GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Perplexer complete sequence, 51264 bp including 10-base 3’ overhang (CG
Analysis Date    :  3/28/24 at 5:35:01
Pages            :  26
Sequence Length  :  51264 bp
GC Content       :   63.71%

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

PS-Version       :  1.2

GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
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Matrix notes & comments

Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

MATRIX INFORMATION

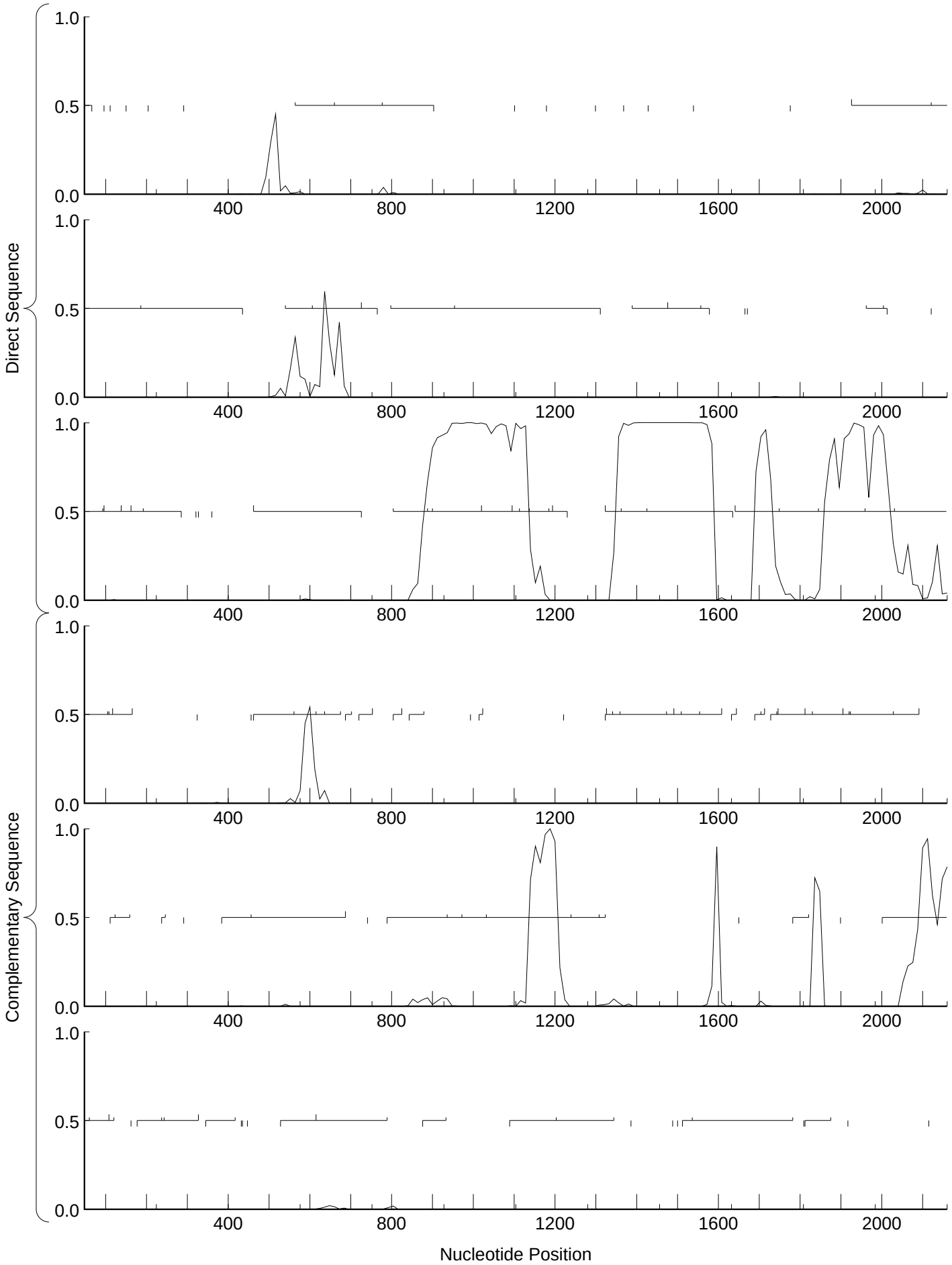
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Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

Order      :  4
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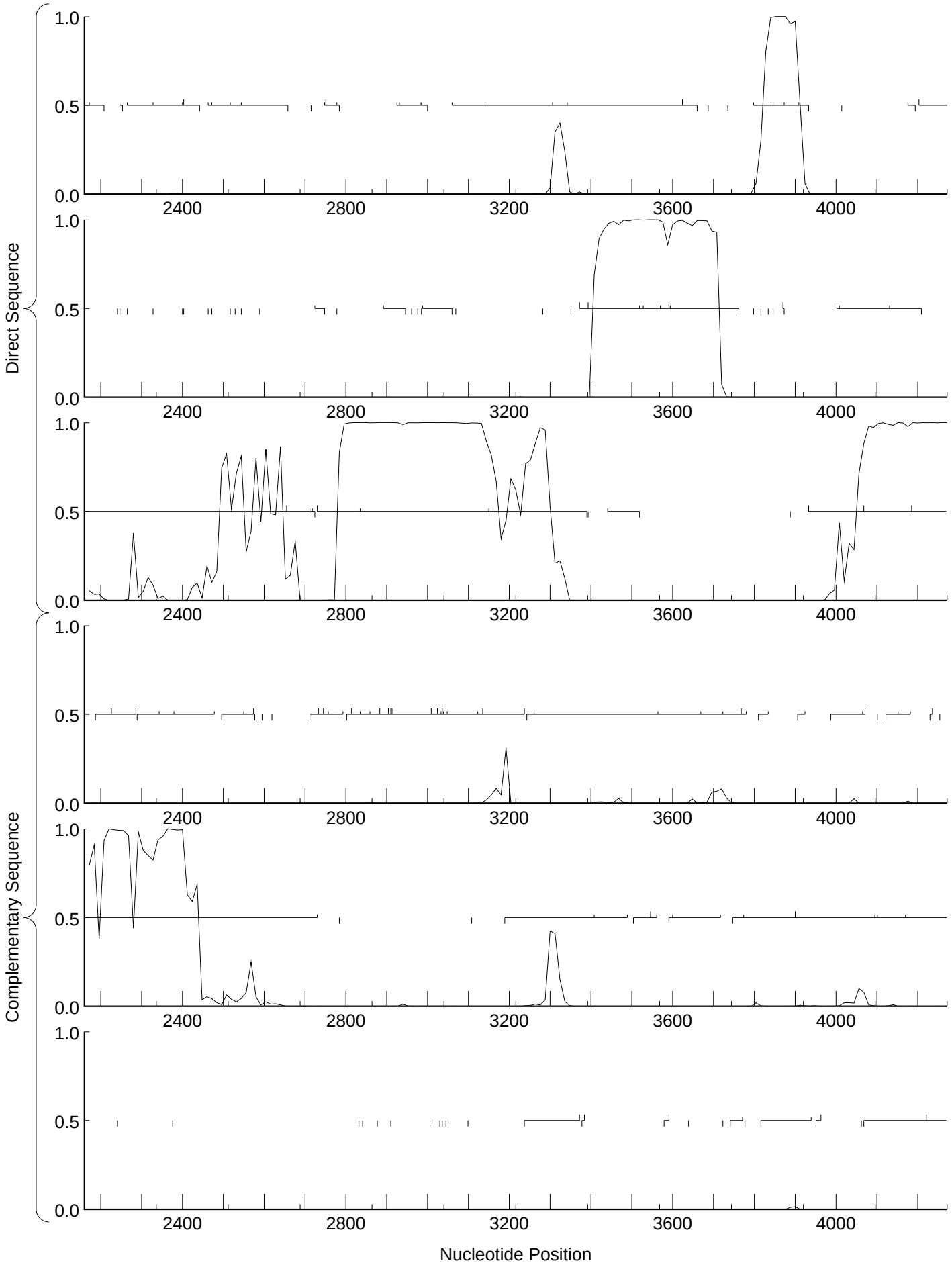

Send questions / comments to:
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Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

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

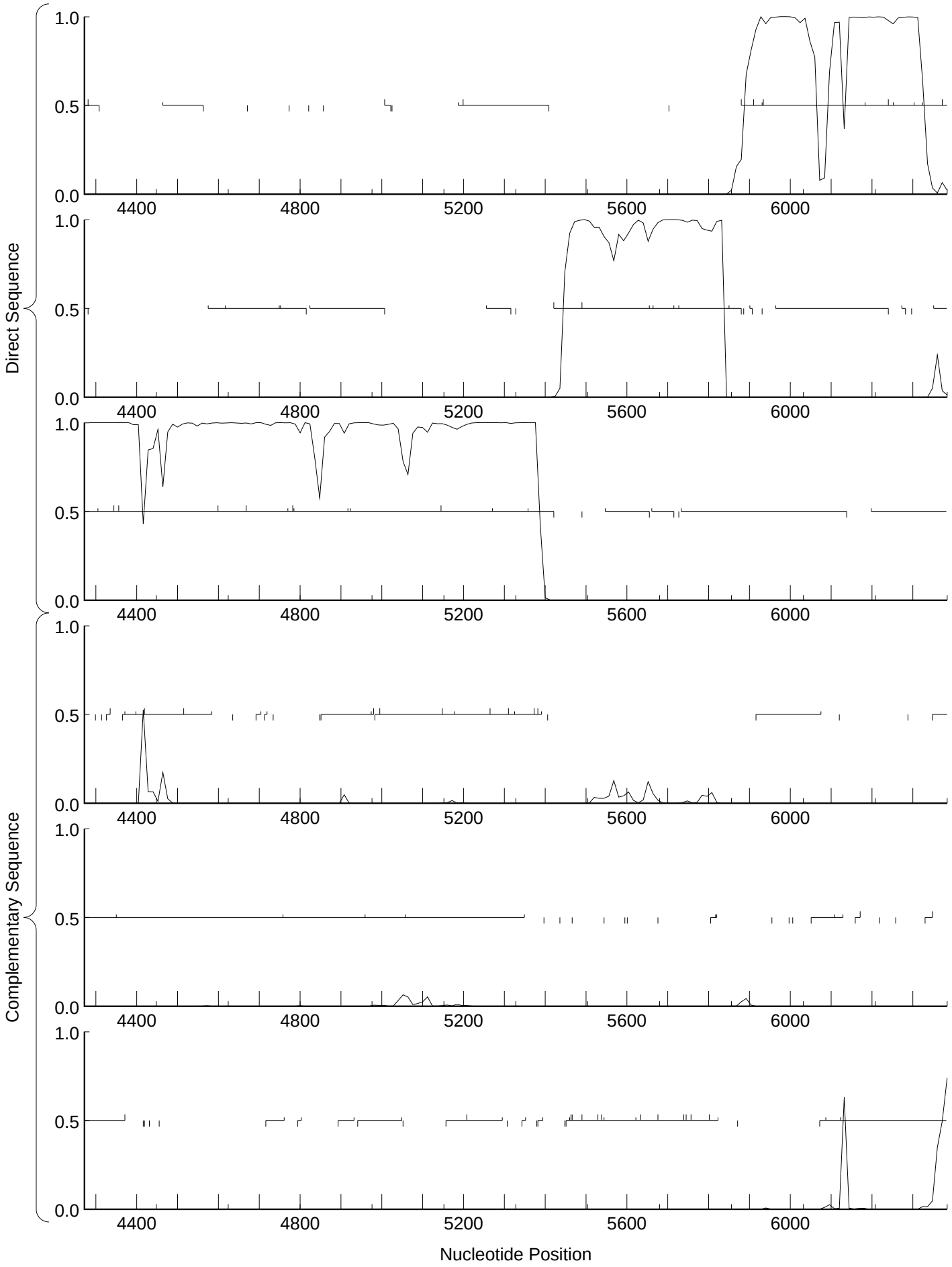
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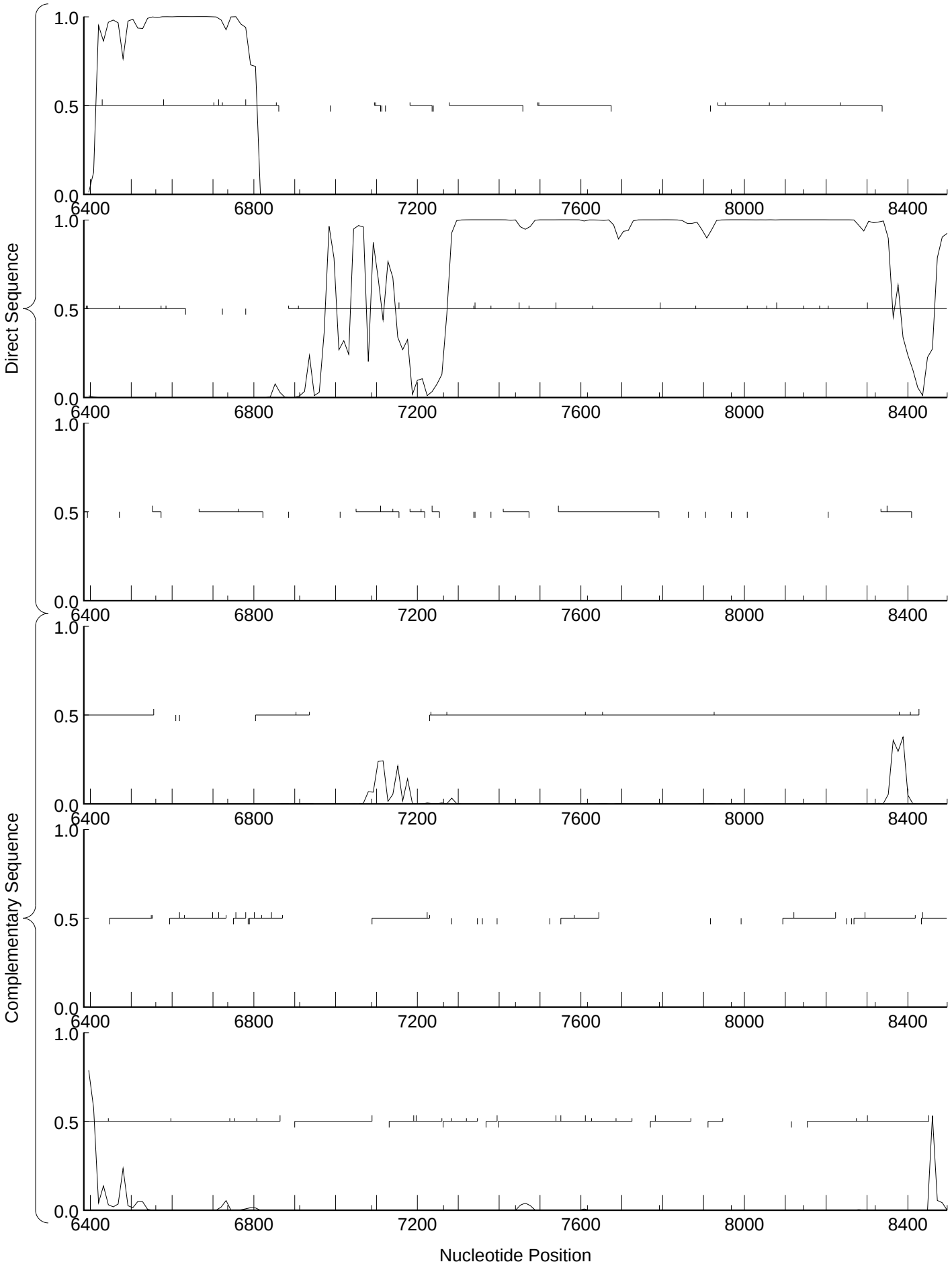
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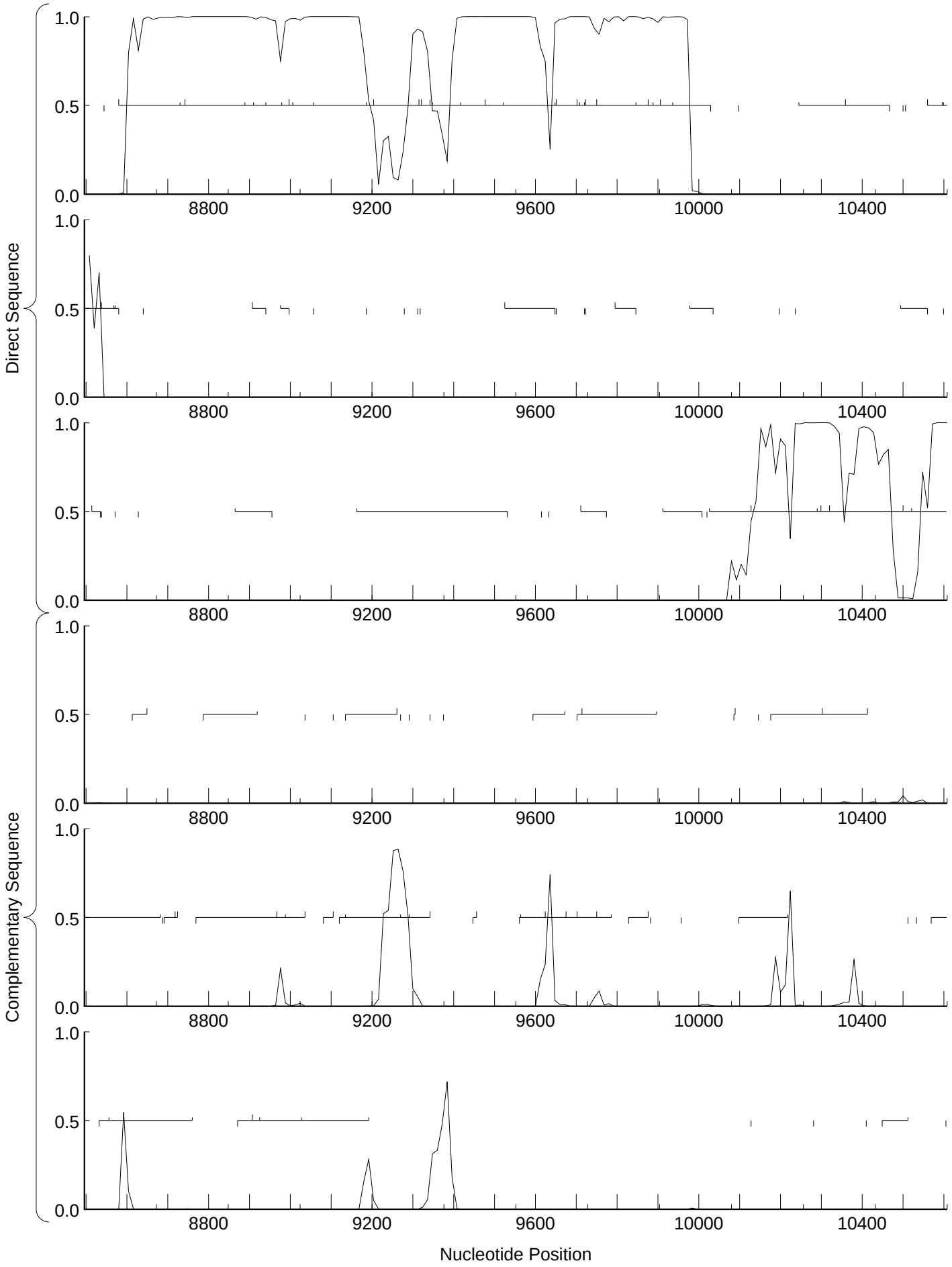
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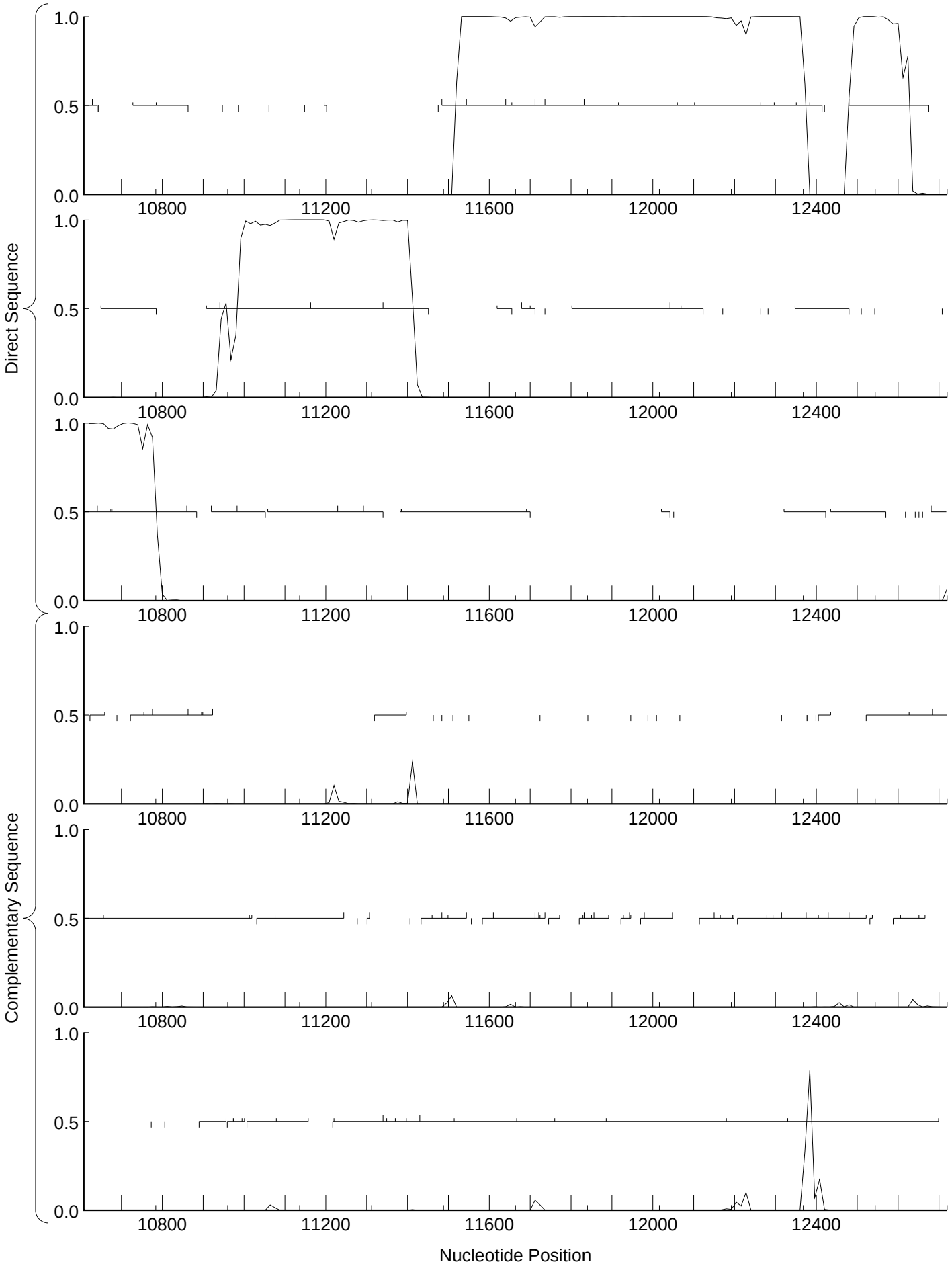
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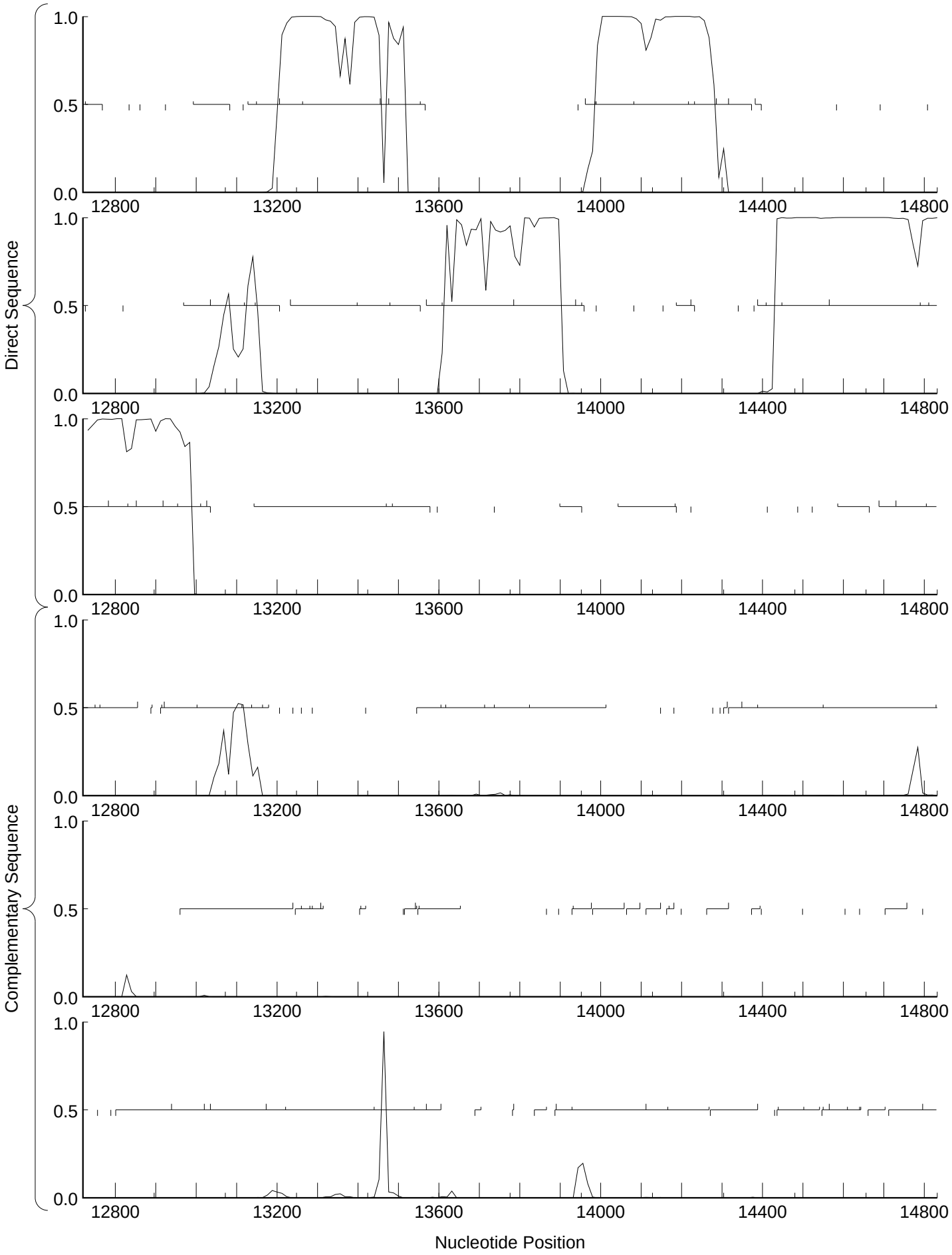
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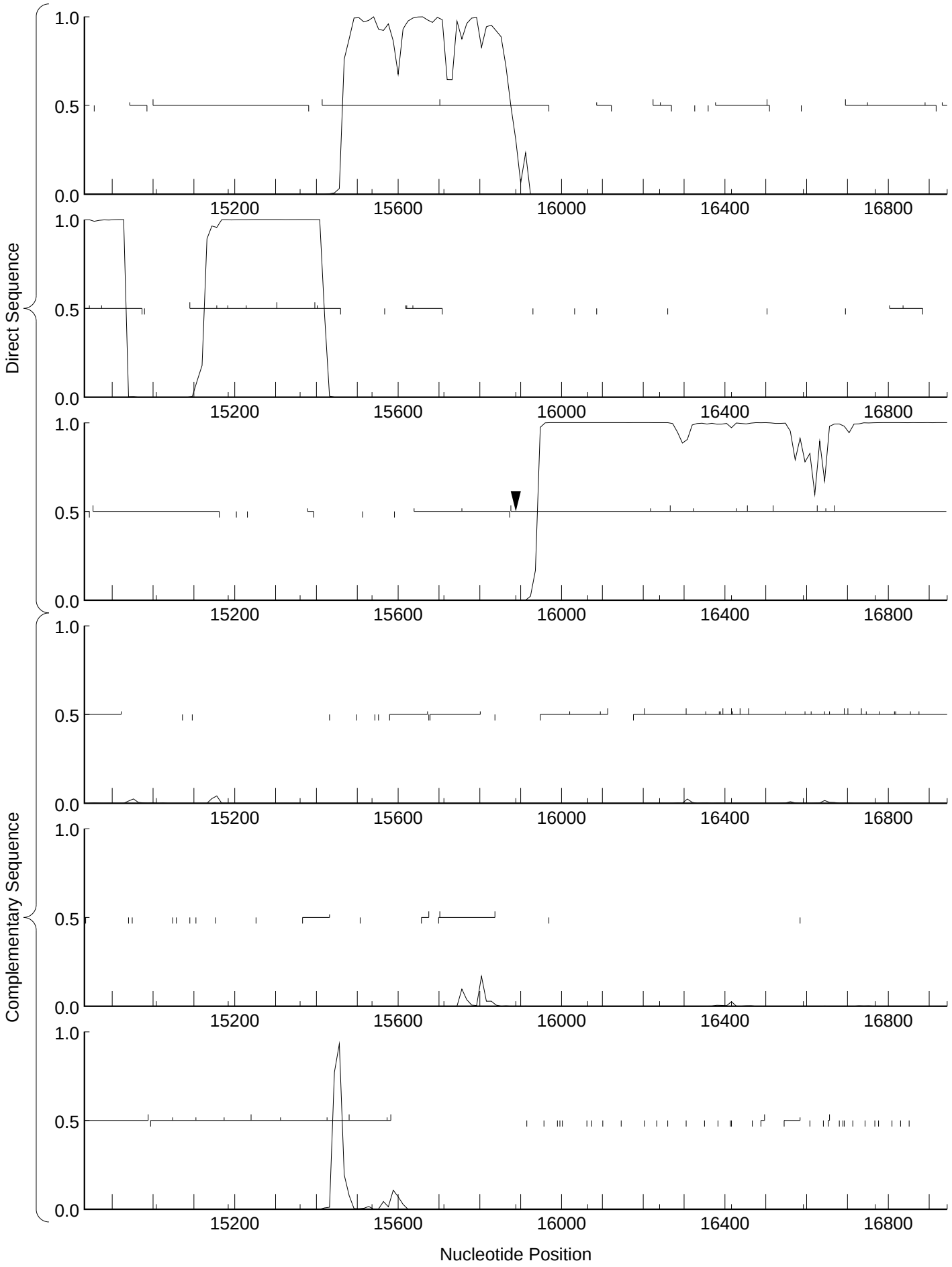
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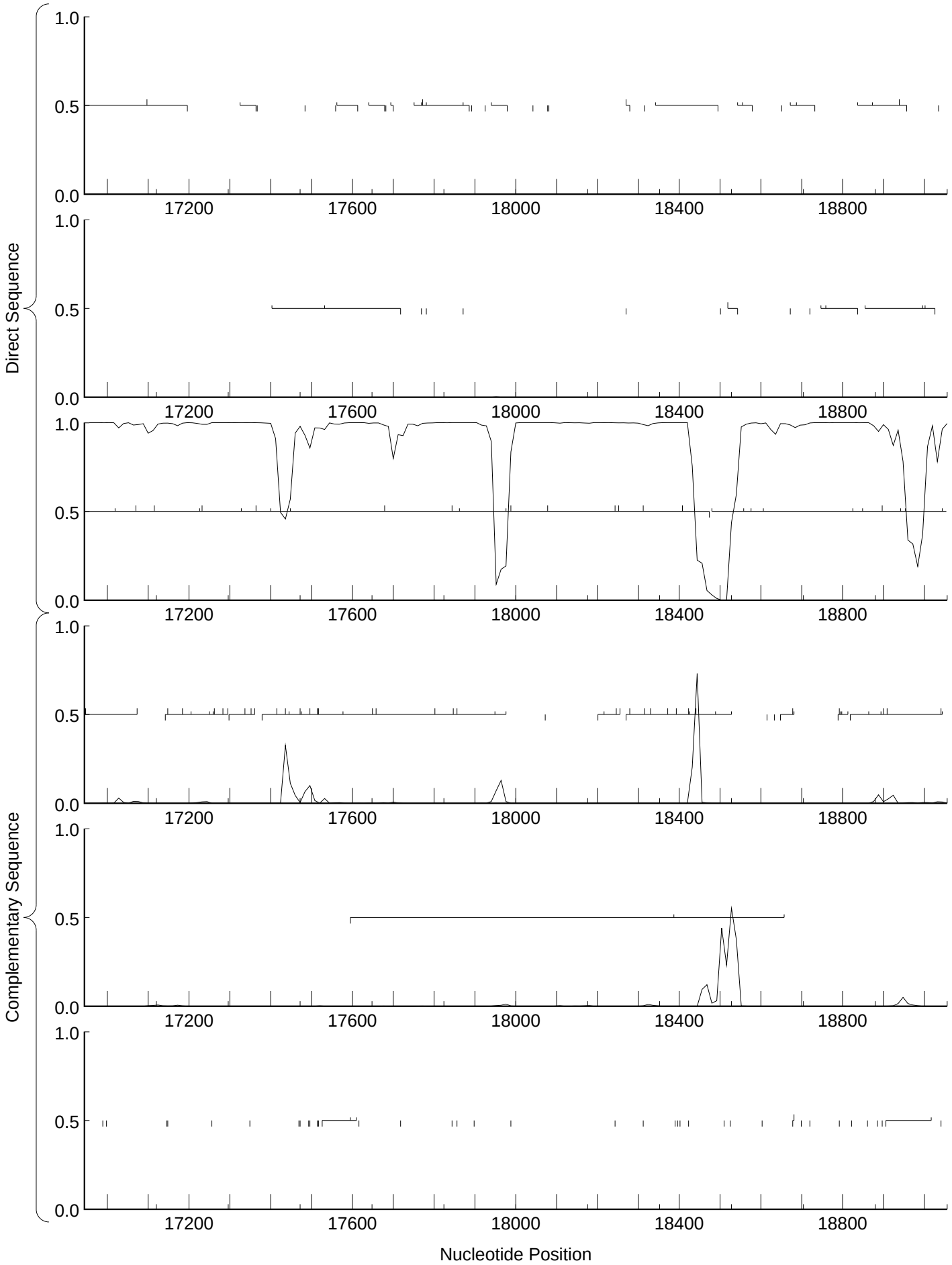
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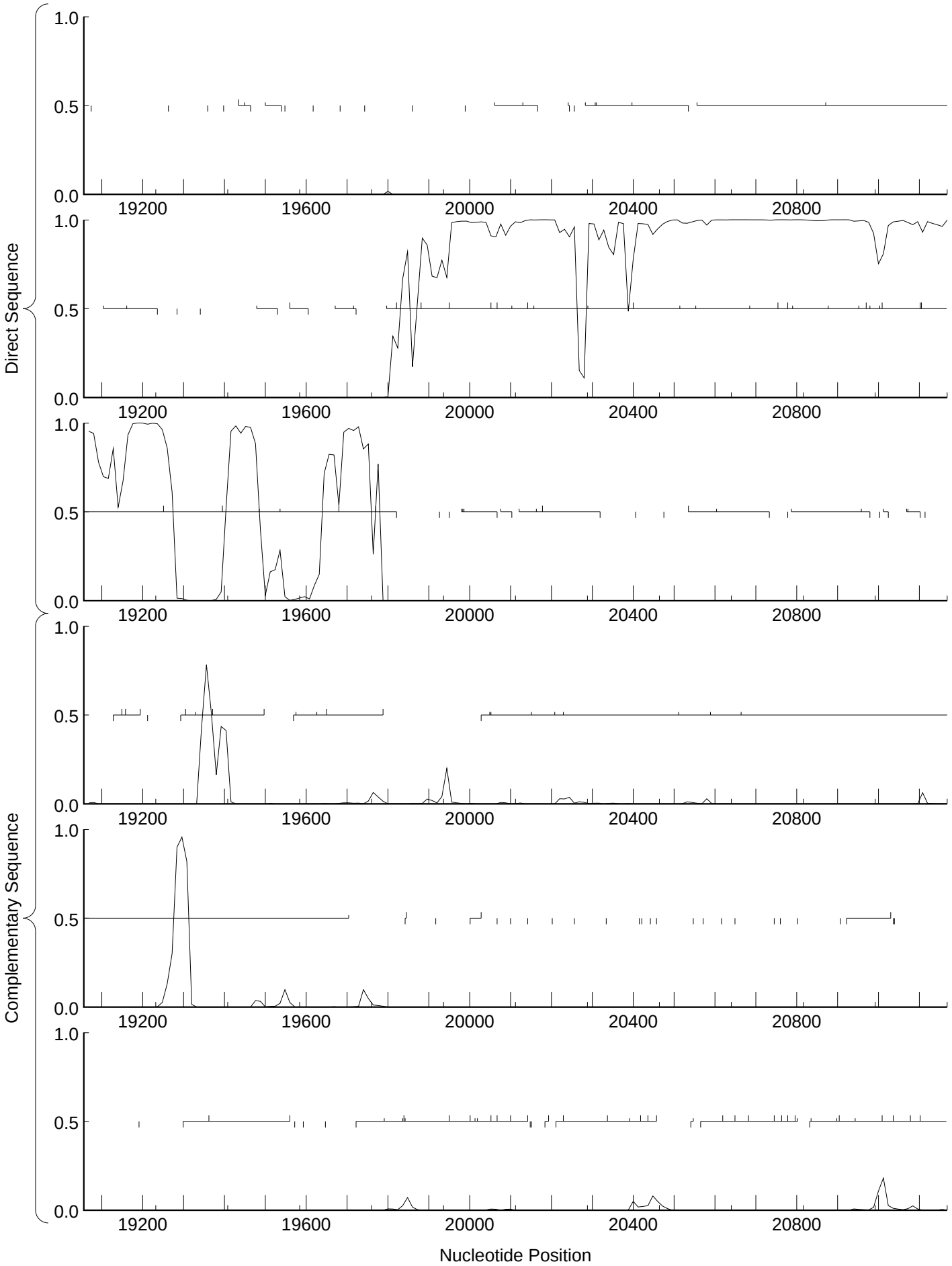
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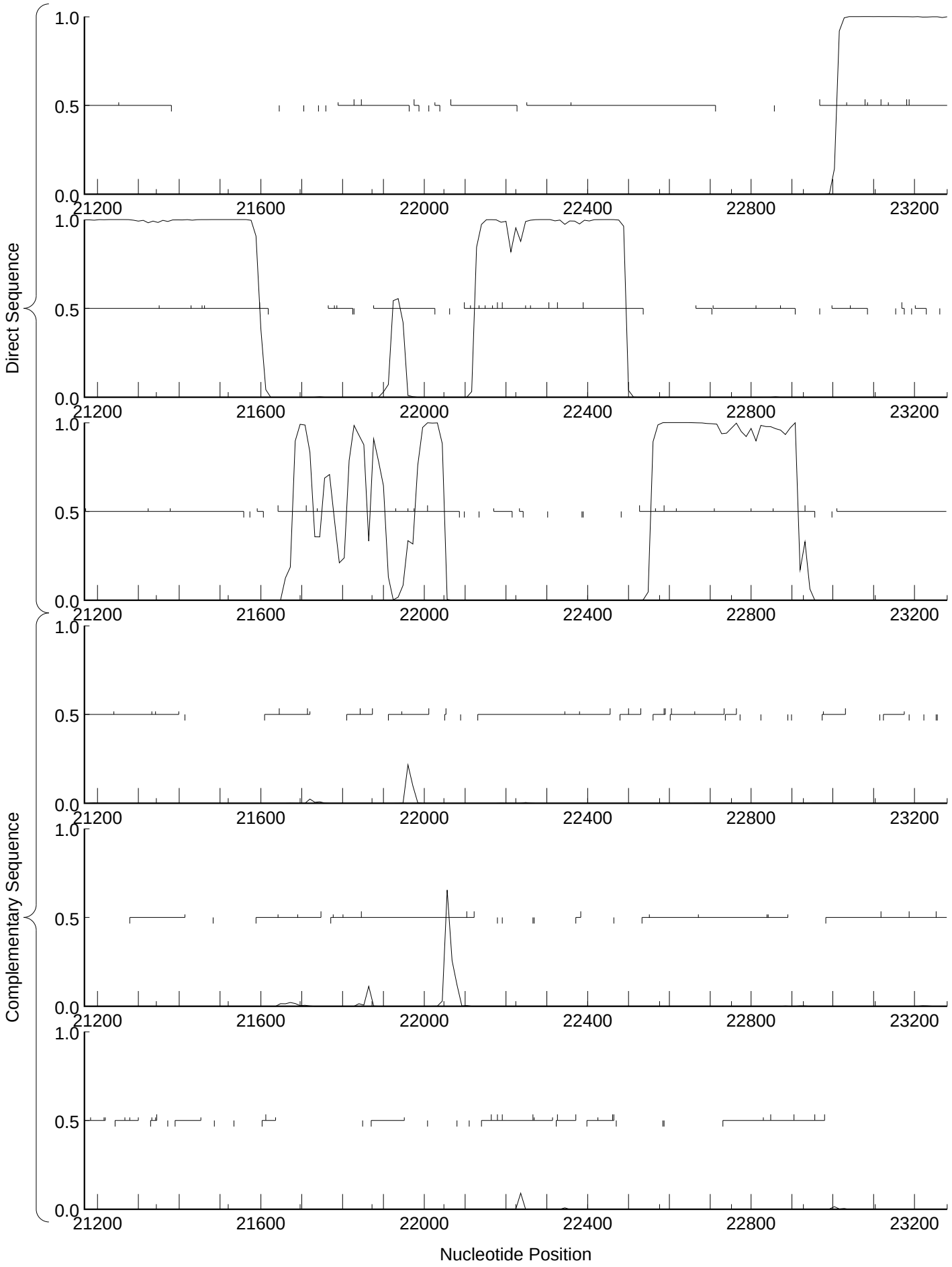
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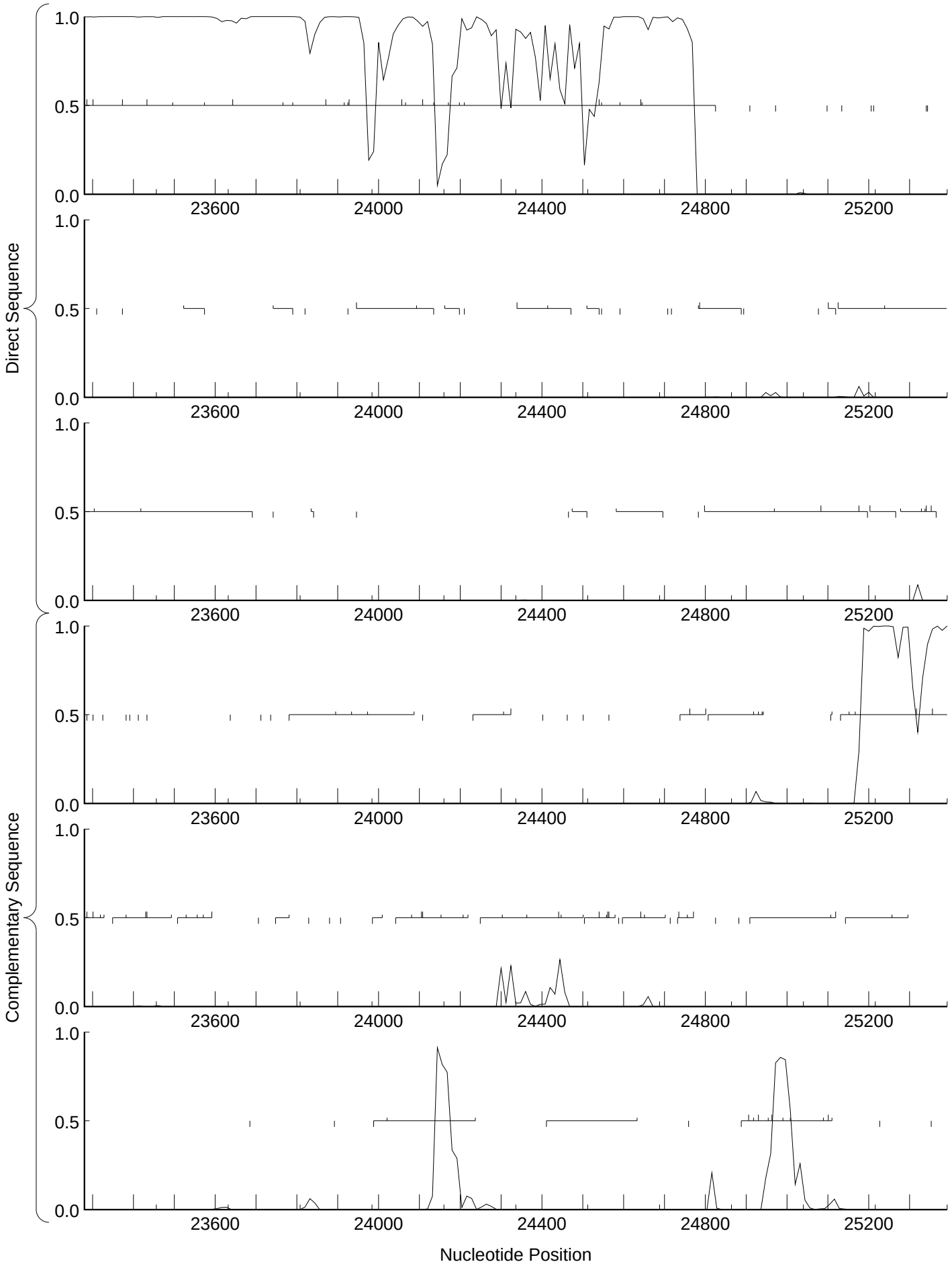
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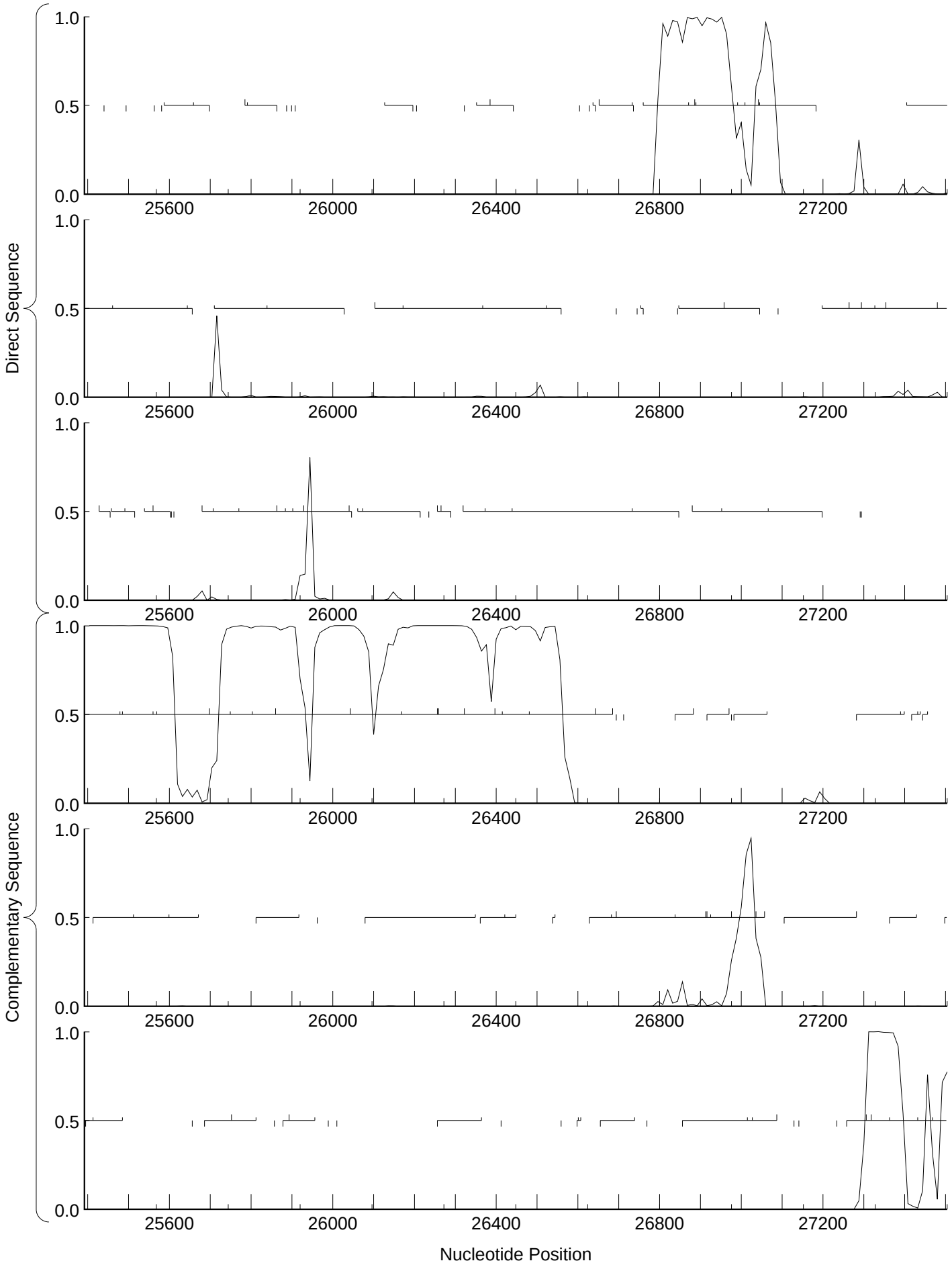
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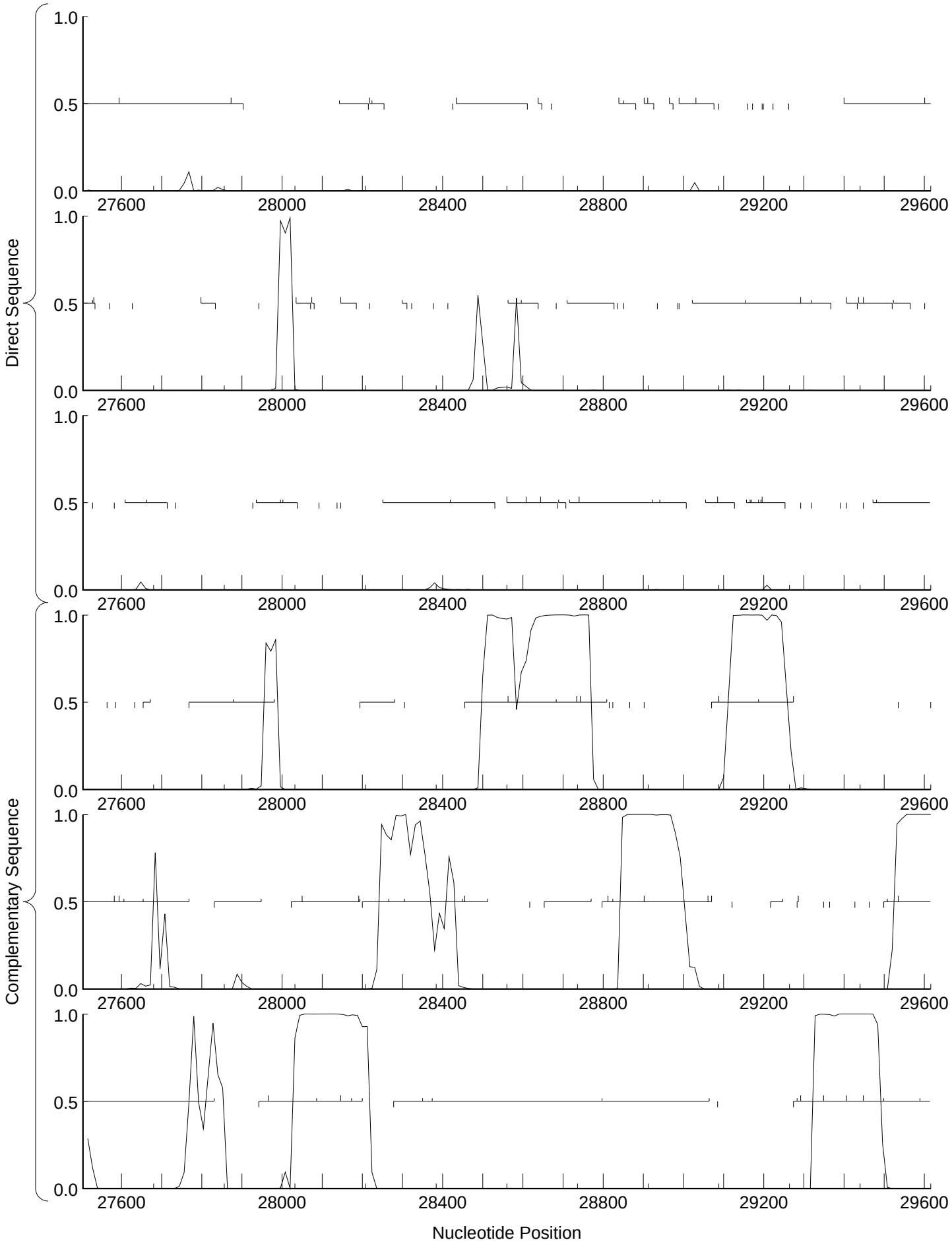
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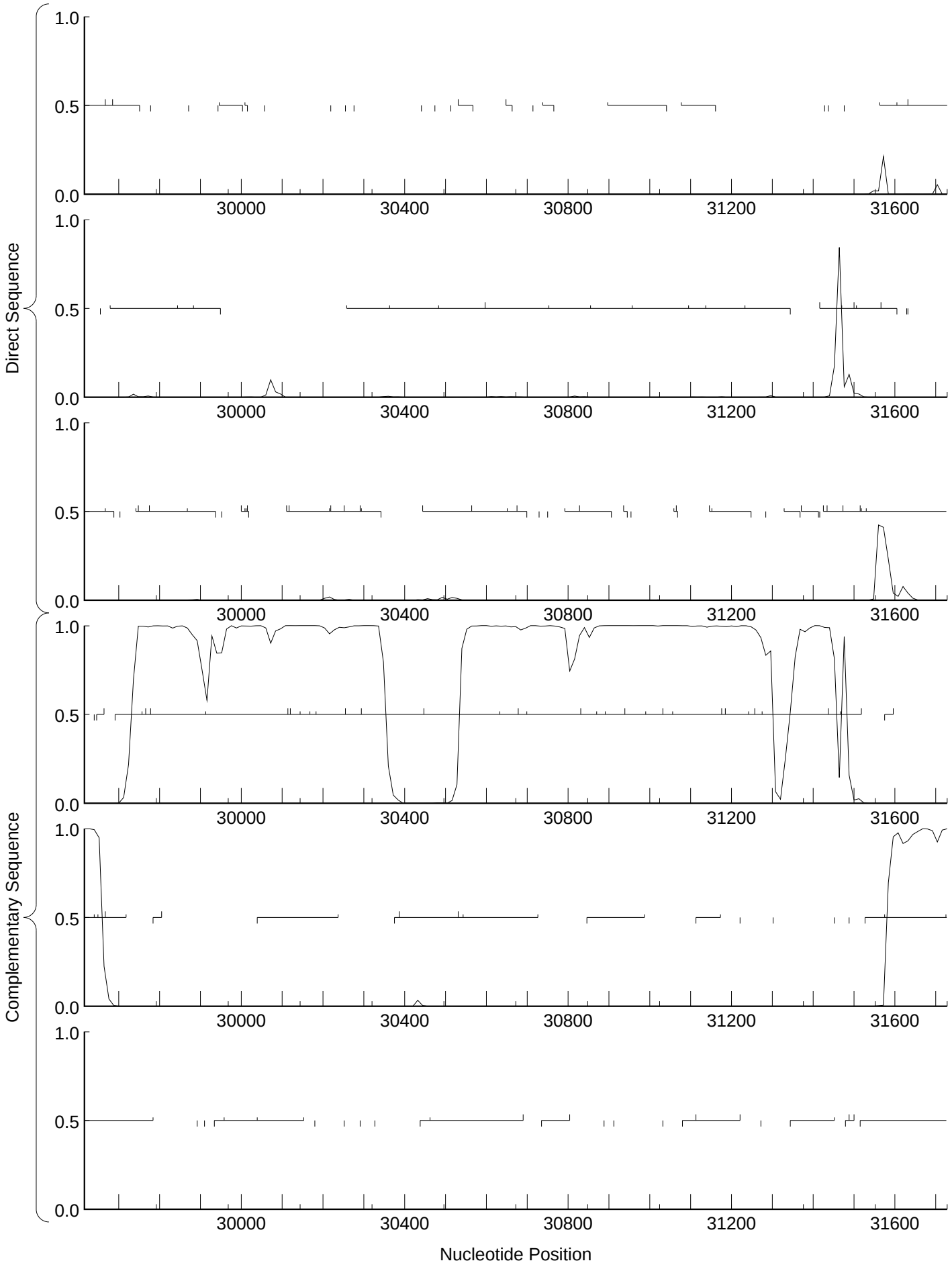
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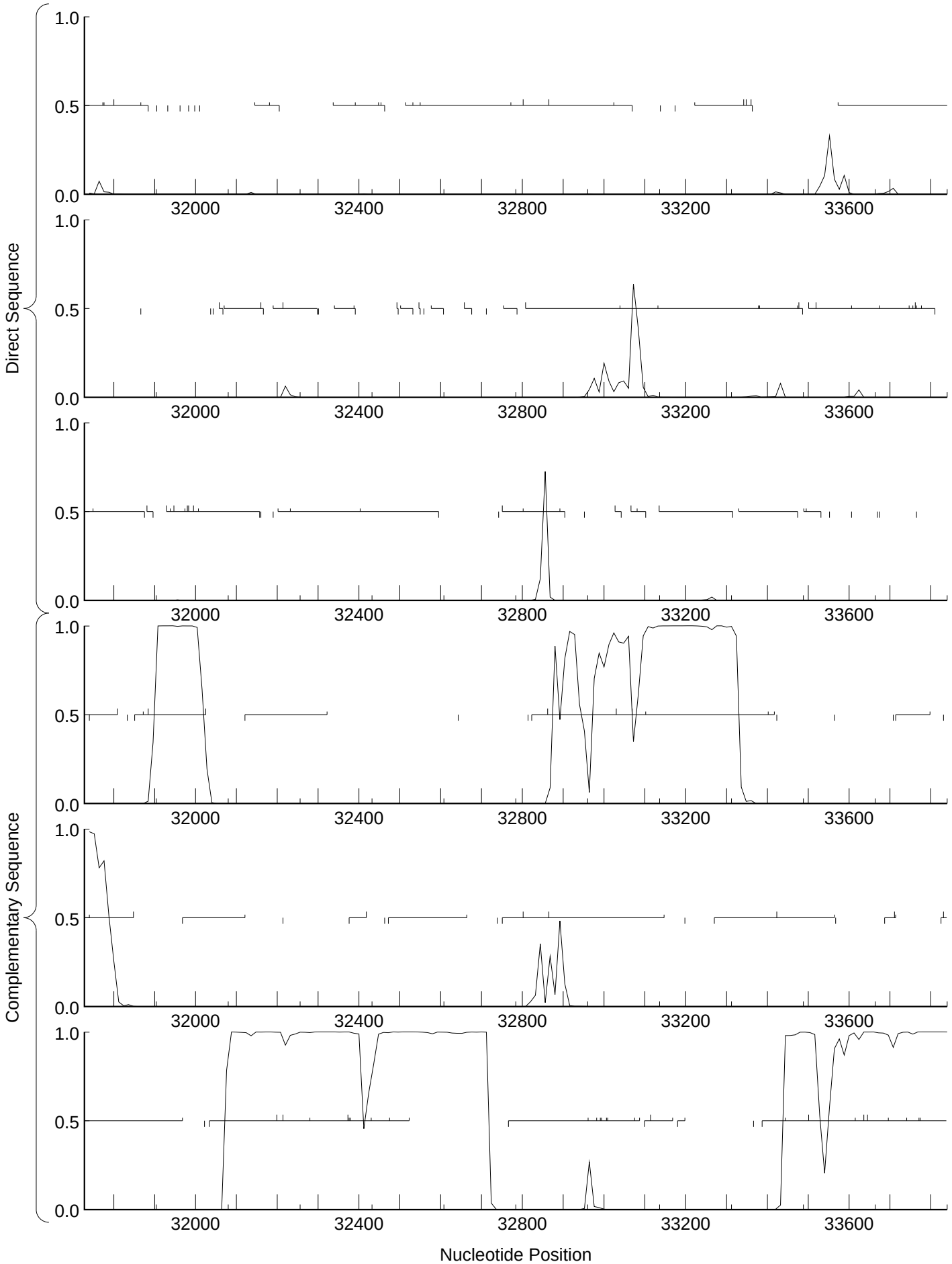
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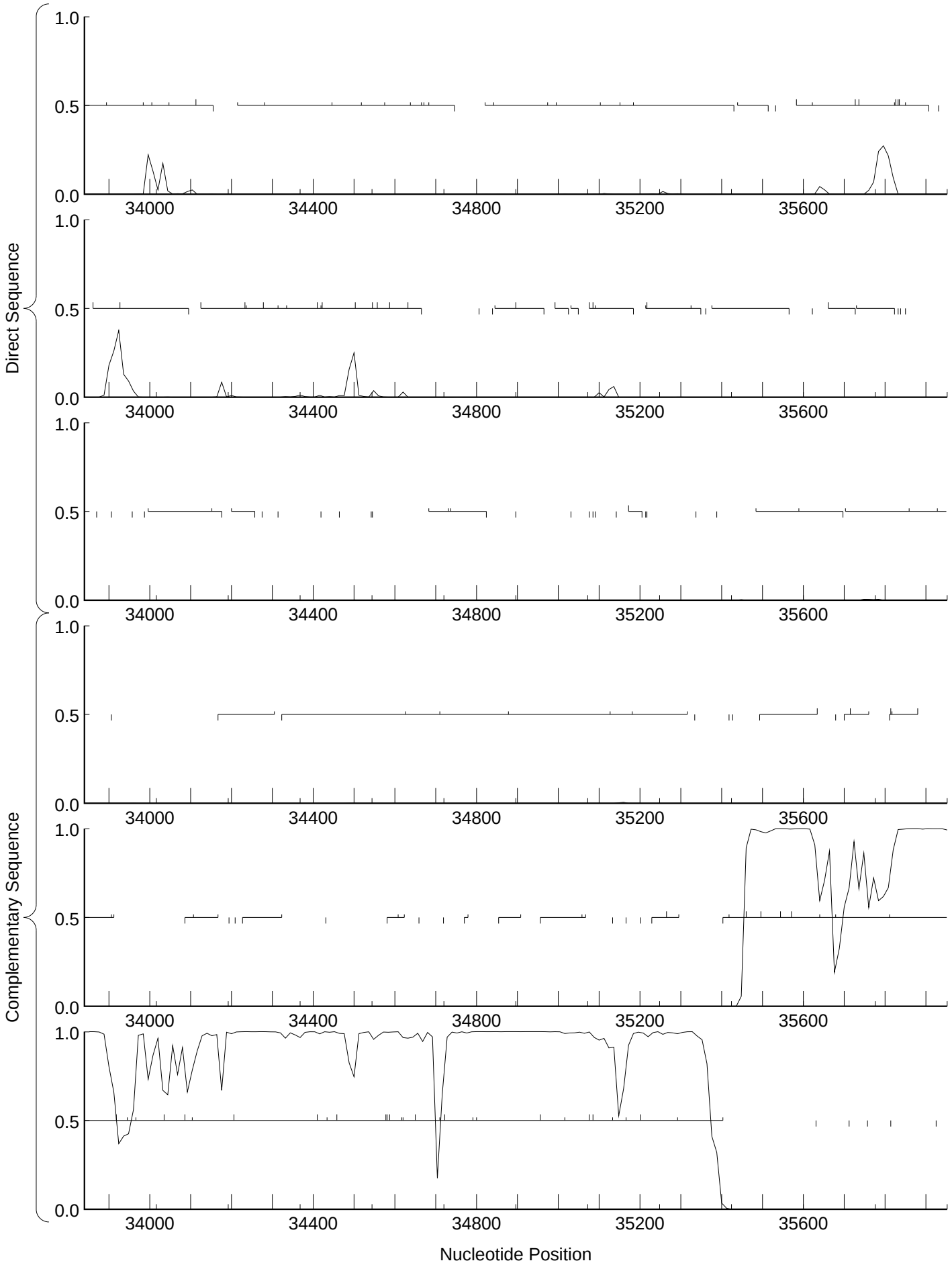
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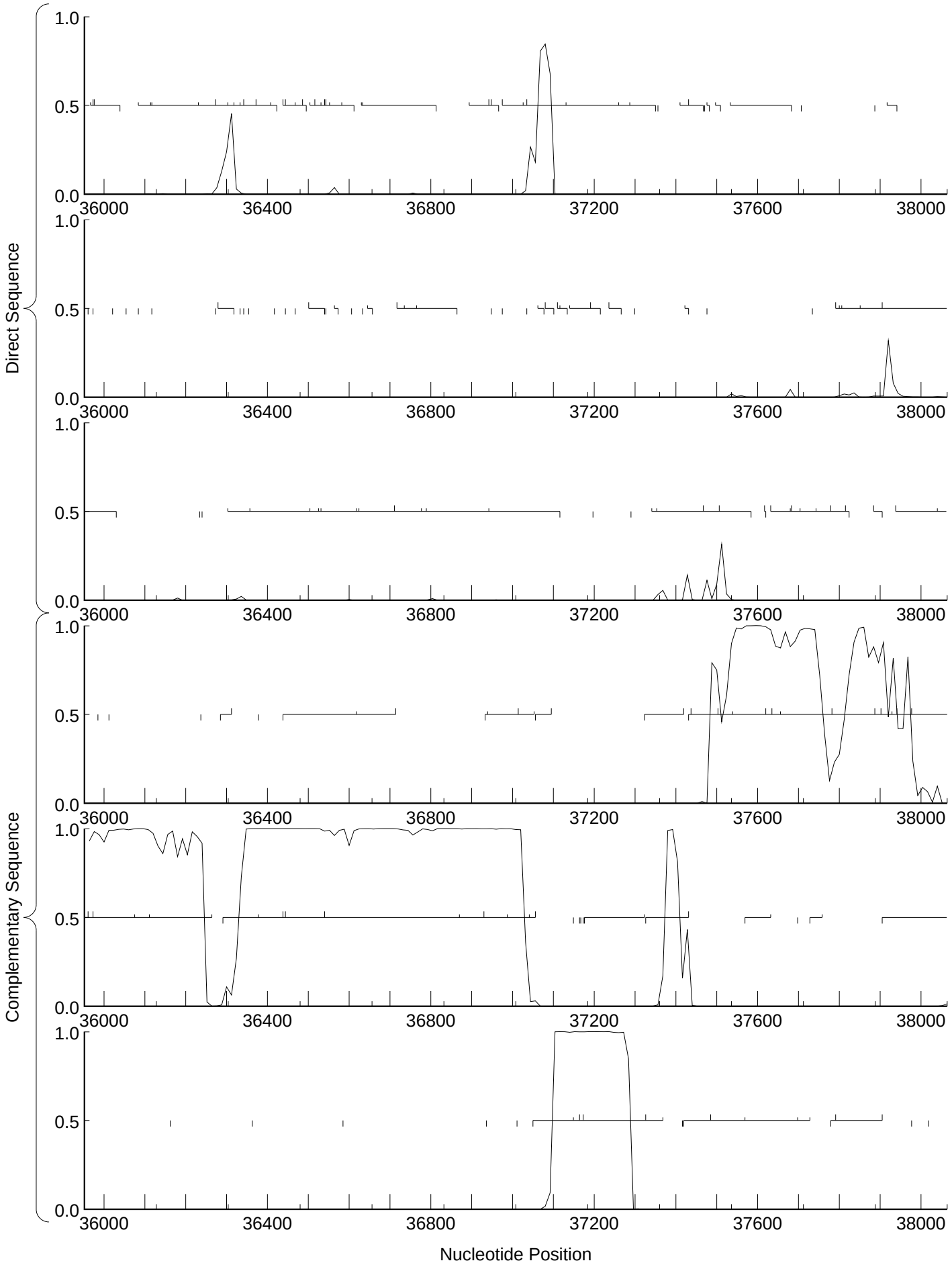
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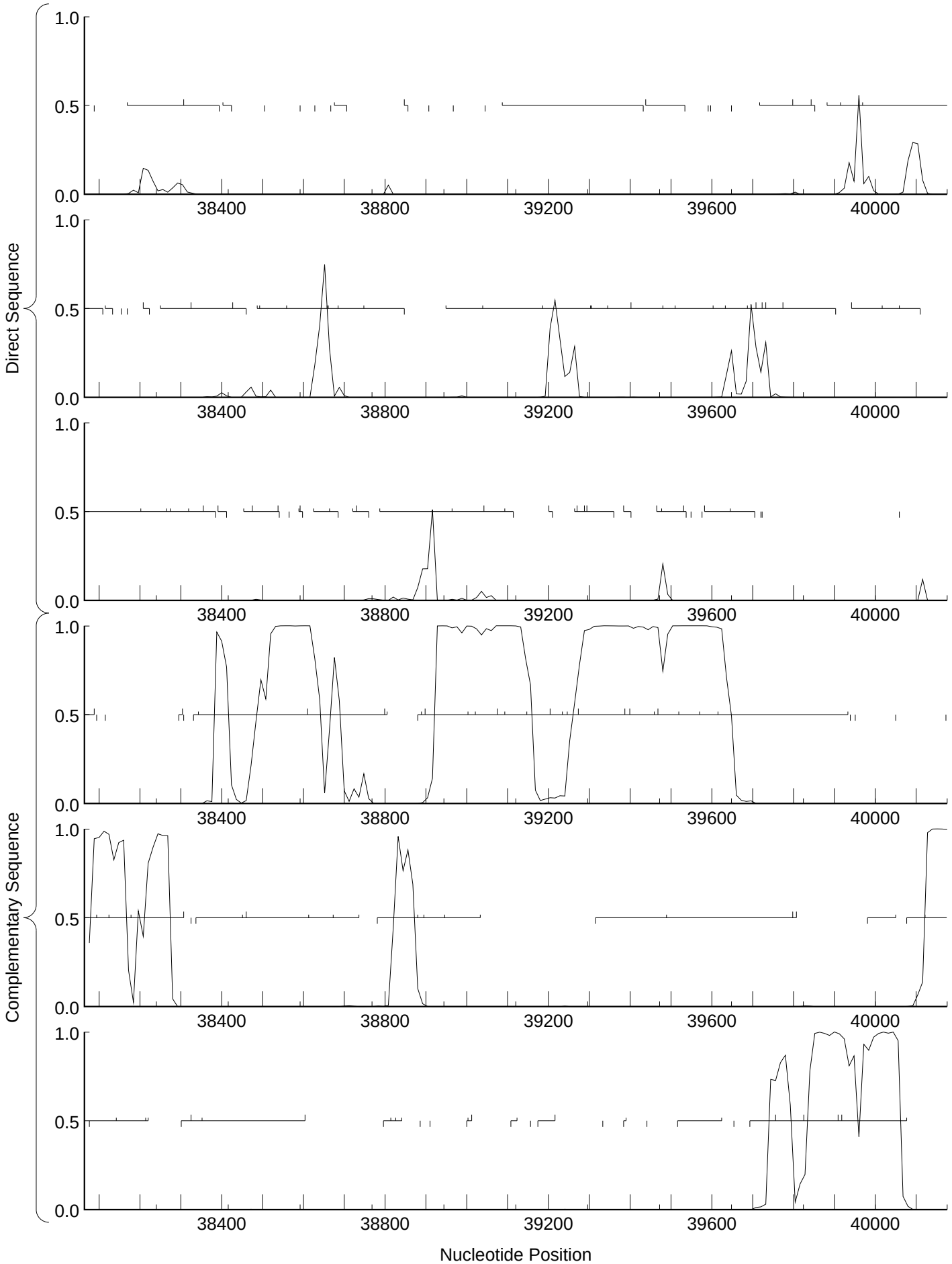
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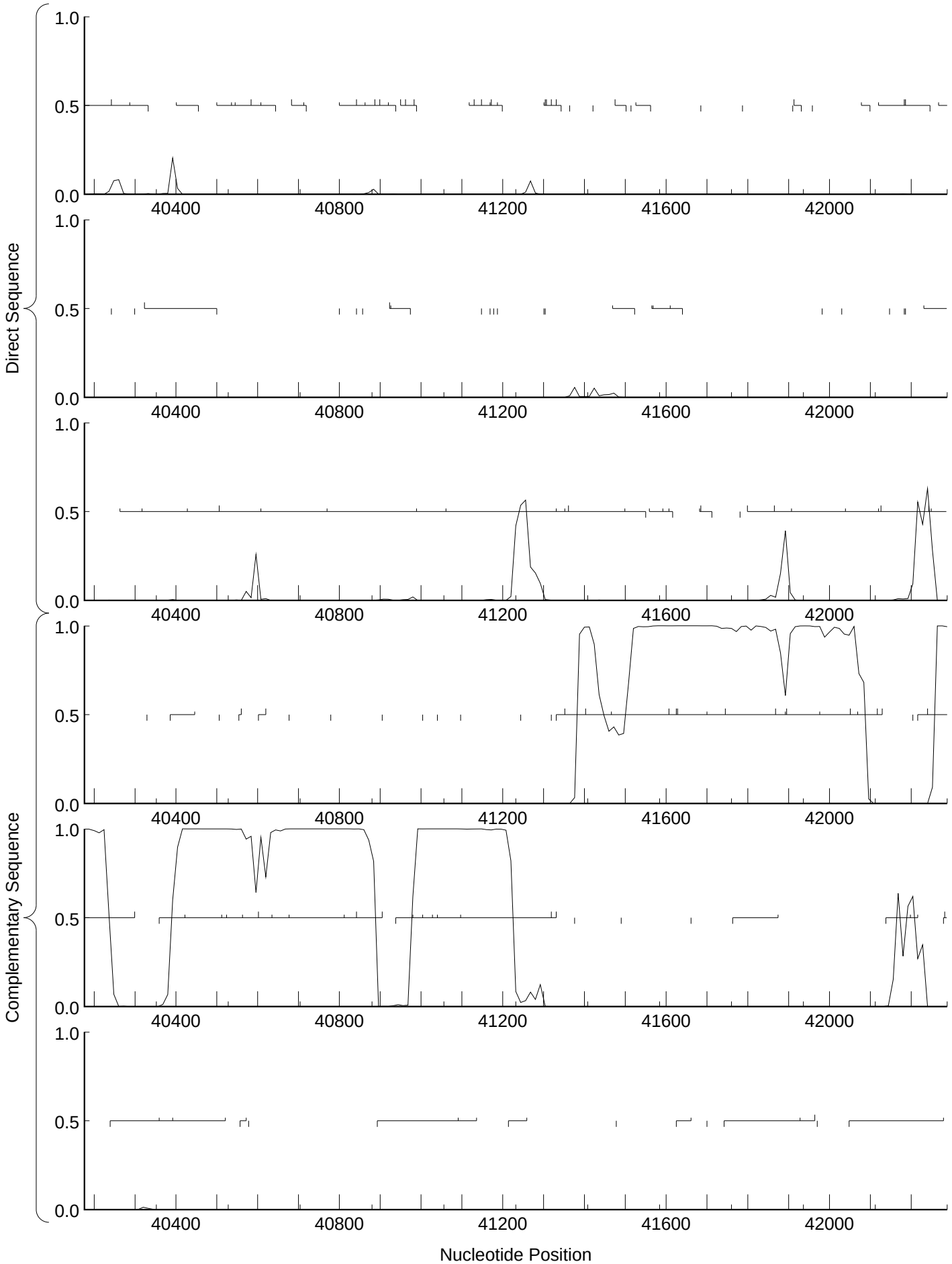
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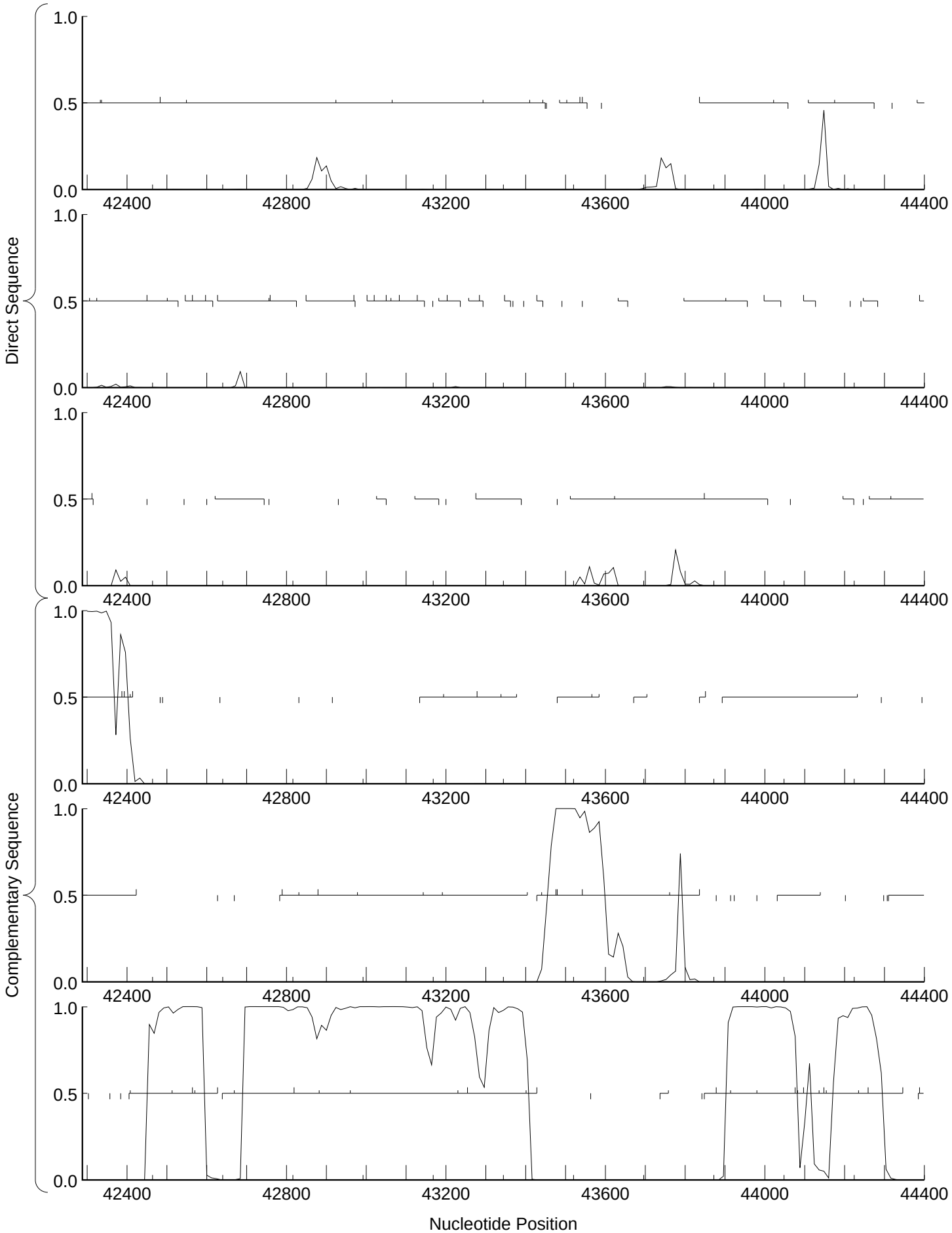
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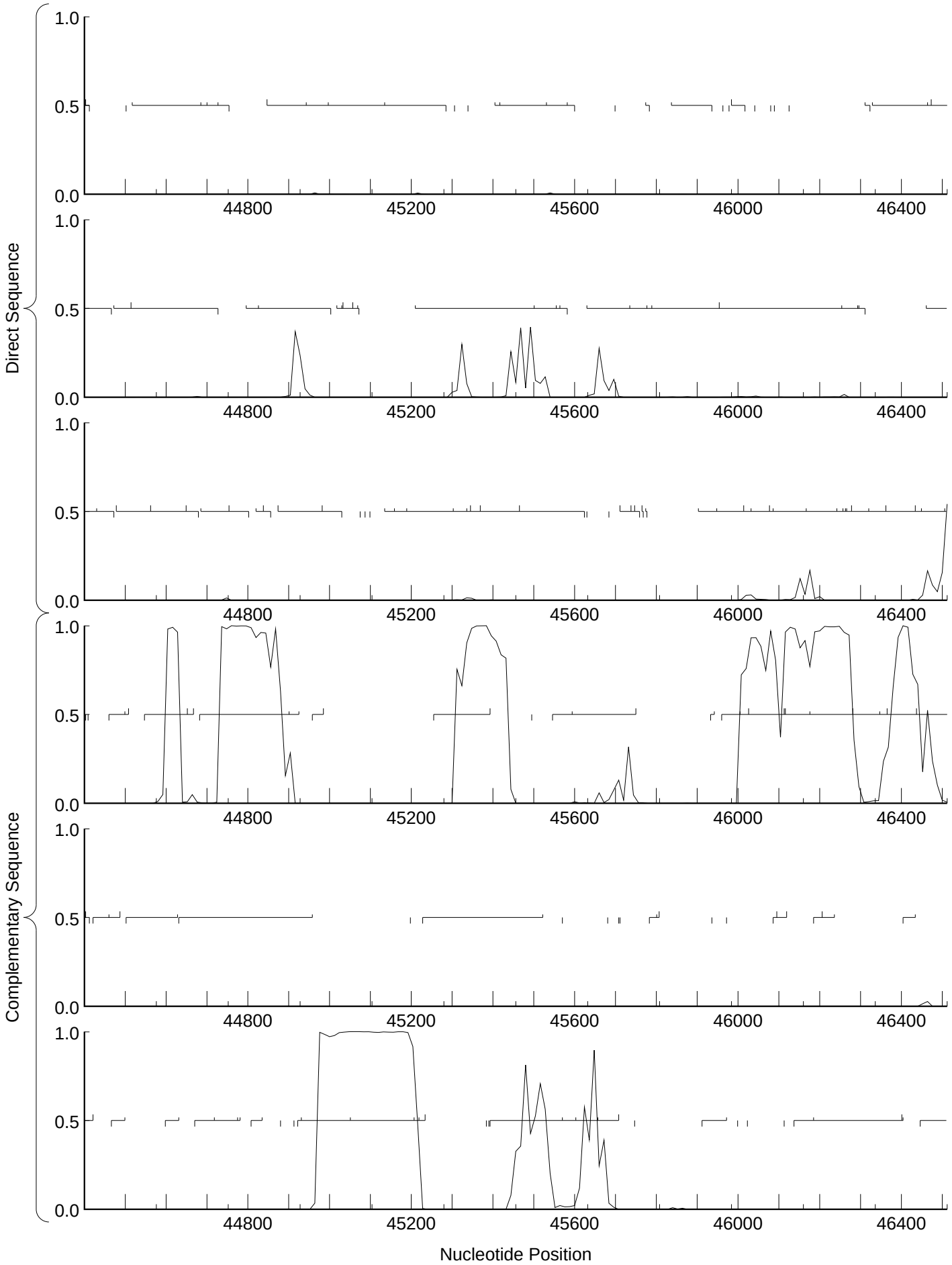
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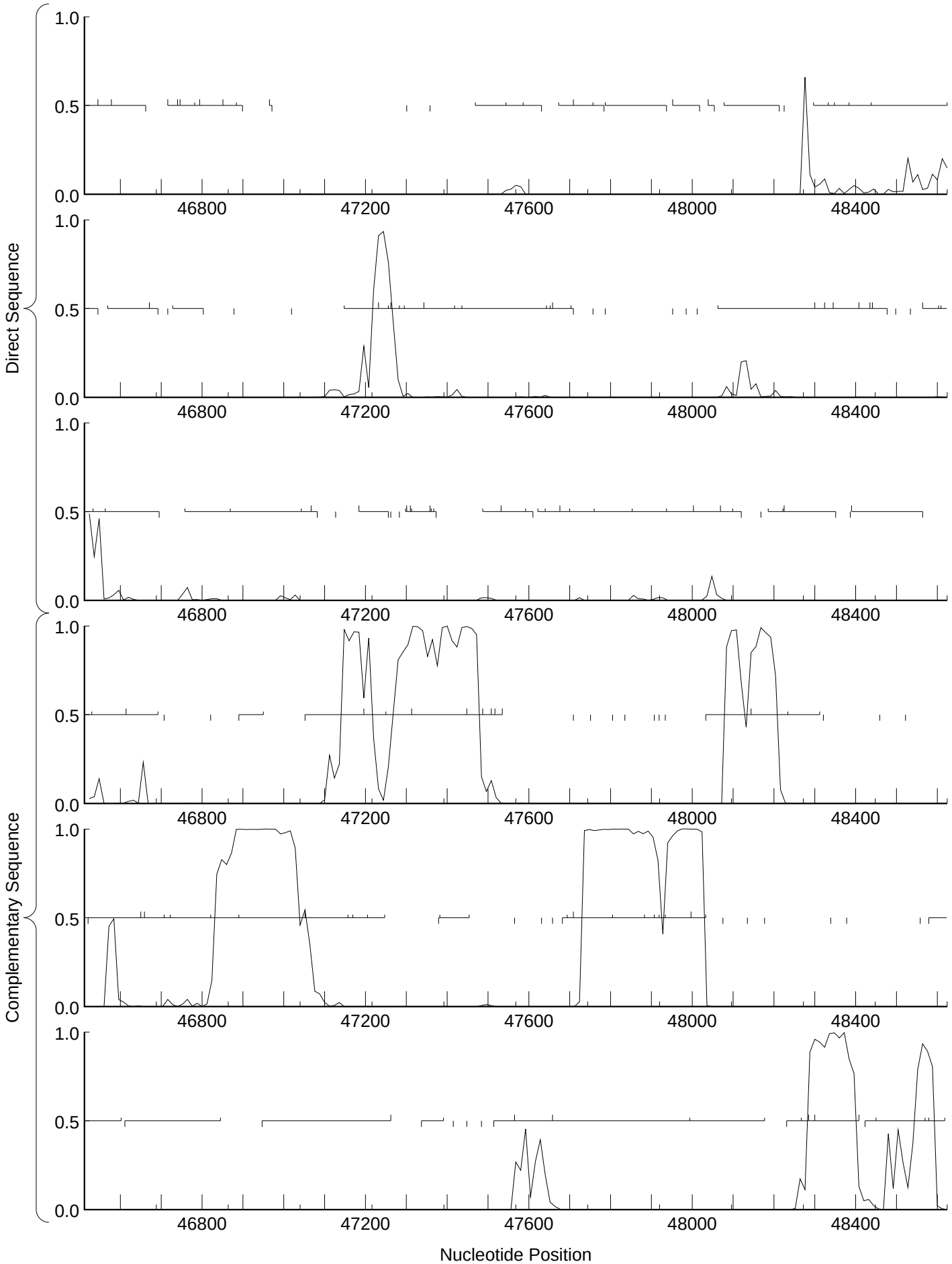
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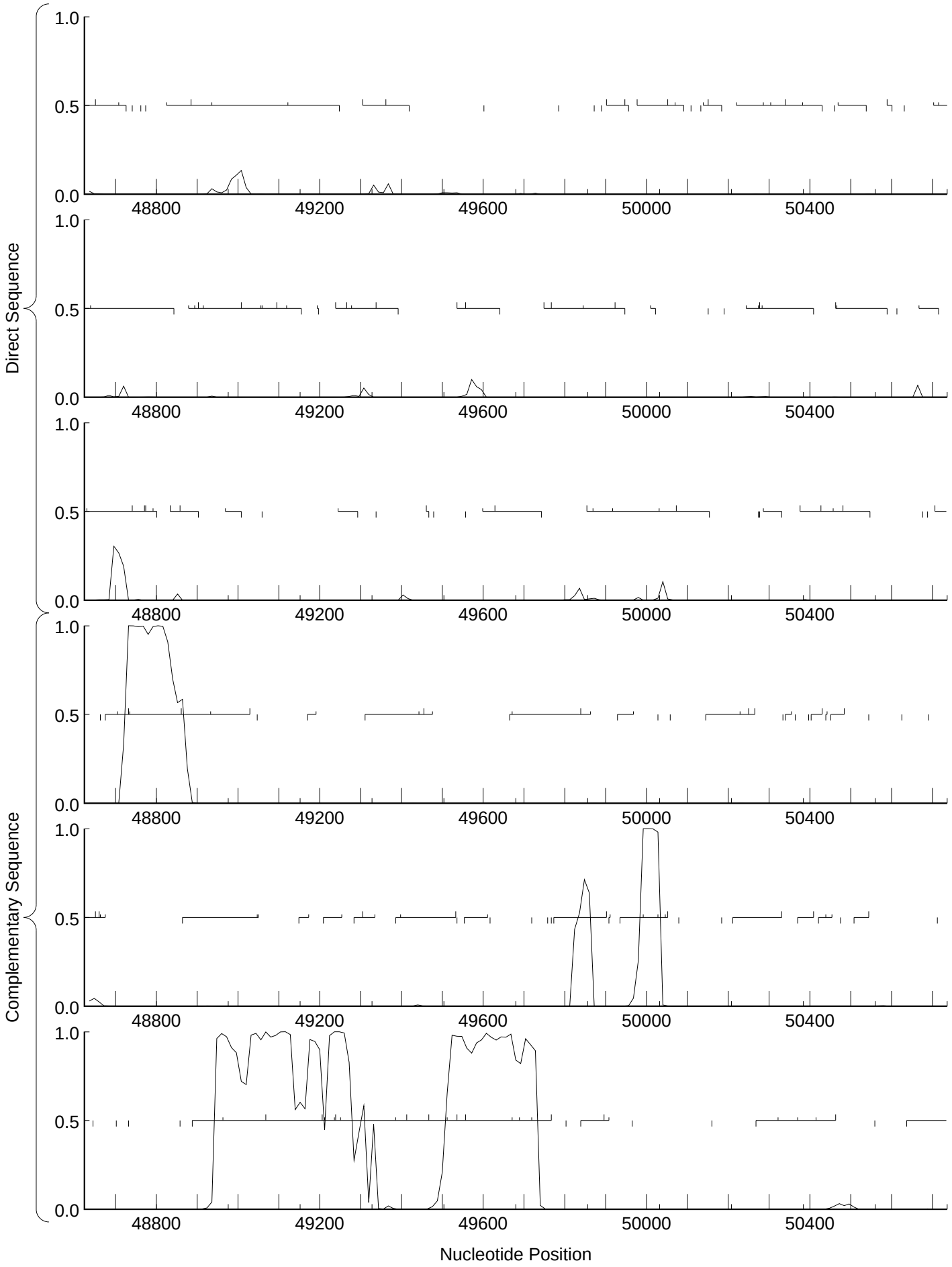
complete sequence, 51264 bp including 10-base 3’ overhang (CGGCCGGTTA), Cluster A4, Order 4, Window 96, Step 12, 23/26



complete sequence, 51264 bp including 10-base 3’ overhang (CGGCCGGTTA), Cluster A4, Order 4, Window 96, Step 12, 24/26



complete sequence, 51264 bp including 10-base 3’ overhang (CGGCCGGTTA), Cluster A4, Order 4, Window 96, Step 12, 25/26



complete sequence, 51264 bp including 10-base 3’ overhang (CGGCCGGTTA), Cluster A4, Order 4, Window 96, Step 12, 26/26

