# GeneMark

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

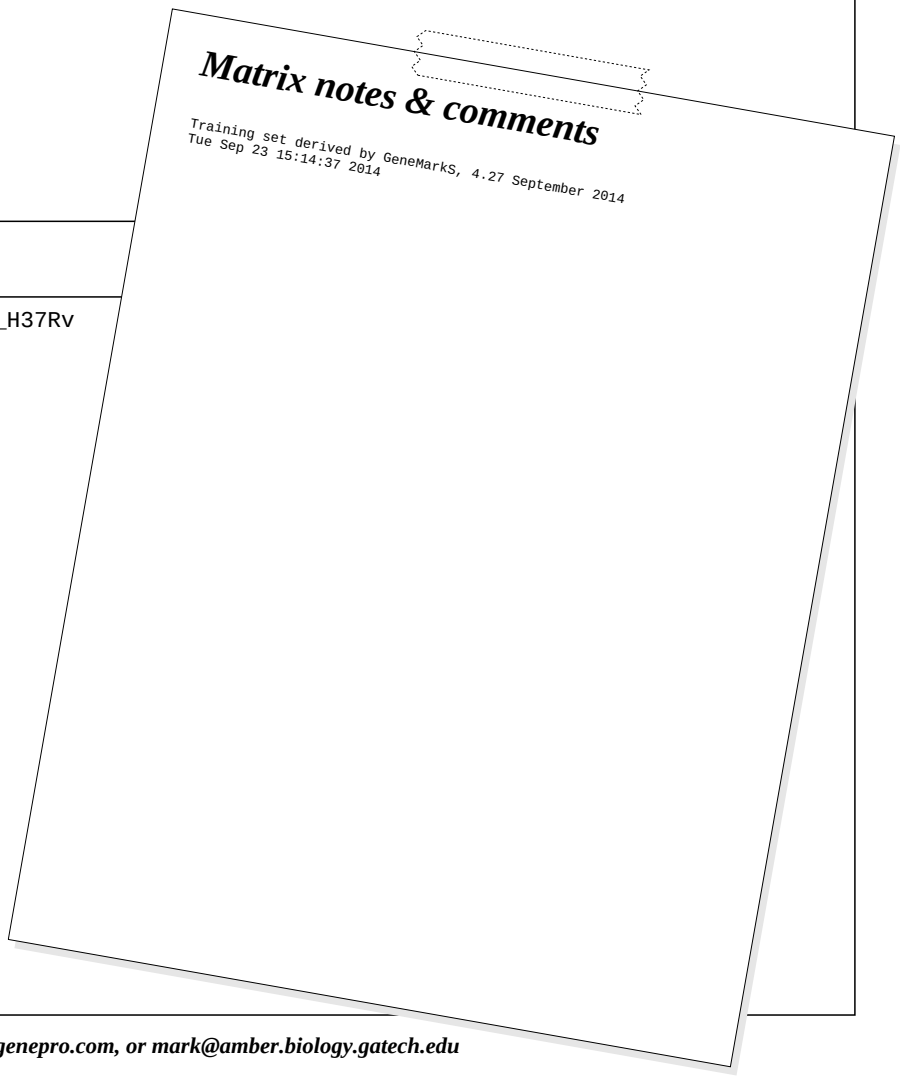
## PROGRAM INFORMATION

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Sequence         :  Mycobacterium phage SDcharge11 complete sequence, 67702 bp, circularly permuted, Cluster B1
Analysis Date    :  3/28/24 at 14:57:49
Pages            :  34
Sequence Length  :  67702 bp
GC Content       :   66.52%

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

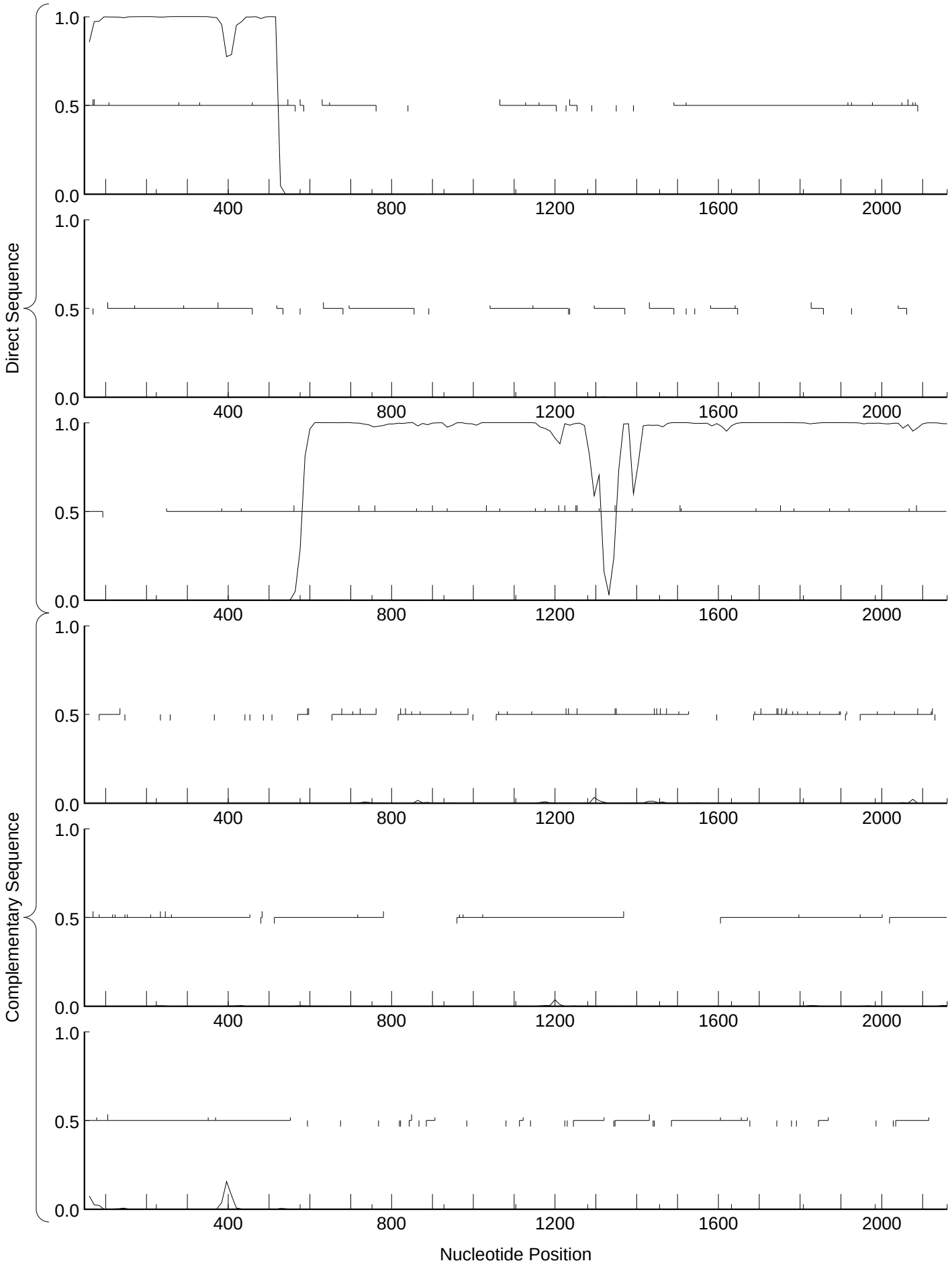
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Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

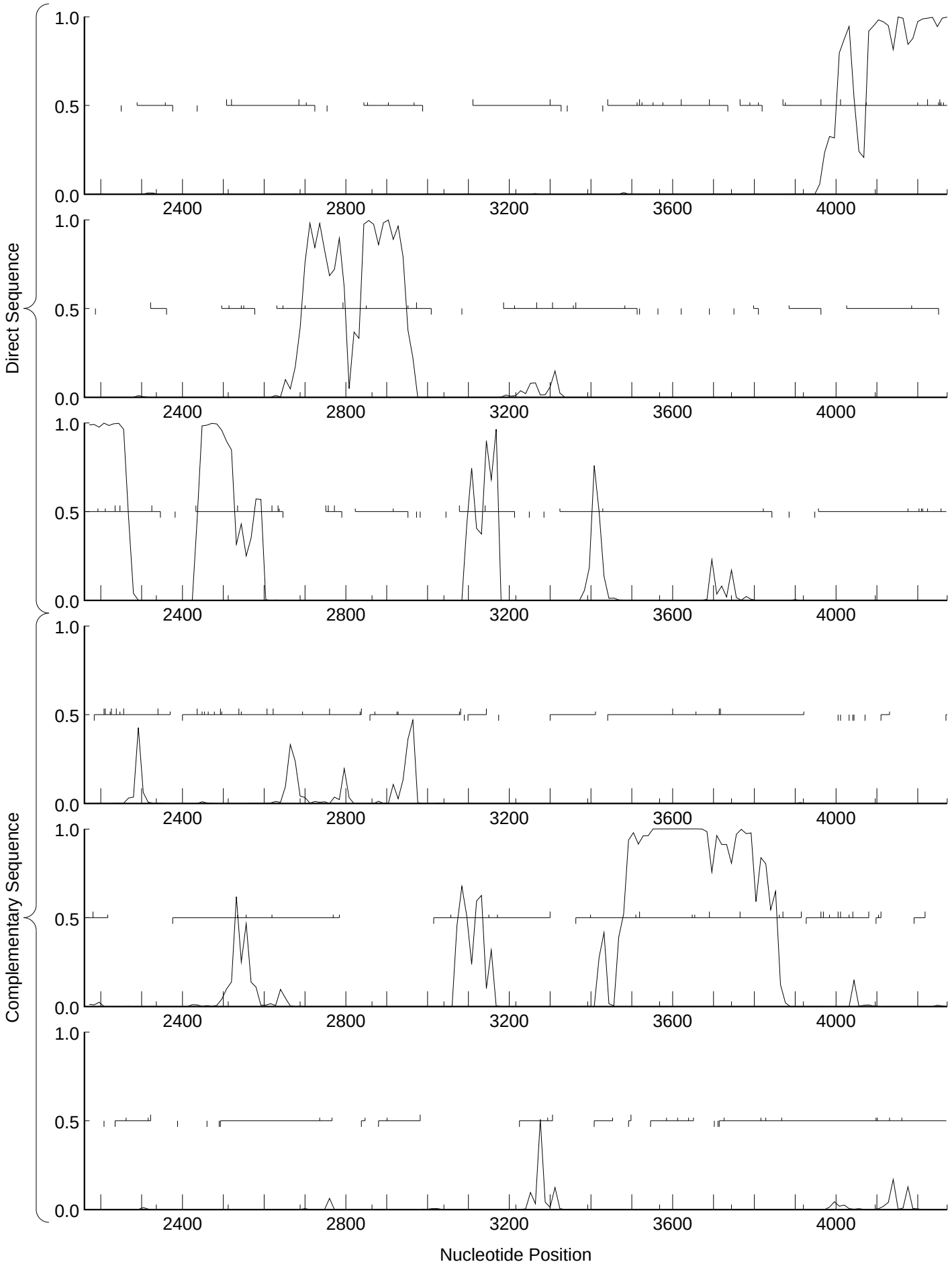
Order      :  4
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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

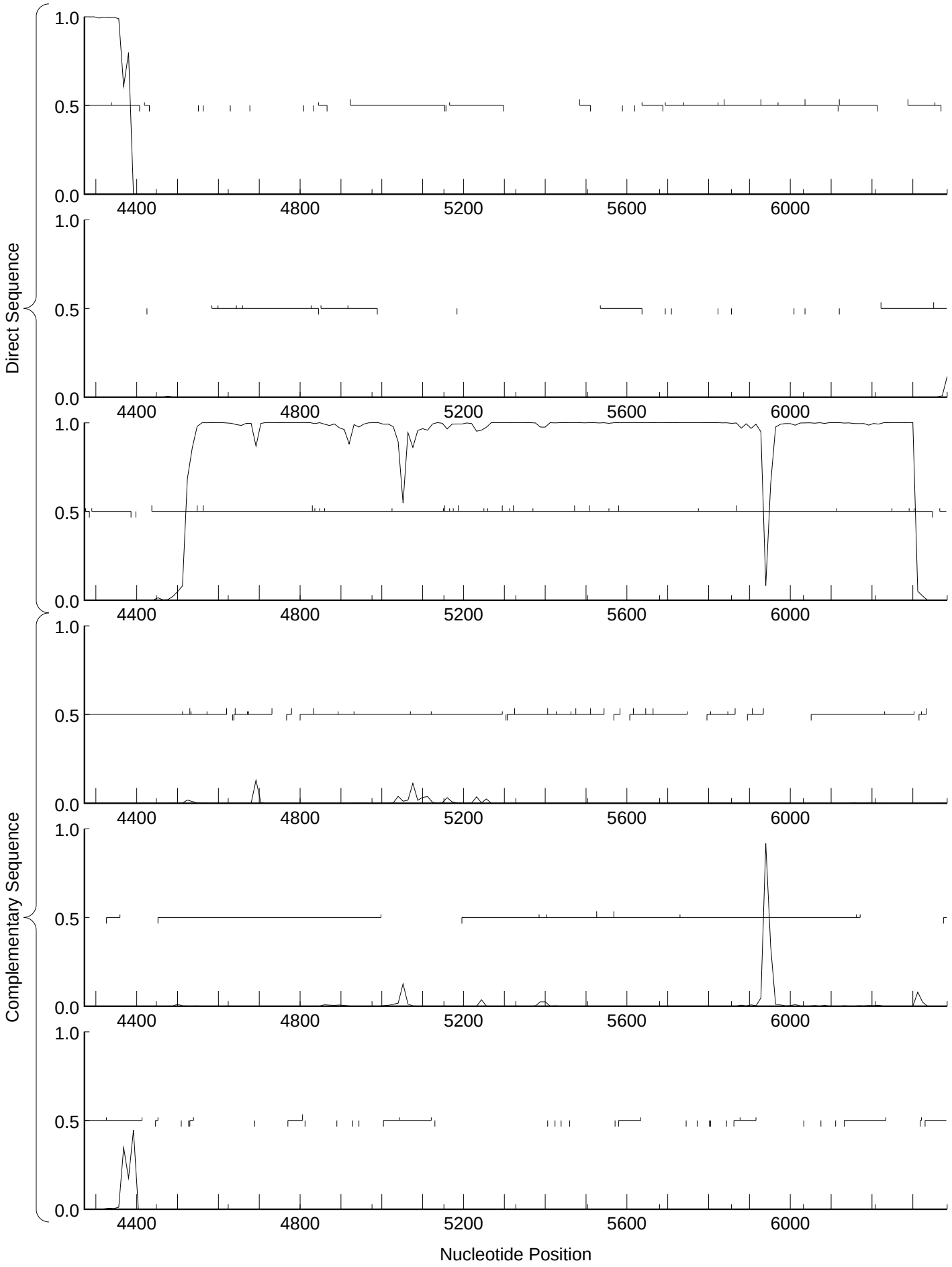

*For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu*

bacterium phage SDcharge11 complete sequence, 67702 bp, circularly permuted, Cluster B1 , Order 4, Window 96, Step 12, 2/34

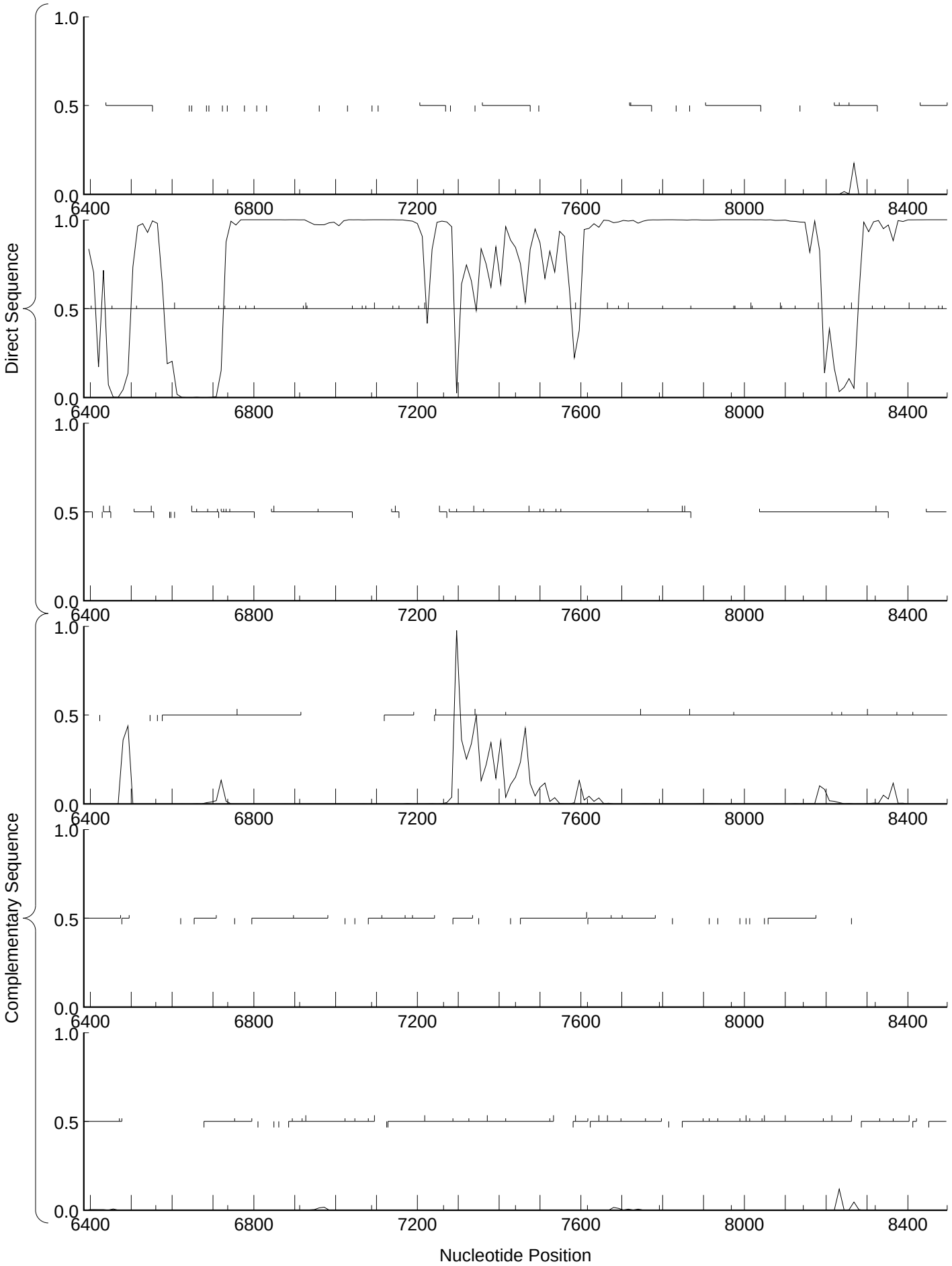


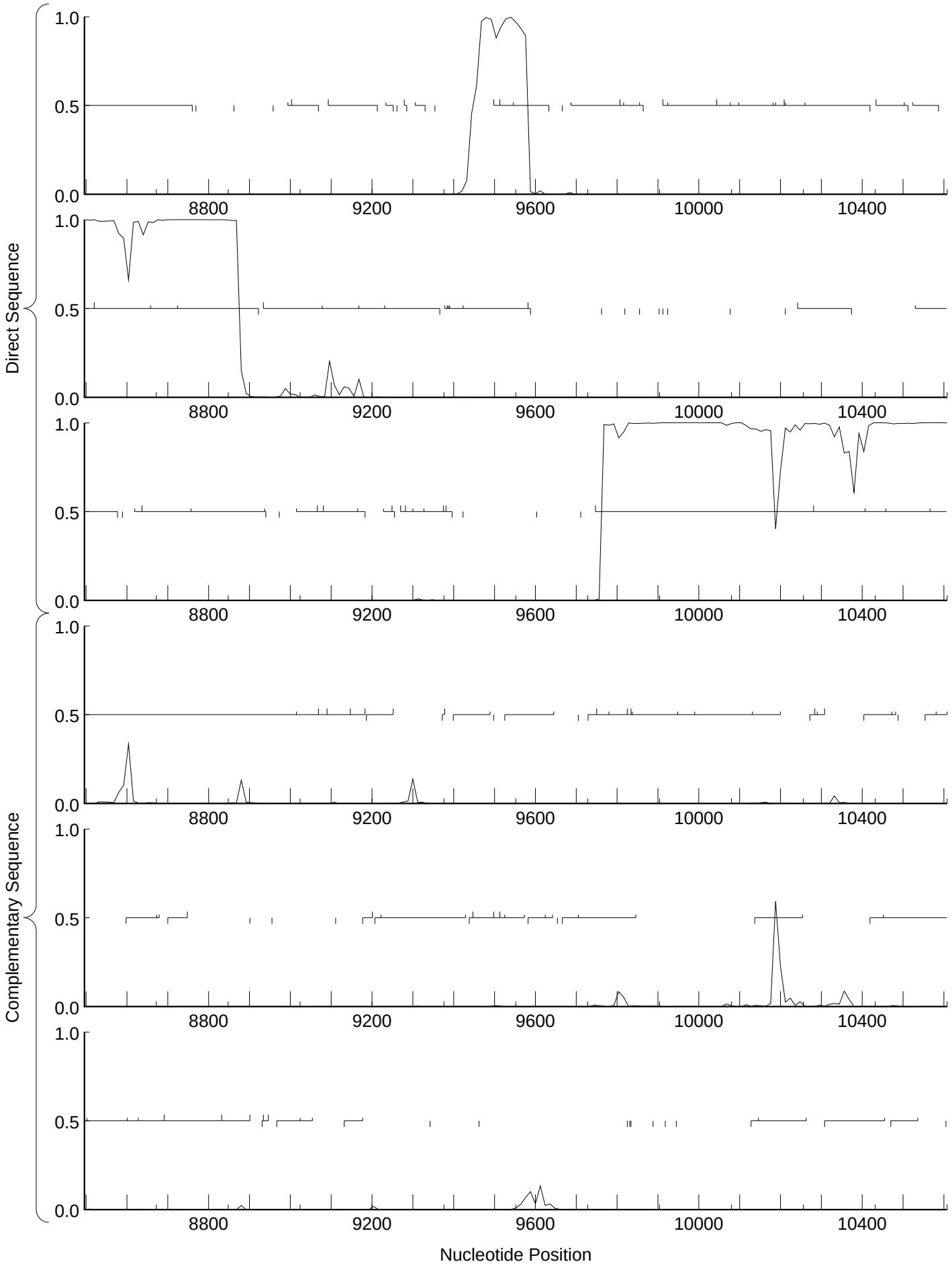

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
2400
2800
3200
3600
4000
Nucleotide Position

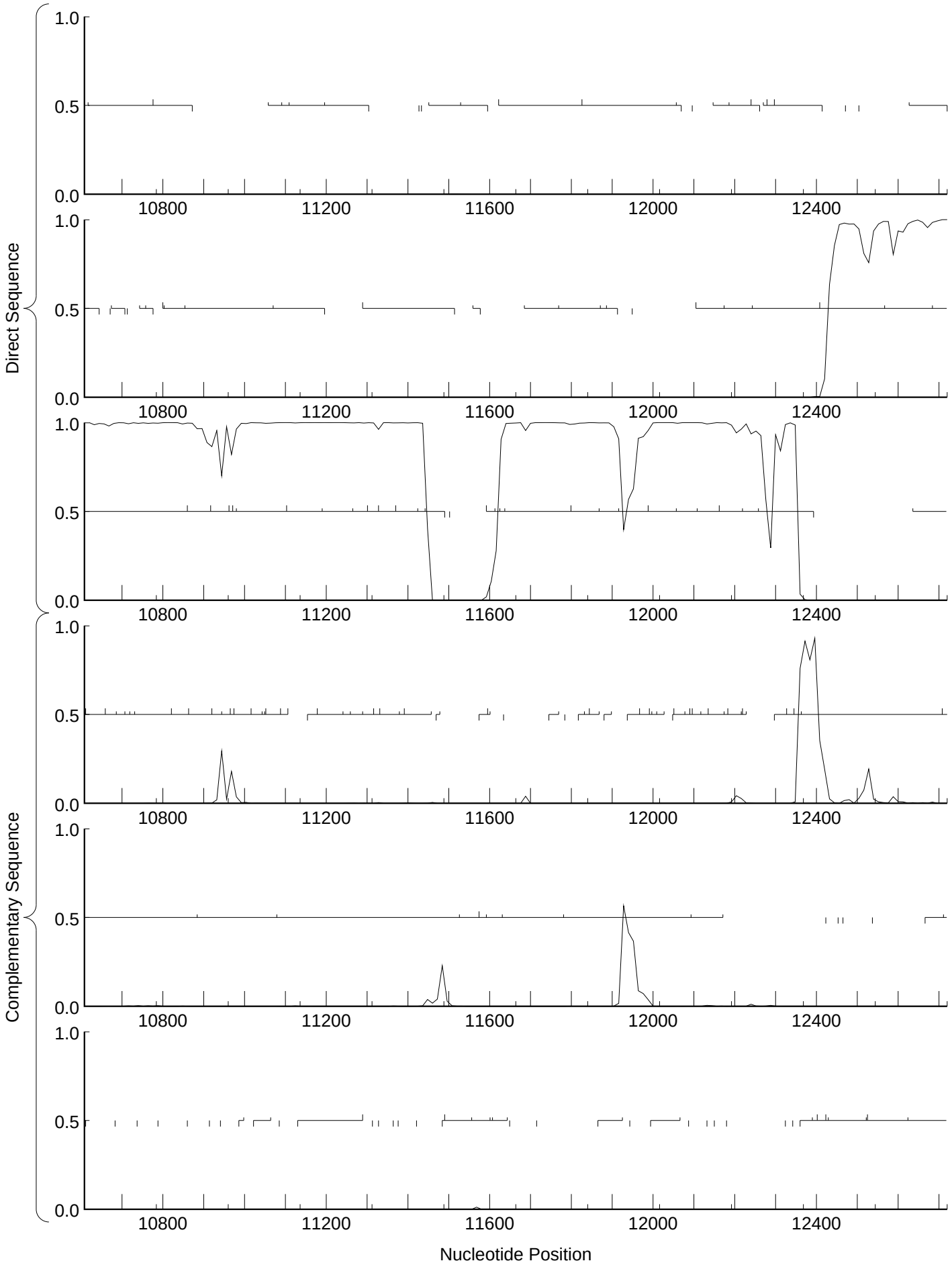
bacterium phage SDcharge11 complete sequence, 67702 bp, circularly permuted, Cluster B1 , Order 4, Window 96, Step 12, 4/34



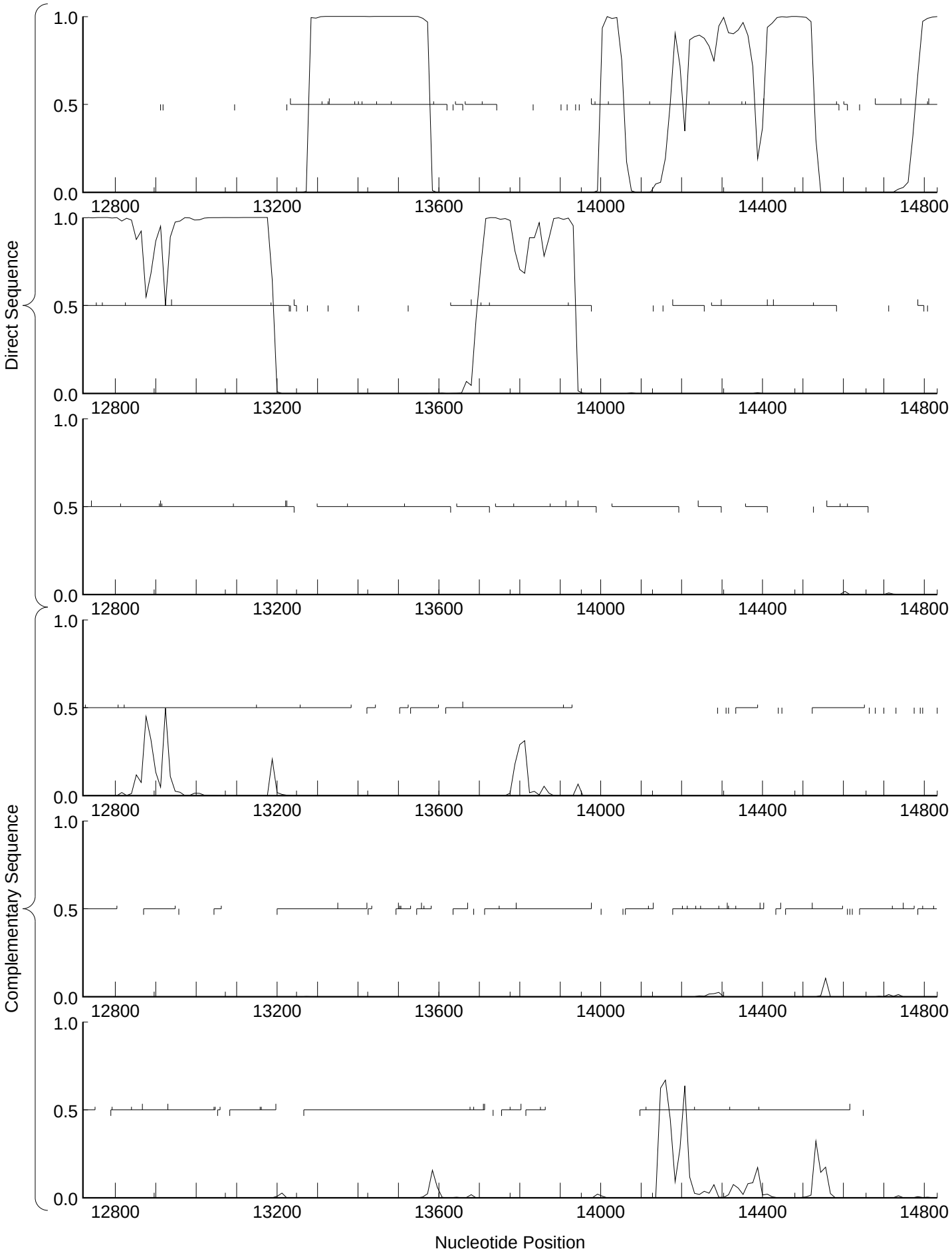
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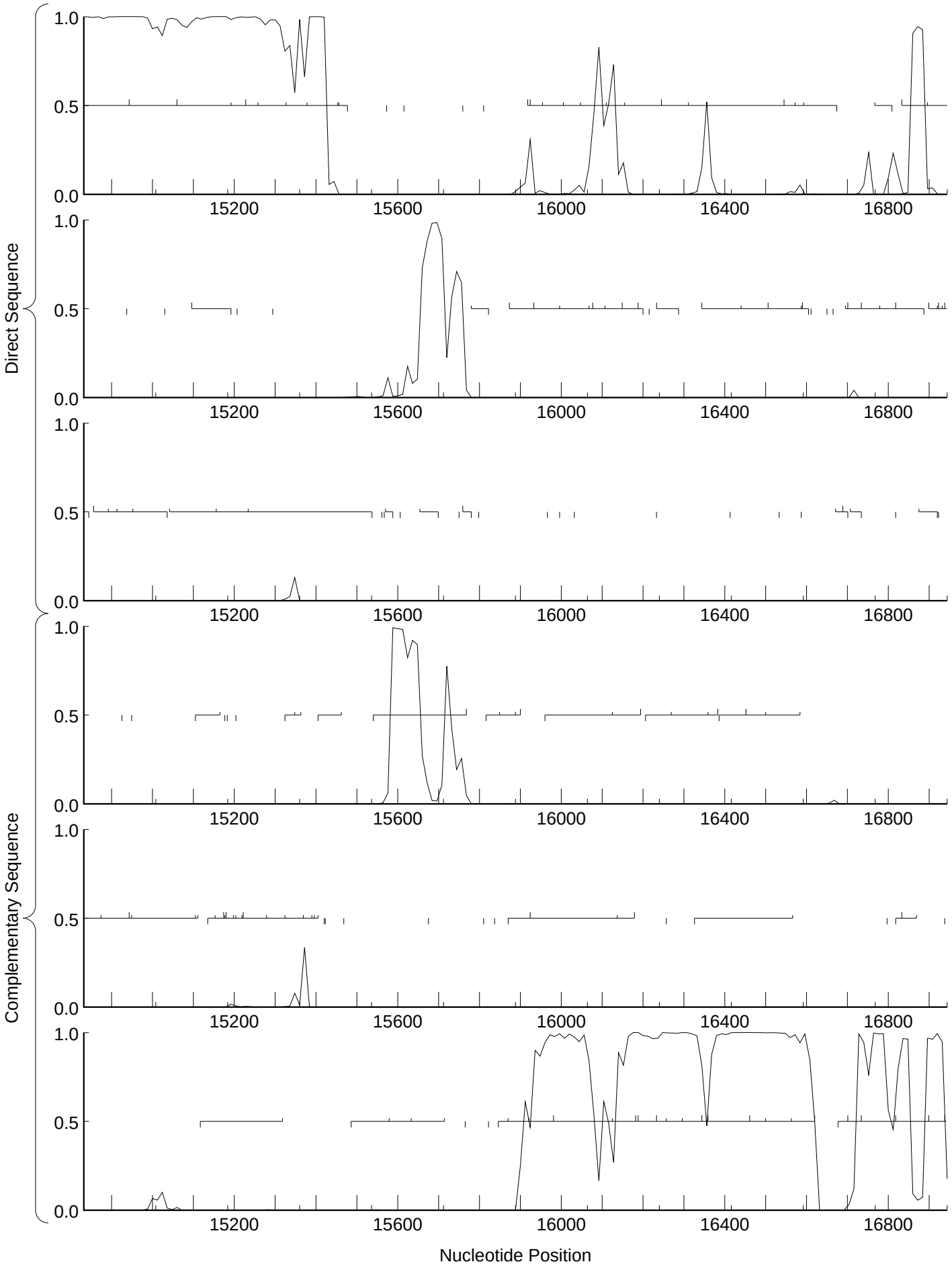

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
8800
9200
9600
10000
10400
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
10800
11200
11600
12000
12400
Nucleotide Position

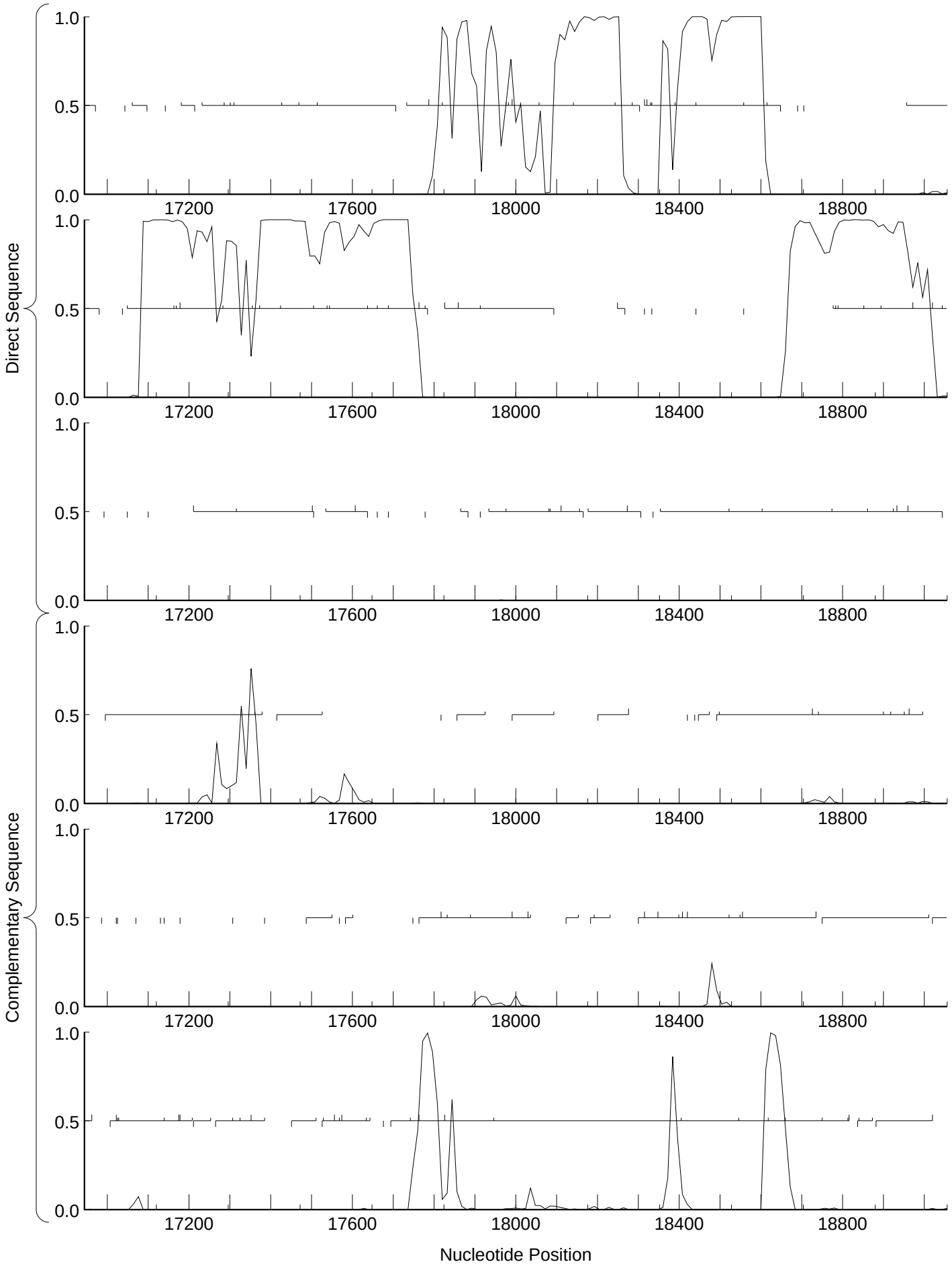
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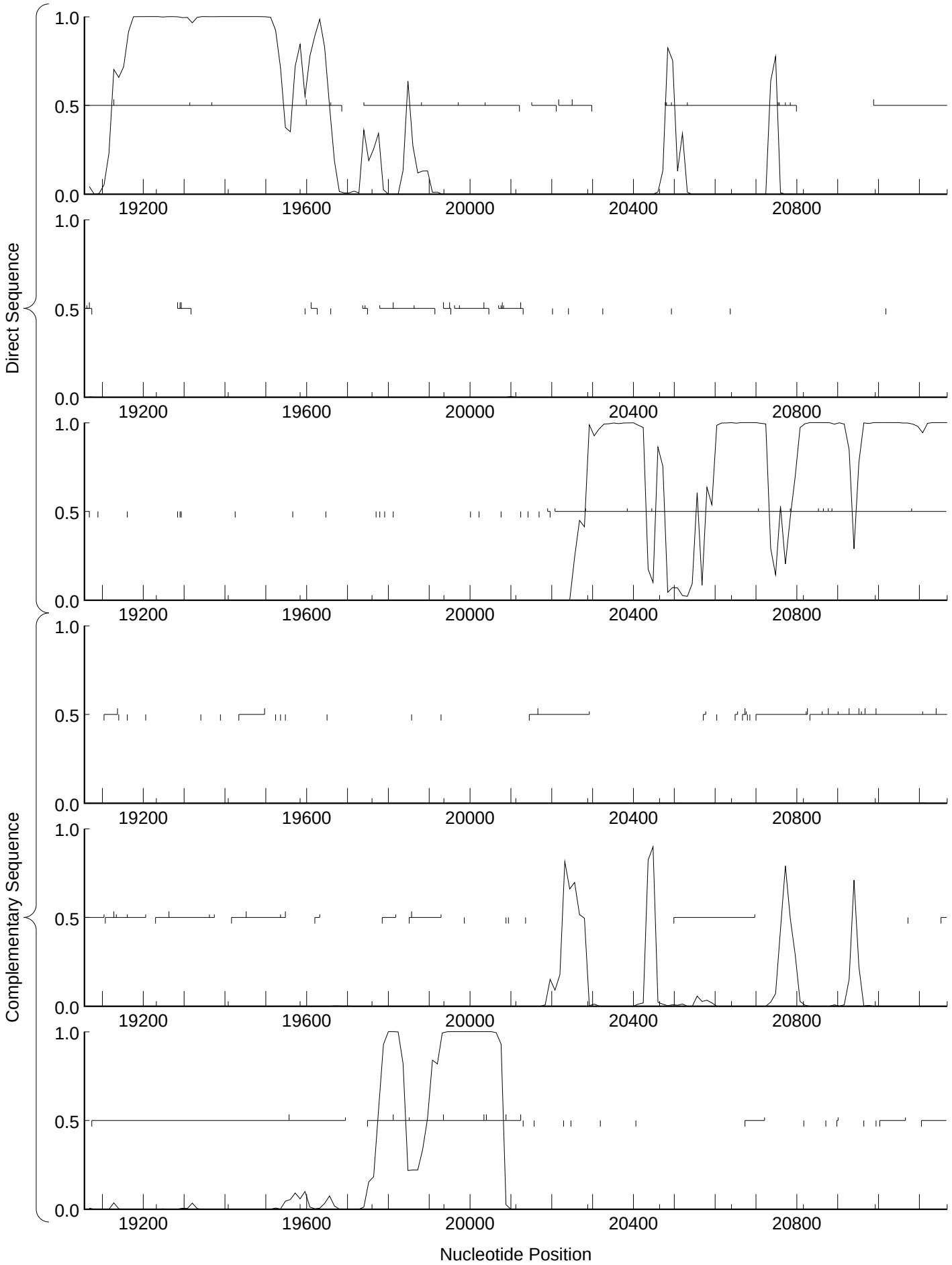
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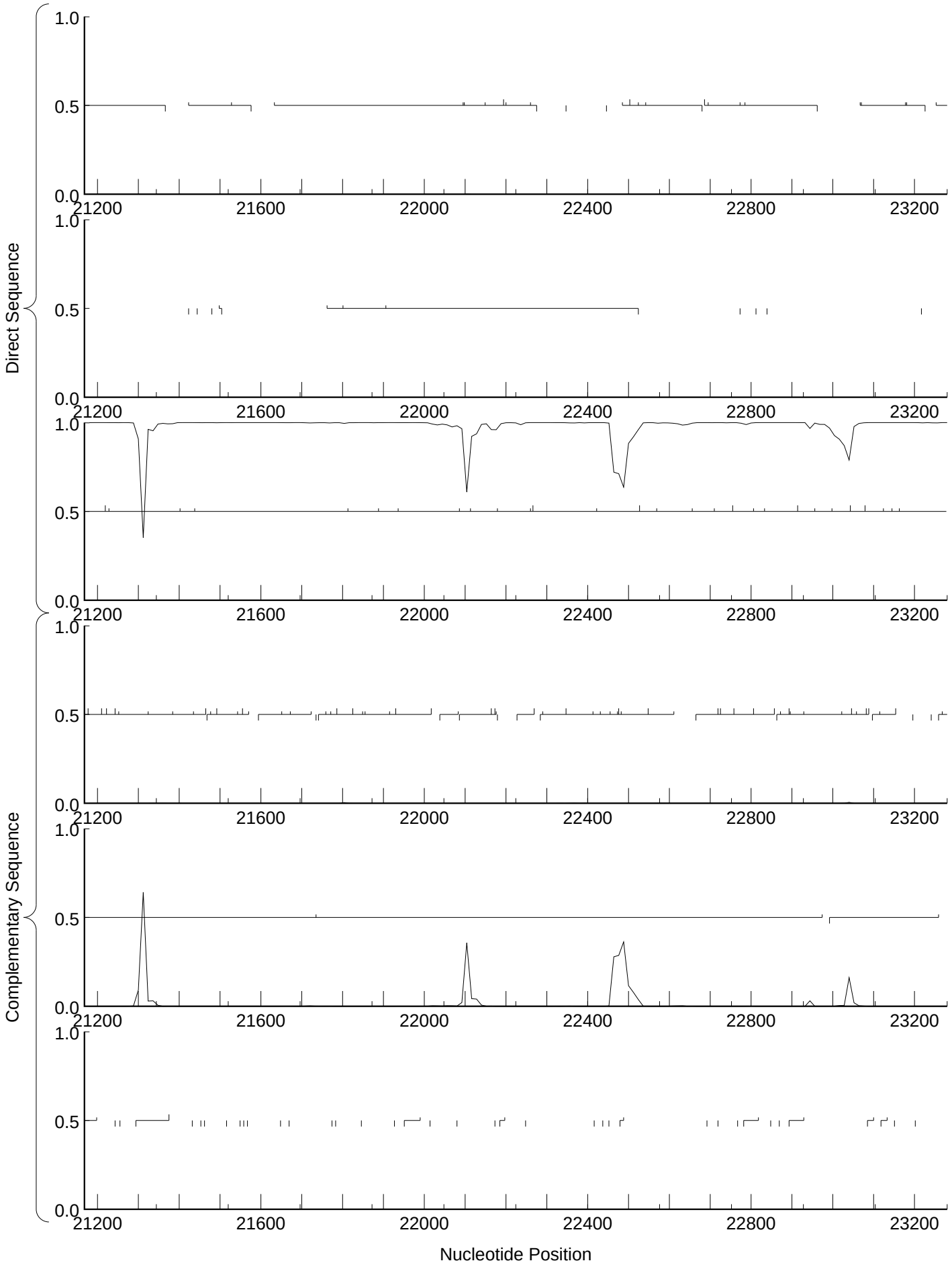
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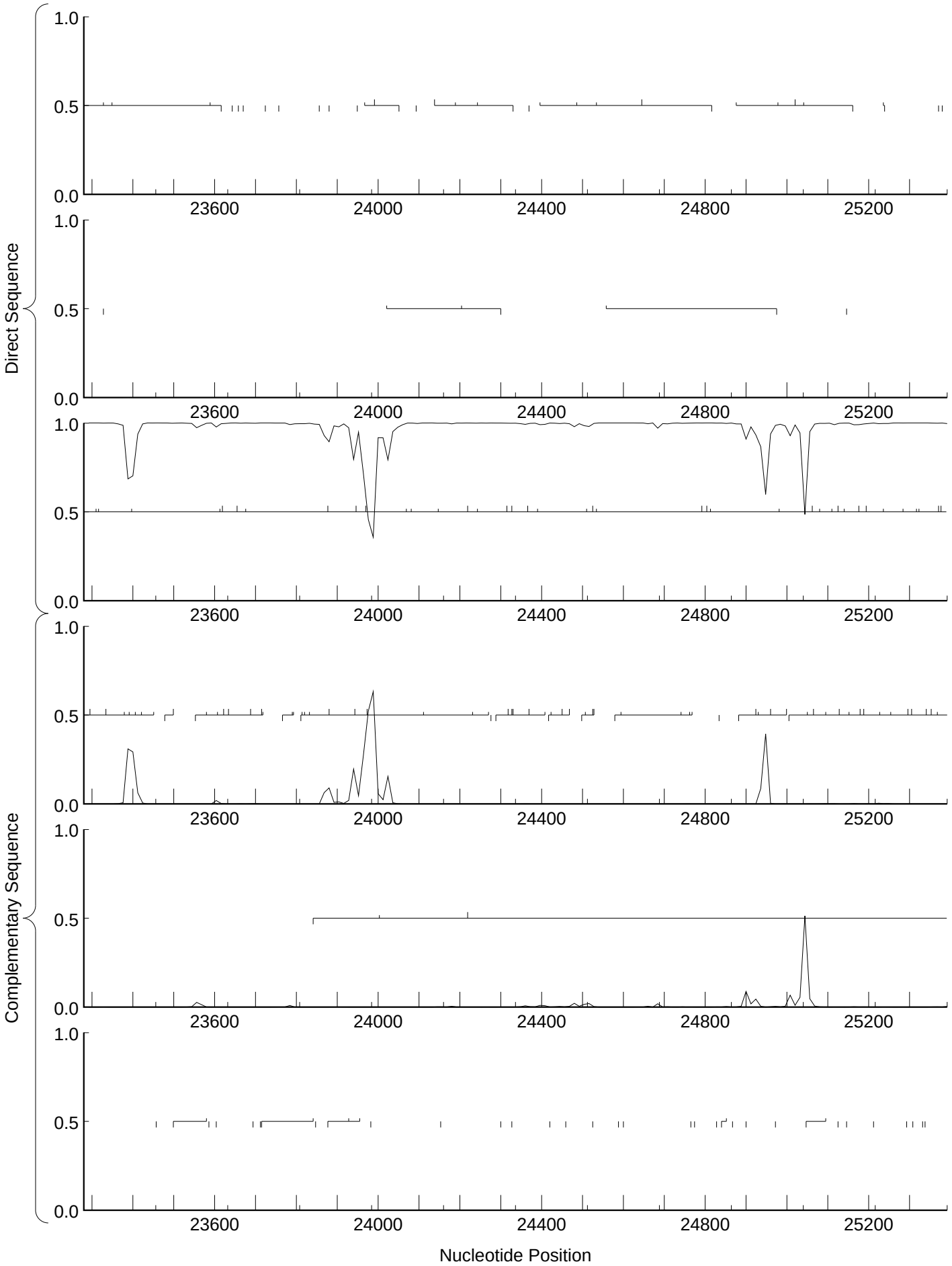
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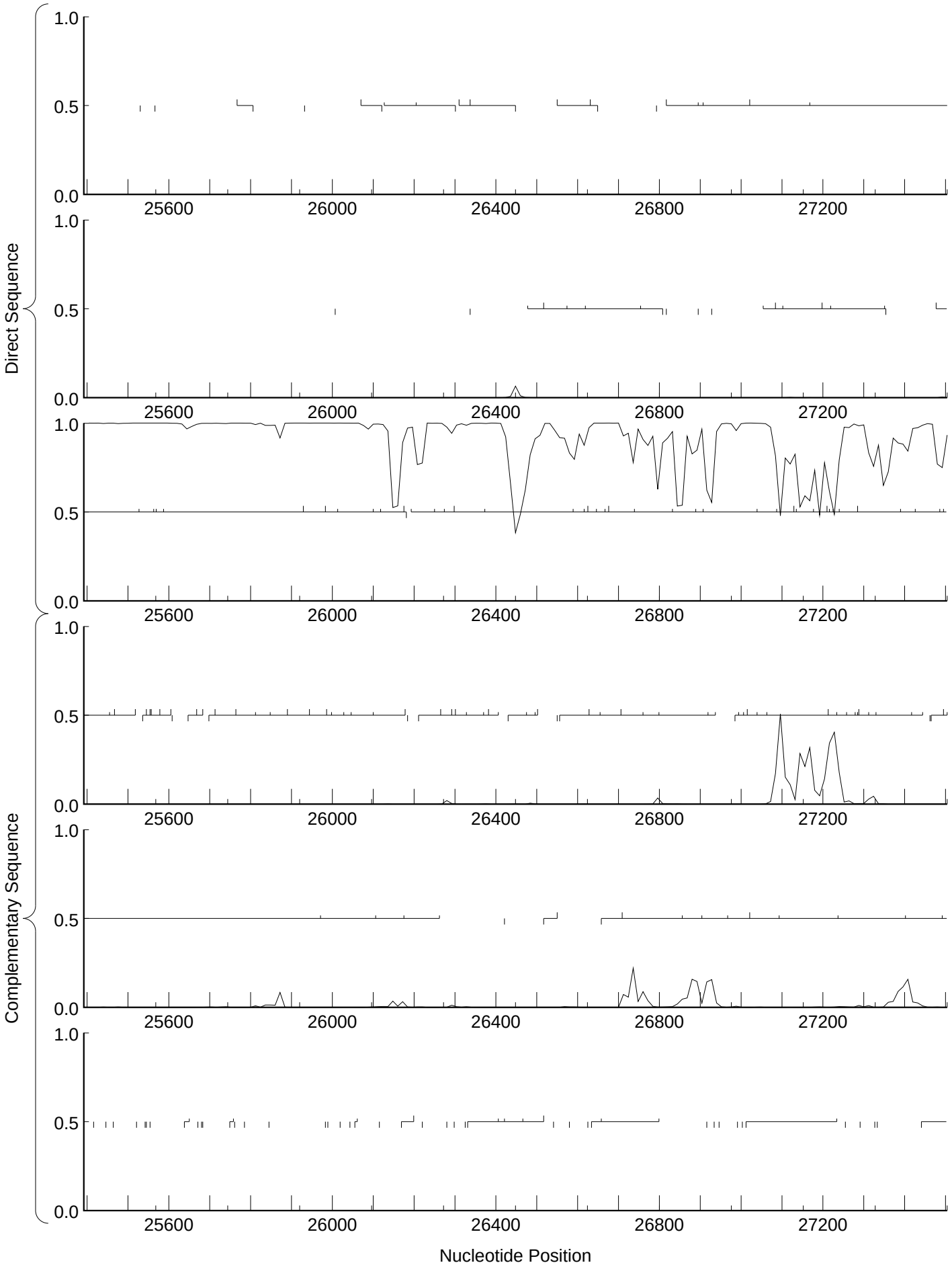
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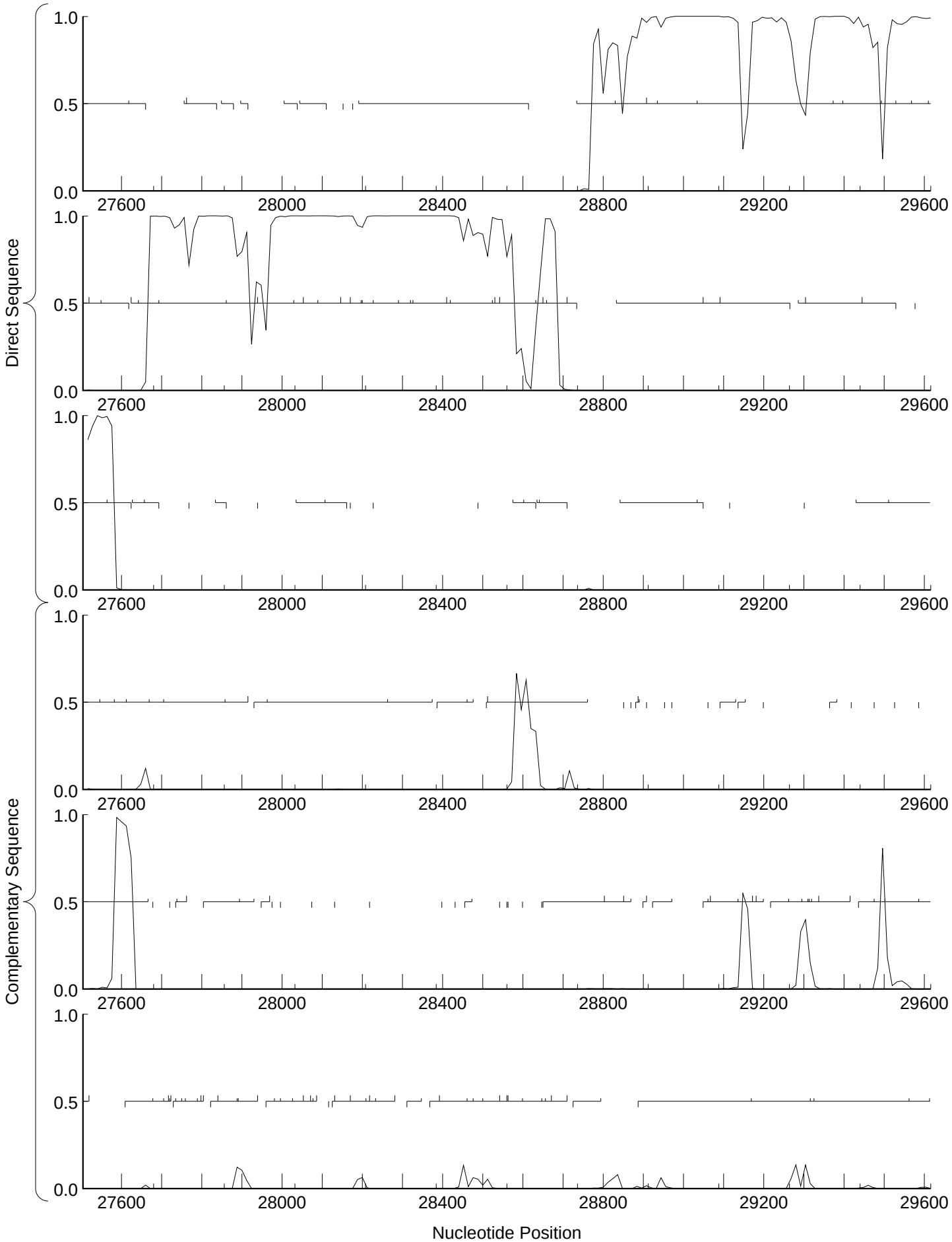
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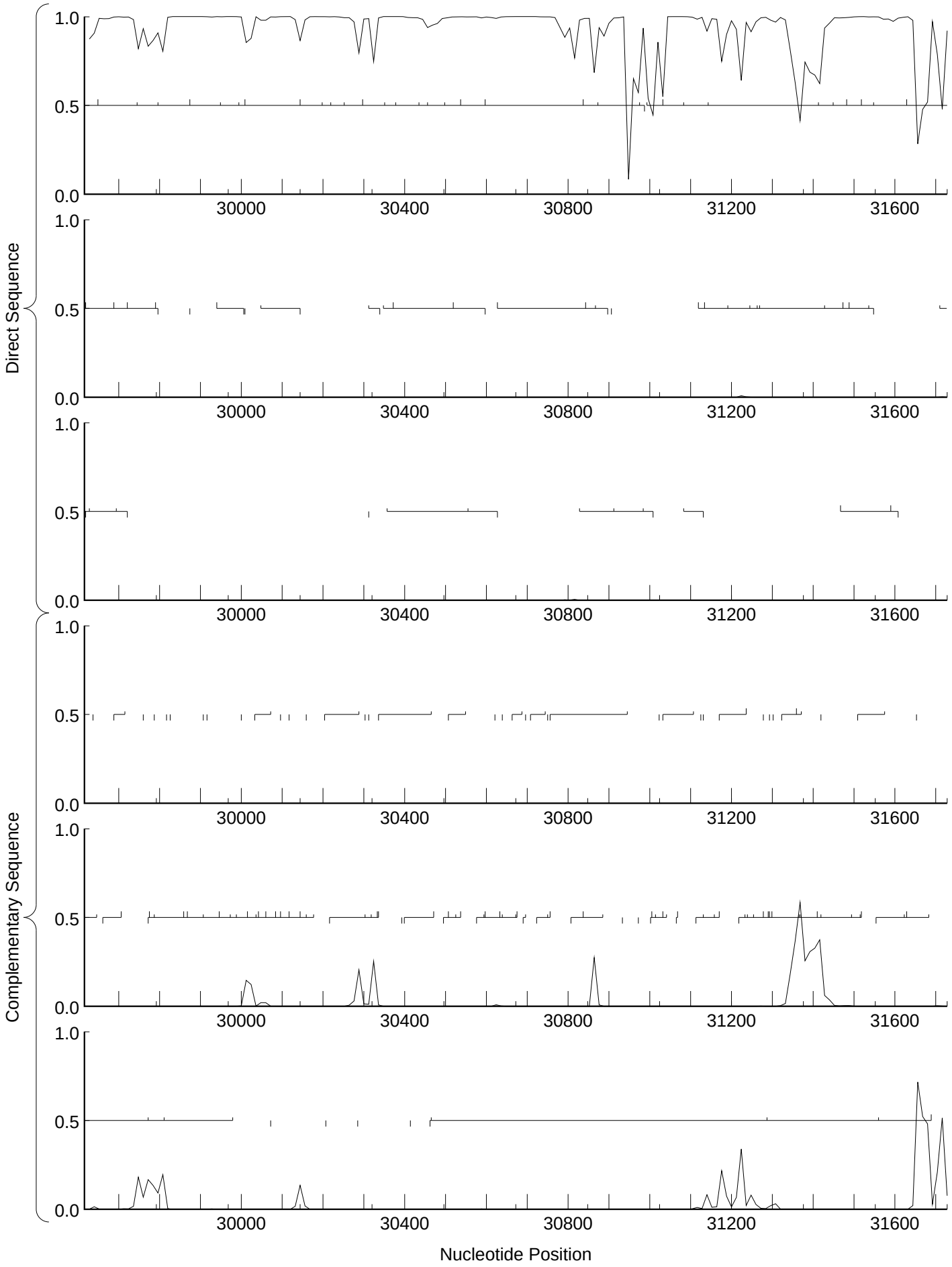
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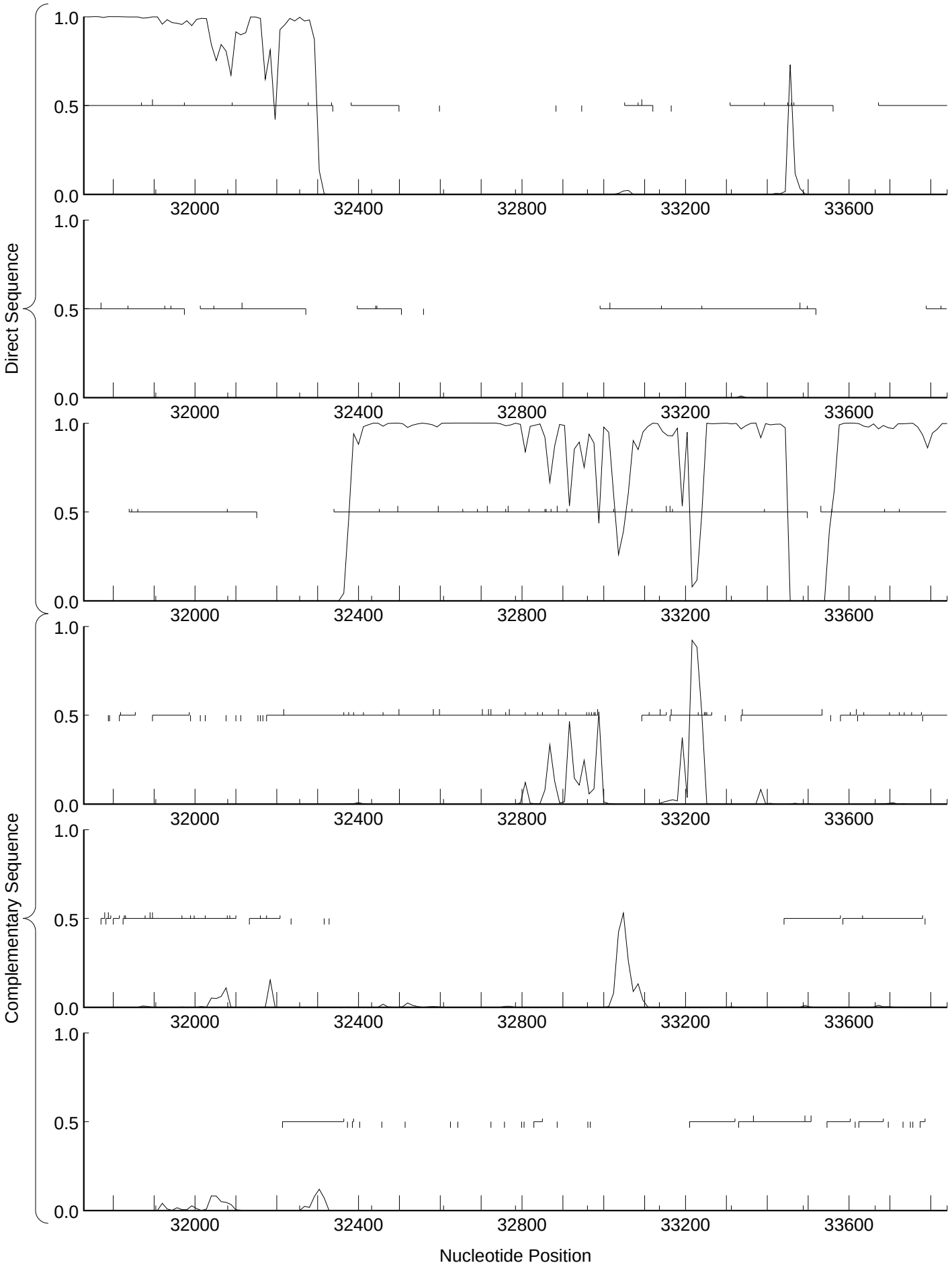
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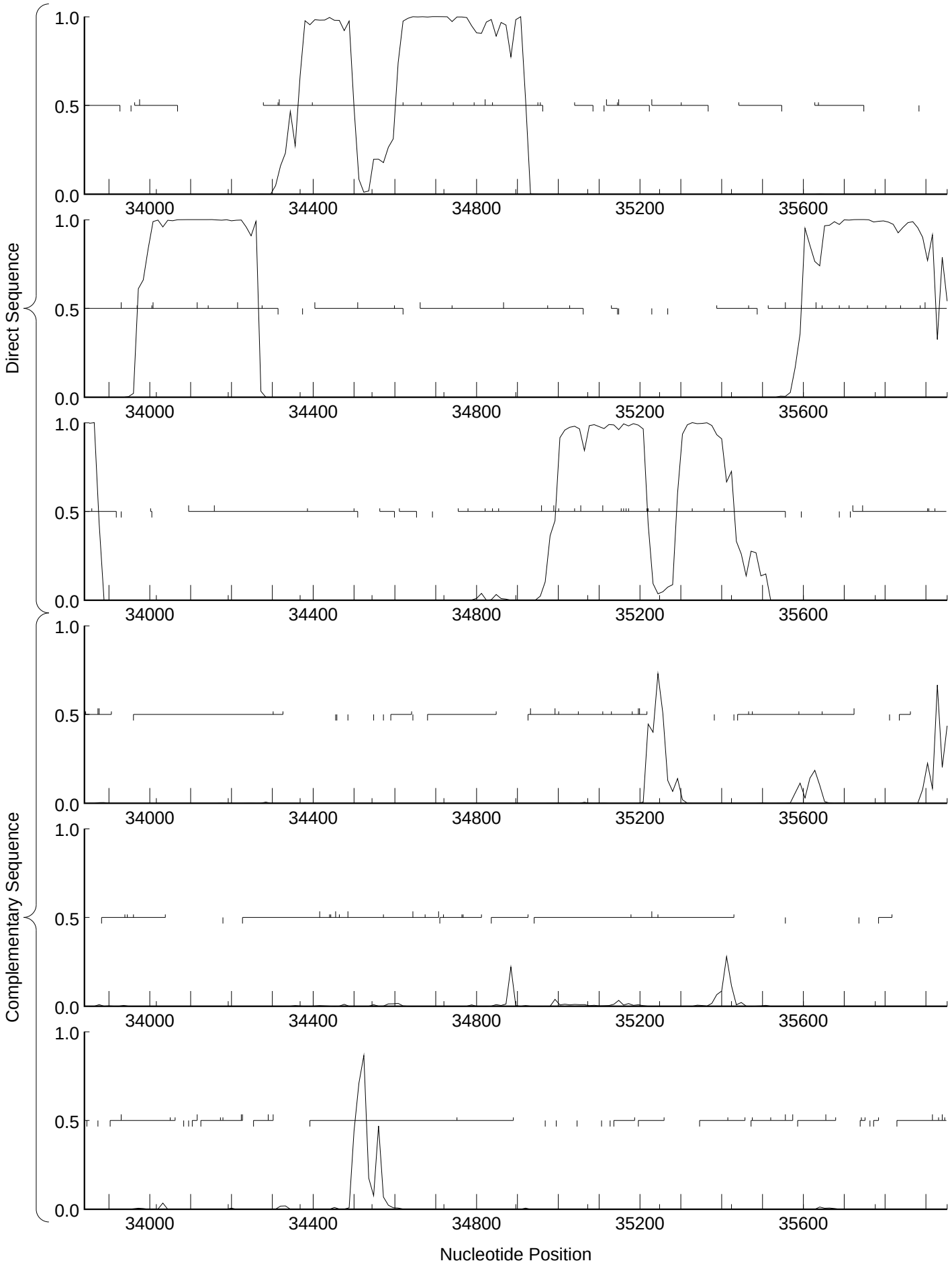
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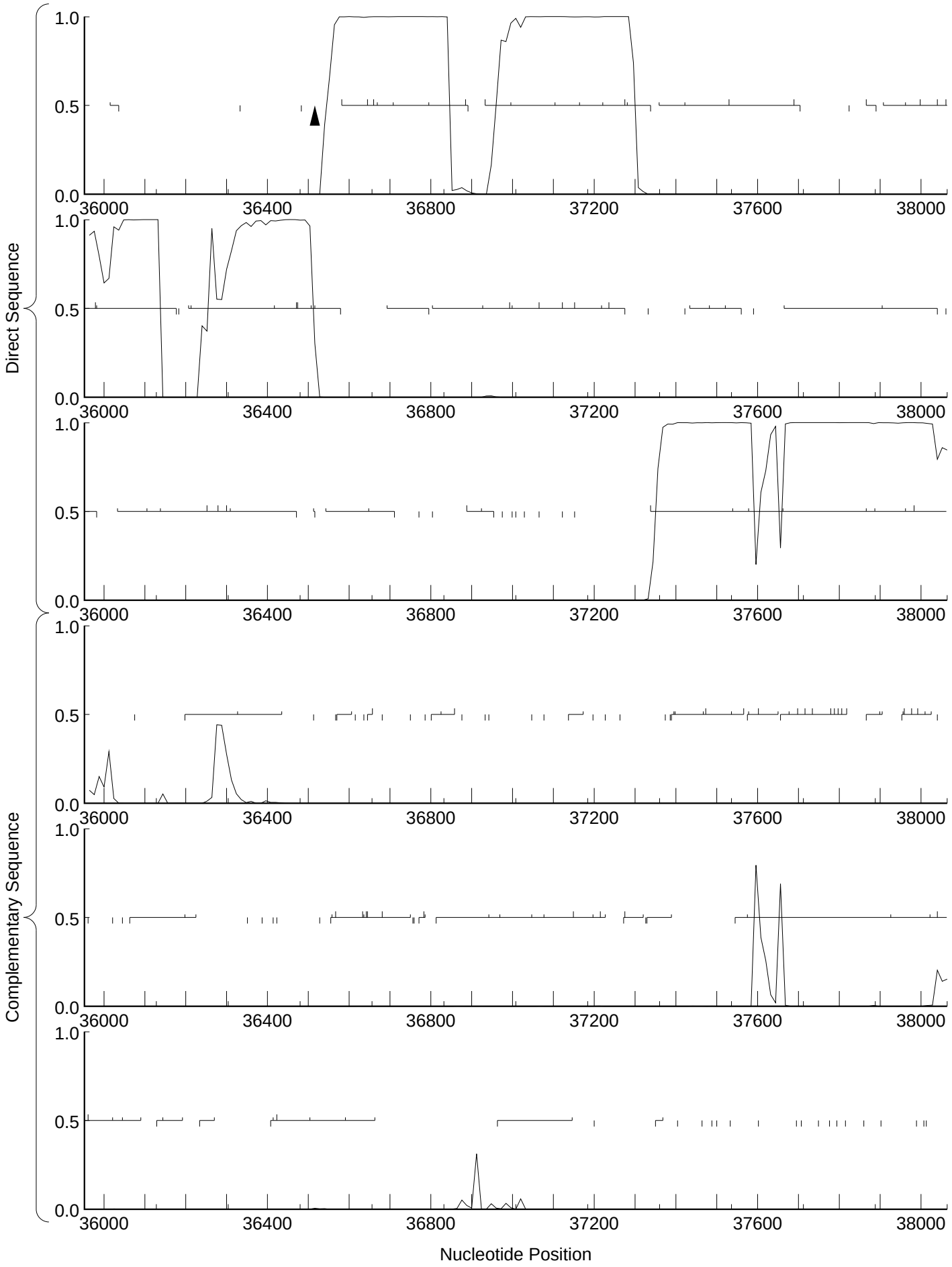
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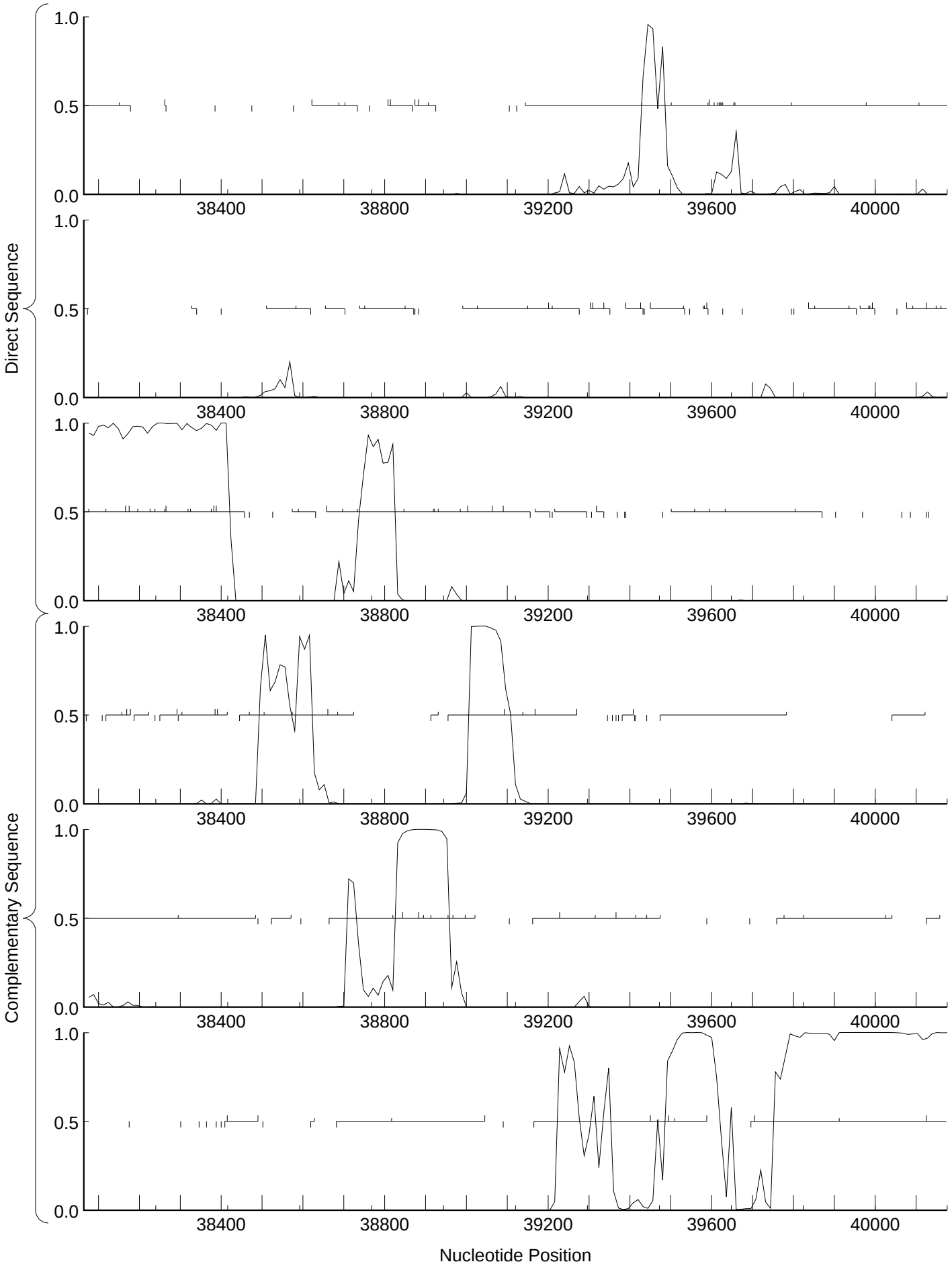
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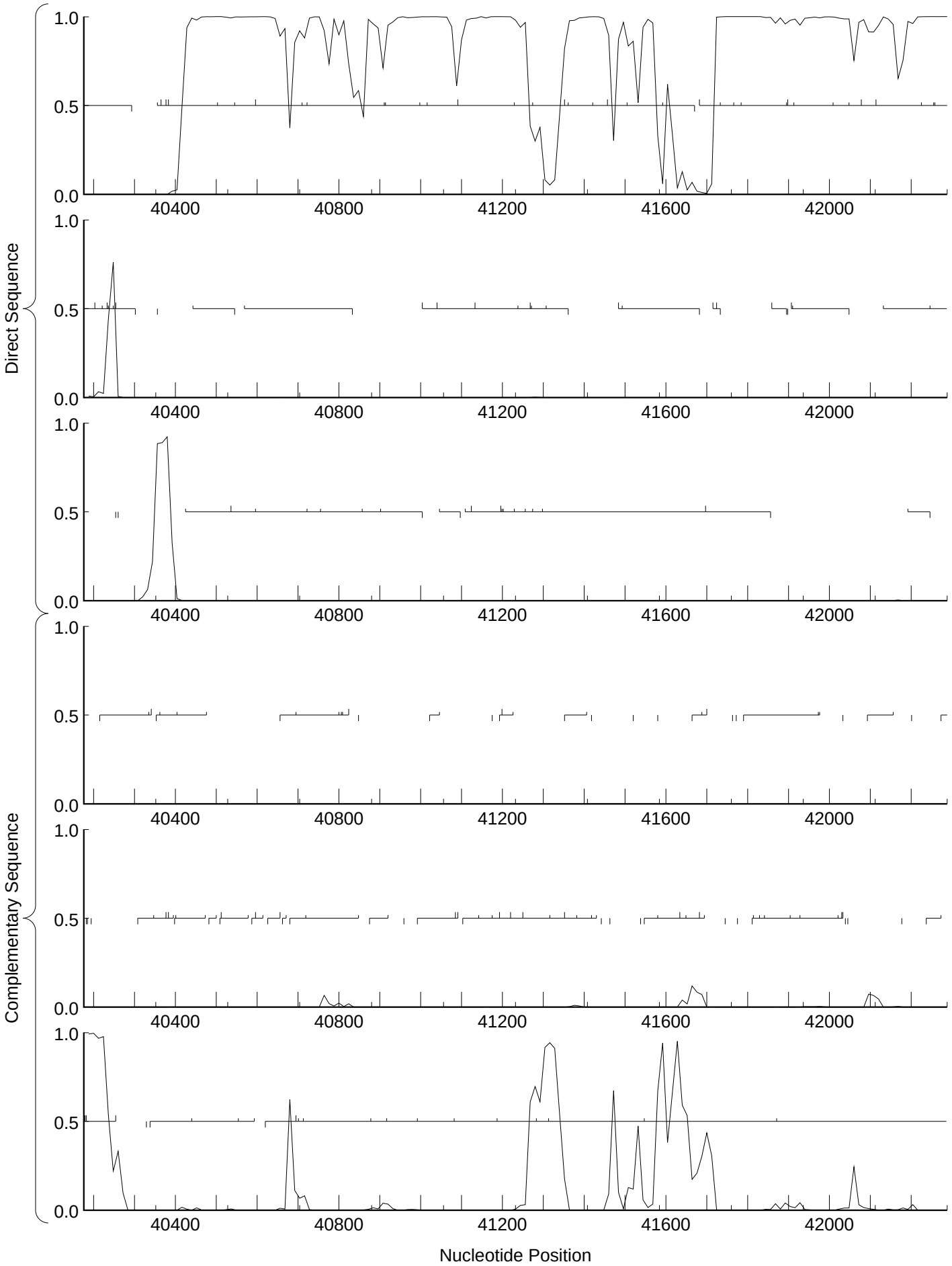
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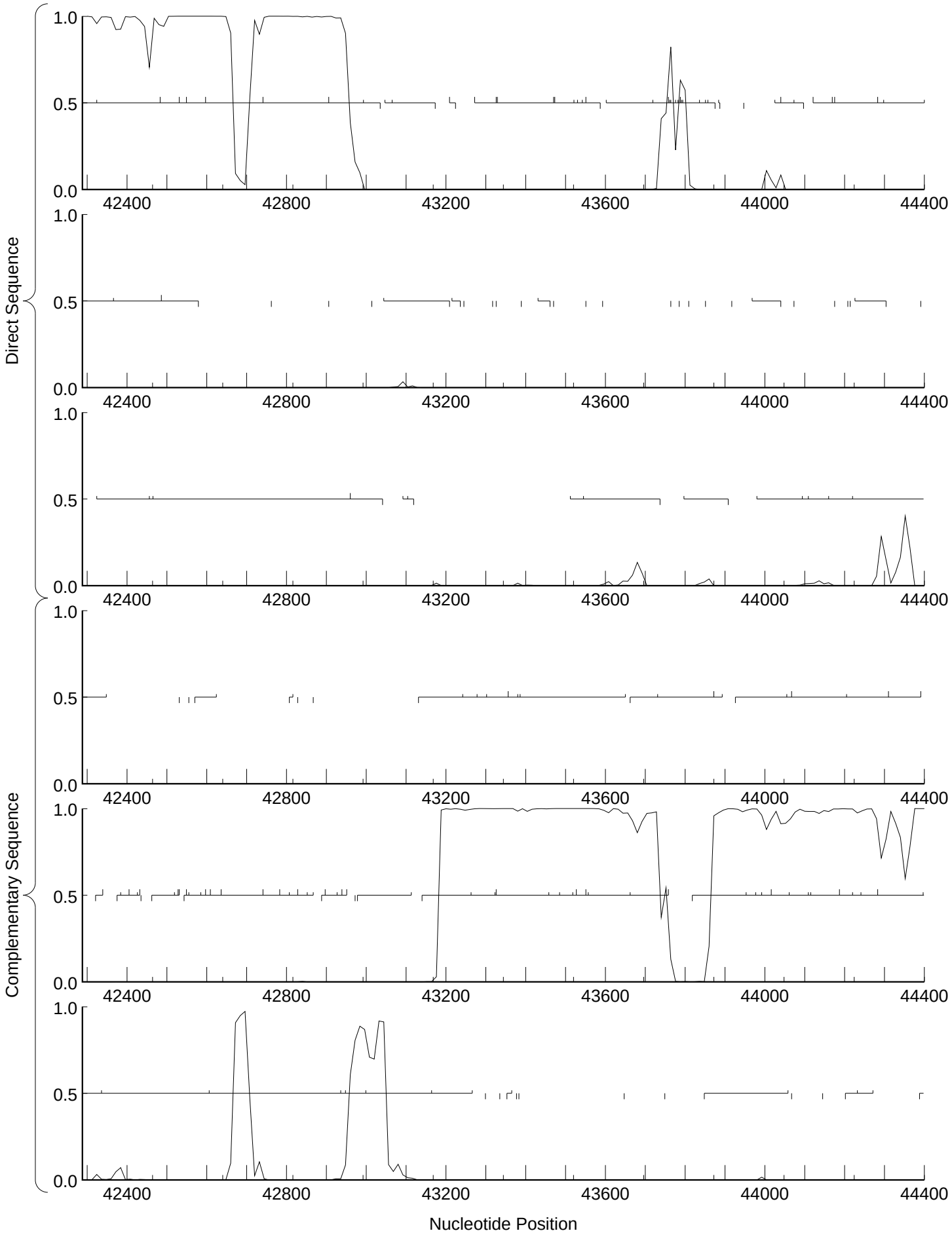
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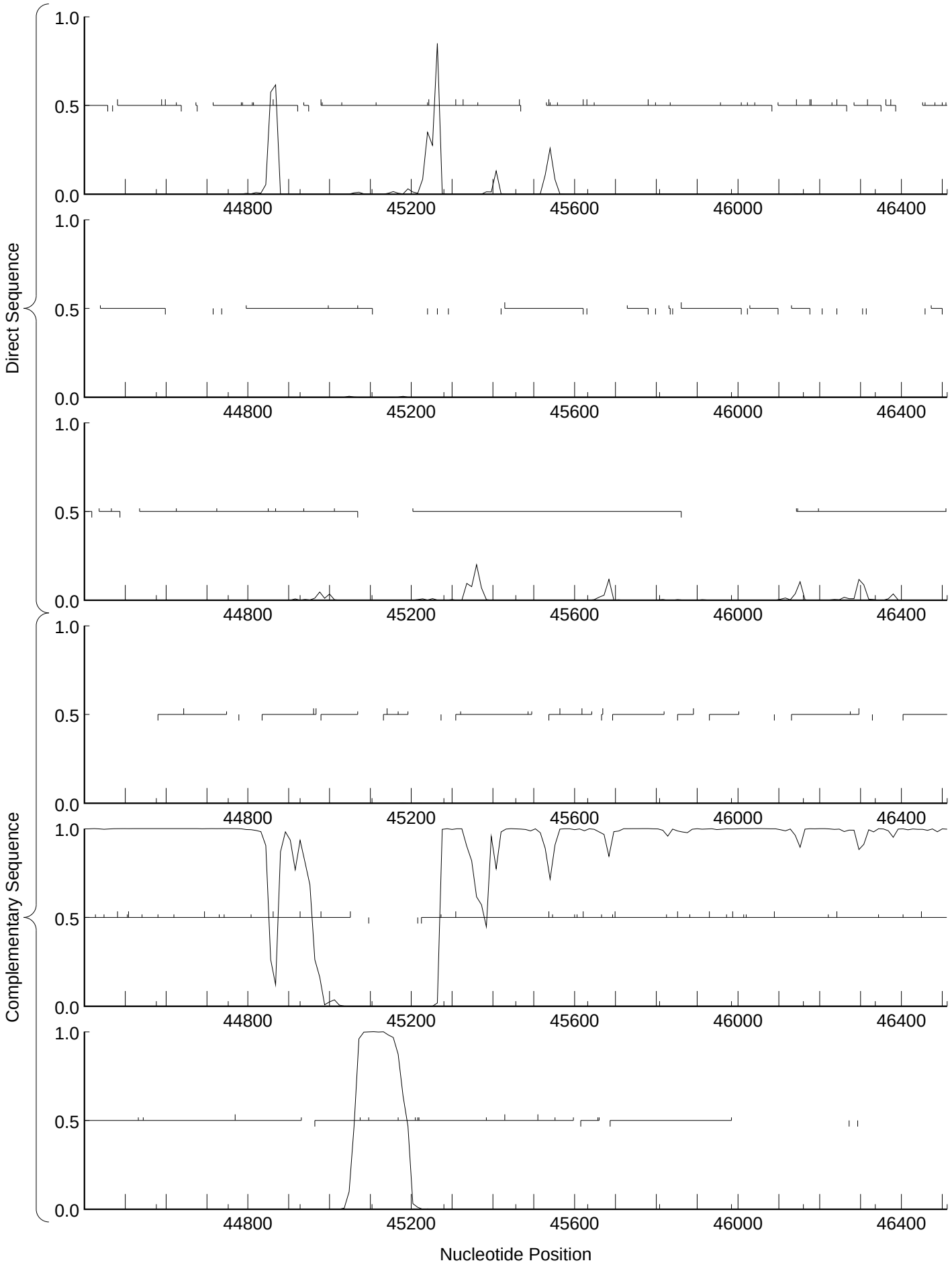
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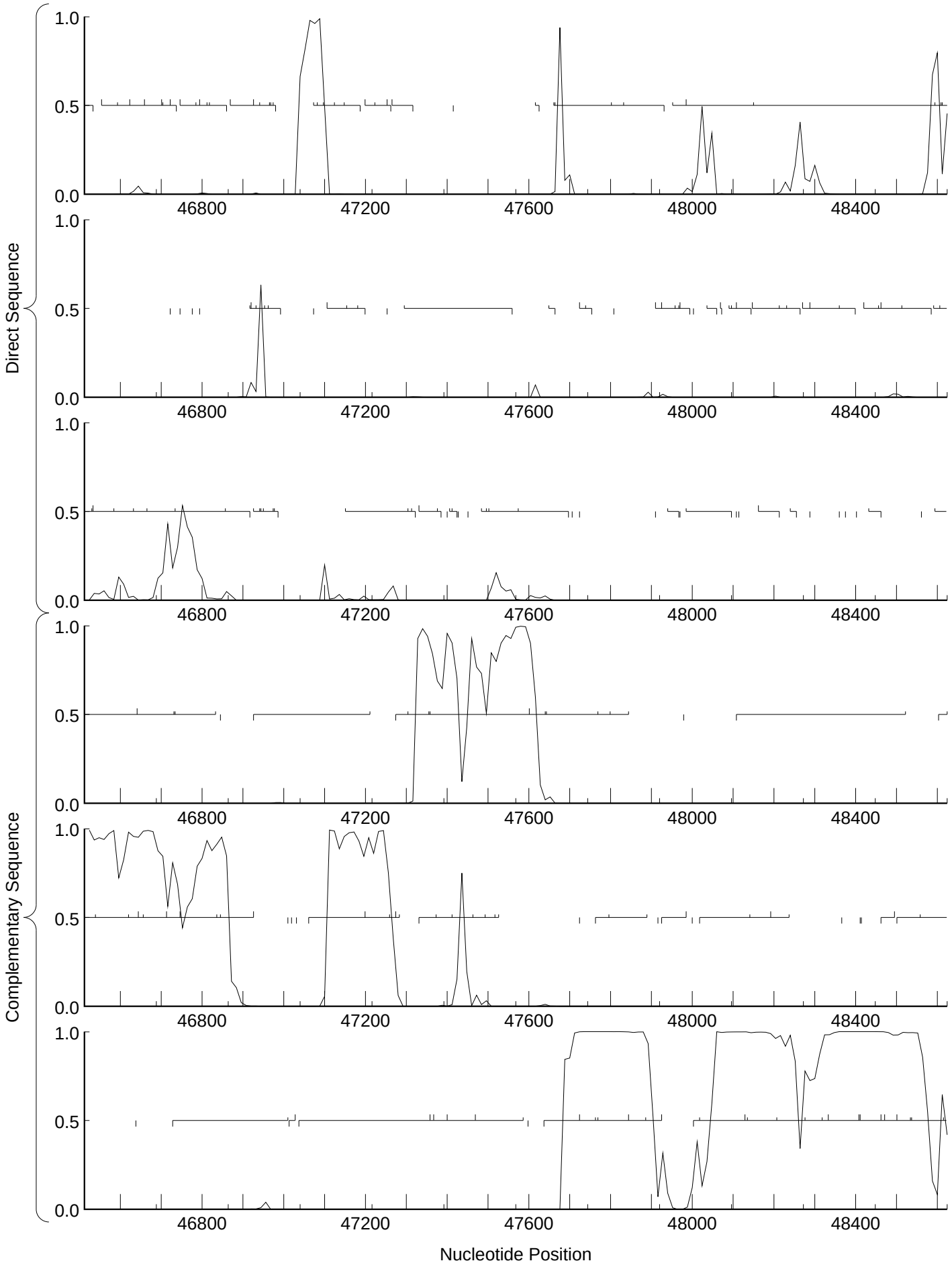
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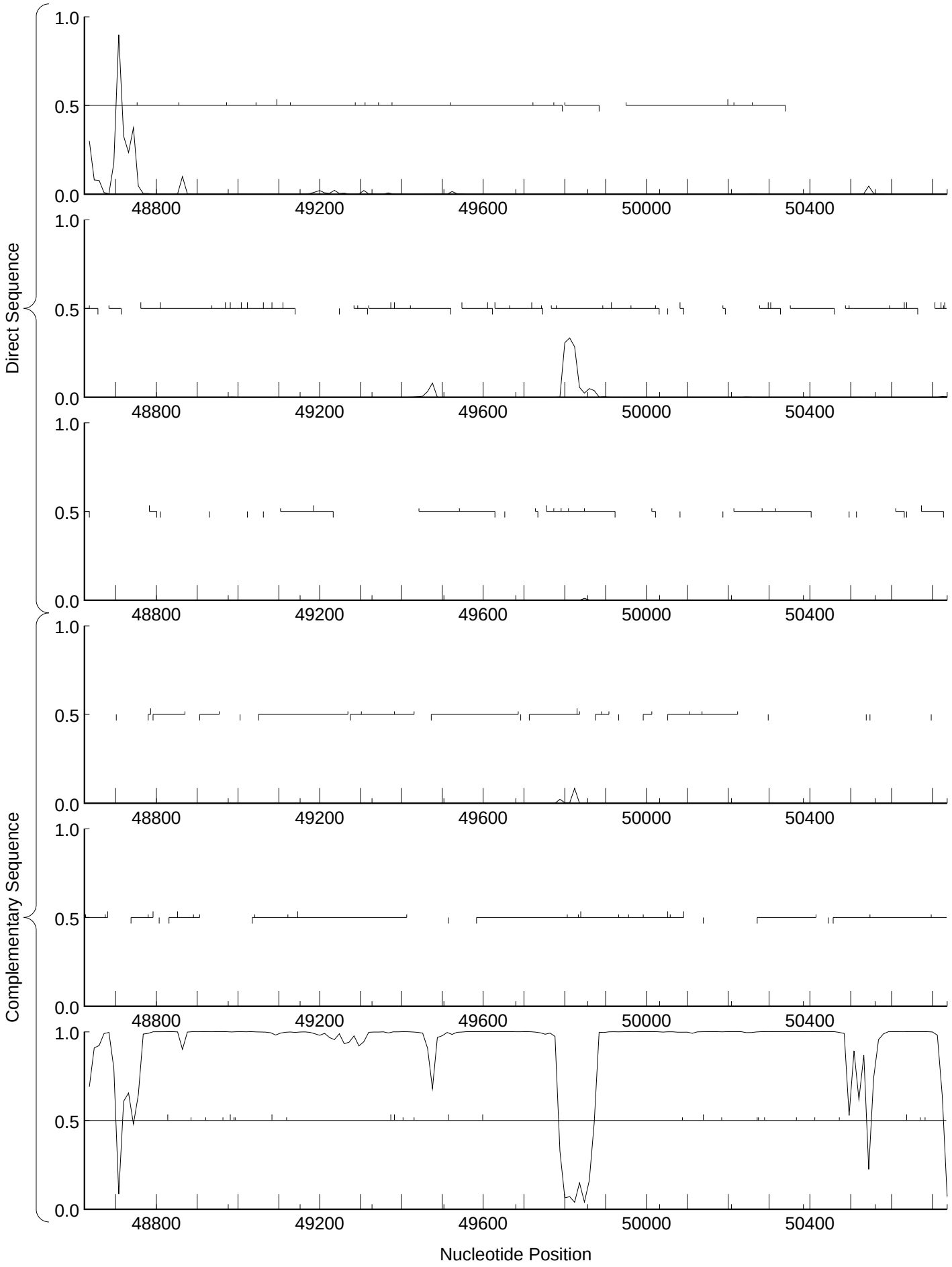
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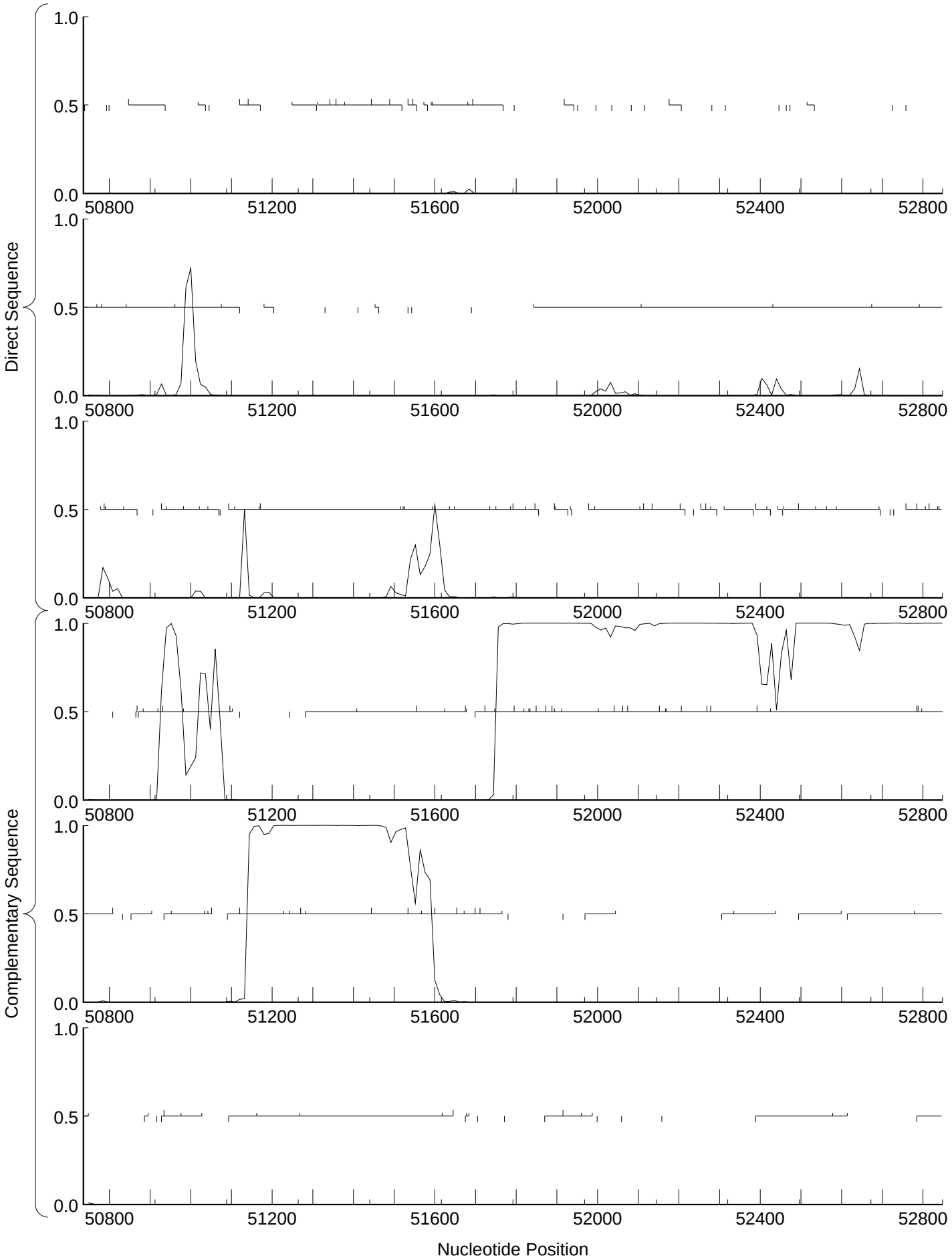
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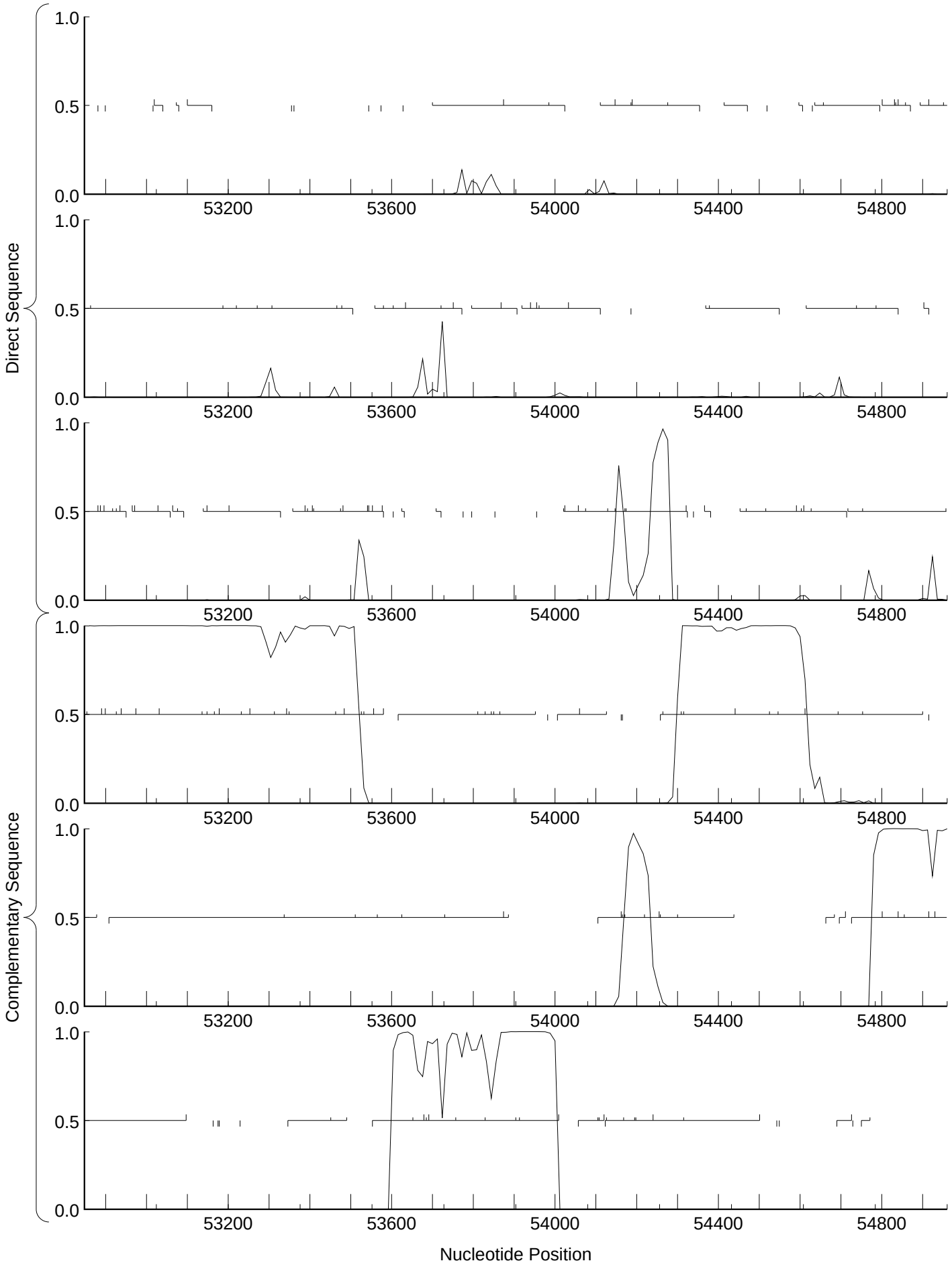
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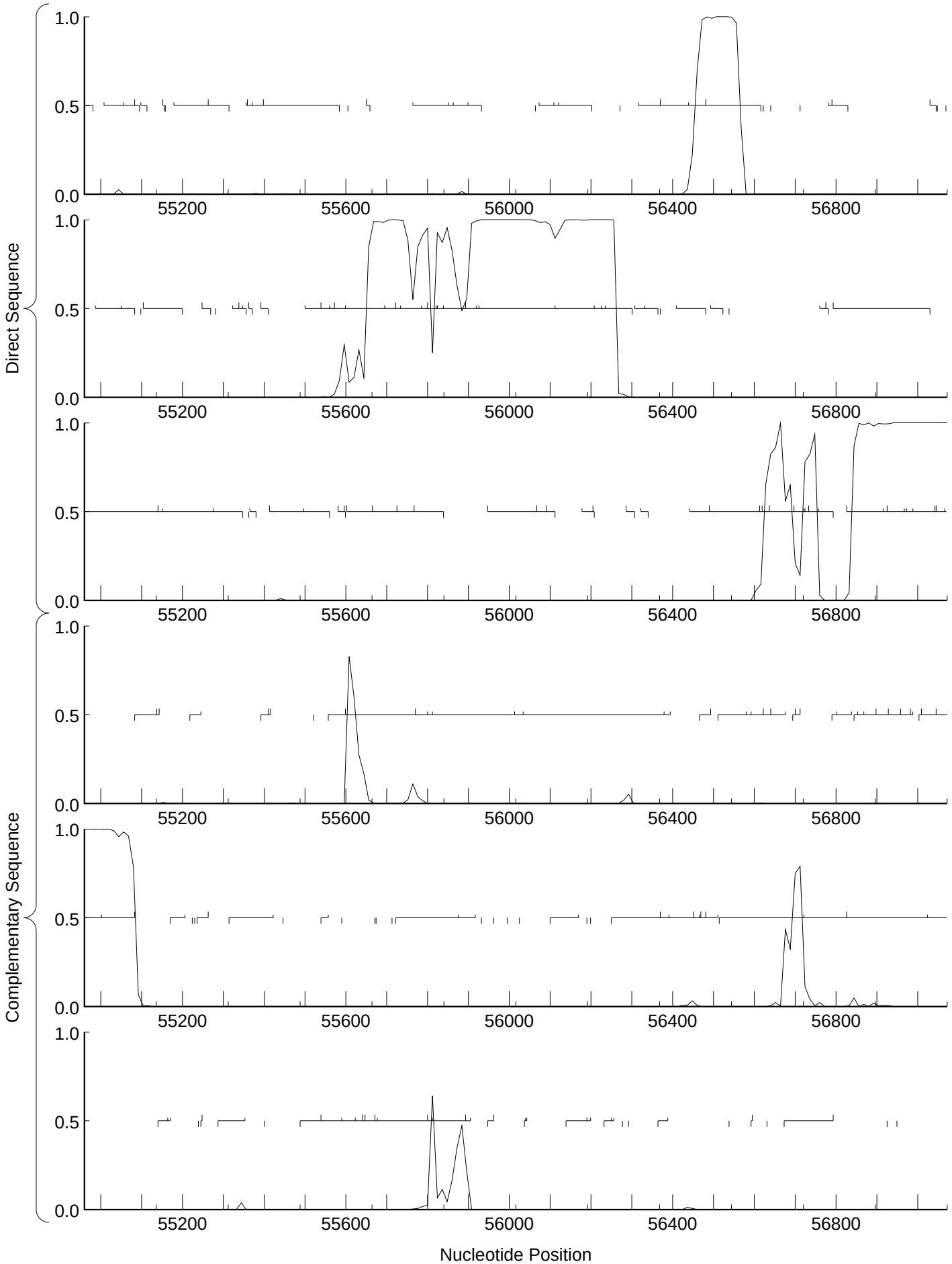
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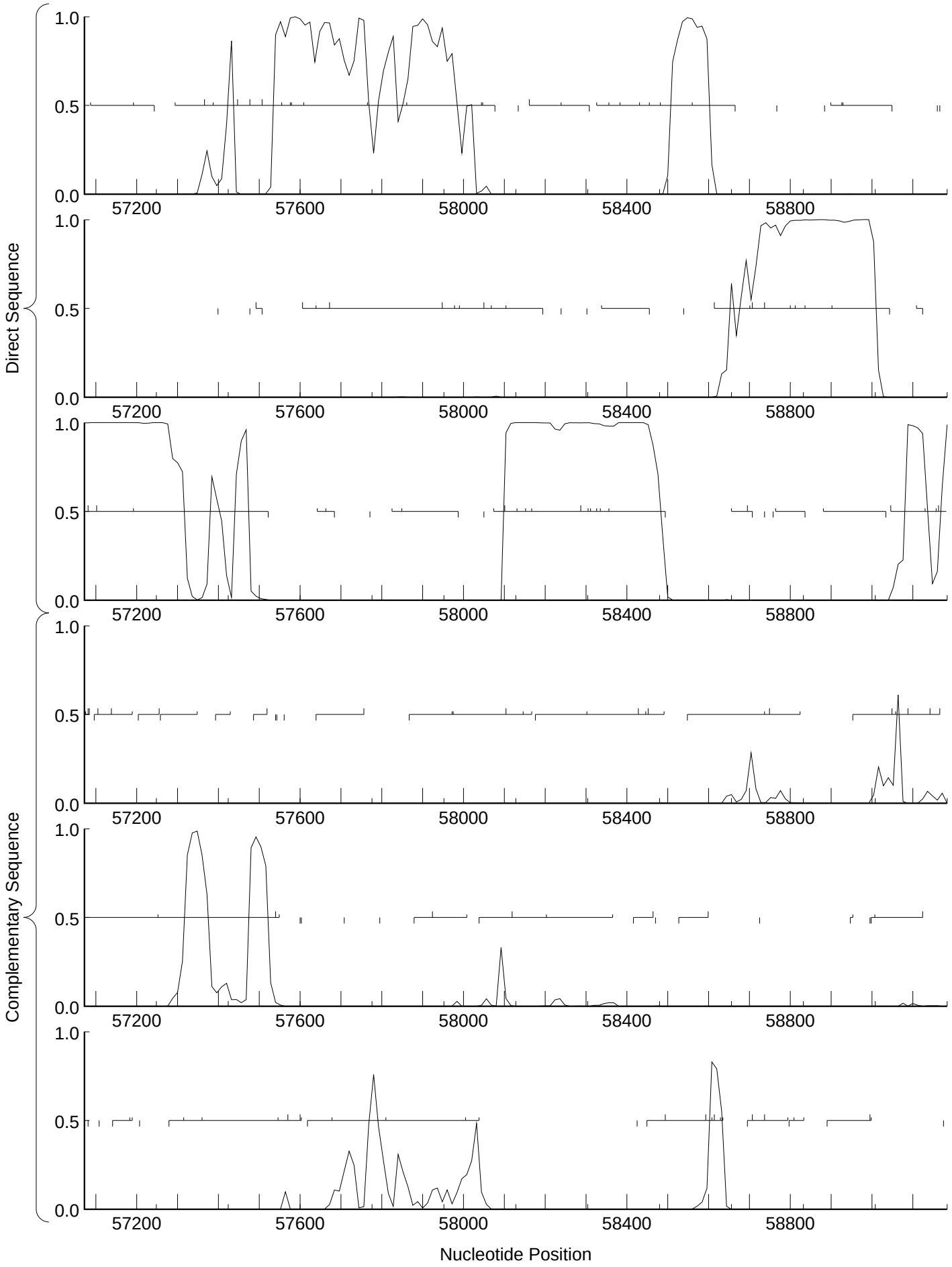
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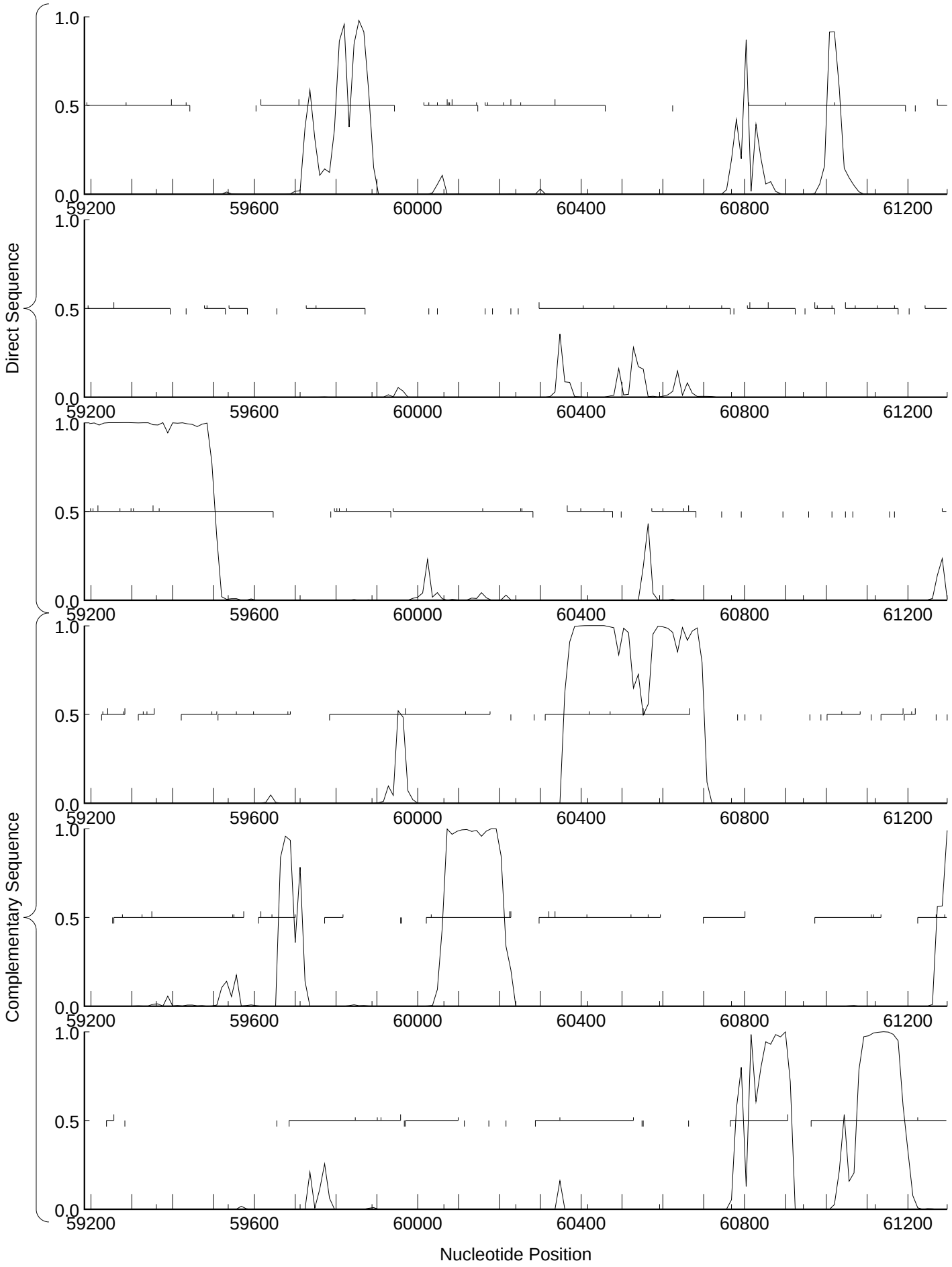
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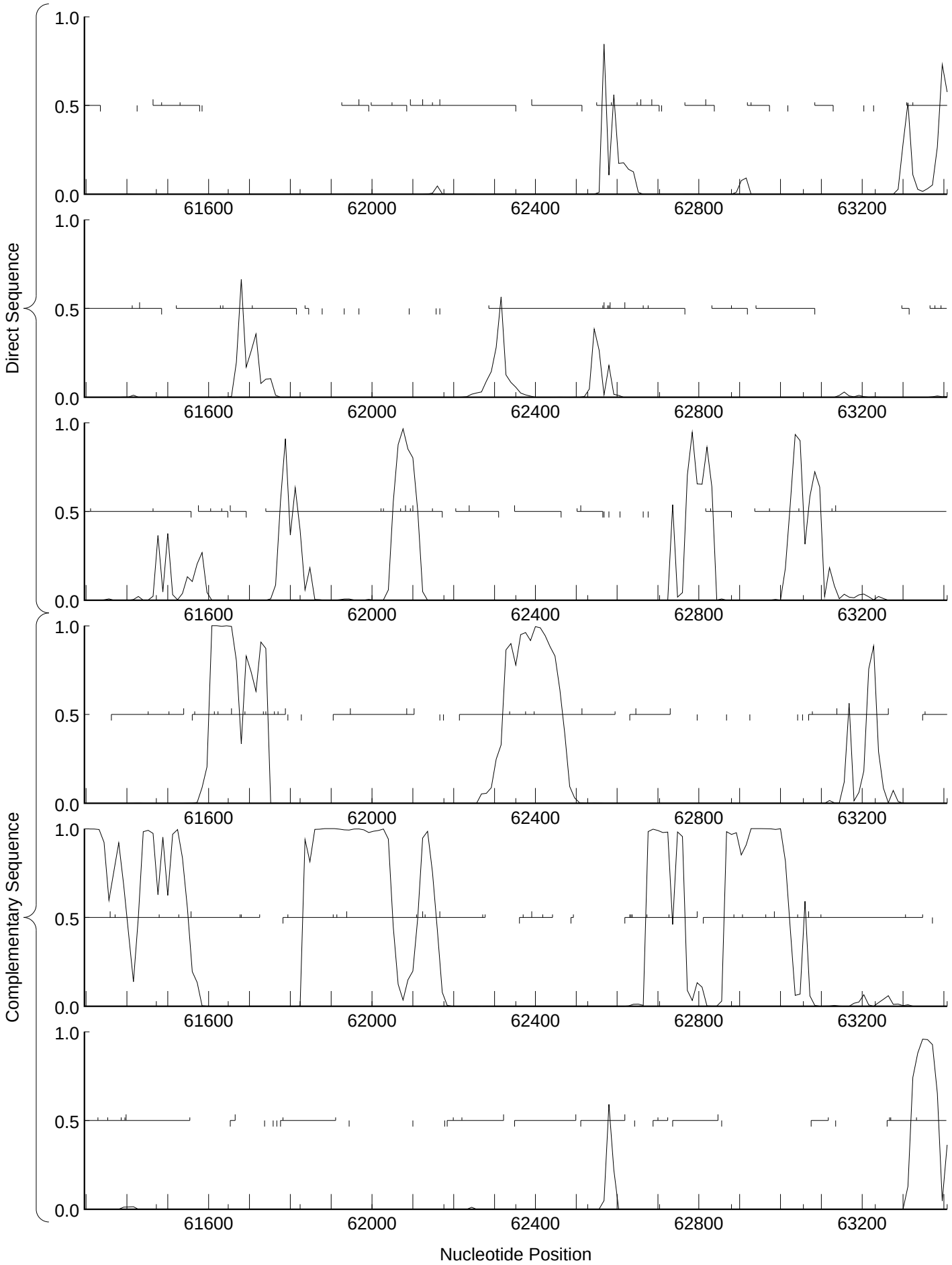
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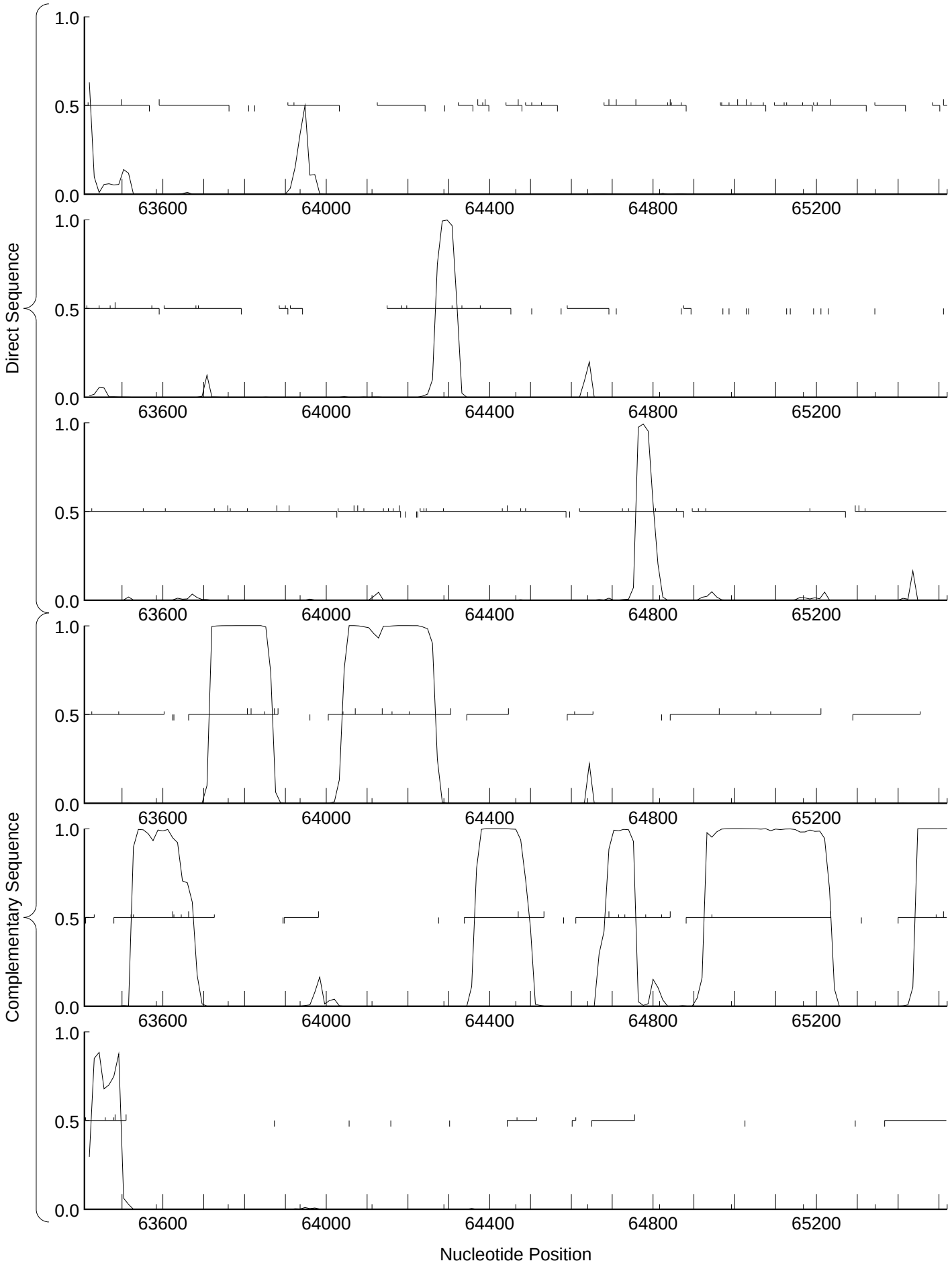
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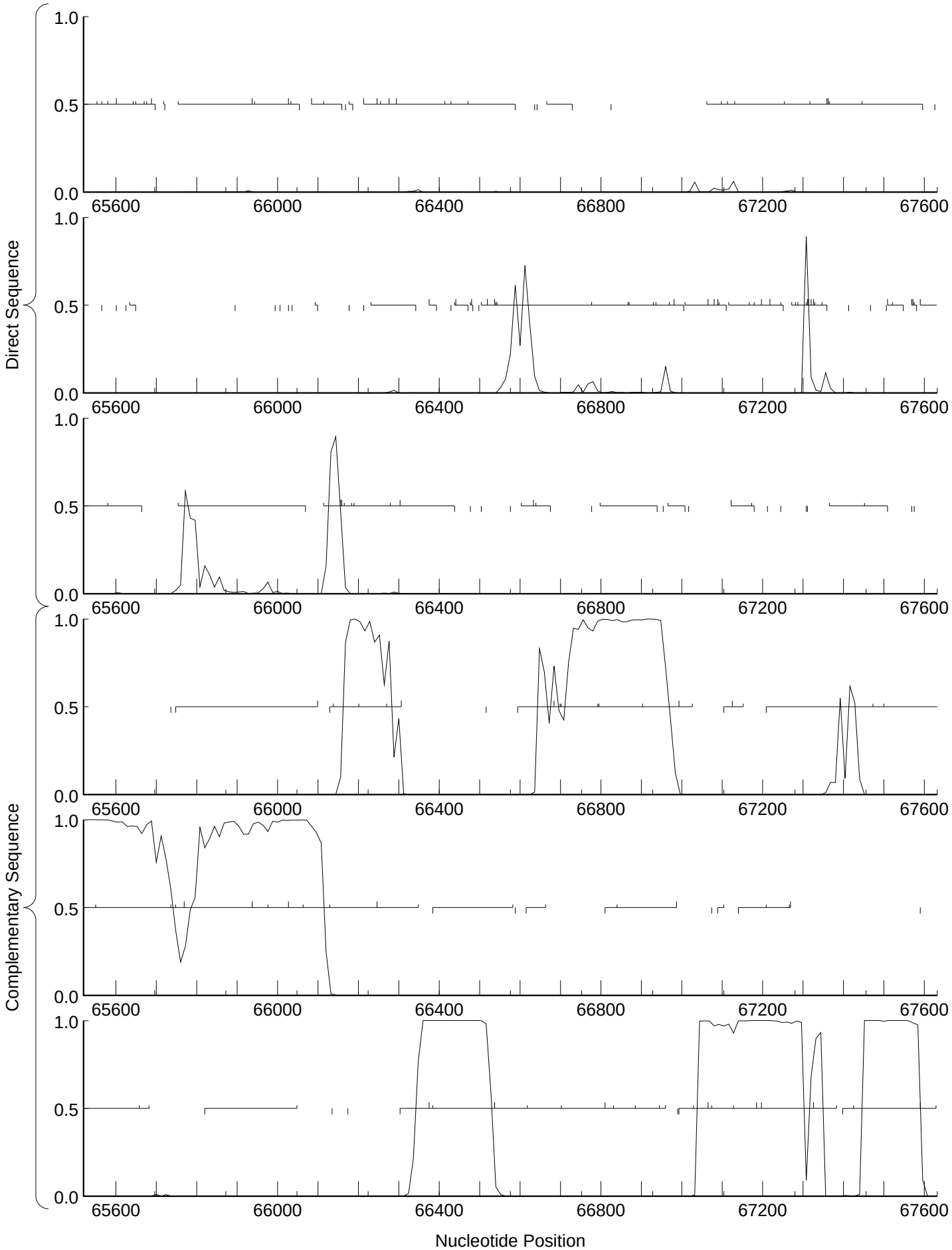
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acterium phage SDcharge11 complete sequence, 67702 bp, circularly permuted, Cluster B1 , Order 4, Window 96, Step 12, 32/34



acterium phage SDcharge11 complete sequence, 67702 bp, circularly permuted, Cluster B1 , Order 4, Window 96, Step 12, 33/34



acterium phage SDcharge11 complete sequence, 67702 bp, circularly permuted, Cluster B1 , Order 4, Window 96, Step 12, 34/34

