GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Kimona complete sequence, 50283 bp including 10-base 3' overhang (CGGGC
Analysis Date    :  3/28/24 at 20:13:34
Pages            :  25
Sequence Length  :  50283 bp
GC Content       :   64.38%

Window Length    :  96 bp
Window Step      :  12 bp
Threshold Value  :  0.500

PS-Version       :  1.2

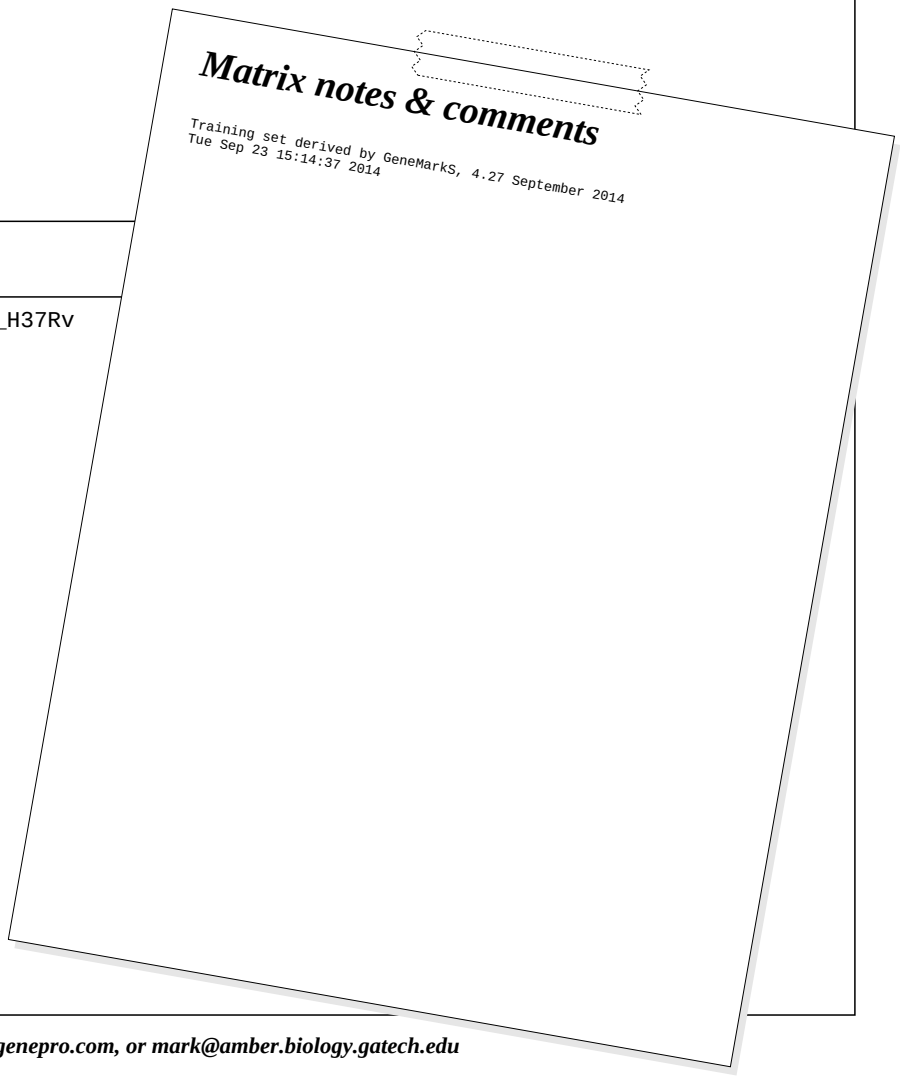
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Author     :  -

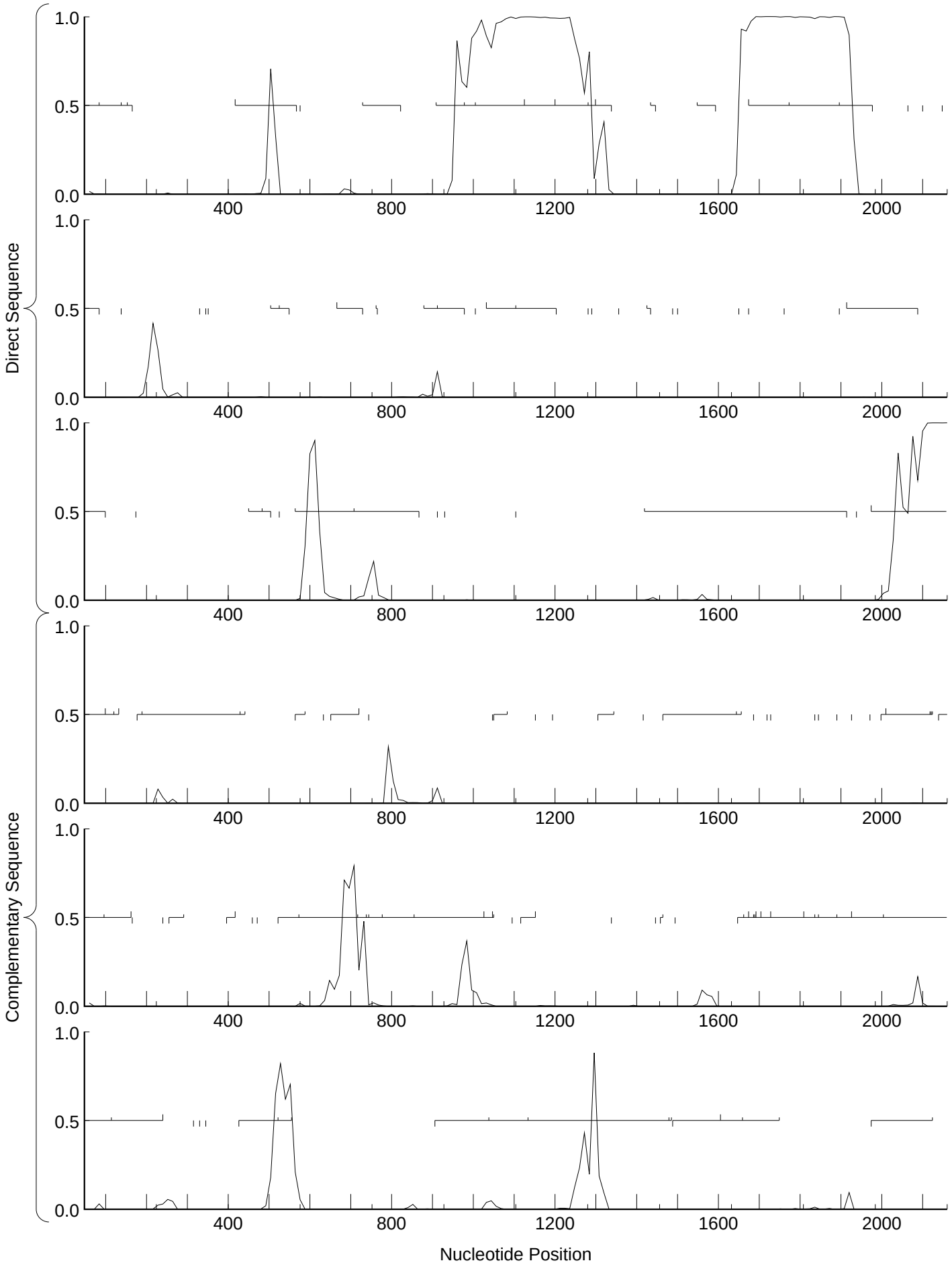
Order      :  4


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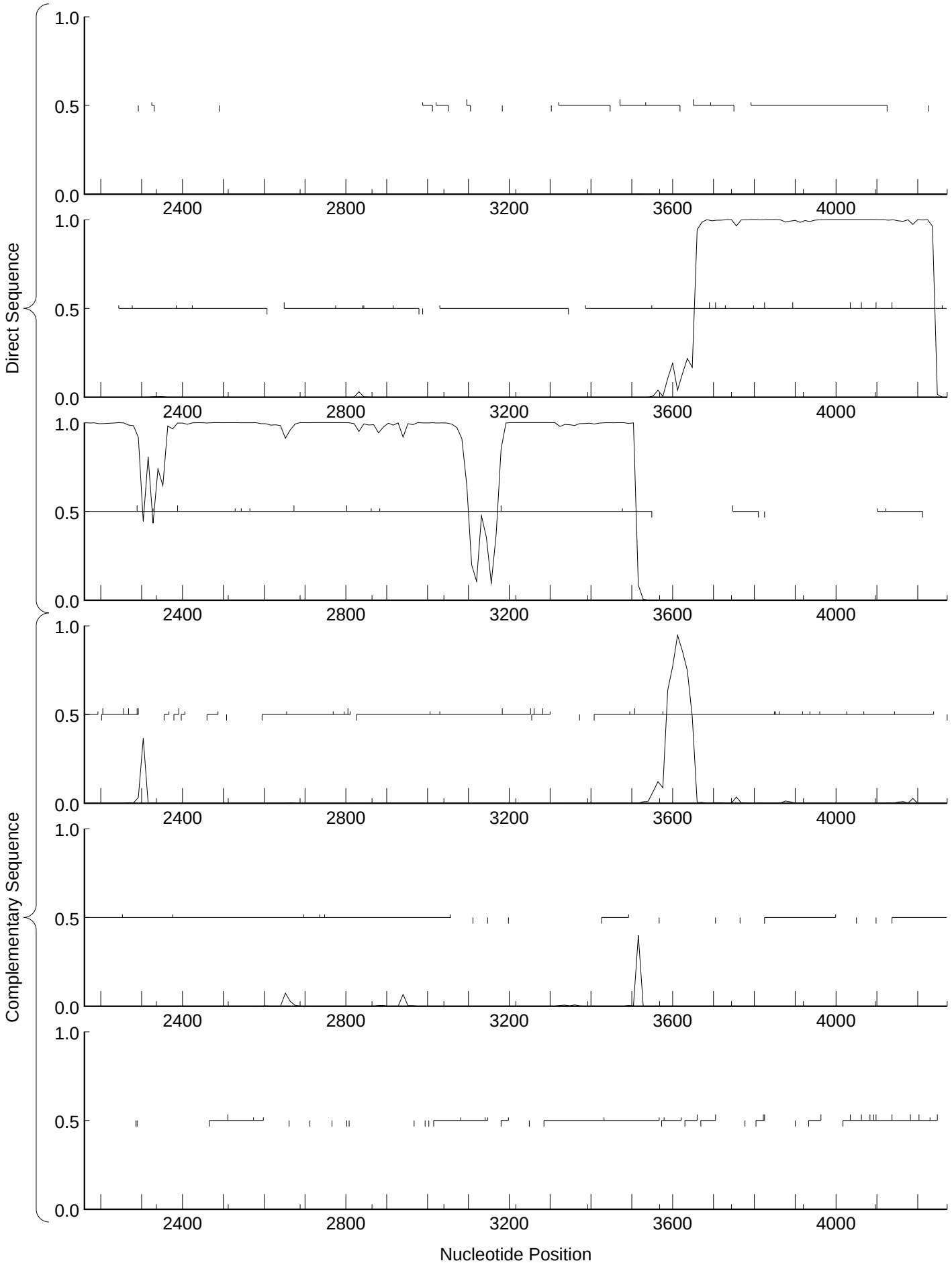


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

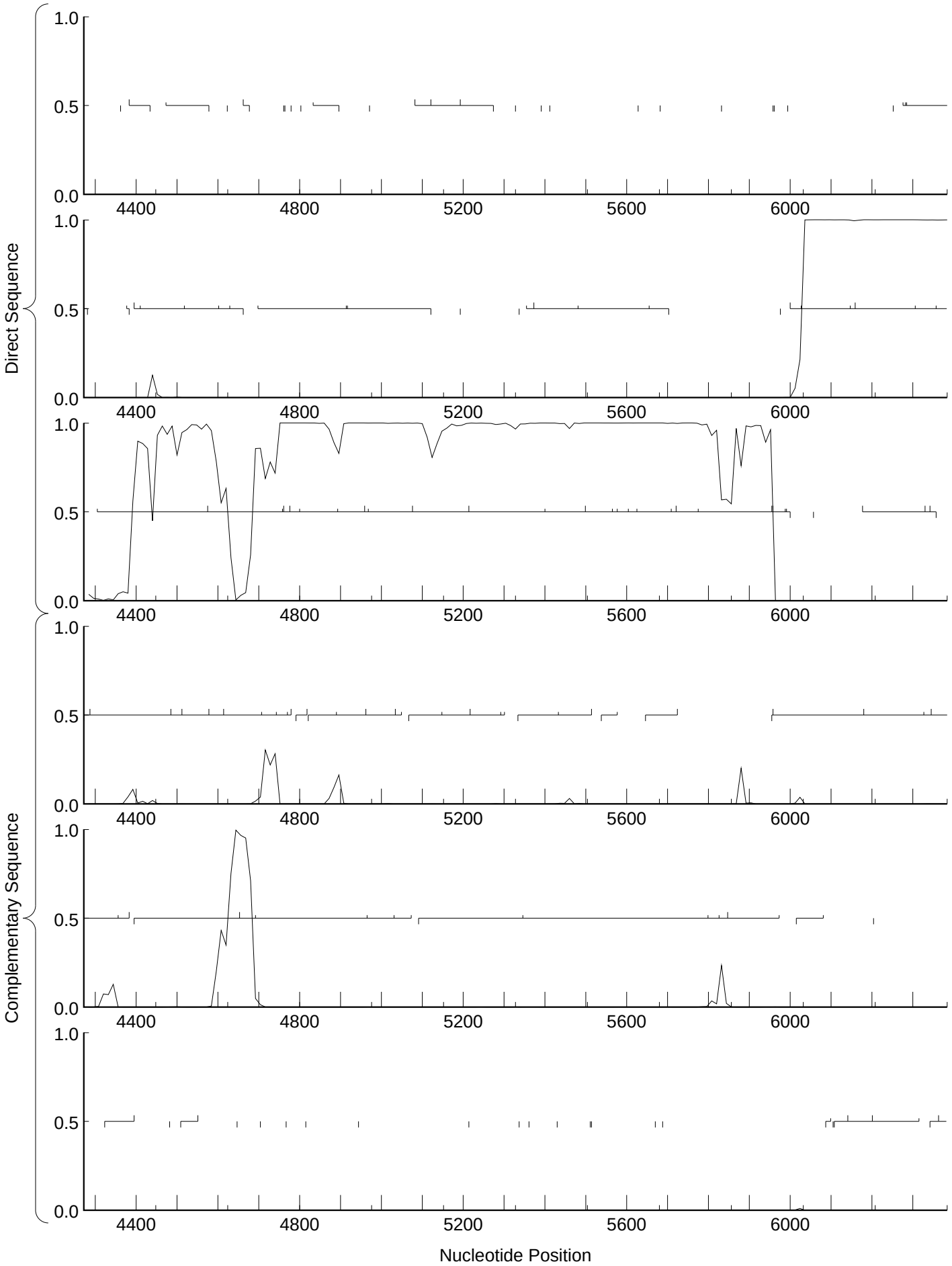
complete sequence, 50283 bp including 10-base 3’ overhang (CGGGCGGTAA), Cluster A19, Order 4, Window 96, Step 12, 2/25



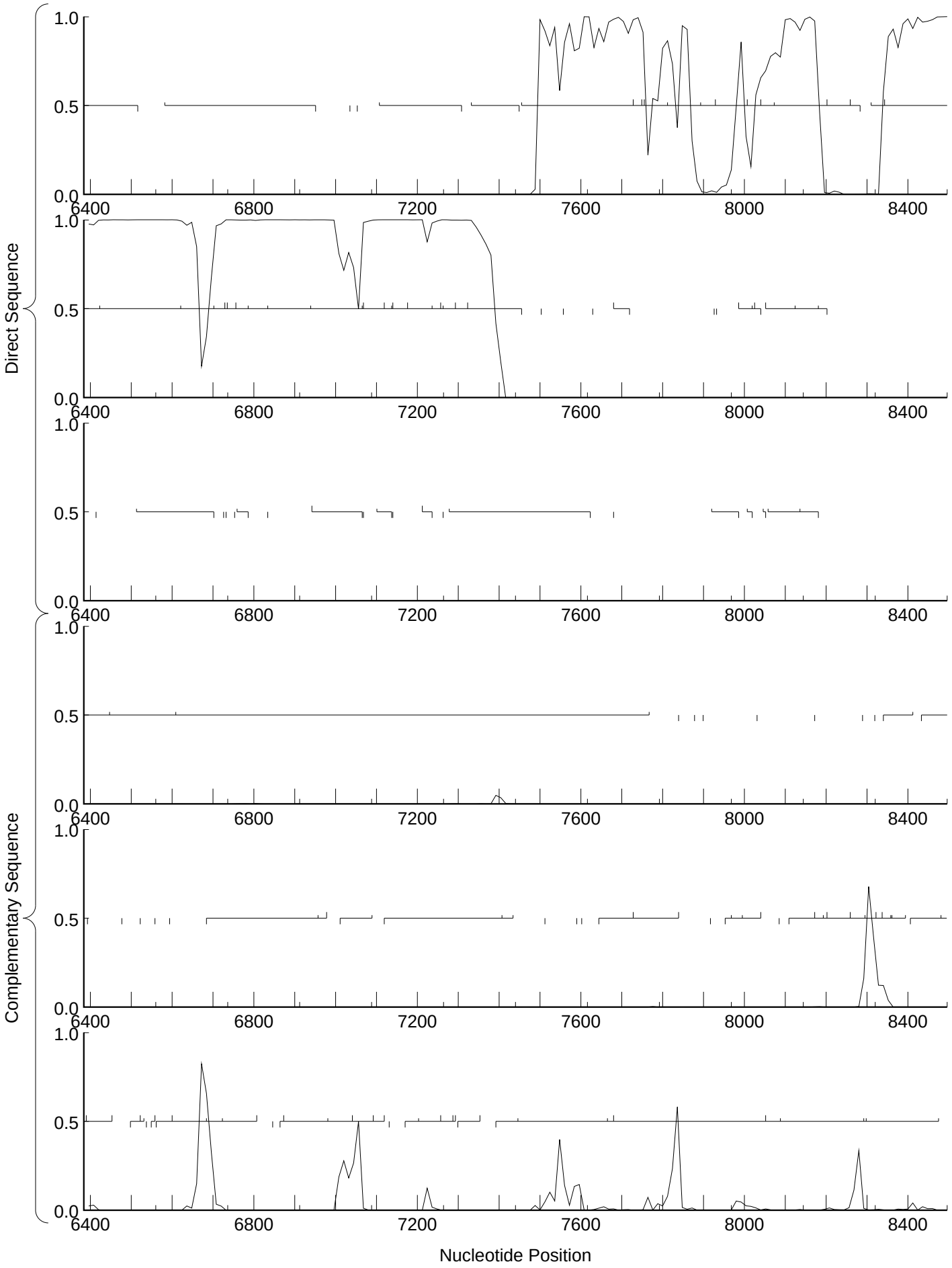
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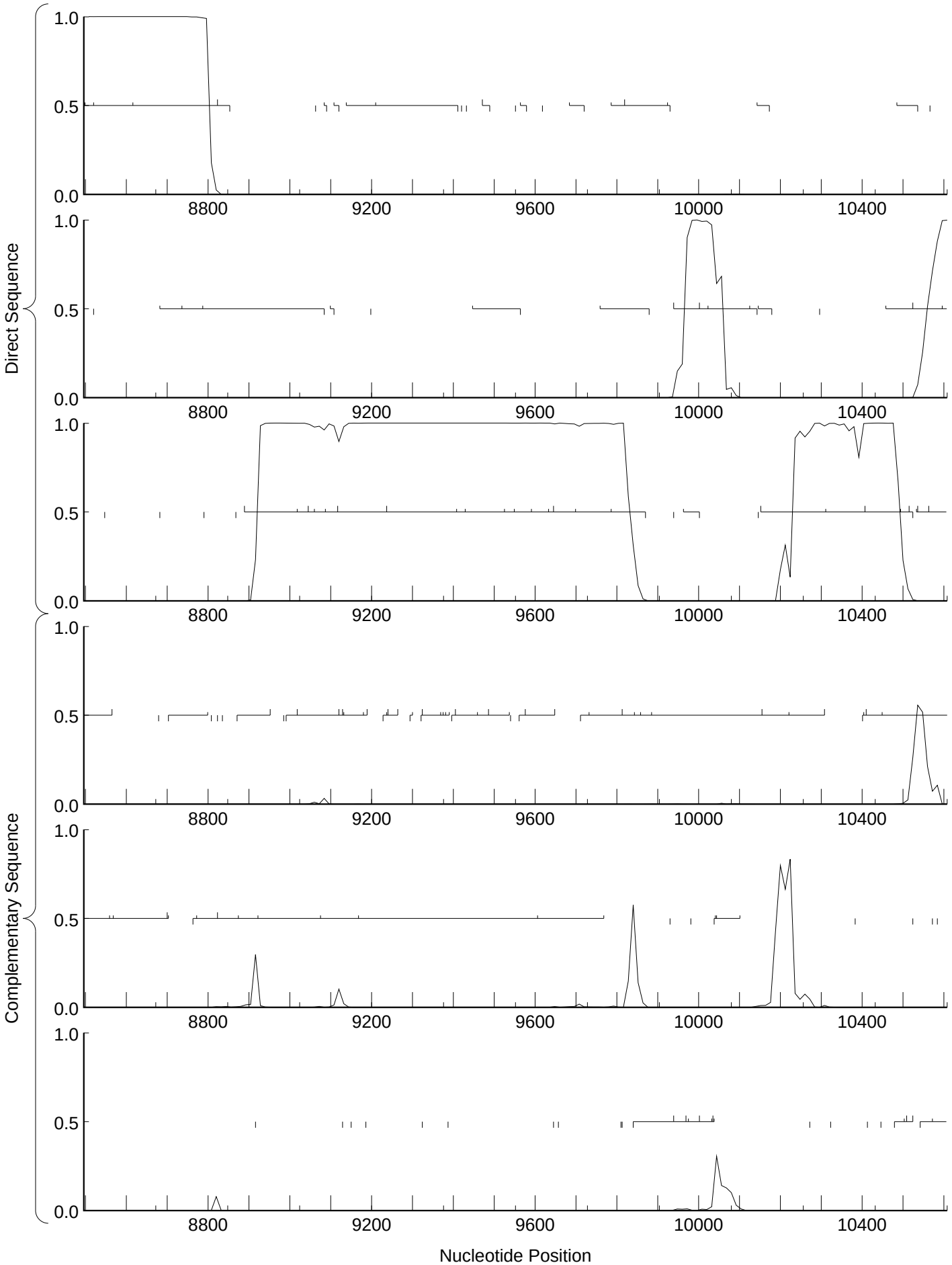
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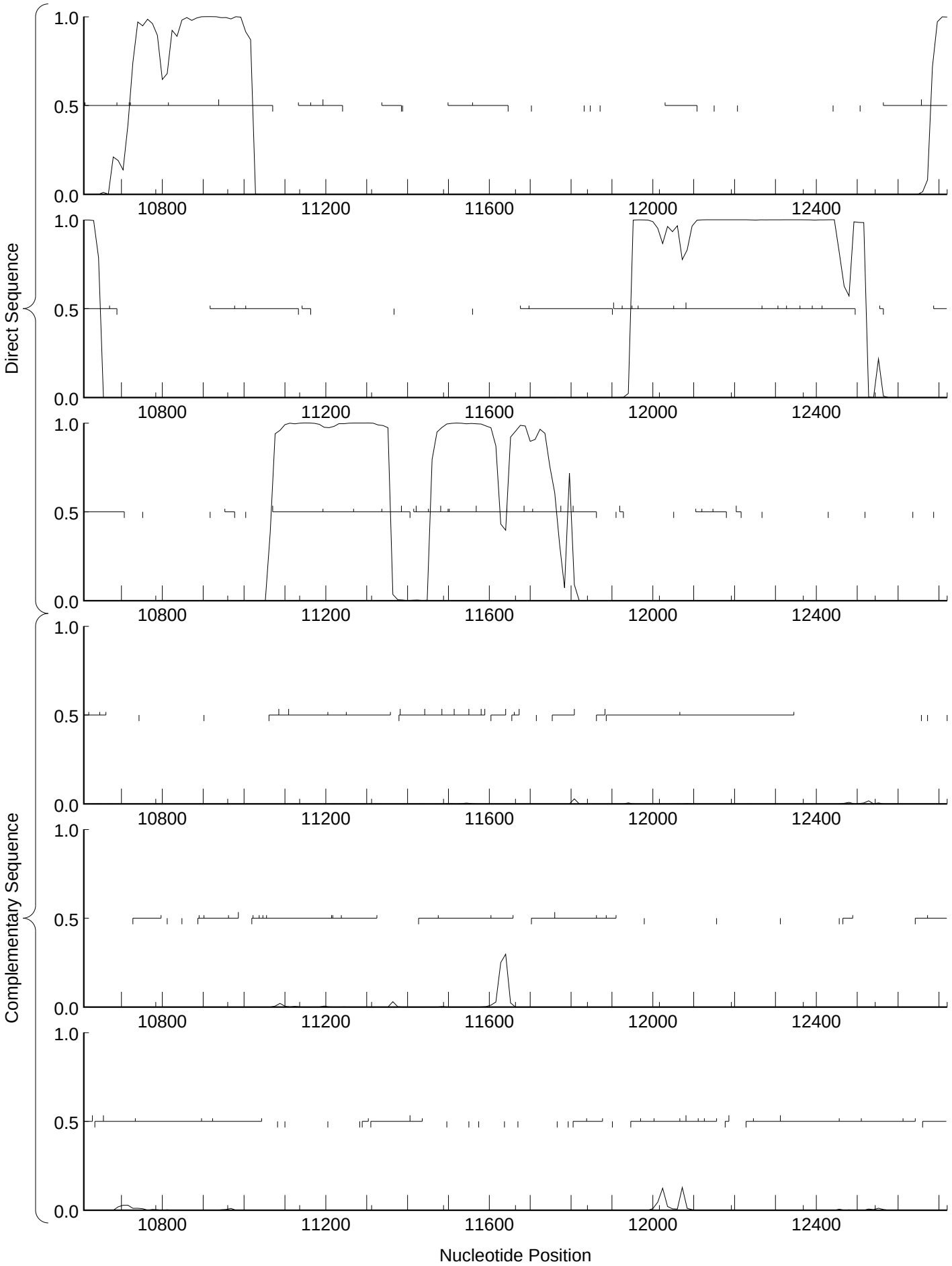
complete sequence, 50283 bp including 10-base 3’ overhang (CGGGCGGTAA), Cluster A19, Order 4, Window 96, Step 12, 5/25



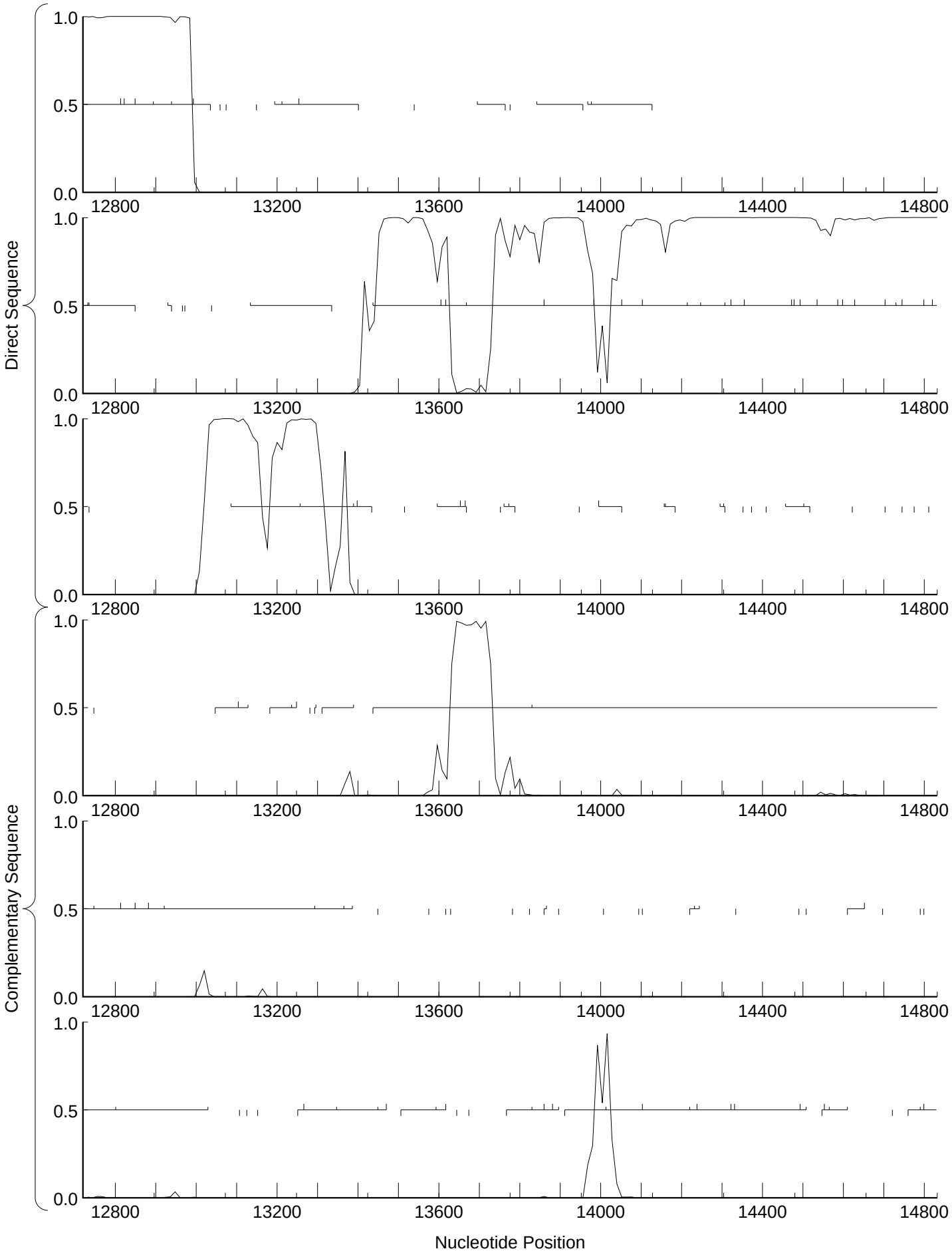
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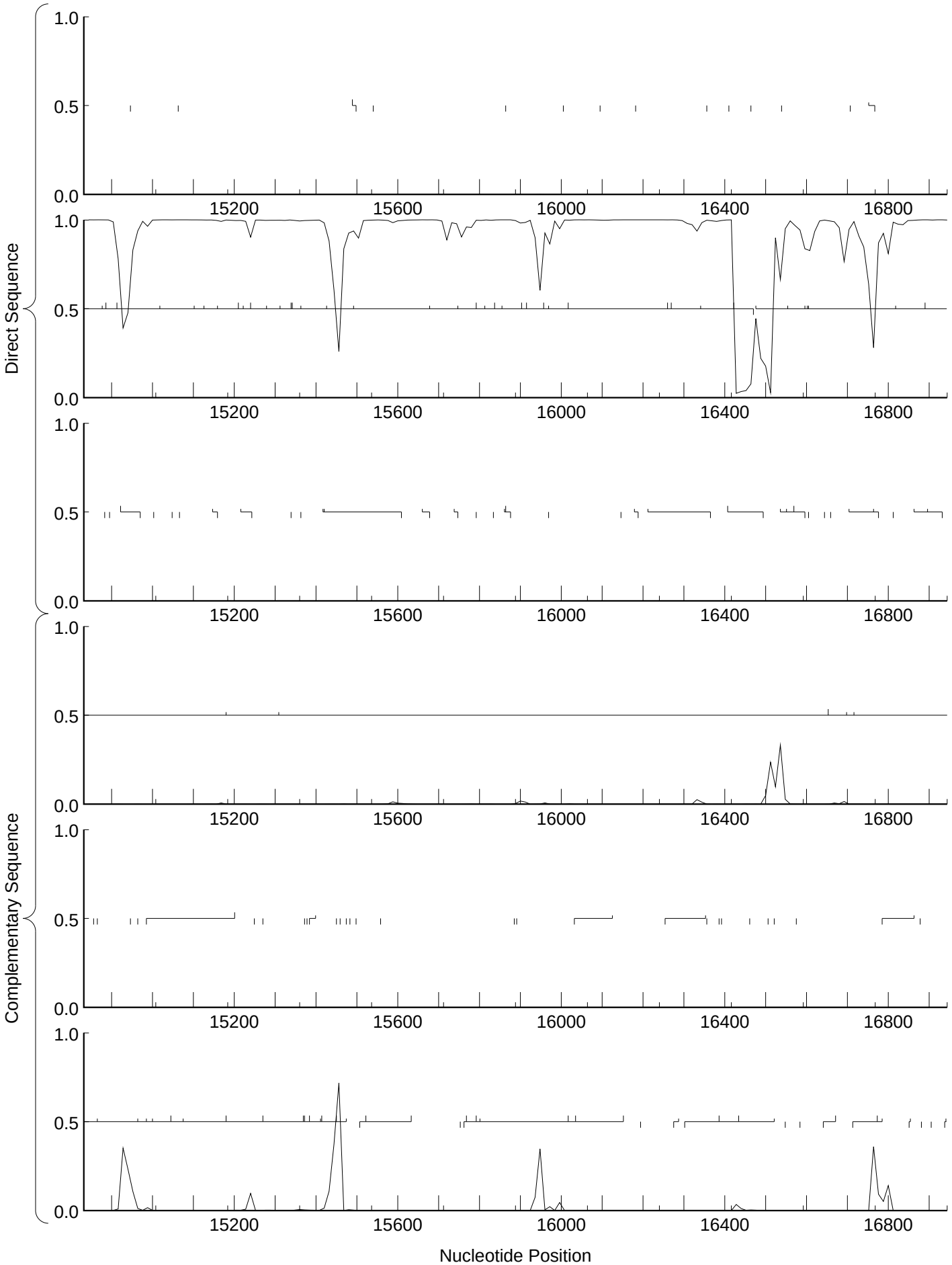
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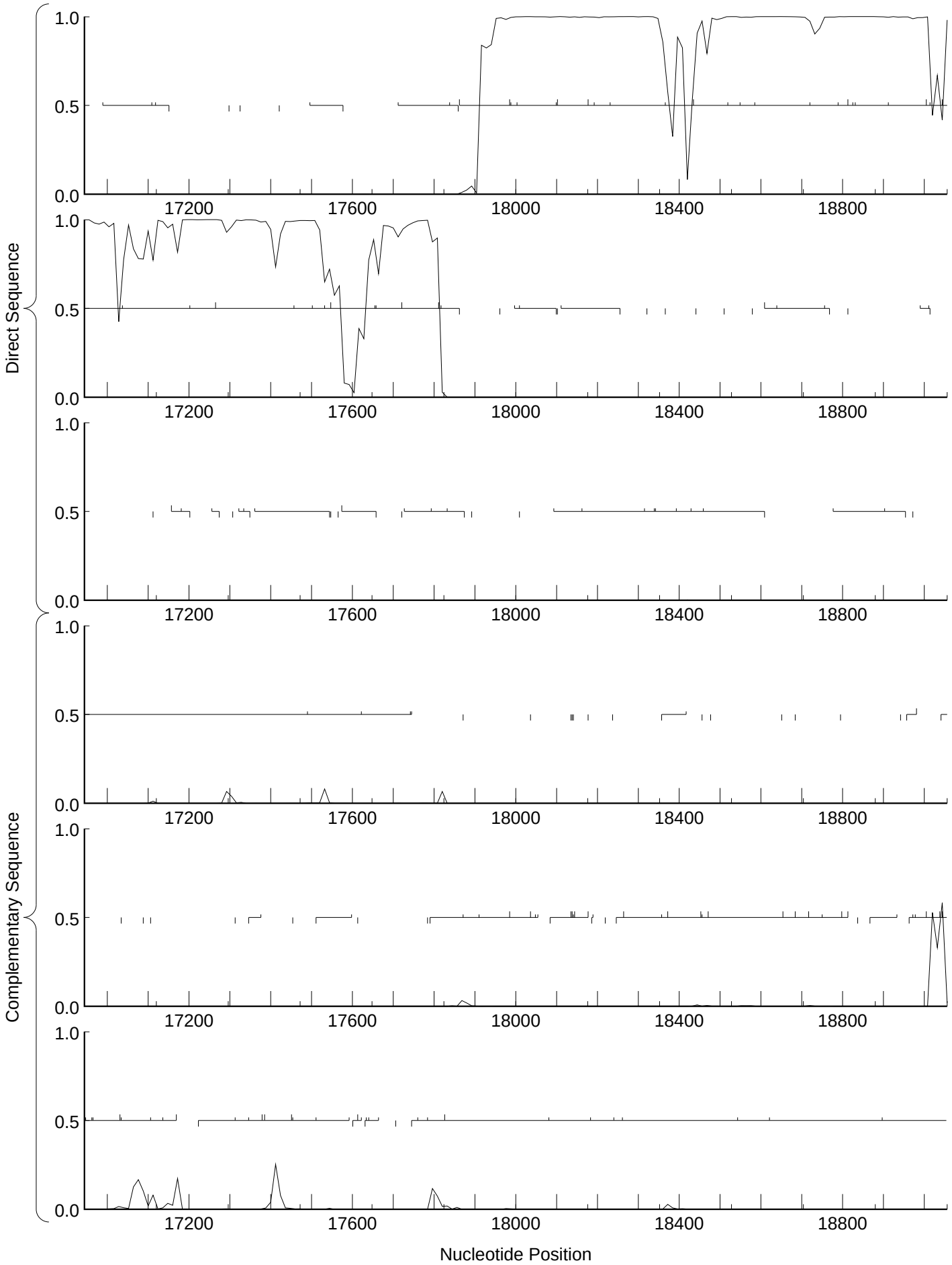
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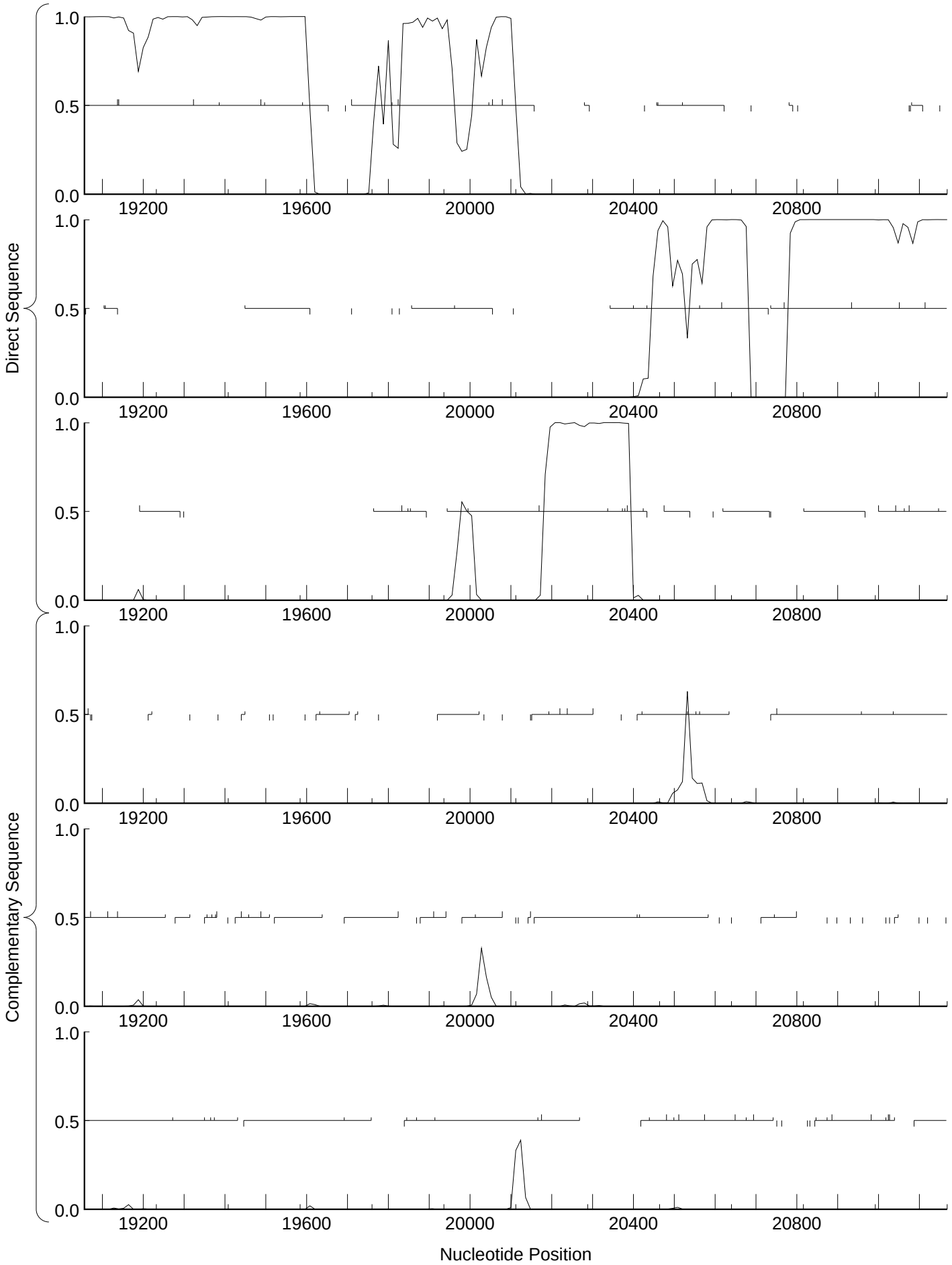
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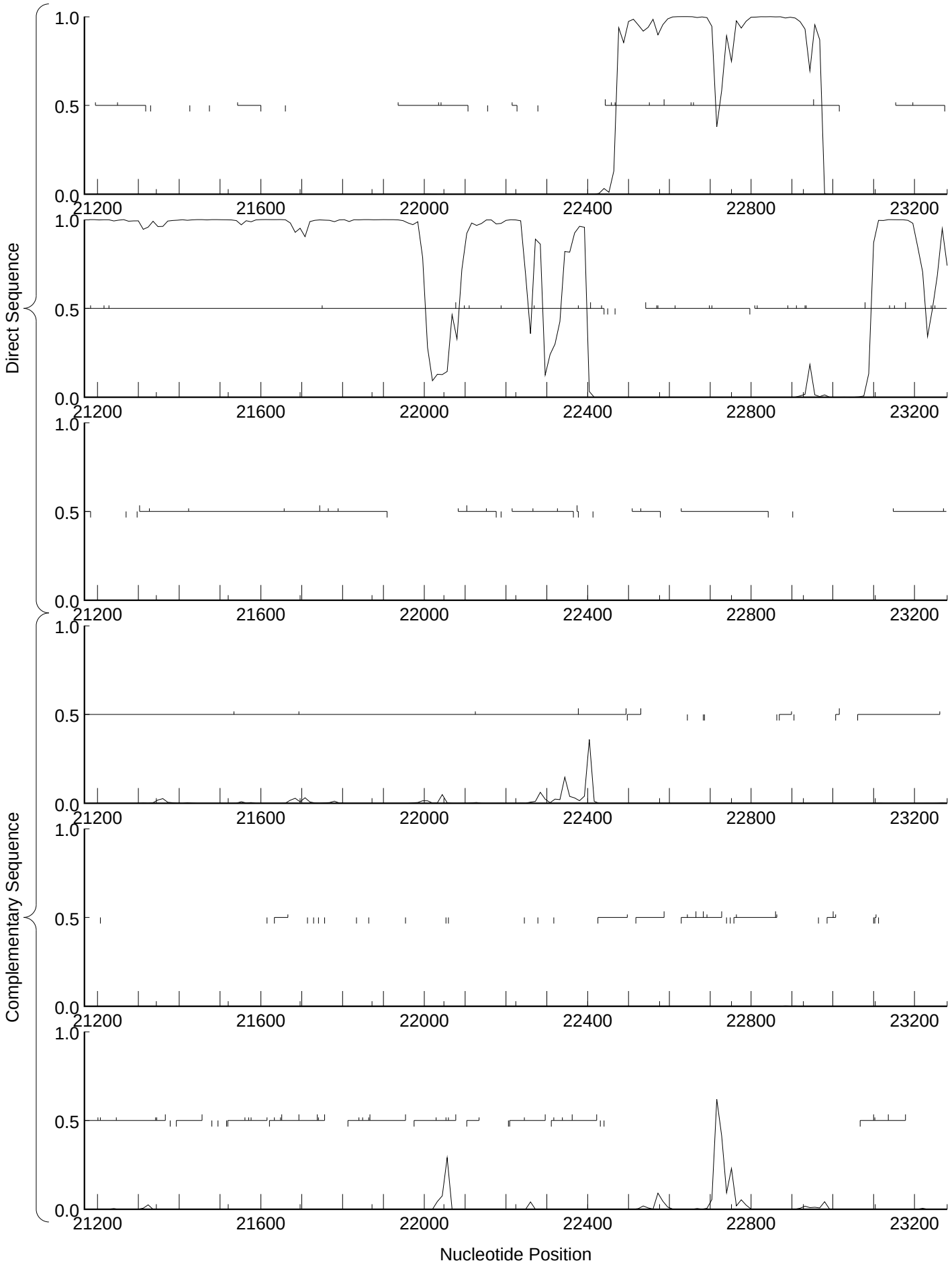
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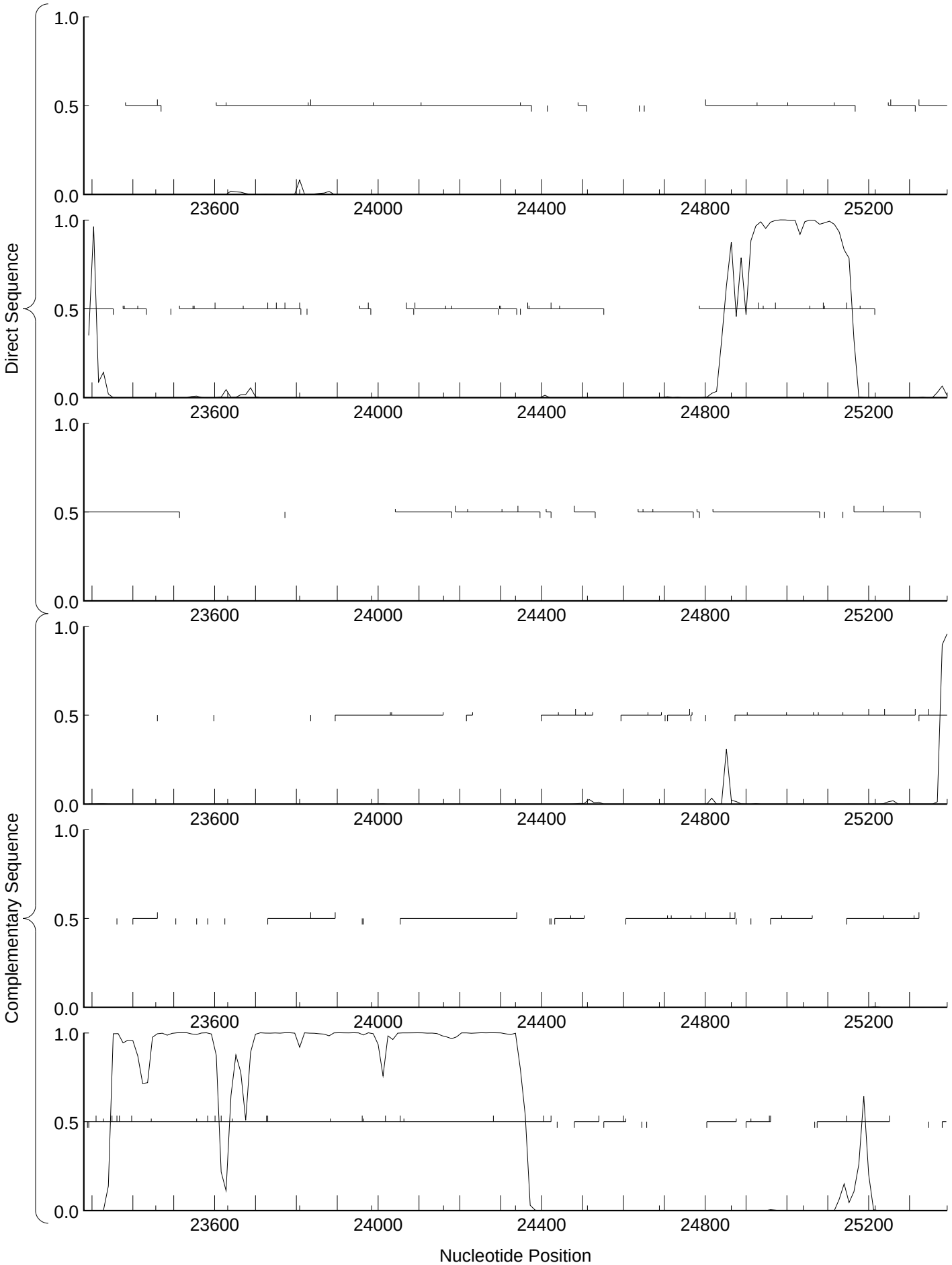
complete sequence, 50283 bp including 10-base 3’ overhang (CGGGCGGTAA), Cluster A19, Order 4, Window 96, Step 12, 11/25

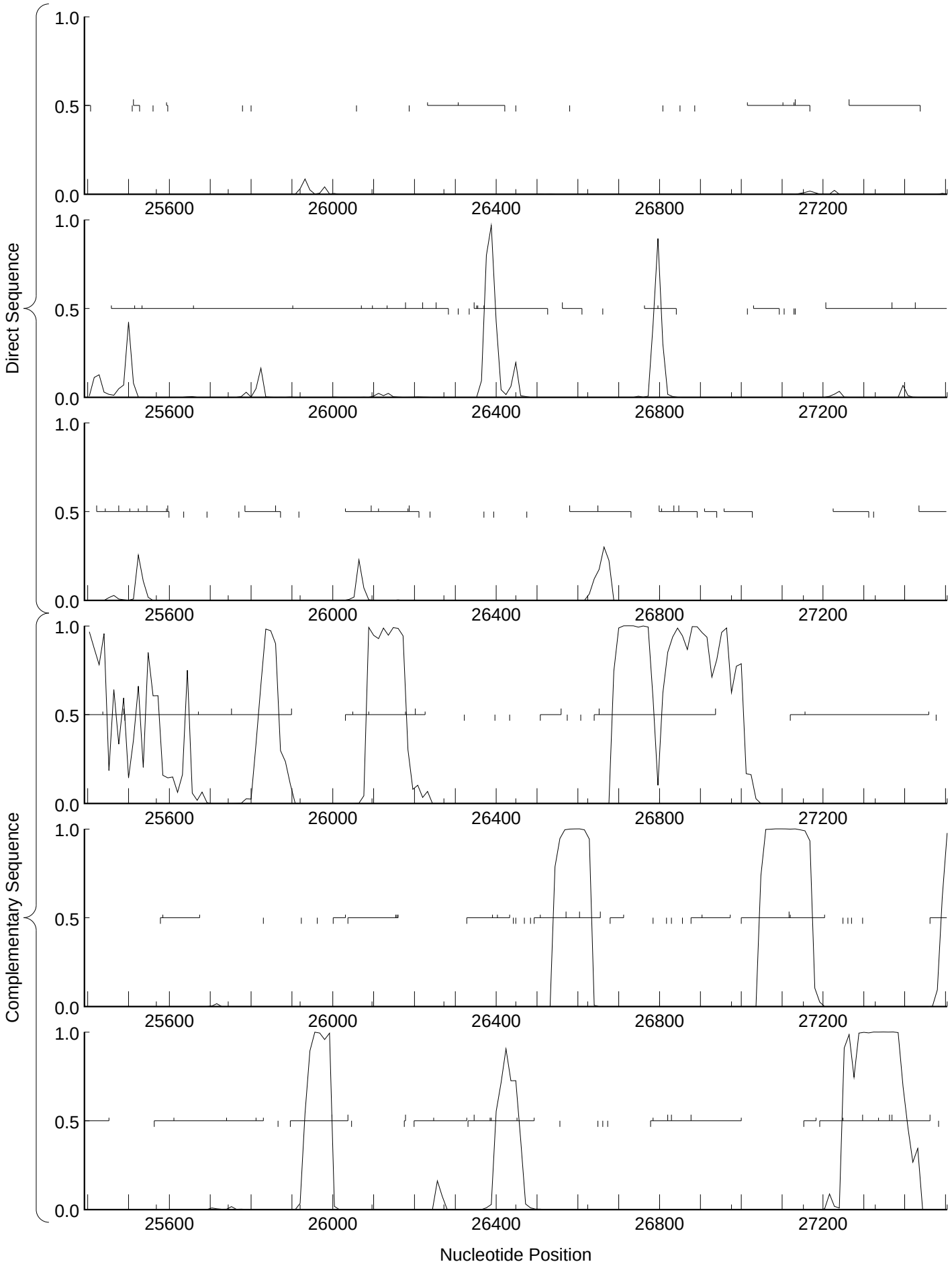


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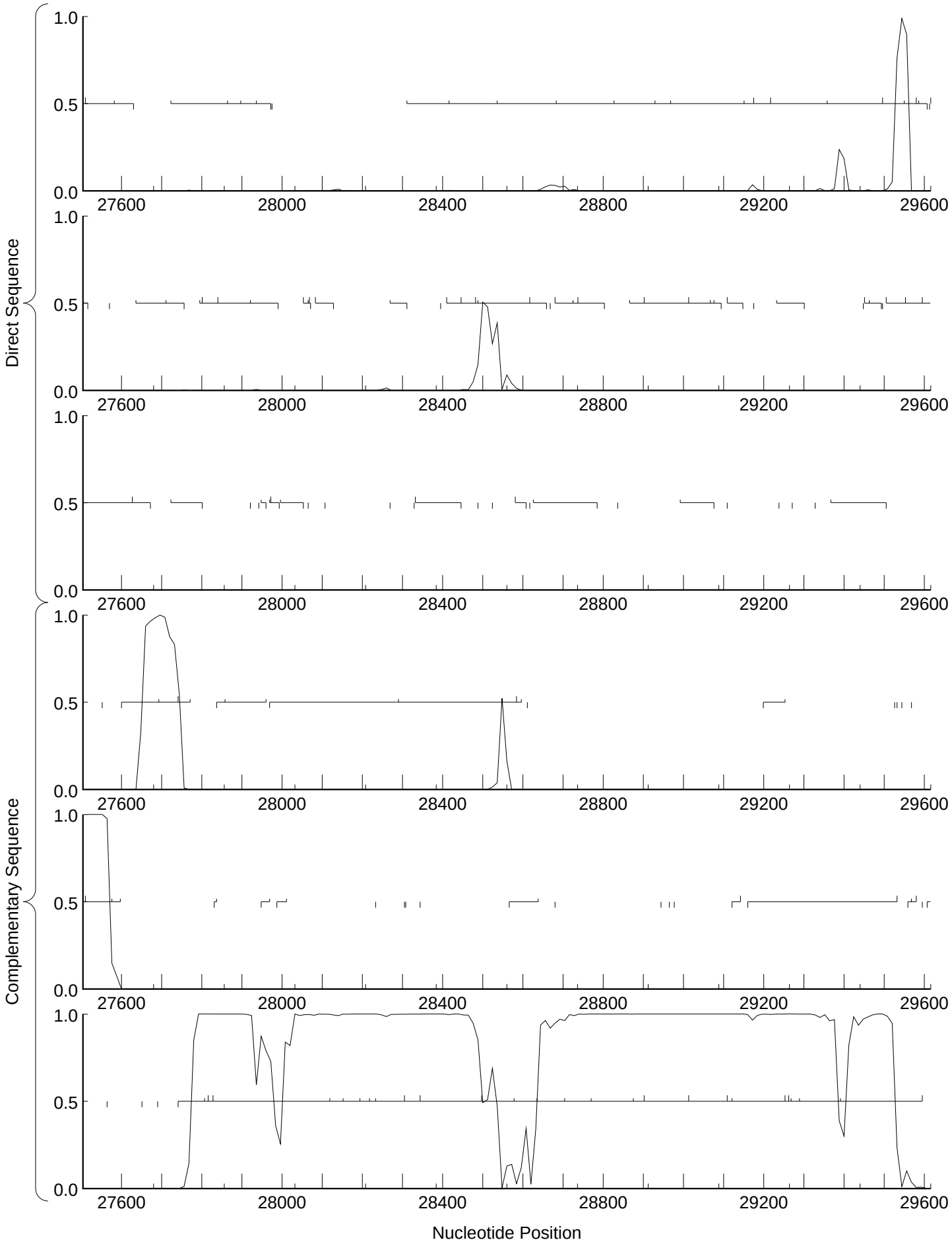


complete sequence, 50283 bp including 10-base 3’ overhang (CGGGCGGTAA), Cluster A19, Order 4, Window 96, Step 12, 13/25

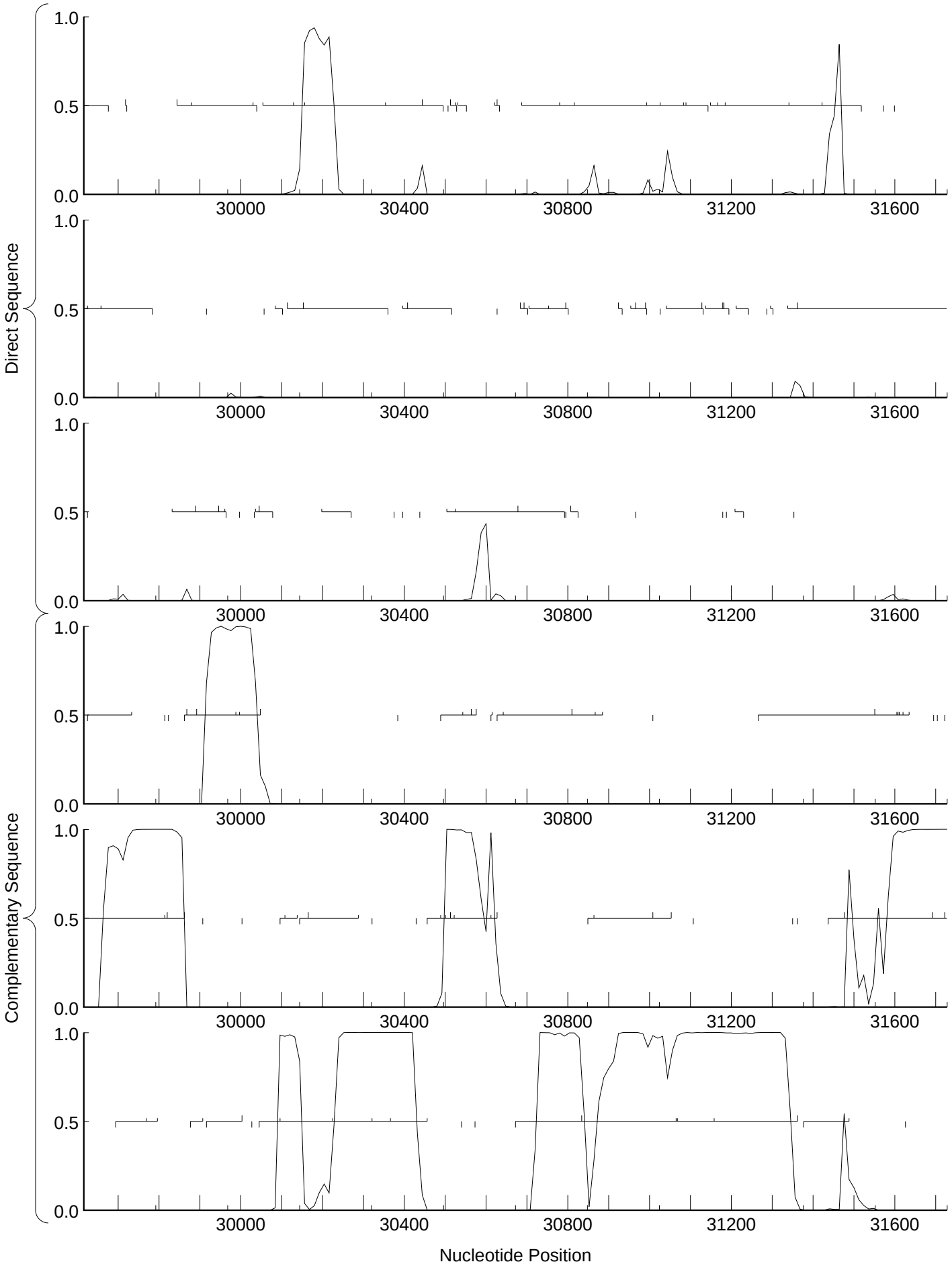


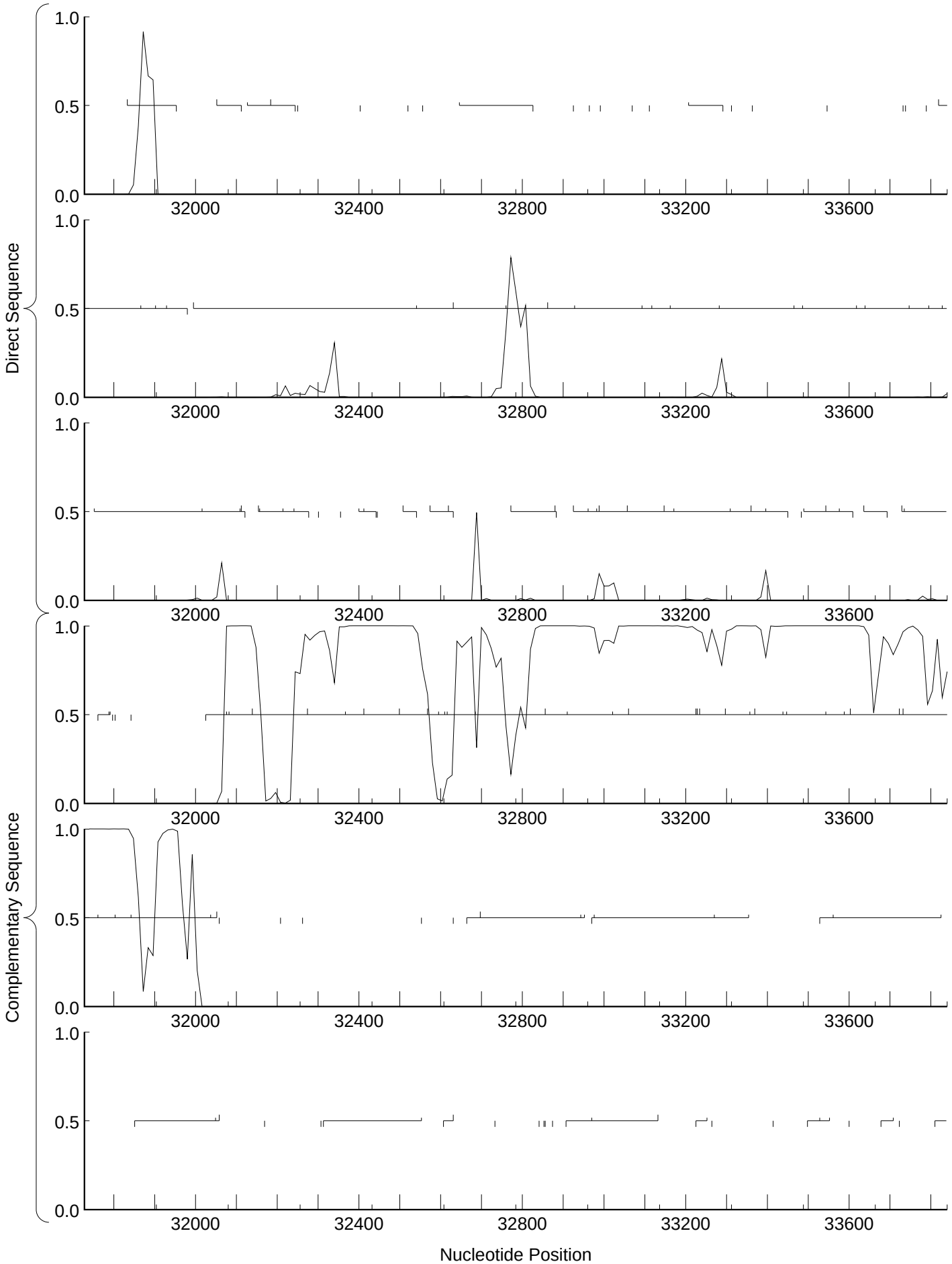

Direct Sequence
Complementary Sequence
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0.0
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26000
26400
26800
27200
Nucleotide Position

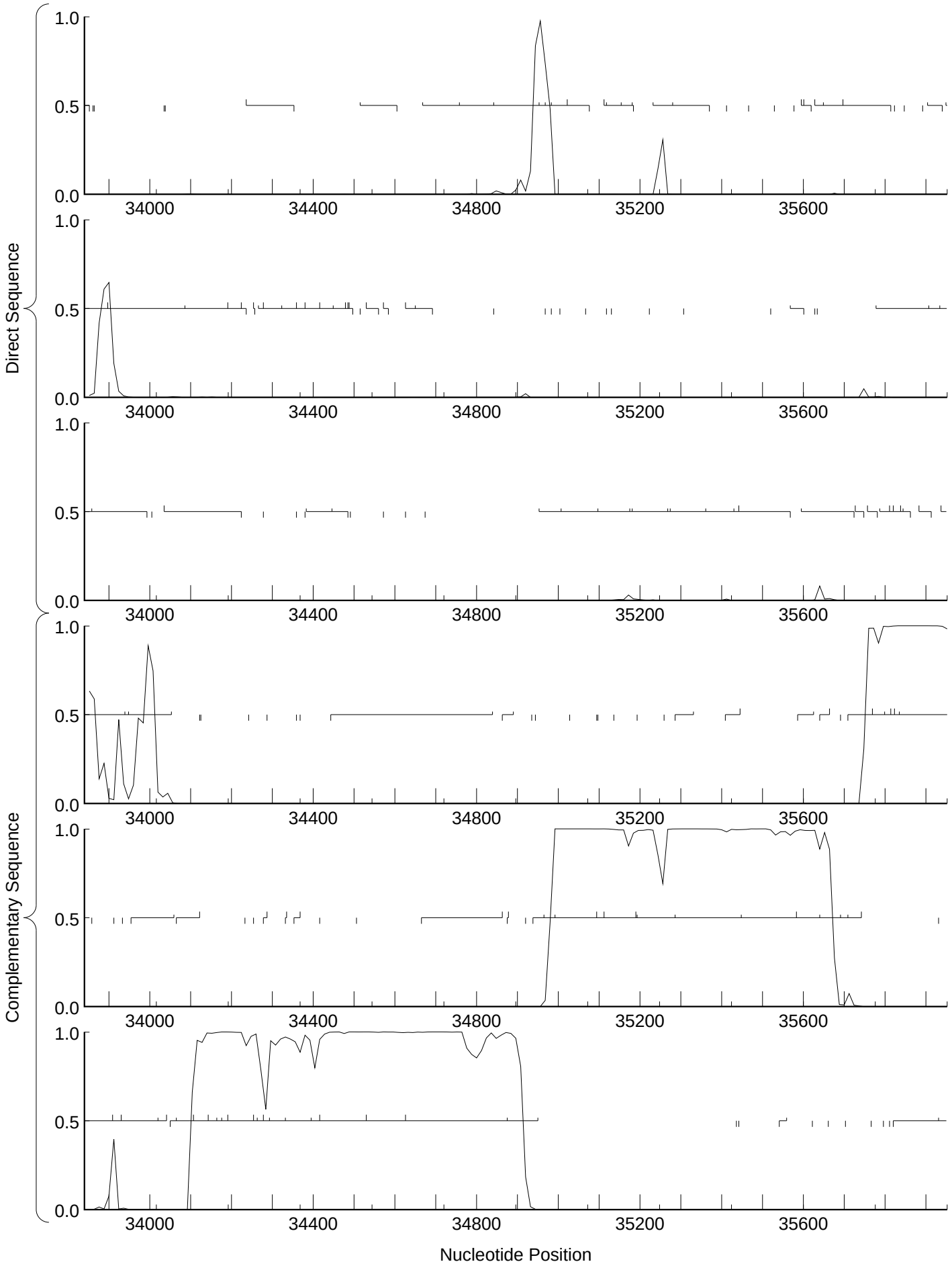
complete sequence, 50283 bp including 10-base 3' overhang (CGGGCGGTAA), Cluster A19, Order 4, Window 96, Step 12, 15/25

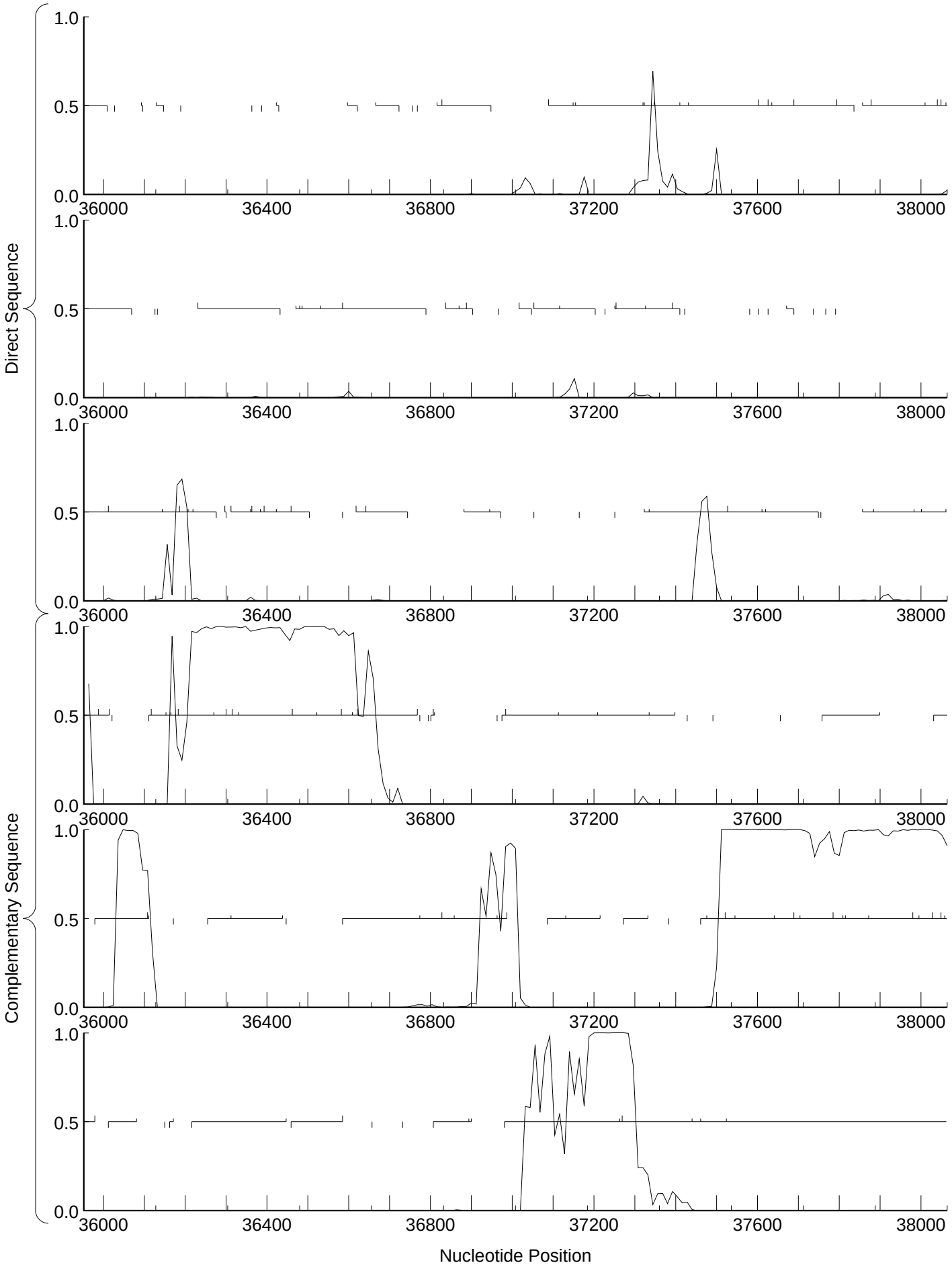


complete sequence, 50283 bp including 10-base 3’ overhang (CGGGCGGTAA), Cluster A19, Order 4, Window 96, Step 12, 16/25

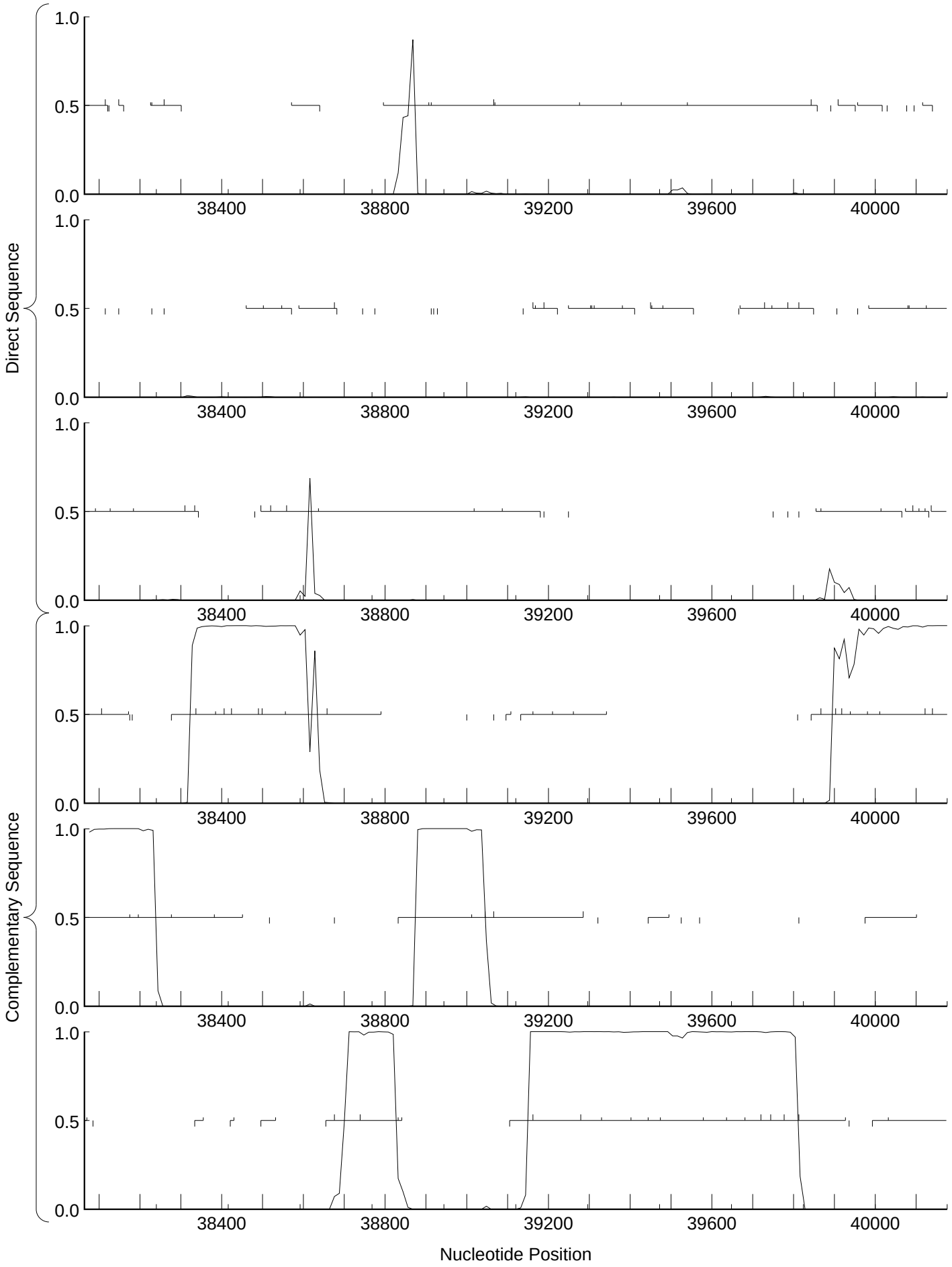



Direct Sequence
Complementary Sequence
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0.5
0.0
32000
32400
32800
33200
33600
Nucleotide Position

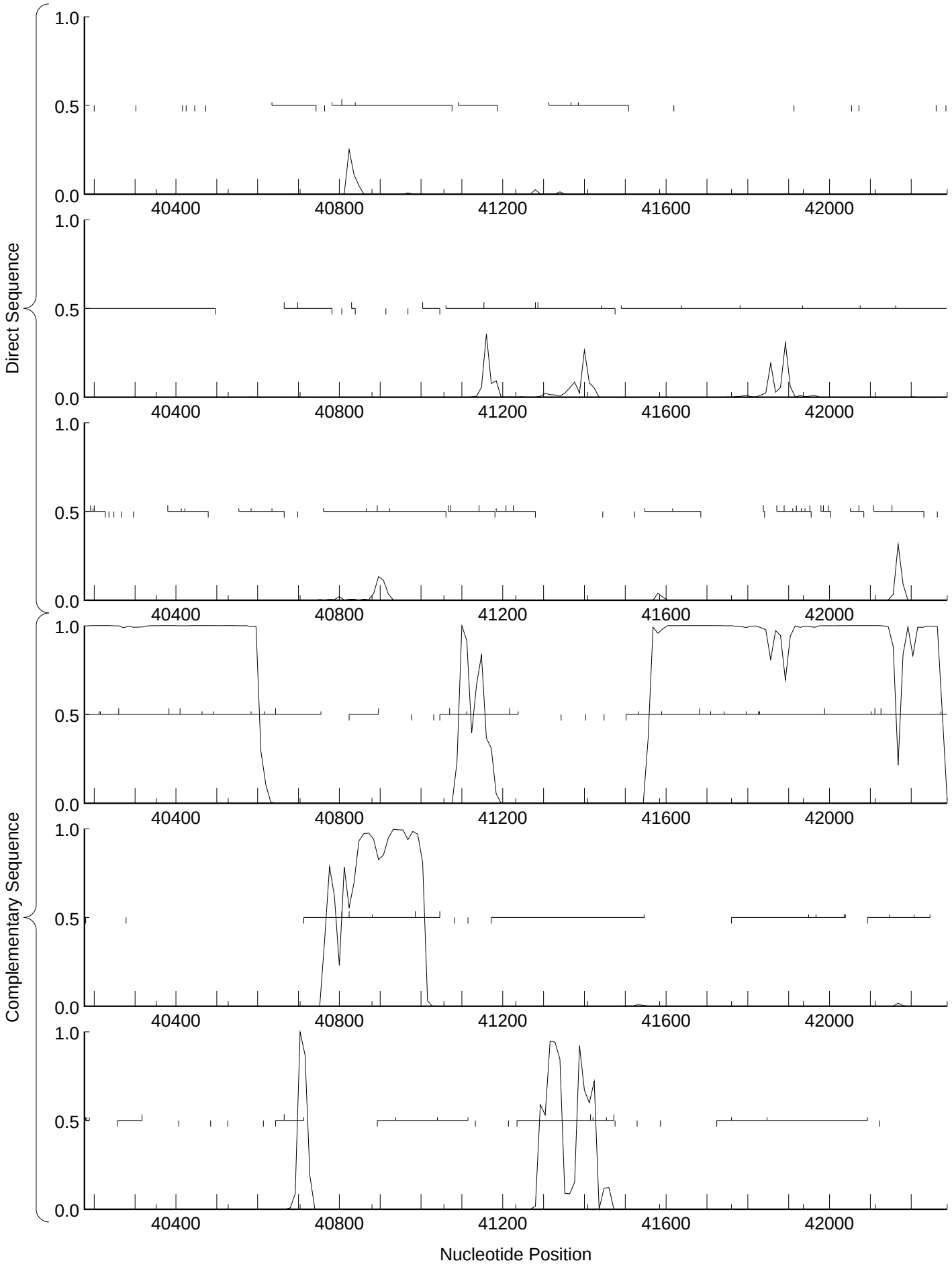

Direct Sequence
Complementary Sequence
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0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

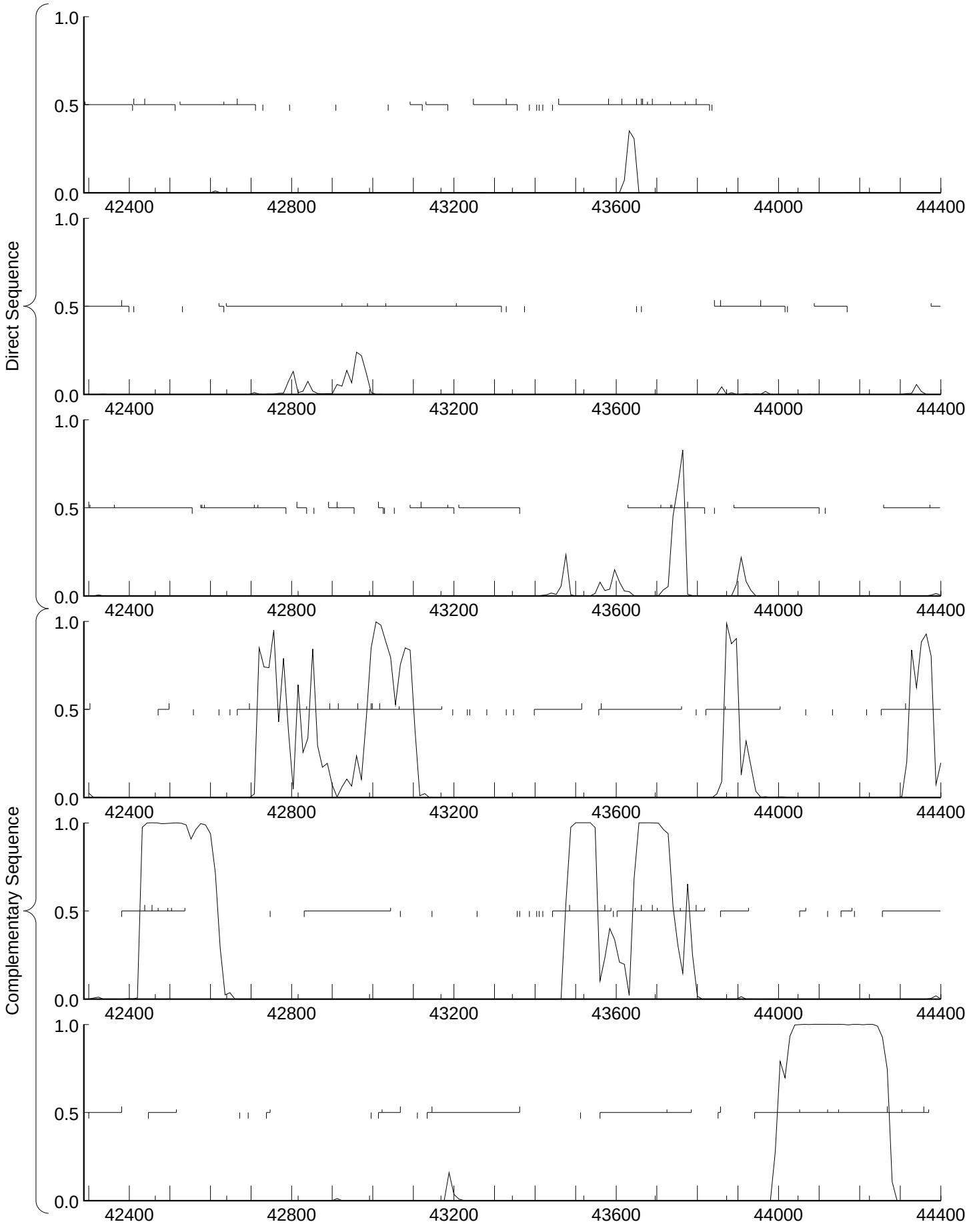

Direct Sequence
Complementary Sequence
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0.5
0.0
36000
36400
36800
37200
37600
38000
Nucleotide Position

complete sequence, 50283 bp including 10-base 3' overhang (CGGGCGGTAA), Cluster A19, Order 4, Window 96, Step 12, 20/25

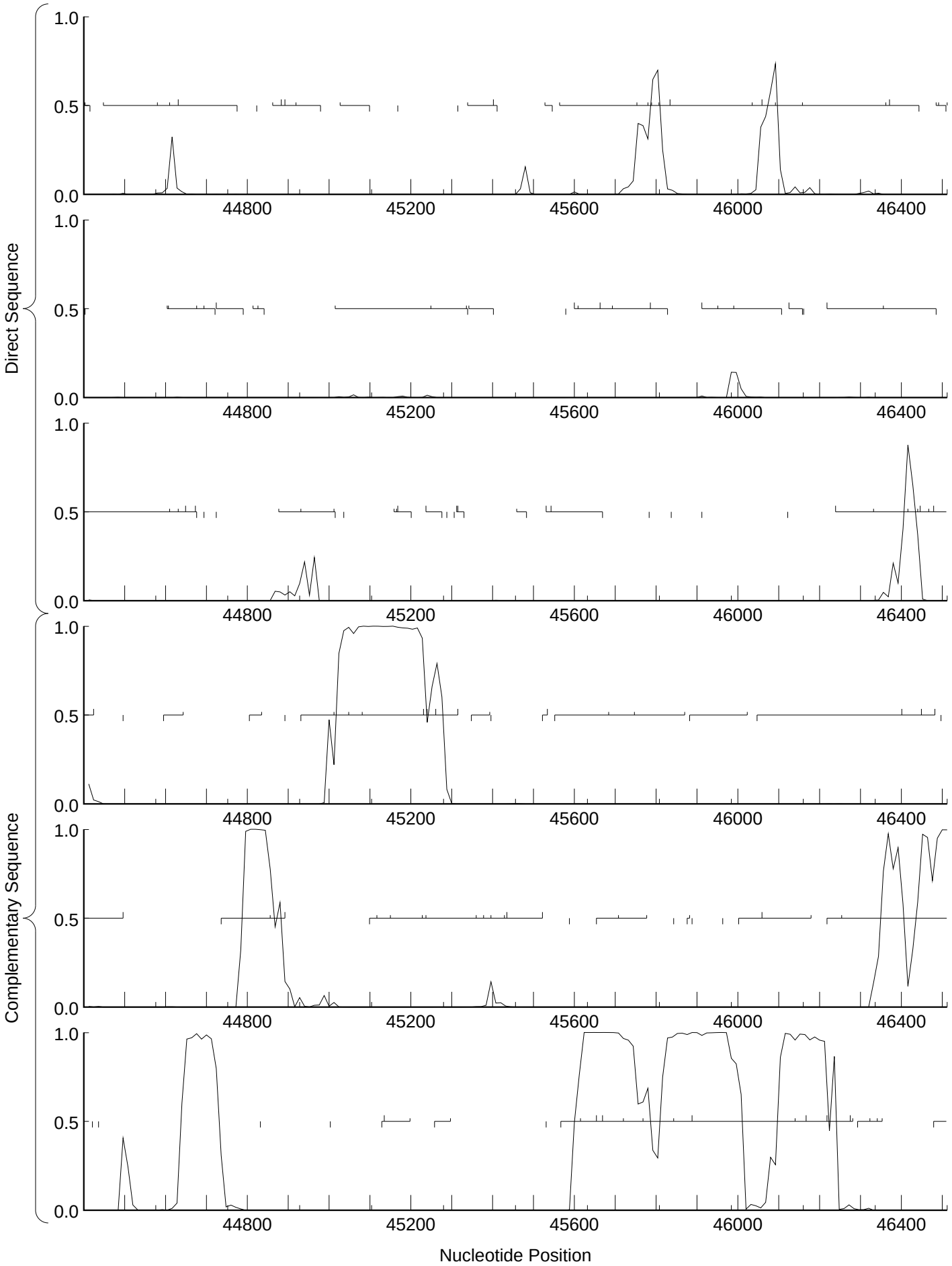


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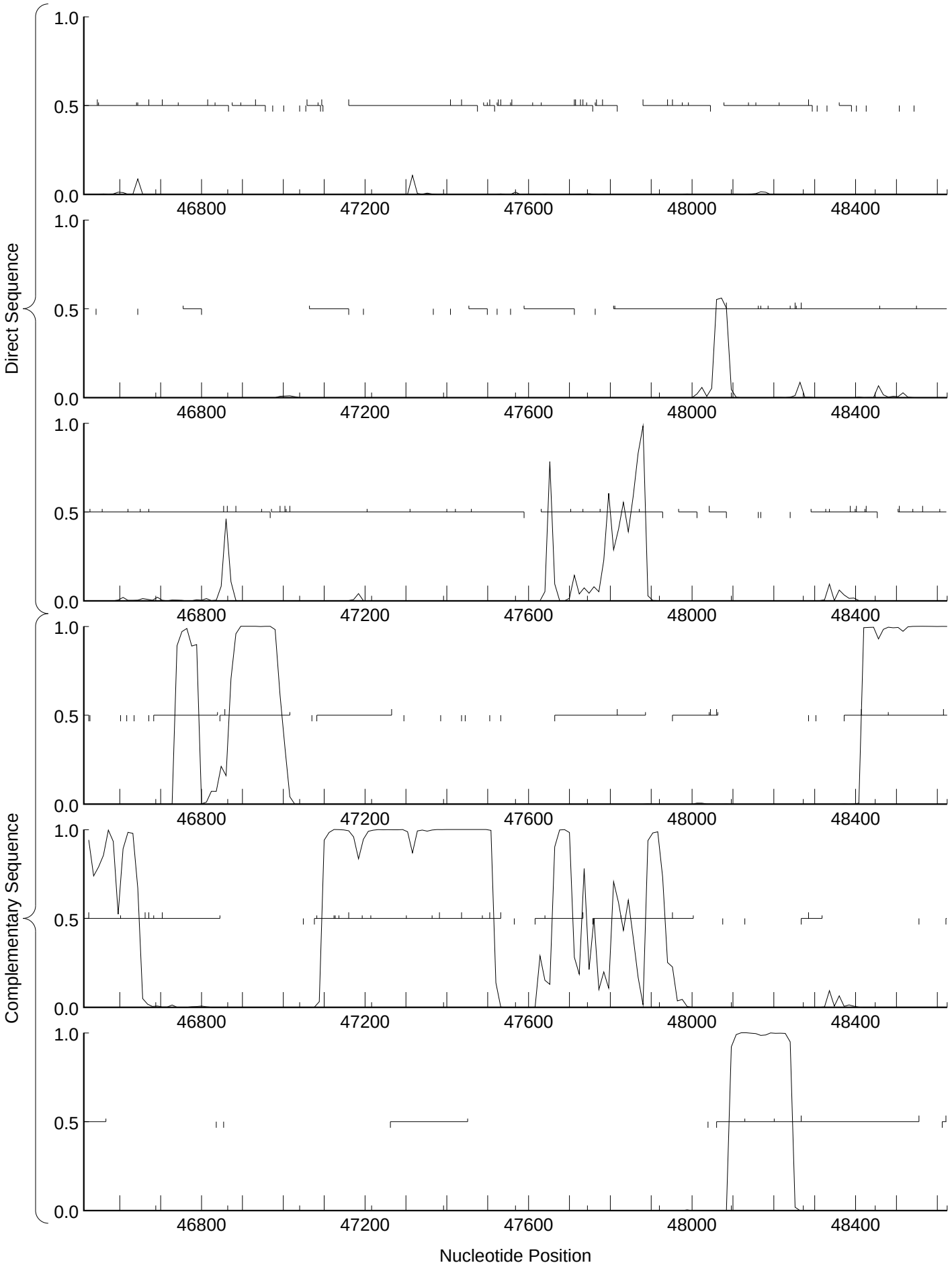



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
42400
42800
43200
43600
44000
44400
Nucleotide Position

complete sequence, 50283 bp including 10-base 3' overhang (CGGGCGGTAA), Cluster A19, Order 4, Window 96, Step 12, 23/25



complete sequence, 50283 bp including 10-base 3' overhang (CGGGCGGTAA), Cluster A19, Order 4, Window 96, Step 12, 24/25



complete sequence, 50283 bp including 10-base 3' overhang (CGGGCGGTAA), Cluster A19, Order 4, Window 96, Step 12, 25/25

