GeneMark

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

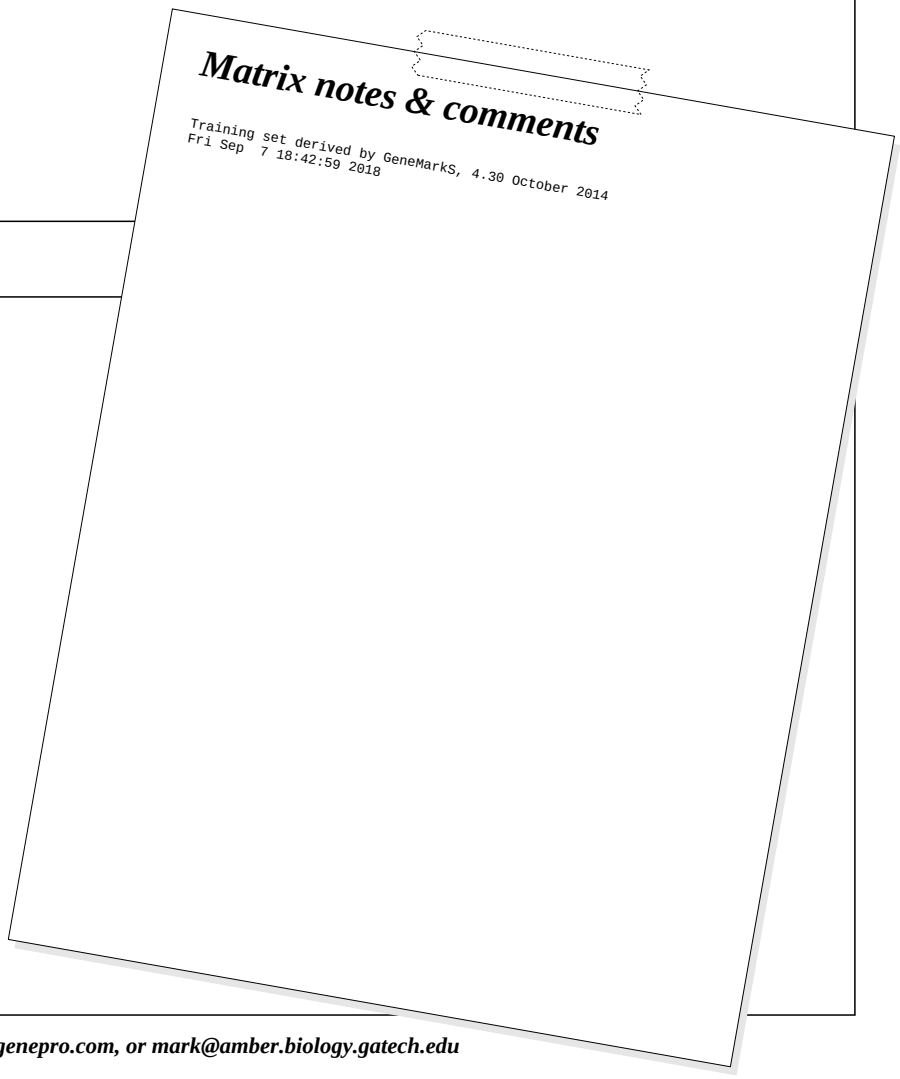
PROGRAM INFORMATION

```
Sequence         :  Microbacterium phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, C
Analysis Date    :  3/28/24 at 7:24:33
Pages            :  31
Sequence Length  :  62358 bp
GC Content       :   66.91%

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

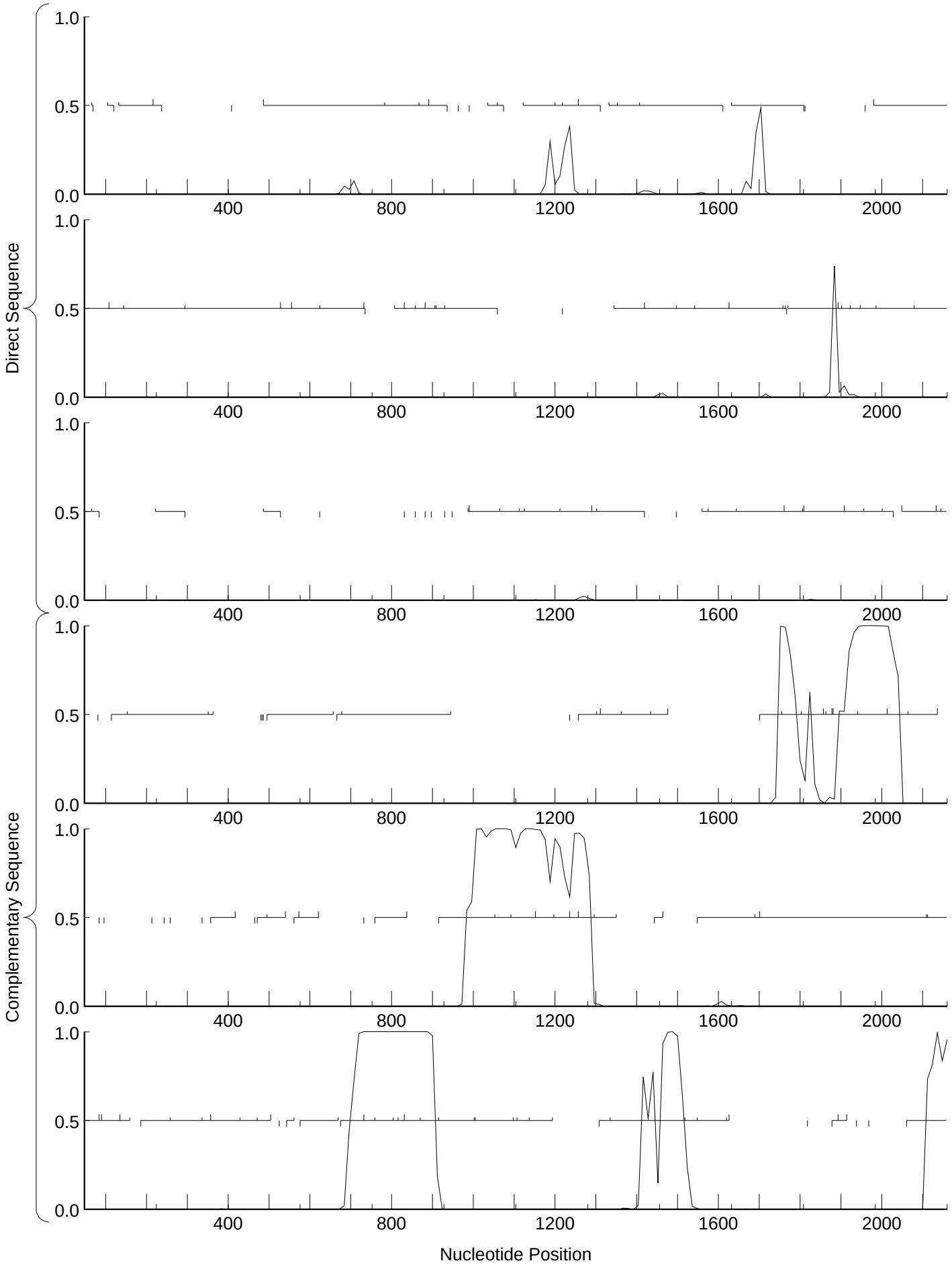
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Matrix     :  -
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Order      :  4
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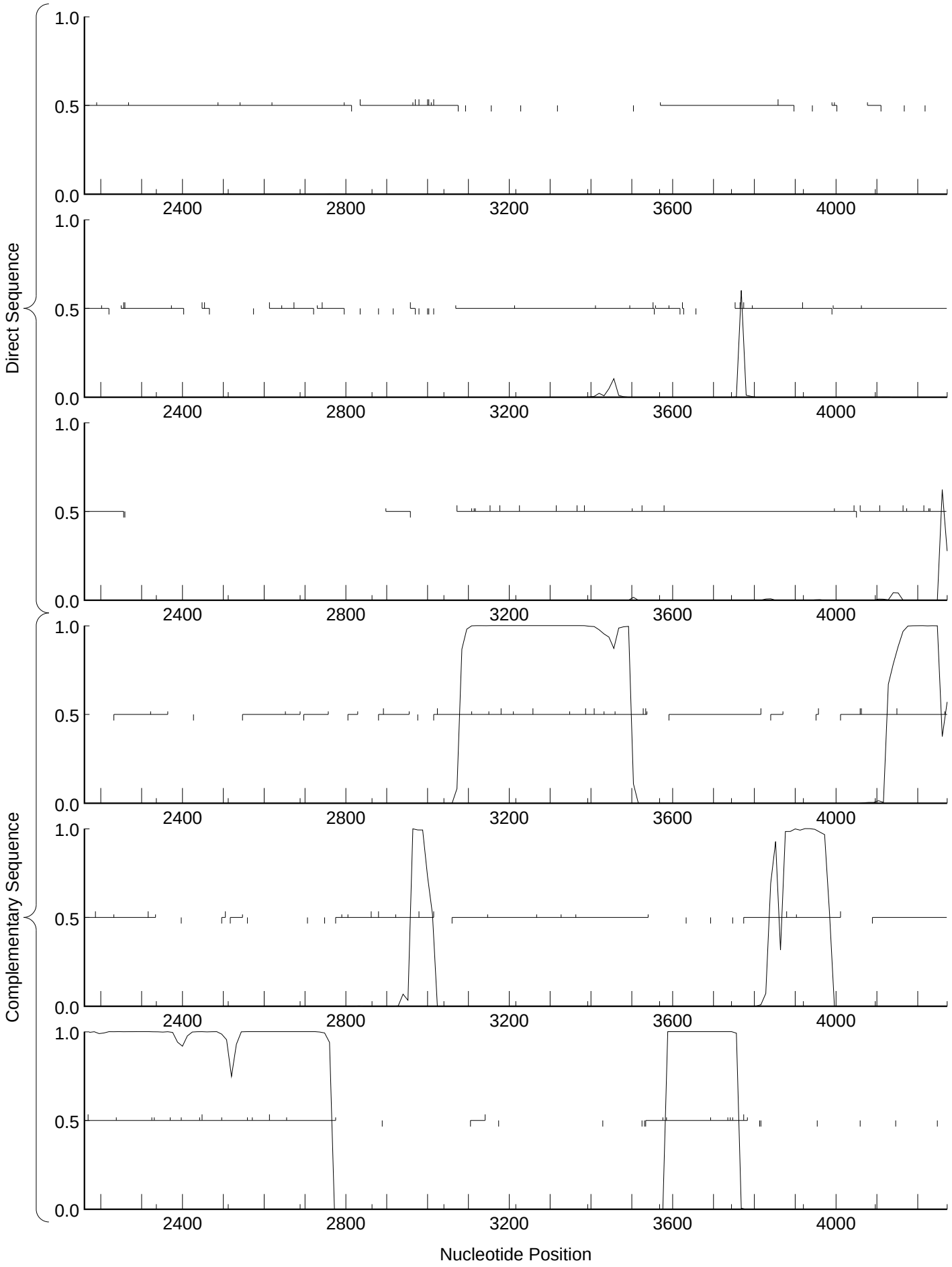

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

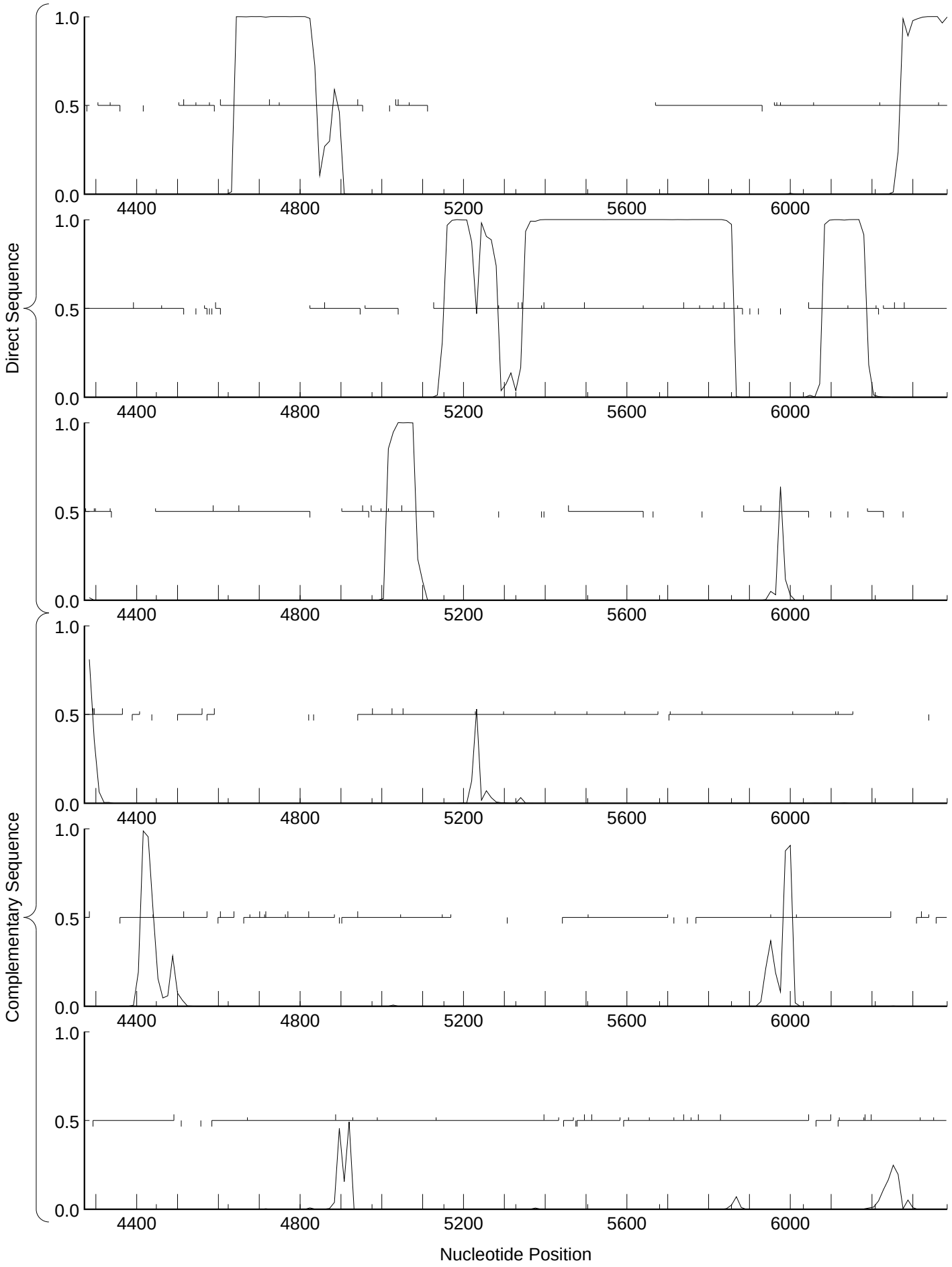
m phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 2/31



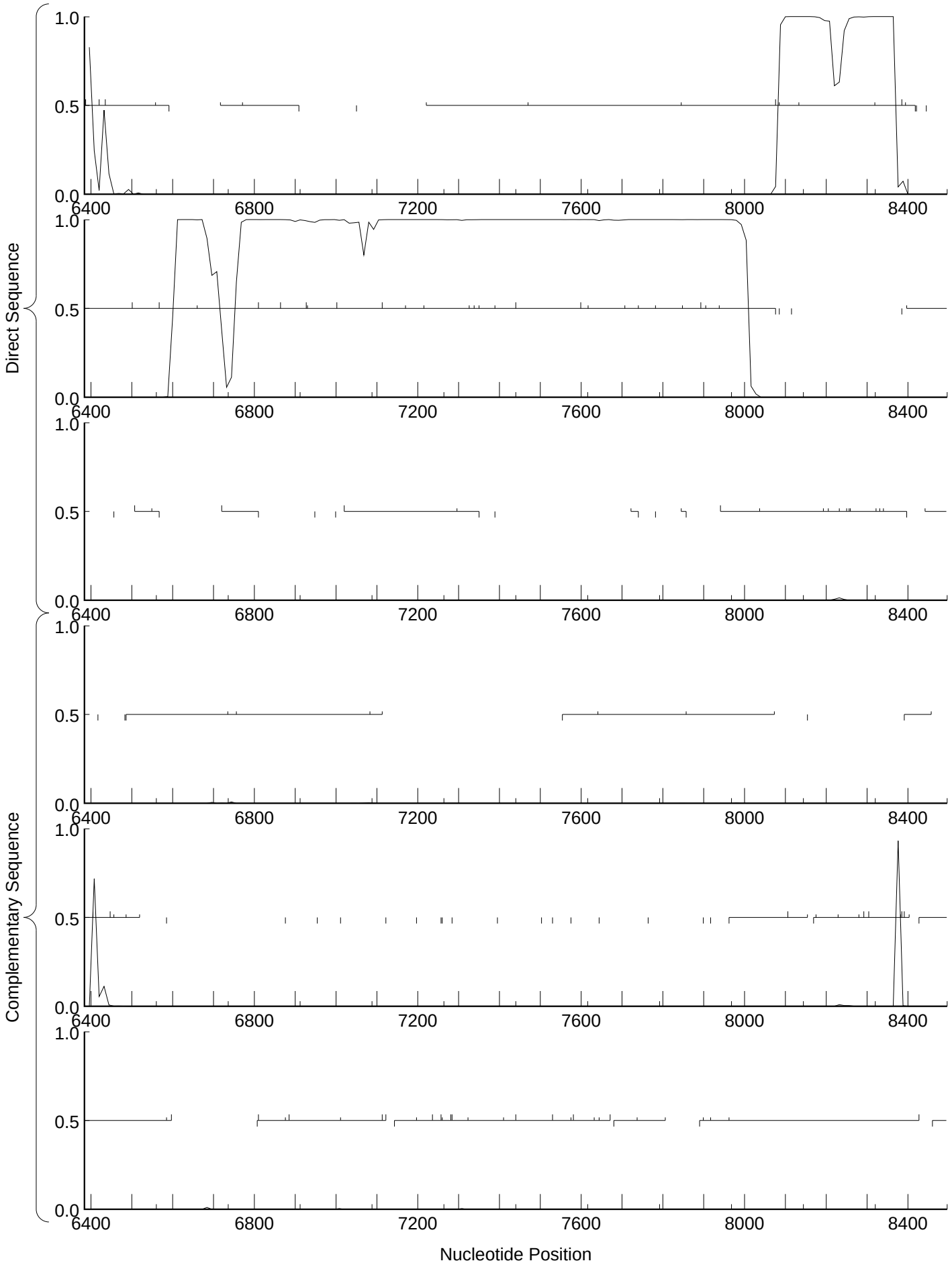
m phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 3/31



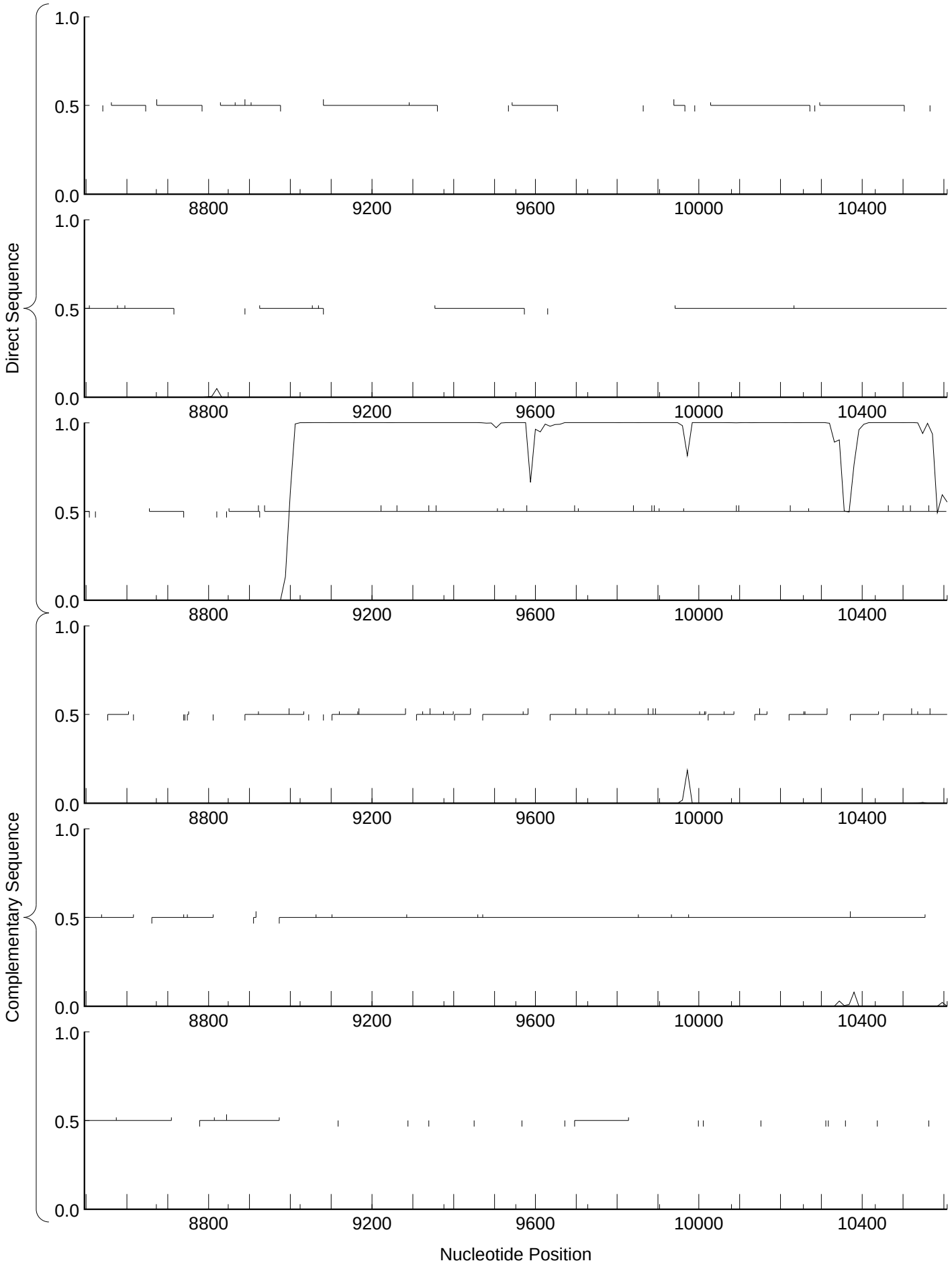
m phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 4/31



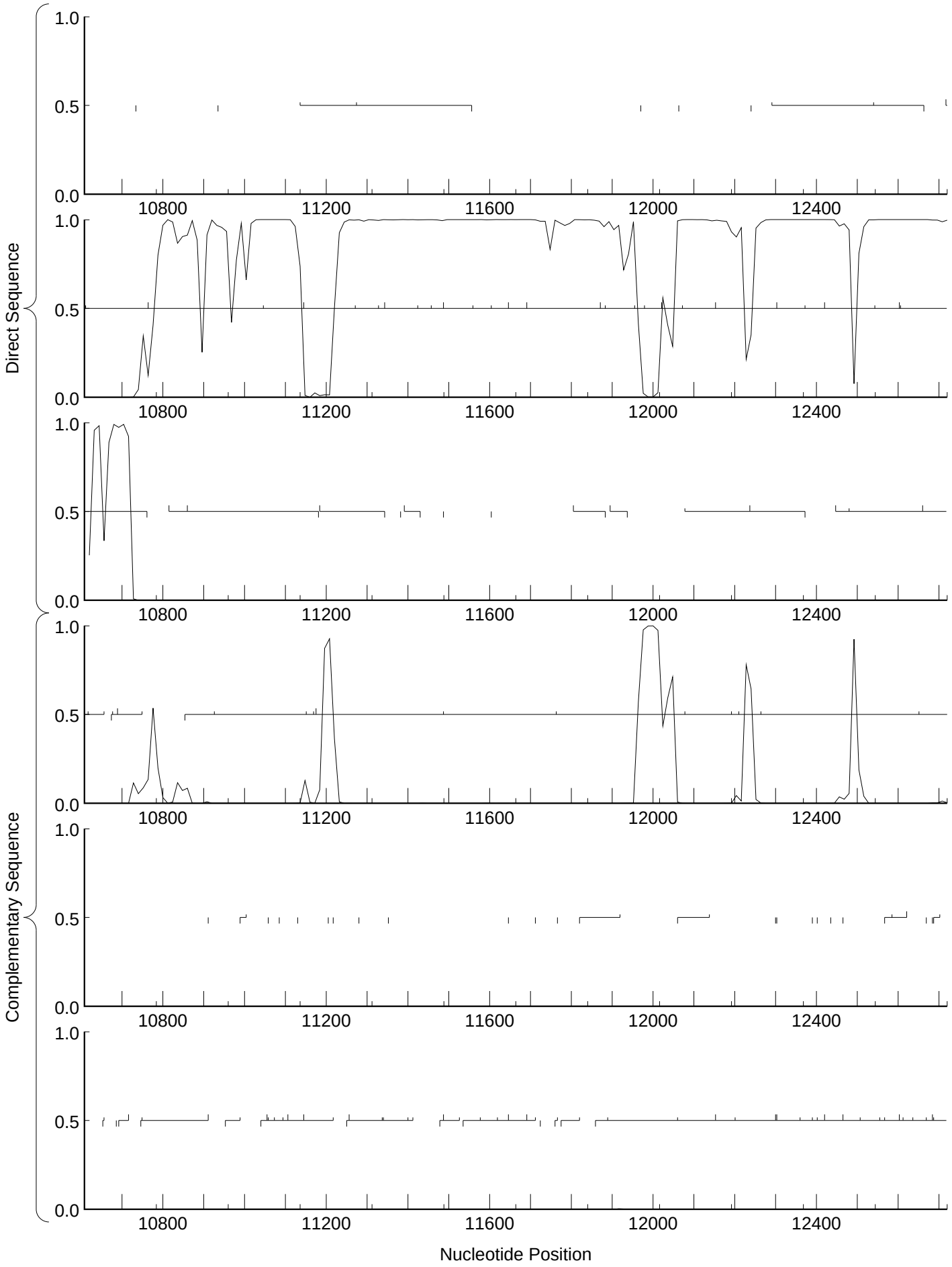
m phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 5/31



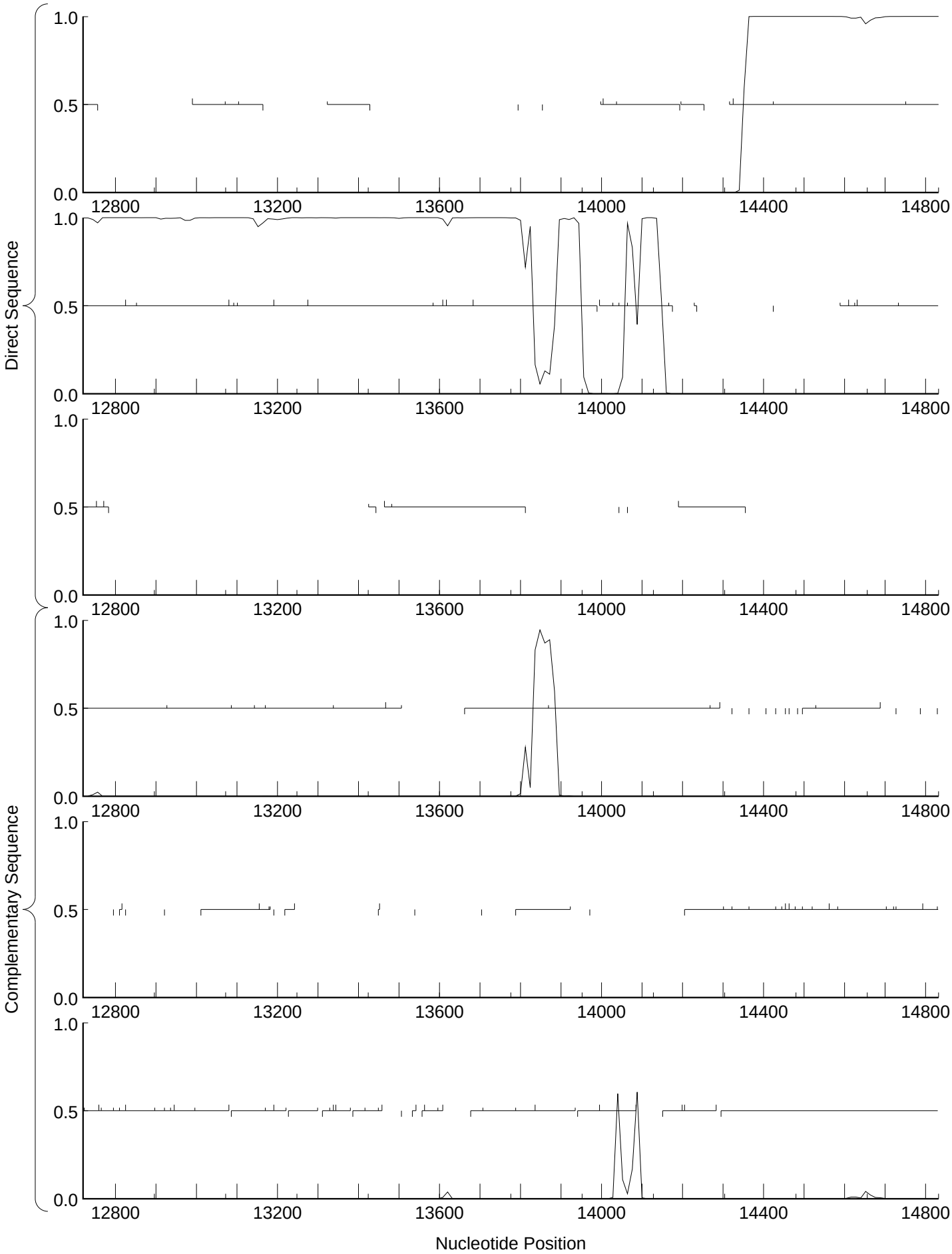
m phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 6/31



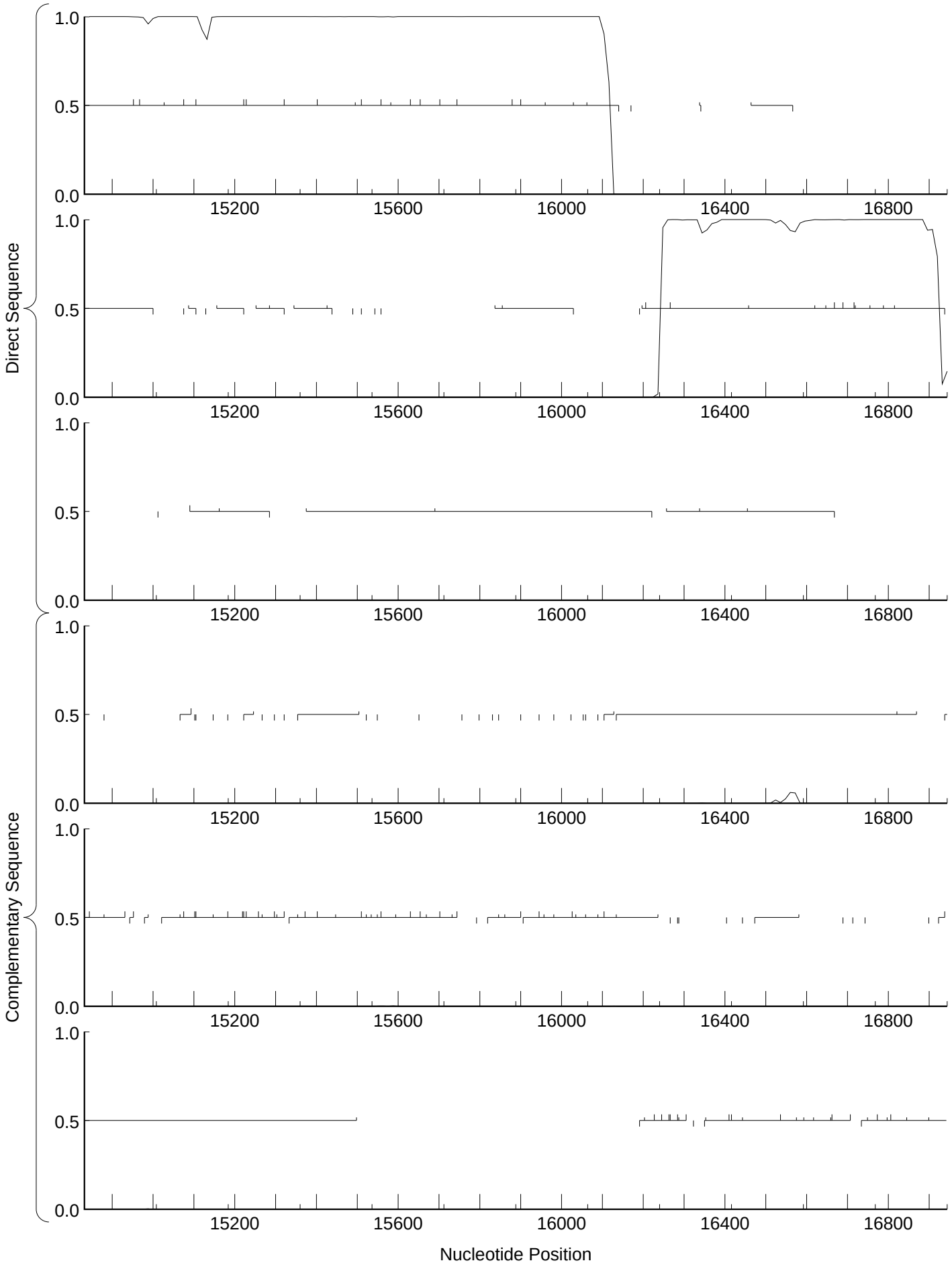
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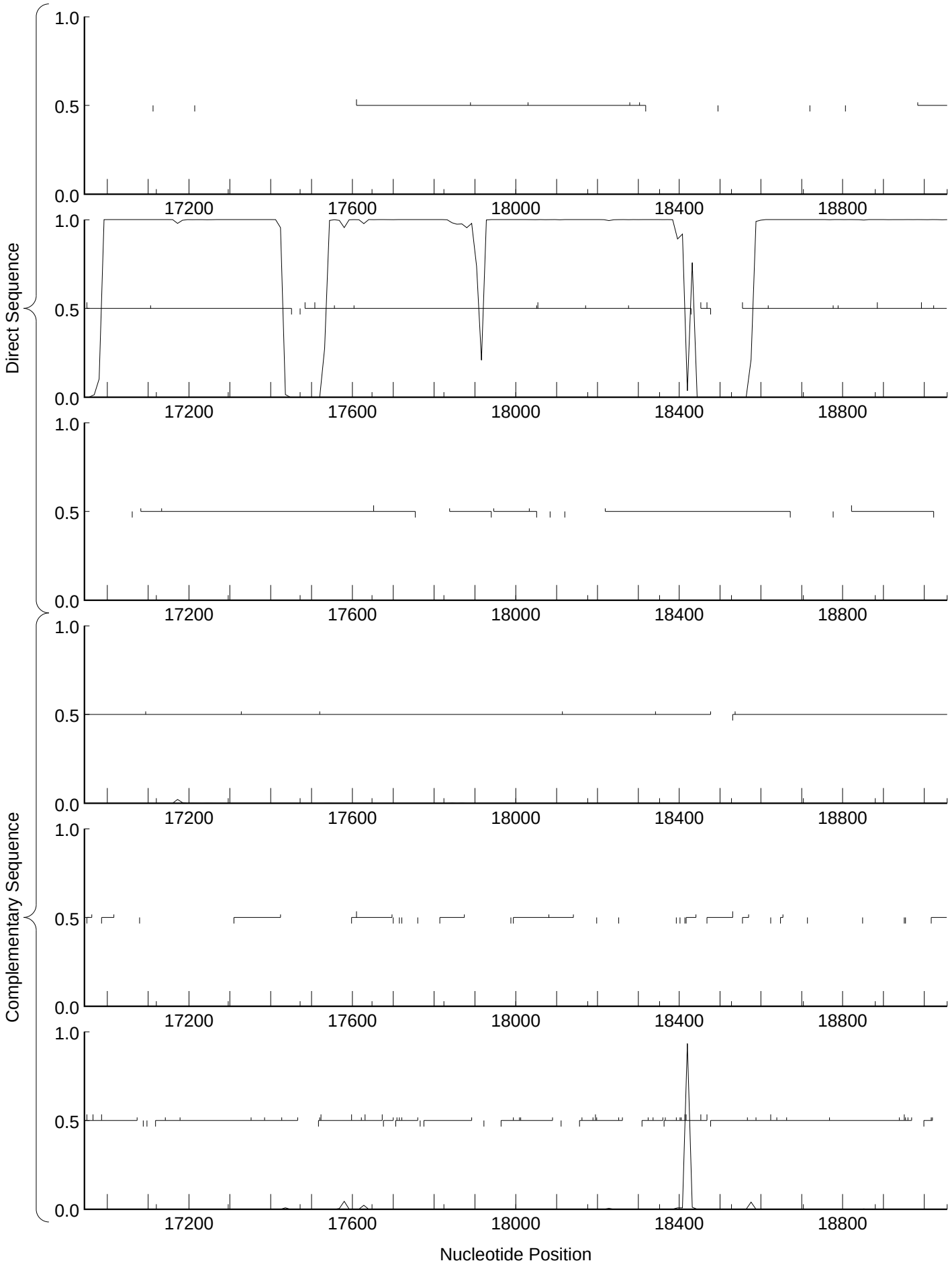


m phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 8/31

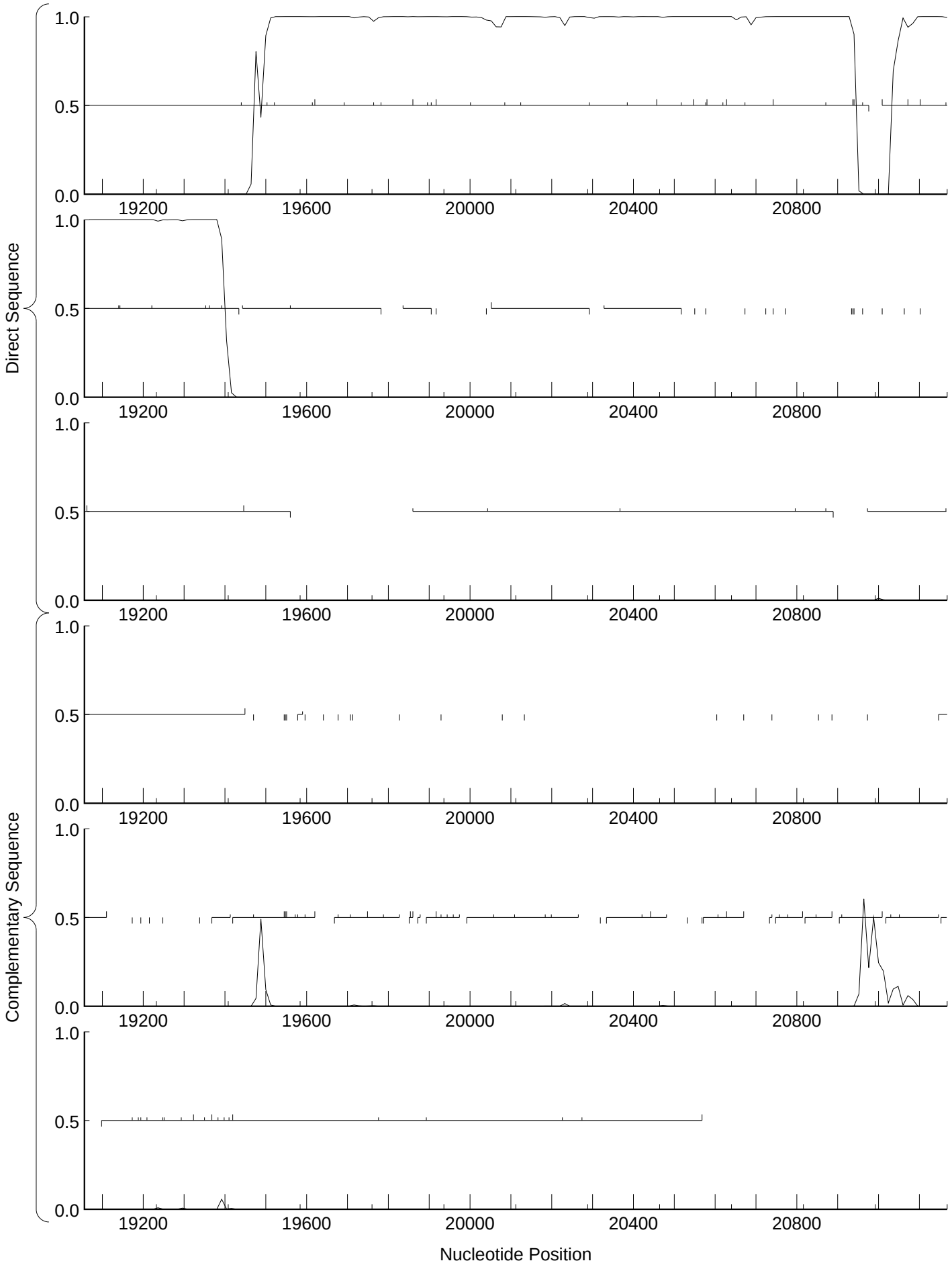


m phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 9/31

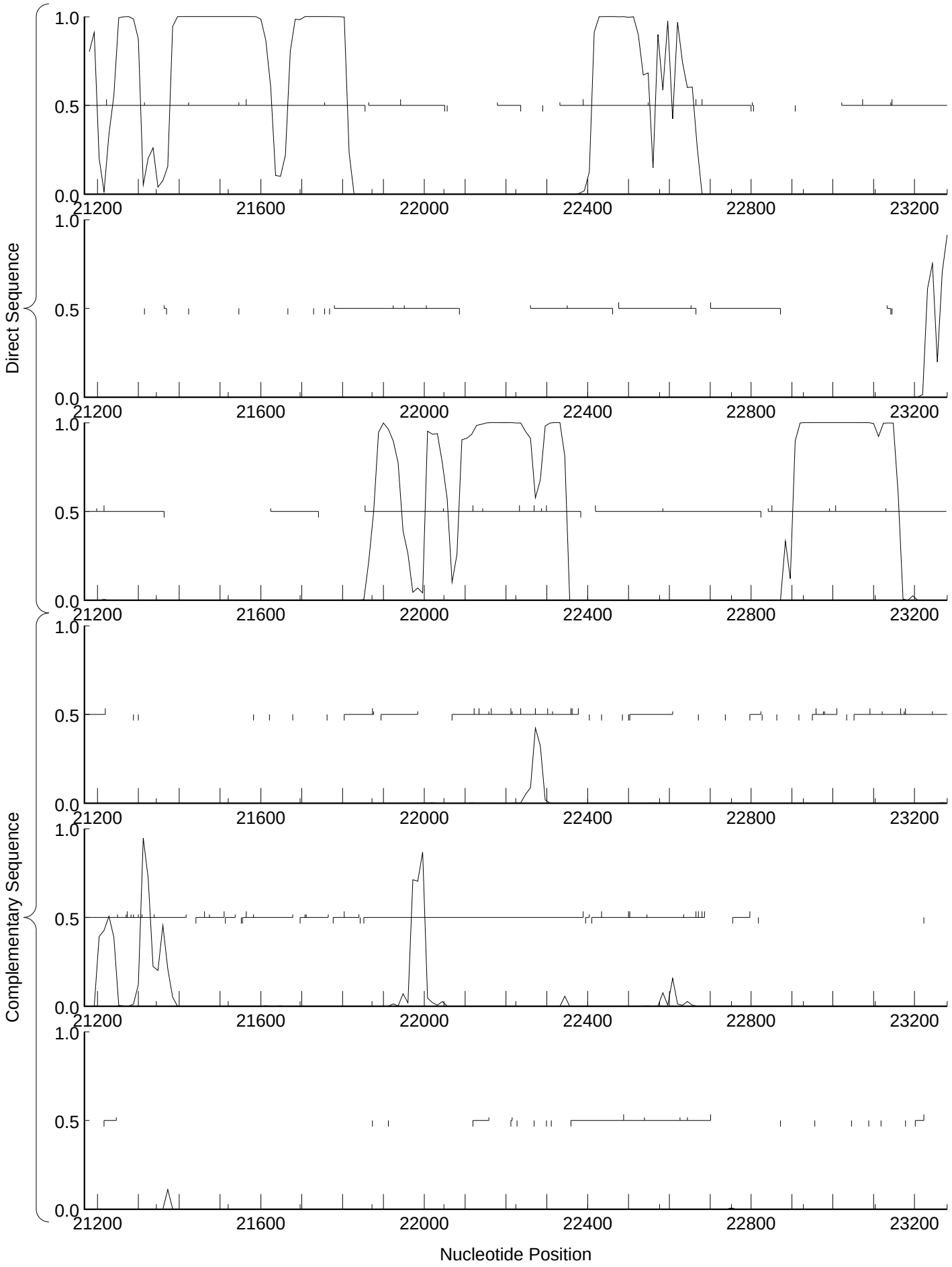



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
17200
17600
18000
18400
18800
Nucleotide Position

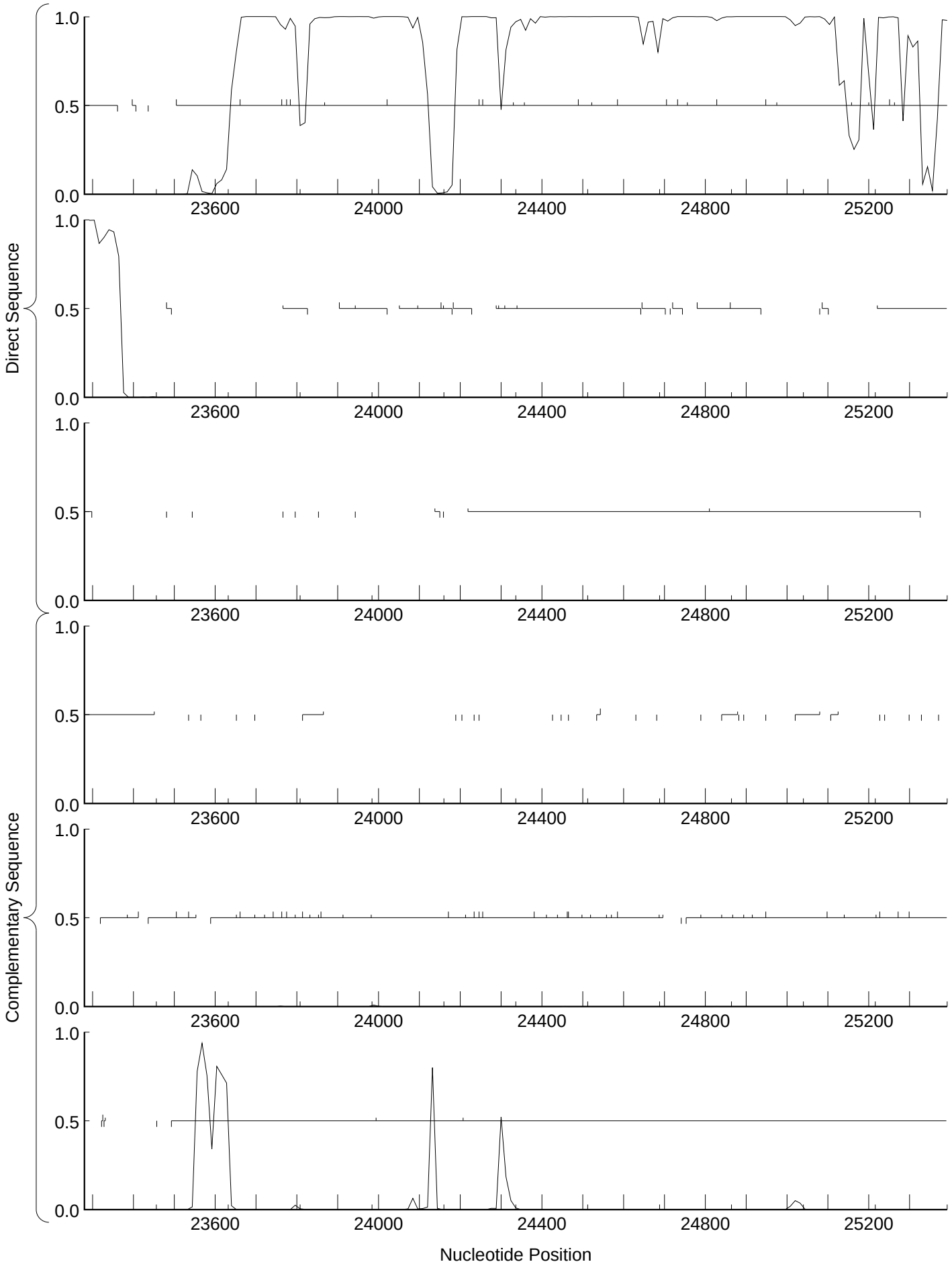
n phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 11/31

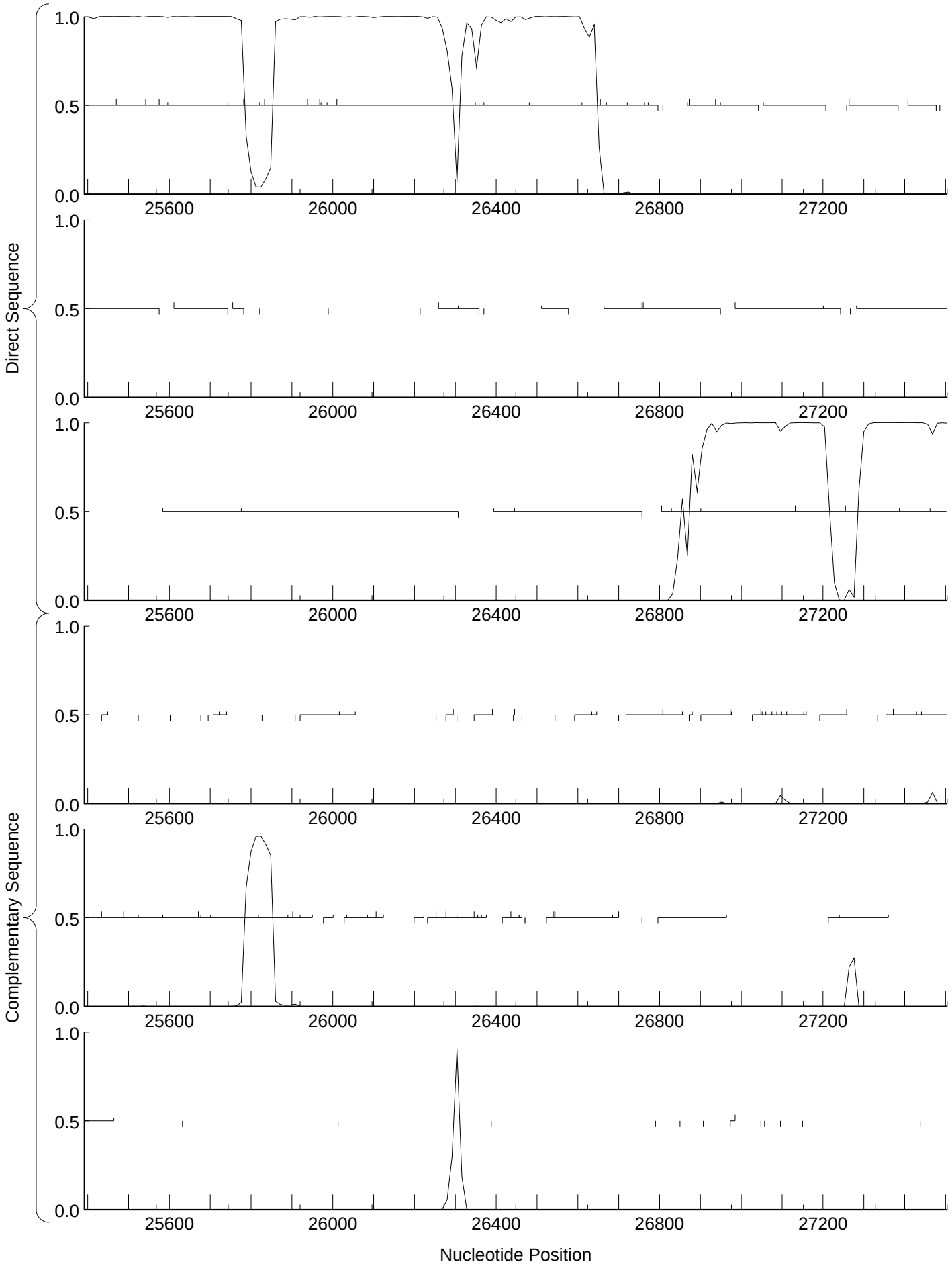


n phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 12/31

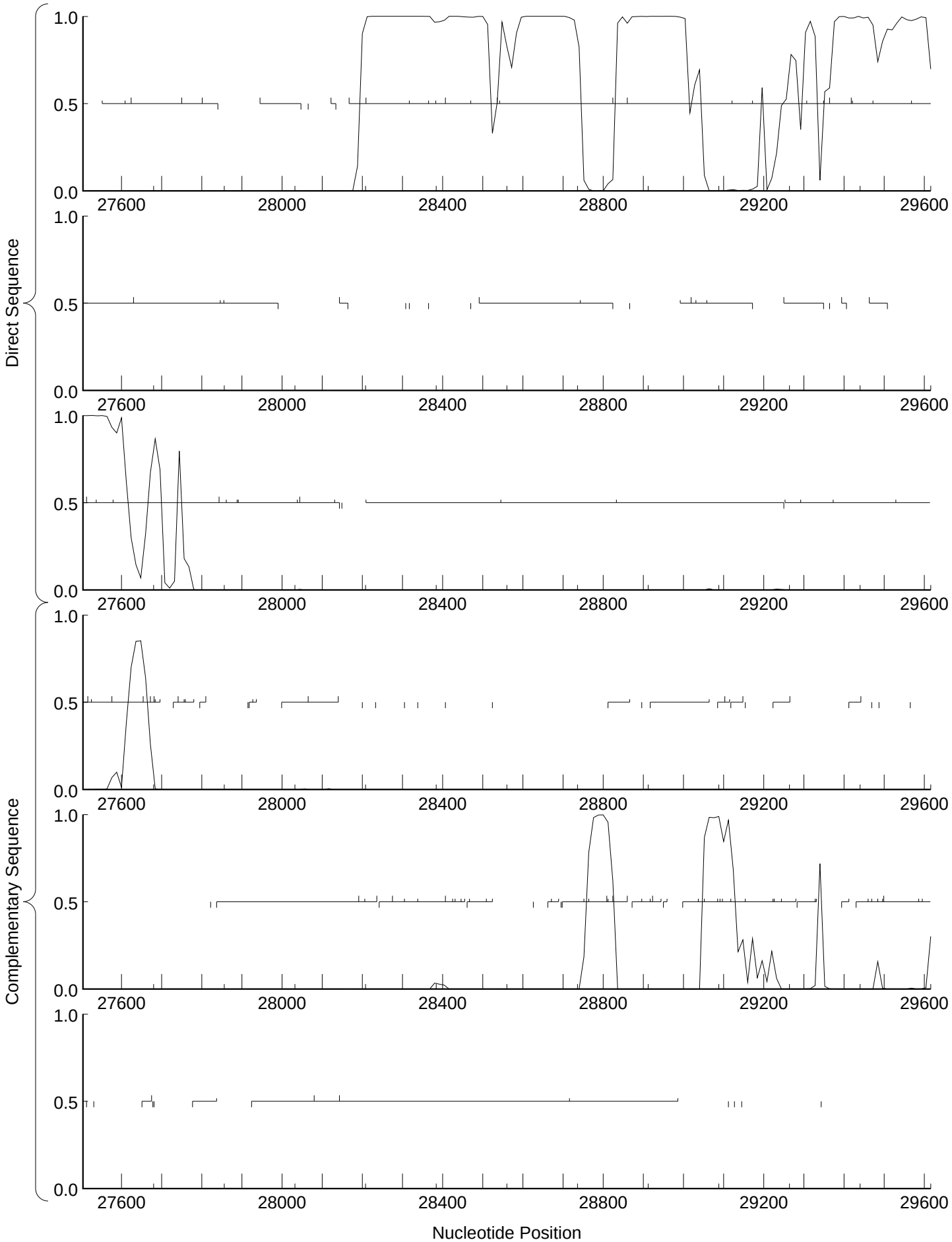


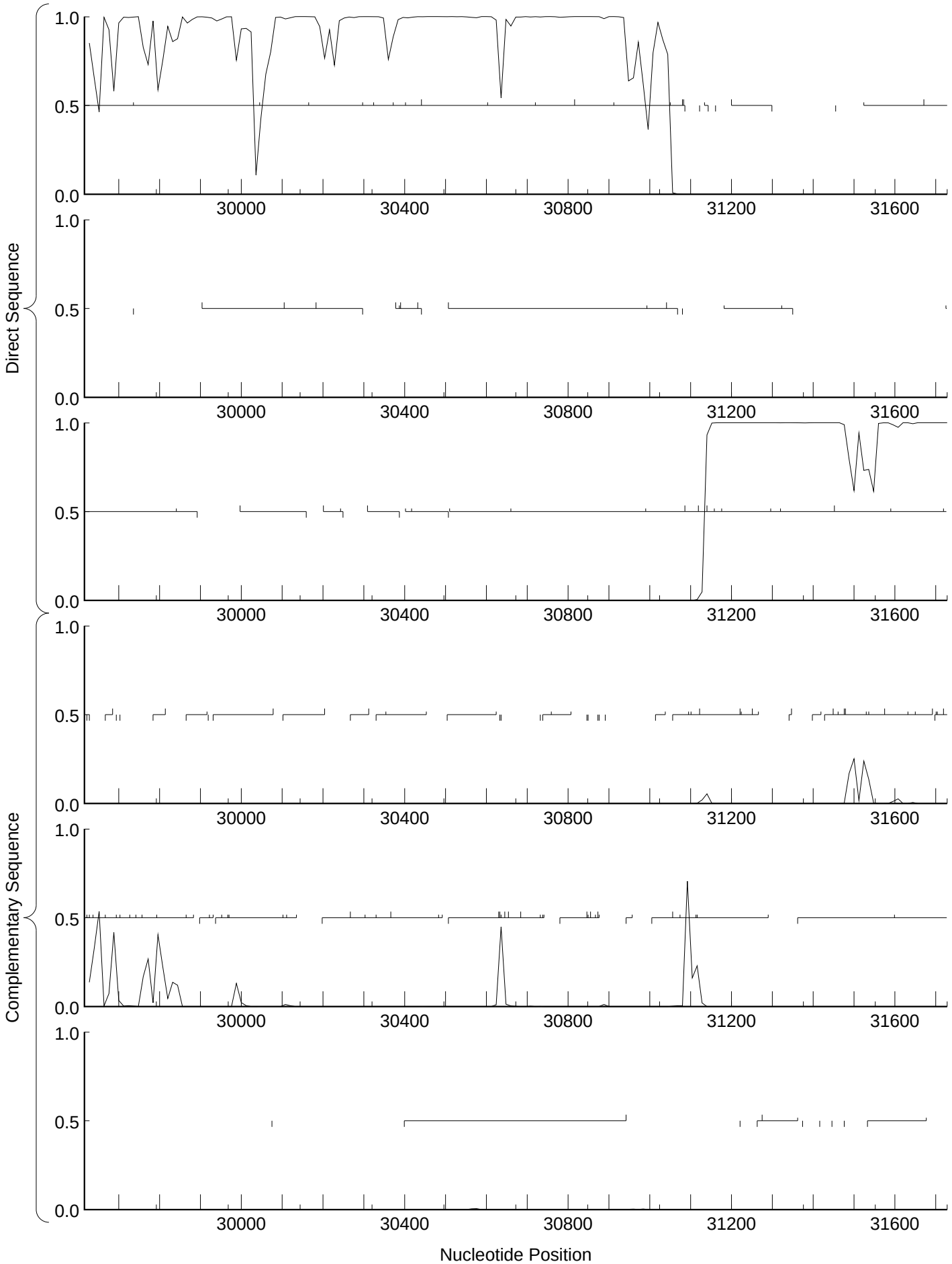
n phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 13/31



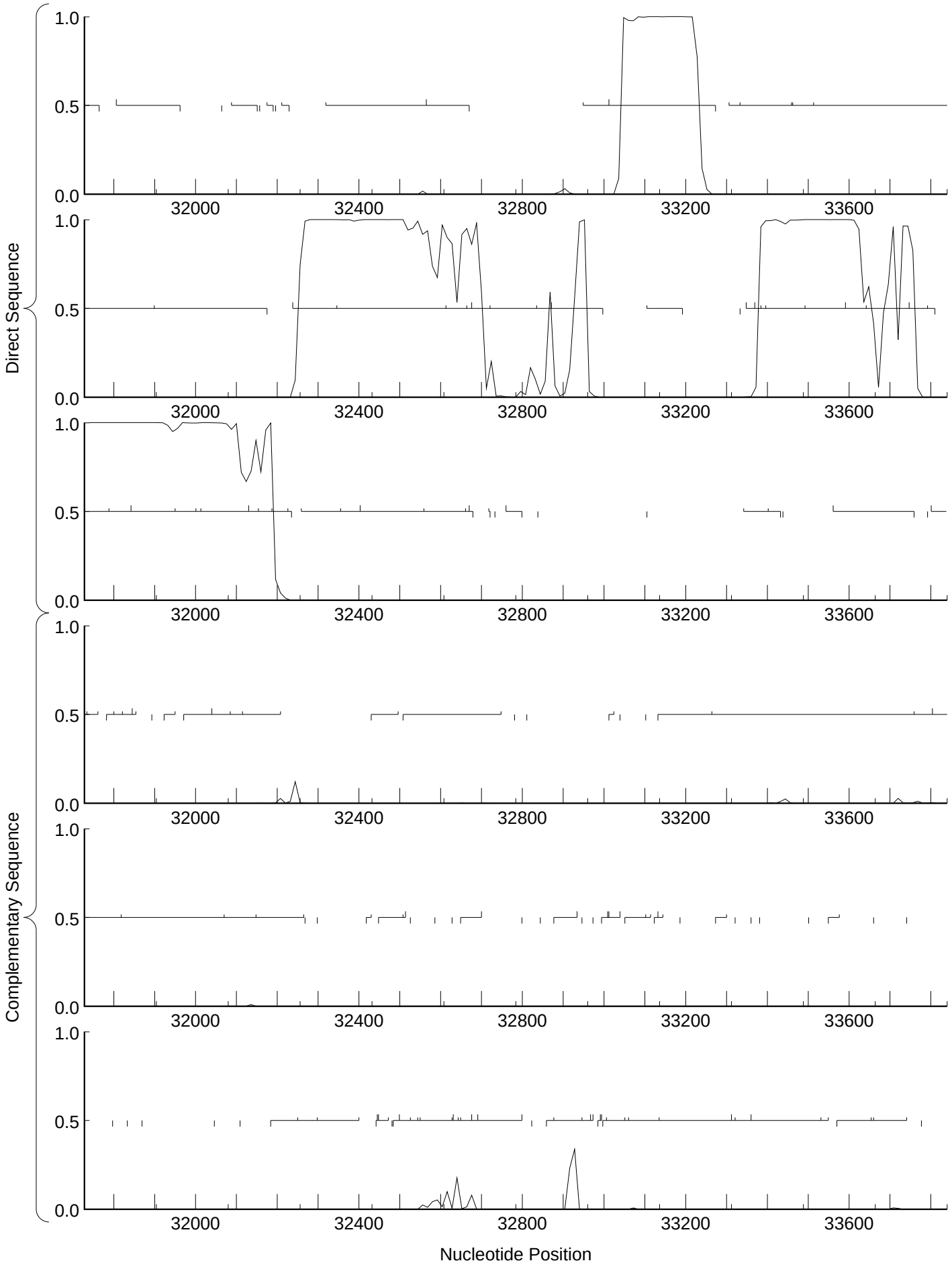

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
25600
26000
26400
26800
27200
Nucleotide Position

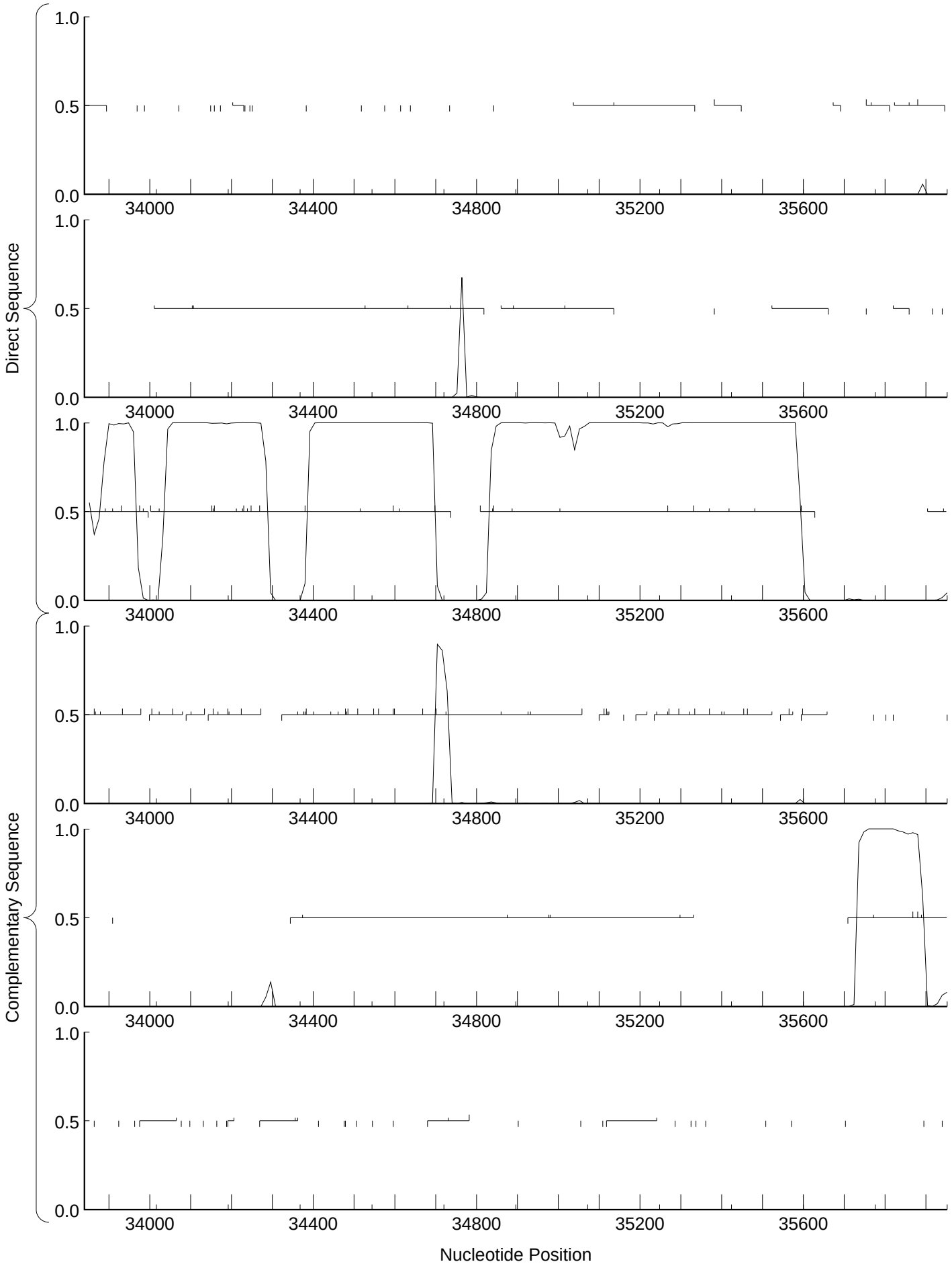
n phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 15/31



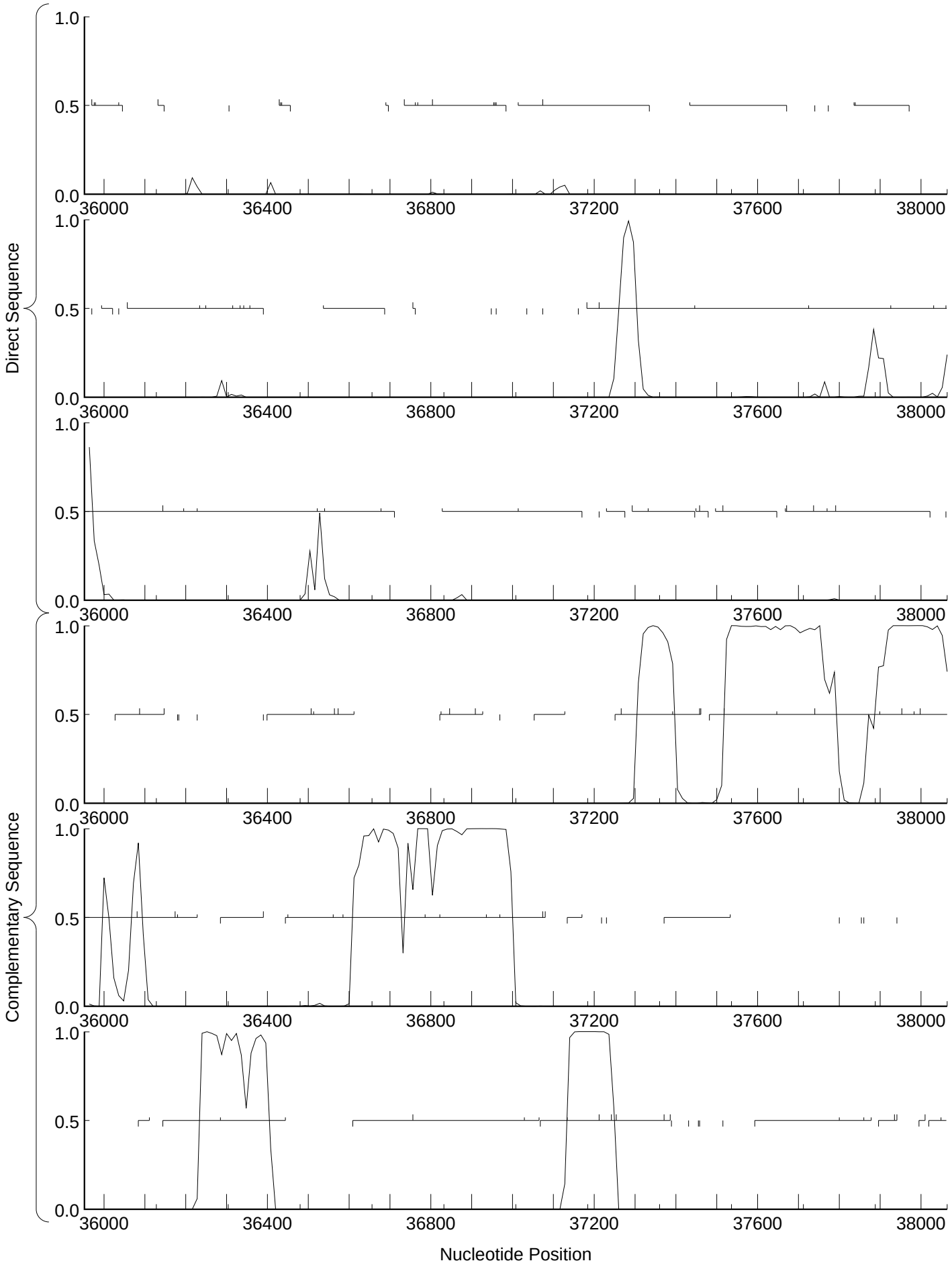

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
30000
30400
30800
31200
31600
Nucleotide Position

n phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 17/31

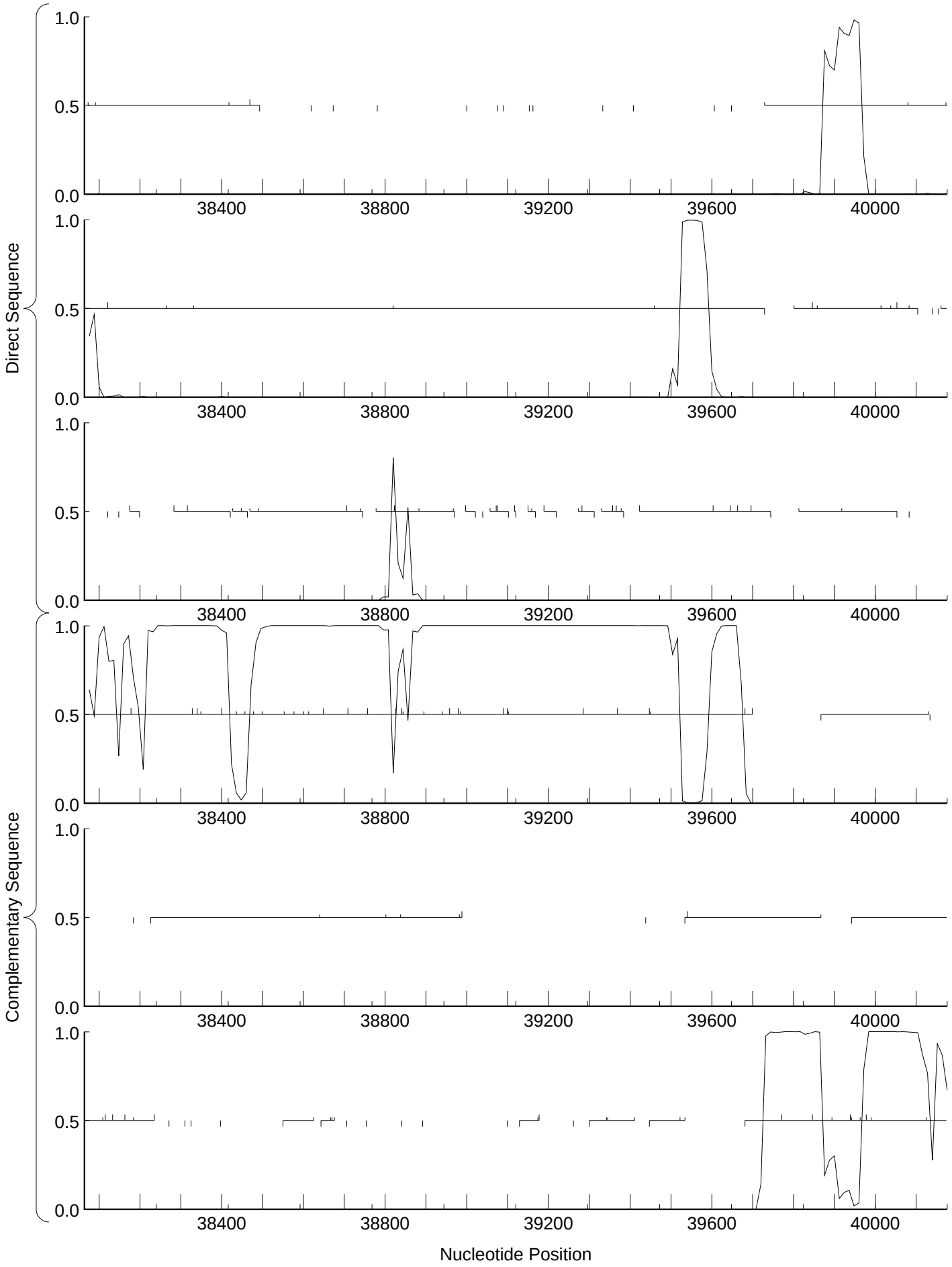



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

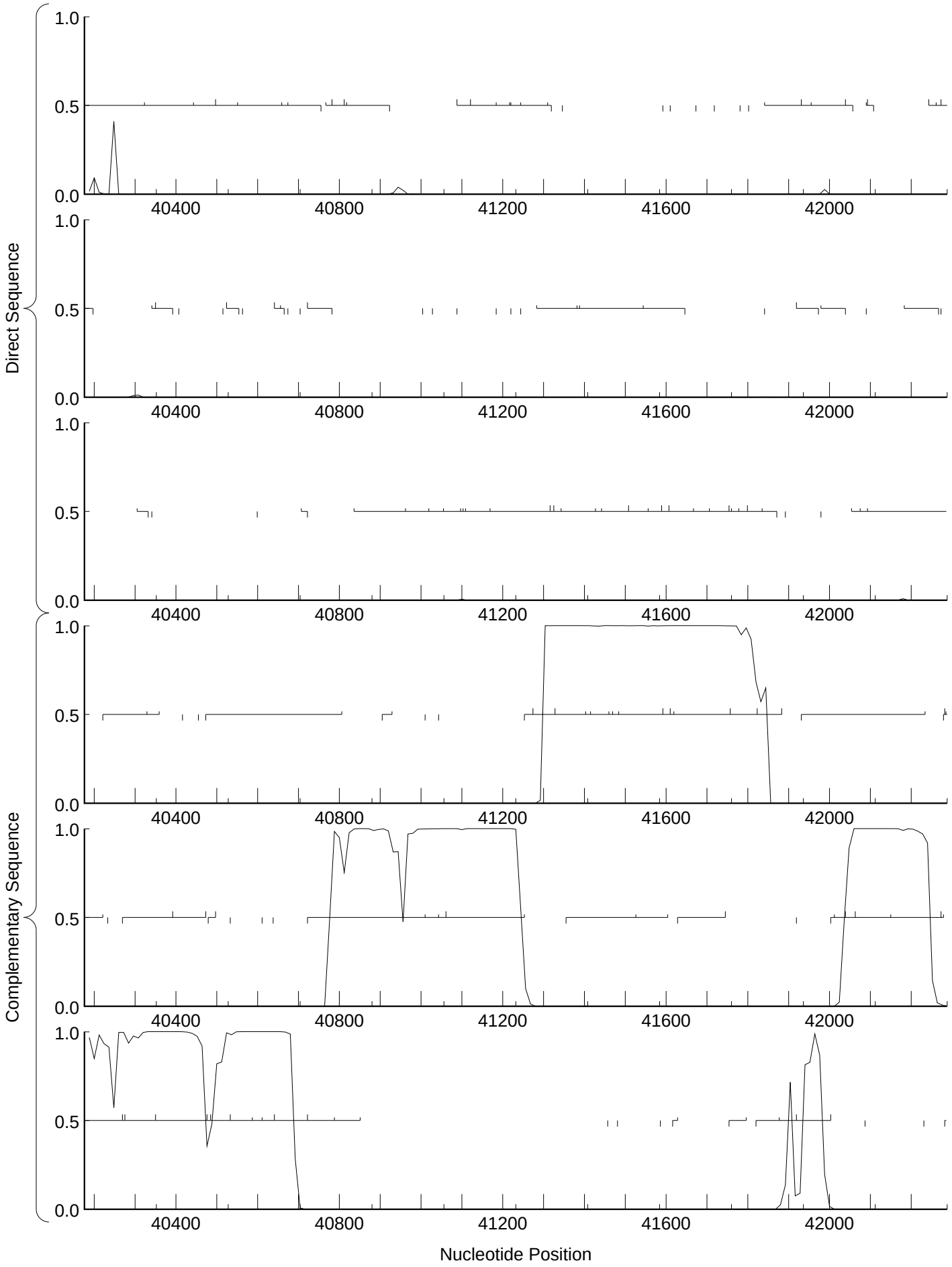
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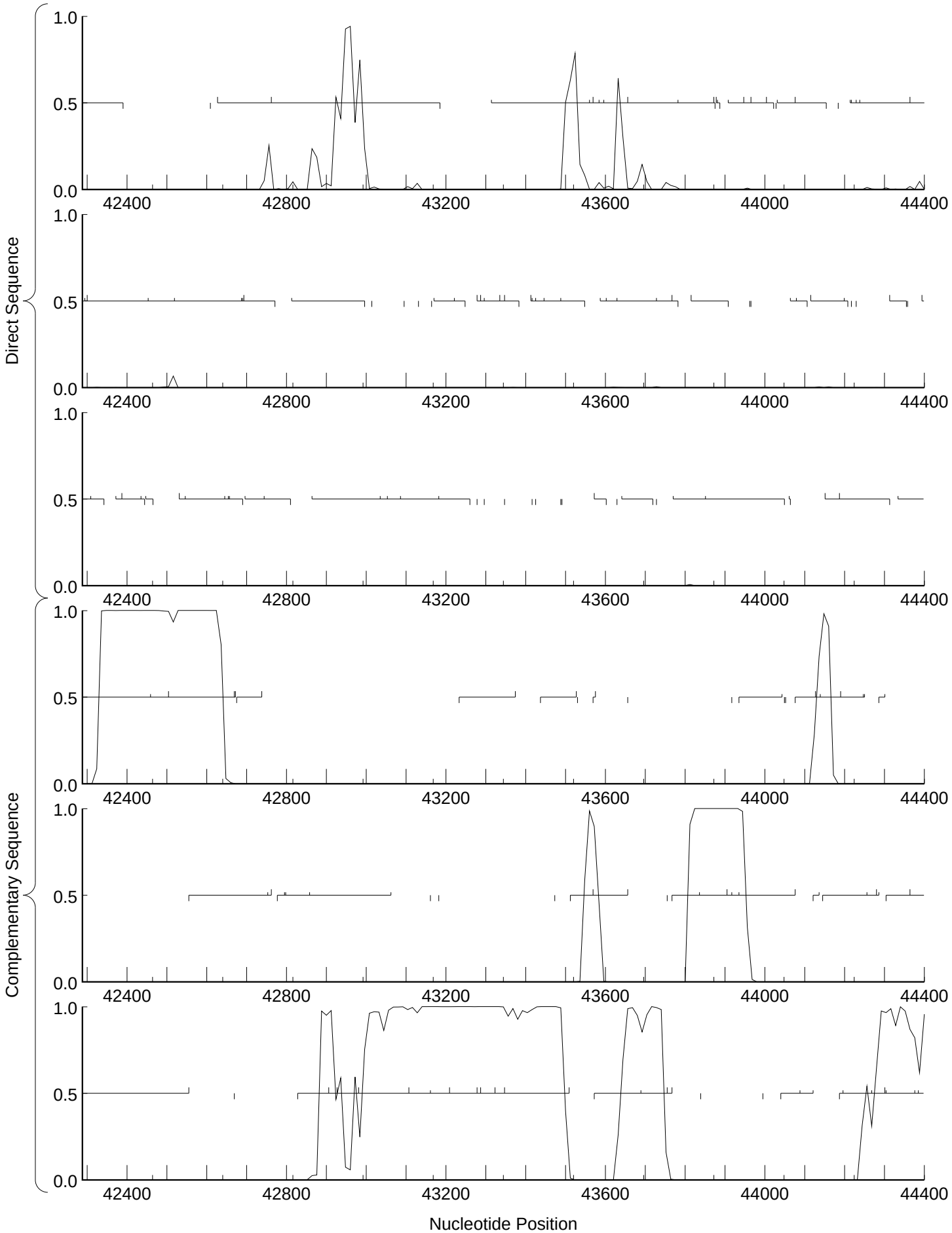


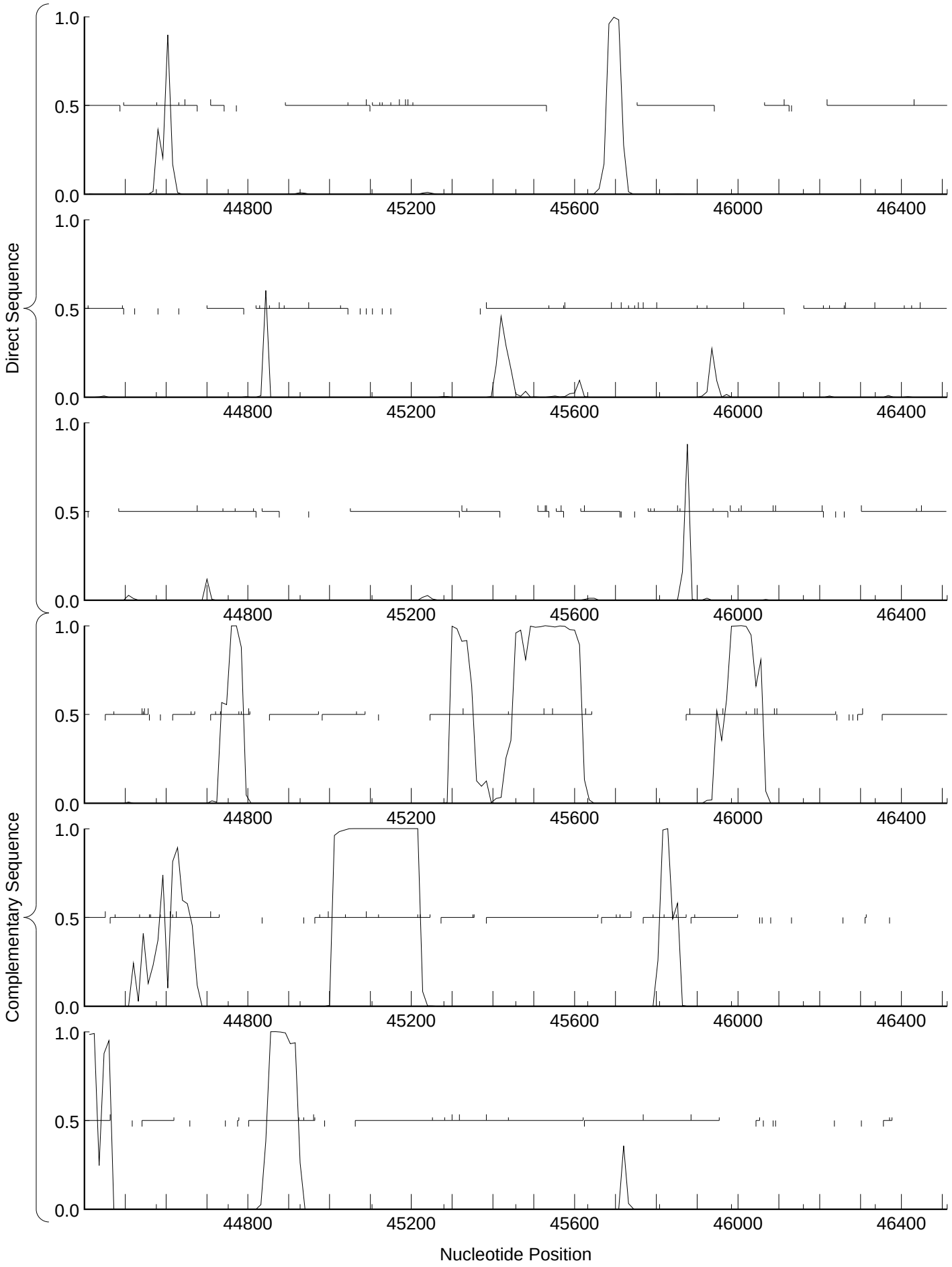
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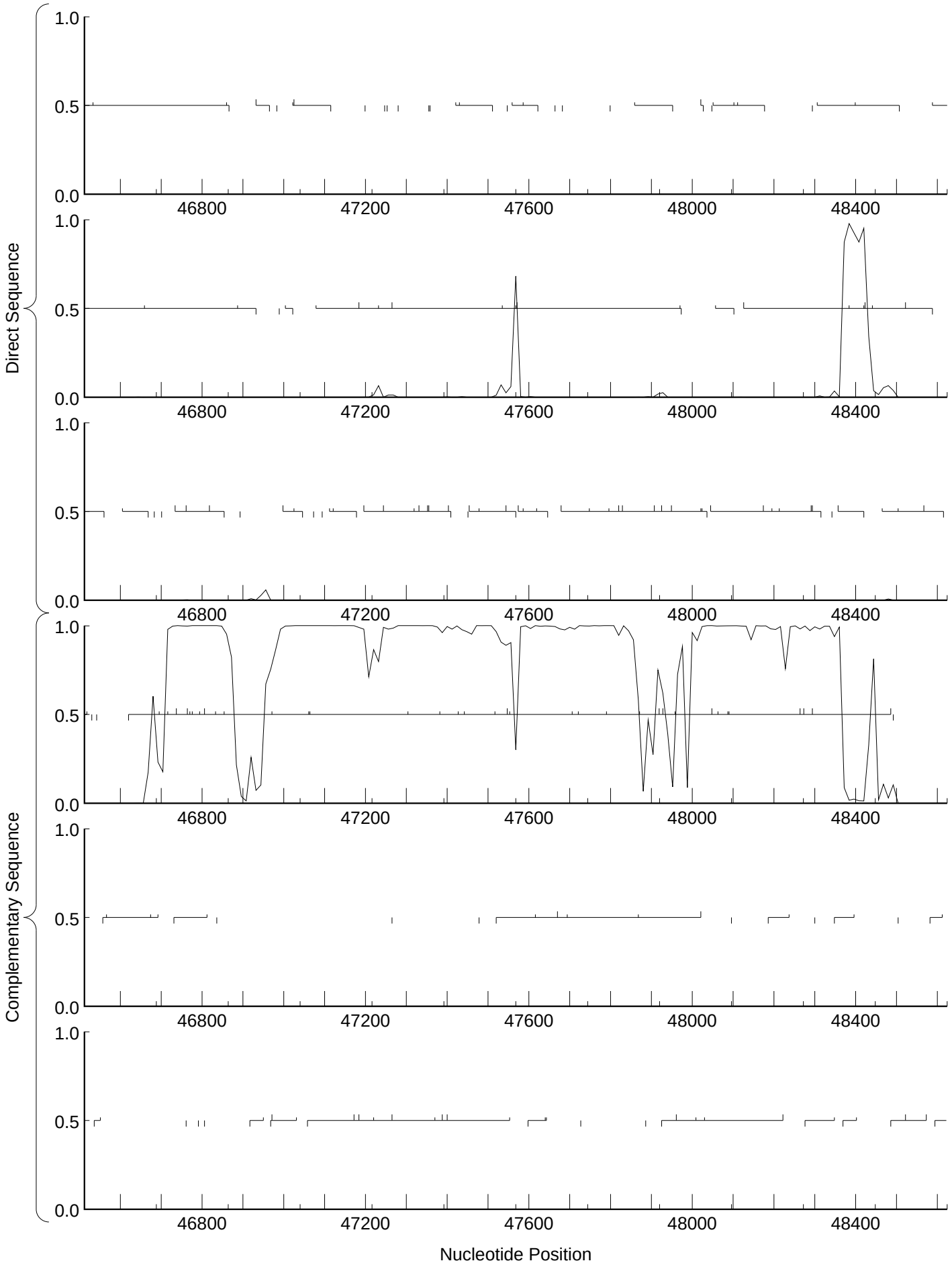
n phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 21/31

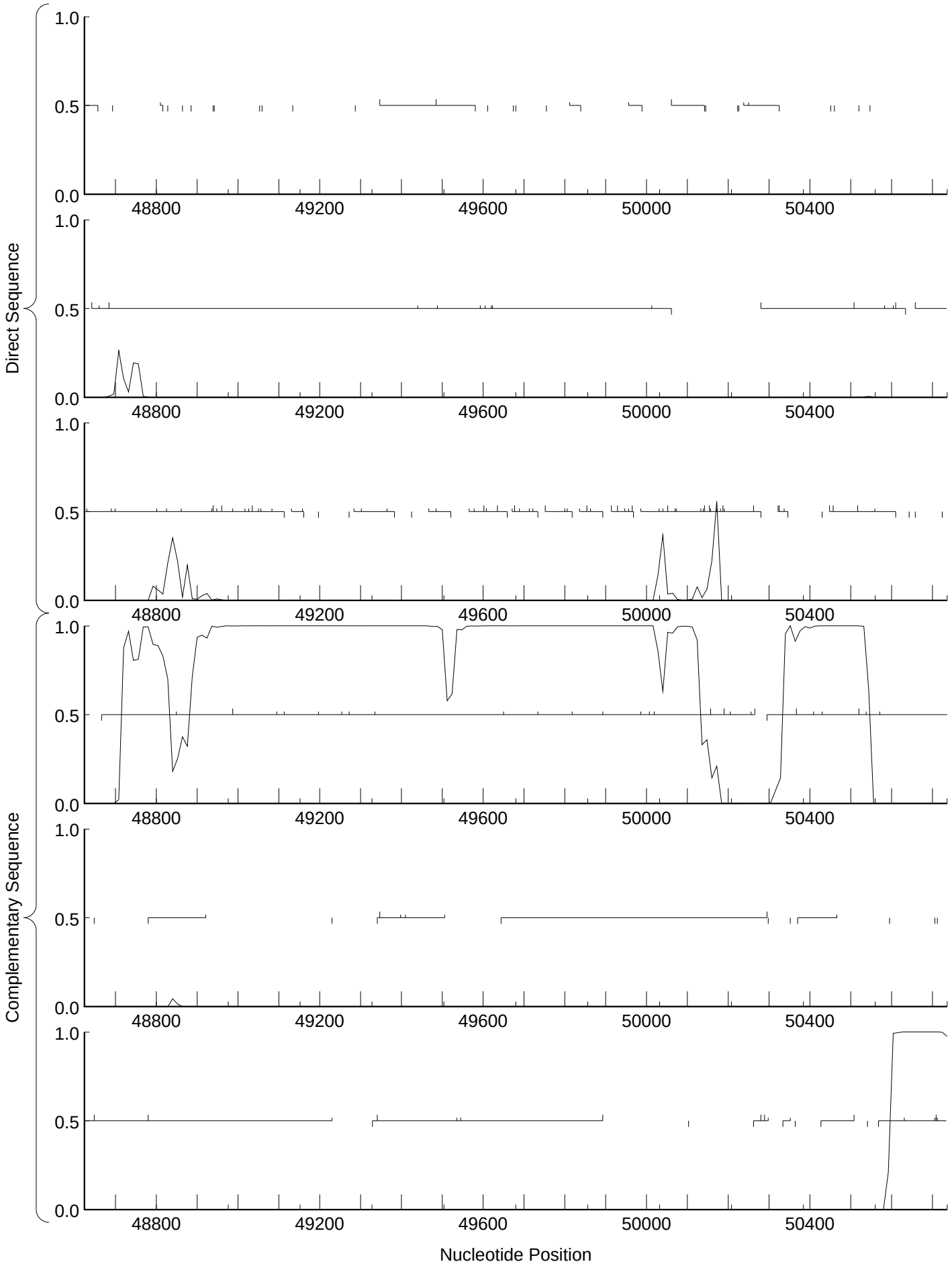



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
42400
42800
43200
43600
44000
44400
Nucleotide Position

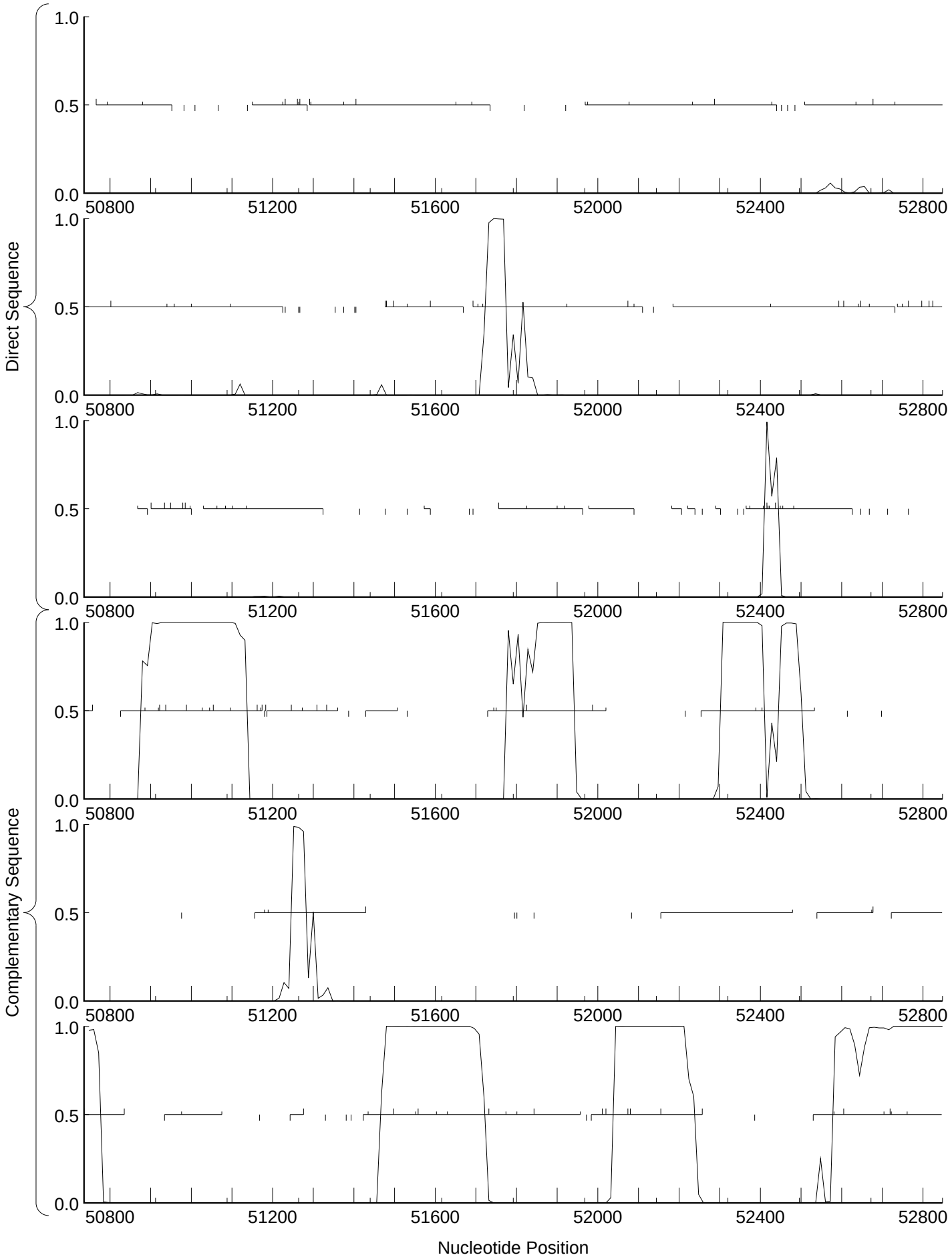

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
44800
45200
45600
46000
46400
Nucleotide Position

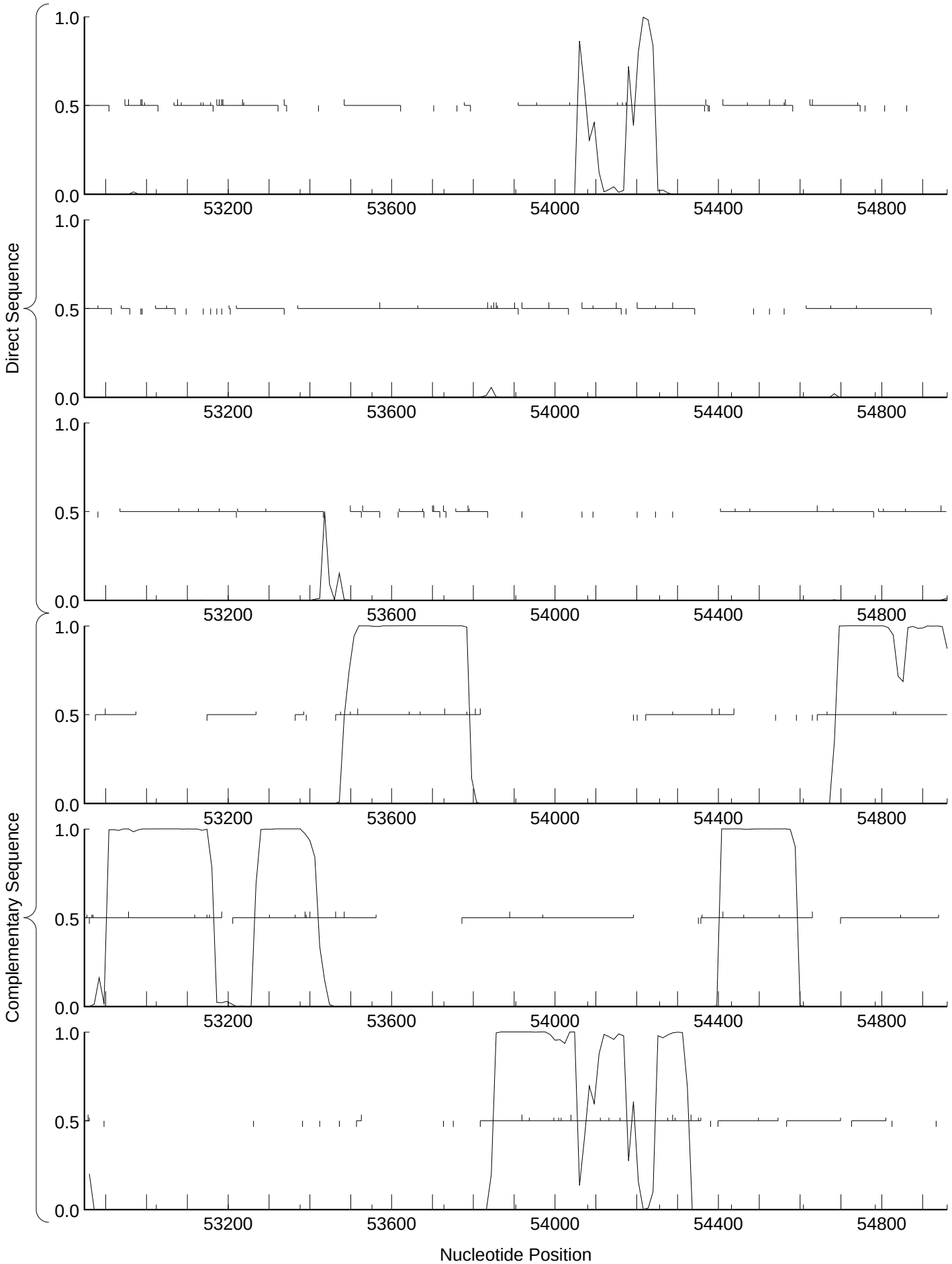
n phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 24/31

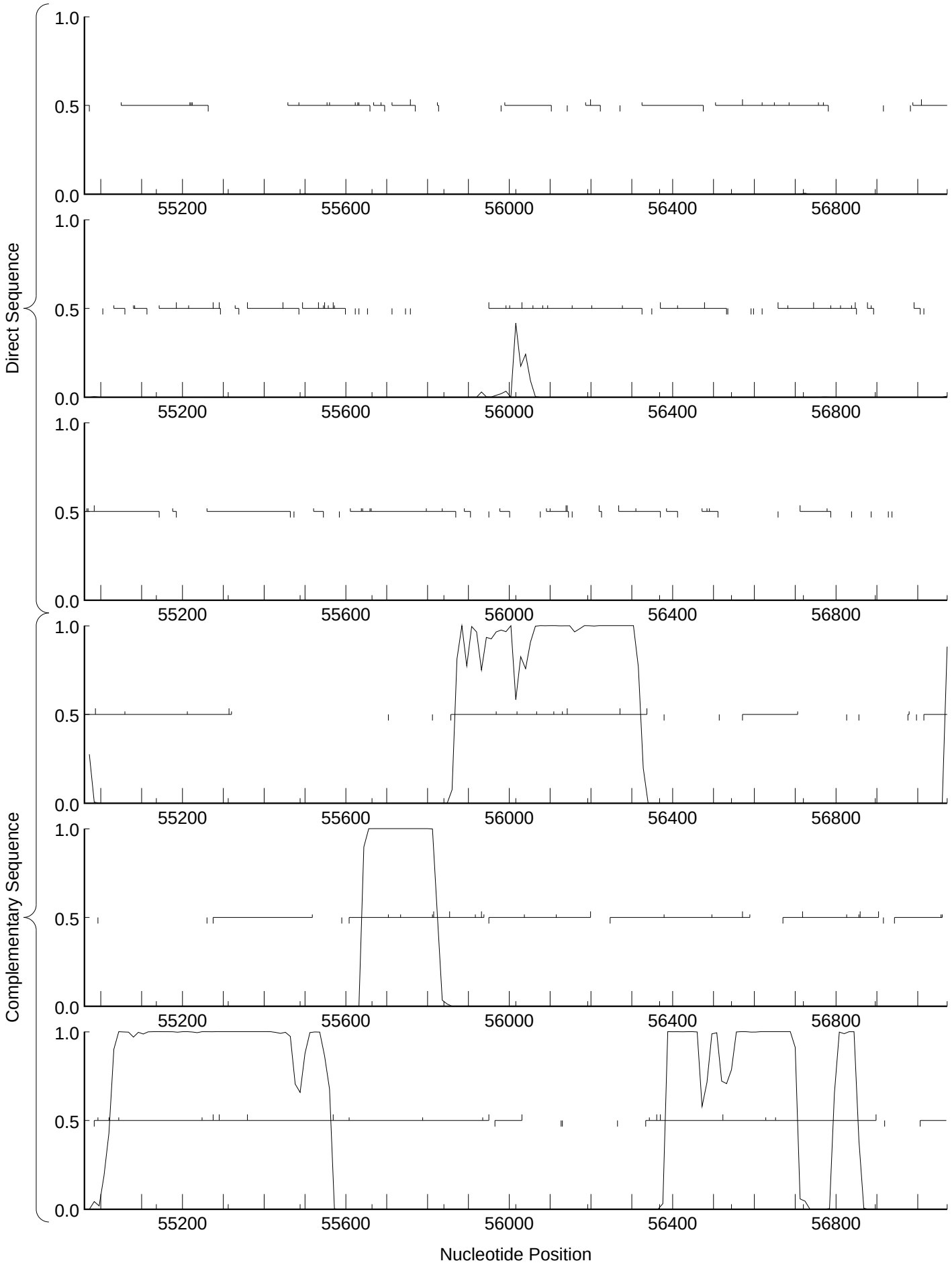


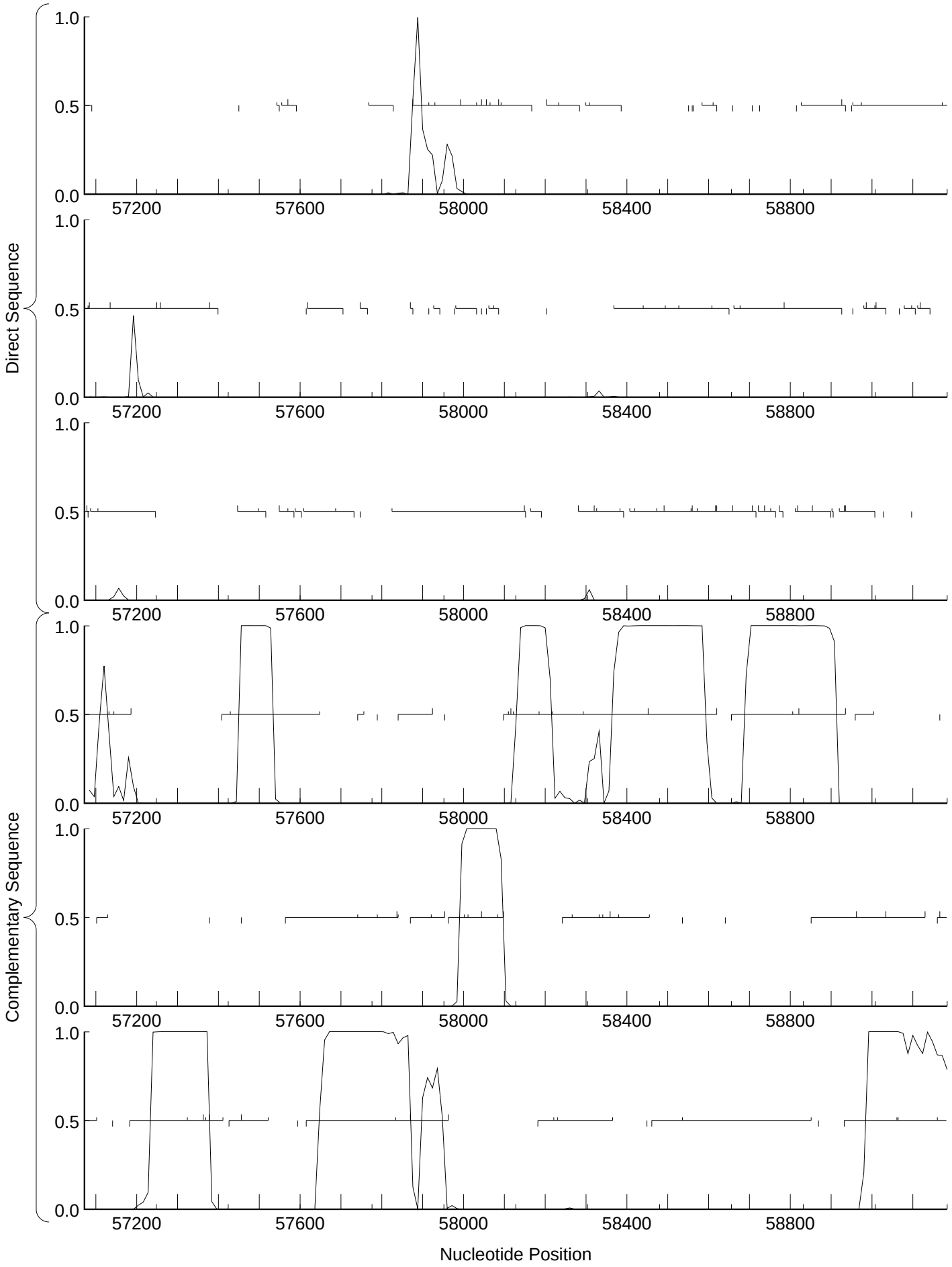

Direct Sequence
Complementary Sequence
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0.5
0.0
48800
49200
49600
50000
50400
Nucleotide Position

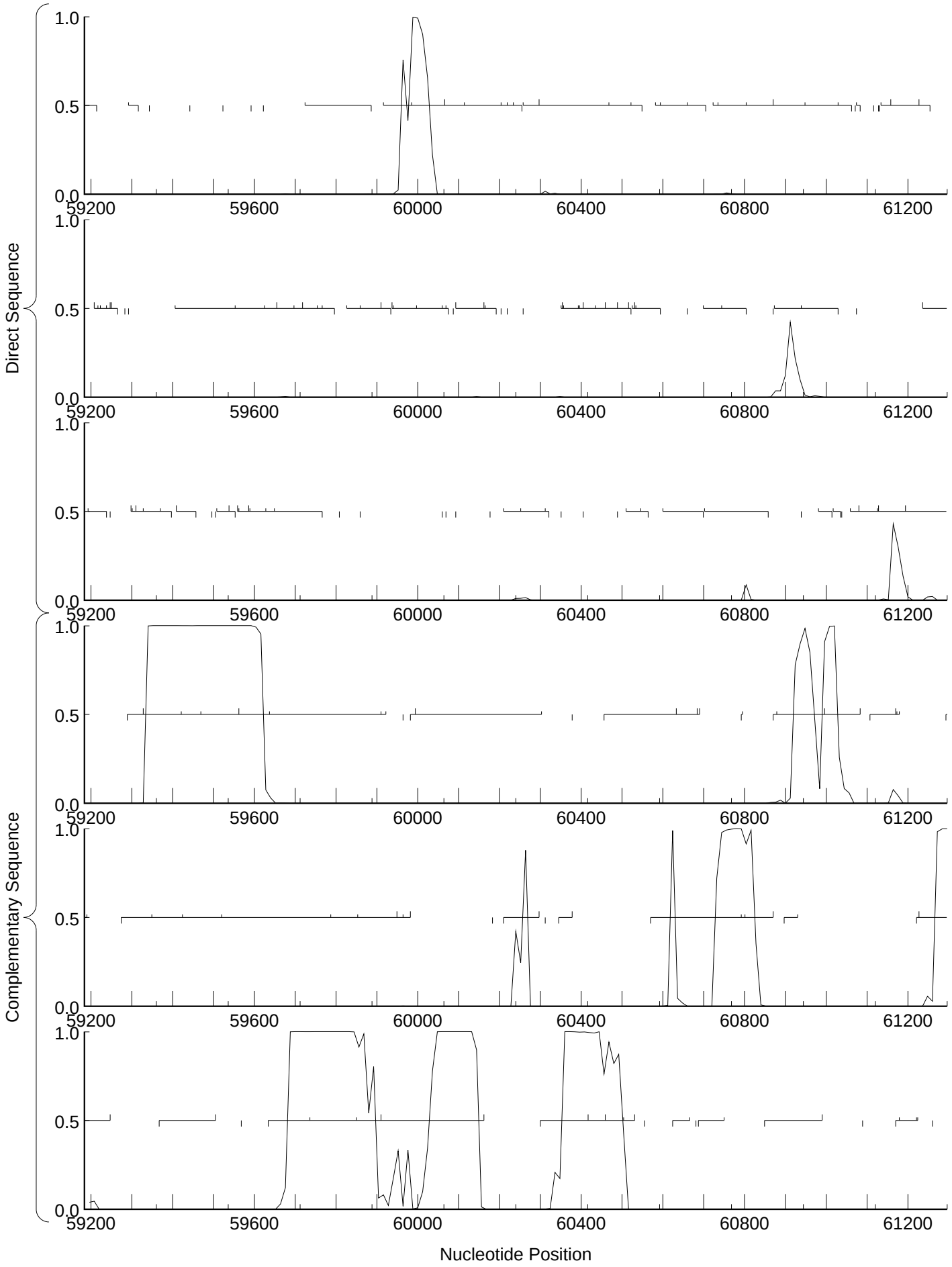
n phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 26/31




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

n phage Gazebo complete sequence, 62358 bp including 200 bp terminal repeat, Cluster EG, Order 4, Window 96, Step 12, 31/31

