GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Duplo complete sequence, 52781 bp including 10-base 3’ overhang (CGGTCG
Analysis Date    :  3/29/24 at 10:57:35
Pages            :  26
Sequence Length  :  52781 bp
GC Content       :   63.48%

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

PS-Version       :  1.2

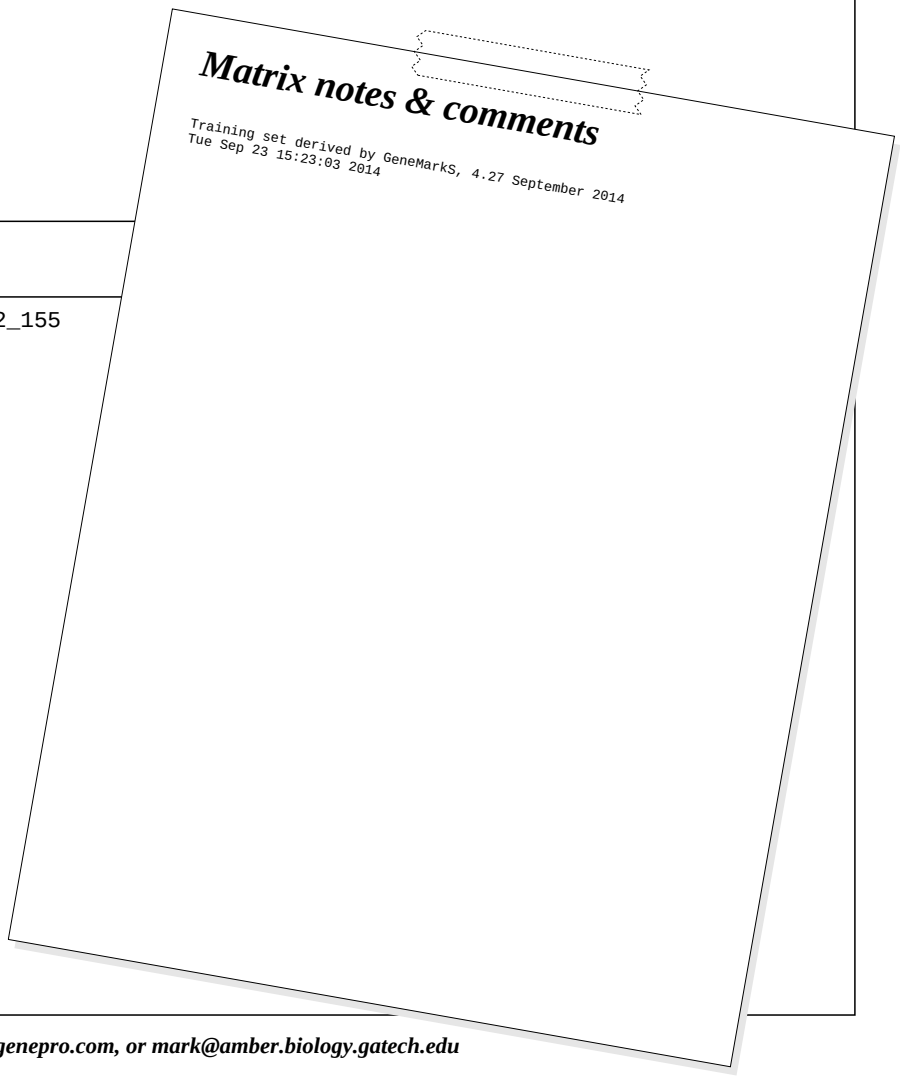
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Author     :  -

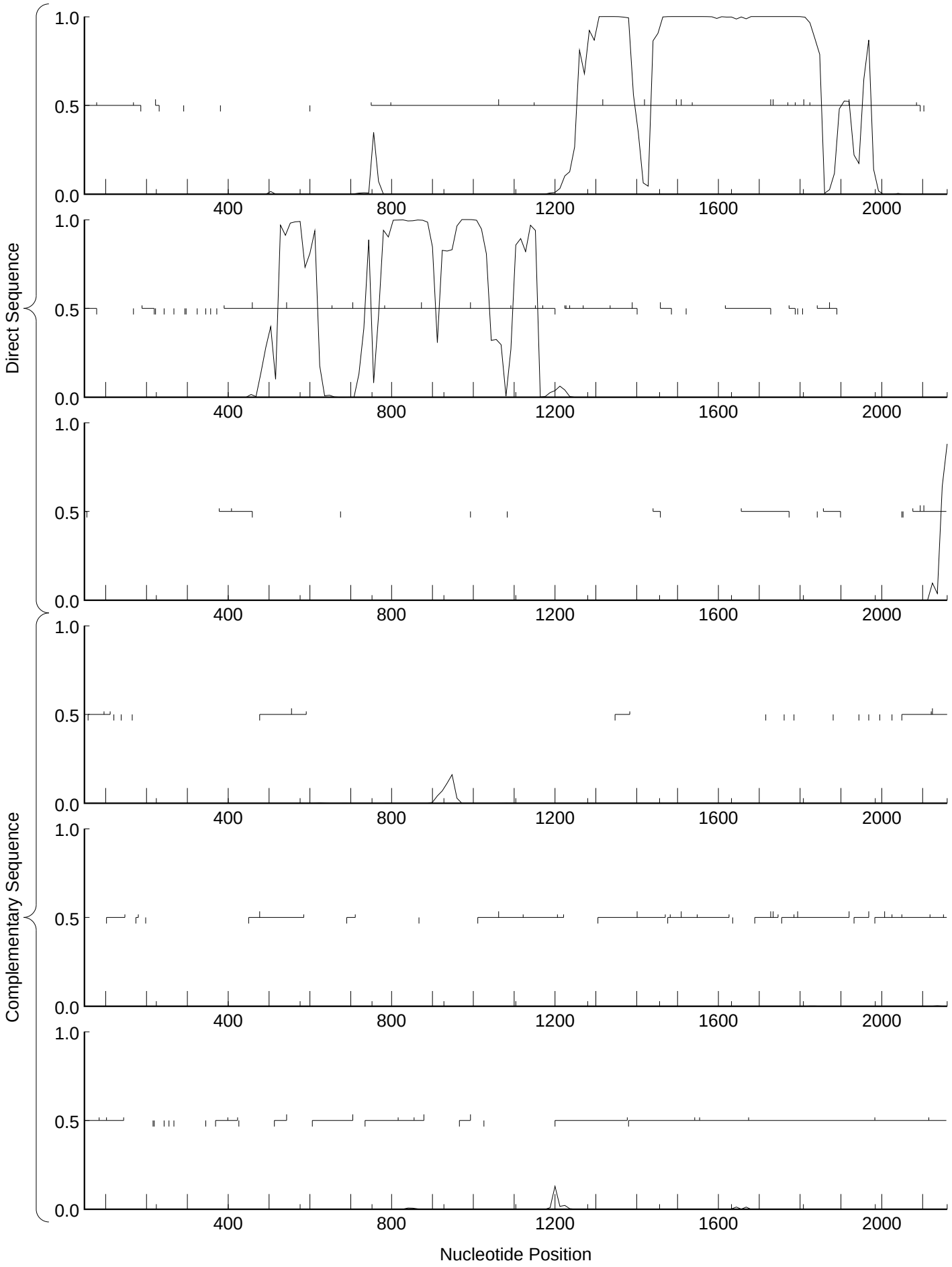
Order      :  4


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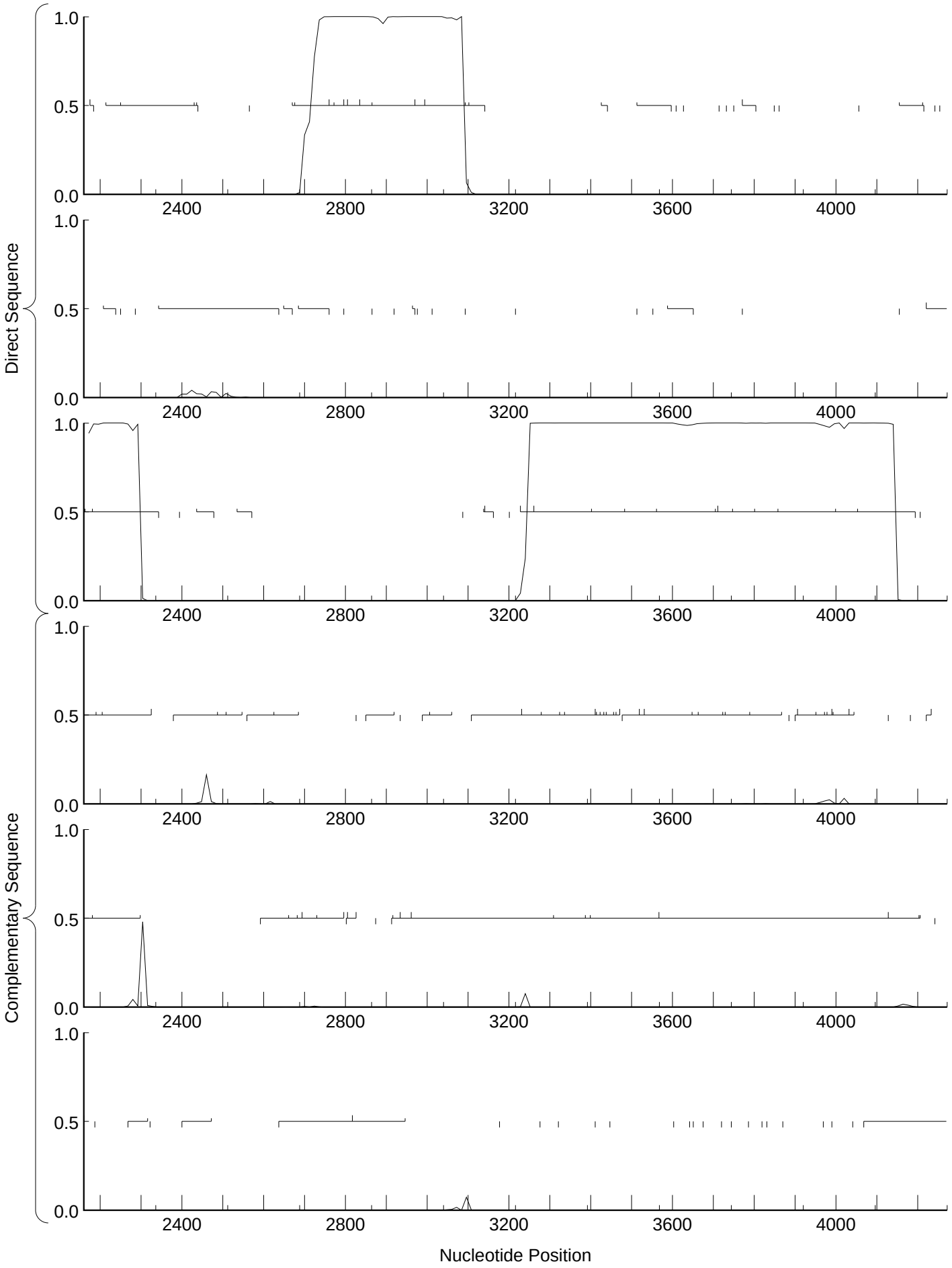


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

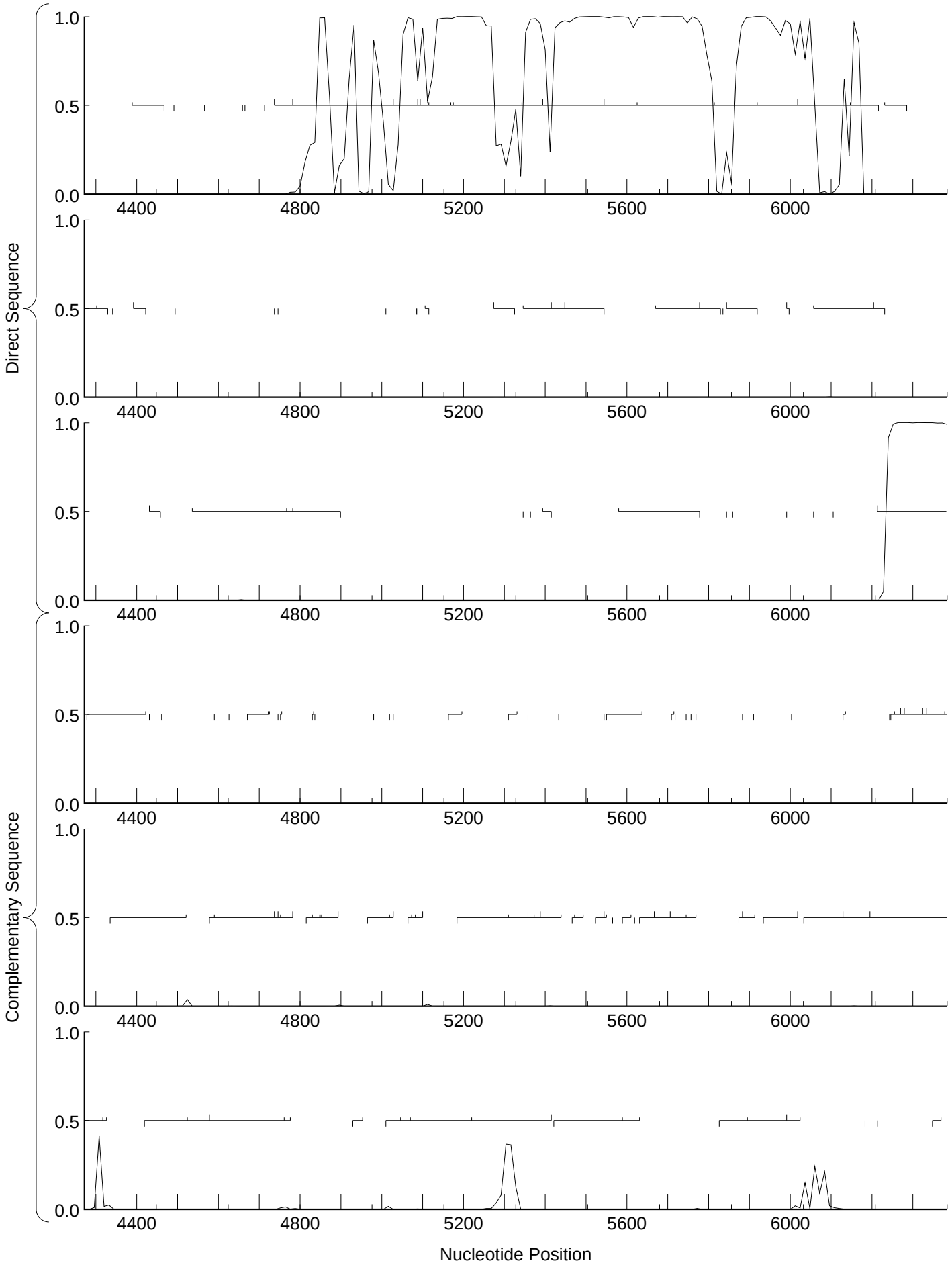
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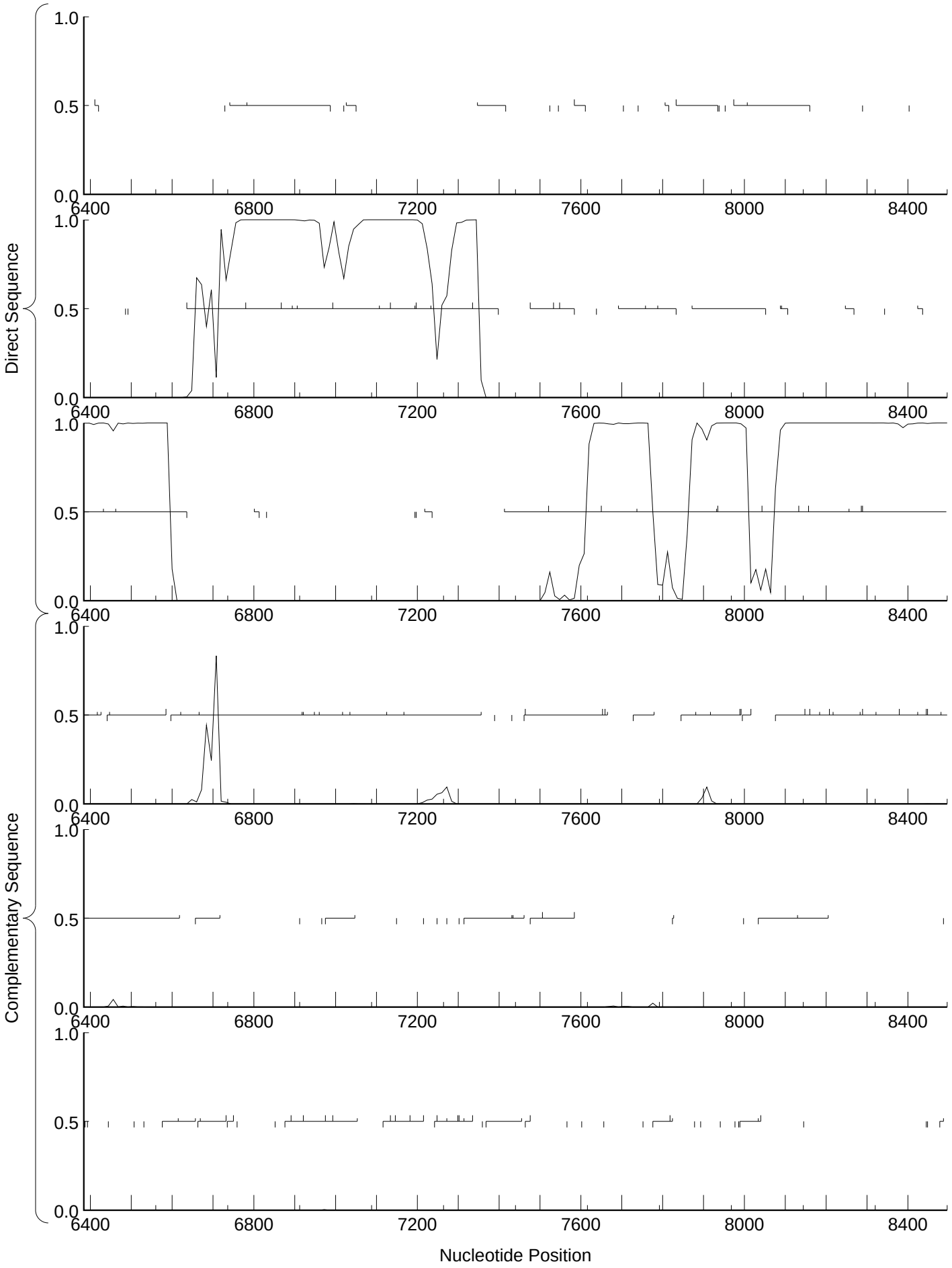
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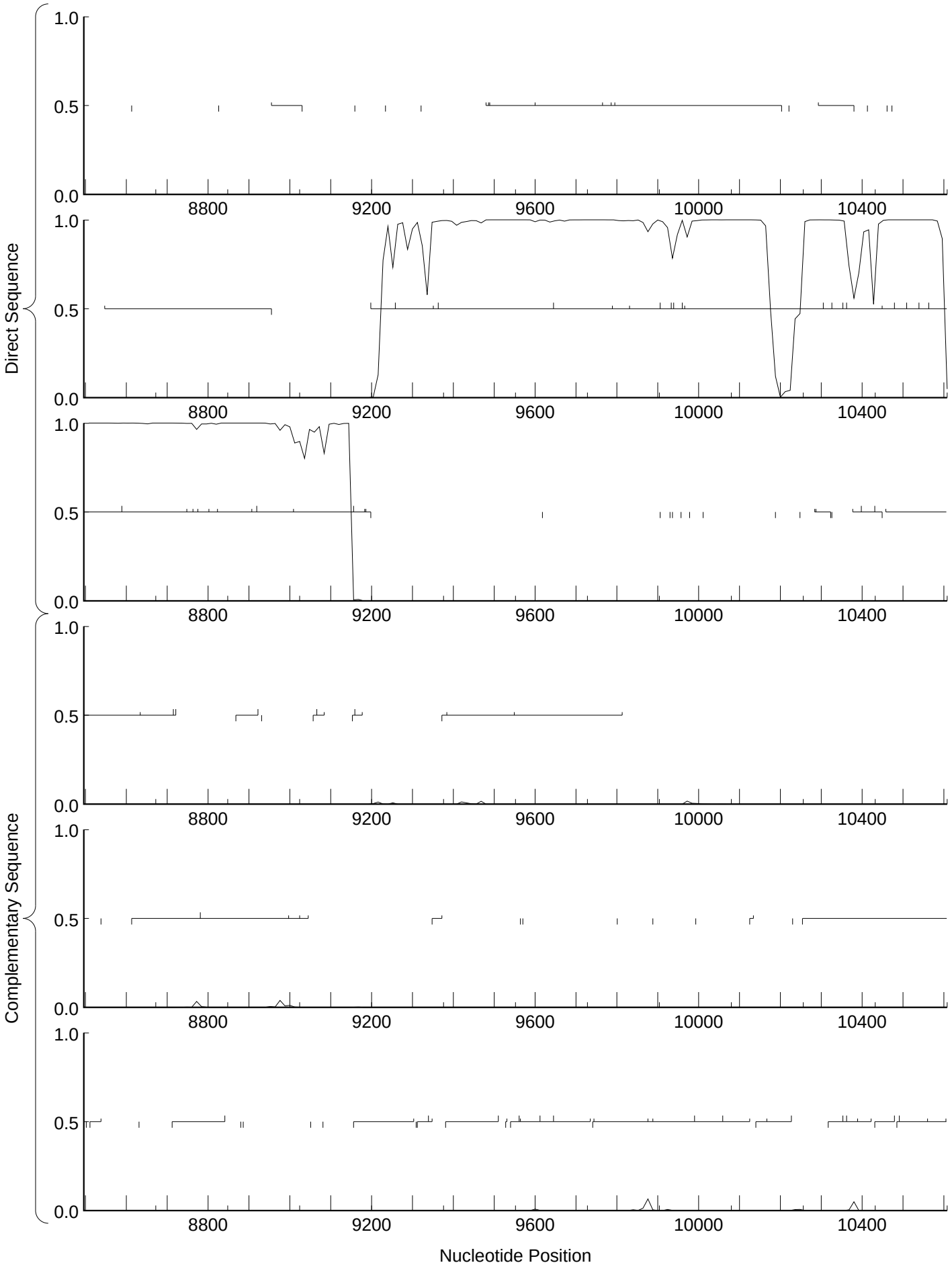
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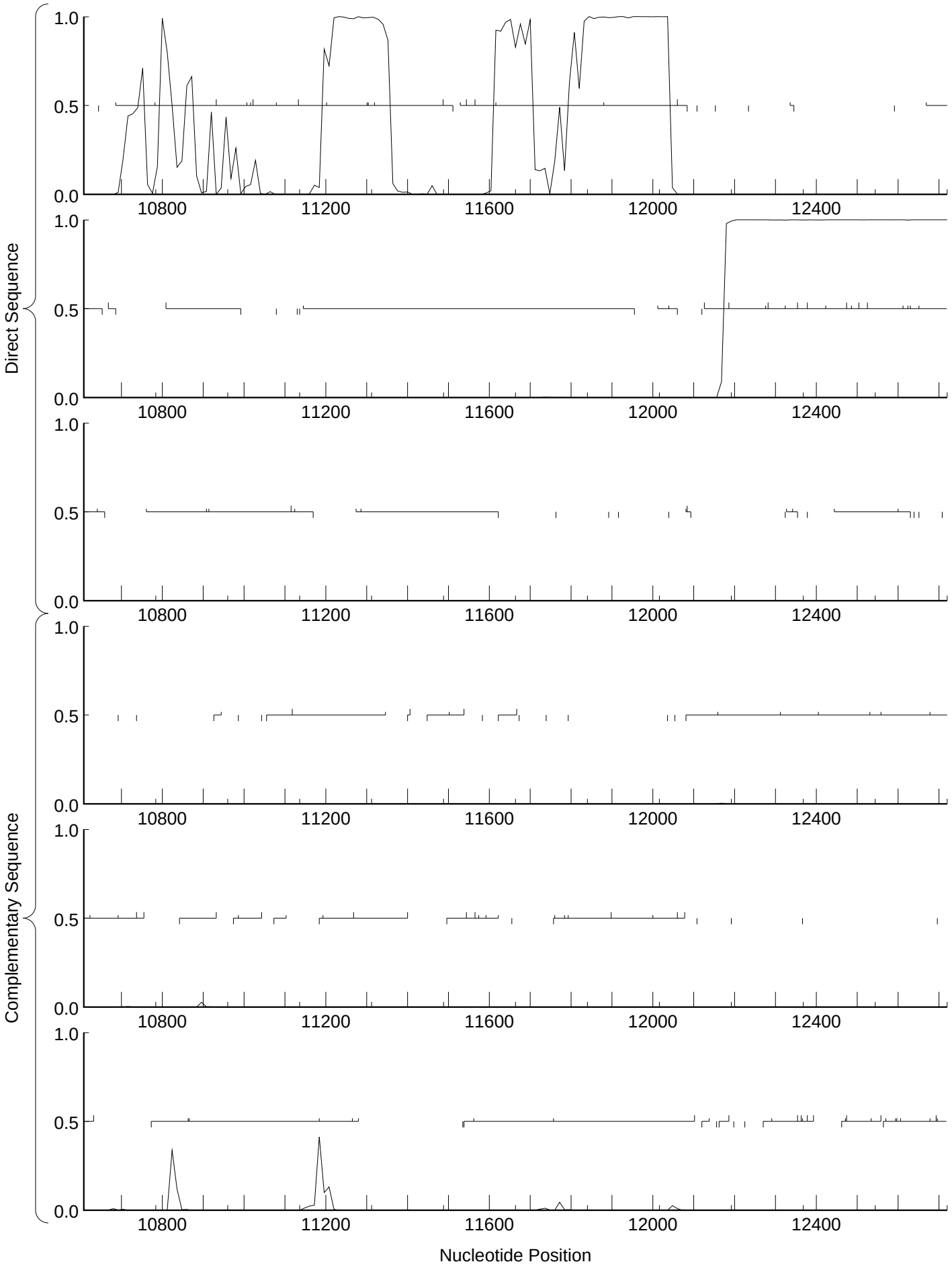
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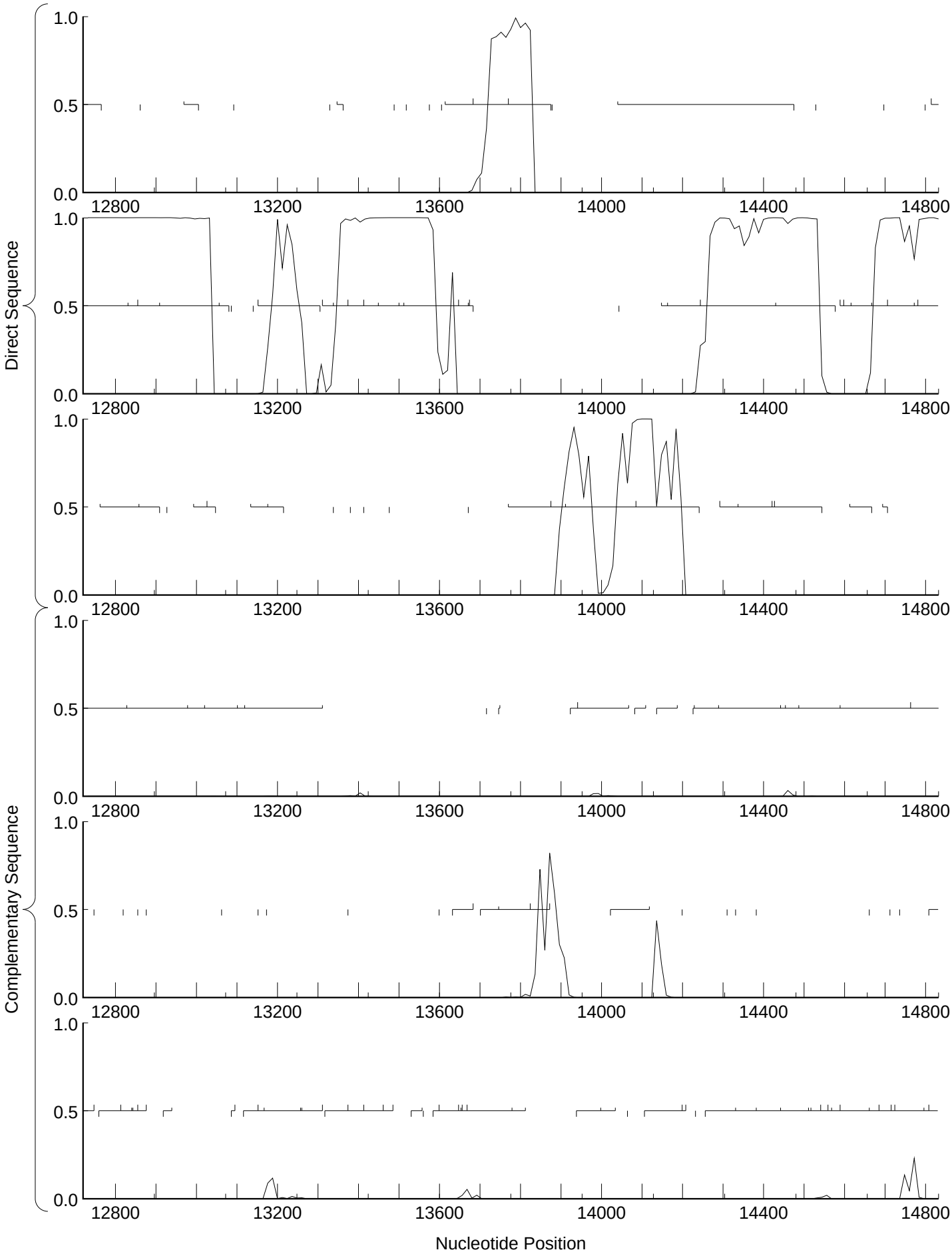
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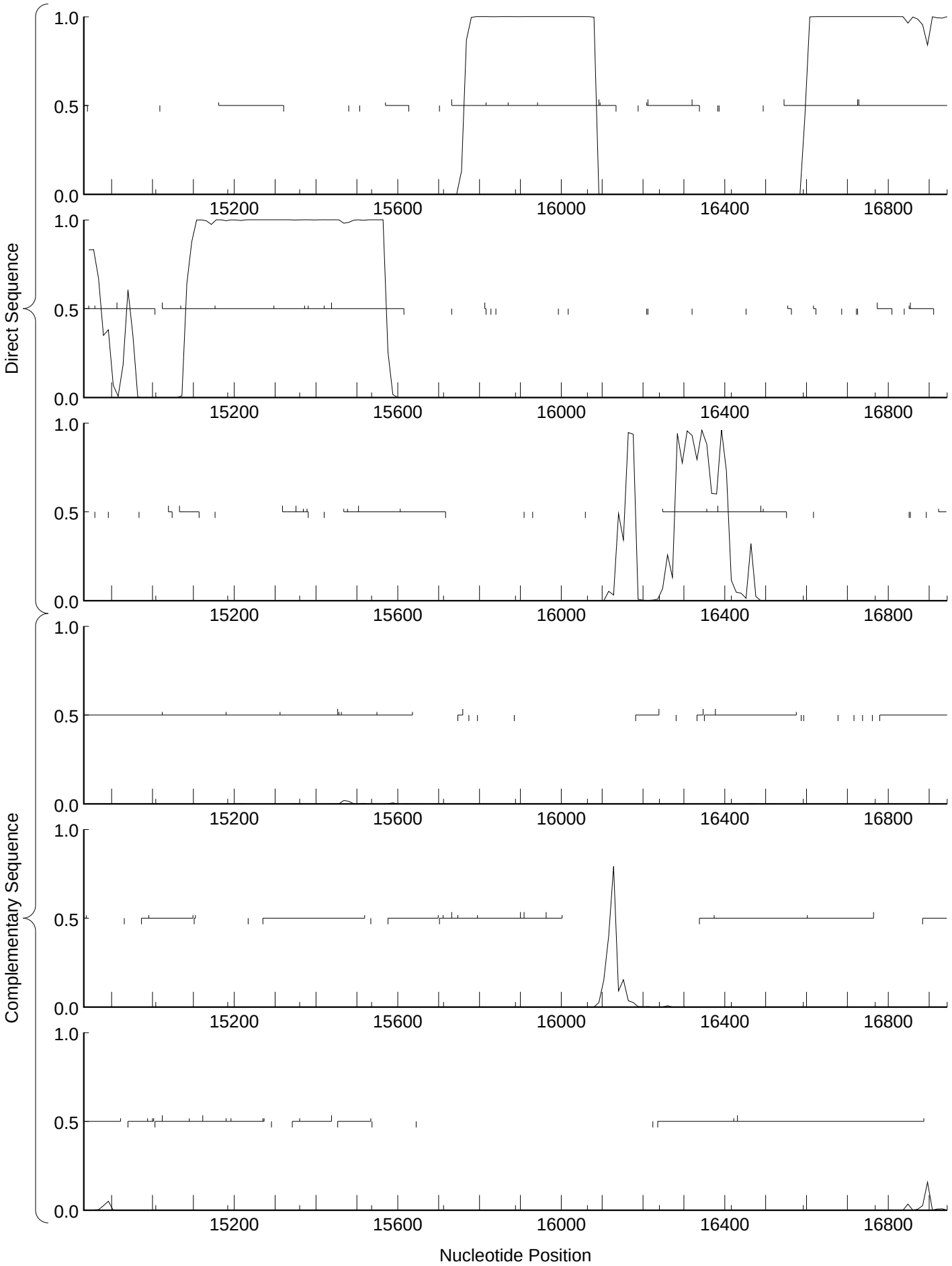
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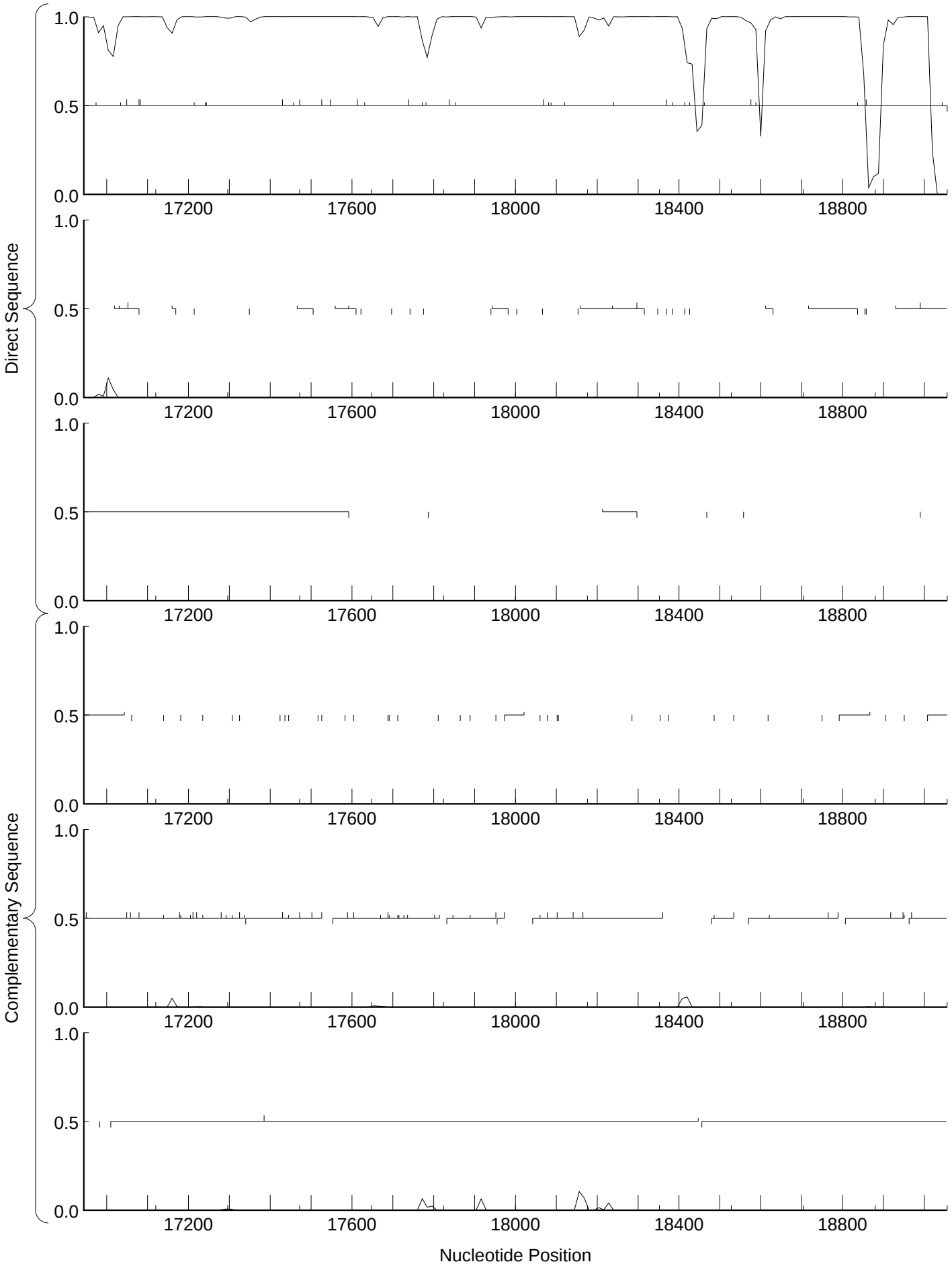


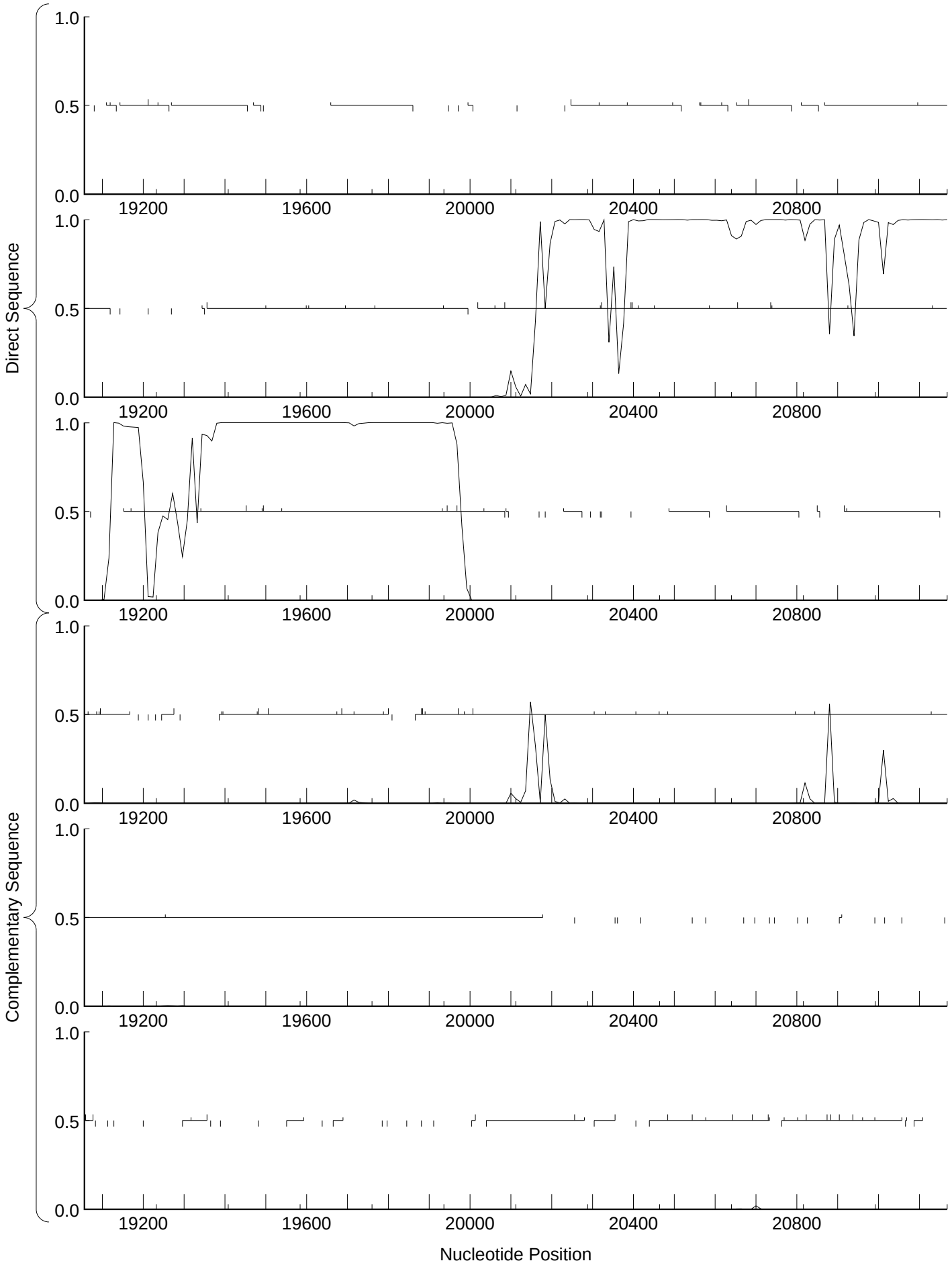
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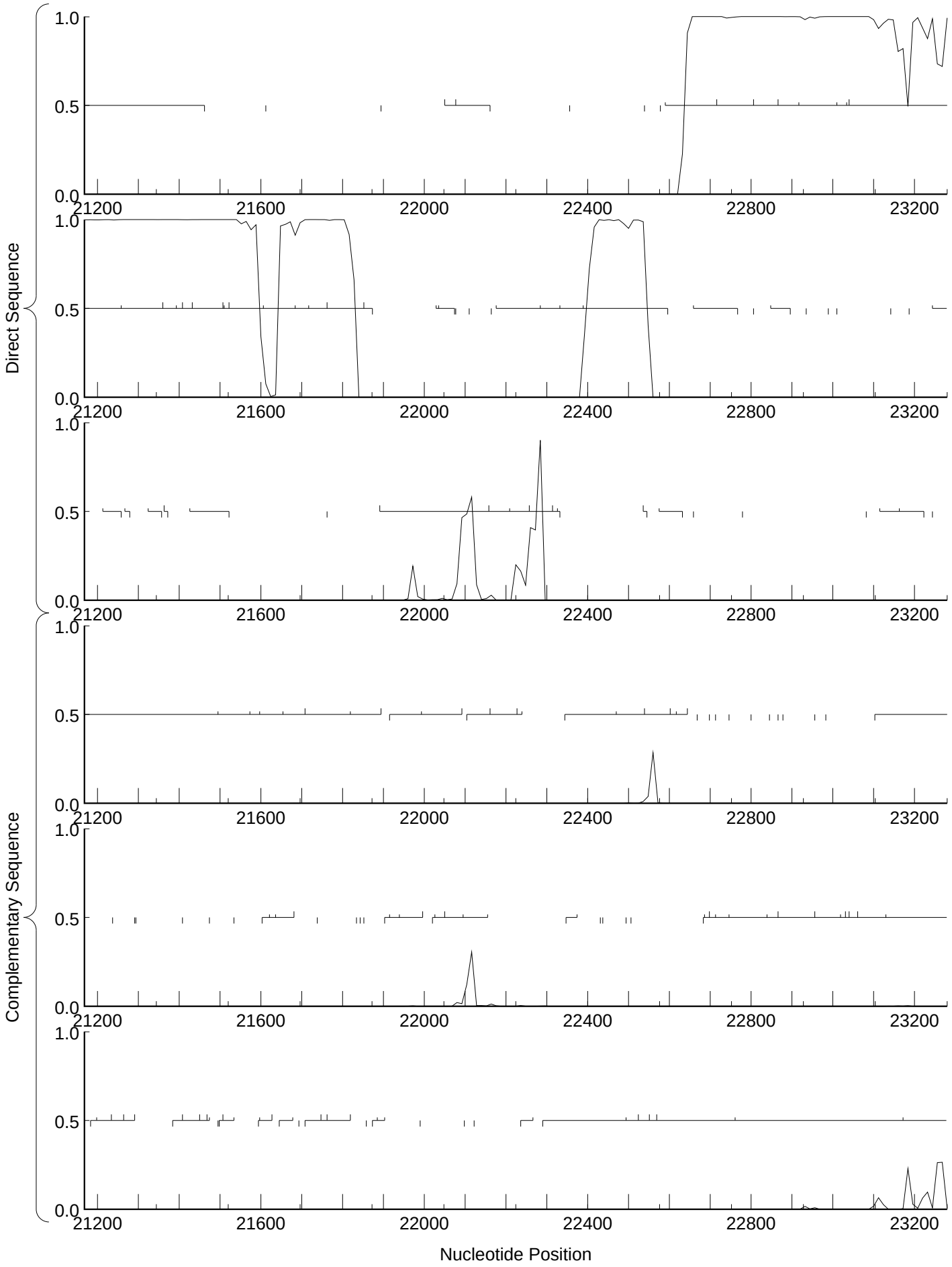


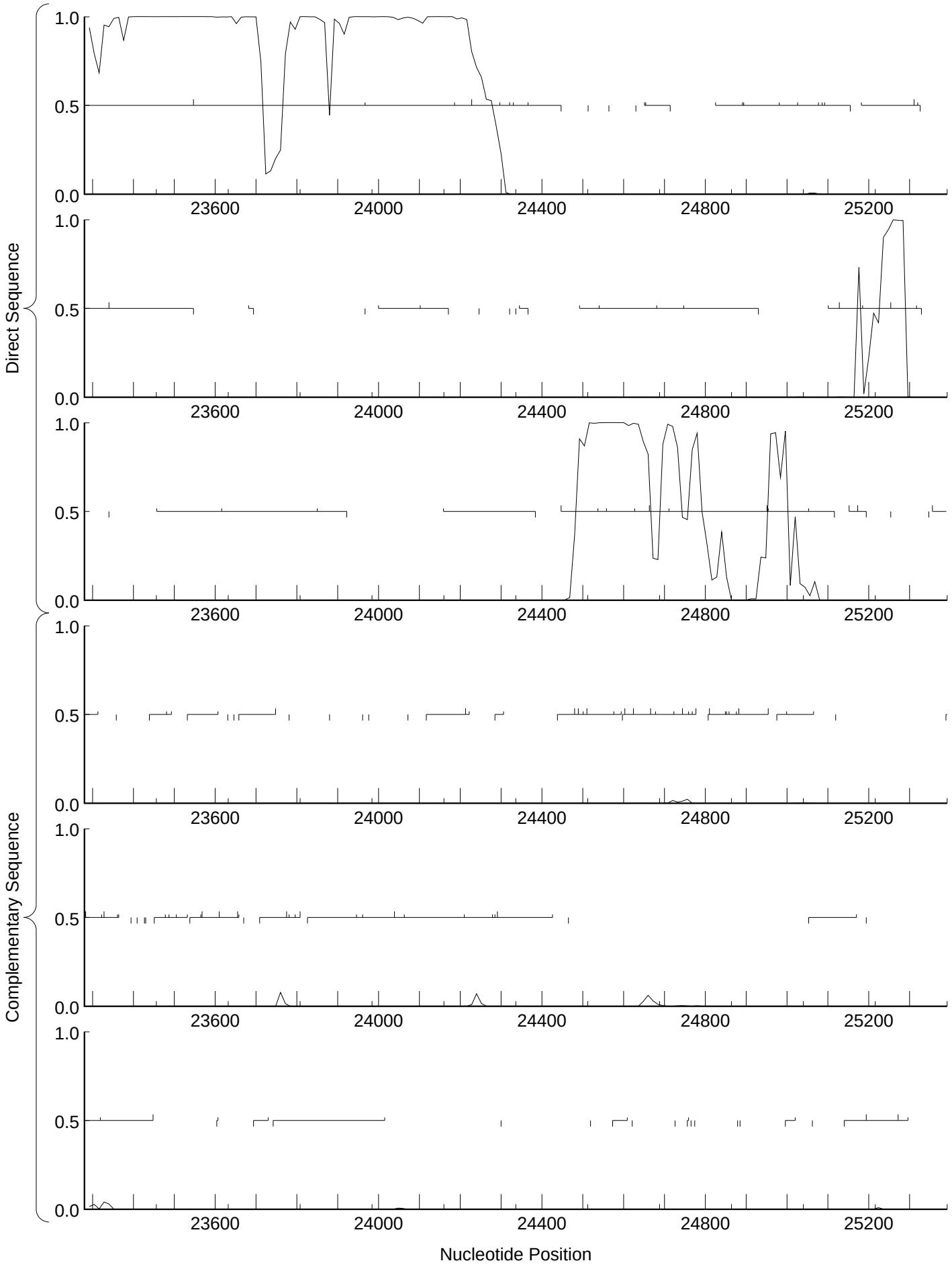
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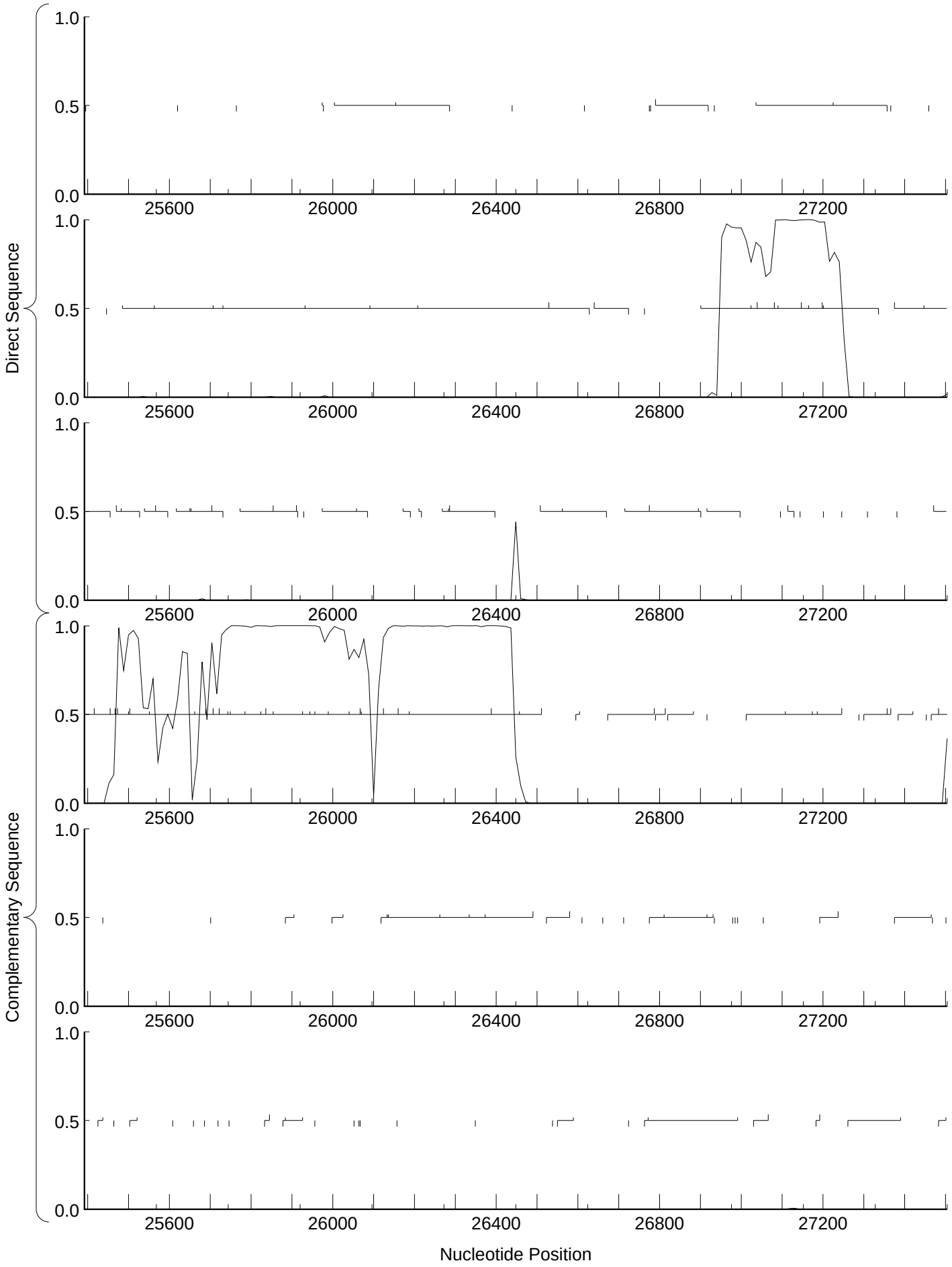


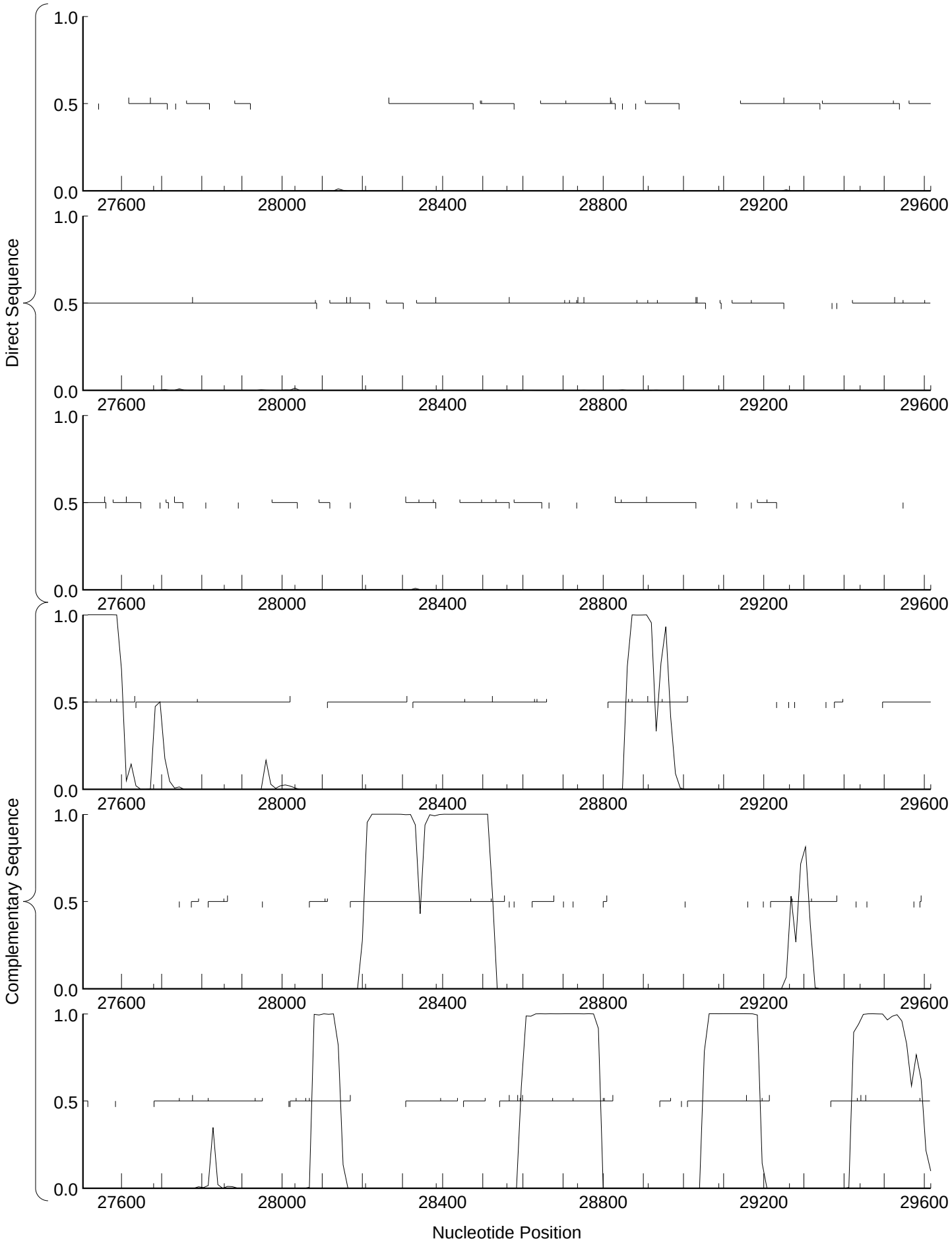

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Complementary Sequence
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0.5
0.0
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17600
18000
18400
18800
Nucleotide Position

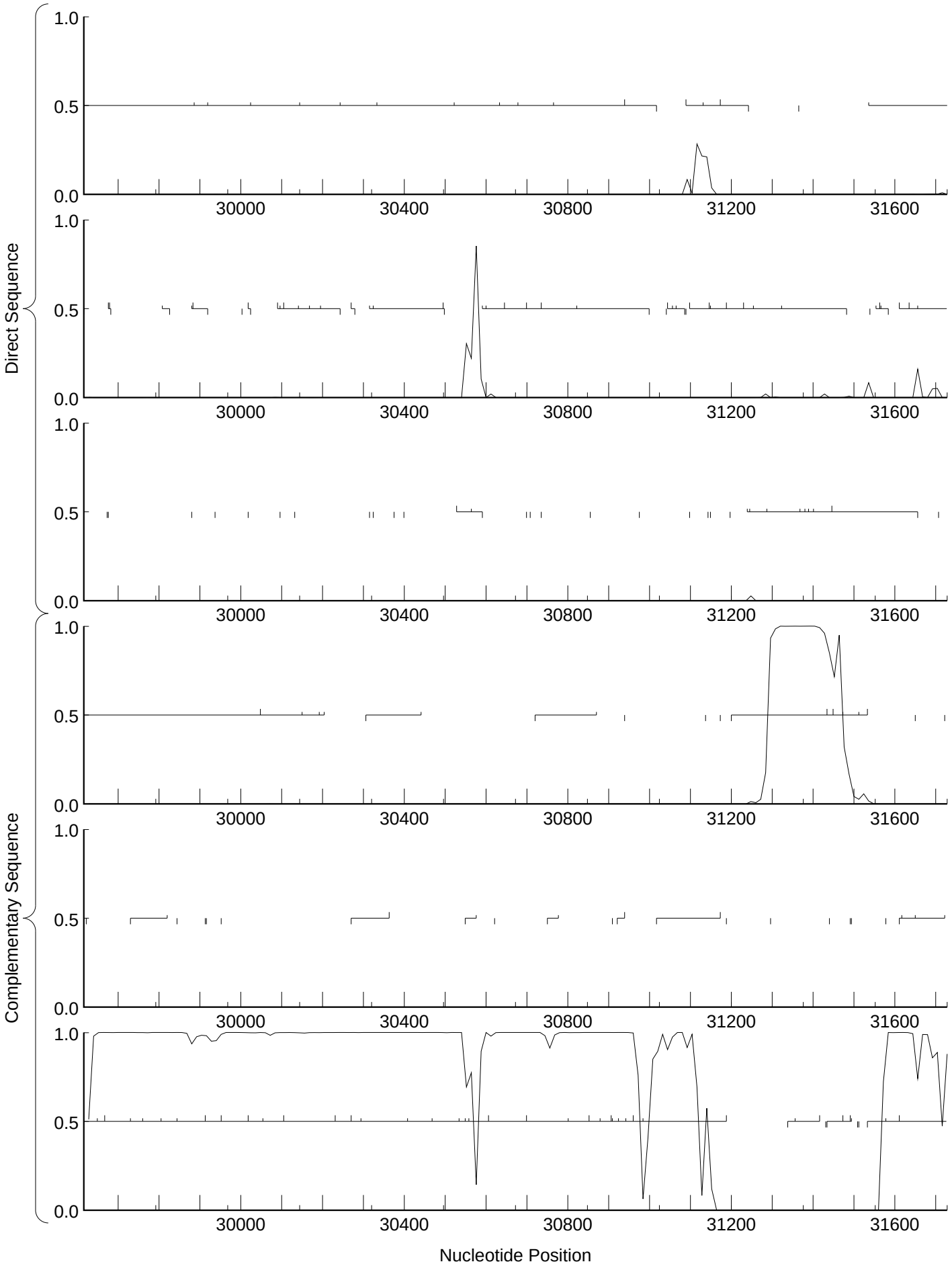

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20000
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20800
Nucleotide Position

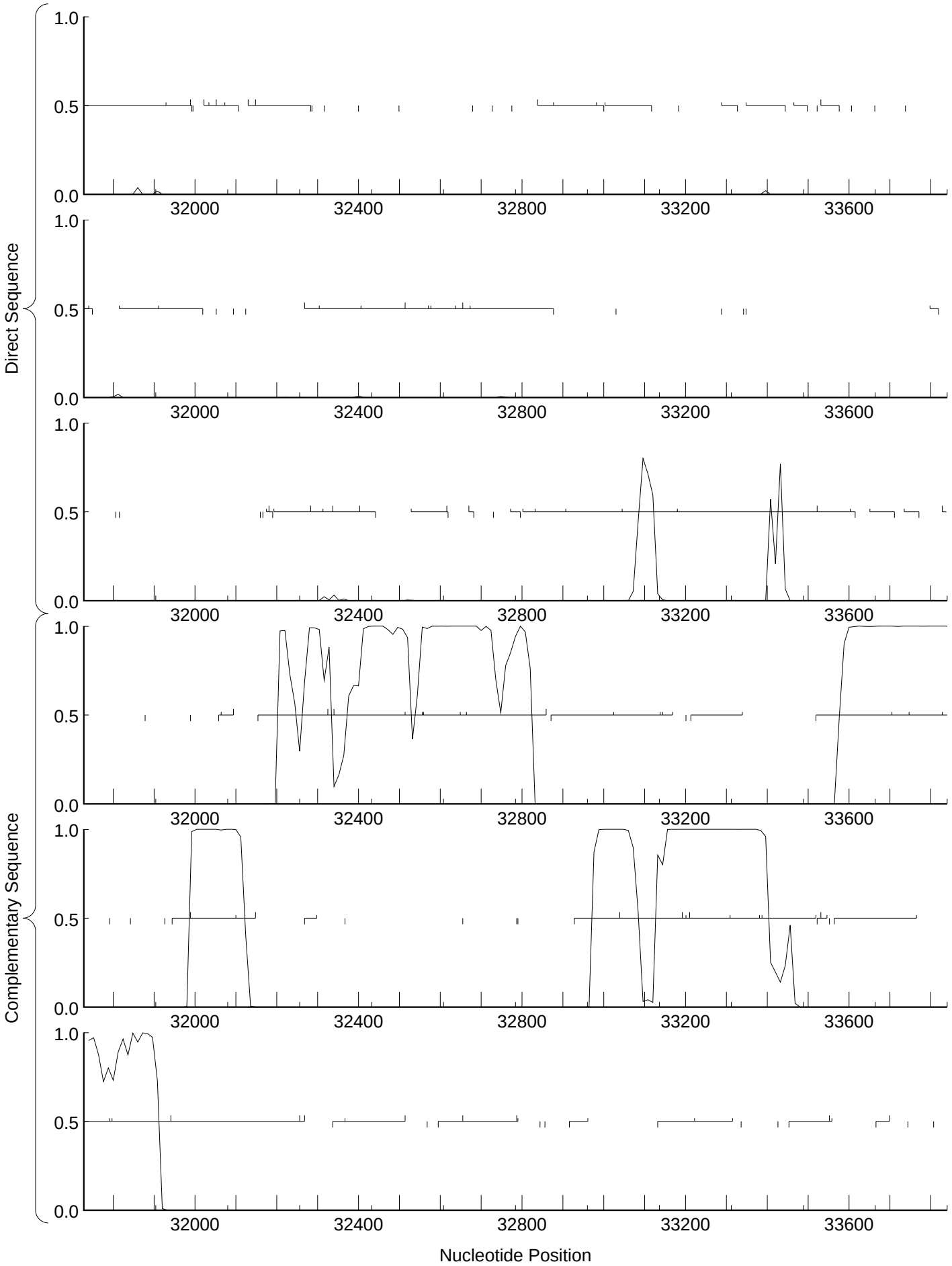

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21600
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22800
23200
Nucleotide Position

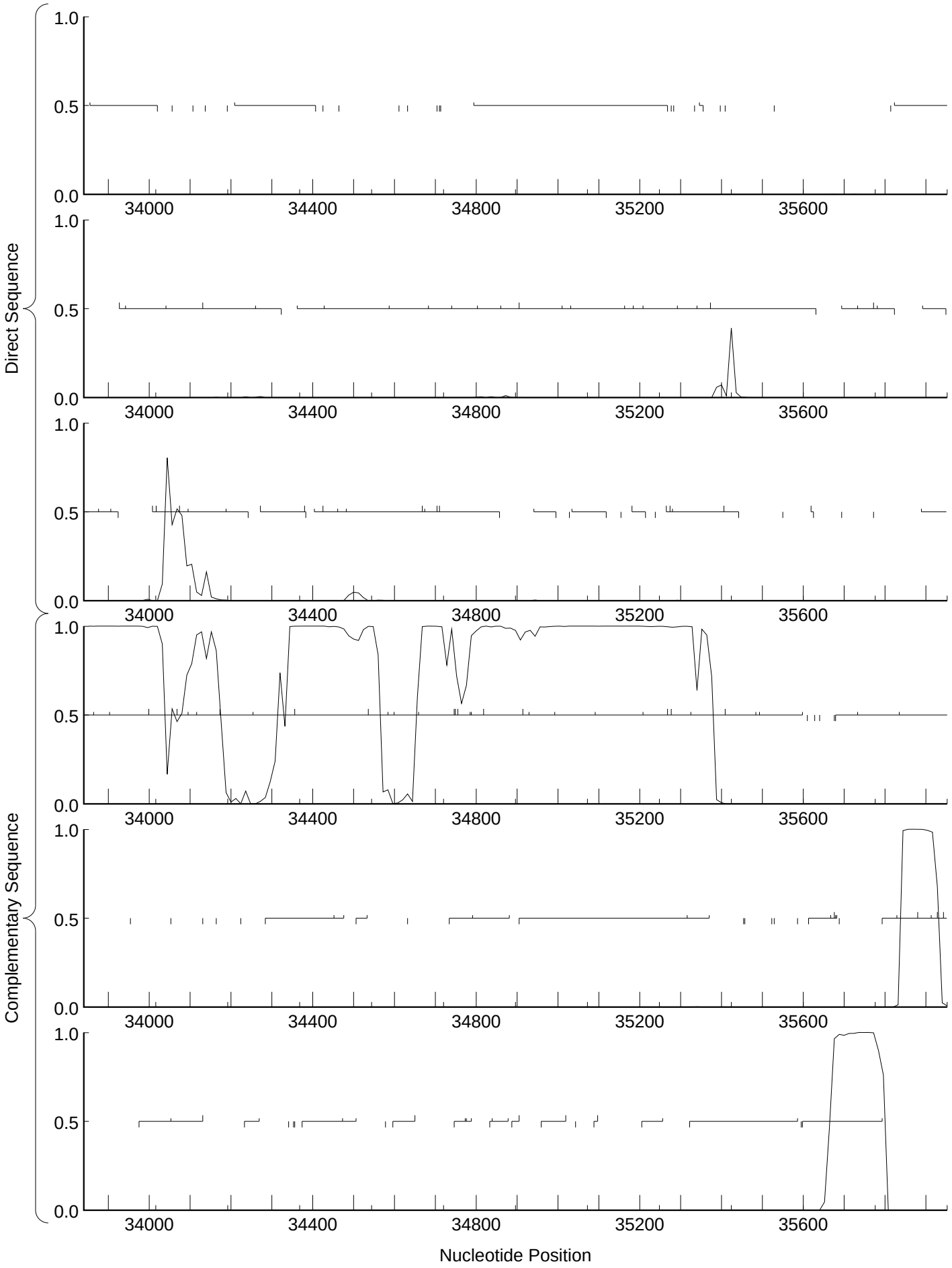

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Complementary Sequence
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24800
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Nucleotide Position

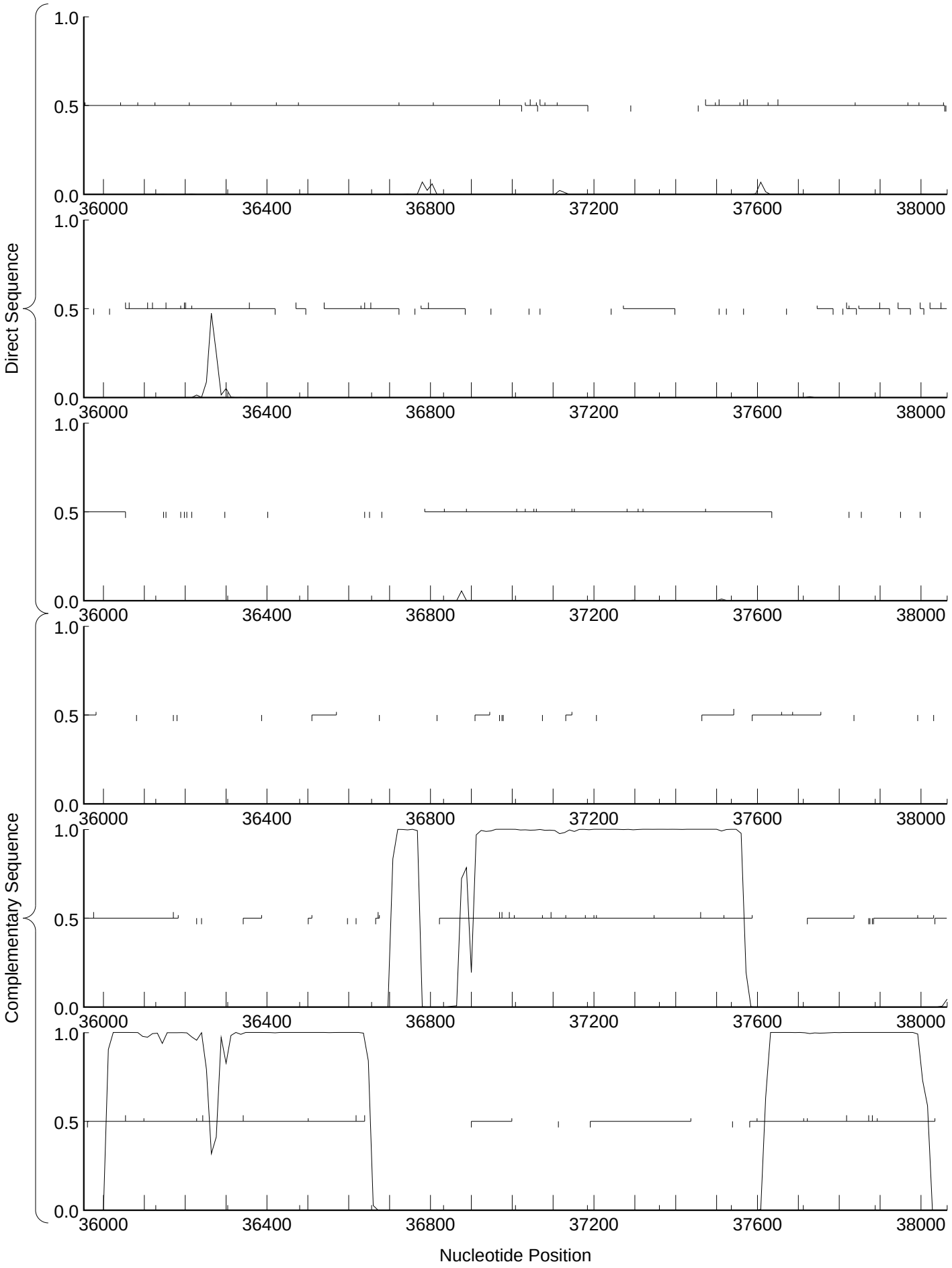

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26400
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Nucleotide Position

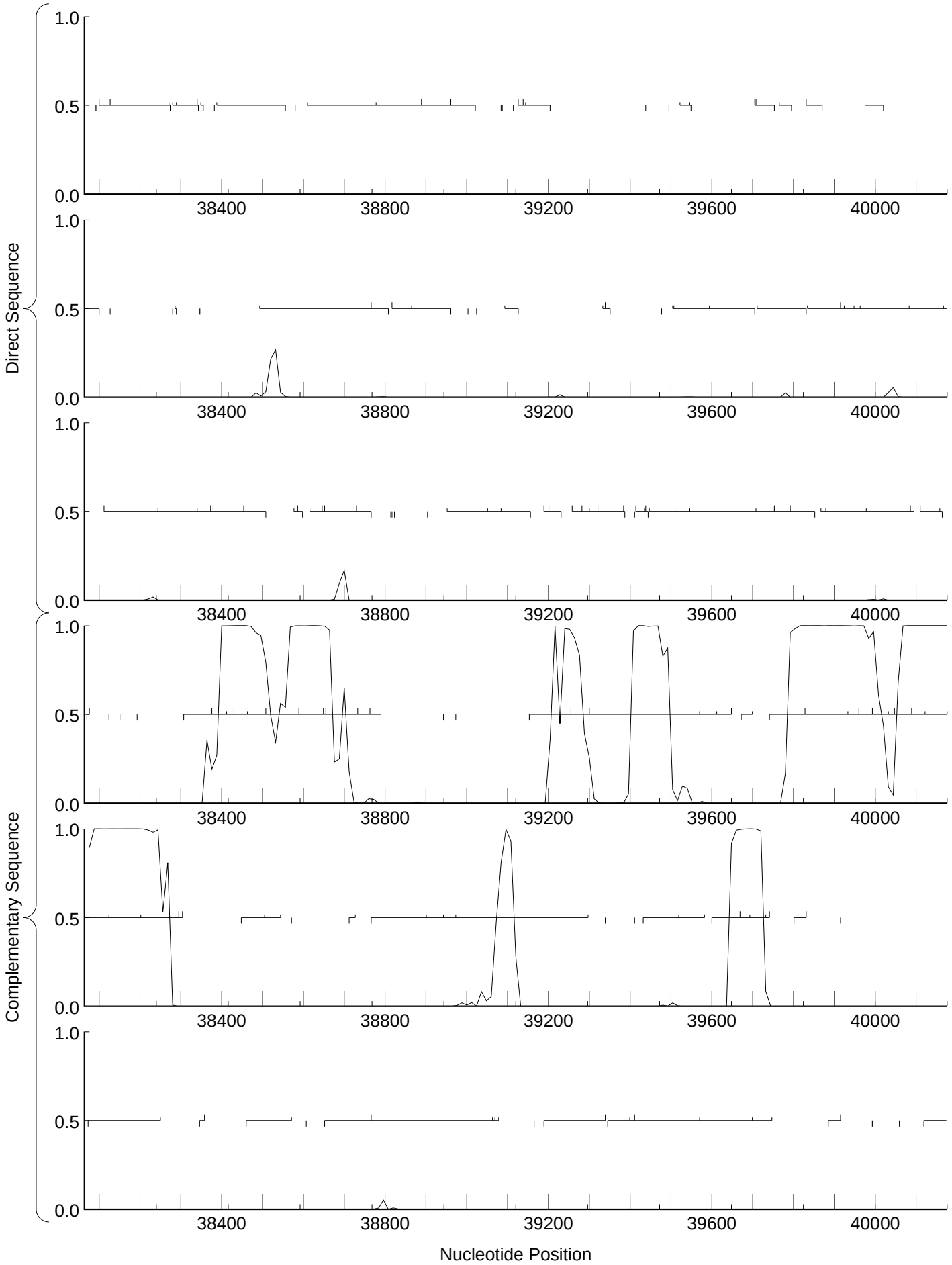

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Nucleotide Position

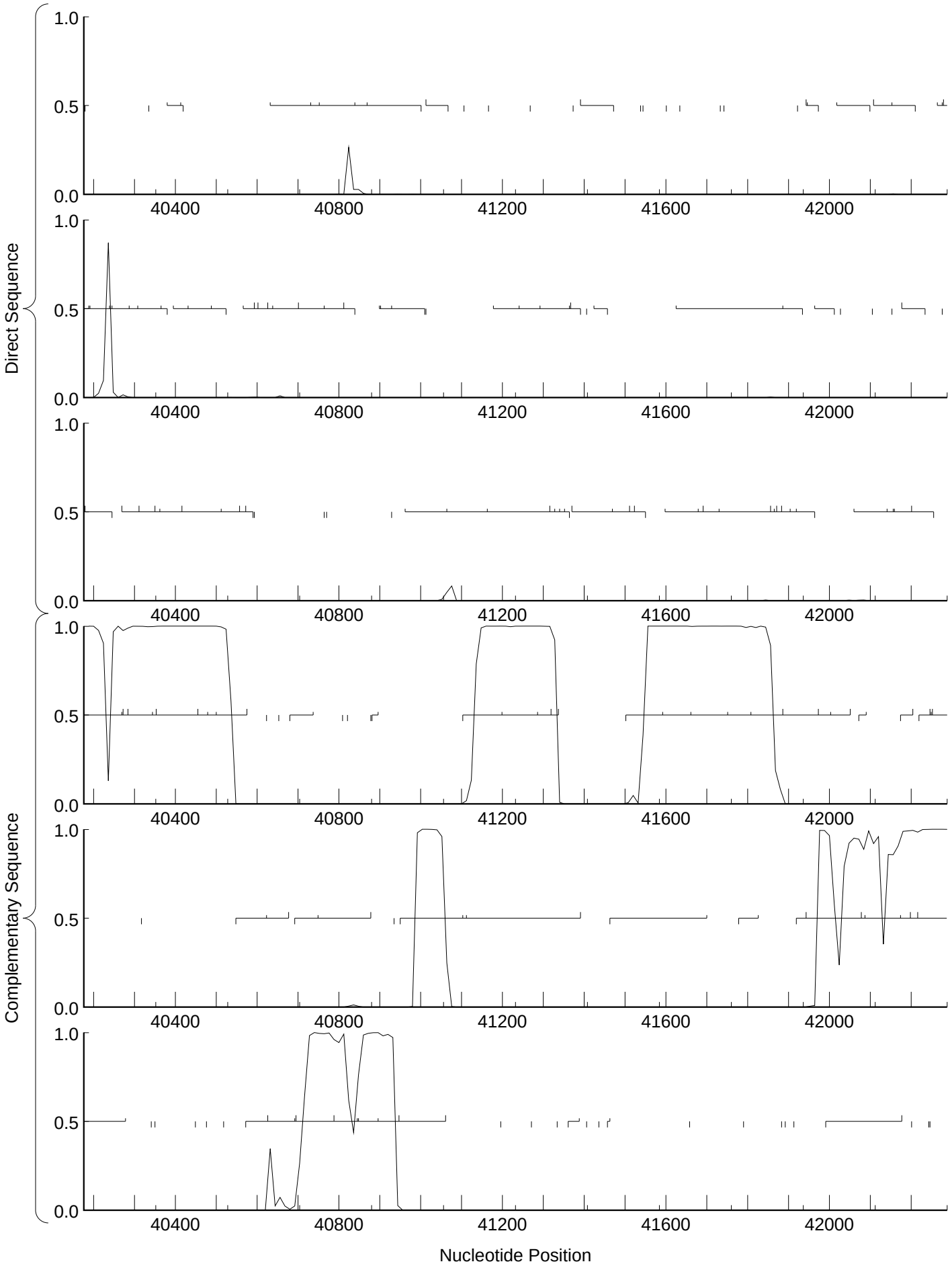

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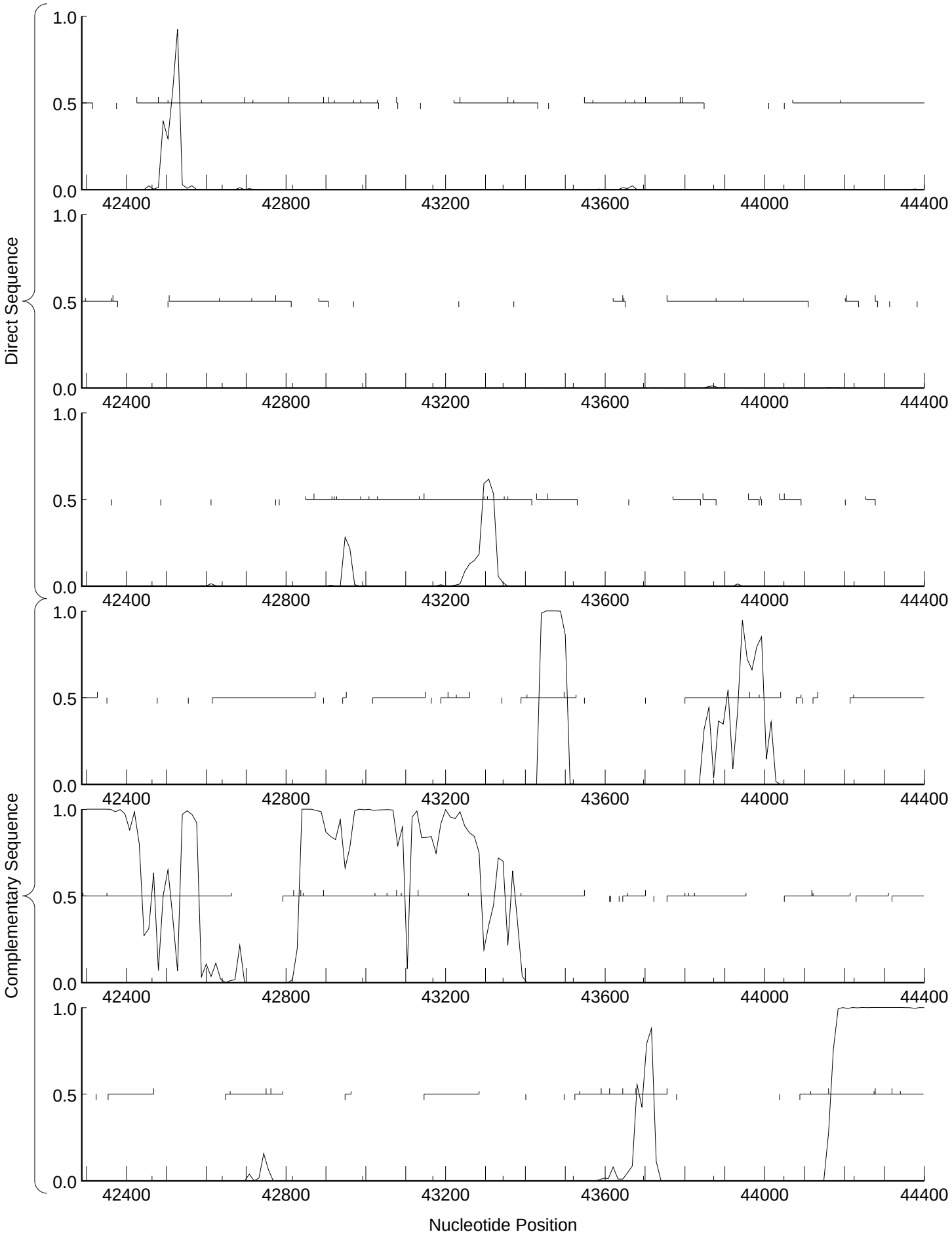

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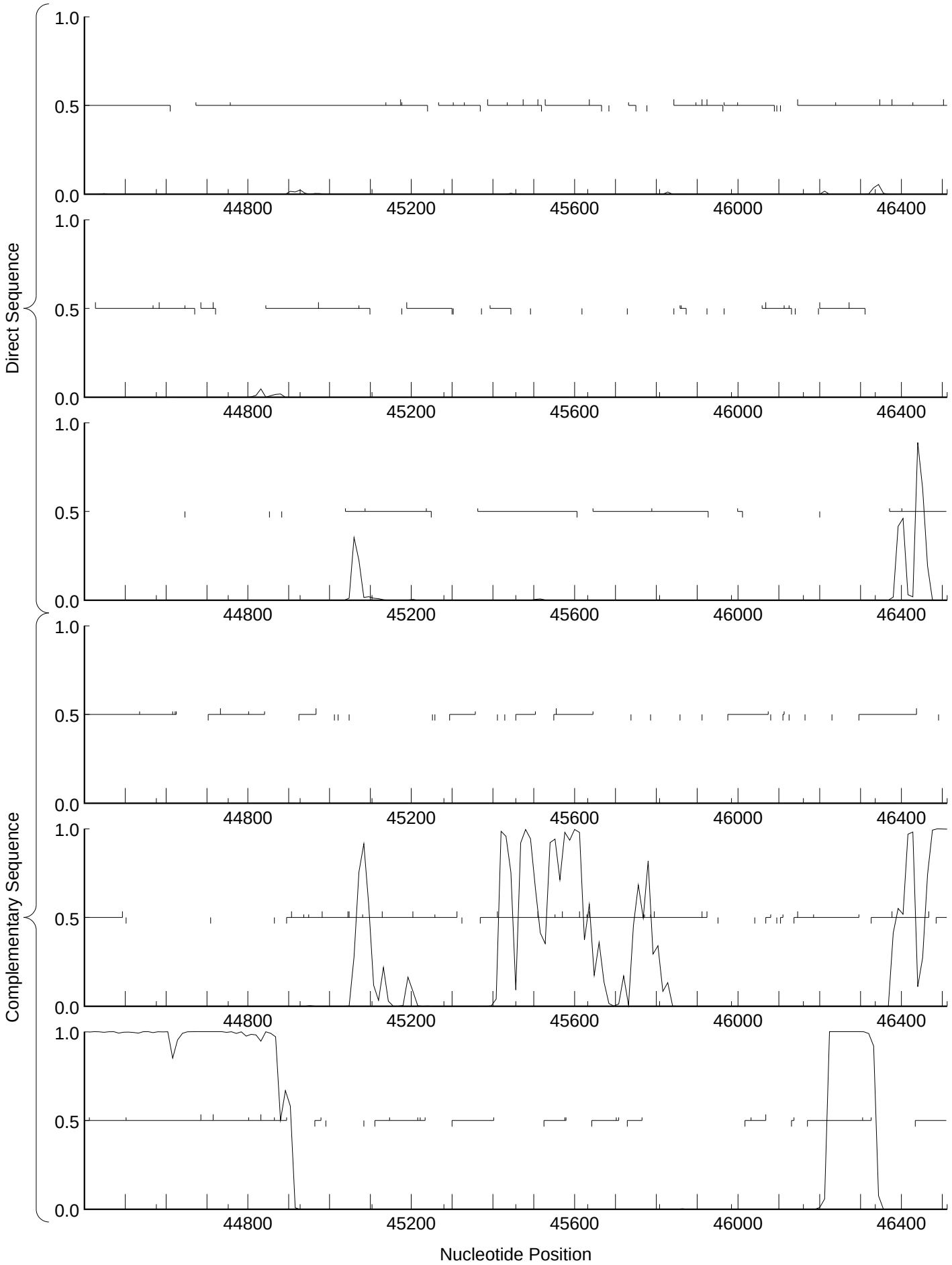

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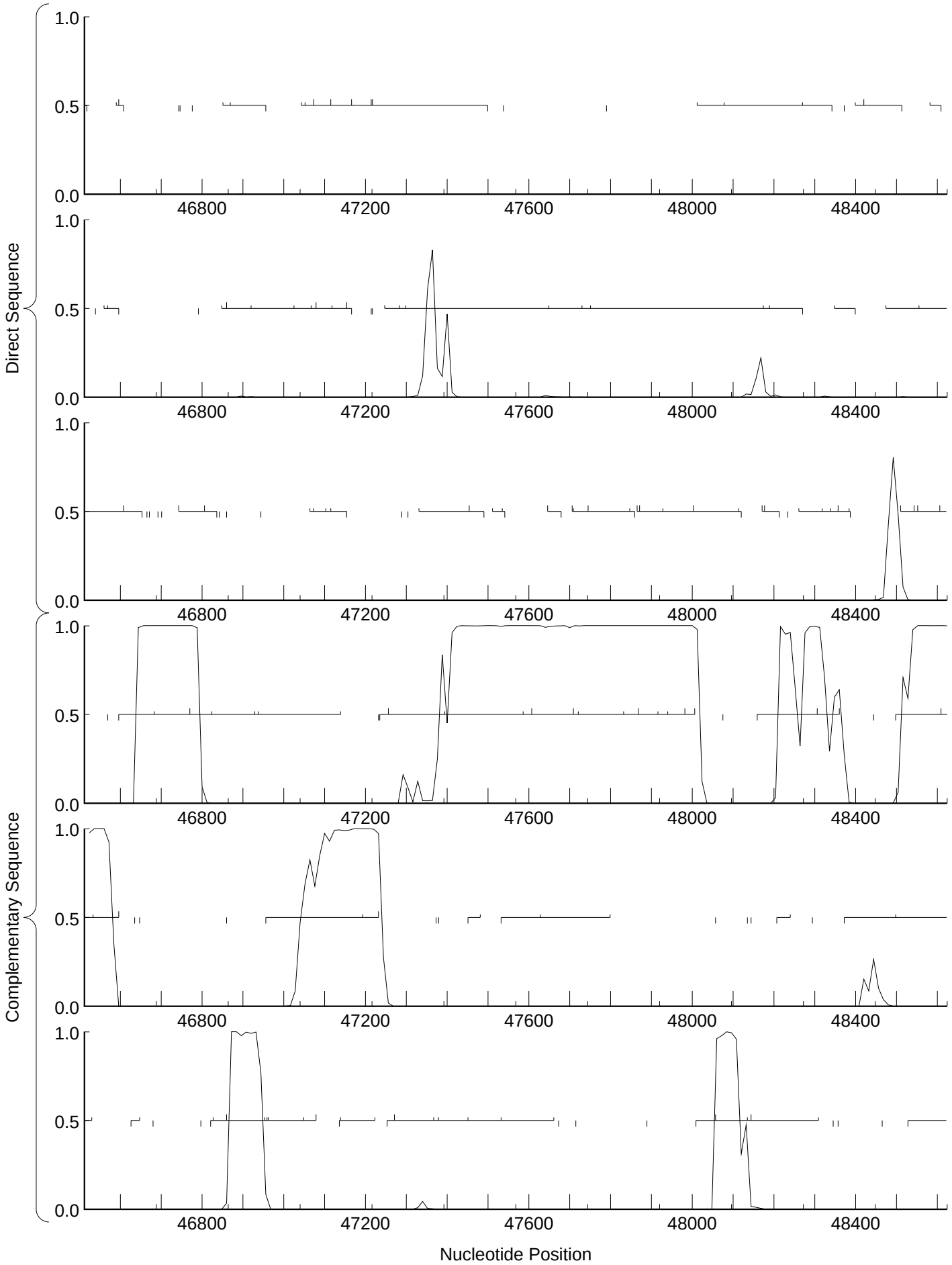

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Nucleotide Position

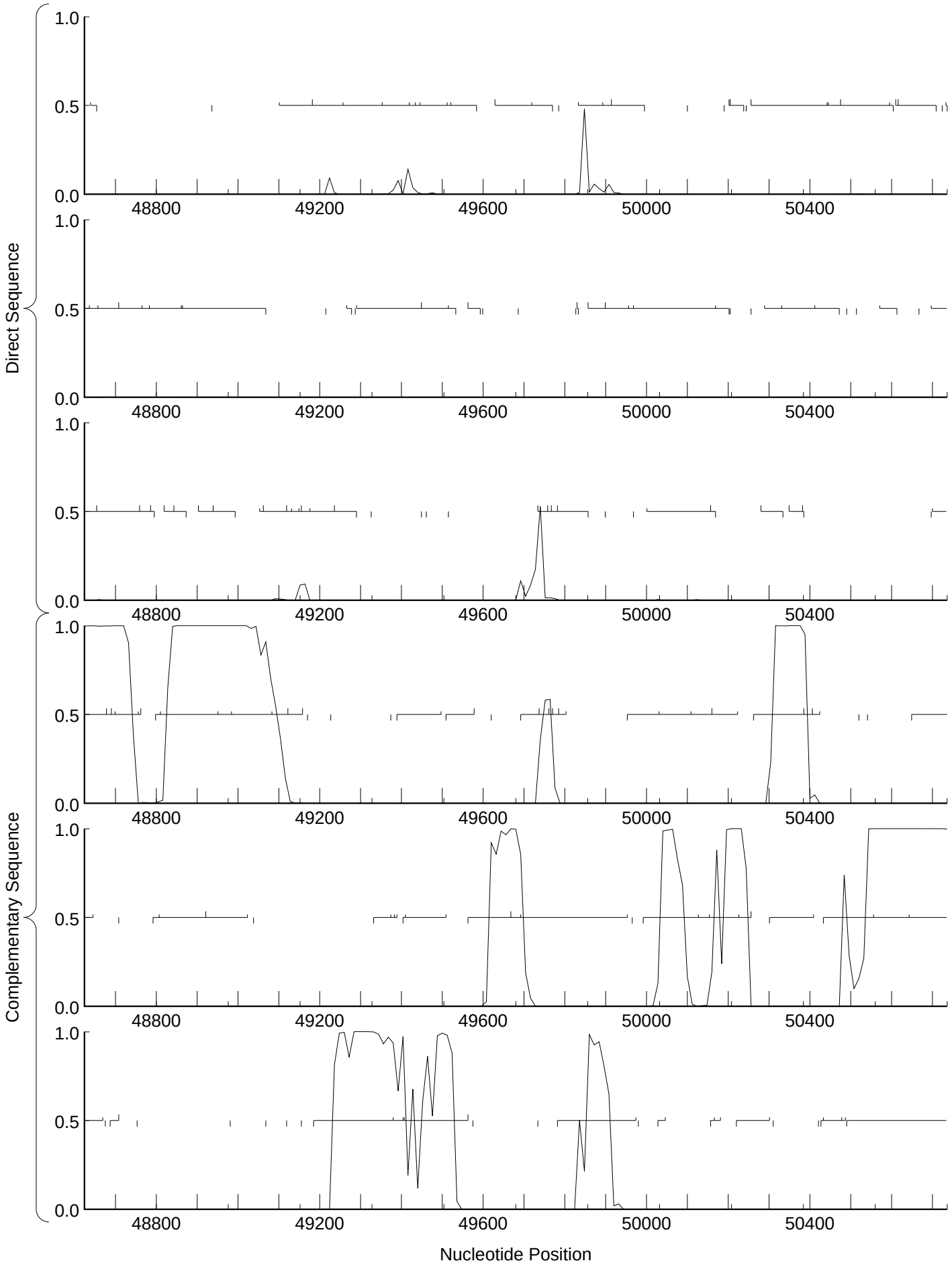

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Complementary Sequence
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39600
40000
Nucleotide Position

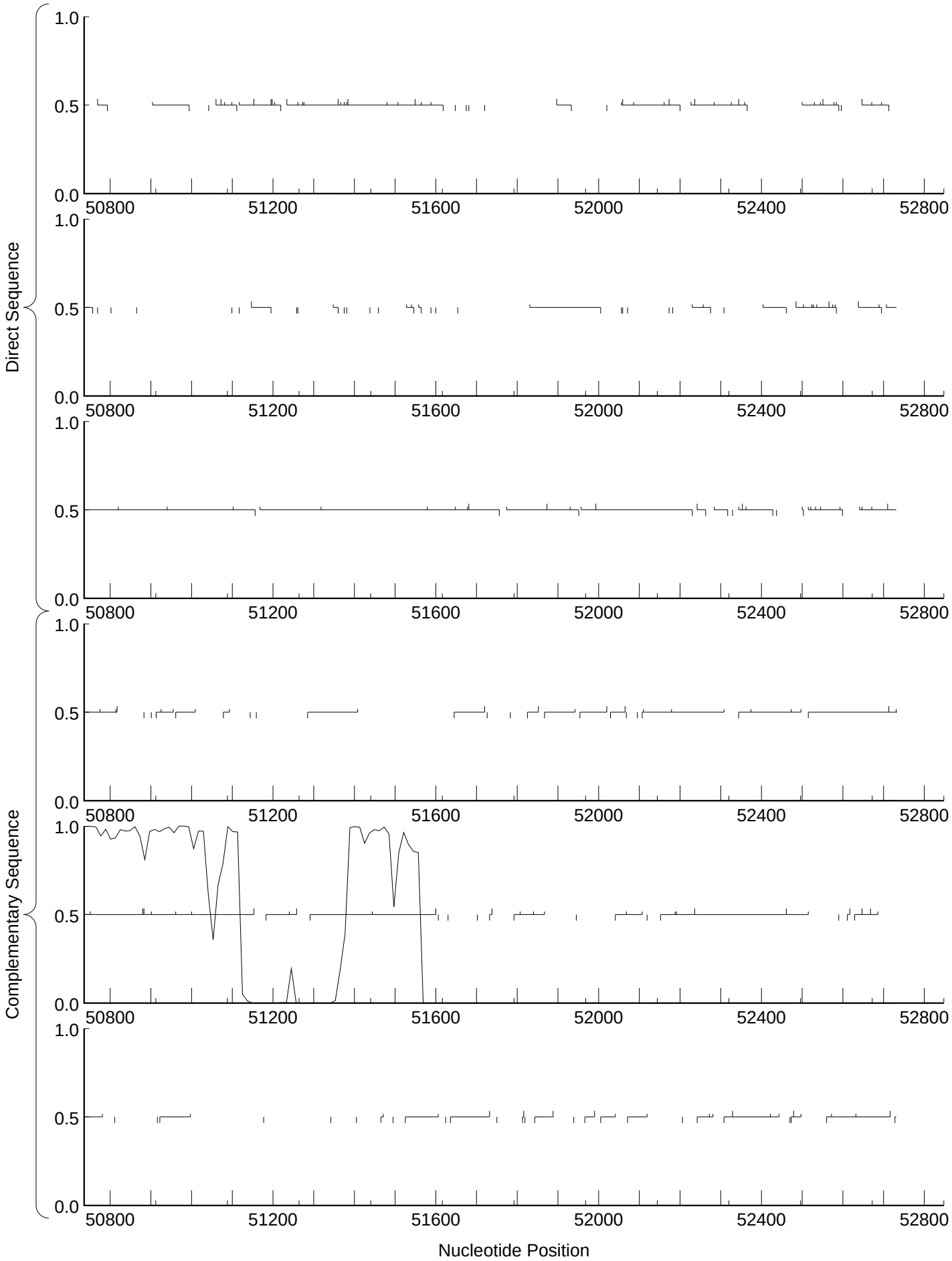

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0.0
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40800
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41600
42000
Nucleotide Position


Direct Sequence
Complementary Sequence
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0.5
0.0
42400
42800
43200
43600
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