GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Streptomyces phage Maya complete sequence, 55713 bp including 9-base 3' overhang (CGCCCGCCT
Analysis Date    :  3/28/24 at 10:40:04
Pages            :  28
Sequence Length  :  55713 bp
GC Content       :   59.50%

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

PS-Version       :  1.2

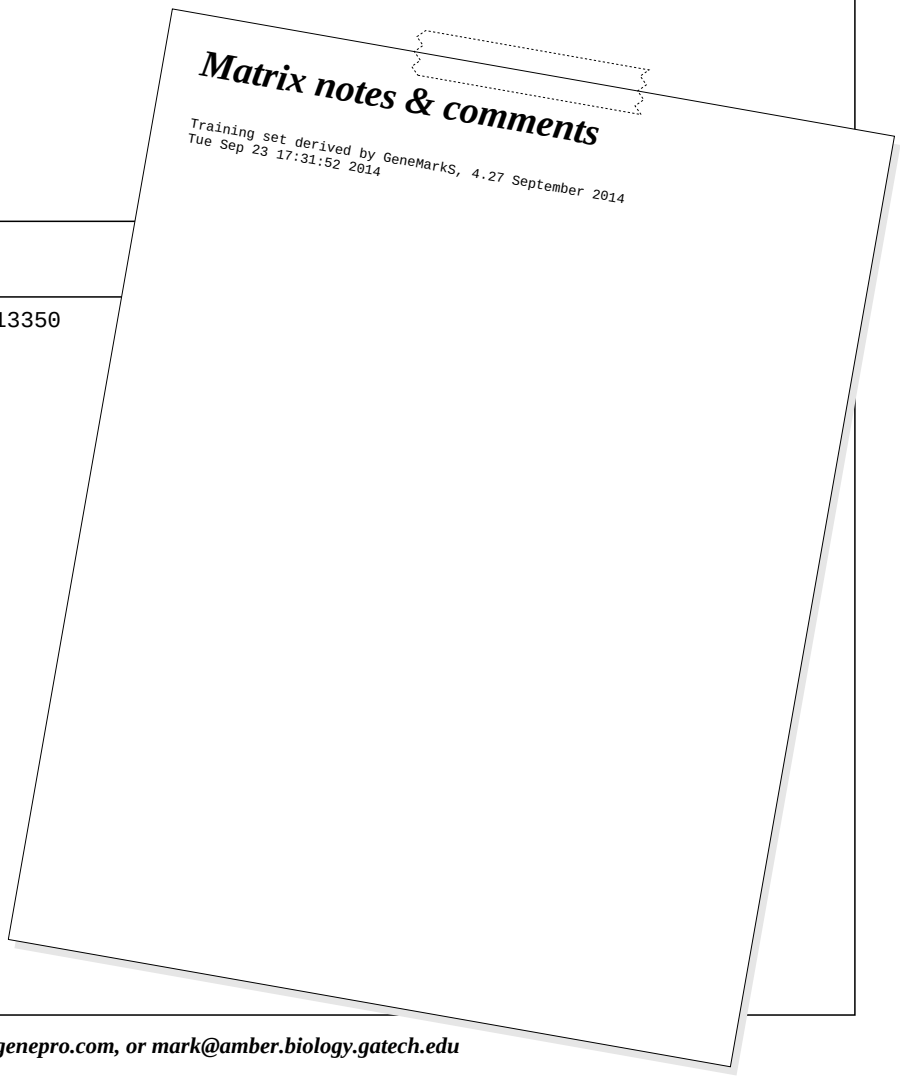
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Author     :  -

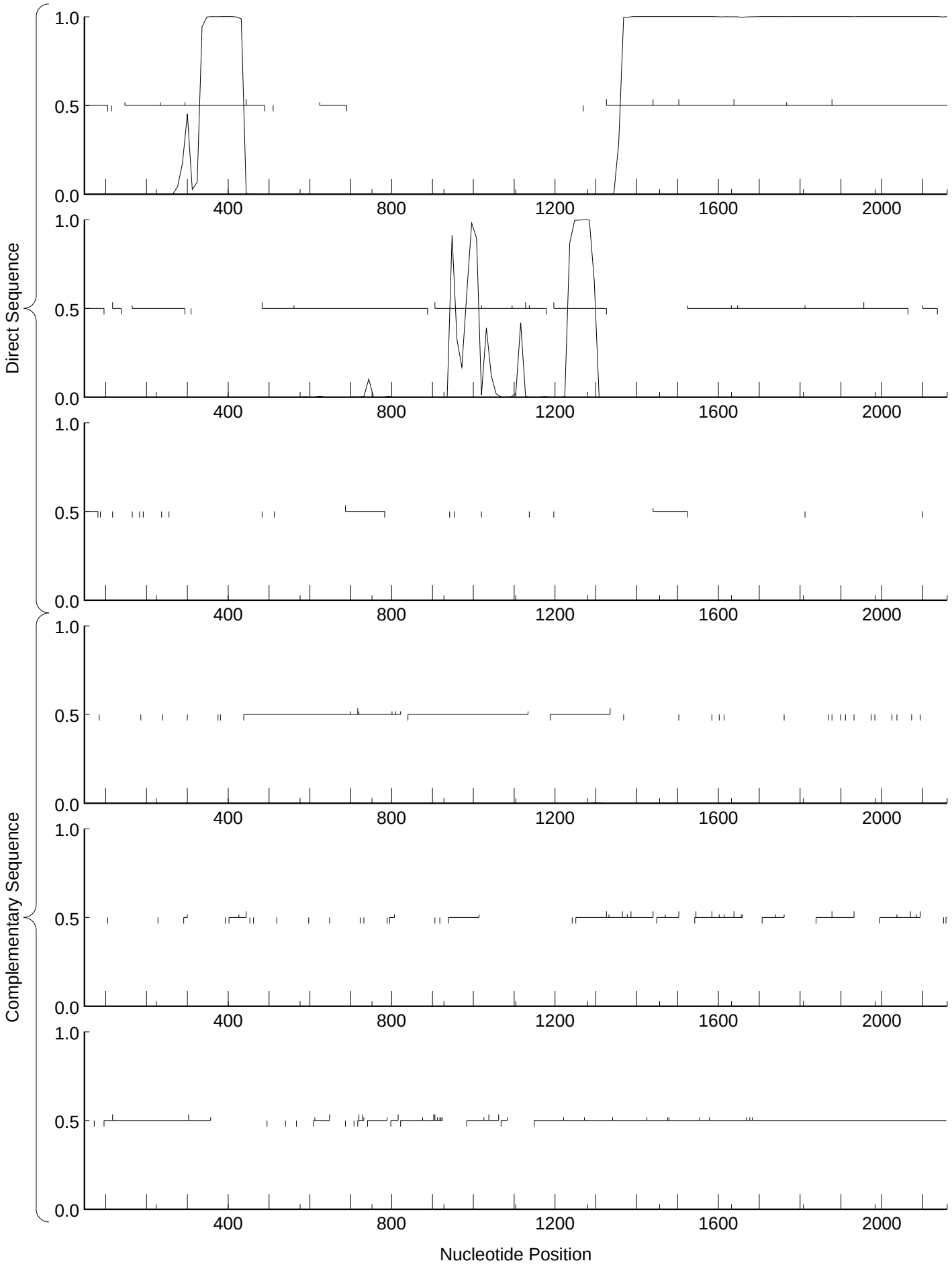
Order      :  4


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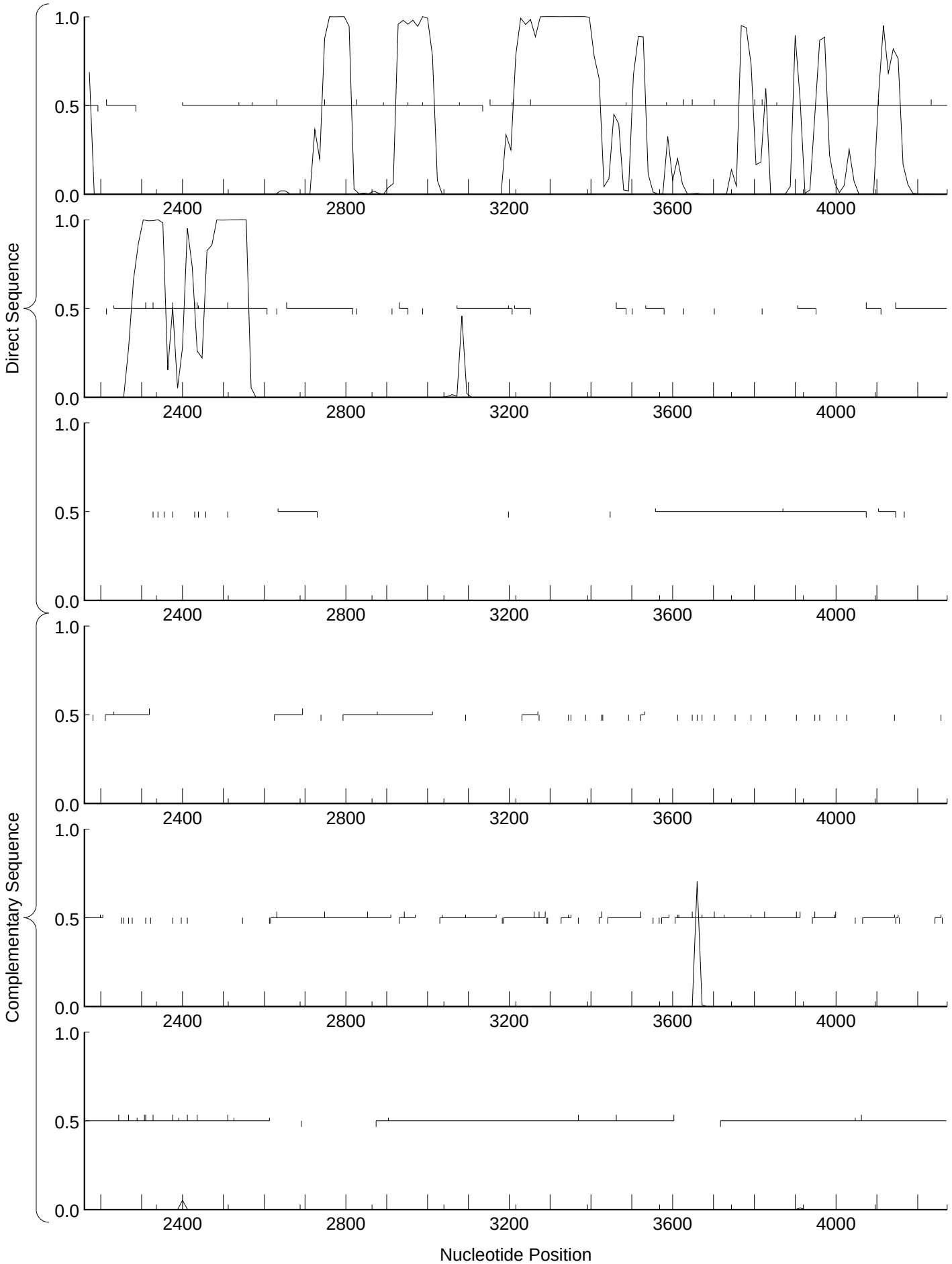


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

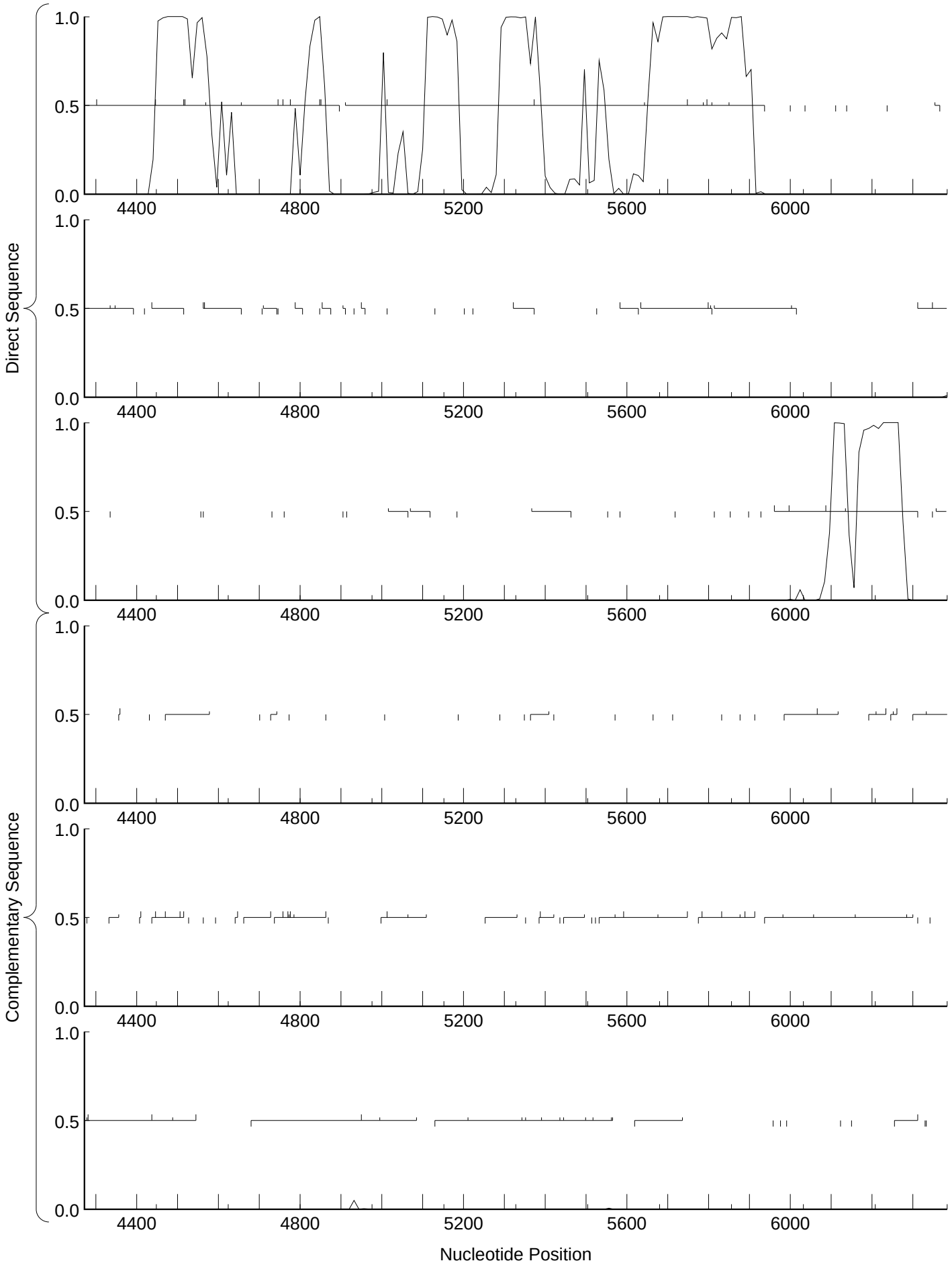
aya complete sequence, 55713 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 2/28



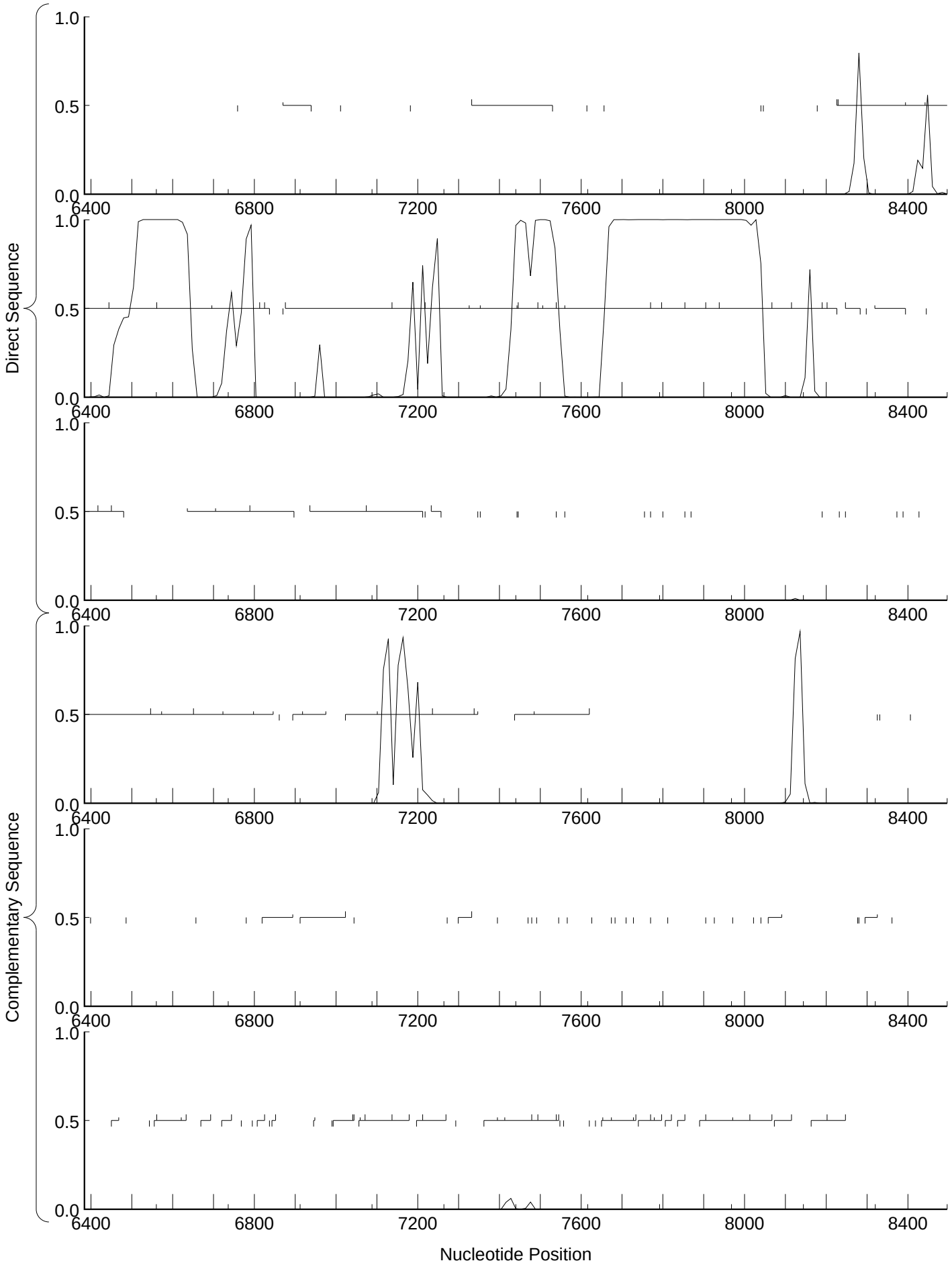
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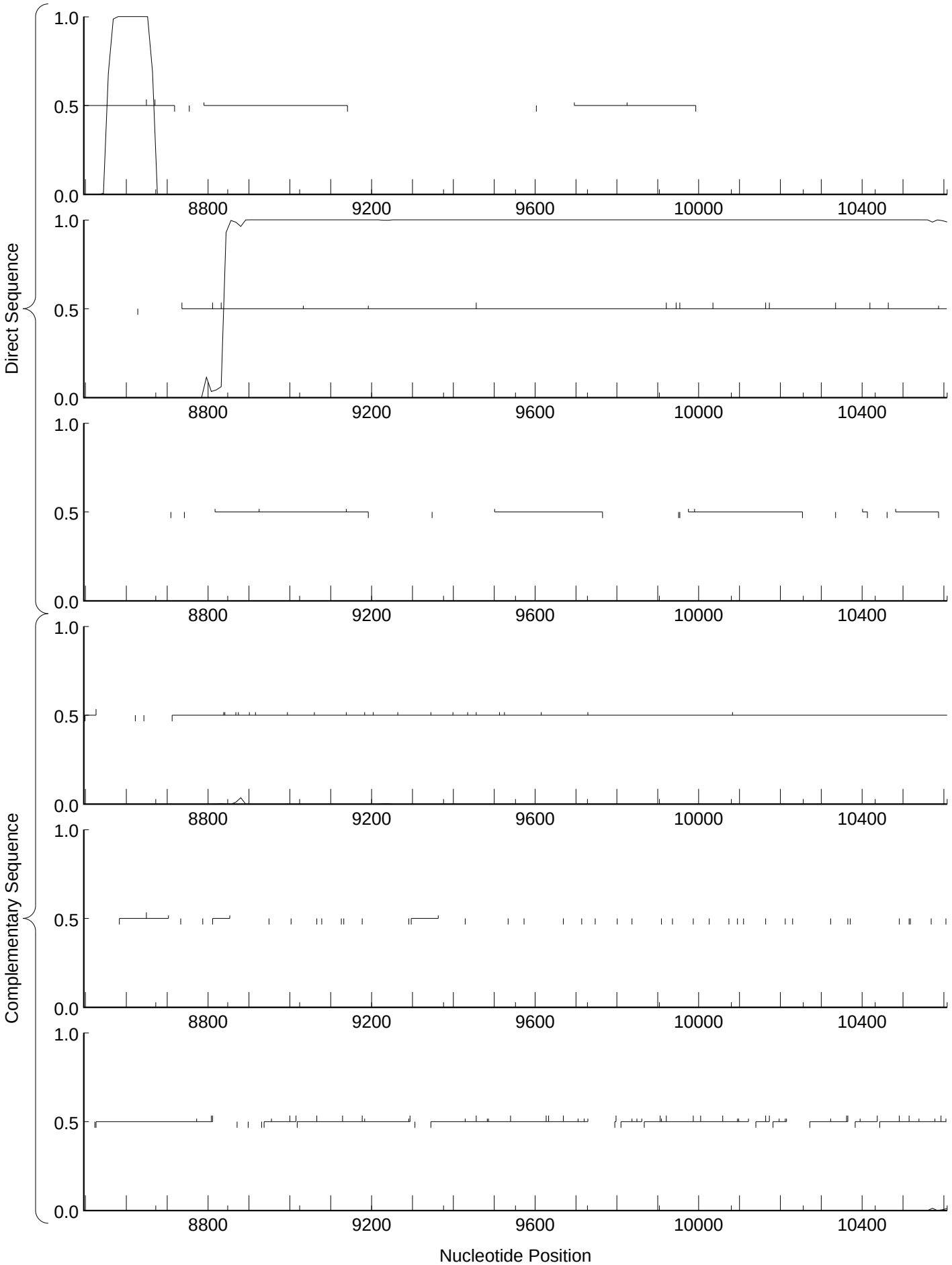
aya complete sequence, 55713 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 4/28



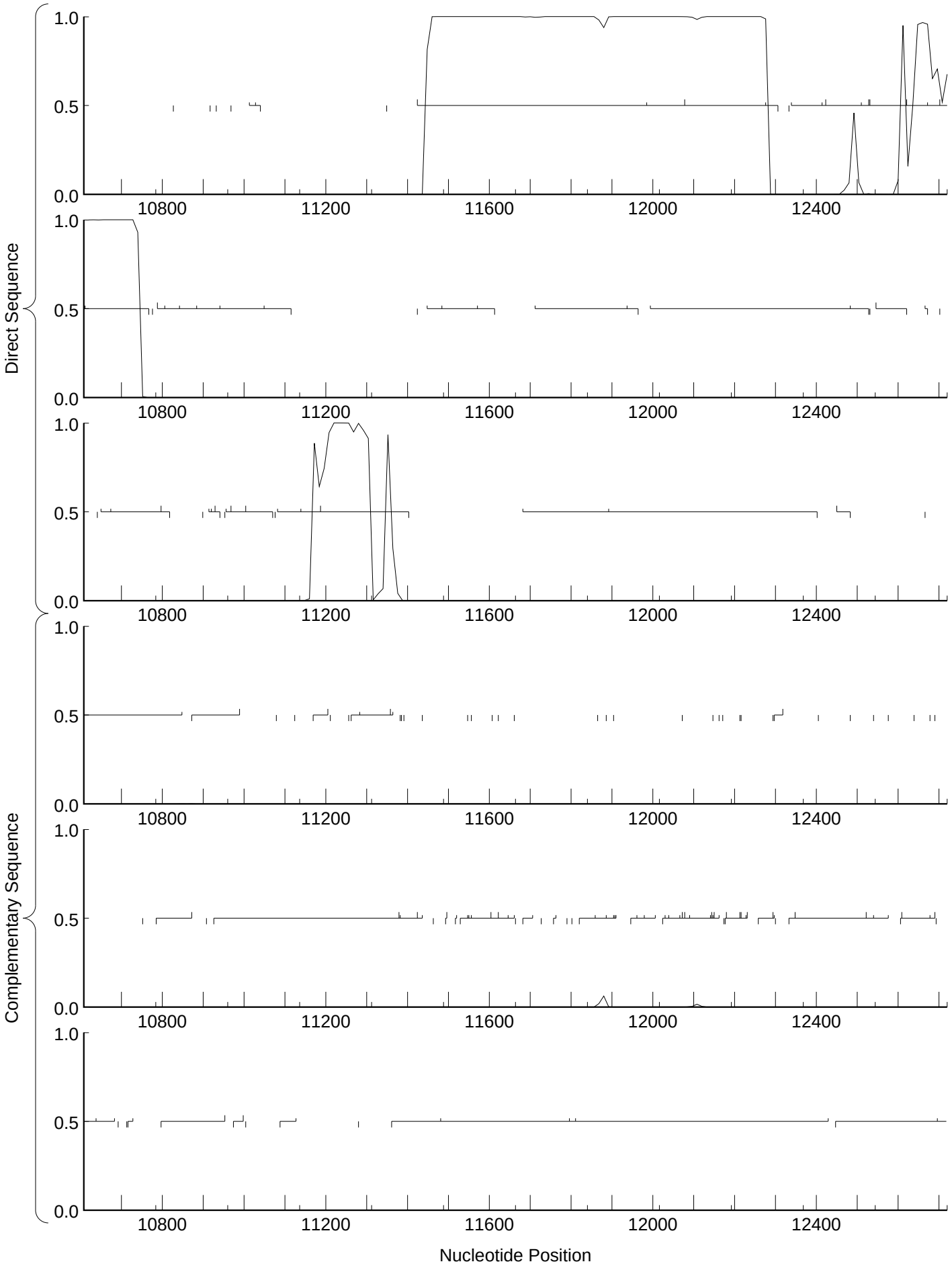
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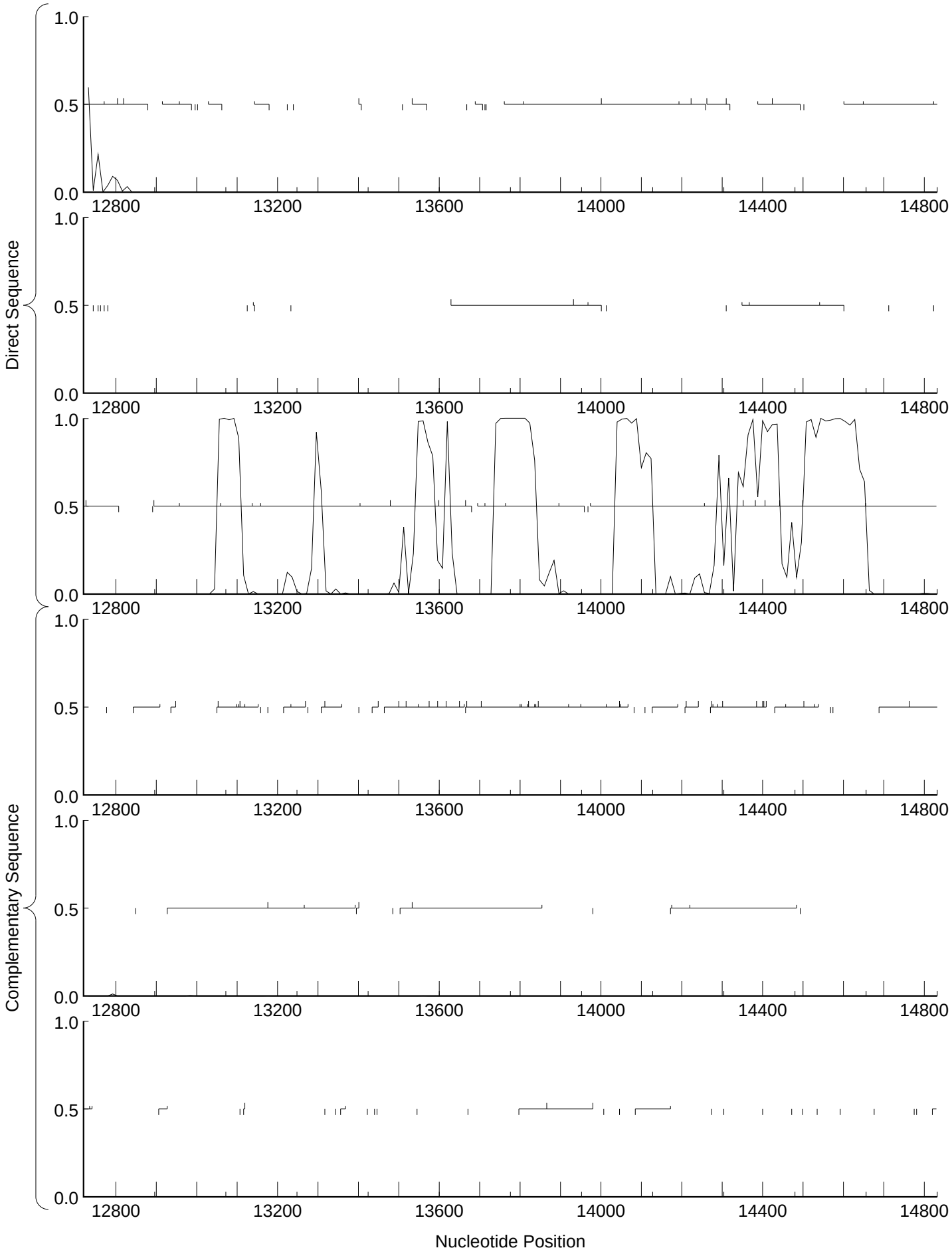
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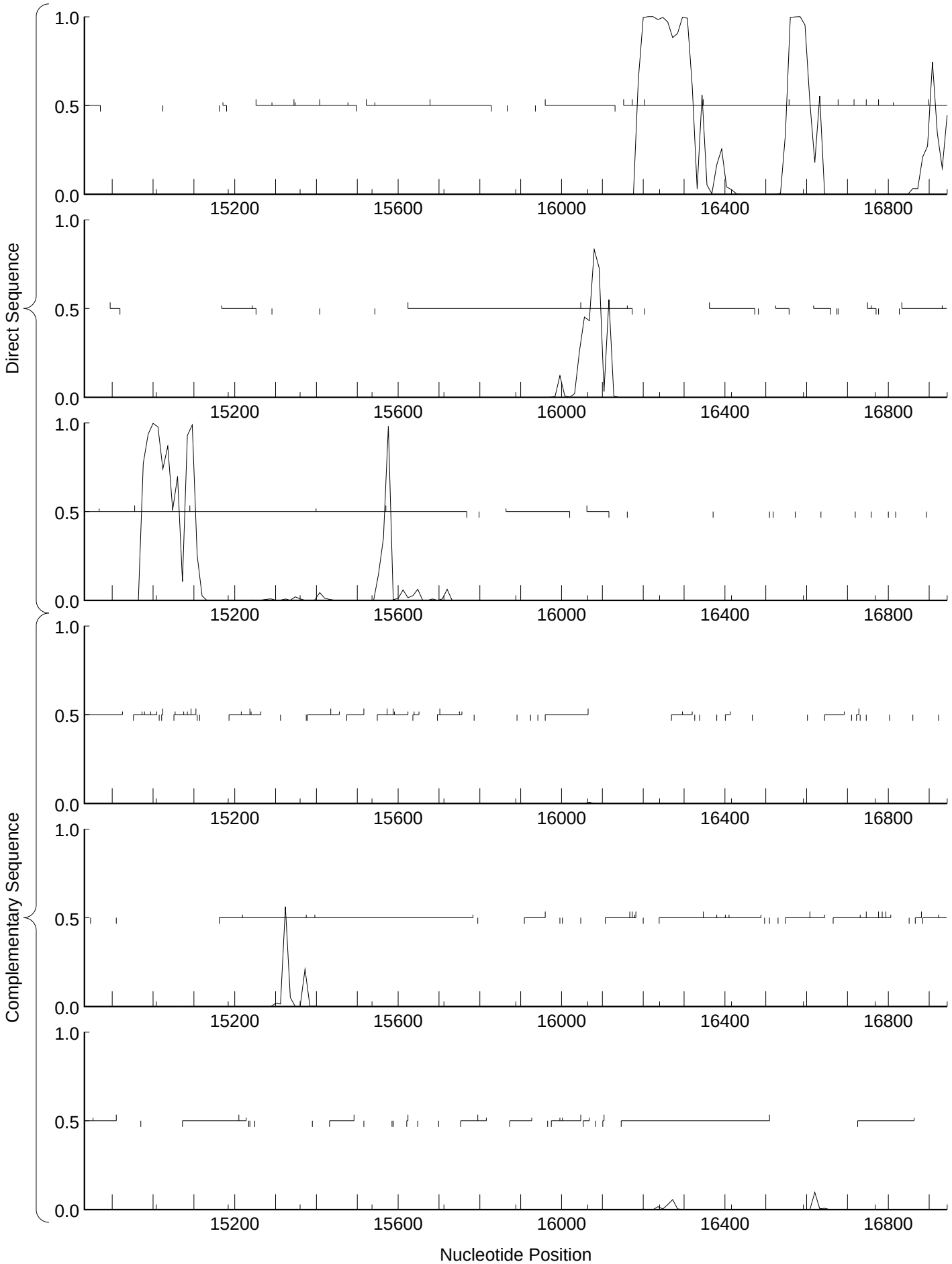
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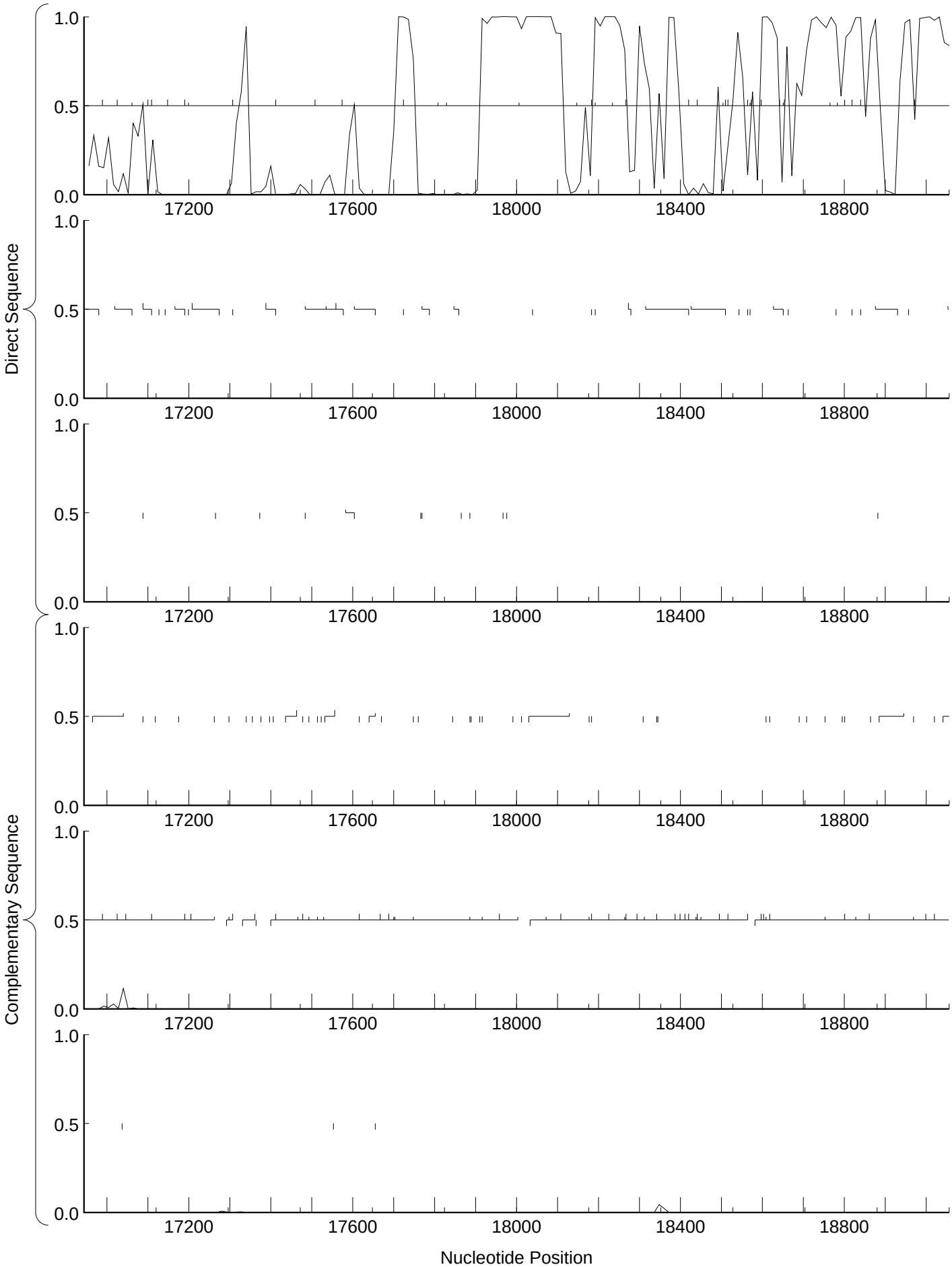
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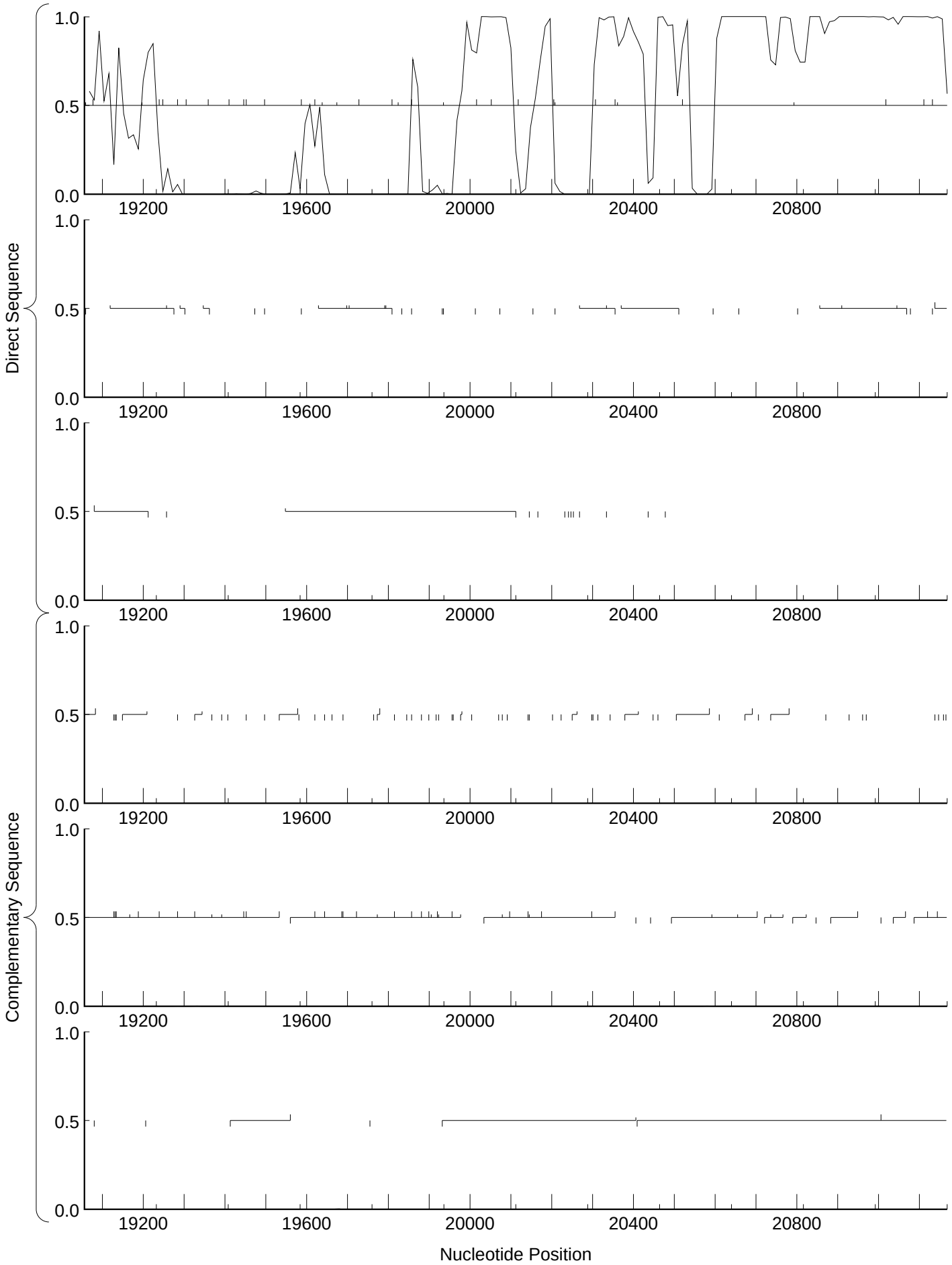
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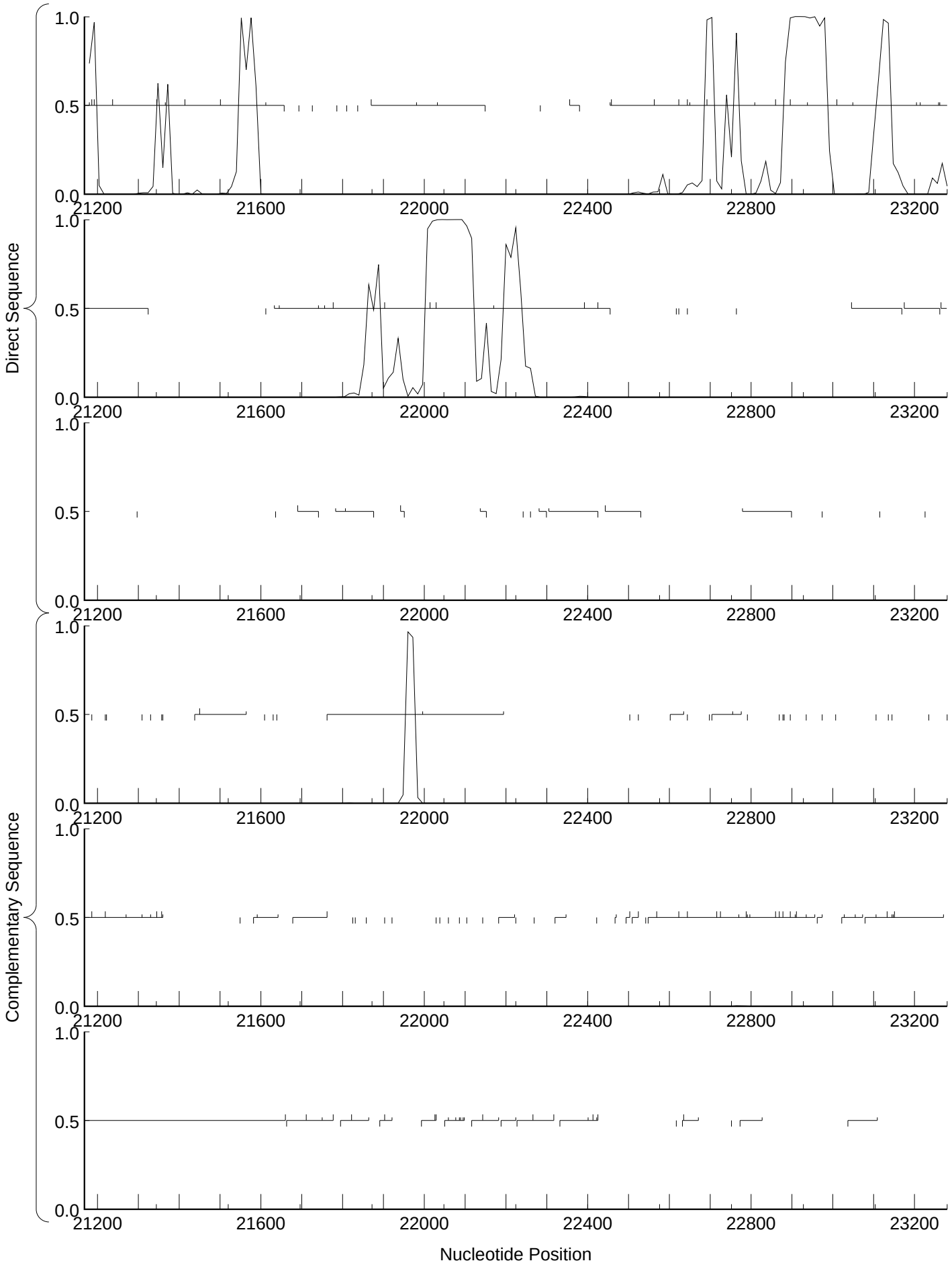
ya complete sequence, 55713 bp including 9-base 3' overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 10/28



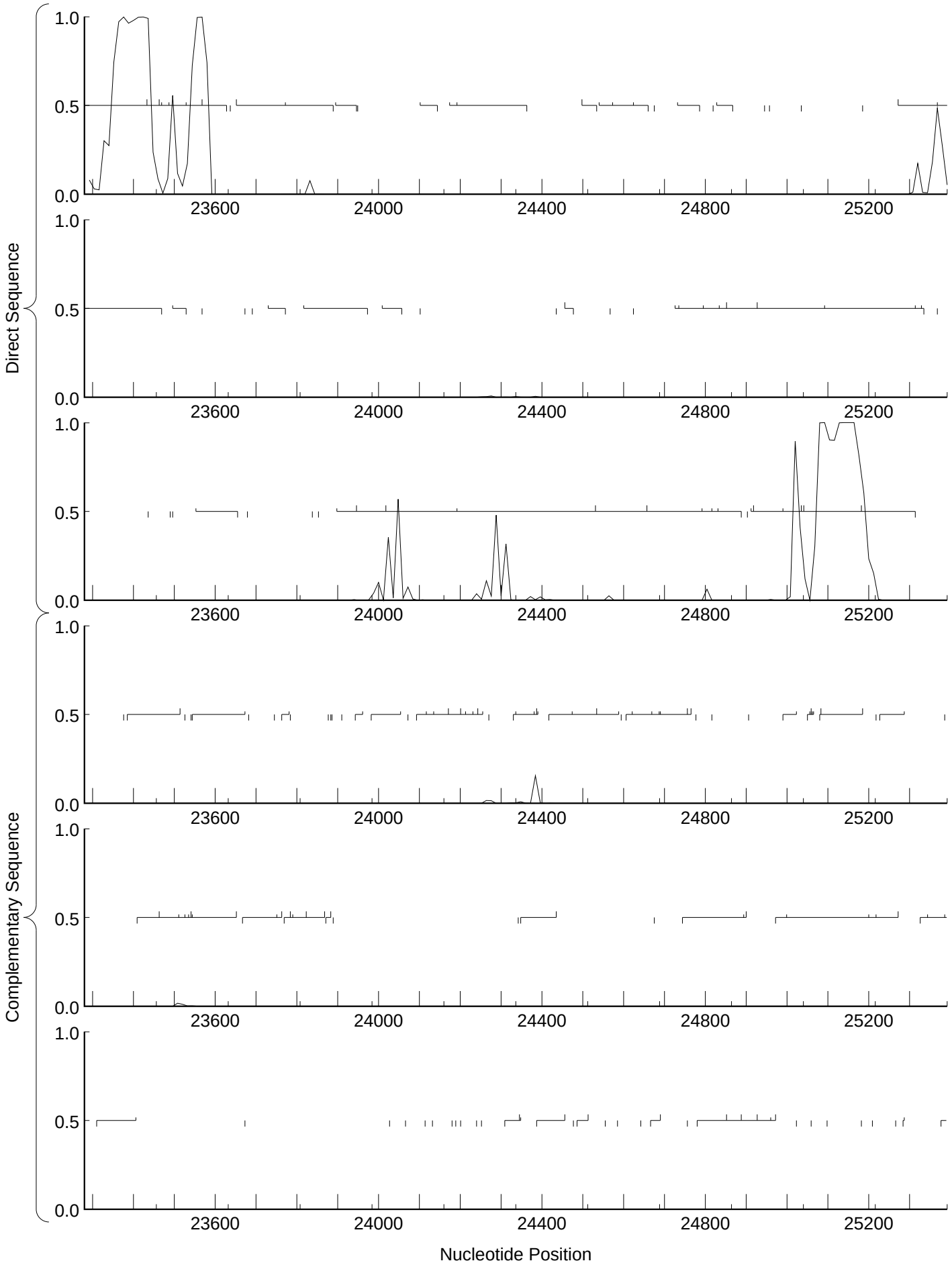
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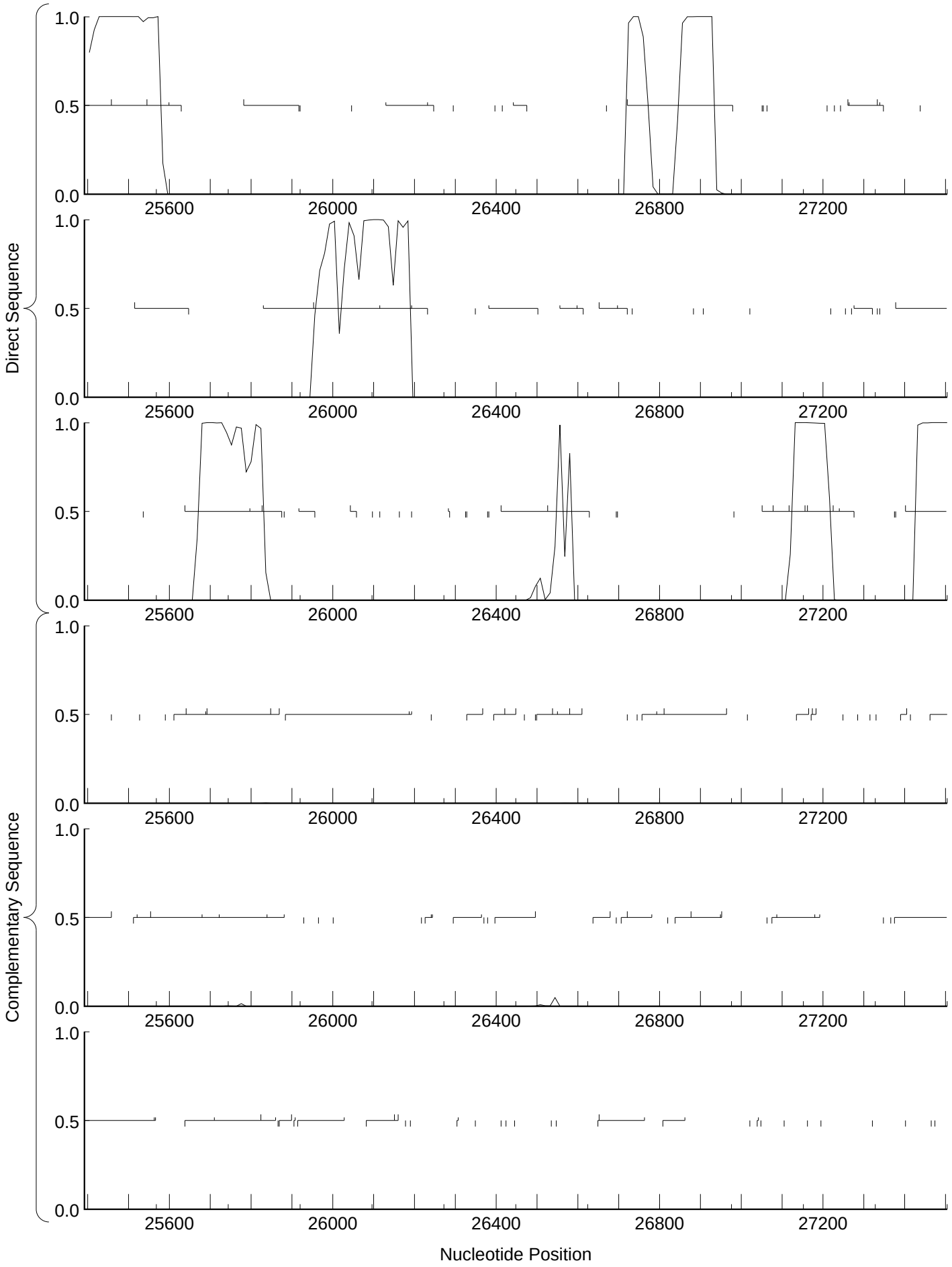


ya complete sequence, 55713 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 12/28

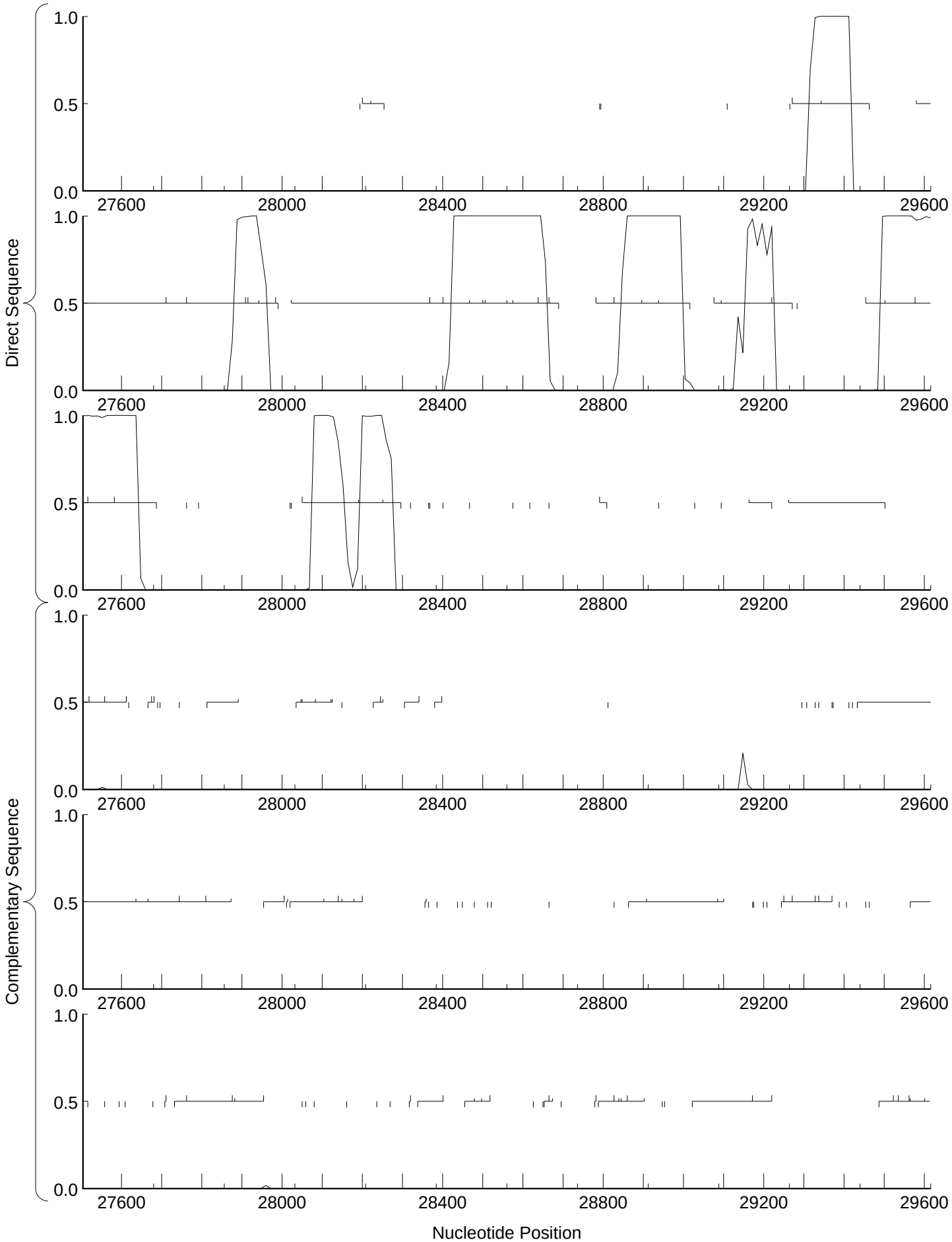


ya complete sequence, 55713 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 13/28

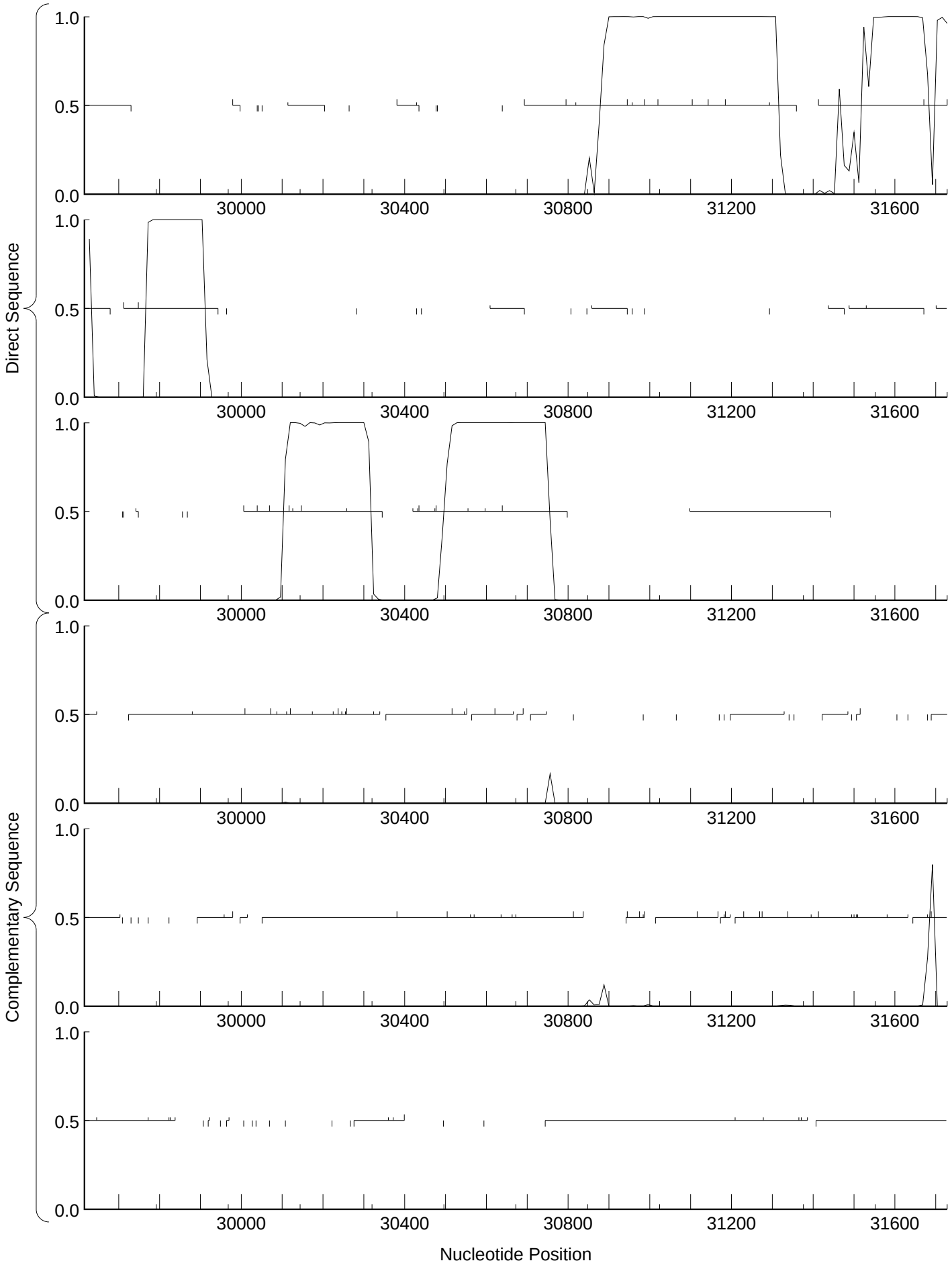



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
25600
26000
26400
26800
27200
Nucleotide Position

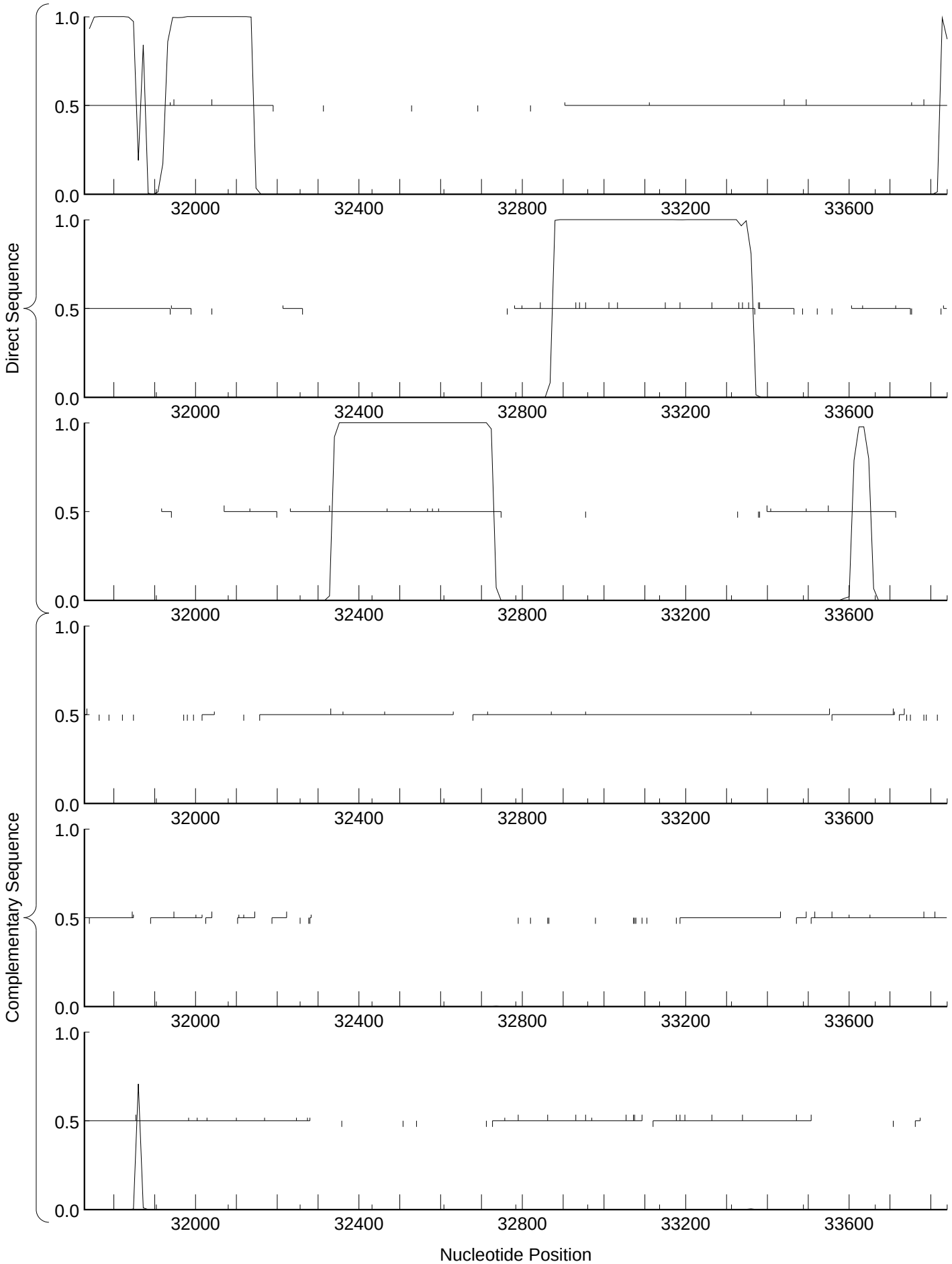
ya complete sequence, 55713 bp including 9-base 3' overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 15/28



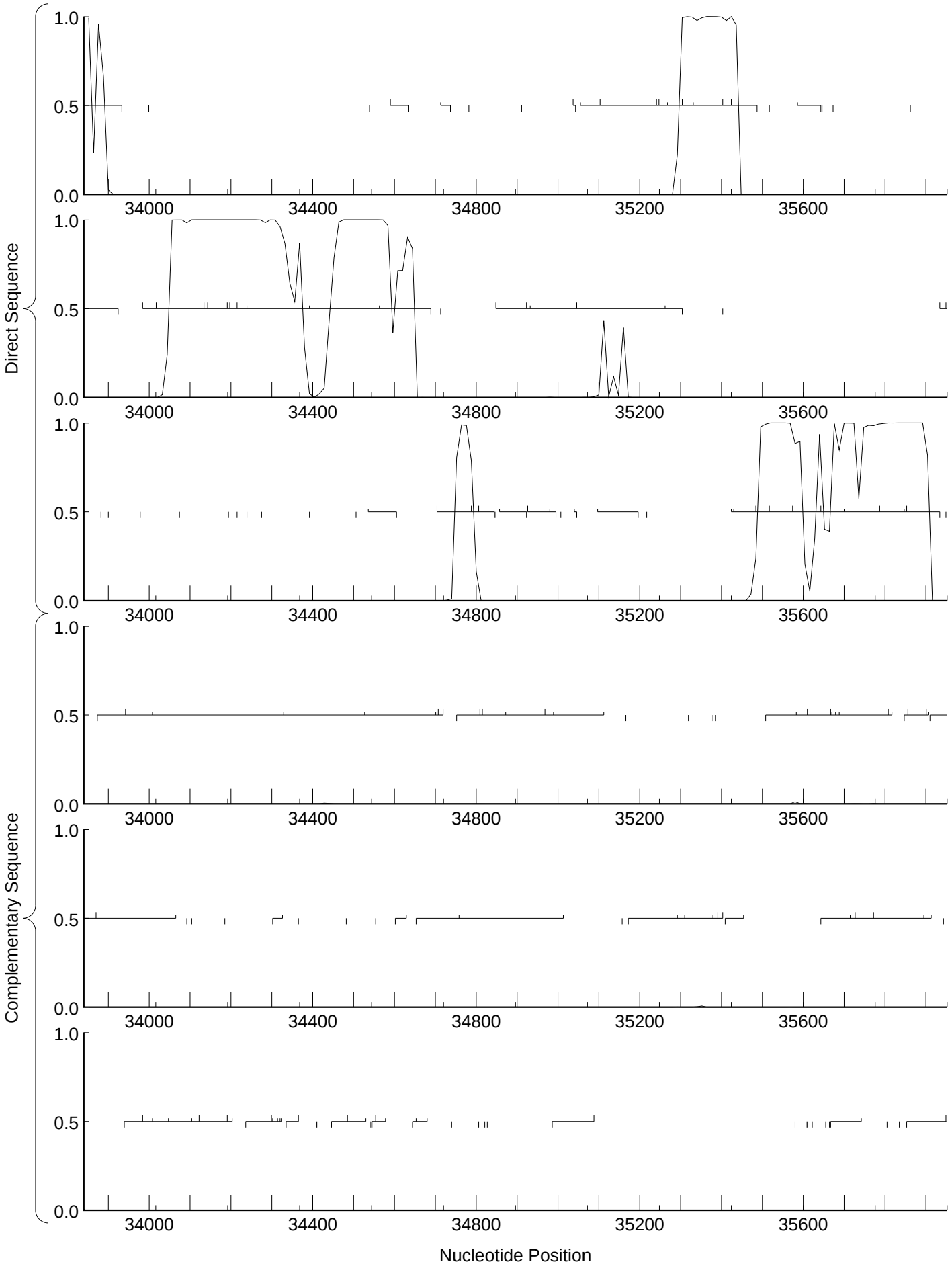
ya complete sequence, 55713 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 16/28



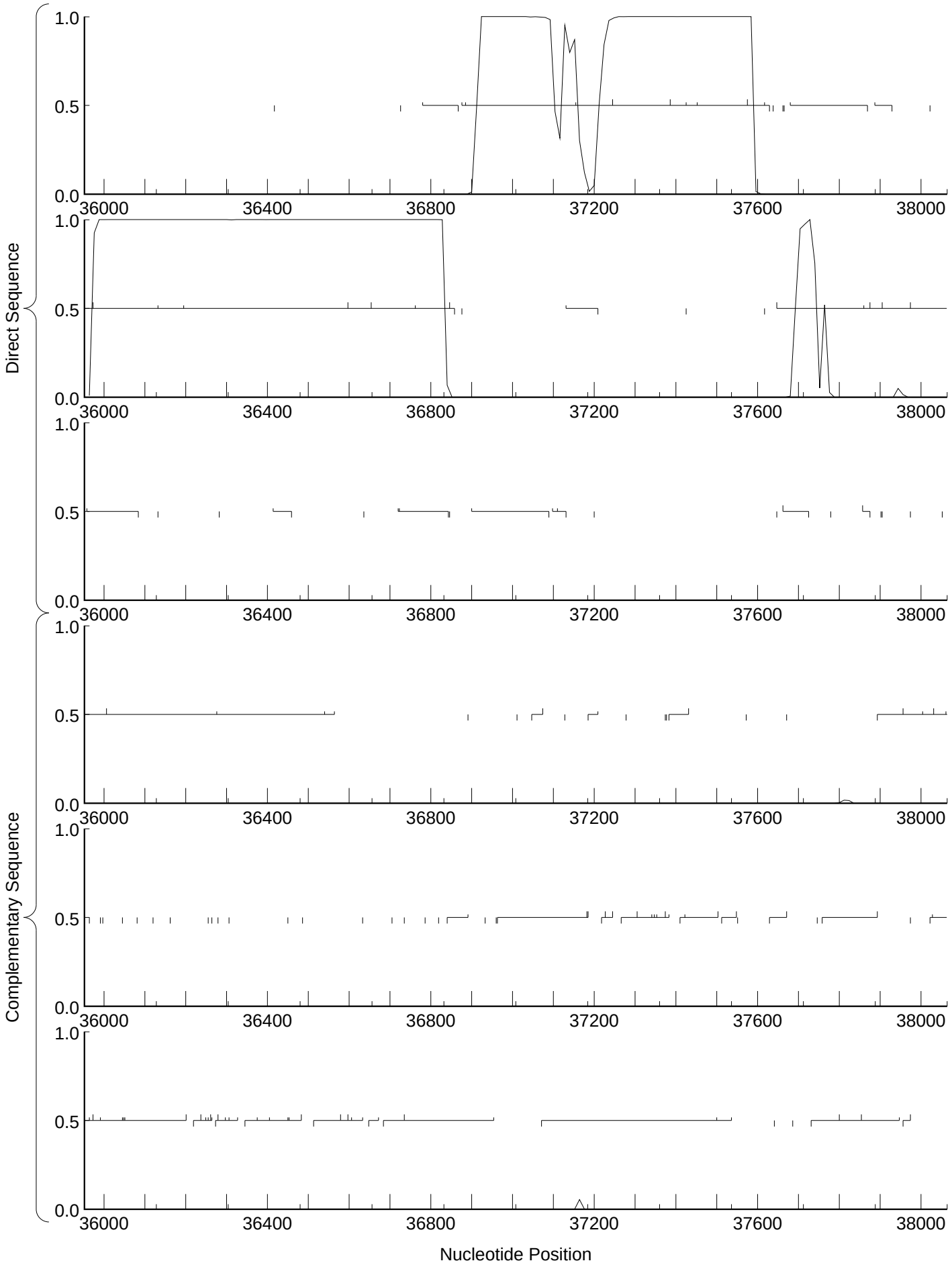
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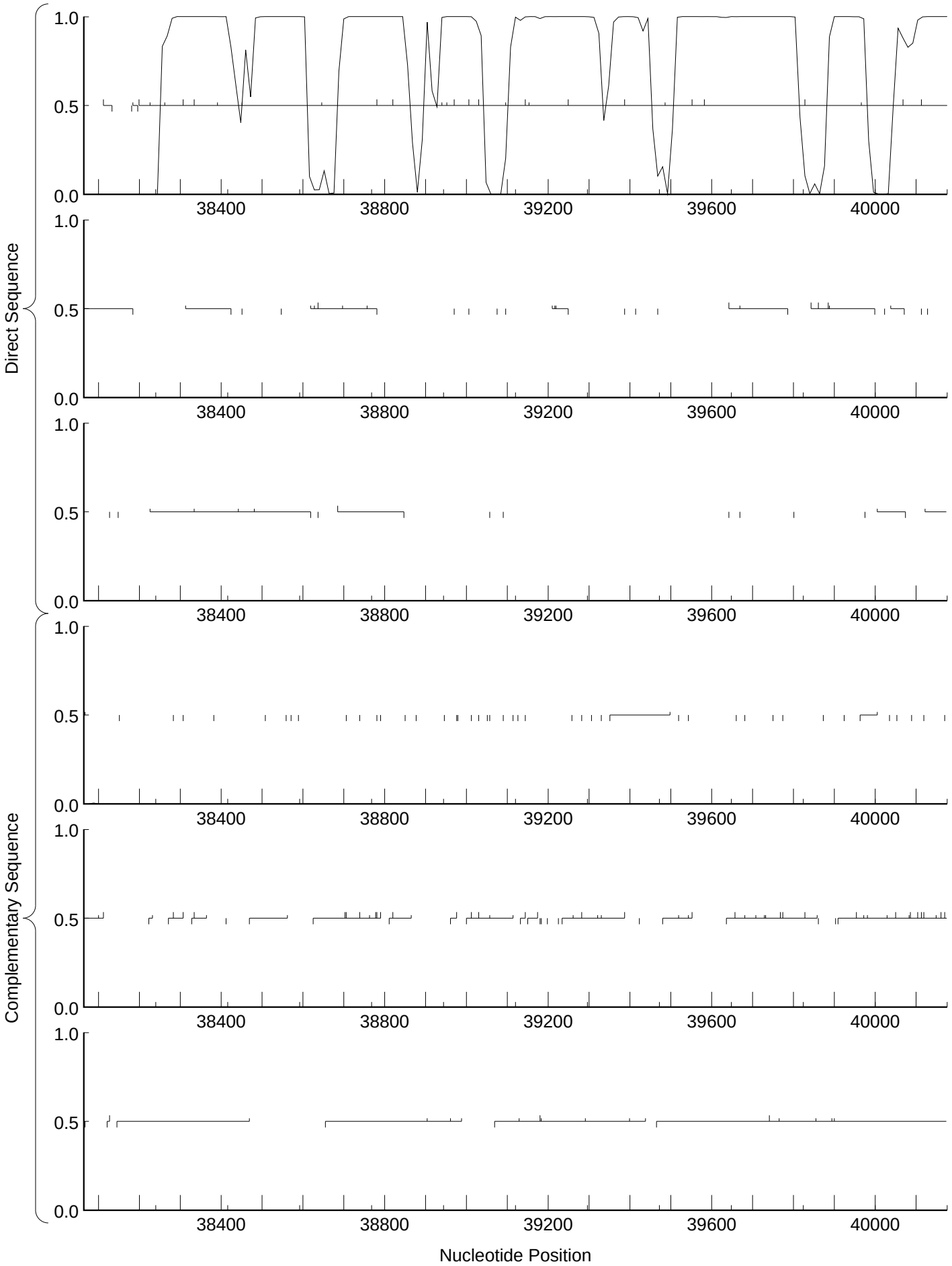
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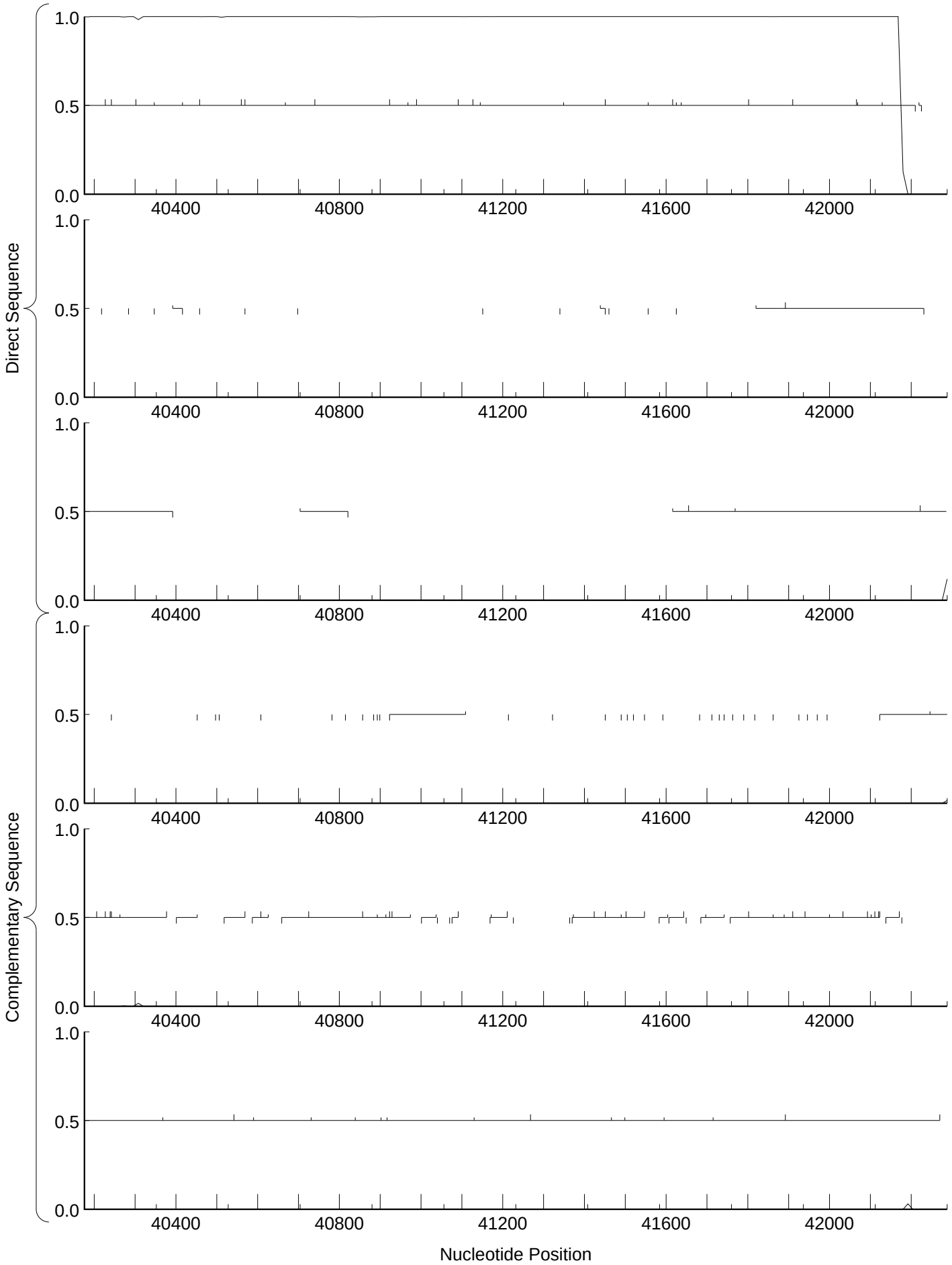
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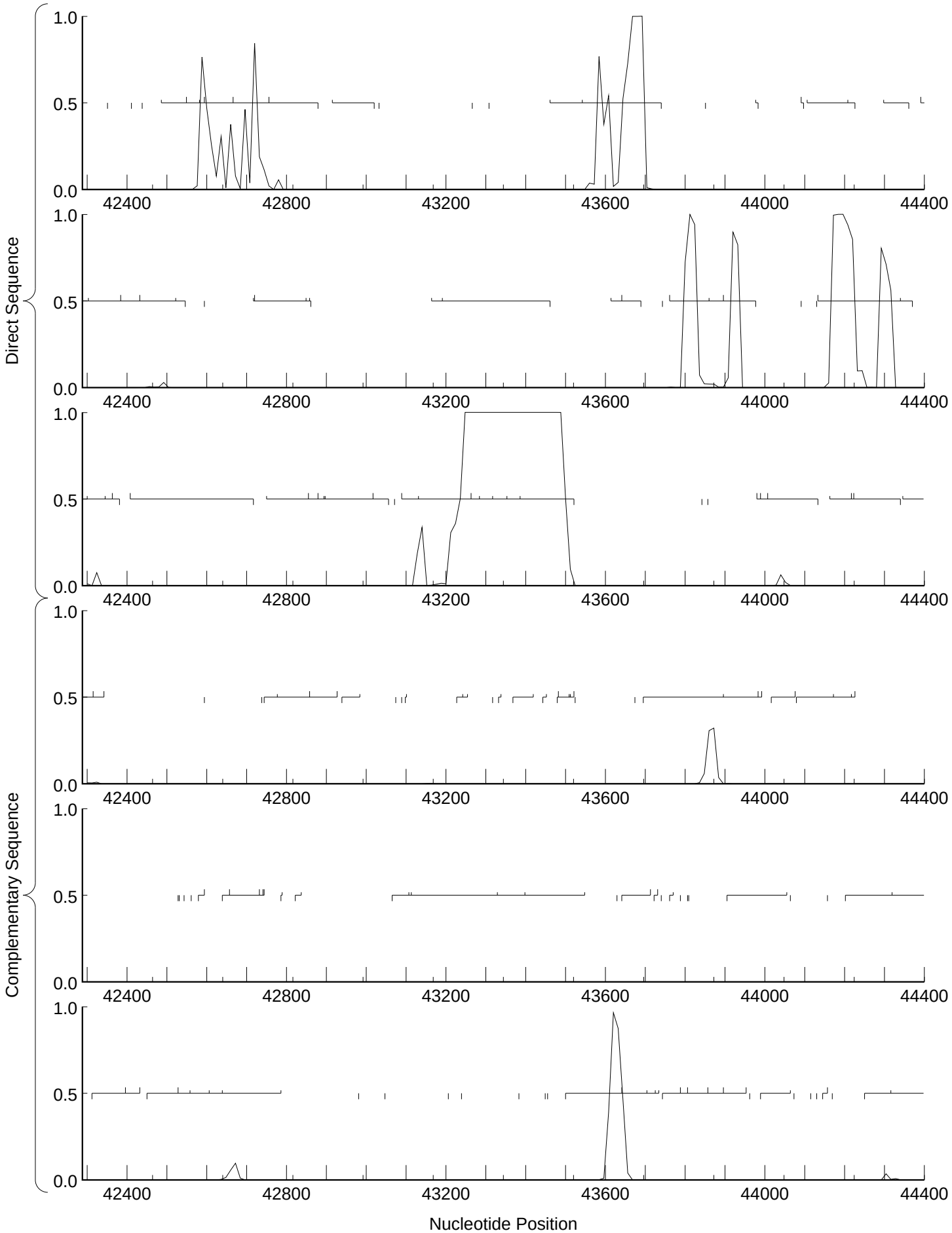
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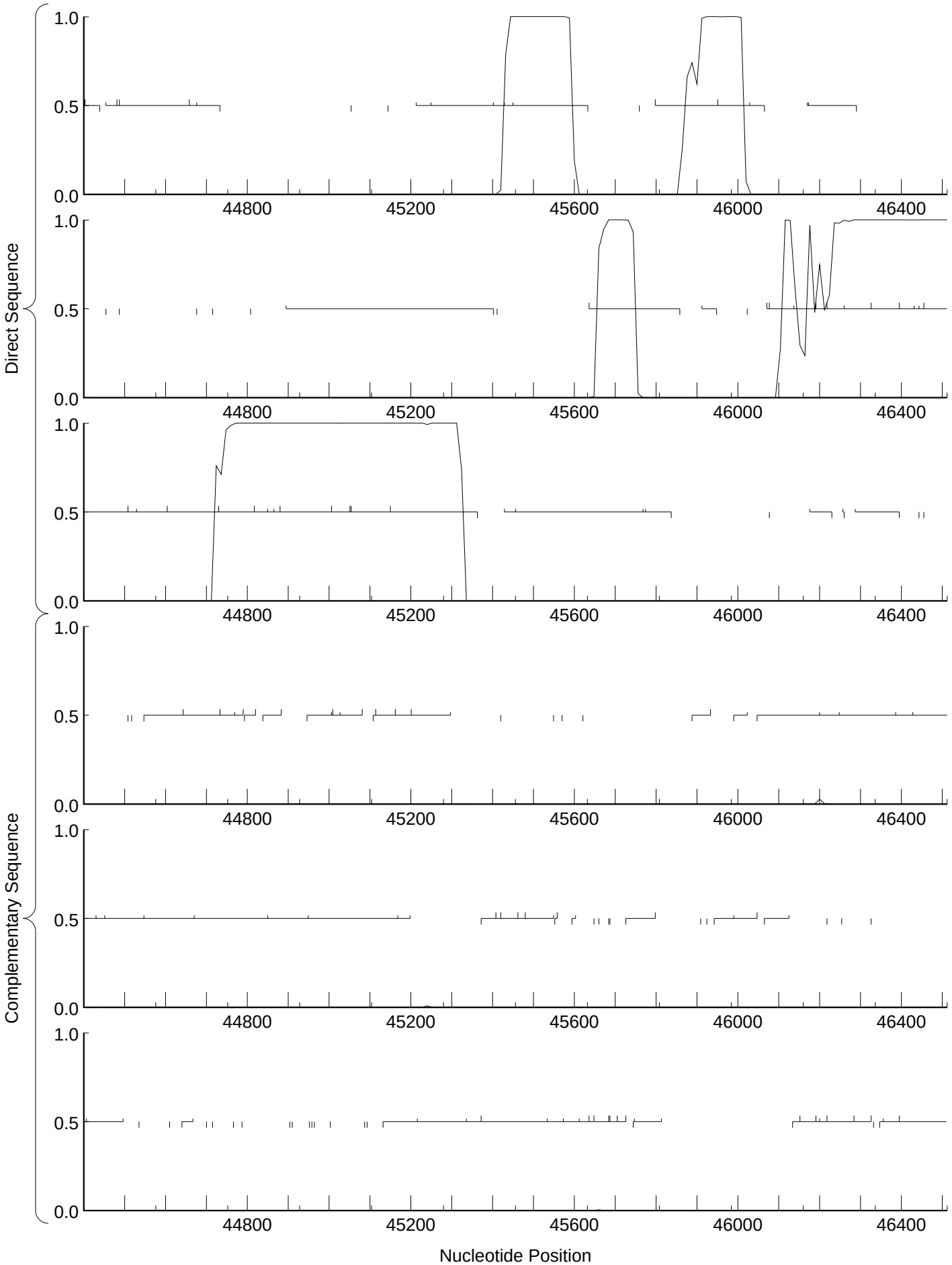
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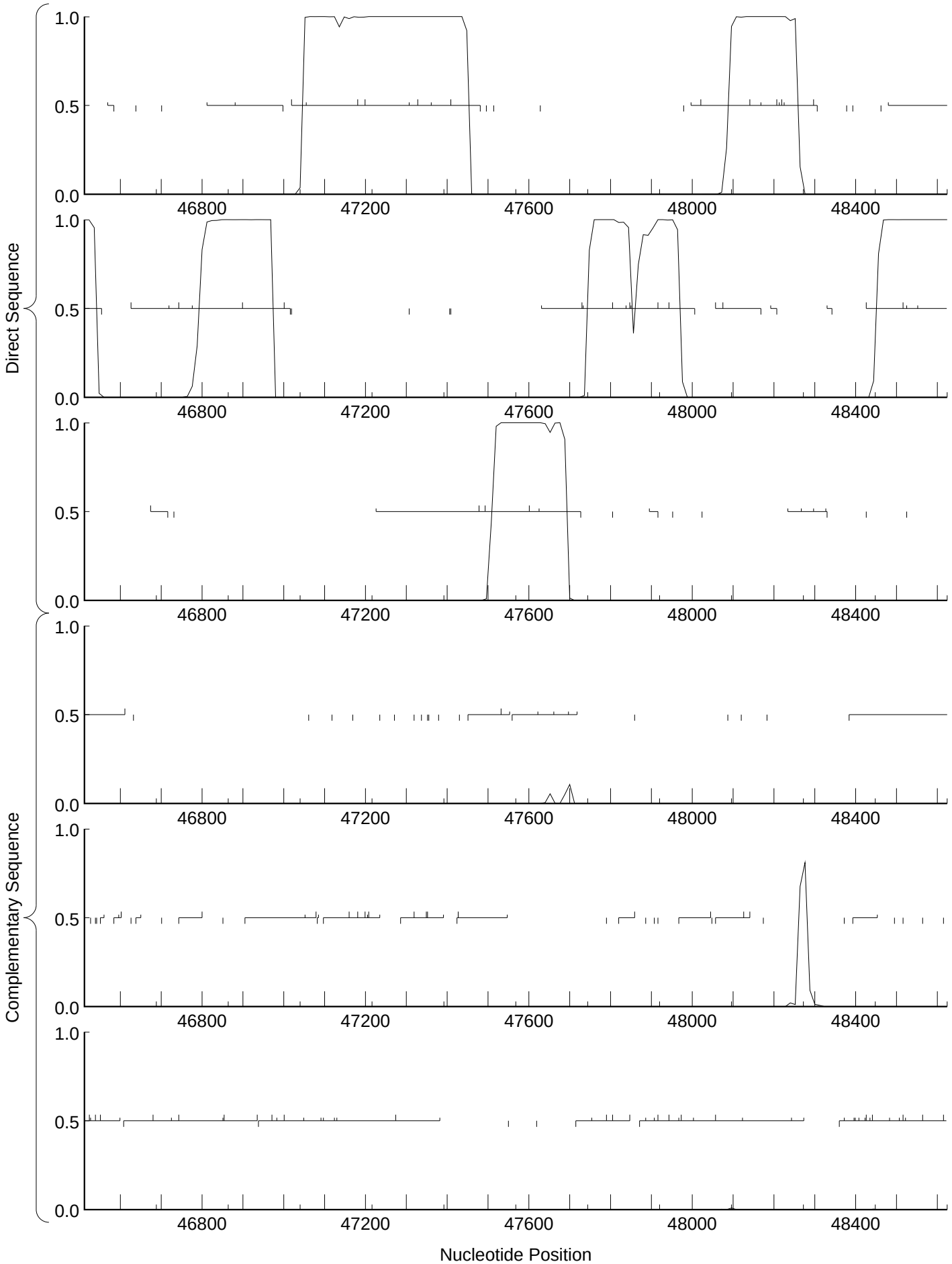
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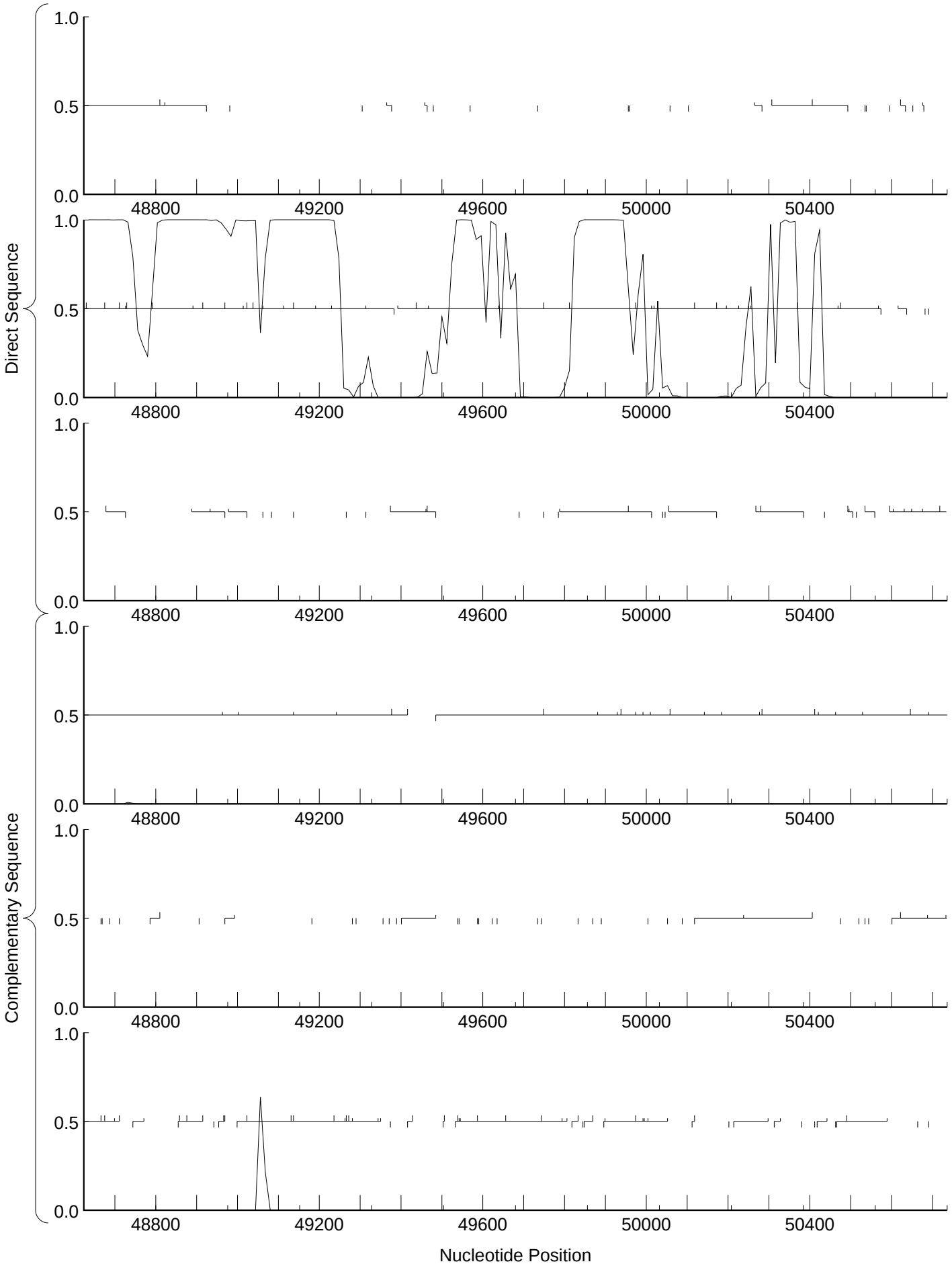
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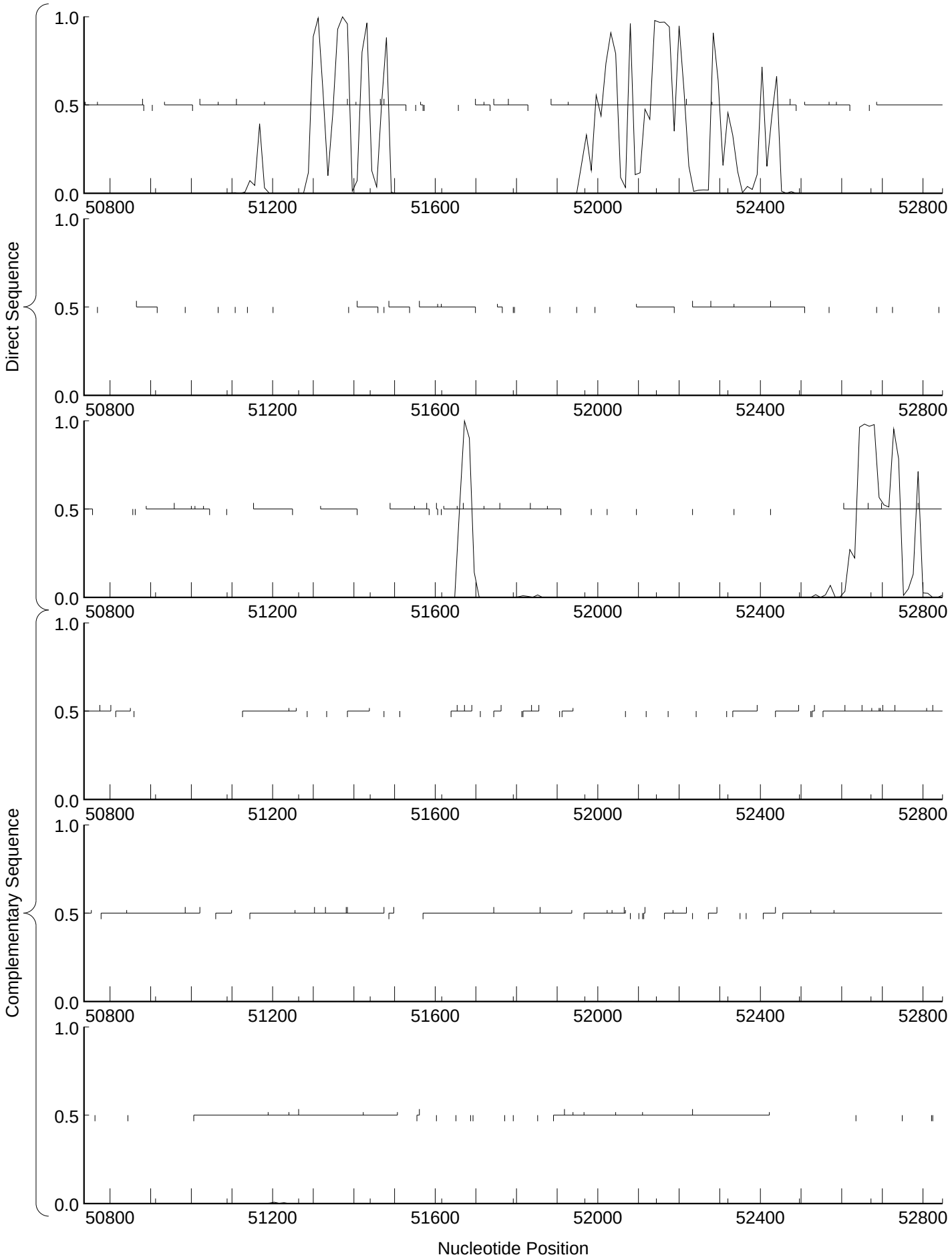


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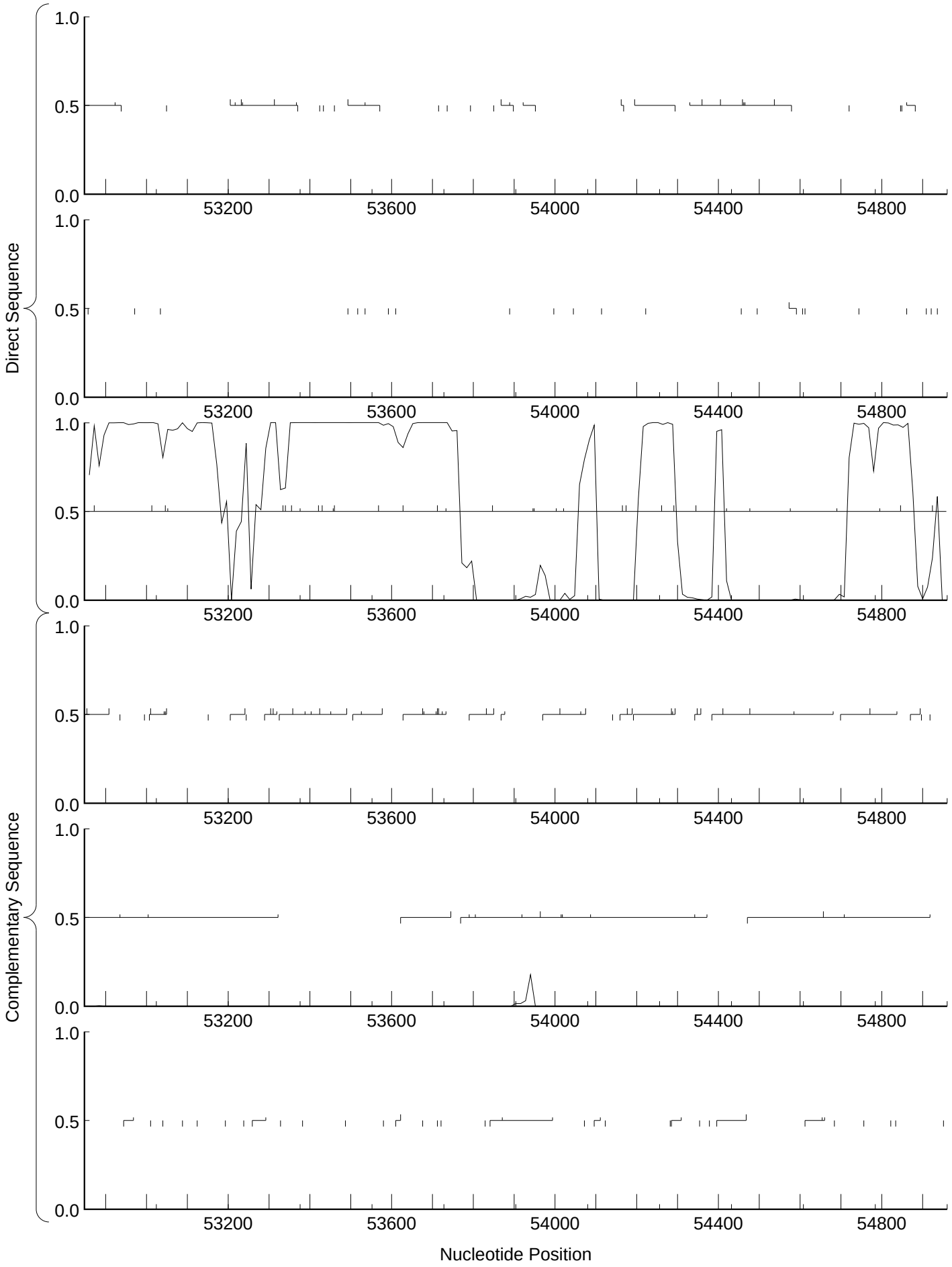


ya complete sequence, 55713 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 25/28




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
50800
51200
51600
52000
52400
52800
Nucleotide Position

ya complete sequence, 55713 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 27/28



ya complete sequence, 55713 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI1, Order 4, Window 96, Step 12, 28/28

