GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Arthrobacter phage Hum25 complete sequence, 42695 bp including 9-base 3’ overhang (CGCCGGTG
Analysis Date    :  3/29/24 at 5:04:21
Pages            :  22
Sequence Length  :  42695 bp
GC Content       :   59.54%

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 notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:22:00 2018

MATRIX INFORMATION

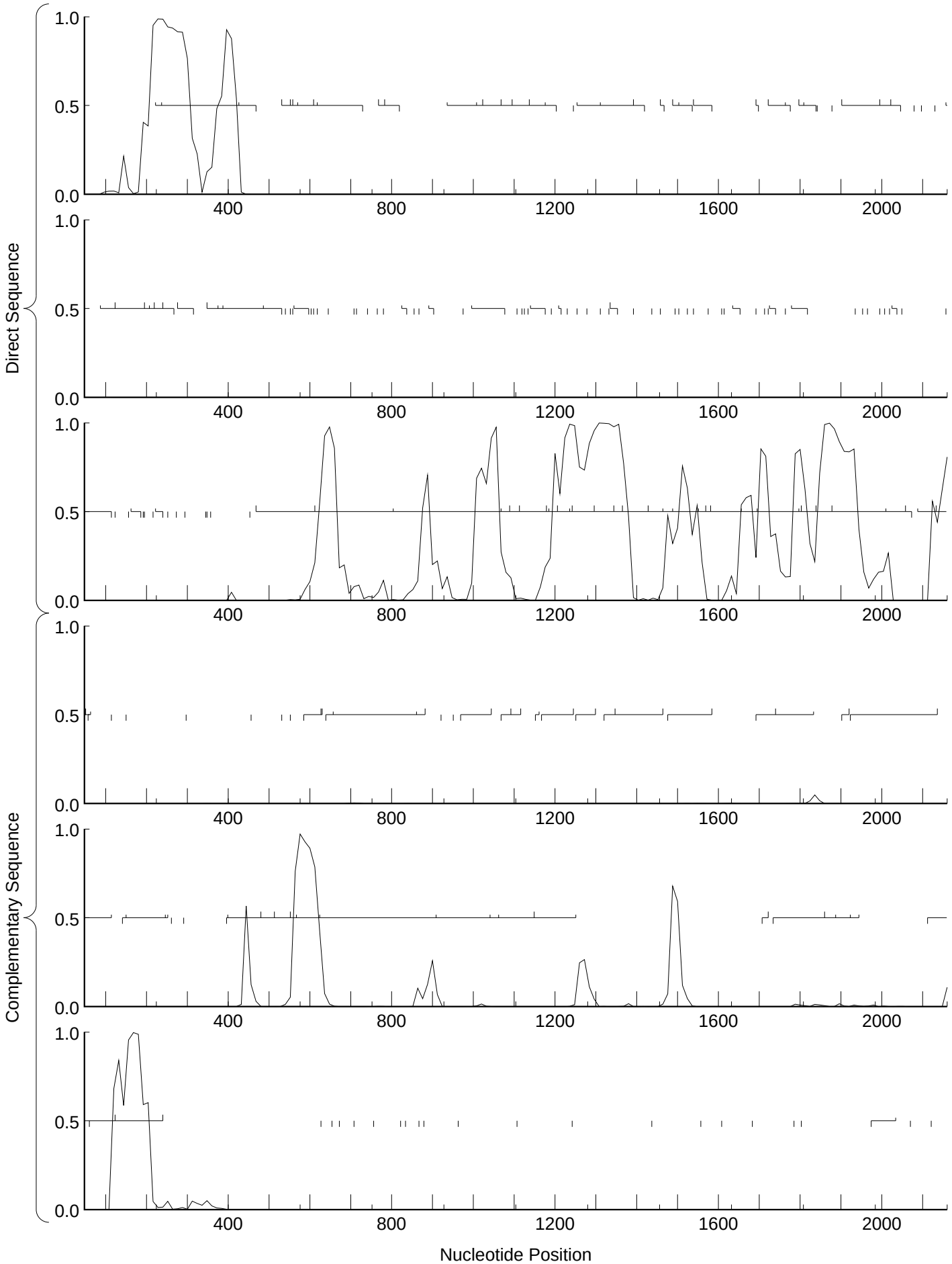
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Matrix     :  -
Author     :  -

Order      :  4
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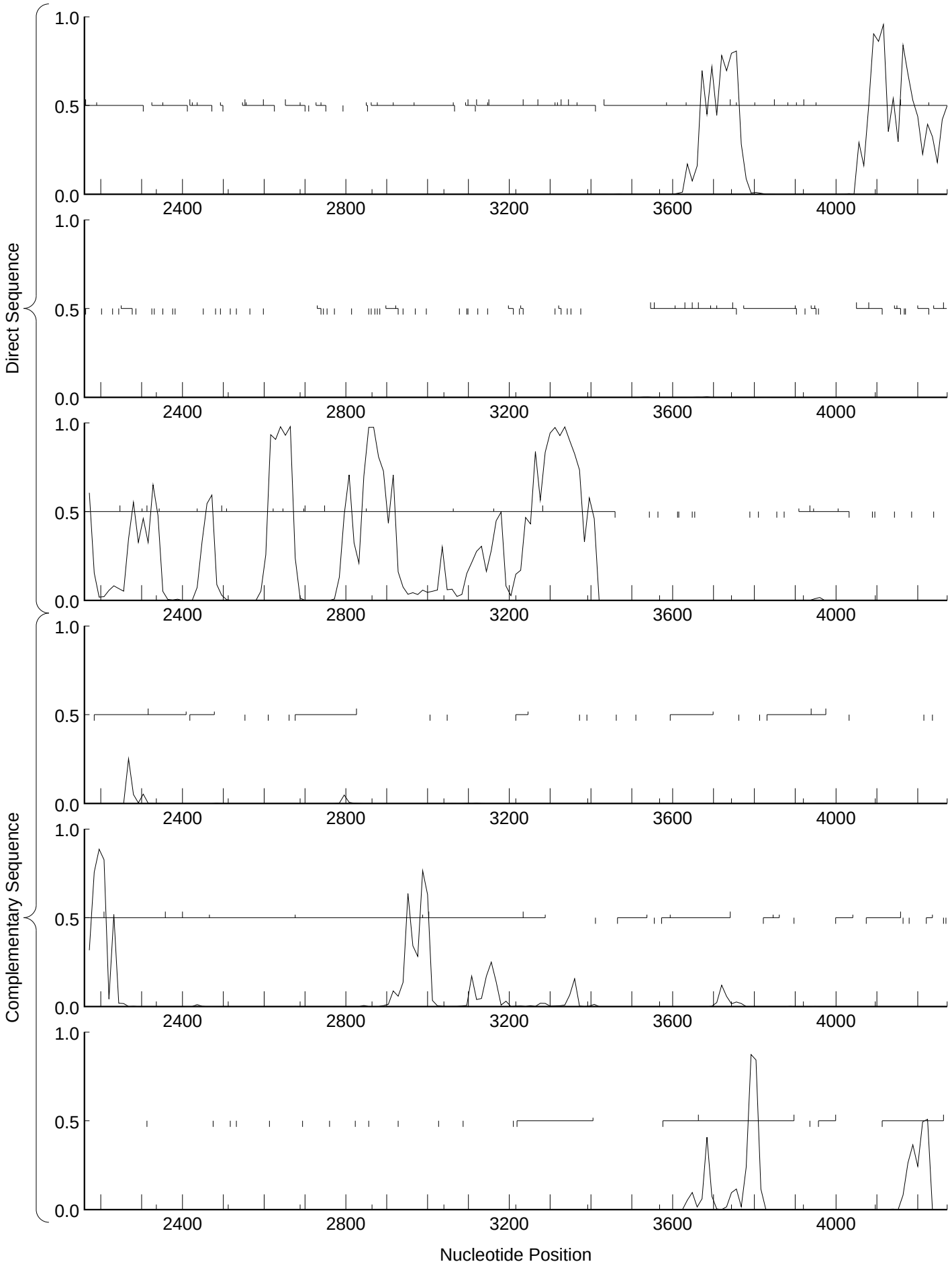

Send questions / comments to:
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Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

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

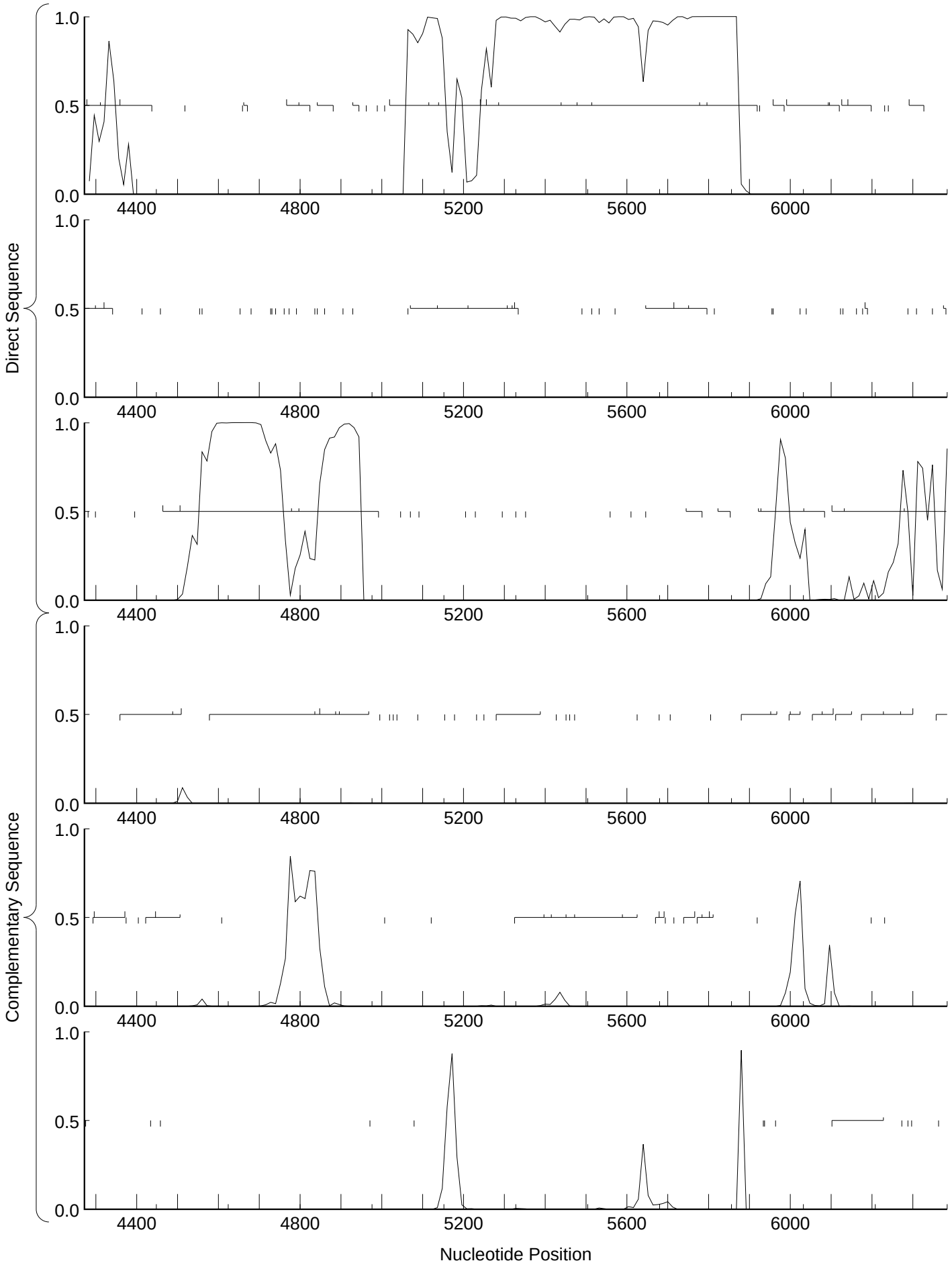
m25 complete sequence, 42695 bp including 9-base 3’ overhang (CGCCGGTGA), Cluster FQ, Order 4, Window 96, Step 12, 2/22



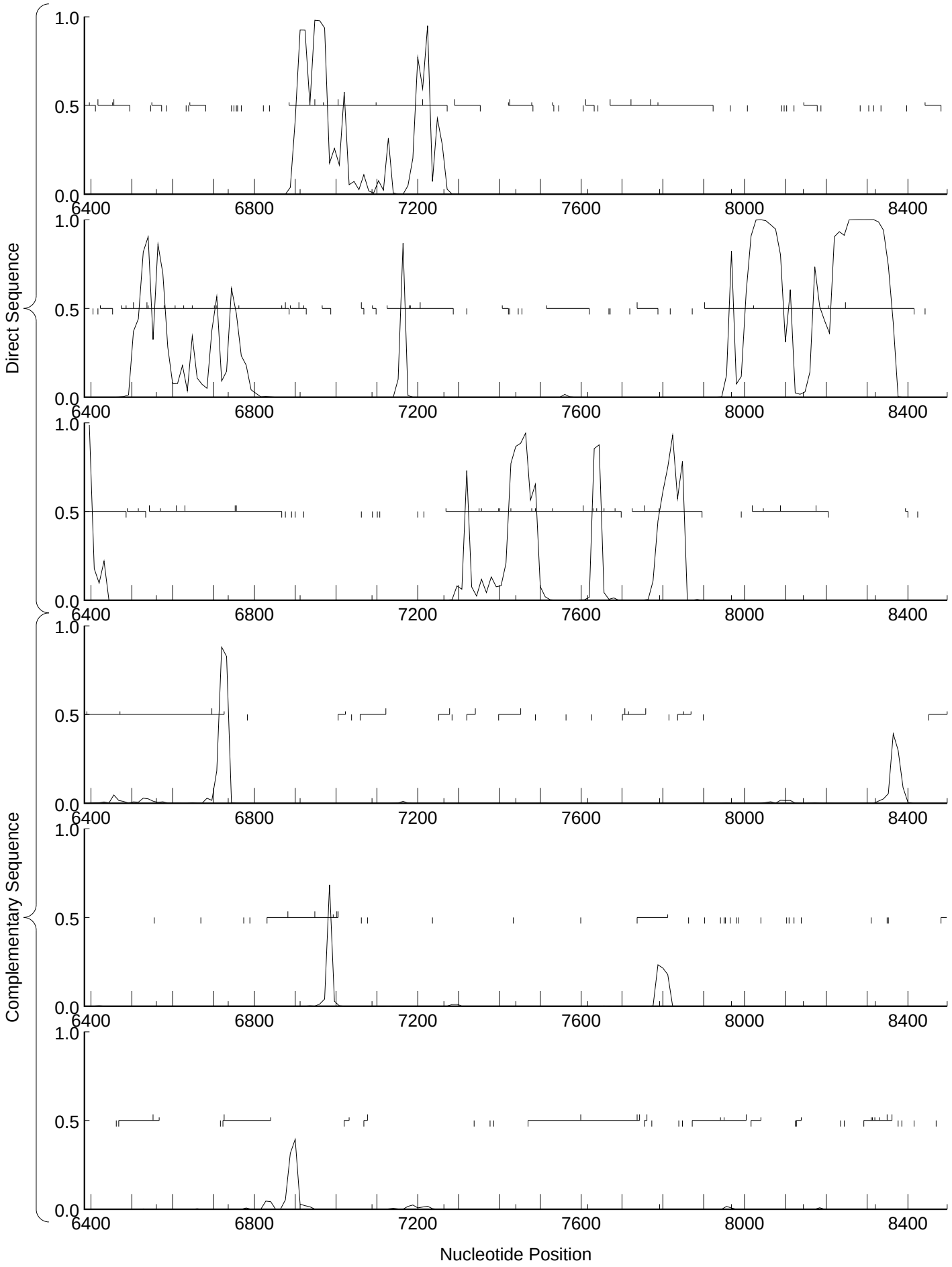
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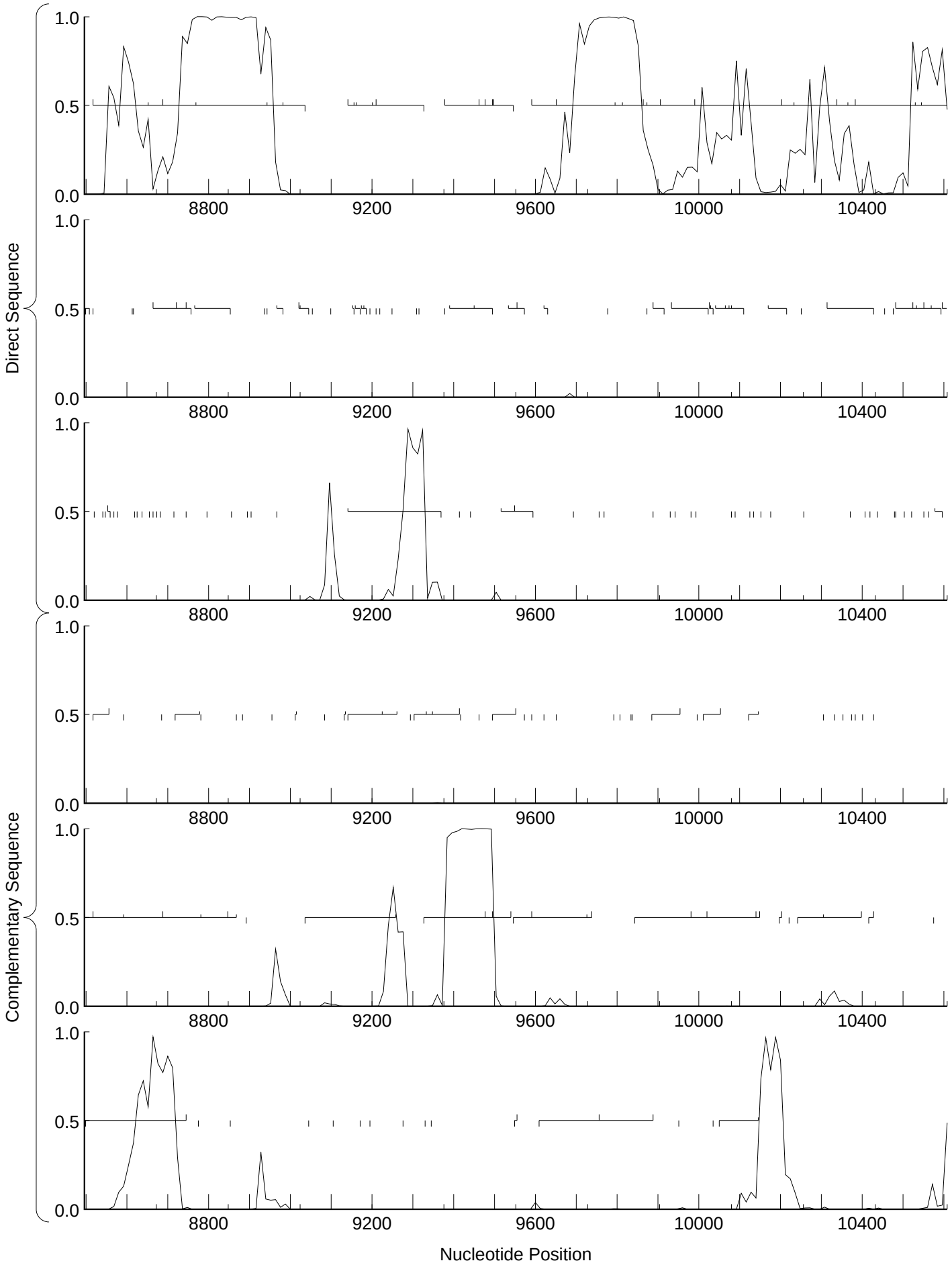
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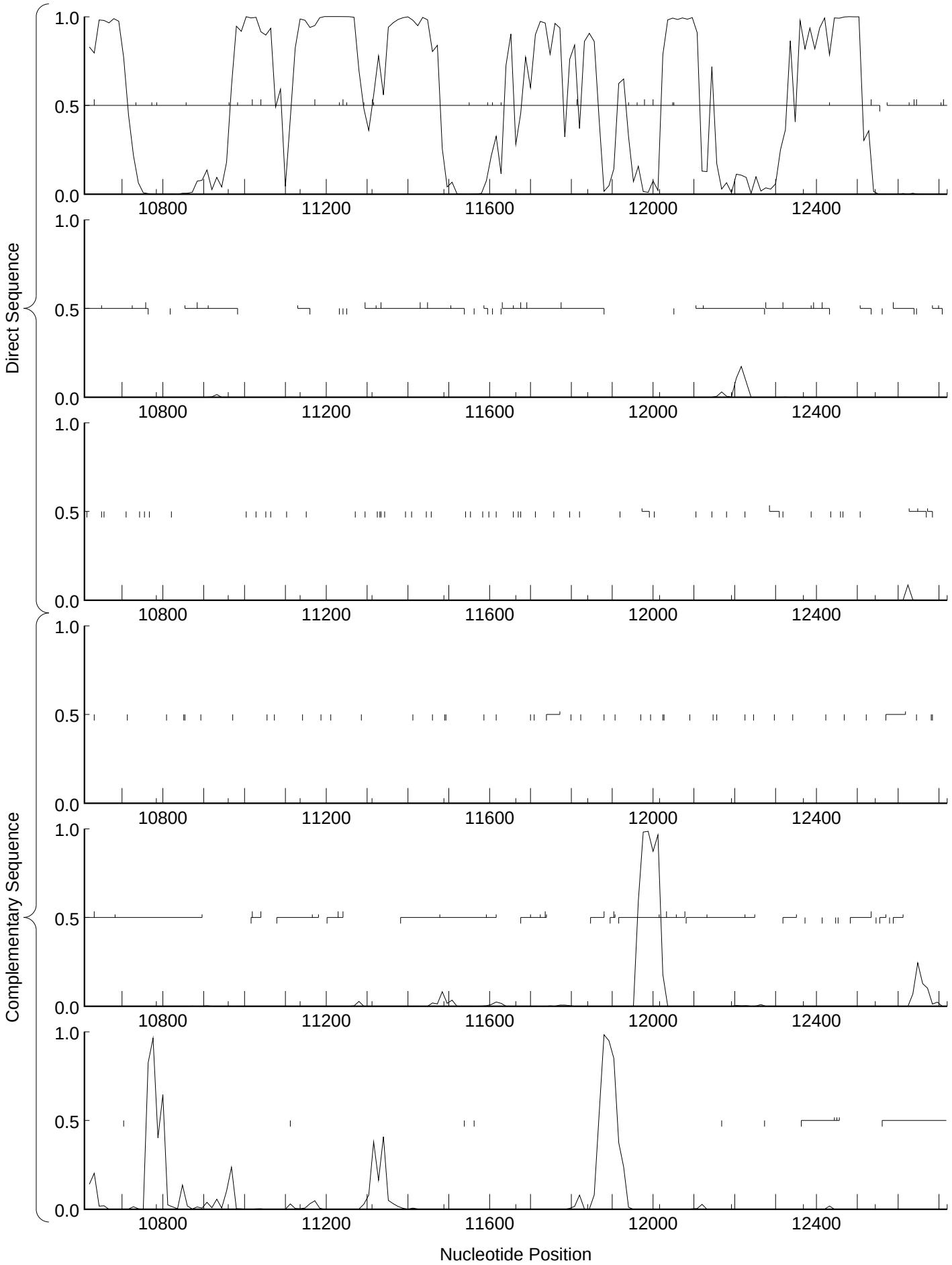
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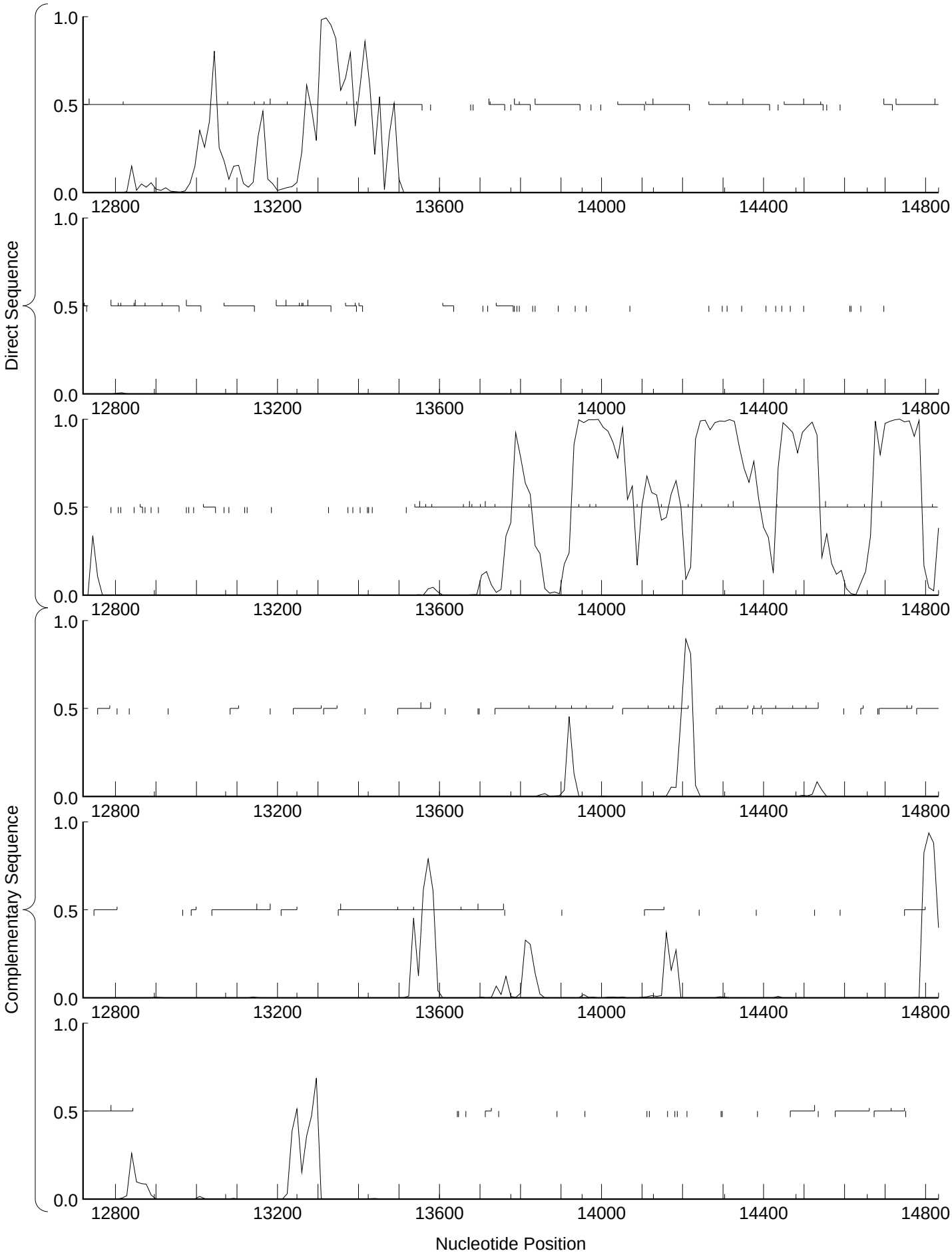
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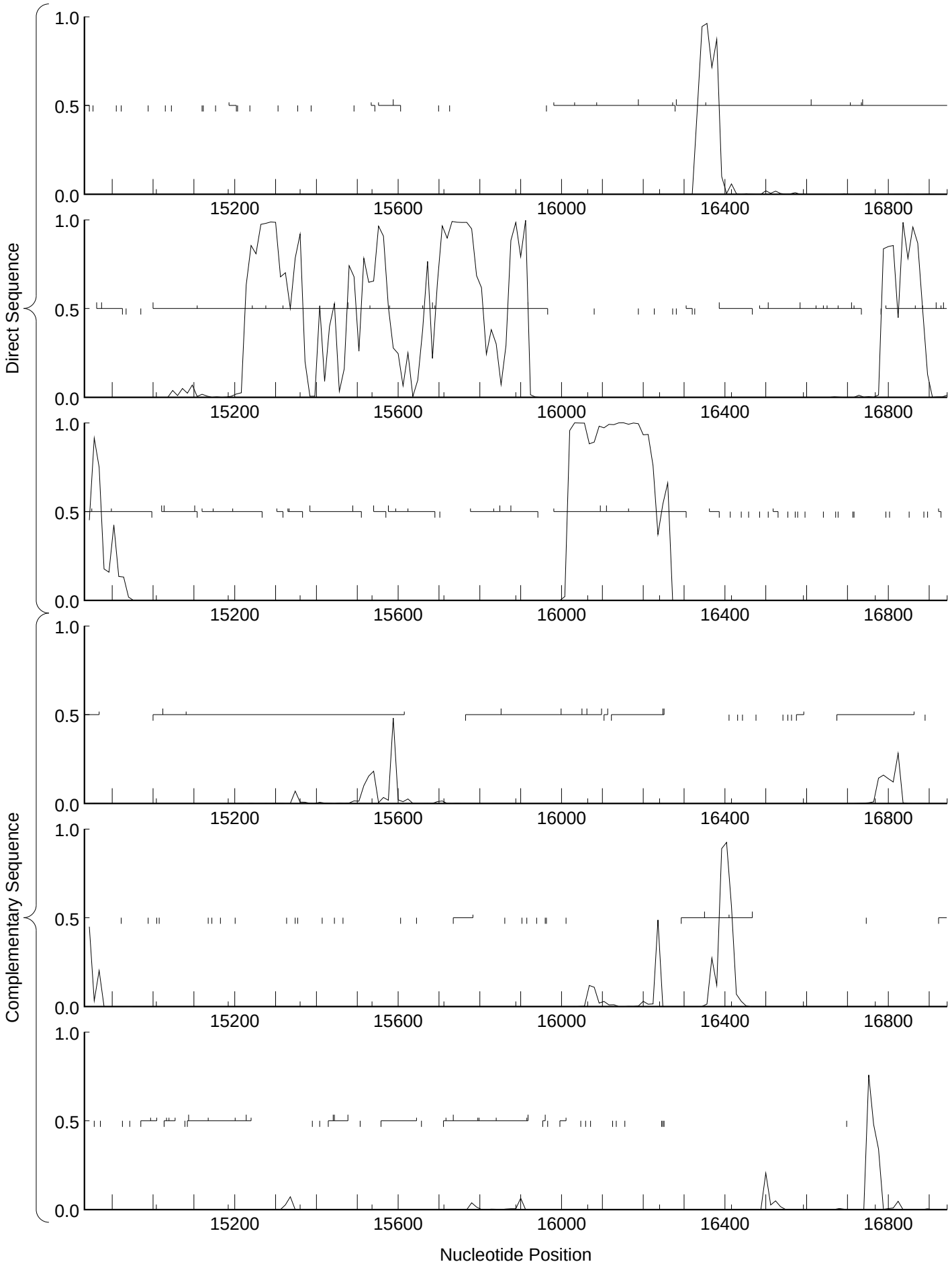
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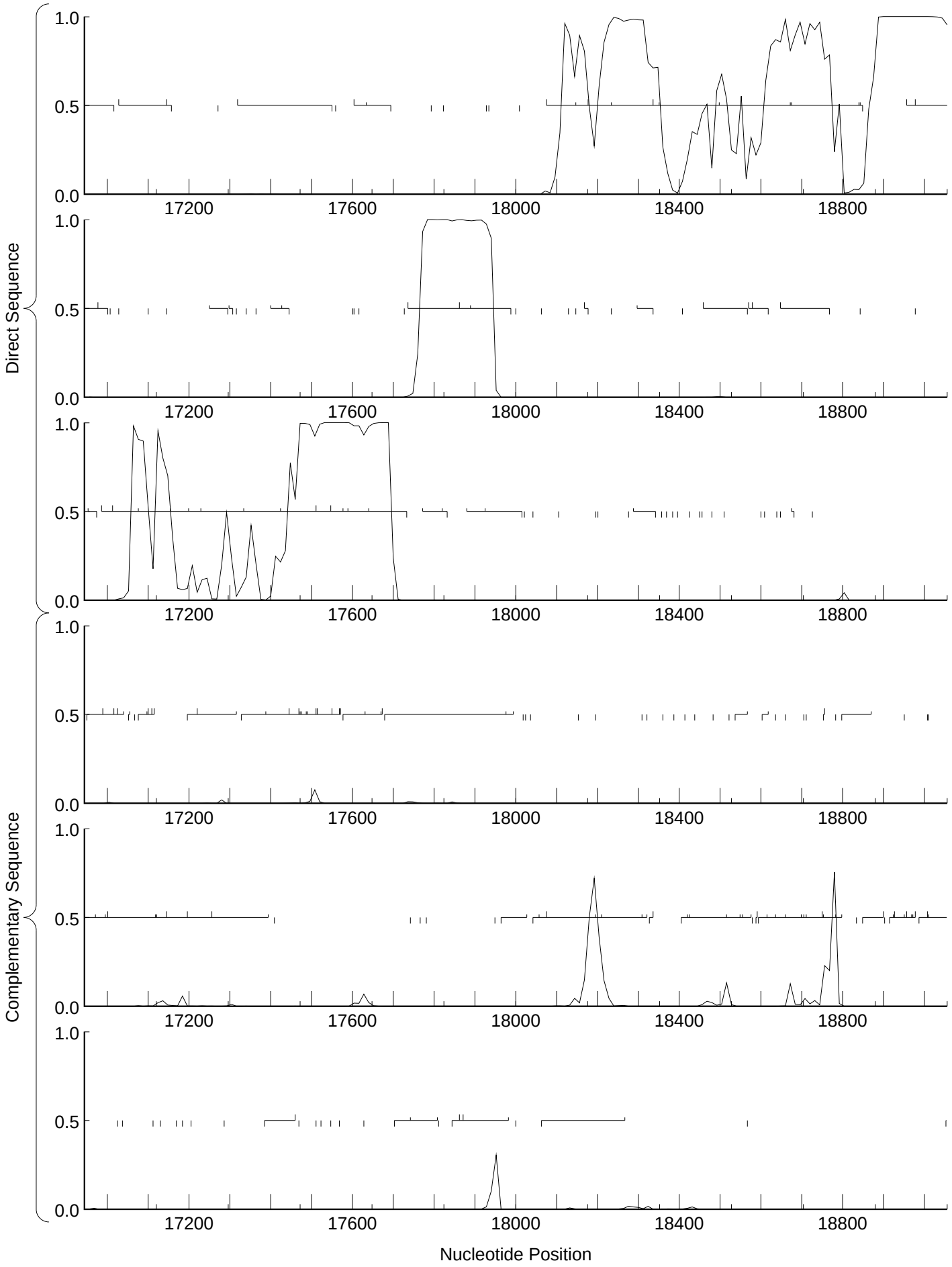
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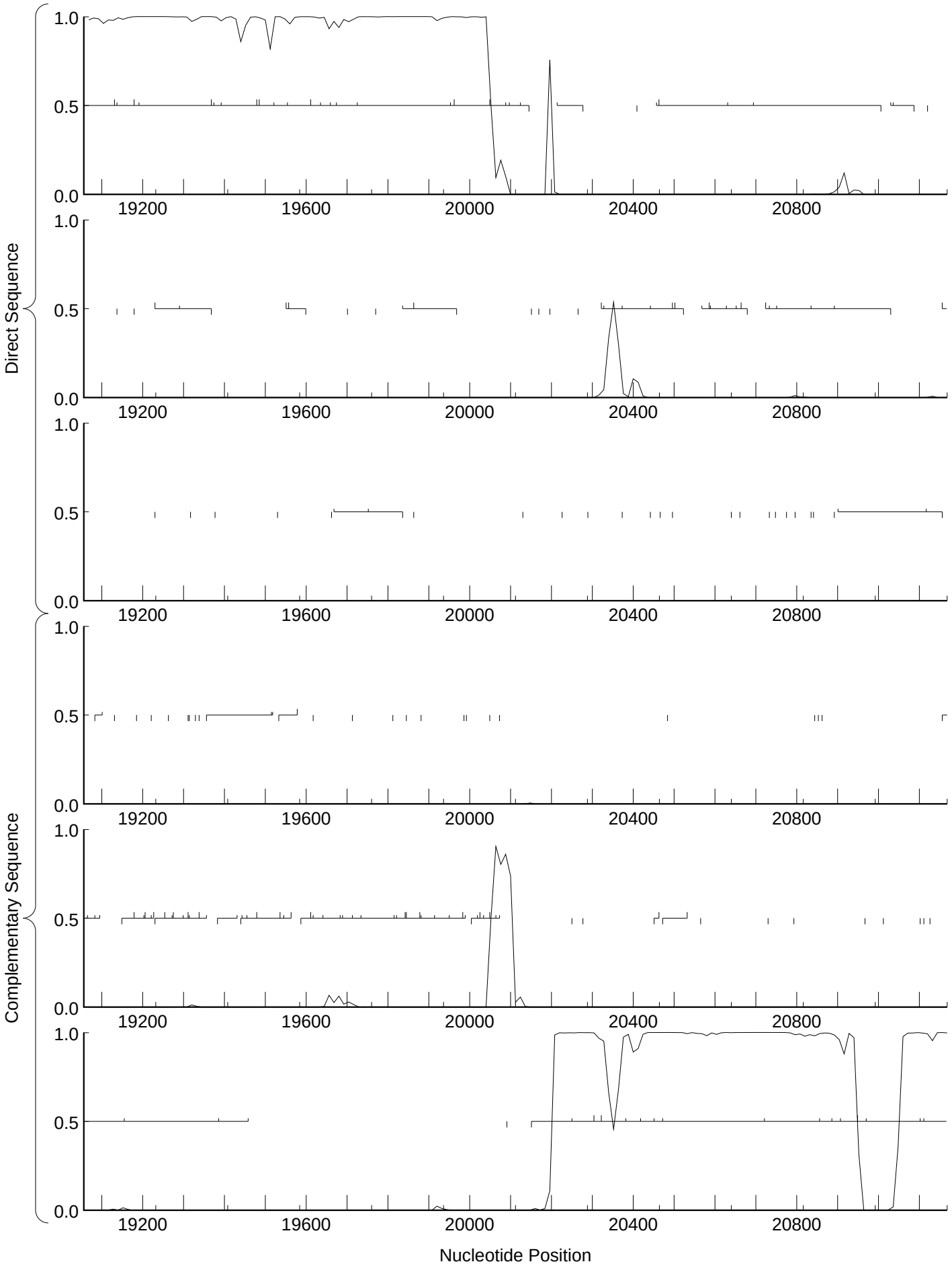
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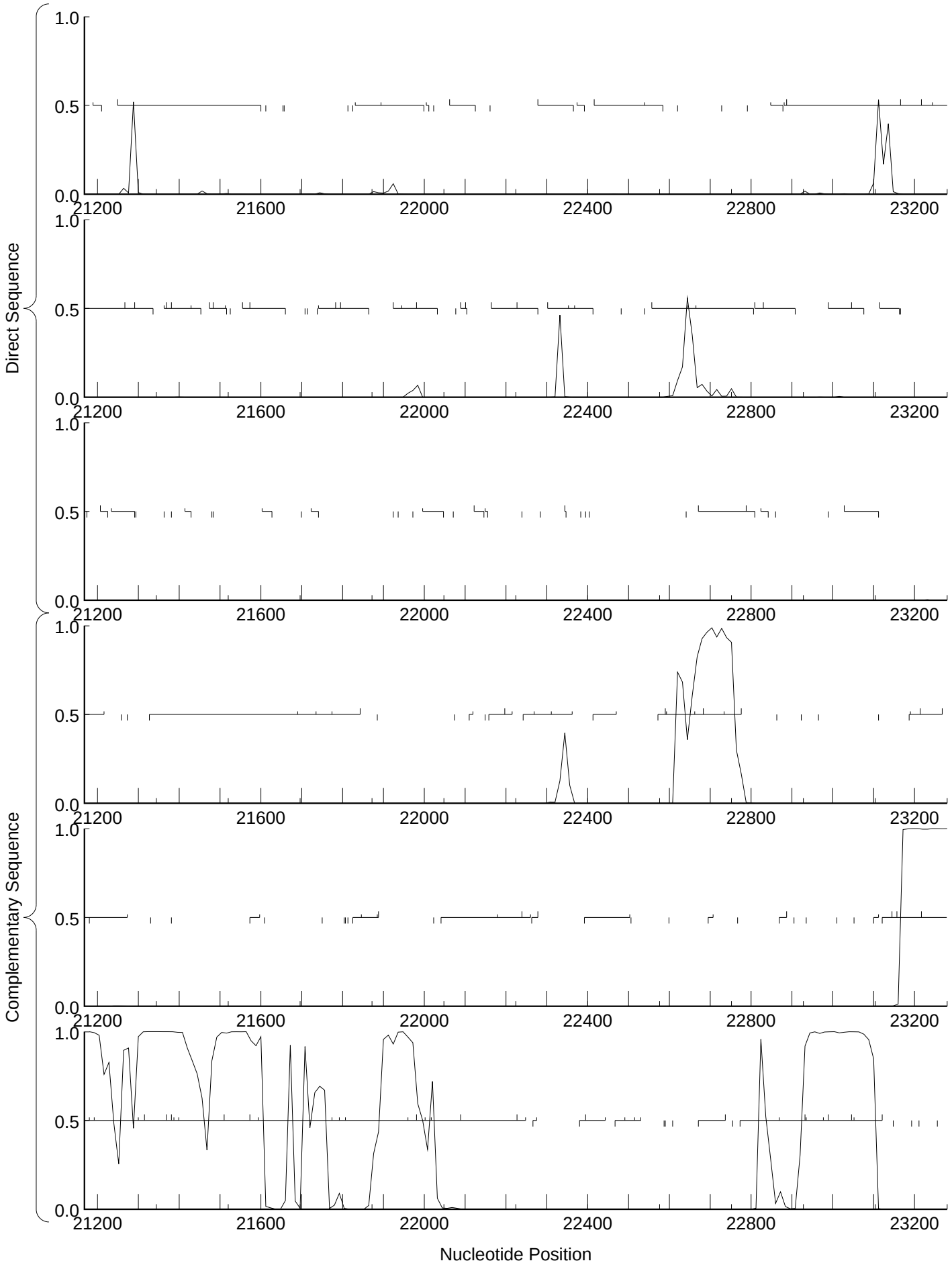
25 complete sequence, 42695 bp including 9-base 3' overhang (CGCCGGTGA), Cluster FQ, Order 4, Window 96, Step 12, 10/22



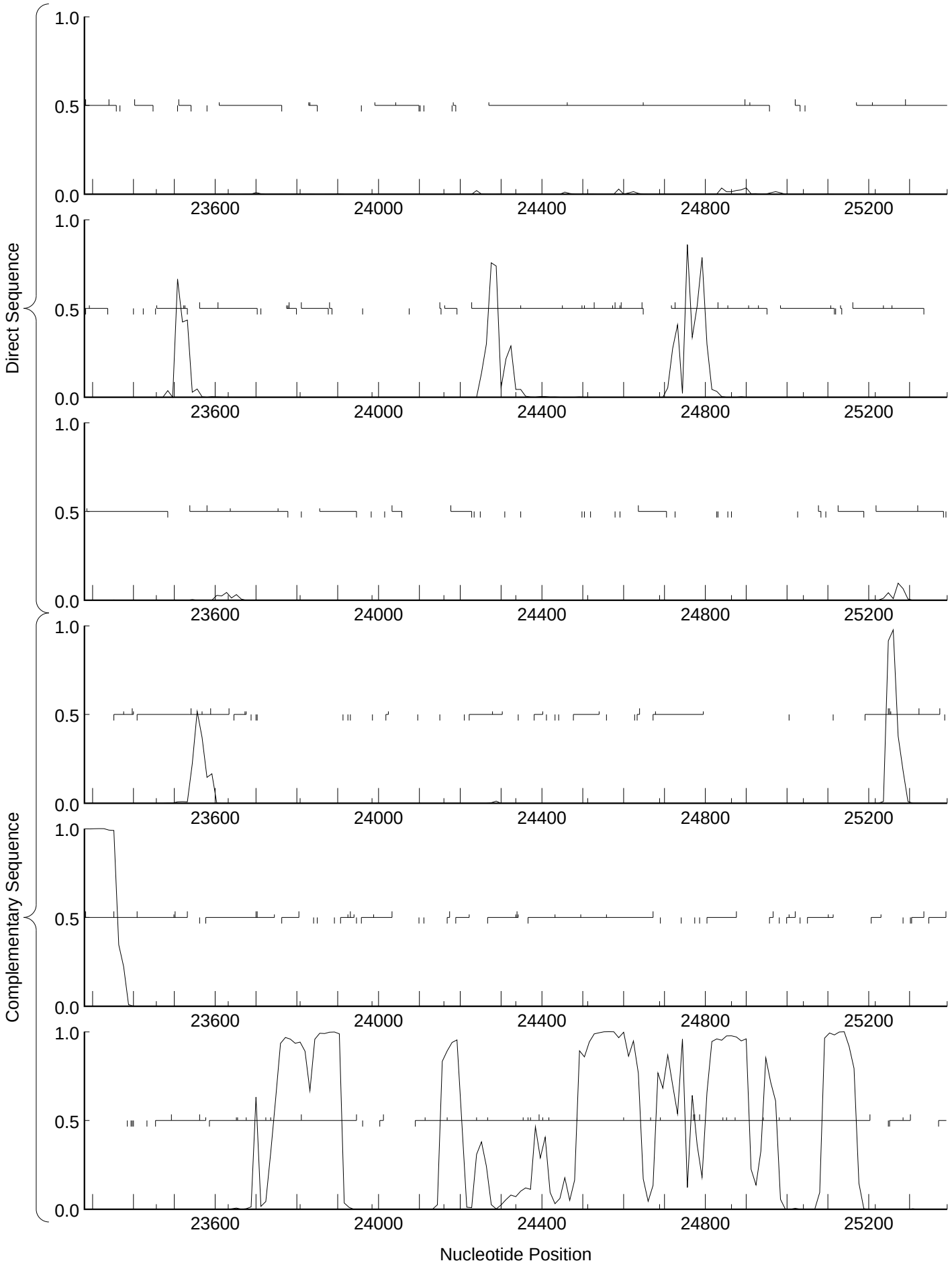
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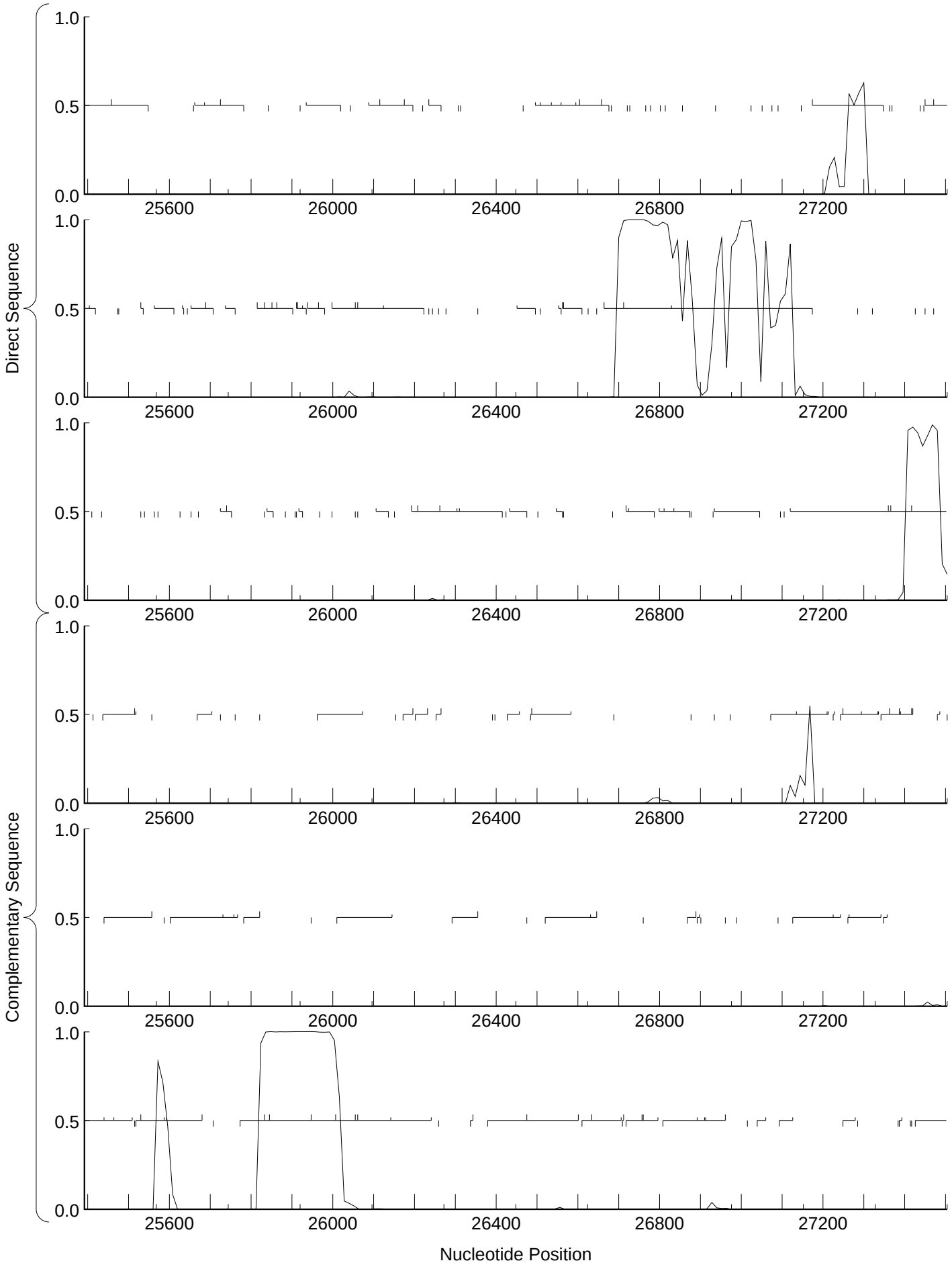


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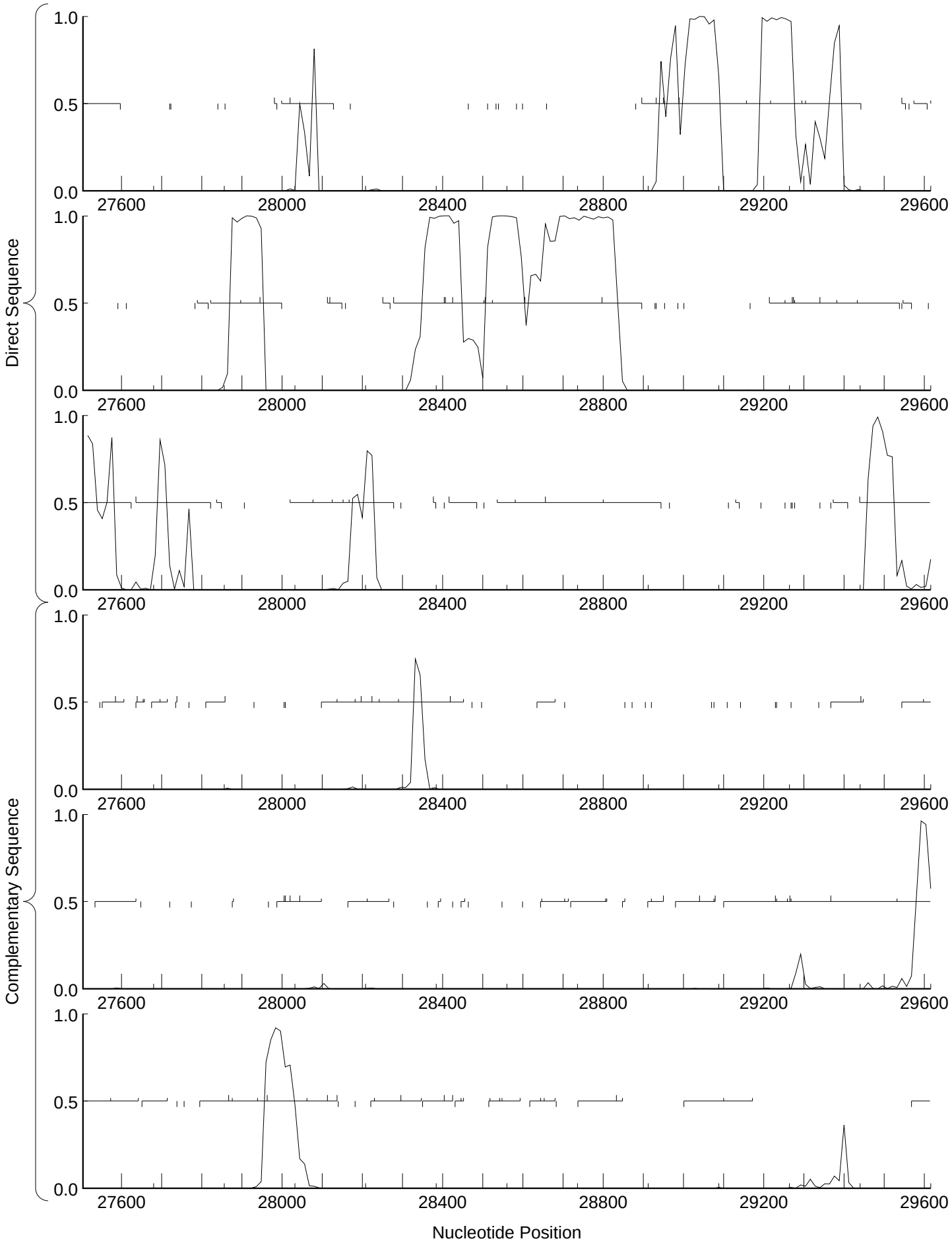


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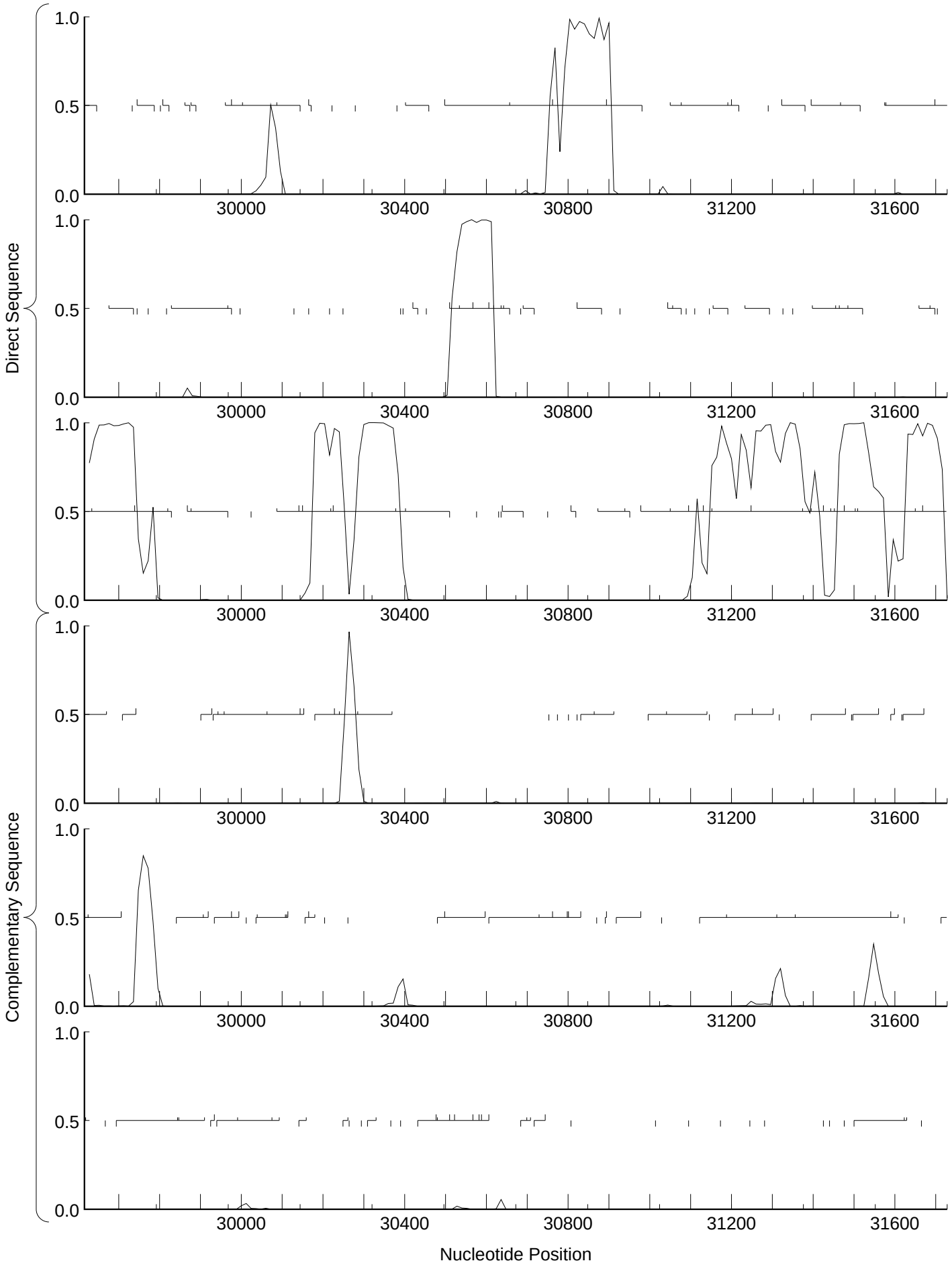



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
25600
26000
26400
26800
27200
Nucleotide Position

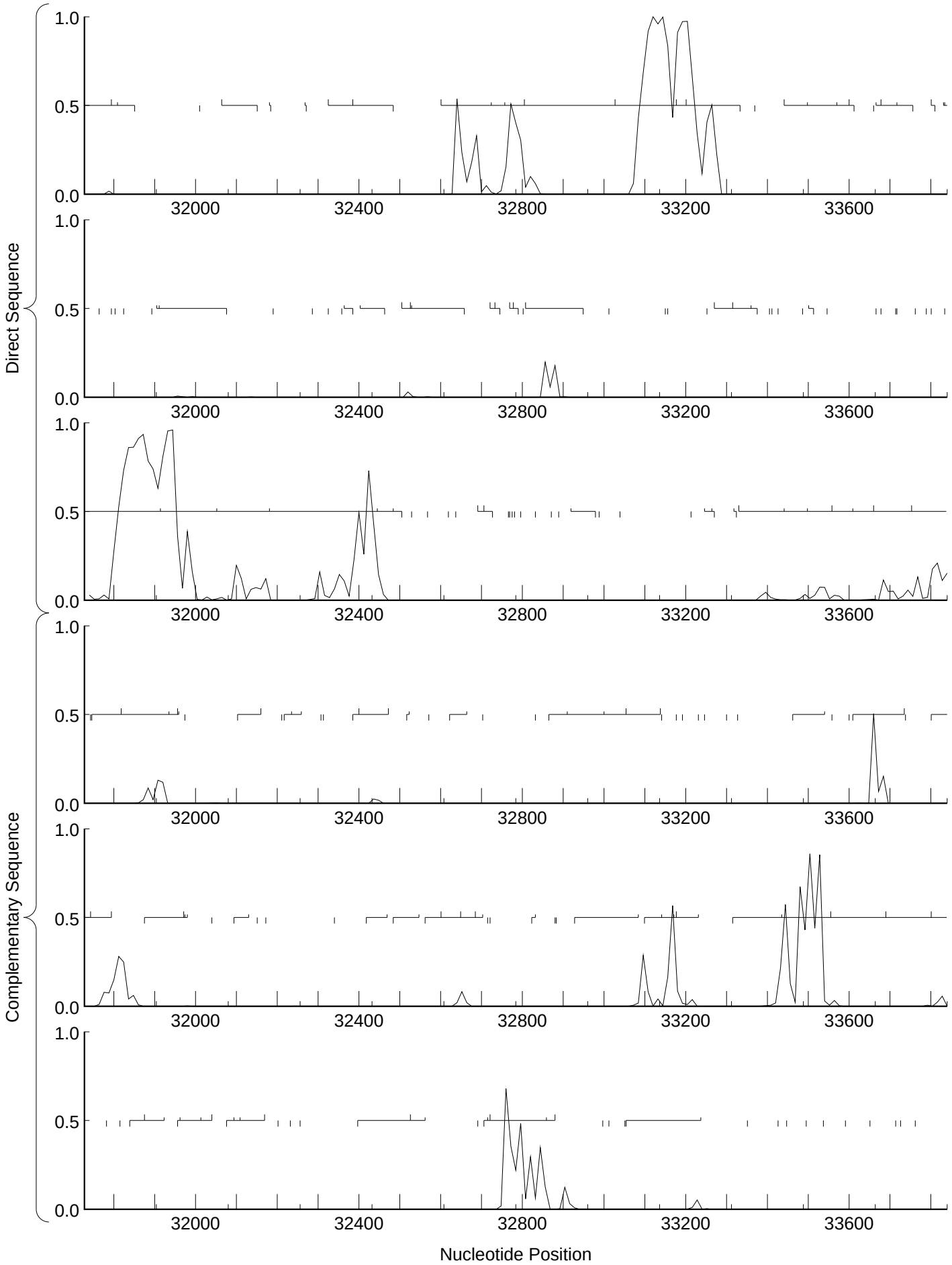
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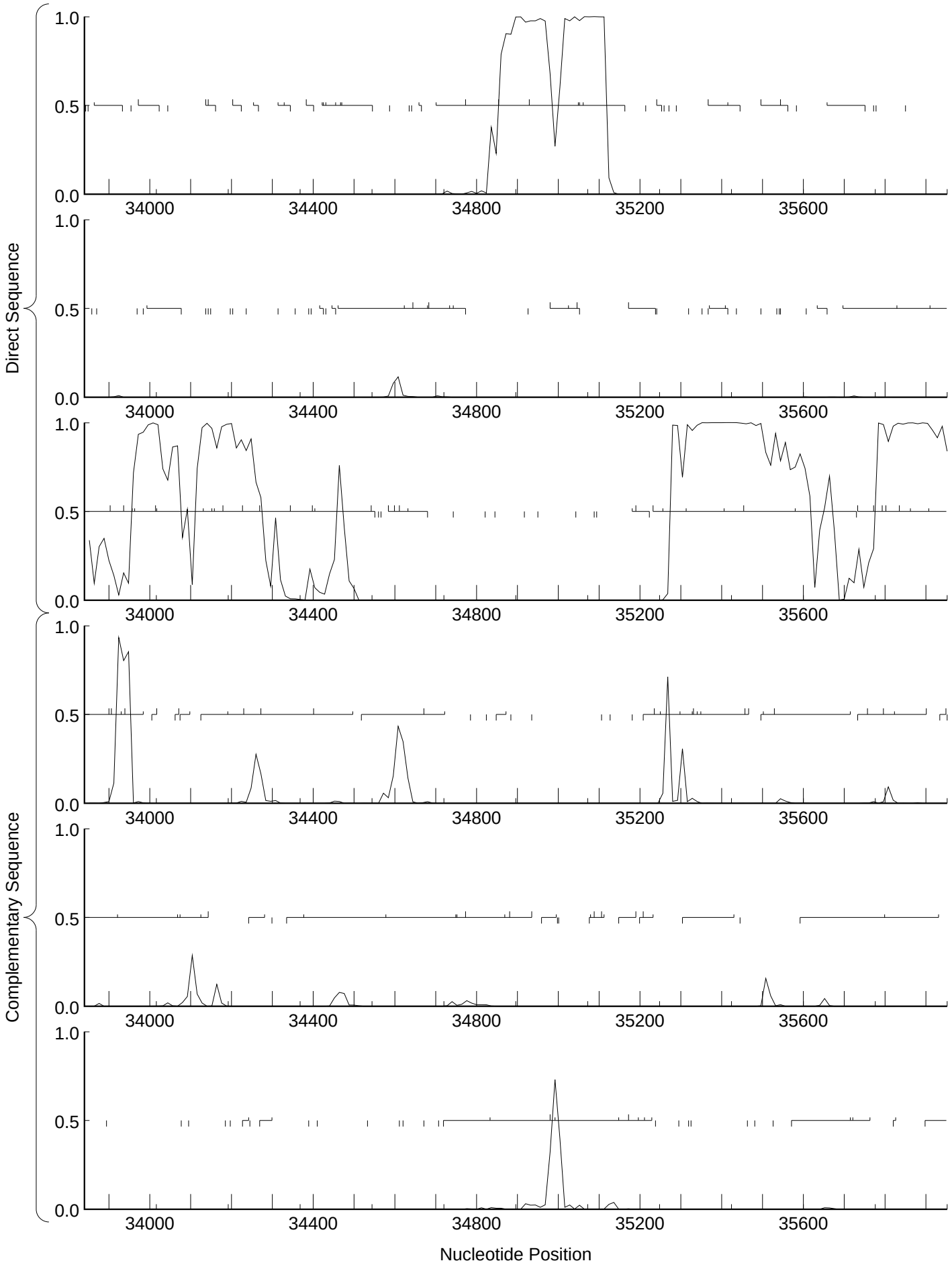


25 complete sequence, 42695 bp including 9-base 3' overhang (CGCCGGTGA), Cluster FQ, Order 4, Window 96, Step 12, 16/22

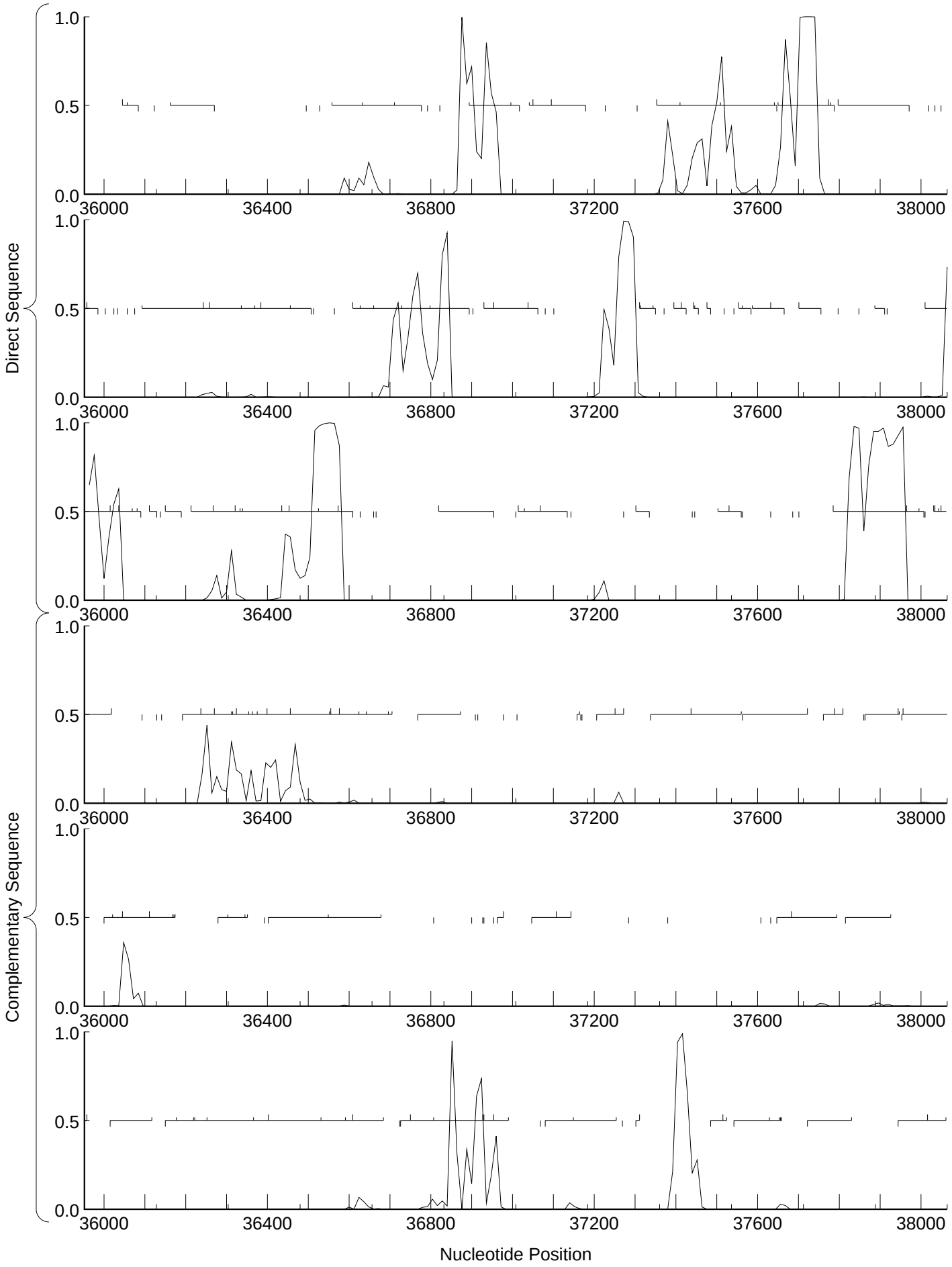


25 complete sequence, 42695 bp including 9-base 3' overhang (CGCCGGTGA), Cluster FQ, Order 4, Window 96, Step 12, 17/22

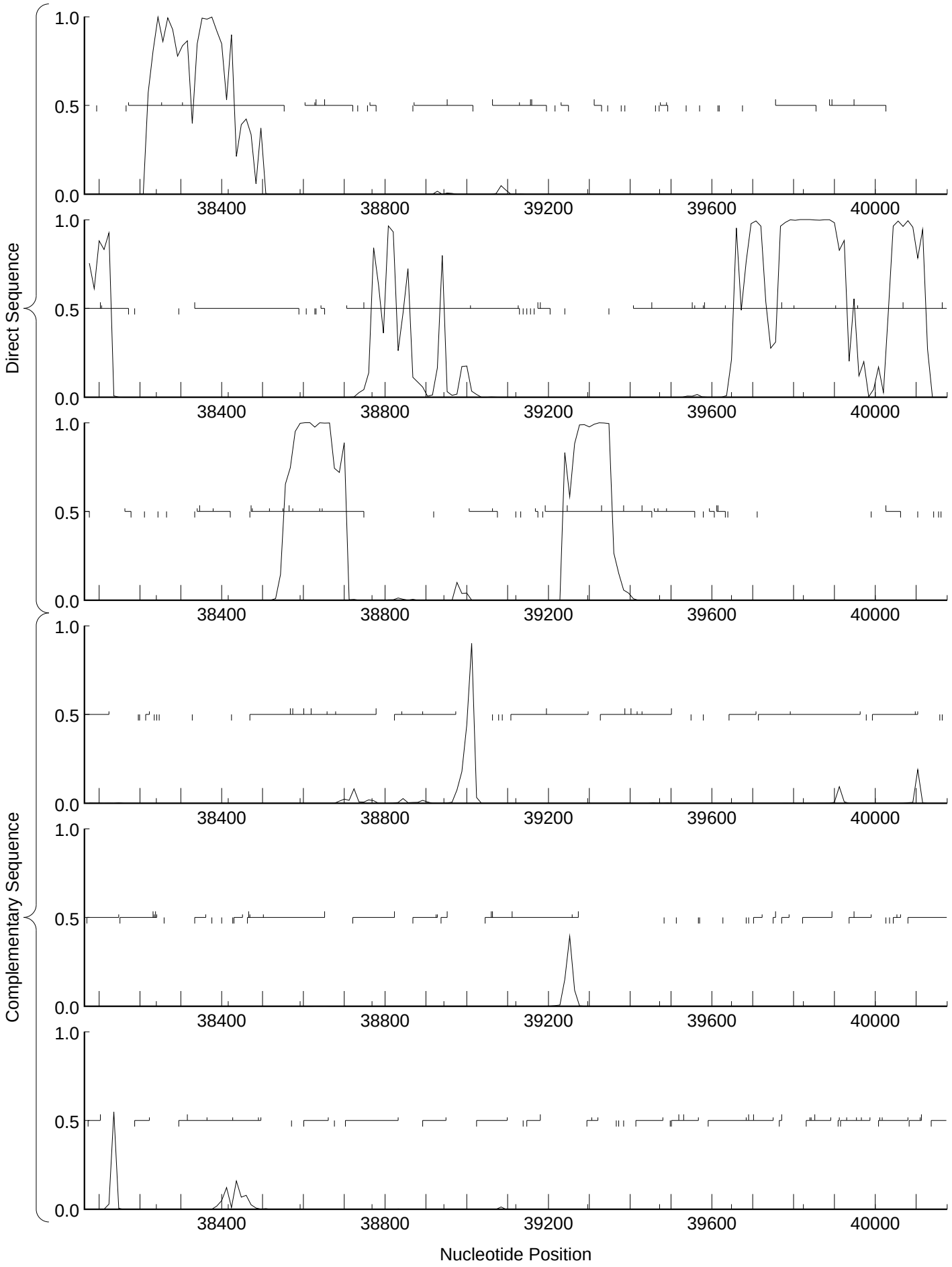



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

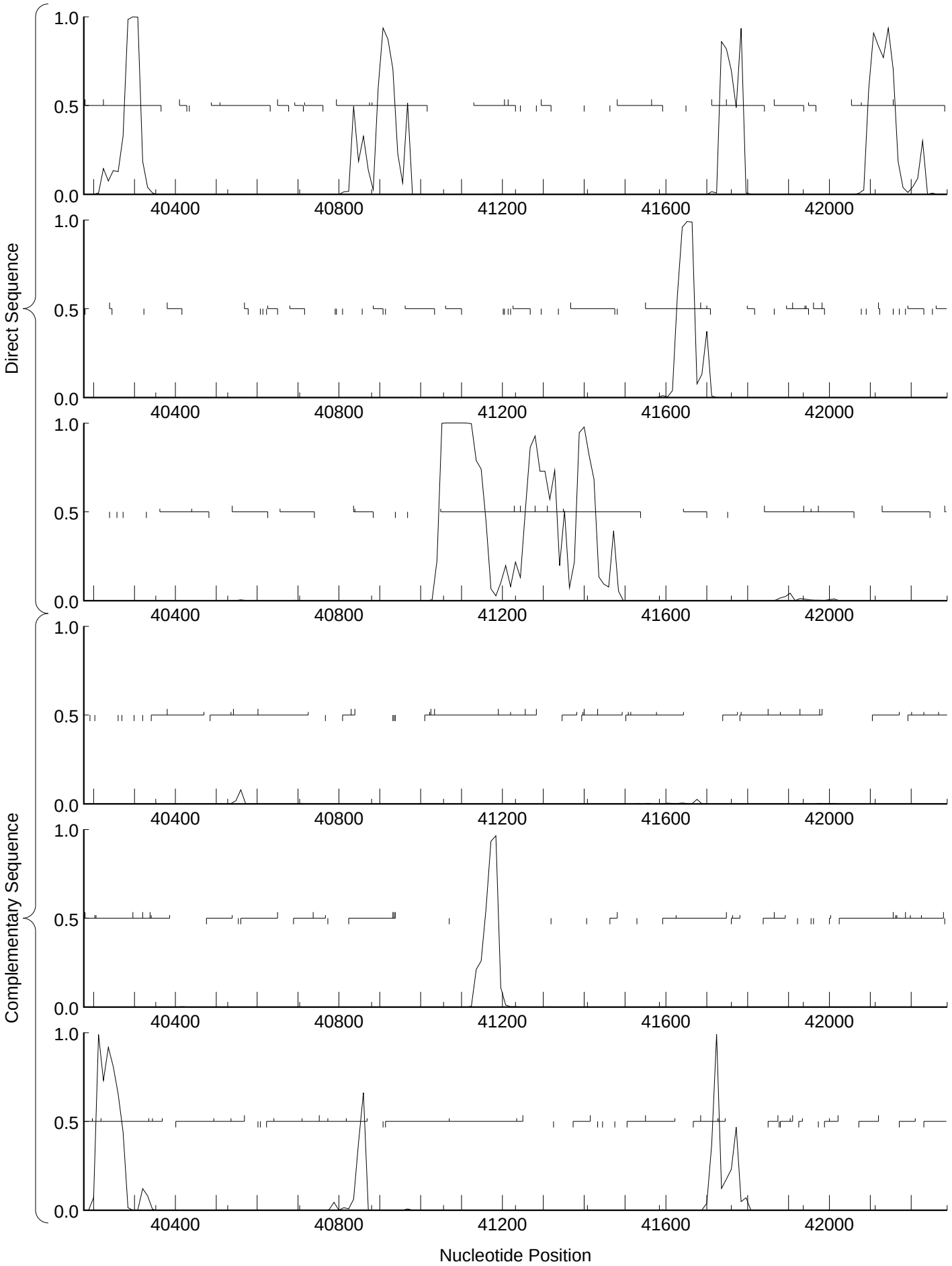
25 complete sequence, 42695 bp including 9-base 3' overhang (CGCCGGTGA), Cluster FQ, Order 4, Window 96, Step 12, 19/22



25 complete sequence, 42695 bp including 9-base 3’ overhang (CGCCGGTGA), Cluster FQ, Order 4, Window 96, Step 12, 20/22



25 complete sequence, 42695 bp including 9-base 3' overhang (CGCCGGTGA), Cluster FQ, Order 4, Window 96, Step 12, 21/22



25 complete sequence, 42695 bp including 9-base 3’ overhang (CGCCGGTGA), Cluster FQ, Order 4, Window 96, Step 12, 22/22

