GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Arthrobacter phage BarretLemon complete sequence, 51290 bp, circularly permuted, Cluster AO
Analysis Date    :  3/28/24 at 20:50:56
Pages            :  26
Sequence Length  :  51290 bp
GC Content       :   60.90%

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

PS-Version       :  1.2

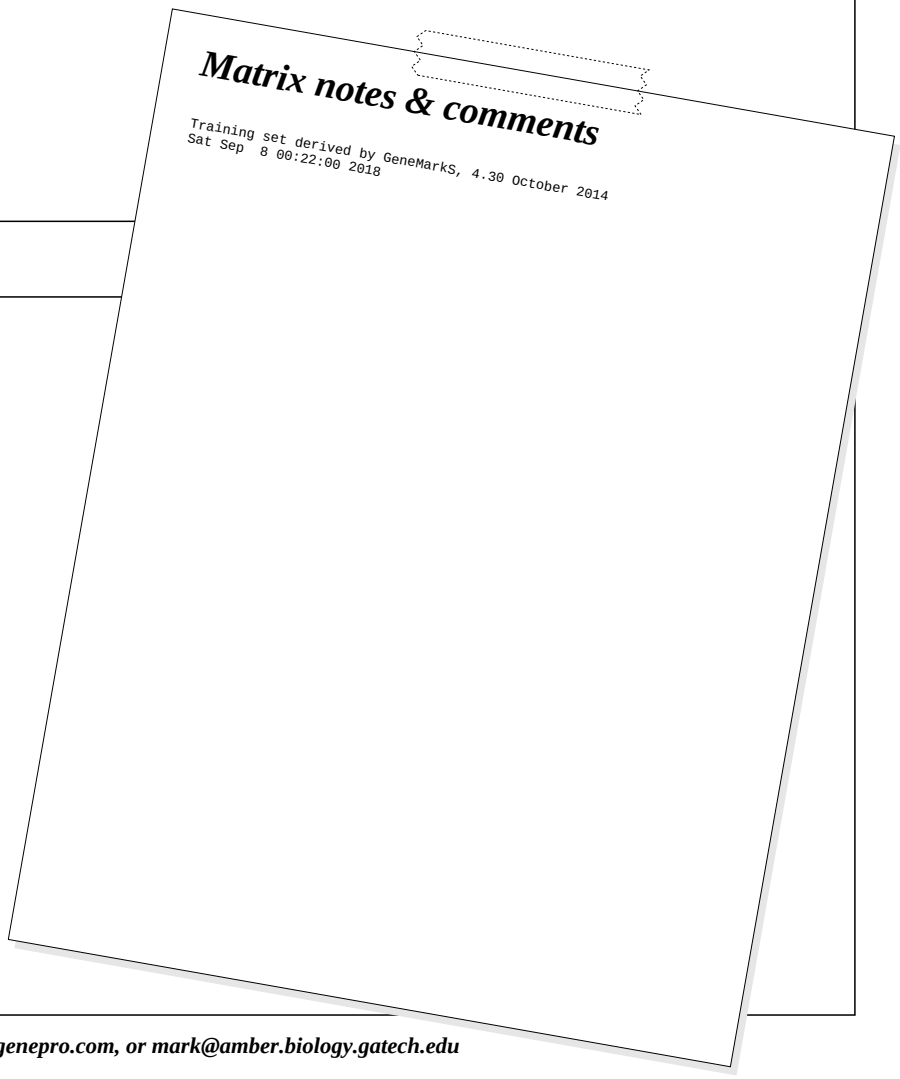
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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                    List regions and/or splice sites,
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MATRIX INFORMATION

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Author     :  -

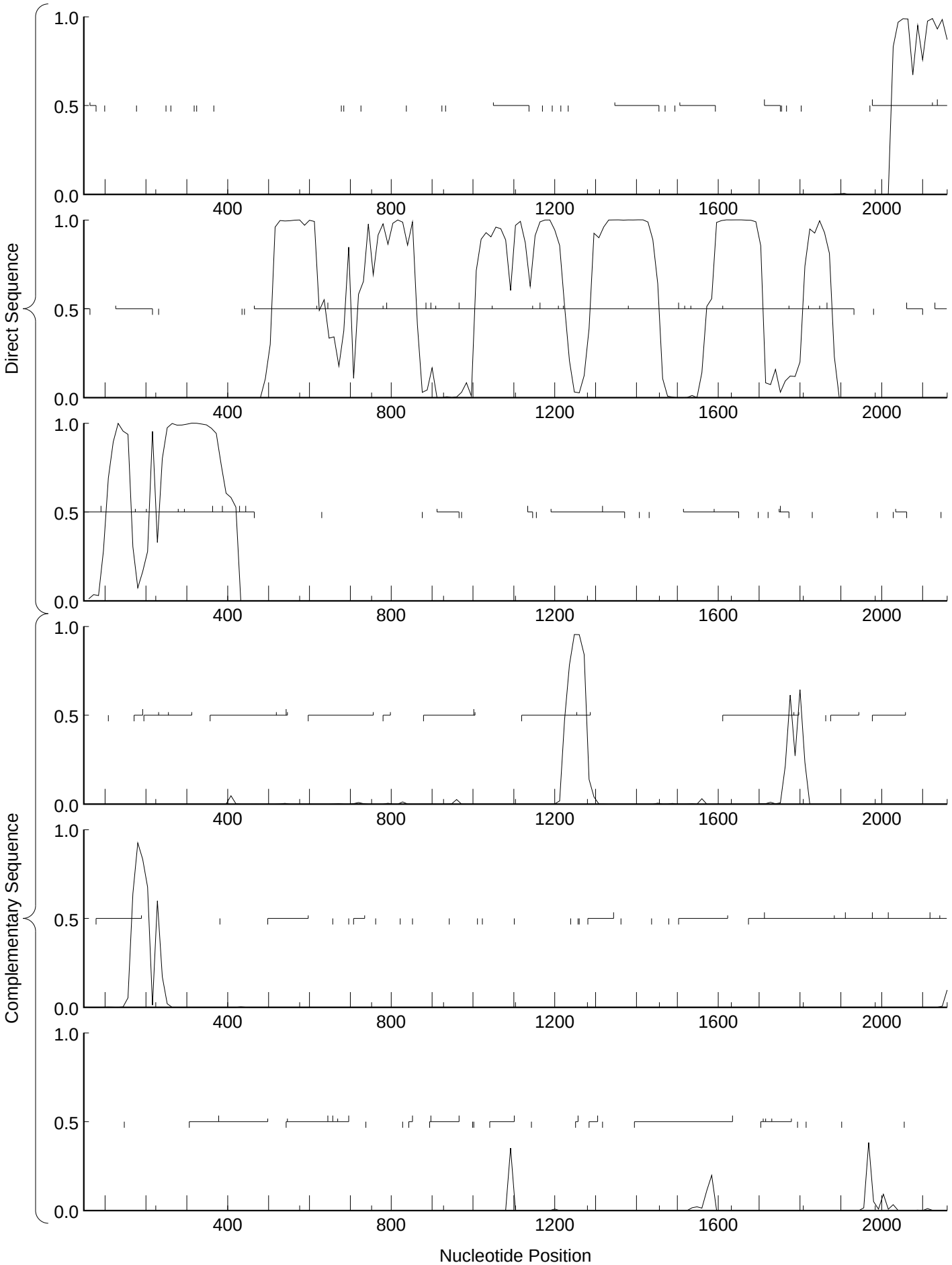
Order      :  4


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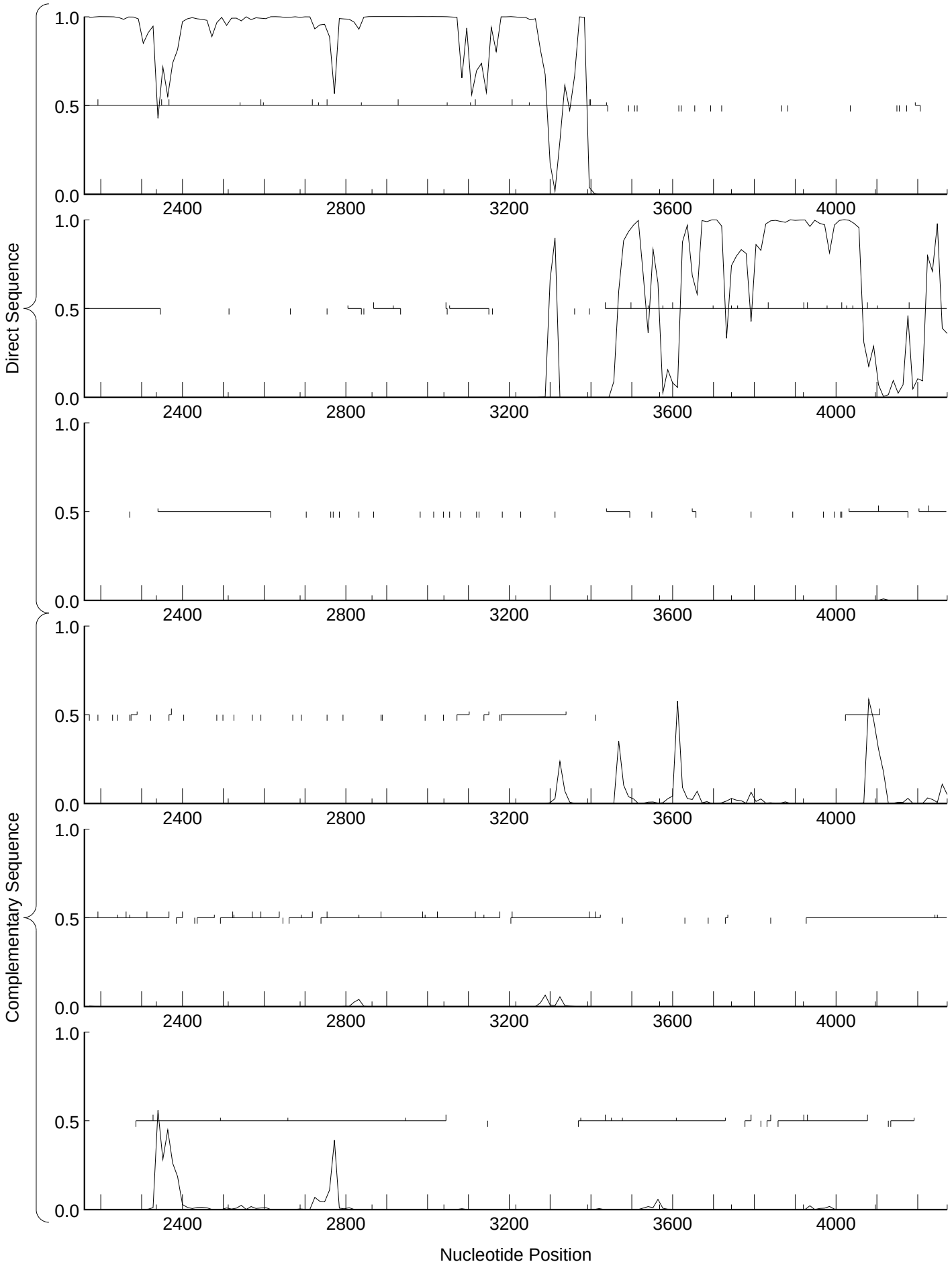


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

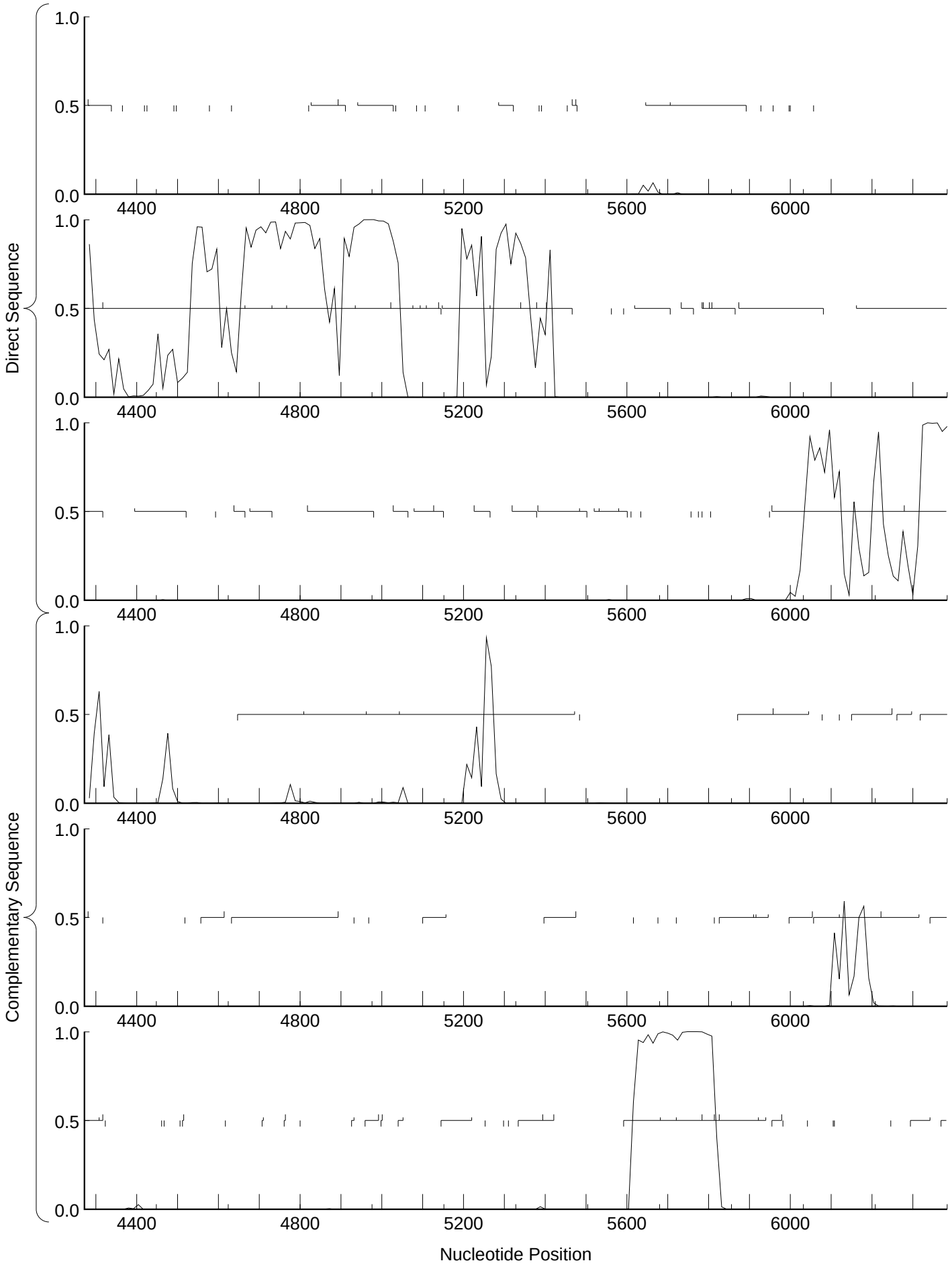
obacter phage BarretLemon complete sequence, 51290 bp, circularly permuted, Cluster AO2, Order 4, Window 96, Step 12, 2/26



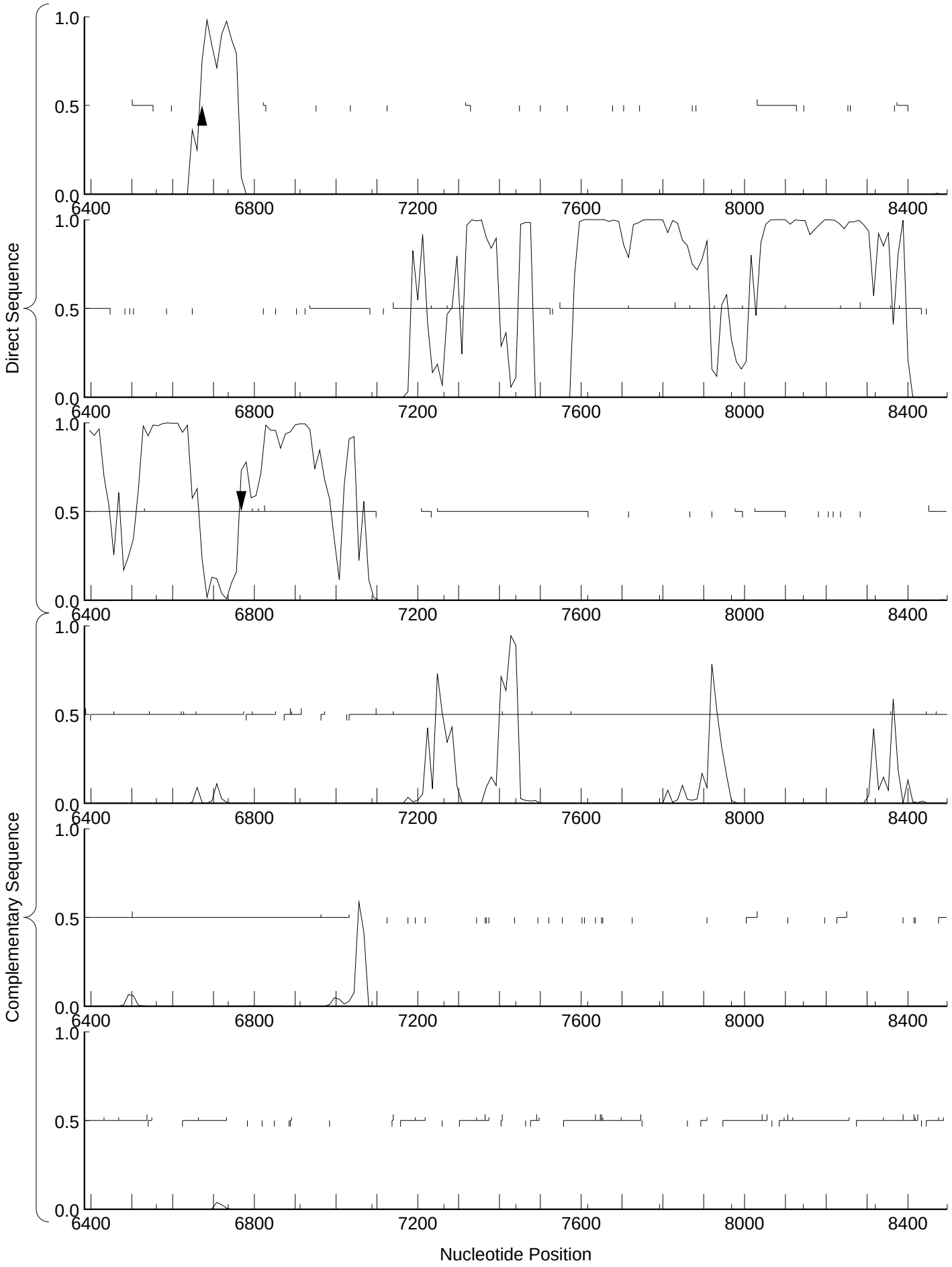
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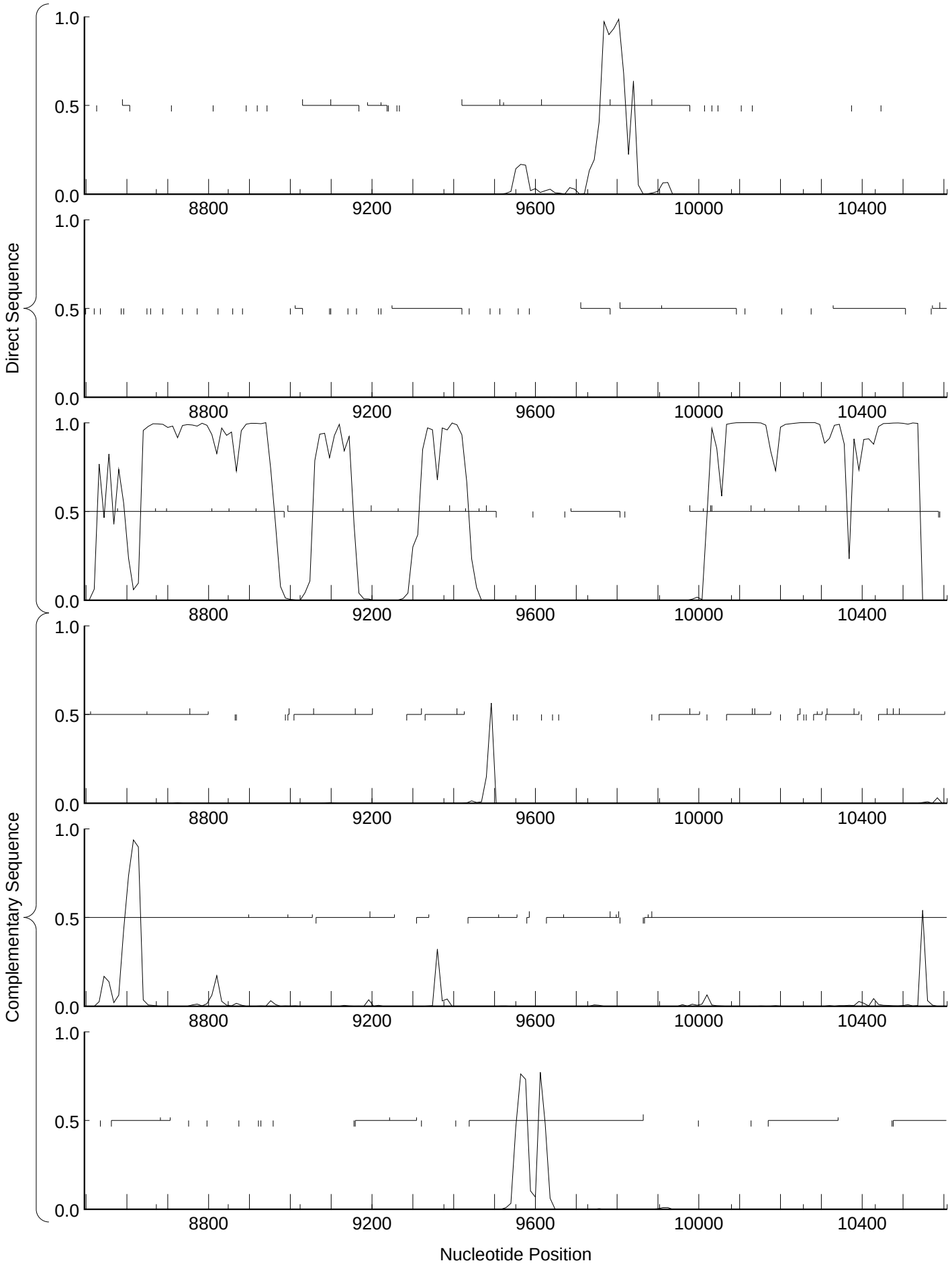
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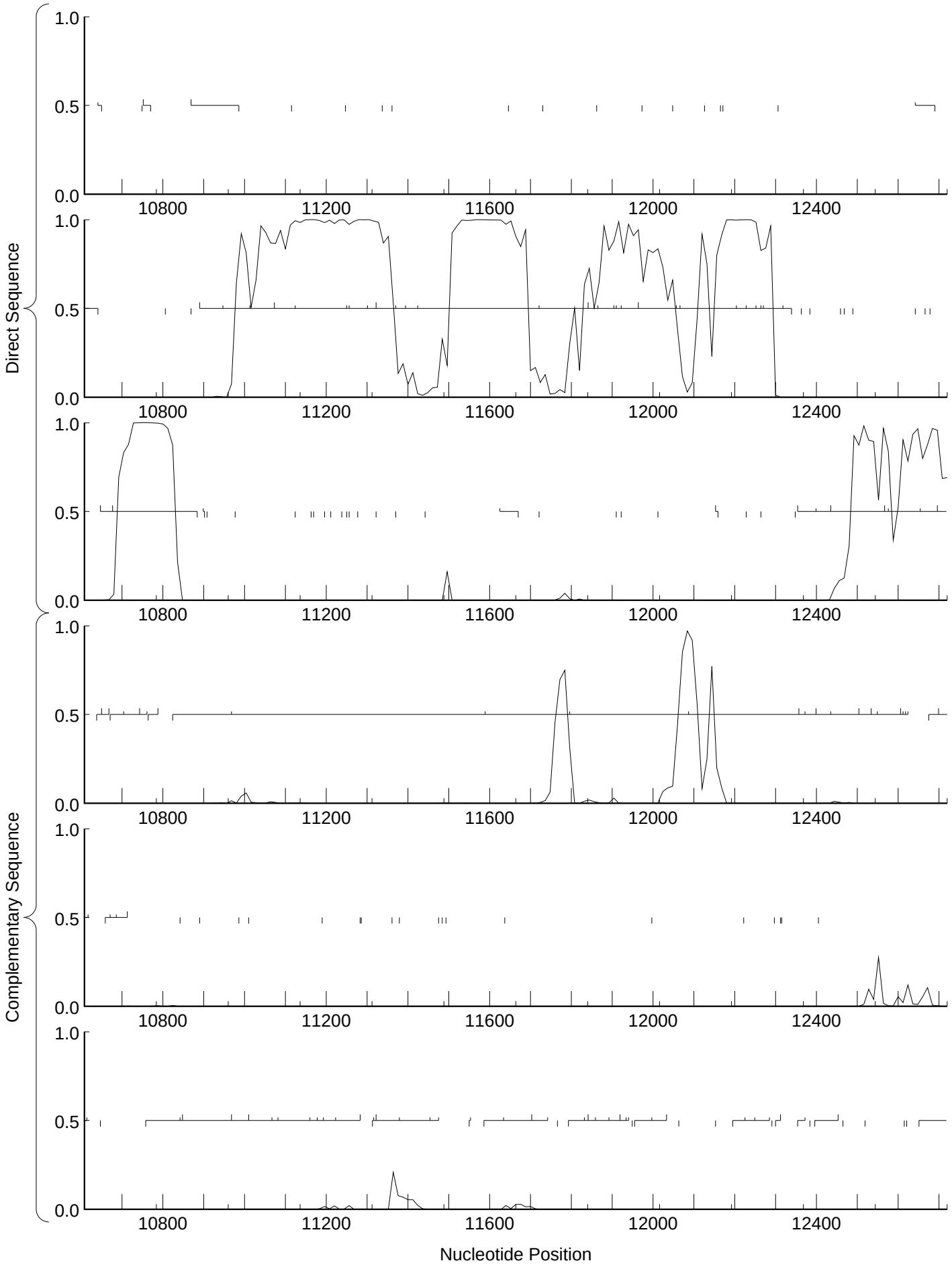
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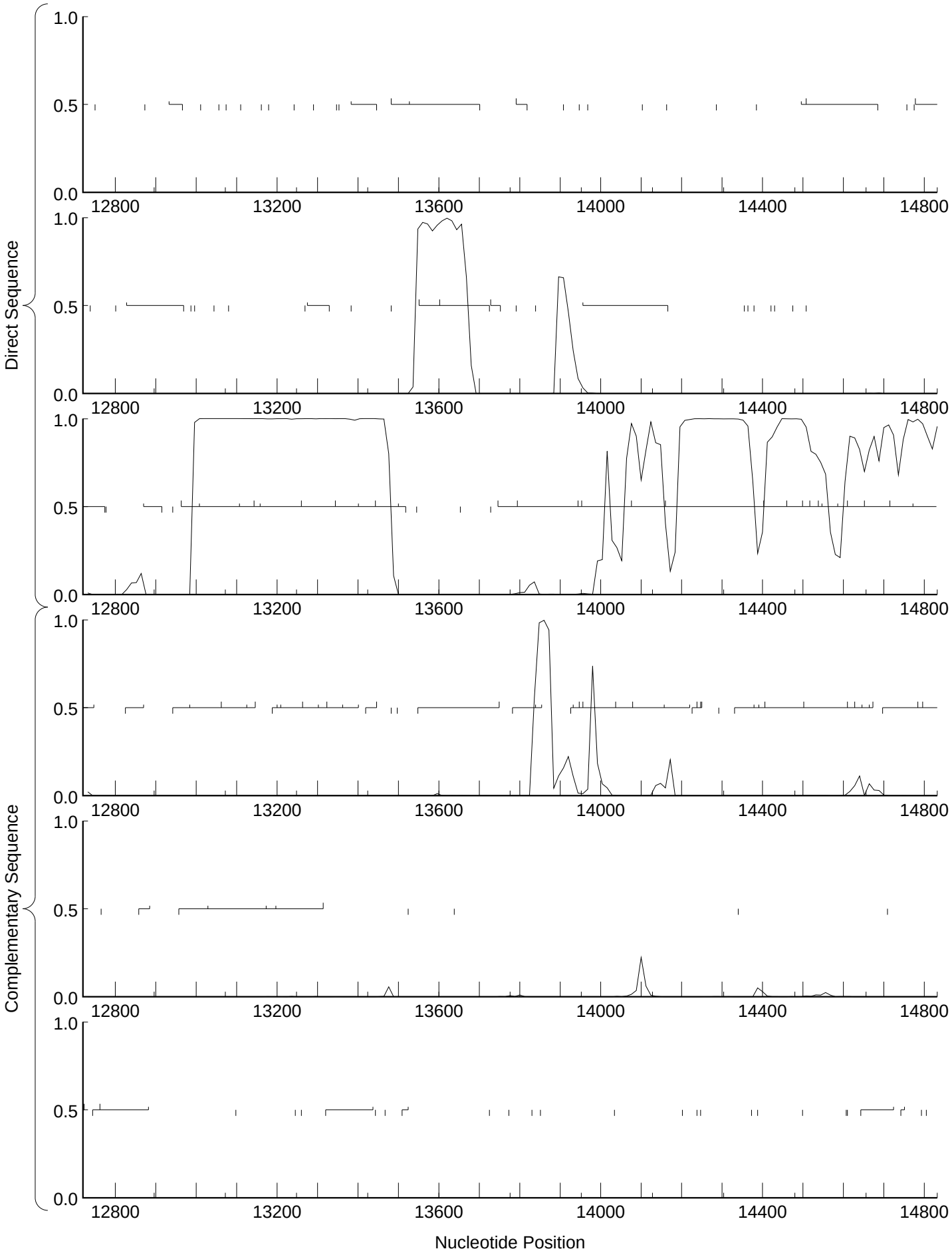
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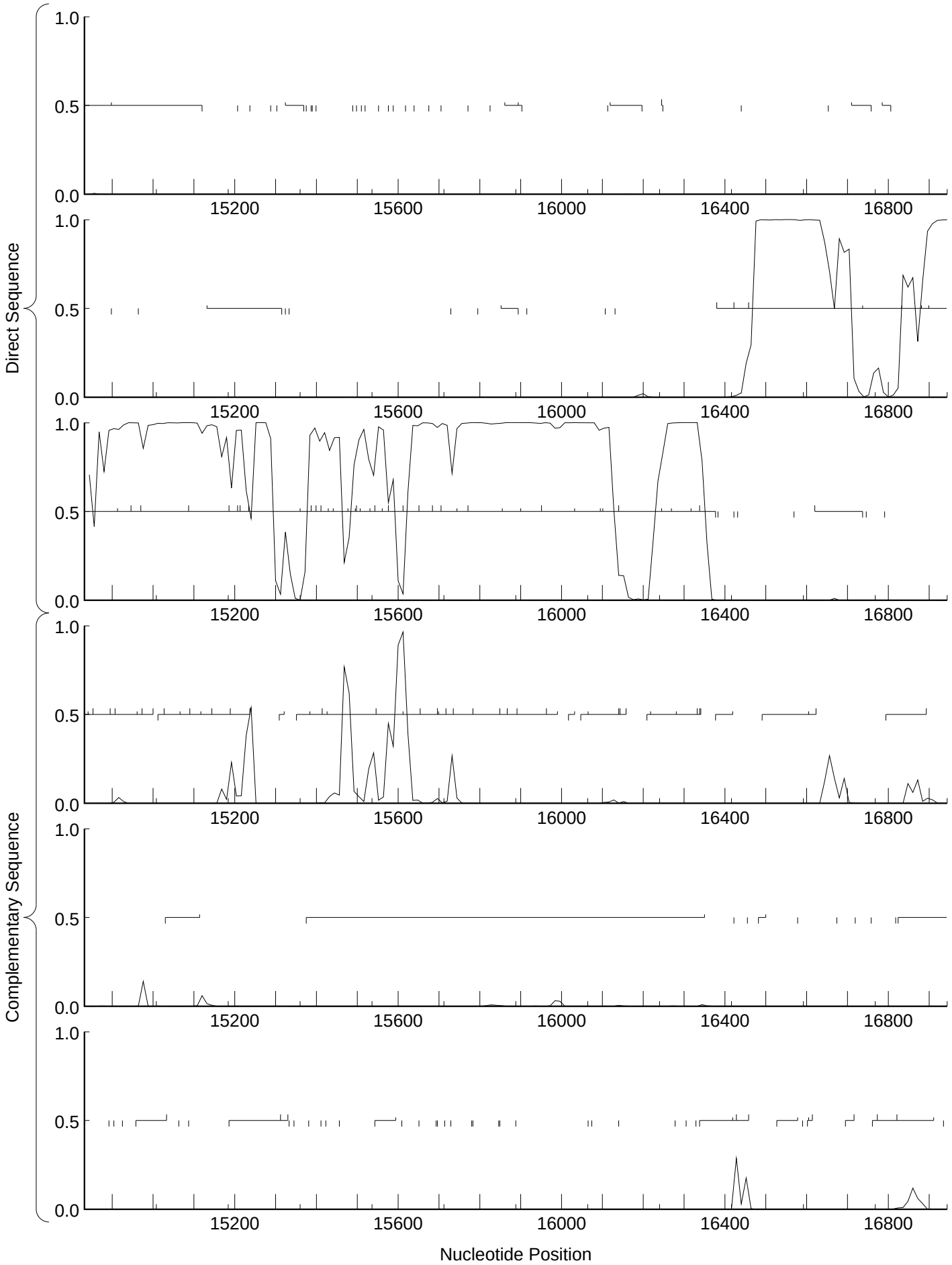
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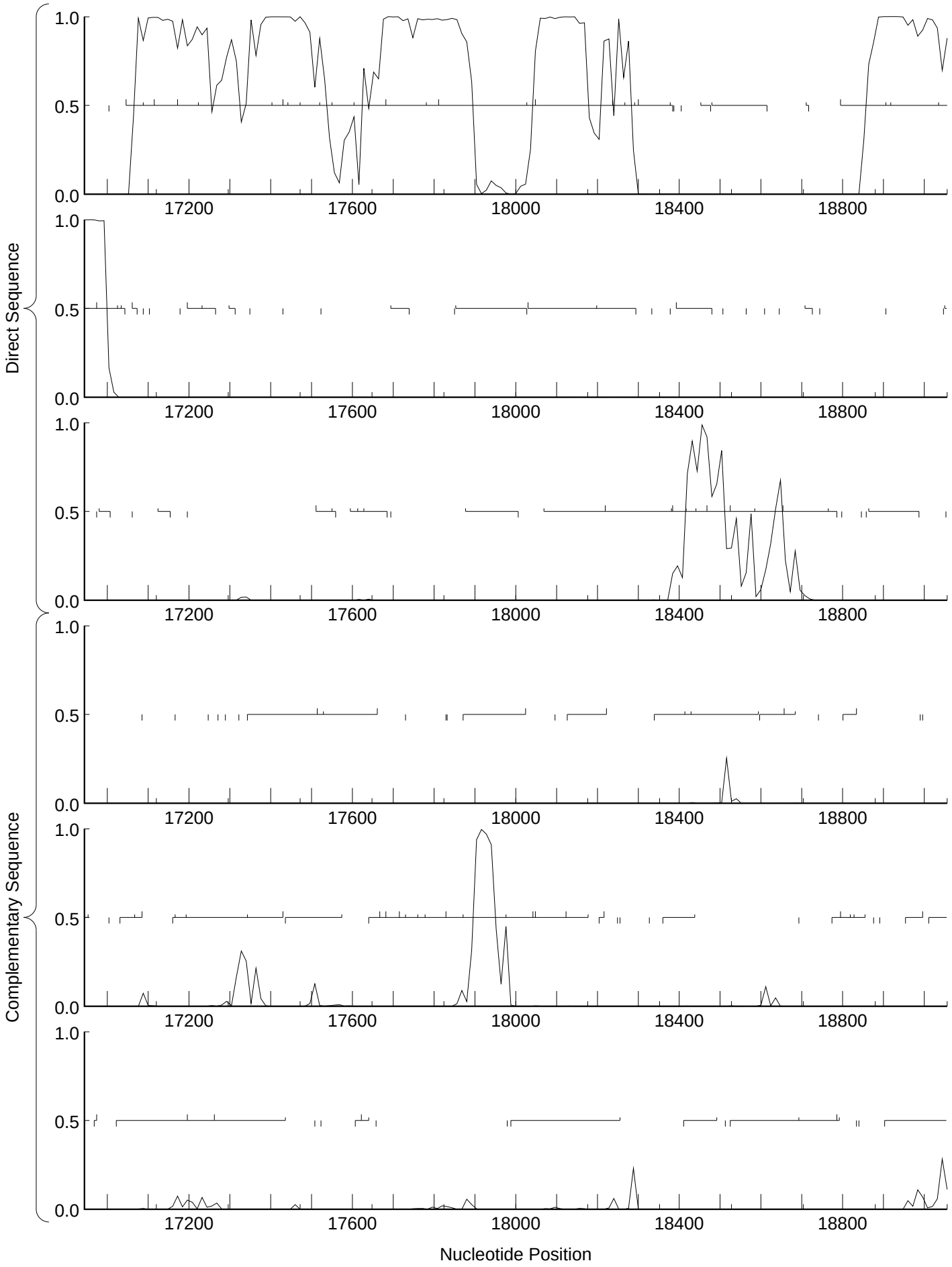
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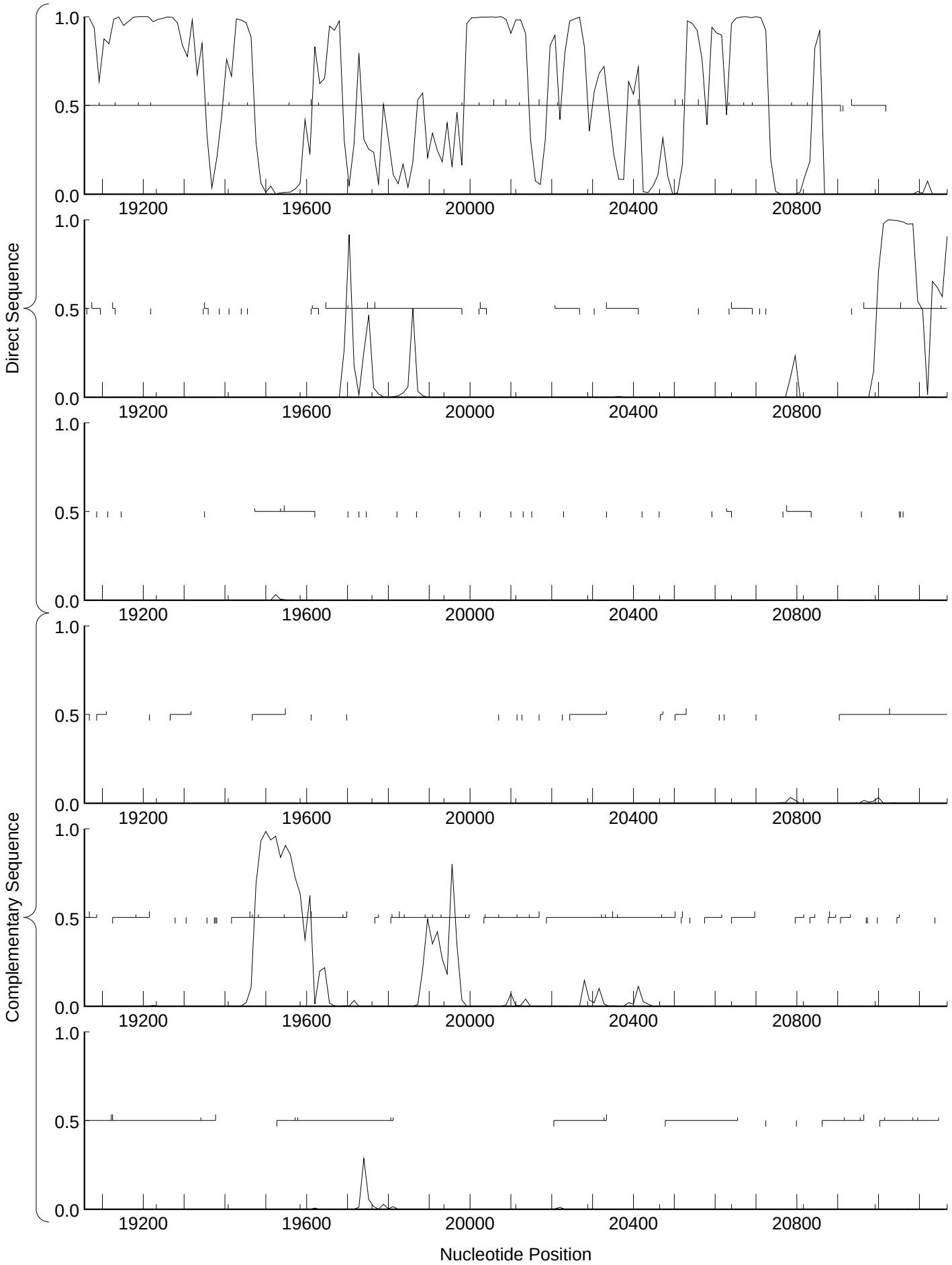
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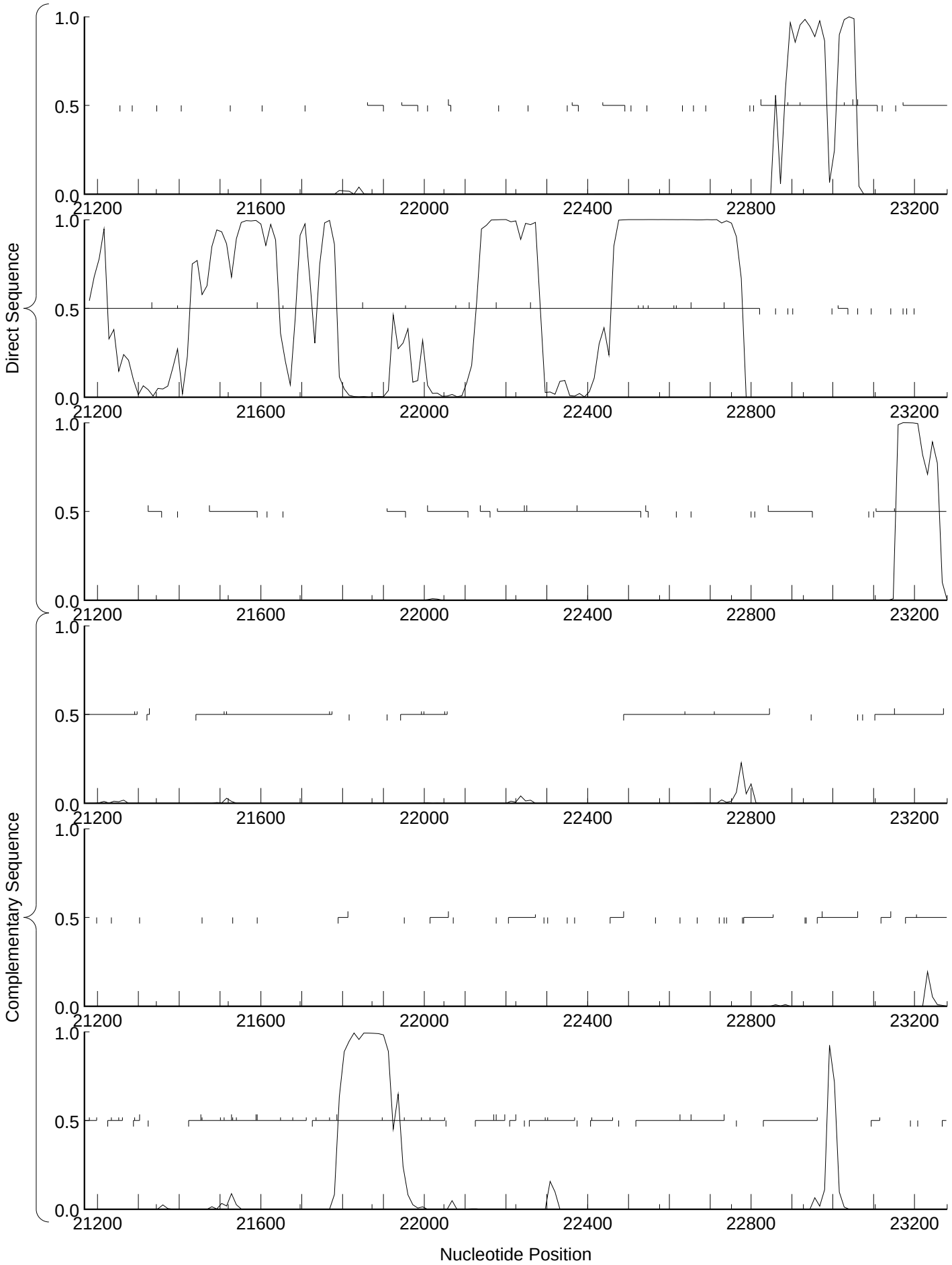
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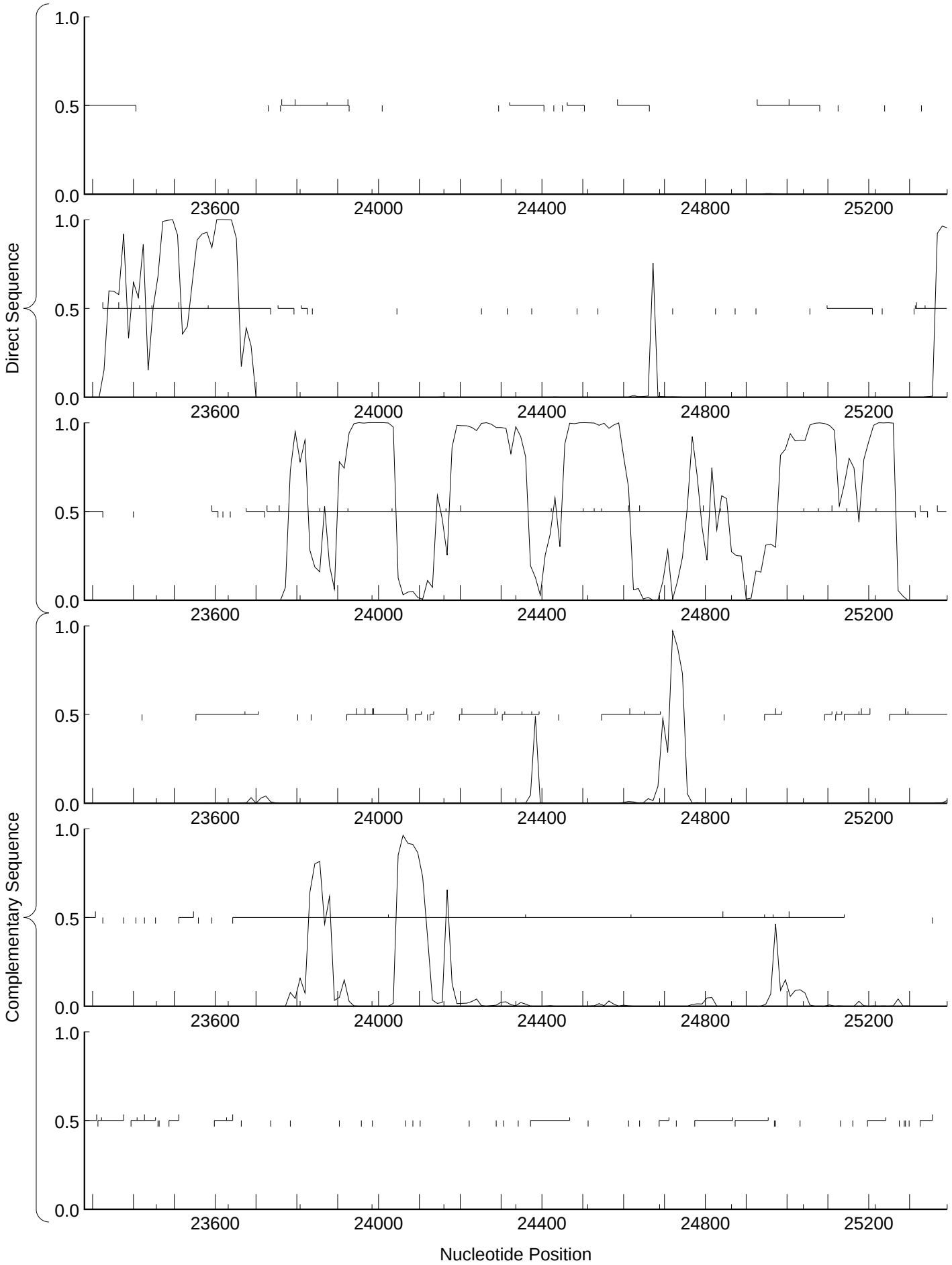
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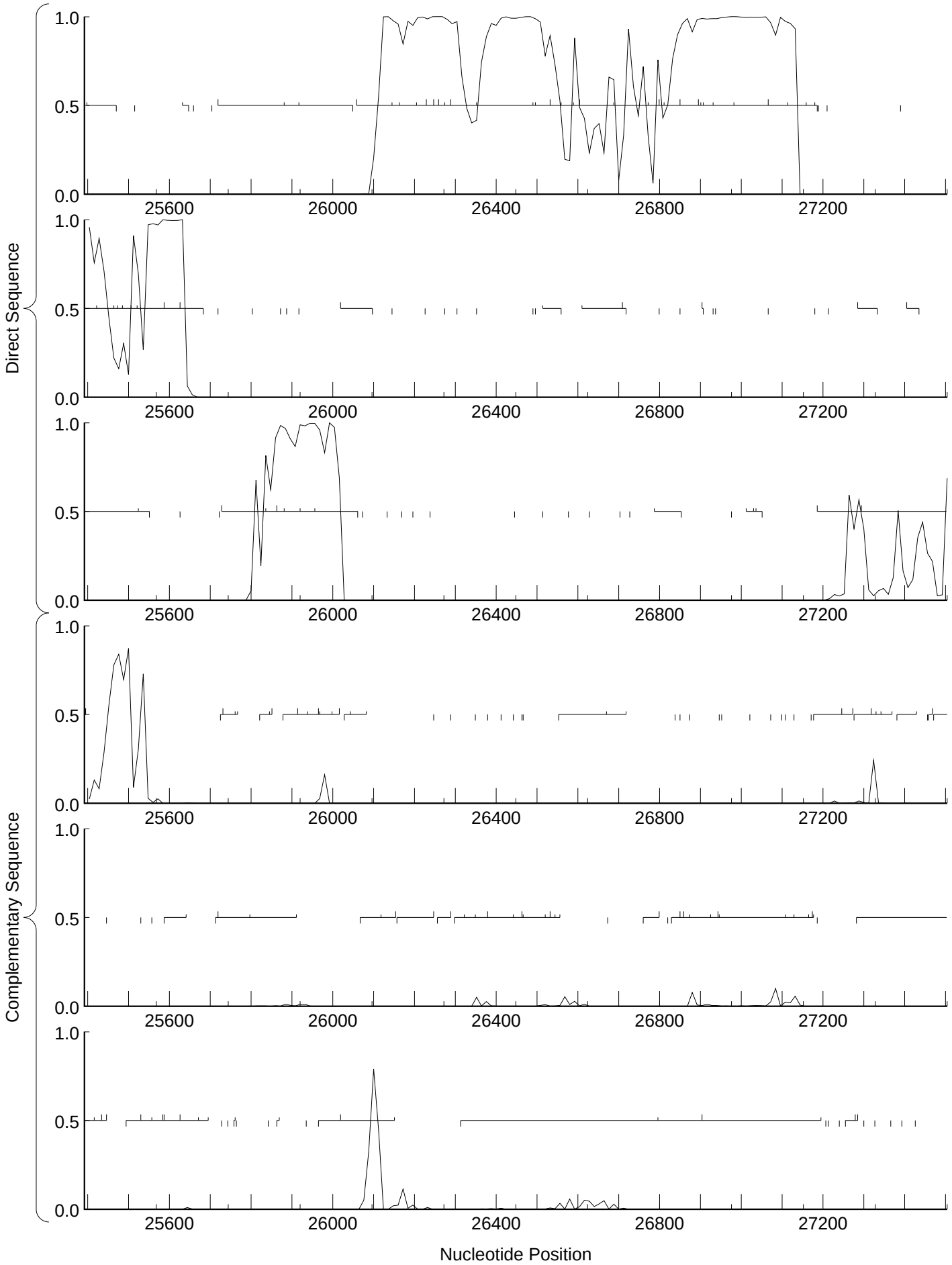
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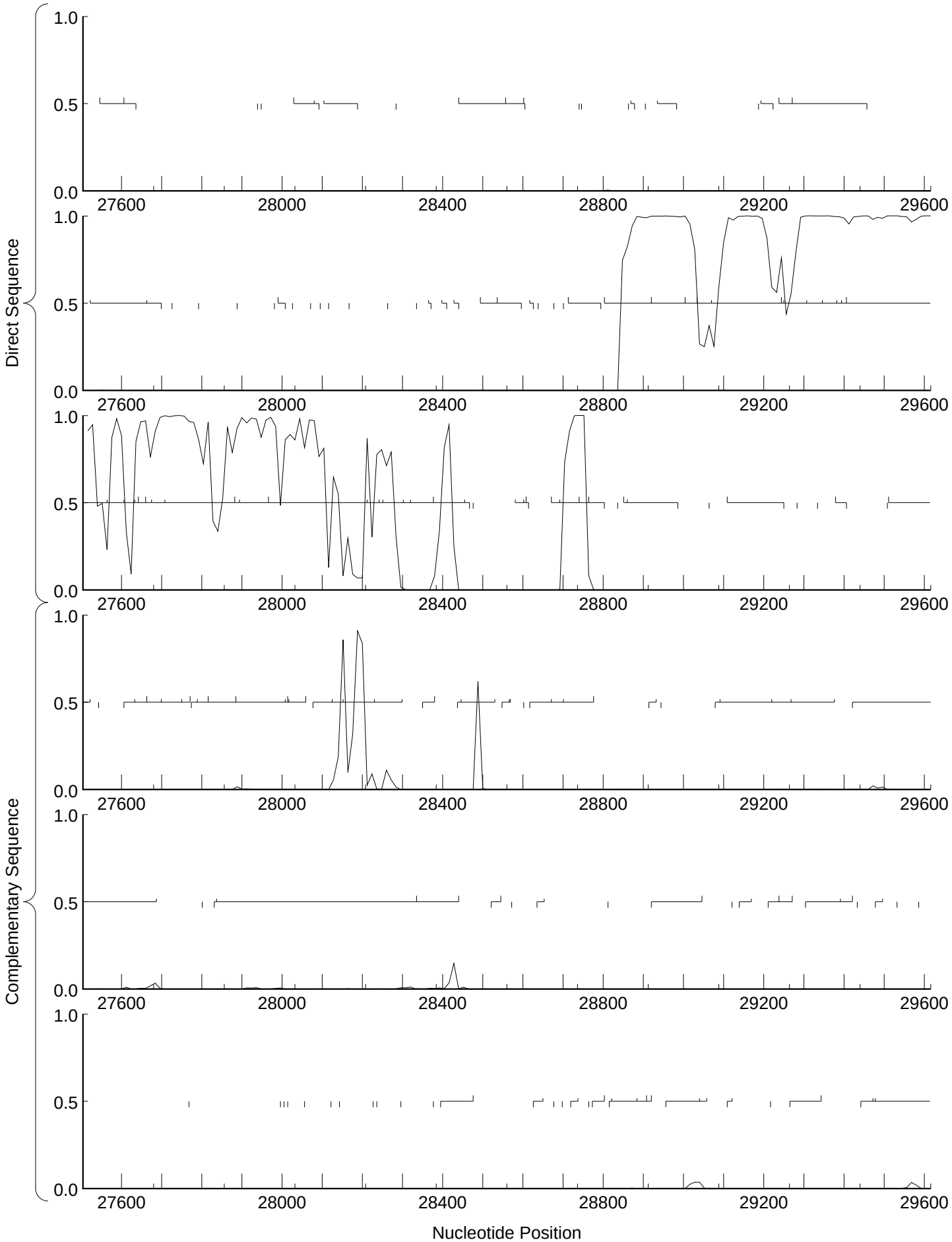
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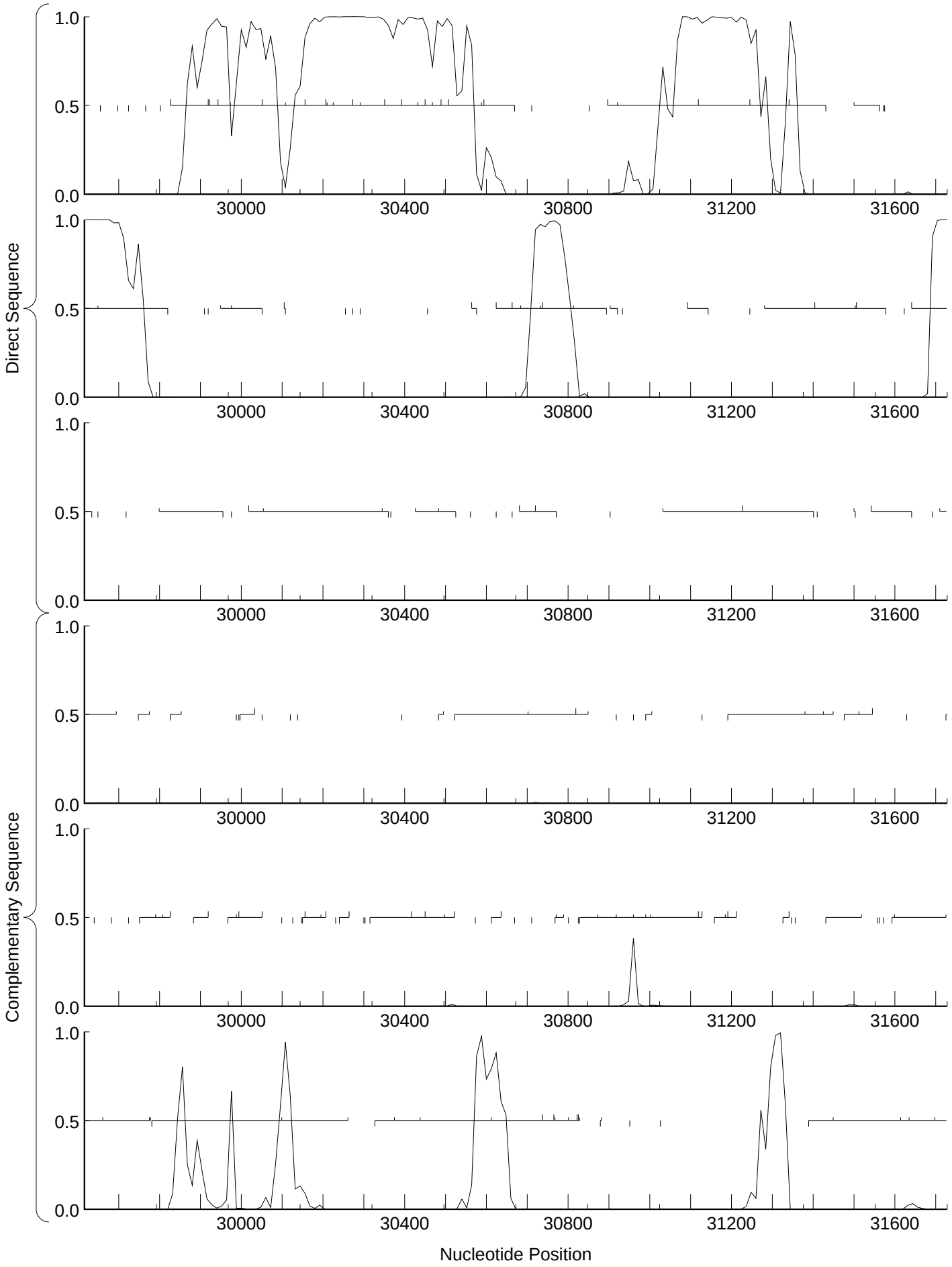
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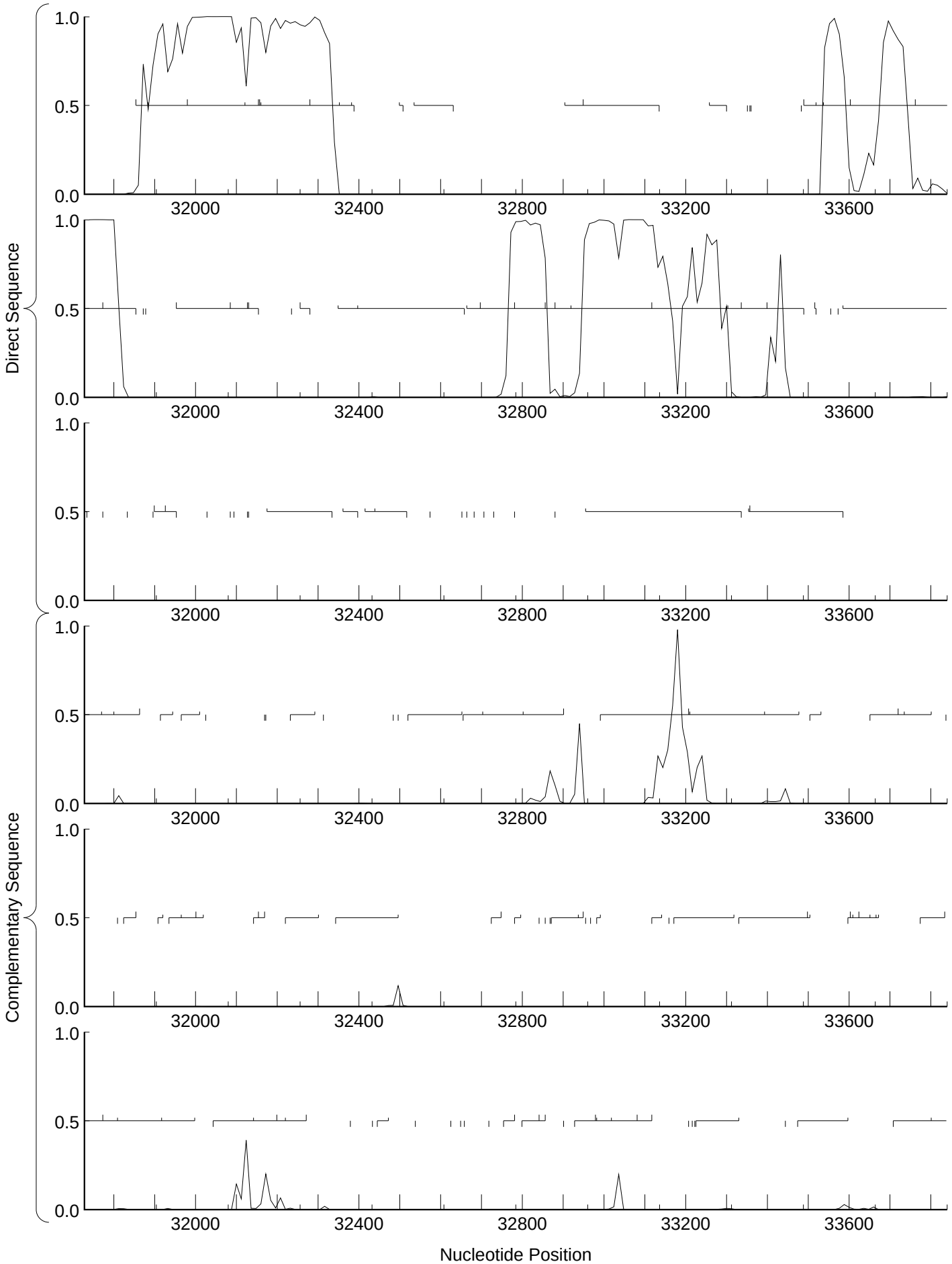
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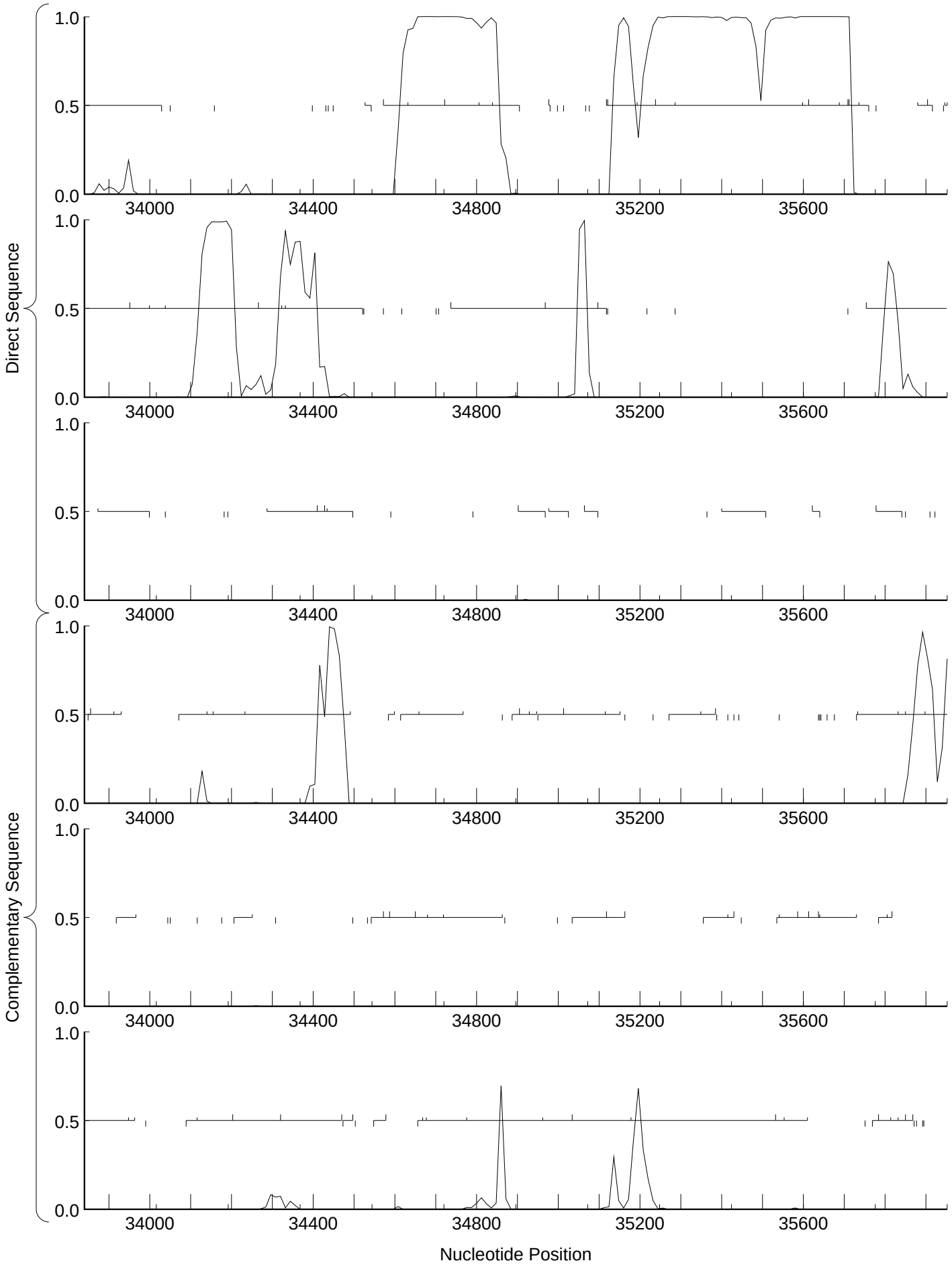
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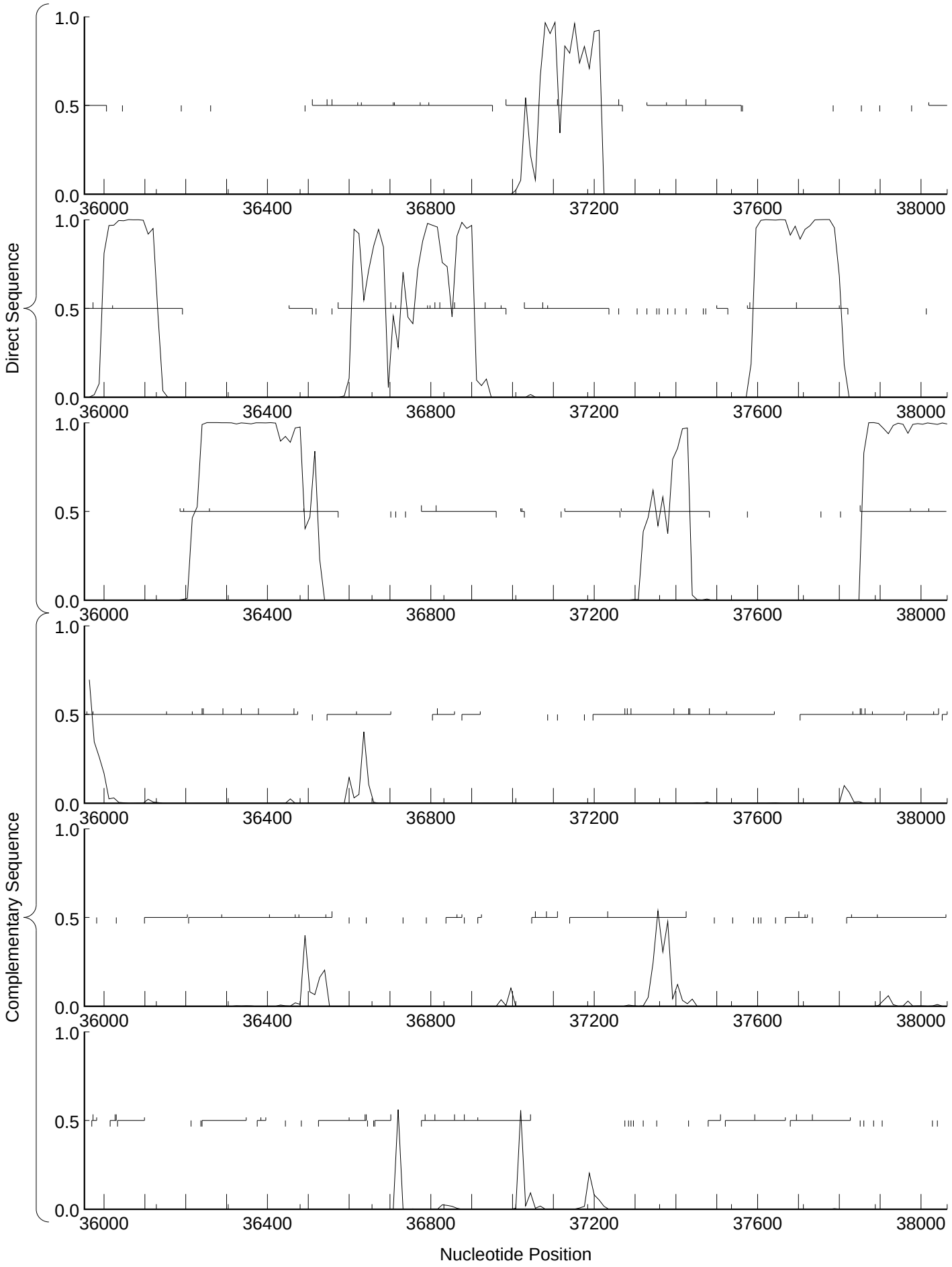
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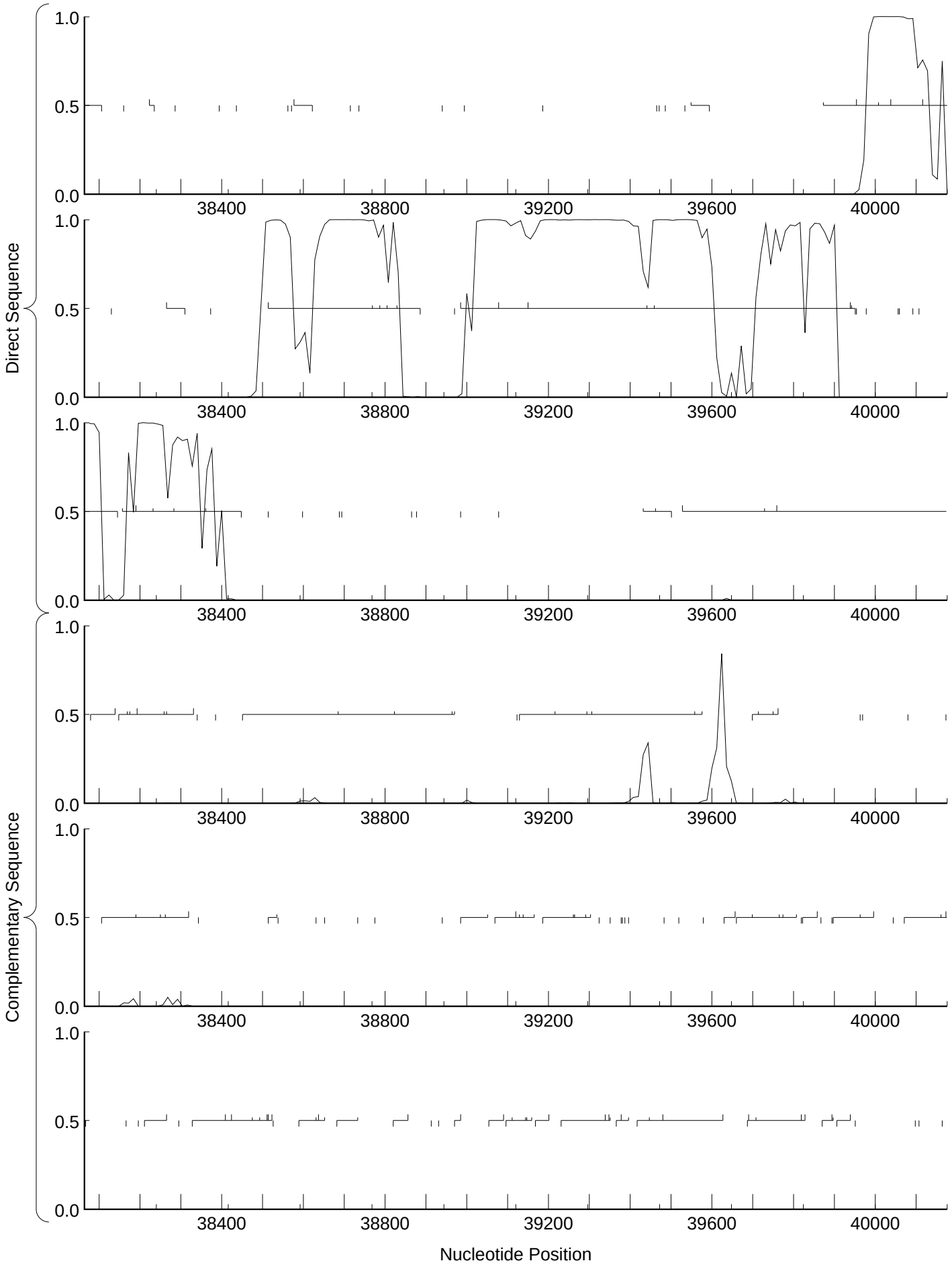
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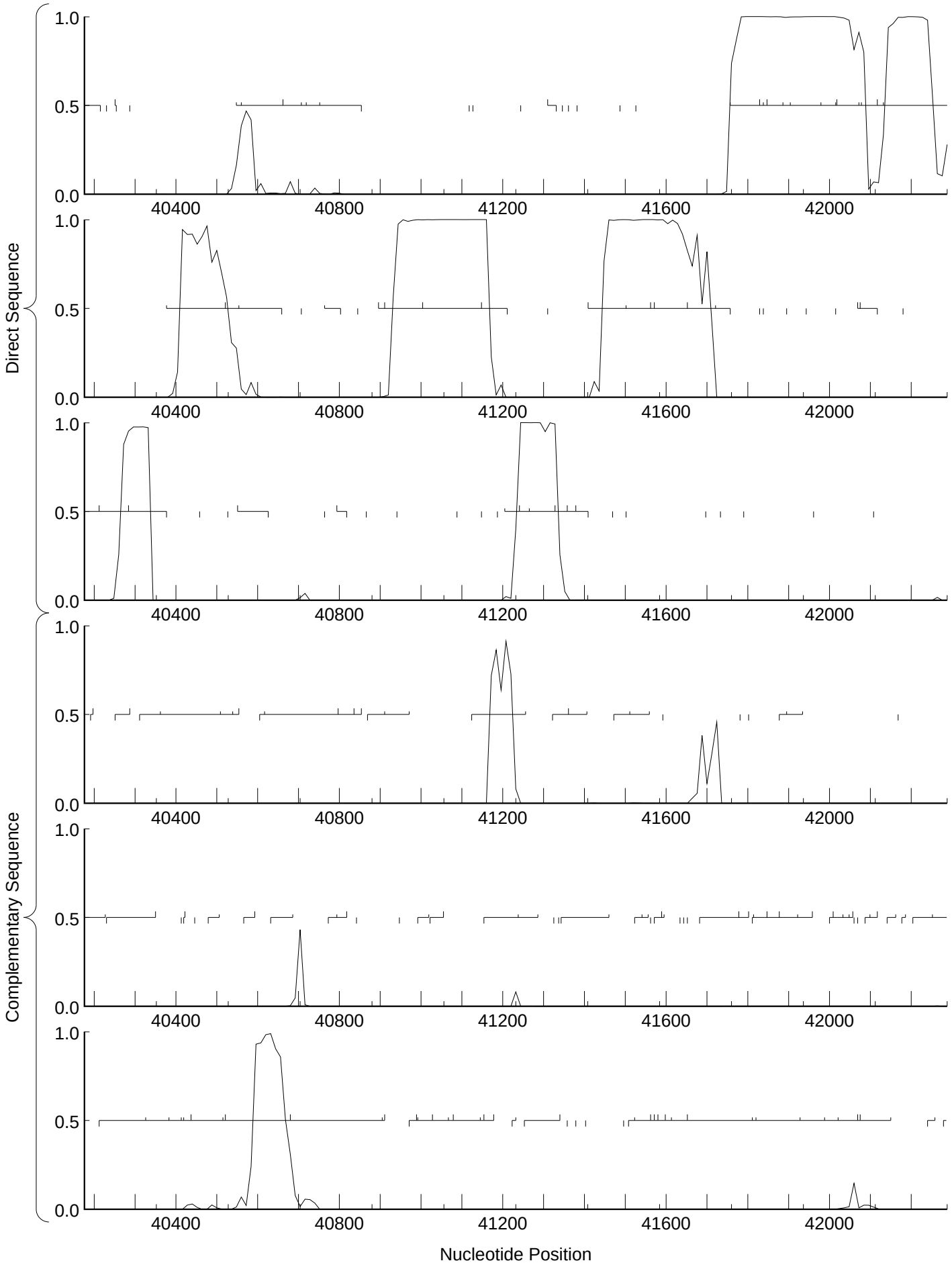
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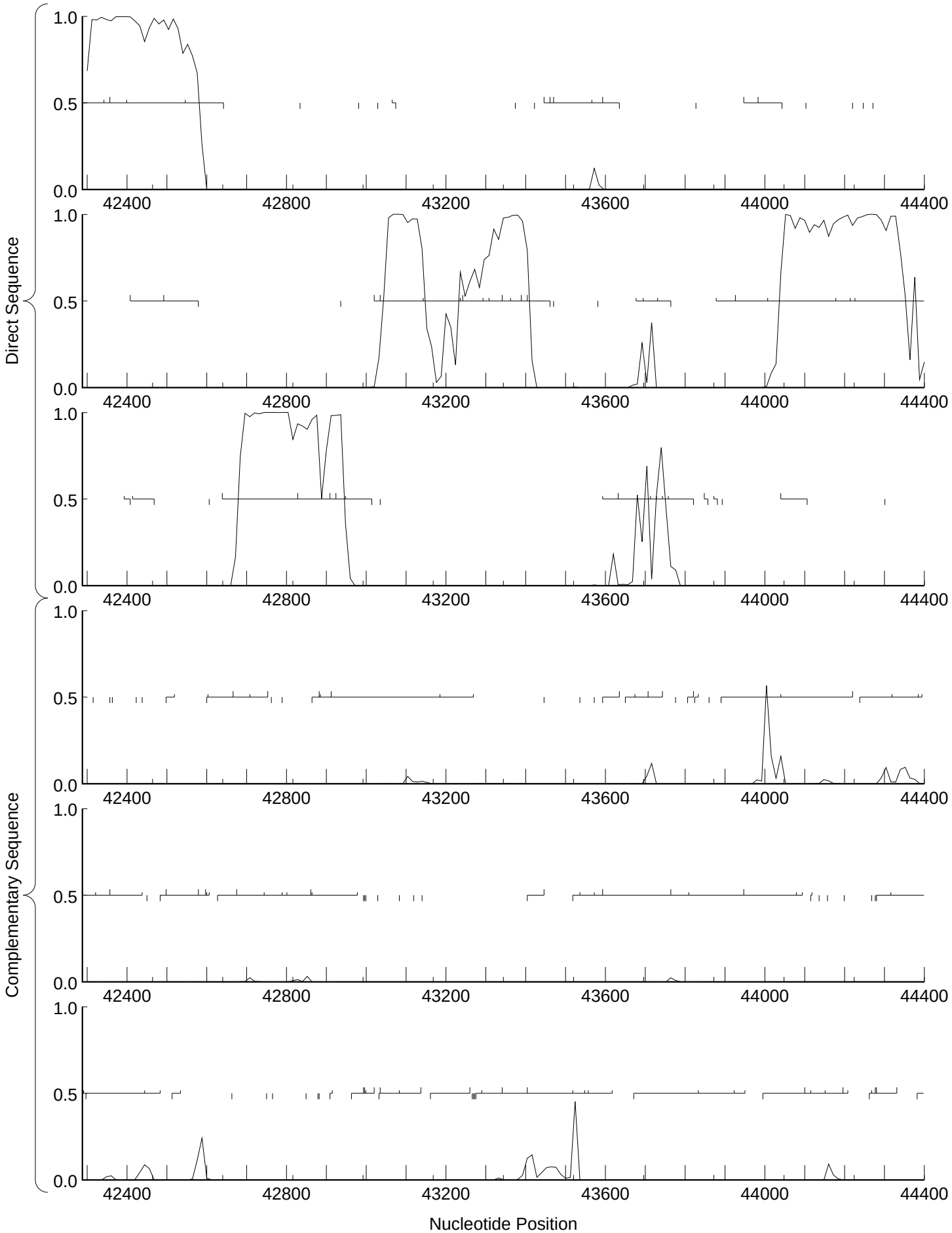


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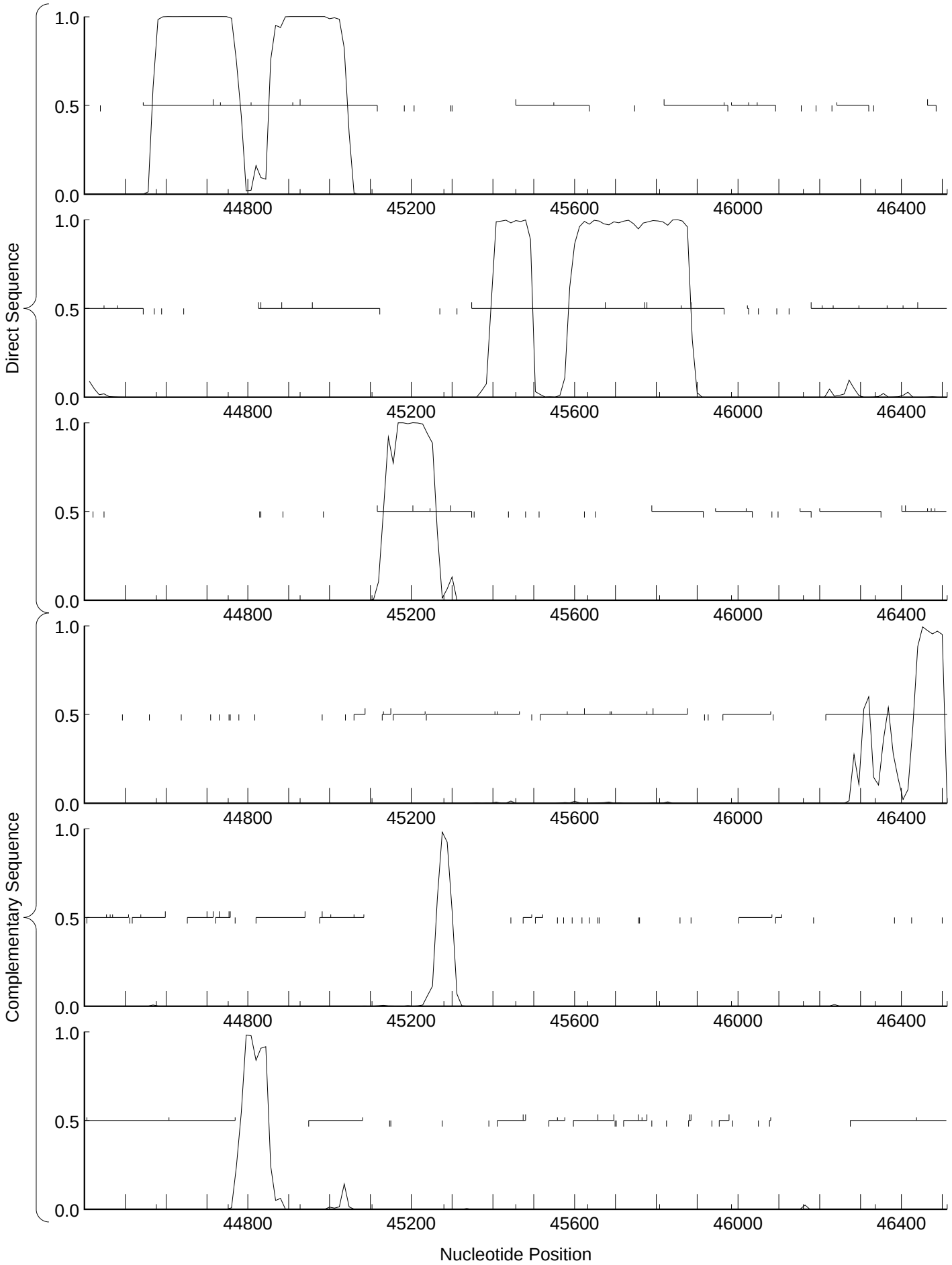


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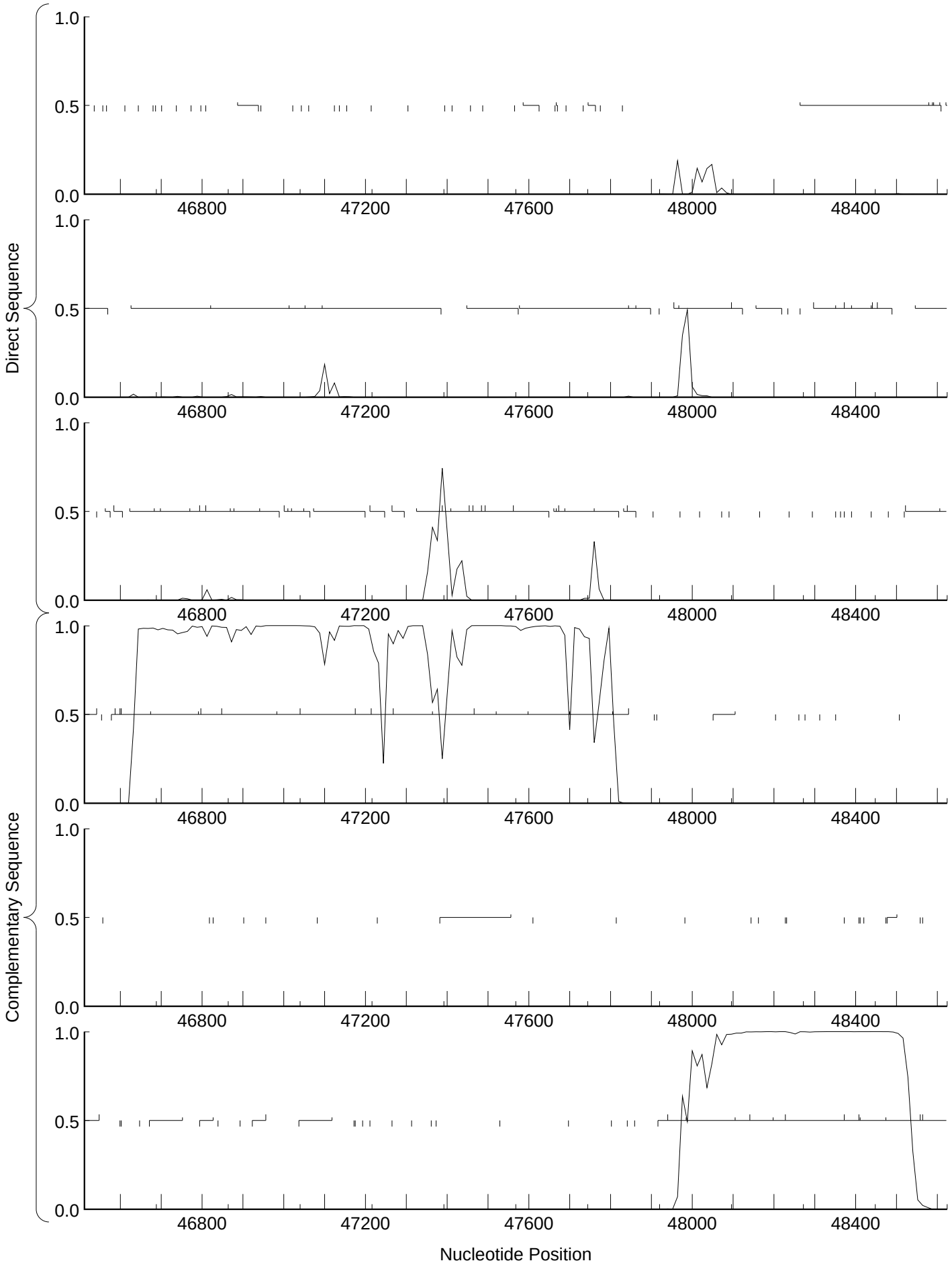



Direct Sequence
Complementary Sequence
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0.5
0.0
42400
42800
43200
43600
44000
44400
Nucleotide Position

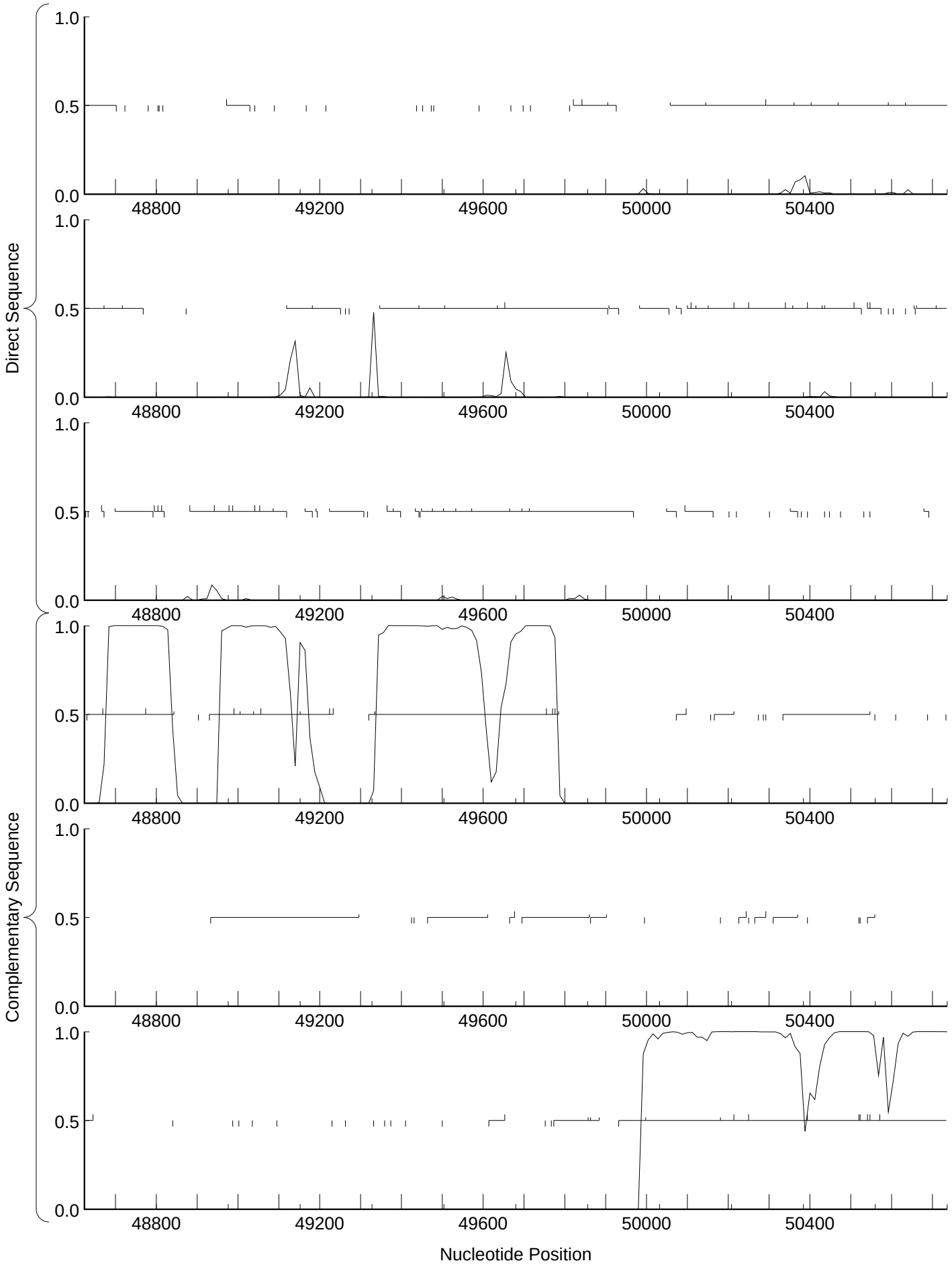
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bacter phage BarretLemon complete sequence, 51290 bp, circularly permuted, Cluster AO2, Order 4, Window 96, Step 12, 24/26



bacter phage BarretLemon complete sequence, 51290 bp, circularly permuted, Cluster AO2, Order 4, Window 96, Step 12, 25/26



bacter phage BarretLemon complete sequence, 51290 bp, circularly permuted, Cluster AO2, Order 4, Window 96, Step 12, 26/26

