GeneMark

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

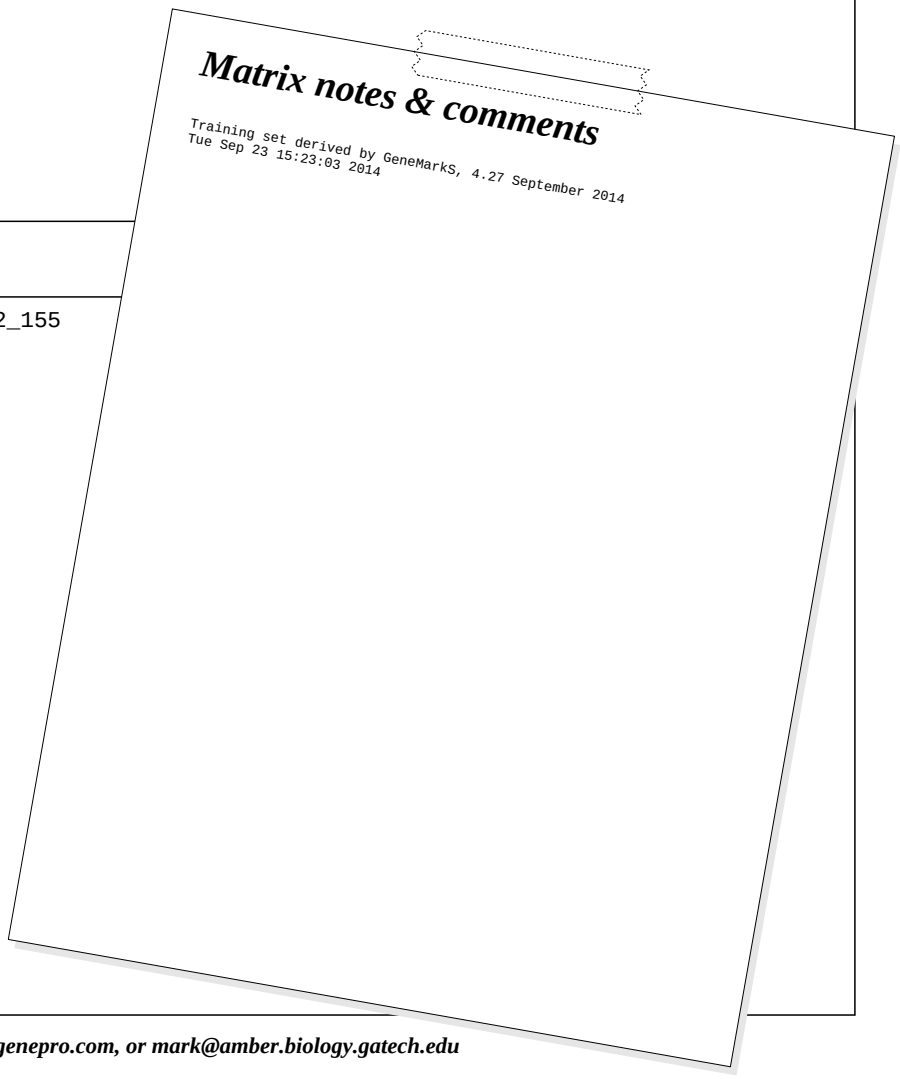
PROGRAM INFORMATION

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Analysis Date    :  3/28/24 at 9:38:24
Pages            :  76
Sequence Length  :  156308 bp
GC Content       :   64.69%

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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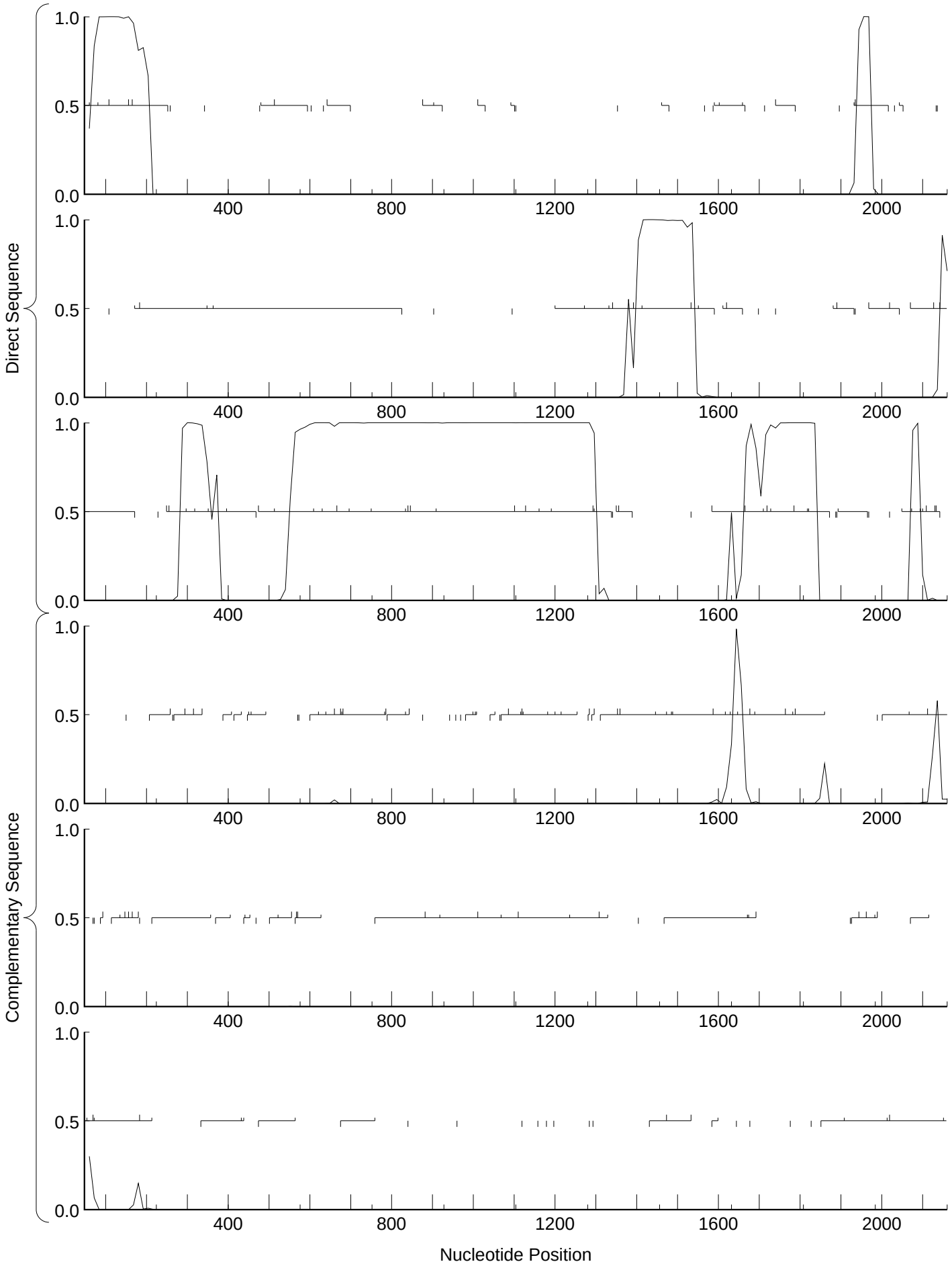
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Order      :  4
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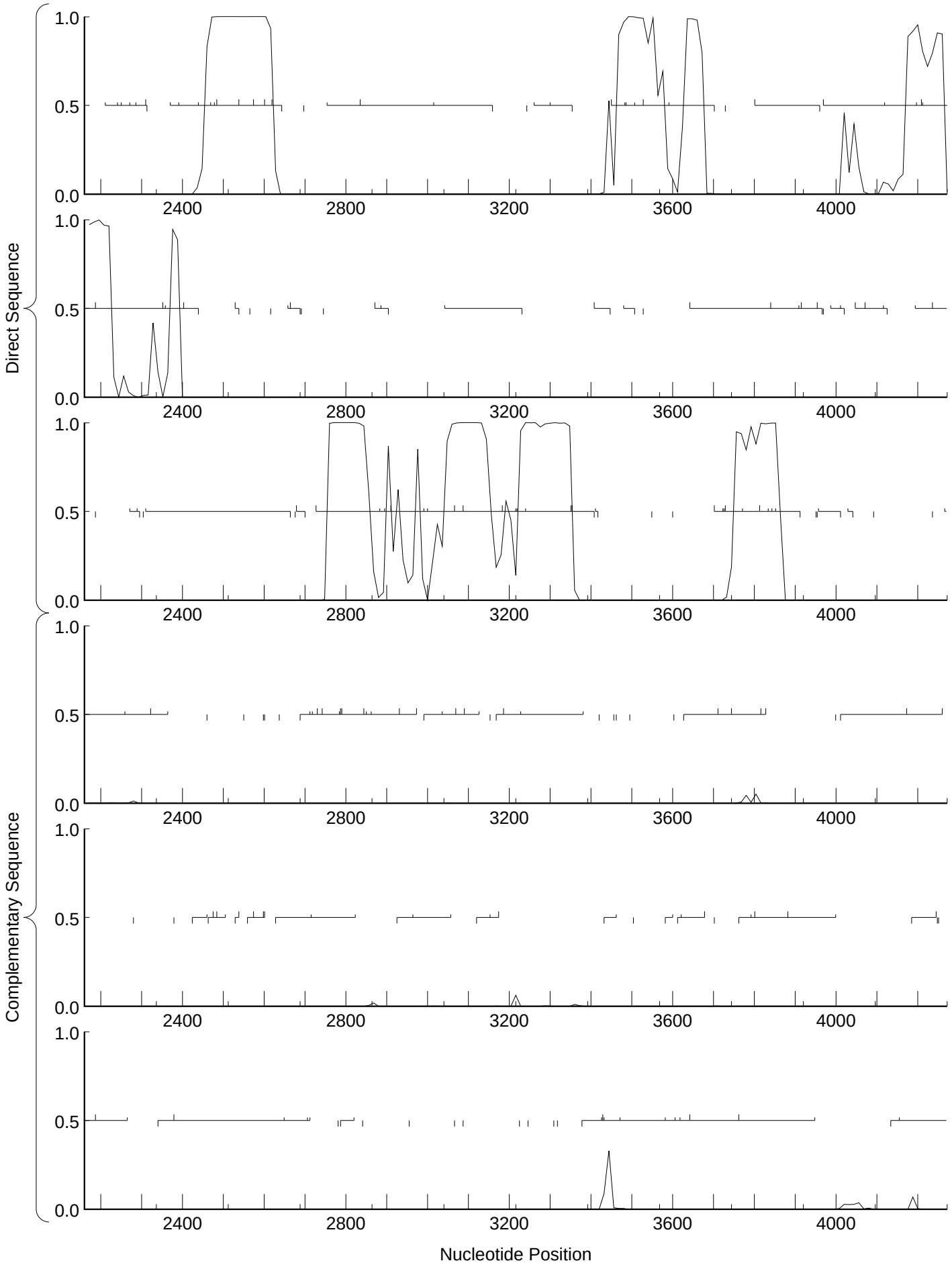

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

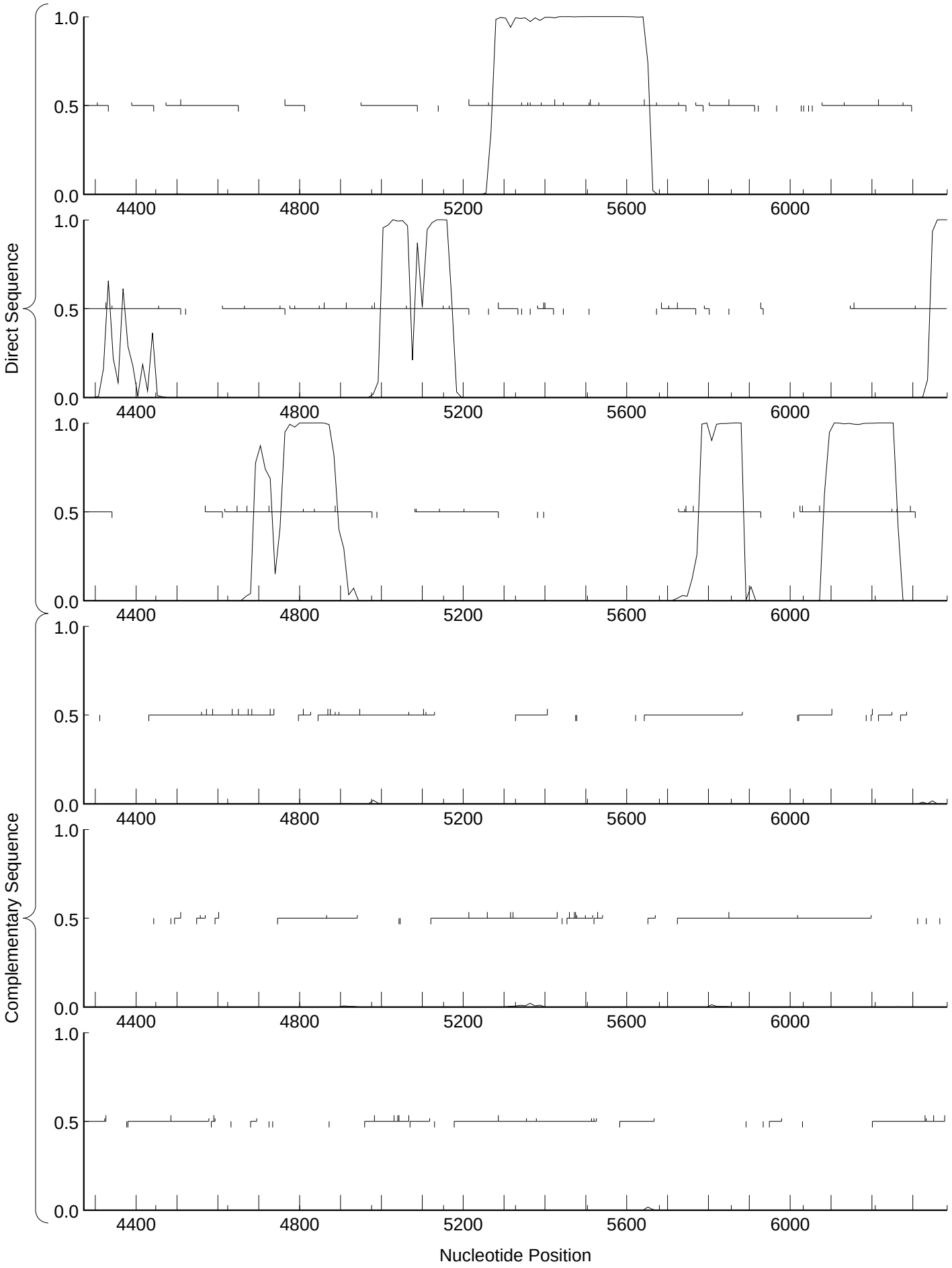
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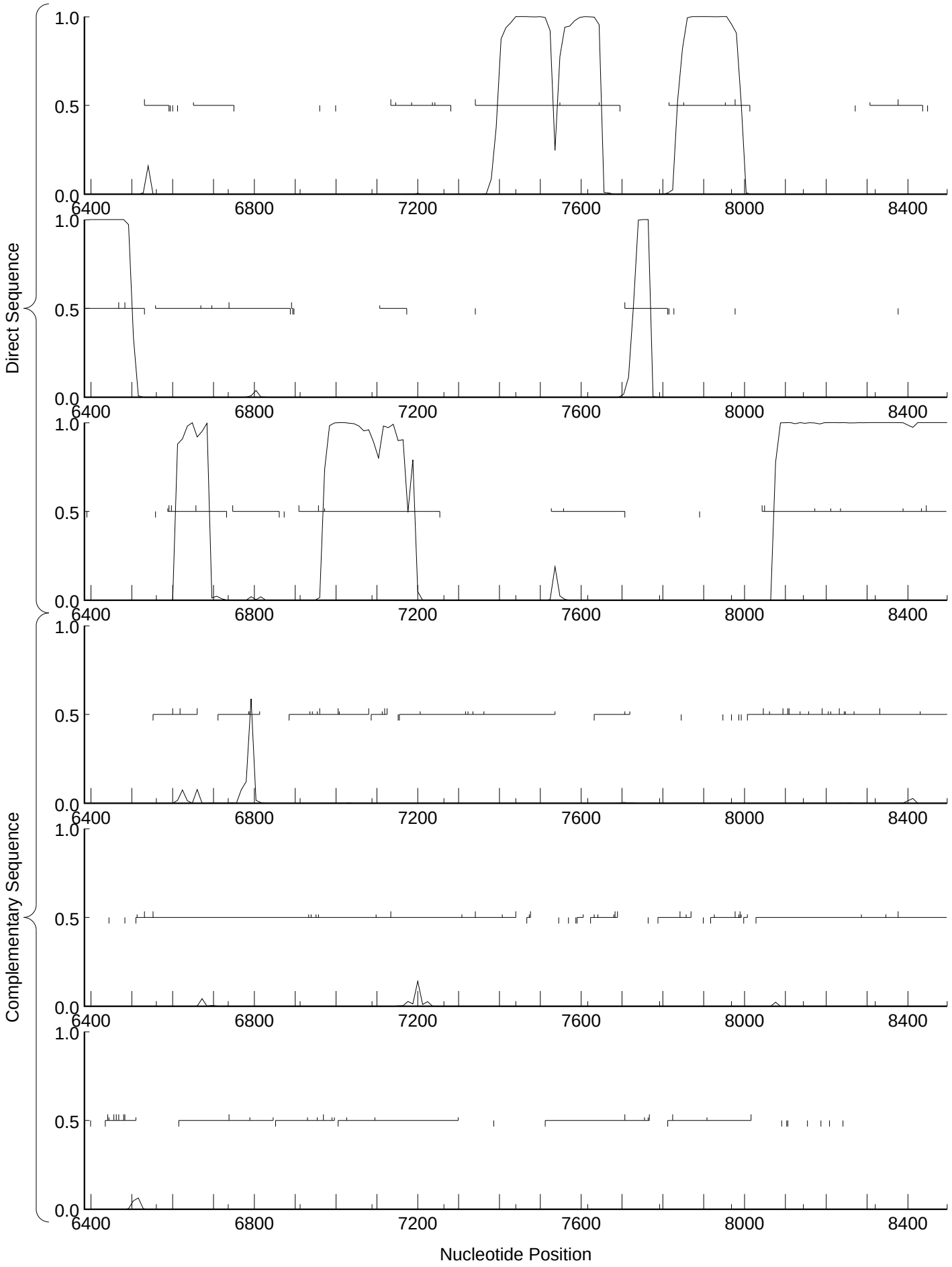
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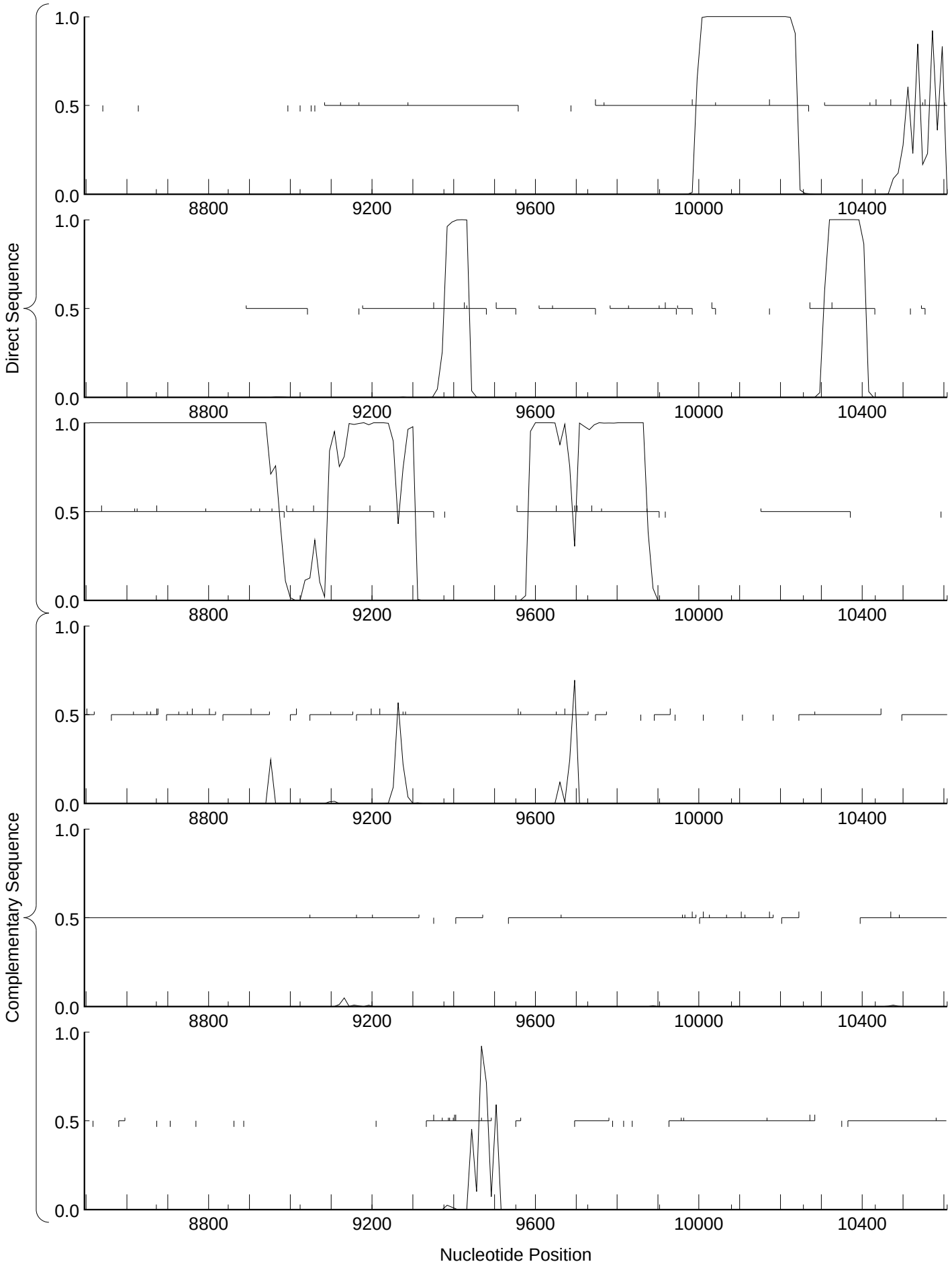
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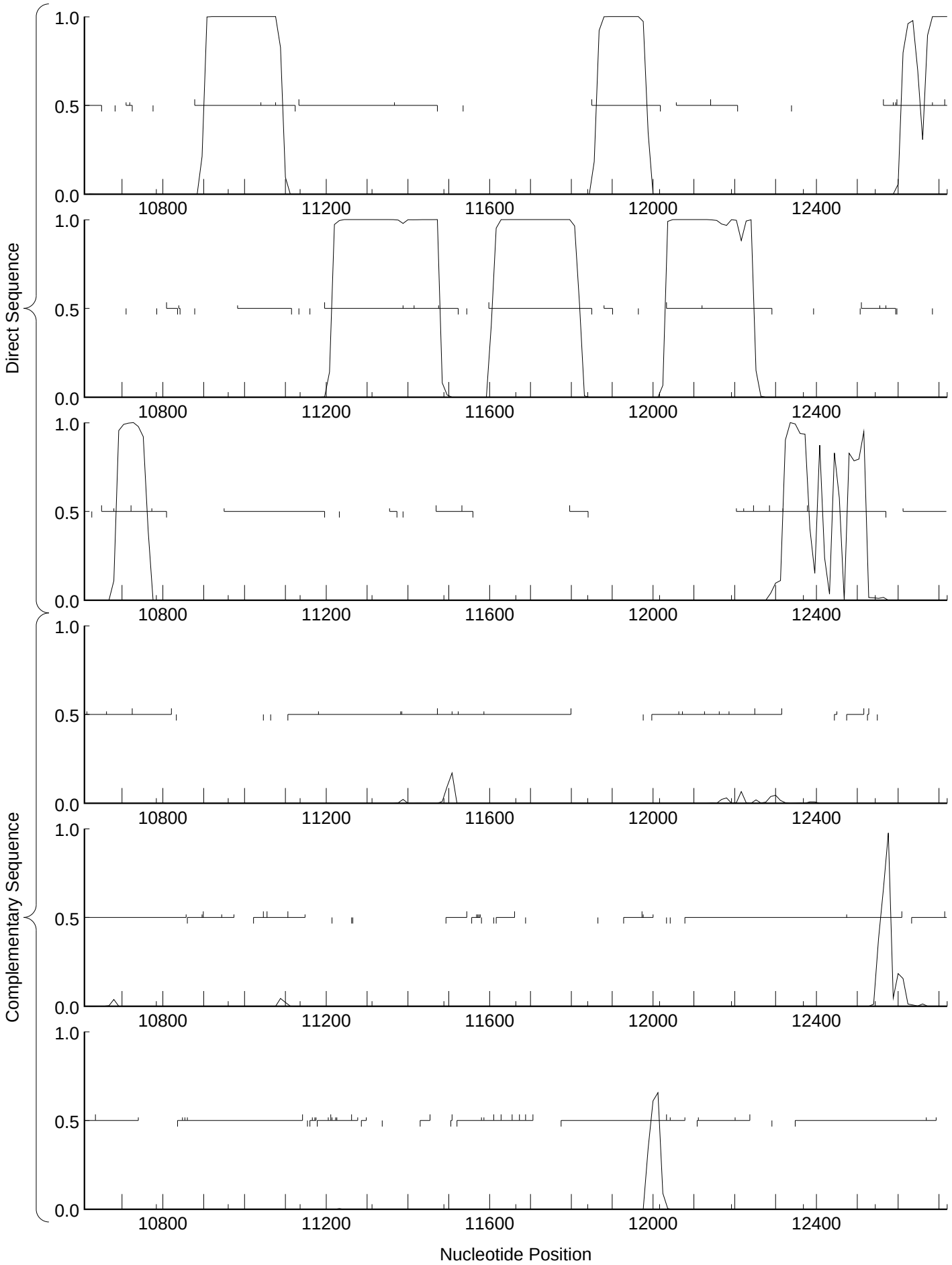
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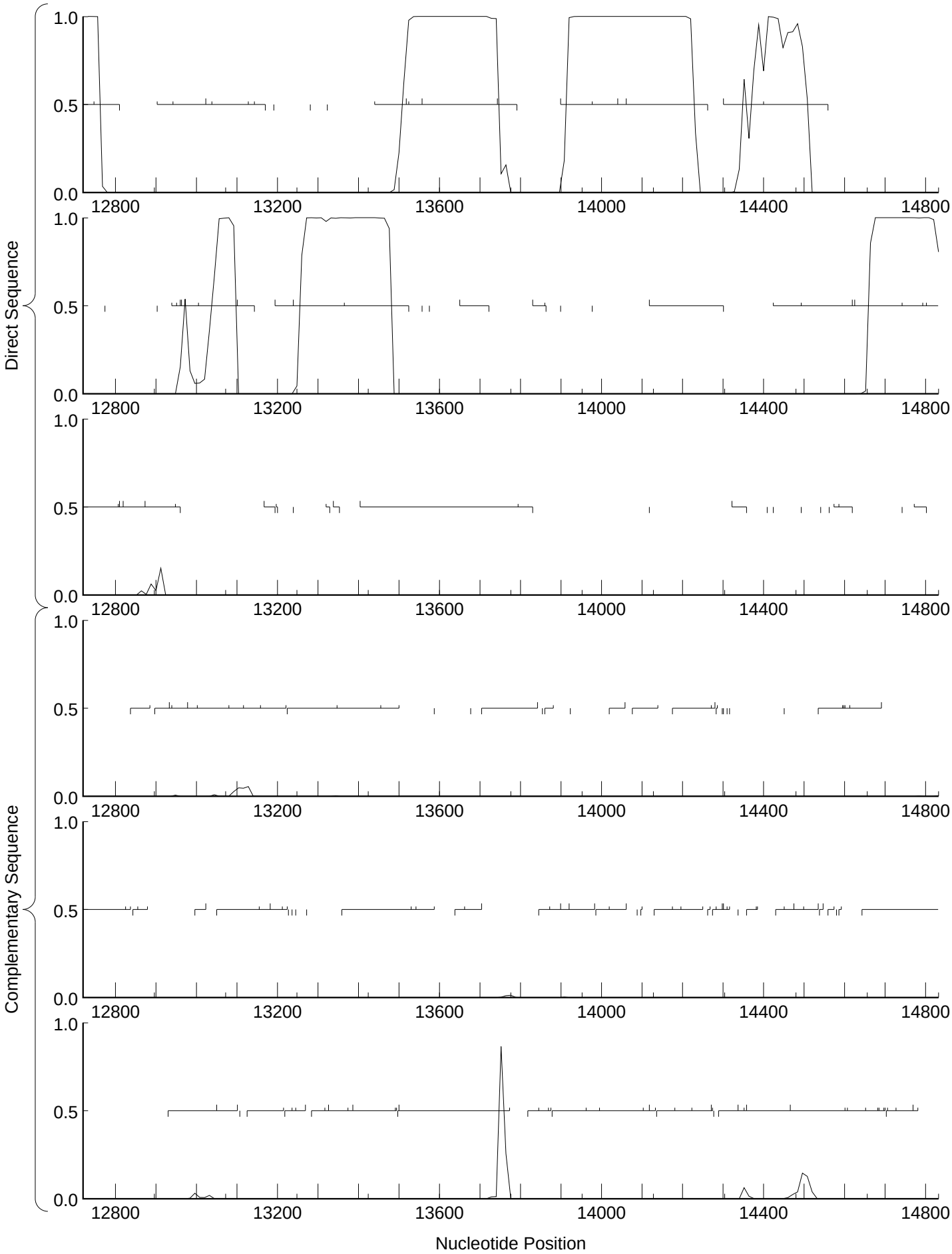
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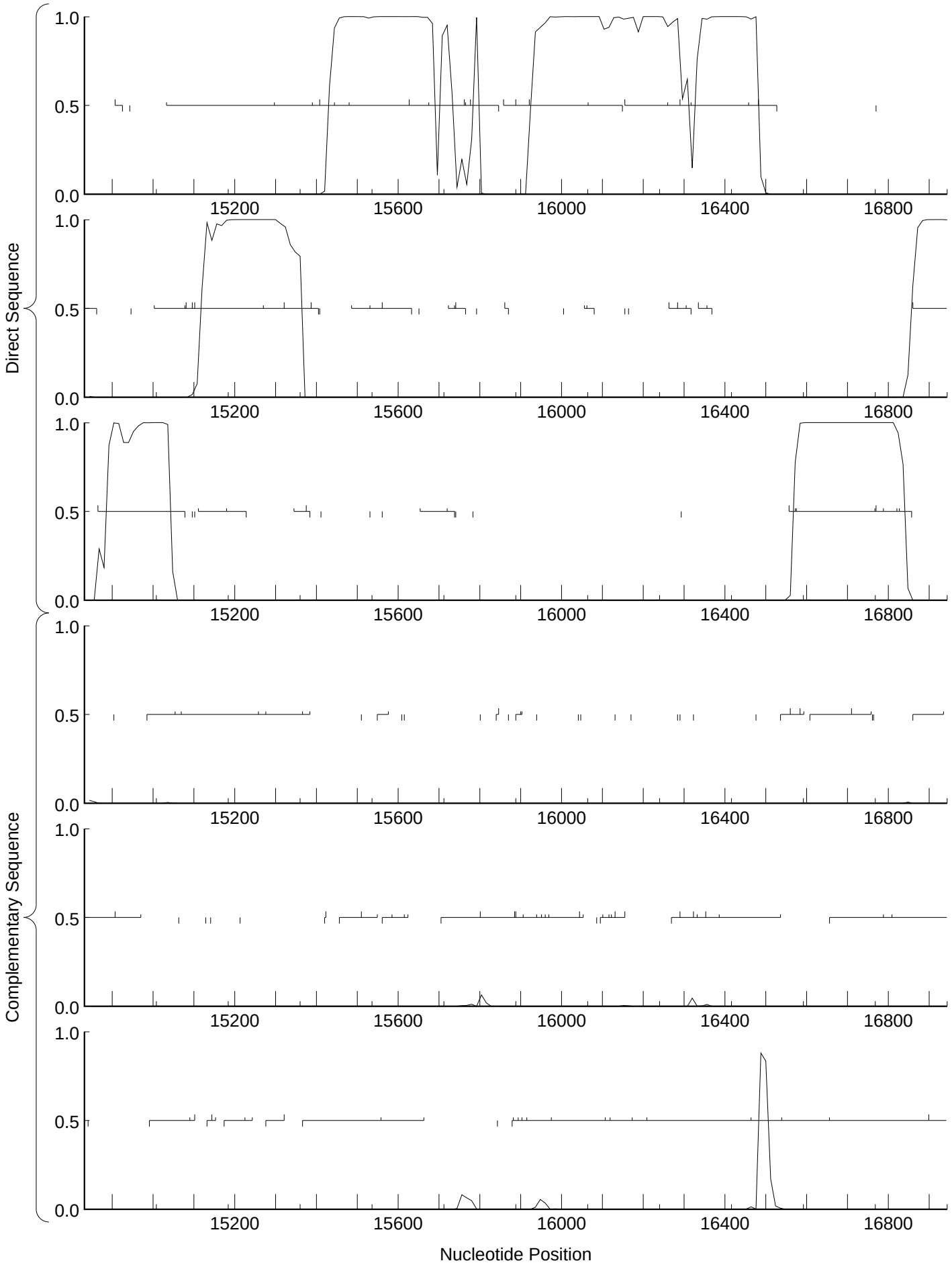
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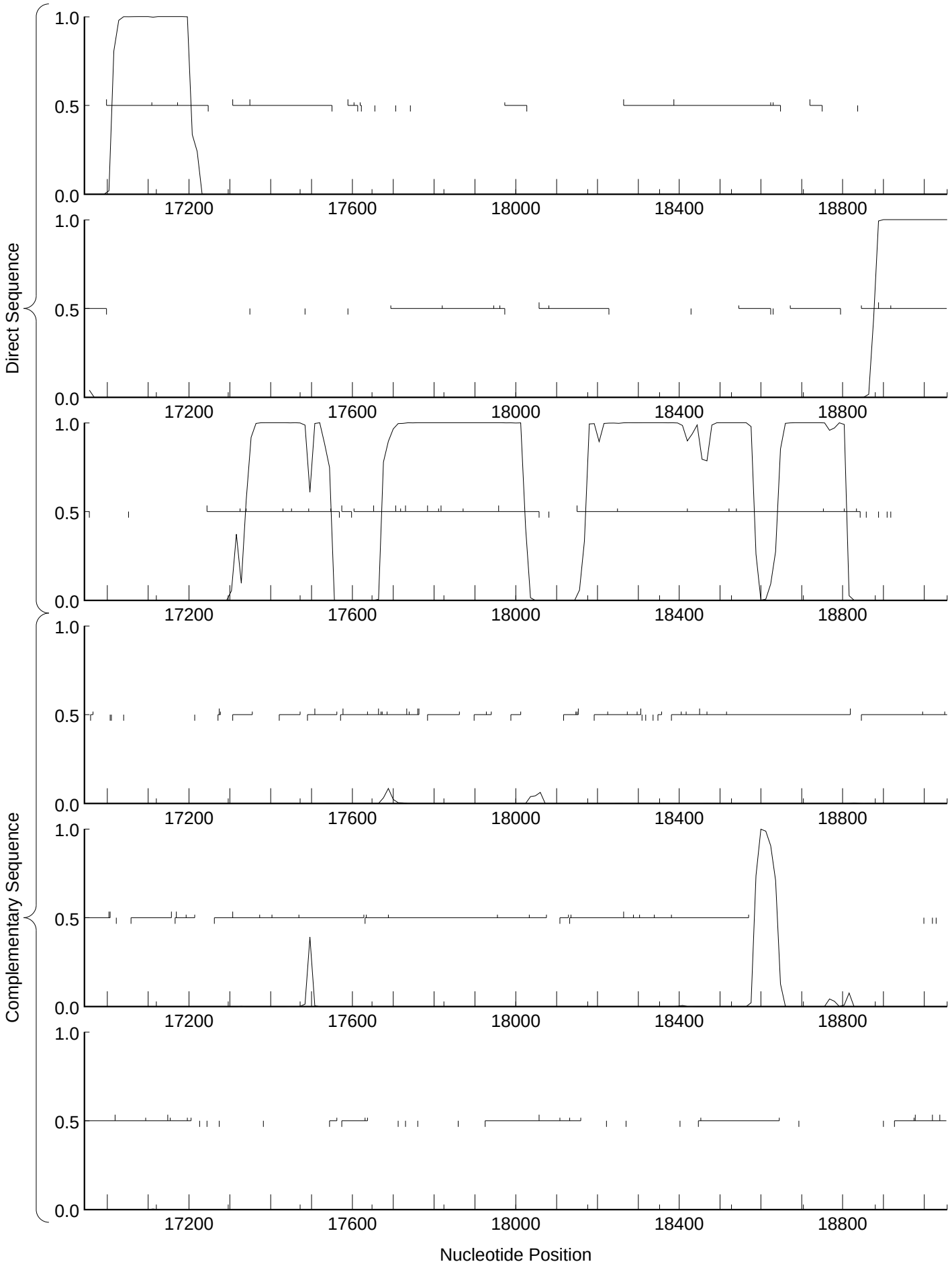
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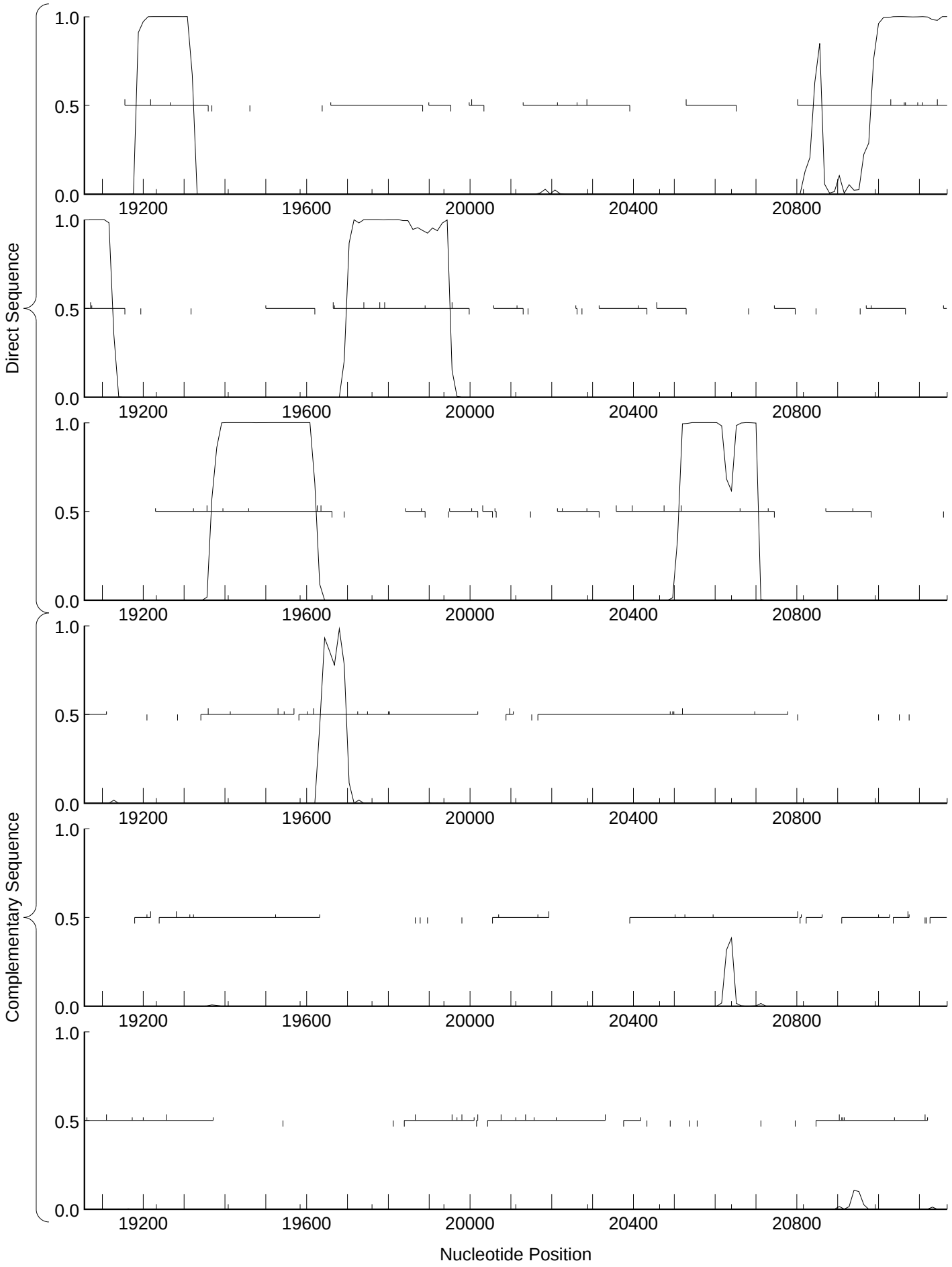
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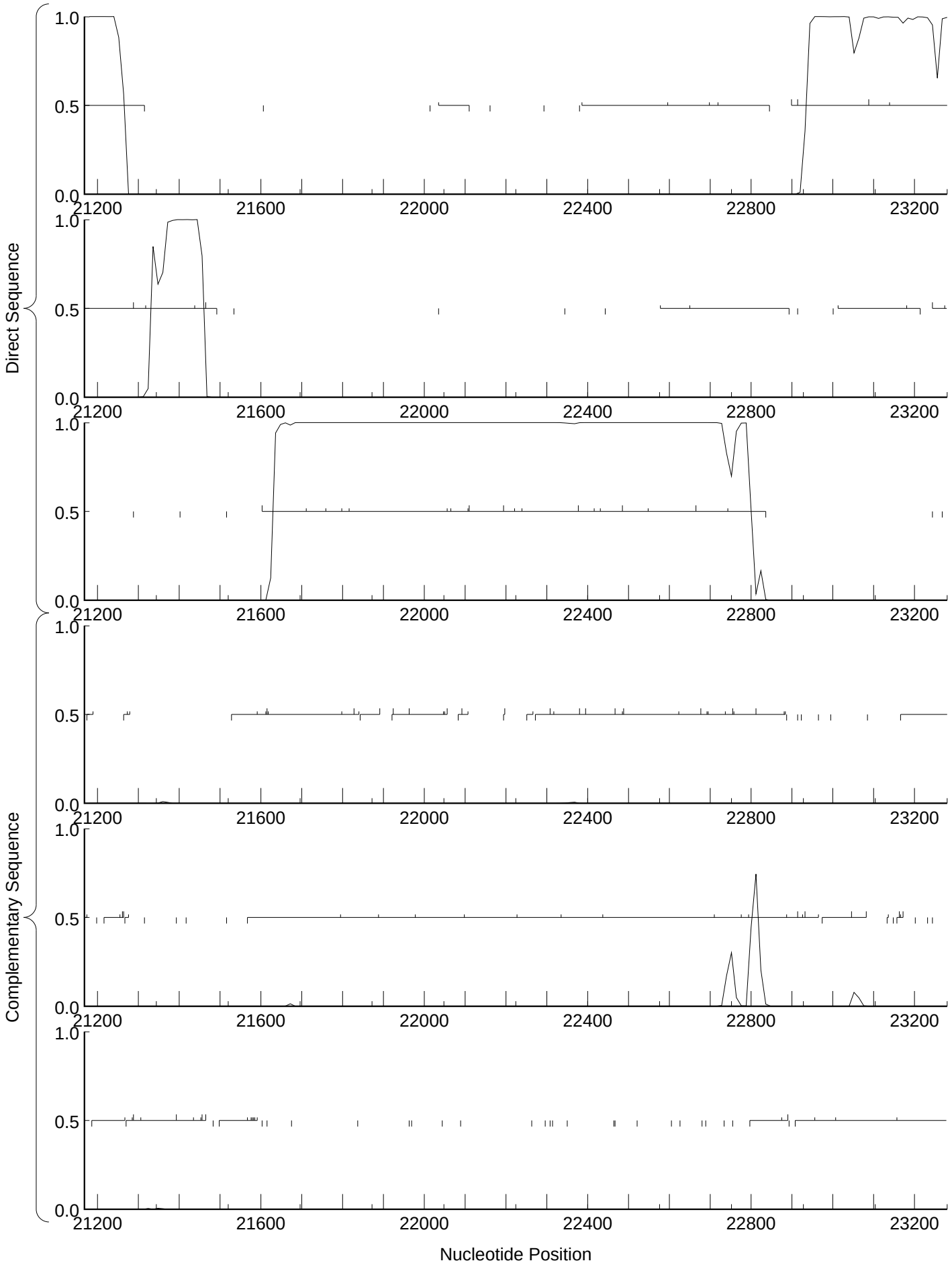
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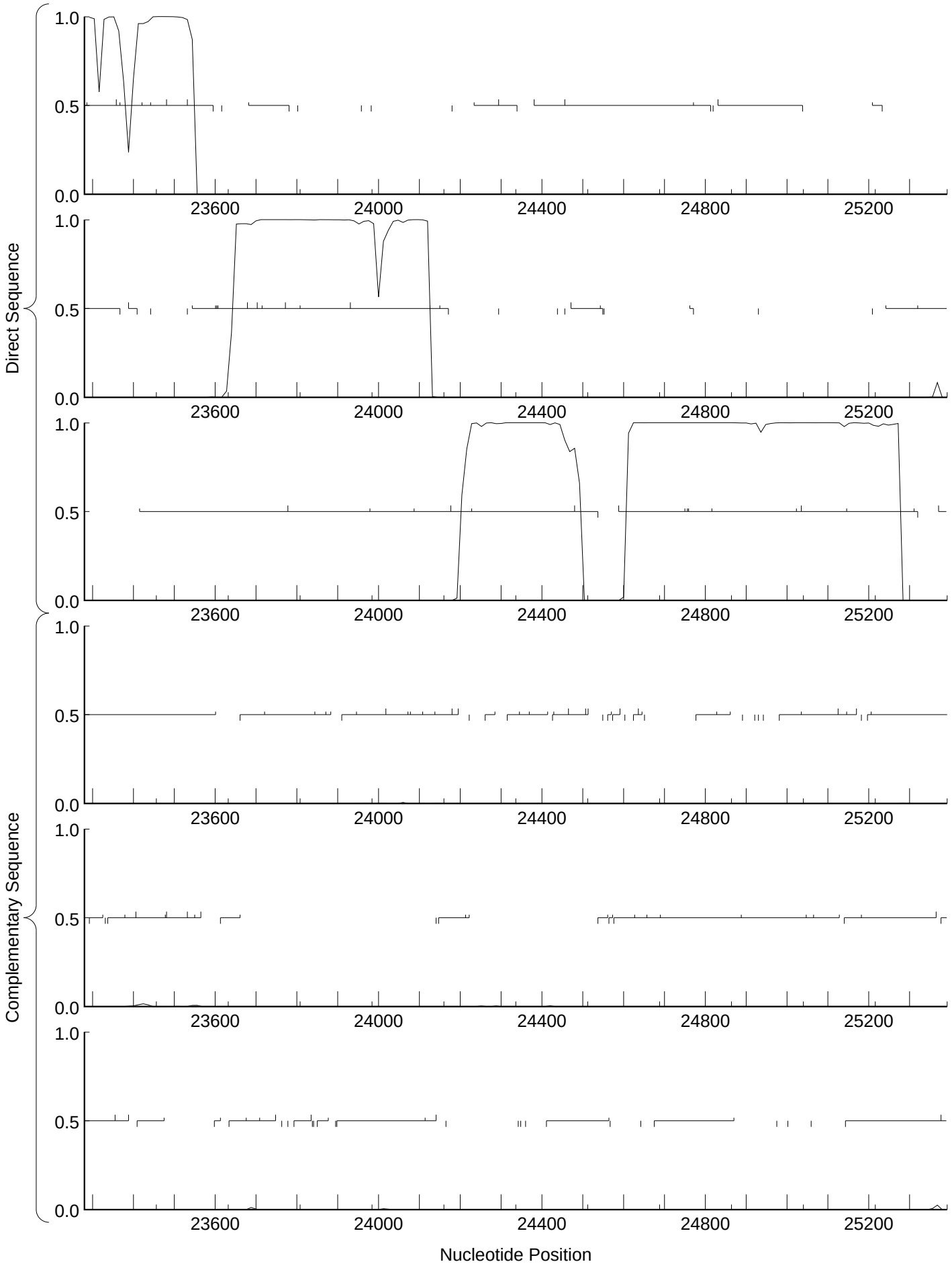
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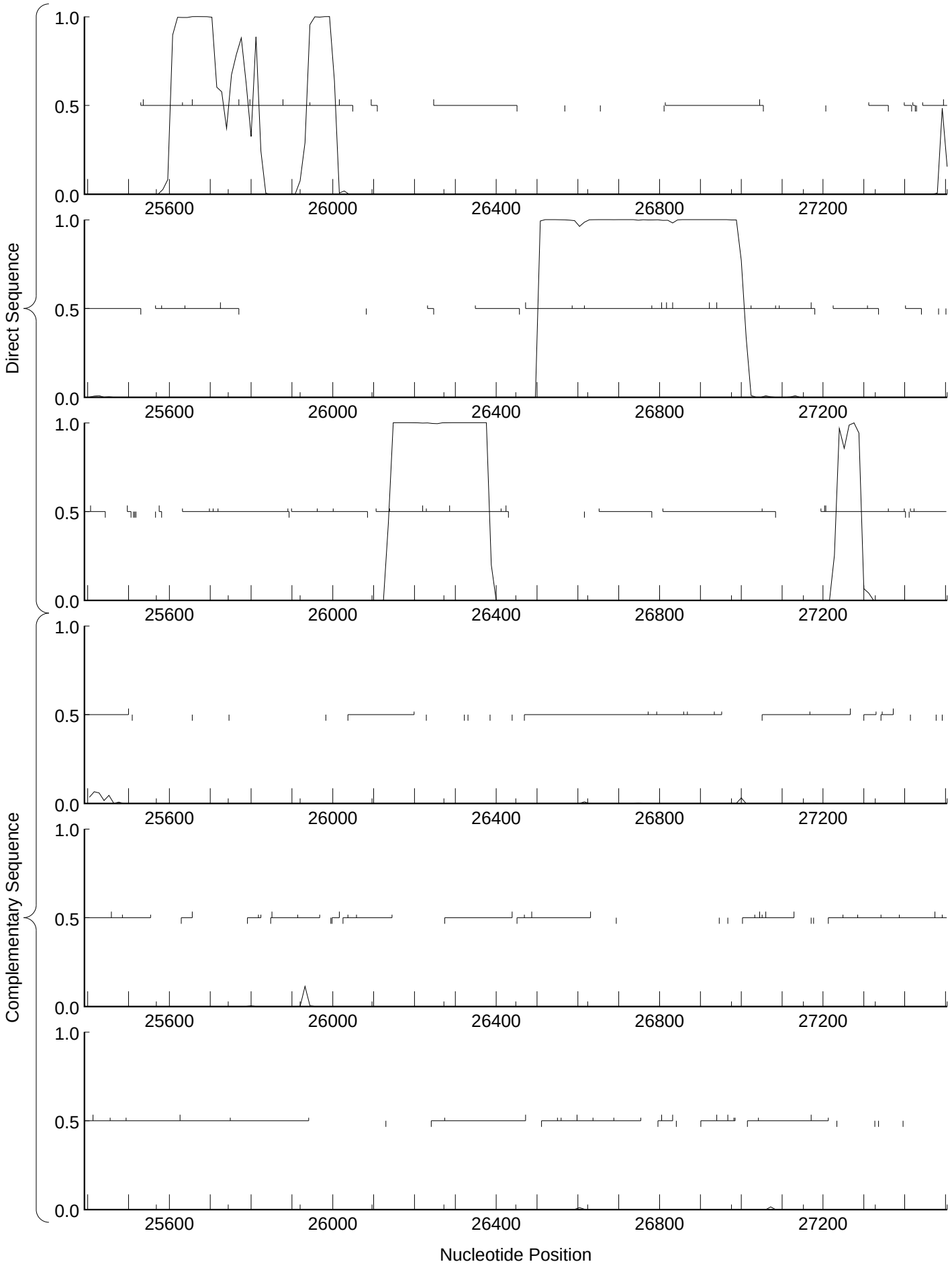
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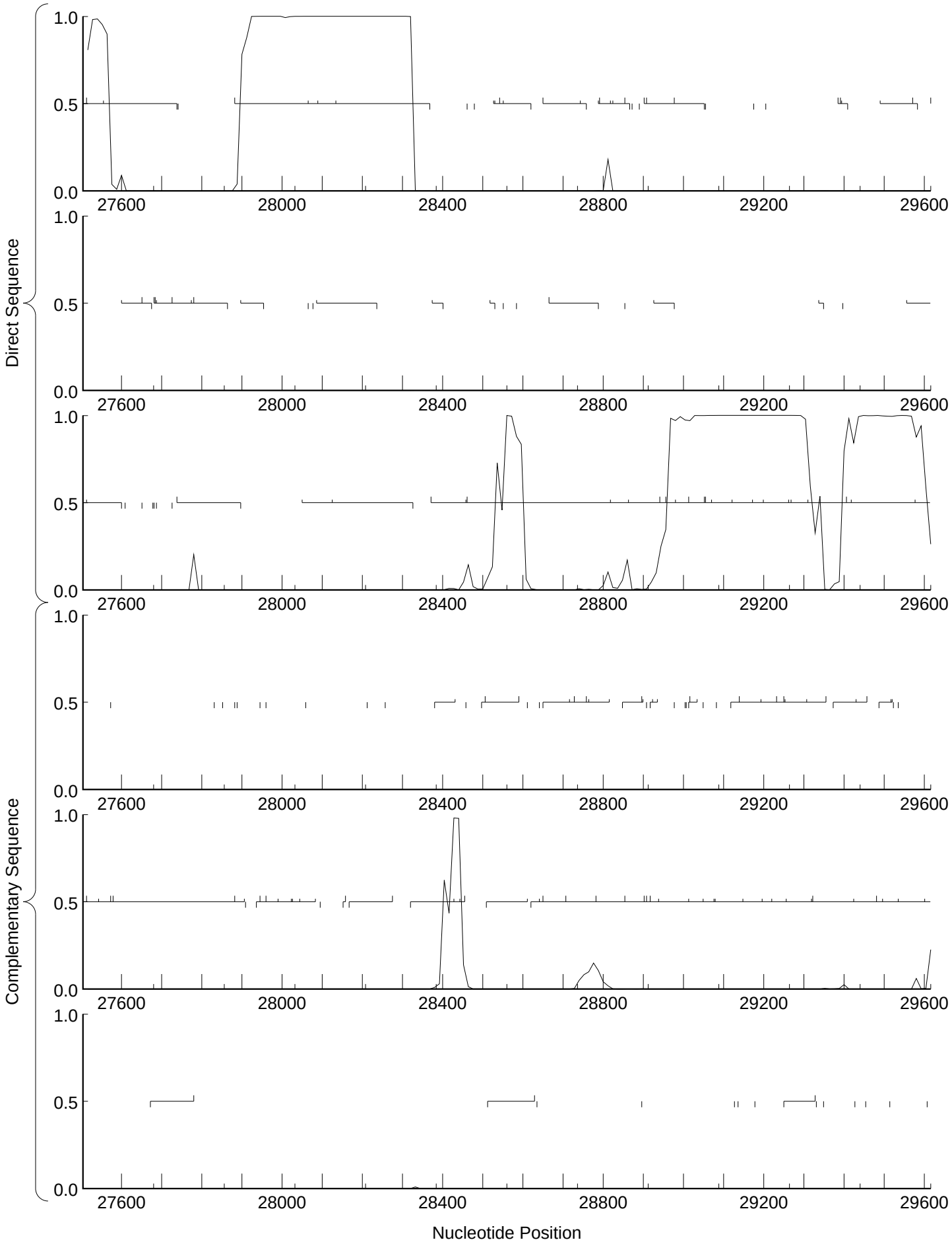
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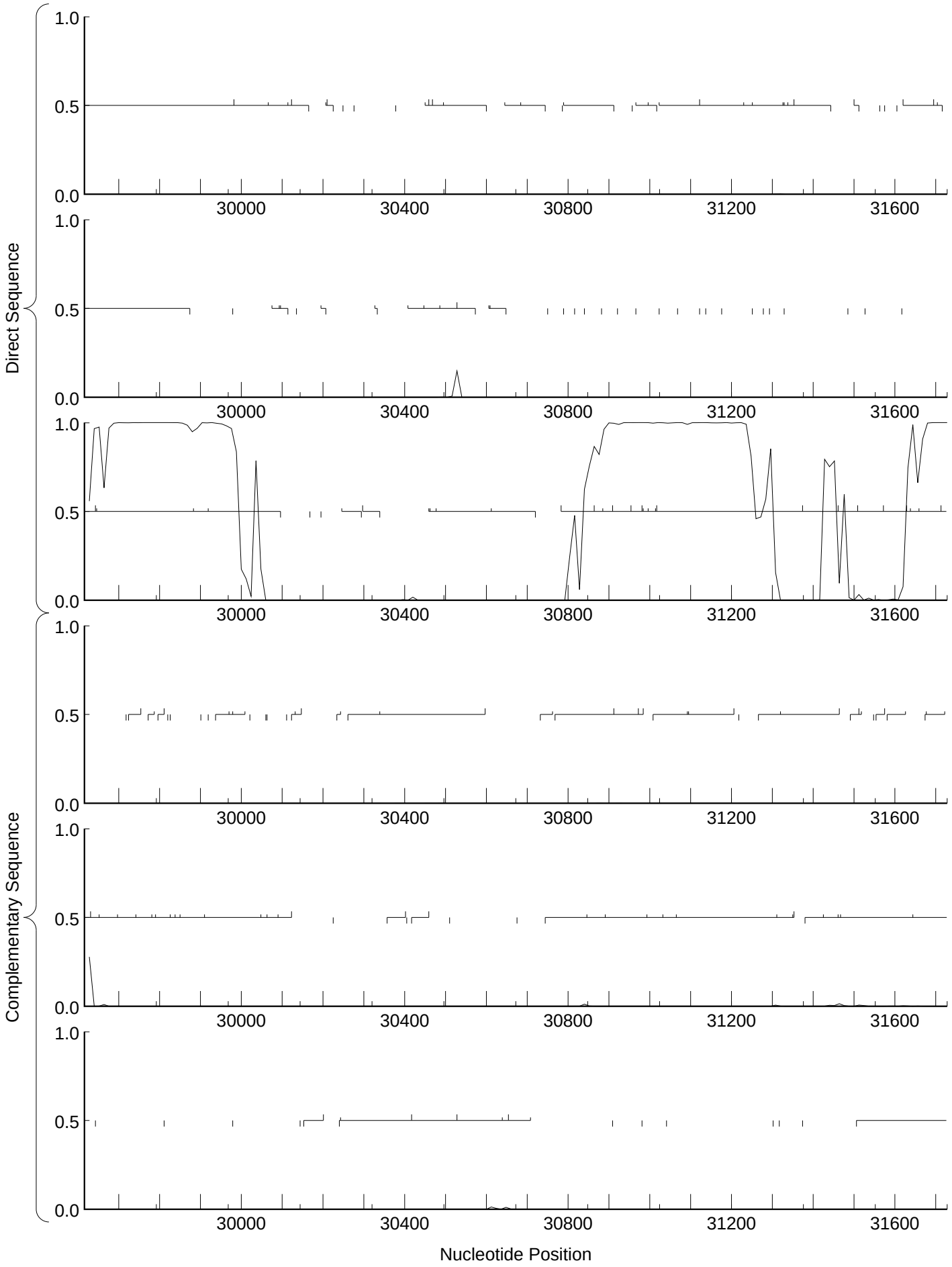
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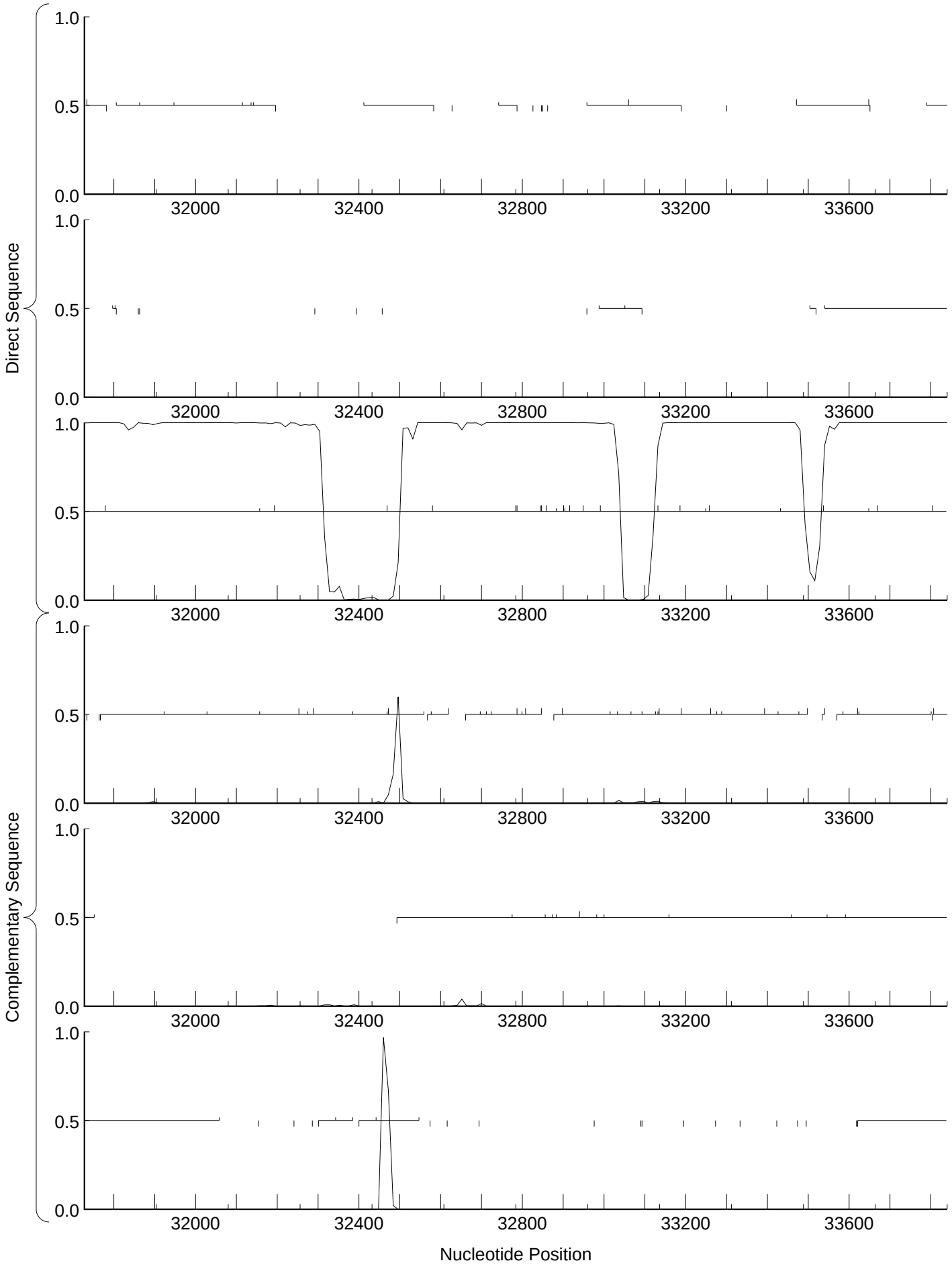
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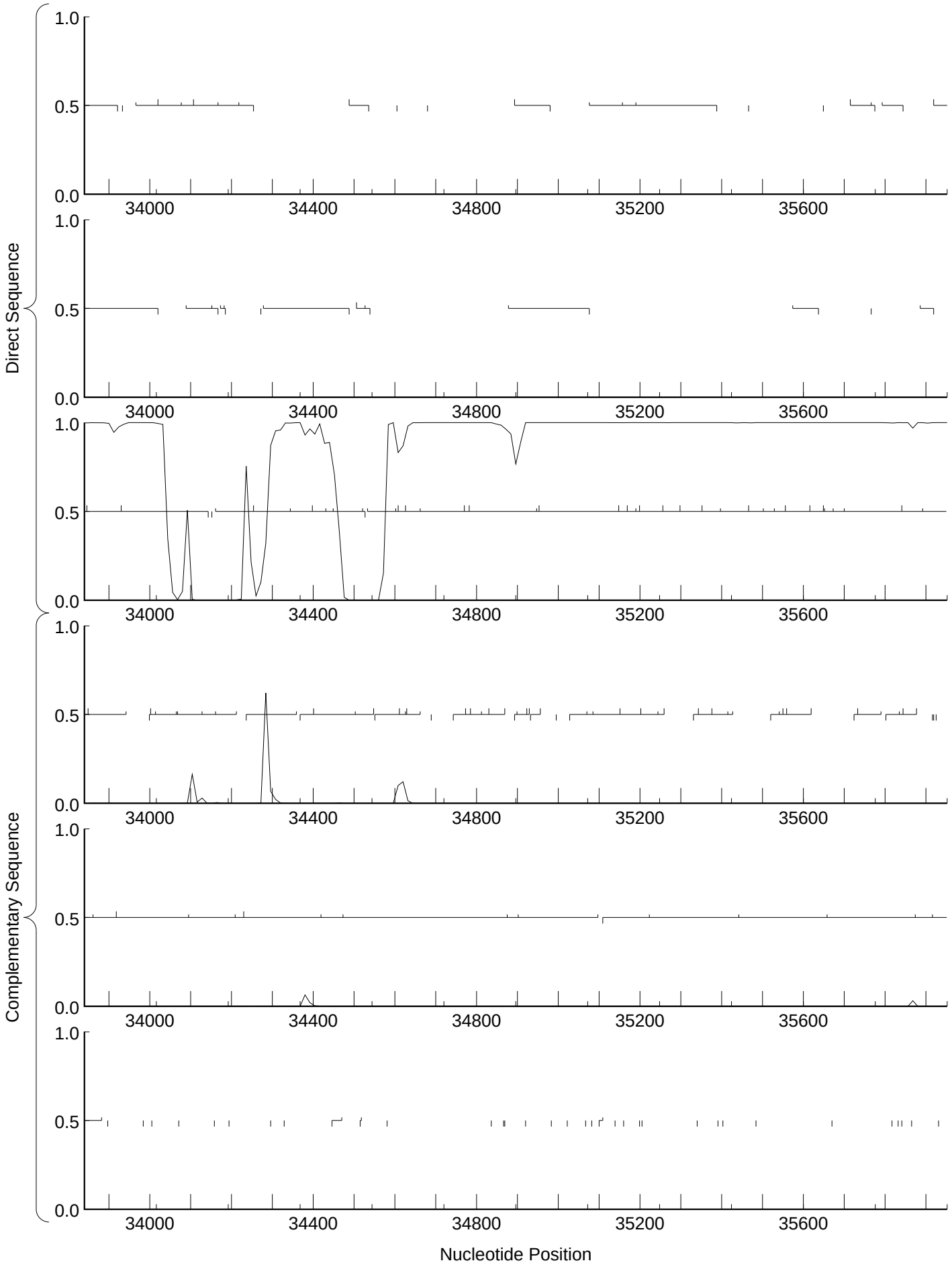
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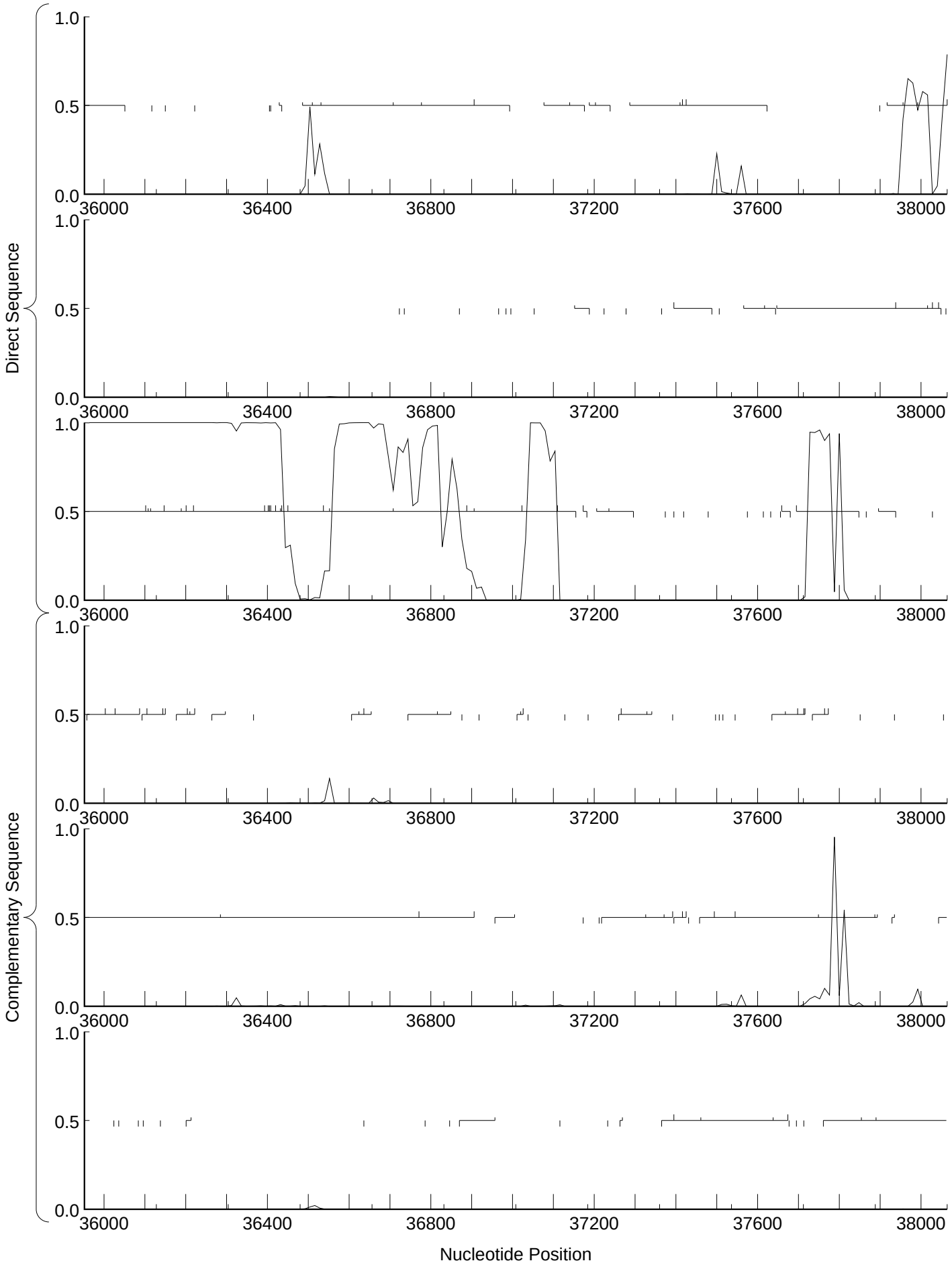
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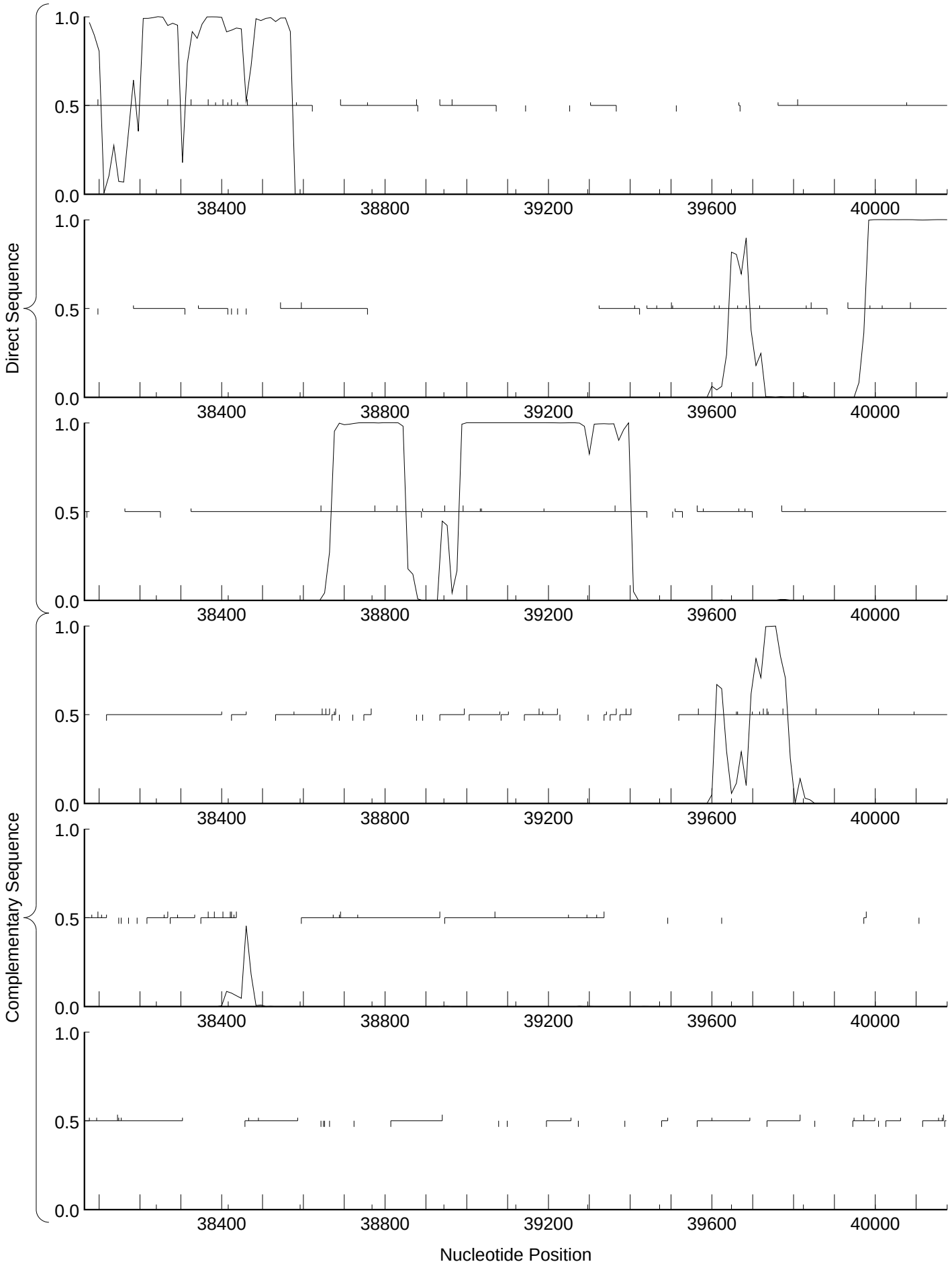


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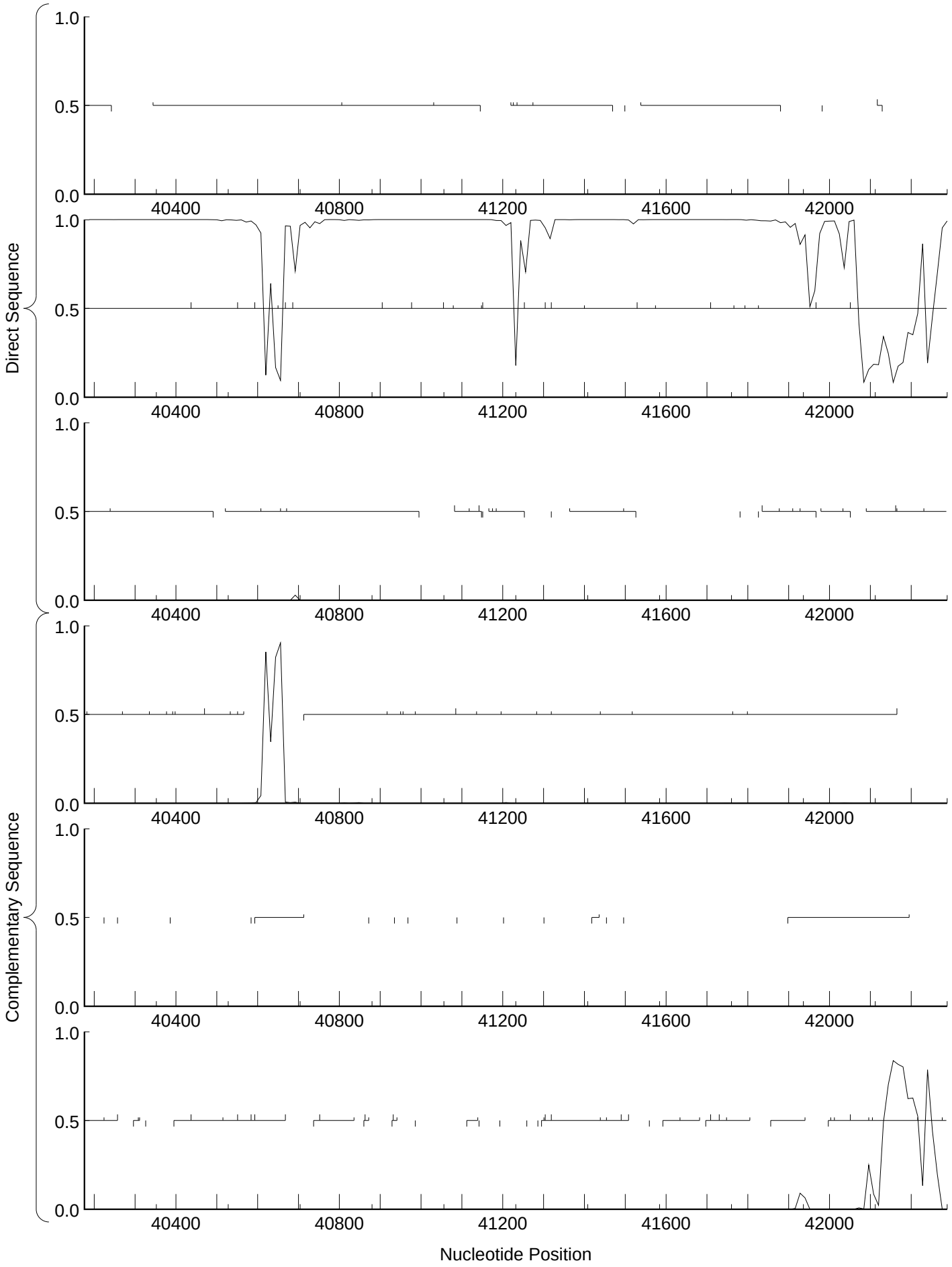


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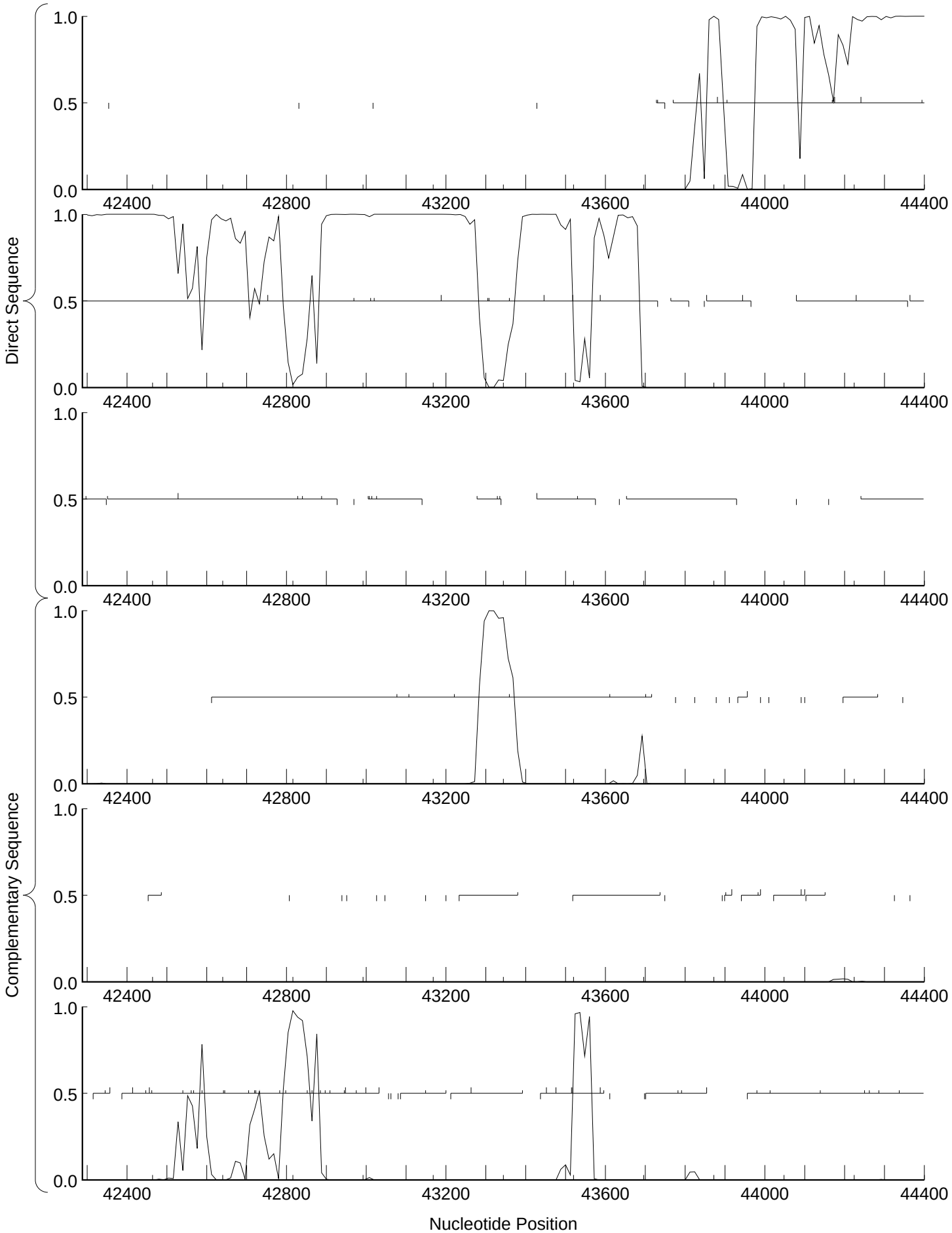



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

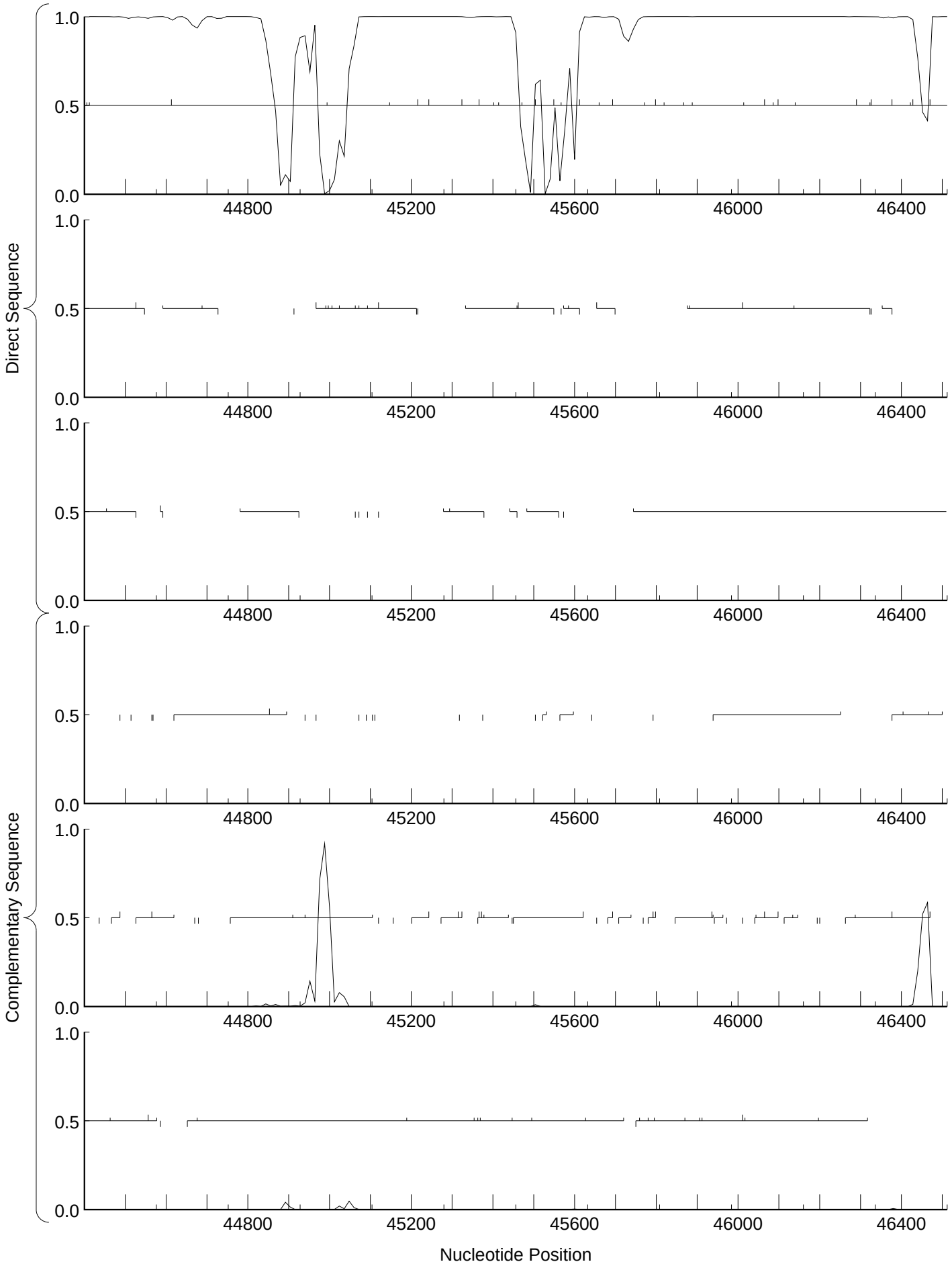
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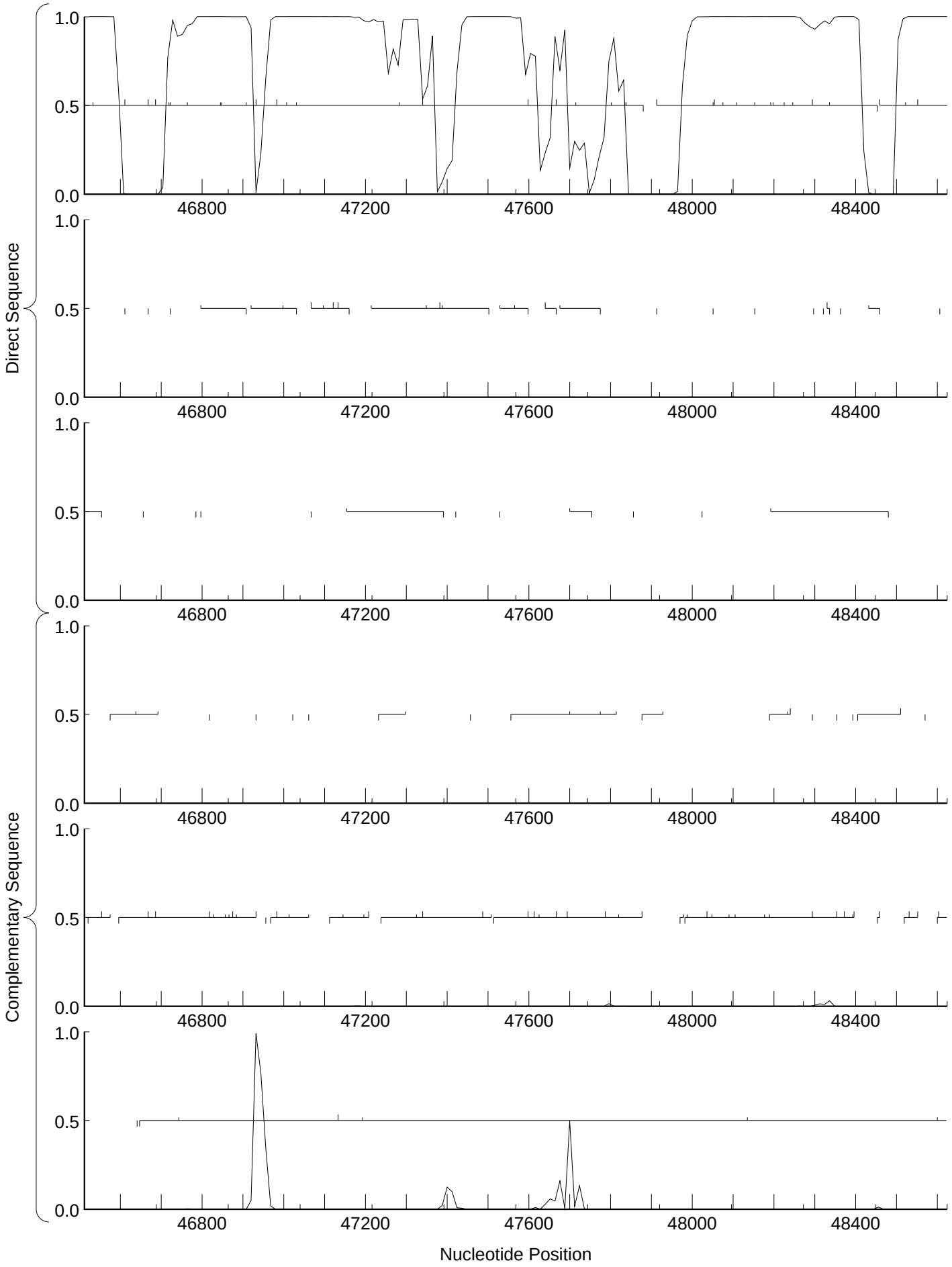
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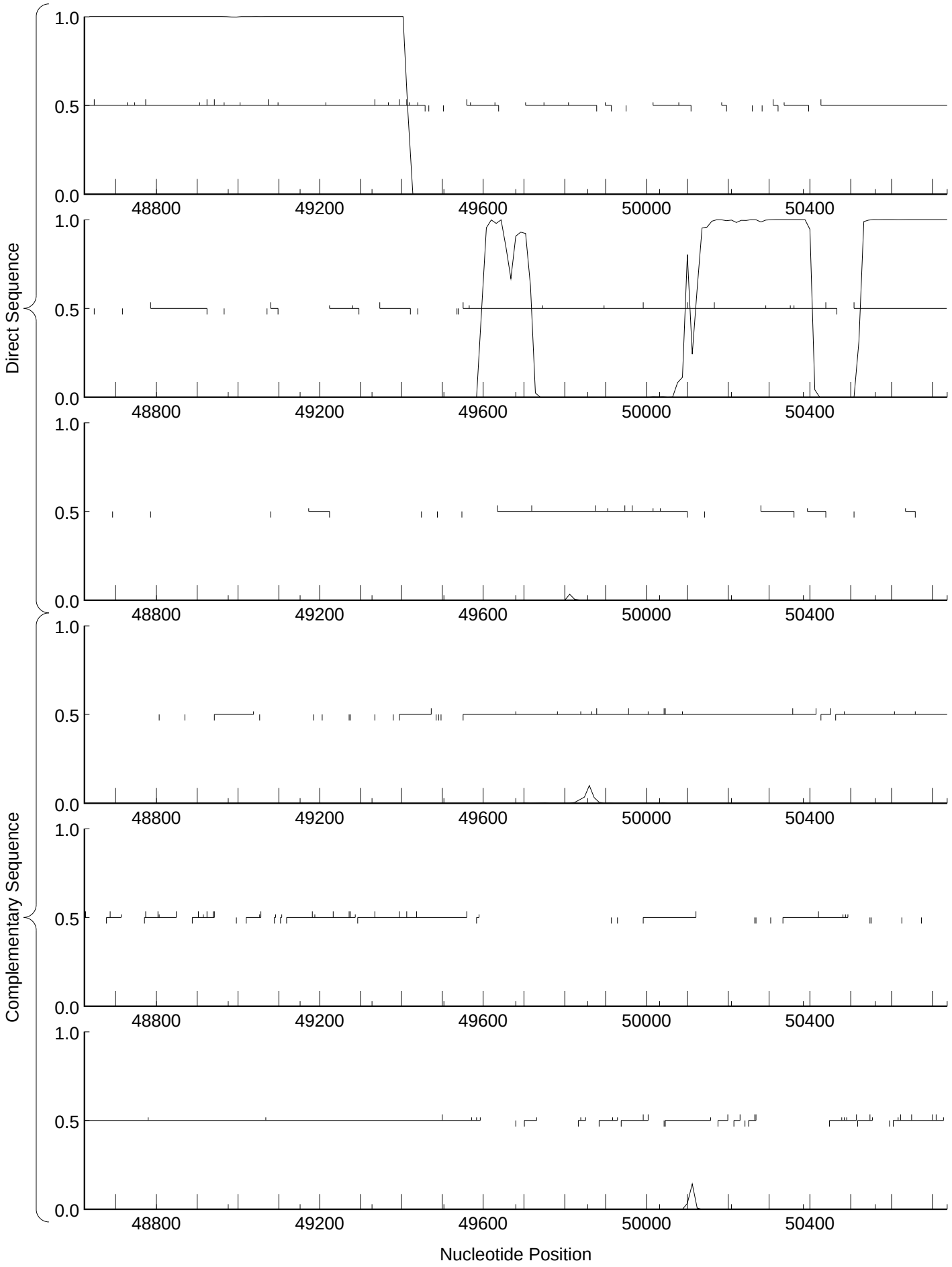
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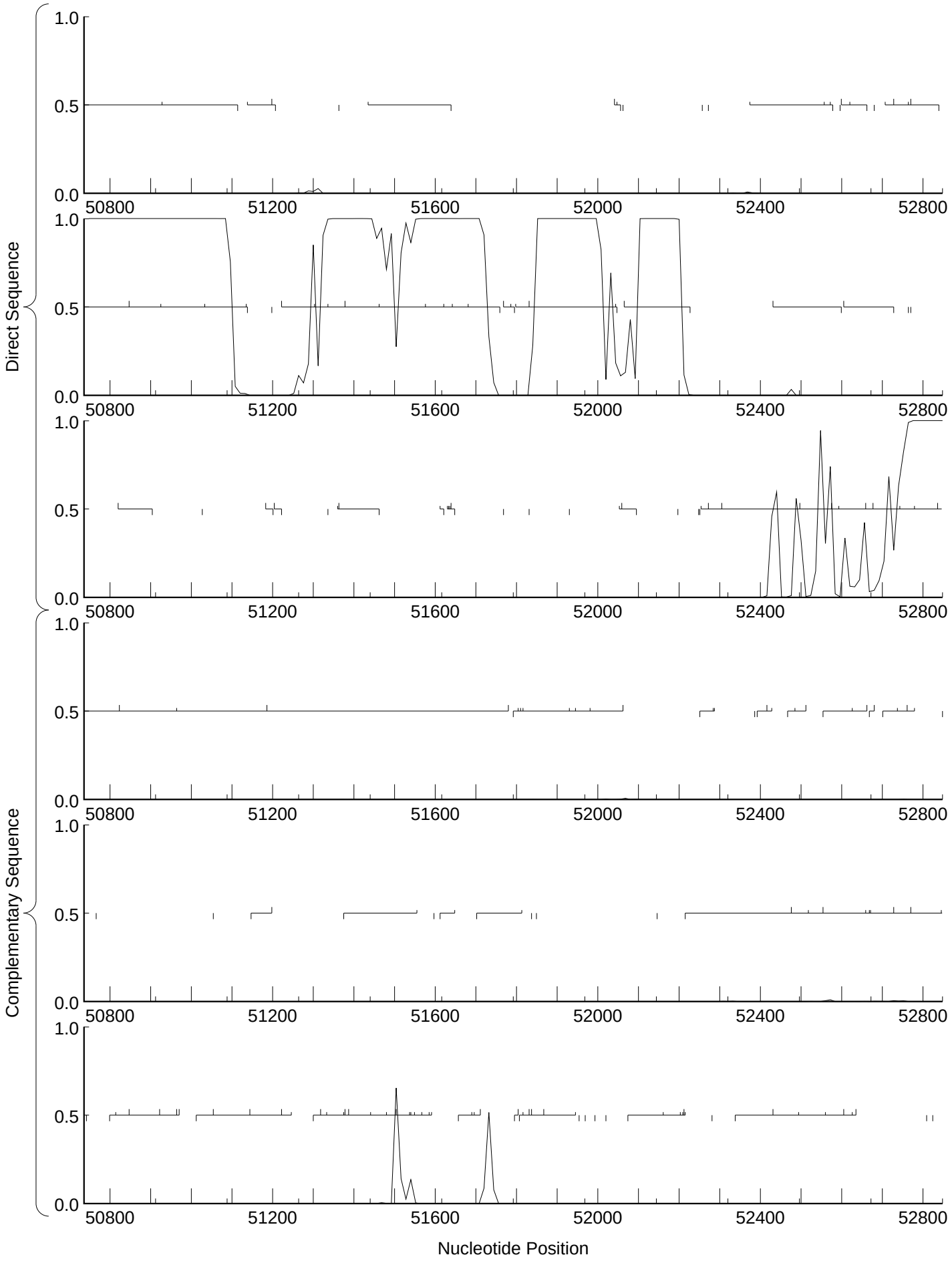
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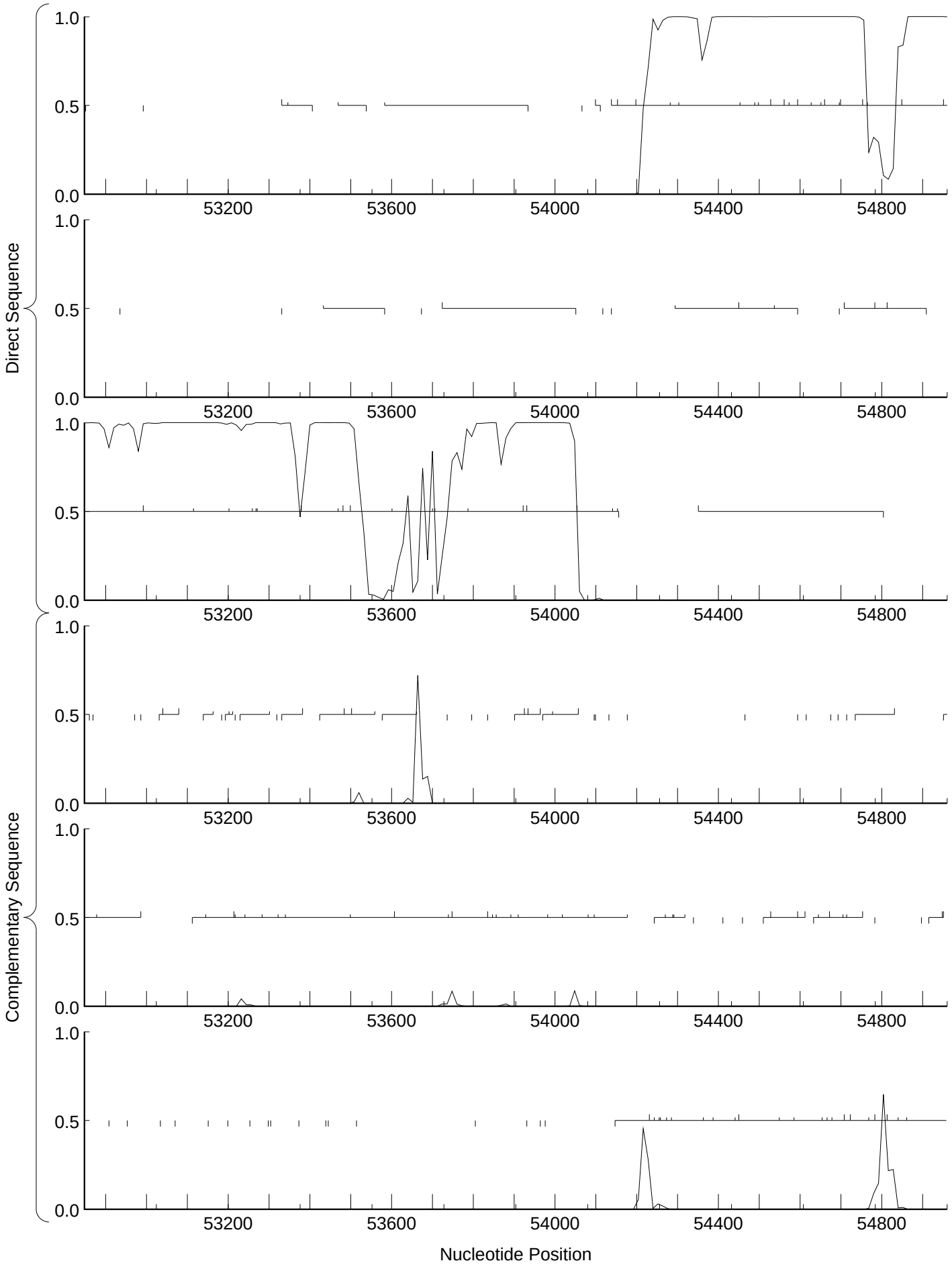
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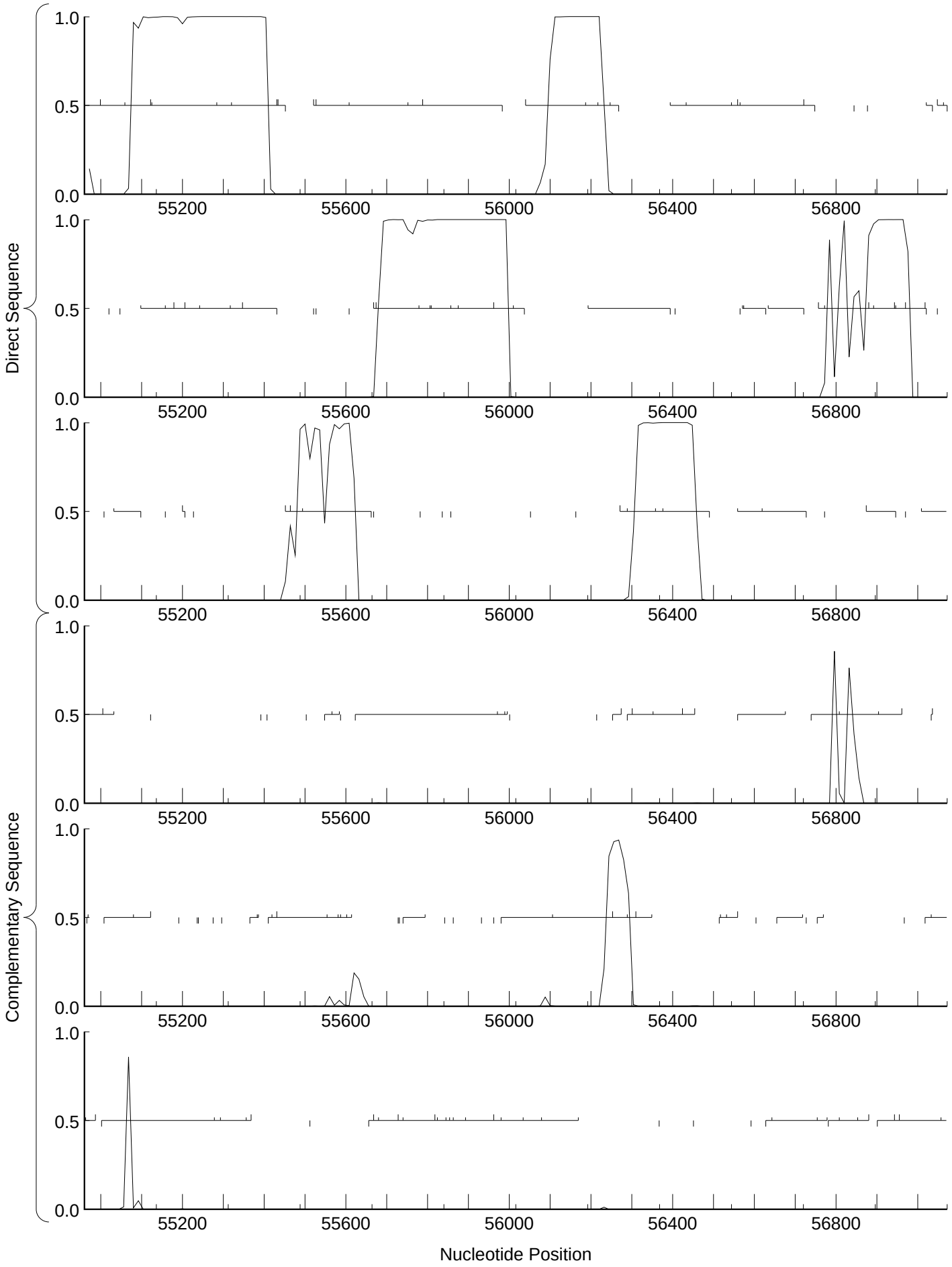
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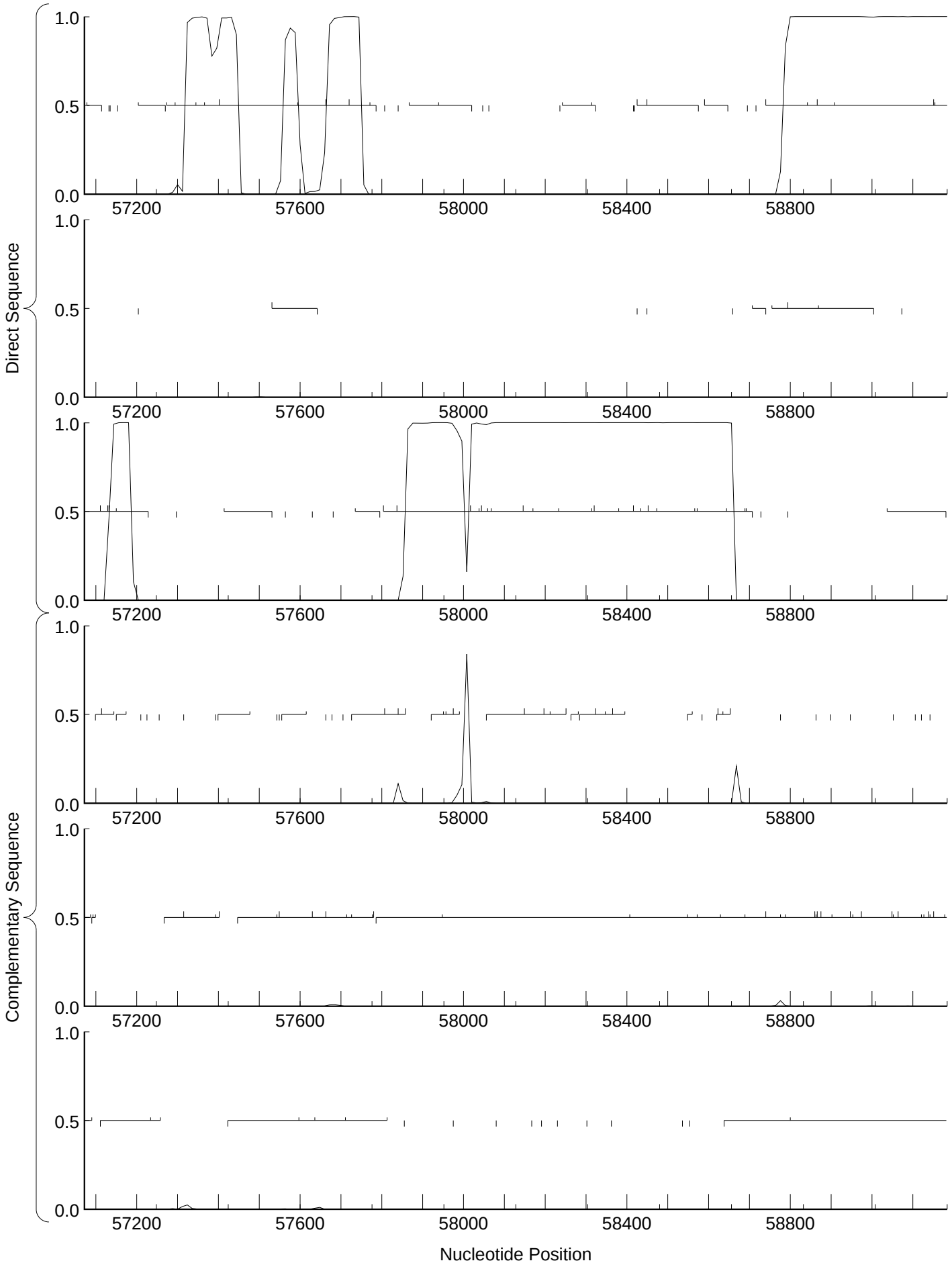
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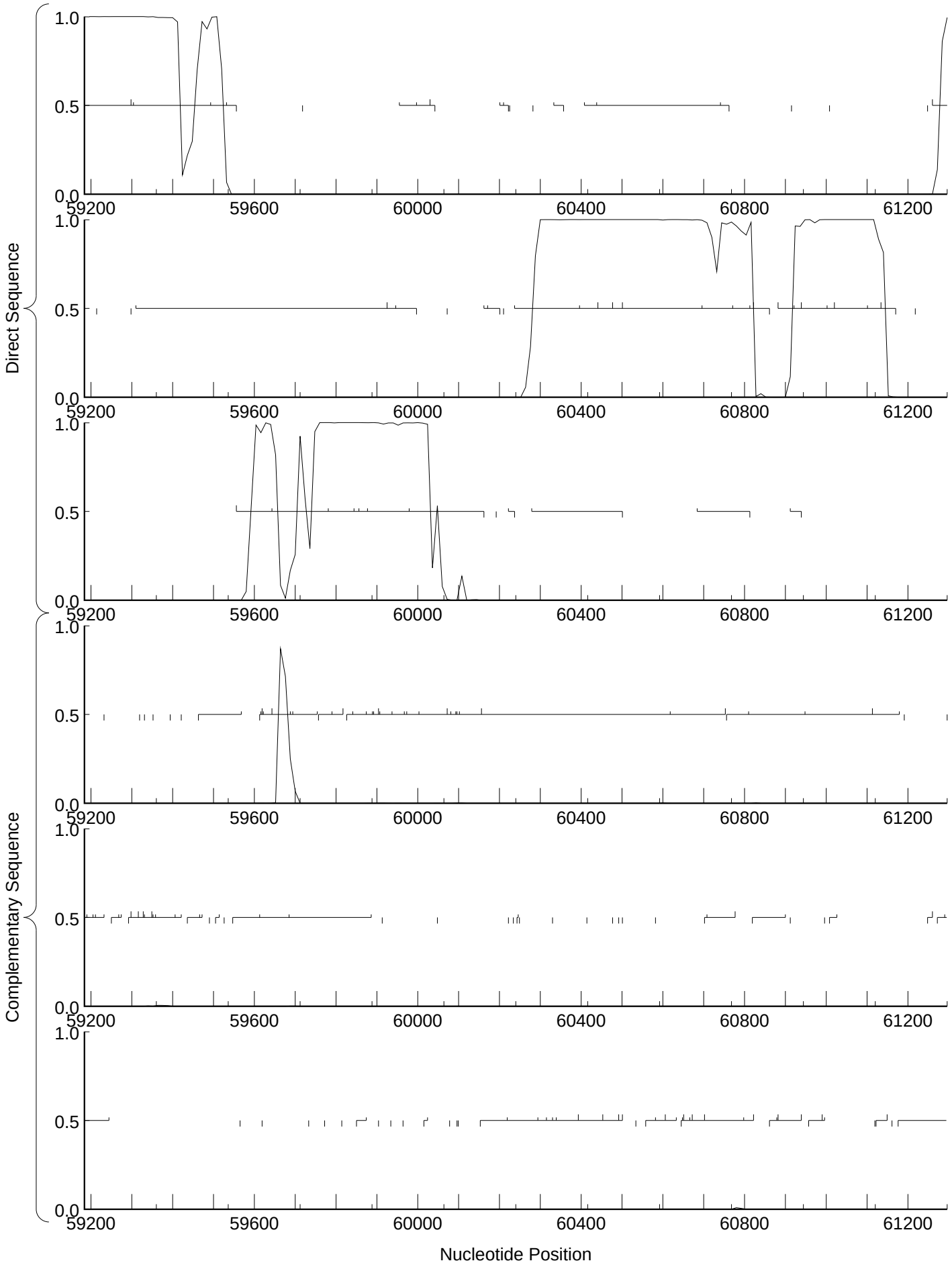
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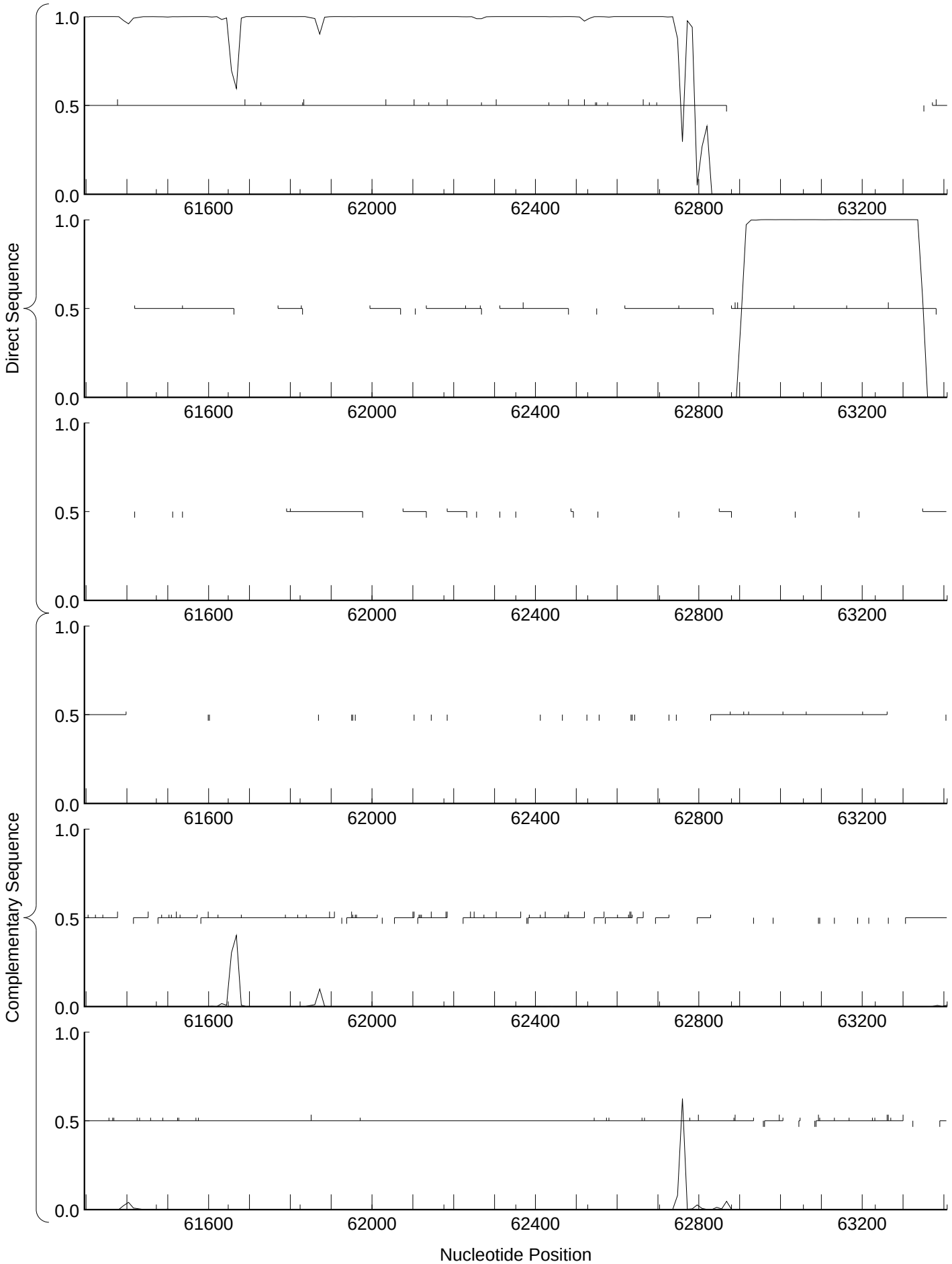
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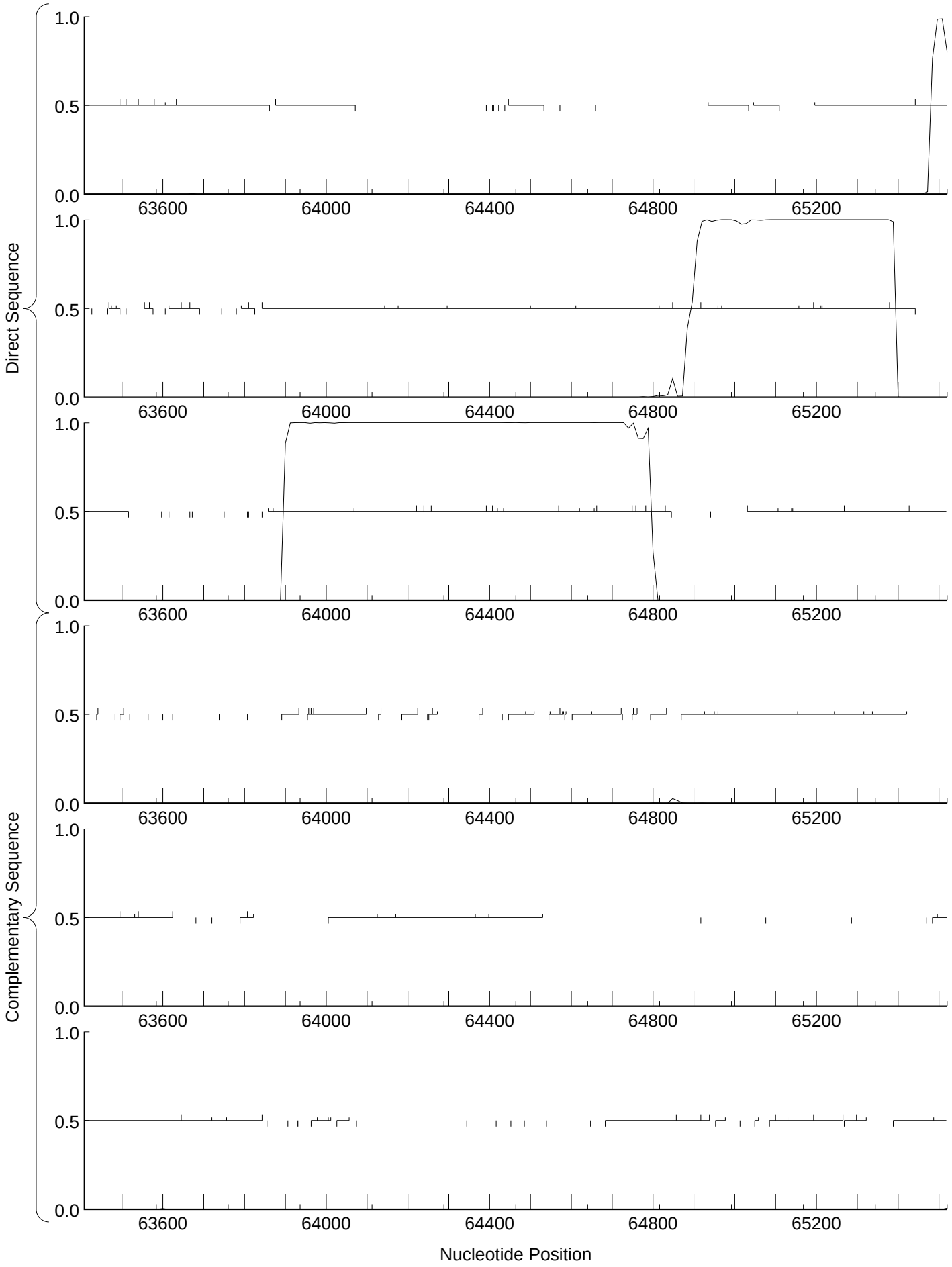
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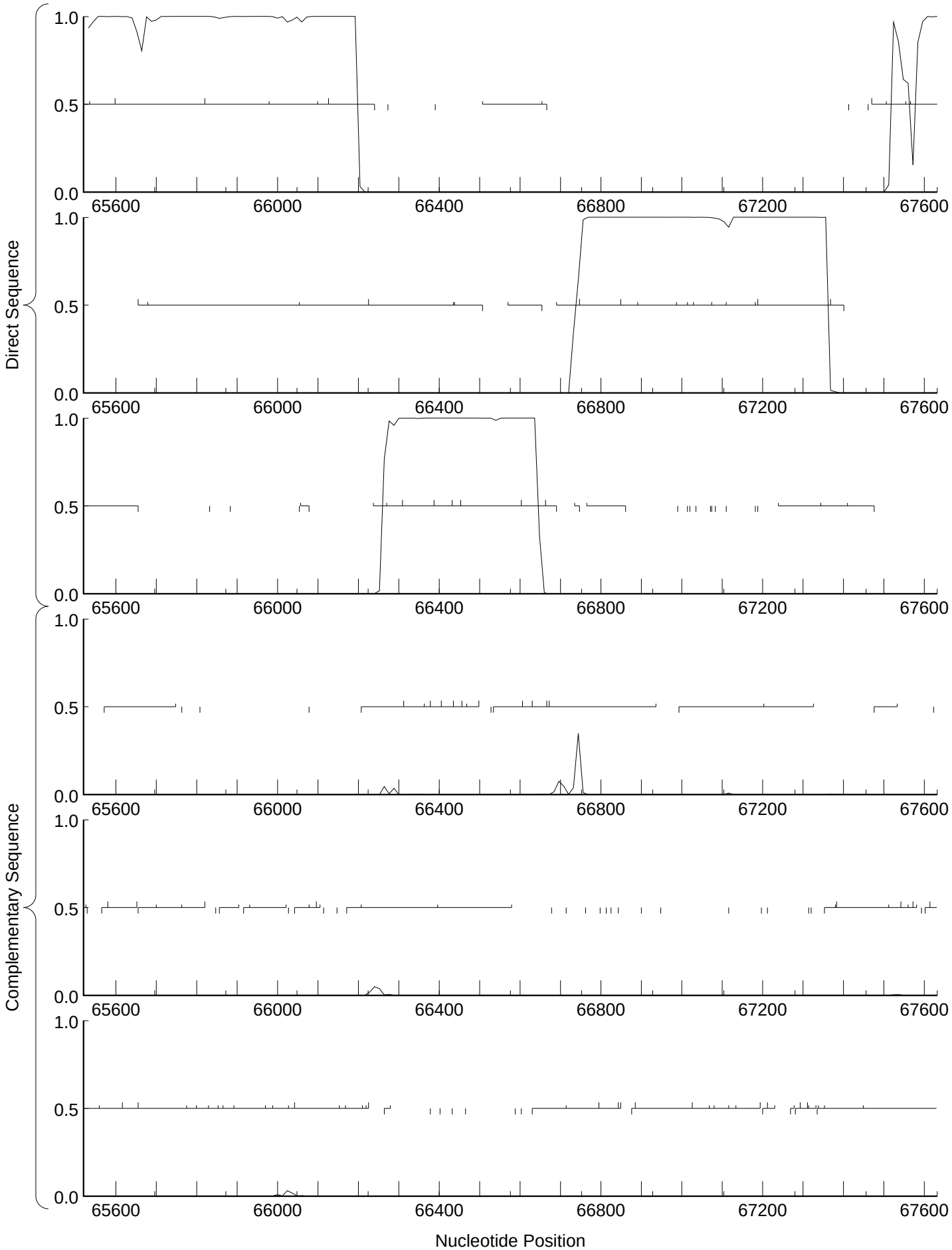
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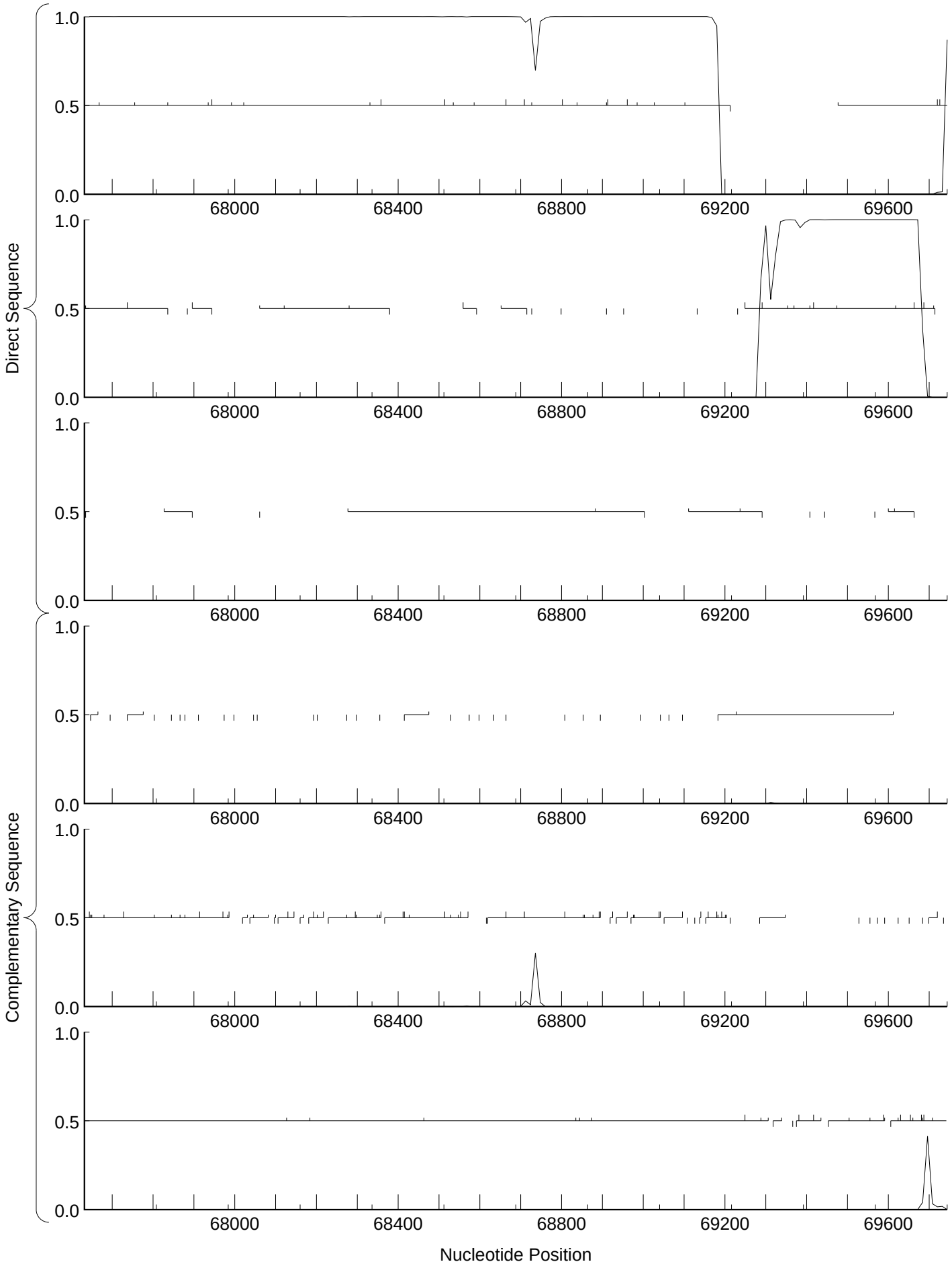
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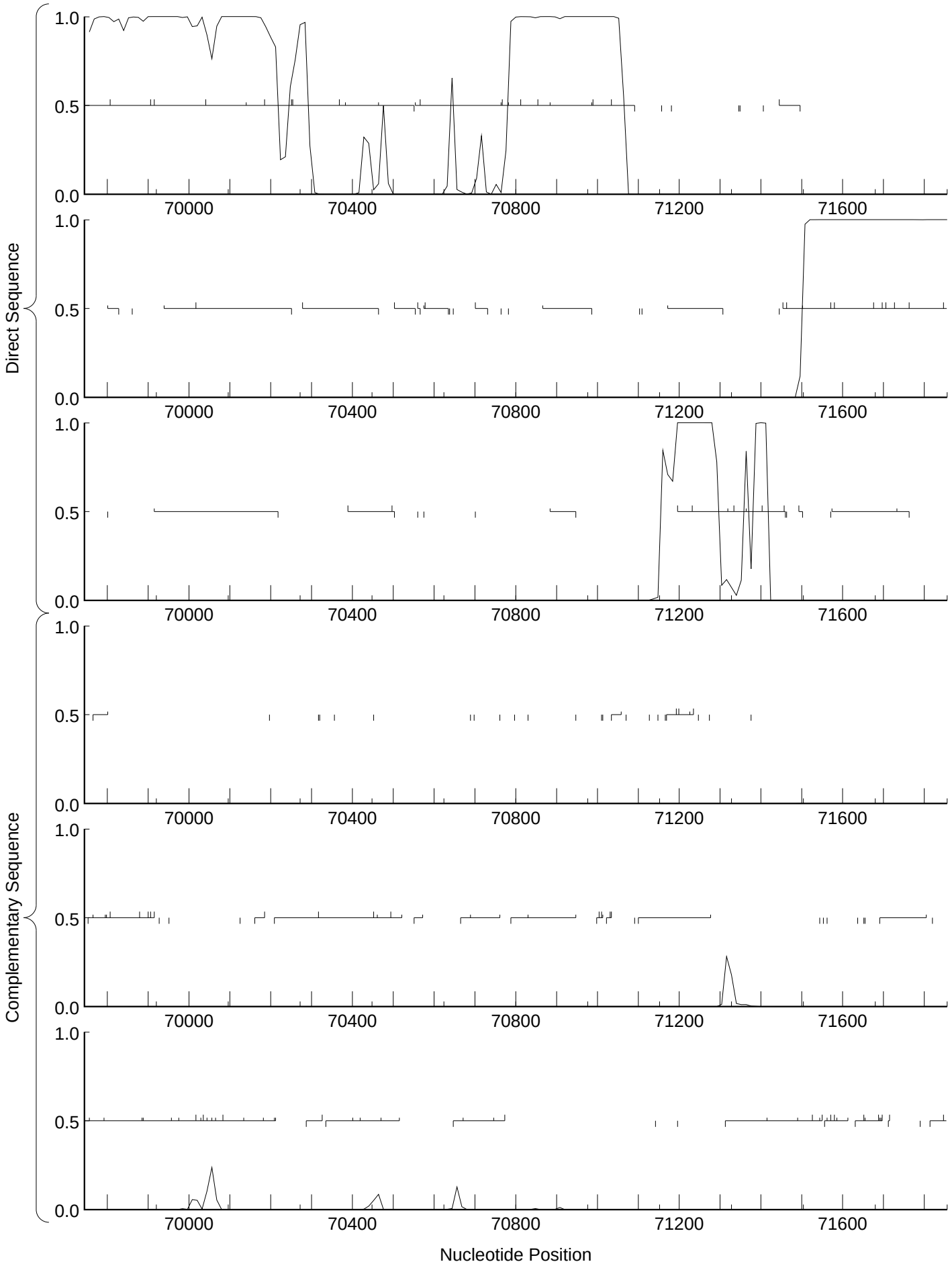
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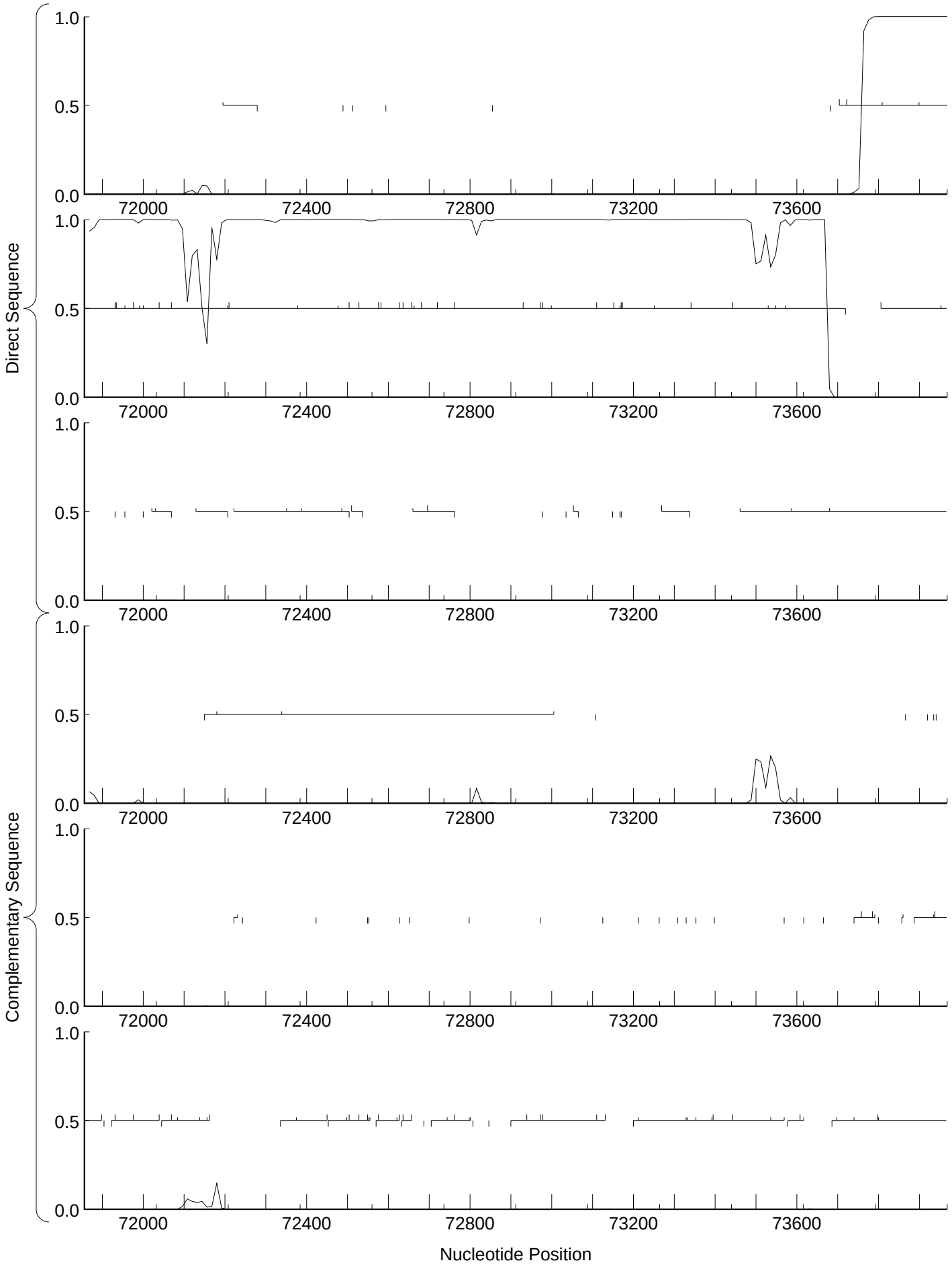
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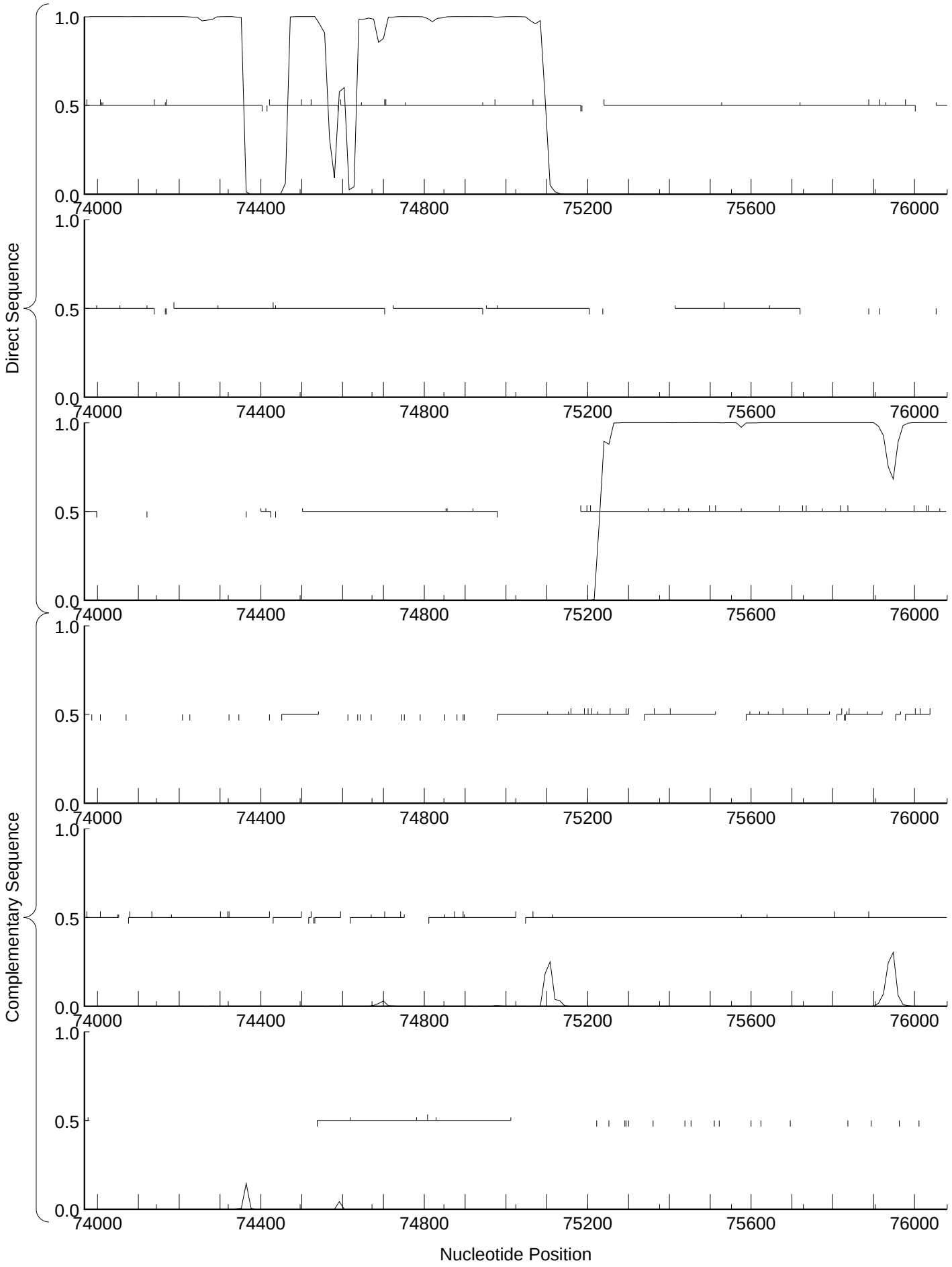
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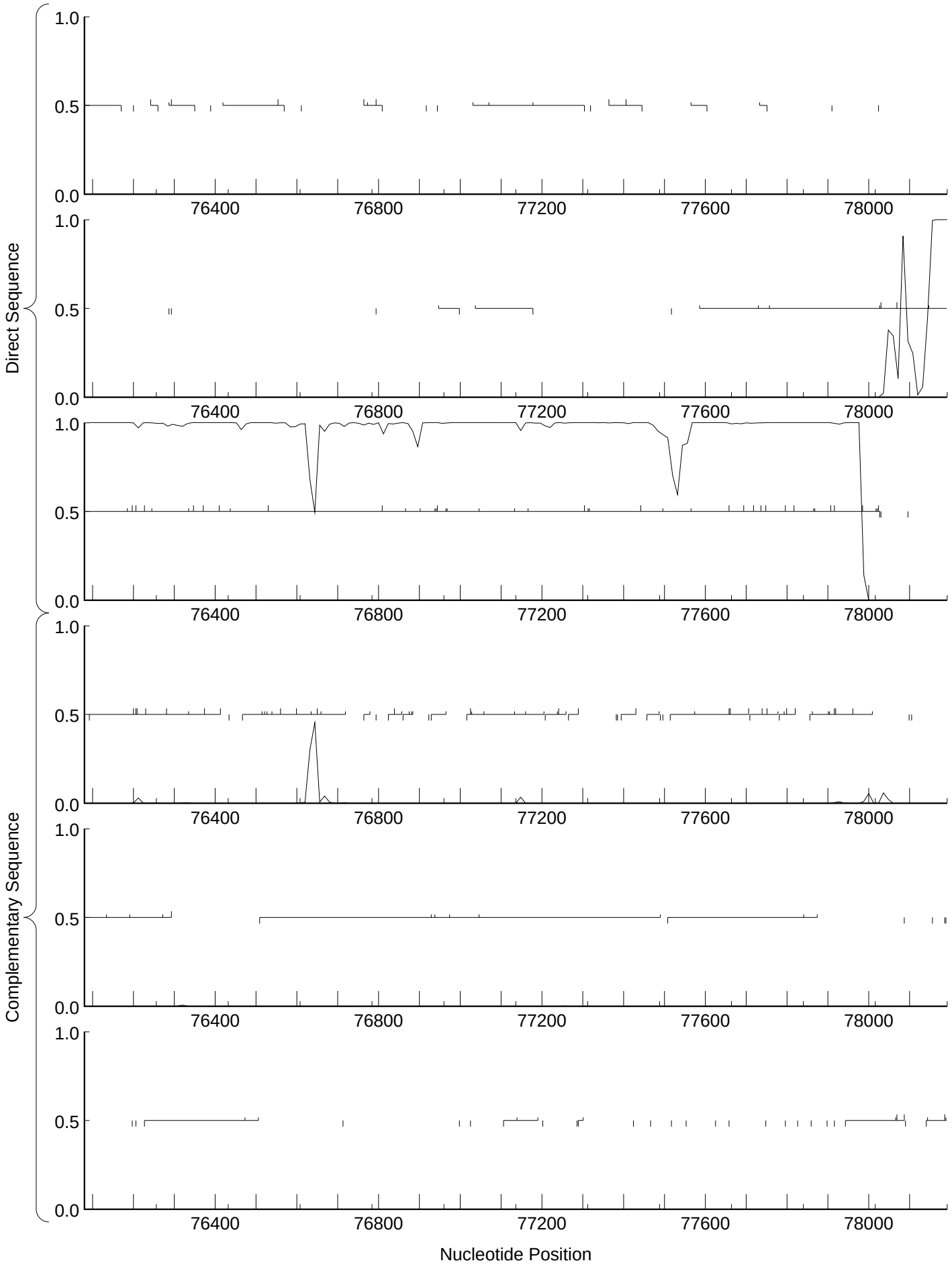
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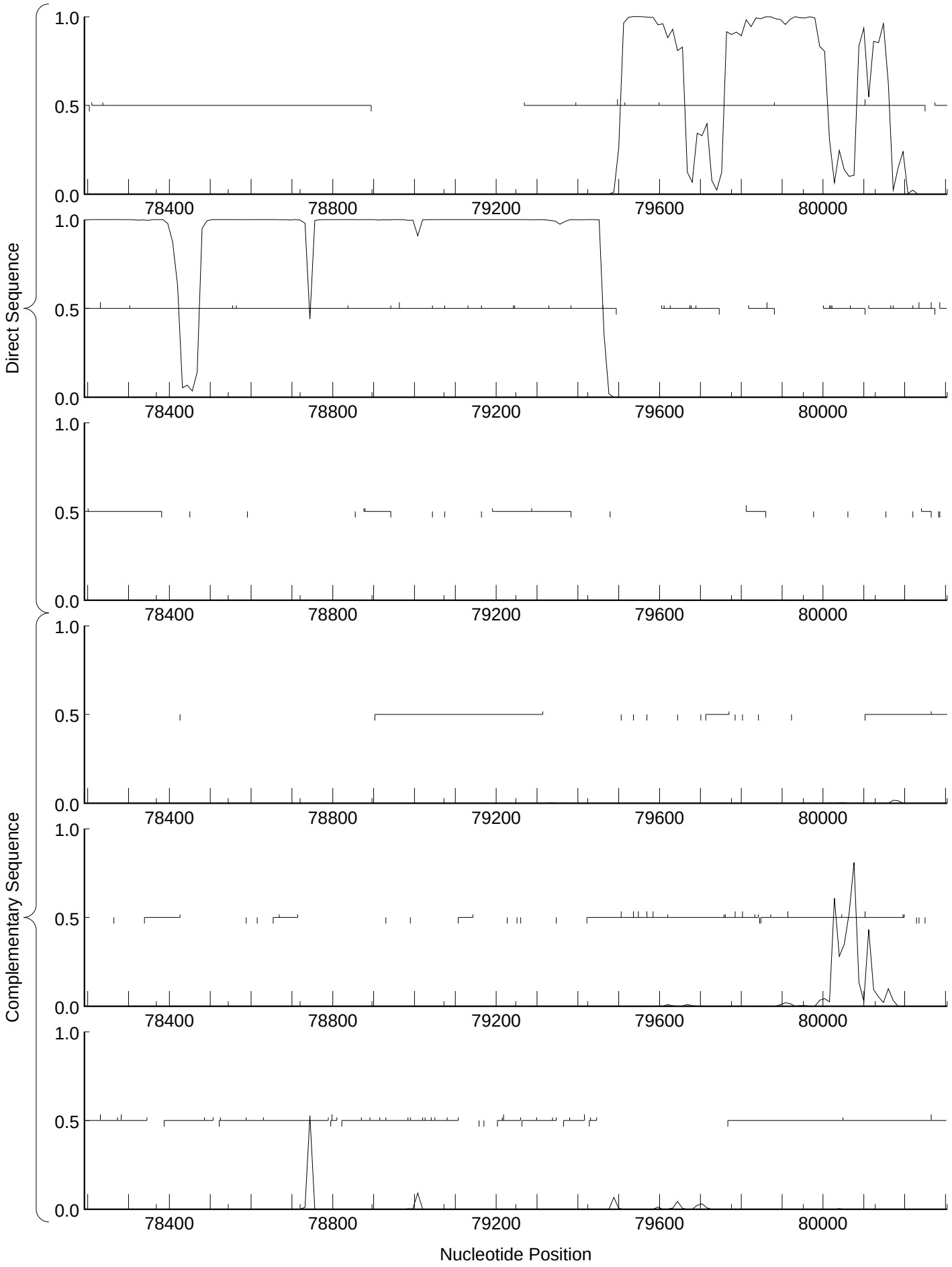
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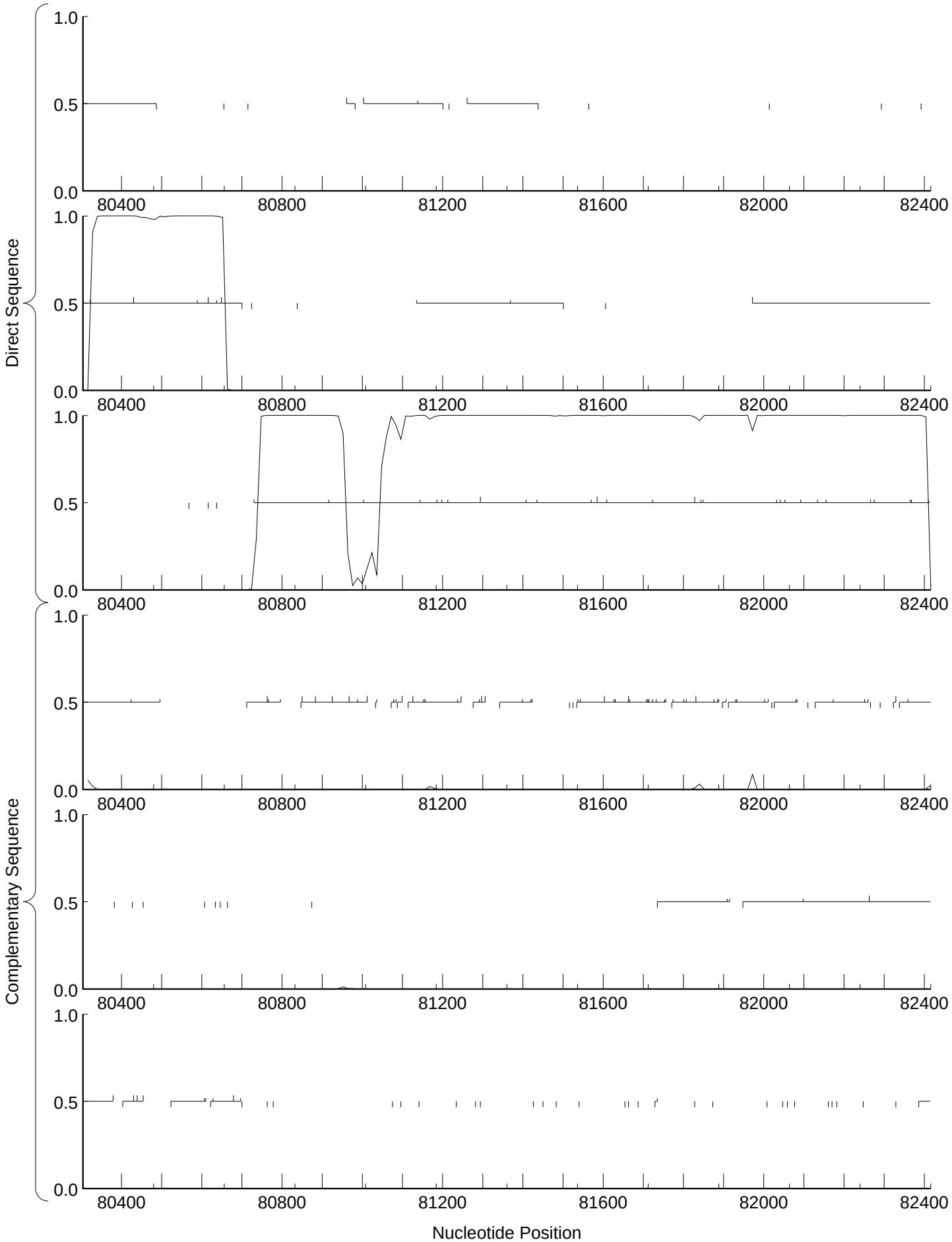
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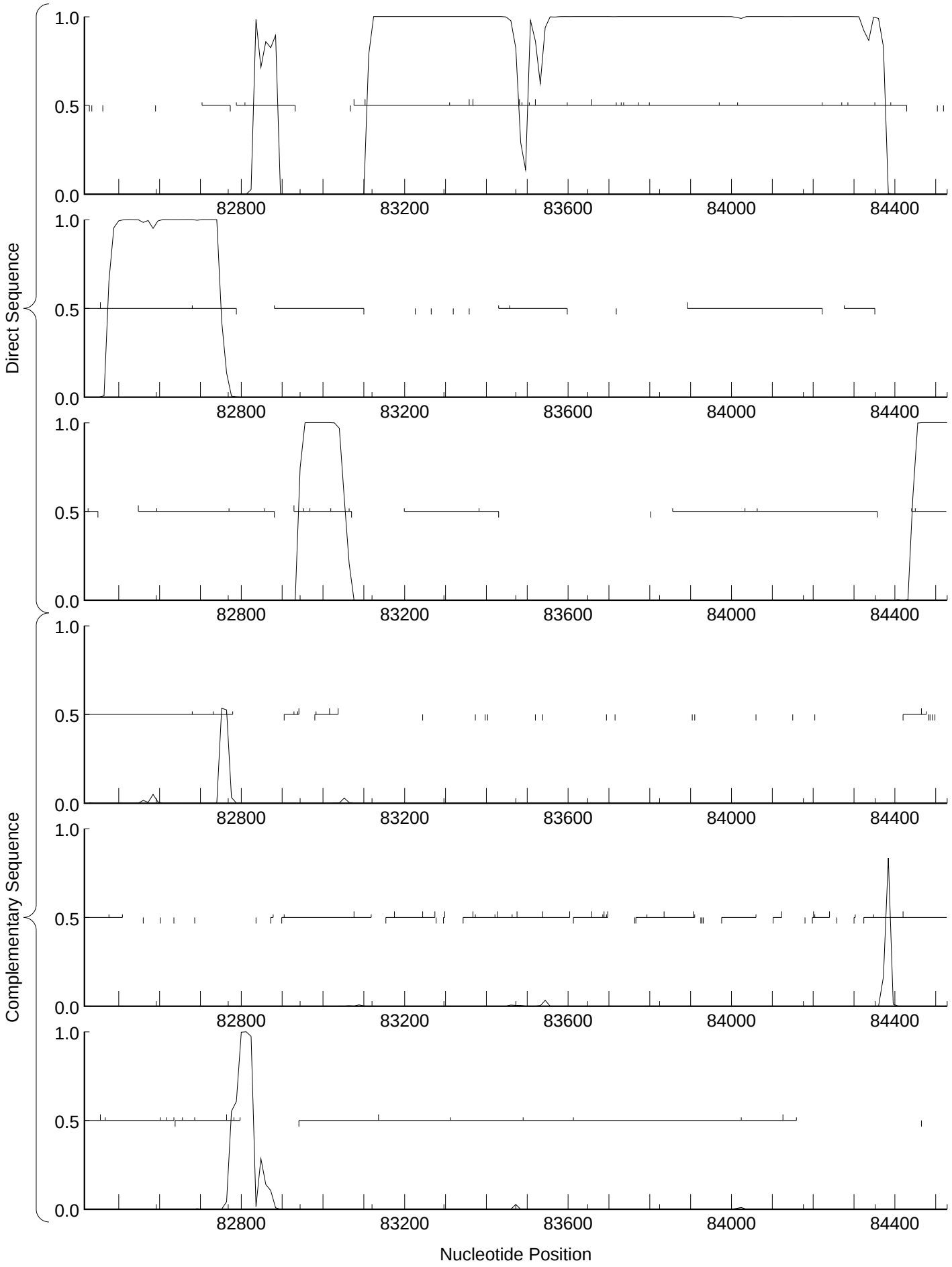
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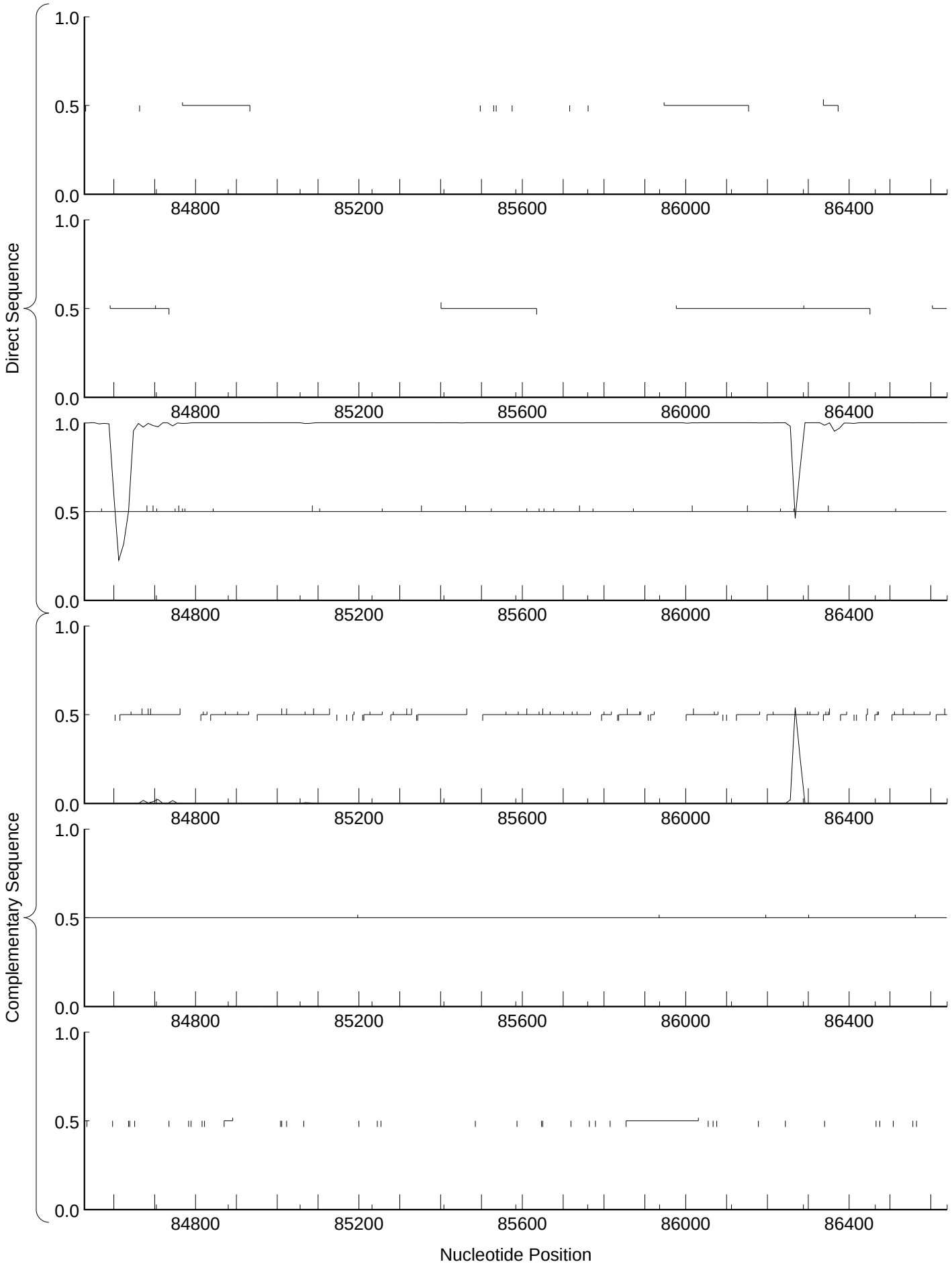
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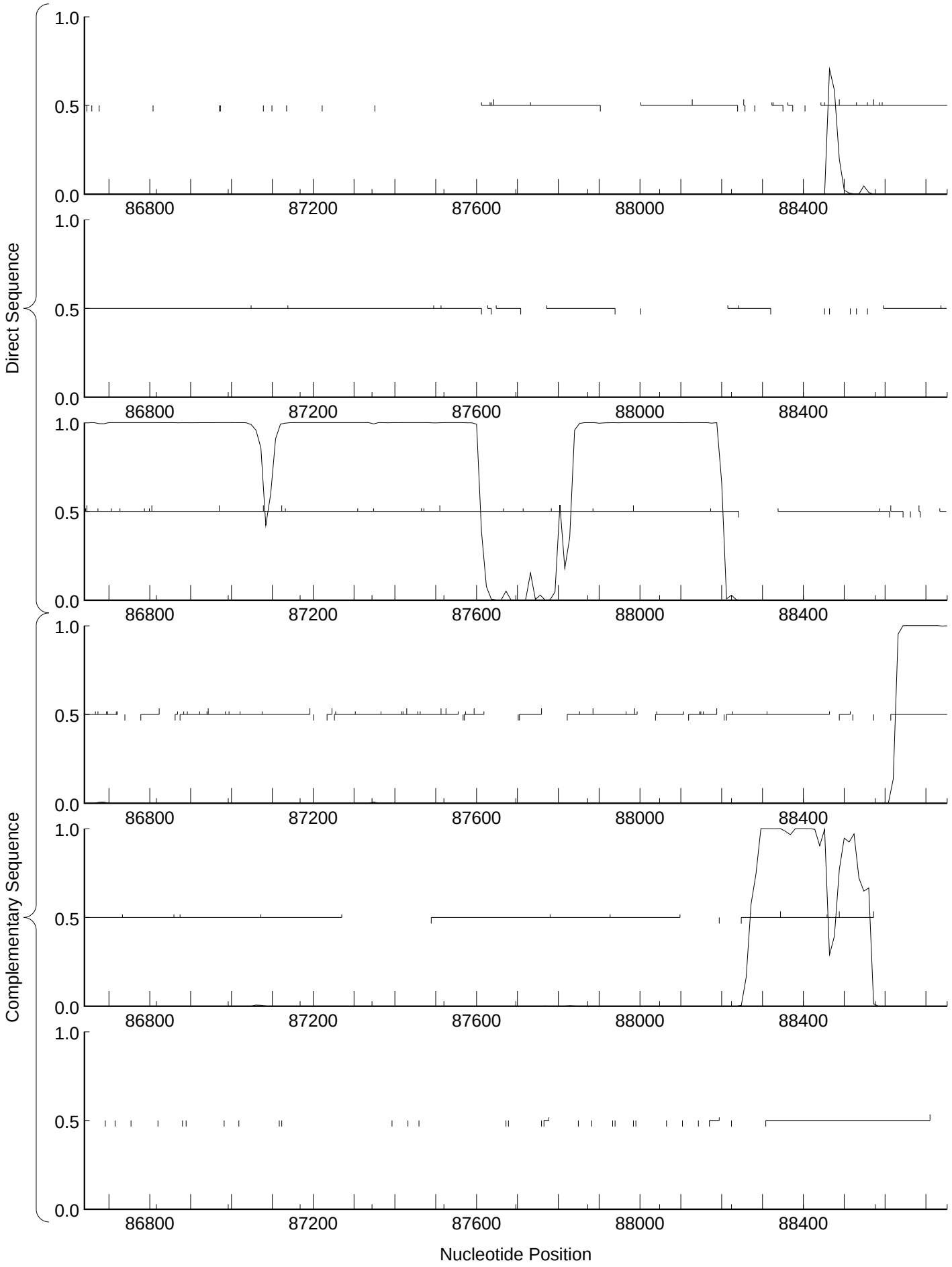
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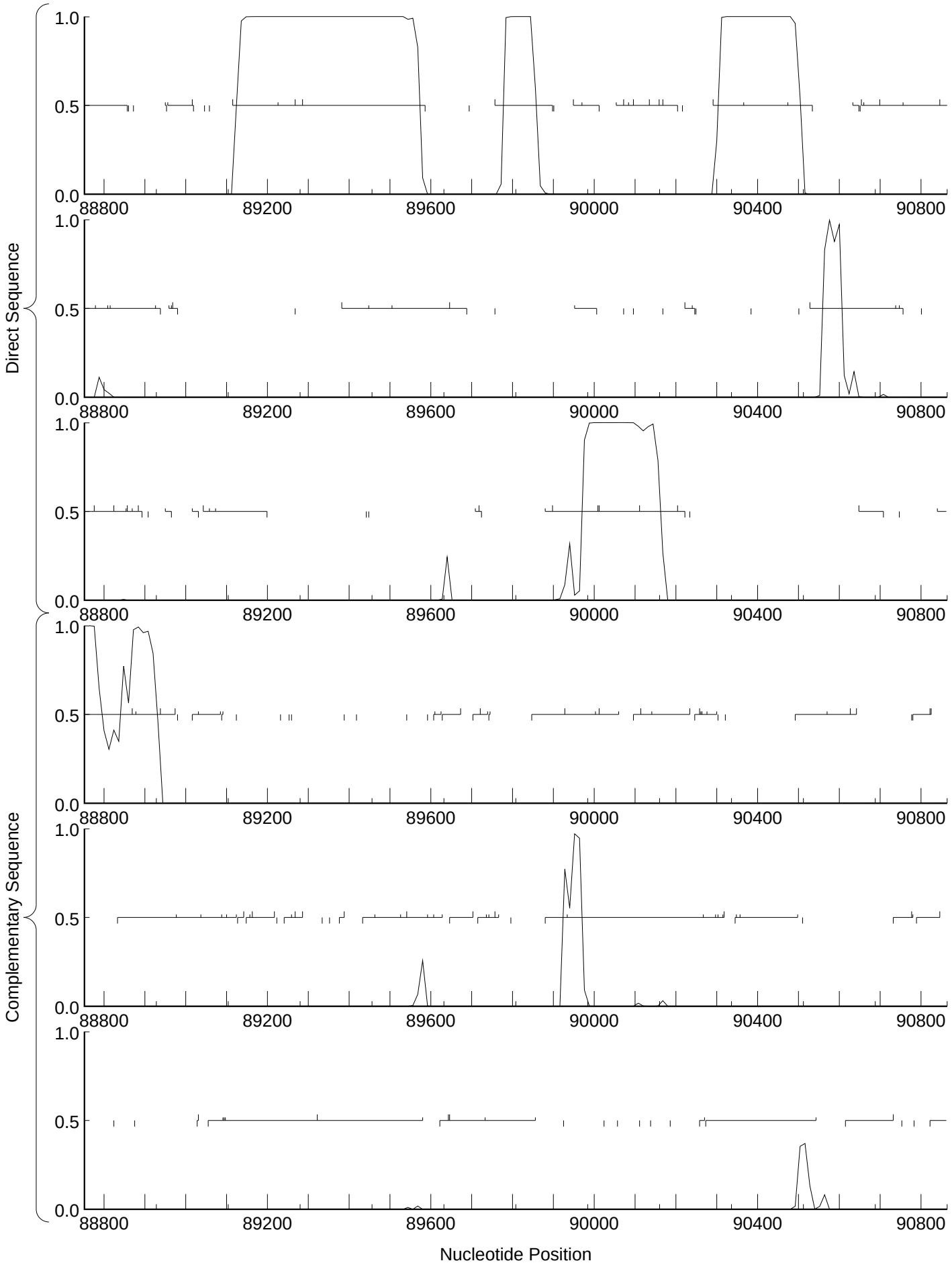


terium phage StubbySaigey complete sequence, 156308 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 42/76

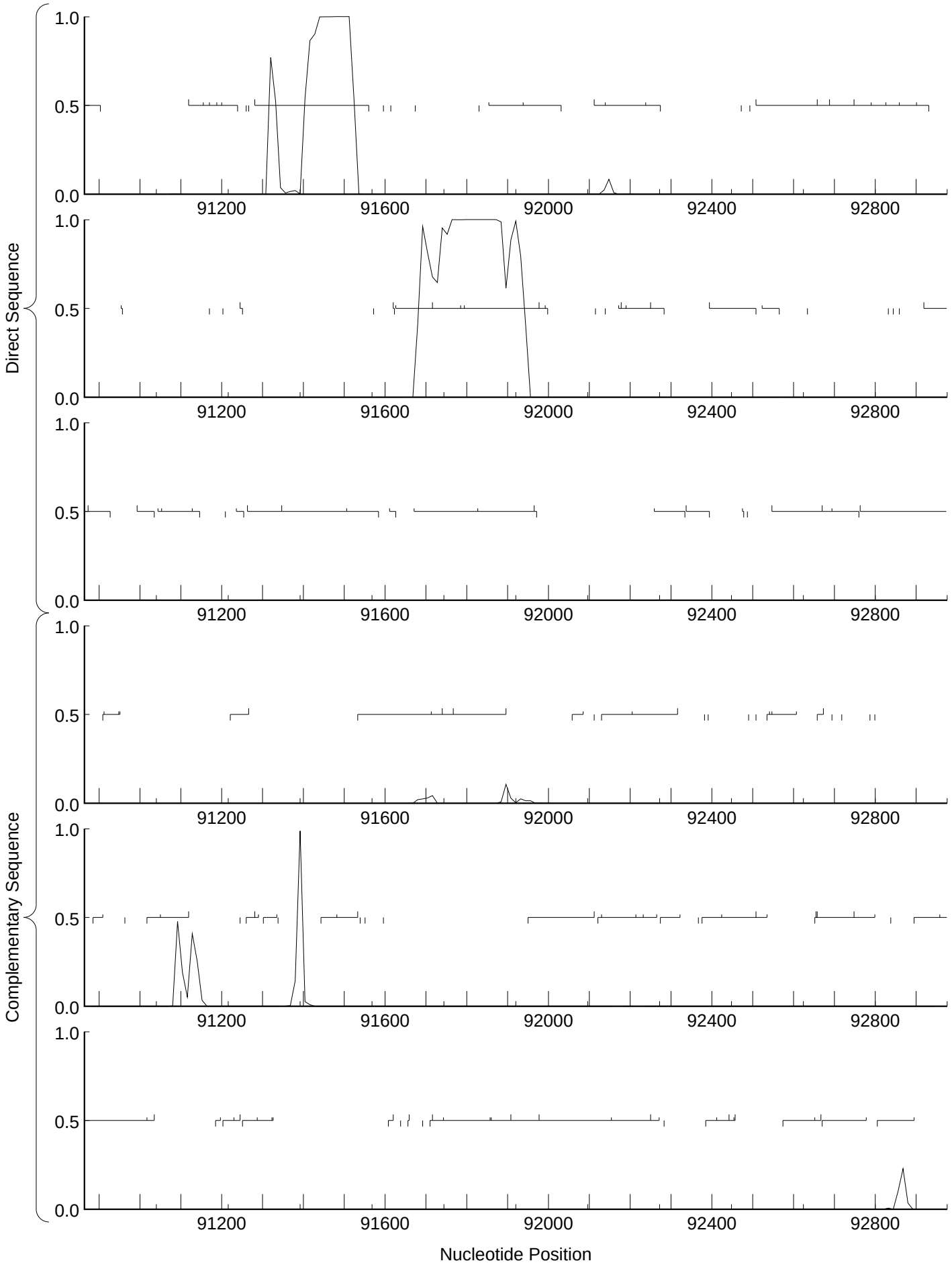


terium phage StubbySaigey complete sequence, 156308 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 43/76

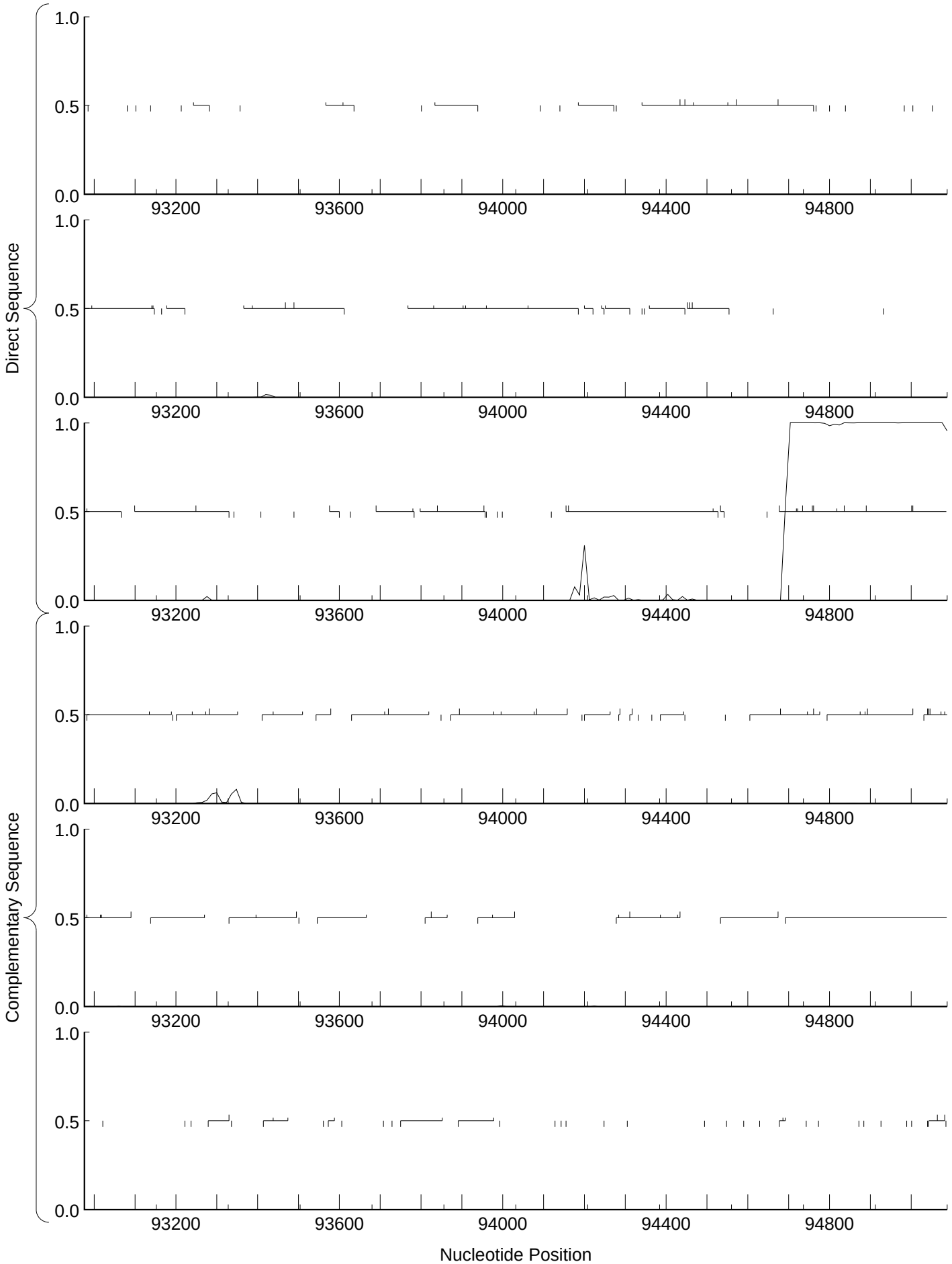



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
88800
89200
89600
90000
90400
90800
Nucleotide Position

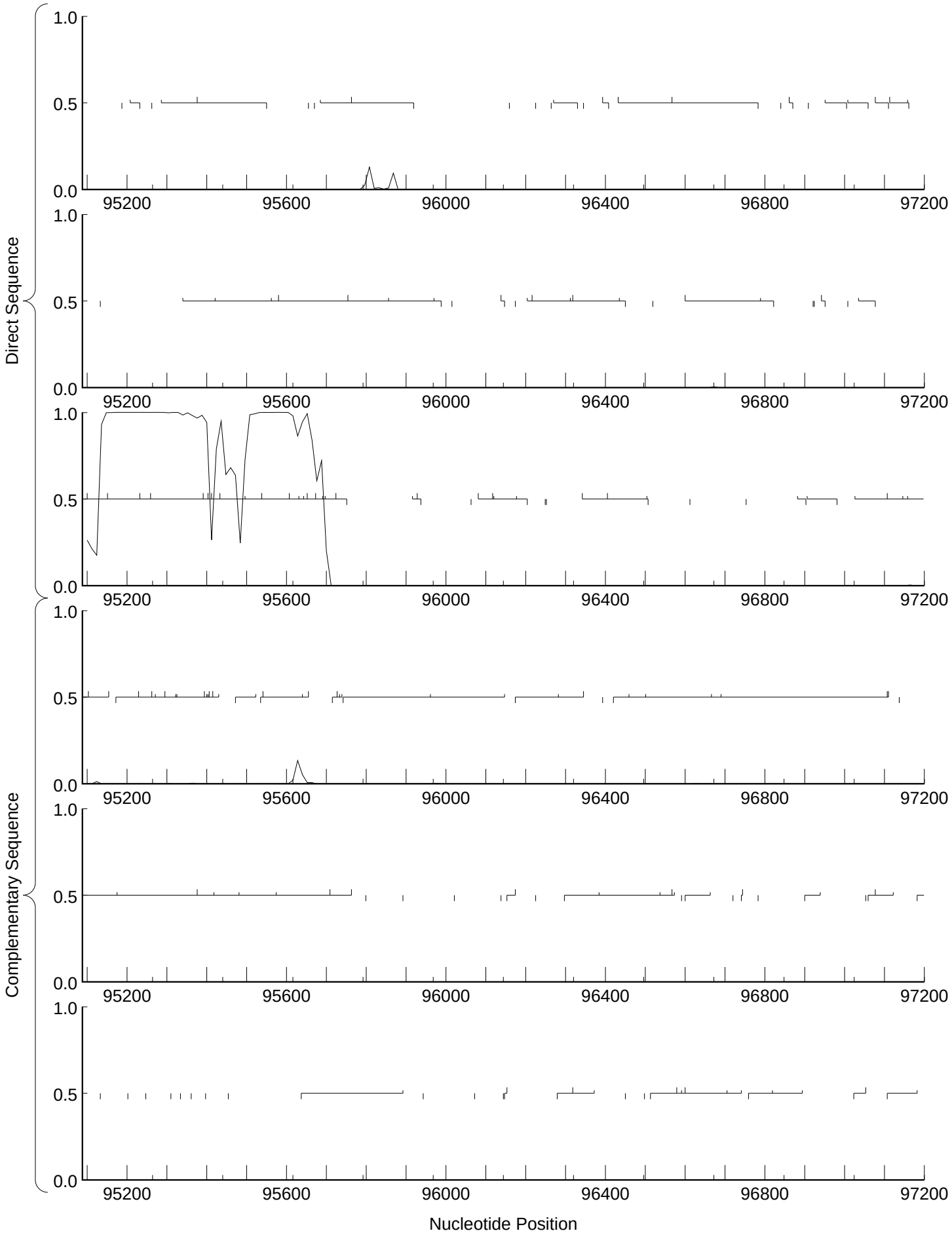
terium phage StubbySaigey complete sequence, 156308 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 45/76



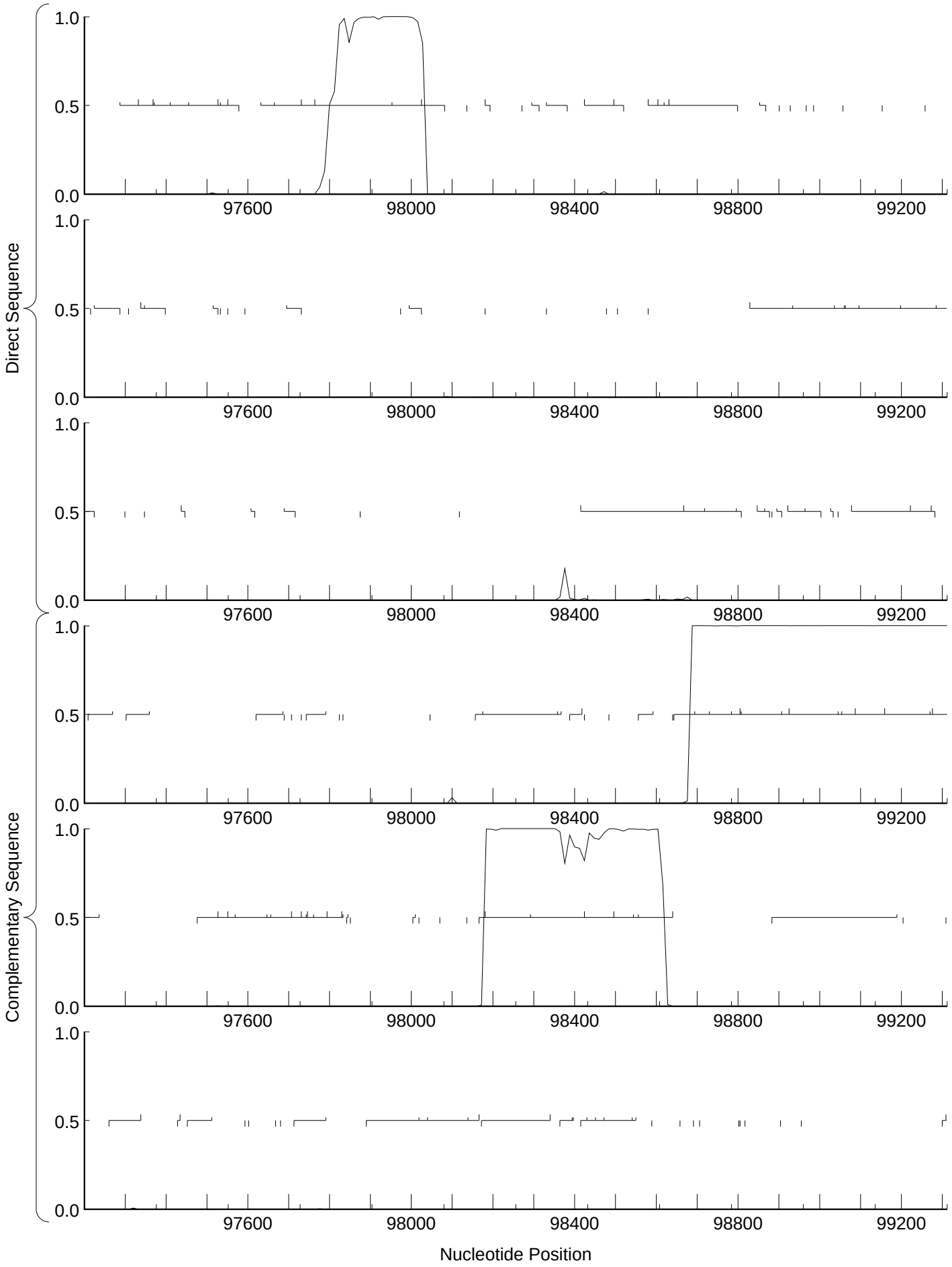
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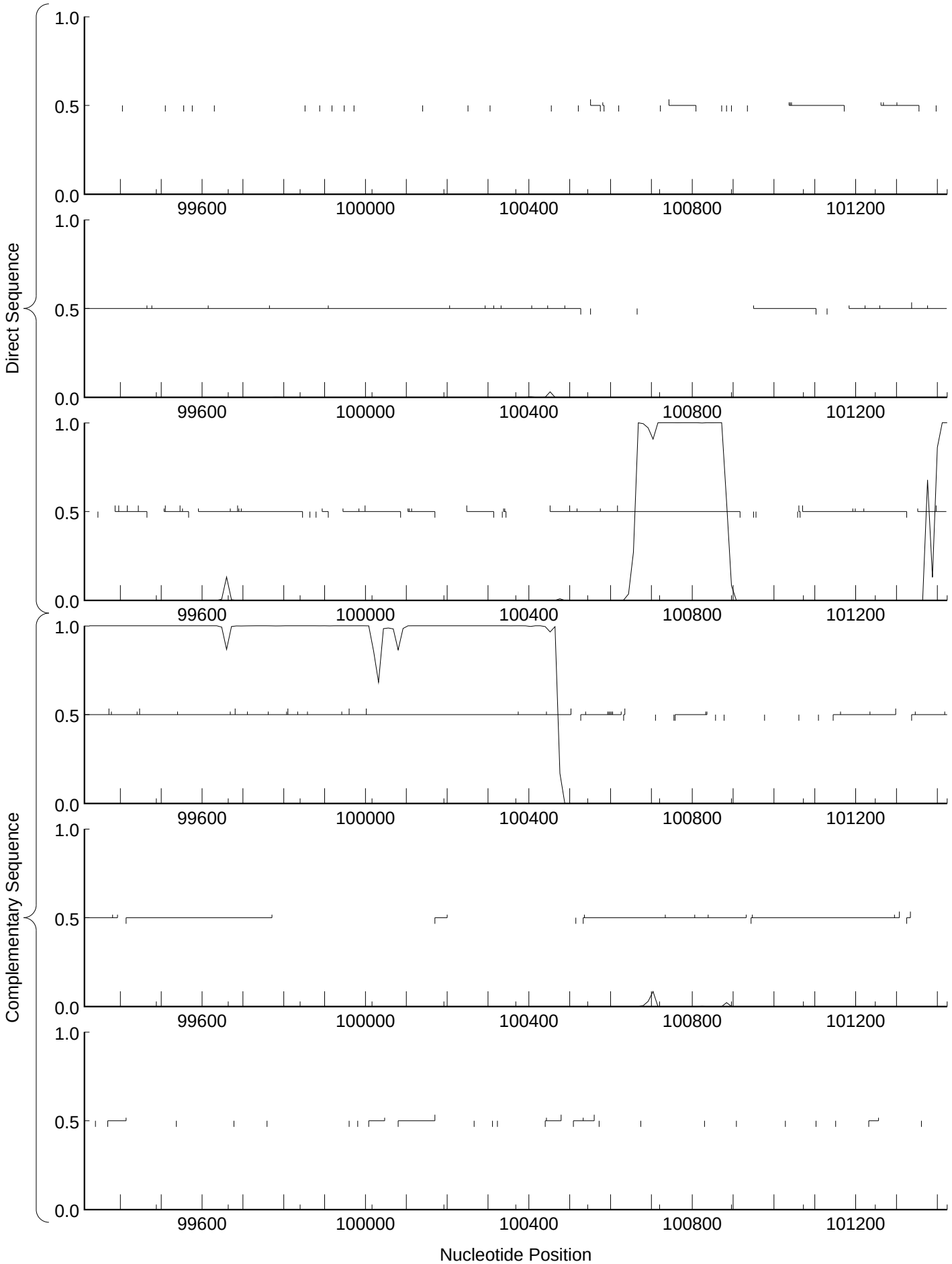
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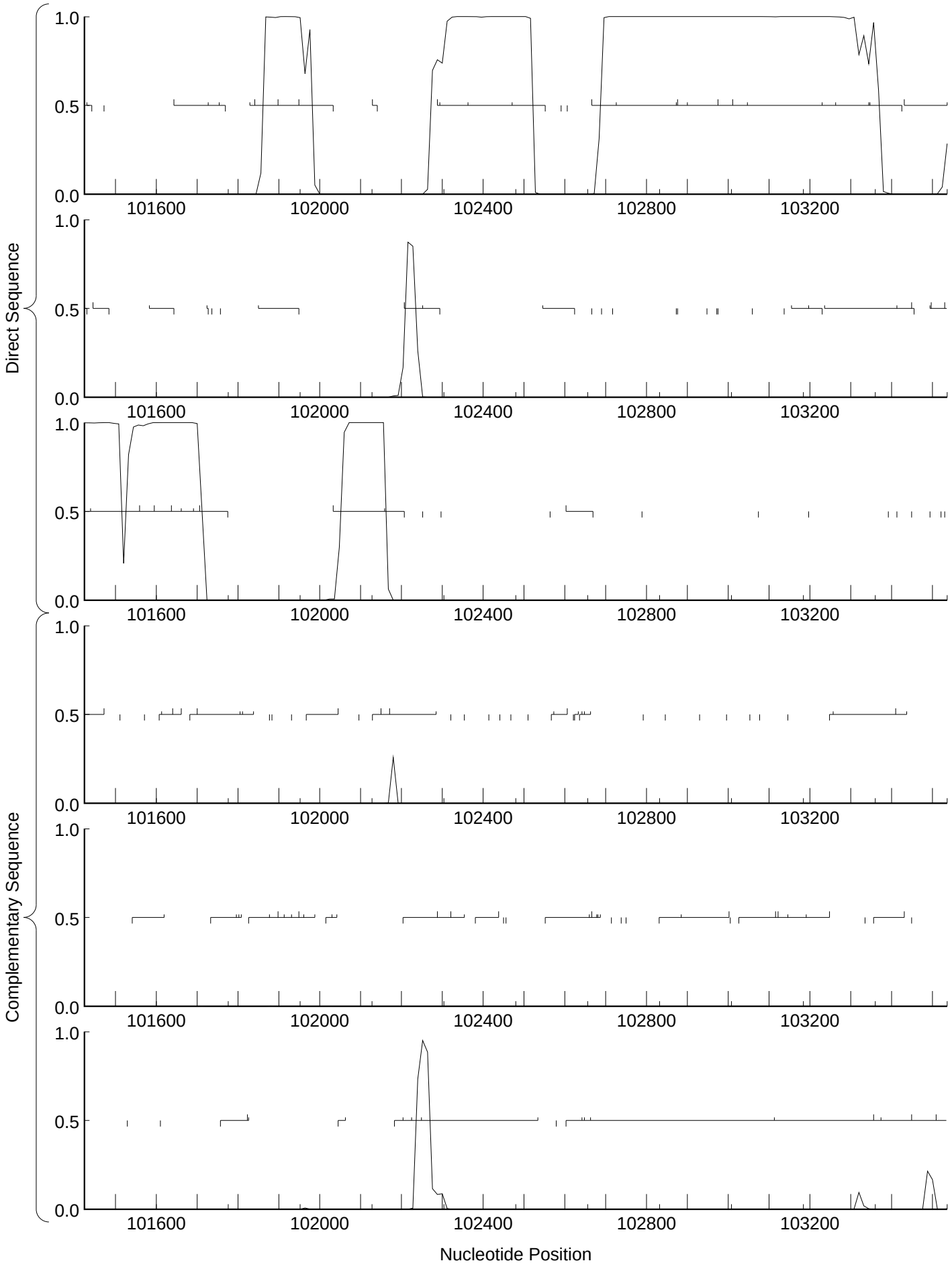
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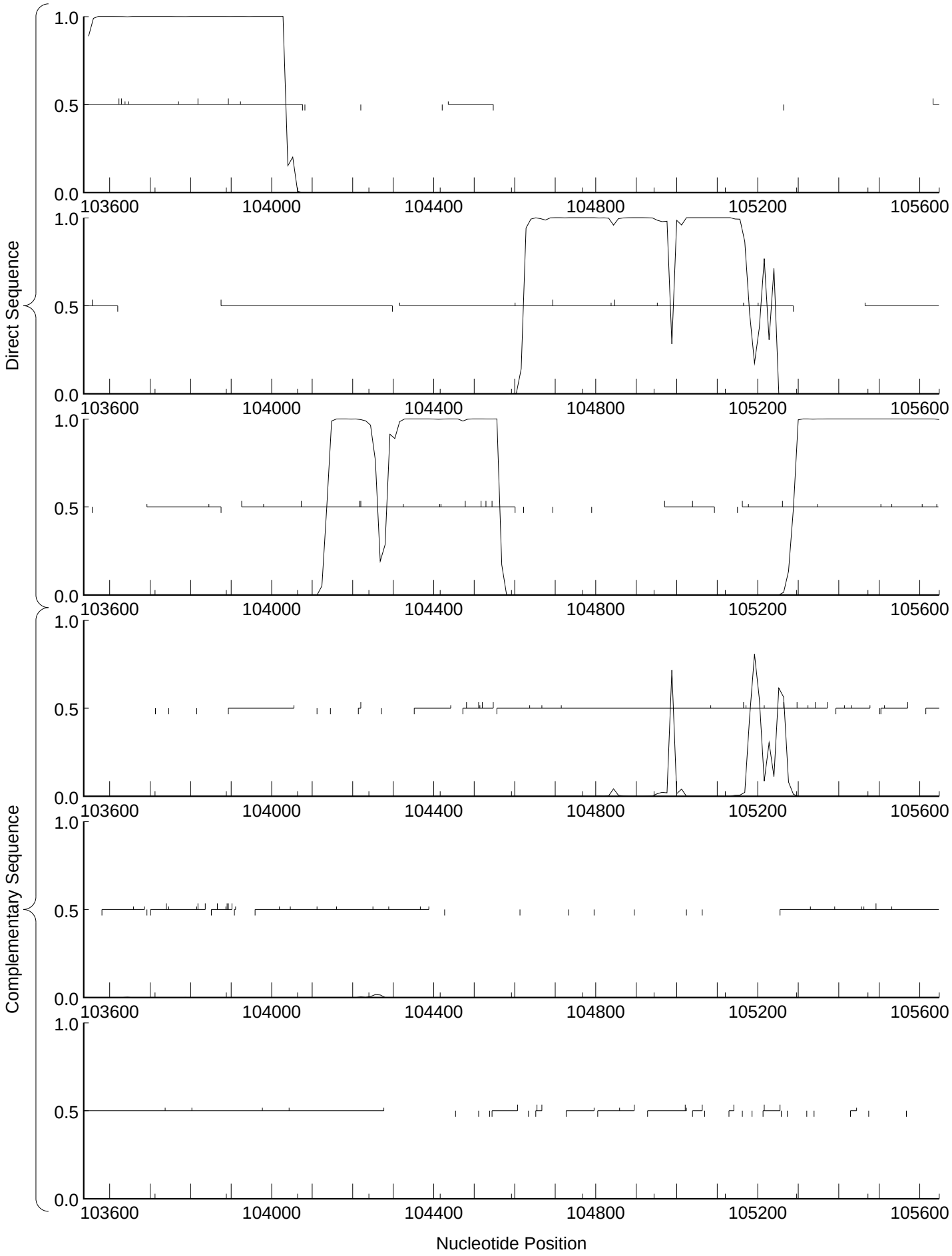
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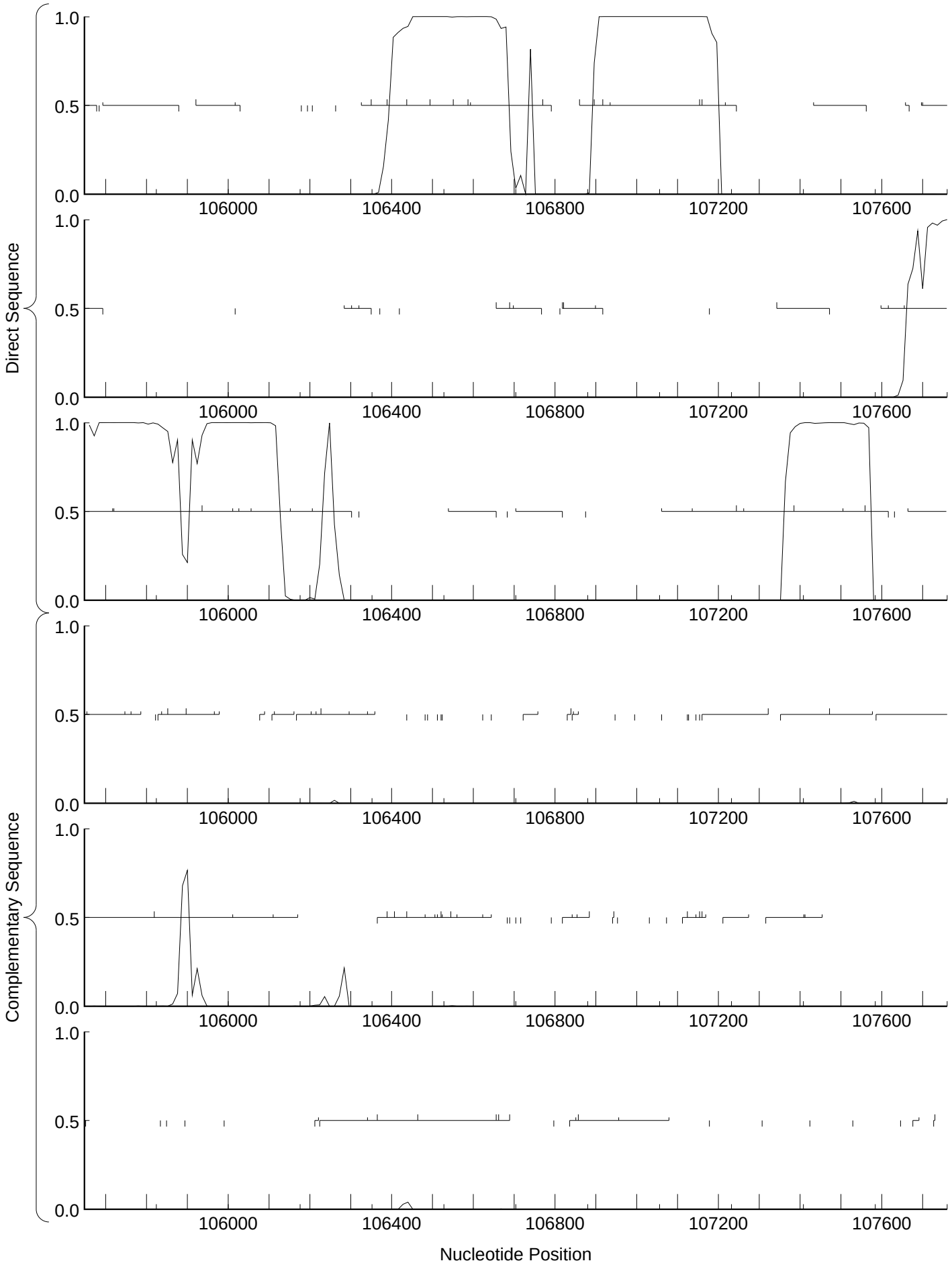
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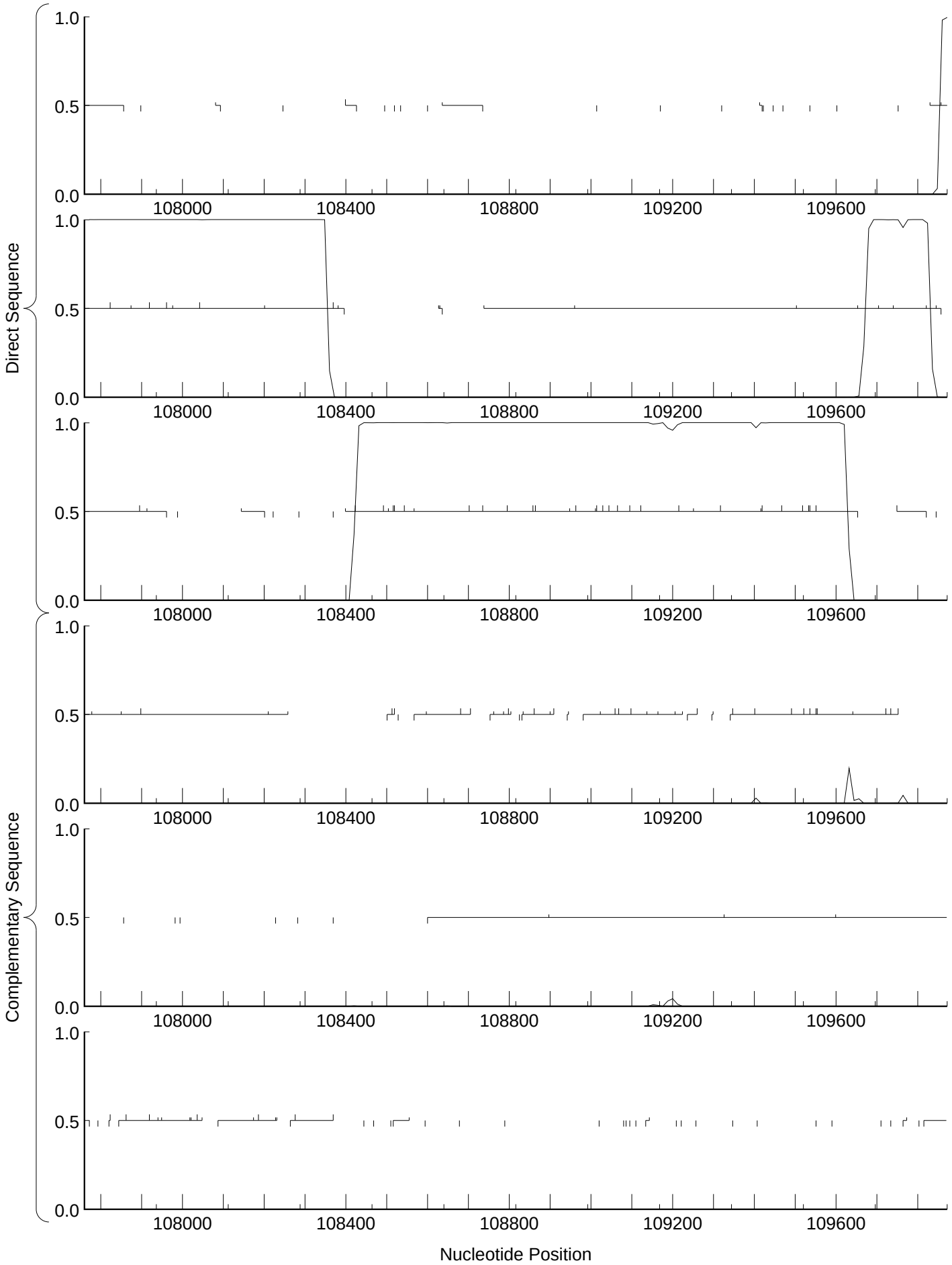
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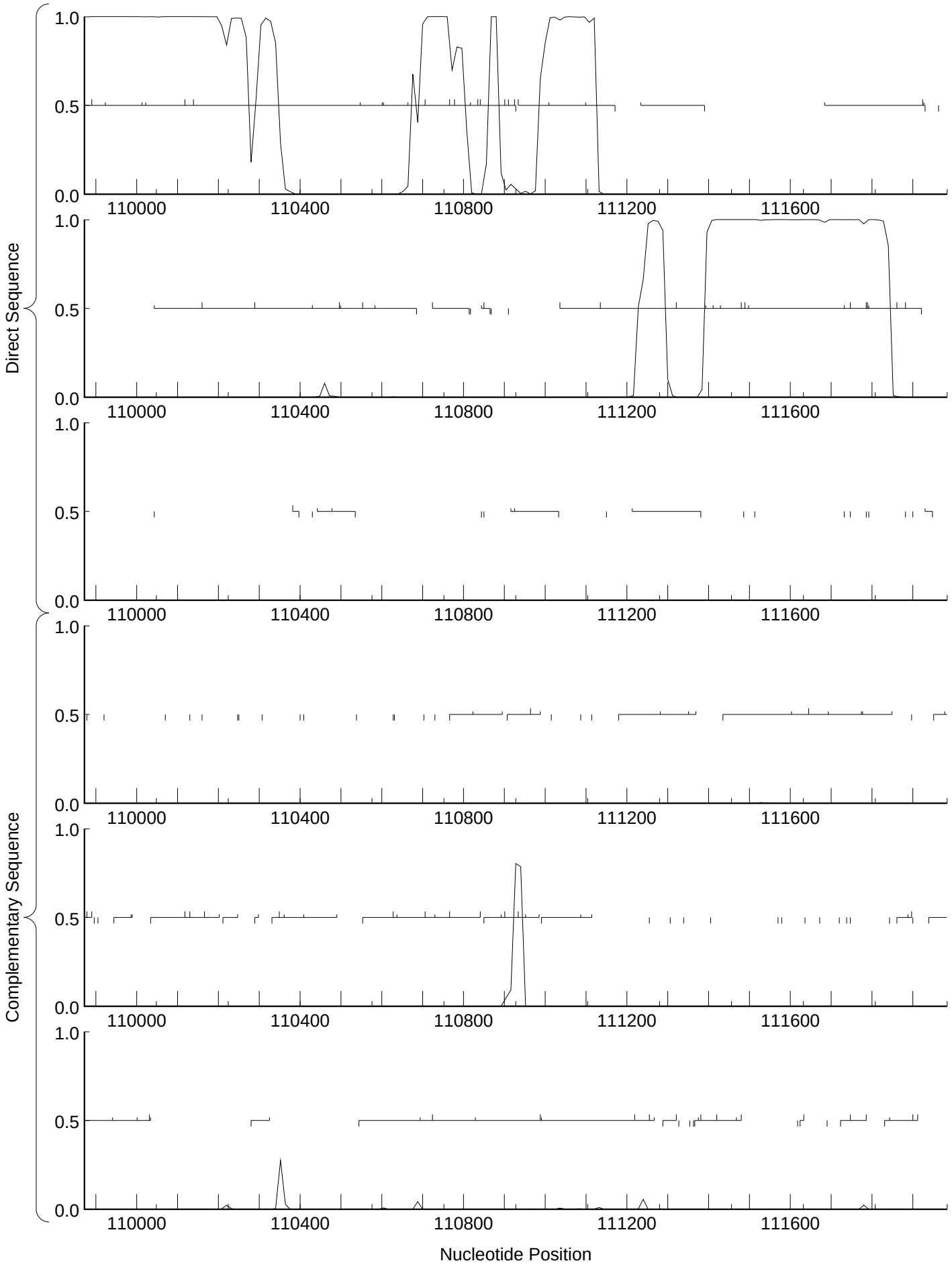
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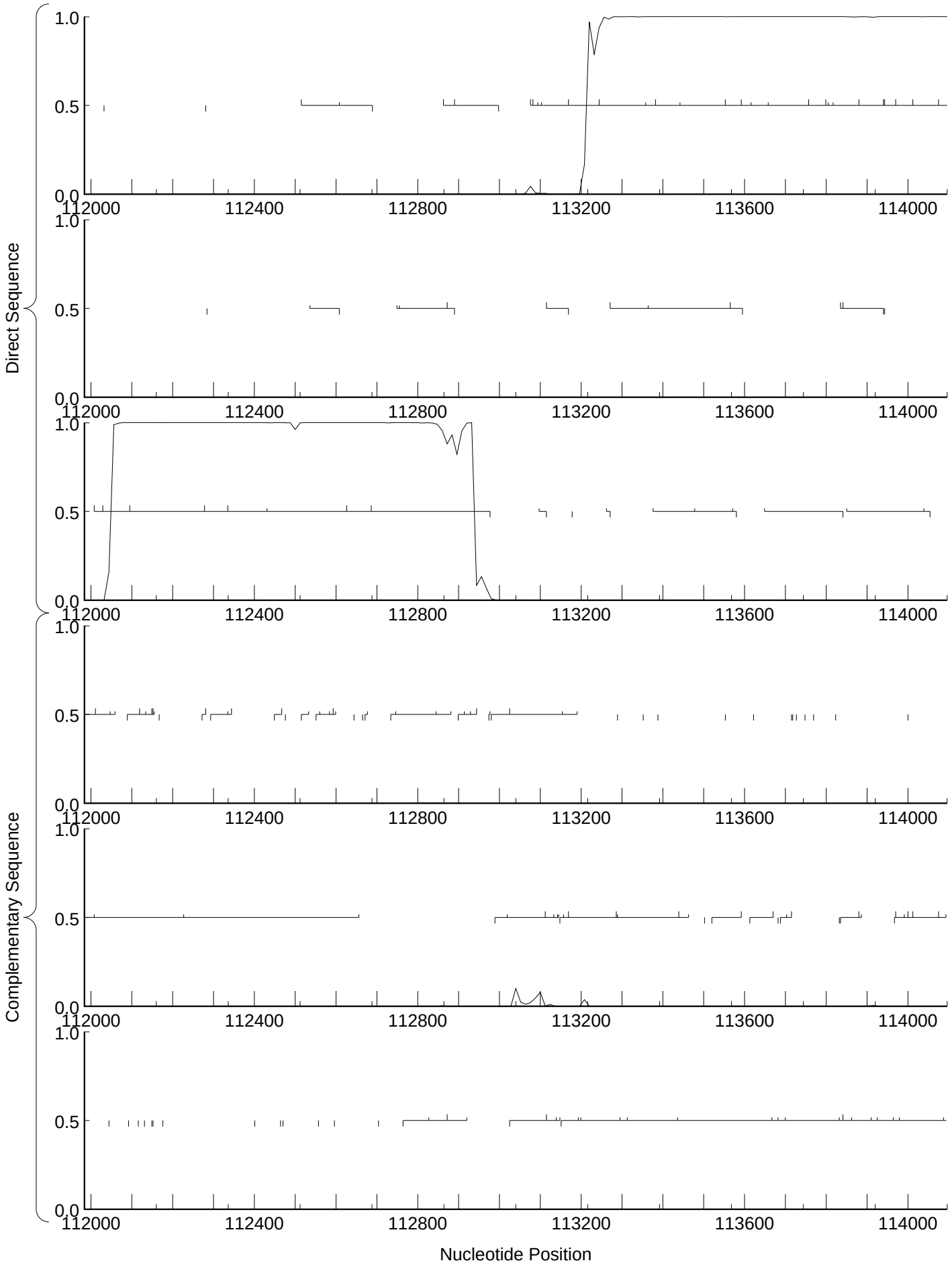
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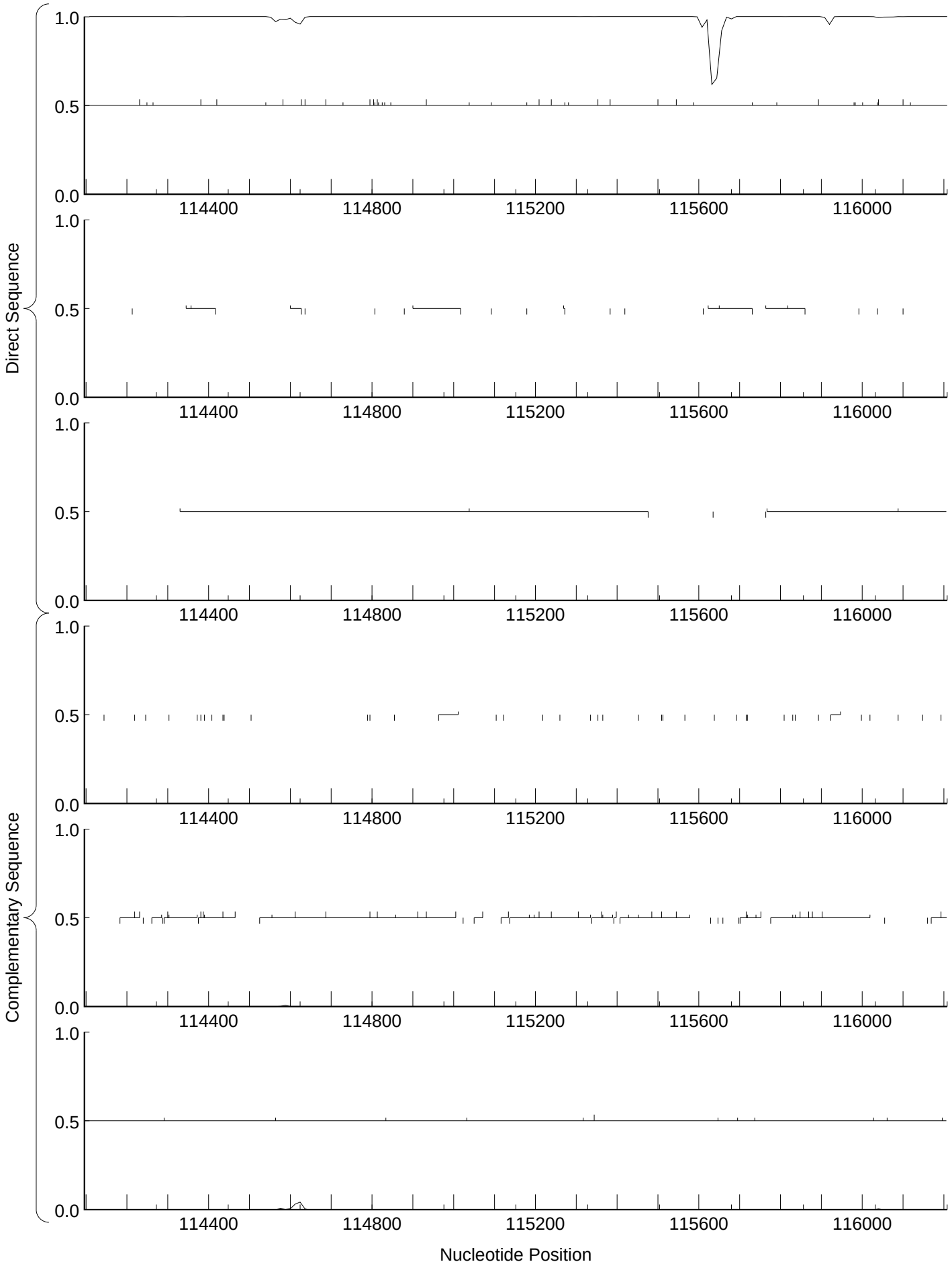
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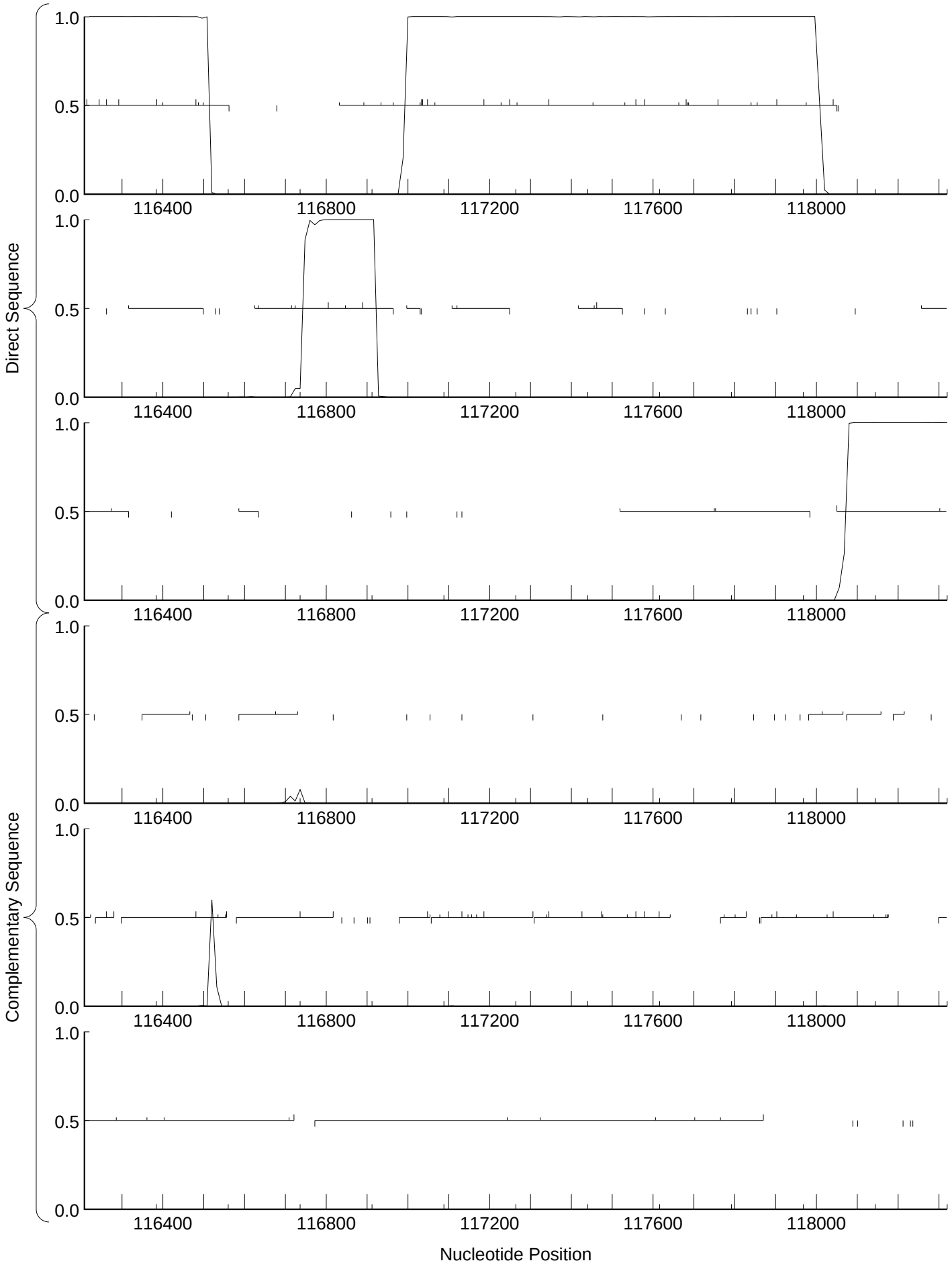
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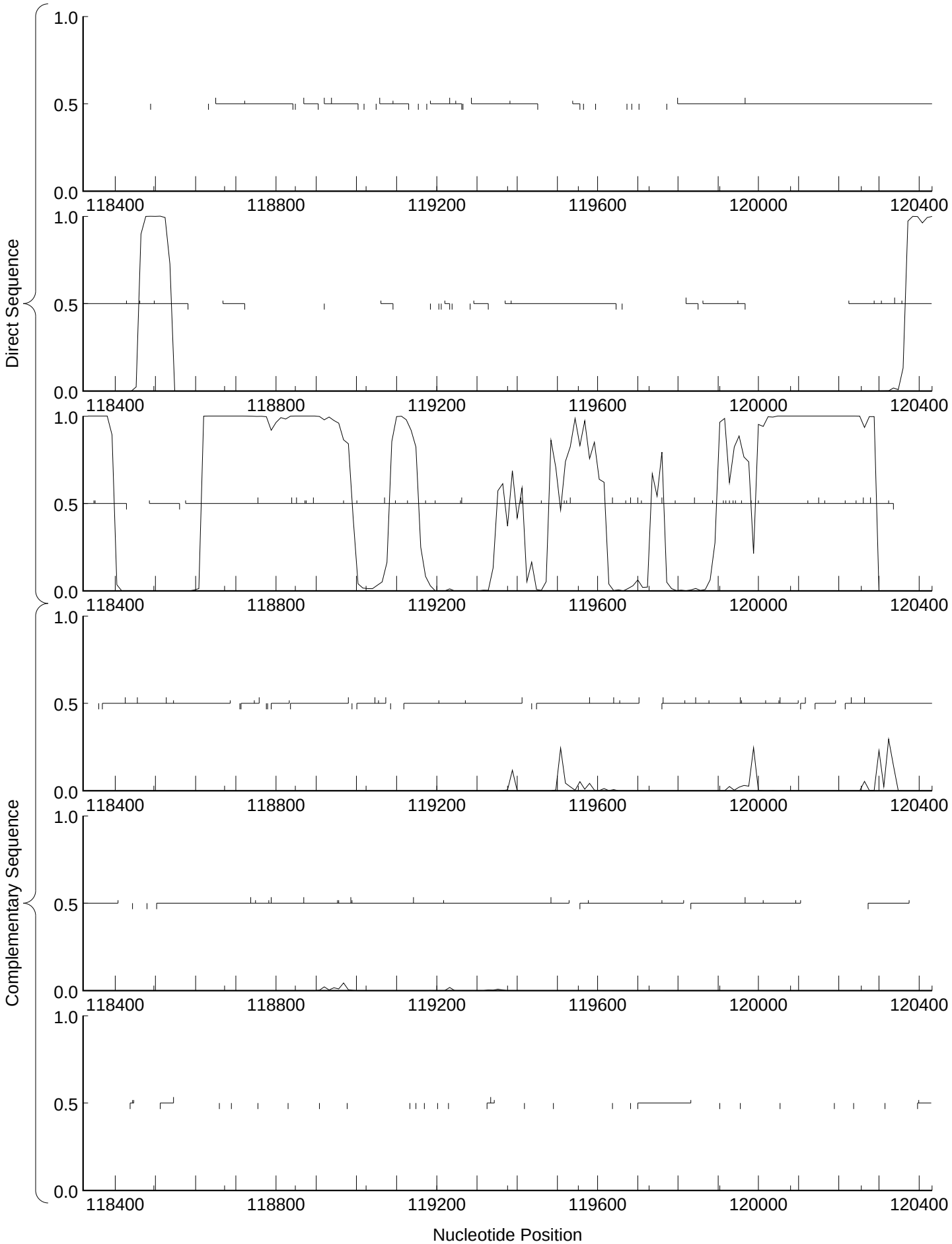
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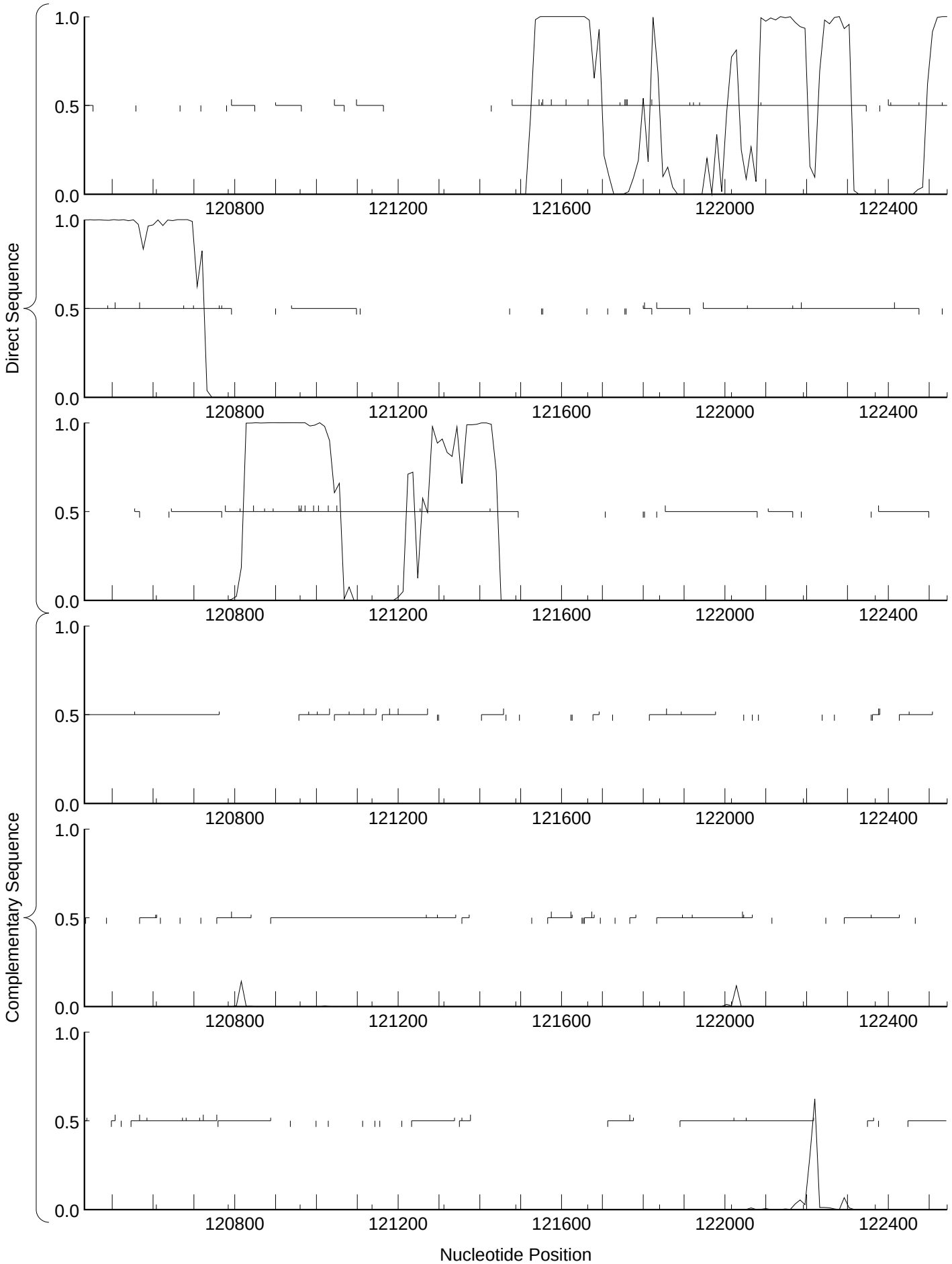
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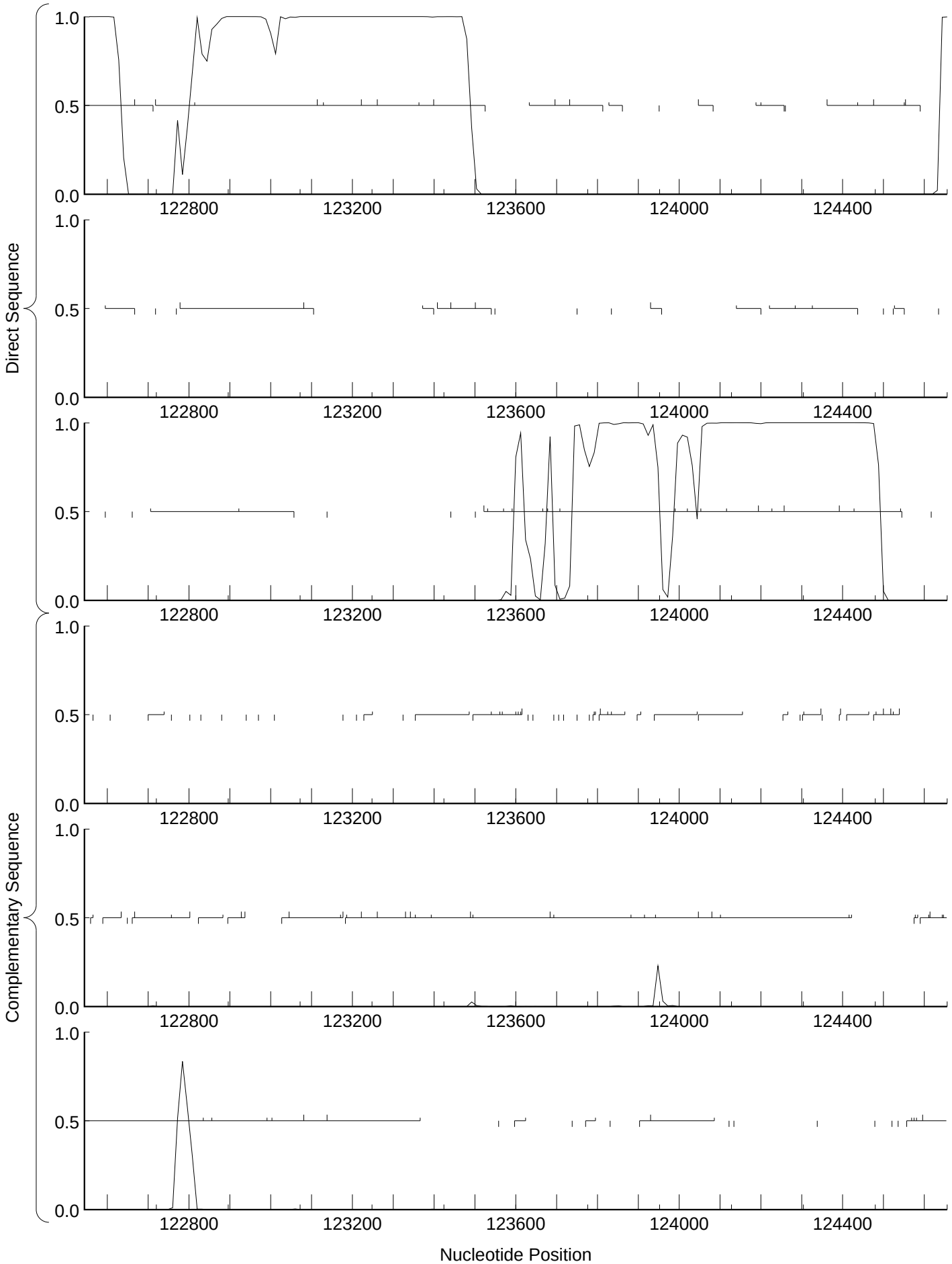
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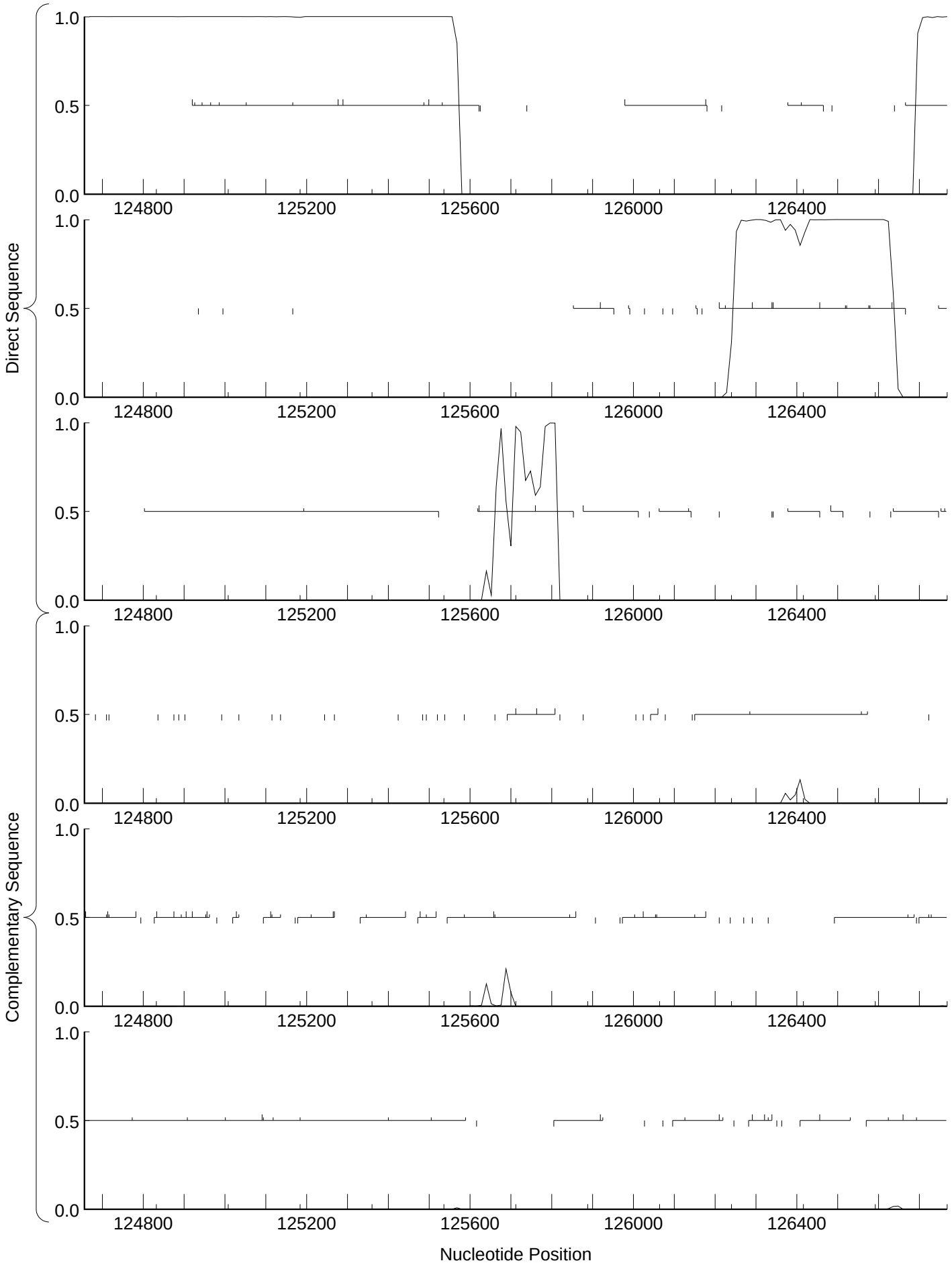
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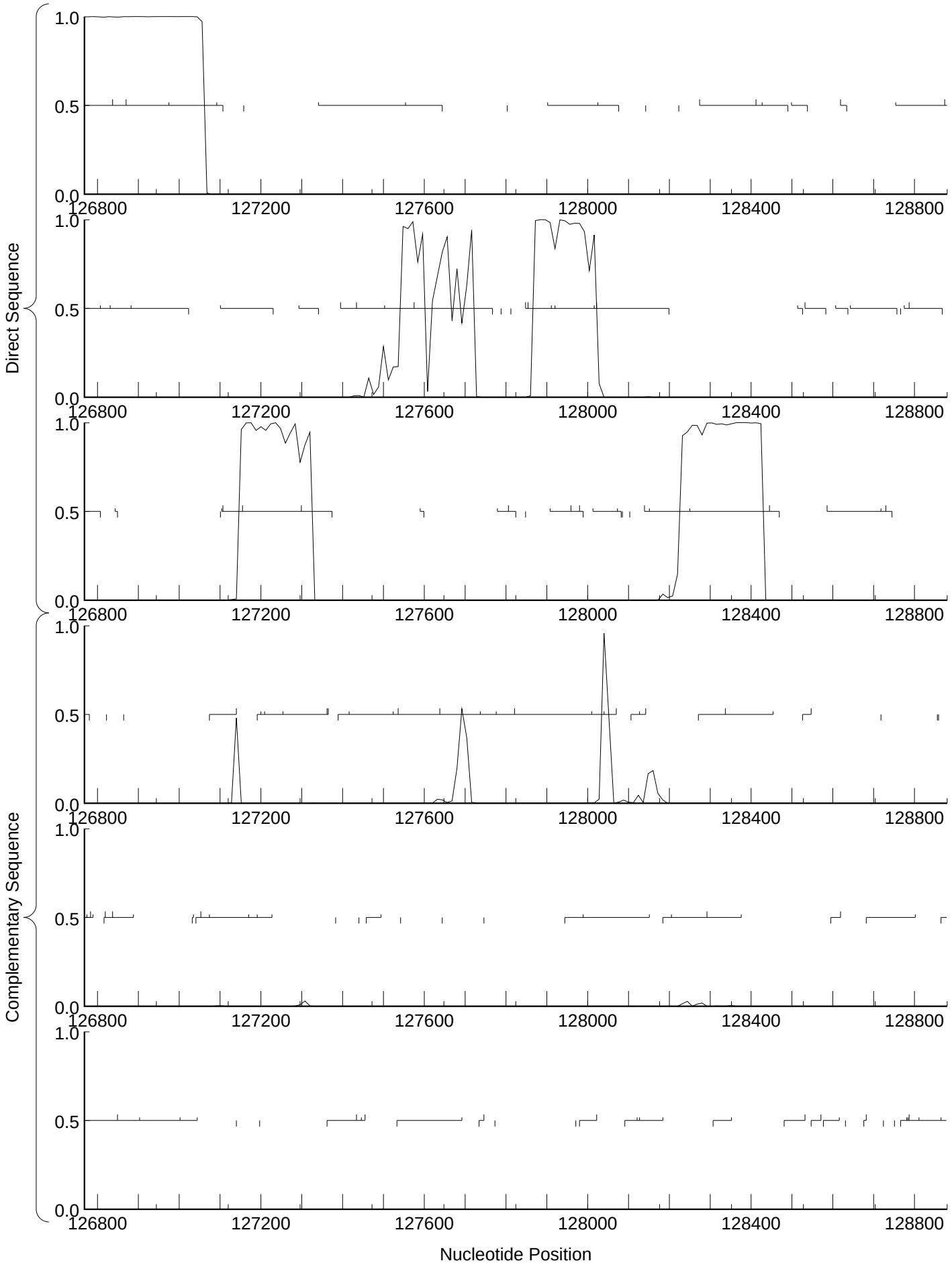
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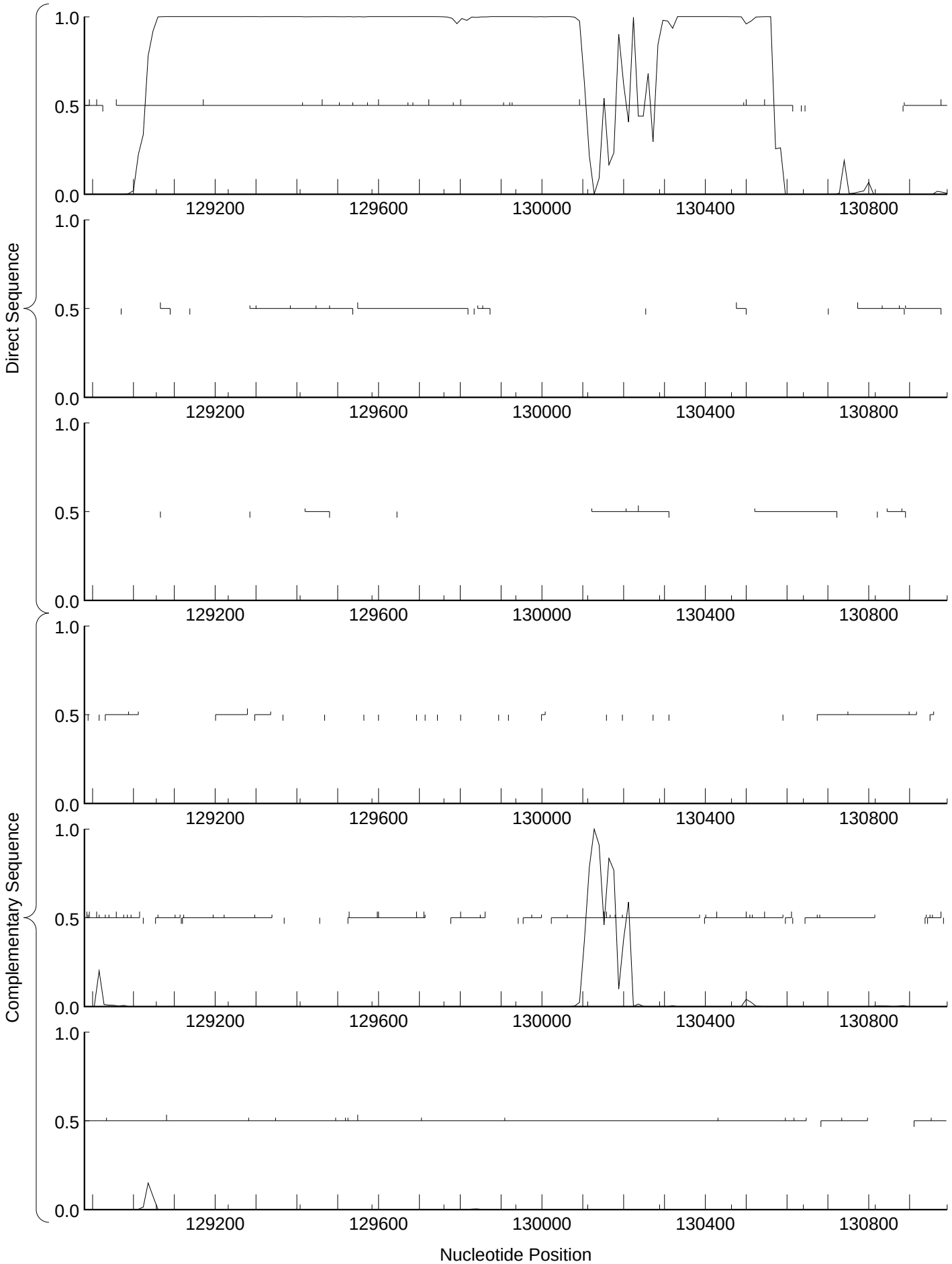
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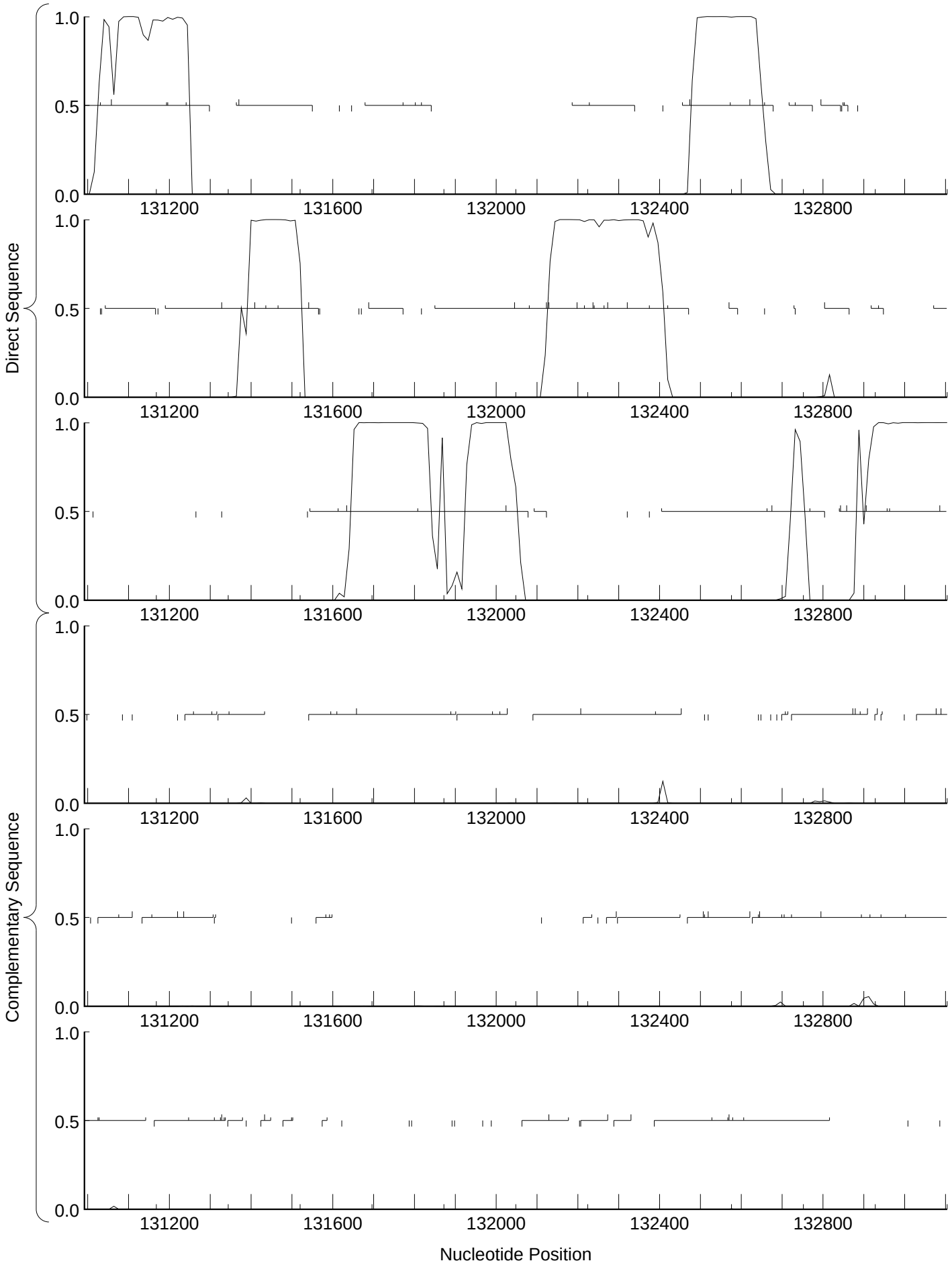
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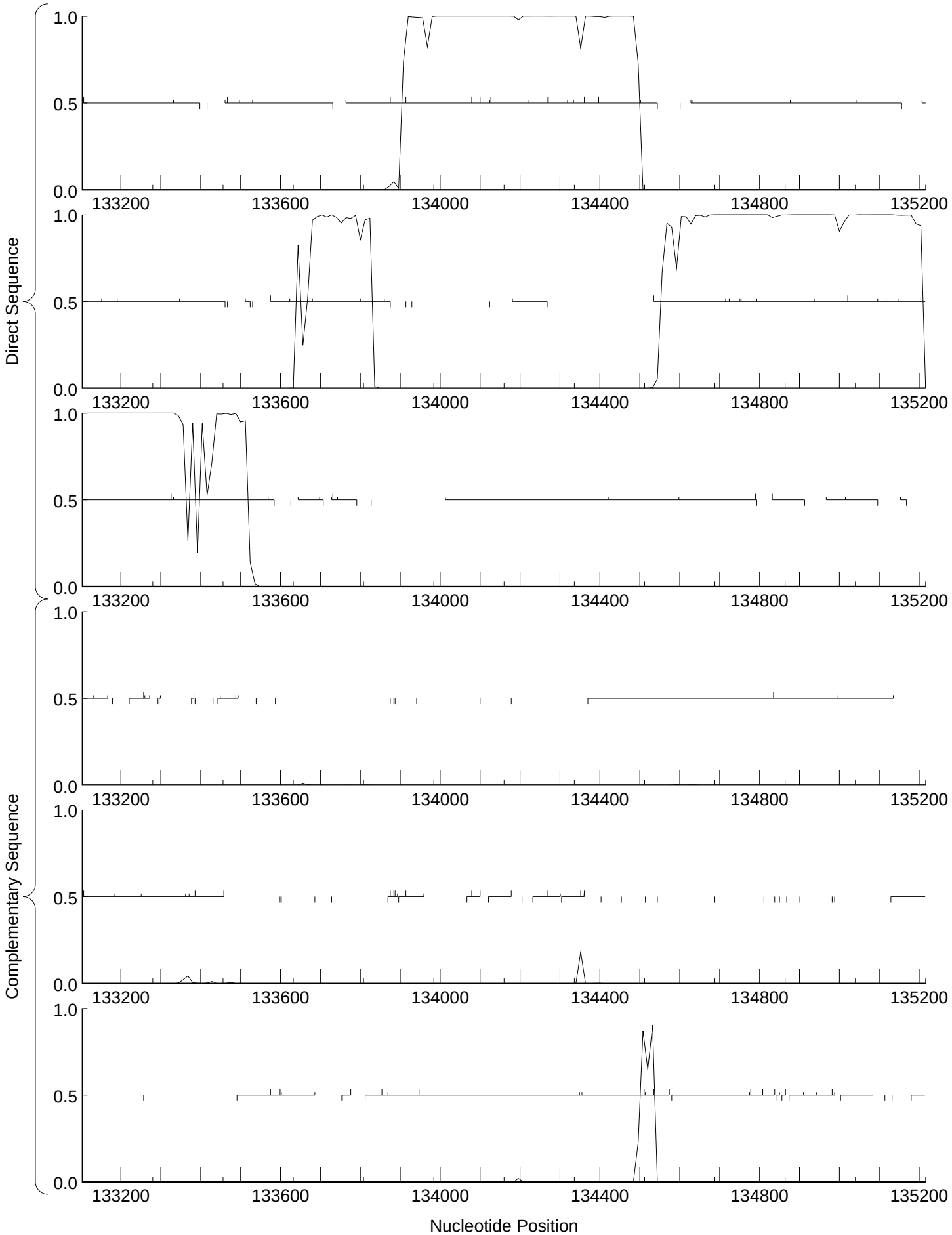
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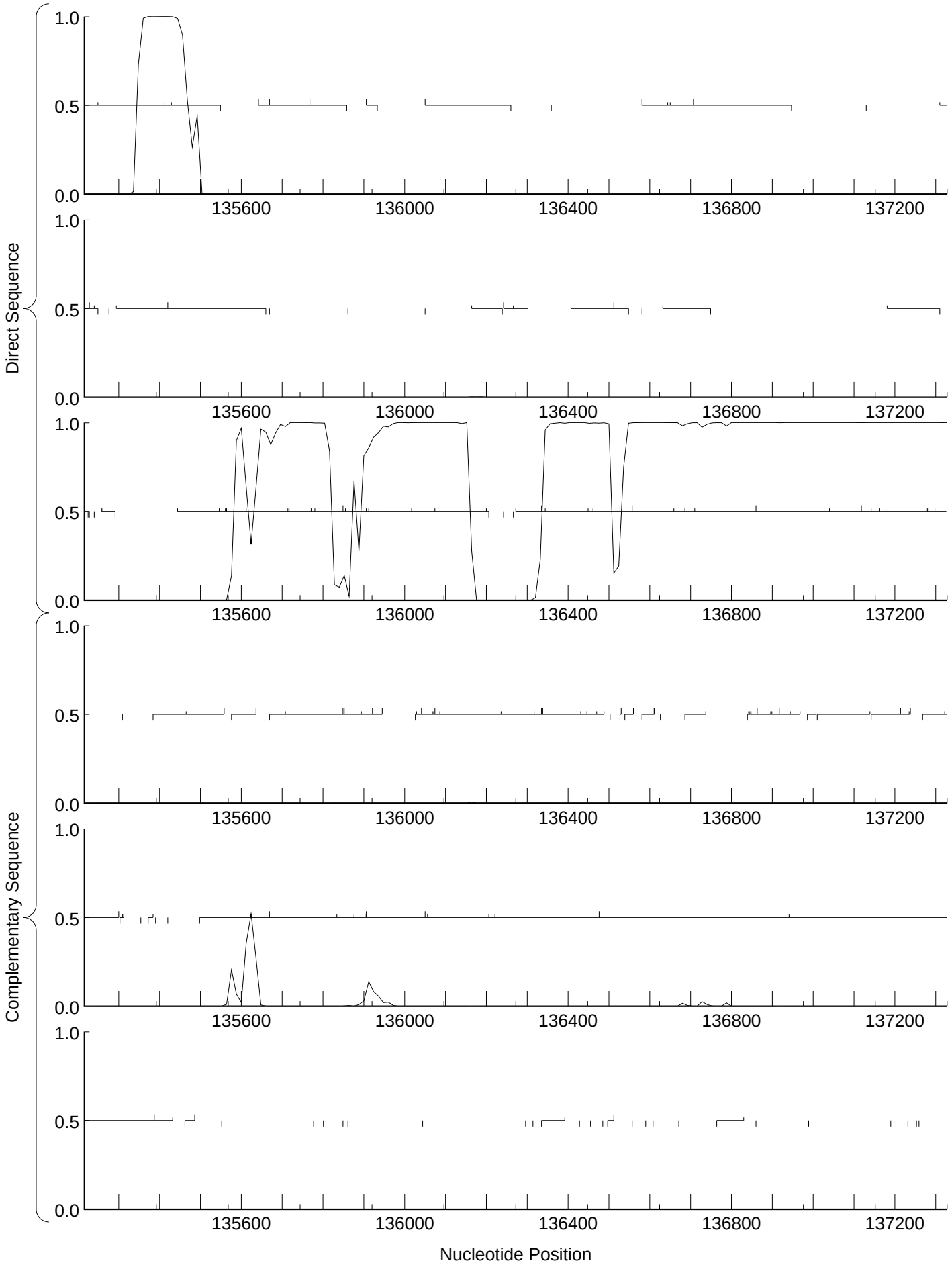
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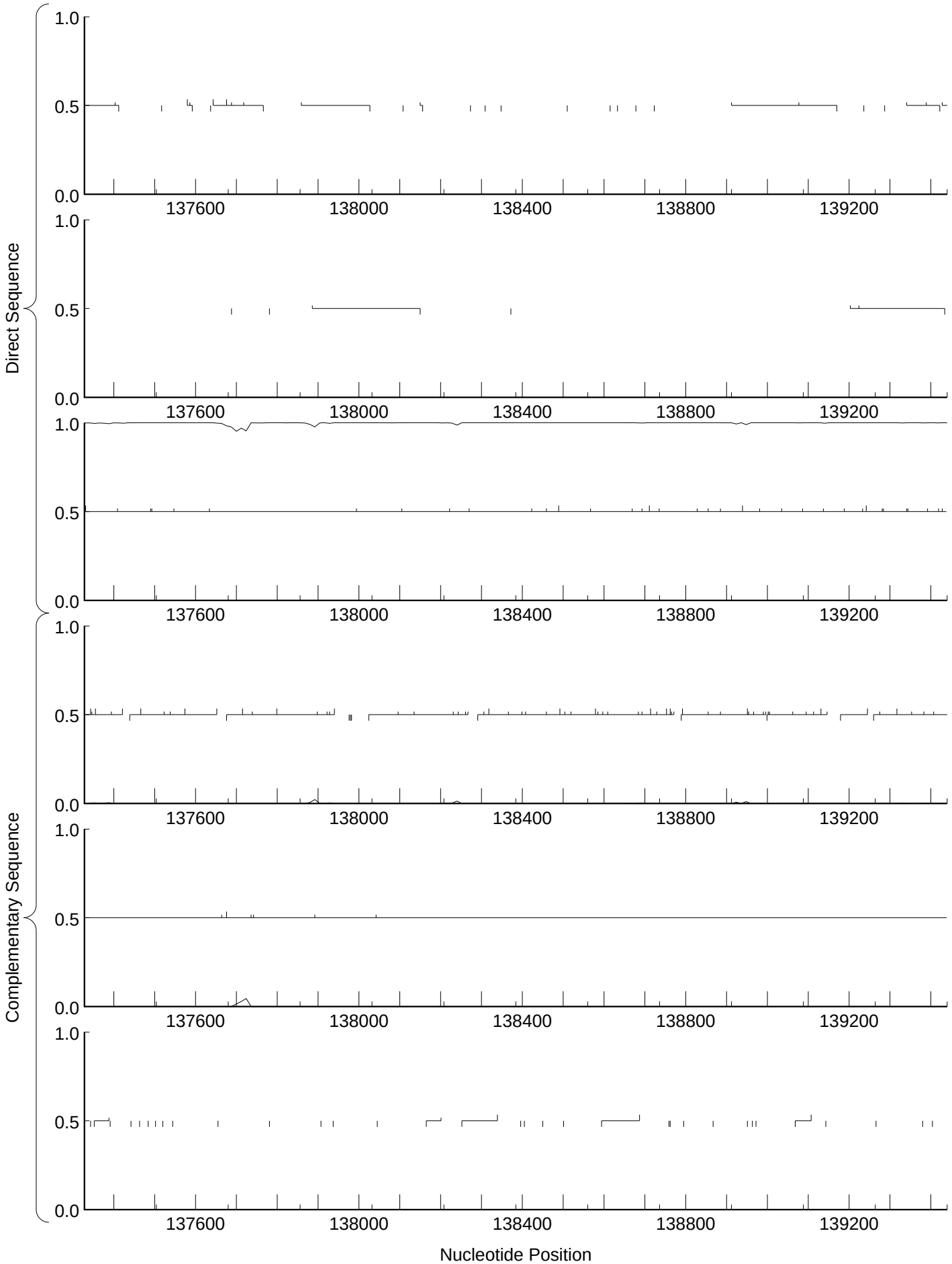
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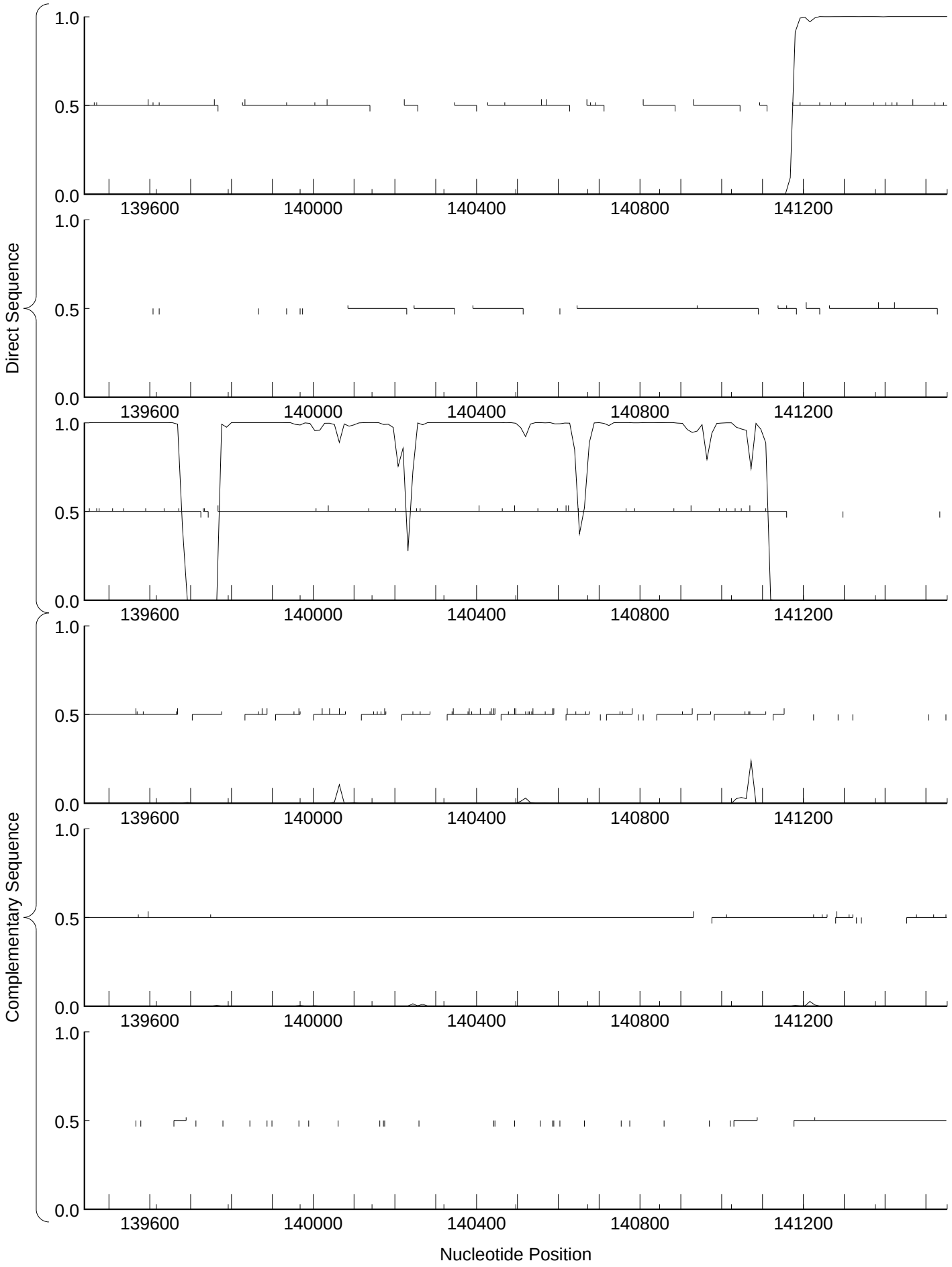
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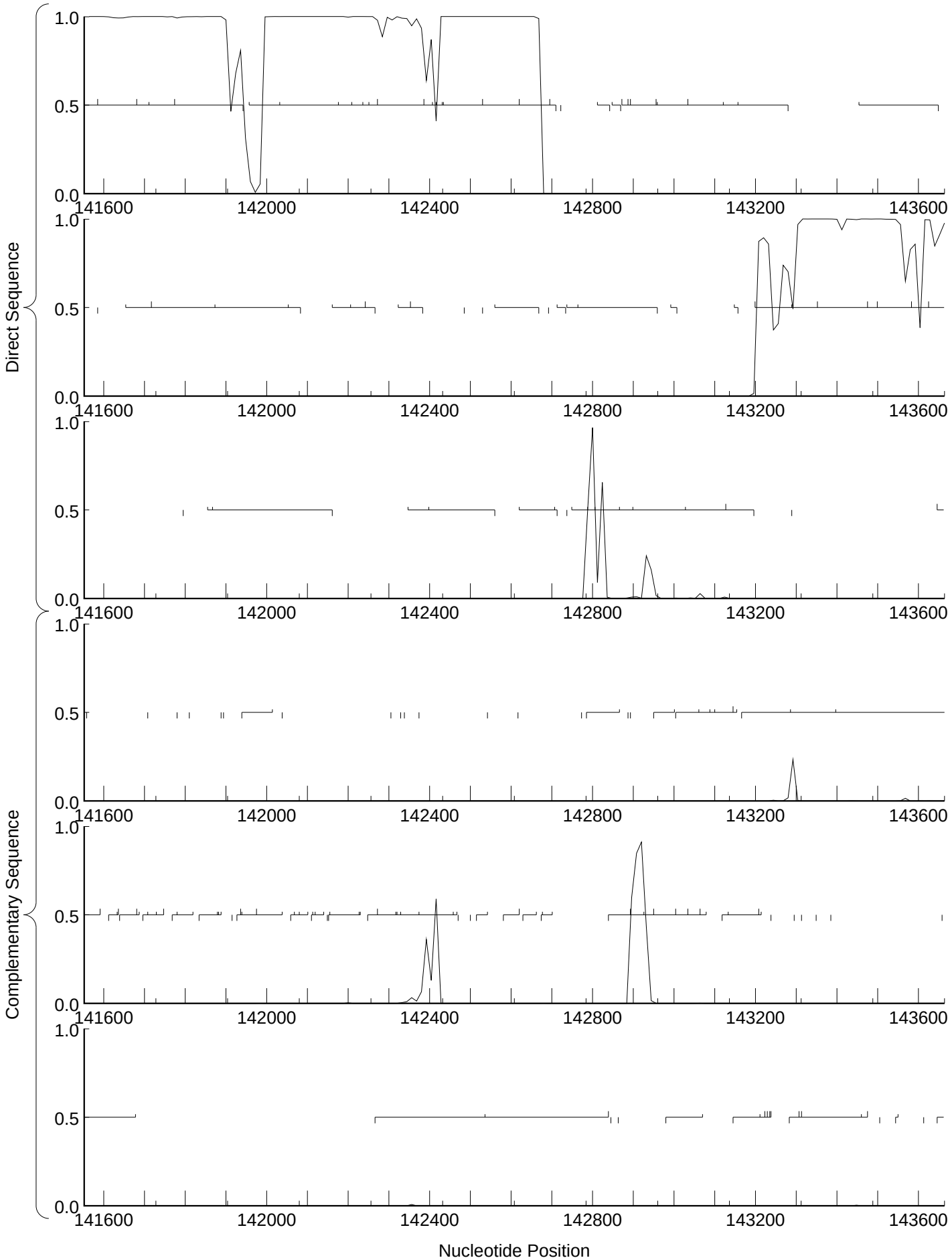
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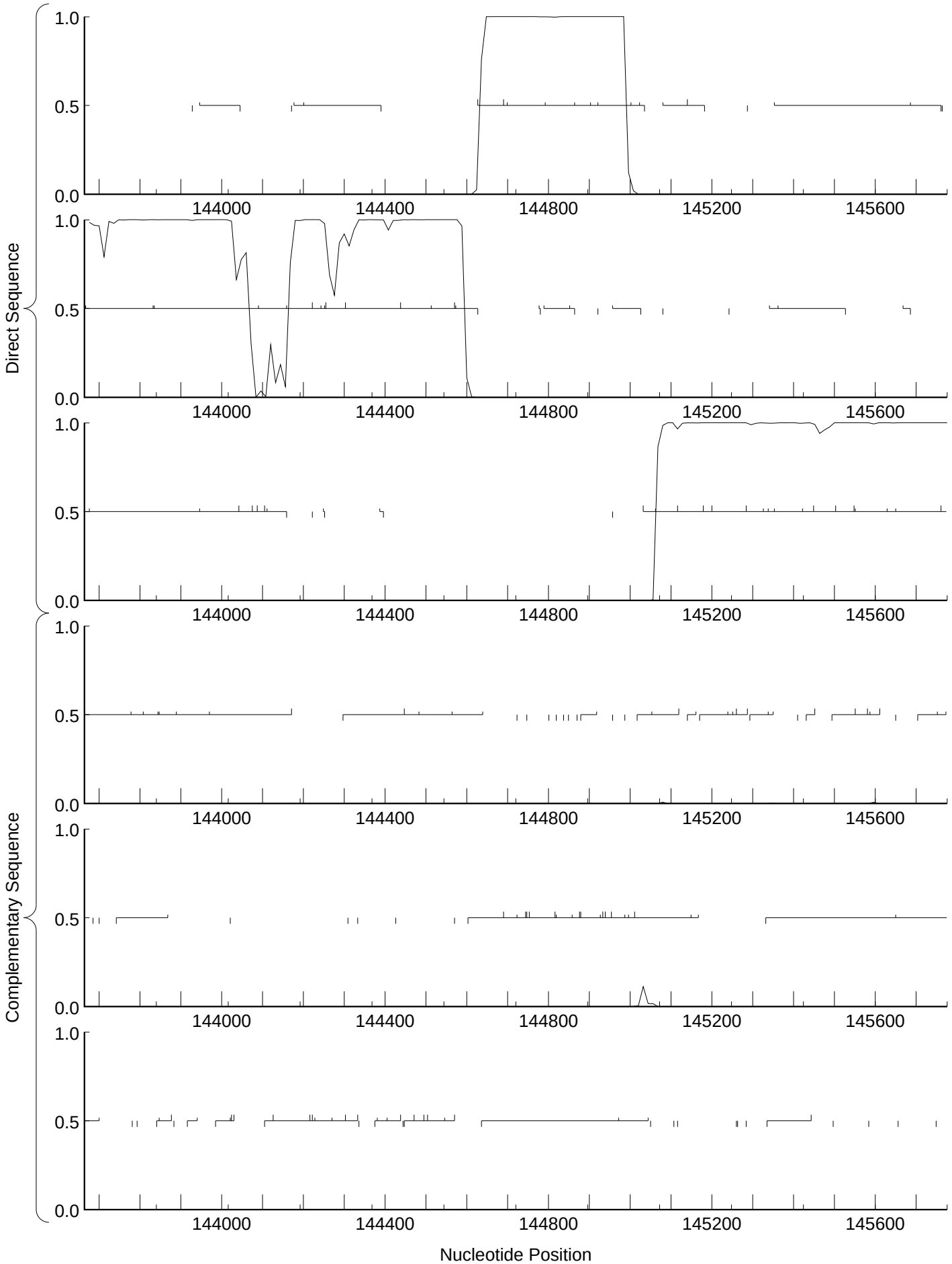
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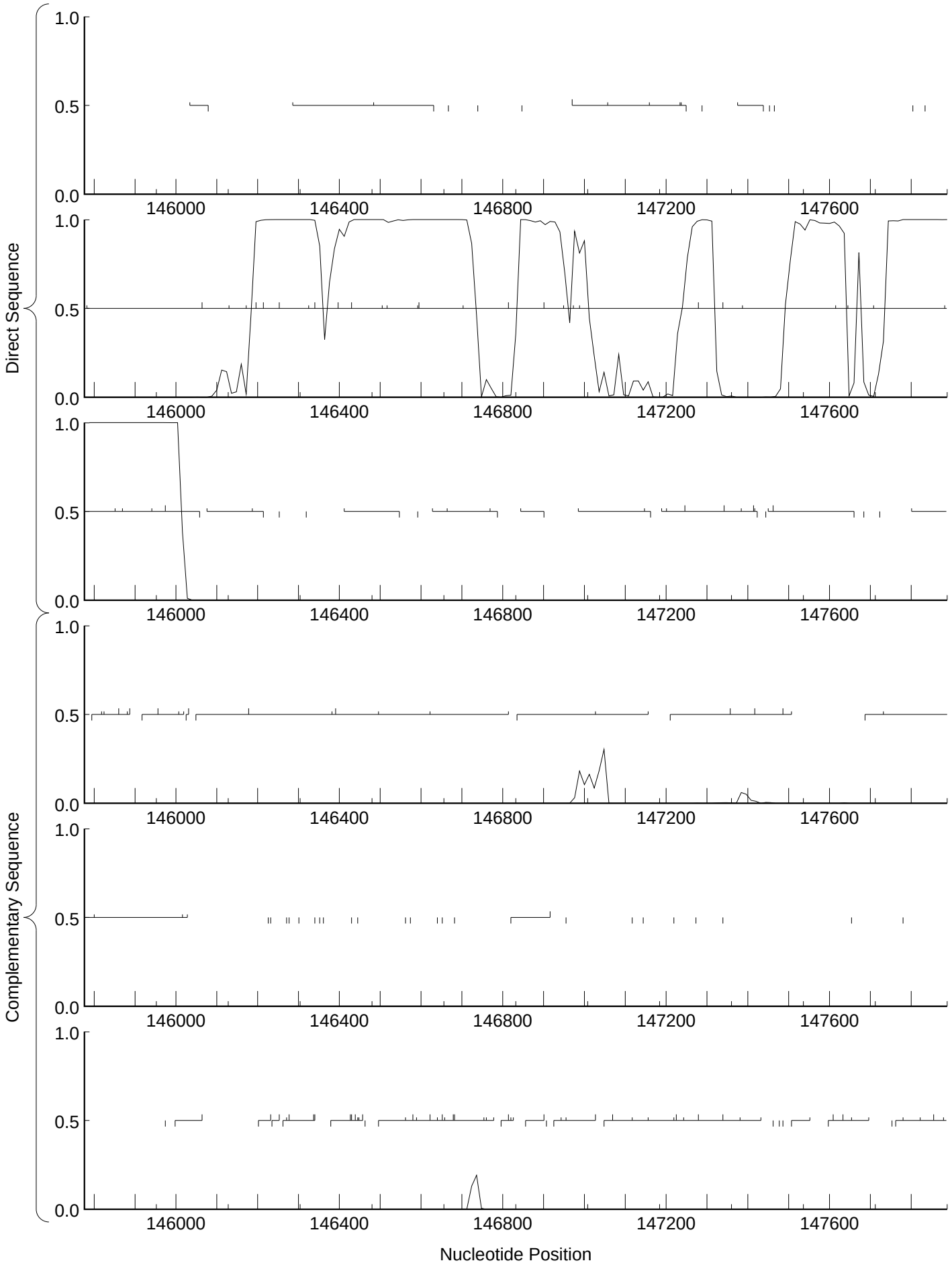
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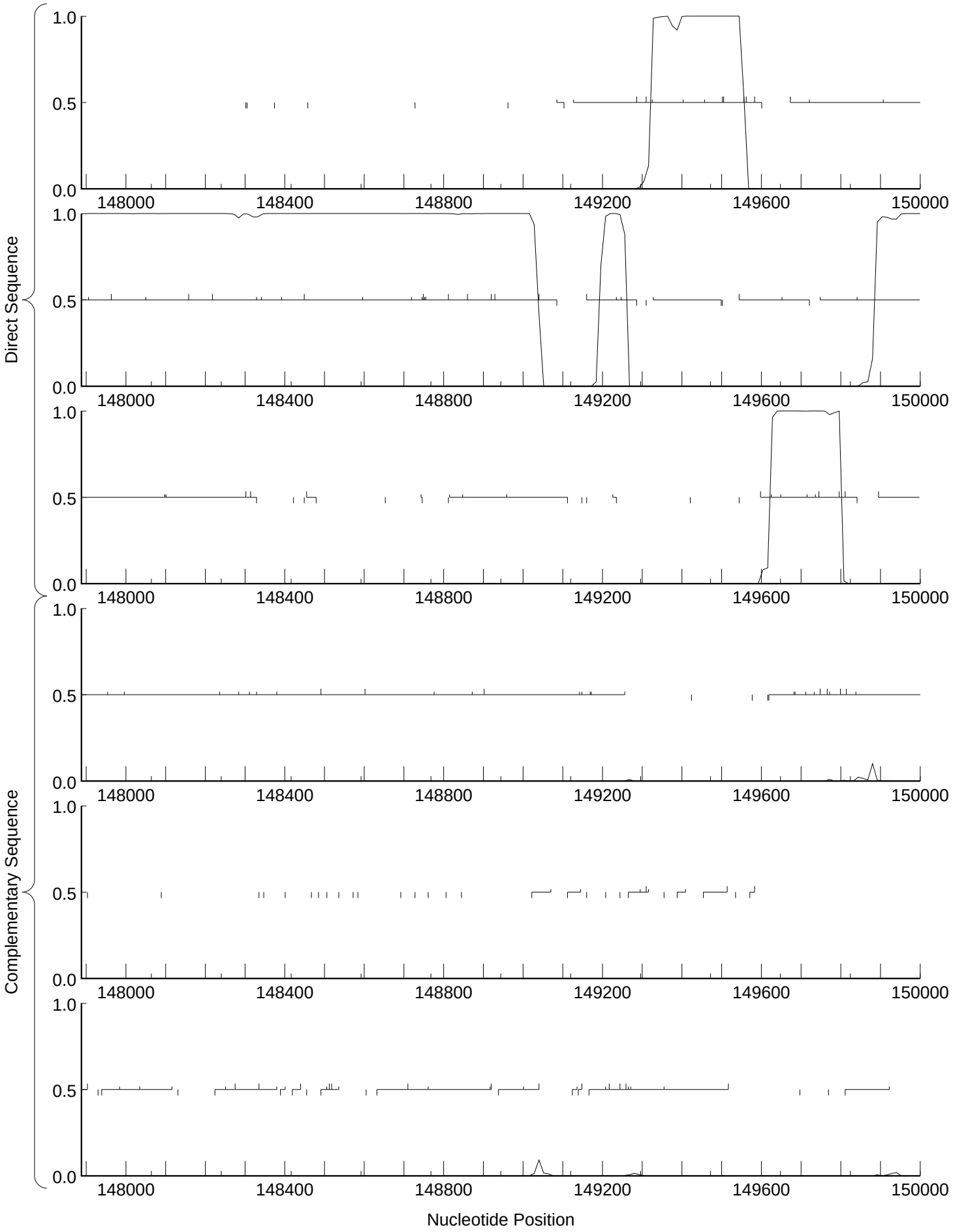
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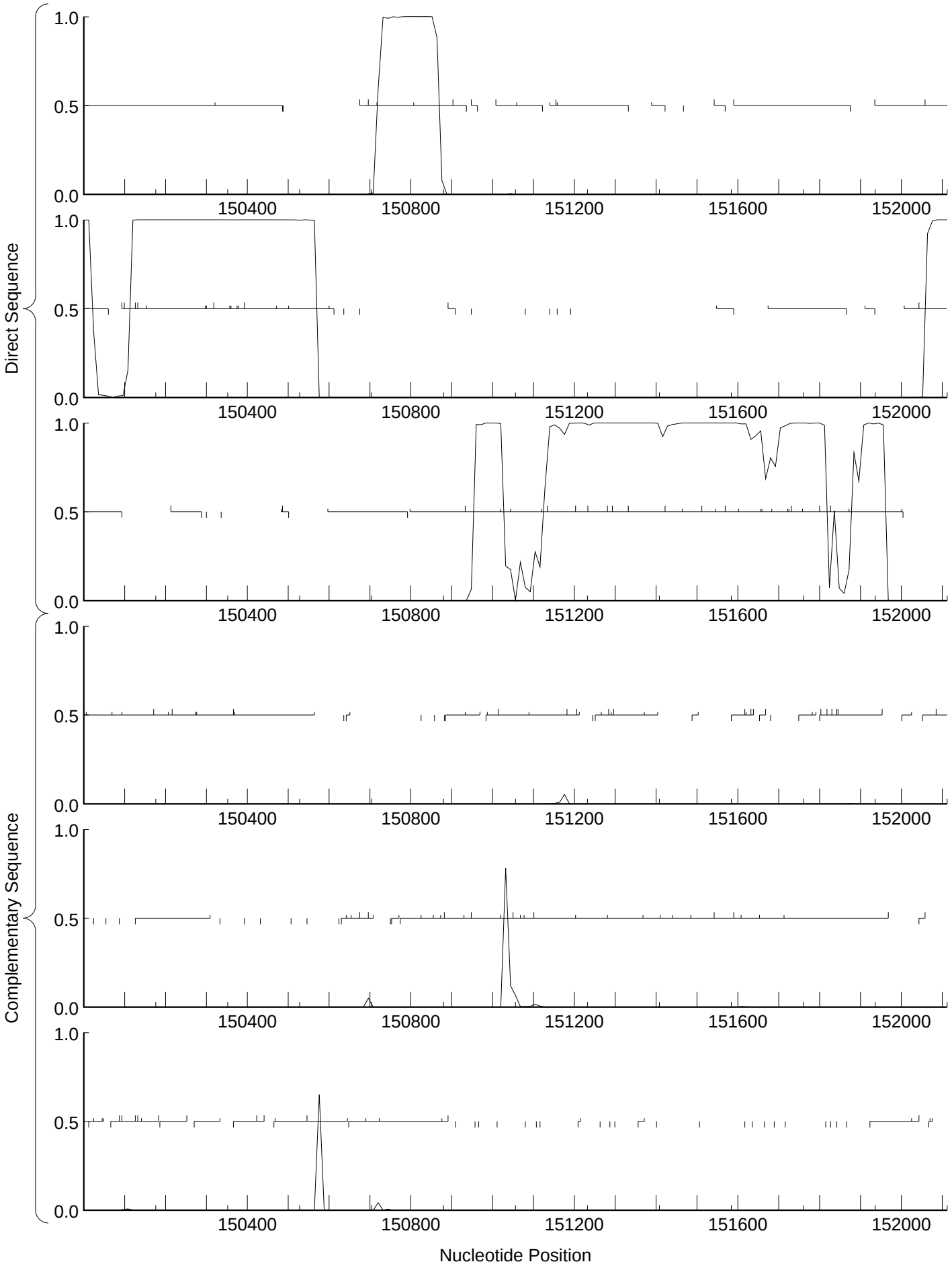
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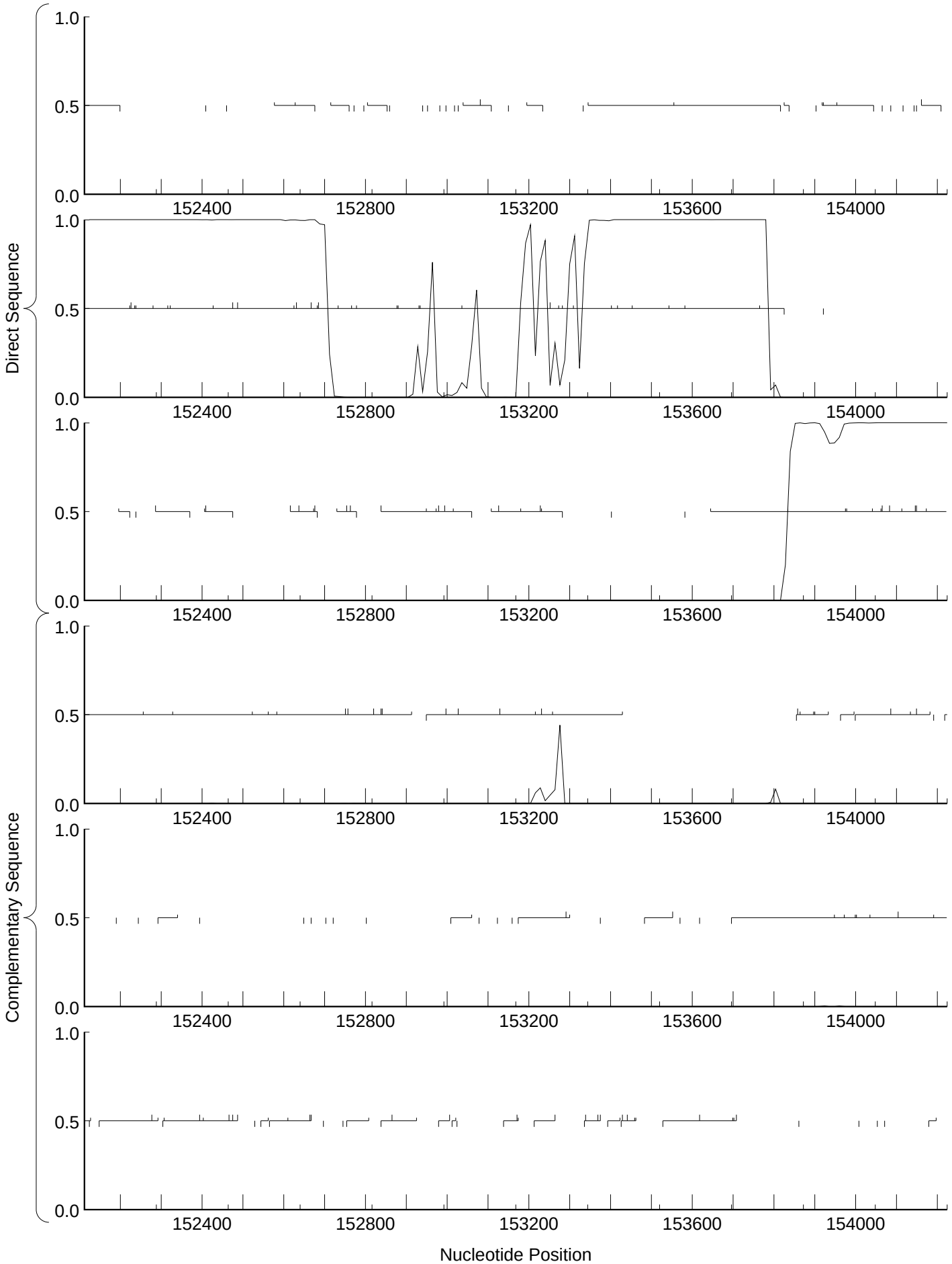
terium phage StubbySaigey complete sequence, 156308 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 72/76



terium phage StubbySaigey complete sequence, 156308 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 73/76



terium phage StubbySaigey complete sequence, 156308 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 74/76



terium phage StubbySaigey complete sequence, 156308 bp, circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 75/76

