GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Qobbit complete sequence, 52911 bp including 10-base 3' overhang (CGGTC
Analysis Date    :  3/28/24 at 6:10:46
Pages            :  27
Sequence Length  :  52911 bp
GC Content       :   62.57%

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

PS-Version       :  1.2

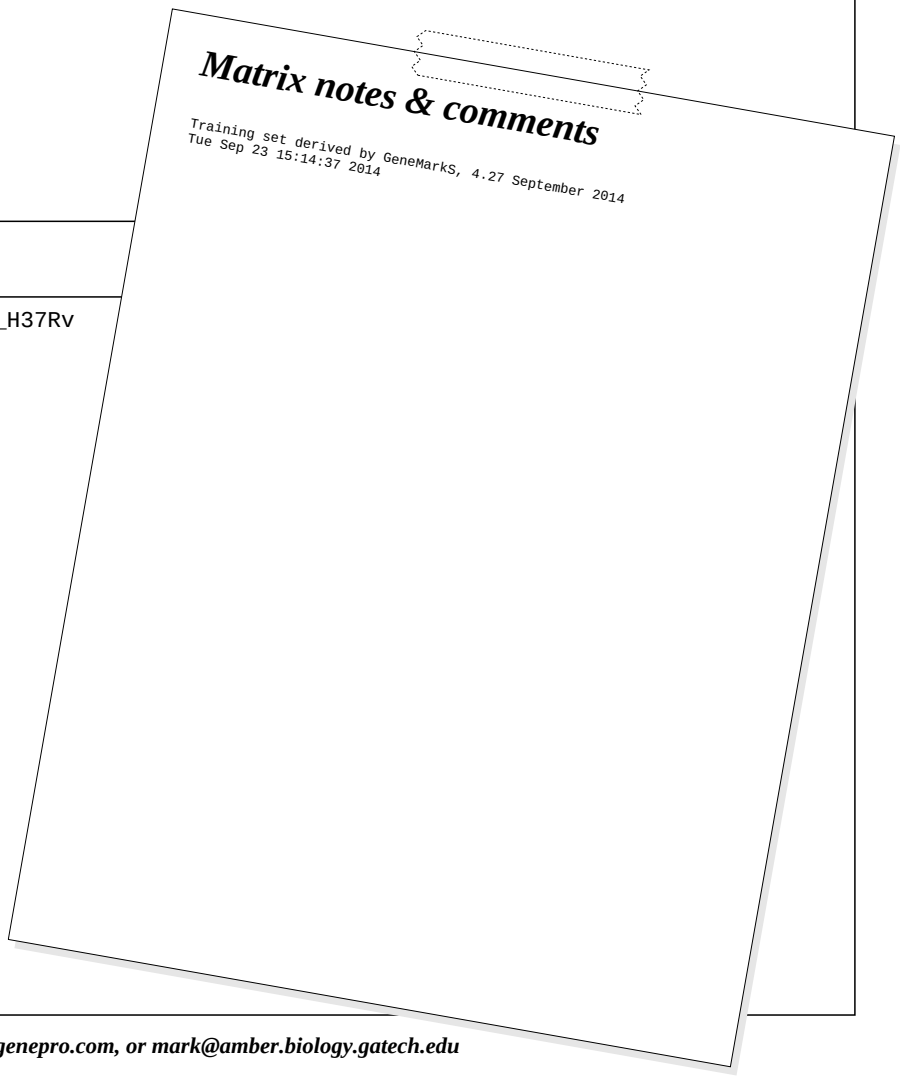
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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                    List regions and/or splice sites,
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MATRIX INFORMATION

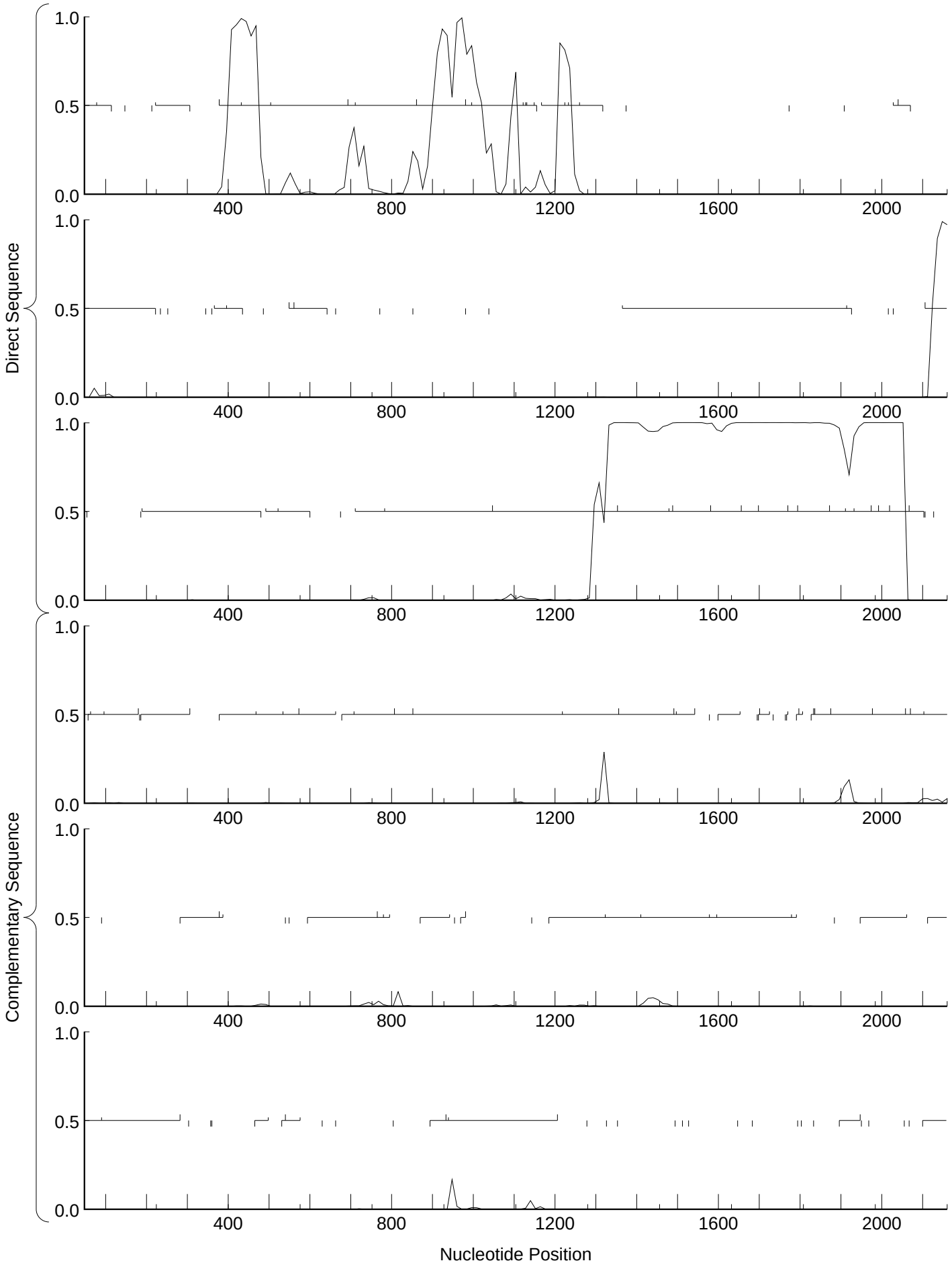
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Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

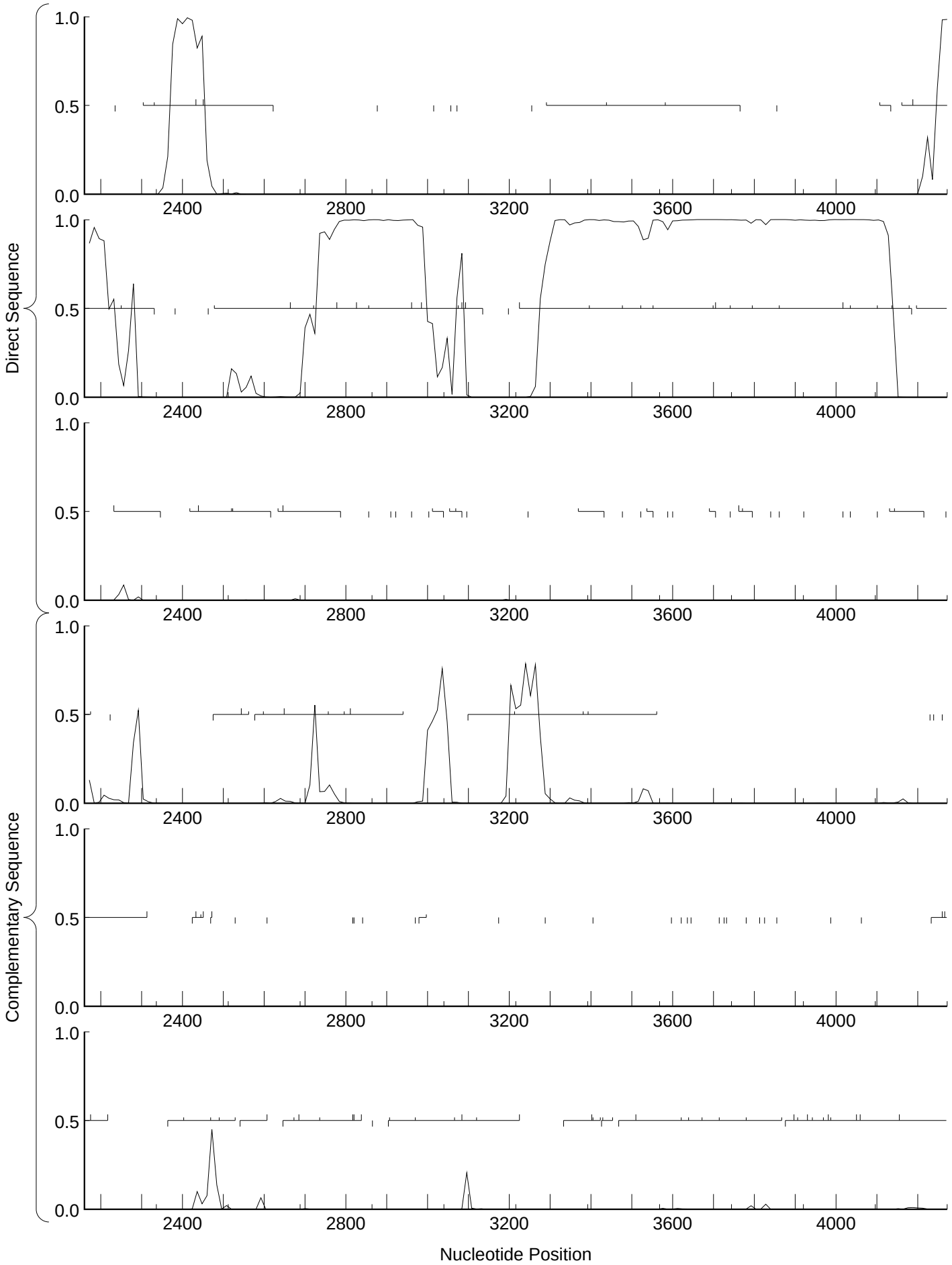
Order      :  4

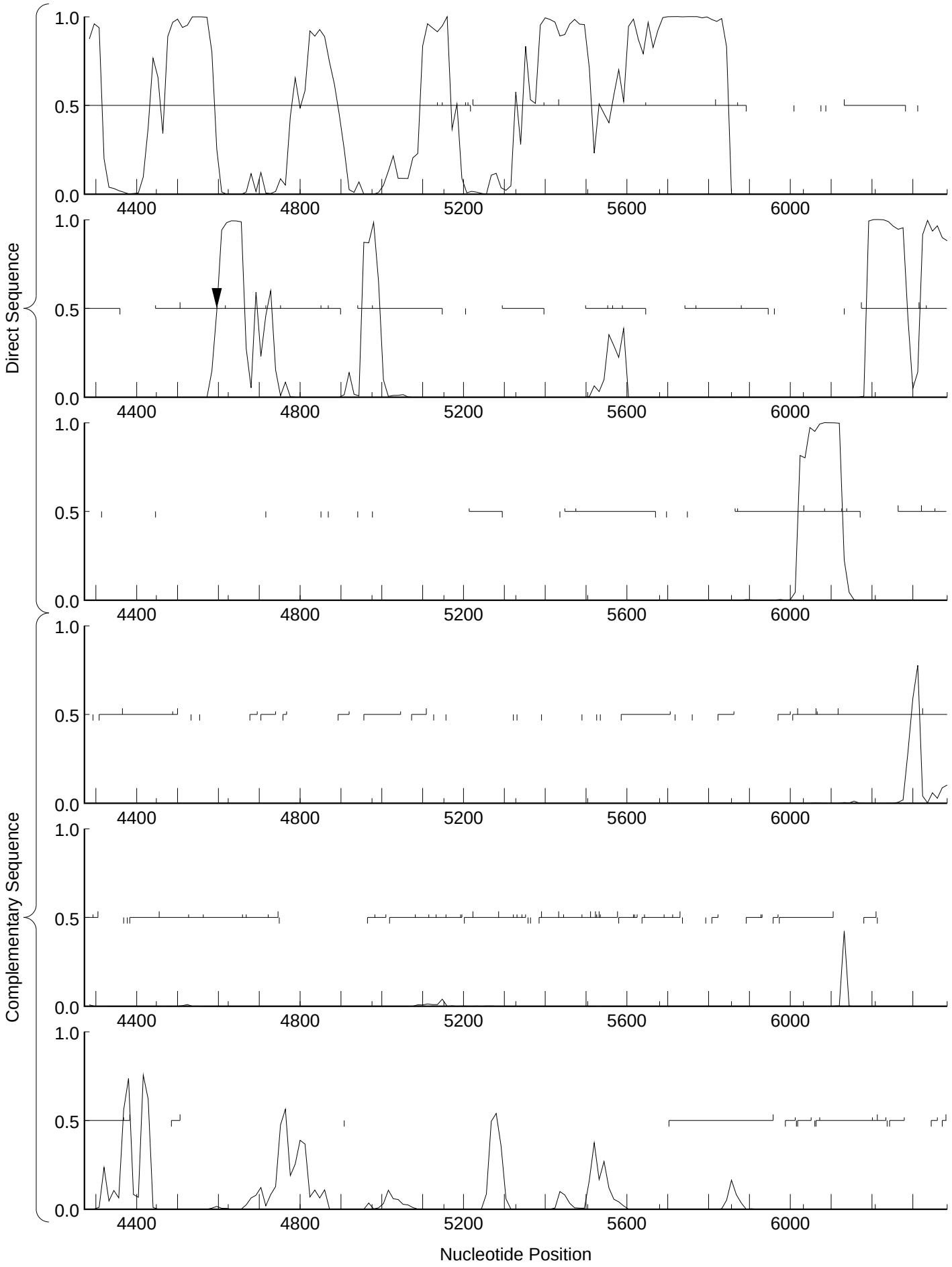

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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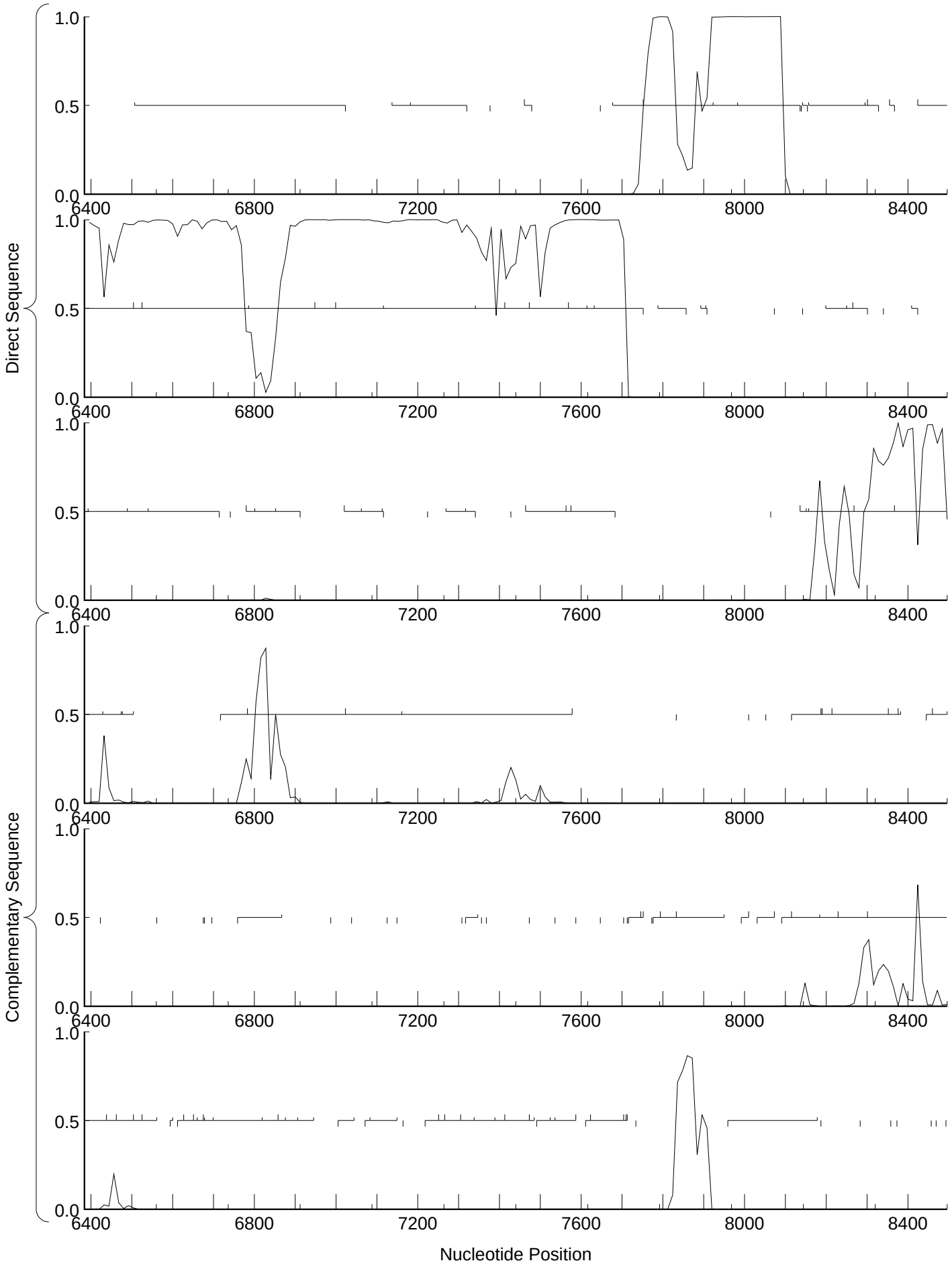


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

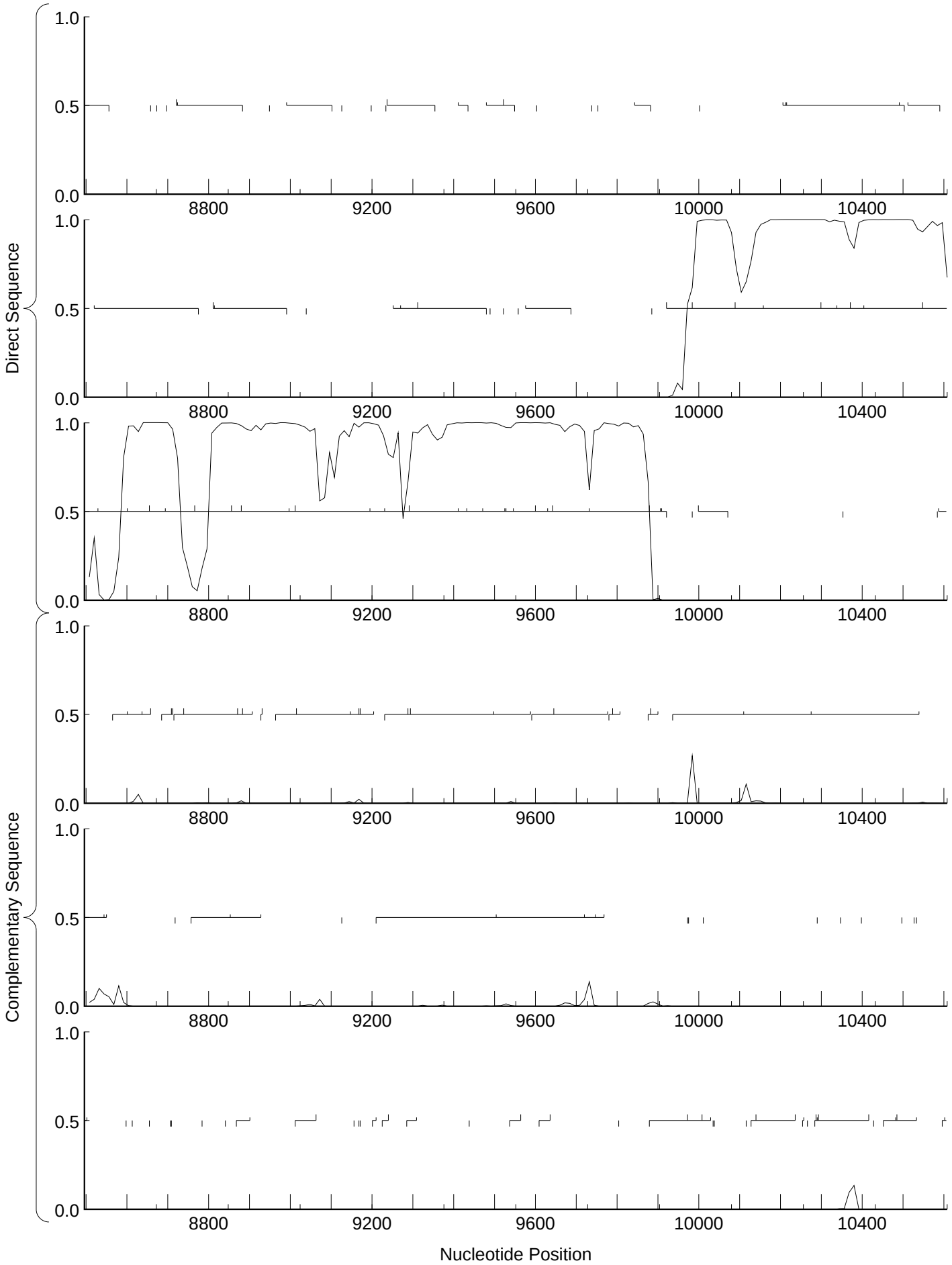

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0.0
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800
1200
1600
2000
Nucleotide Position

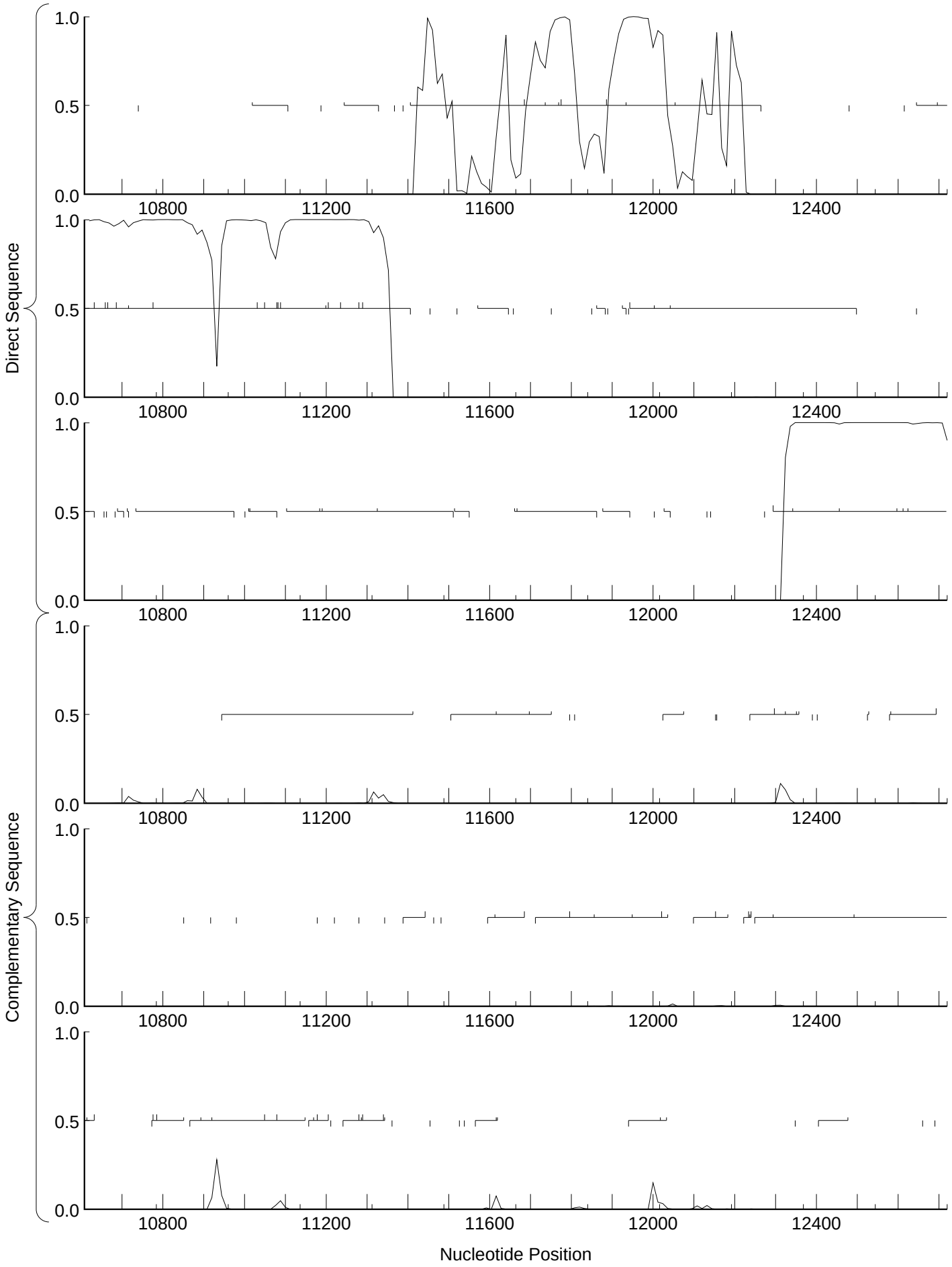

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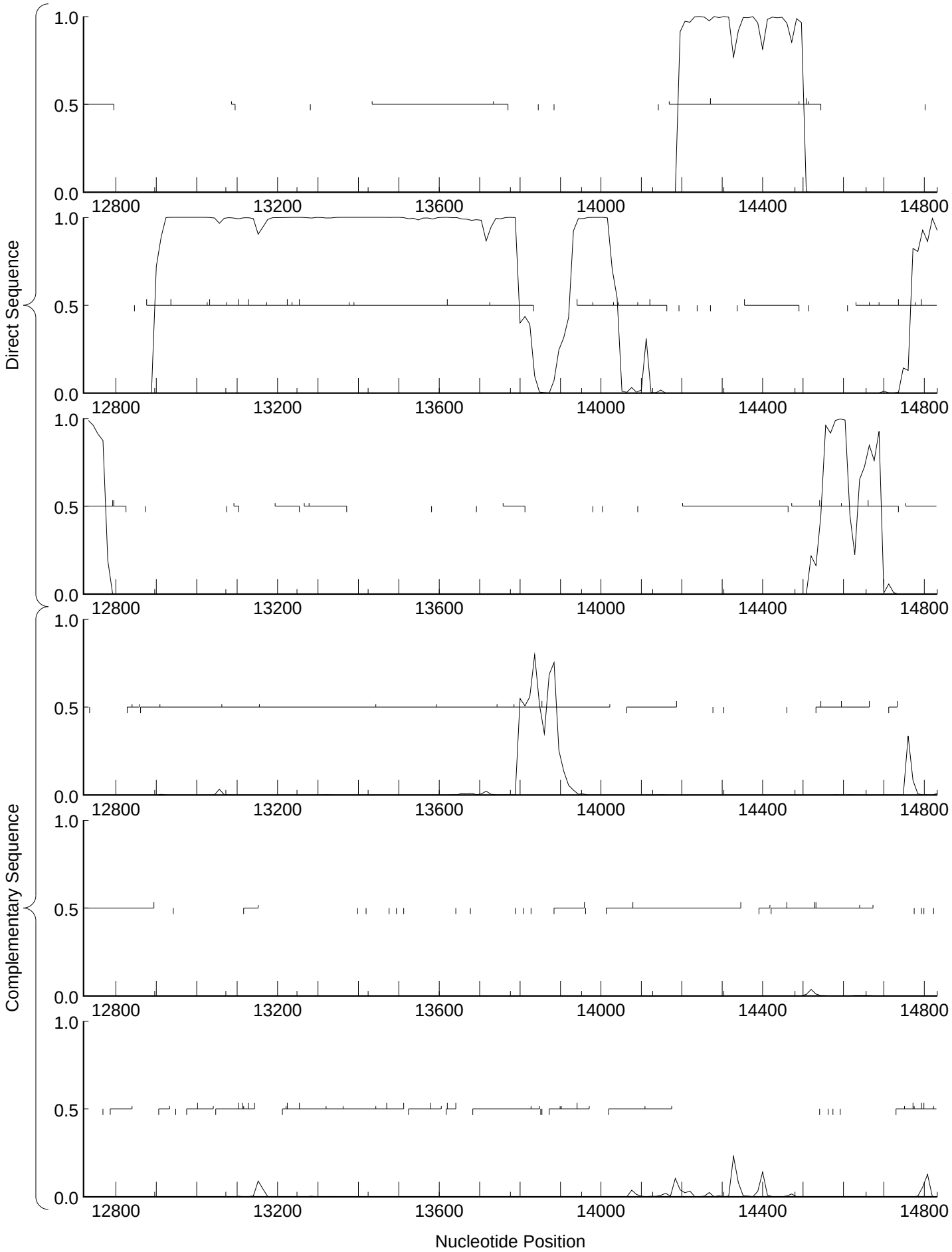

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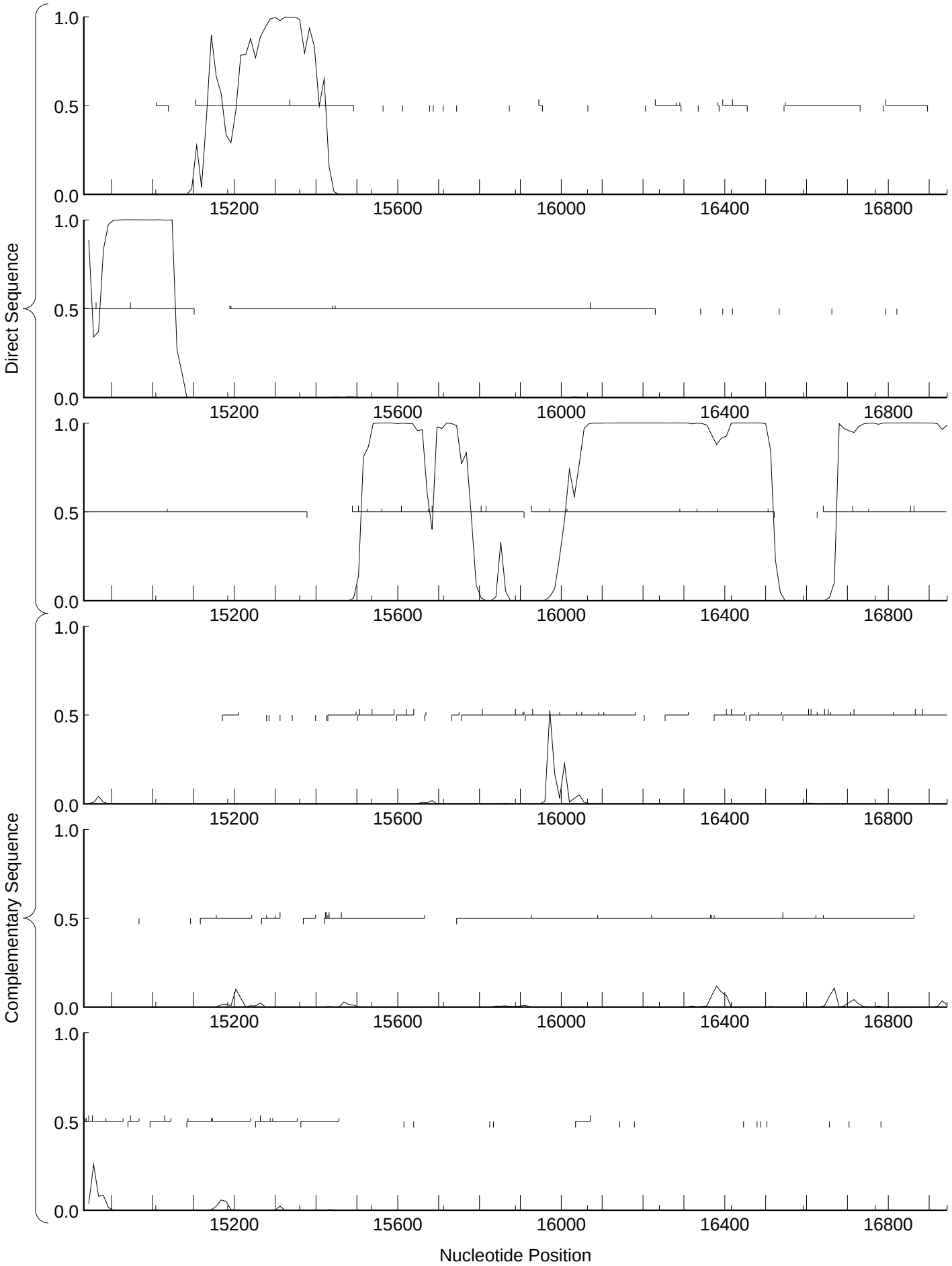

Direct Sequence
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6800
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7600
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8400
Nucleotide Position

bit complete sequence, 52911 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 6/27

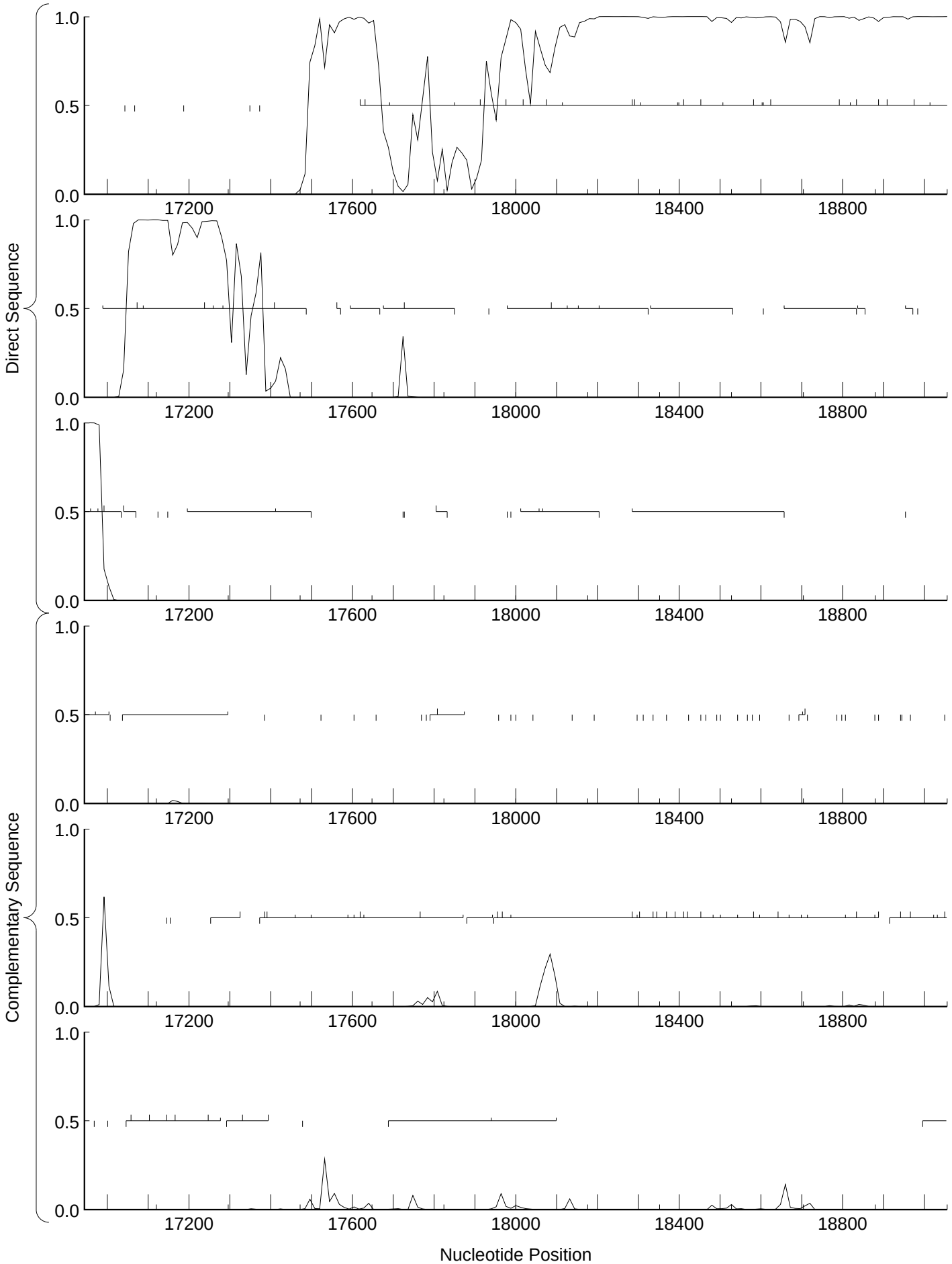



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11200
11600
12000
12400
Nucleotide Position

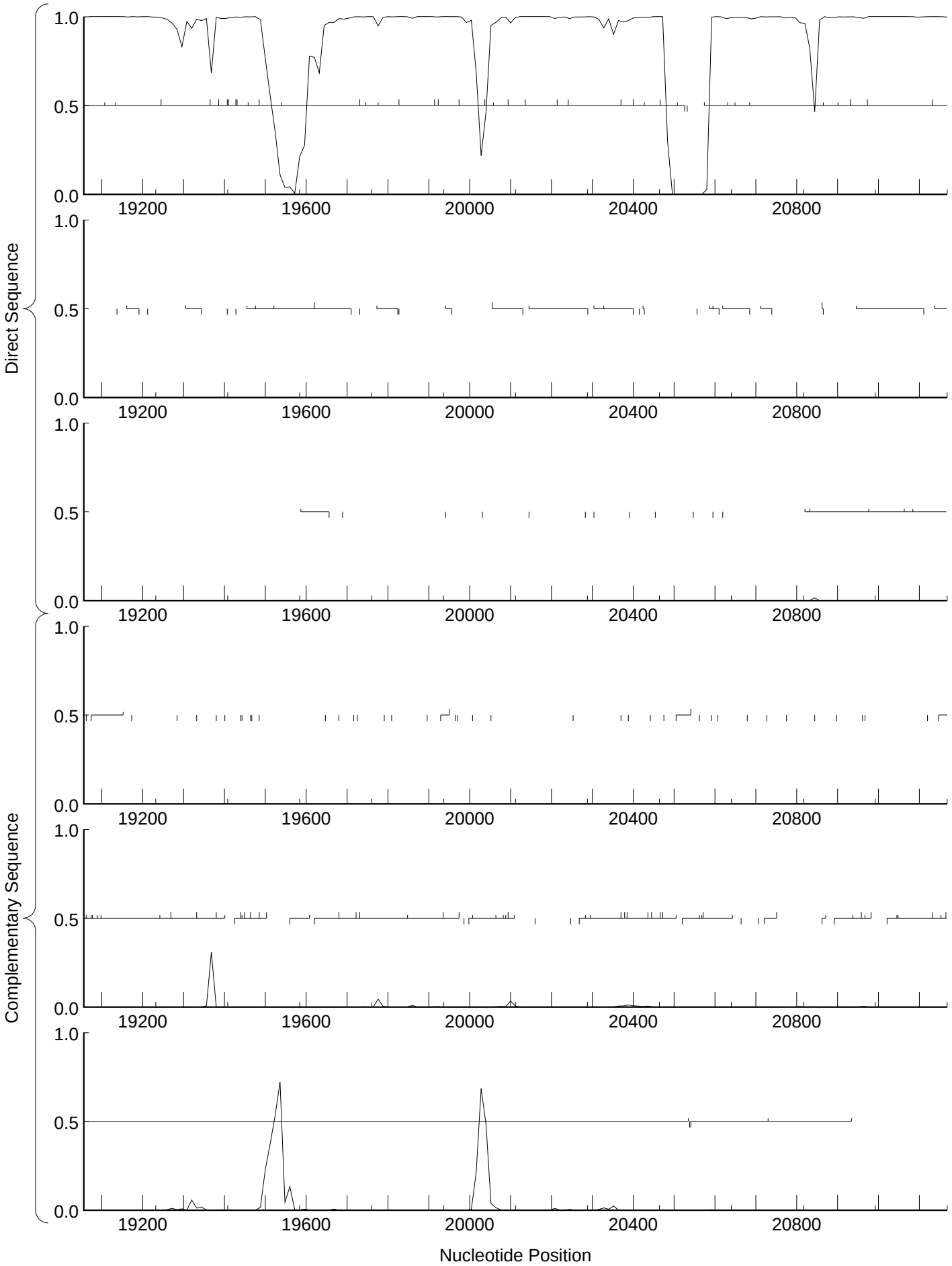

Direct Sequence
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13200
13600
14000
14400
14800
Nucleotide Position


Direct Sequence
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Nucleotide Position

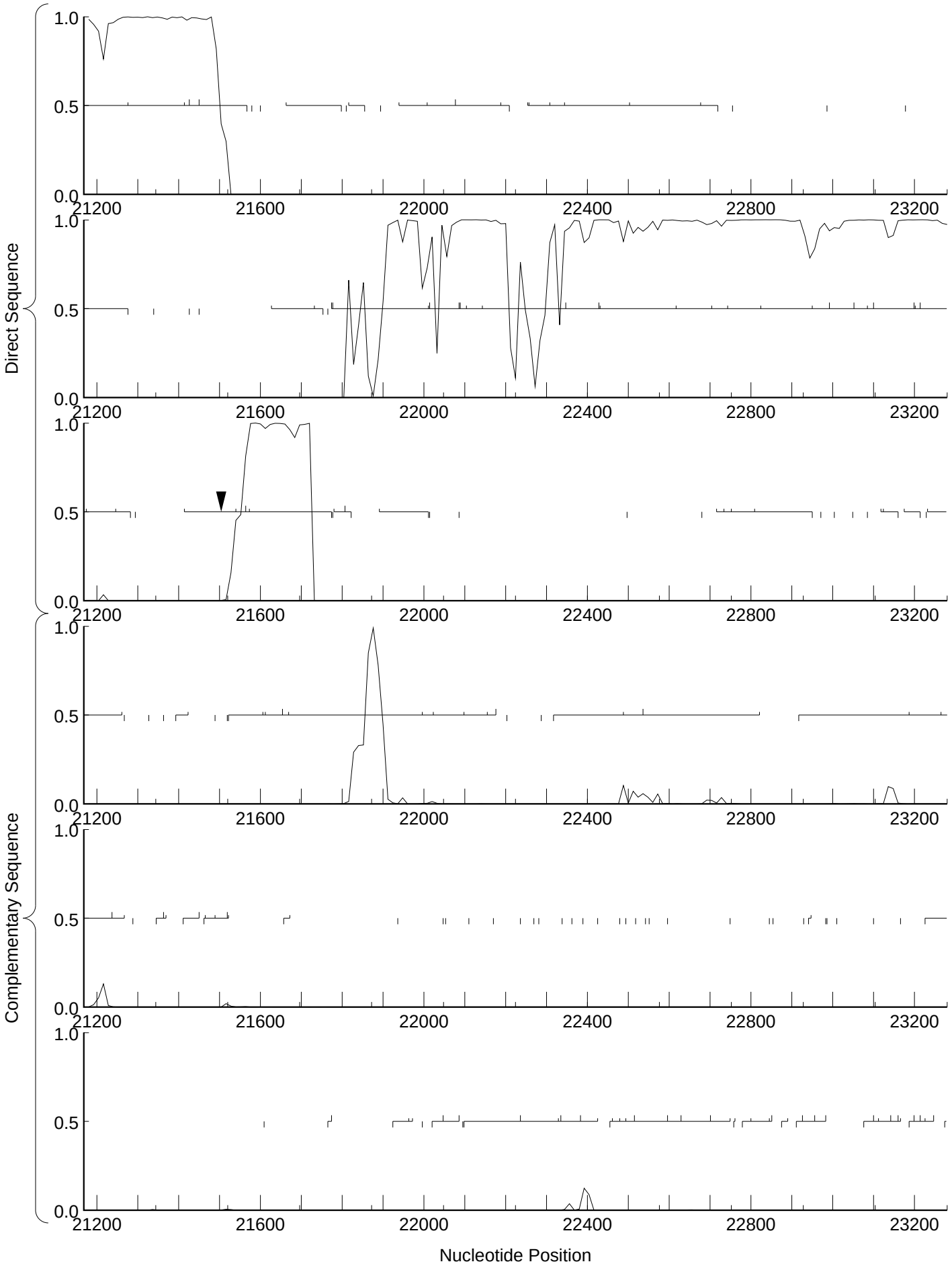
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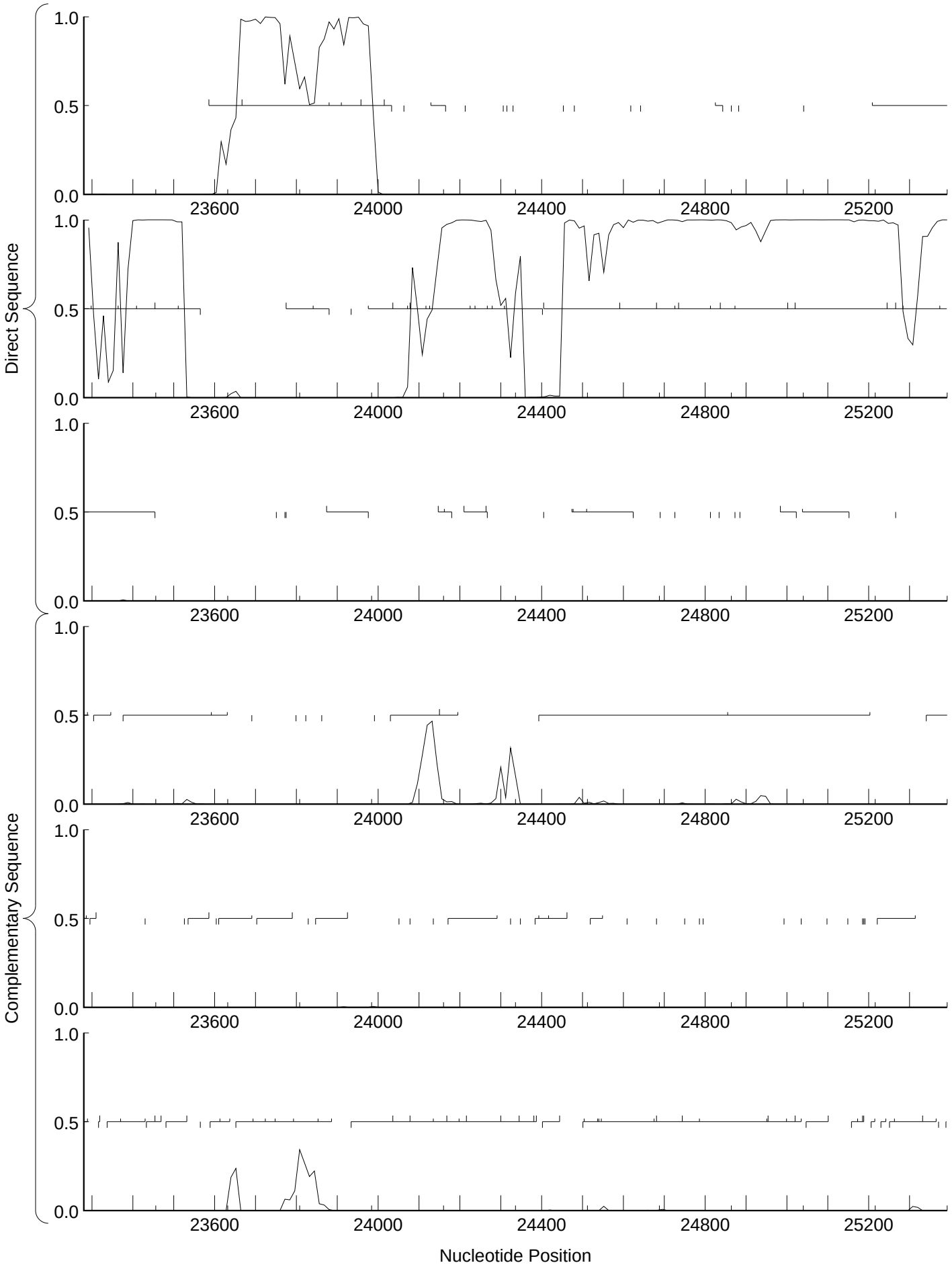
t complete sequence, 52911 bp including 10-base 3' overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 11/27



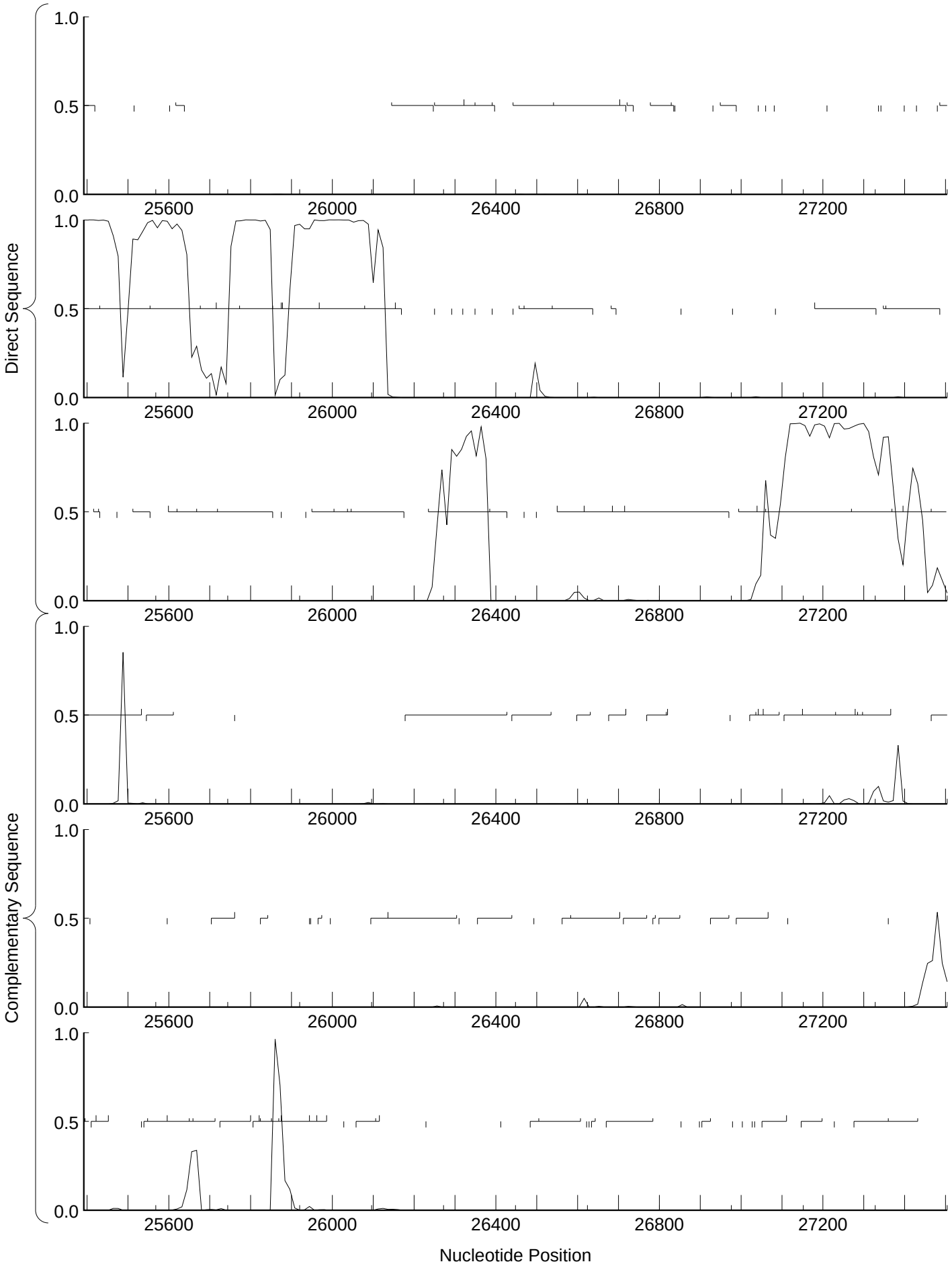
t complete sequence, 52911 bp including 10-base 3' overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 12/27



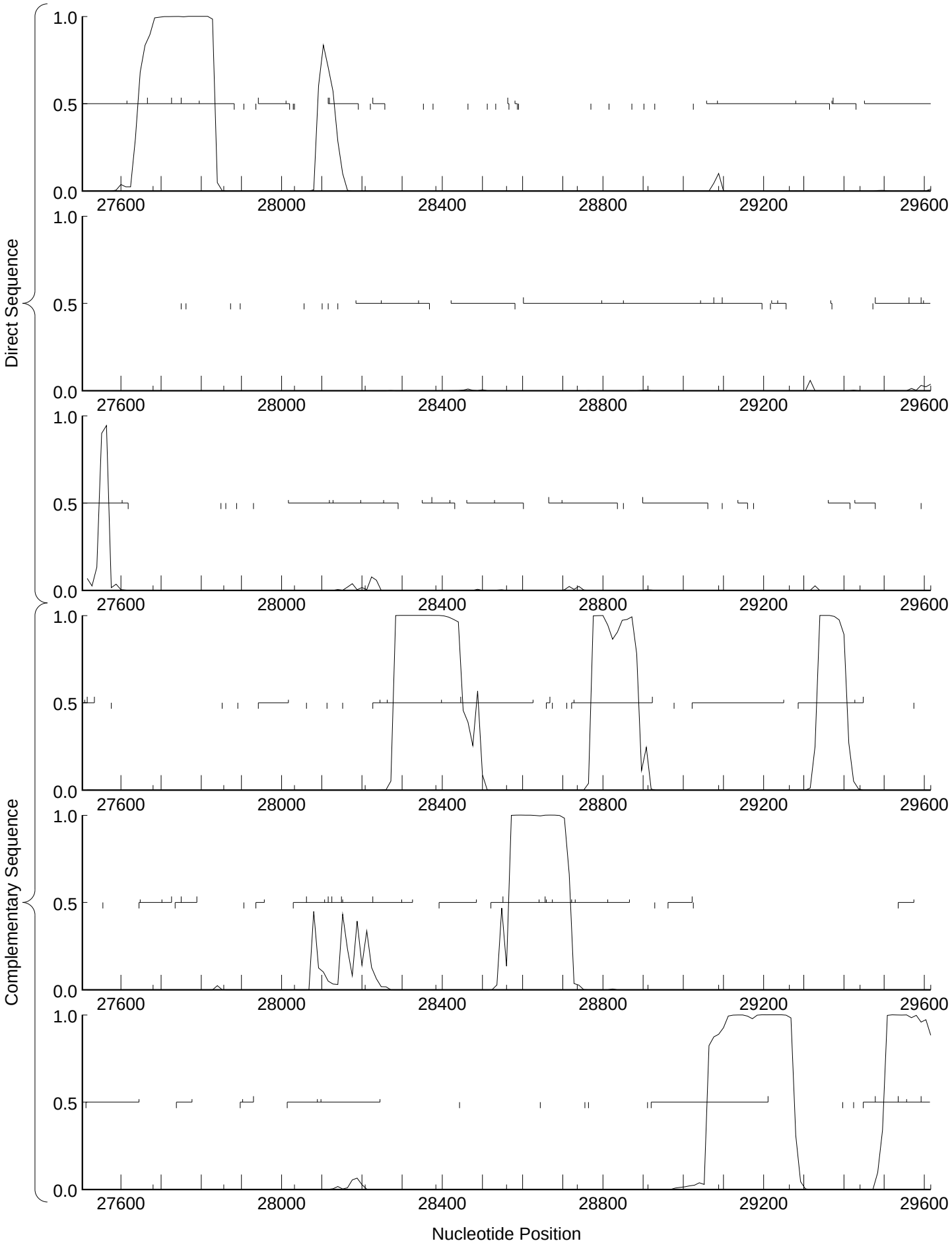
t complete sequence, 52911 bp including 10-base 3' overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 13/27



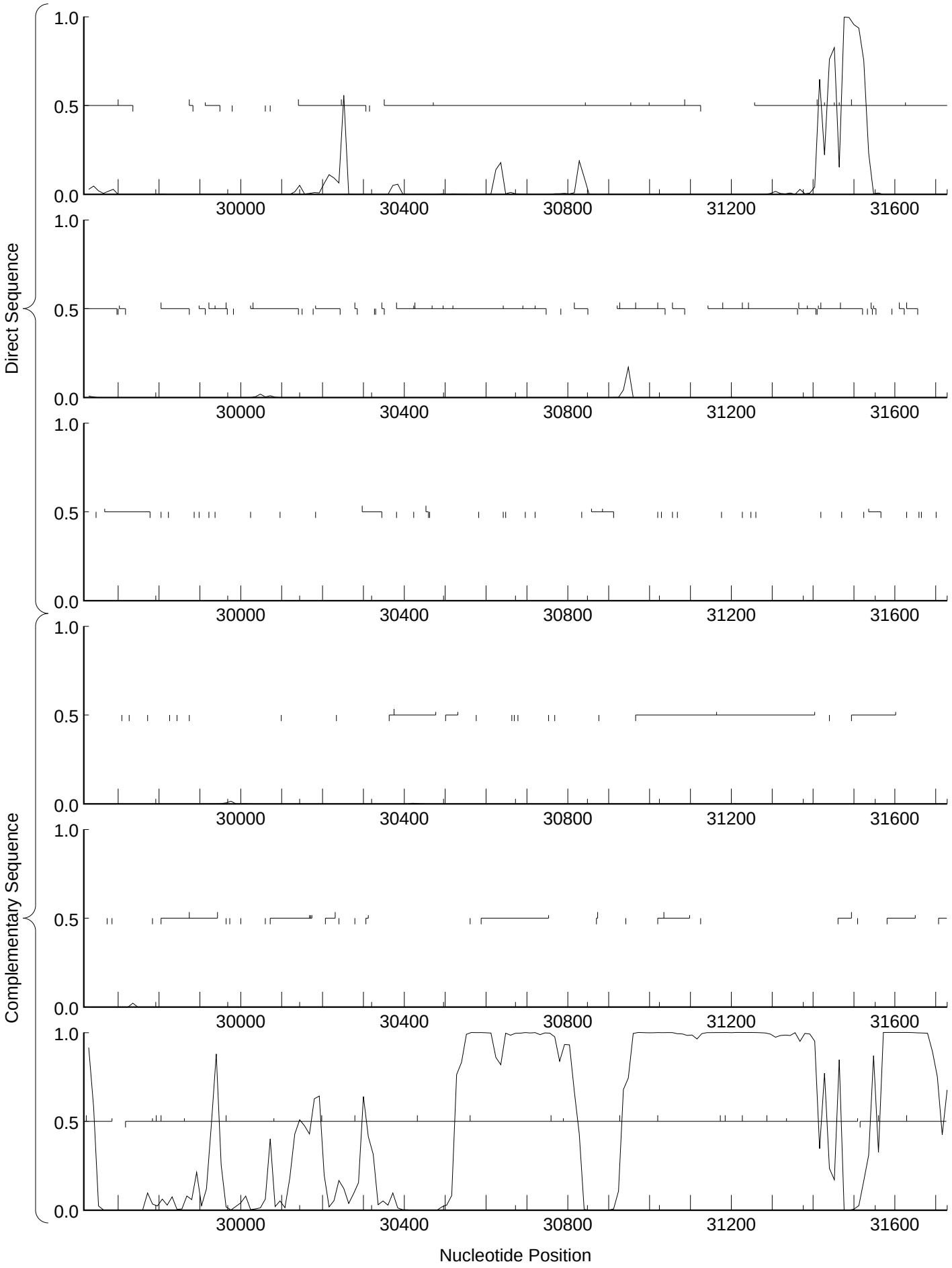
t complete sequence, 52911 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 14/27



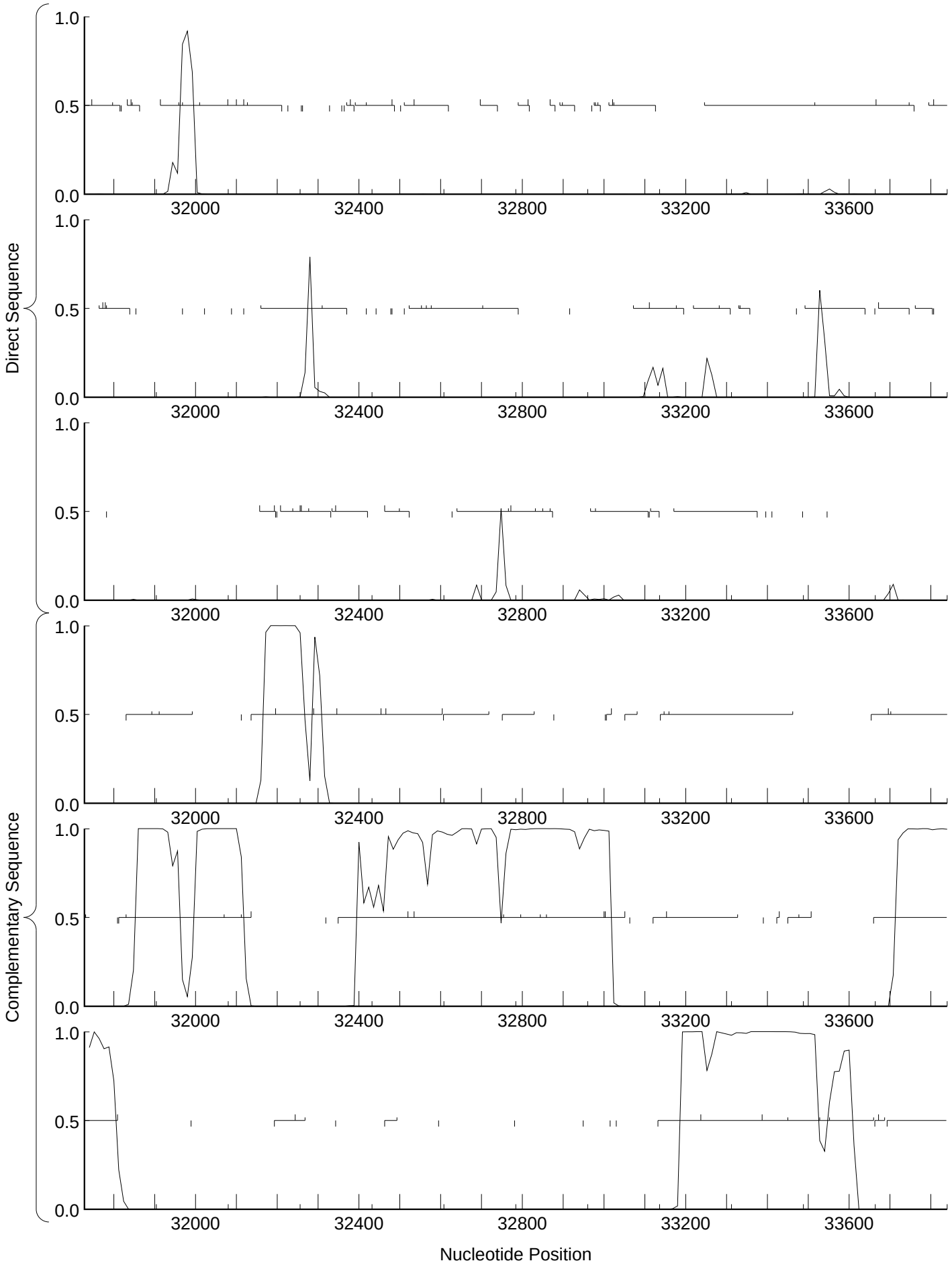
t complete sequence, 52911 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 15/27



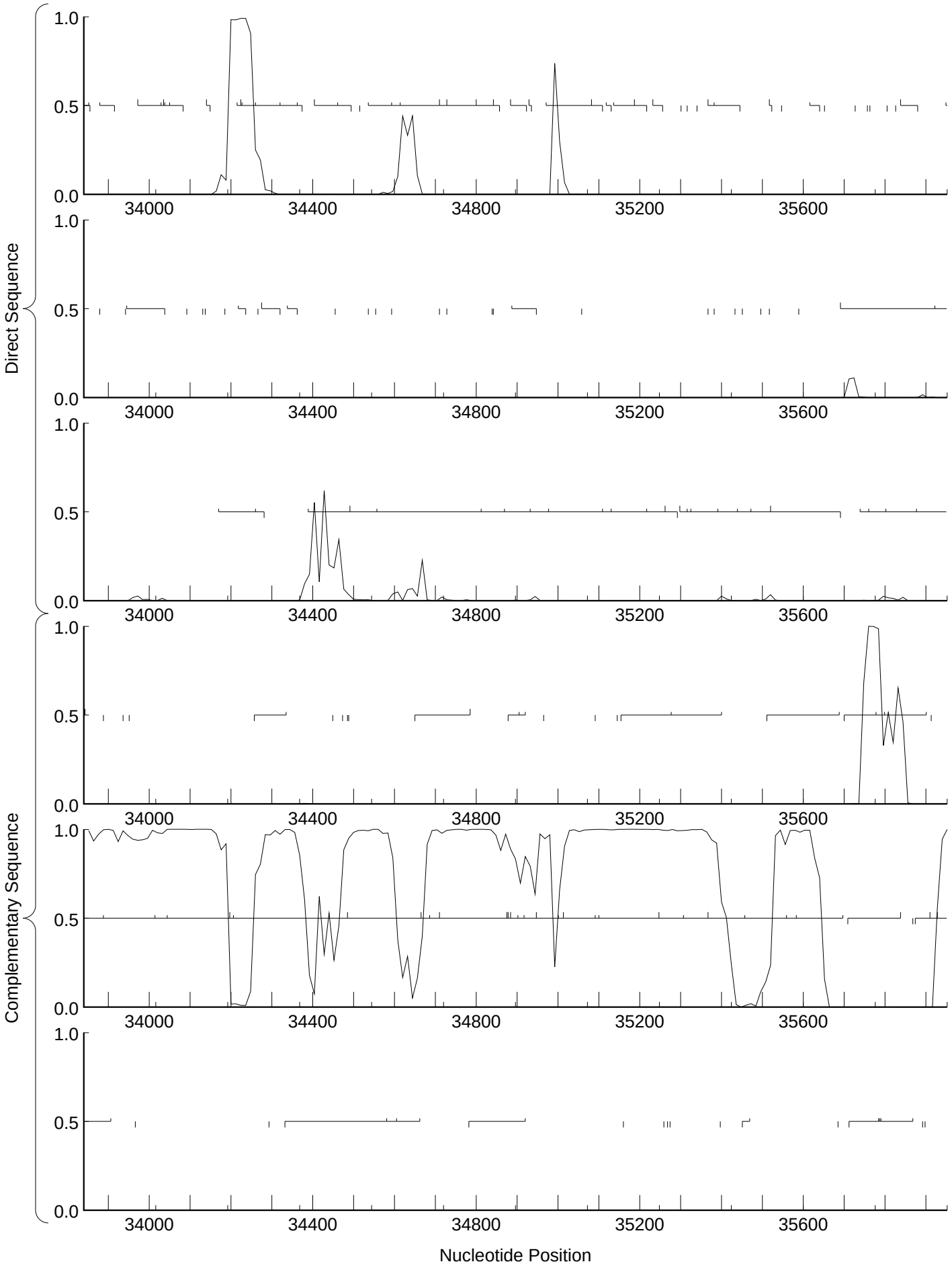
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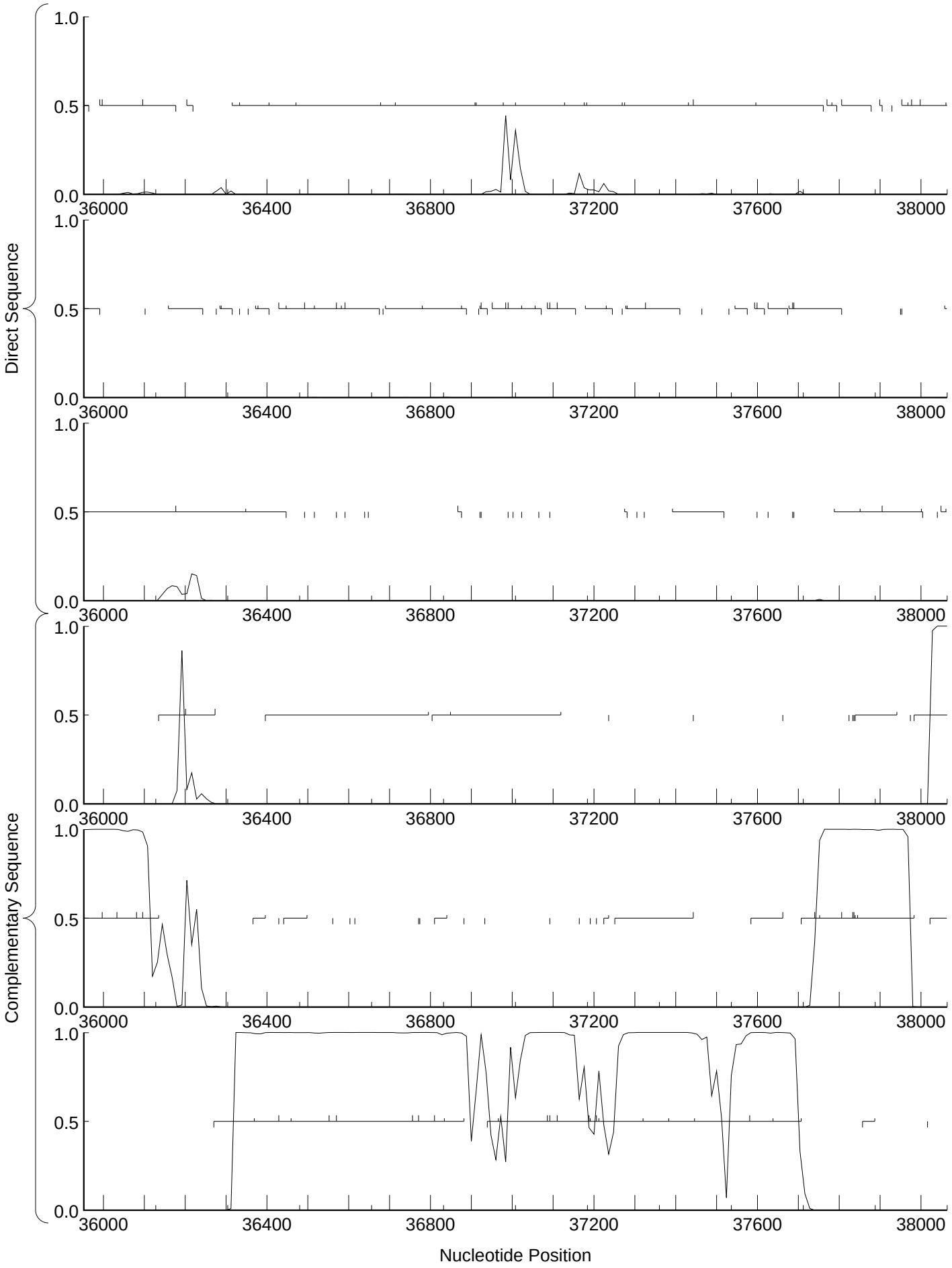
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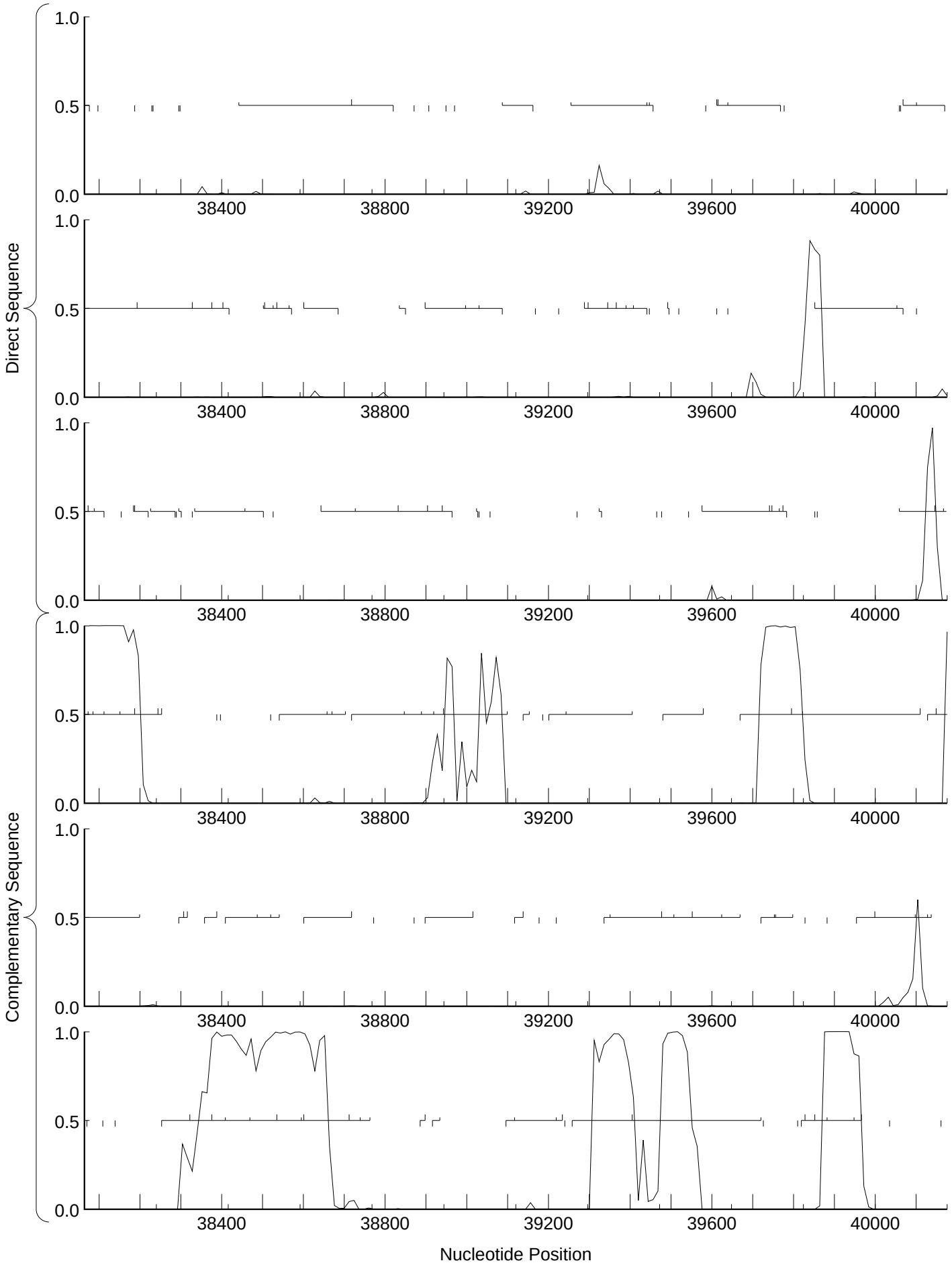
t complete sequence, 52911 bp including 10-base 3' overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 18/27



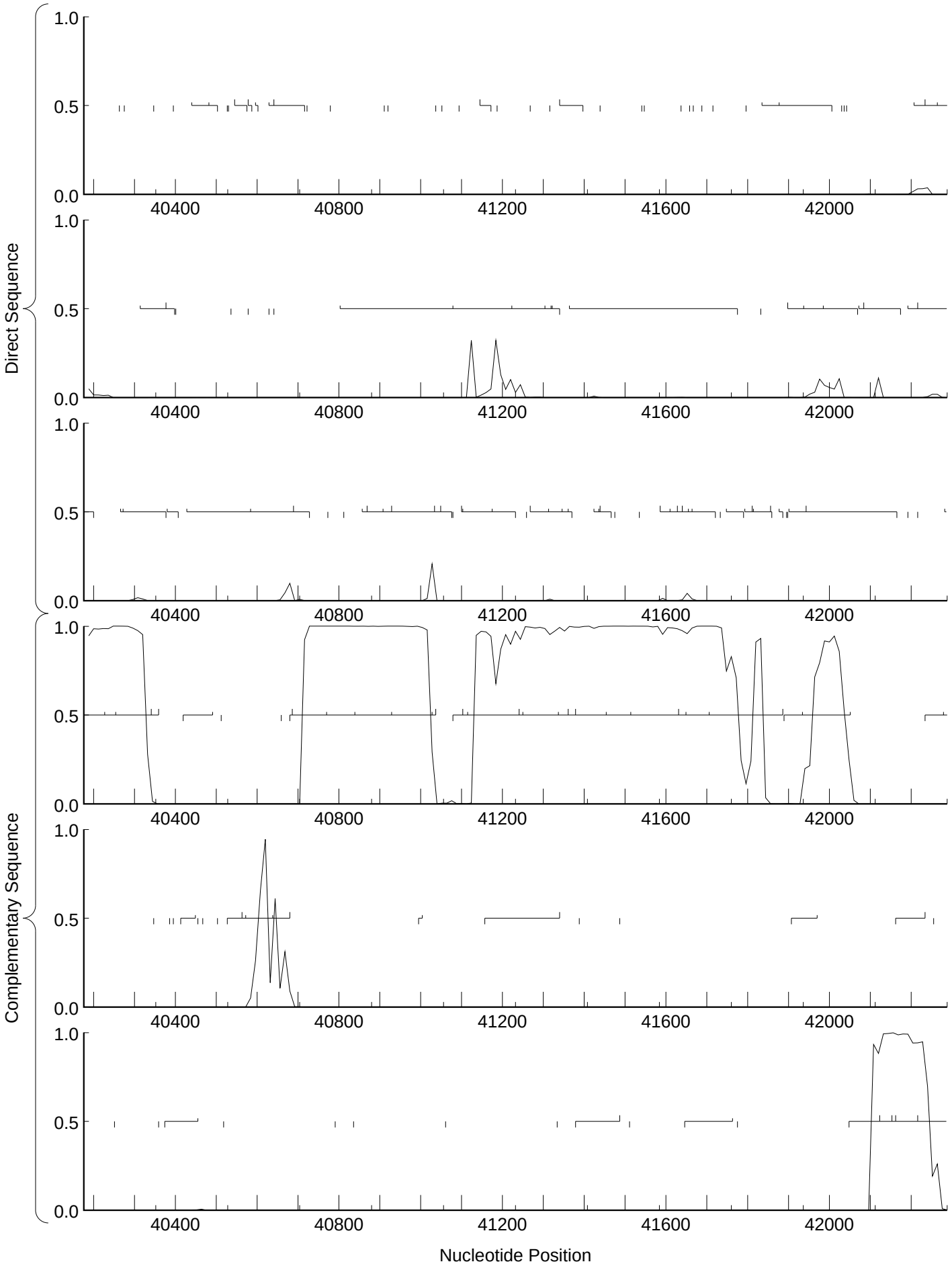
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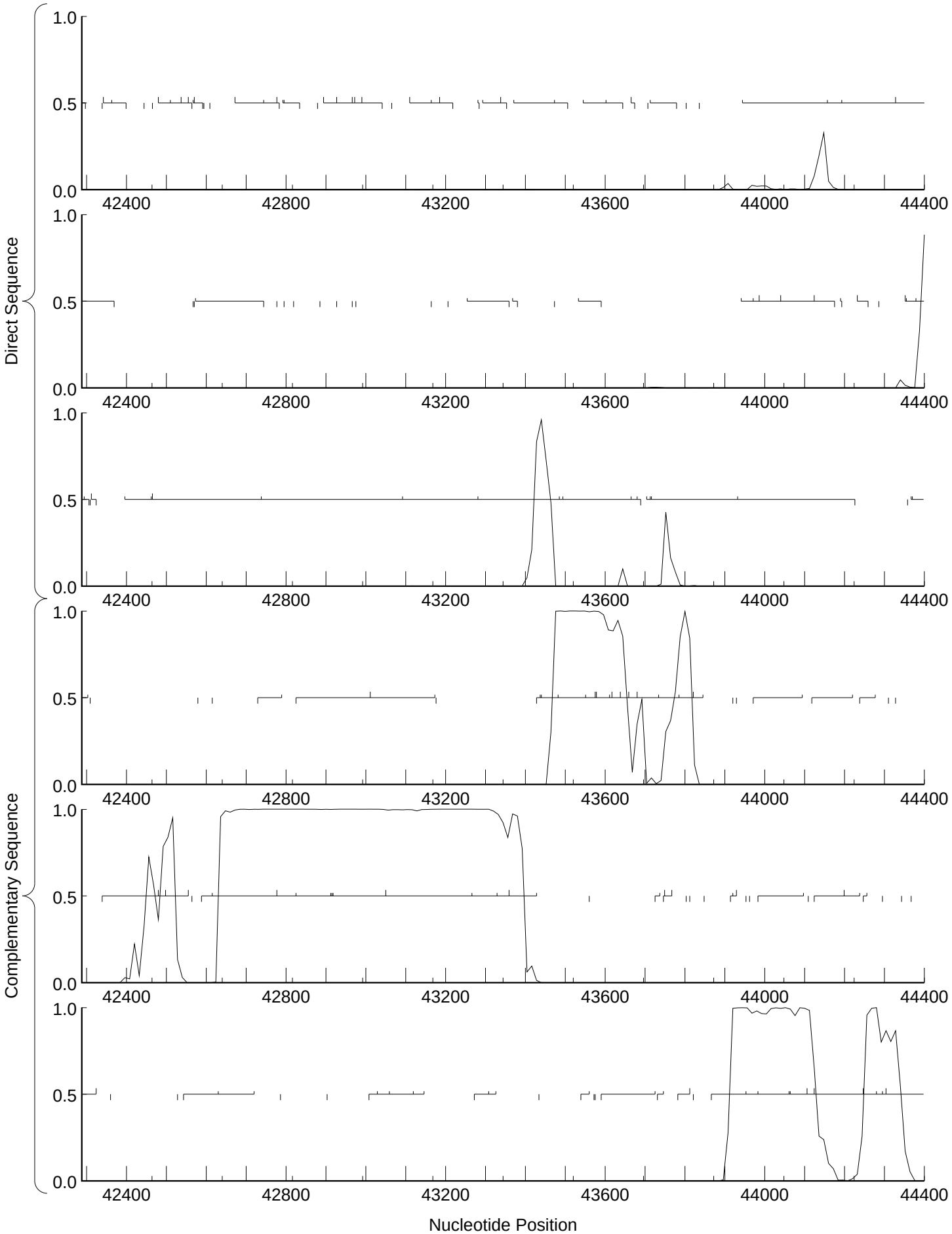
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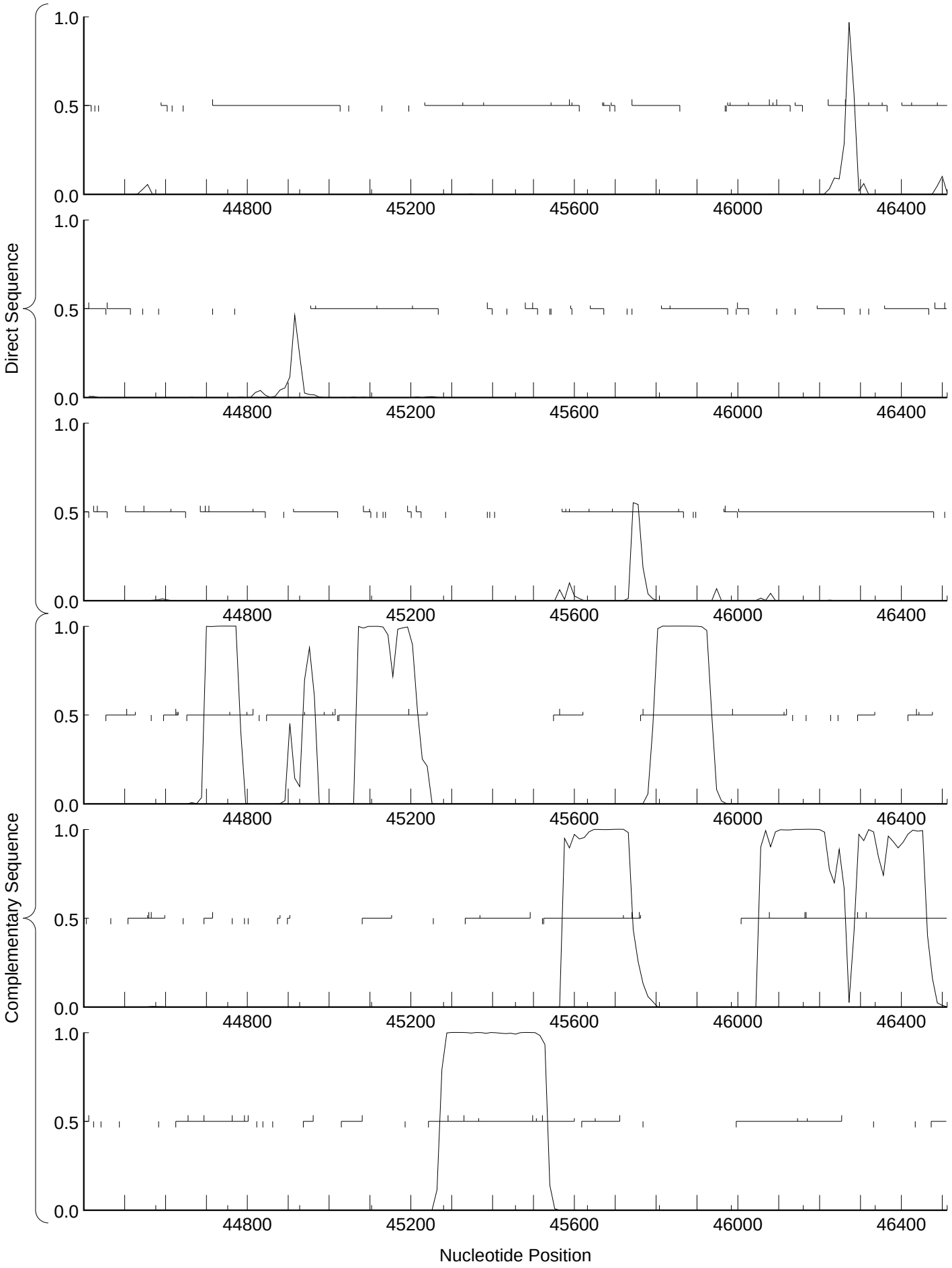
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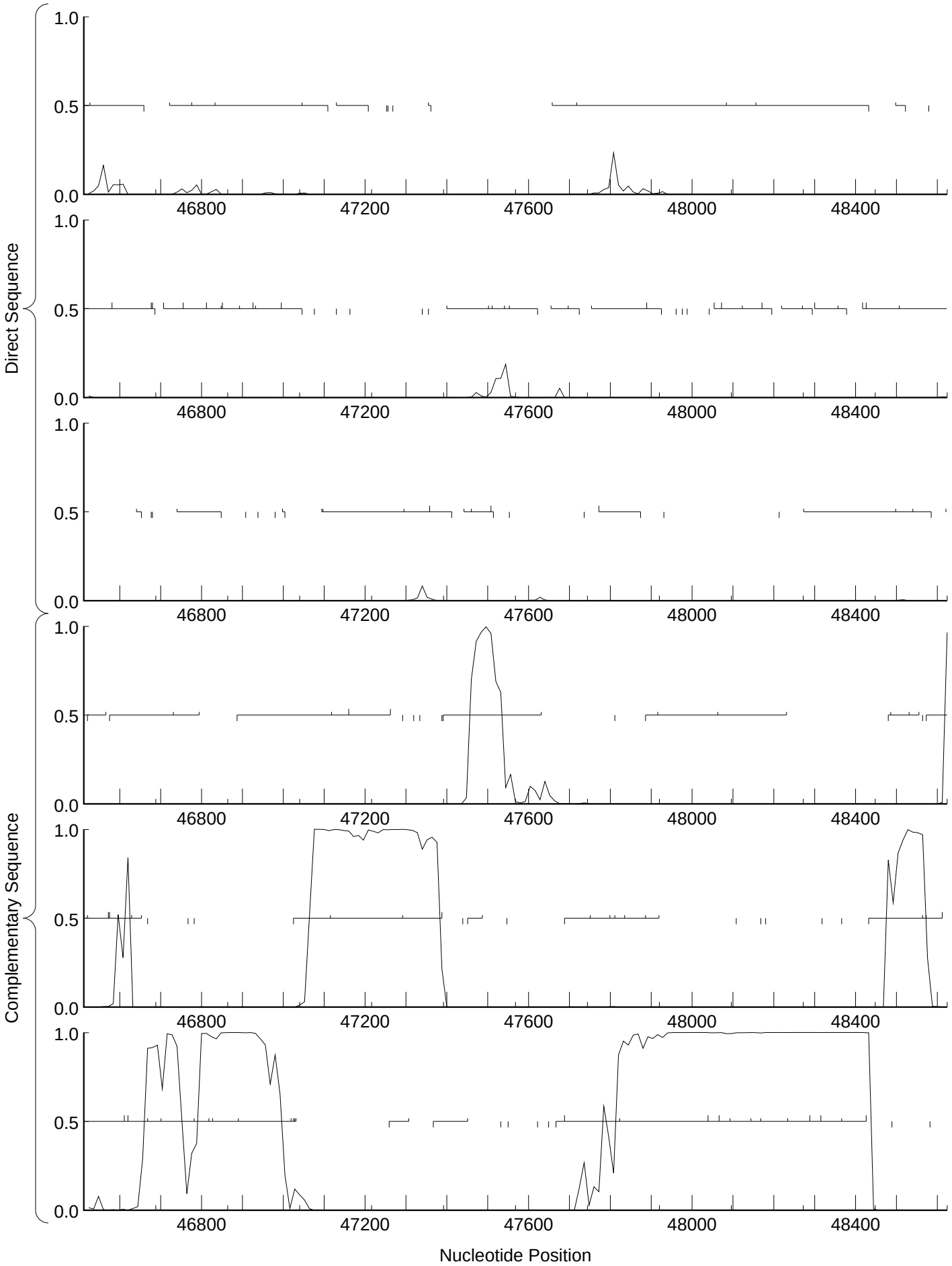
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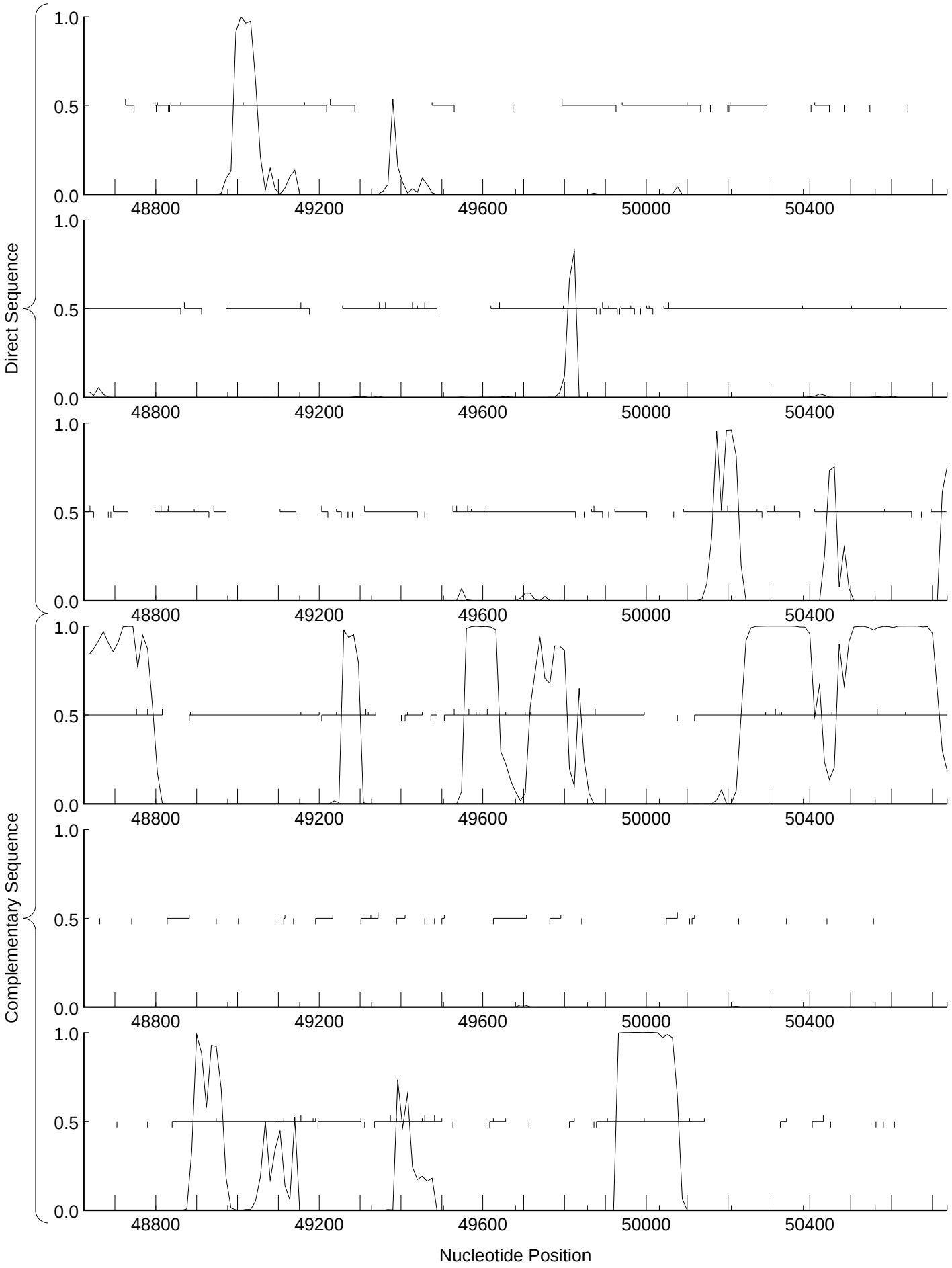
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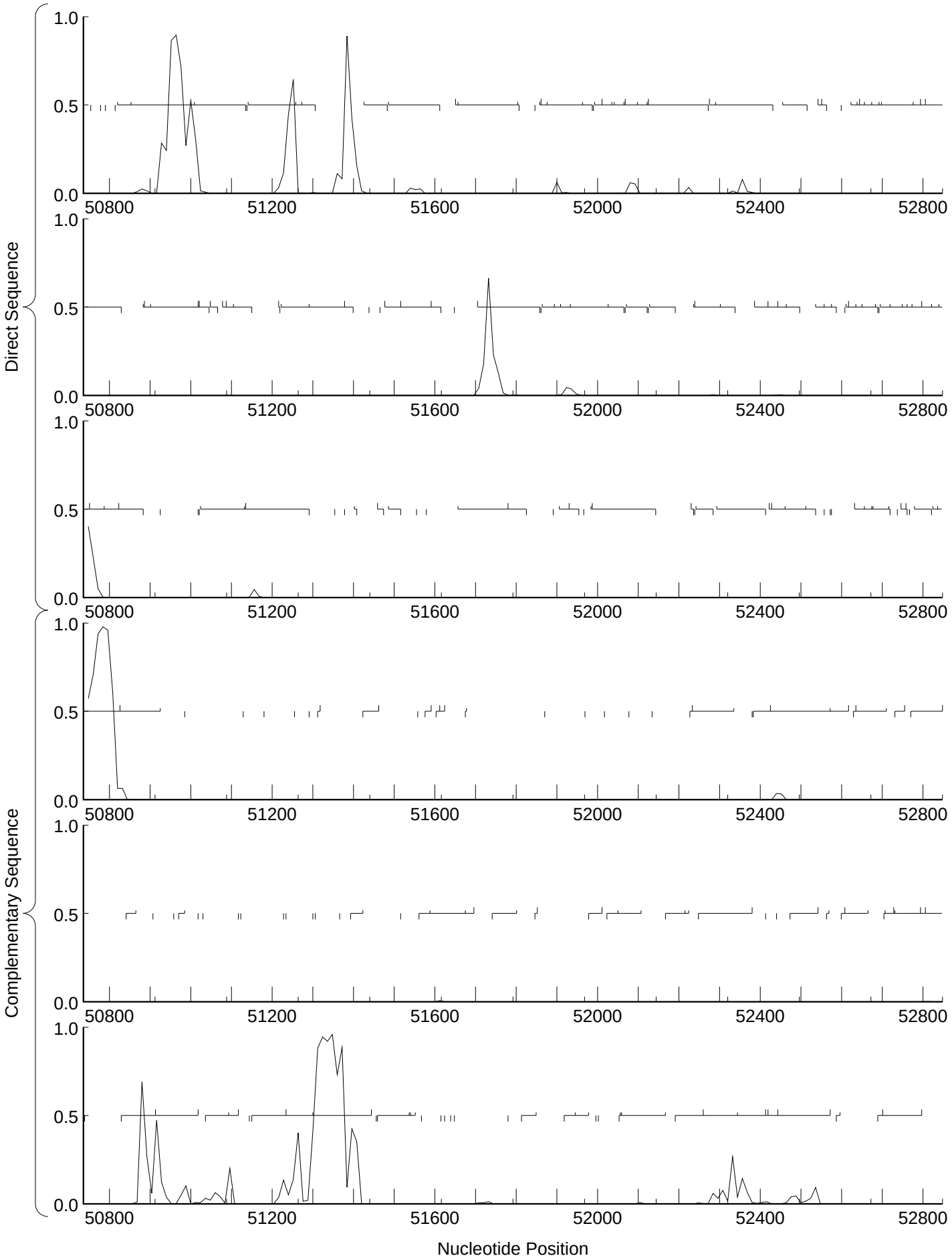
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t complete sequence, 52911 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 26/27



t complete sequence, 52911 bp including 10-base 3’ overhang (CGGTCGGTTA), Cluster A9, Order 4, Window 96, Step 12, 27/27

