GeneMark

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

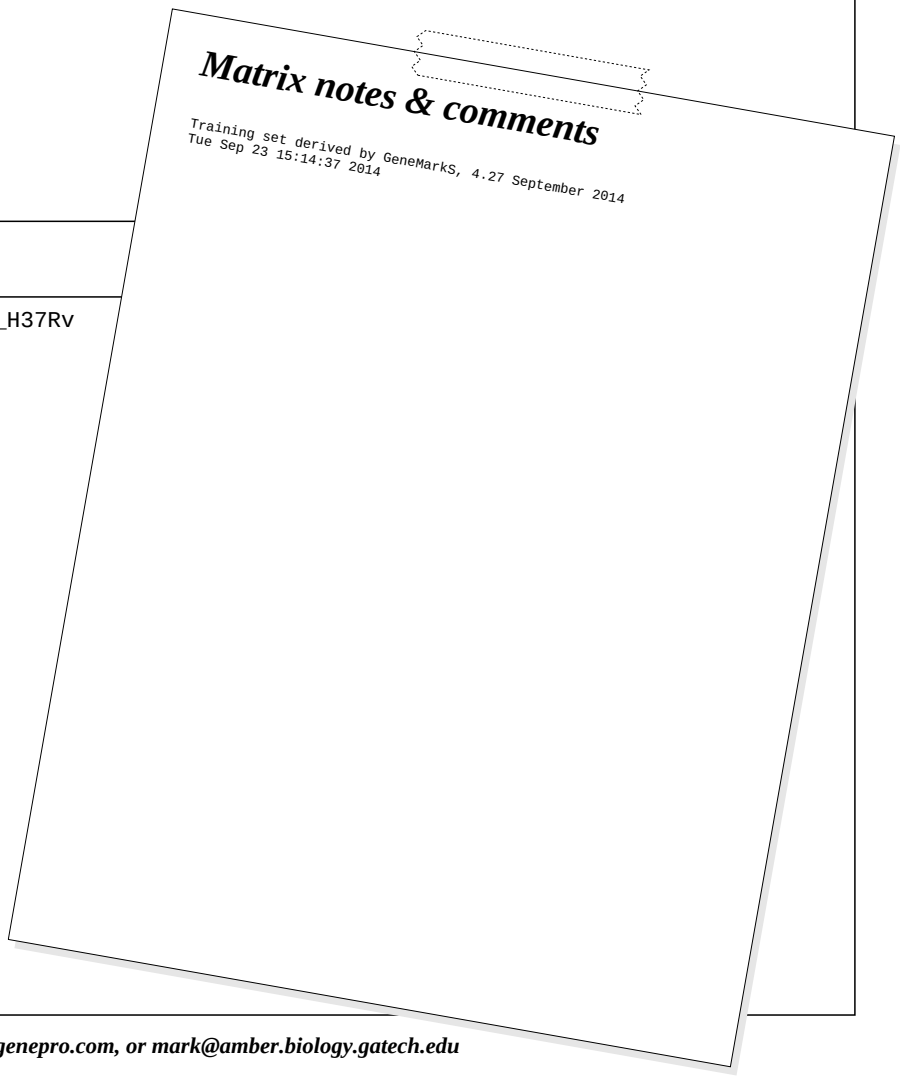
PROGRAM INFORMATION

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Sequence          :  Mycobacterium phage ItsyBitsy1 complete sequence, 67570 bp, circularly permuted, Cluster B2
Analysis Date     :  3/29/24 at 4:19:02
Pages             :  33
Sequence Length   :  67570 bp
GC Content        :   68.93%

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

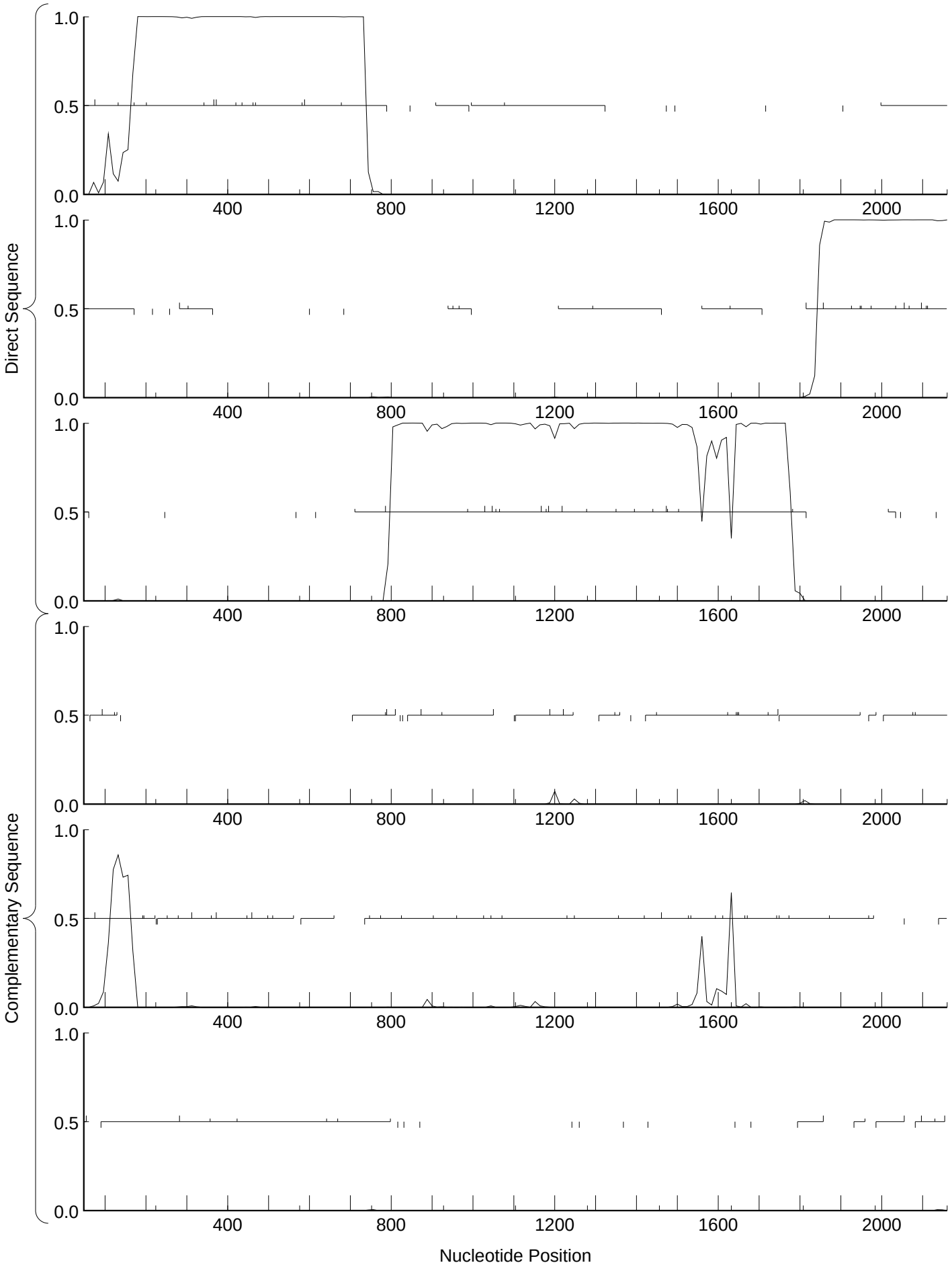
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Matrix      :  Mycobacterium_tuberculosis_H37Rv
Author      :  -

Order       :  4
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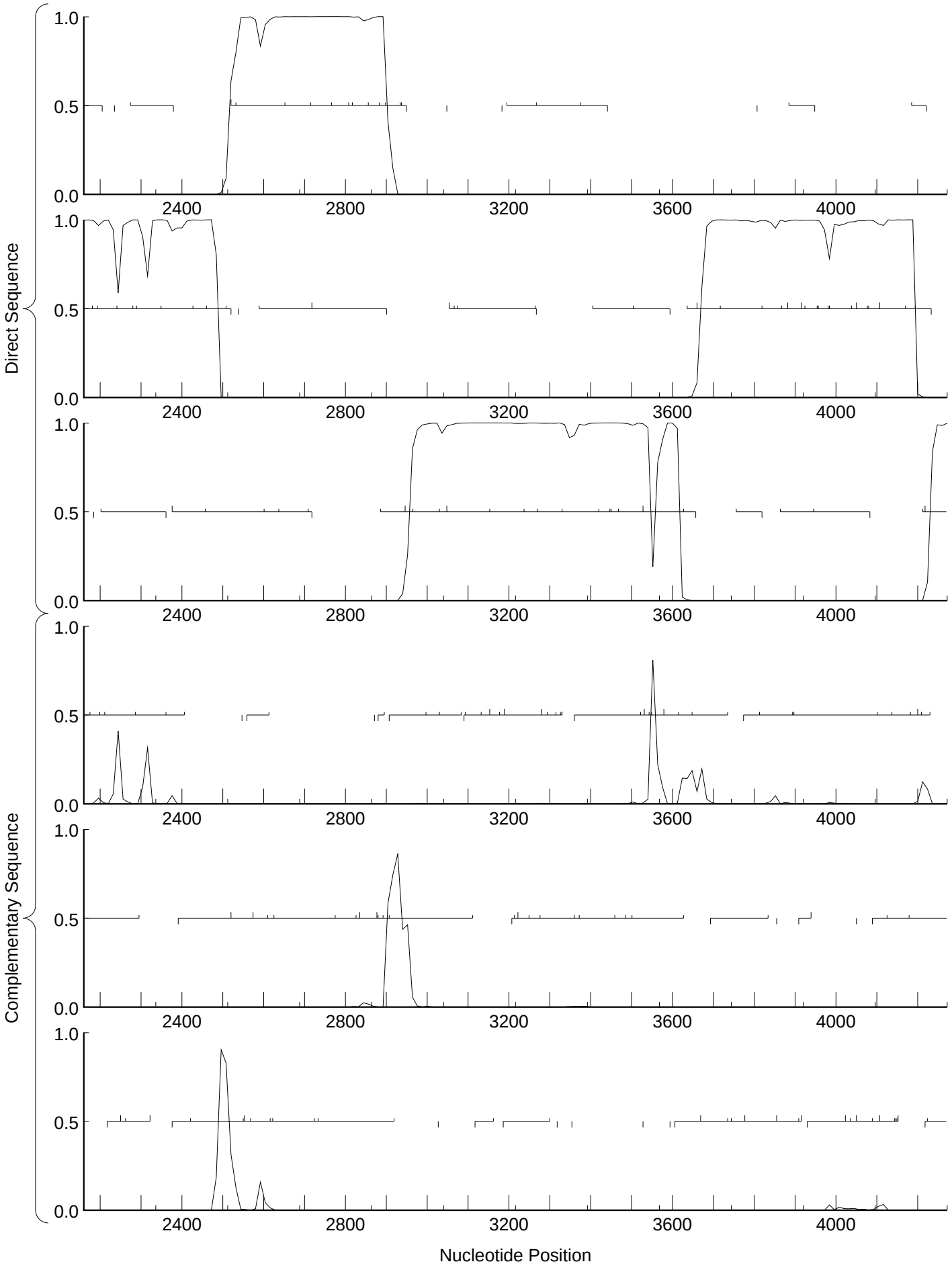

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

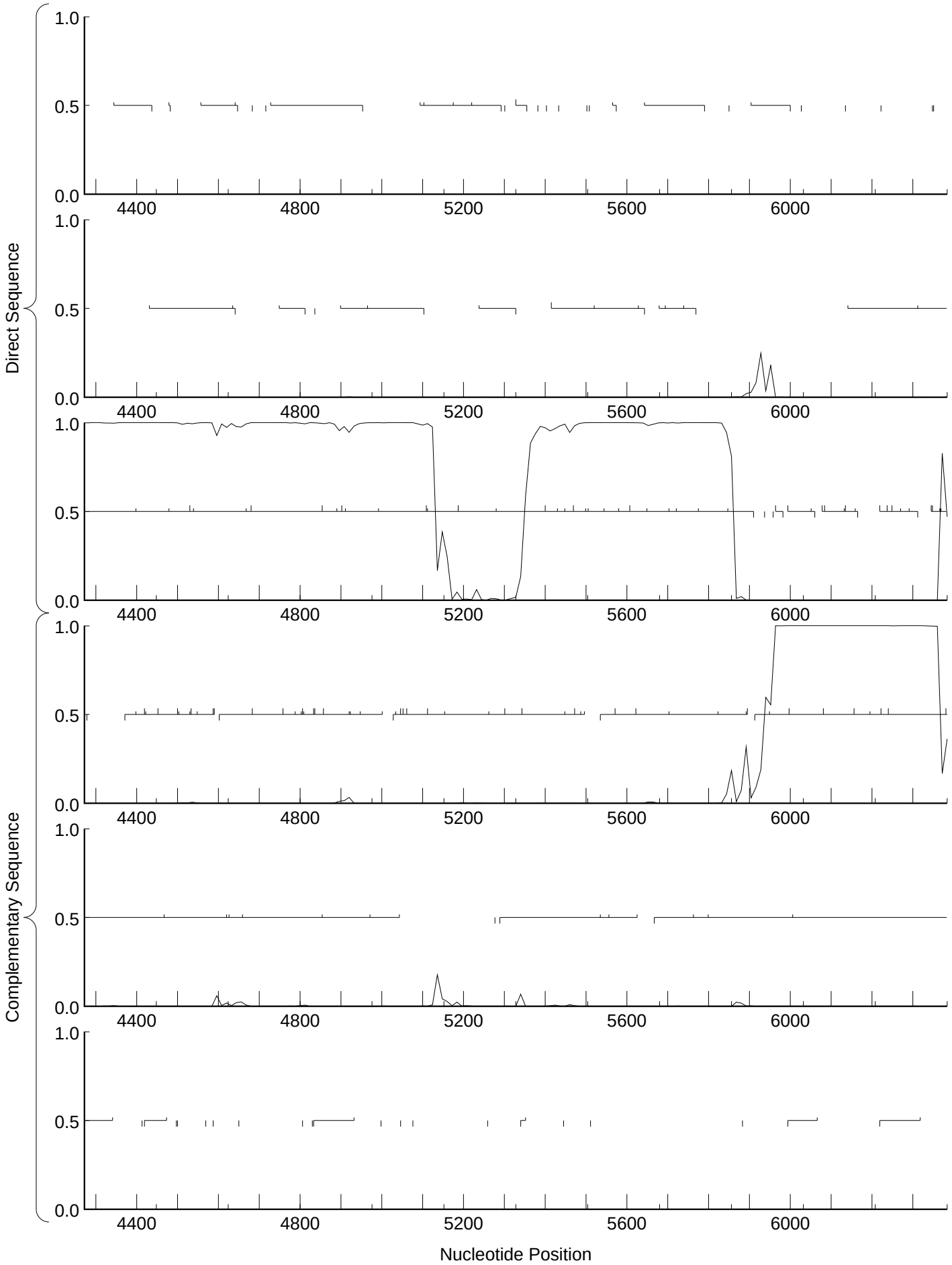
ycobacterium phage ItsyBitsy1 complete sequence, 67570 bp, circularly permuted, Cluster B2, Order 4, Window 96, Step 12, 2/33

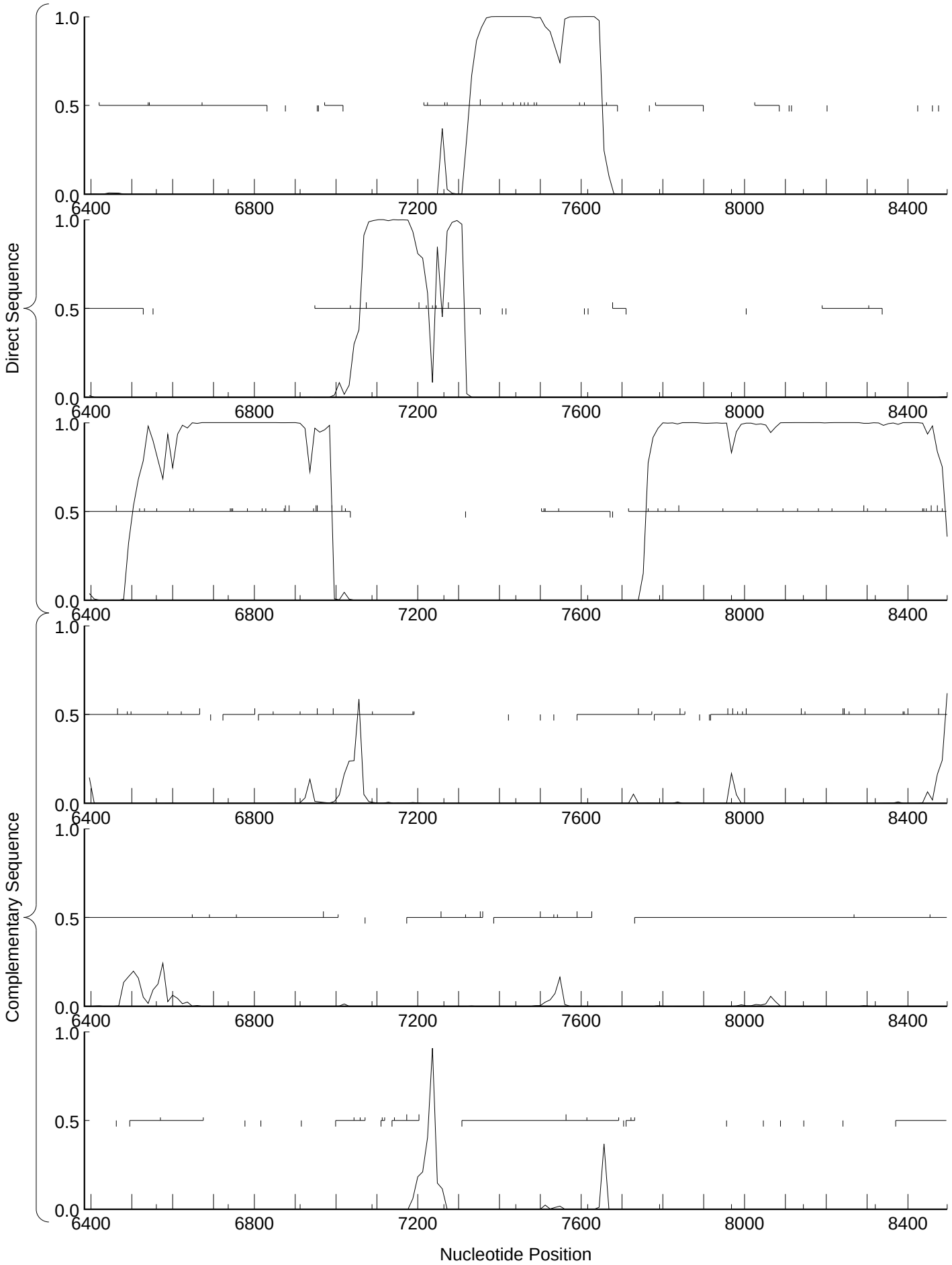


ycobacterium phage ItsyBitsy1 complete sequence, 67570 bp, circularly permuted, Cluster B2, Order 4, Window 96, Step 12, 3/33

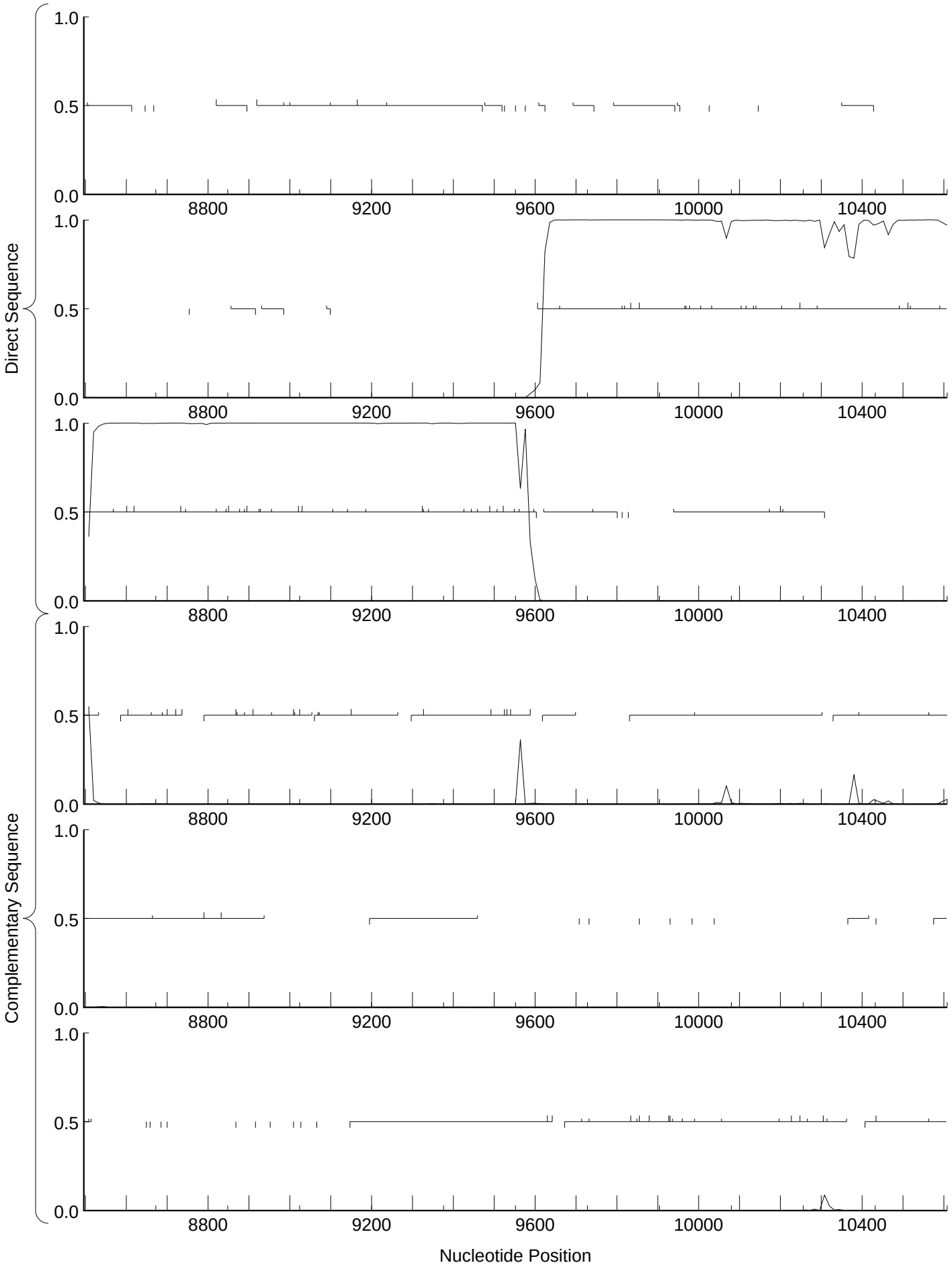


ycobacterium phage ItsyBitsy1 complete sequence, 67570 bp, circularly permuted, Cluster B2, Order 4, Window 96, Step 12, 4/33

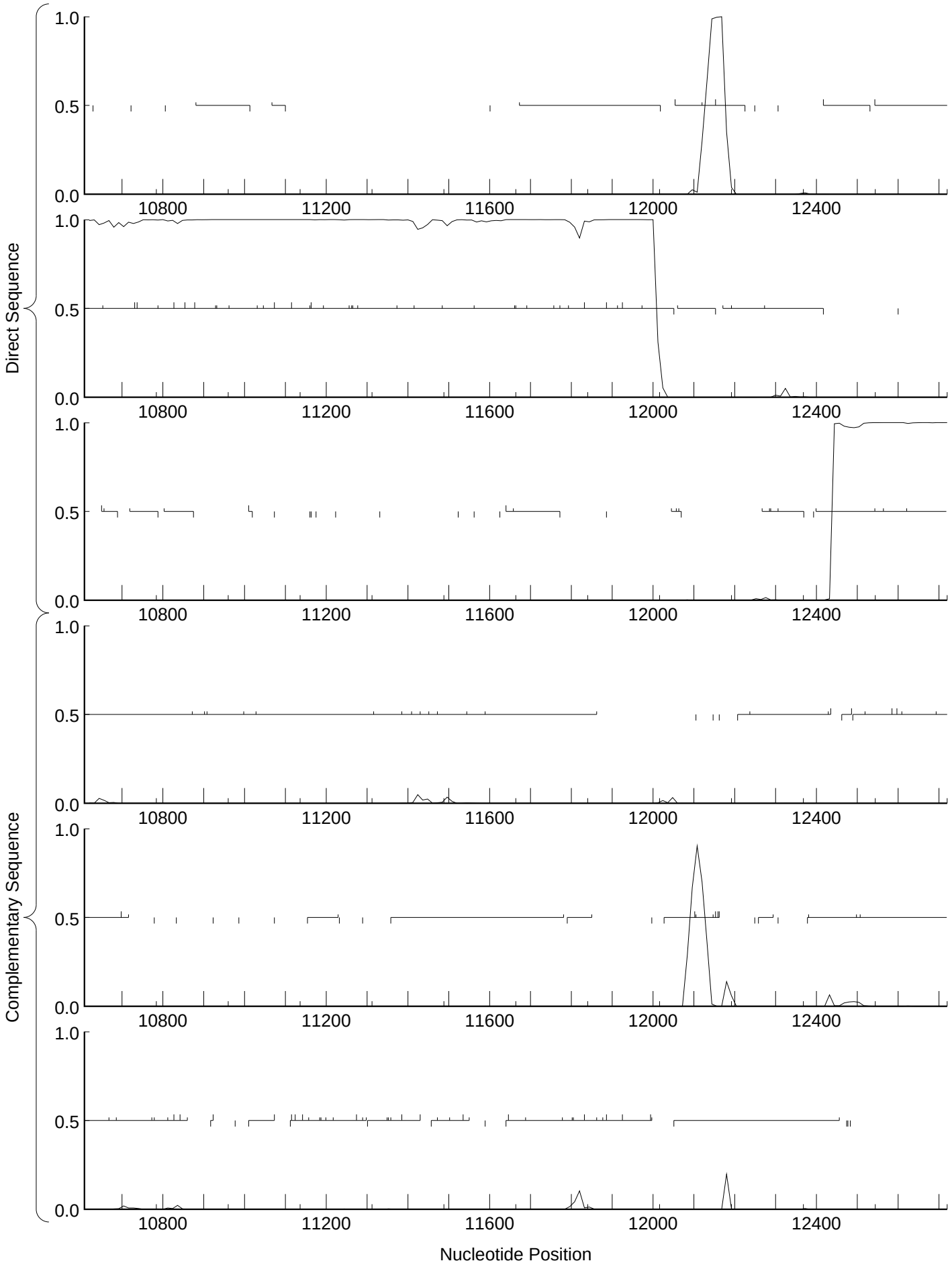



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
6400
6800
7200
7600
8000
8400
Nucleotide Position

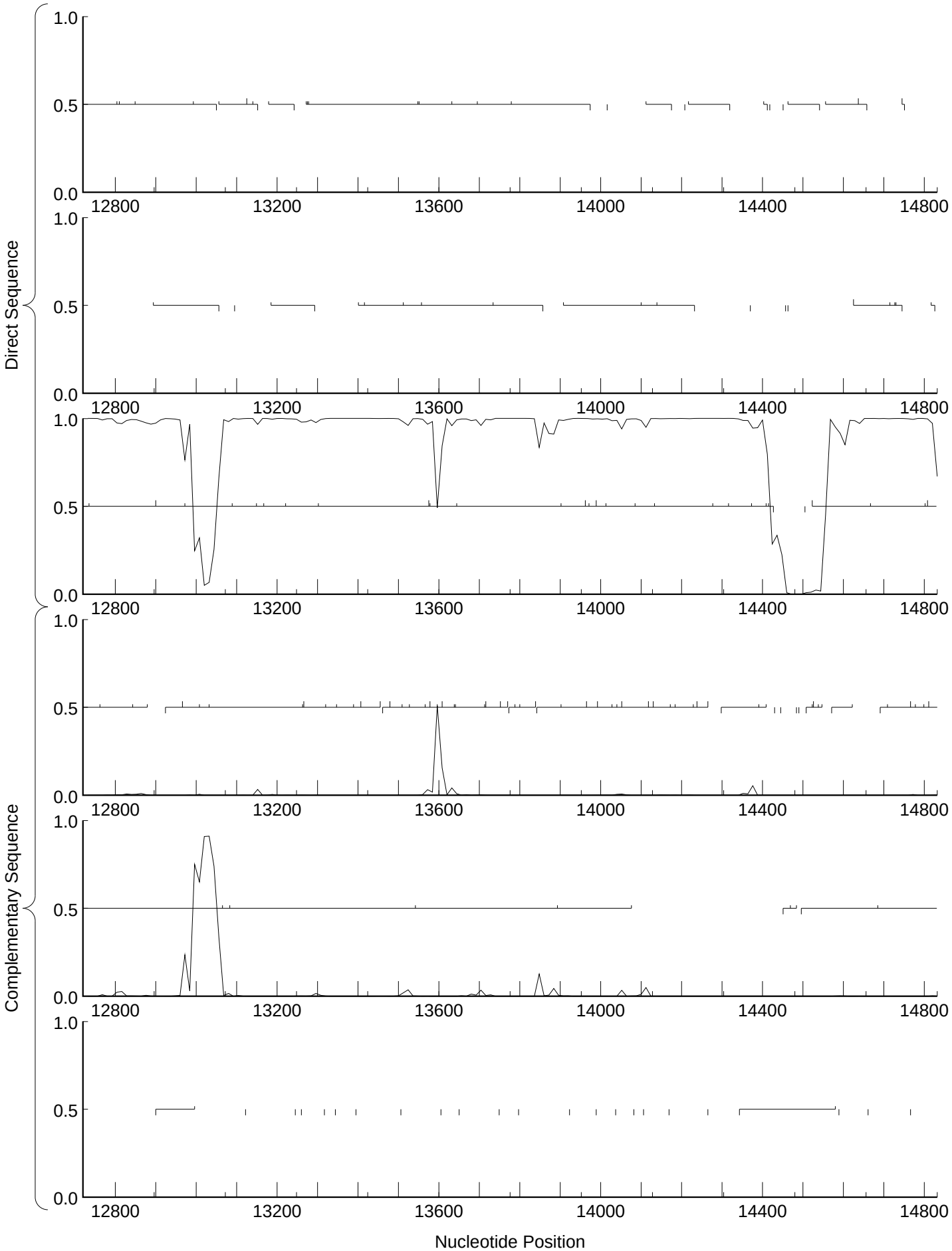
ycobacterium phage ItsyBitsy1 complete sequence, 67570 bp, circularly permuted, Cluster B2, Order 4, Window 96, Step 12, 6/33



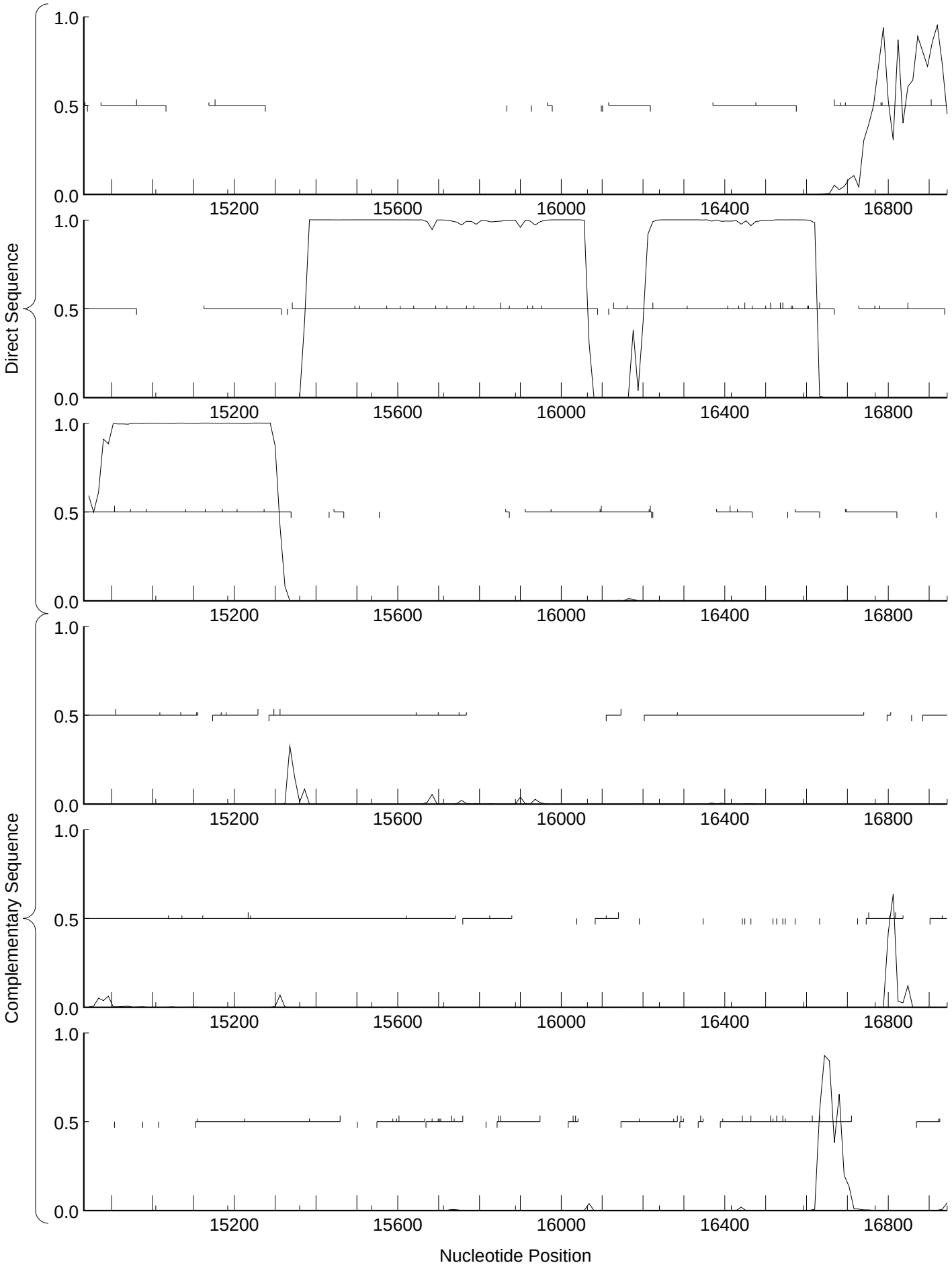
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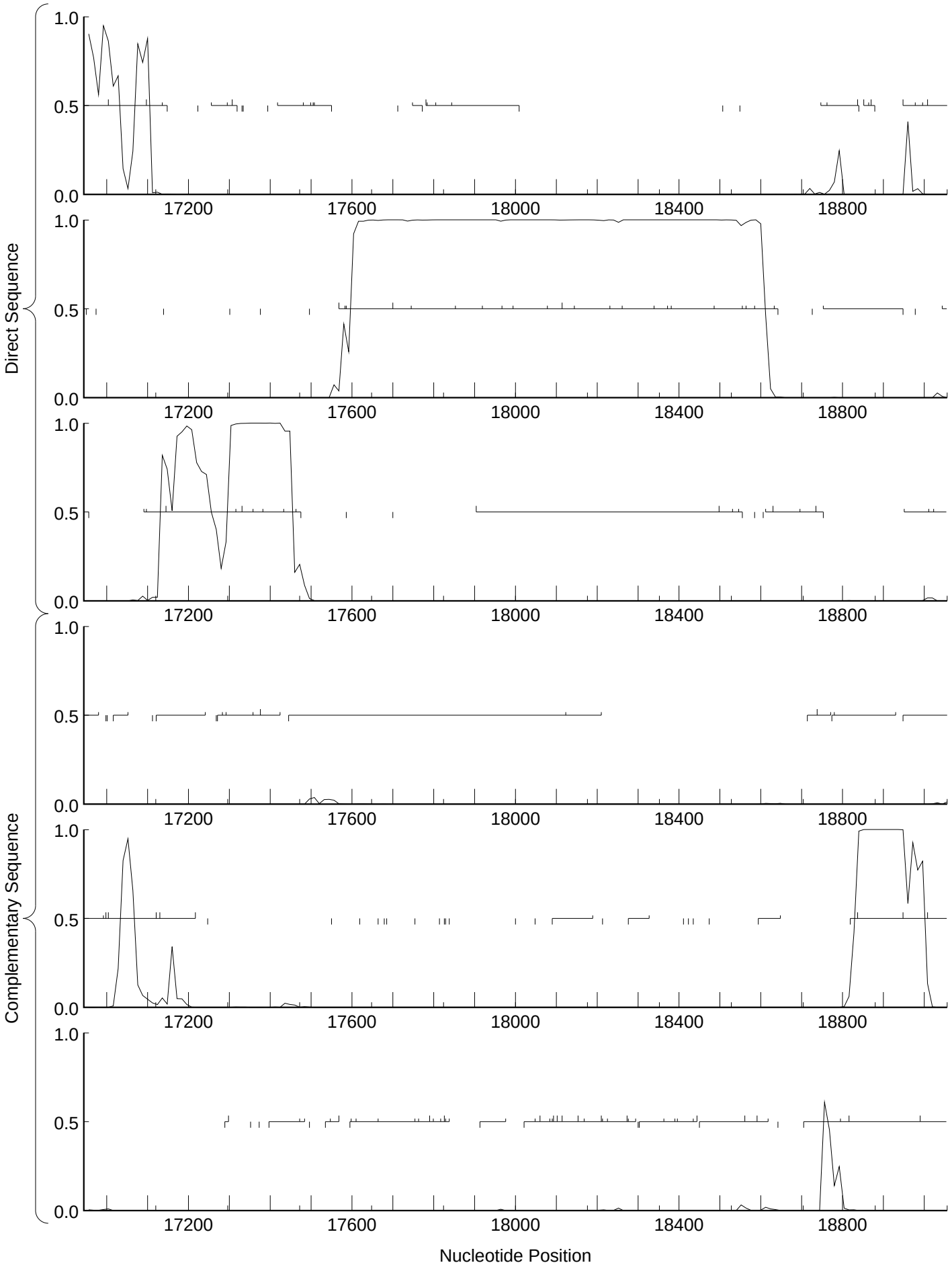
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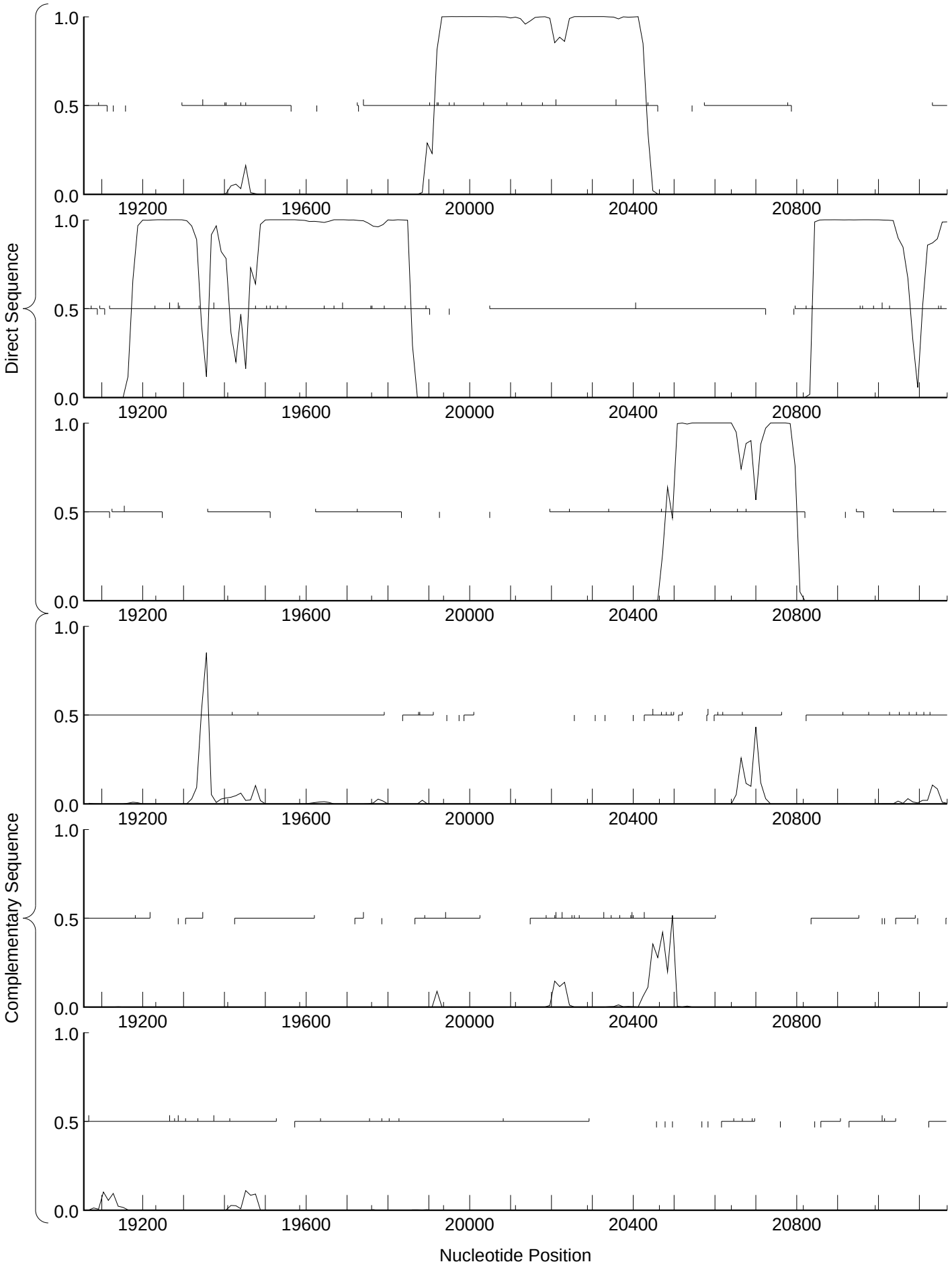
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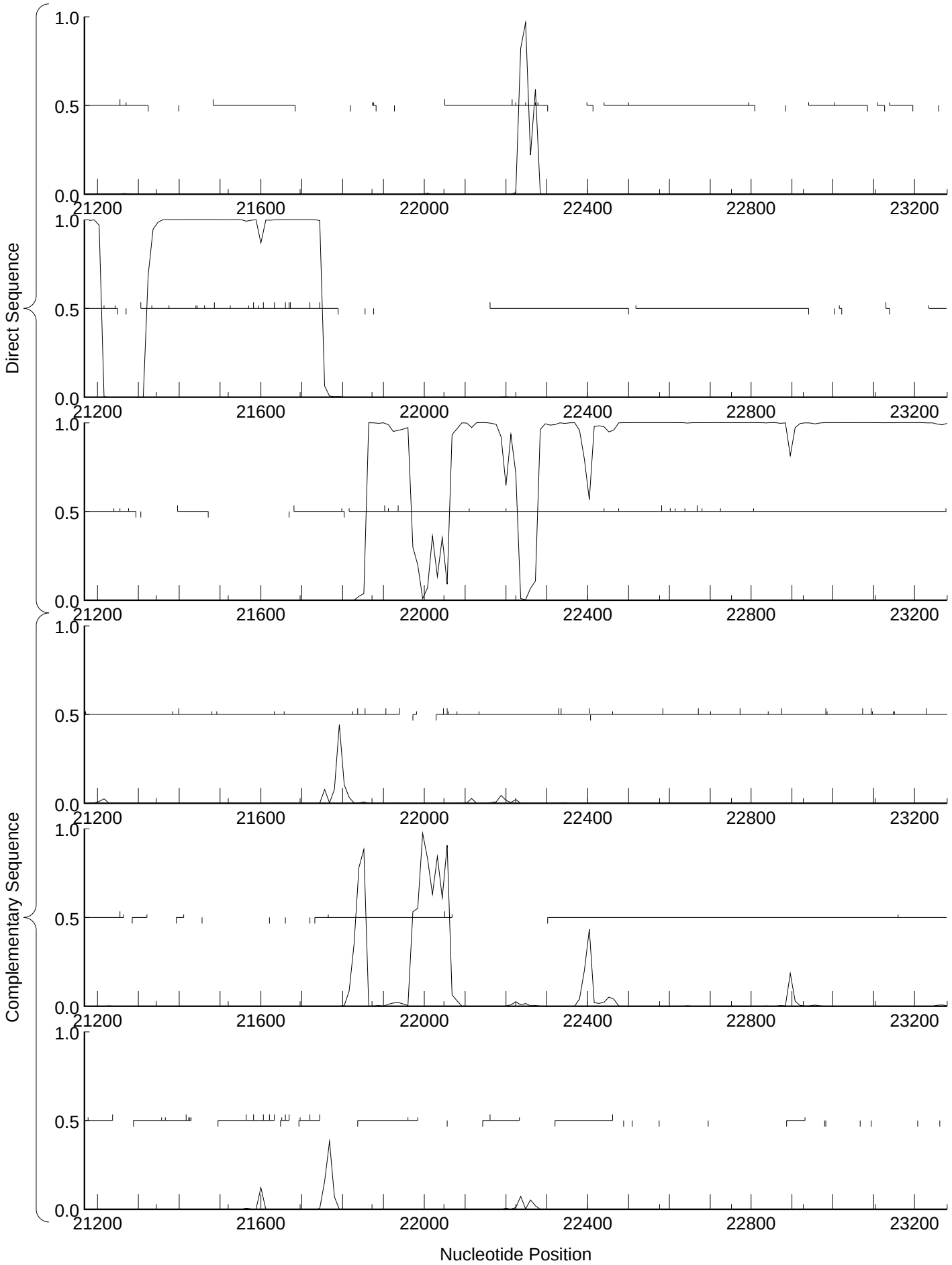
cobacterium phage ItsyBitsy1 complete sequence, 67570 bp, circularly permuted, Cluster B2, Order 4, Window 96, Step 12, 10/33



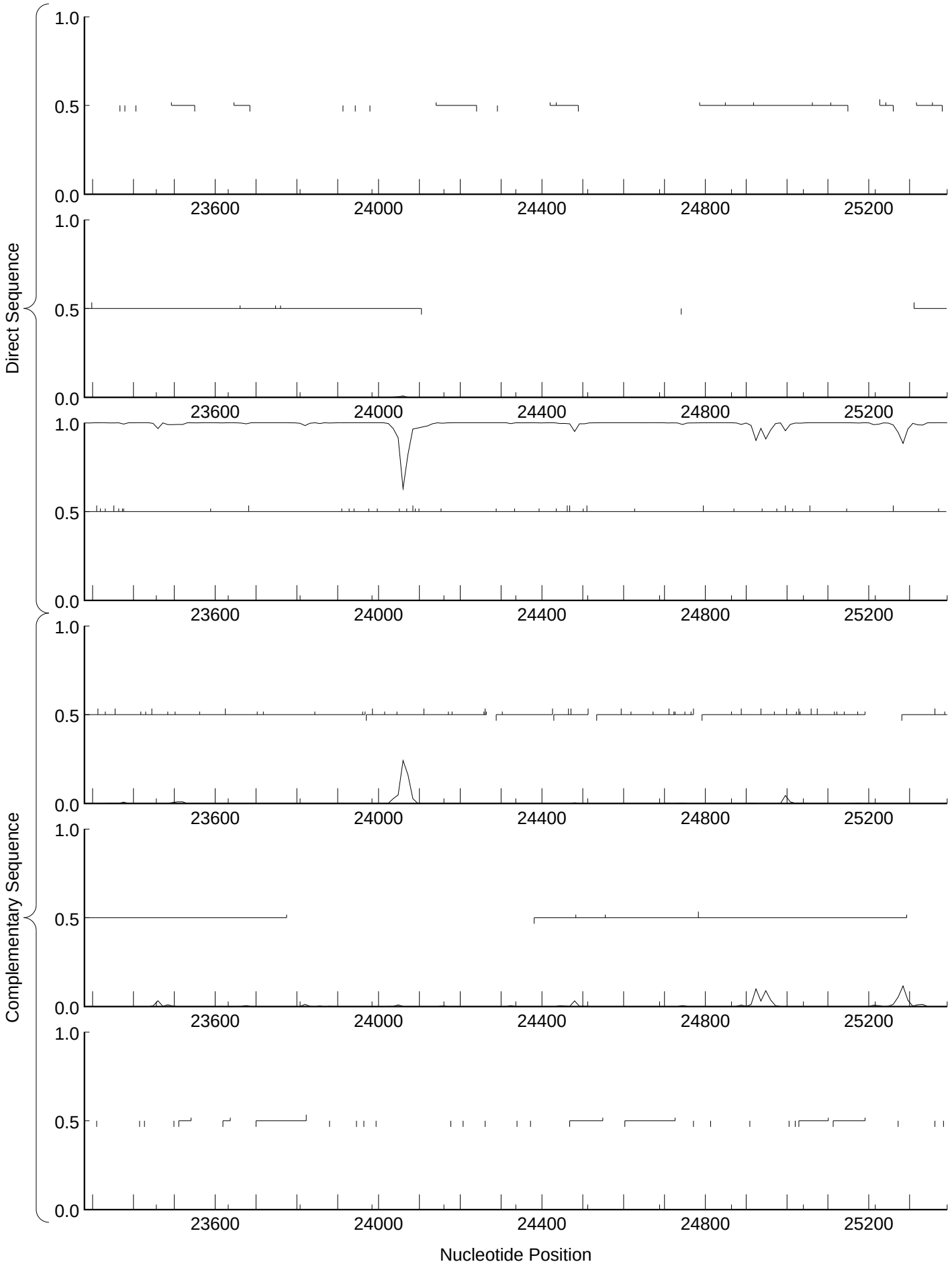
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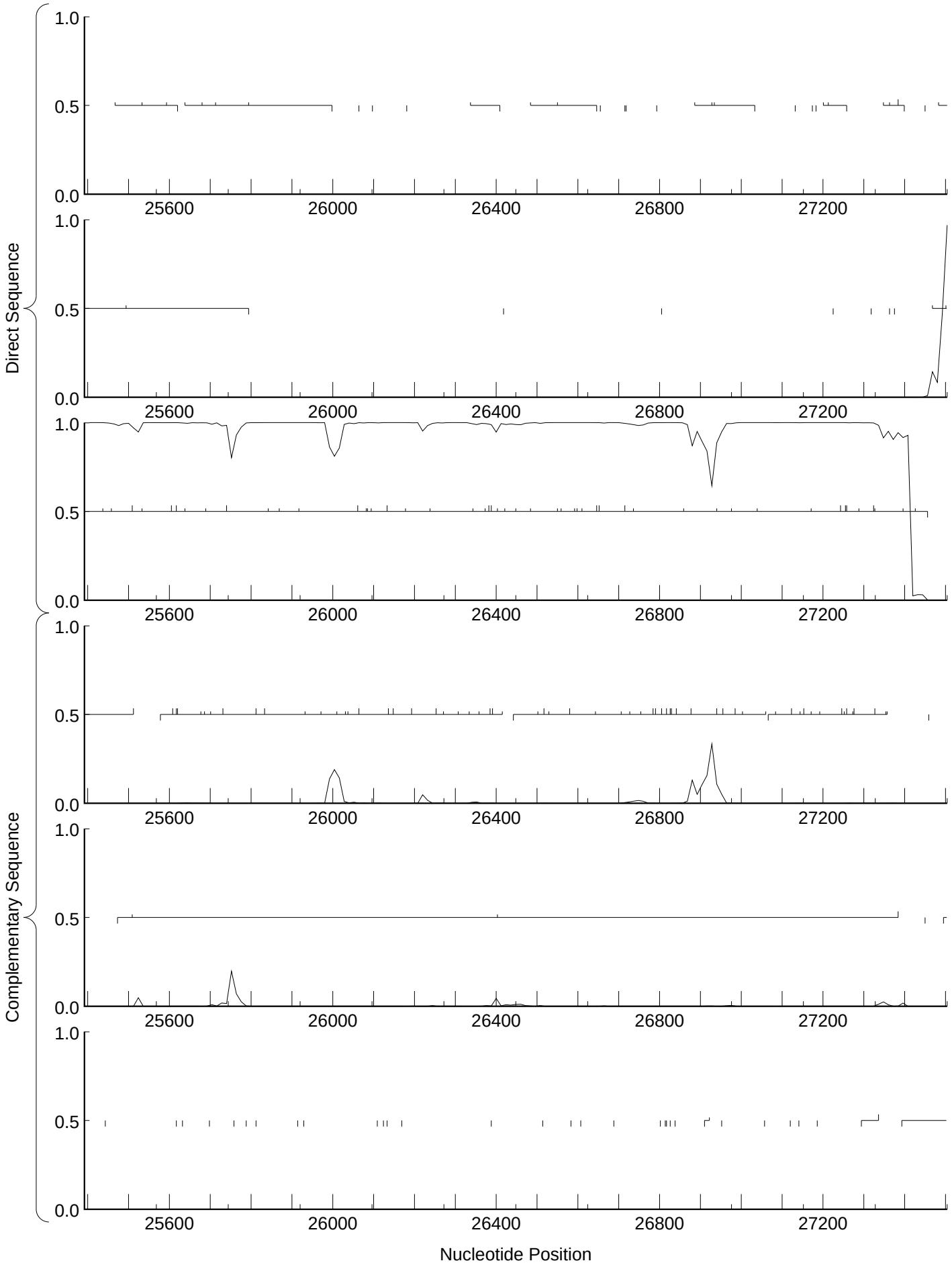
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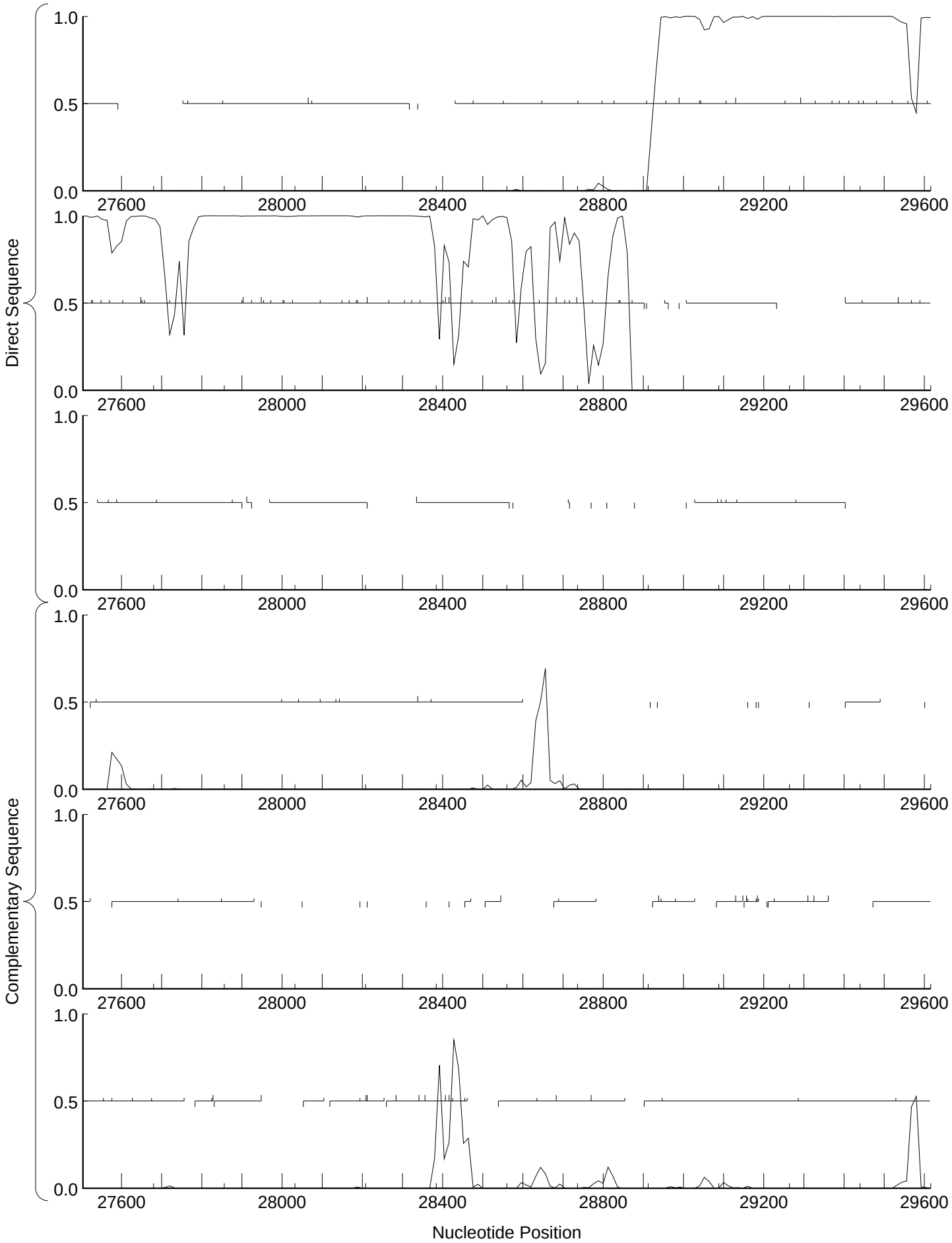
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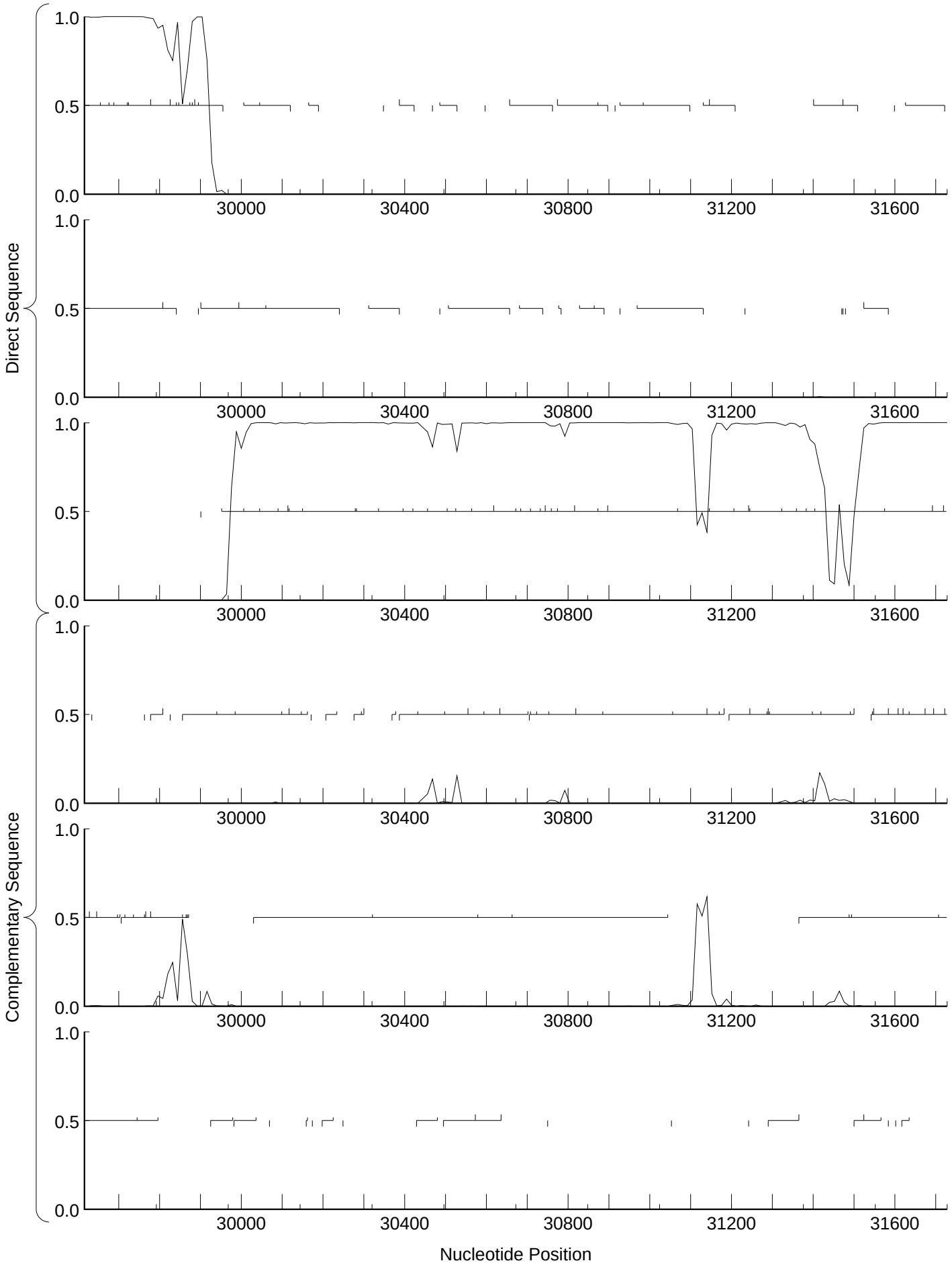
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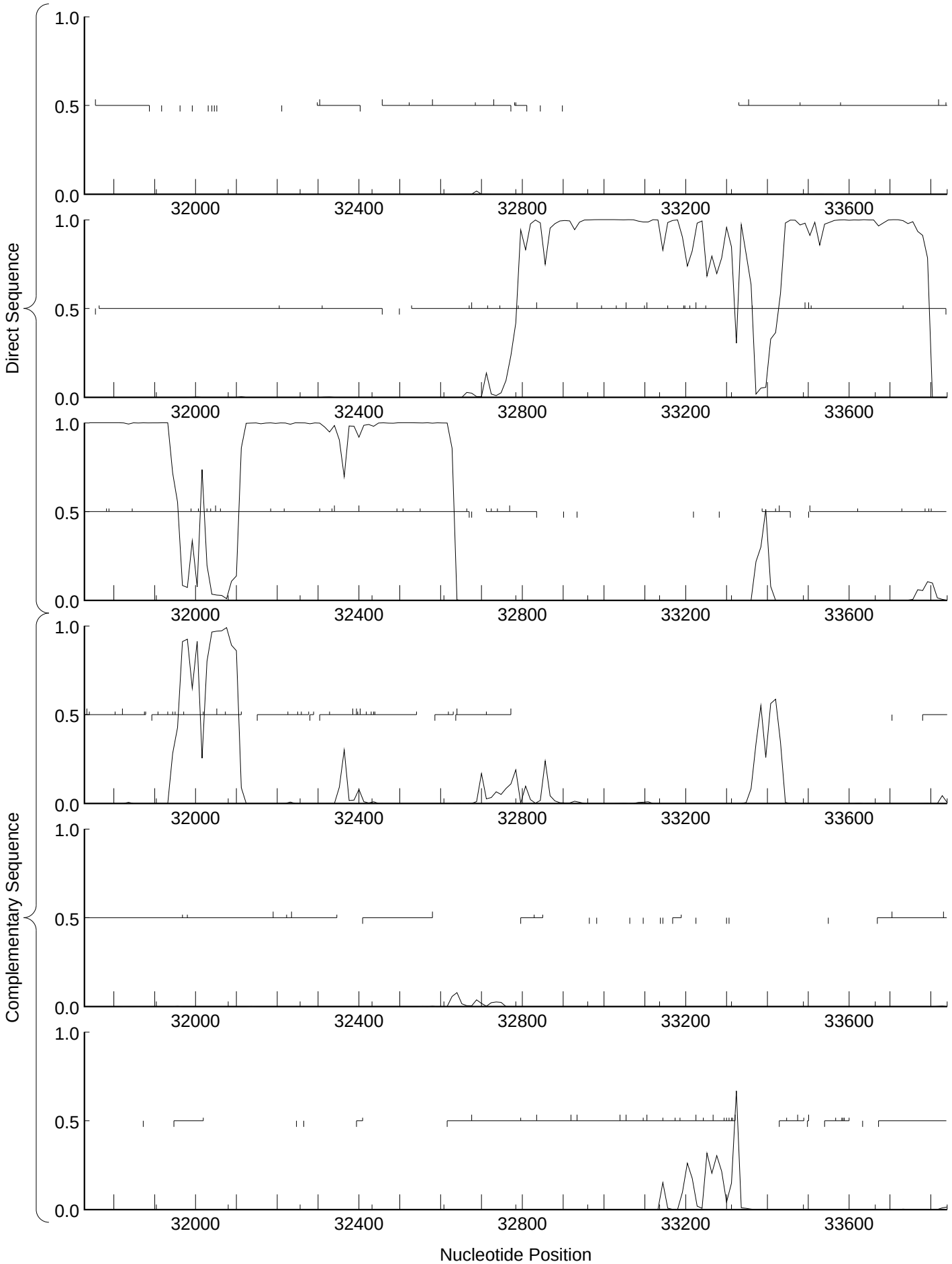
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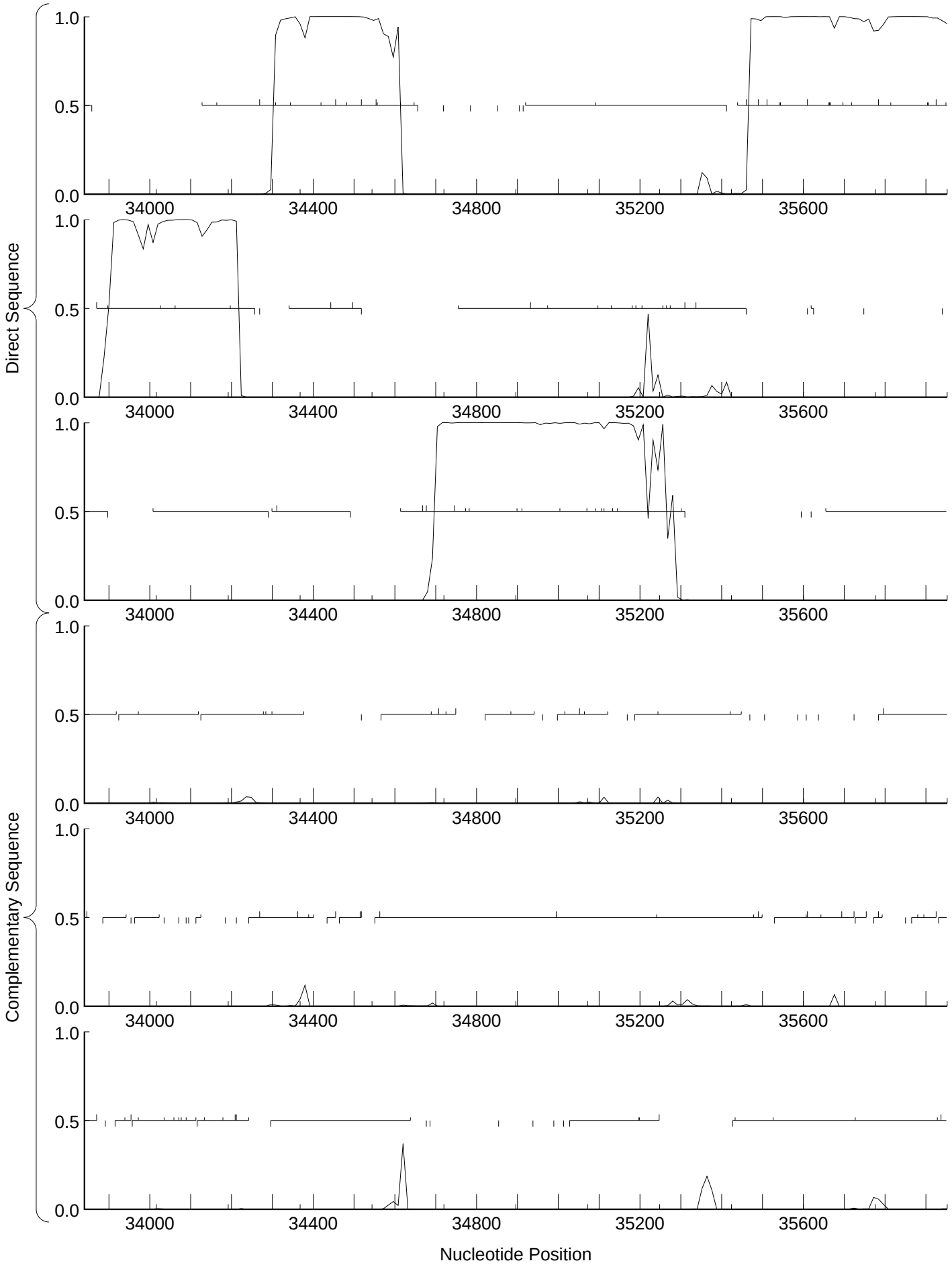
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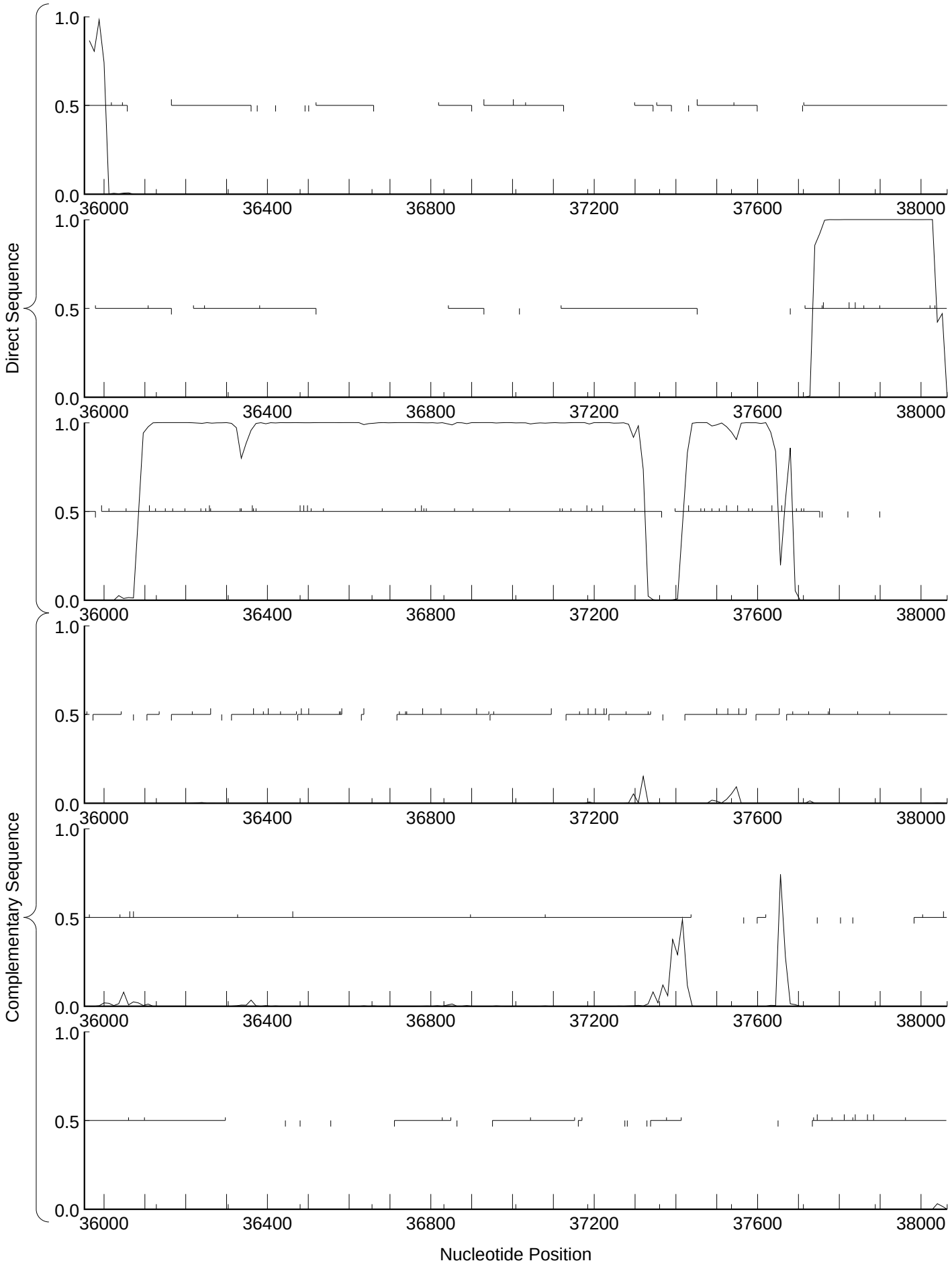
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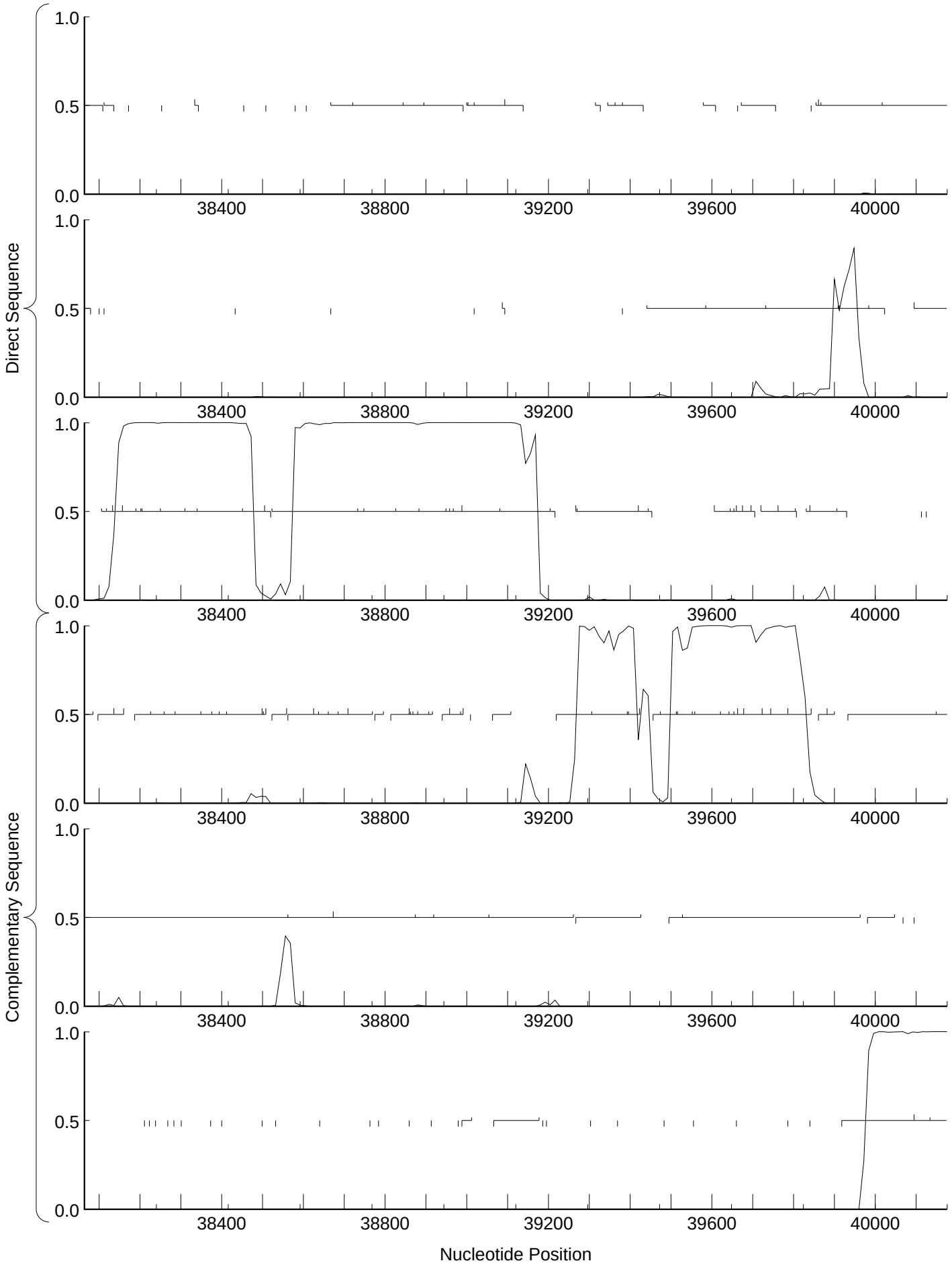
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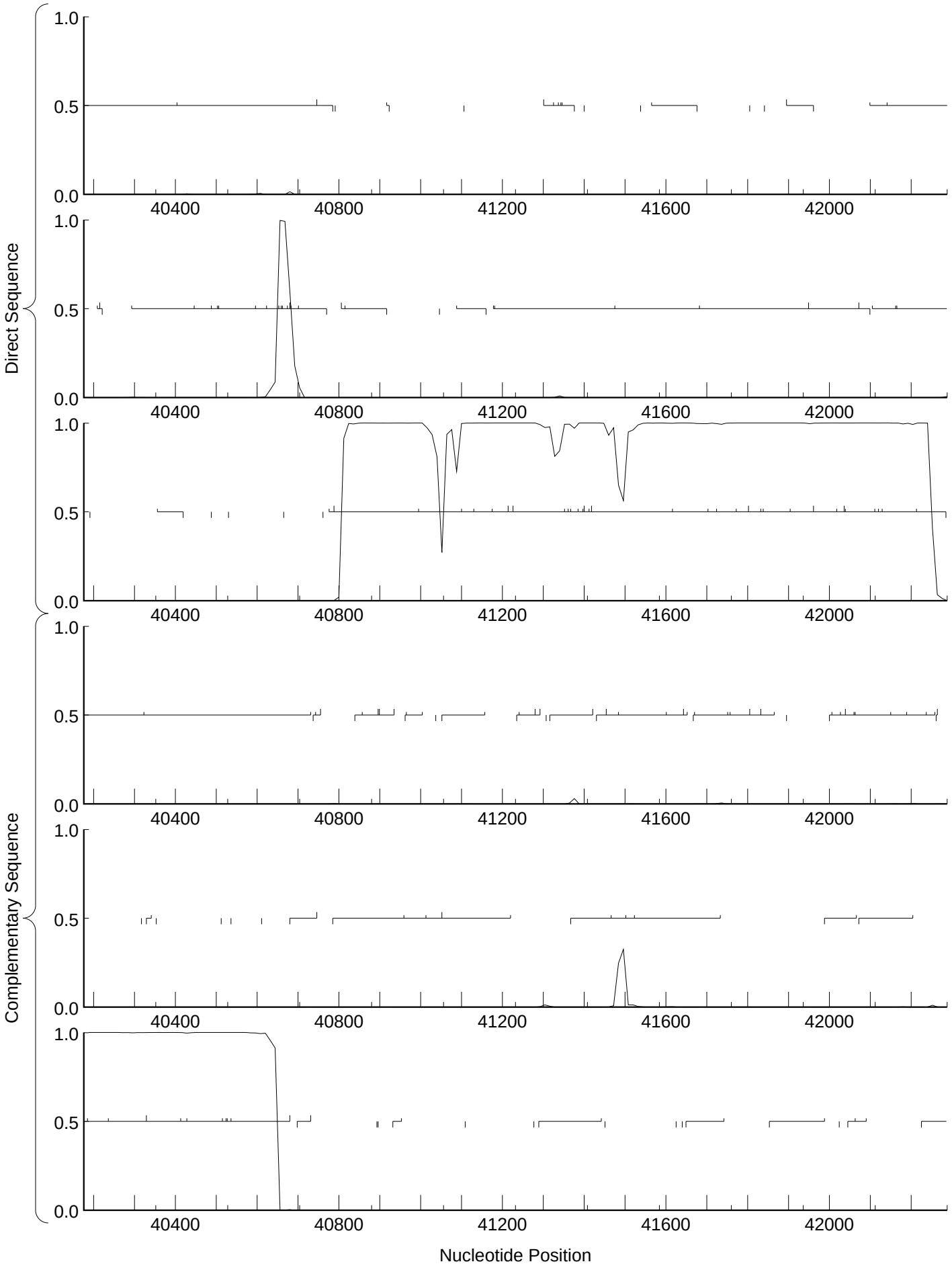
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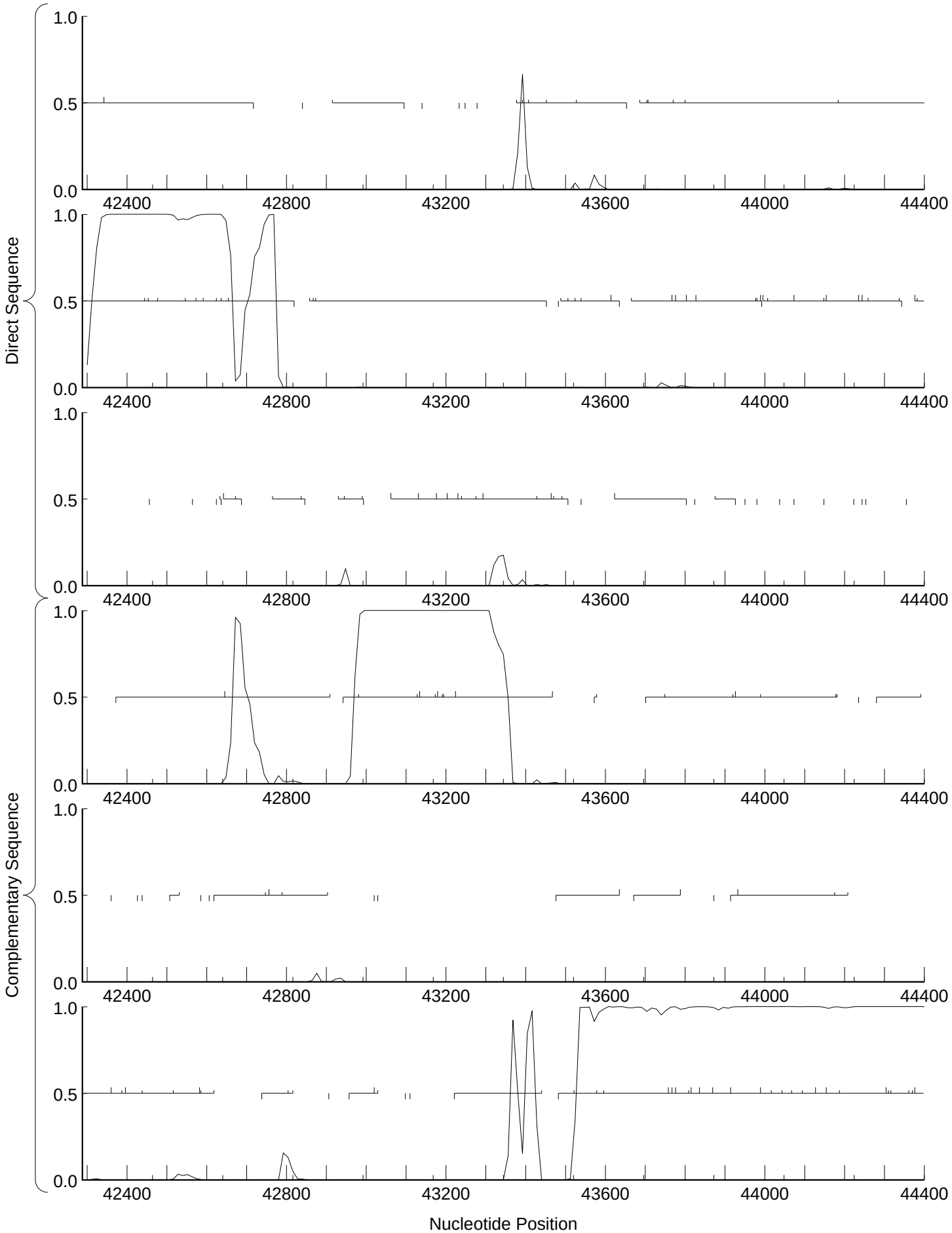
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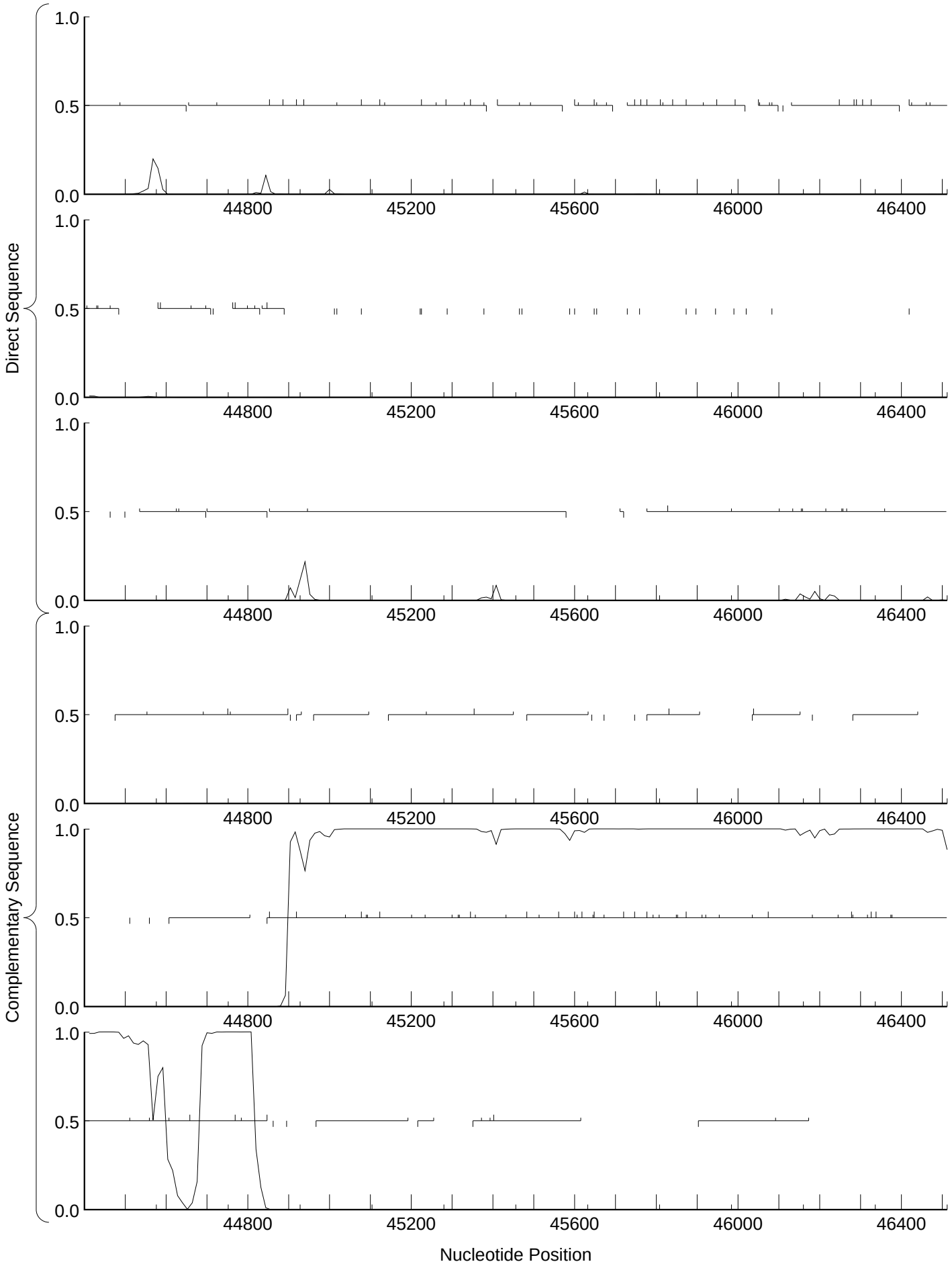
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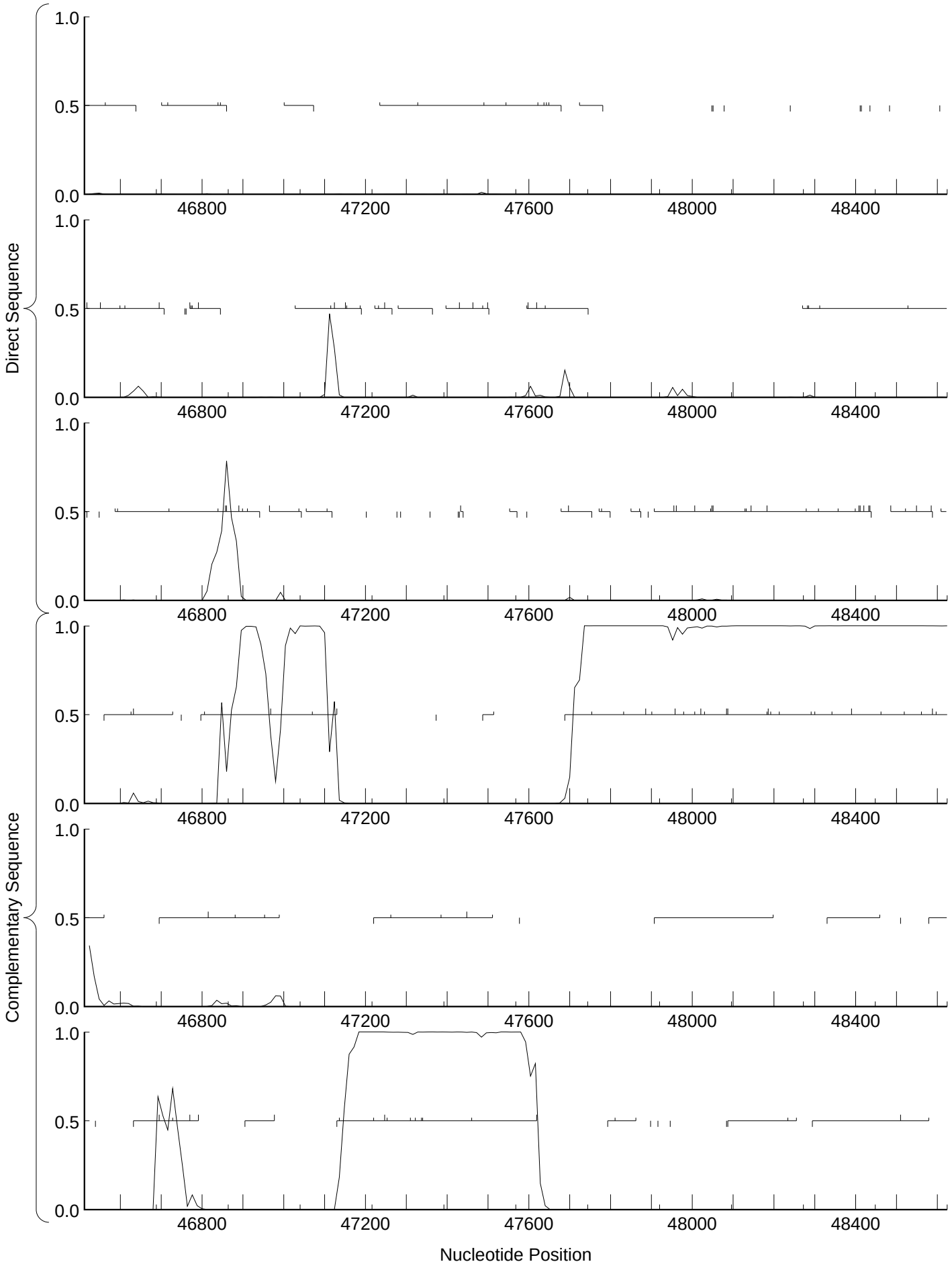
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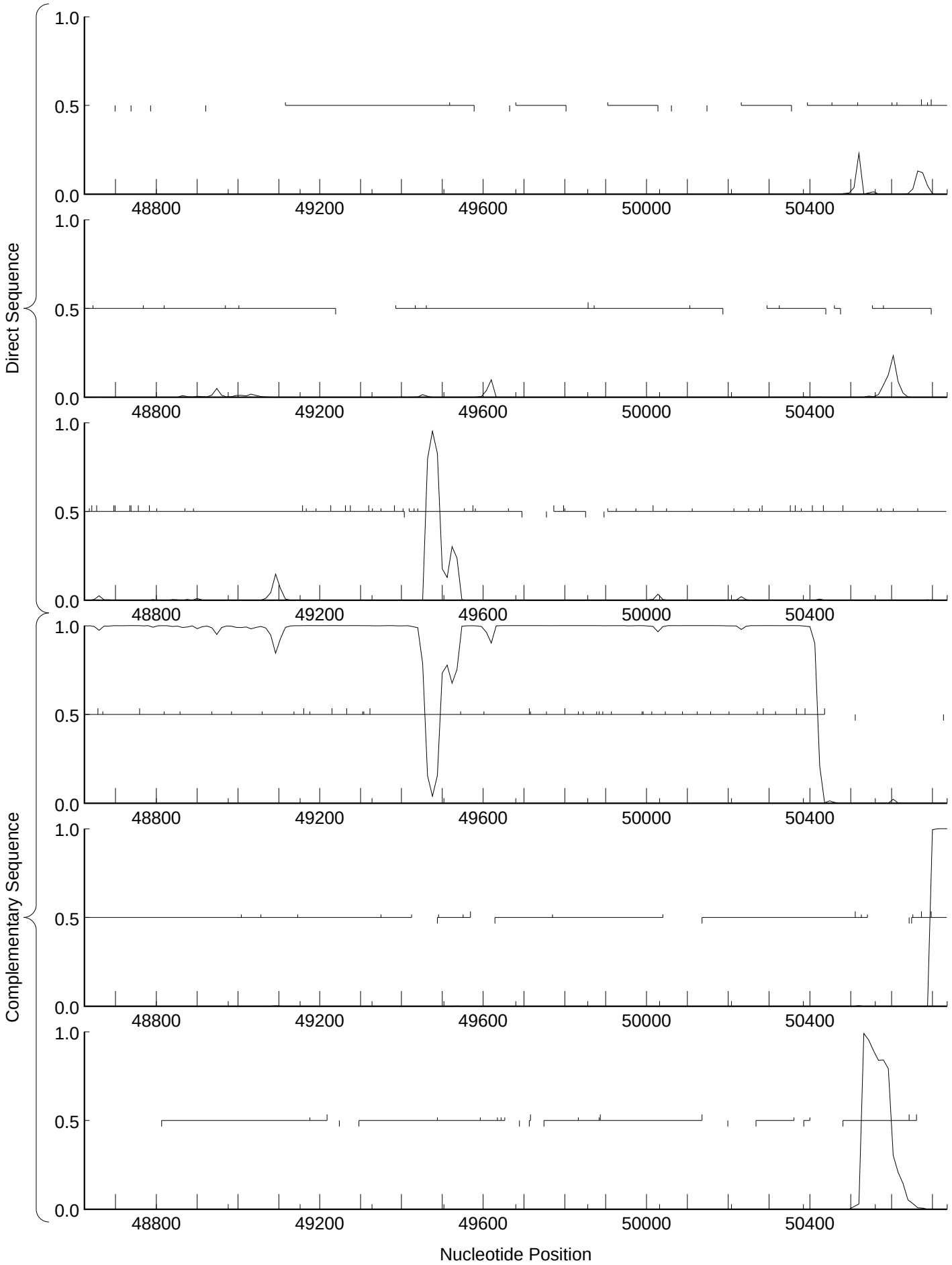
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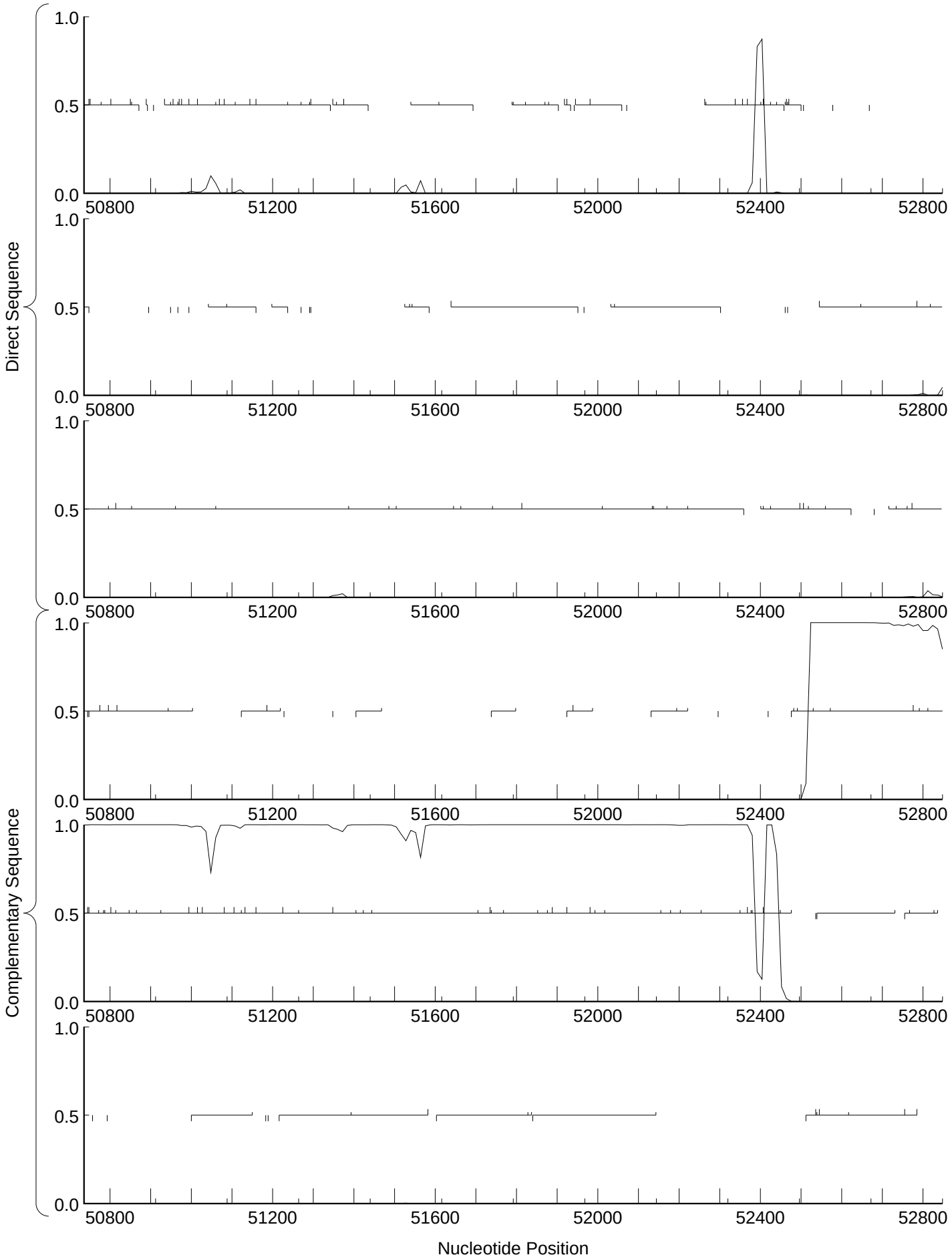
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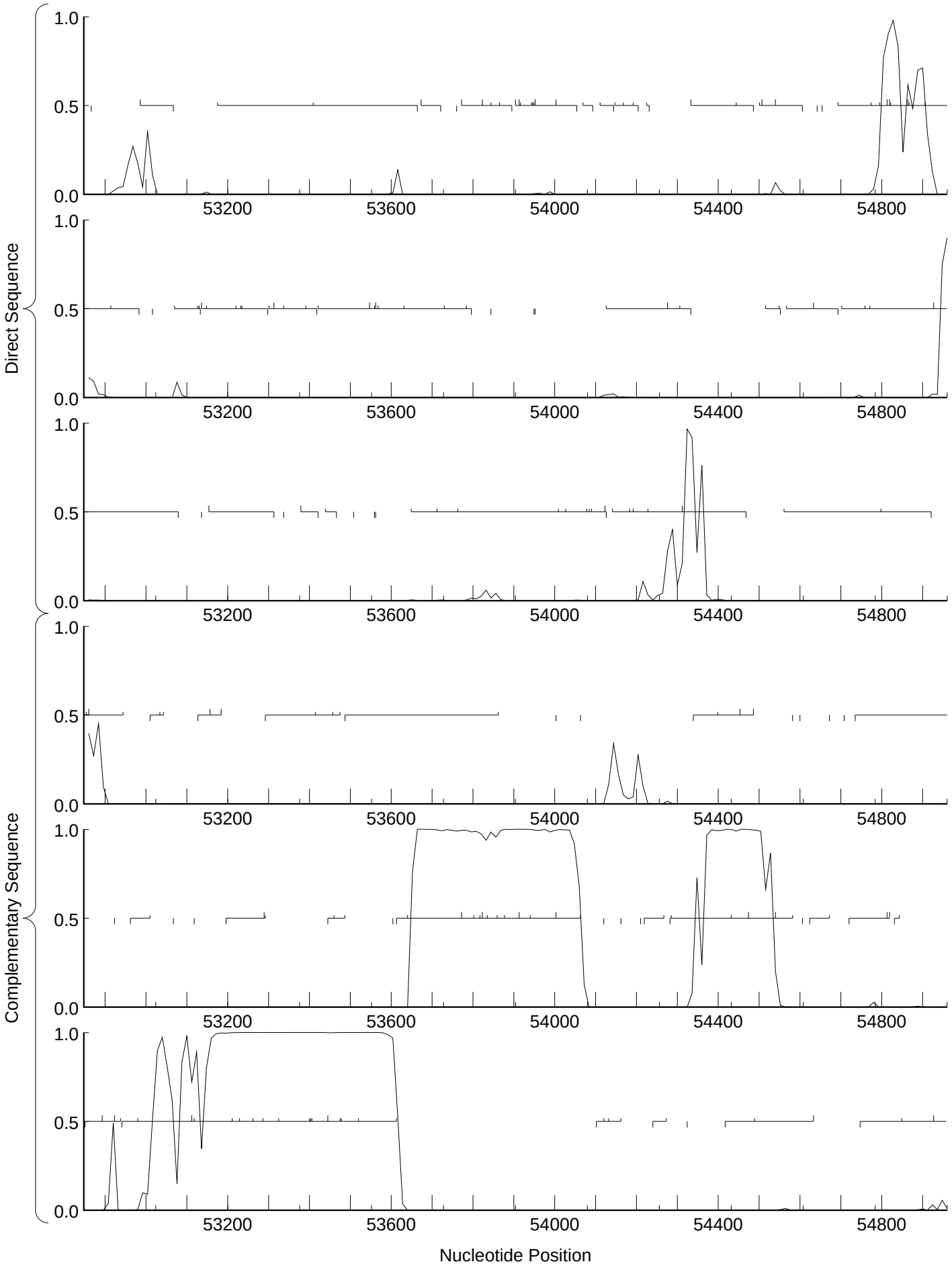
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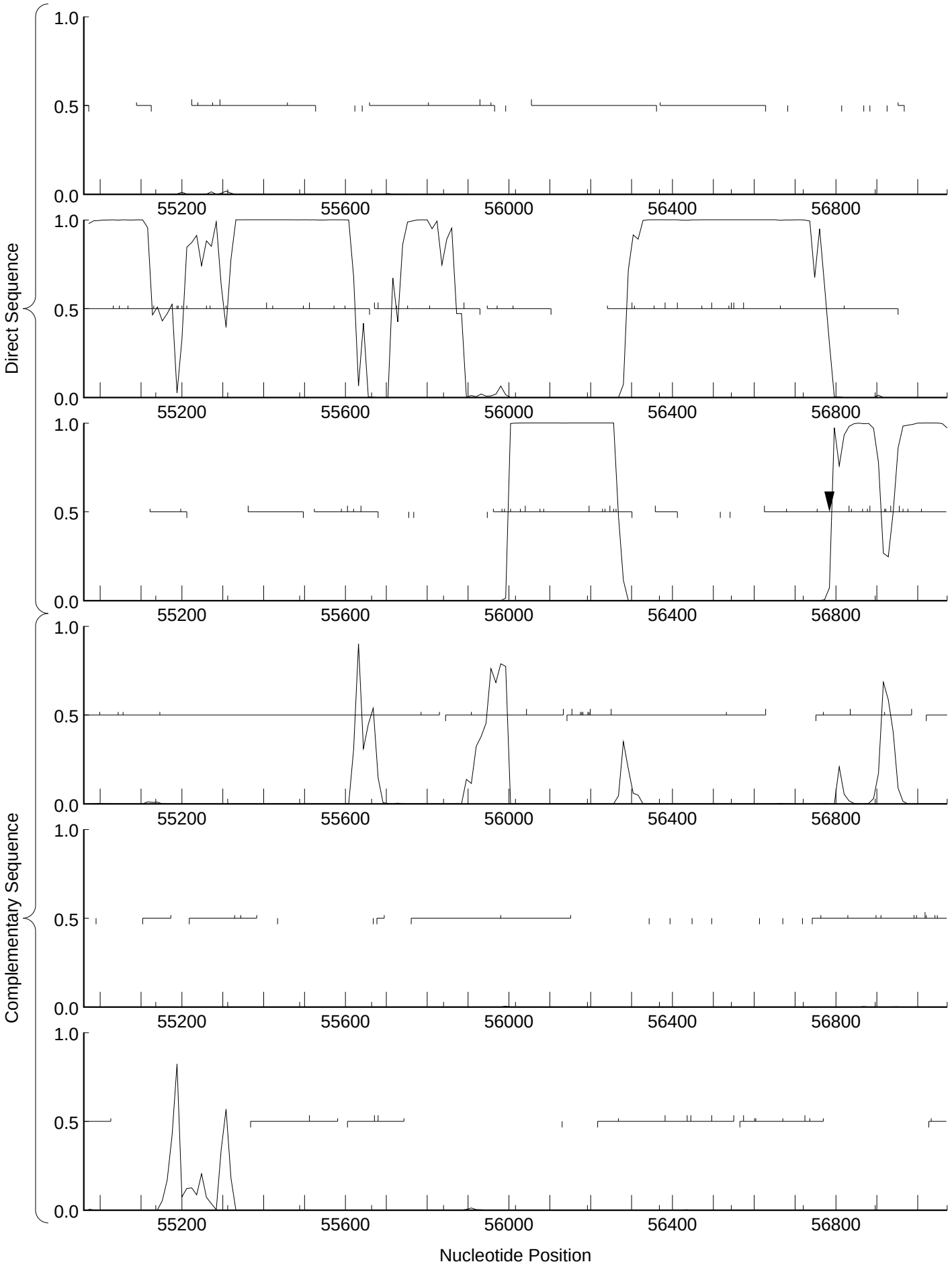
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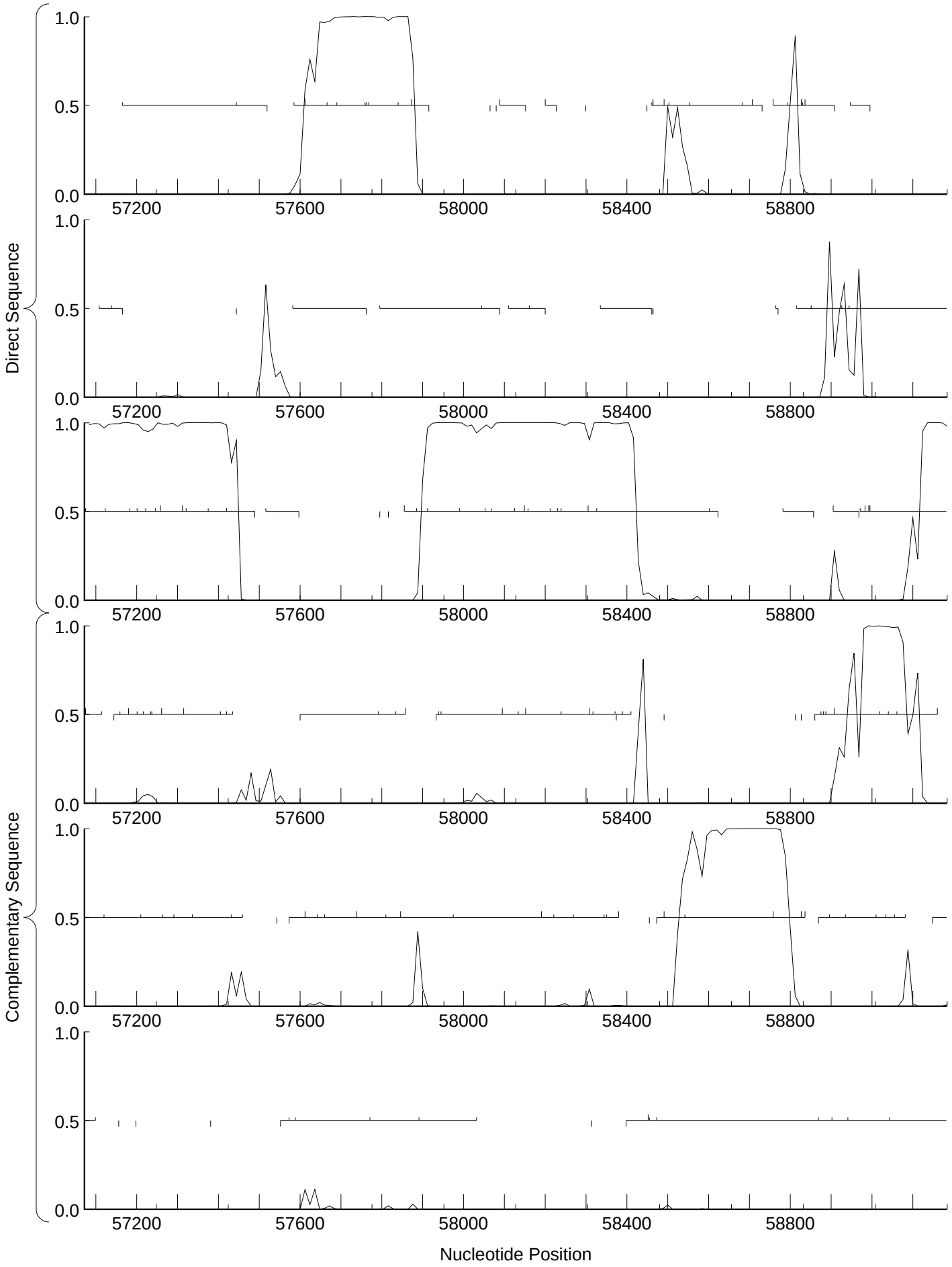
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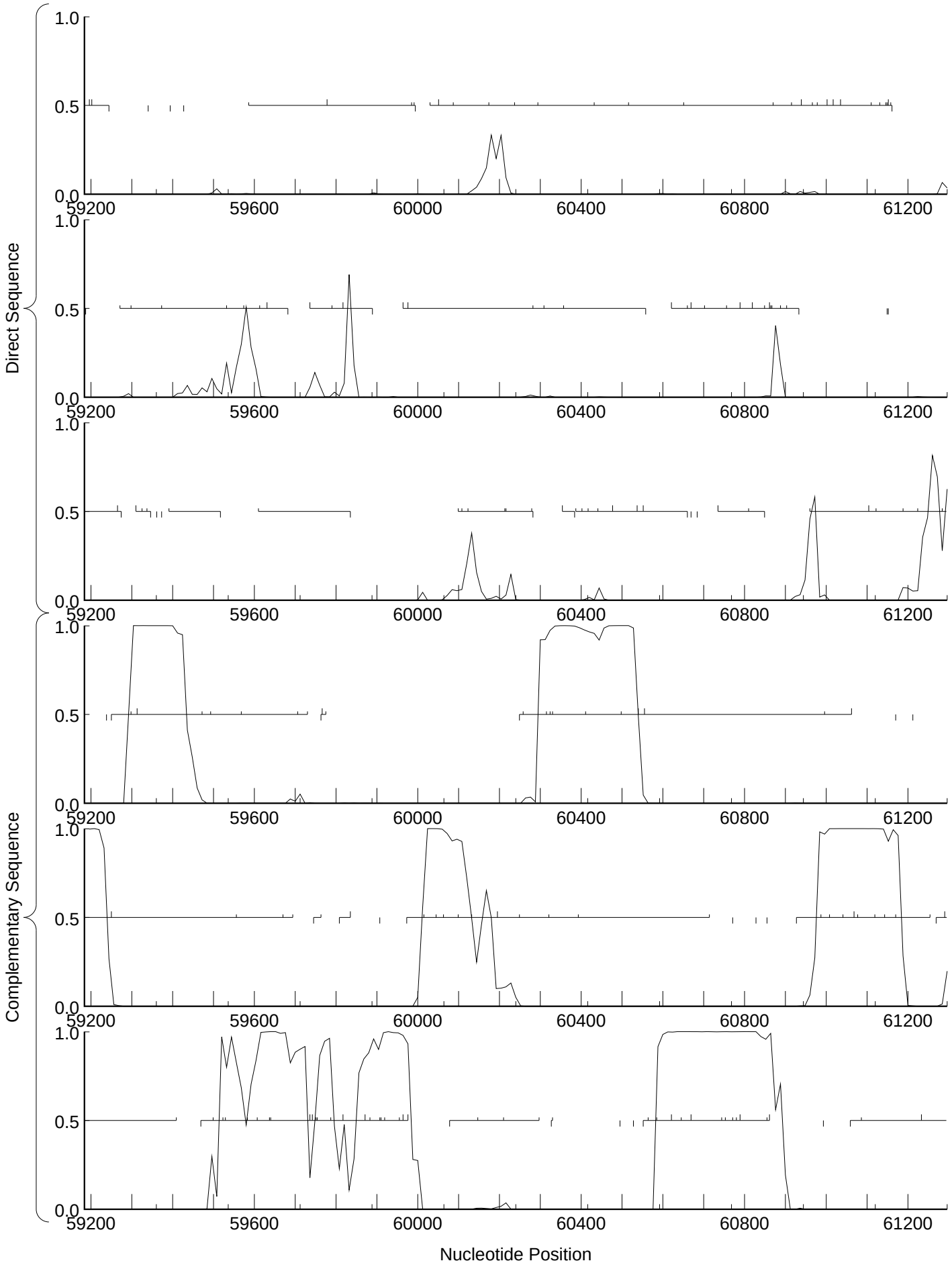
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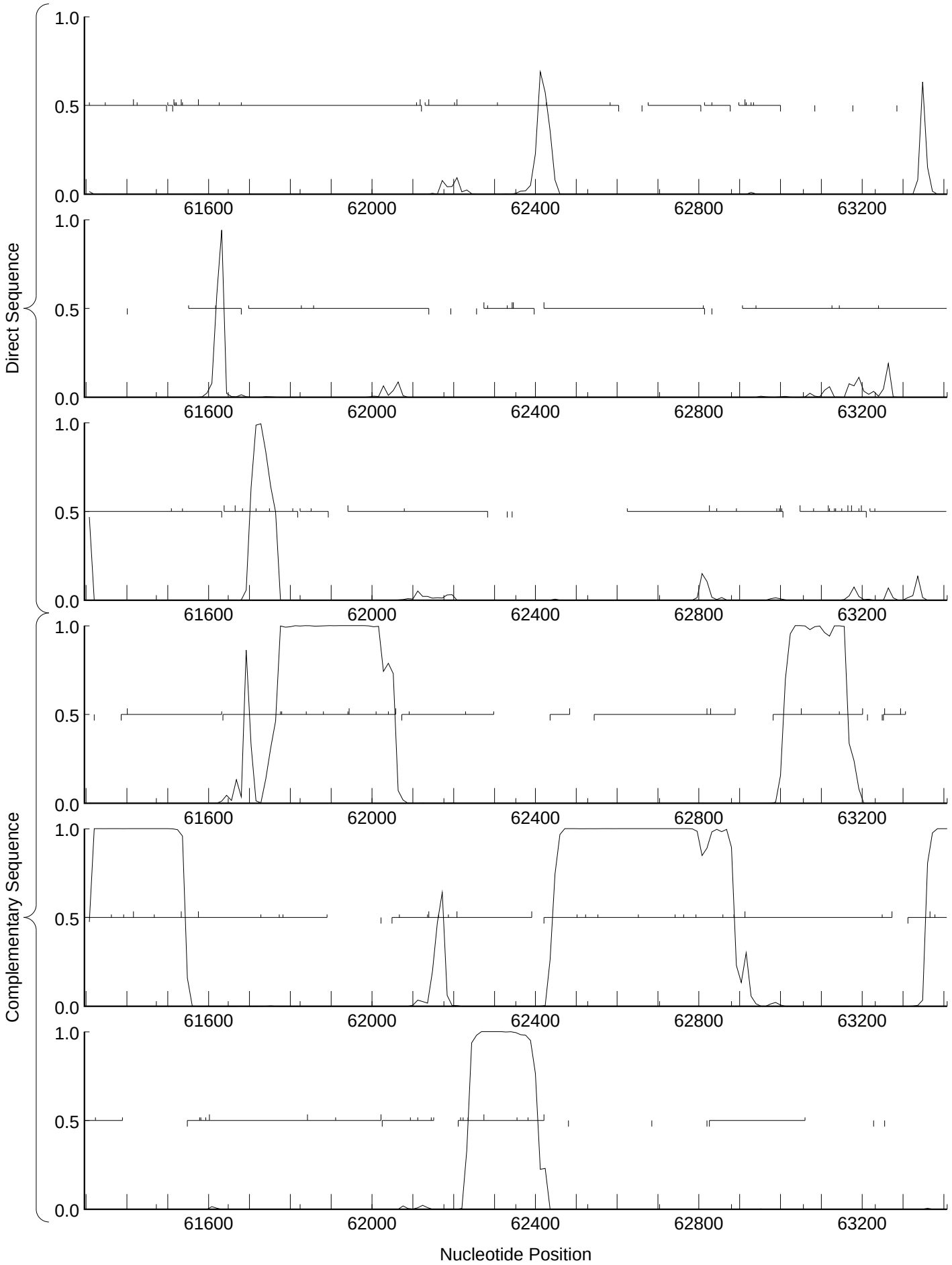
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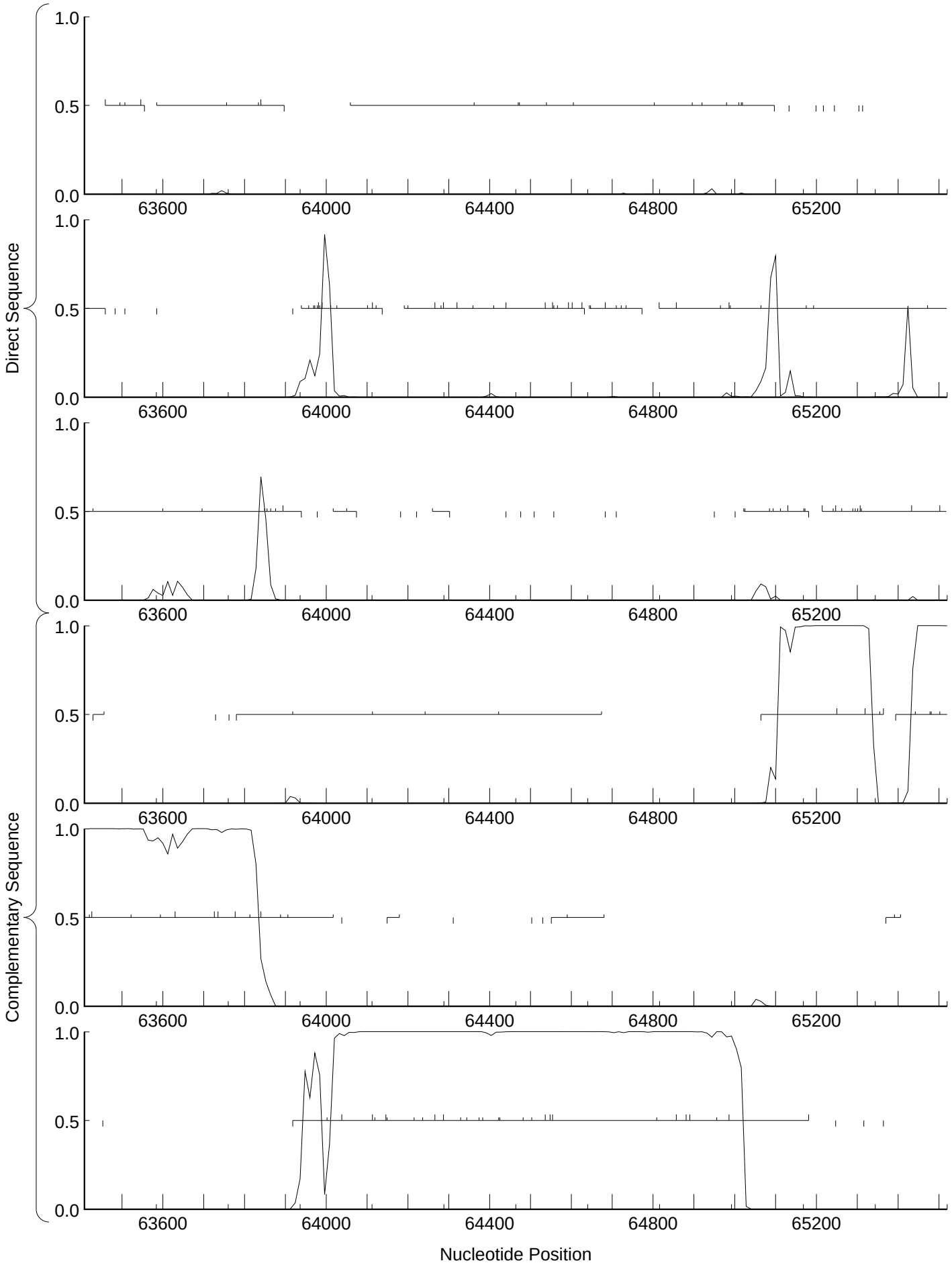
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cobacterium phage ItsyBitsy1 complete sequence, 67570 bp, circularly permuted, Cluster B2, Order 4, Window 96, Step 12, 31/33



cobacterium phage ItsyBitsy1 complete sequence, 67570 bp, circularly permuted, Cluster B2, Order 4, Window 96, Step 12, 32/33



cobacterium phage ItsyBitsy1 complete sequence, 67570 bp, circularly permuted, Cluster B2, Order 4, Window 96, Step 12, 33/33

