GeneMark

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

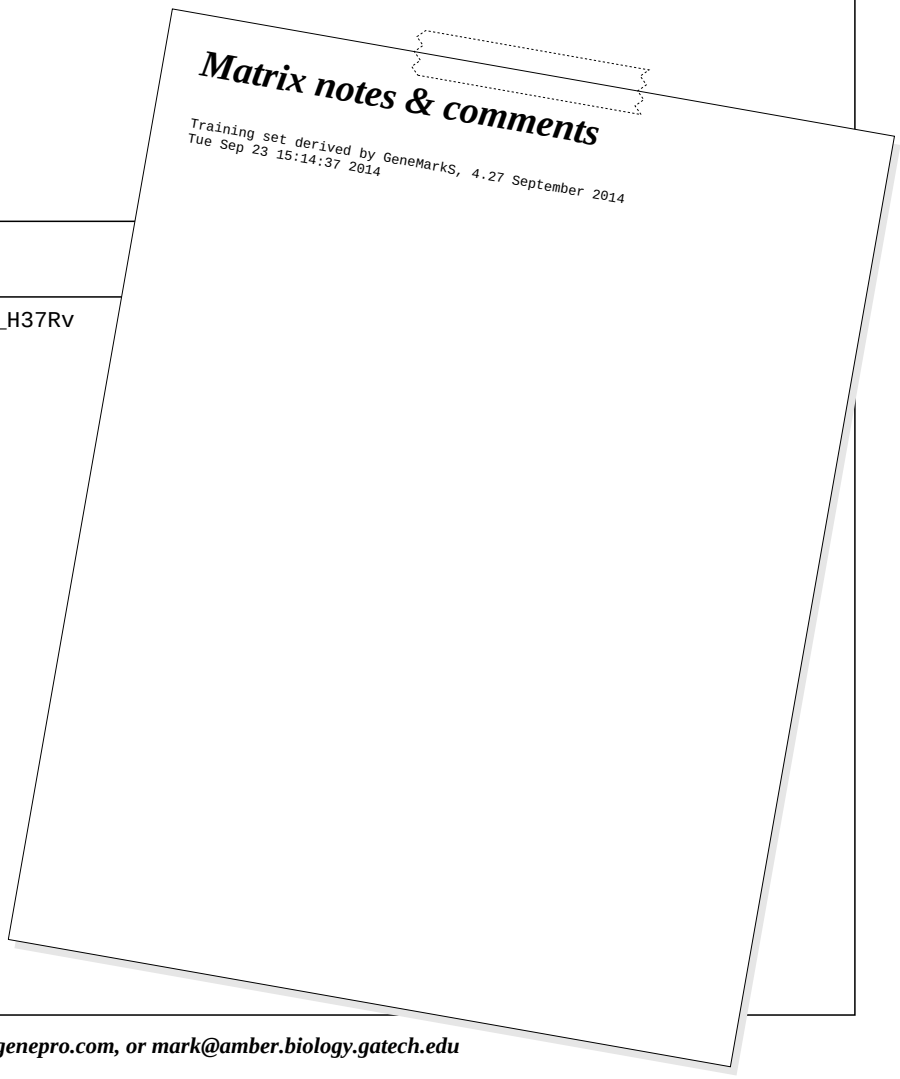
PROGRAM INFORMATION

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Sequence         :  Mycobacterium phage Datway complete sequence, 51303 bp including 10-base 3’ overhang (CGGCC
Analysis Date    :  3/29/24 at 2:37:22
Pages            :  26
Sequence Length  :  51303 bp
GC Content       :   63.67%

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

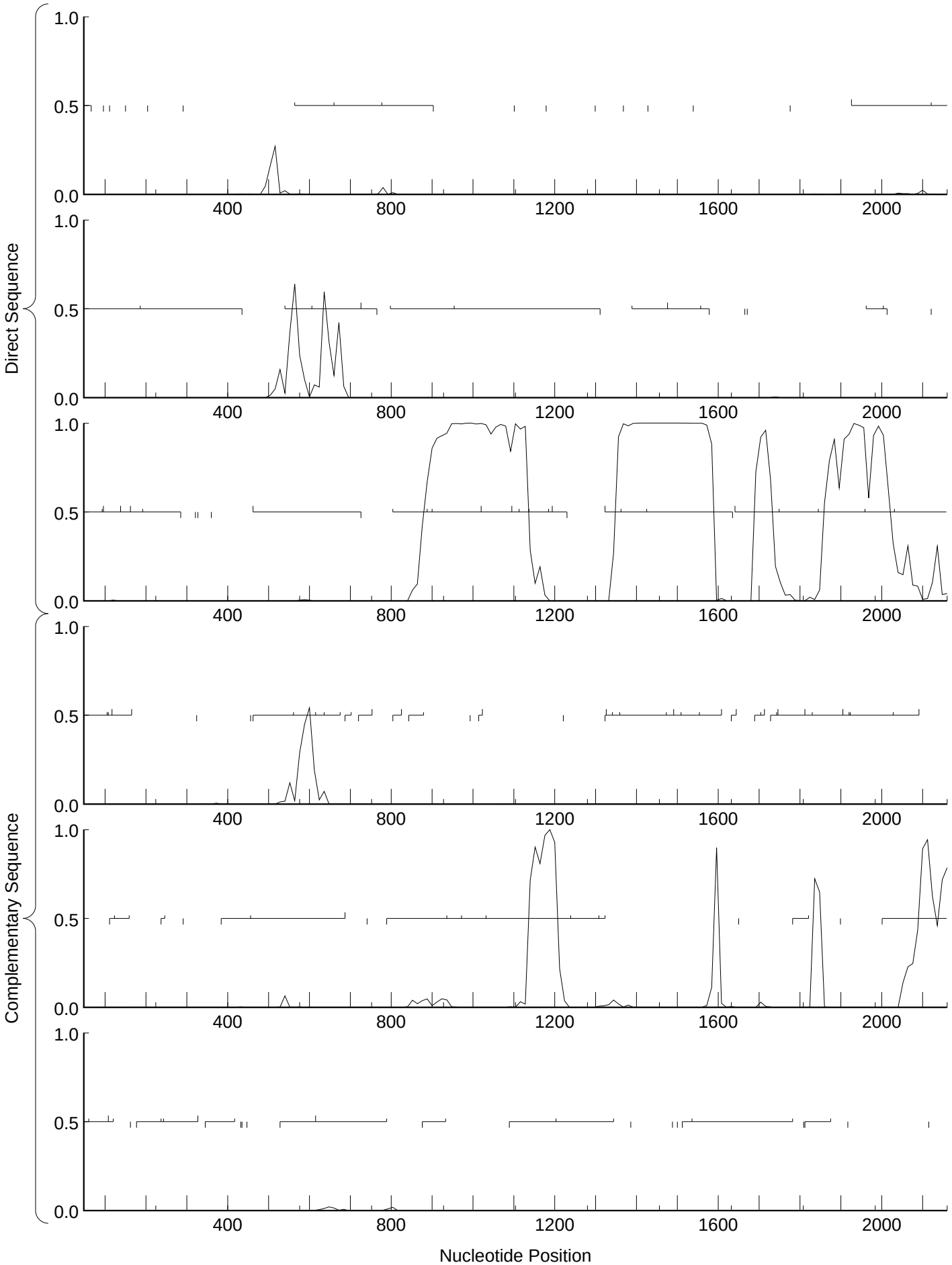
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Author     :  -

Order      :  4

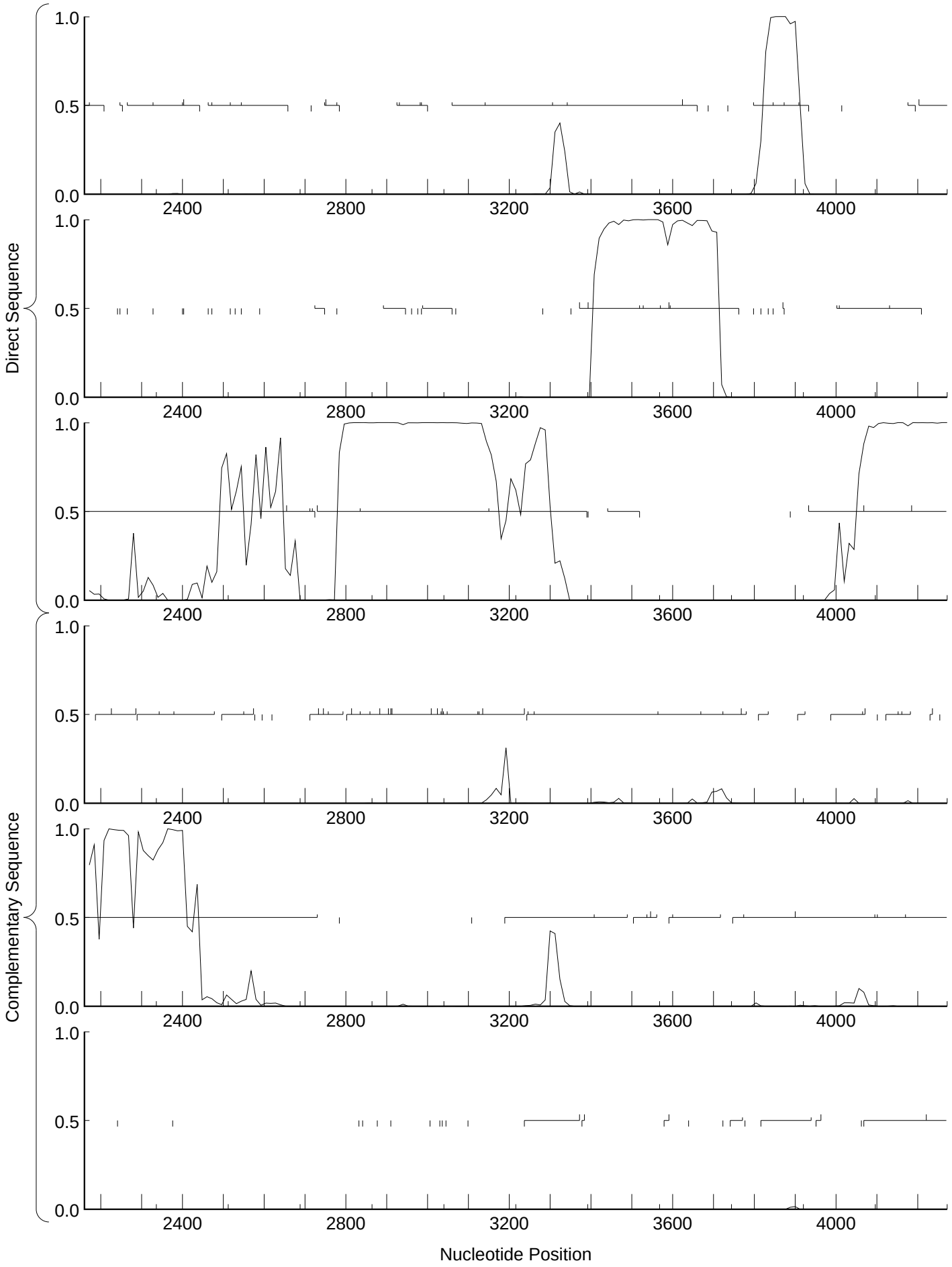

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

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

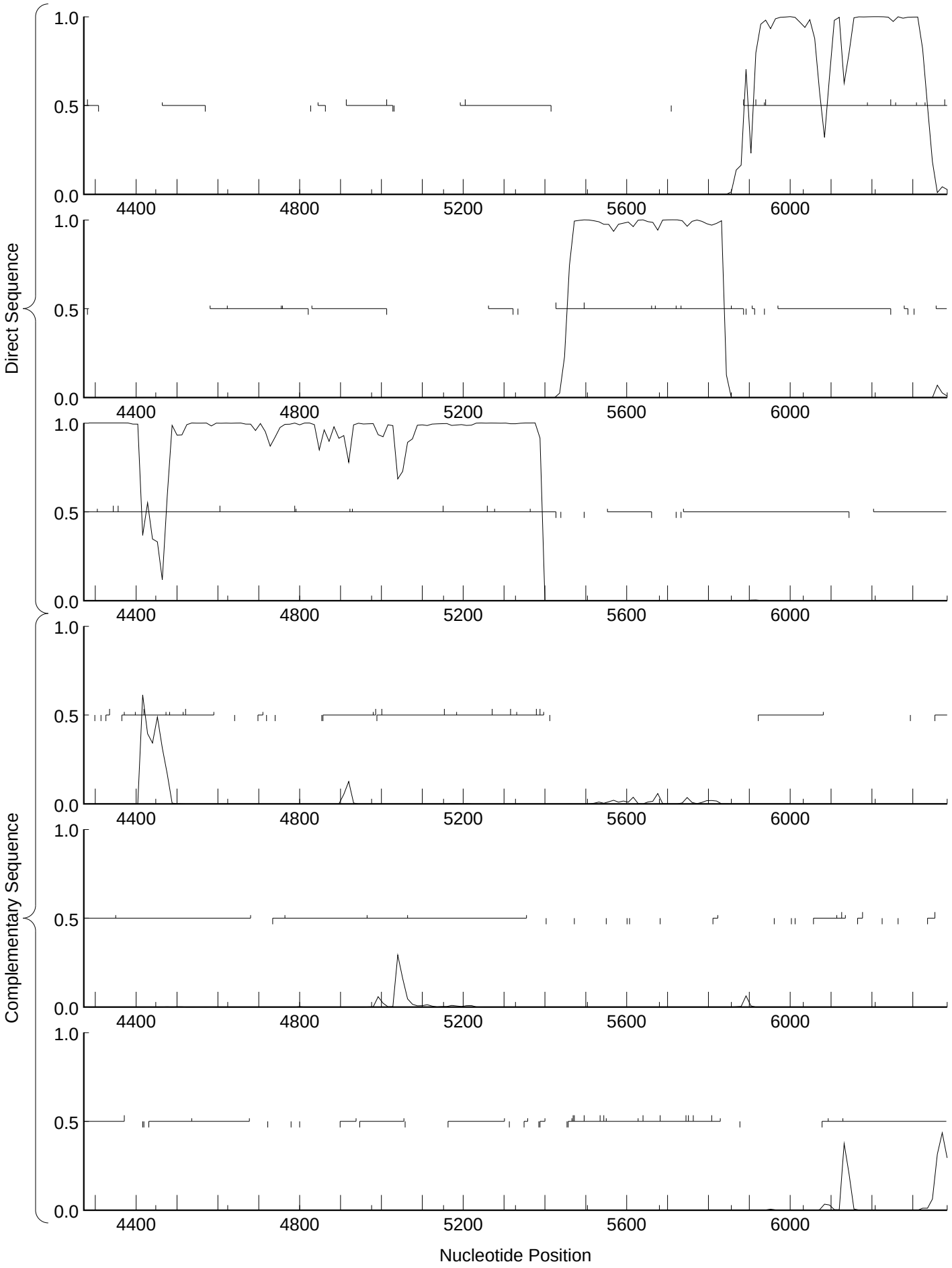
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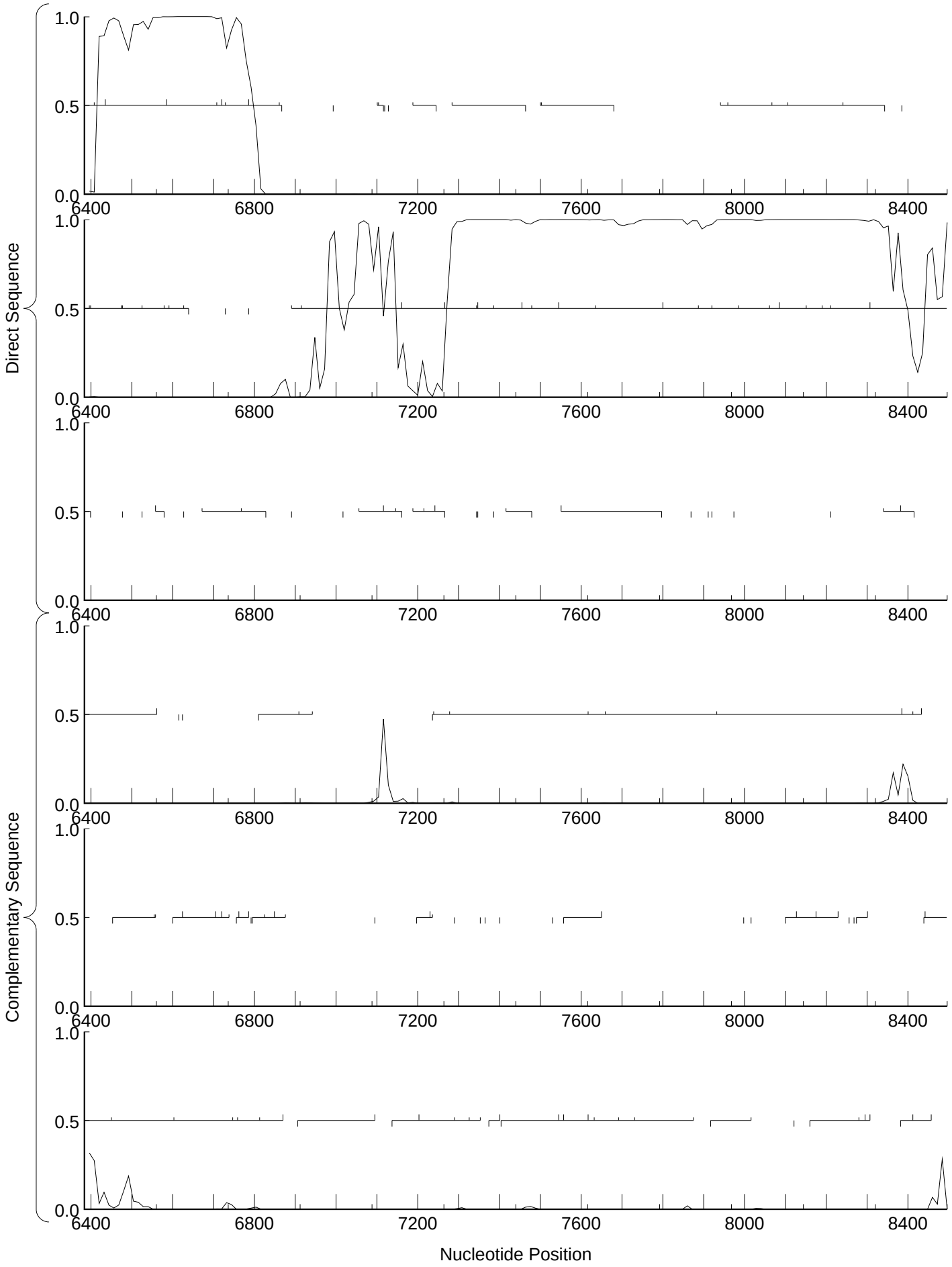
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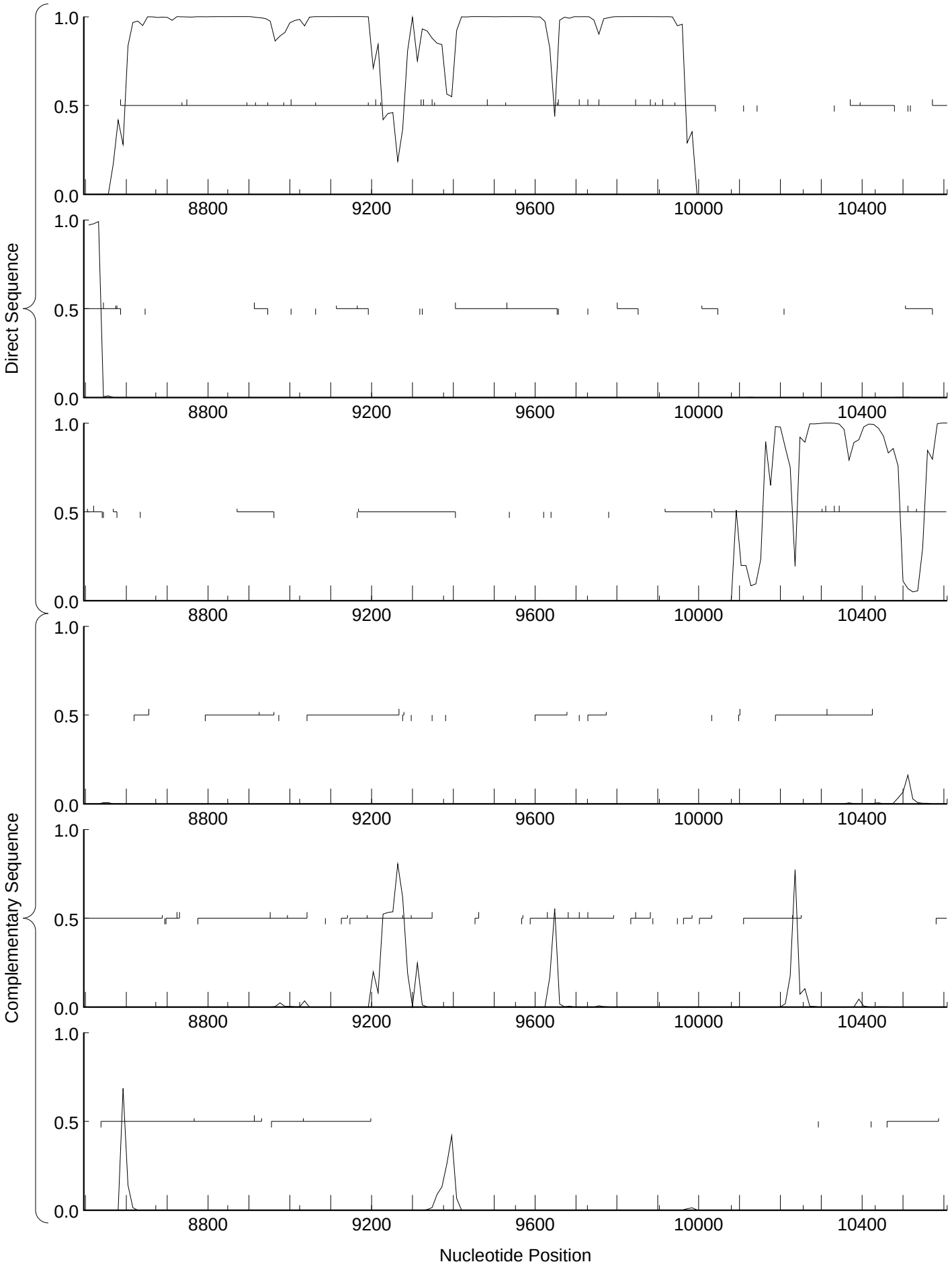
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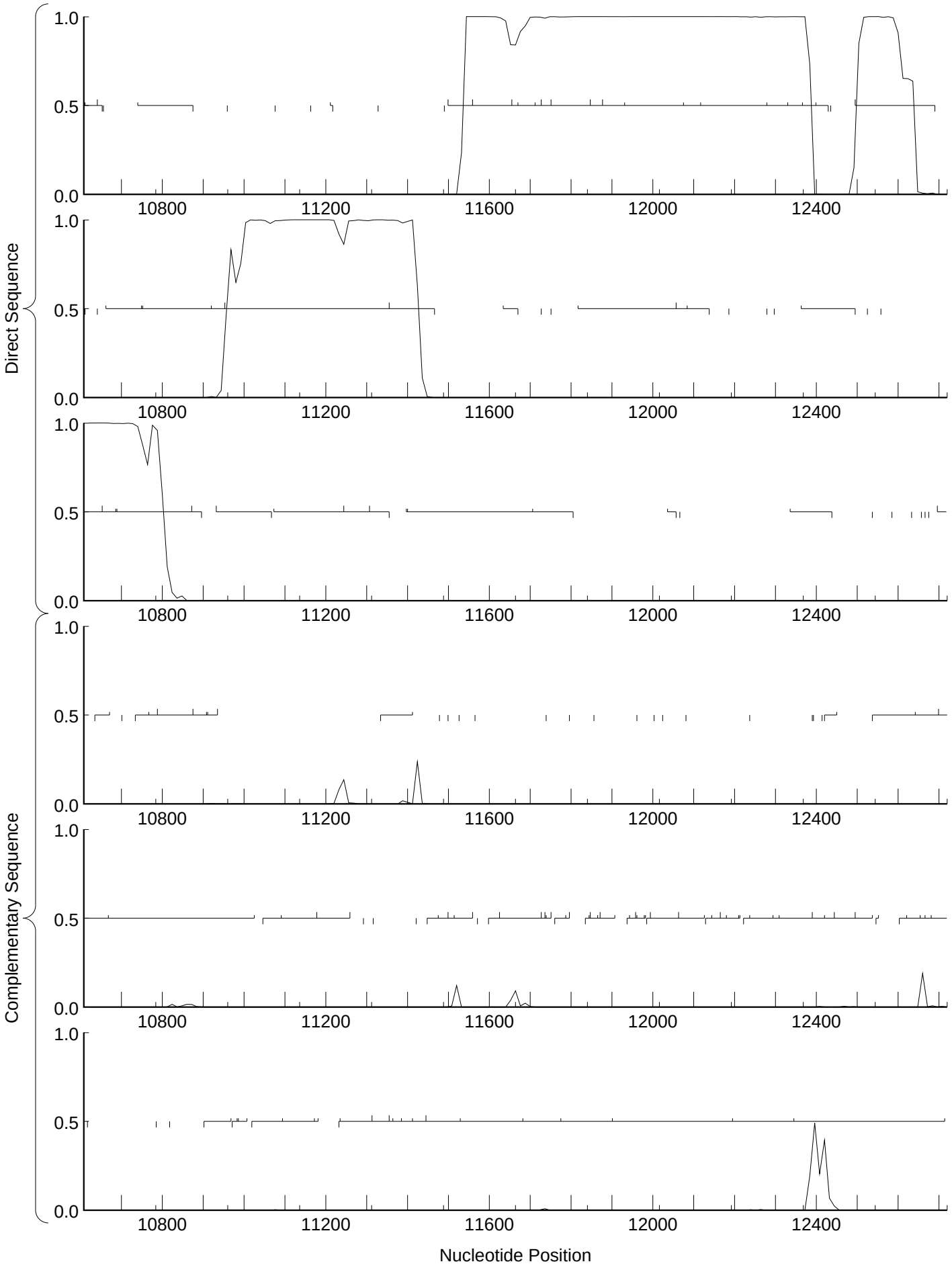
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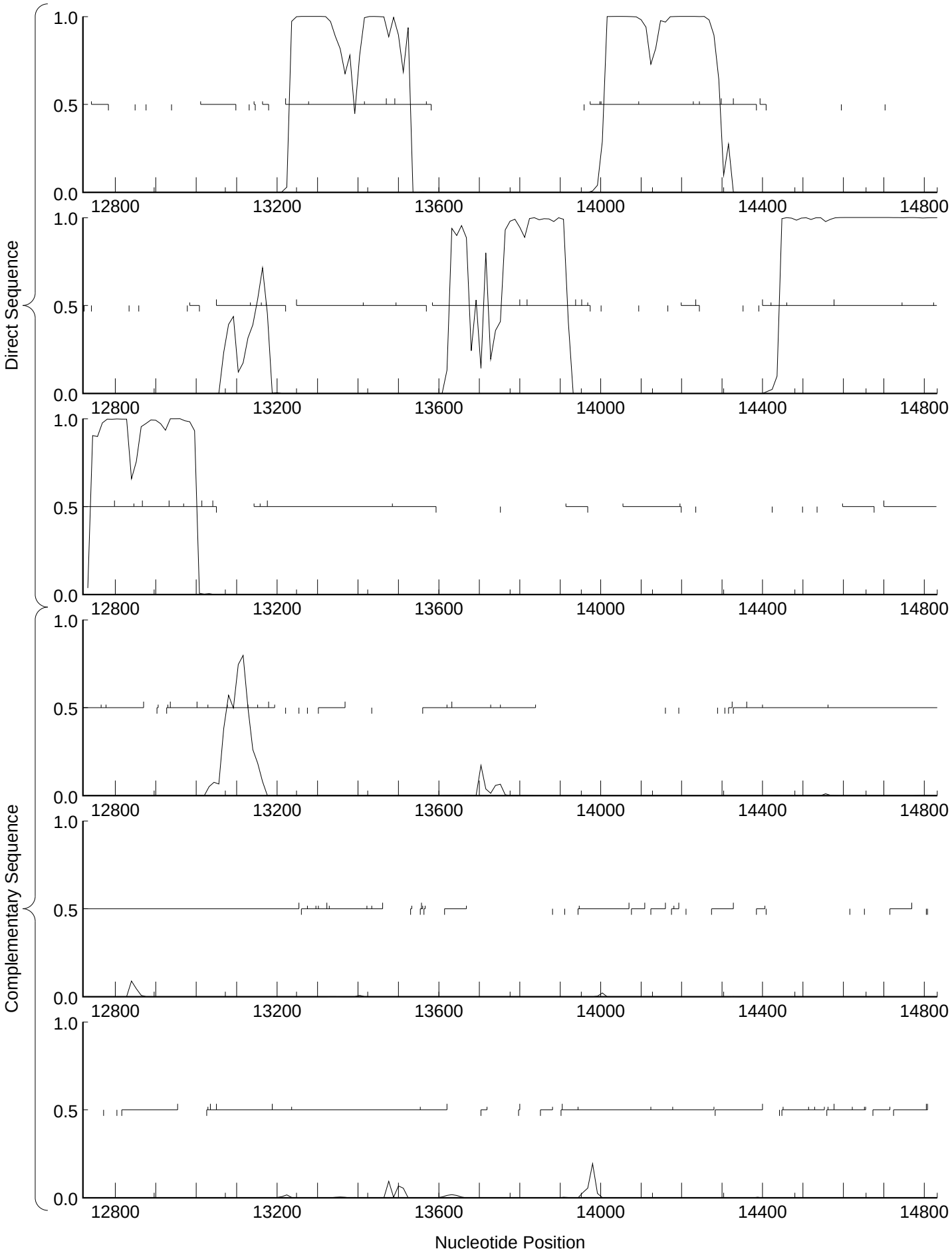
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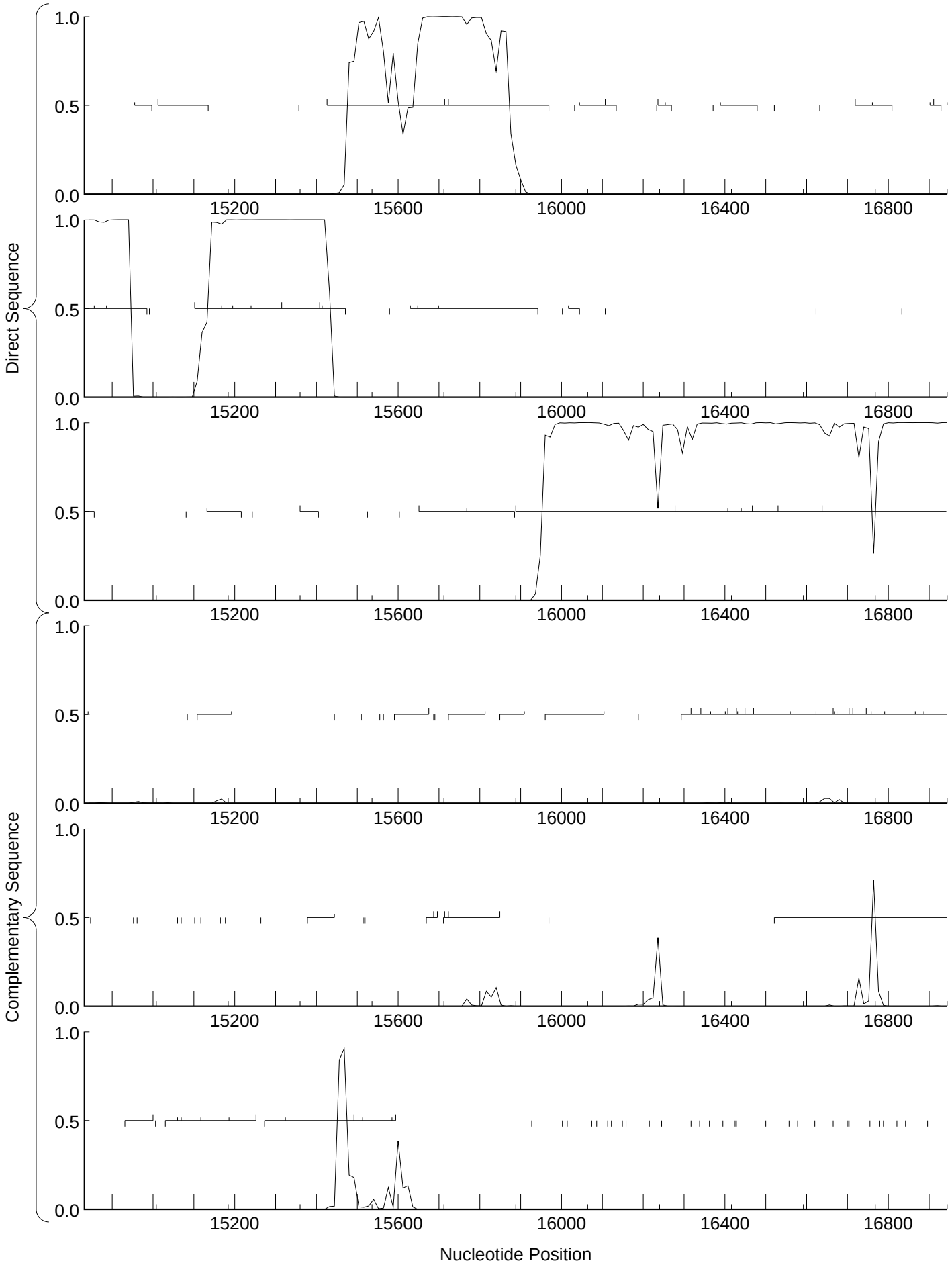
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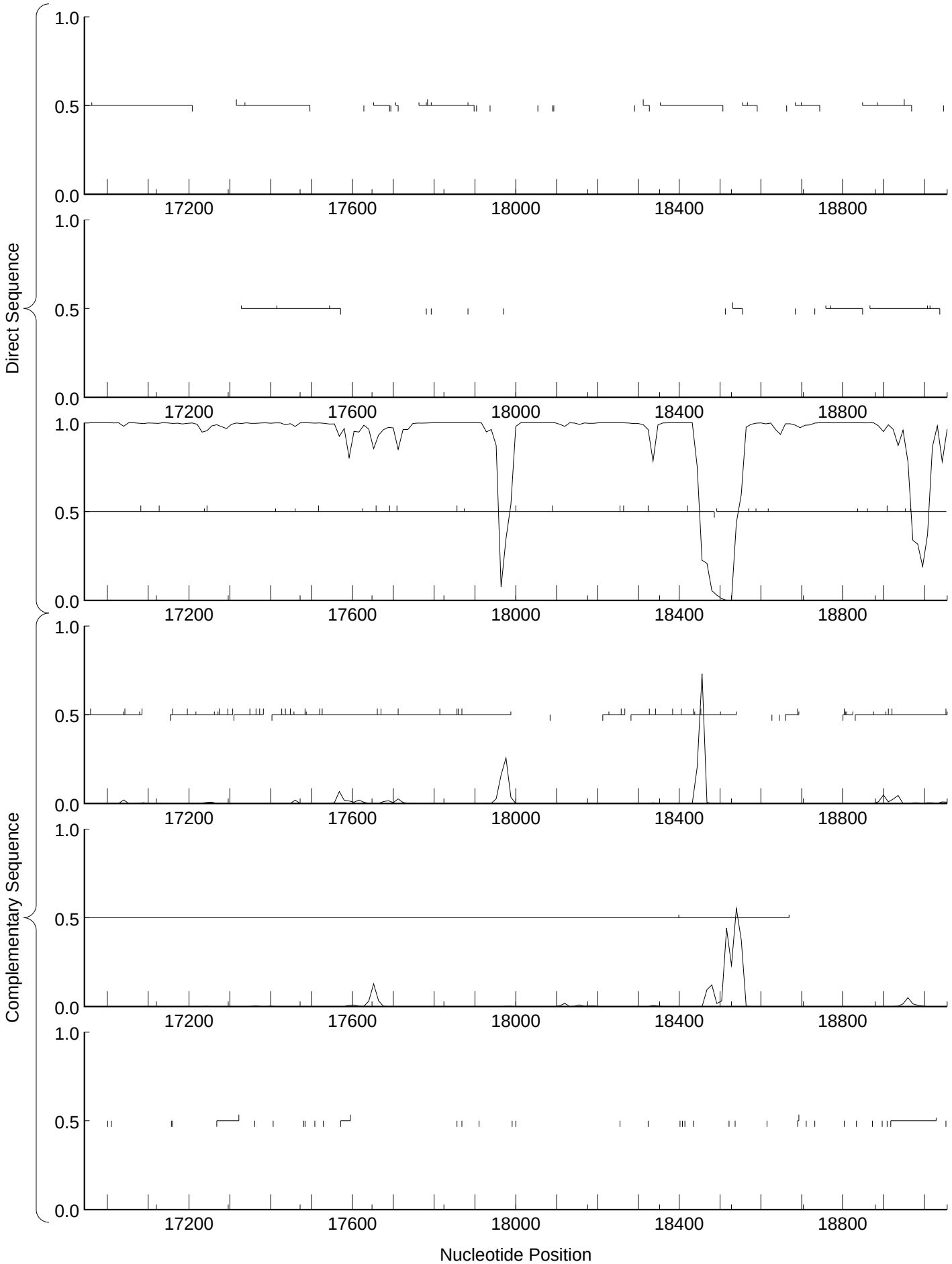
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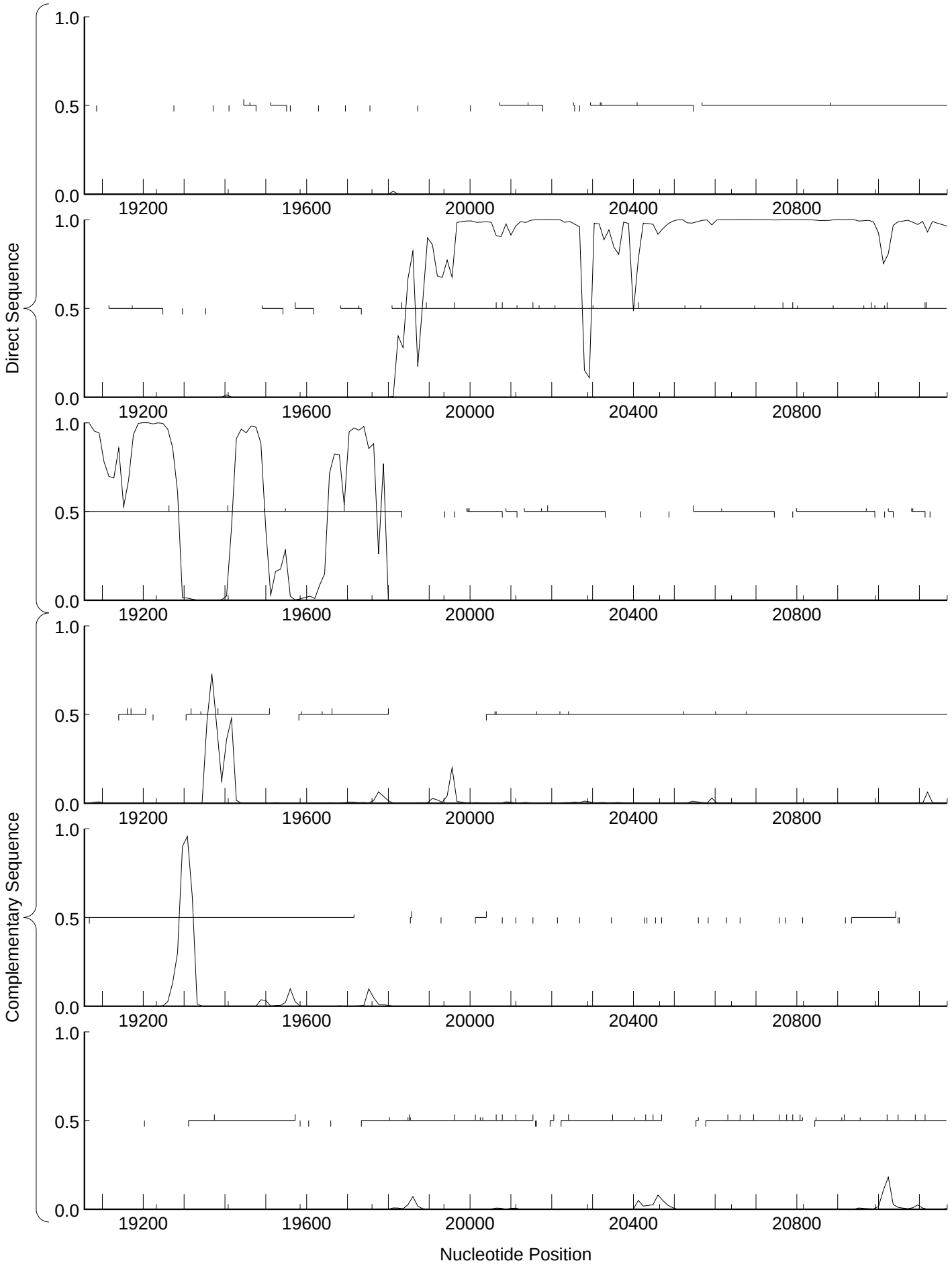
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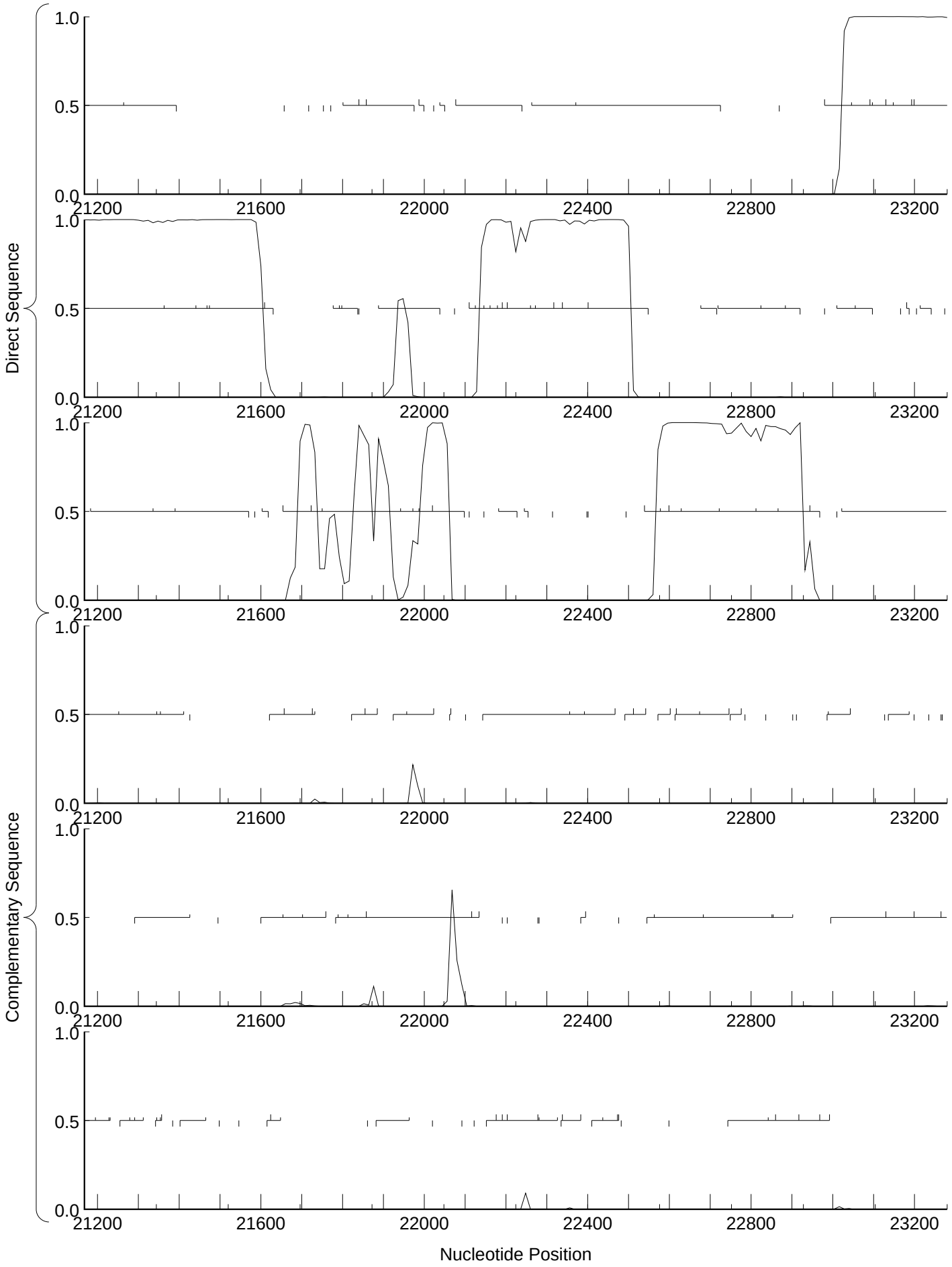
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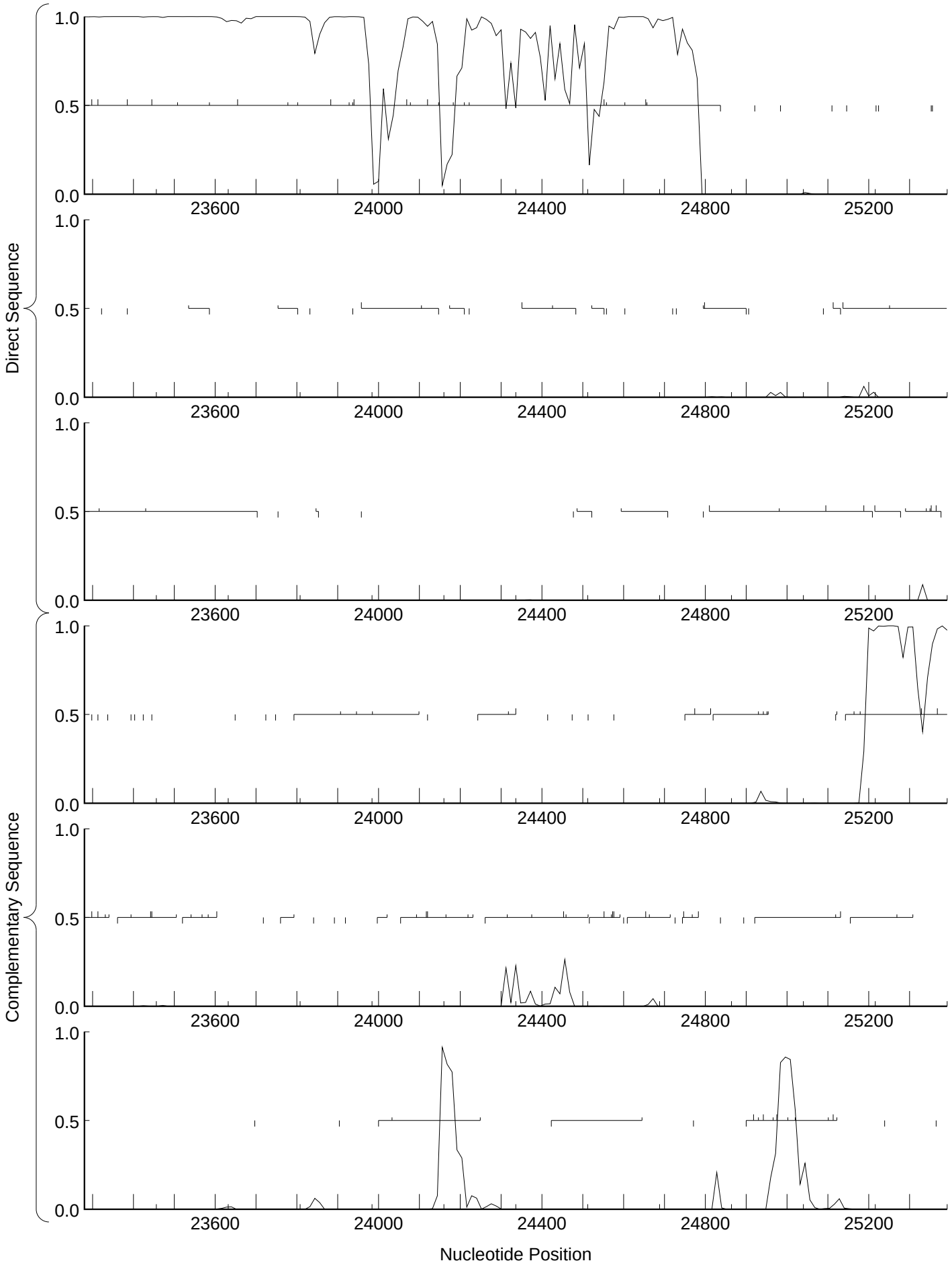
r complete sequence, 51303 bp including 10-base 3’ overhang (CGGCCGGTTA), Cluster A4, Order 4, Window 96, Step 12, 11/26



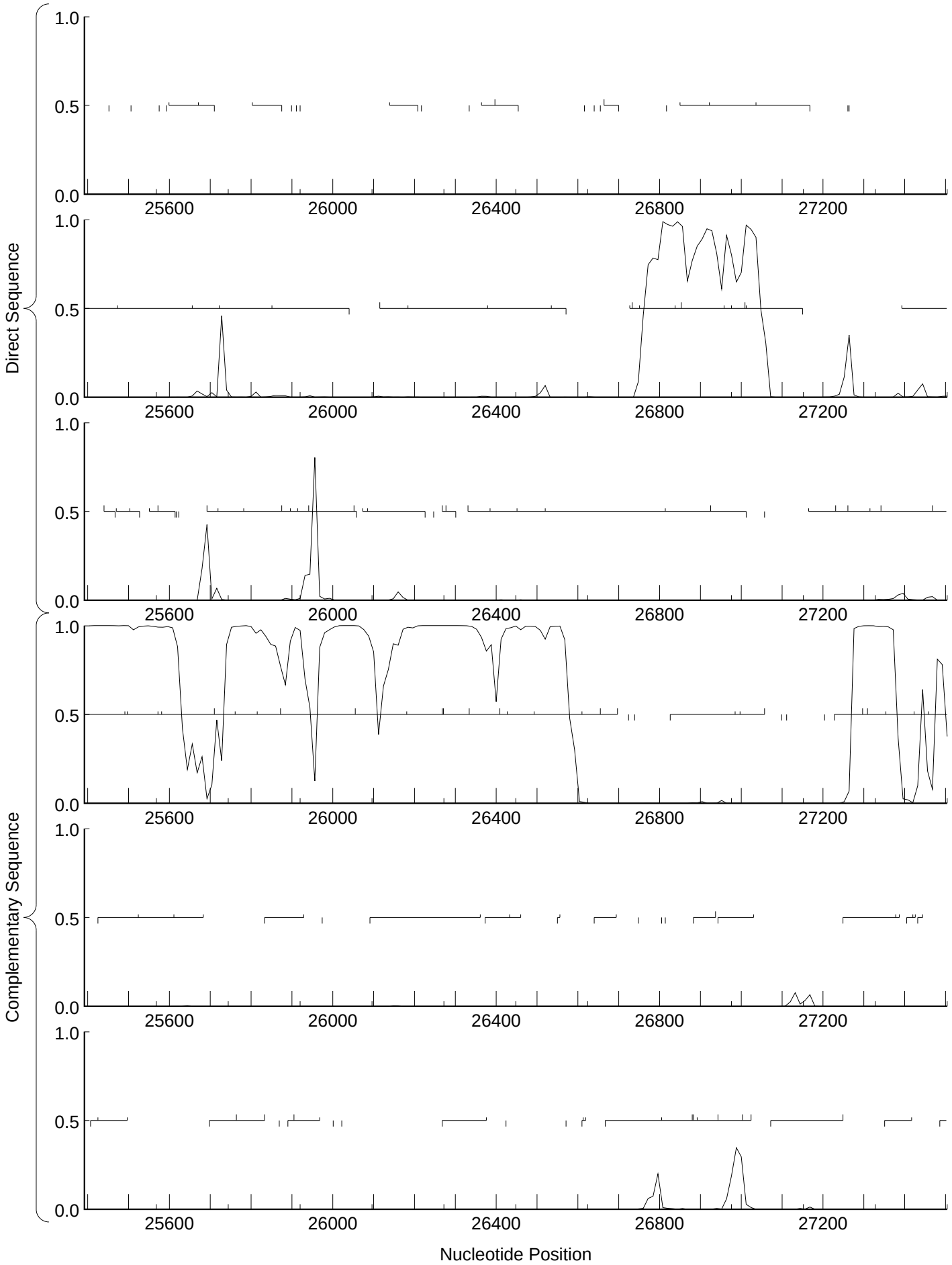
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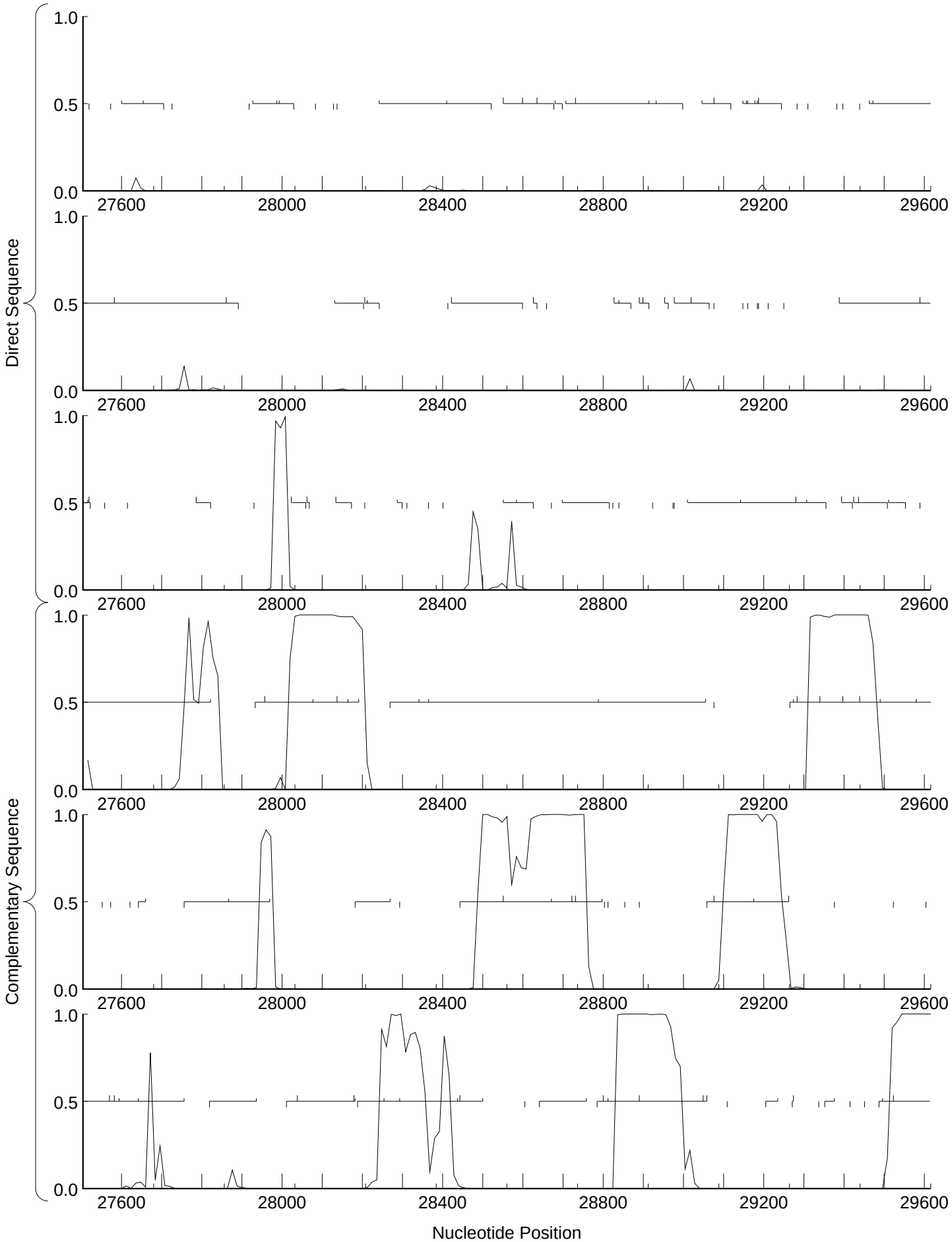
complete sequence, 51303 bp including 10-base 3' overhang (CGGCCGGTTA), Cluster A4, Order 4, Window 96, Step 12, 13/26

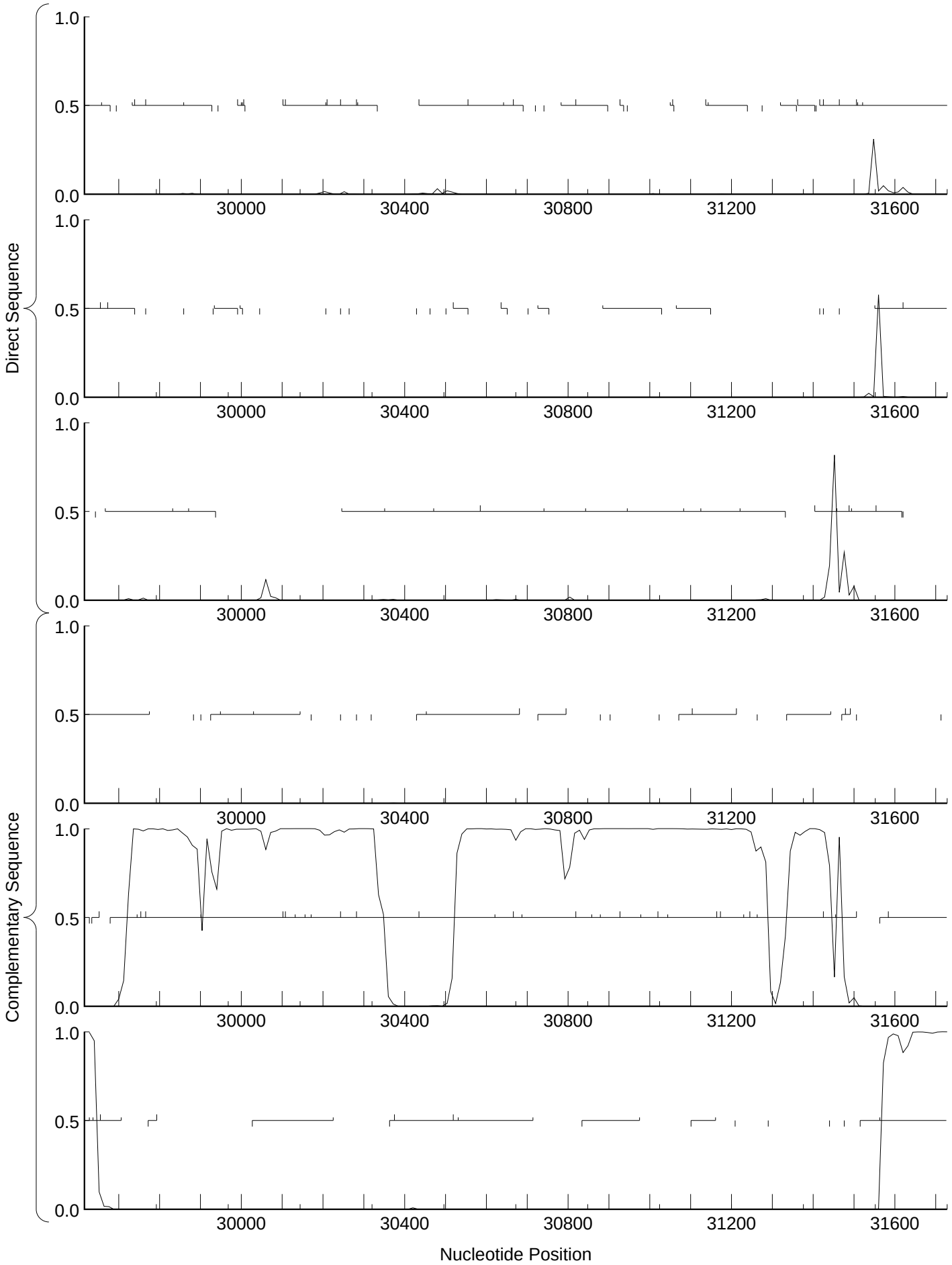


complete sequence, 51303 bp including 10-base 3’ overhang (CGGCCGGTTA), Cluster A4, Order 4, Window 96, Step 12, 14/26

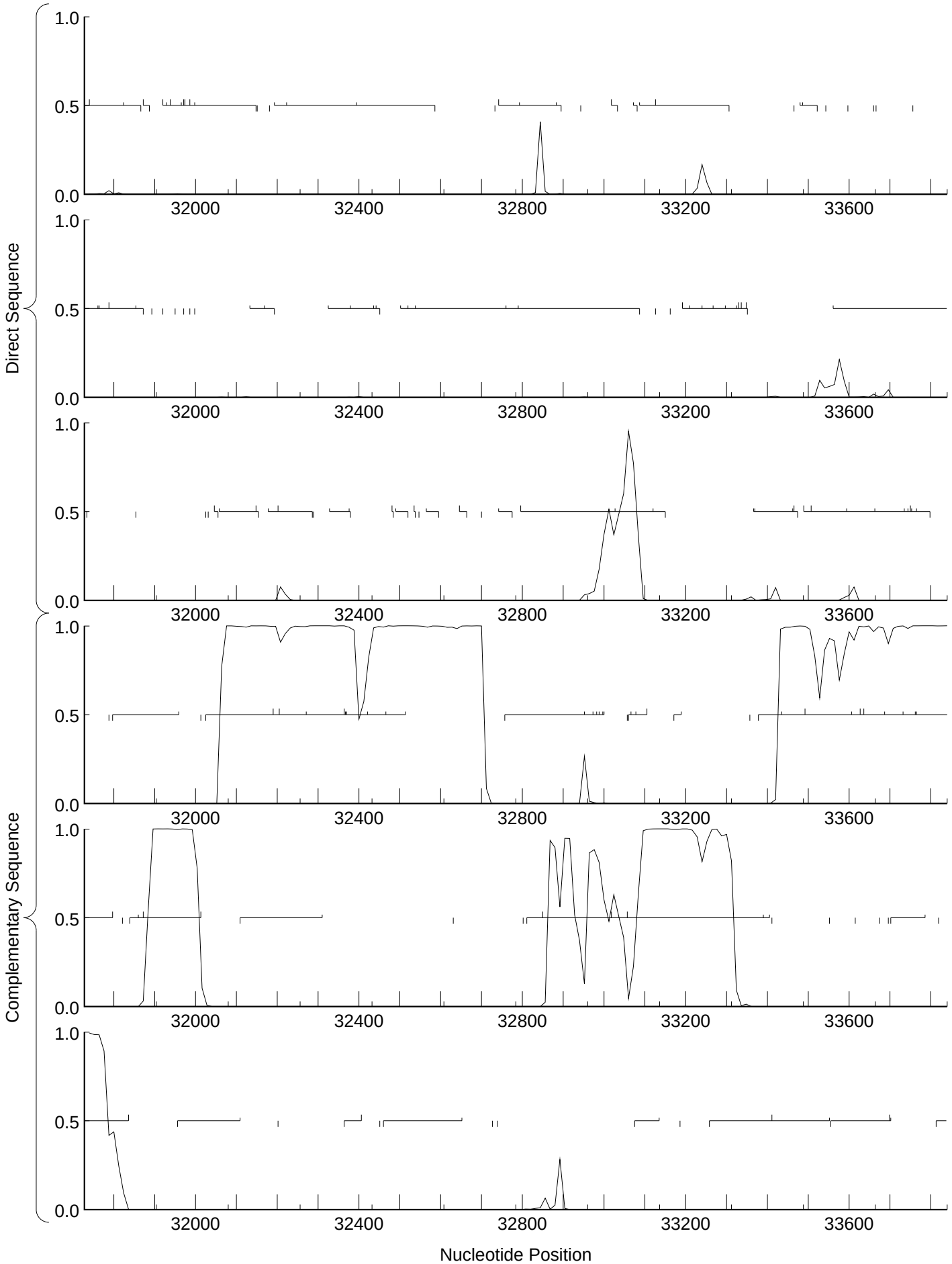


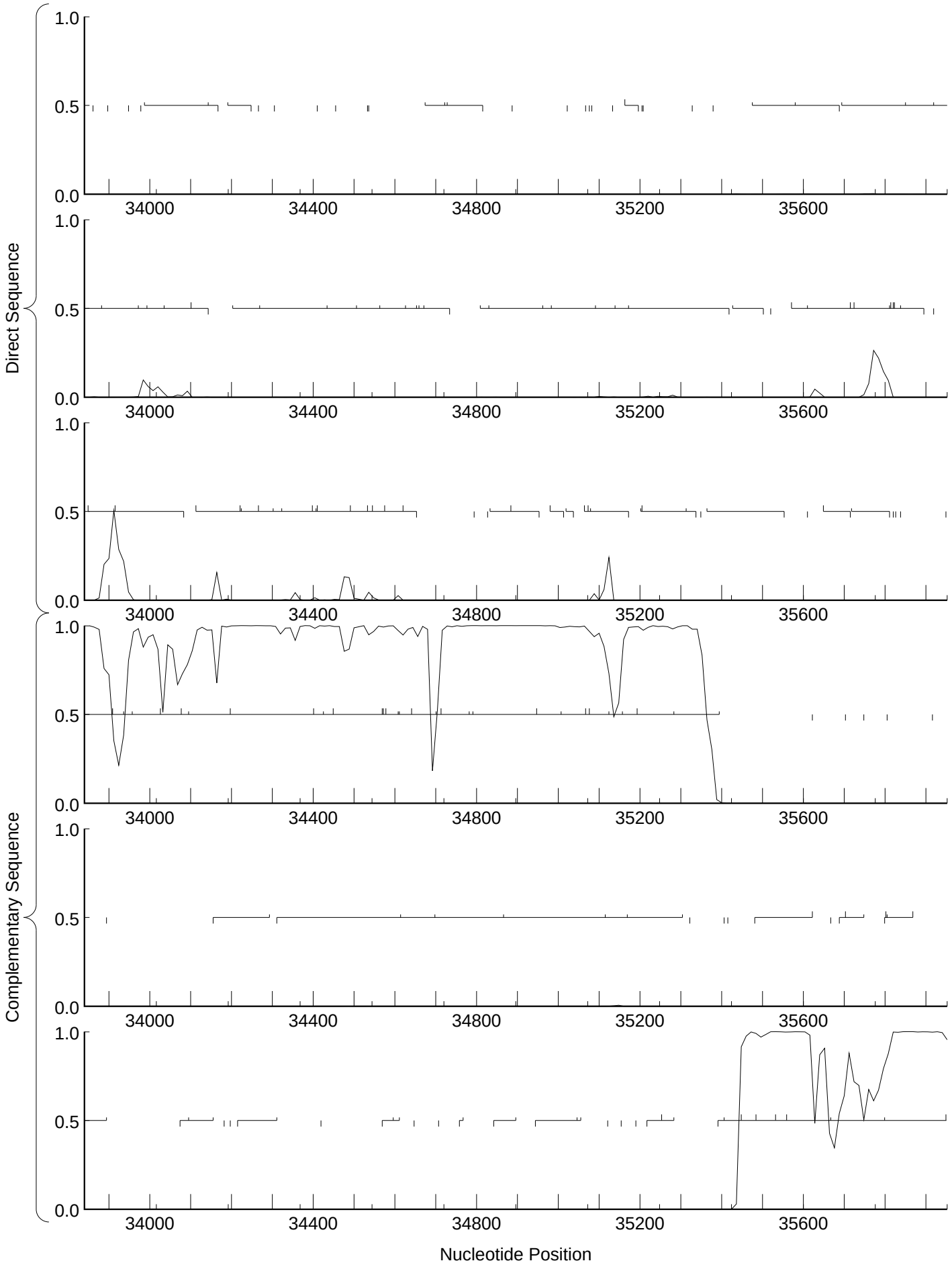
complete sequence, 51303 bp including 10-base 3’ overhang (CGGCCGGTTA), Cluster A4, Order 4, Window 96, Step 12, 15/26



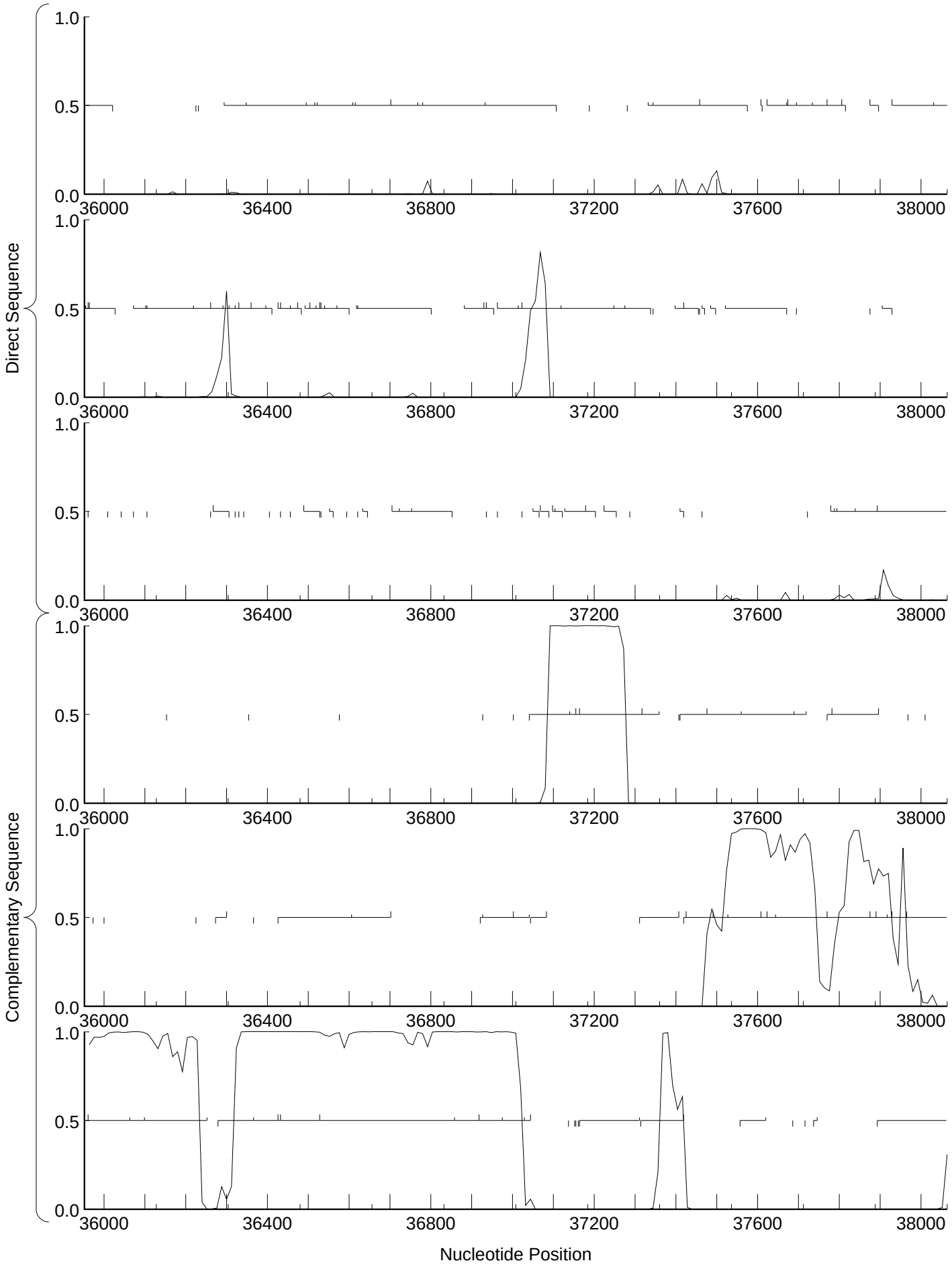

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
30000
30400
30800
31200
31600
Nucleotide Position

complete sequence, 51303 bp including 10-base 3’ overhang (CGGCCGGTTA), Cluster A4, Order 4, Window 96, Step 12, 17/26

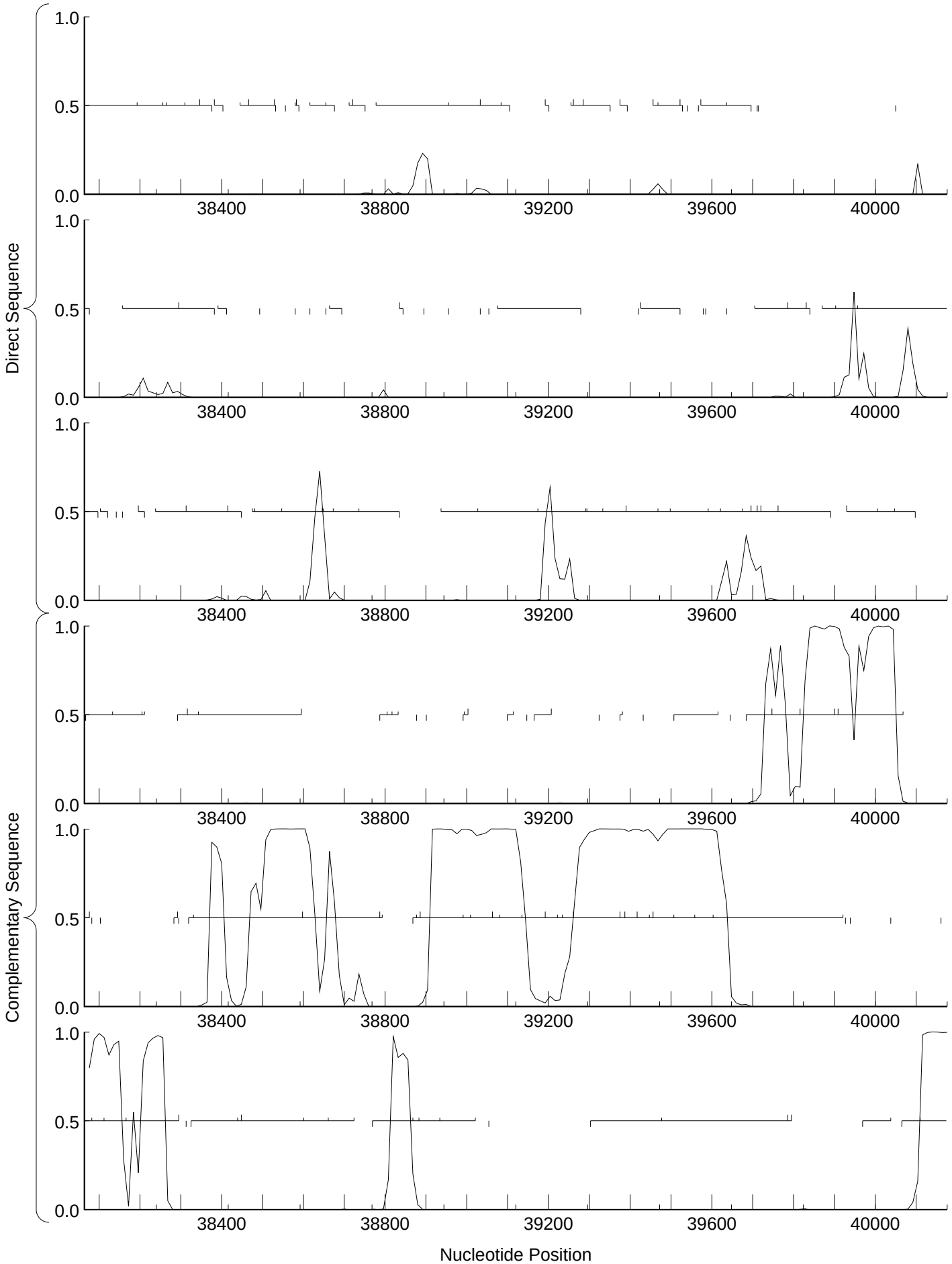



Direct Sequence
Complementary Sequence
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0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

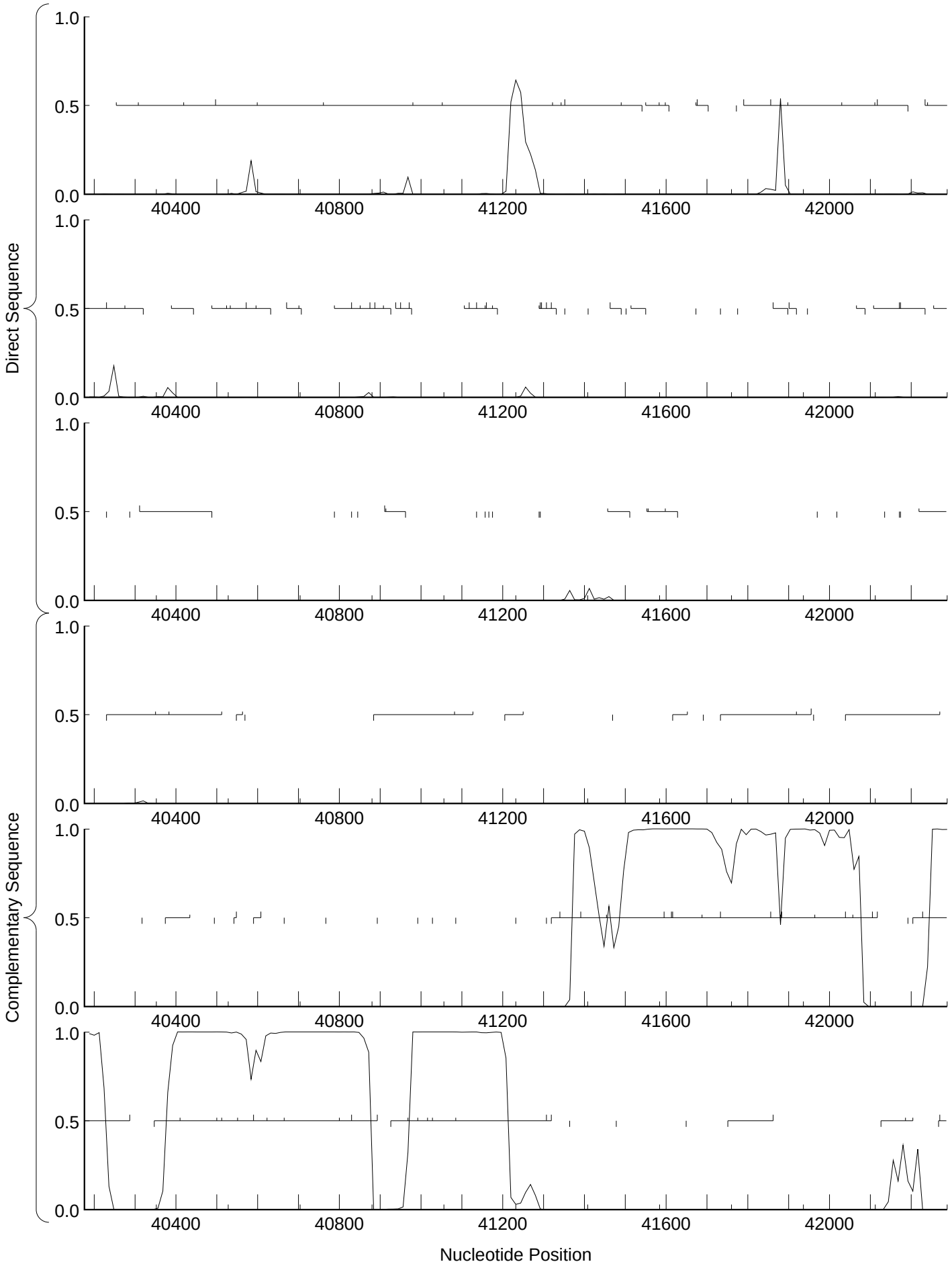
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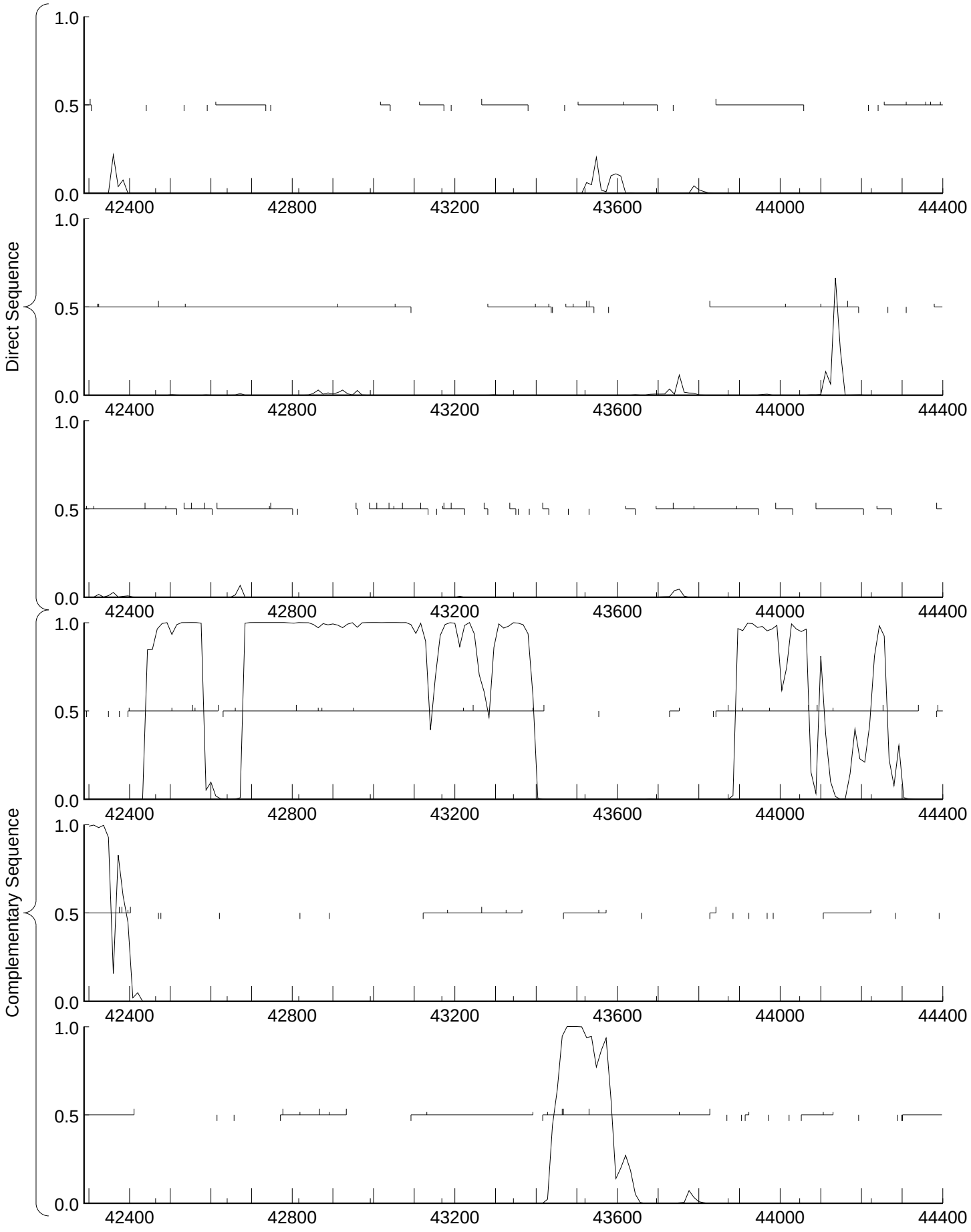


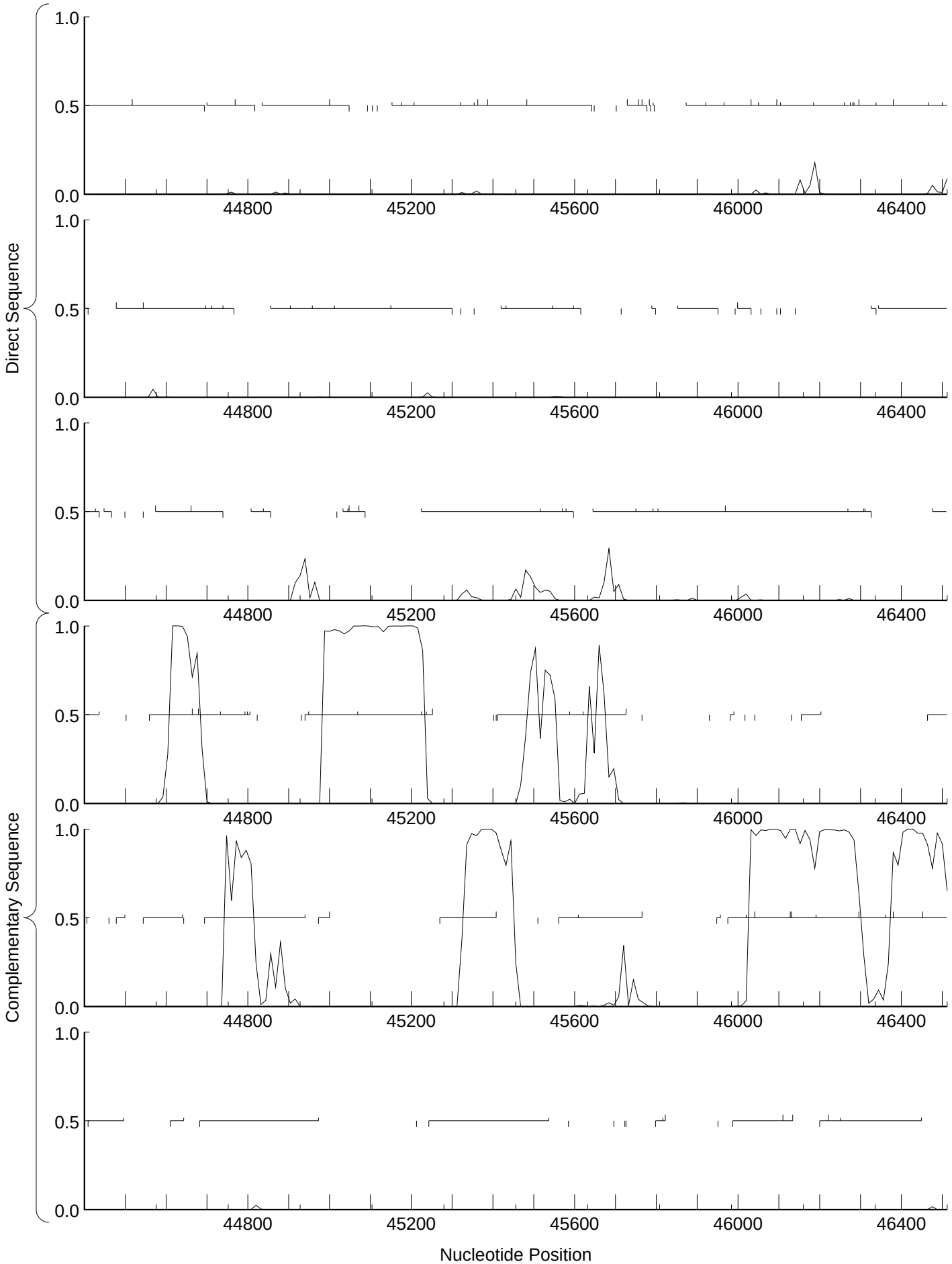
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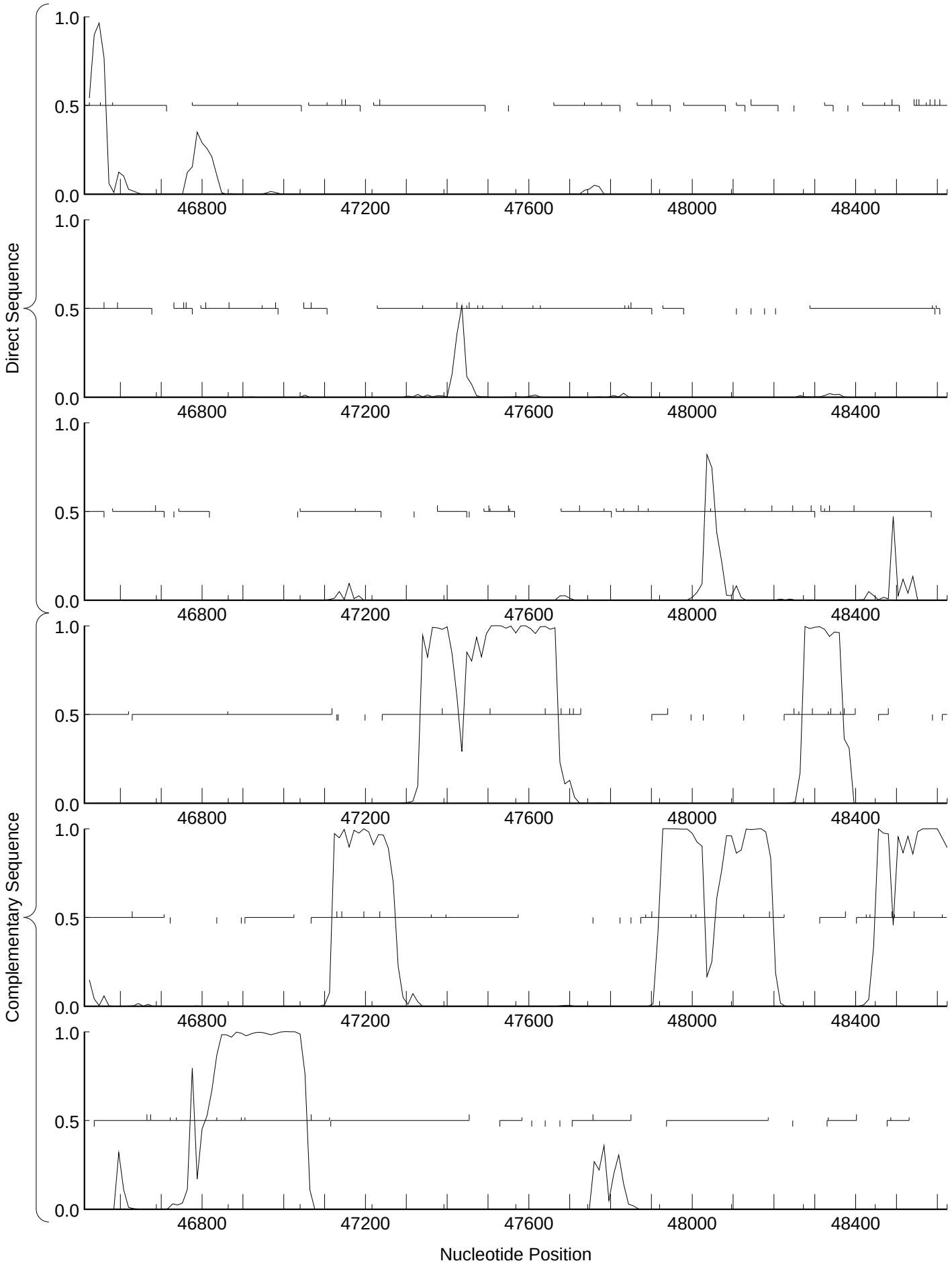
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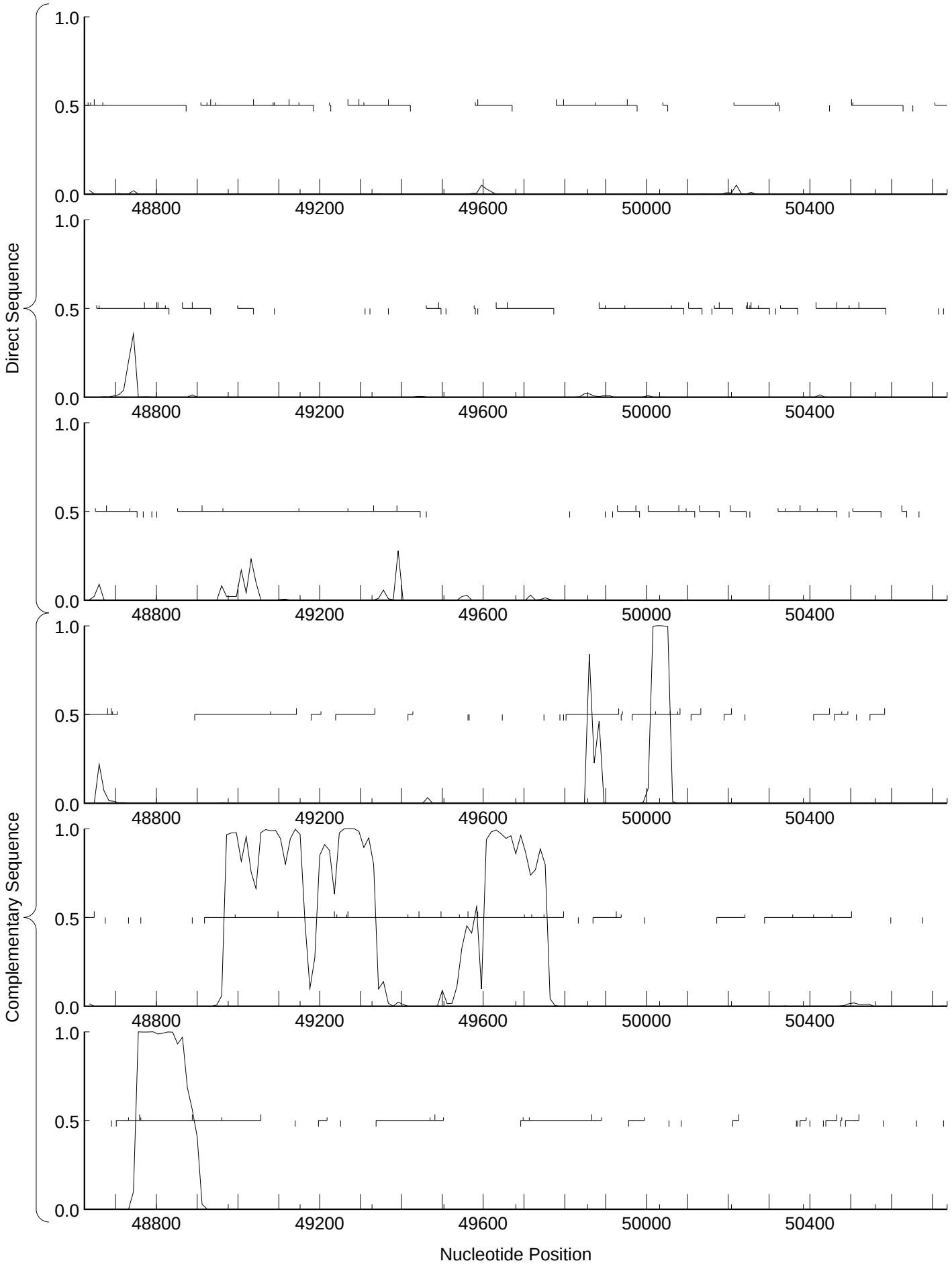

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
42400
42800
43200
43600
44000
44400
Nucleotide Position


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

complete sequence, 51303 bp including 10-base 3’ overhang (CGGCCGGTTA), Cluster A4, Order 4, Window 96, Step 12, 24/26



complete sequence, 51303 bp including 10-base 3’ overhang (CGGCCGGTTA), Cluster A4, Order 4, Window 96, Step 12, 25/26



r complete sequence, 51303 bp including 10-base 3’ overhang (CGGCCGGTTA), Cluster A4, Order 4, Window 96, Step 12, 26/26

