# GeneMark

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

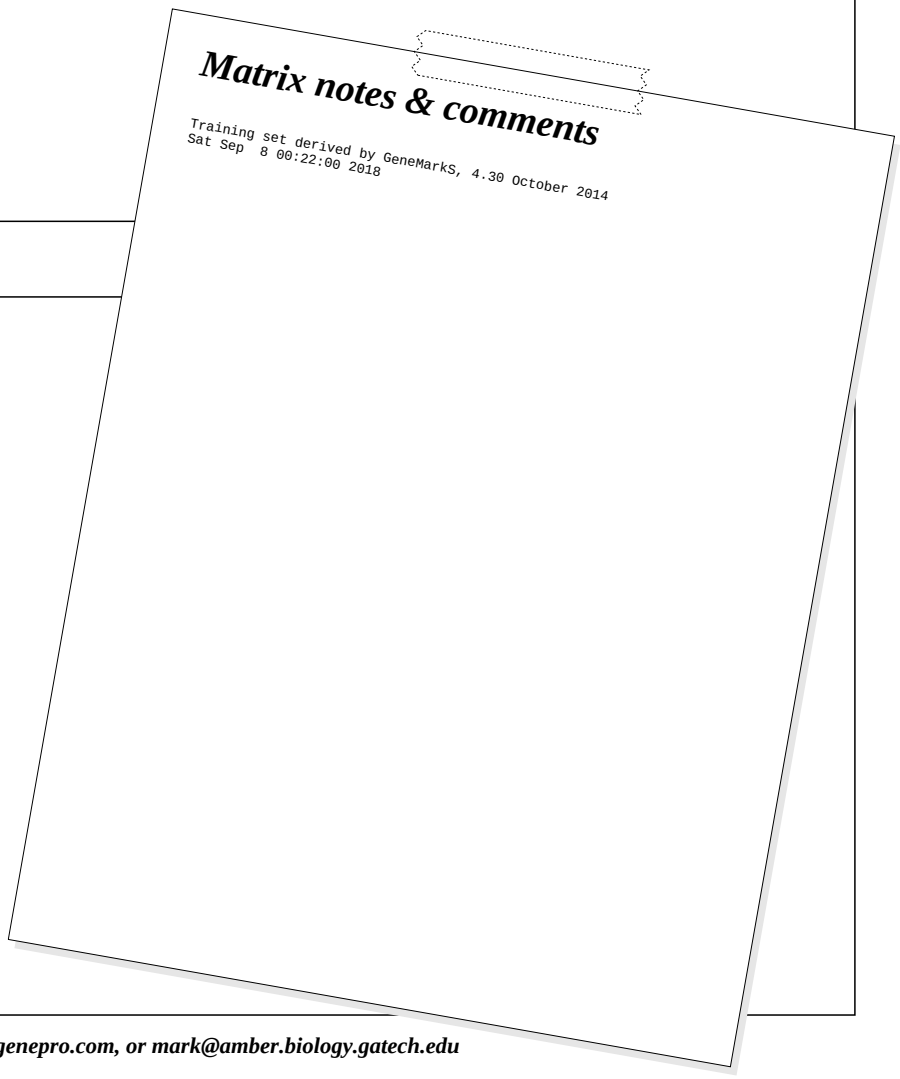
## PROGRAM INFORMATION

```
Sequence         :  Arthrobacter phage Eesa complete sequence, 39792 bp including 12-base 3’ overhang (GAGTCGCC
Analysis Date    :  3/29/24 at 11:07:28
Pages            :  20
Sequence Length  :  39792 bp
GC Content       :   67.09%

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

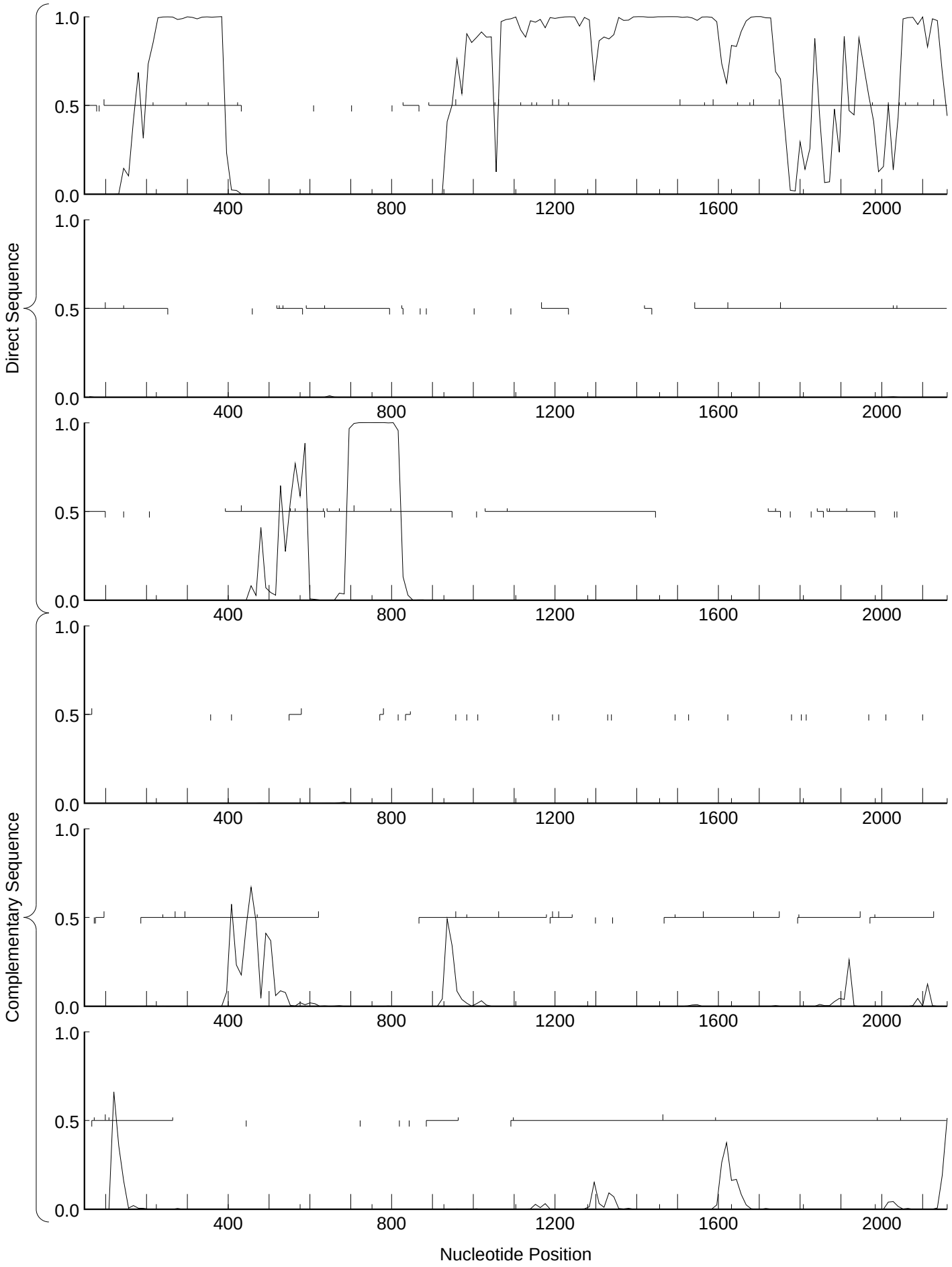
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Matrix     :  -
Author     :  -

Order      :  4

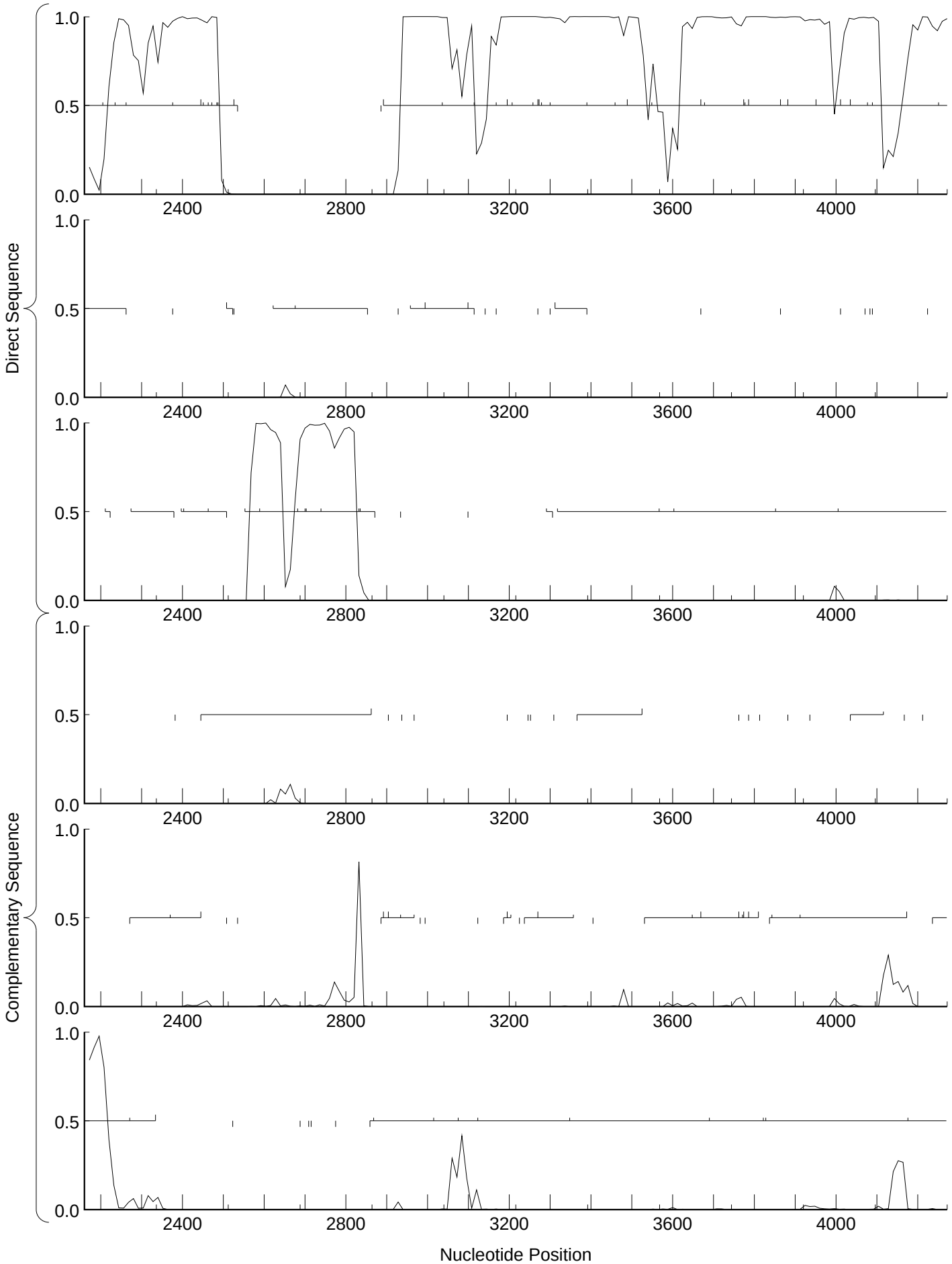

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

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

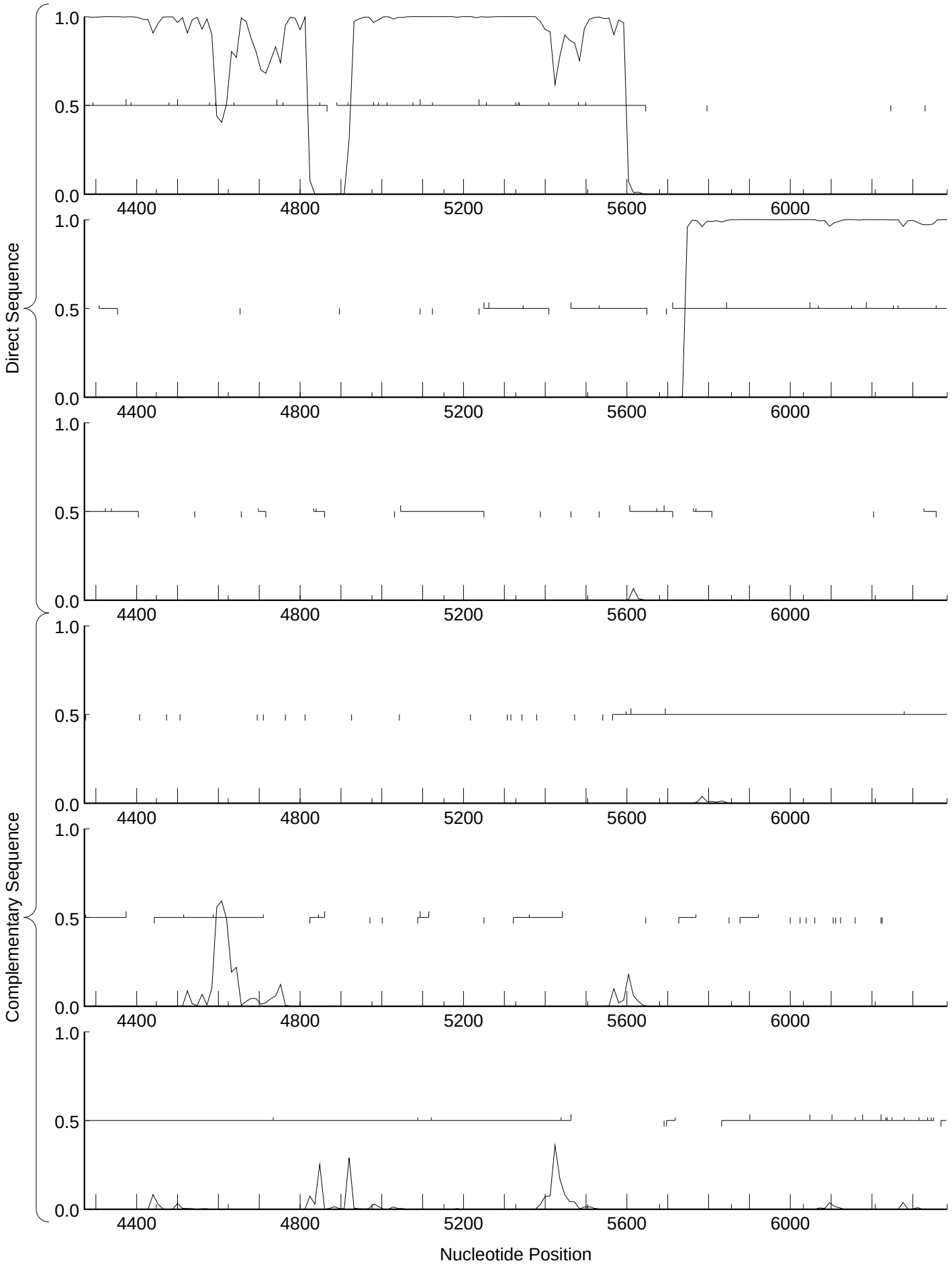
mplete sequence, 39792 bp including 12-base 3' overhang (GAGTCGCCGGCA), Cluster AS1, Order 4, Window 96, Step 12, 2/20



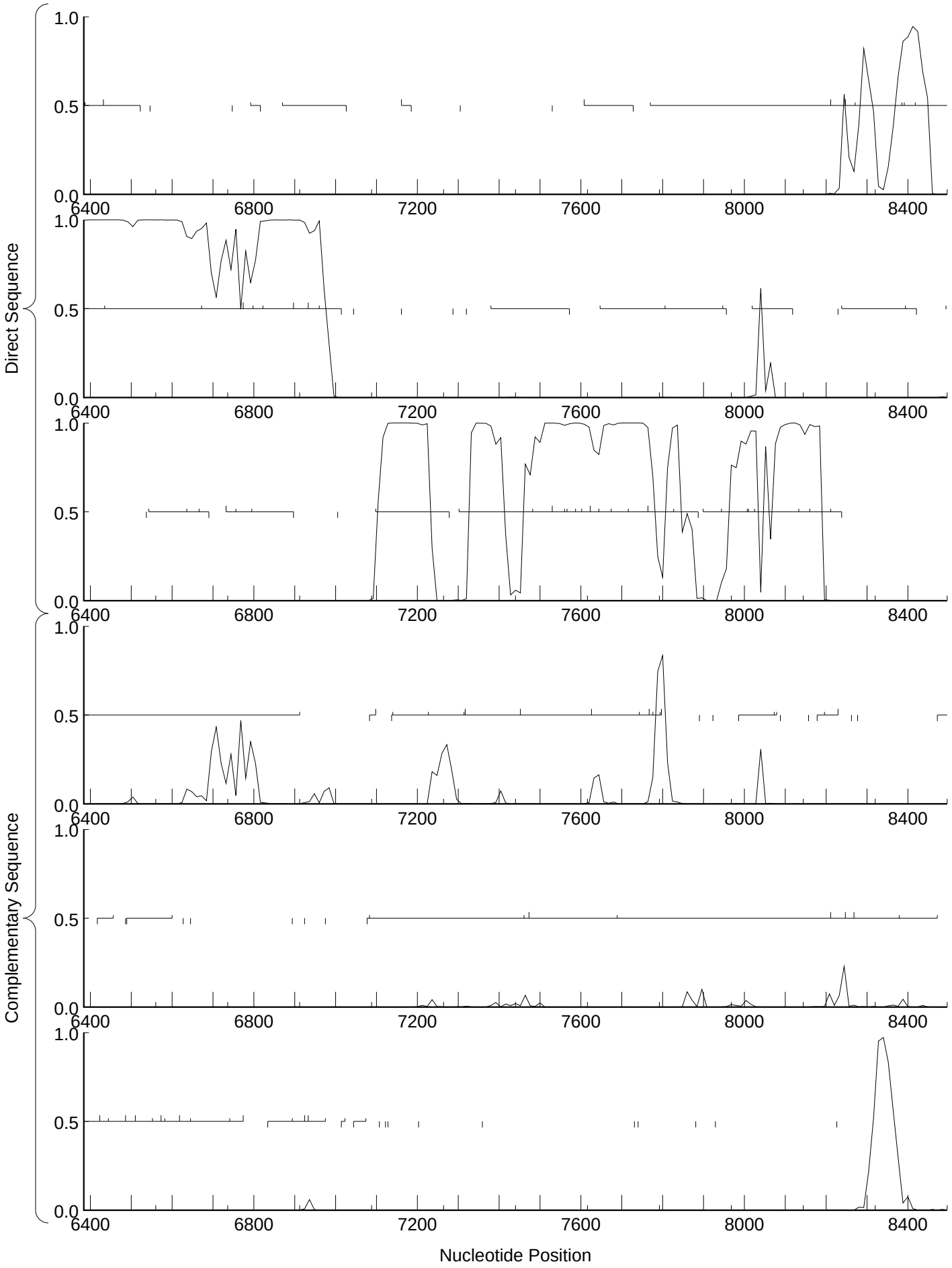
mplete sequence, 39792 bp including 12-base 3’ overhang (GAGTCGCCGGCA), Cluster AS1, Order 4, Window 96, Step 12, 3/20



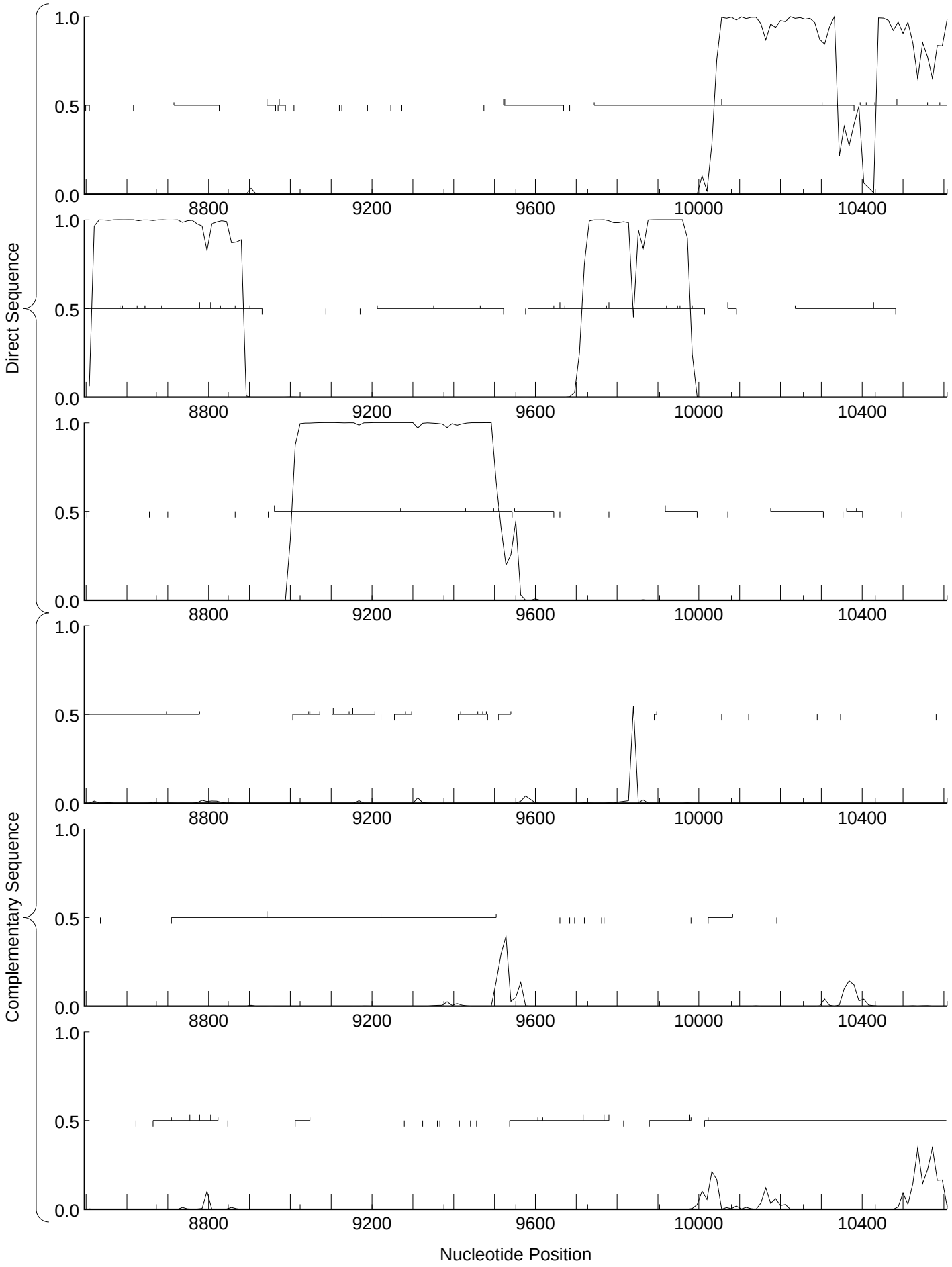
mplete sequence, 39792 bp including 12-base 3’ overhang (GAGTCGCCGGCA), Cluster AS1, Order 4, Window 96, Step 12, 4/20



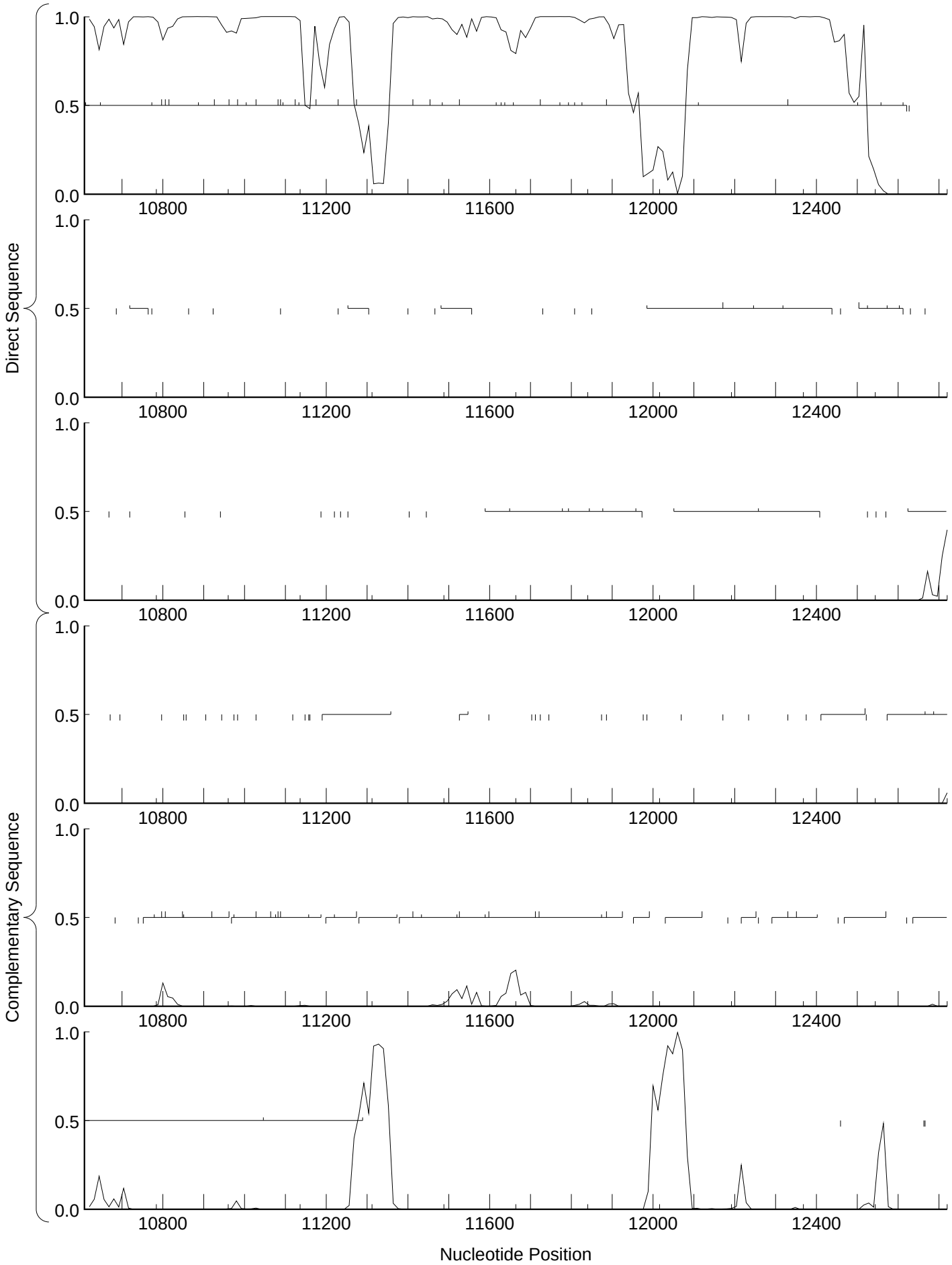
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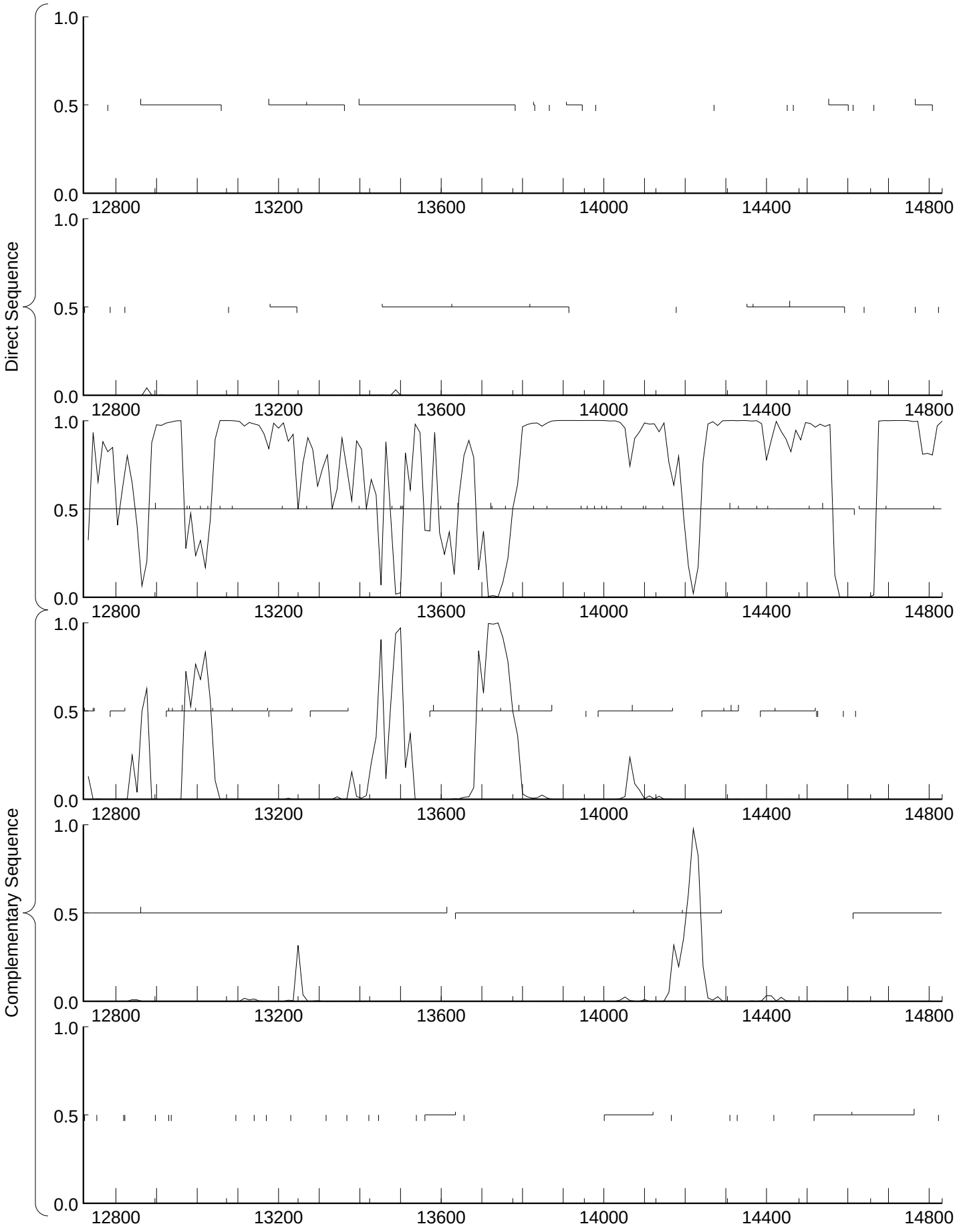
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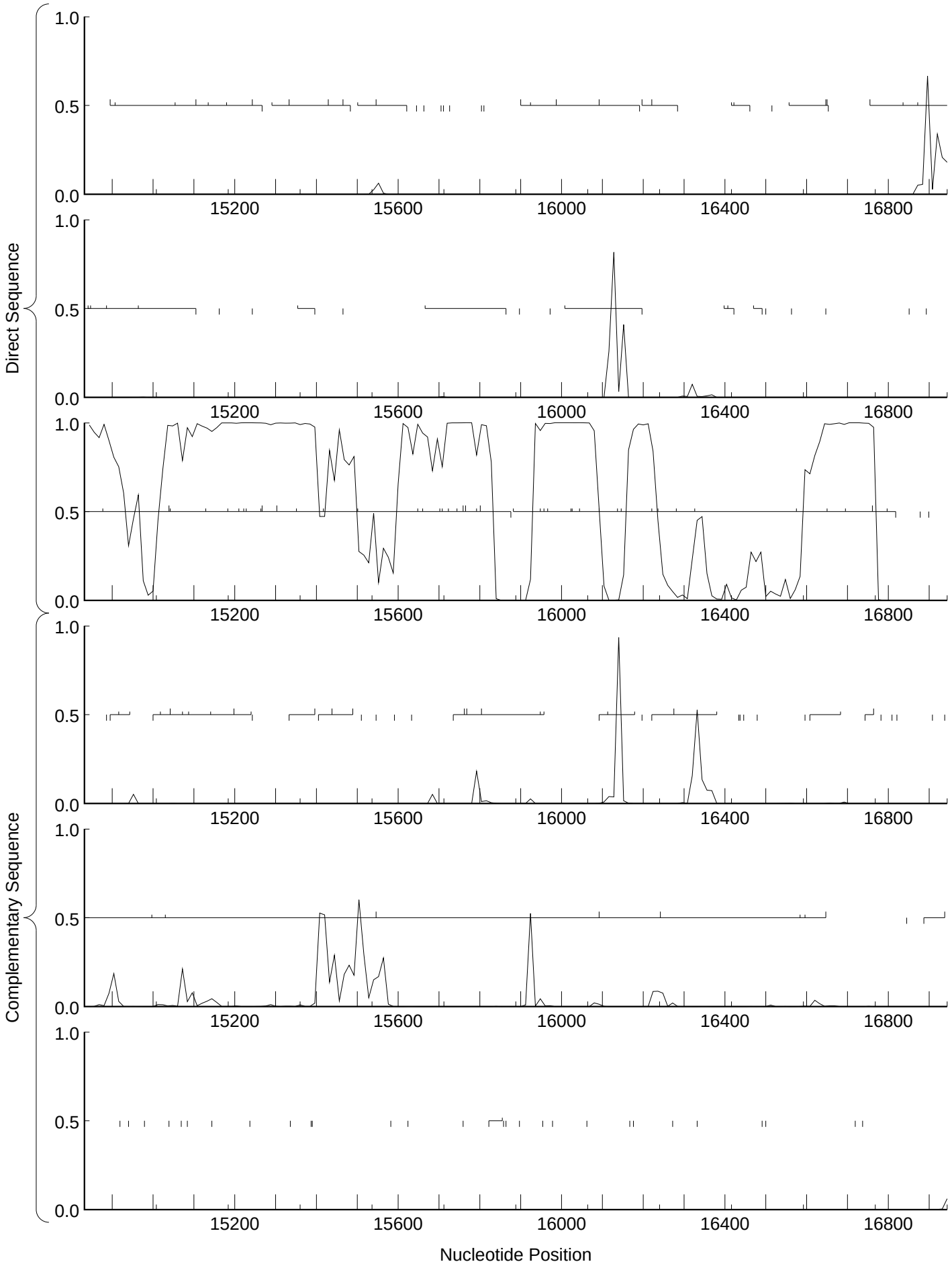
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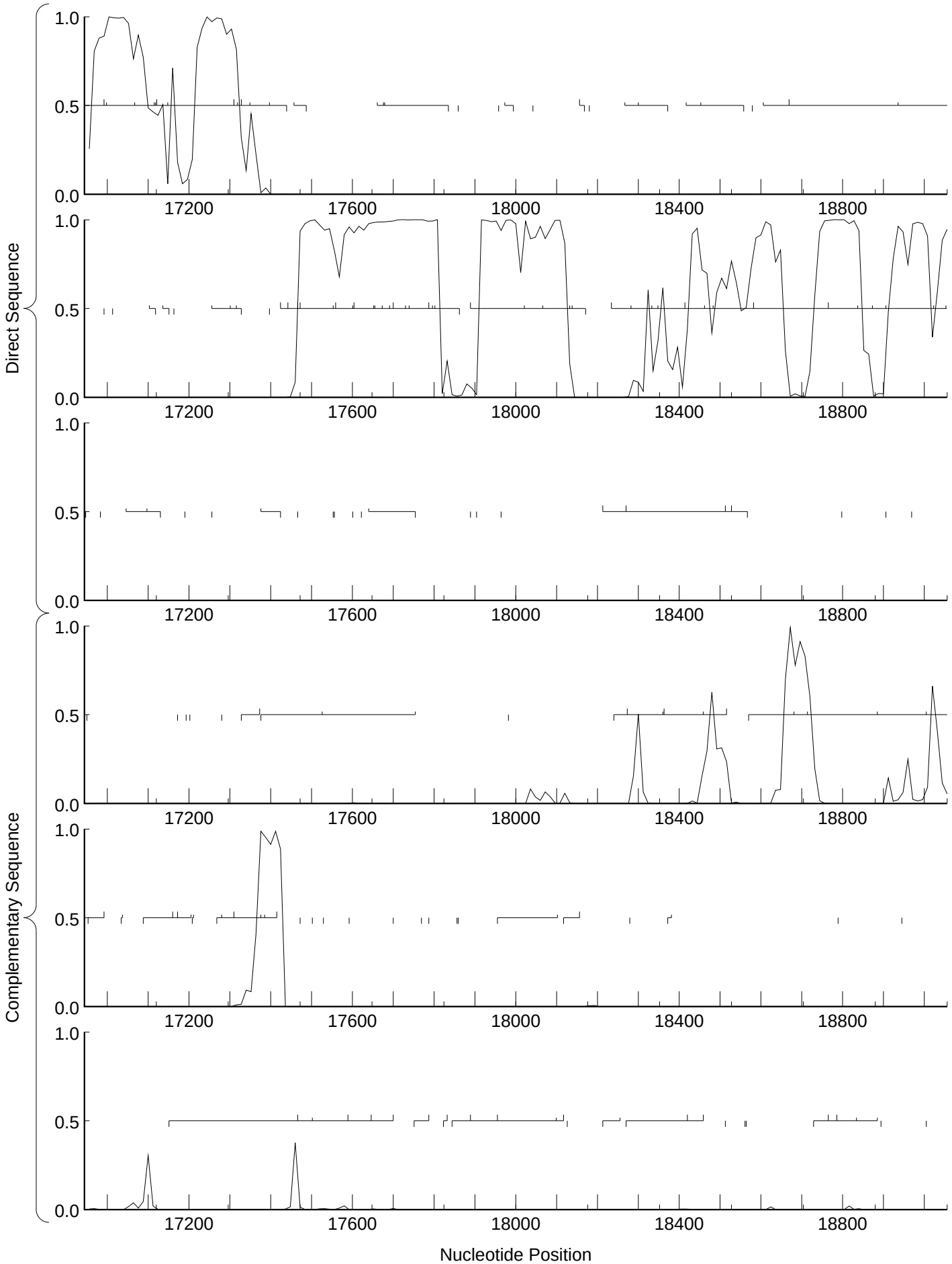


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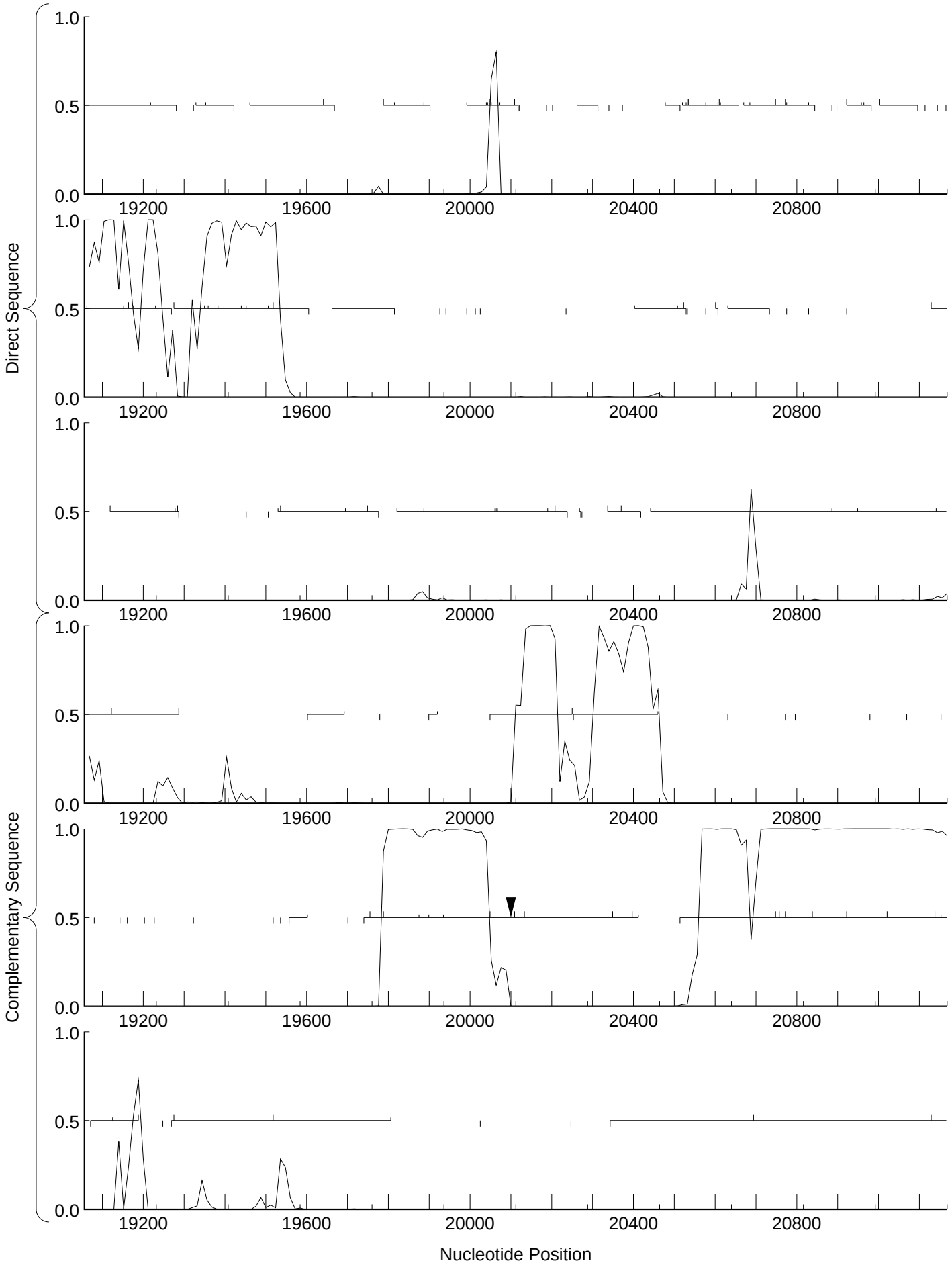


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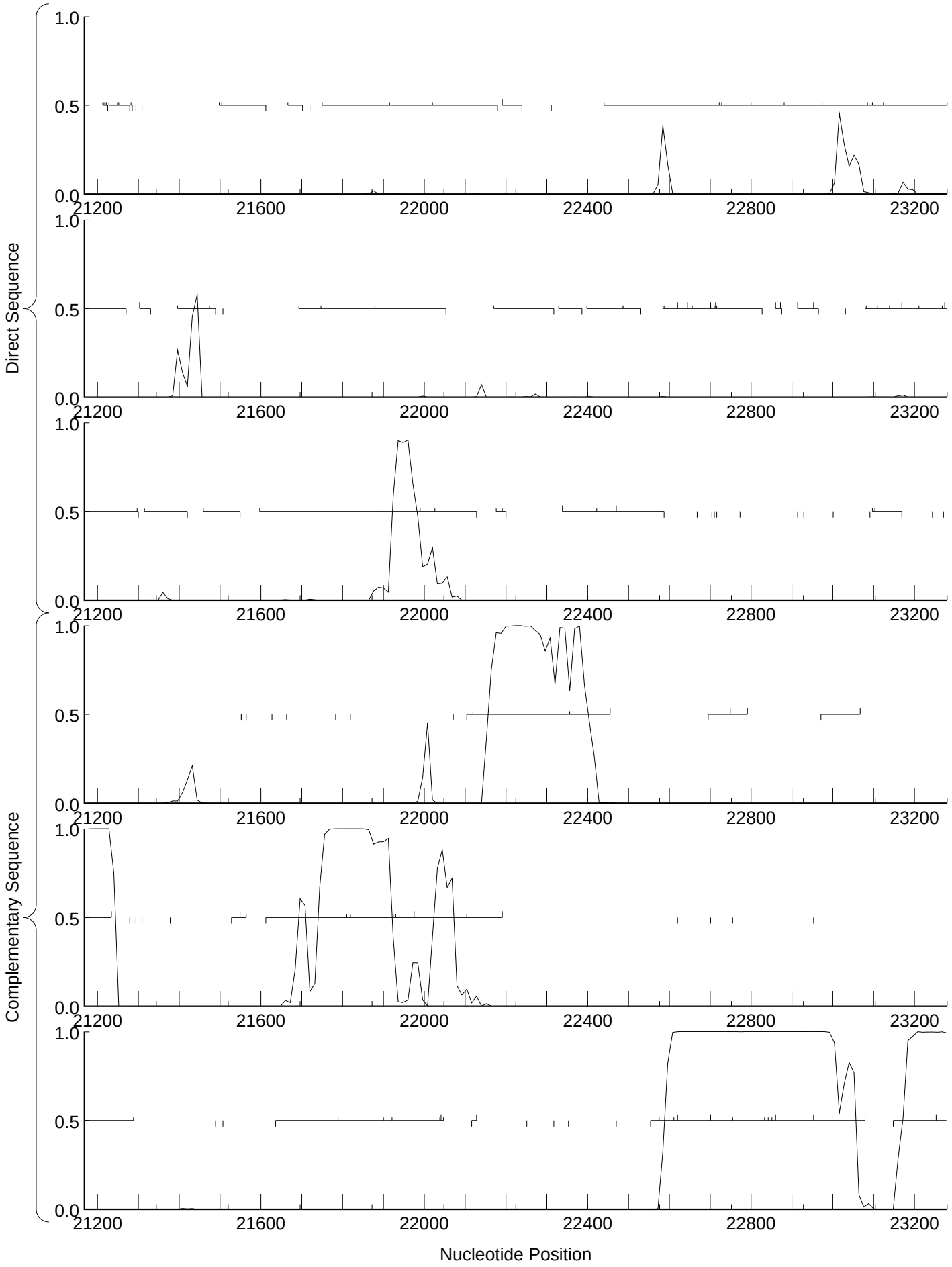



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
17200
17600
18000
18400
18800
Nucleotide Position

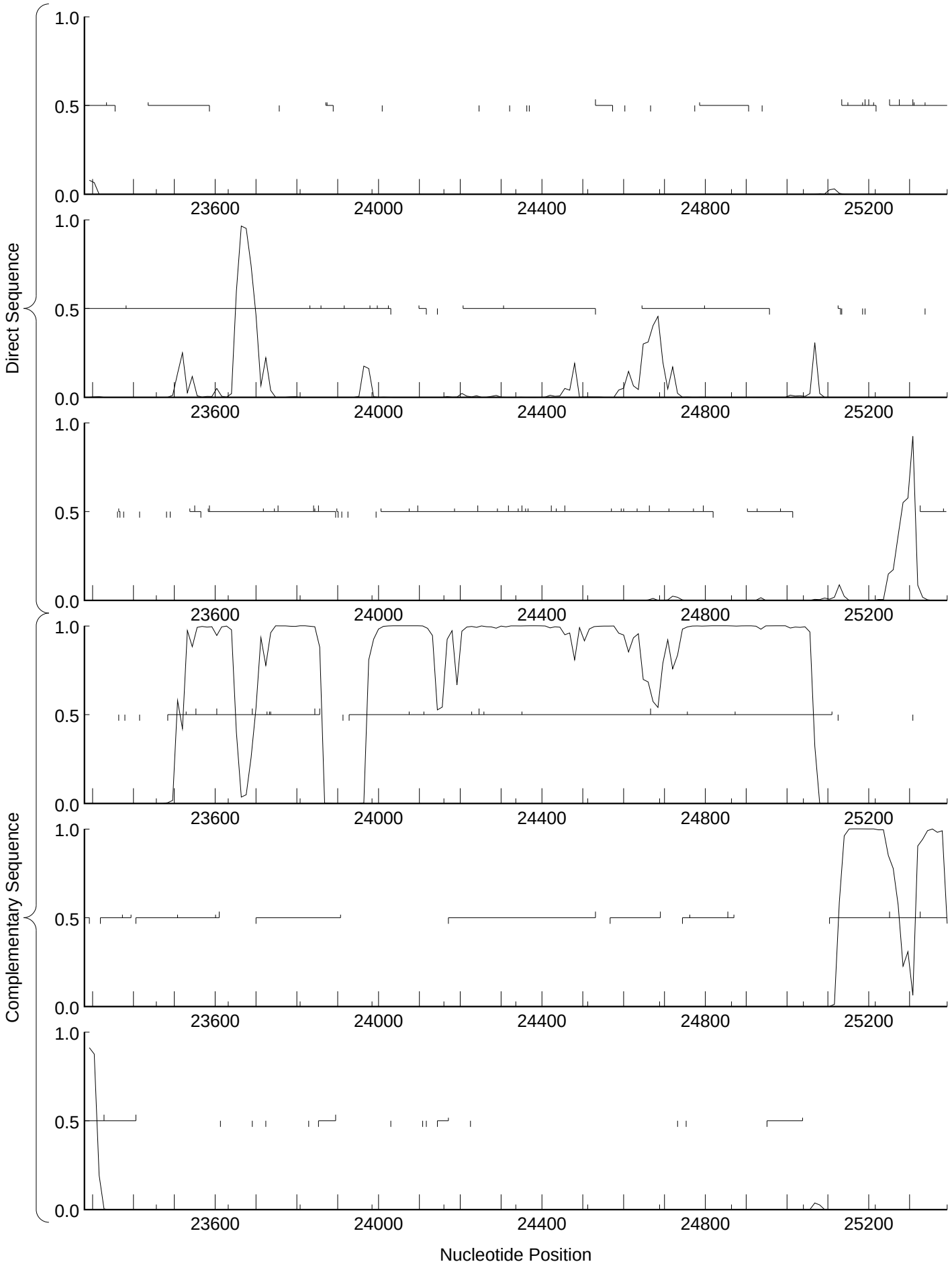
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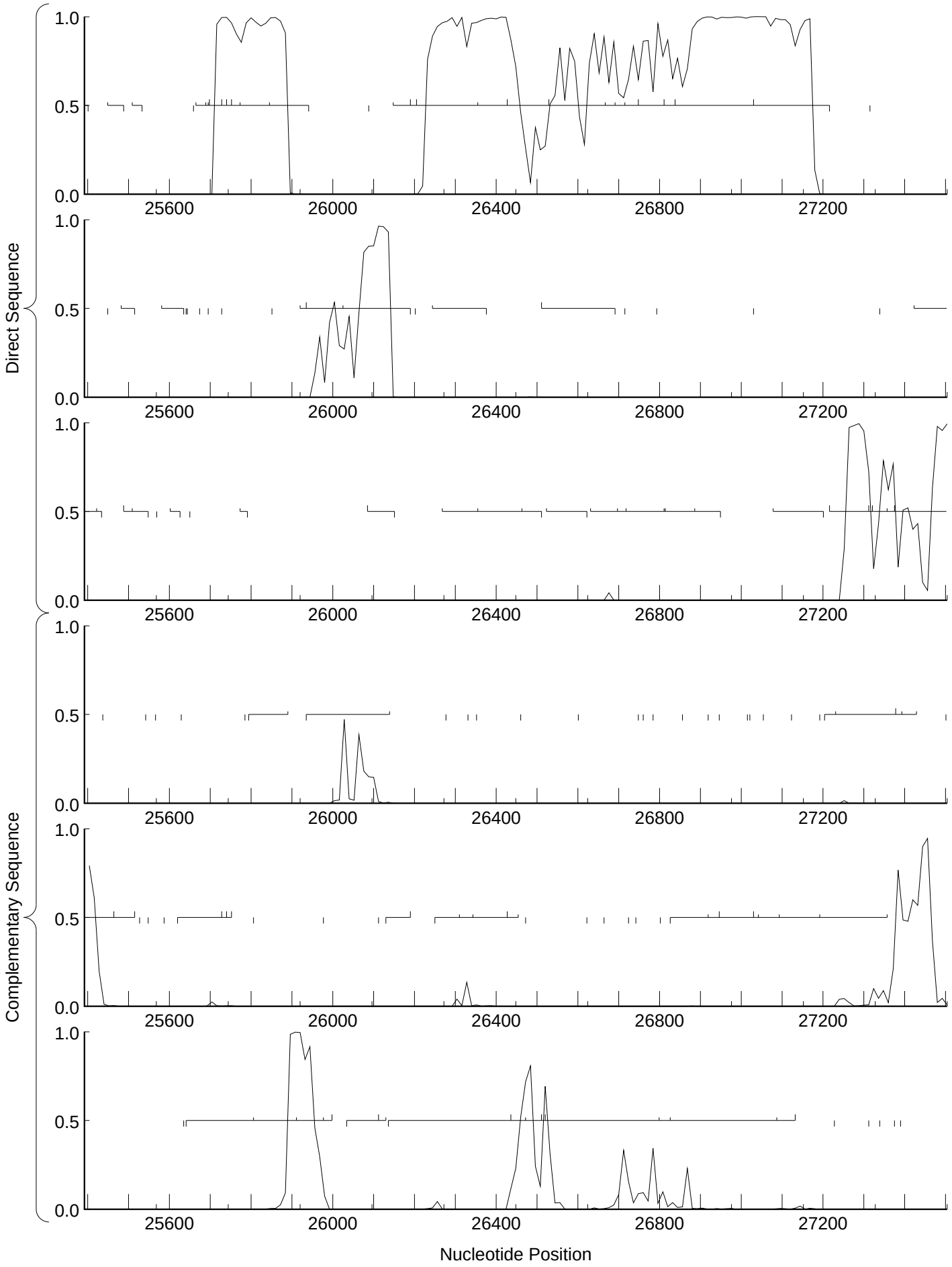


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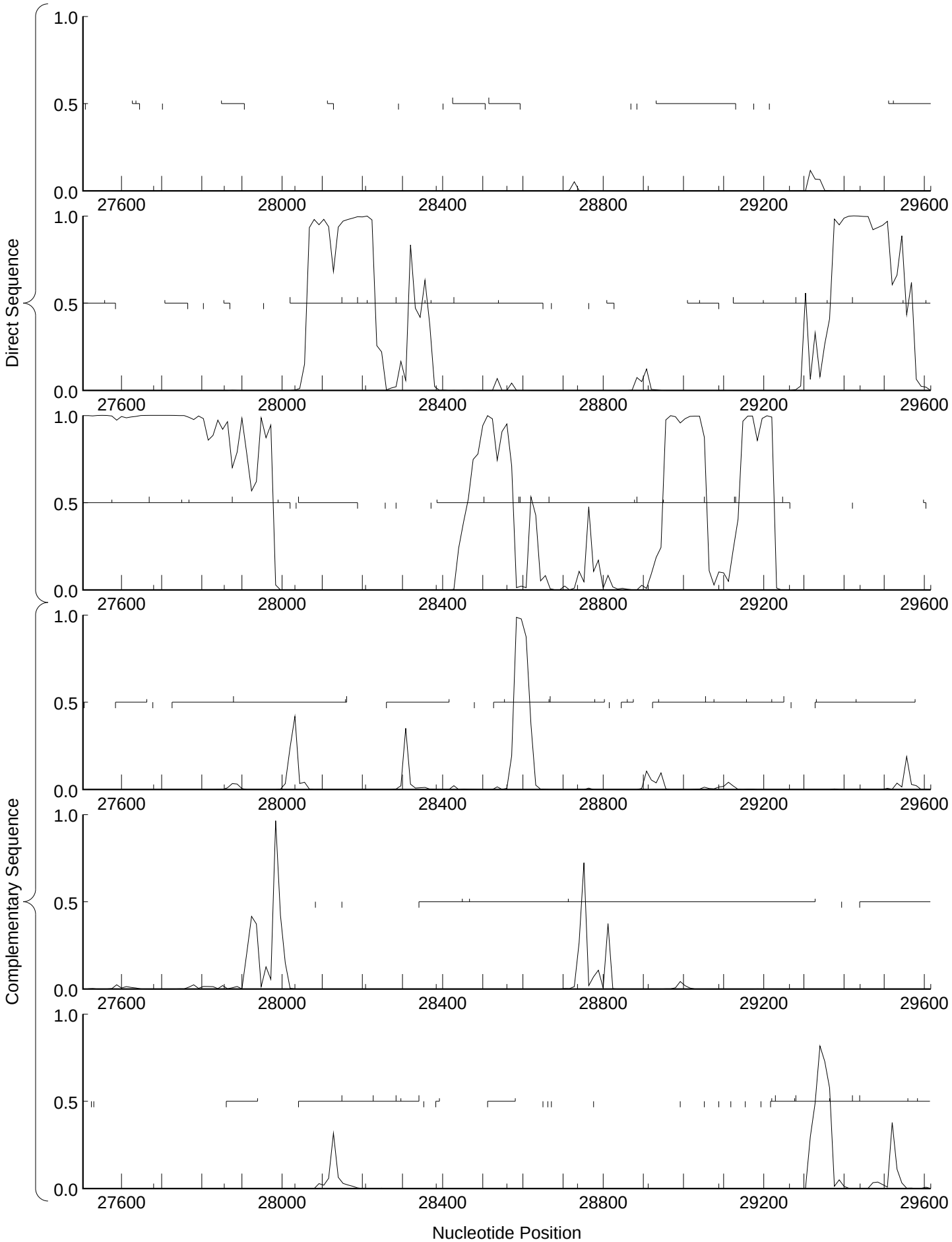


plete sequence, 39792 bp including 12-base 3’ overhang (GAGTCGCCGGCA), Cluster AS1, Order 4, Window 96, Step 12, 13/20

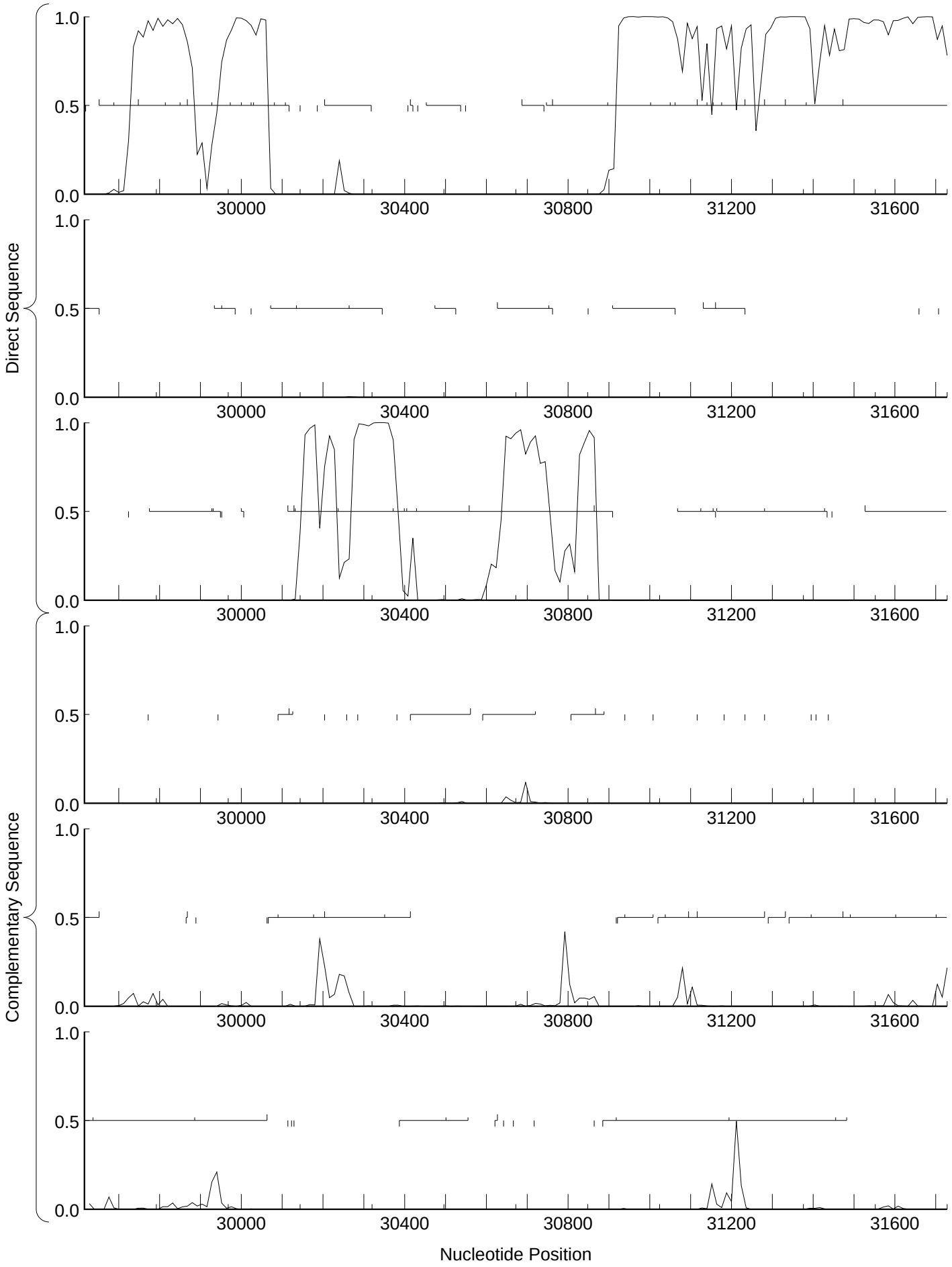



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
25600
26000
26400
26800
27200
Nucleotide Position

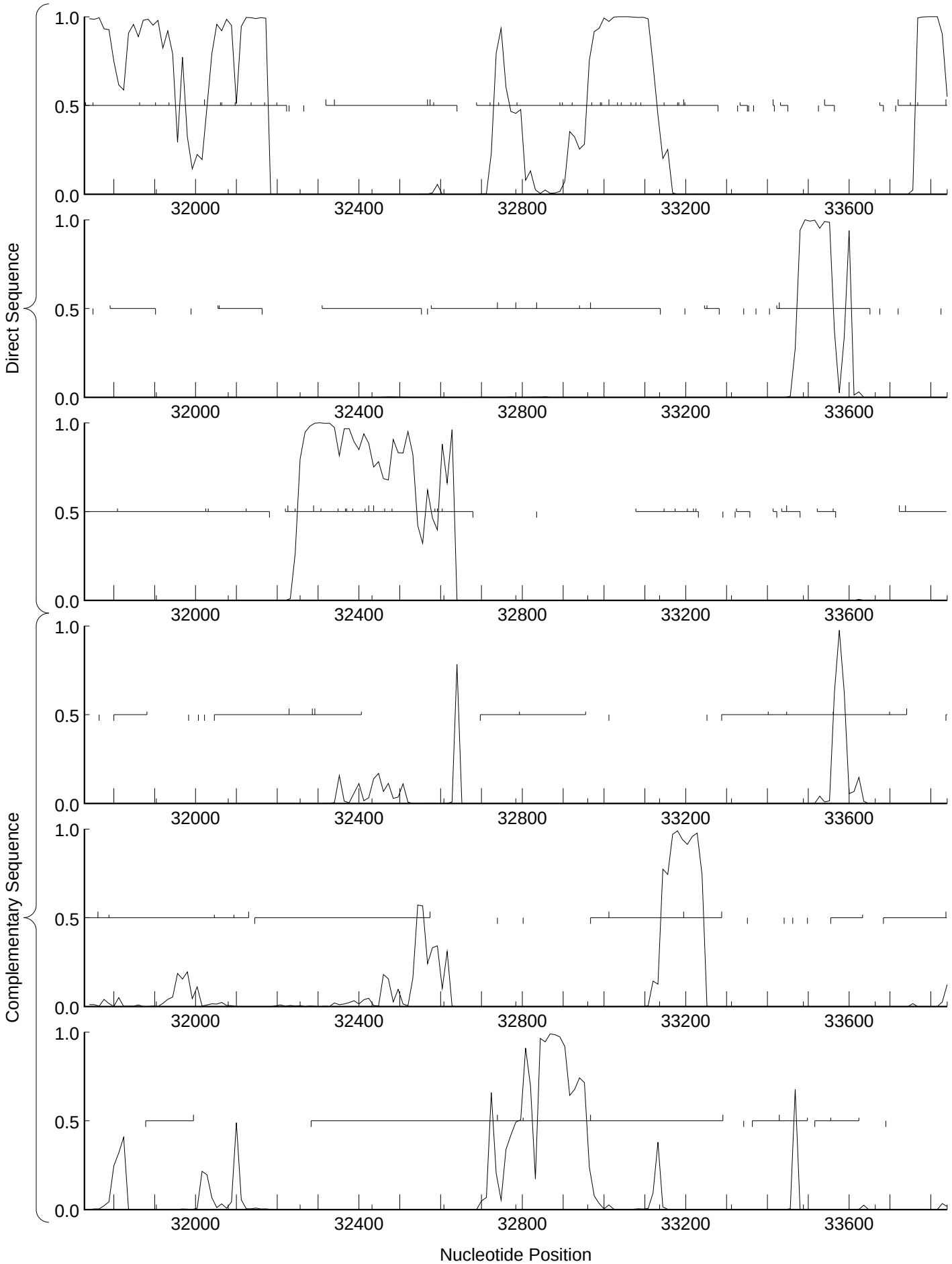
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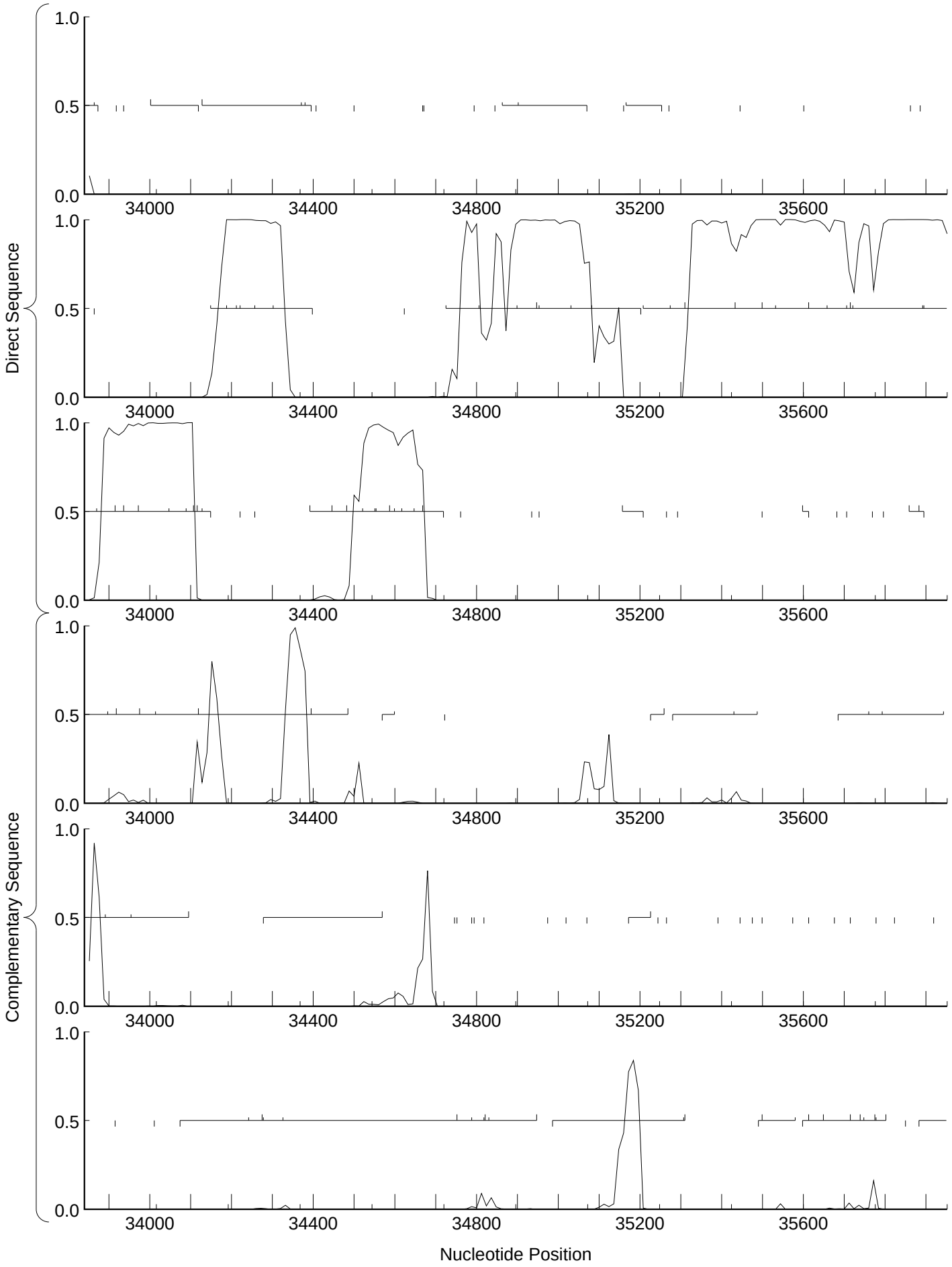
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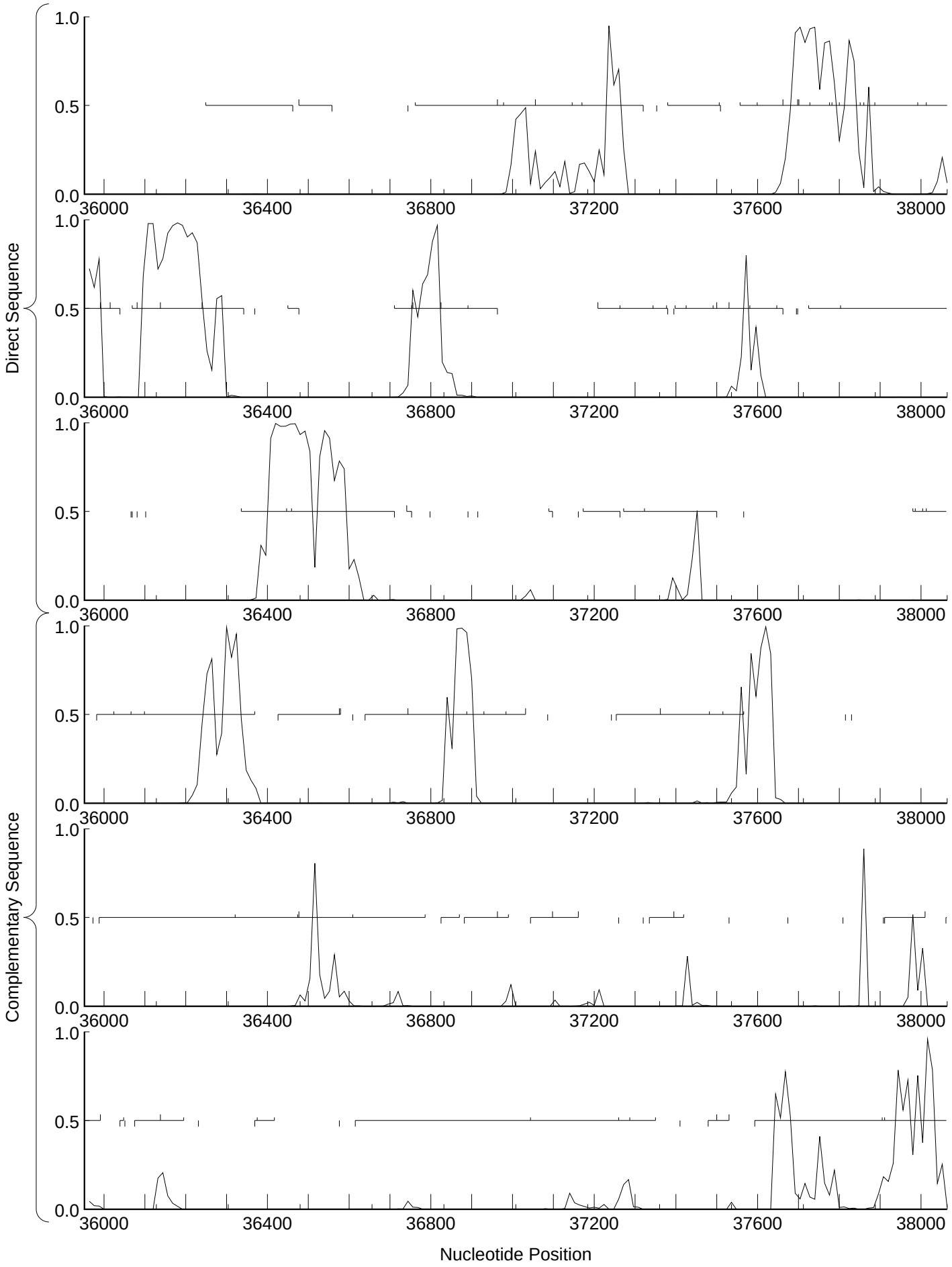
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plete sequence, 39792 bp including 12-base 3’ overhang (GAGTCGCCGGCA), Cluster AS1, Order 4, Window 96, Step 12, 18/20



plete sequence, 39792 bp including 12-base 3’ overhang (GAGTCGCCGGCA), Cluster AS1, Order 4, Window 96, Step 12, 19/20



plete sequence, 39792 bp including 12-base 3’ overhang (GAGTCGCCGGCA), Cluster AS1, Order 4, Window 96, Step 12, 20/20

