GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage DaWorst complete sequence, 56827 bp including 10-base 3’ overhang (CCGA
Analysis Date    :  3/28/24 at 15:16:03
Pages            :  28
Sequence Length  :  56827 bp
GC Content       :   60.98%

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

PS-Version       :  1.2

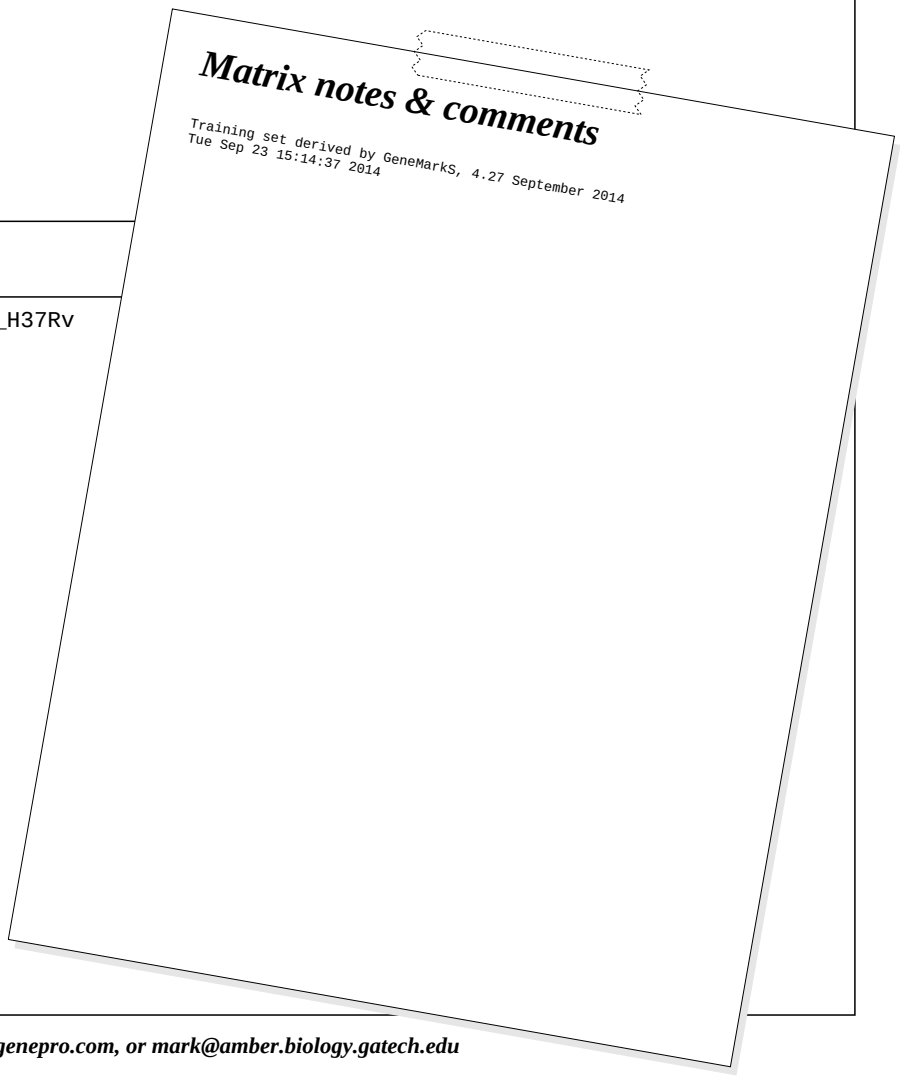
GeneMark Options :  PostScript graph,
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MATRIX INFORMATION

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Author     :  -

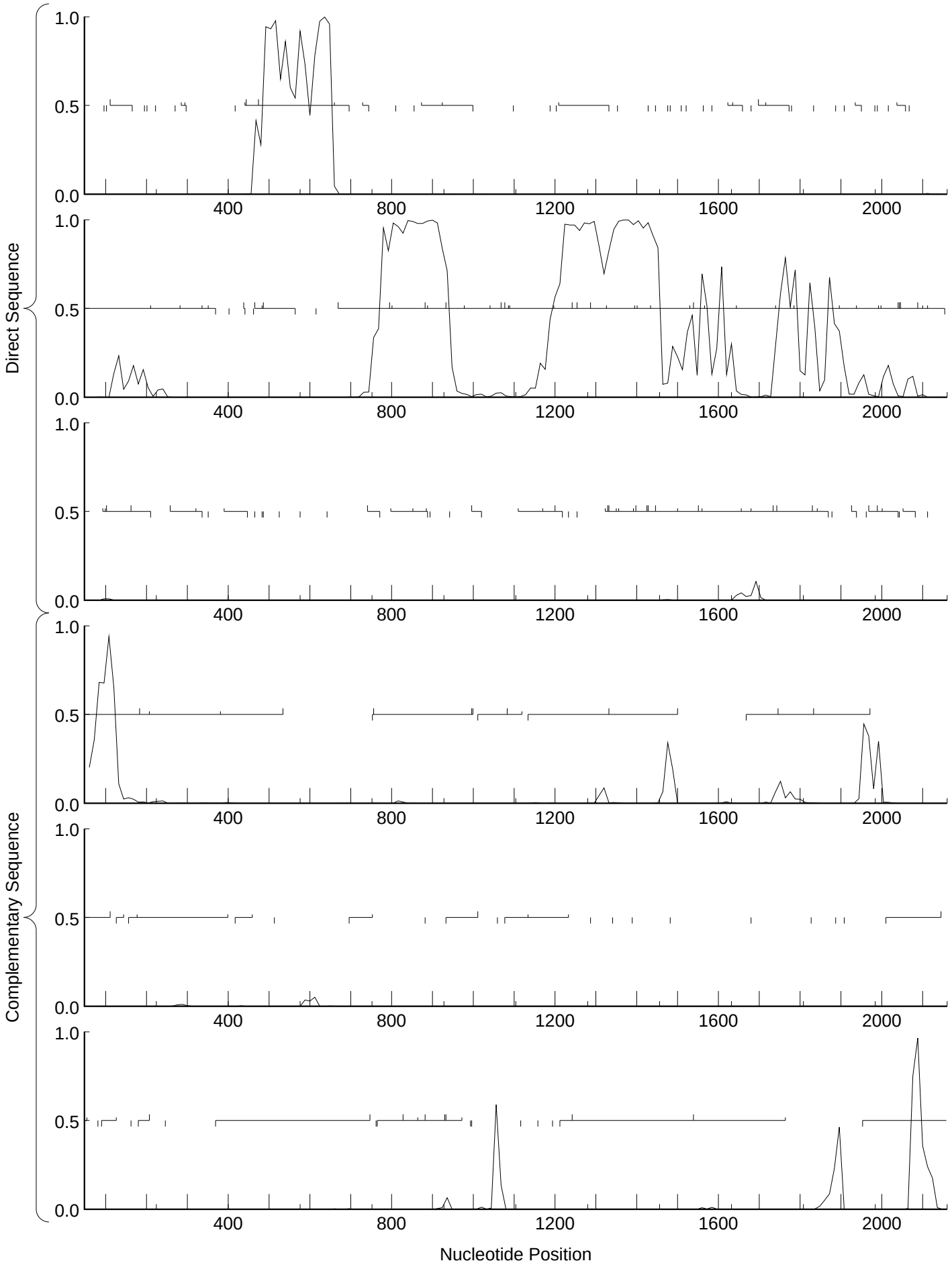
Order      :  4


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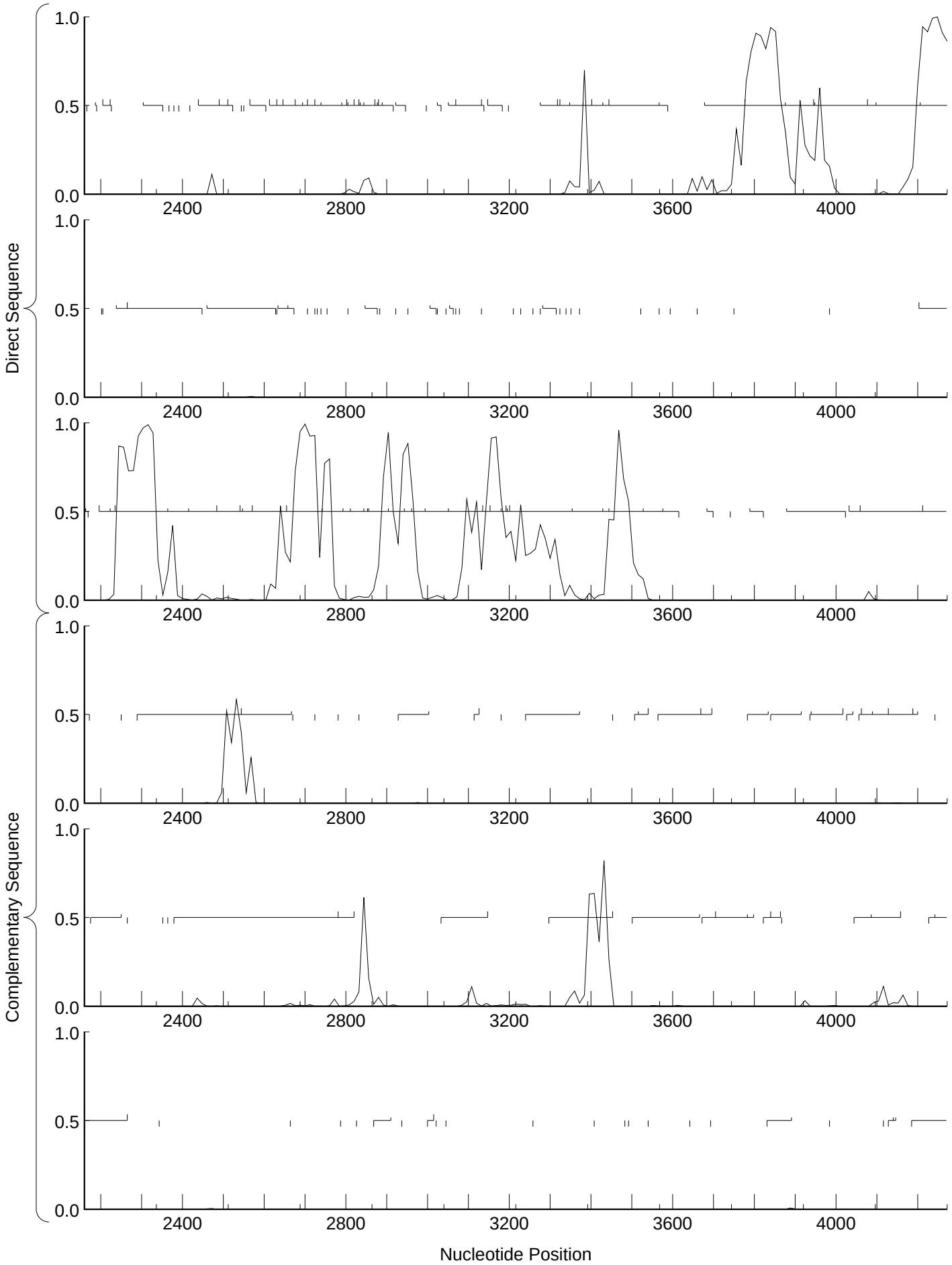


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

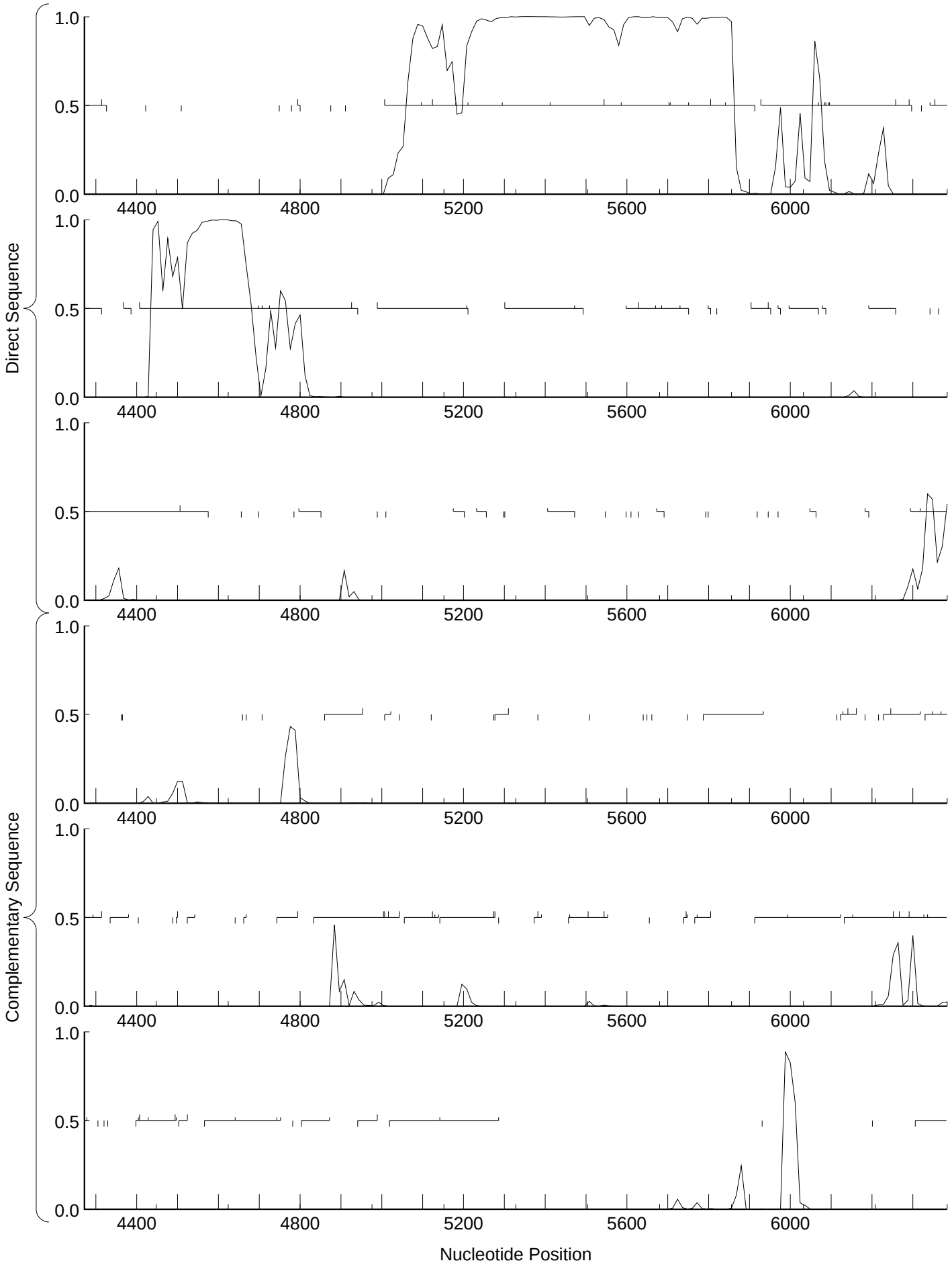
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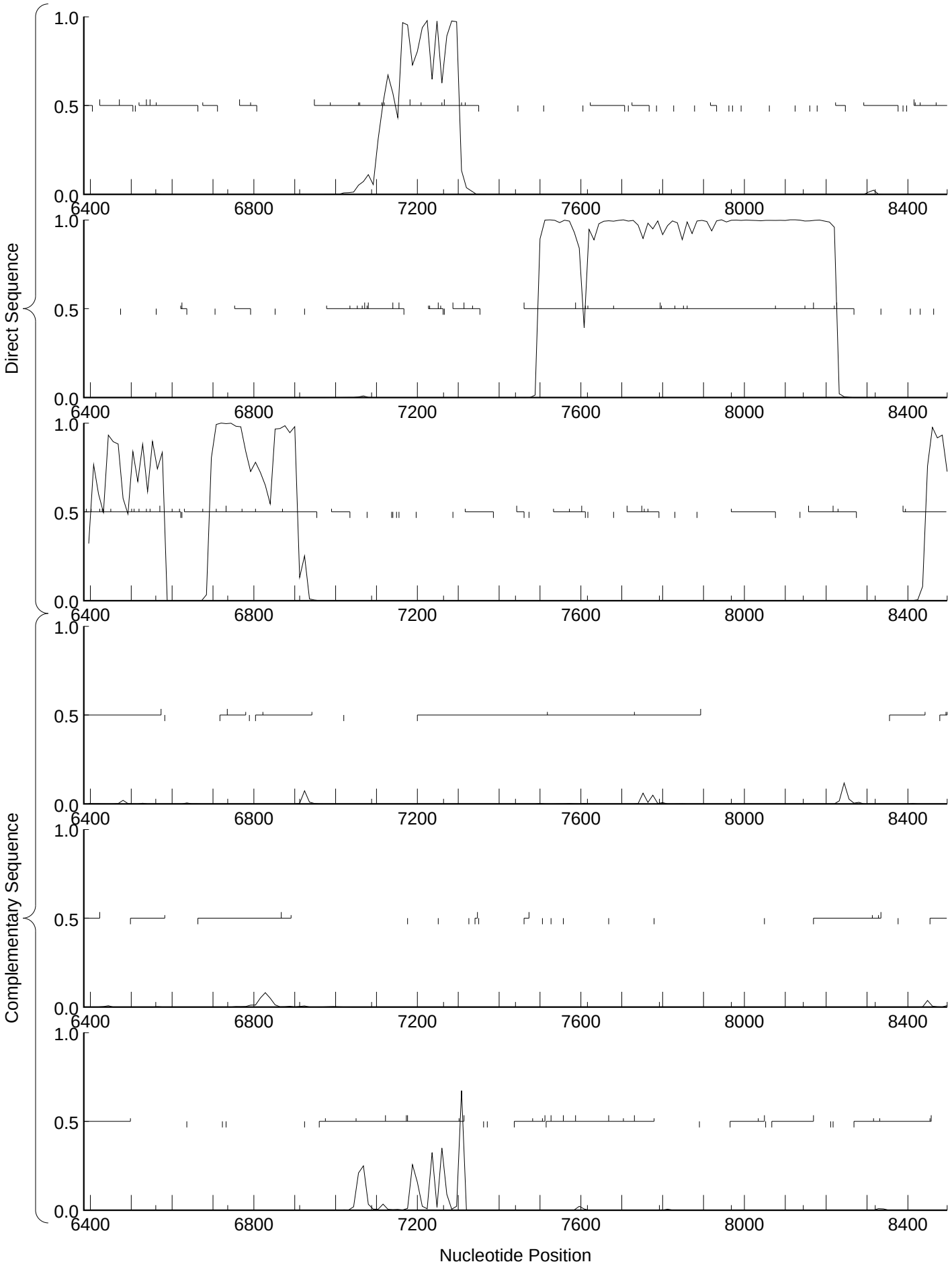
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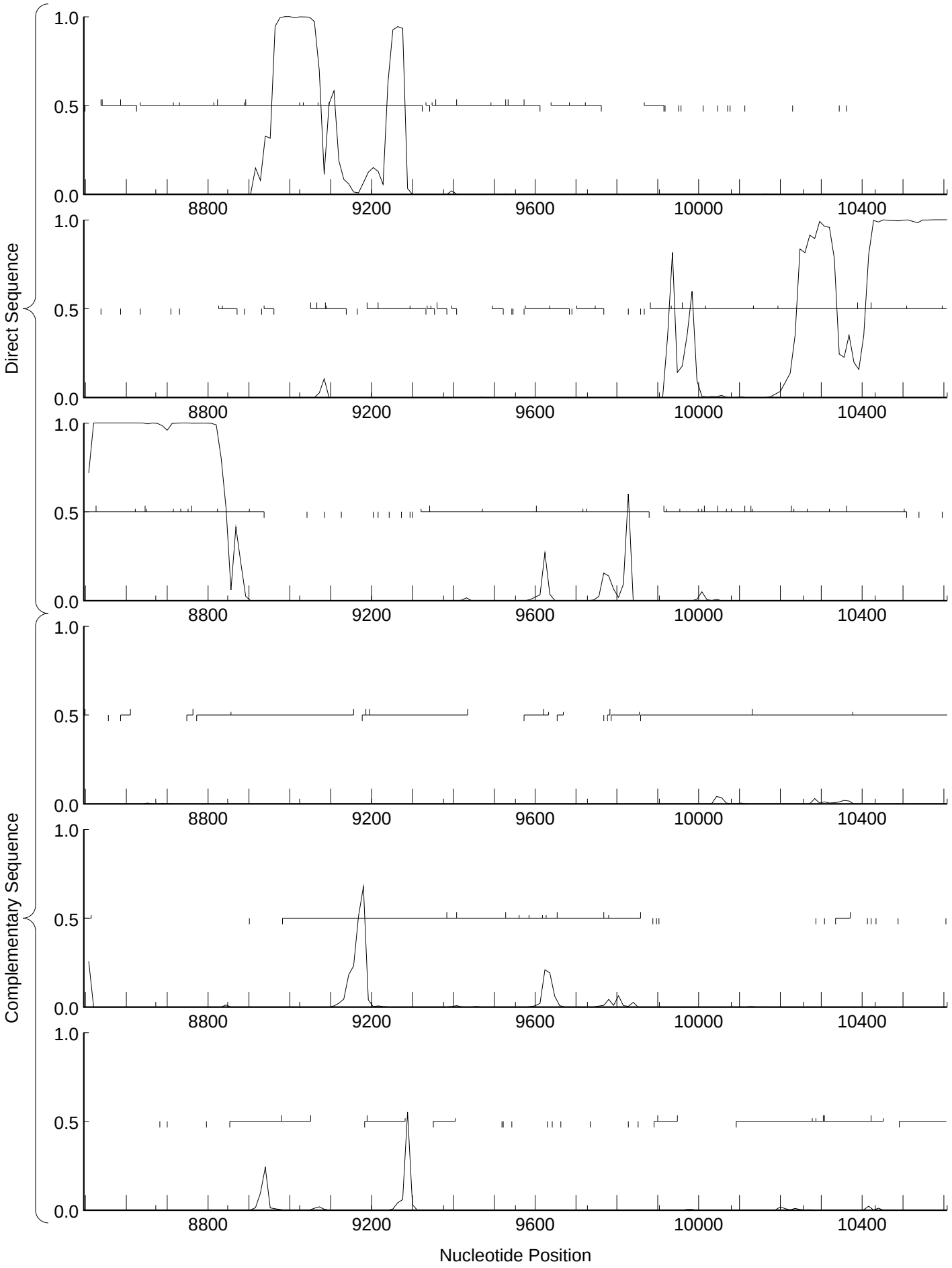
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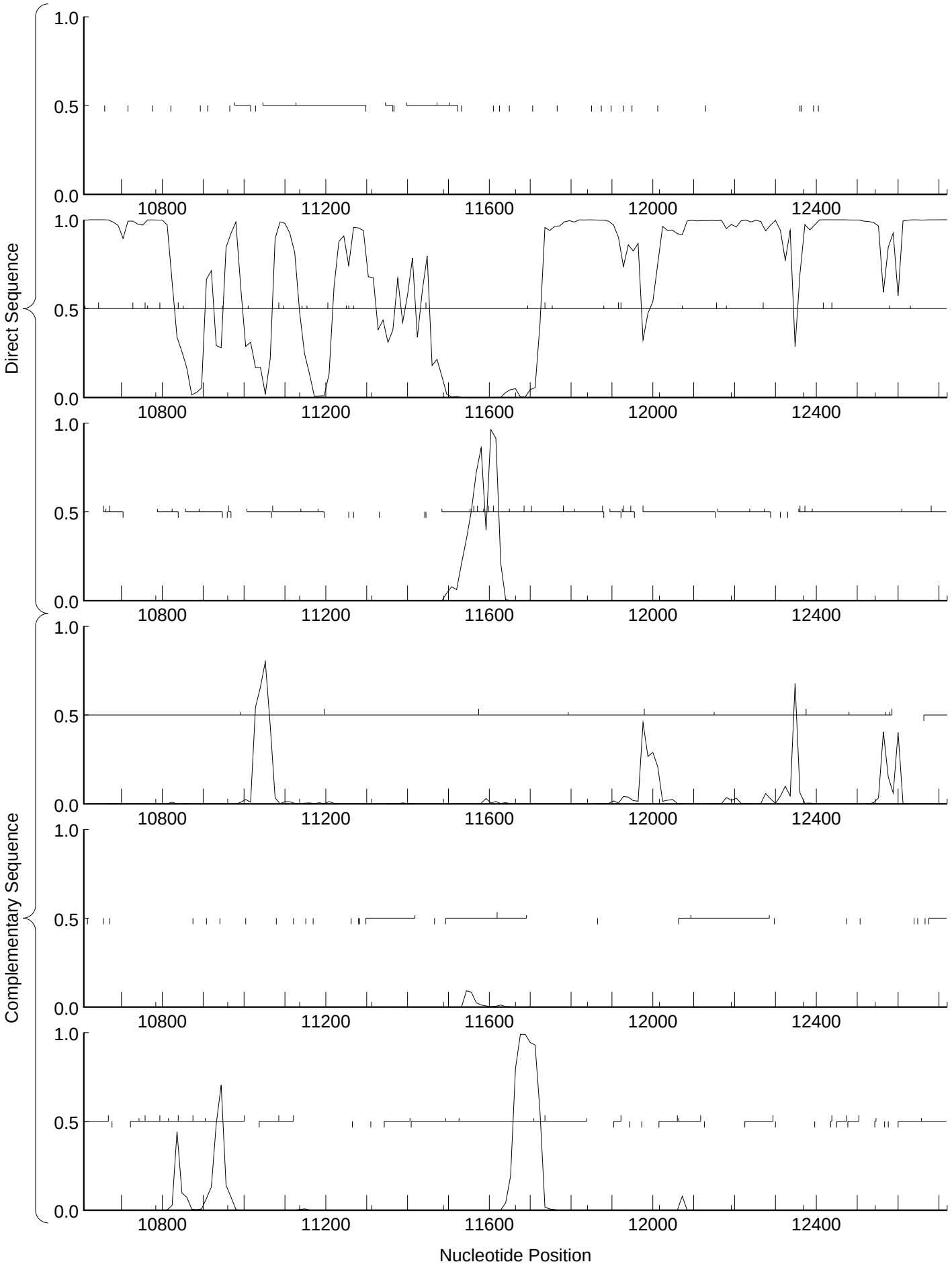
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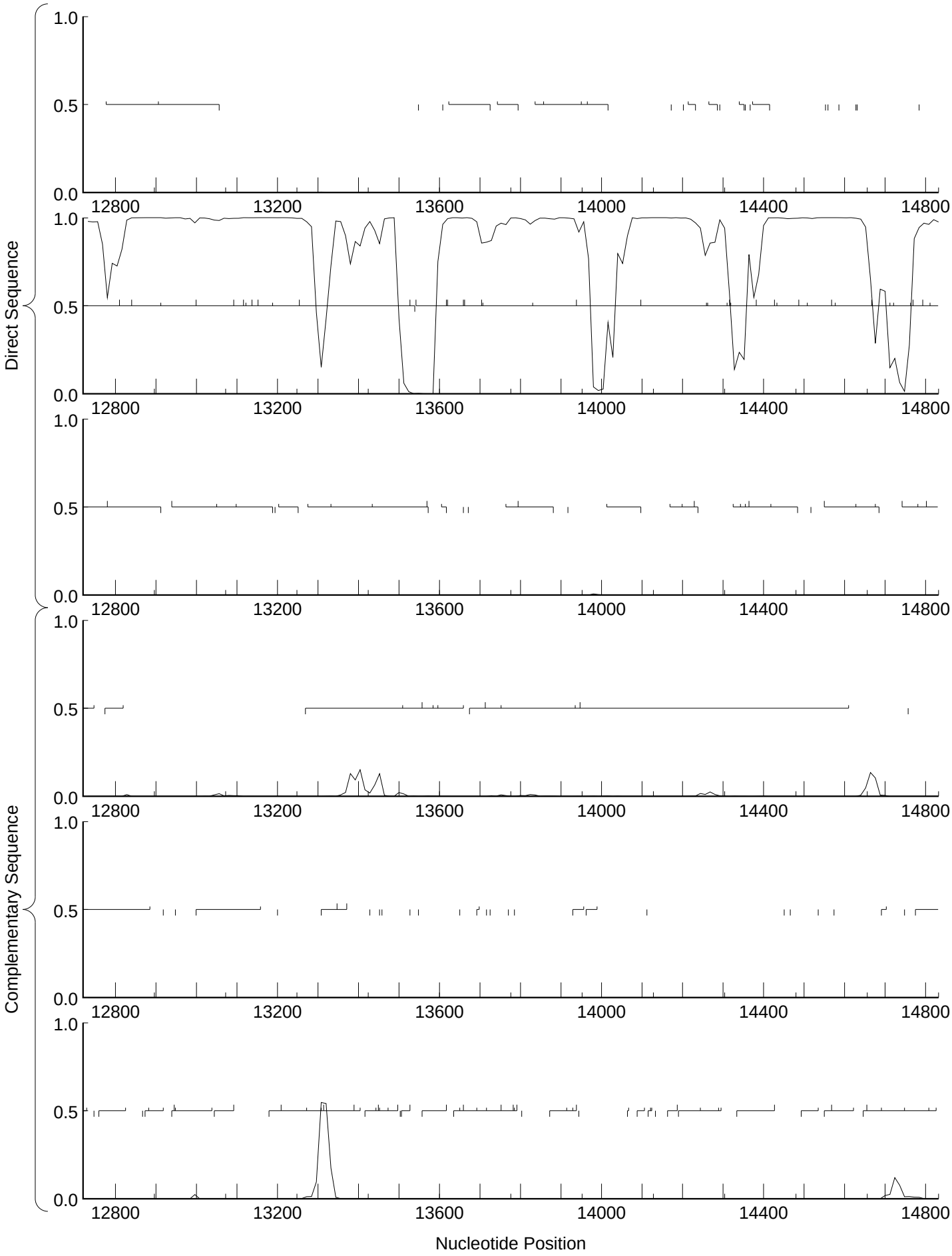
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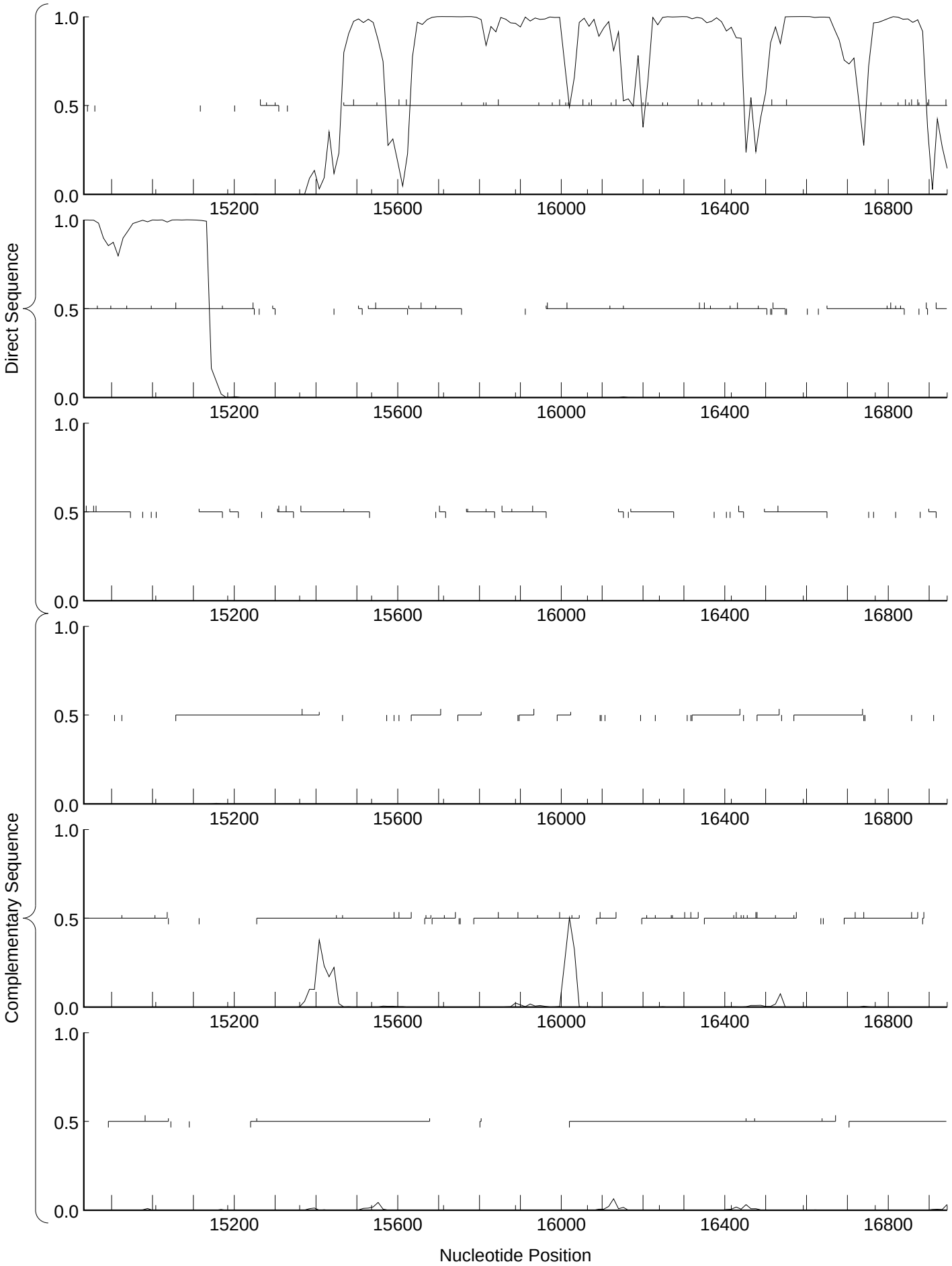
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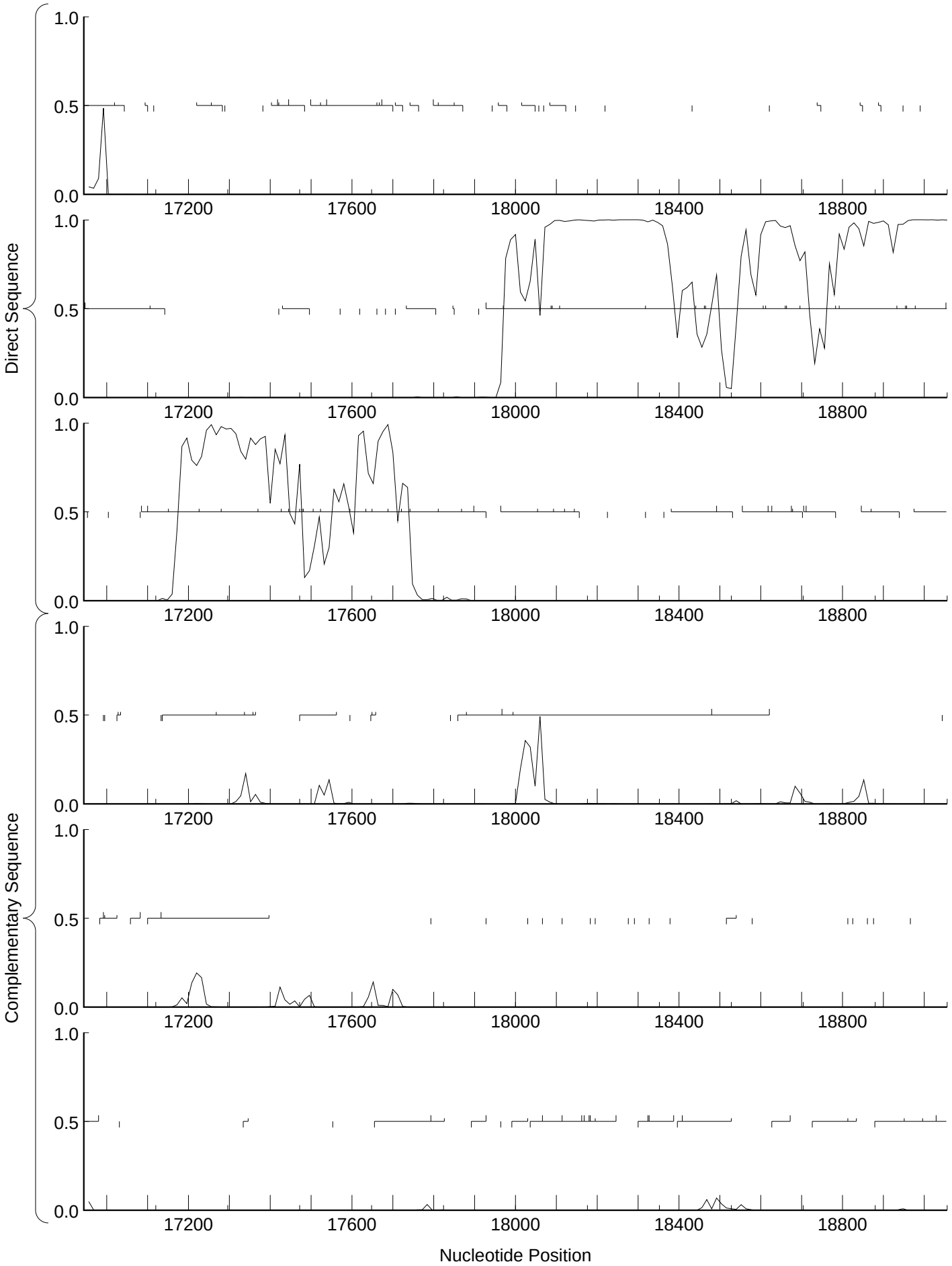
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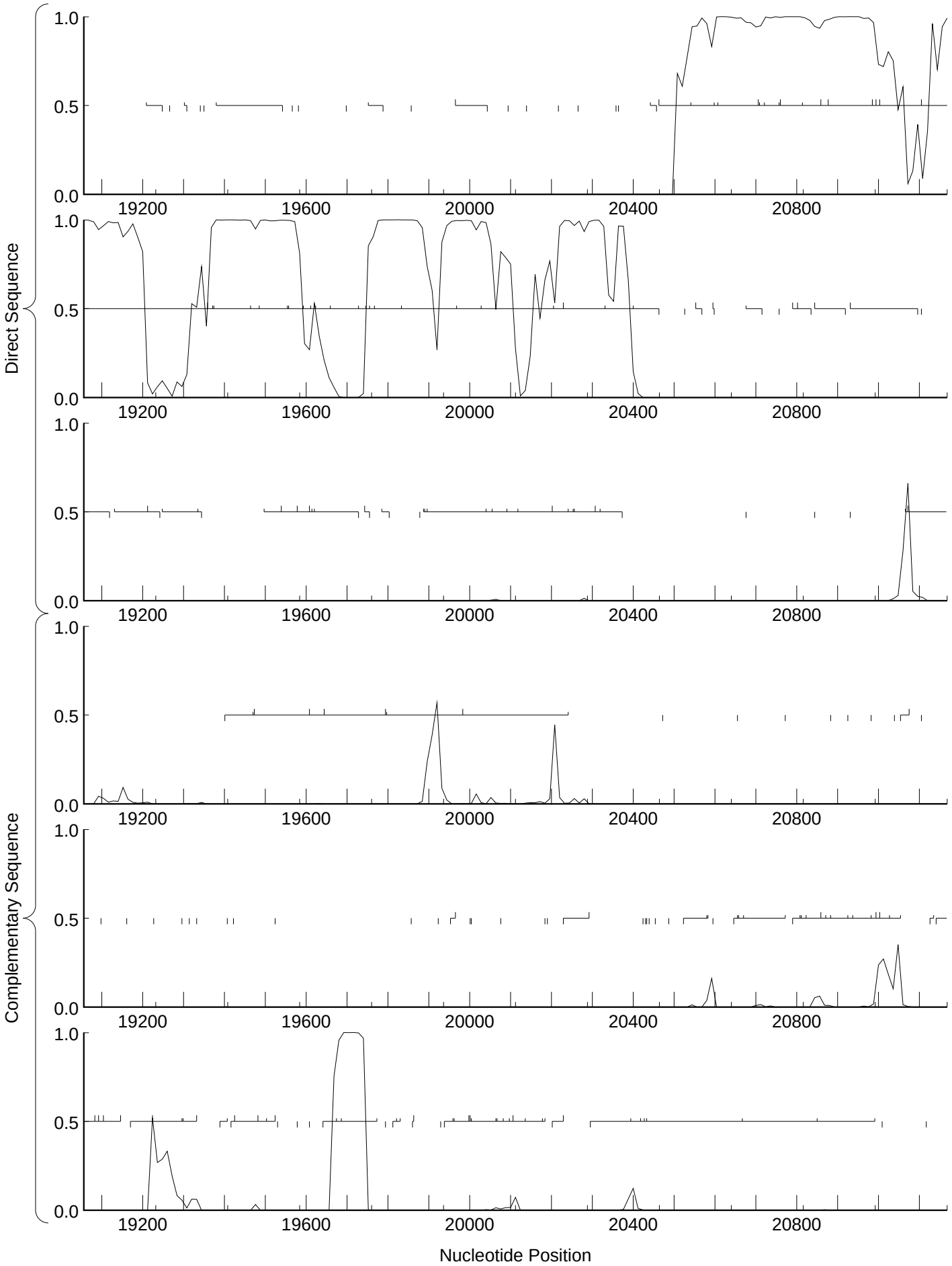
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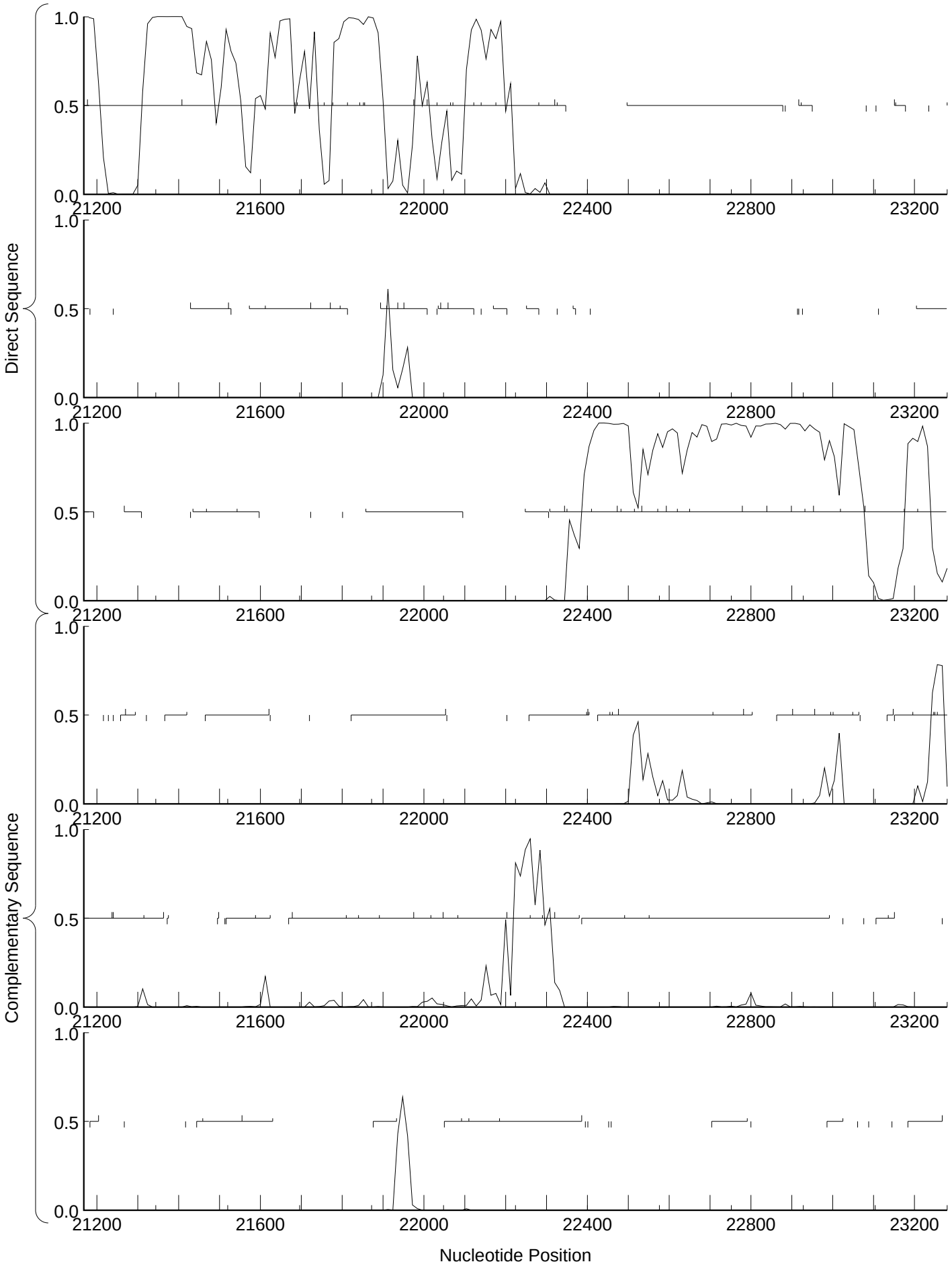
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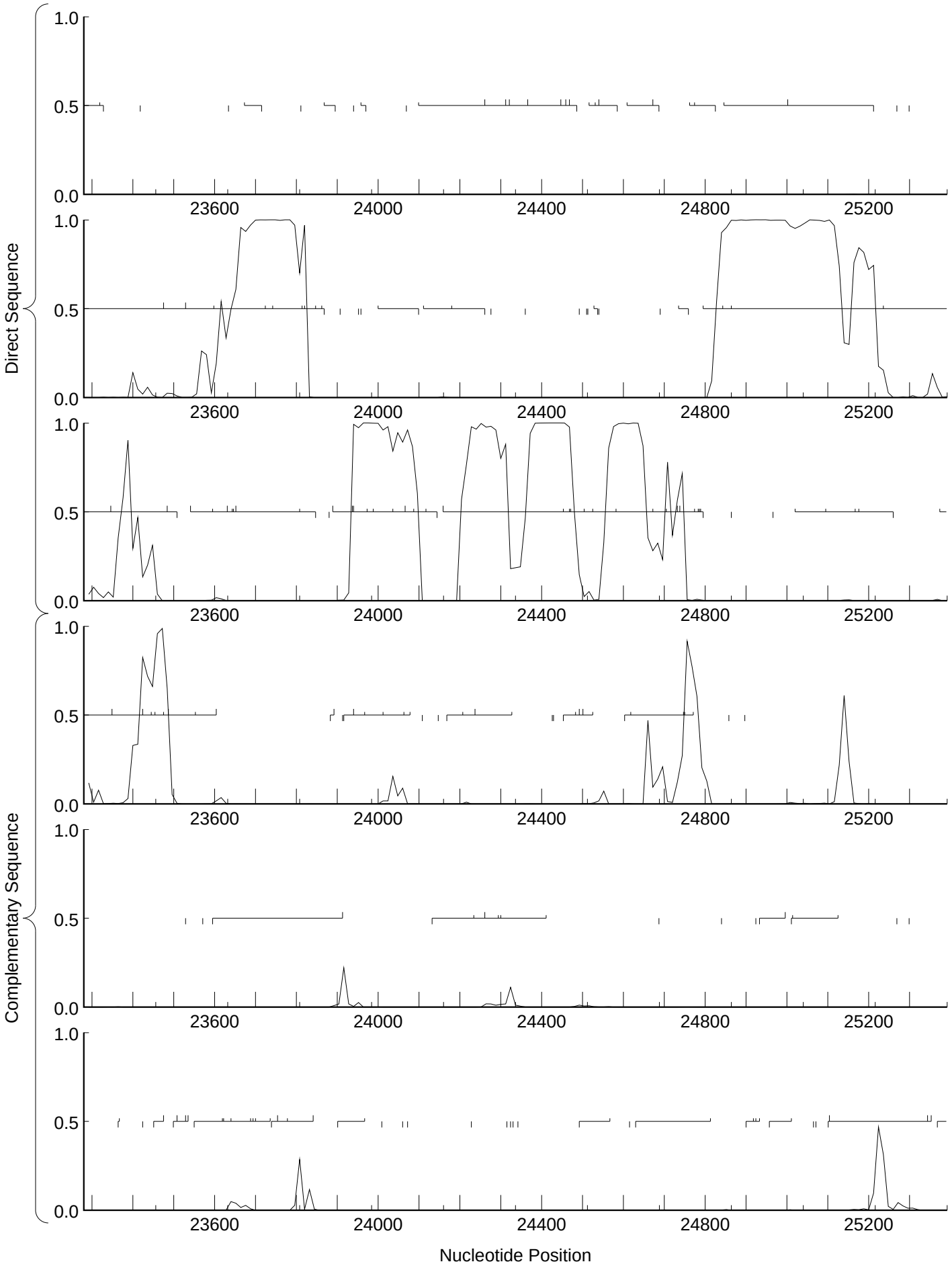
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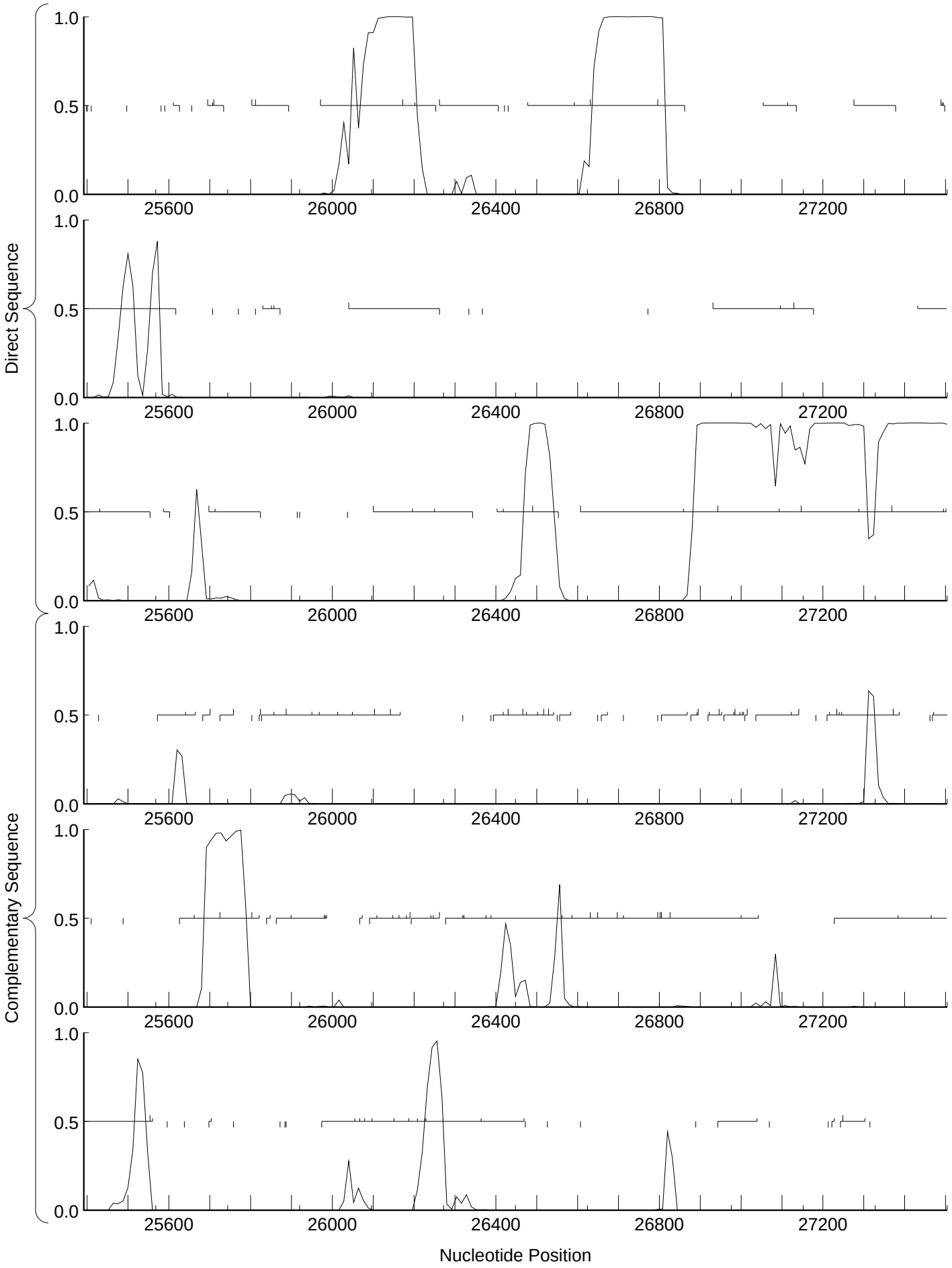
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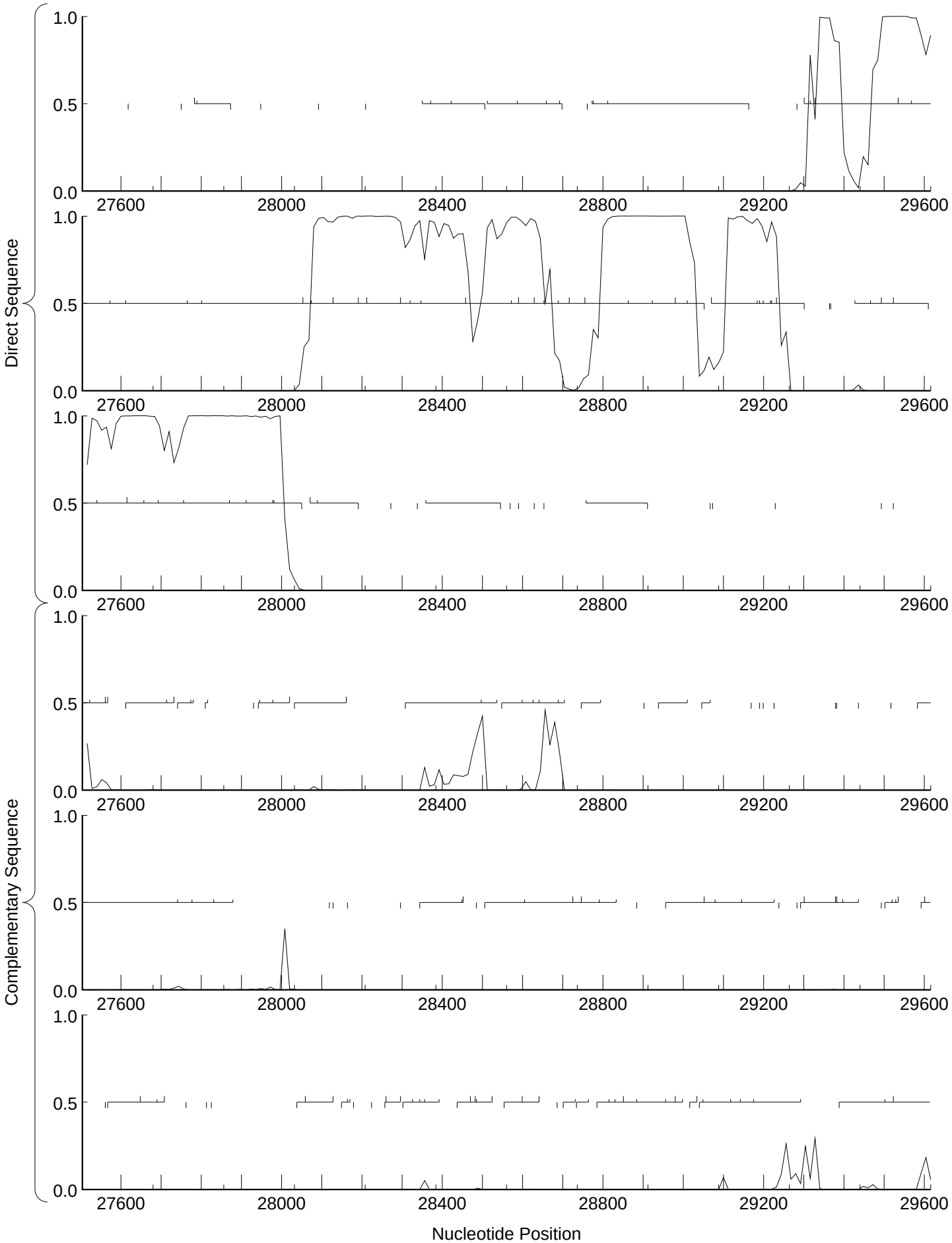
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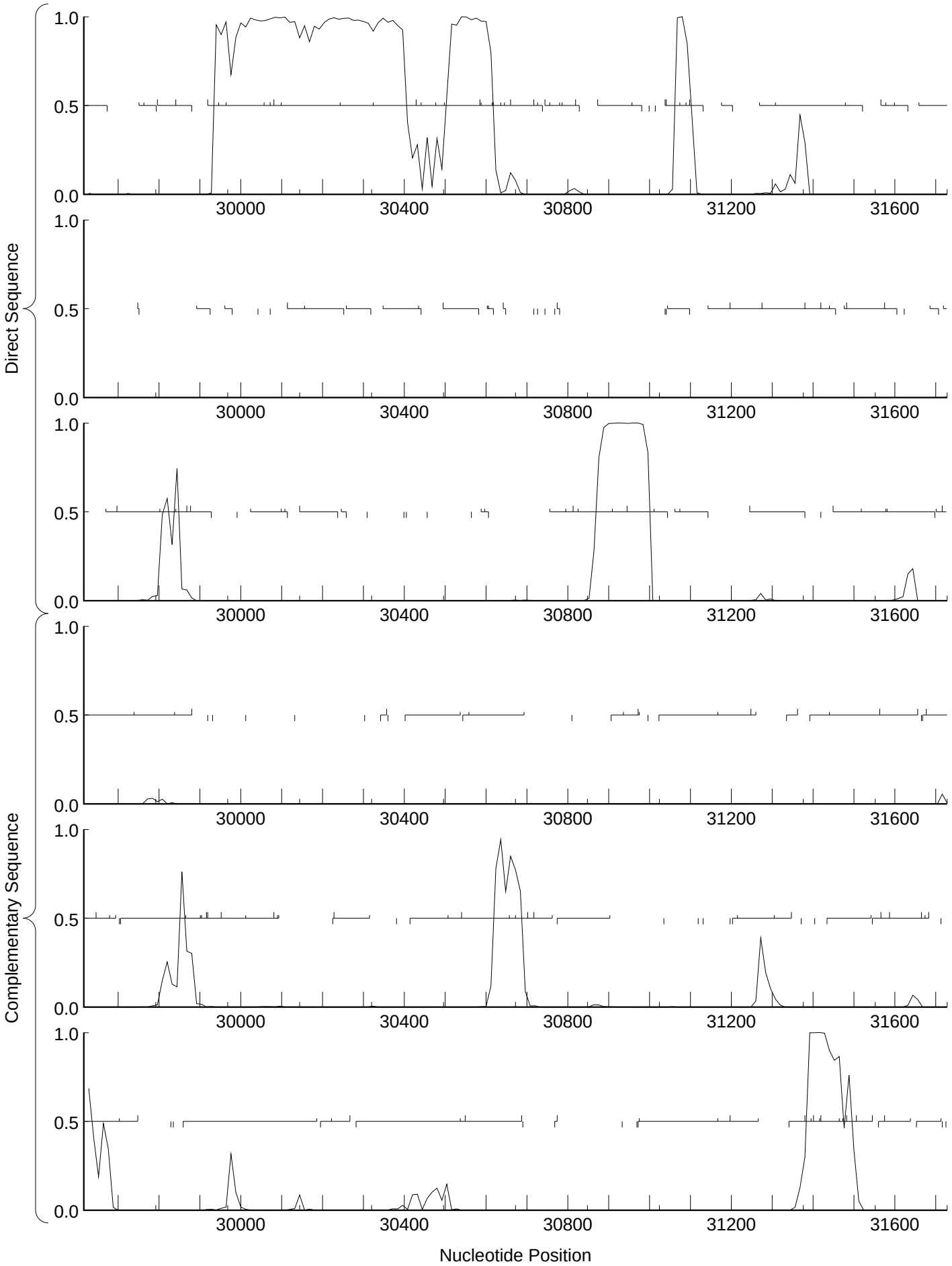
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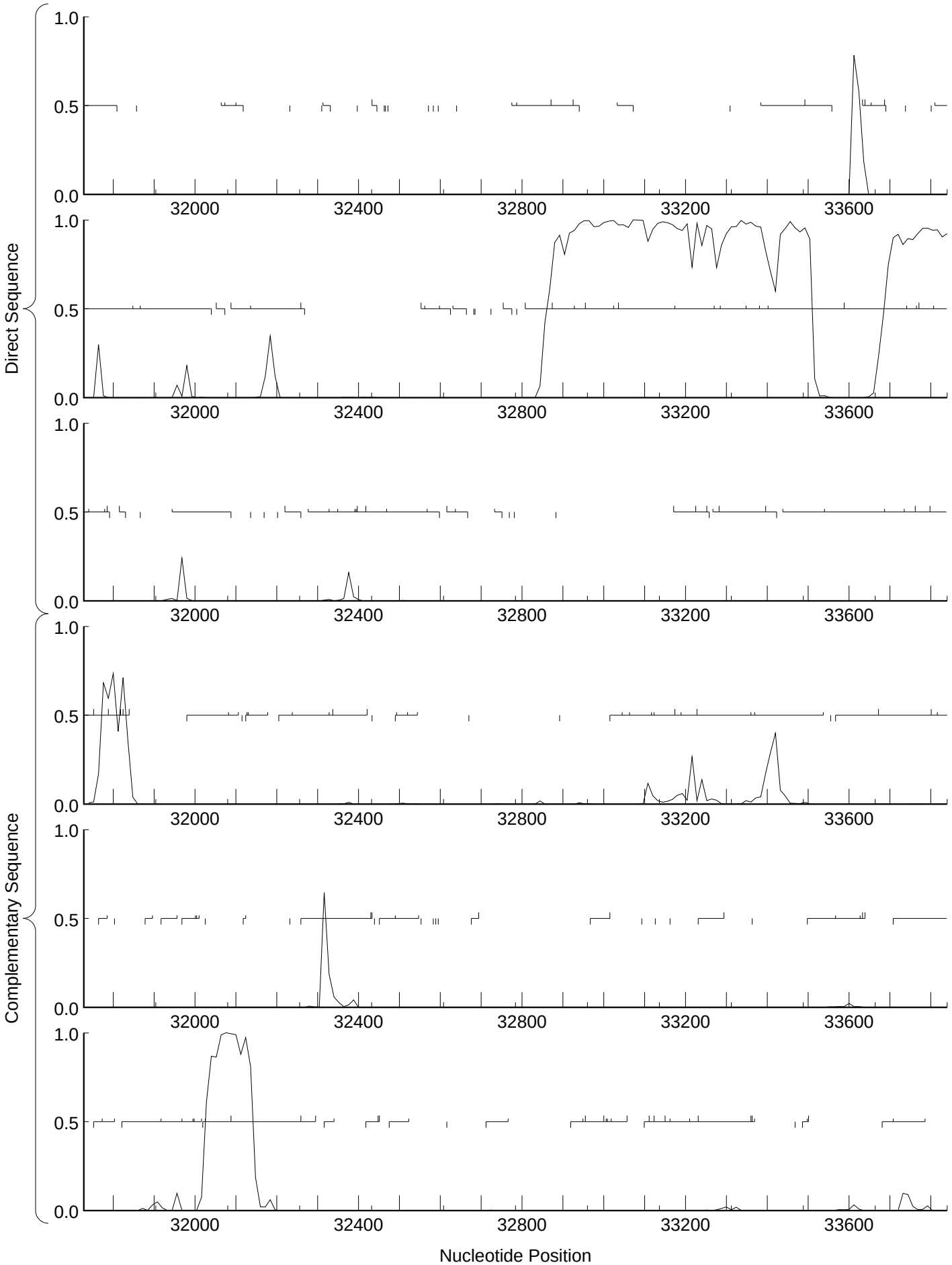
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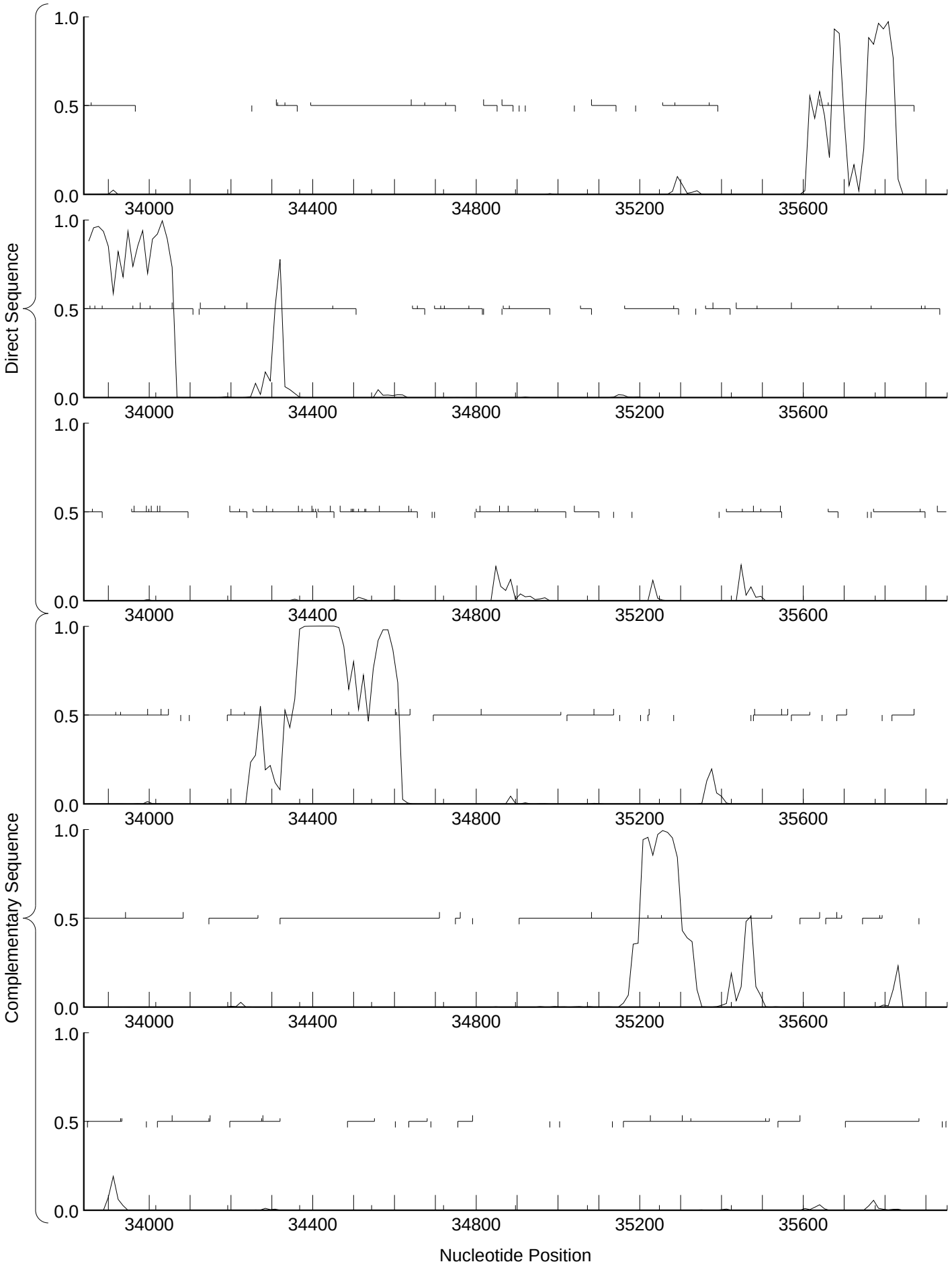
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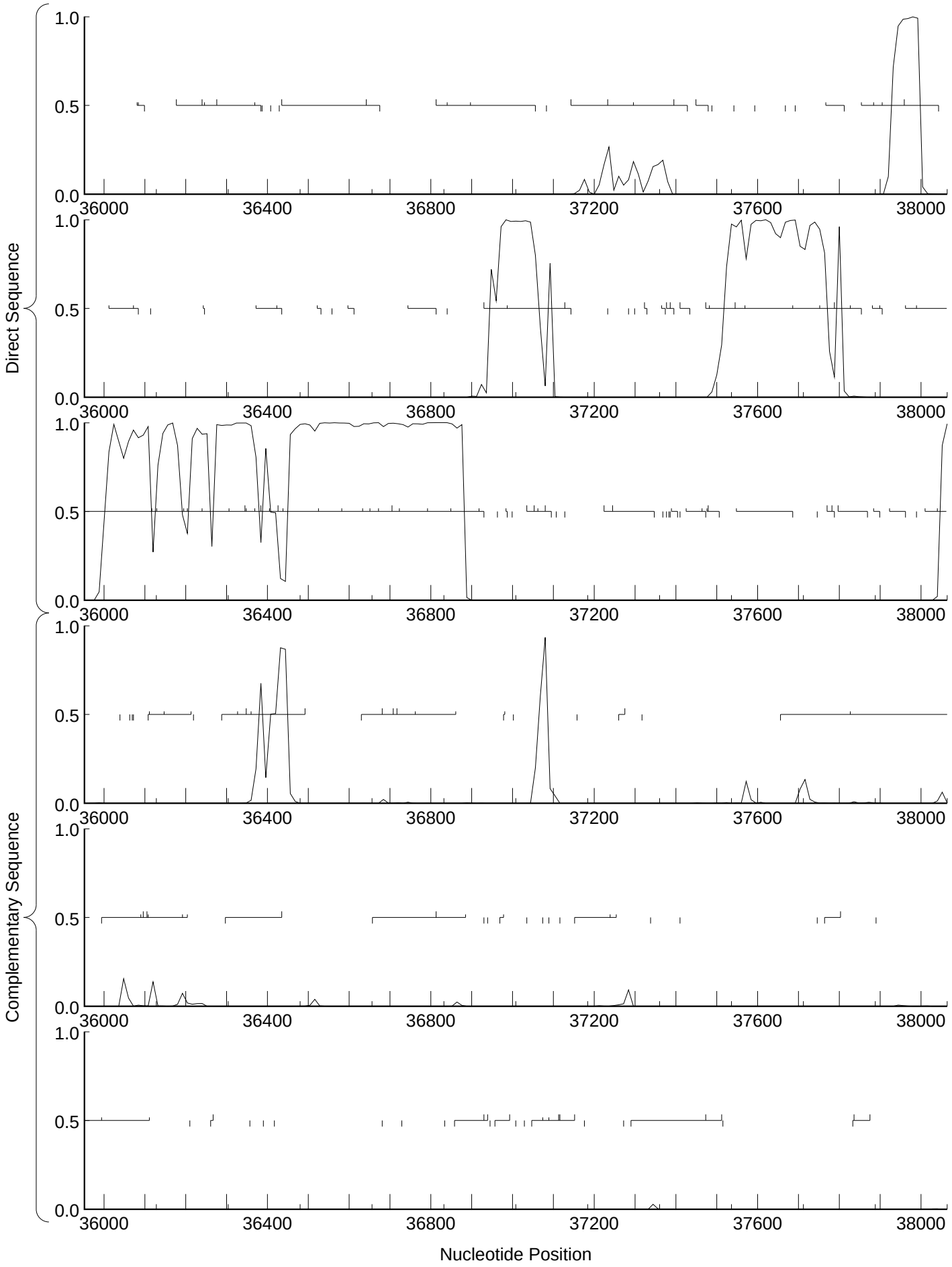
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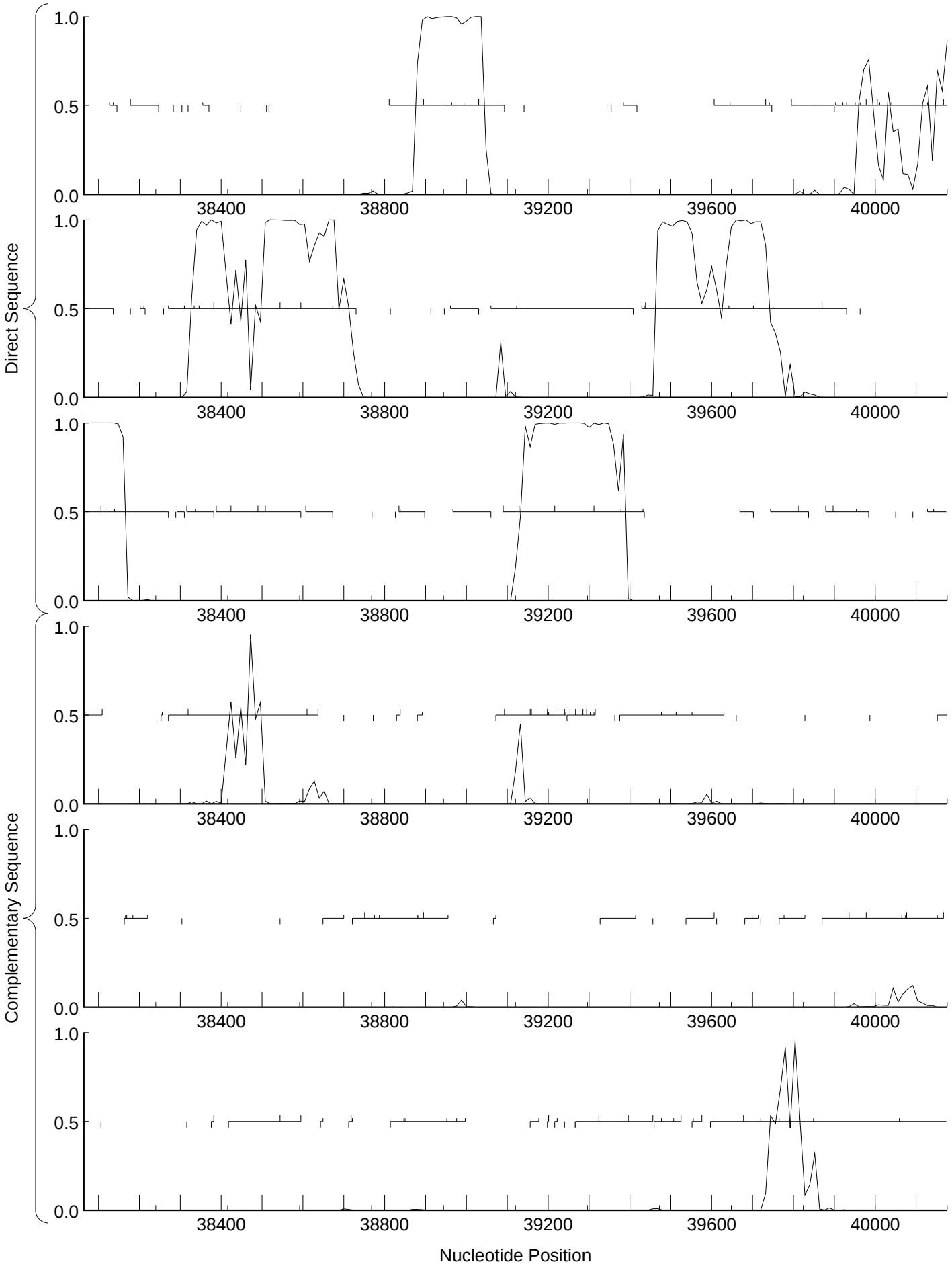
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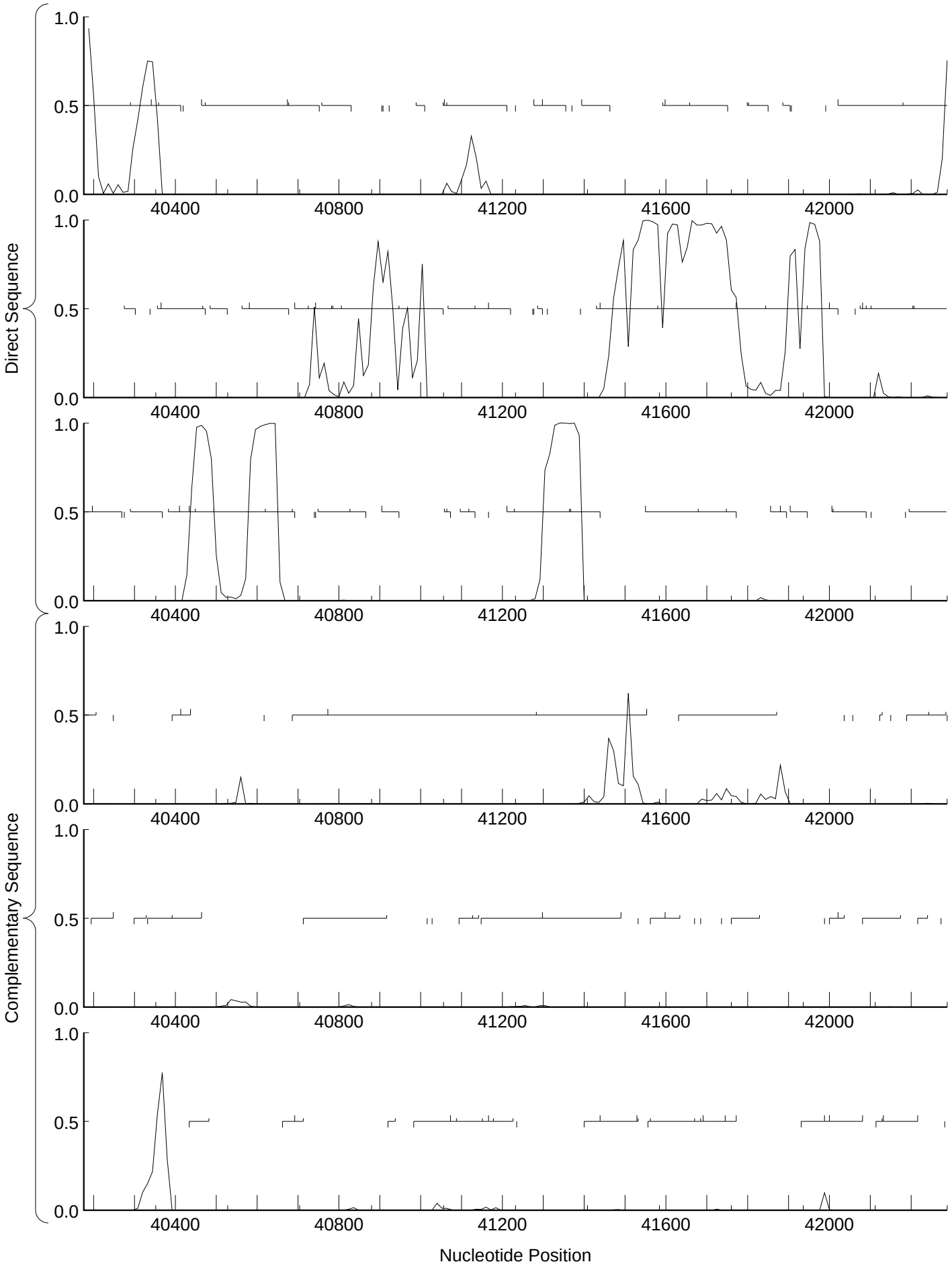
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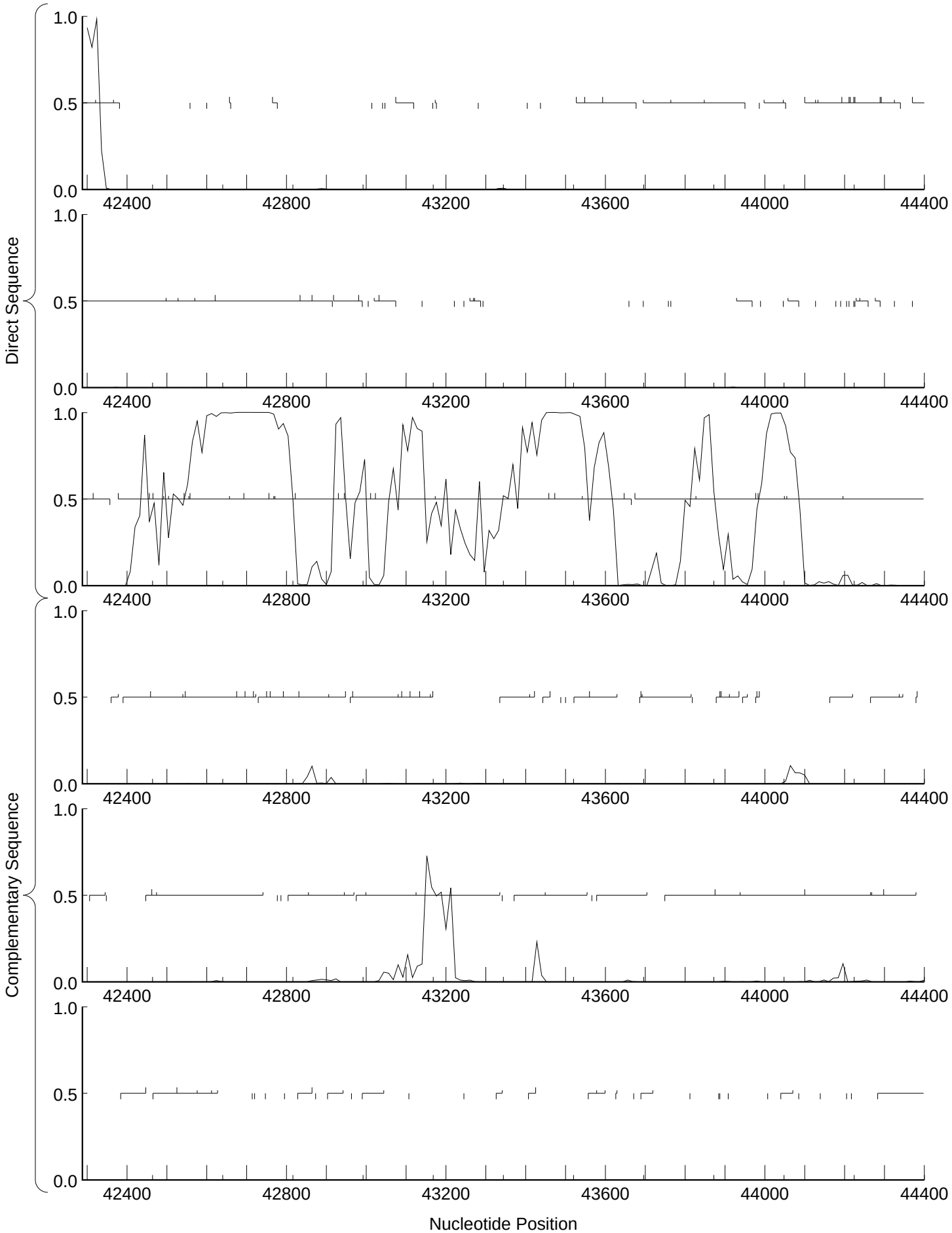
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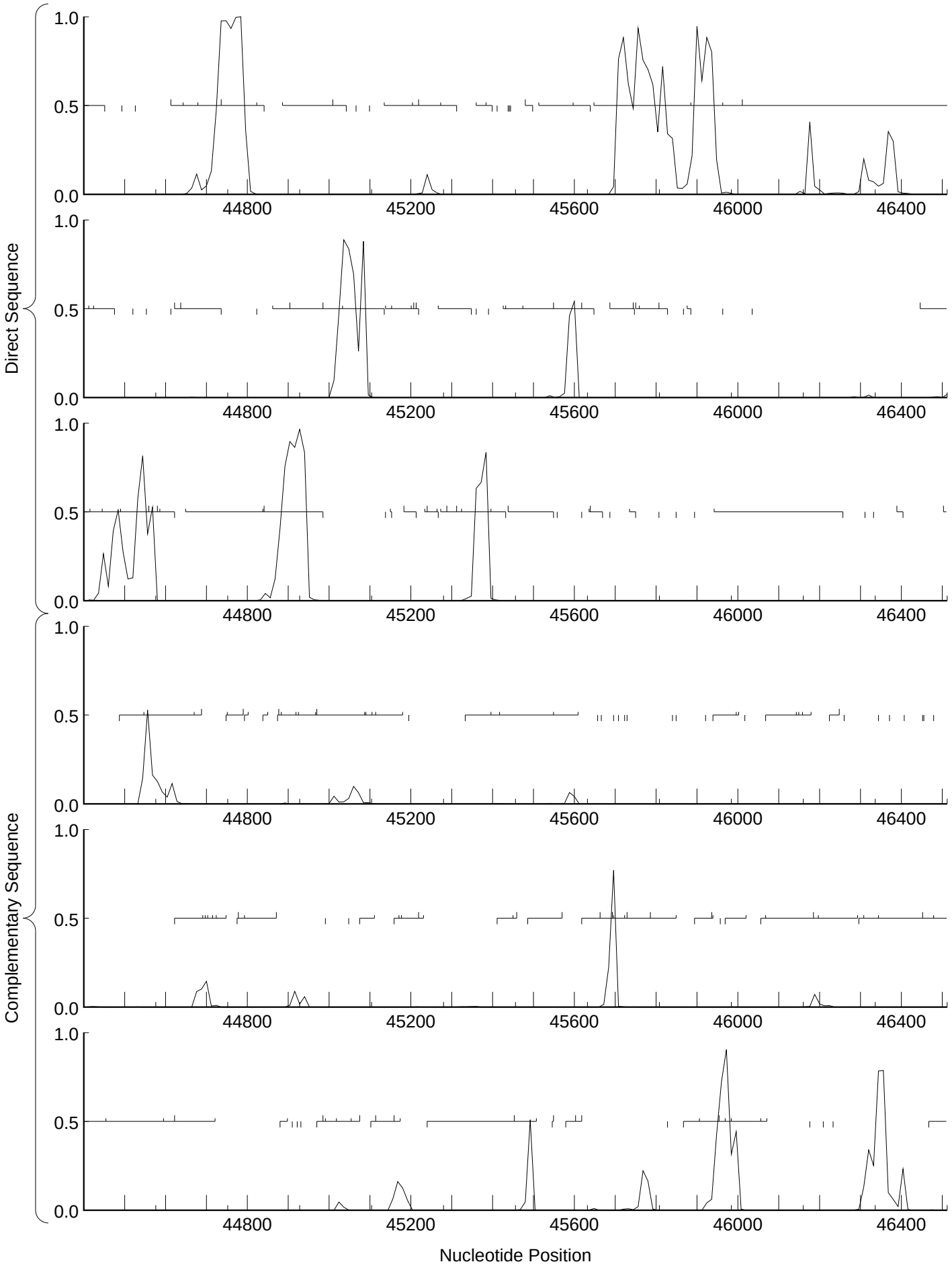
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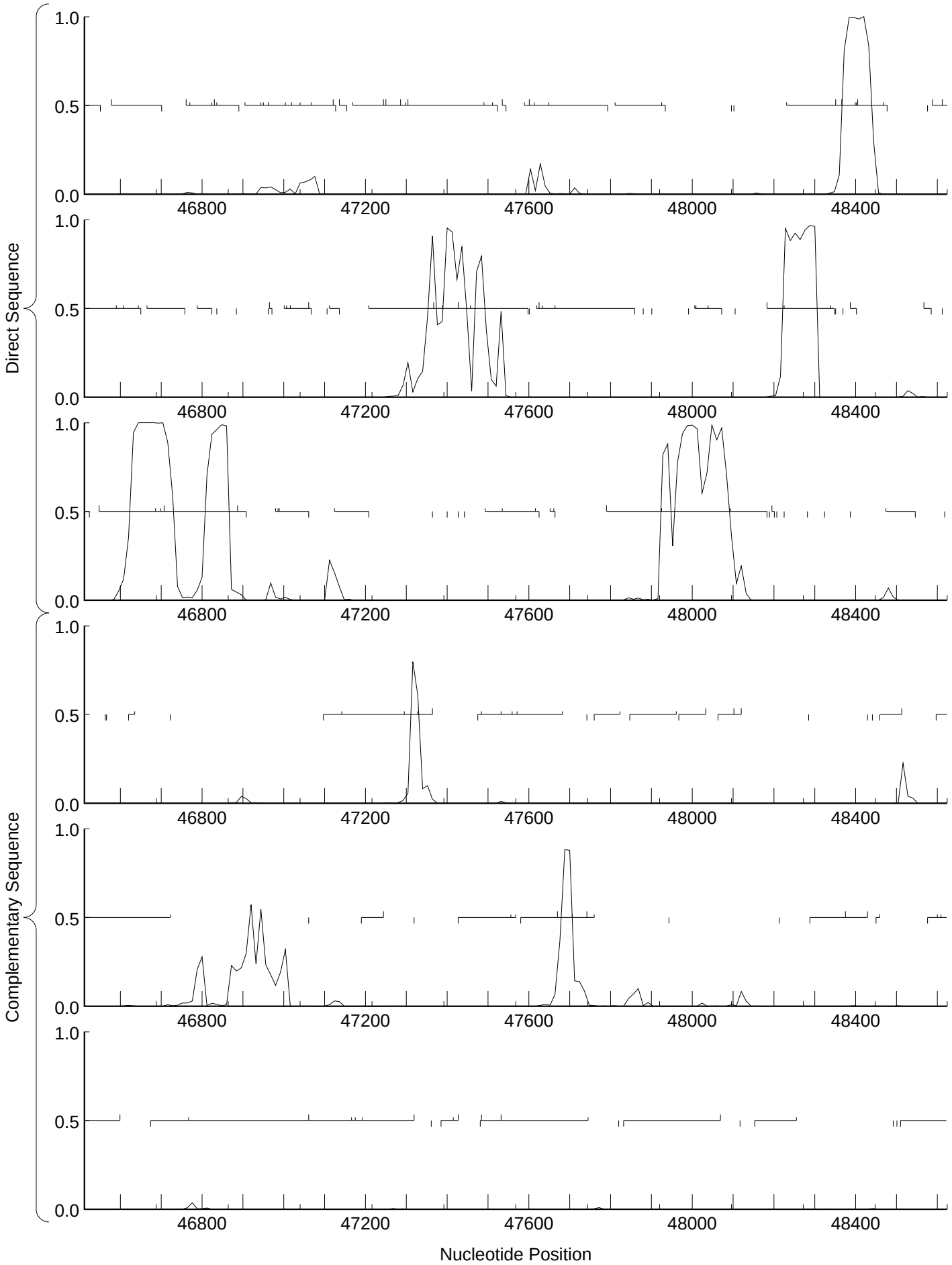
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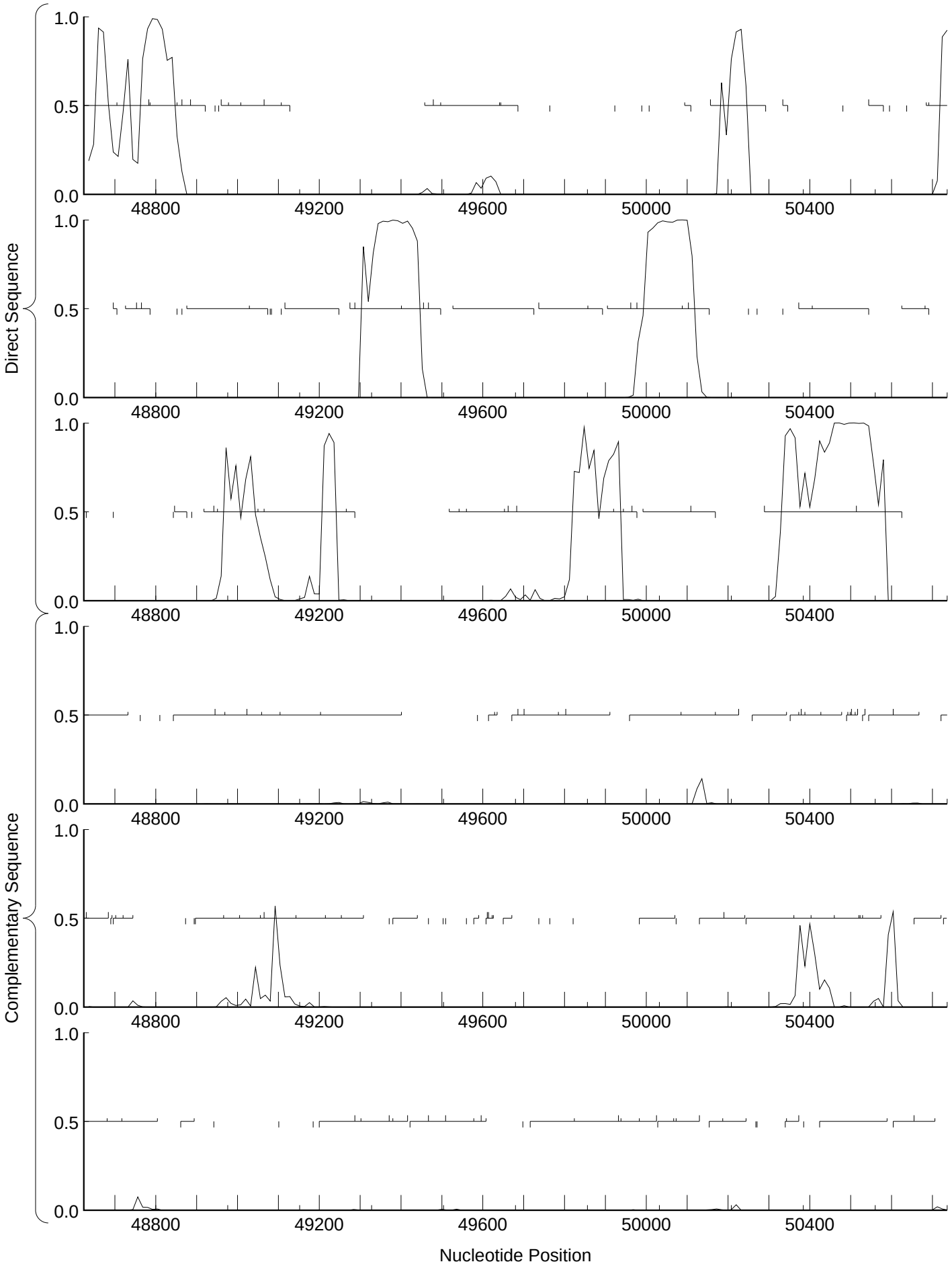
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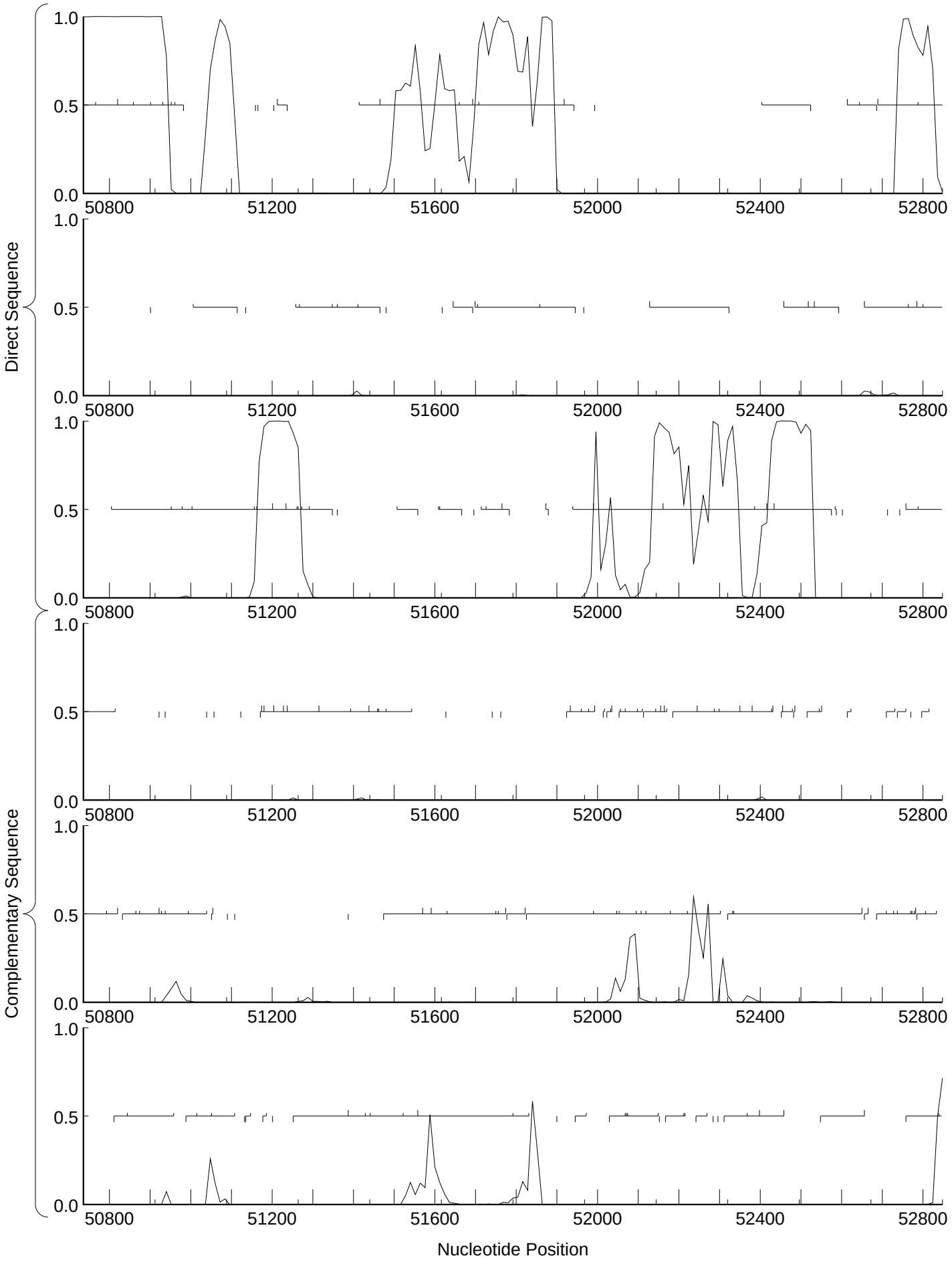
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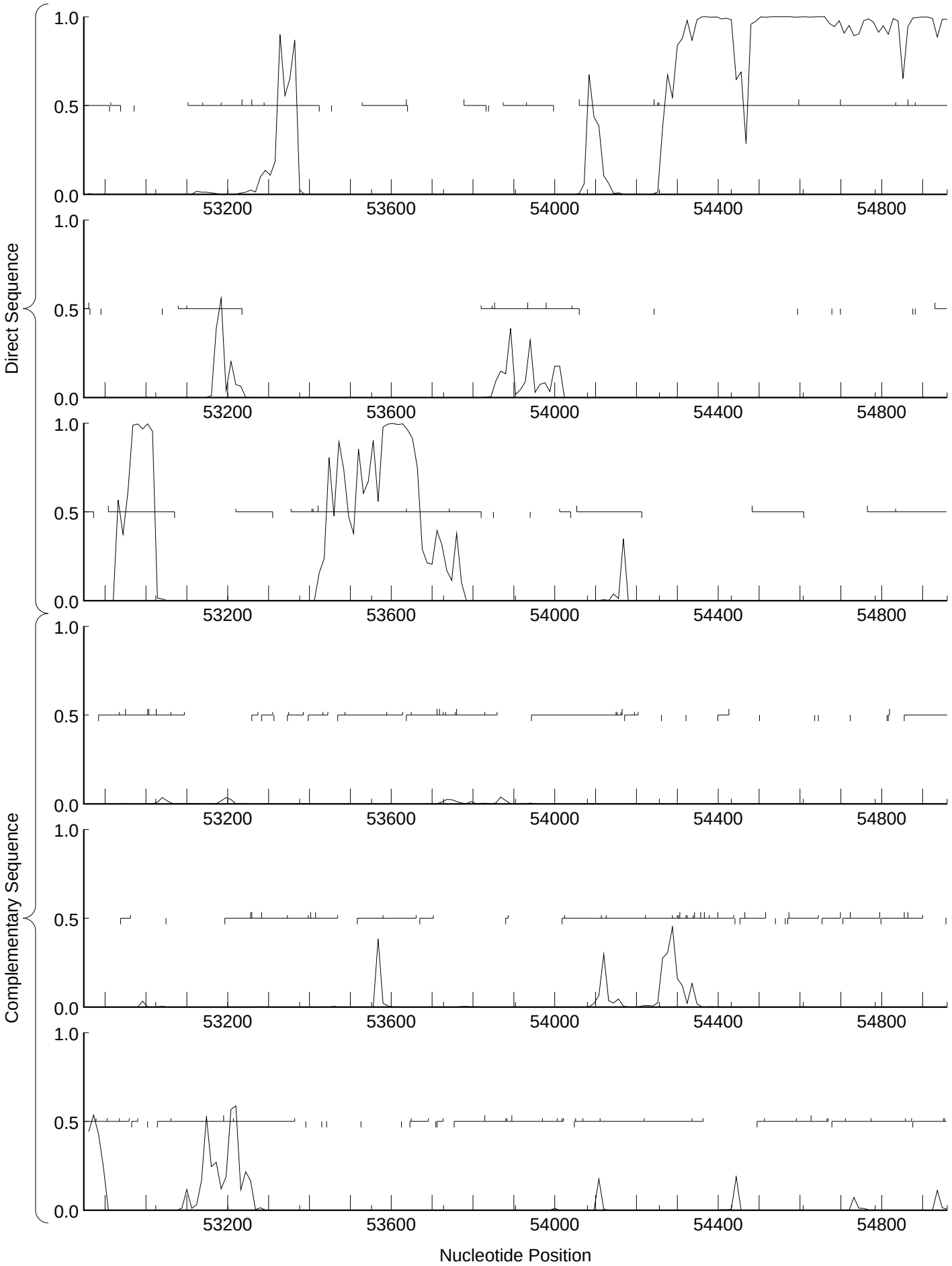
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t complete sequence, 56827 bp including 10-base 3’ overhang (CCGATGGCAT), Cluster F1, Order 4, Window 96, Step 12, 28/28

