GeneMark

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

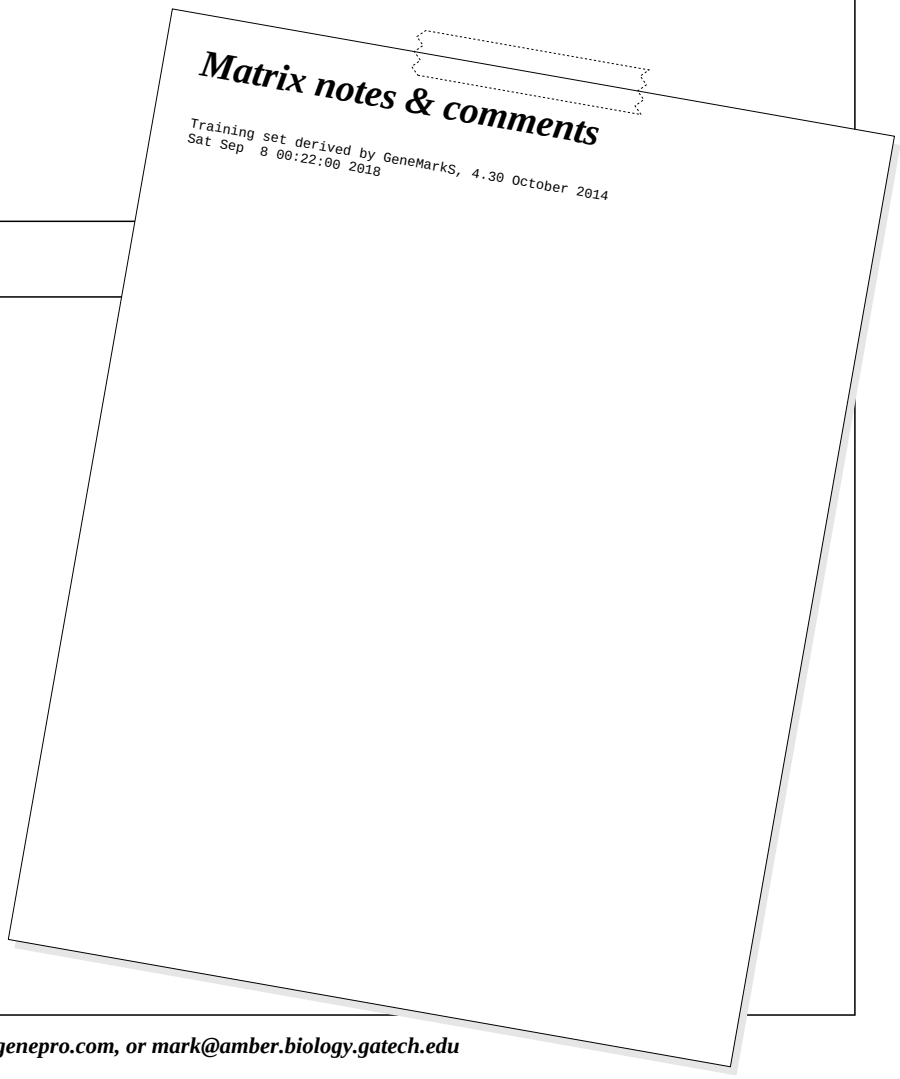
PROGRAM INFORMATION

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Sequence         :  Arthrobacter phage GoCrazy complete sequence, 56809 bp including 9-base 3’ overhang (CGCCGG
Analysis Date    :  3/29/24 at 4:02:11
Pages            :  28
Sequence Length  :  56809 bp
GC Content       :   45.43%

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 INFORMATION

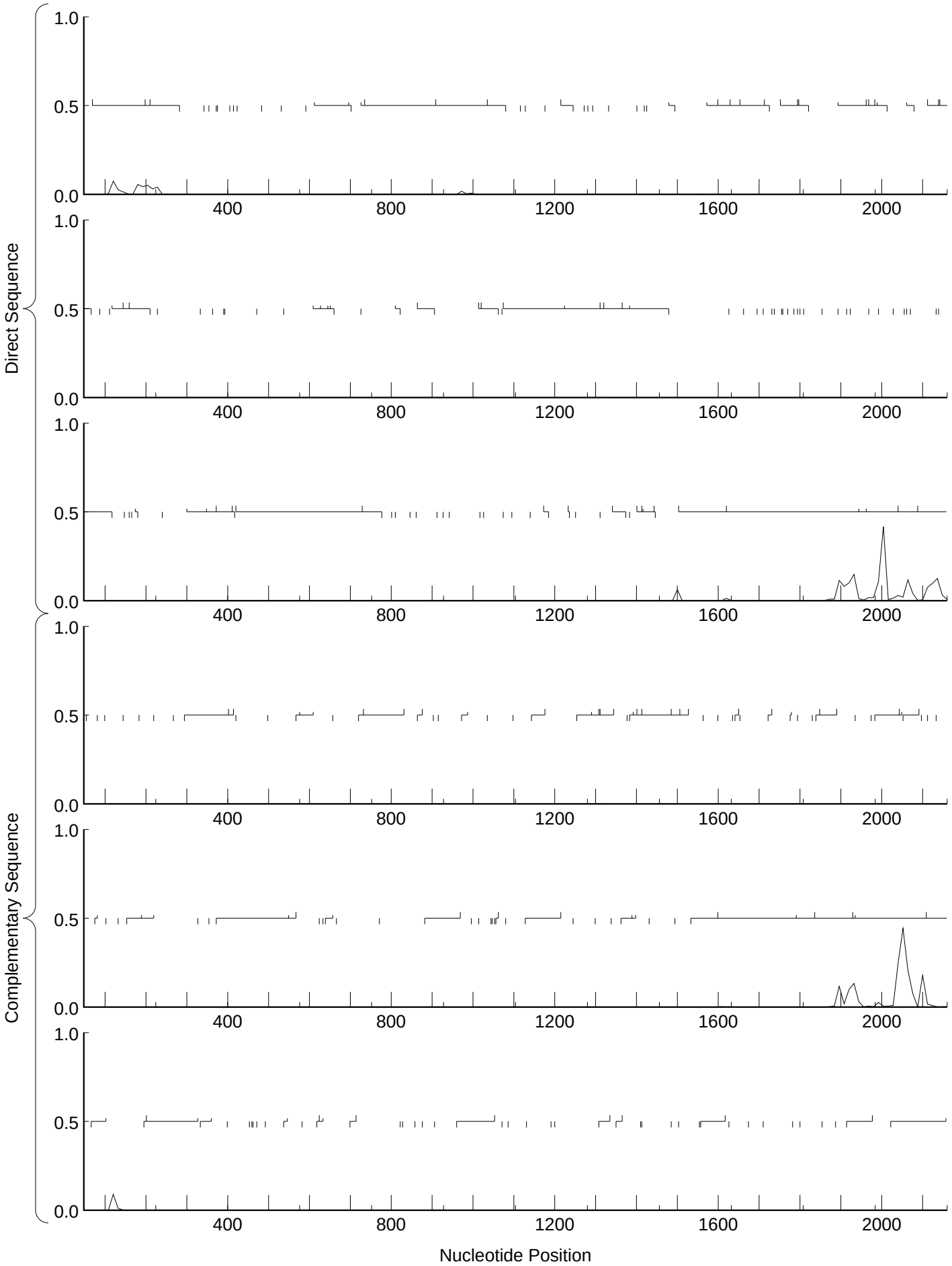
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Matrix     :  -
Author     :  -

Order      :  4

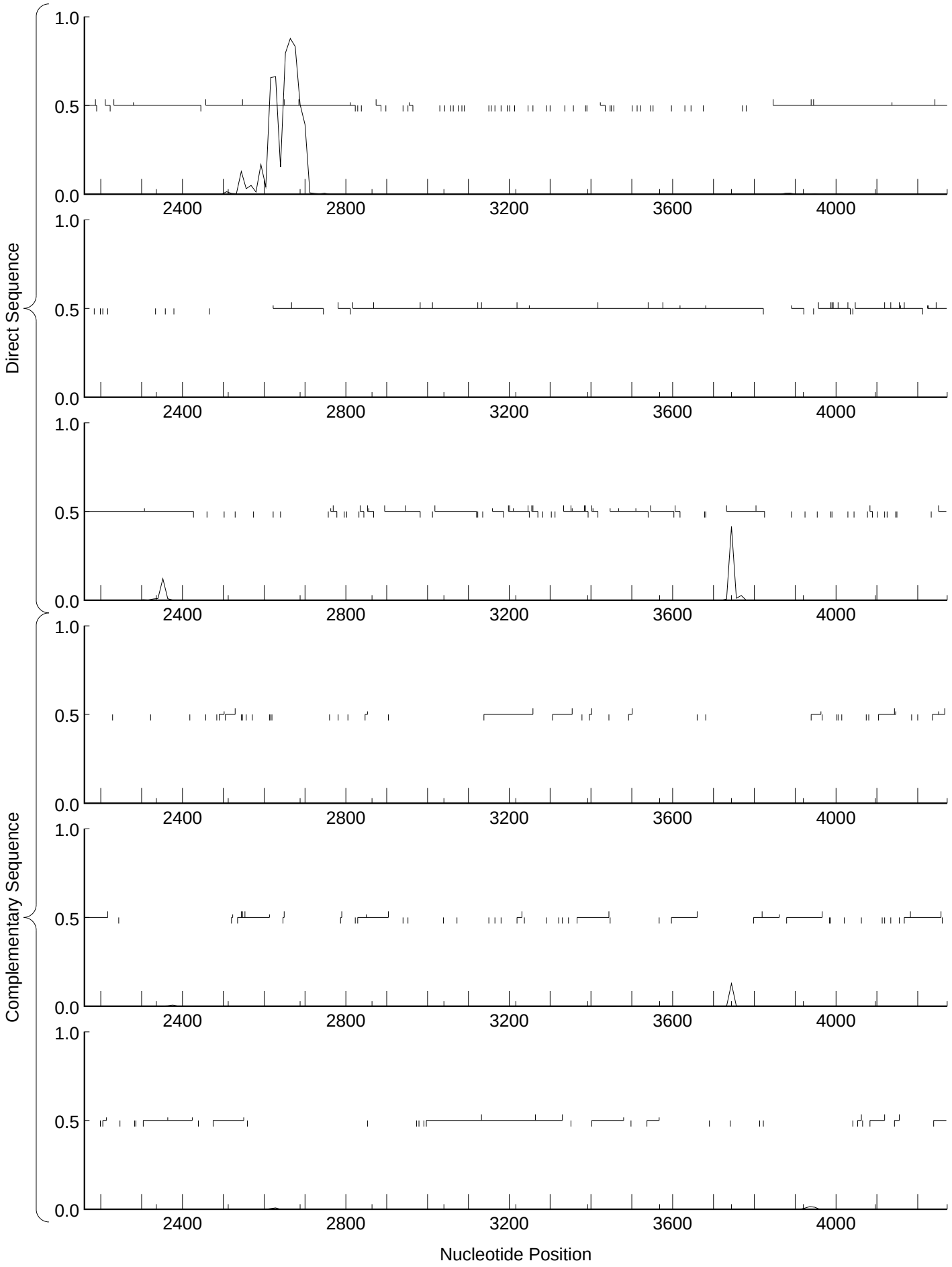

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

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

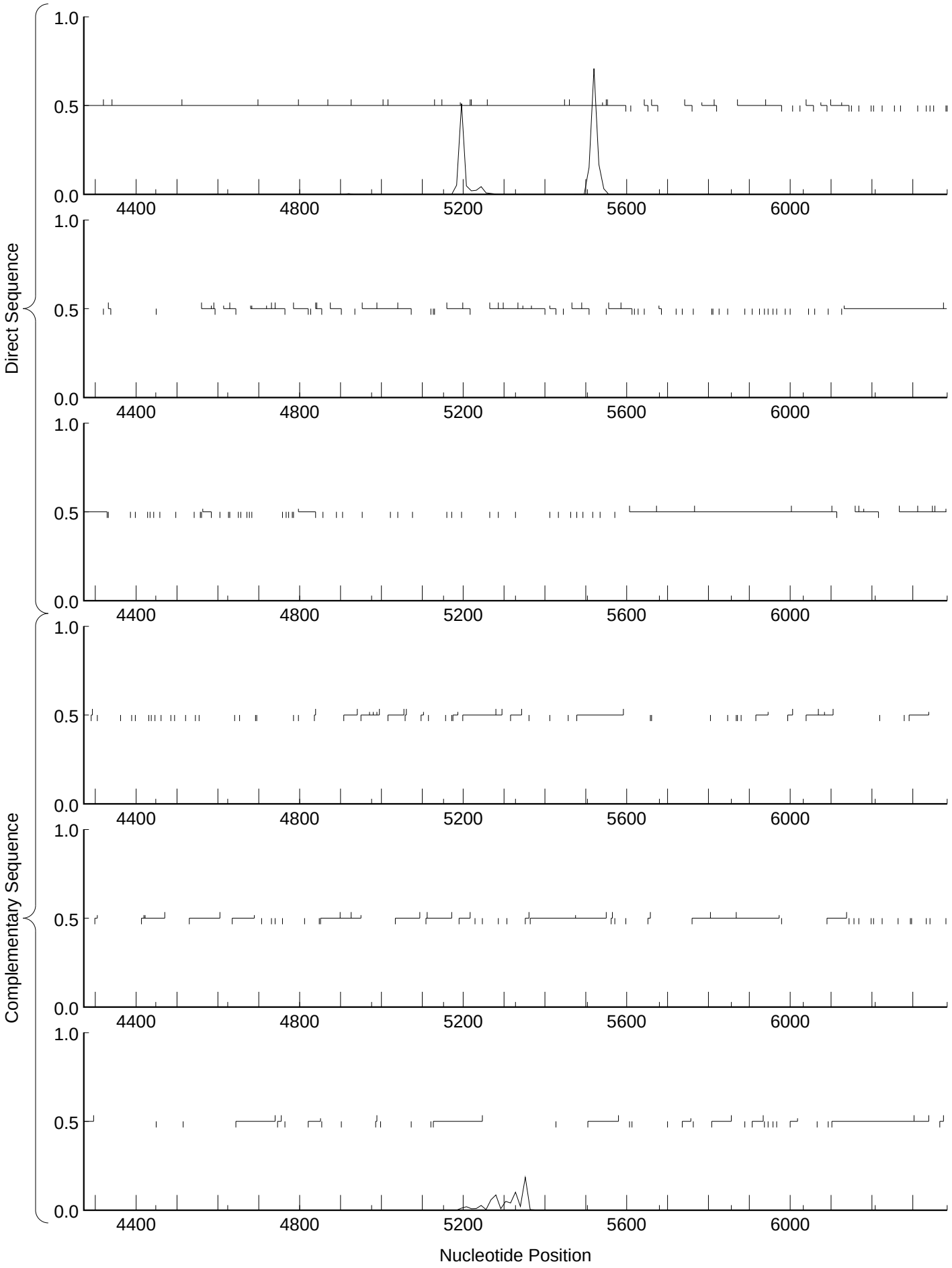
azy complete sequence, 56809 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AM, Order 4, Window 96, Step 12, 2/28



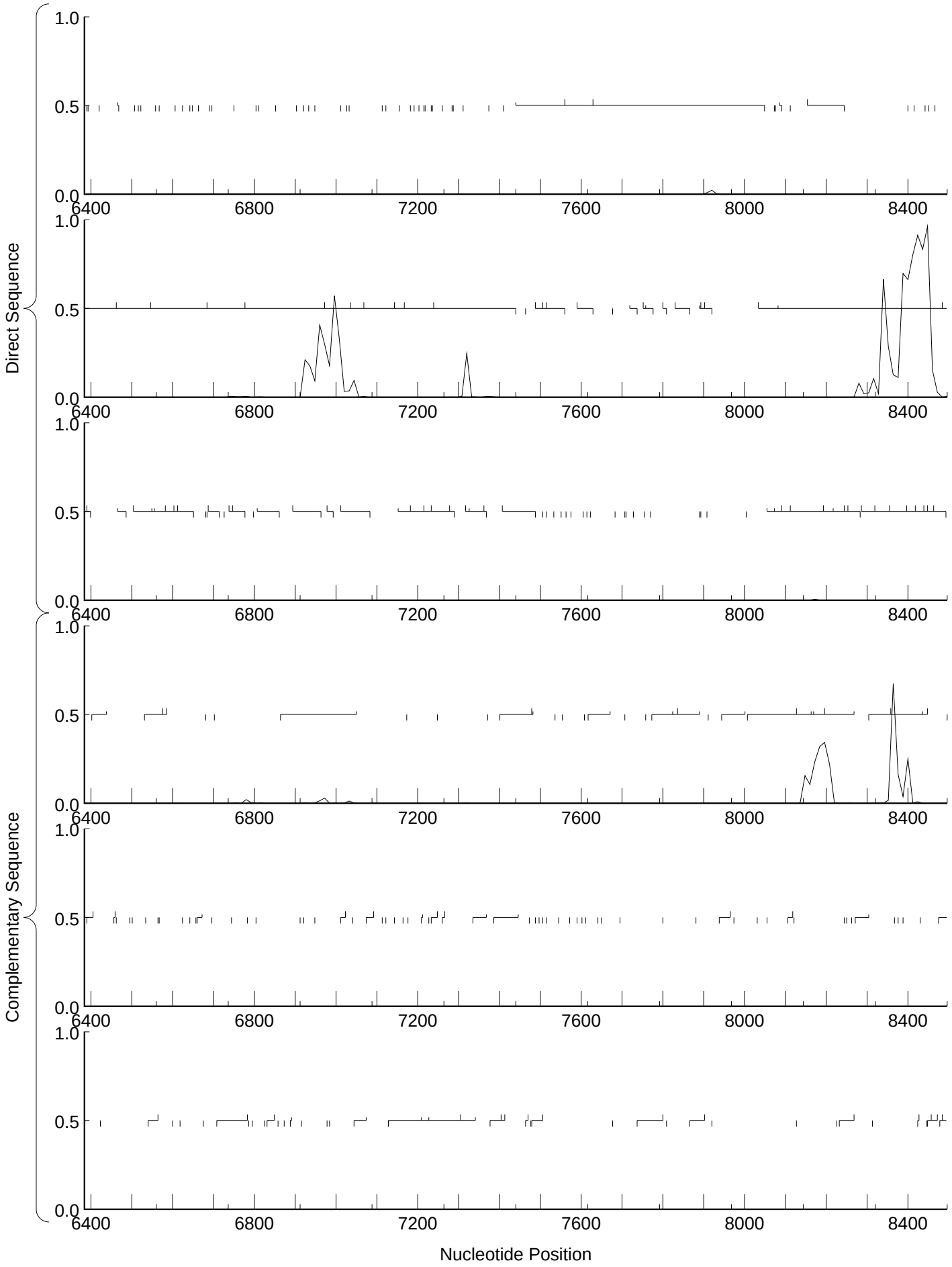
azy complete sequence, 56809 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AM, Order 4, Window 96, Step 12, 3/28



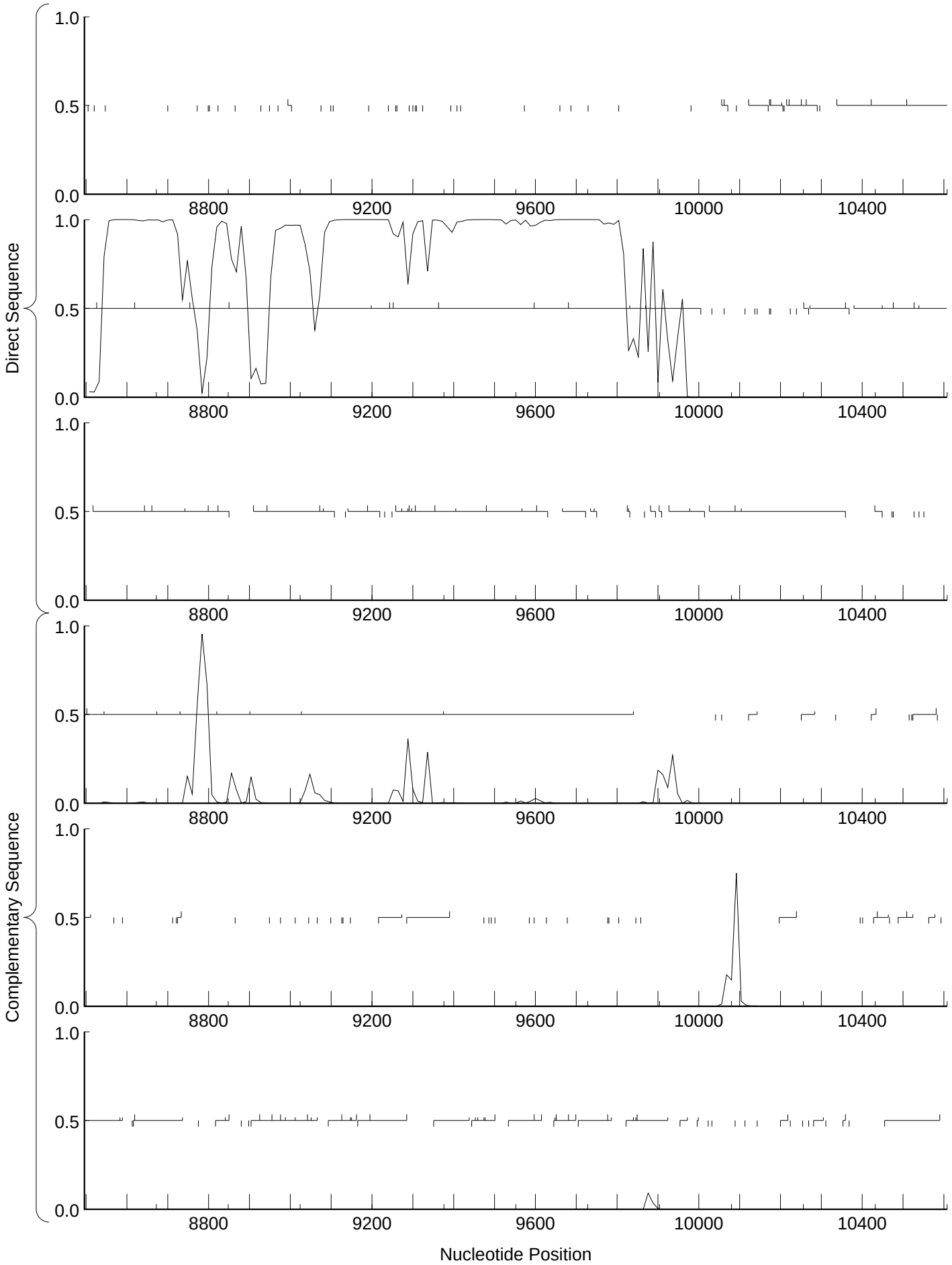
azy complete sequence, 56809 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AM, Order 4, Window 96, Step 12, 4/28



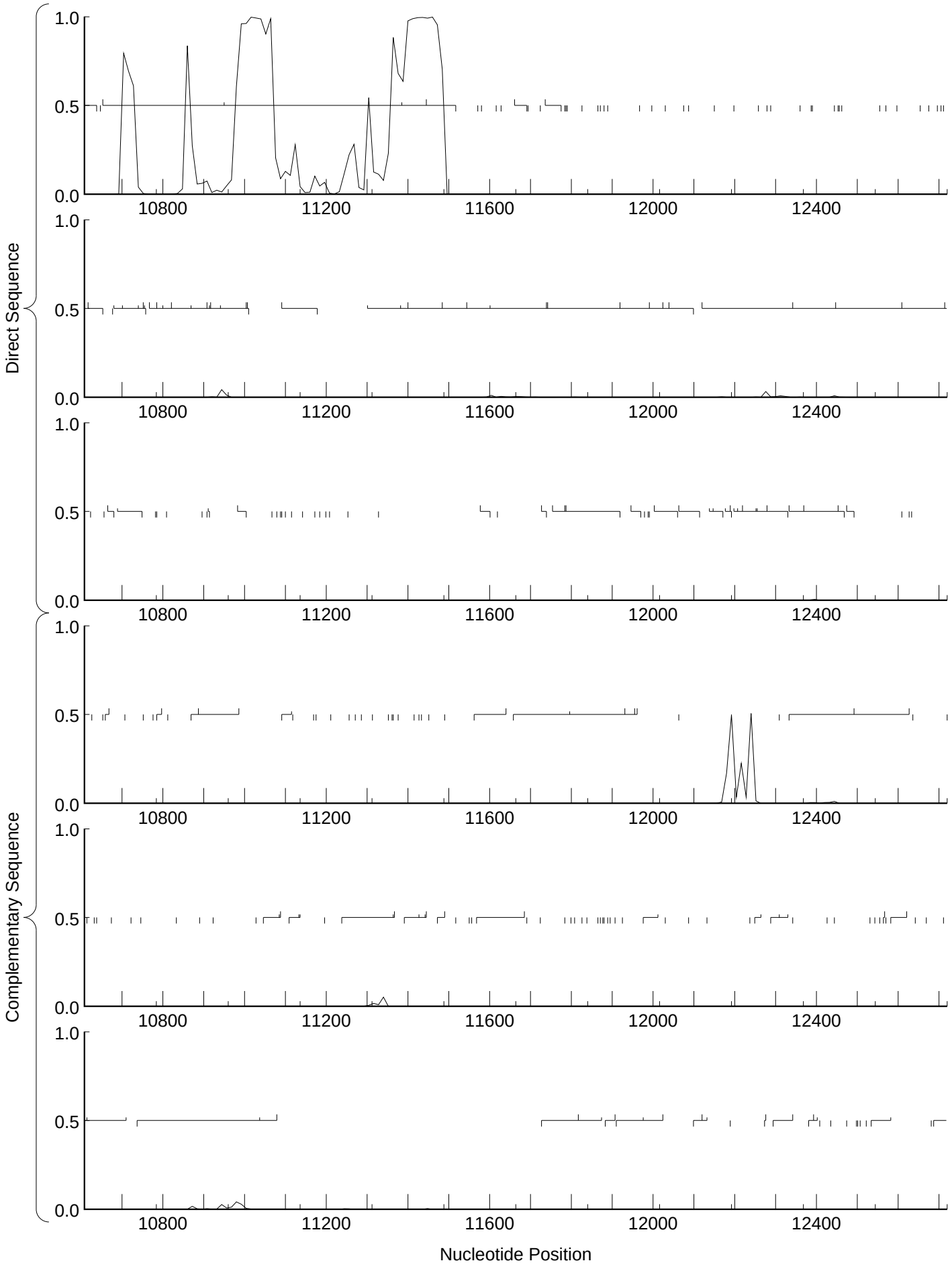
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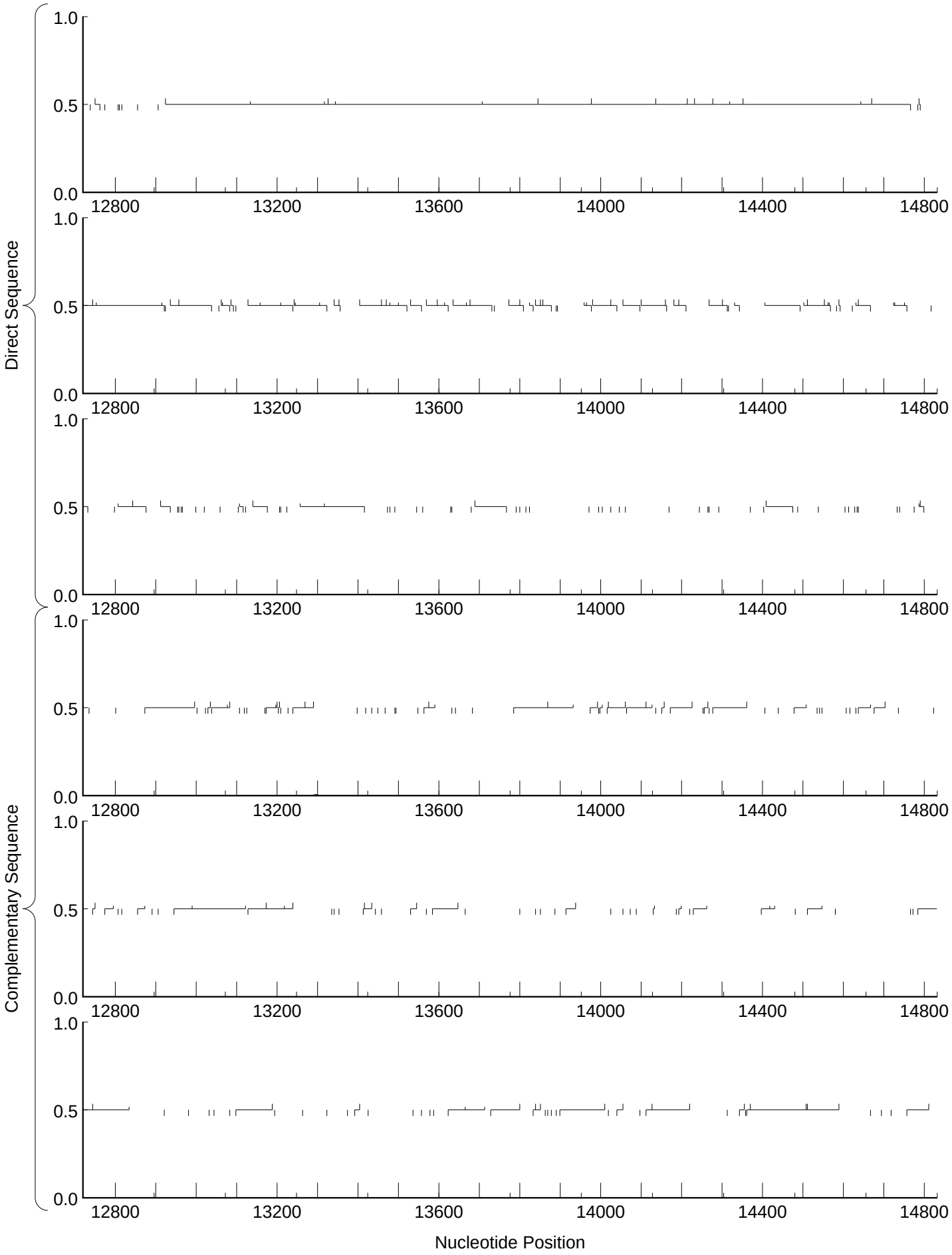
azy complete sequence, 56809 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AM, Order 4, Window 96, Step 12, 6/28



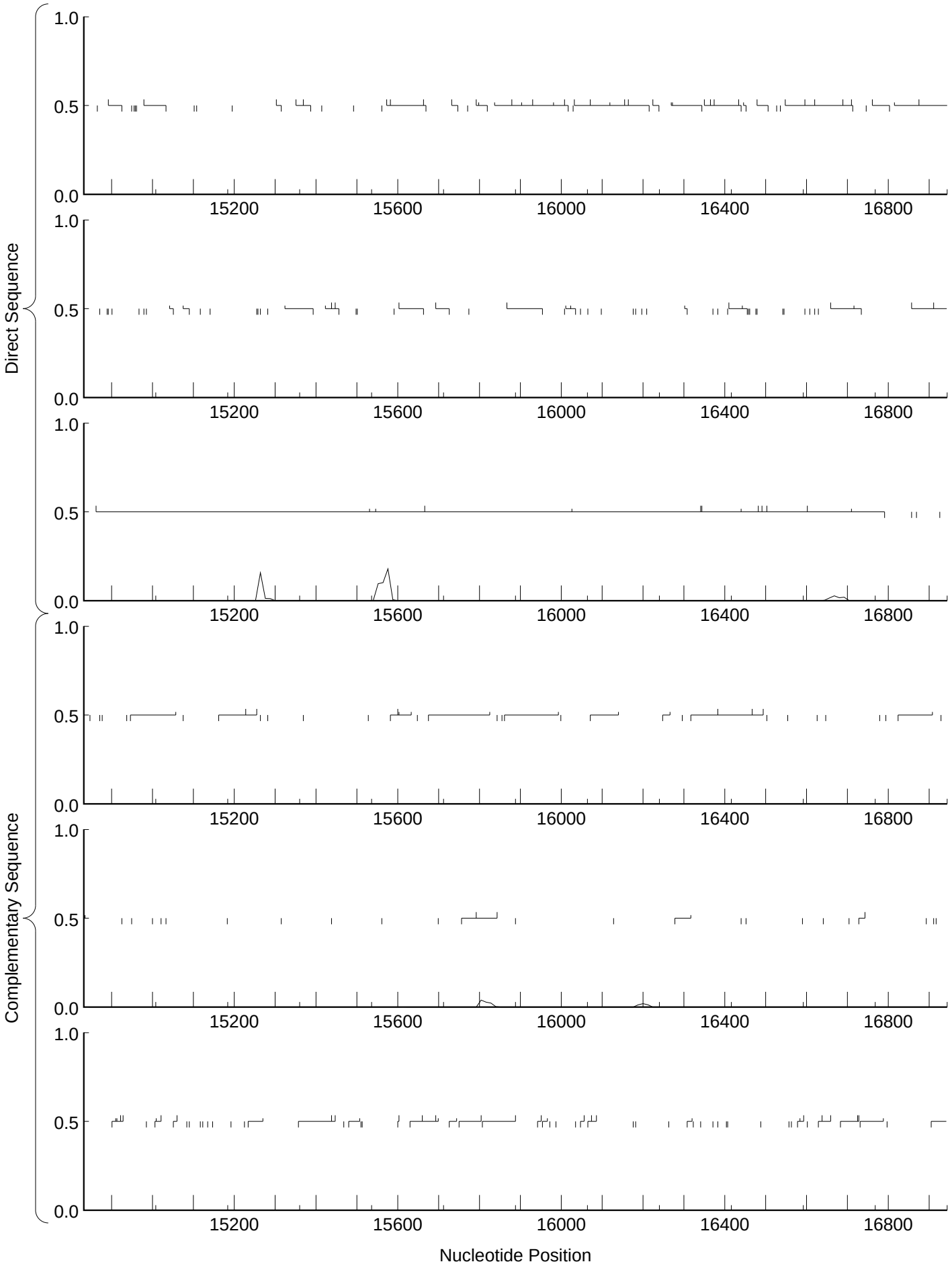
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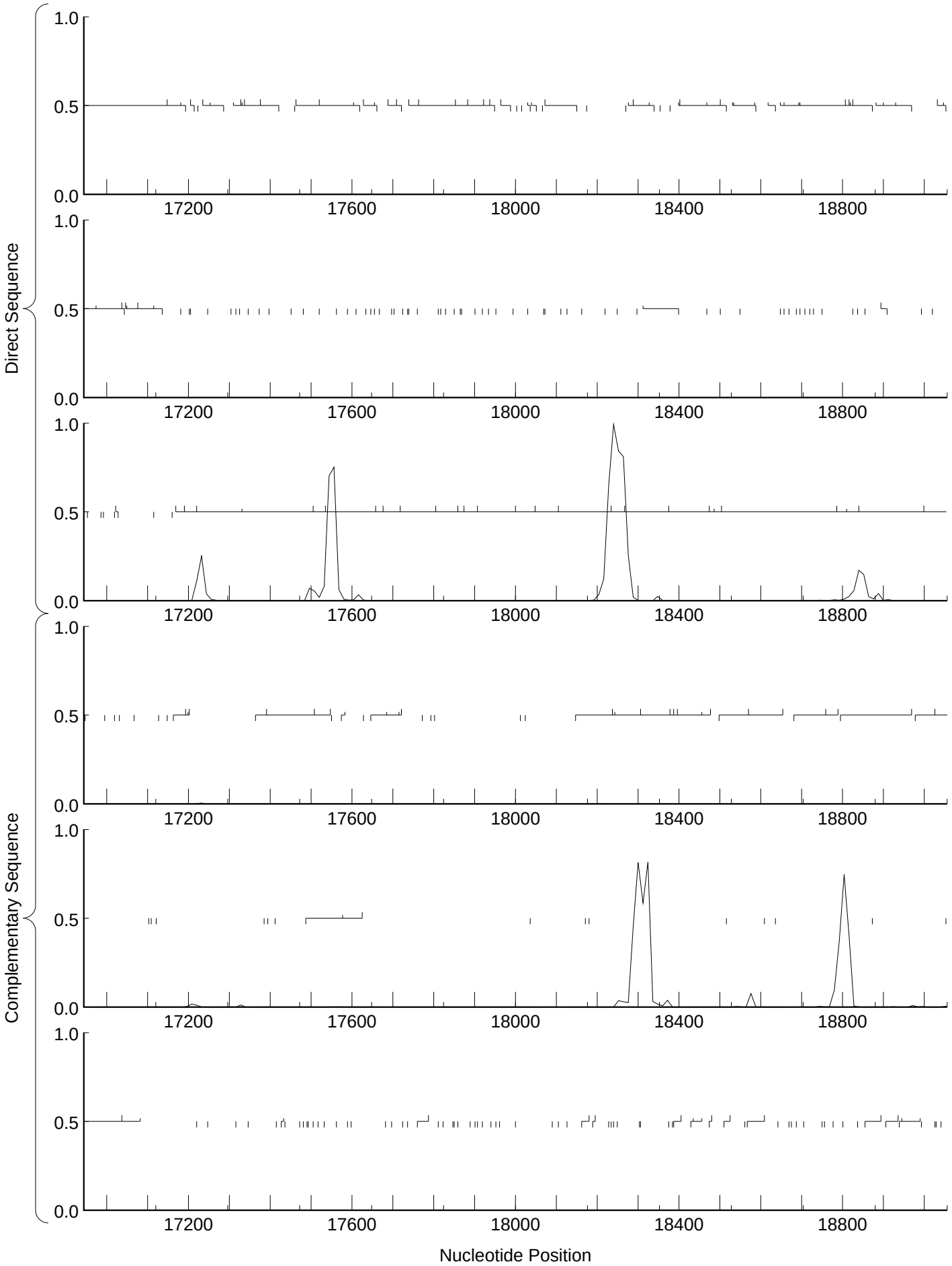
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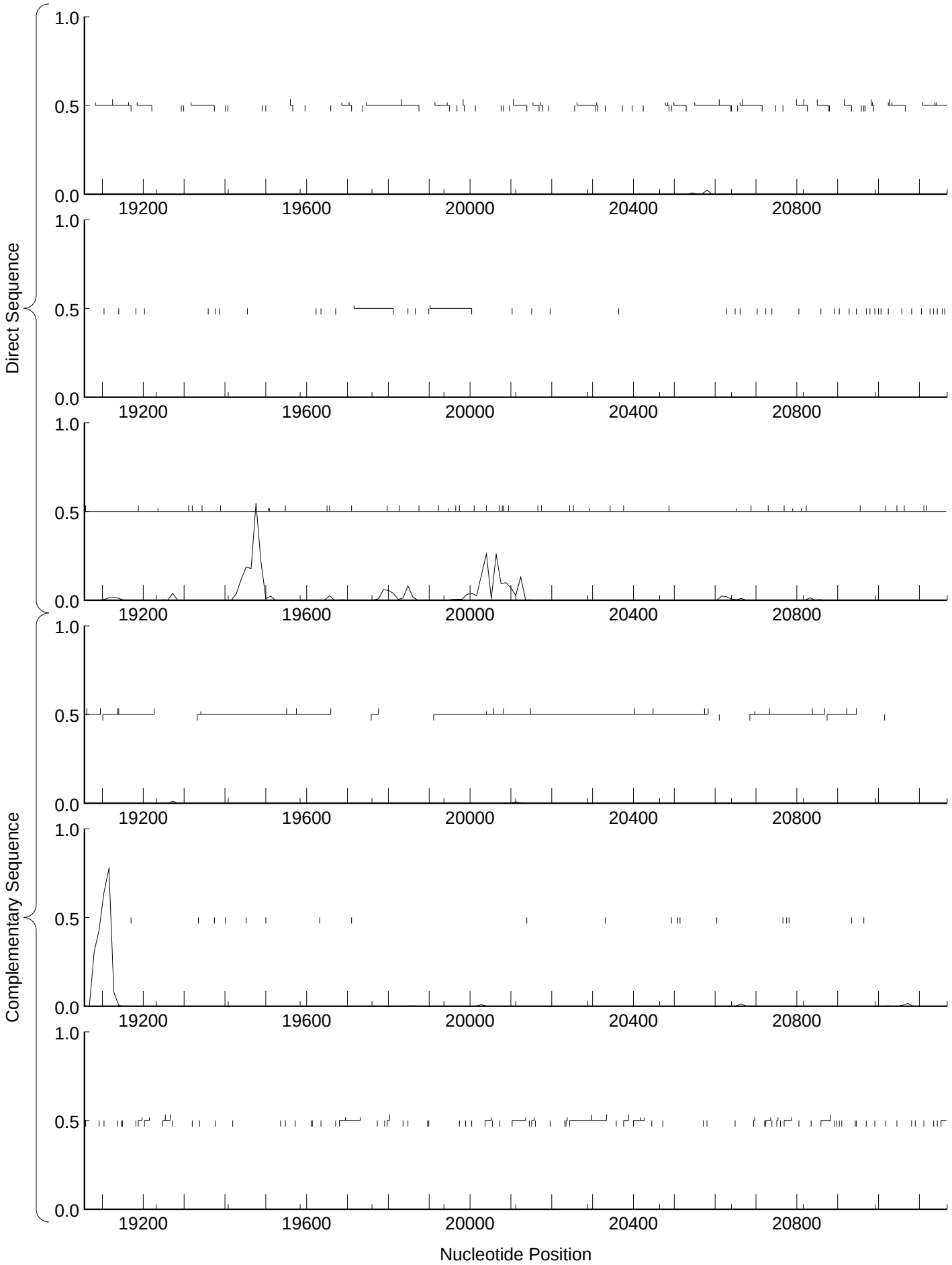
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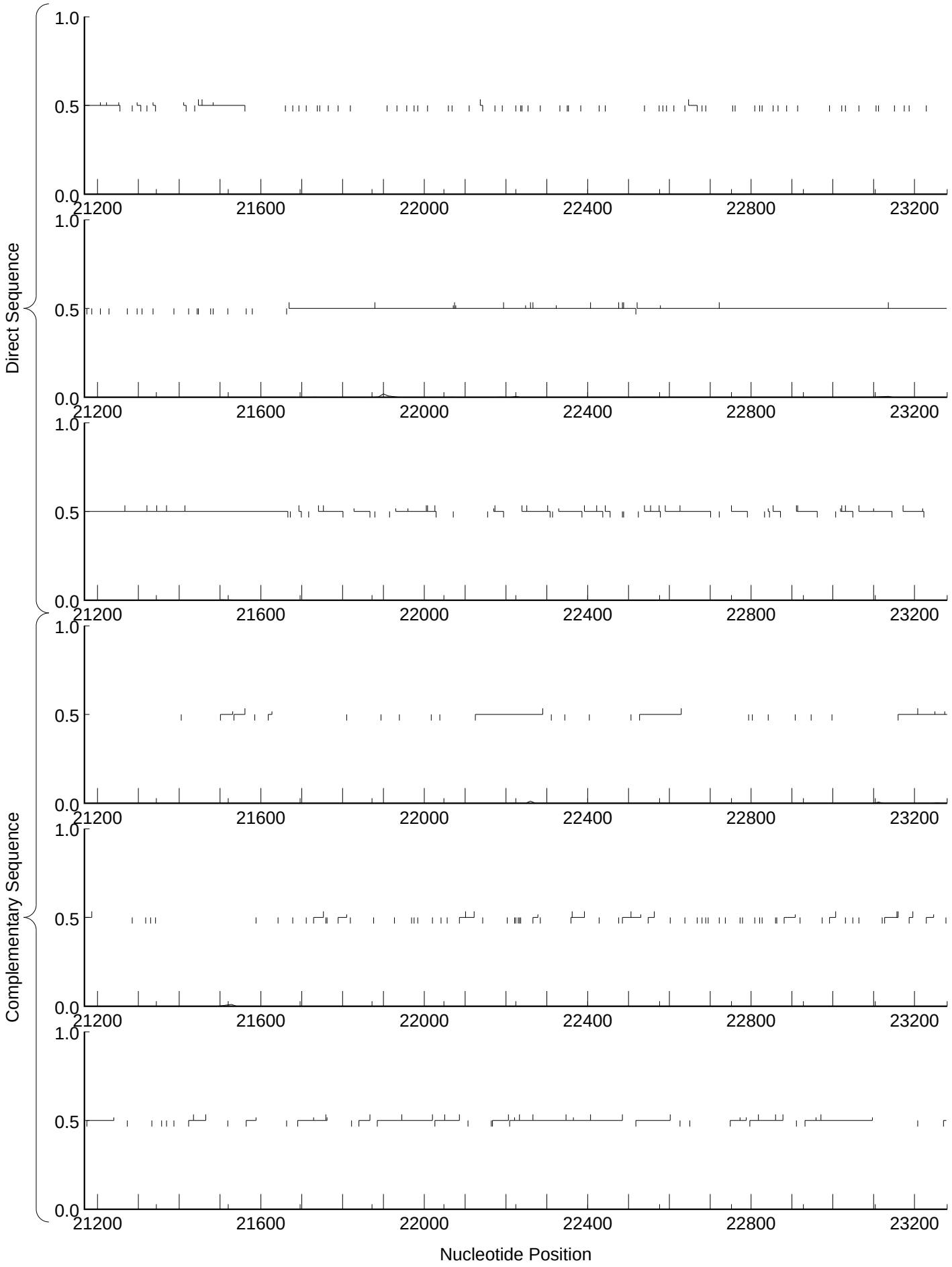
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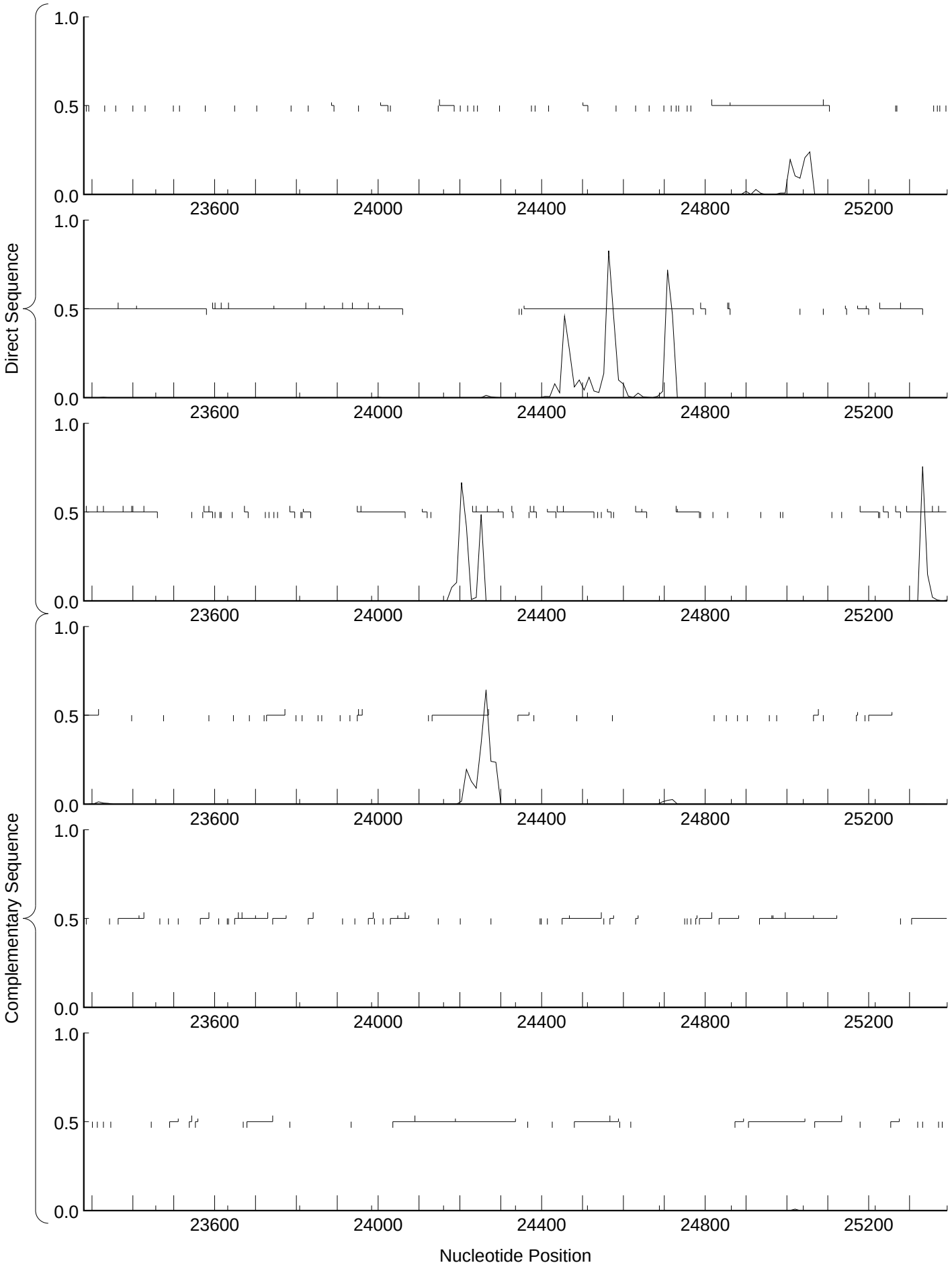
zy complete sequence, 56809 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AM, Order 4, Window 96, Step 12, 11/28



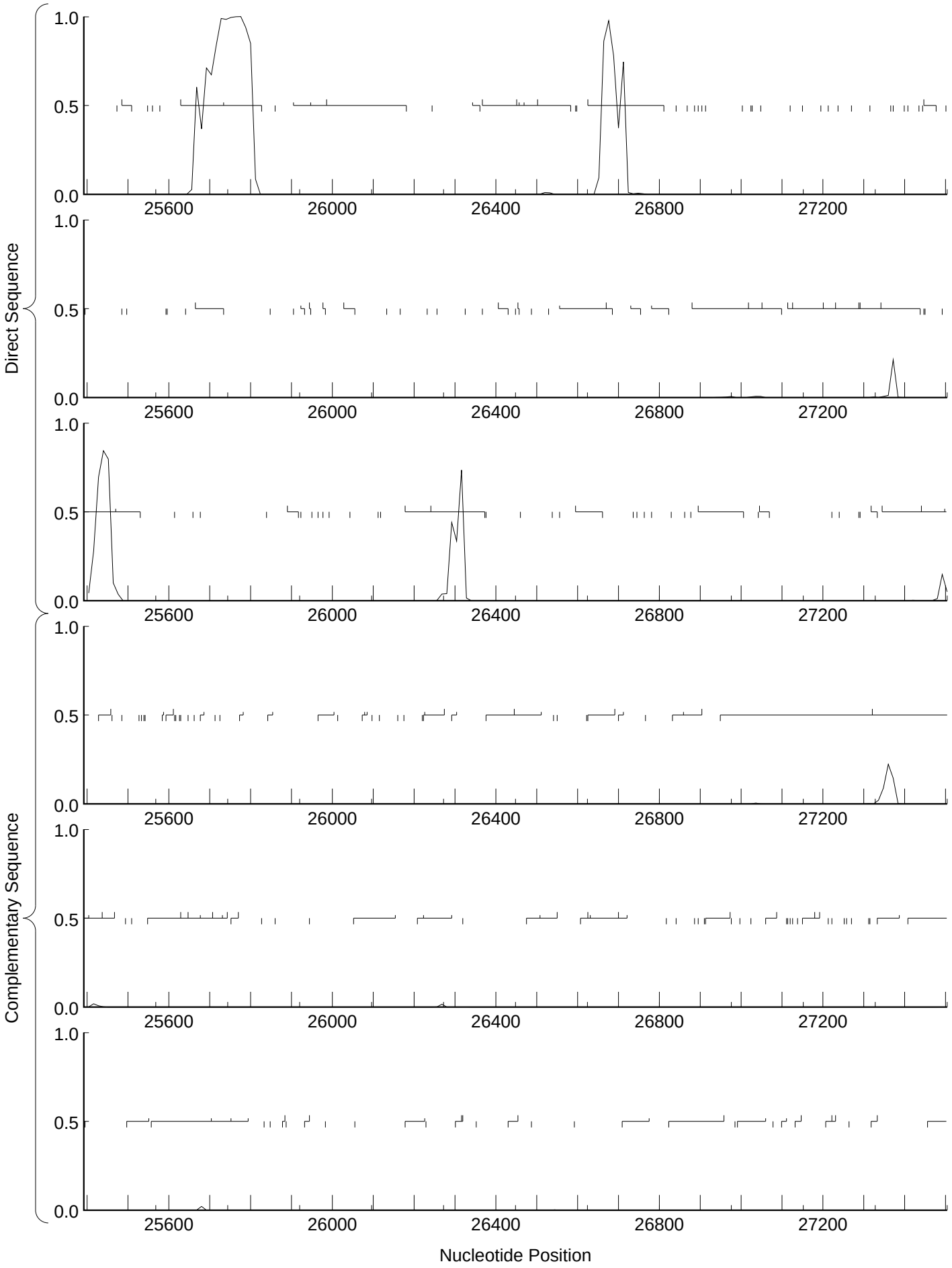
zy complete sequence, 56809 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AM, Order 4, Window 96, Step 12, 12/28



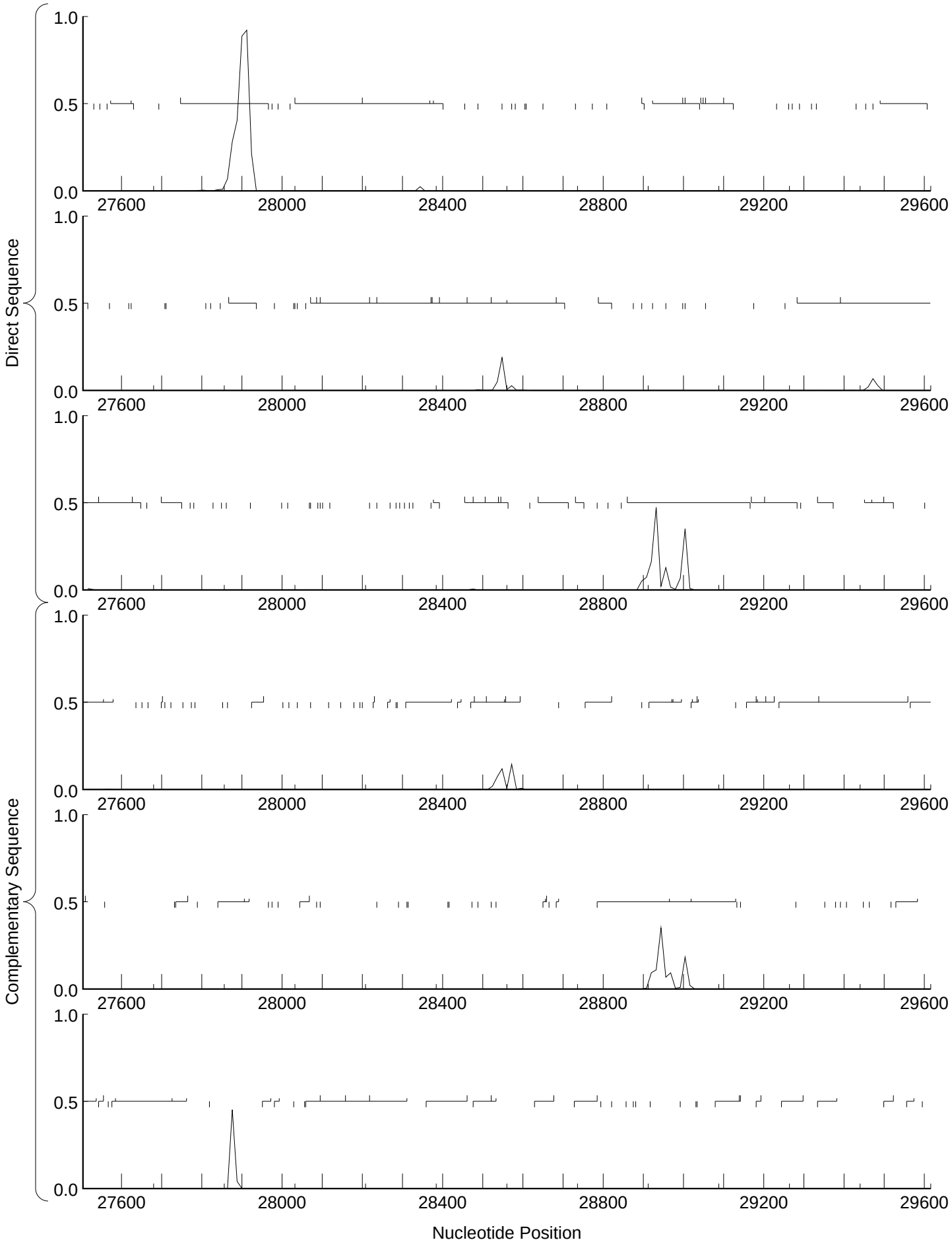
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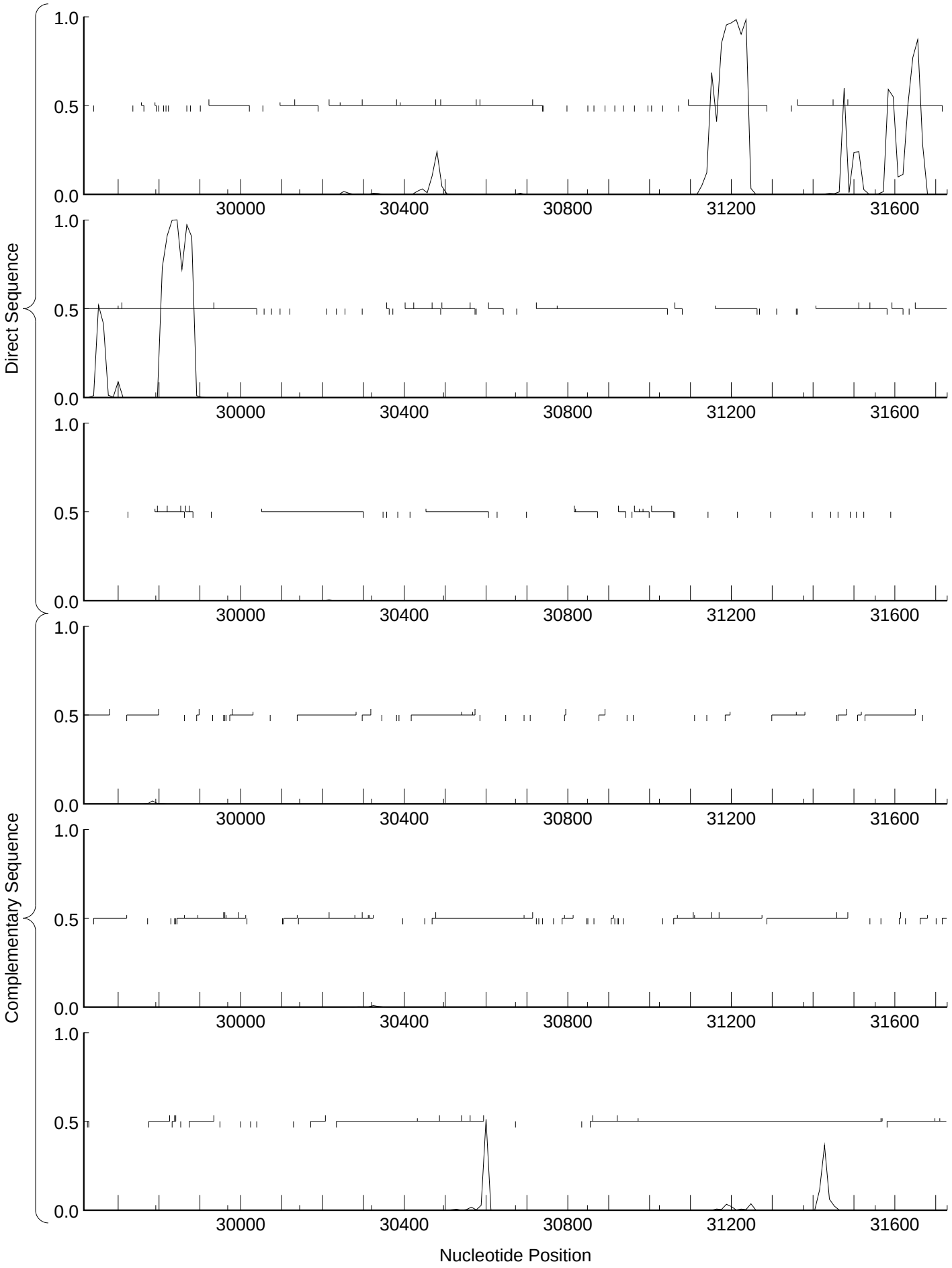


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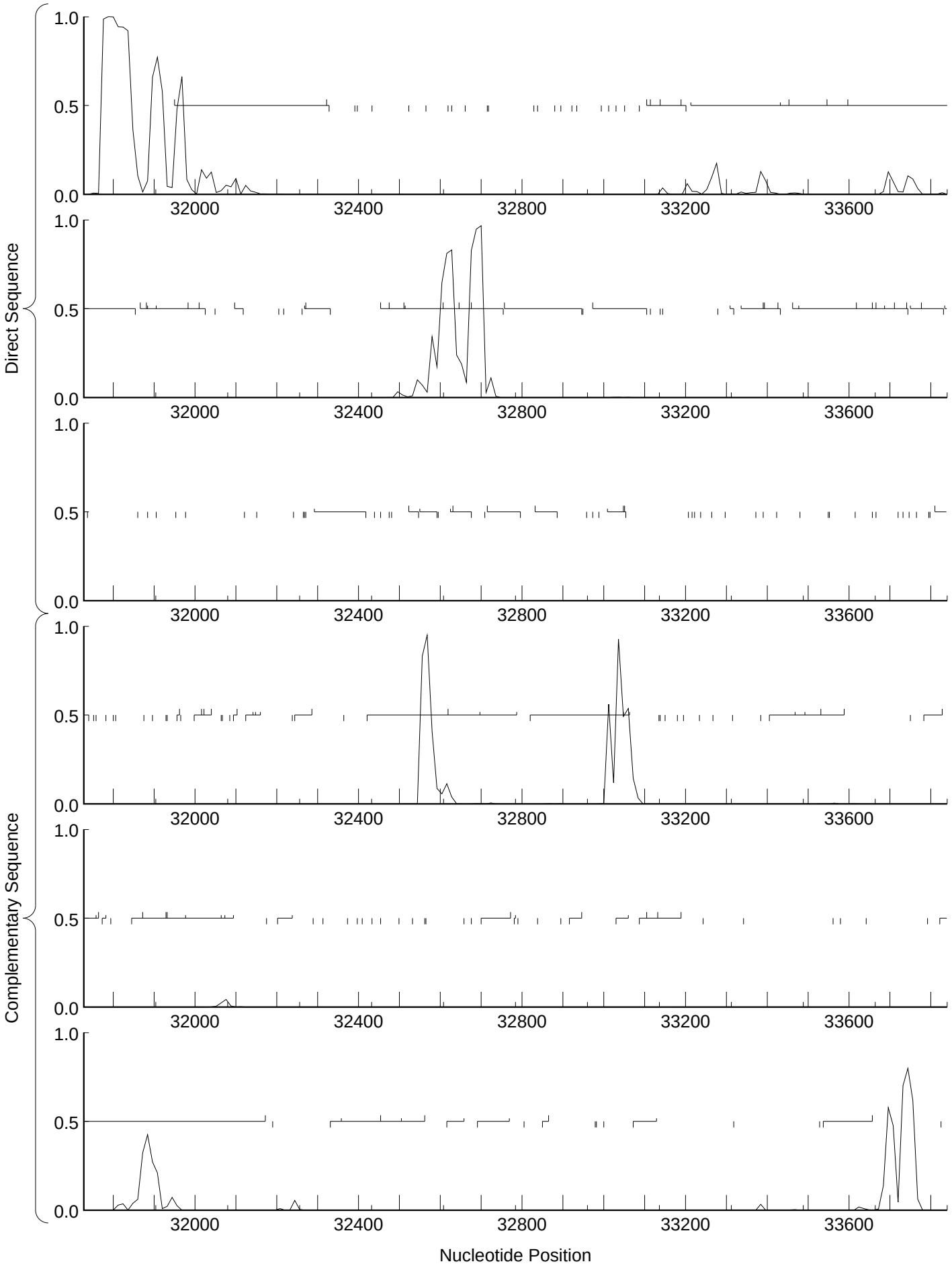


zy complete sequence, 56809 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AM, Order 4, Window 96, Step 12, 15/28

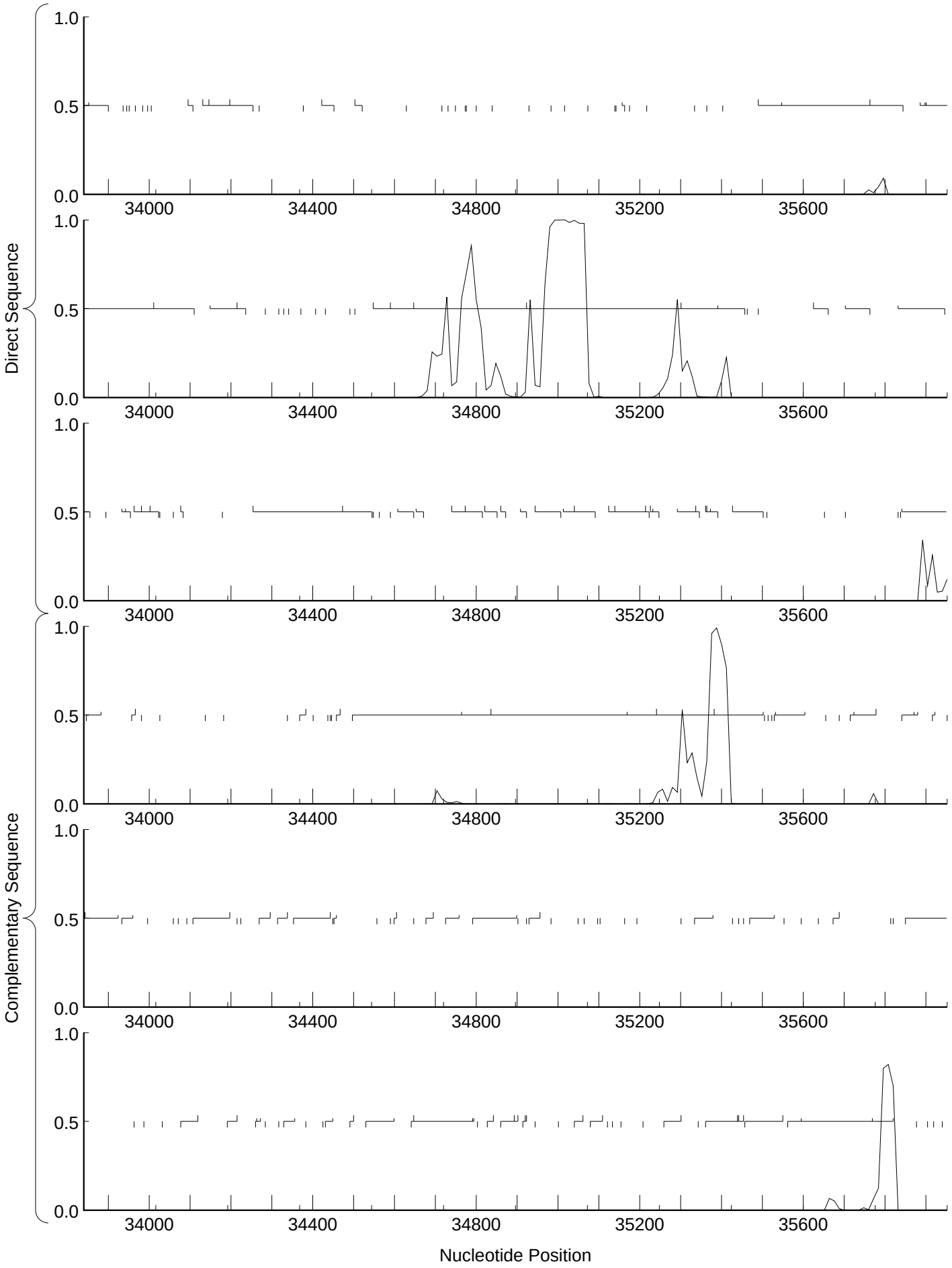



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
30000
30400
30800
31200
31600
Nucleotide Position

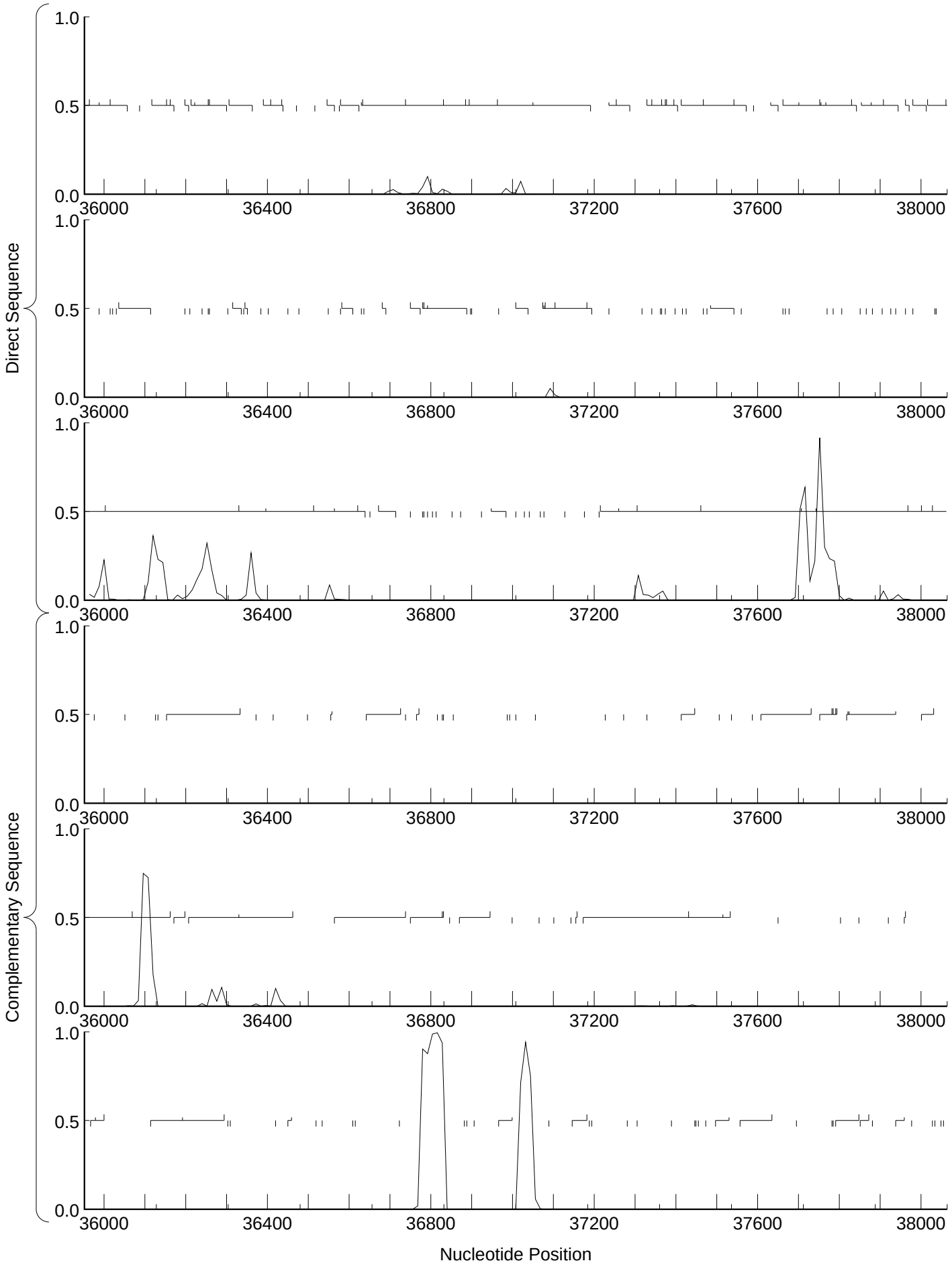
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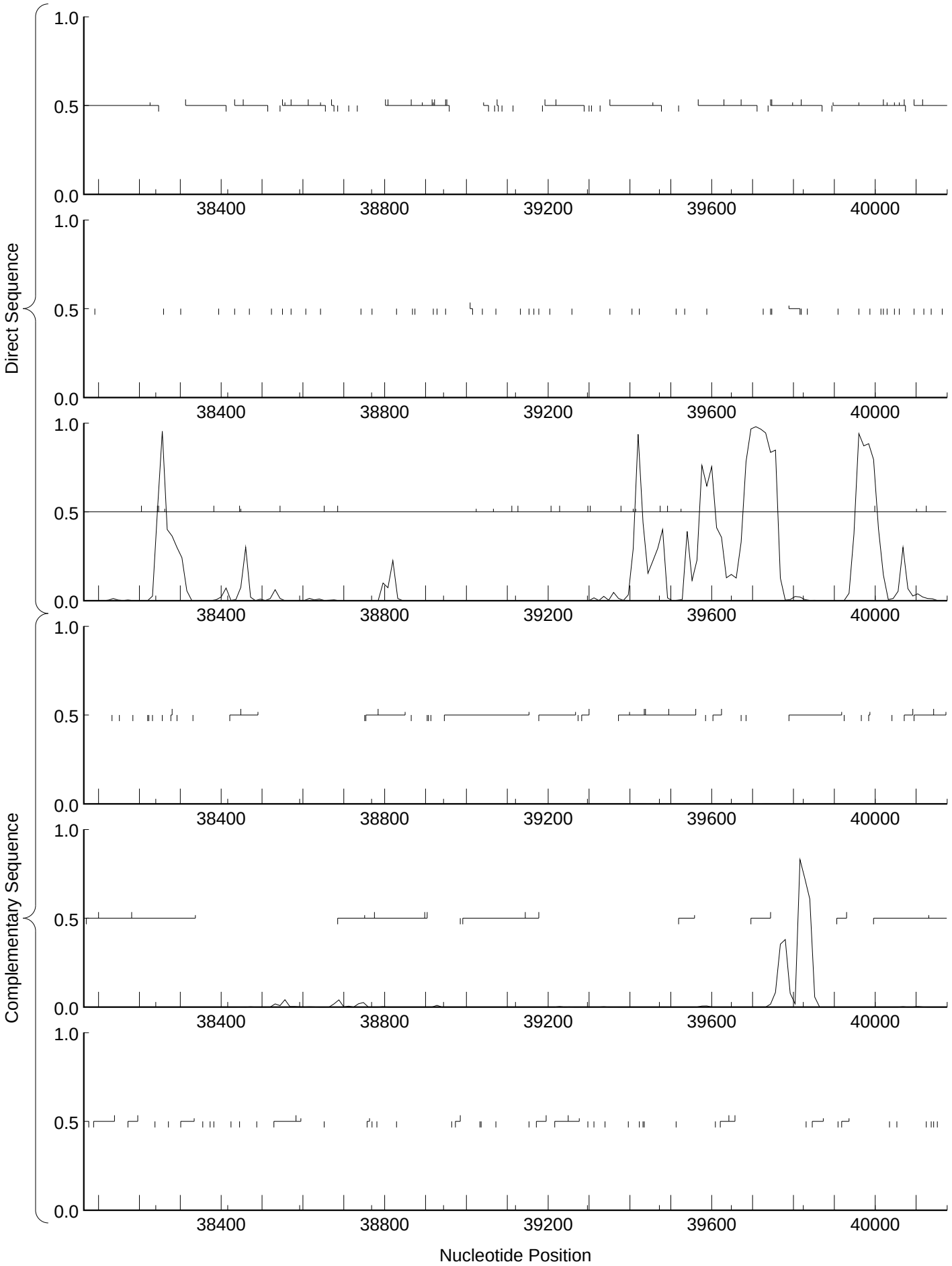
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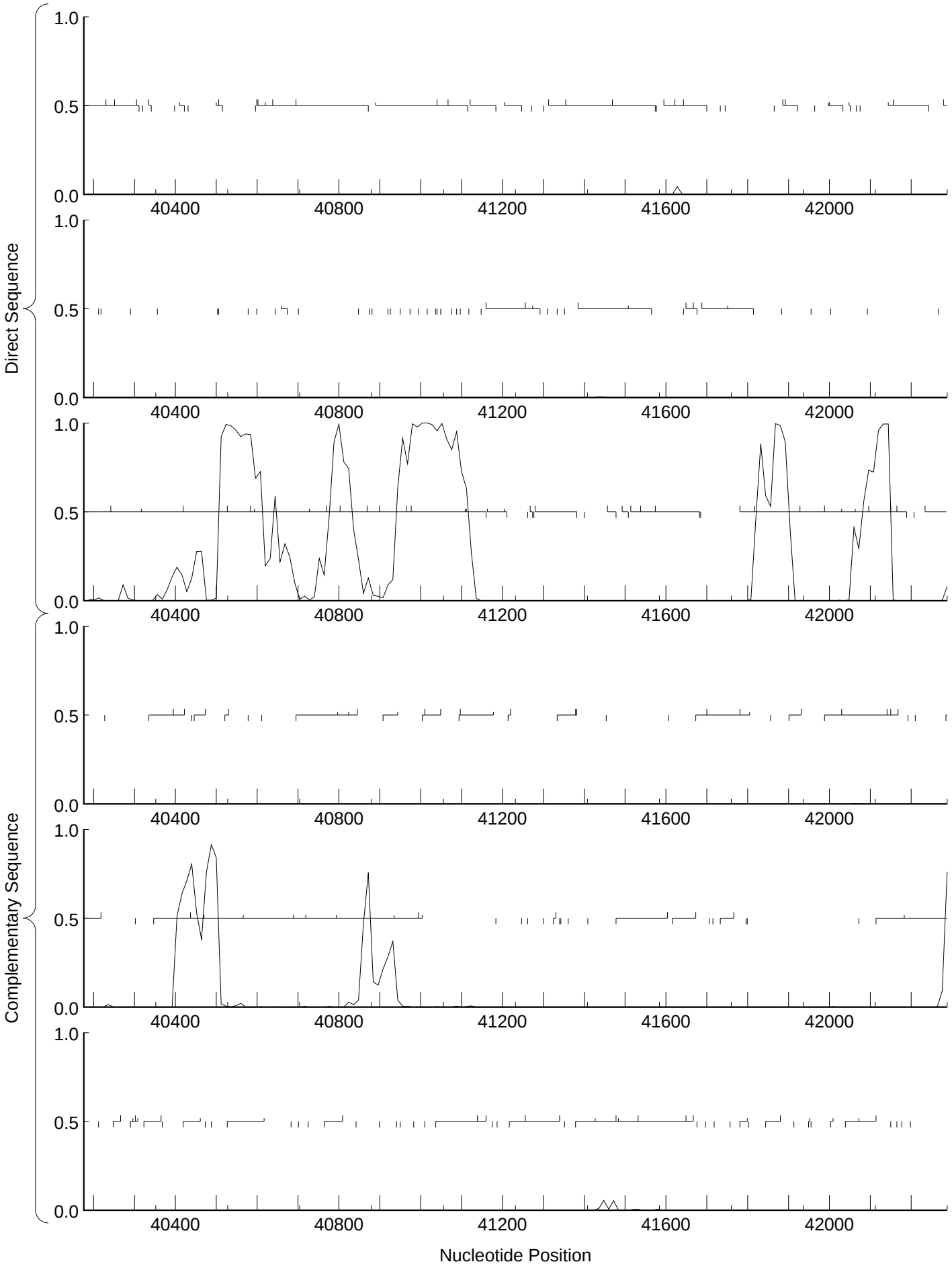
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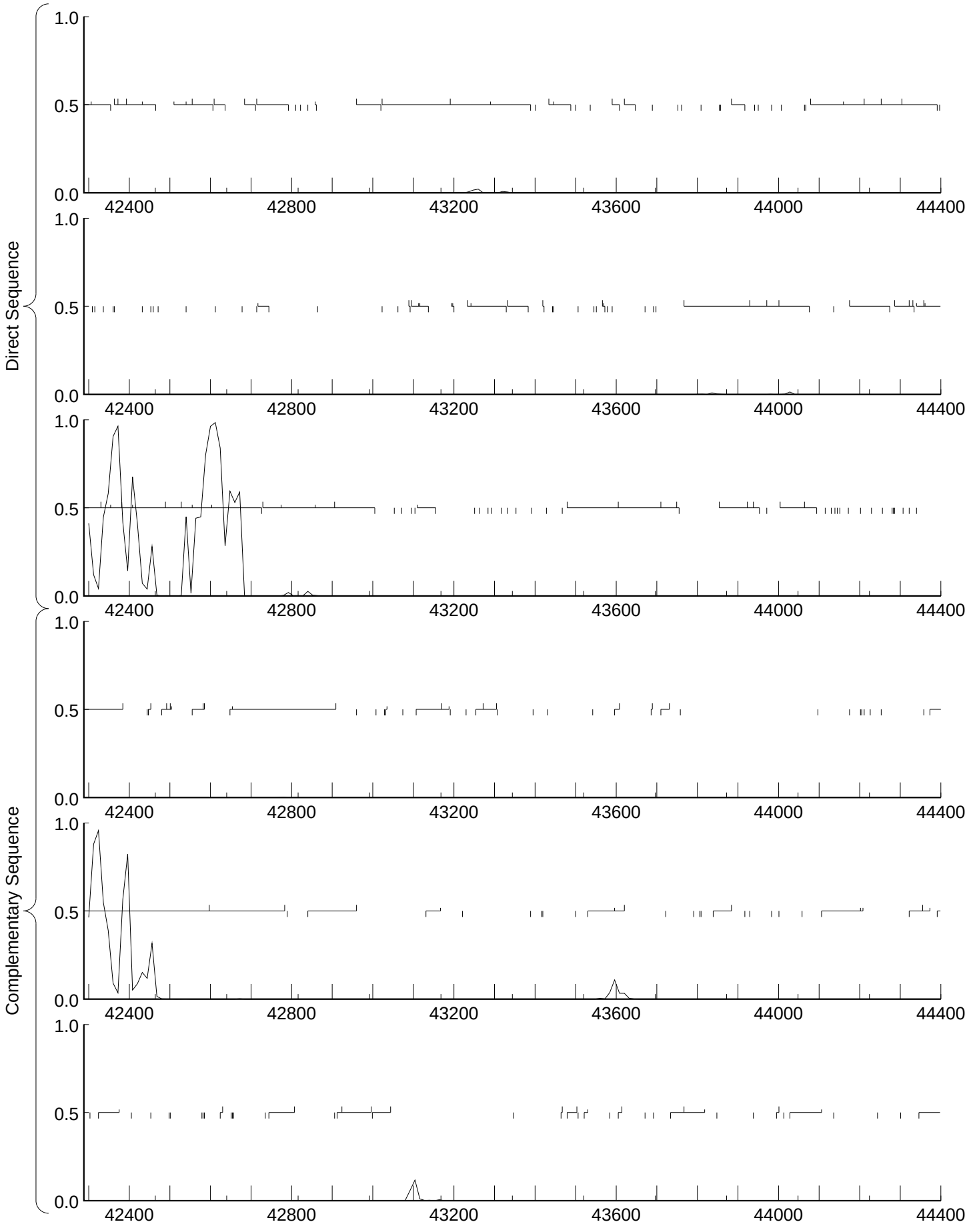


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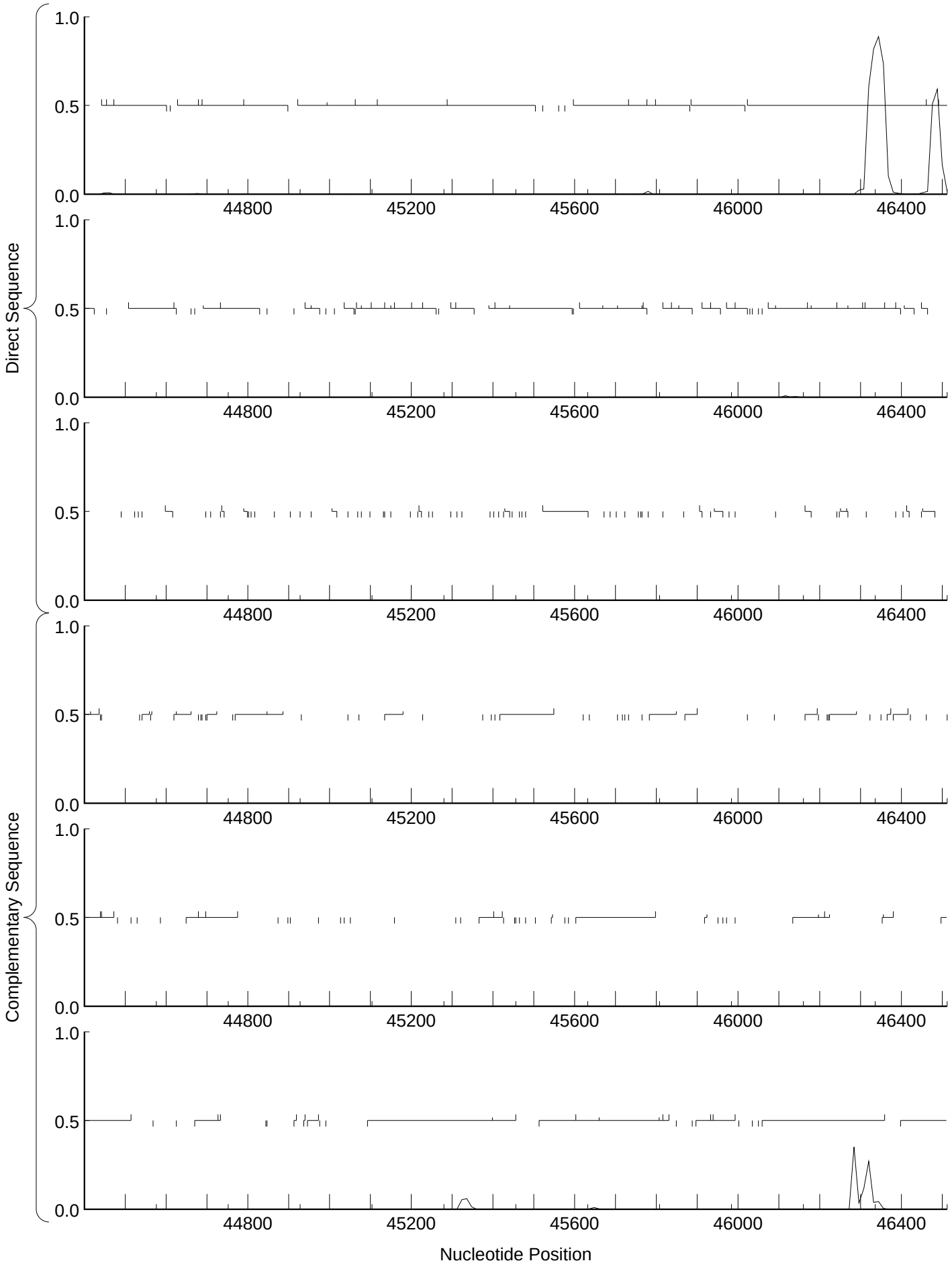


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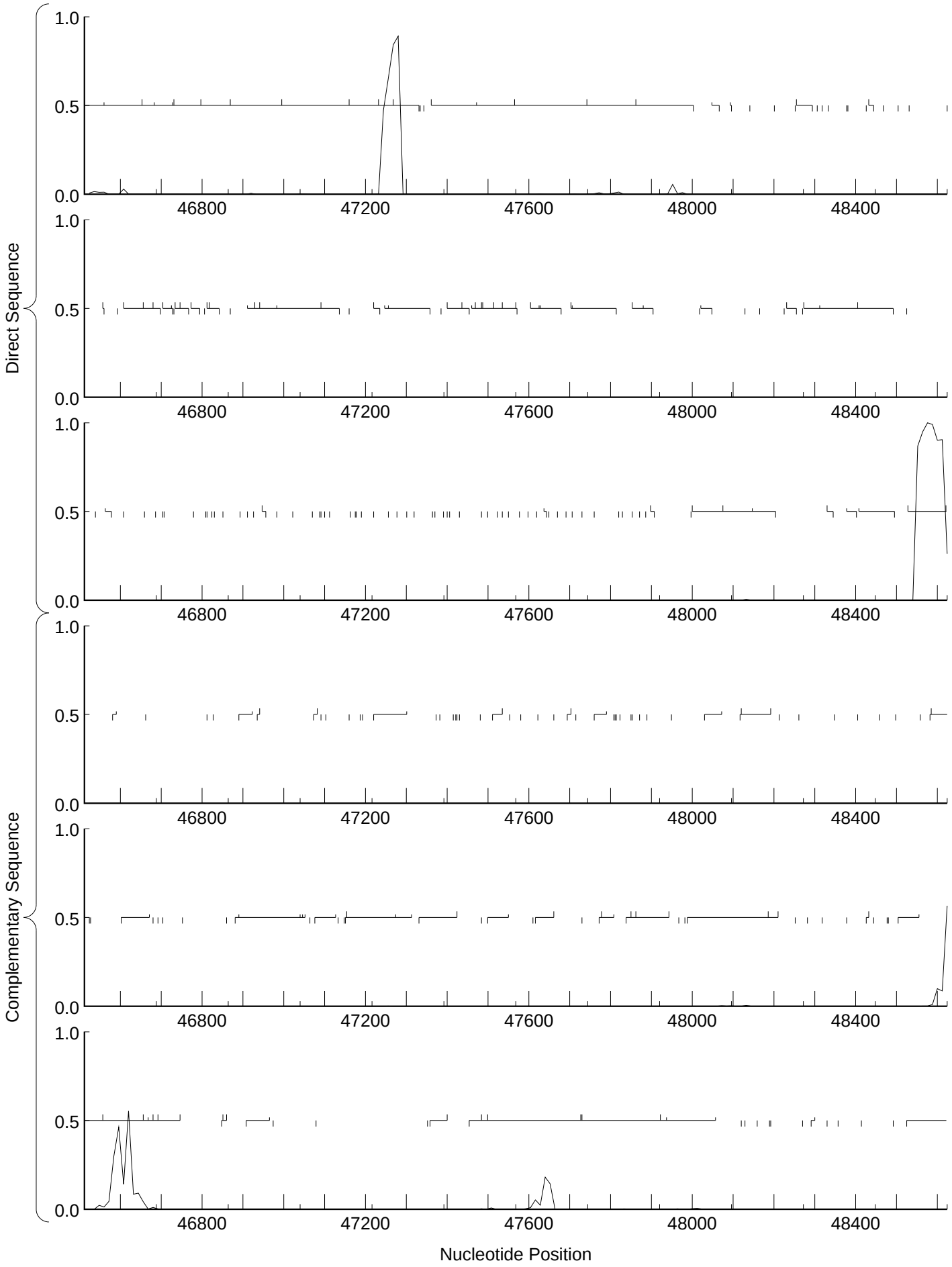



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
42400
42800
43200
43600
44000
44400
Nucleotide Position

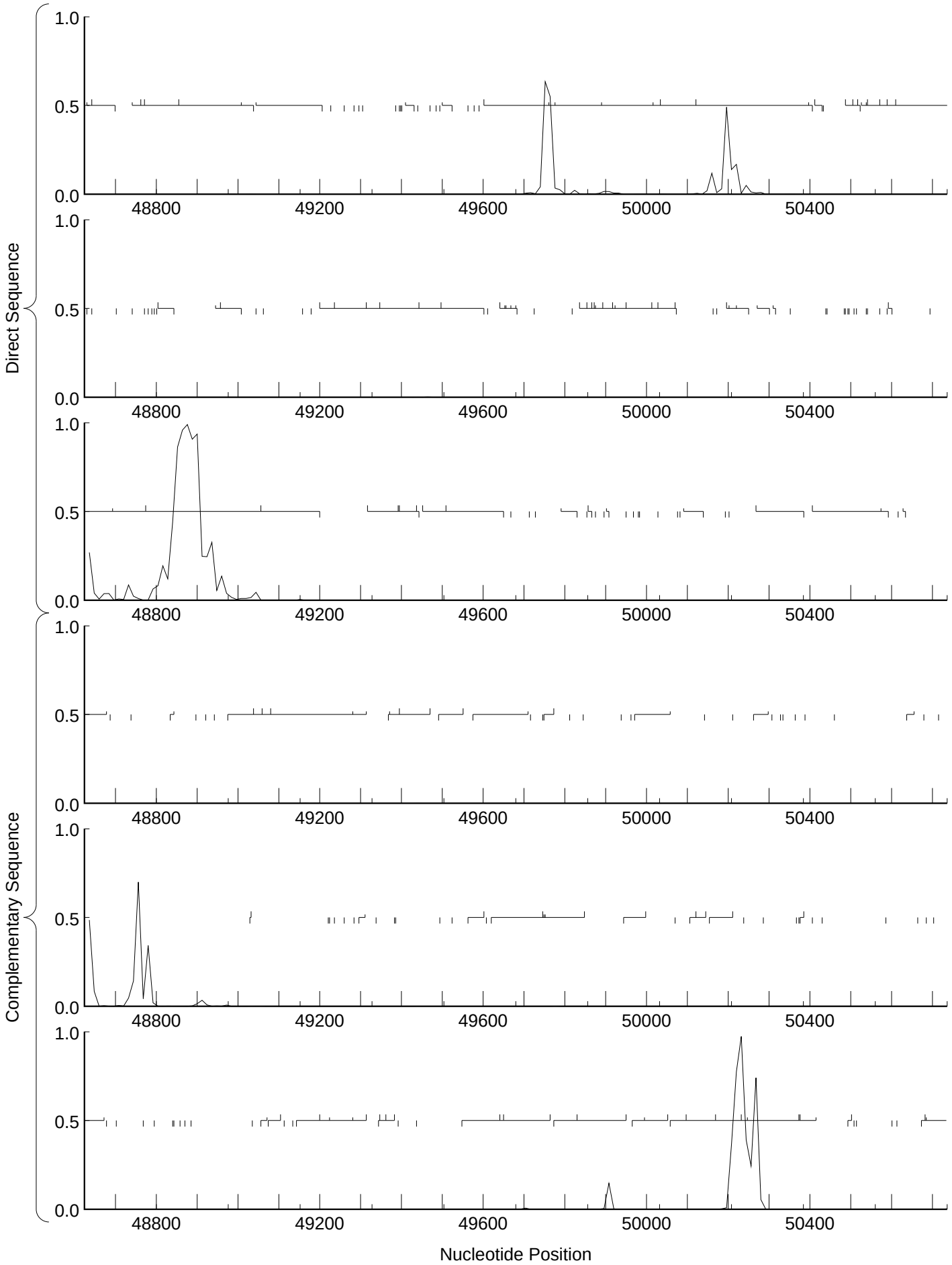
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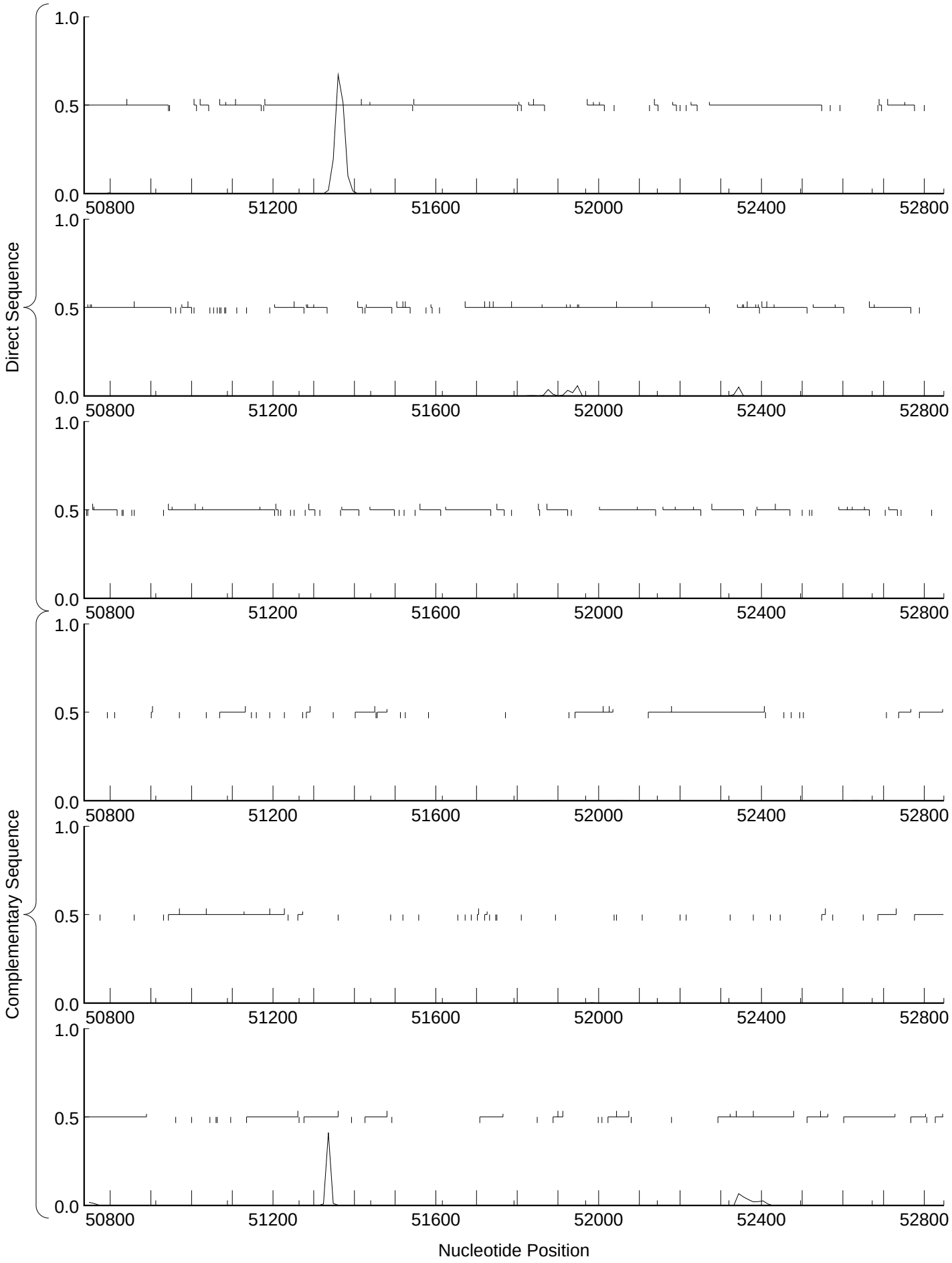


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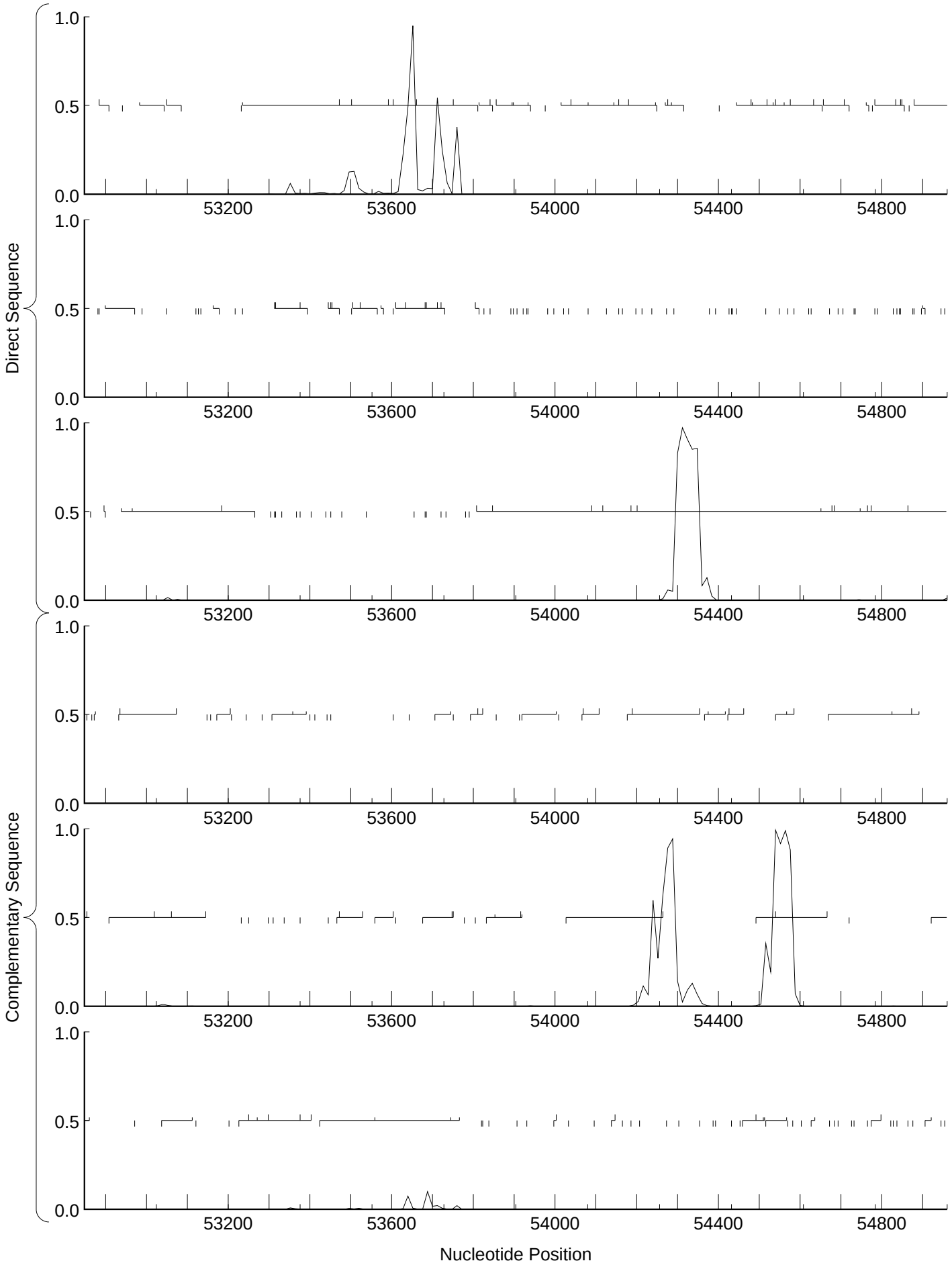


zy complete sequence, 56809 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AM, Order 4, Window 96, Step 12, 25/28




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

zy complete sequence, 56809 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AM, Order 4, Window 96, Step 12, 27/28



zy complete sequence, 56809 bp including 9-base 3’ overhang (CGCCGGCCT), Cluster AM, Order 4, Window 96, Step 12, 28/28

