GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage KilKor complete sequence, 48916 bp including 13-base 3' overhang (CCTGC
Analysis Date    :  3/29/24 at 3:30:39
Pages            :  25
Sequence Length  :  48916 bp
GC Content       :   67.21%

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

PS-Version       :  1.2

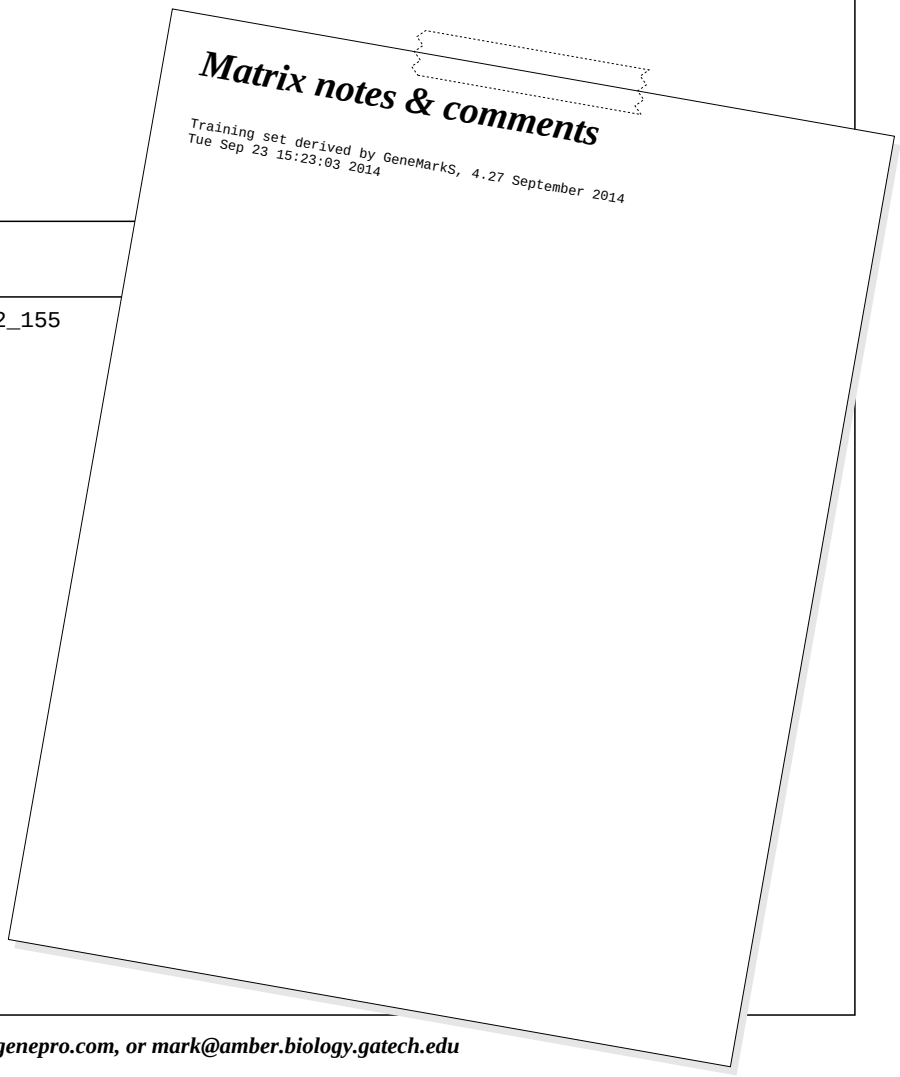
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Author     :  -

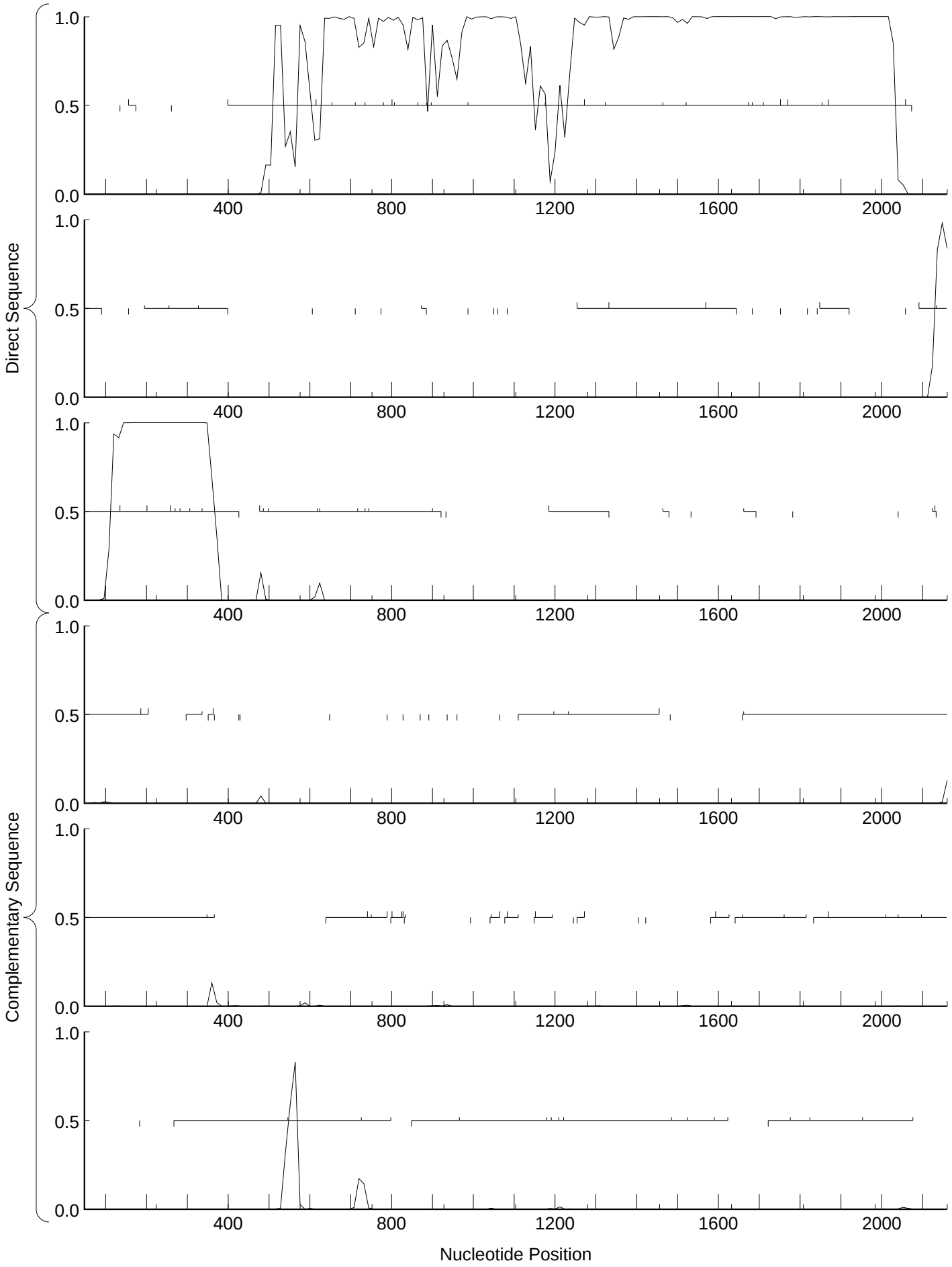
Order      :  4


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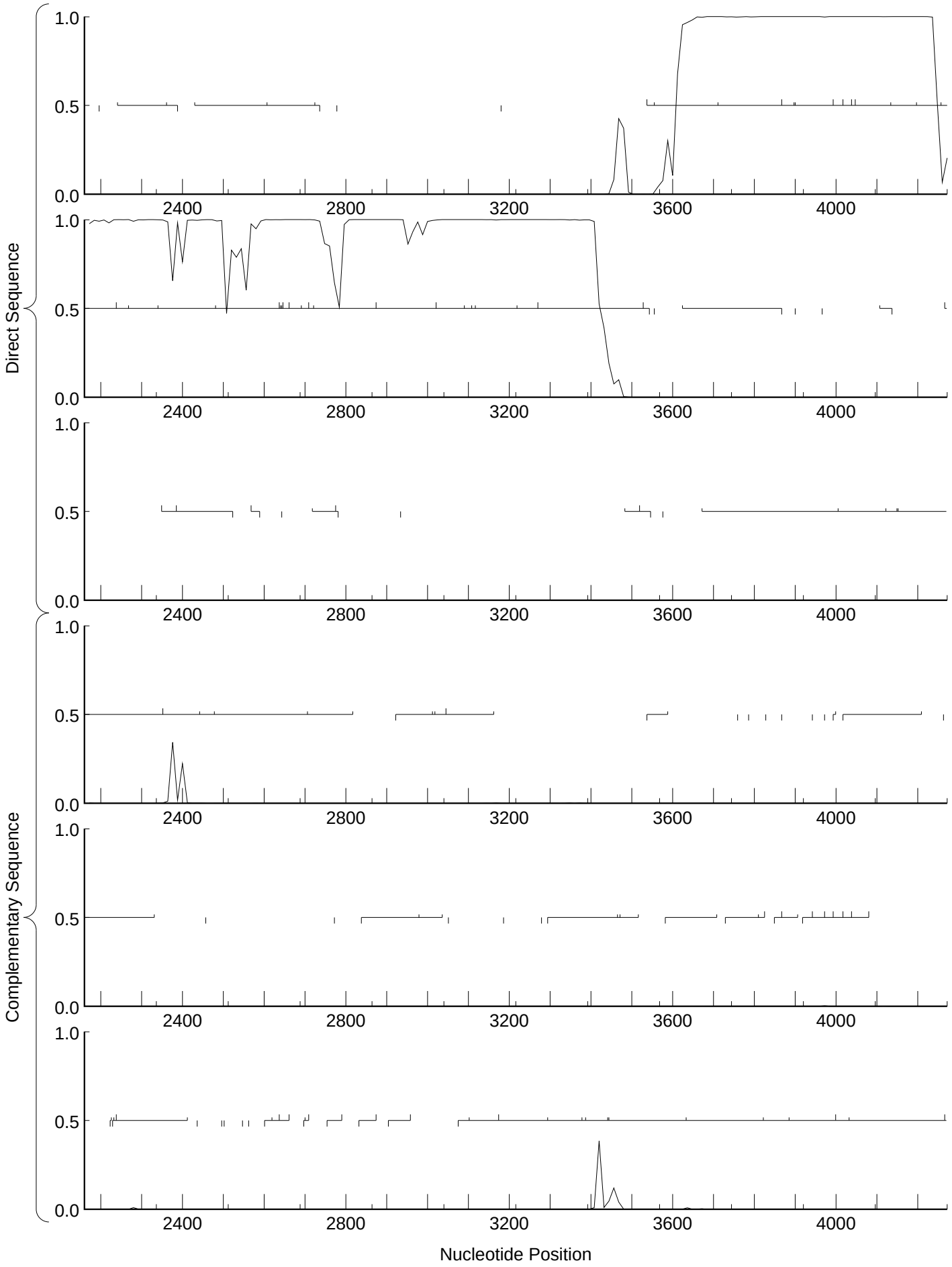


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

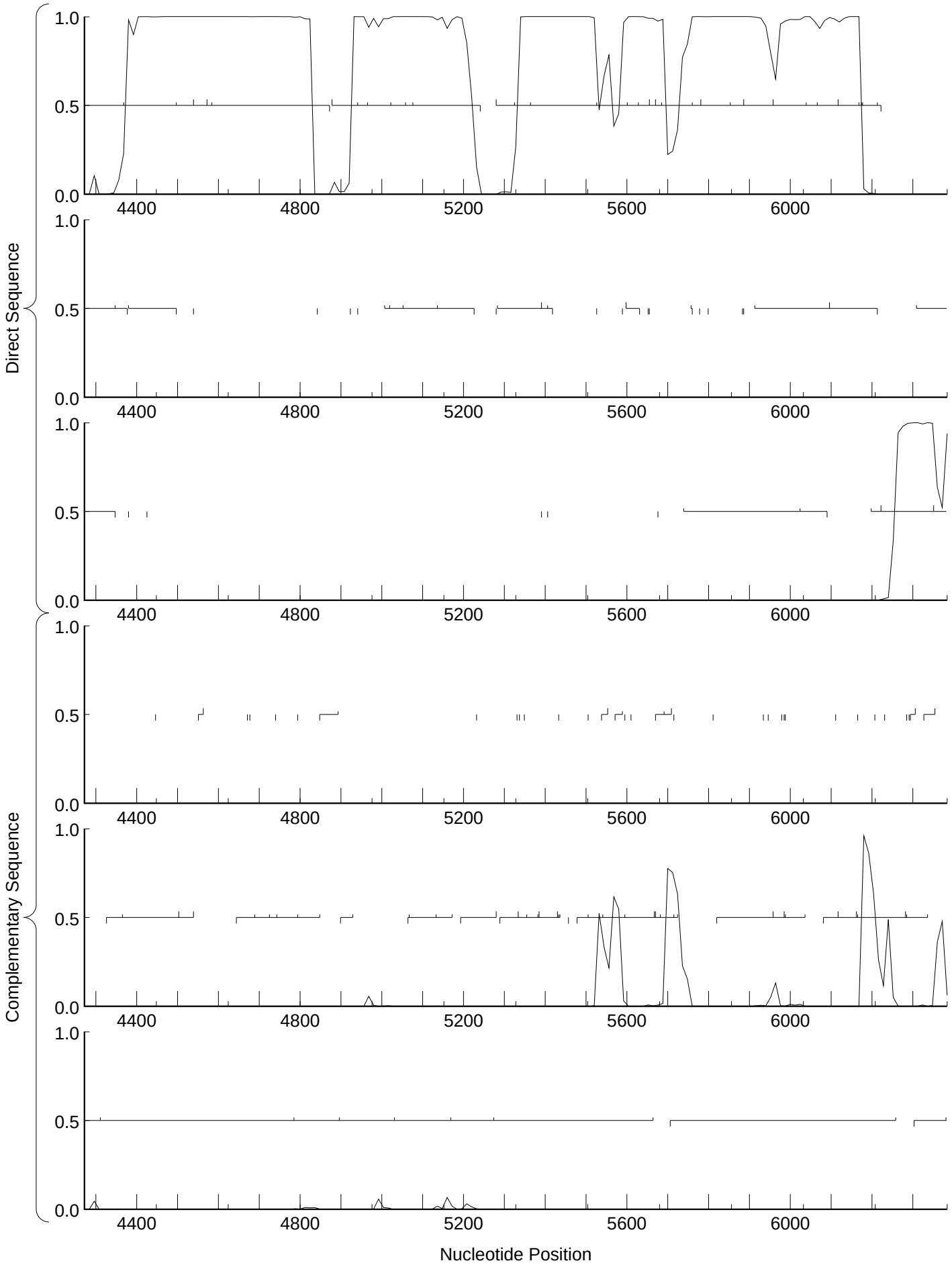
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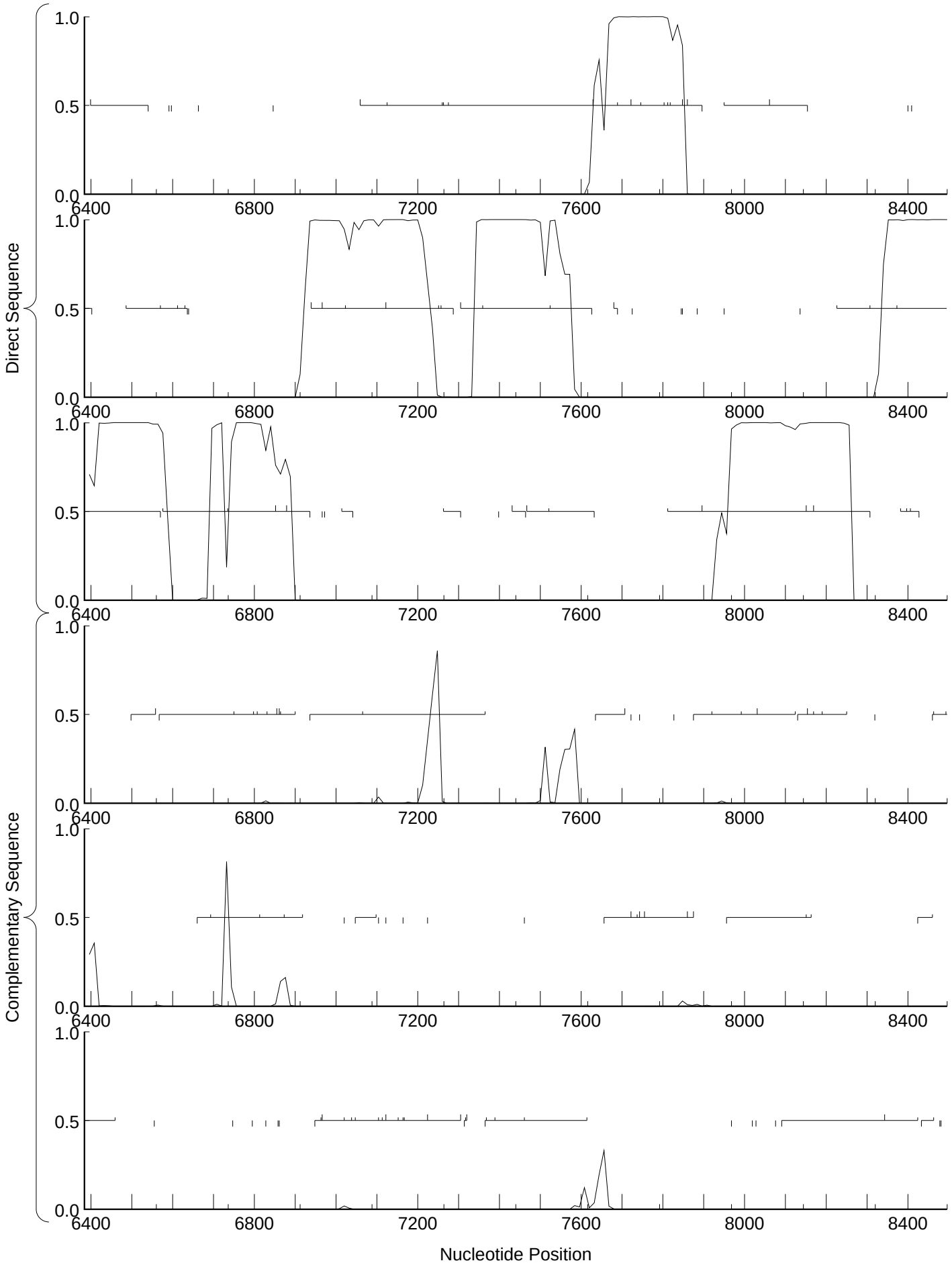
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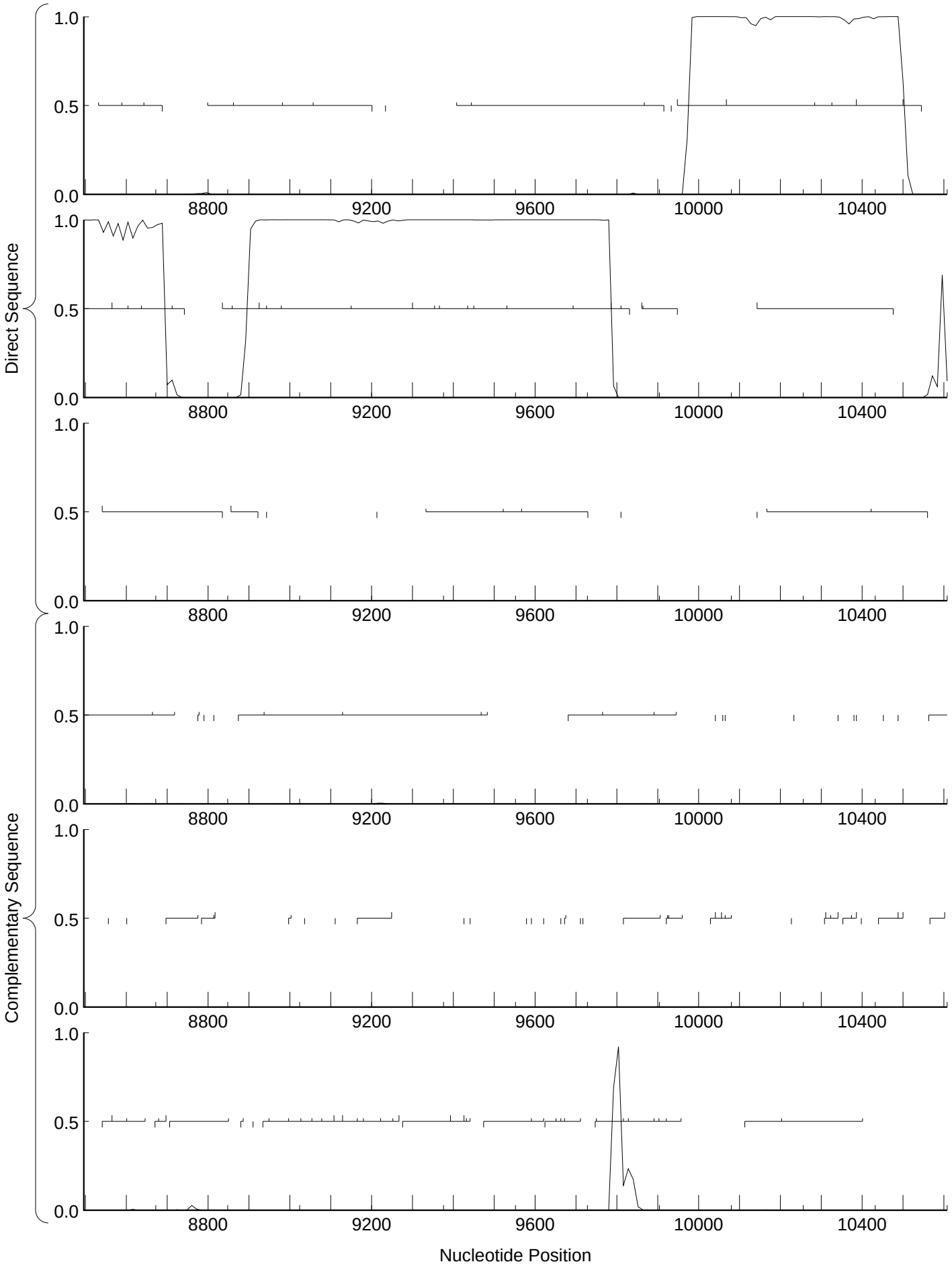
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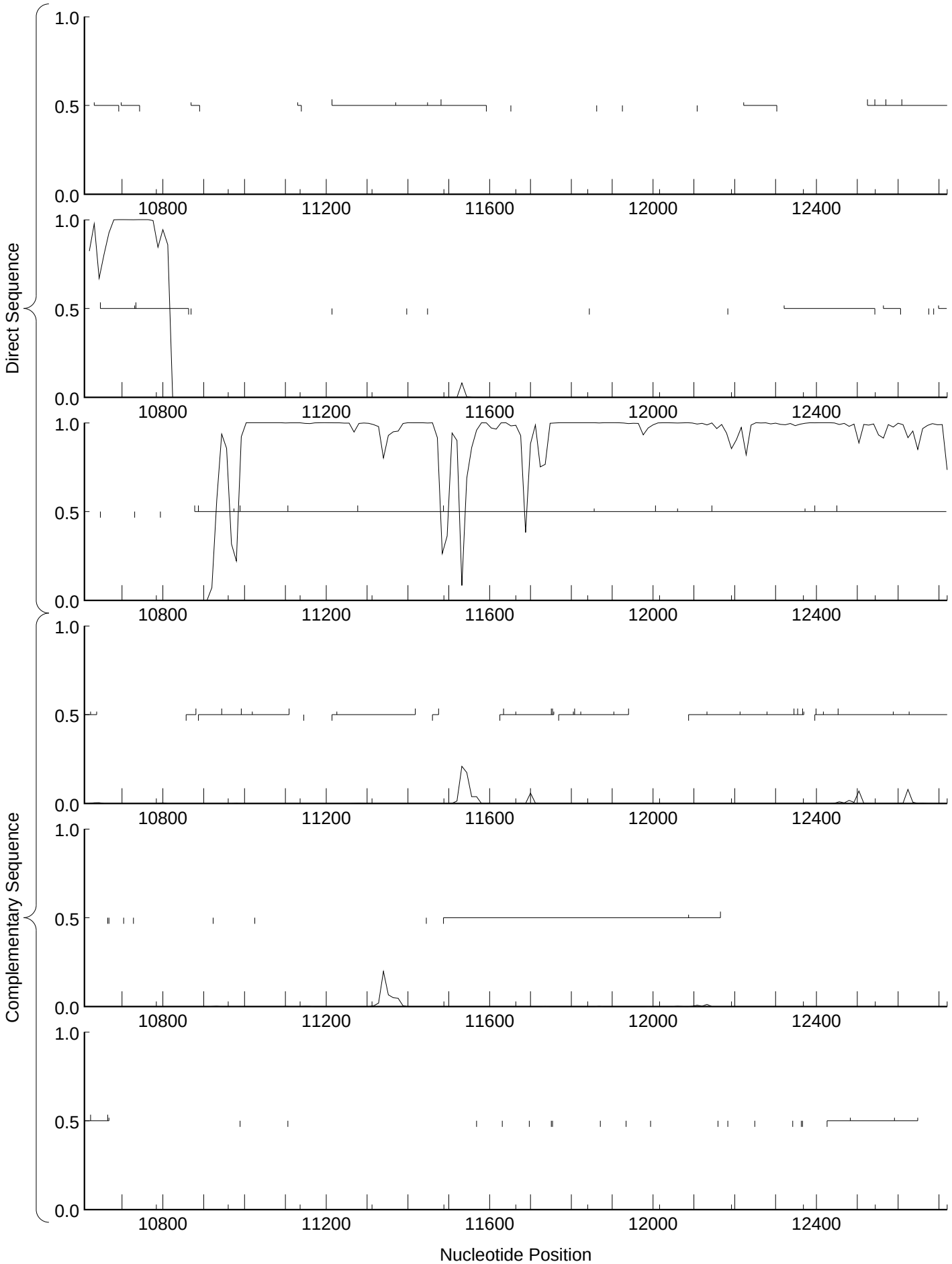
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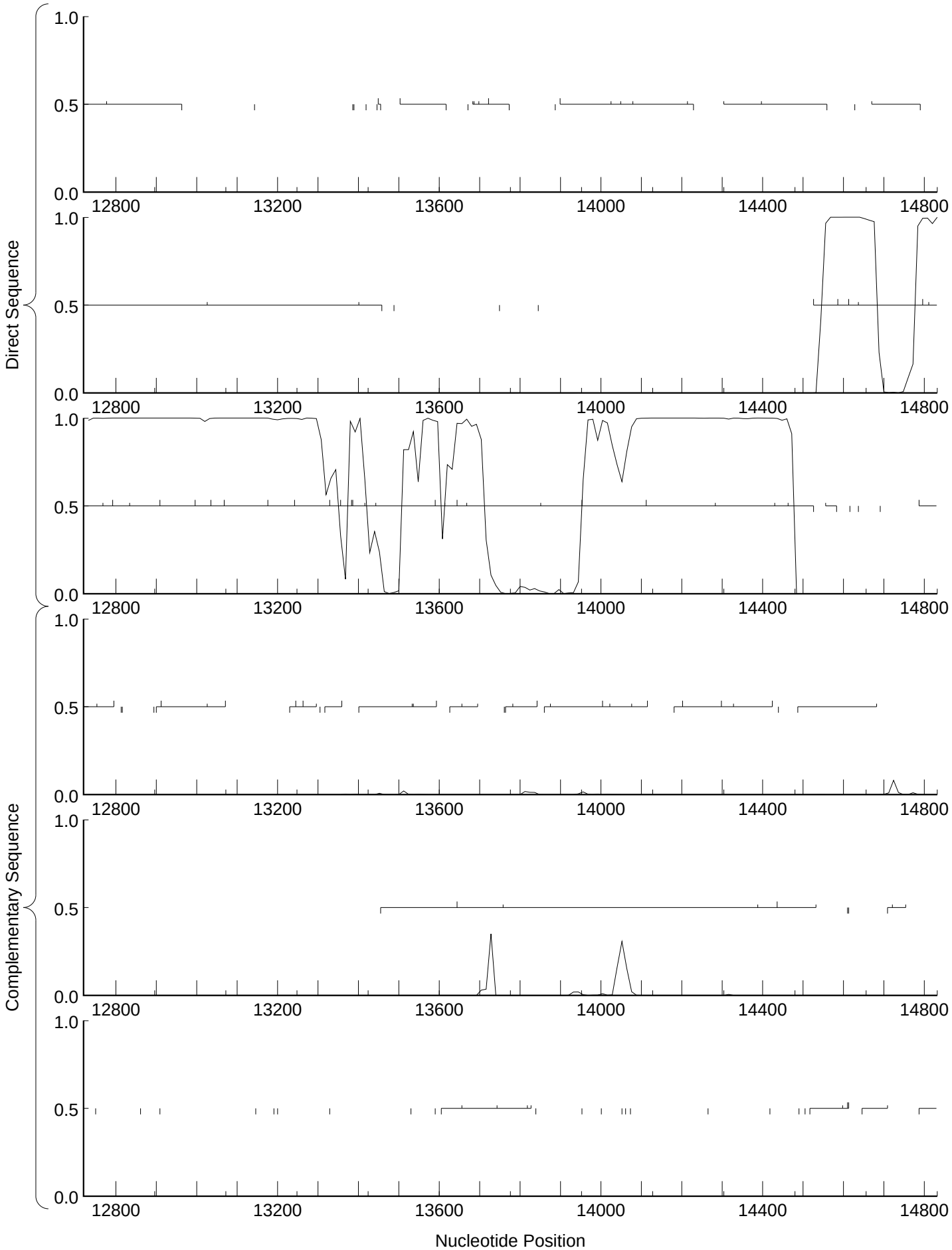
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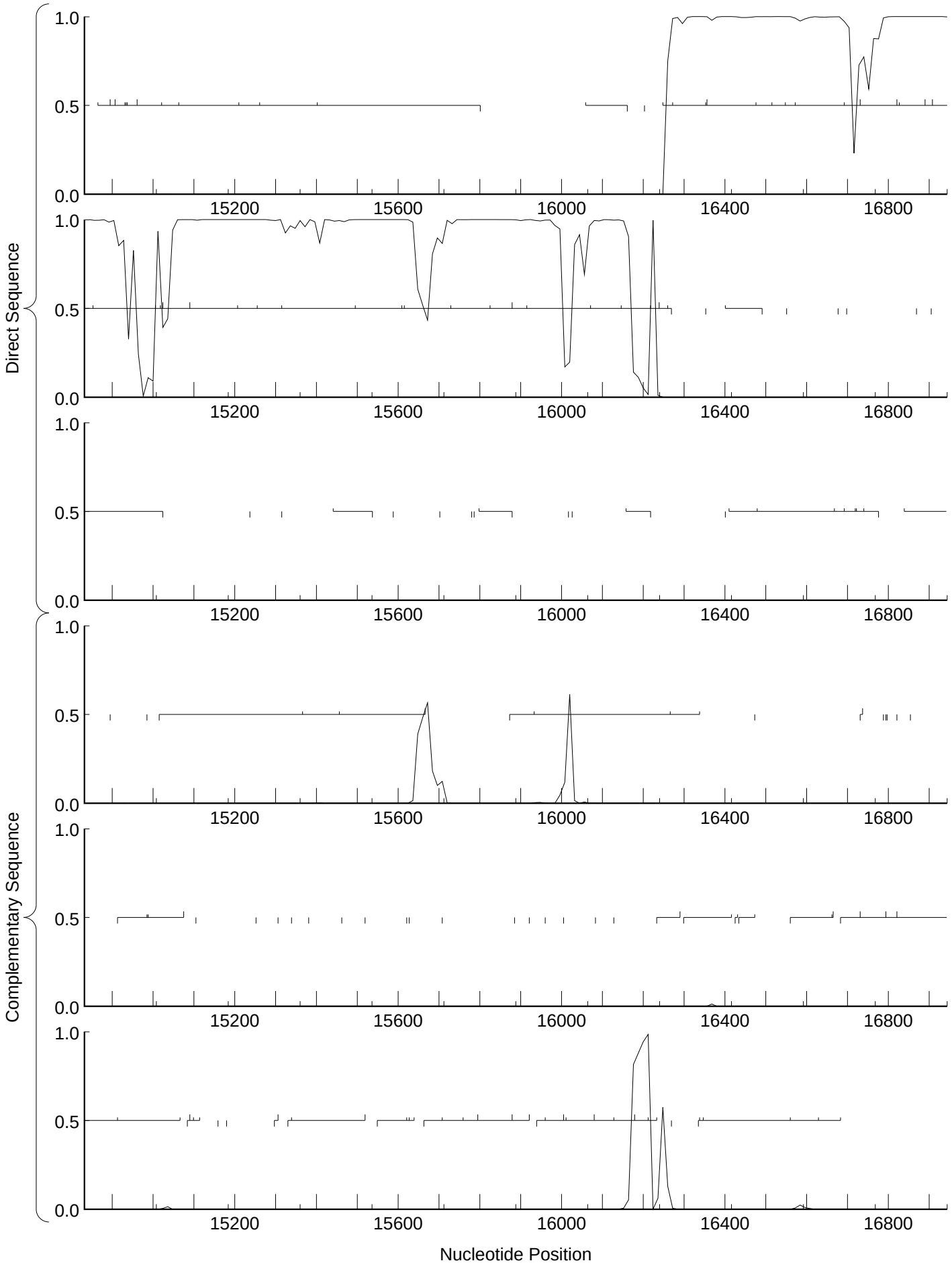
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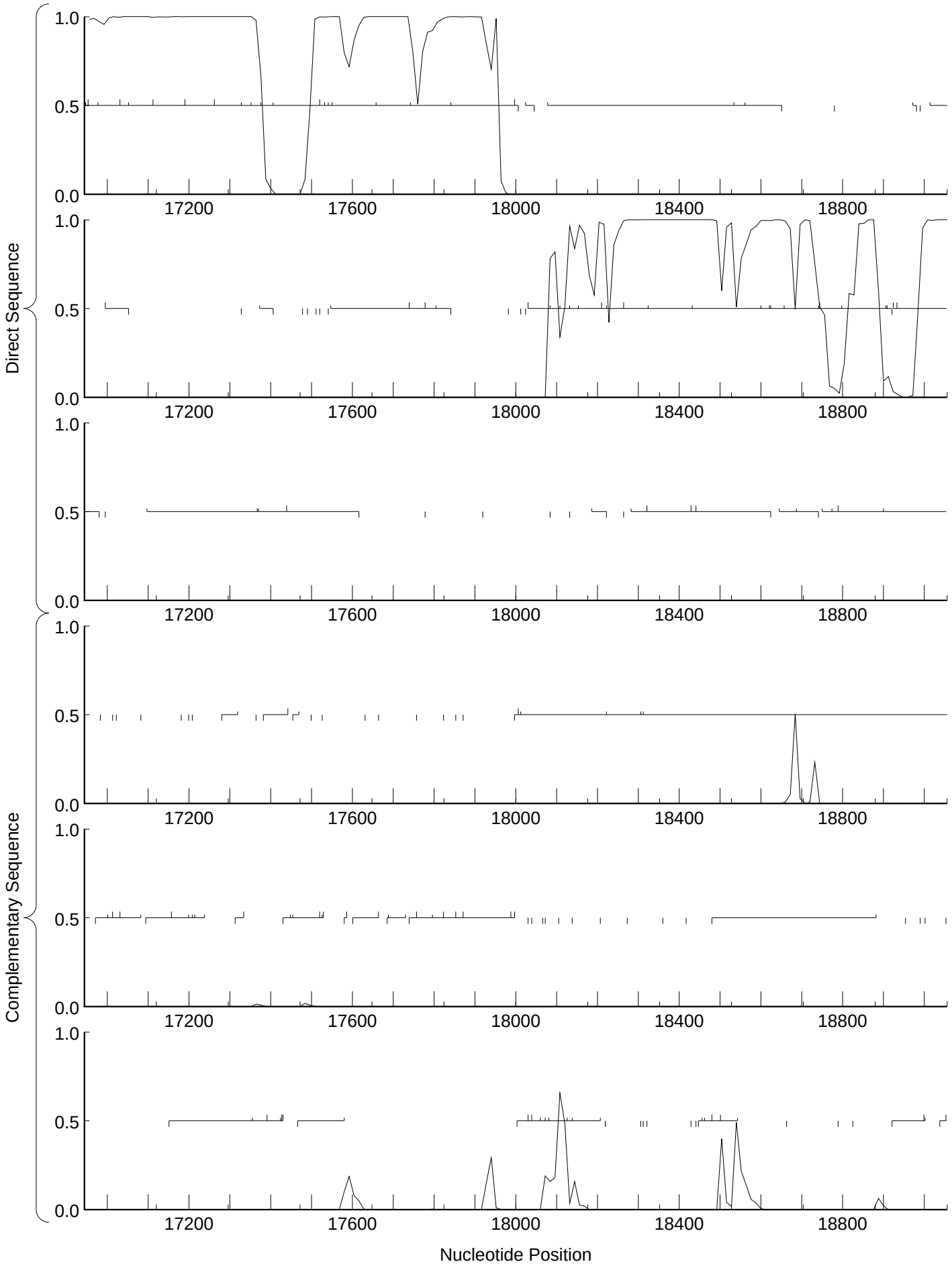
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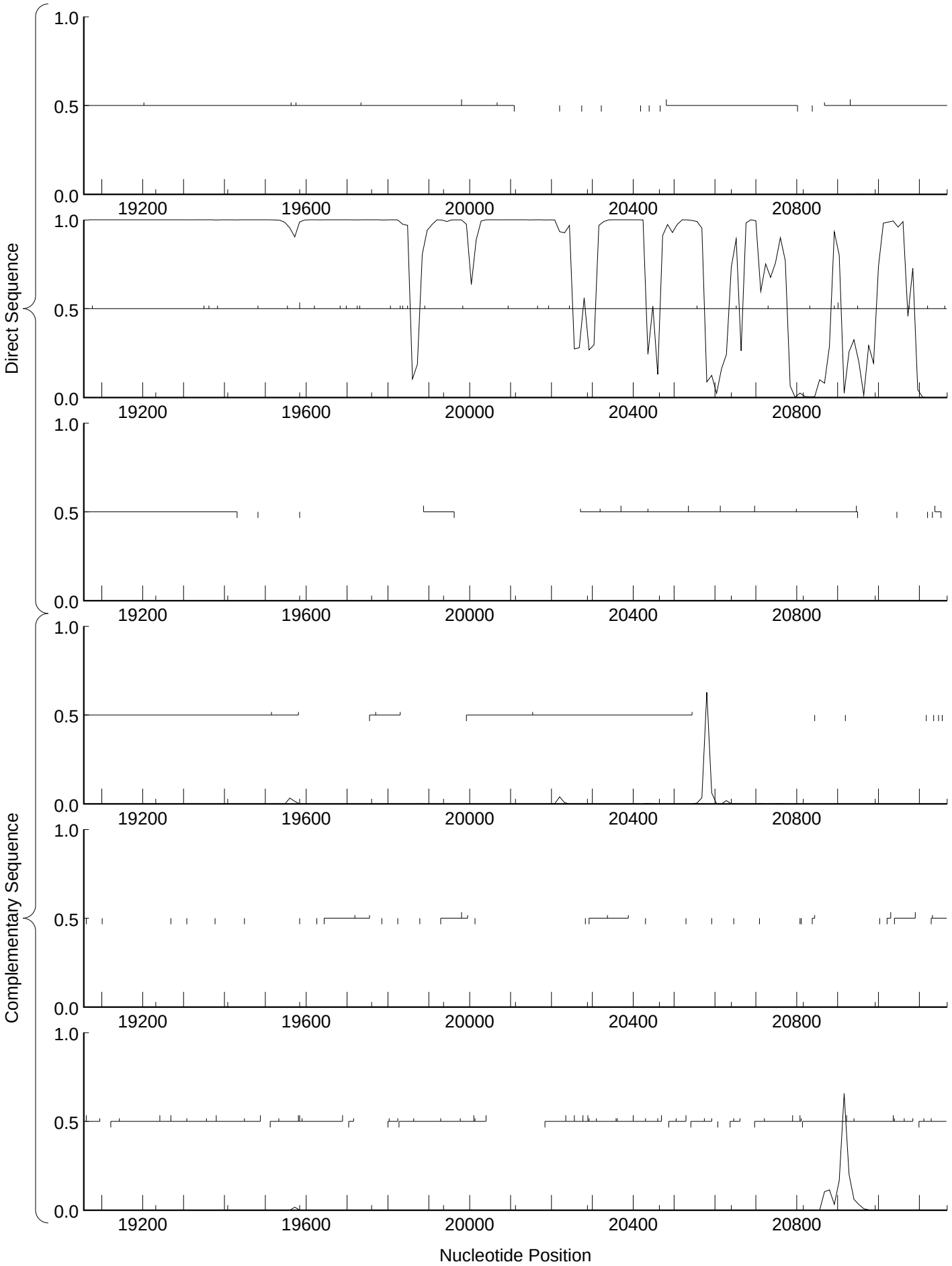
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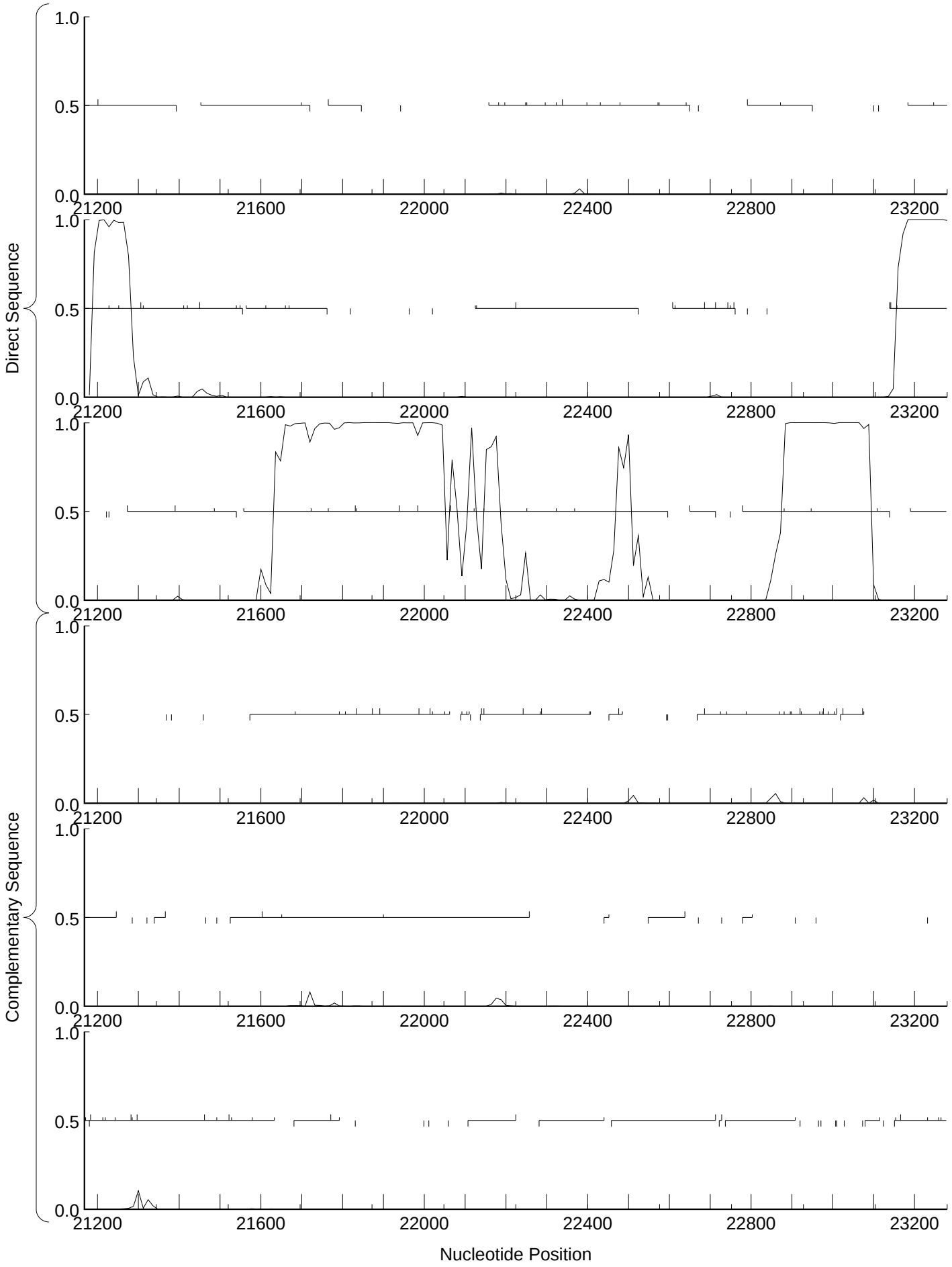
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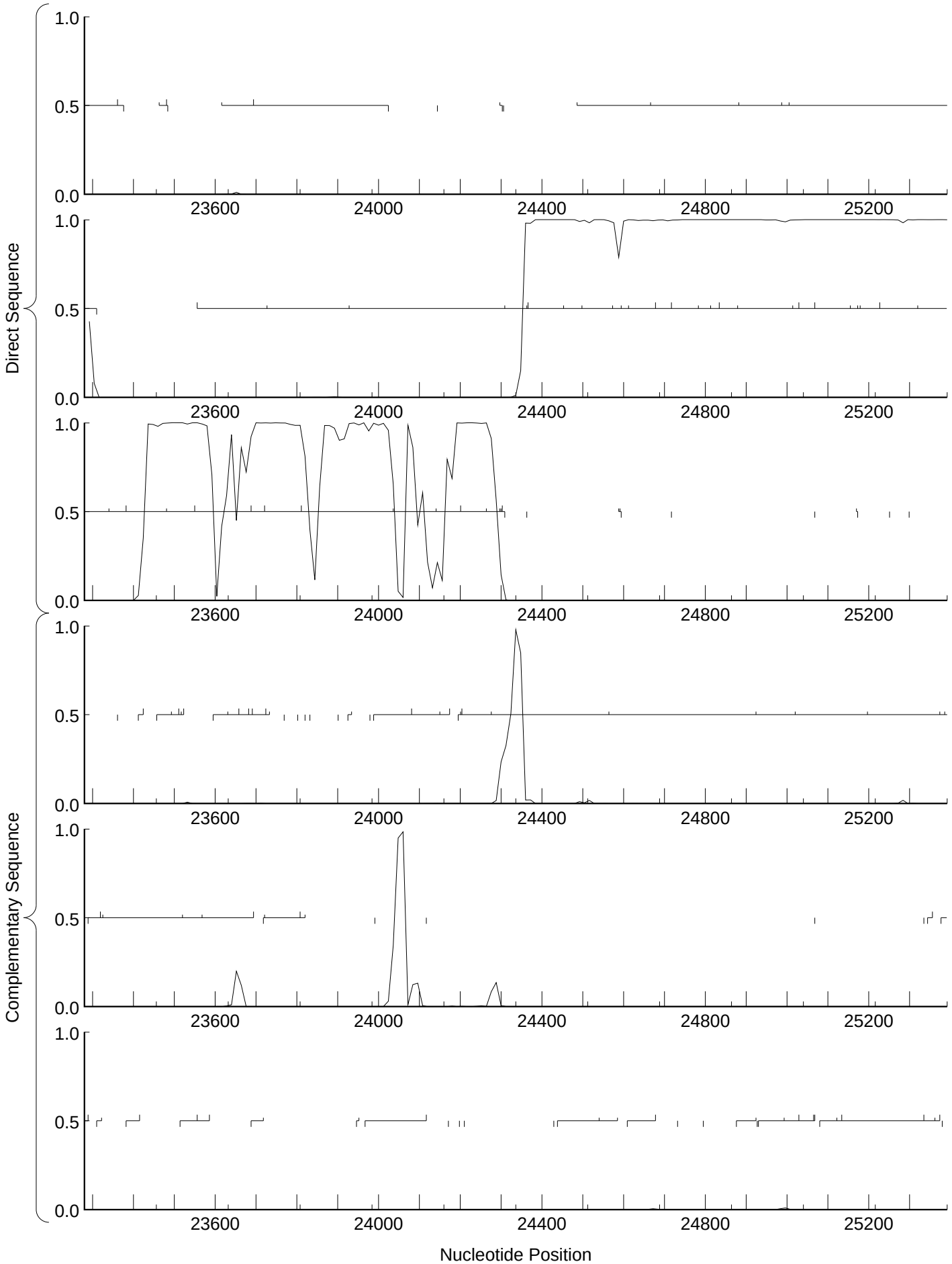
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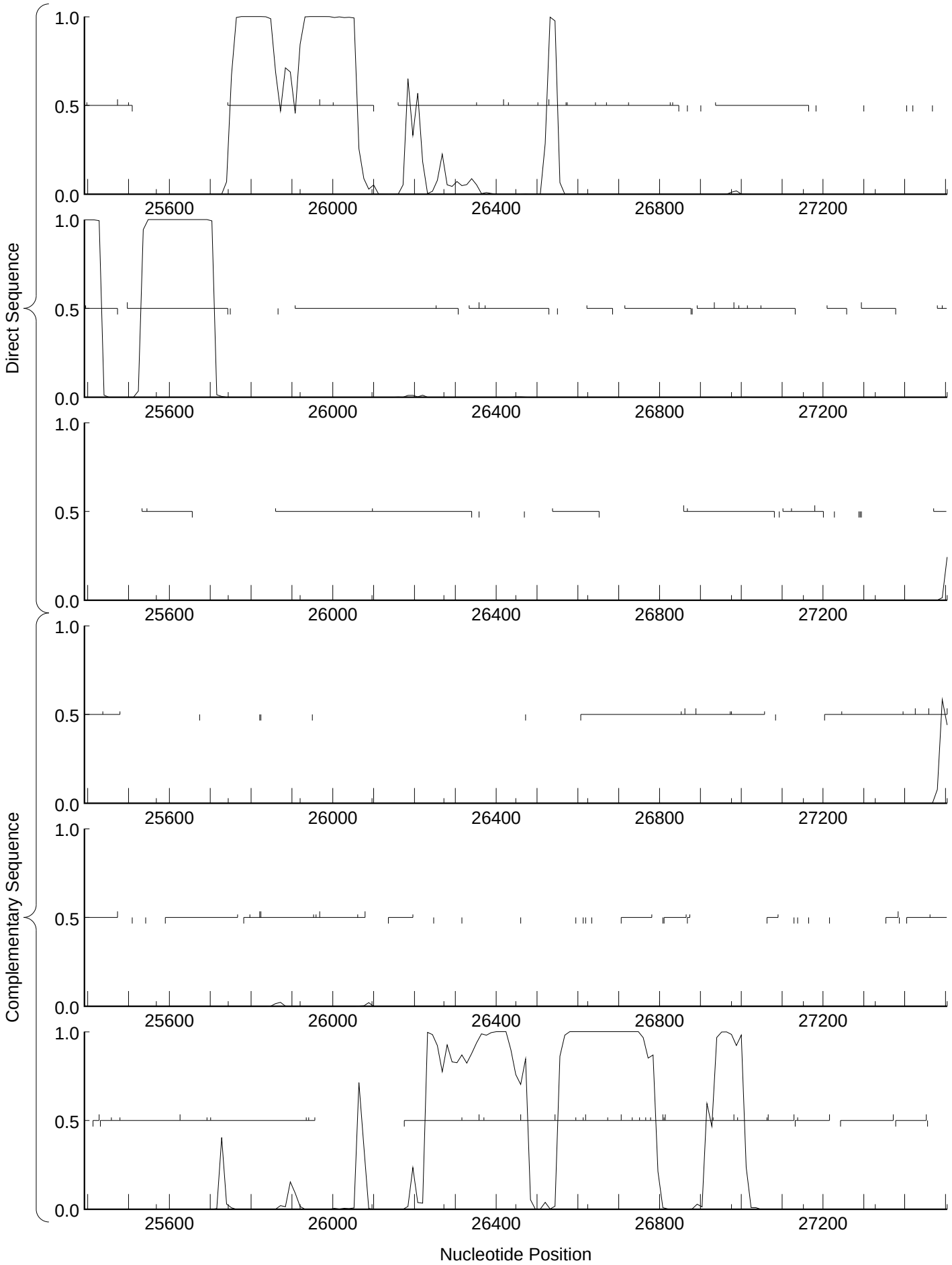
plete sequence, 48916 bp including 13-base 3’ overhang (CCTGCCGCCCCGA), Cluster P1, Order 4, Window 96, Step 12, 12/25

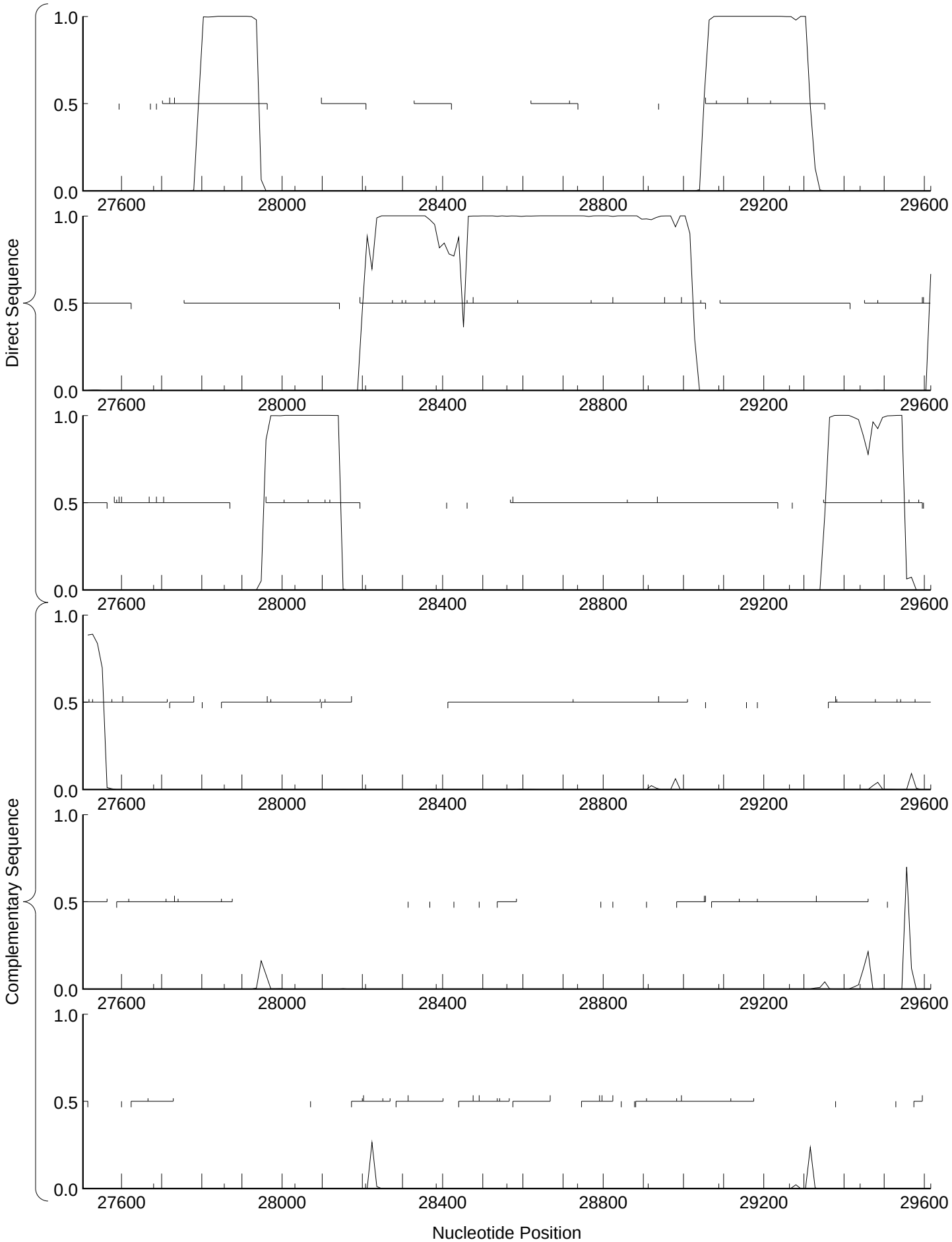


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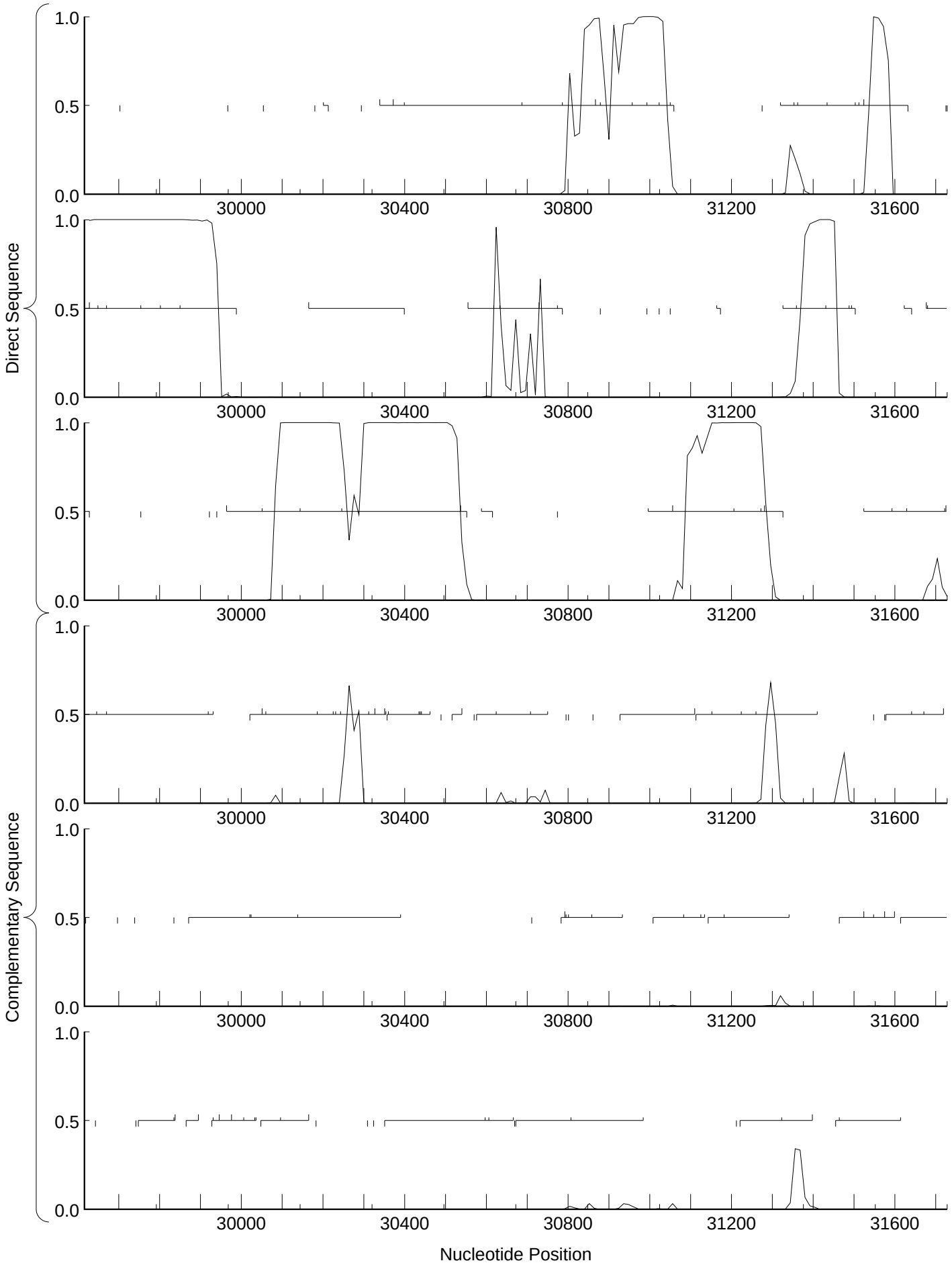


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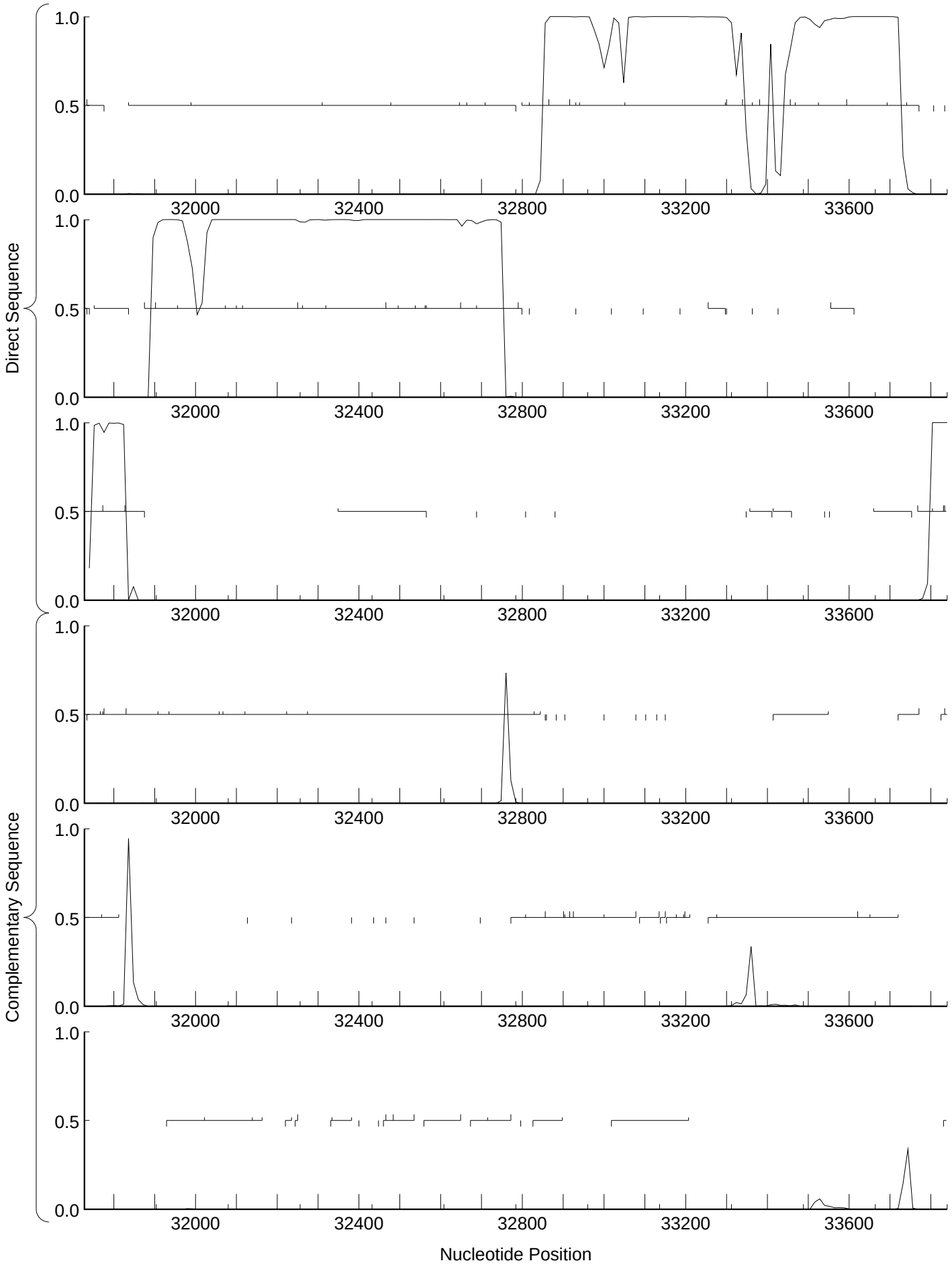



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
27600
28000
28400
28800
29200
29600
Nucleotide Position

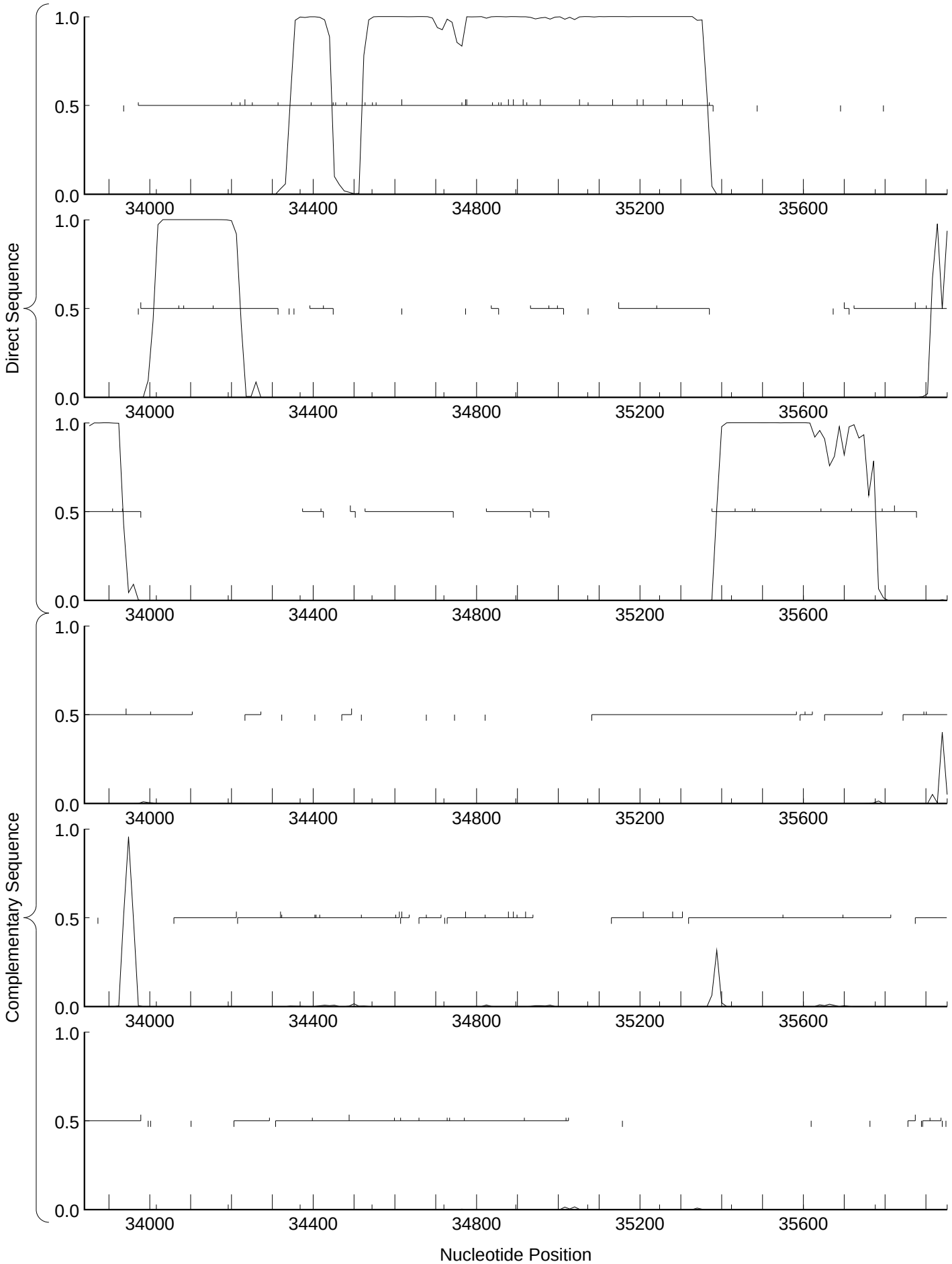
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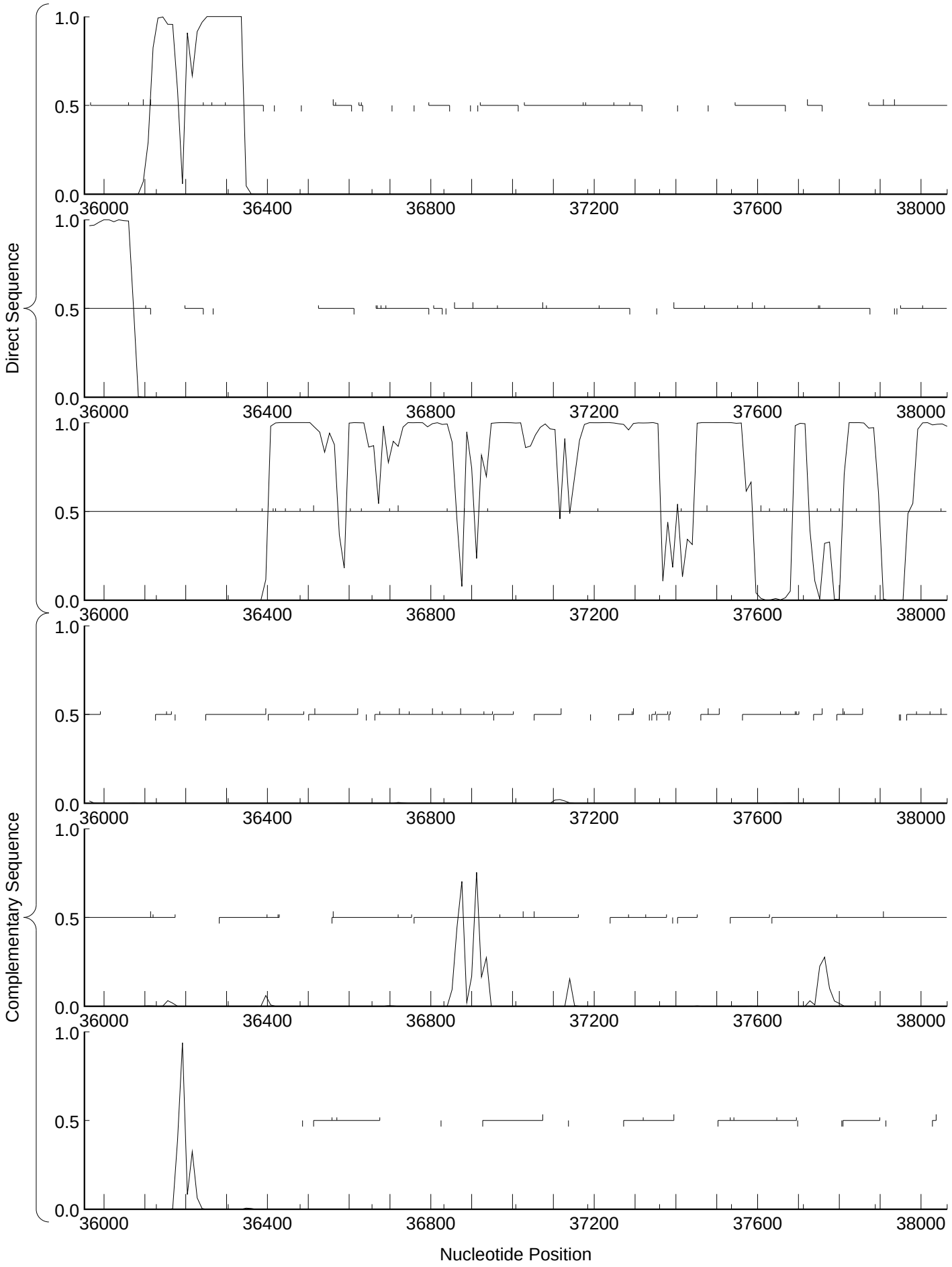
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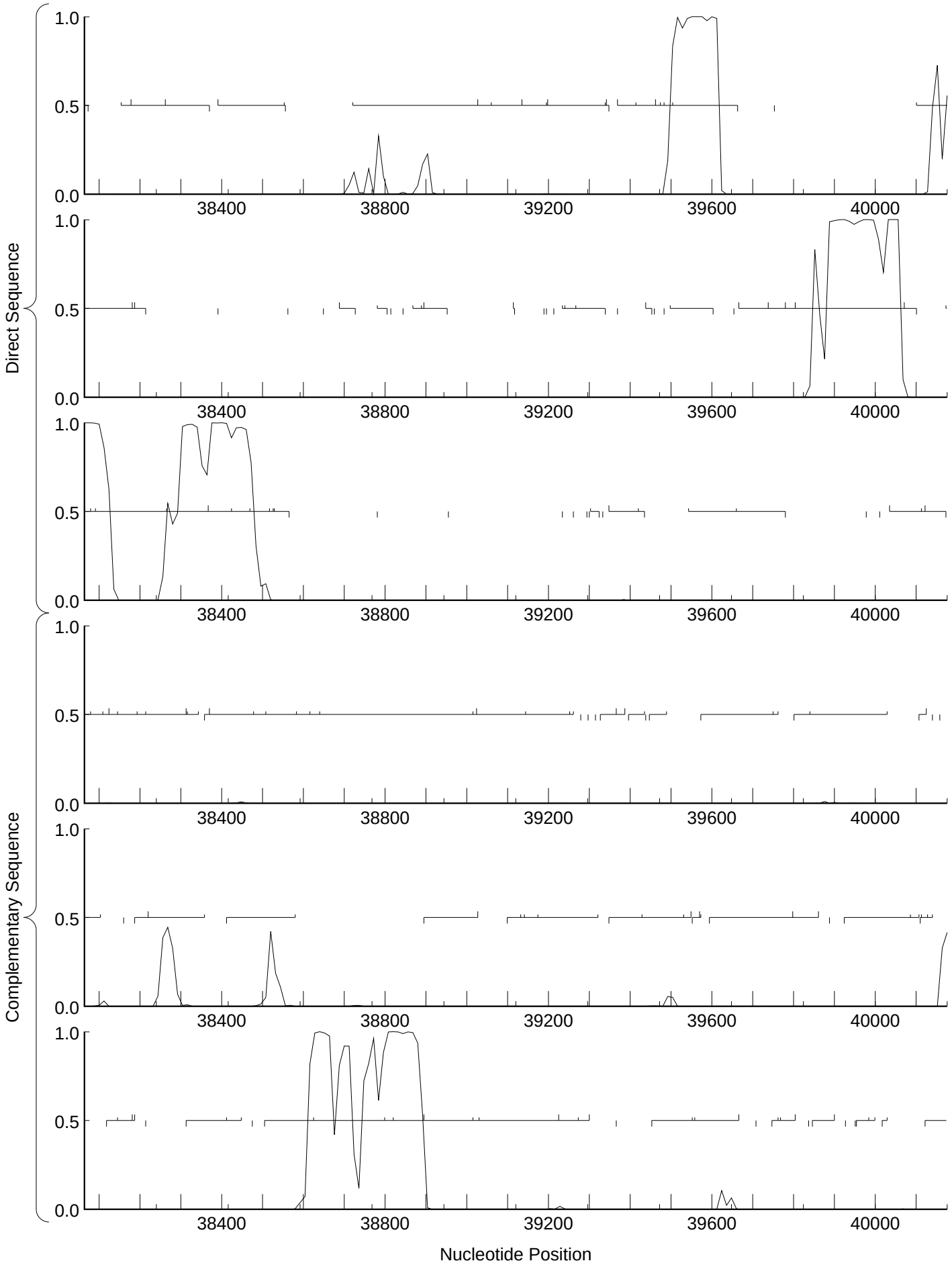


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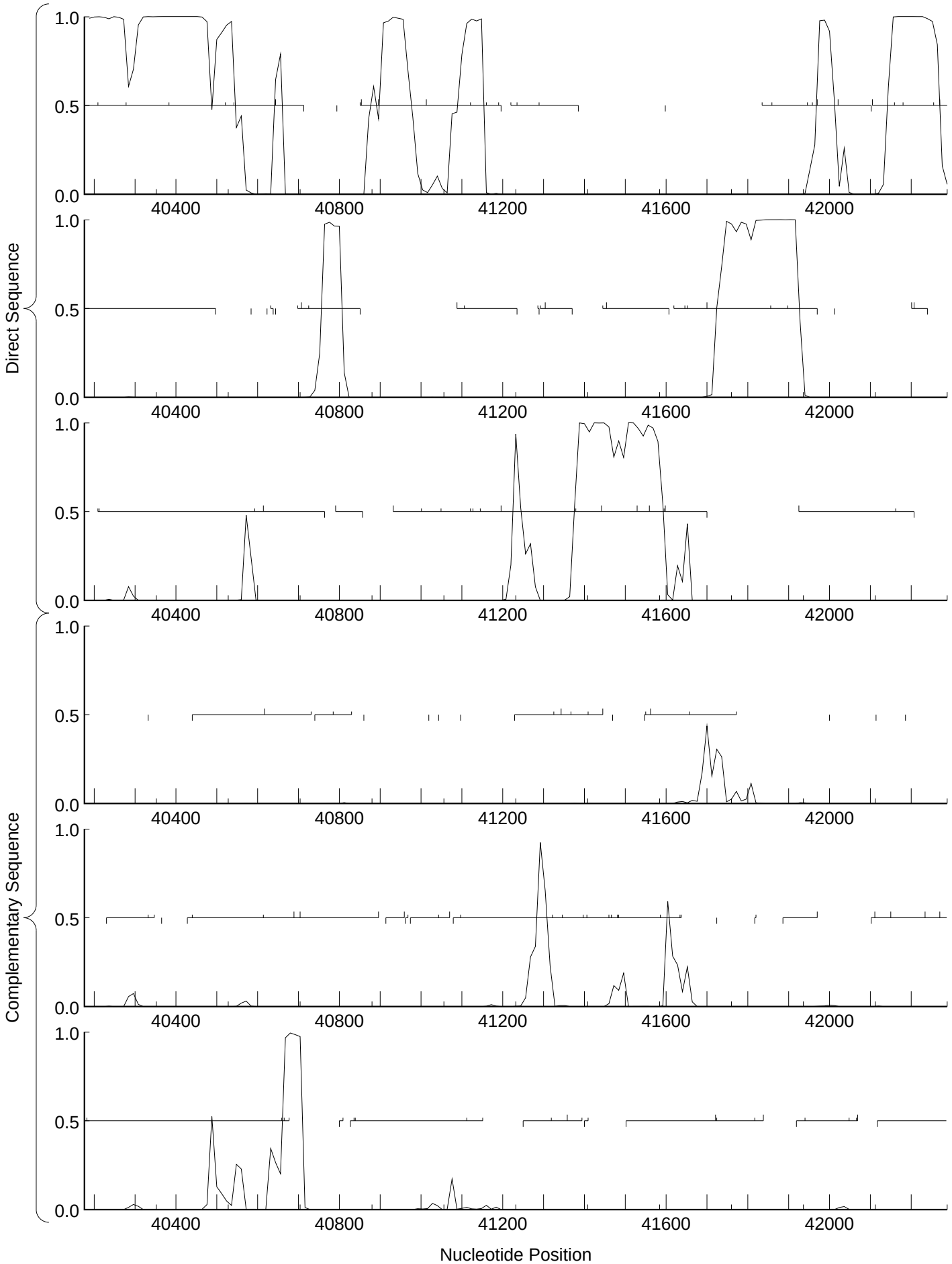


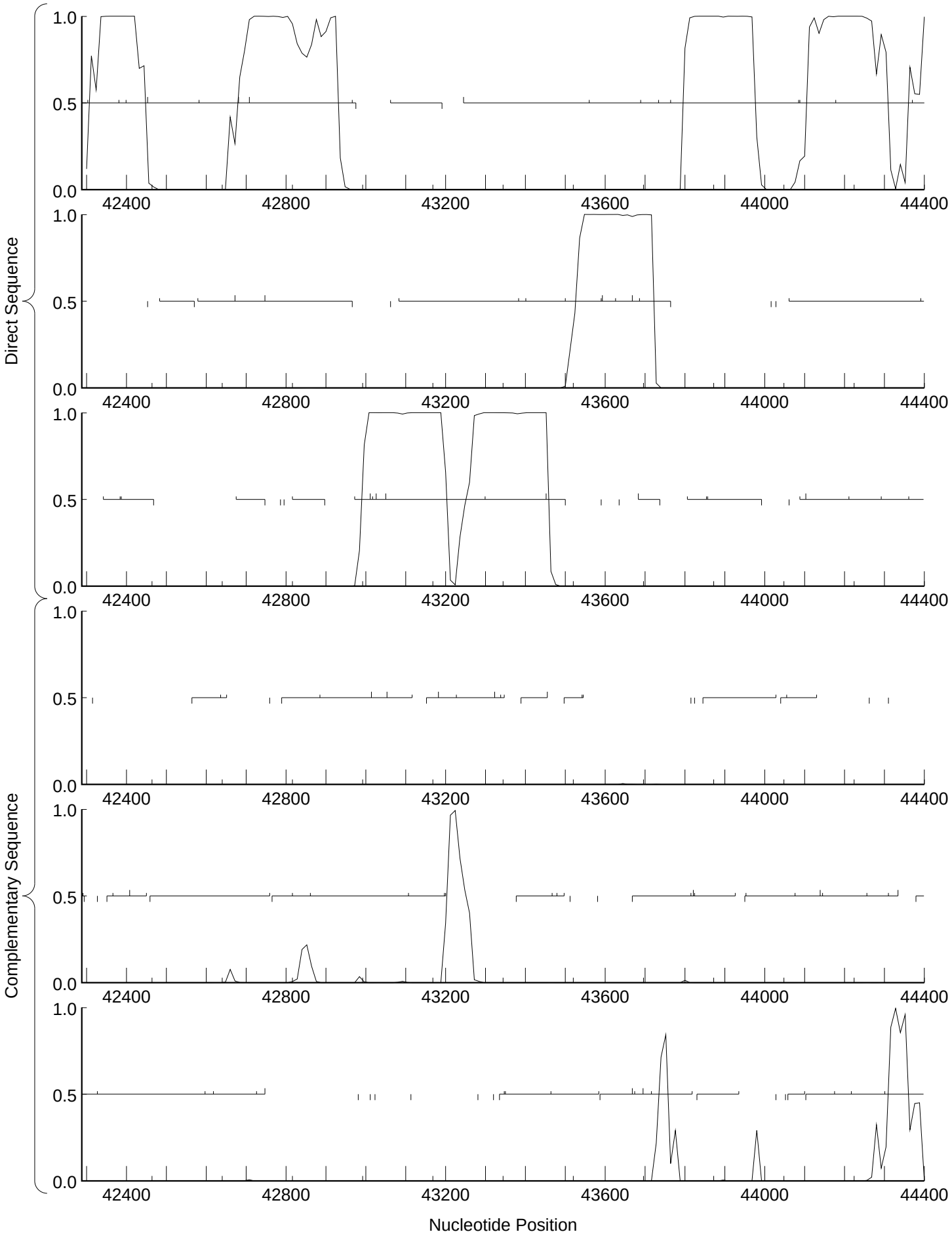
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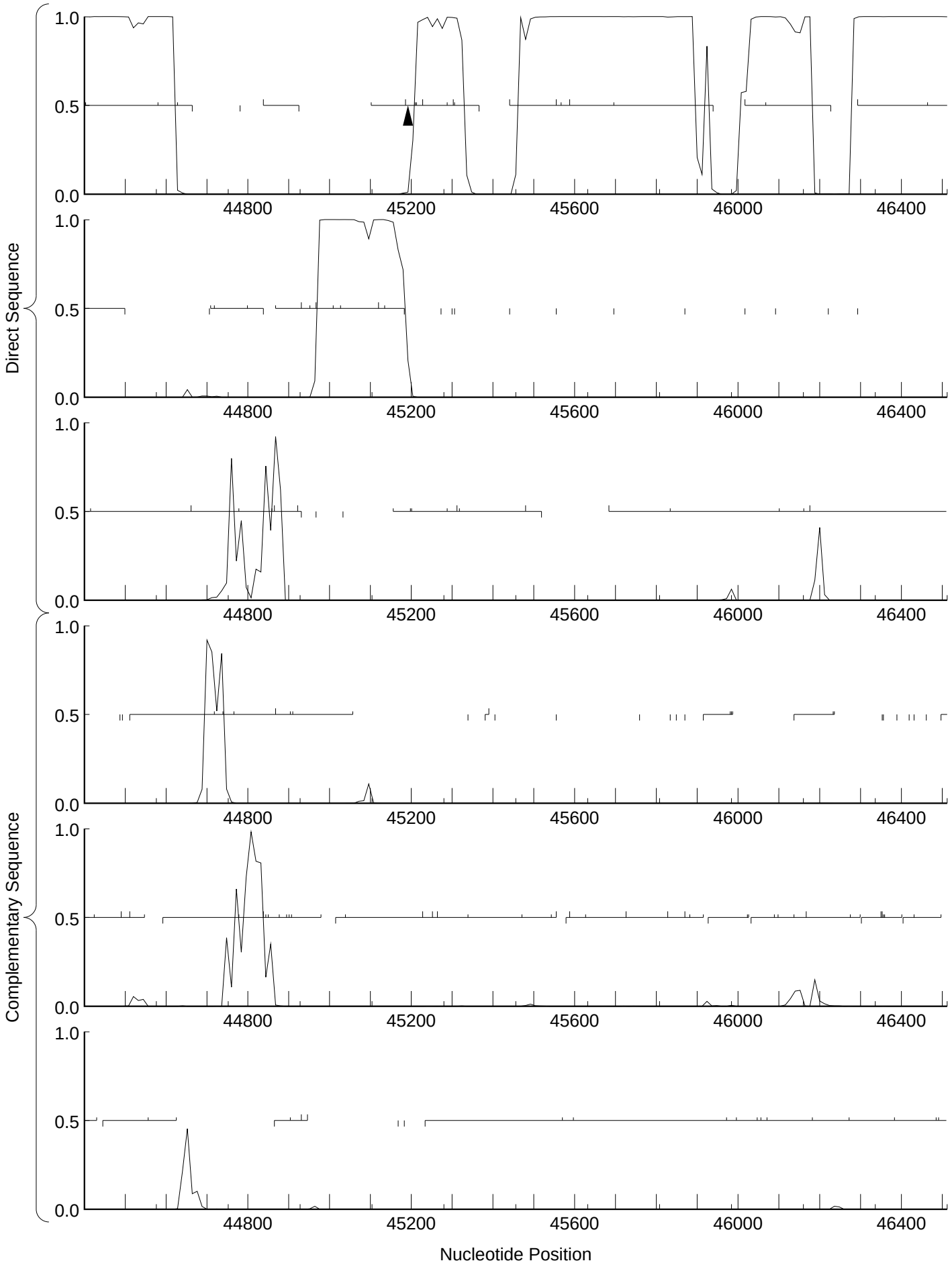

Direct Sequence
Complementary Sequence
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0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

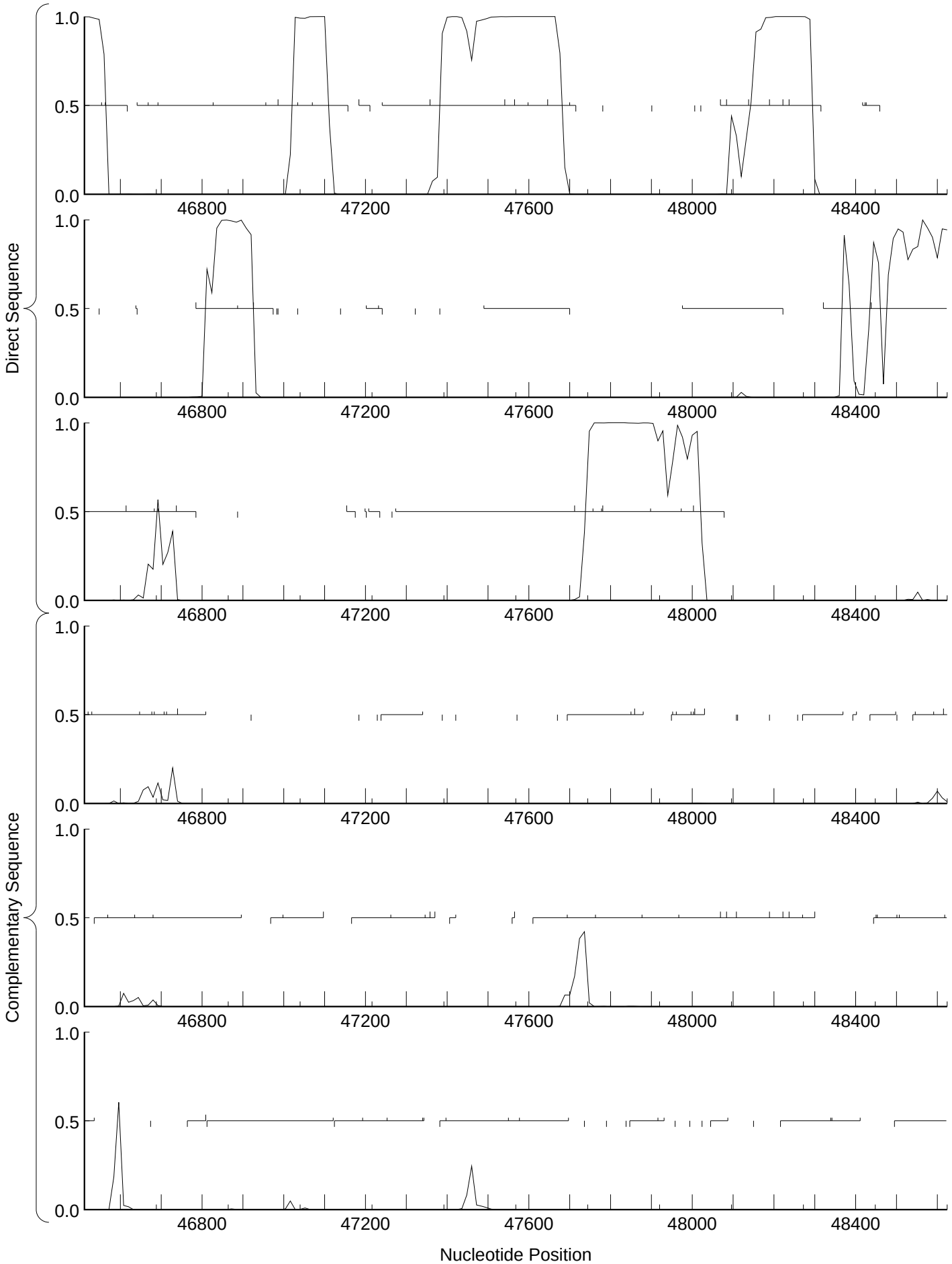
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Direct Sequence
Complementary Sequence
1.0
0.5
0.0
42400
42800
43200
43600
44000
44400
Nucleotide Position

plete sequence, 48916 bp including 13-base 3’ overhang (CCTGCCGCCCCGA), Cluster P1, Order 4, Window 96, Step 12, 23/25




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
46800
47200
47600
48000
48400
Nucleotide Position

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