GeneMark

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

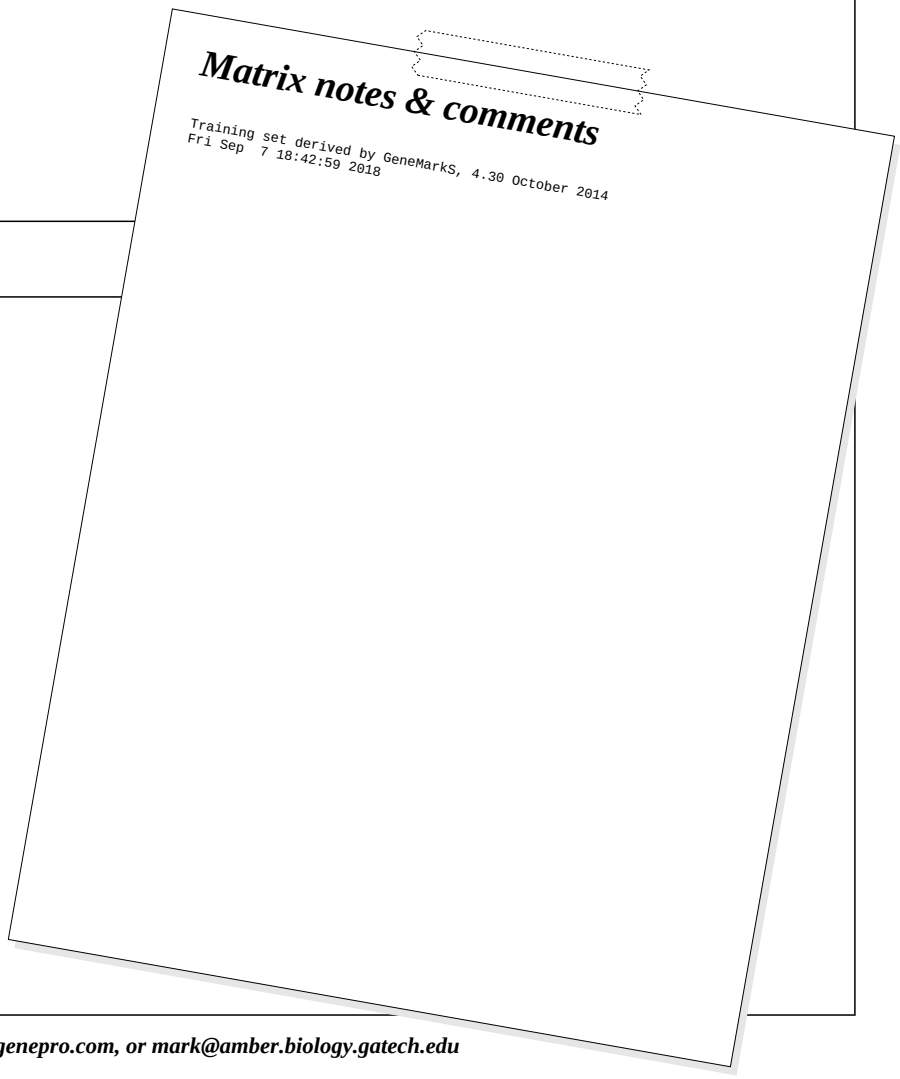
PROGRAM INFORMATION

```
Sequence         :  Microbacterium phage RobinRose complete sequence, 63011 bp including 203 bp terminal repeat
Analysis Date    :  3/29/24 at 11:08:32
Pages            :  31
Sequence Length  :  63011 bp
GC Content       :   66.99%

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

PS-Version       :  1.2

GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
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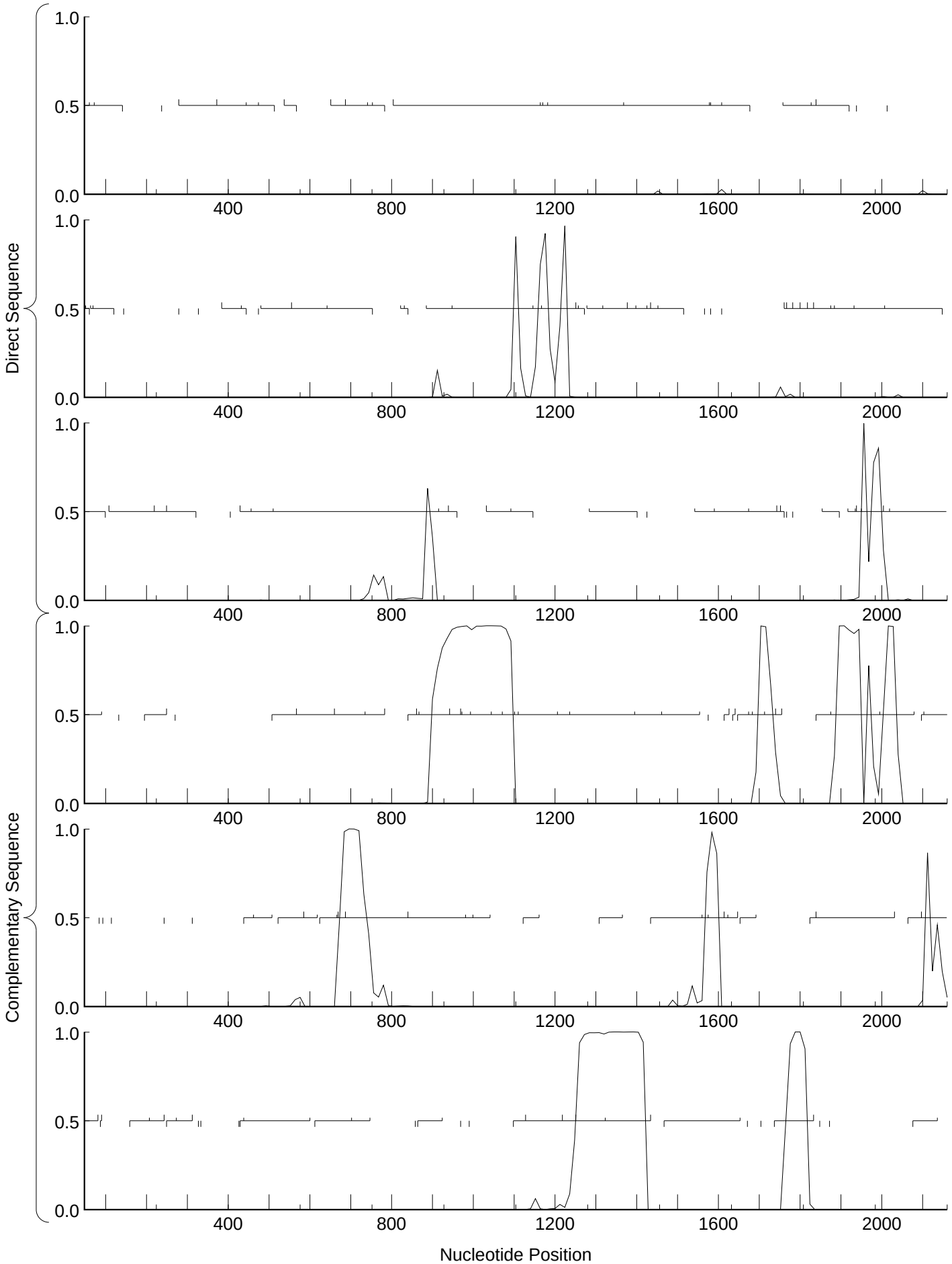
MATRIX INFORMATION

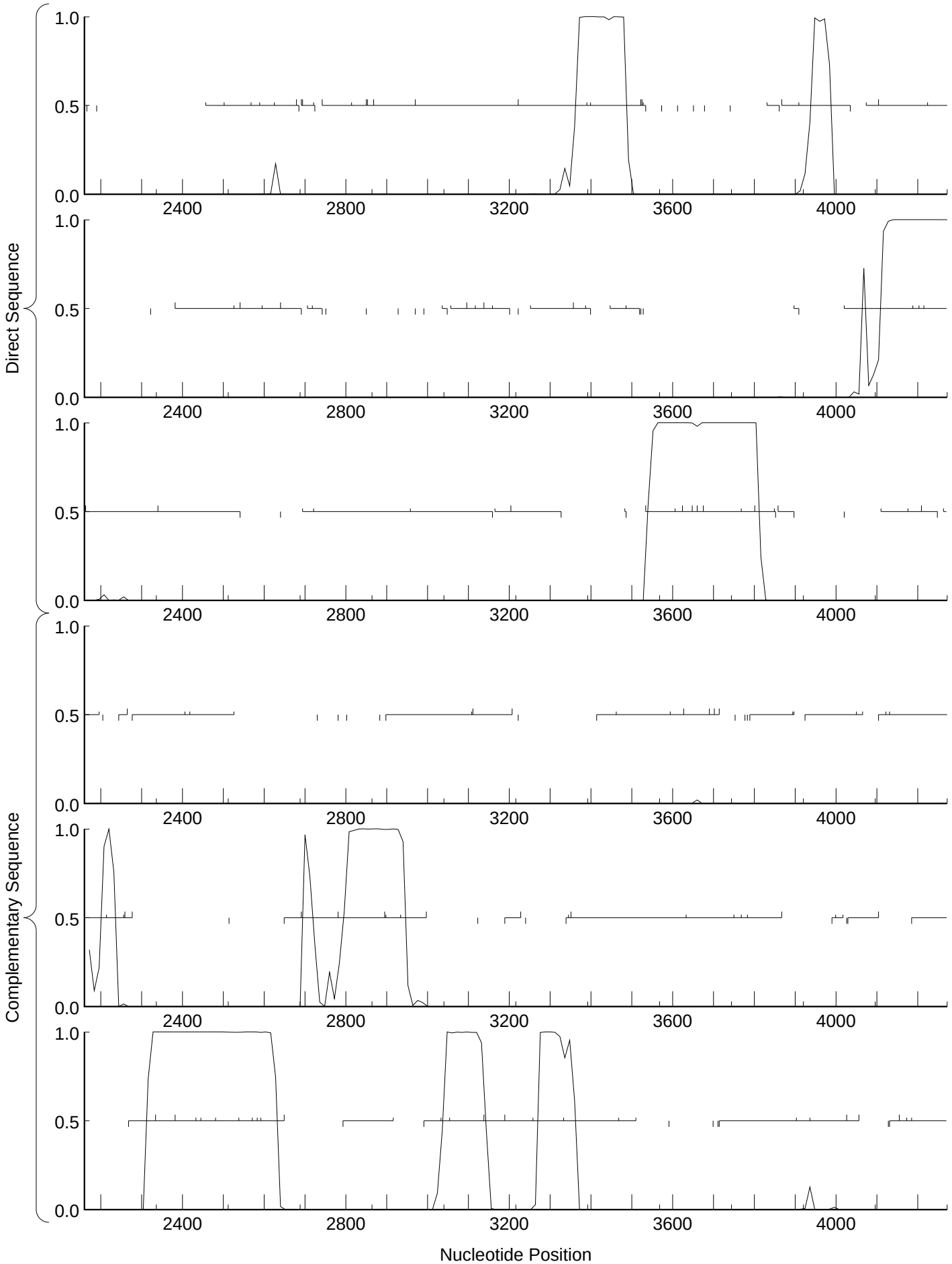
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Matrix     :  -
Author     :  -

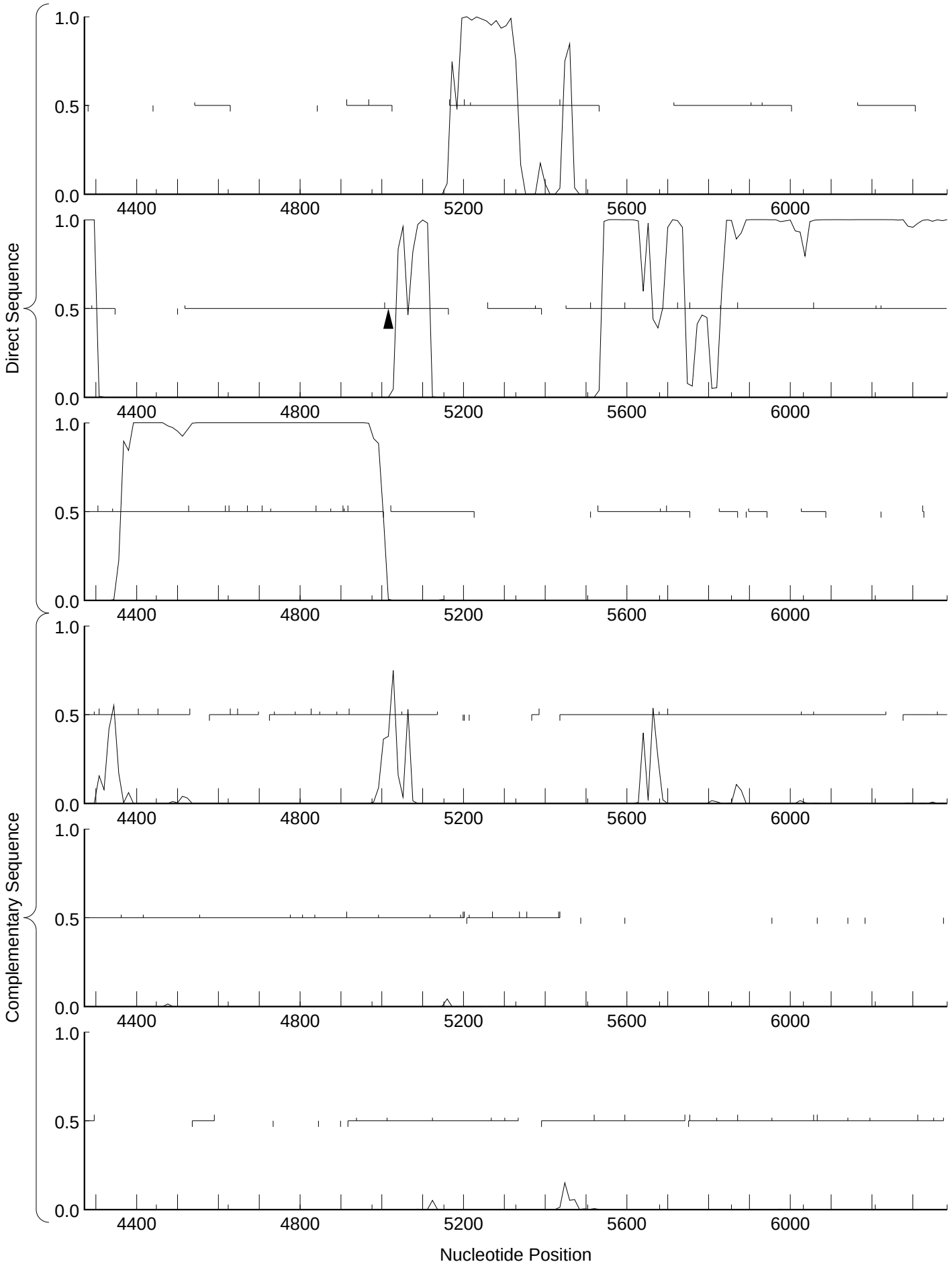
Order      :  4

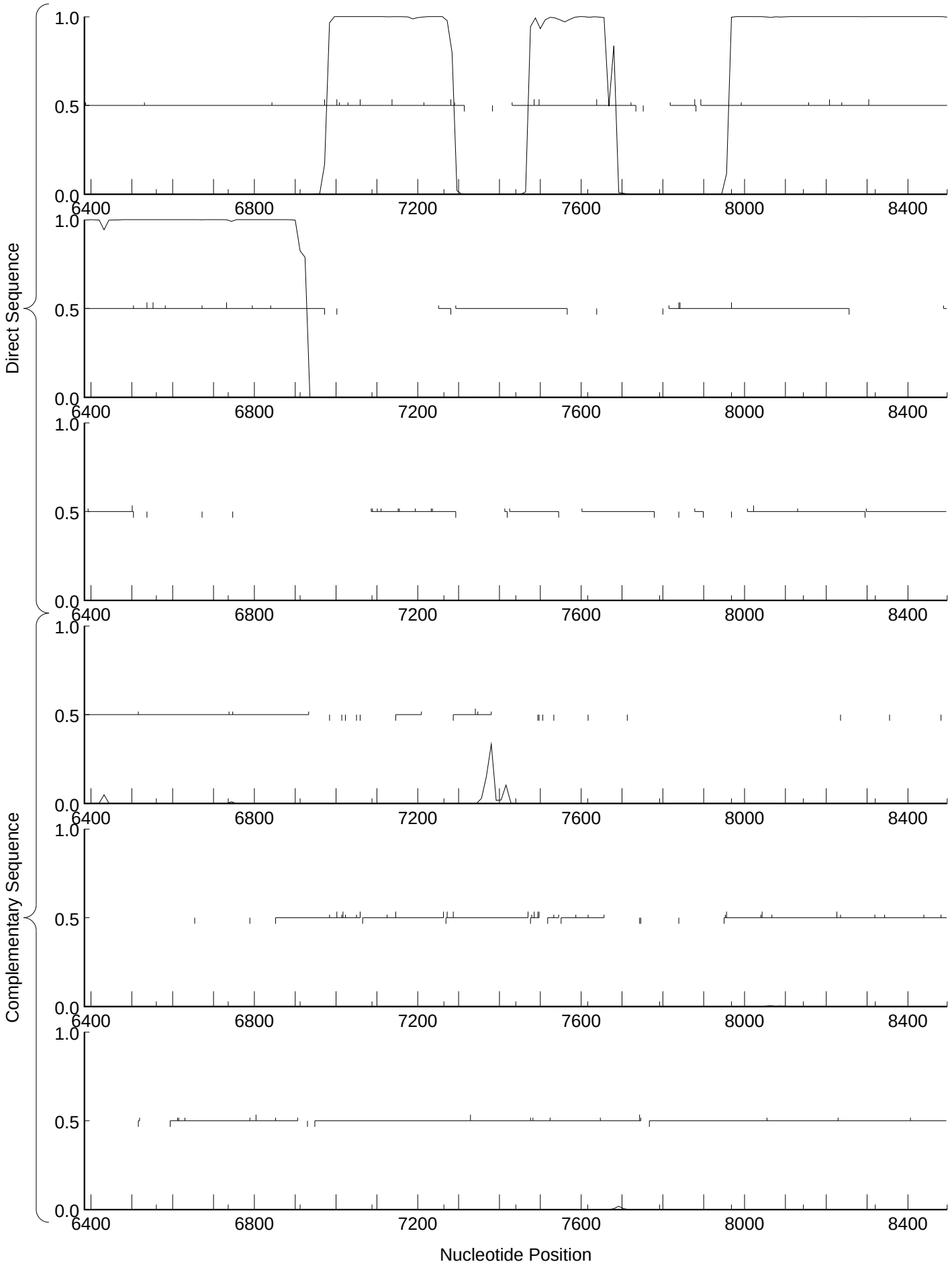

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

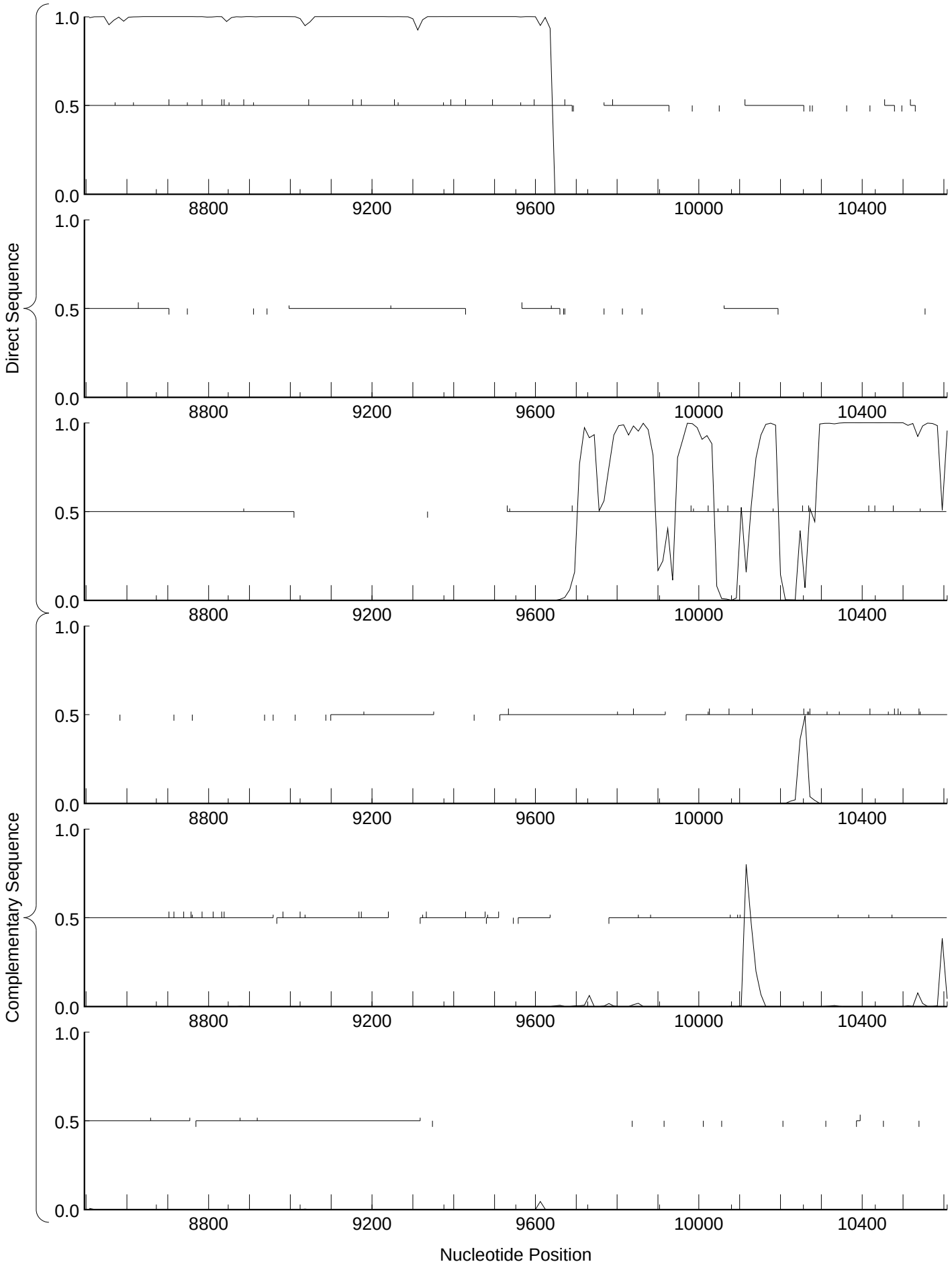
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

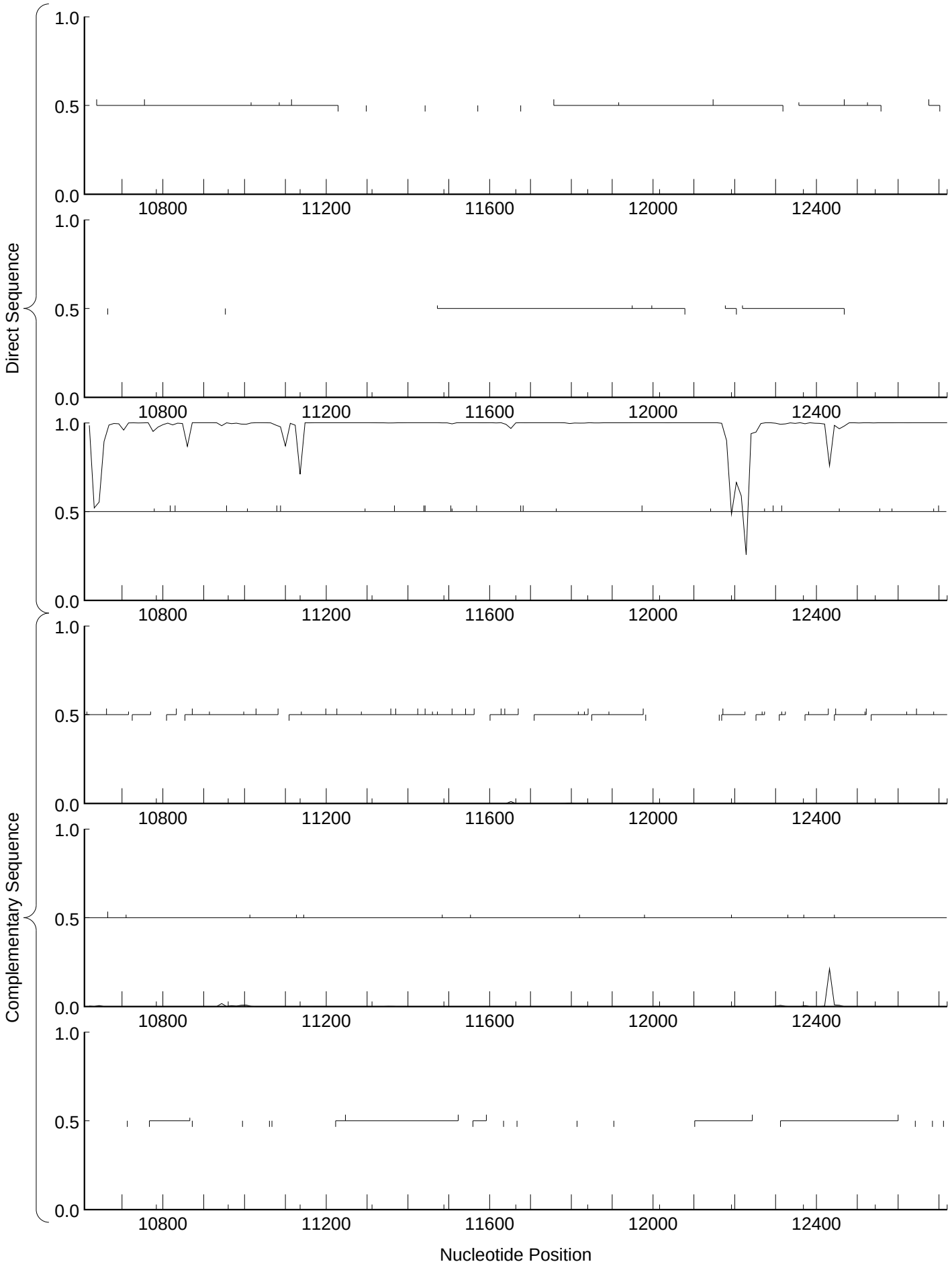

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
400
800
1200
1600
2000
Nucleotide Position

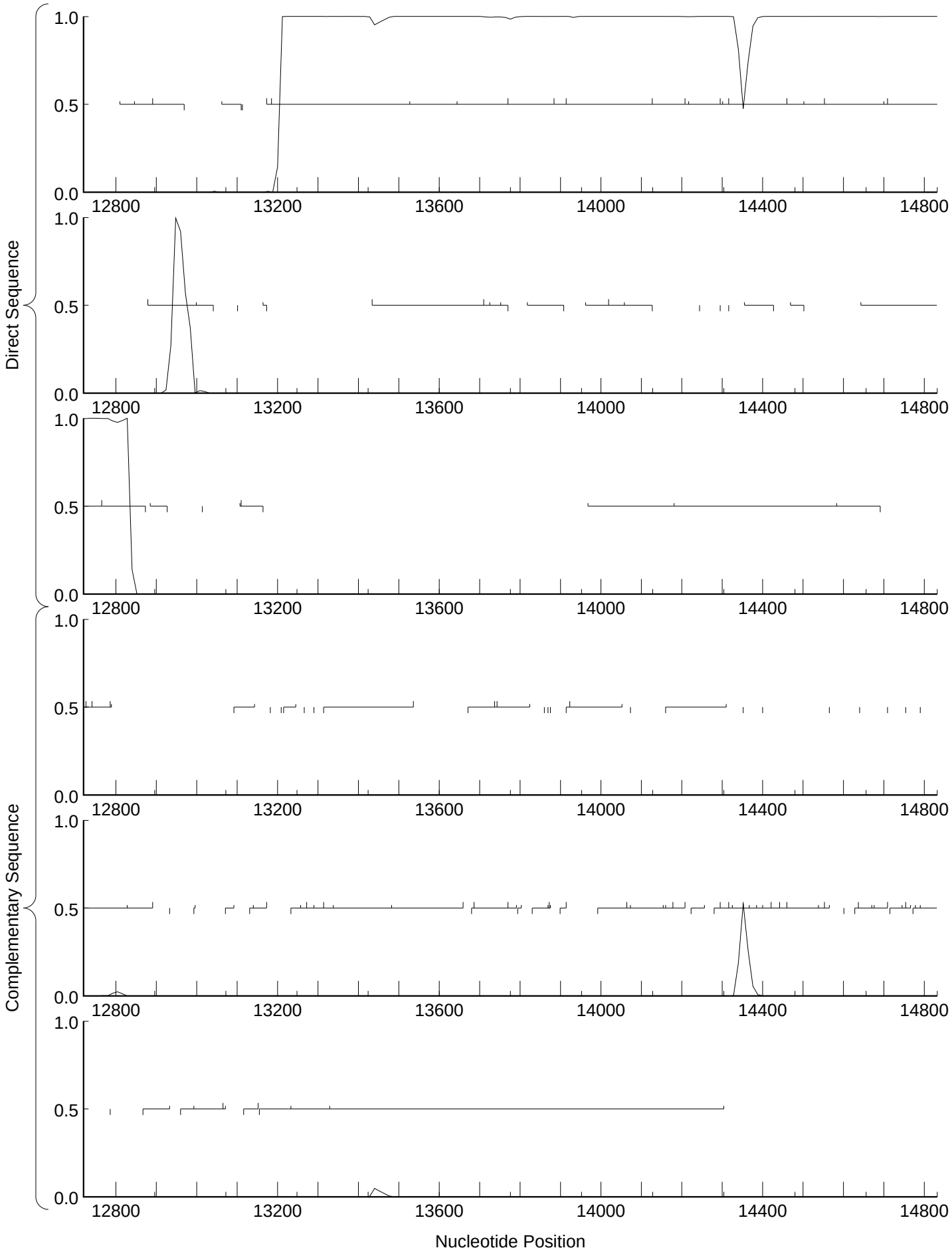

Direct Sequence
Complementary Sequence
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2800
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3600
4000
Nucleotide Position

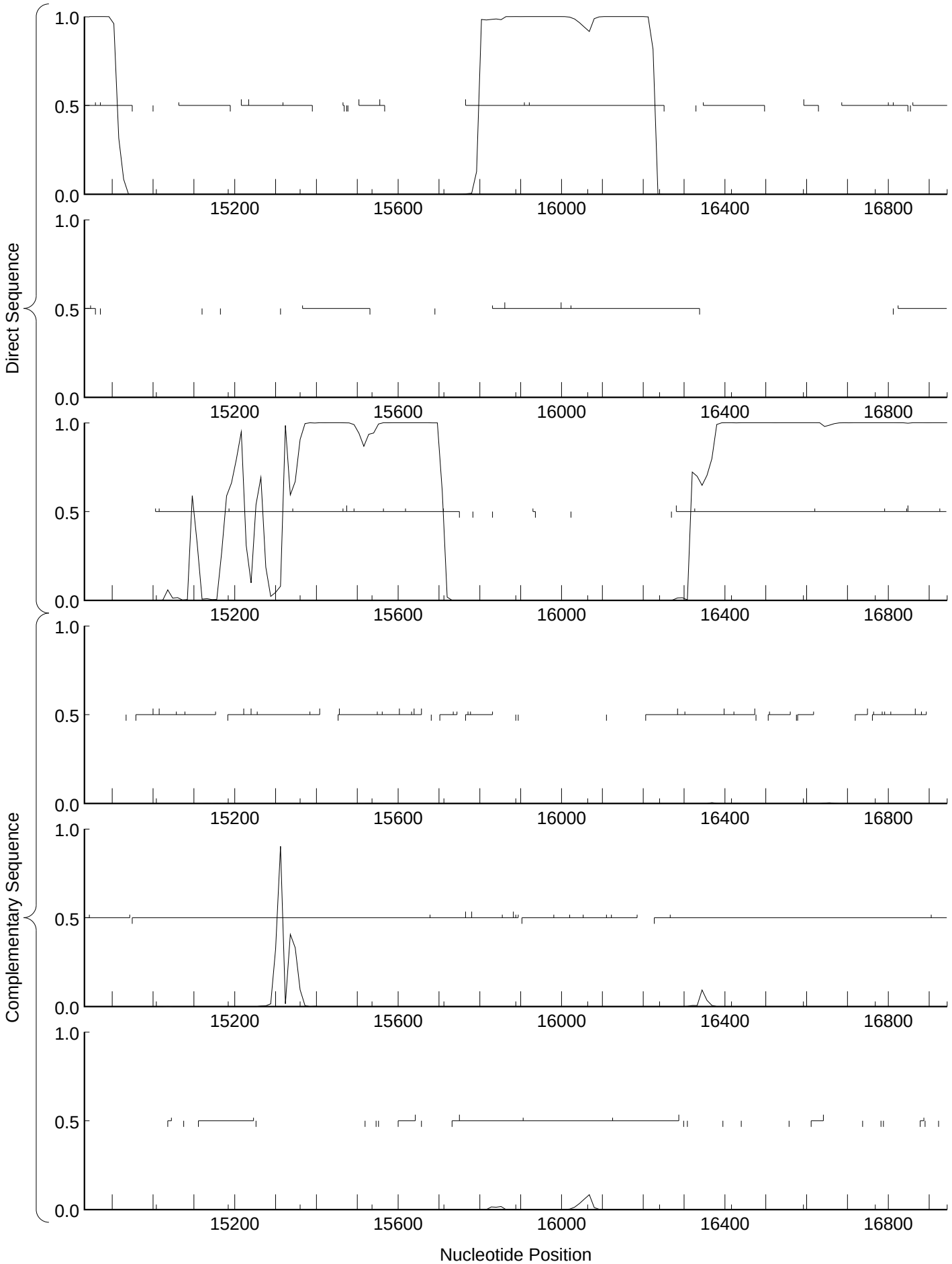

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Complementary Sequence
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5600
6000
Nucleotide Position

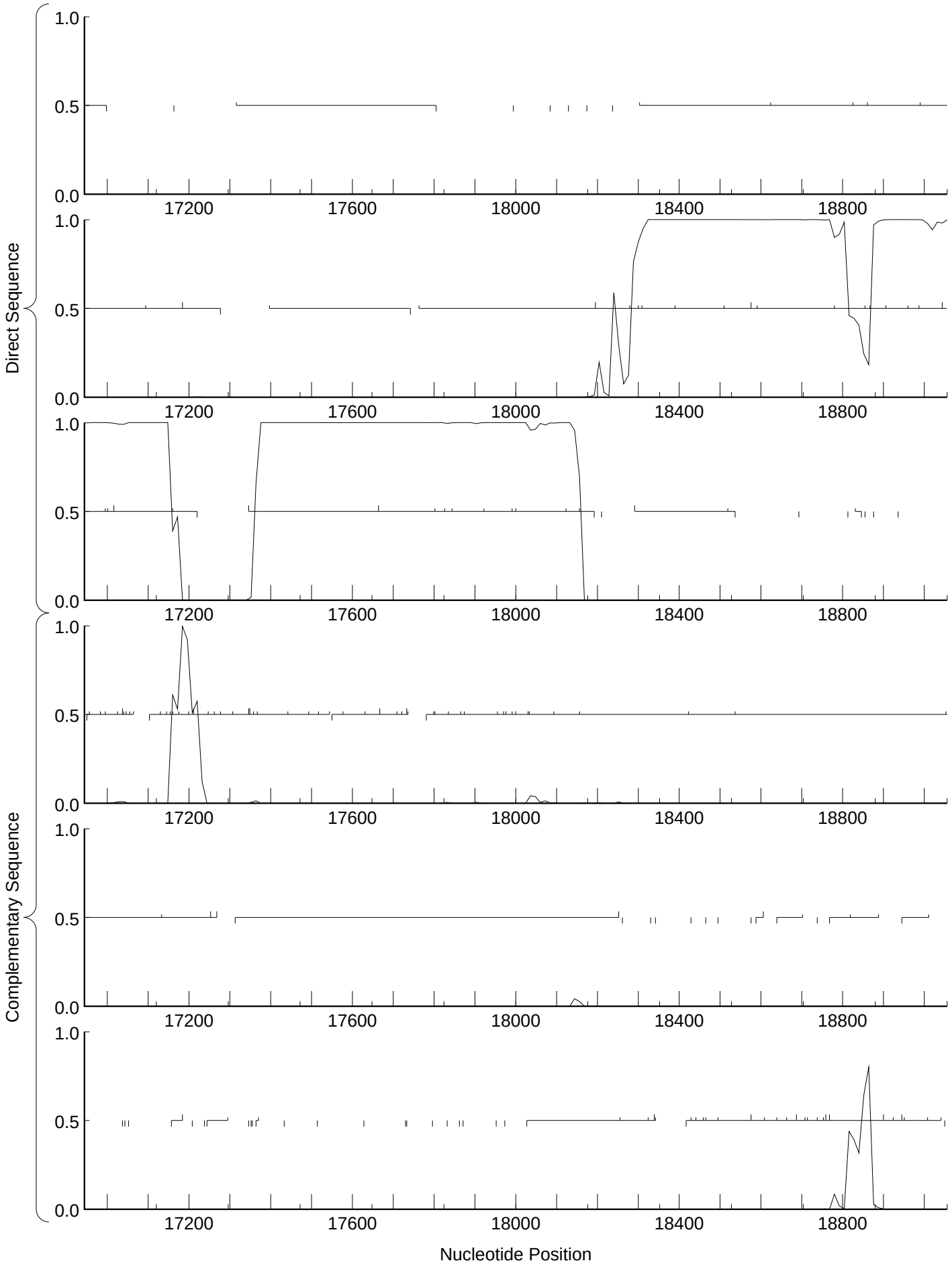

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8000
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Nucleotide Position

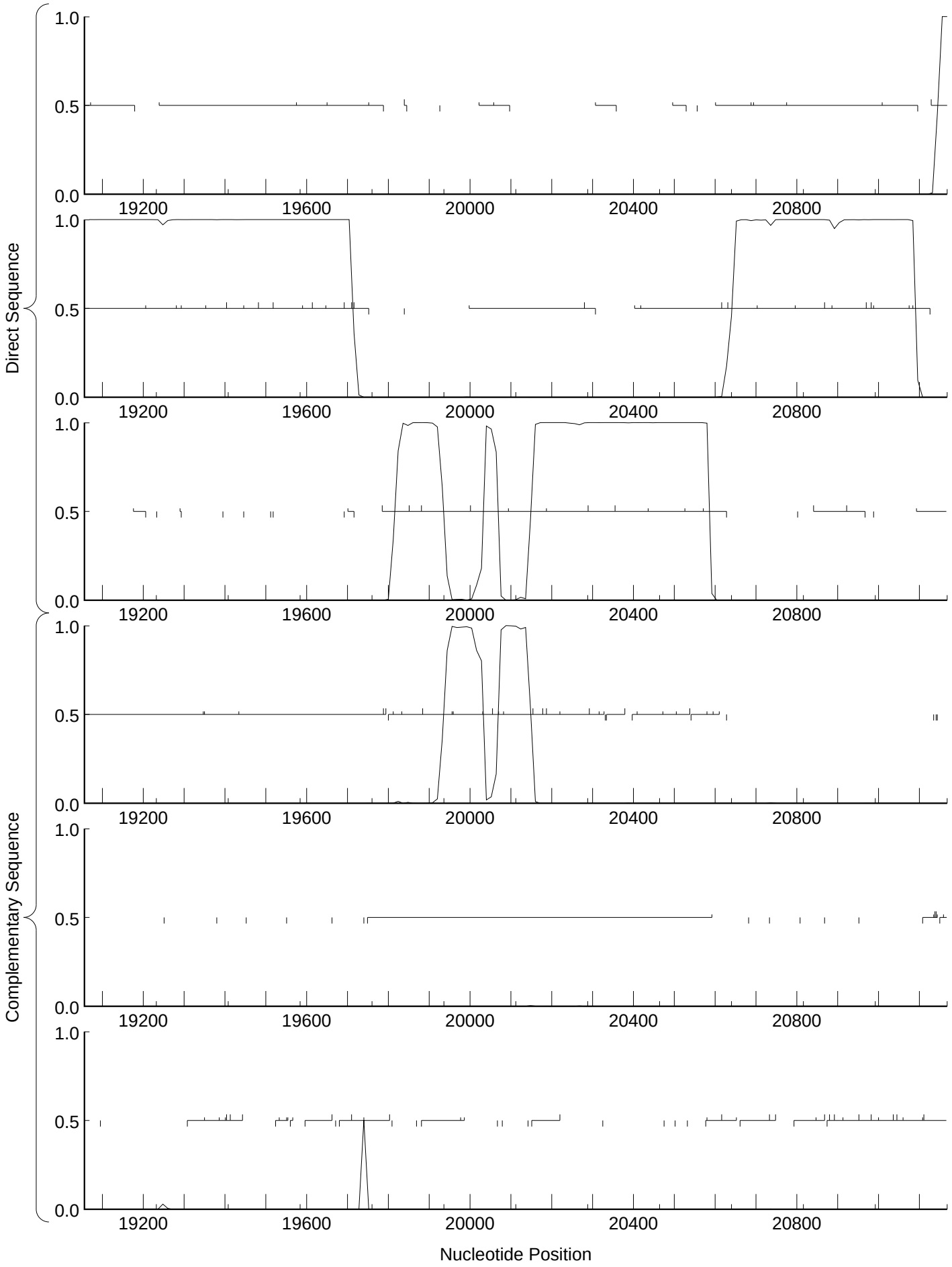

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Nucleotide Position

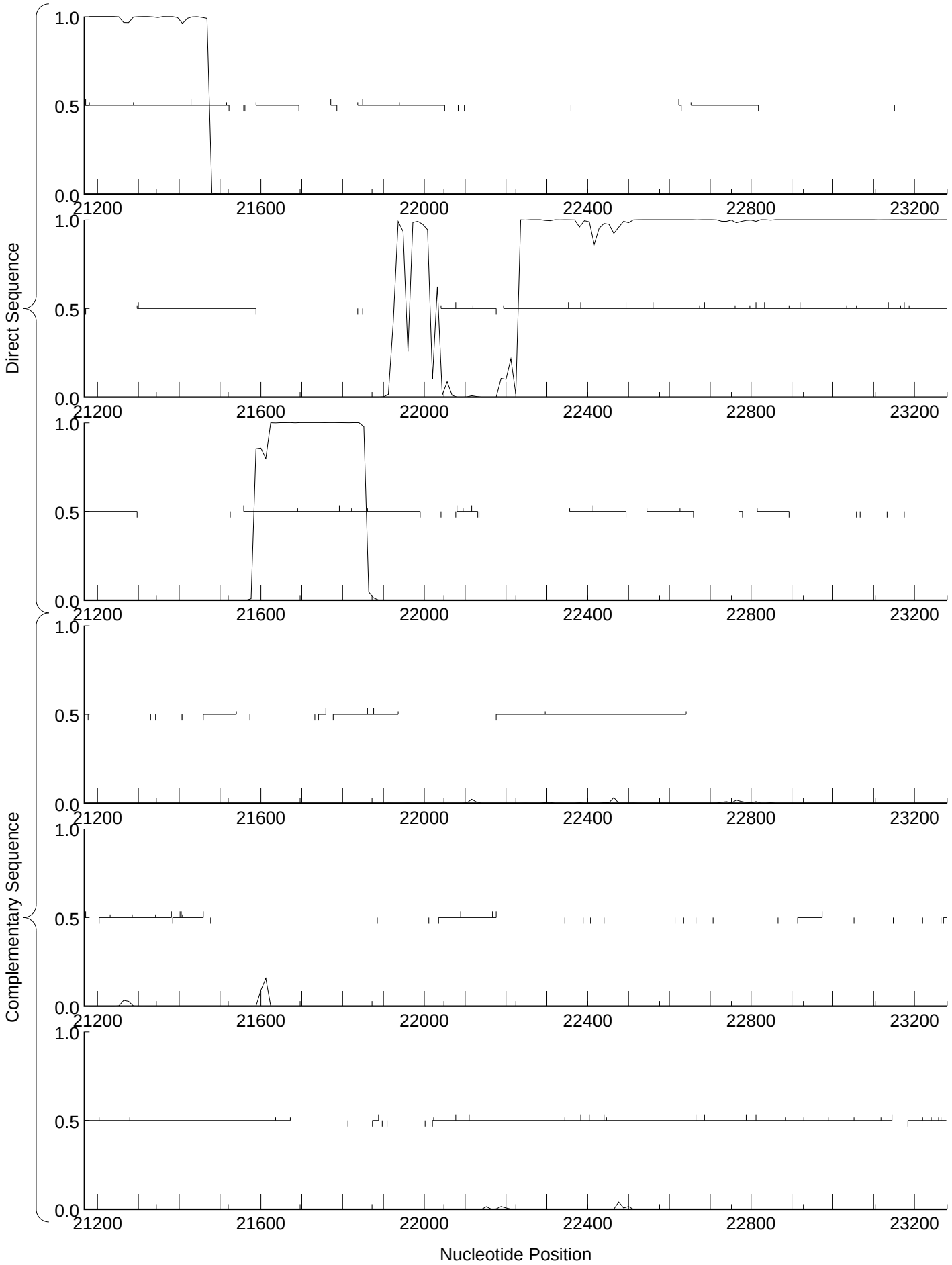

Direct Sequence
Complementary Sequence
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11200
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12000
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Nucleotide Position

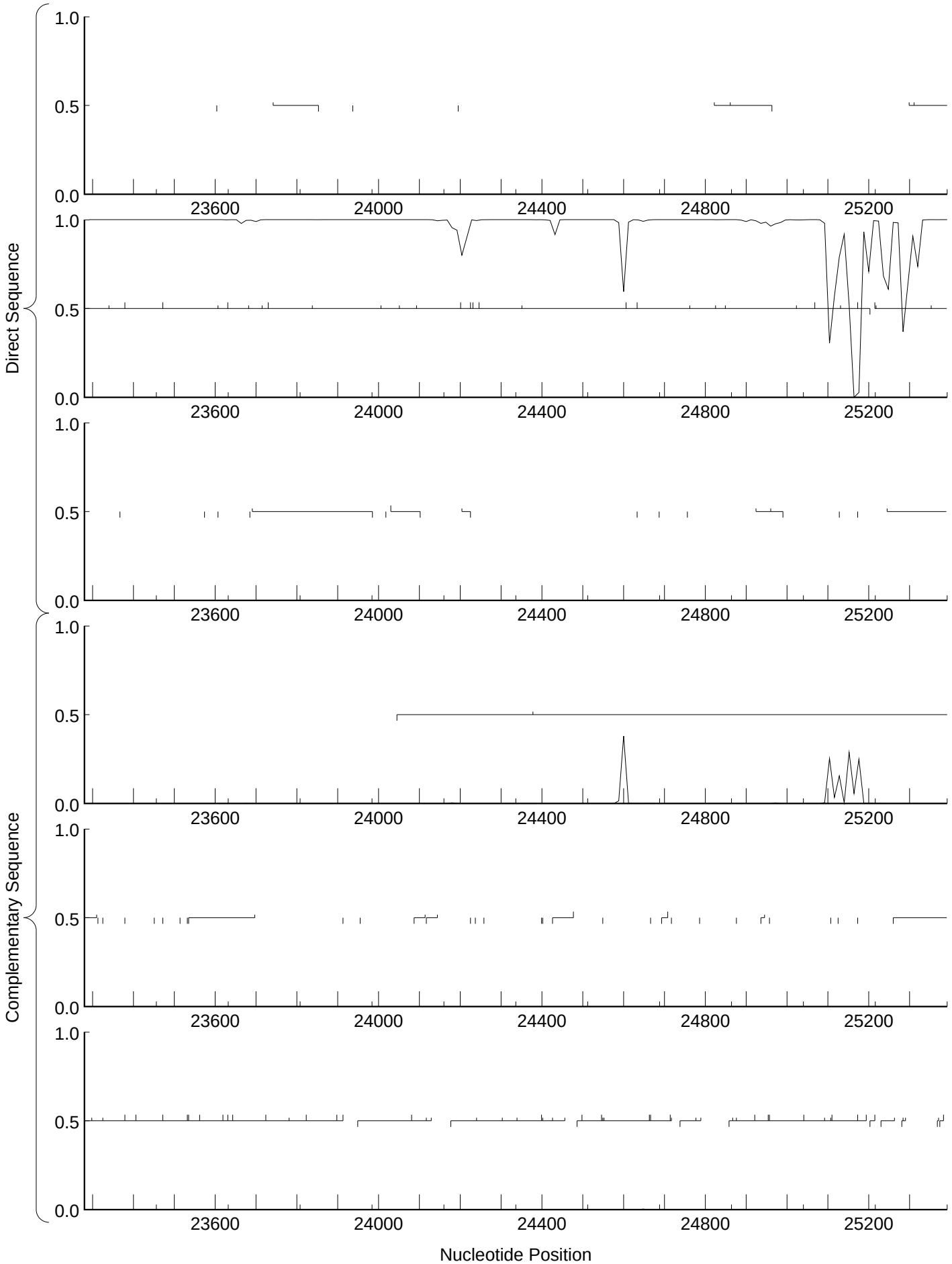

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Complementary Sequence
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Nucleotide Position

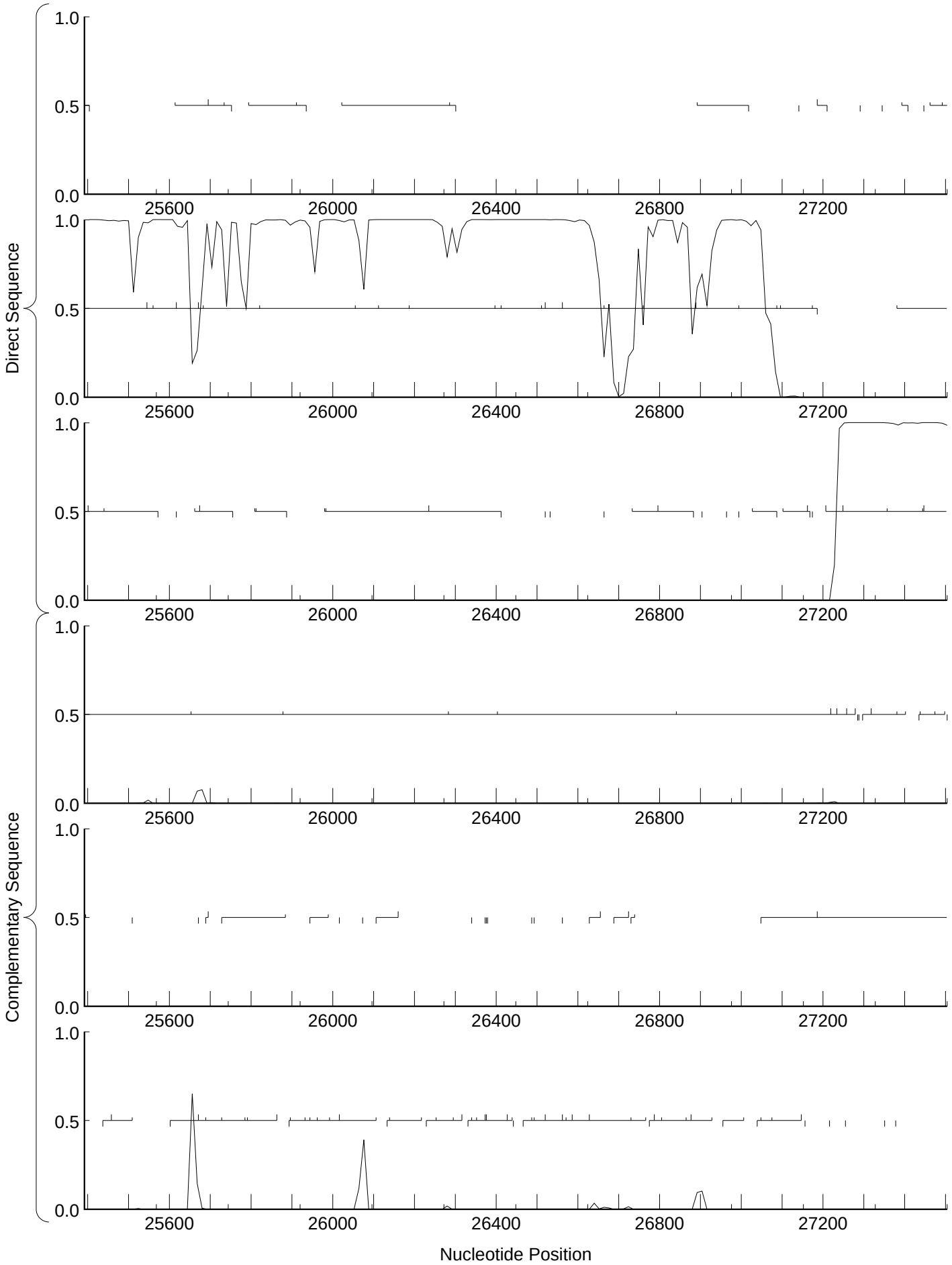

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Complementary Sequence
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Nucleotide Position

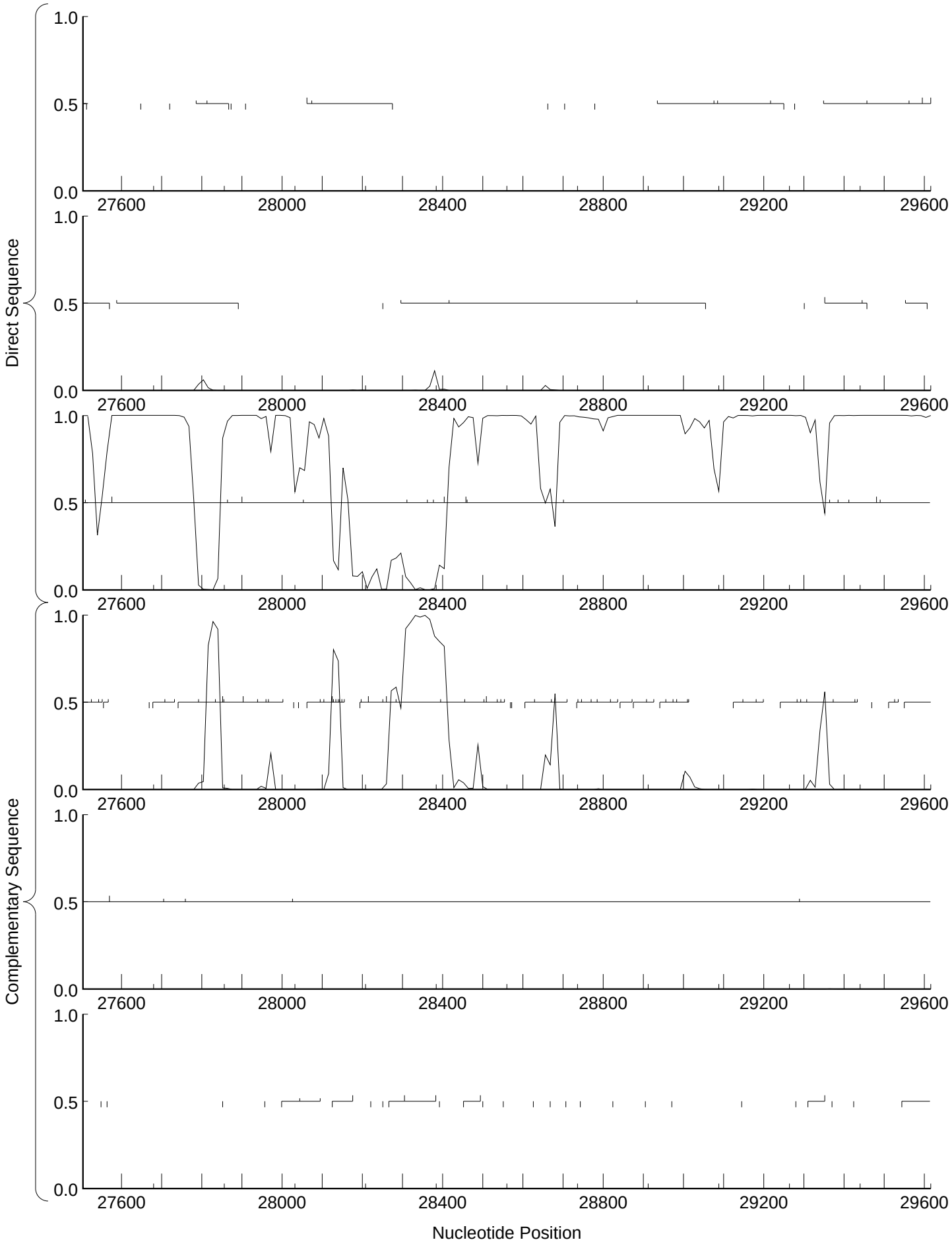

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Complementary Sequence
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18800
Nucleotide Position

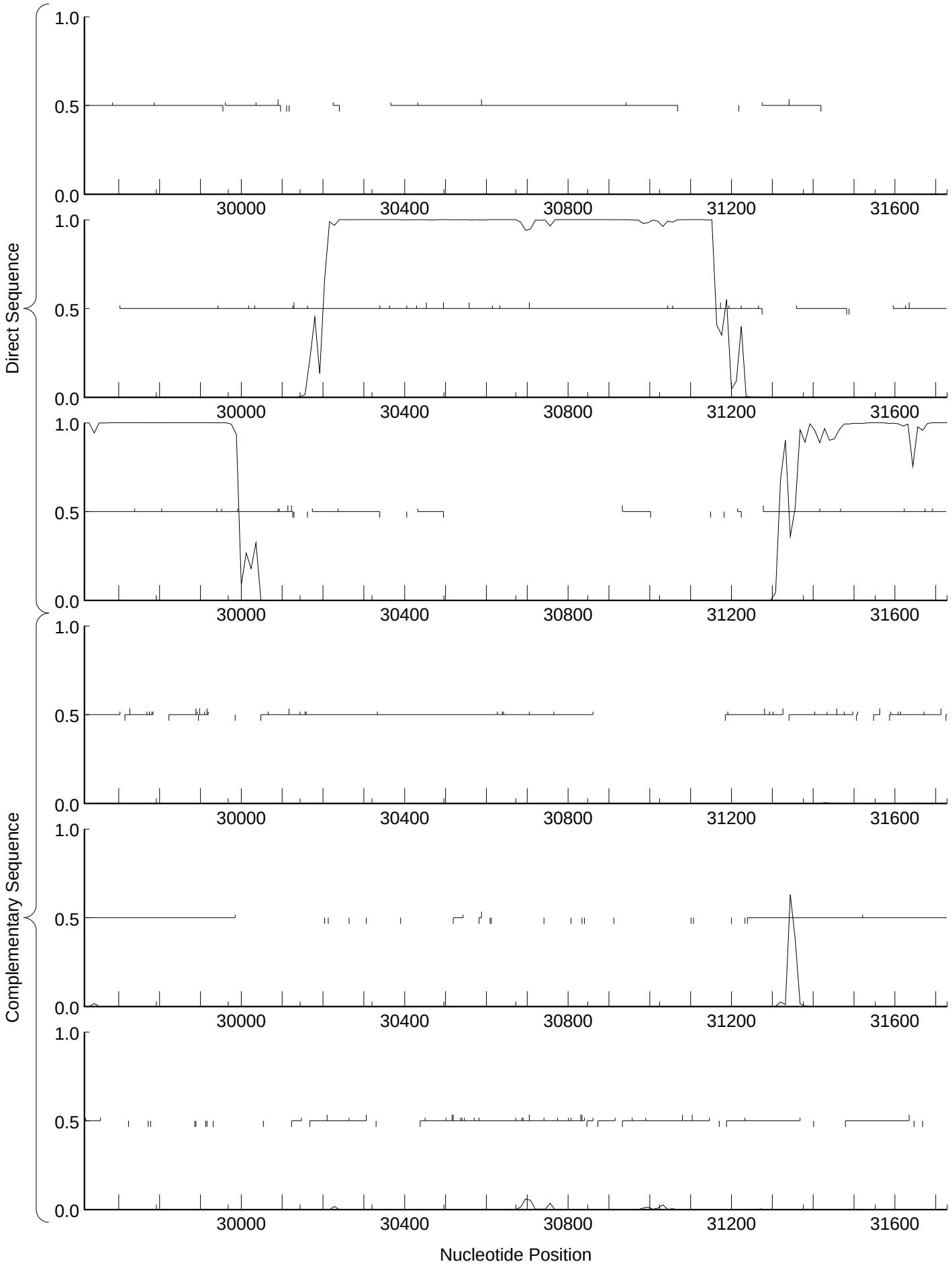

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Complementary Sequence
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Nucleotide Position

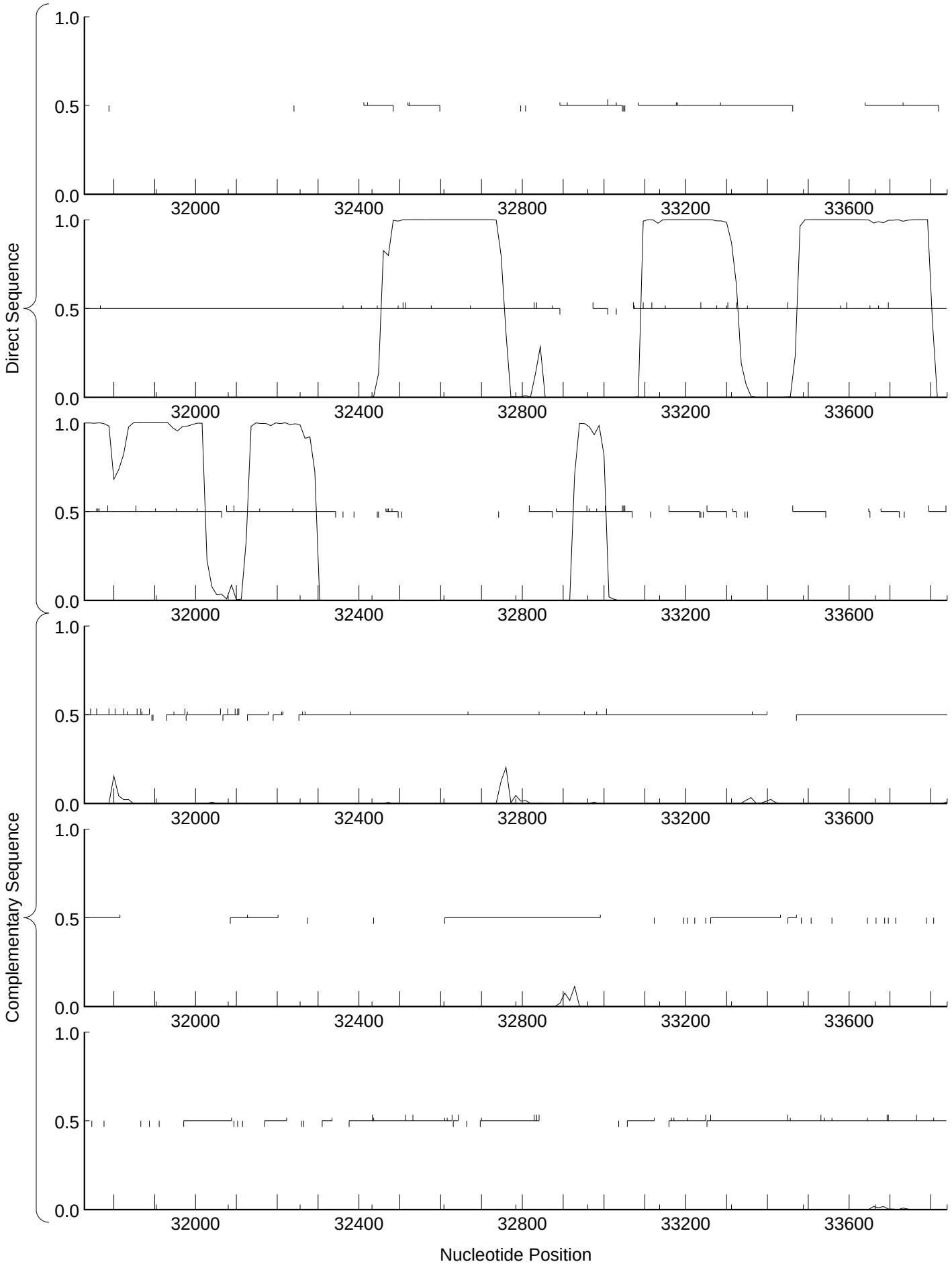

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Complementary Sequence
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22800
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Nucleotide Position

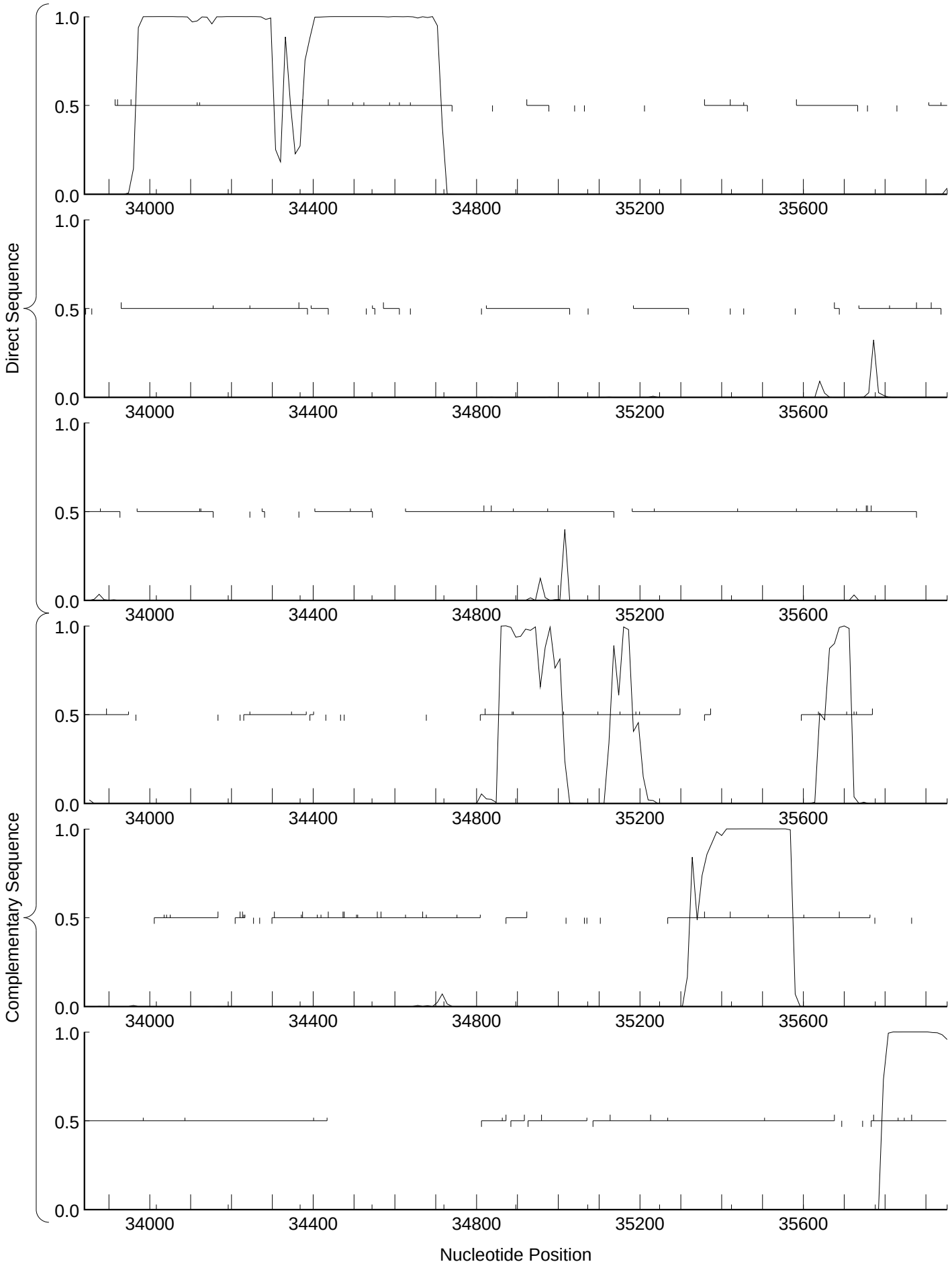

Direct Sequence
Complementary Sequence
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Nucleotide Position

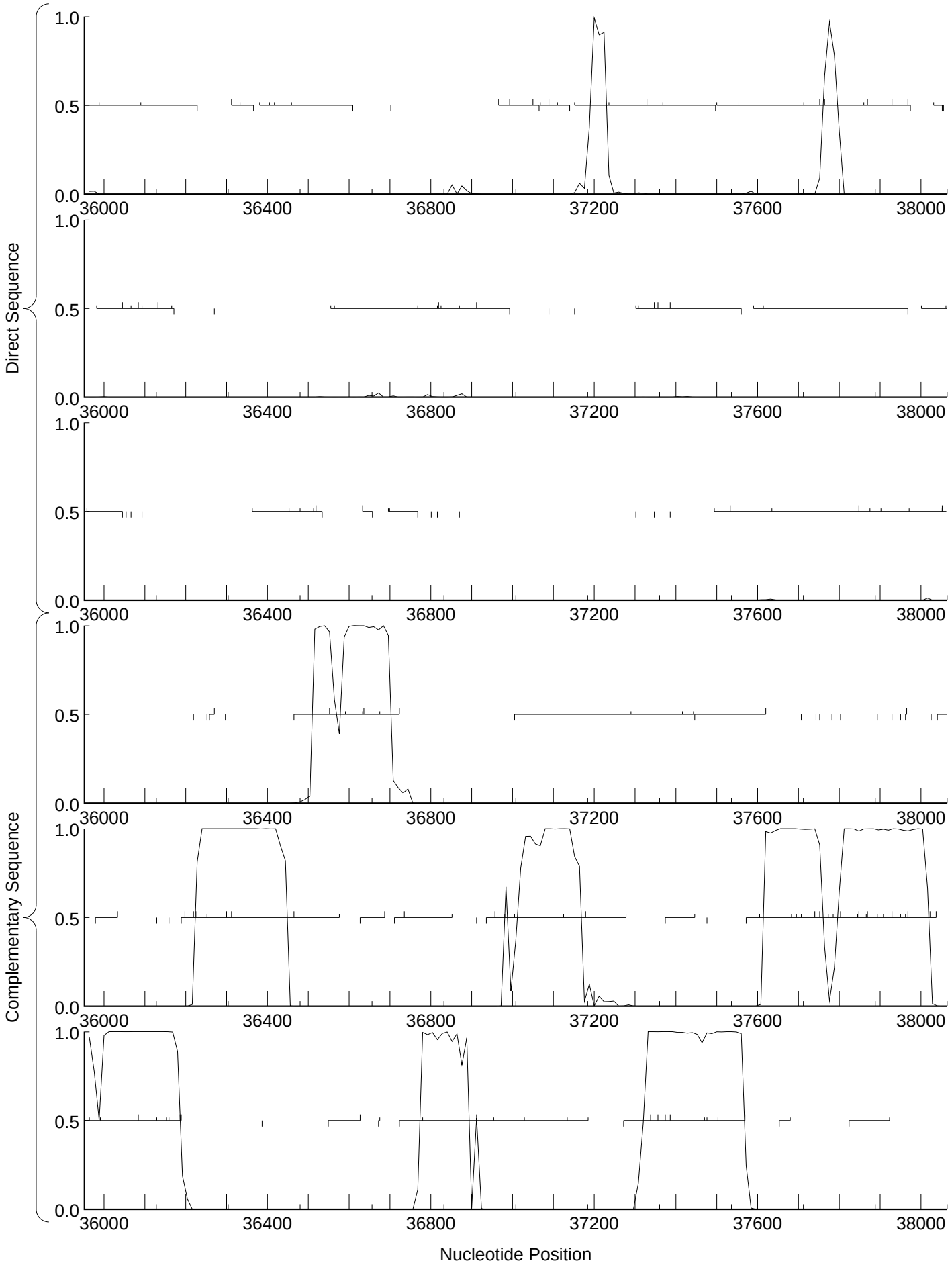

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Complementary Sequence
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Nucleotide Position

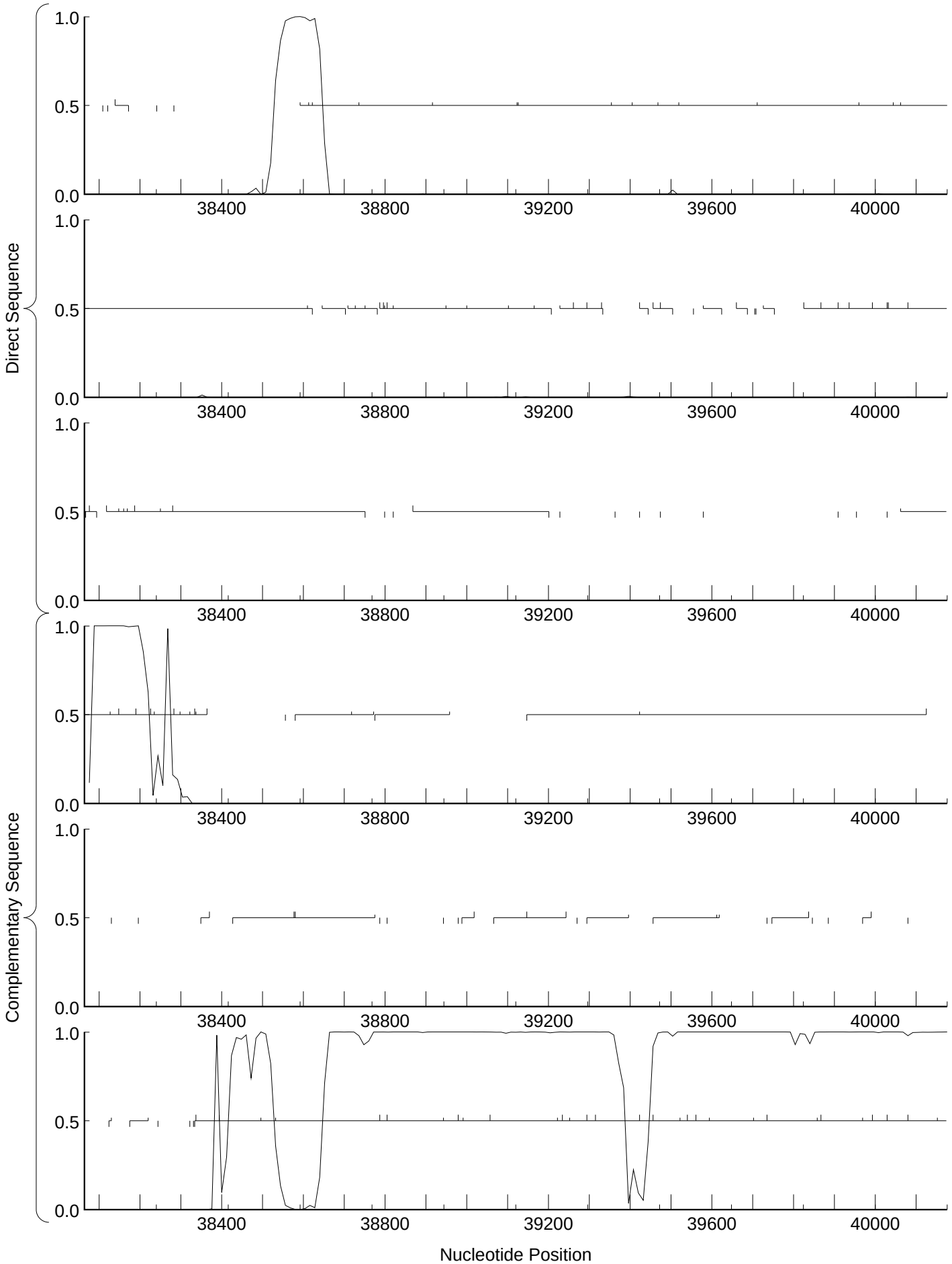

Direct Sequence
Complementary Sequence
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29200
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Nucleotide Position

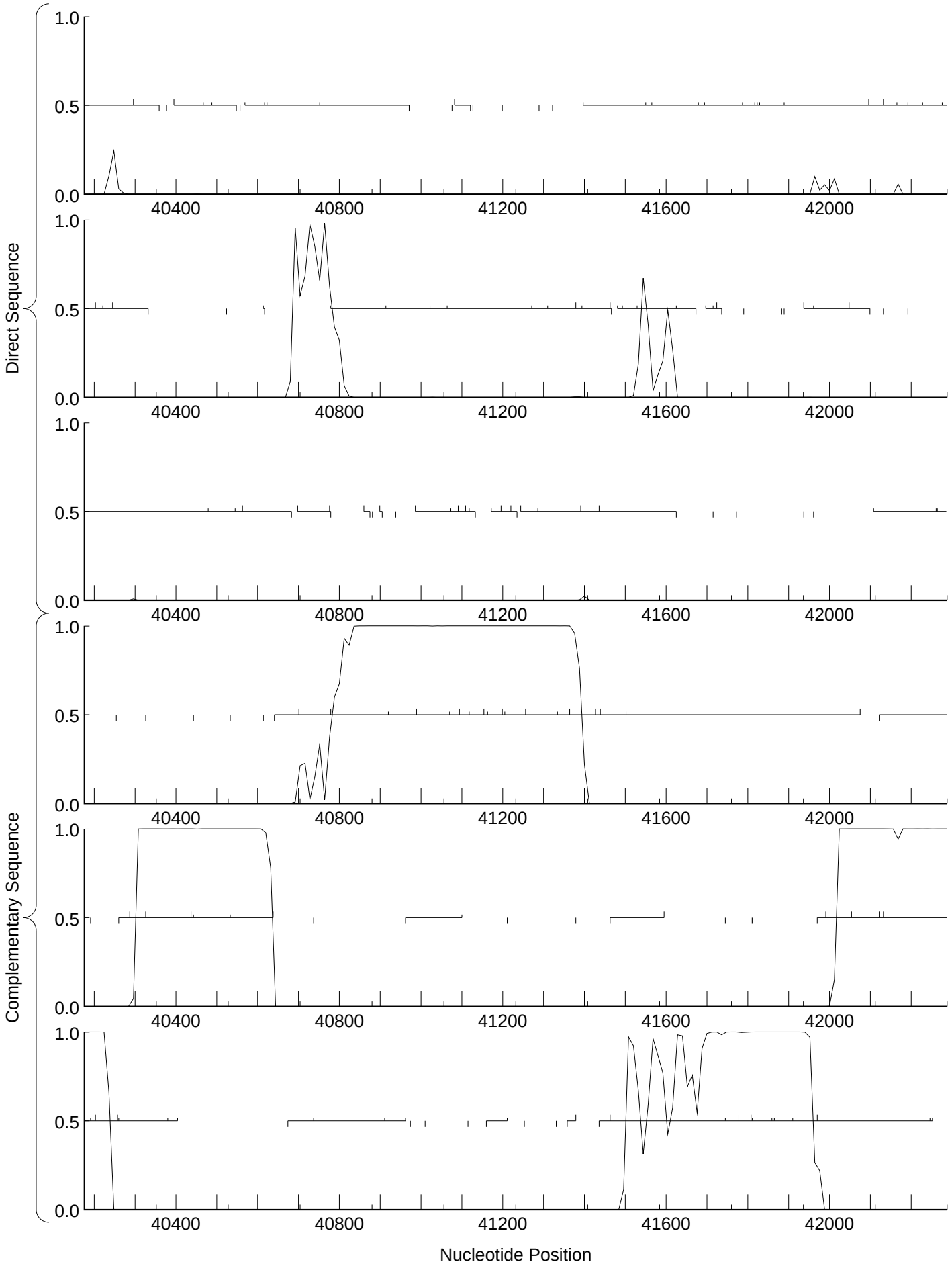

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Complementary Sequence
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Nucleotide Position

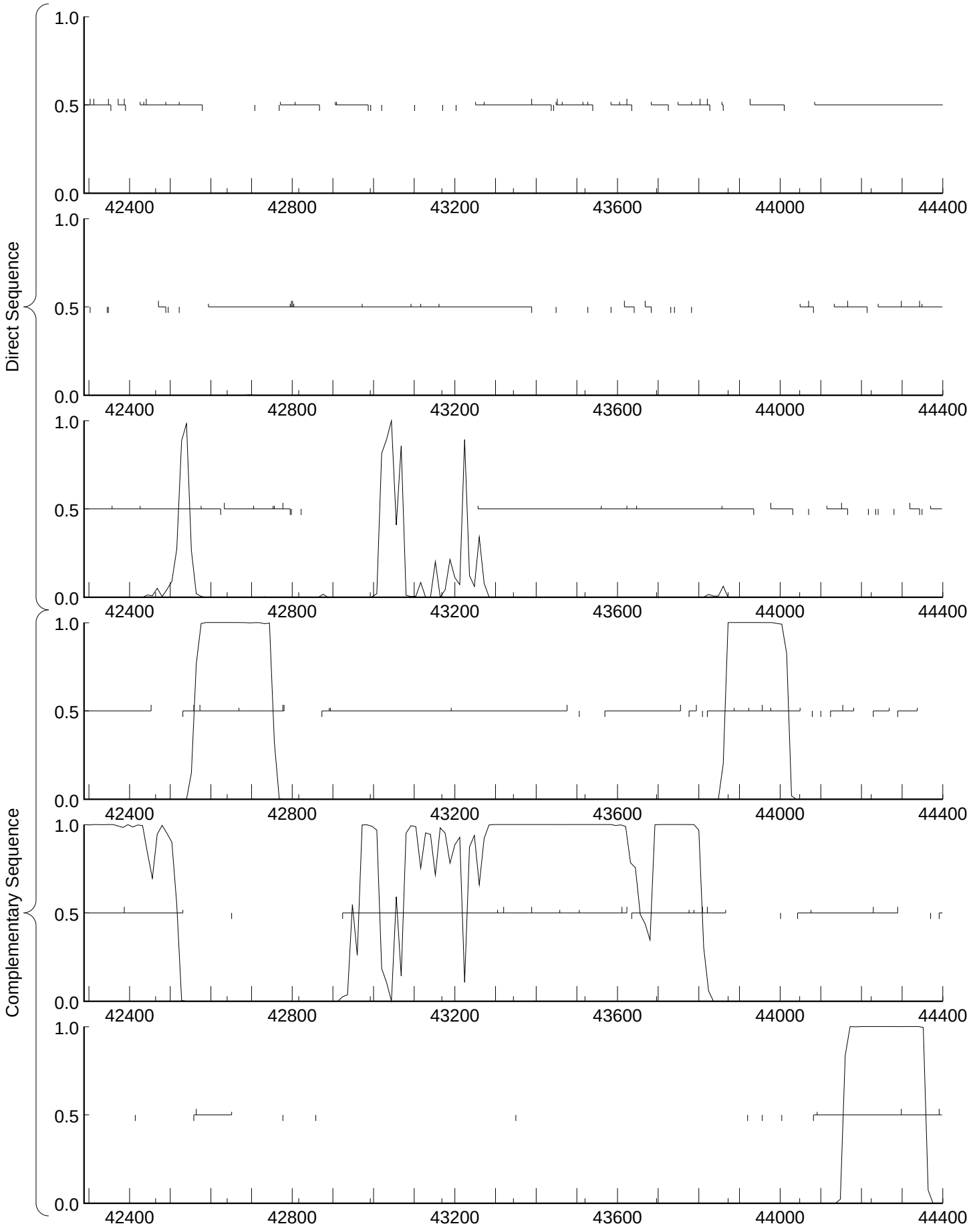

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Nucleotide Position

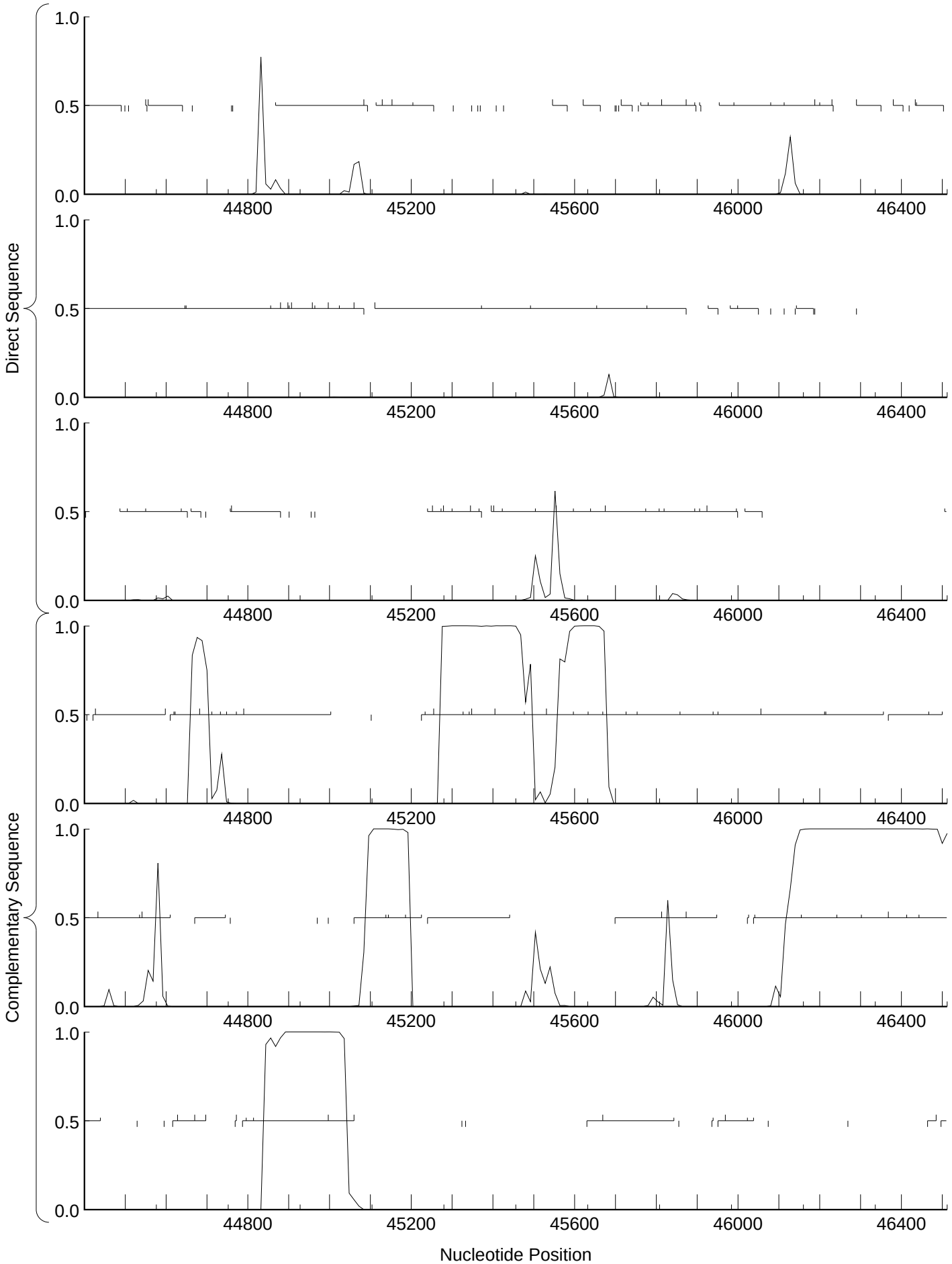

Direct Sequence
Complementary Sequence
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Nucleotide Position

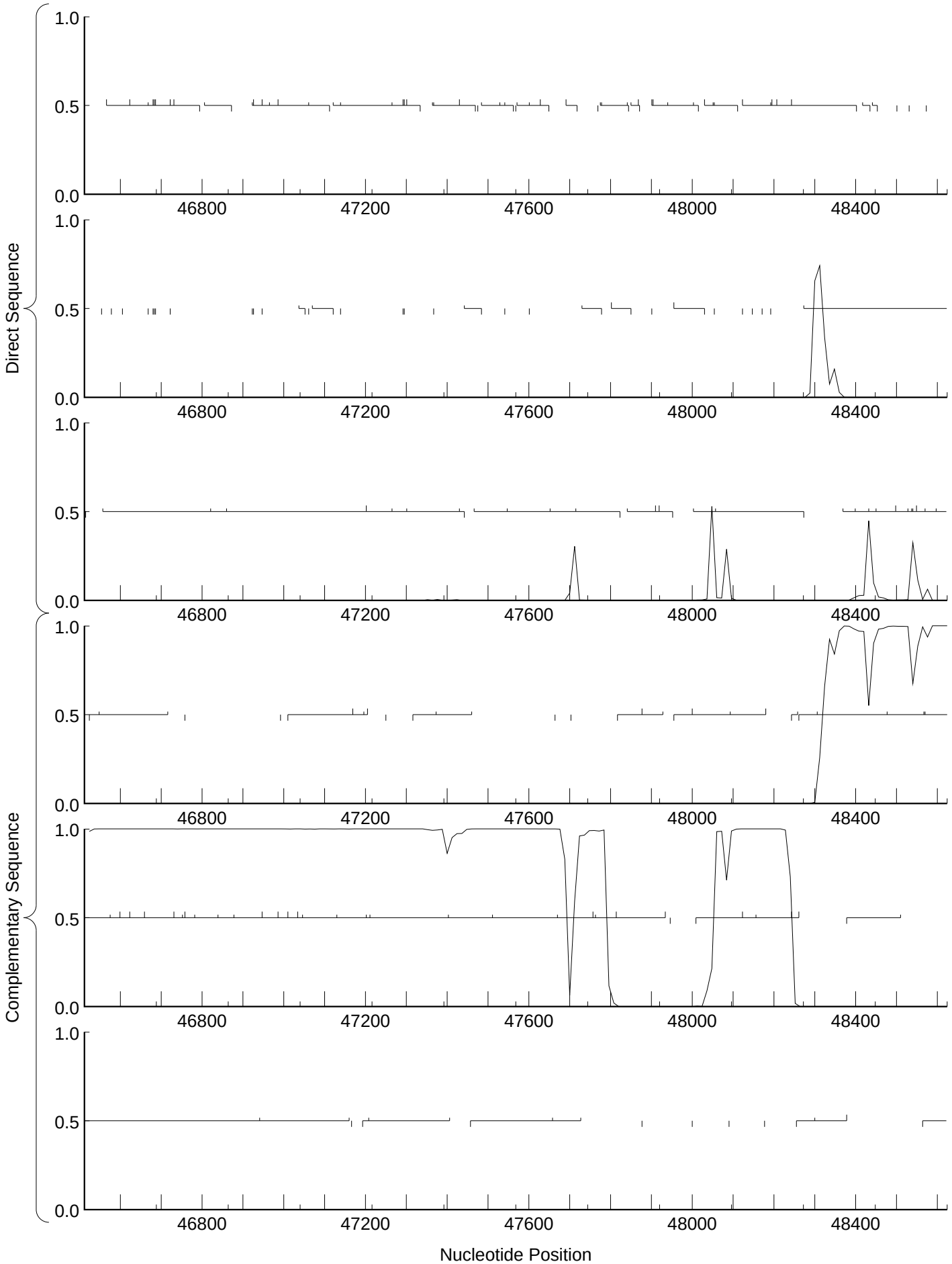

Direct Sequence
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37600
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Nucleotide Position

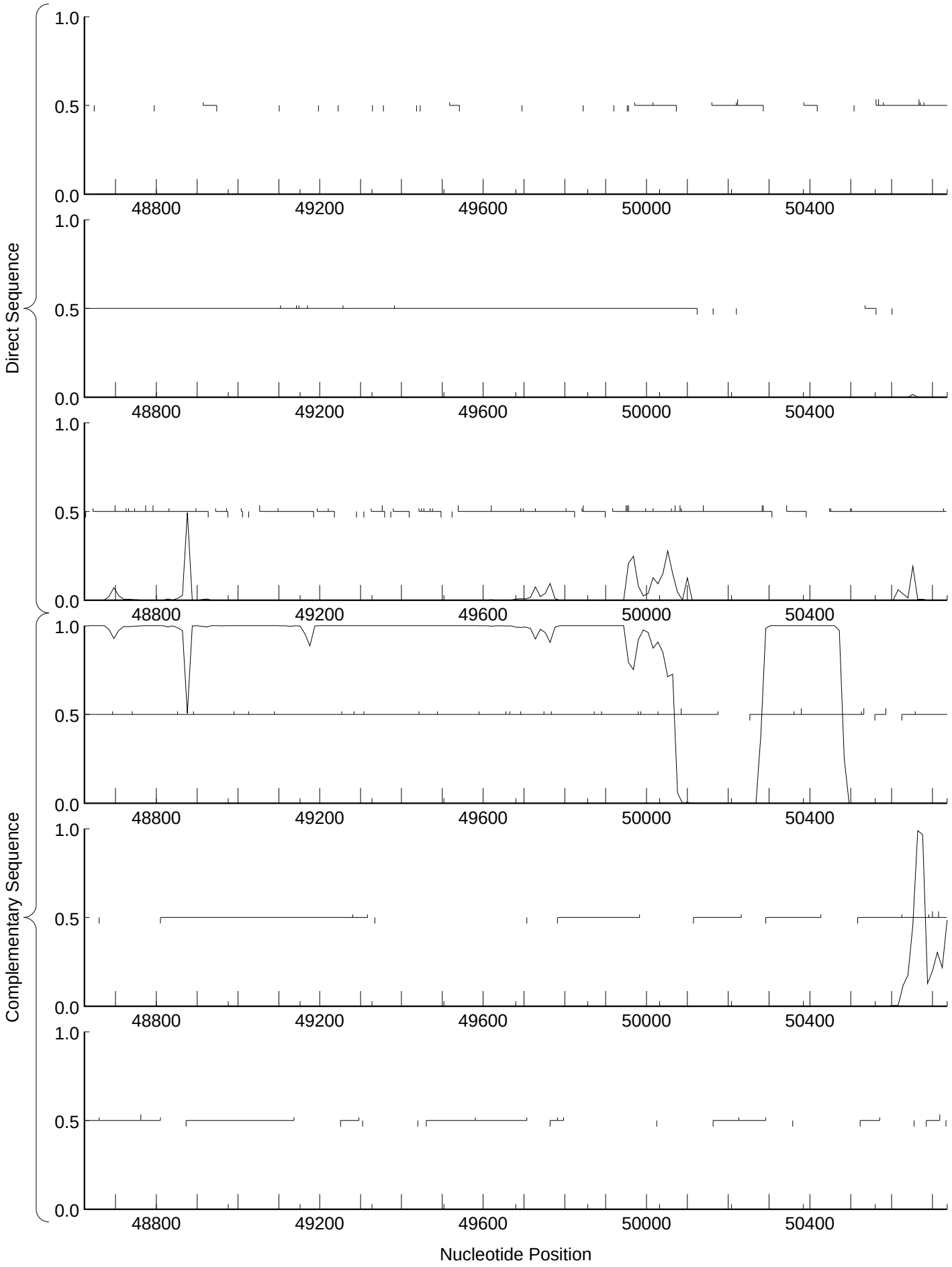

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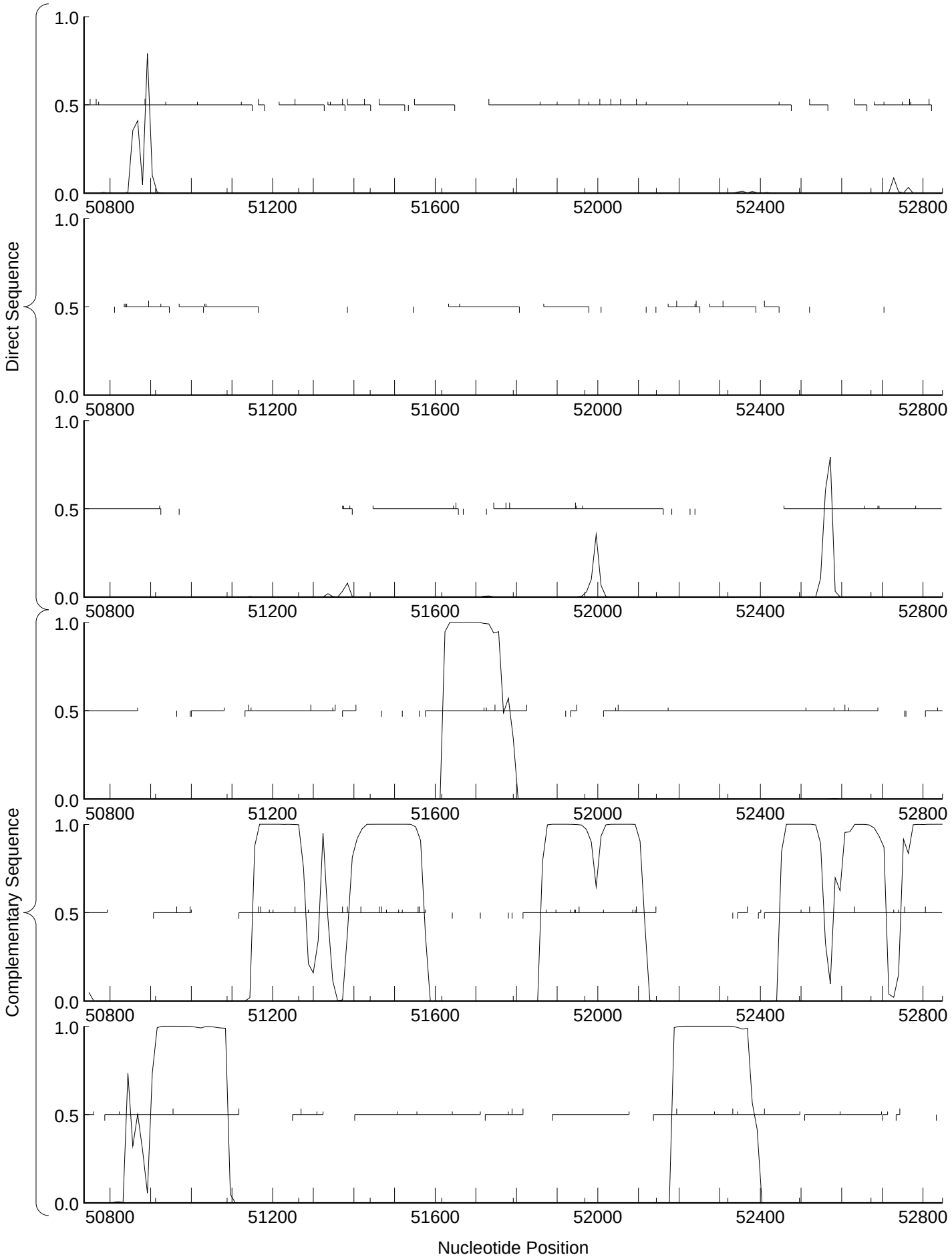

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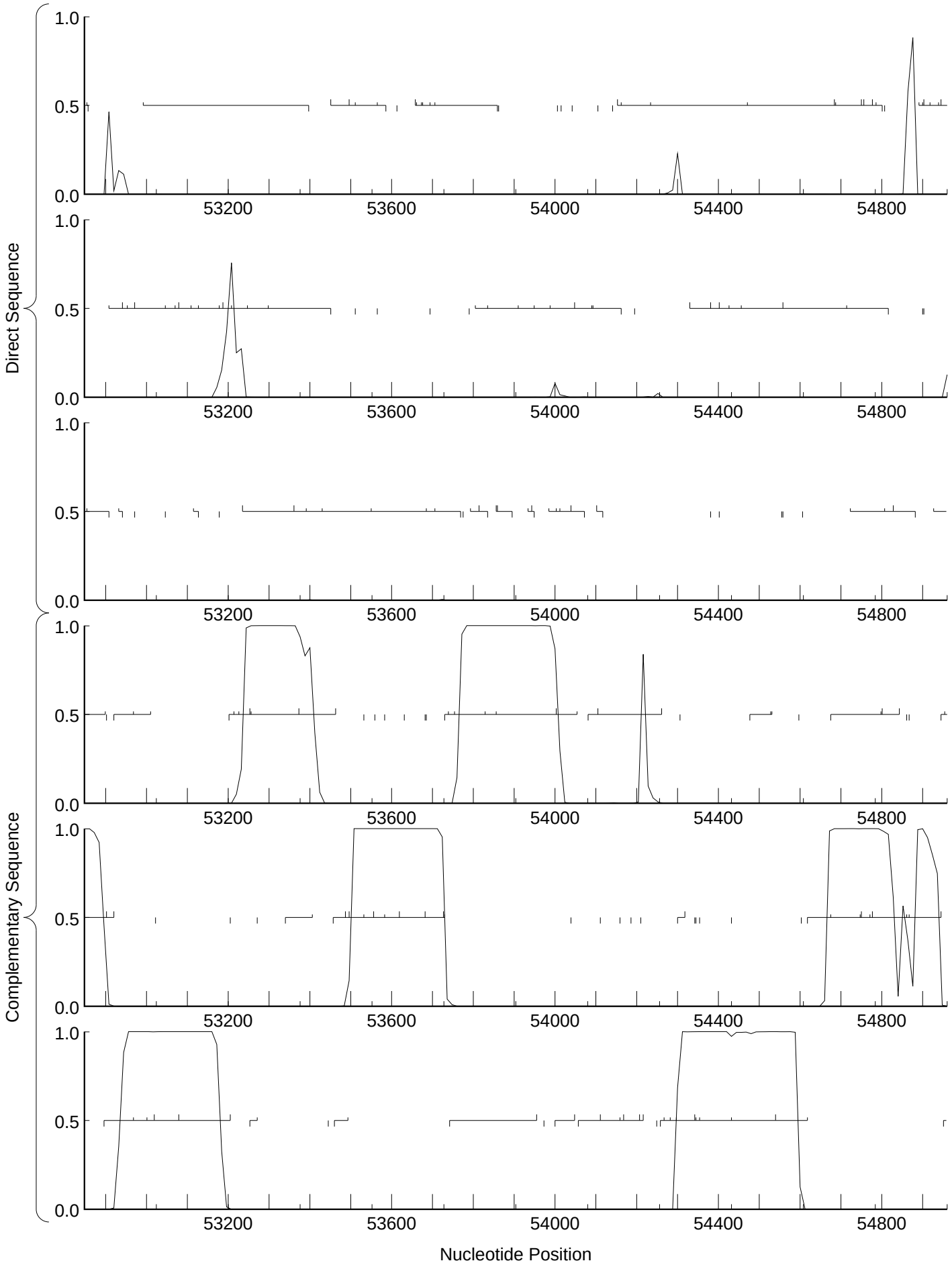

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Nucleotide Position

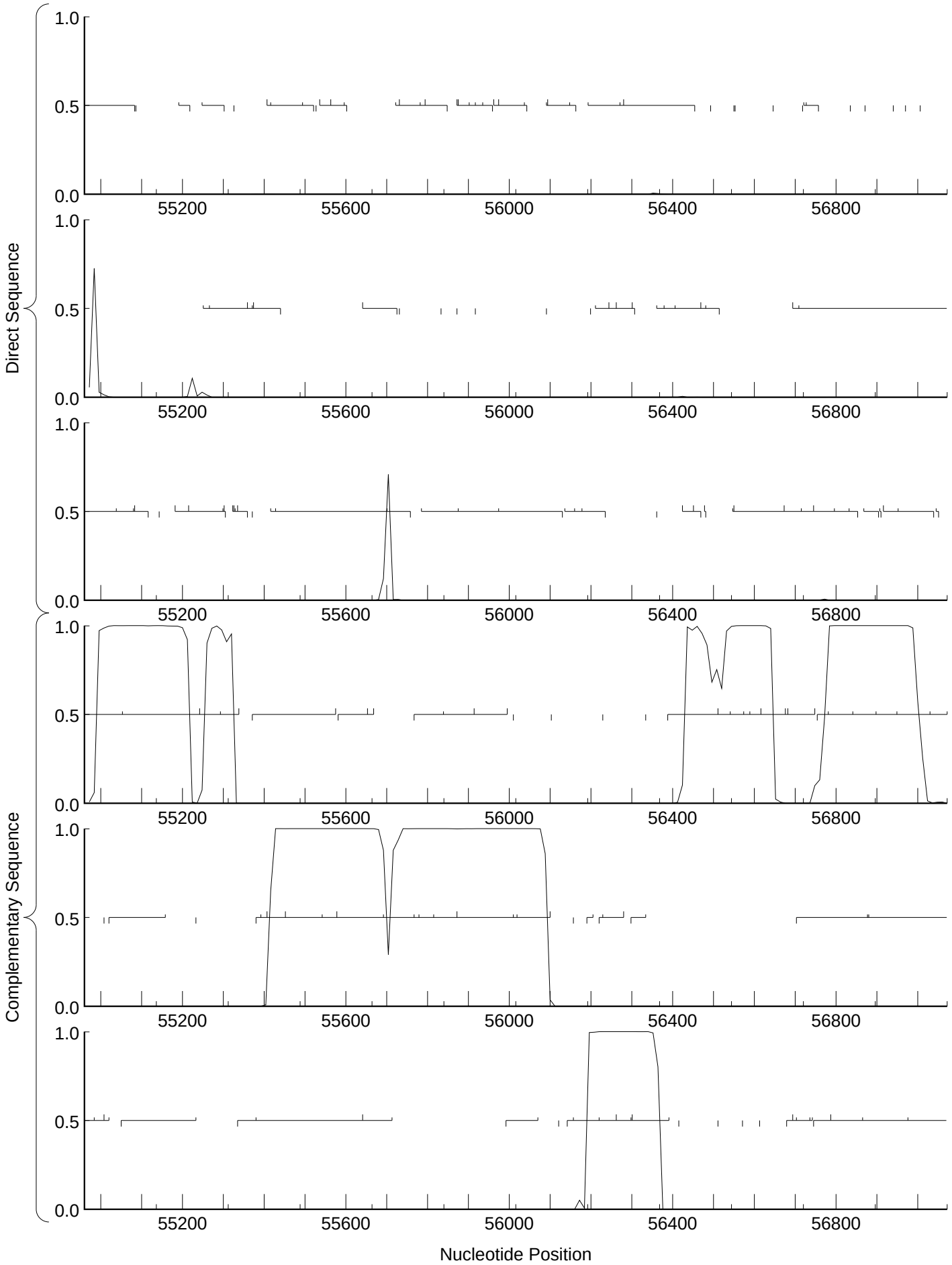

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Nucleotide Position

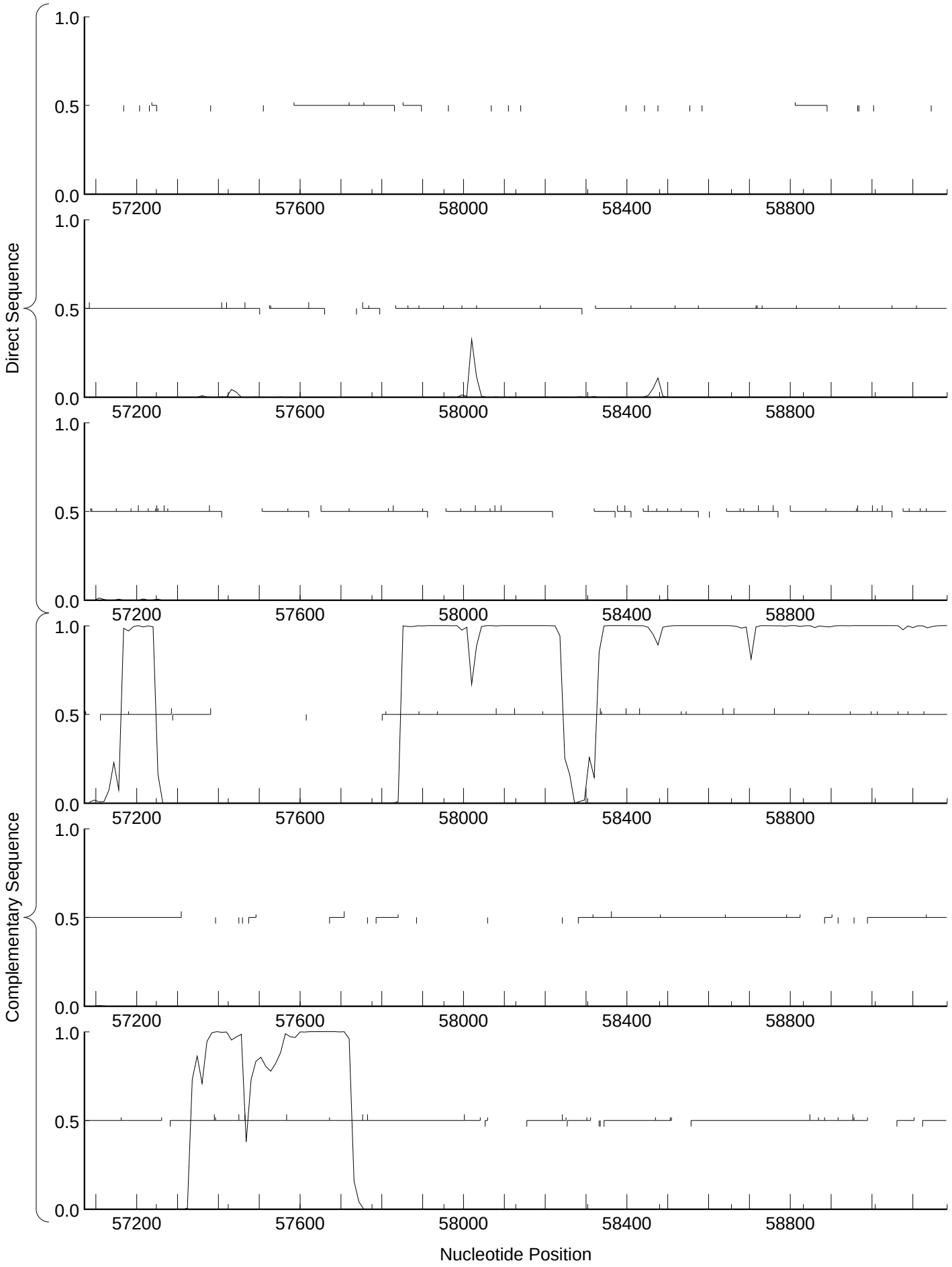

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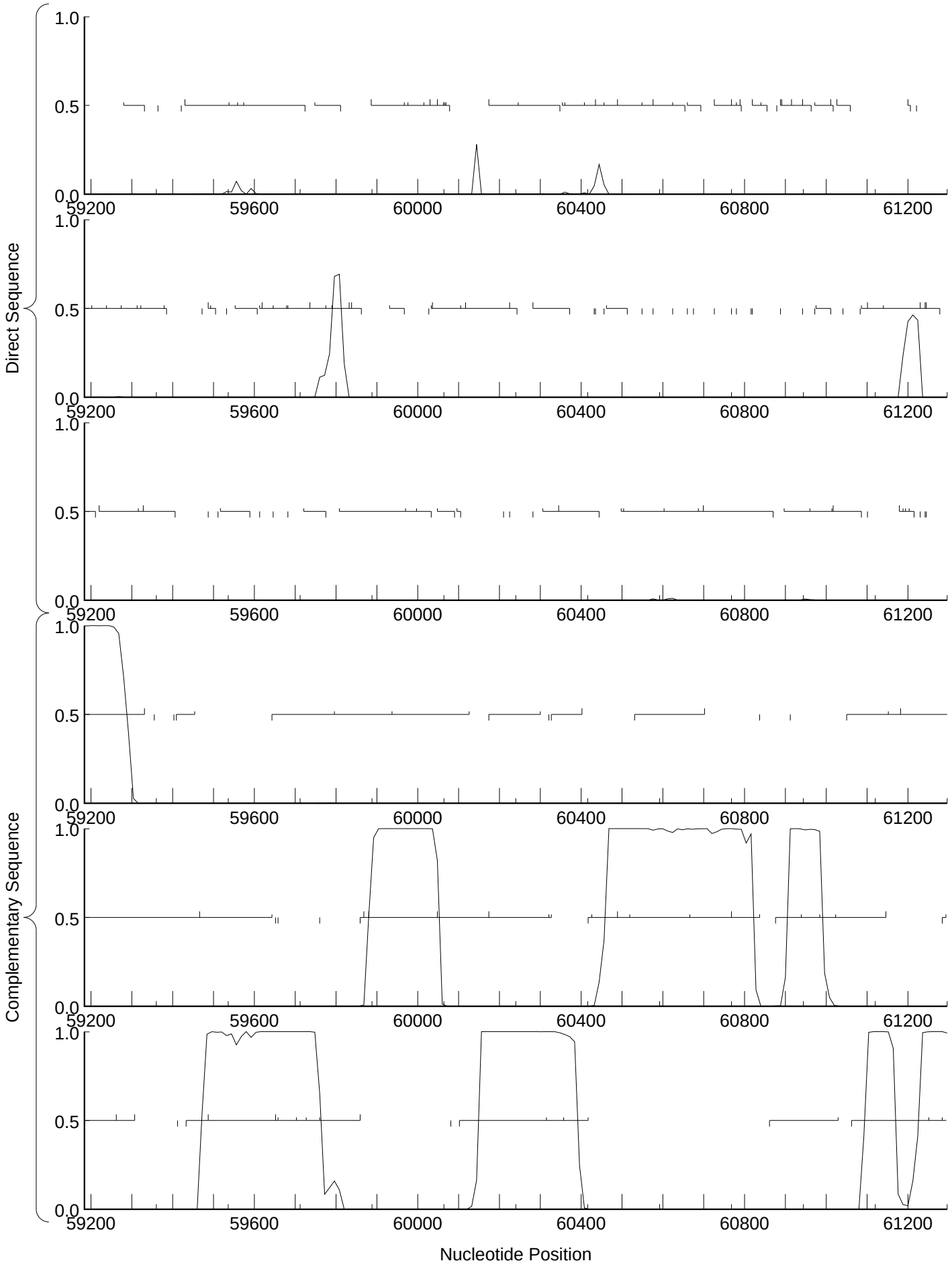

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Nucleotide Position

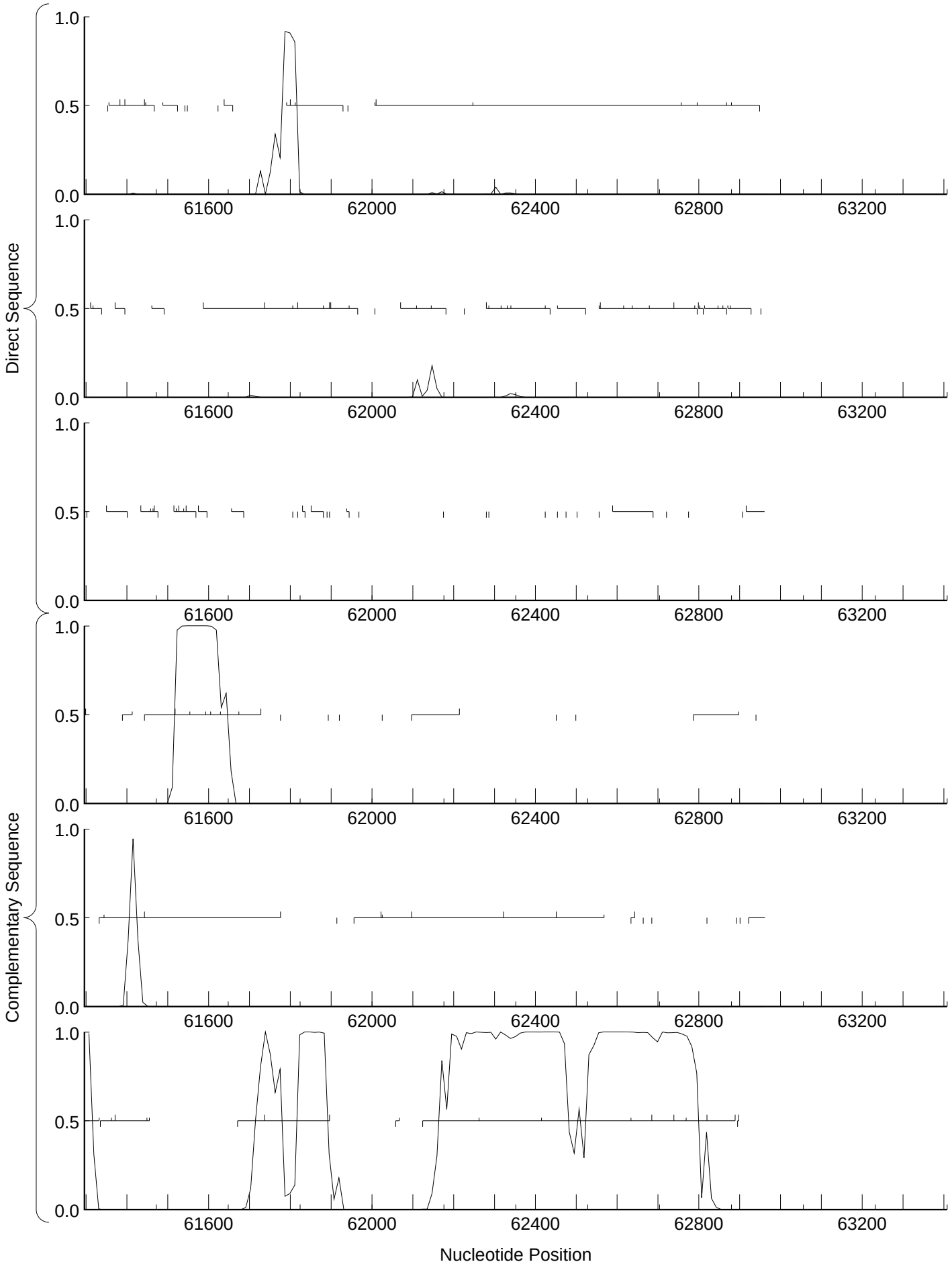

Direct Sequence
Complementary Sequence
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0.0
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52000
52400
52800
Nucleotide Position


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


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
55200
55600
56000
56400
56800
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
57200
57600
58000
58400
58800
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
59200
59600
60000
60400
60800
61200
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
61600
62000
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63200
Nucleotide Position