GeneMark

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

PROGRAM INFORMATION

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Sequence         :  Arthrobacter phage Lymara complete sequence, 68762 bp, circularly permuted, Cluster AR
Analysis Date    :  3/28/24 at 22:23:54
Pages            :  34
Sequence Length  :  68762 bp
GC Content       :   61.49%

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

PS-Version       :  1.2

GeneMark Options :  PostScript graph,
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MATRIX INFORMATION

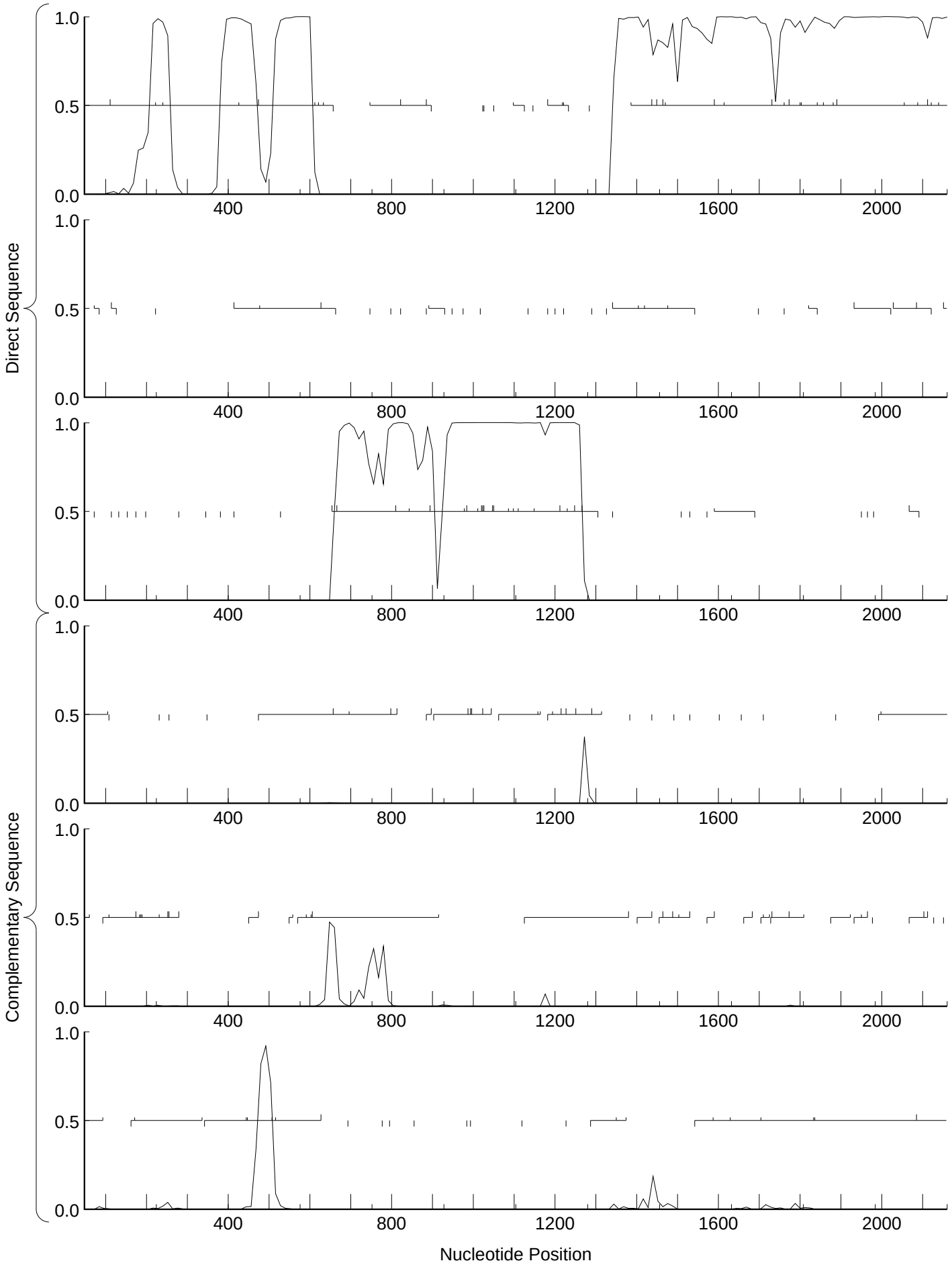
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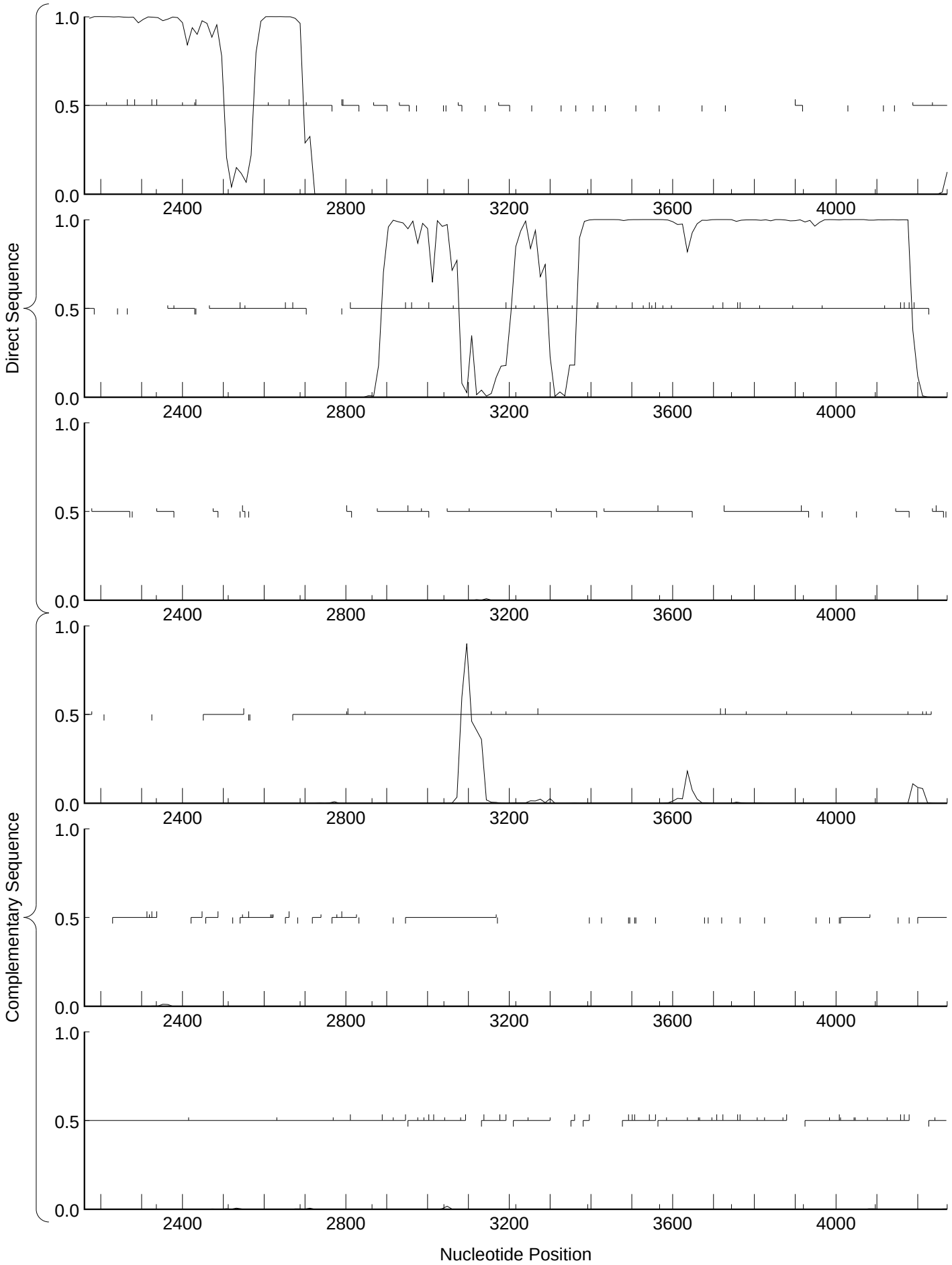

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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

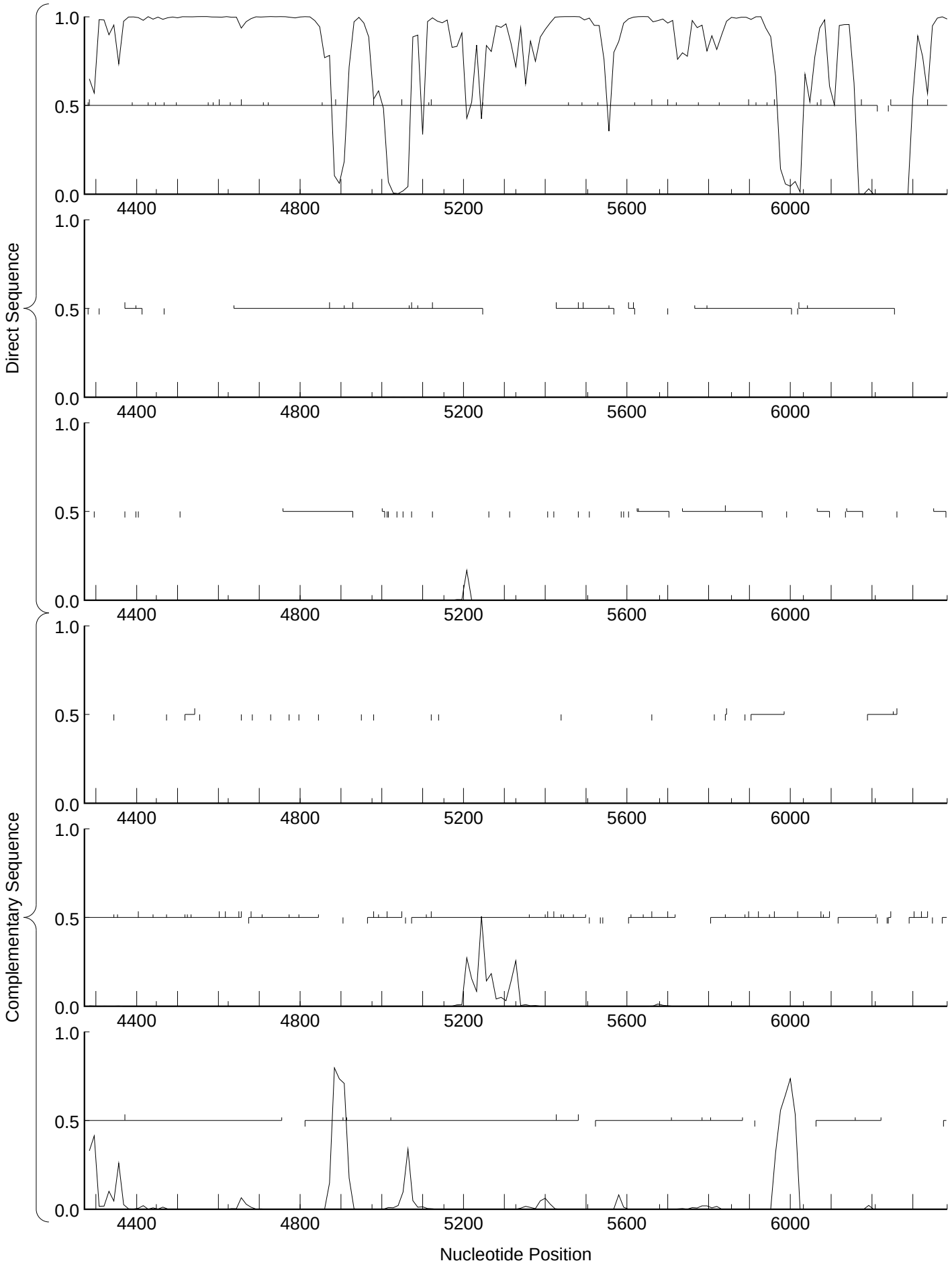
Arthrobacter phage Lymara complete sequence, 68762 bp, circularly permuted, Cluster AR, Order 4, Window 96, Step 12, 2/34



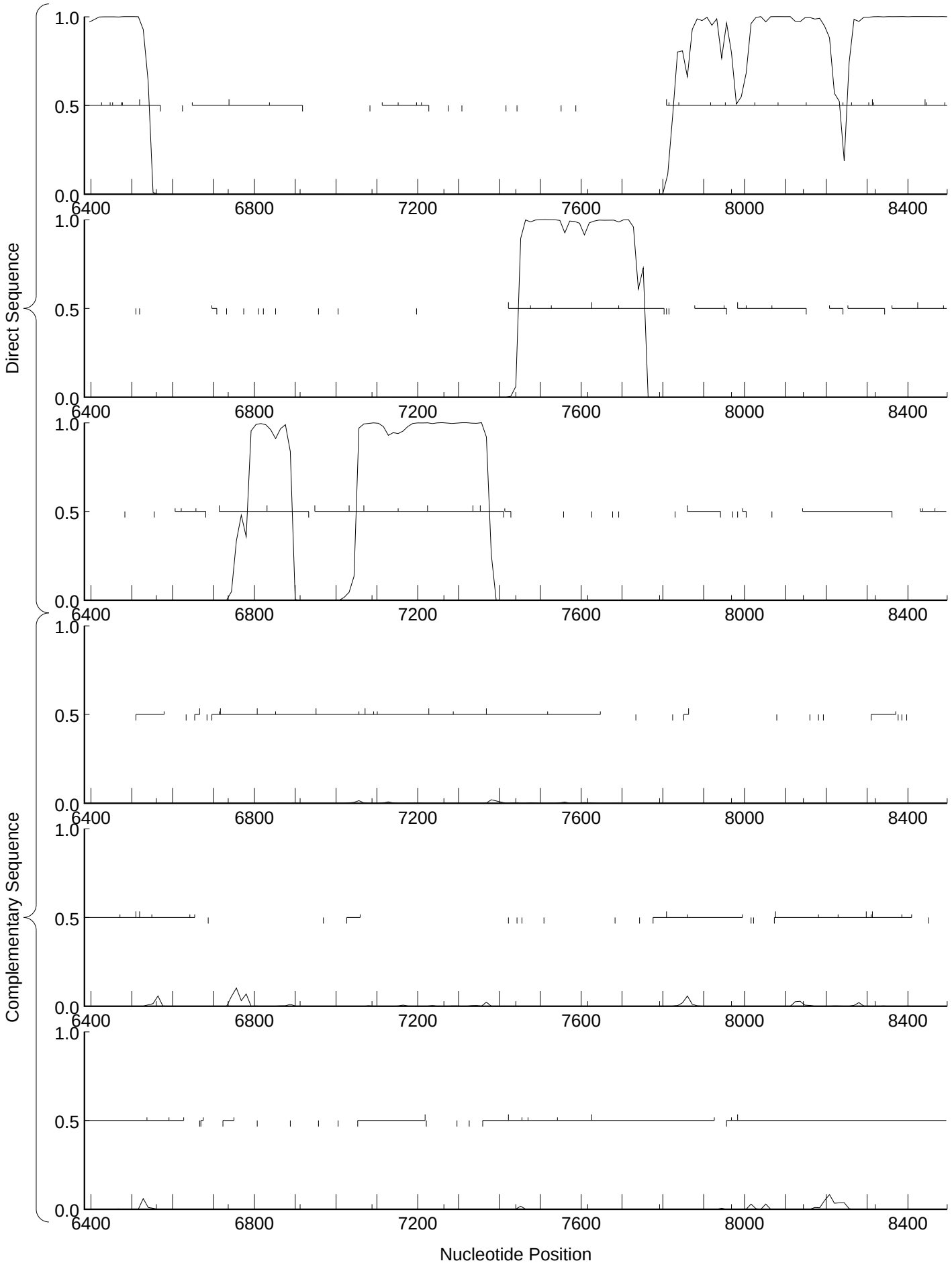
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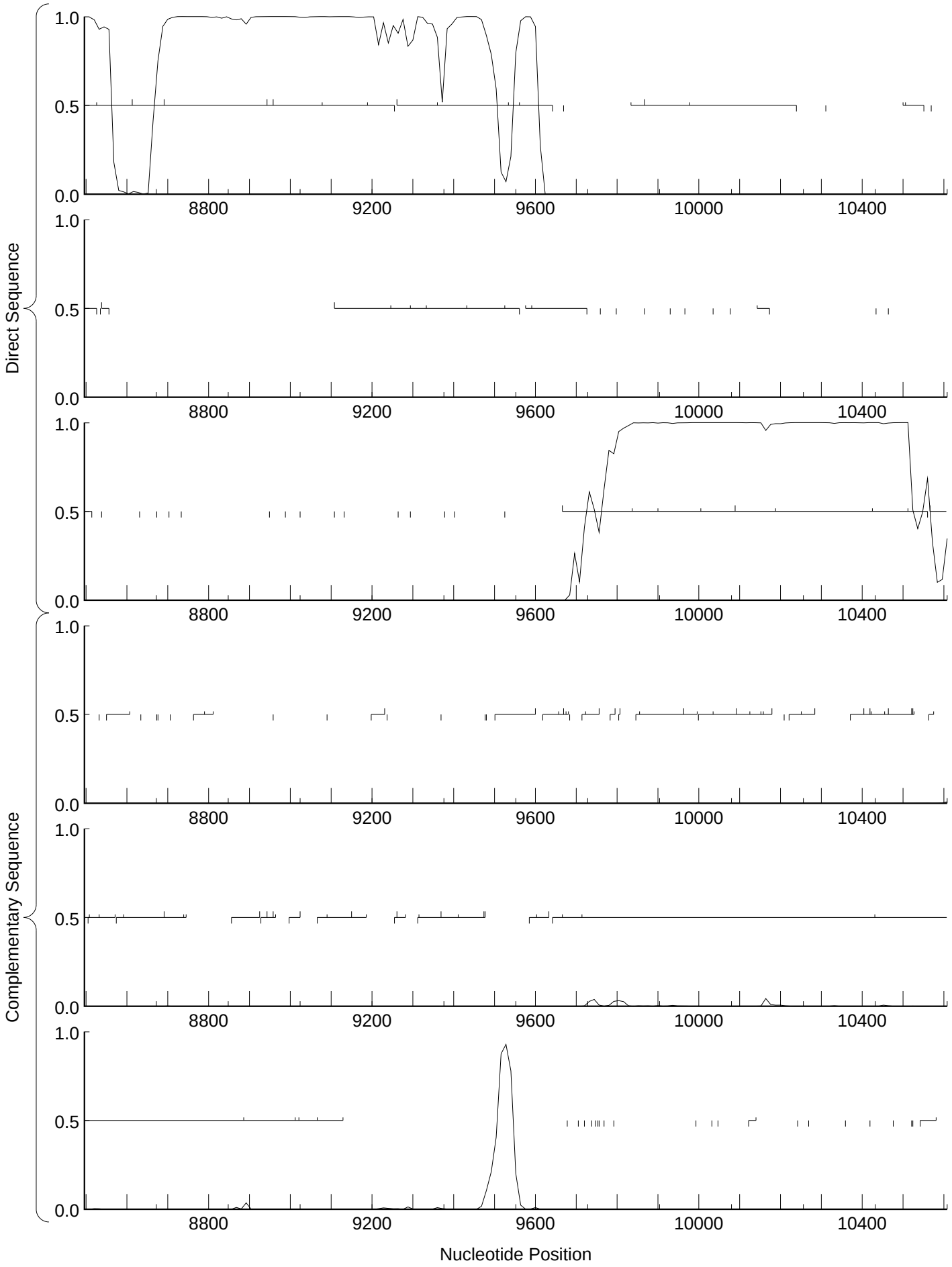
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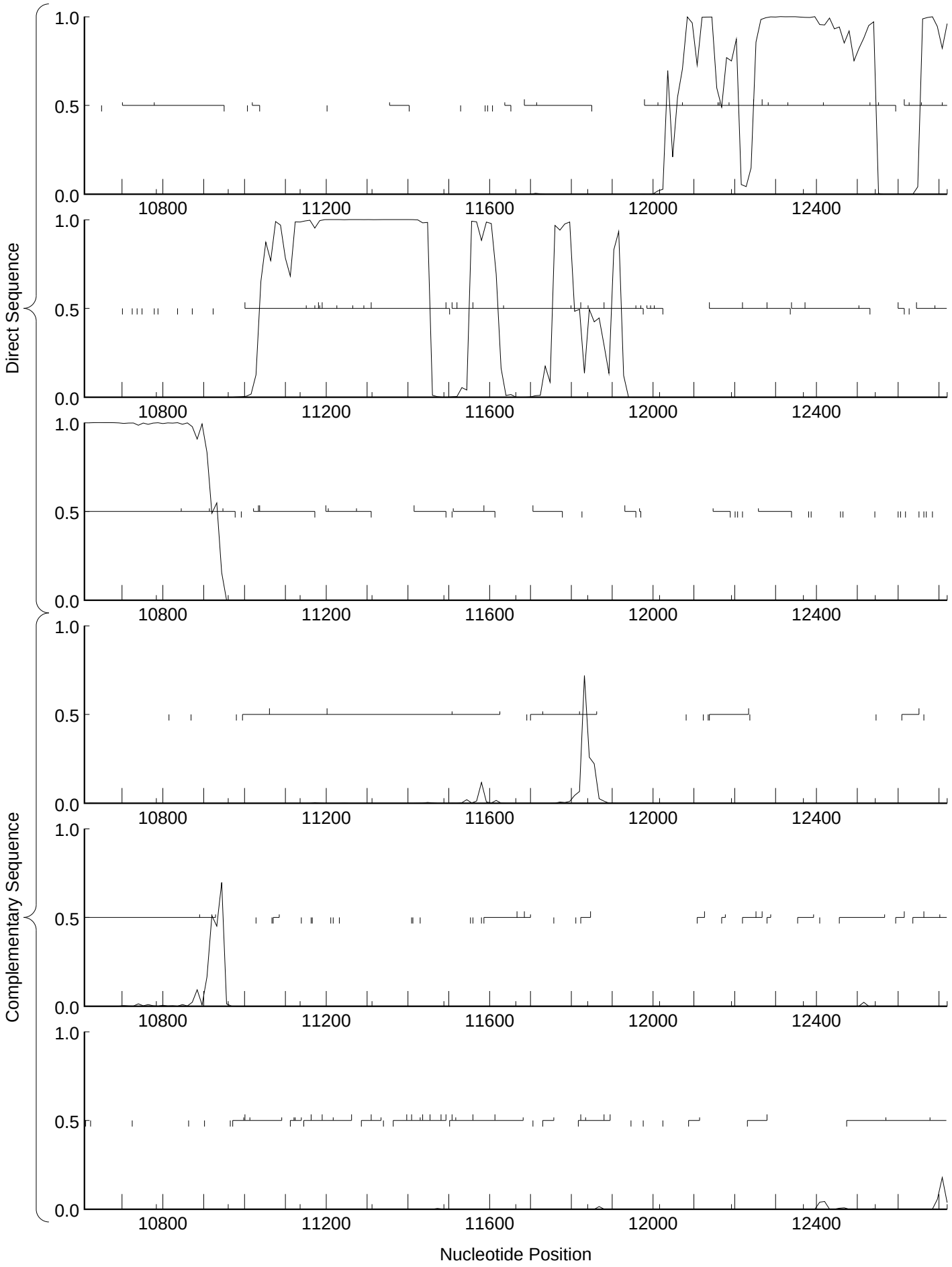
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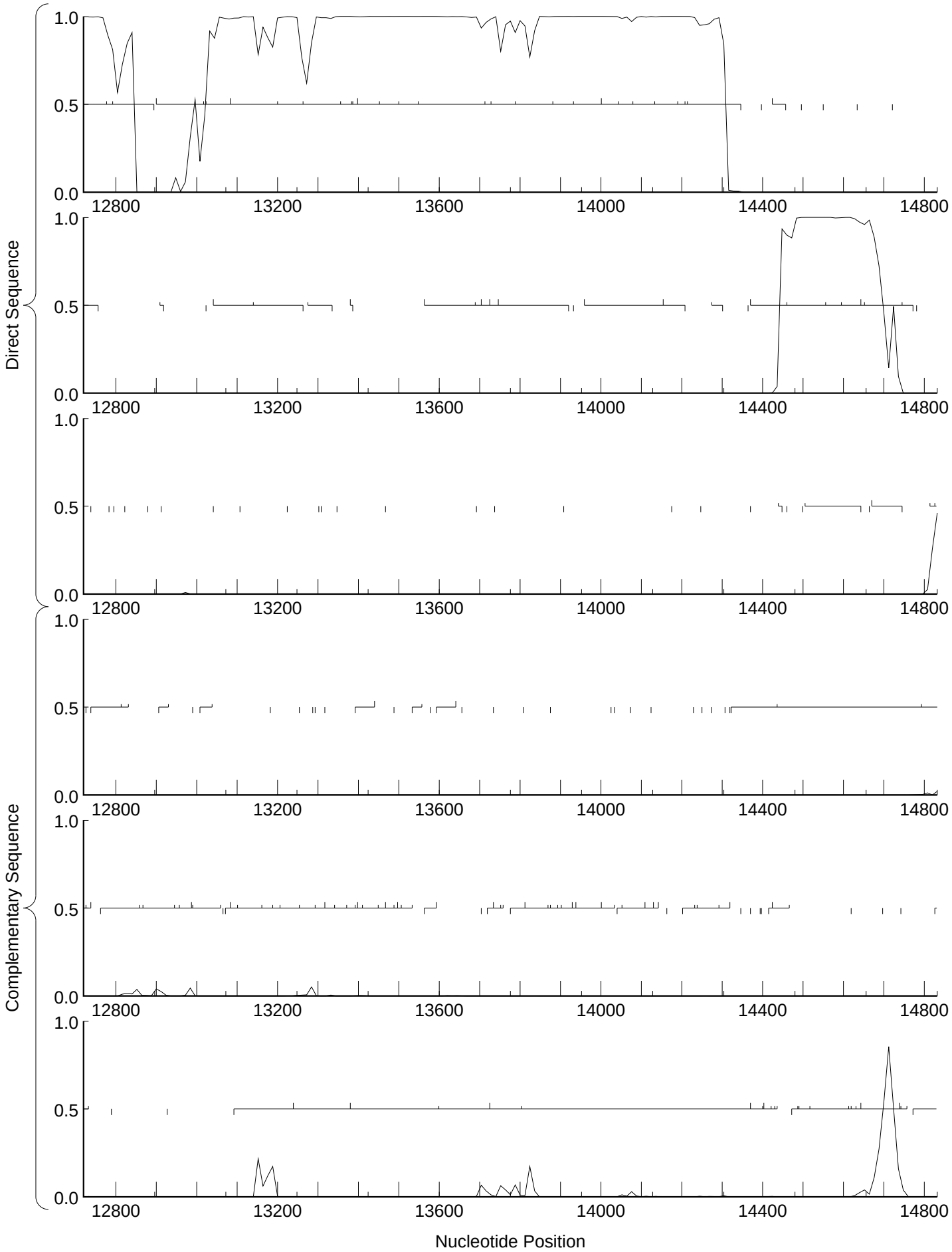
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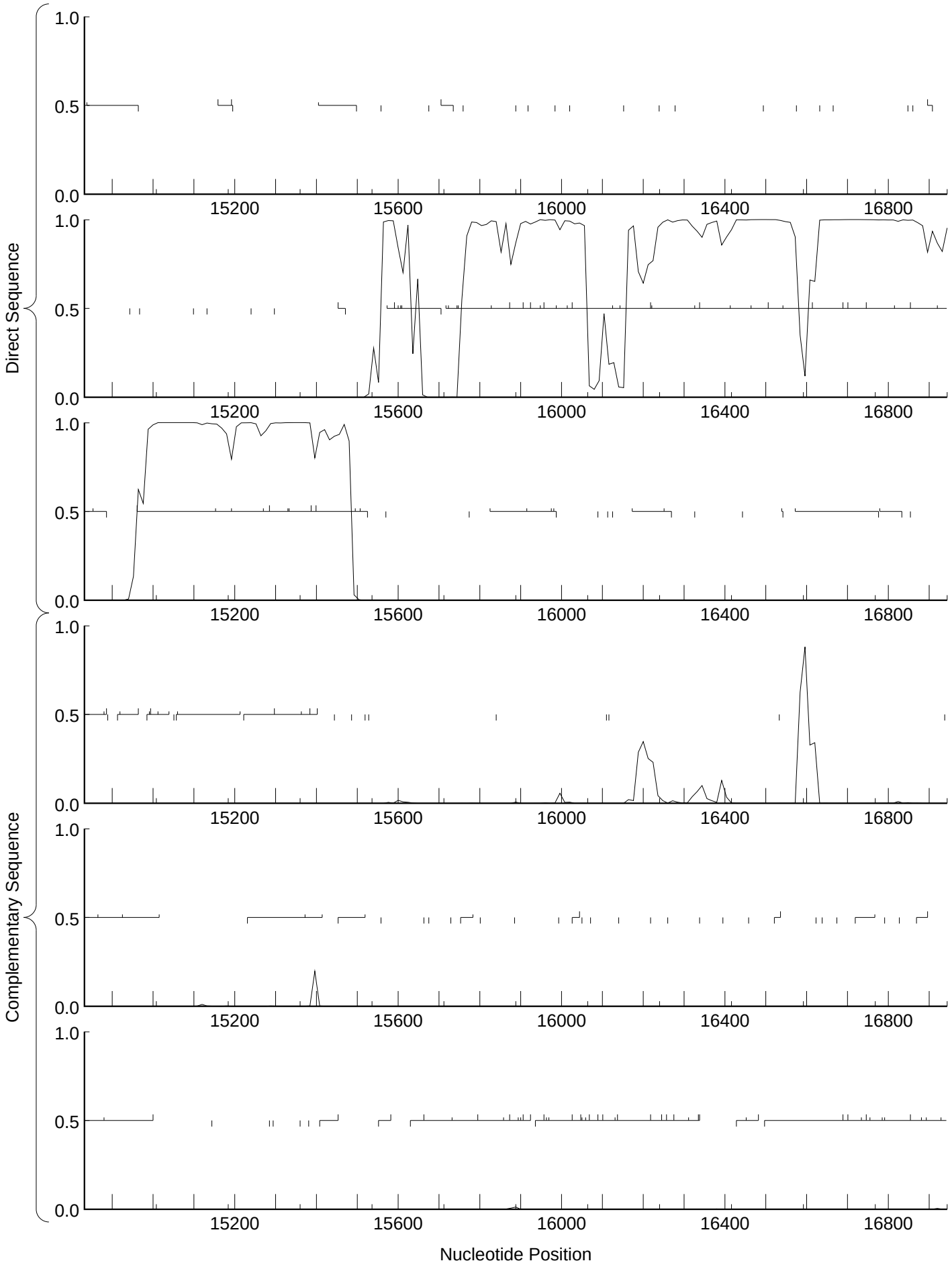
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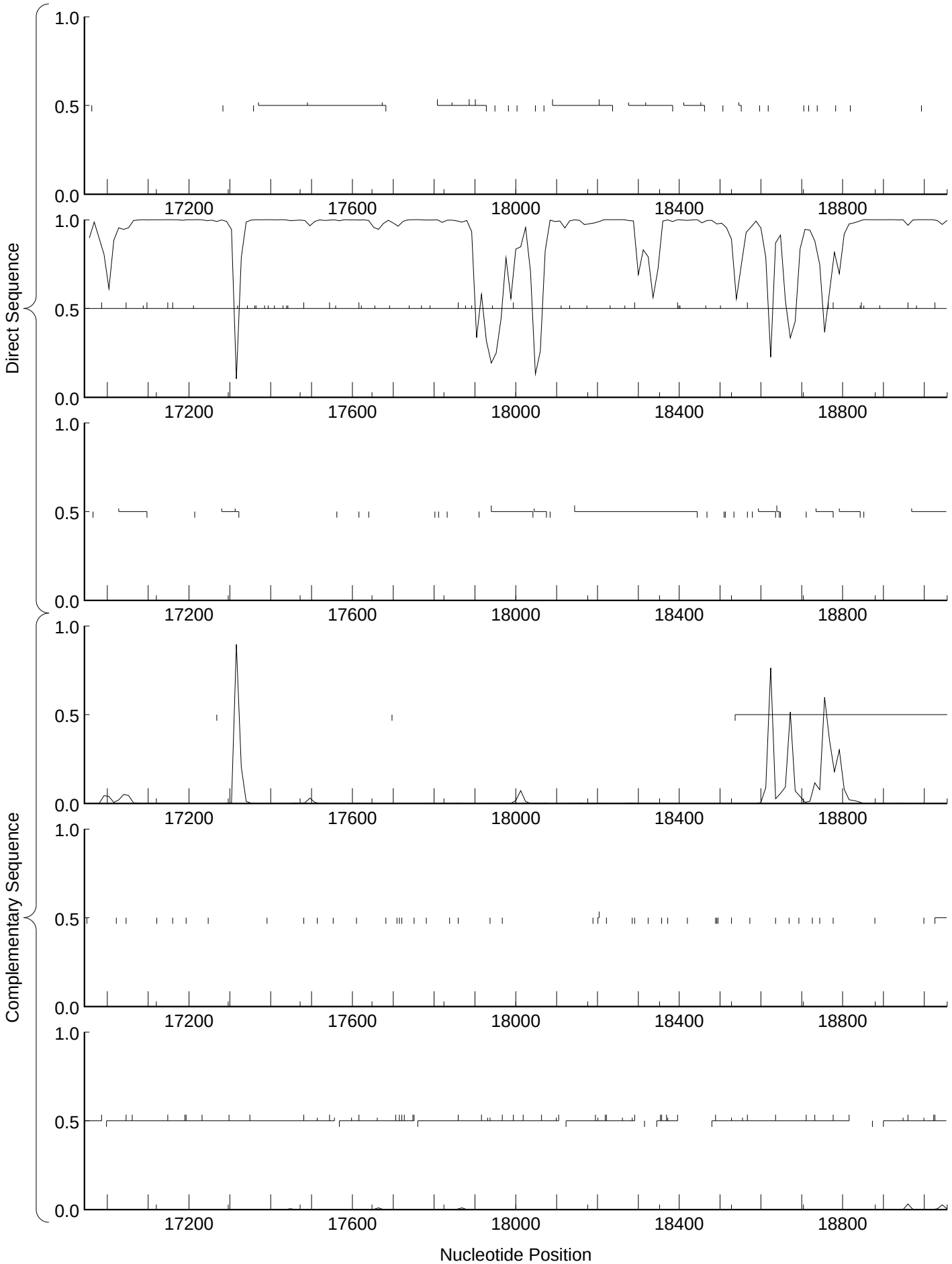
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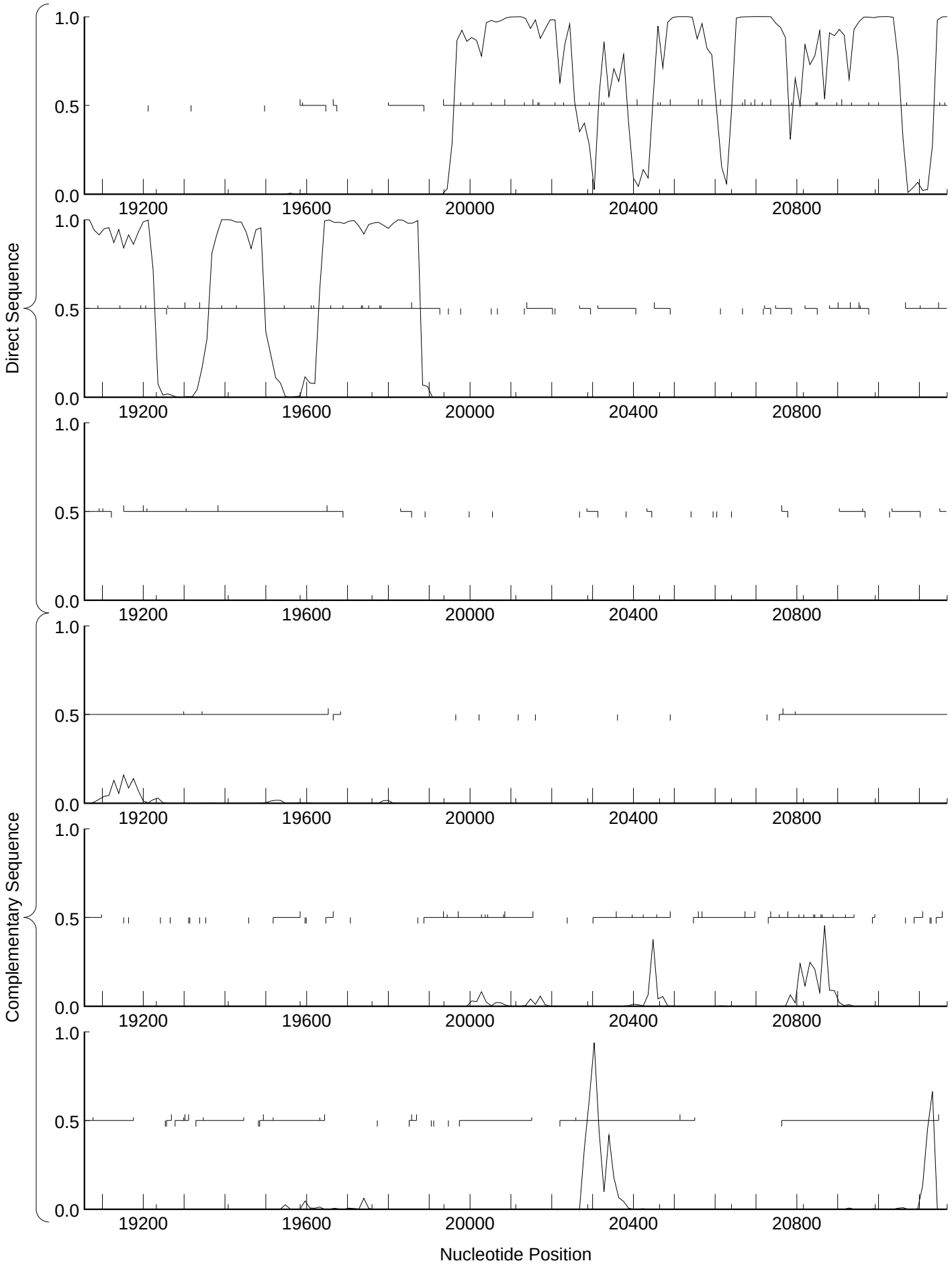
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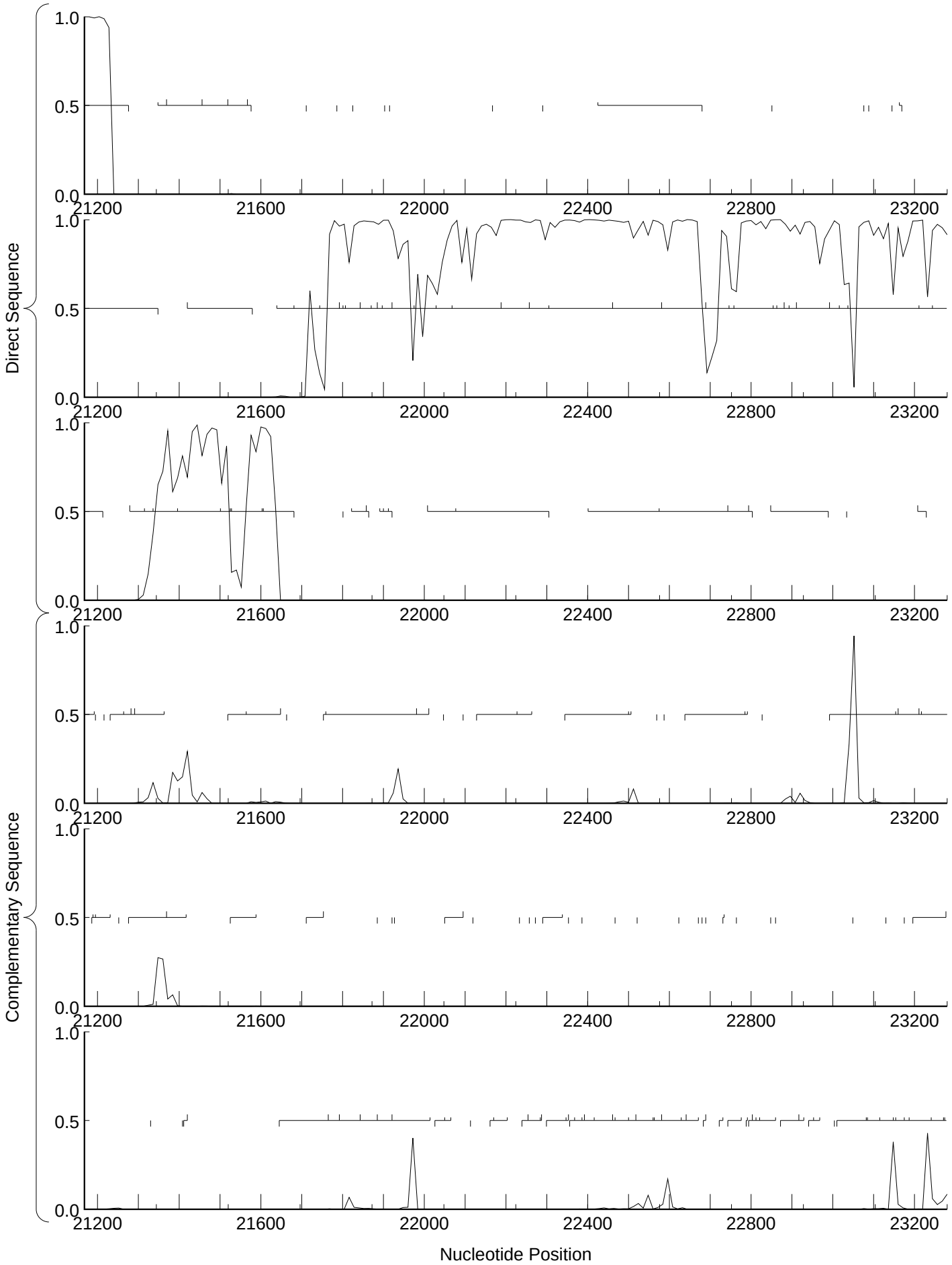
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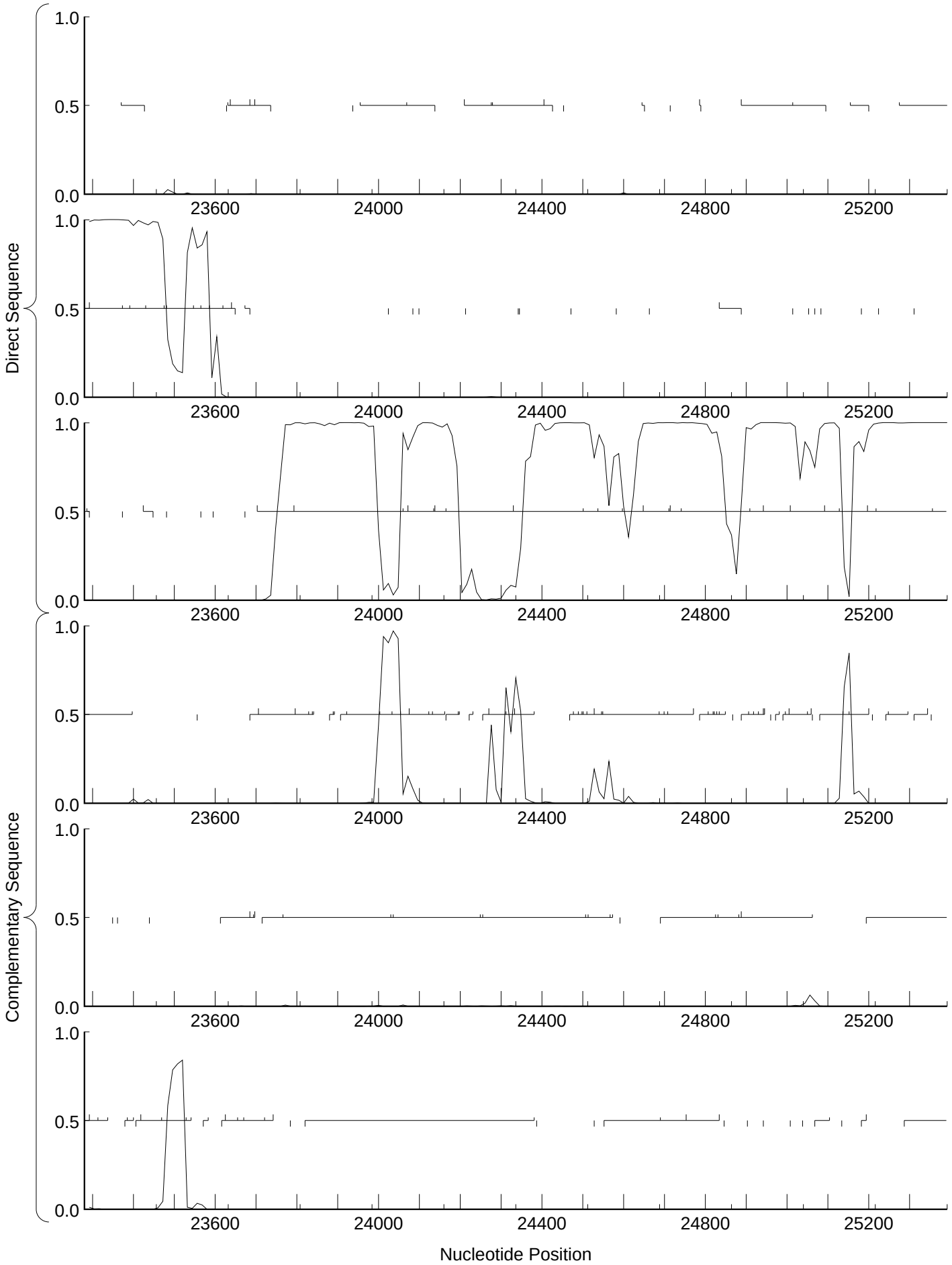
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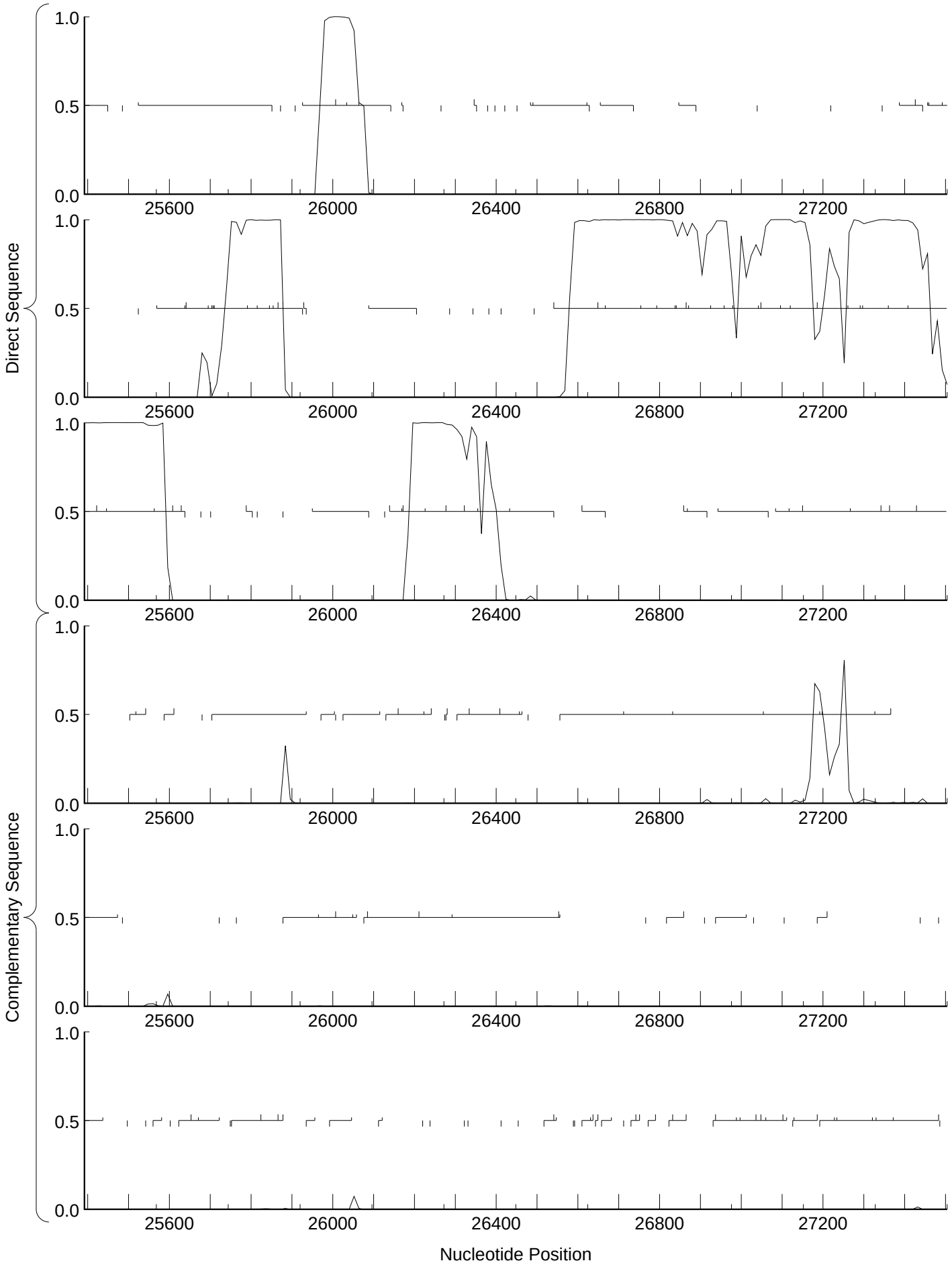
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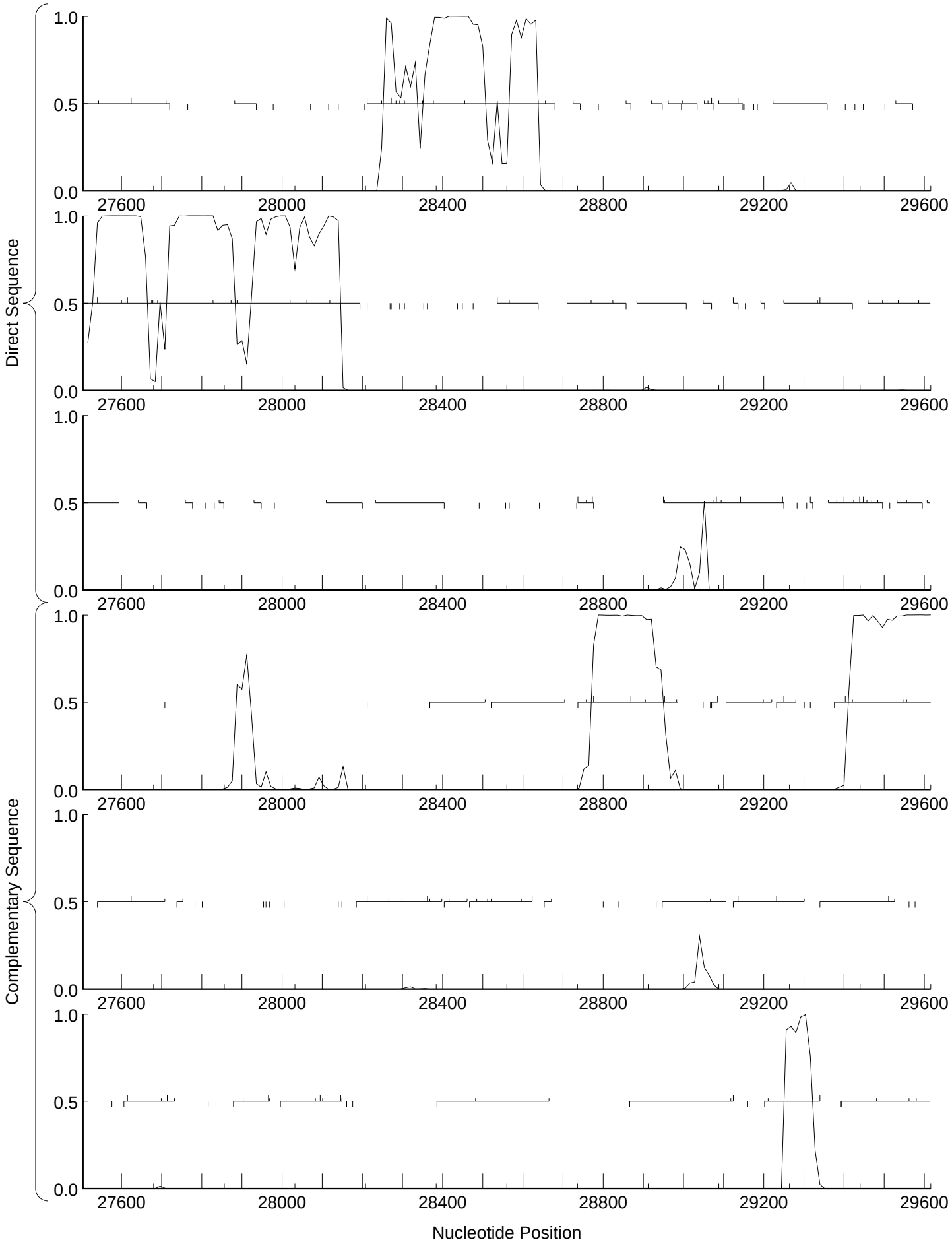
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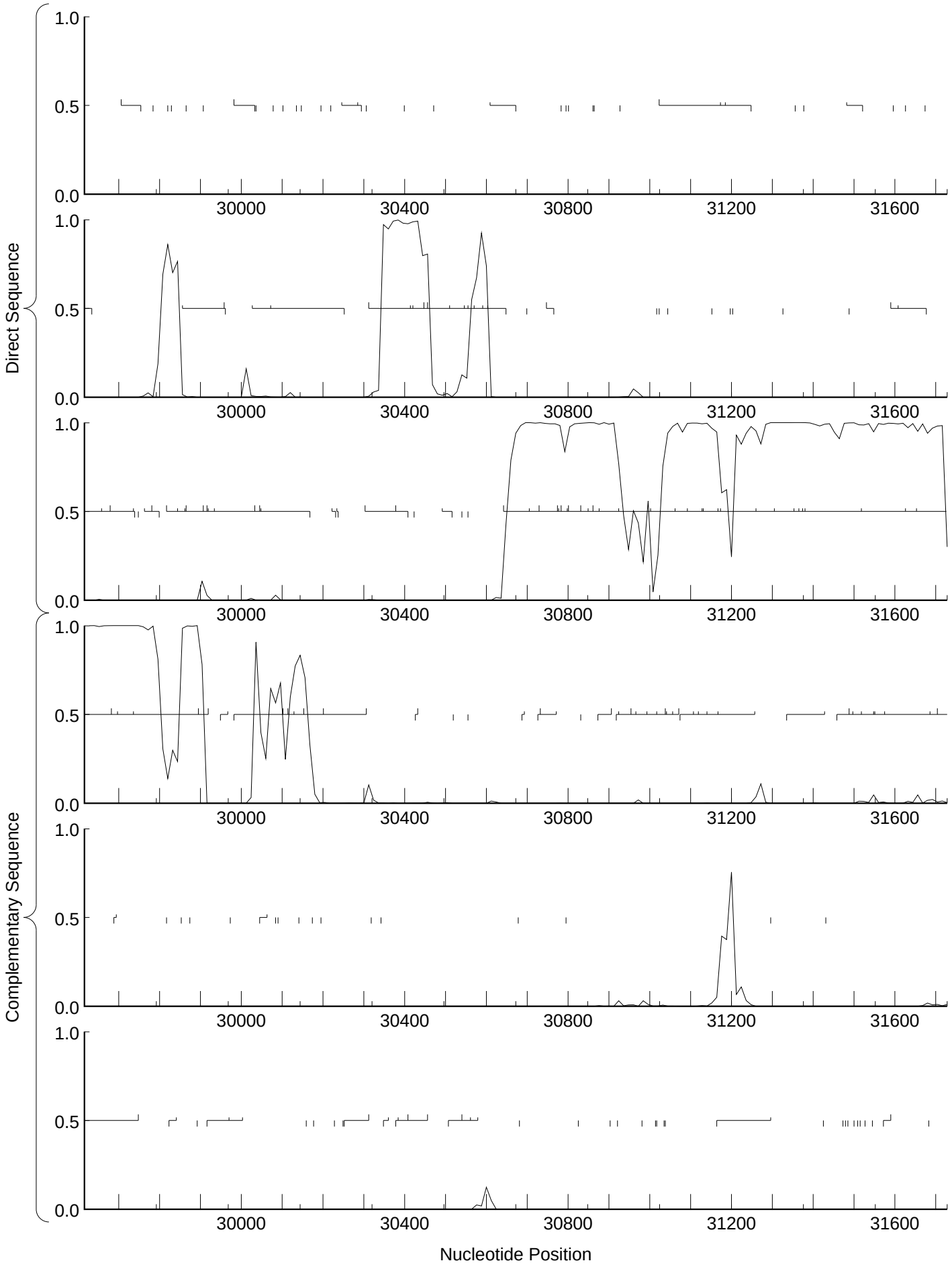
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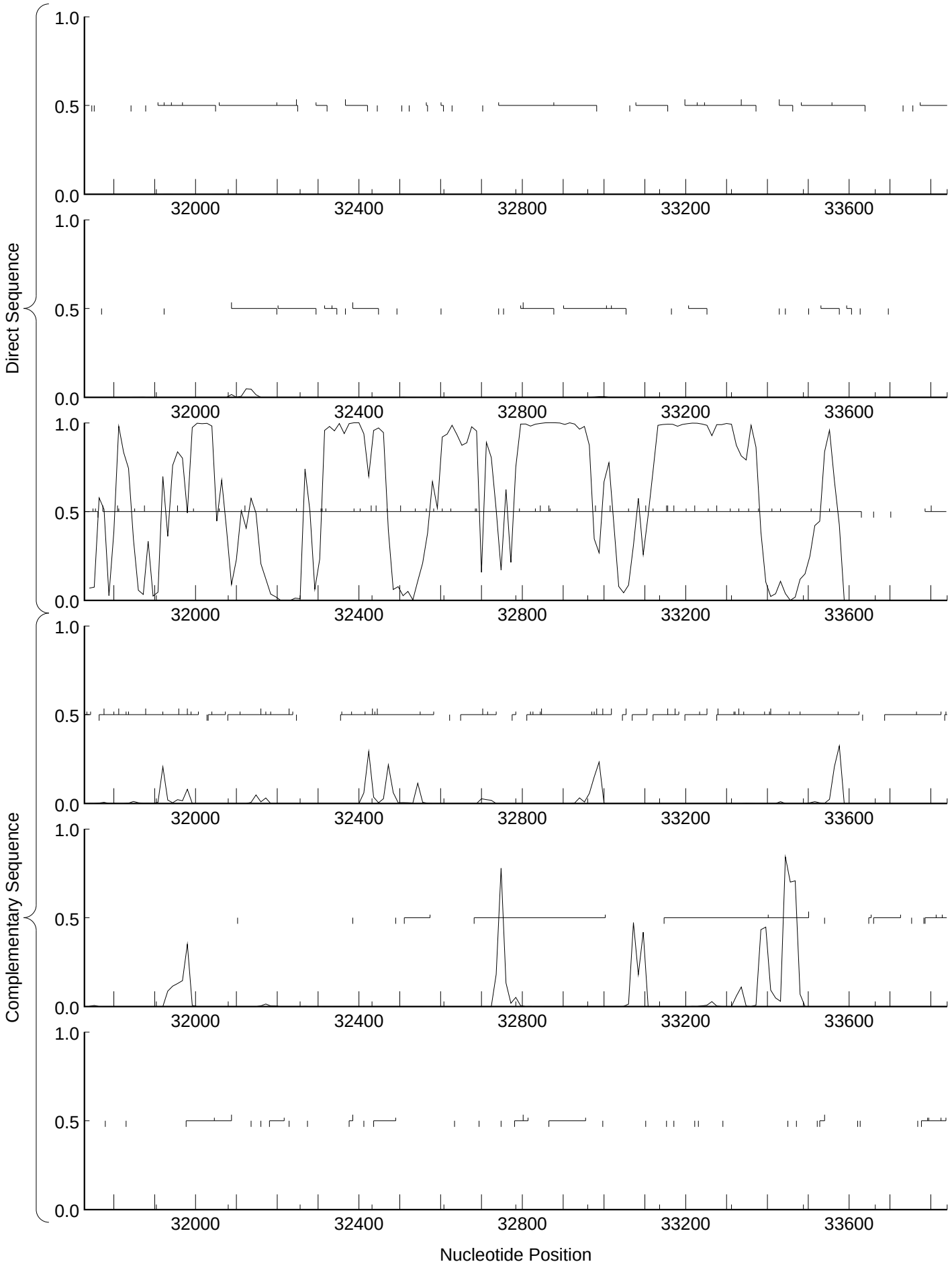
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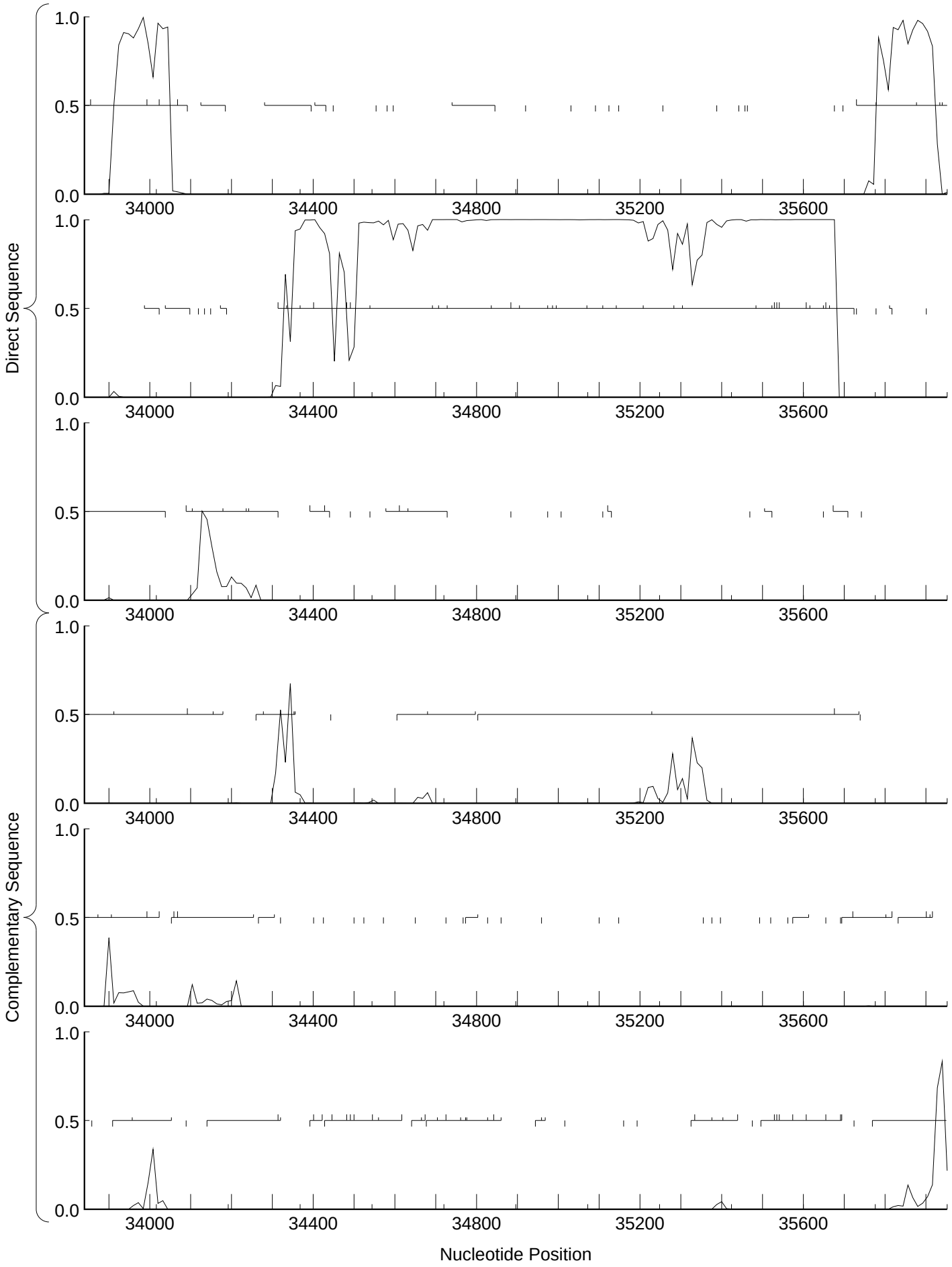
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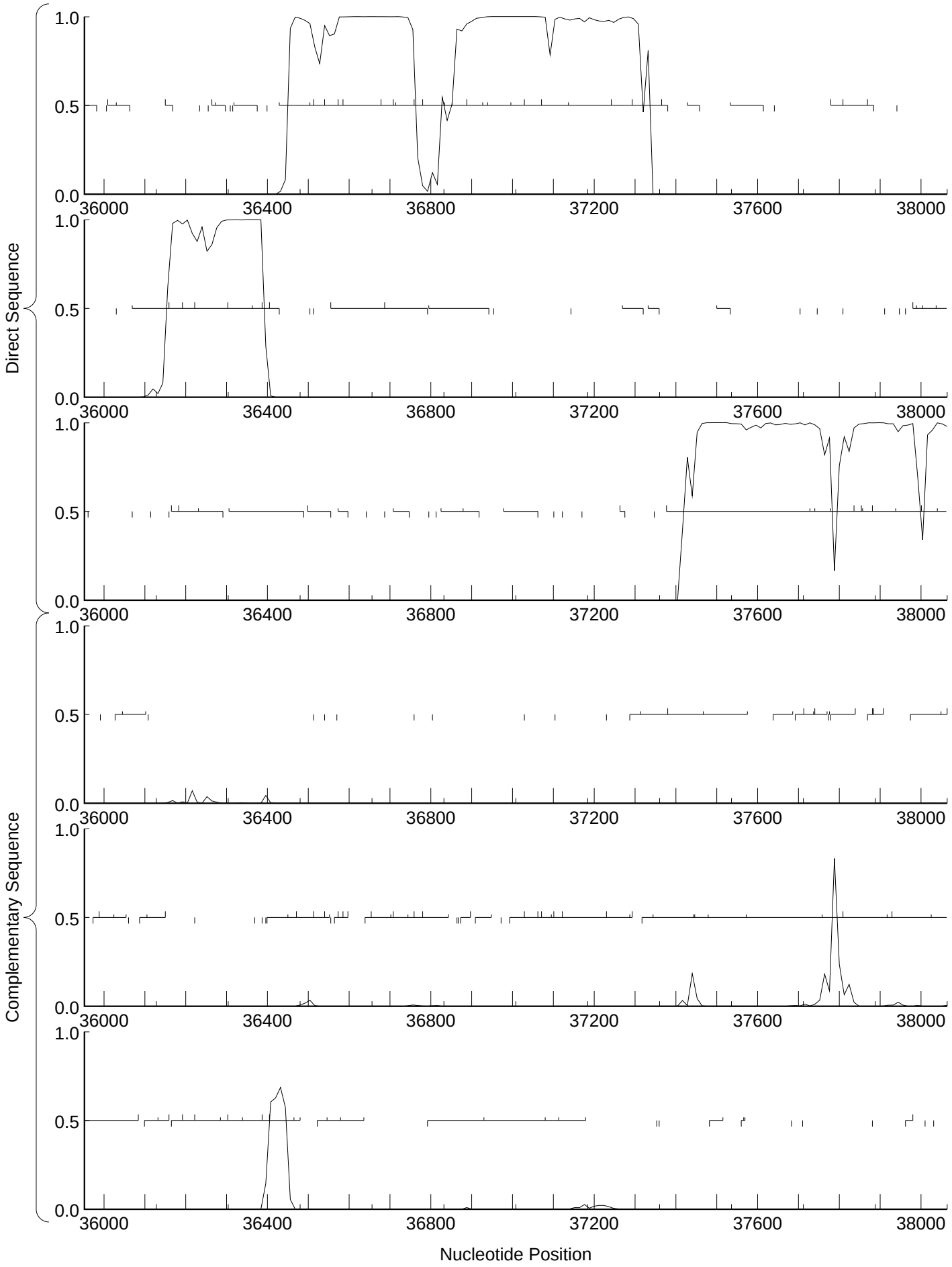
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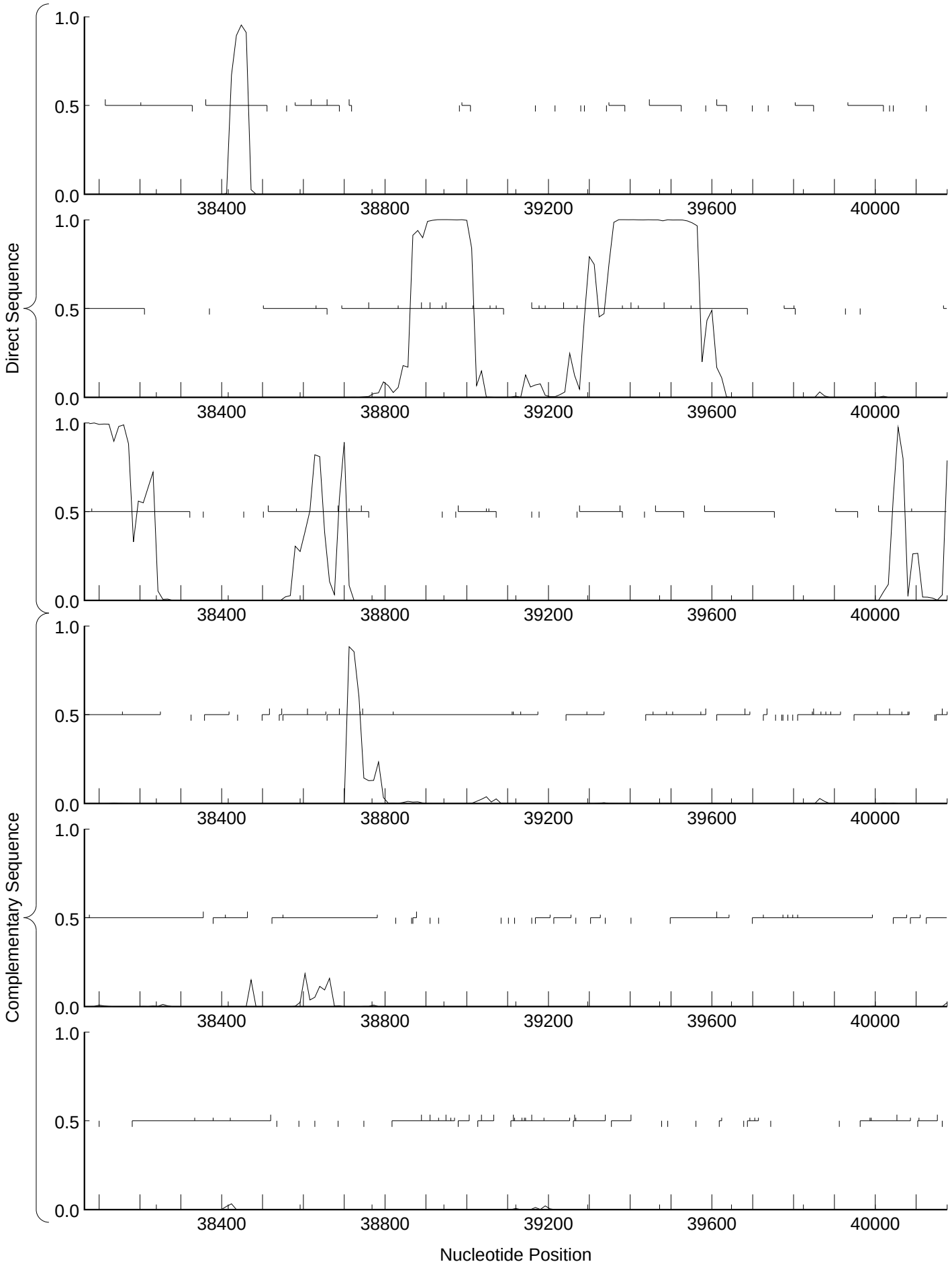
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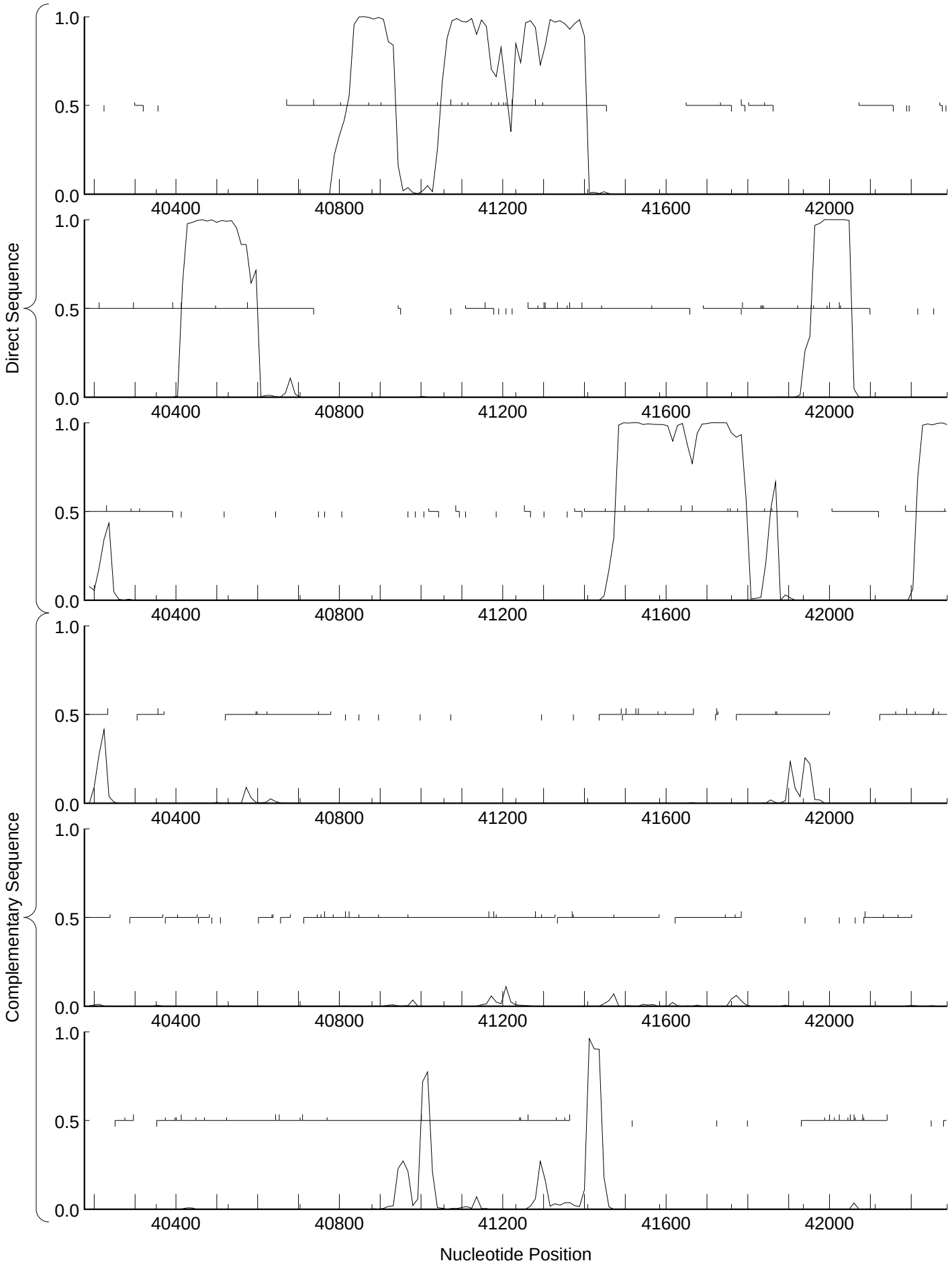
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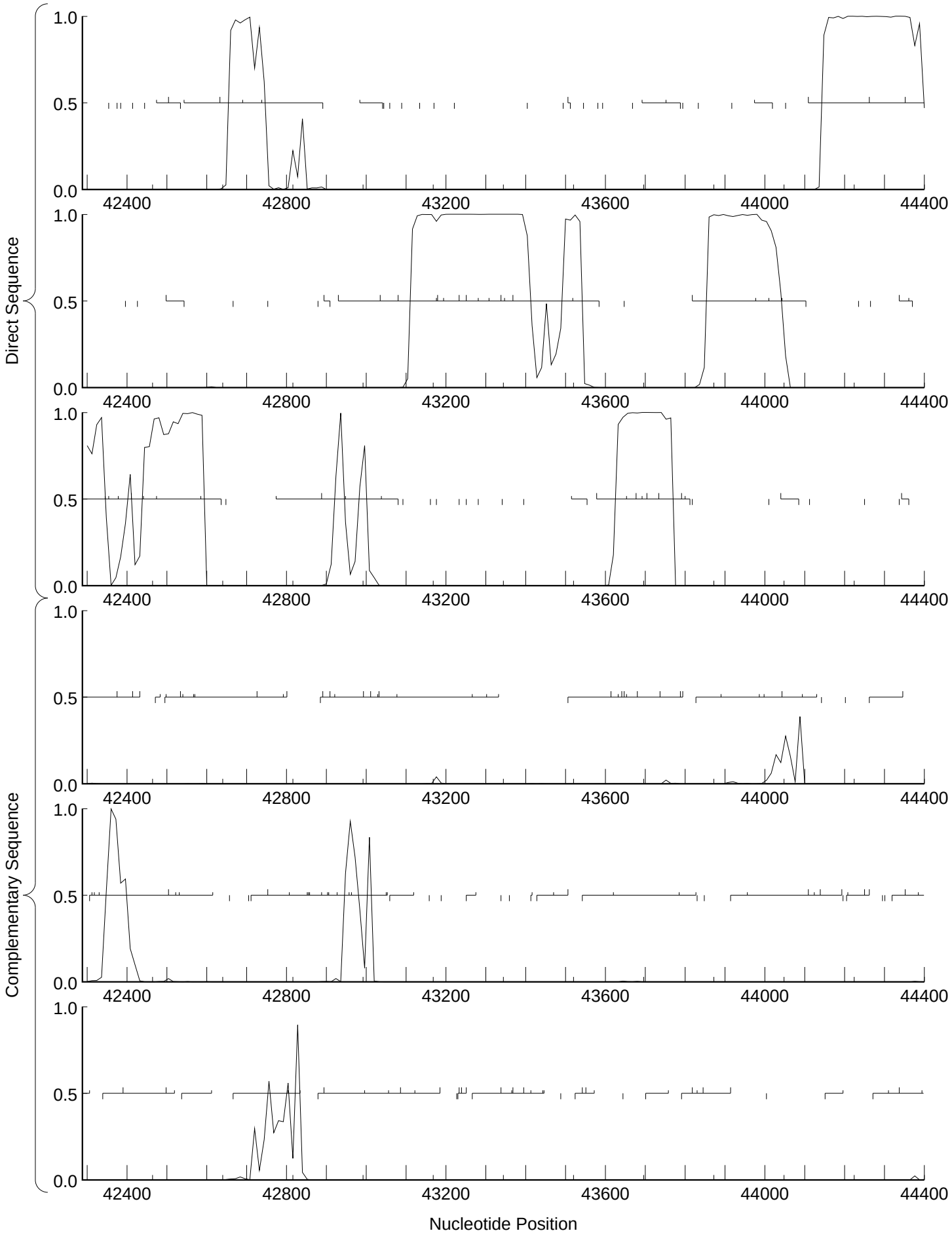
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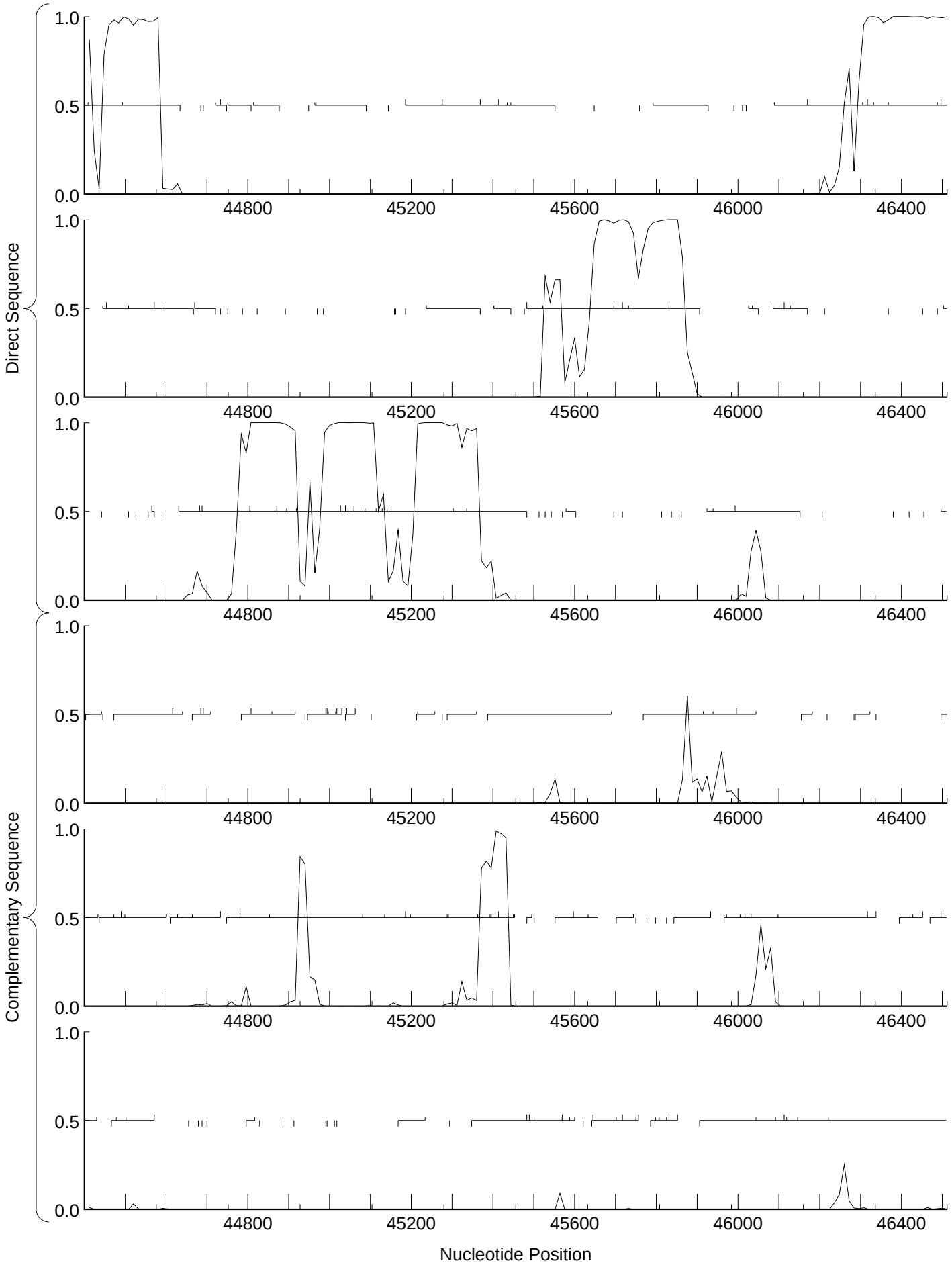
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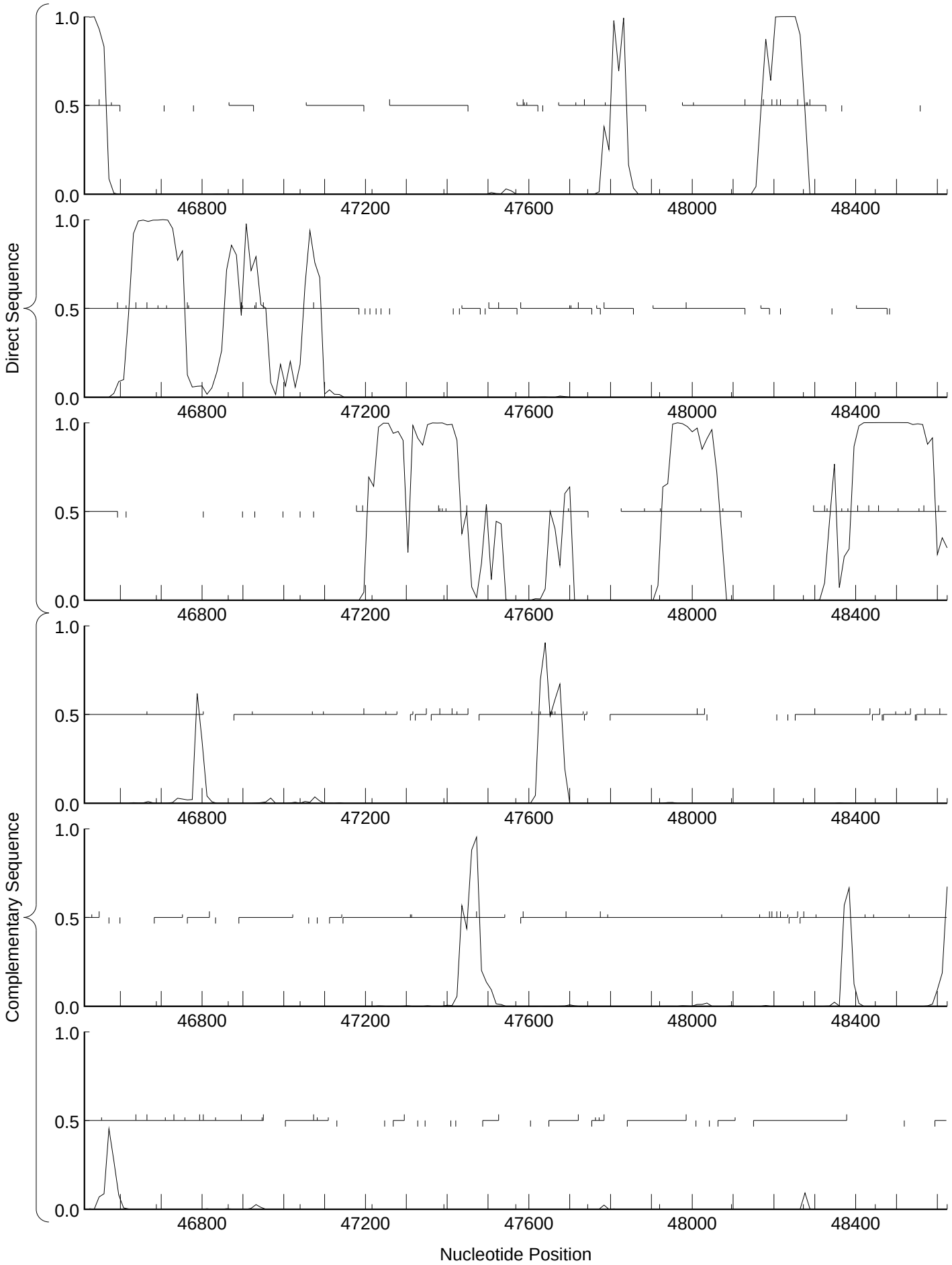
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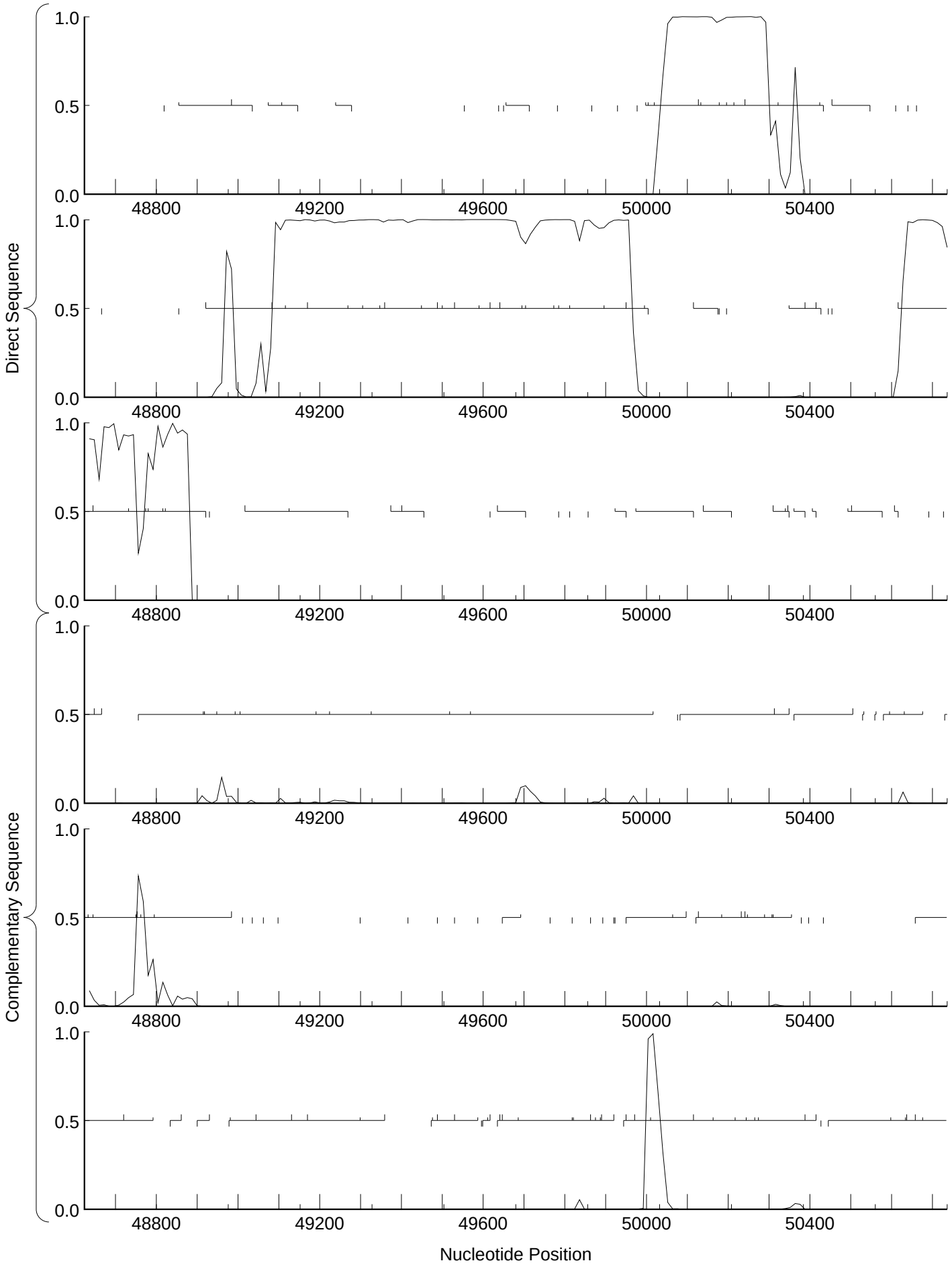
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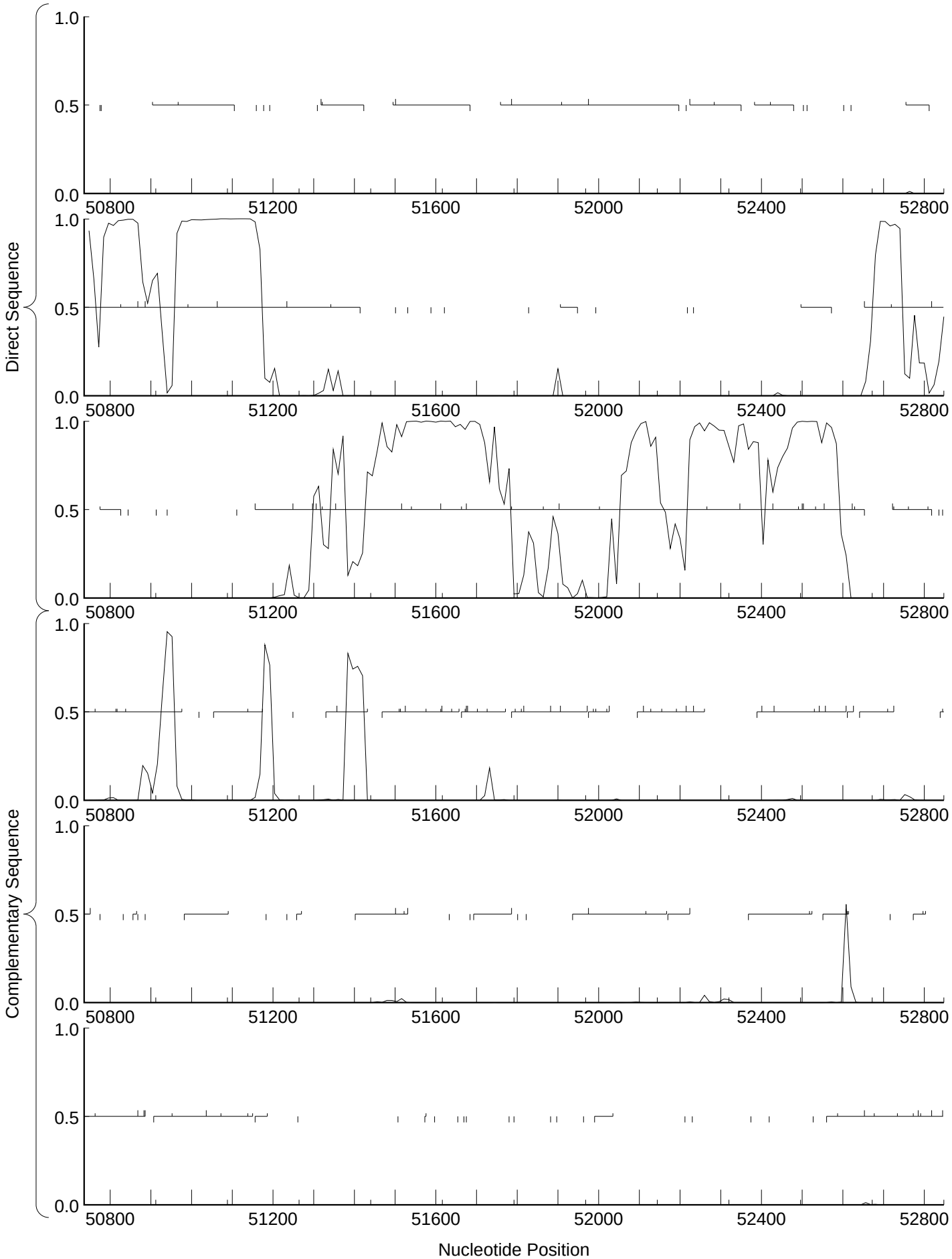
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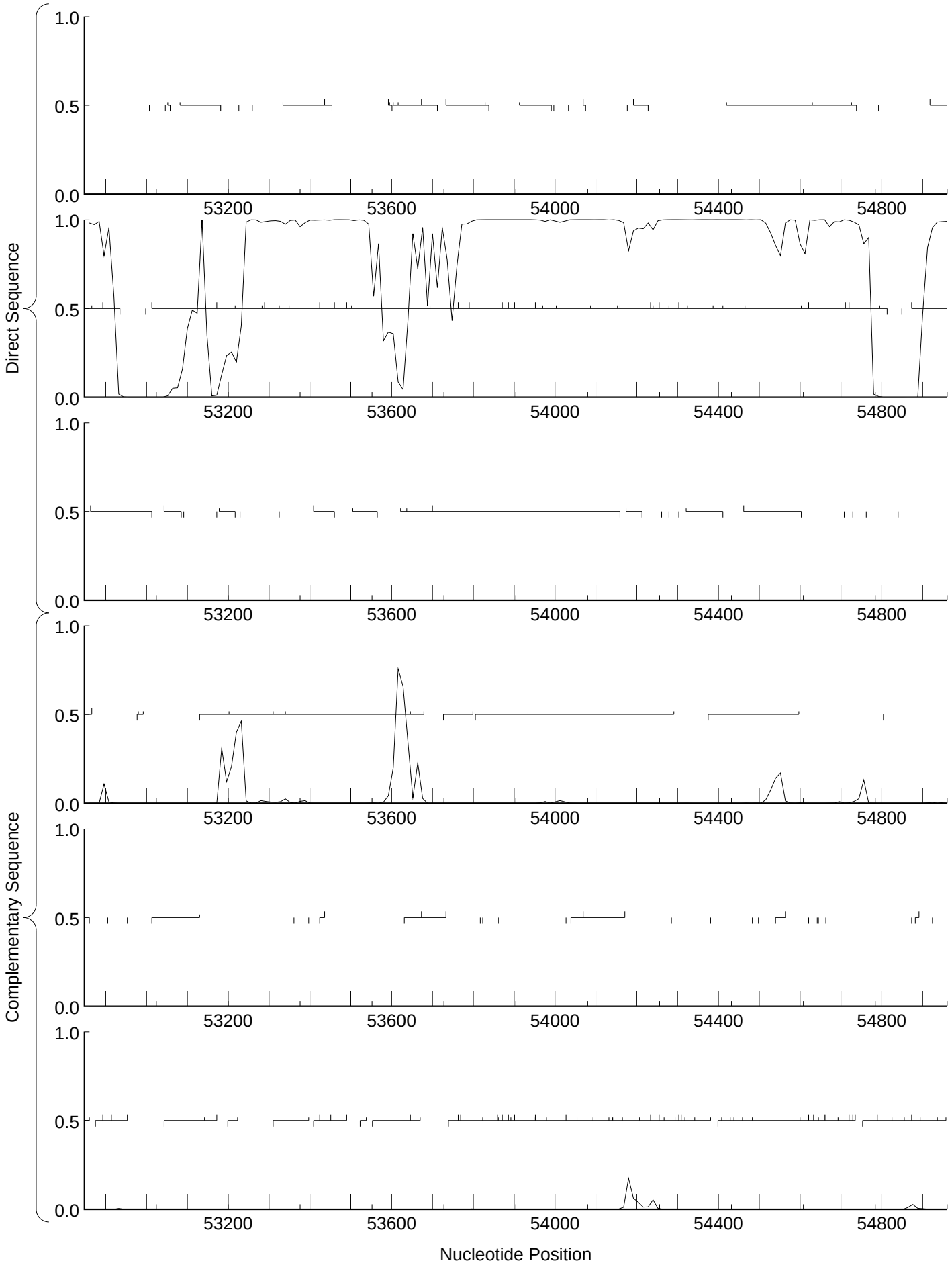
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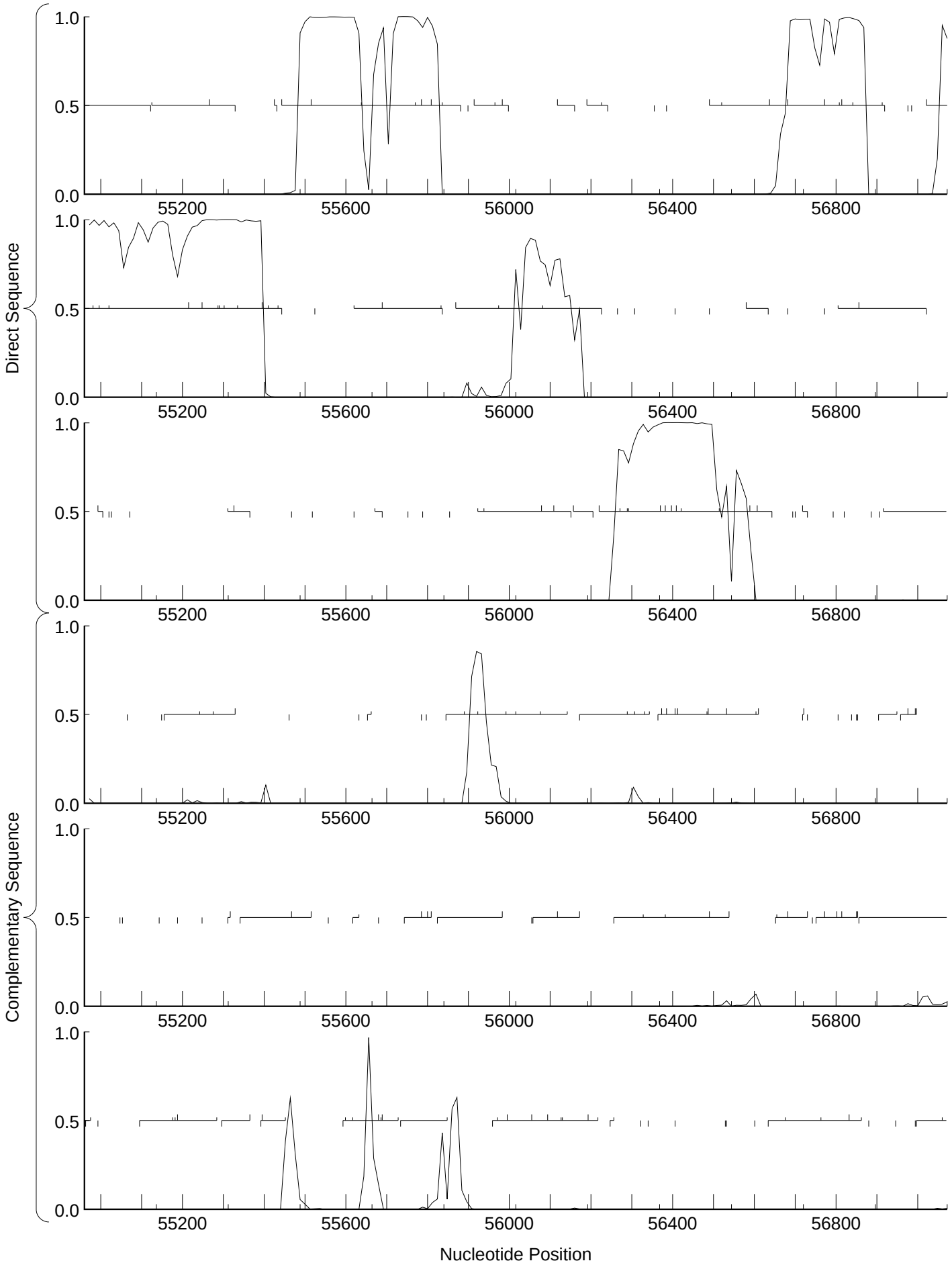
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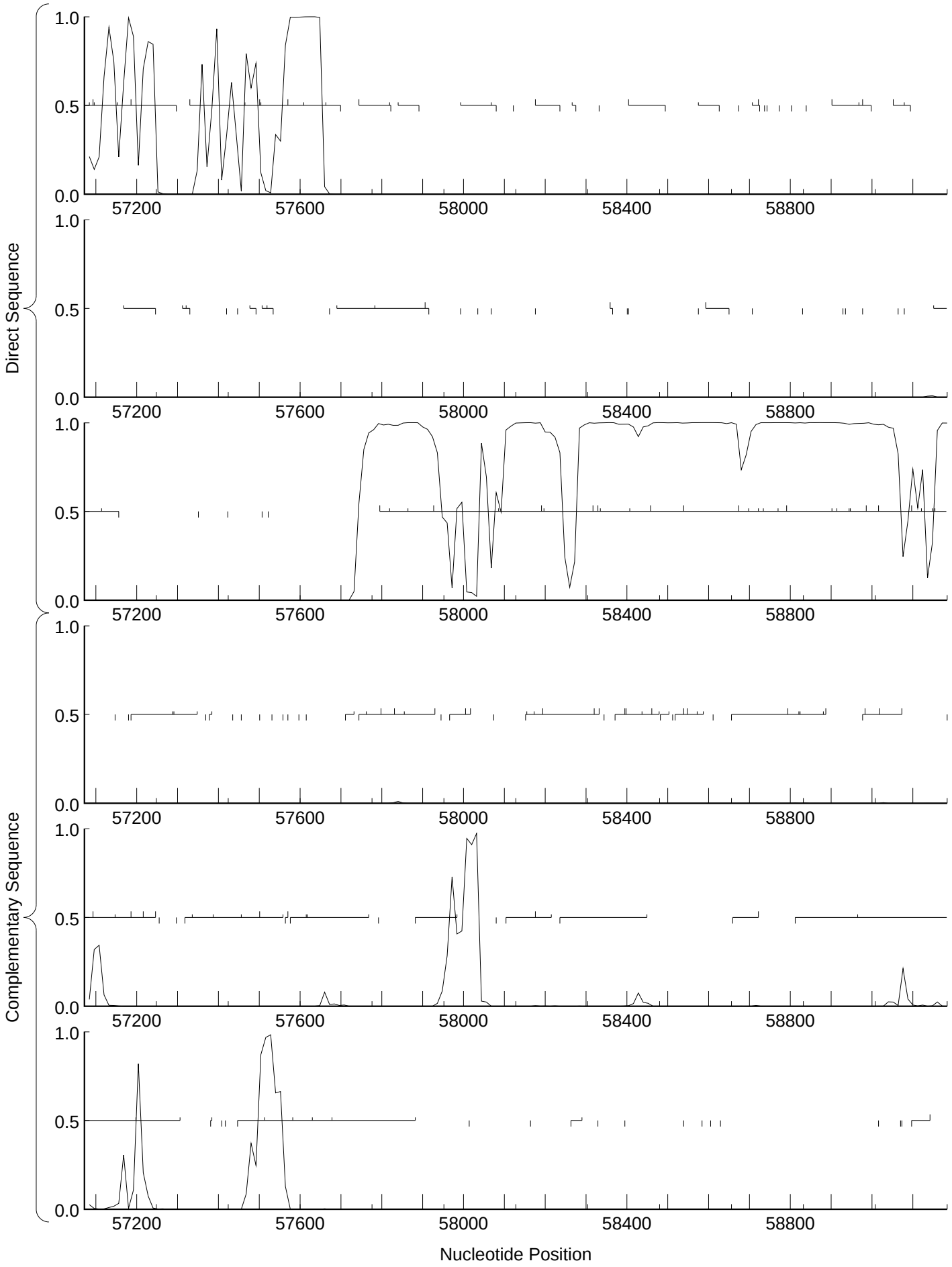
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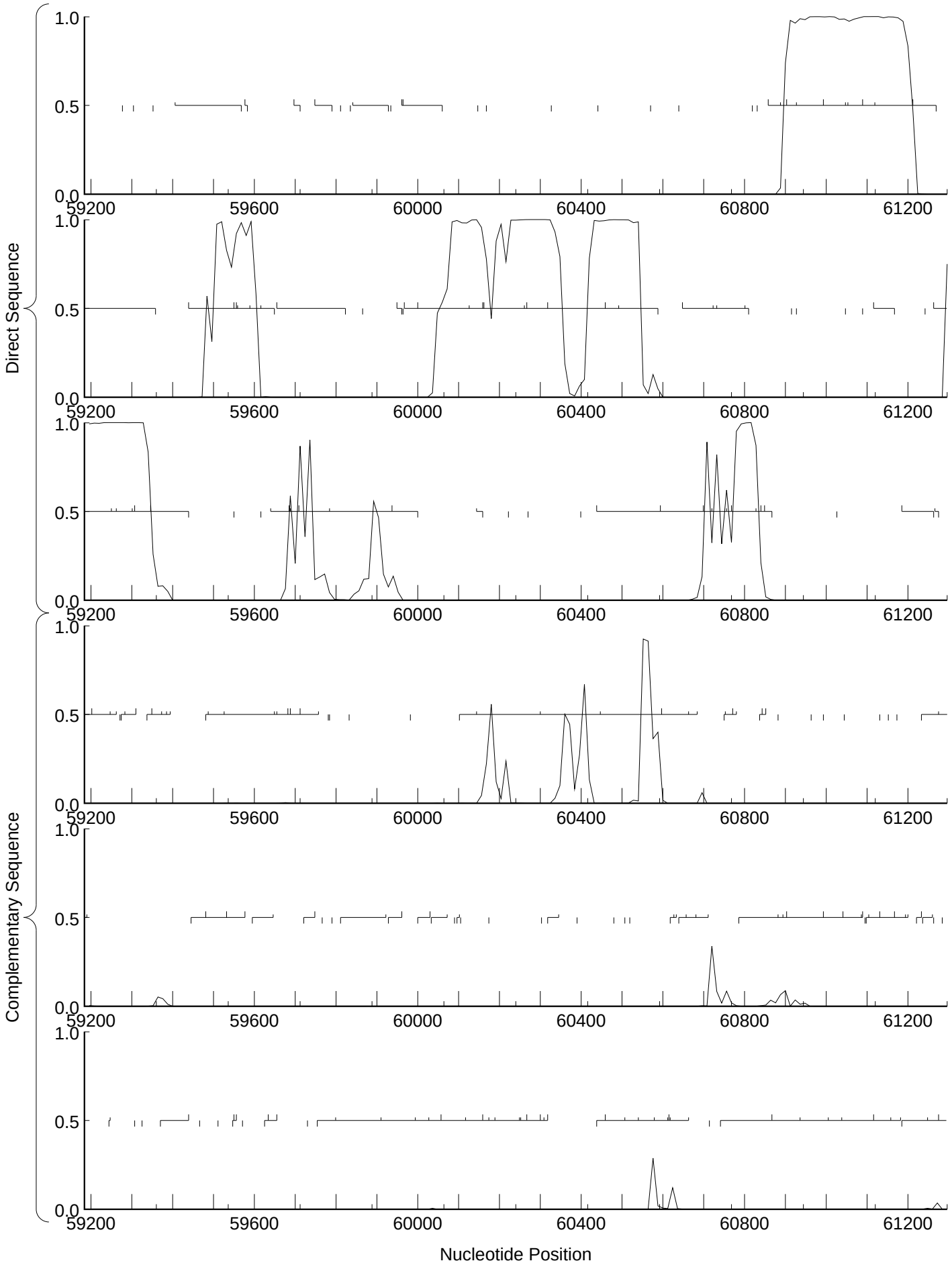
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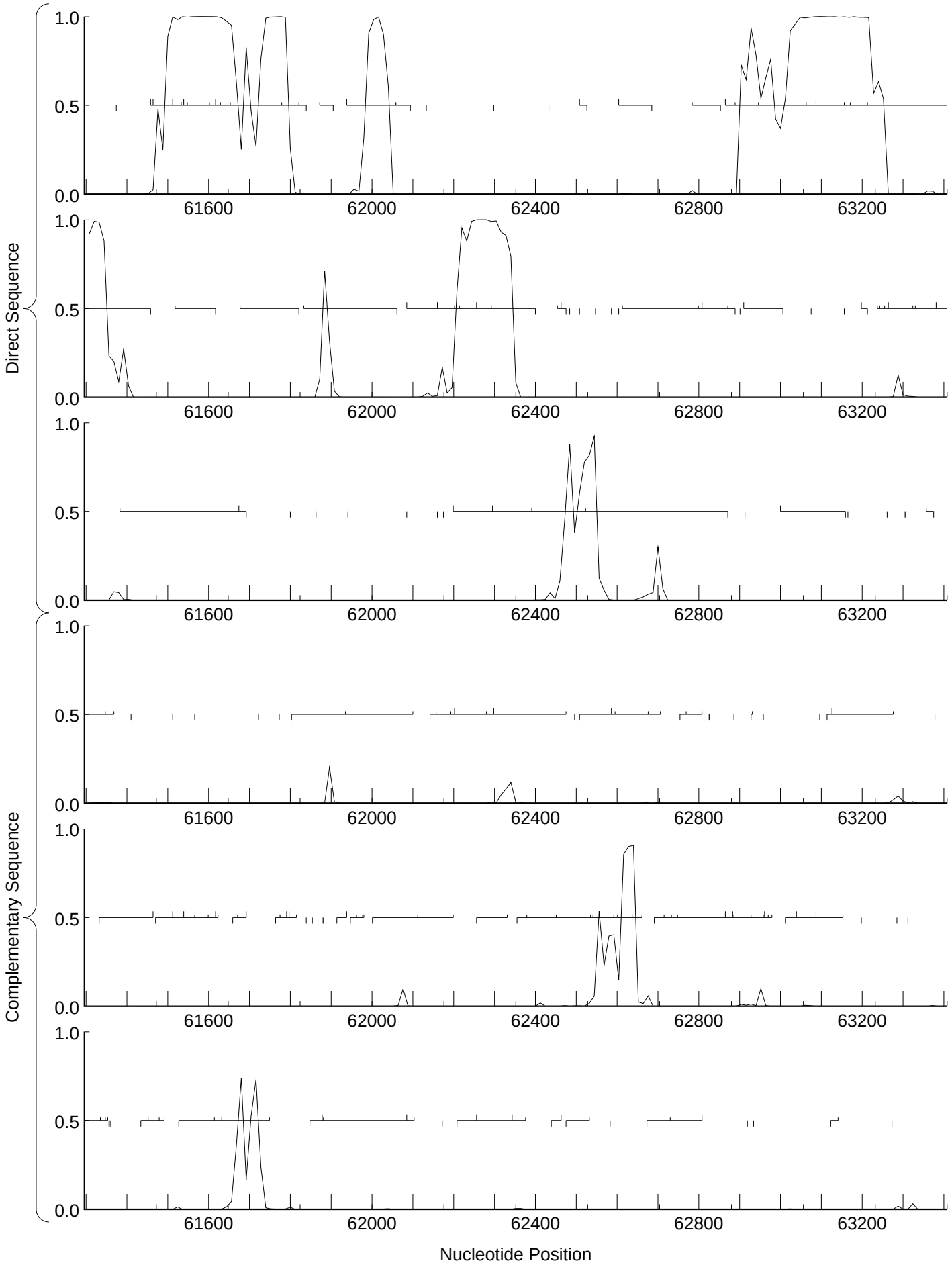
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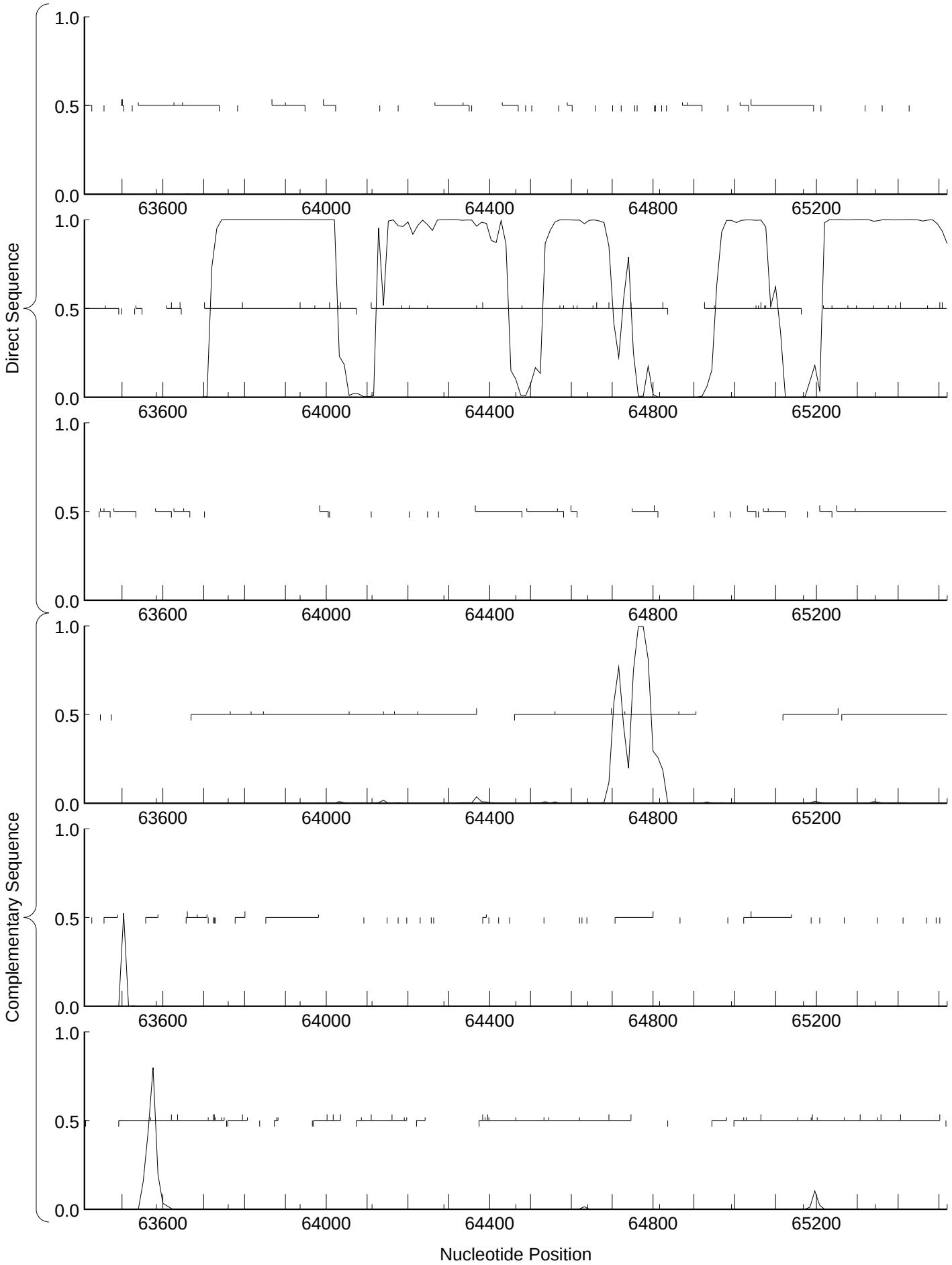
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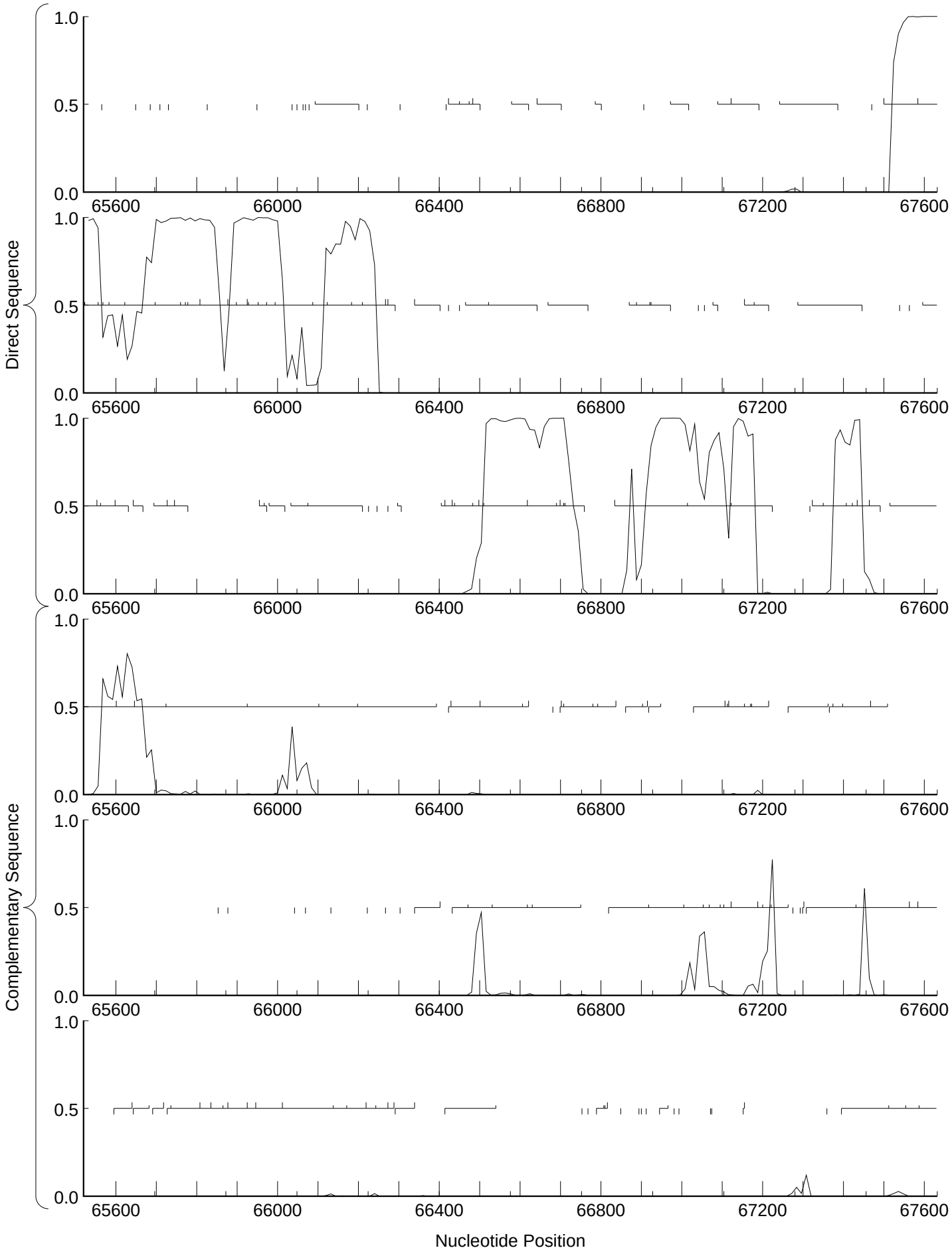
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Arthrobacter phage Lymara complete sequence, 68762 bp, circularly permuted, Cluster AR, Order 4, Window 96, Step 12, 32/34



Arthrobacter phage Lymara complete sequence, 68762 bp, circularly permuted, Cluster AR, Order 4, Window 96, Step 12, 33/34



Arthrobacter phage Lymara complete sequence, 68762 bp, circularly permuted, Cluster AR, Order 4, Window 96, Step 12, 34/34

