GeneMark

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

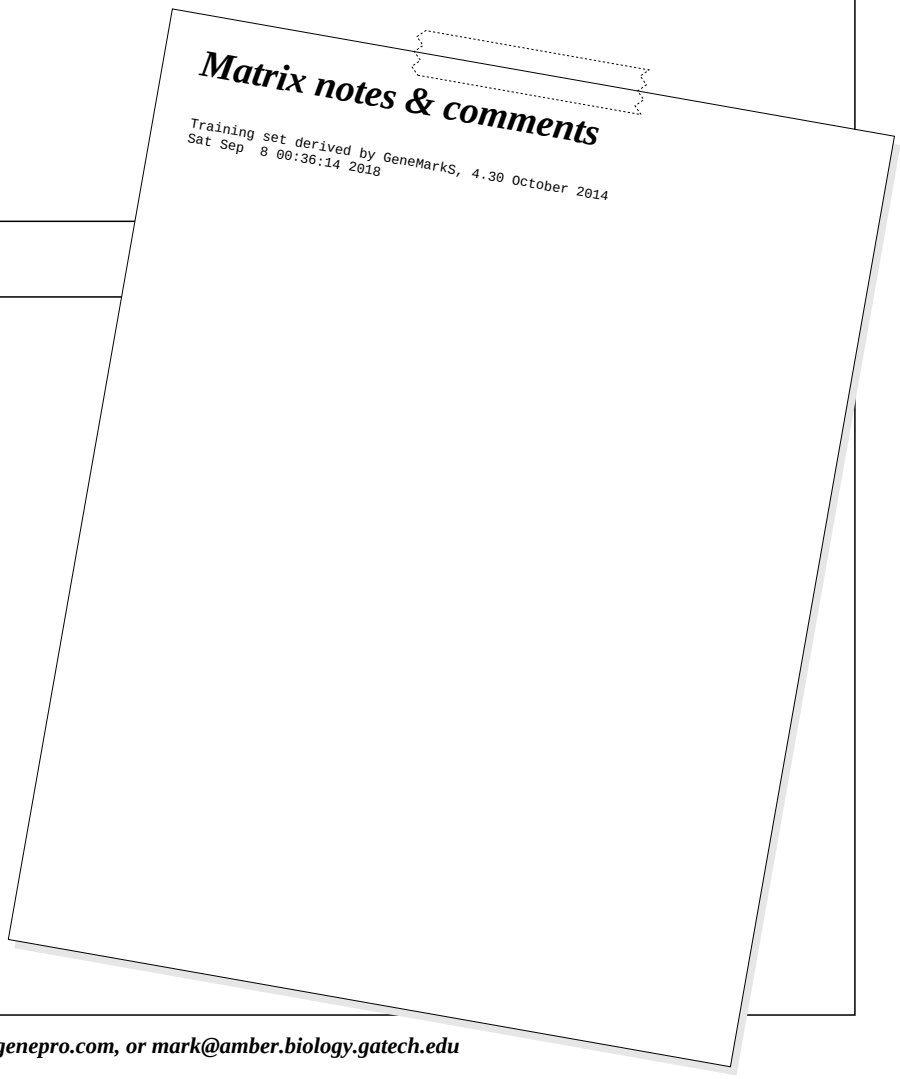
PROGRAM INFORMATION

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Sequence         :  Gordonia phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster
Analysis Date    :  3/29/24 at 5:26:25
Pages            :  38
Sequence Length  :  77394 bp
GC Content       :   58.88%

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

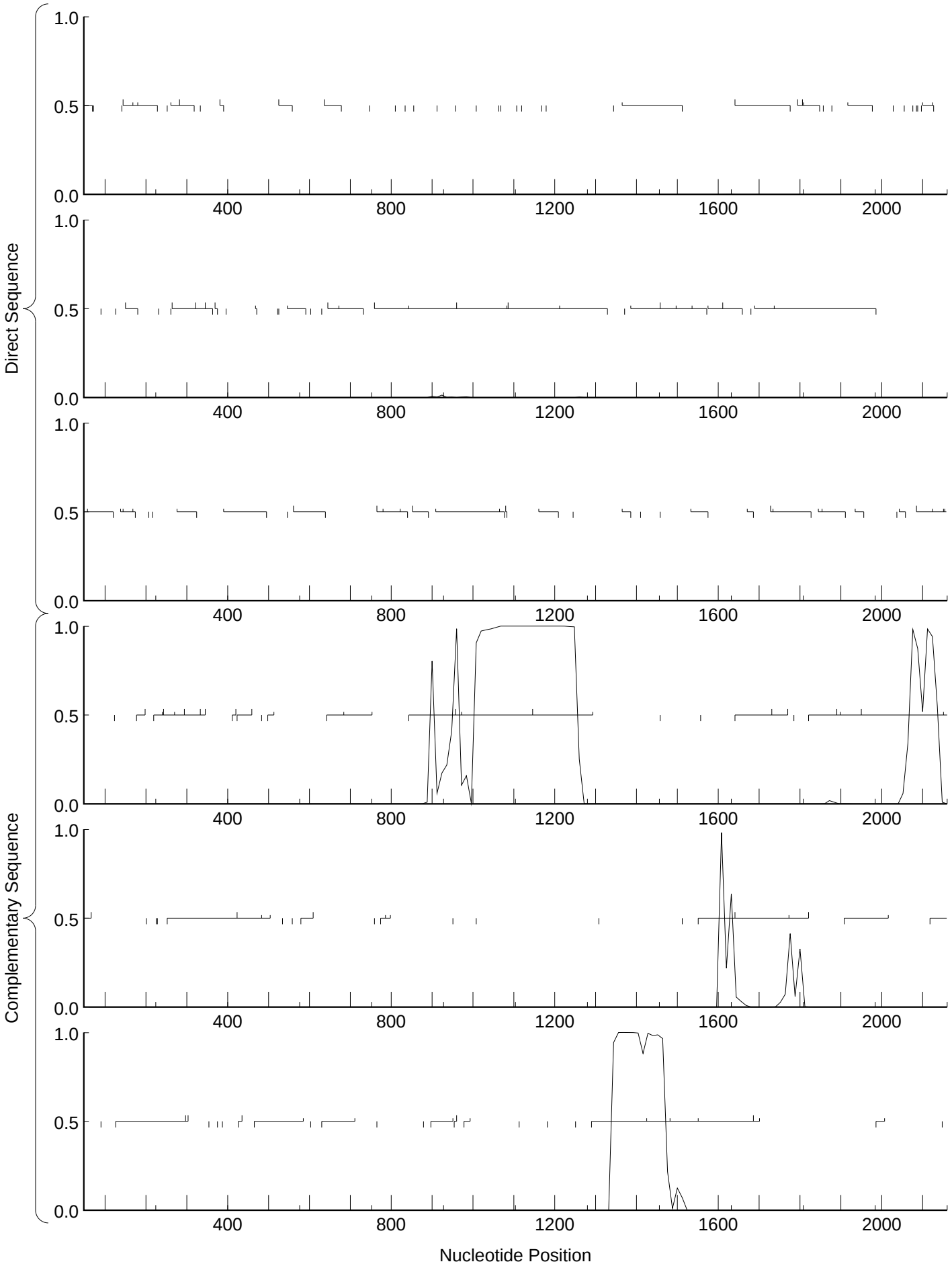
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Matrix     :  -
Author     :  -

Order      :  4
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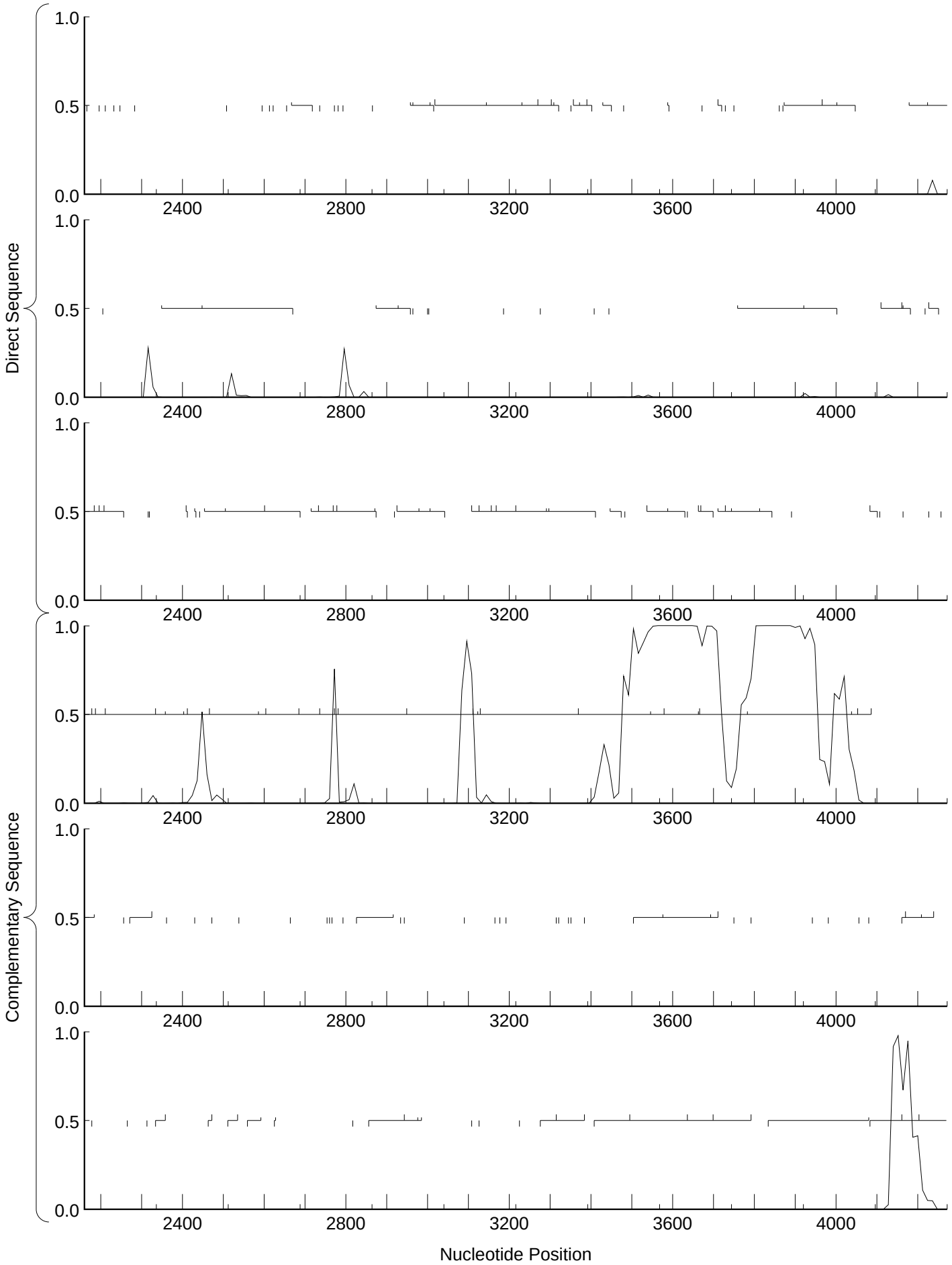

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

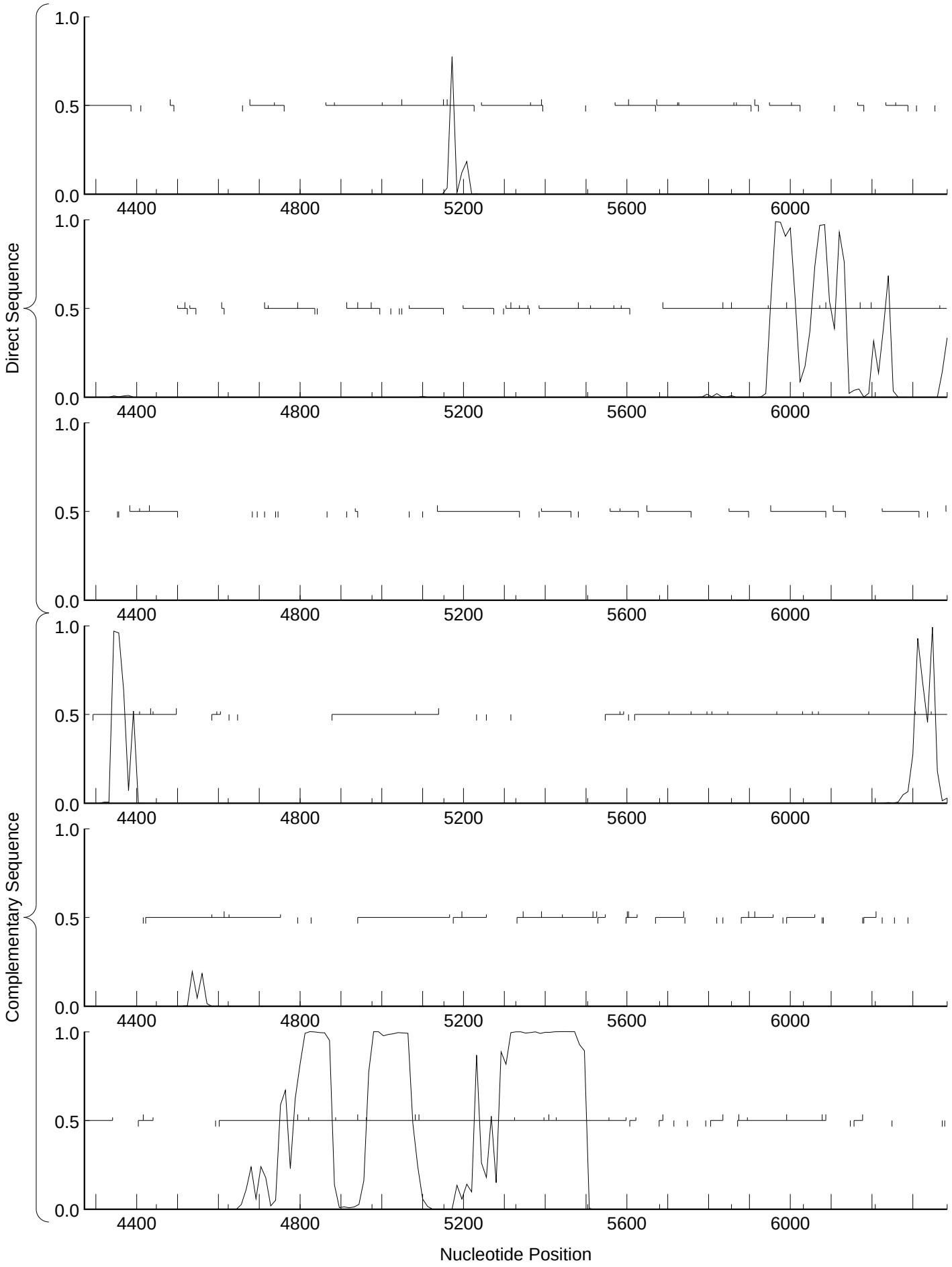
a phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 2/38



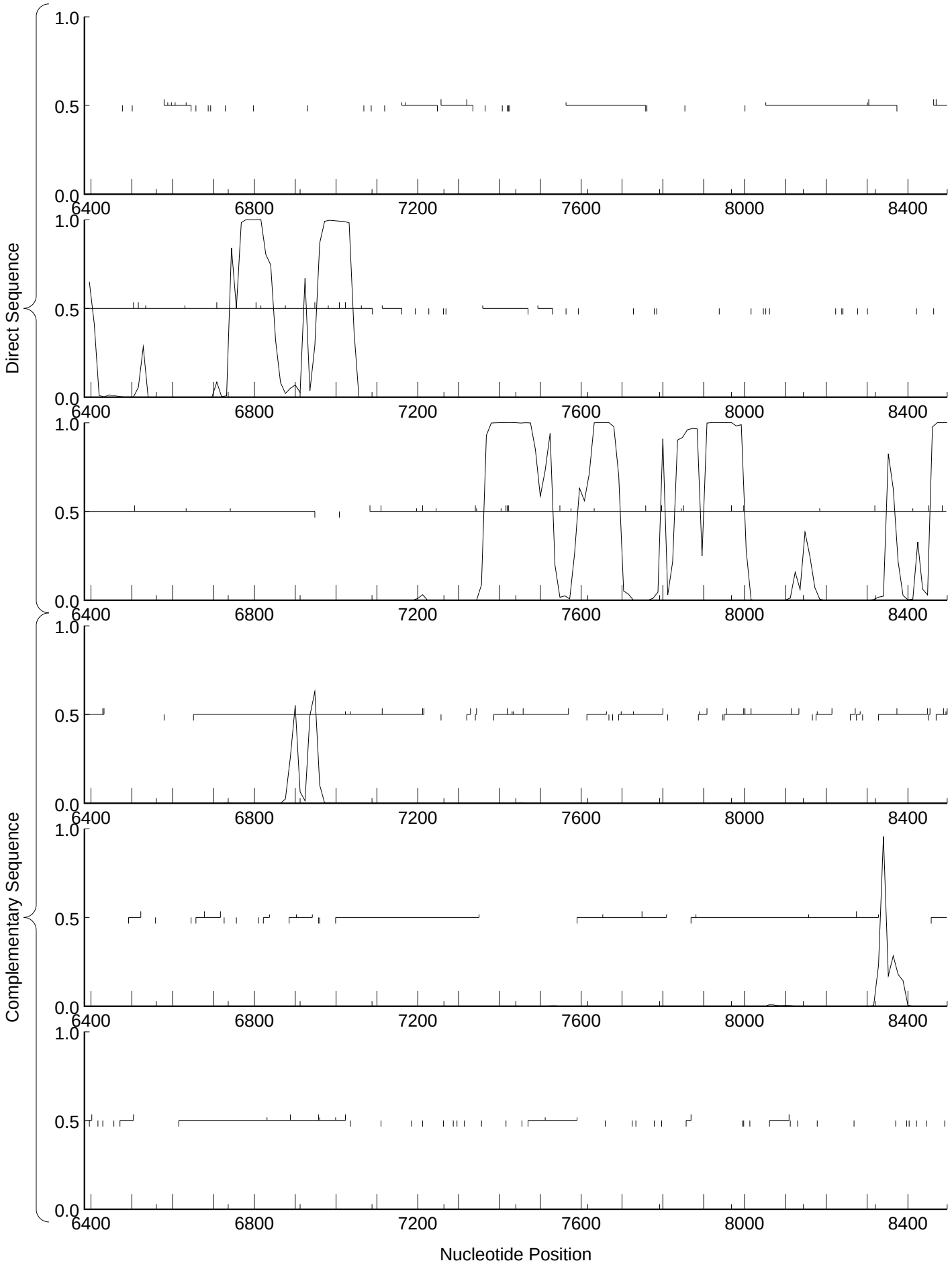
a phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 3/38



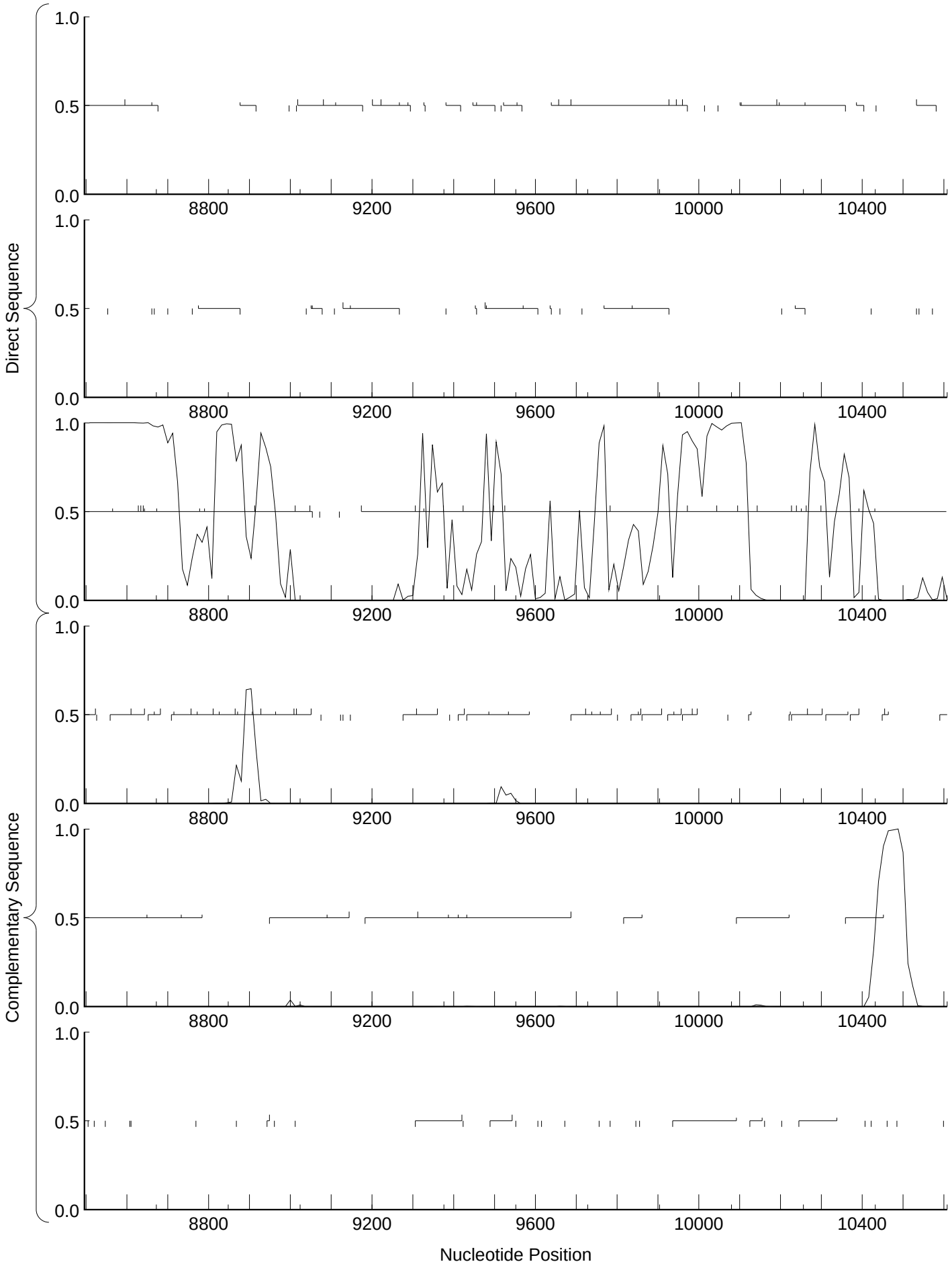
a phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 4/38



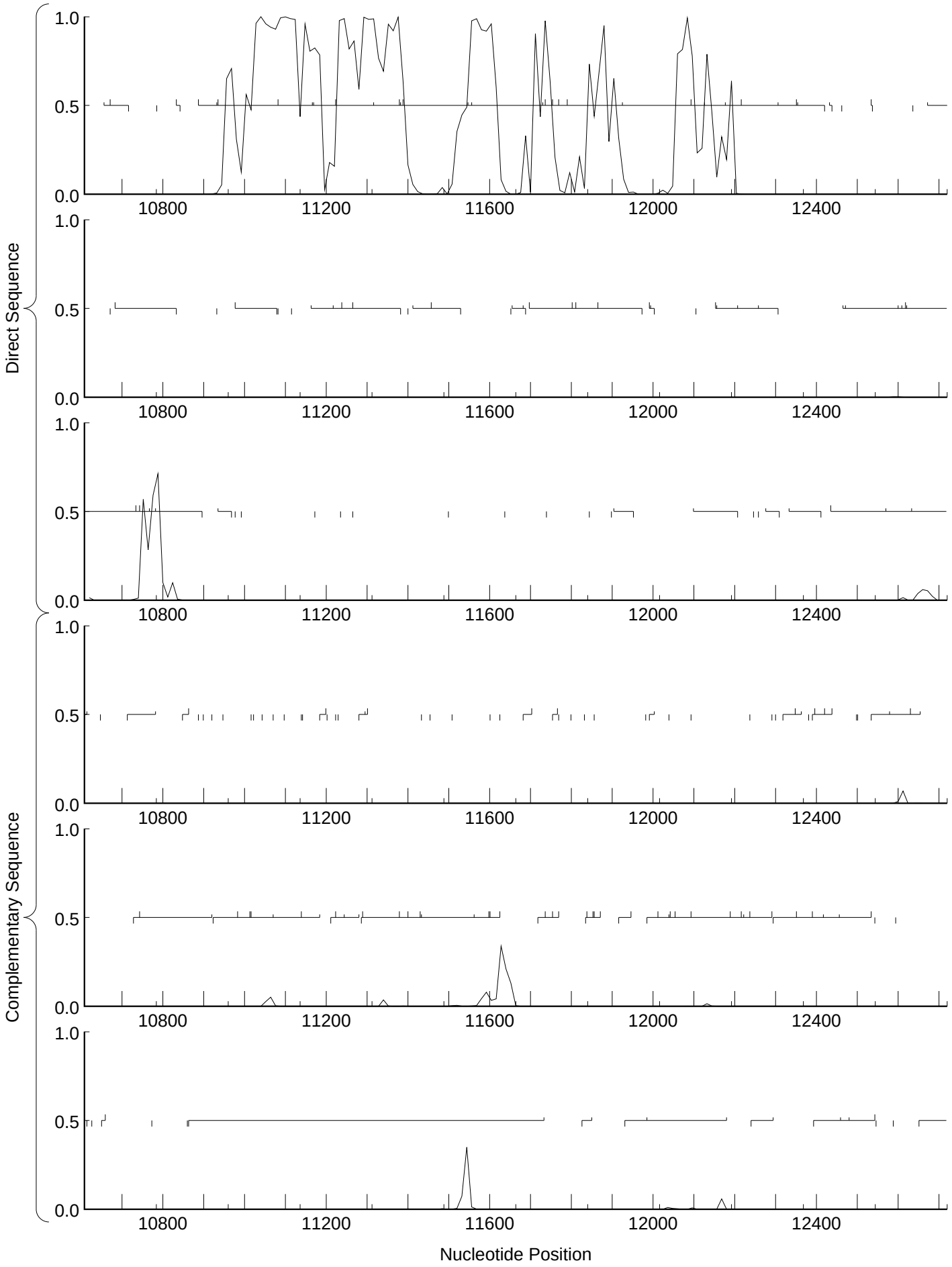
a phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 5/38



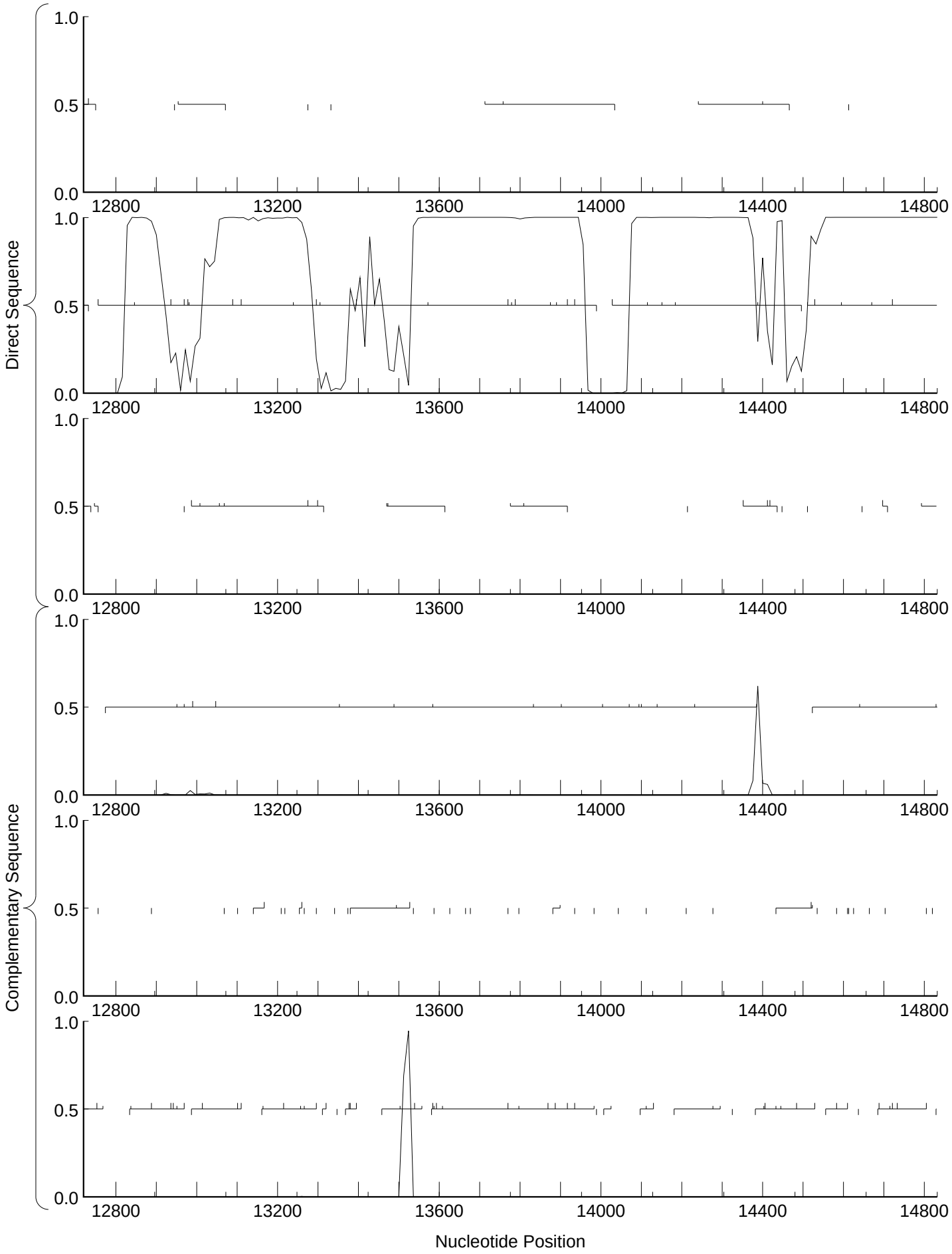
a phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 6/38



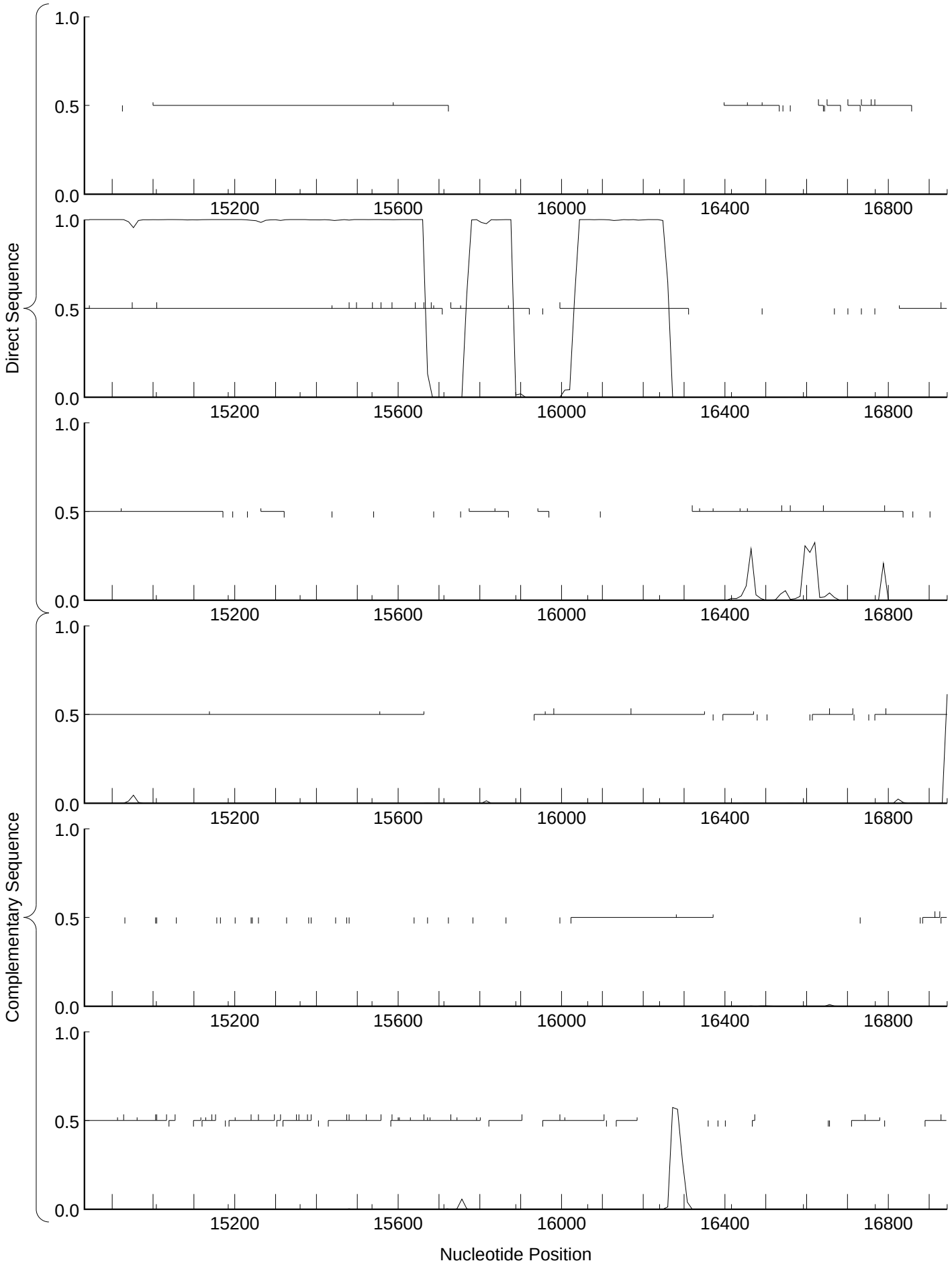
a phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 7/38



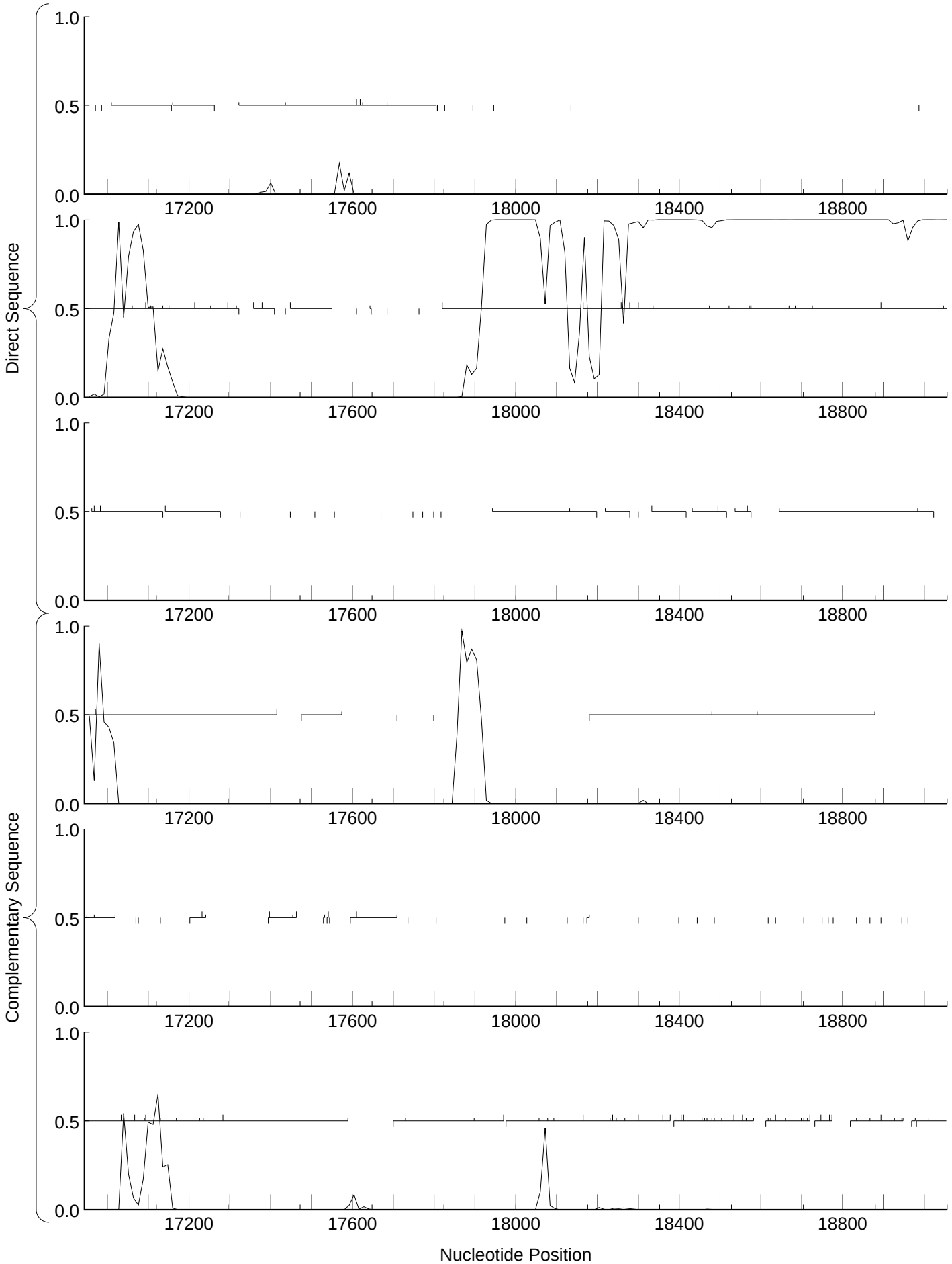
a phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 8/38



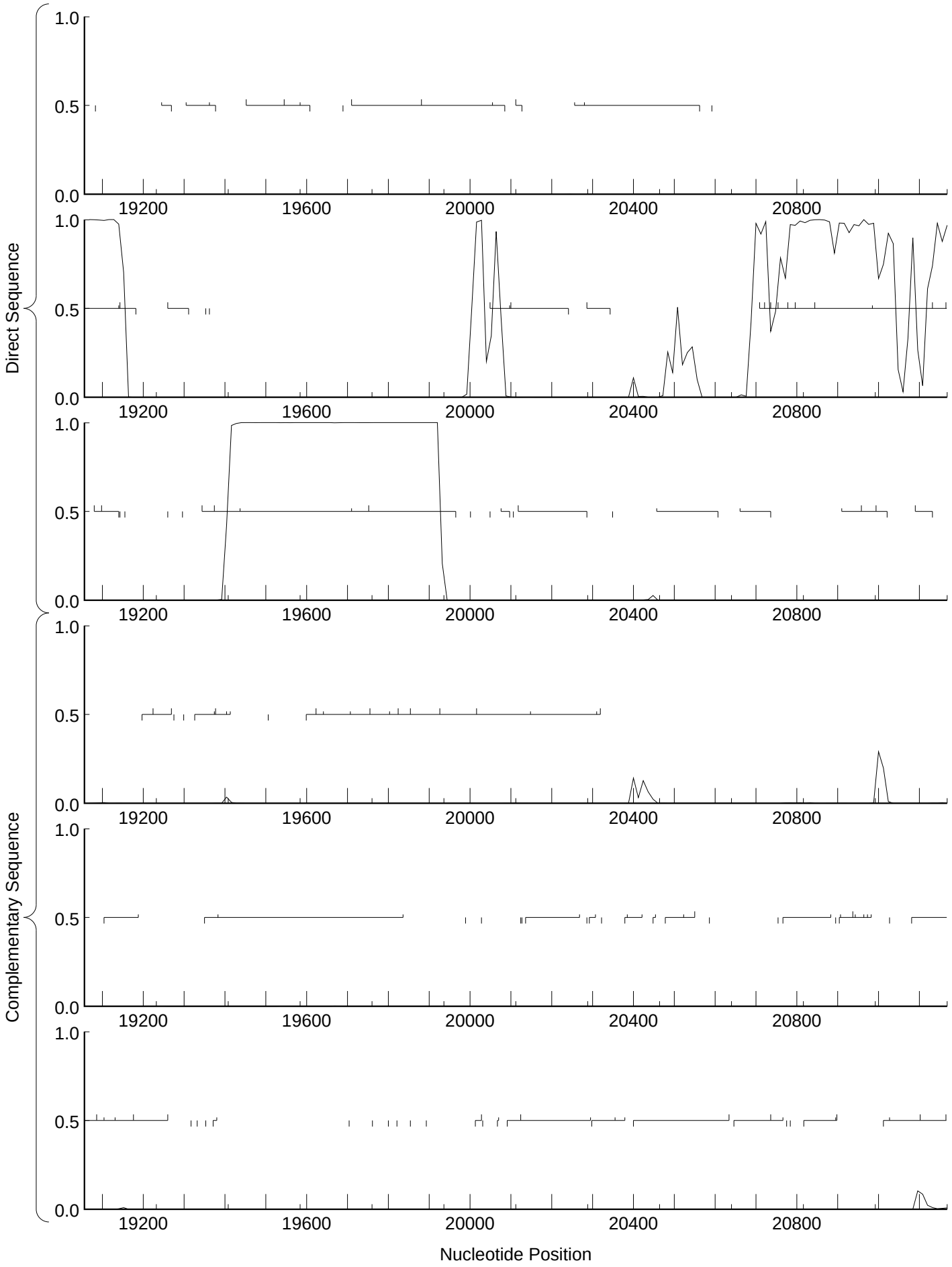
a phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 9/38



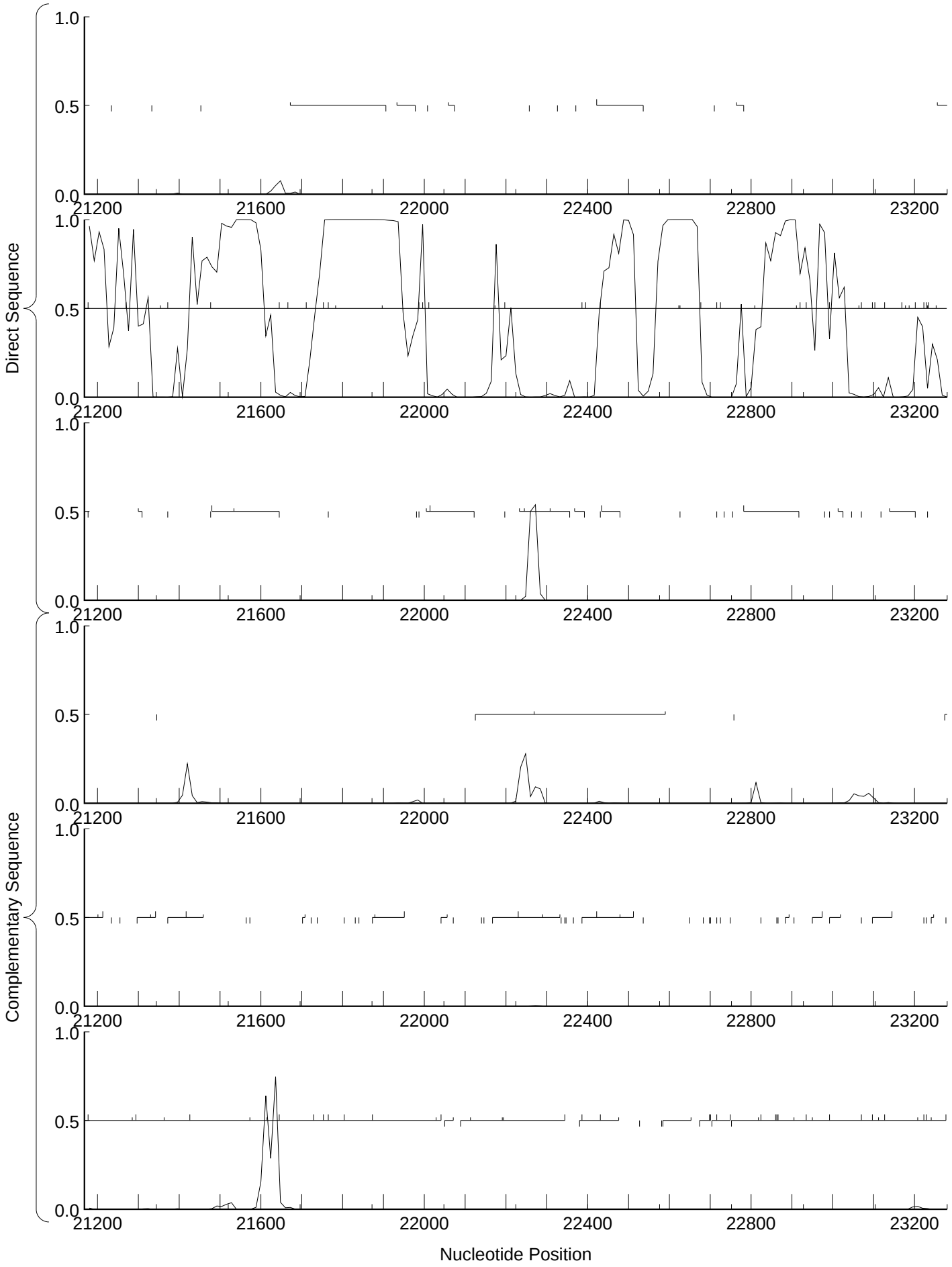
, phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 10/38

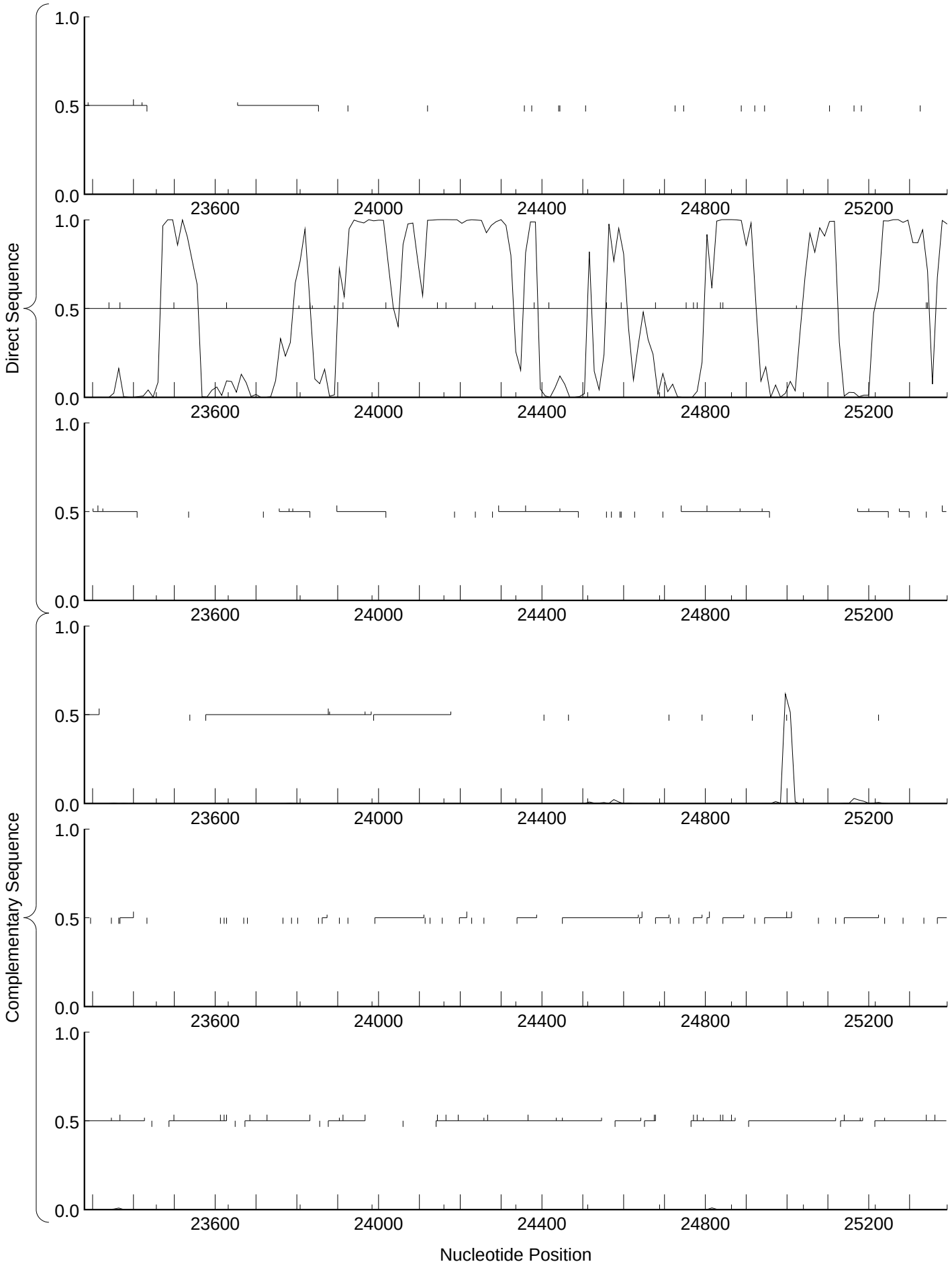


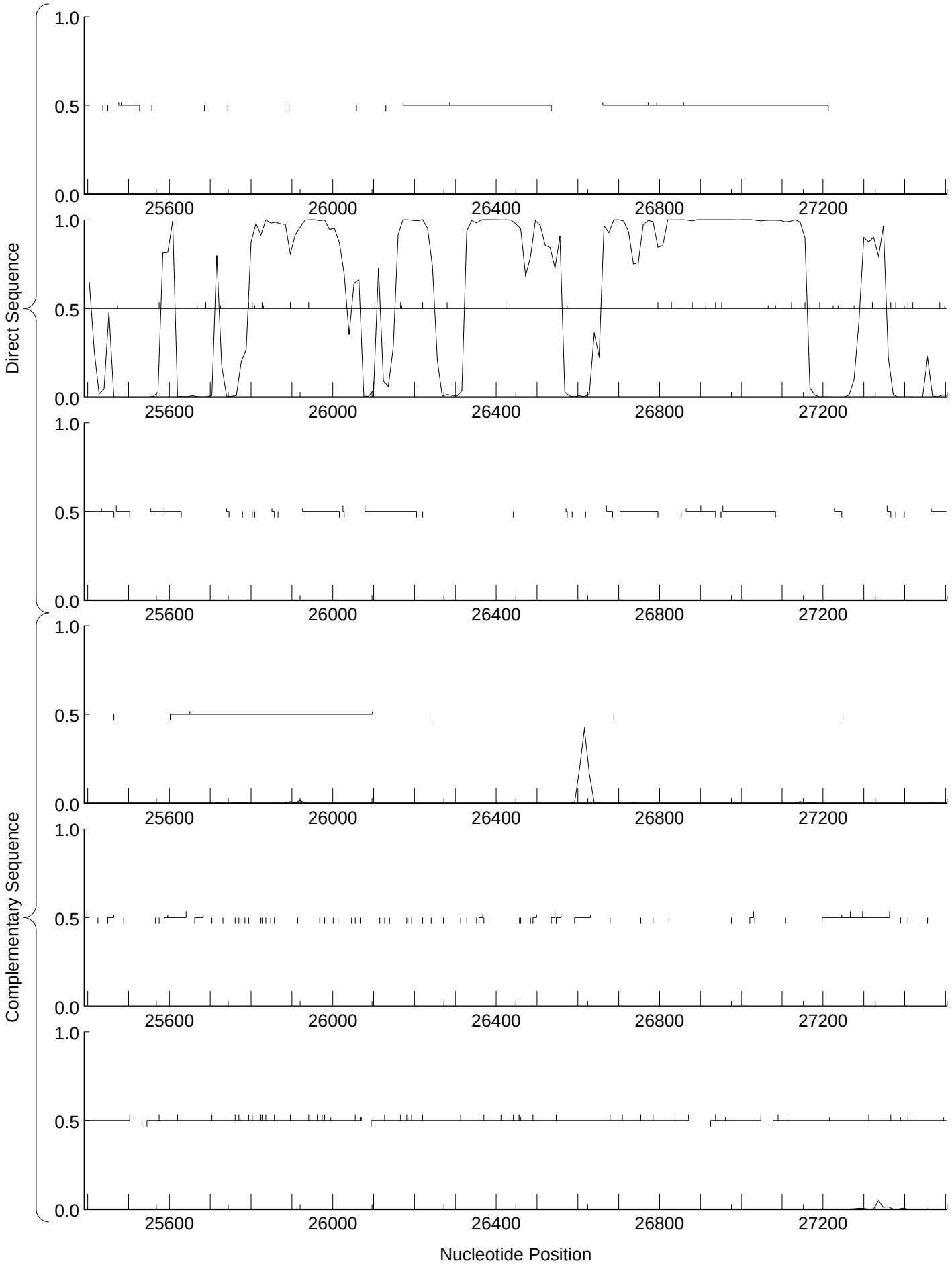
, phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 11/38



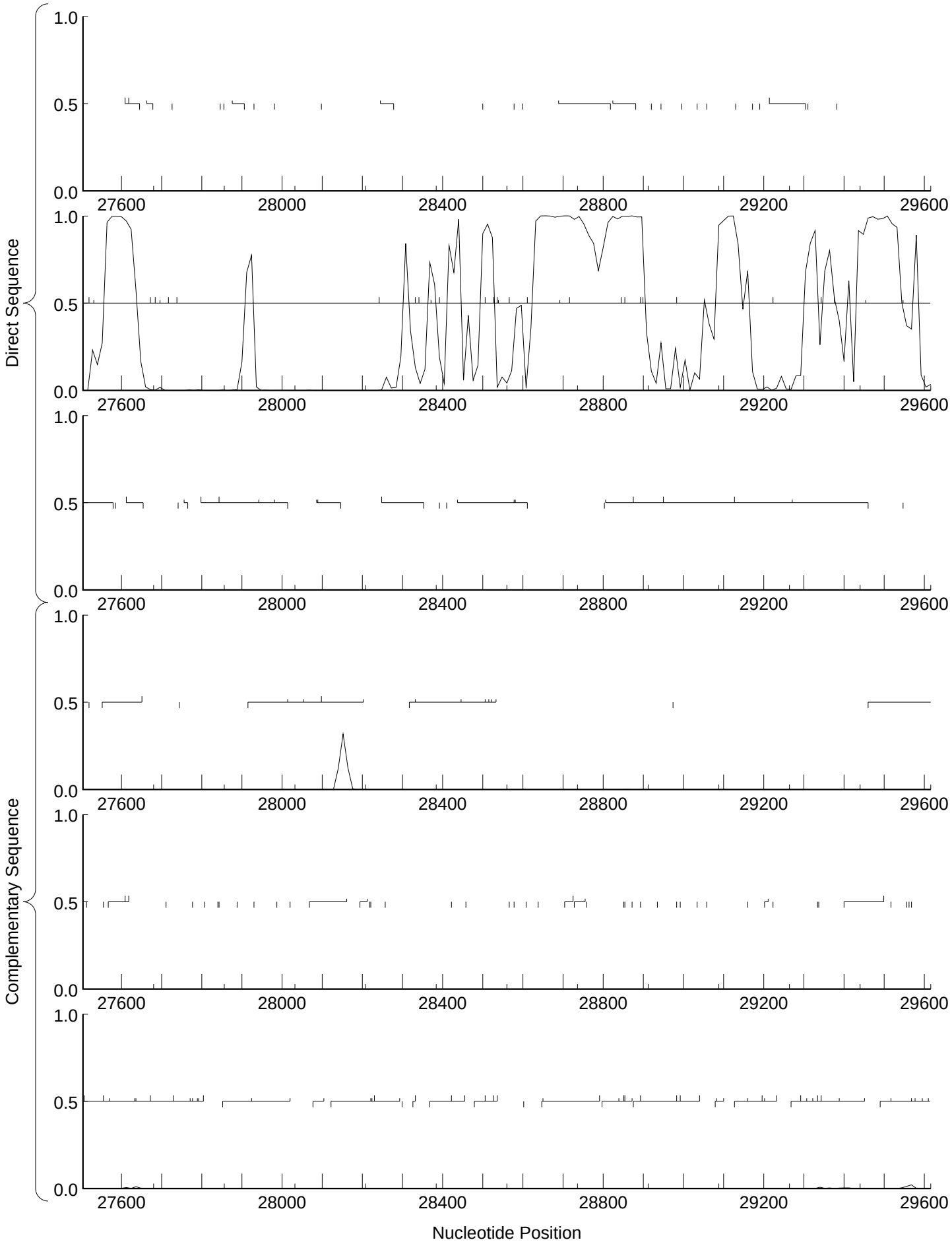
phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 12/38



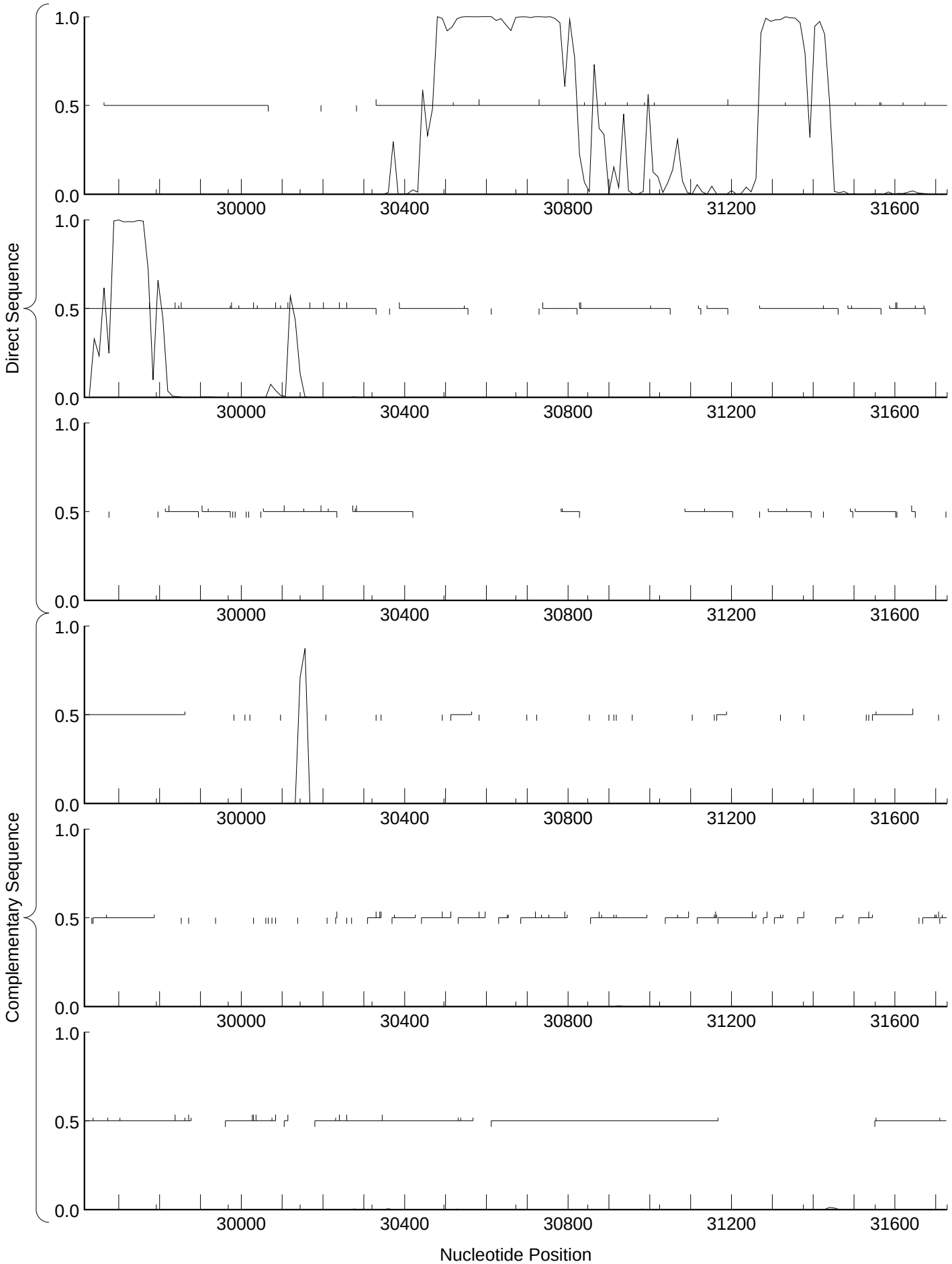

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
23600
24000
24400
24800
25200
Nucleotide Position


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

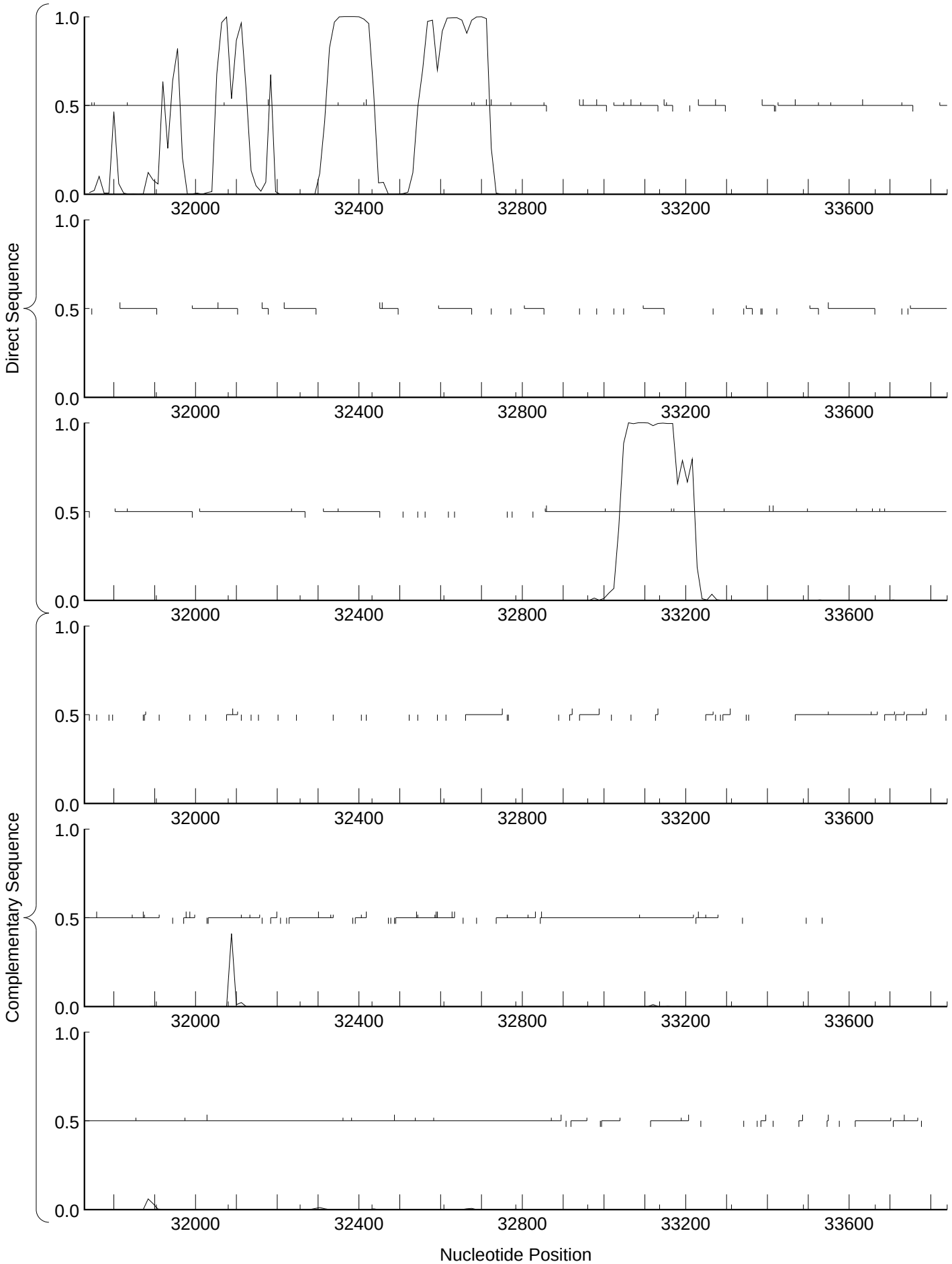
, phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 15/38



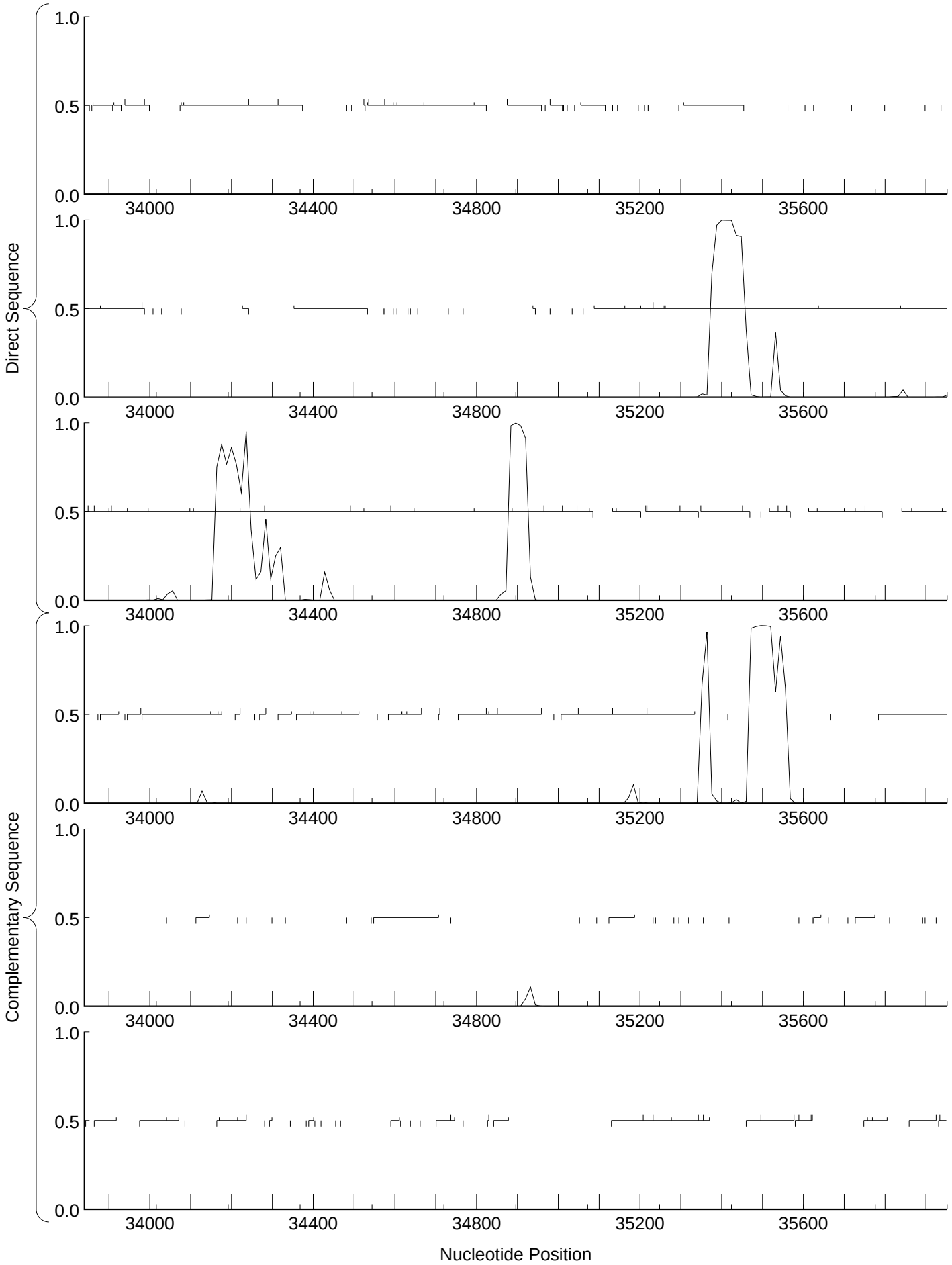
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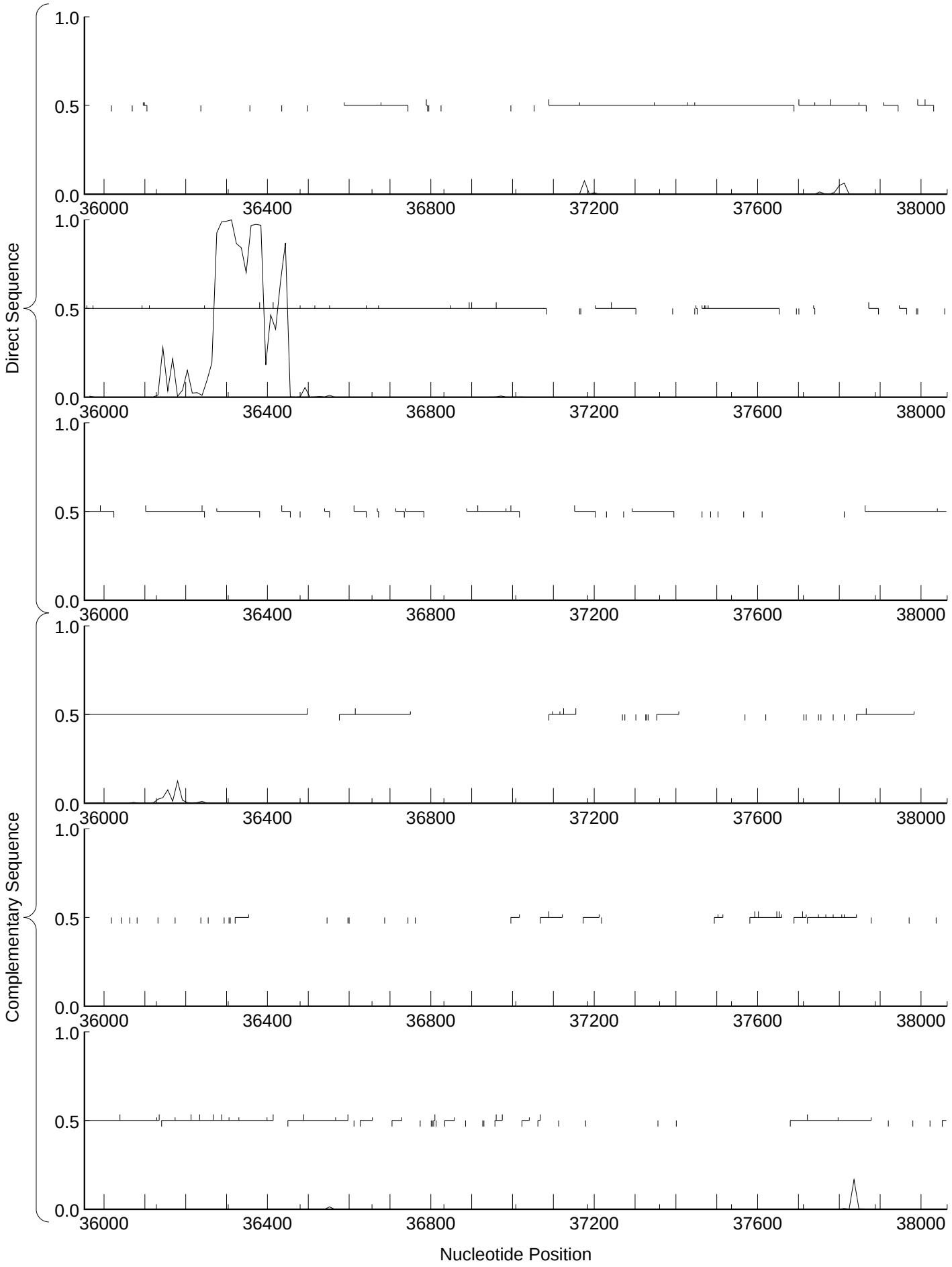
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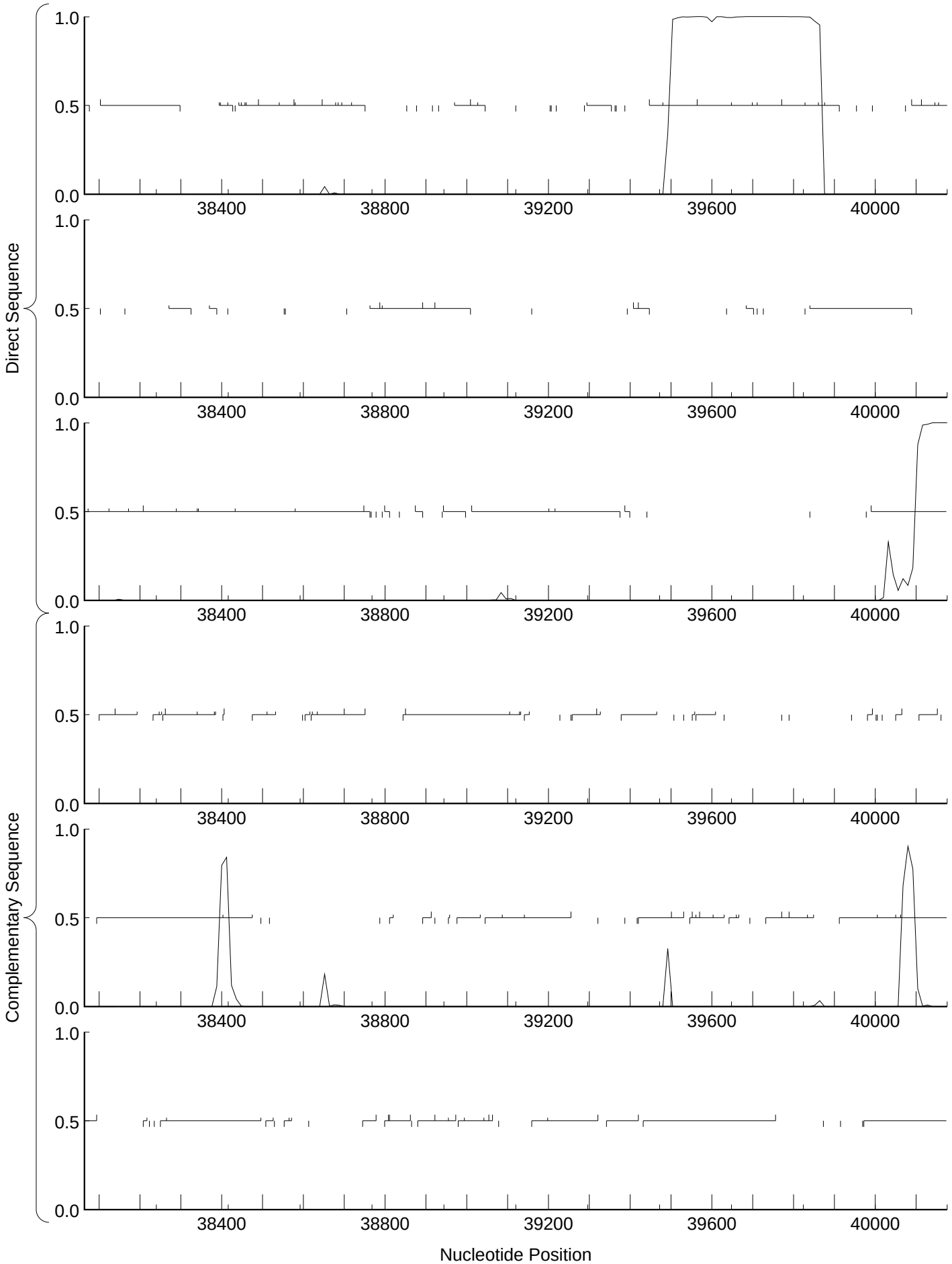


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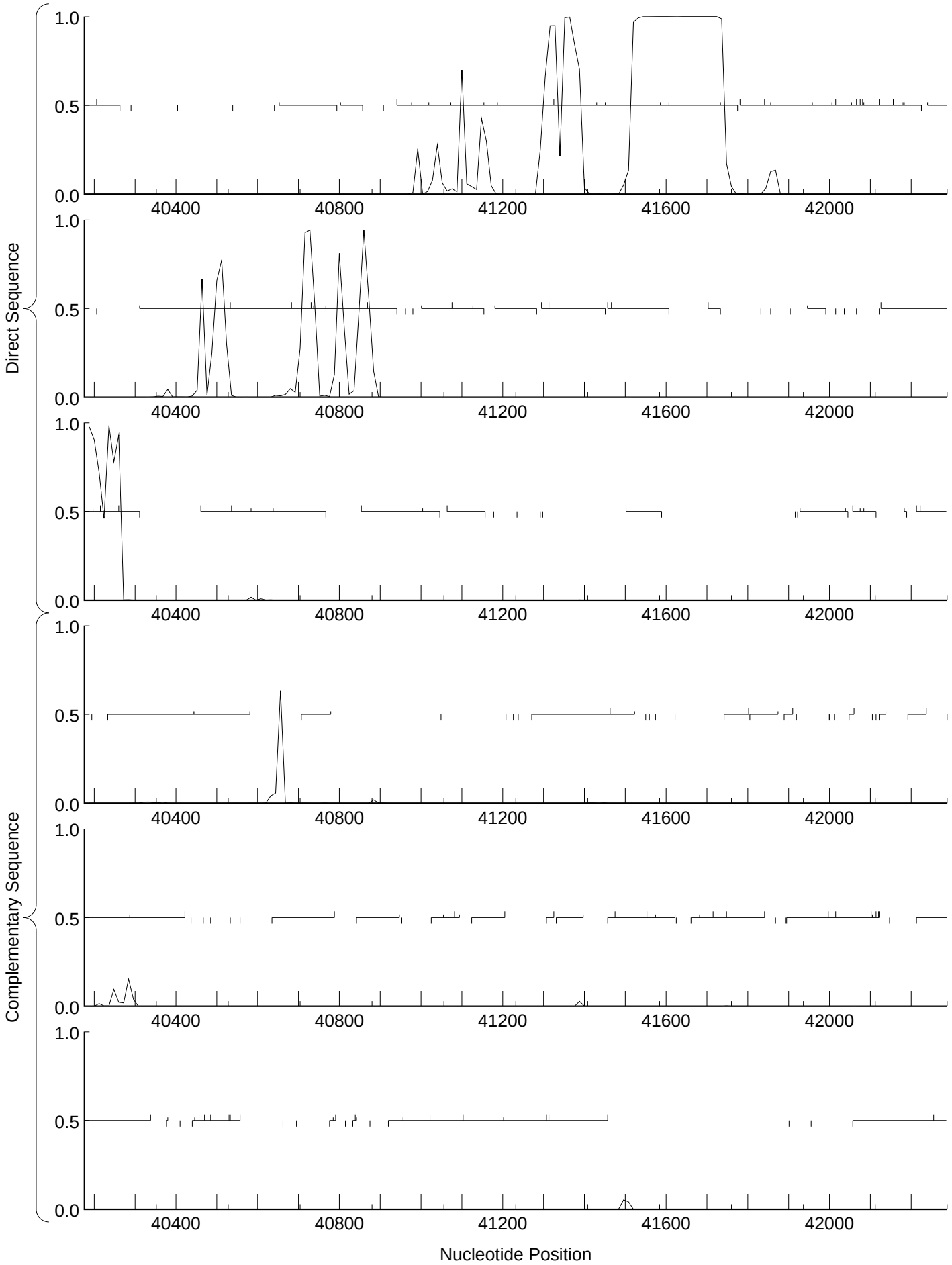


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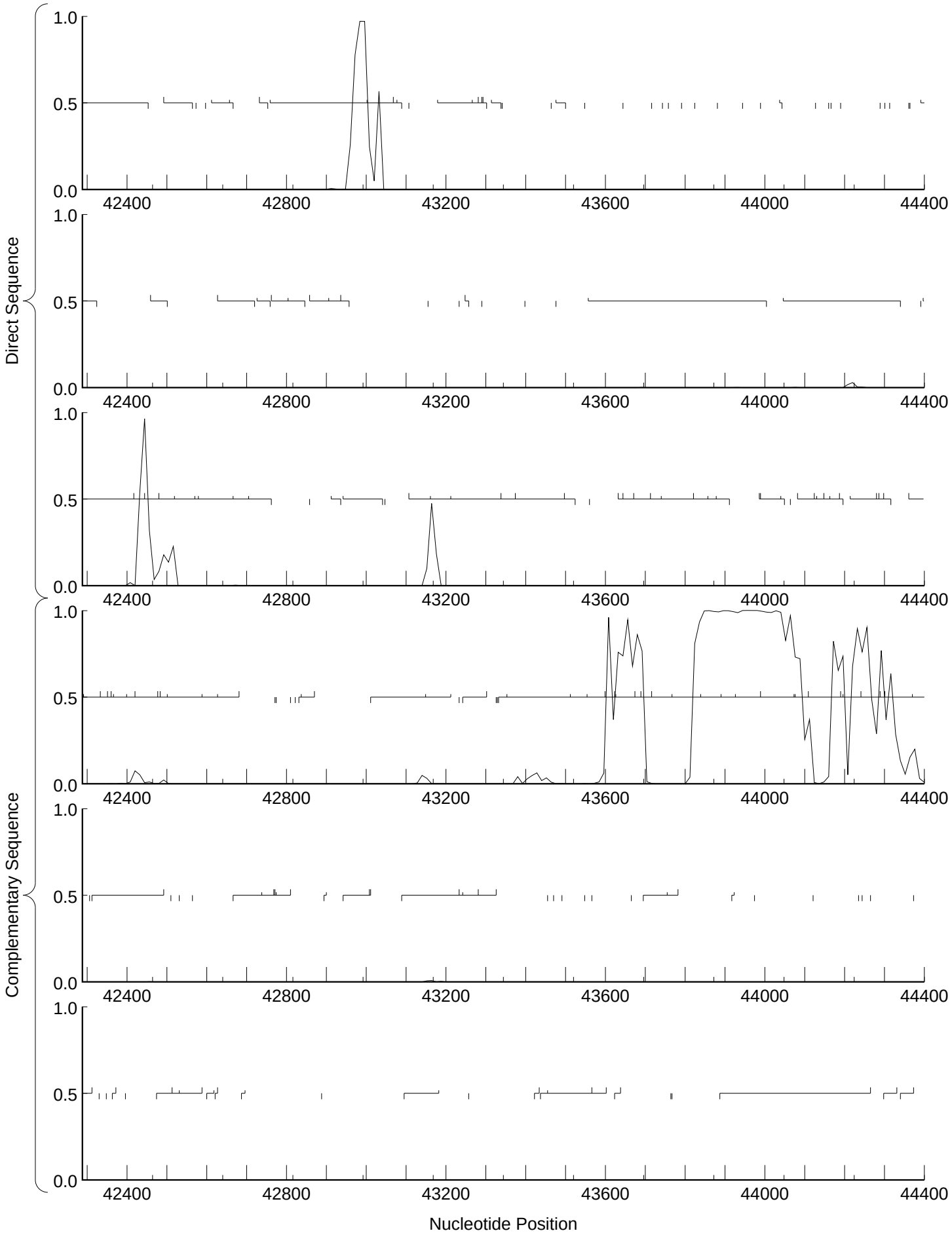


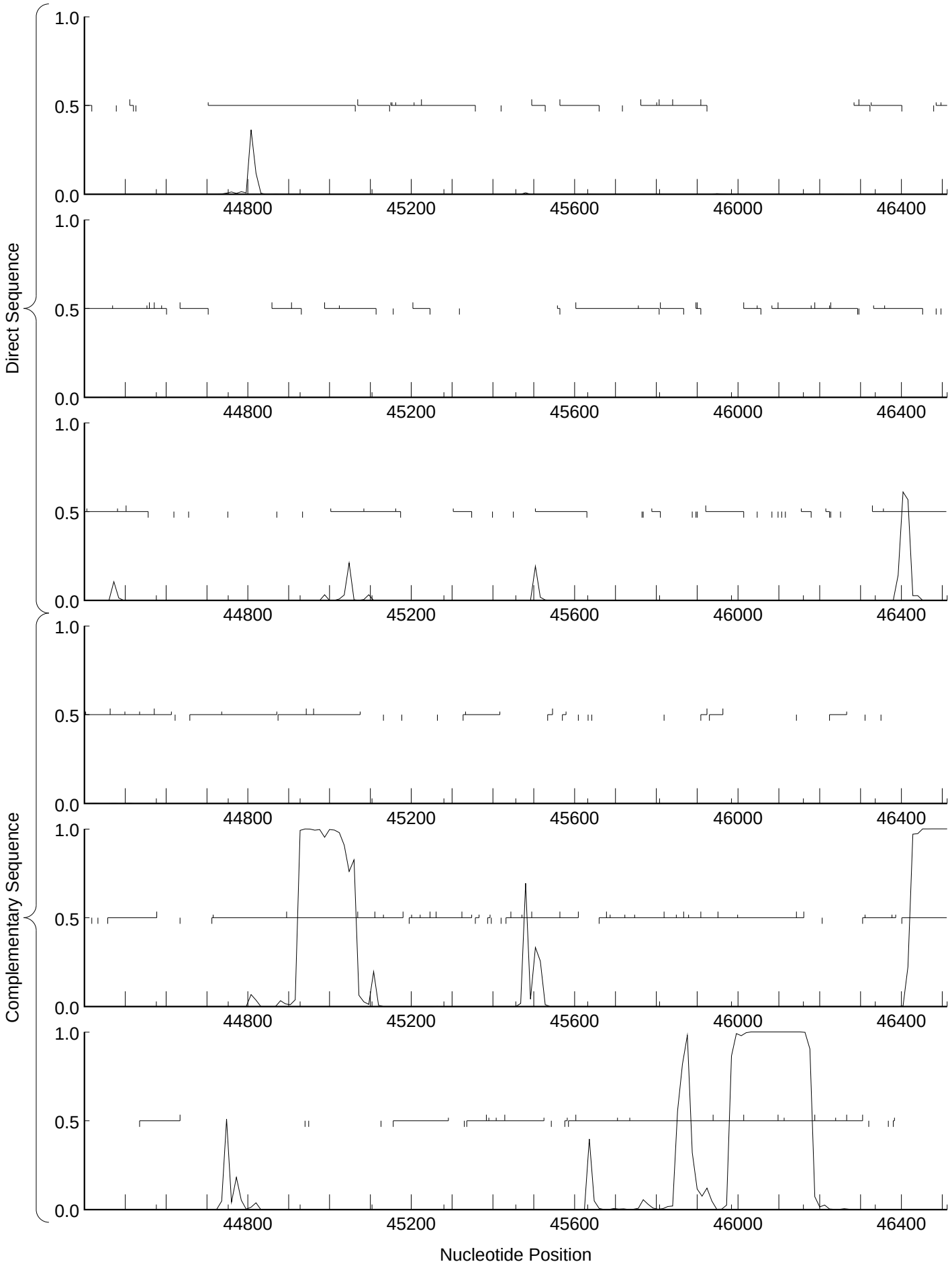

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

, phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 21/38

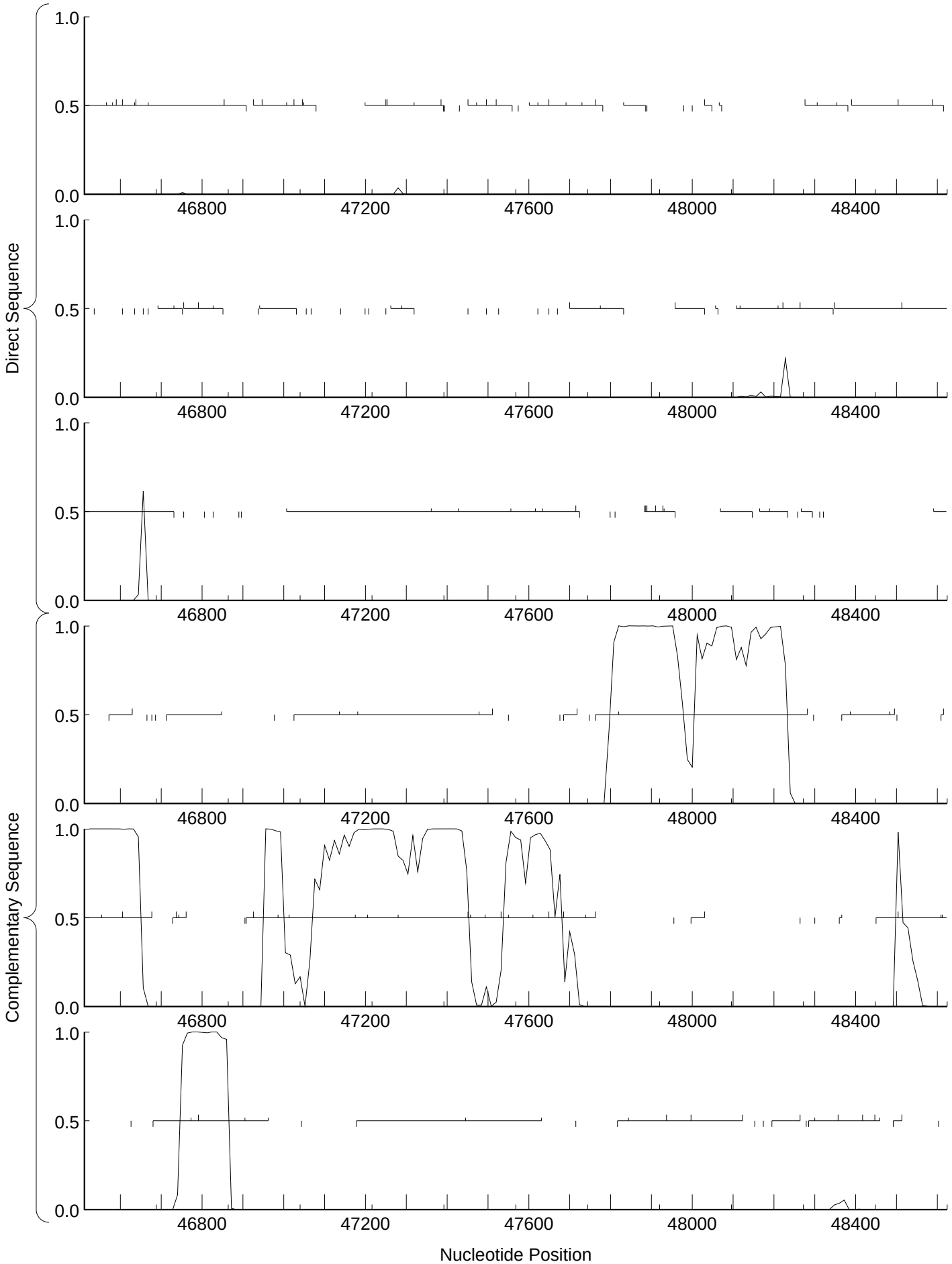


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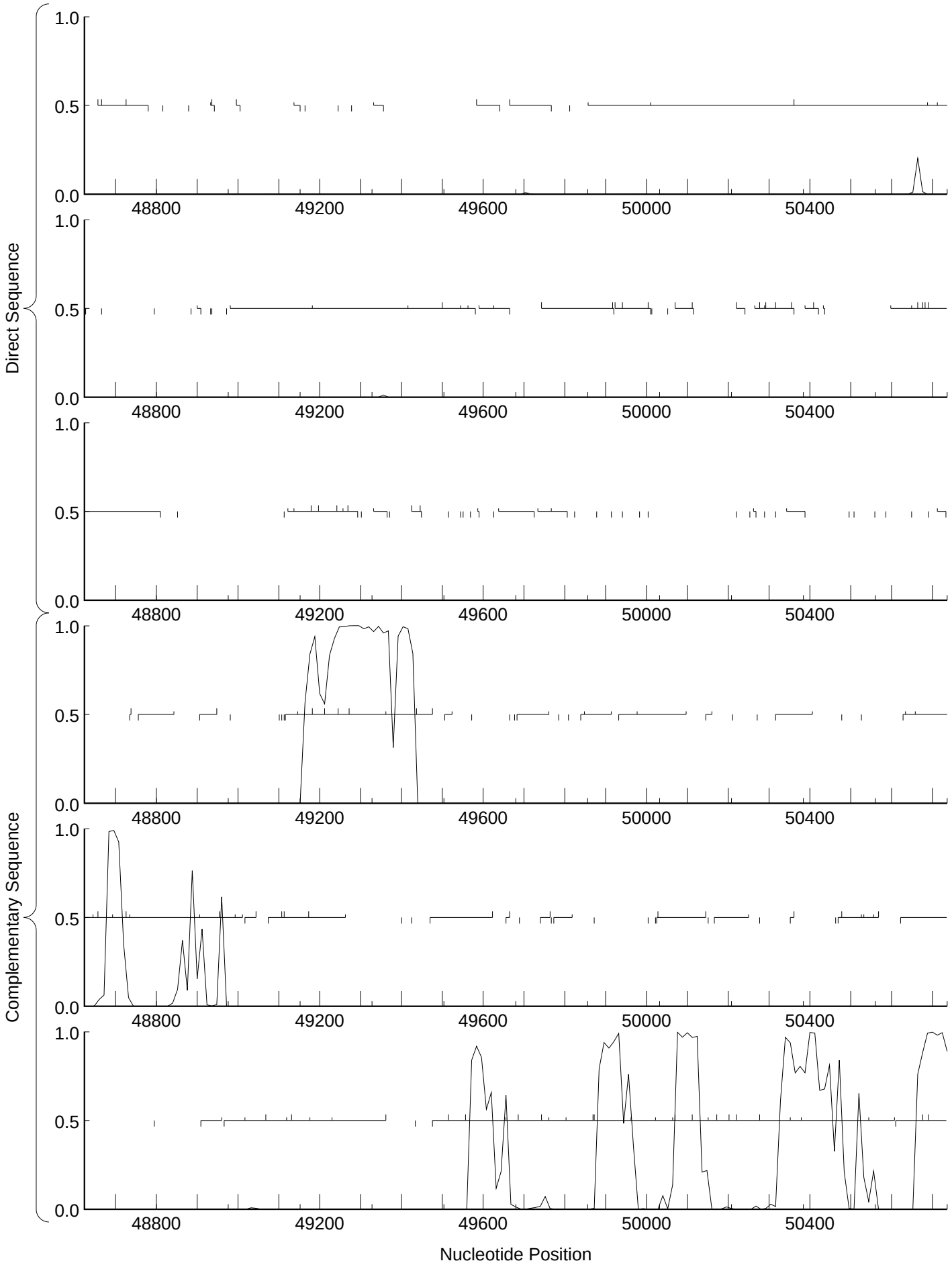



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
44800
45200
45600
46000
46400
Nucleotide Position

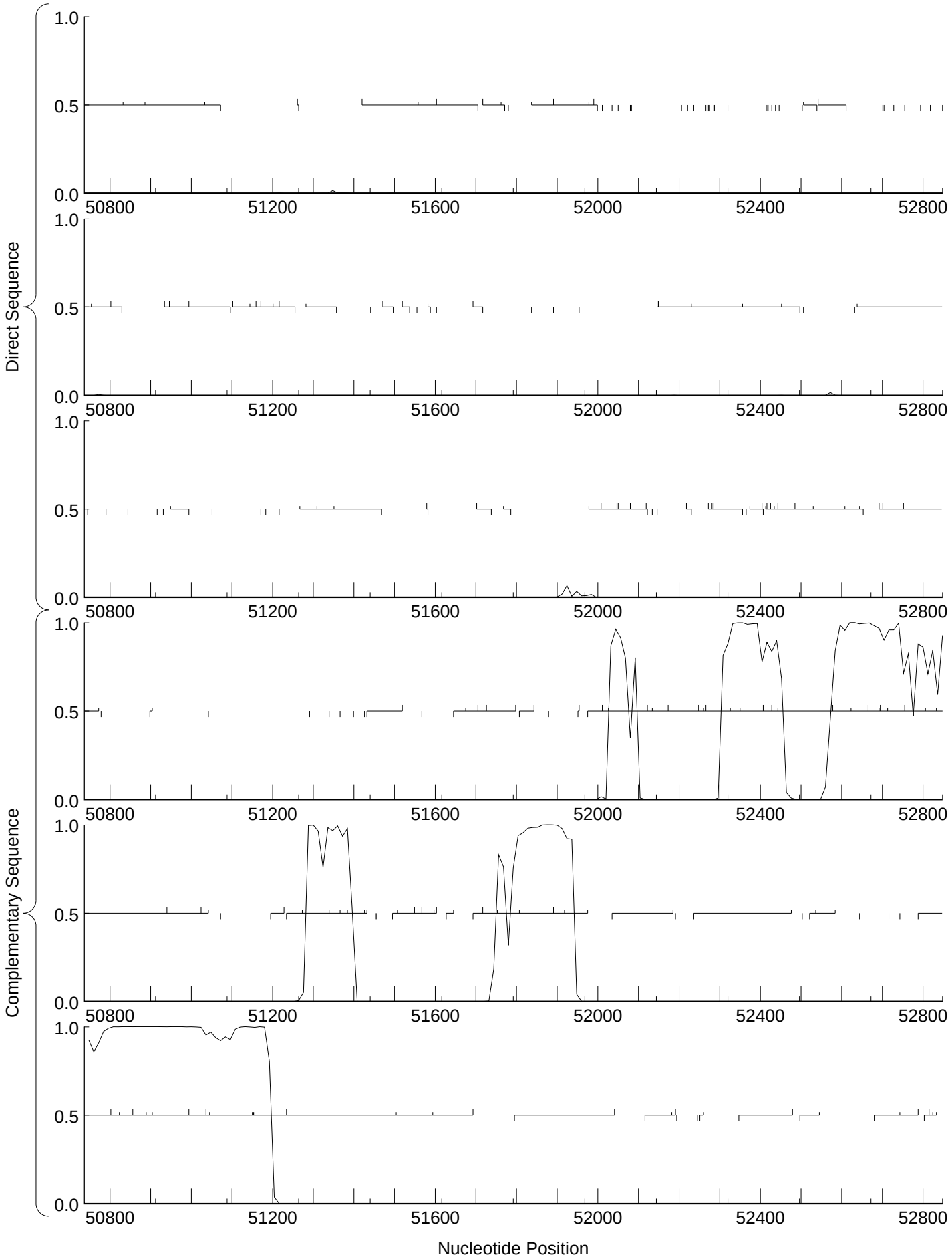
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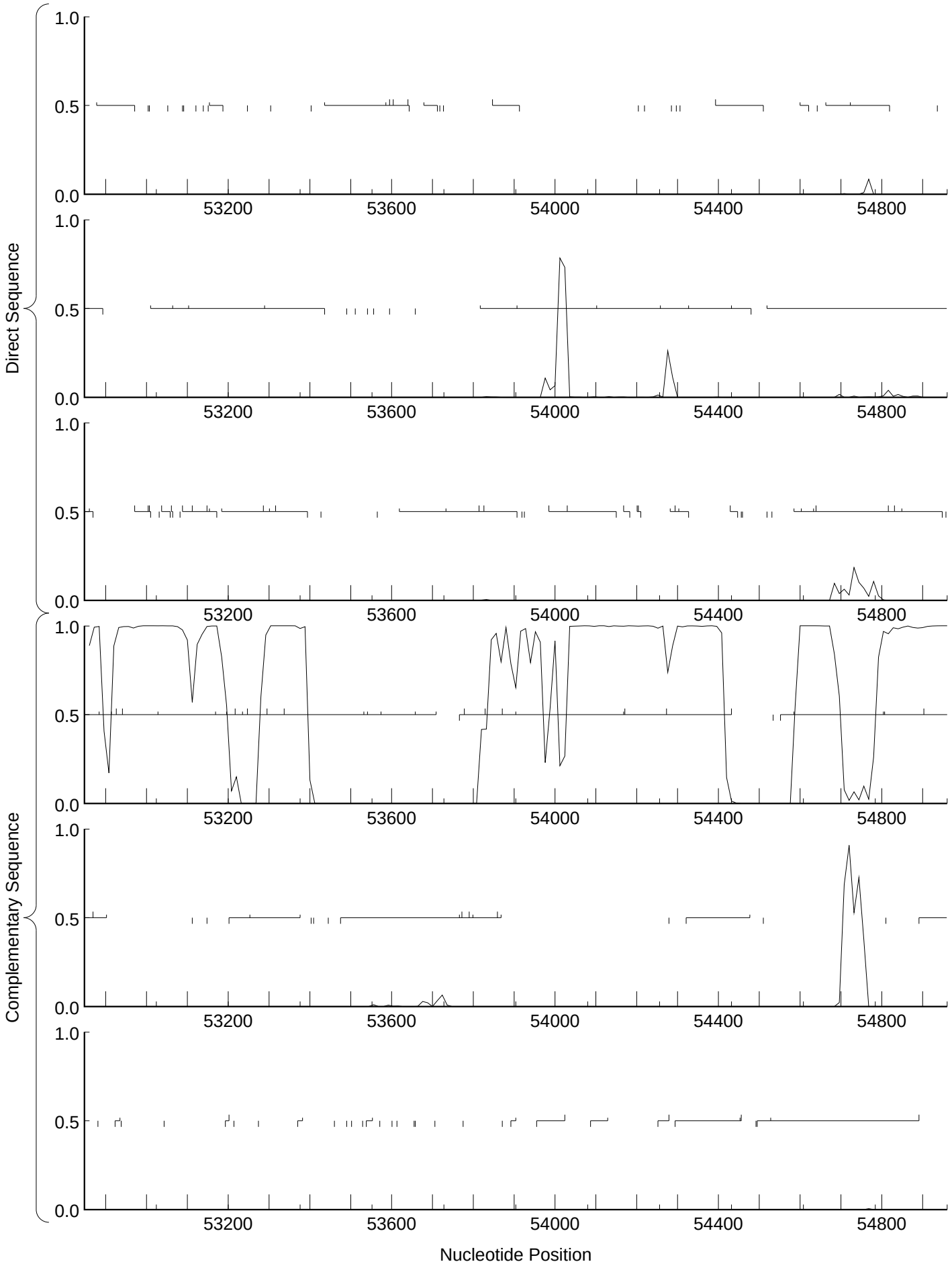


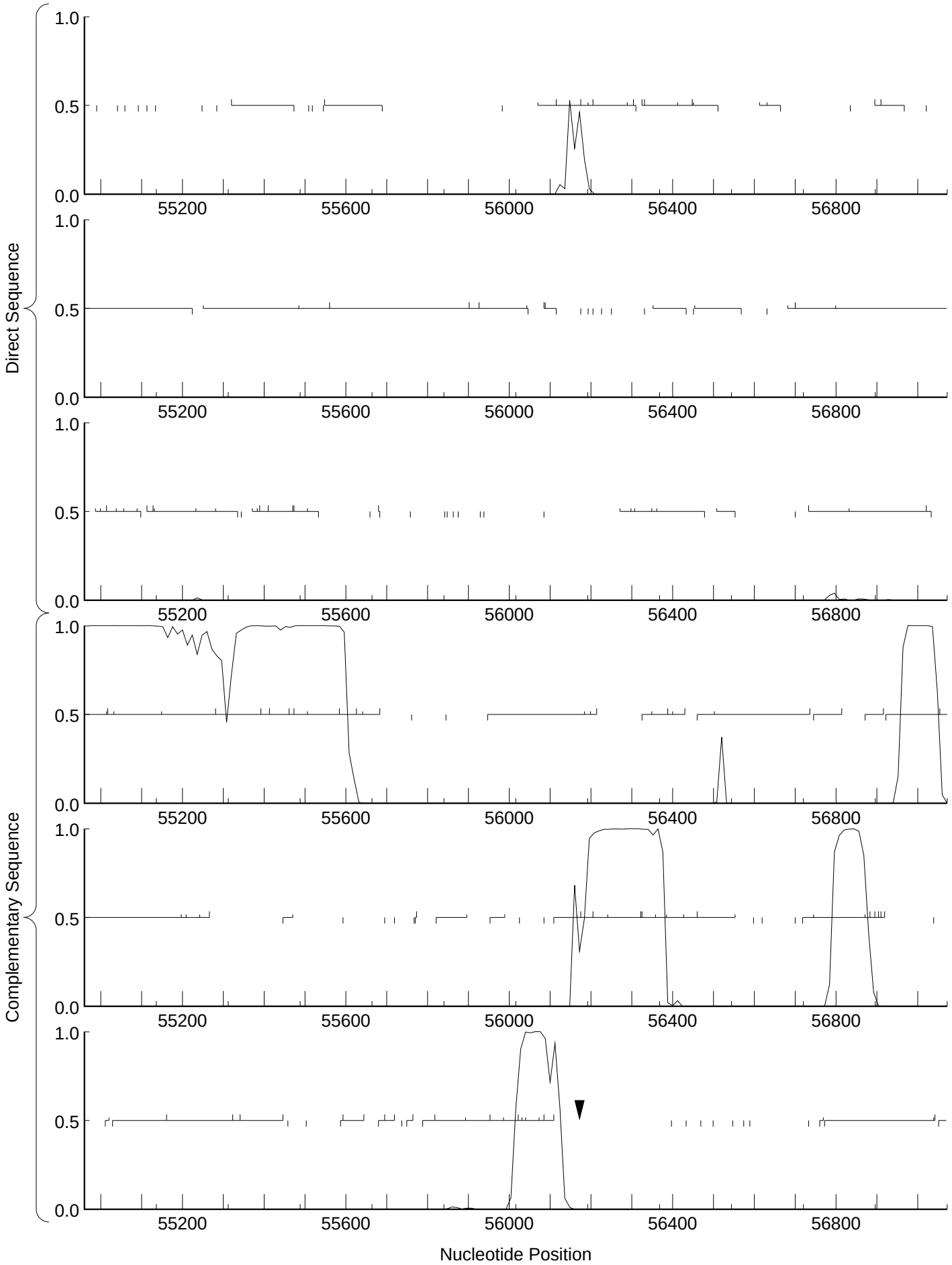
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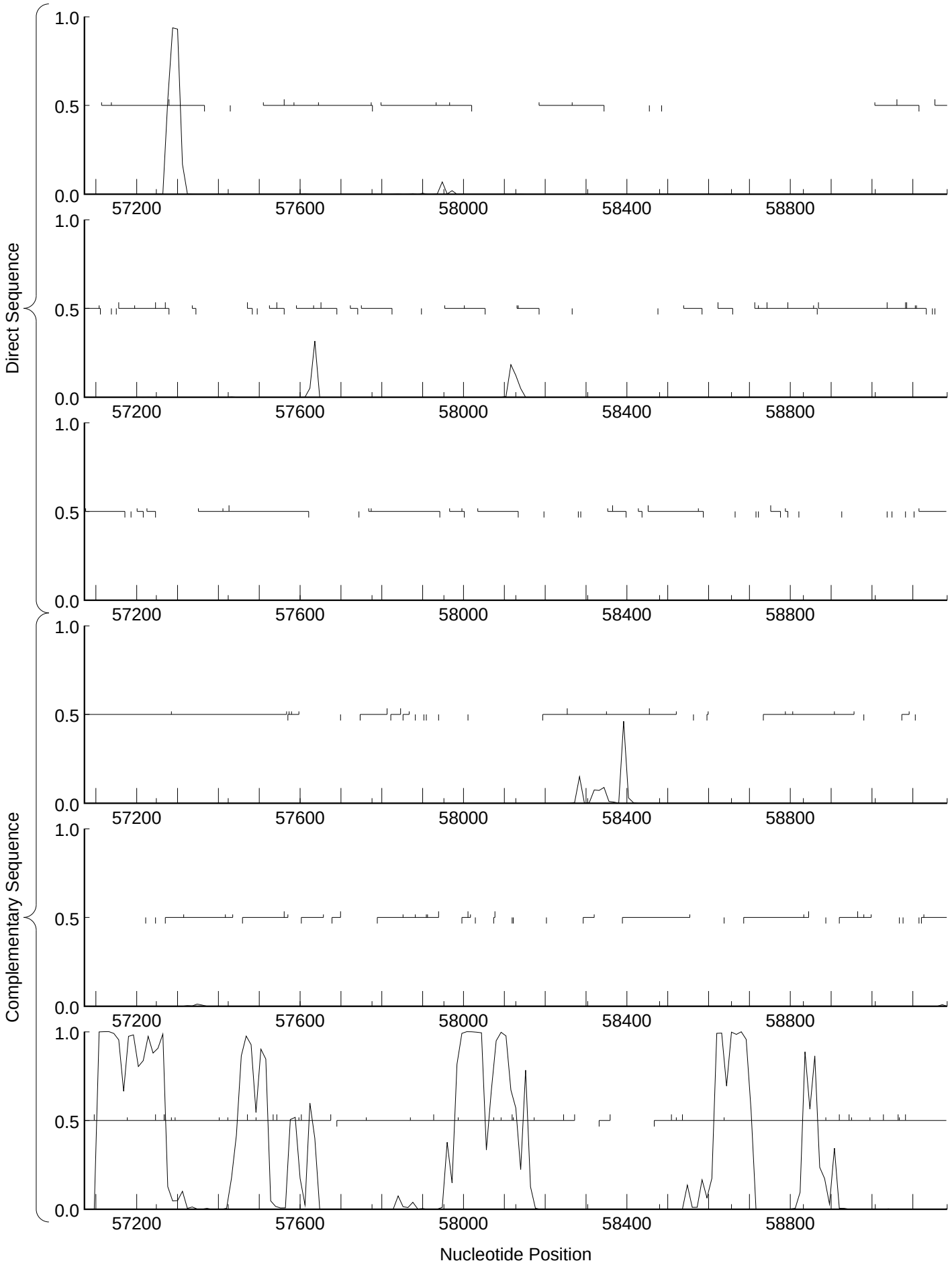
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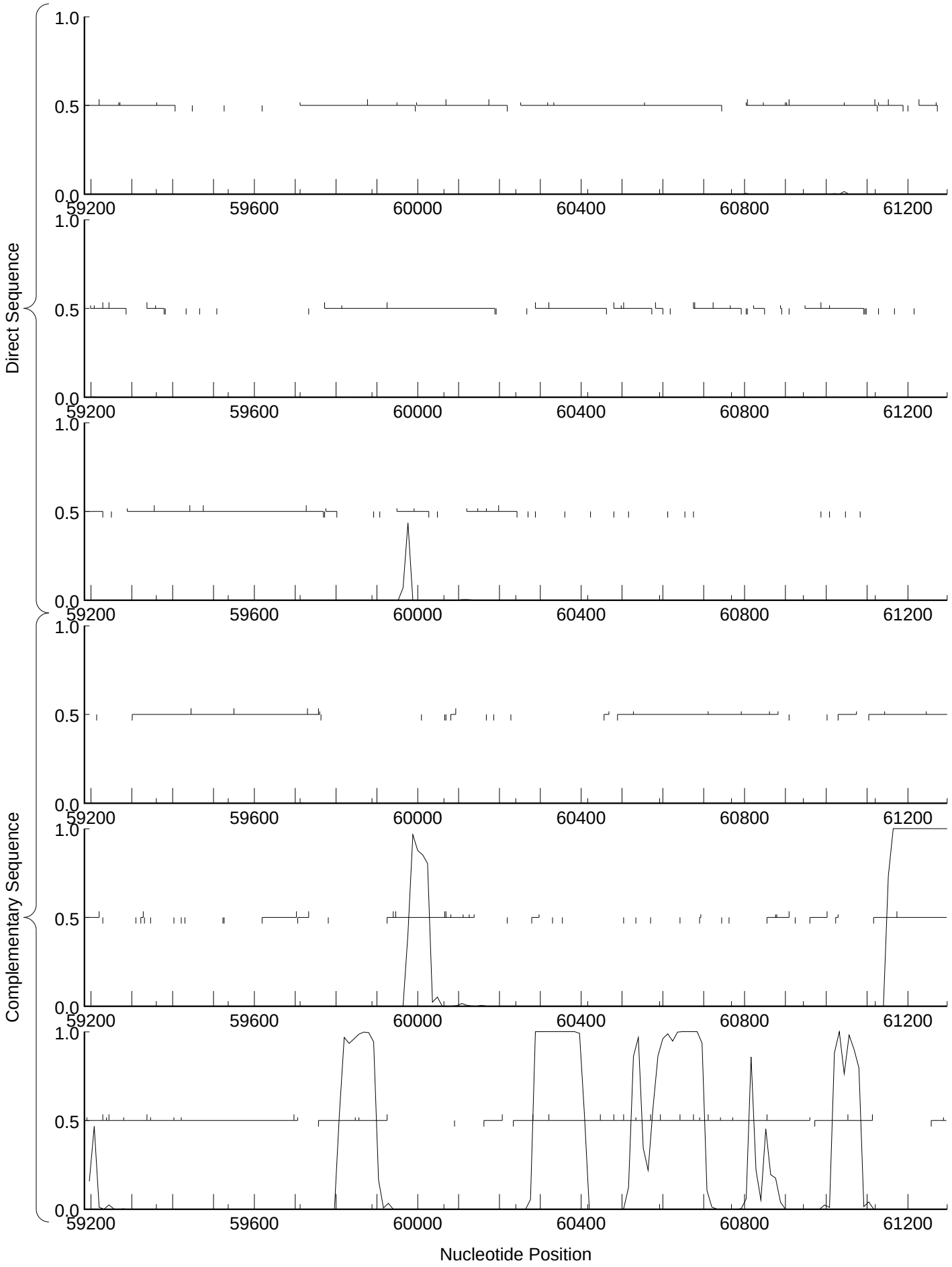

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
53200
53600
54000
54400
54800
Nucleotide Position


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

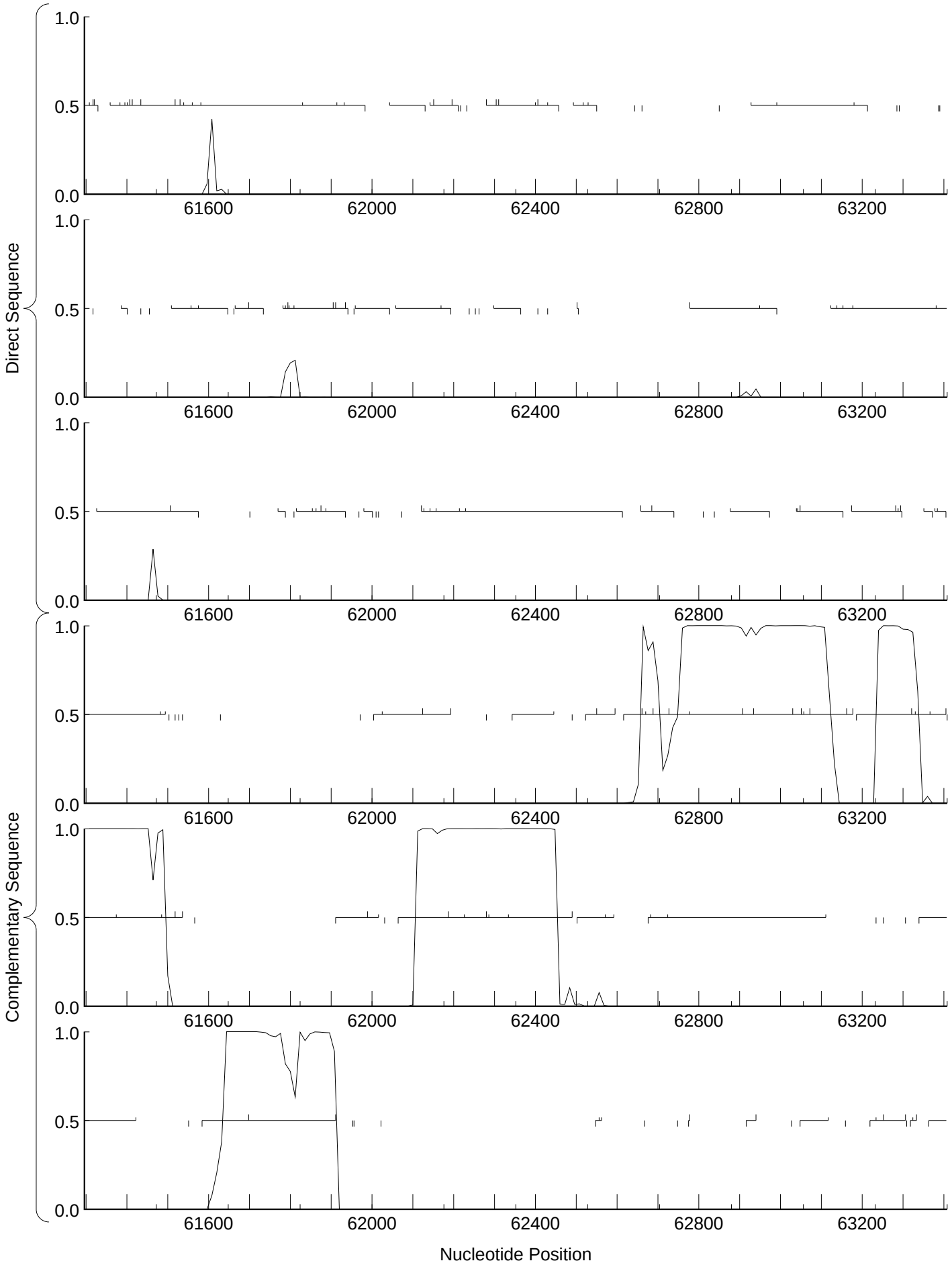
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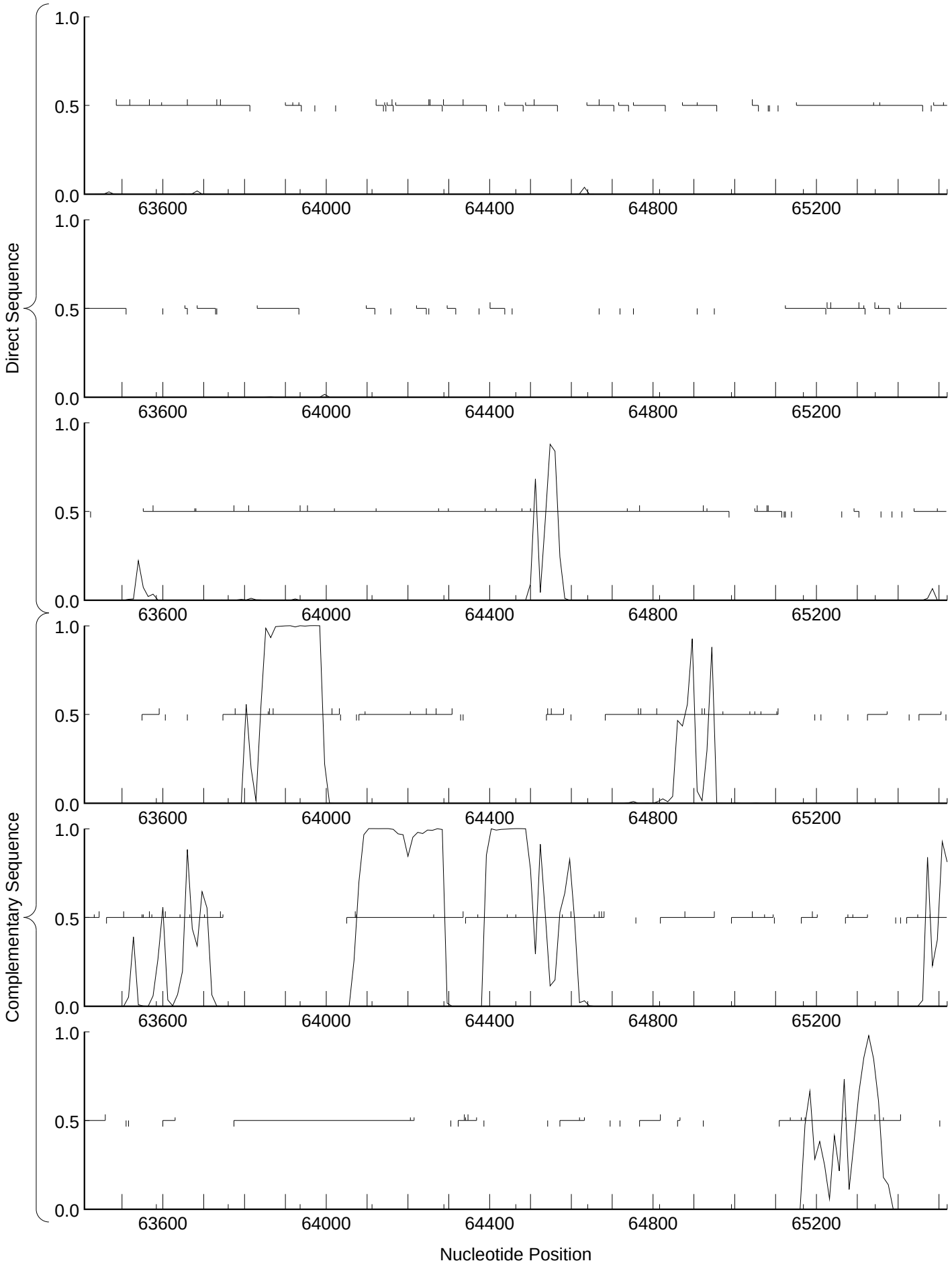
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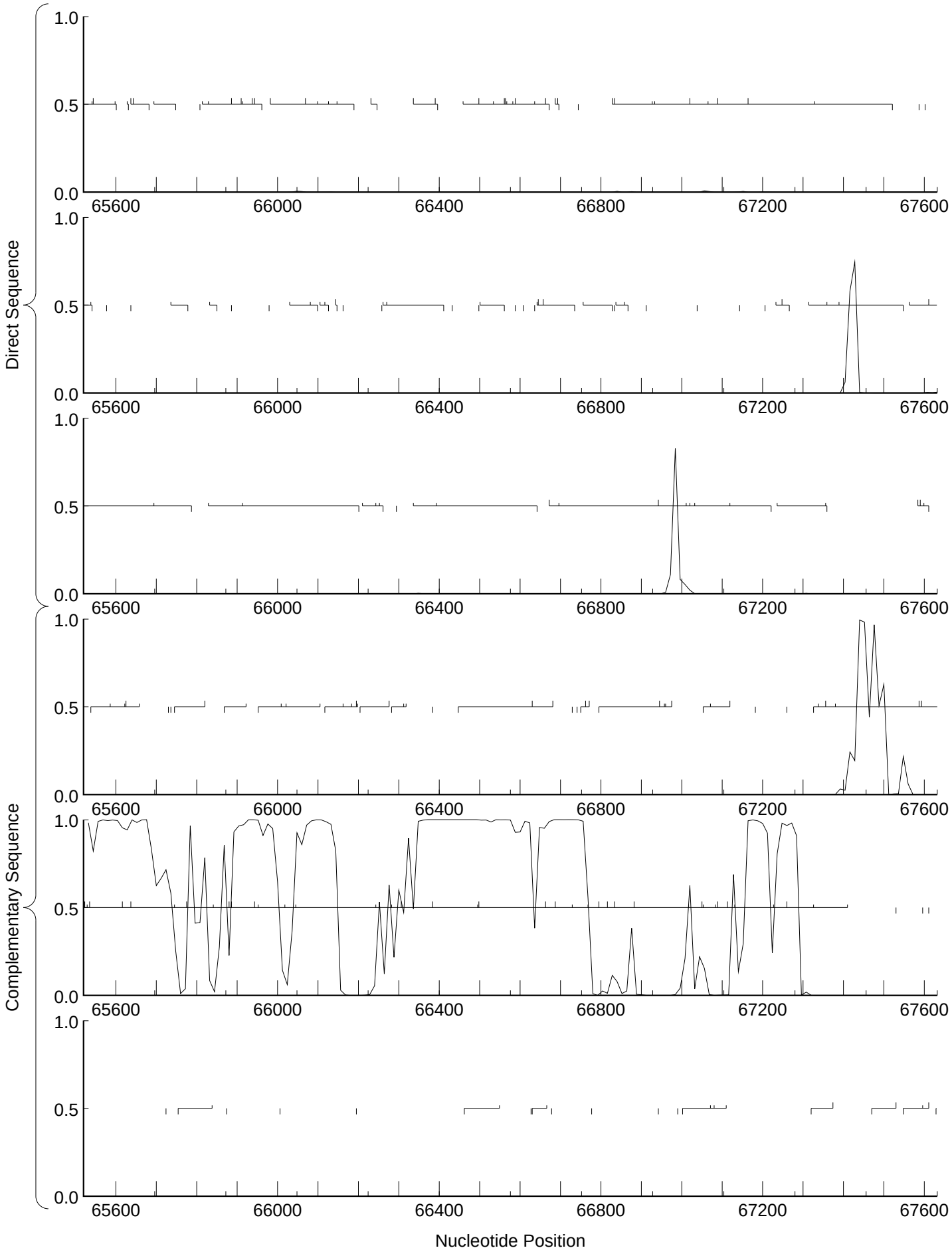


phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 31/38

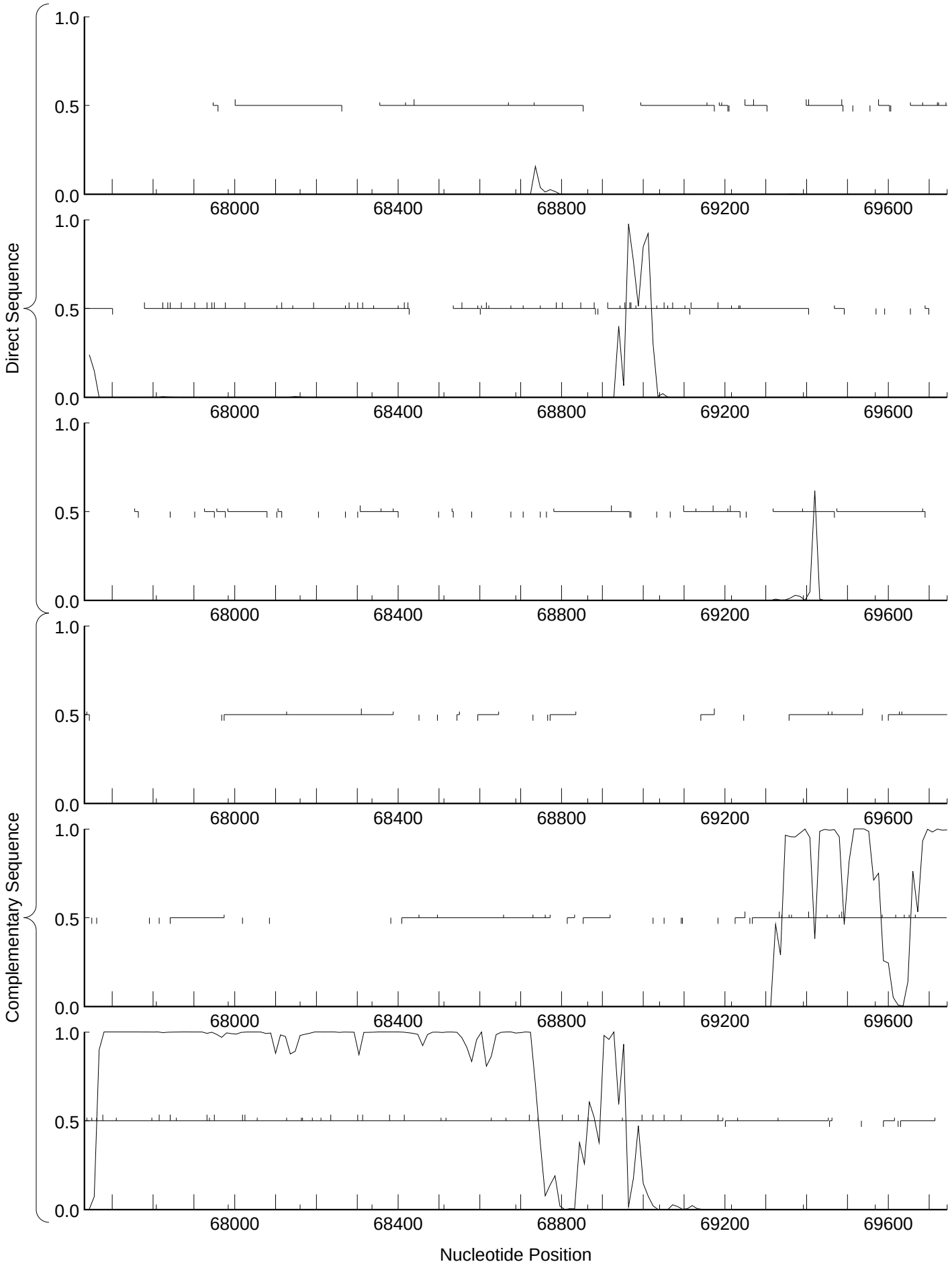


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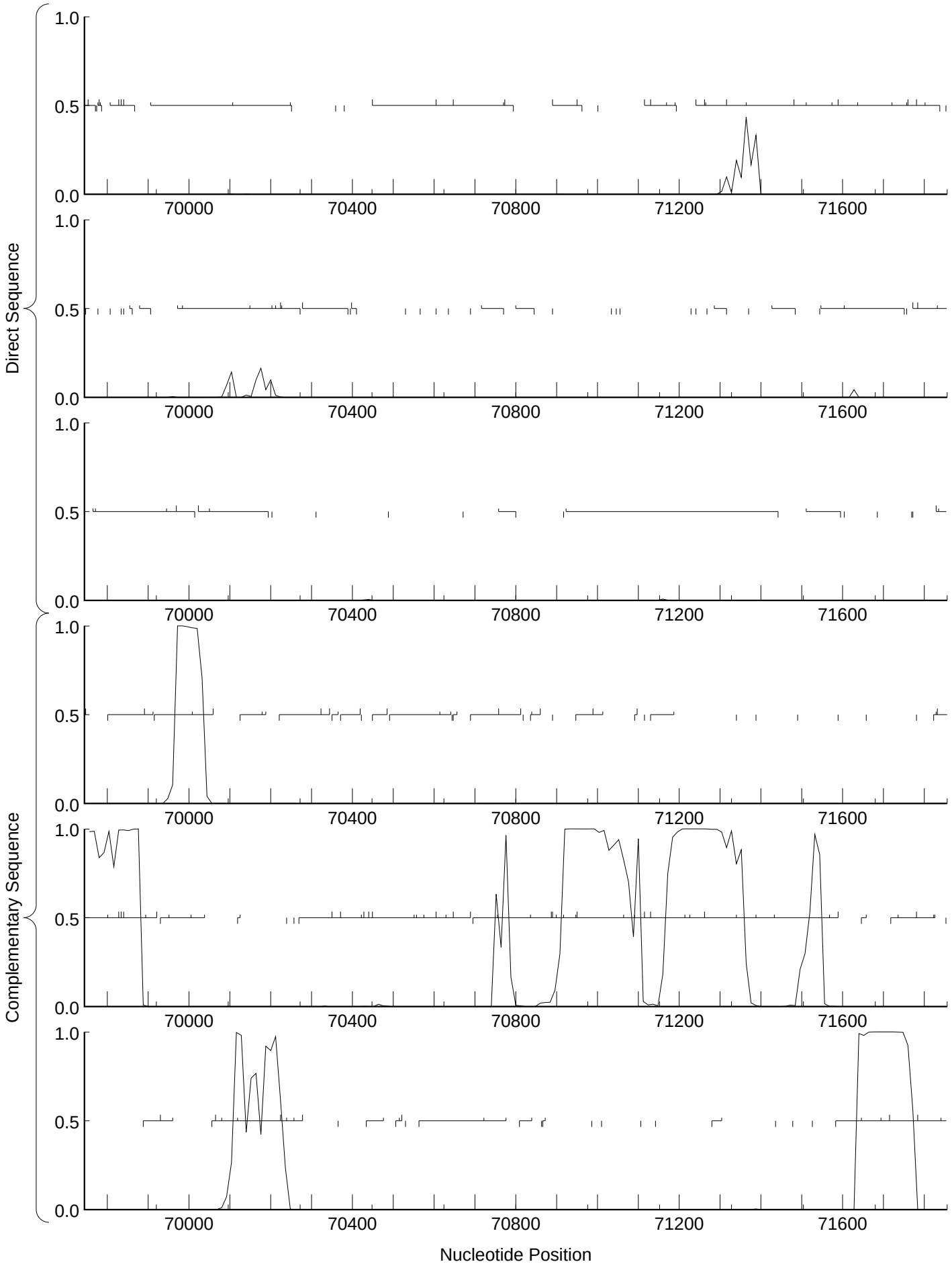


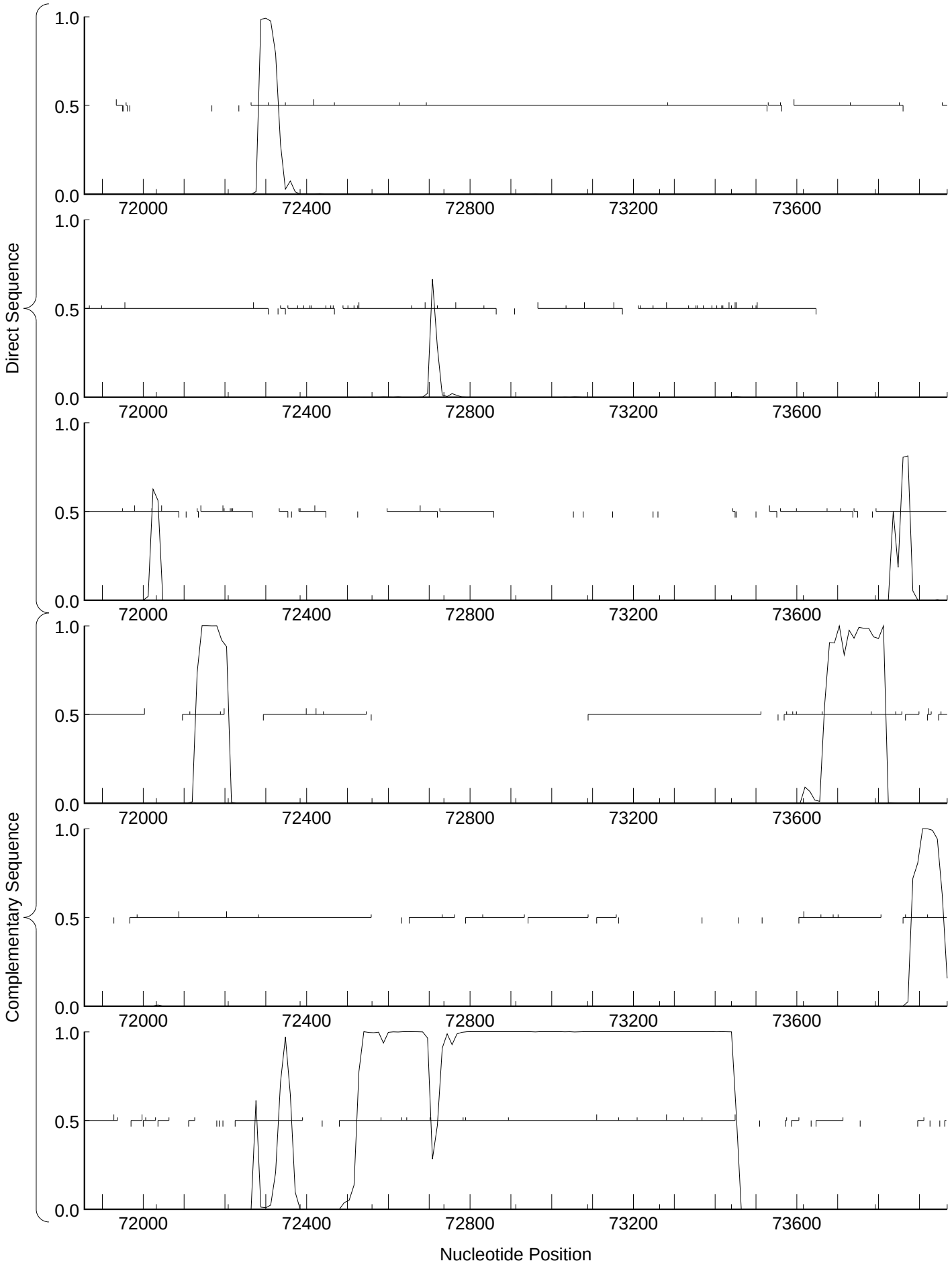

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
65600
66000
66400
66800
67200
67600
Nucleotide Position

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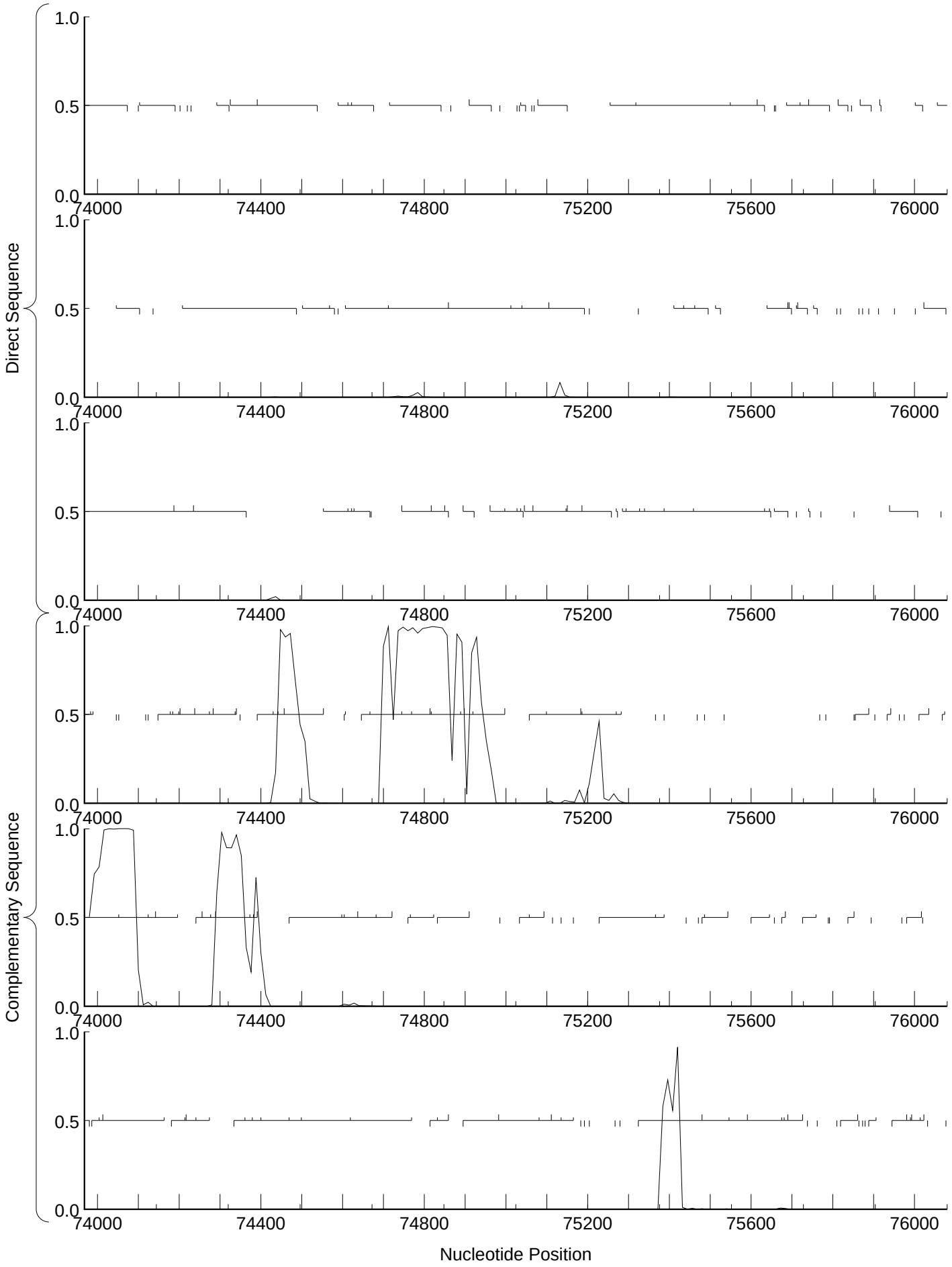


, phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 35/38




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
72000
72400
72800
73200
73600
Nucleotide Position

phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 37/38



, phage Beaver complete sequence, 77394 bp including 191 bp terminal repeat, Cluster CS2, Order 4, Window 96, Step 12, 38/38

