GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage LugYA complete sequence, 50878 bp including 10-base 3’ overhang (CGGGTG
Analysis Date    :  3/29/24 at 8:34:57
Pages            :  26
Sequence Length  :  50878 bp
GC Content       :   64.00%

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

PS-Version       :  1.2

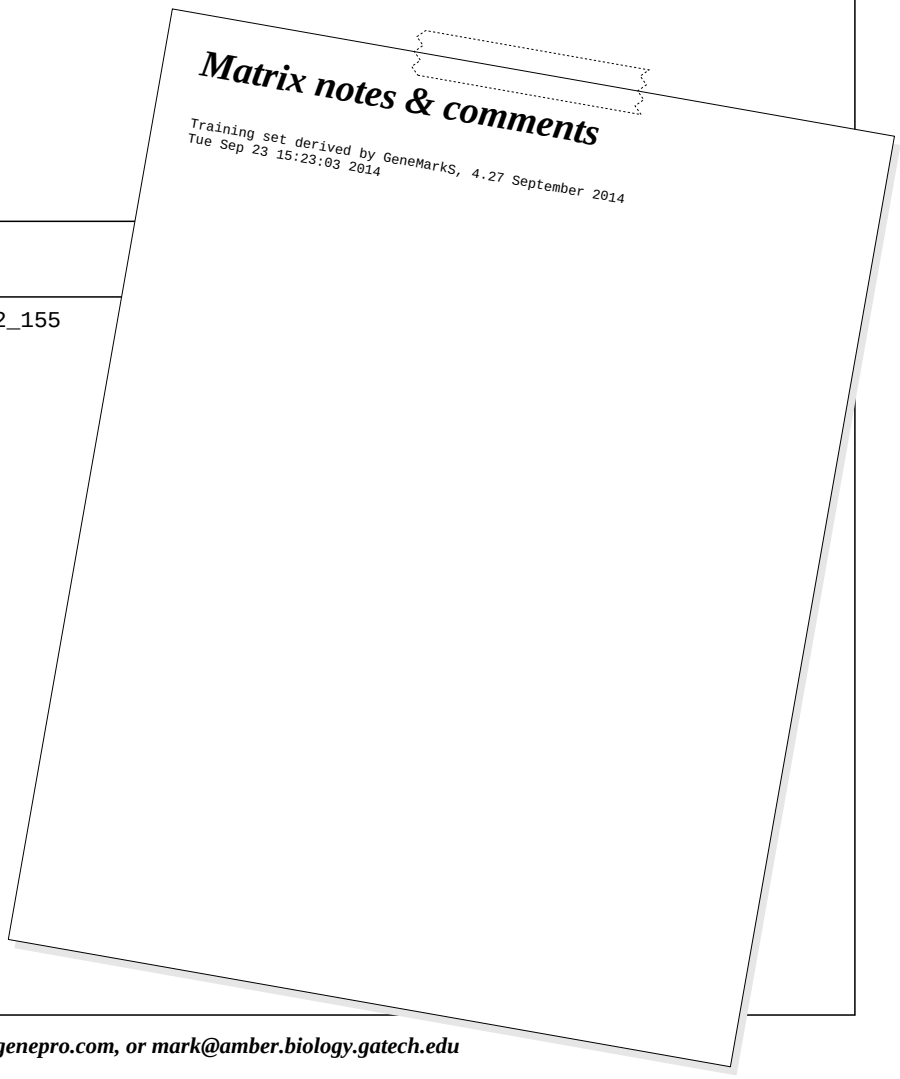
GeneMark Options :  PostScript graph,
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MATRIX INFORMATION

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Author     :  -

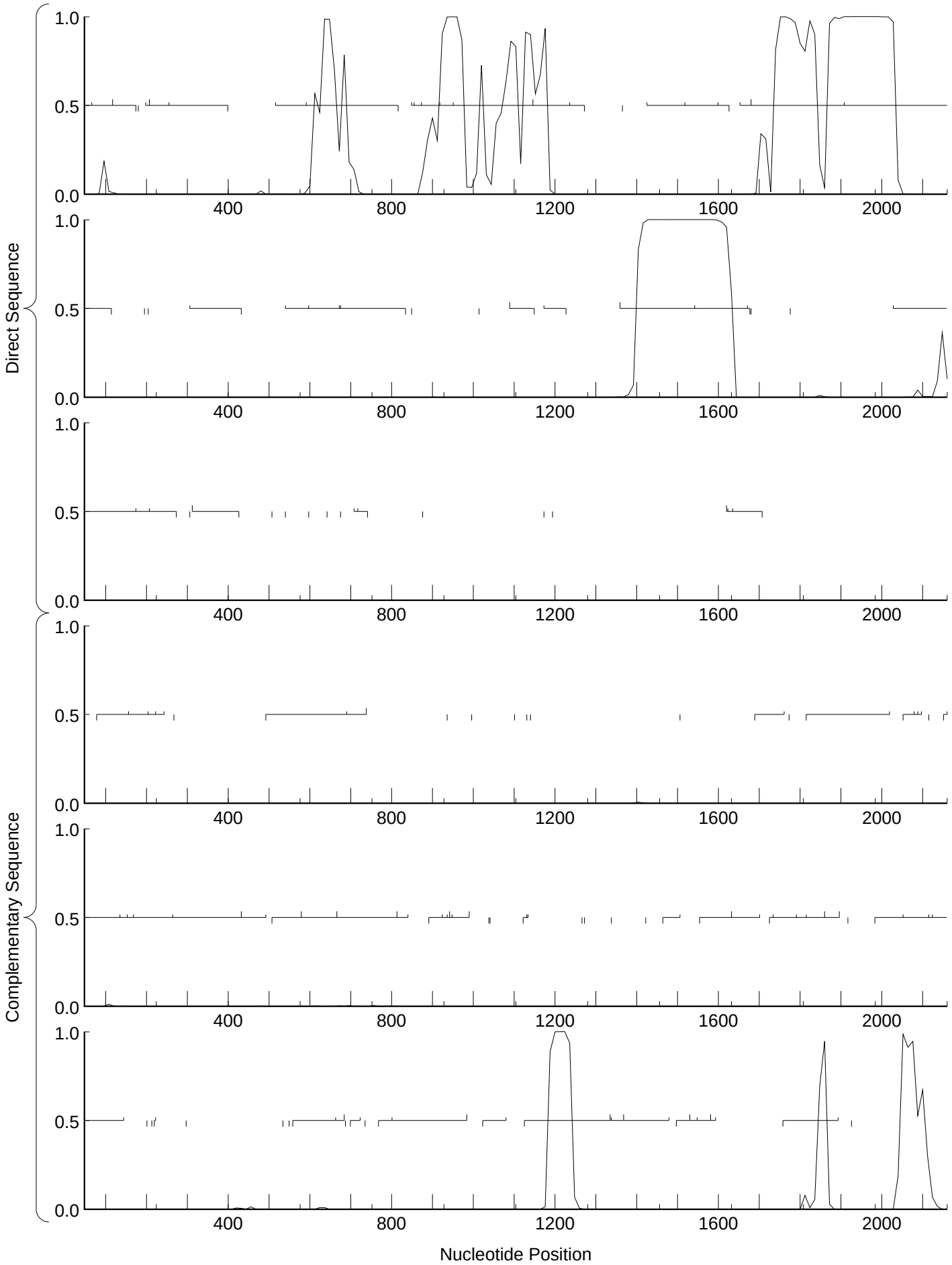
Order      :  4


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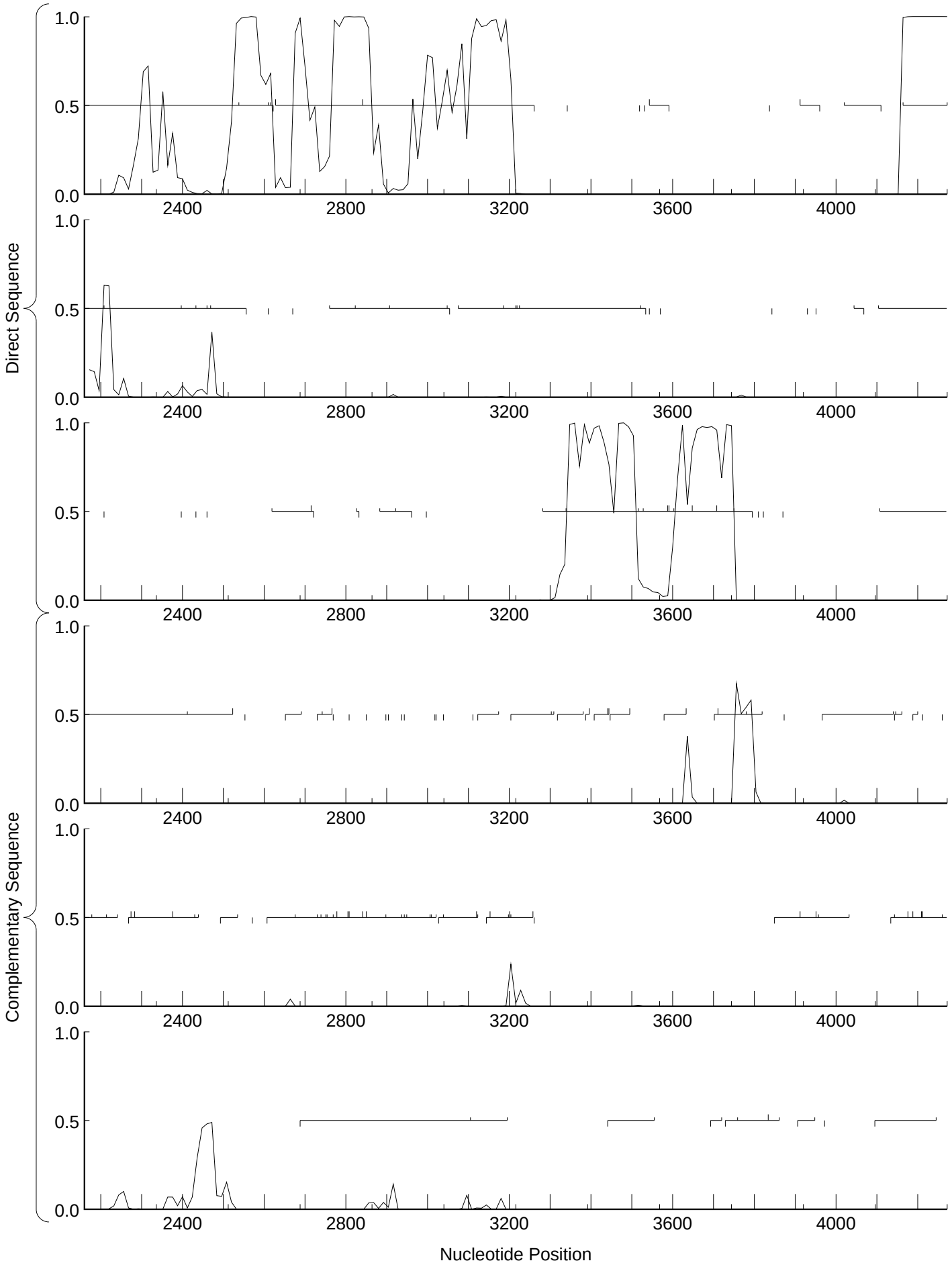


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

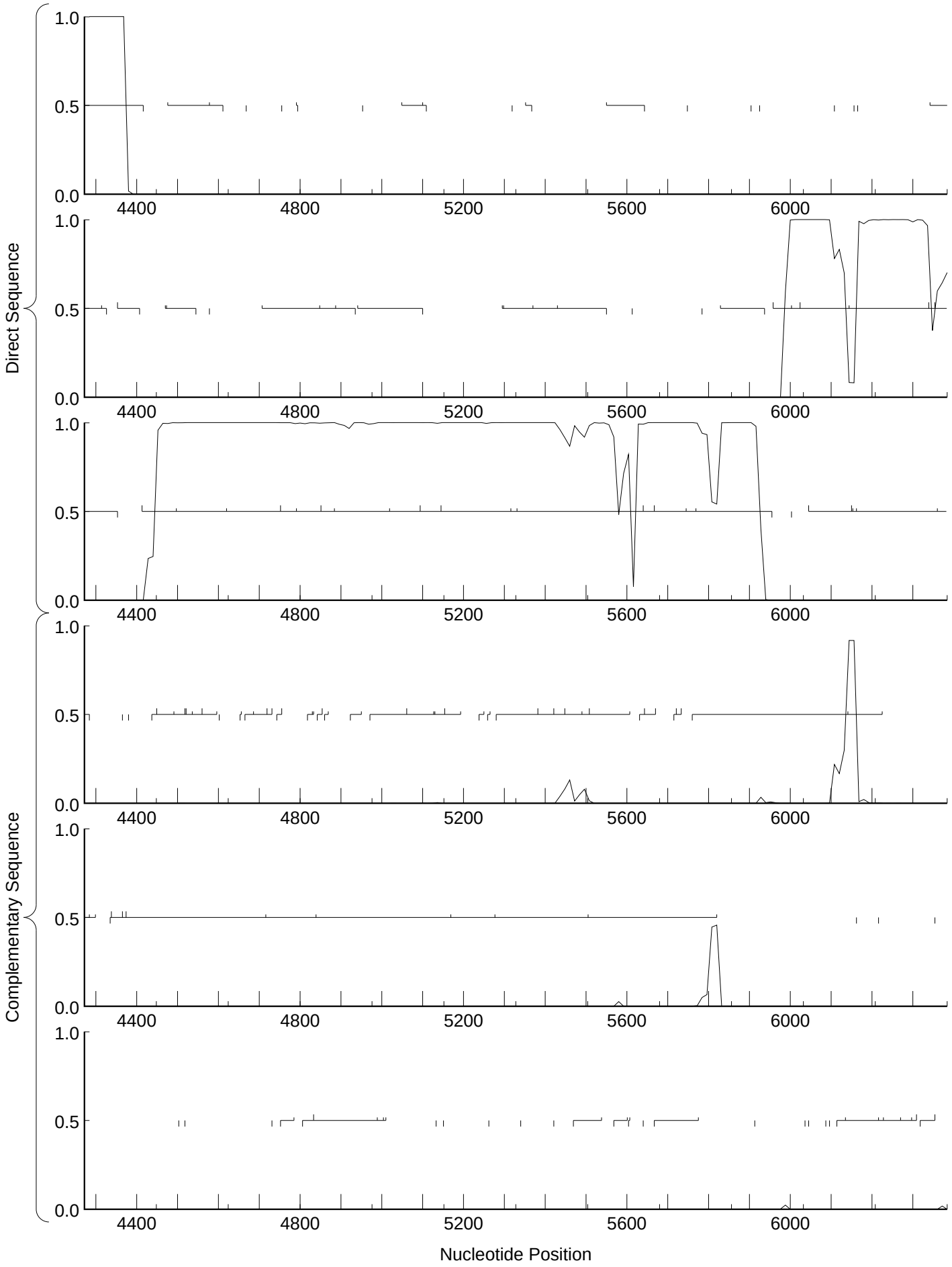
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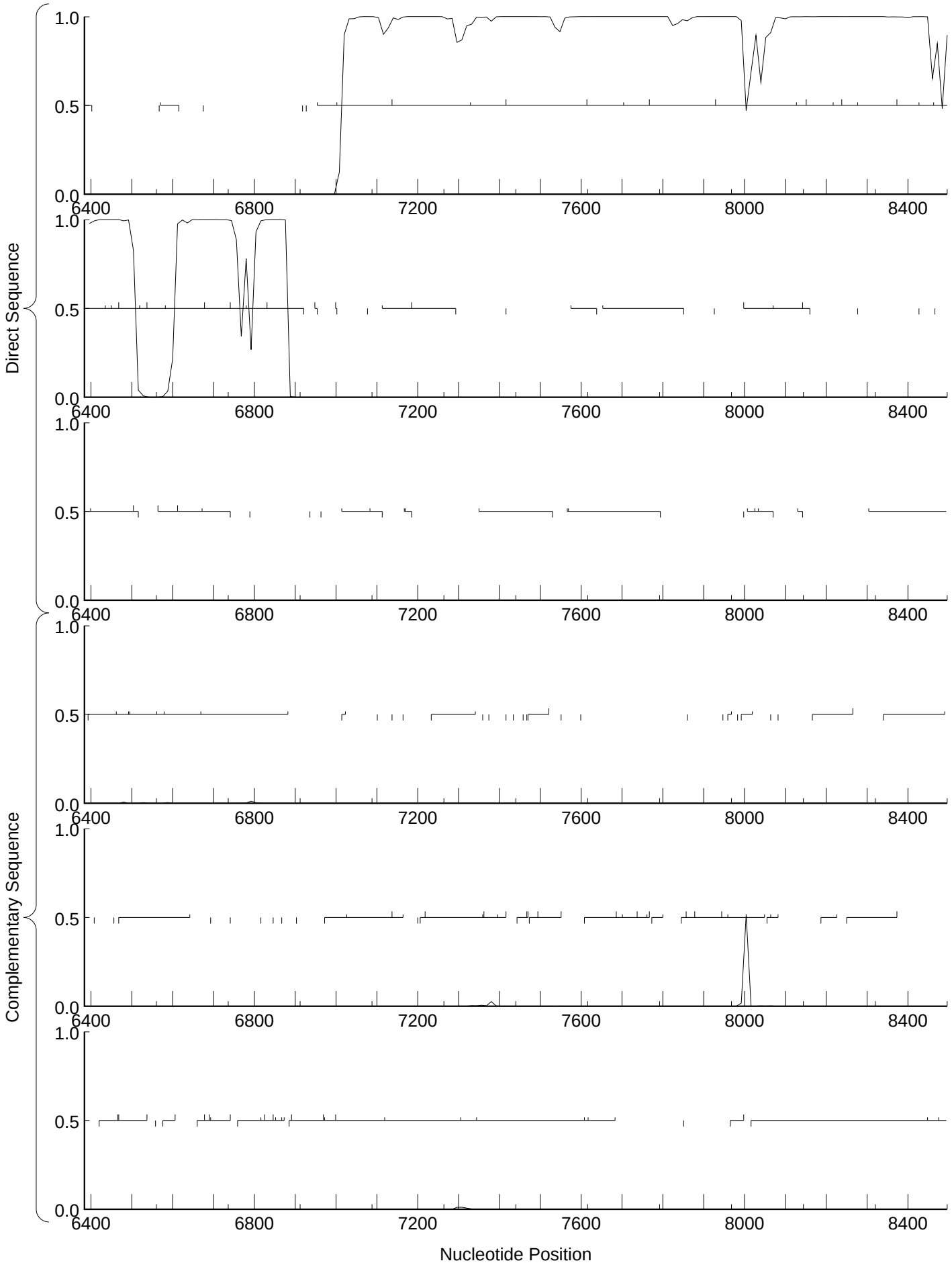
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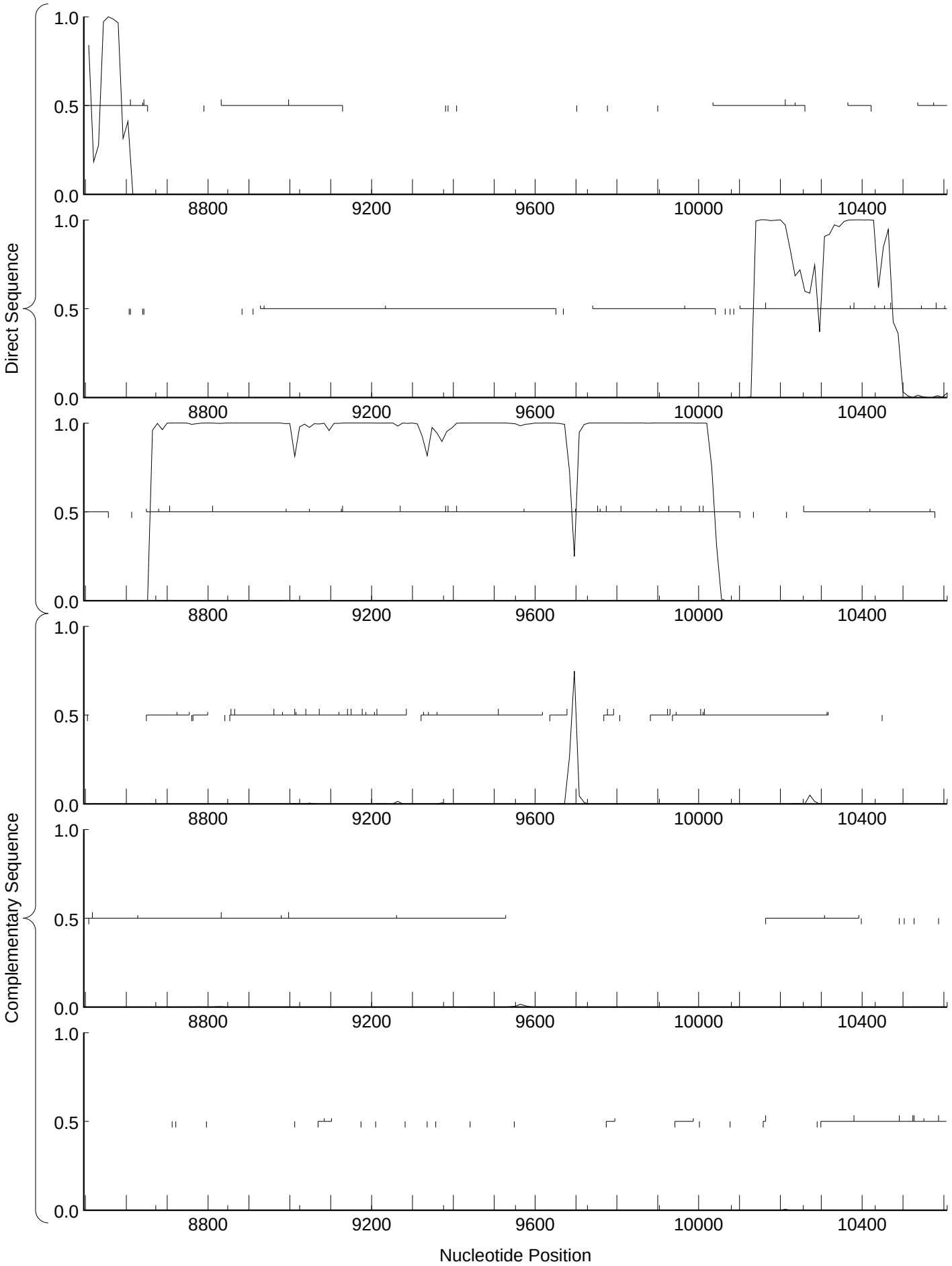
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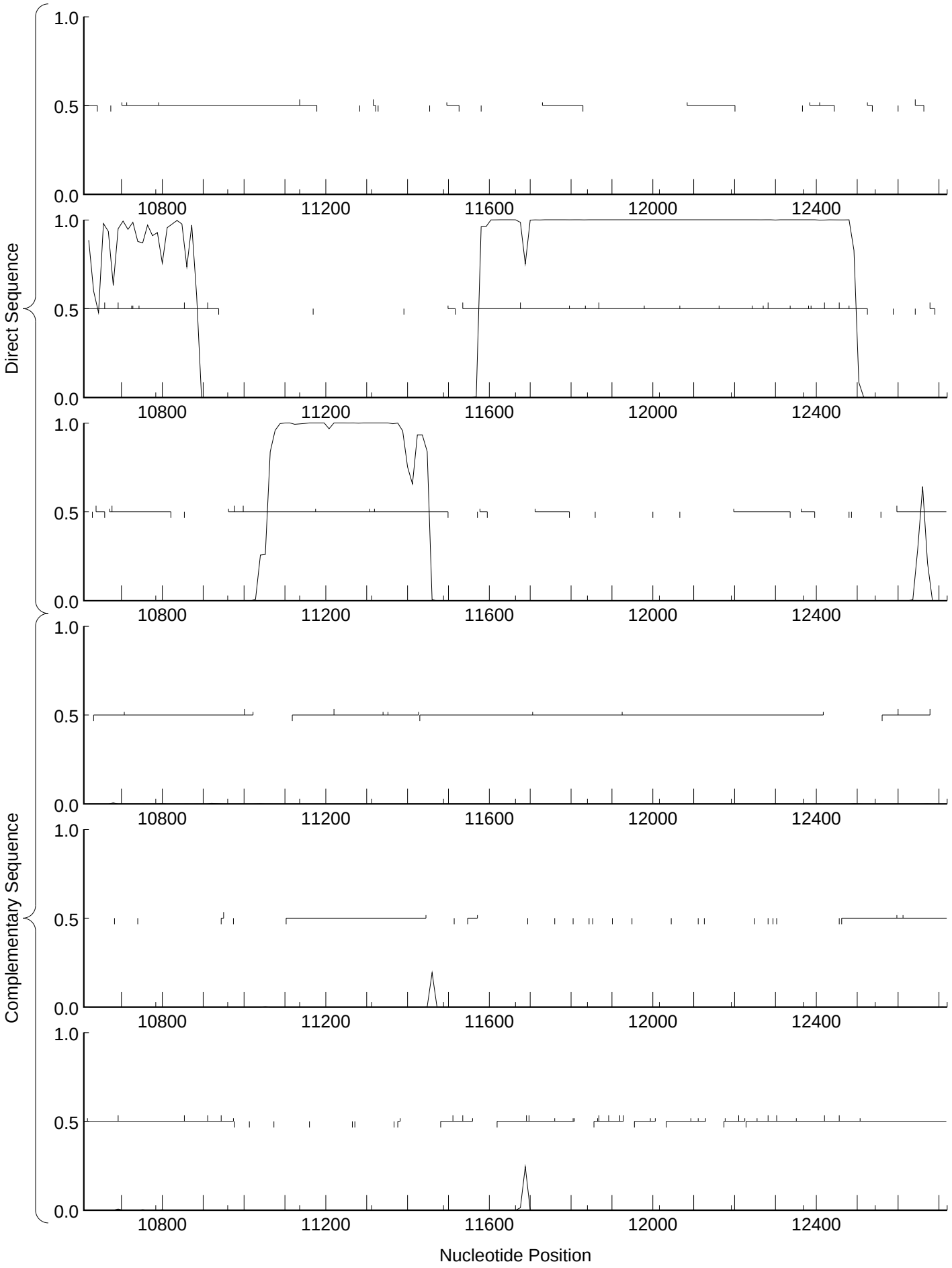
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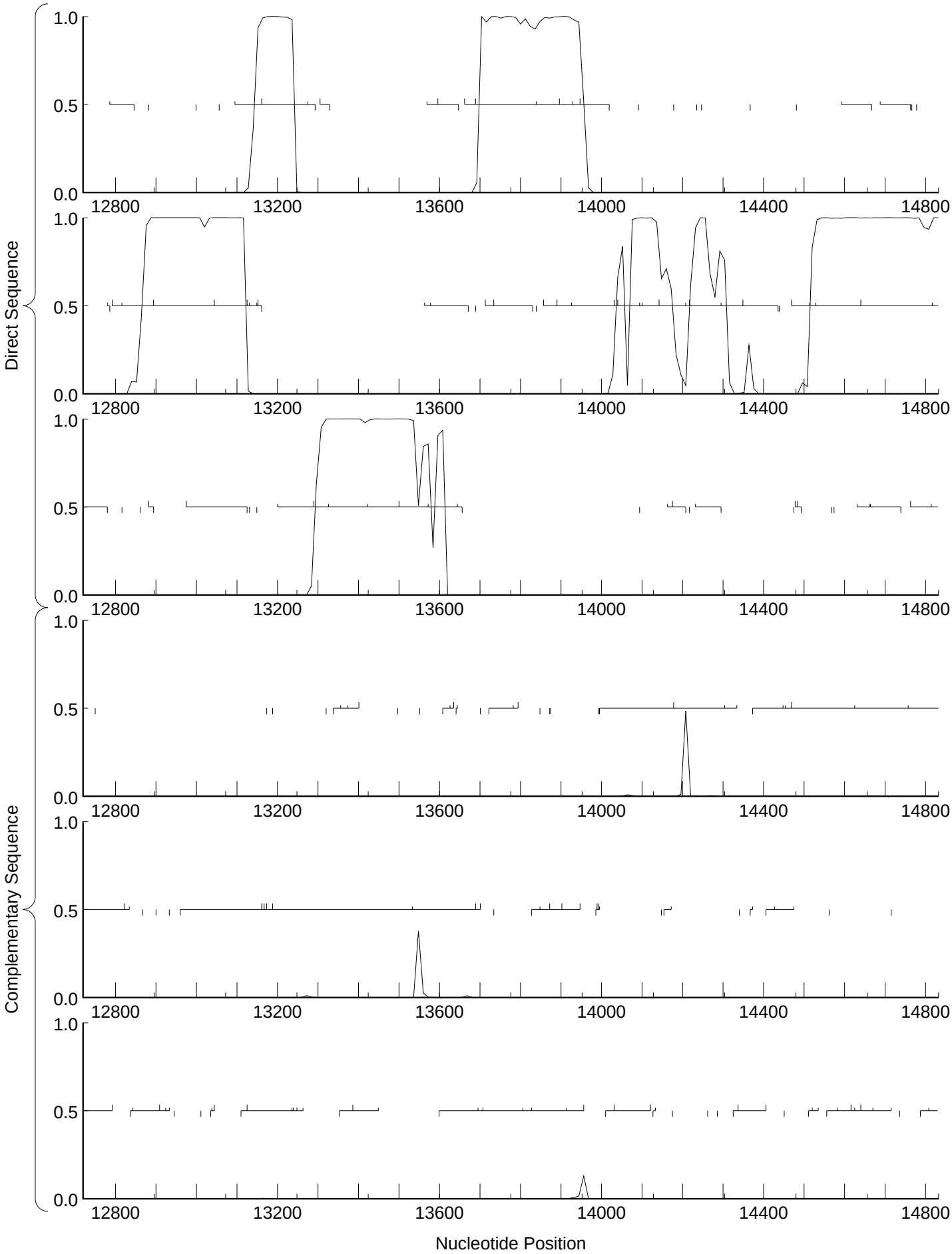
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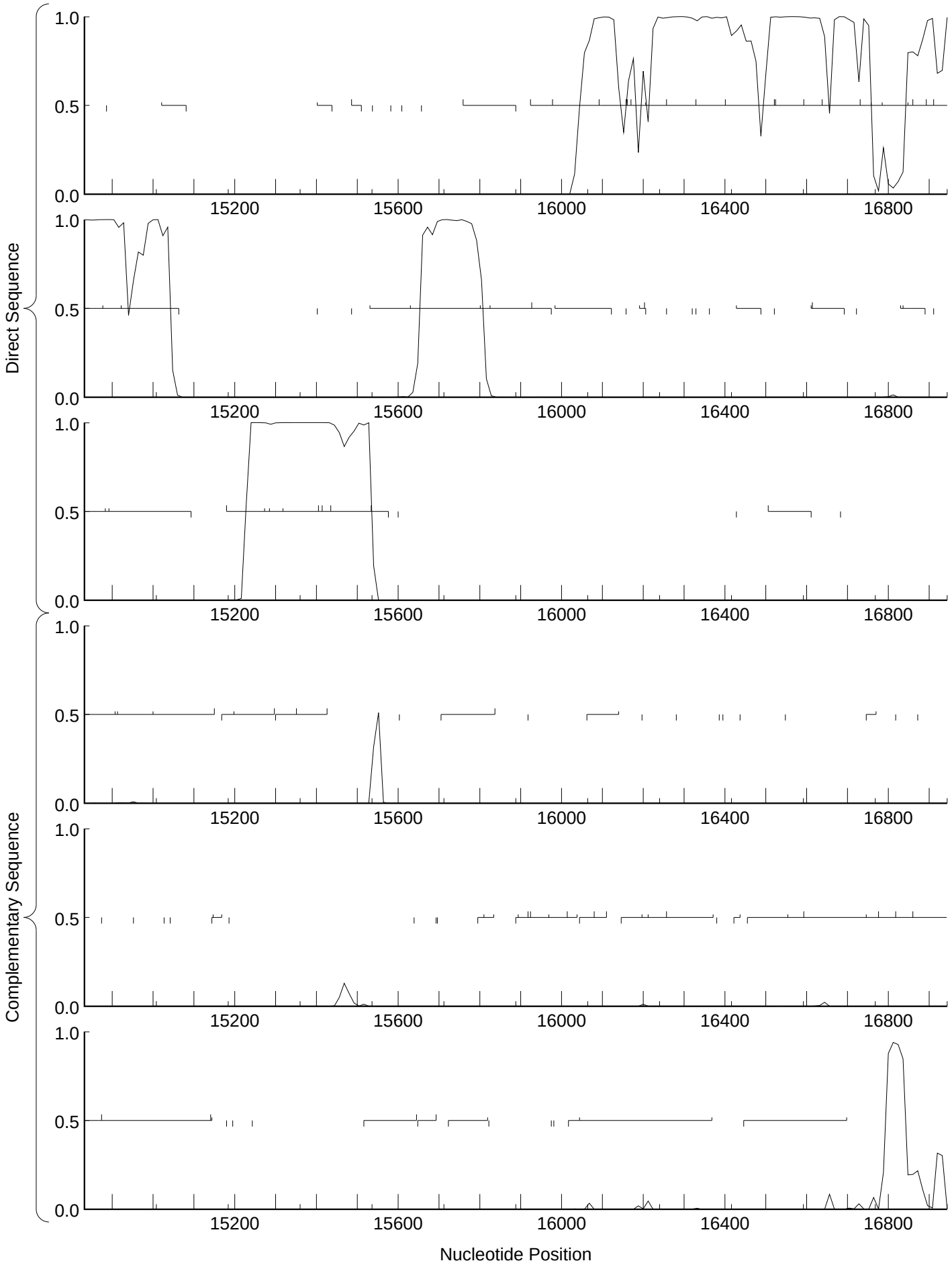
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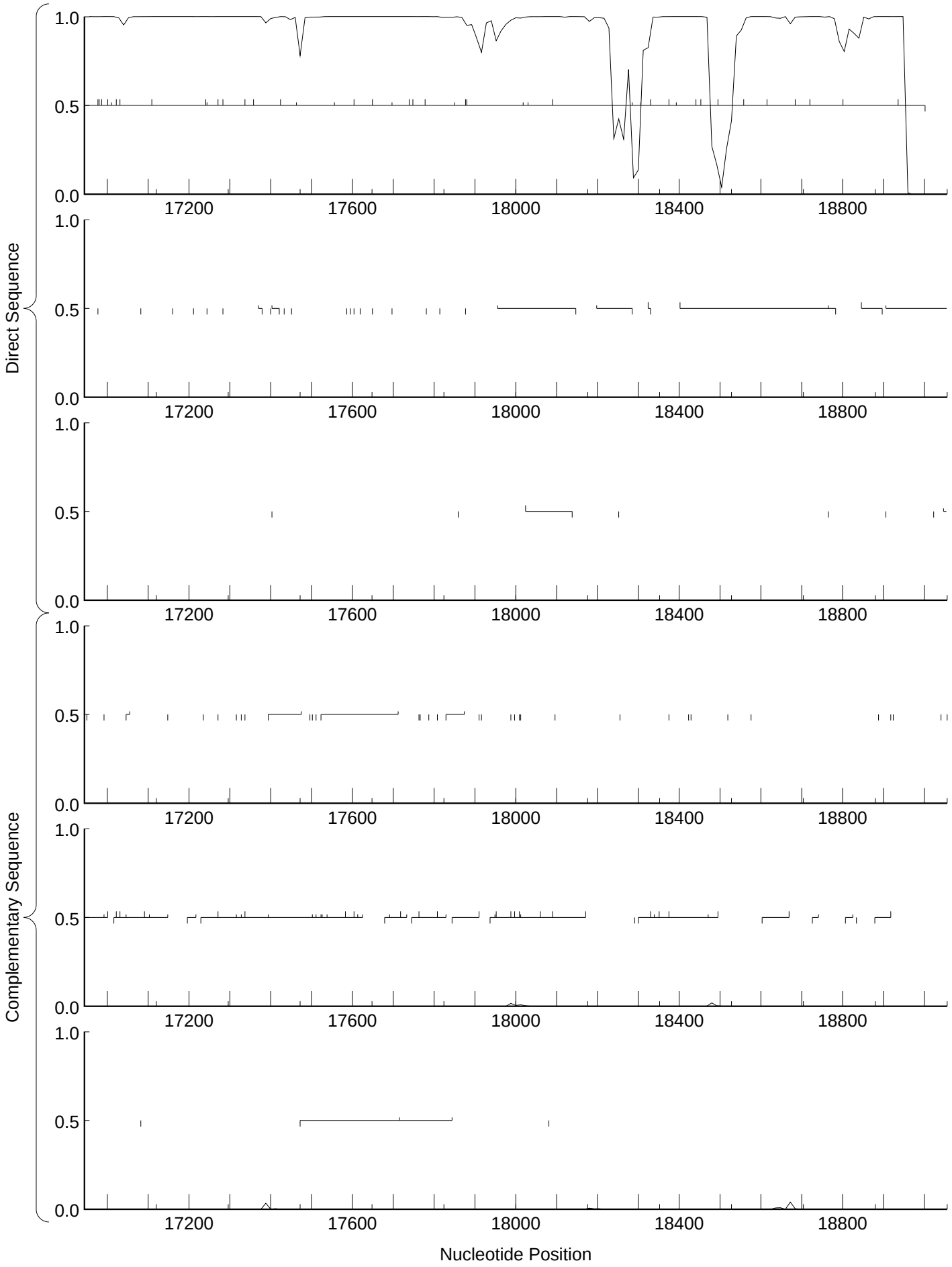
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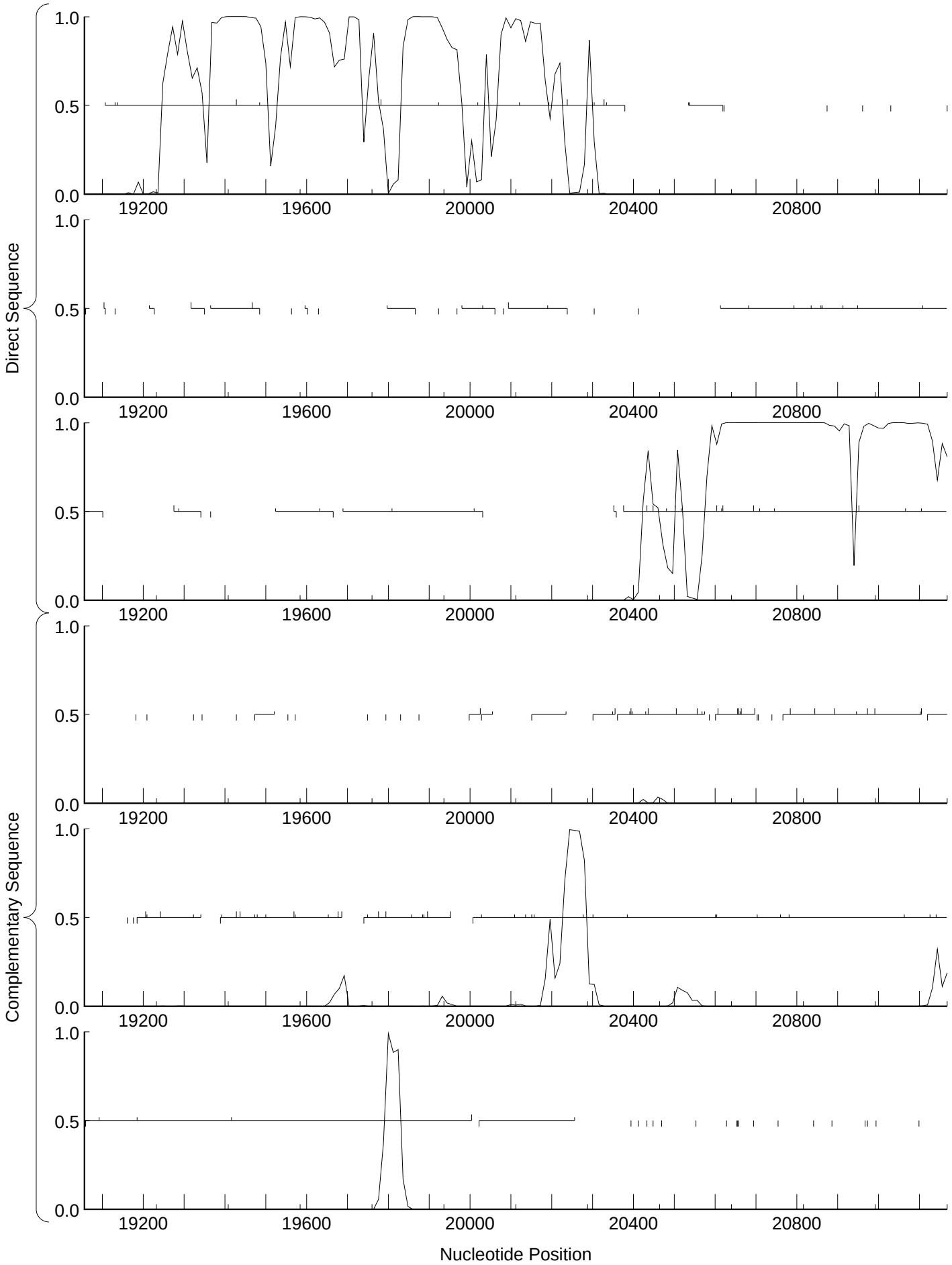
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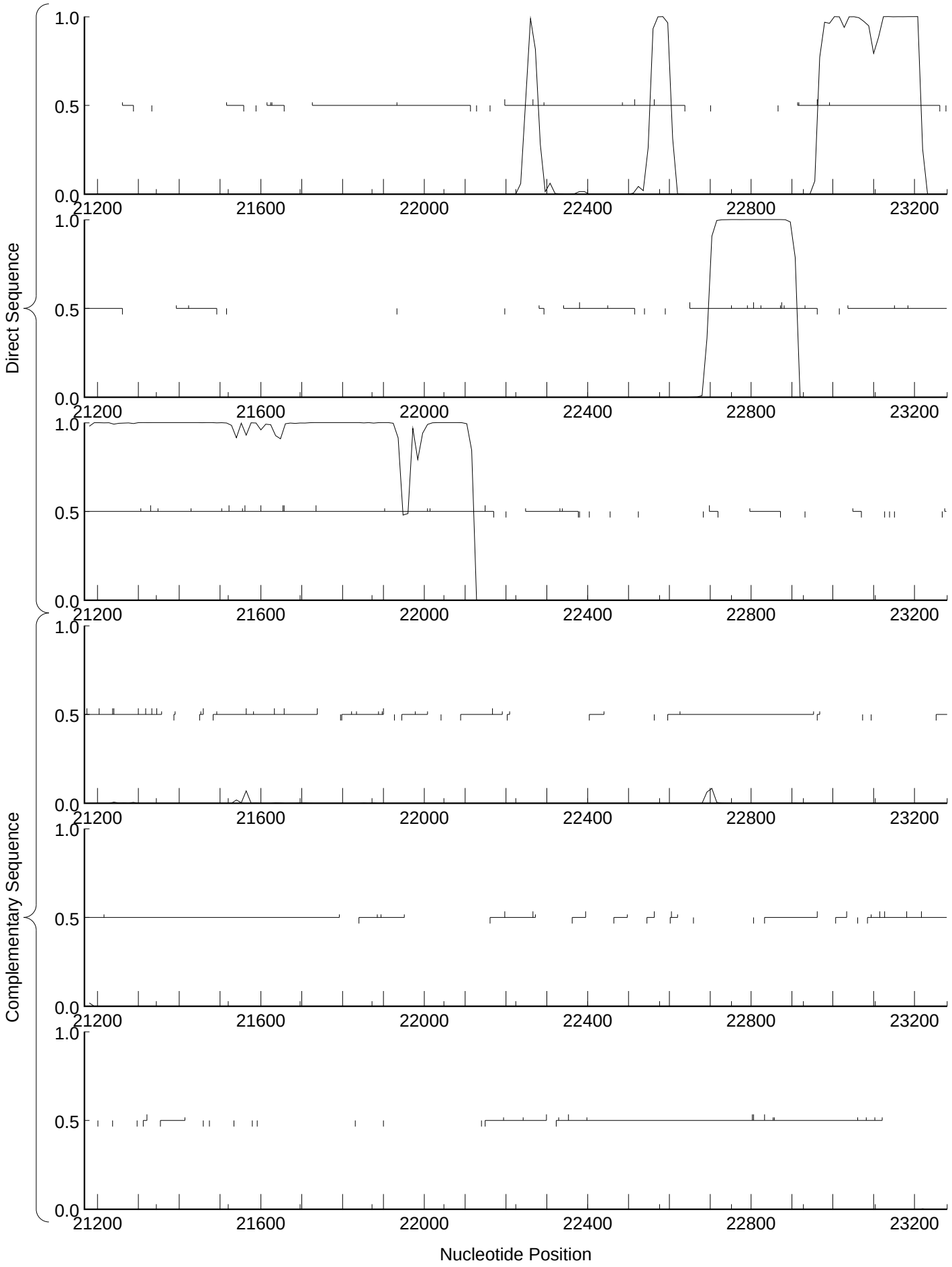
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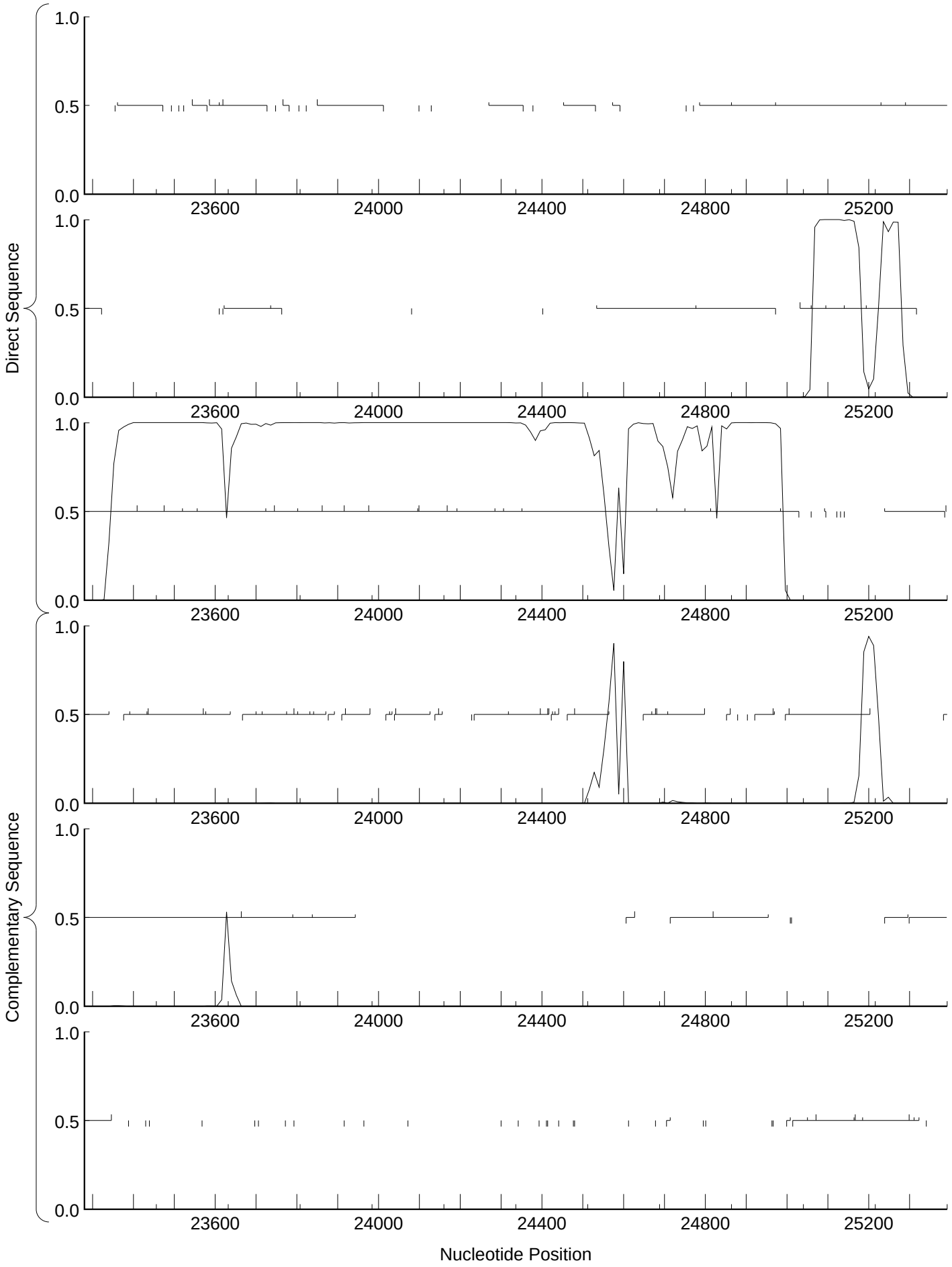
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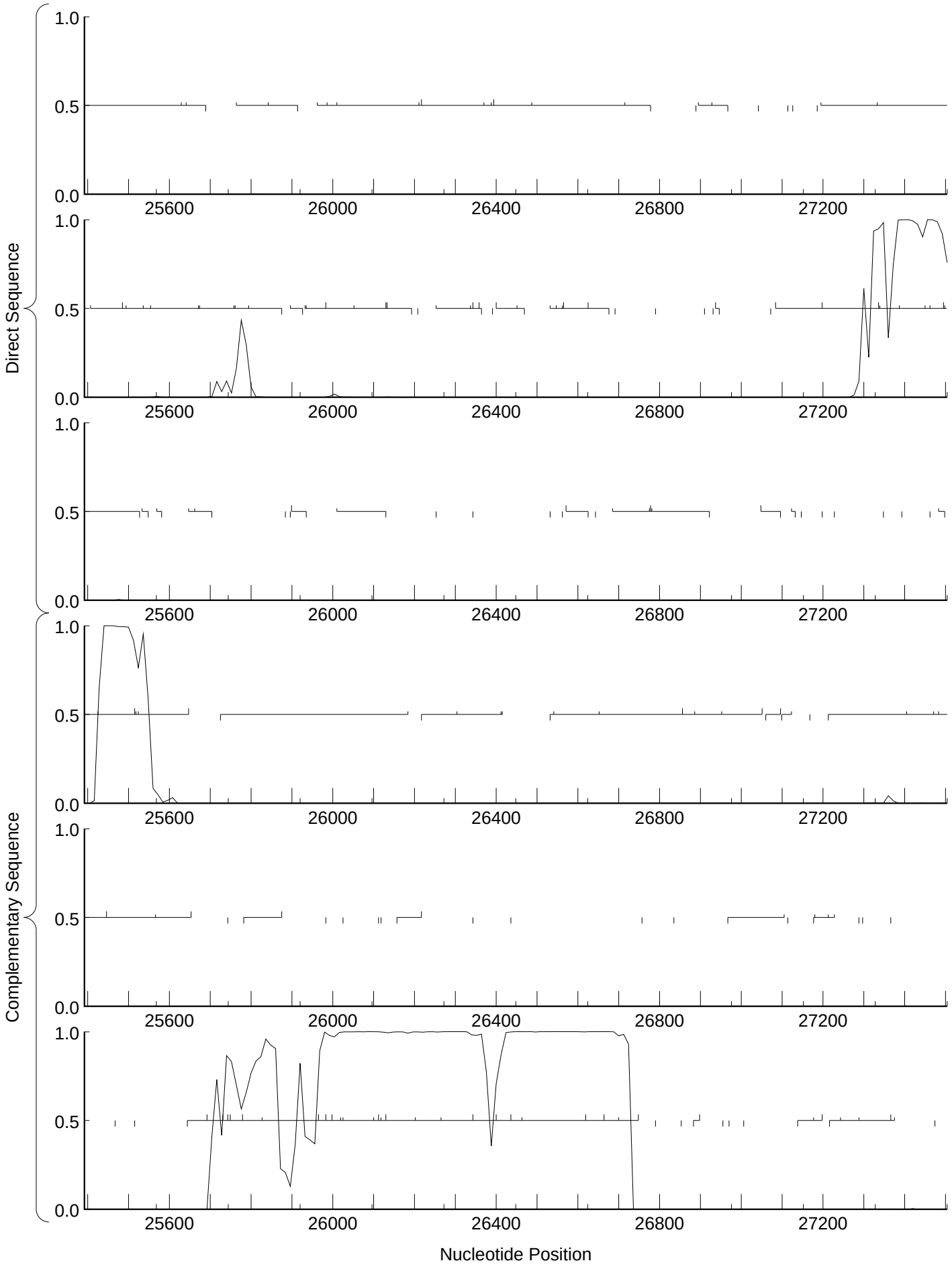
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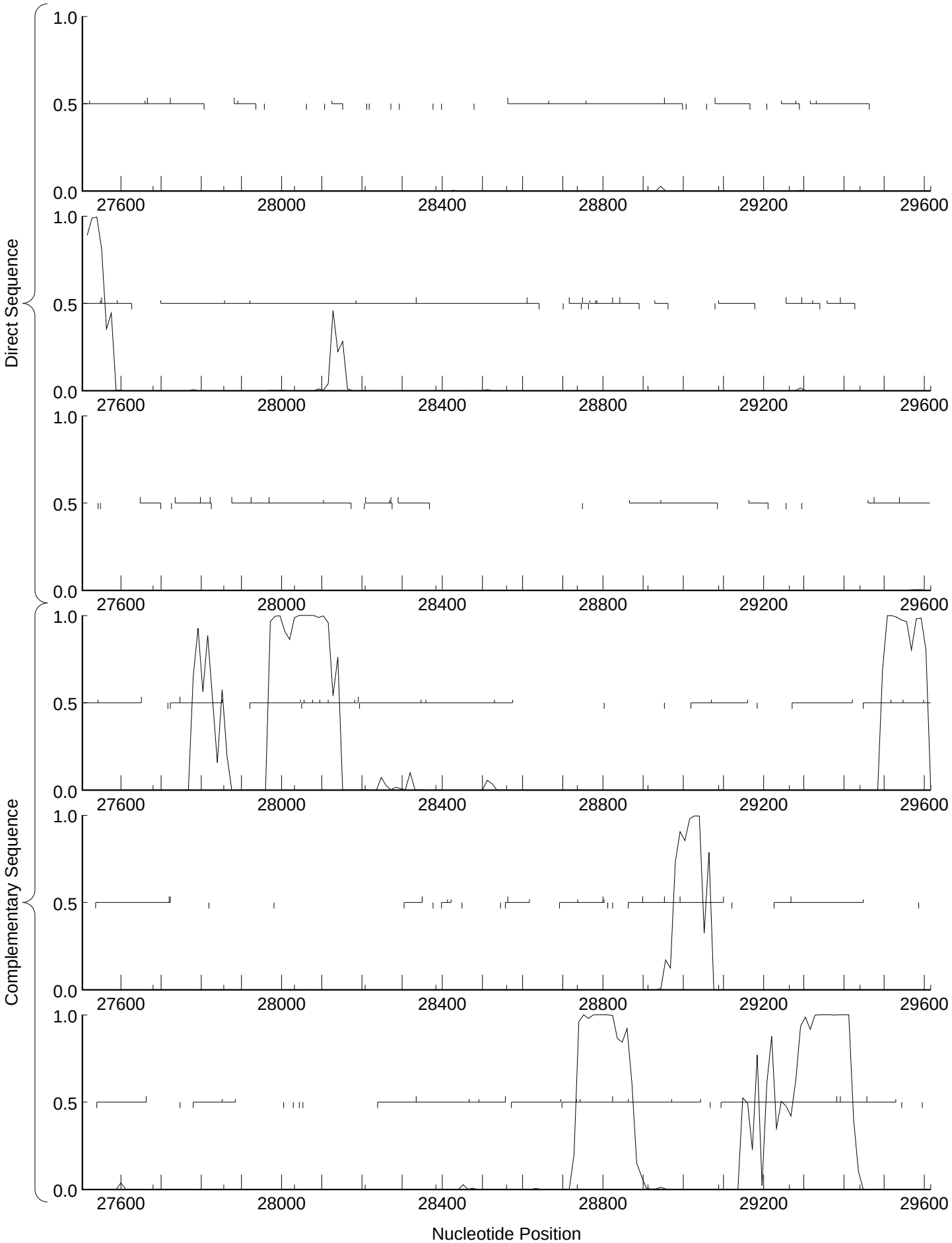
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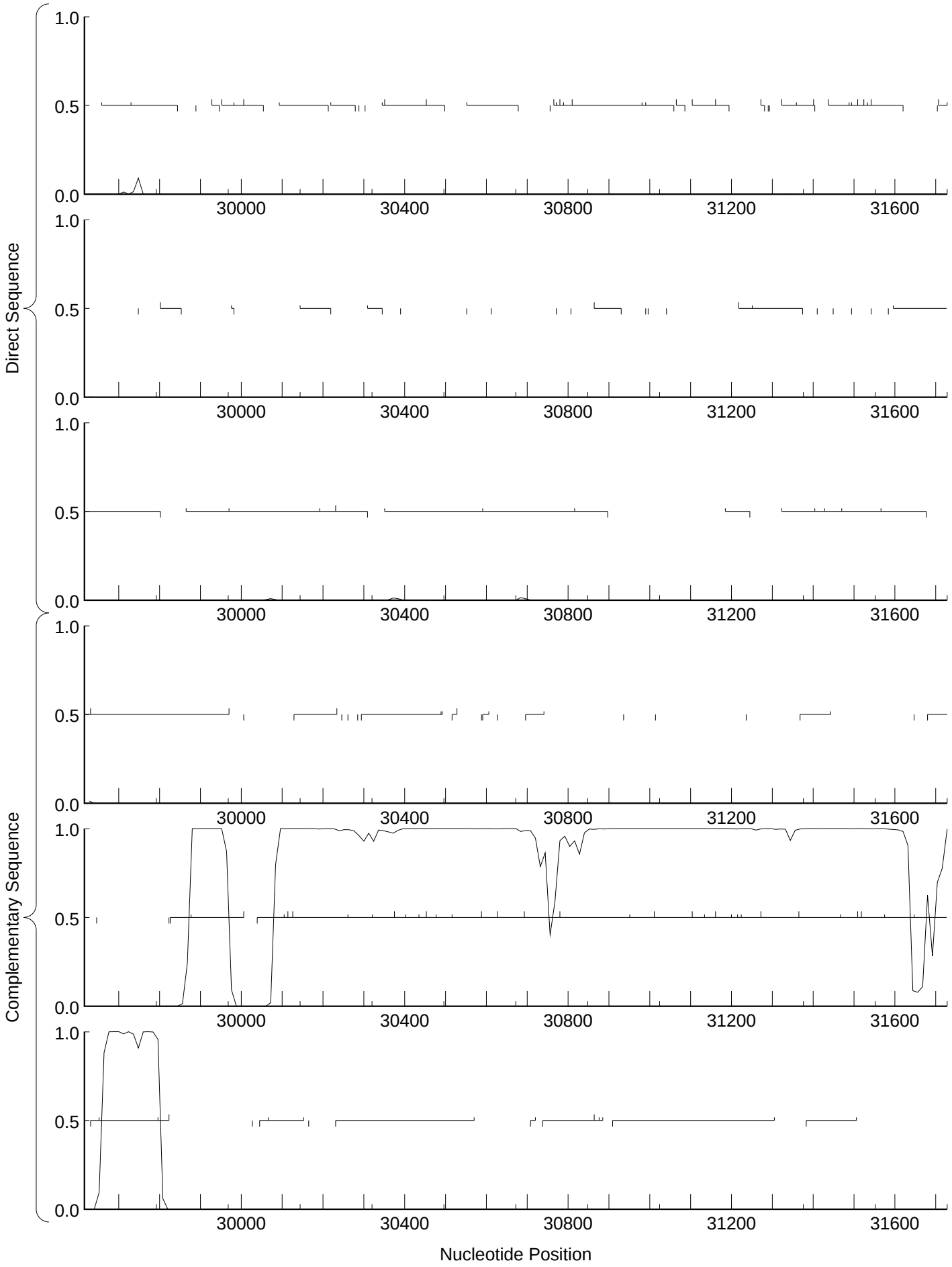
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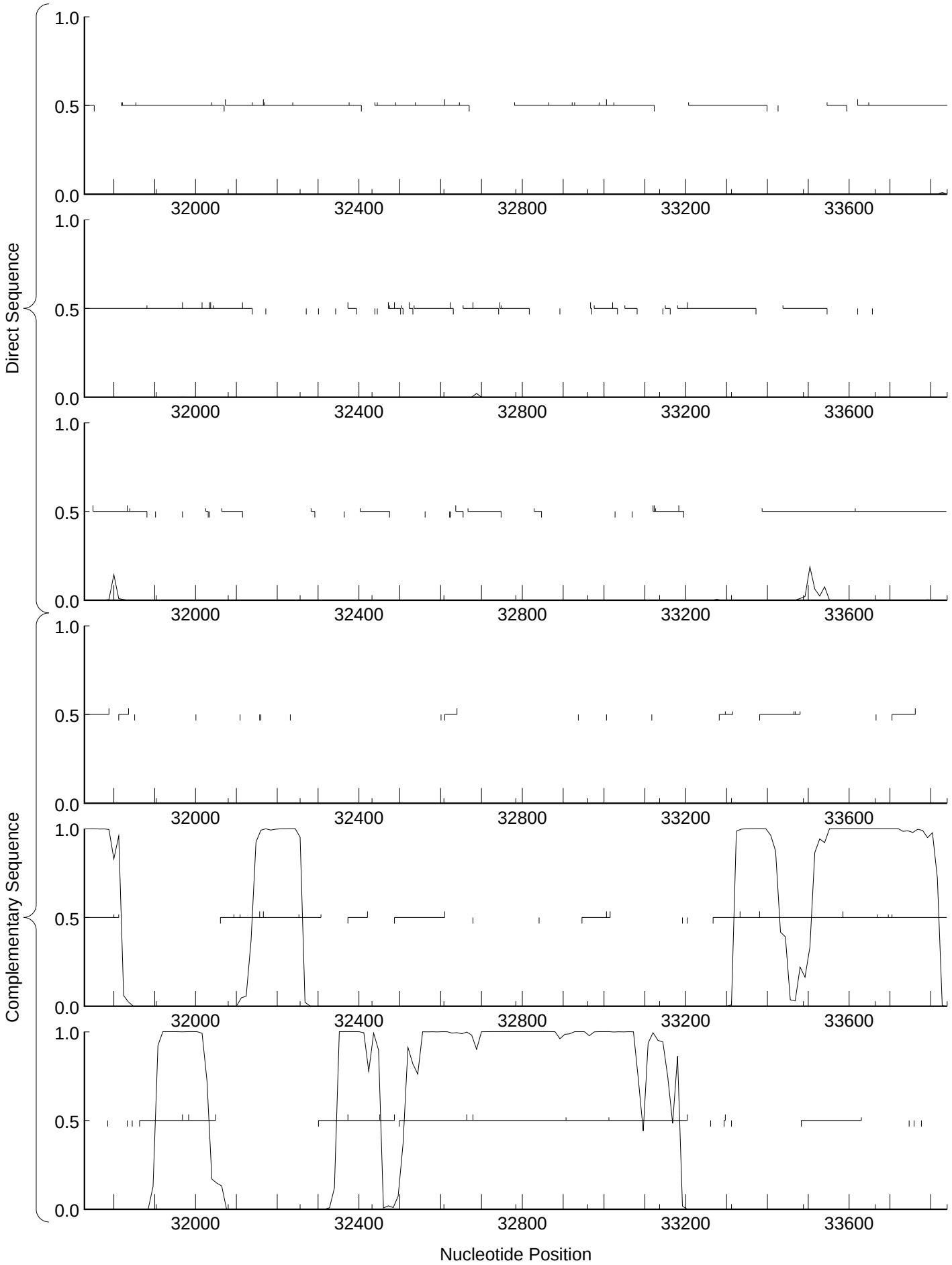
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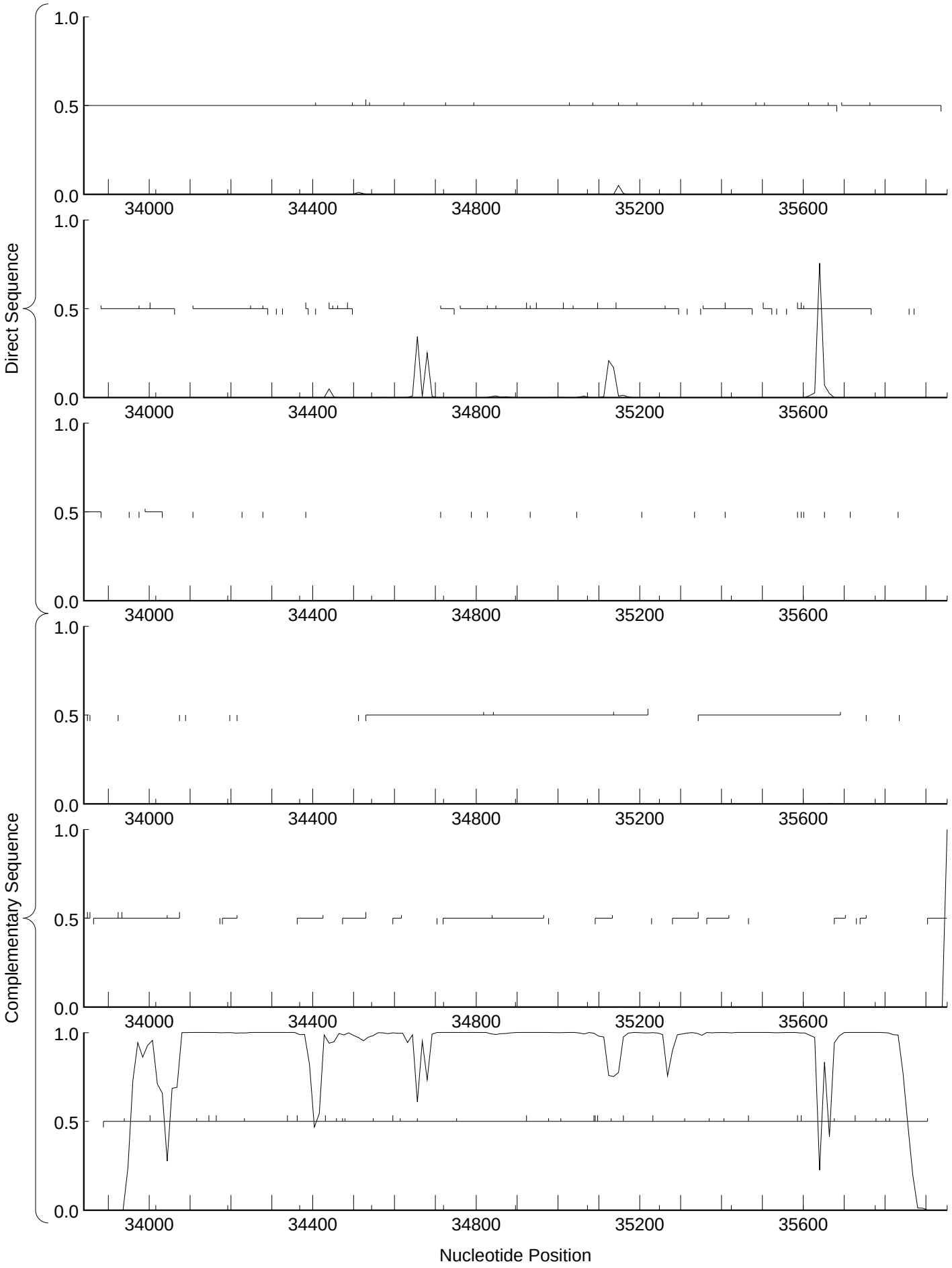
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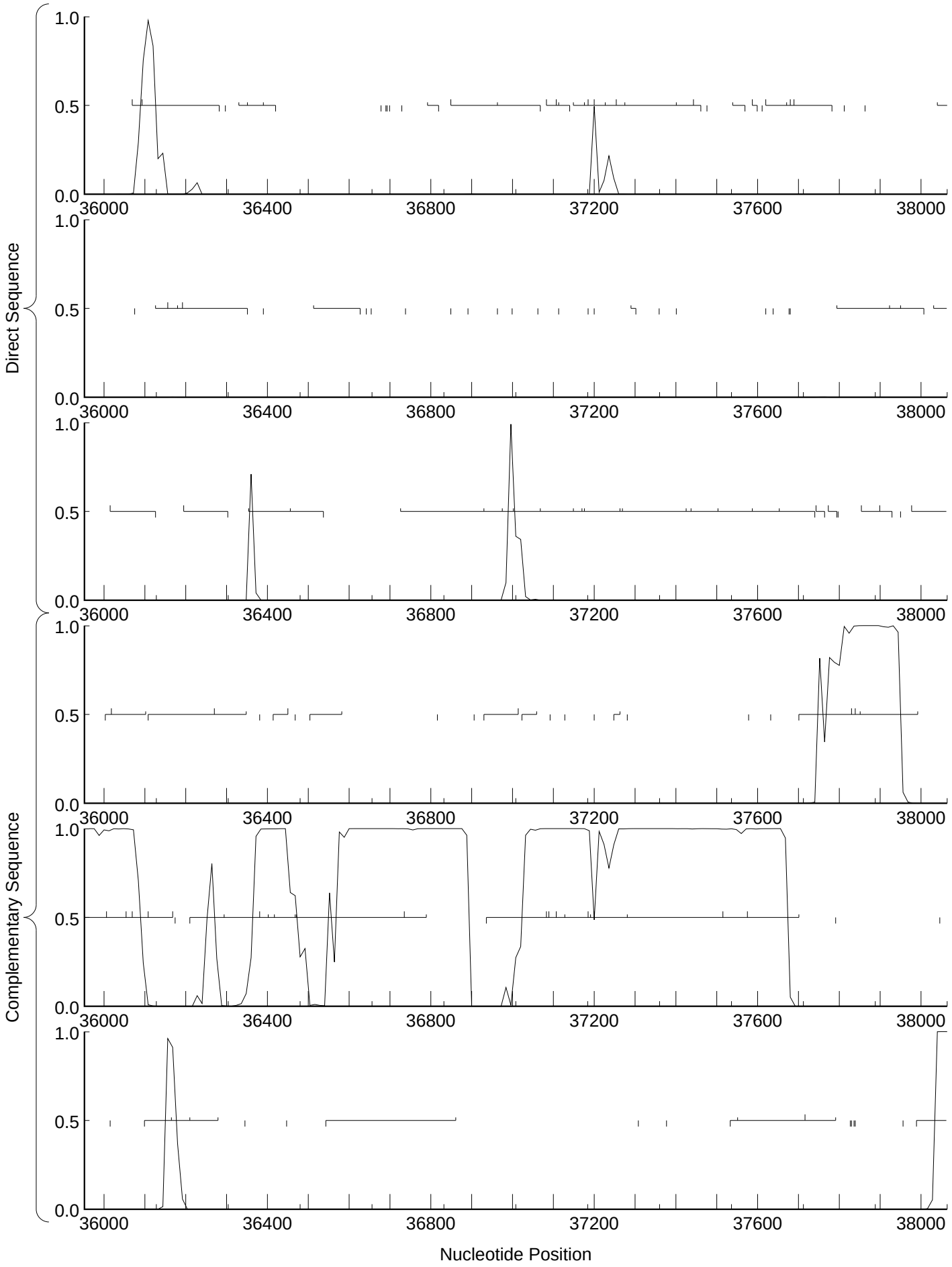
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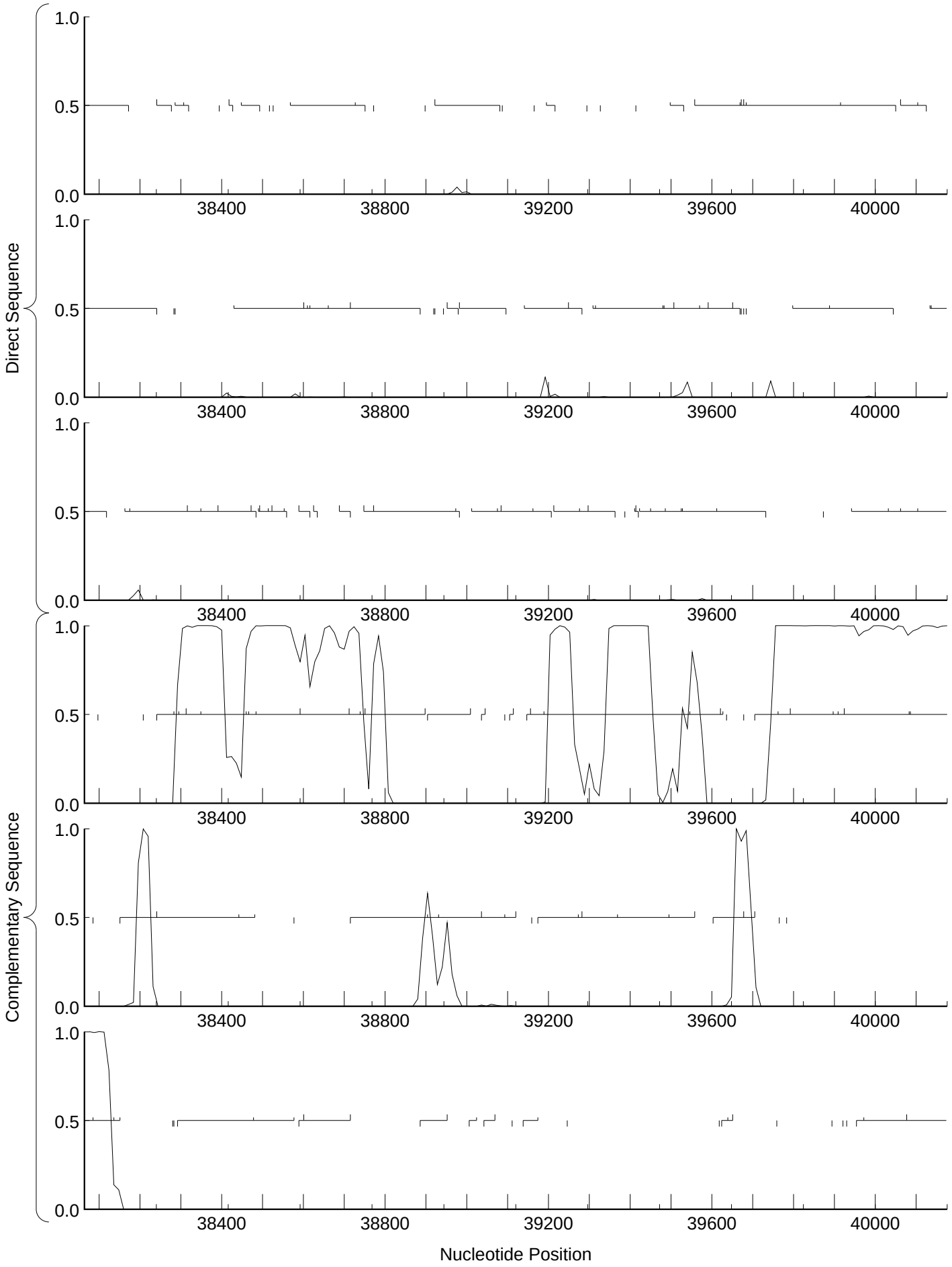
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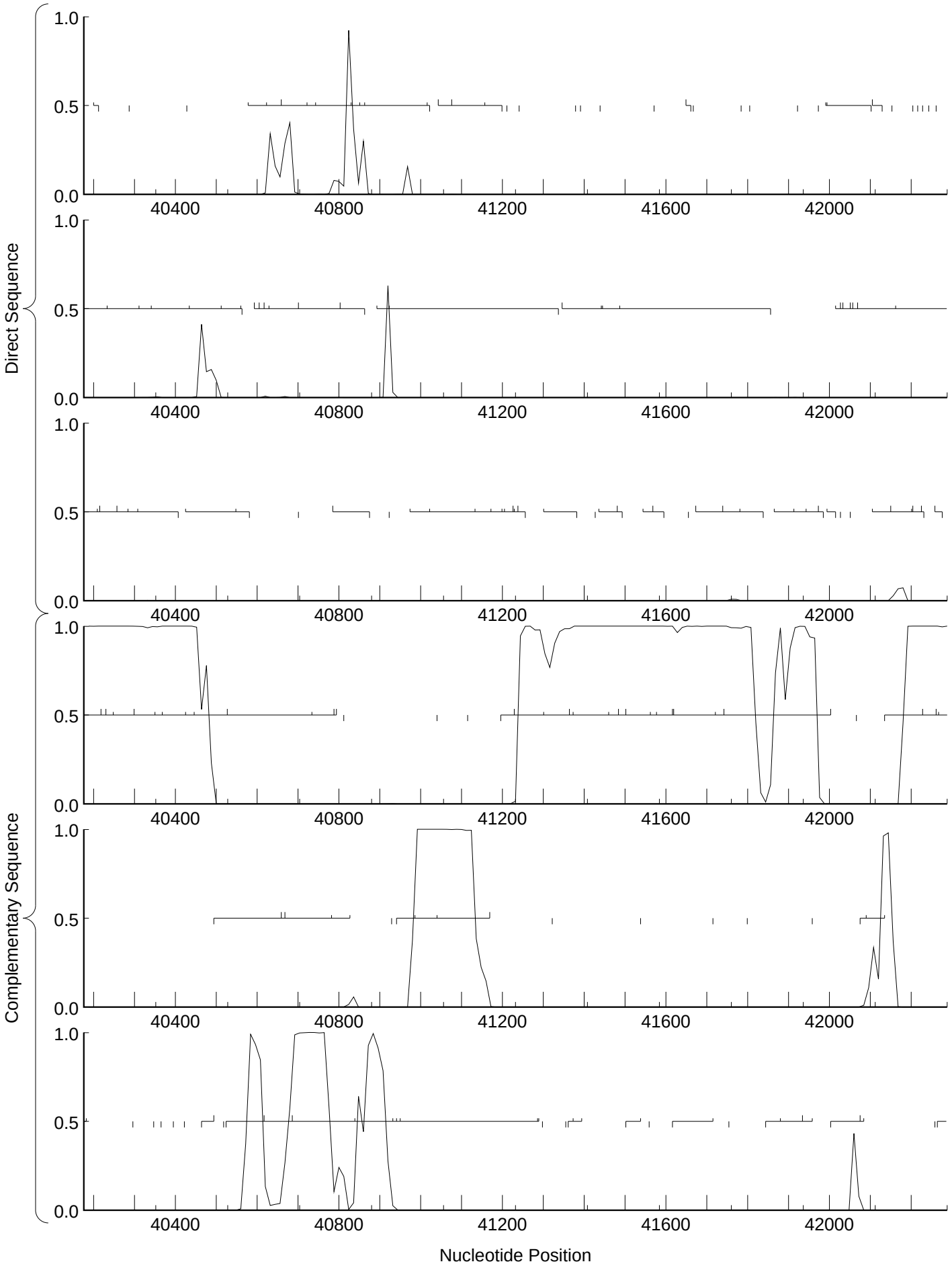
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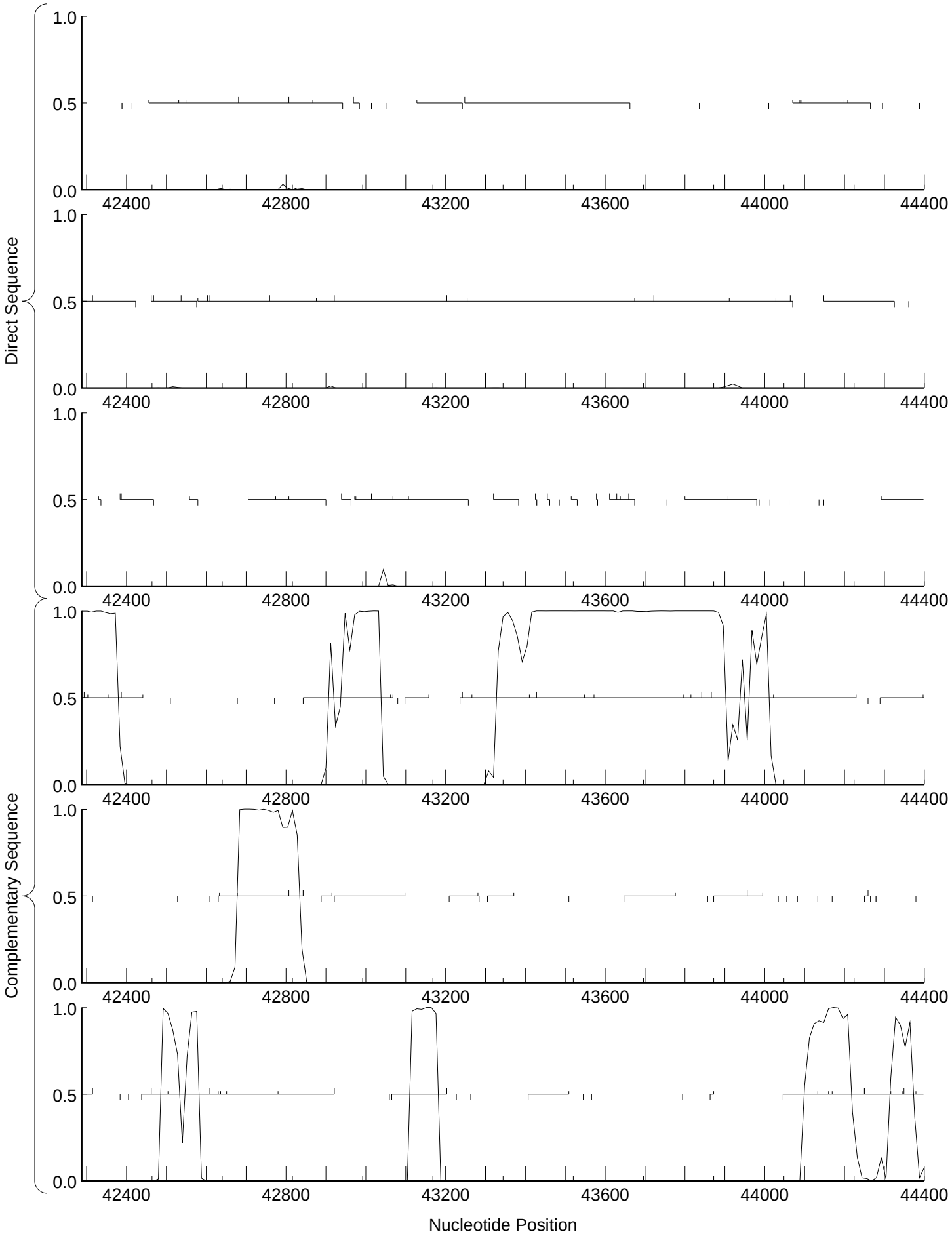
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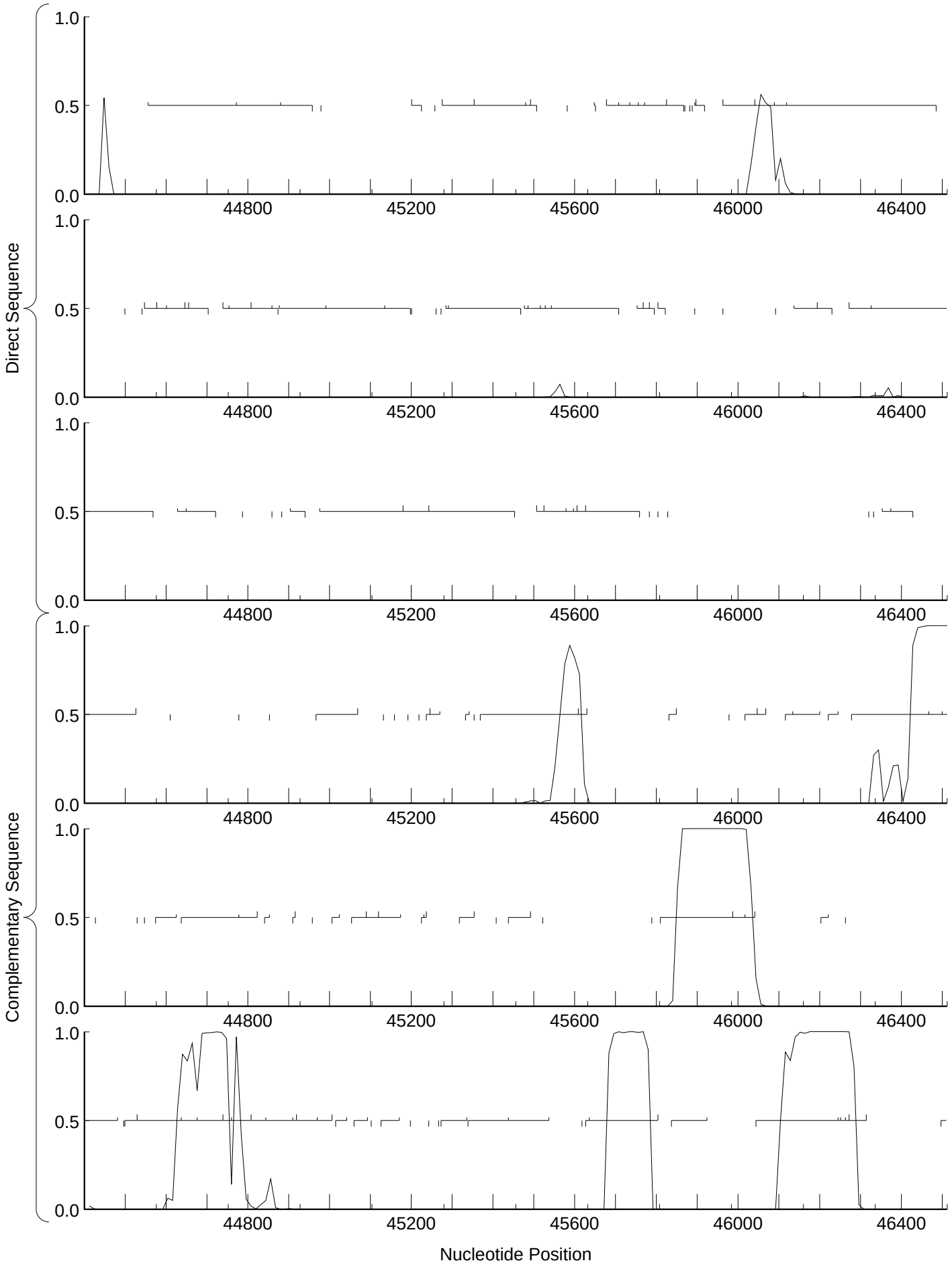
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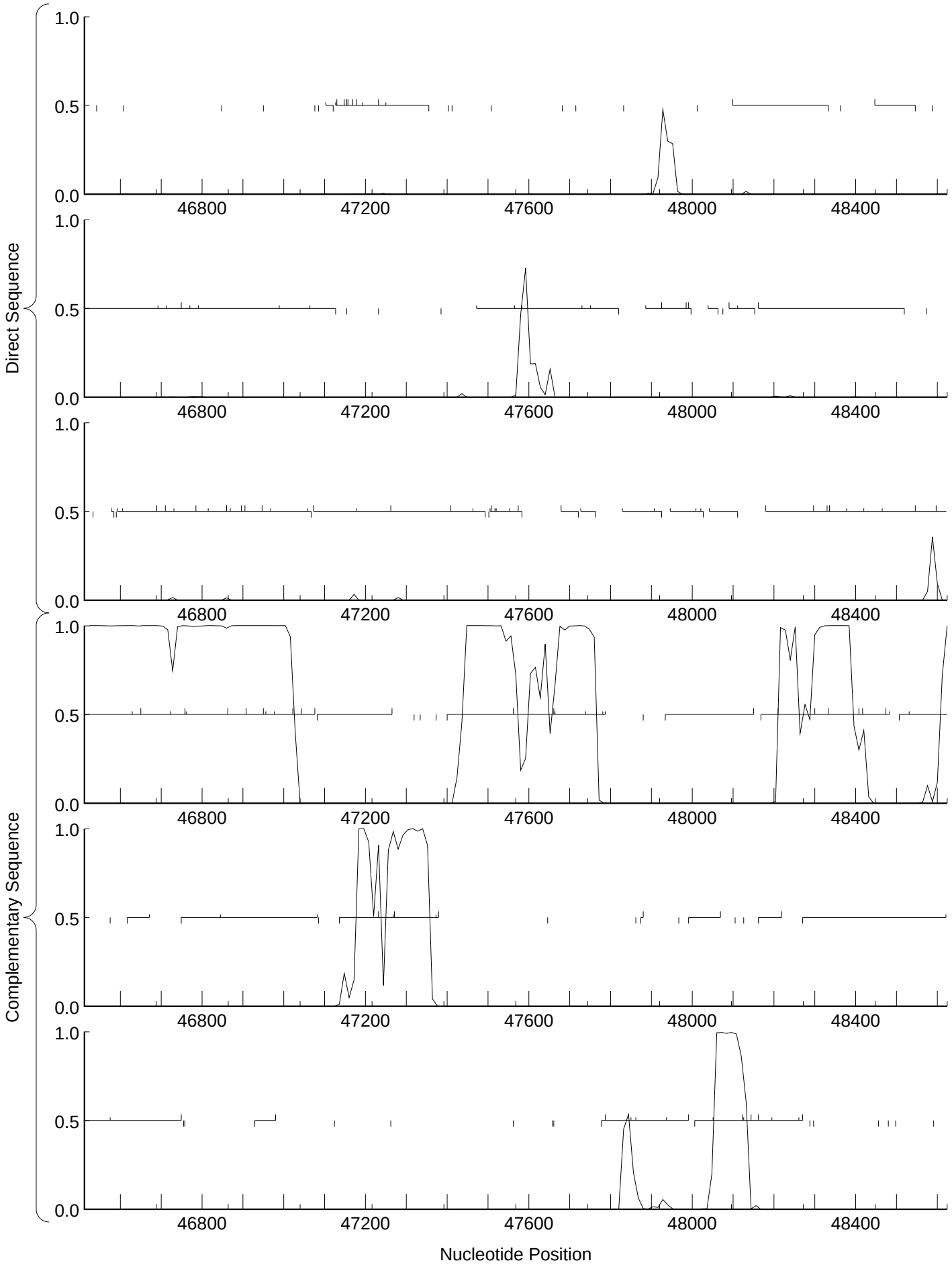
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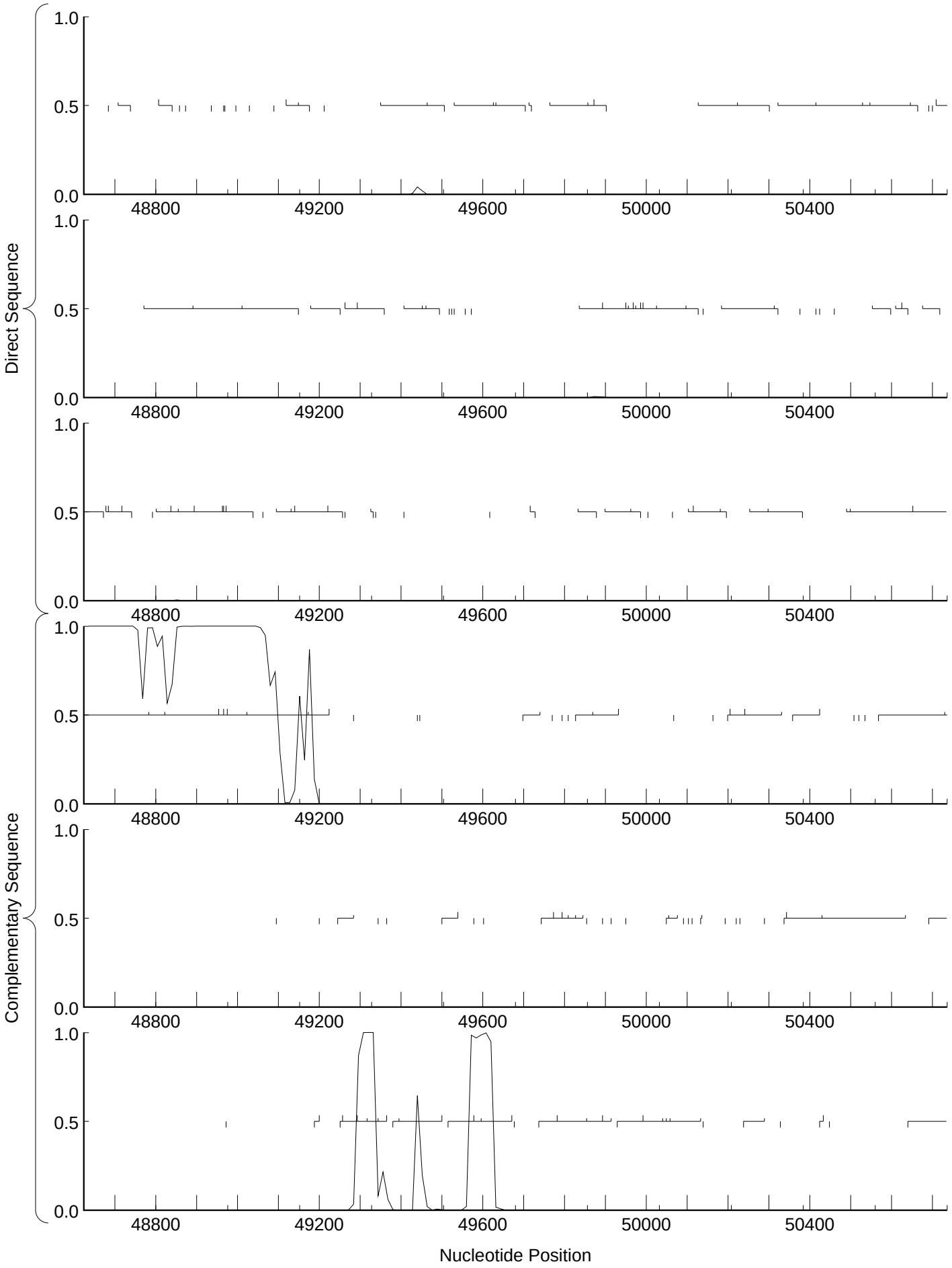
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complete sequence, 50878 bp including 10-base 3’ overhang (CGGGTGGTAA), Cluster A3, Order 4, Window 96, Step 12, 25/26



complete sequence, 50878 bp including 10-base 3’ overhang (CGGGTGGTAA), Cluster A3, Order 4, Window 96, Step 12, 26/26

