GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Easy2Say complete sequence, 75598 bp including 9-base 3’ overhang (CGCT
Analysis Date    :  3/28/24 at 19:14:01
Pages            :  37
Sequence Length  :  75598 bp
GC Content       :   62.92%

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

PS-Version       :  1.2

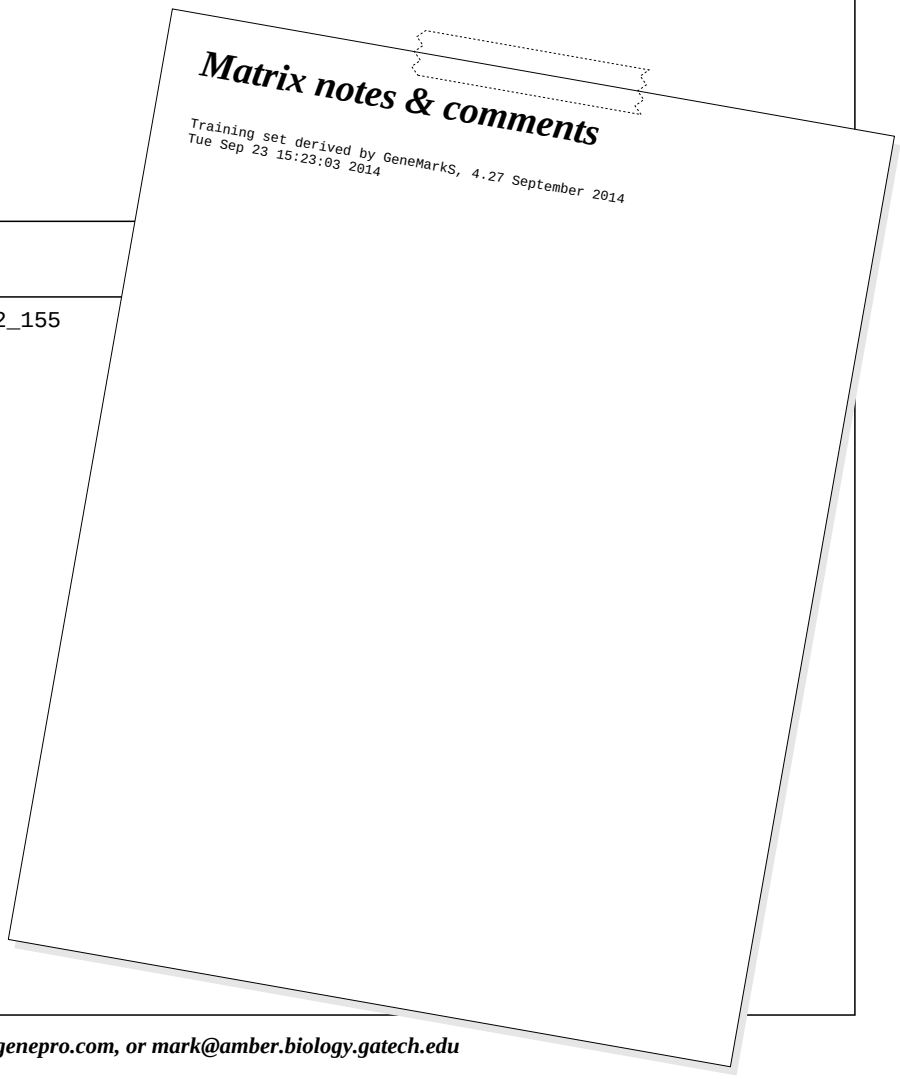
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Author     :  -

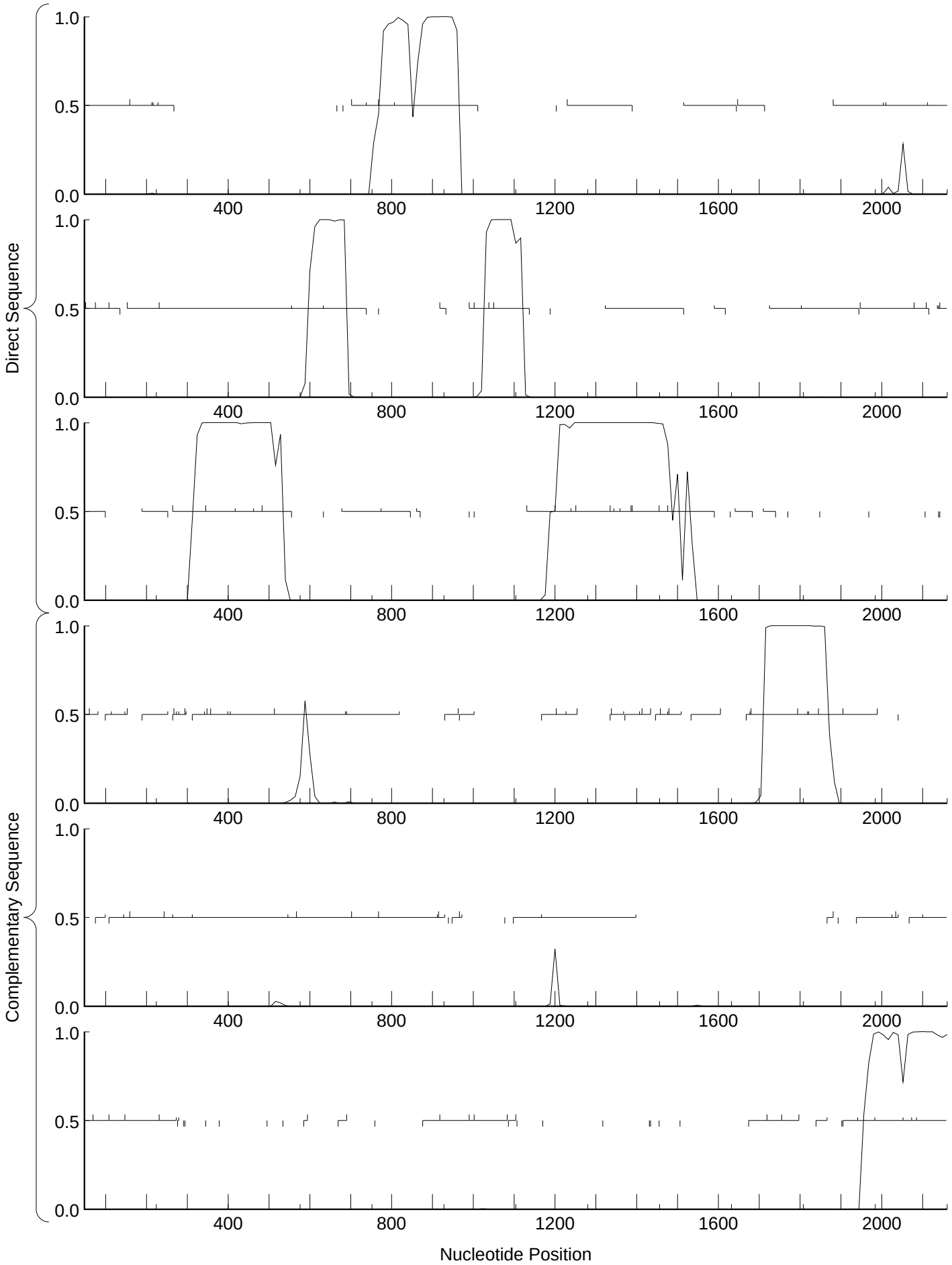
Order      :  4


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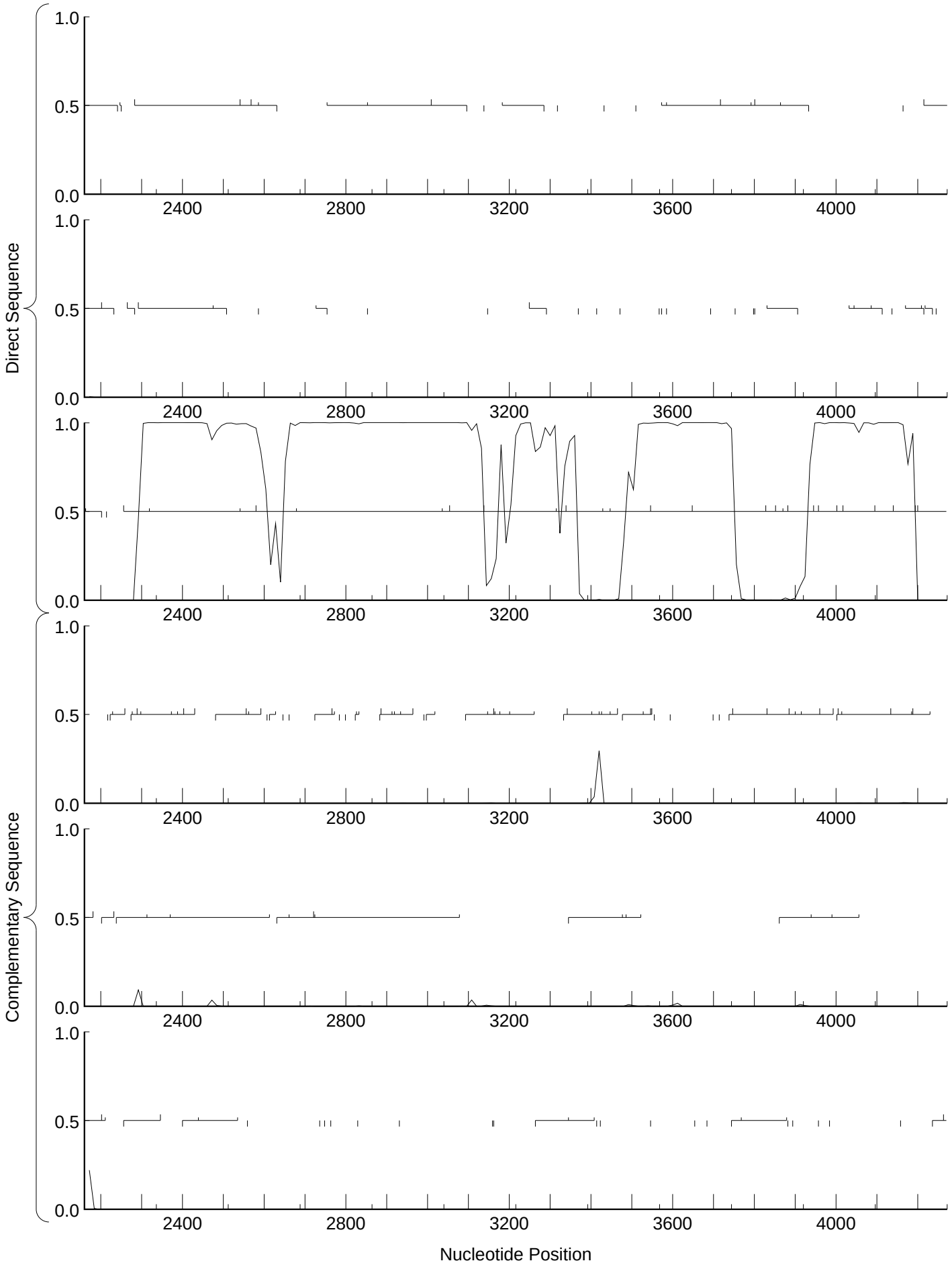


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

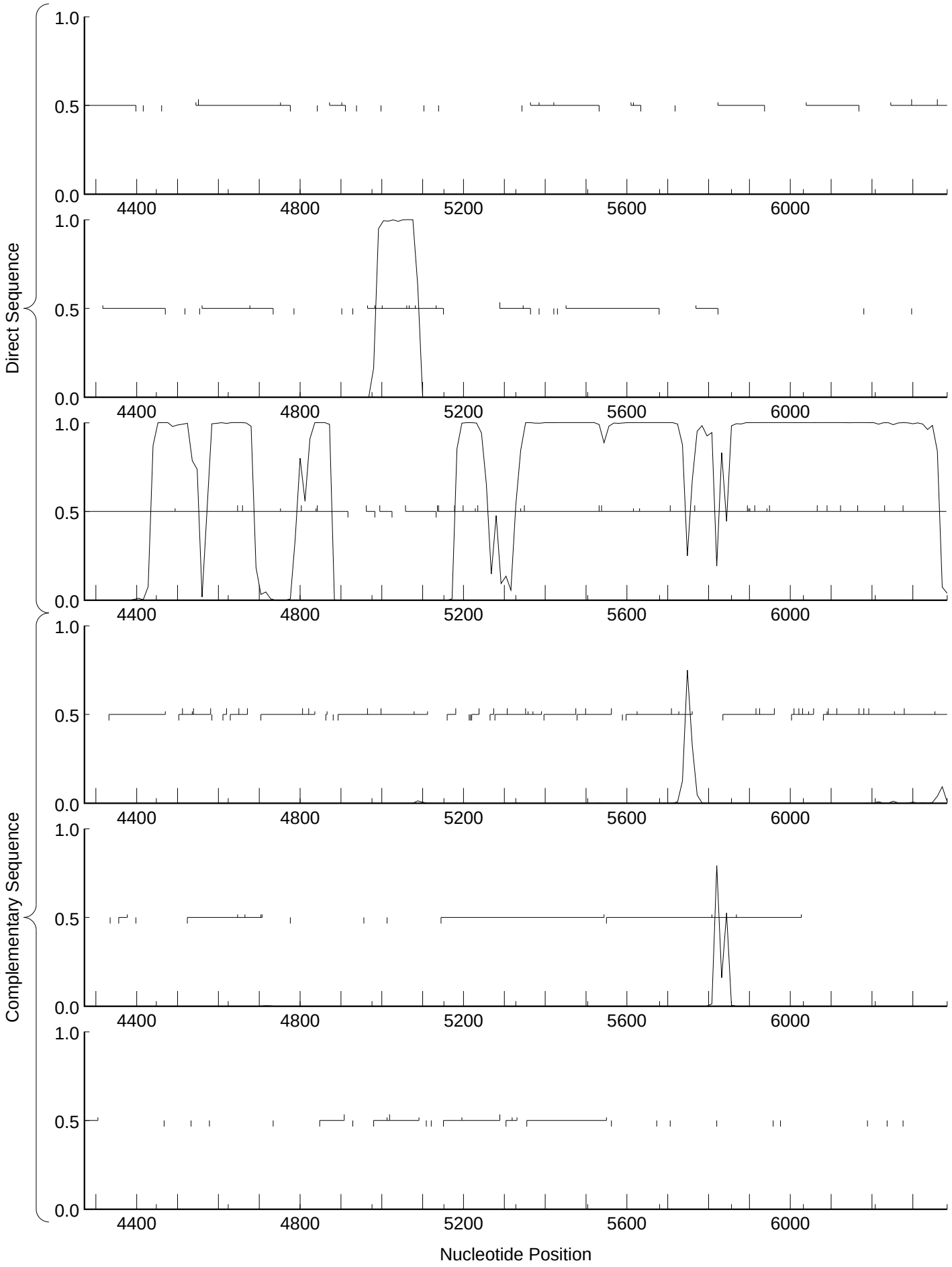
y2Say complete sequence, 75598 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 2/37



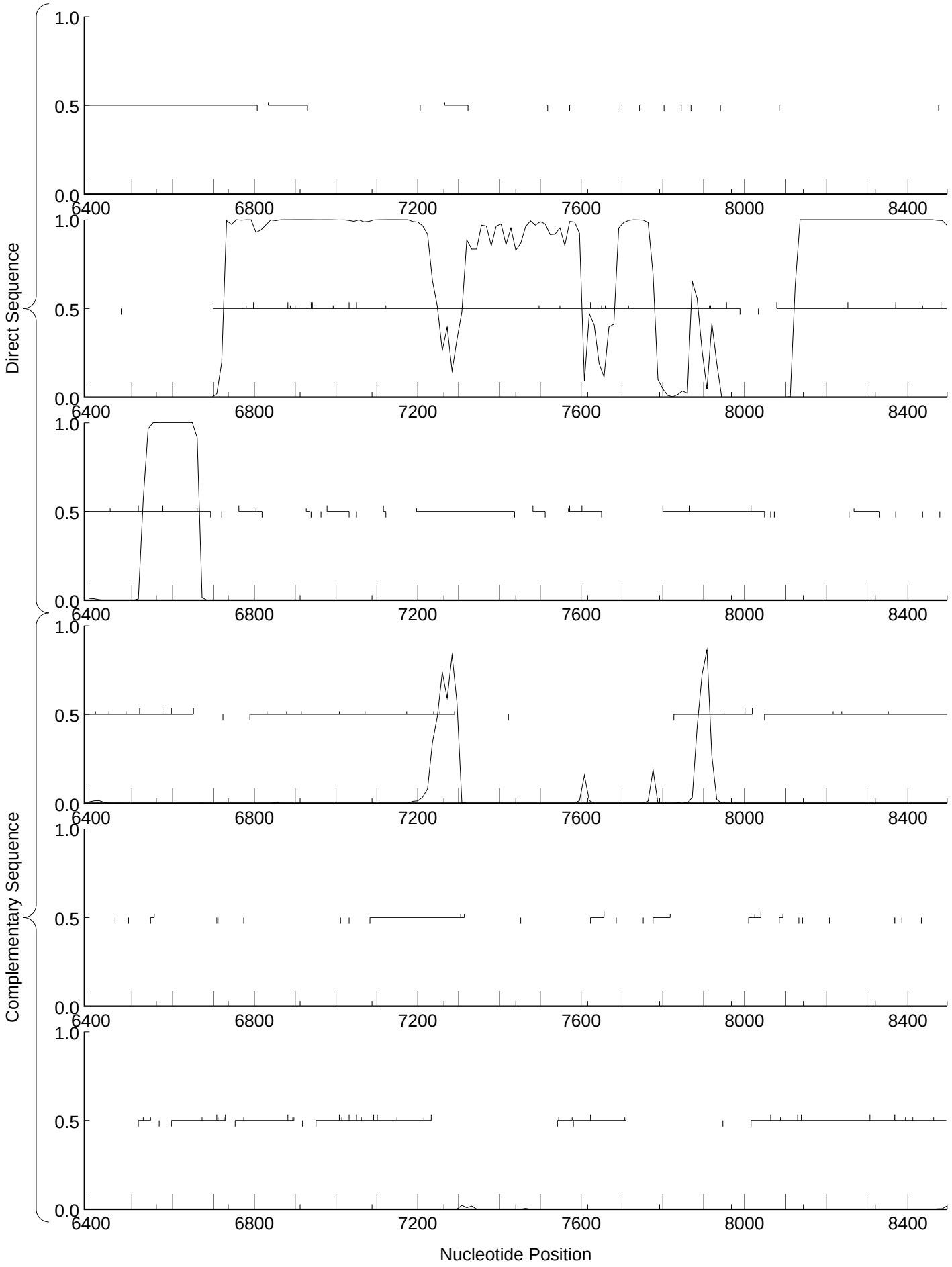
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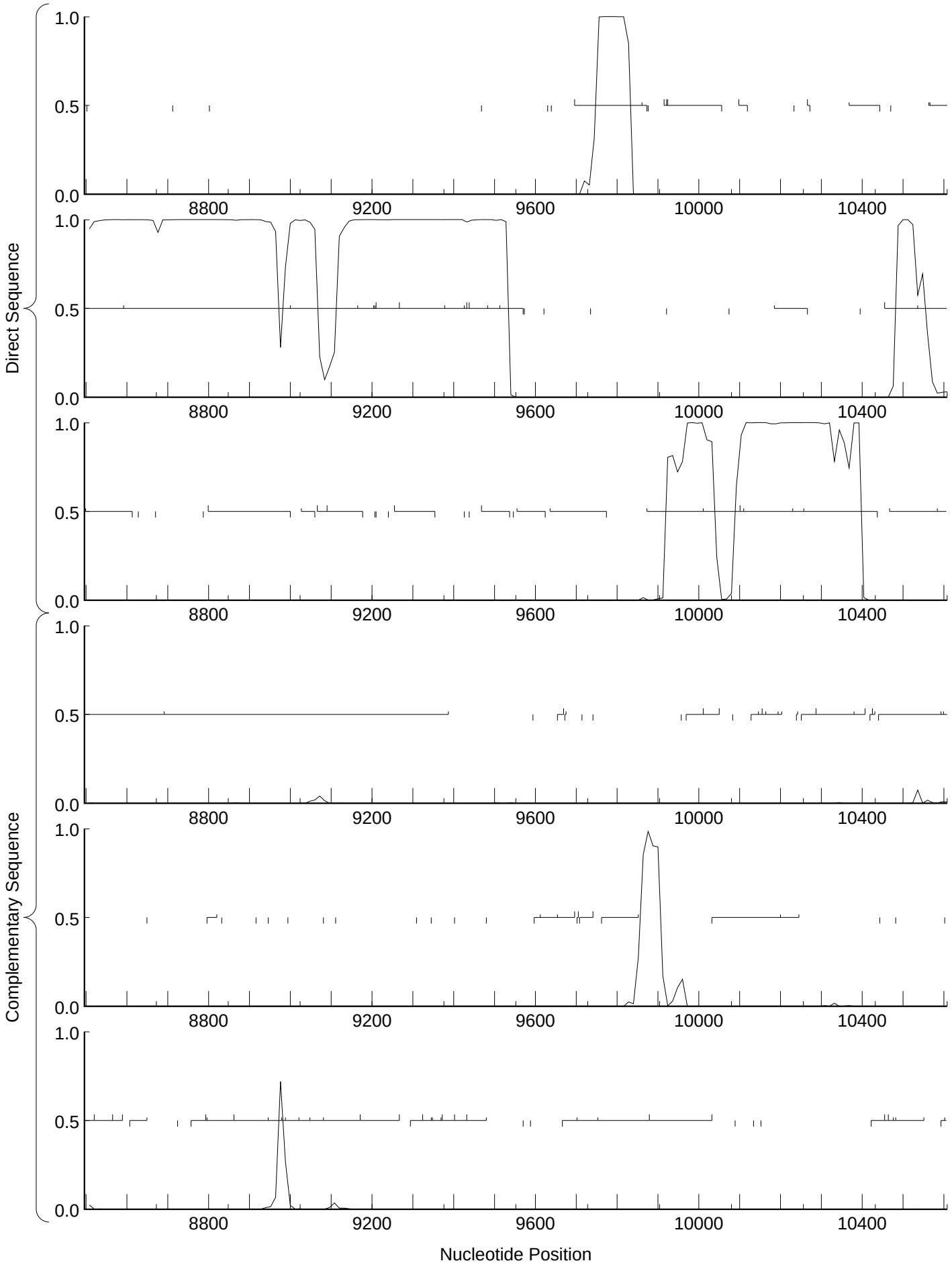
y2Say complete sequence, 75598 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 4/37



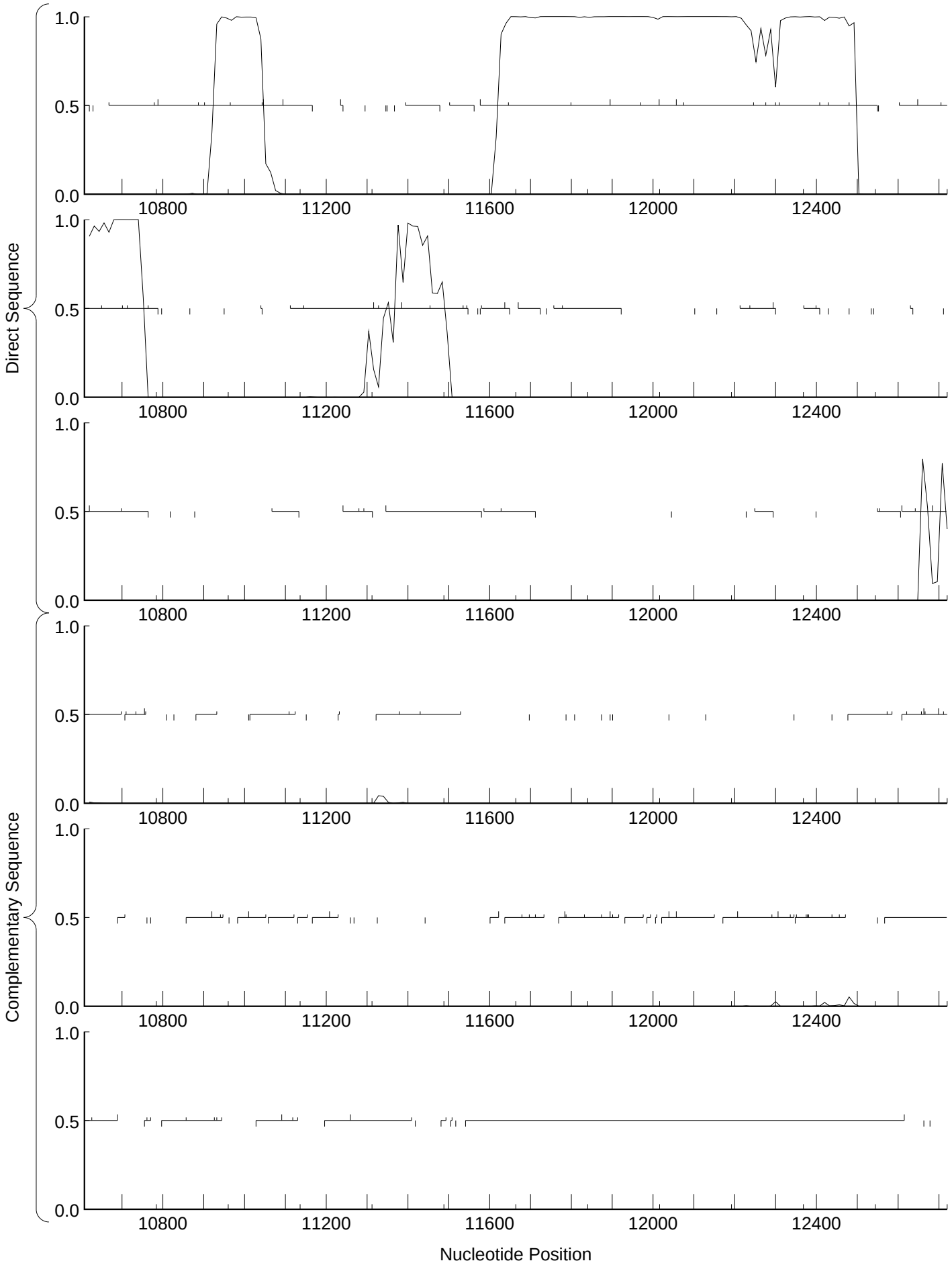
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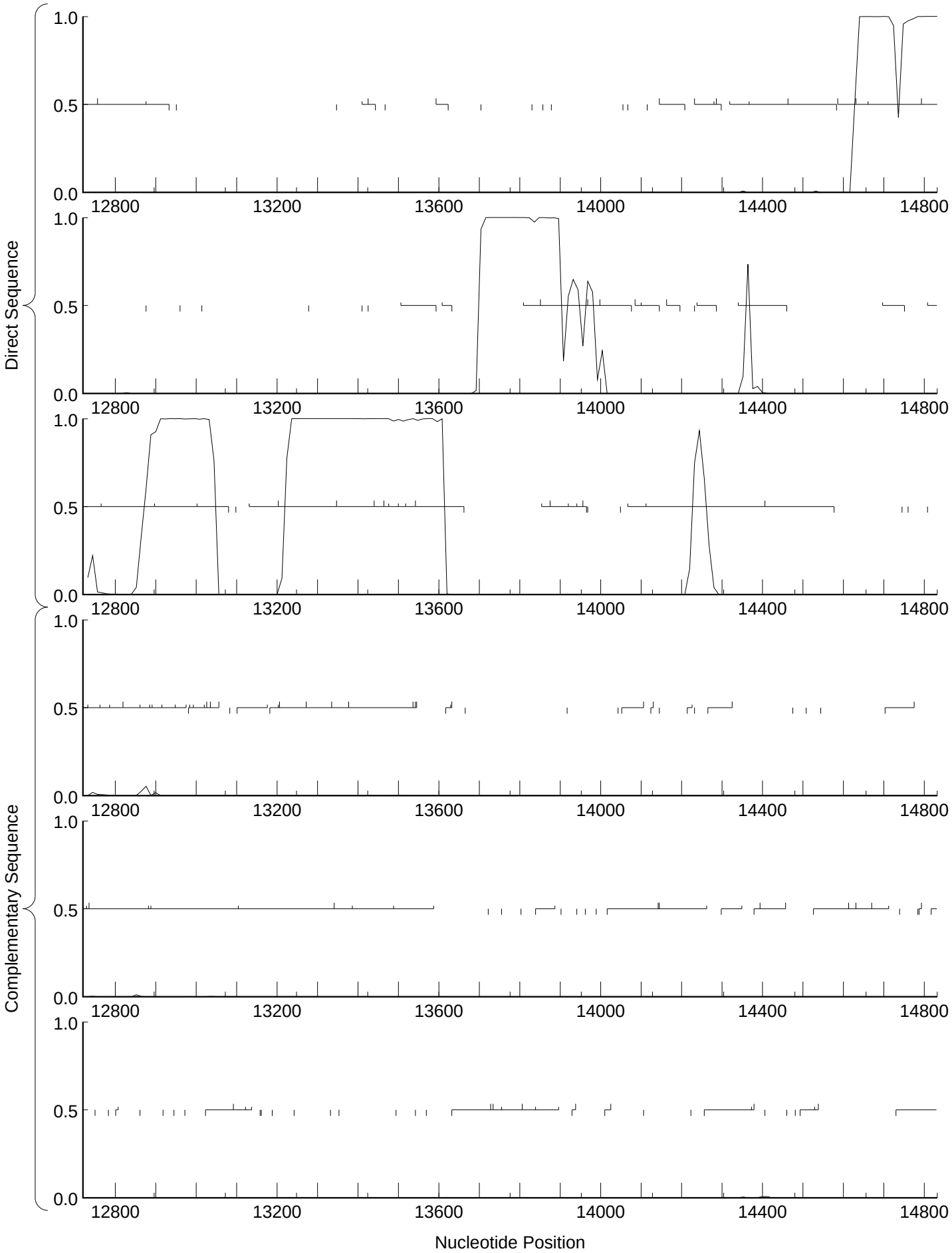
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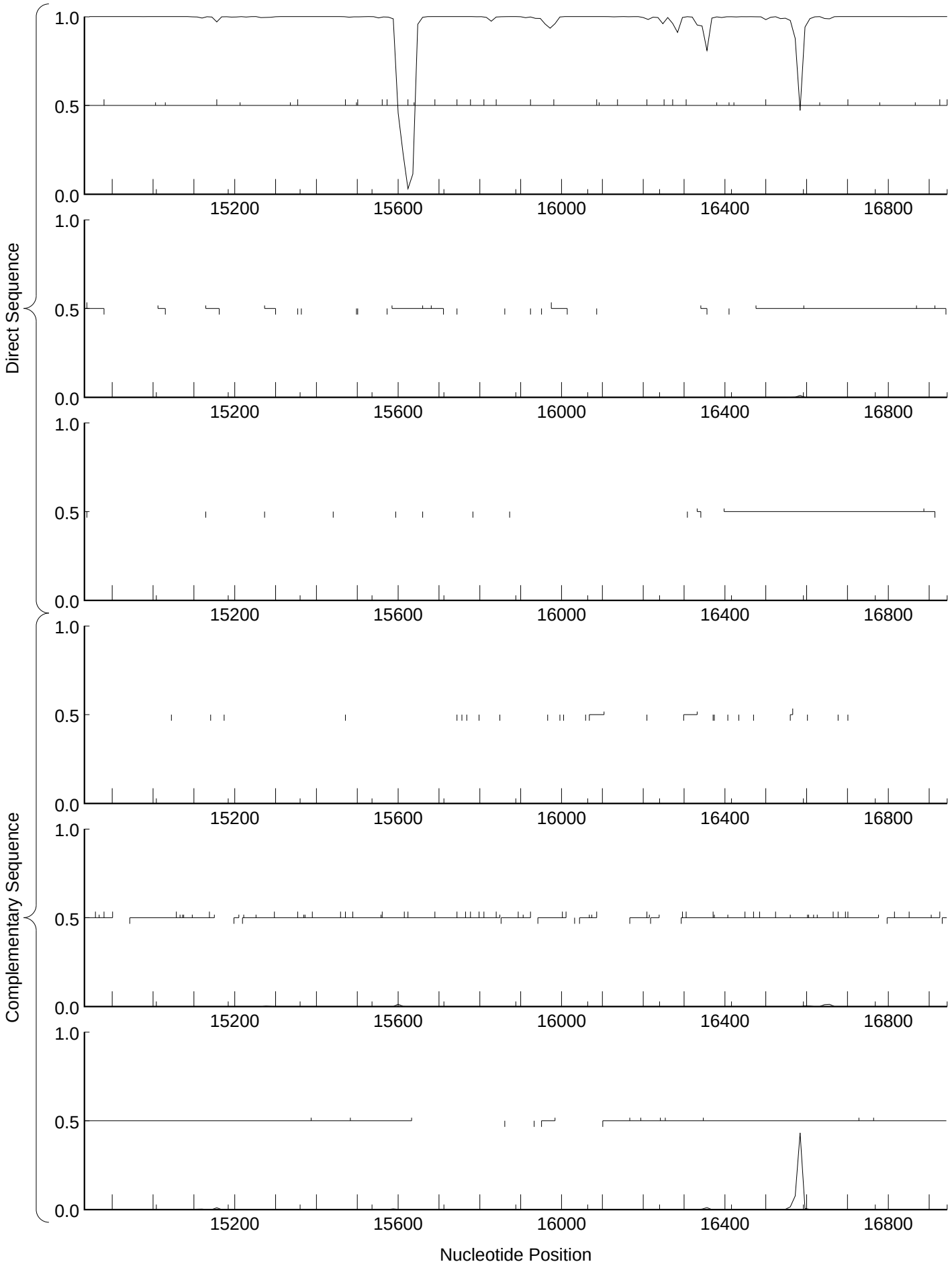
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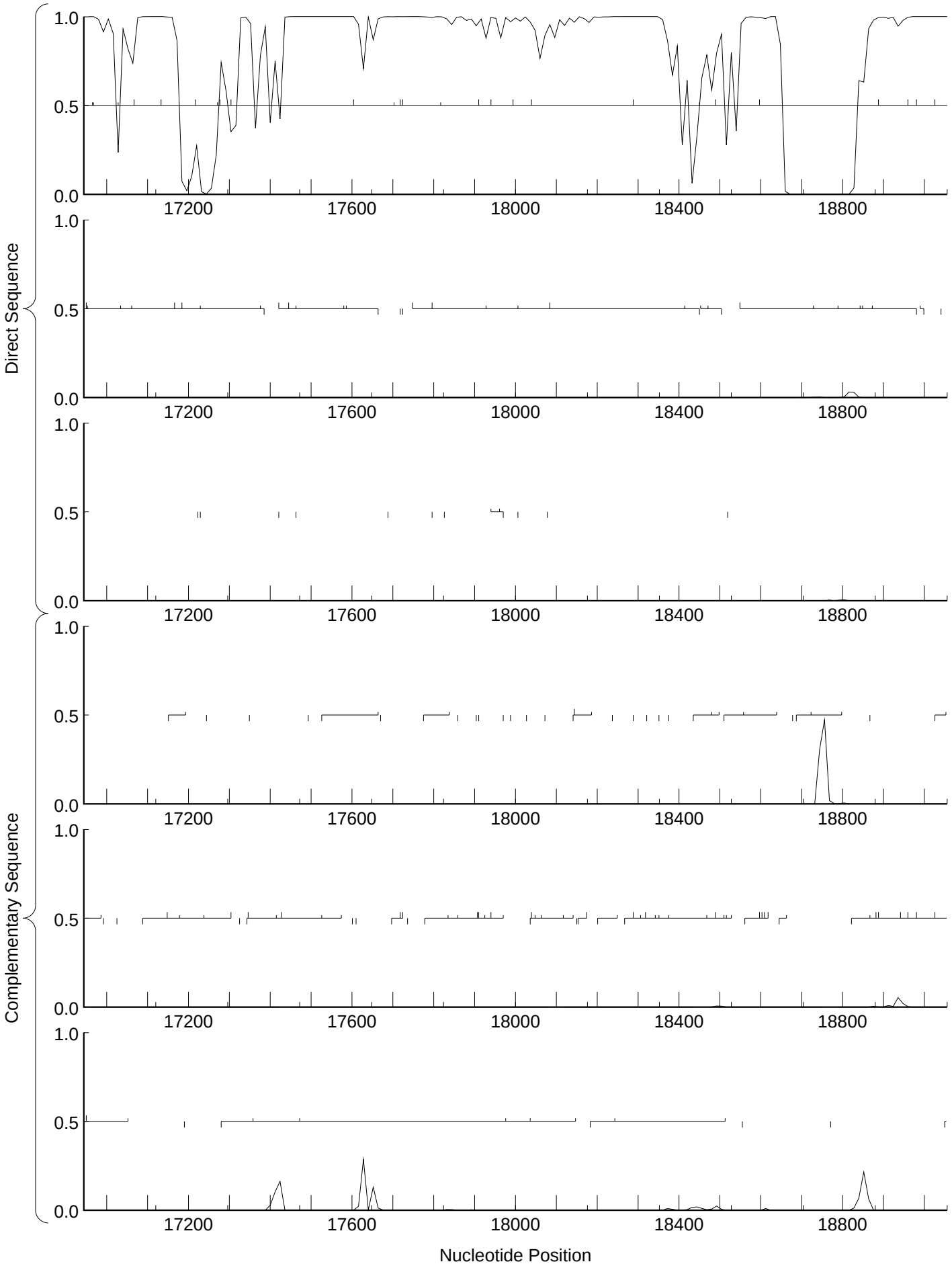


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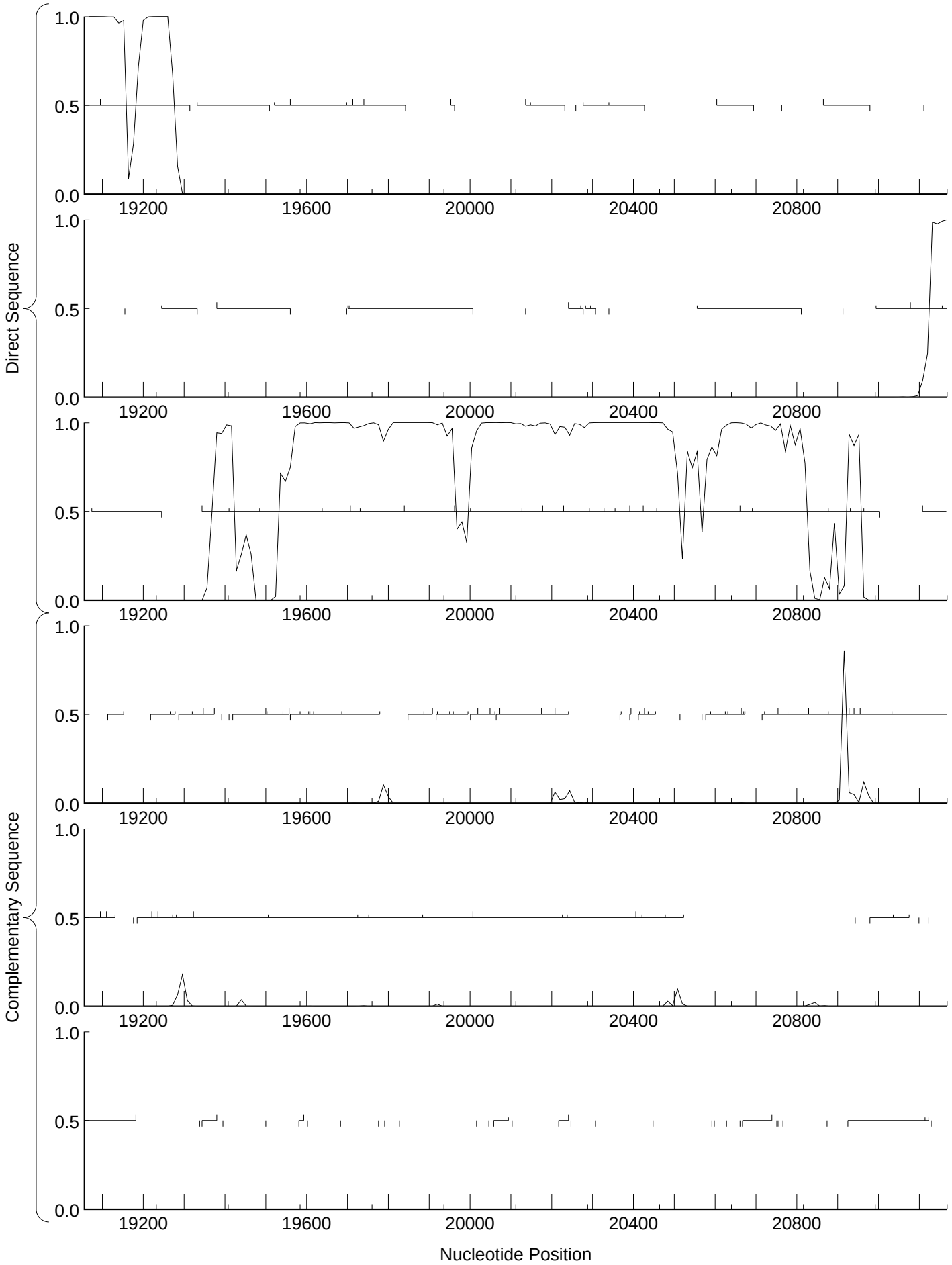


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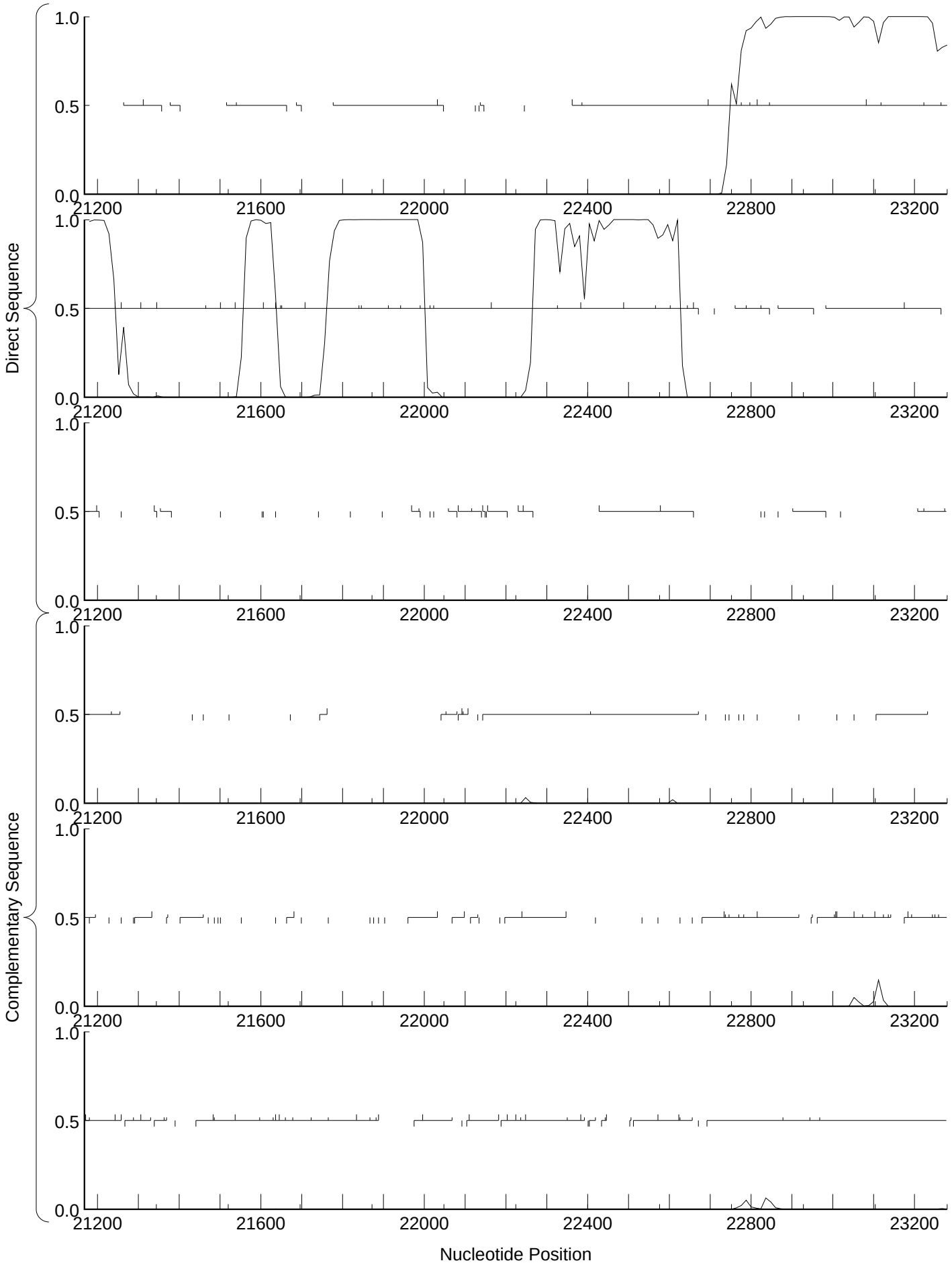


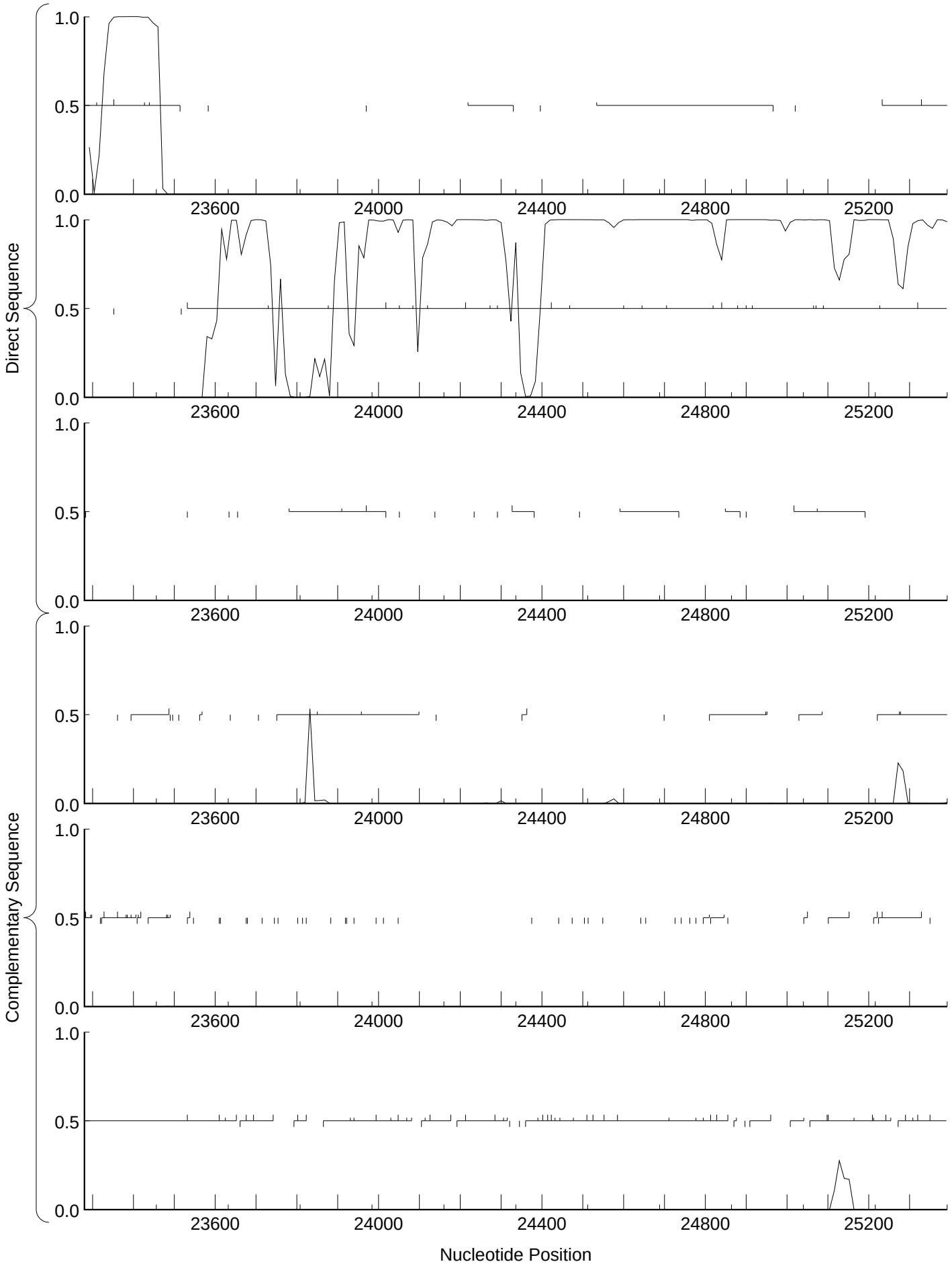

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Complementary Sequence
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Nucleotide Position

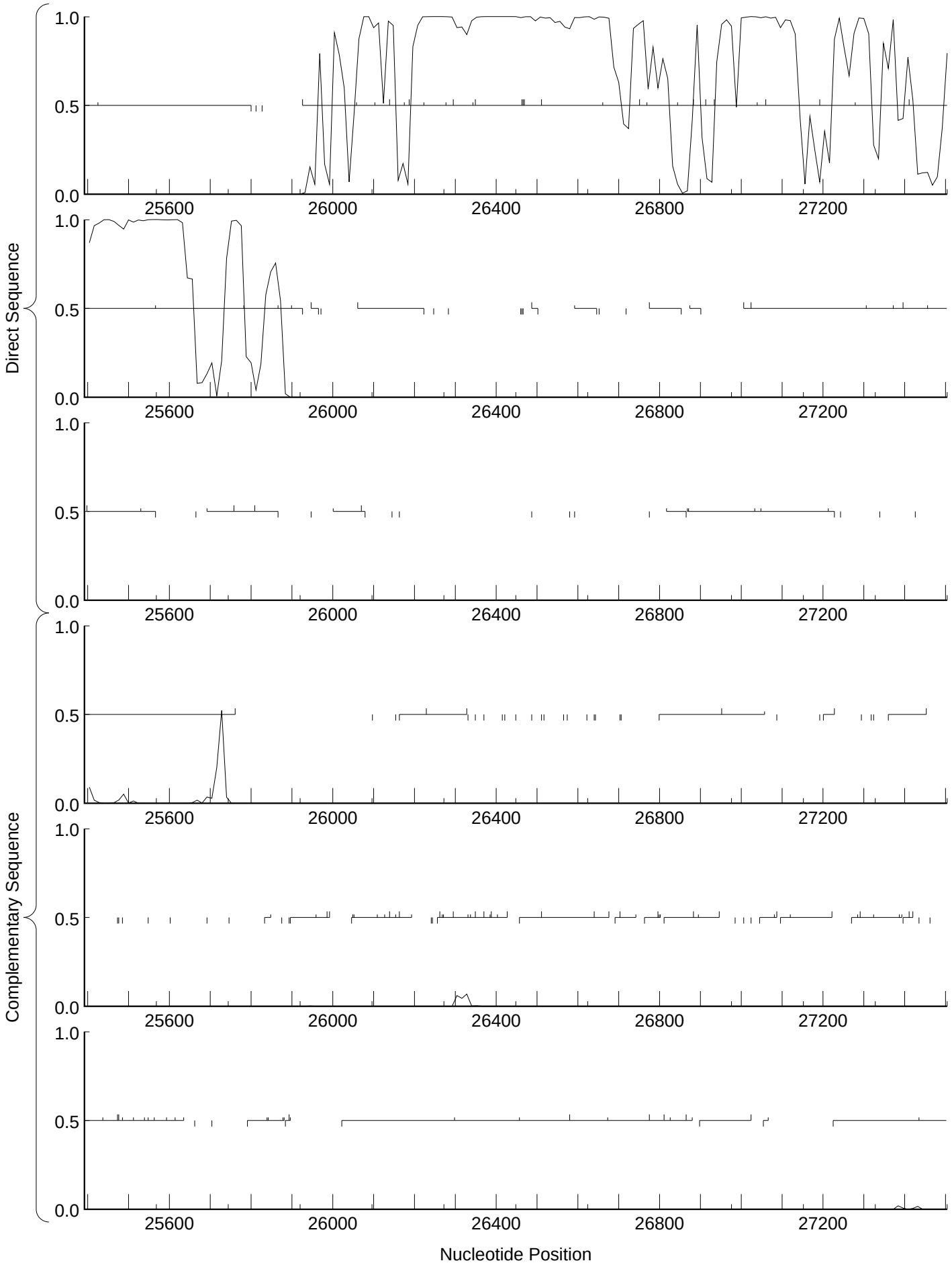
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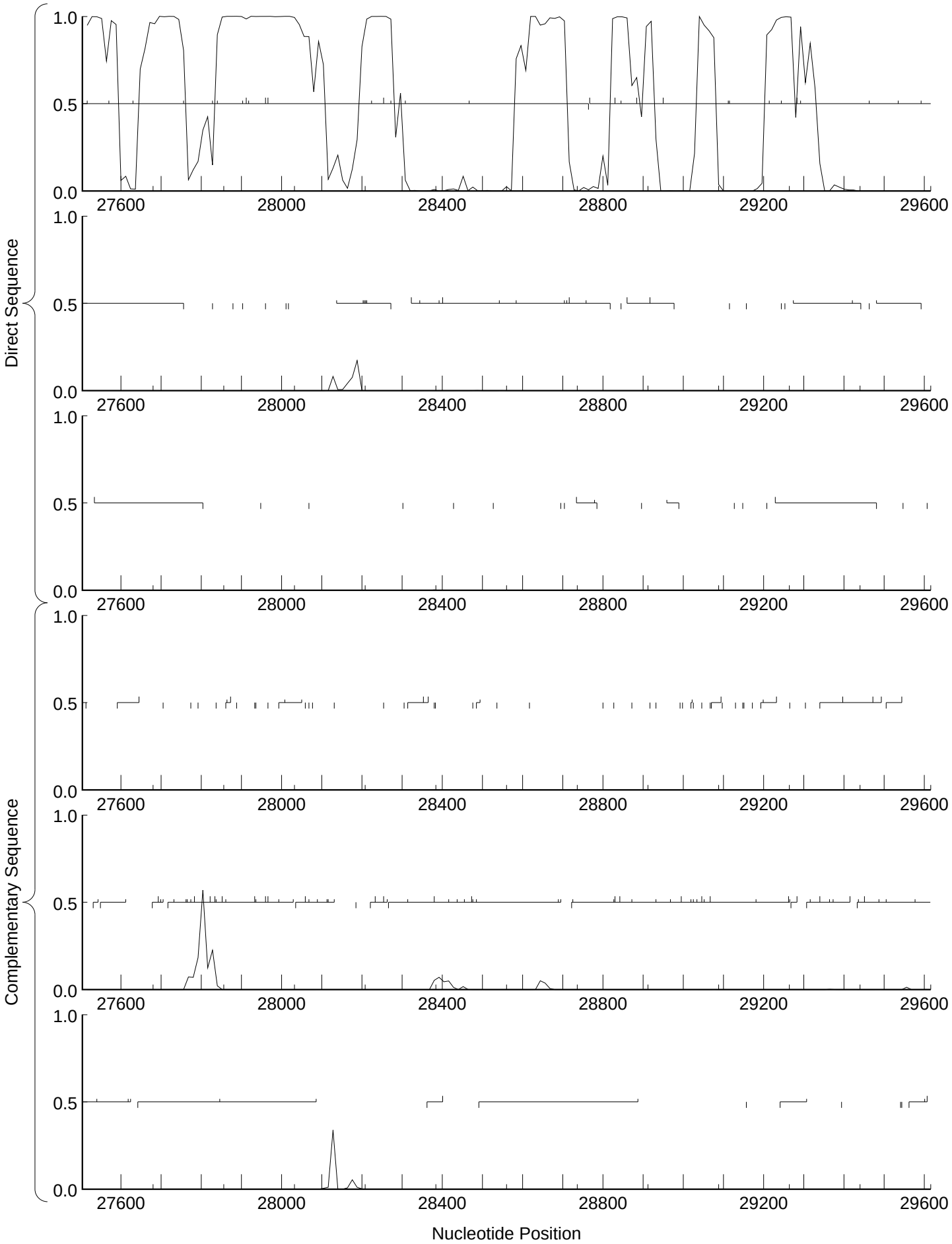
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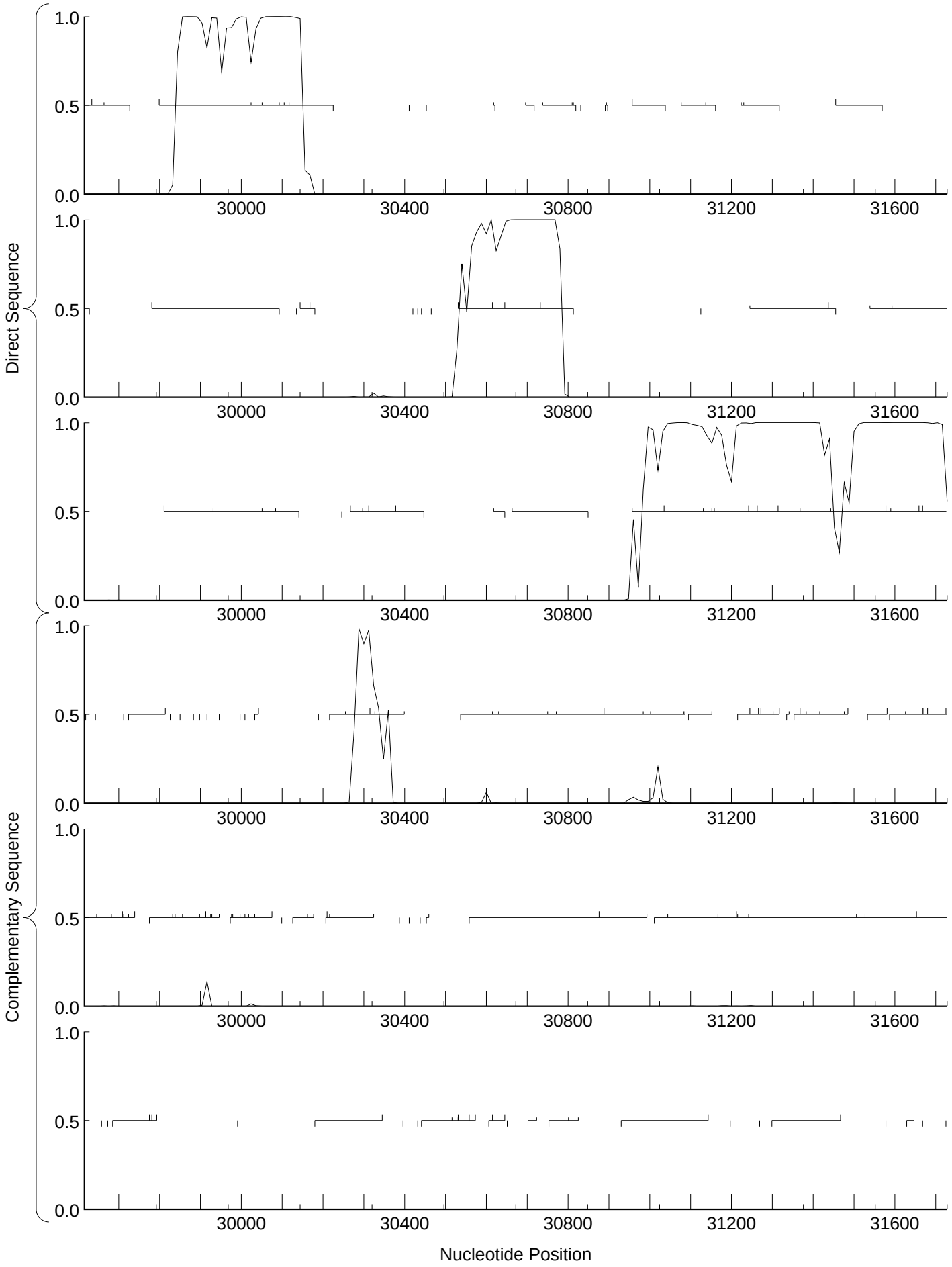

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Complementary Sequence
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27200
Nucleotide Position

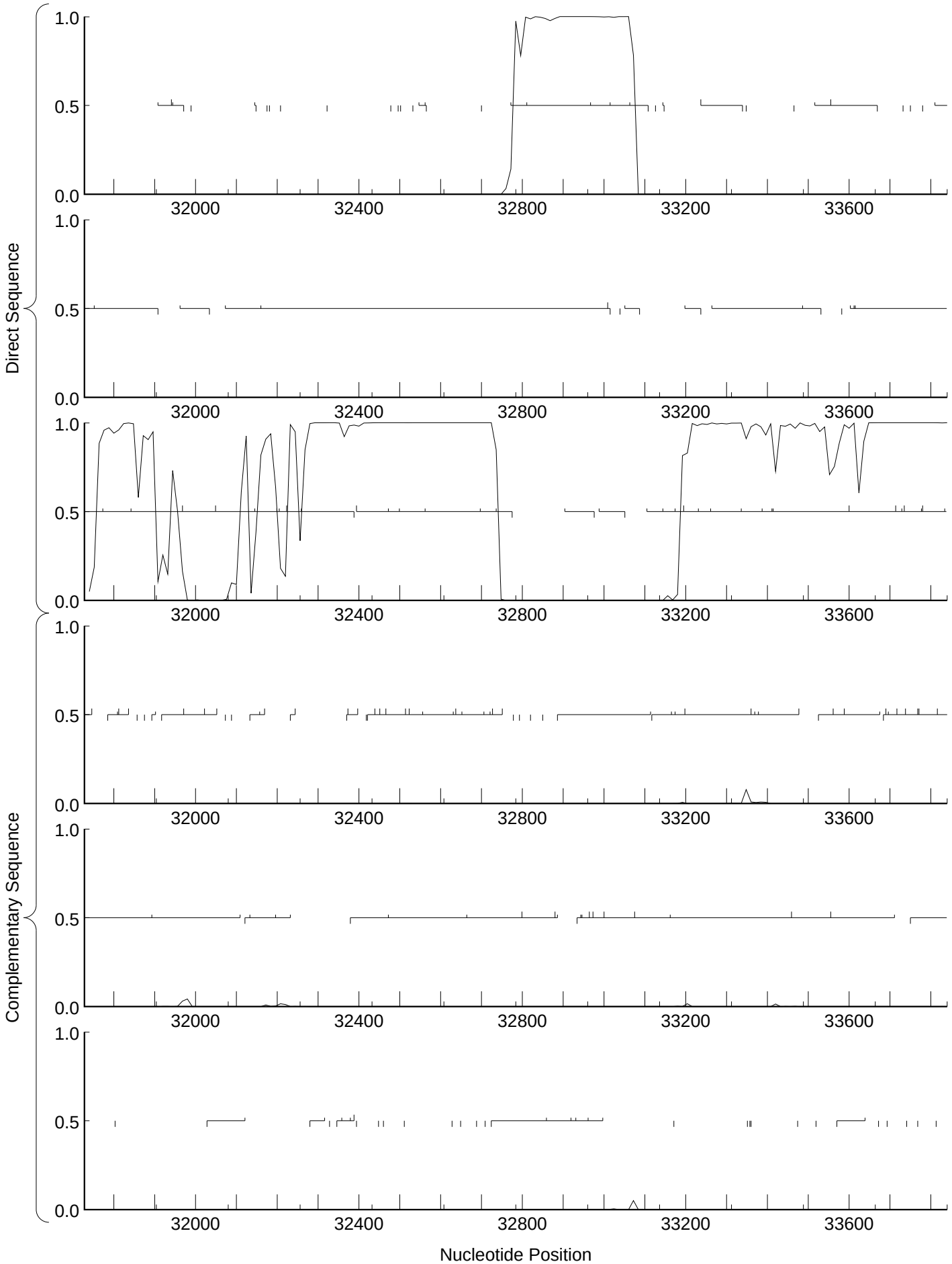
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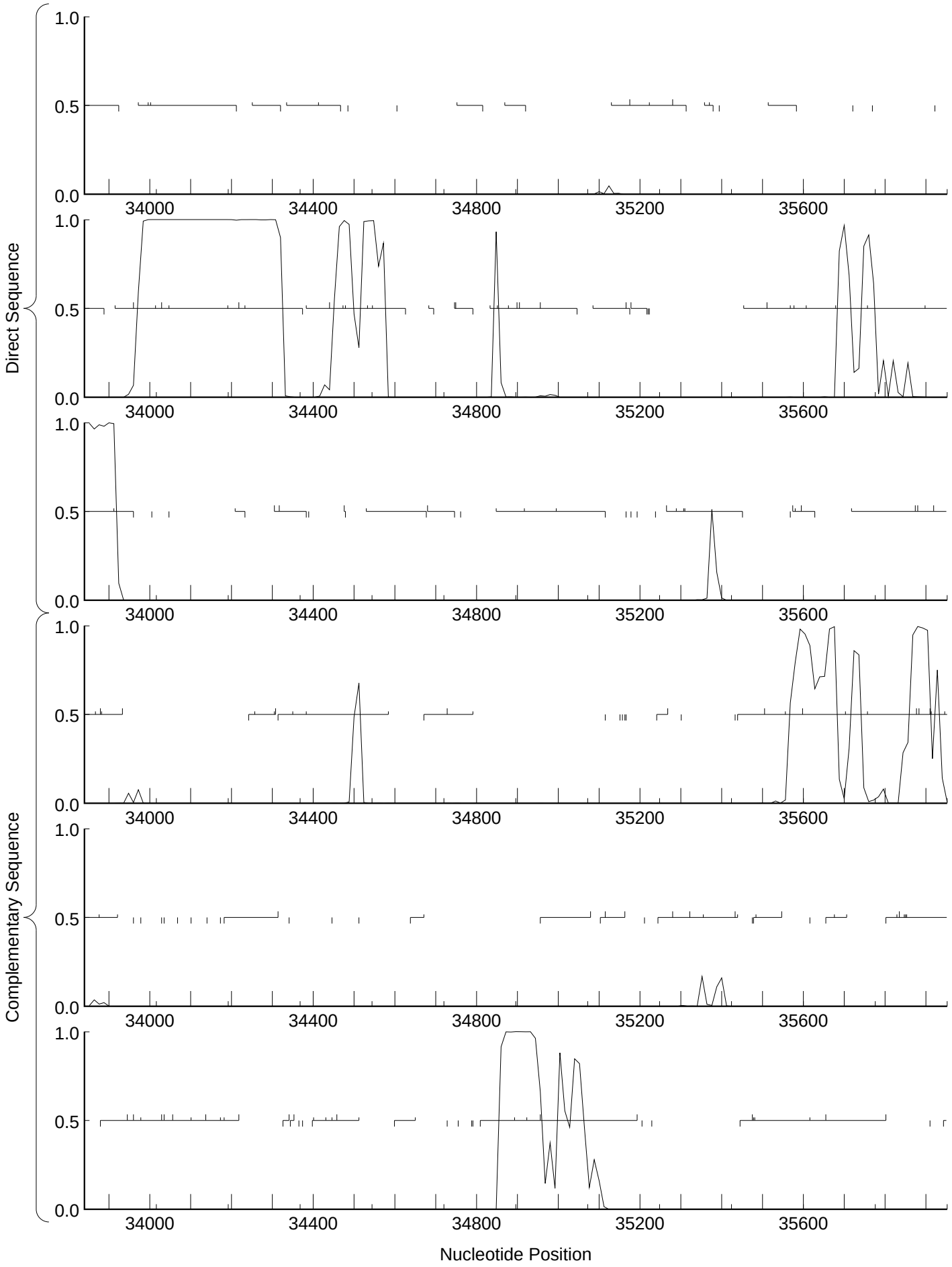


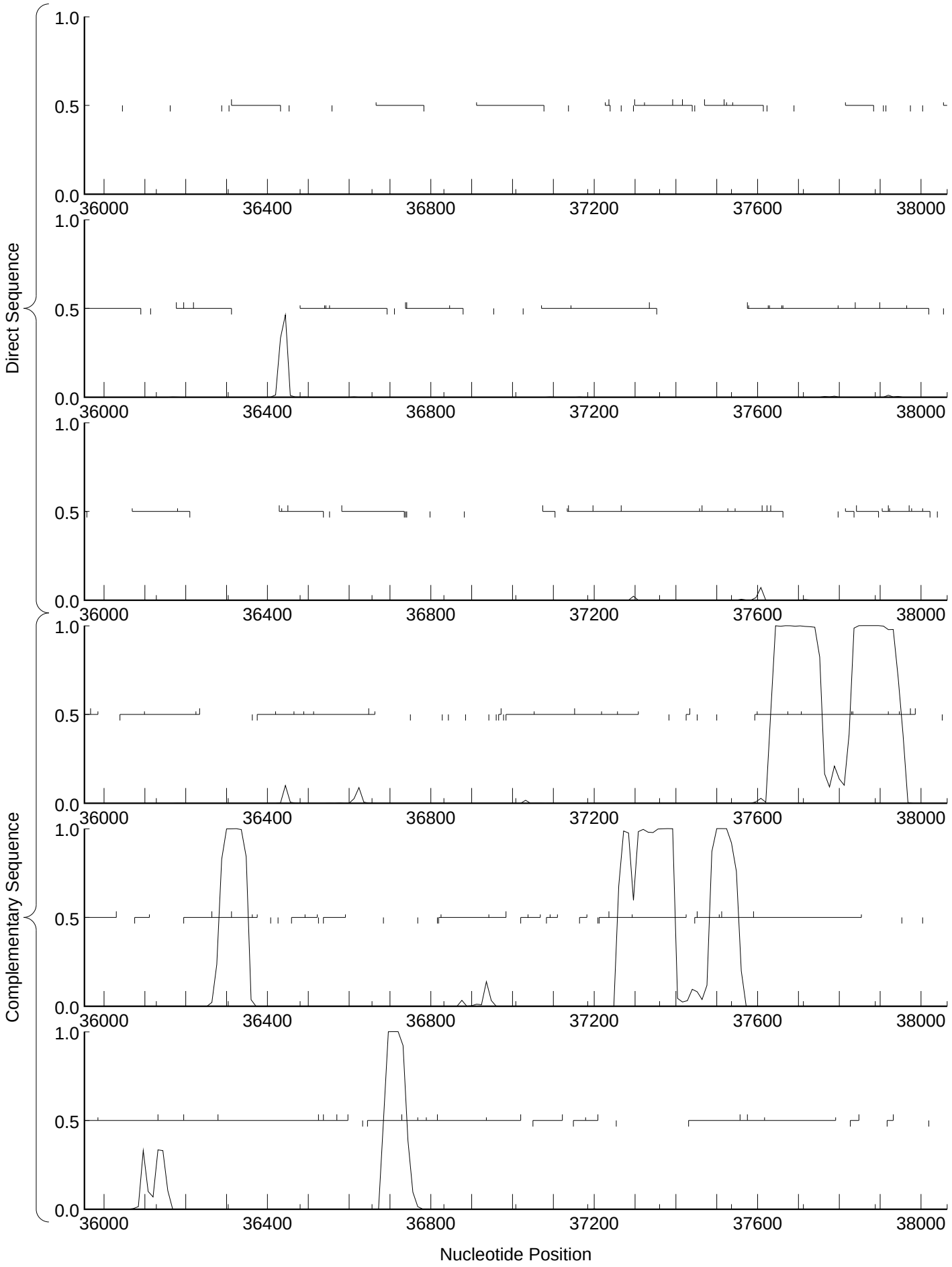
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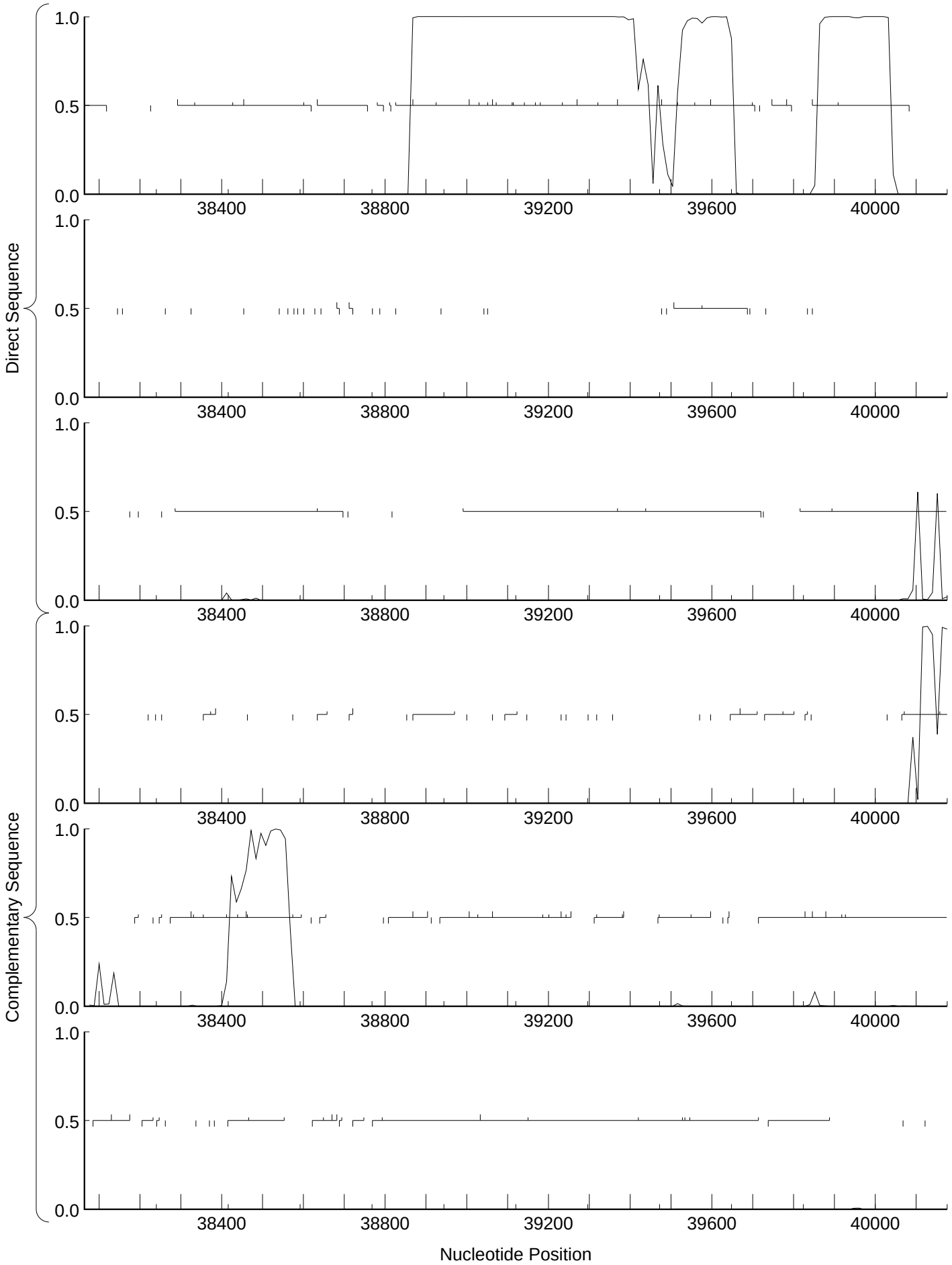


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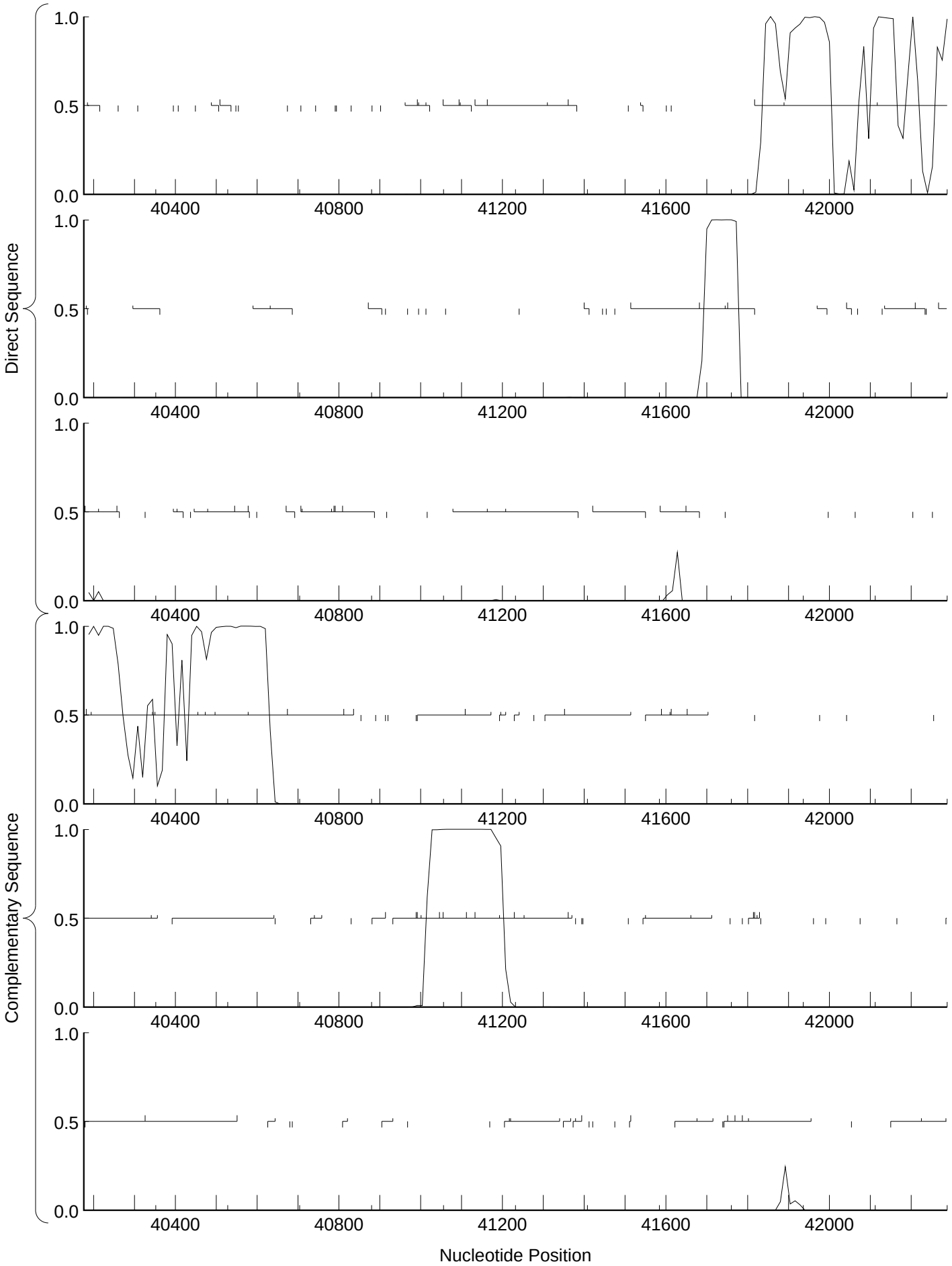


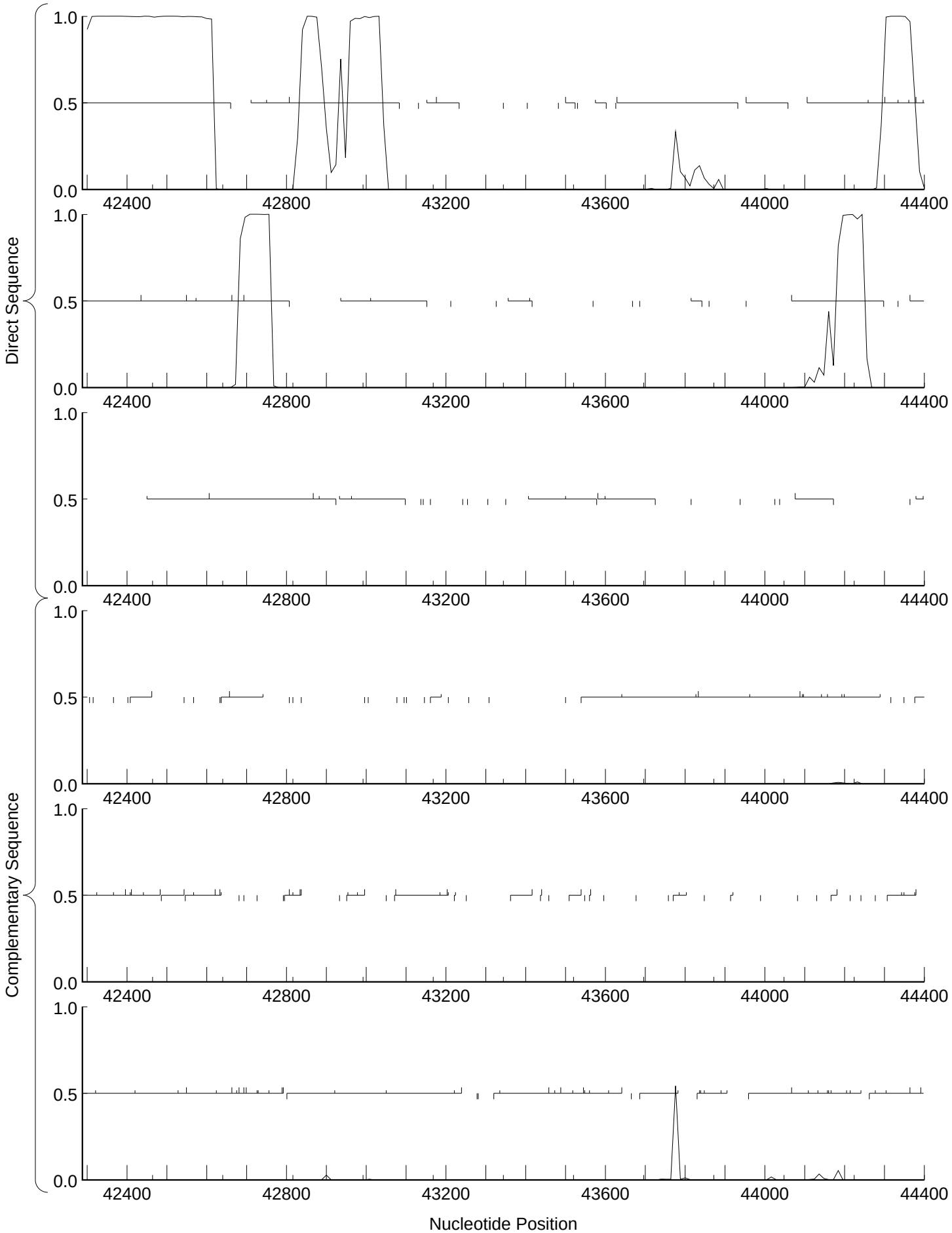

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35600
Nucleotide Position

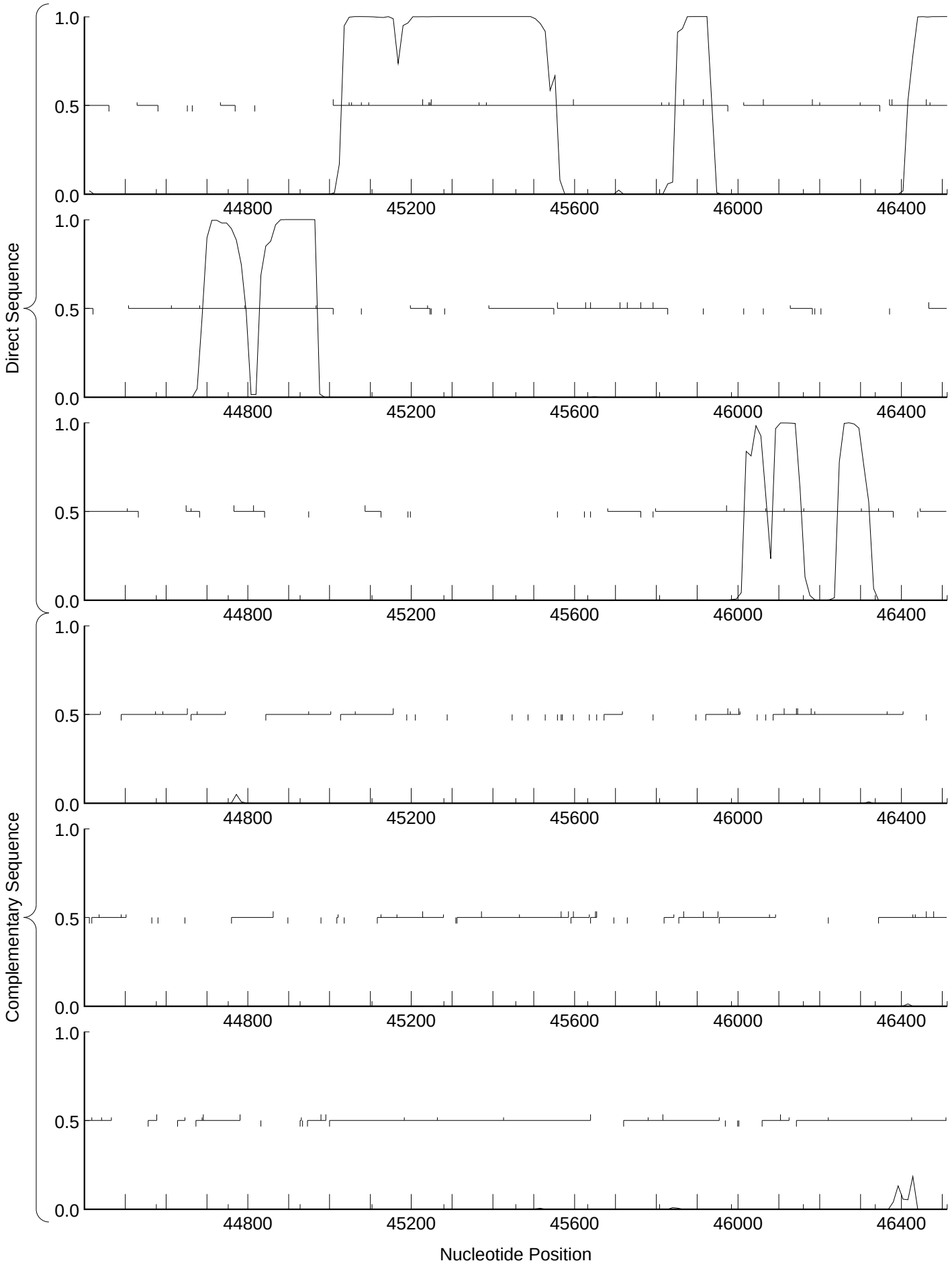

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Complementary Sequence
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0.0
36000
36400
36800
37200
37600
38000
Nucleotide Position


Direct Sequence
Complementary Sequence
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38800
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39600
40000
Nucleotide Position

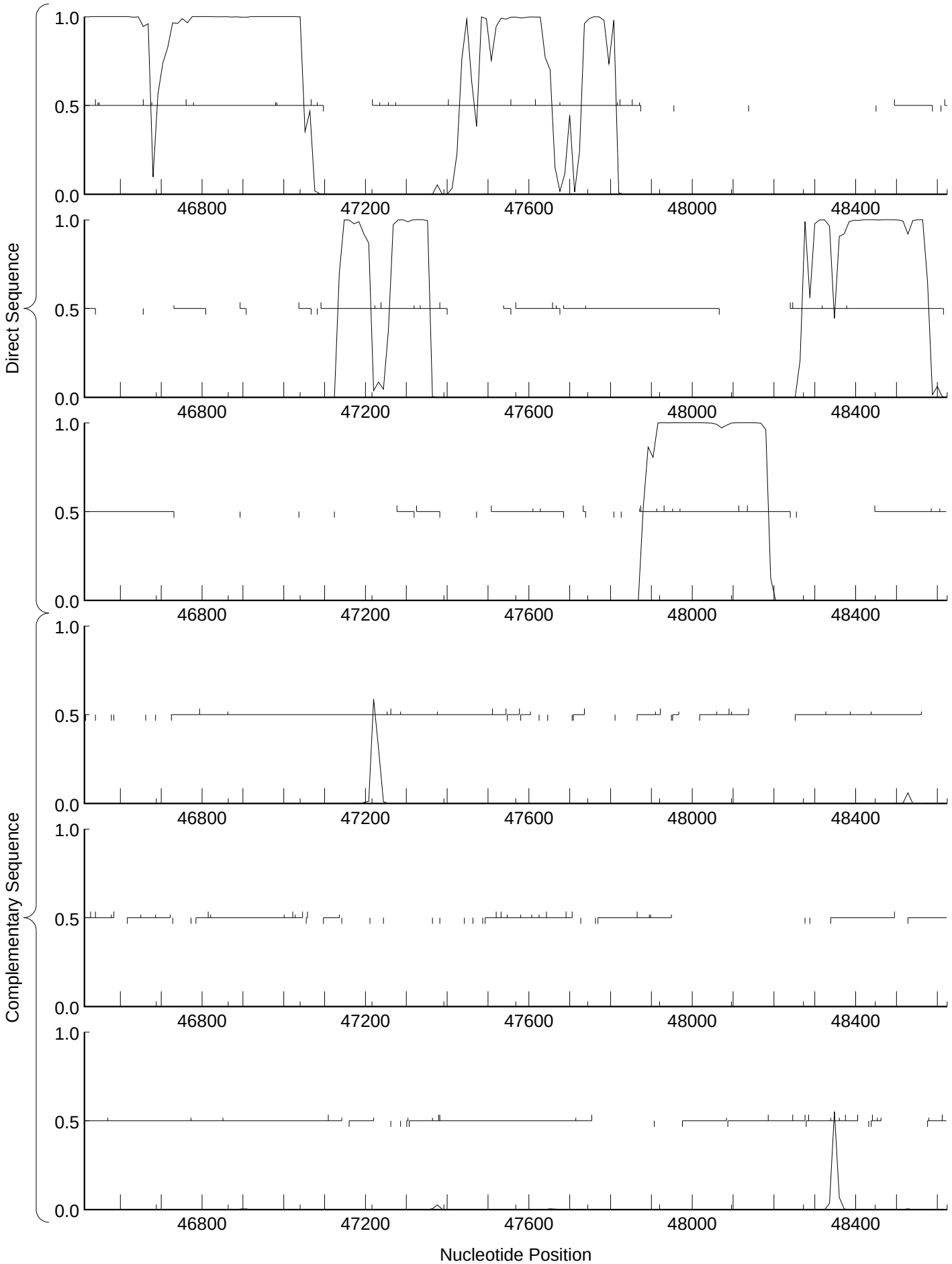
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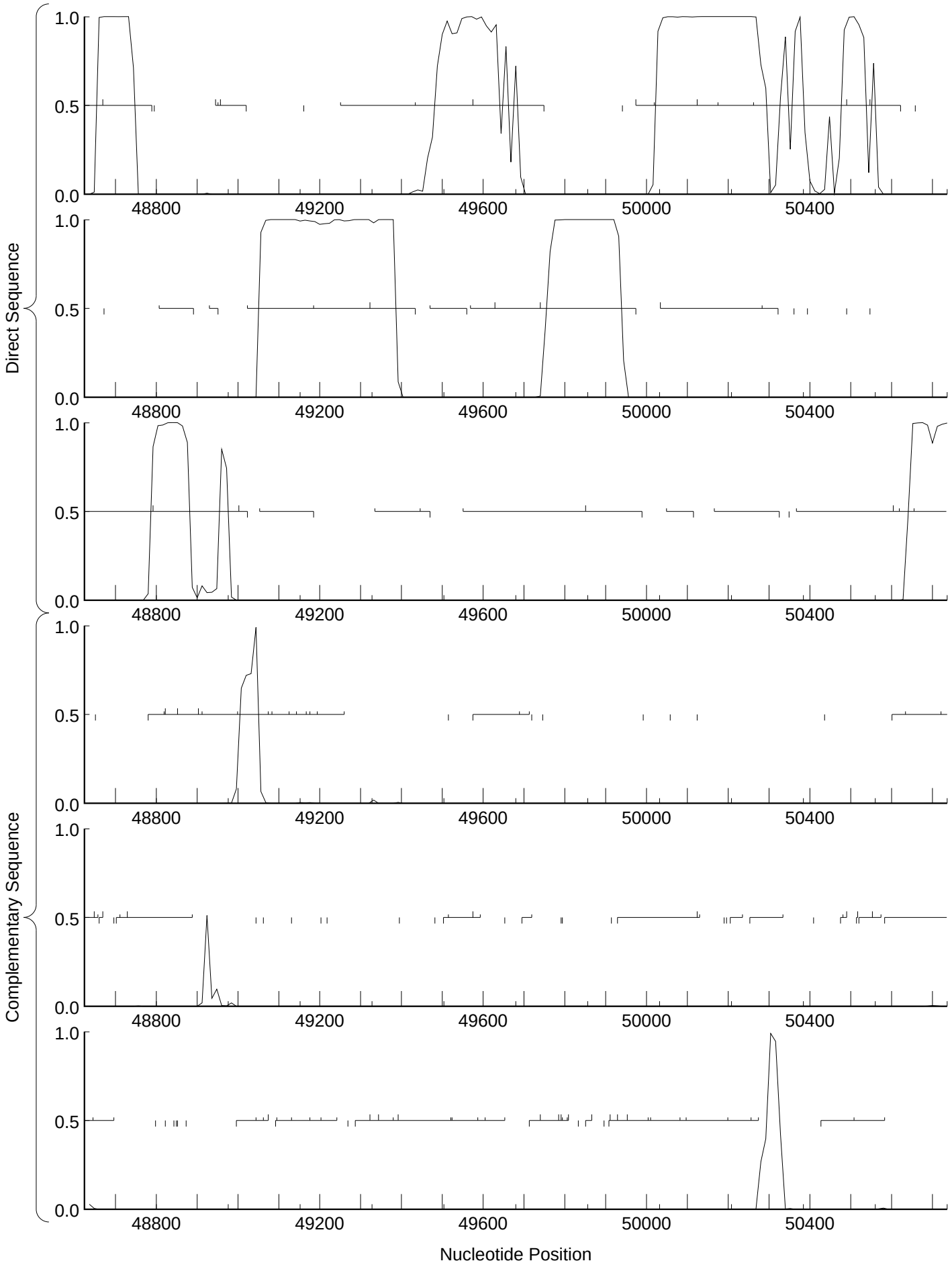


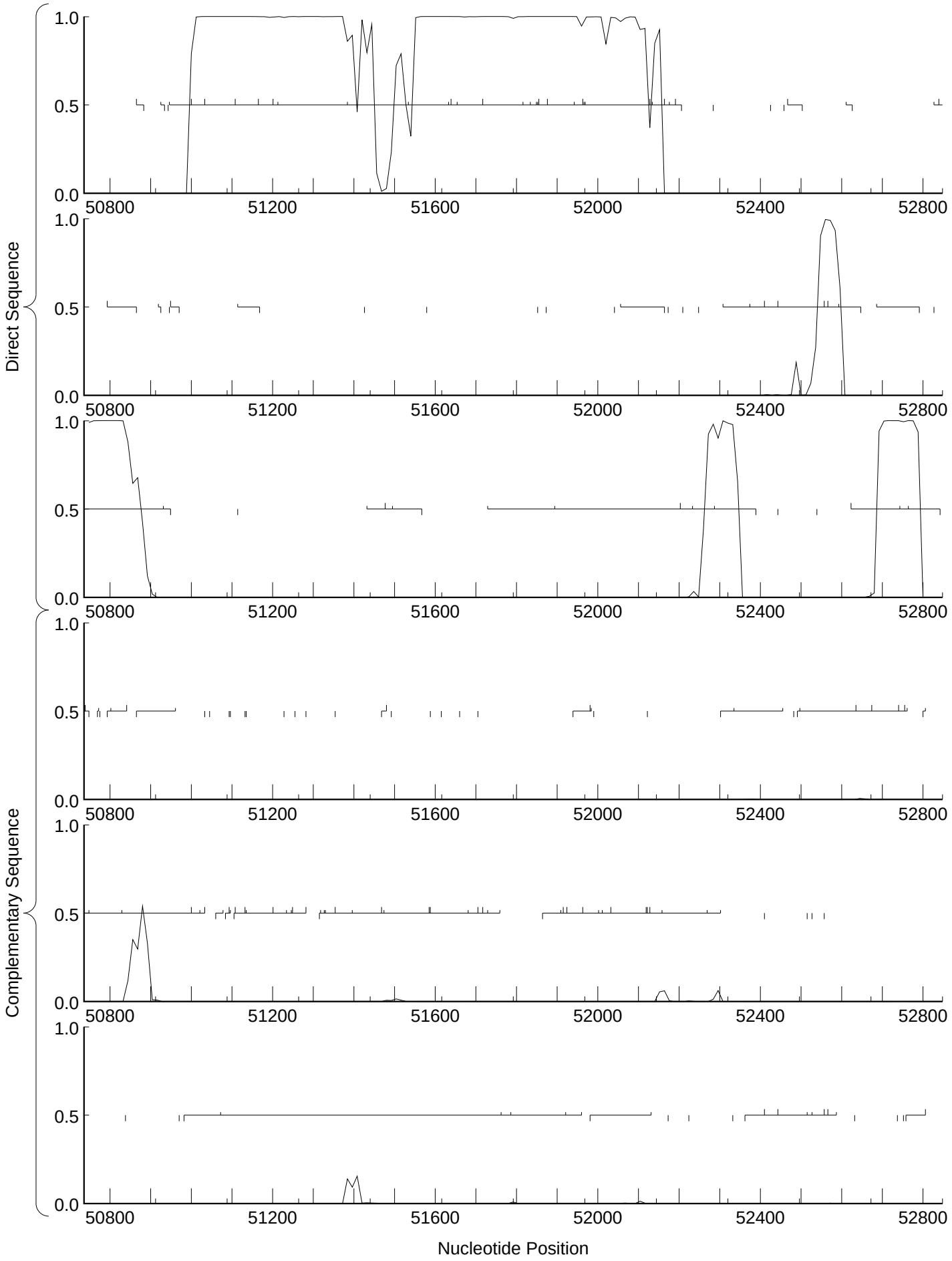

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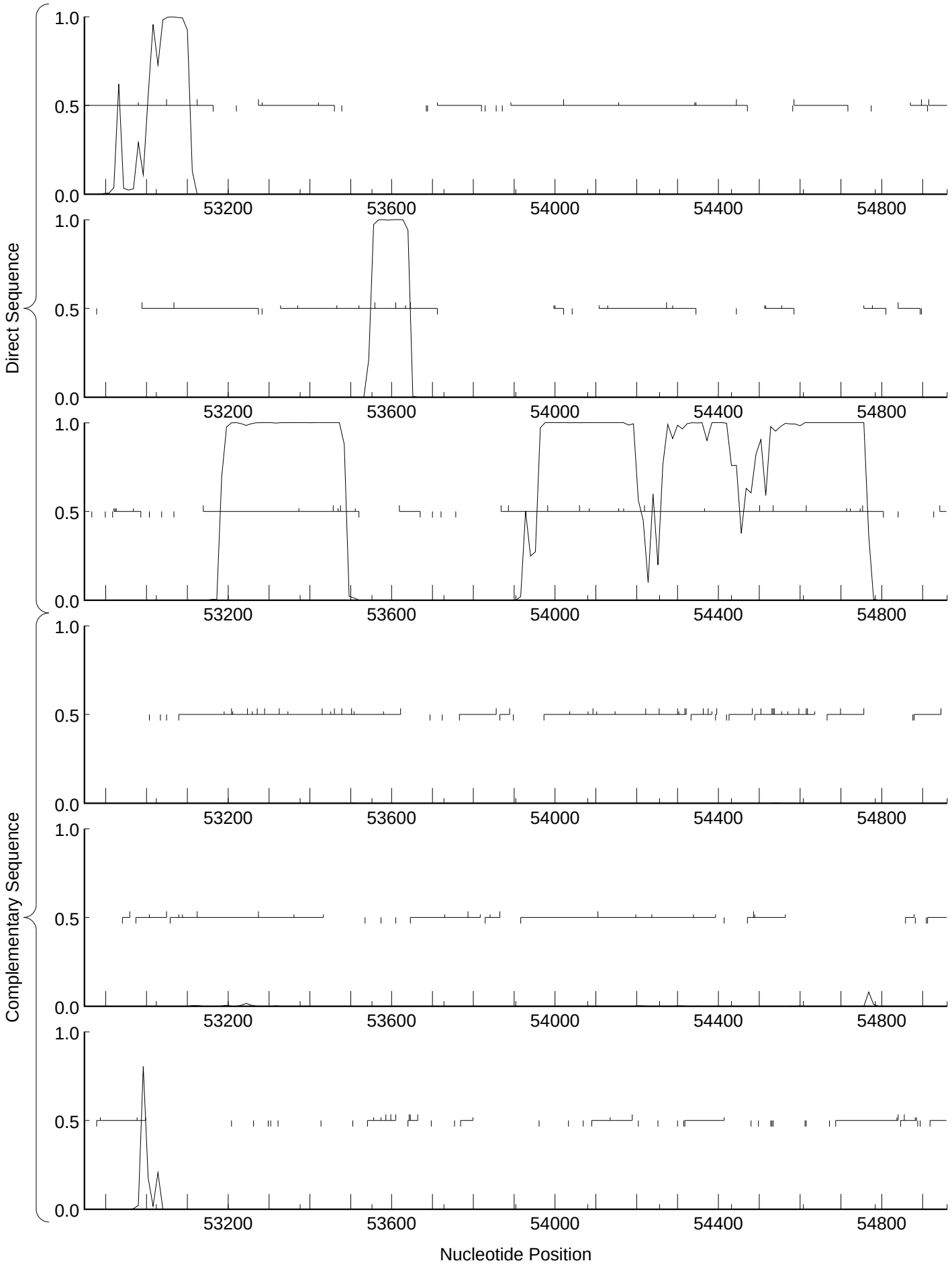

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Nucleotide Position

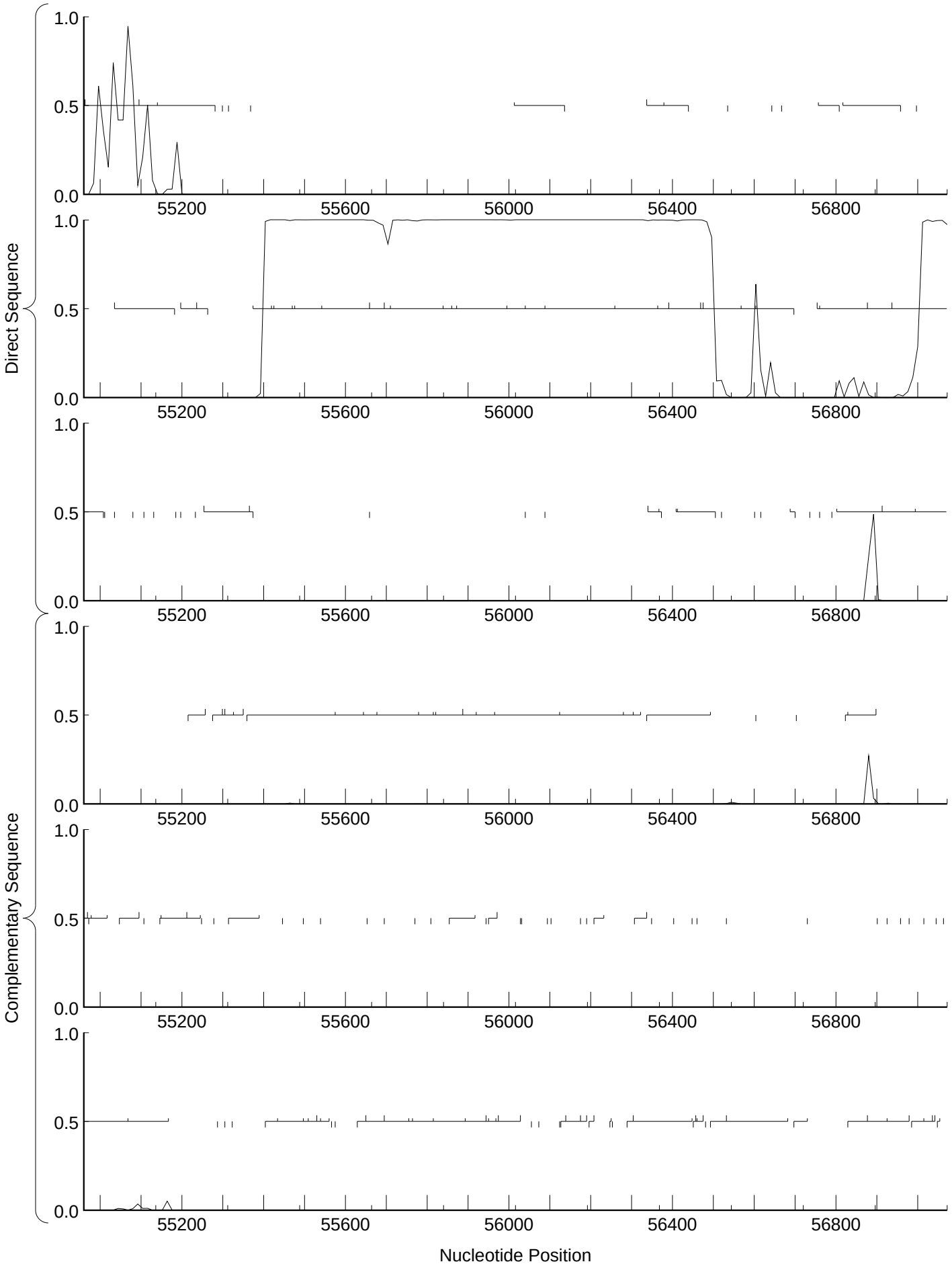
2Say complete sequence, 75598 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 24/37

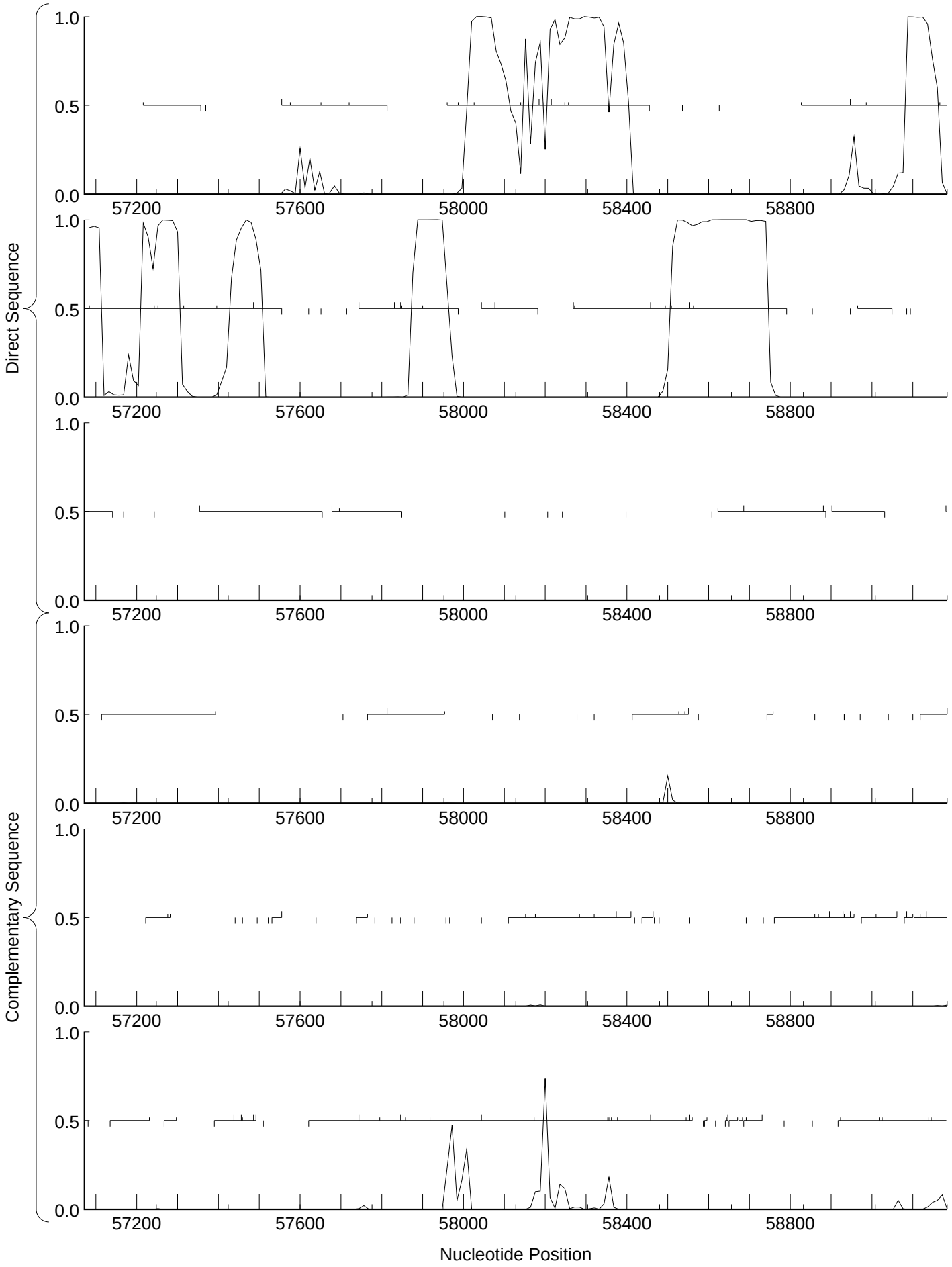



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Nucleotide Position

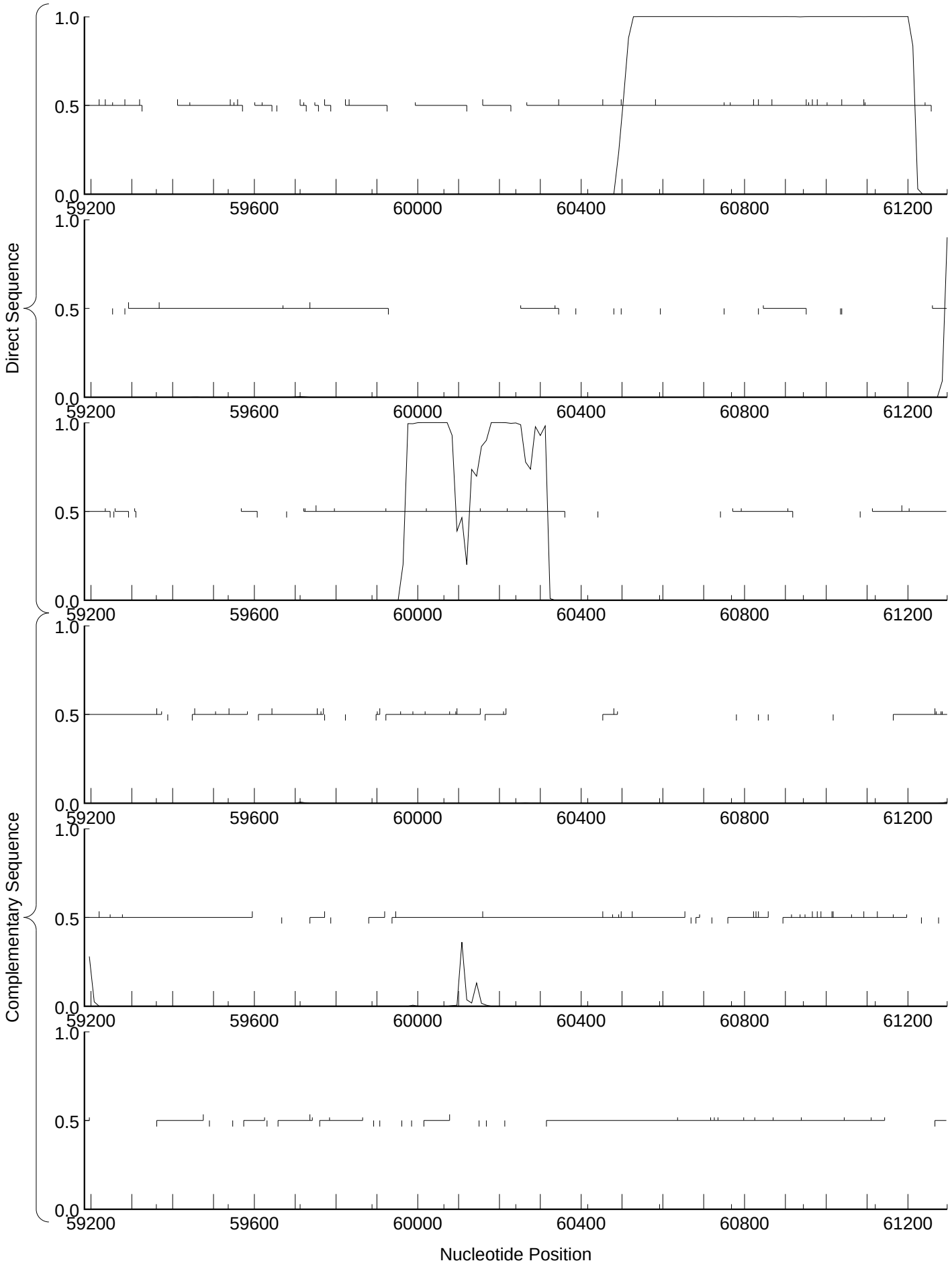

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52400
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54400
54800
Nucleotide Position

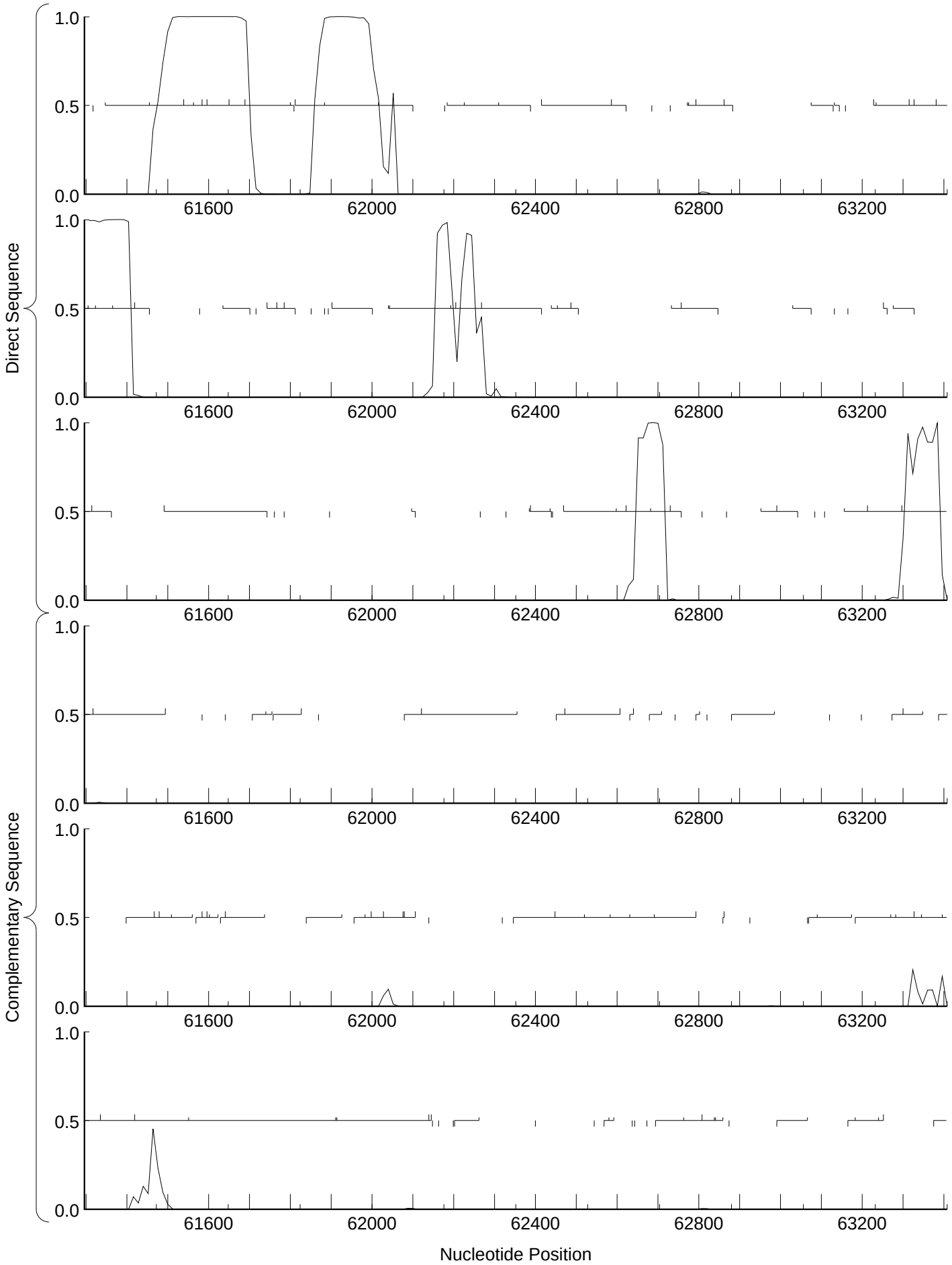

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Nucleotide Position

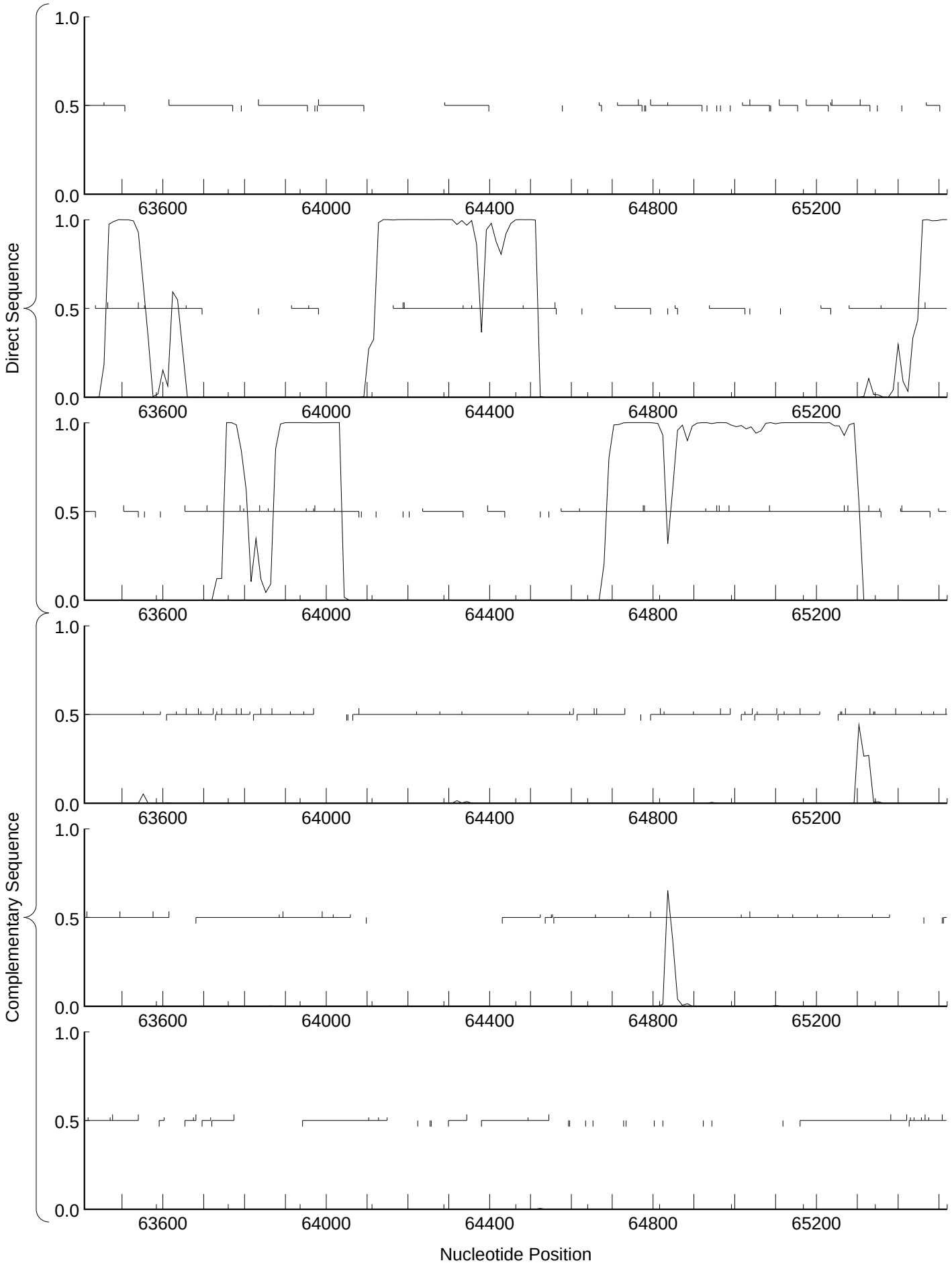

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Complementary Sequence
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58000
58400
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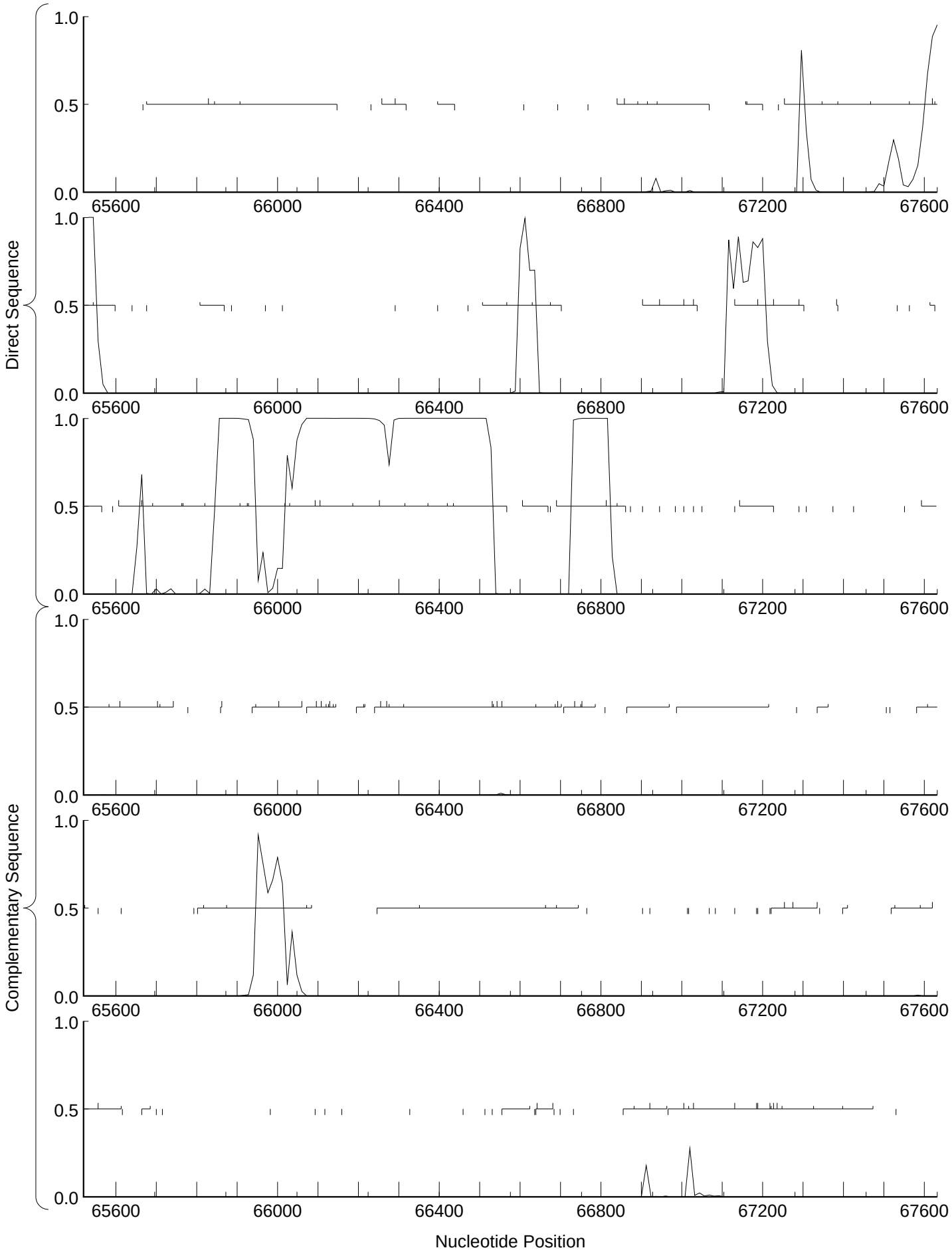
2Say complete sequence, 75598 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 30/37



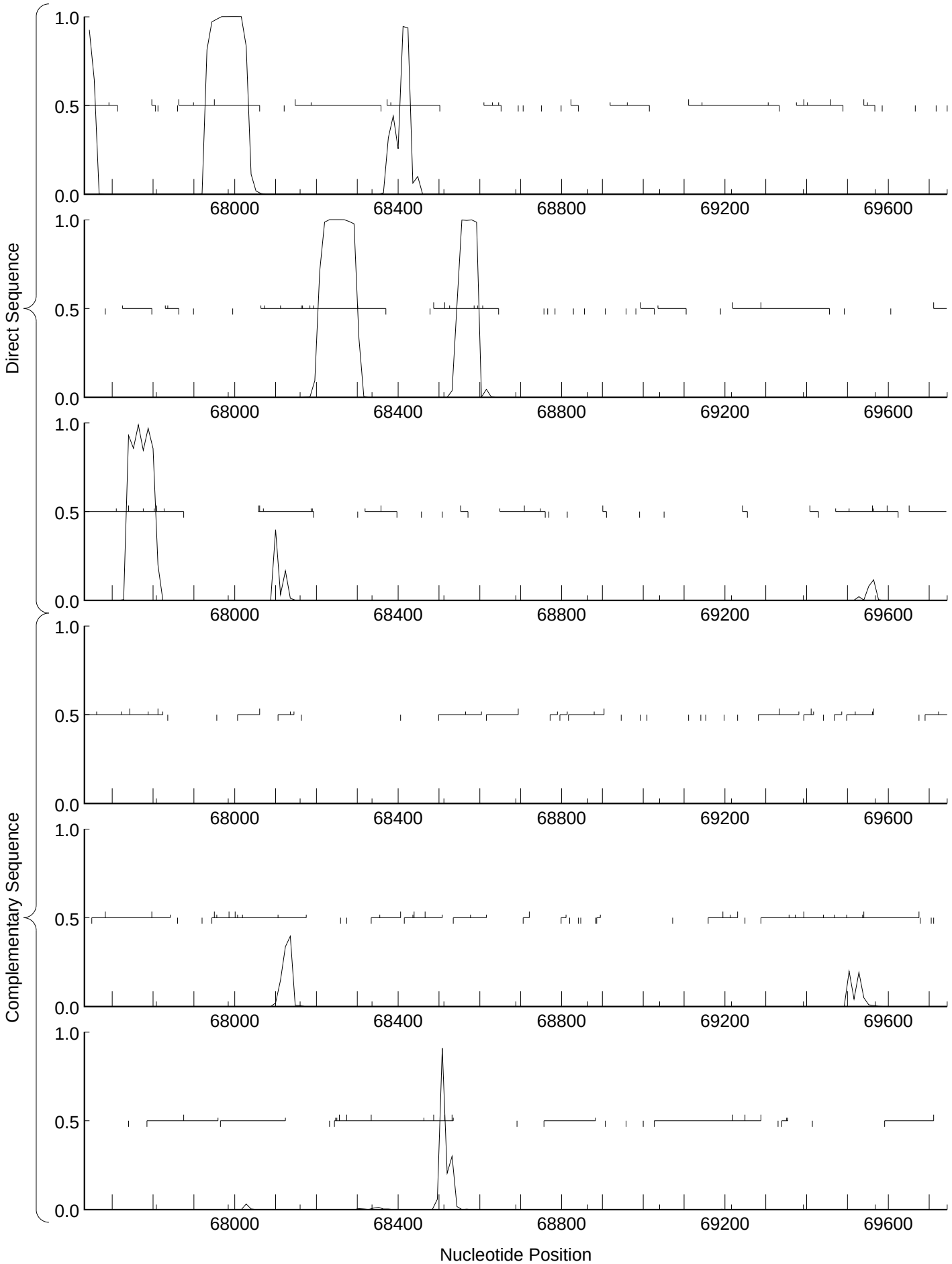
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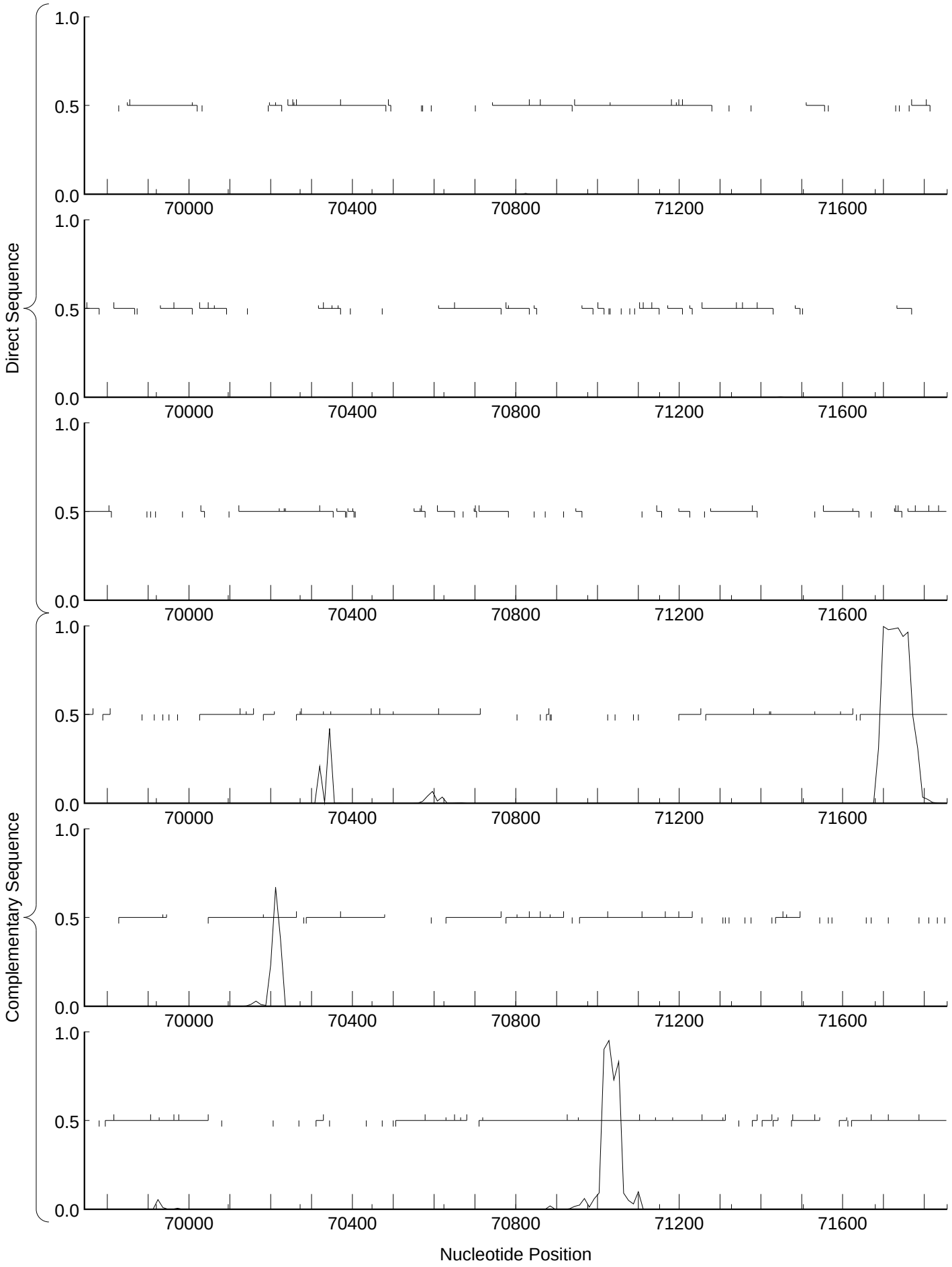

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Nucleotide Position

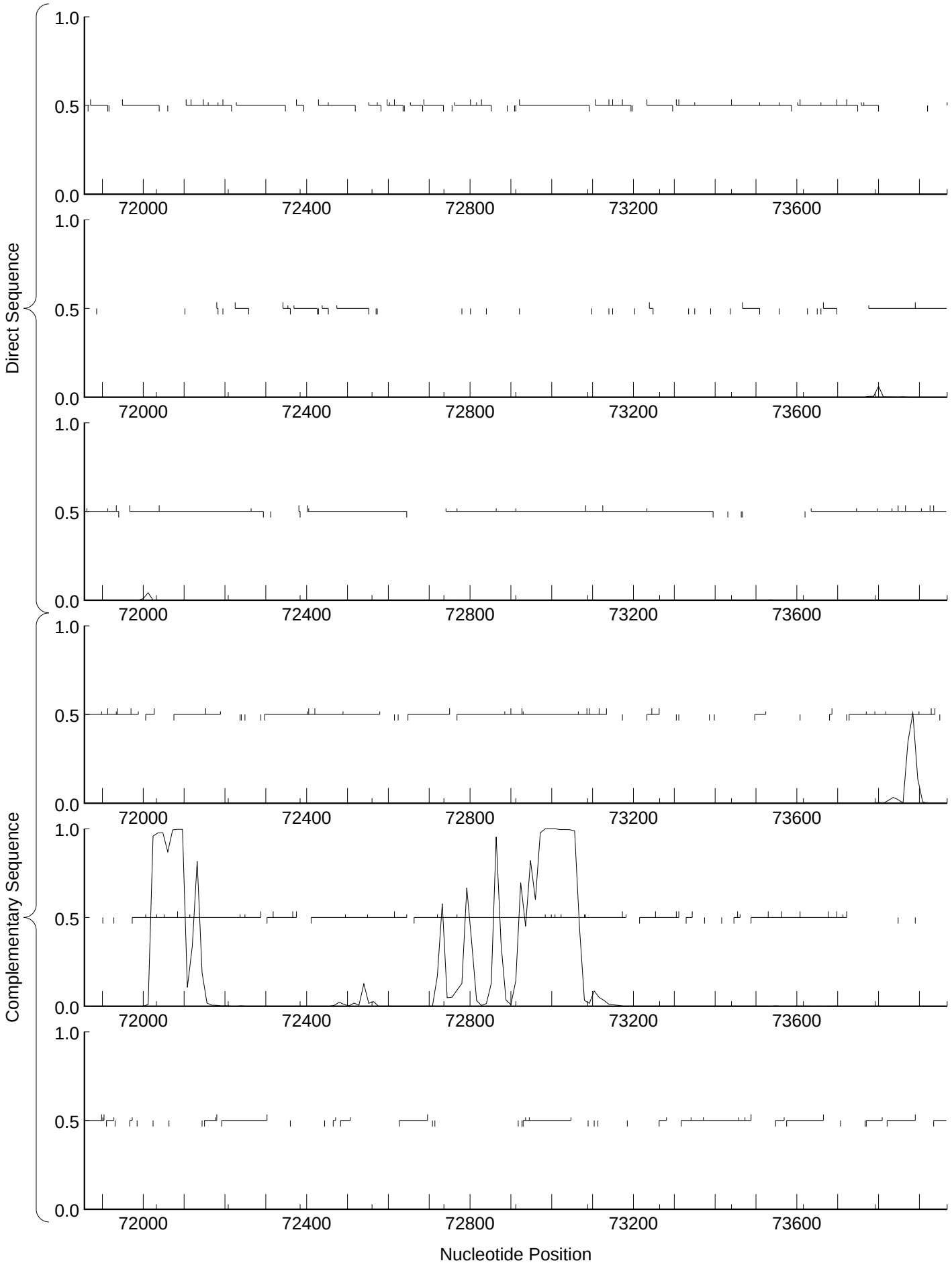
2Say complete sequence, 75598 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 34/37



2Say complete sequence, 75598 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 35/37



2Say complete sequence, 75598 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 36/37



2Say complete sequence, 75598 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 37/37

