GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage SnapTap complete sequence, 53051 bp including 10-base 3’ overhang (CGGT
Analysis Date    :  3/28/24 at 18:44:44
Pages            :  27
Sequence Length  :  53051 bp
GC Content       :   63.45%

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

PS-Version       :  1.2

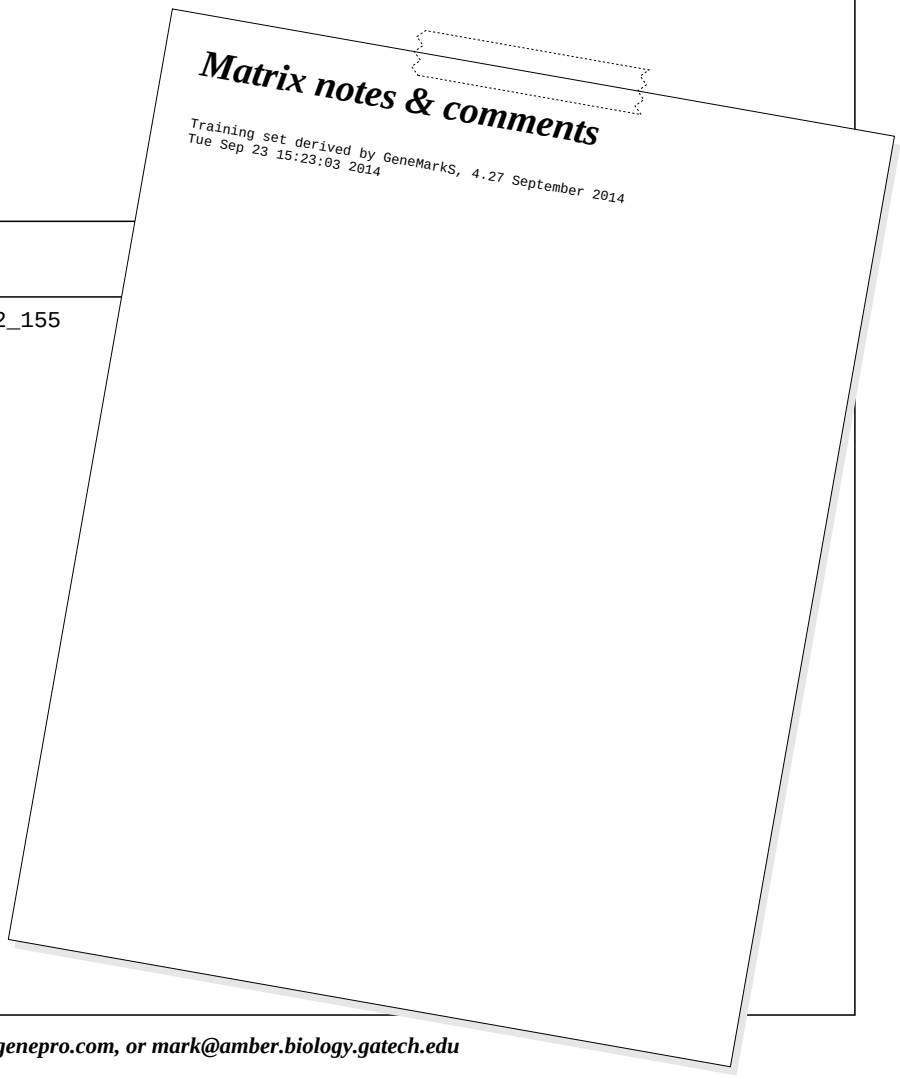
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Author     :  -

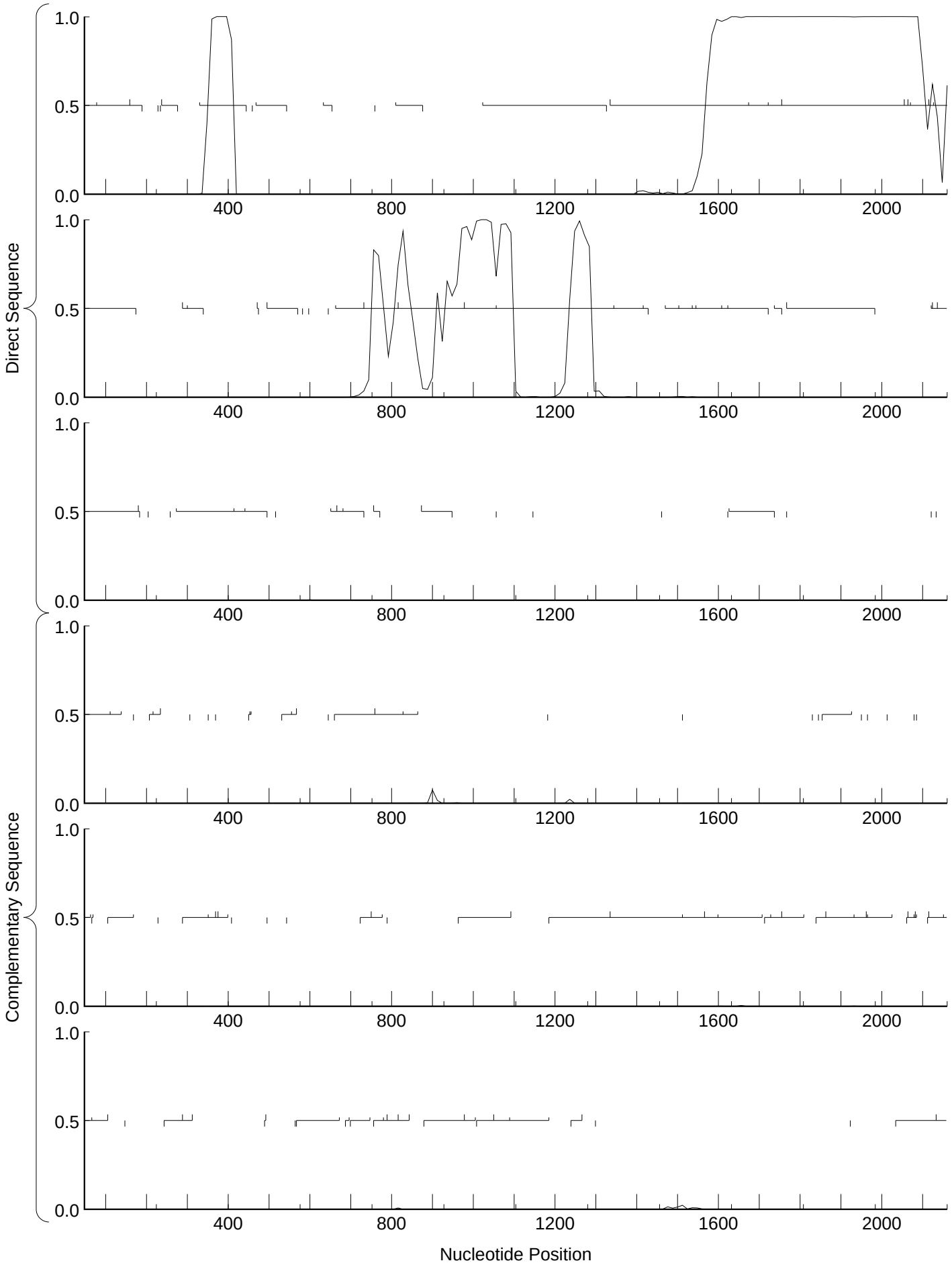
Order      :  4


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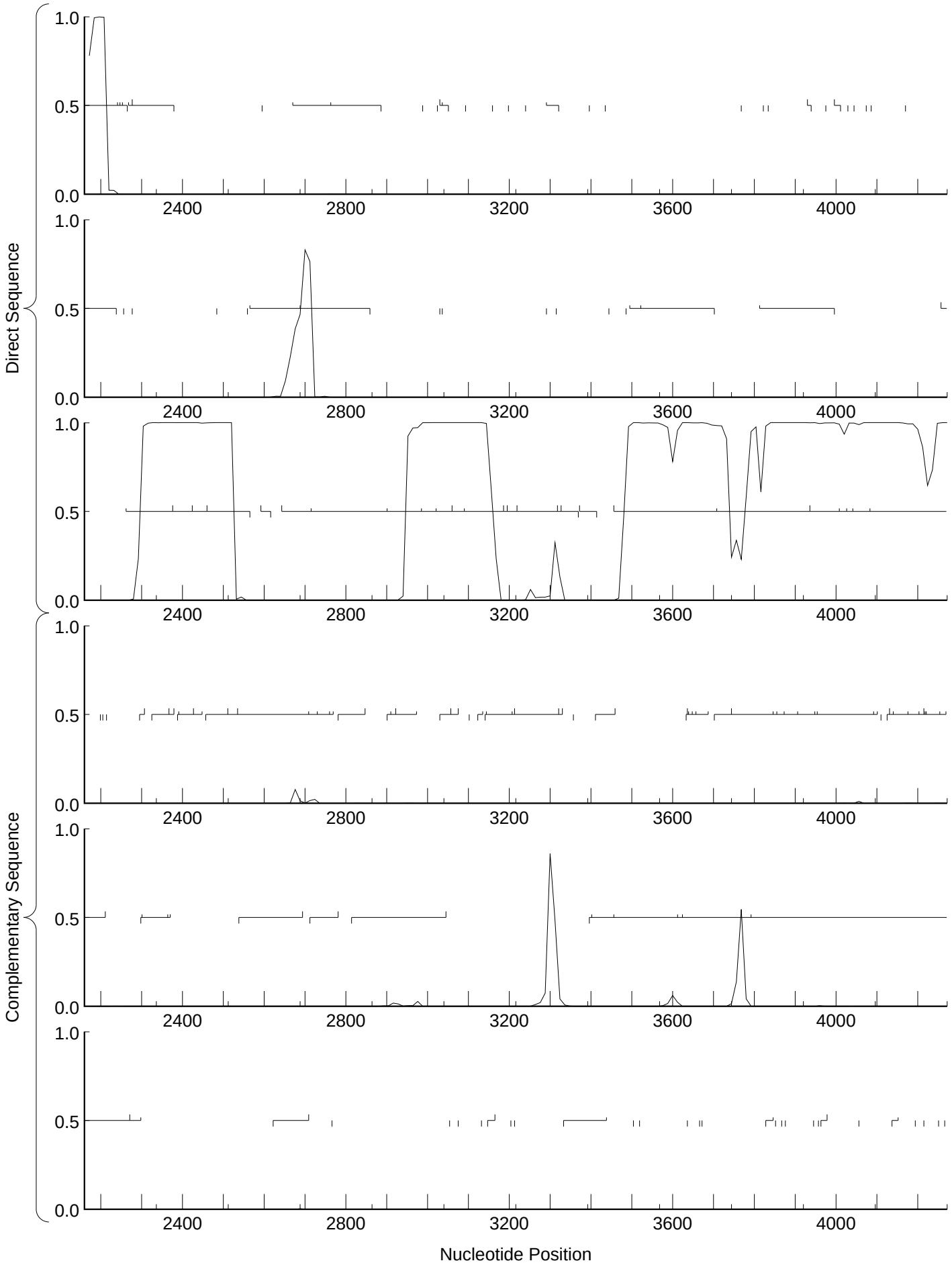


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

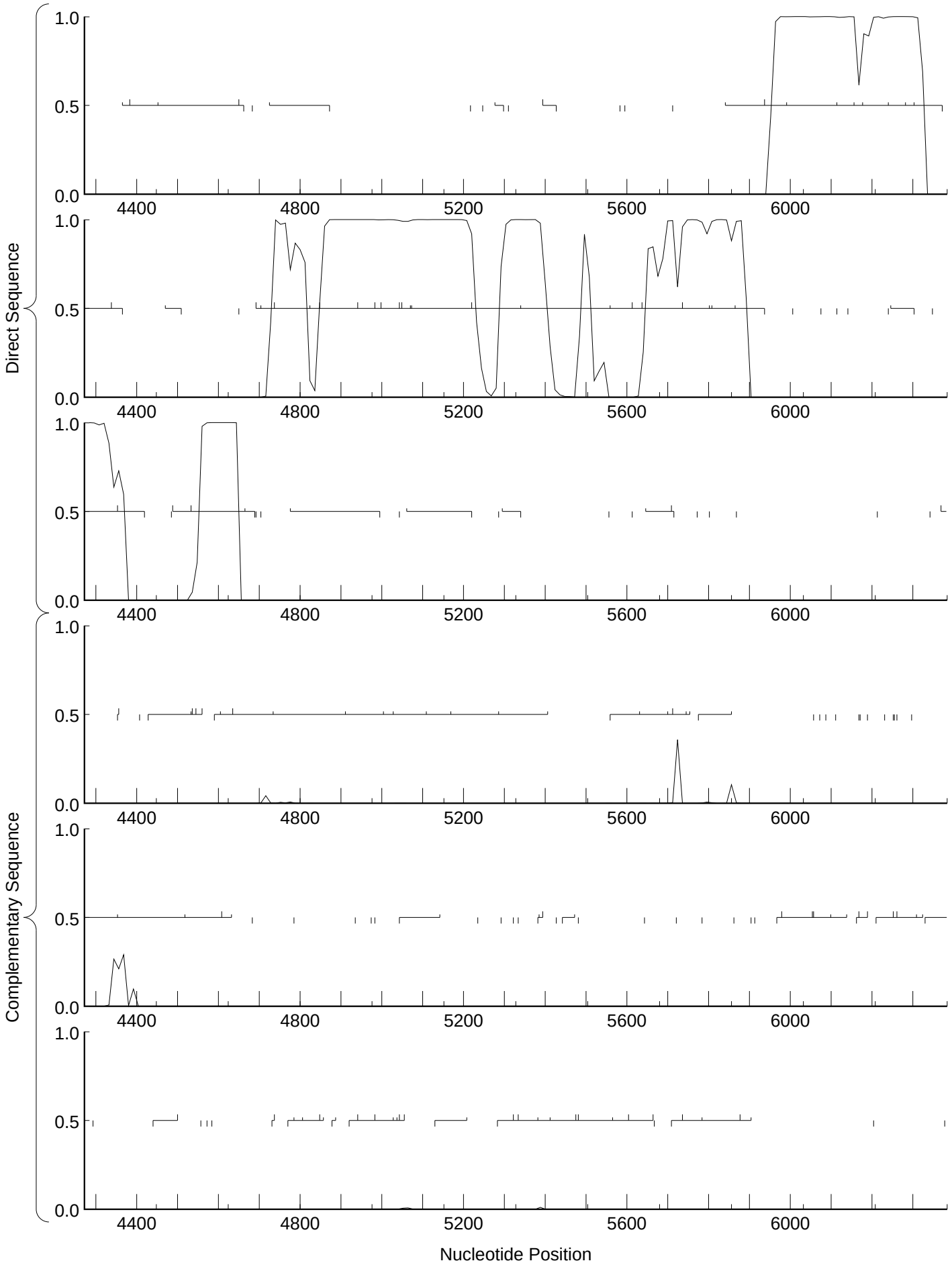
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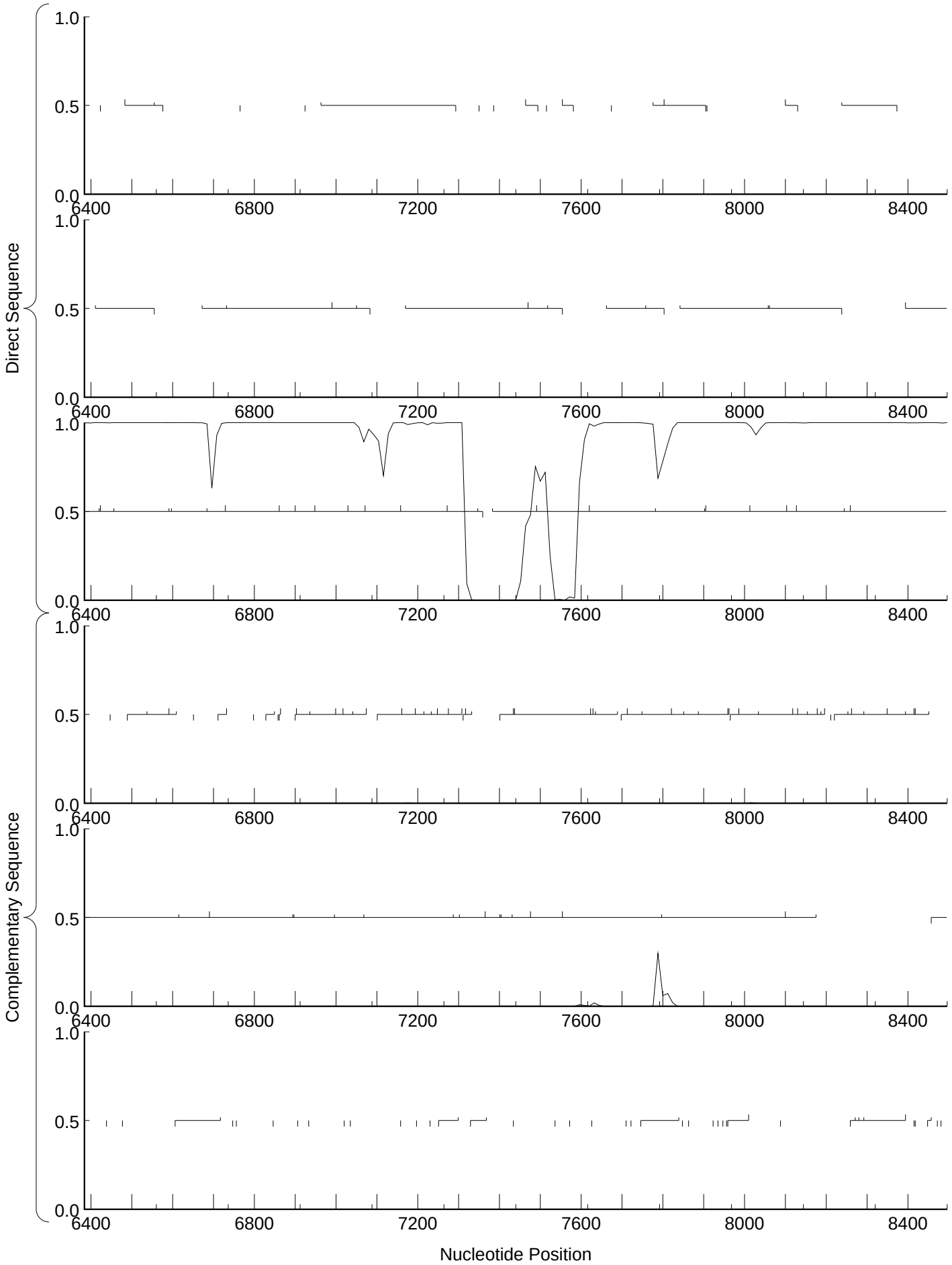
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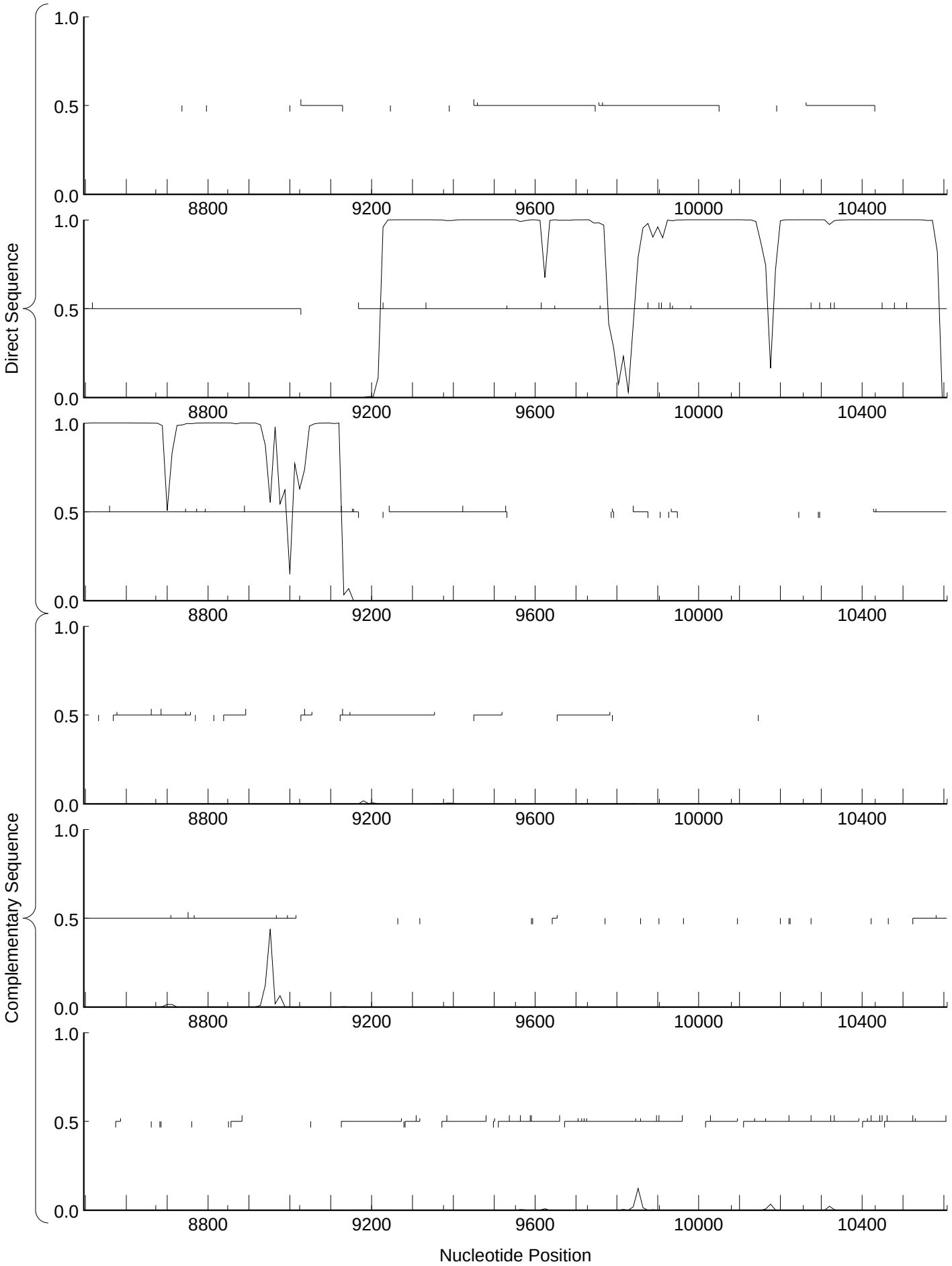
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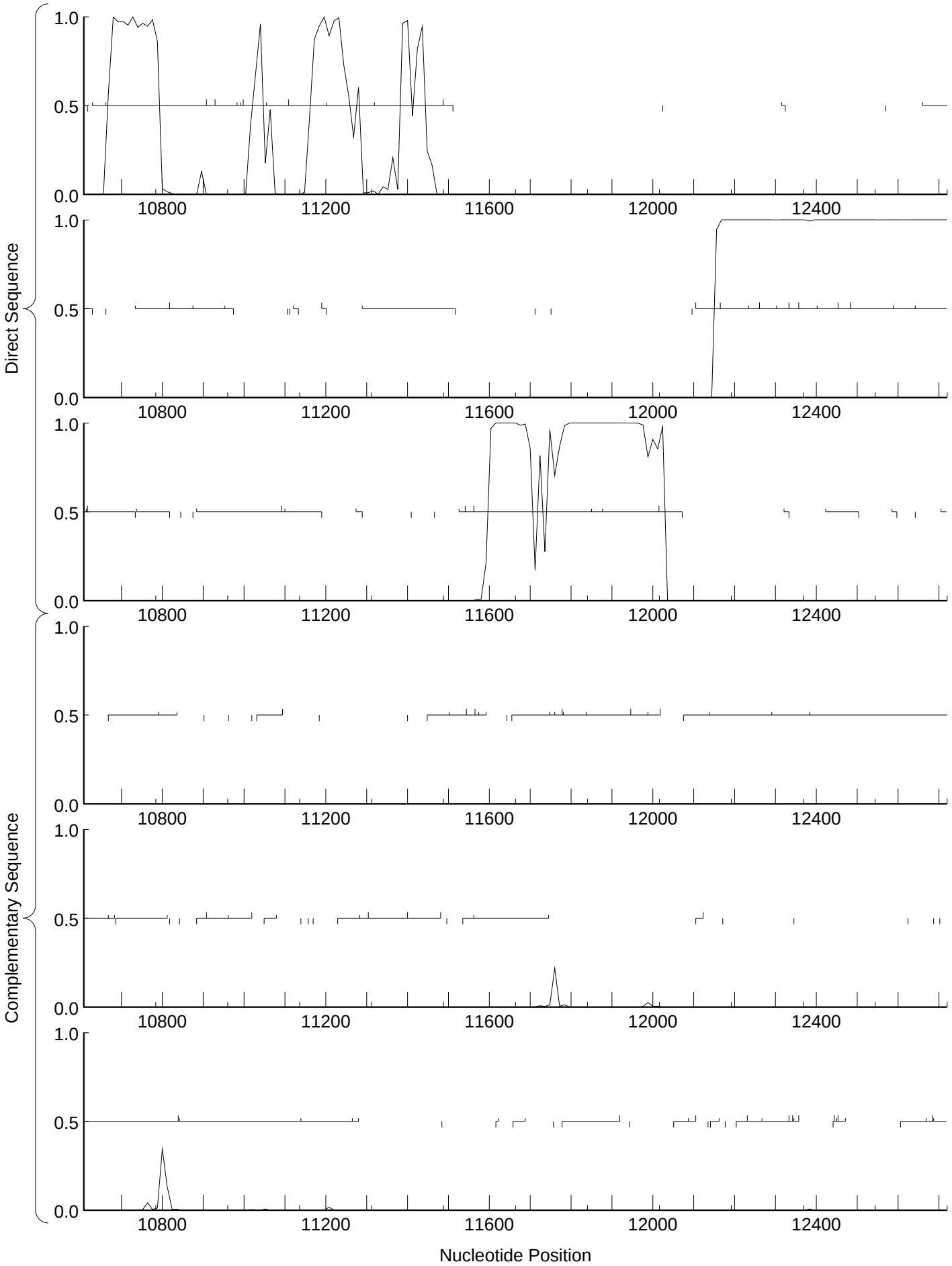
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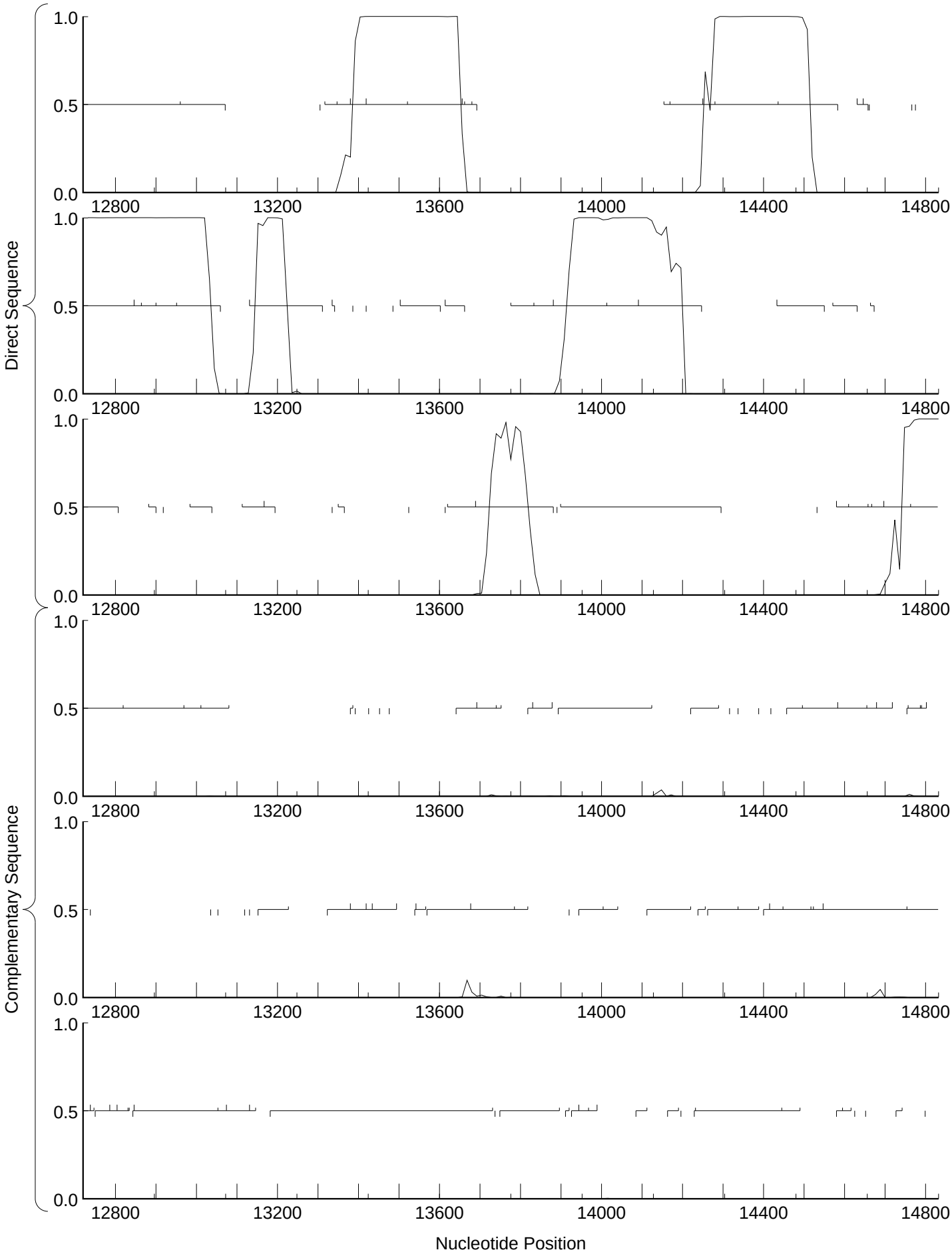
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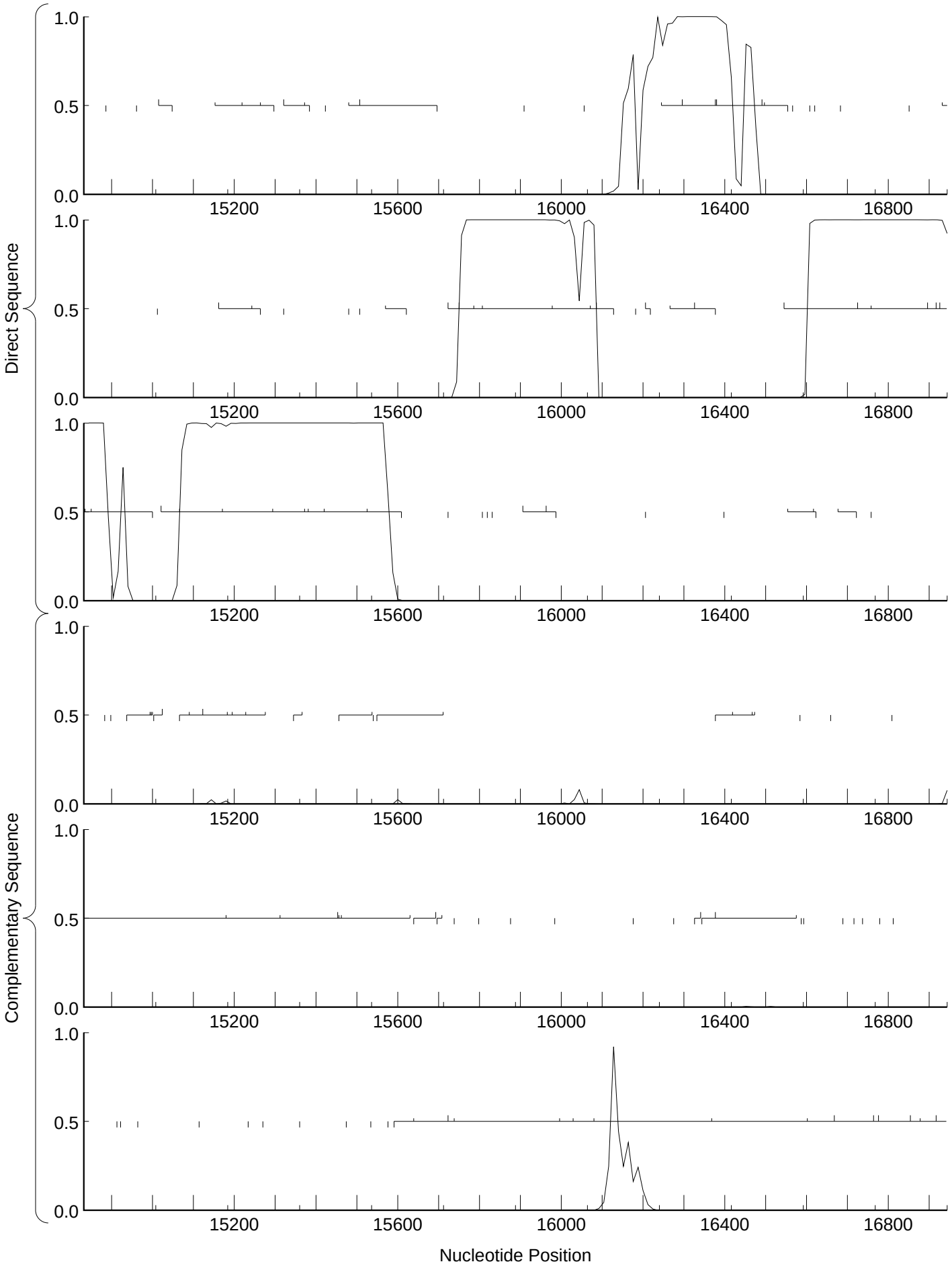
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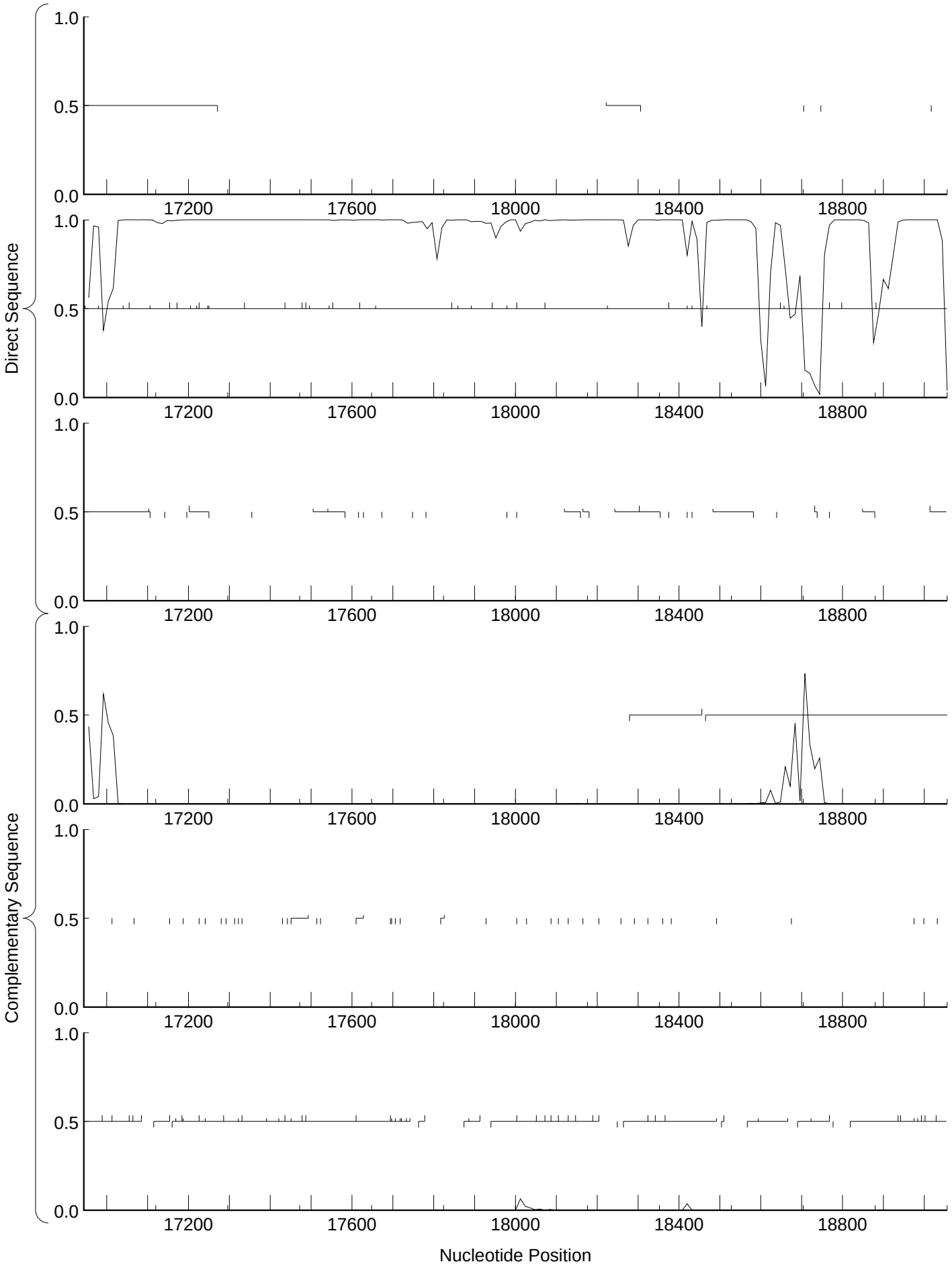


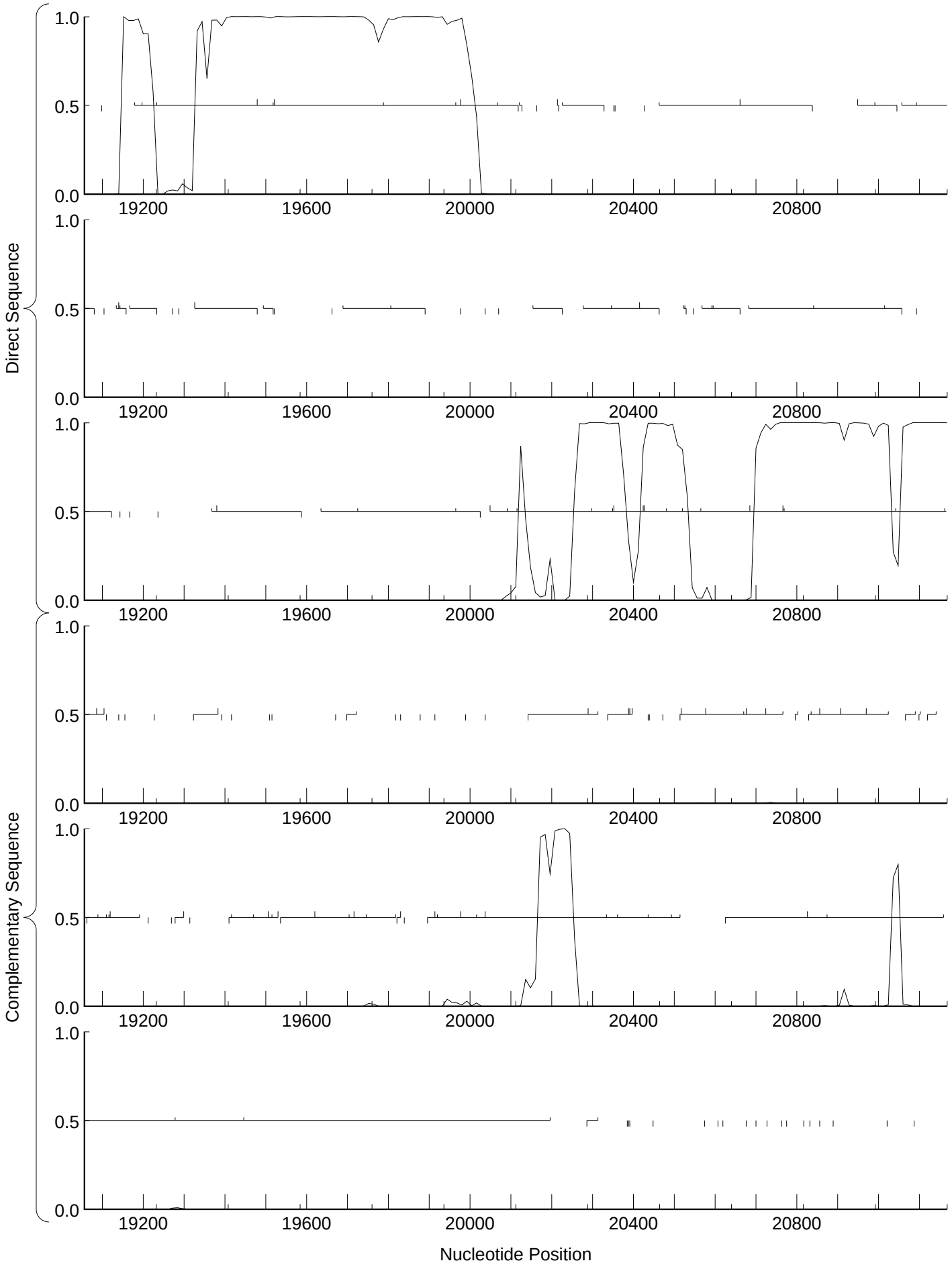
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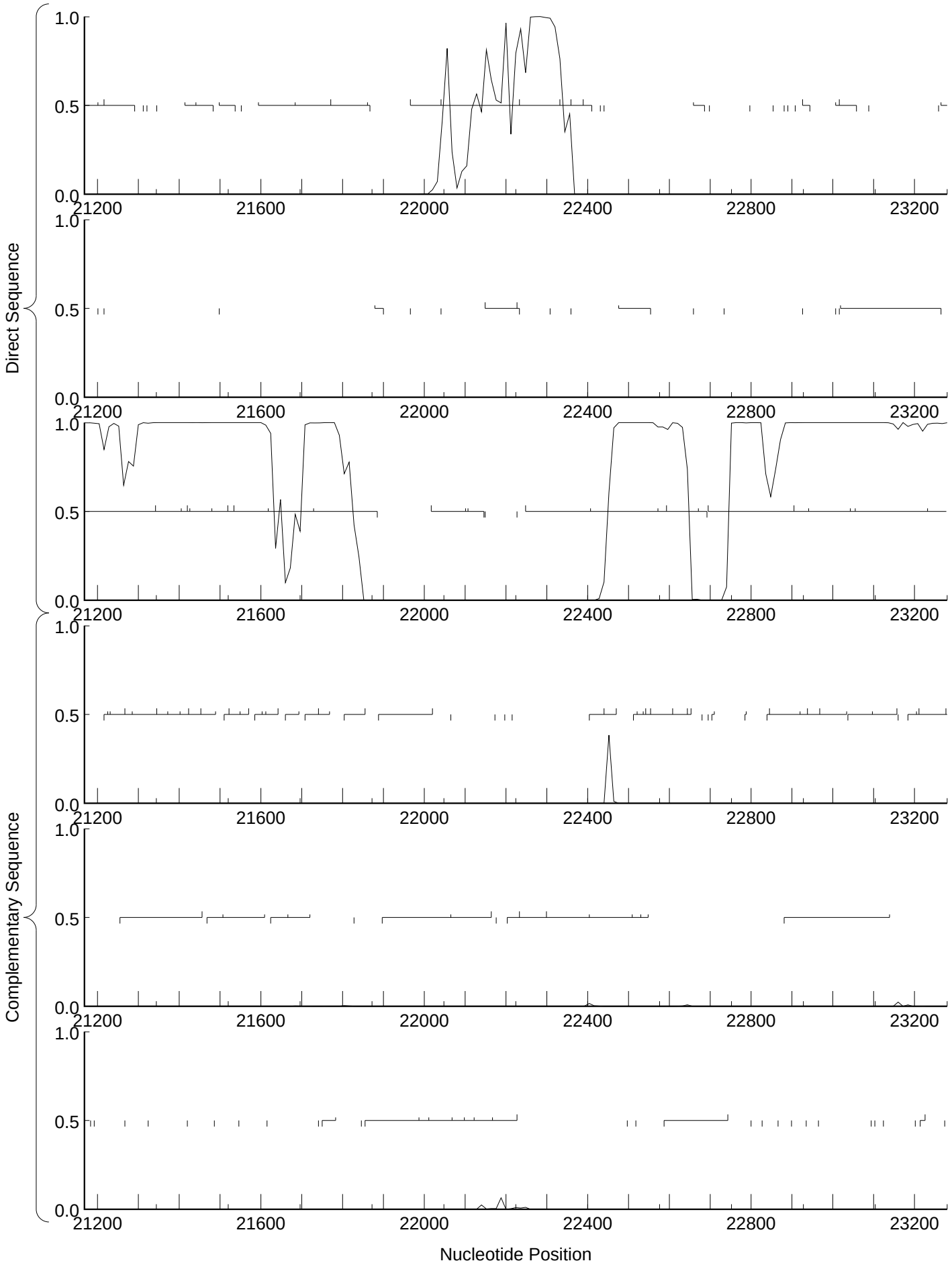


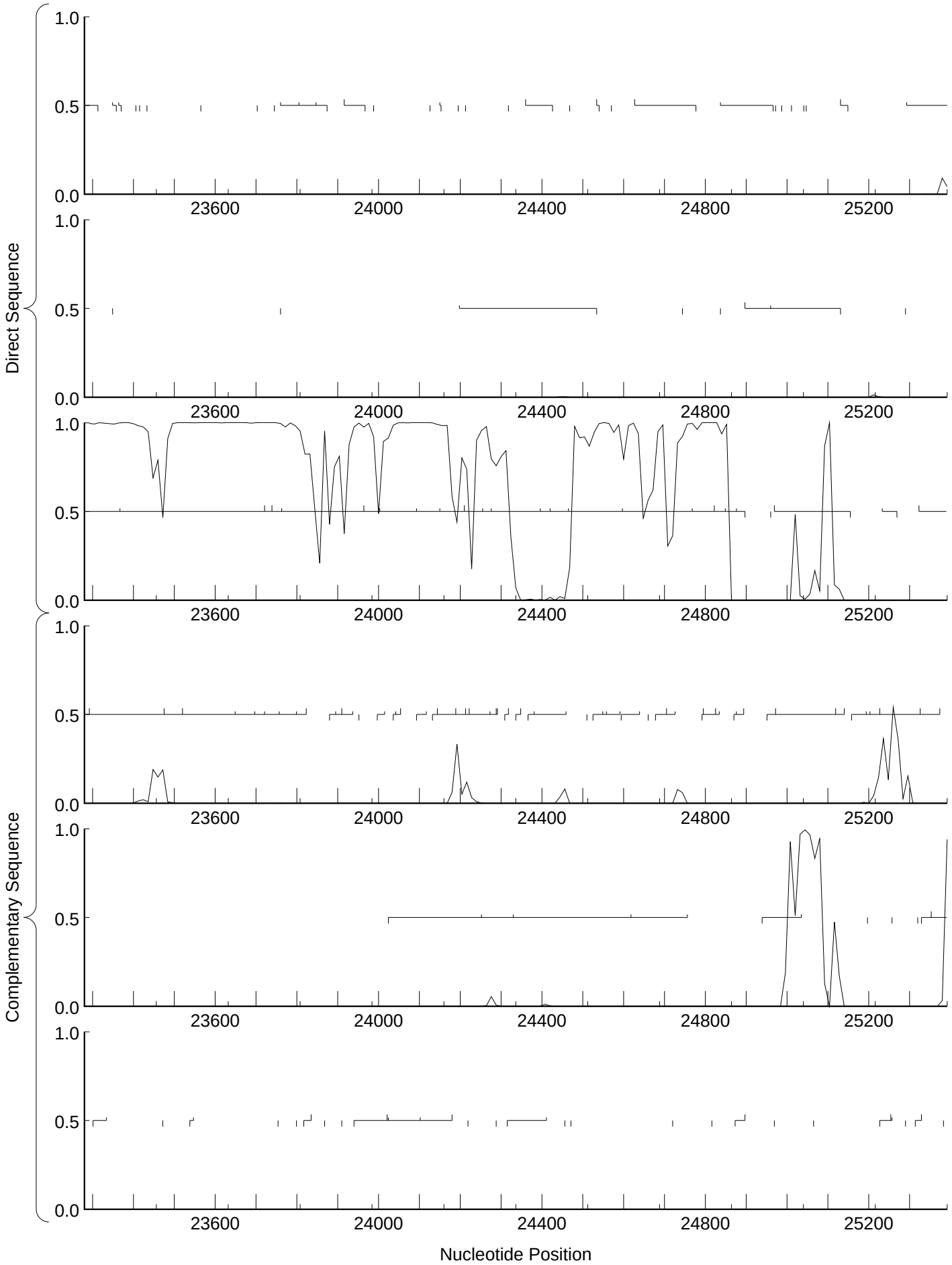
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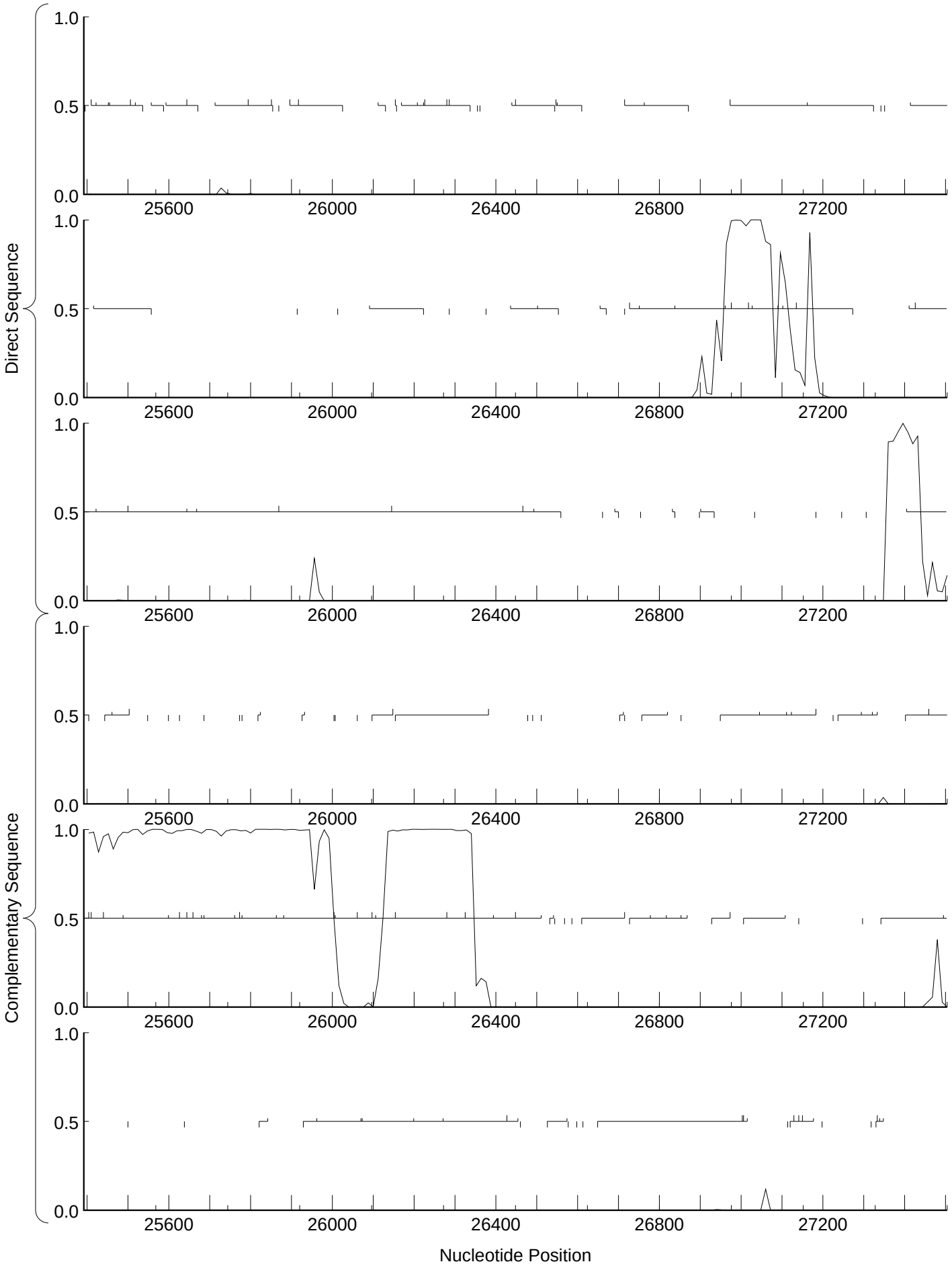


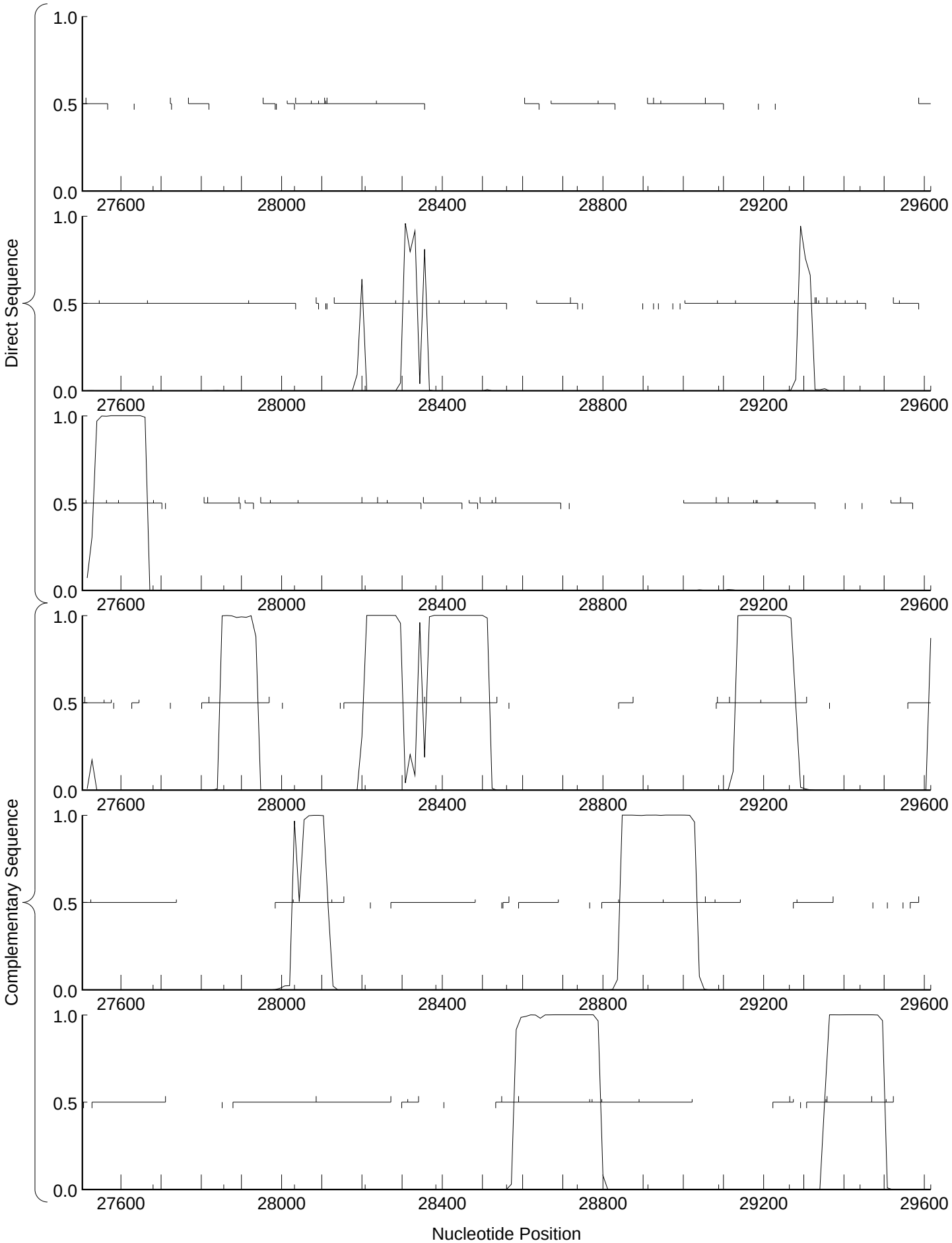

Direct Sequence
Complementary Sequence
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0.0
17200
17600
18000
18400
18800
Nucleotide Position

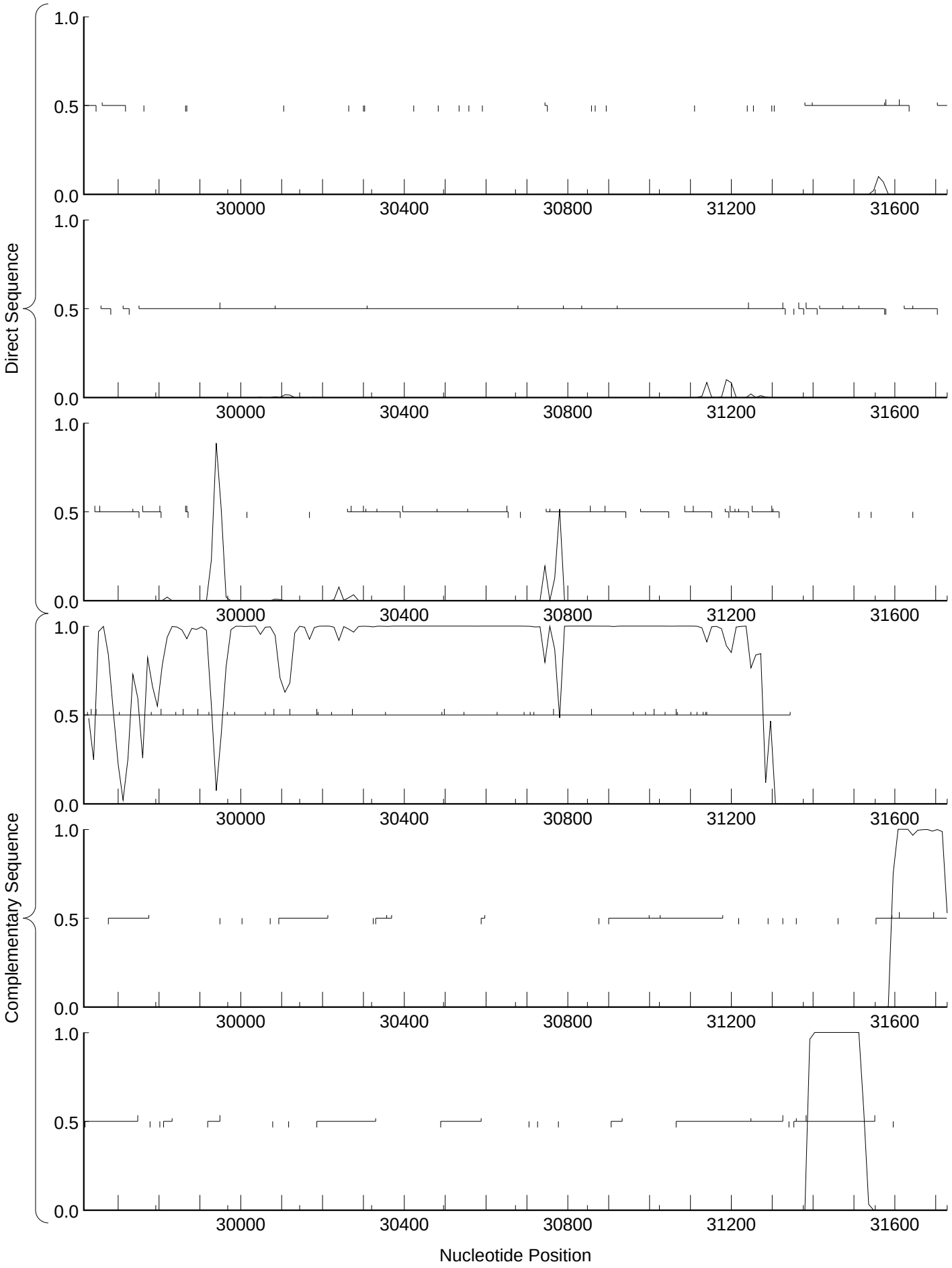

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Complementary Sequence
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20000
20400
20800
Nucleotide Position

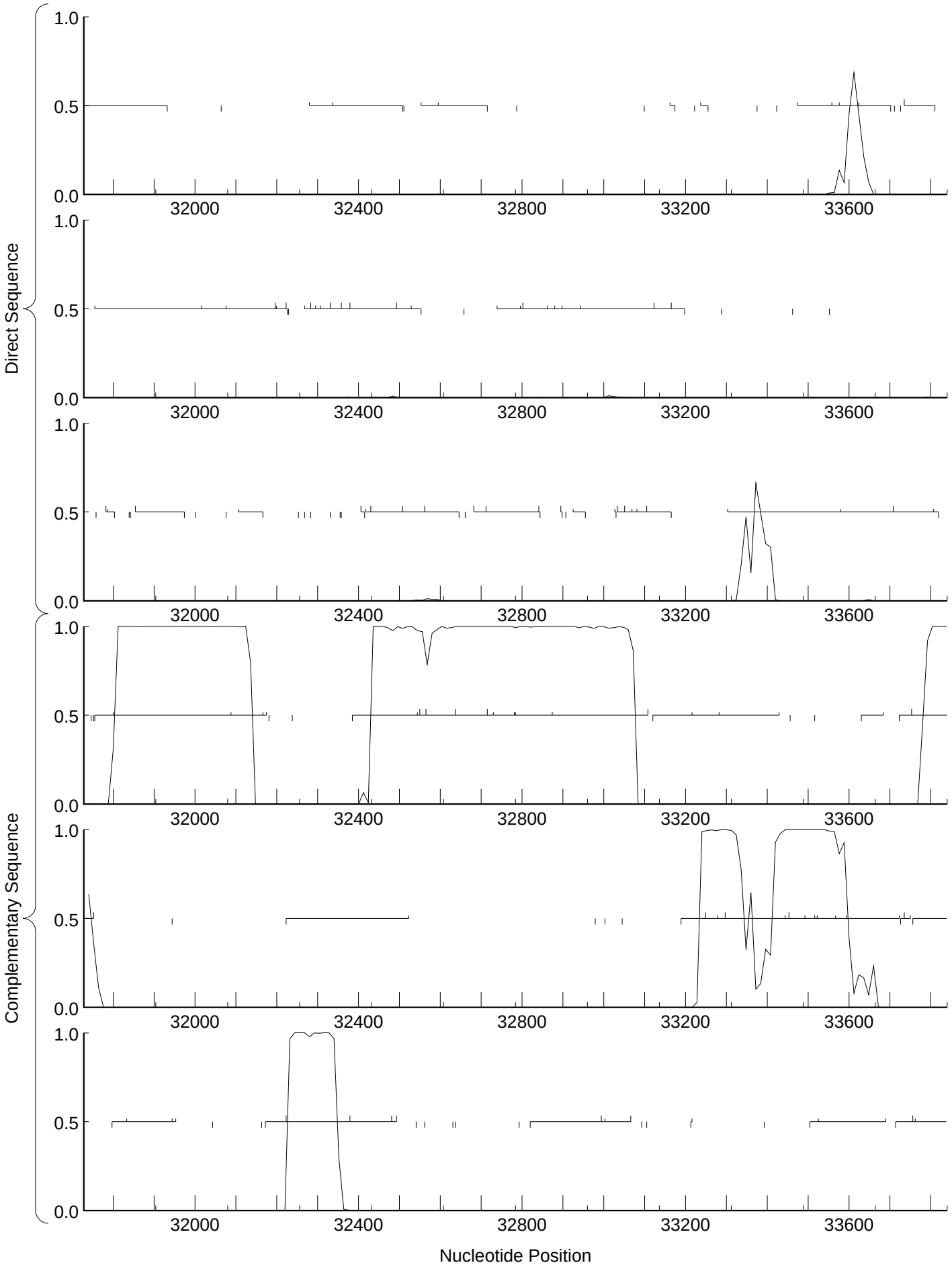

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22000
22400
22800
23200
Nucleotide Position

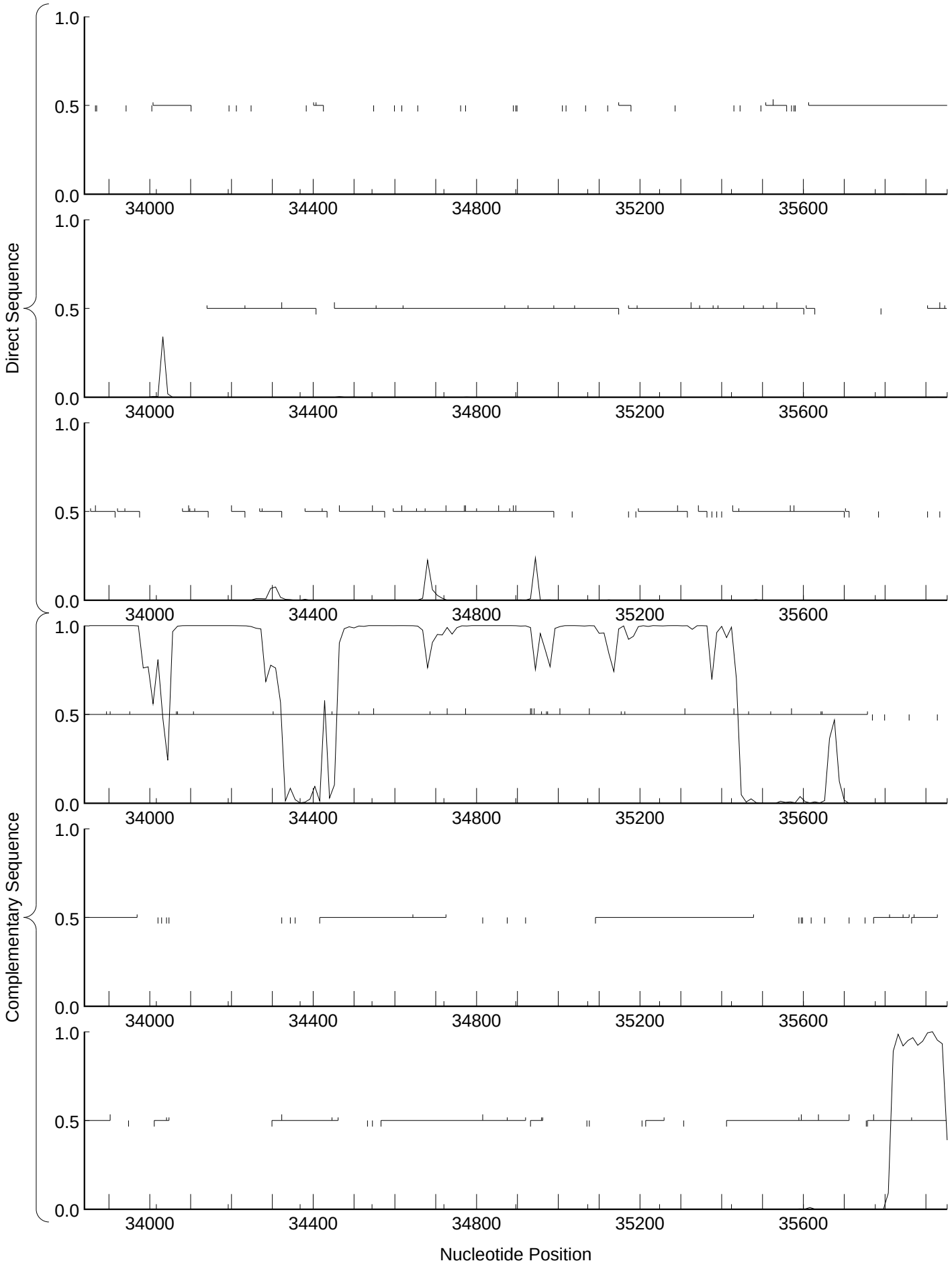

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Complementary Sequence
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24400
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Nucleotide Position

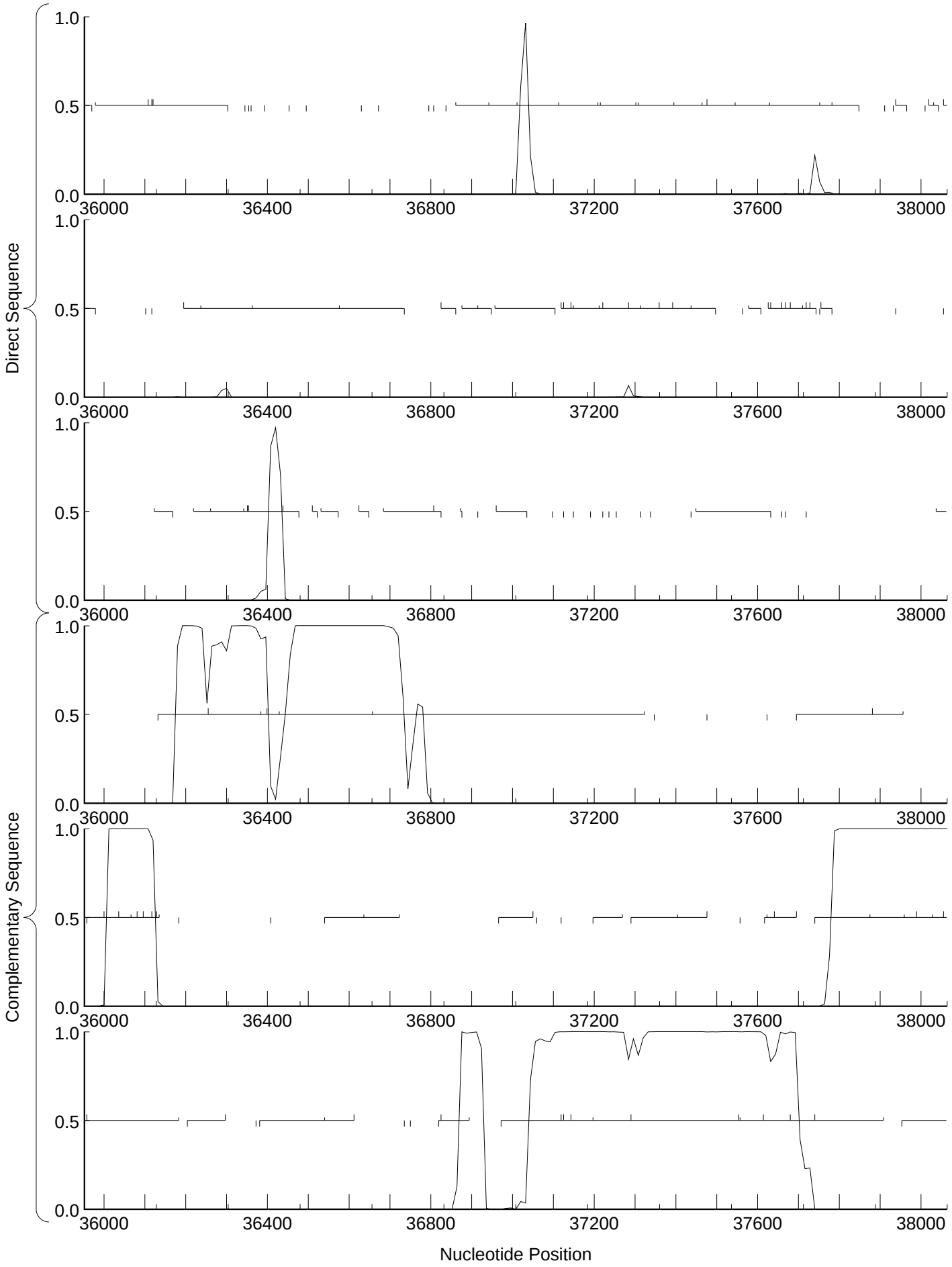

Direct Sequence
Complementary Sequence
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26800
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Nucleotide Position

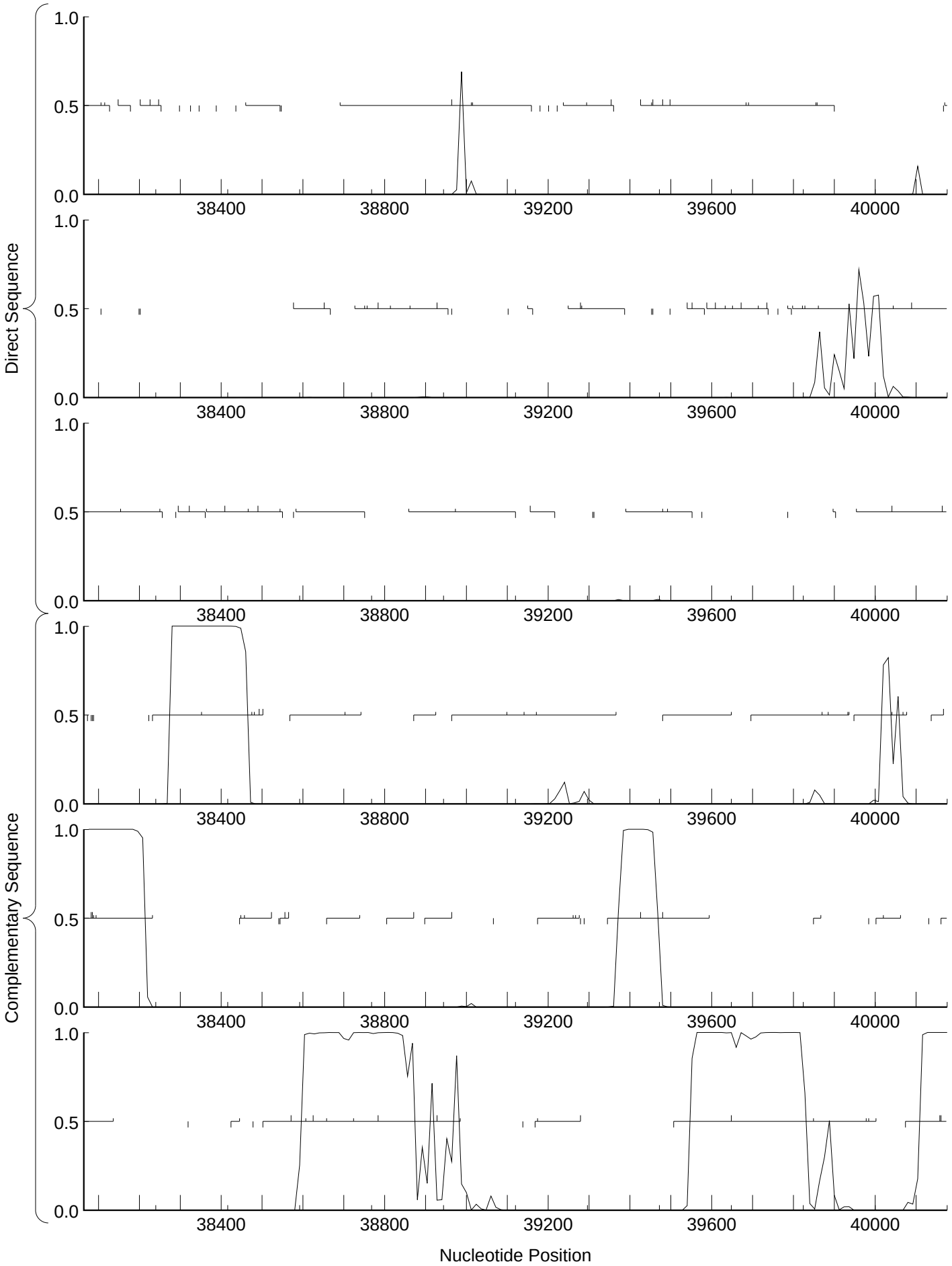

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Complementary Sequence
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Nucleotide Position

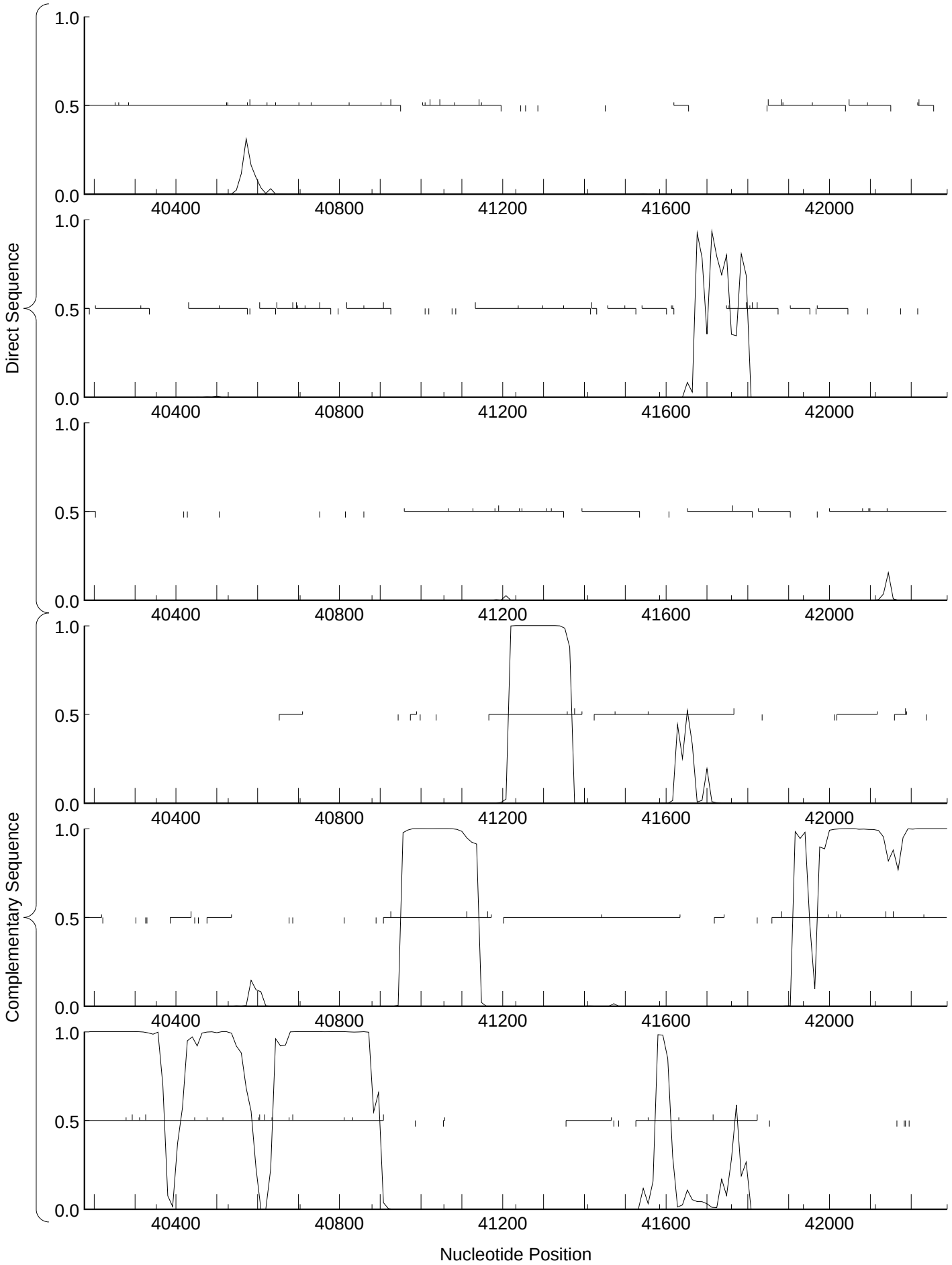

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31200
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Nucleotide Position

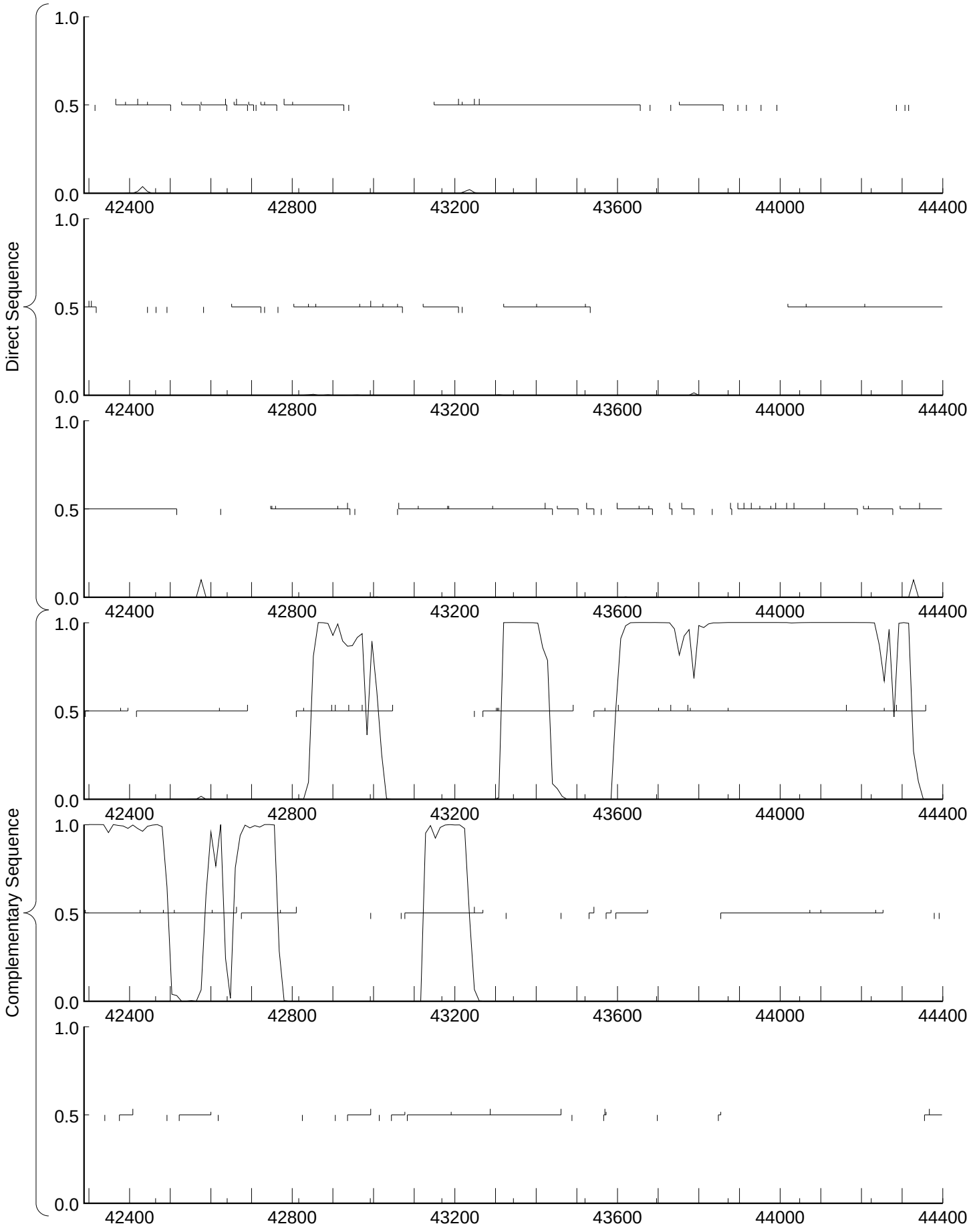

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33600
Nucleotide Position

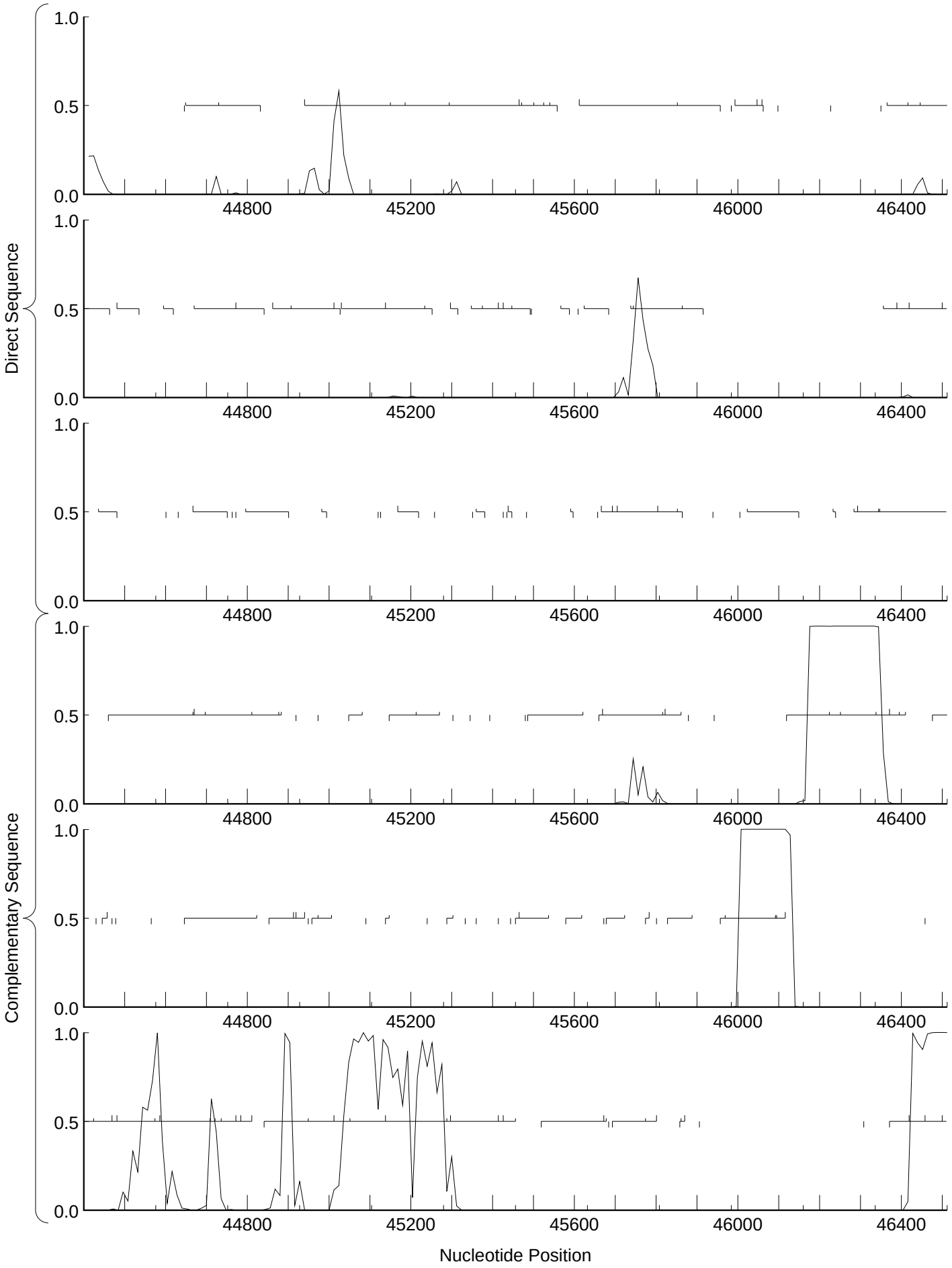

Direct Sequence
Complementary Sequence
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Nucleotide Position

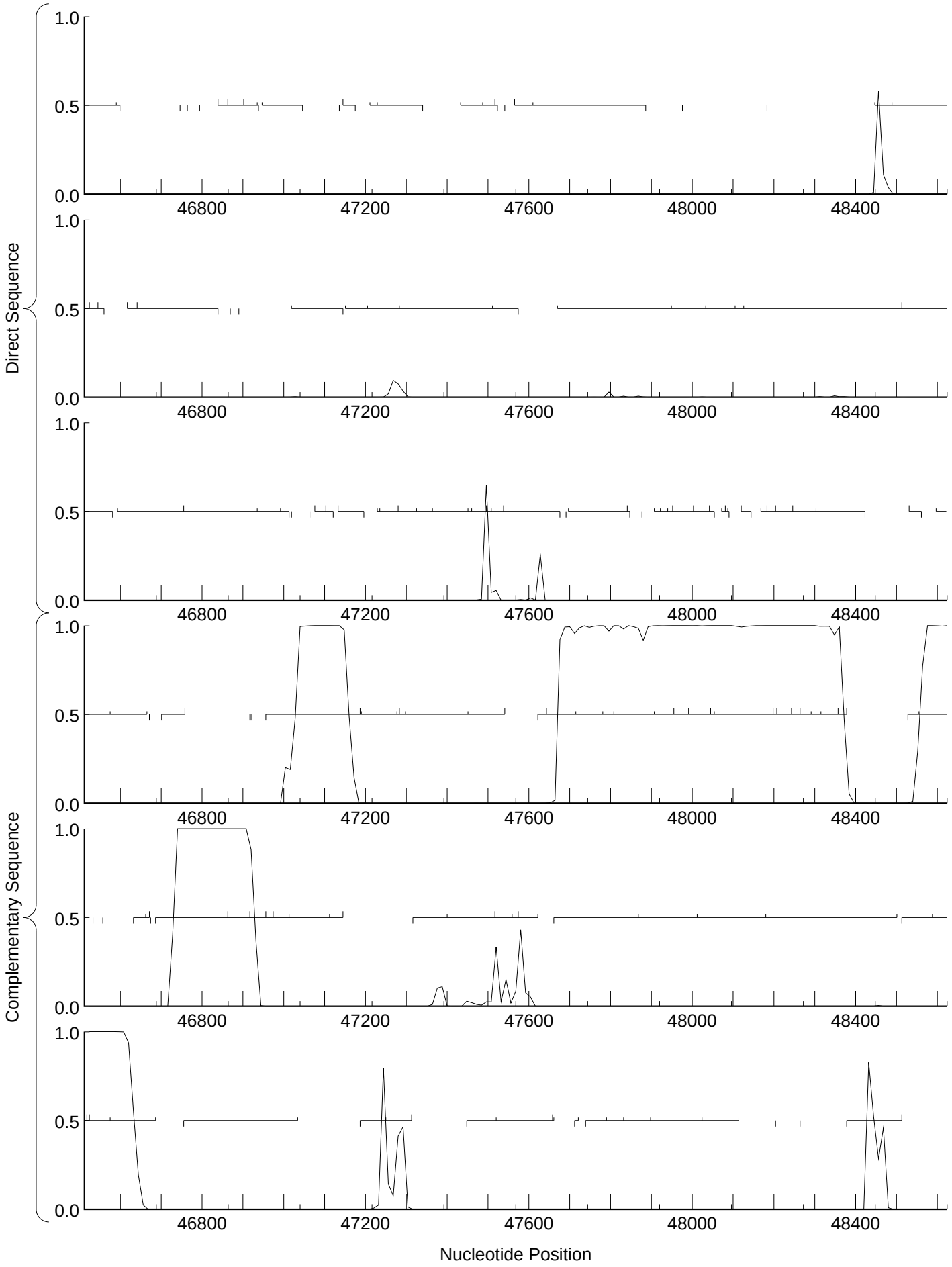

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Complementary Sequence
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37200
37600
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Nucleotide Position

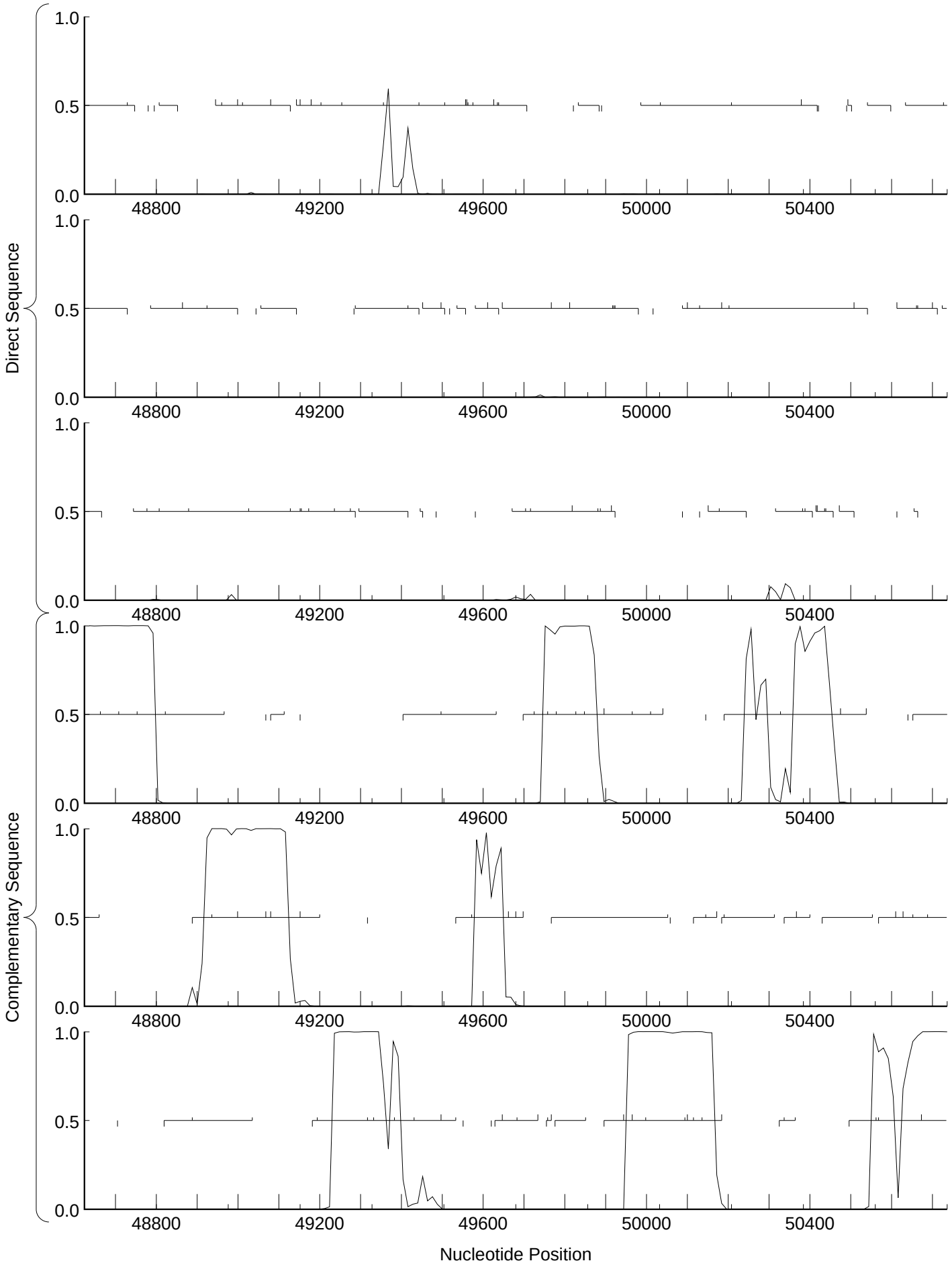

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Complementary Sequence
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0.0
38400
38800
39200
39600
40000
Nucleotide Position

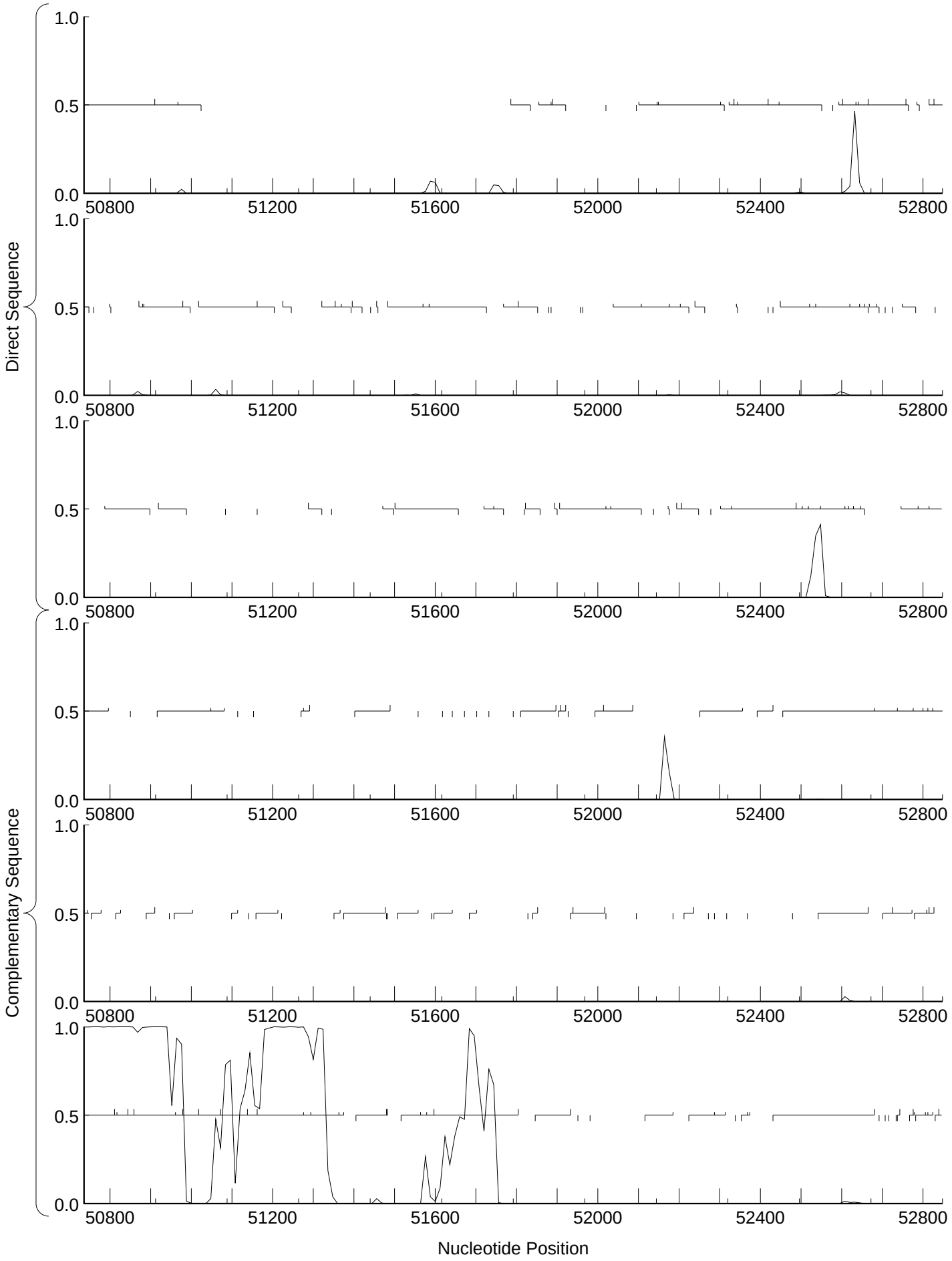

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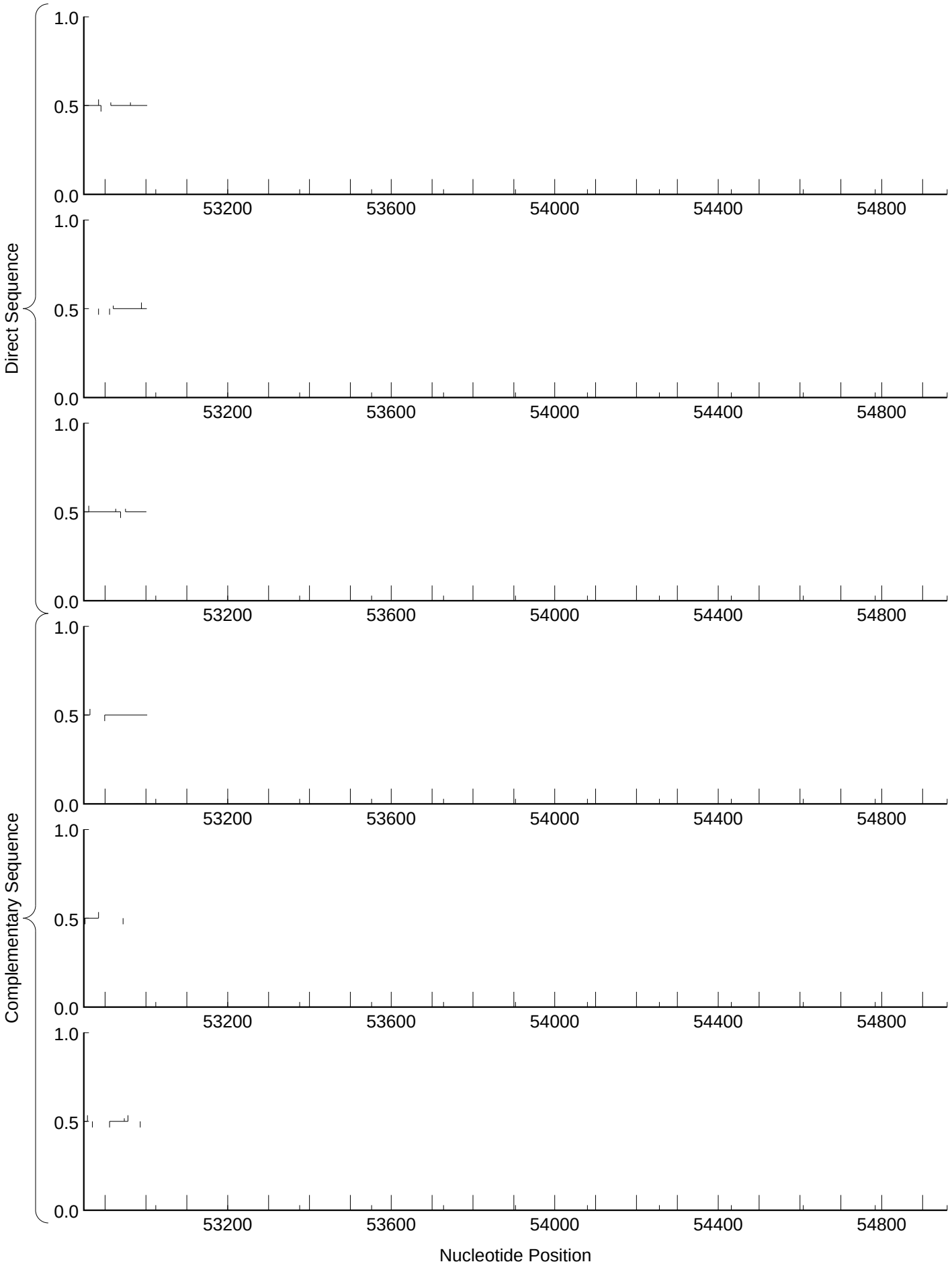

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0.0
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43200
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44400
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
44800
45200
45600
46000
46400
Nucleotide Position


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


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
48800
49200
49600
50000
50400
Nucleotide Position


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


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
53200
53600
54000
54400
54800
Nucleotide Position