GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Gordonia phage Feastonyeet complete sequence, 47166 bp including 13-base 3' overhang (CGGCG
Analysis Date    :  3/28/24 at 9:21:39
Pages            :  24
Sequence Length  :  47166 bp
GC Content       :   60.71%

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

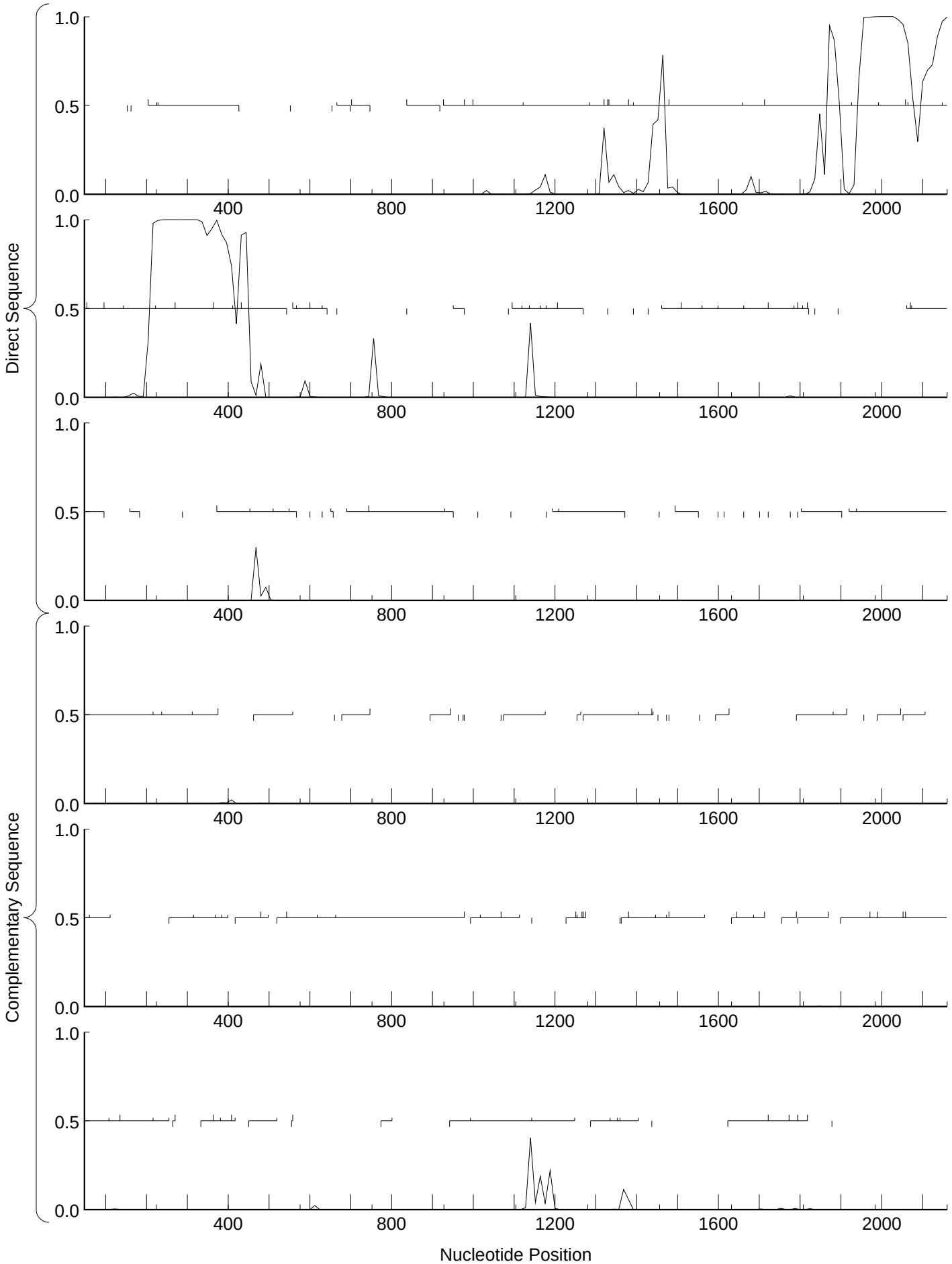

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

Matrix notes & comments

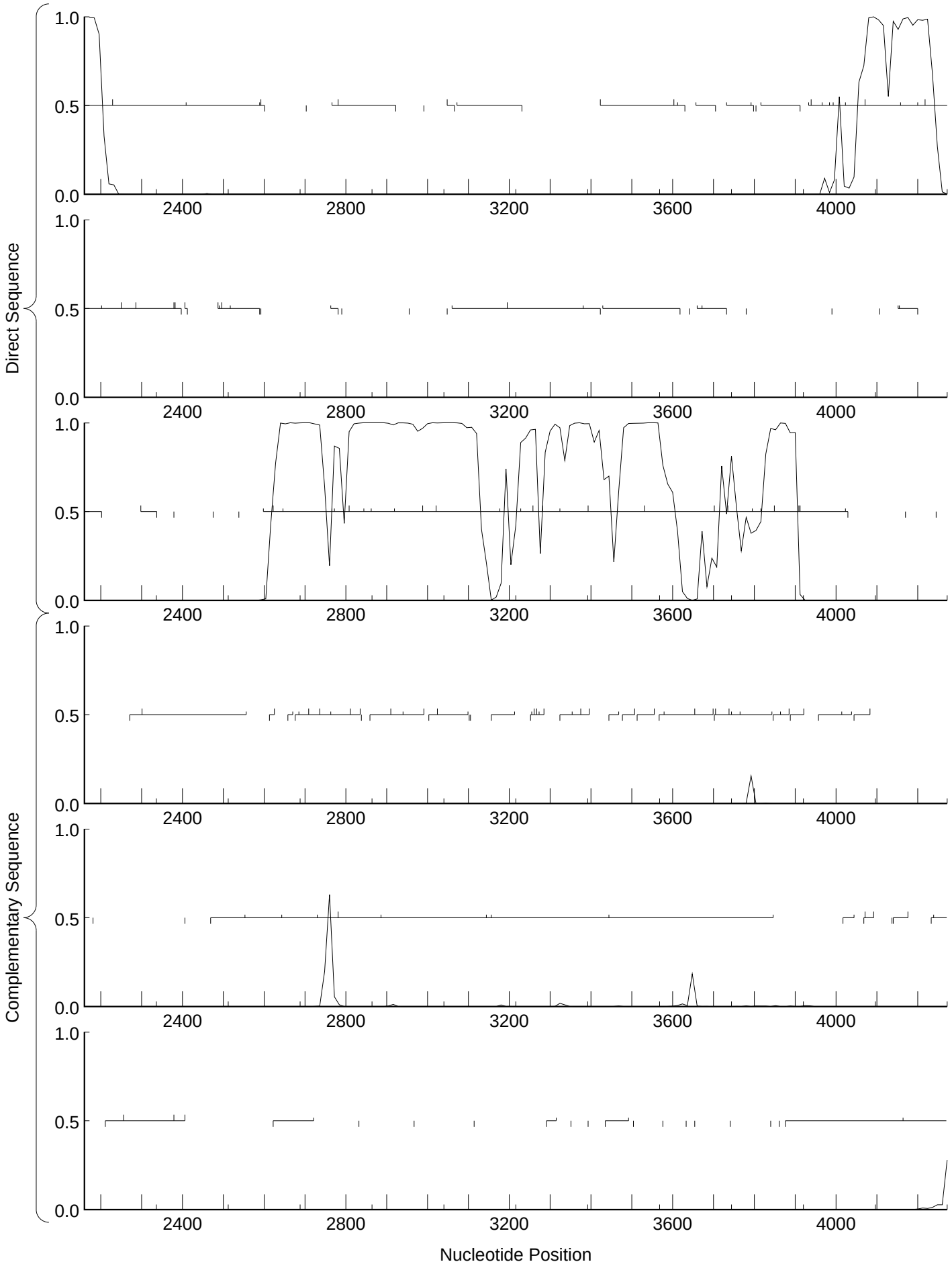
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Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep  8 00:36:14 2018
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For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

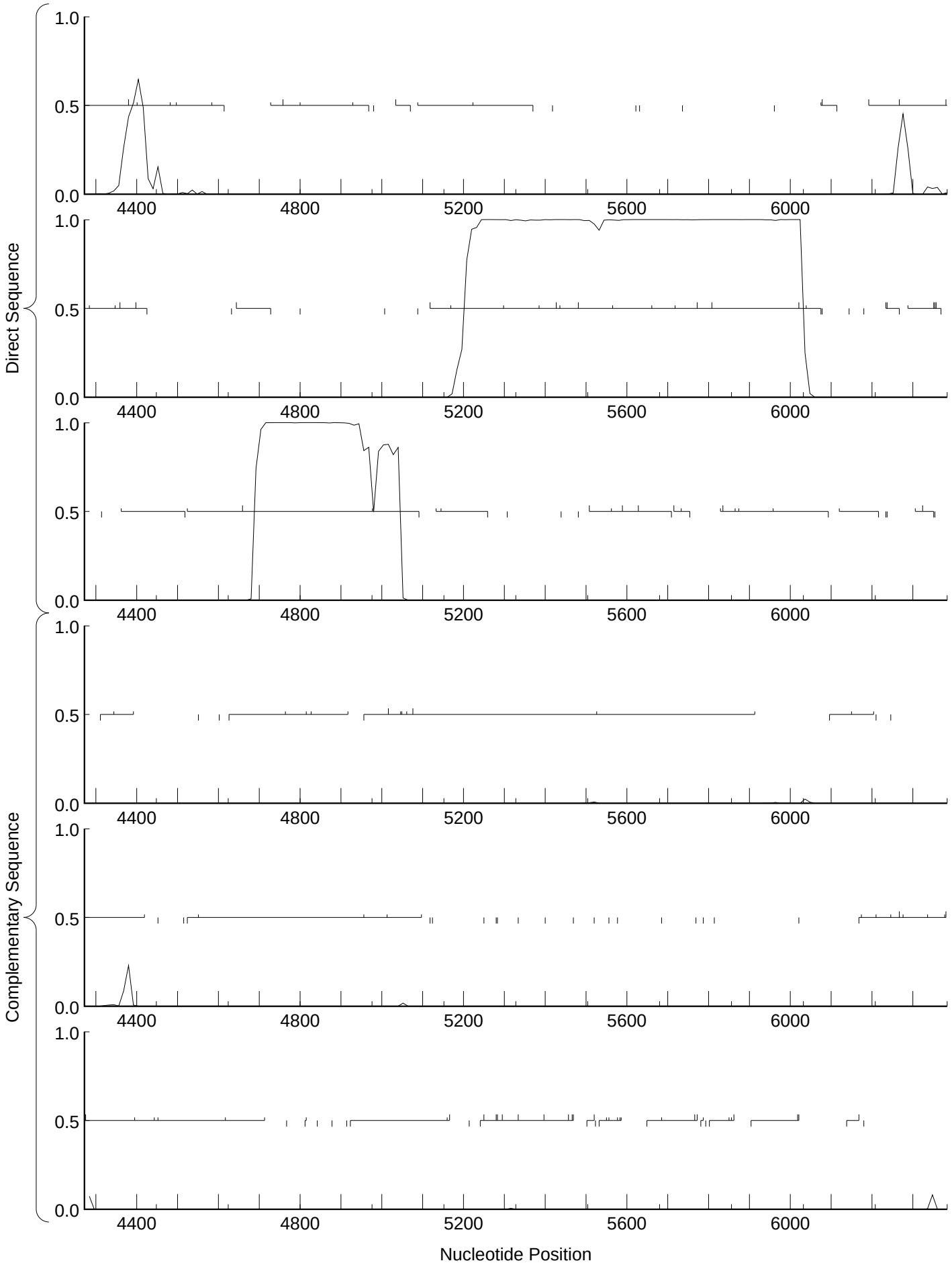
mplete sequence, 47166 bp including 13-base 3’ overhang (CGGCGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 2/24



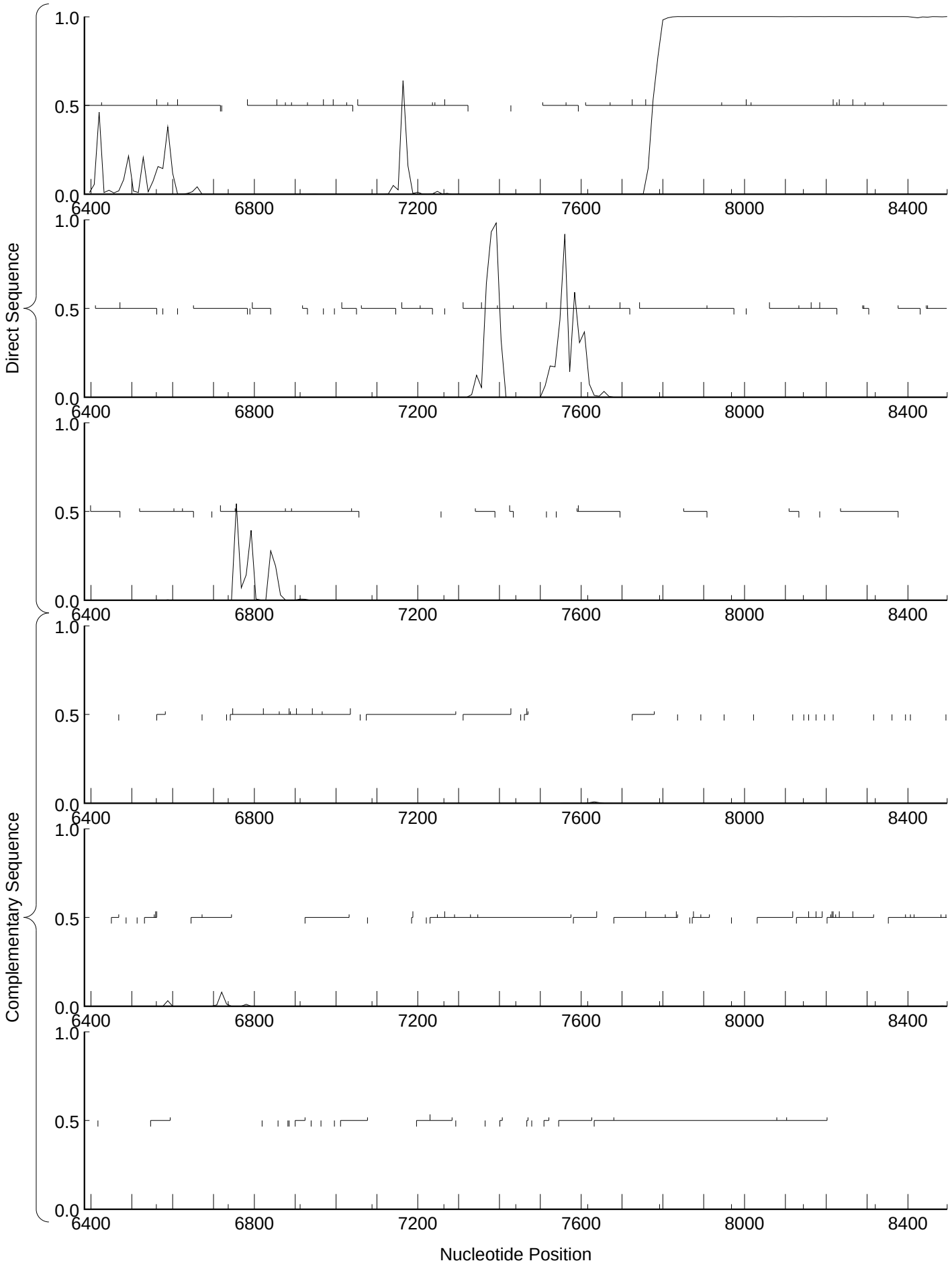
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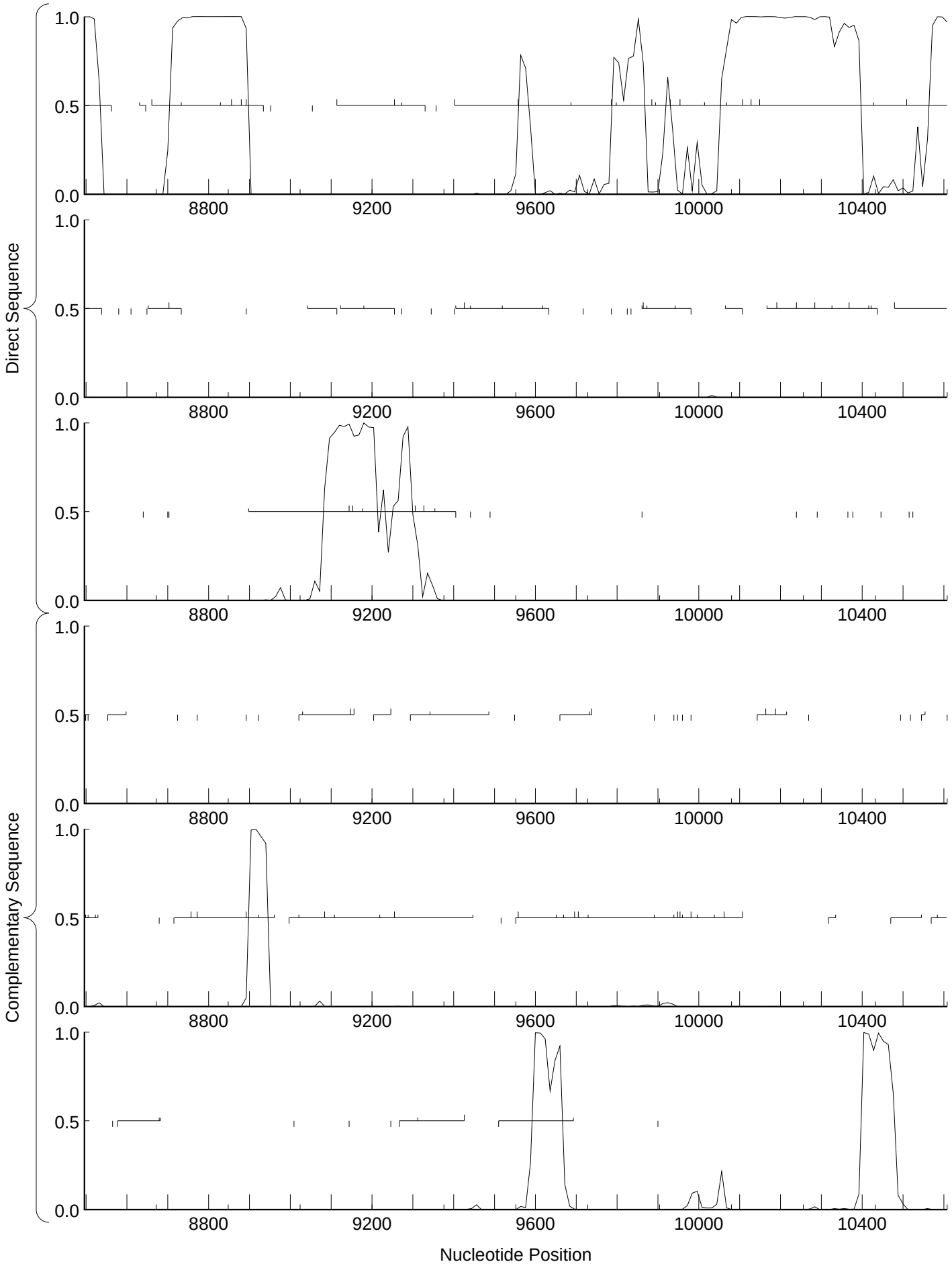
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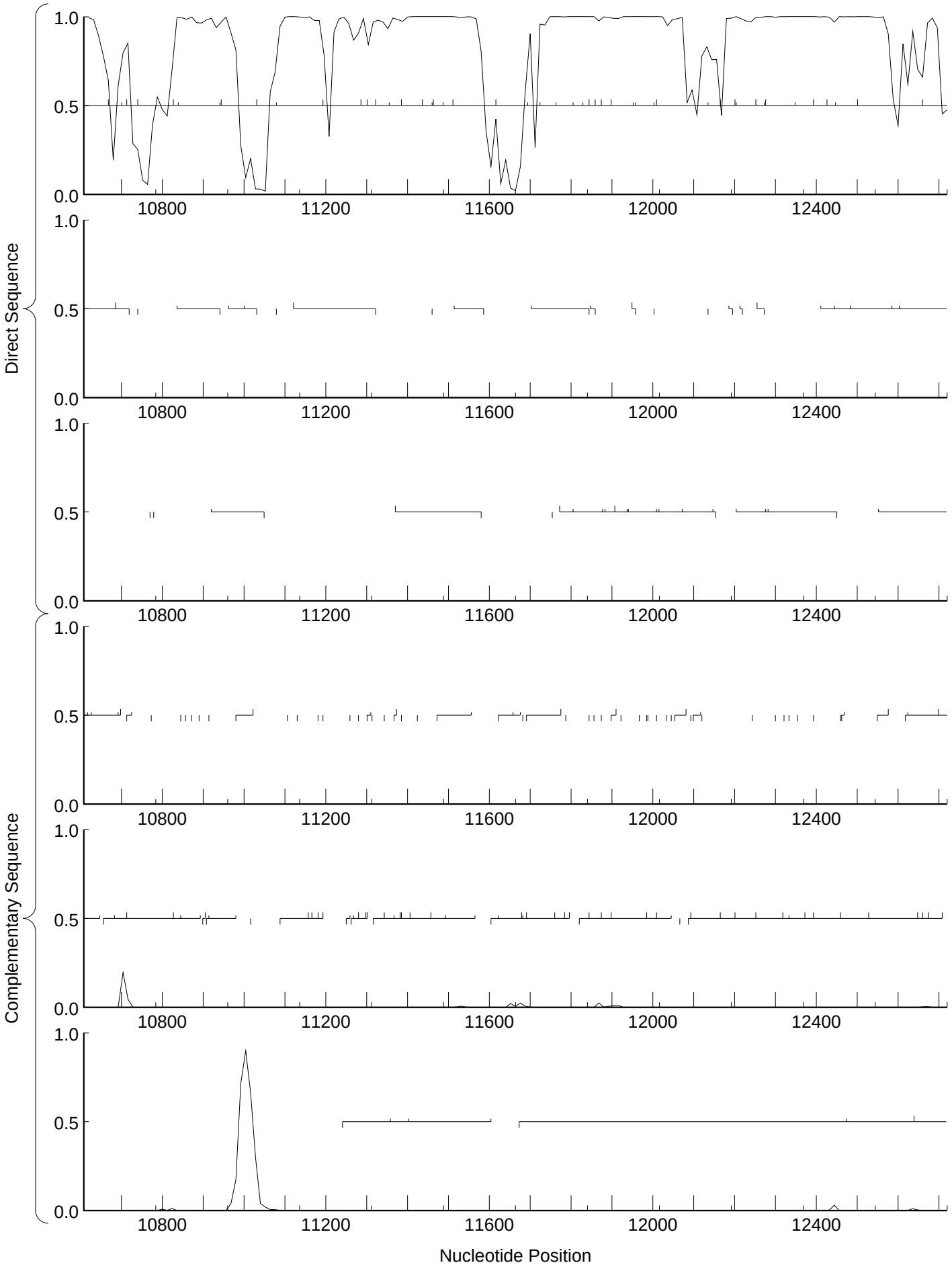
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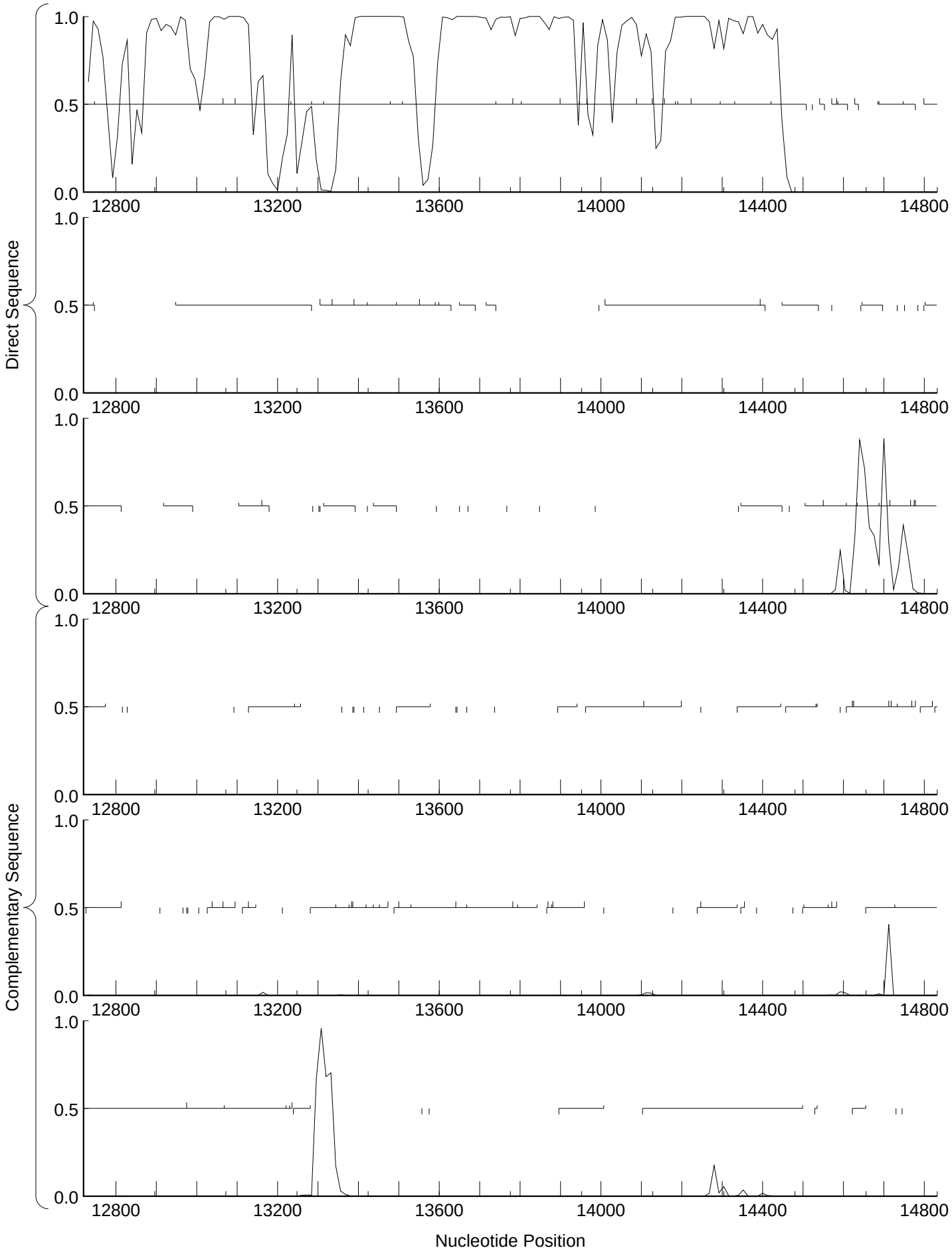
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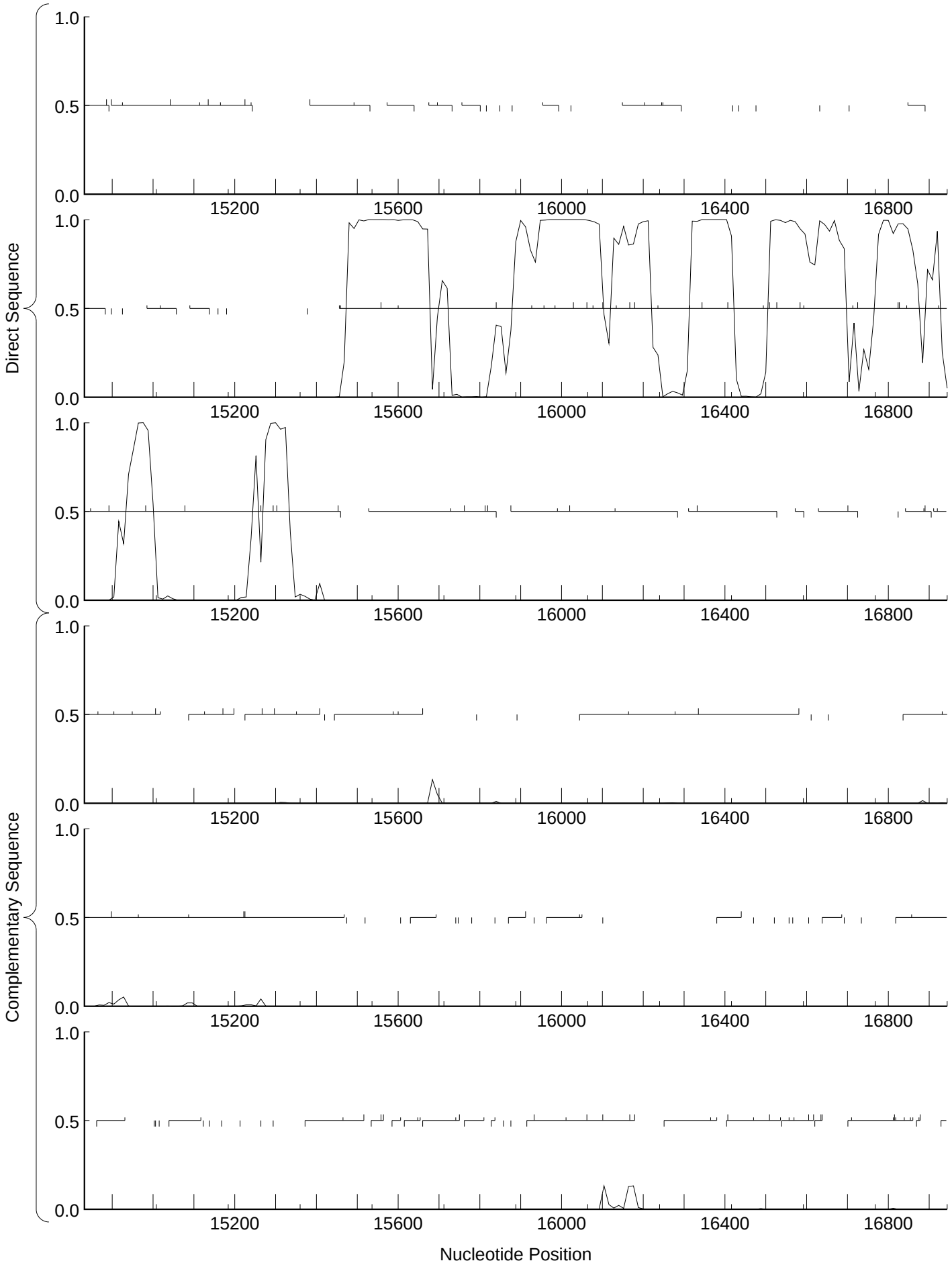
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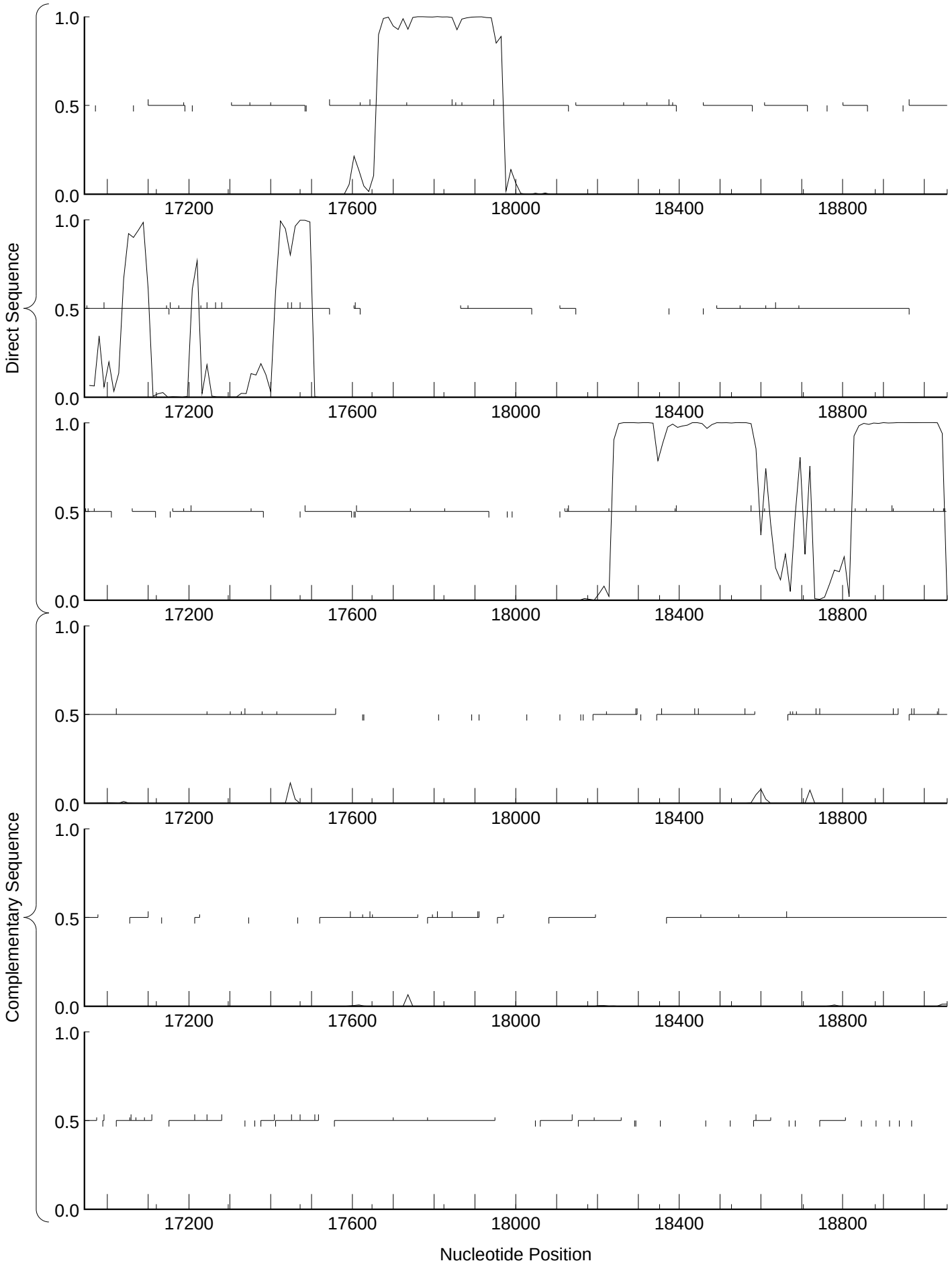
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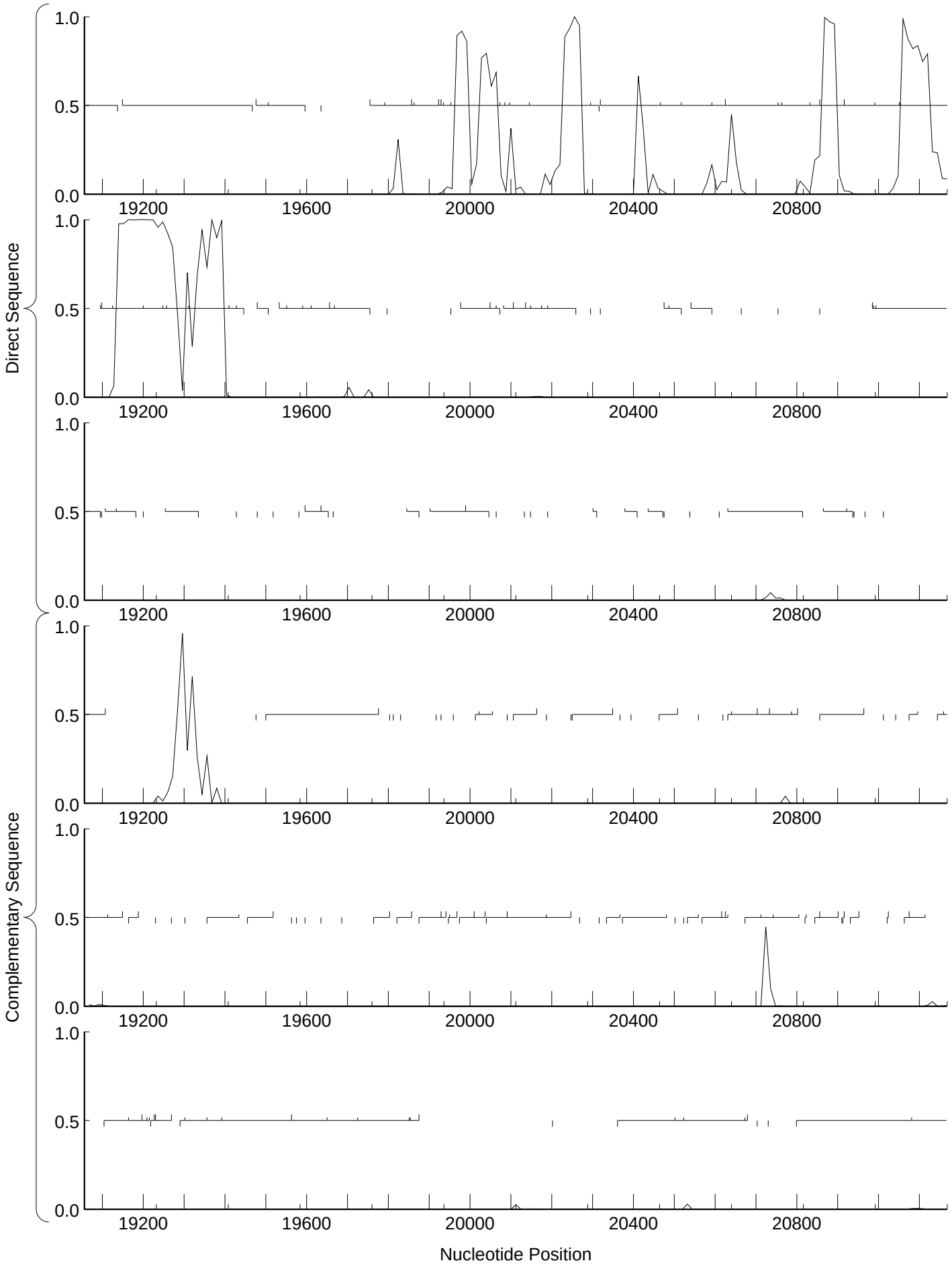
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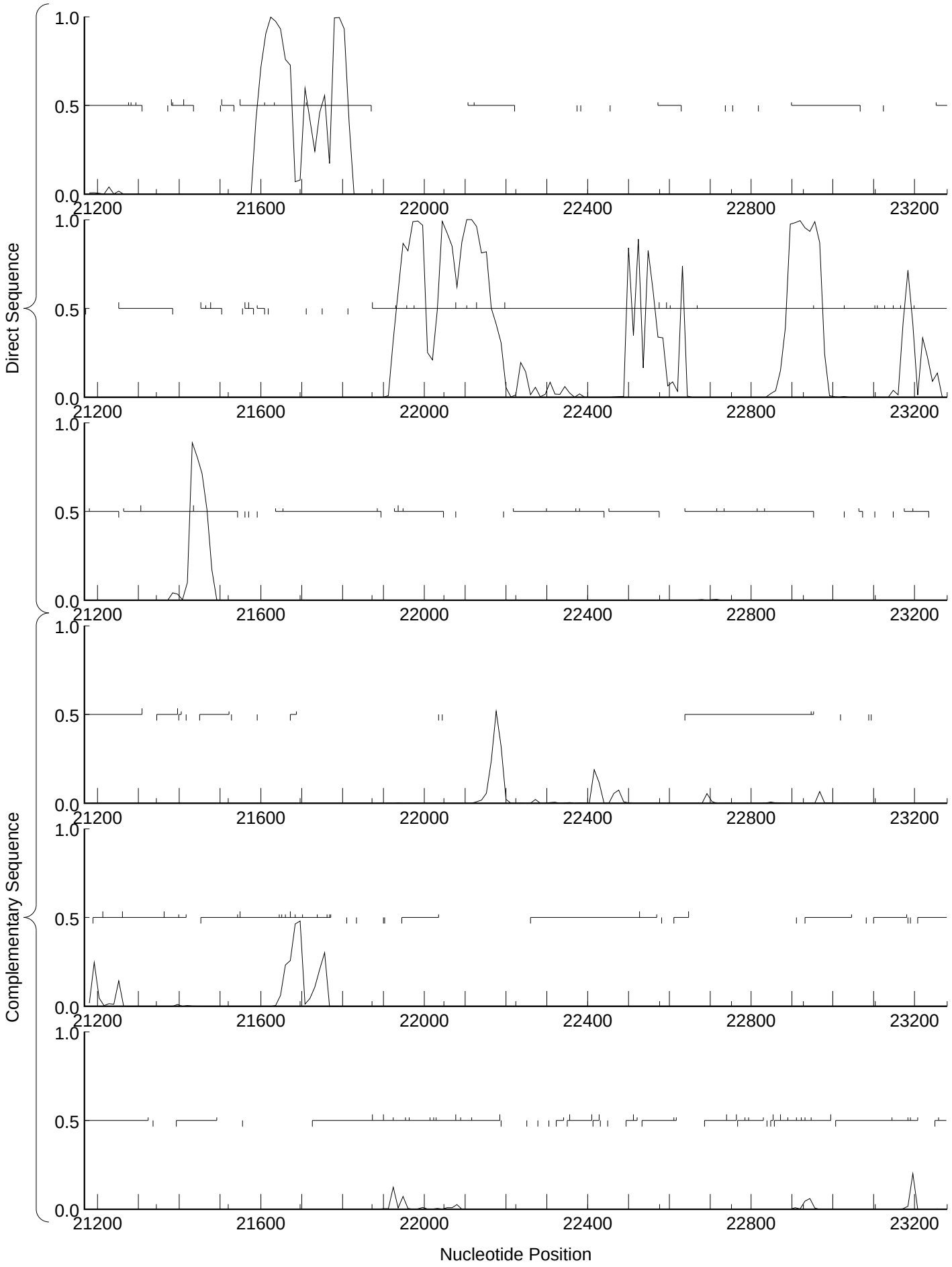
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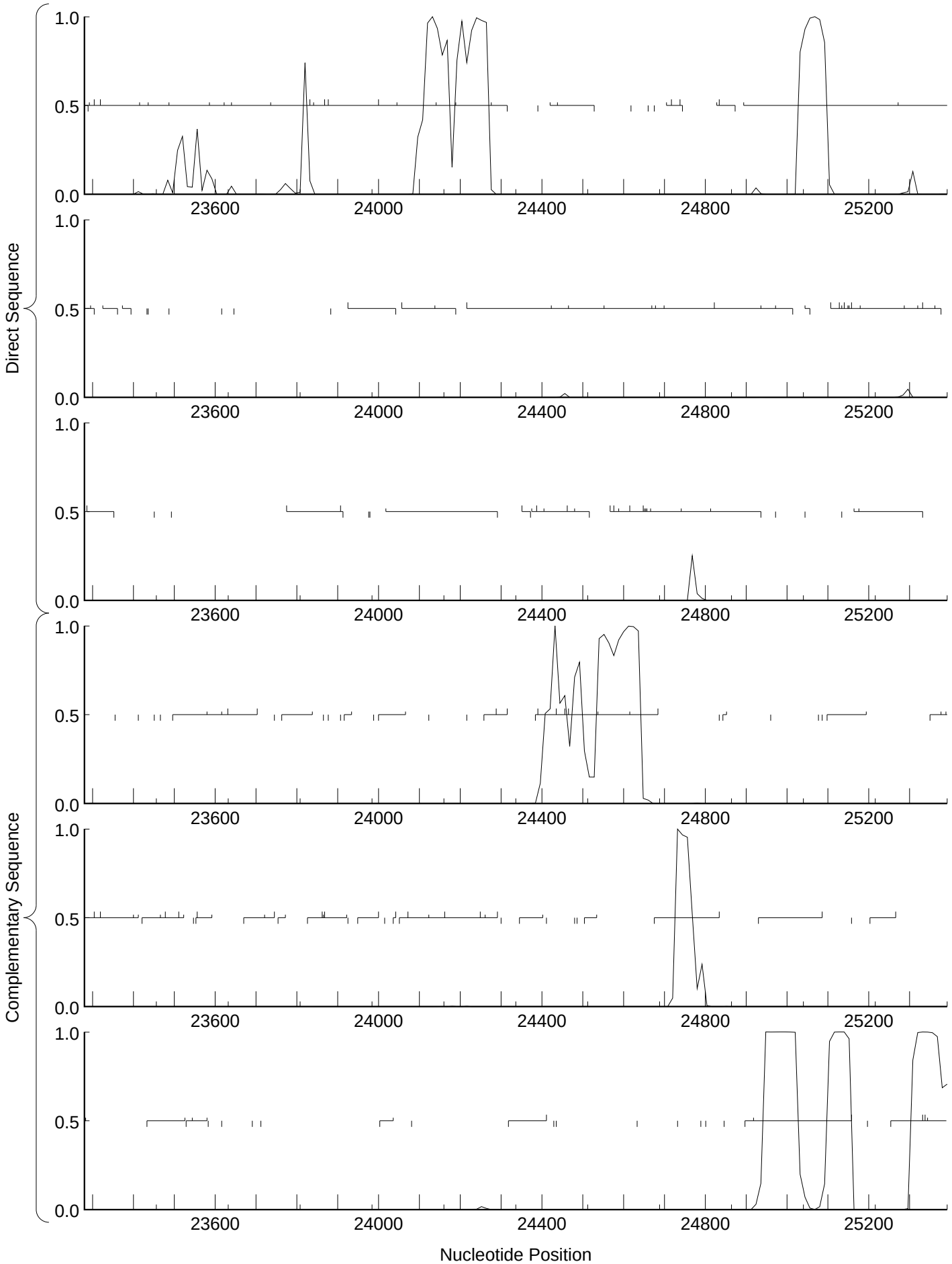
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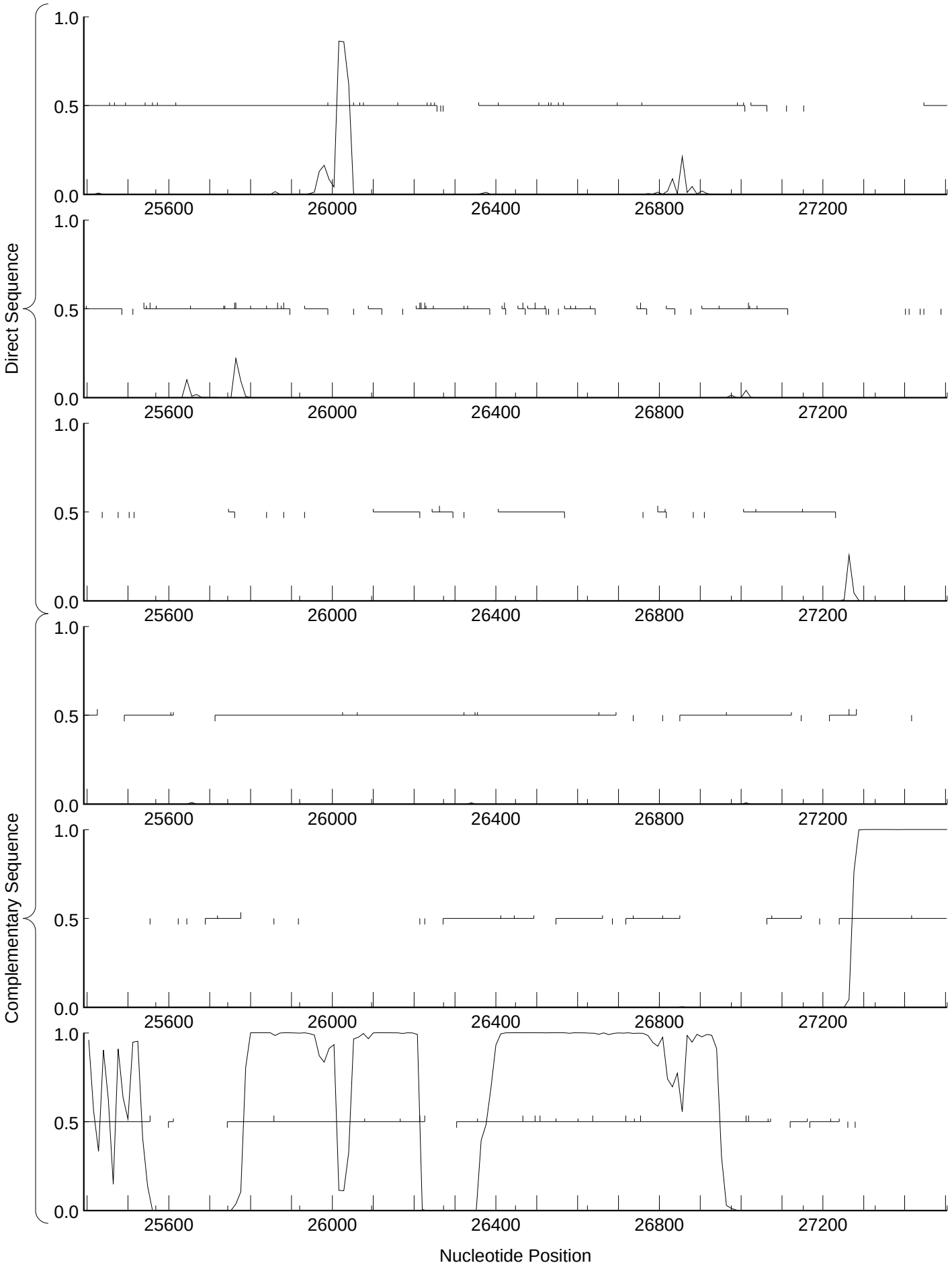
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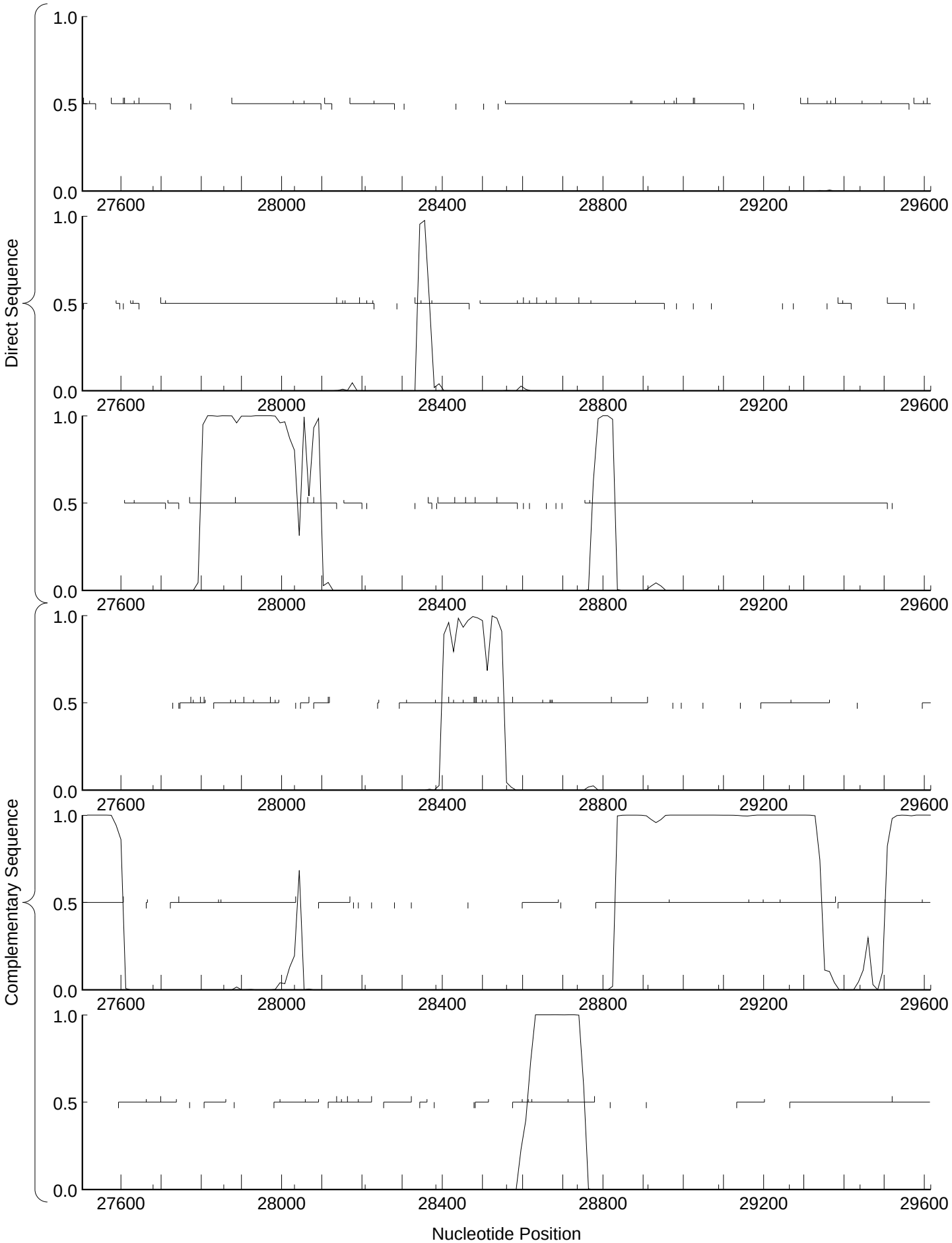
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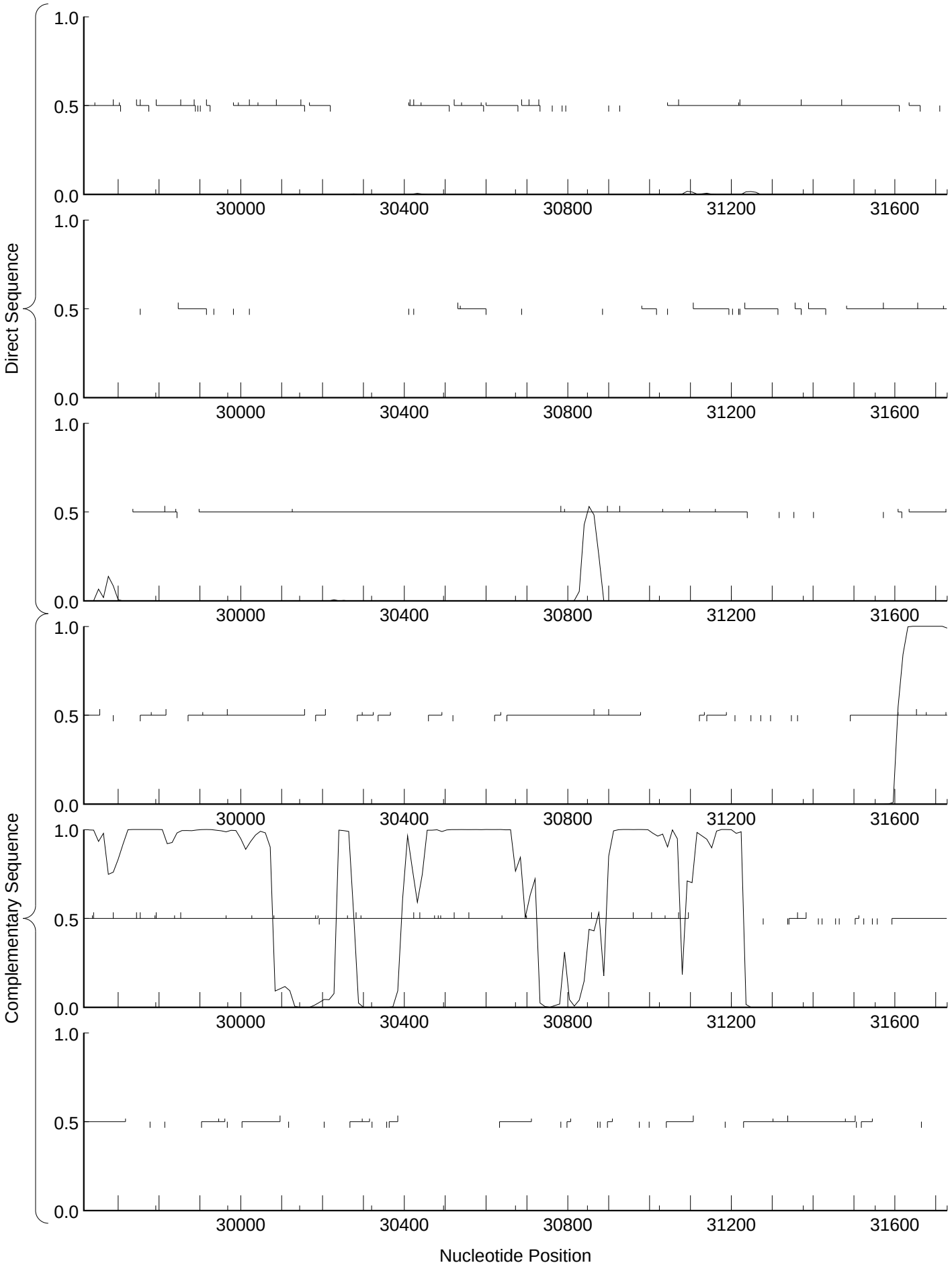
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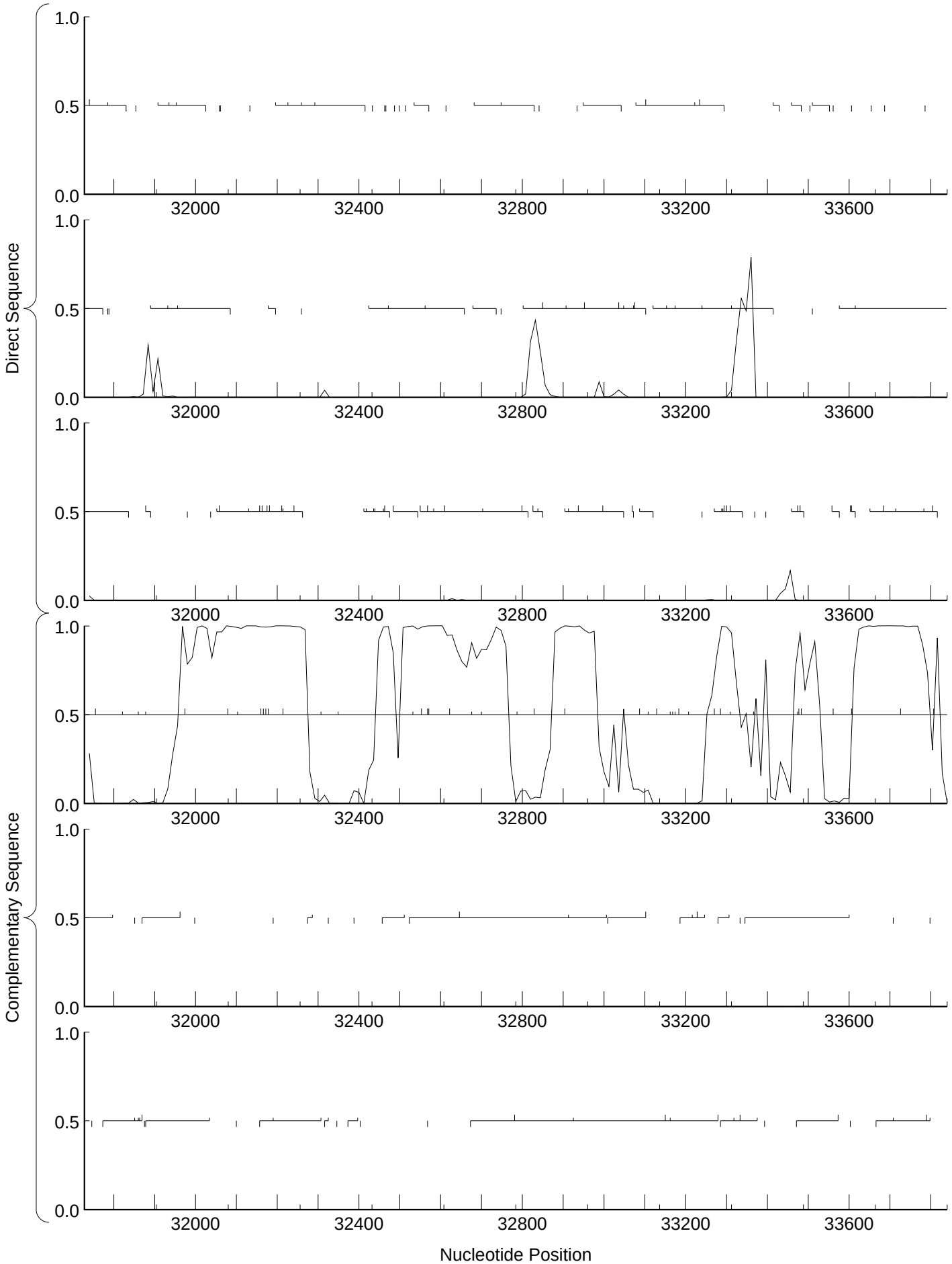
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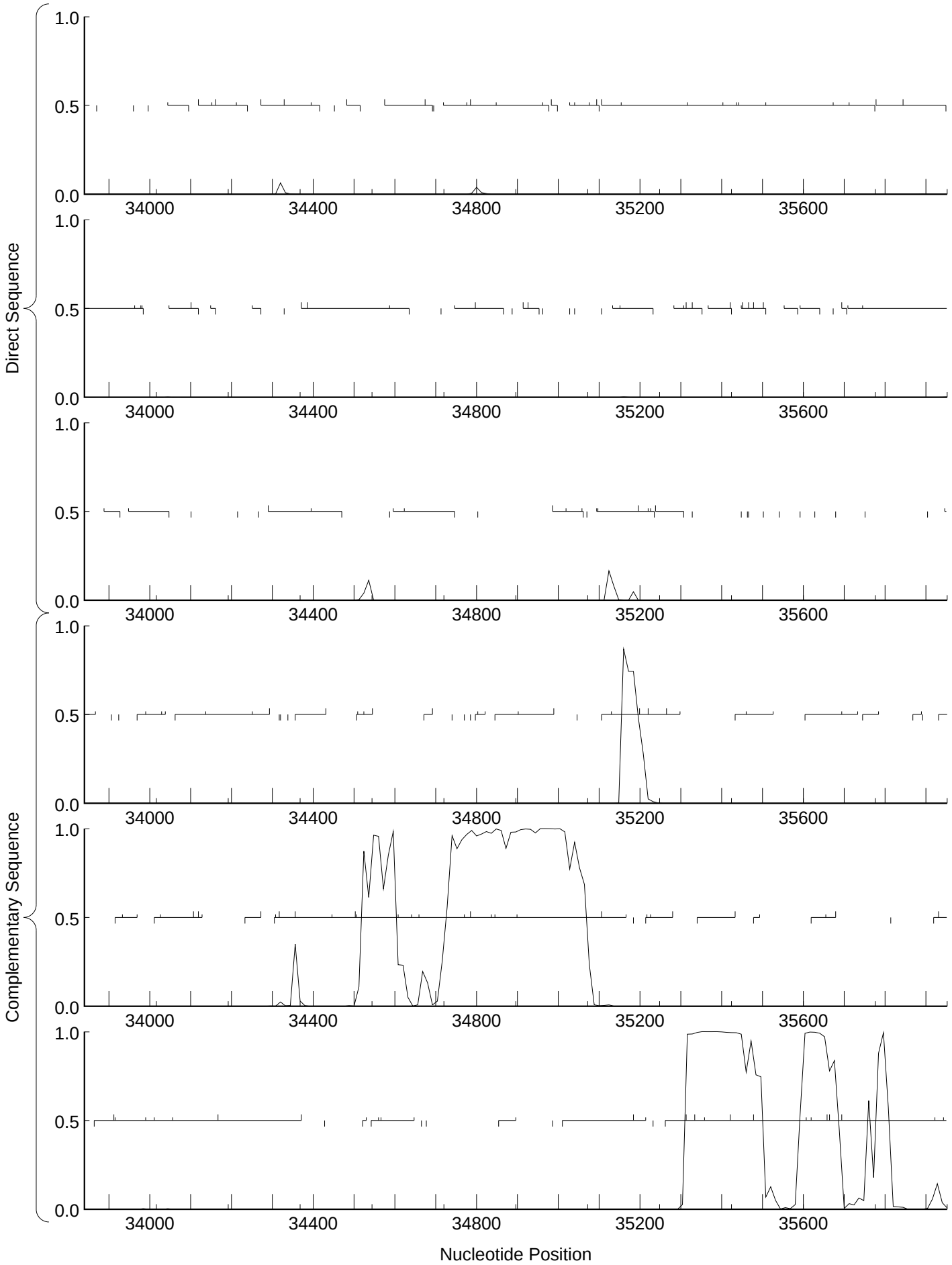


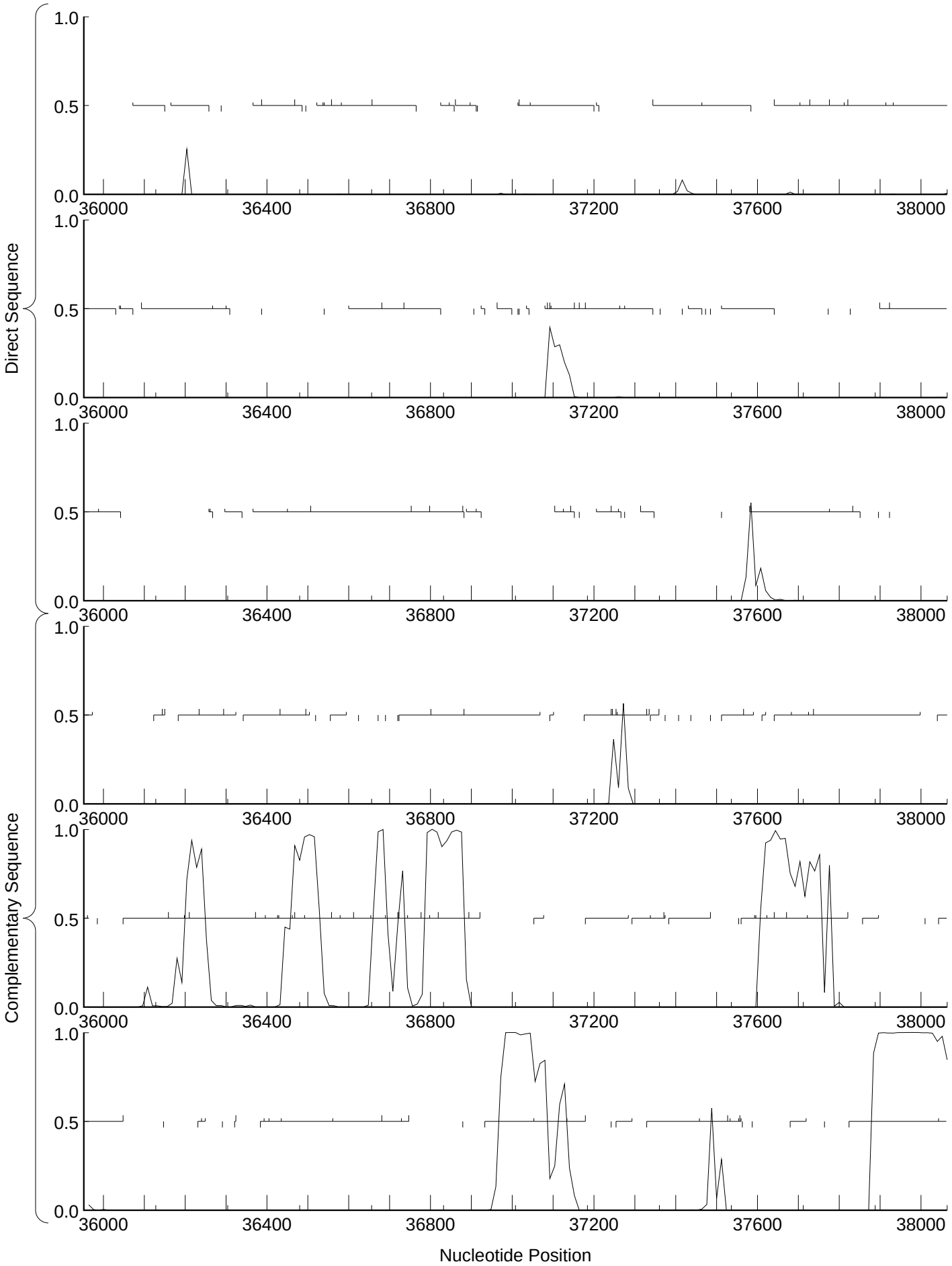
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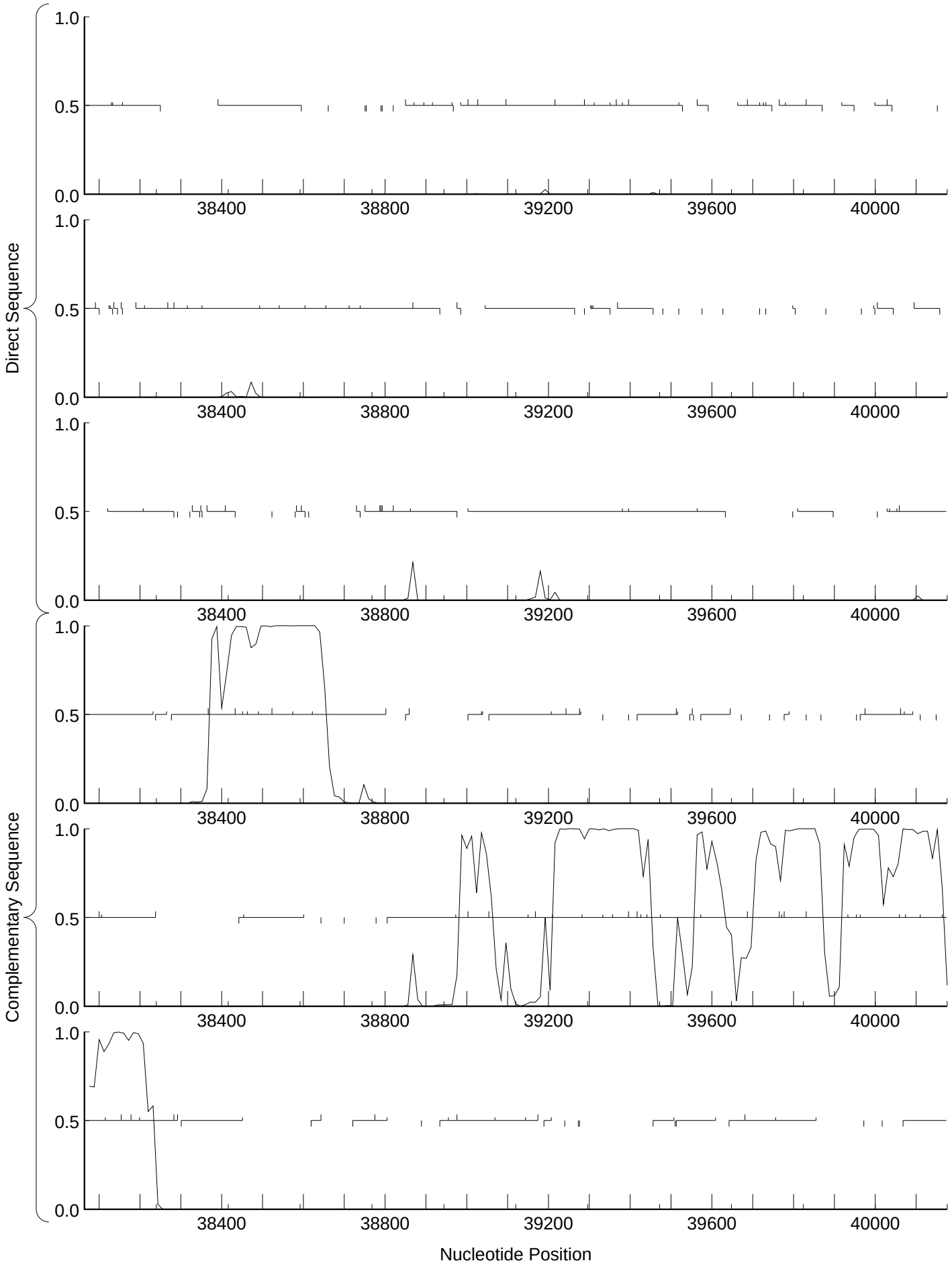


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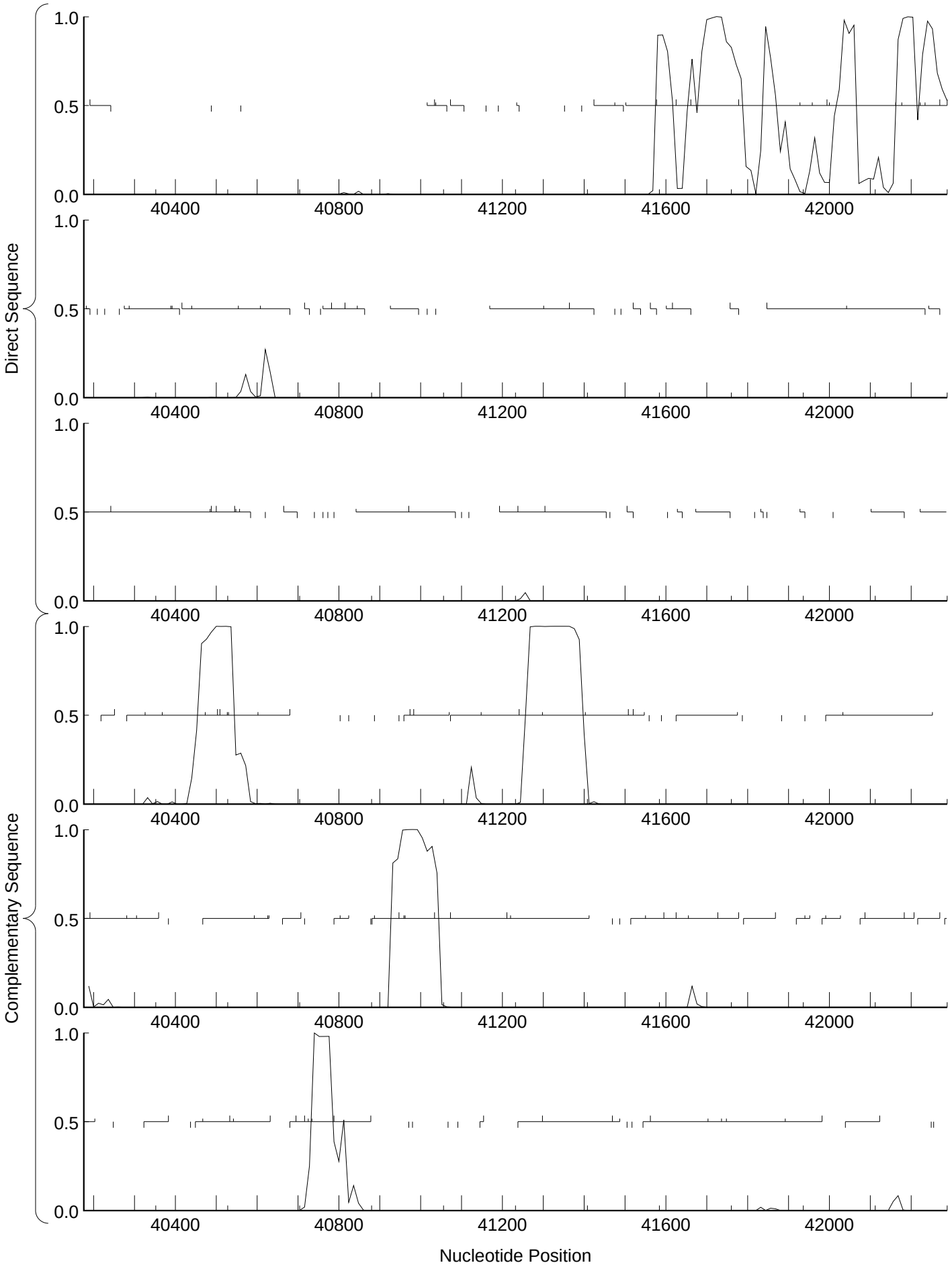



Direct Sequence
Complementary Sequence
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0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

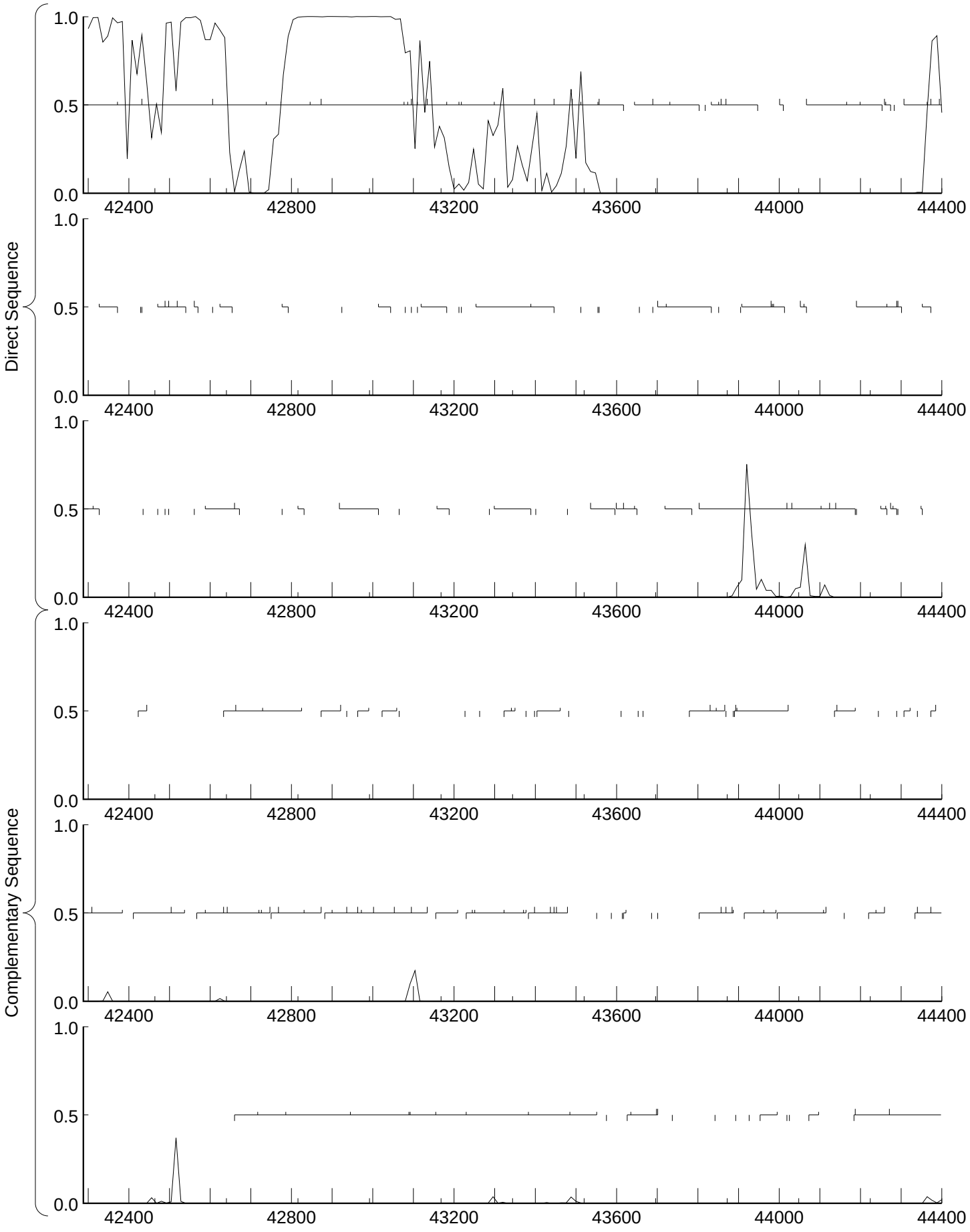

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
36000
36400
36800
37200
37600
38000
Nucleotide Position

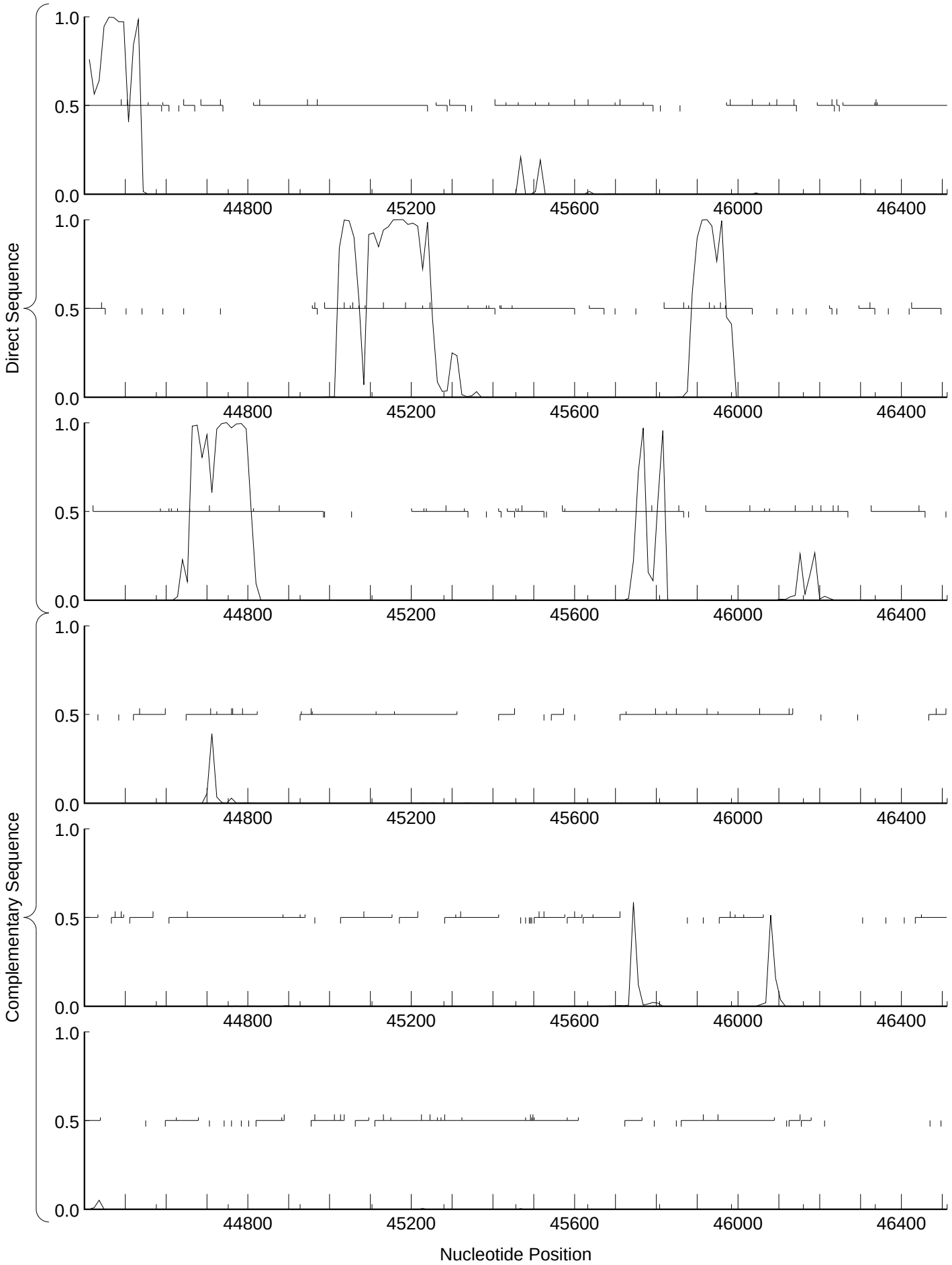

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

plete sequence, 47166 bp including 13-base 3’ overhang (CGGCGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 21/24



plete sequence, 47166 bp including 13-base 3’ overhang (CGGCGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 22/24




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
44800
45200
45600
46000
46400
Nucleotide Position

plete sequence, 47166 bp including 13-base 3’ overhang (CGGCGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 24/24

