GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage HarryOW complete sequence, 52935 bp including 10-base 3’ overhang (CGGA
Analysis Date    :  3/28/24 at 6:32:50
Pages            :  27
Sequence Length  :  52935 bp
GC Content       :   63.77%

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

PS-Version       :  1.2

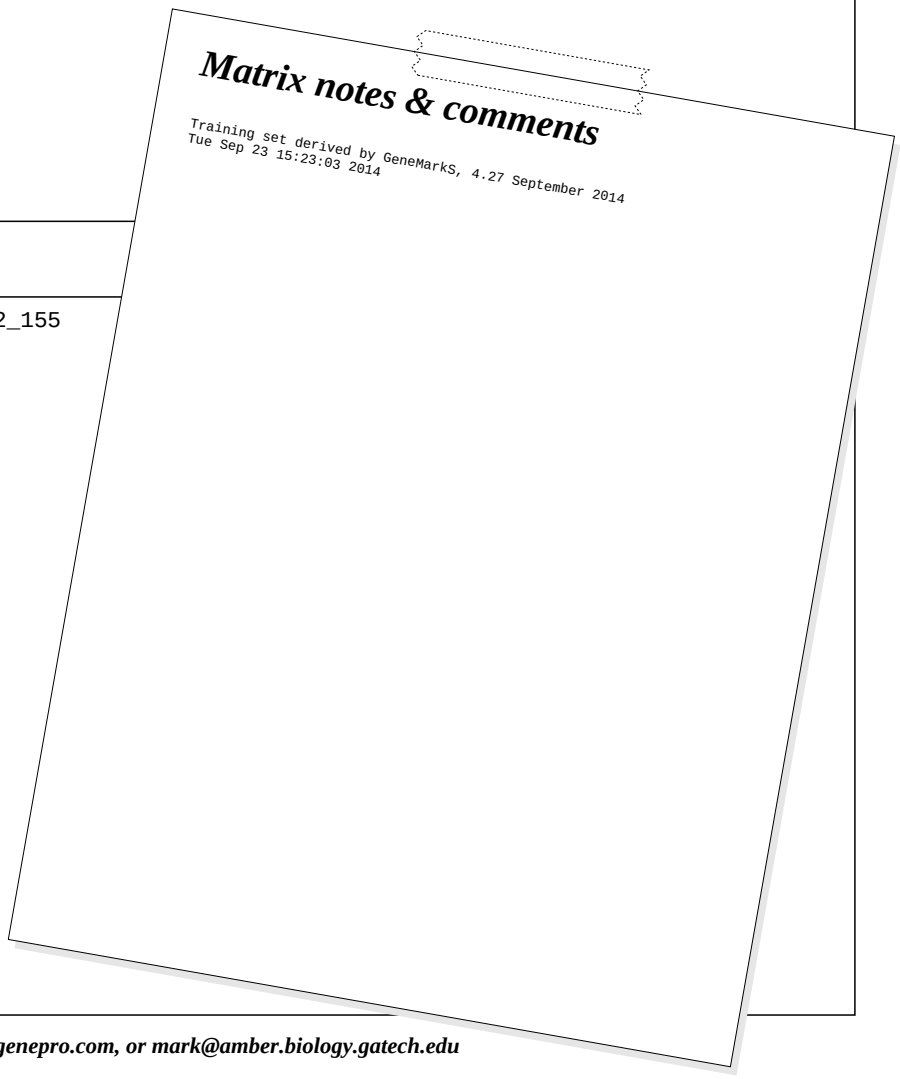
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Author     :  -

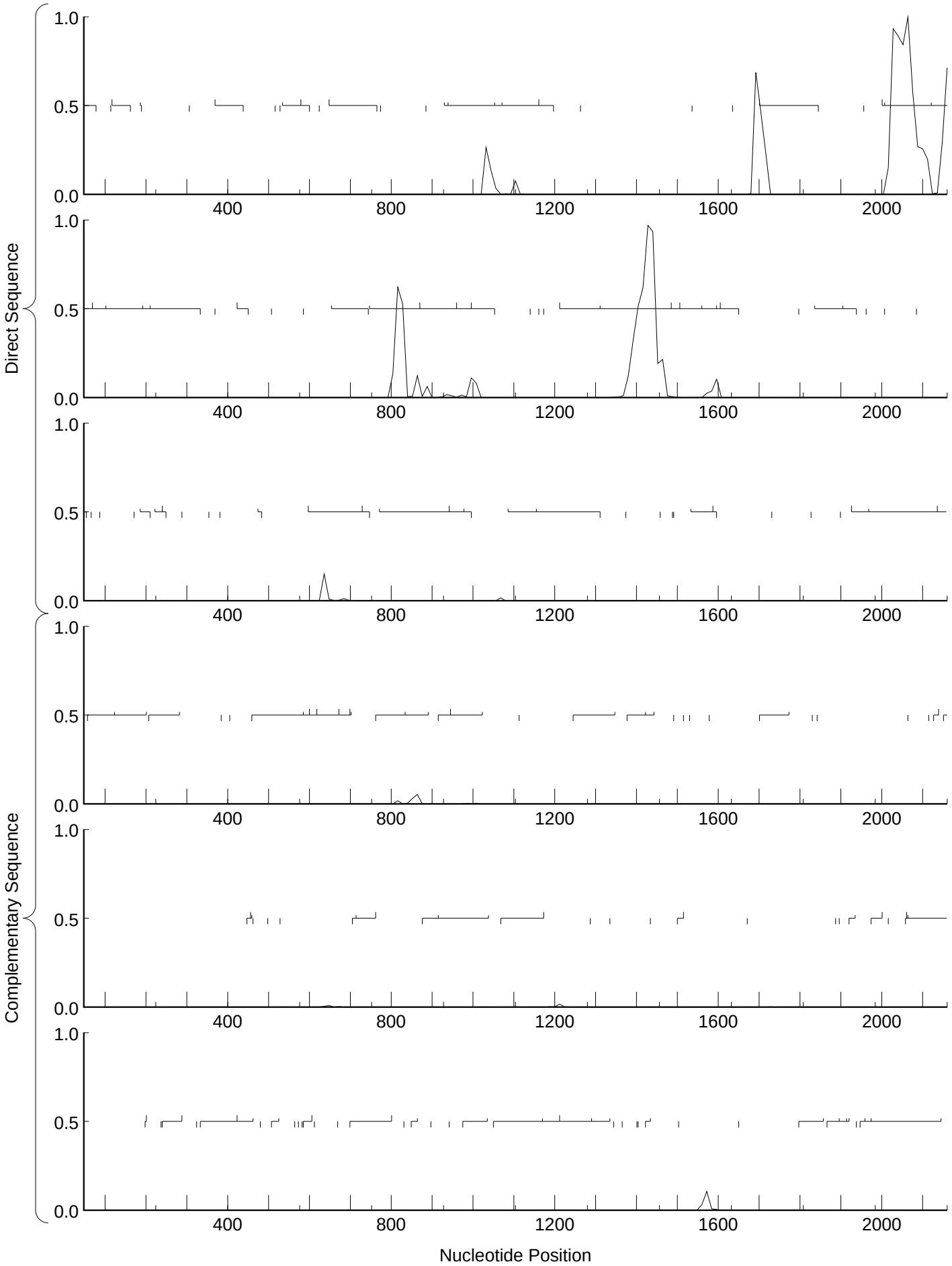
Order      :  4


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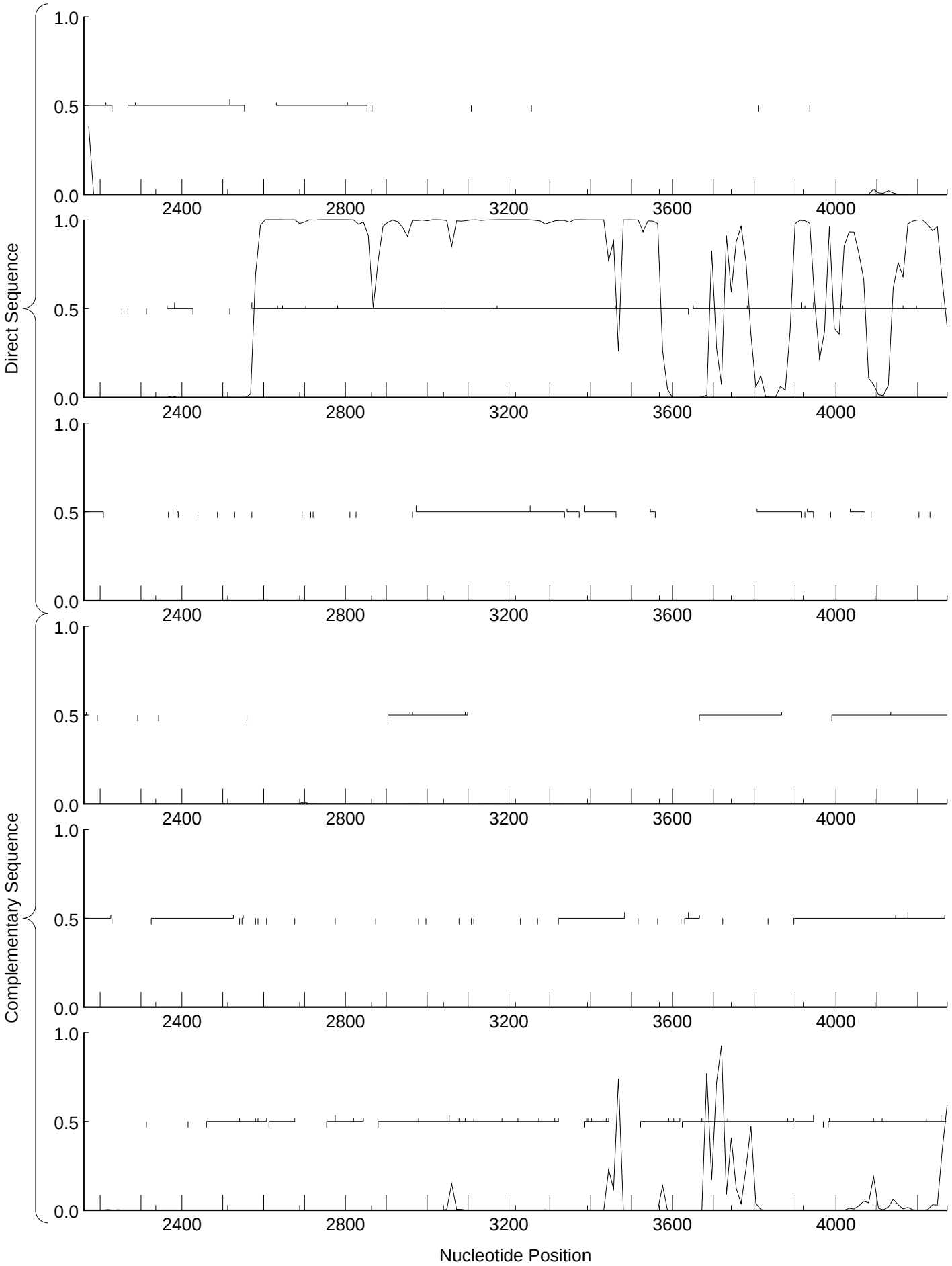


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

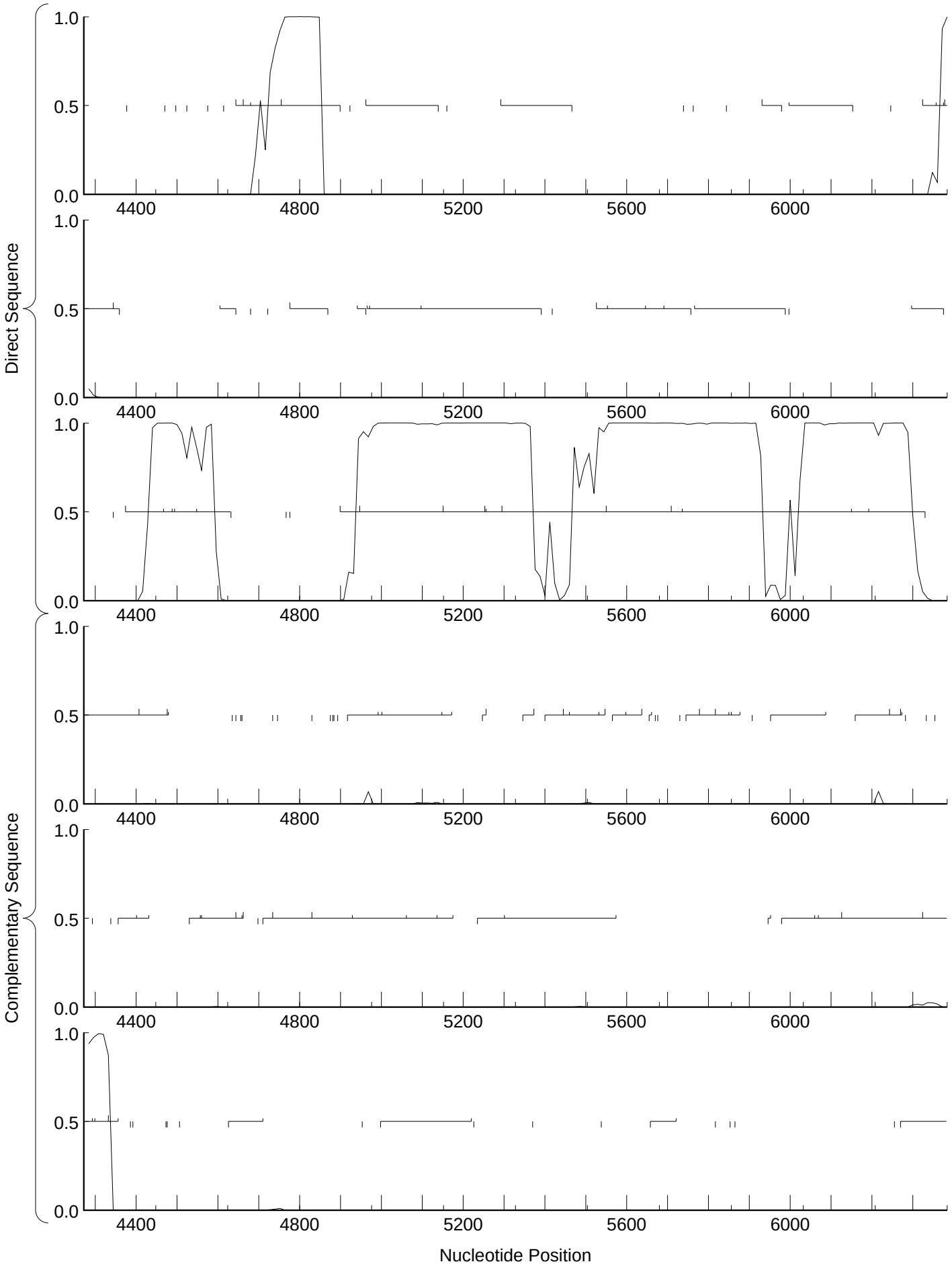
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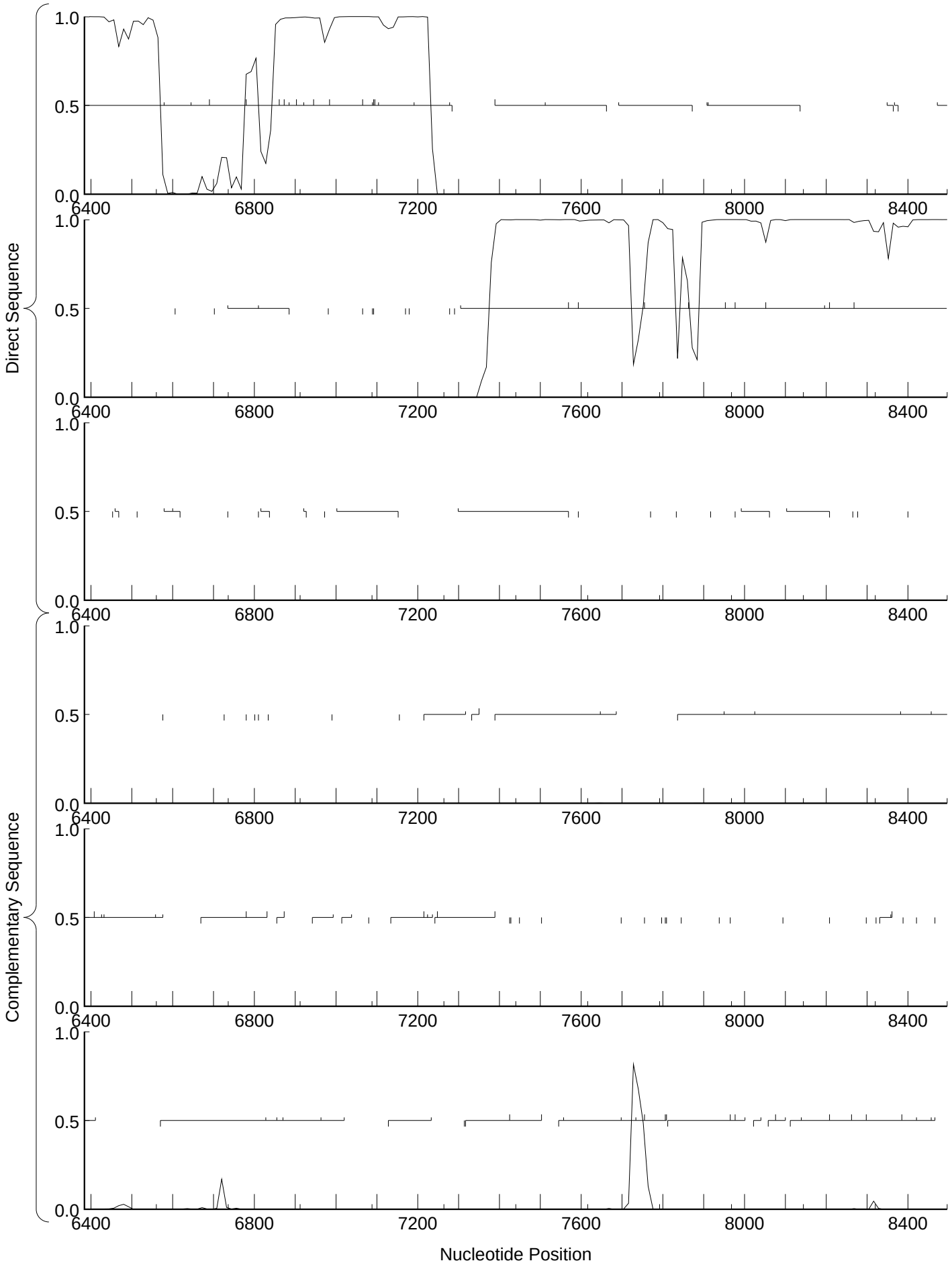
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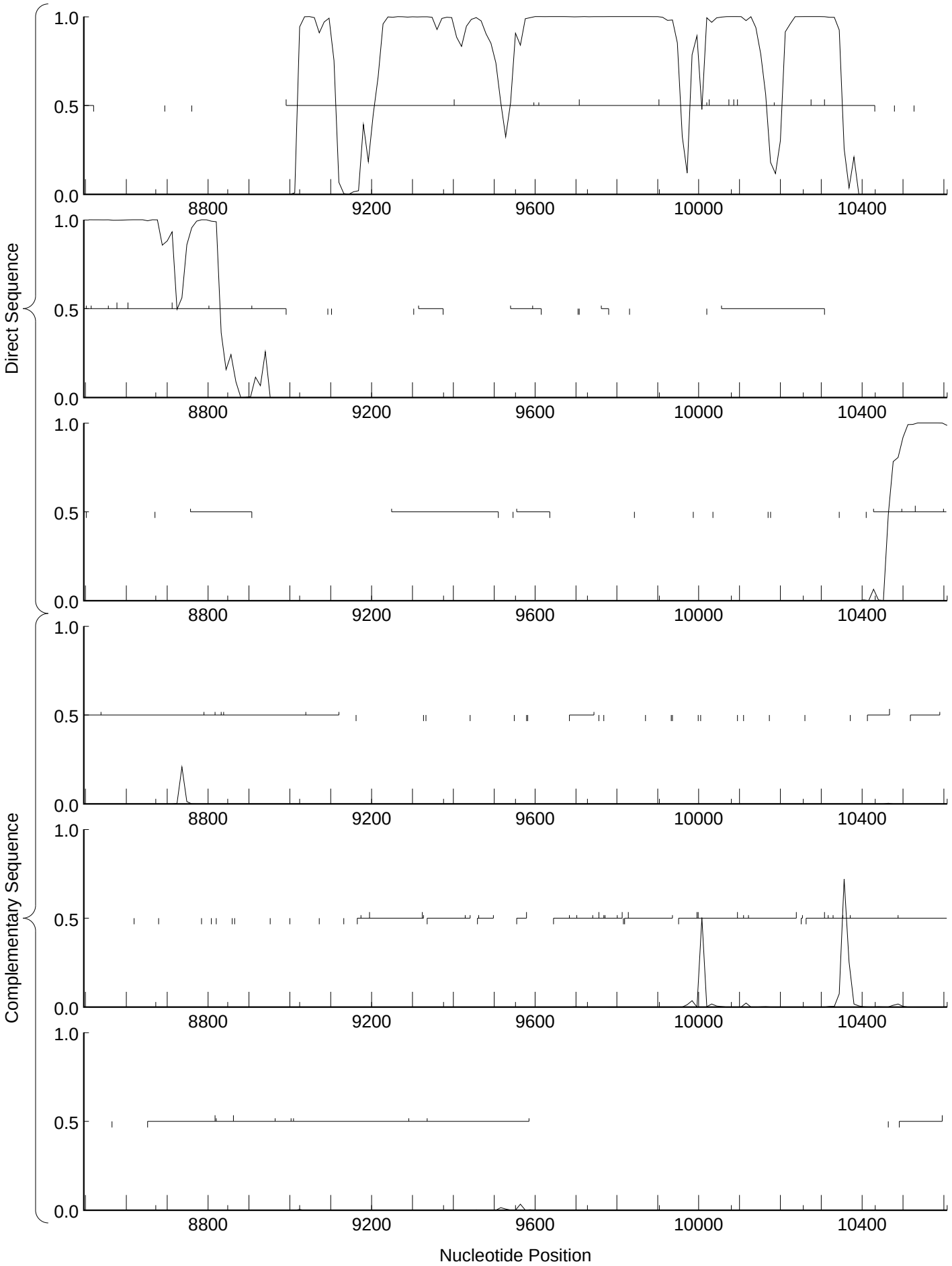
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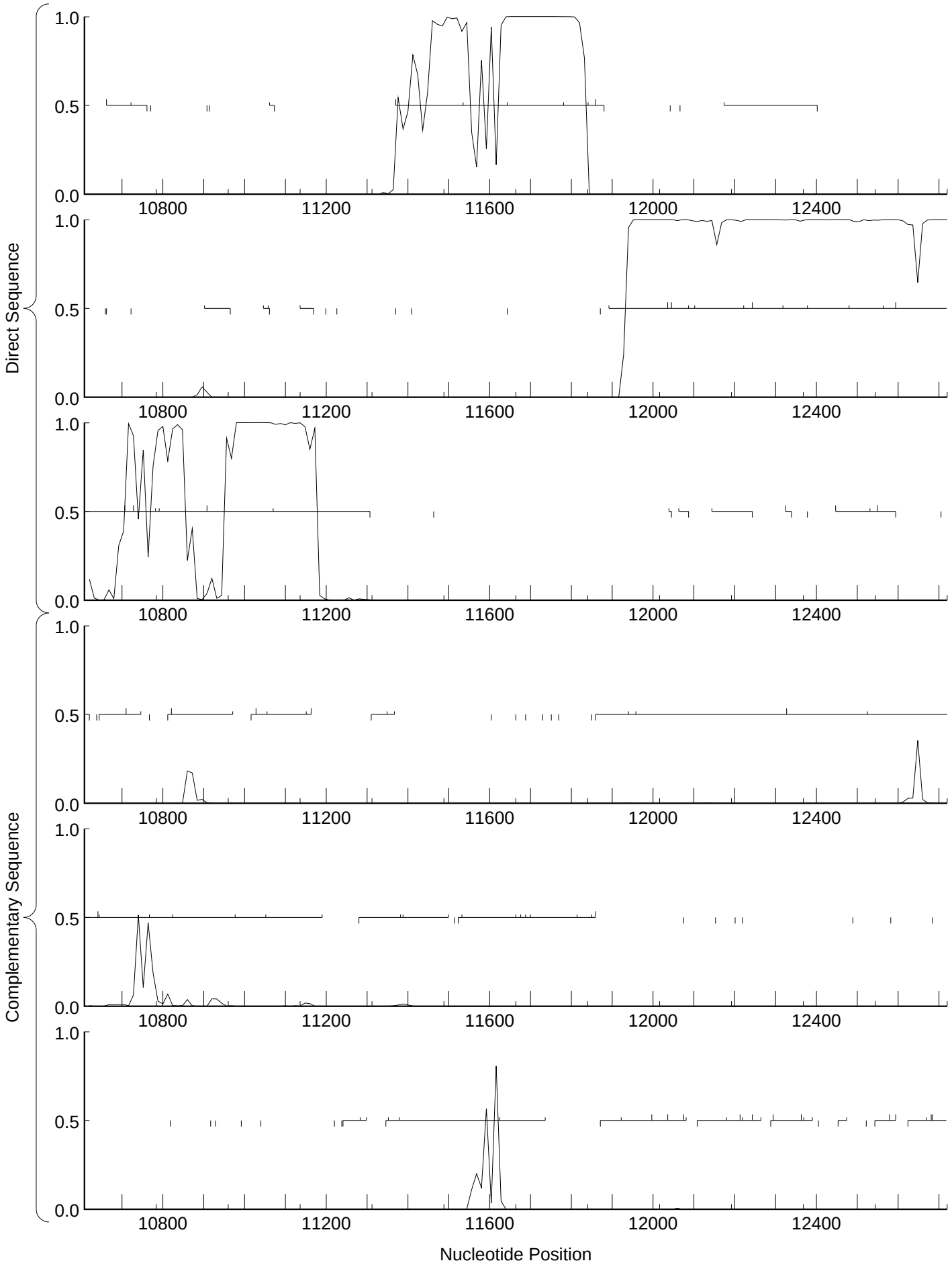
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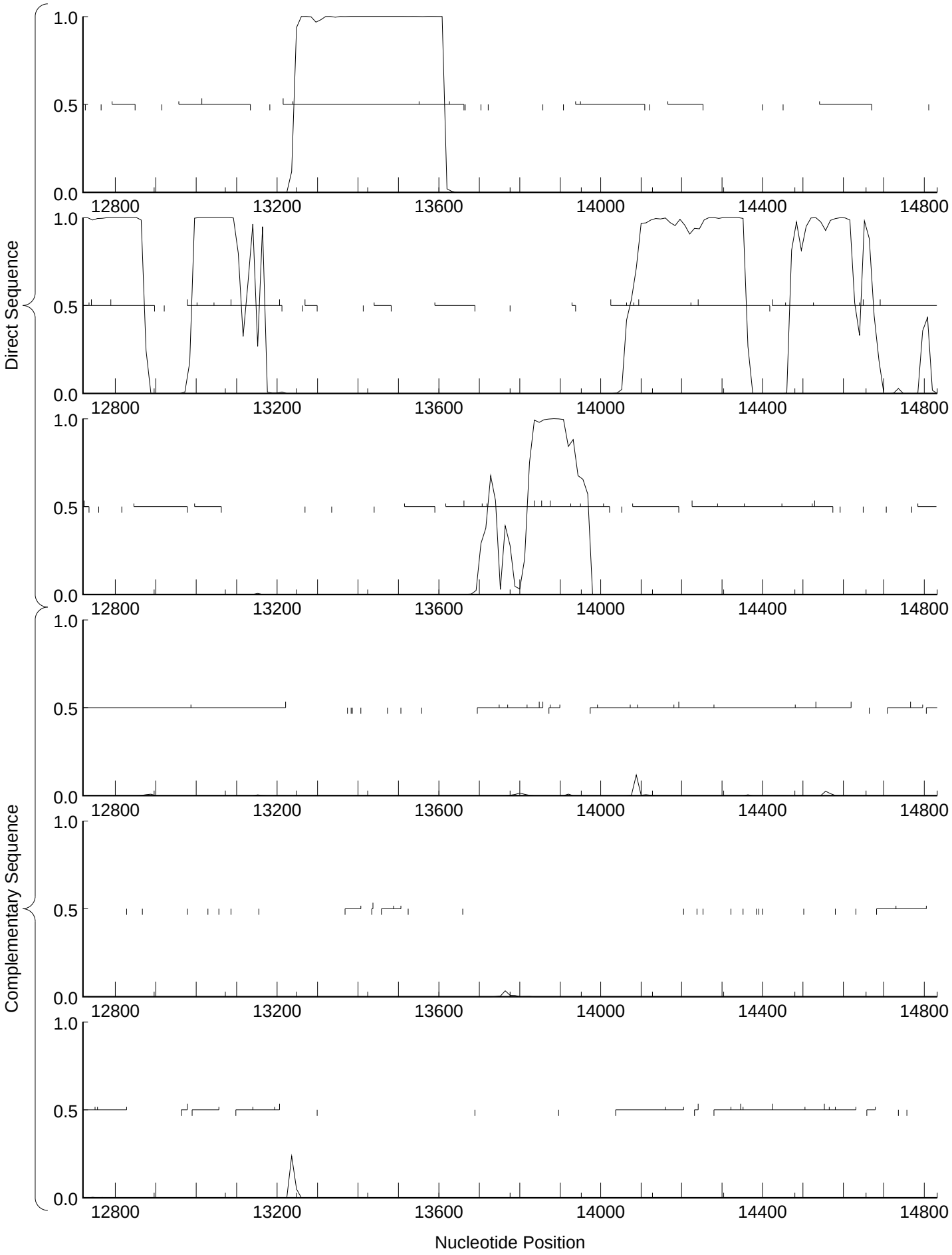
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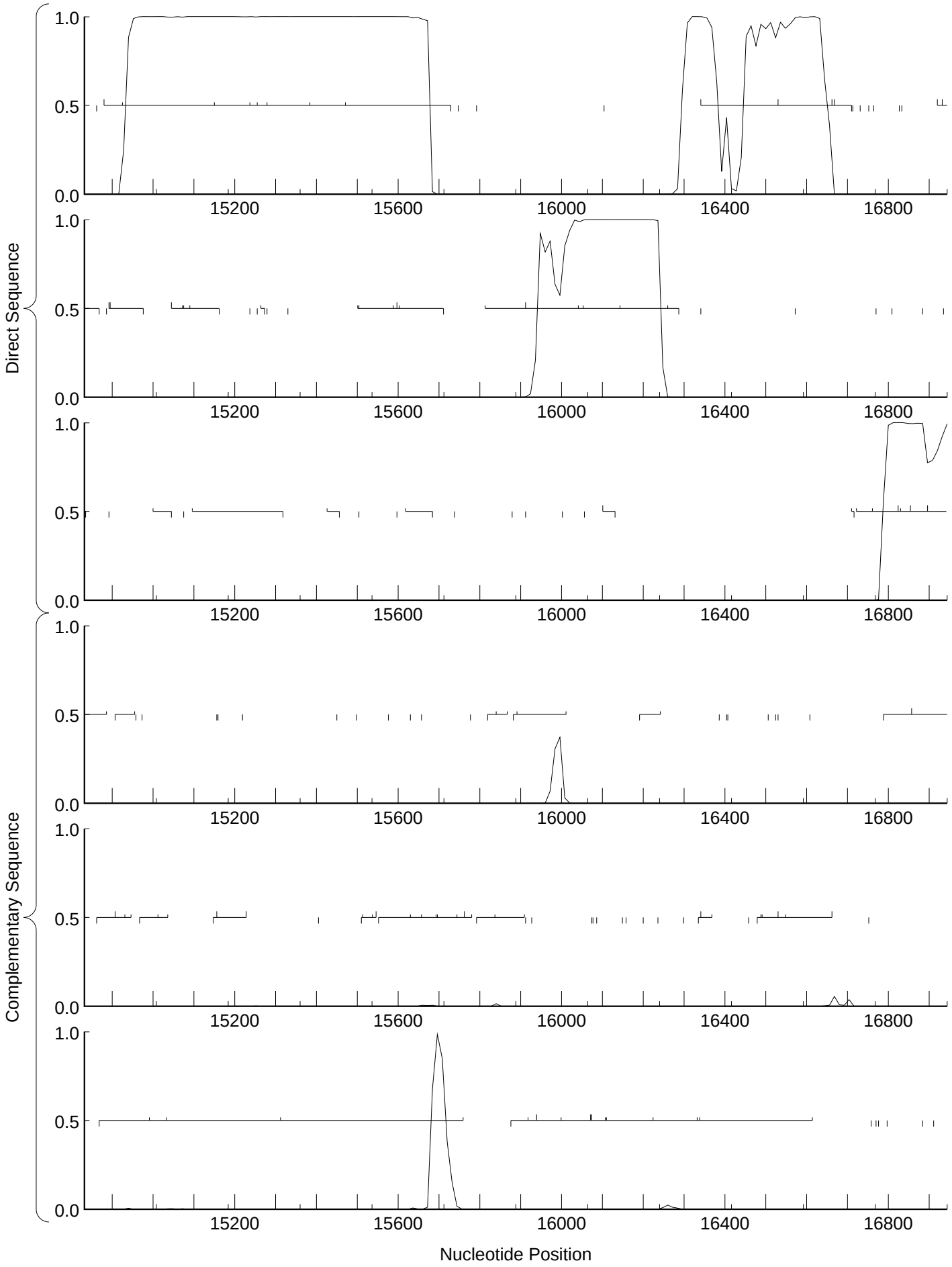
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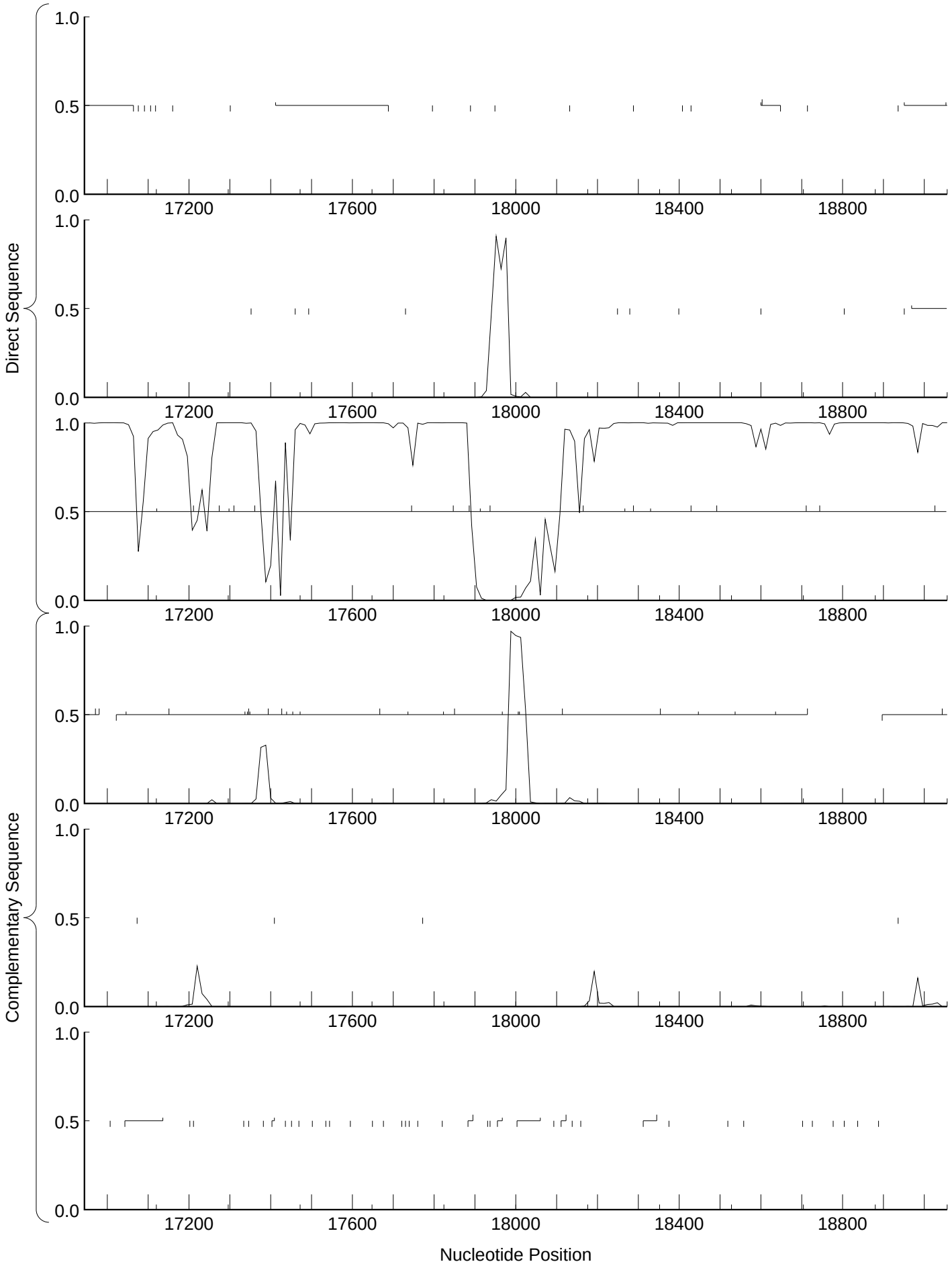


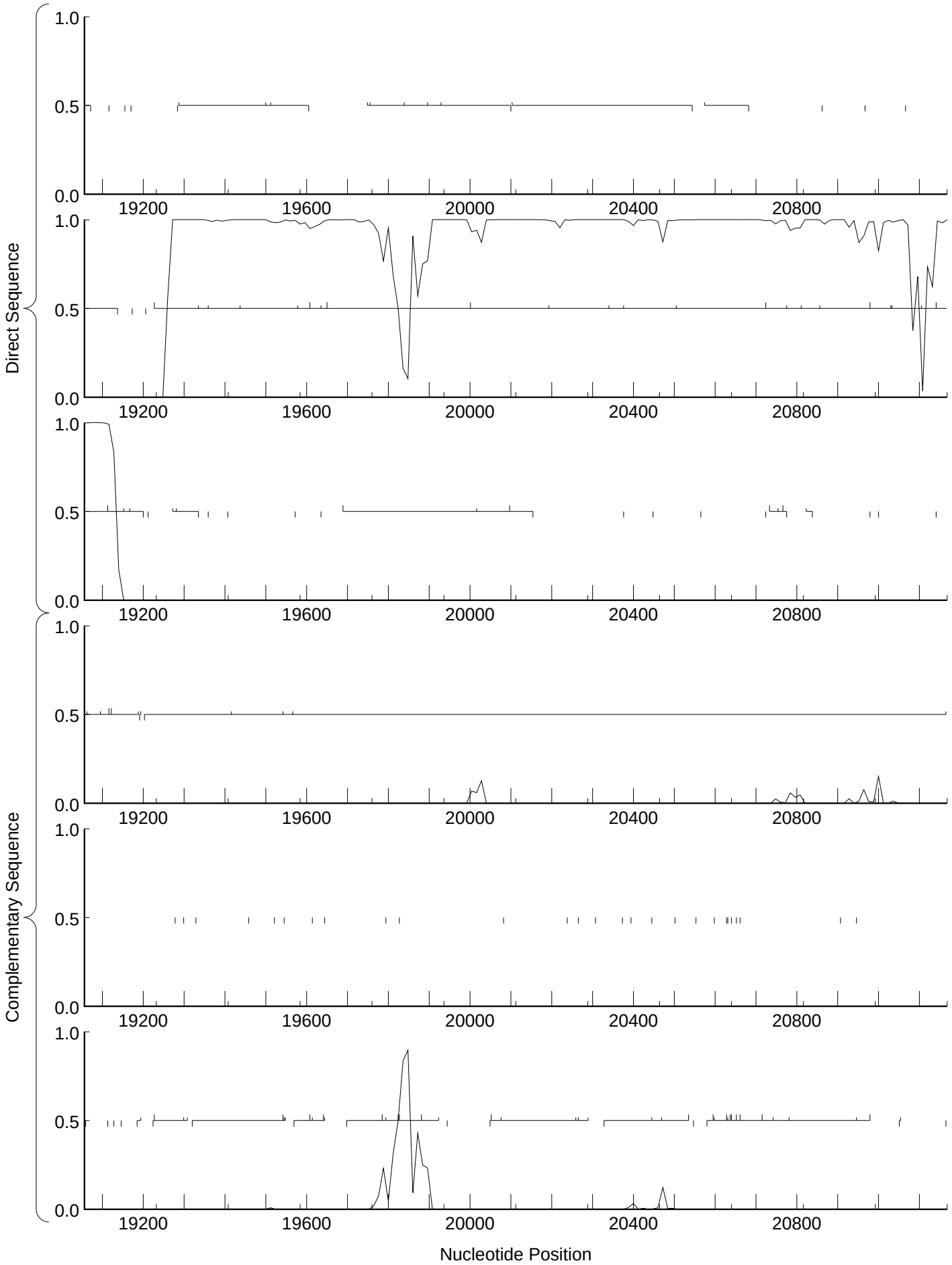
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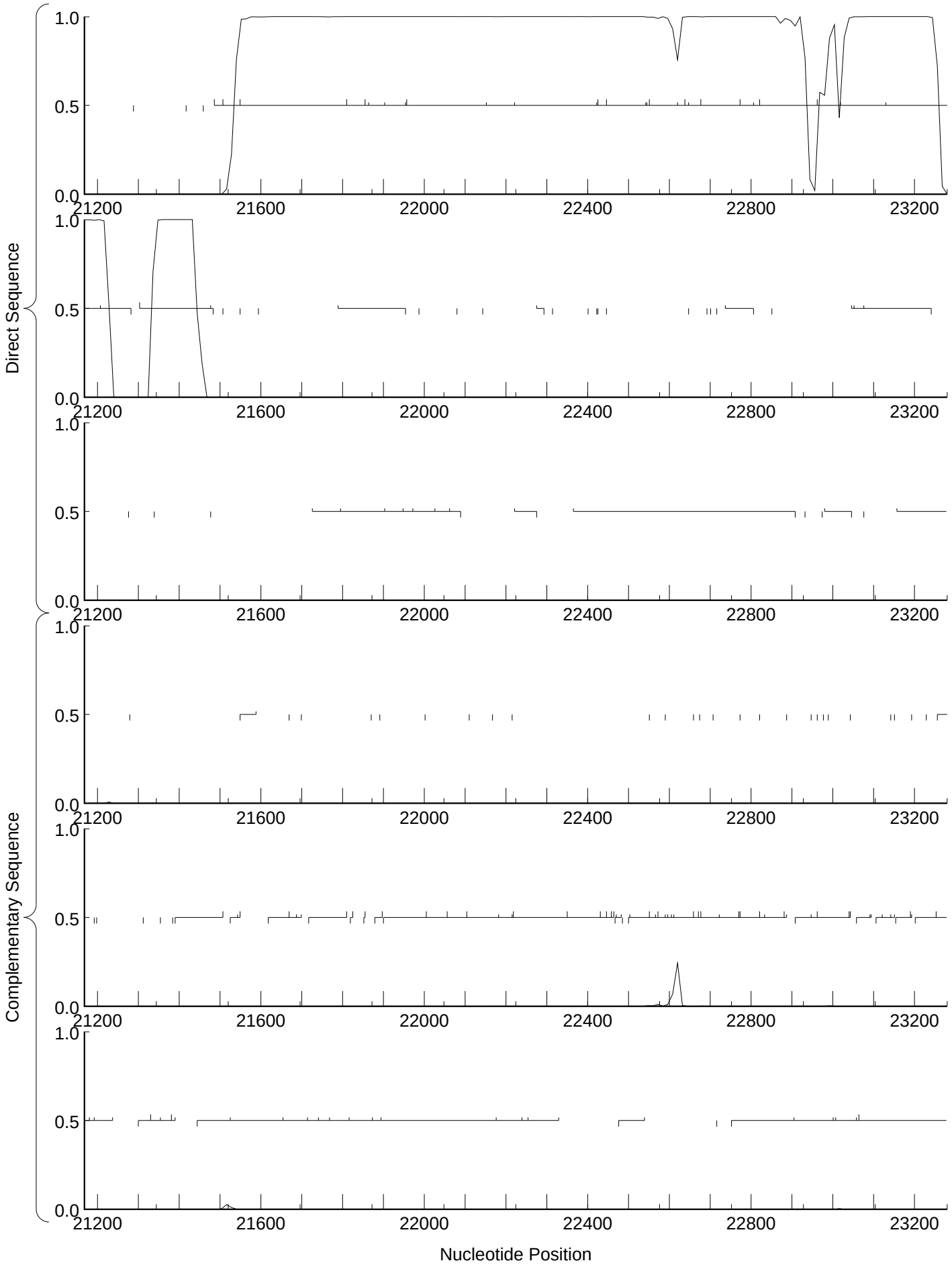


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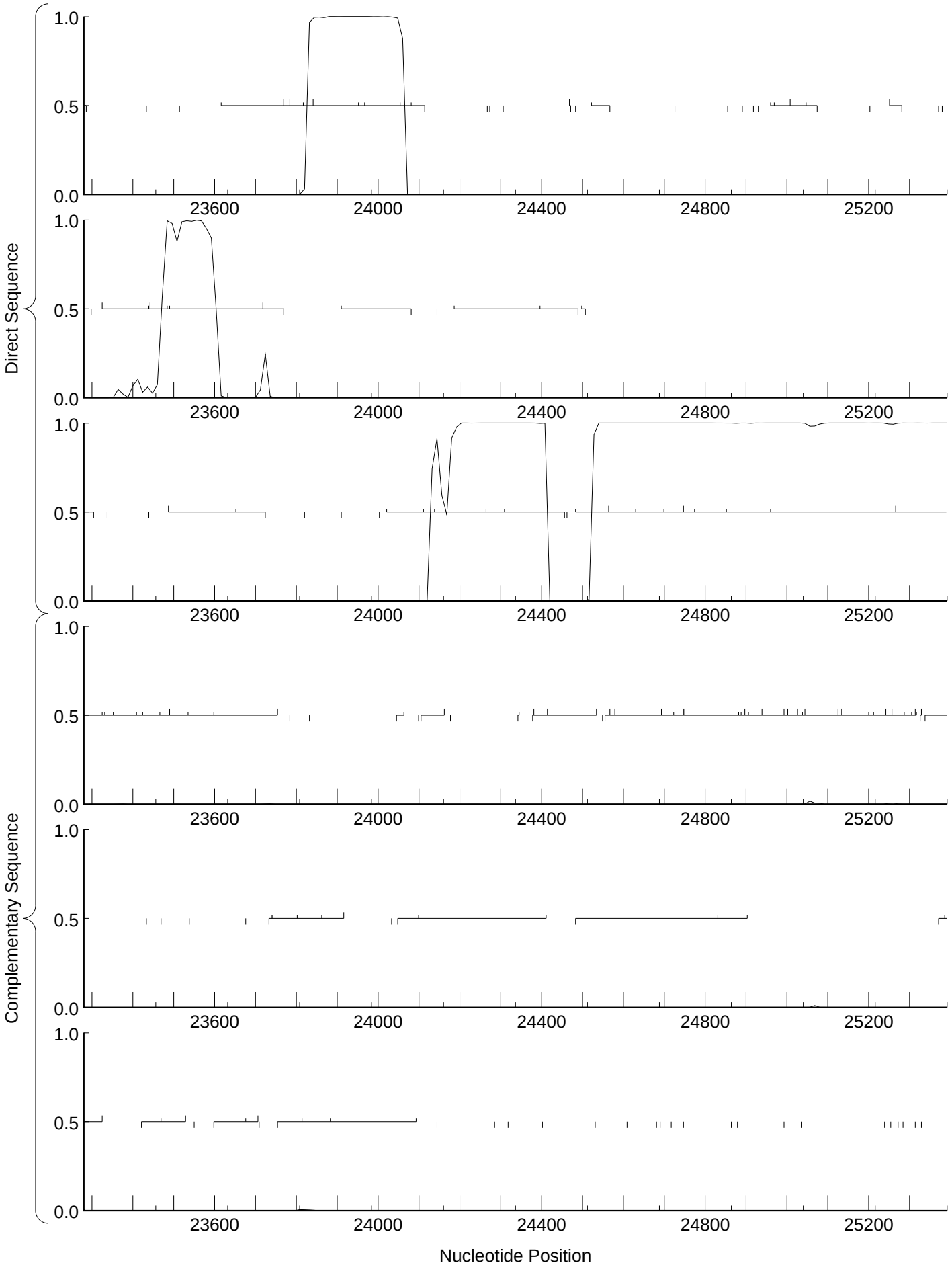


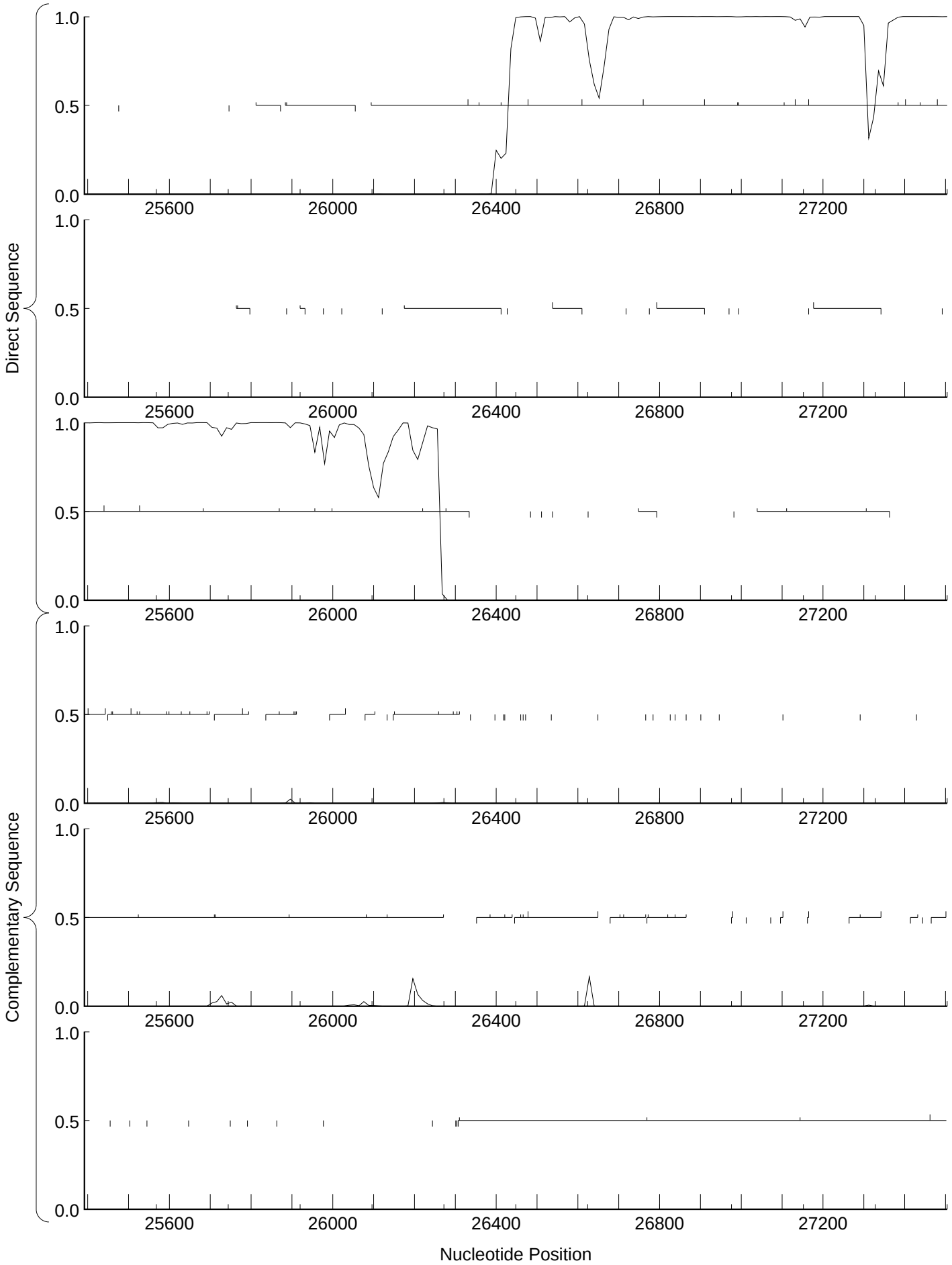

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Complementary Sequence
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0.5
0.0
17200
17600
18000
18400
18800
Nucleotide Position

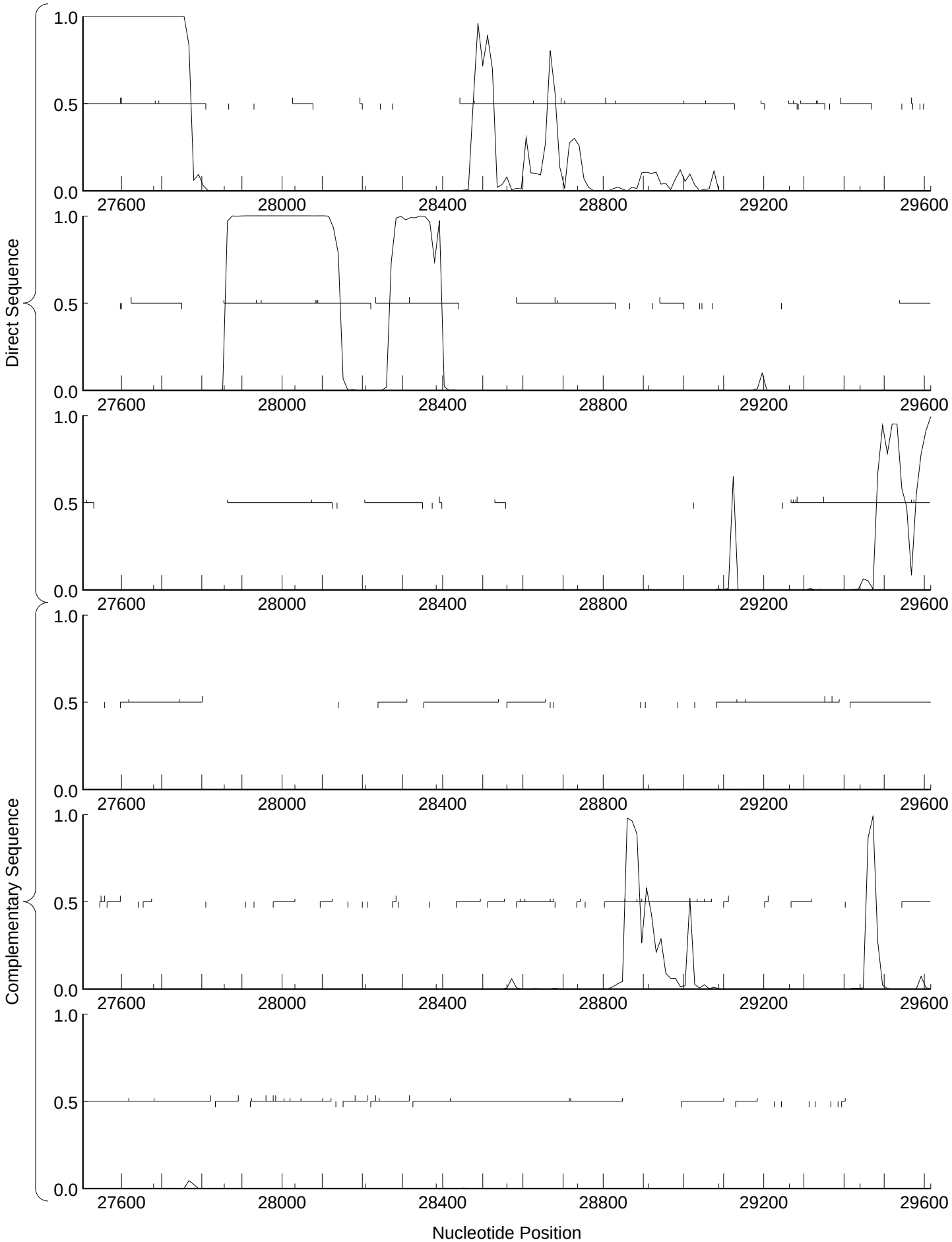

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Complementary Sequence
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20000
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Nucleotide Position

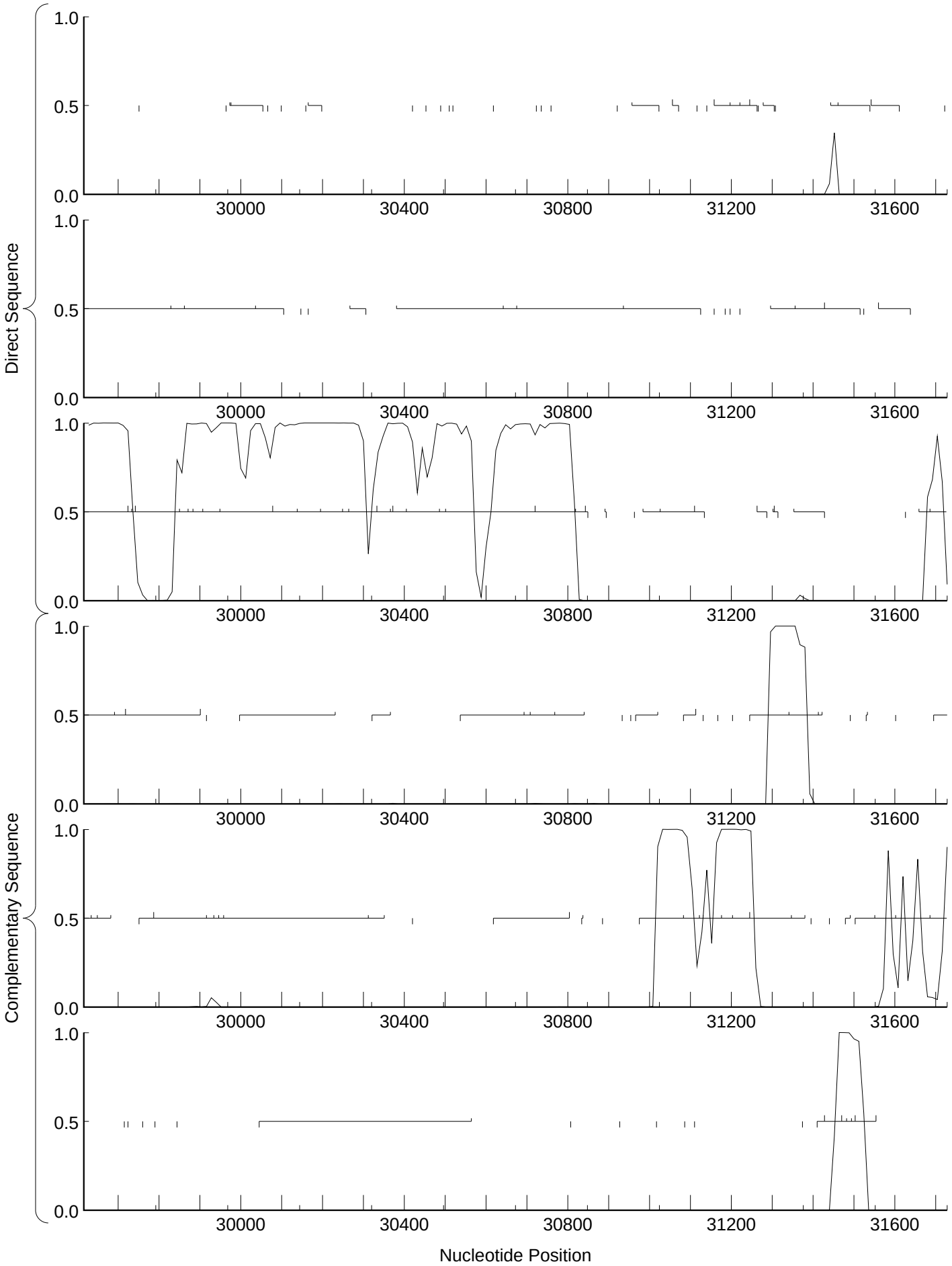

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Complementary Sequence
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21600
22000
22400
22800
23200
Nucleotide Position

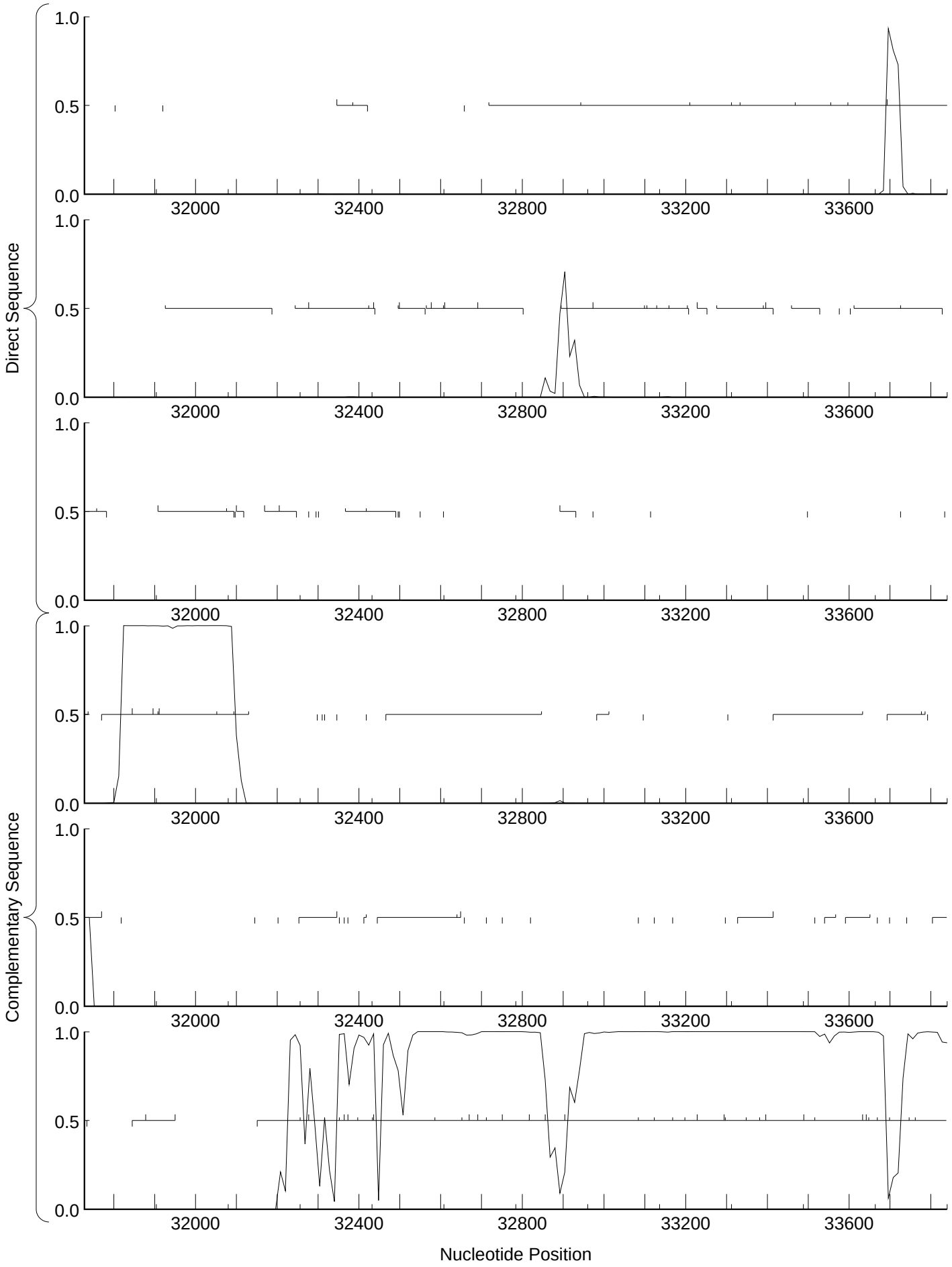
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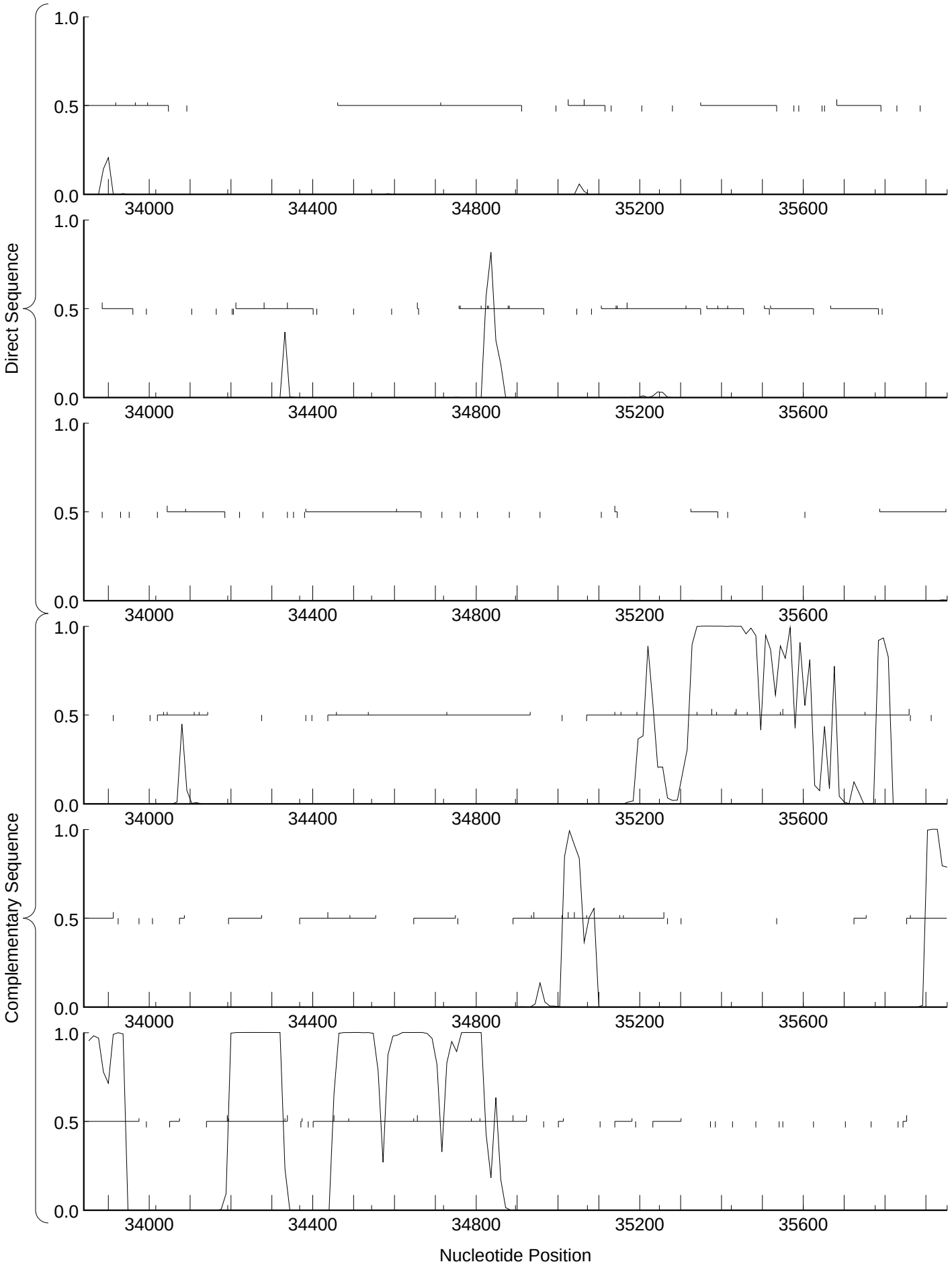


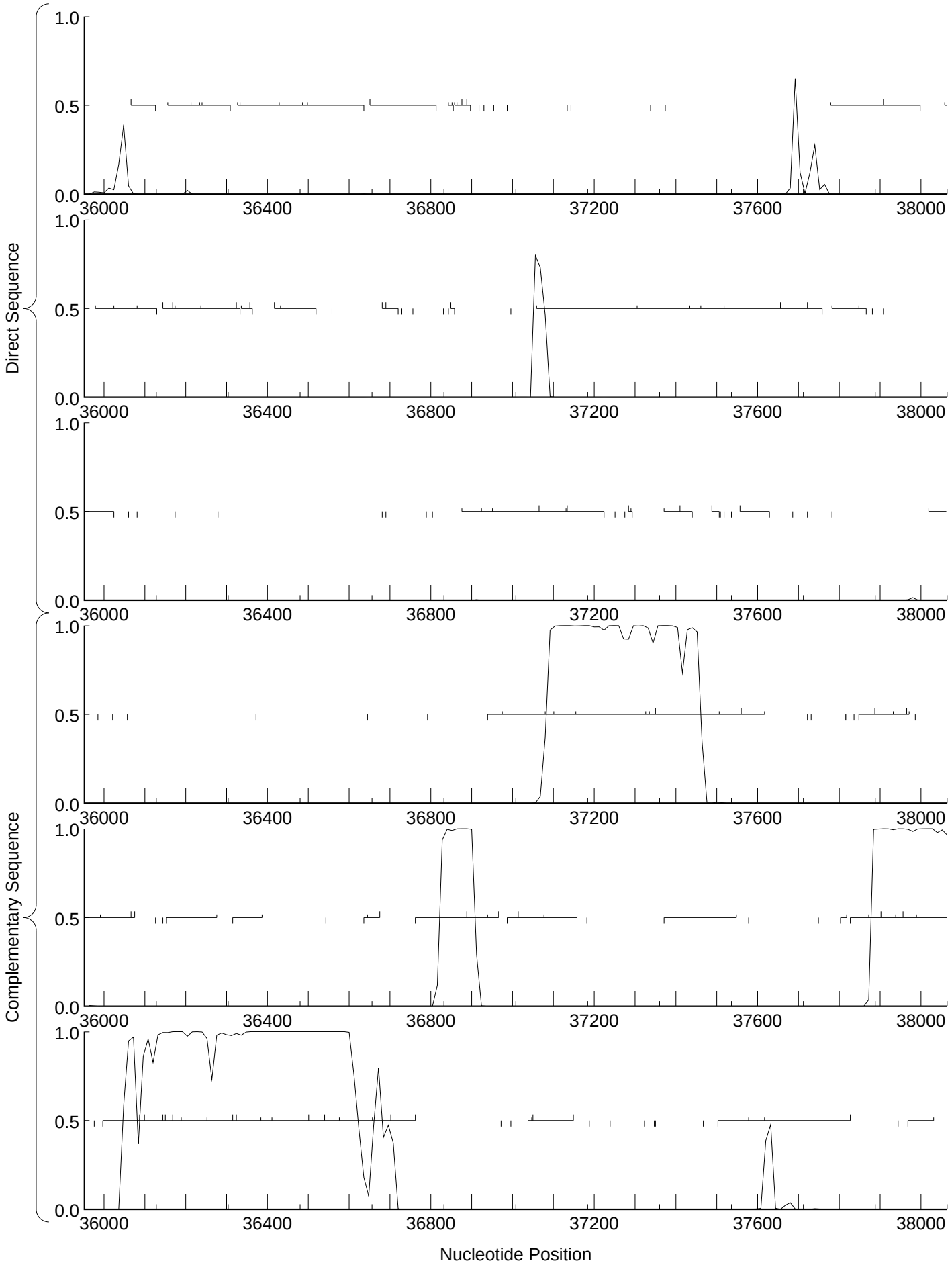

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Complementary Sequence
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0.0
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26000
26400
26800
27200
Nucleotide Position

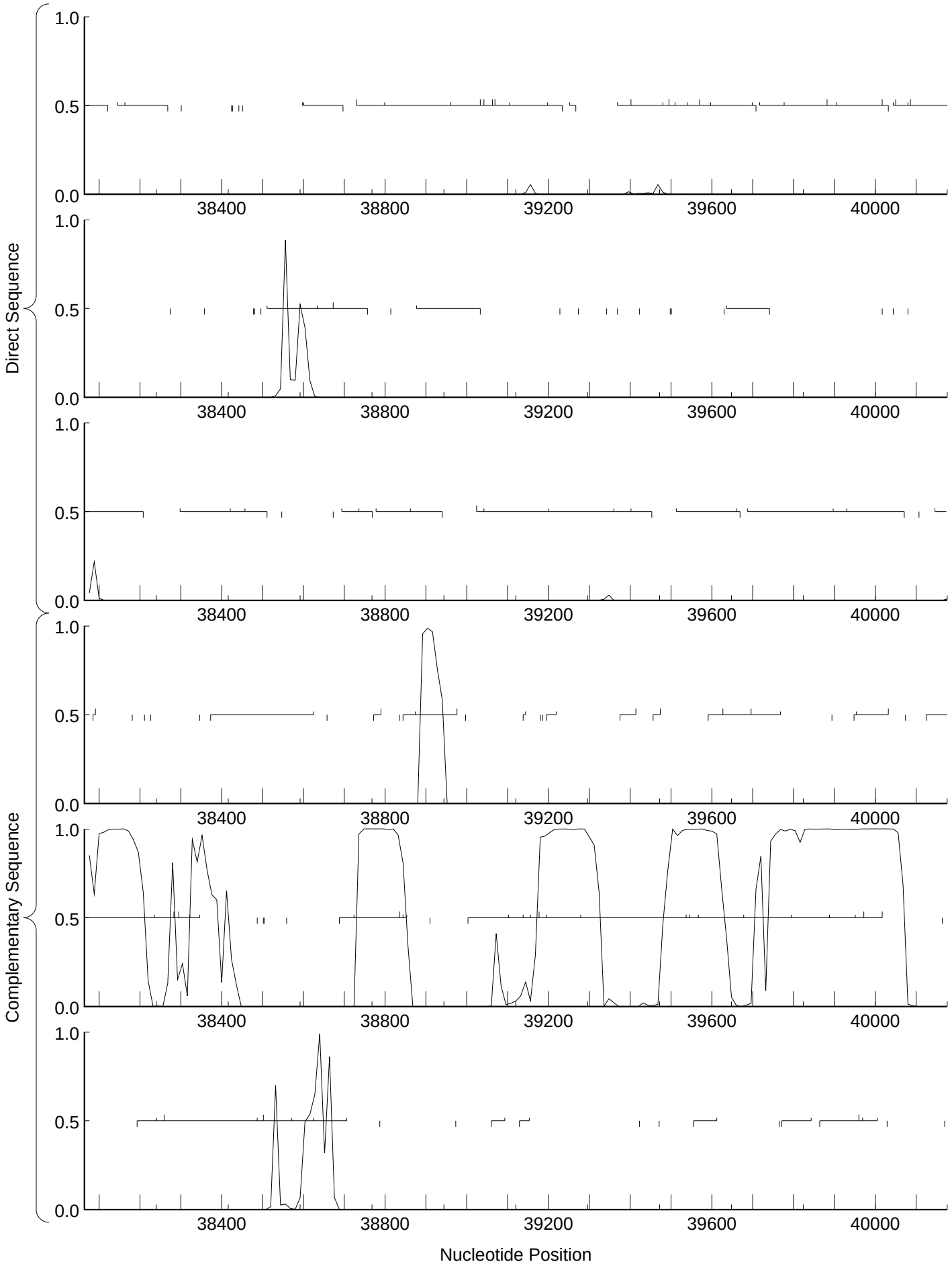

Direct Sequence
Complementary Sequence
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0.0
27600
28000
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28800
29200
29600
Nucleotide Position

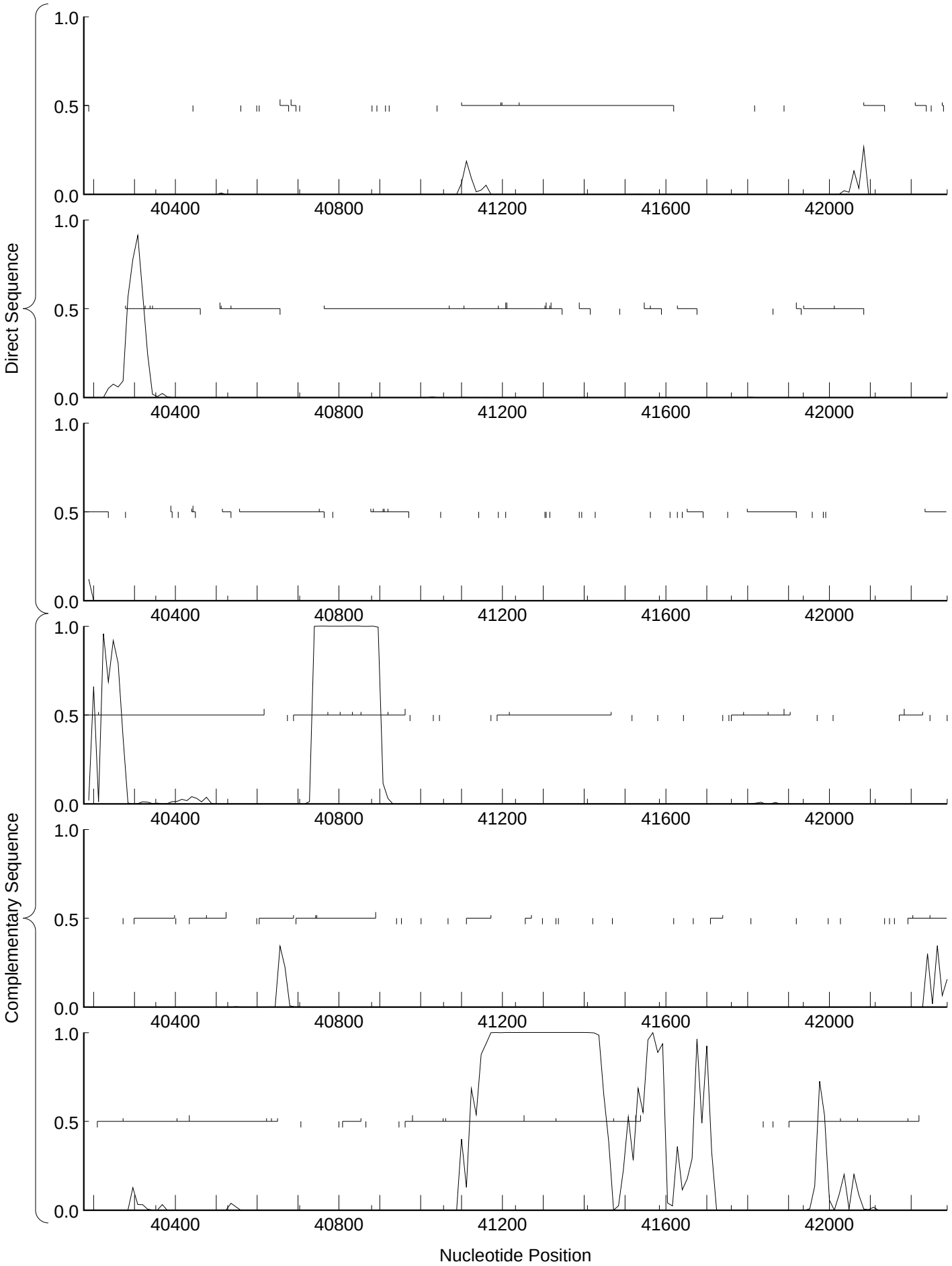

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Complementary Sequence
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31200
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Nucleotide Position

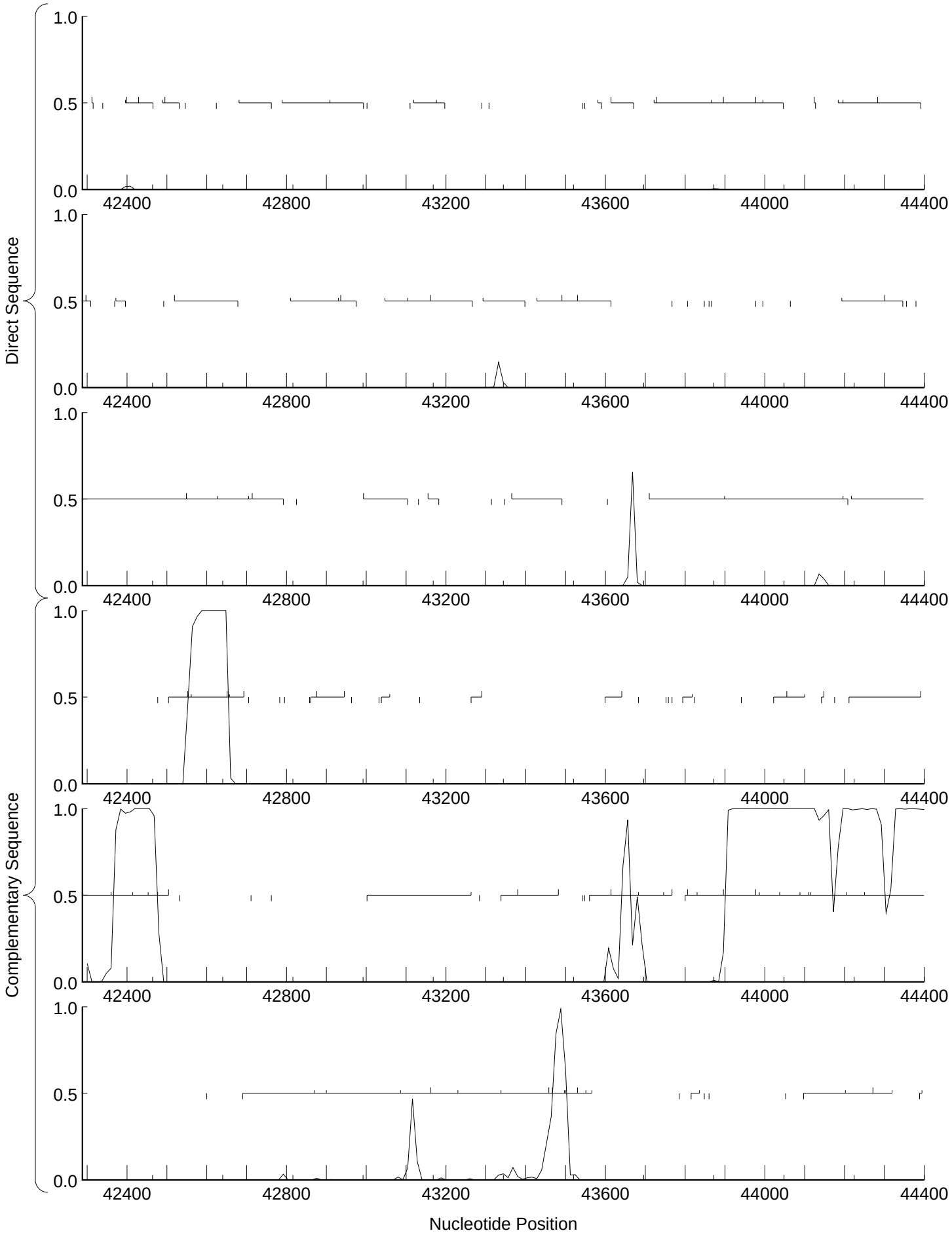

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32800
33200
33600
Nucleotide Position

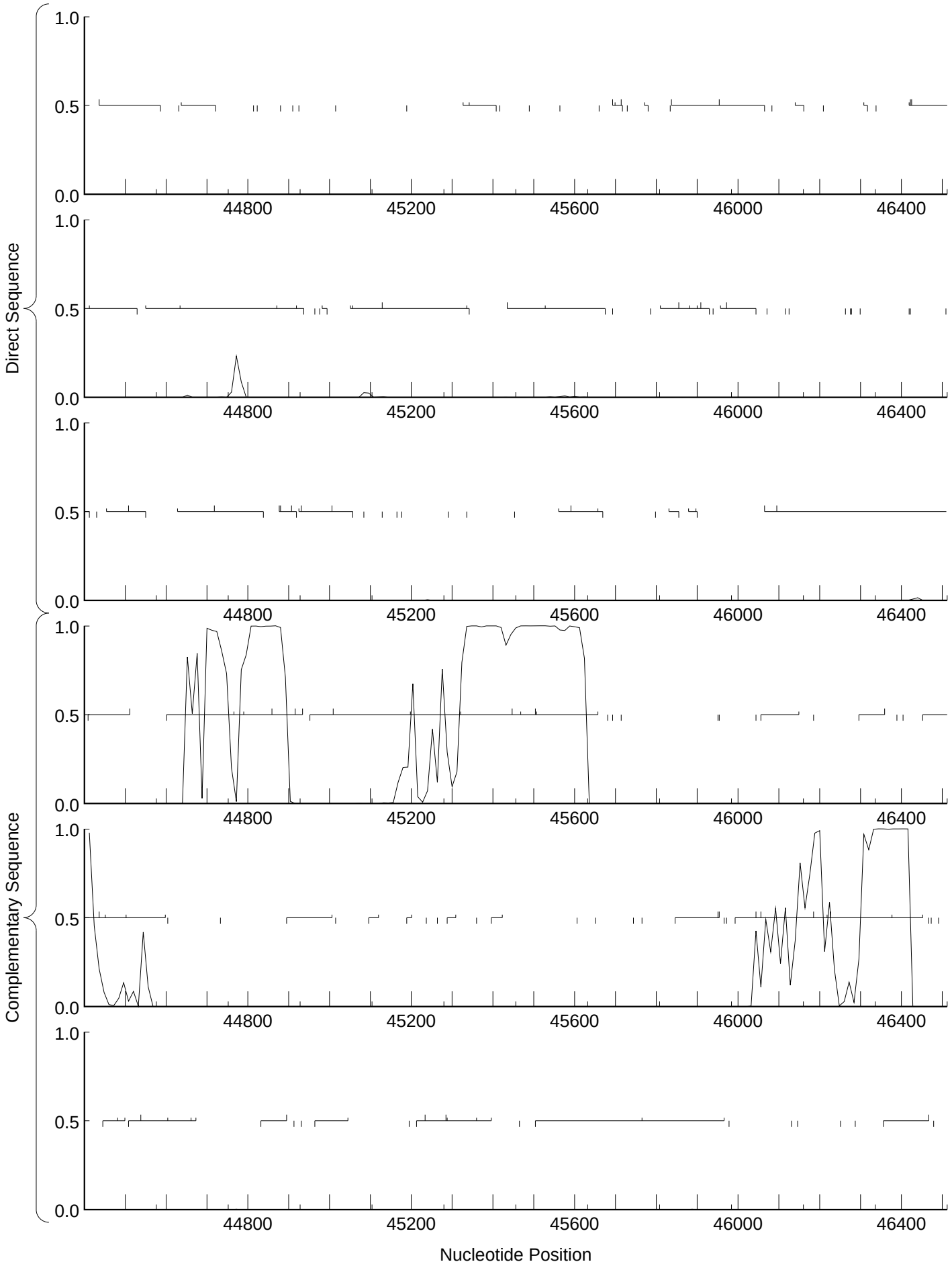

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Complementary Sequence
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Nucleotide Position

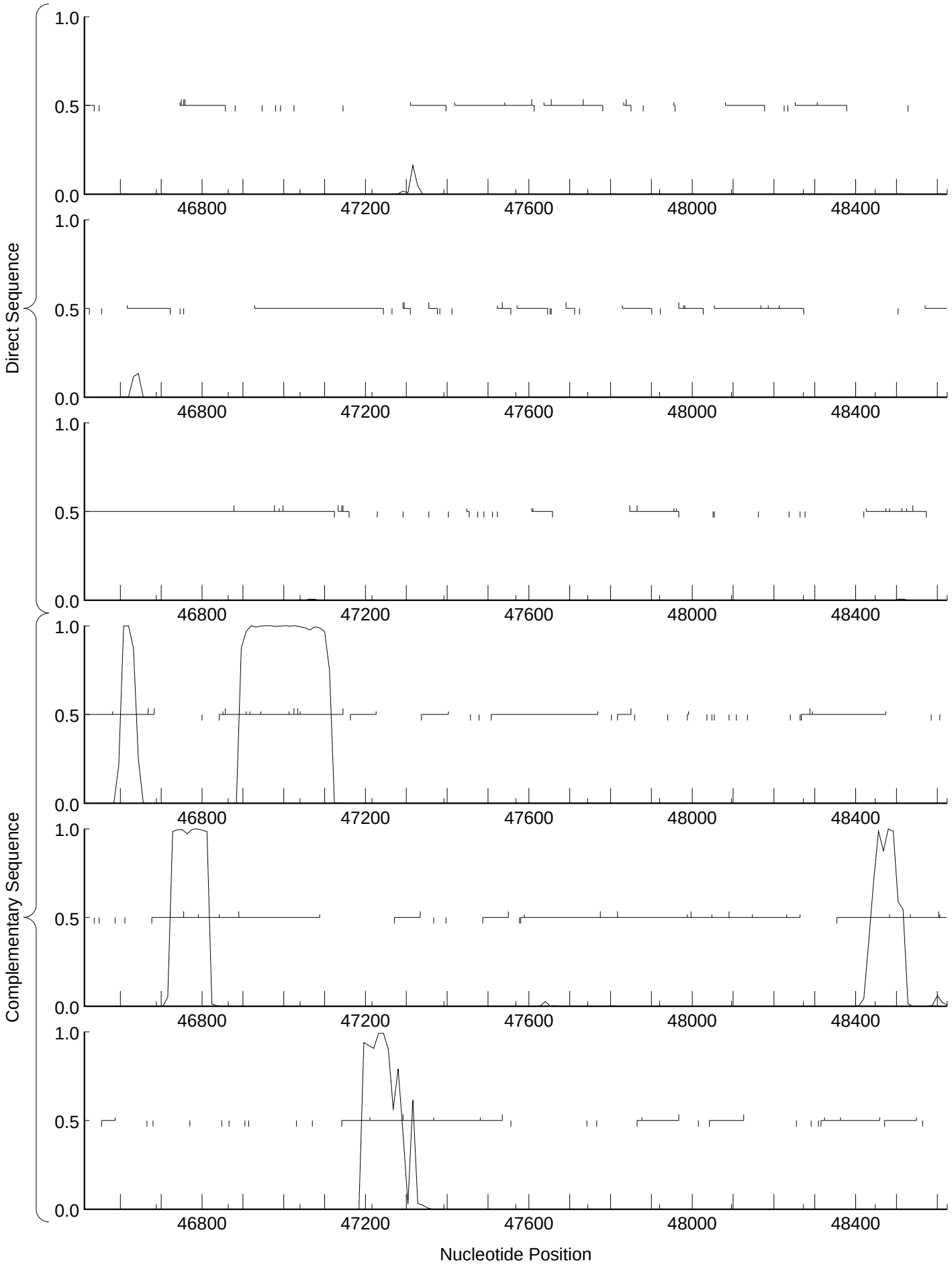

Direct Sequence
Complementary Sequence
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36000
36400
36800
37200
37600
38000
Nucleotide Position

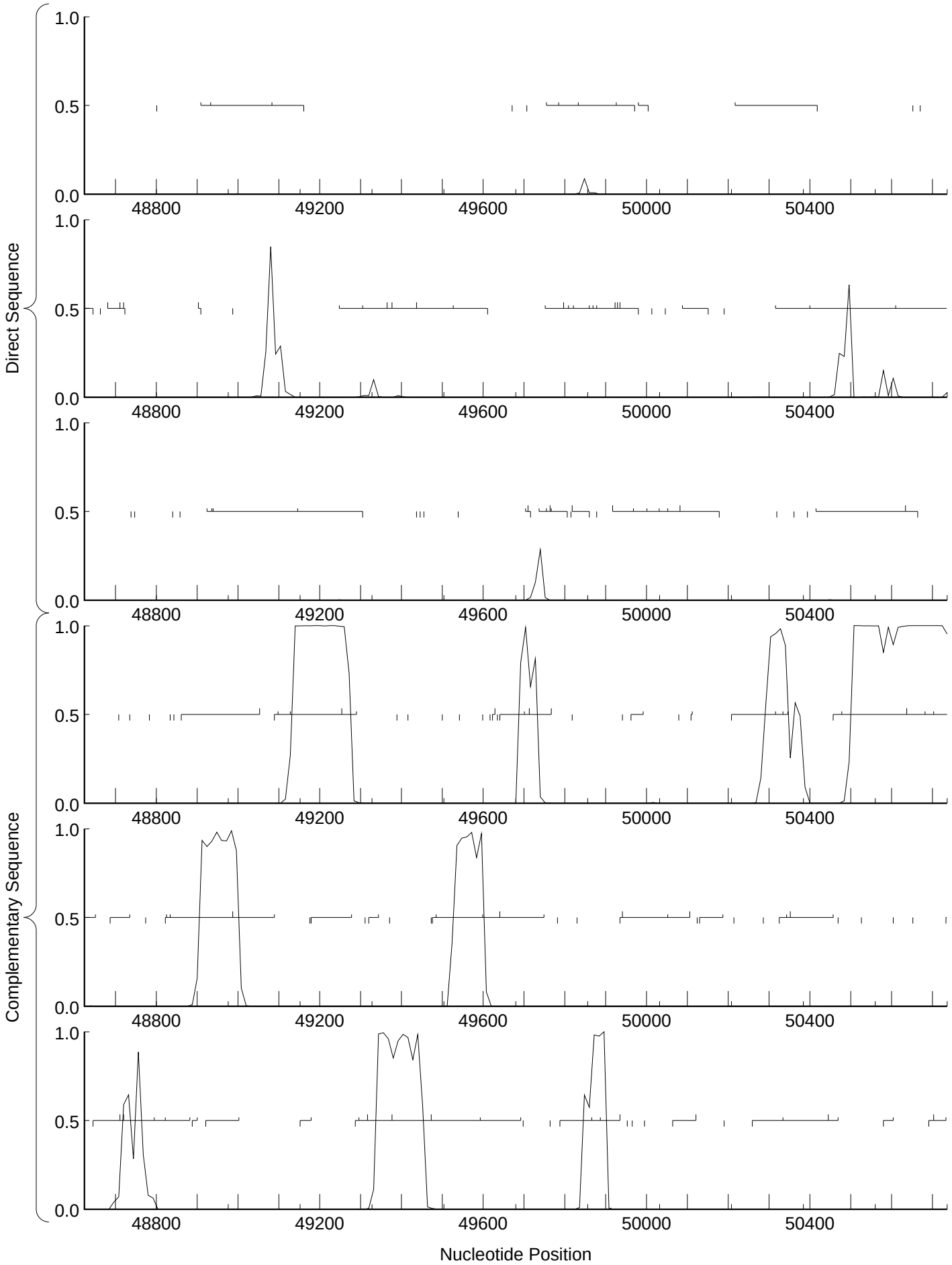

Direct Sequence
Complementary Sequence
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0.0
38400
38800
39200
39600
40000
Nucleotide Position

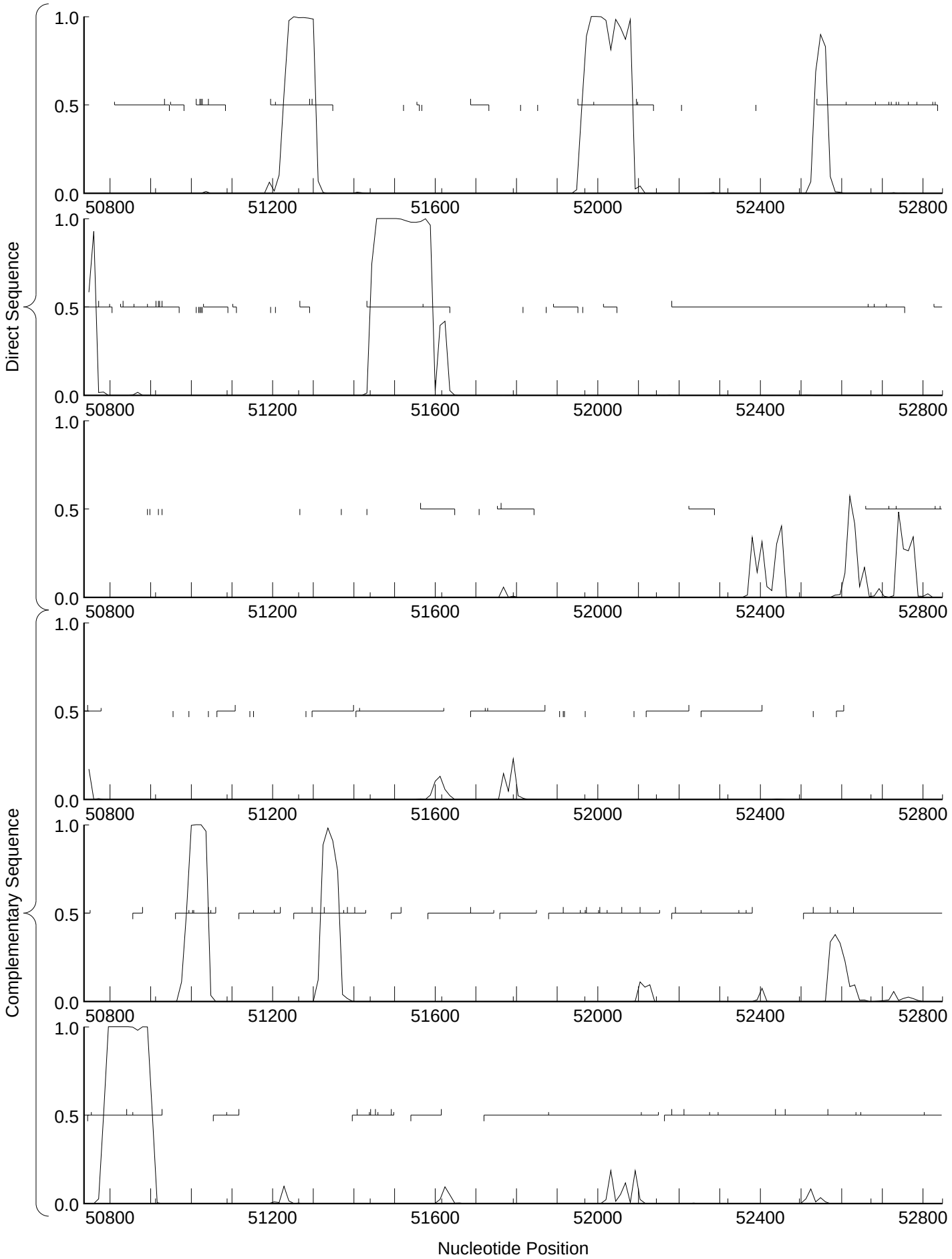

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Nucleotide Position

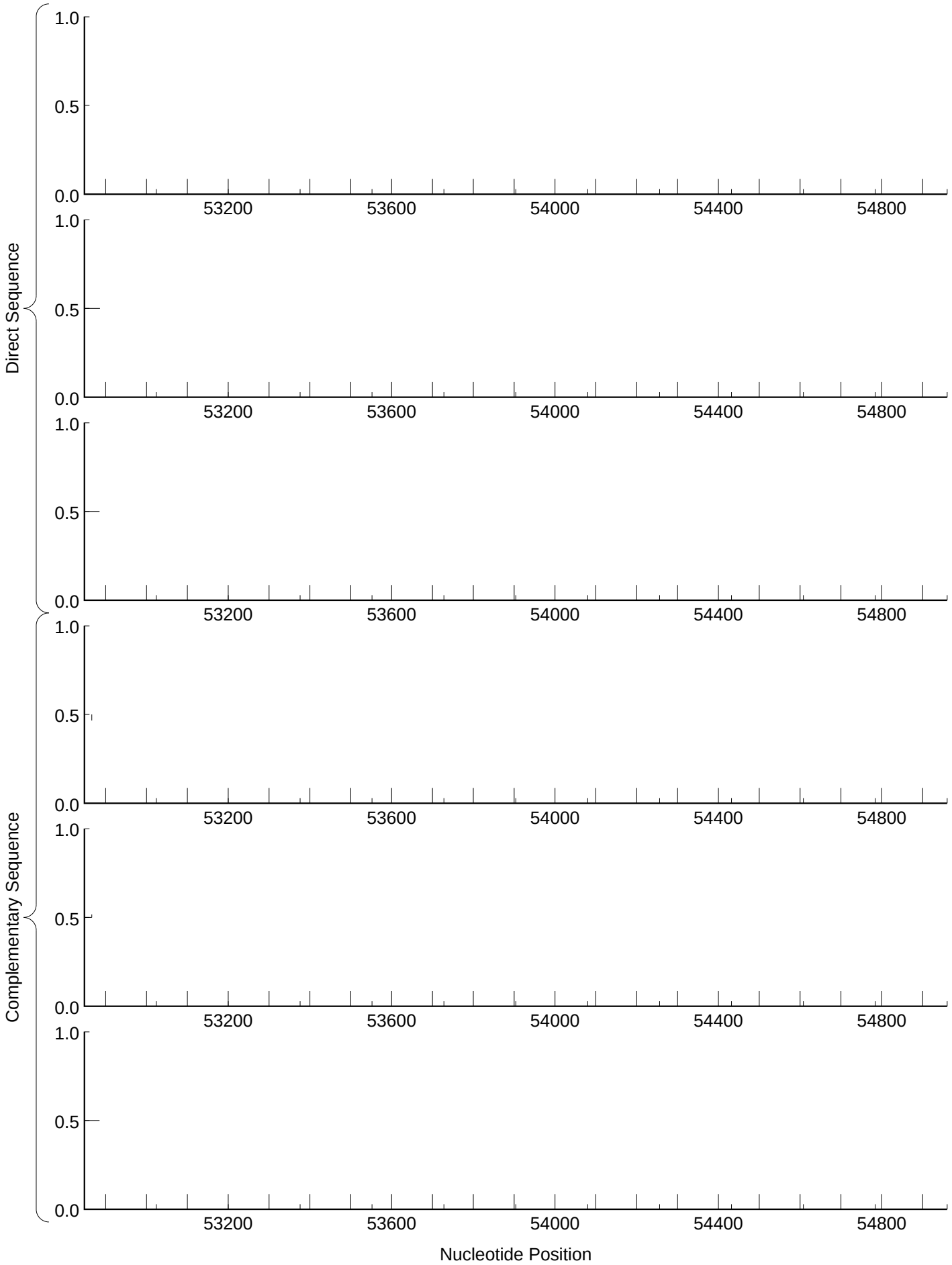

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Complementary Sequence
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0.0
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43200
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44000
44400
Nucleotide Position


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