GeneMark

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

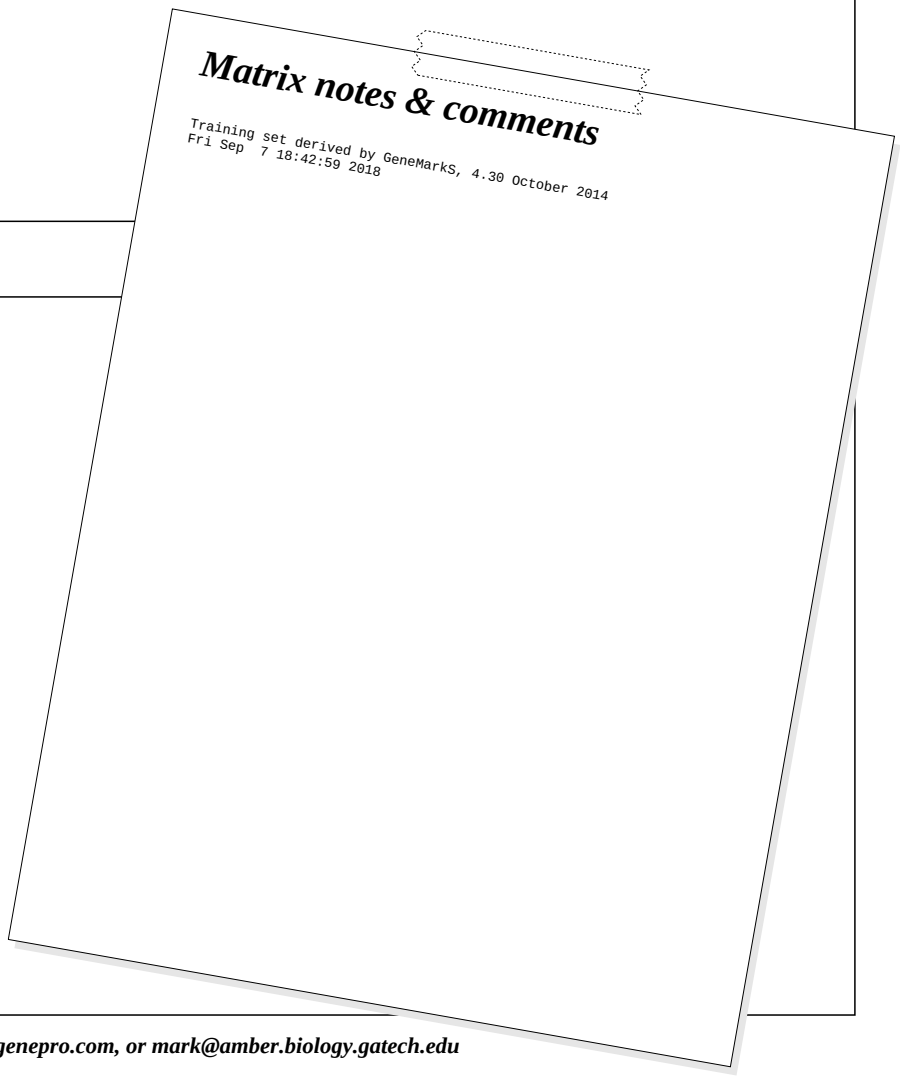
PROGRAM INFORMATION

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Sequence         :  Microbacterium phage OneinaGillian complete sequence, 61703 bp including 203 bp terminal re
Analysis Date    :  3/28/24 at 8:47:30
Pages            :  31
Sequence Length  :  61703 bp
GC Content       :   67.07%

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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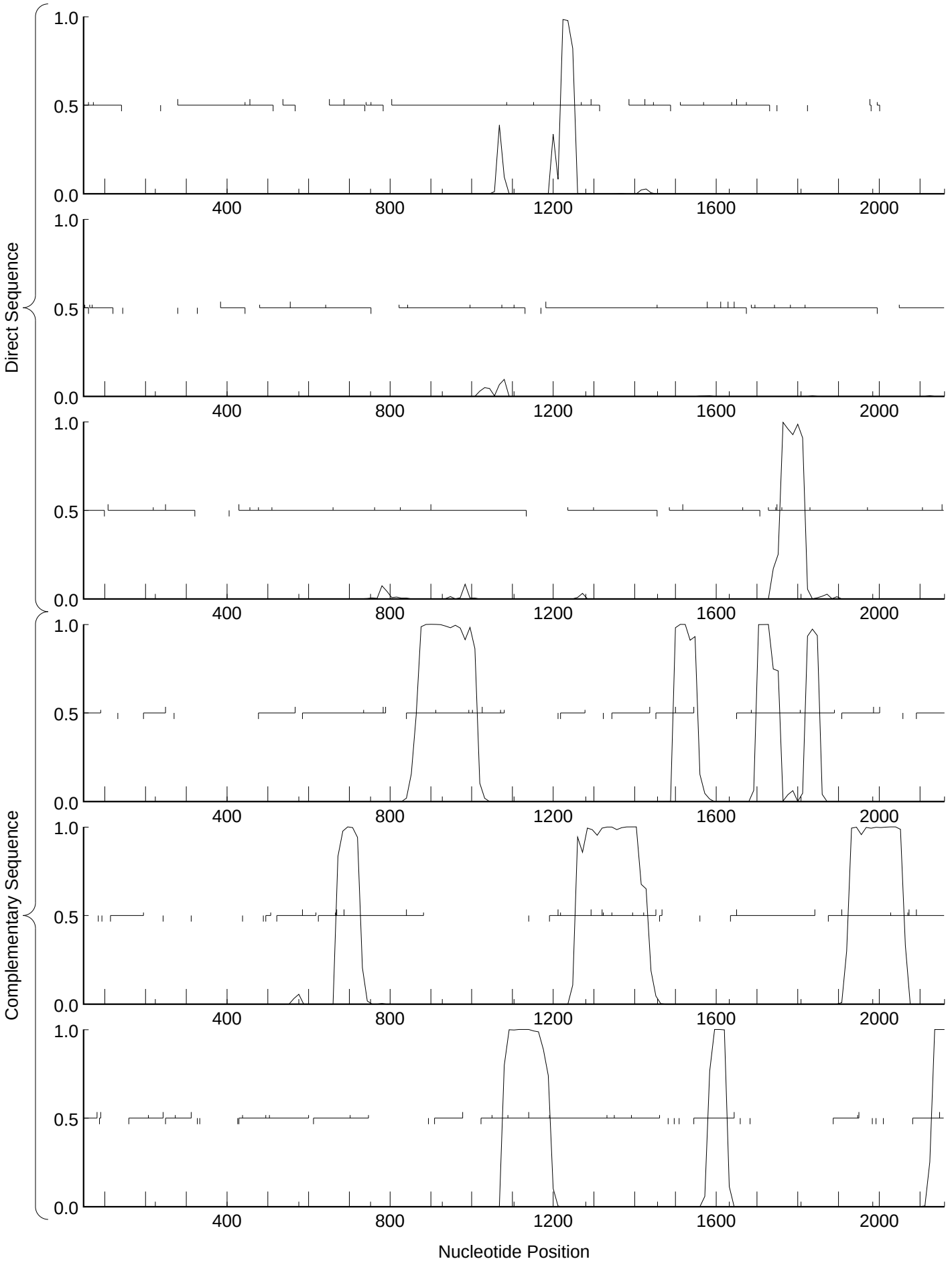
MATRIX INFORMATION

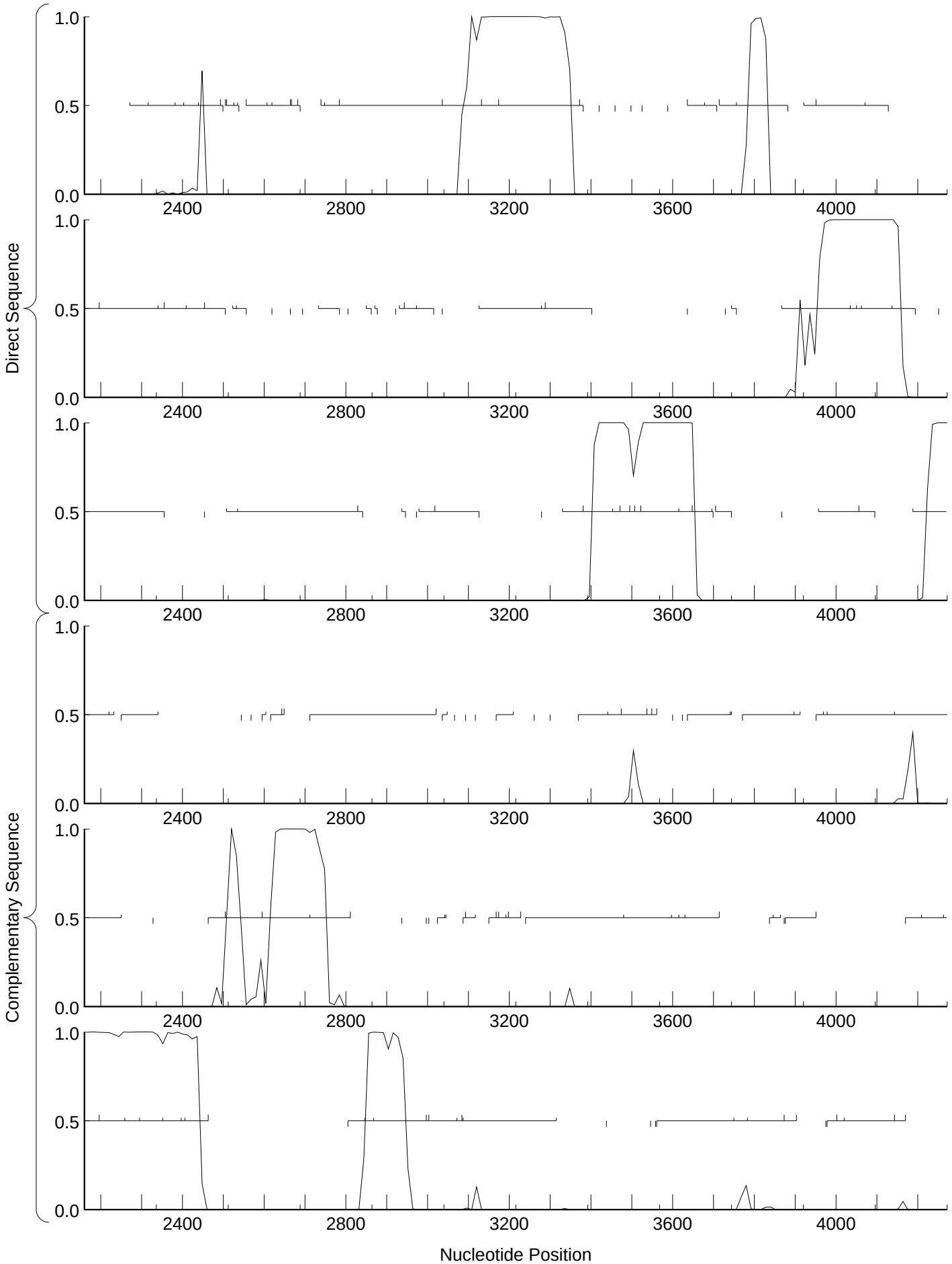
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Matrix     :  -
Author     :  -

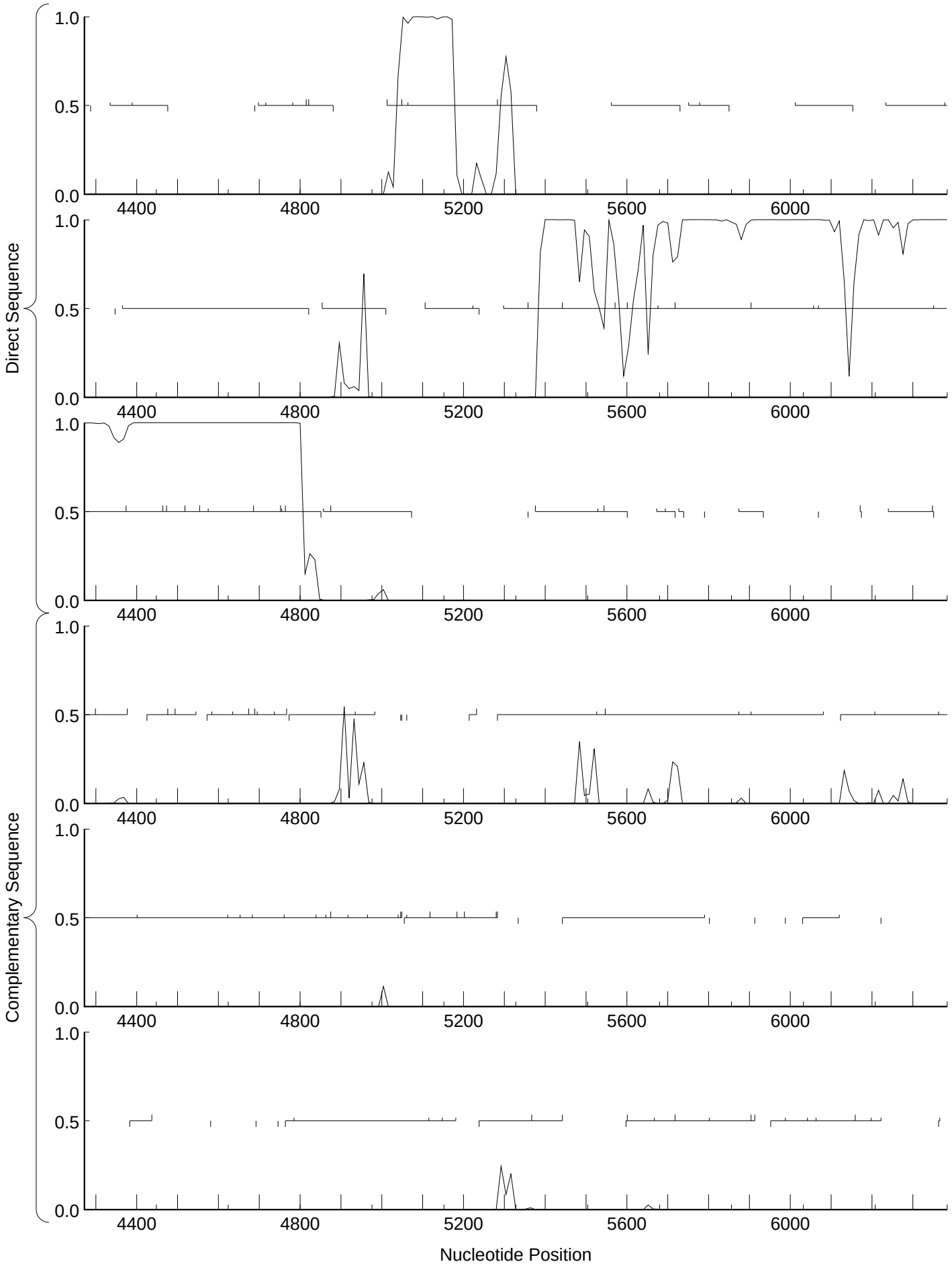
Order      :  4
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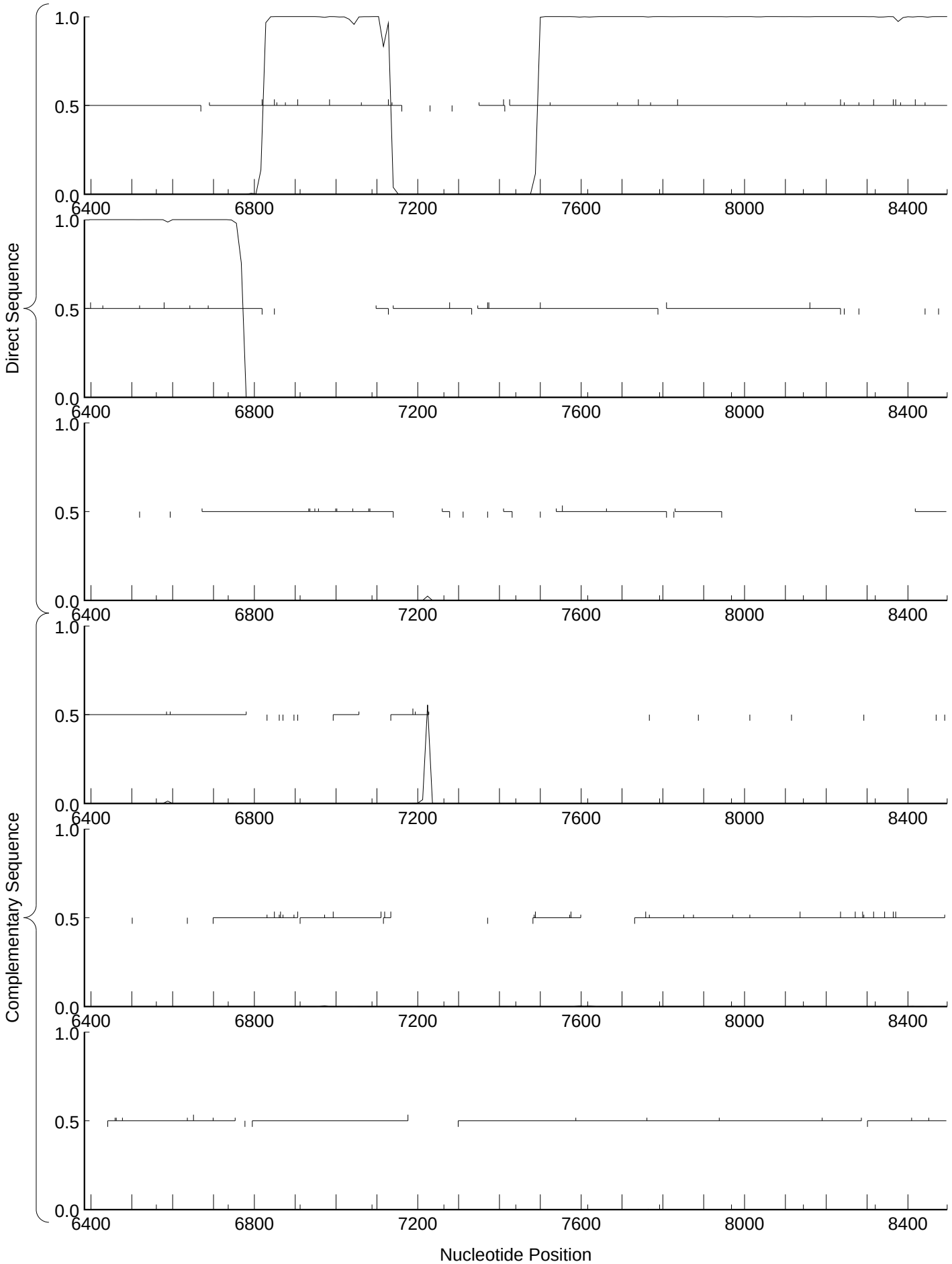

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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

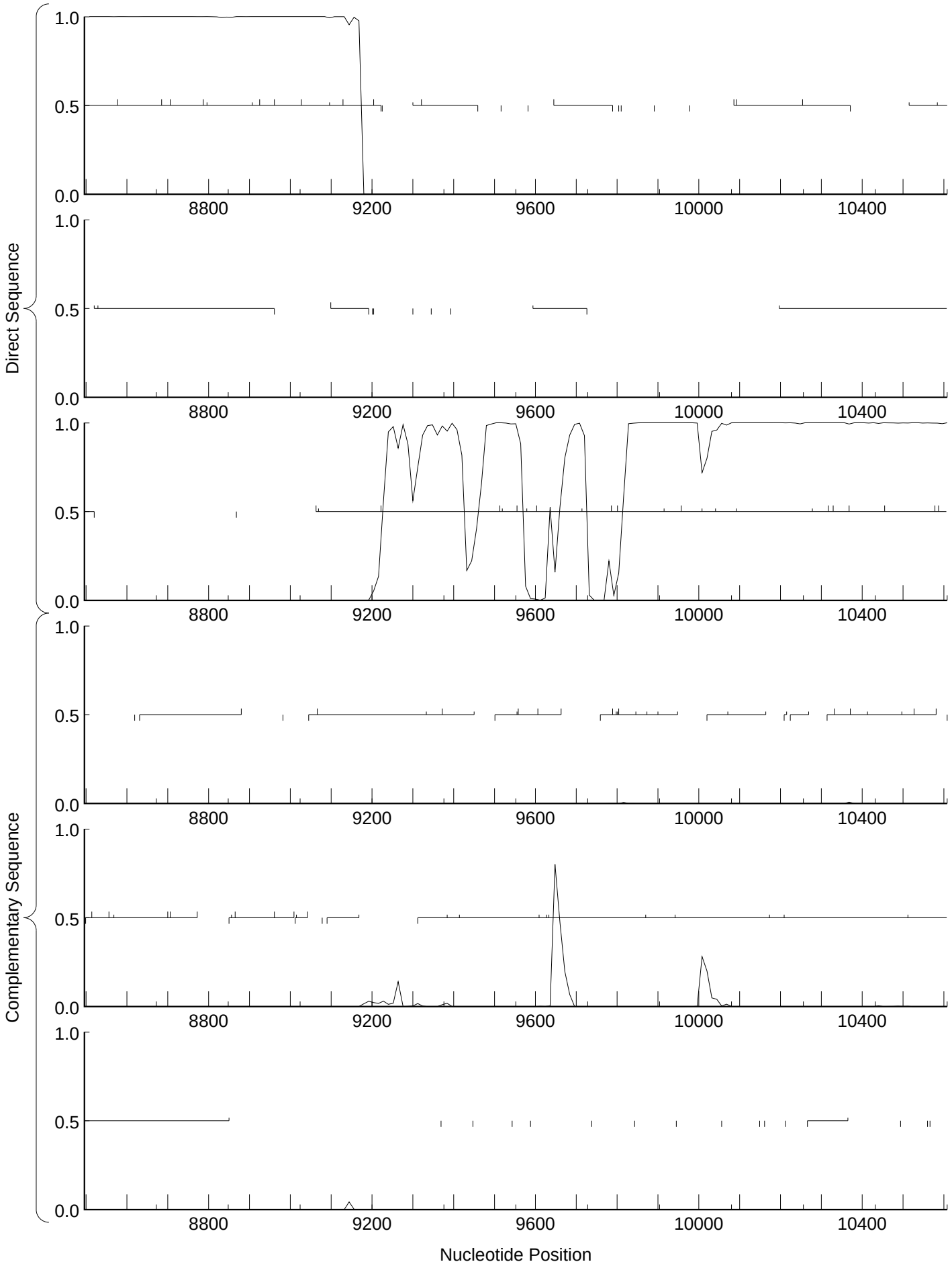

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

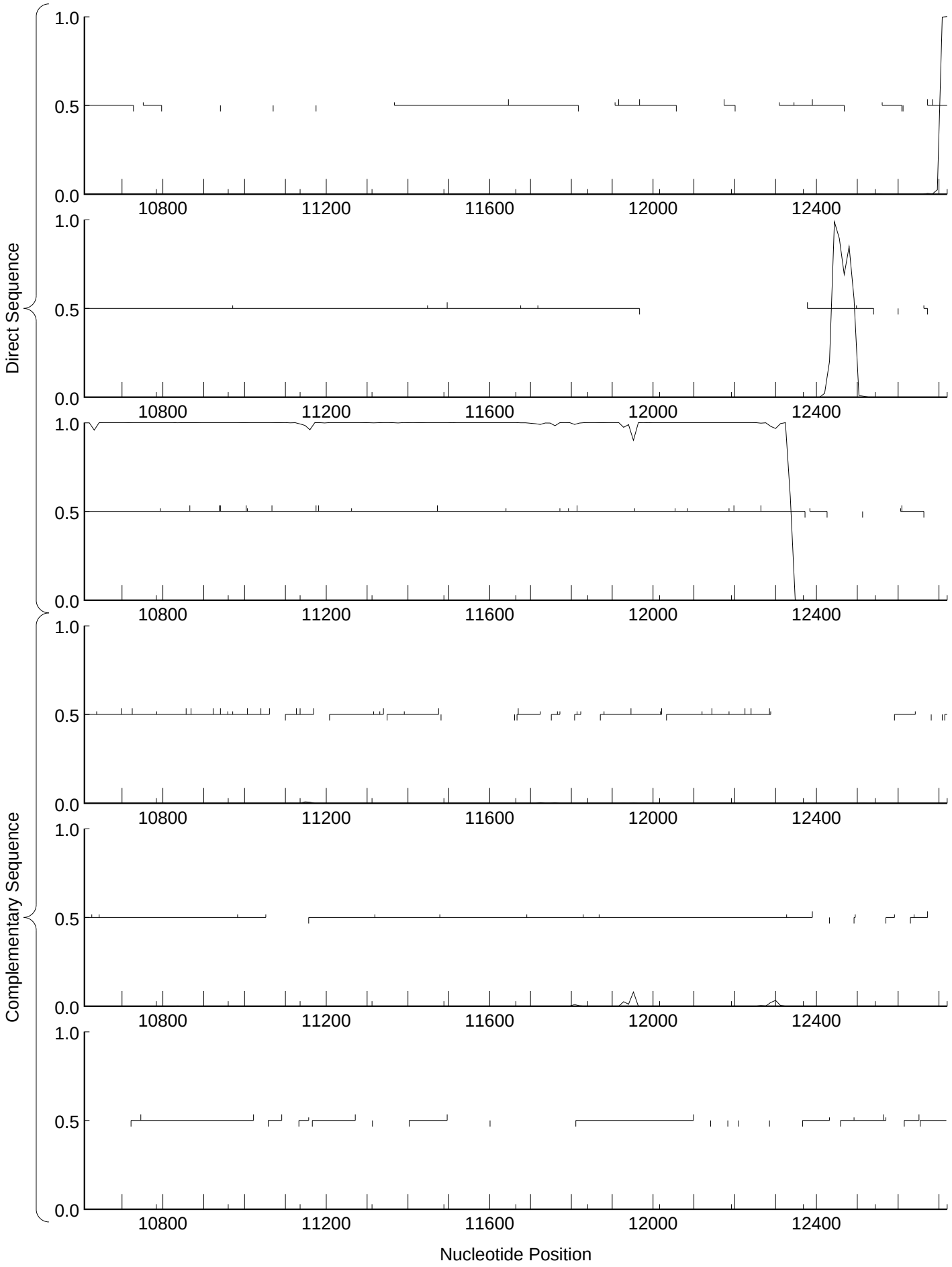

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Complementary Sequence
1.0
0.5
0.0
400
800
1200
1600
2000
Nucleotide Position

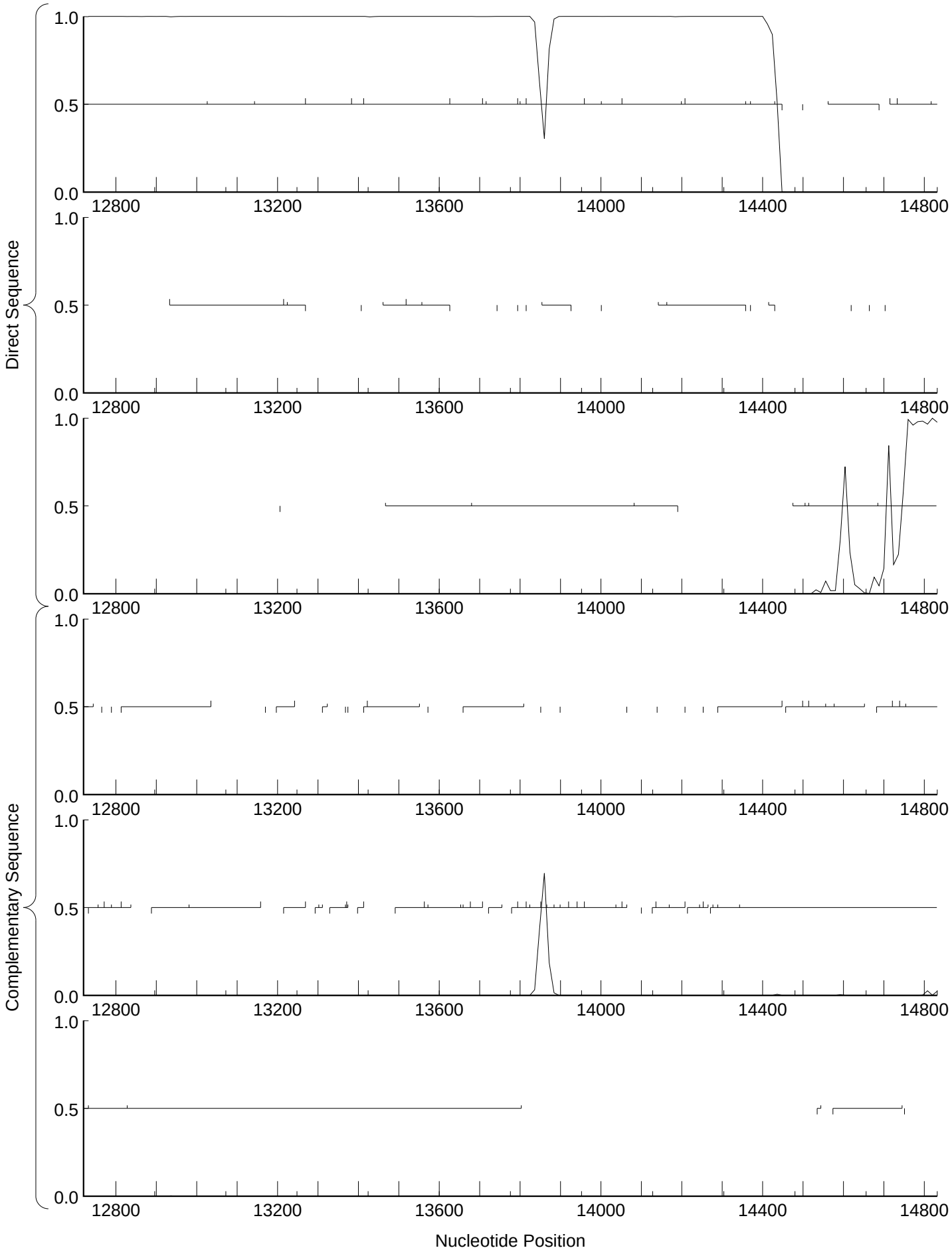

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Complementary Sequence
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0.0
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3200
3600
4000
Nucleotide Position

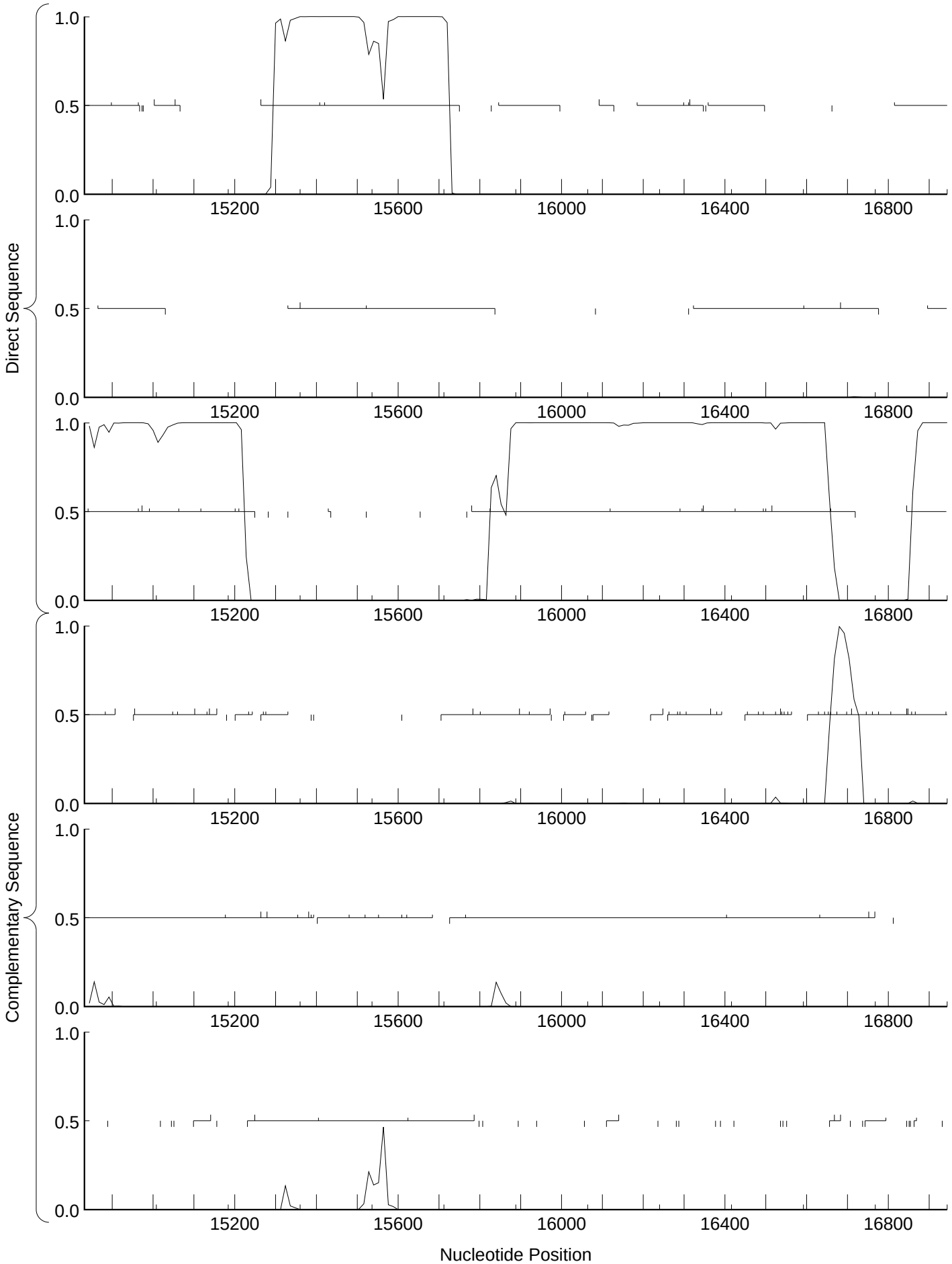

Direct Sequence
Complementary Sequence
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0.5
0.0
4400
4800
5200
5600
6000
Nucleotide Position


Direct Sequence
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6400
6800
7200
7600
8000
8400
Nucleotide Position

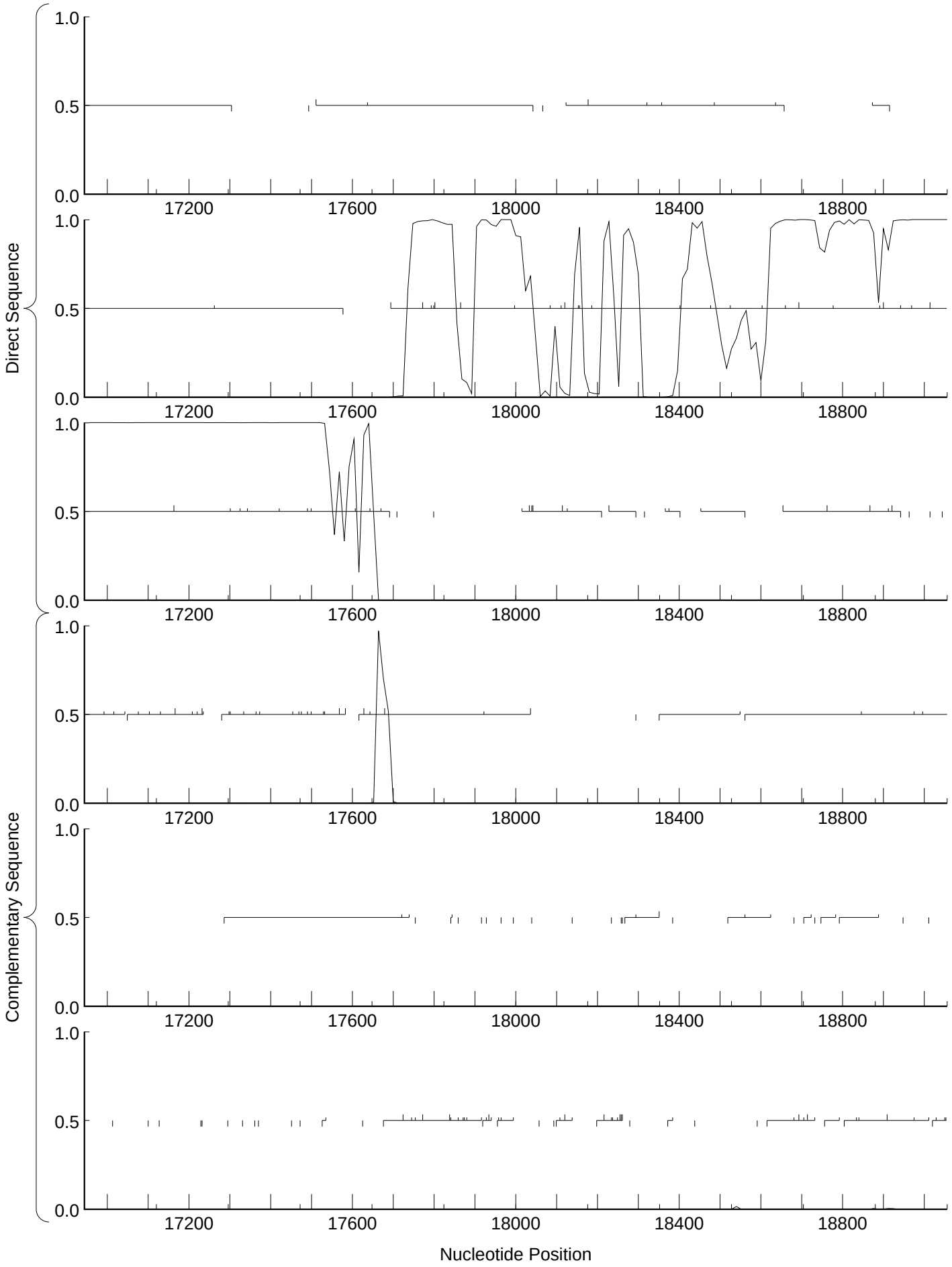

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
8800
9200
9600
10000
10400
Nucleotide Position


Direct Sequence
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10800
11200
11600
12000
12400
Nucleotide Position

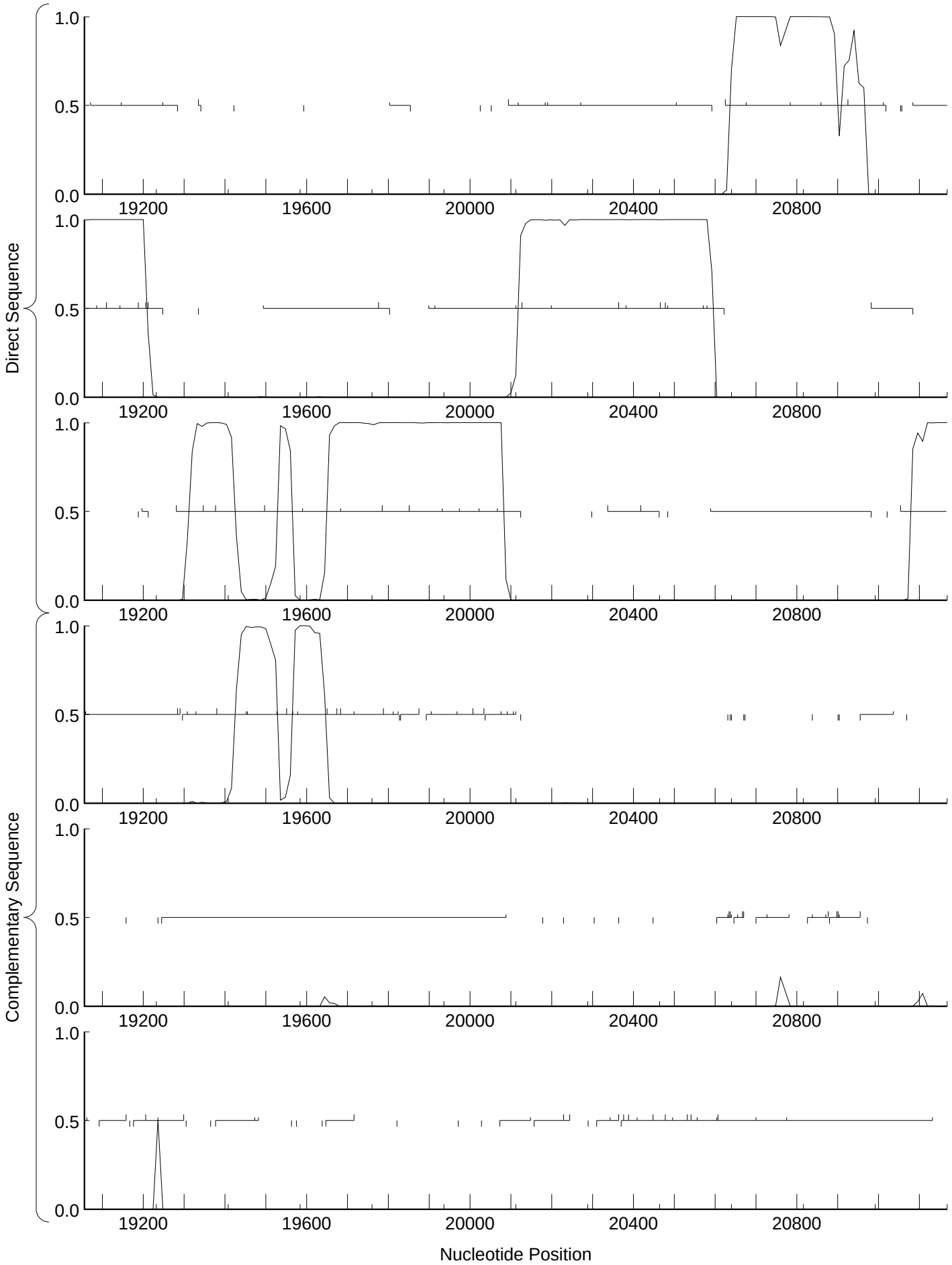

Direct Sequence
Complementary Sequence
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0.5
0.0
12800
13200
13600
14000
14400
14800
Nucleotide Position


Direct Sequence
Complementary Sequence
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0.5
0.0
15200
15600
16000
16400
16800
Nucleotide Position

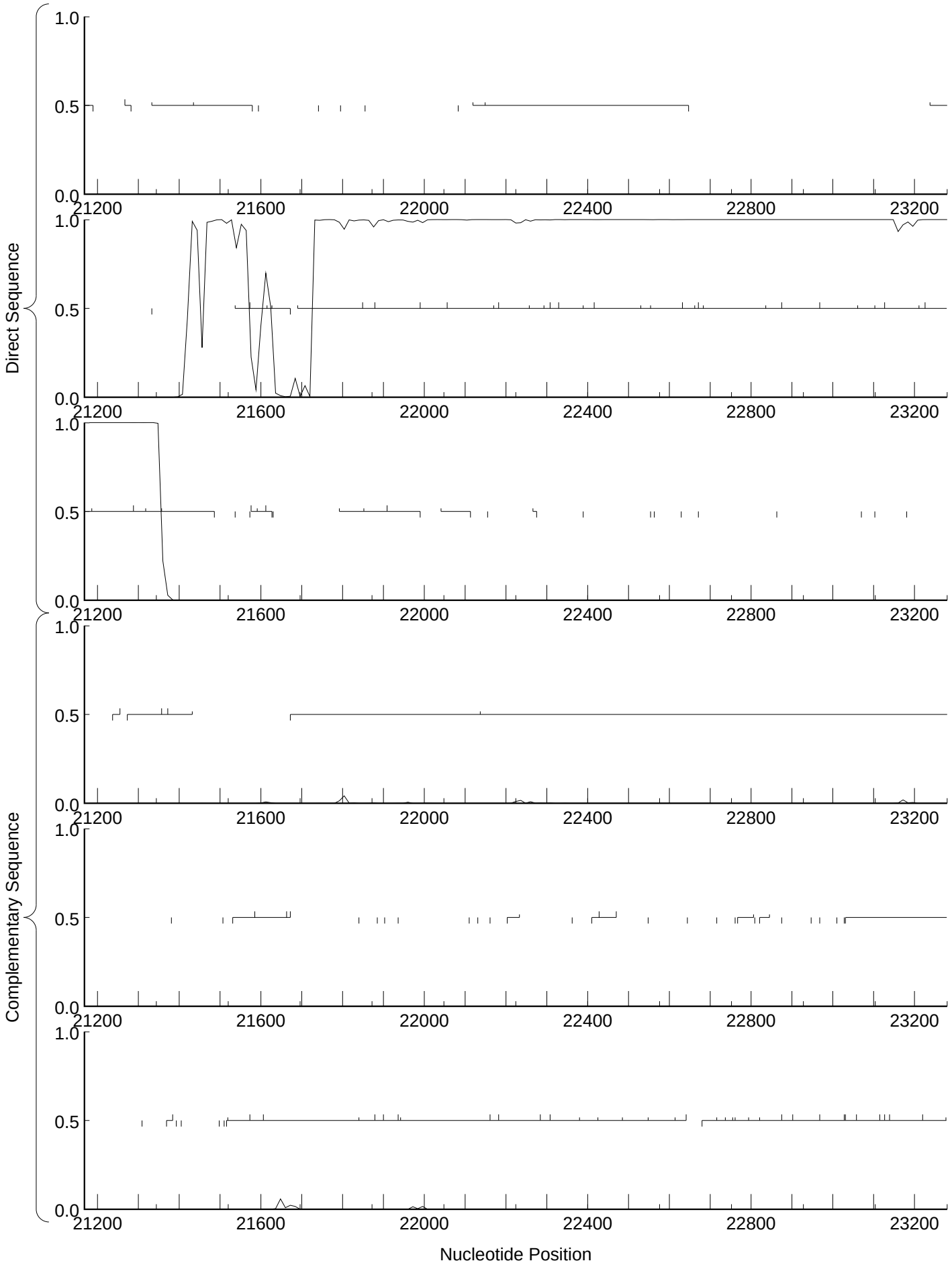
ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 10/31



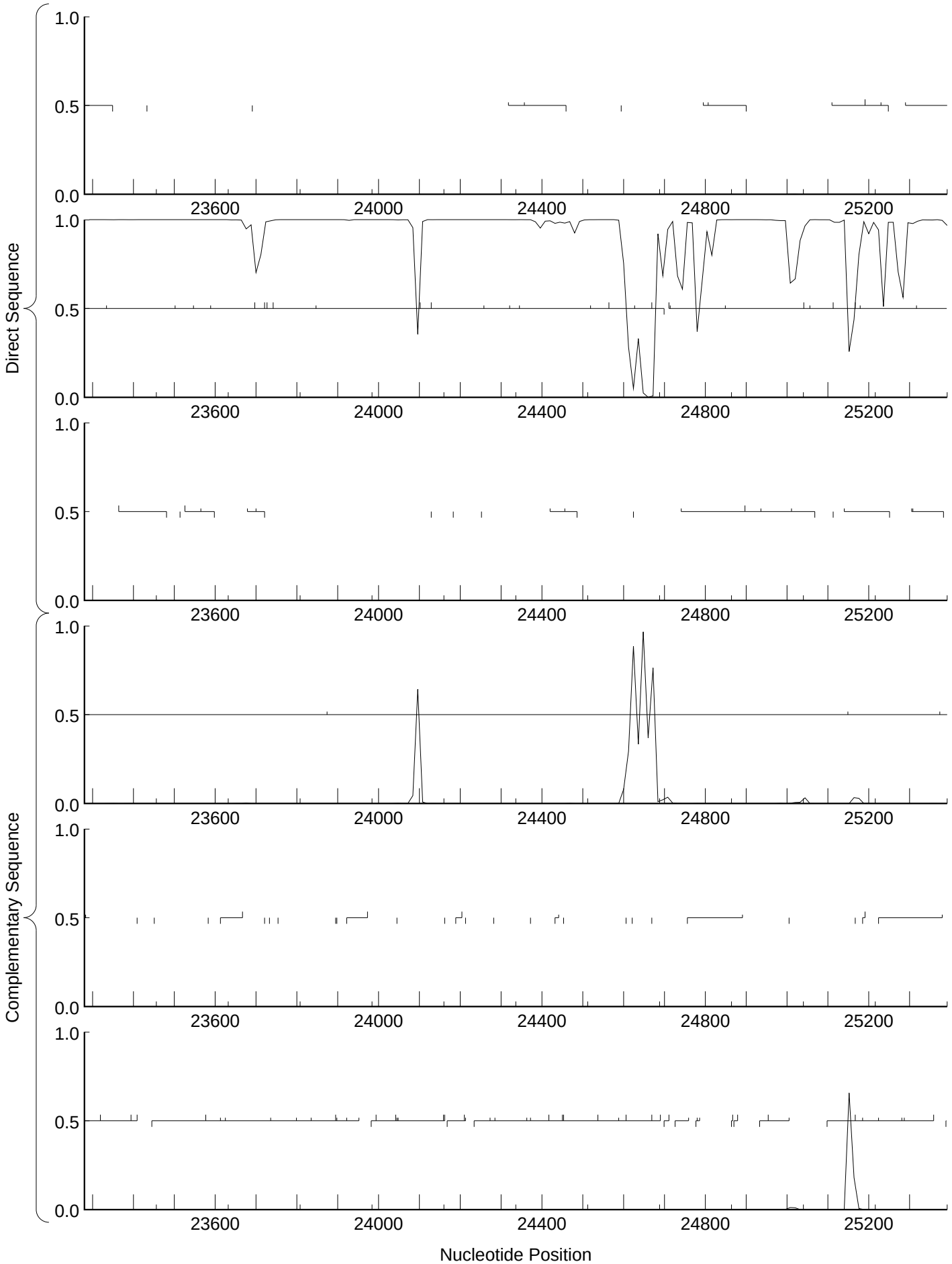
ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 11/31



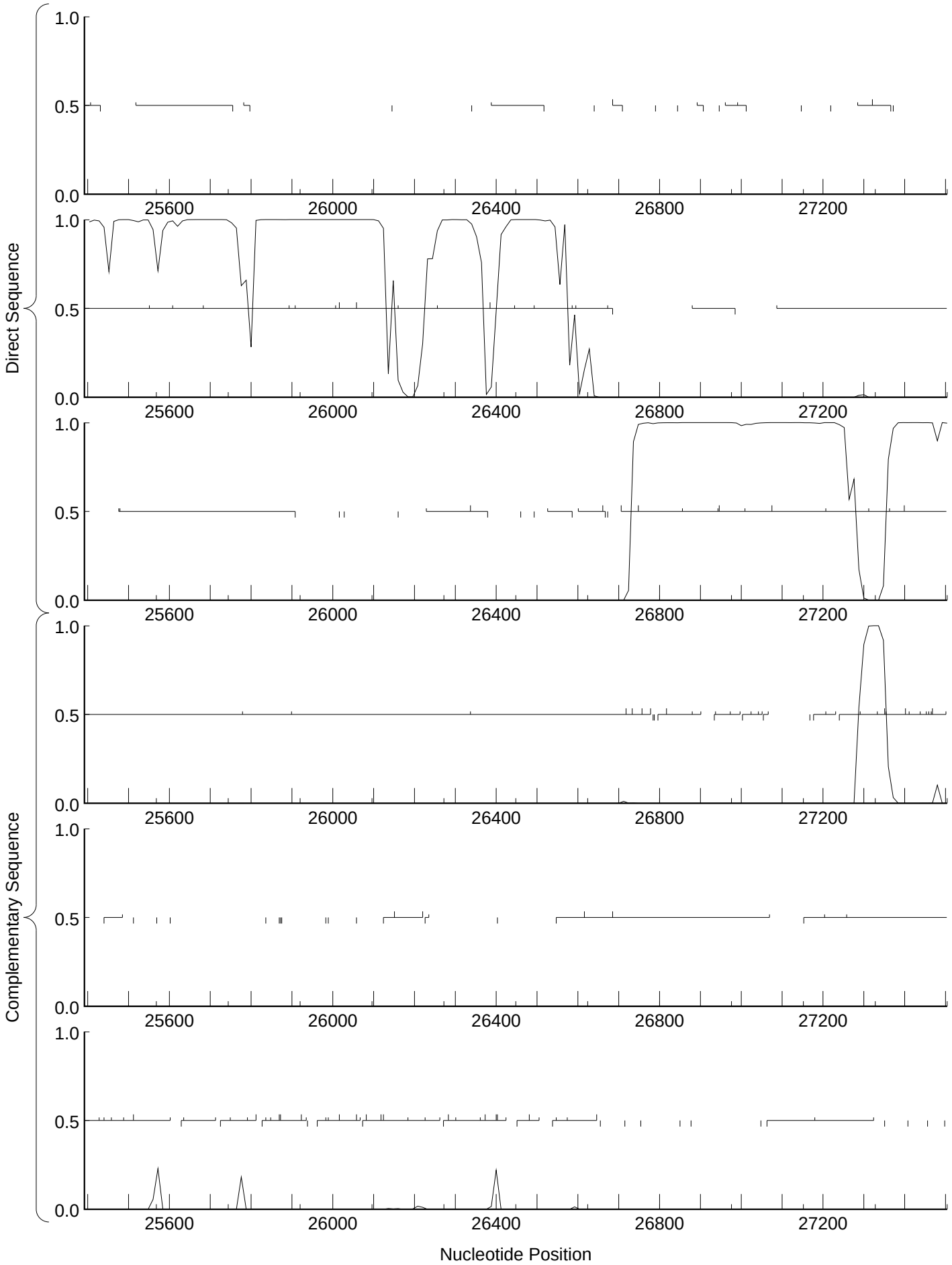
ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 12/31



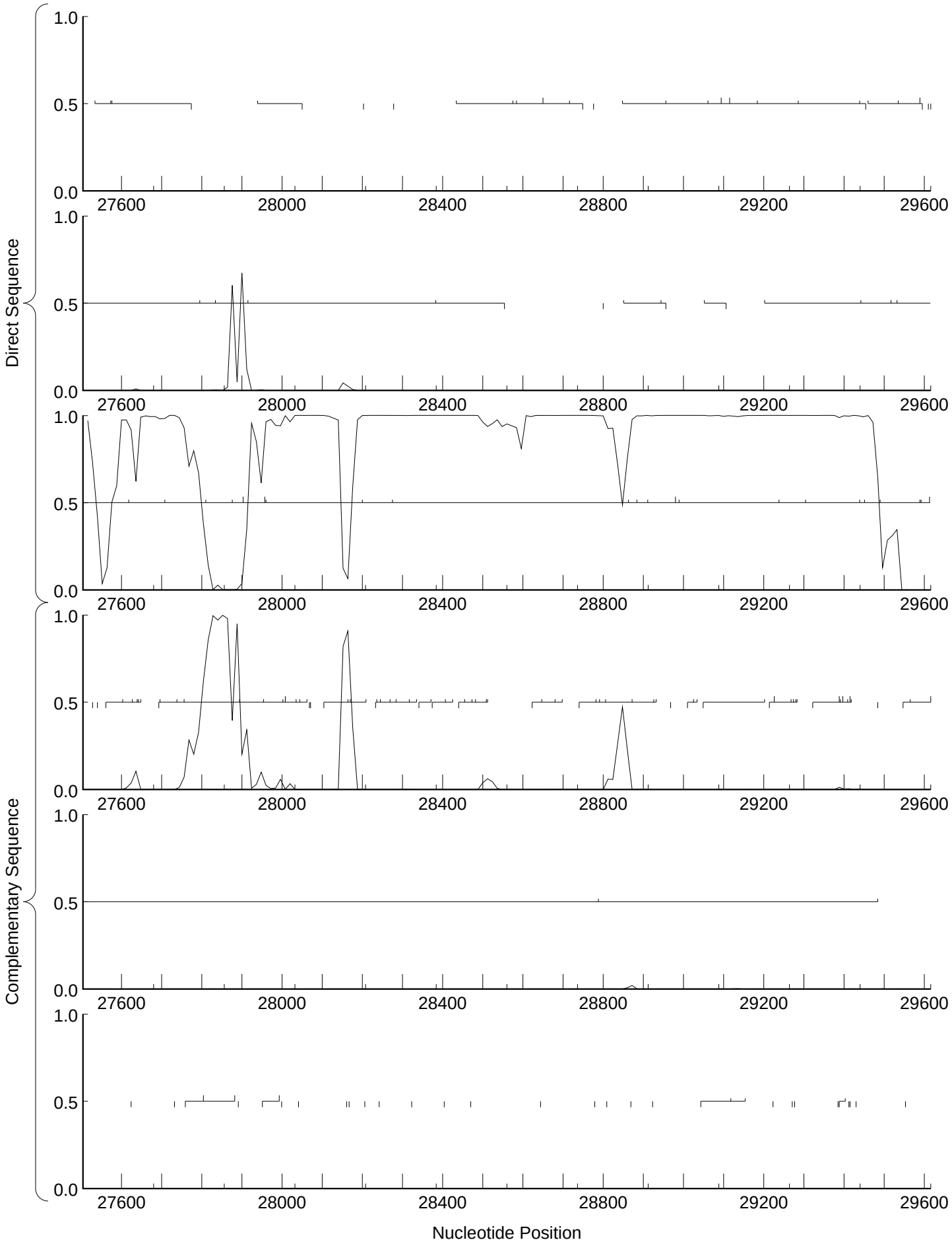
ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 13/31



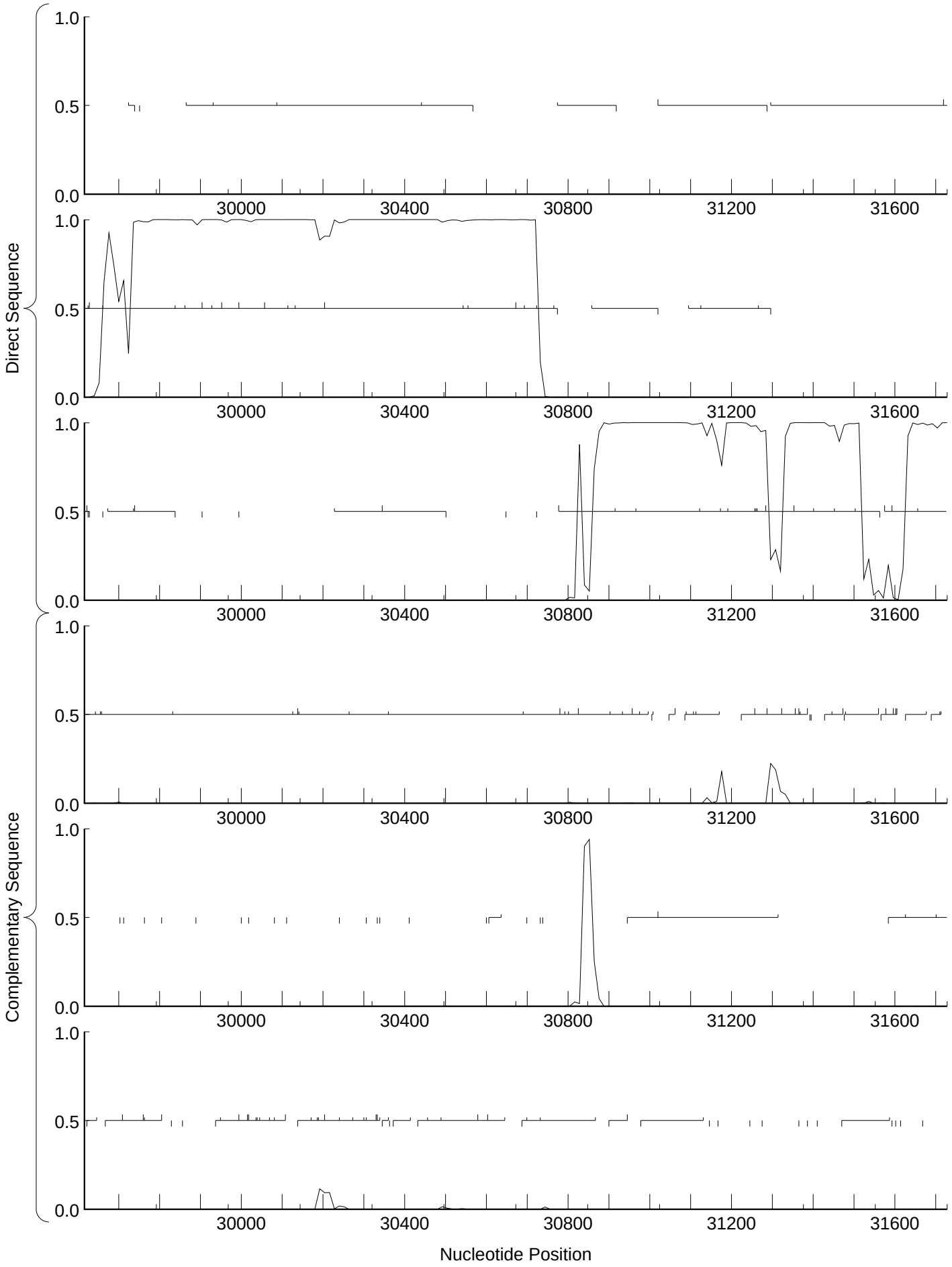
ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 14/31



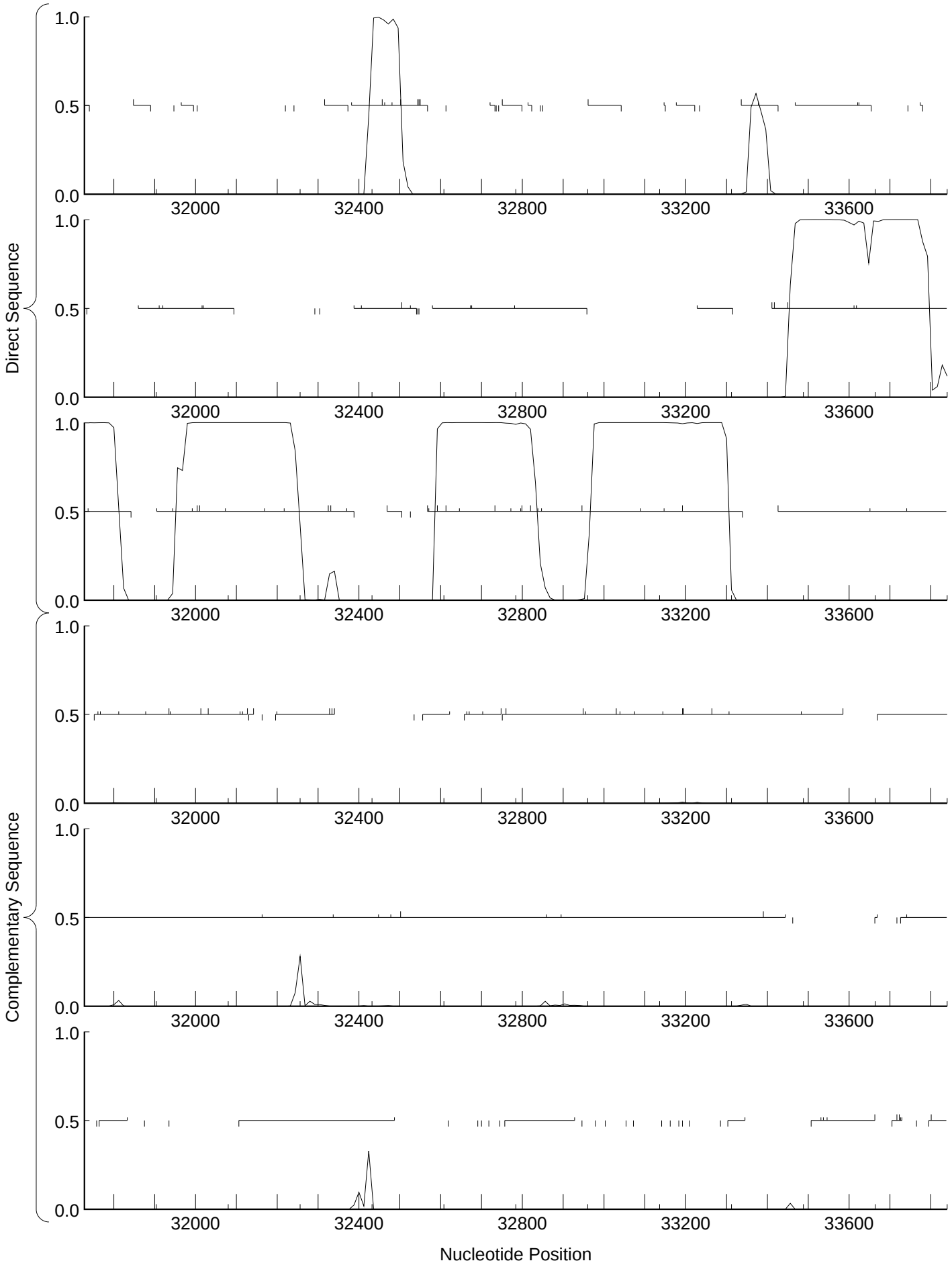
ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 15/31



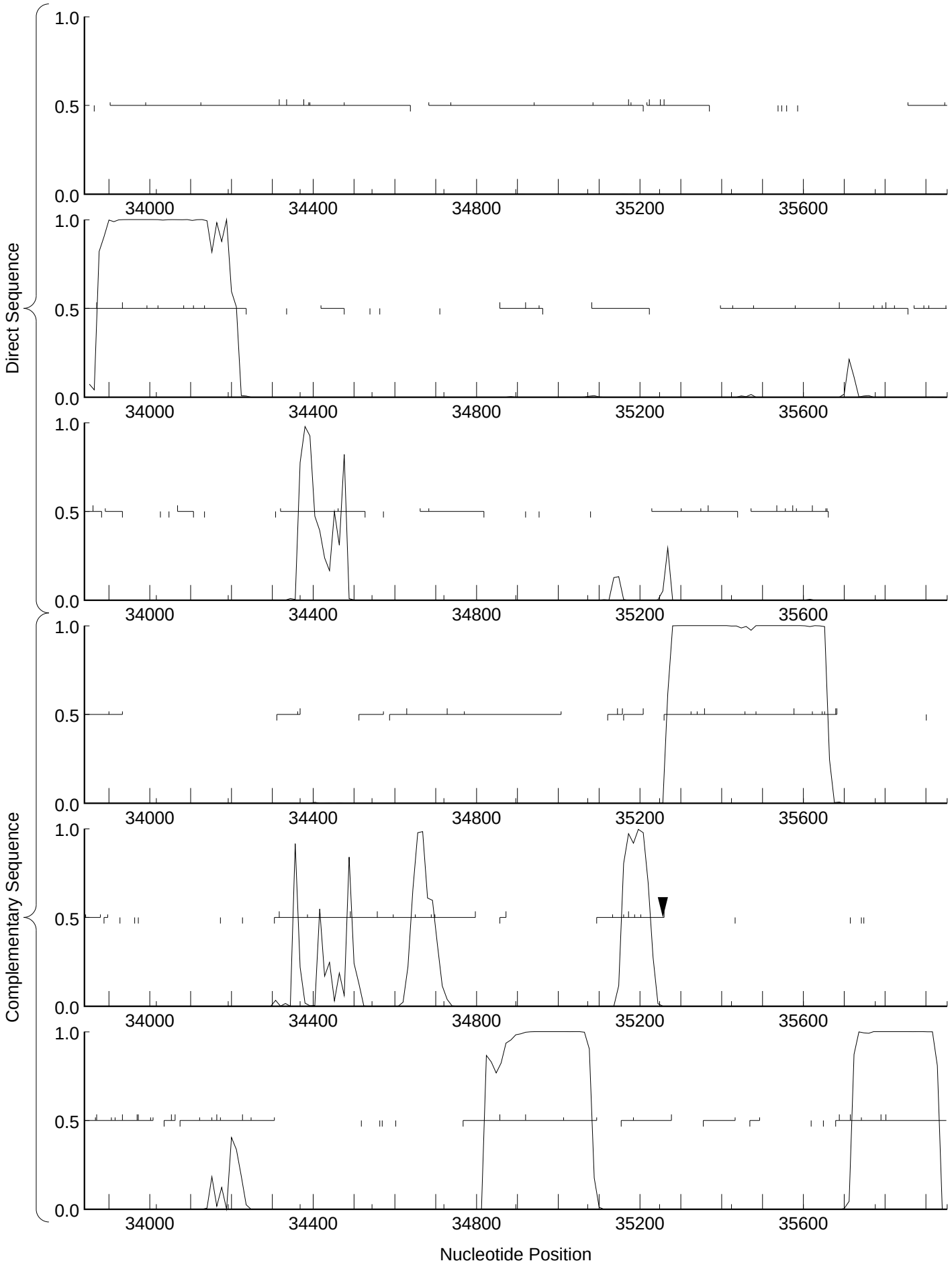
ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 16/31



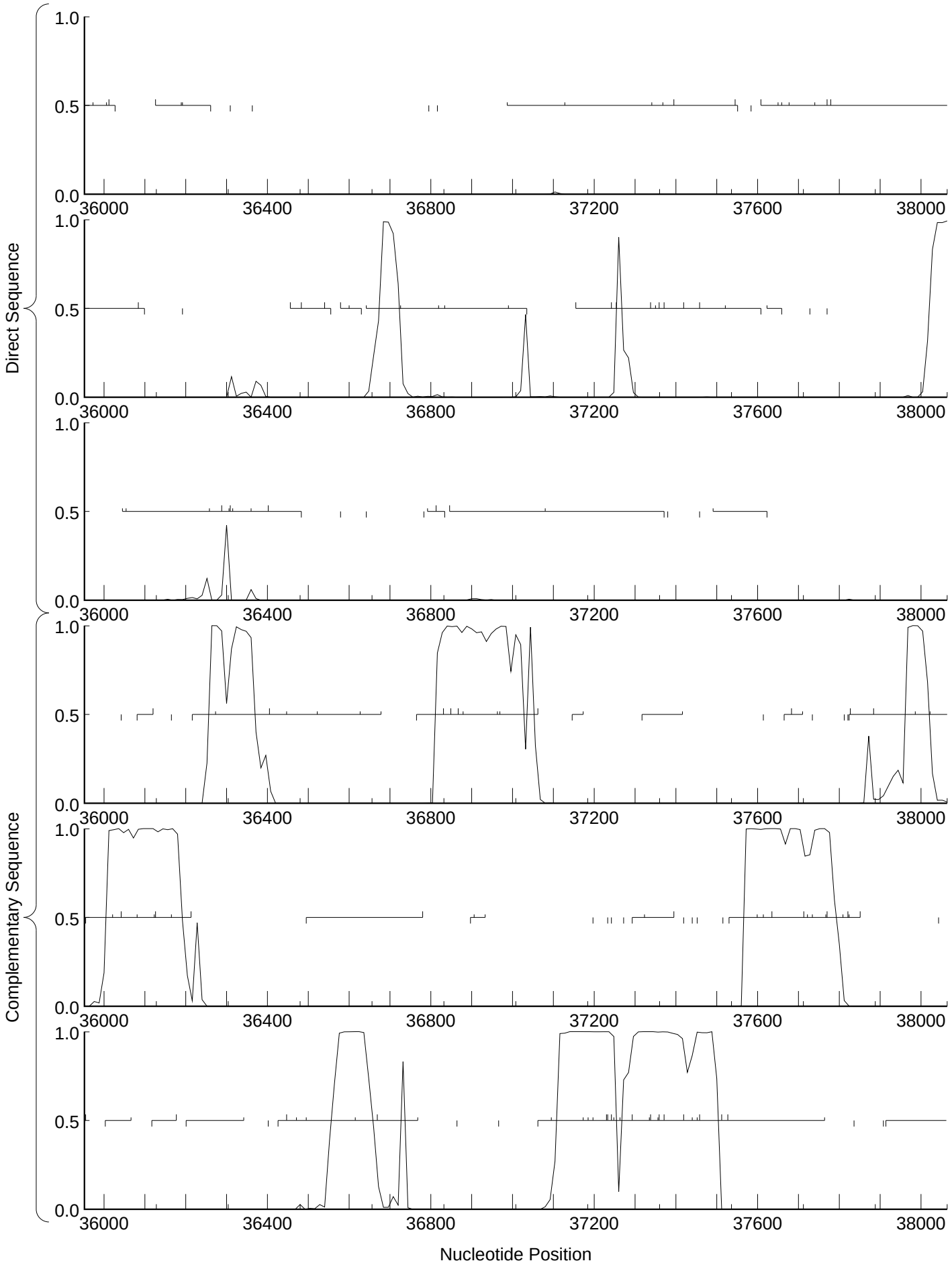
ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 17/31



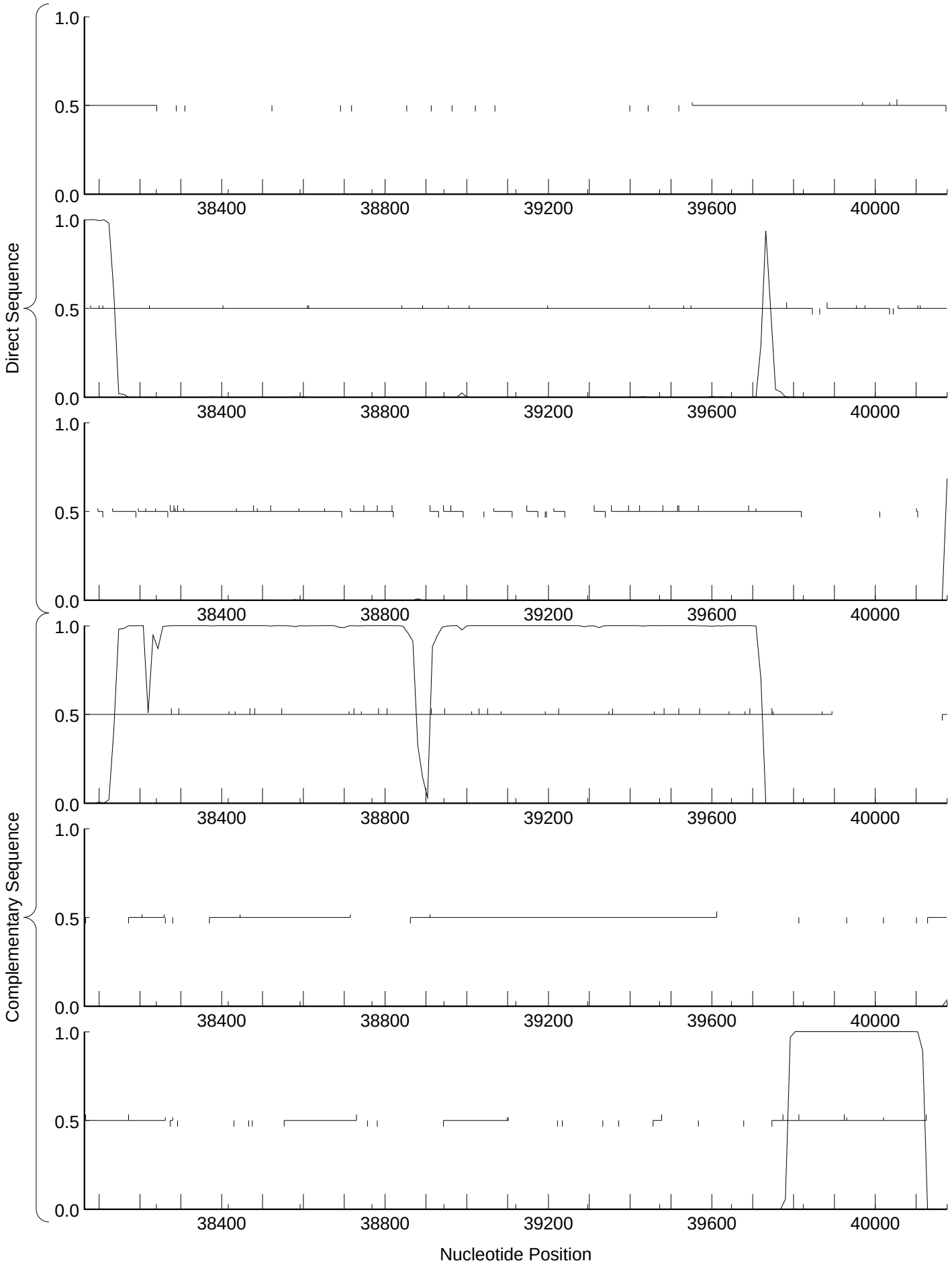
ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 18/31



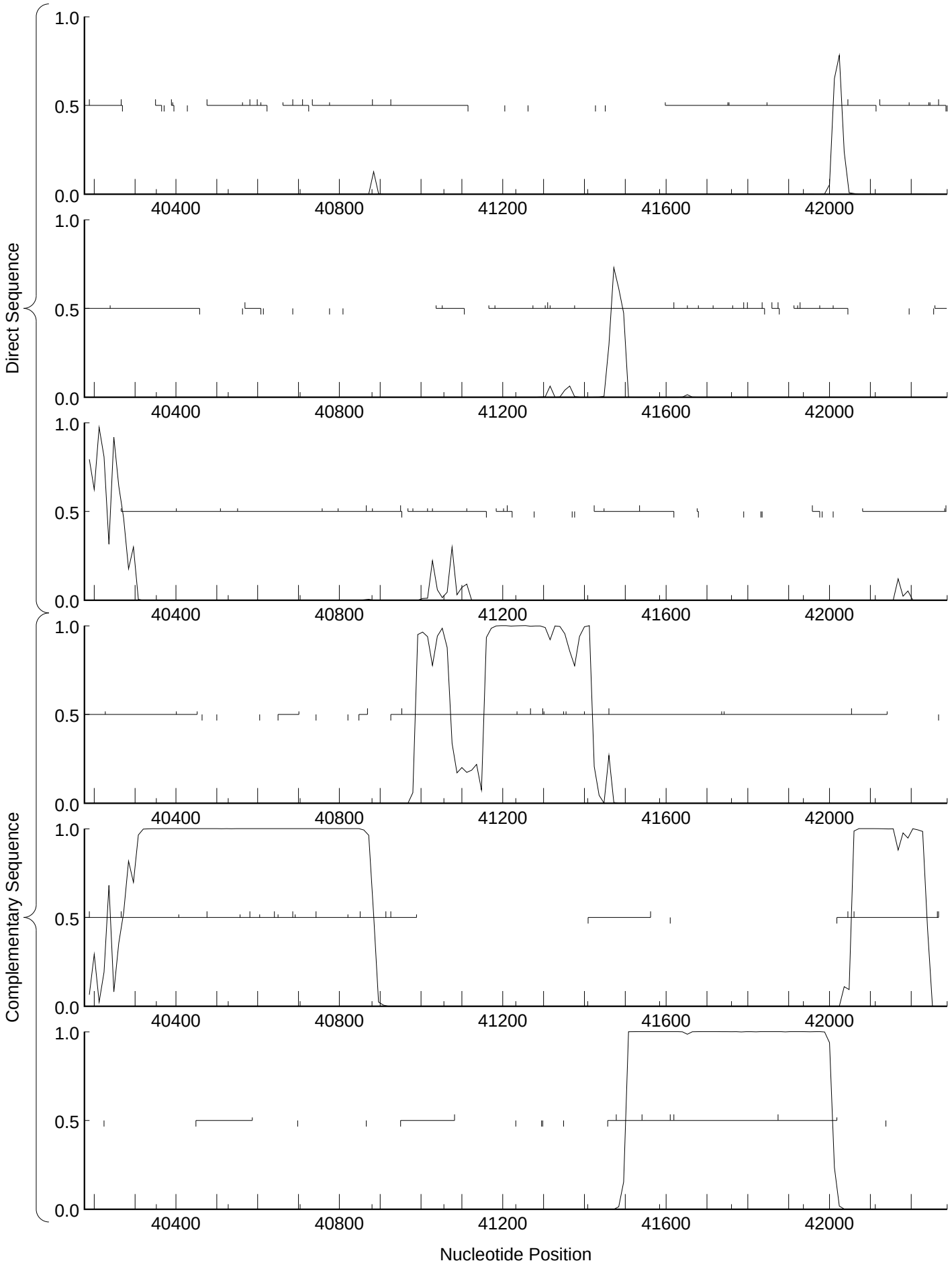
ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 19/31



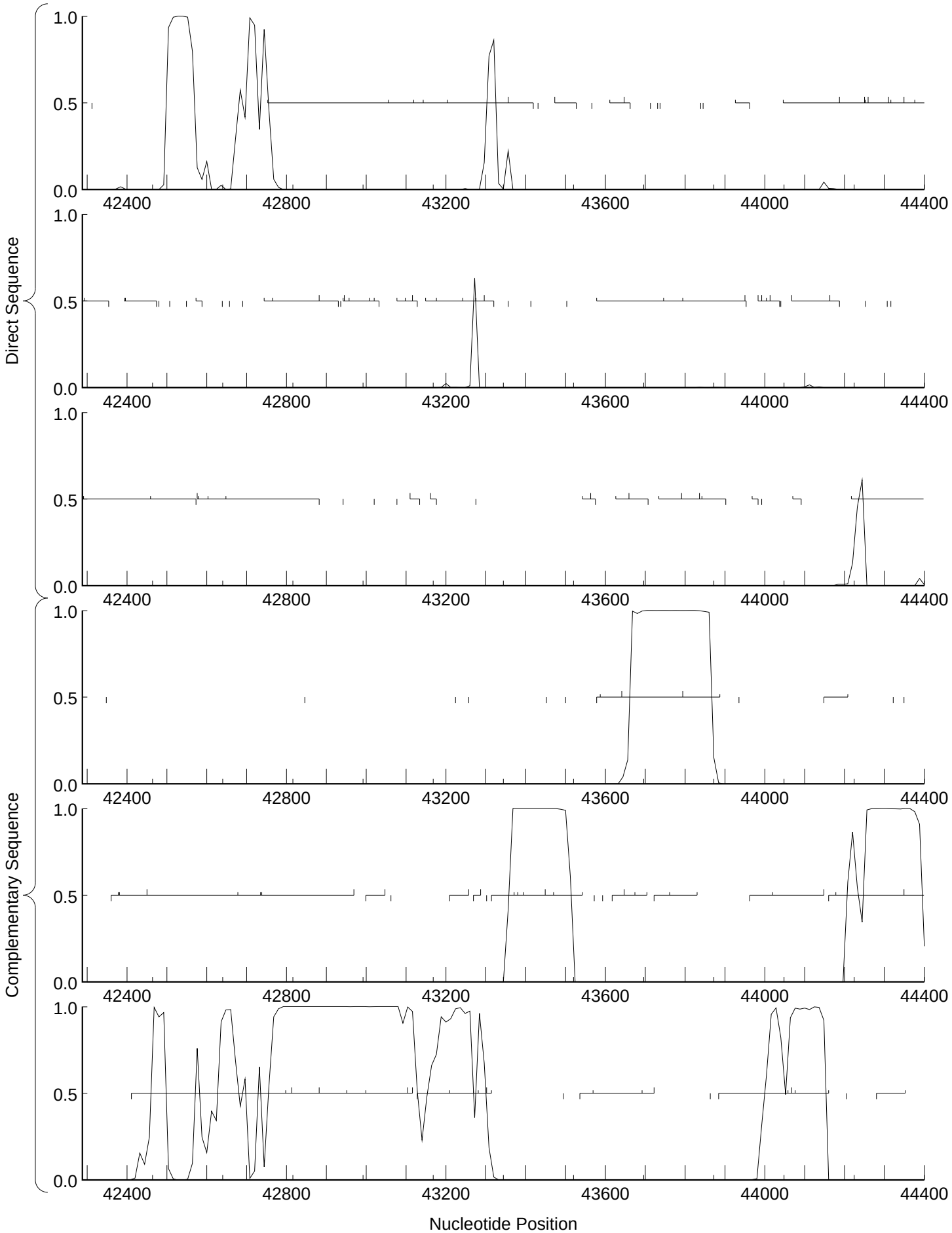
ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 20/31



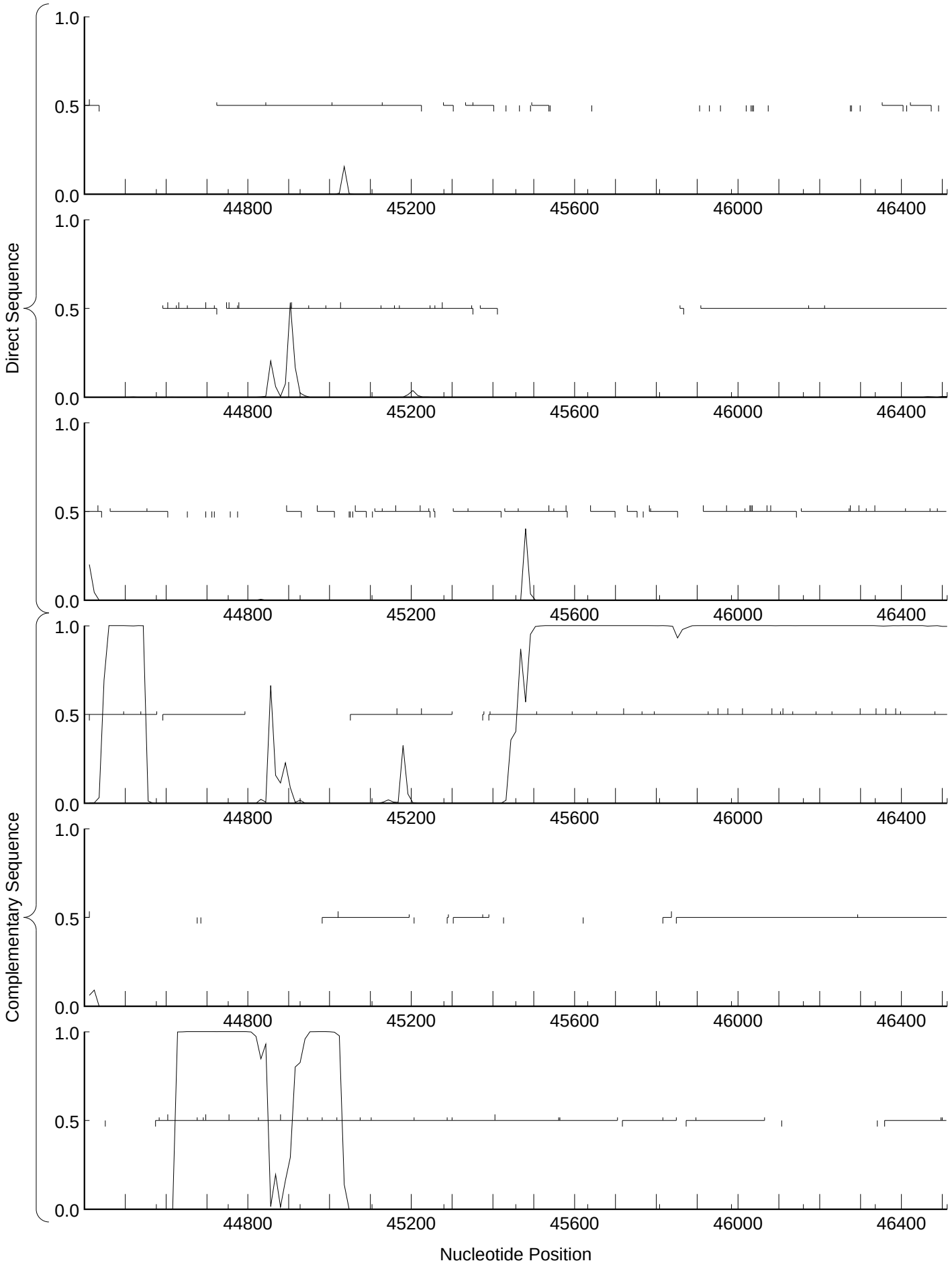
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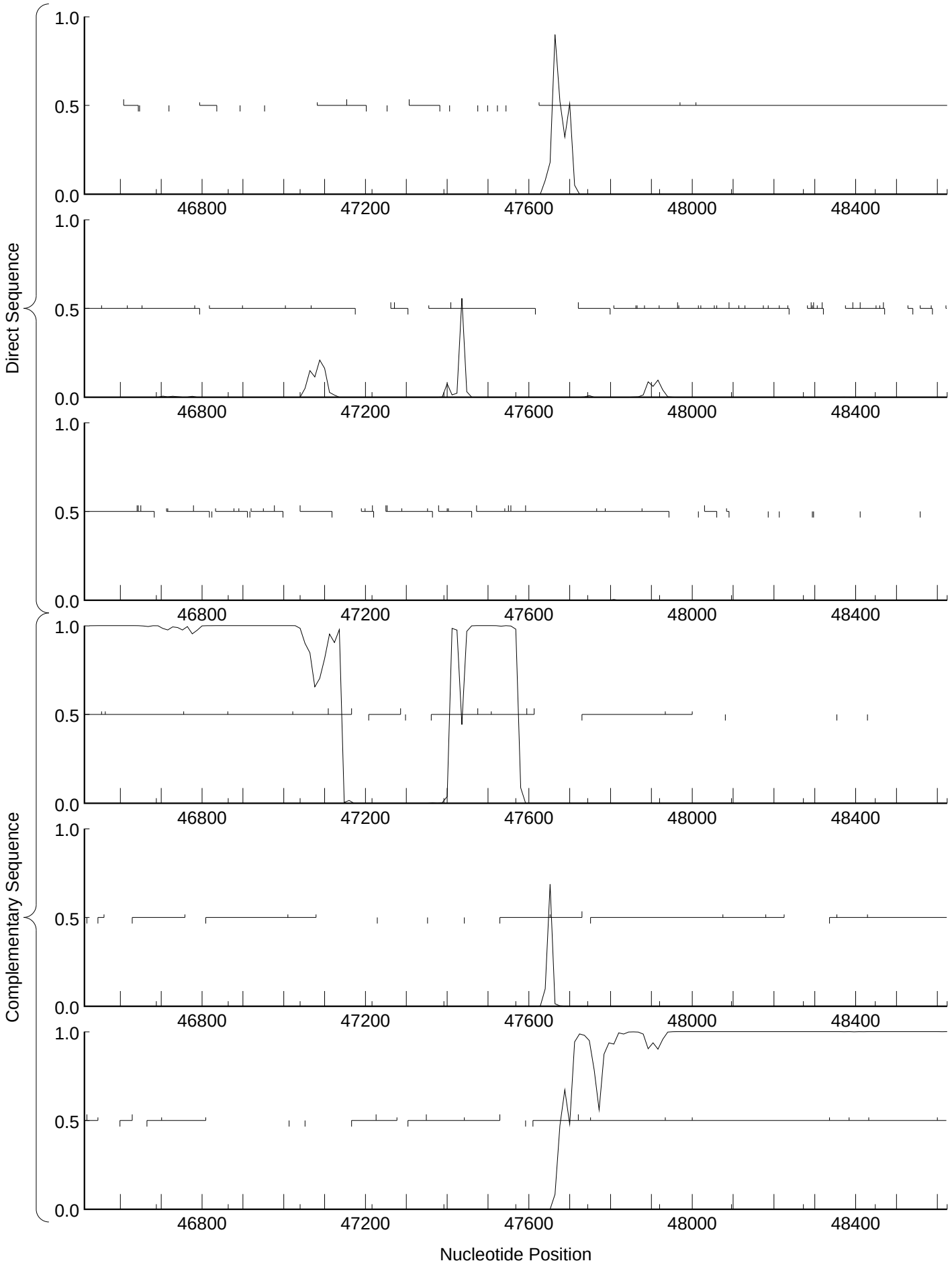
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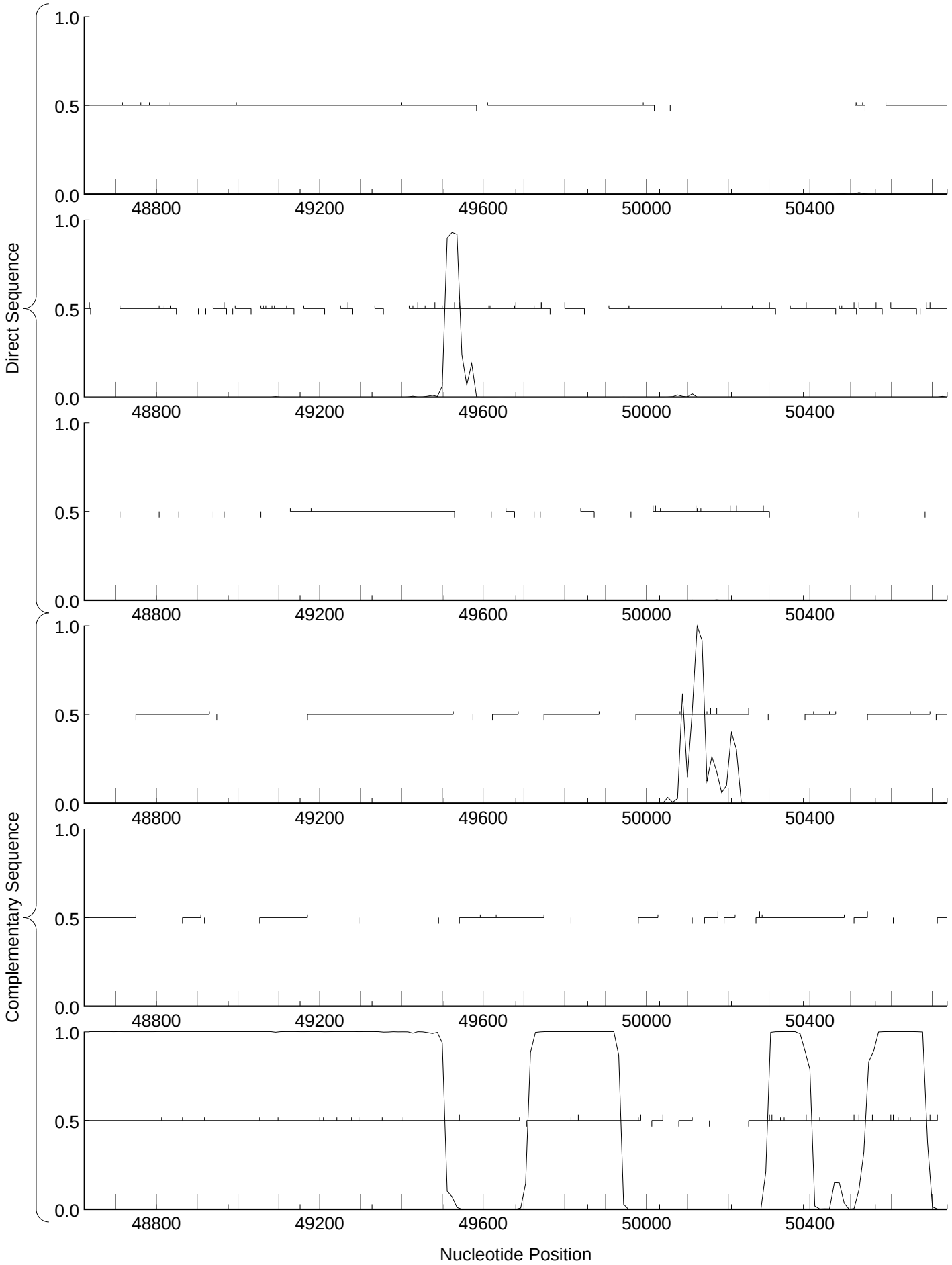
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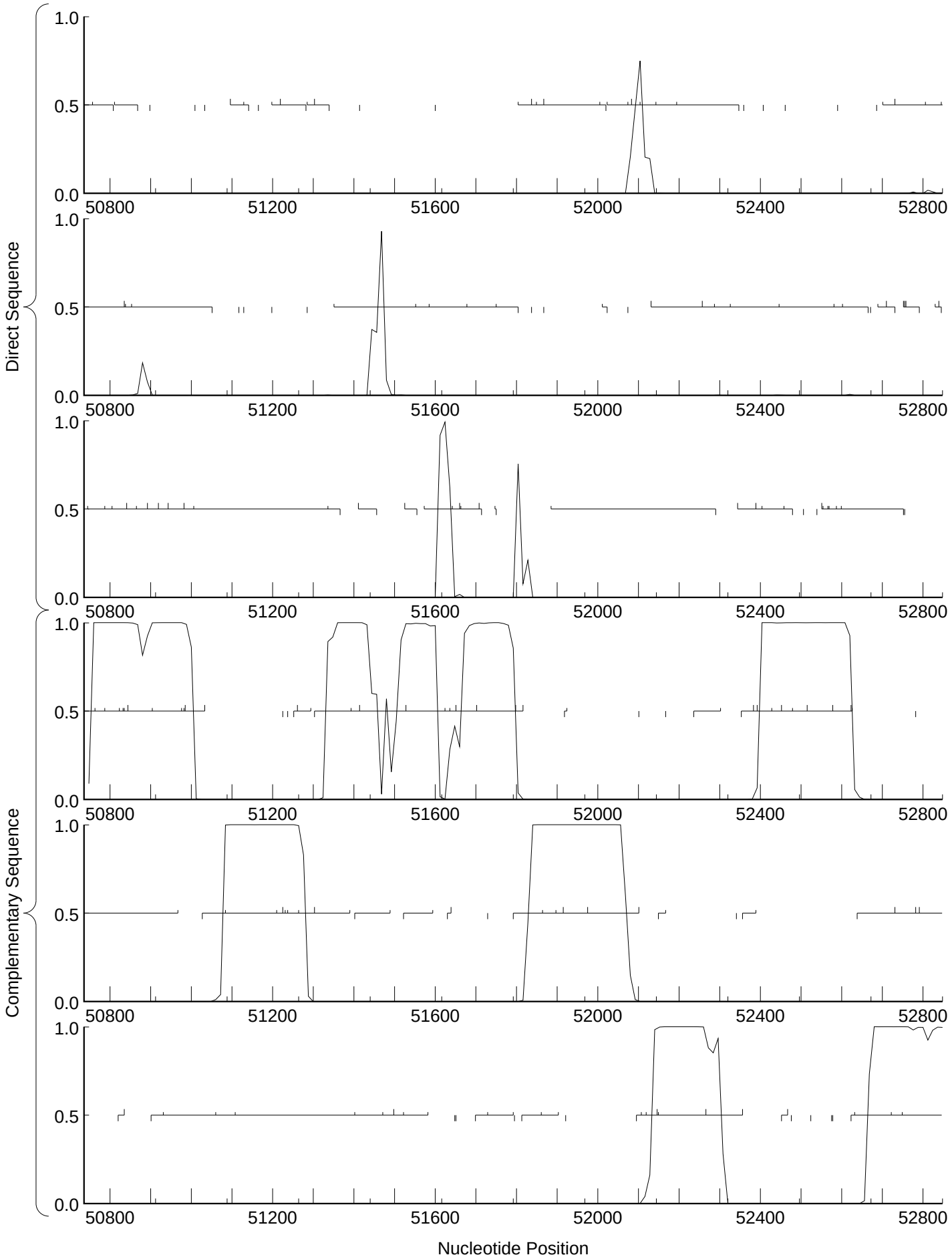
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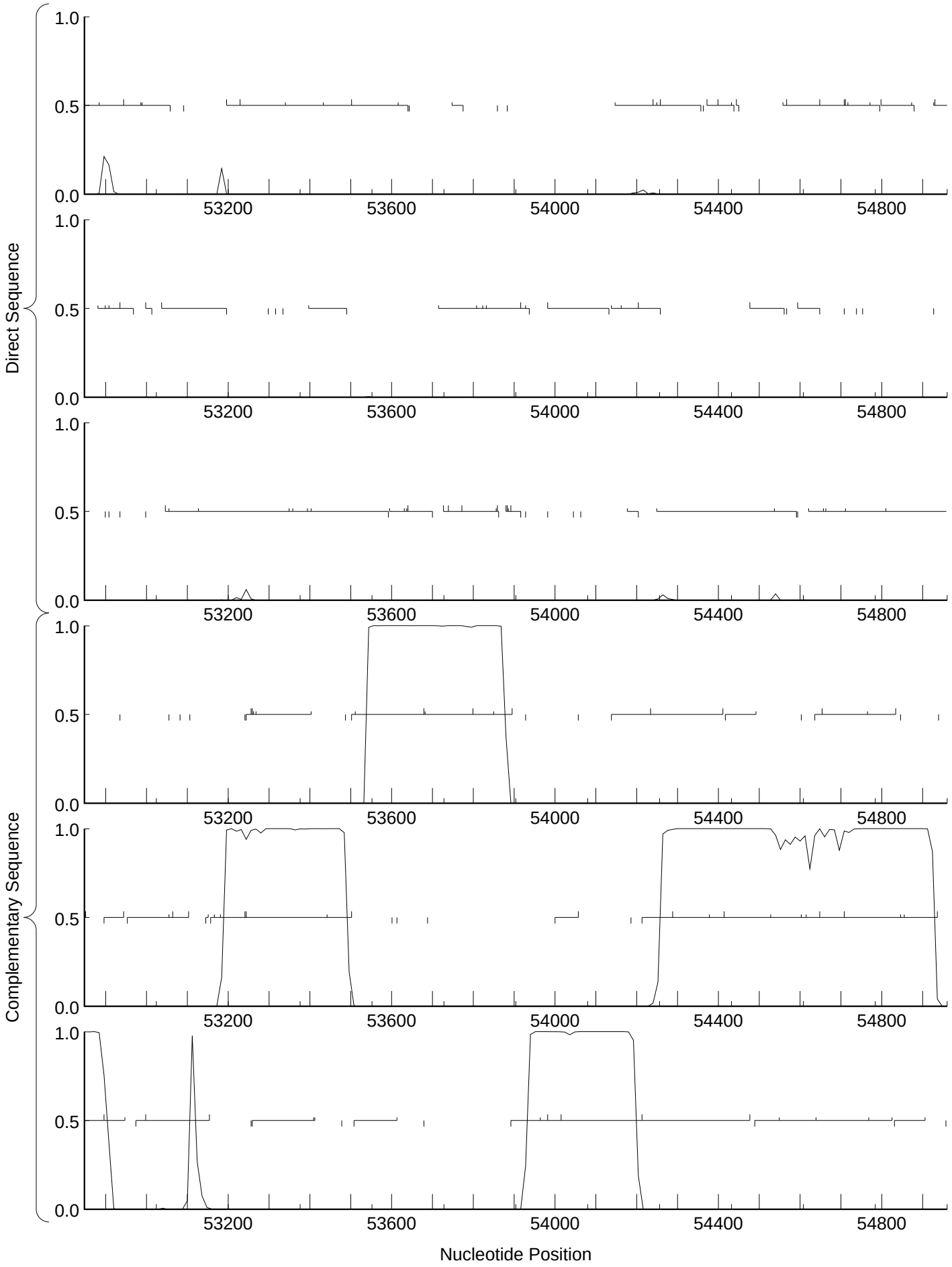
ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 25/31

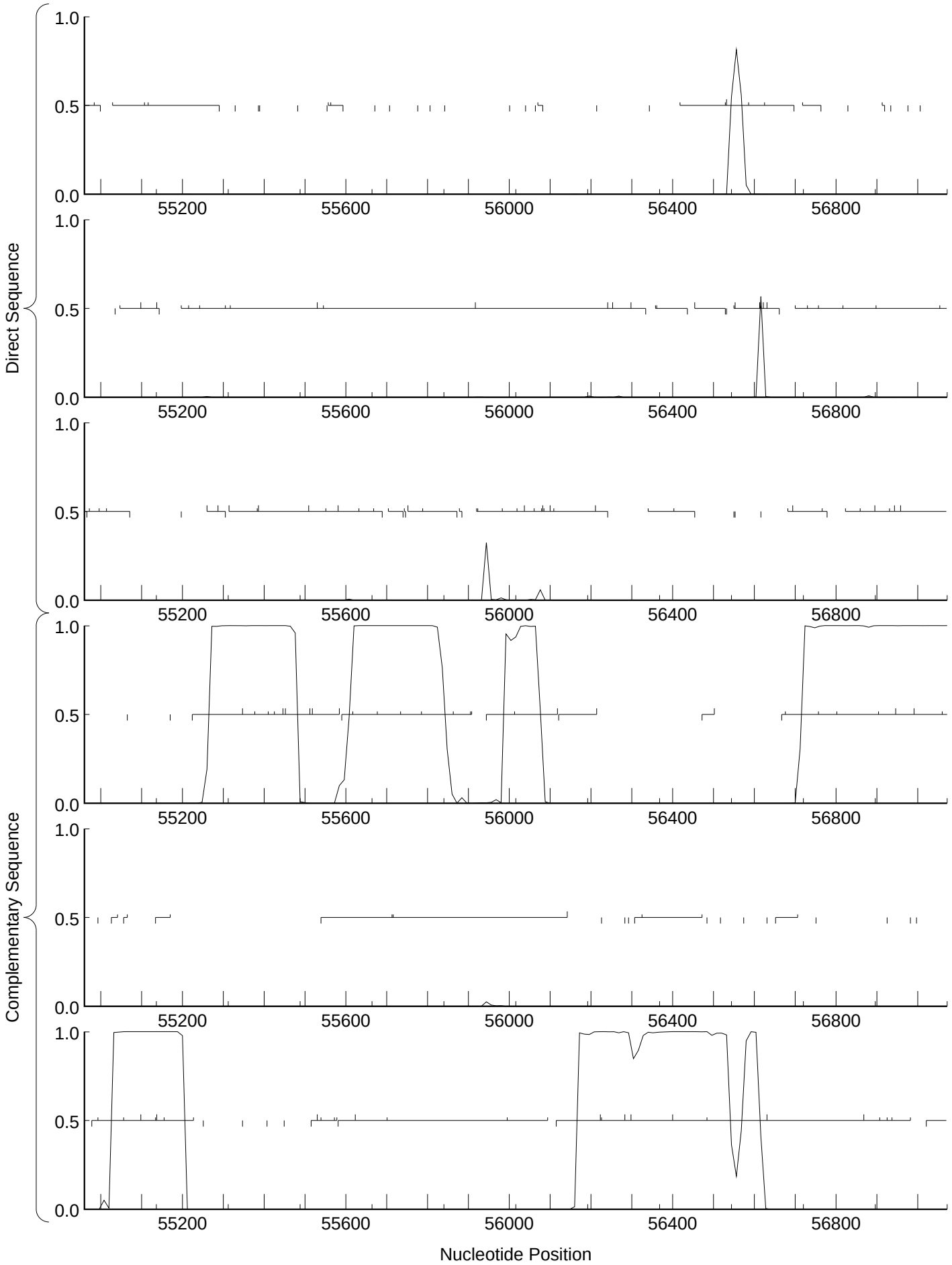


ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 26/31

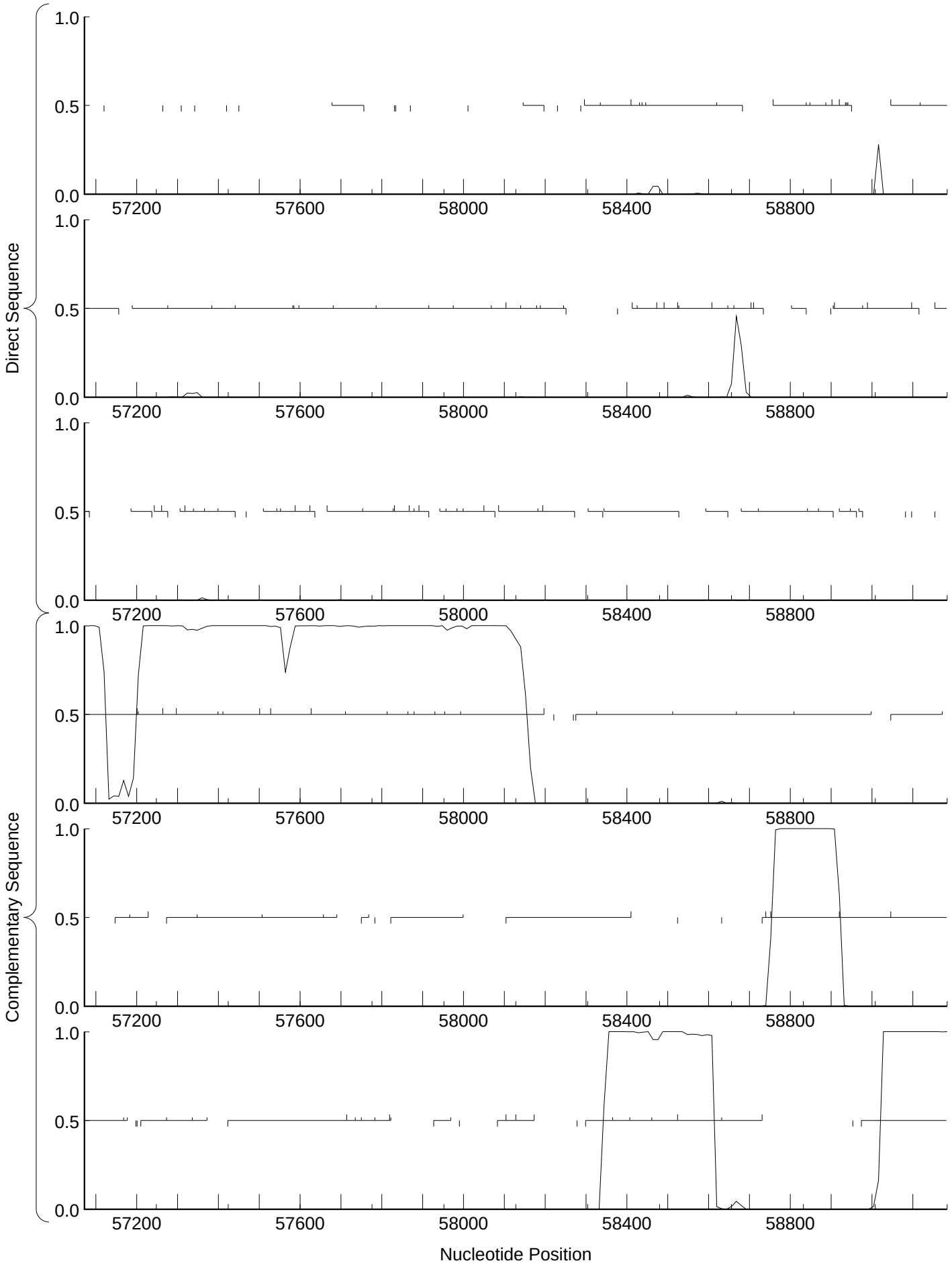


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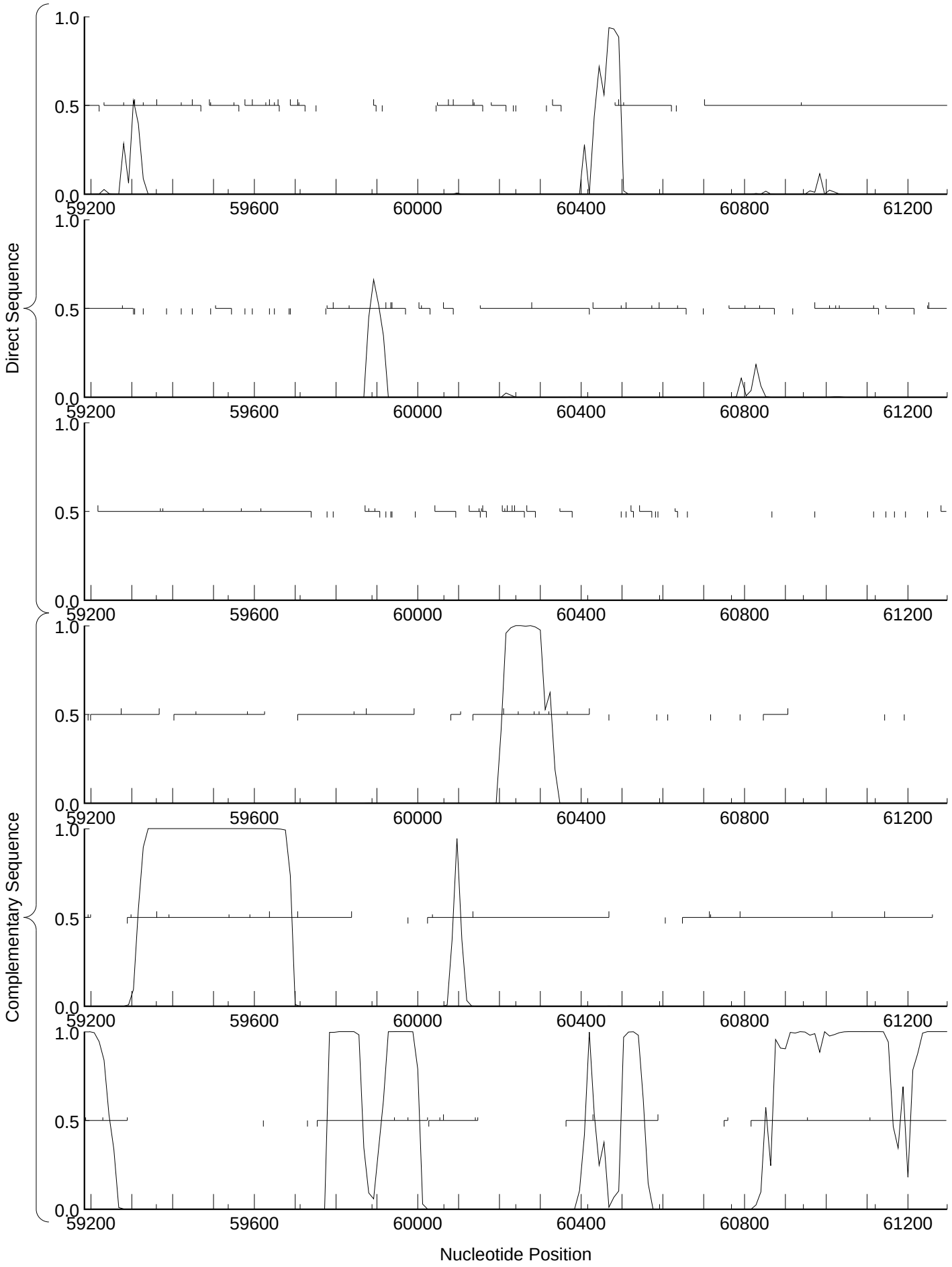



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
55200
55600
56000
56400
56800
Nucleotide Position

ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 29/31



ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 30/31



ge OneinaGillian complete sequence, 61703 bp including 203 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 31/31

