GeneMark

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

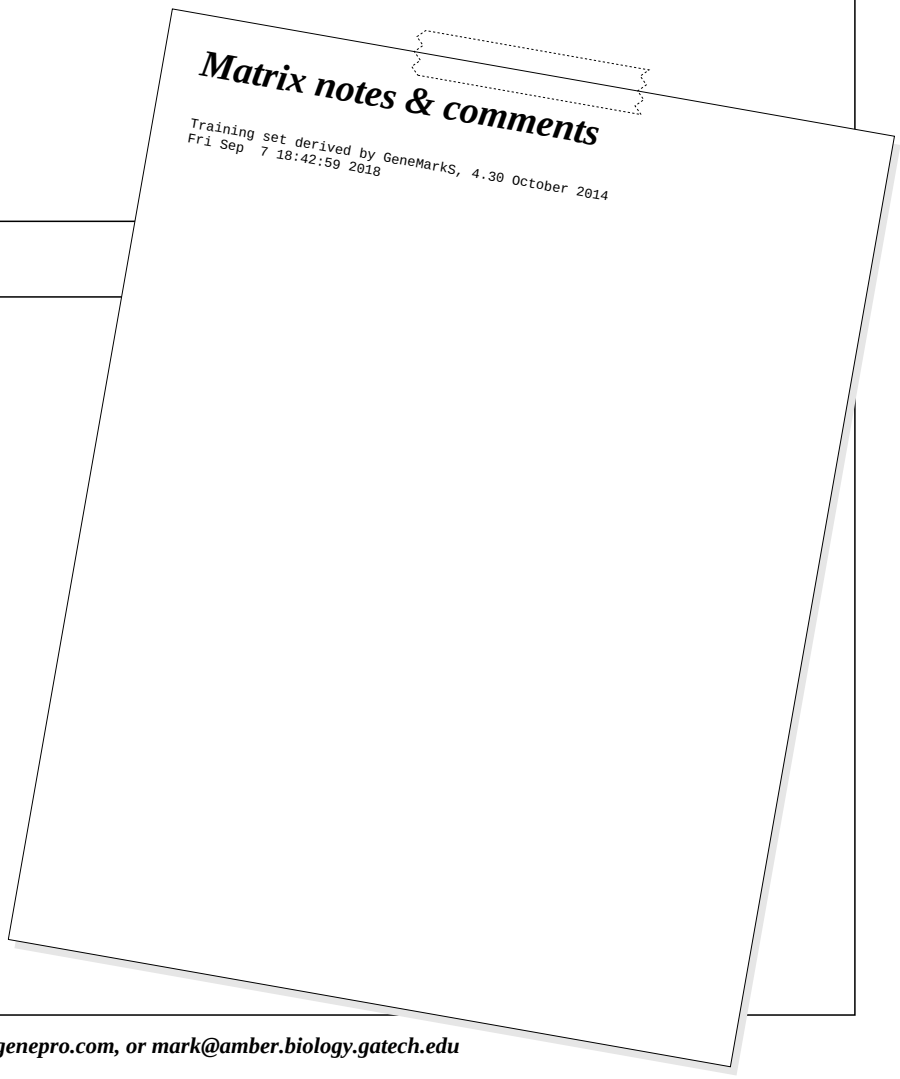
PROGRAM INFORMATION

```
Sequence         :  Microbacterium phage Didgeridoo complete sequence, 42655 bp including 10-base 3' overhang (
Analysis Date    :  3/28/24 at 21:39:10
Pages            :  22
Sequence Length  :  42655 bp
GC Content       :   66.07%

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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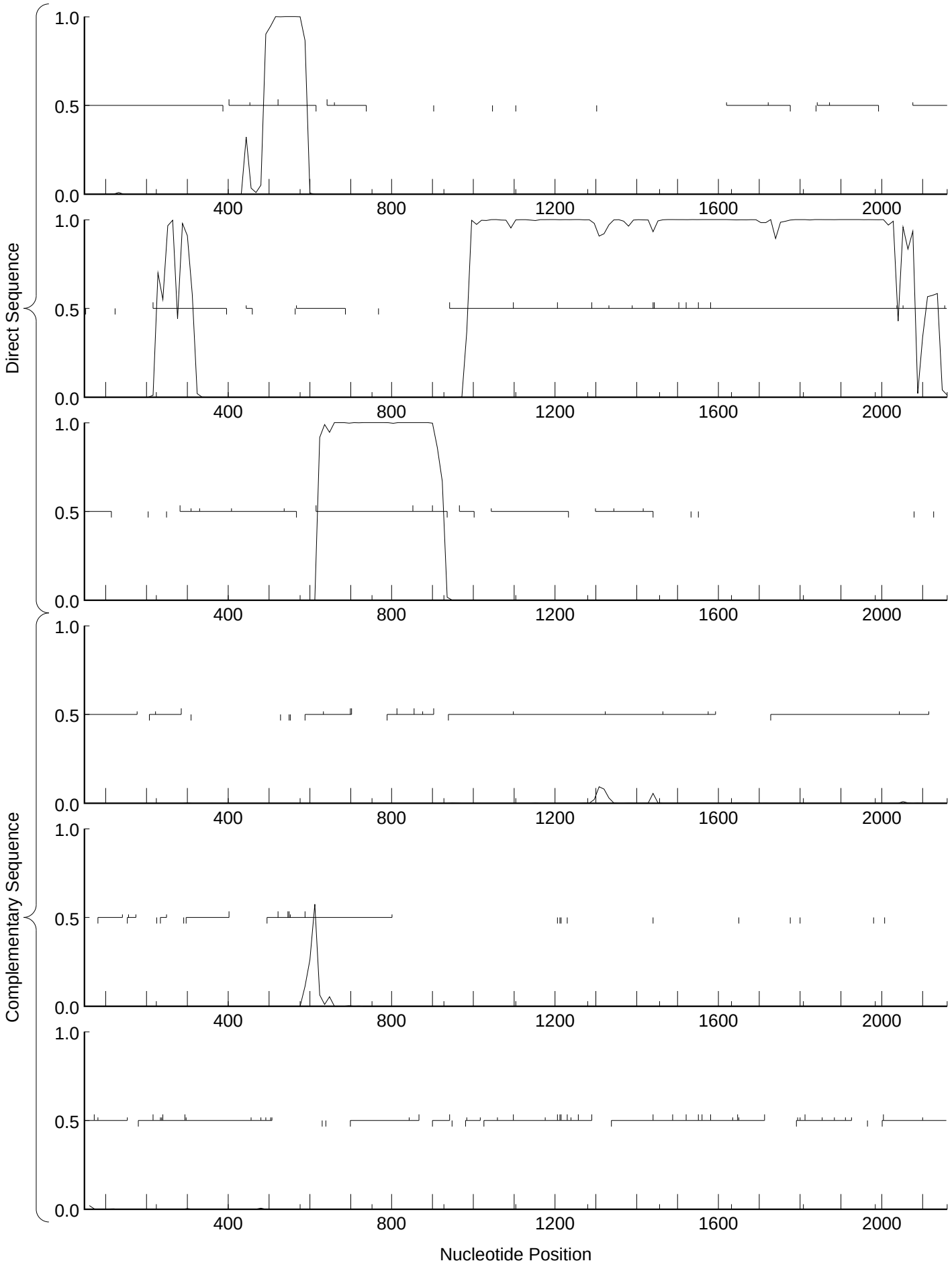
MATRIX INFORMATION

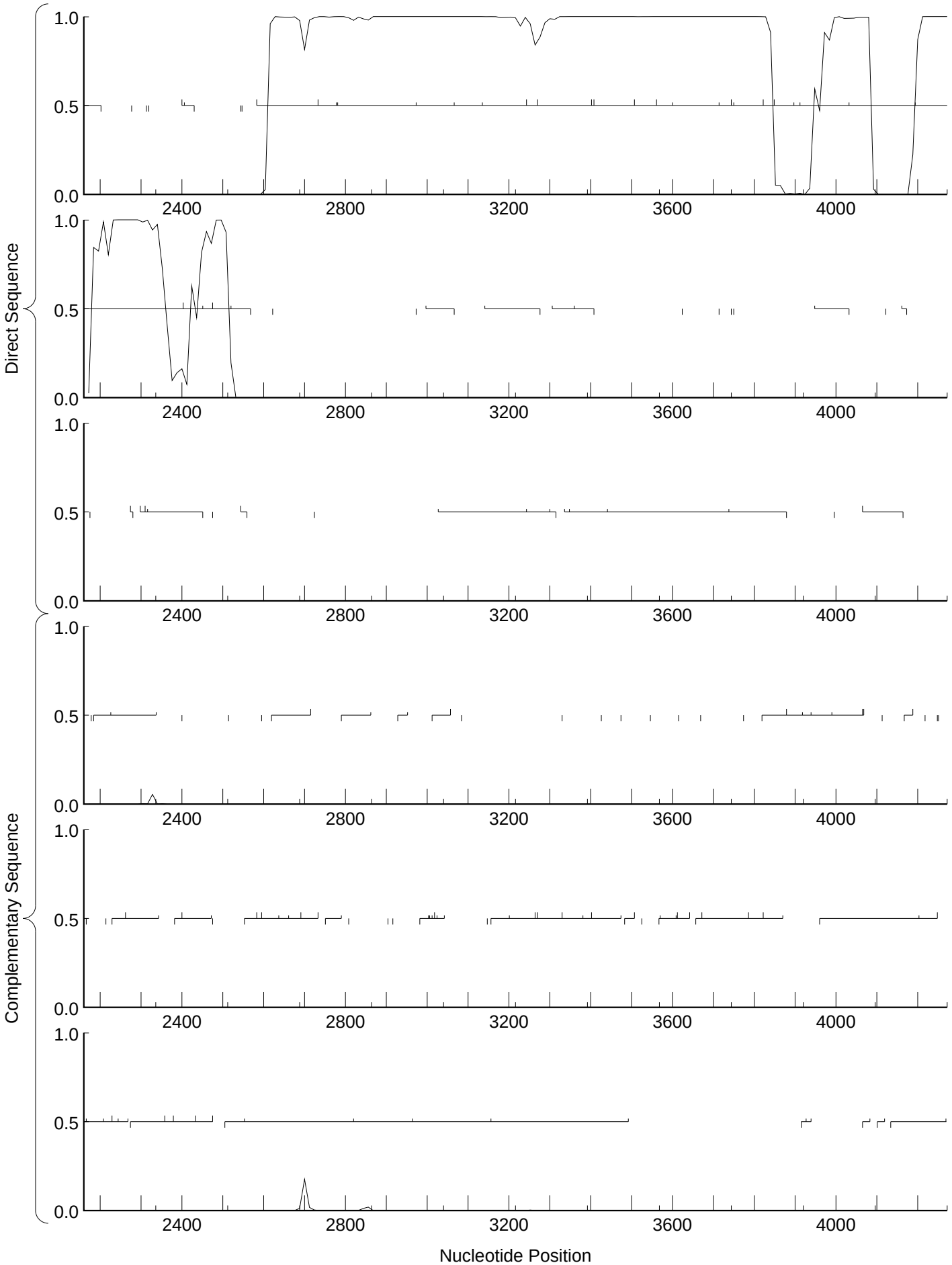
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Matrix     :  -
Author     :  -

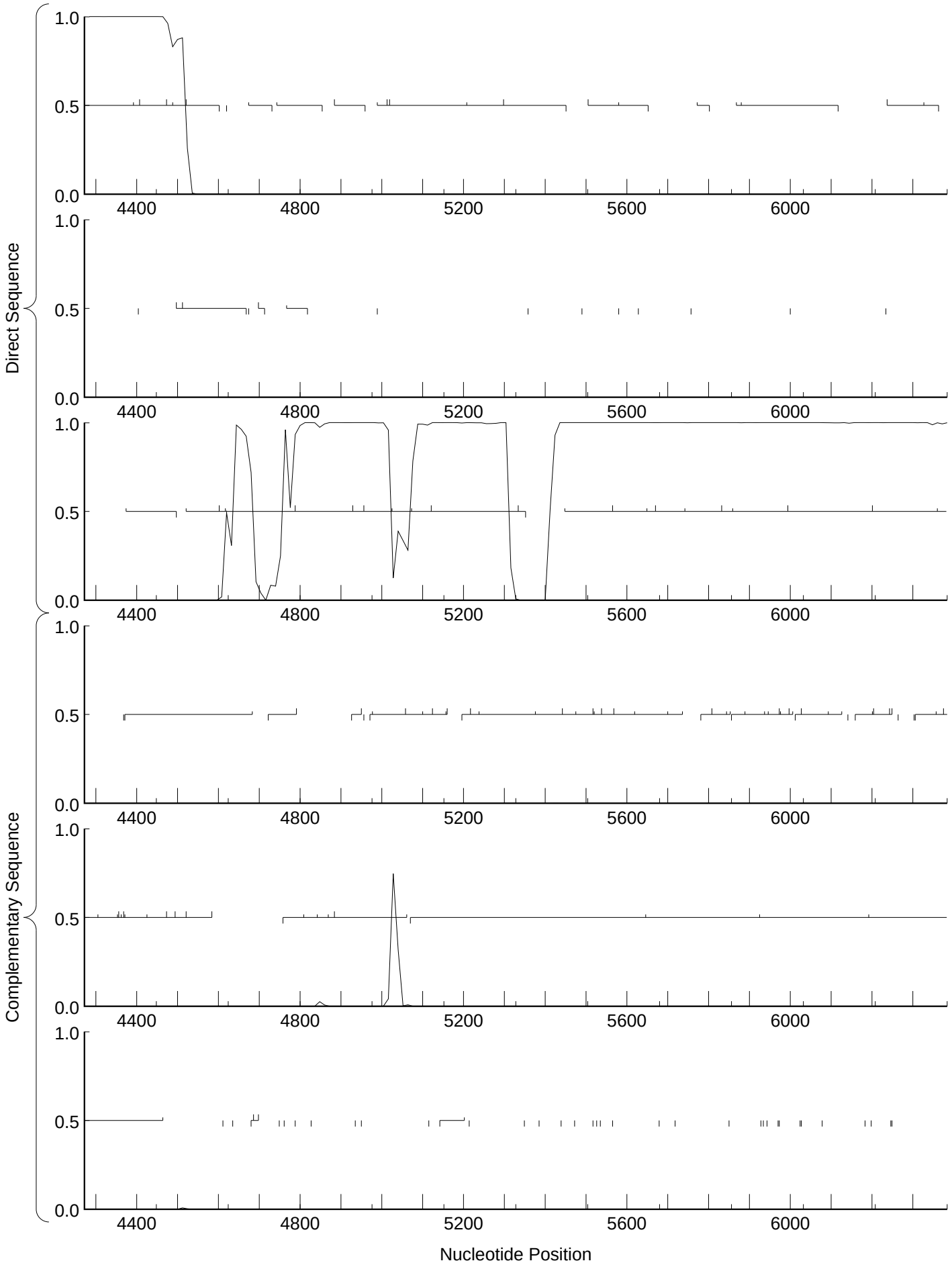
Order      :  4
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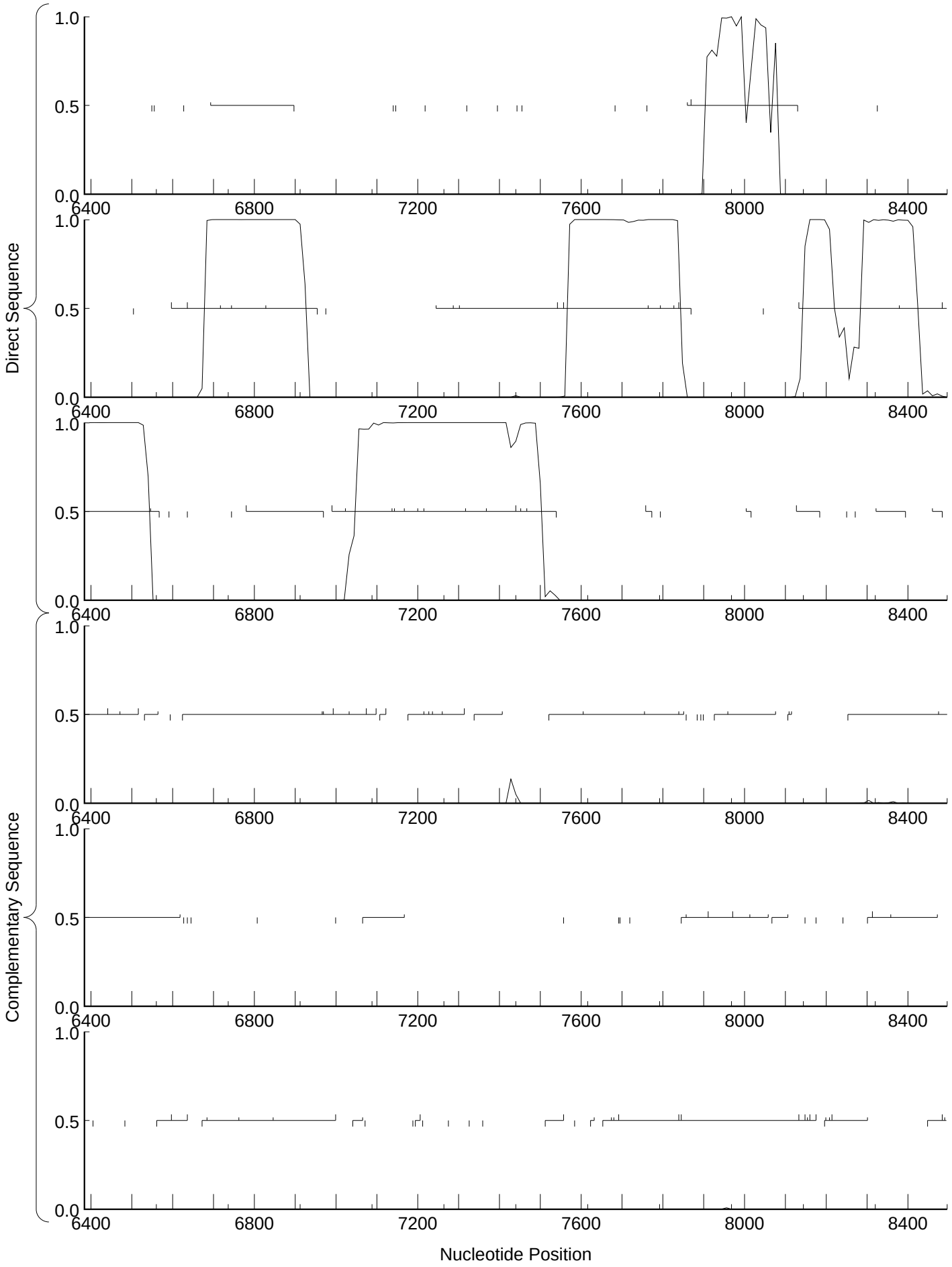

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

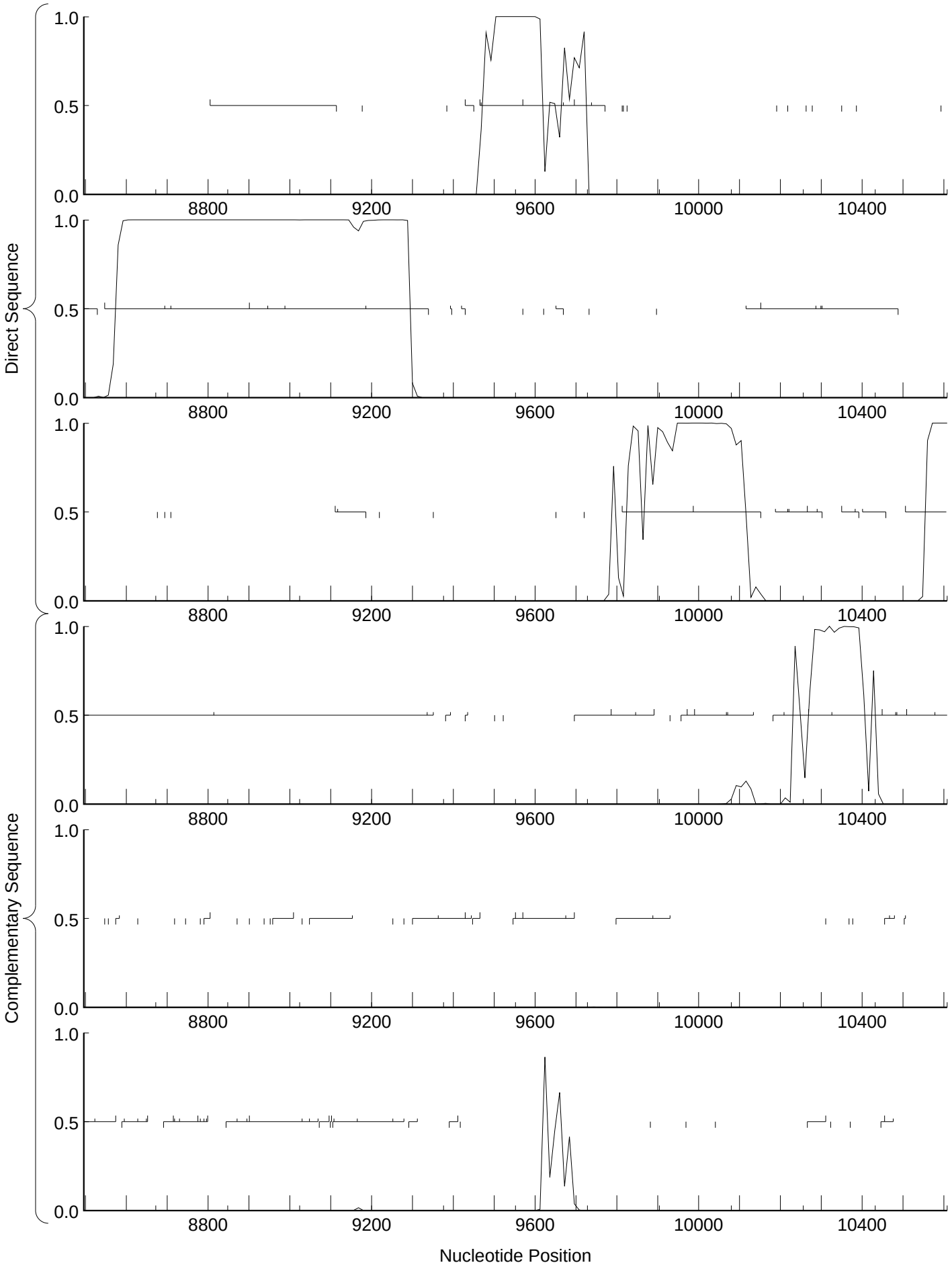

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

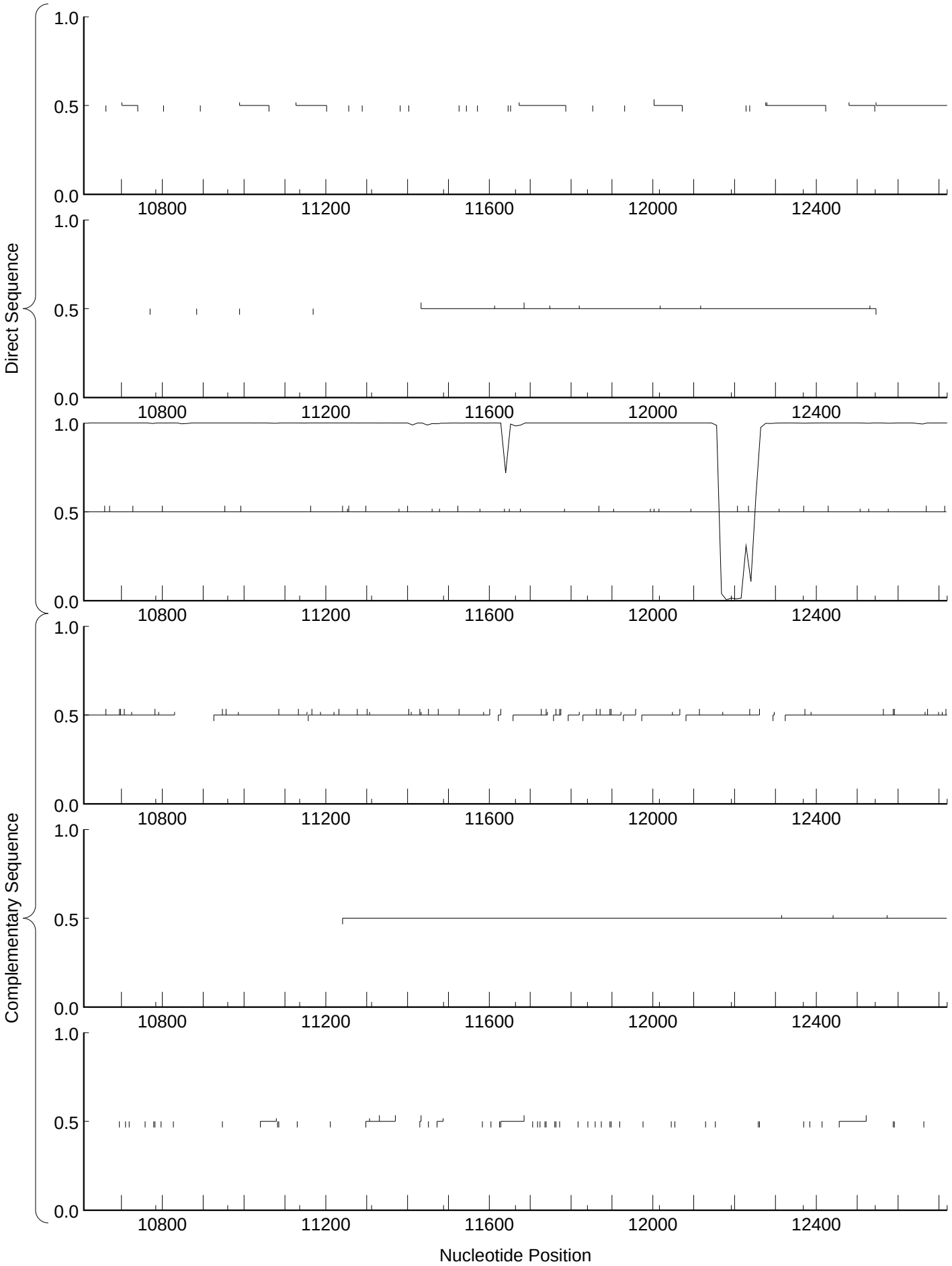

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
400
800
1200
1600
2000
Nucleotide Position

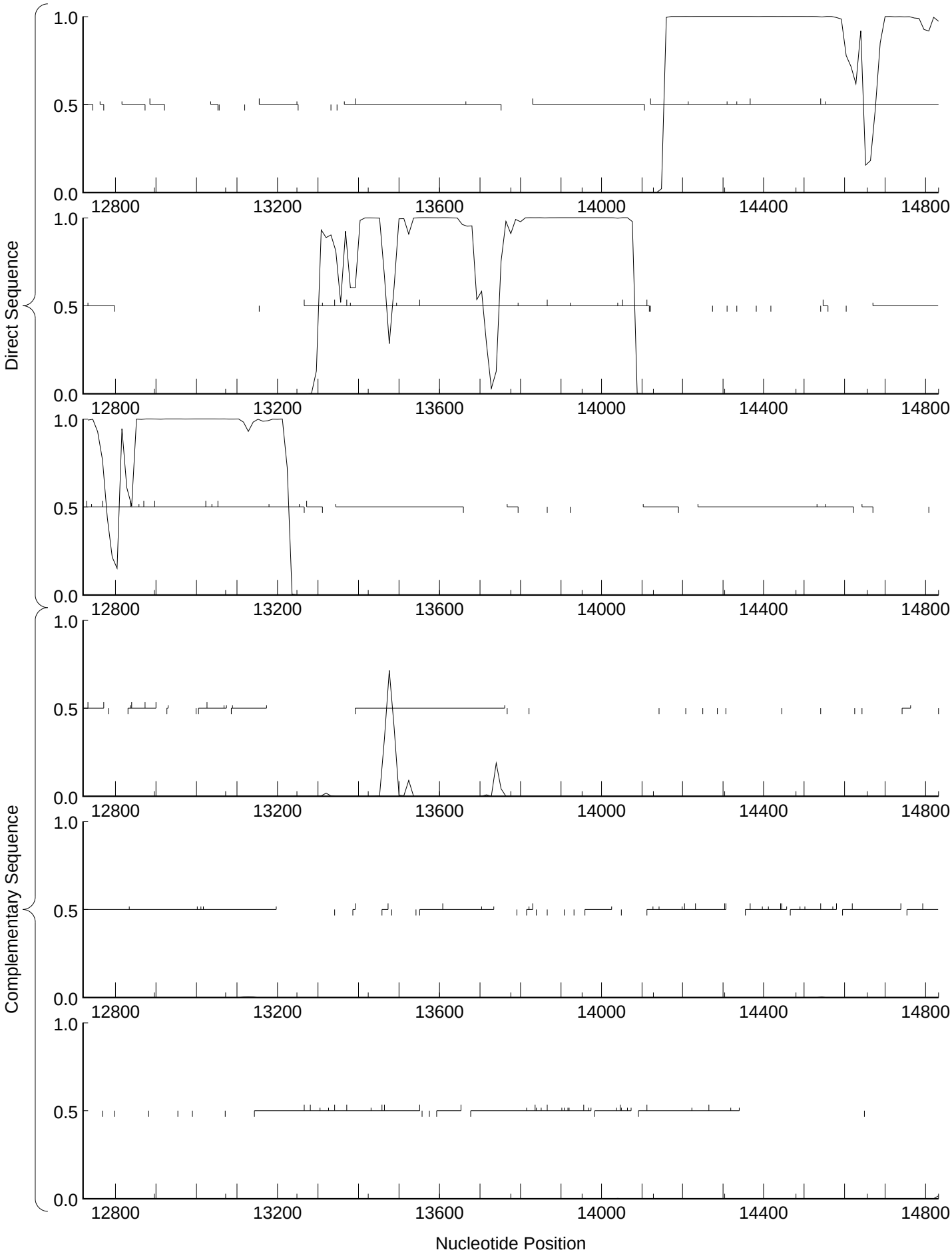

Direct Sequence
Complementary Sequence
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3600
4000
Nucleotide Position

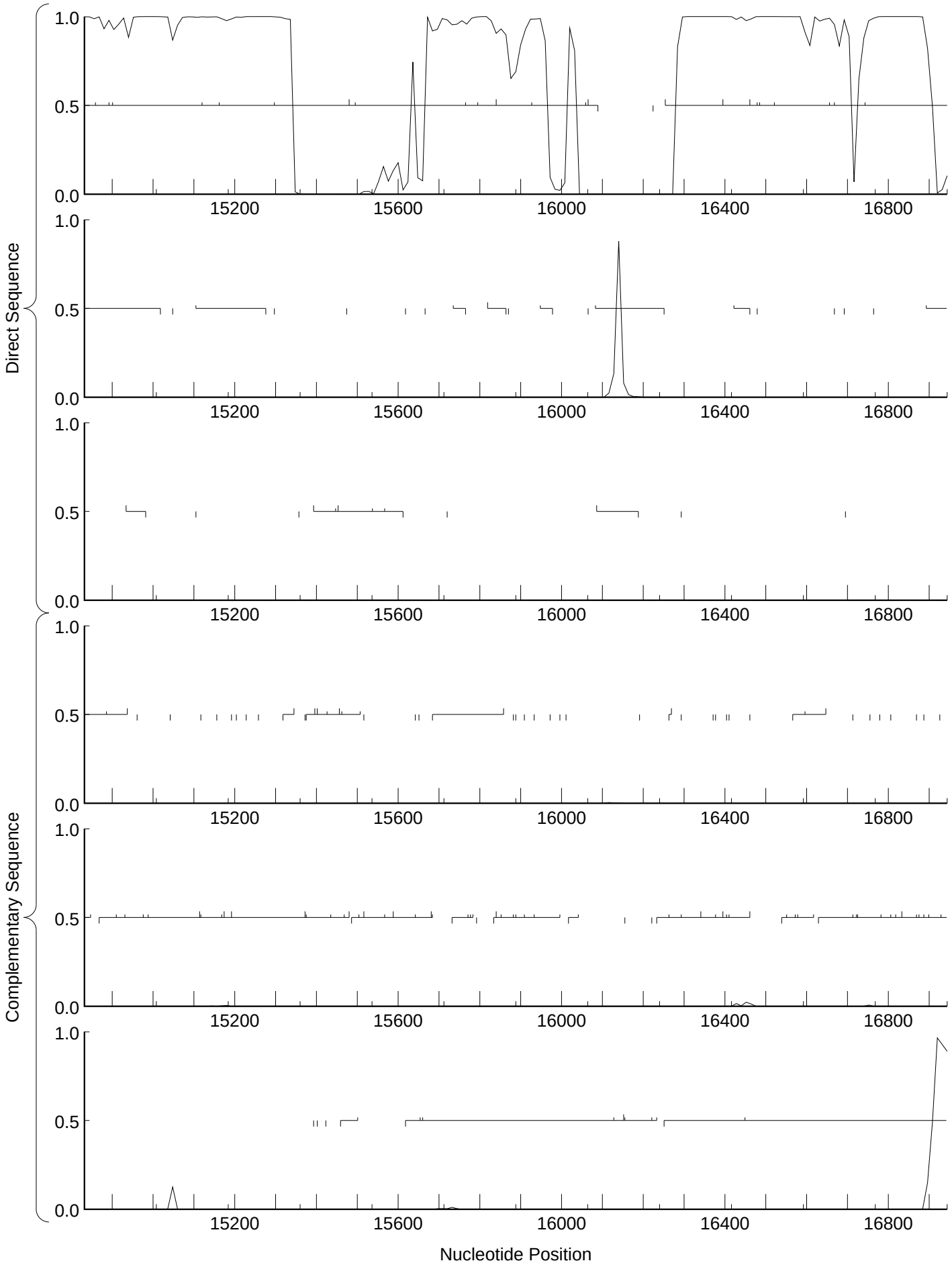

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Complementary Sequence
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5200
5600
6000
Nucleotide Position

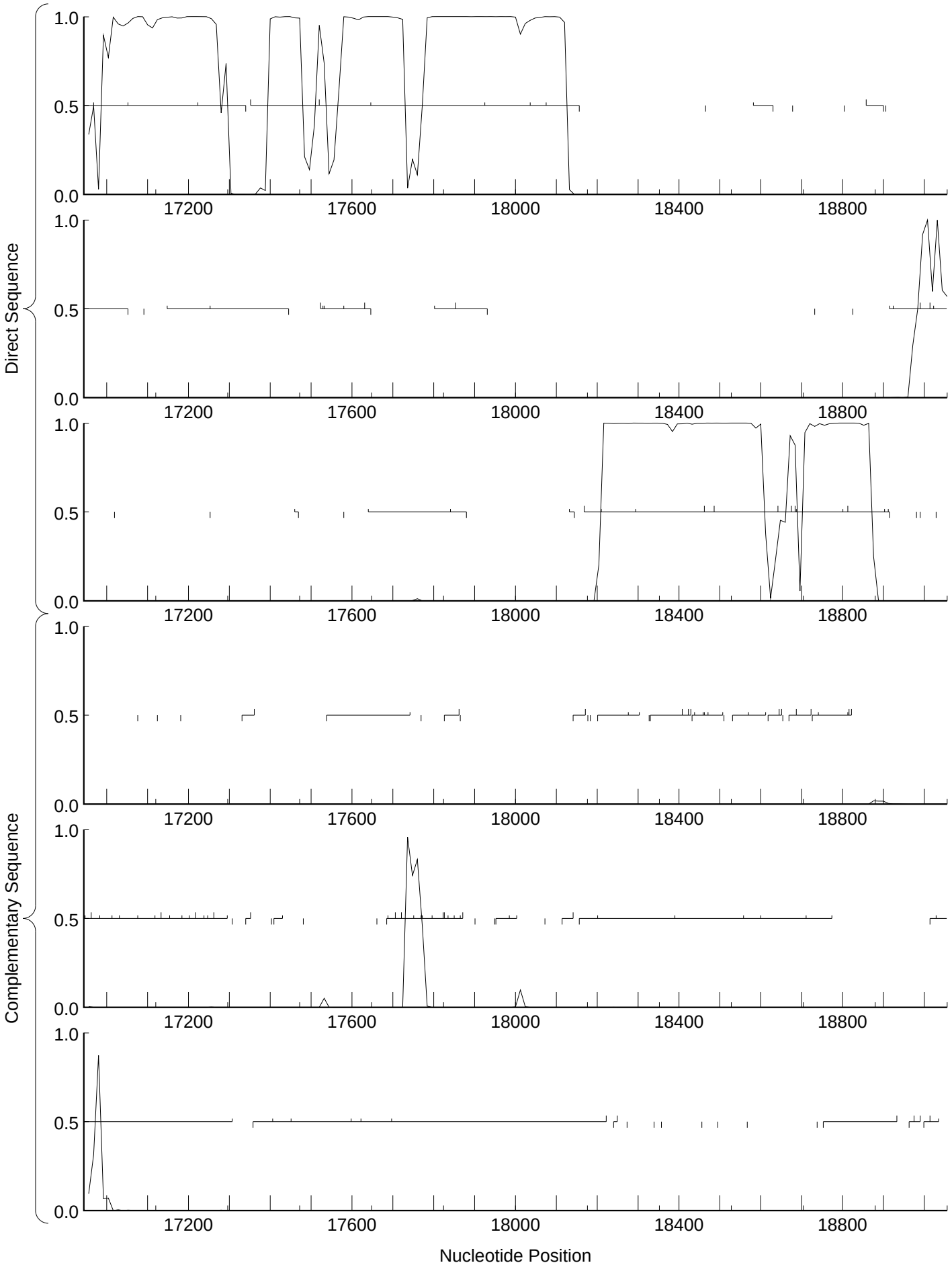

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Complementary Sequence
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6800
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7600
8000
8400
Nucleotide Position

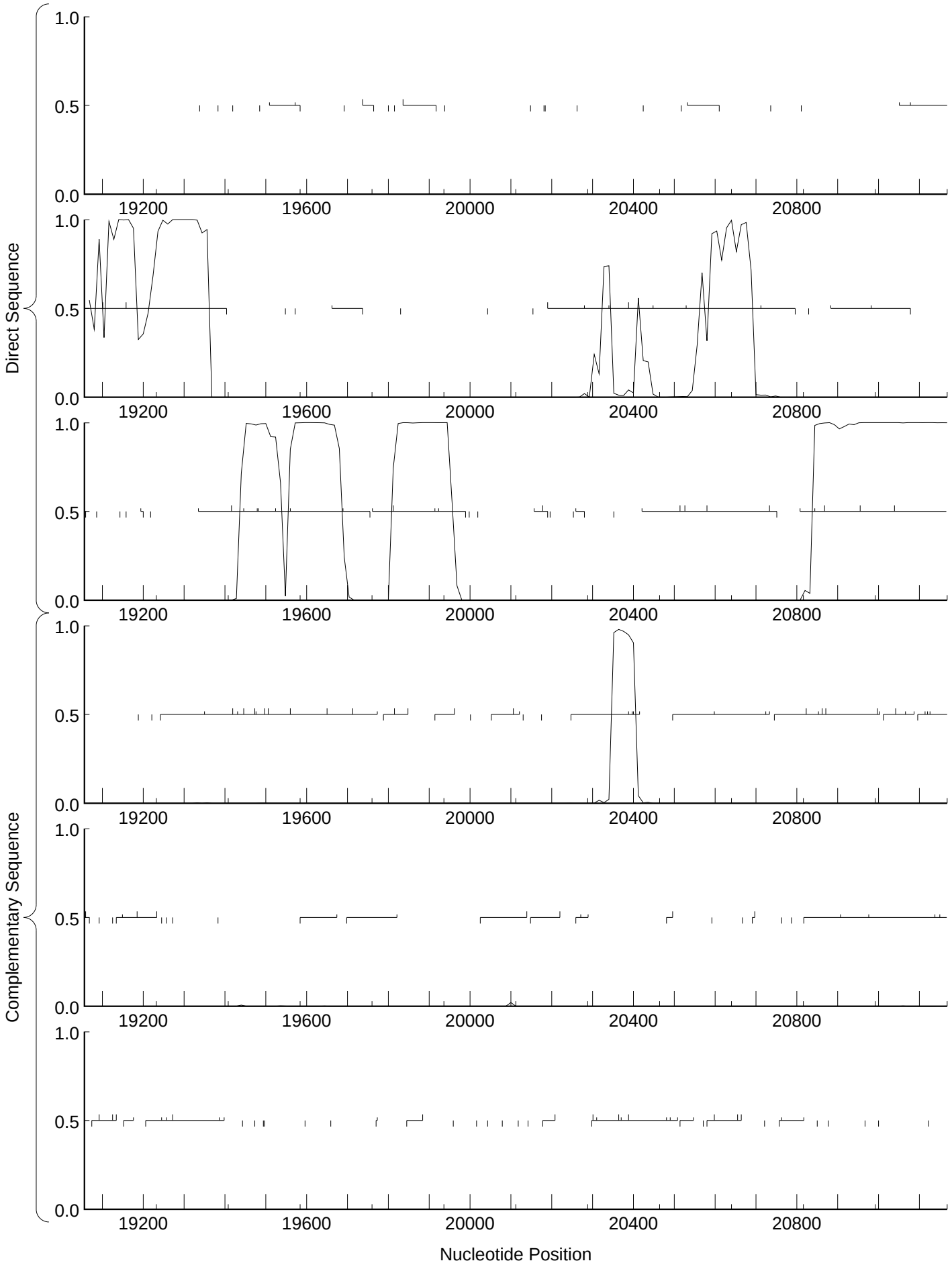

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Complementary Sequence
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10000
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Nucleotide Position

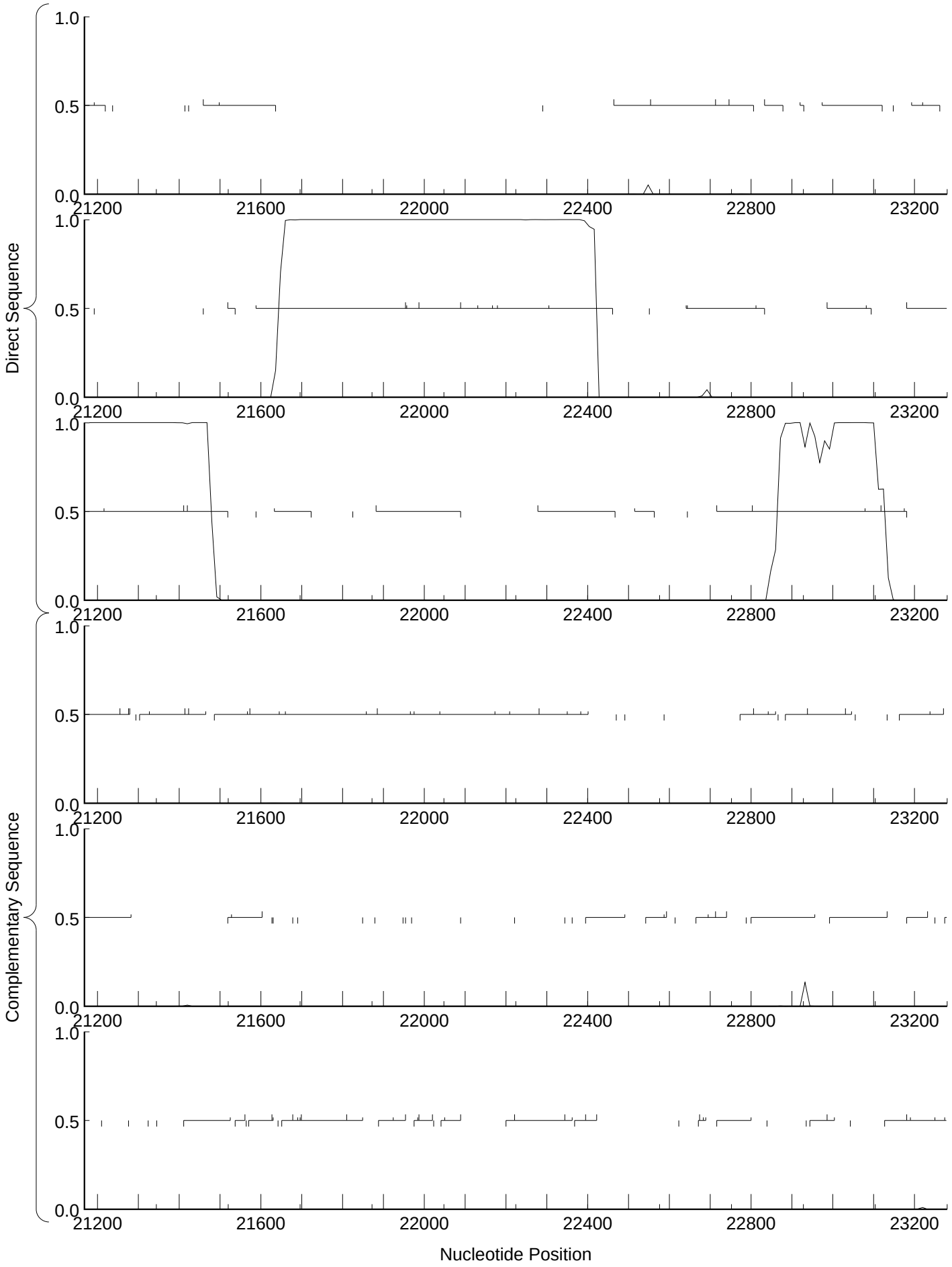

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Complementary Sequence
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0.0
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11200
11600
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Nucleotide Position

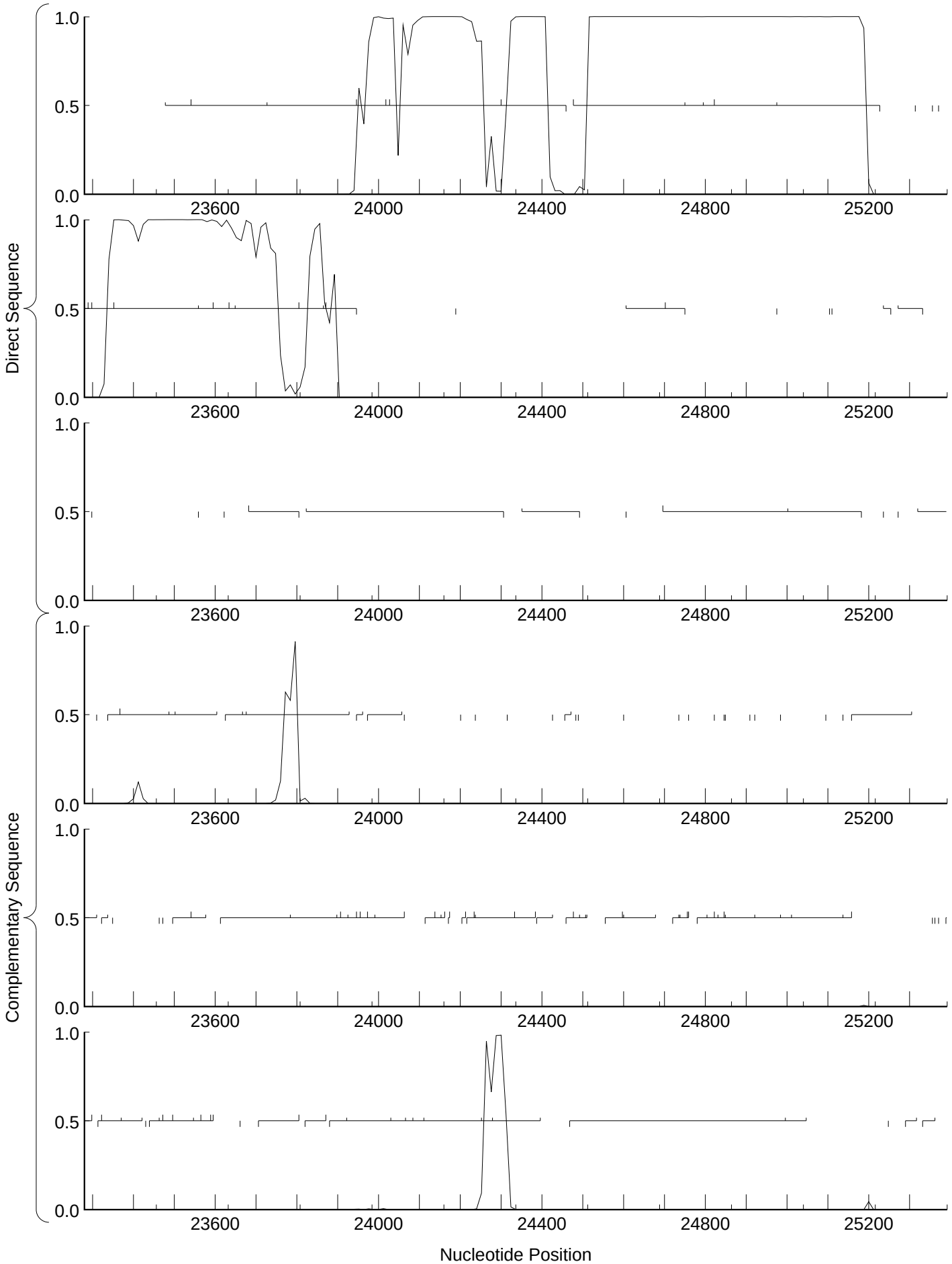

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Complementary Sequence
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14400
14800
Nucleotide Position

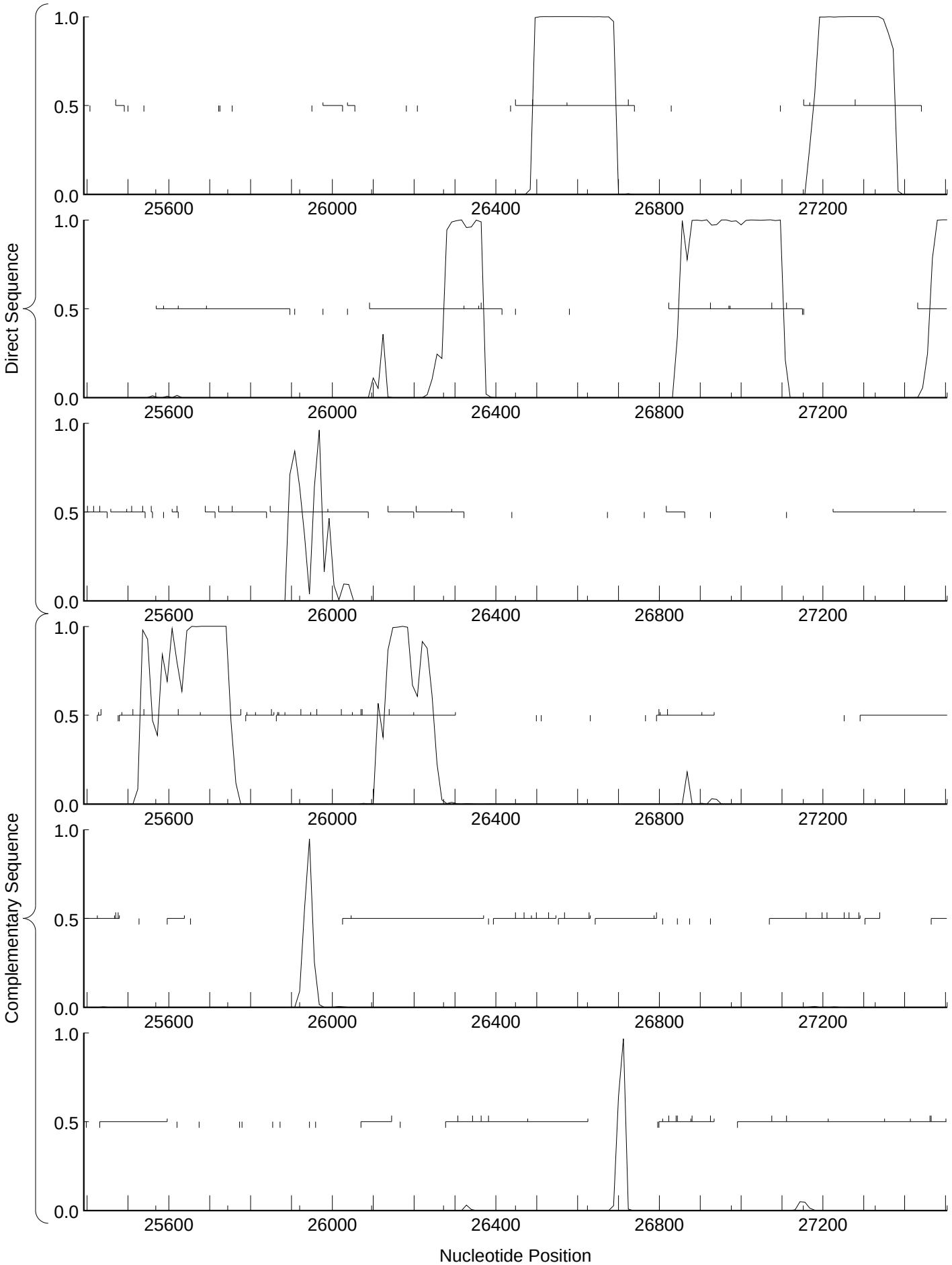

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Complementary Sequence
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Nucleotide Position

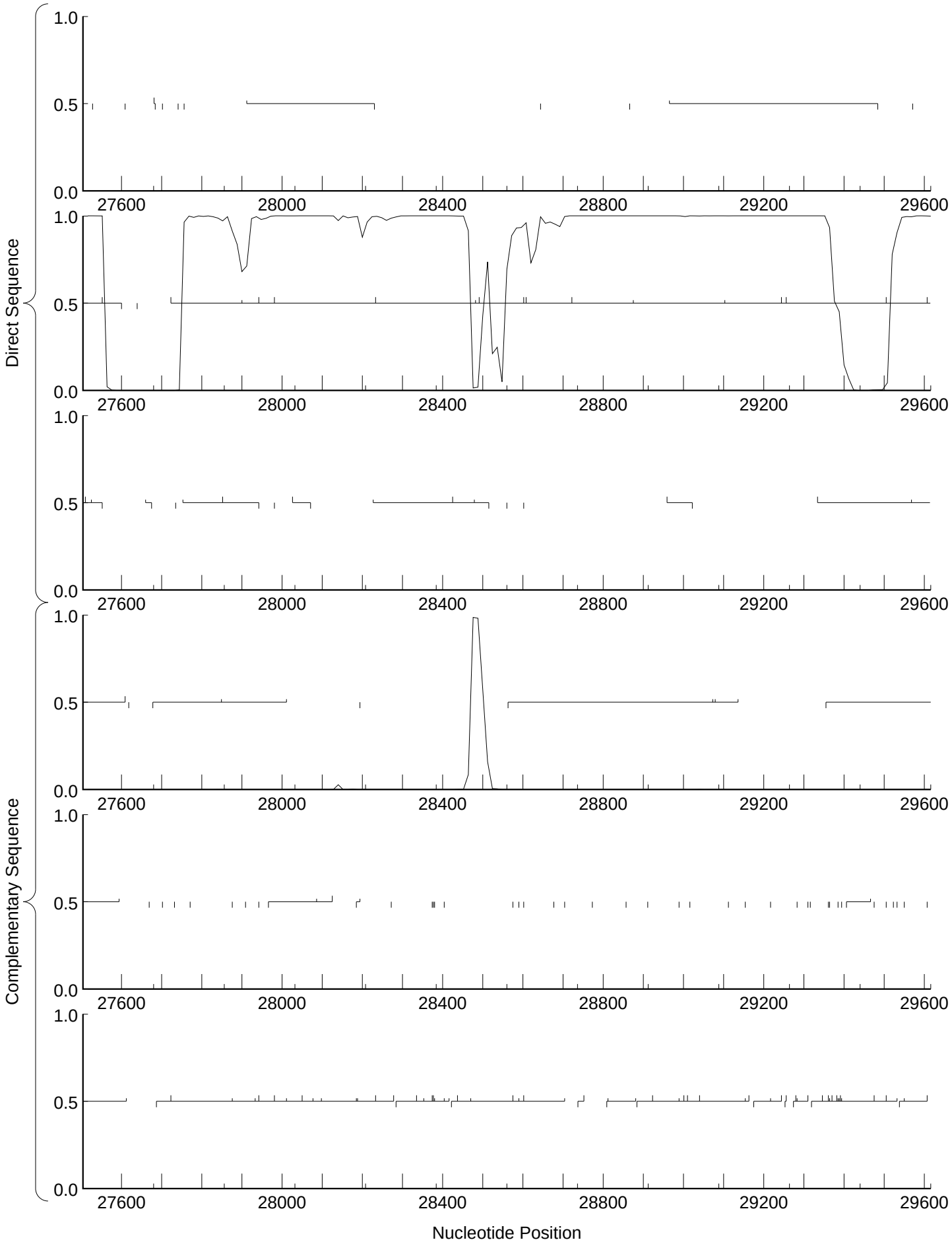

Direct Sequence
Complementary Sequence
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0.5
0.0
17200
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18000
18400
18800
Nucleotide Position

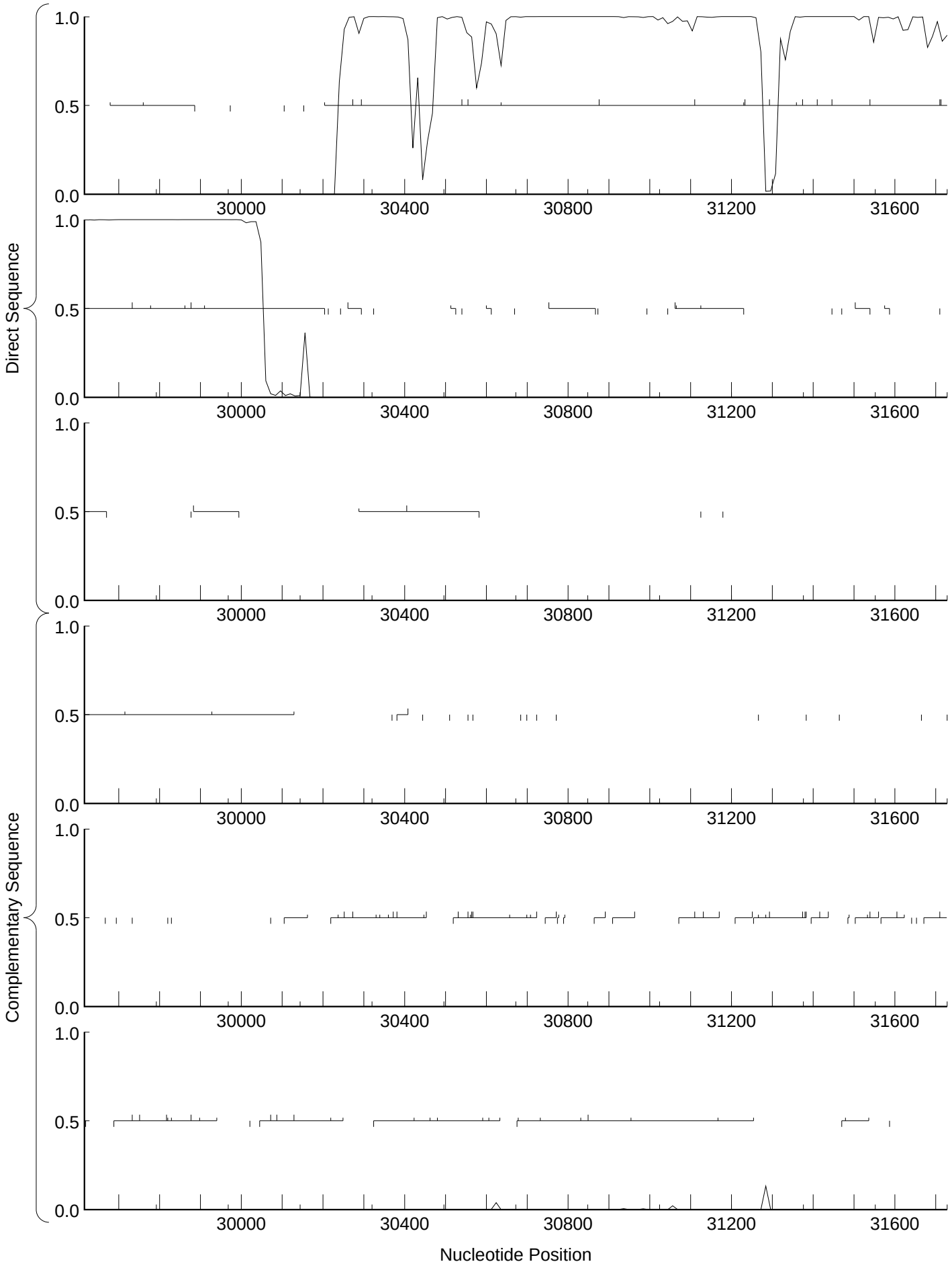

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Complementary Sequence
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20800
Nucleotide Position

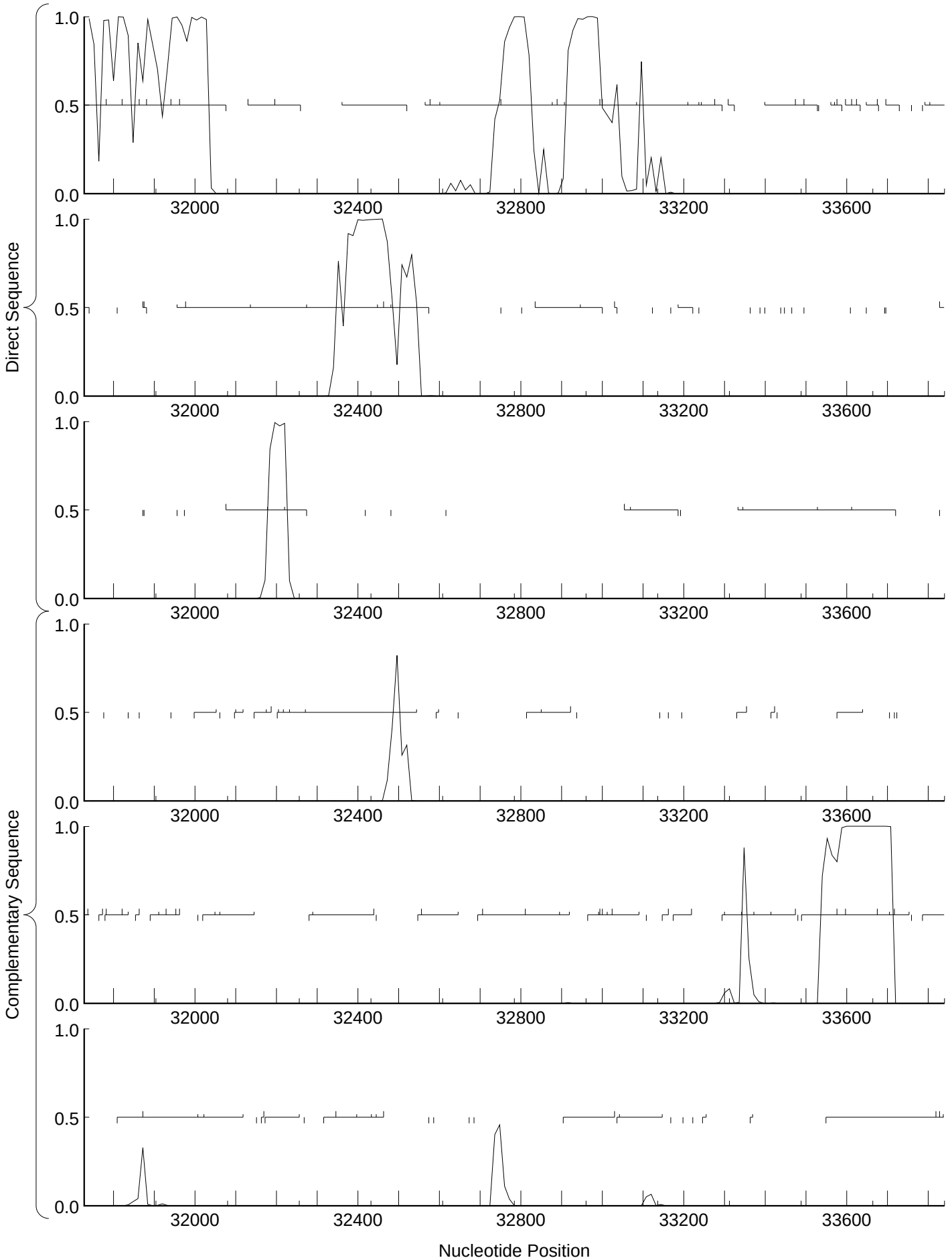

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Complementary Sequence
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23200
Nucleotide Position

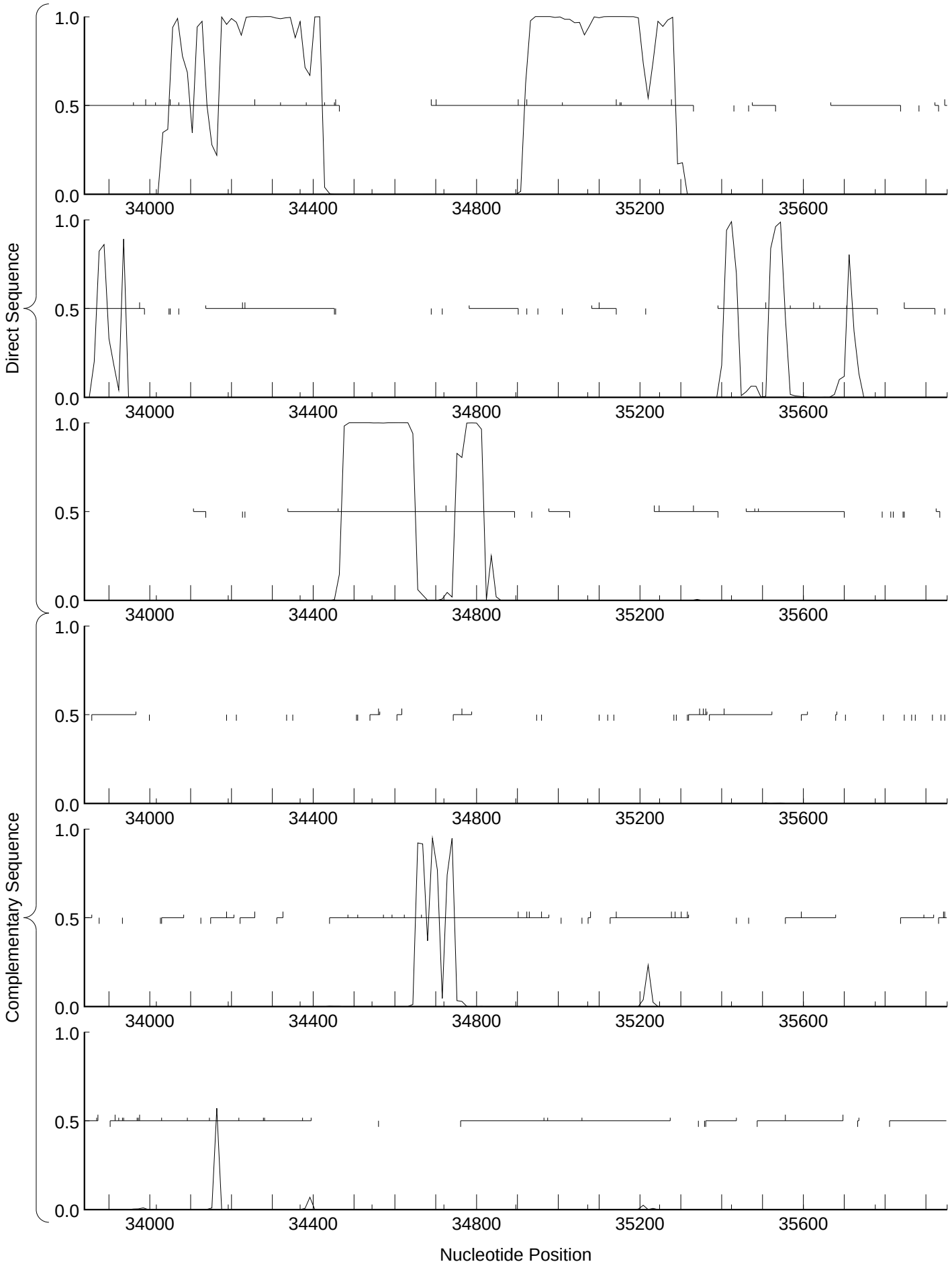

Direct Sequence
Complementary Sequence
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23600
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24800
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Nucleotide Position

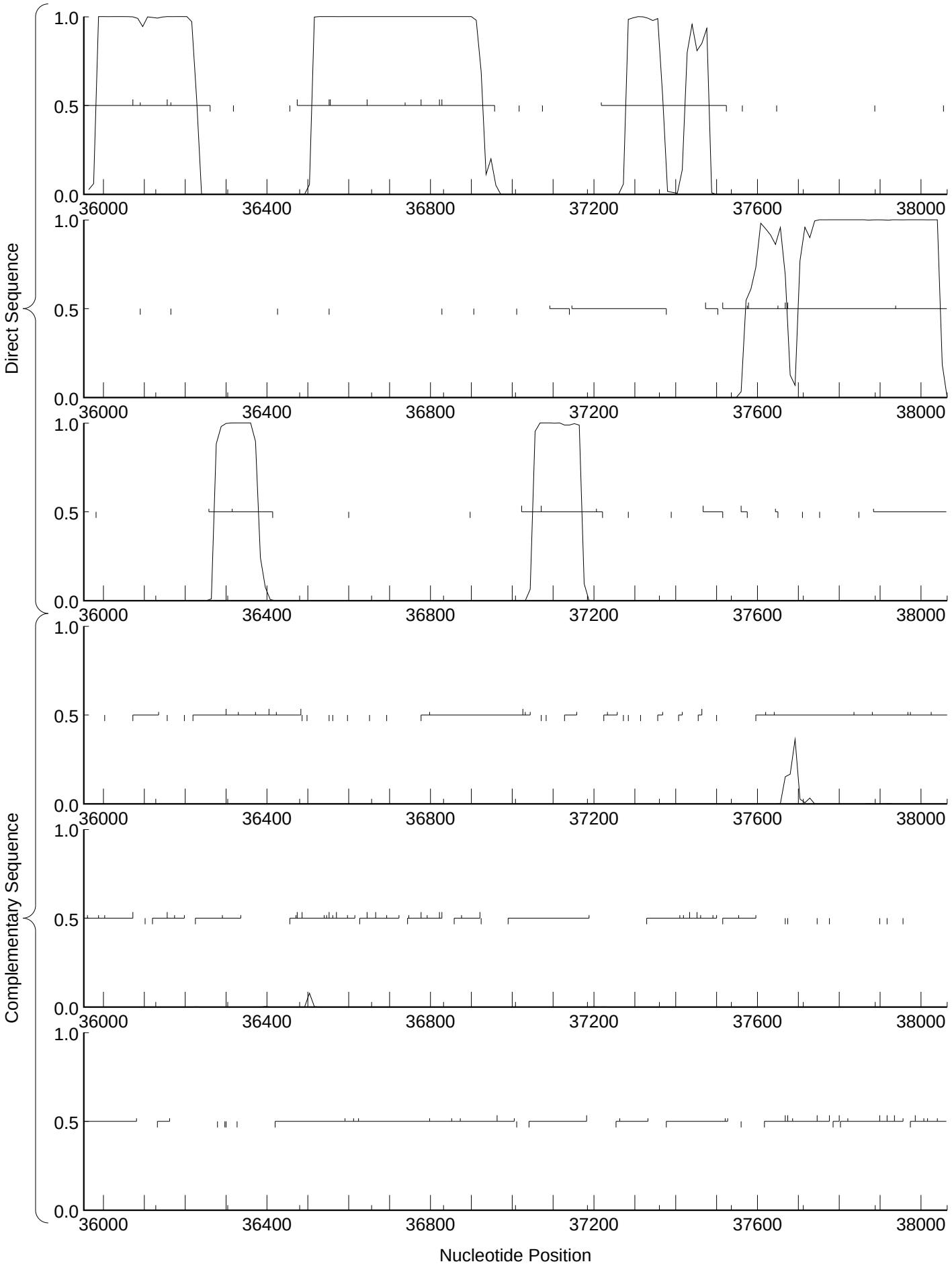

Direct Sequence
Complementary Sequence
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0.0
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26800
27200
Nucleotide Position

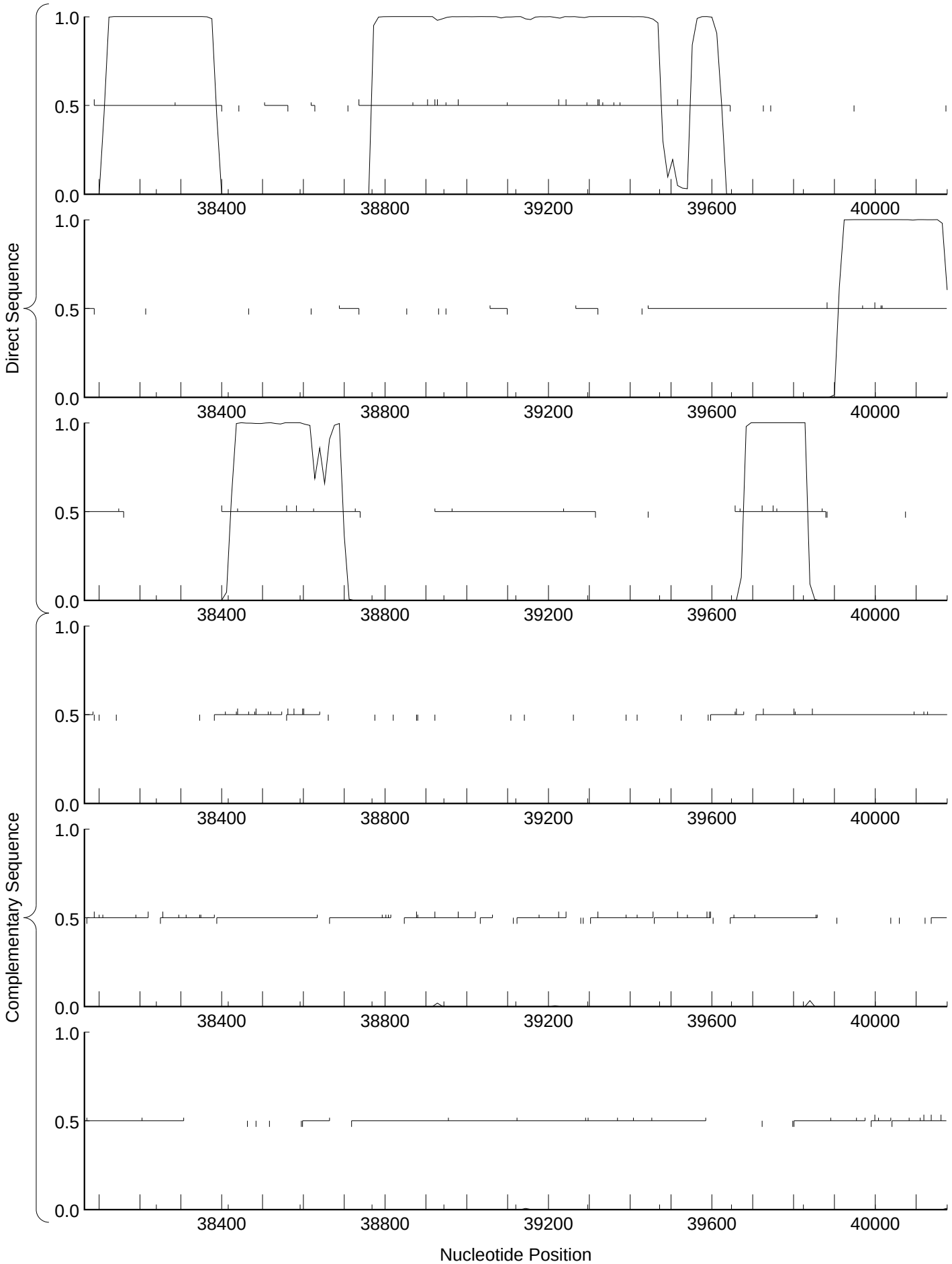

Direct Sequence
Complementary Sequence
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0.0
27600
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28400
28800
29200
29600
Nucleotide Position

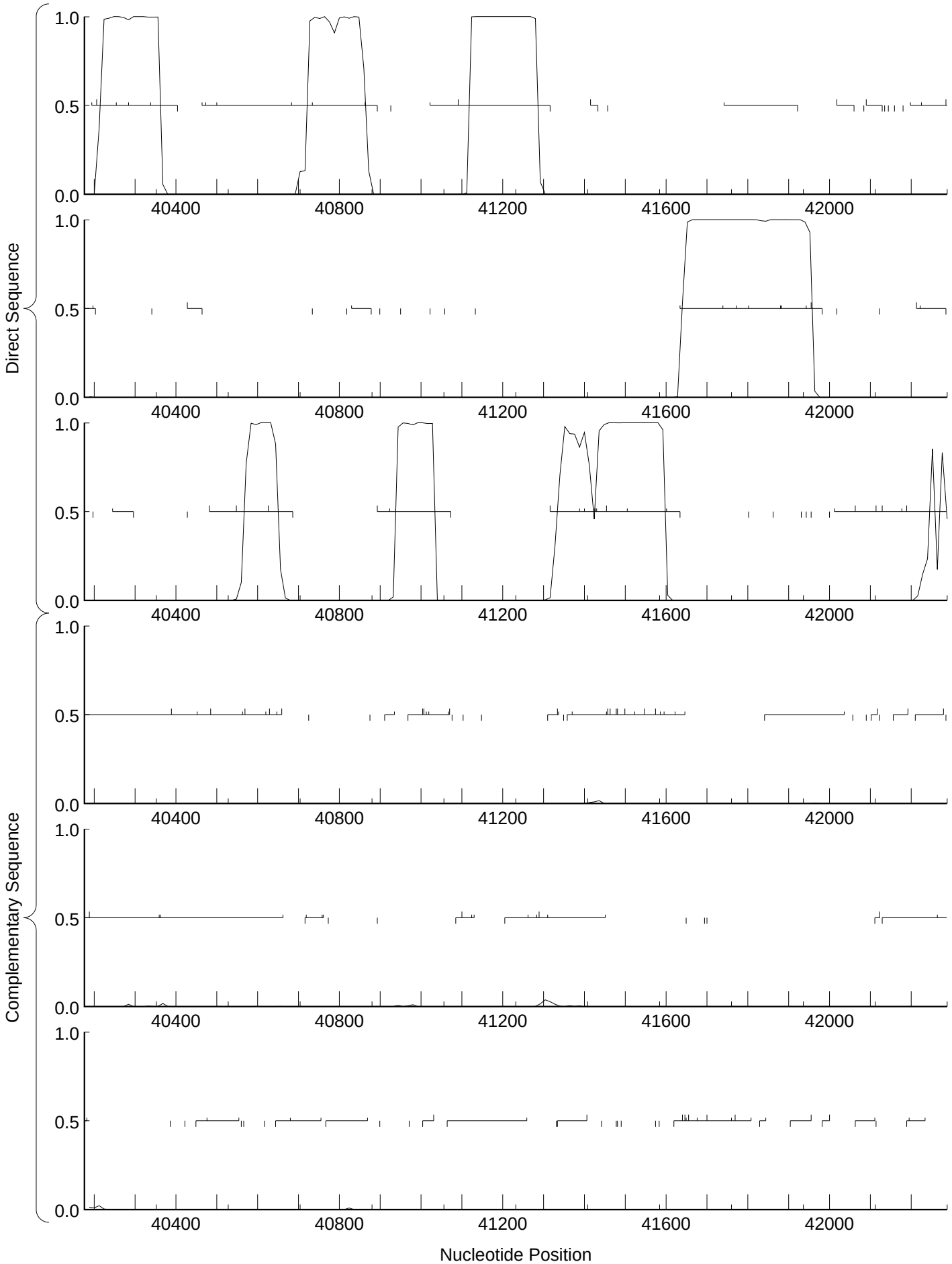

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Nucleotide Position

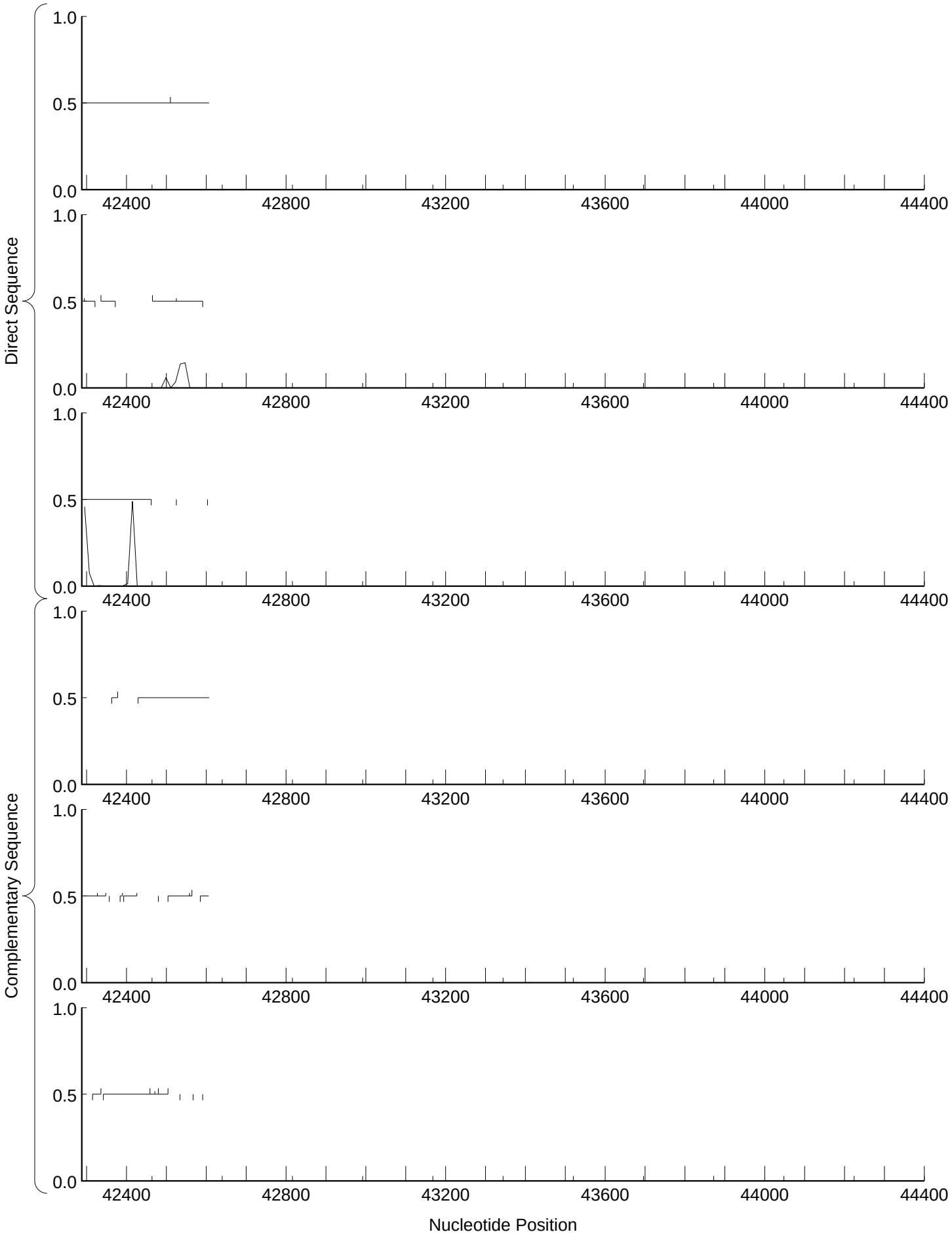

Direct Sequence
Complementary Sequence
1.0
0.5
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32000
32400
32800
33200
33600
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
36000
36400
36800
37200
37600
38000
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
40400
40800
41200
41600
42000
Nucleotide Position


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