GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage BobaPhett complete sequence, 59815 bp including 10-base 3’ overhang (CG
Analysis Date    :  3/28/24 at 10:49:19
Pages            :  30
Sequence Length  :  59815 bp
GC Content       :   61.33%

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

PS-Version       :  1.2

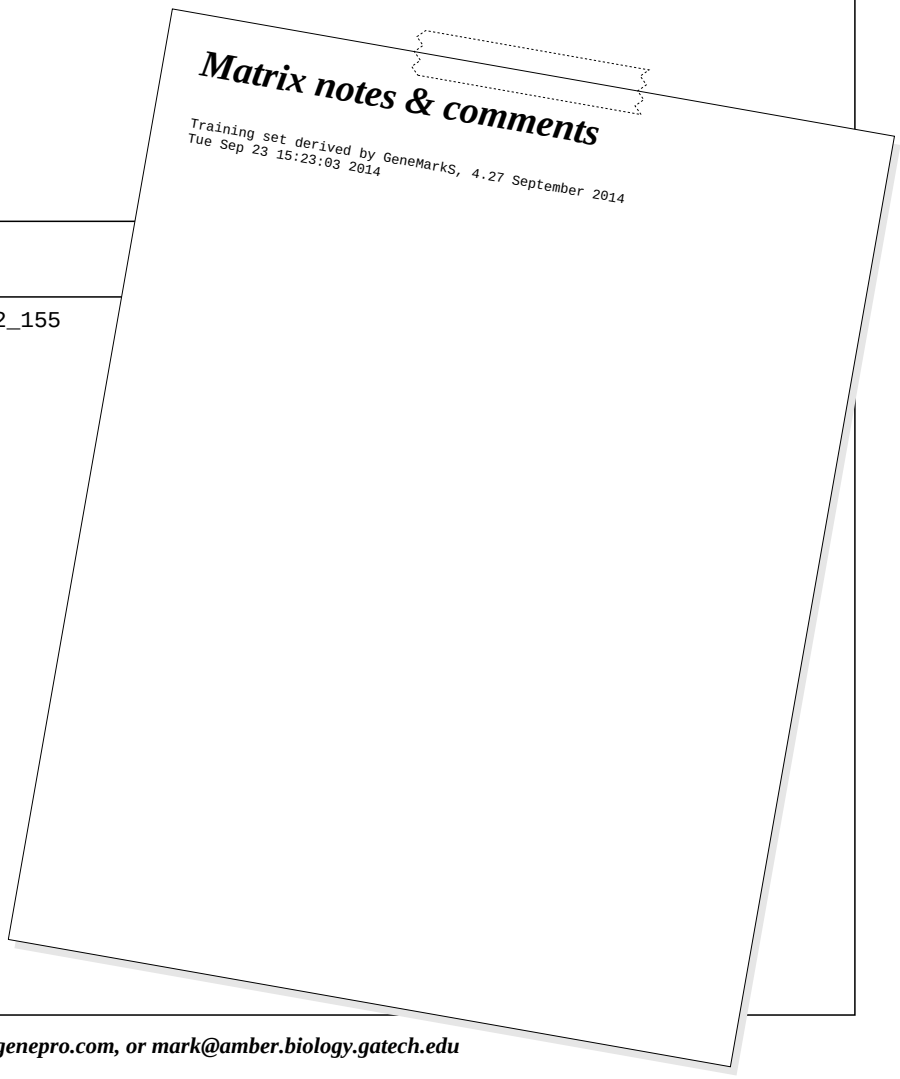
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

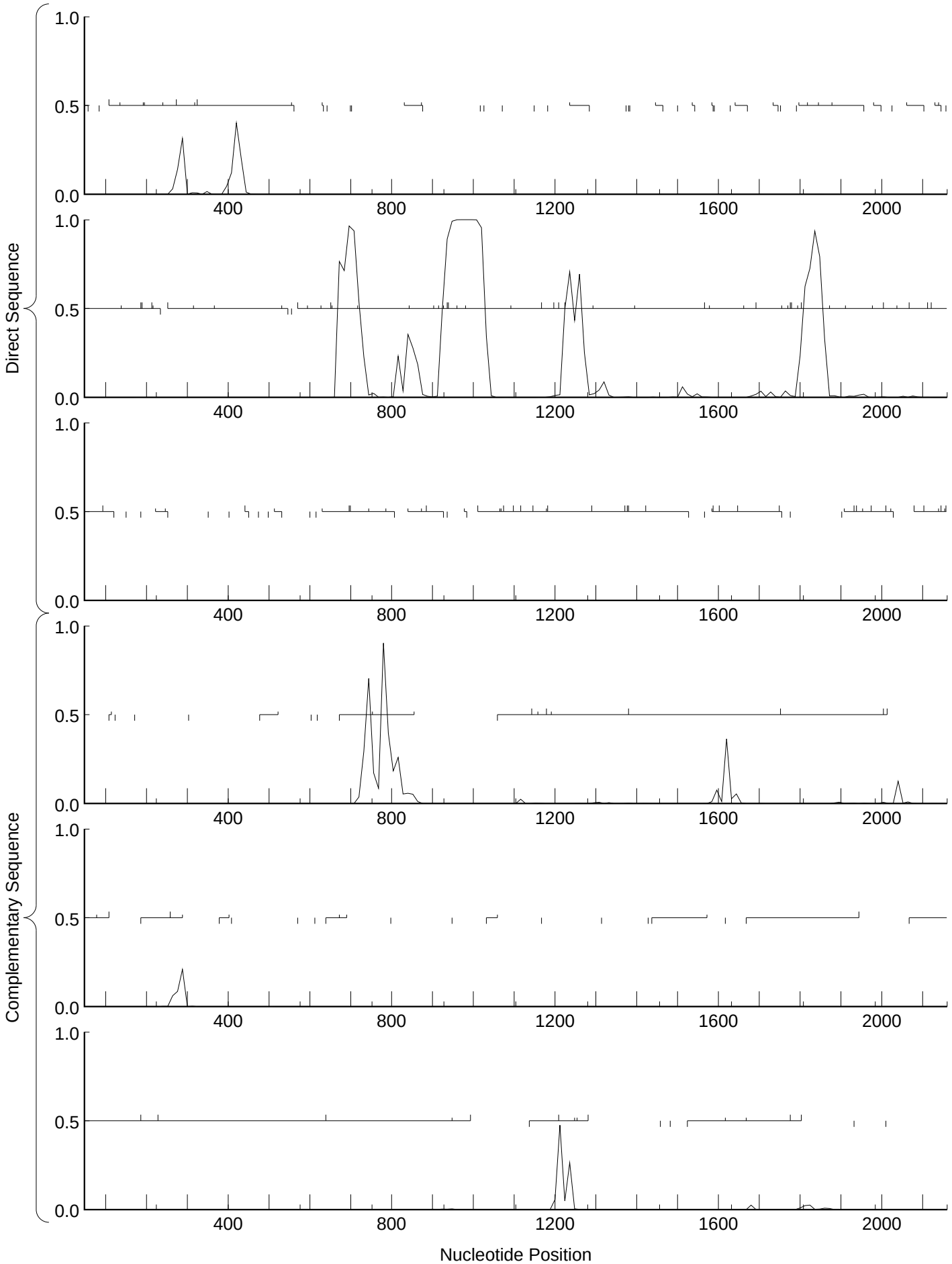
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Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

Order      :  4

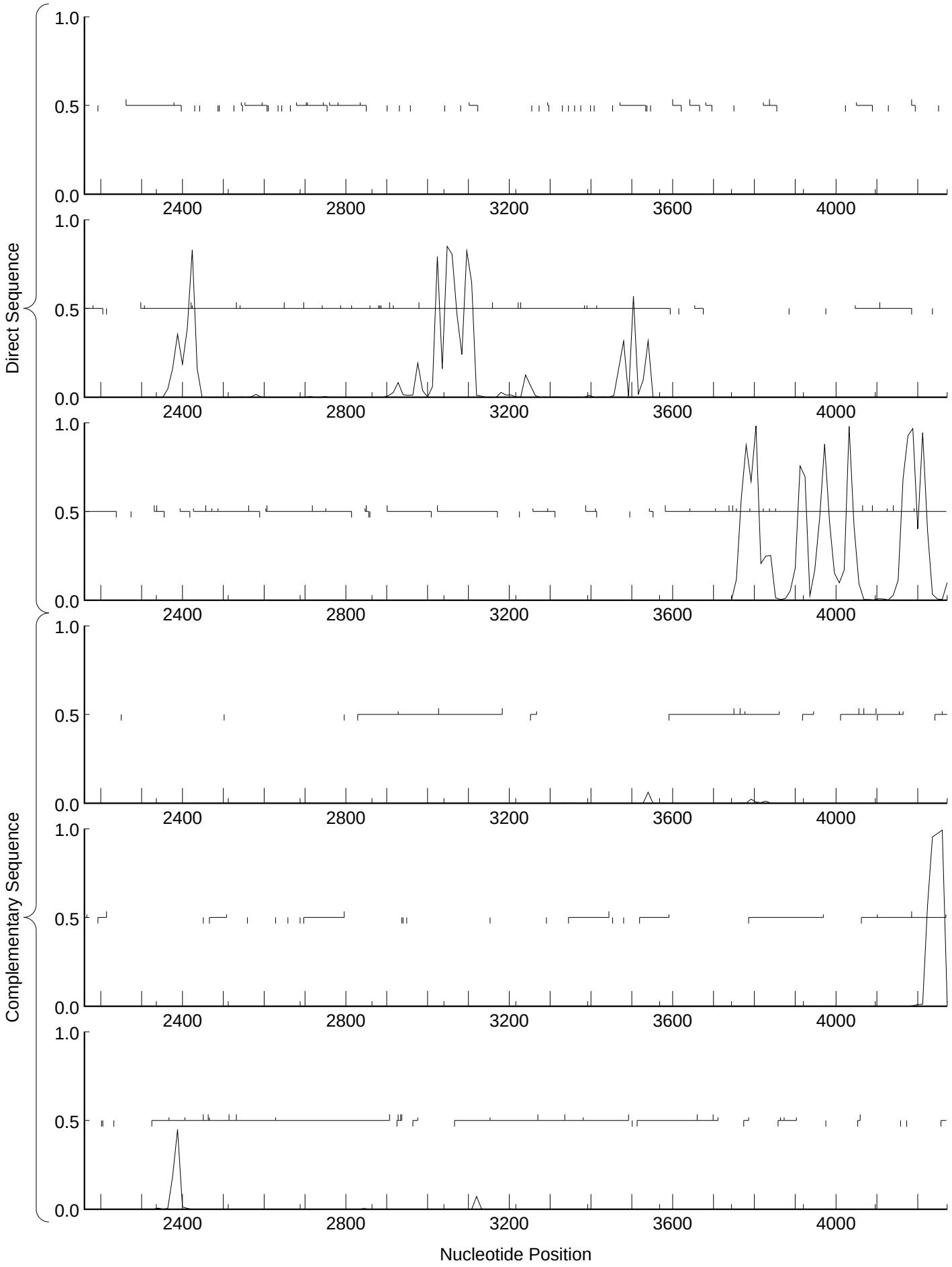

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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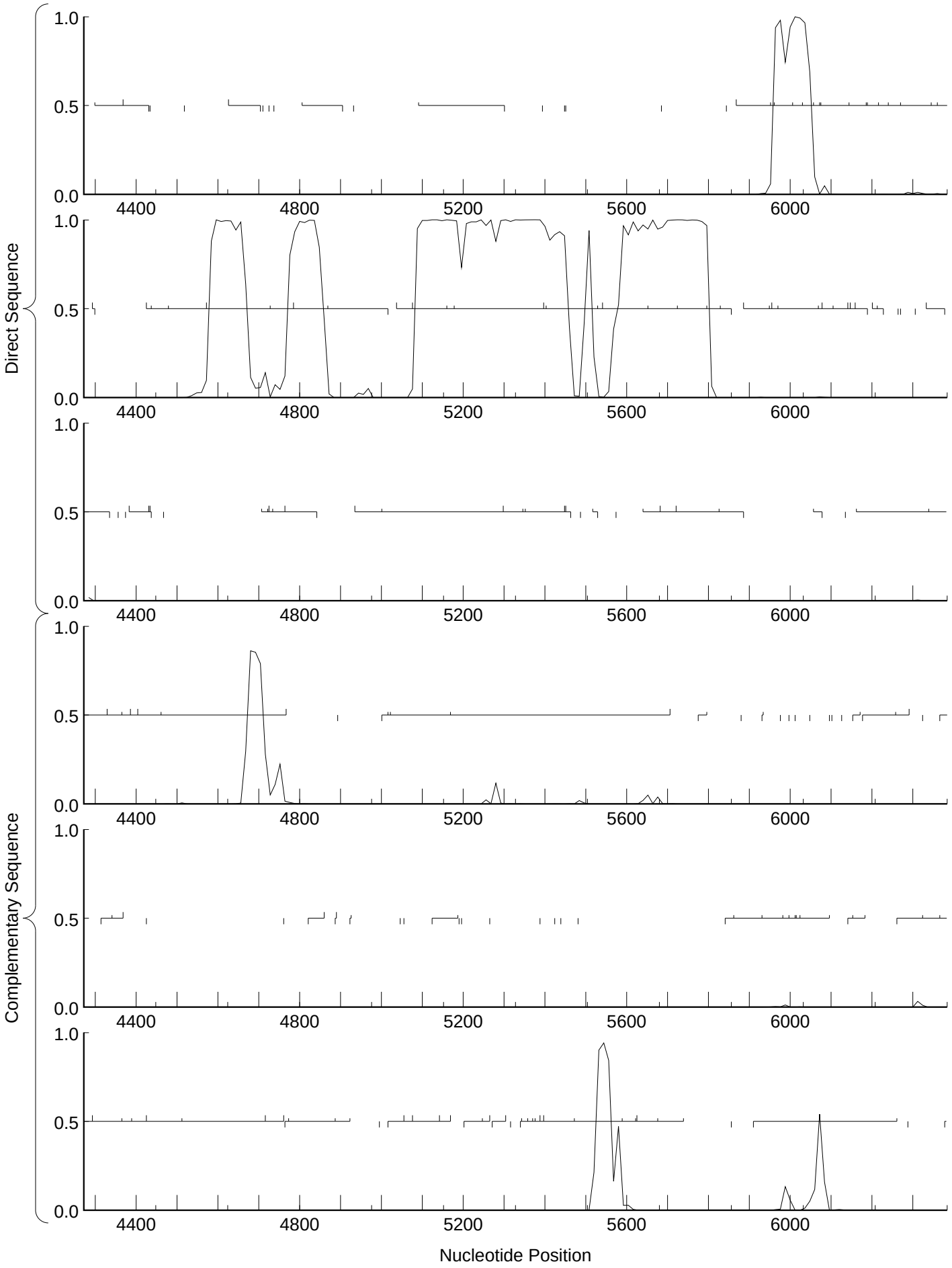
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

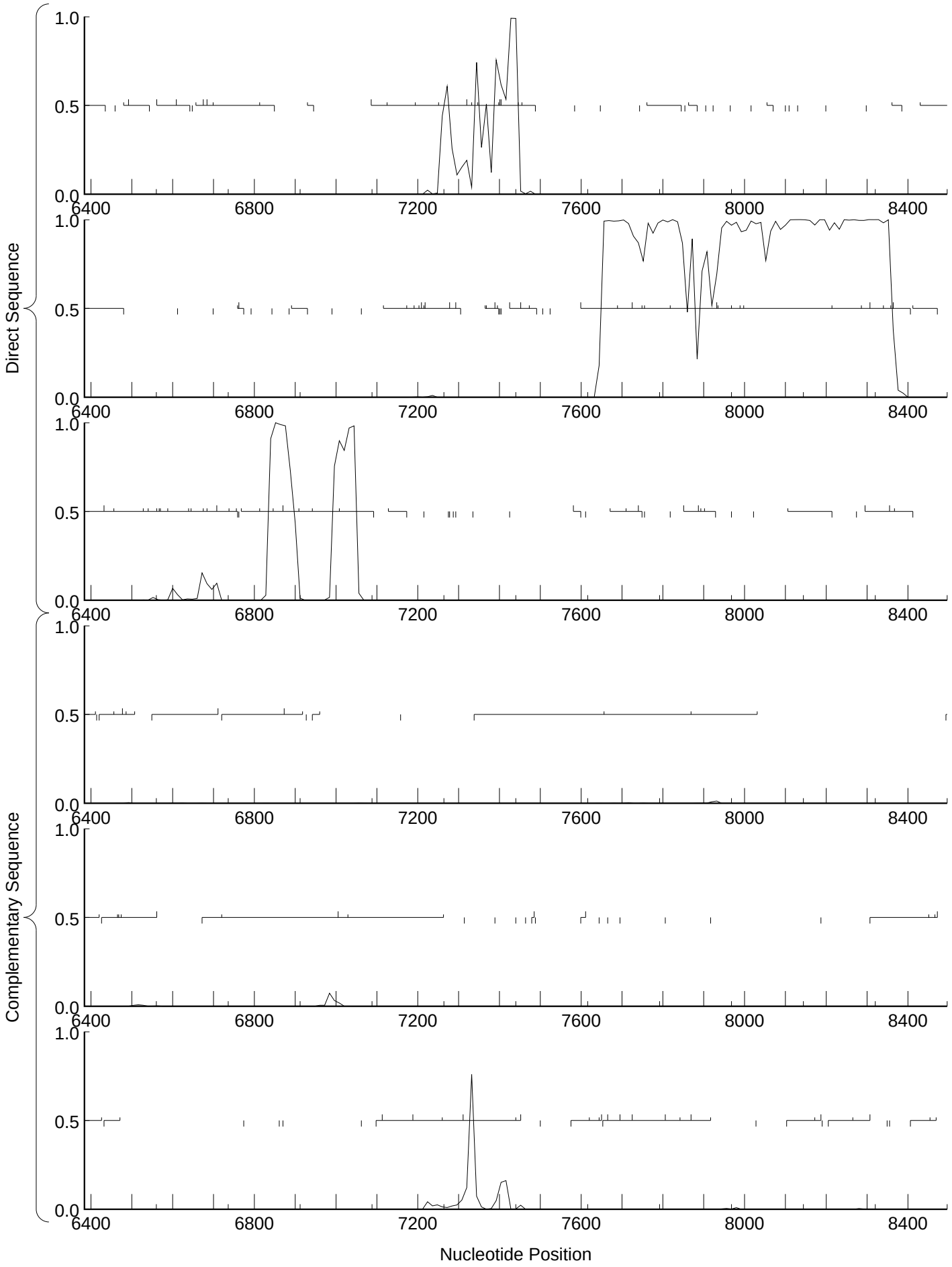

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1200
1600
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Nucleotide Position

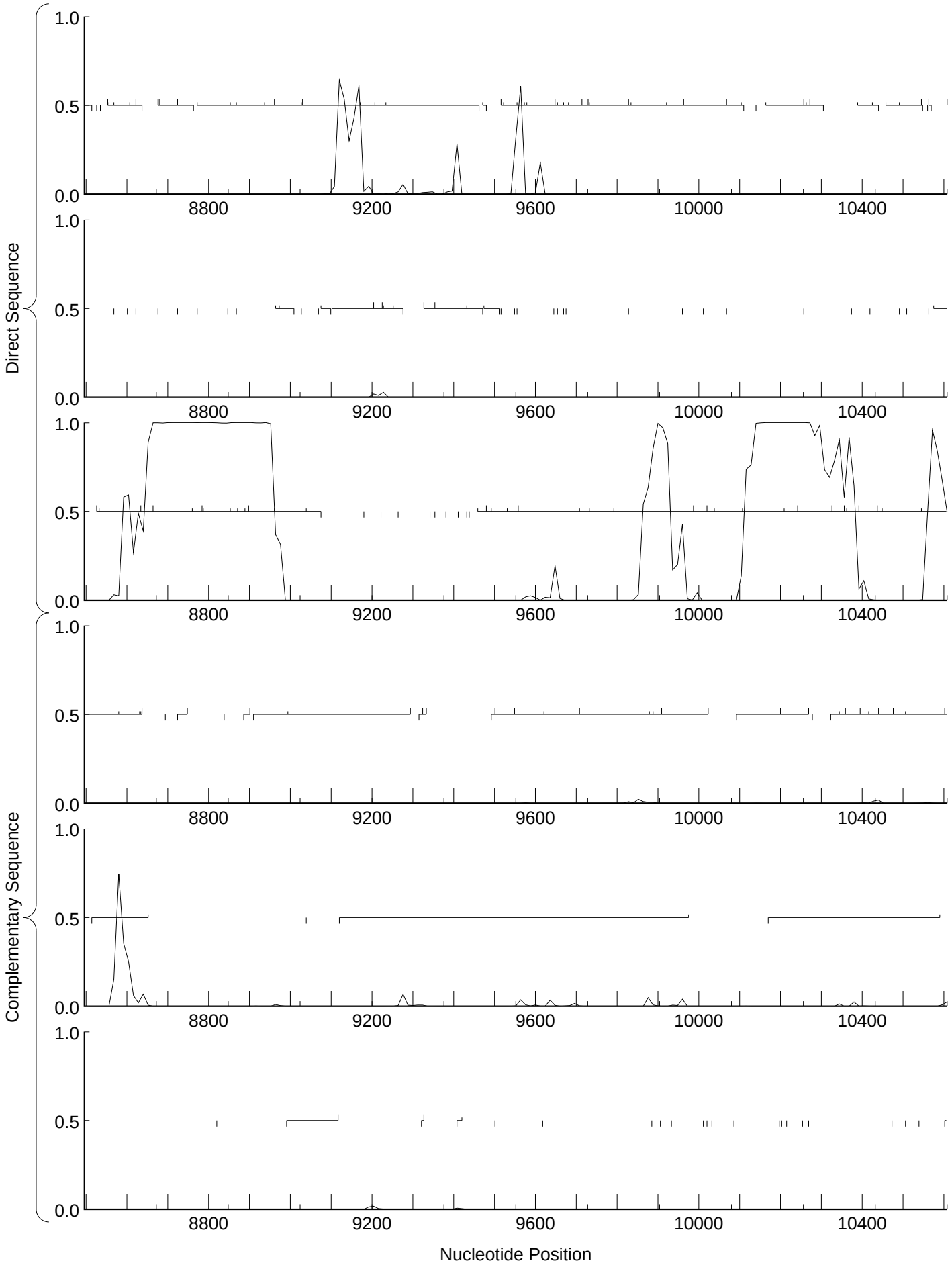
tt complete sequence, 59815 bp including 10-base 3’ overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 3/30

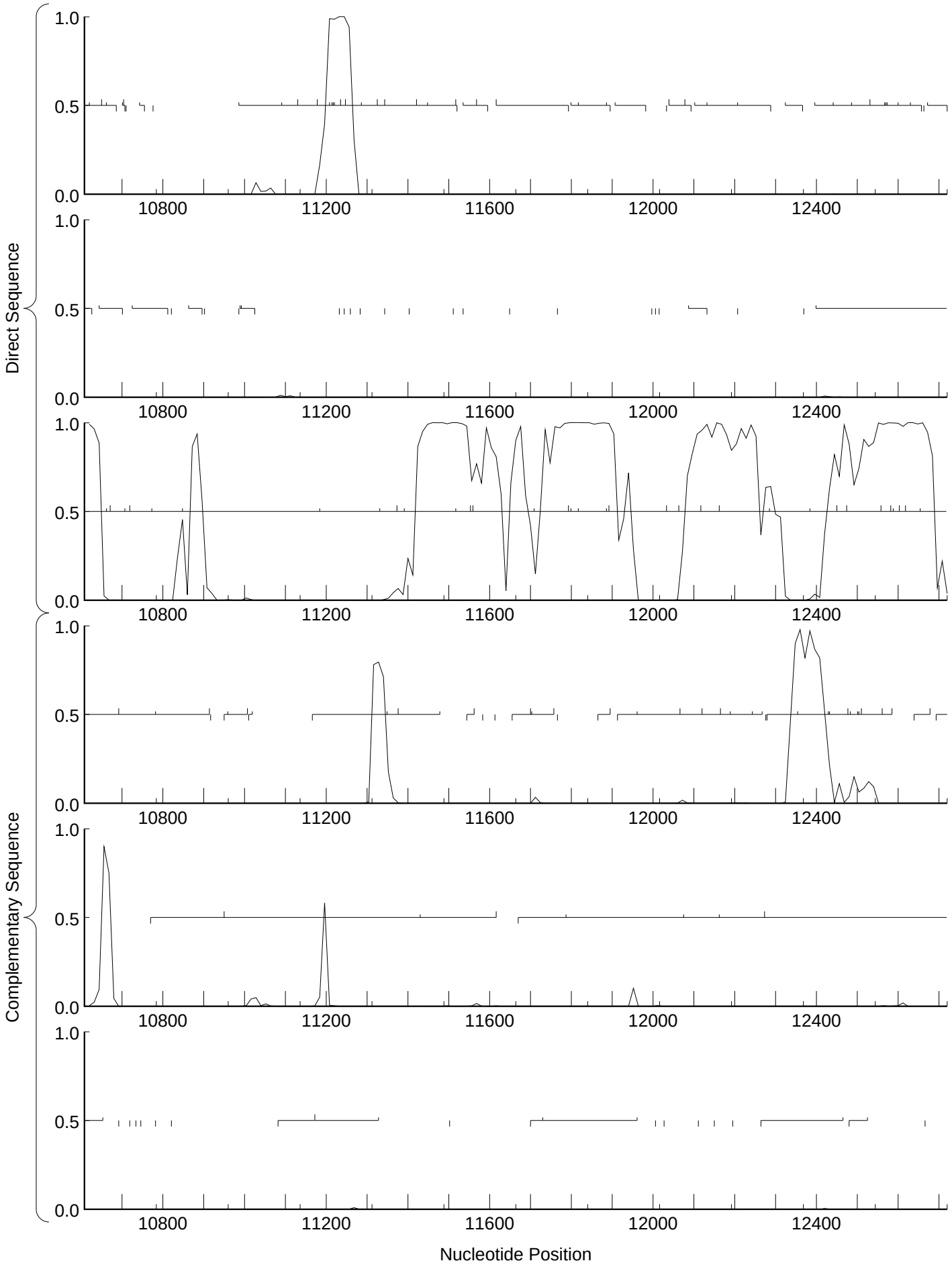


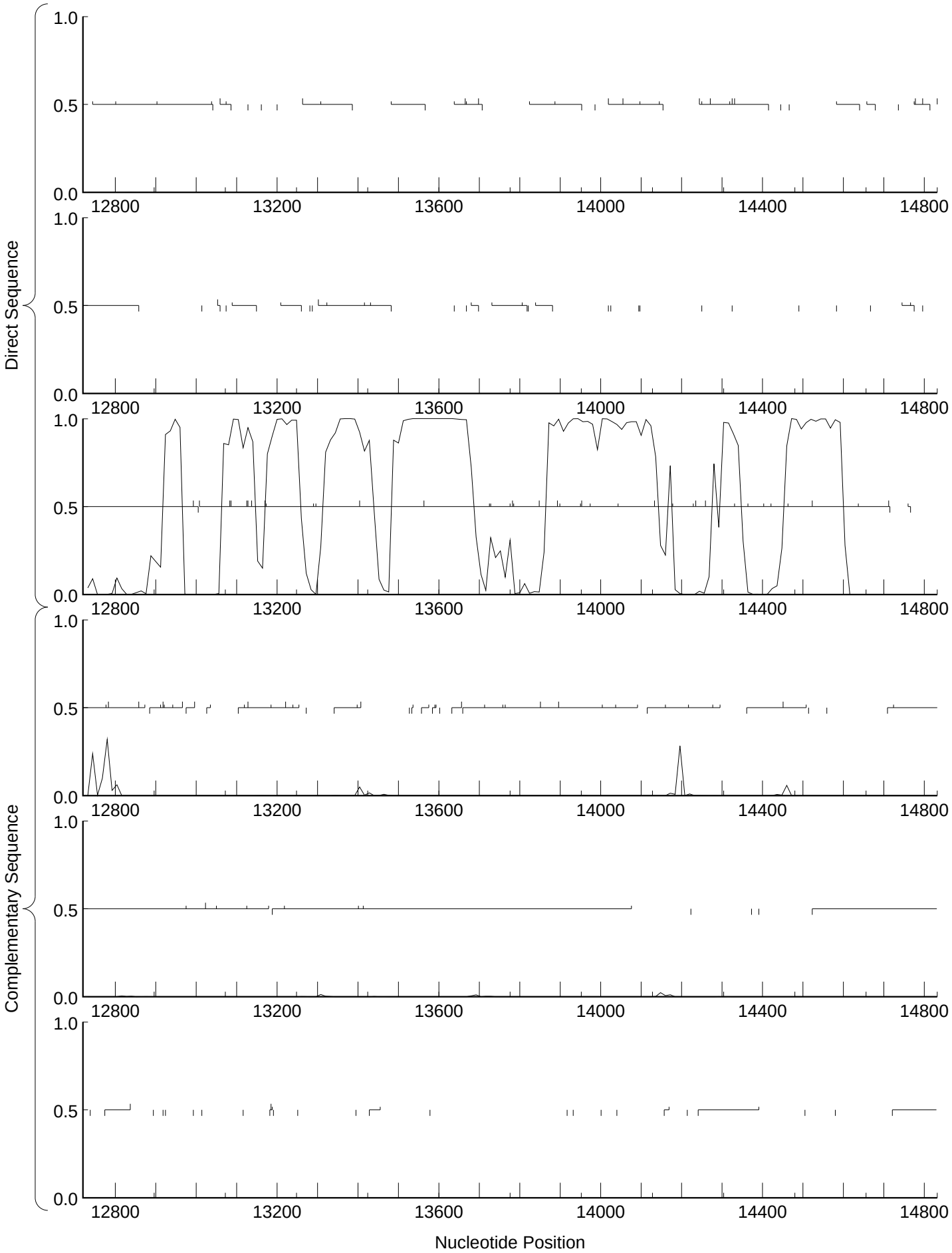
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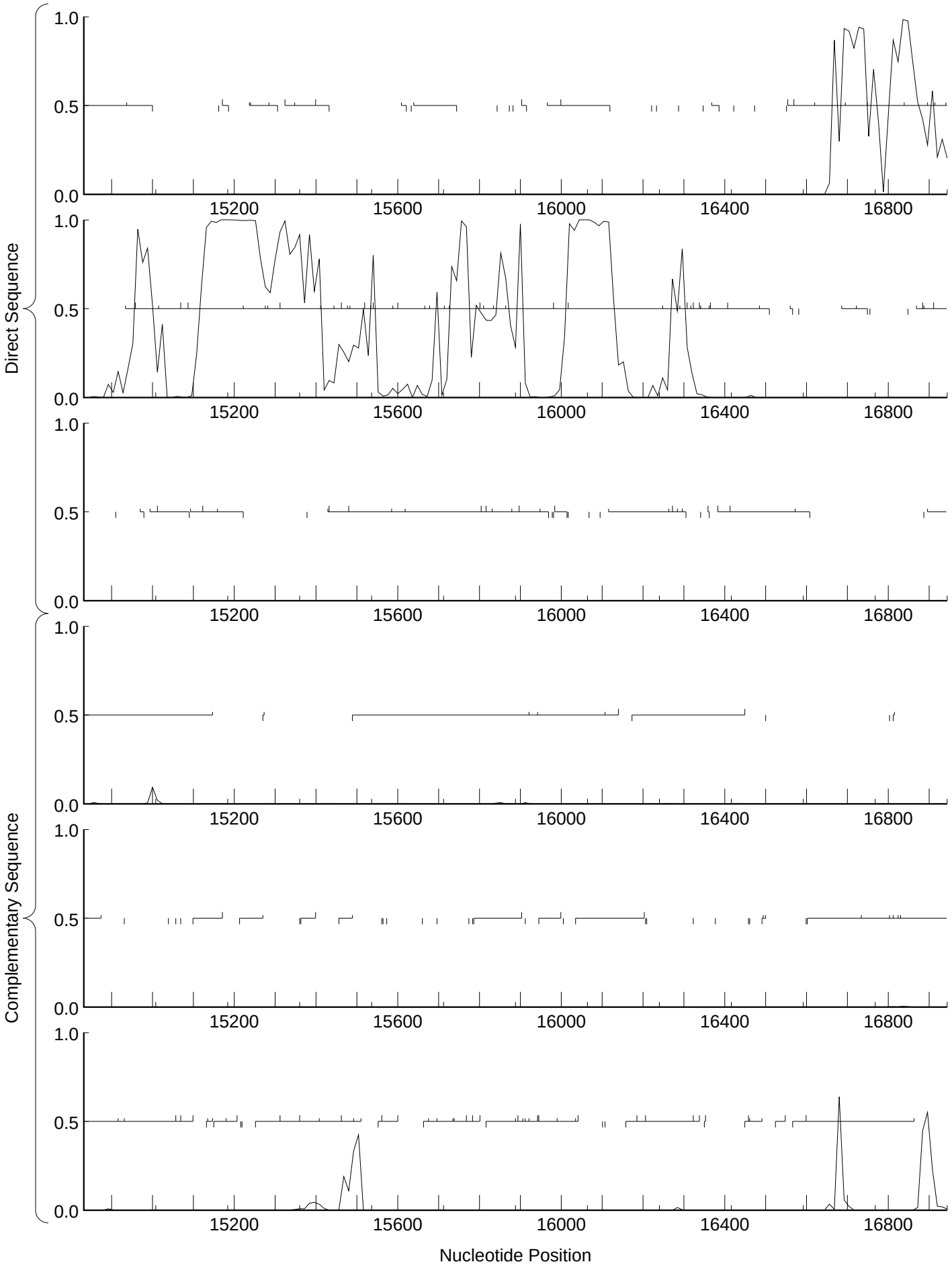

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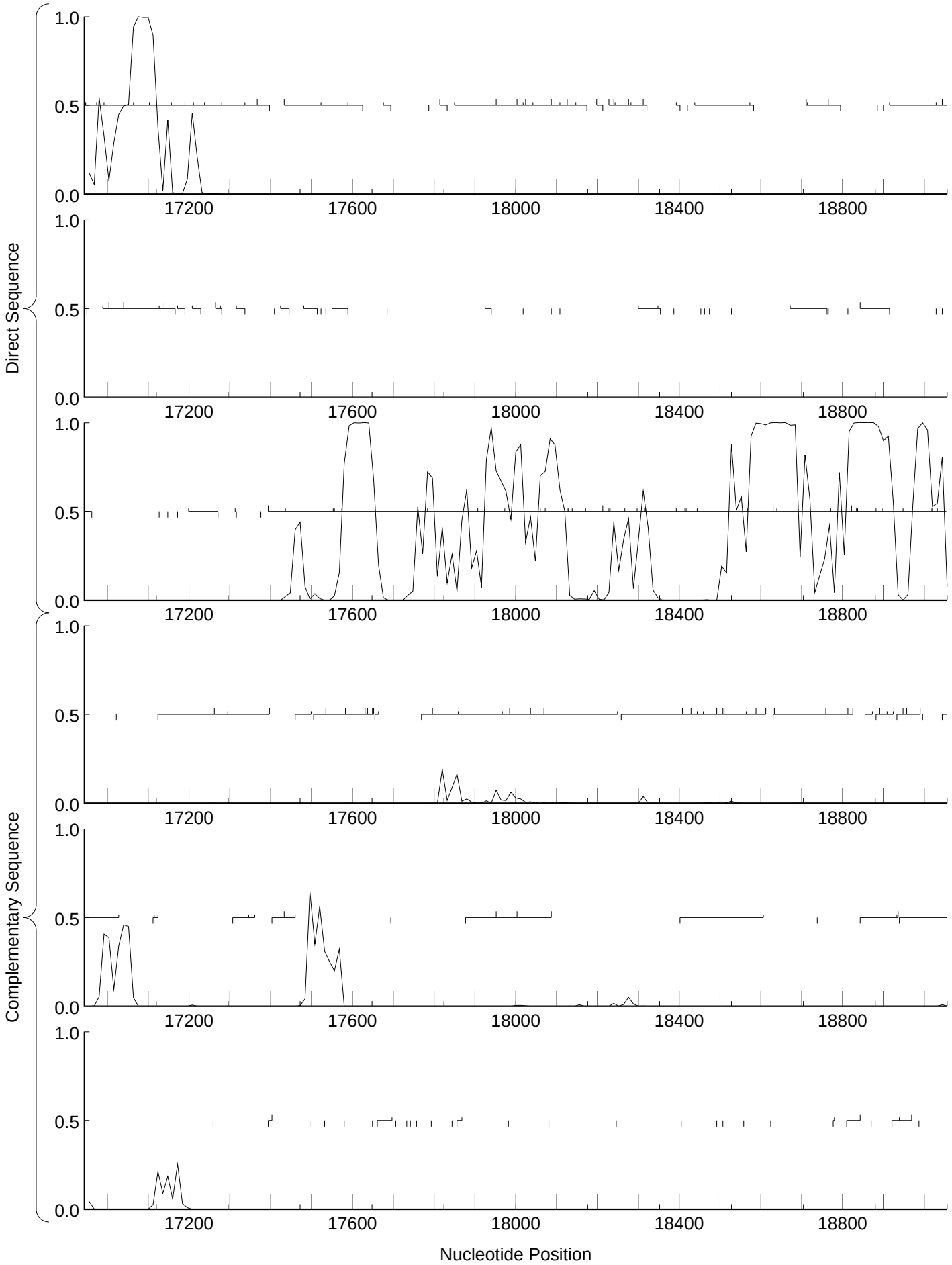

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Complementary Sequence
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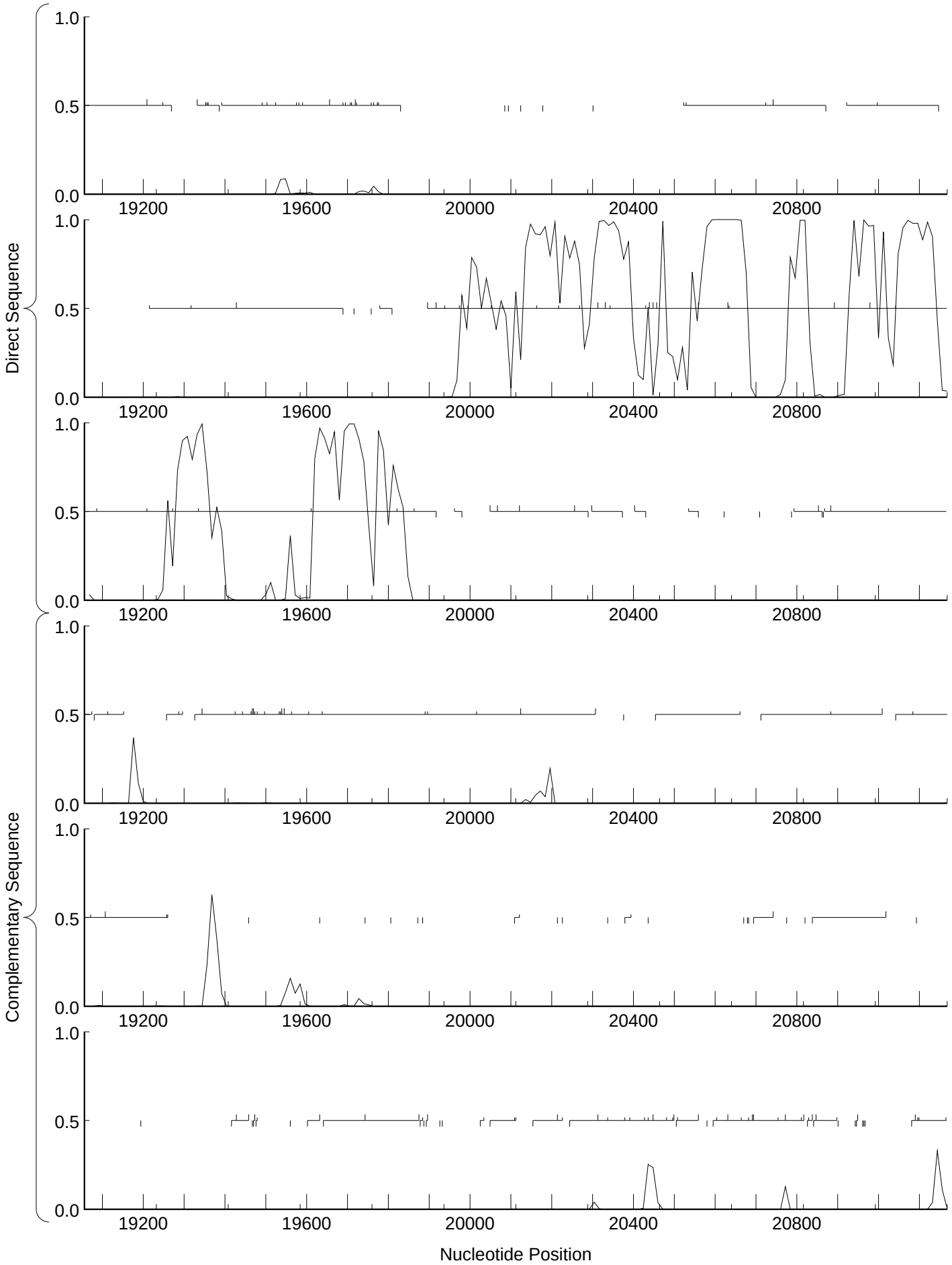
t complete sequence, 59815 bp including 10-base 3’ overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 9/30



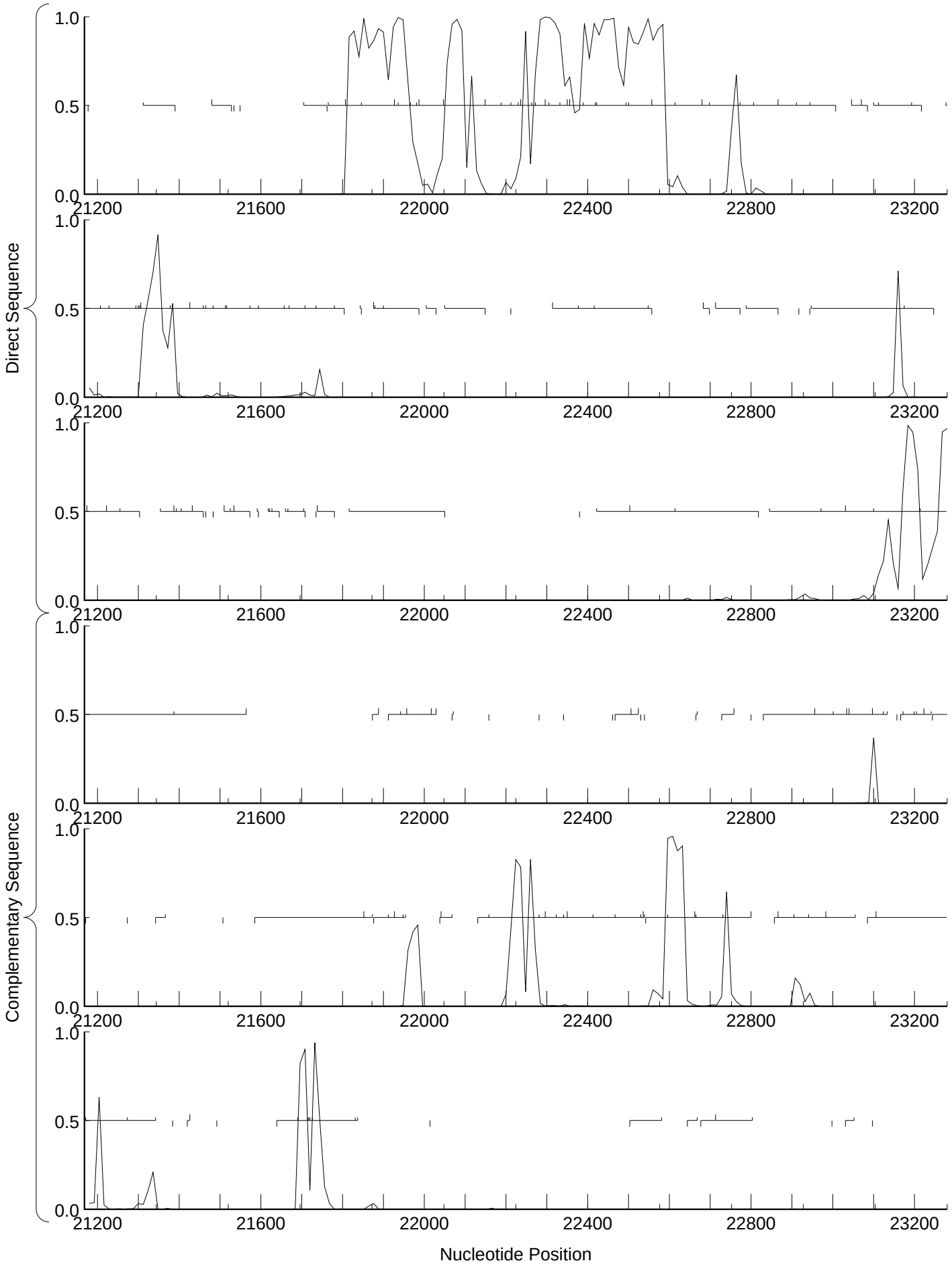
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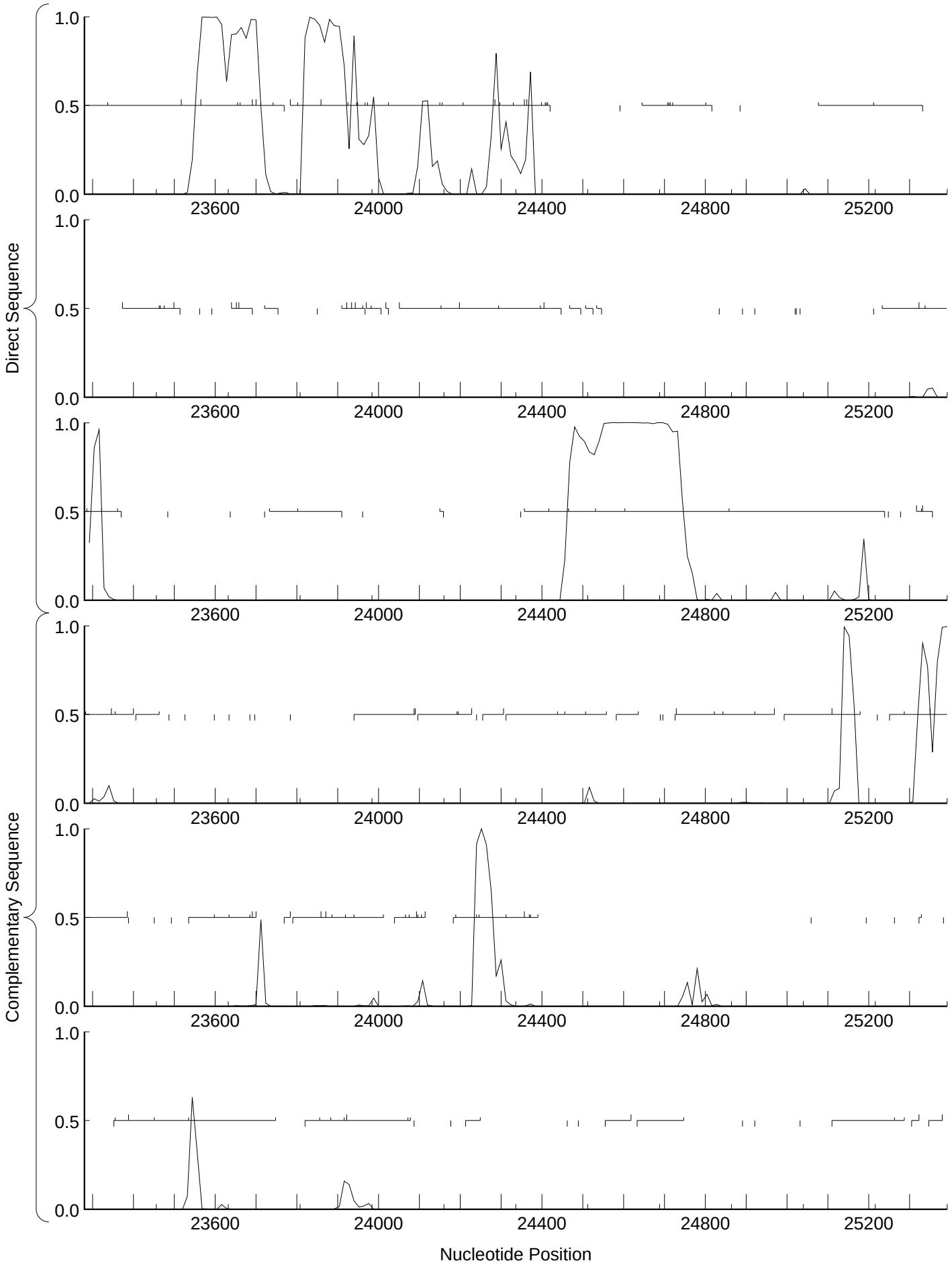
complete sequence, 59815 bp including 10-base 3’ overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 11/30



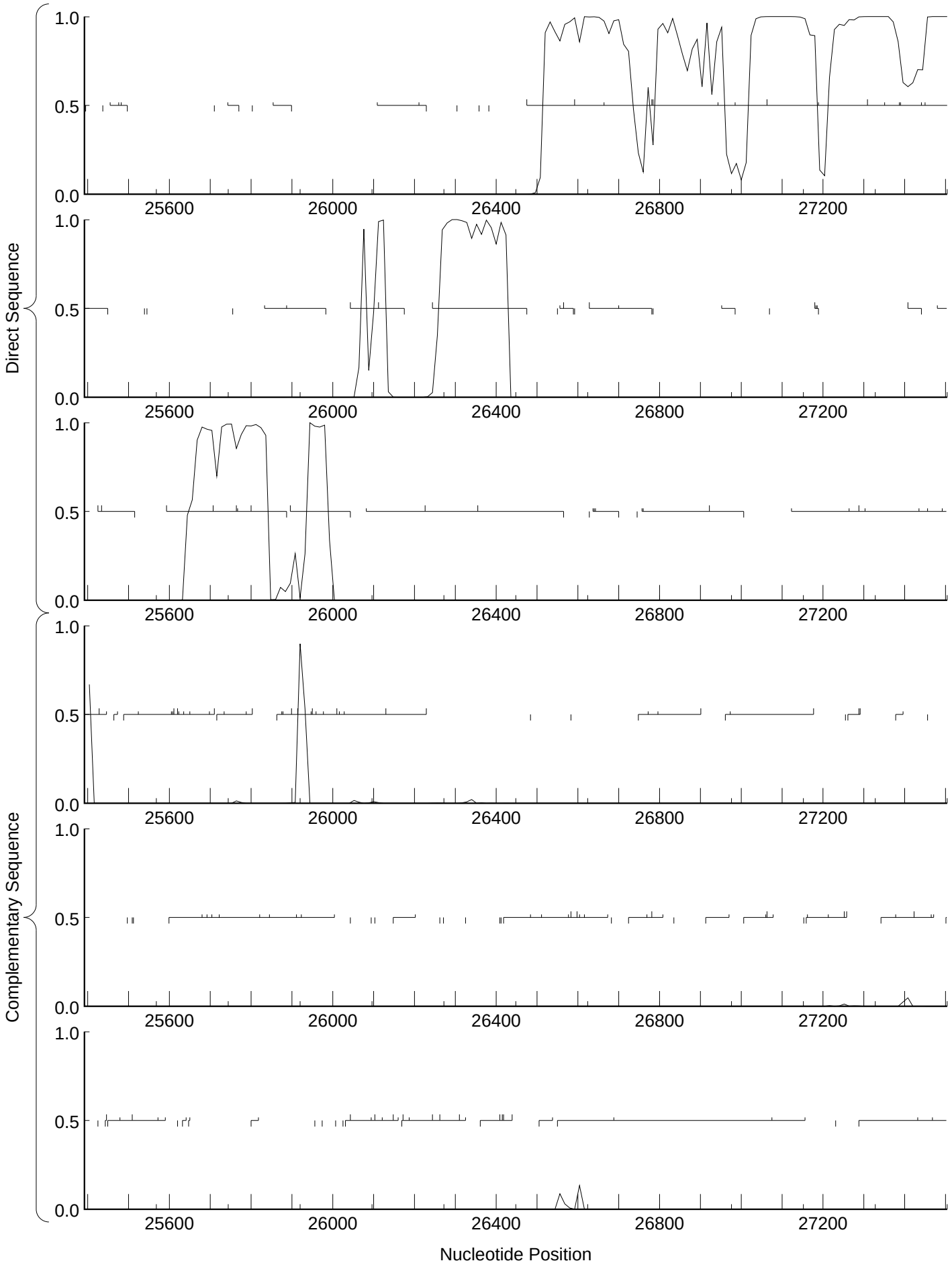
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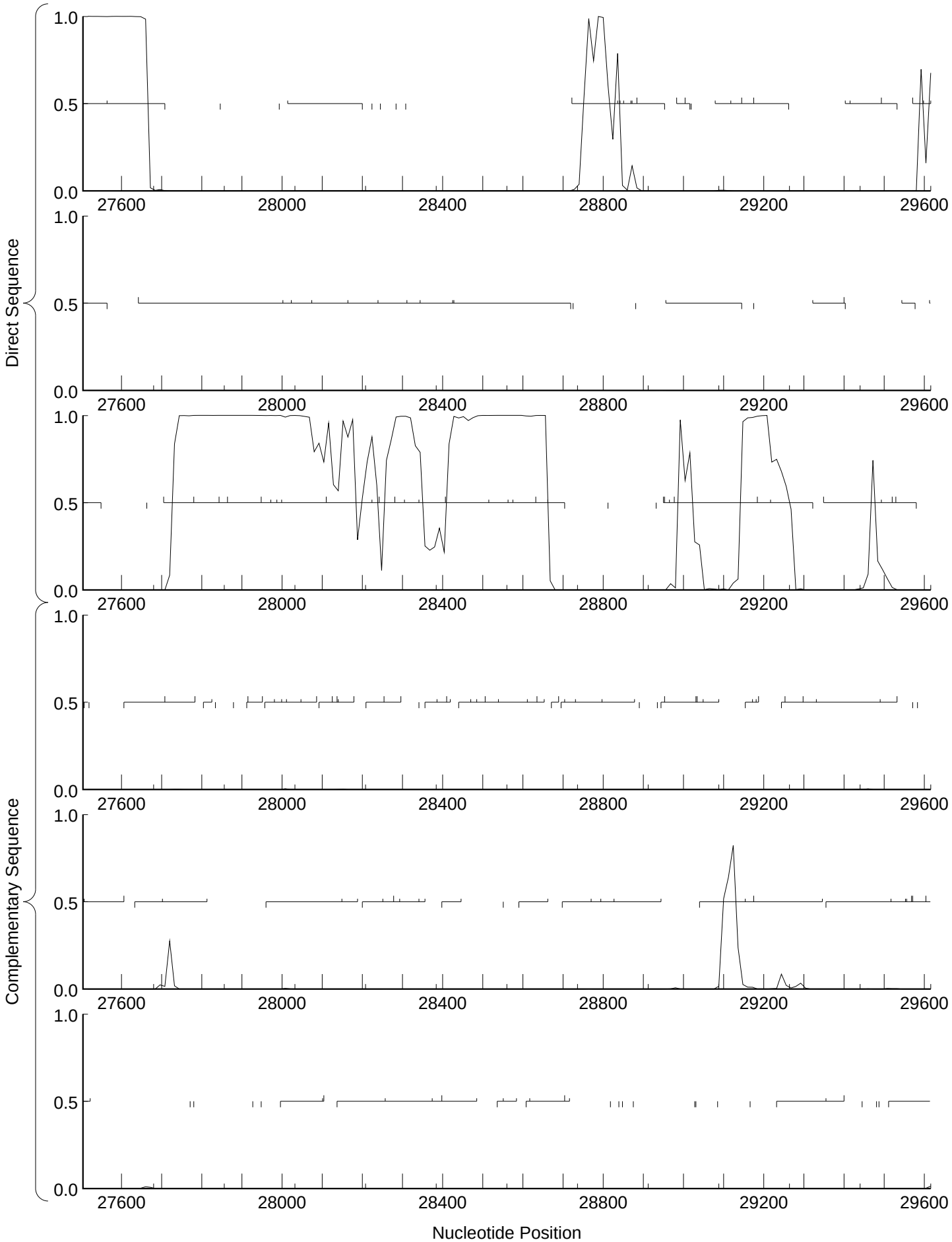
complete sequence, 59815 bp including 10-base 3’ overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 13/30



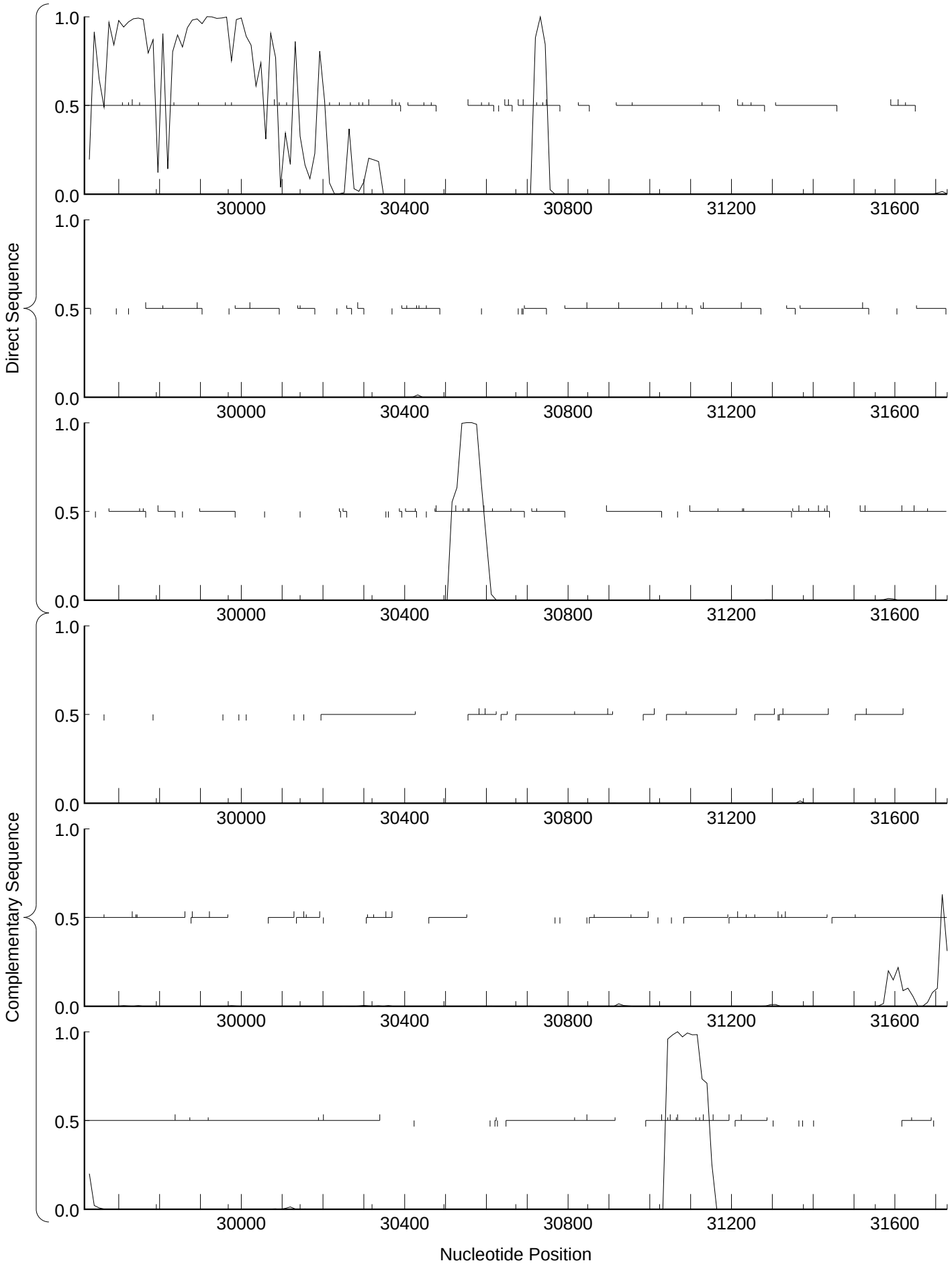
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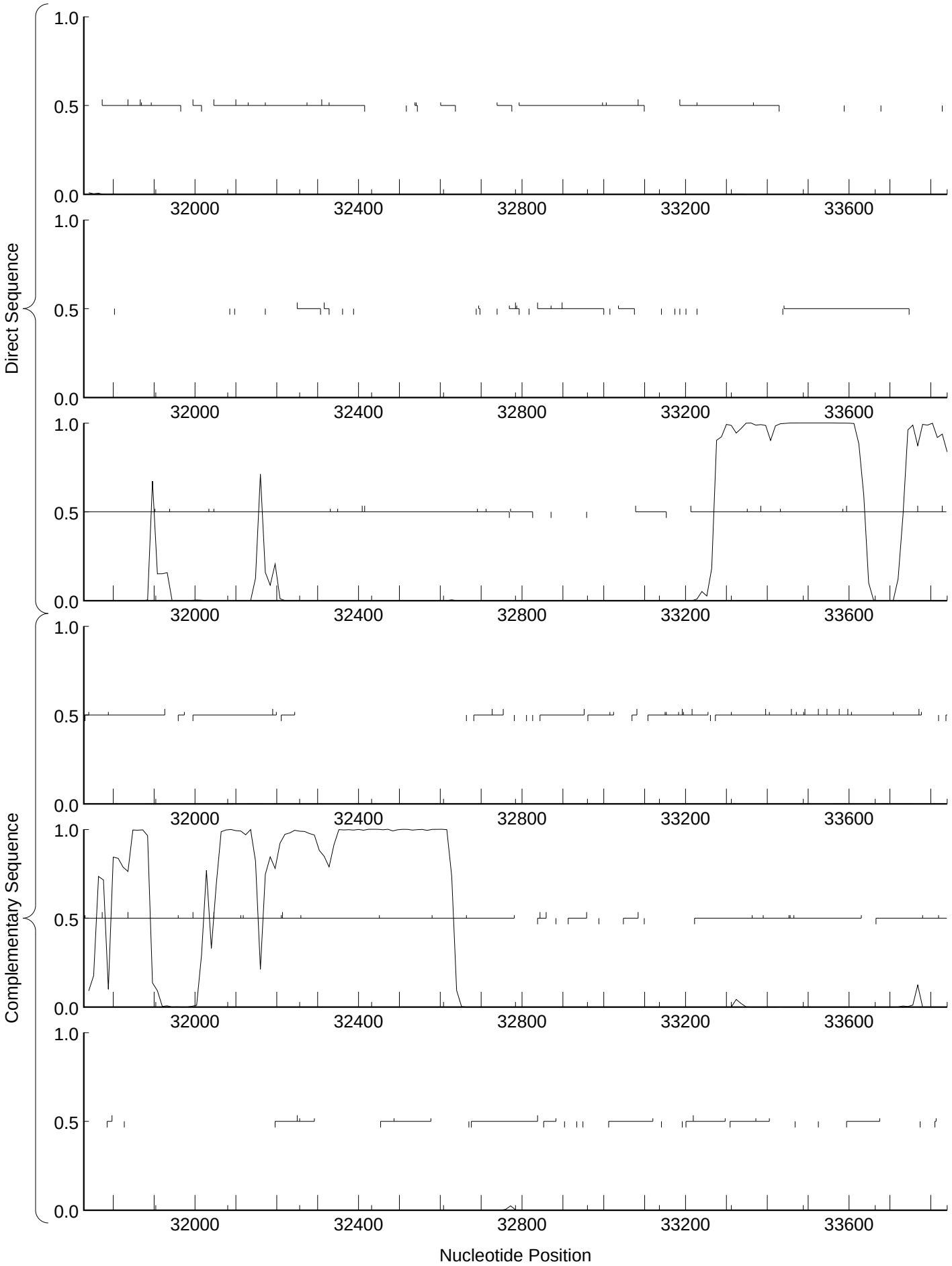
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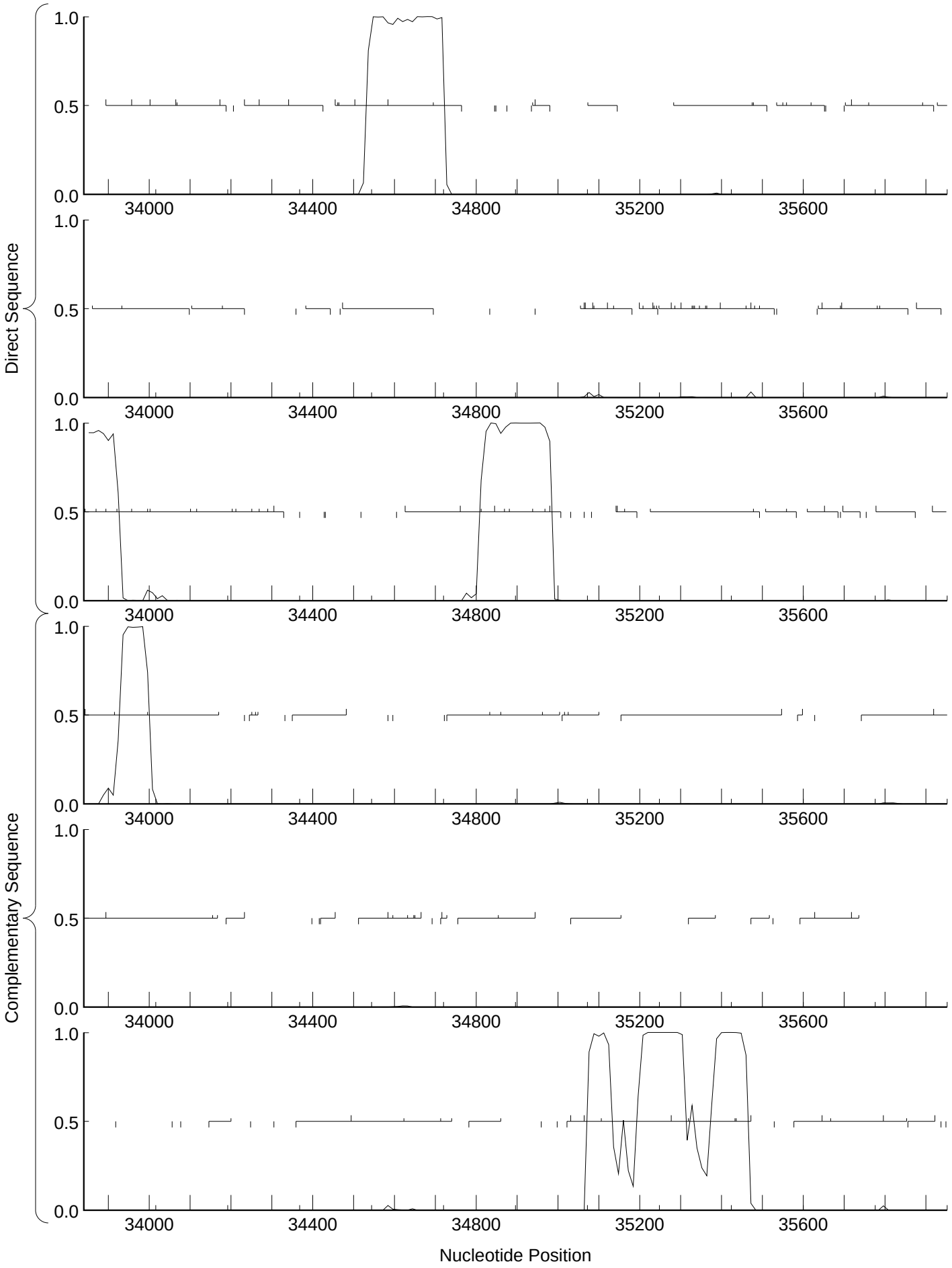
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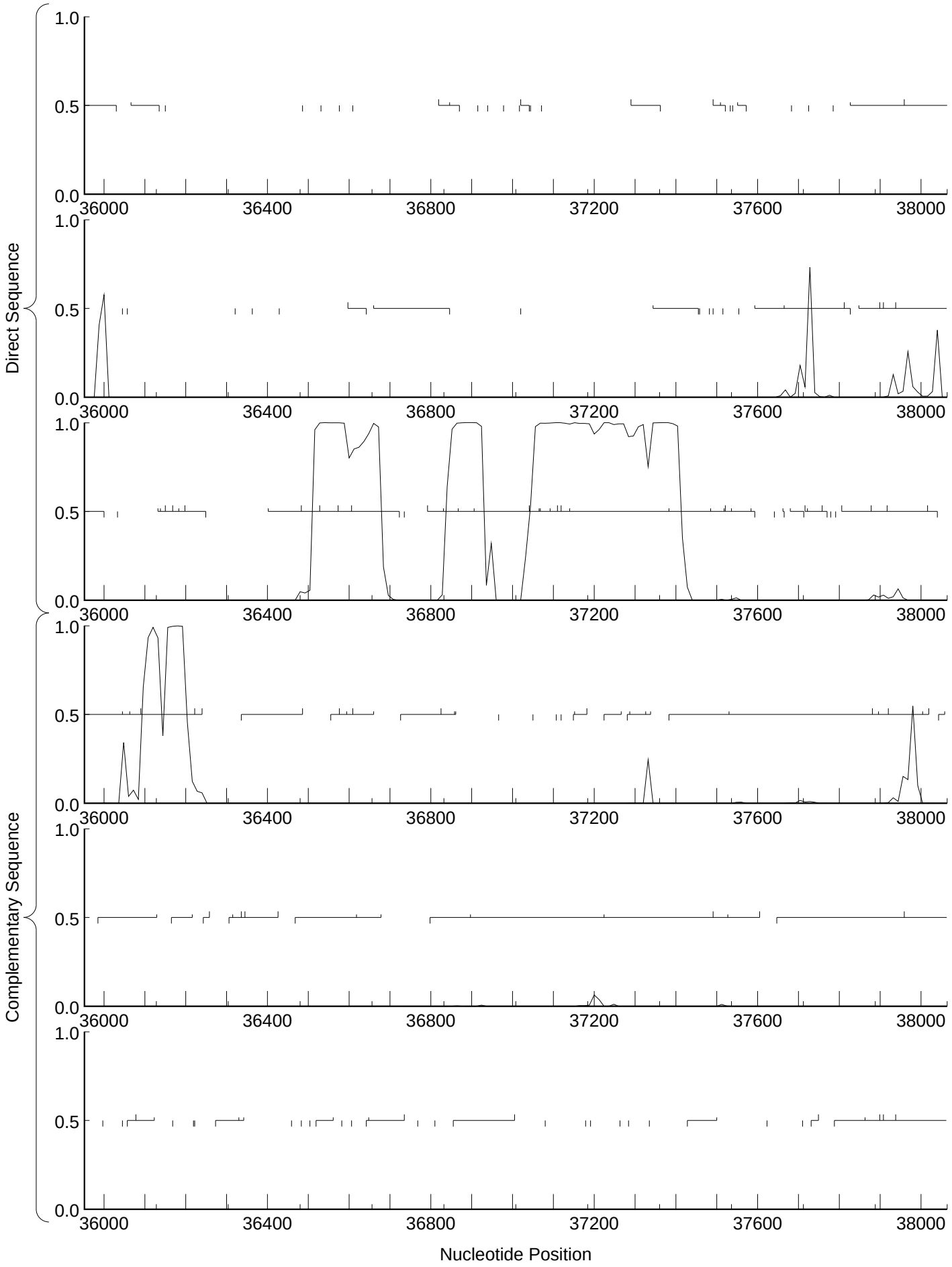
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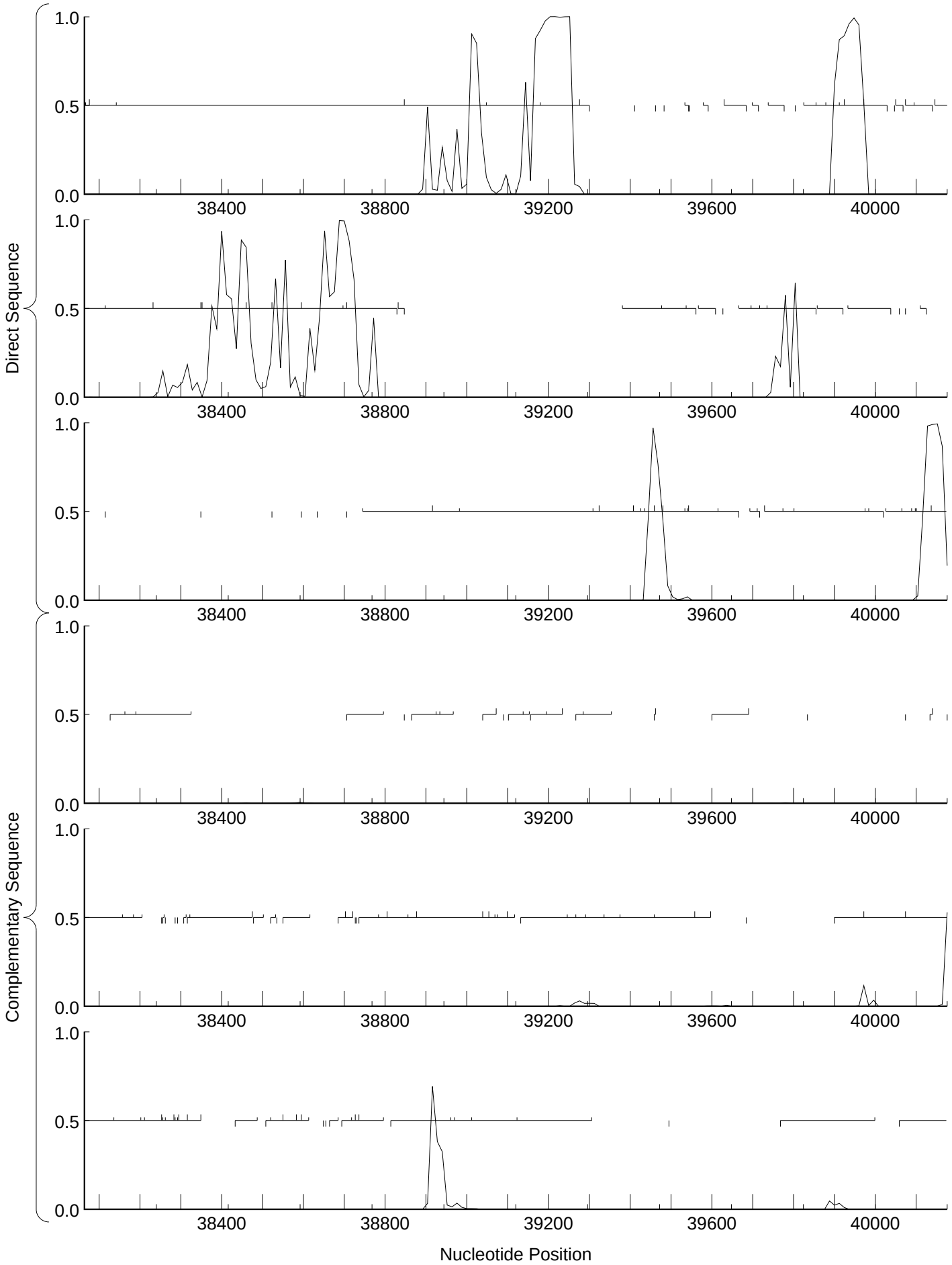
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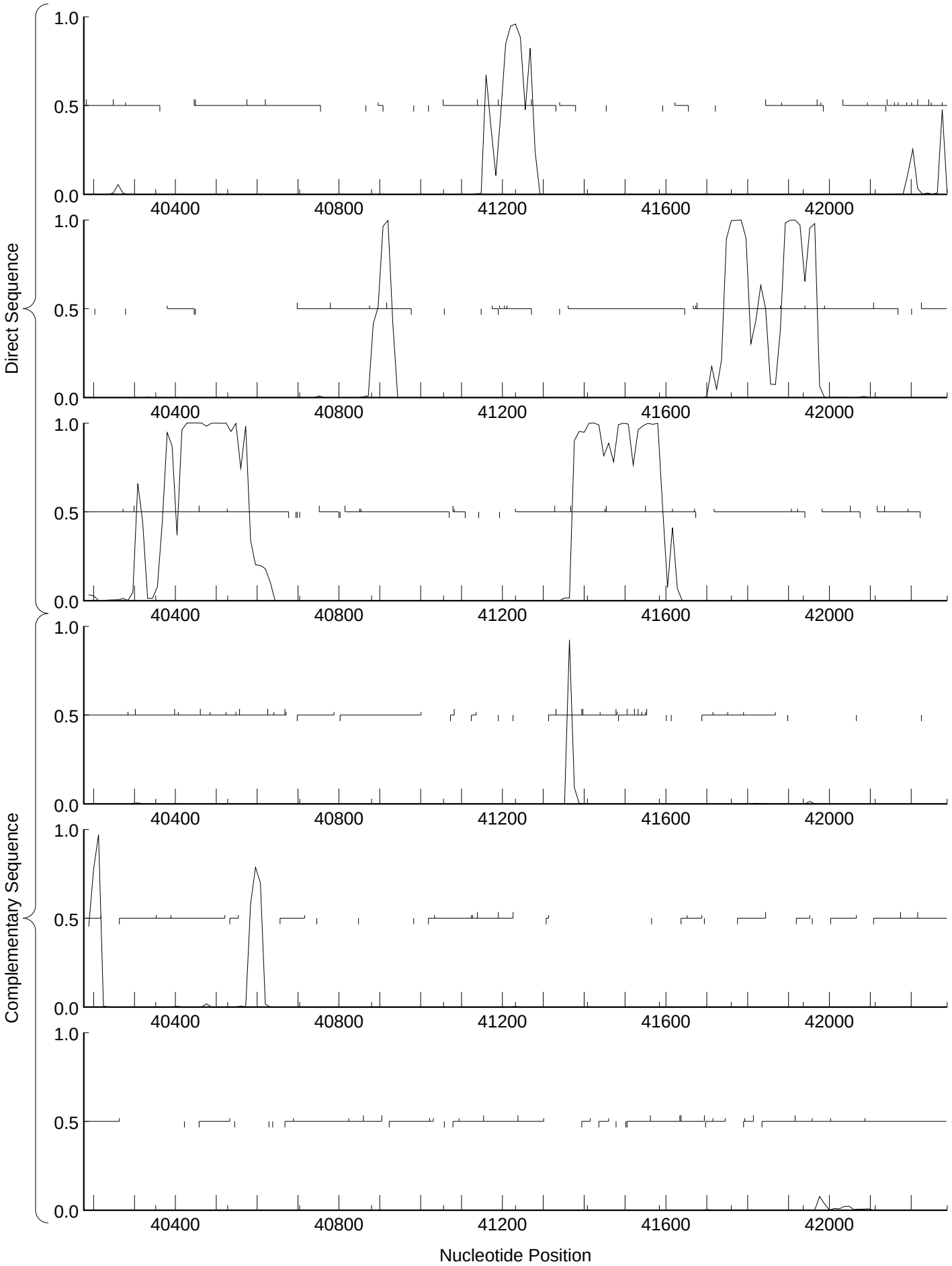
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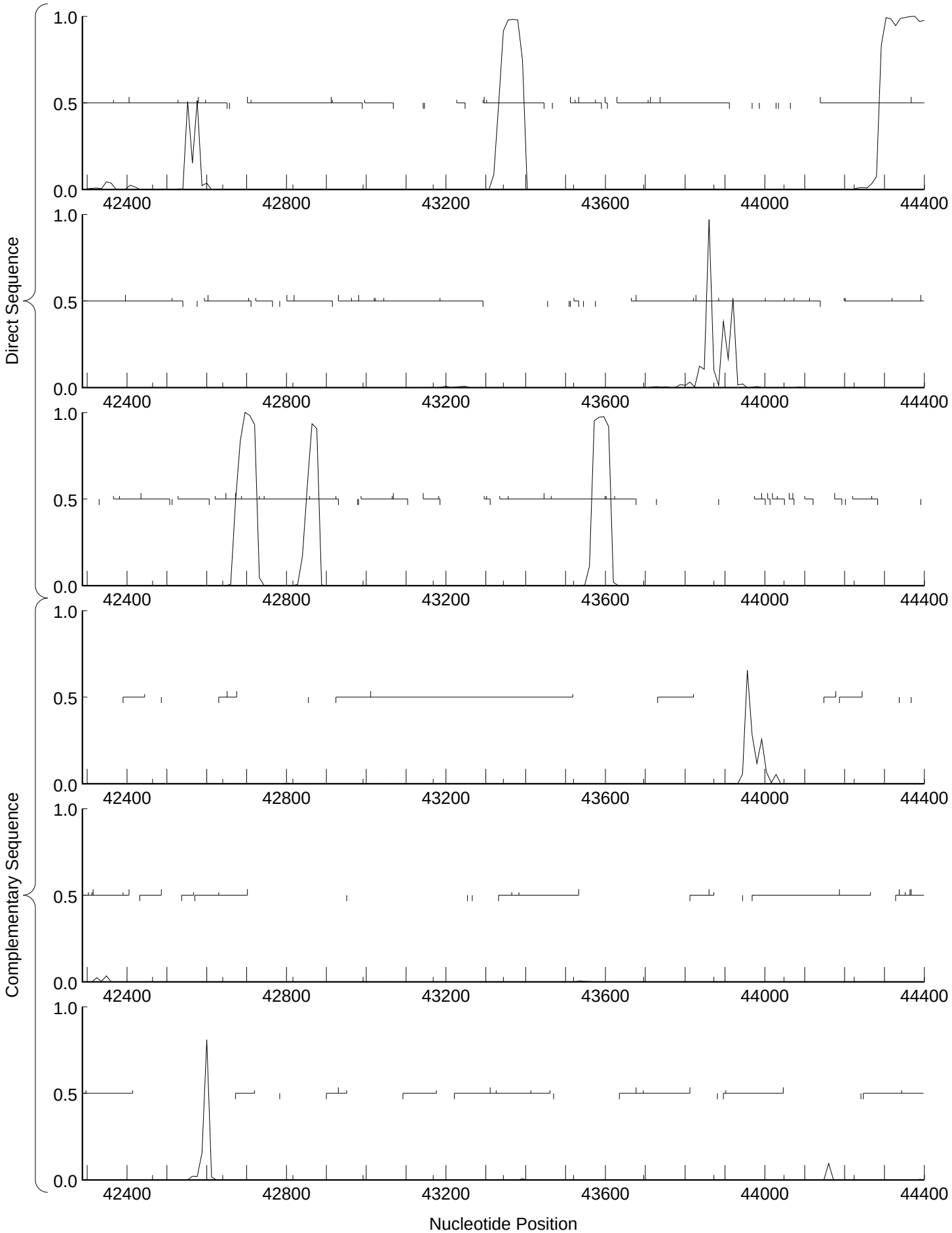
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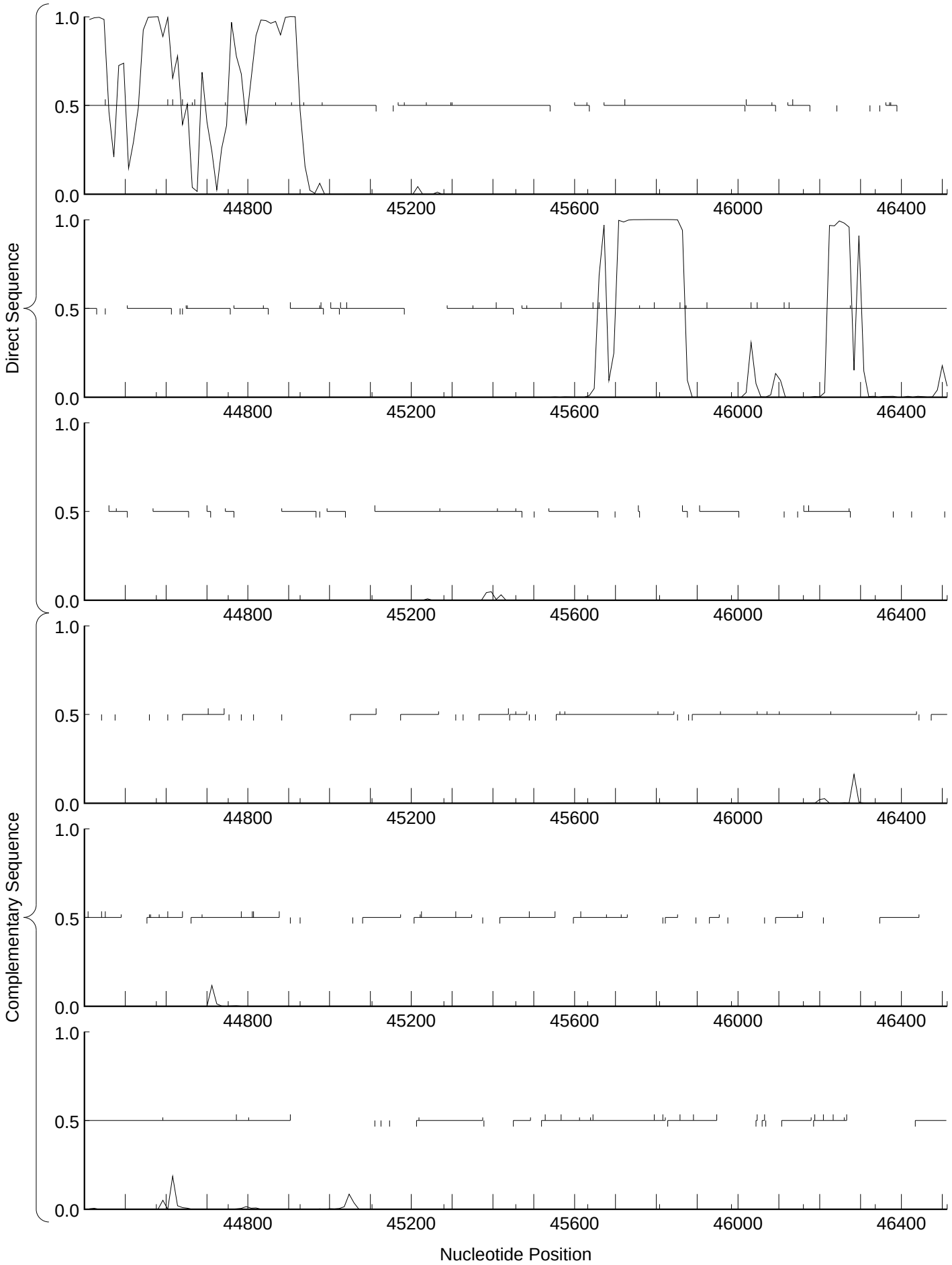
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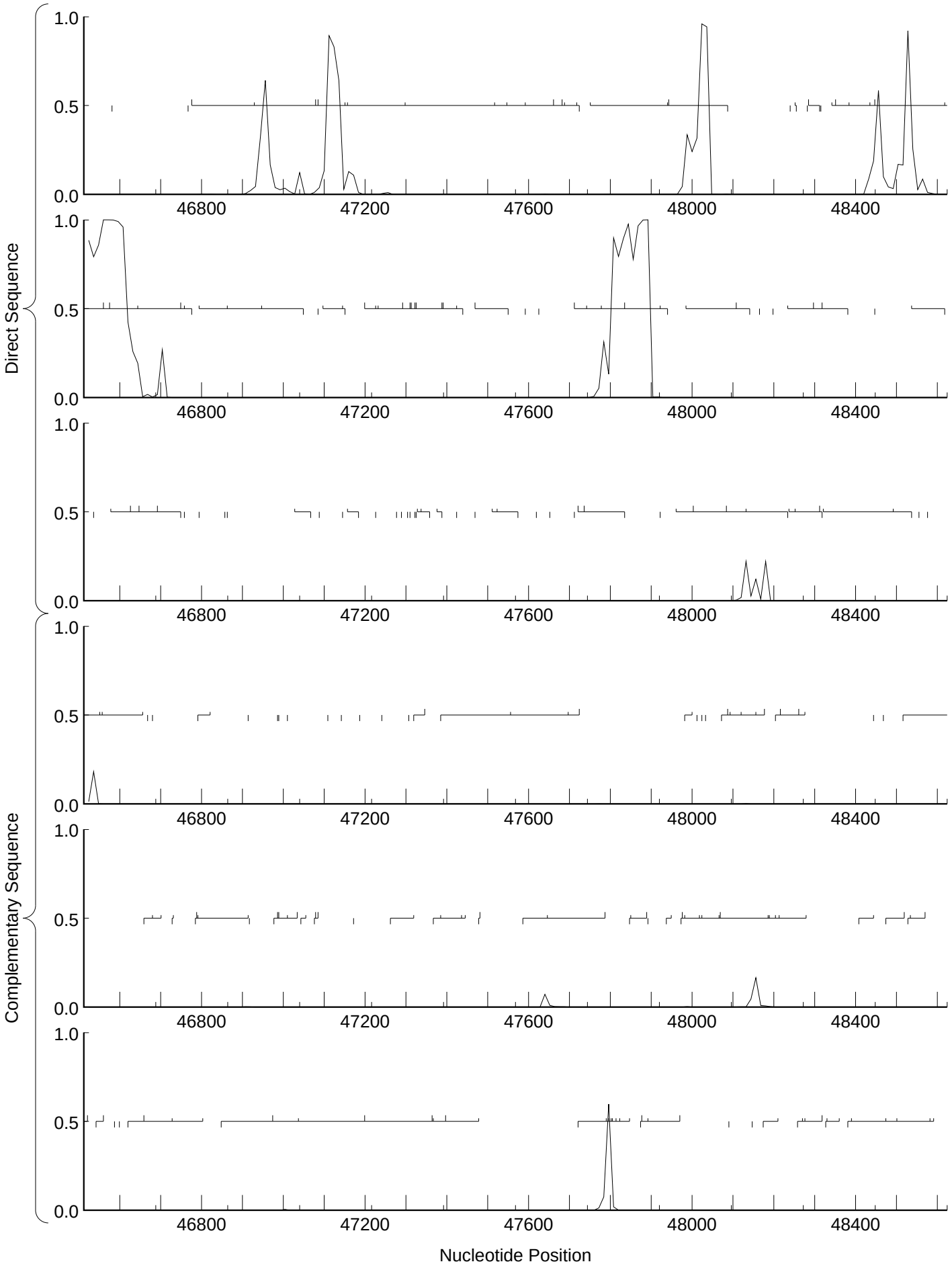
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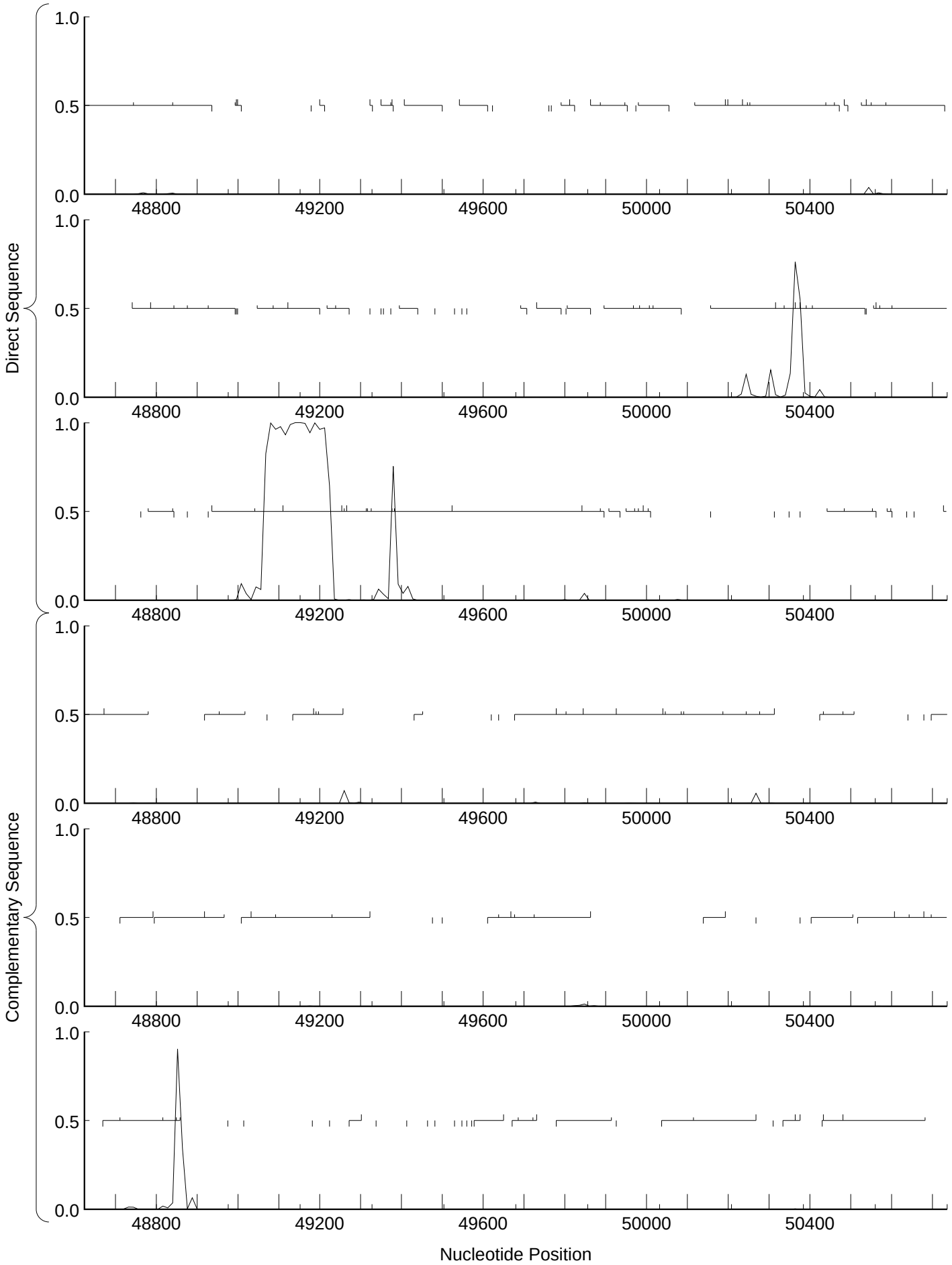
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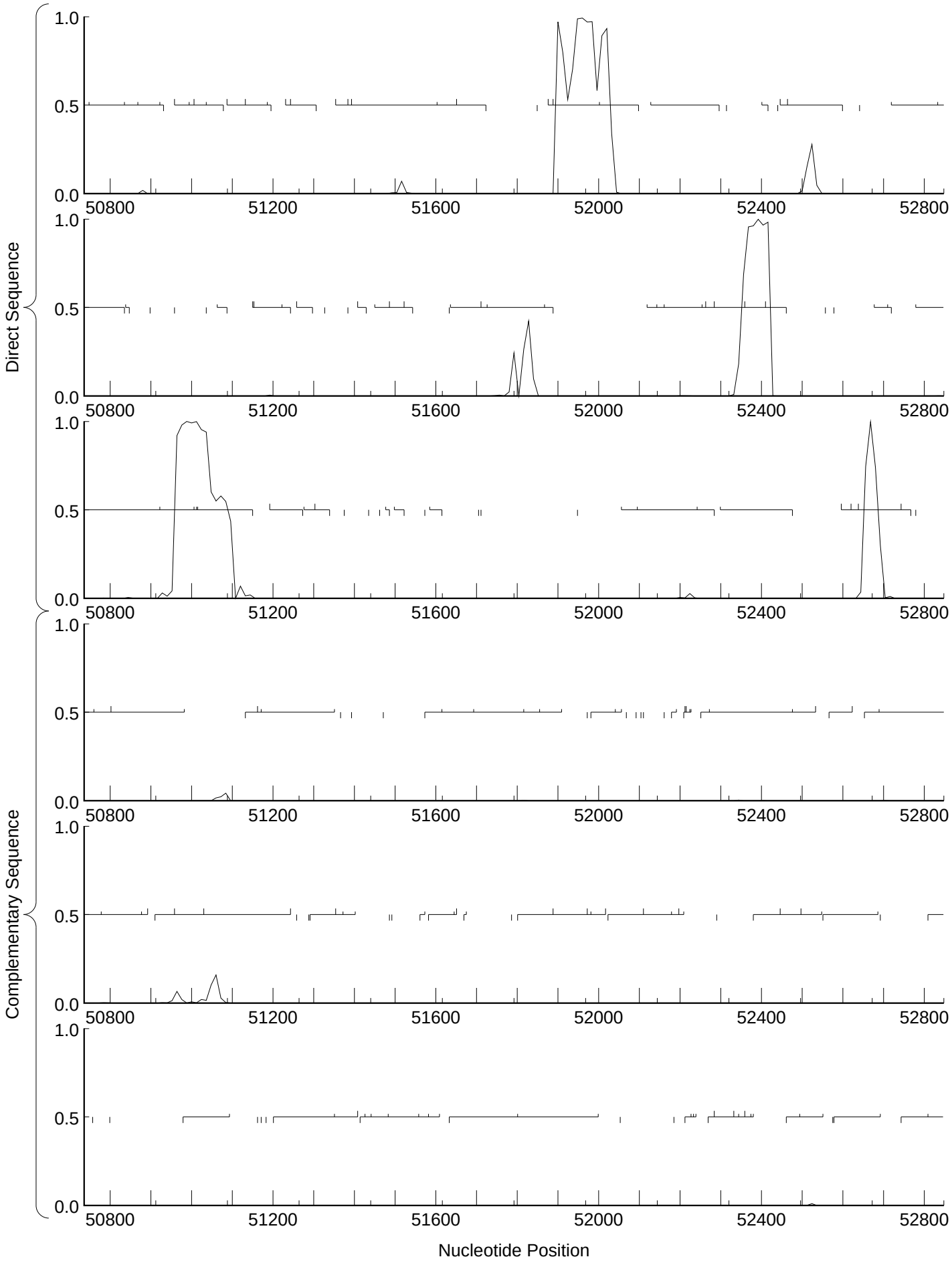
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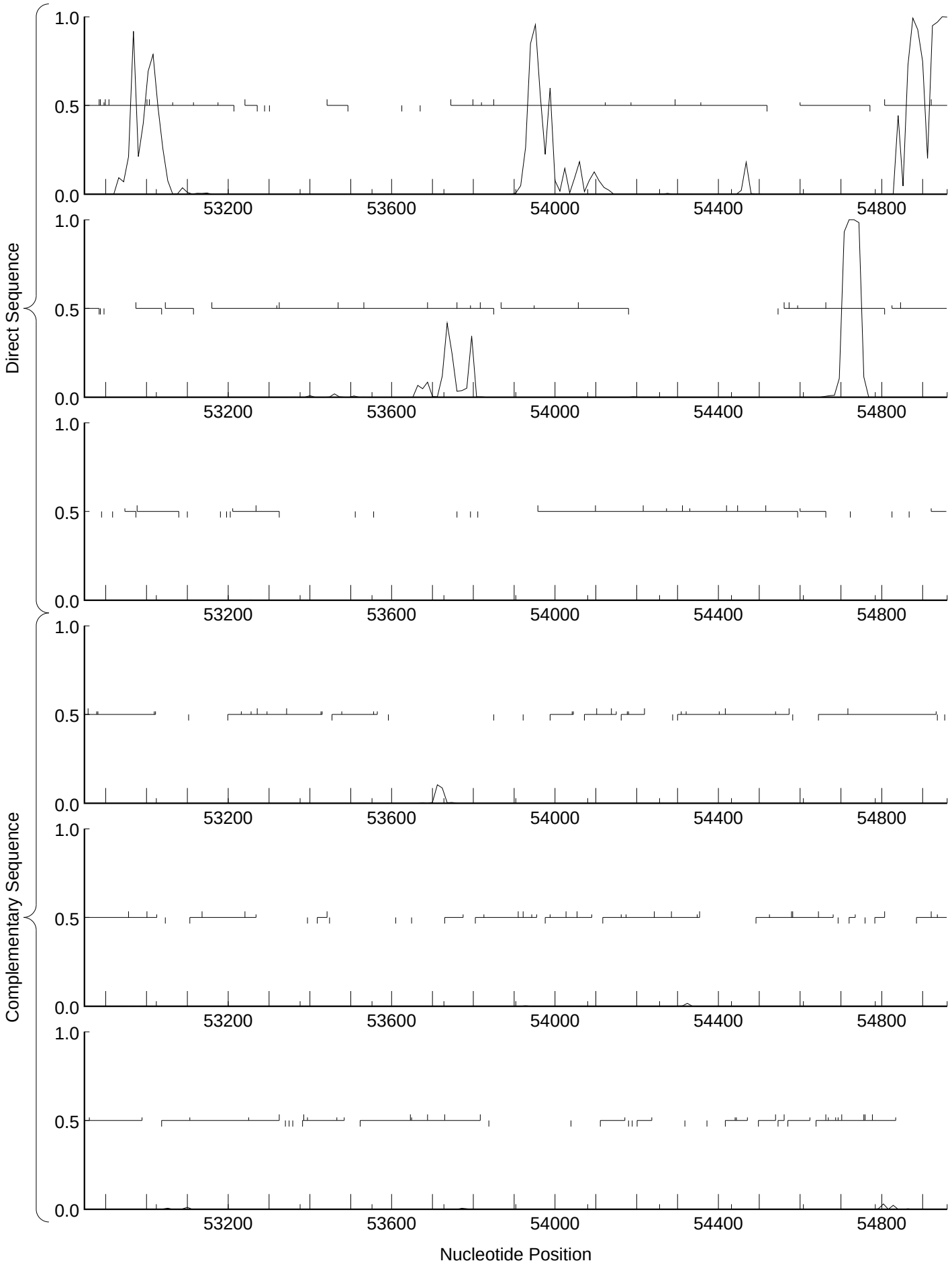
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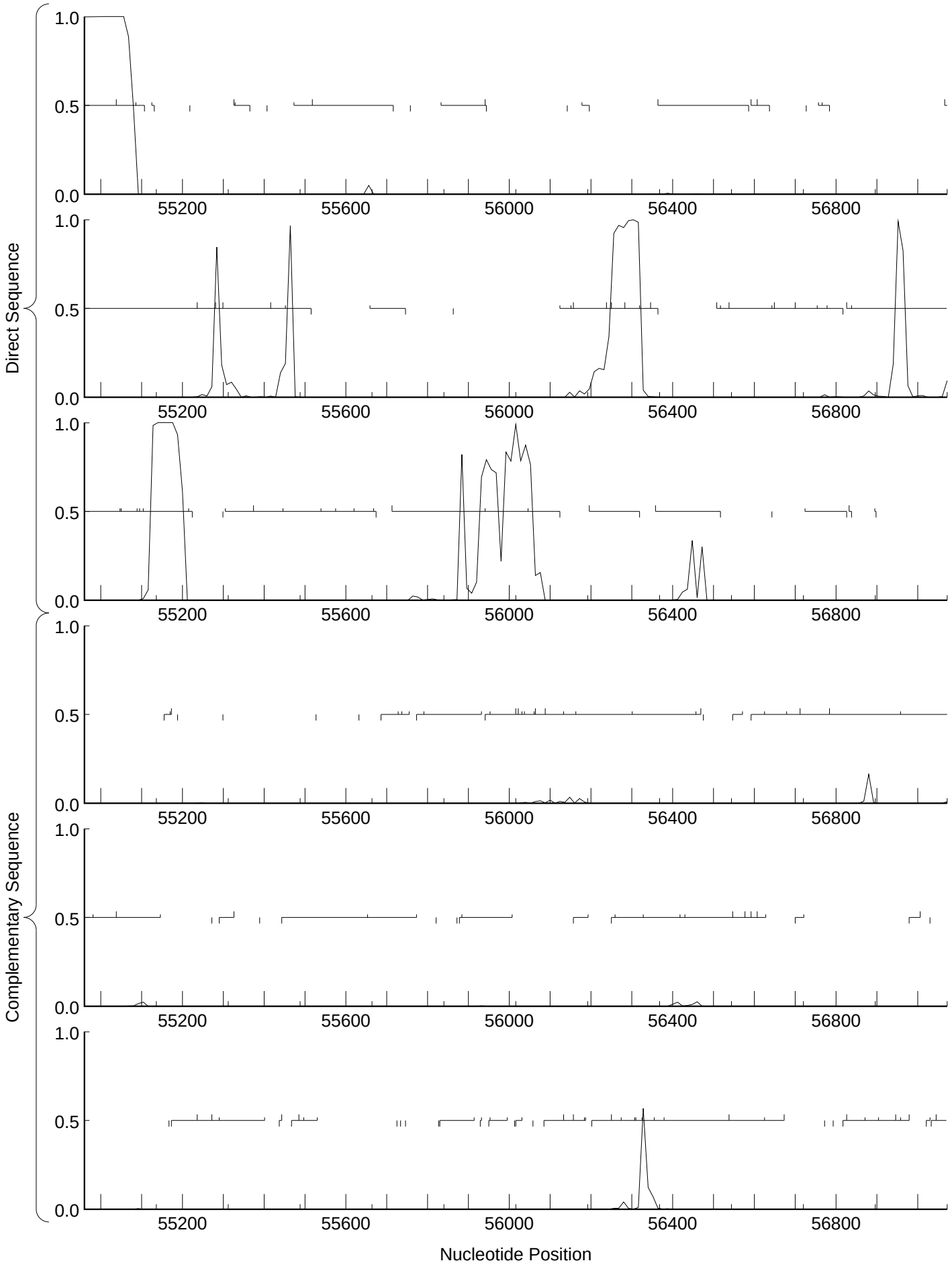
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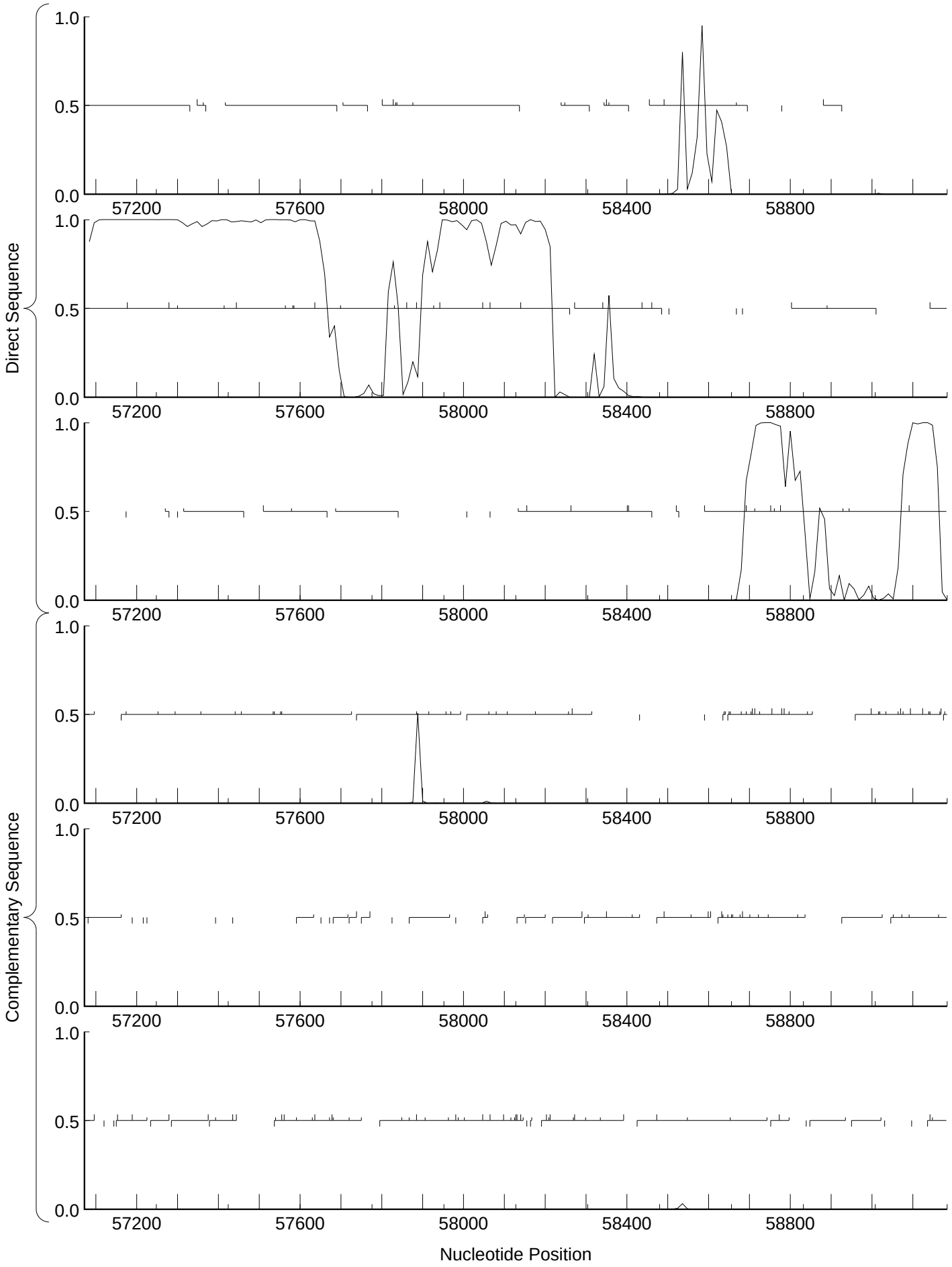
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complete sequence, 59815 bp including 10-base 3’ overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 28/30



complete sequence, 59815 bp including 10-base 3’ overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 29/30



complete sequence, 59815 bp including 10-base 3’ overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 30/30

