GeneMark

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

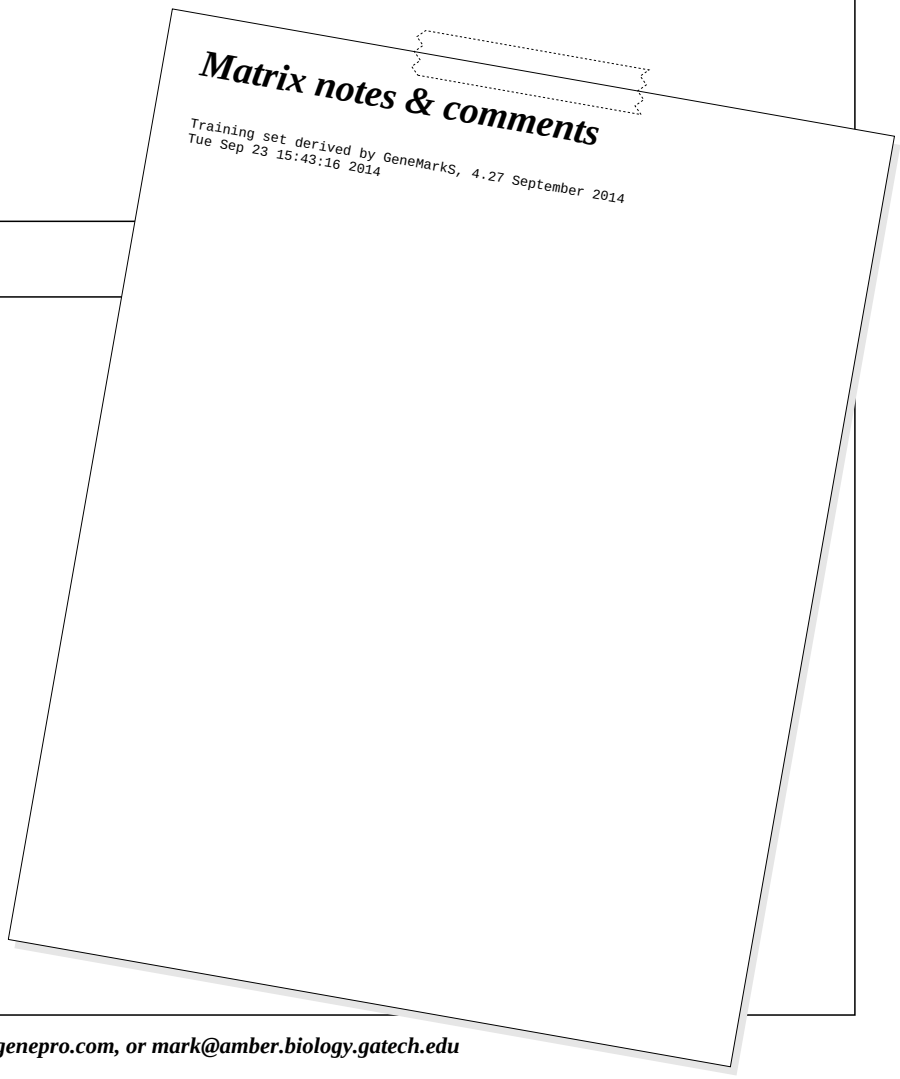
PROGRAM INFORMATION

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Analysis Date    :  3/28/24 at 21:54:16
Pages            :  65
Sequence Length  :  134105 bp
GC Content       :   49.42%

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

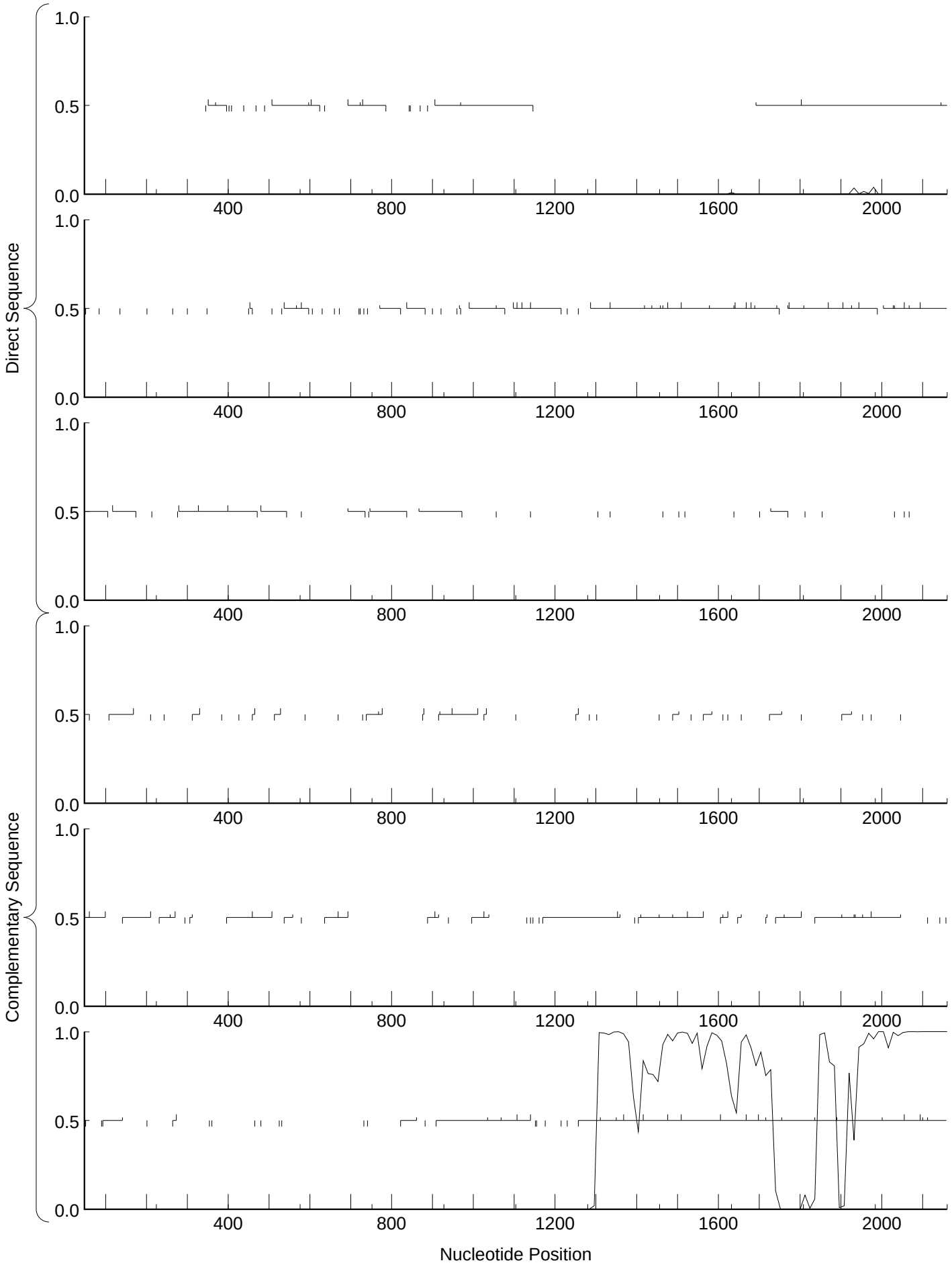
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Author     :  -

Order      :  4
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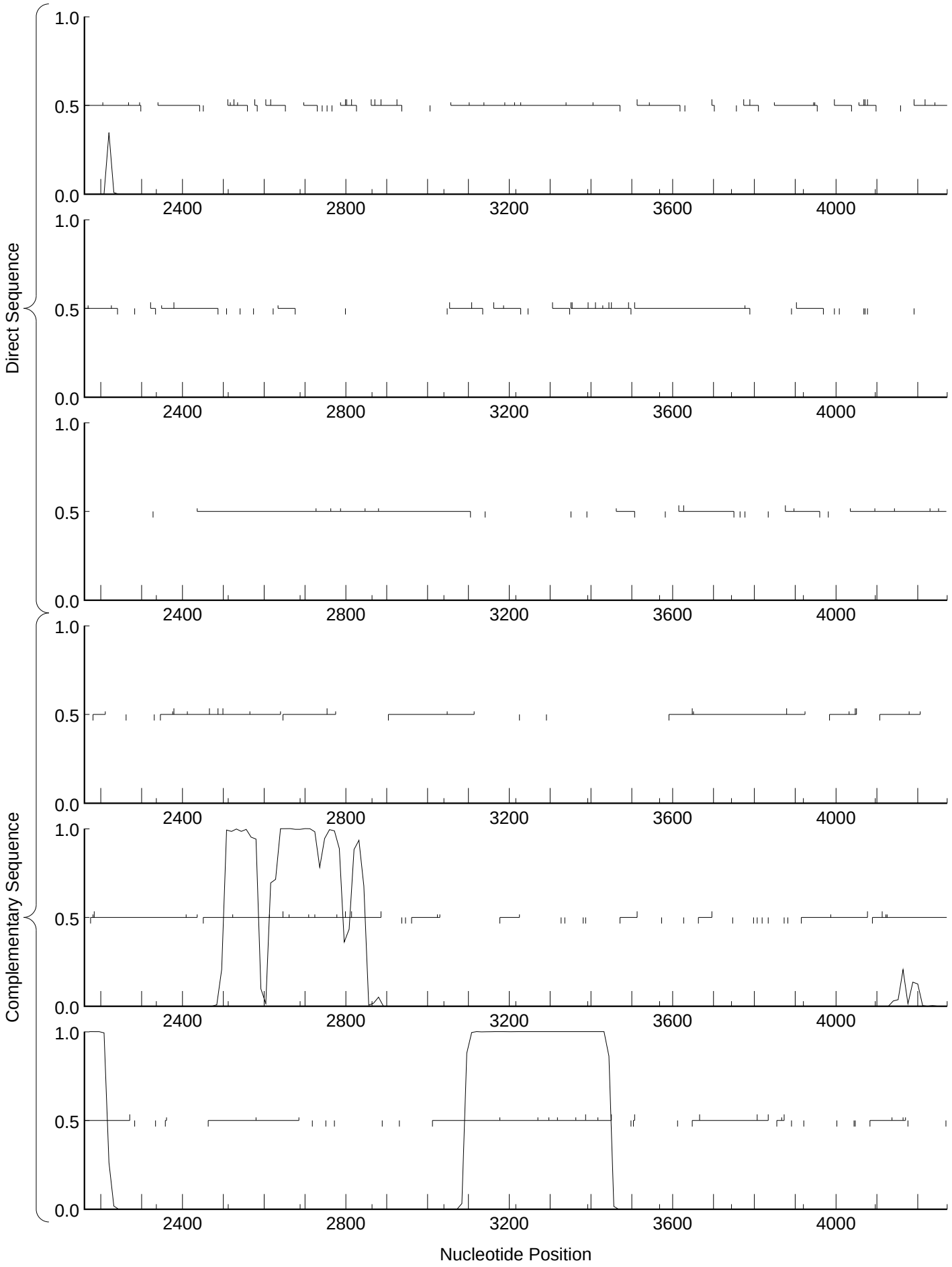

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

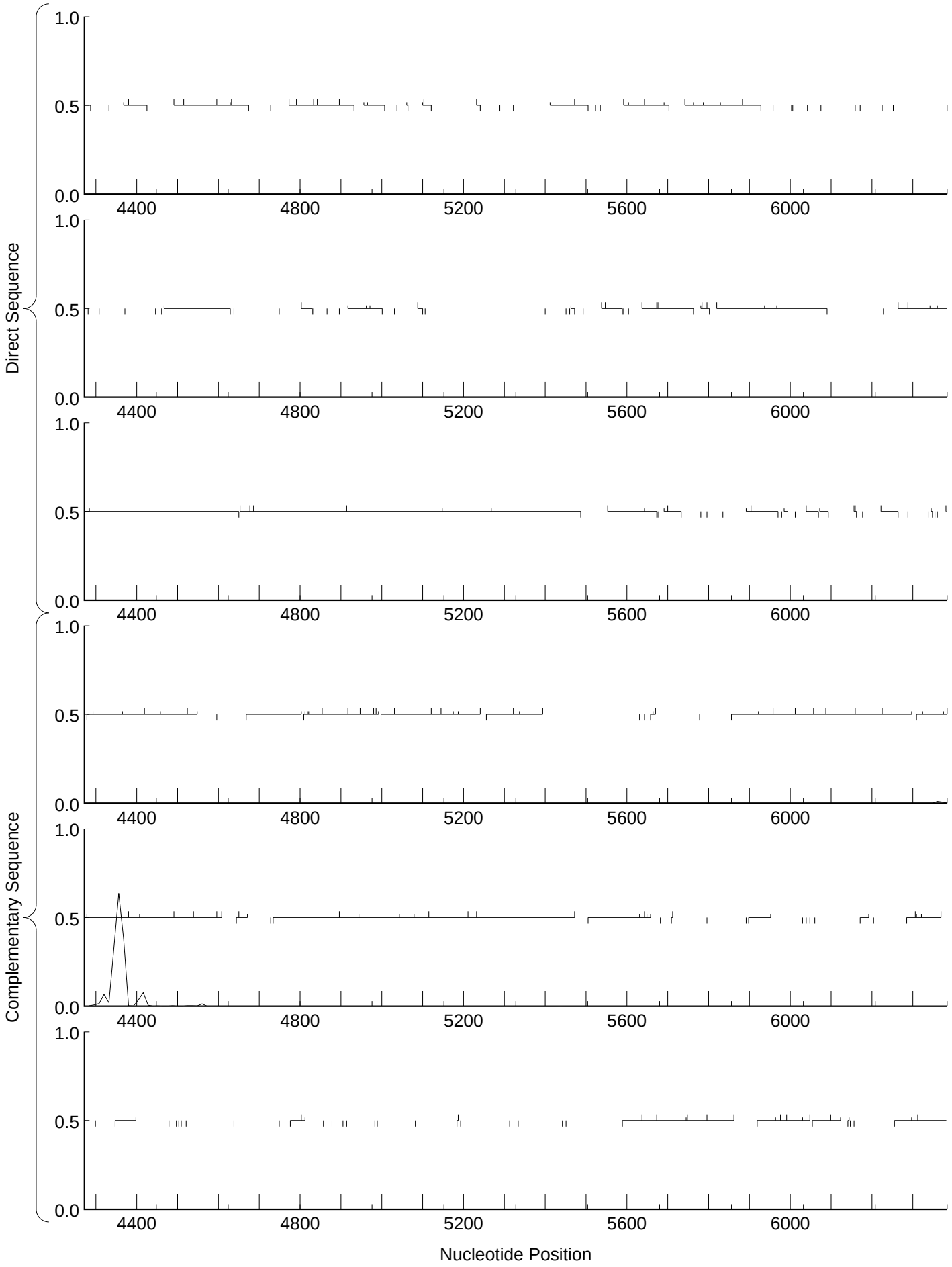
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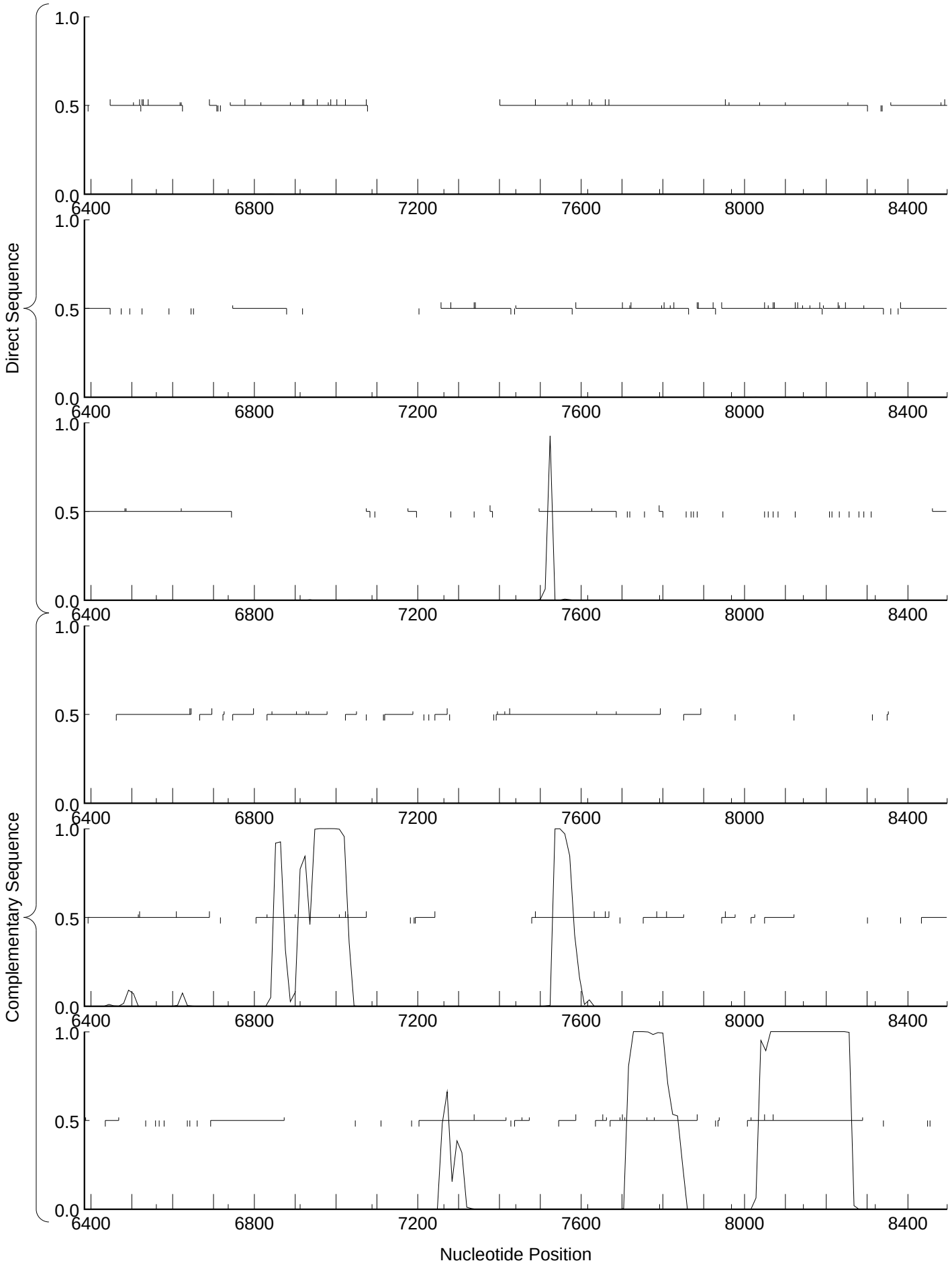
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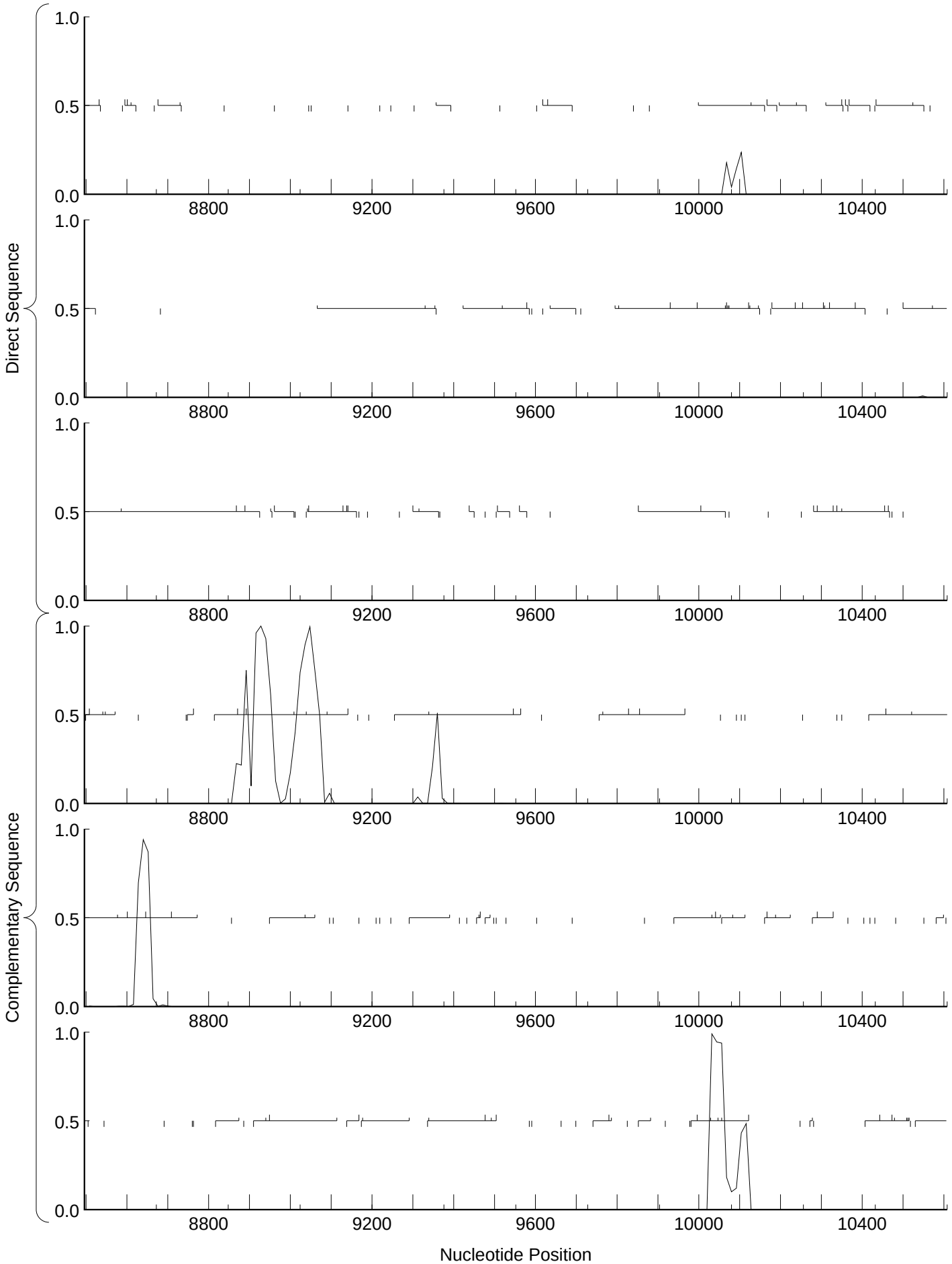
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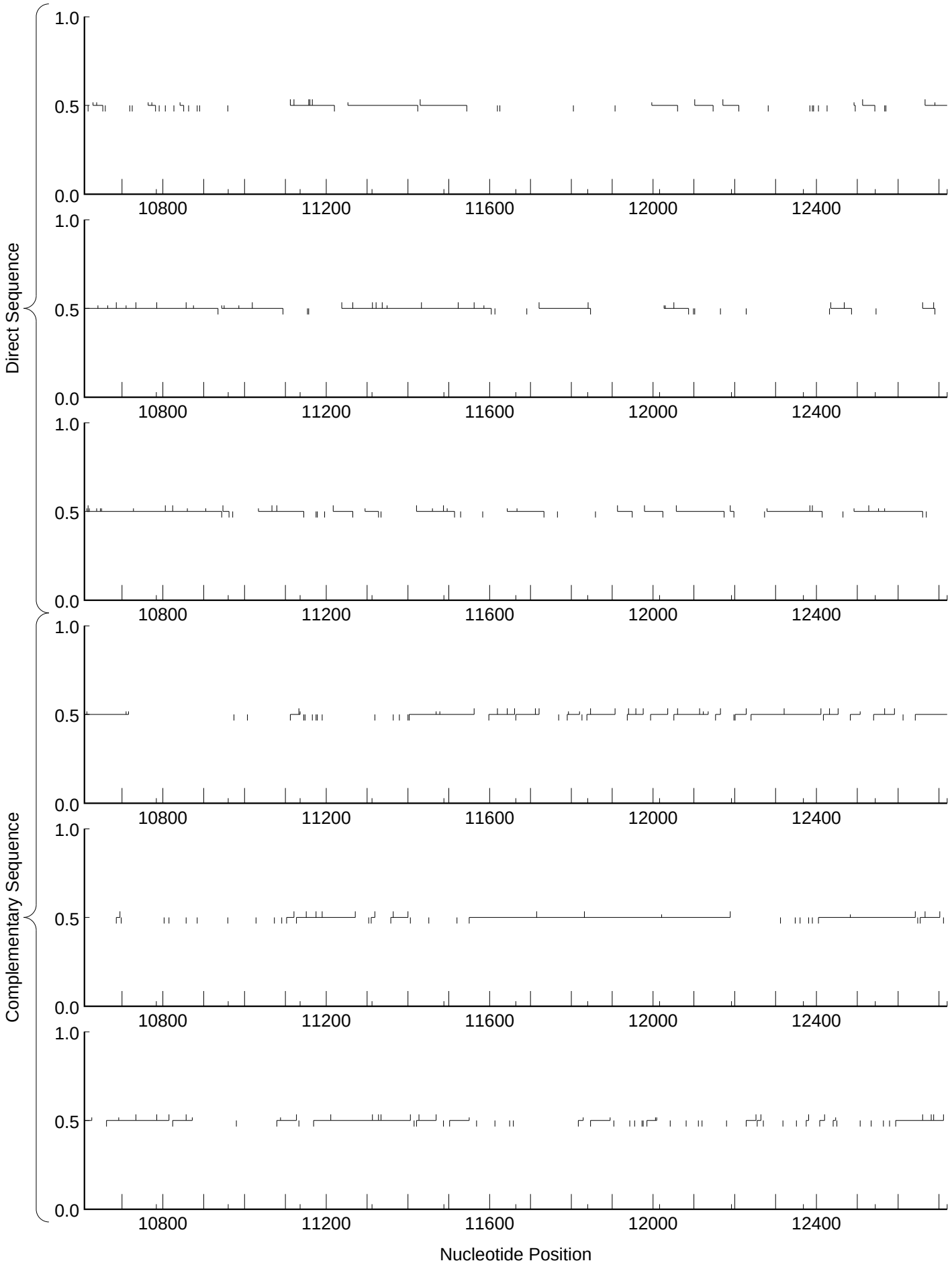
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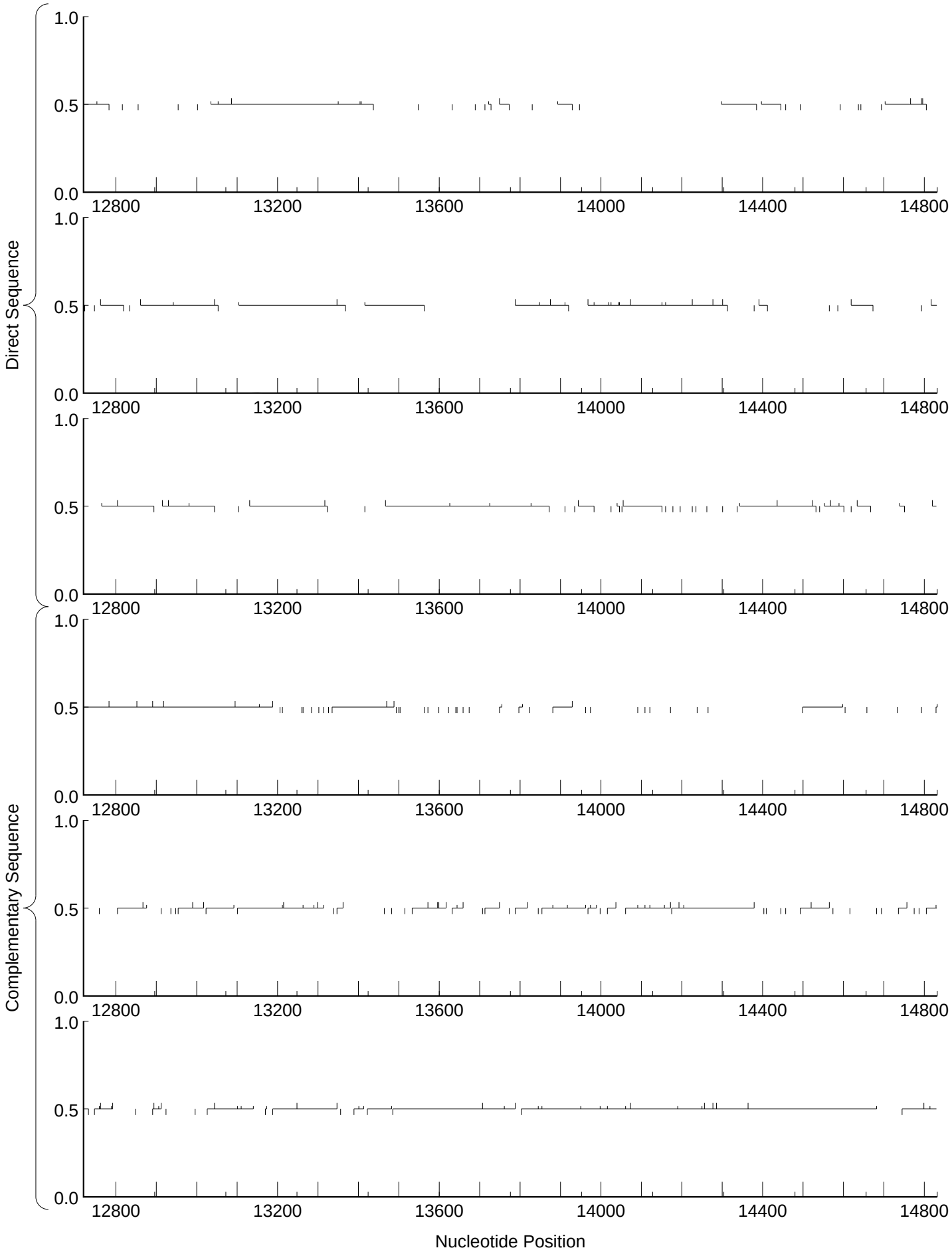
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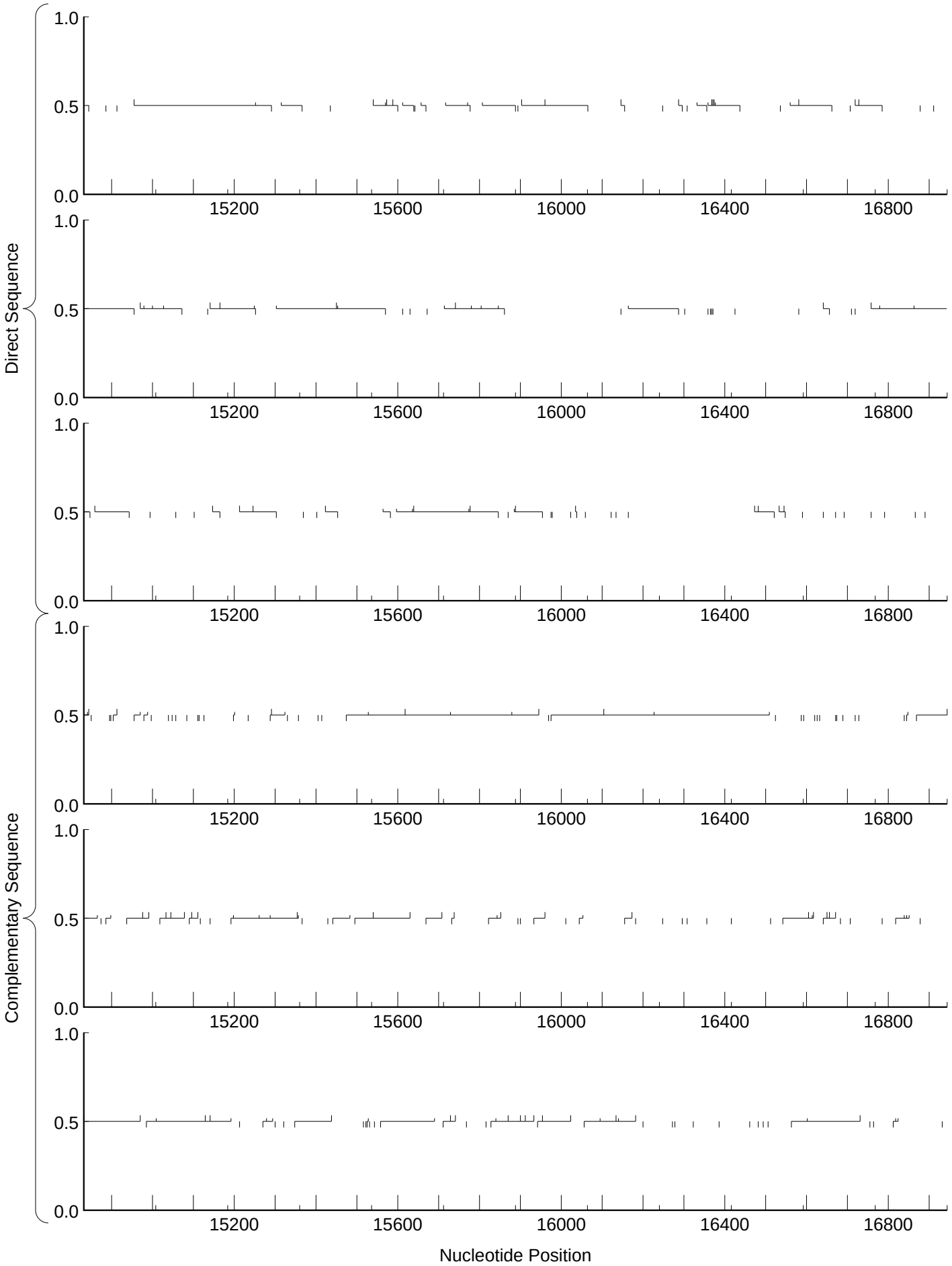
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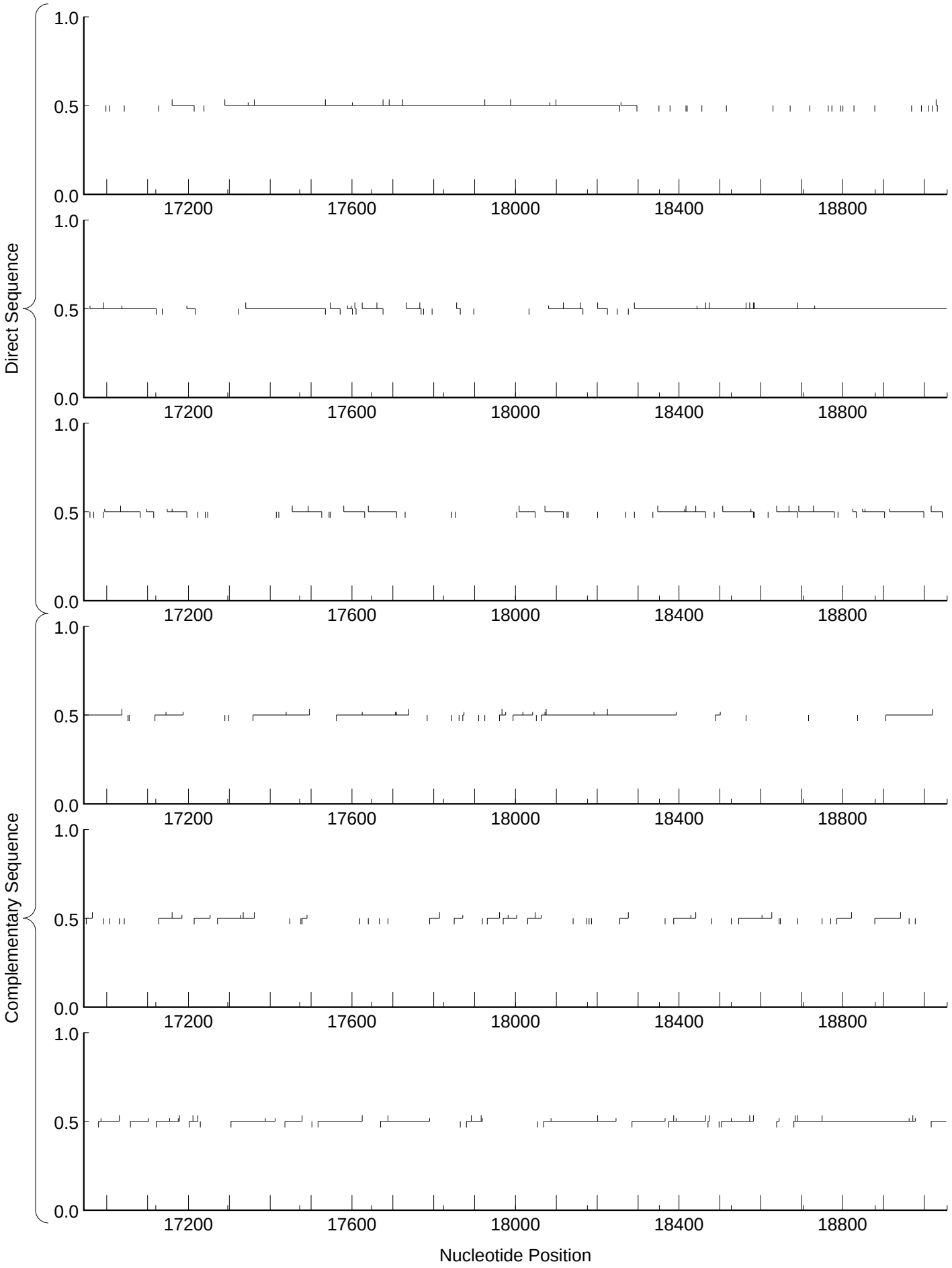
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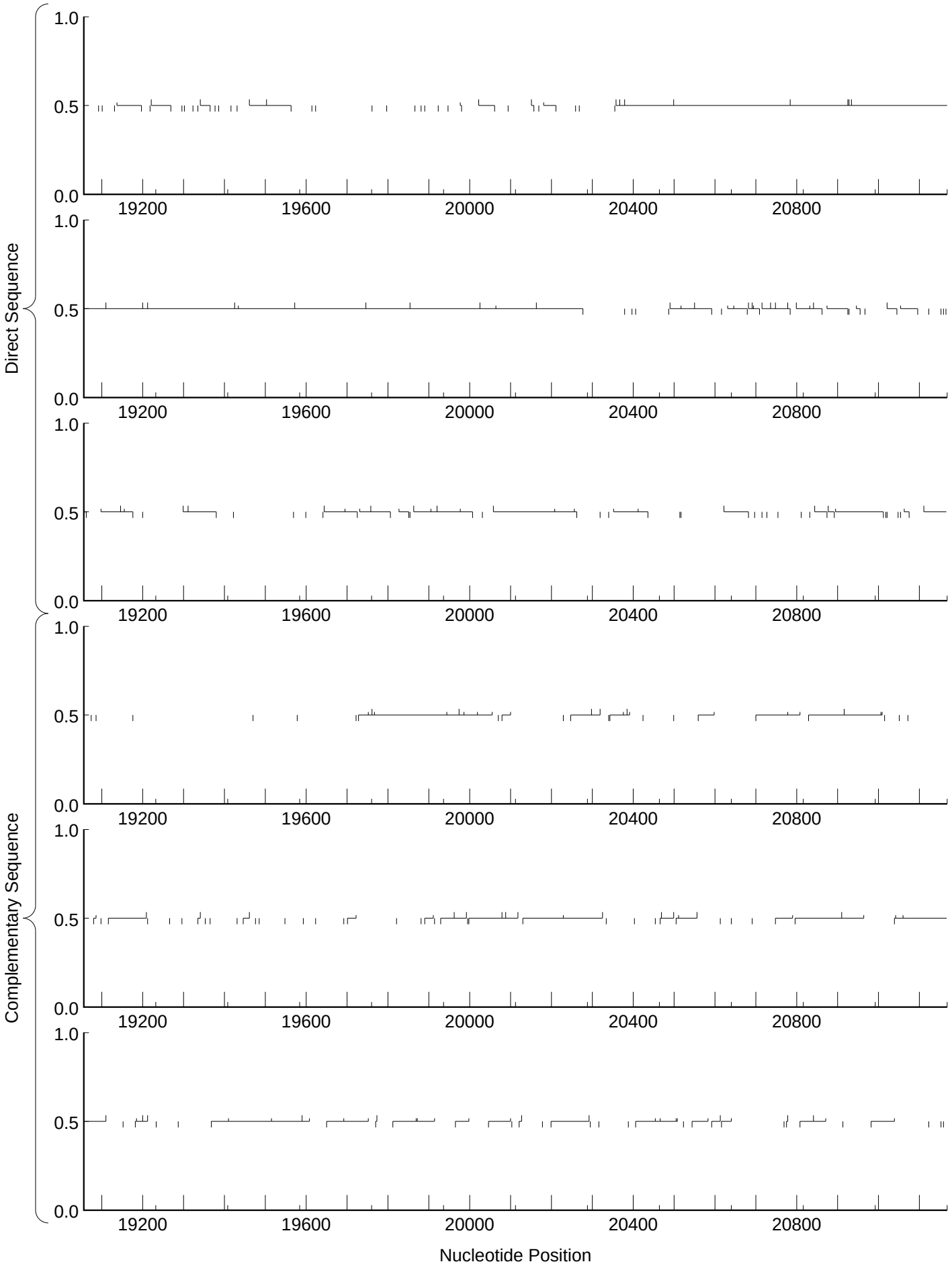
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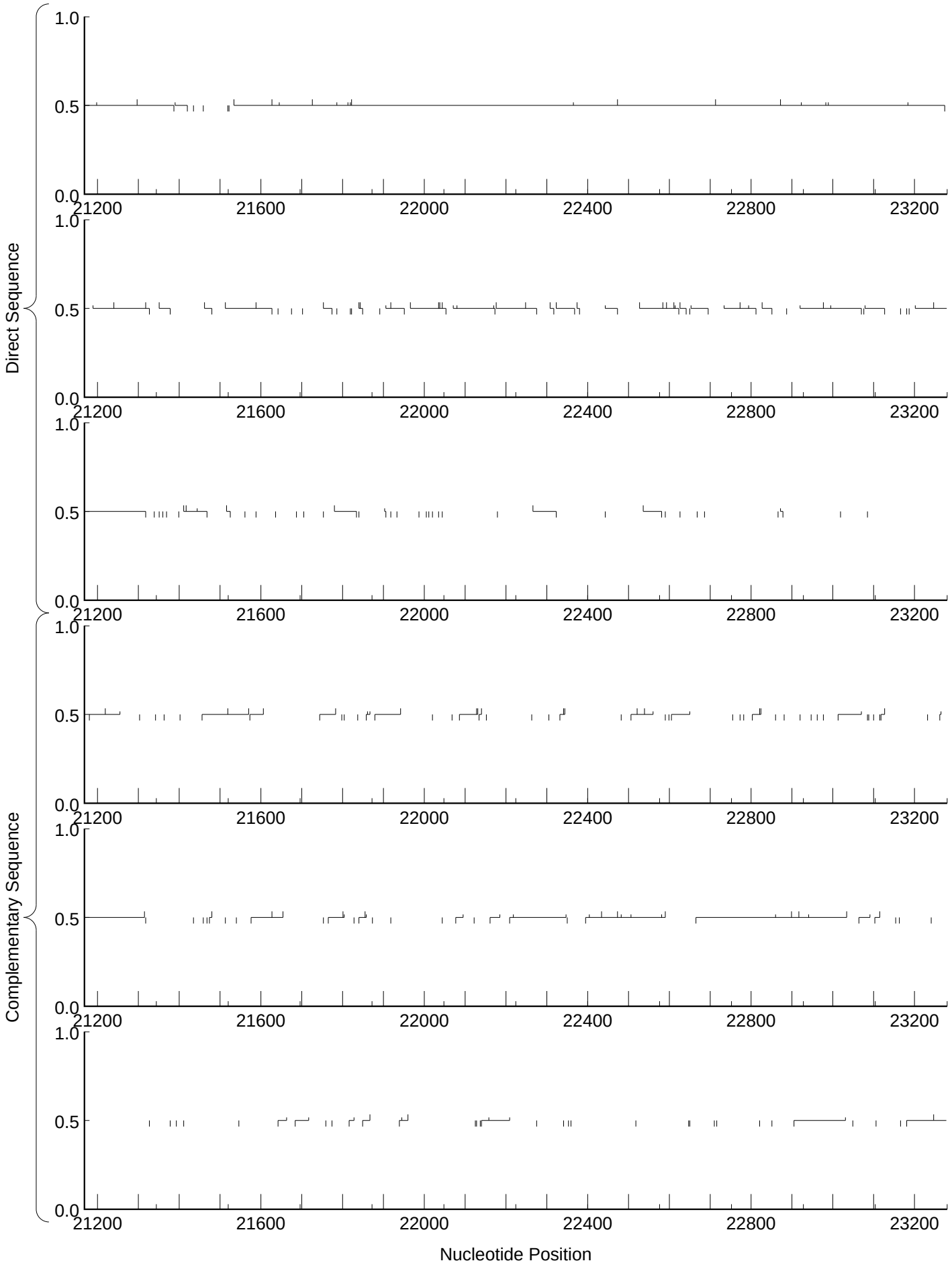
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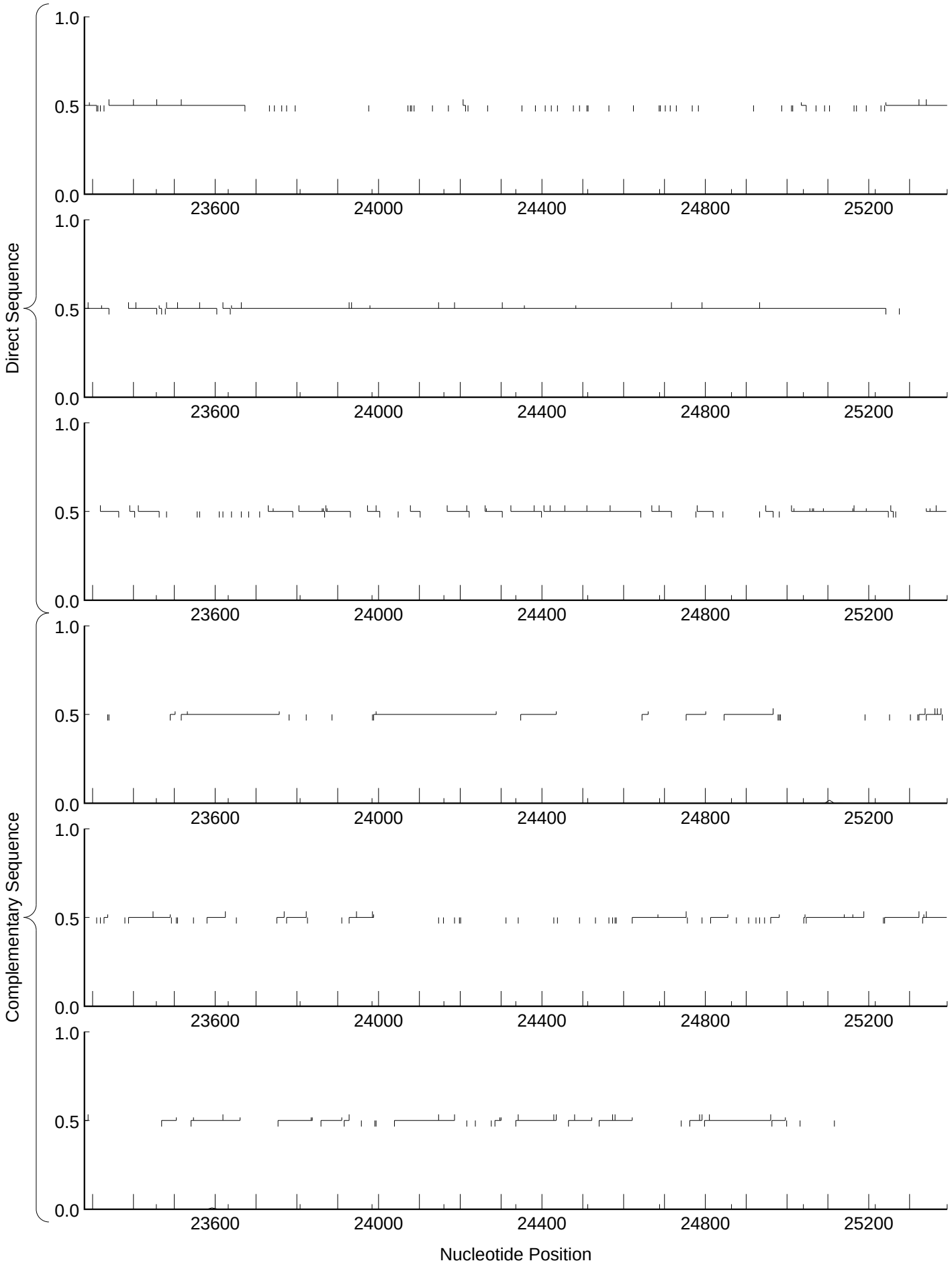
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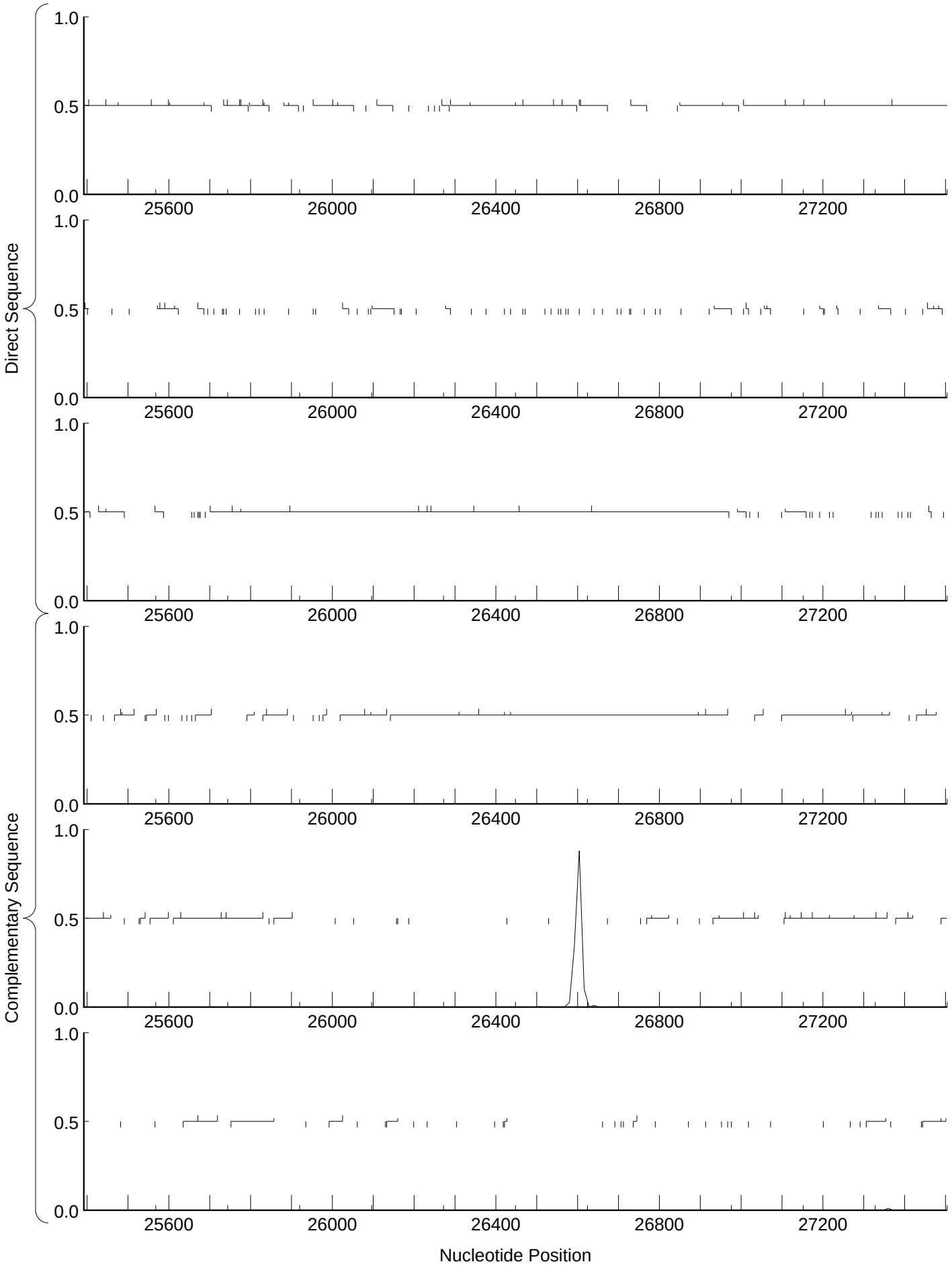
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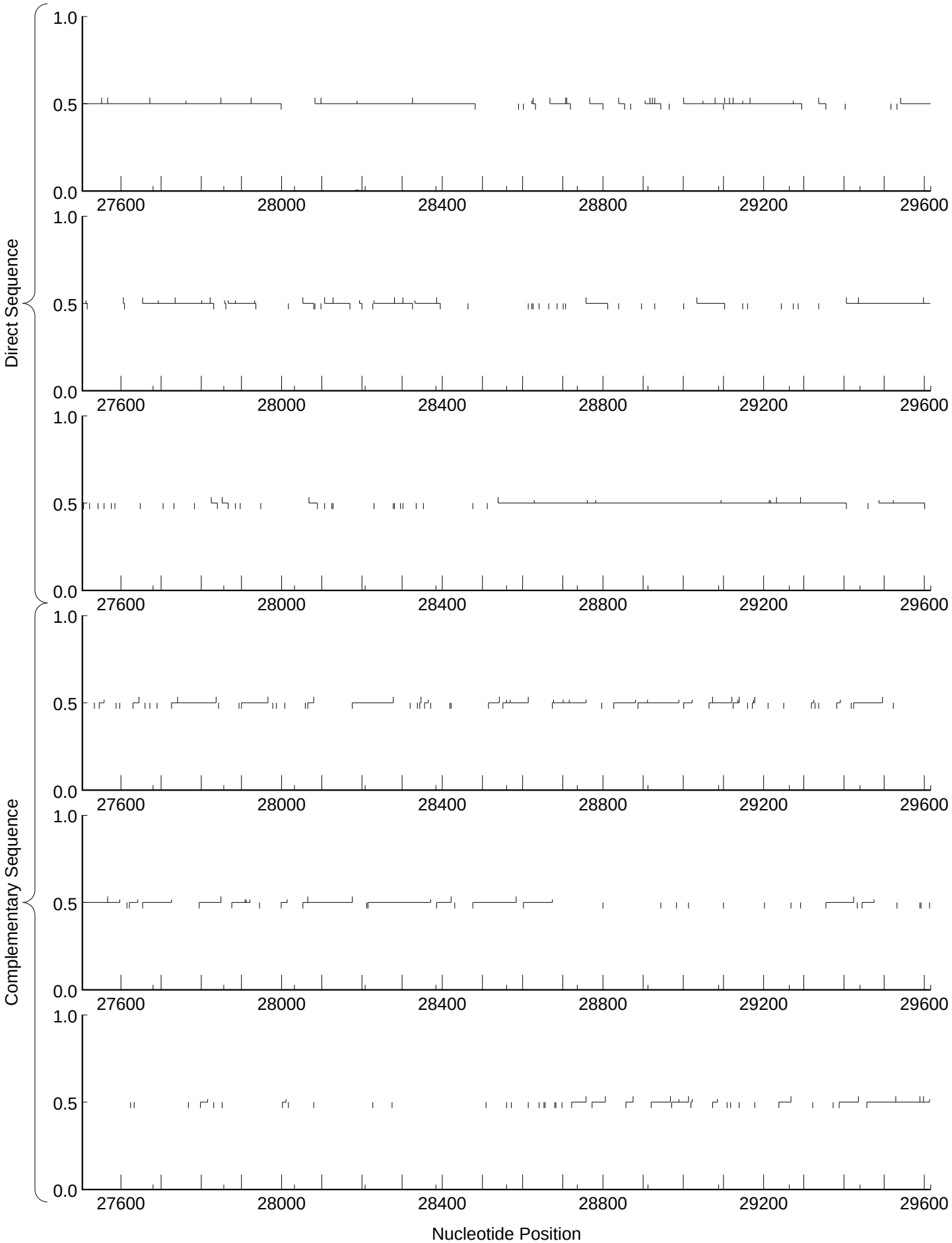
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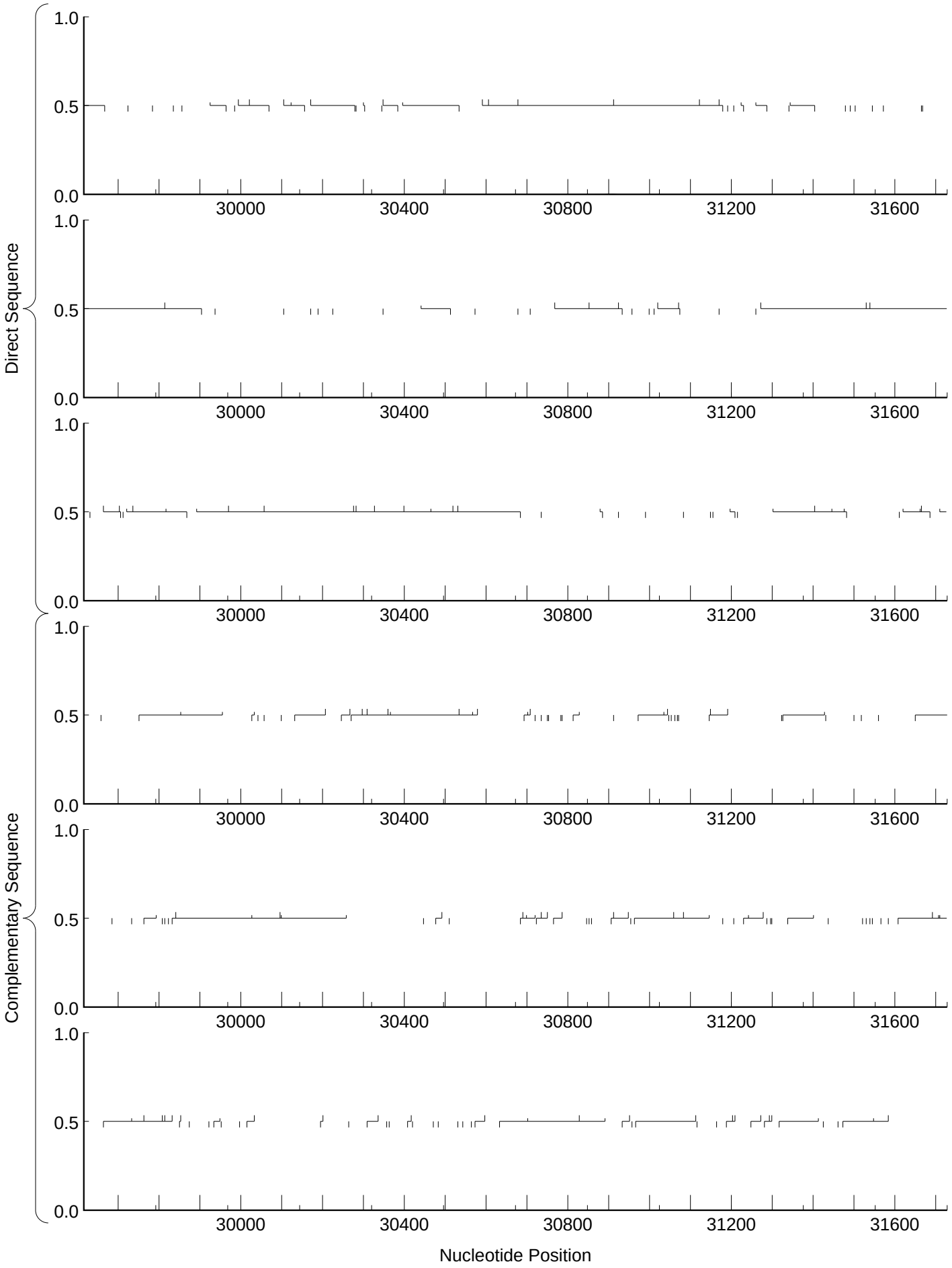
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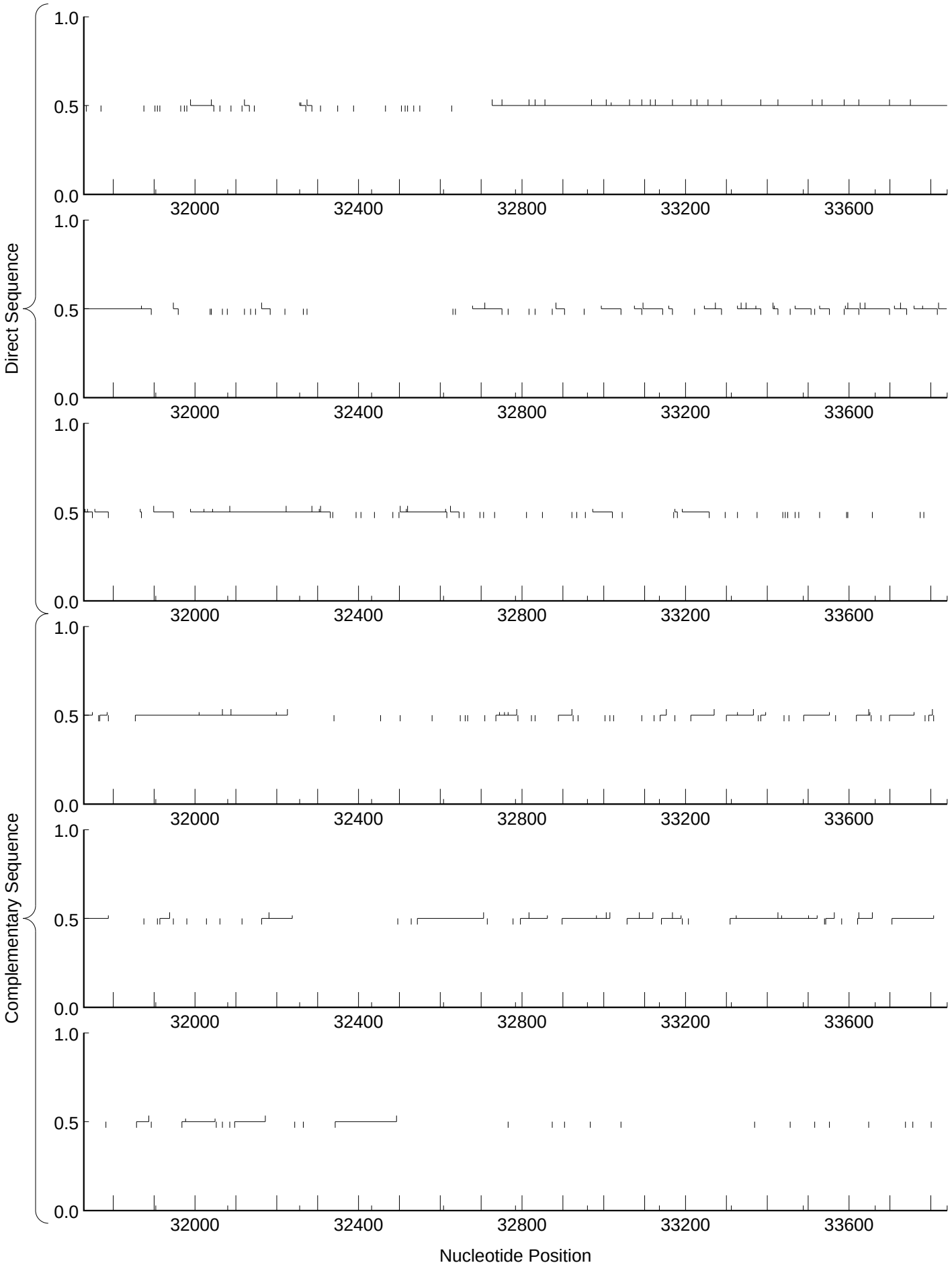
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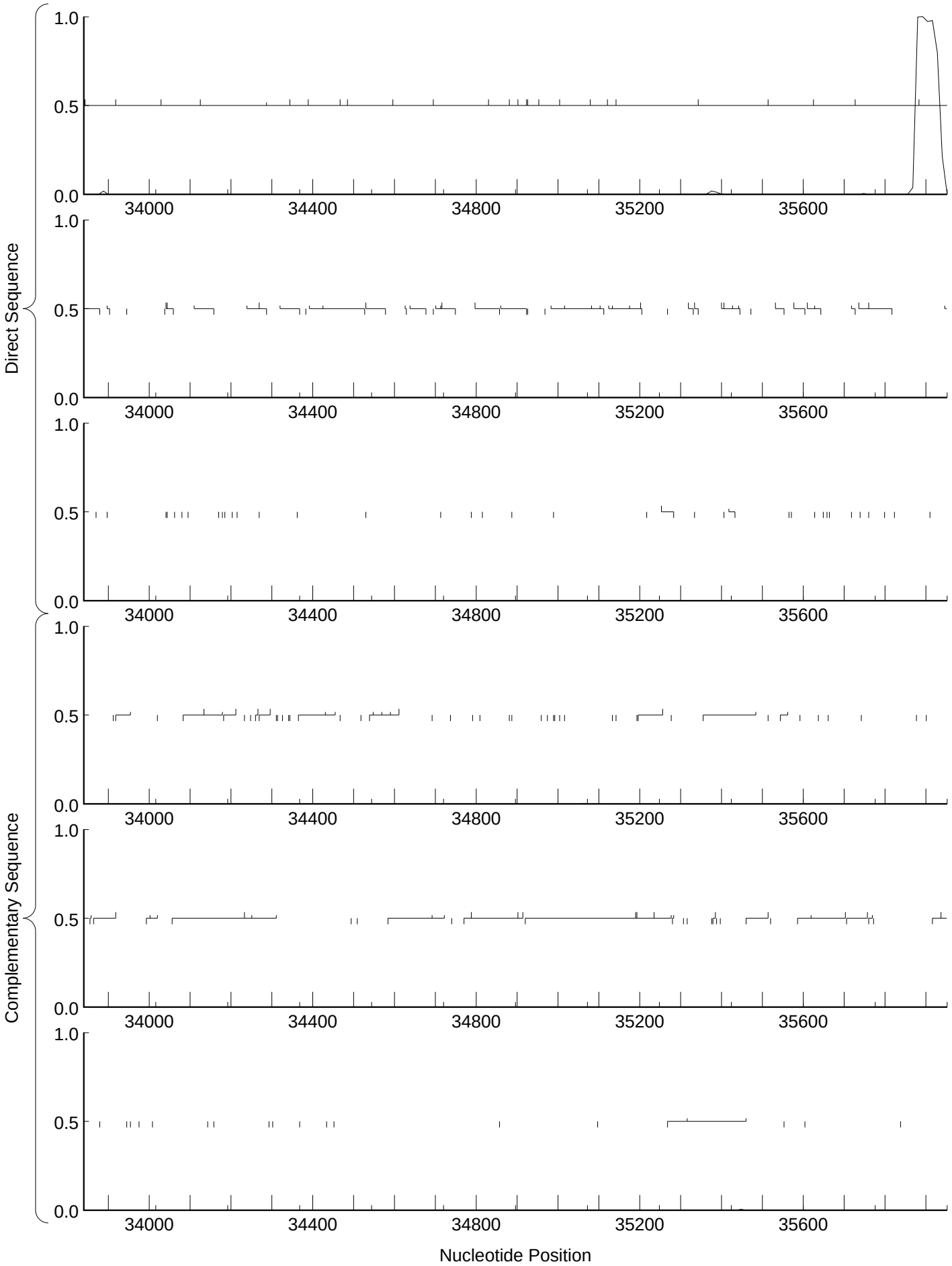
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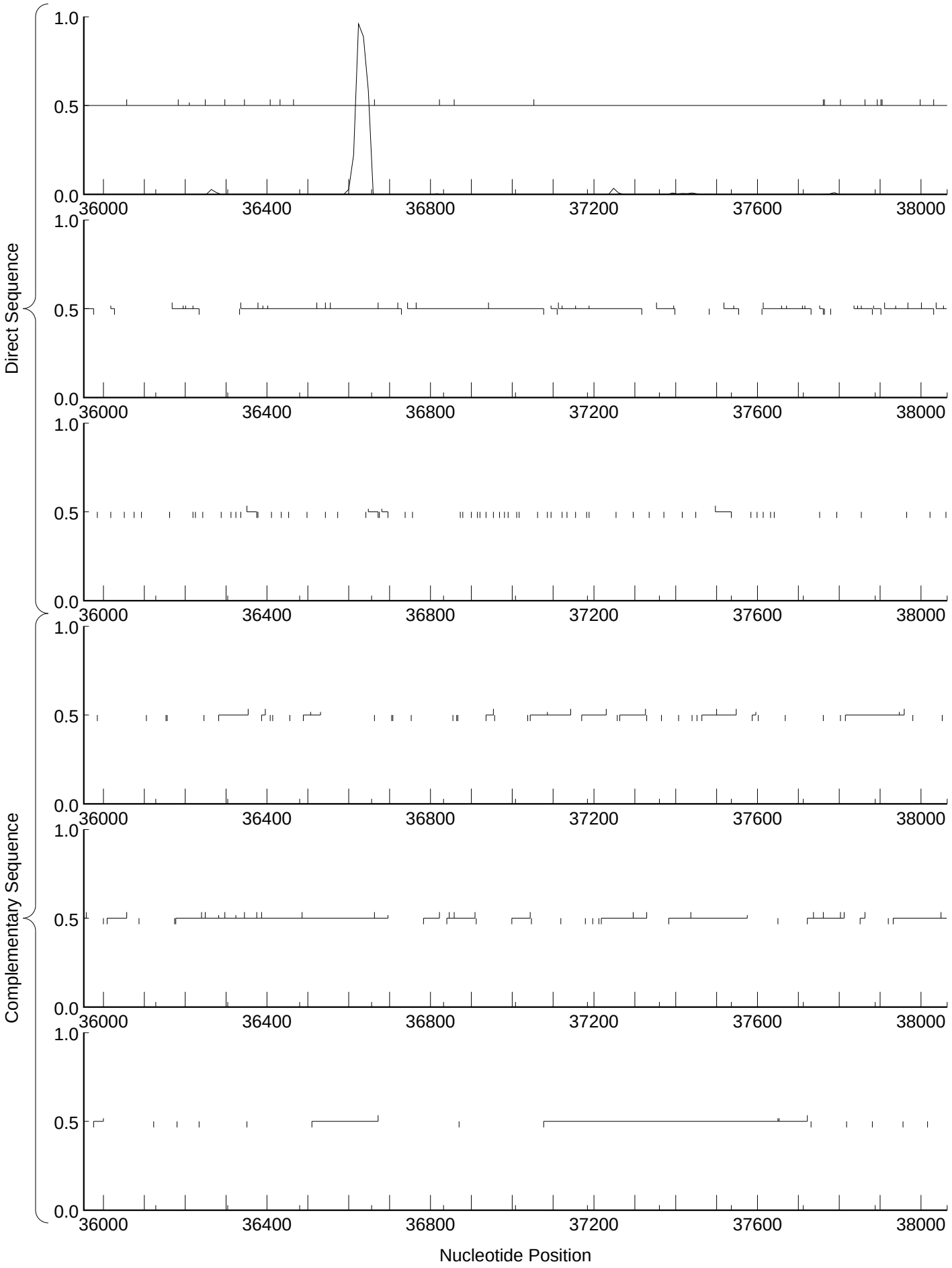
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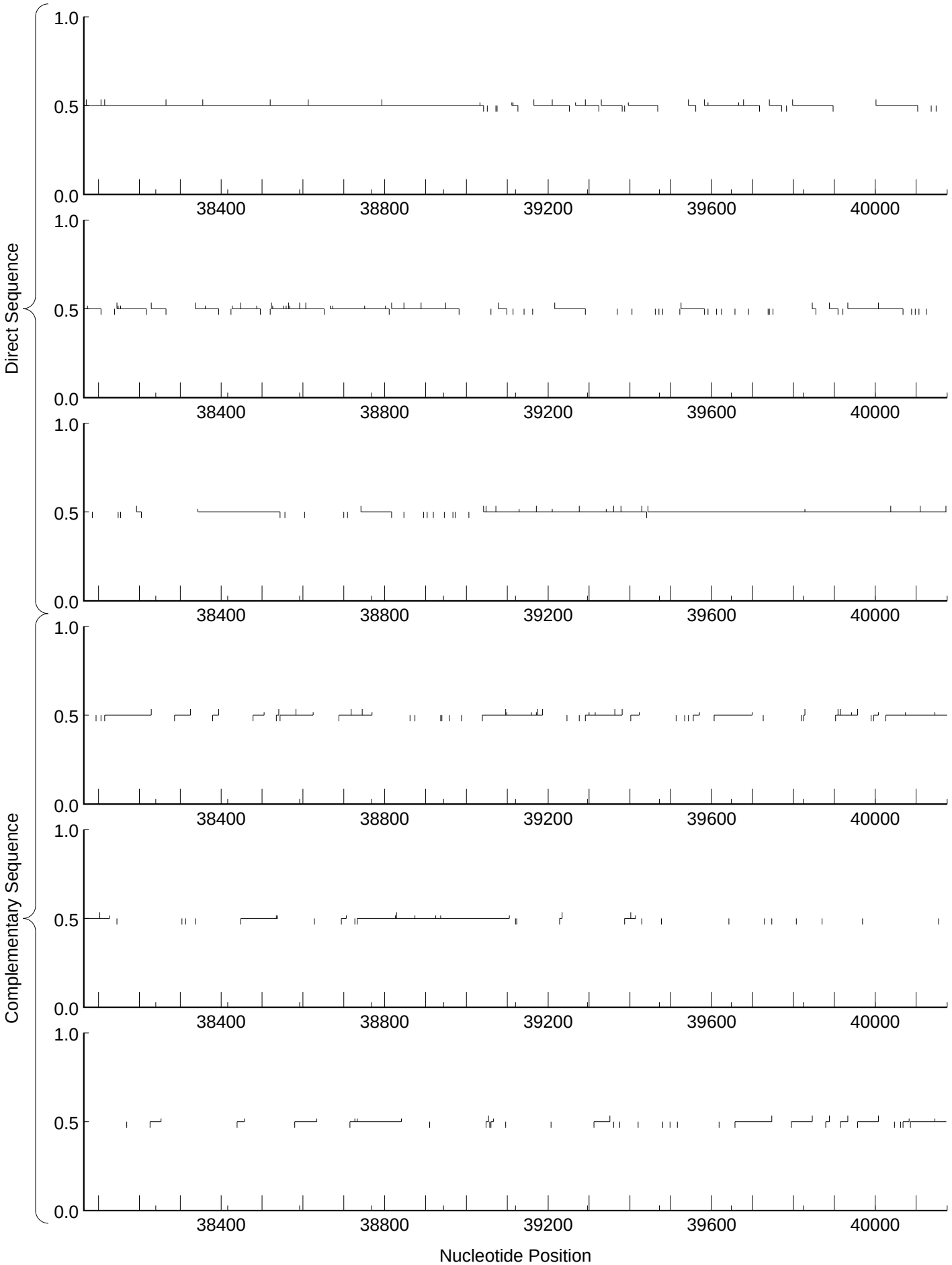
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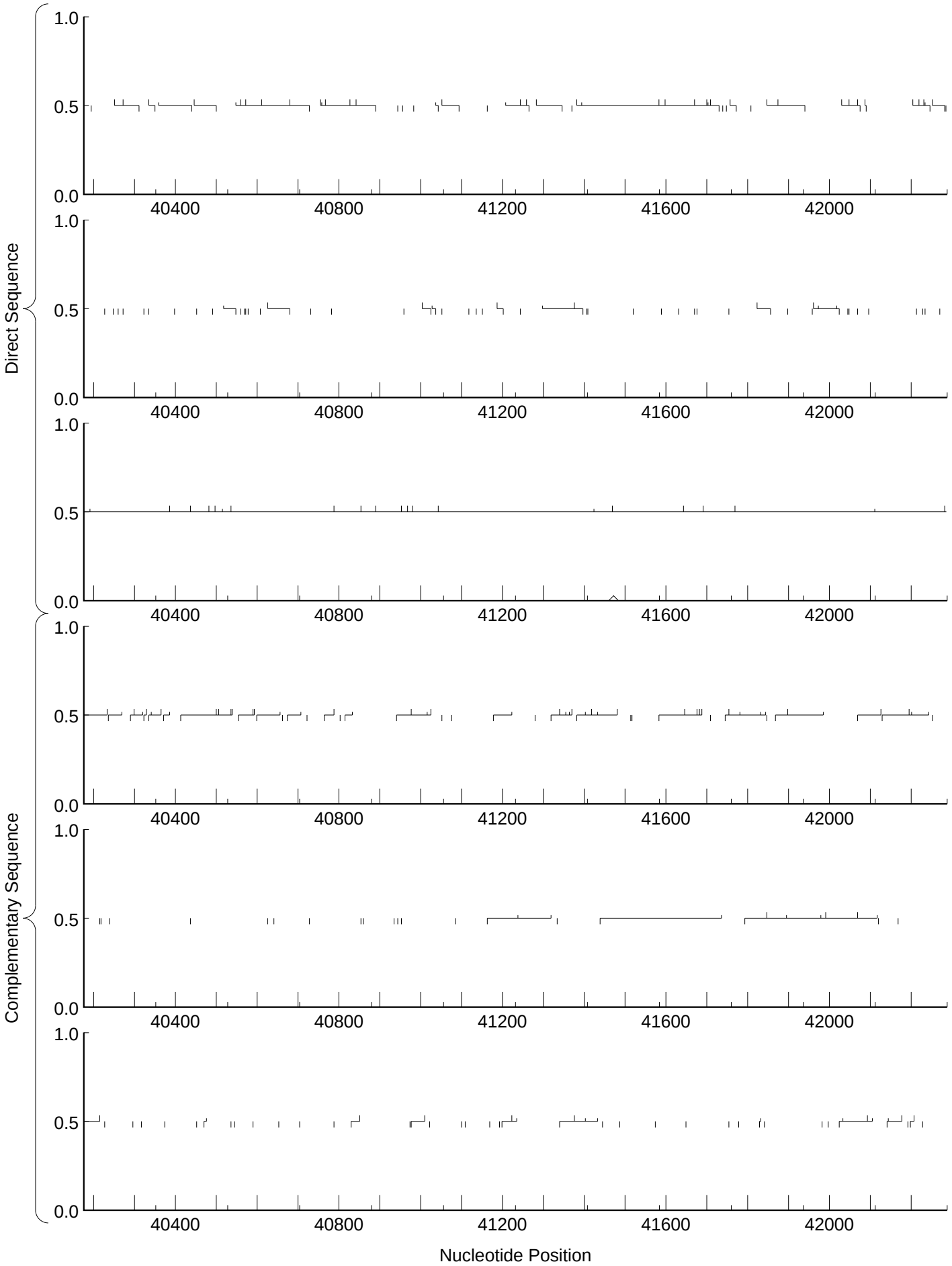
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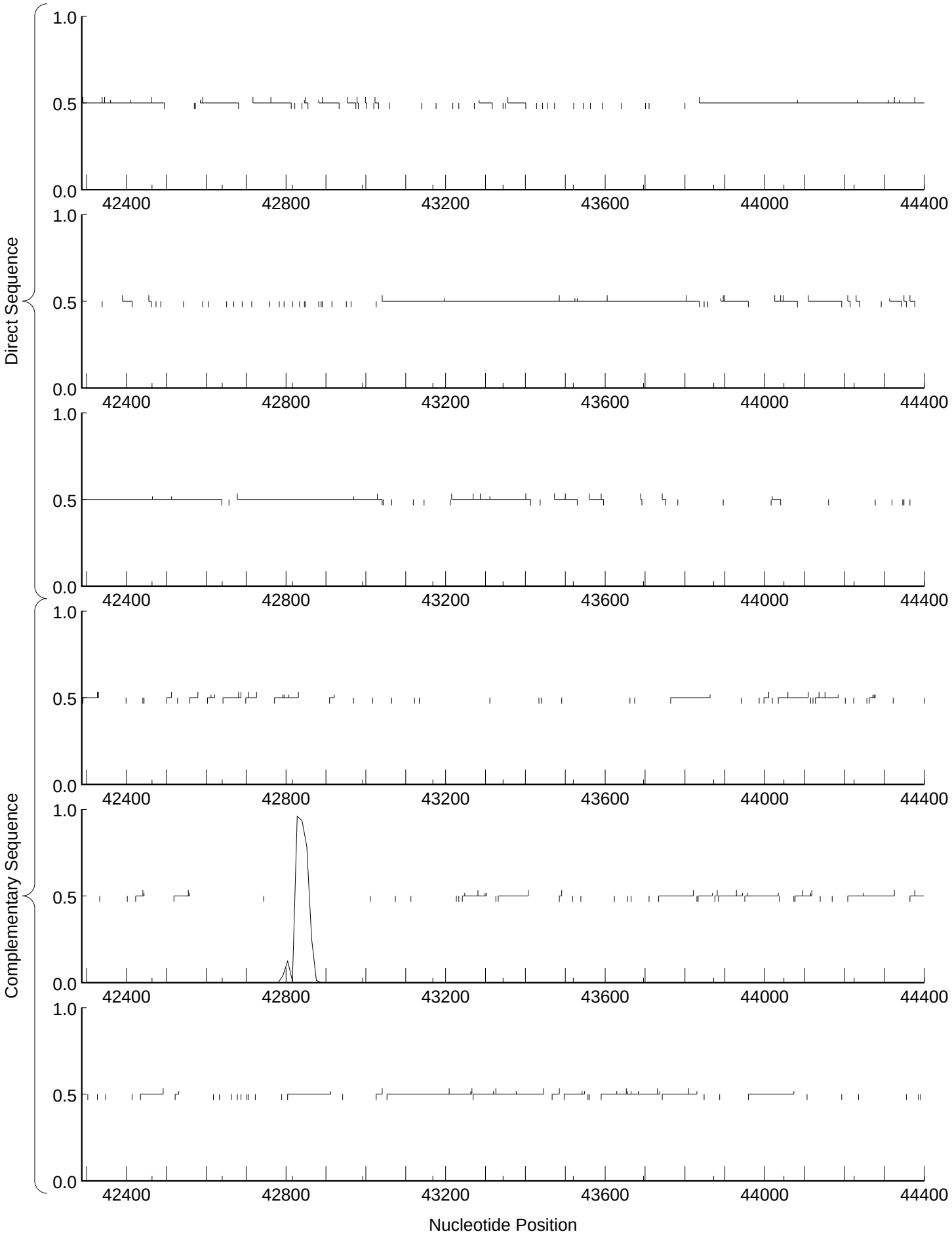
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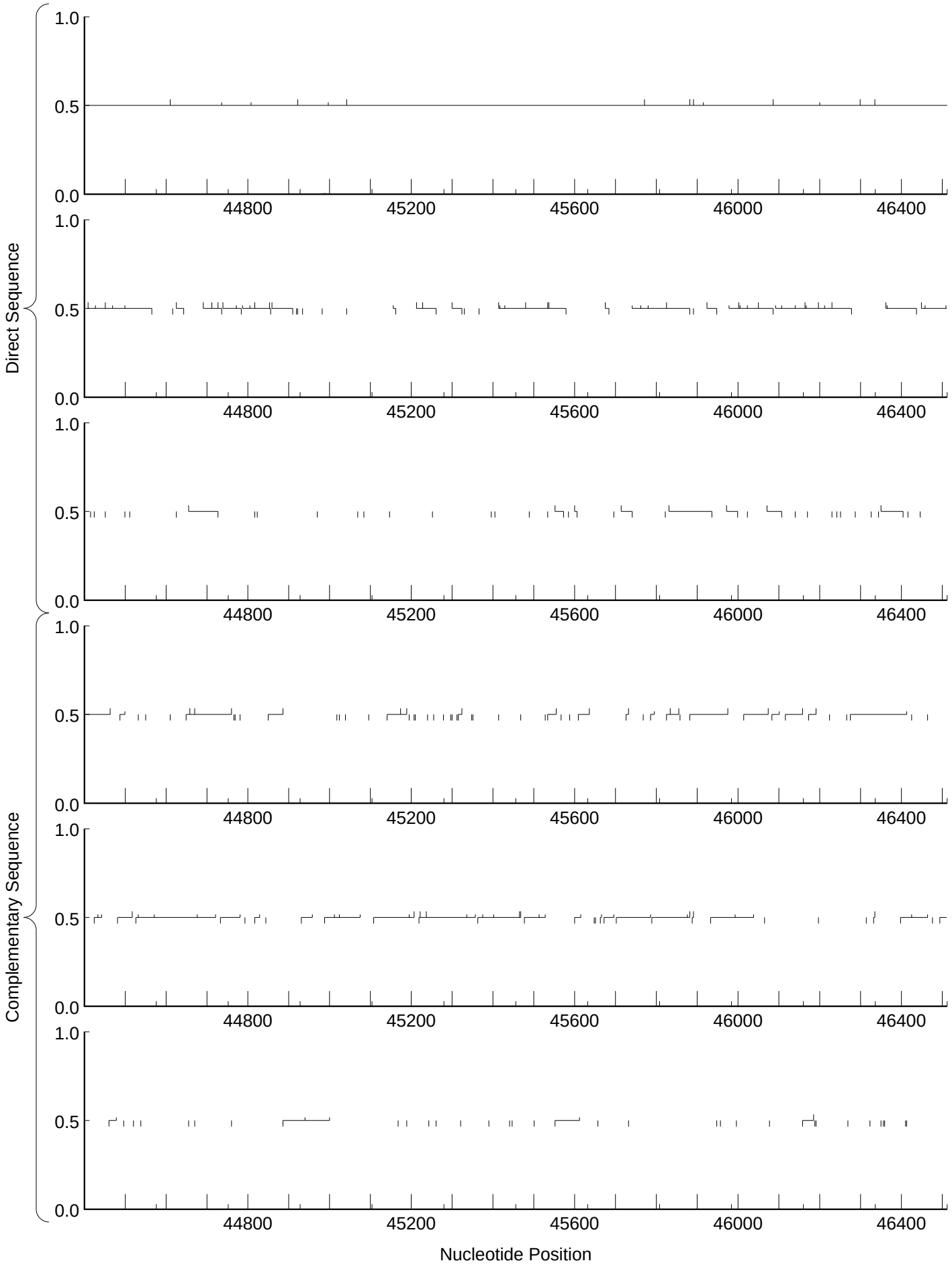
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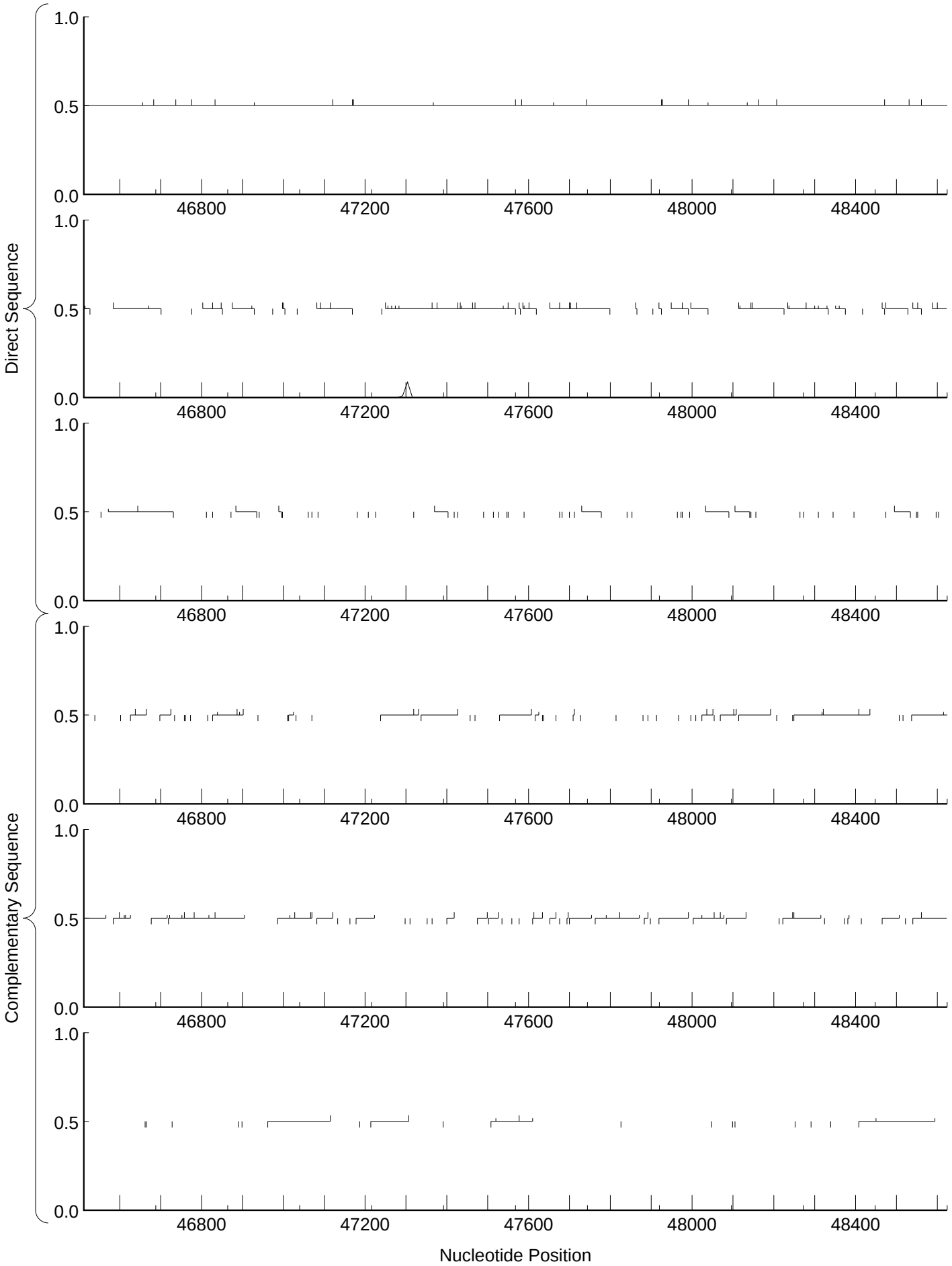
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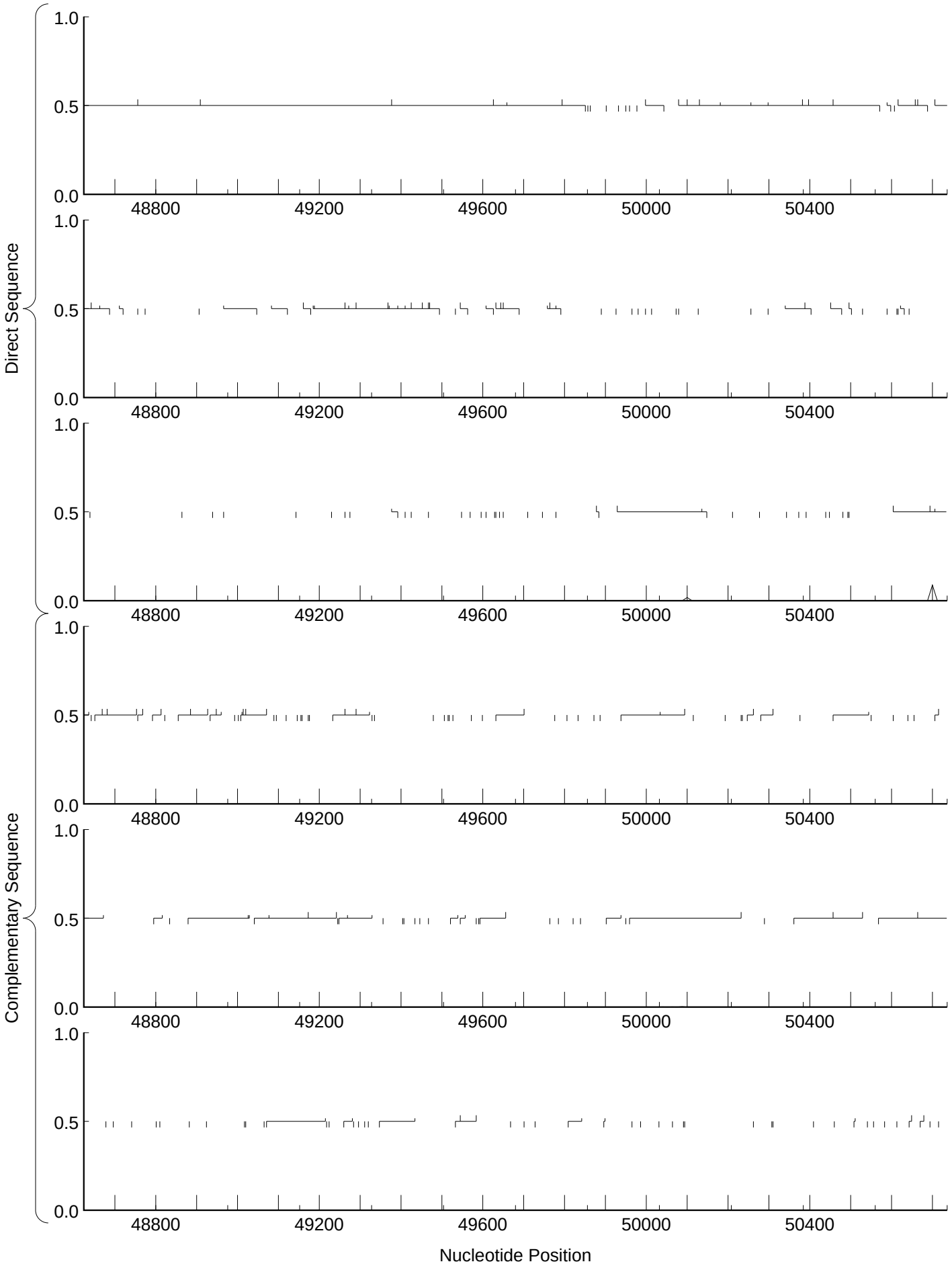
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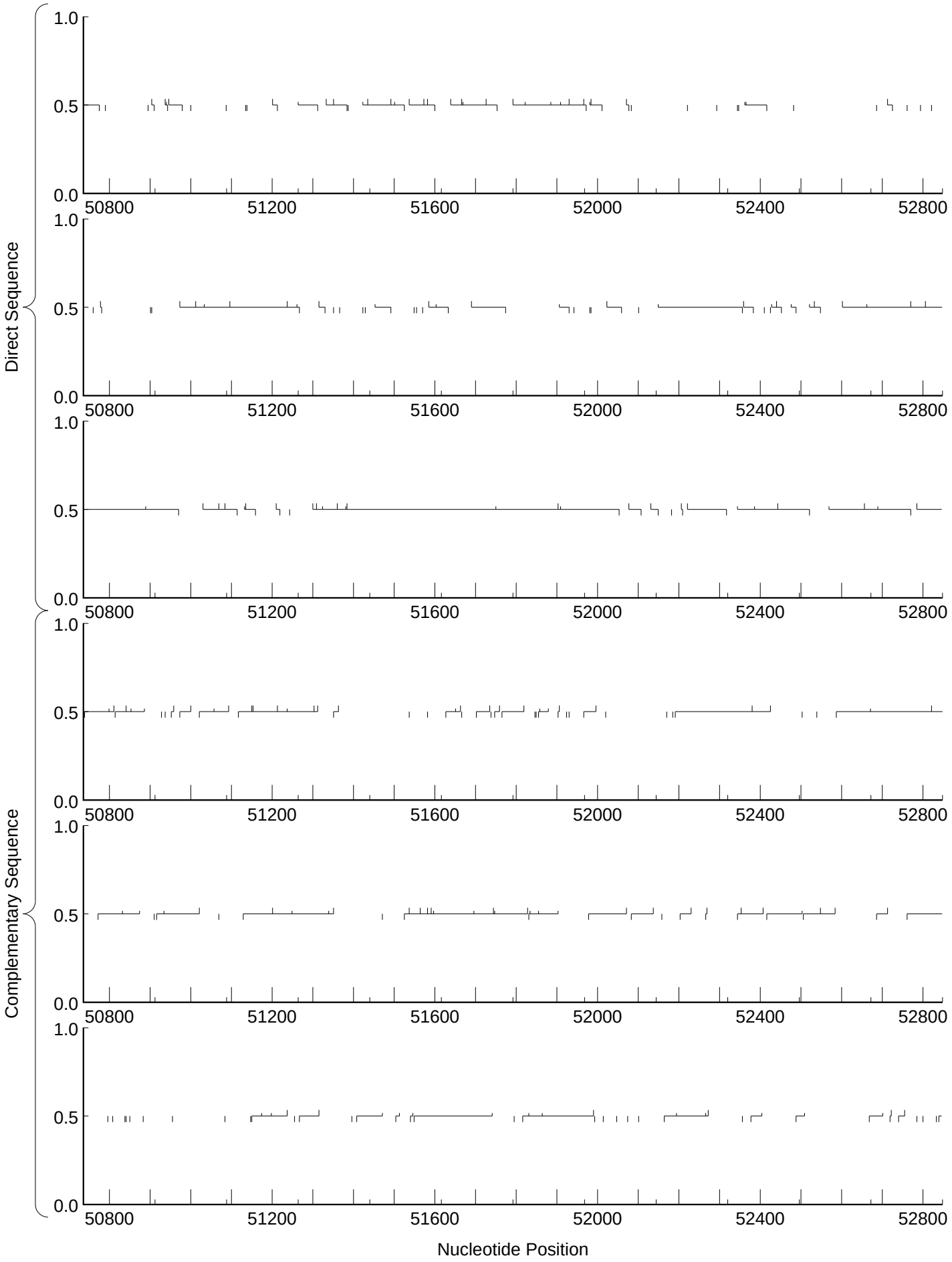
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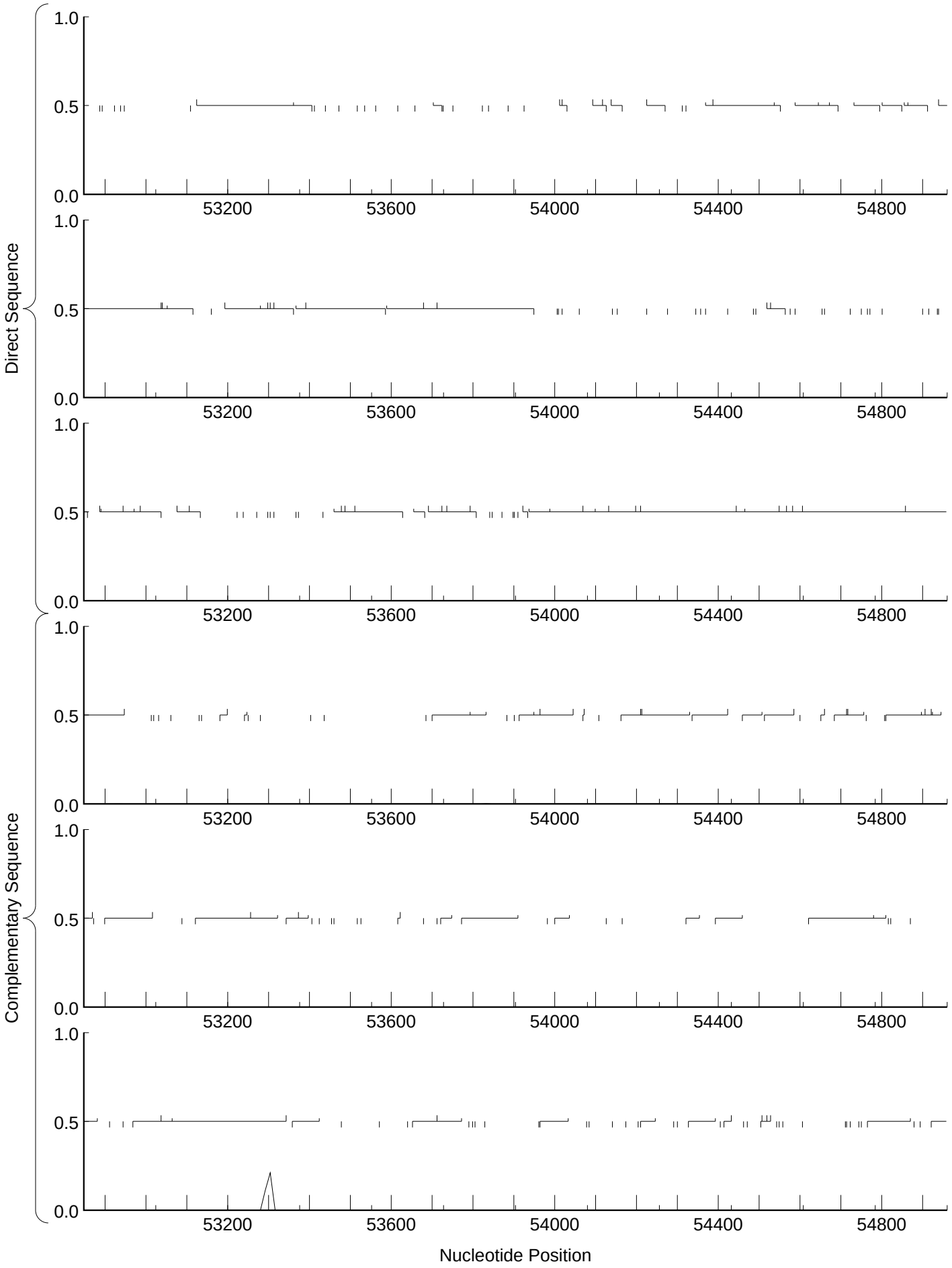
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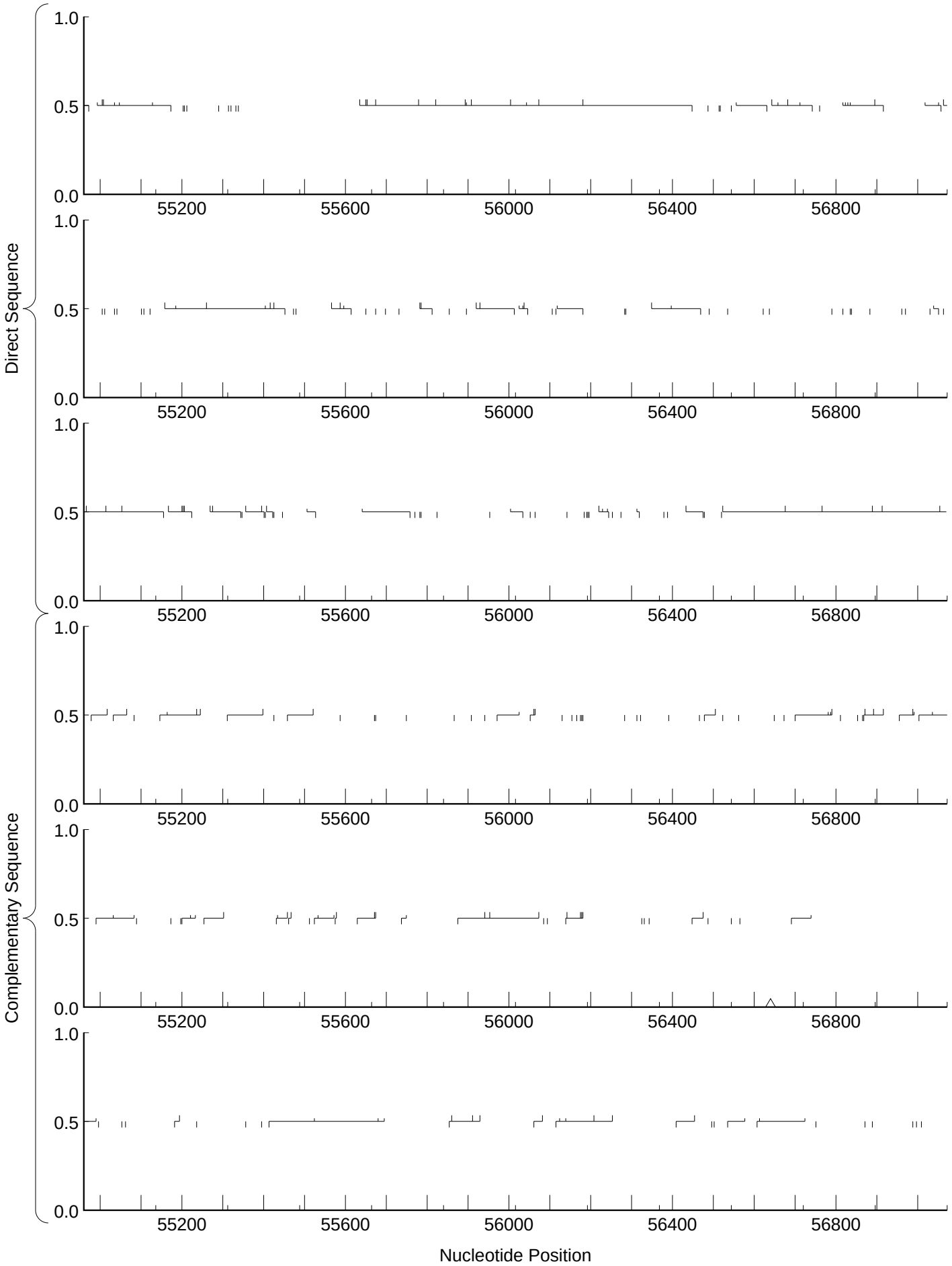
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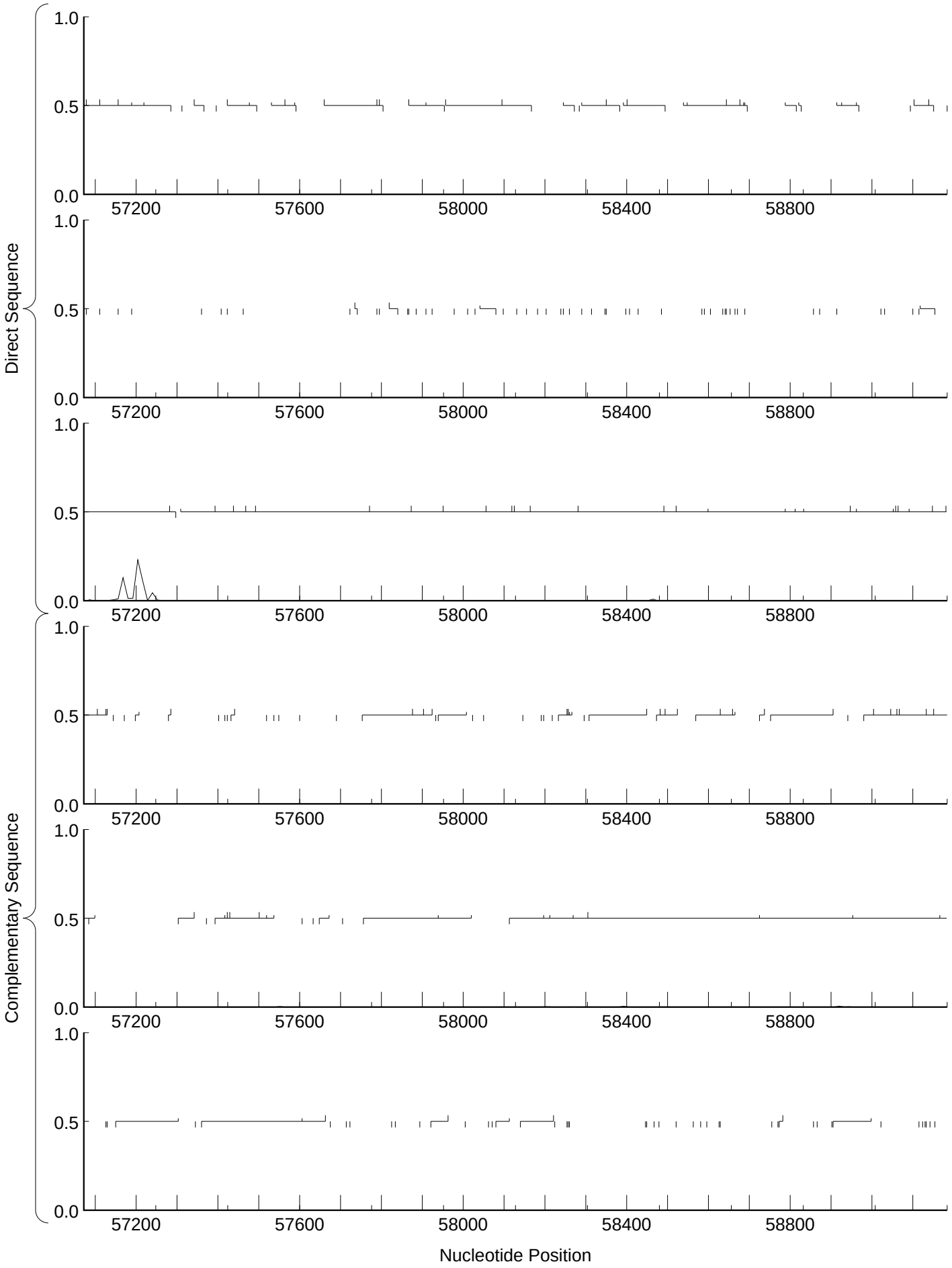
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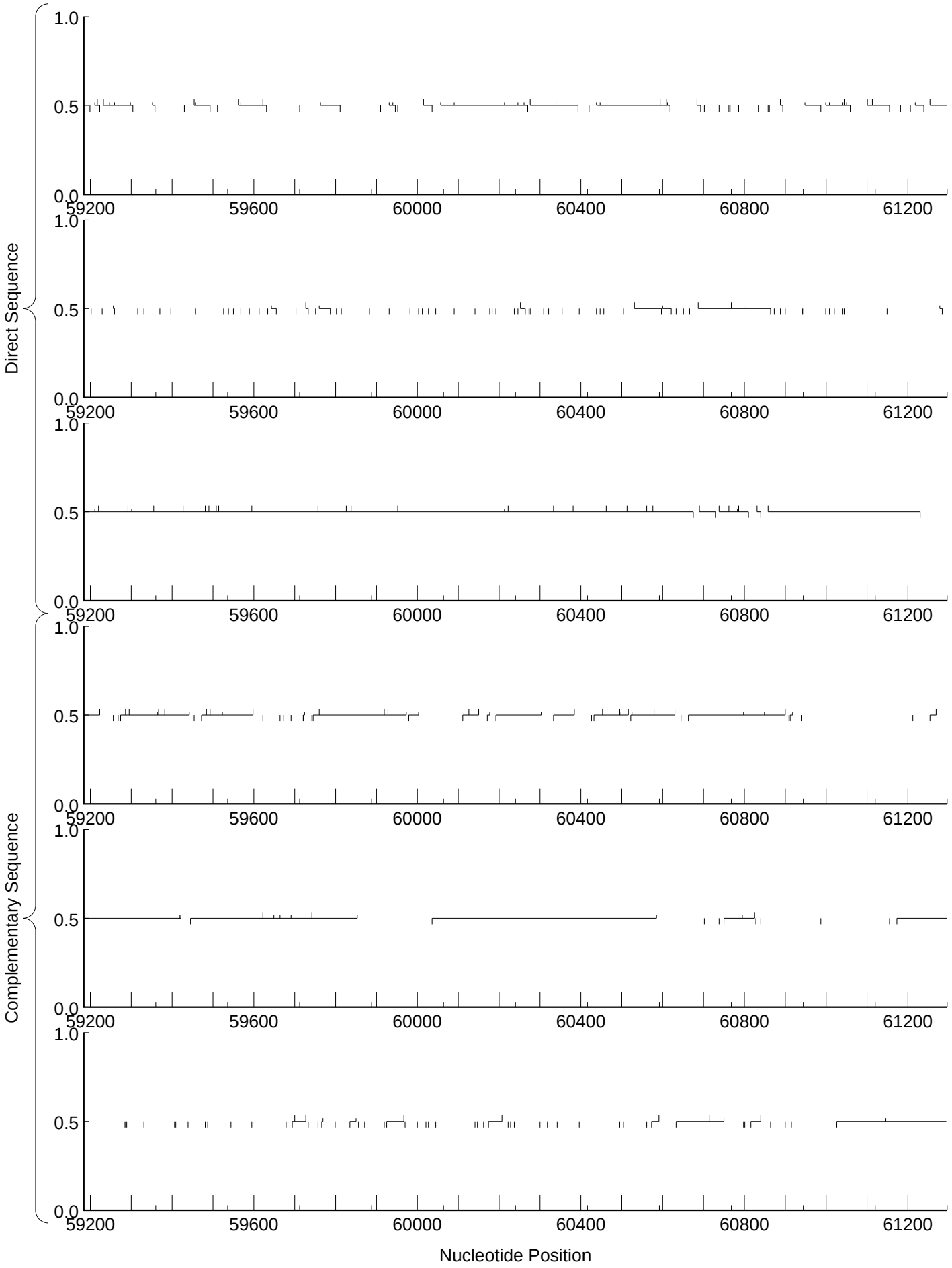
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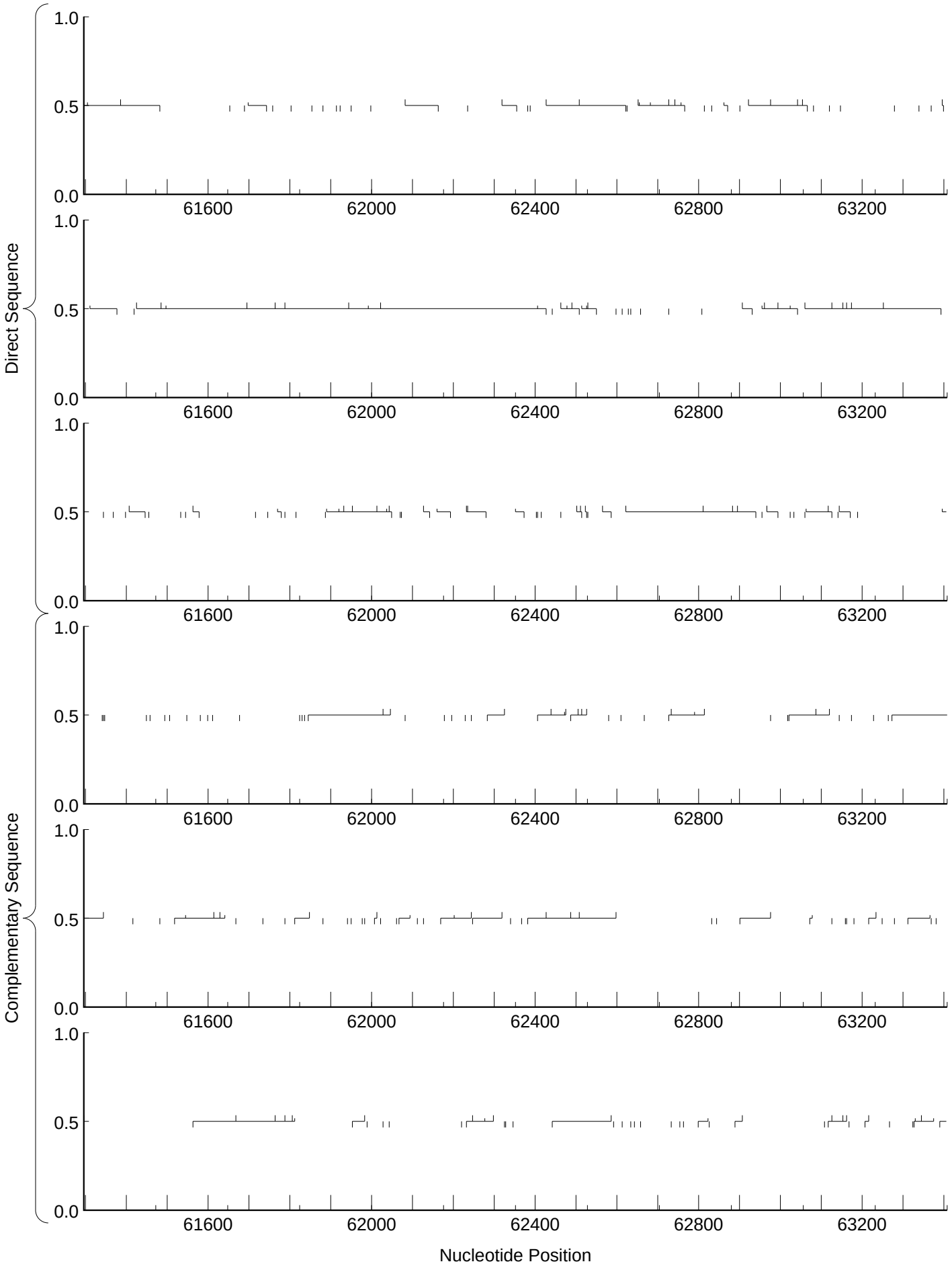
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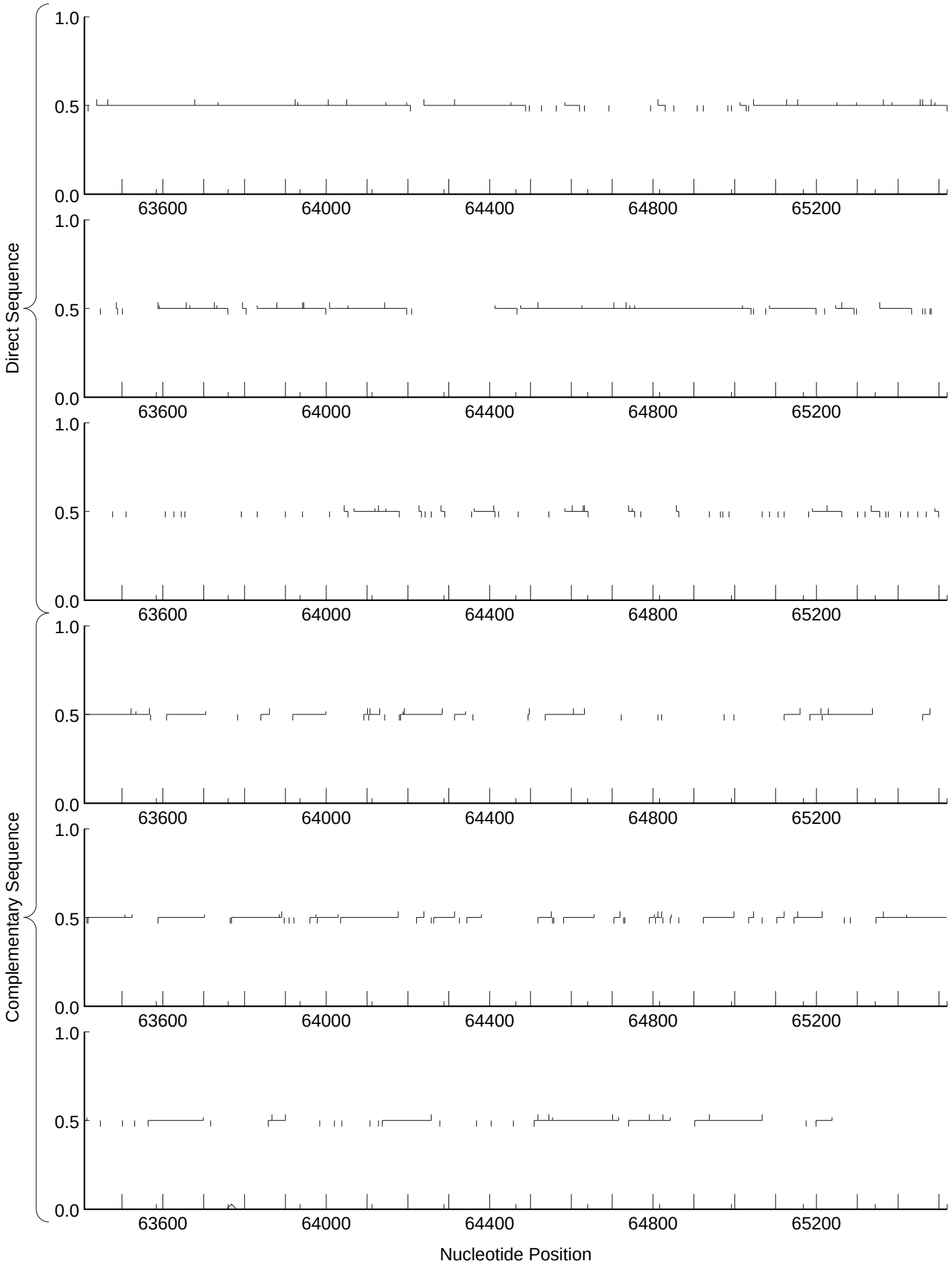
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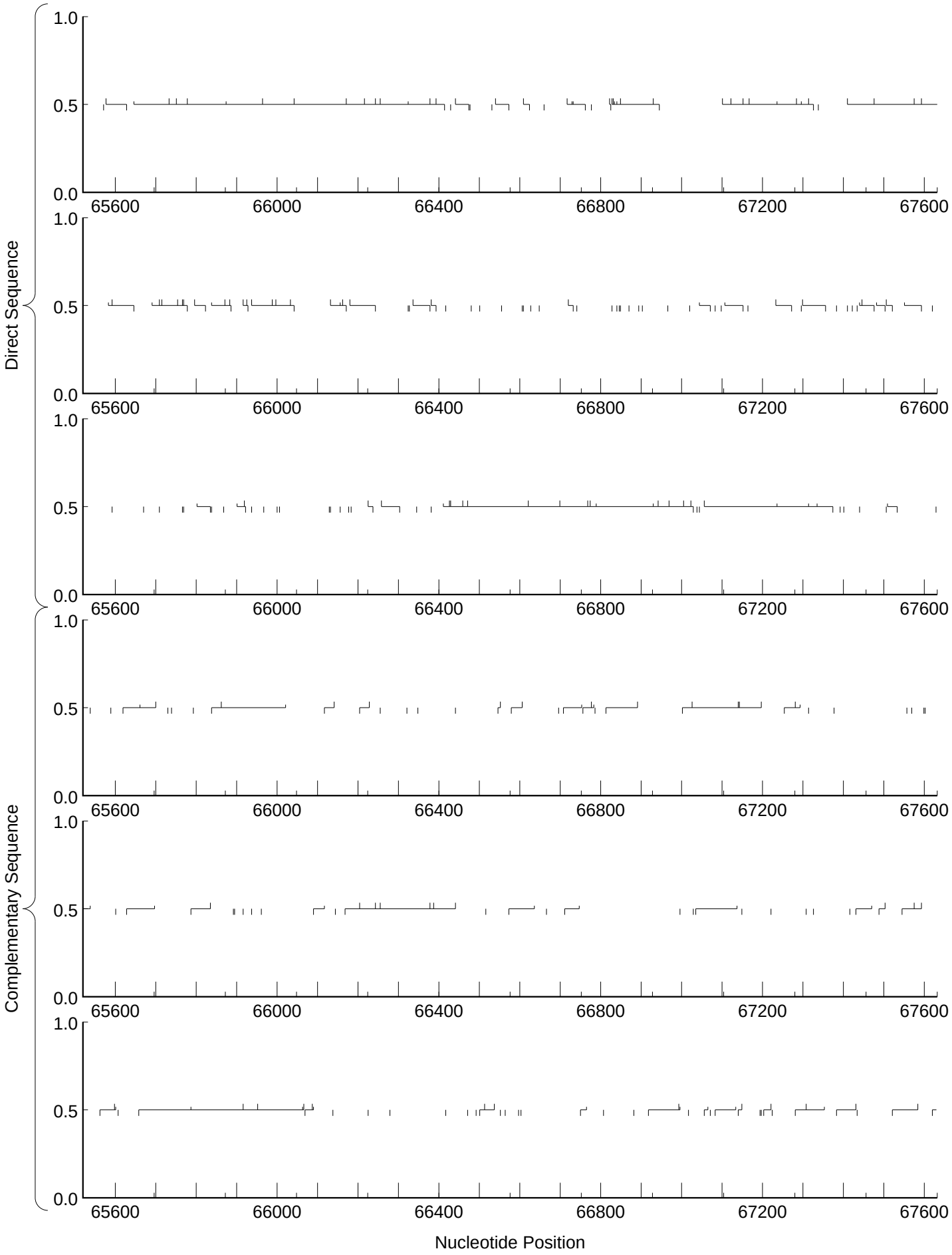
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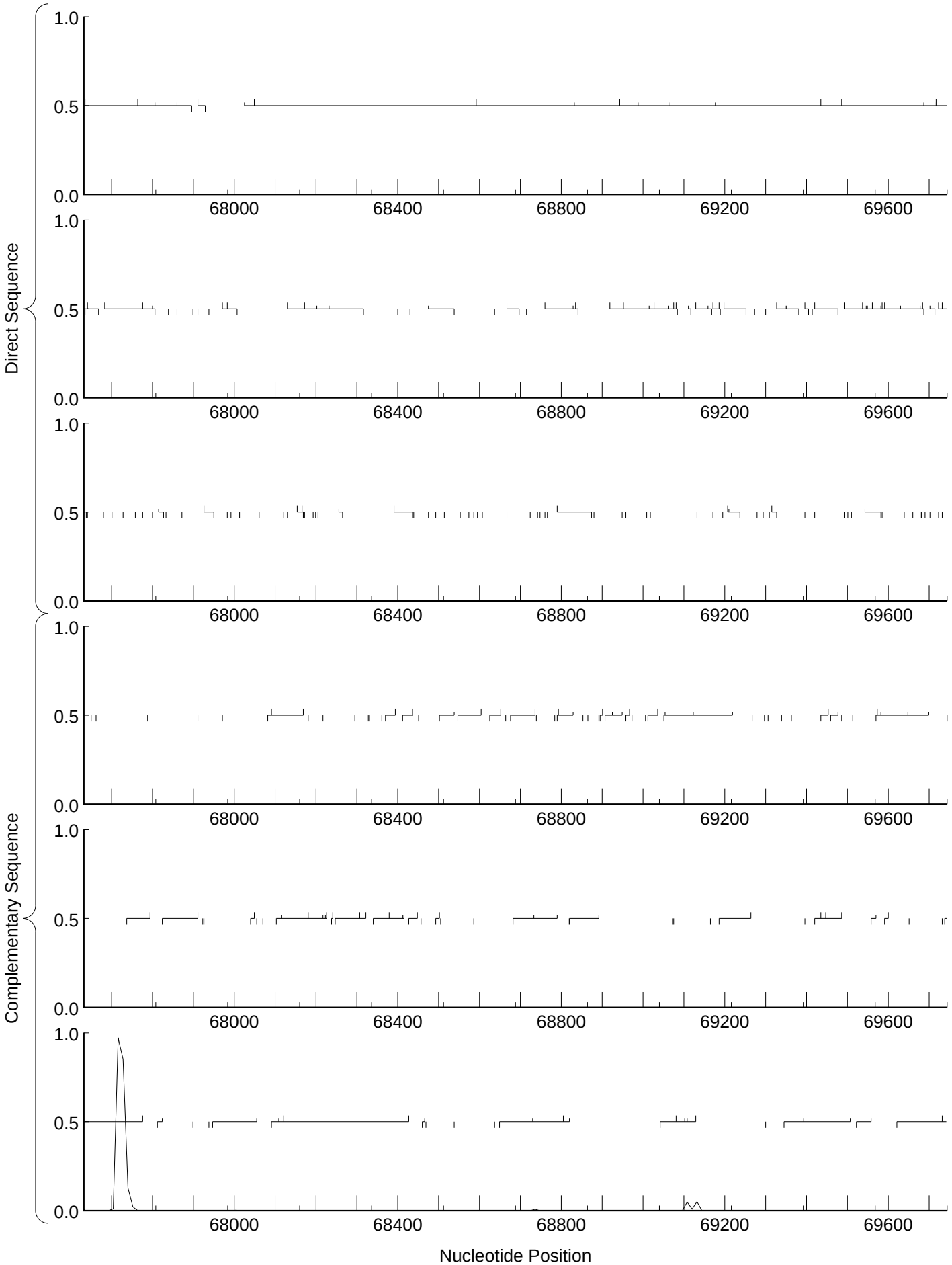
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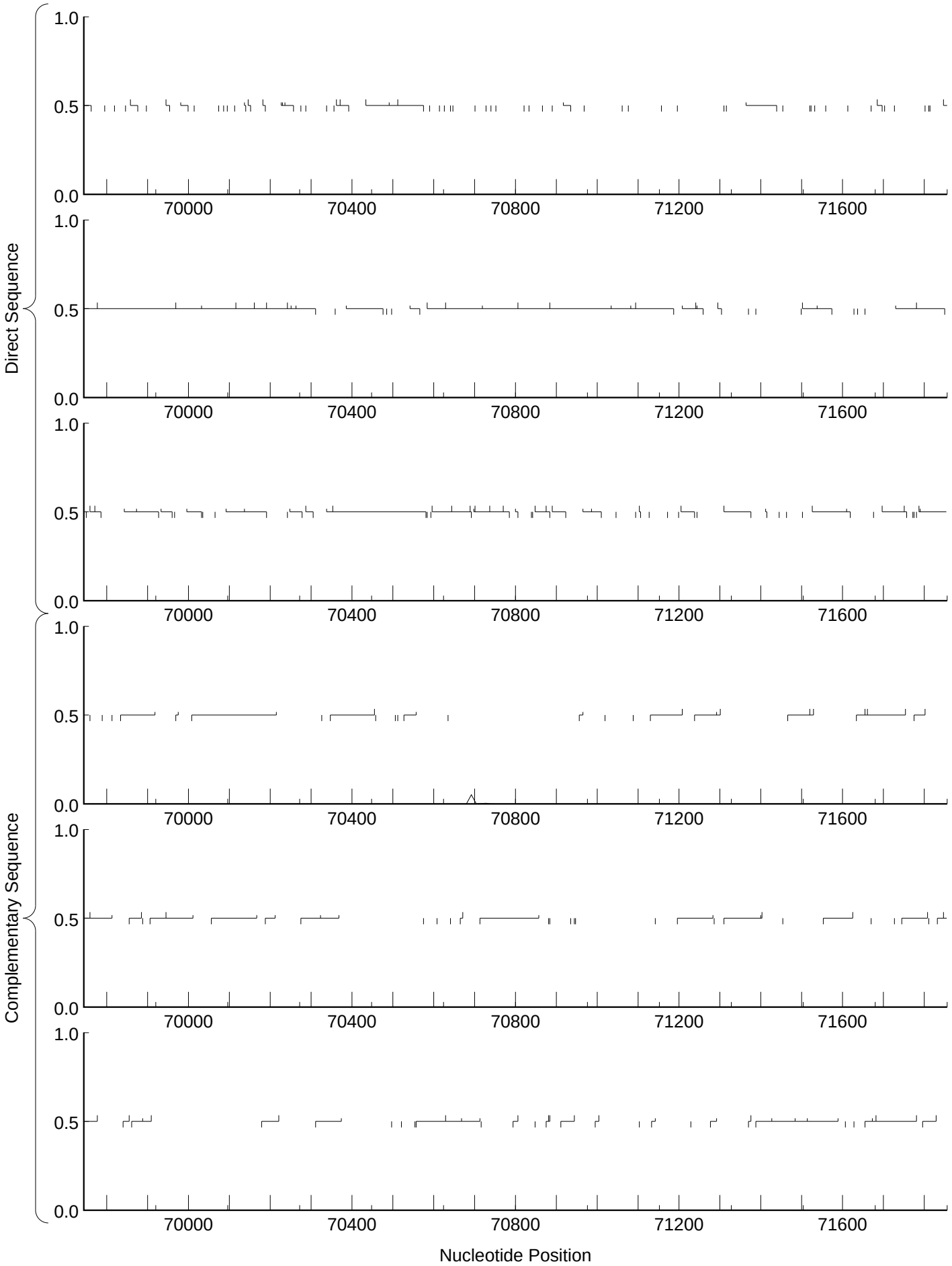
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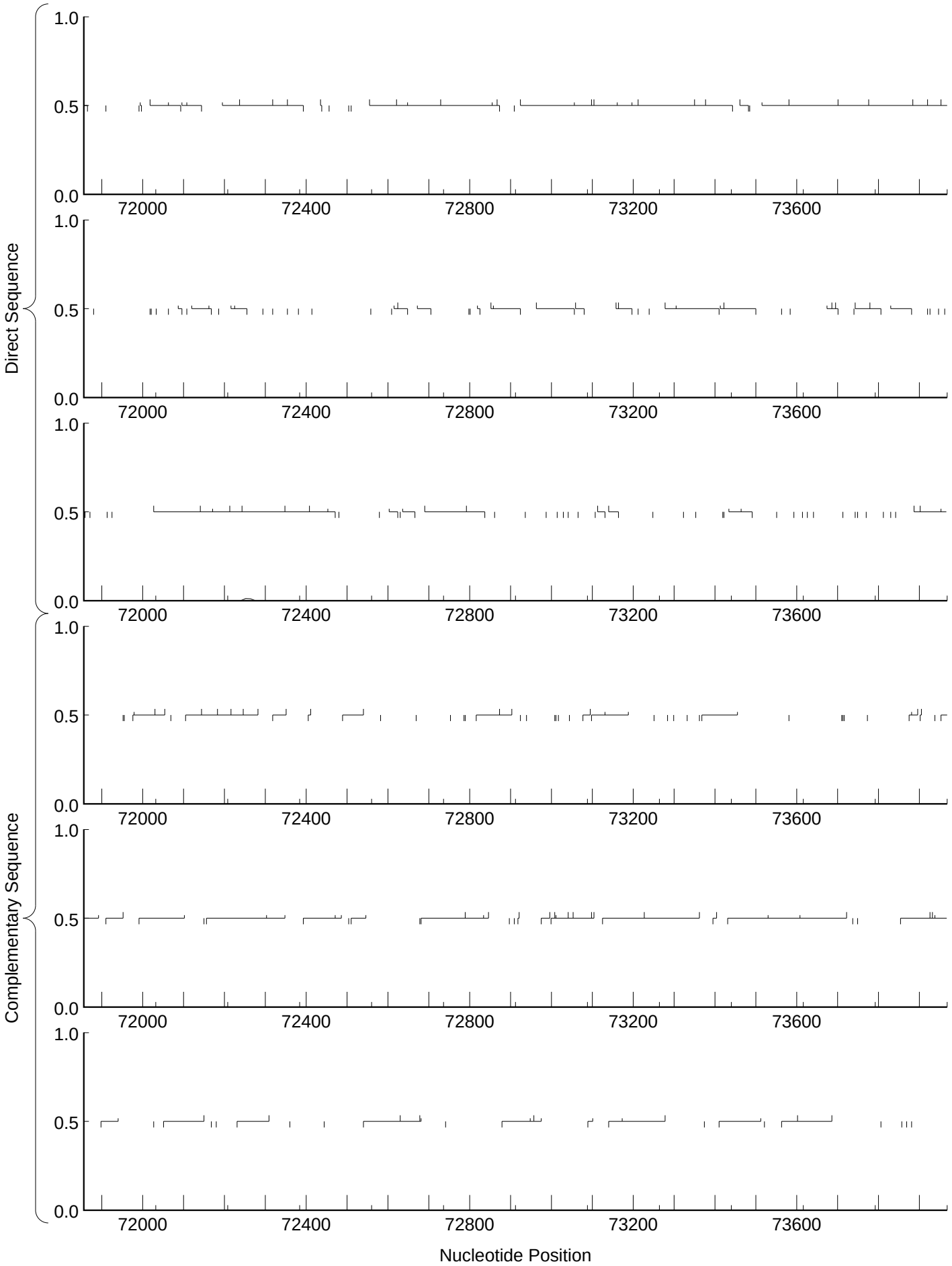
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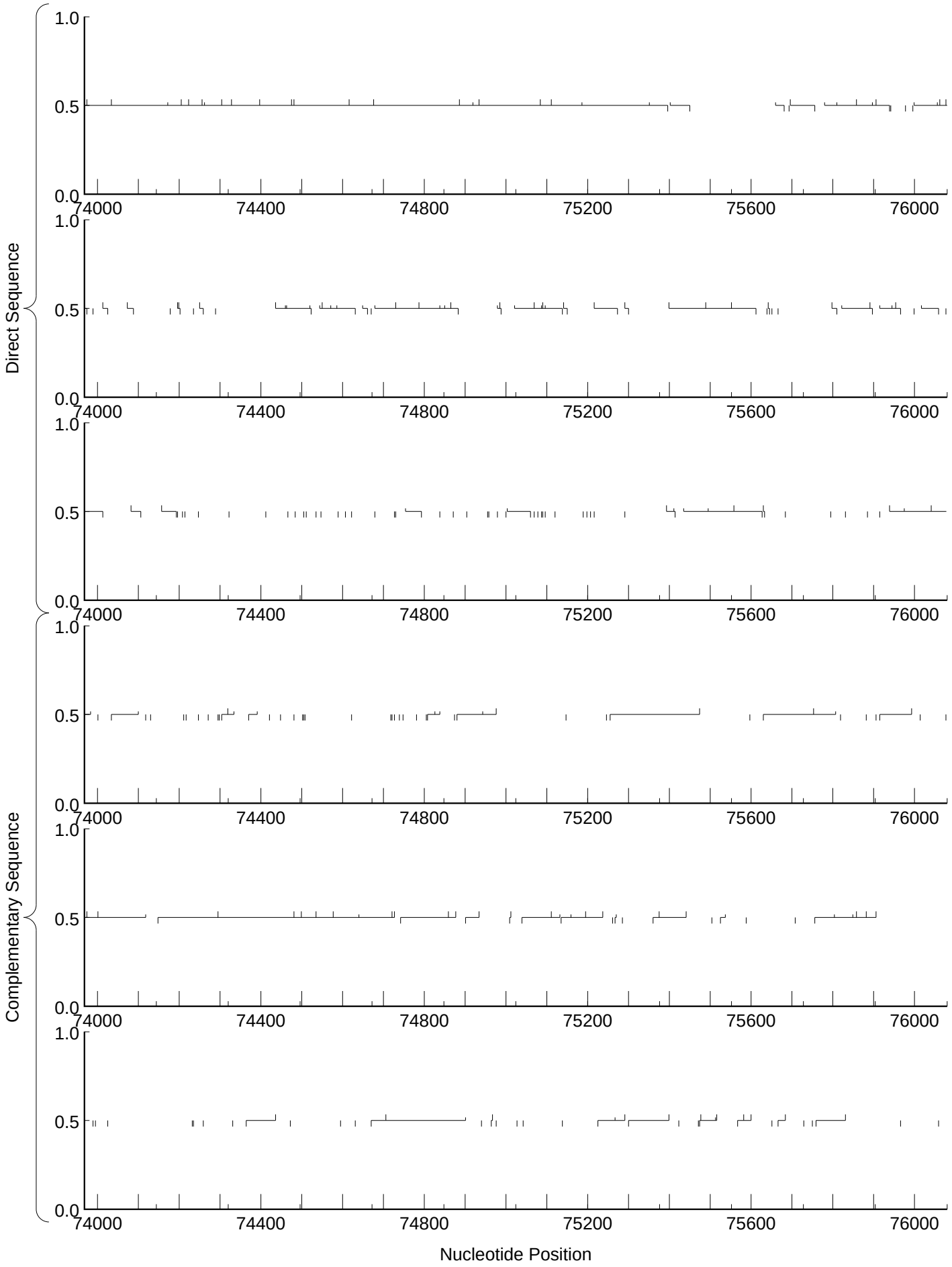
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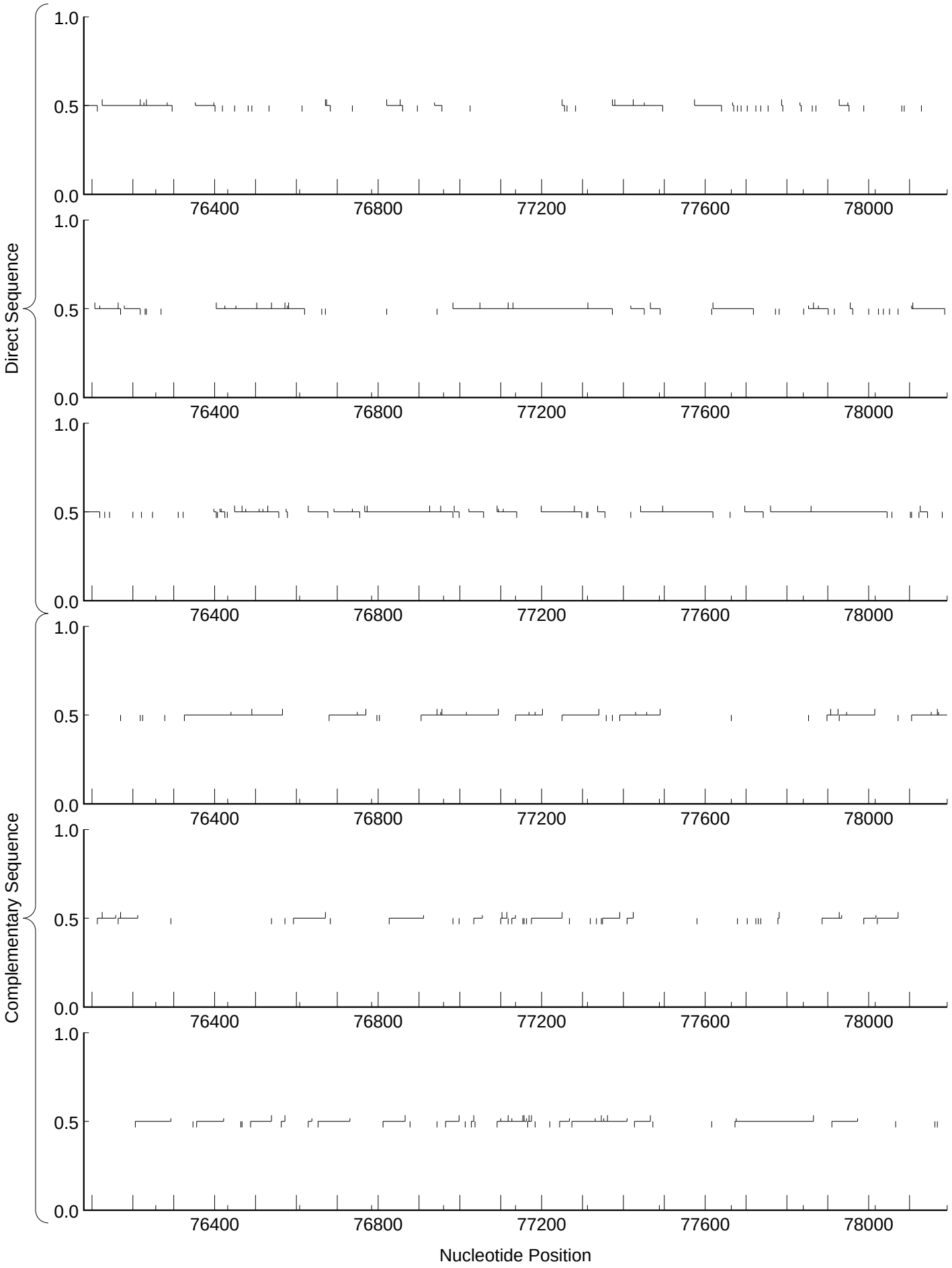
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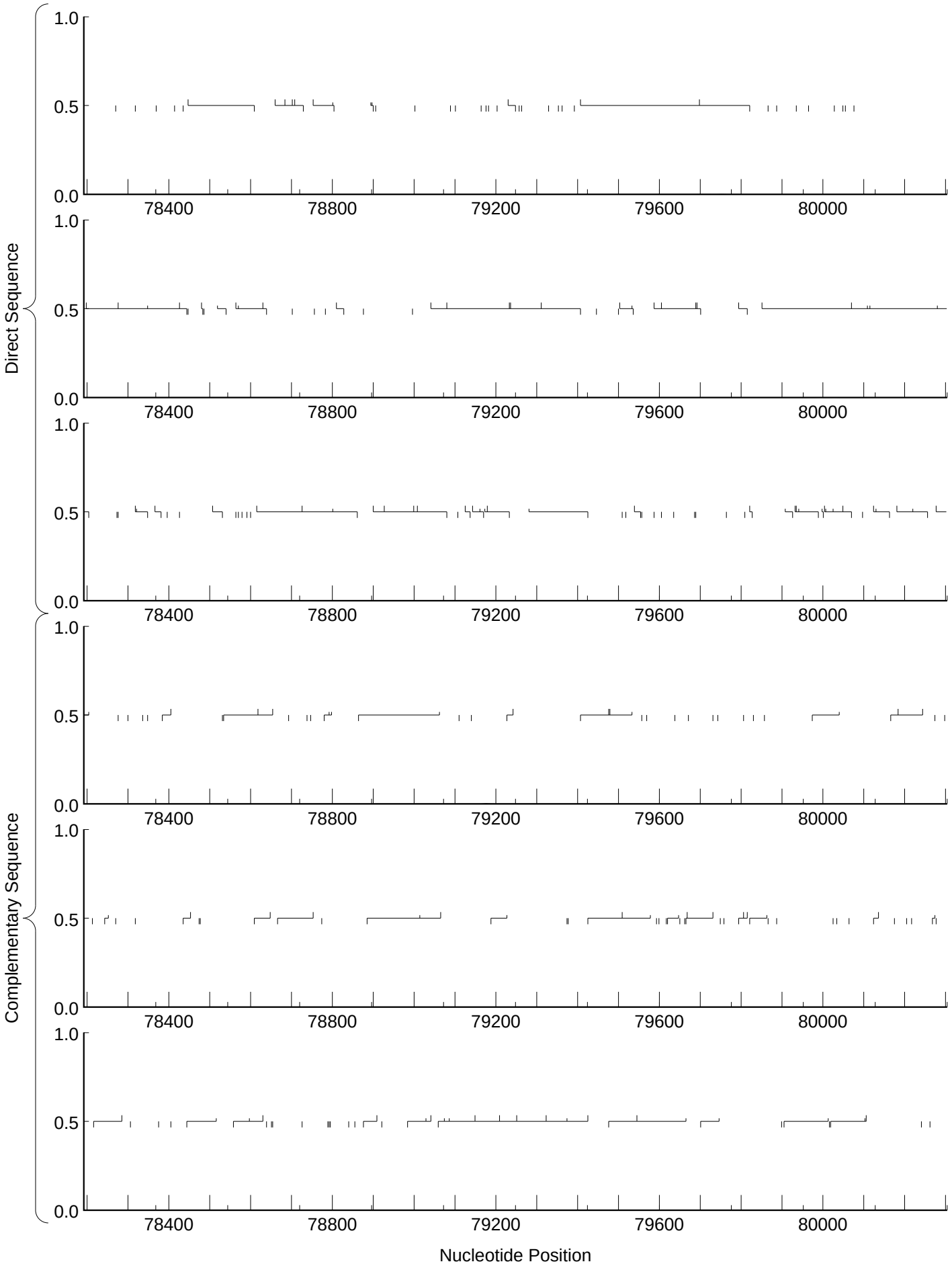
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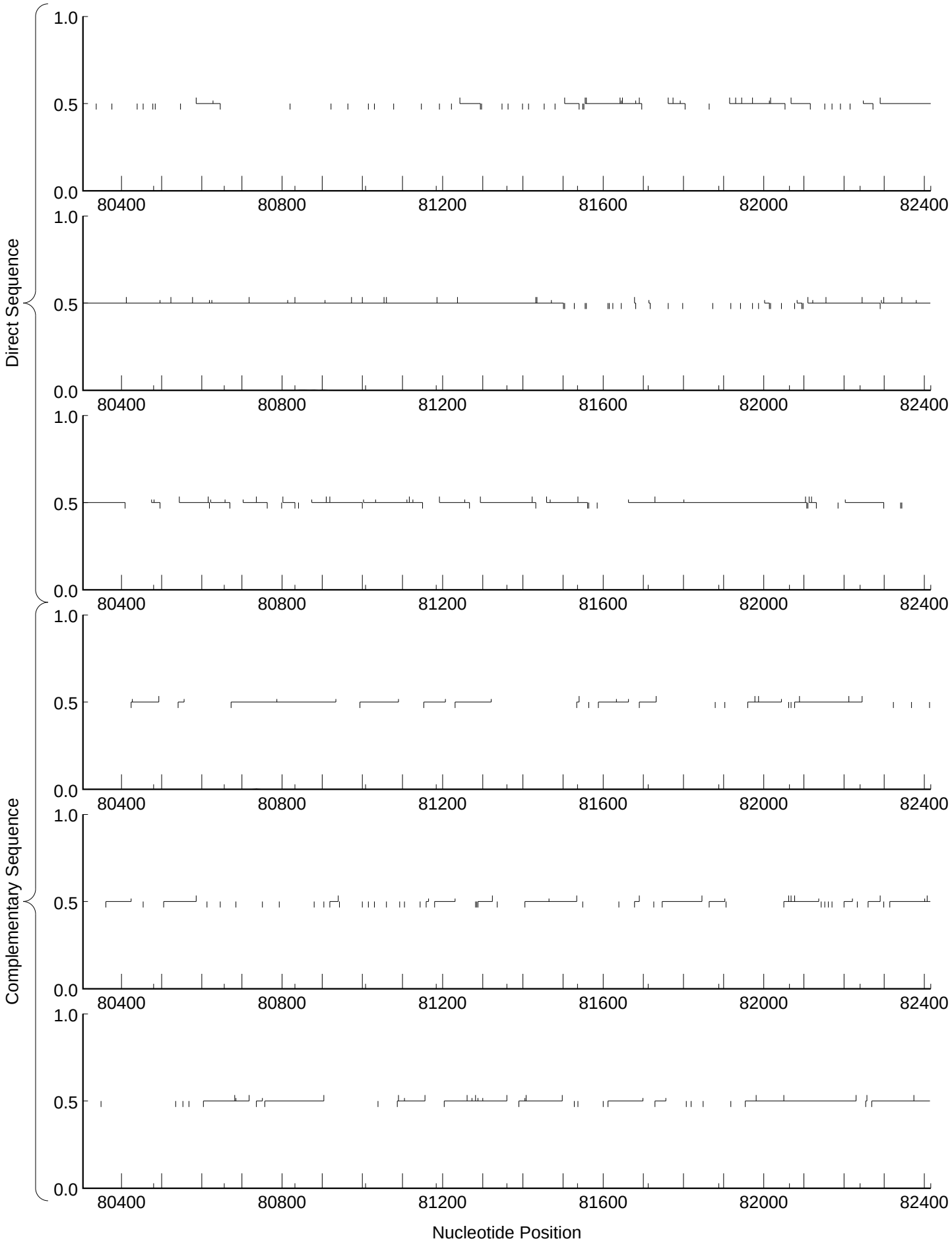
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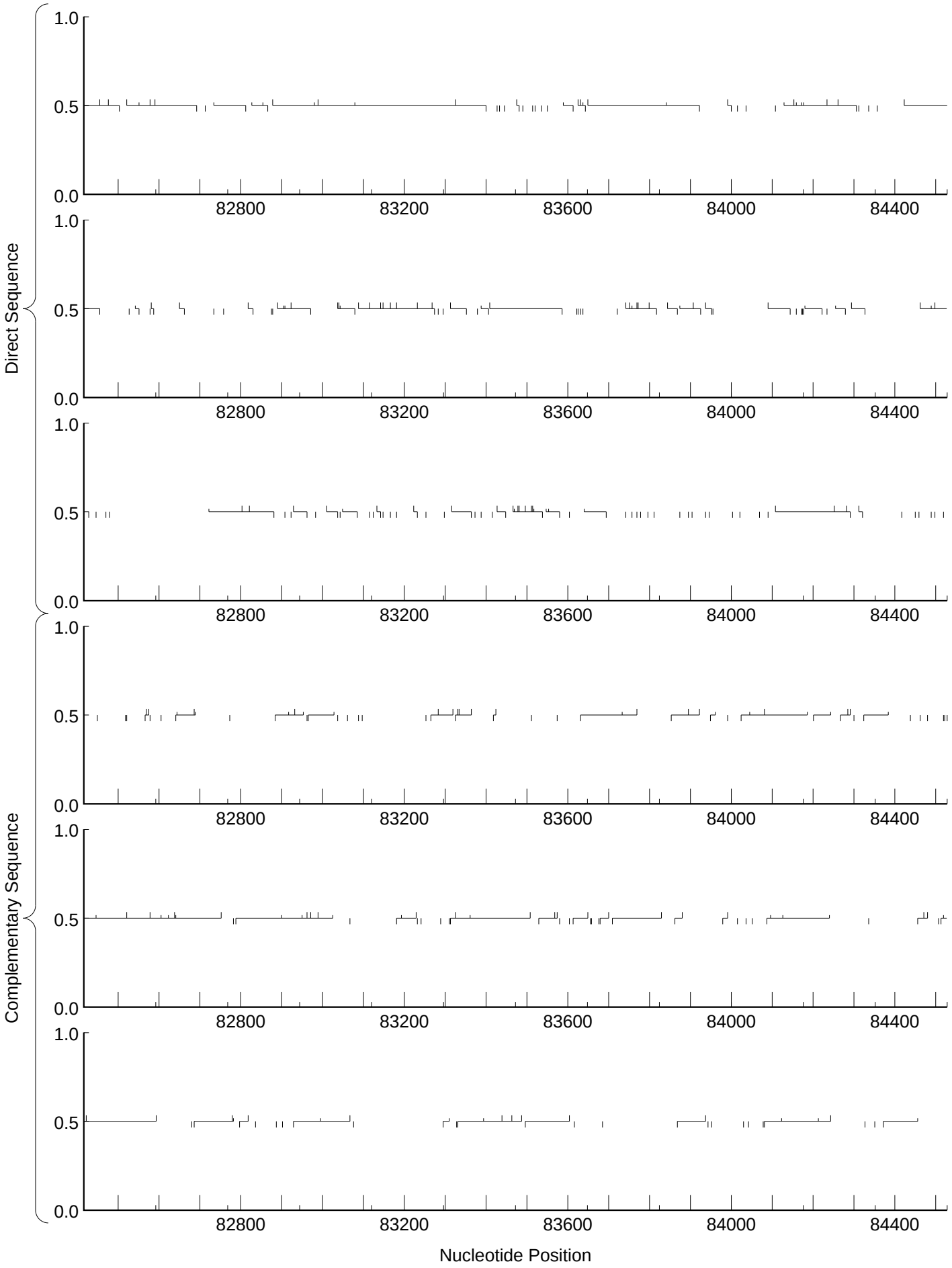
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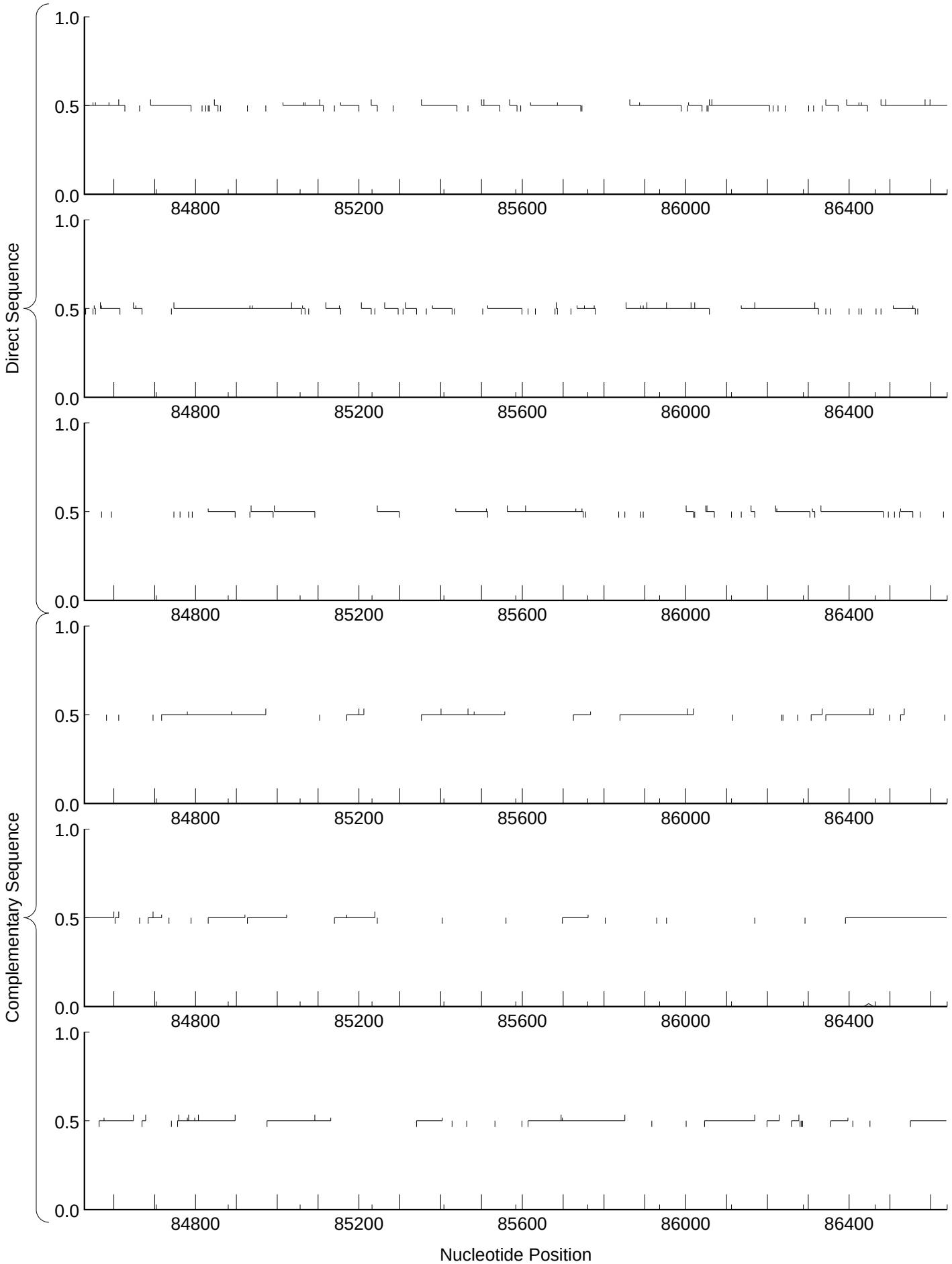
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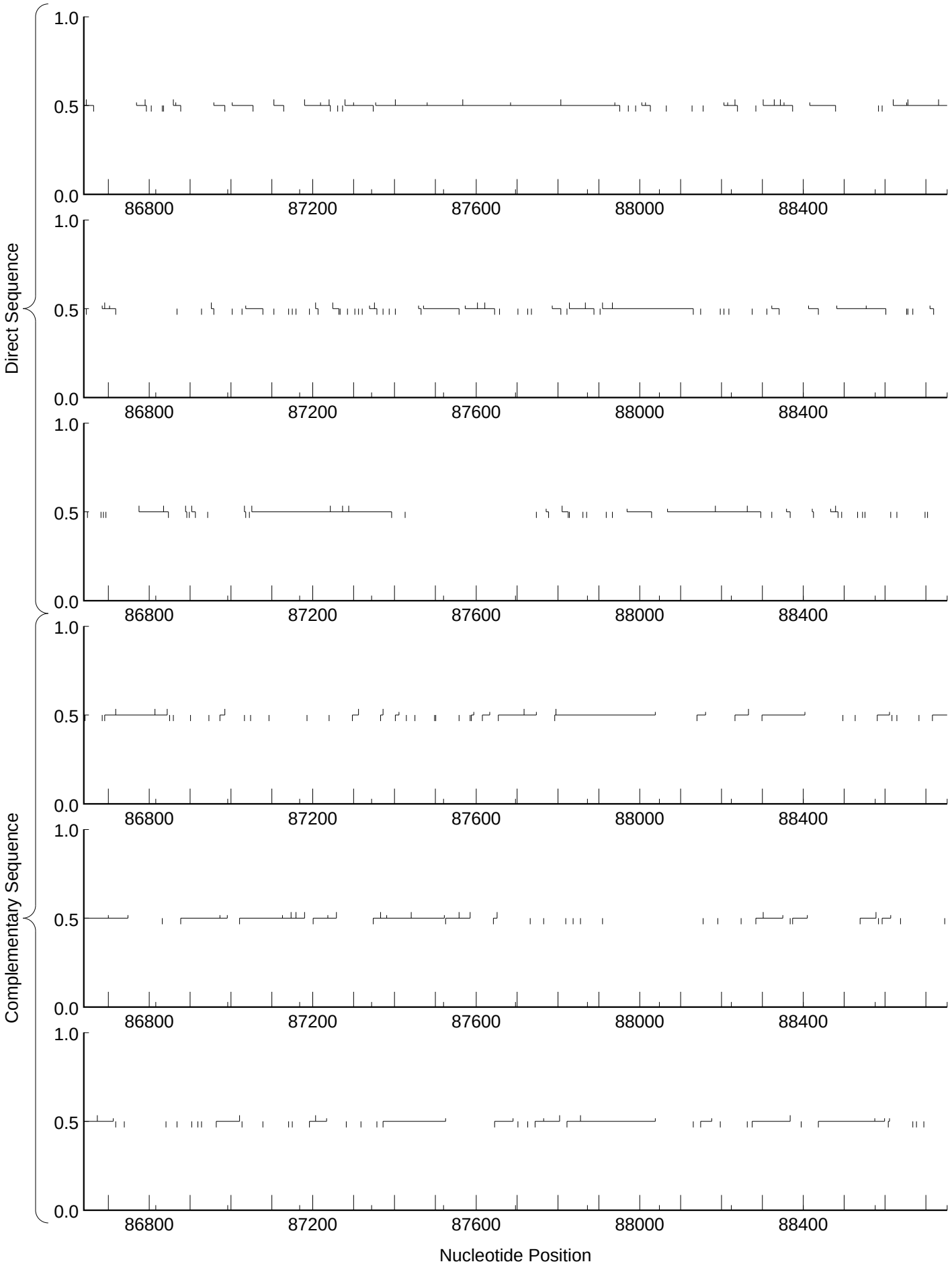
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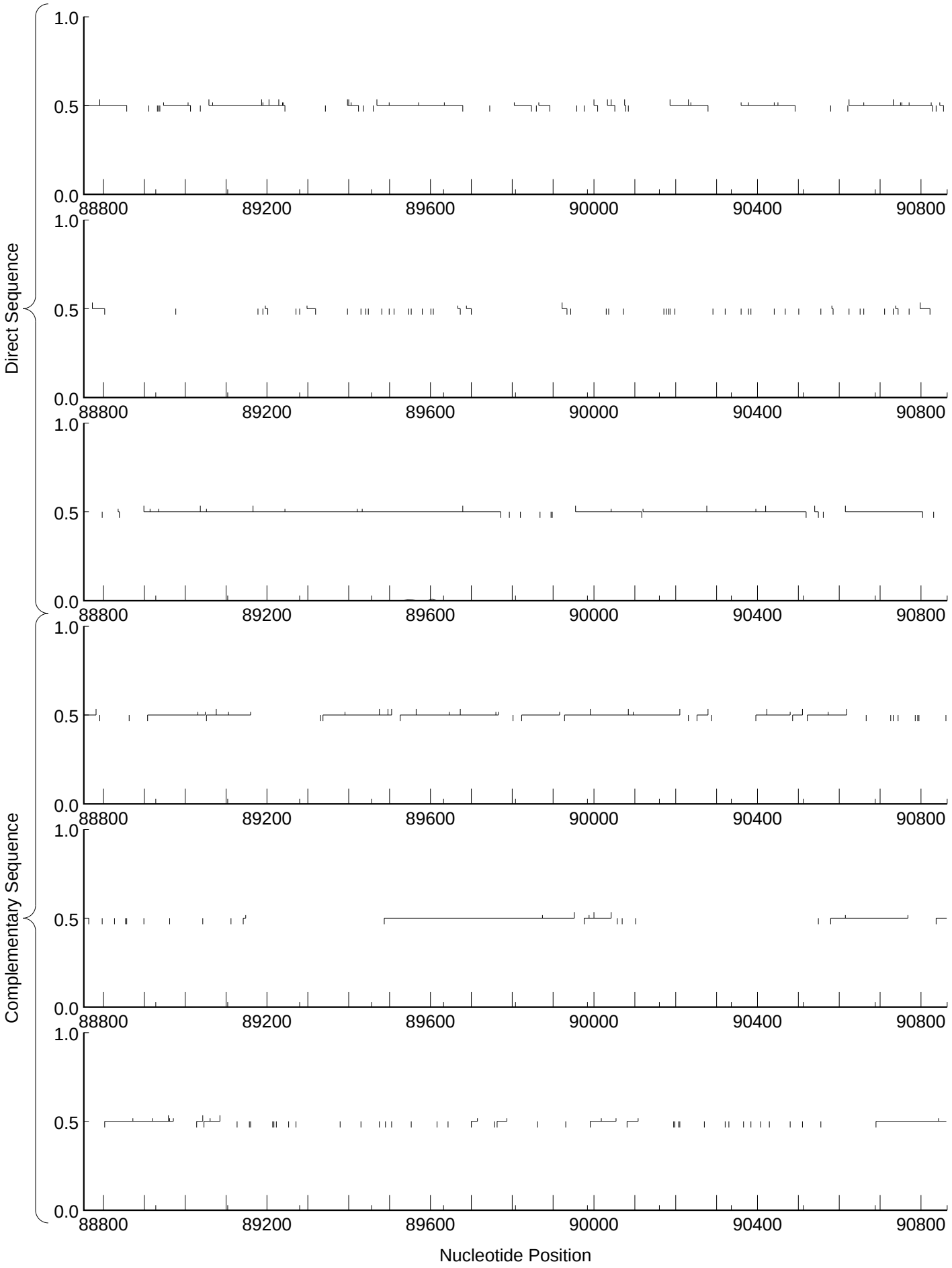
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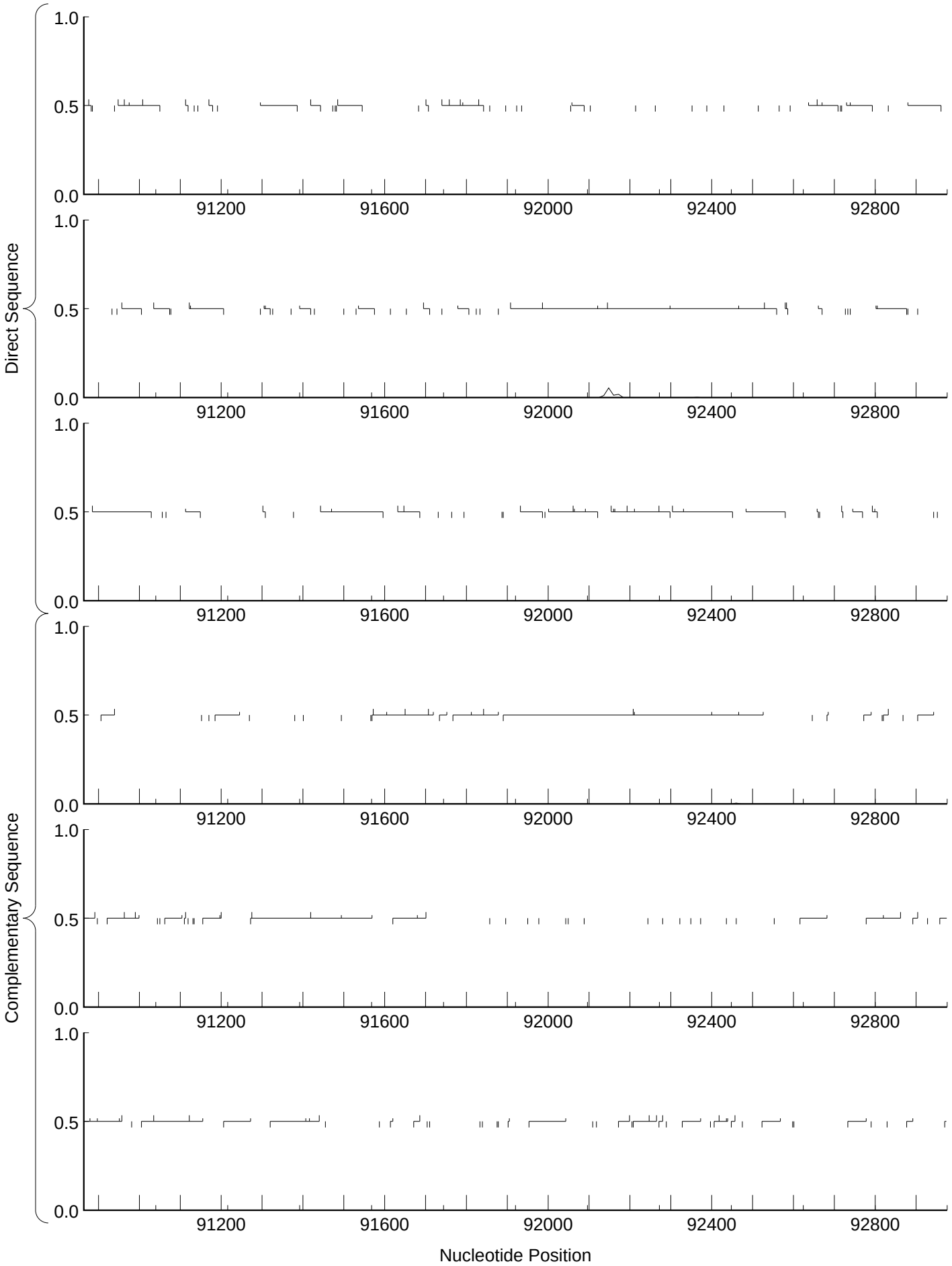
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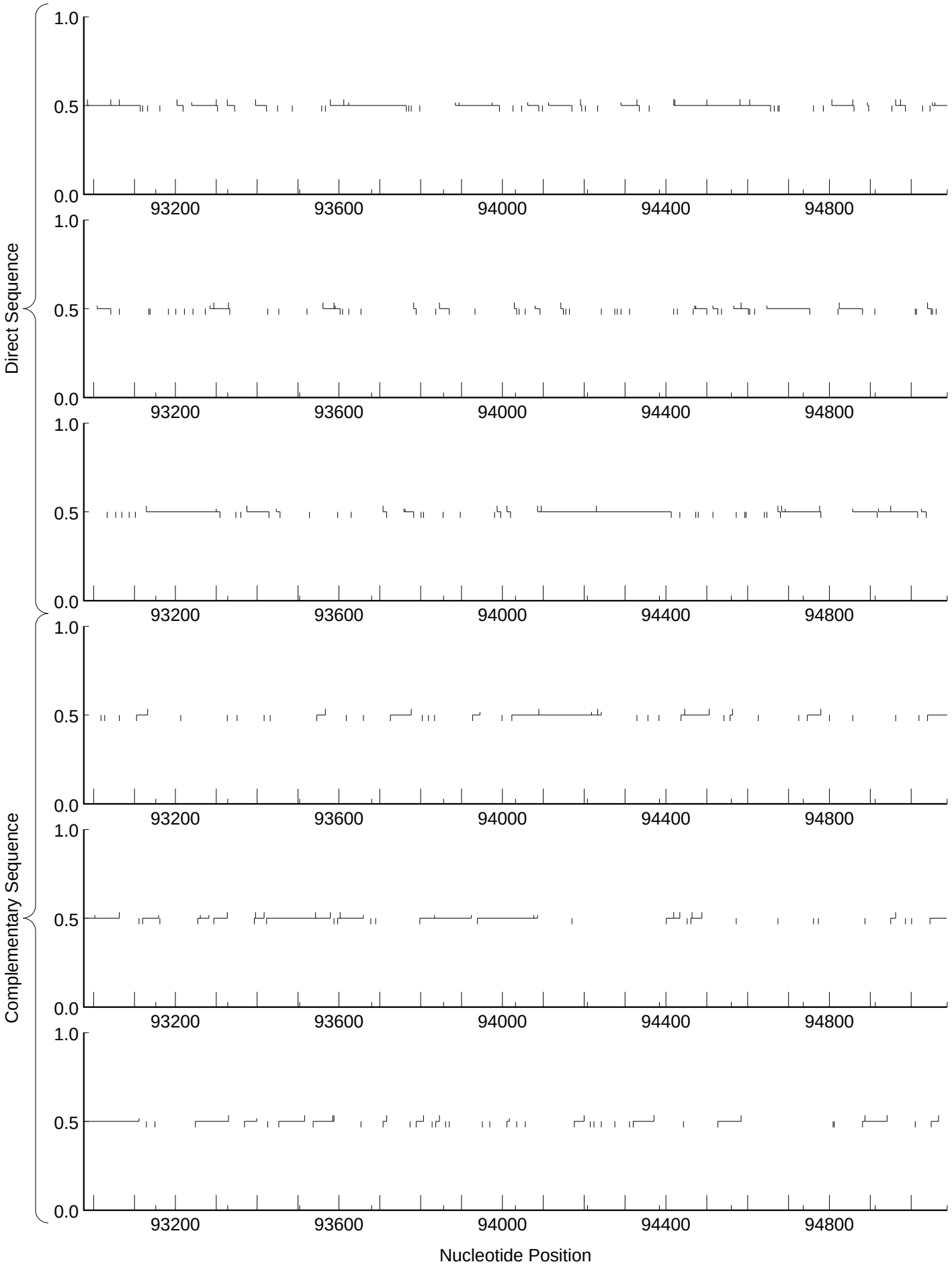
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 44/65



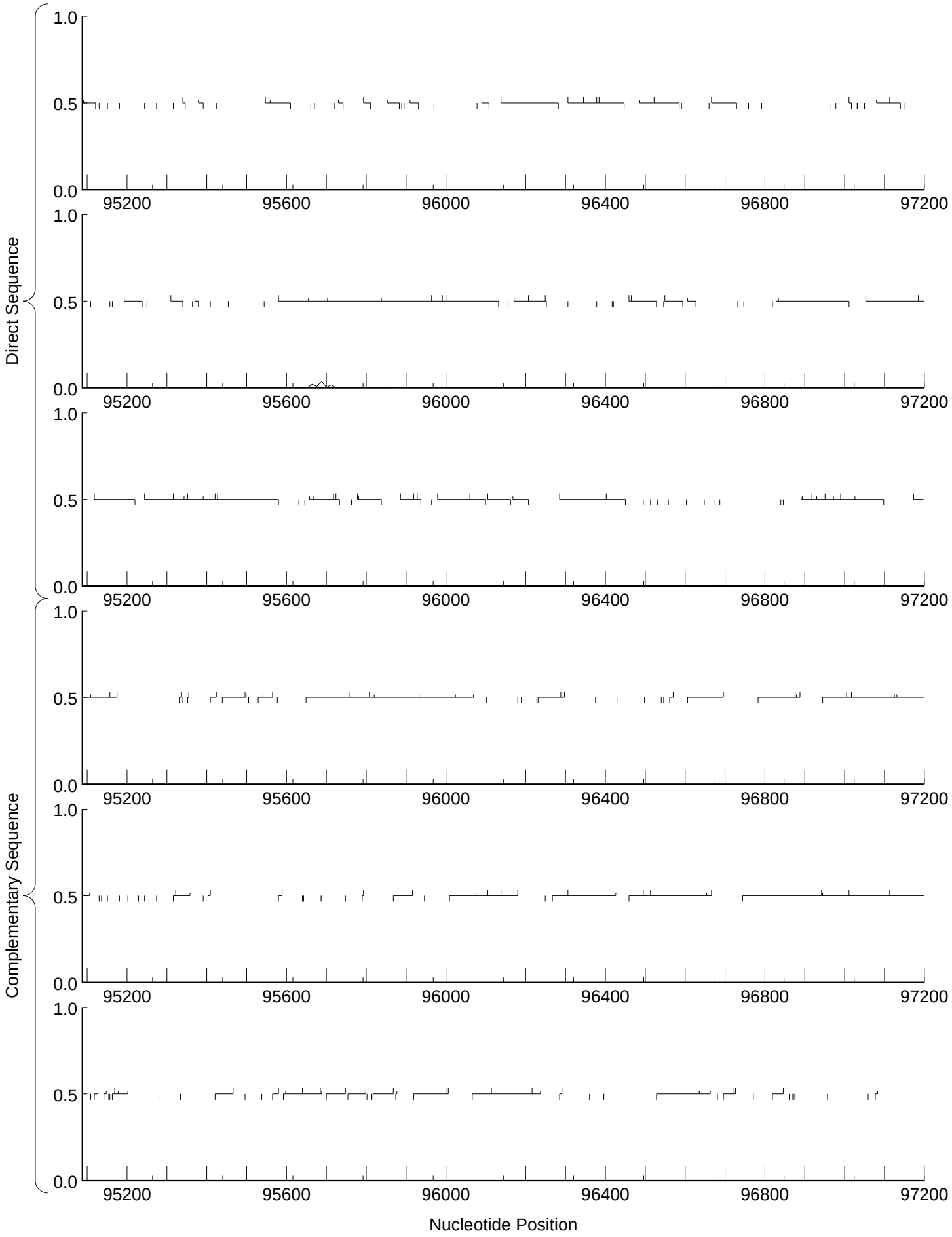
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 45/65



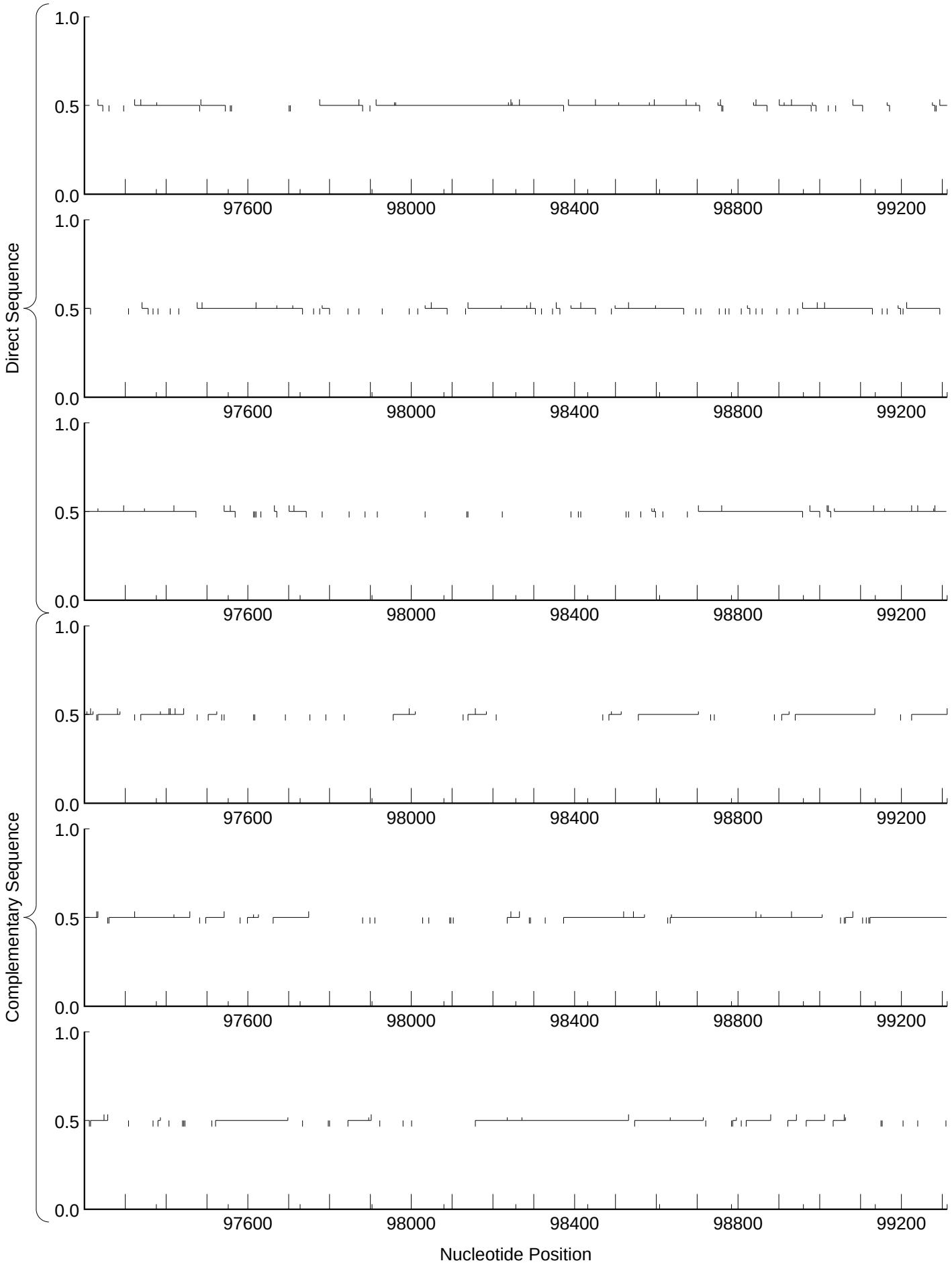
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 46/65



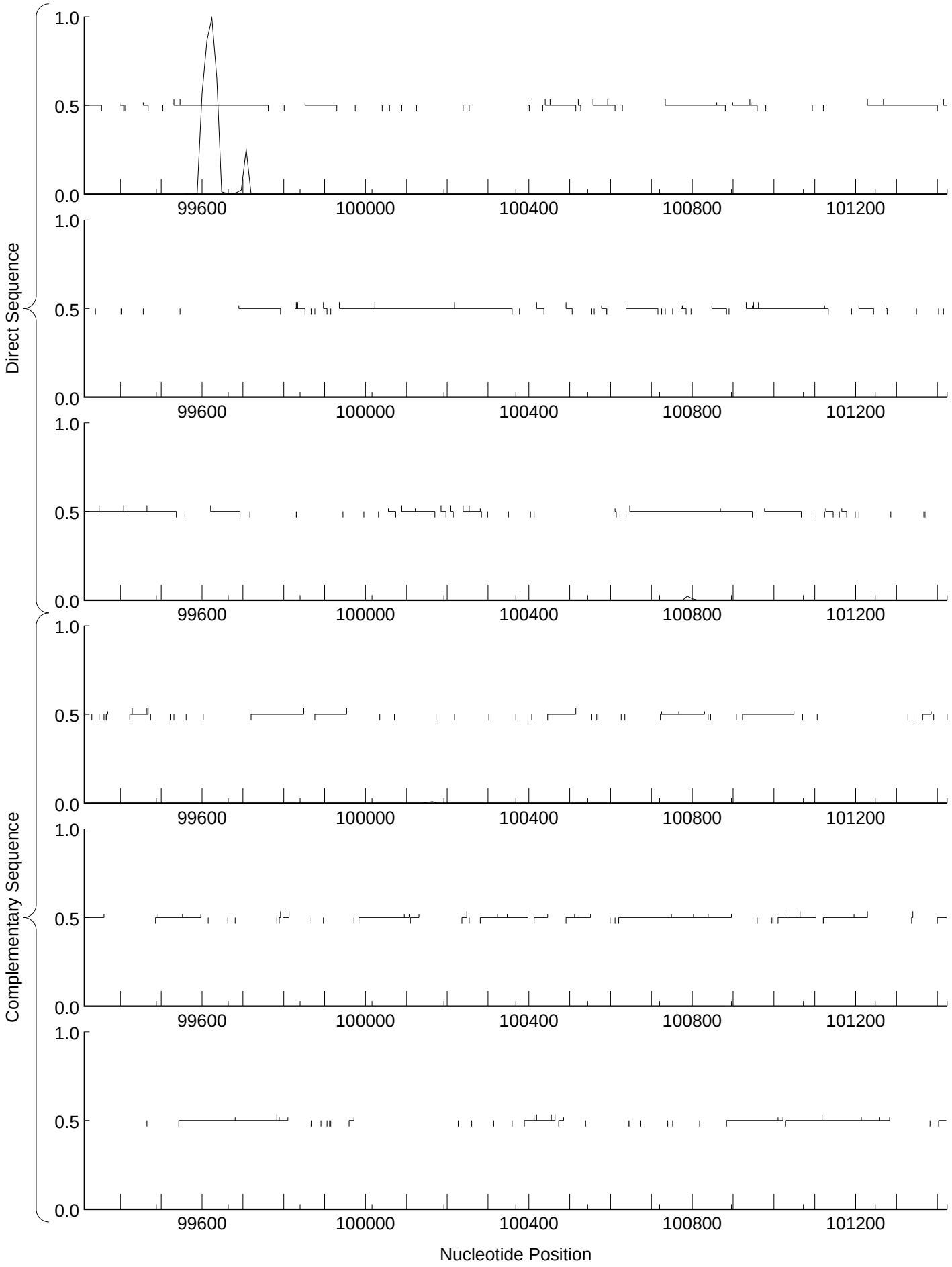
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 47/65



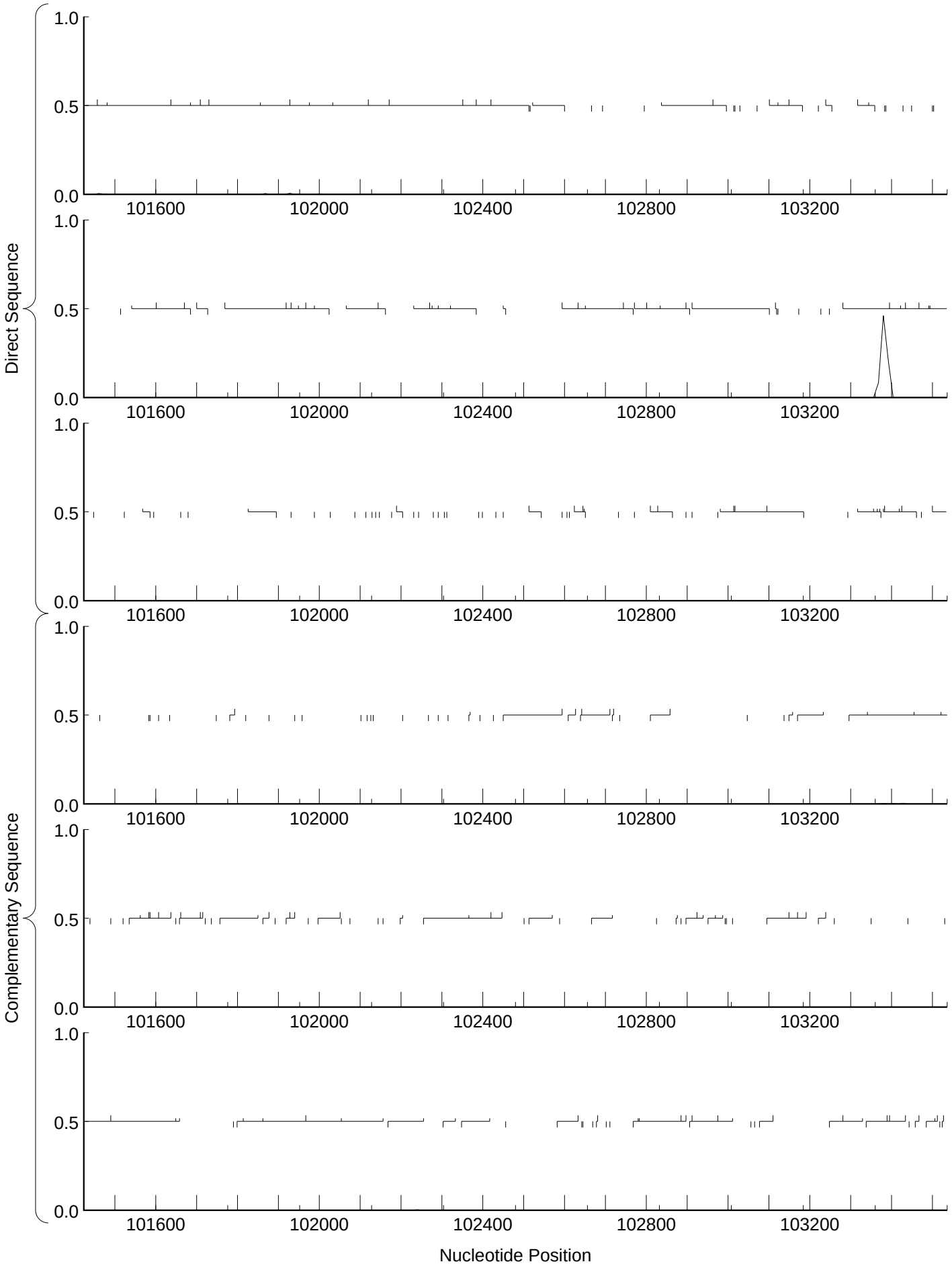
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 48/65



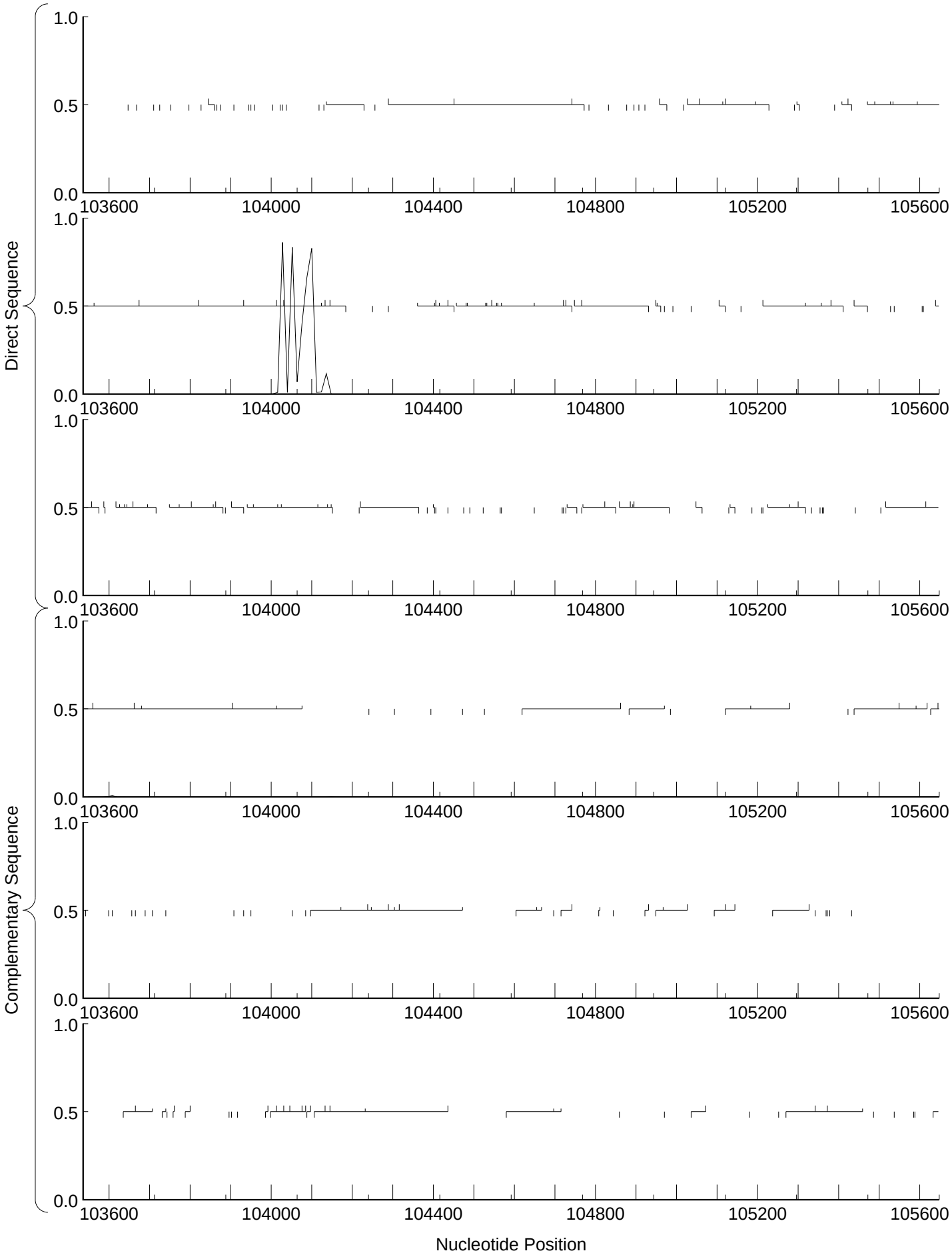
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 49/65



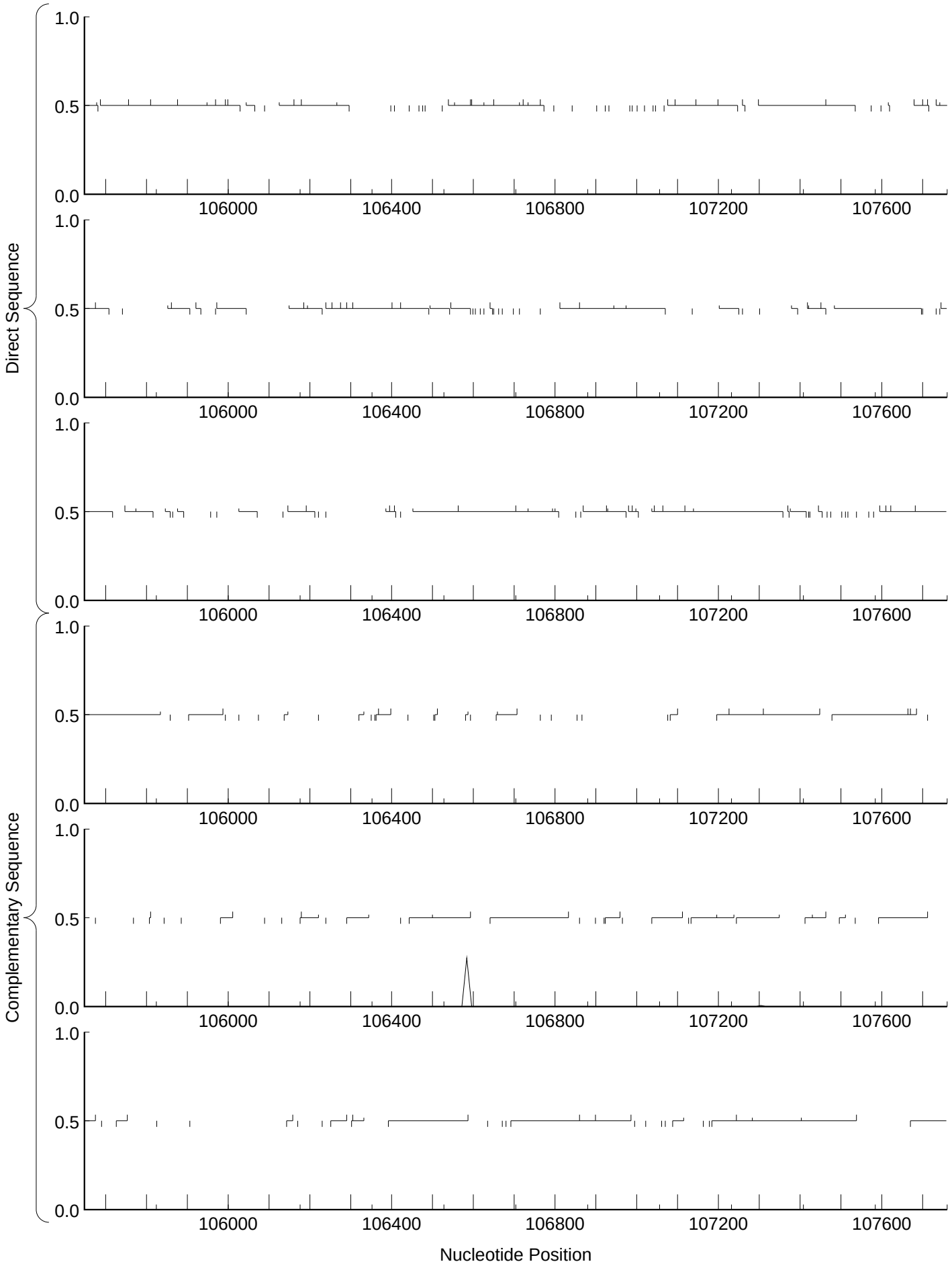
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 50/65



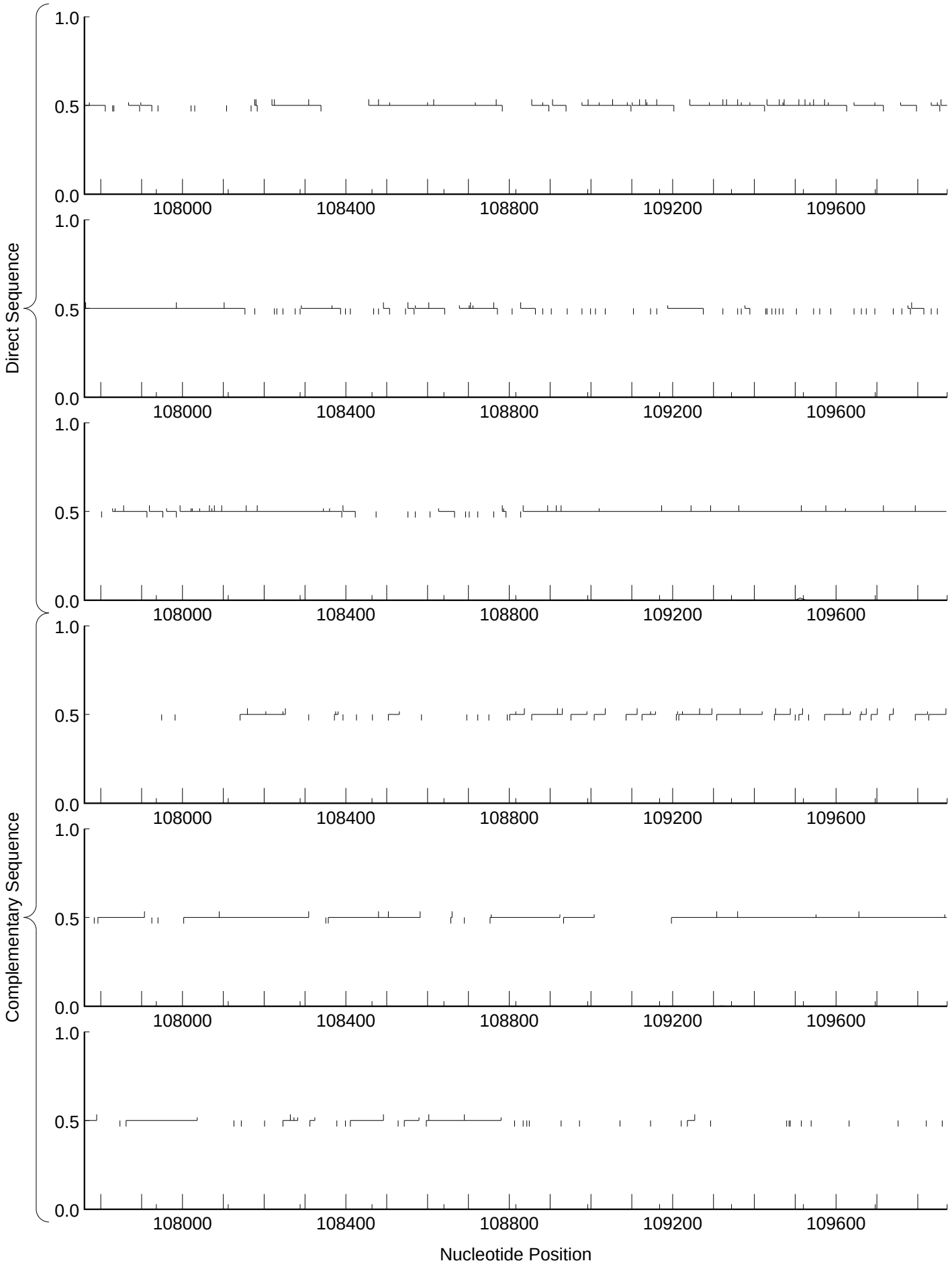
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 51/65



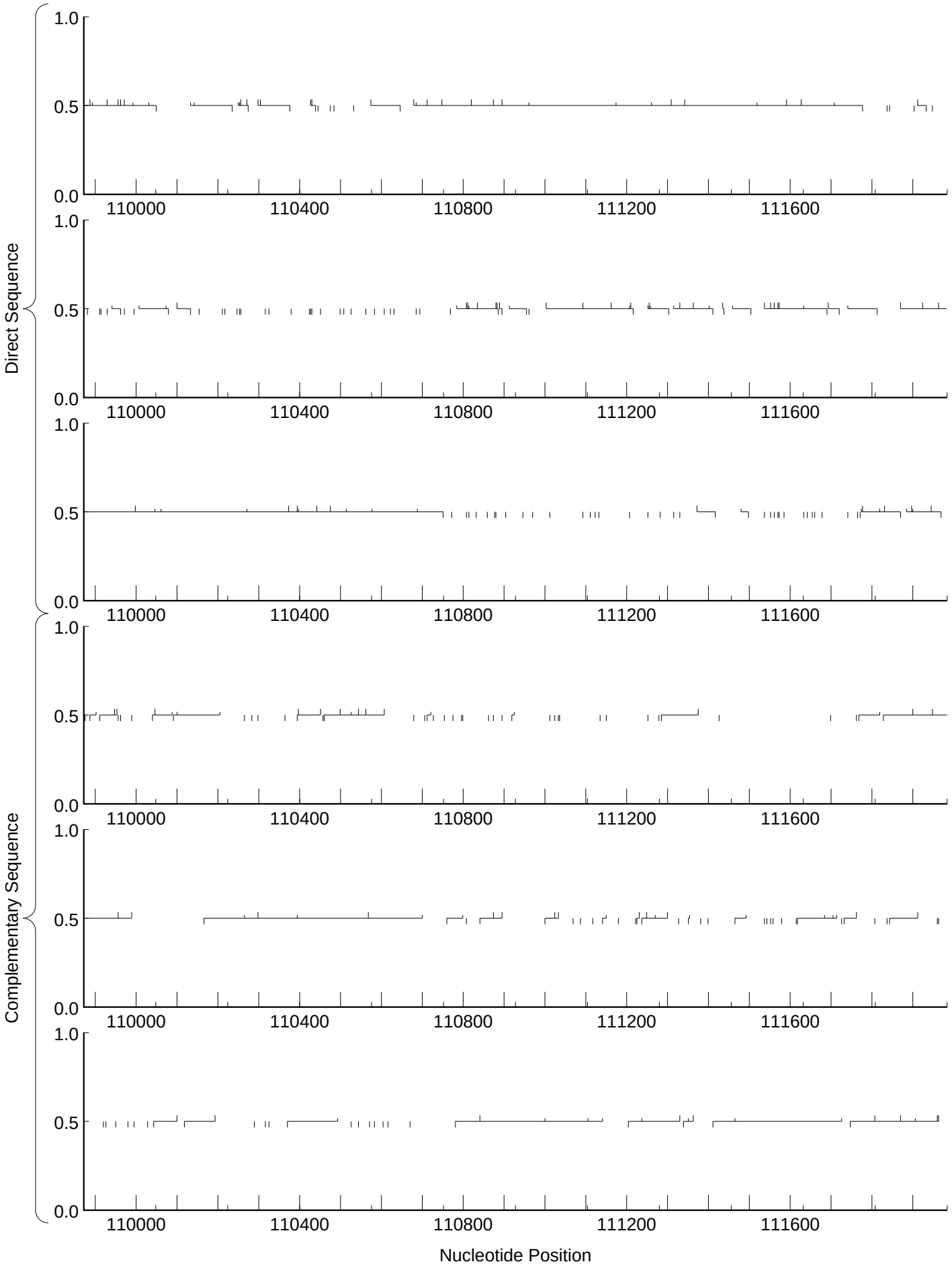
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 52/65



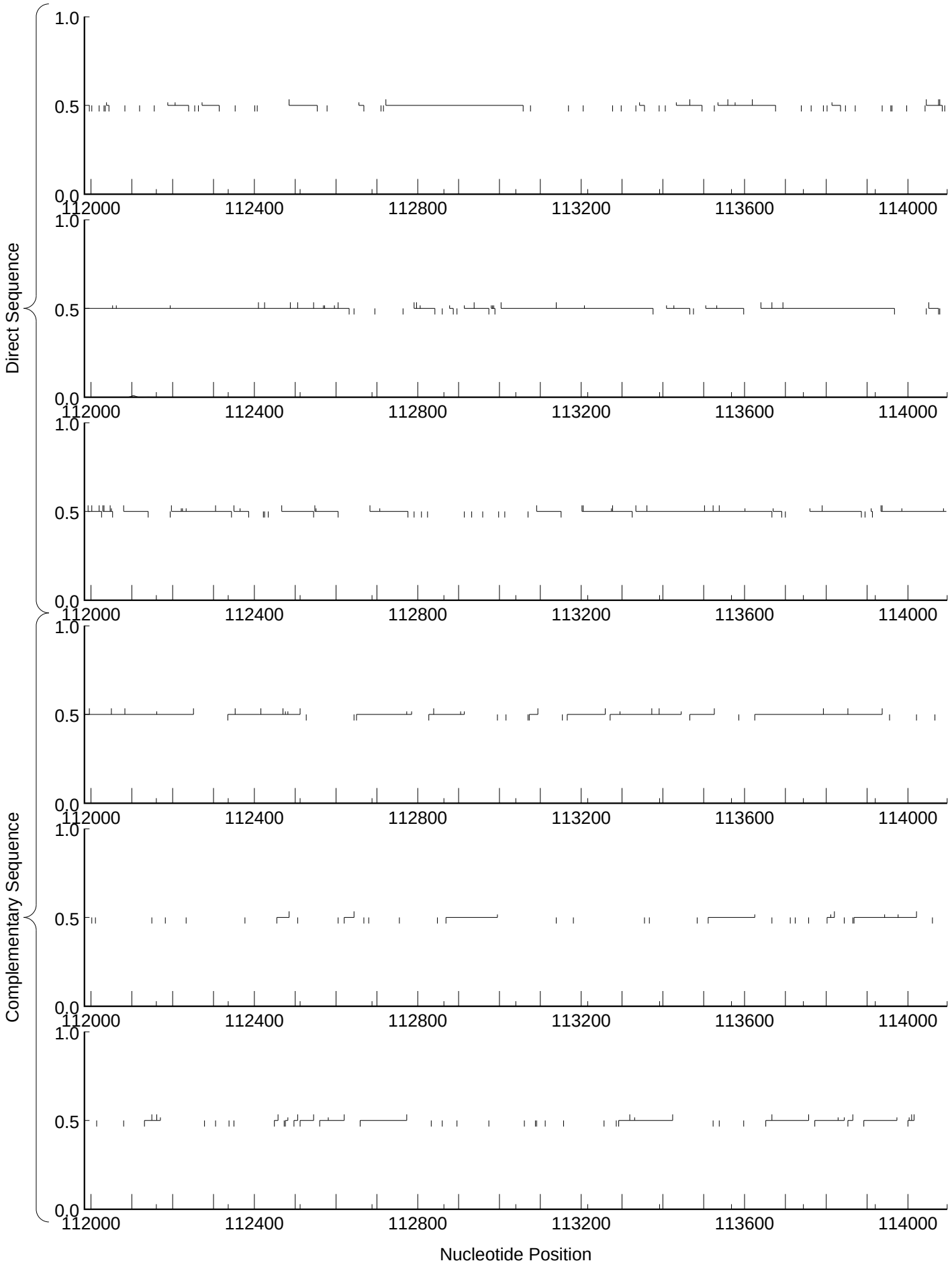
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 53/65



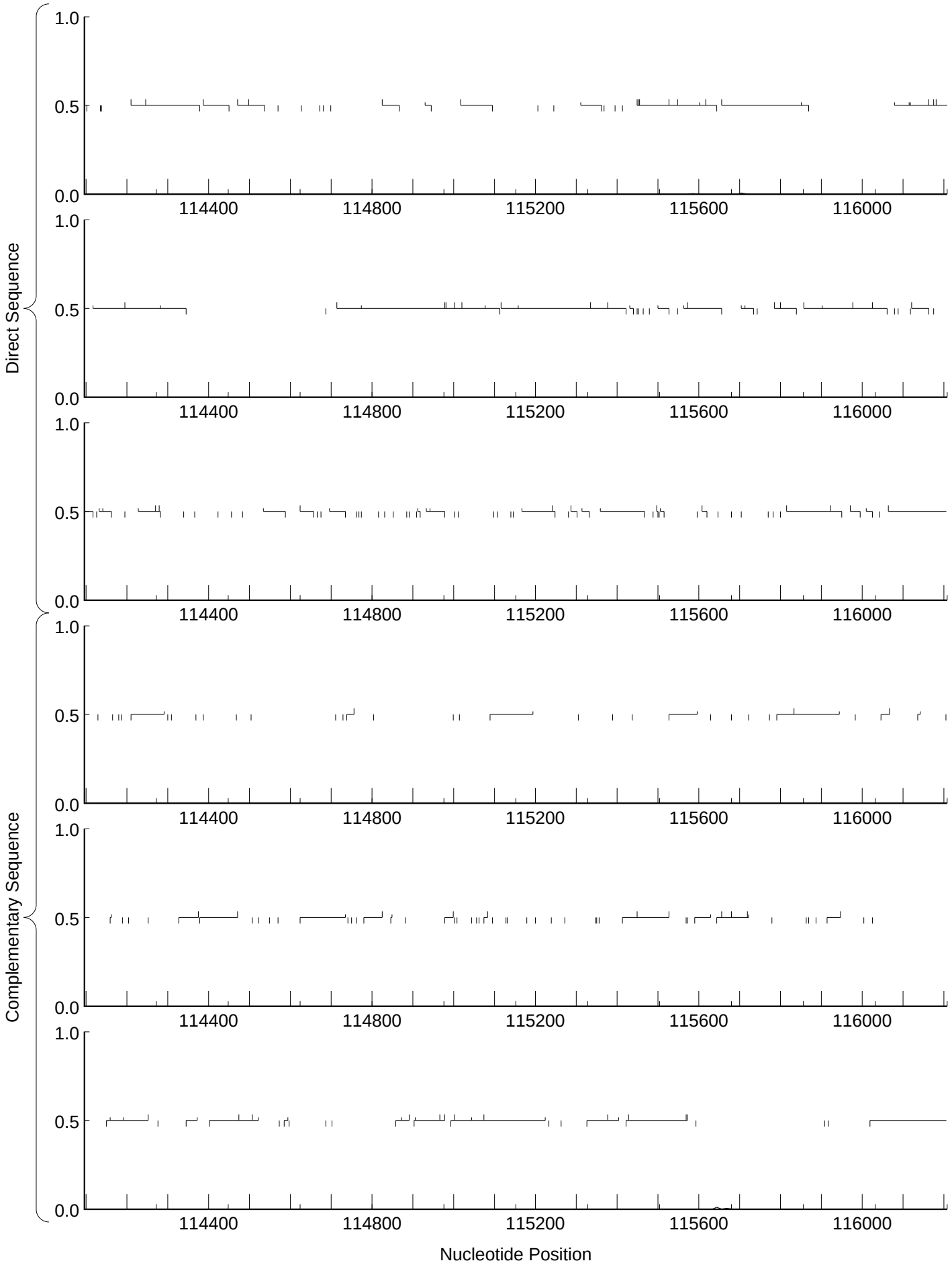
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 54/65



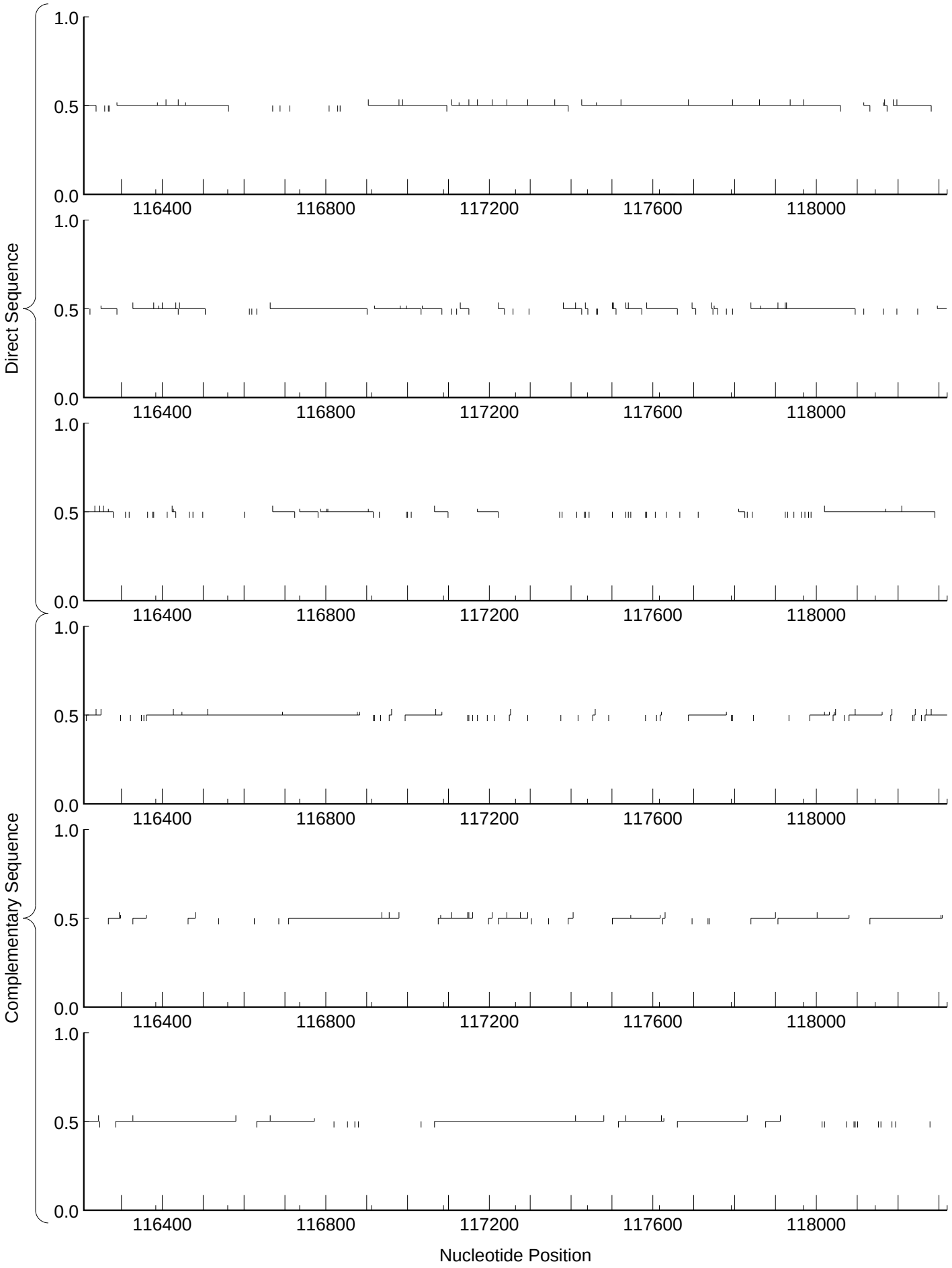
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 55/65



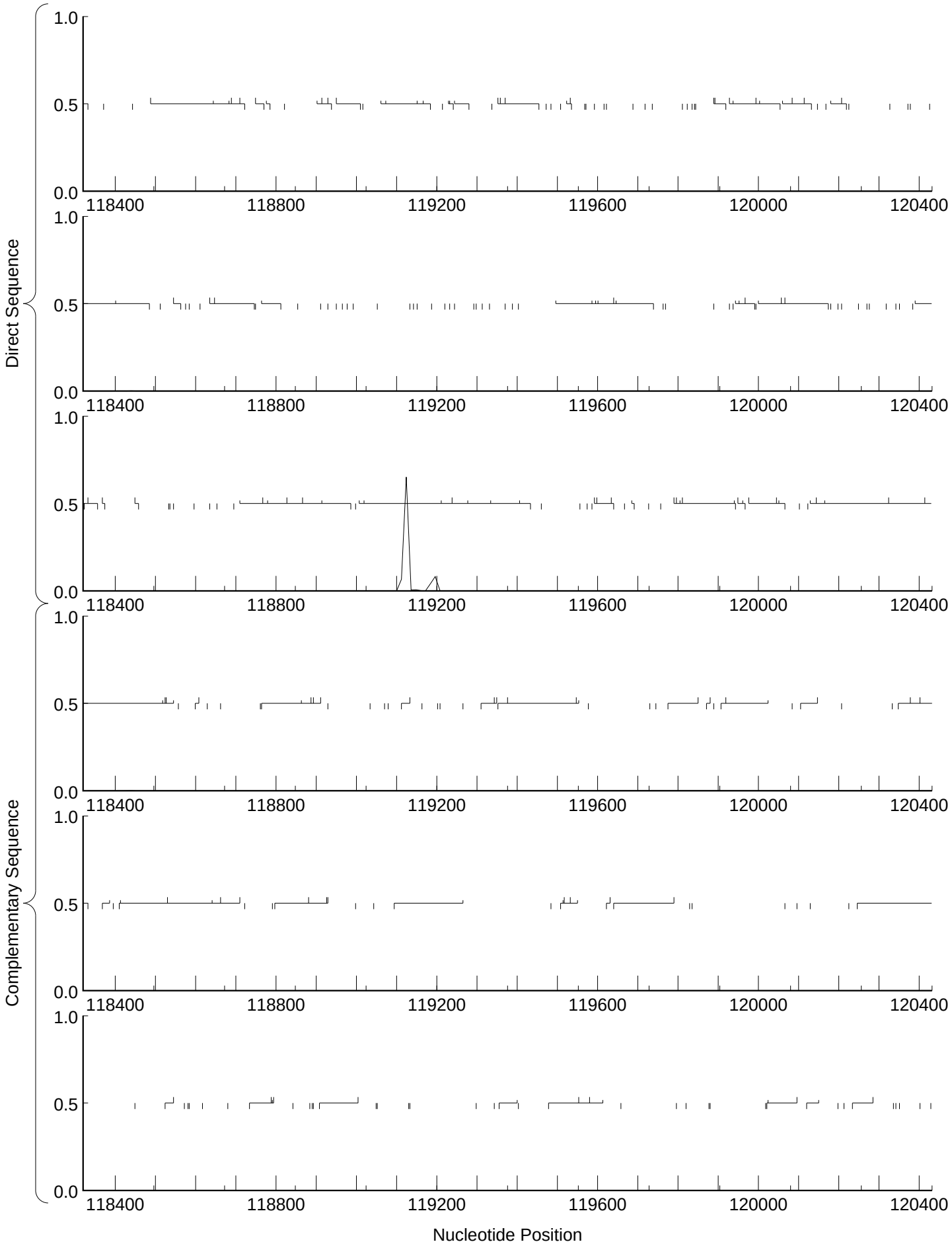
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 56/65



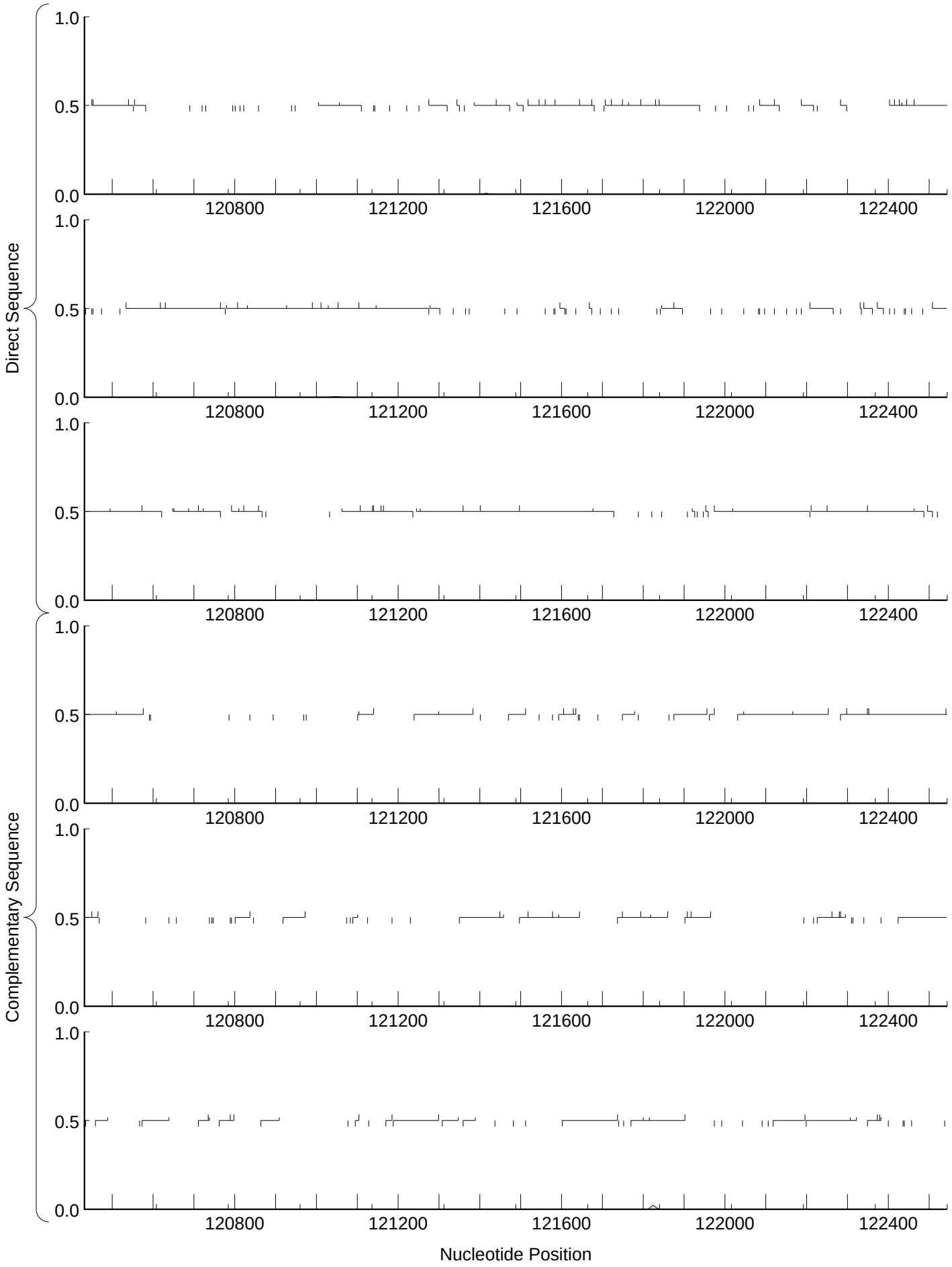
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 57/65



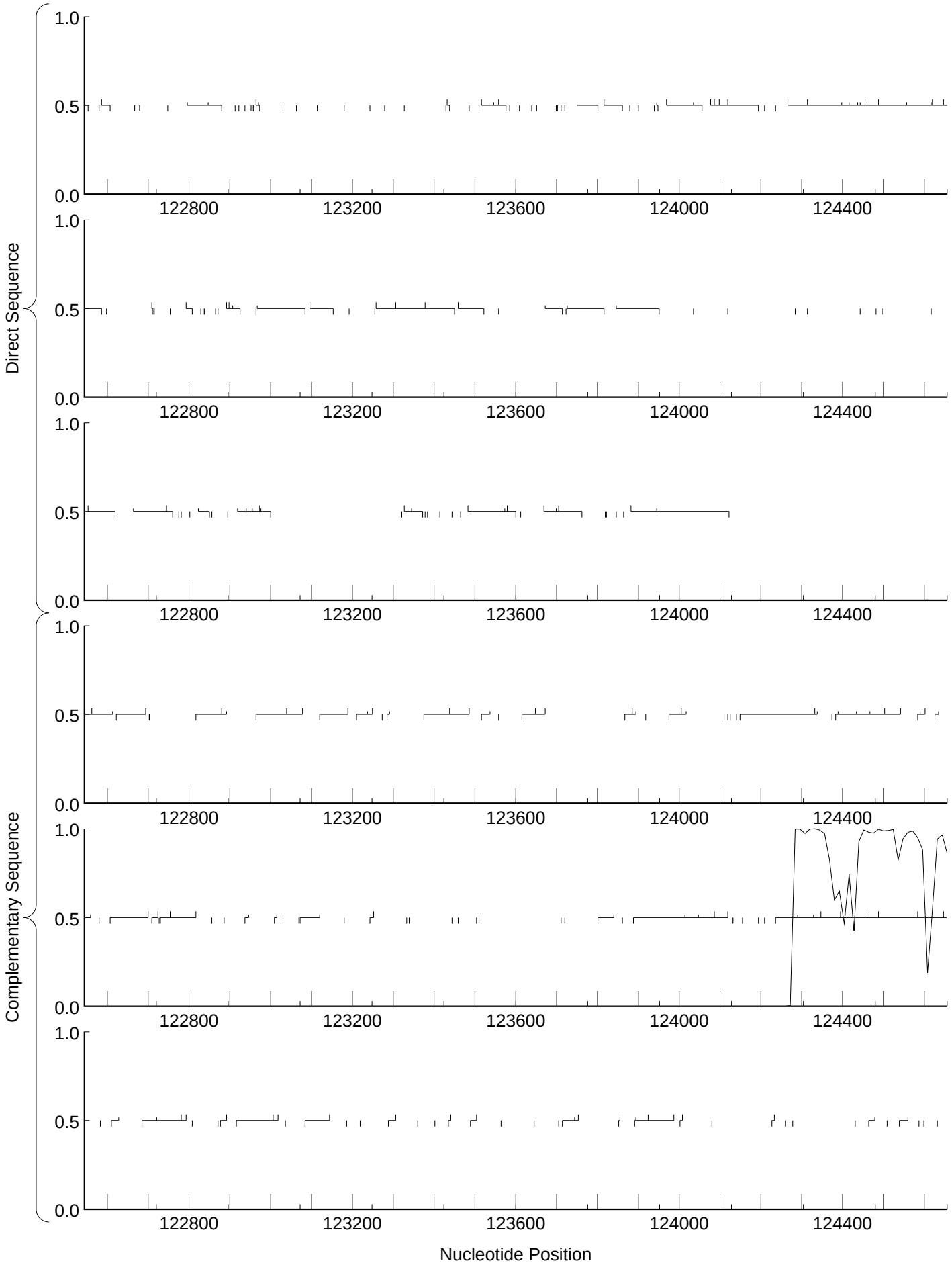
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 58/65



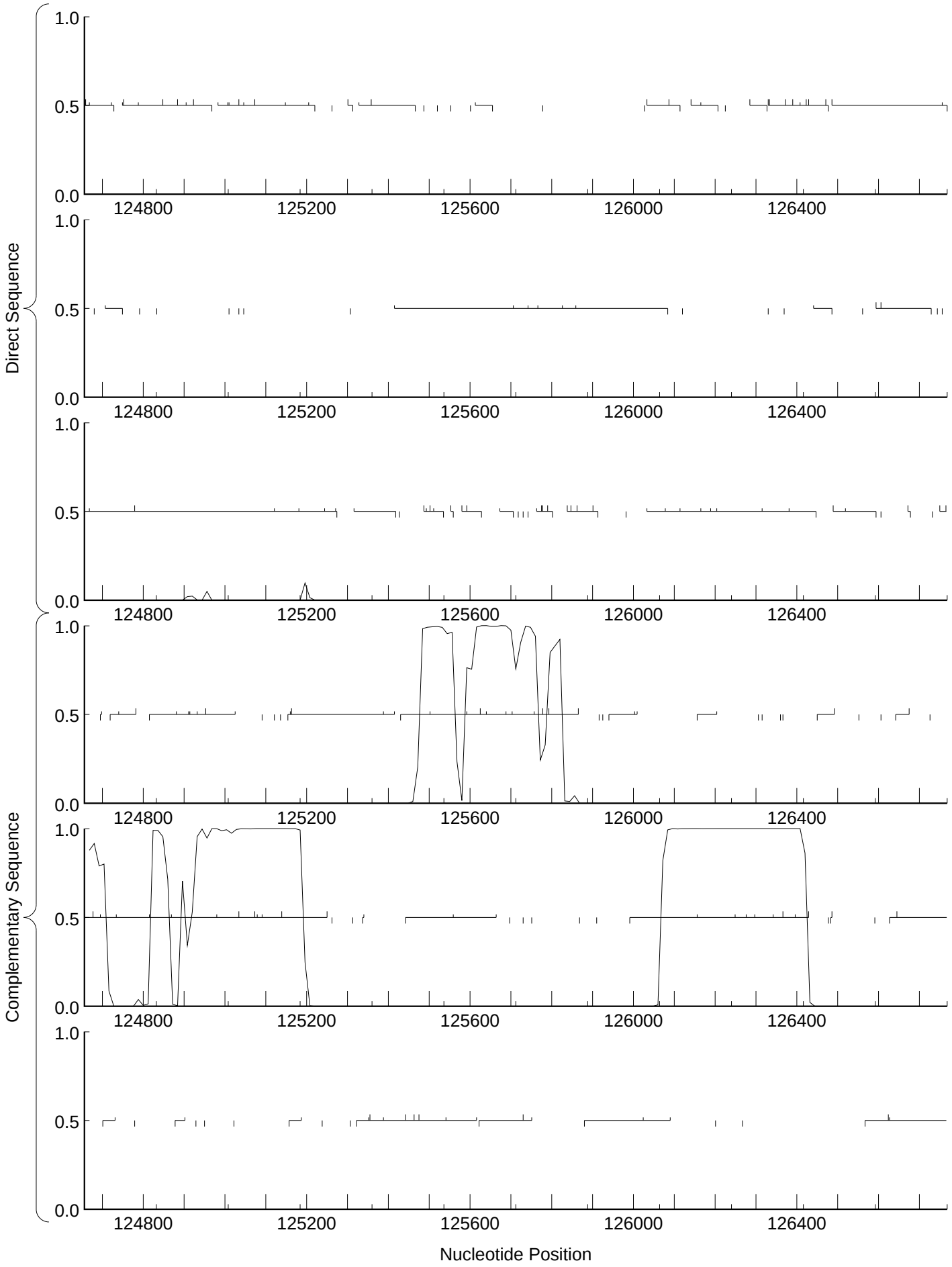
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 59/65



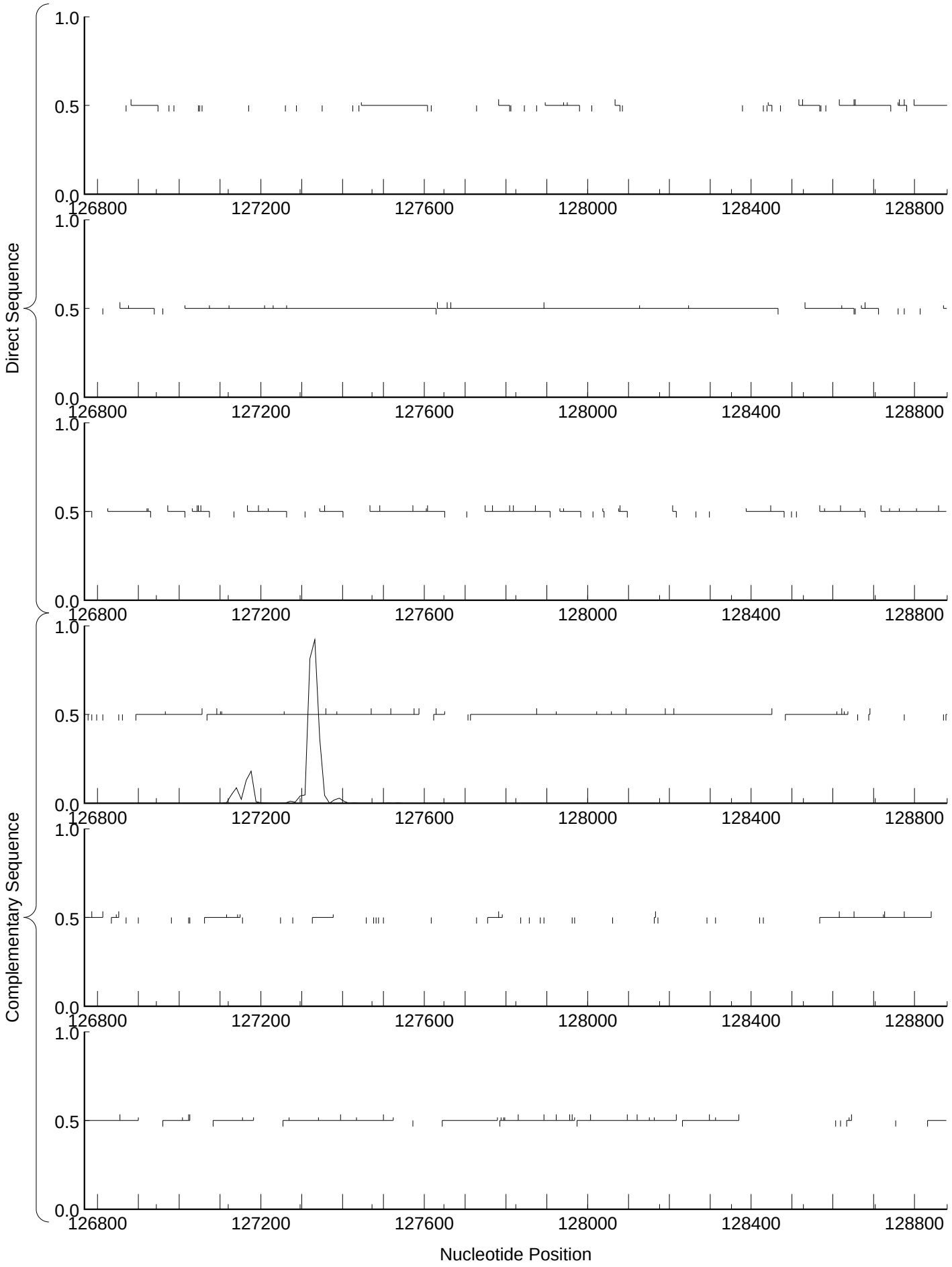
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 60/65



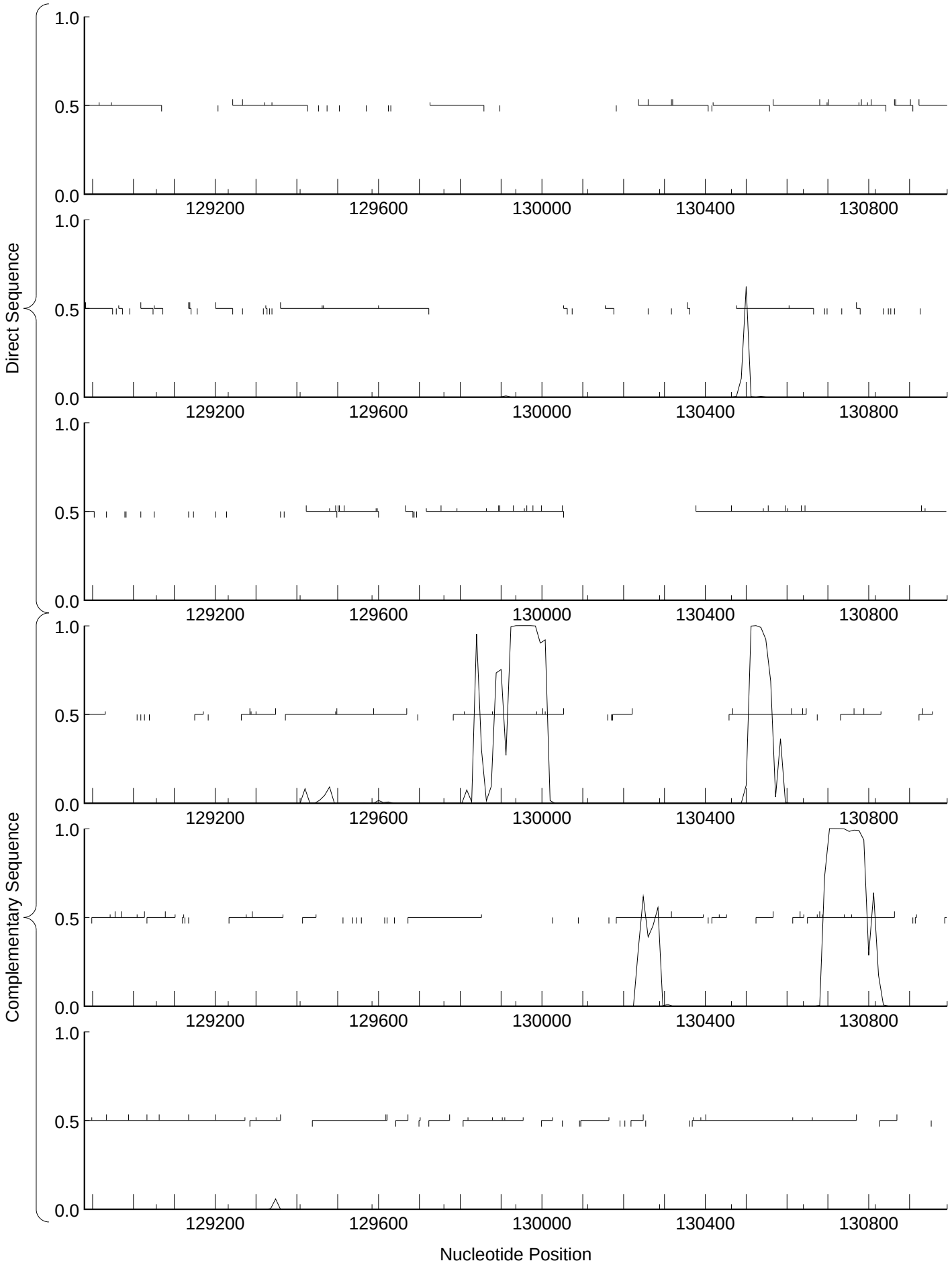
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 61/65



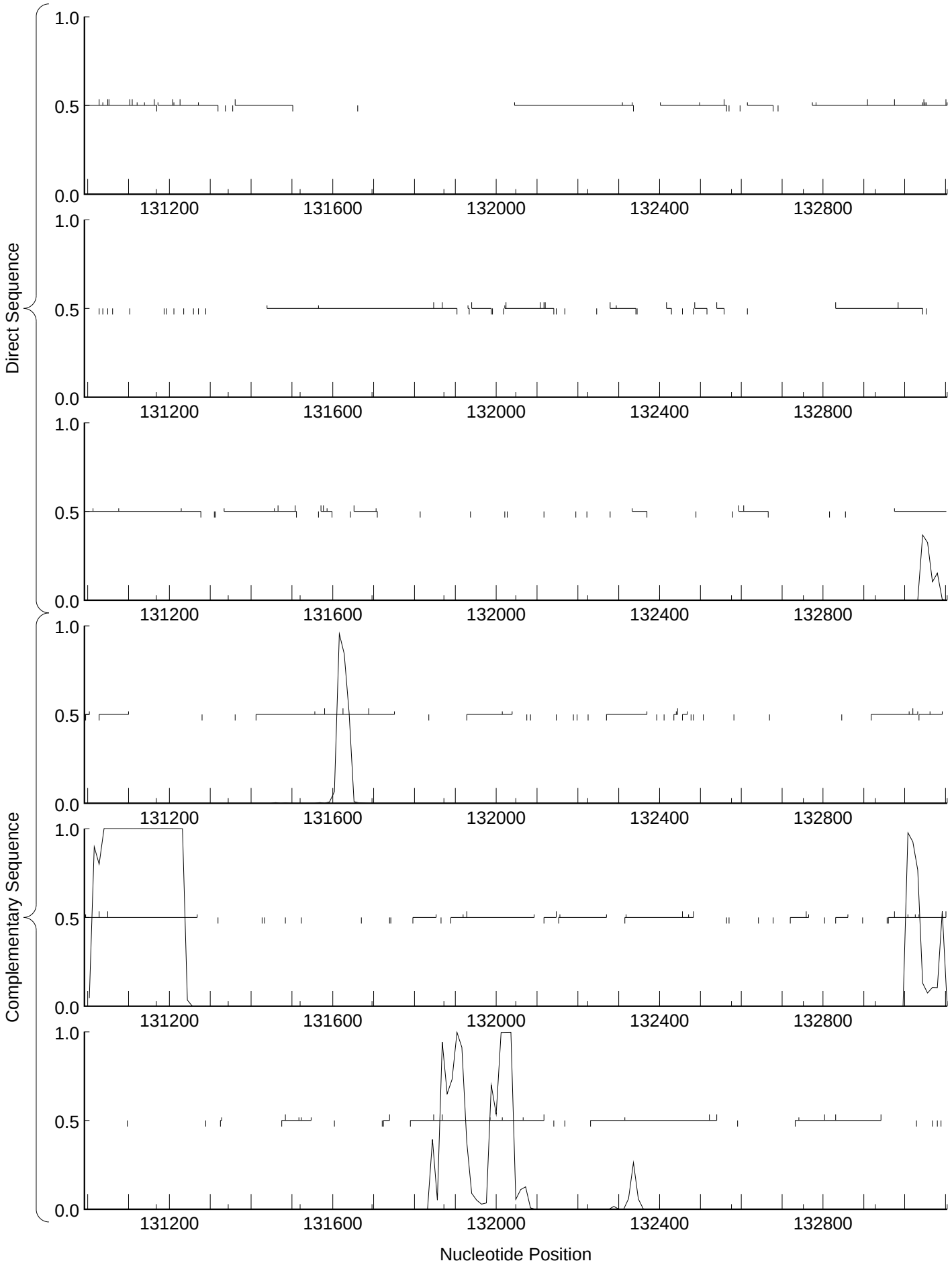
chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 62/65



chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 63/65



chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 64/65



chMansion complete sequence, 134105 bp including 11127 bp terminal repeat, Cluster BE1, Order 4, Window 96, Step 12, 65/65

