GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Kwksand96 complete sequence, 68822 bp, circularly permuted, Cluster B1
Analysis Date    :  3/28/24 at 8:17:58
Pages            :  34
Sequence Length  :  68822 bp
GC Content       :   66.35%

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

PS-Version       :  1.2

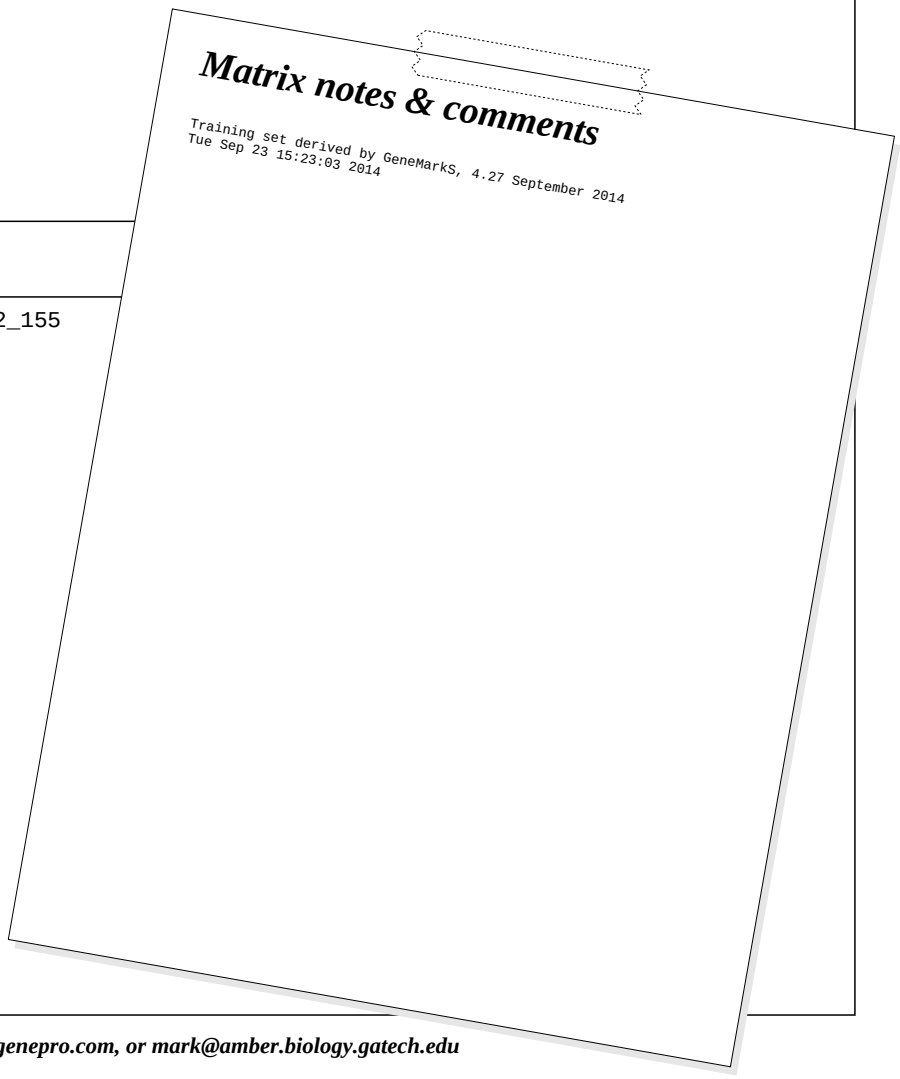
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Author     :  -

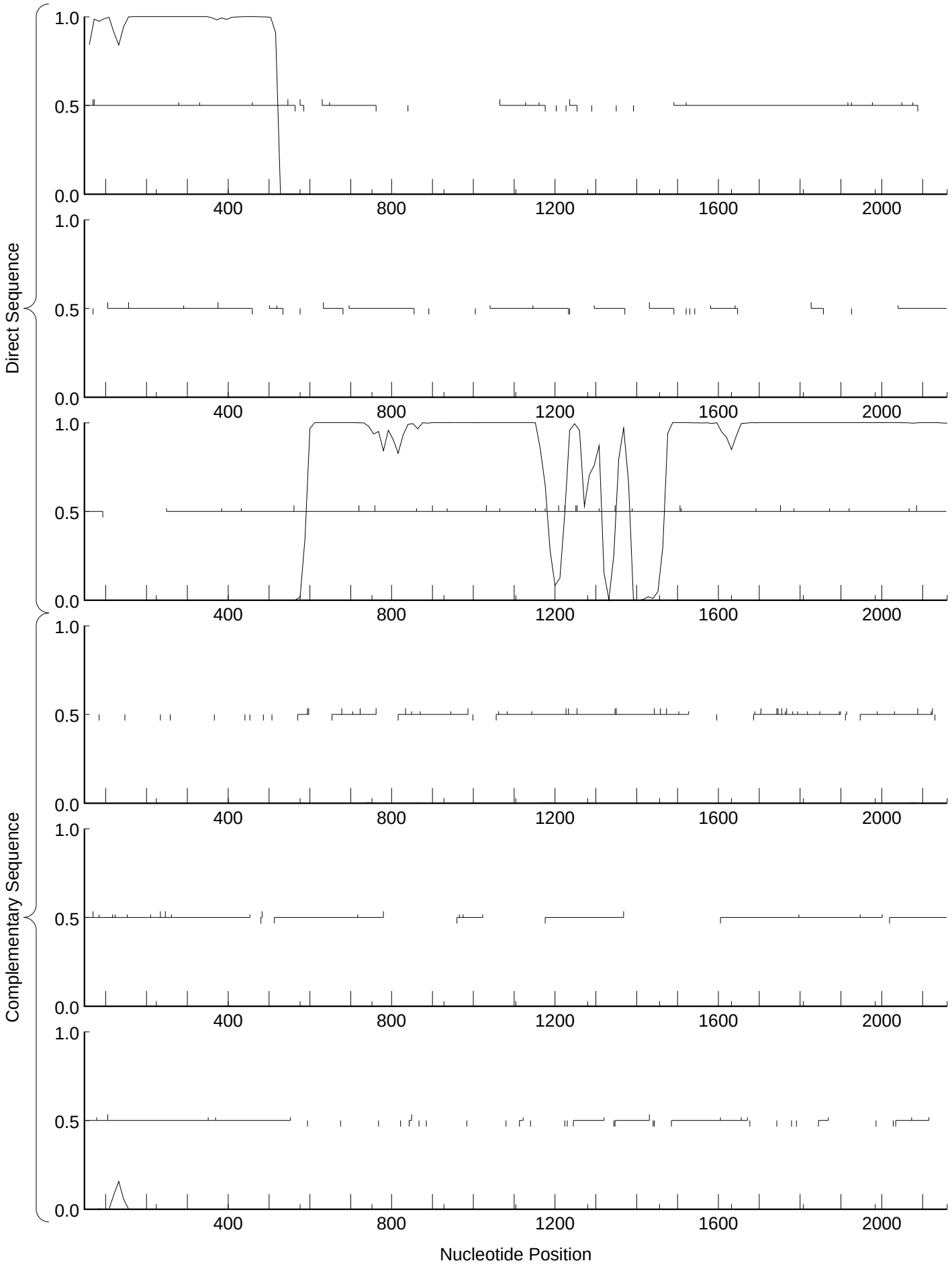
Order      :  4


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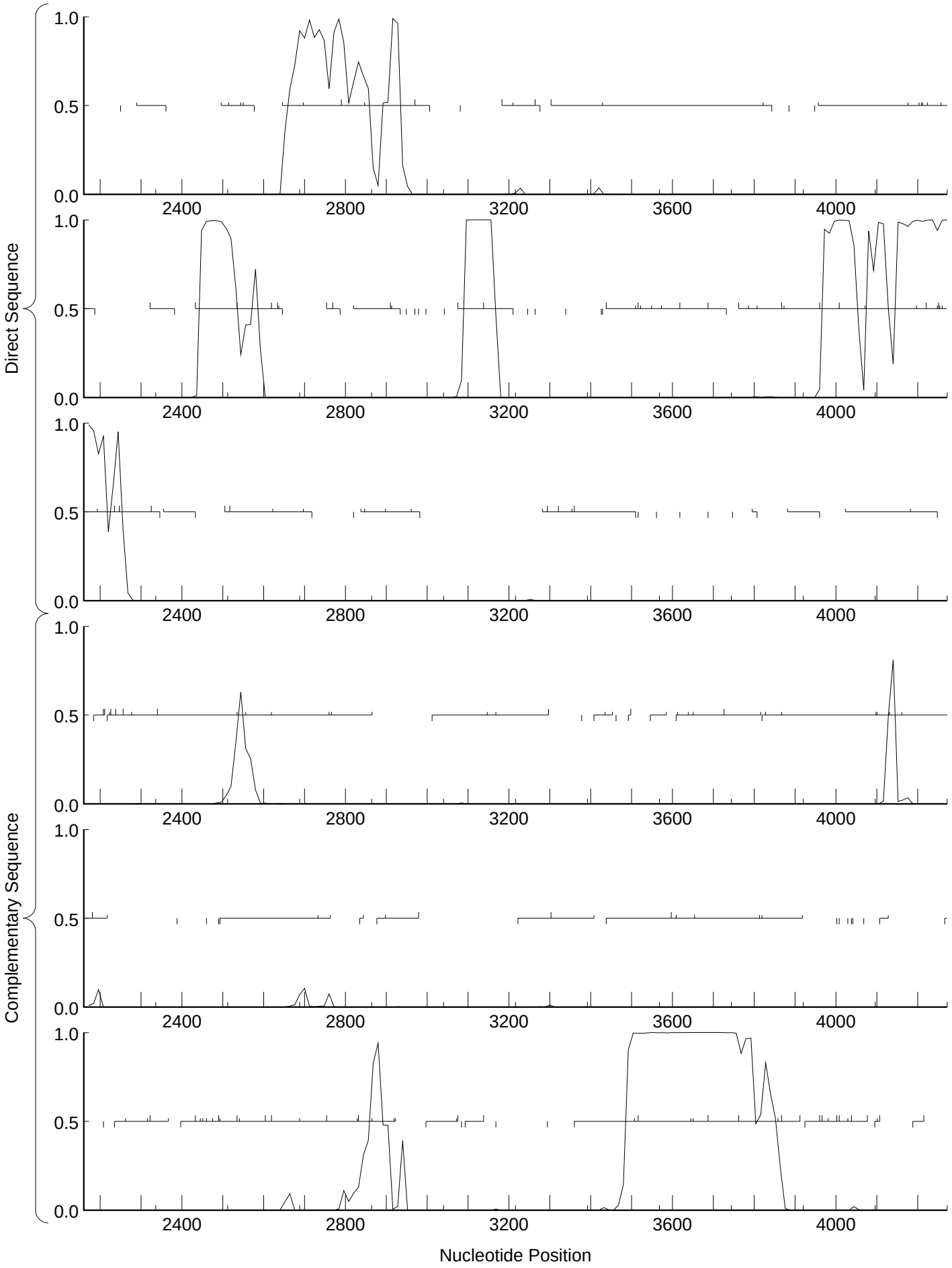


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

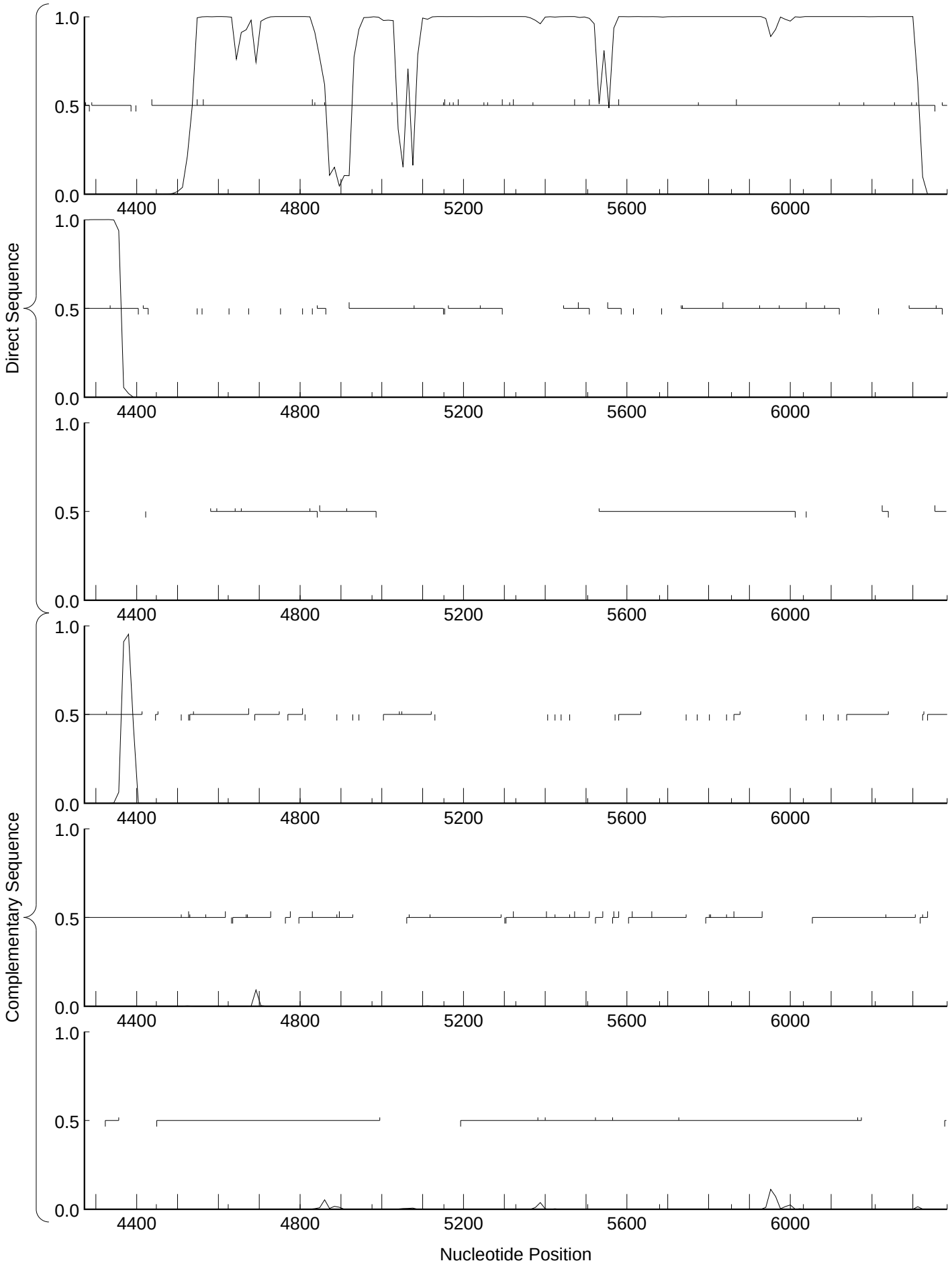
obacterium phage Kwksand96 complete sequence, 68822 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 2/34



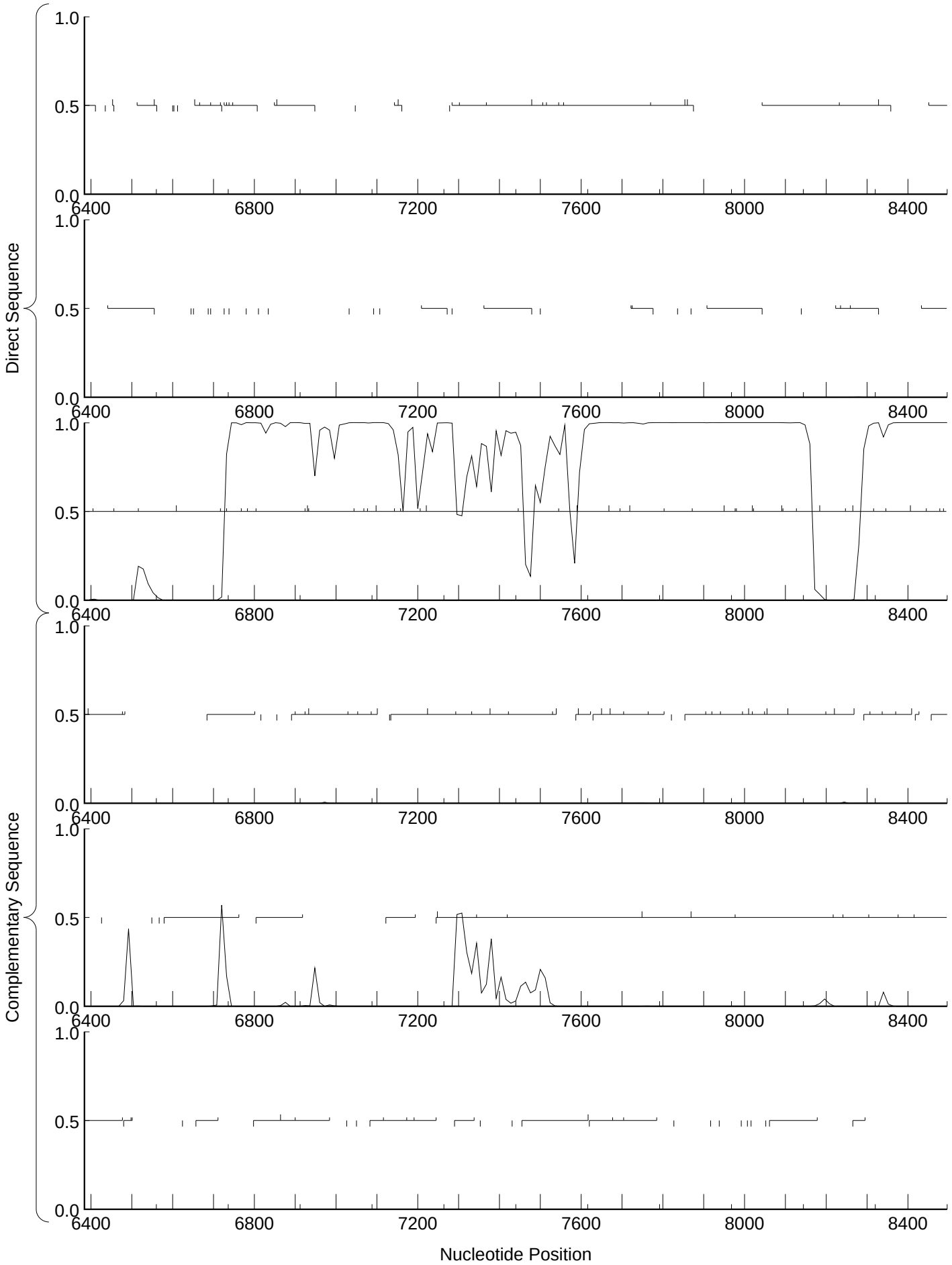
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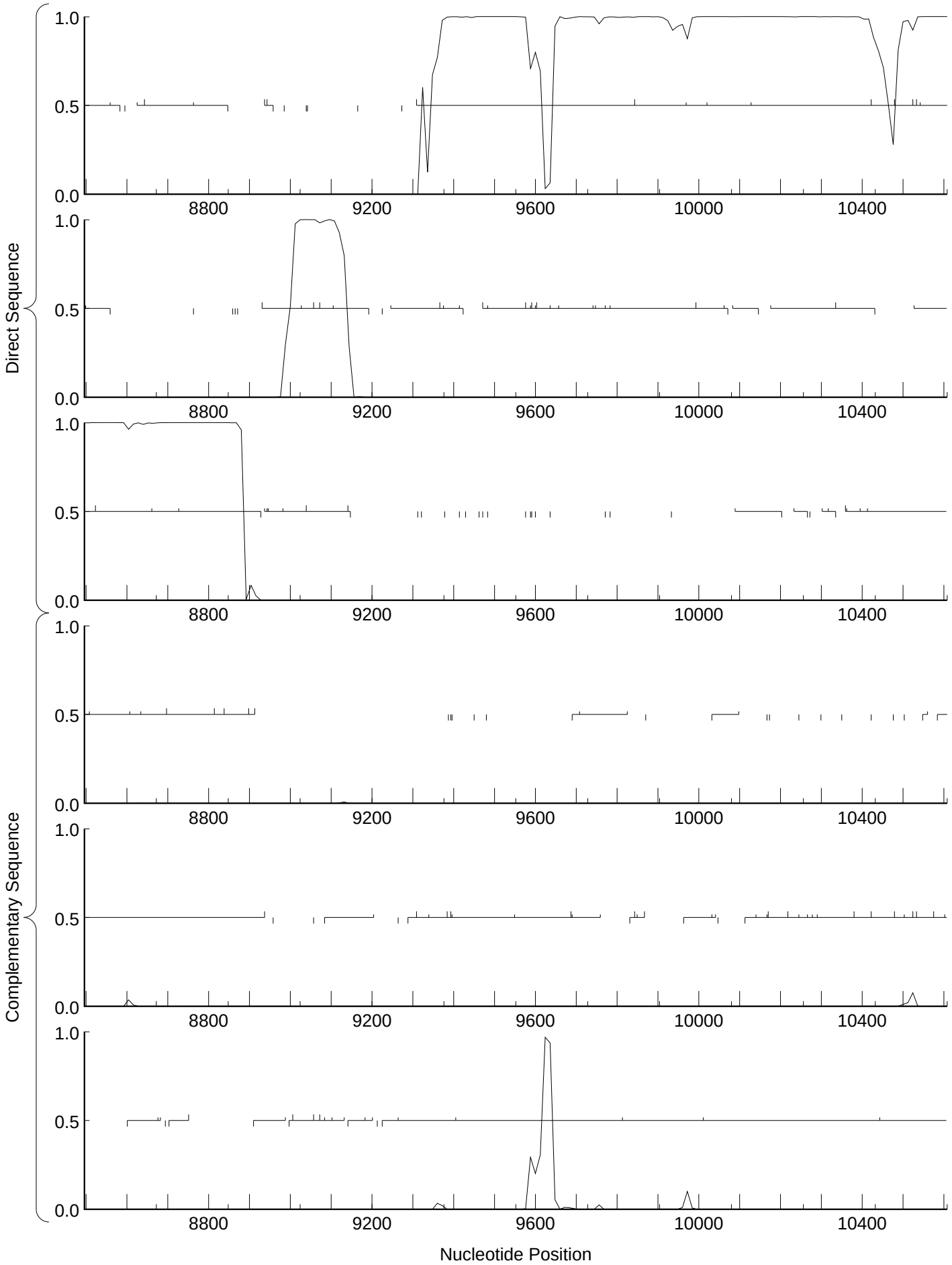
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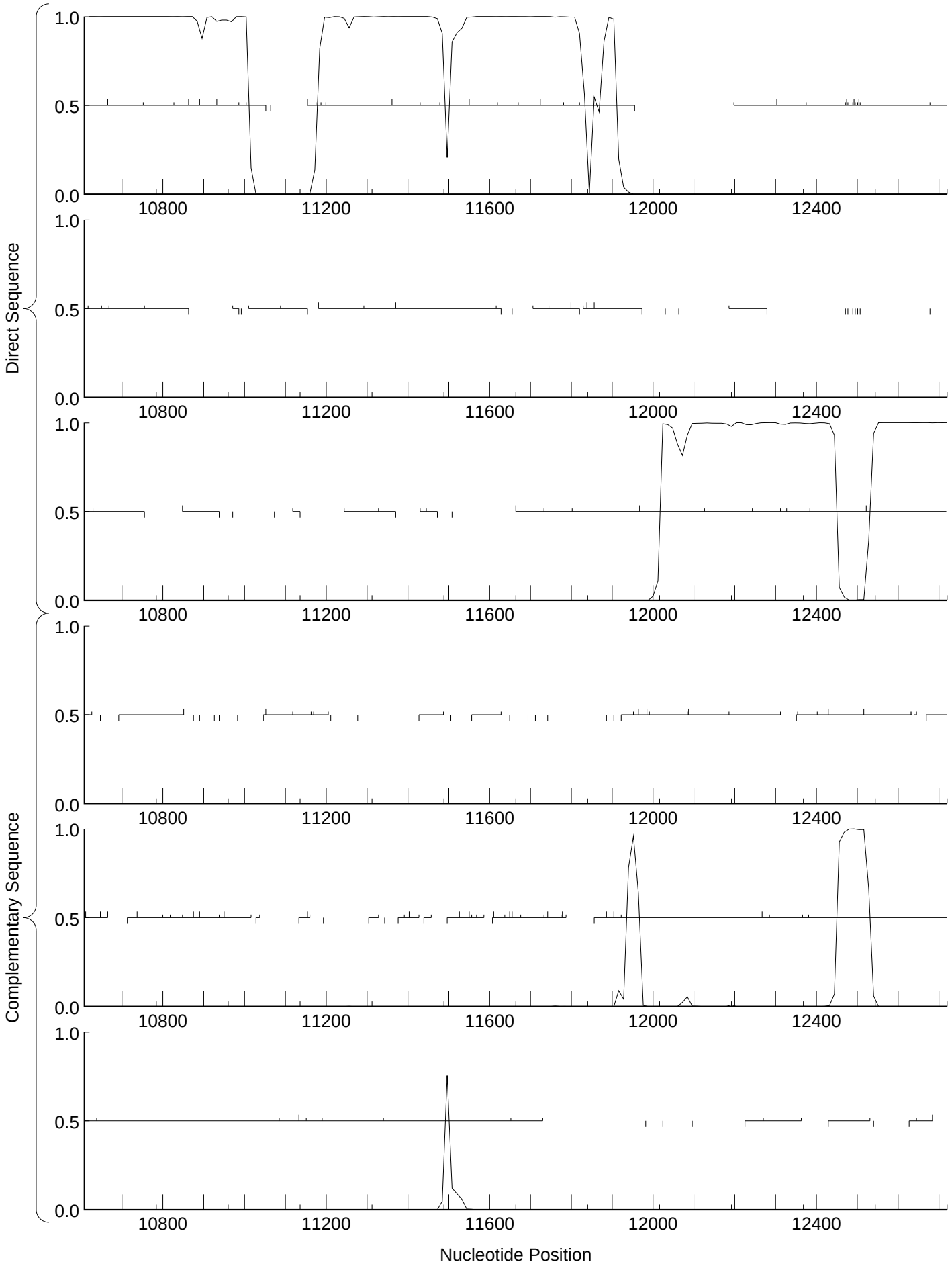
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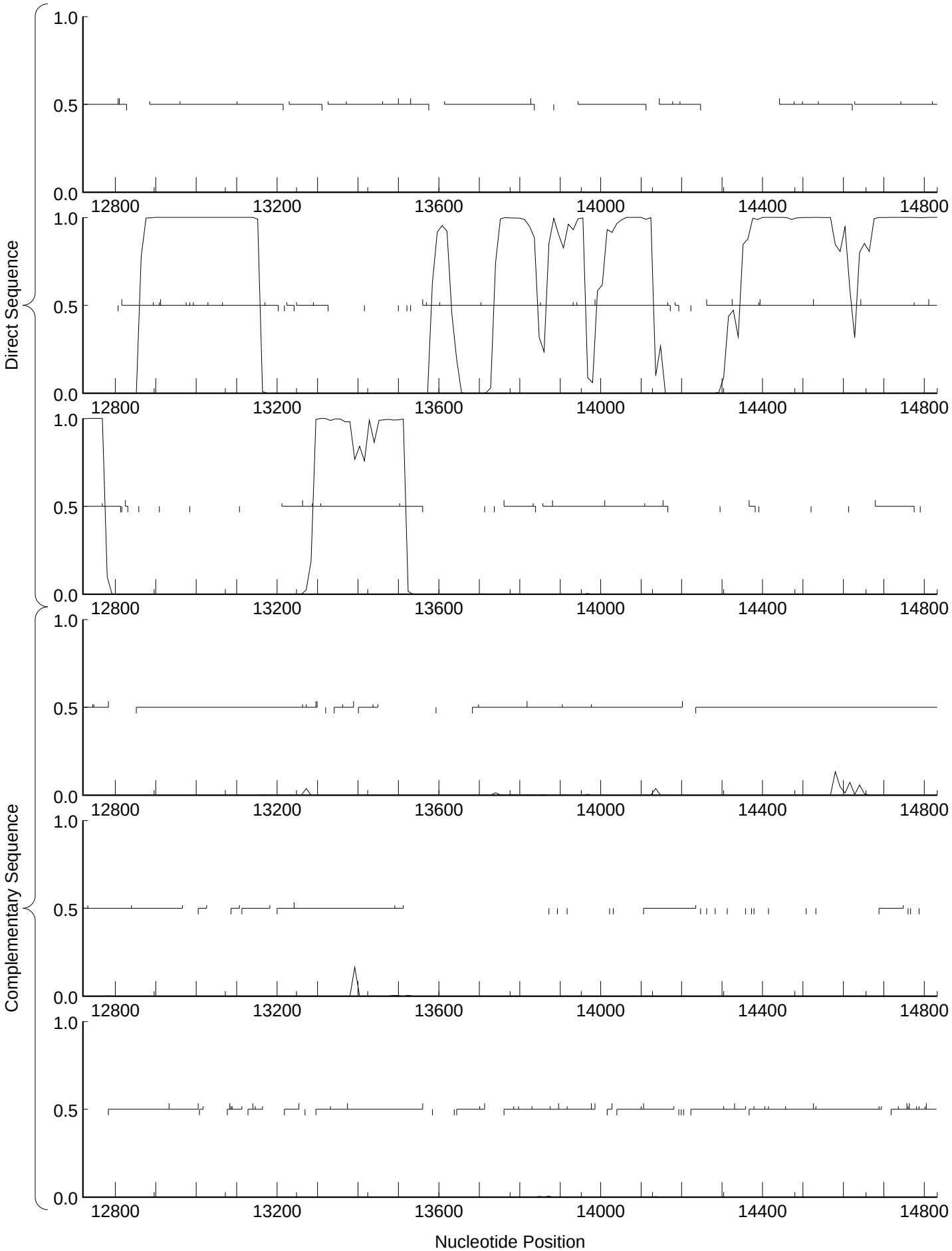
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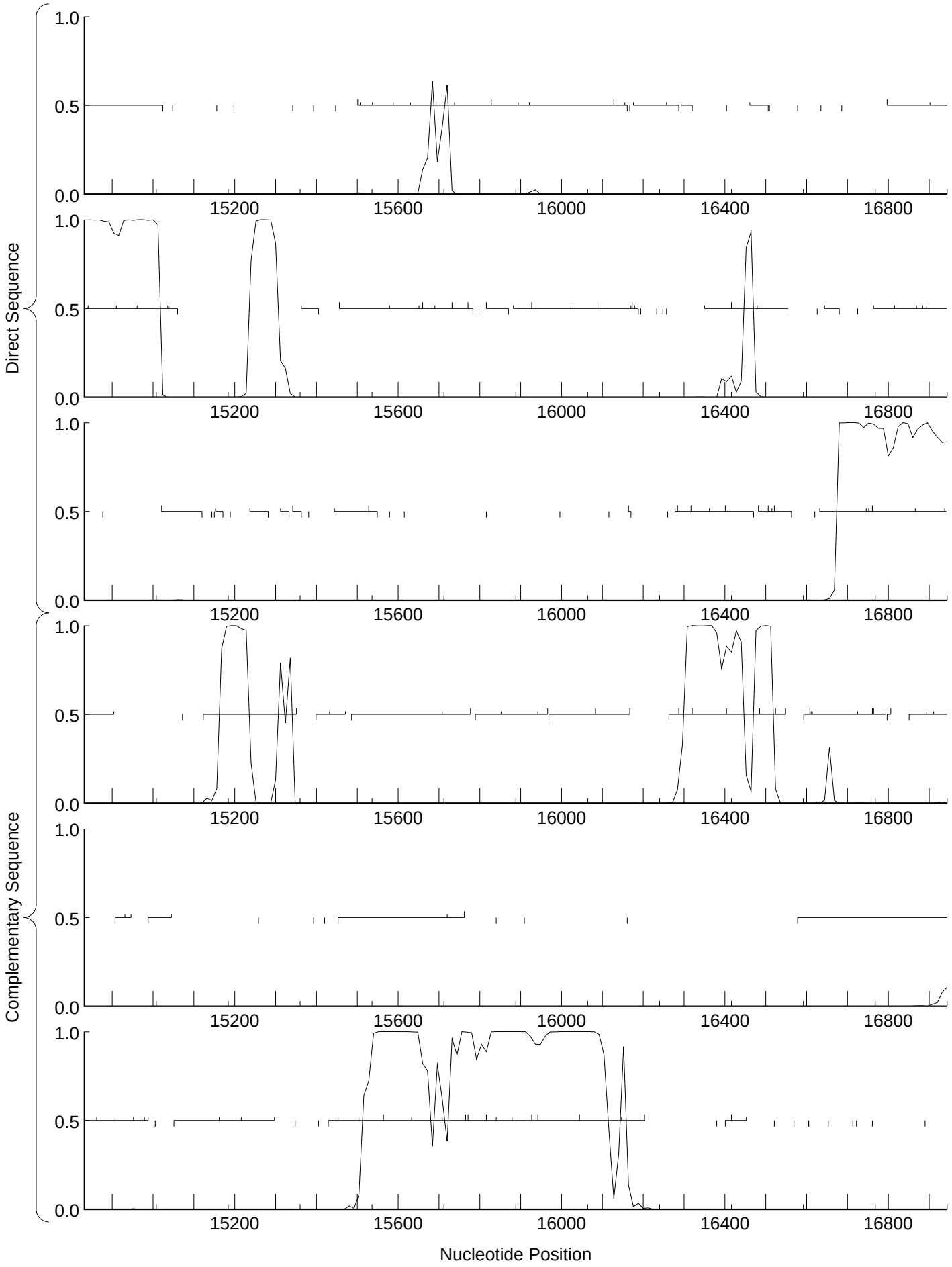
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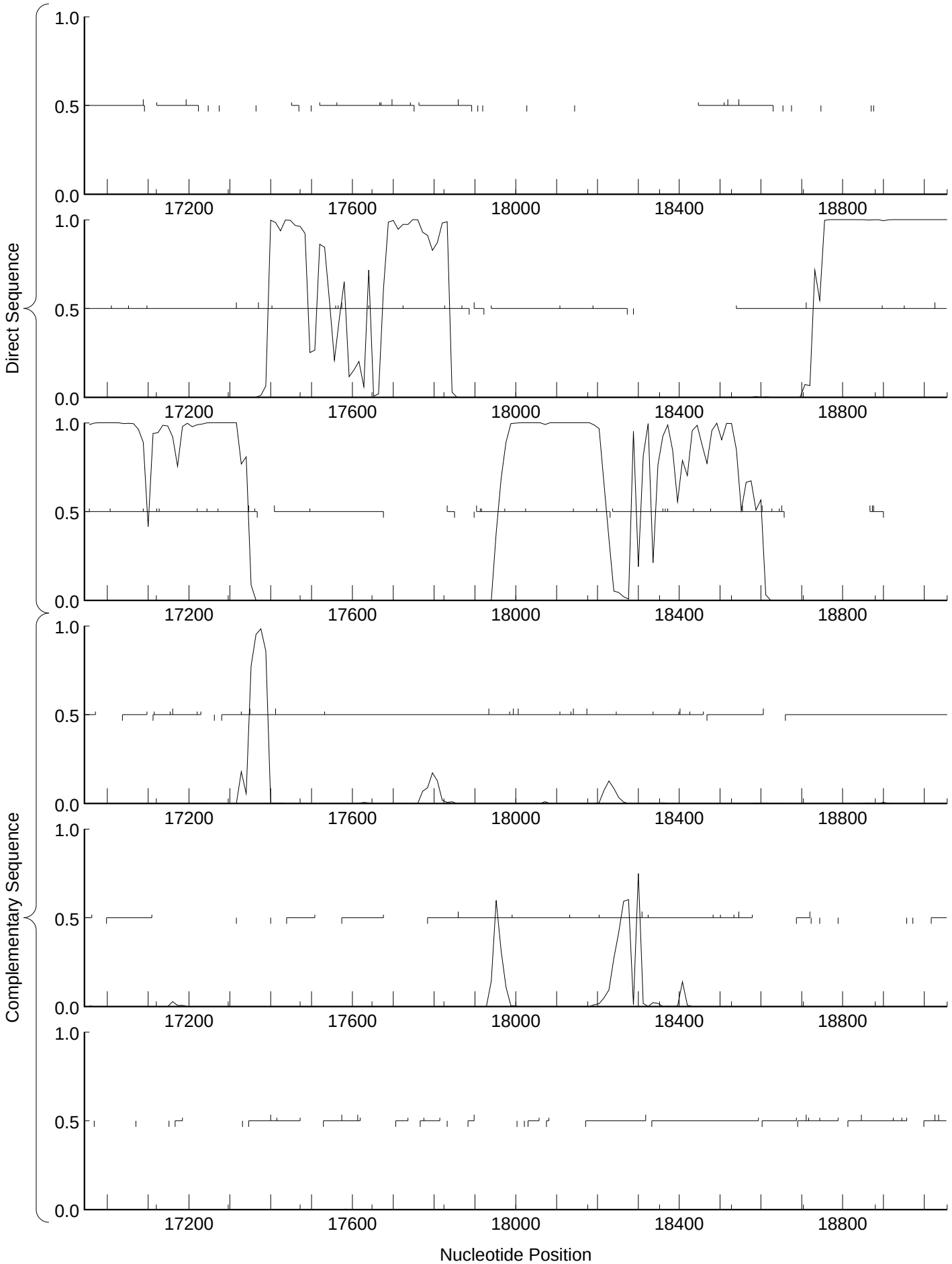
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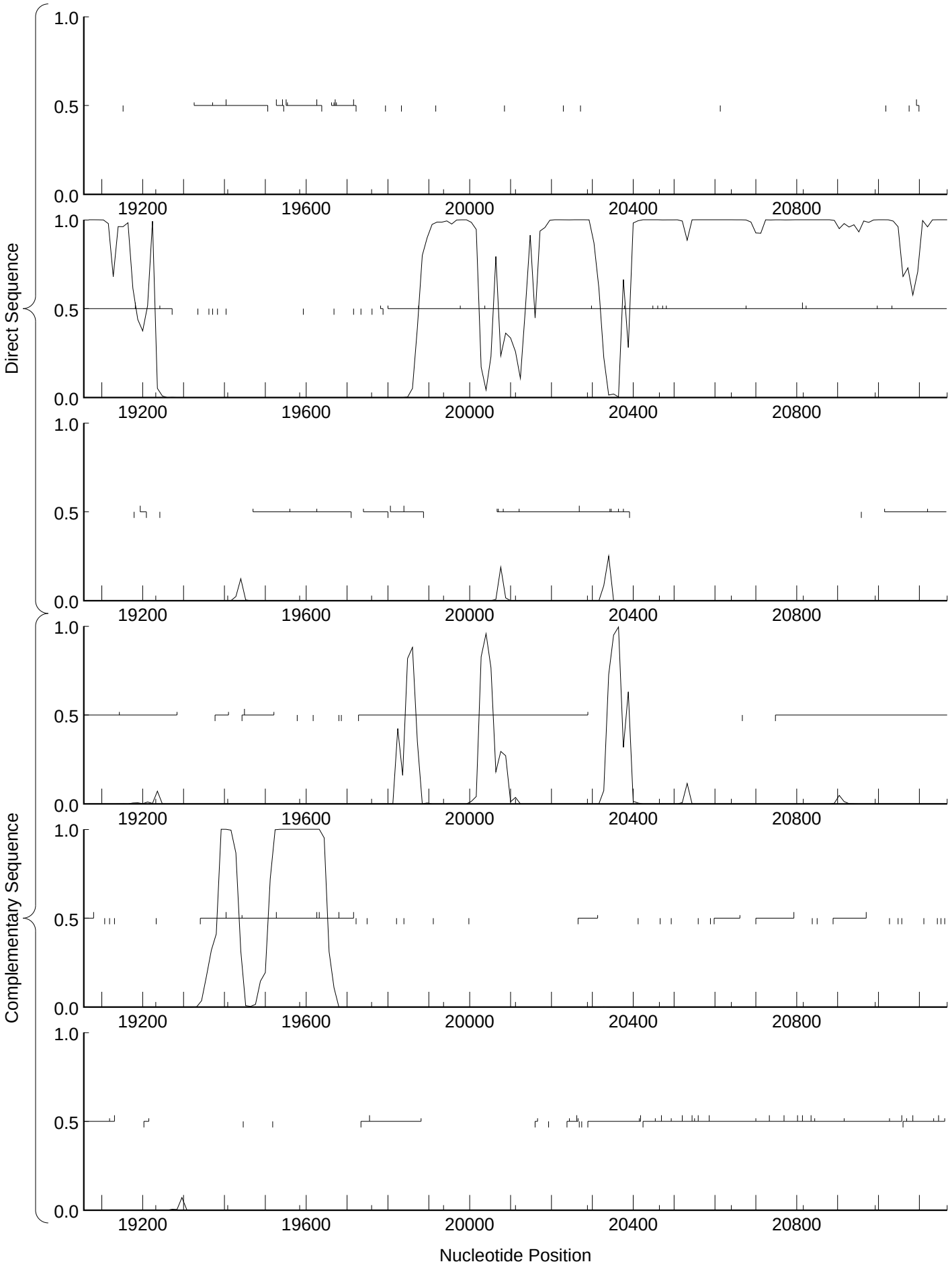
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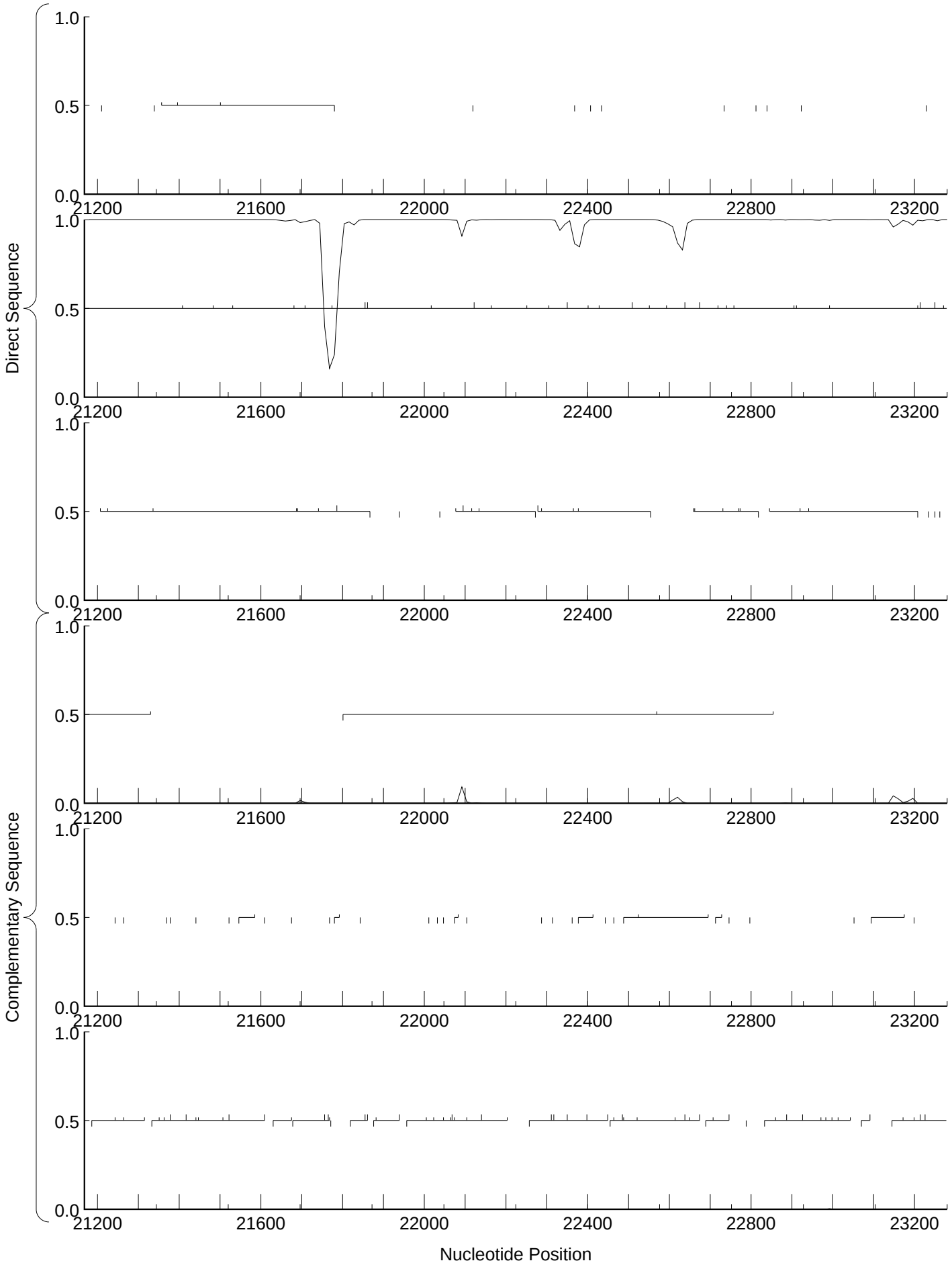
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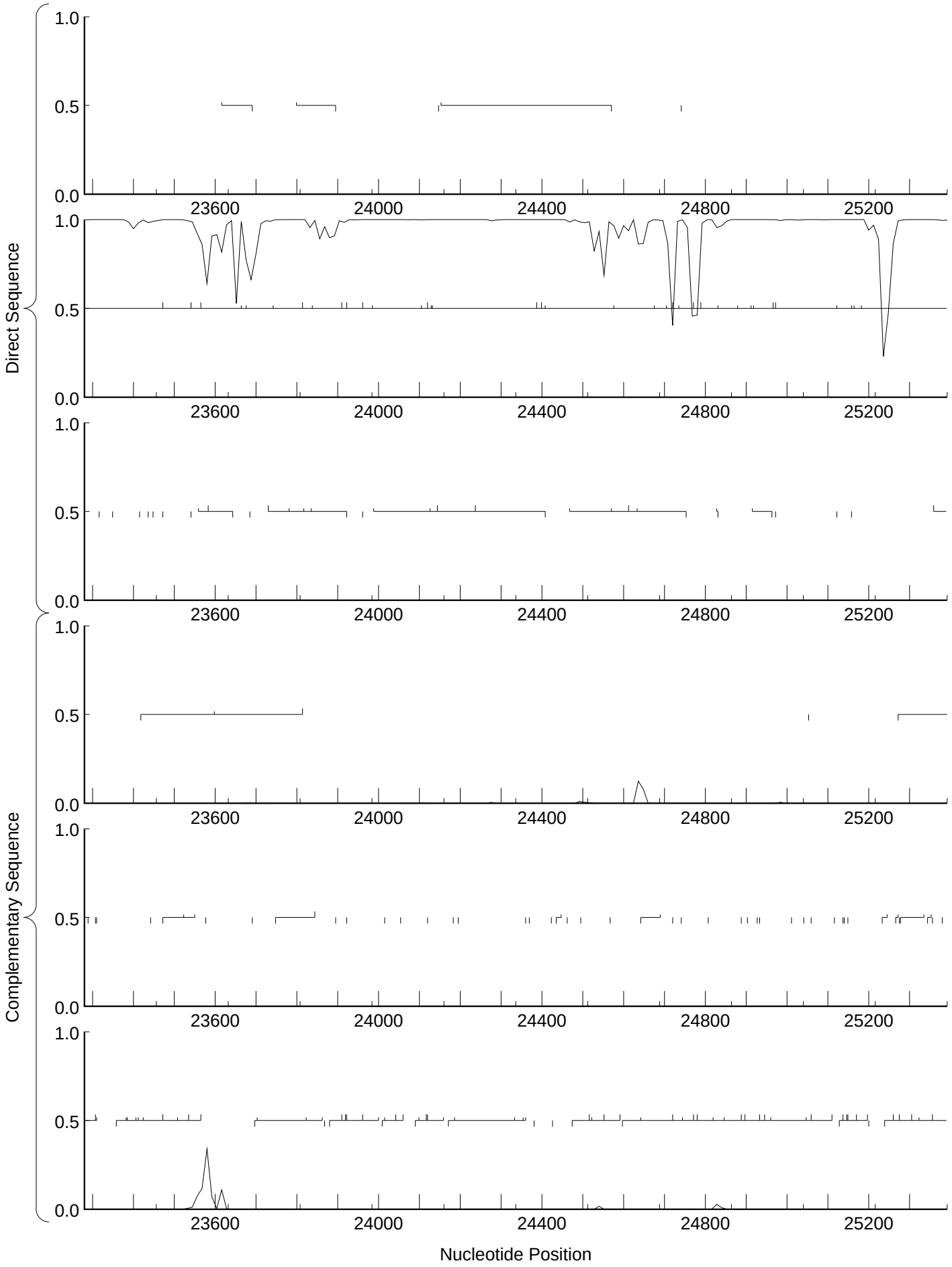
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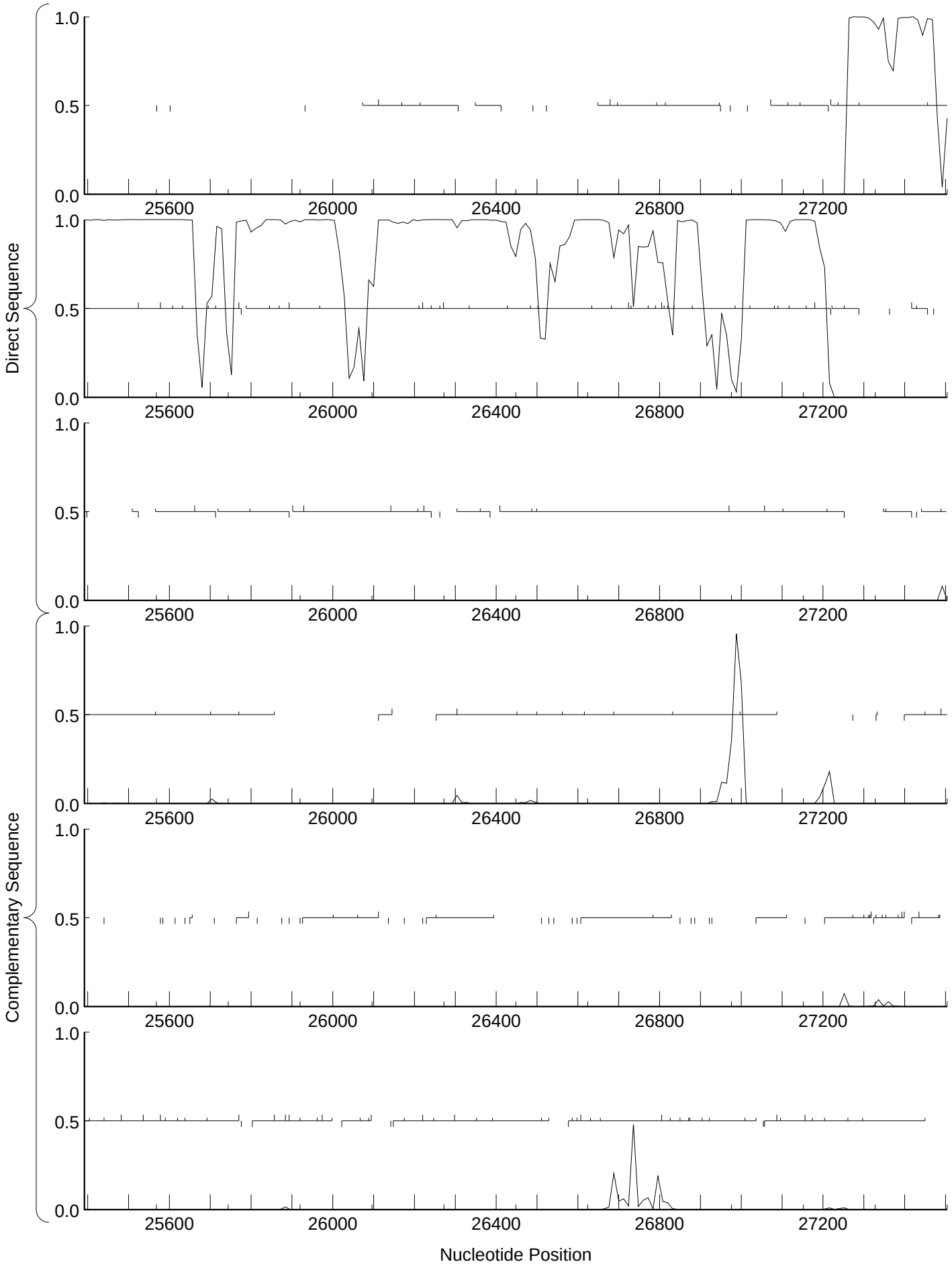


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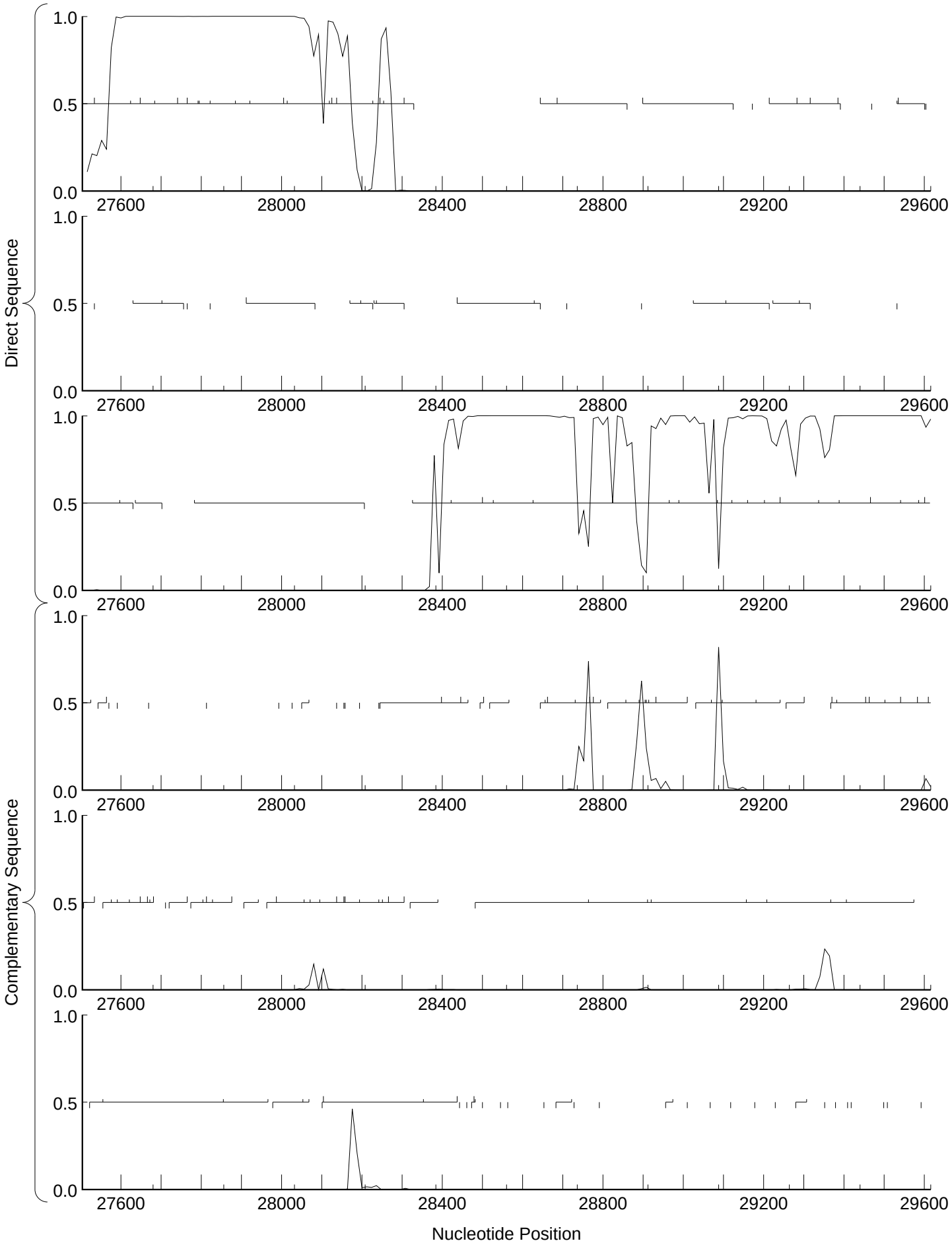


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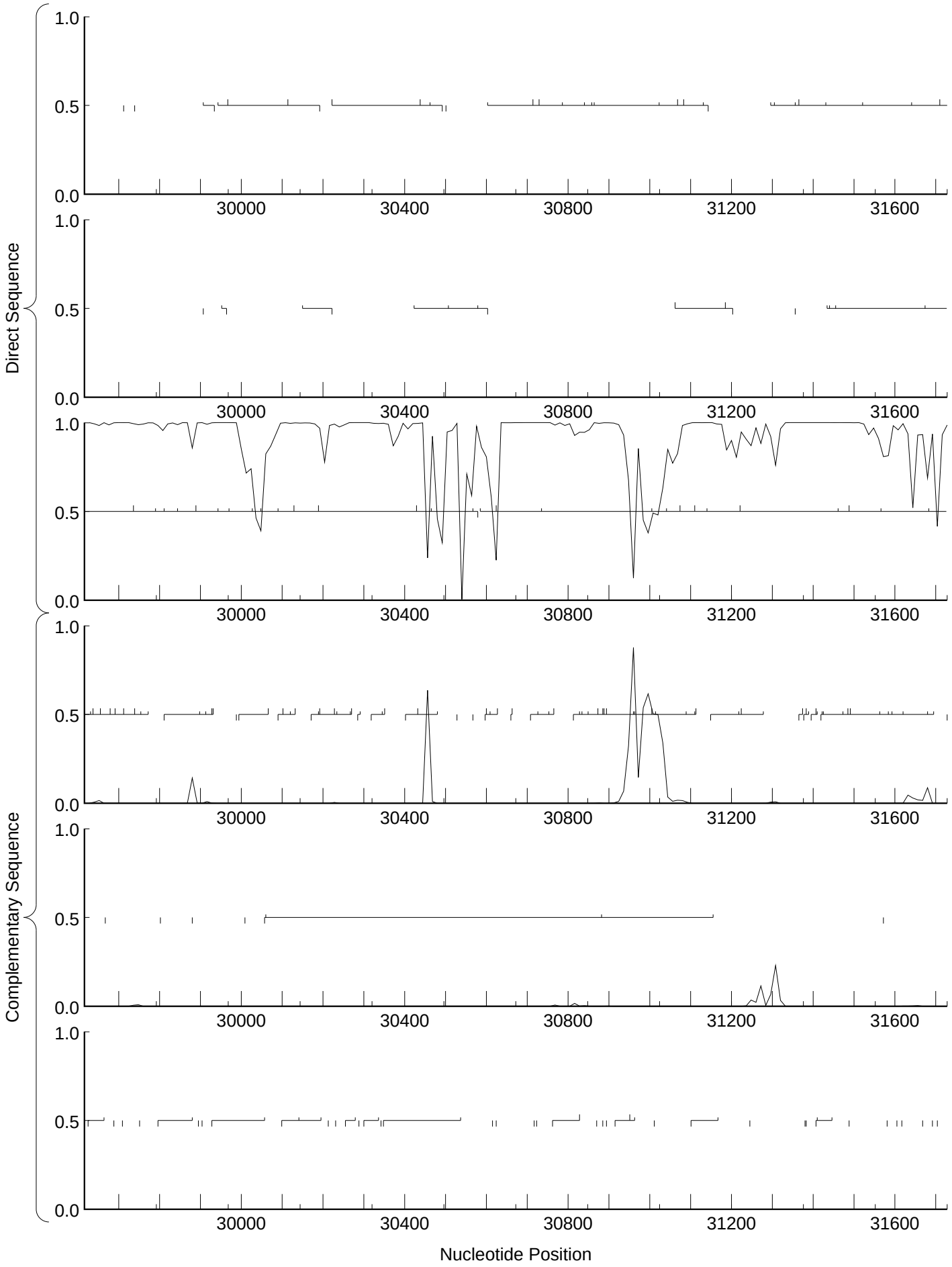



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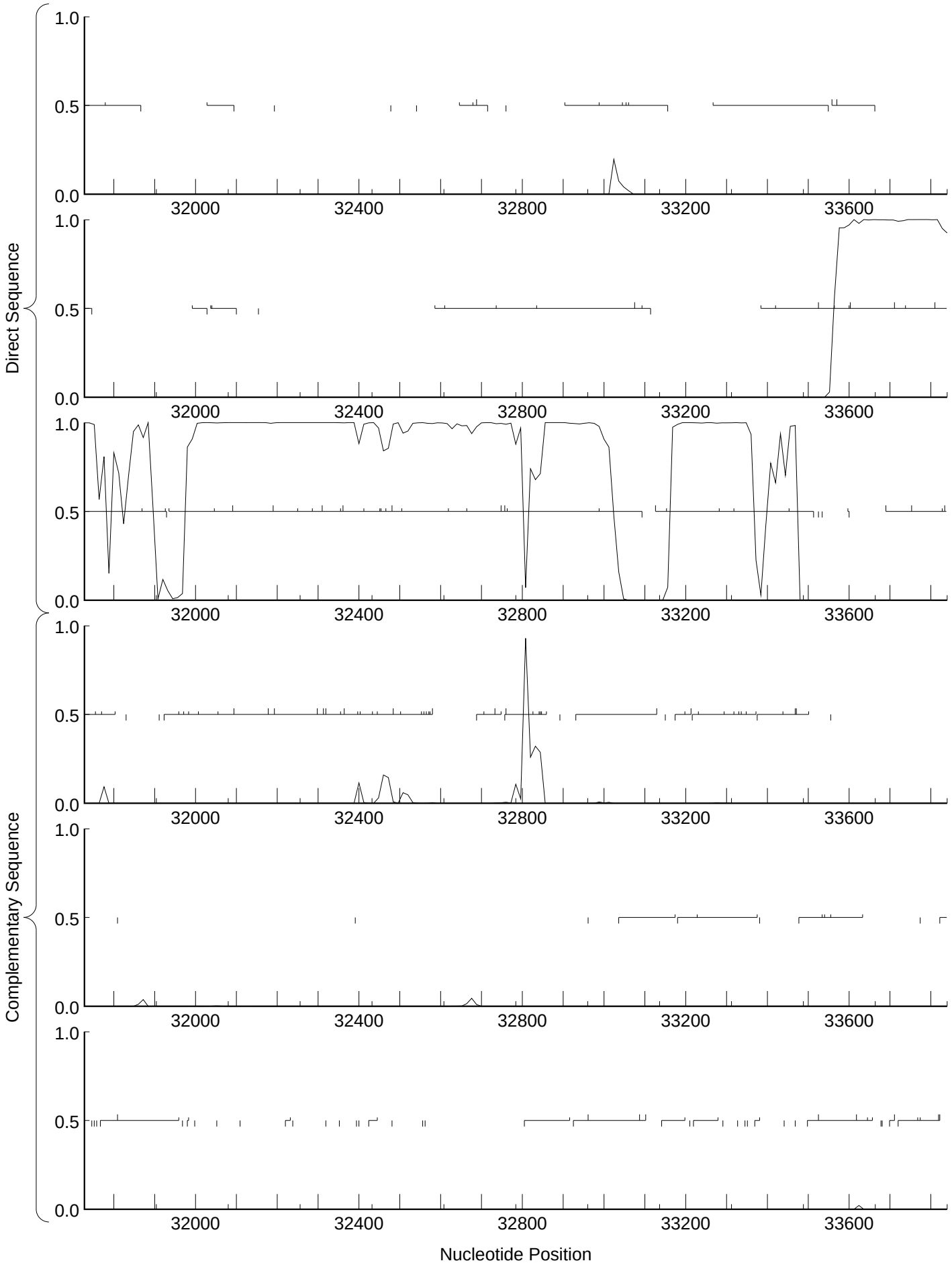
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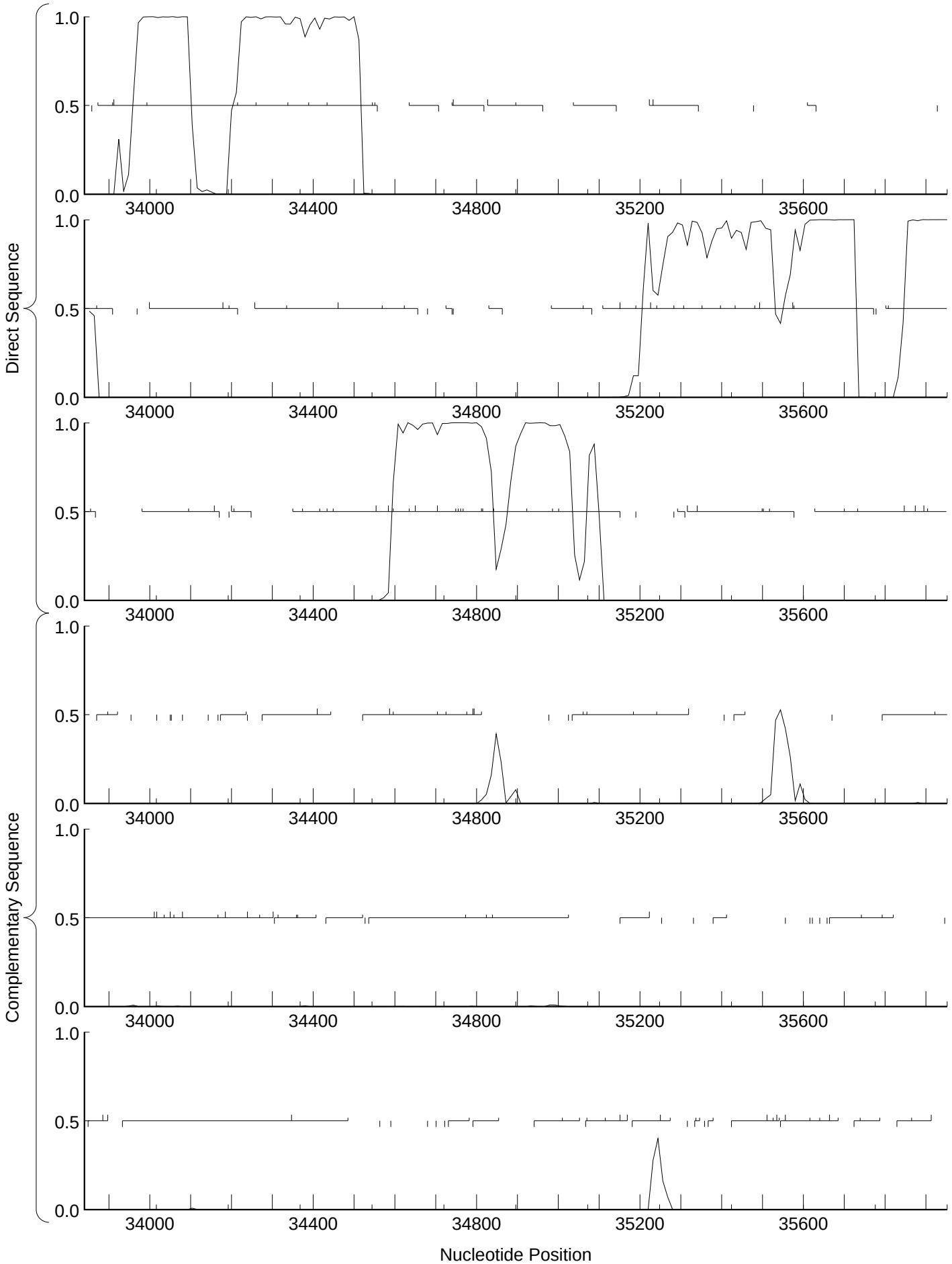


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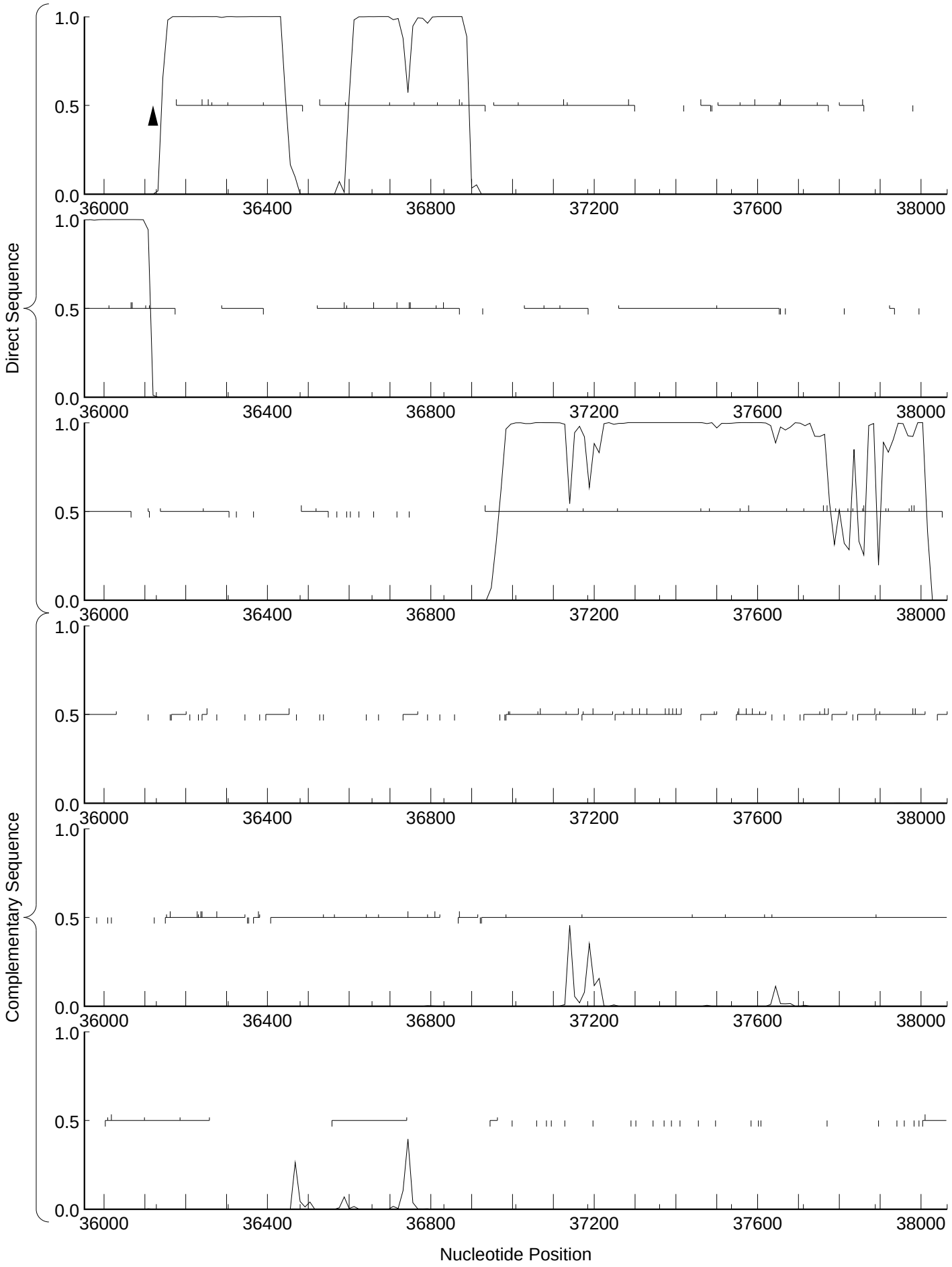


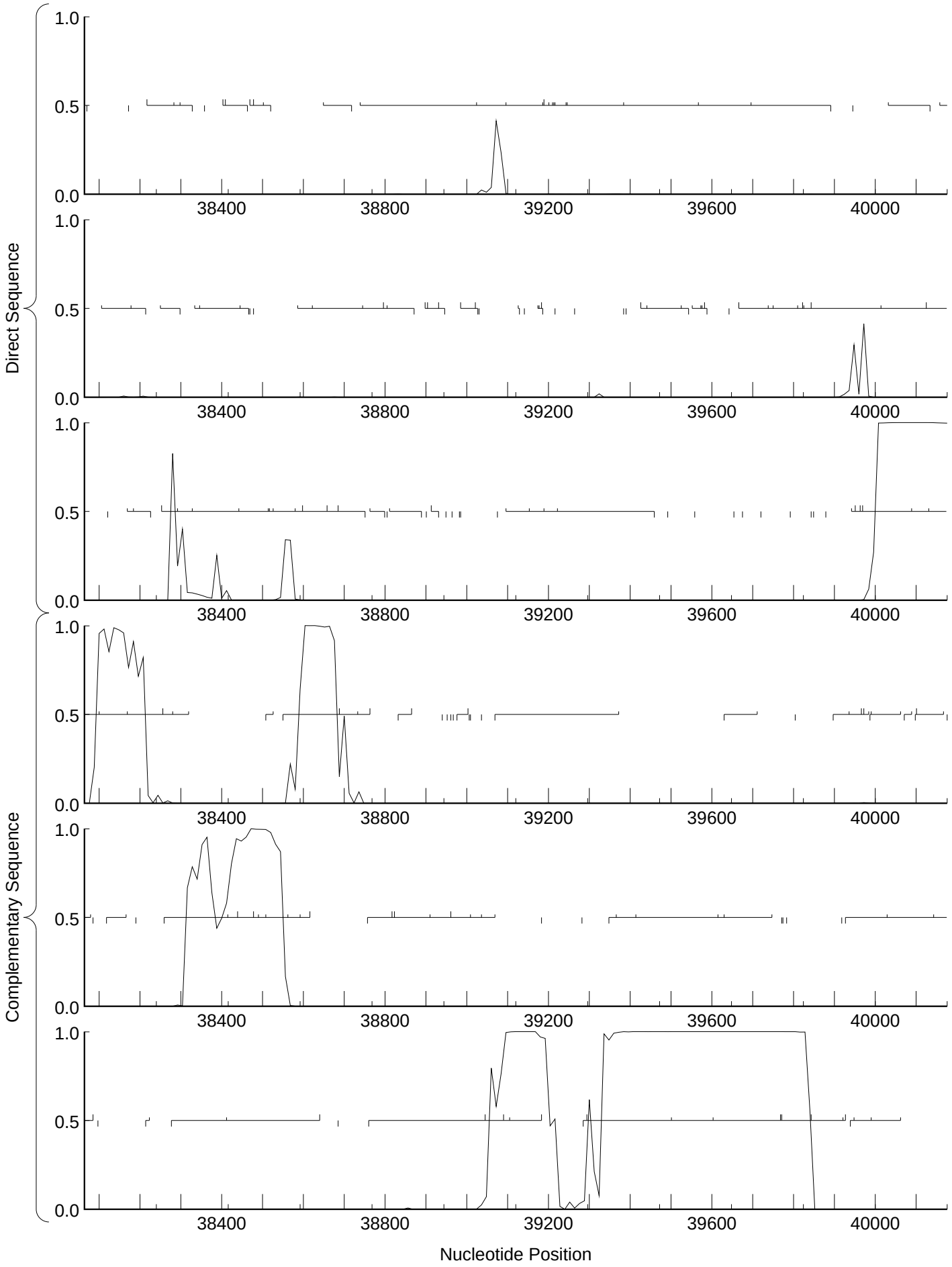
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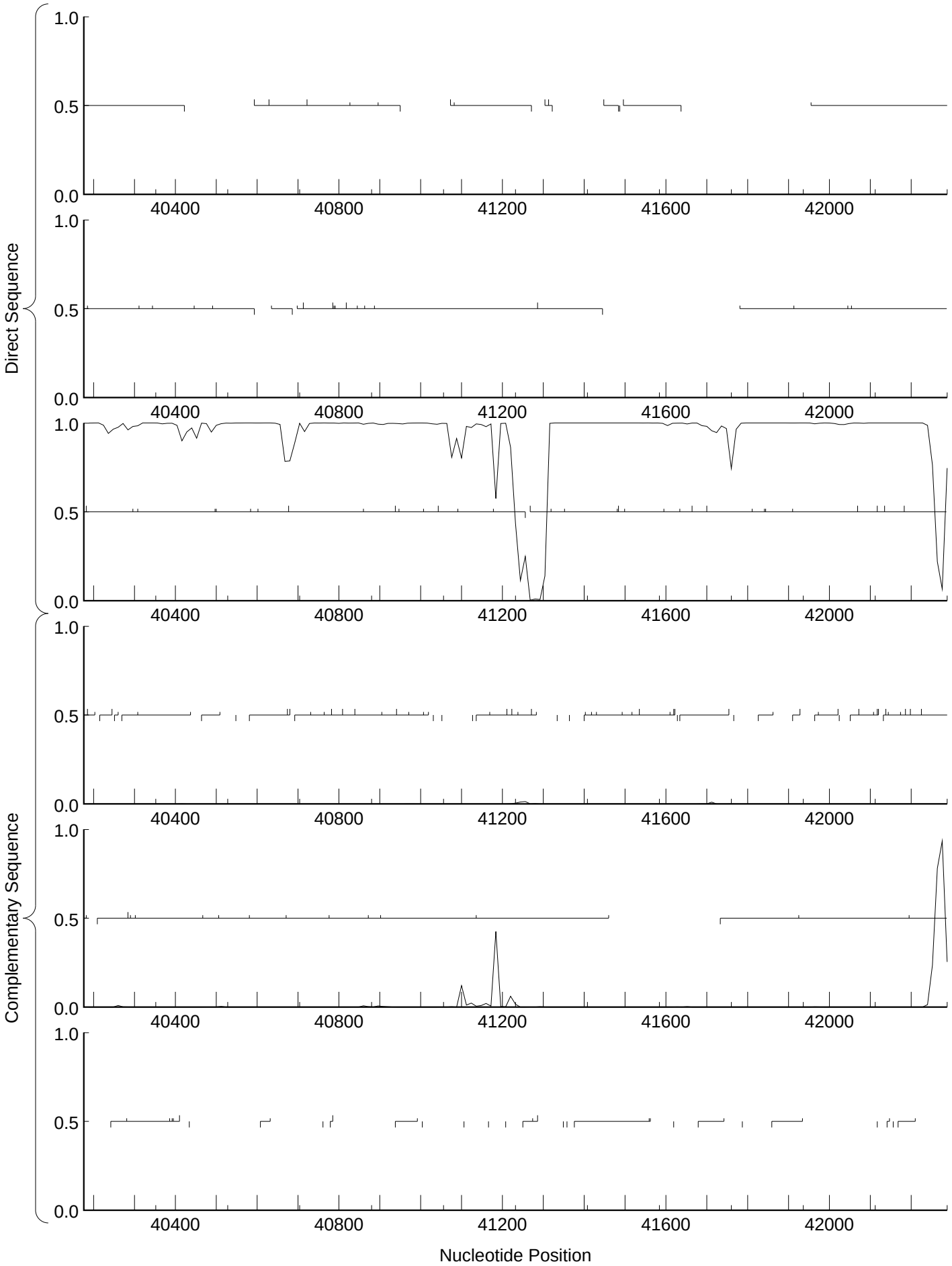

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Complementary Sequence
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Nucleotide Position

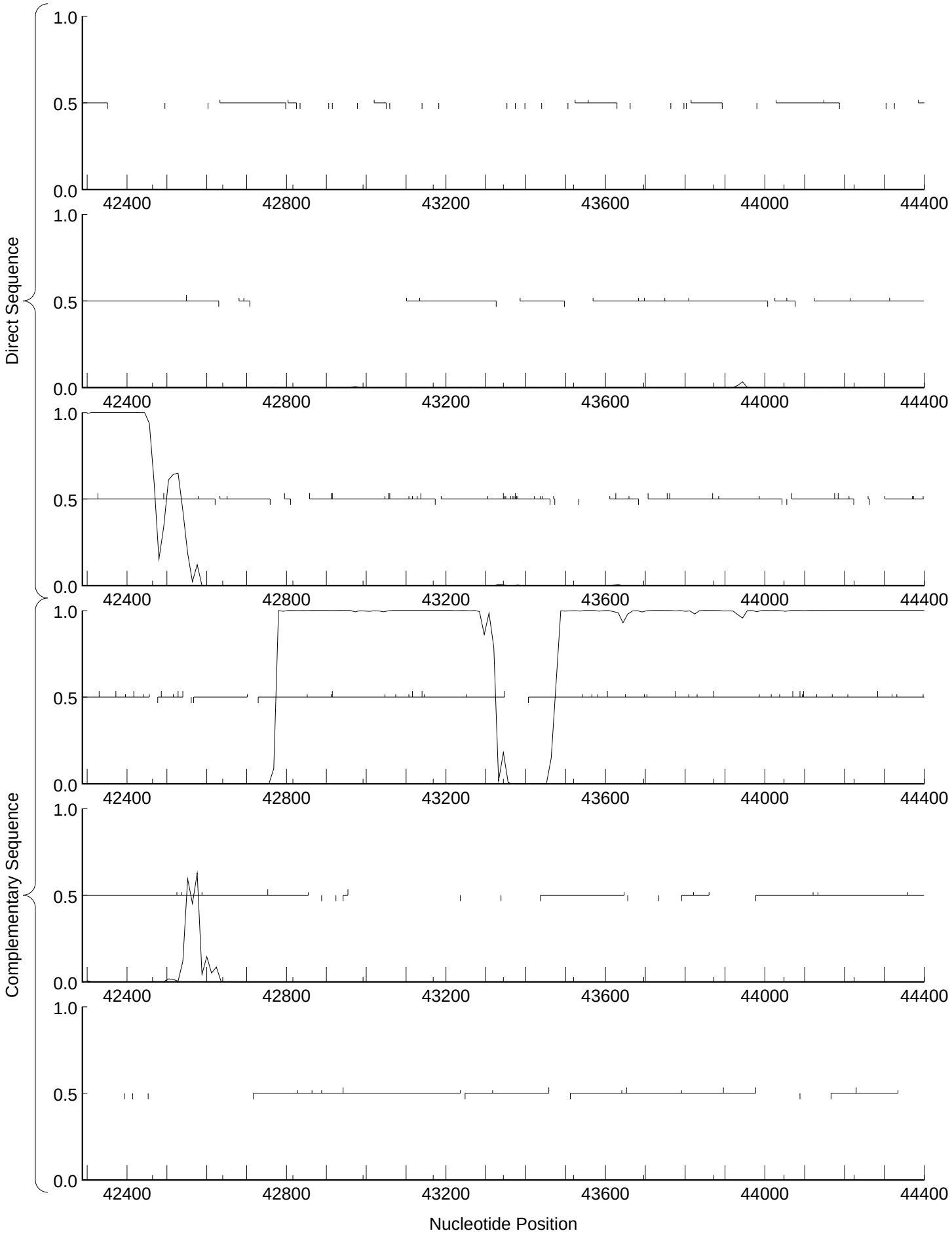
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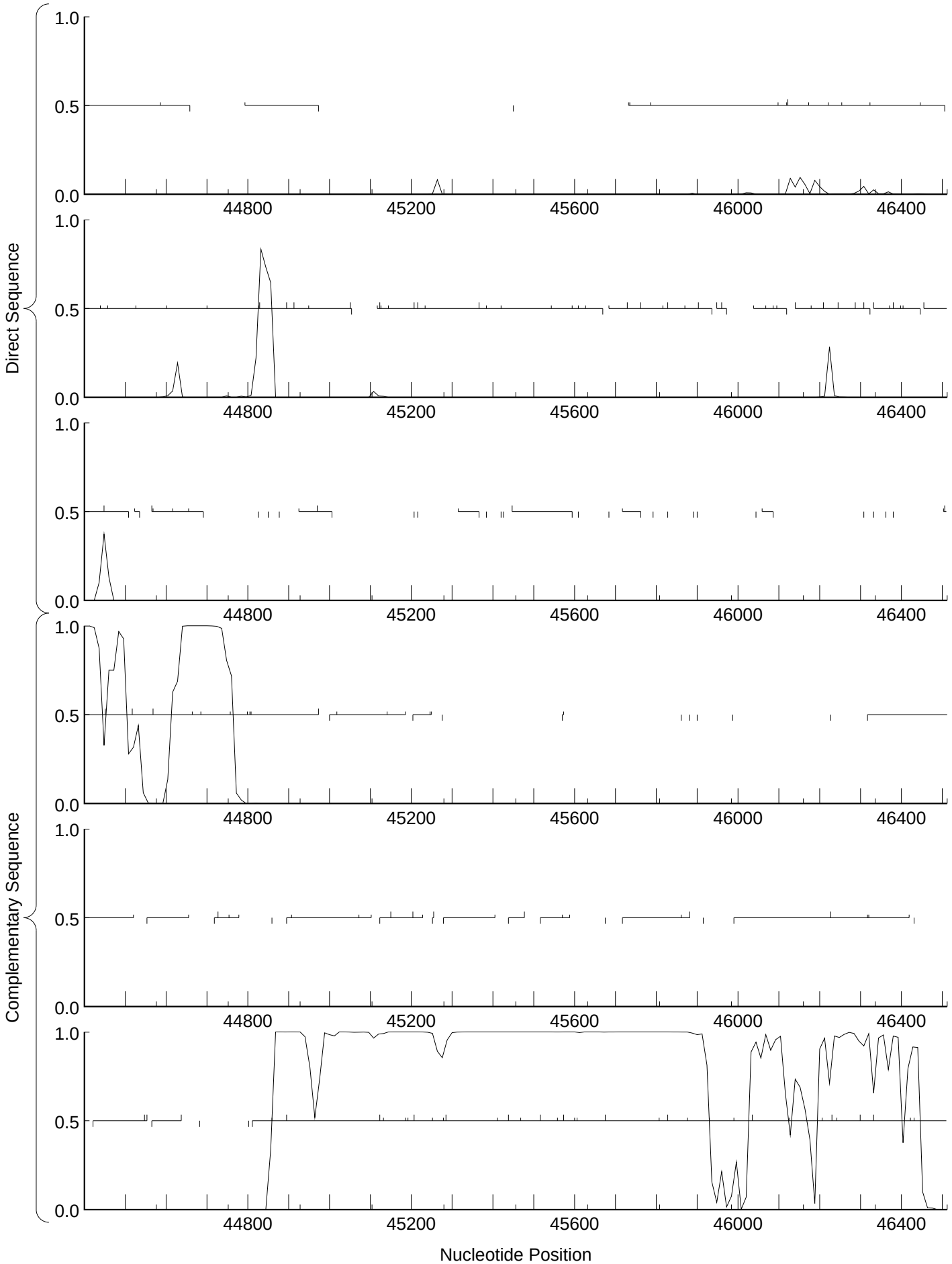



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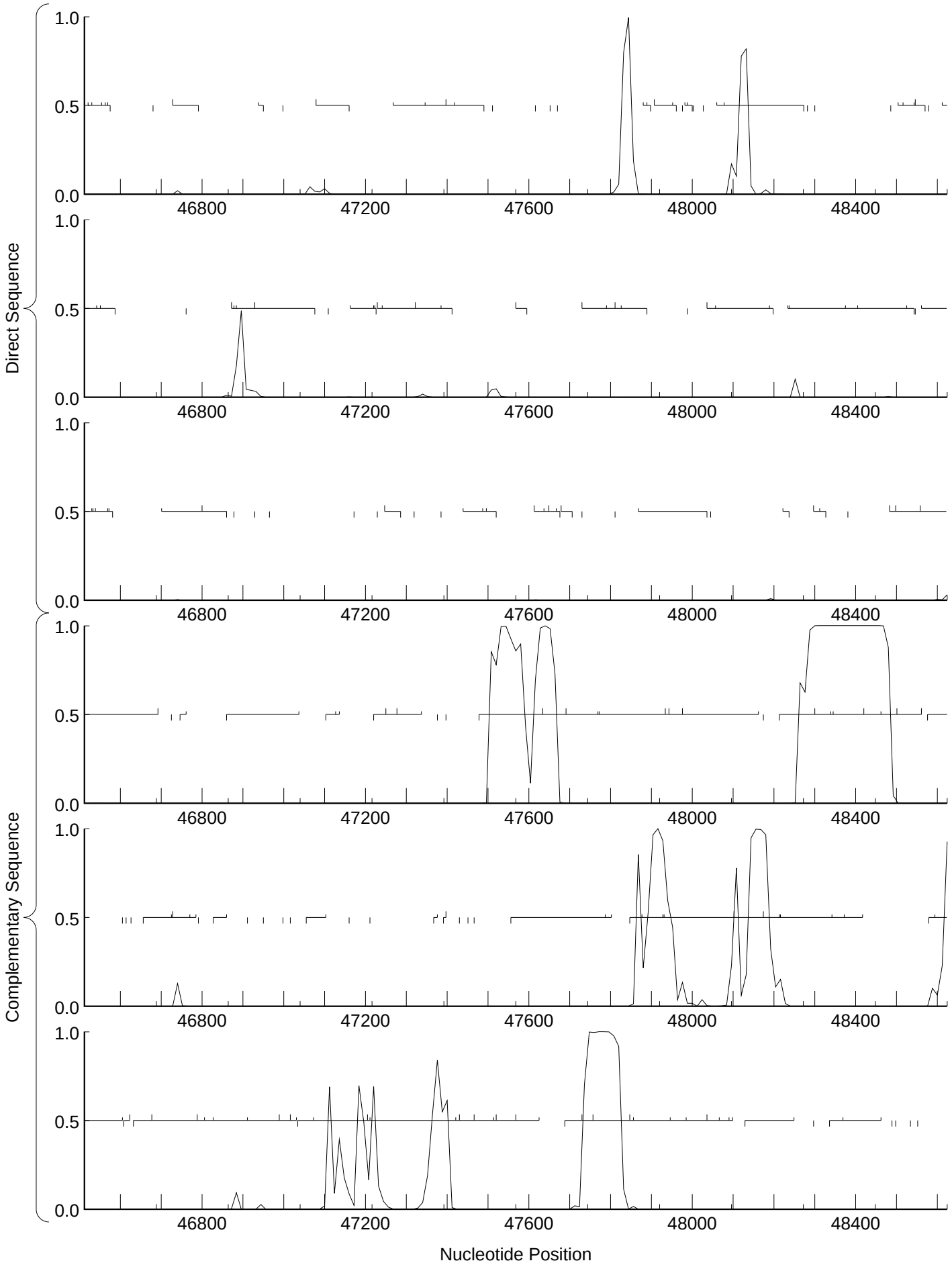
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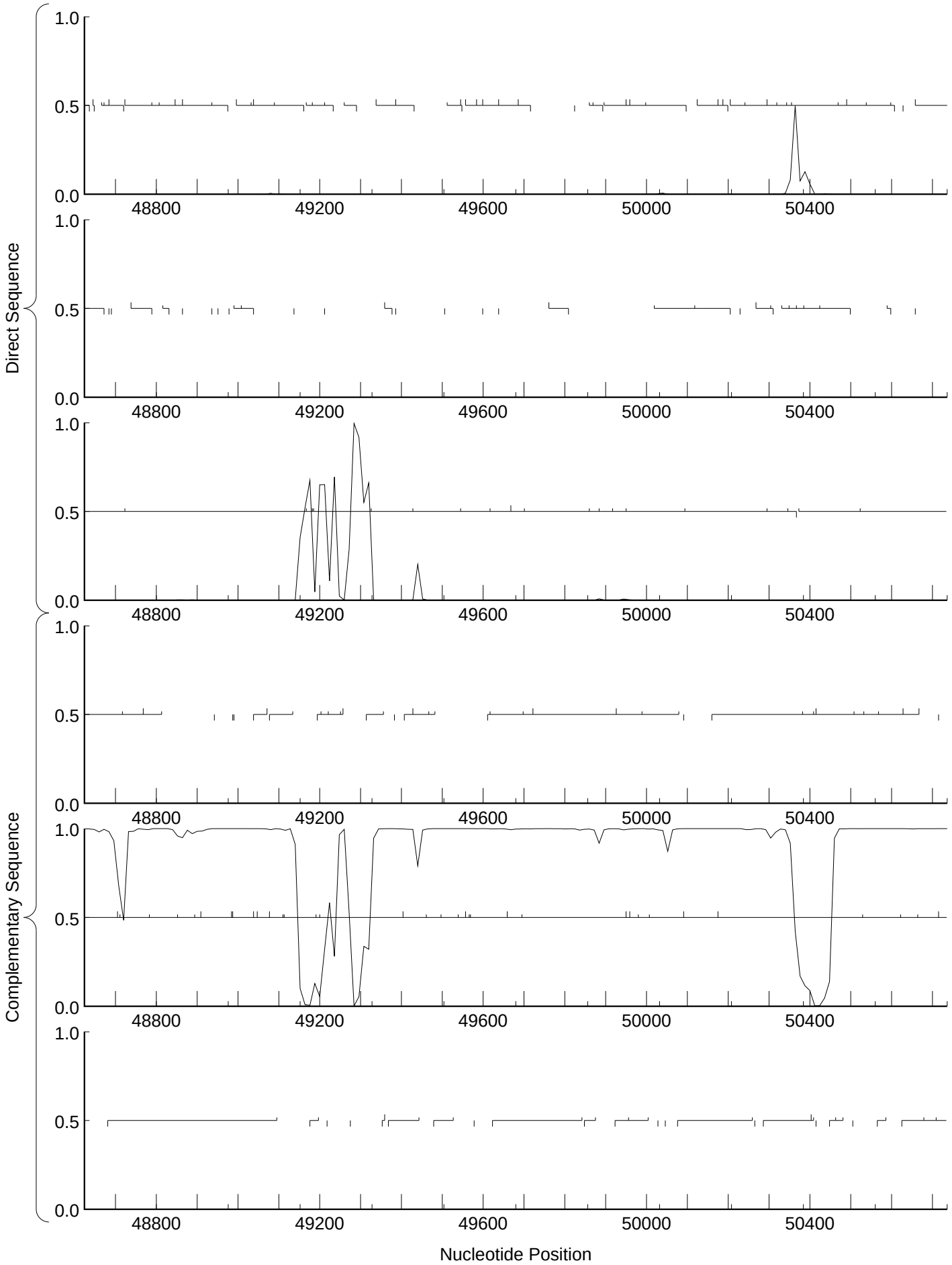

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Nucleotide Position

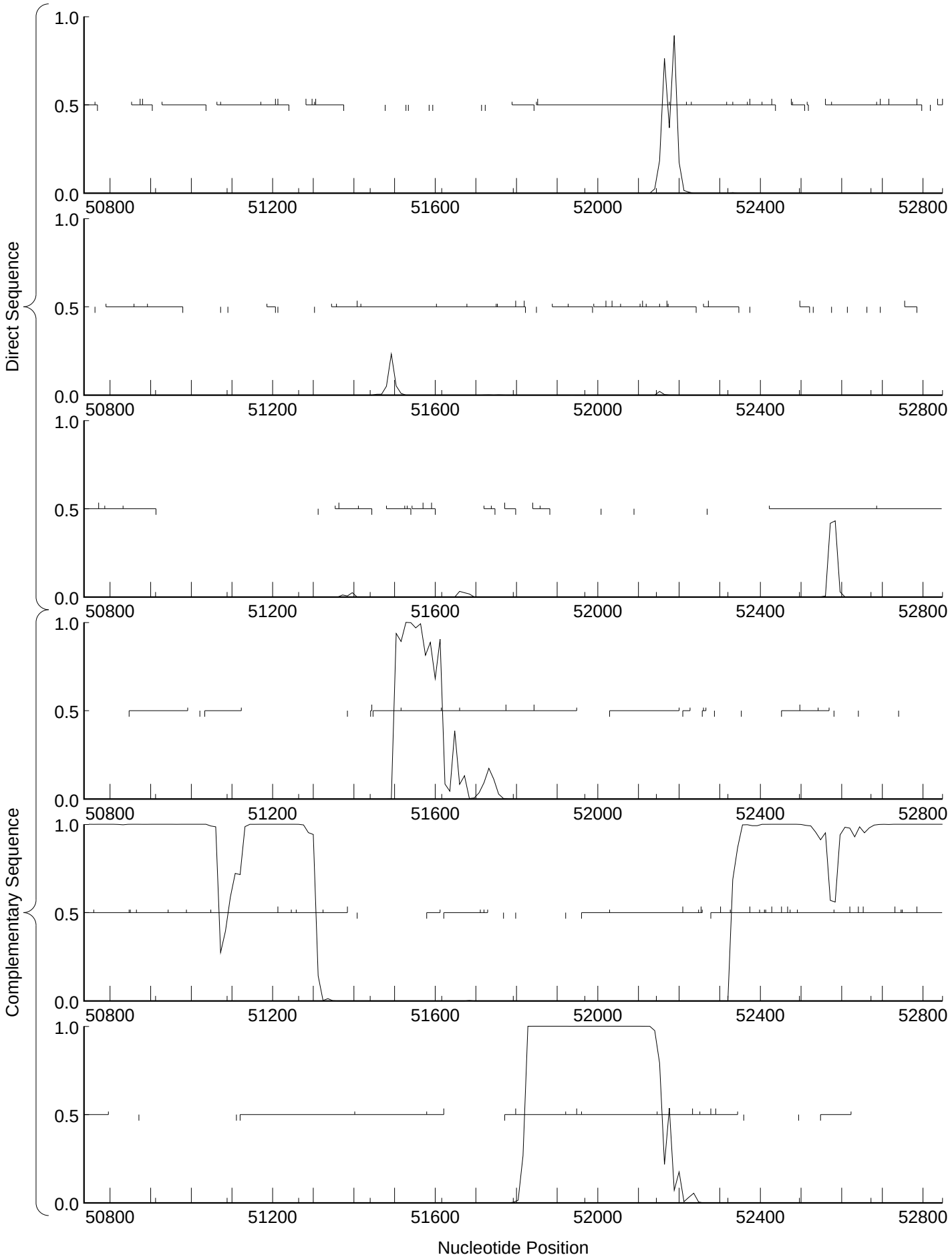

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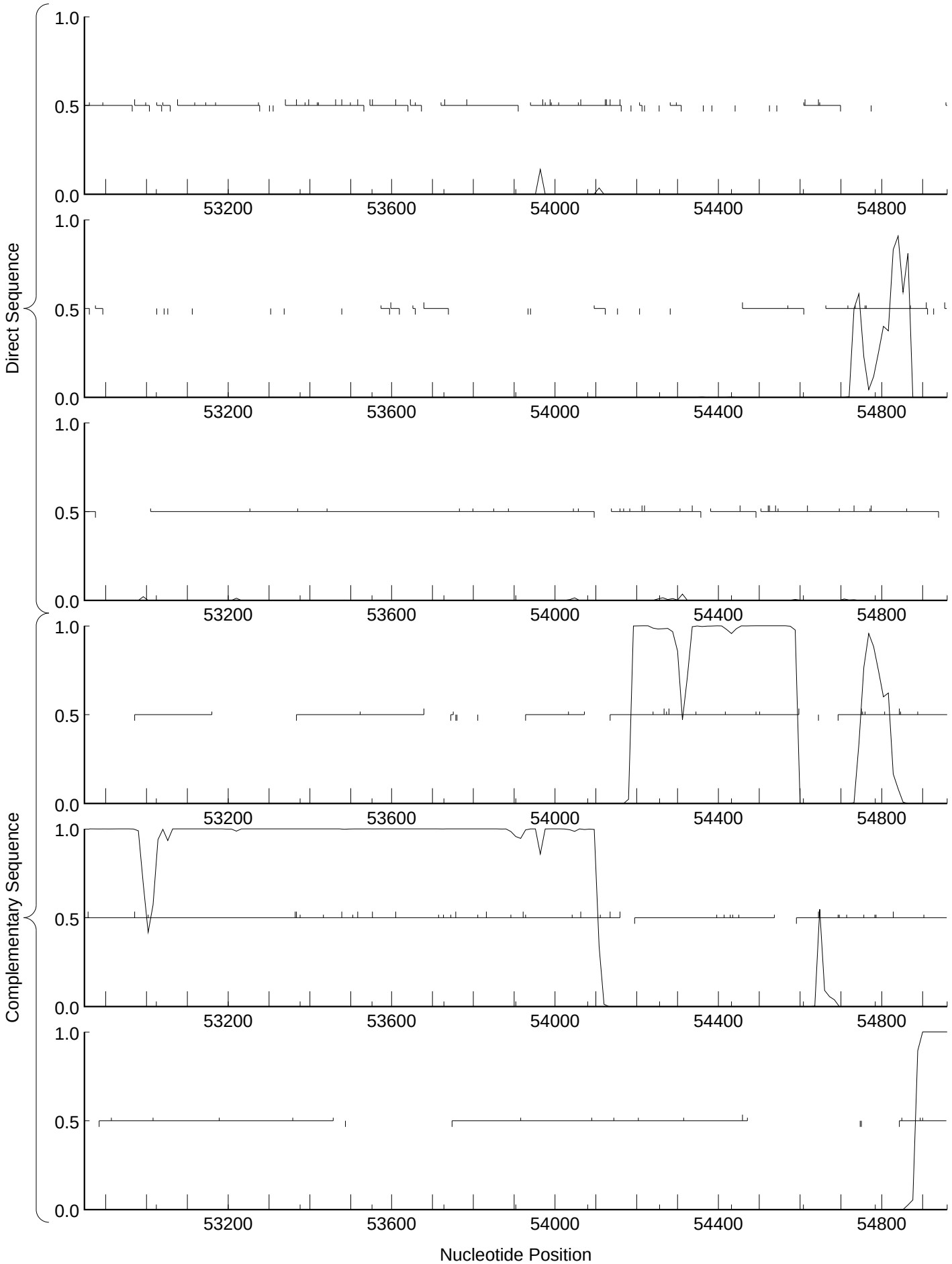
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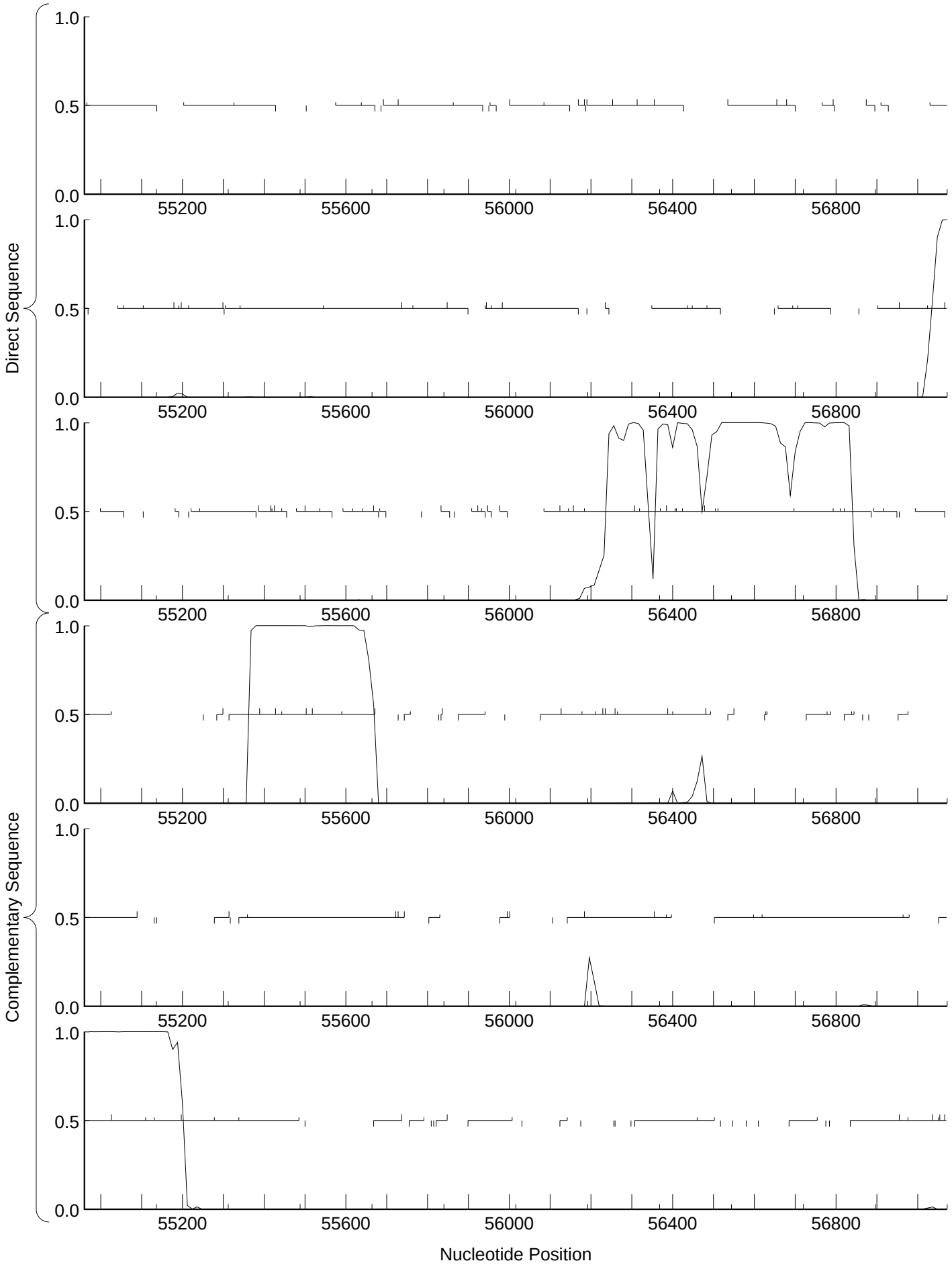
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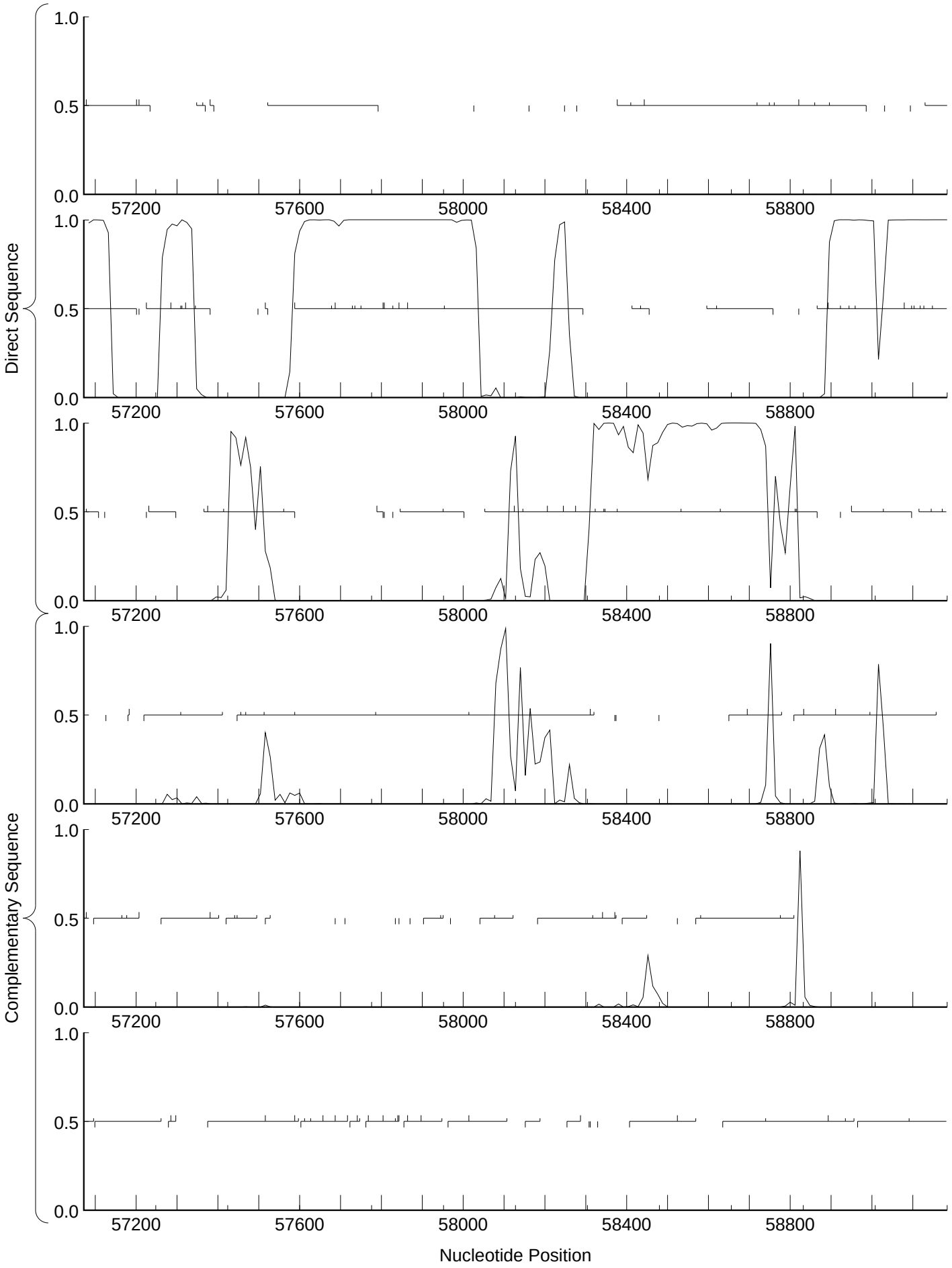

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Nucleotide Position

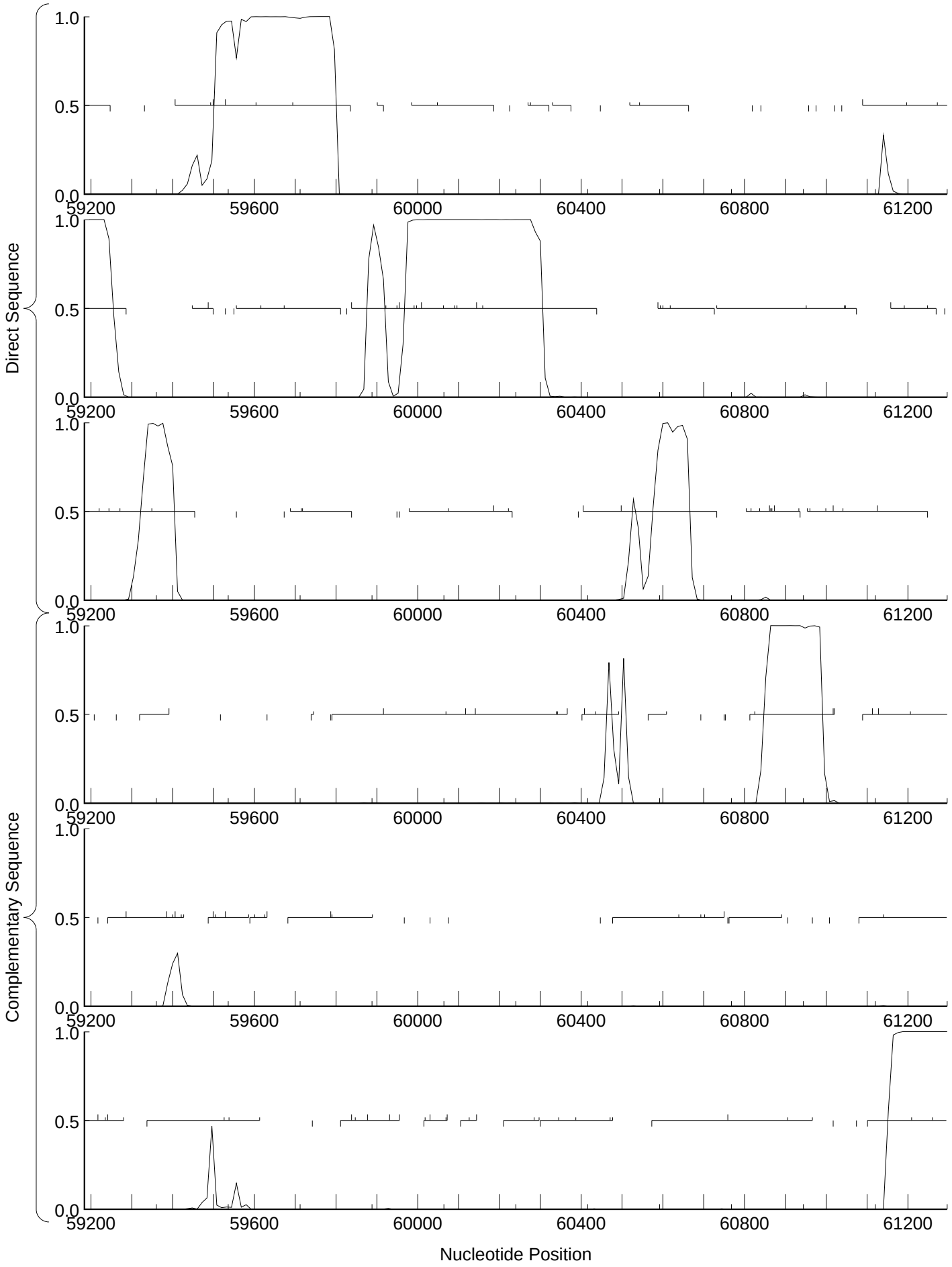
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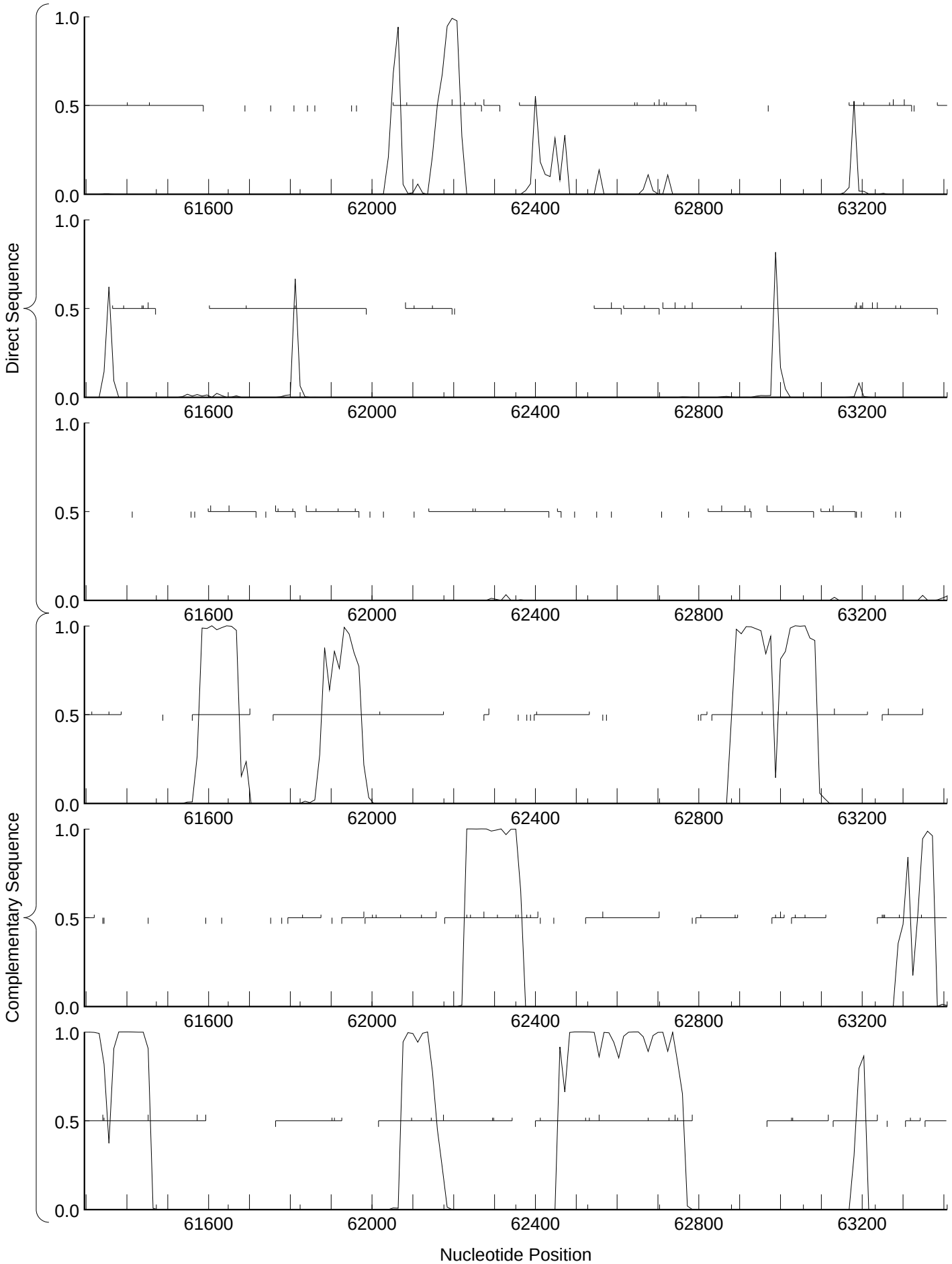


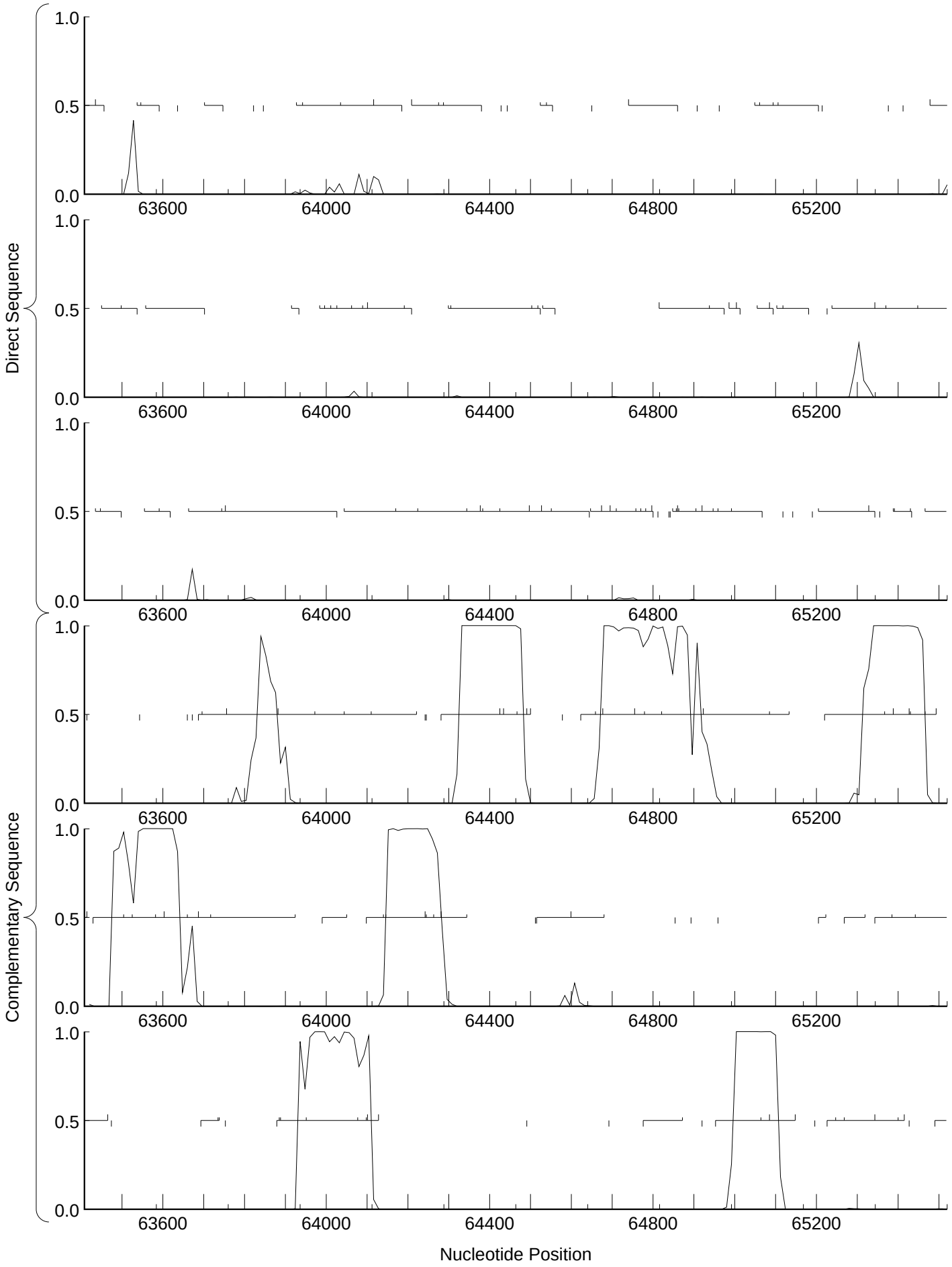
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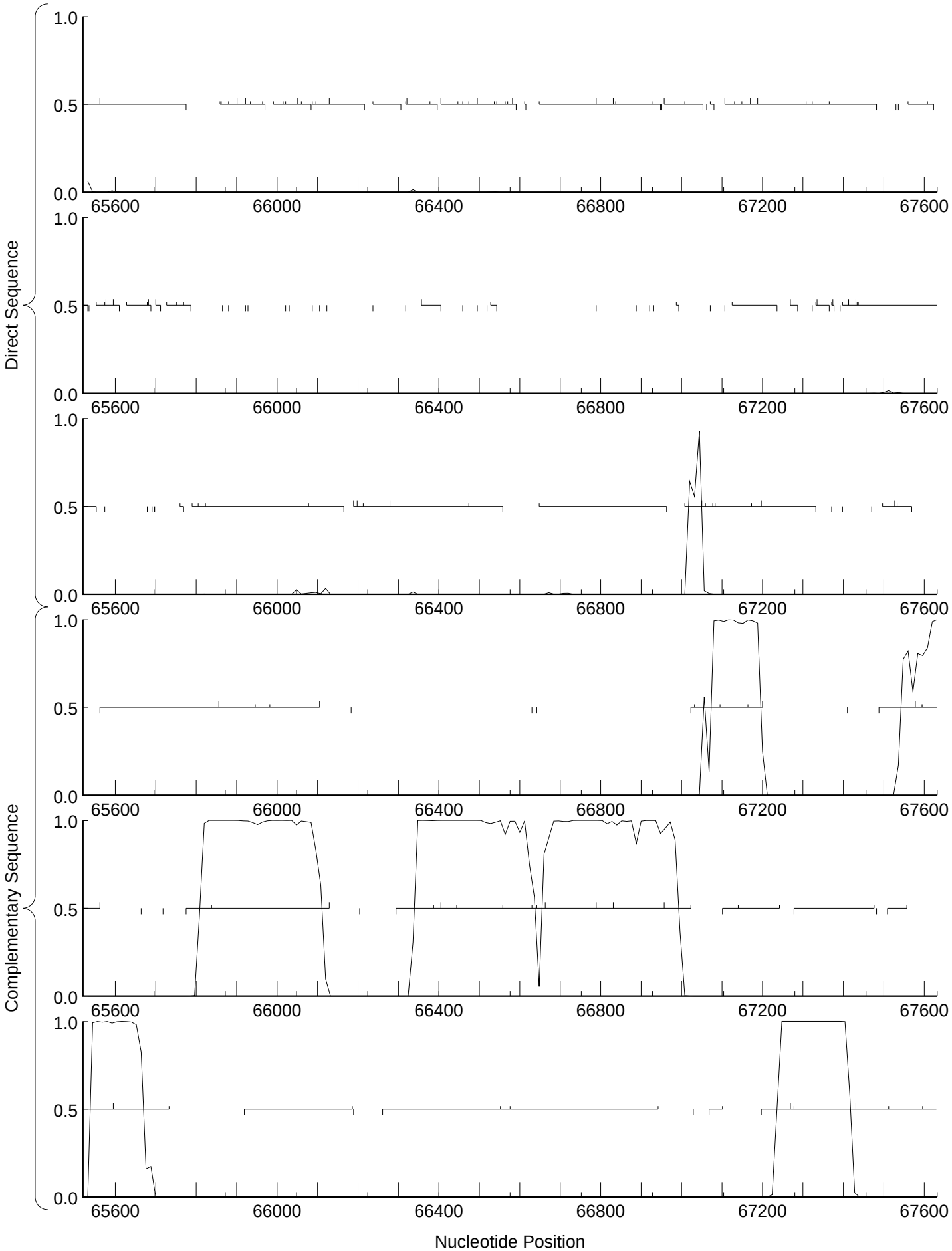
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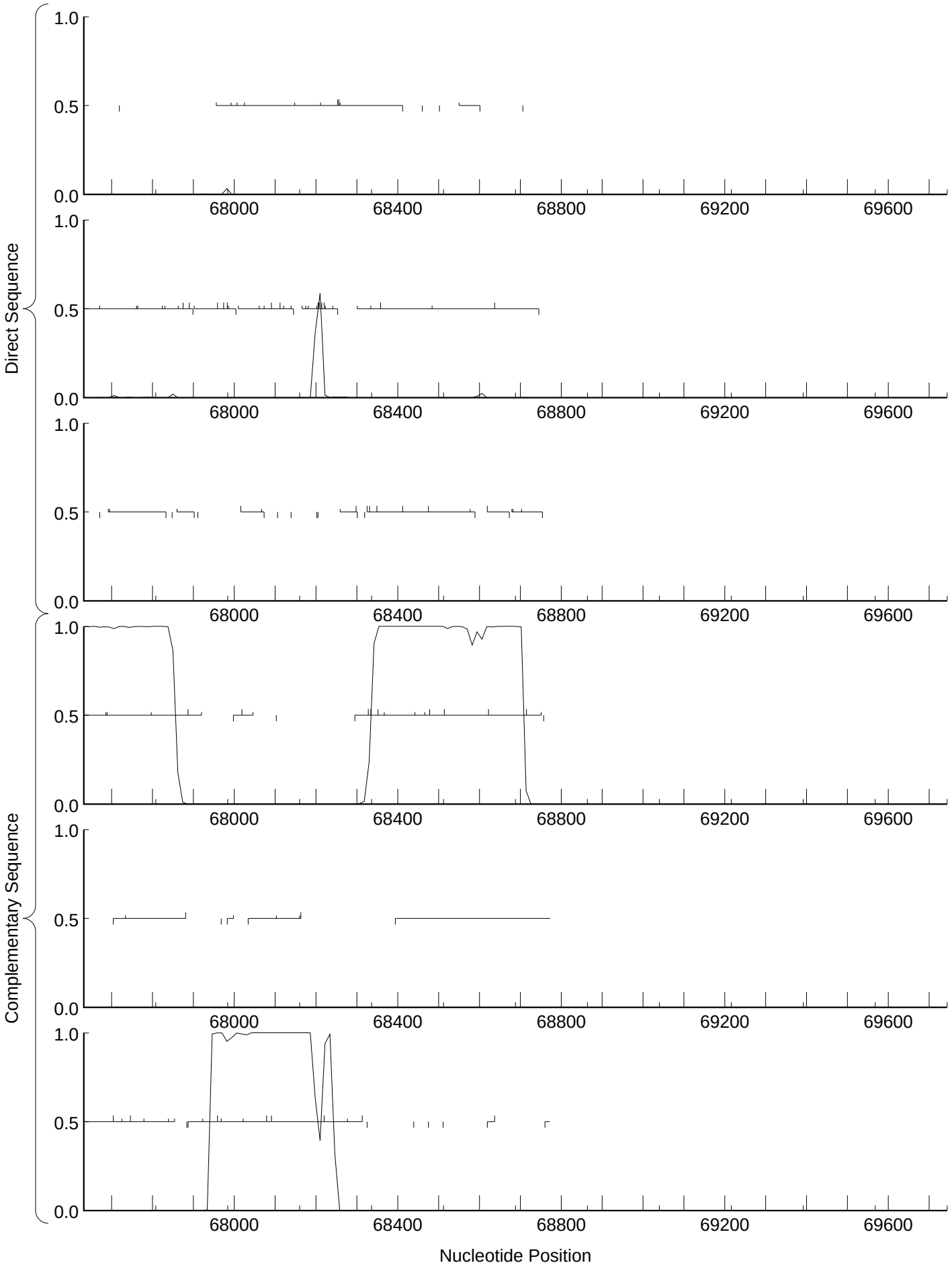



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Direct Sequence
Complementary Sequence
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0.0
63600
64000
64400
64800
65200
Nucleotide Position

bacterium phage Kwksand96 complete sequence, 68822 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 33/34




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
68000
68400
68800
69200
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