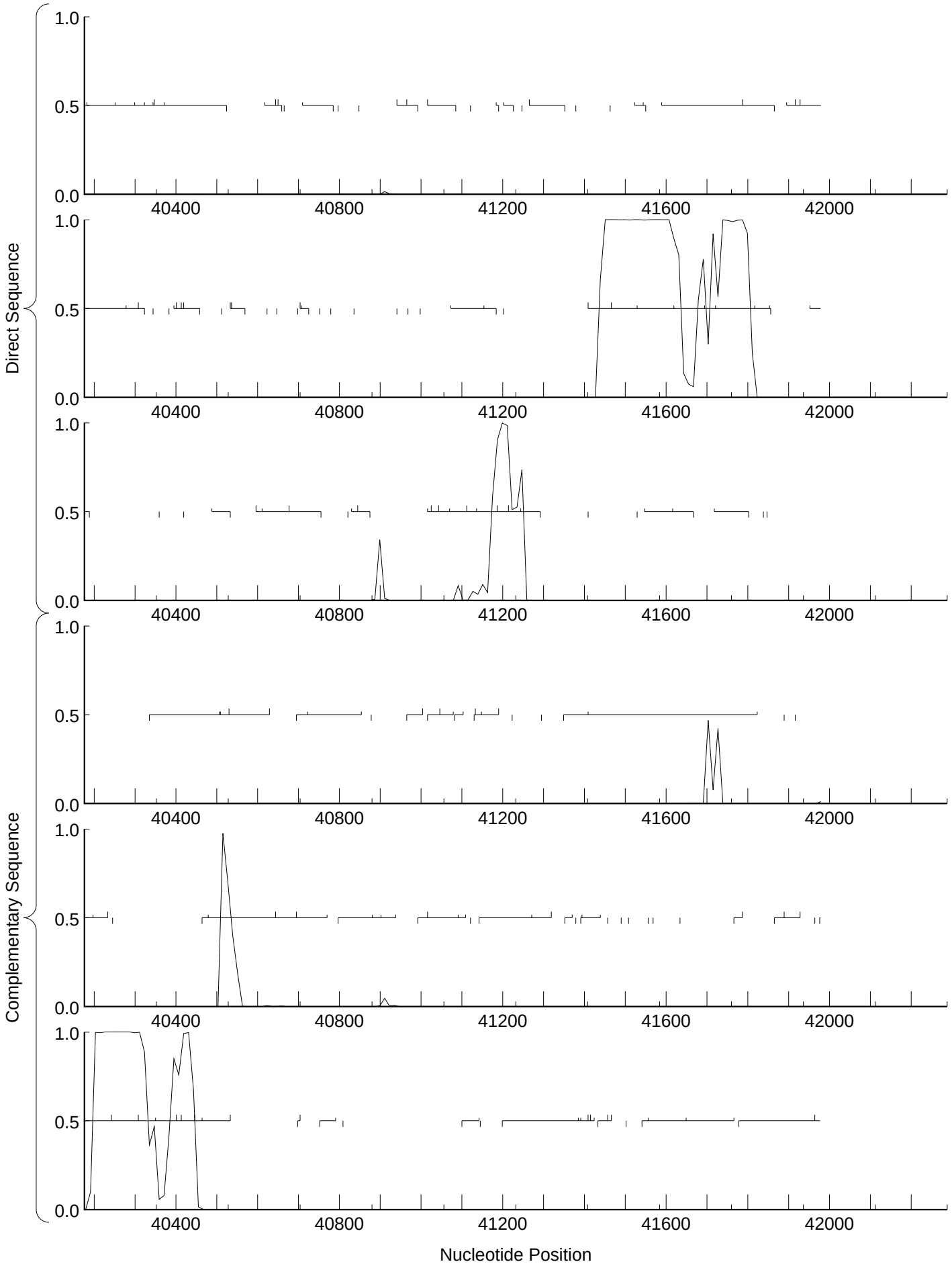

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

robacterium phage BeeBee8 complete sequence, 42027 bp, circularly permuted, Cluster EA, Order 4, Window 96, Step 12, 19/21



robacterium phage BeeBee8 complete sequence, 42027 bp, circularly permuted, Cluster EA, Order 4, Window 96, Step 12, 20/21




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
40400
40800
41200
41600
42000
Nucleotide Position