GeneMark

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

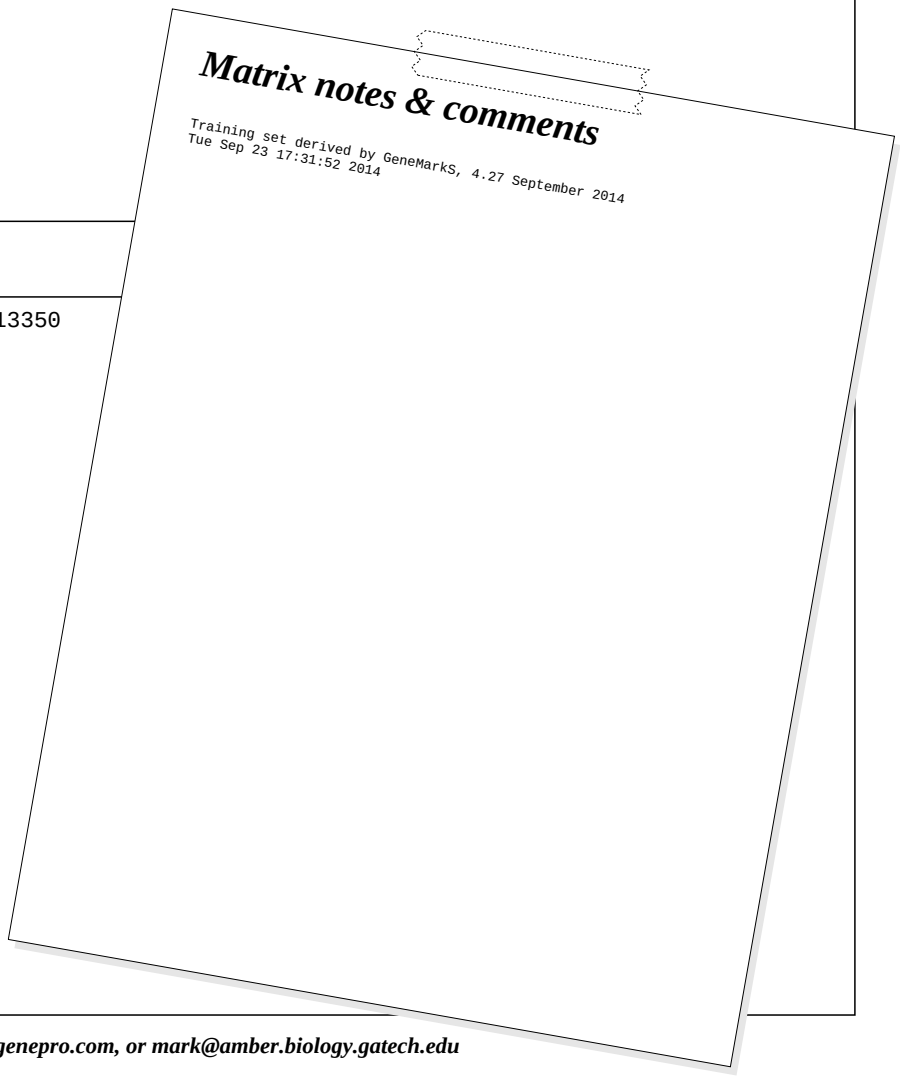
PROGRAM INFORMATION

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Pages            :  63
Sequence Length  :  130258 bp
GC Content       :   49.47%

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

PS-Version       :  1.2

GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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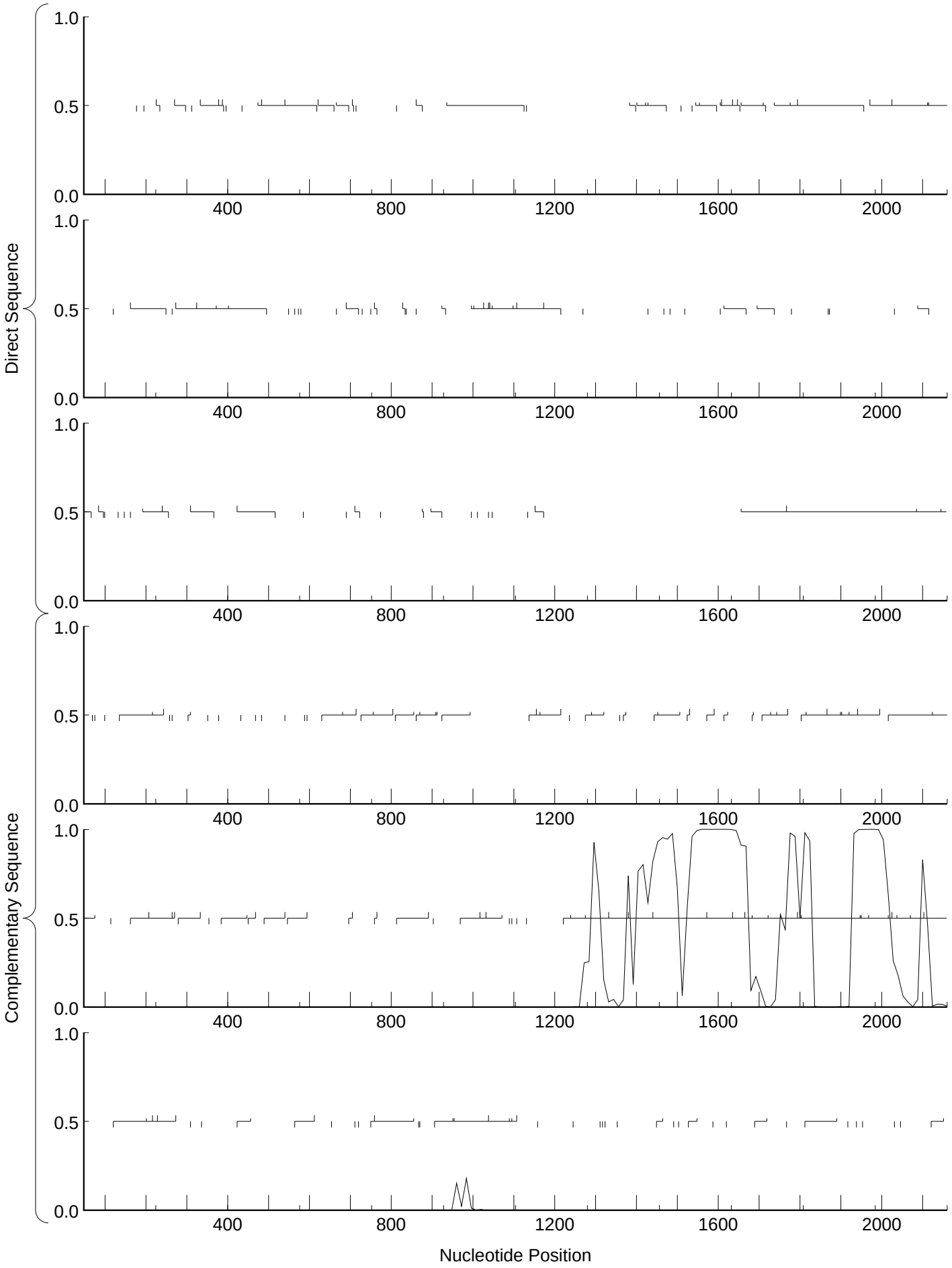
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Author     :  -

Order      :  4
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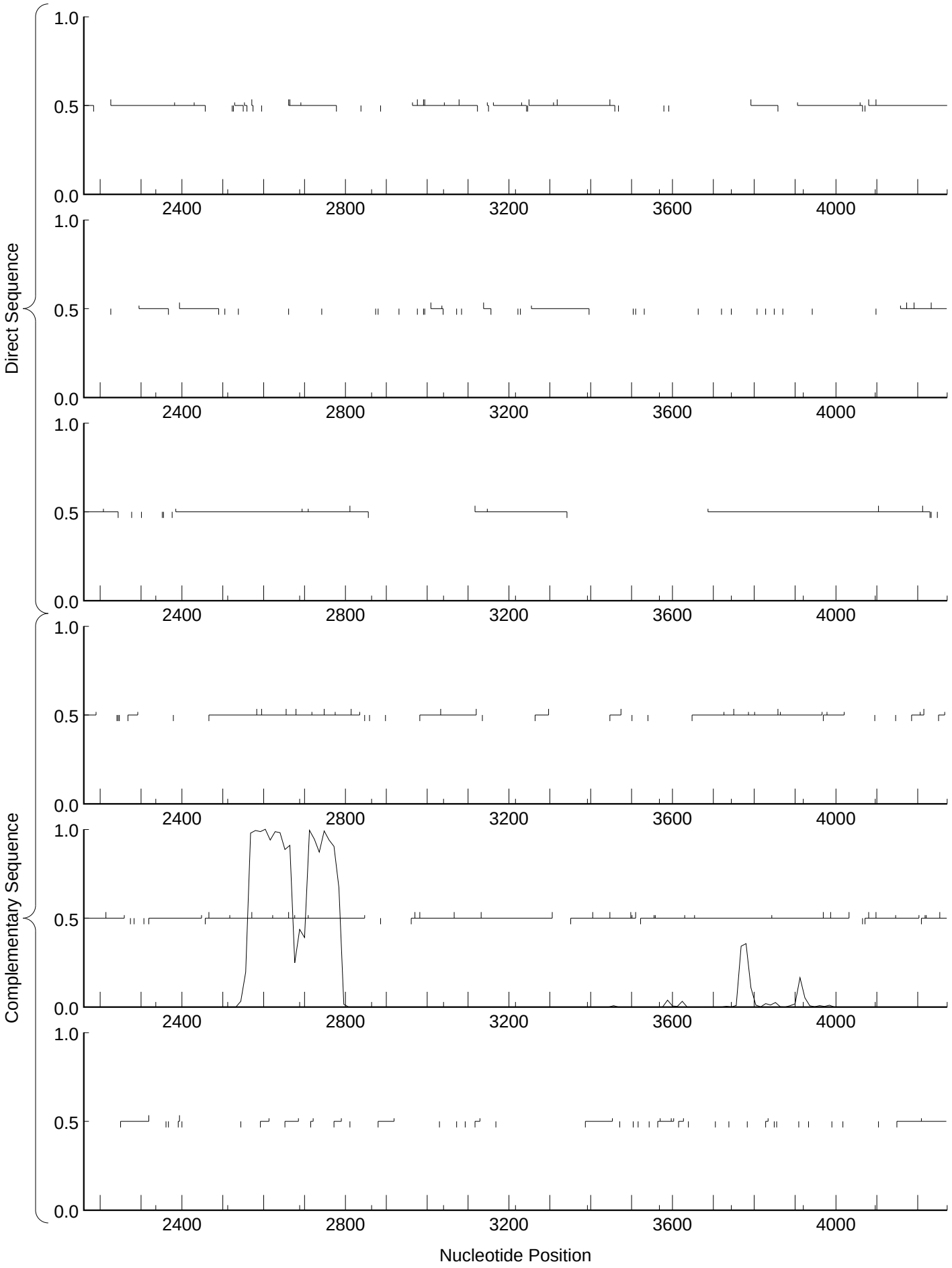

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

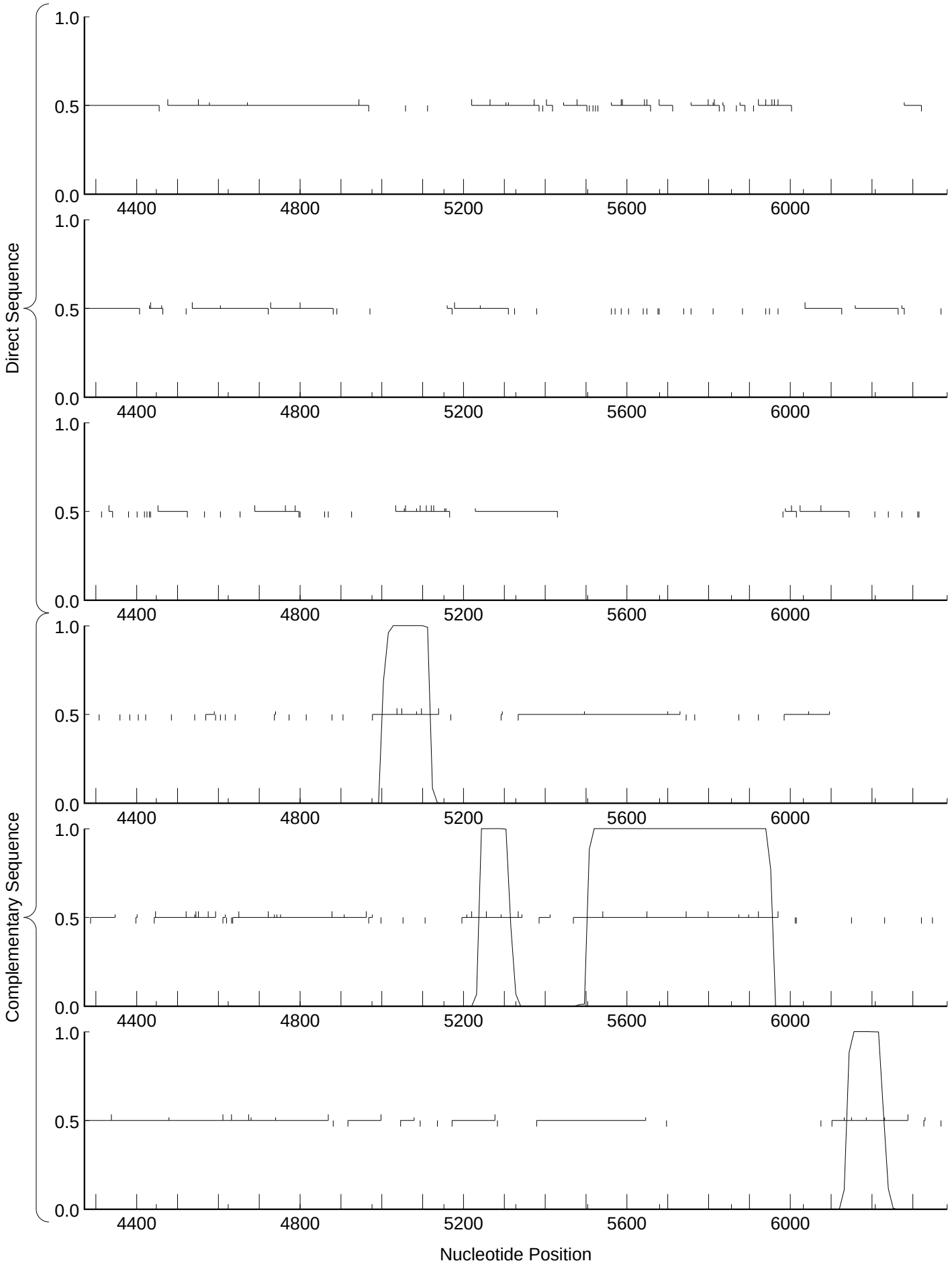
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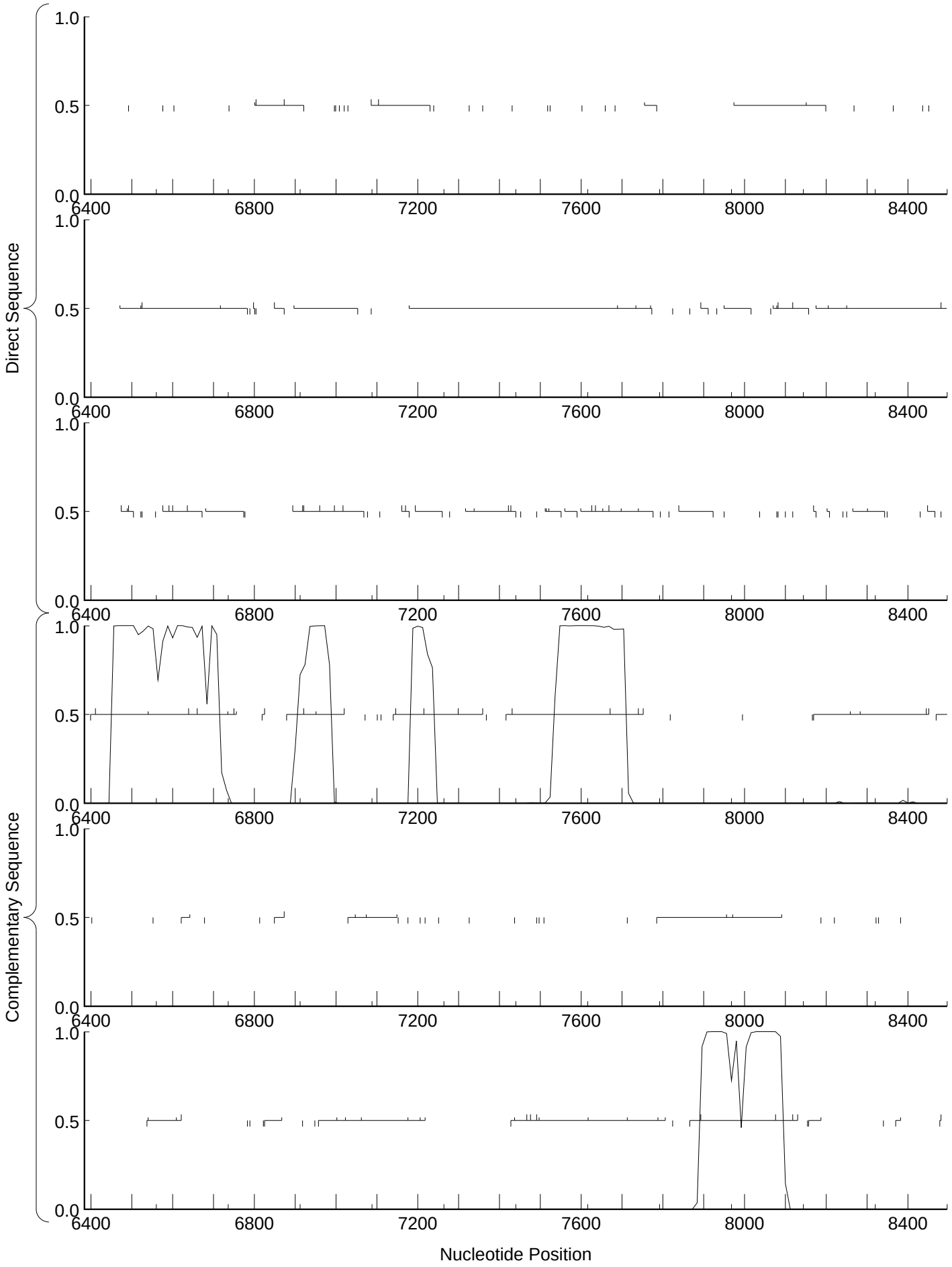
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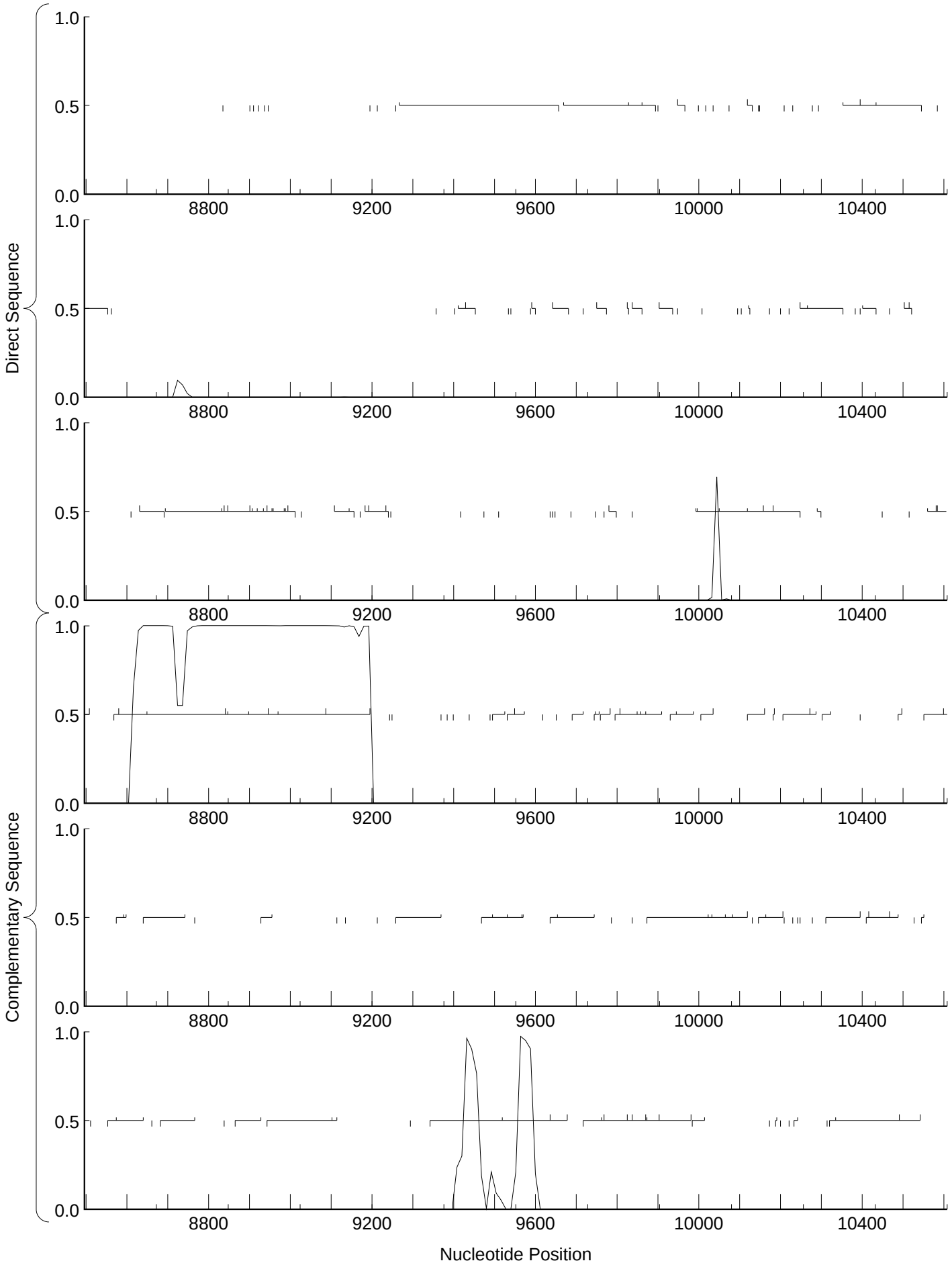
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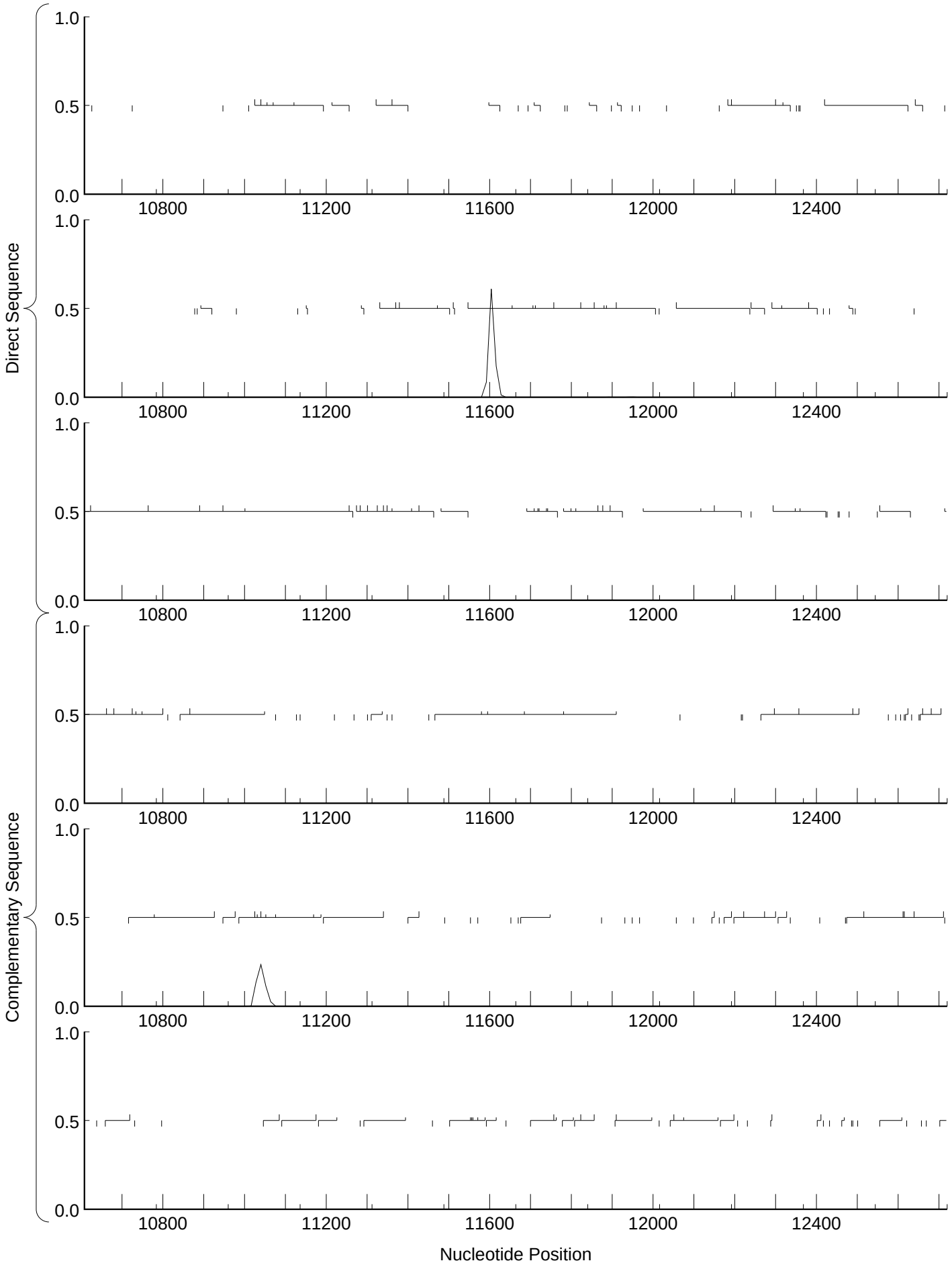
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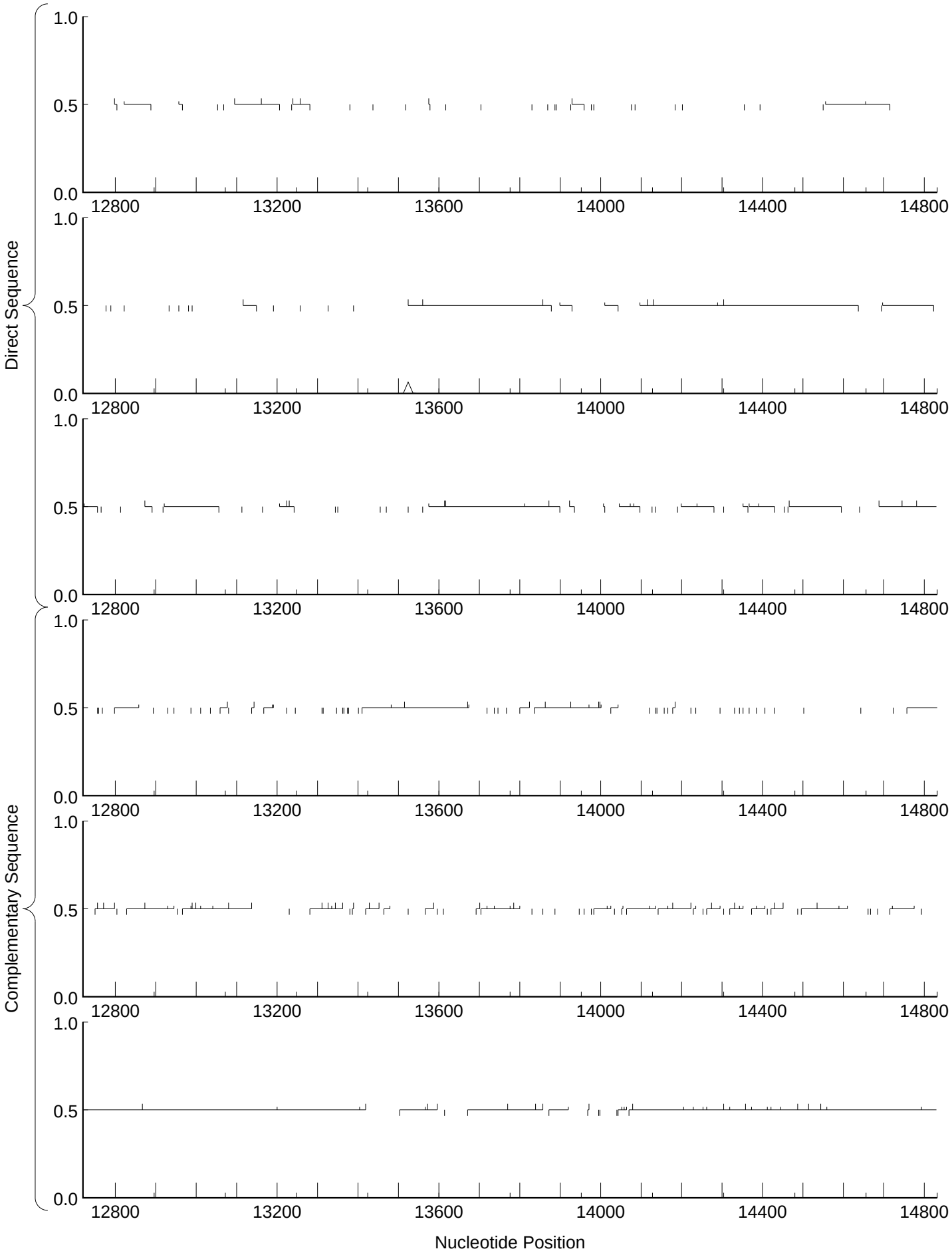
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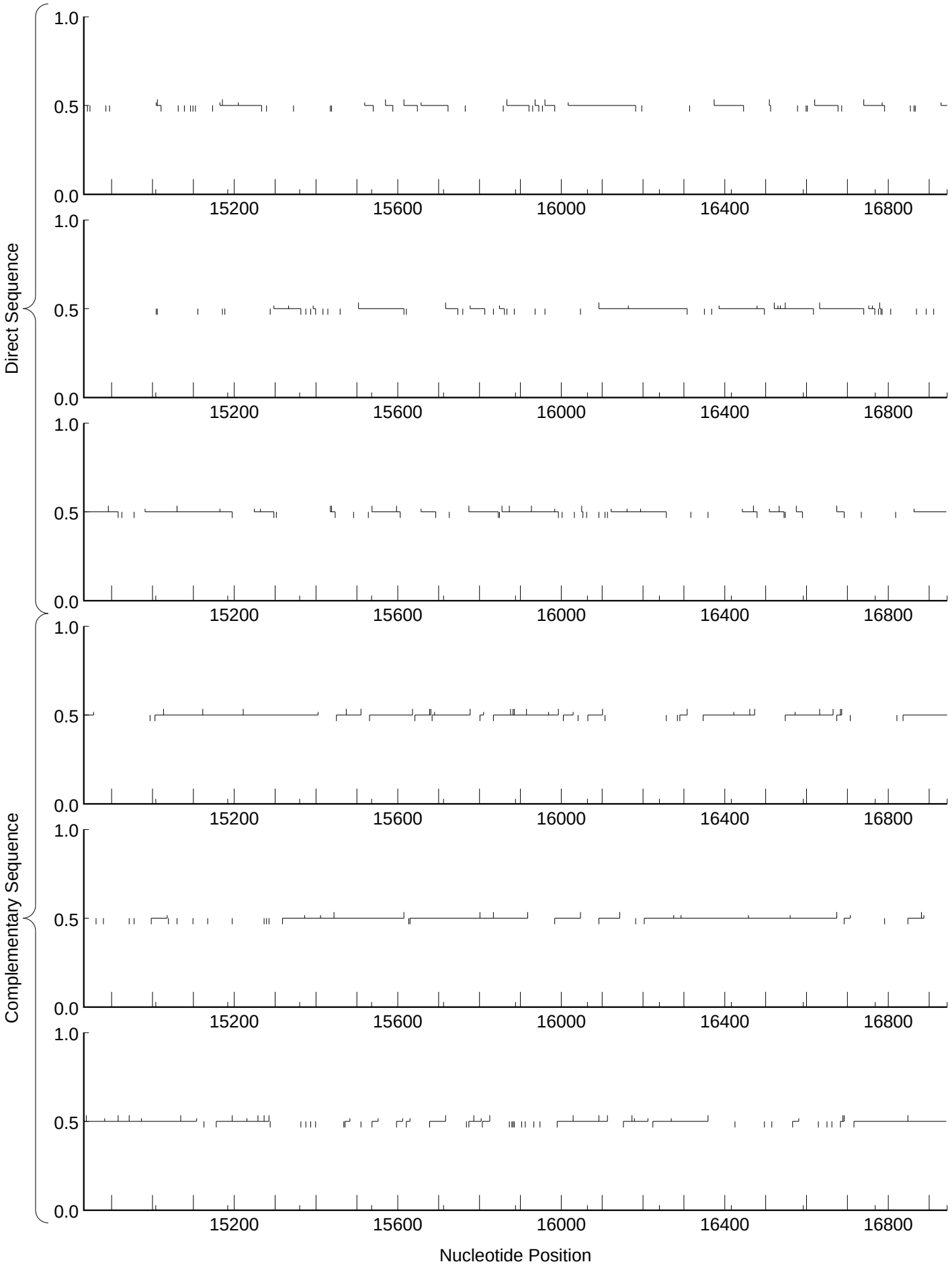
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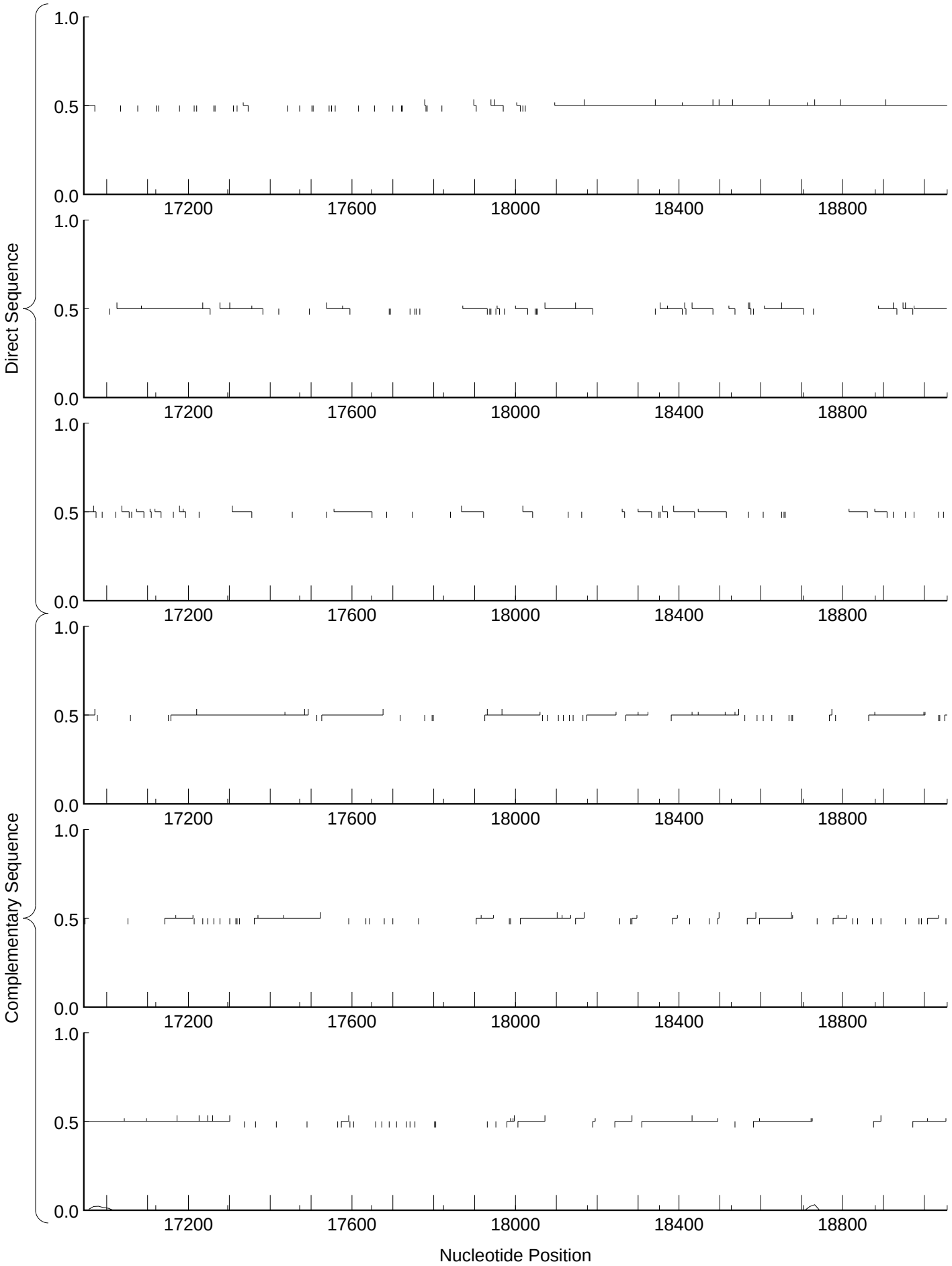
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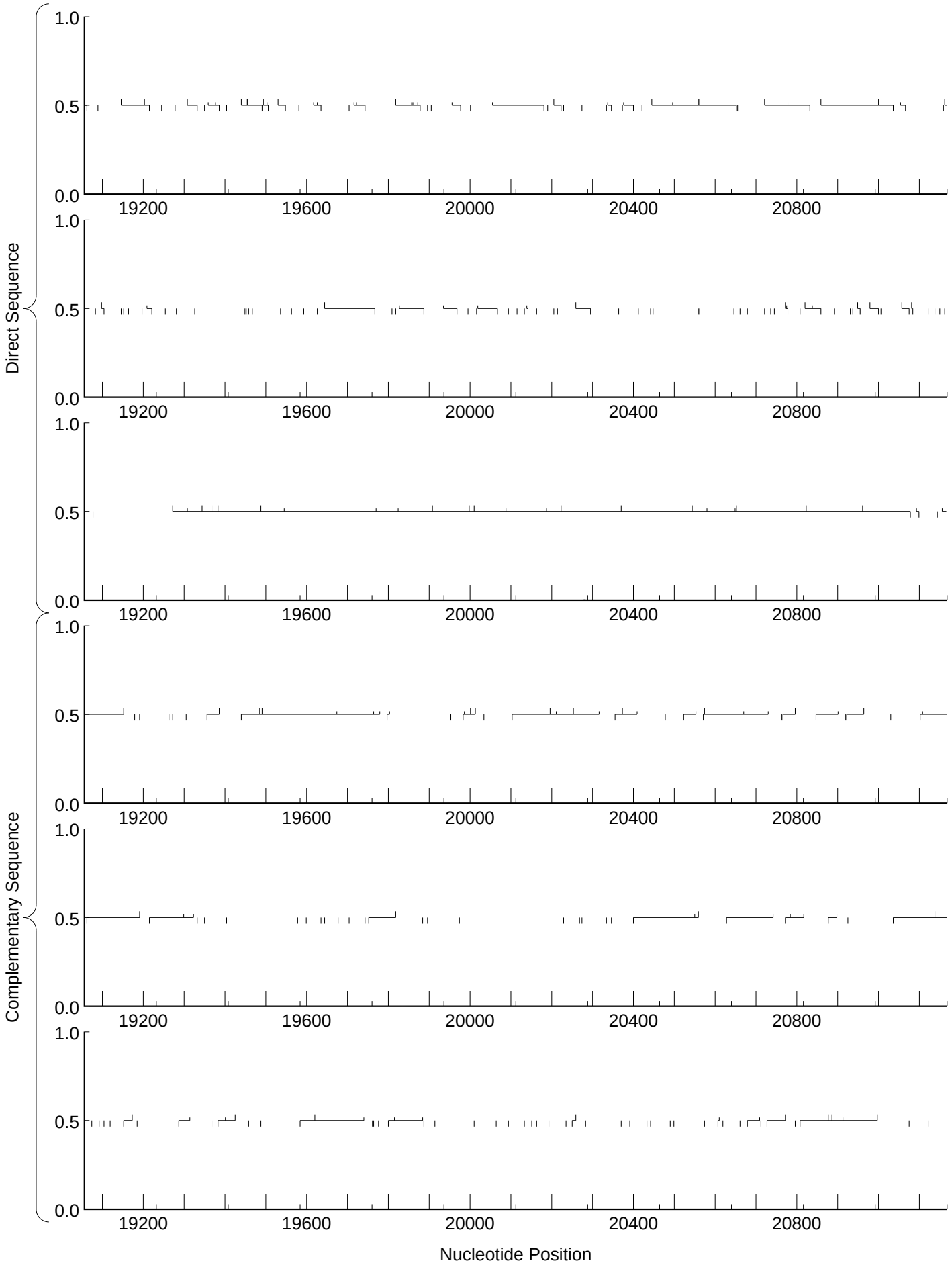
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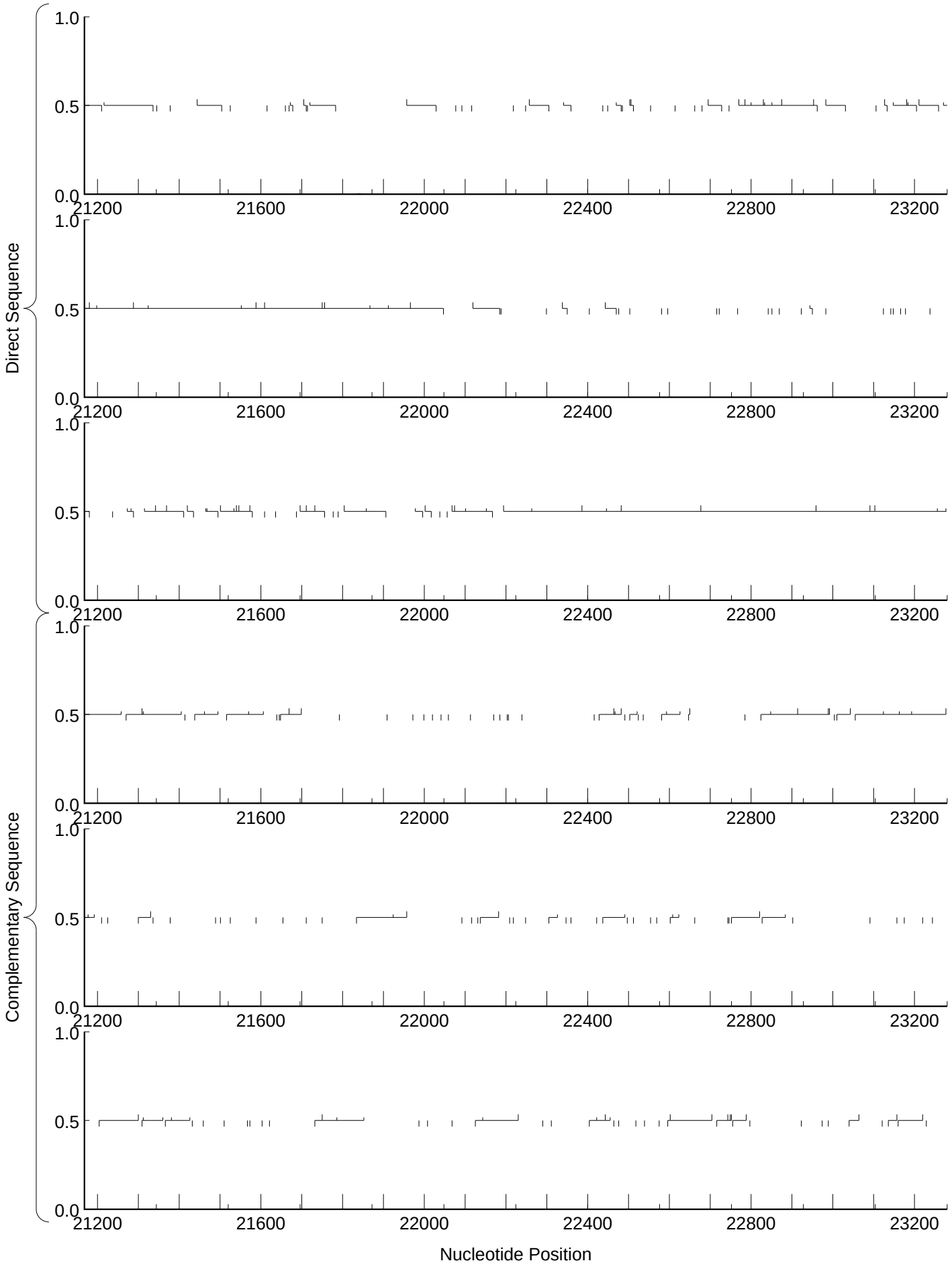
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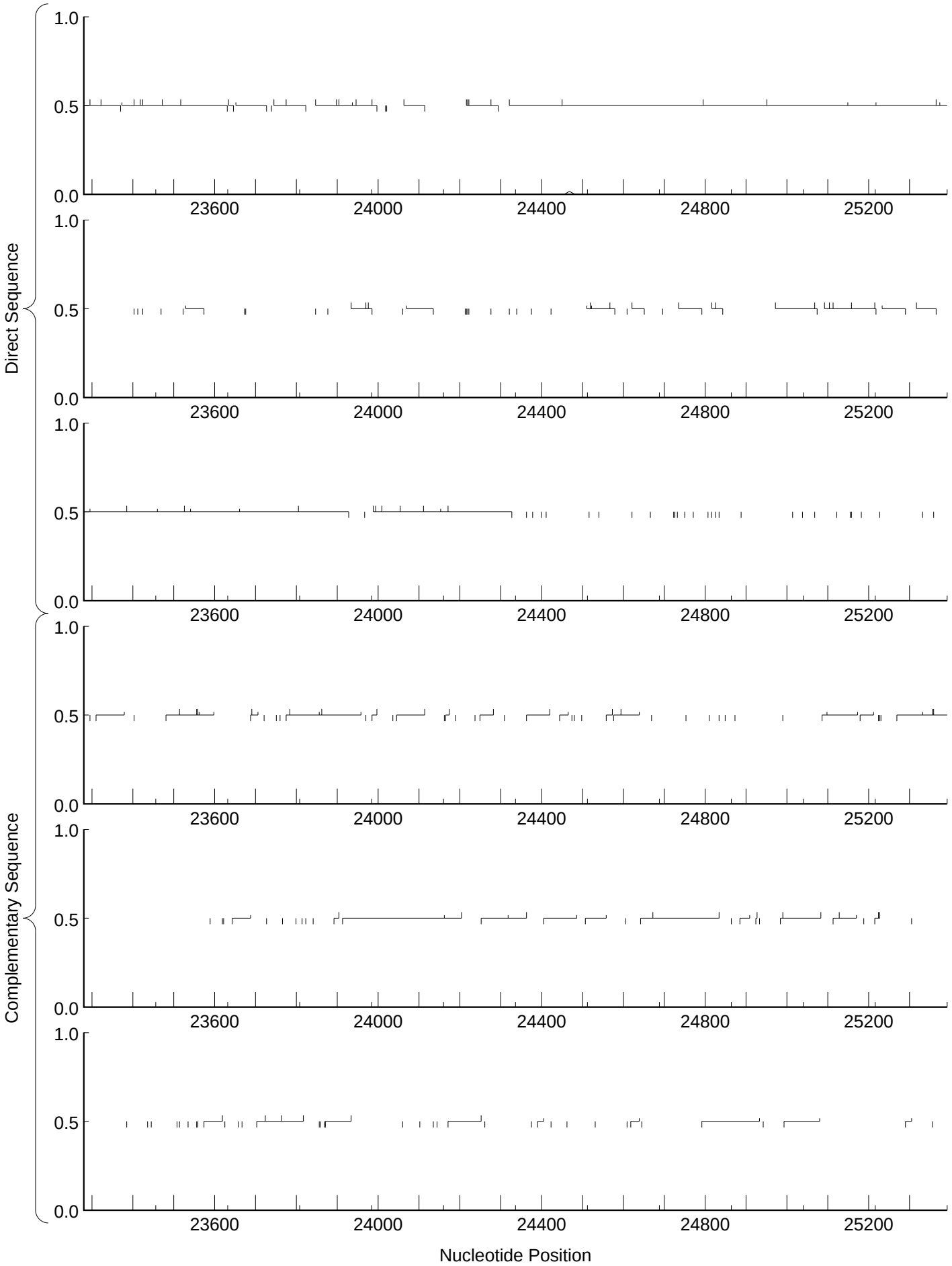
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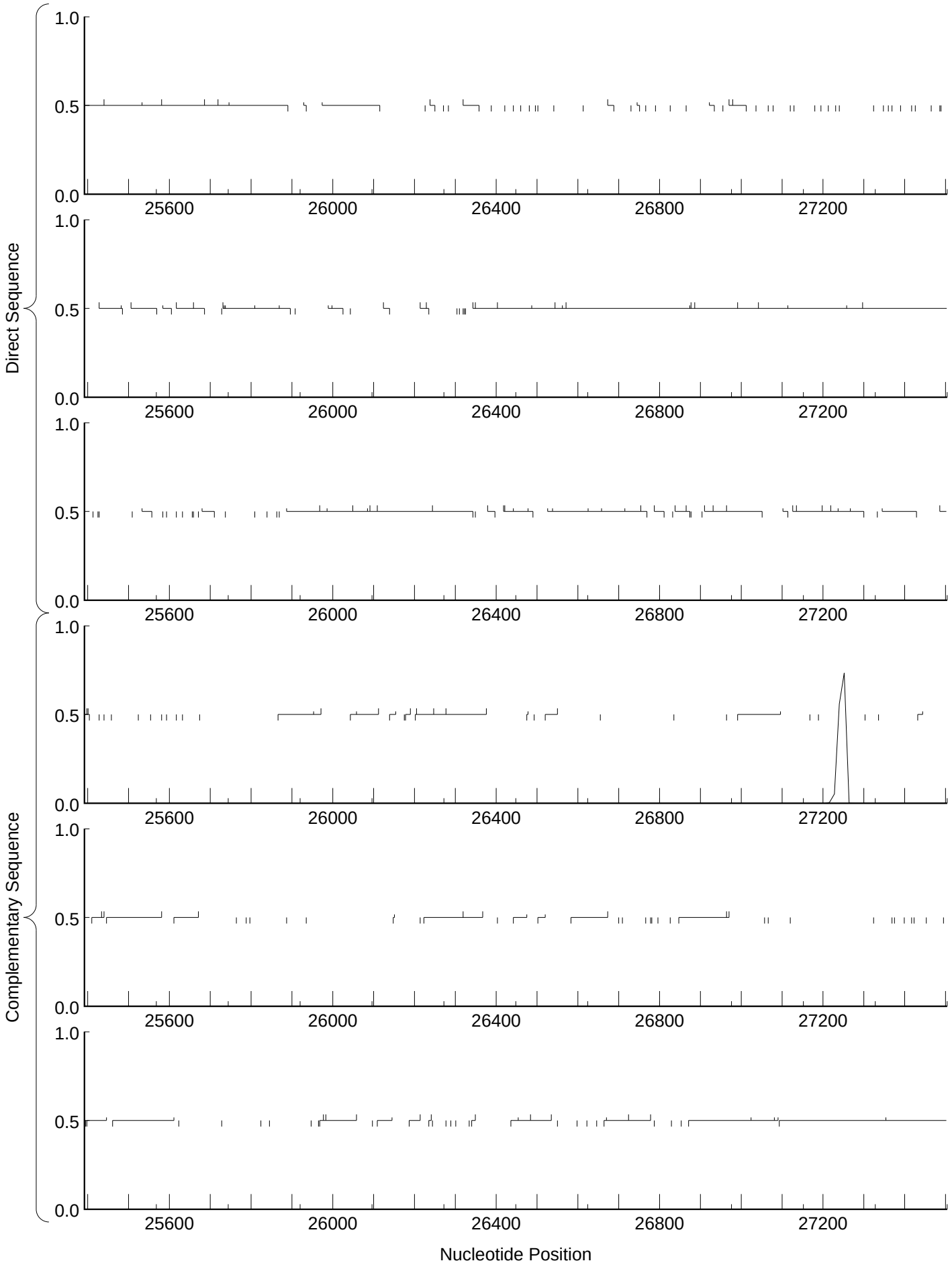
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 12/63



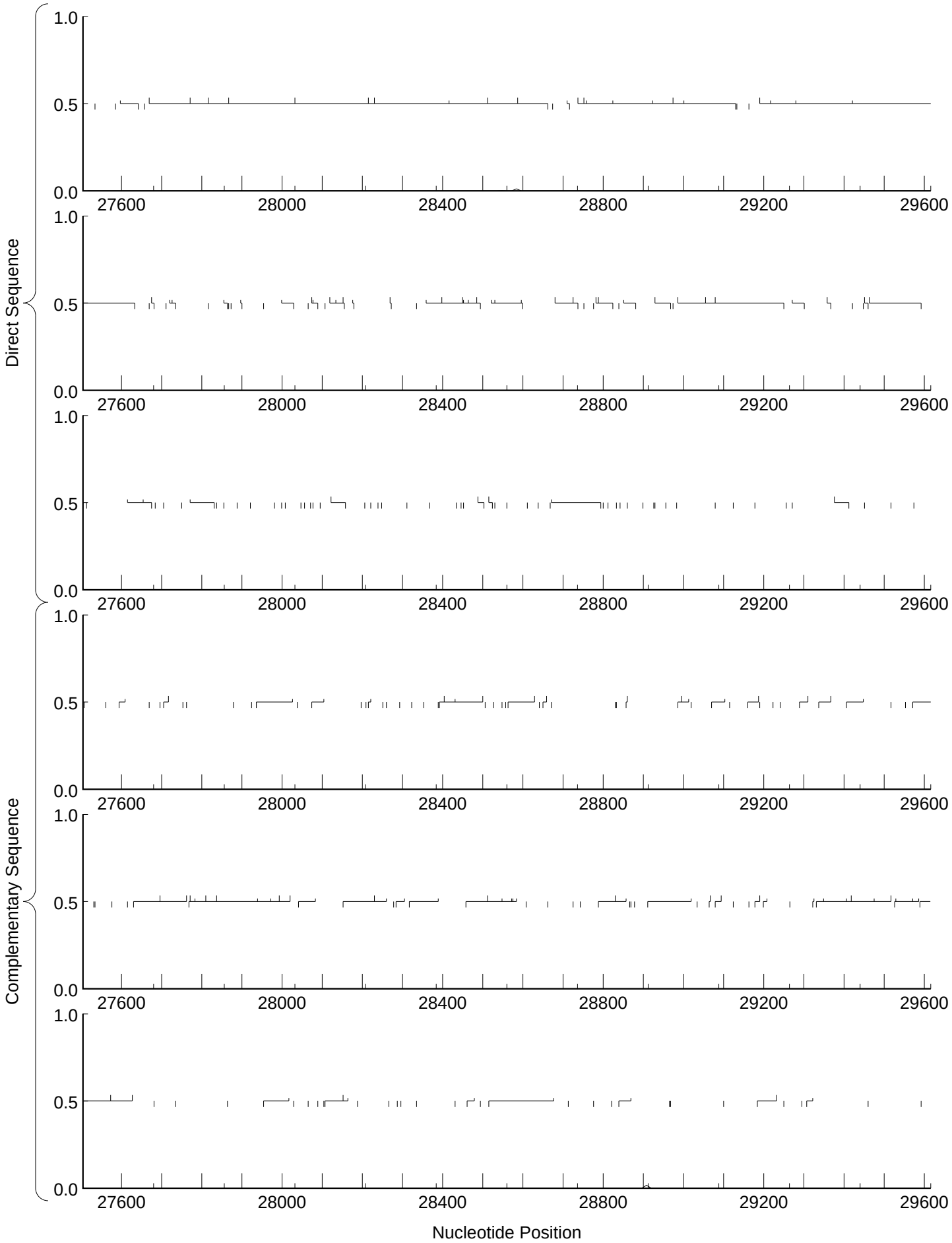
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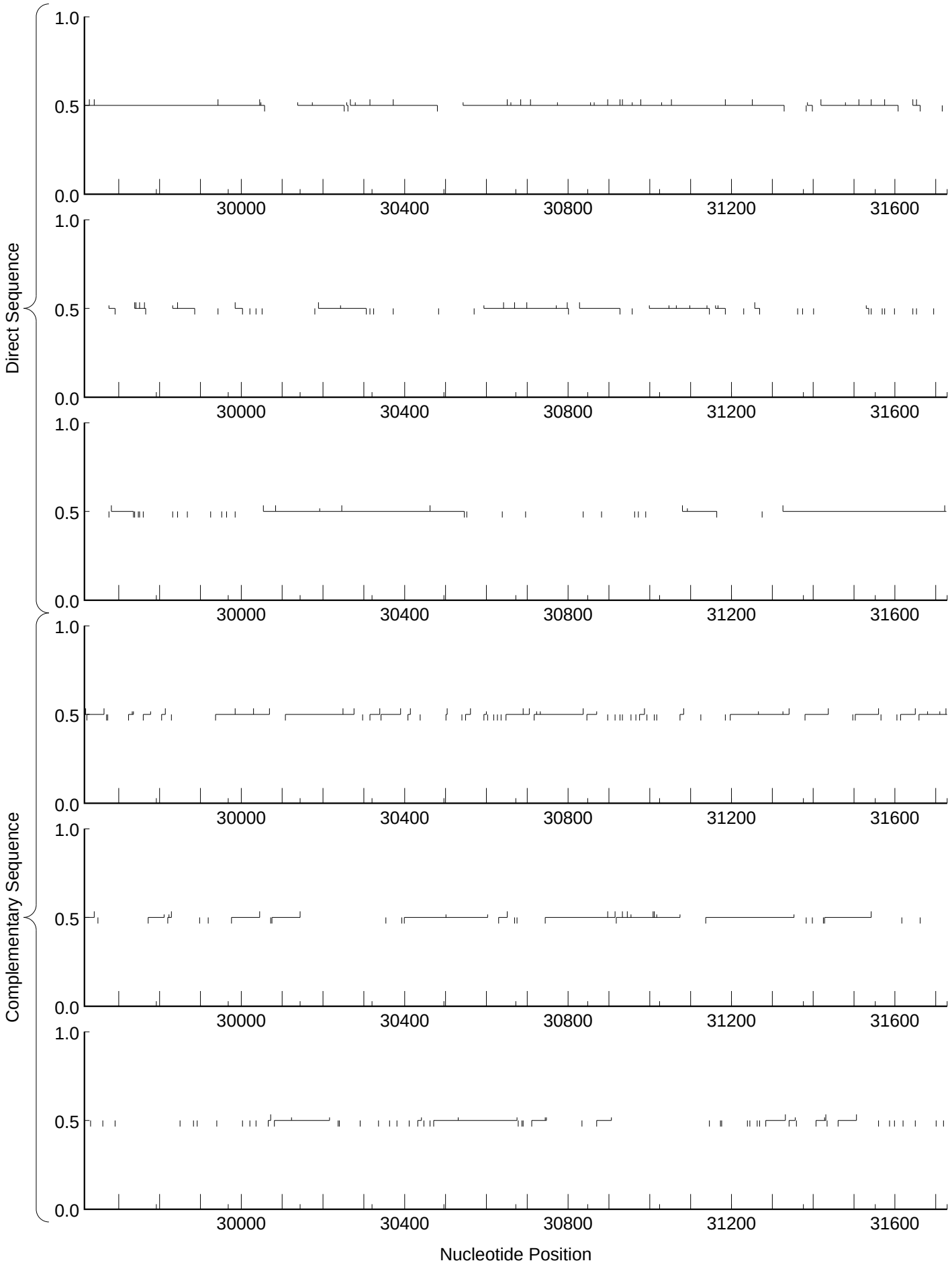
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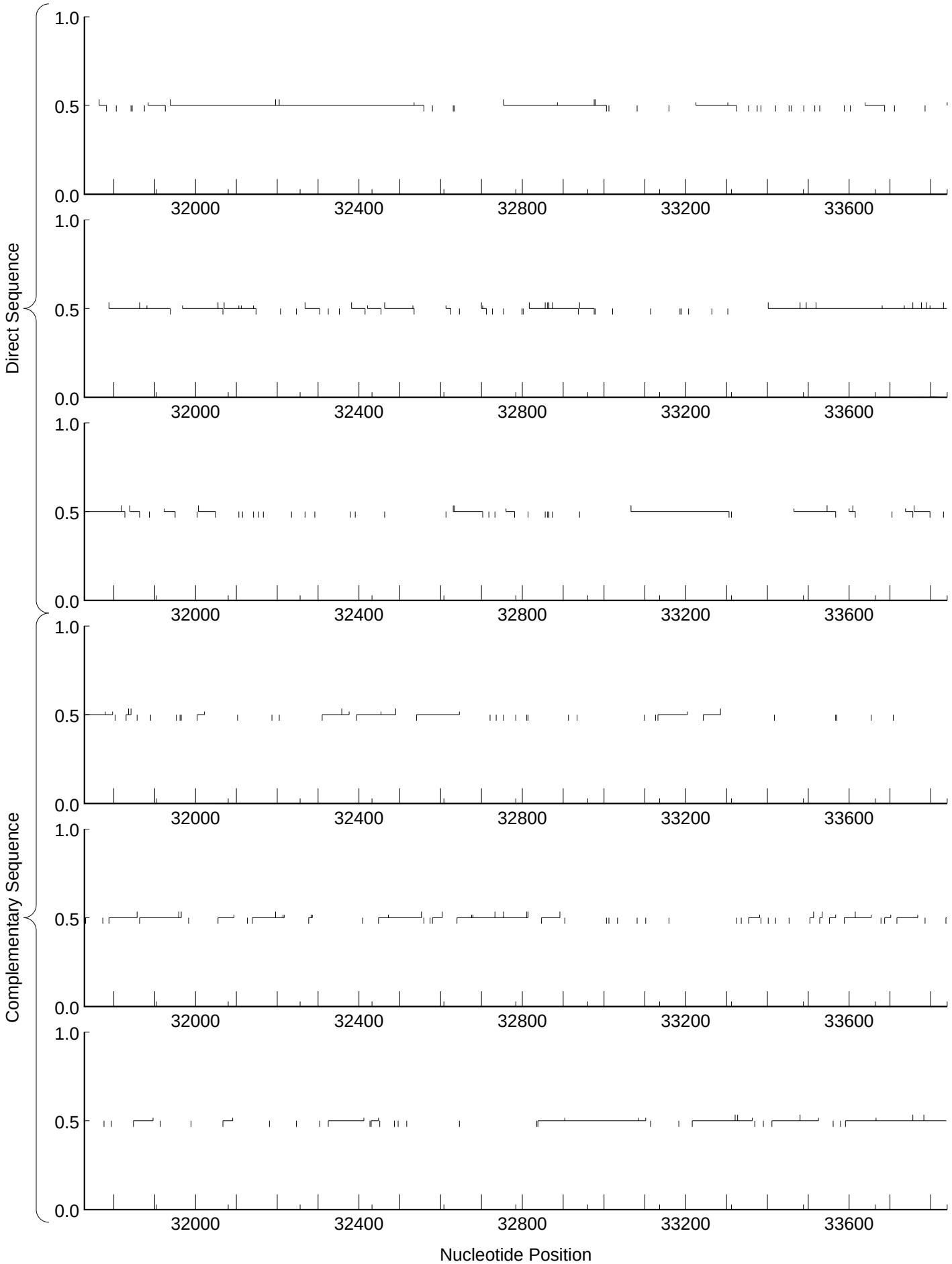
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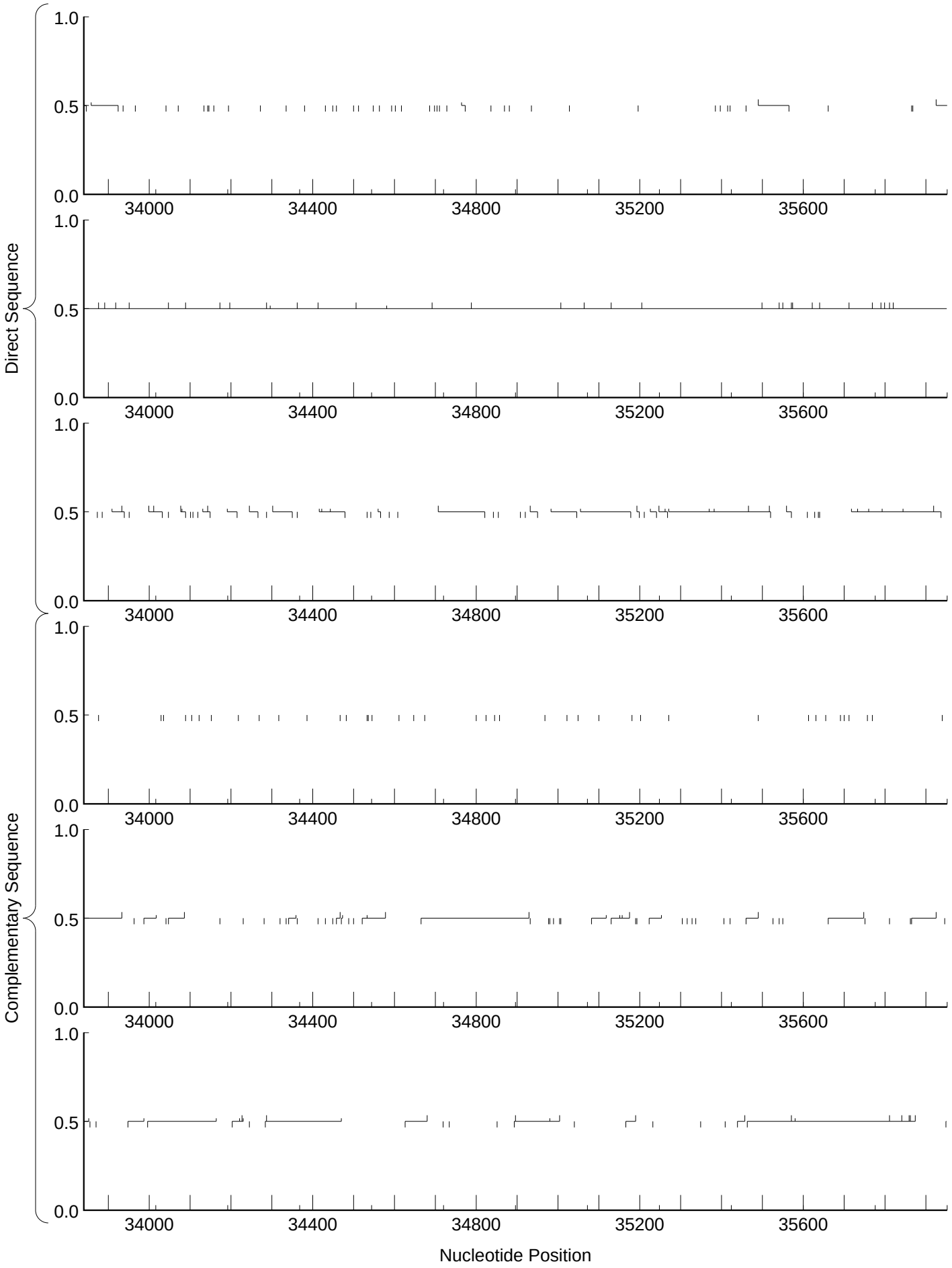
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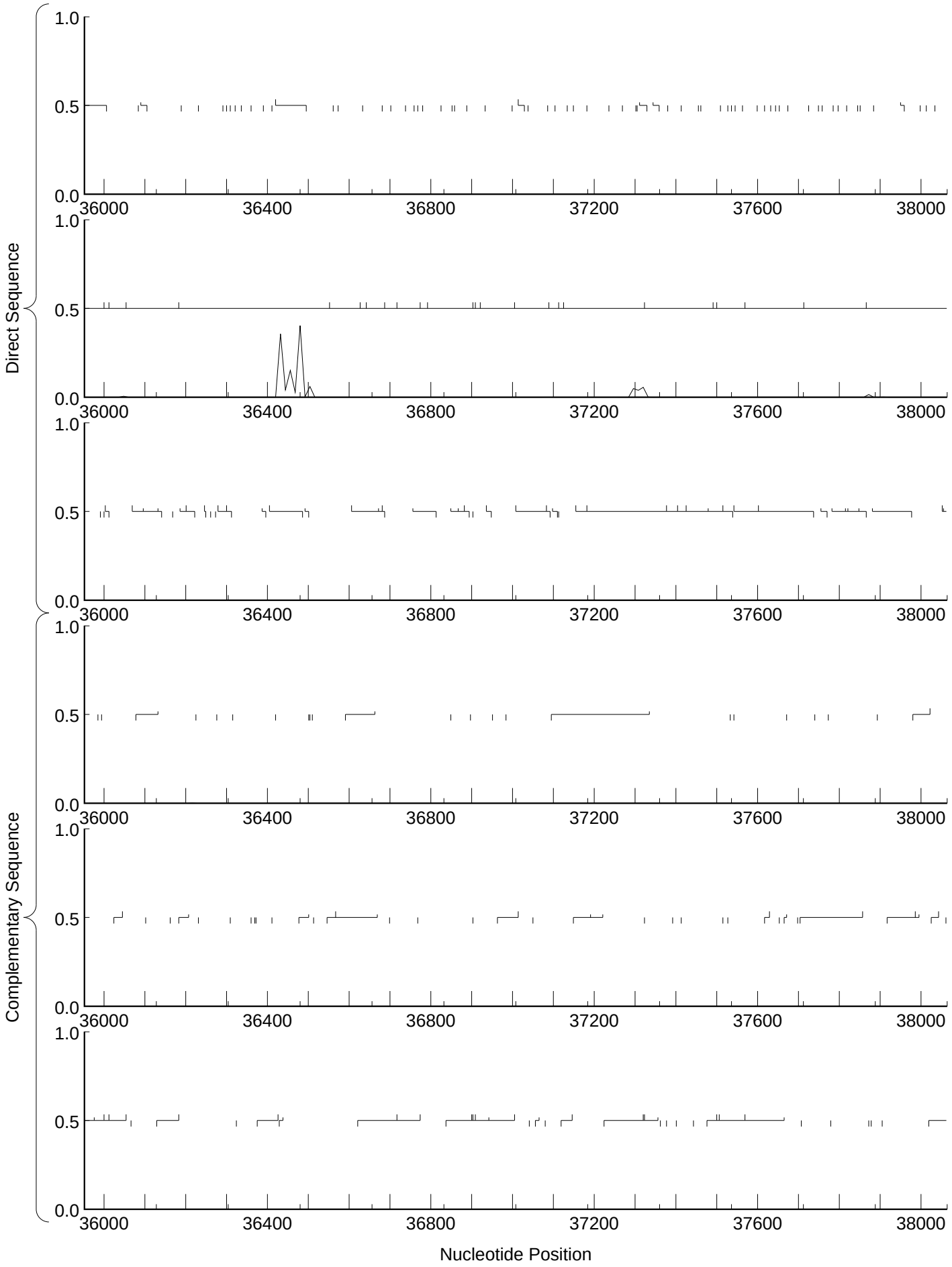
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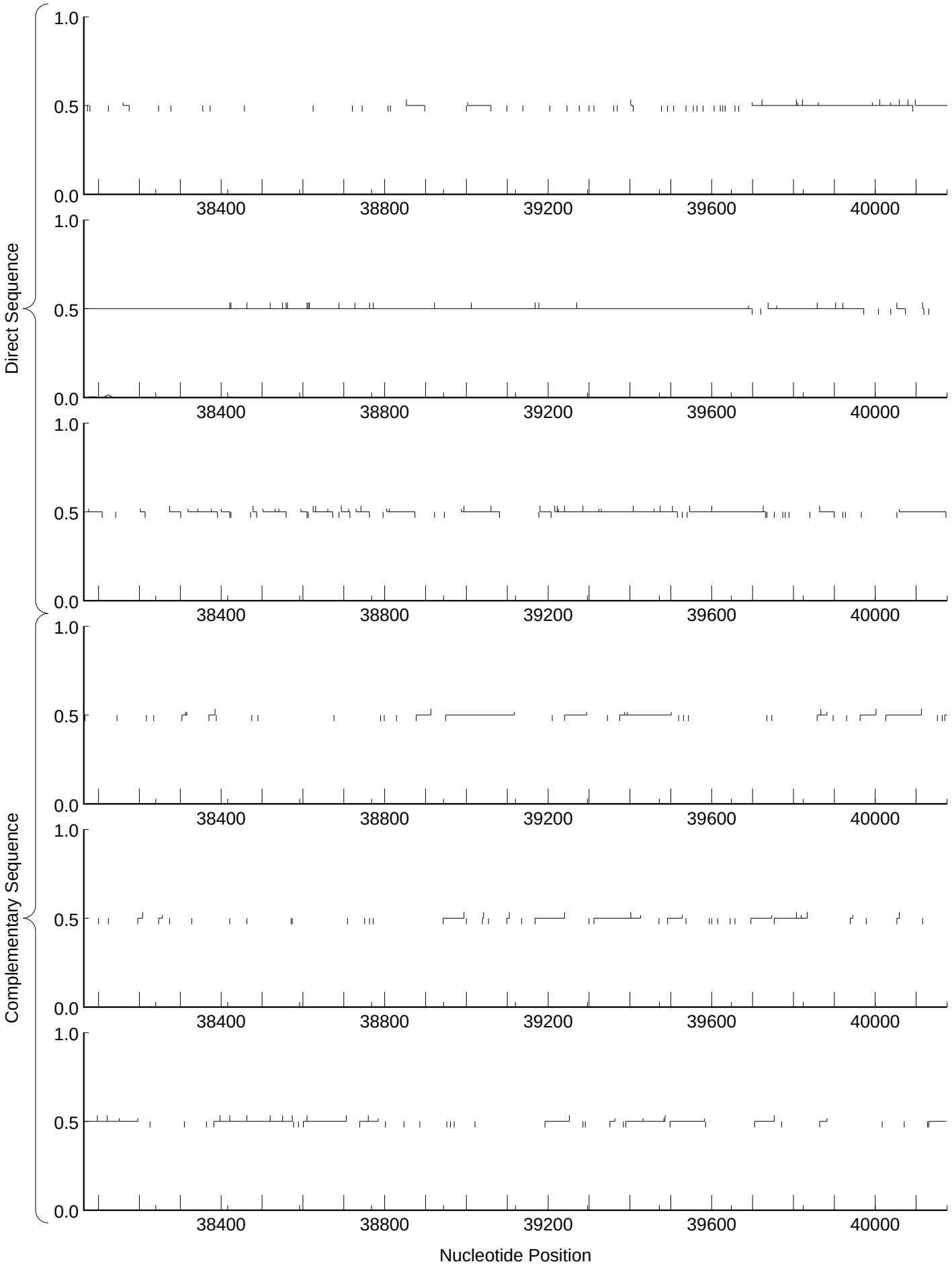
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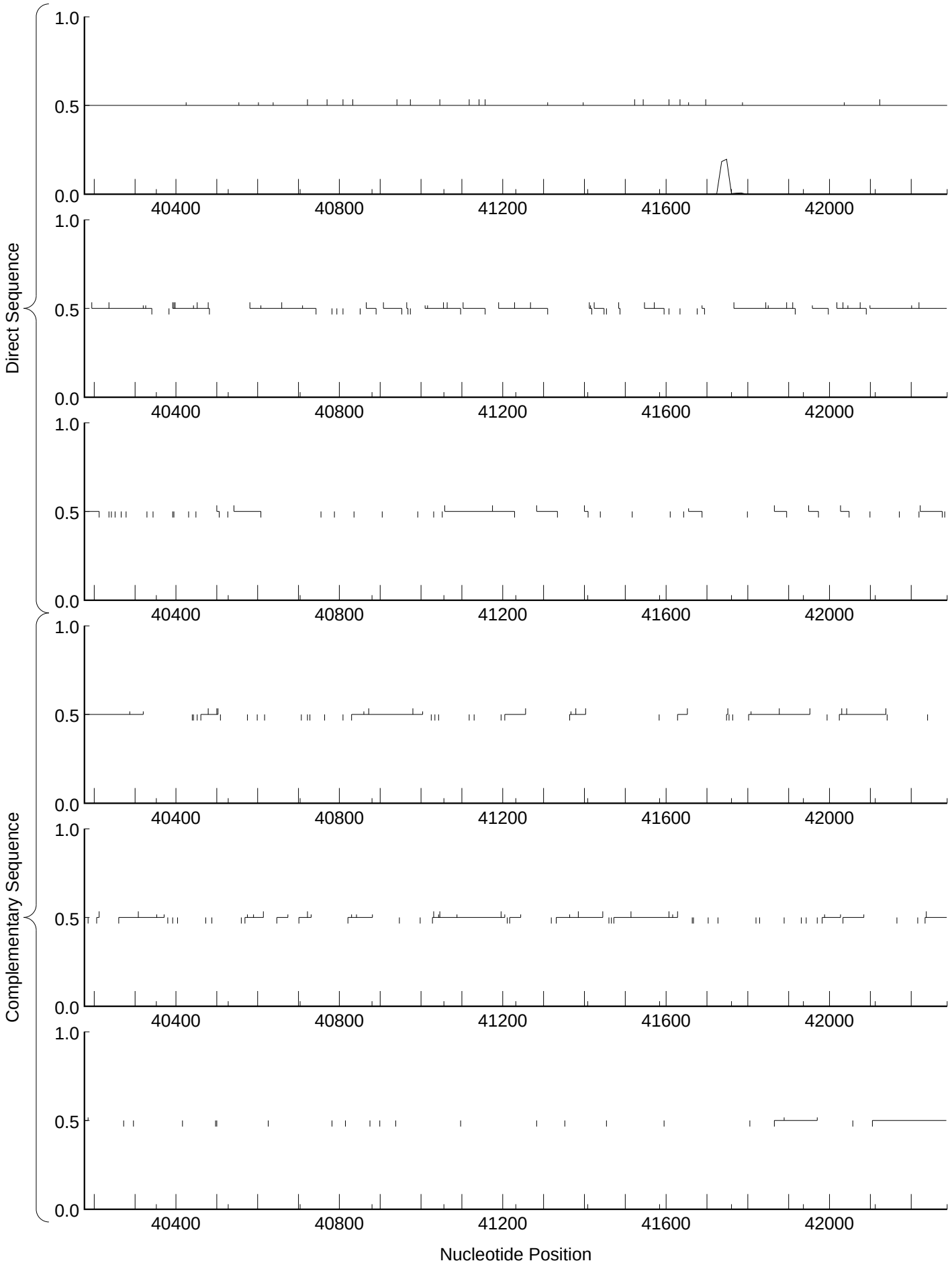
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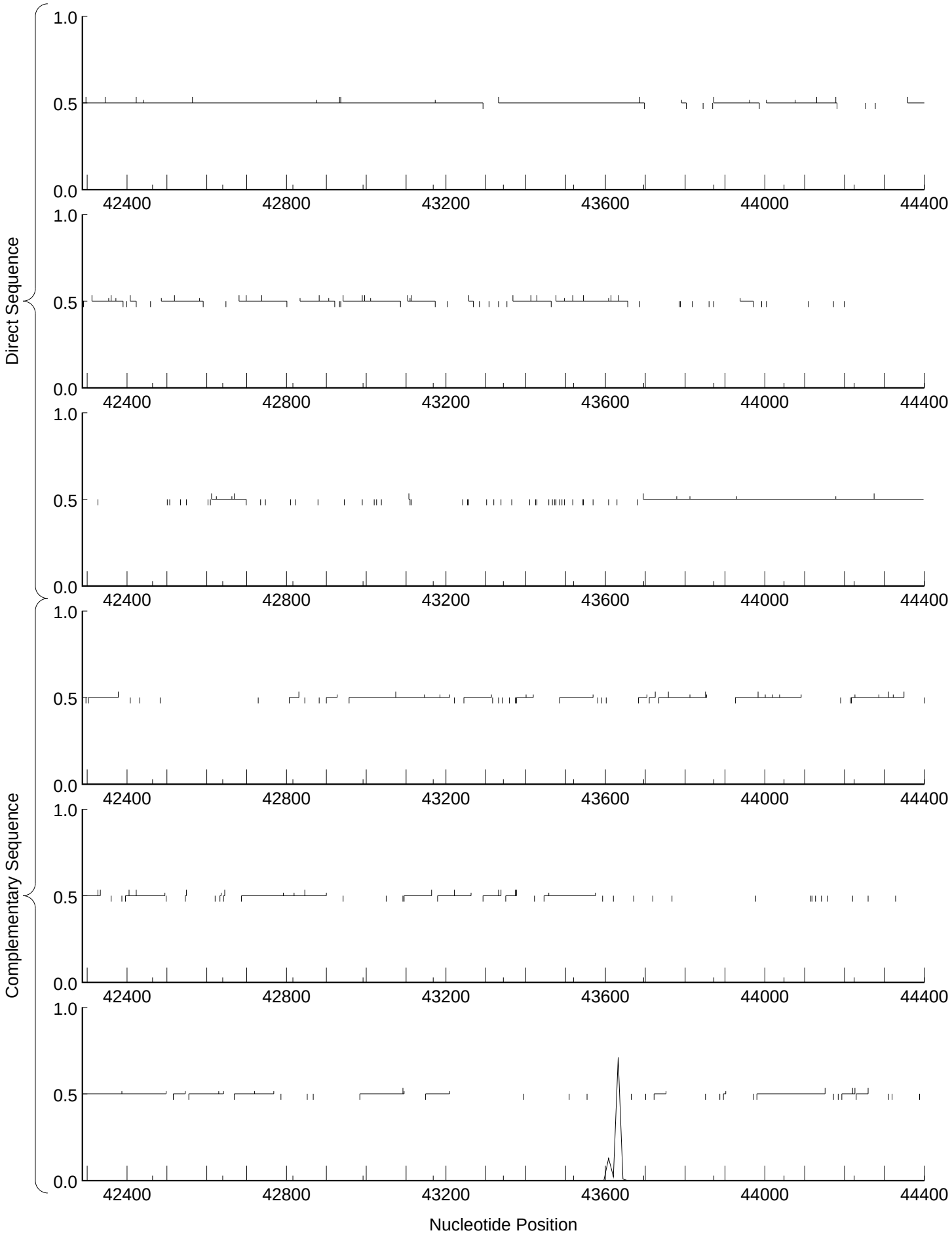
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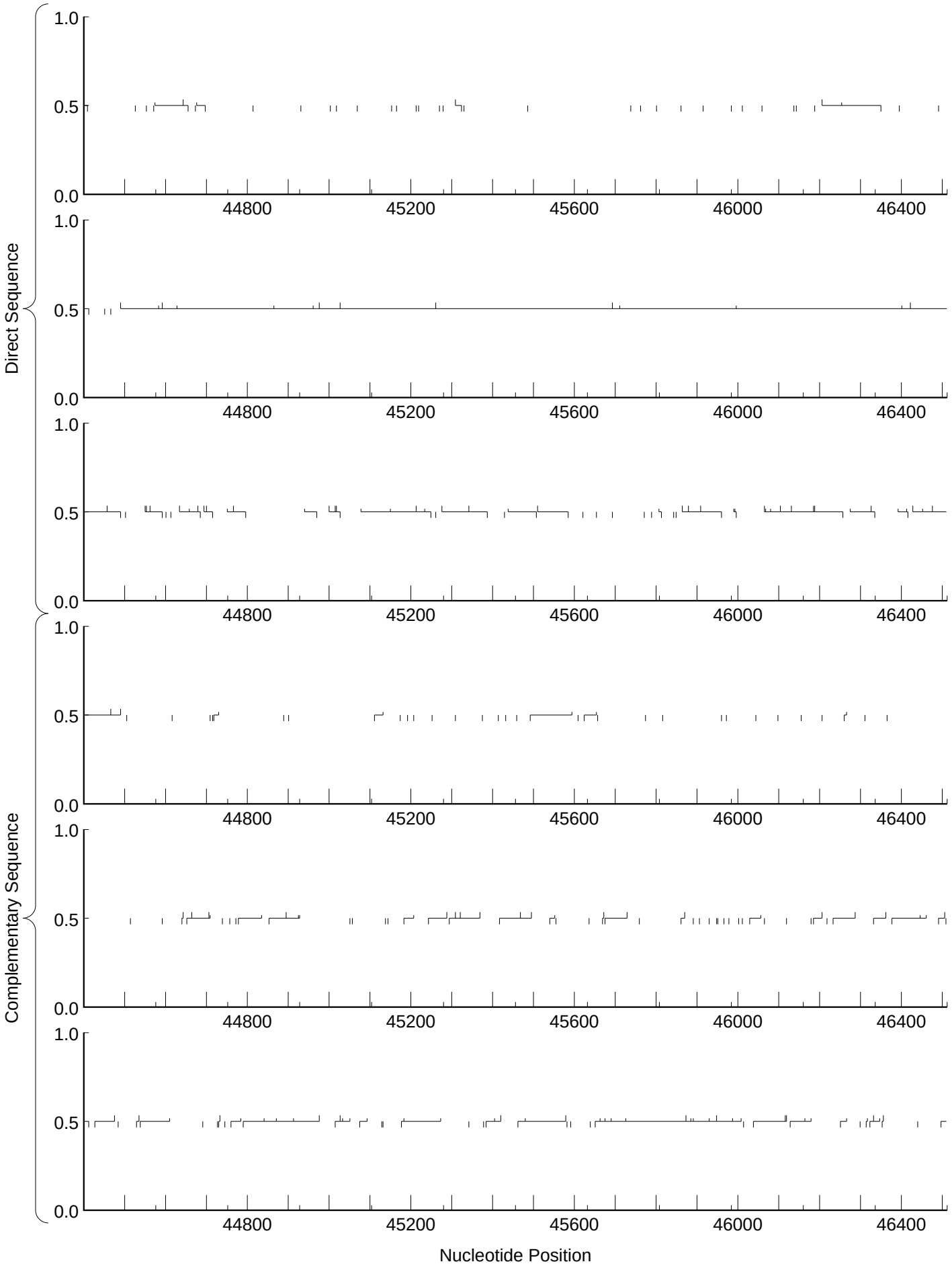
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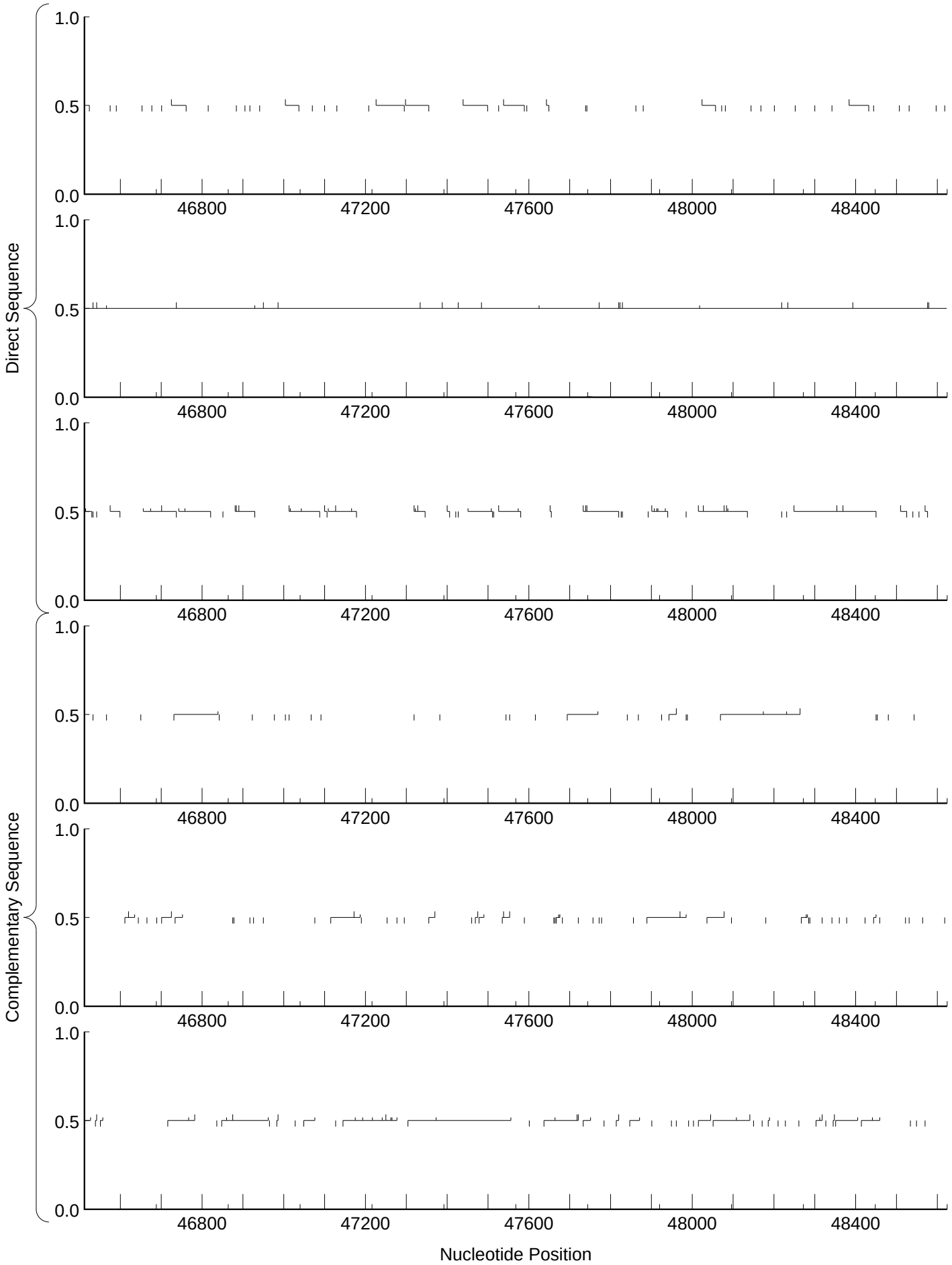
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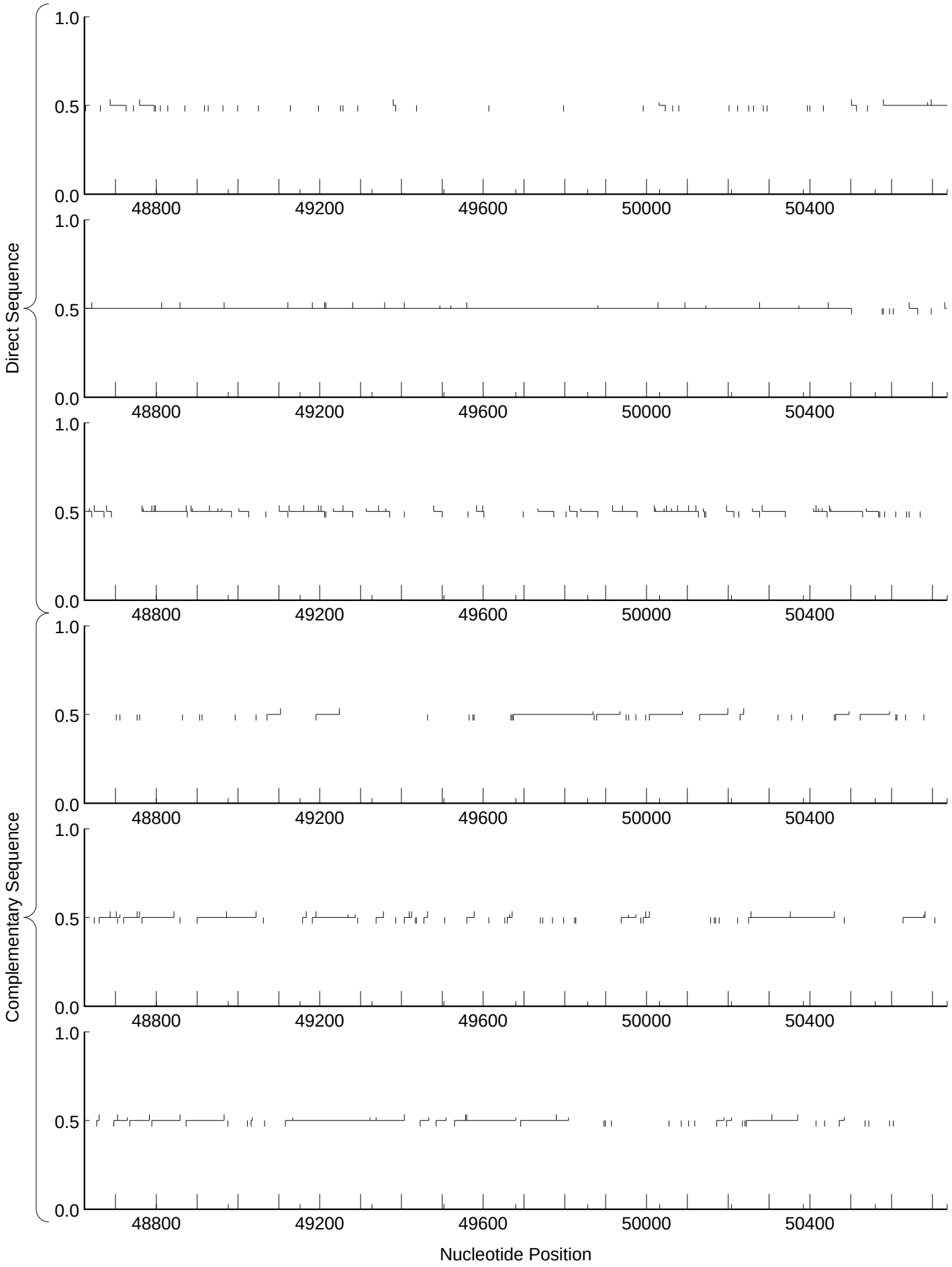
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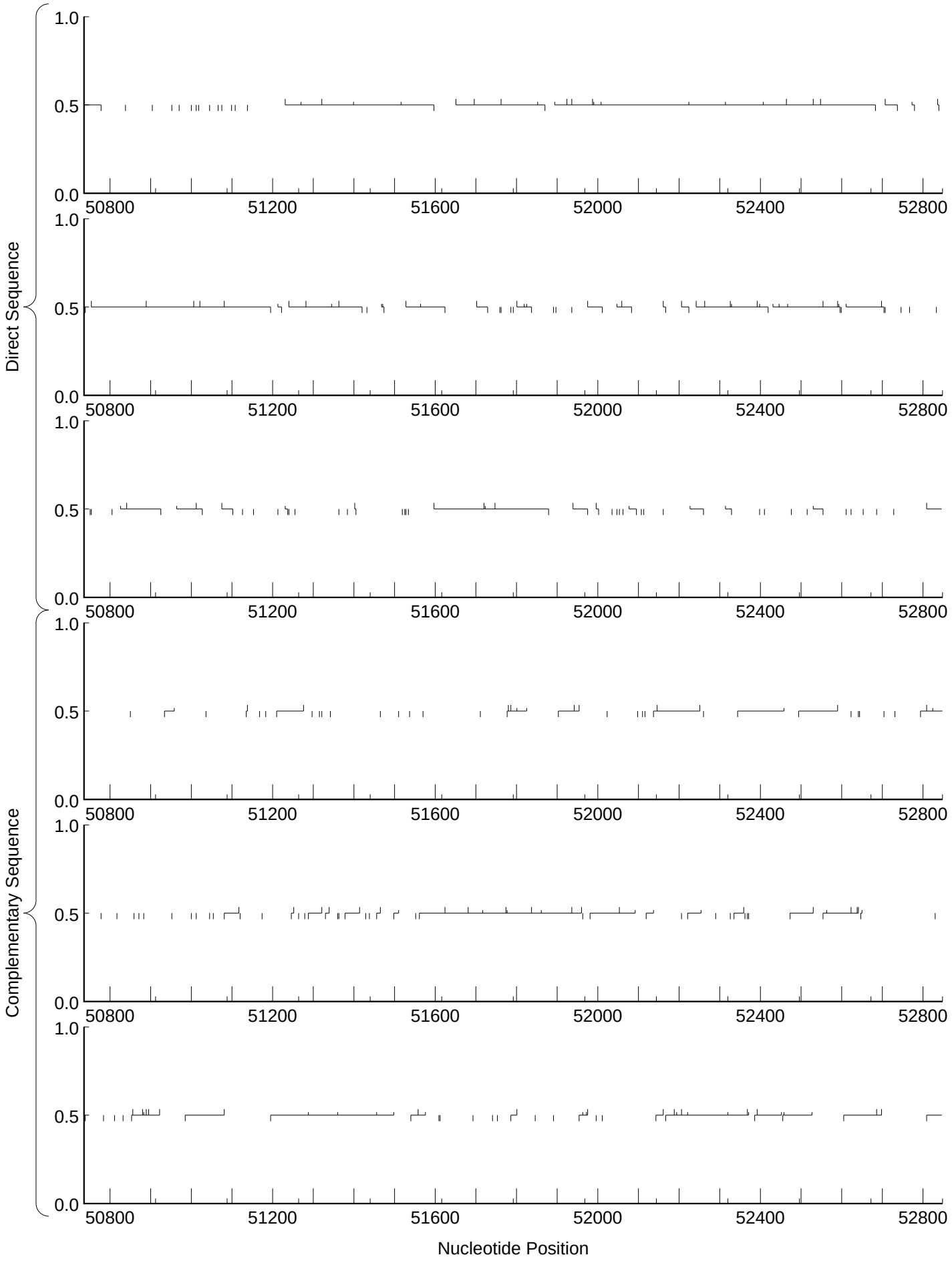
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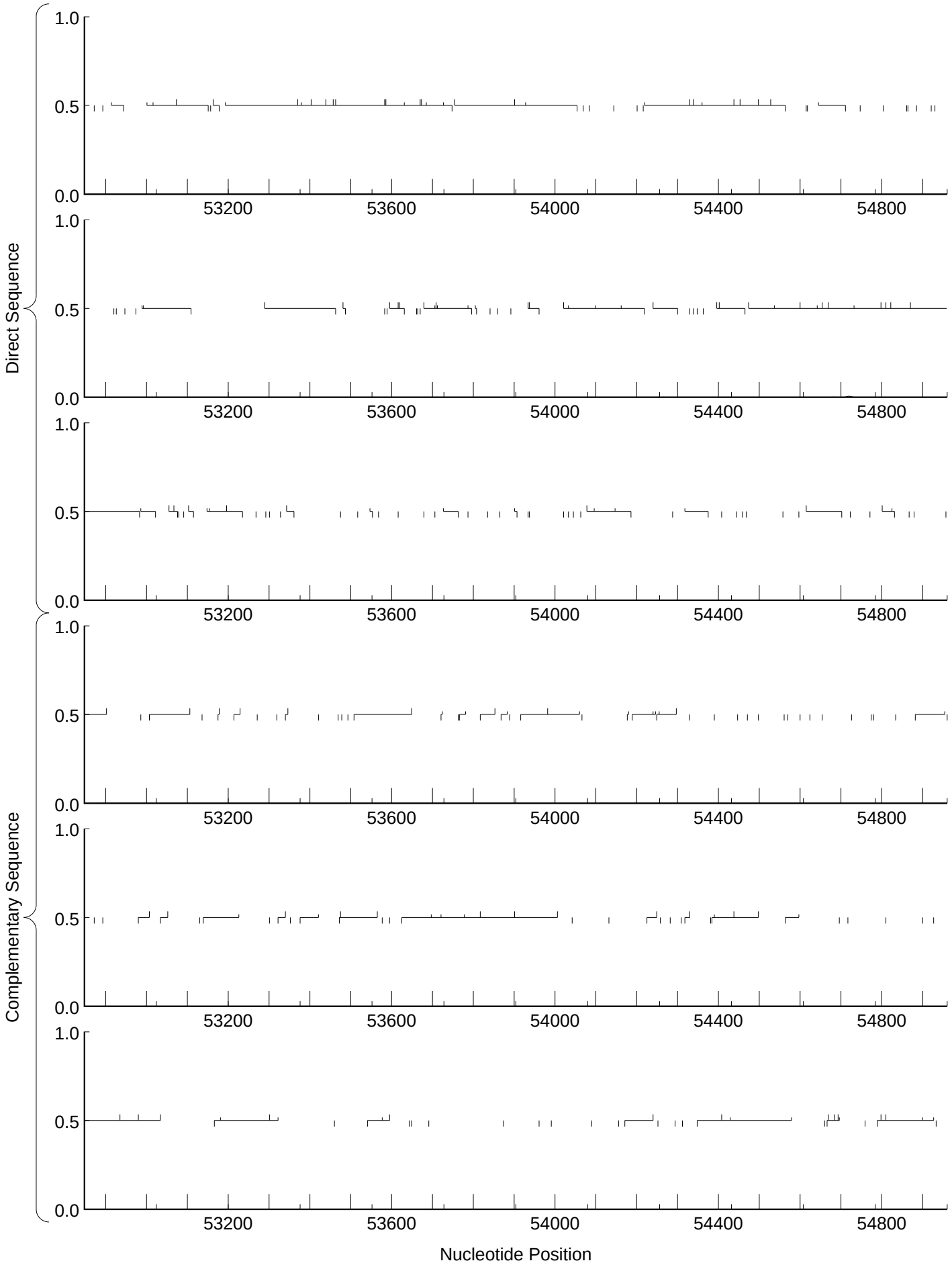
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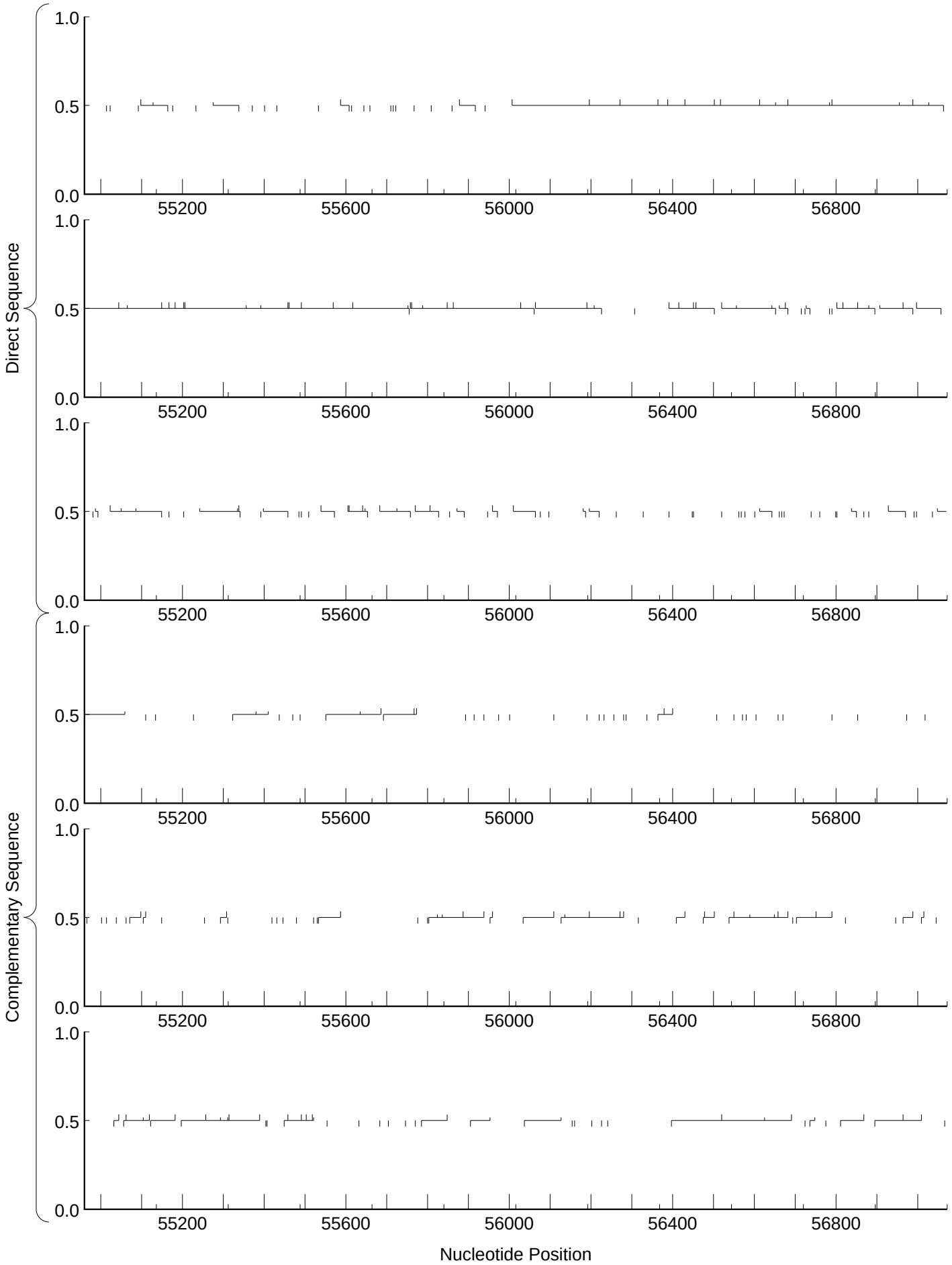
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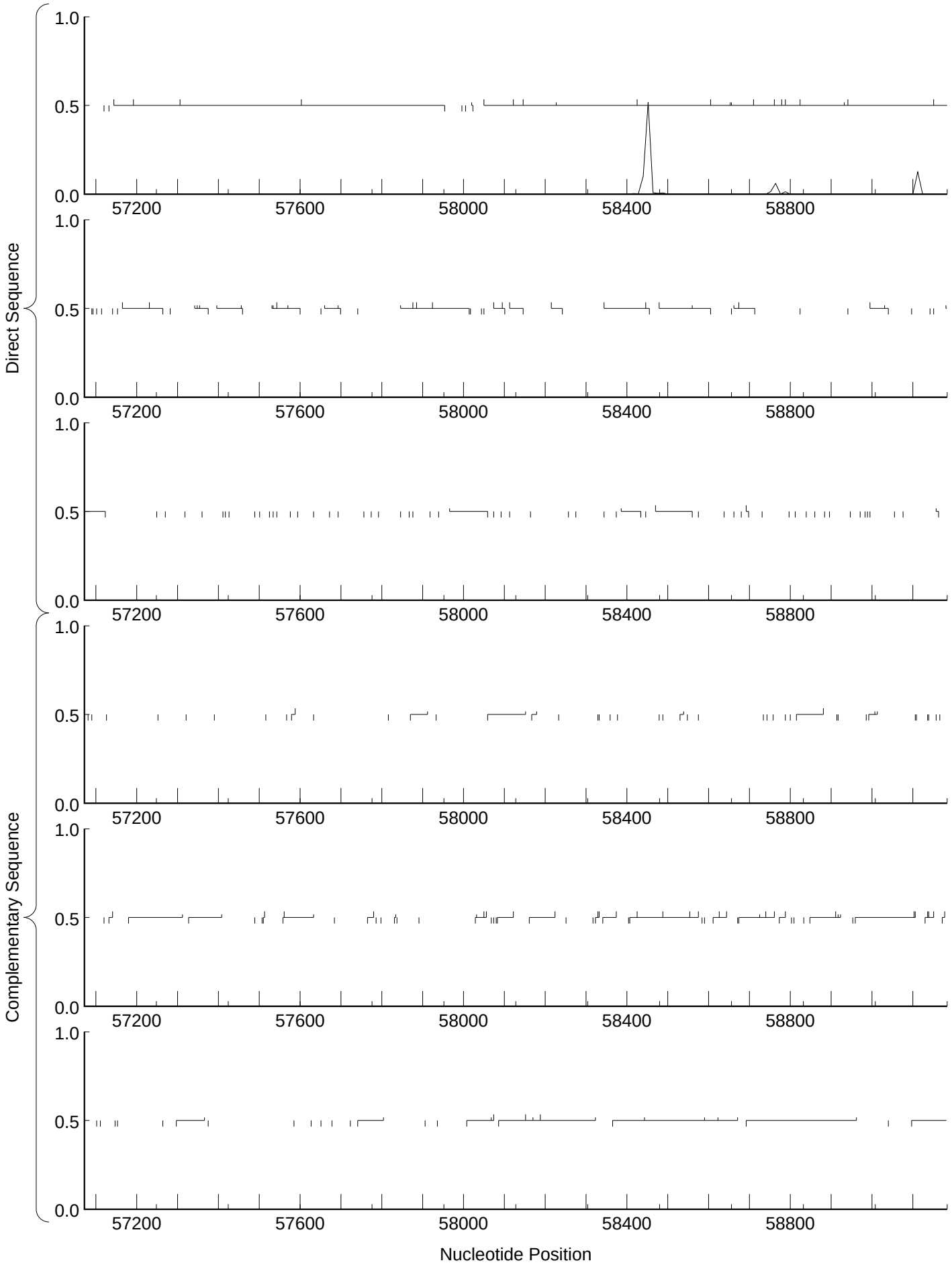
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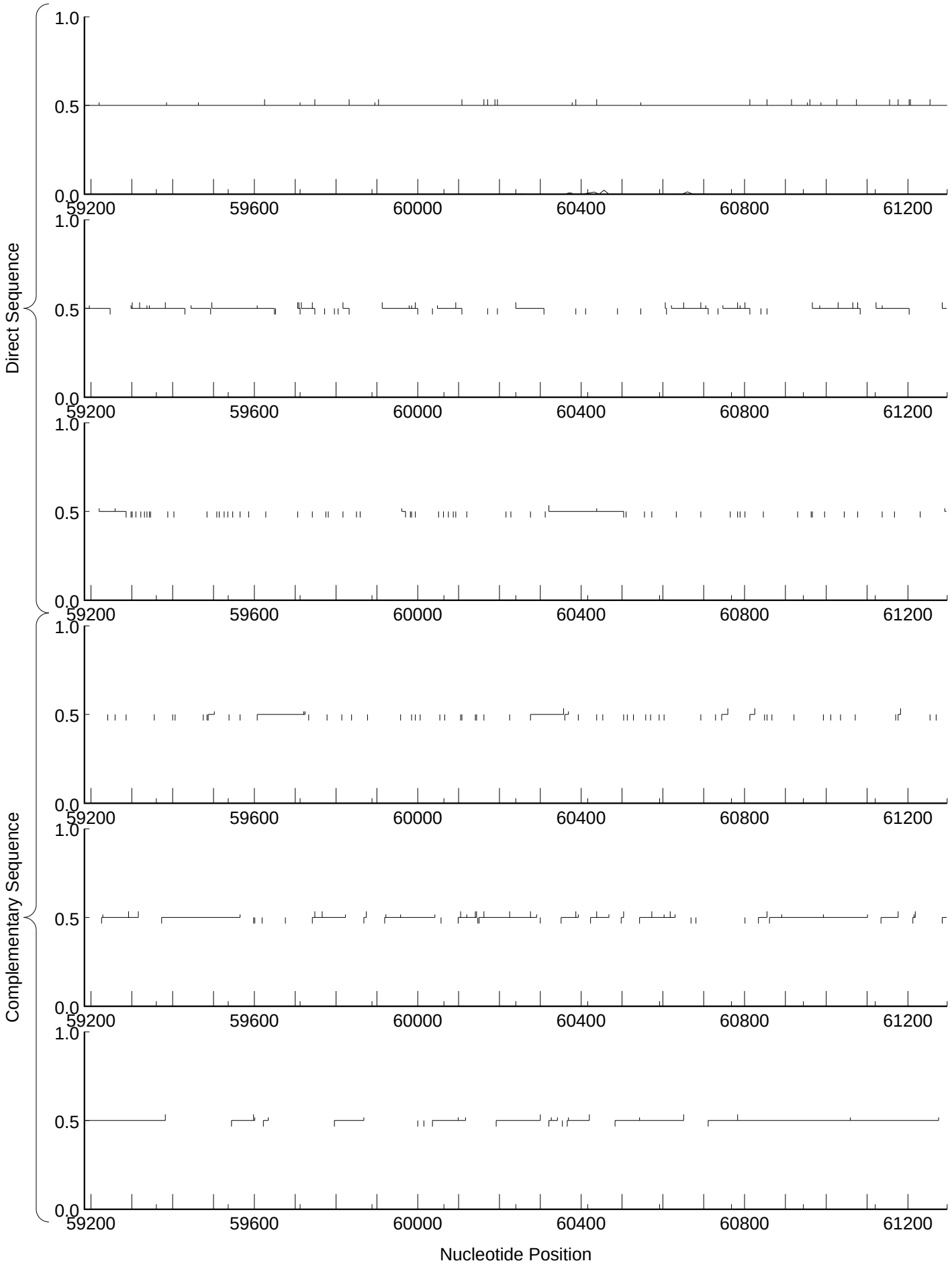
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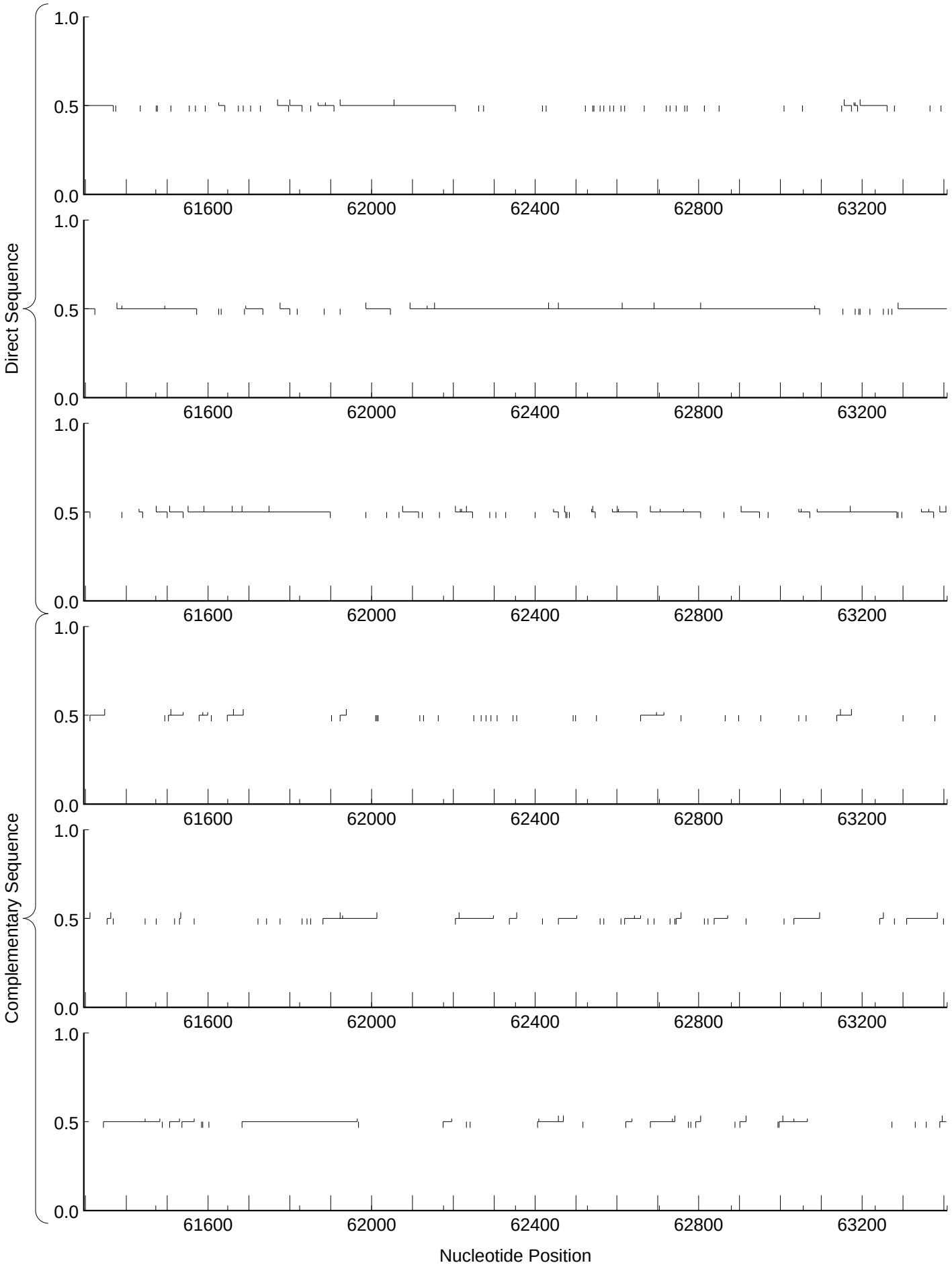
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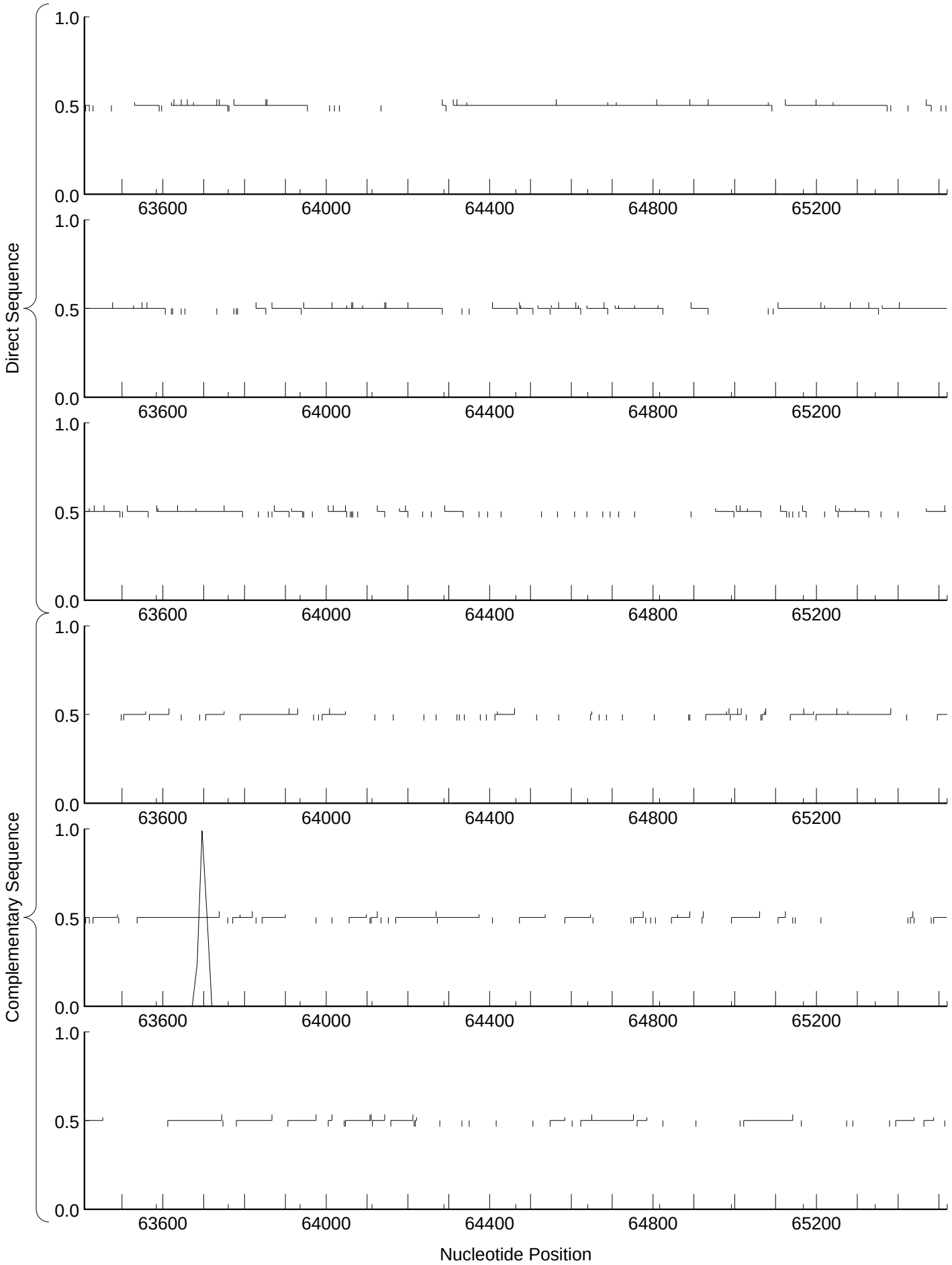
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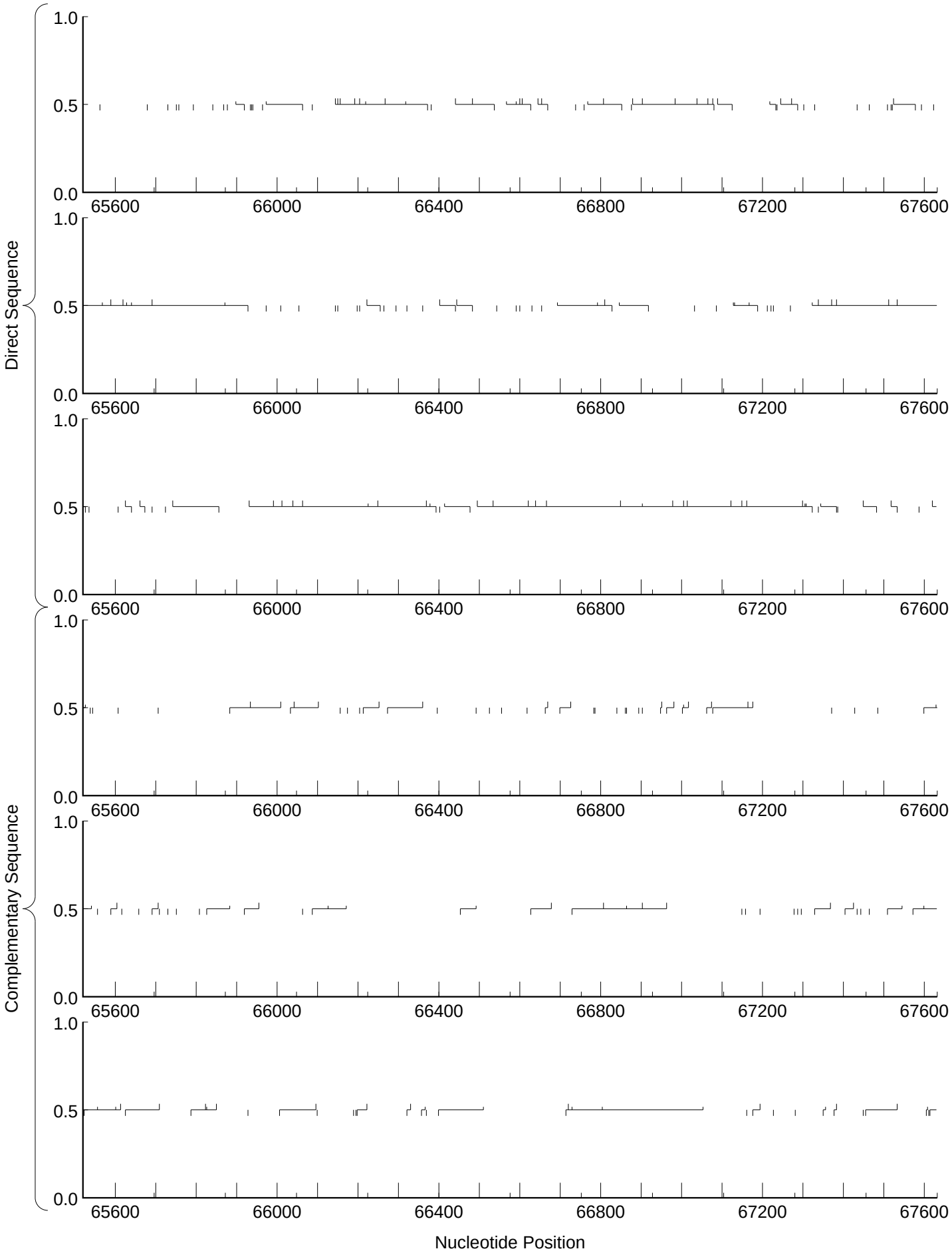
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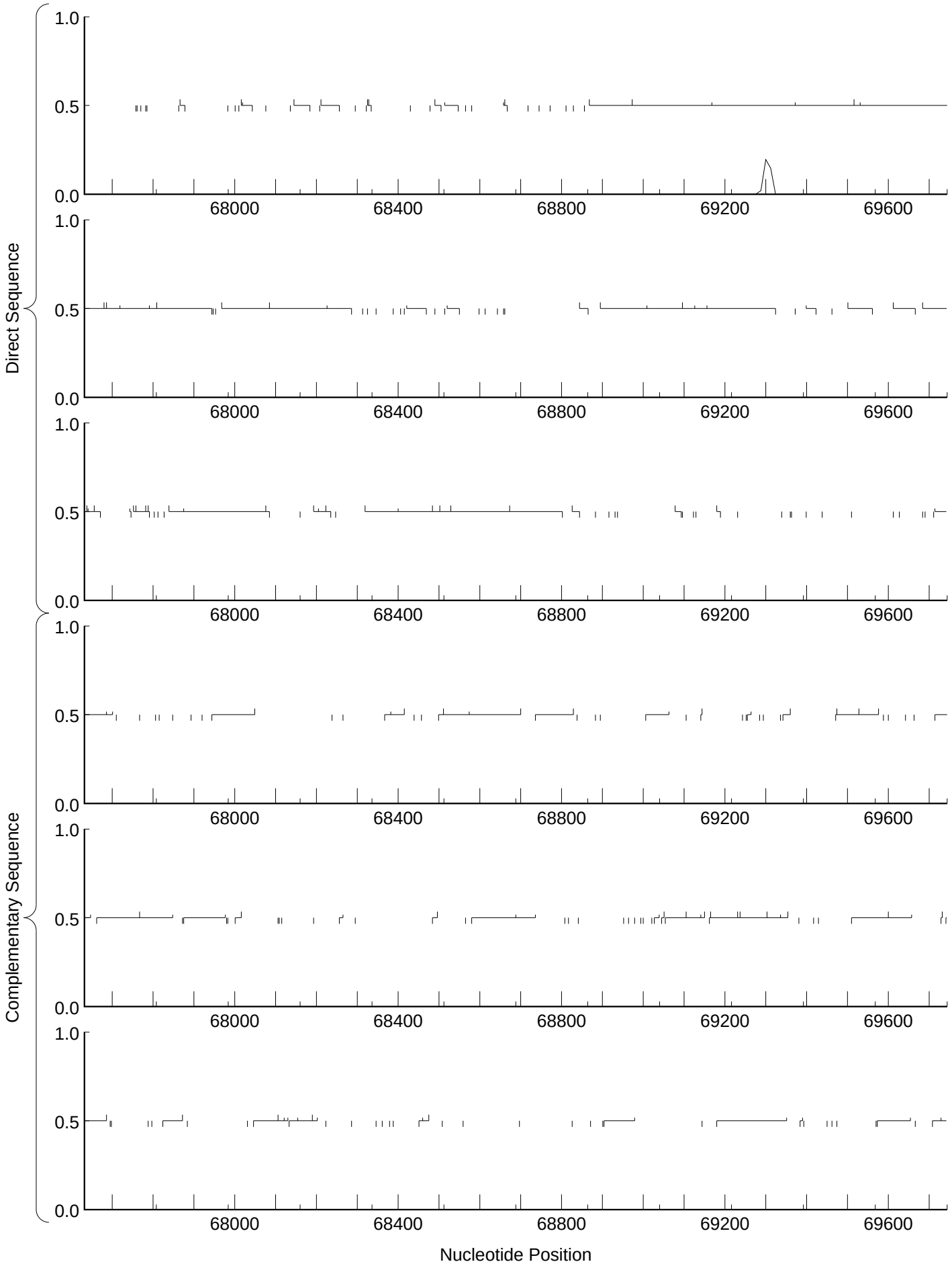
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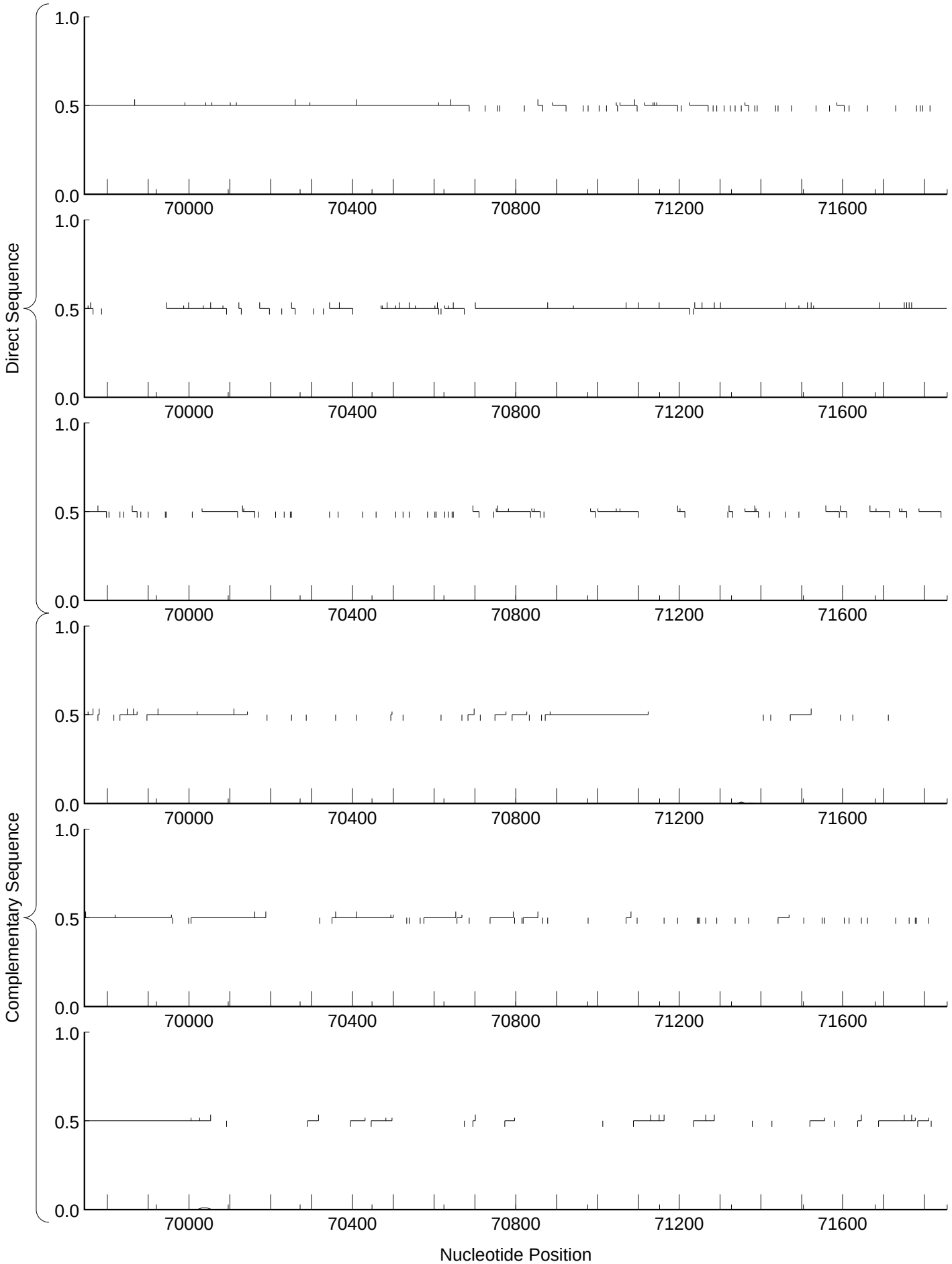
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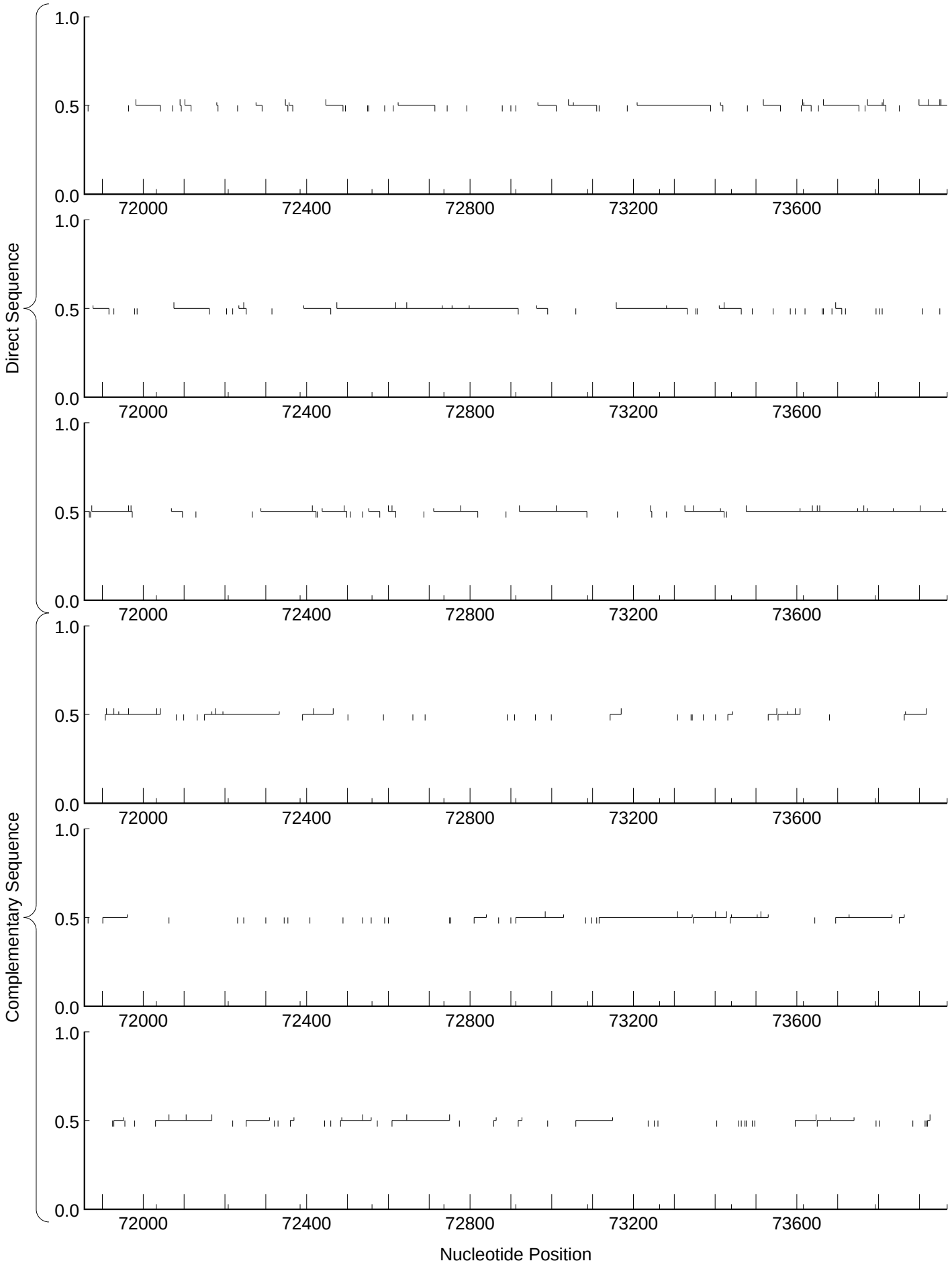
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 34/63



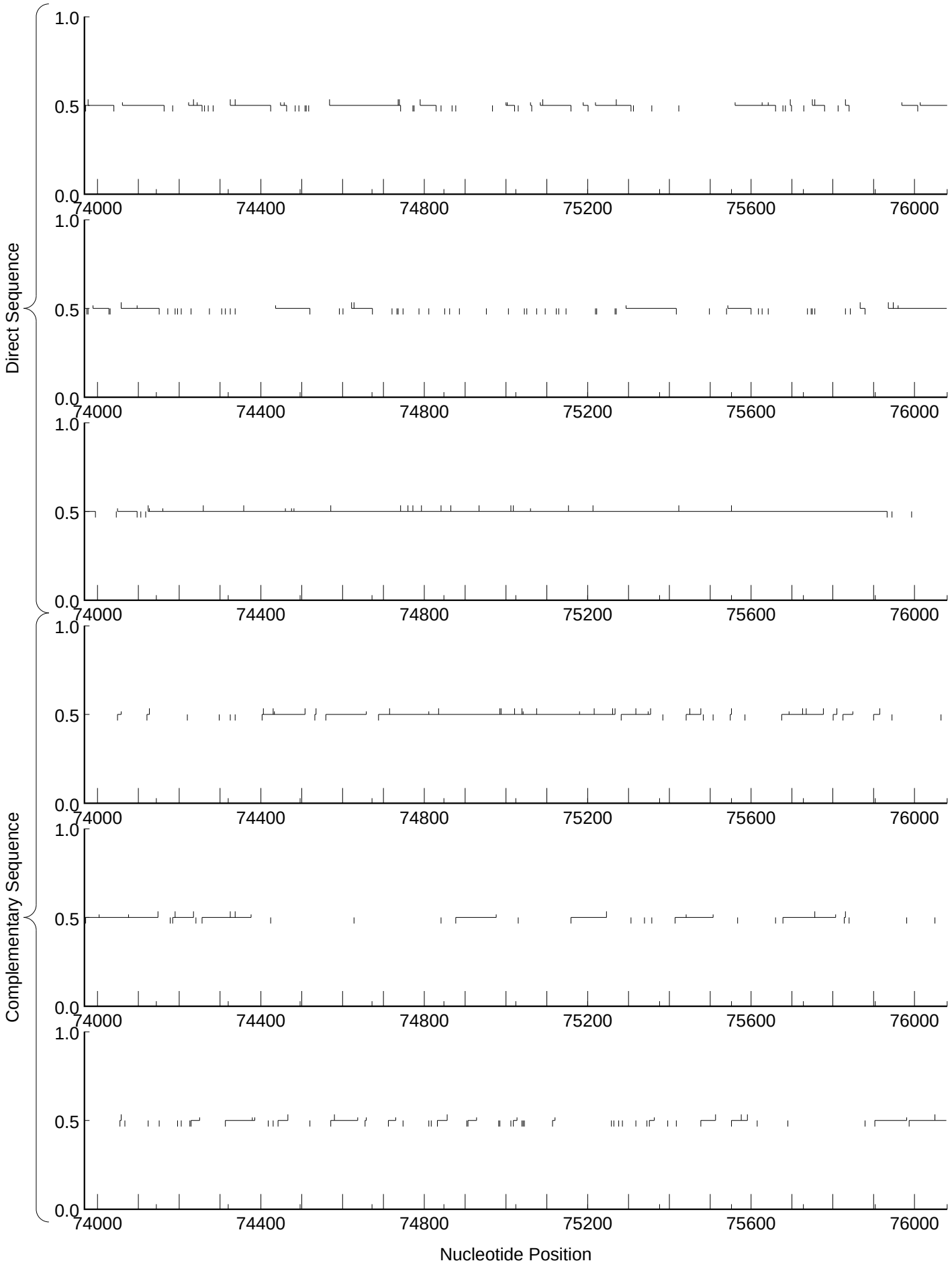
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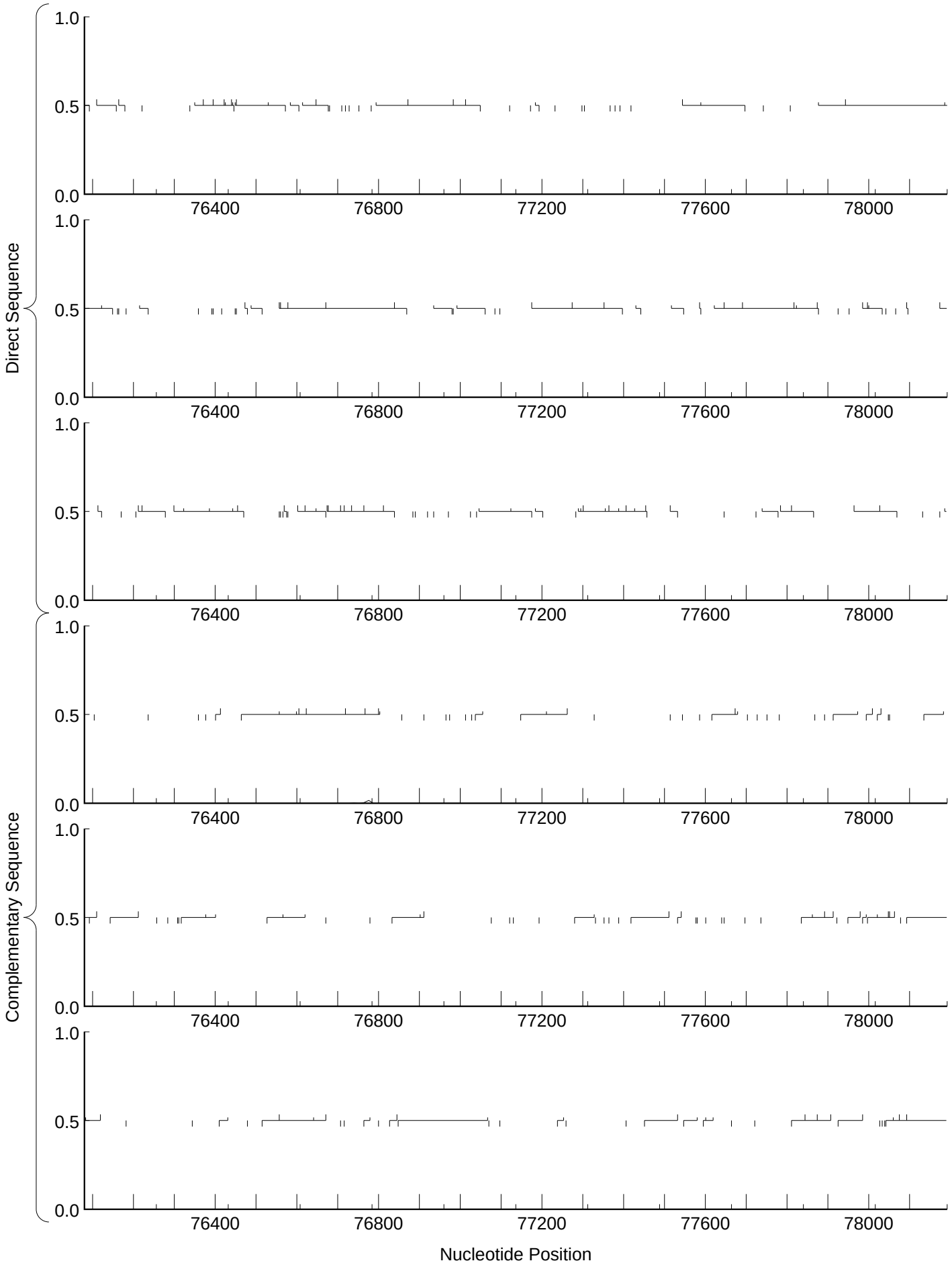
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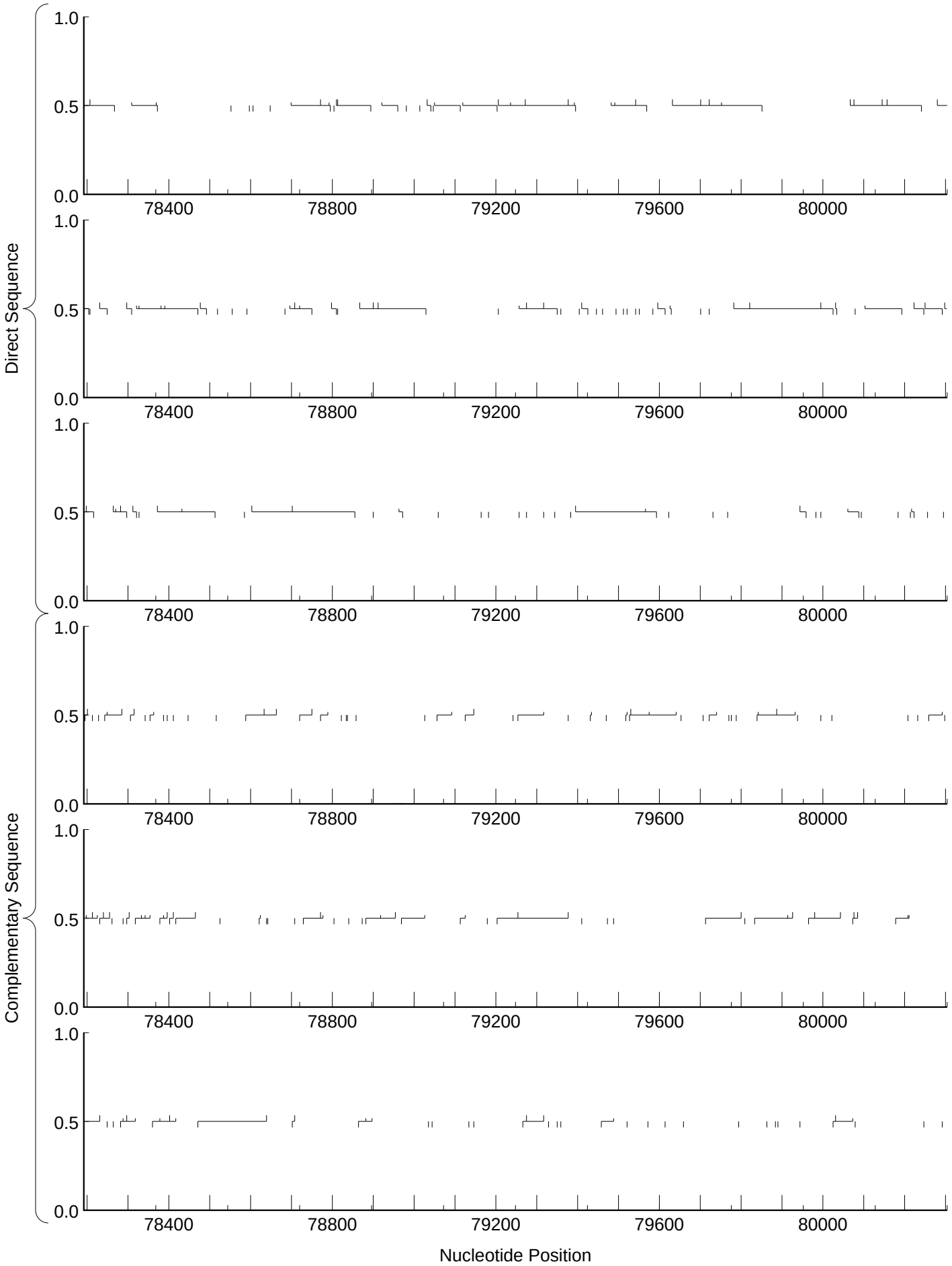
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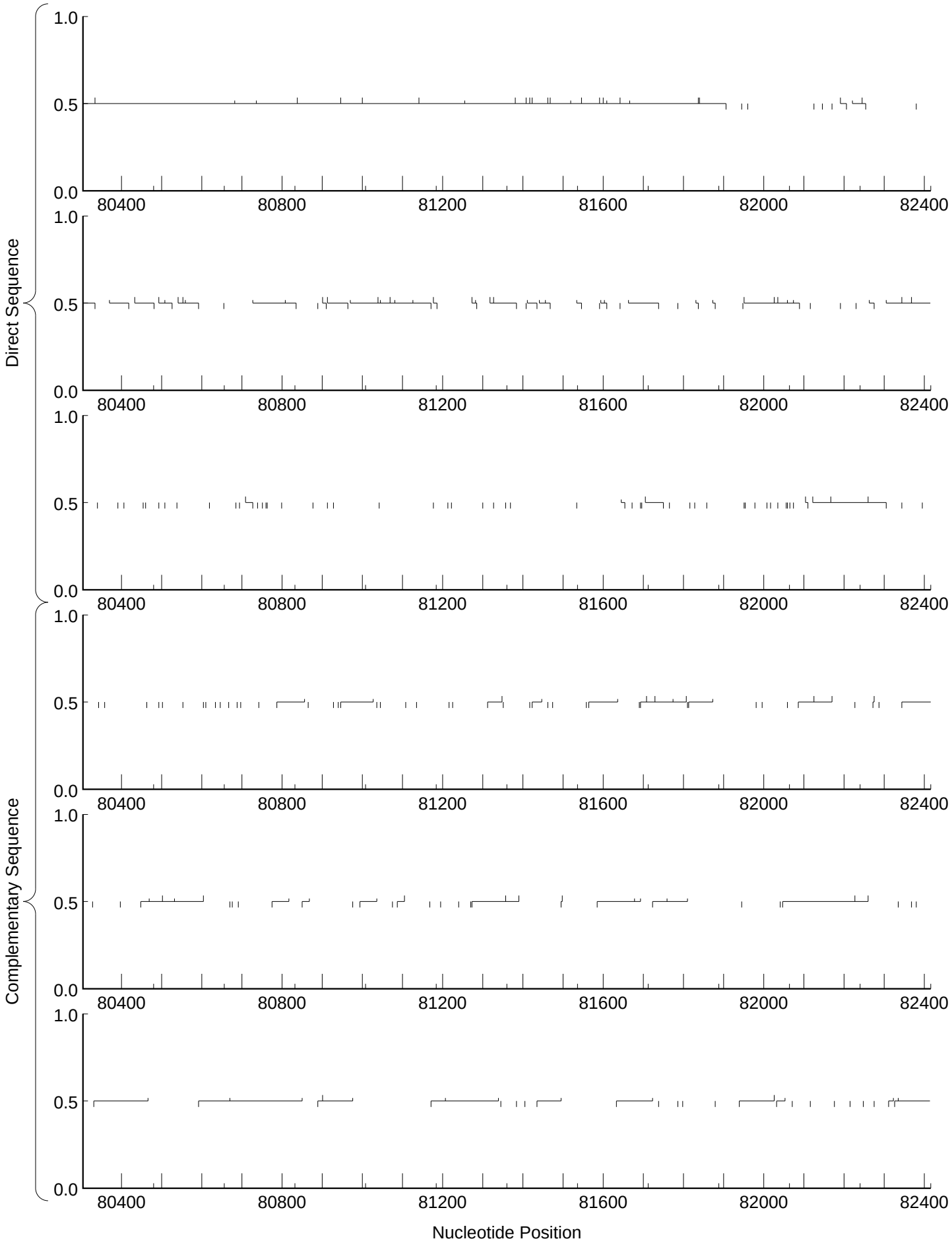
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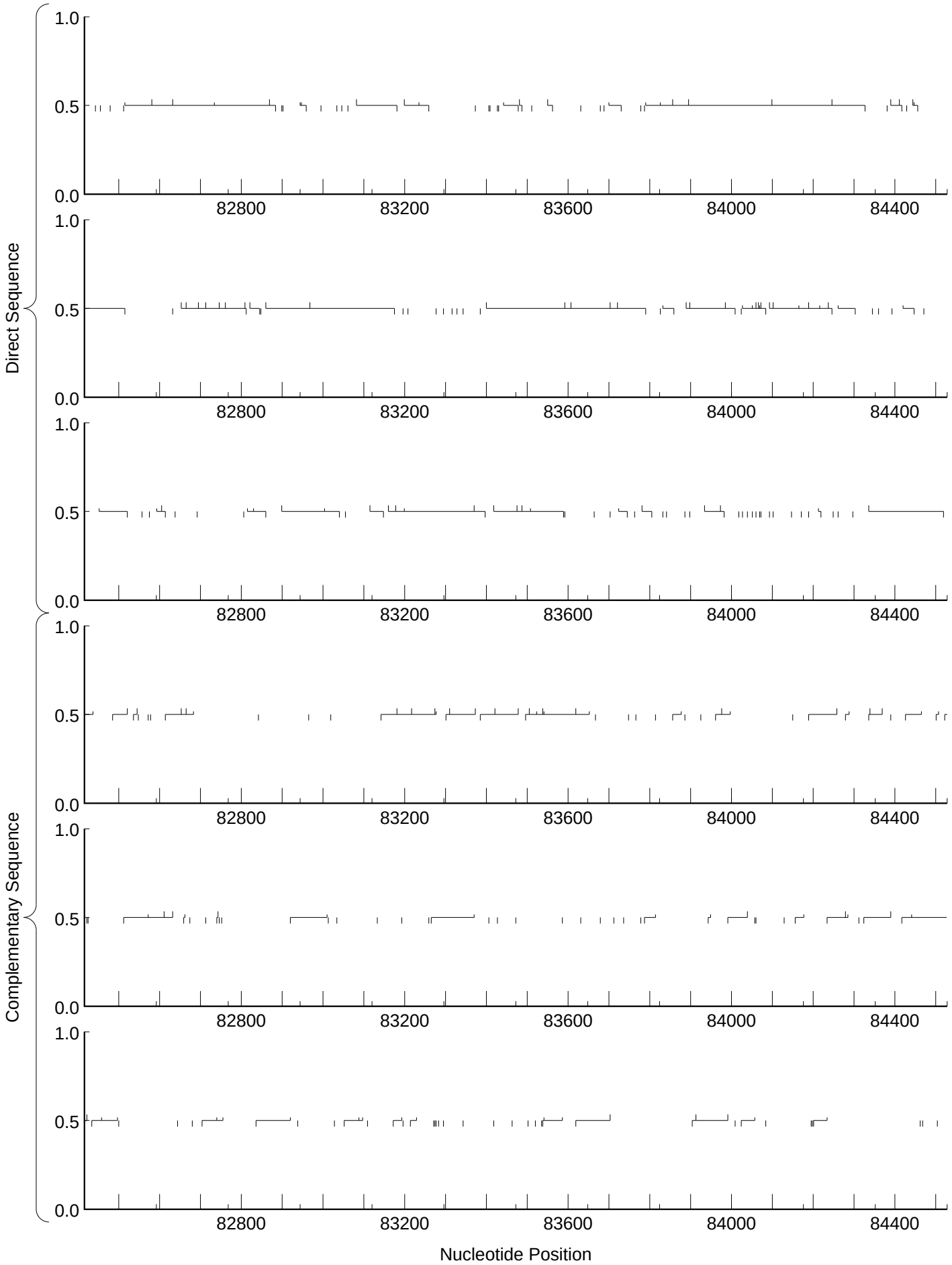
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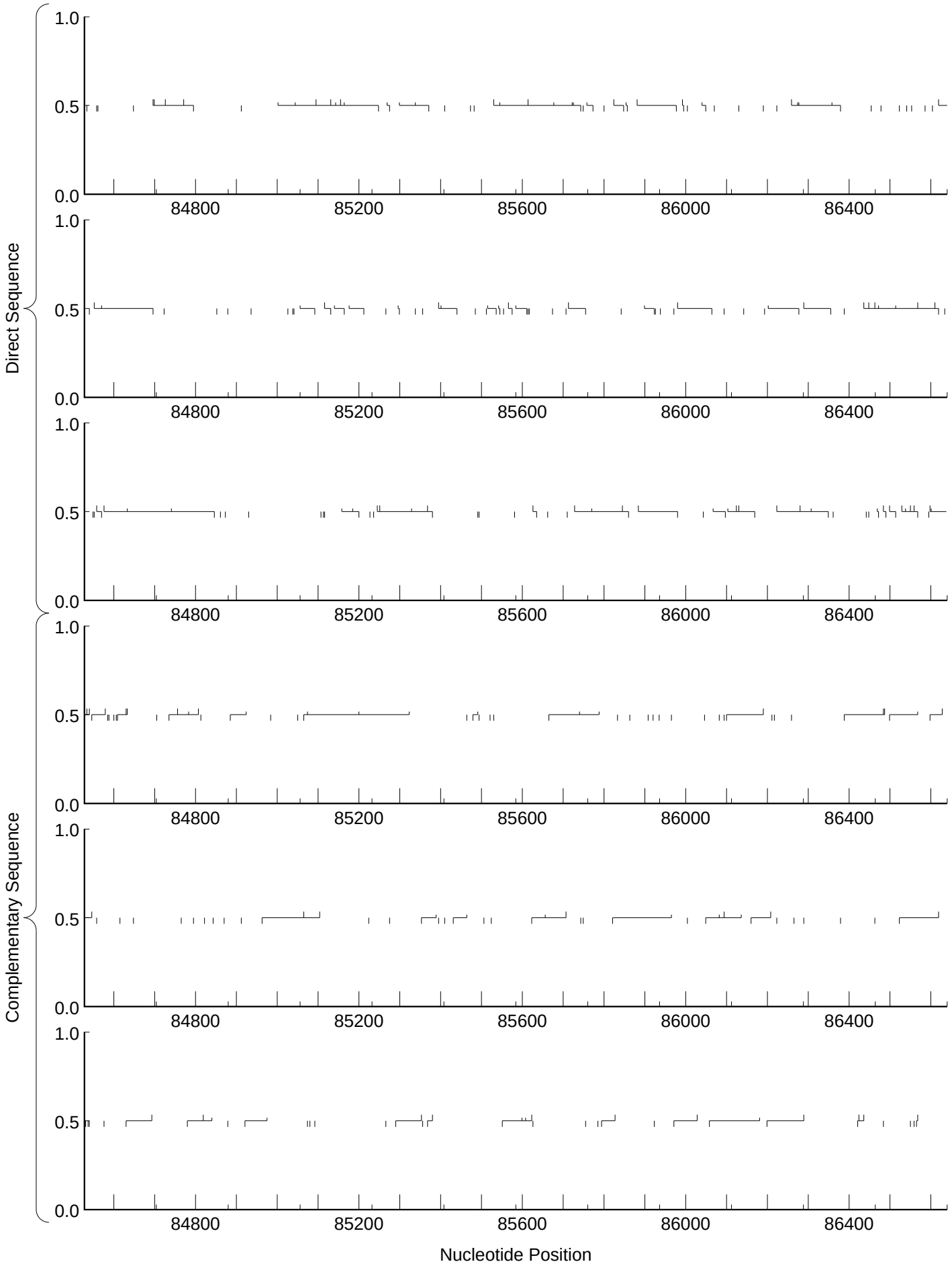
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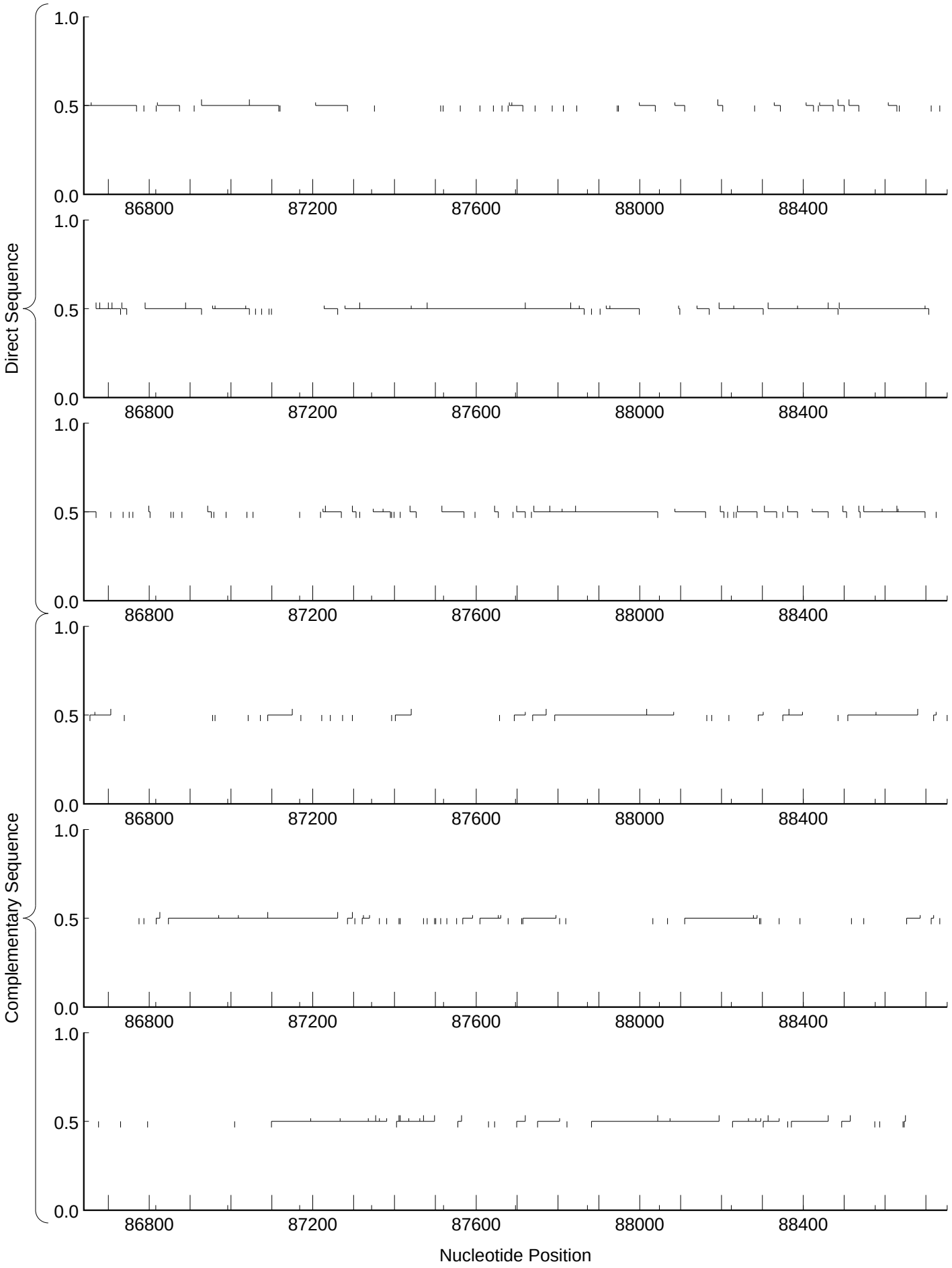
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 41/63



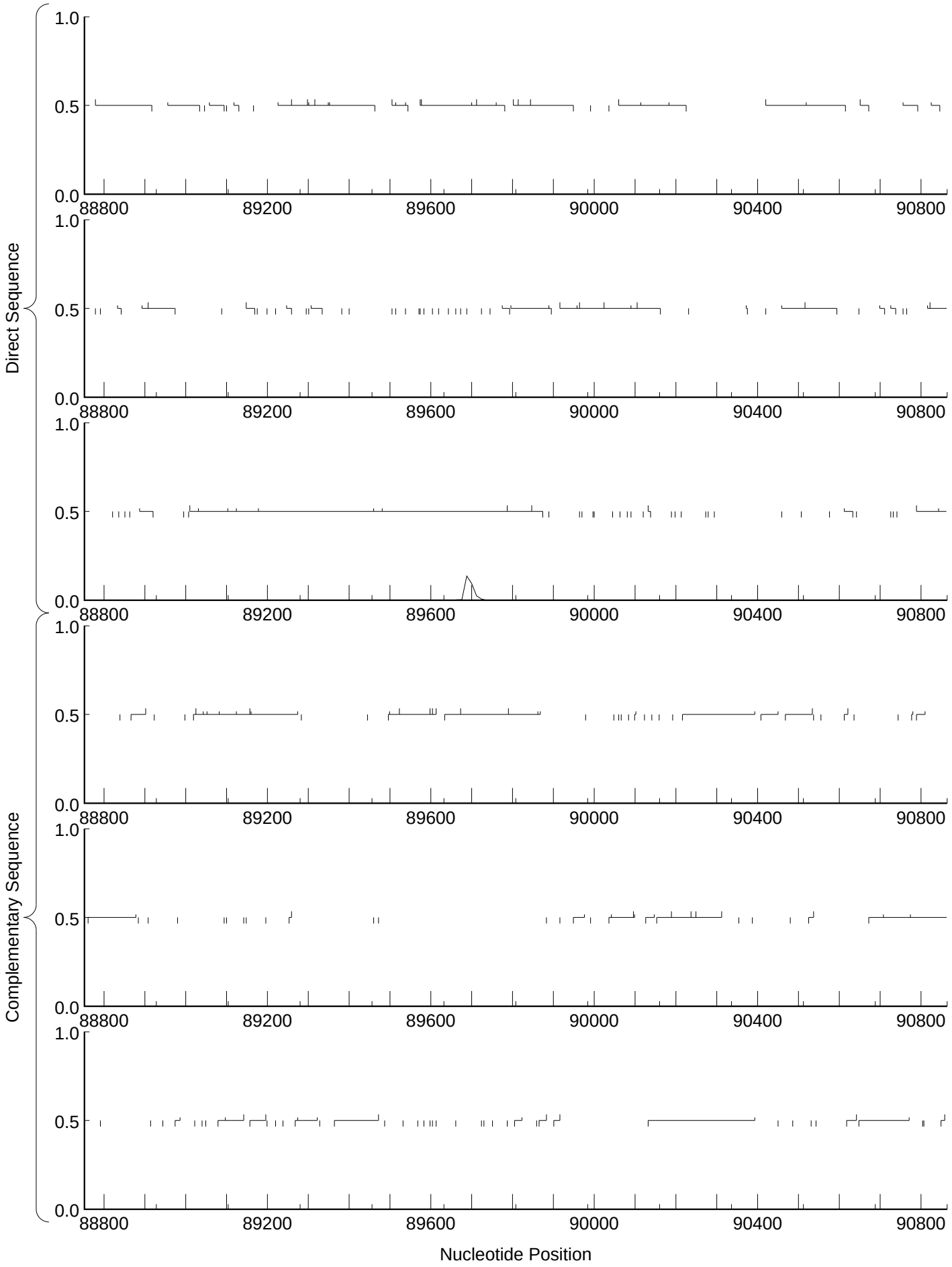
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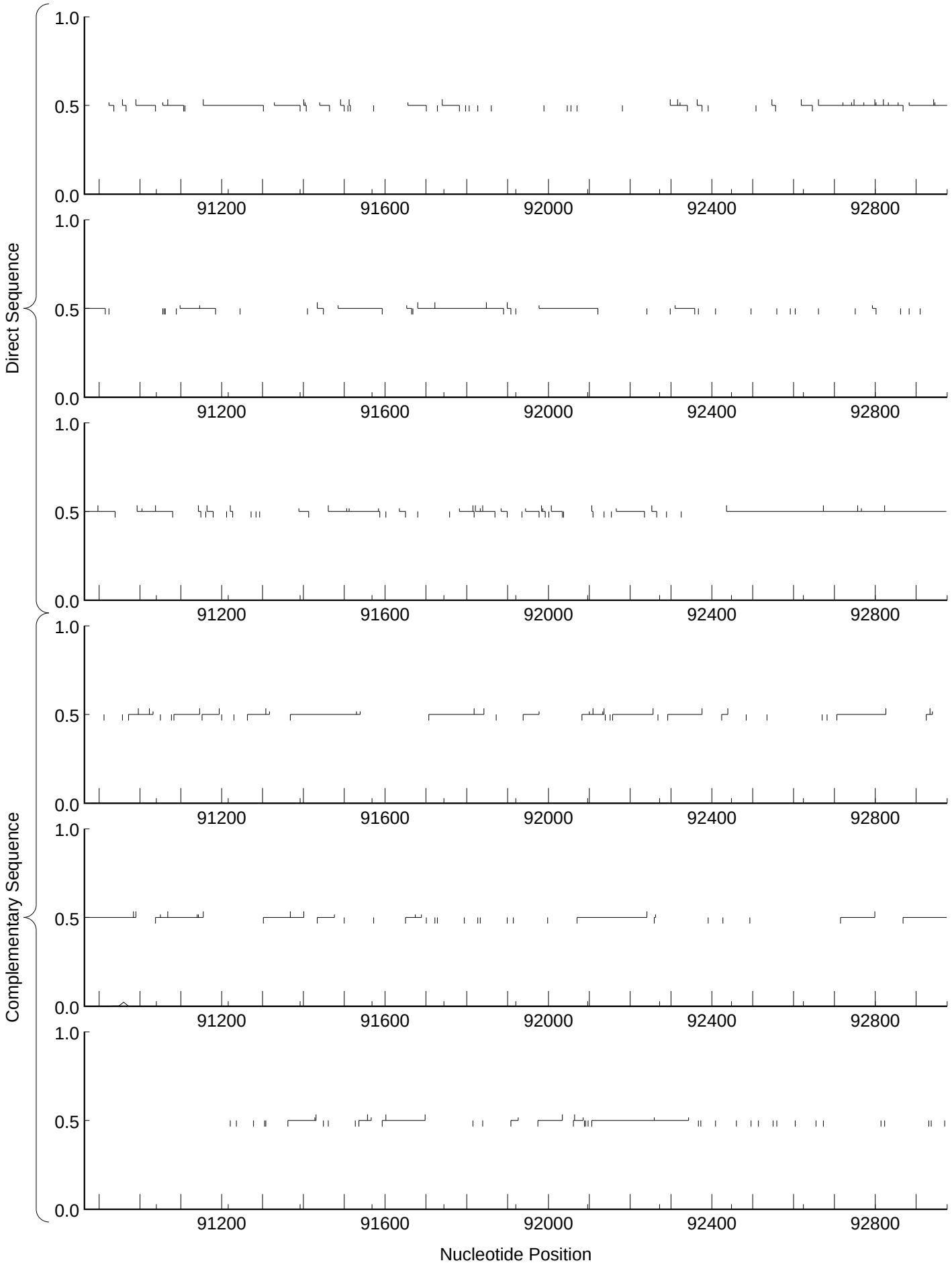
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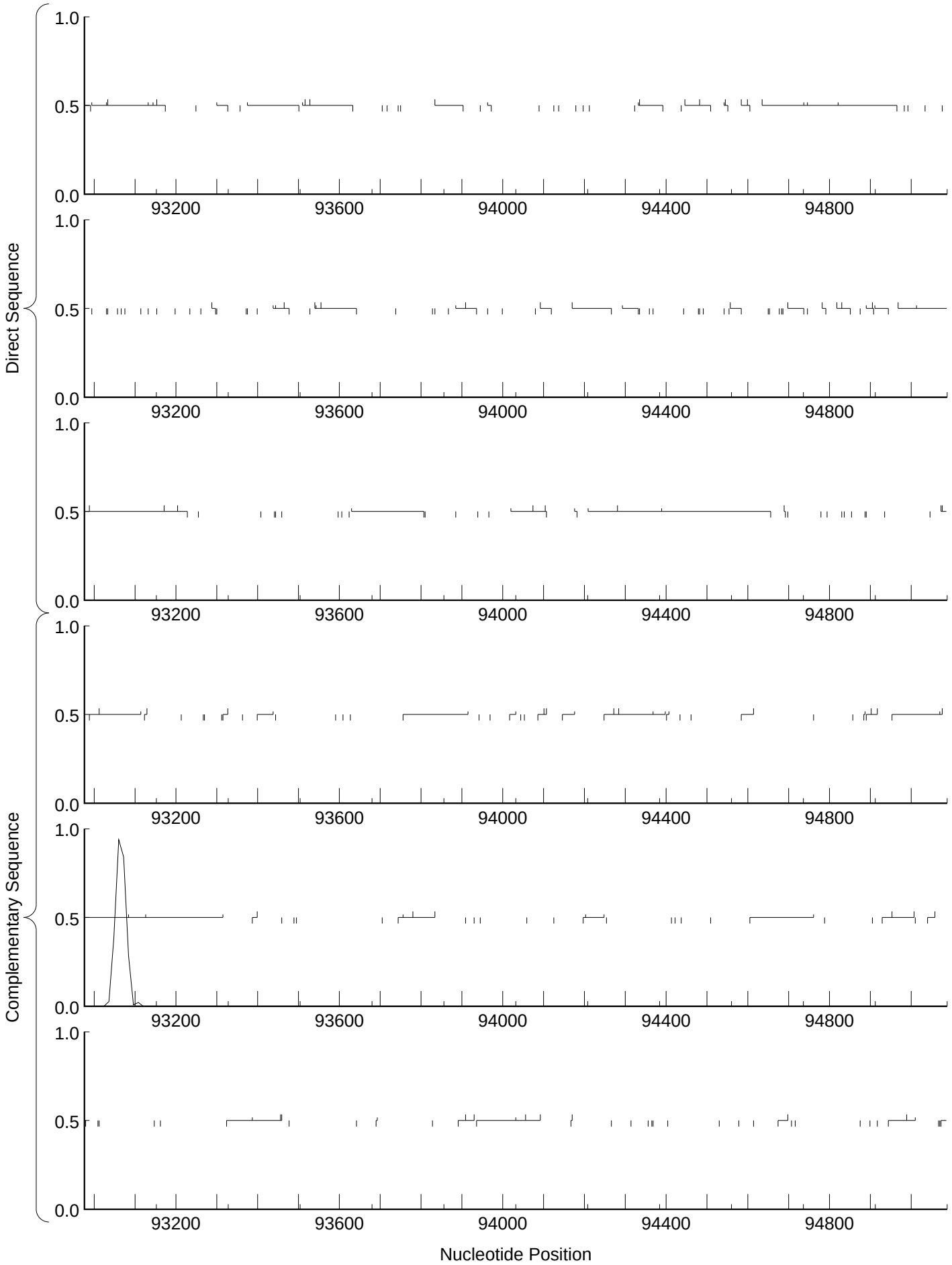
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 44/63



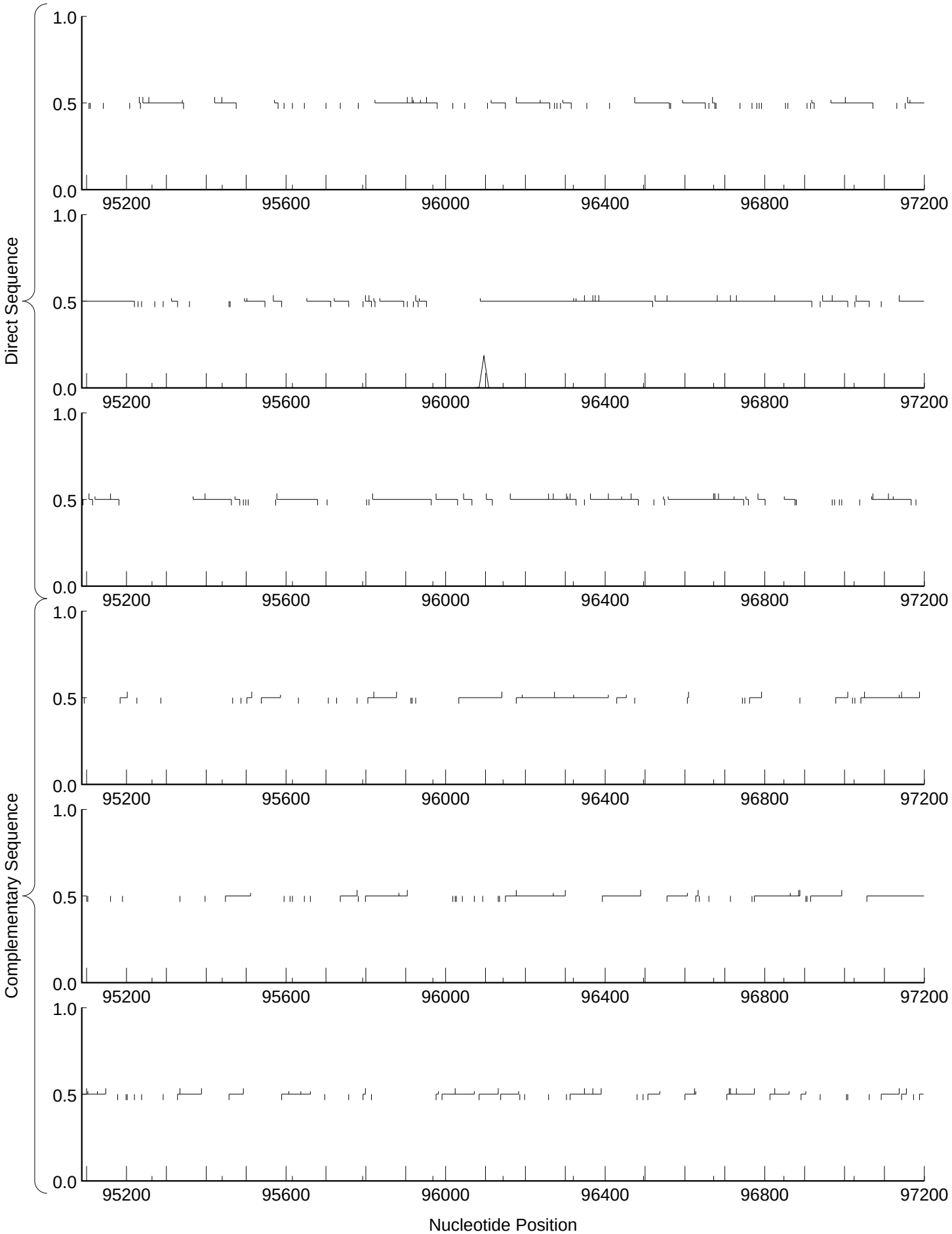
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 45/63



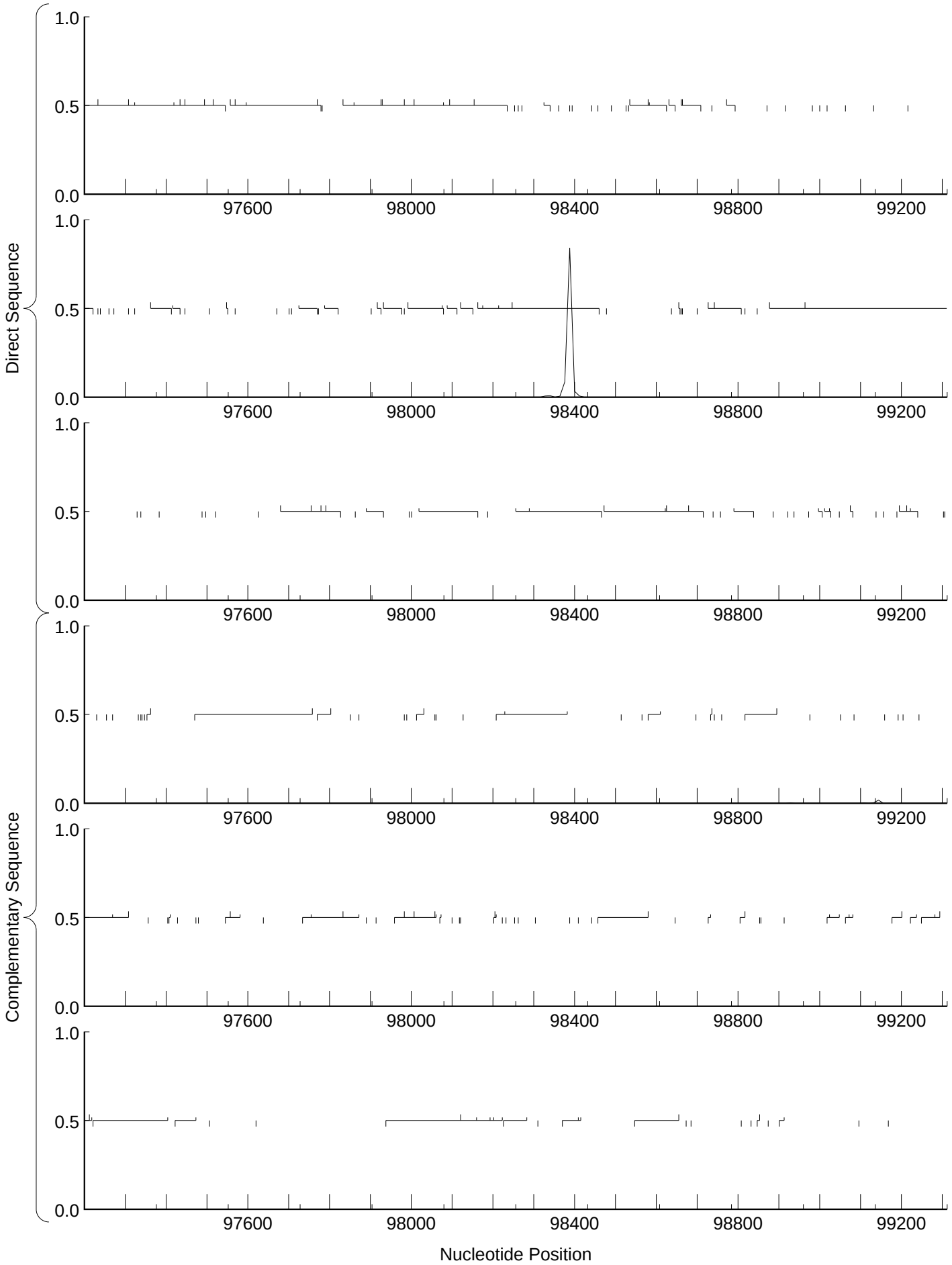
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 46/63



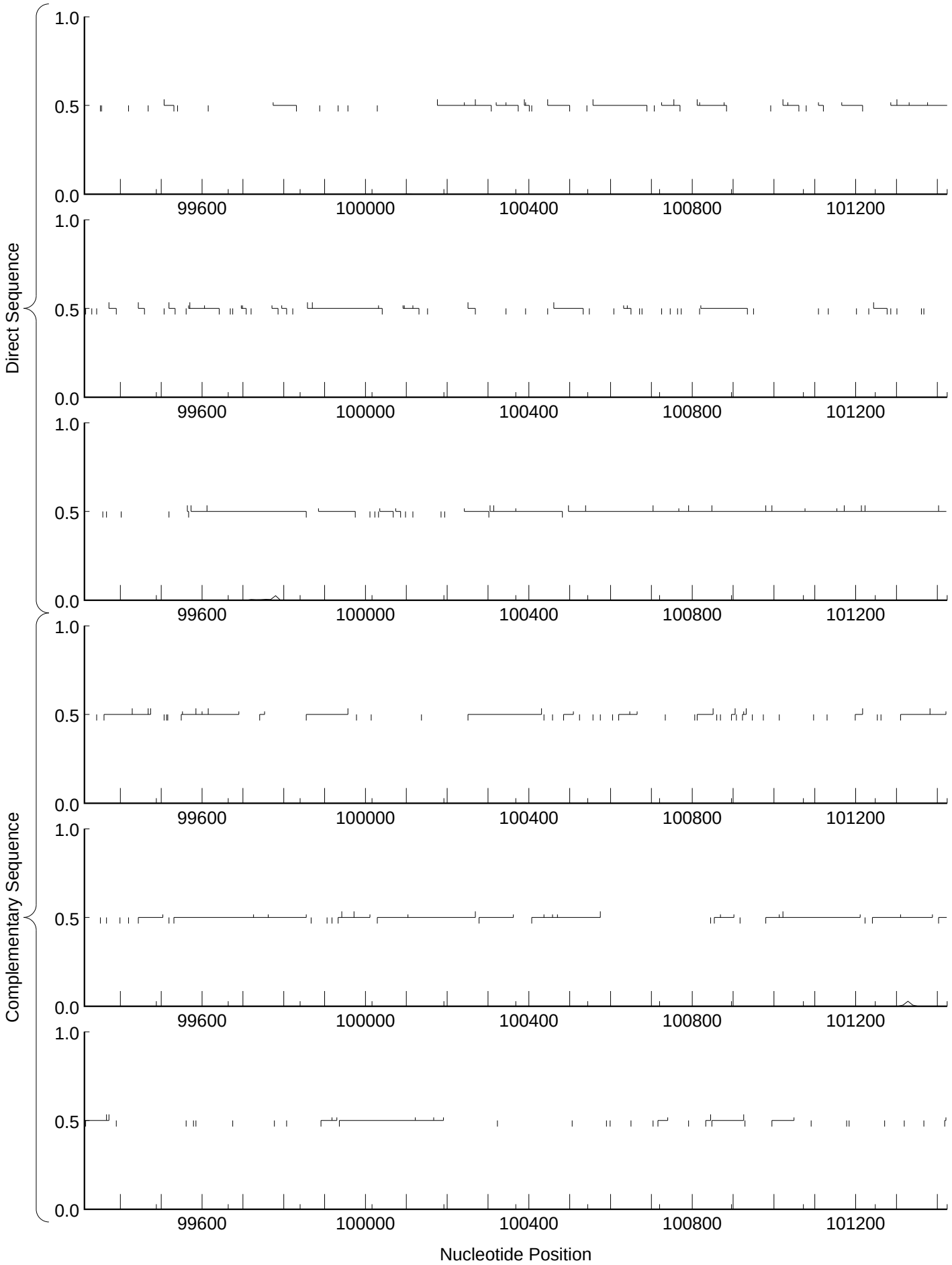
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 47/63



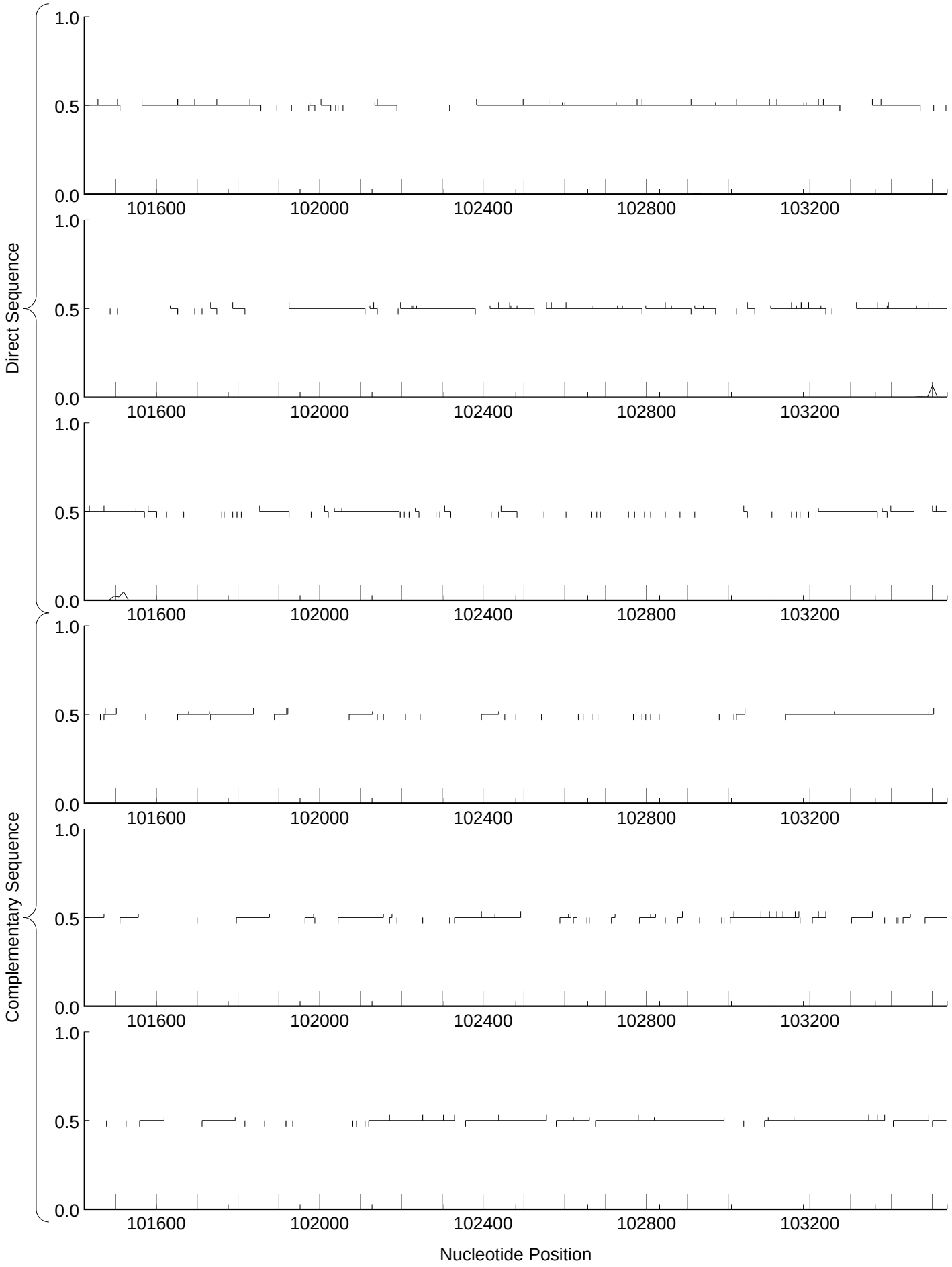
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 48/63



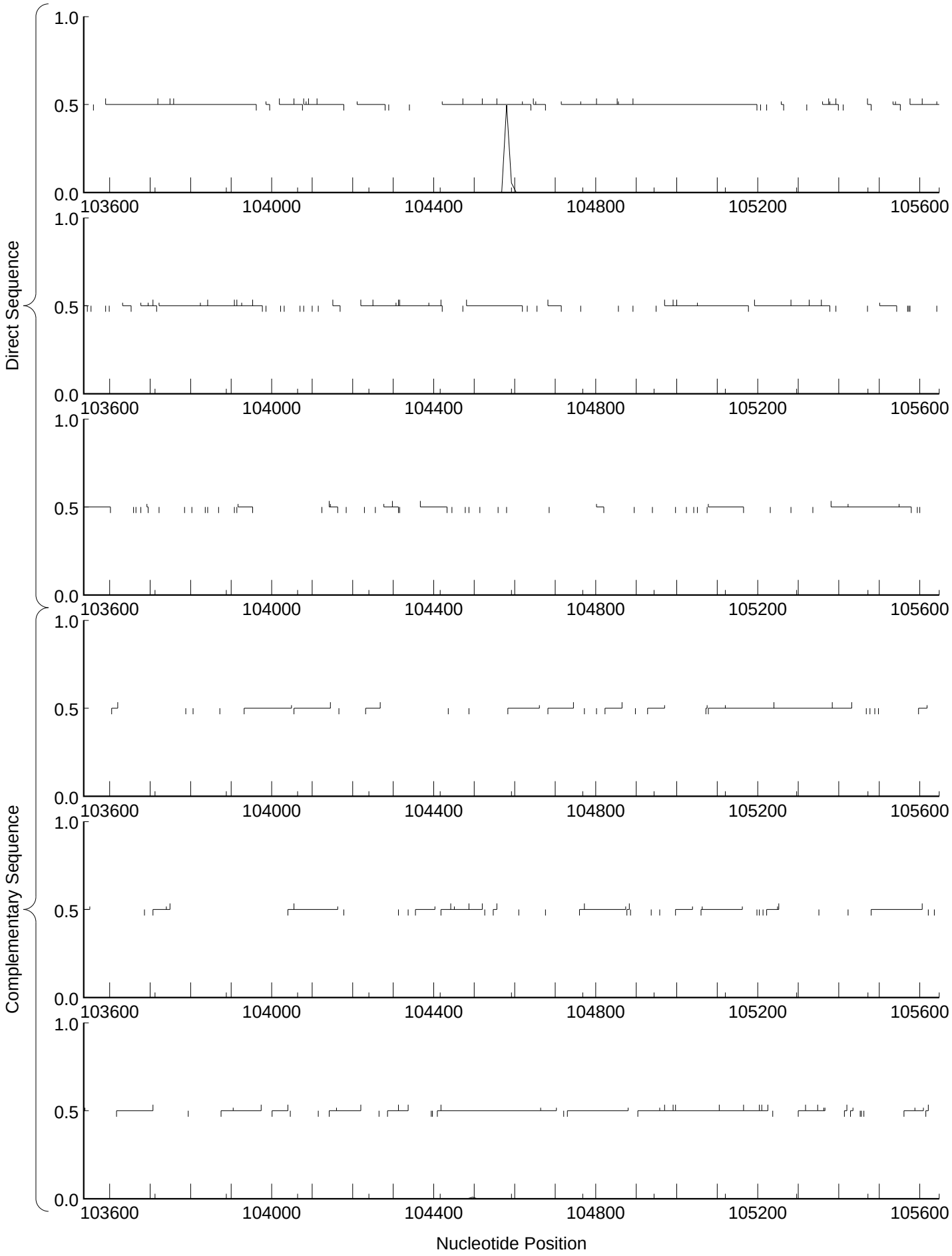
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 49/63



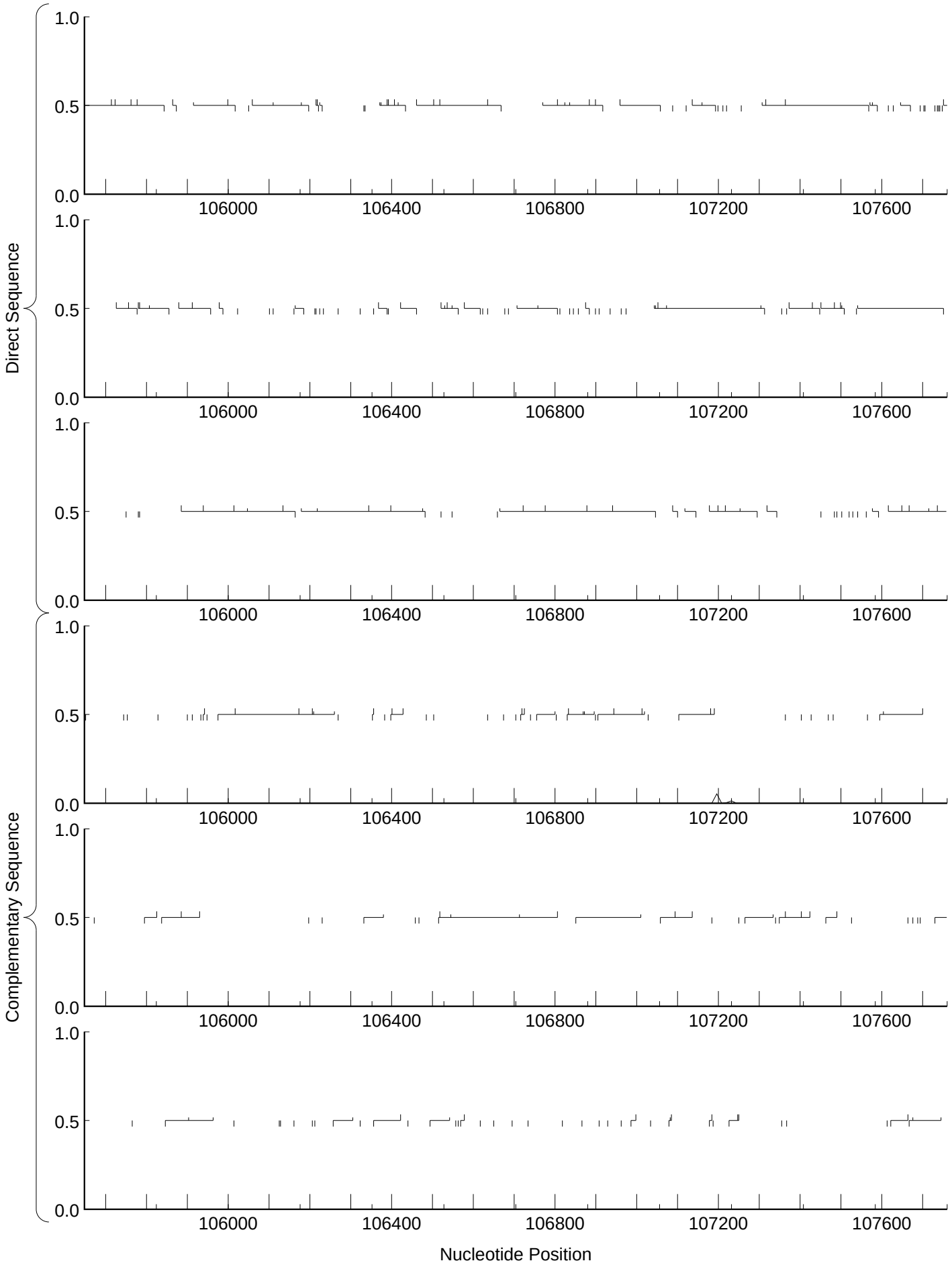
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 50/63



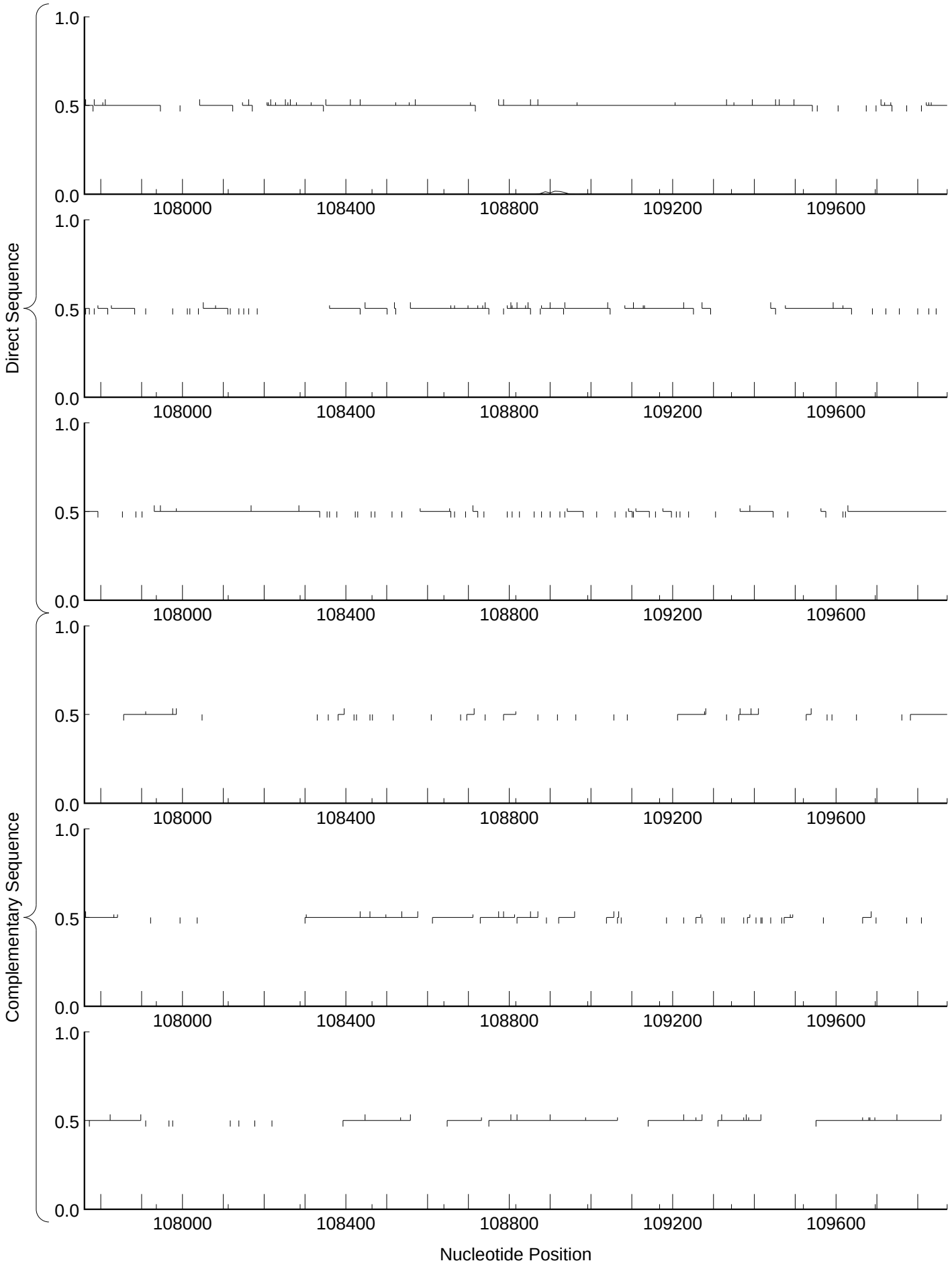
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 51/63



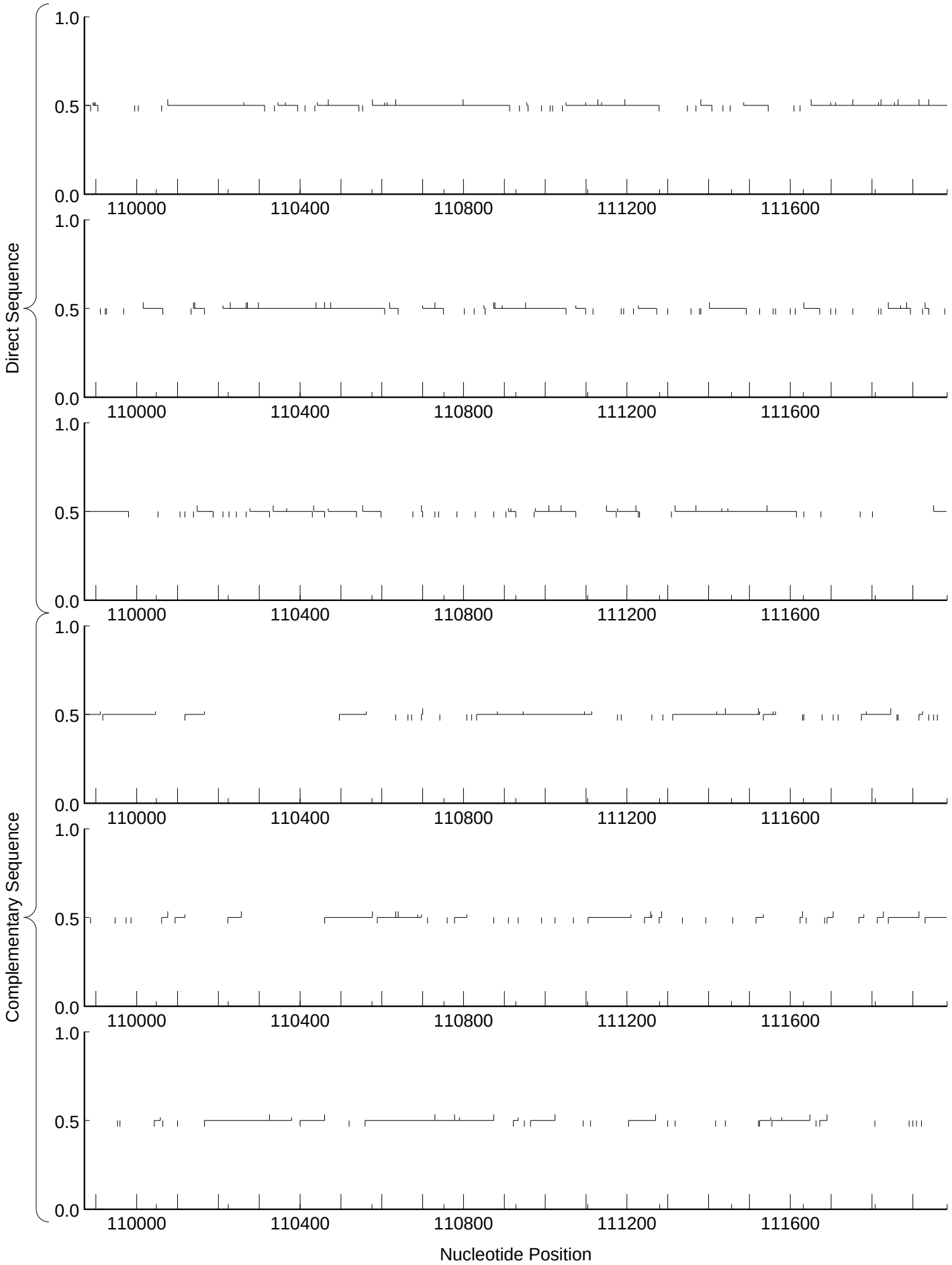
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 52/63



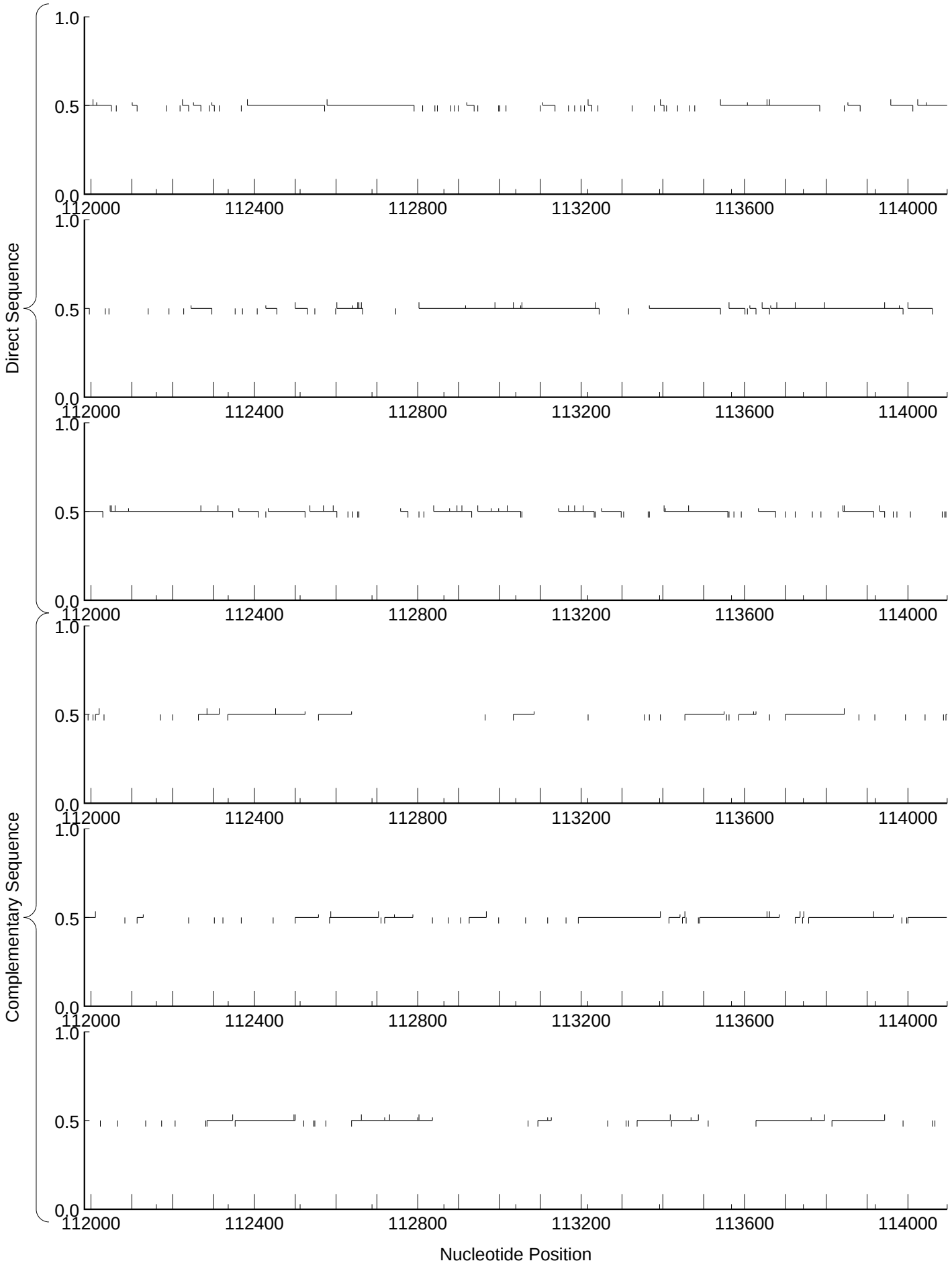
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 53/63



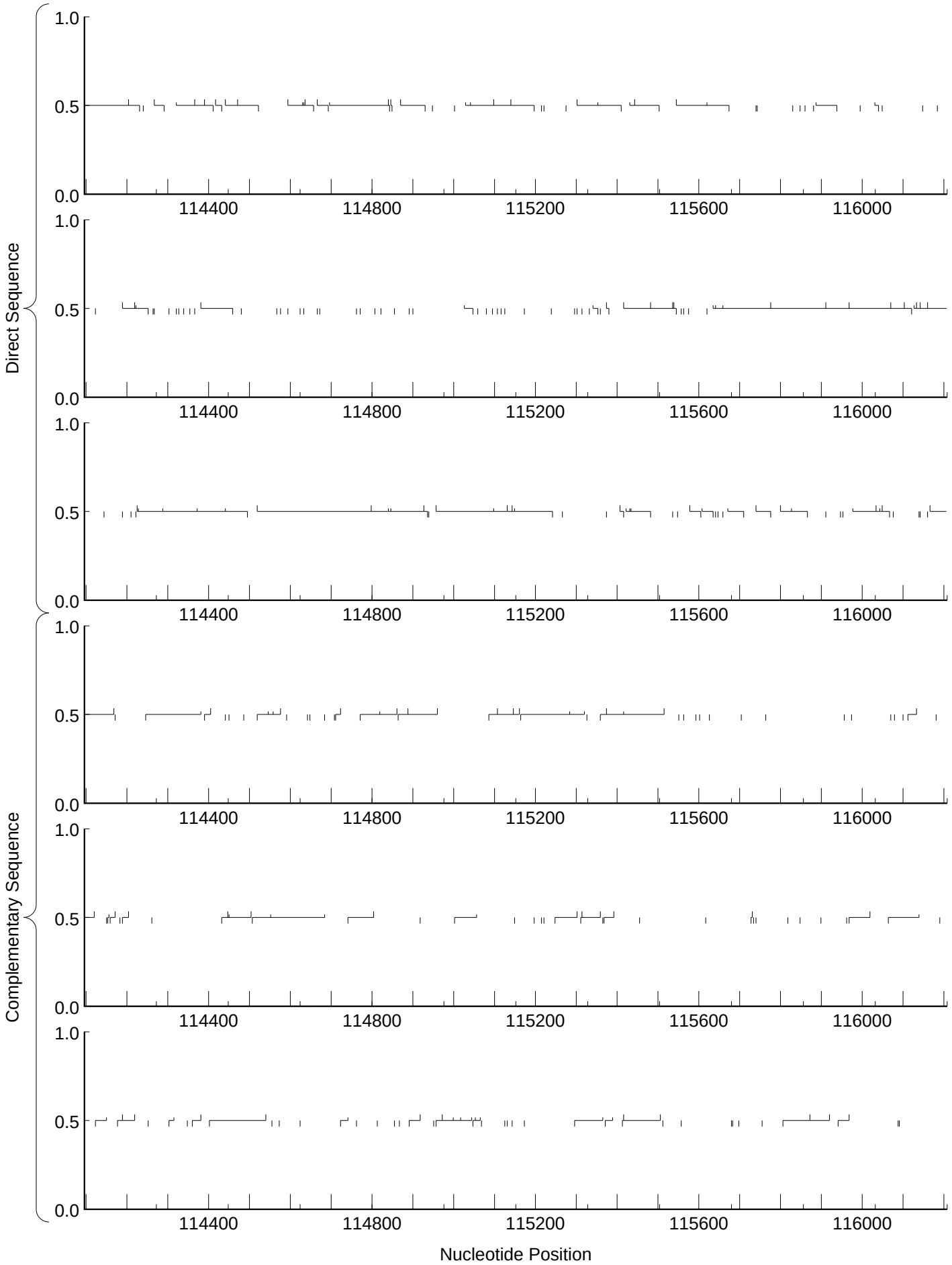
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 54/63



MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 55/63

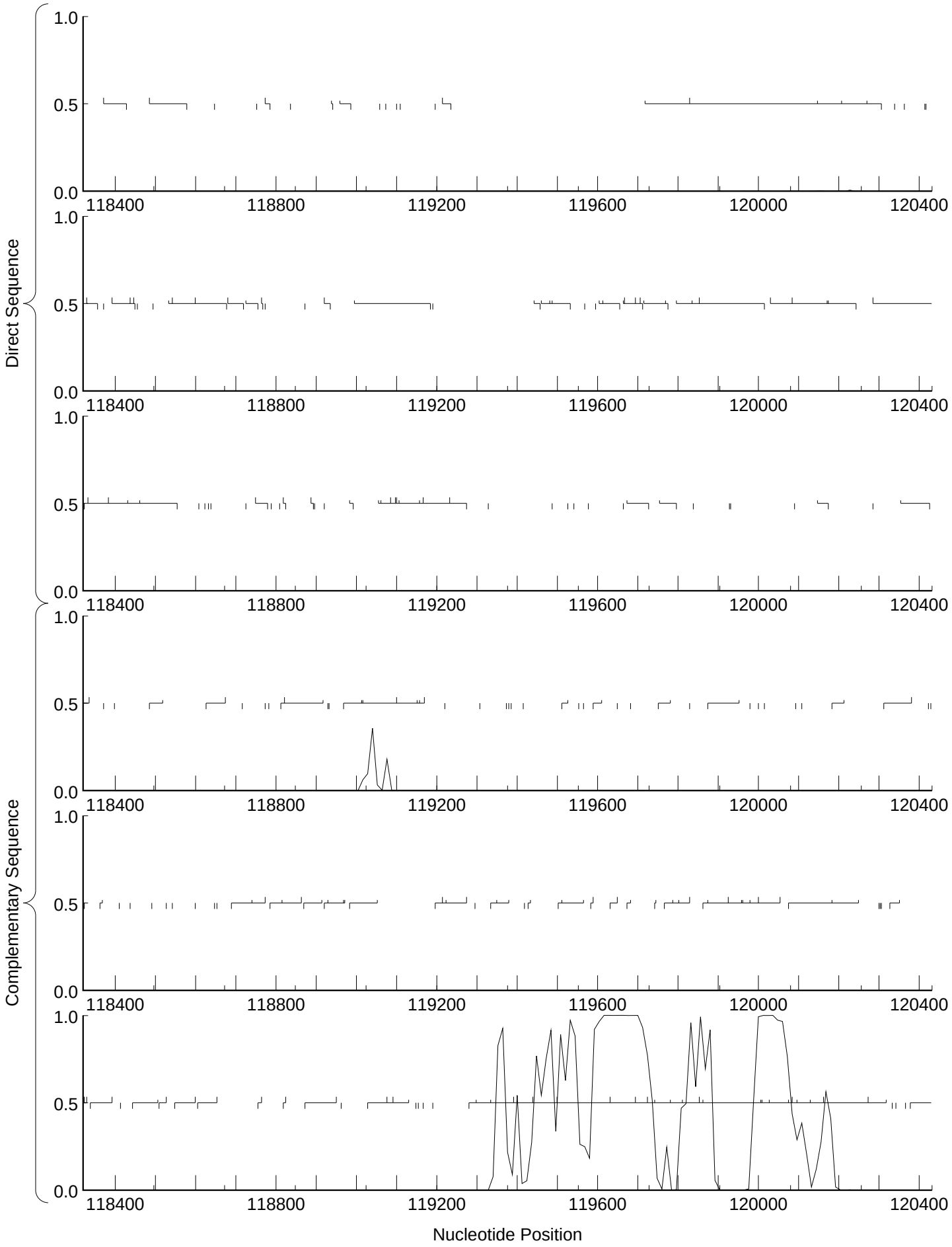


MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 56/63

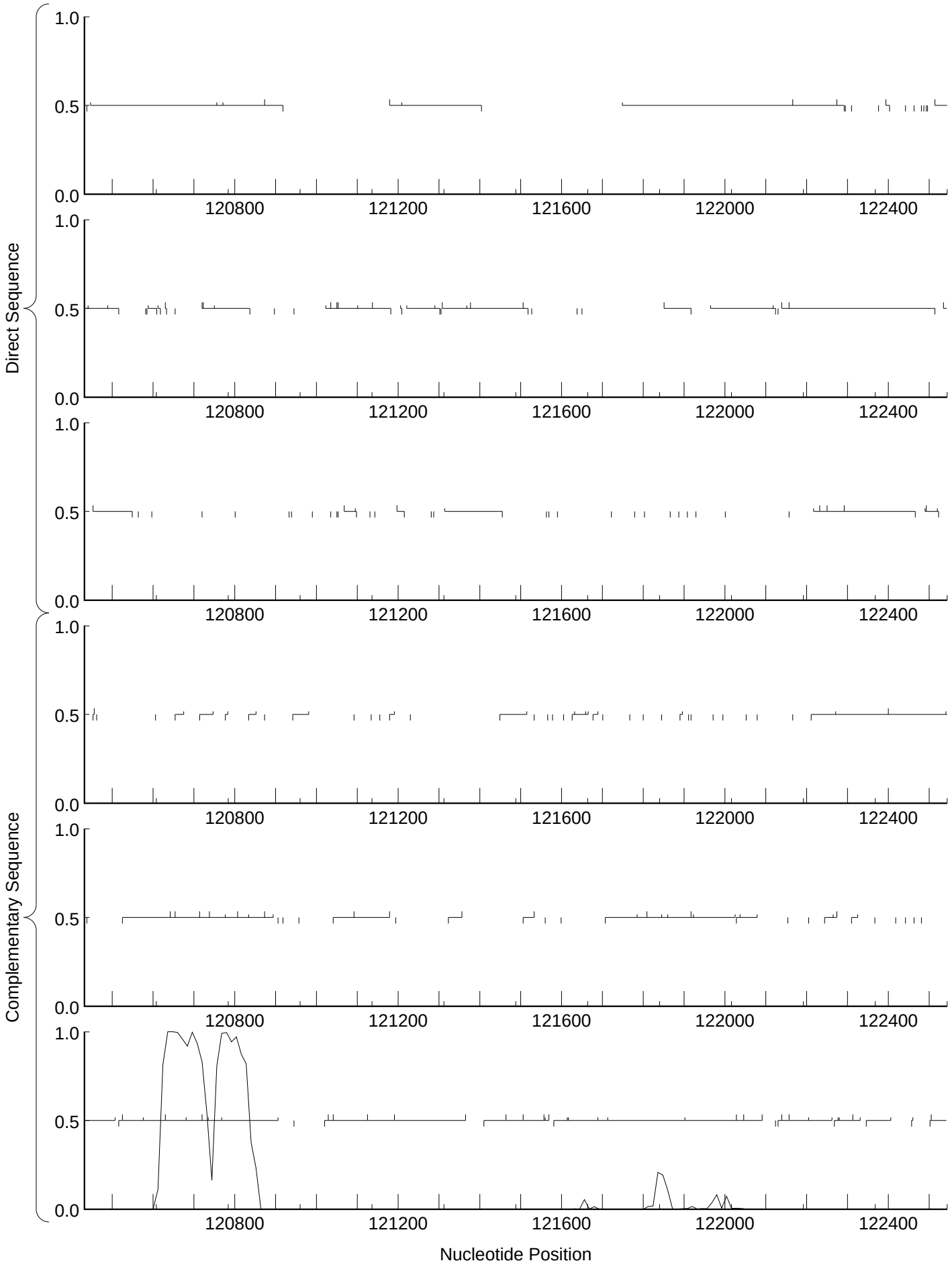


MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 57/63
Direct Sequence
Complementary Sequence
1.0
0.5
0.0
116400
116800
117200
117600
118000
Nucleotide Position

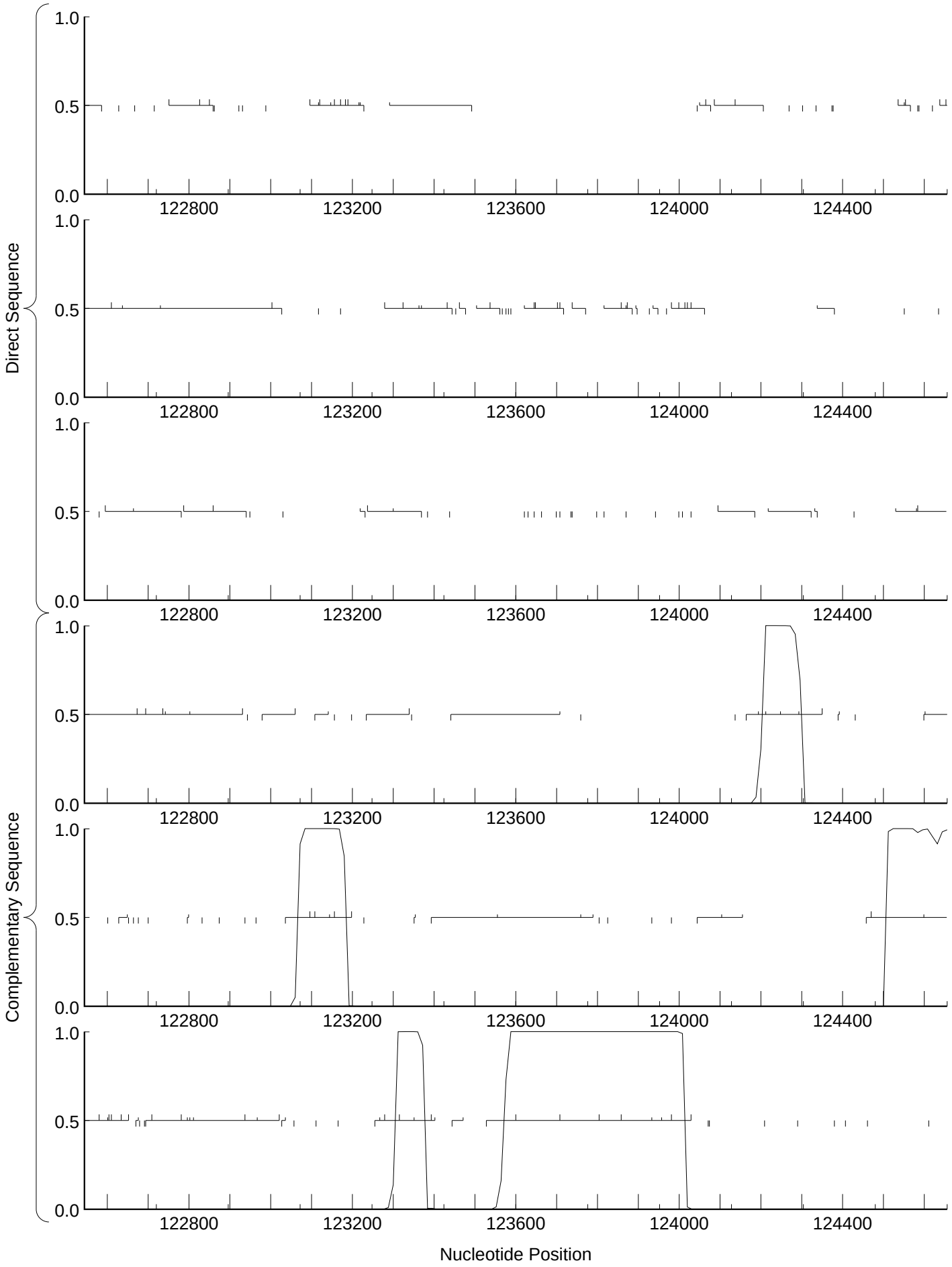
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 58/63



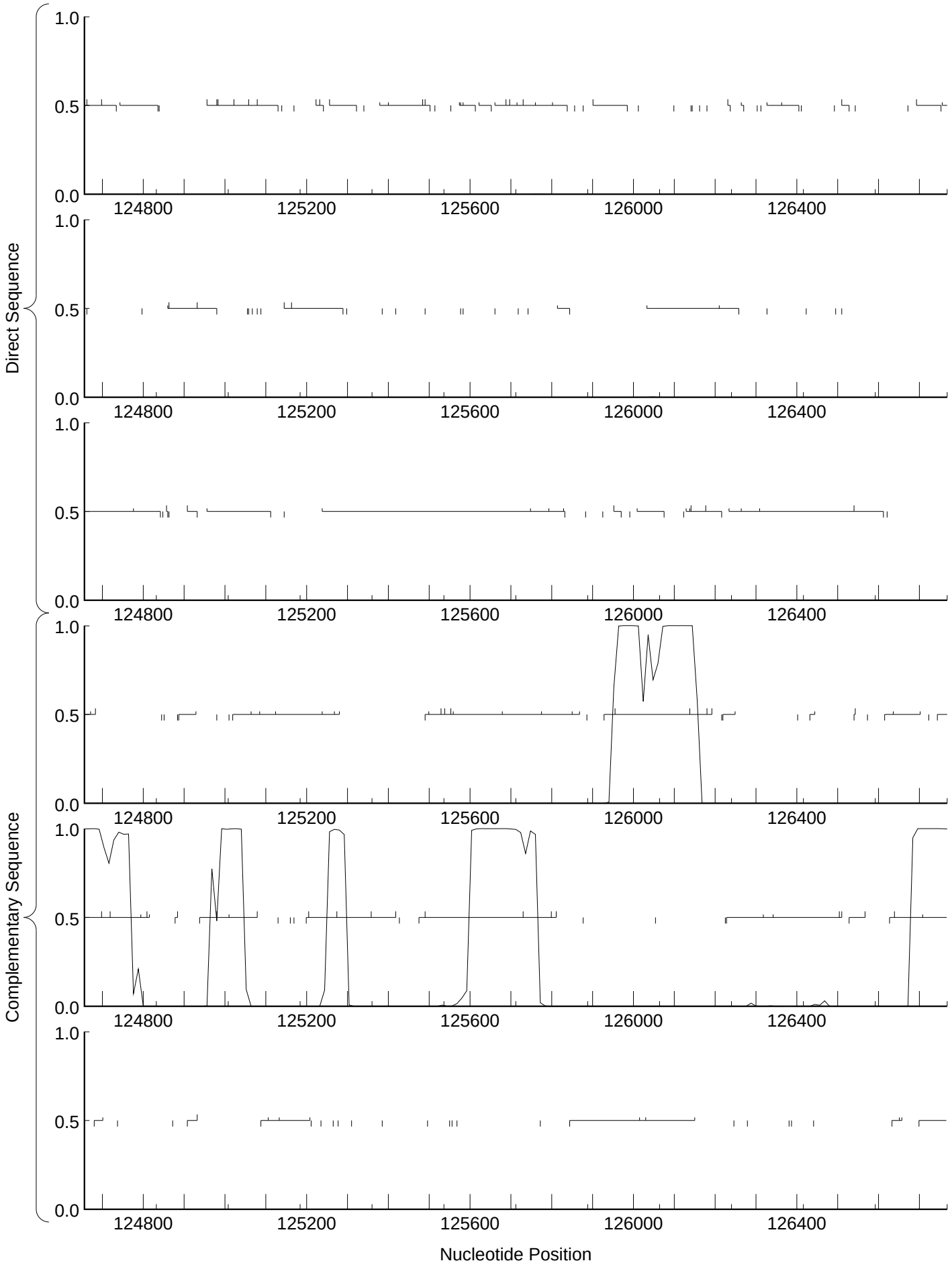
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 59/63



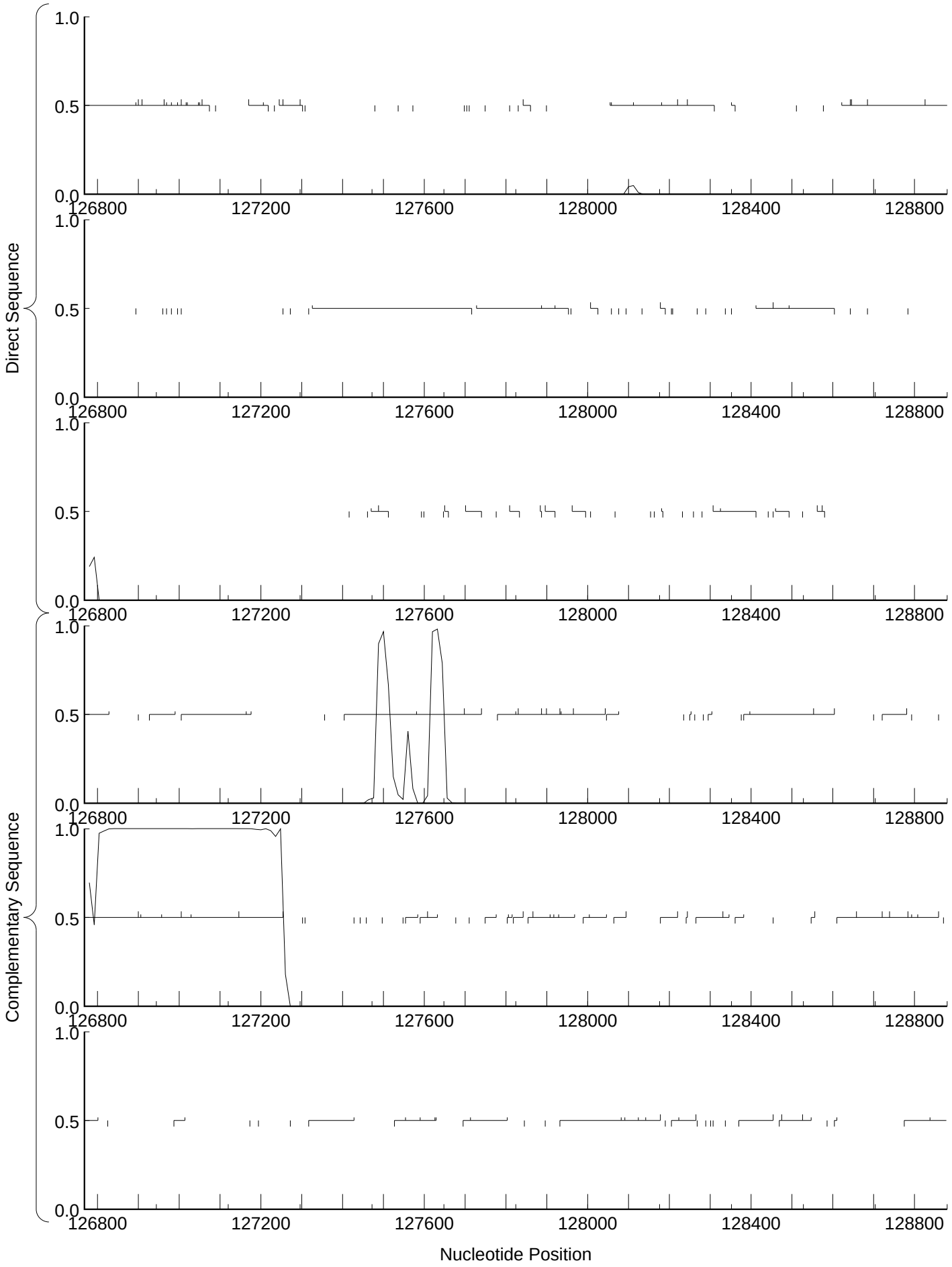
MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 60/63



MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 61/63



MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 62/63



MindFlayer complete sequence, 130258 bp including 12198 bp terminal repeat, Cluster BE2, Order 4, Window 96, Step 12, 63/63

