GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage LilDestine complete sequence, 75440 bp including 10-base 3' overhang (T
Analysis Date    :  3/29/24 at 1:11:11
Pages            :  37
Sequence Length  :  75440 bp
GC Content       :   58.95%

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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                    List regions and/or splice sites,
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MATRIX INFORMATION

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Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

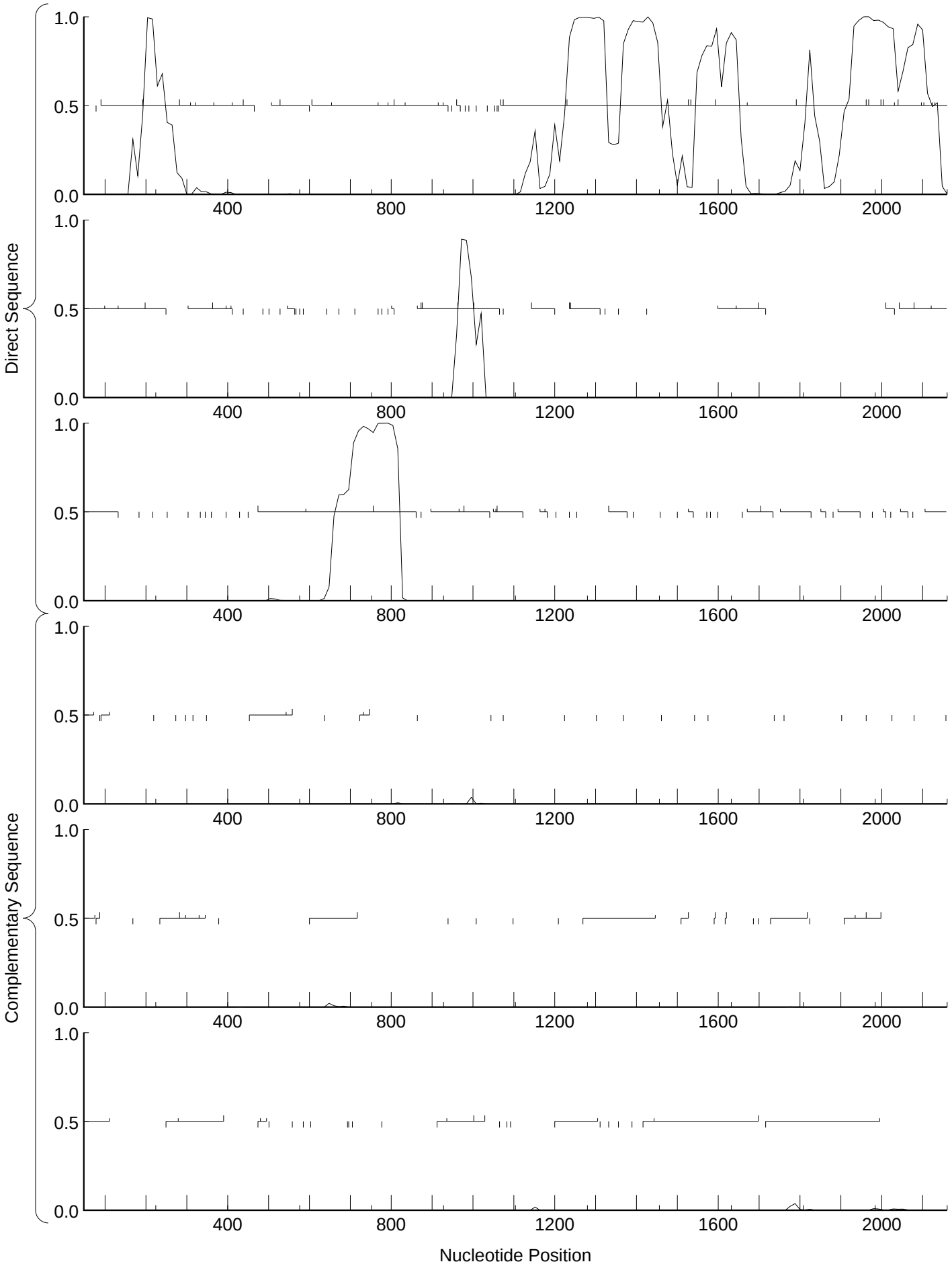
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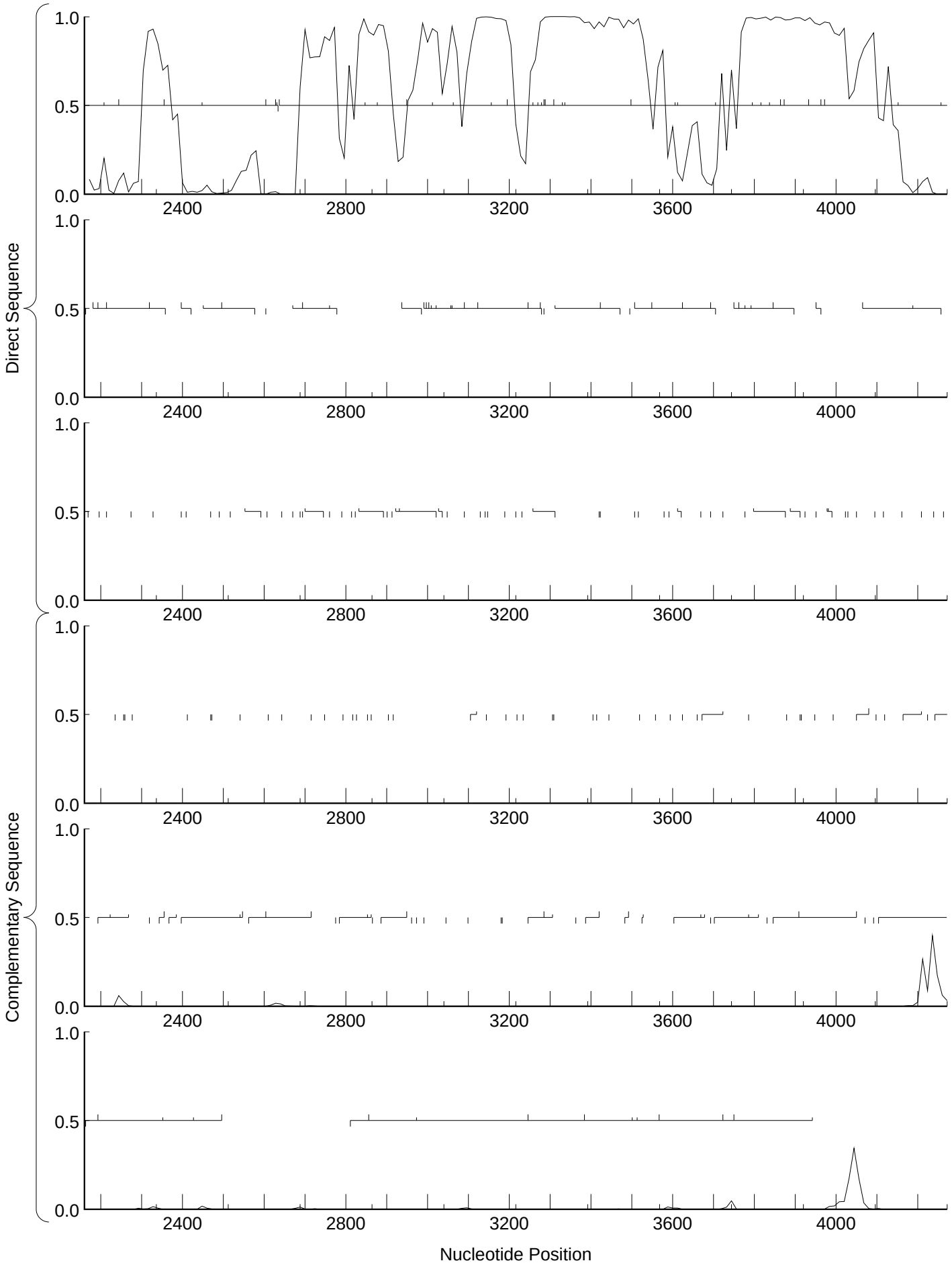

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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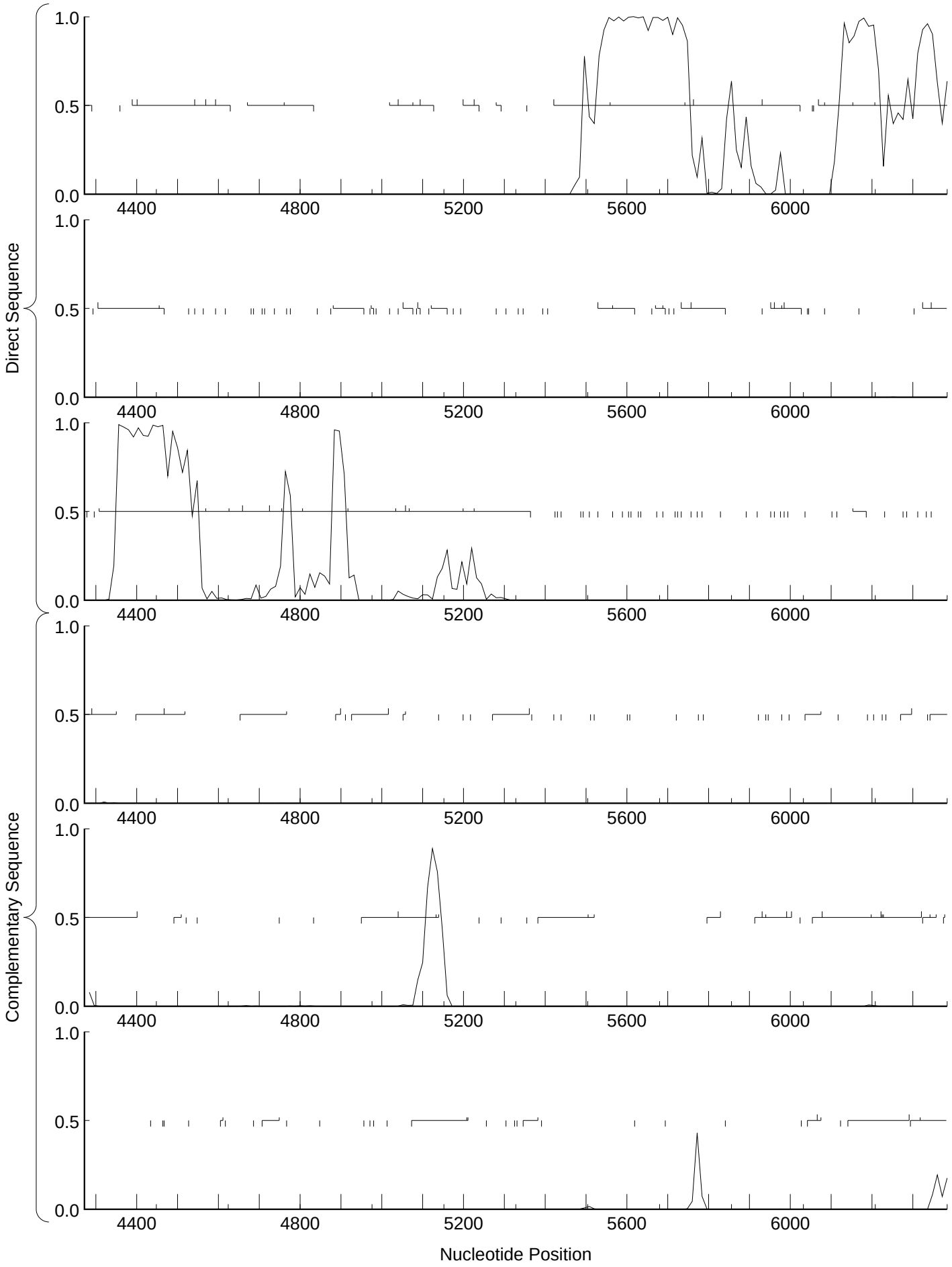
Matrix notes & comments

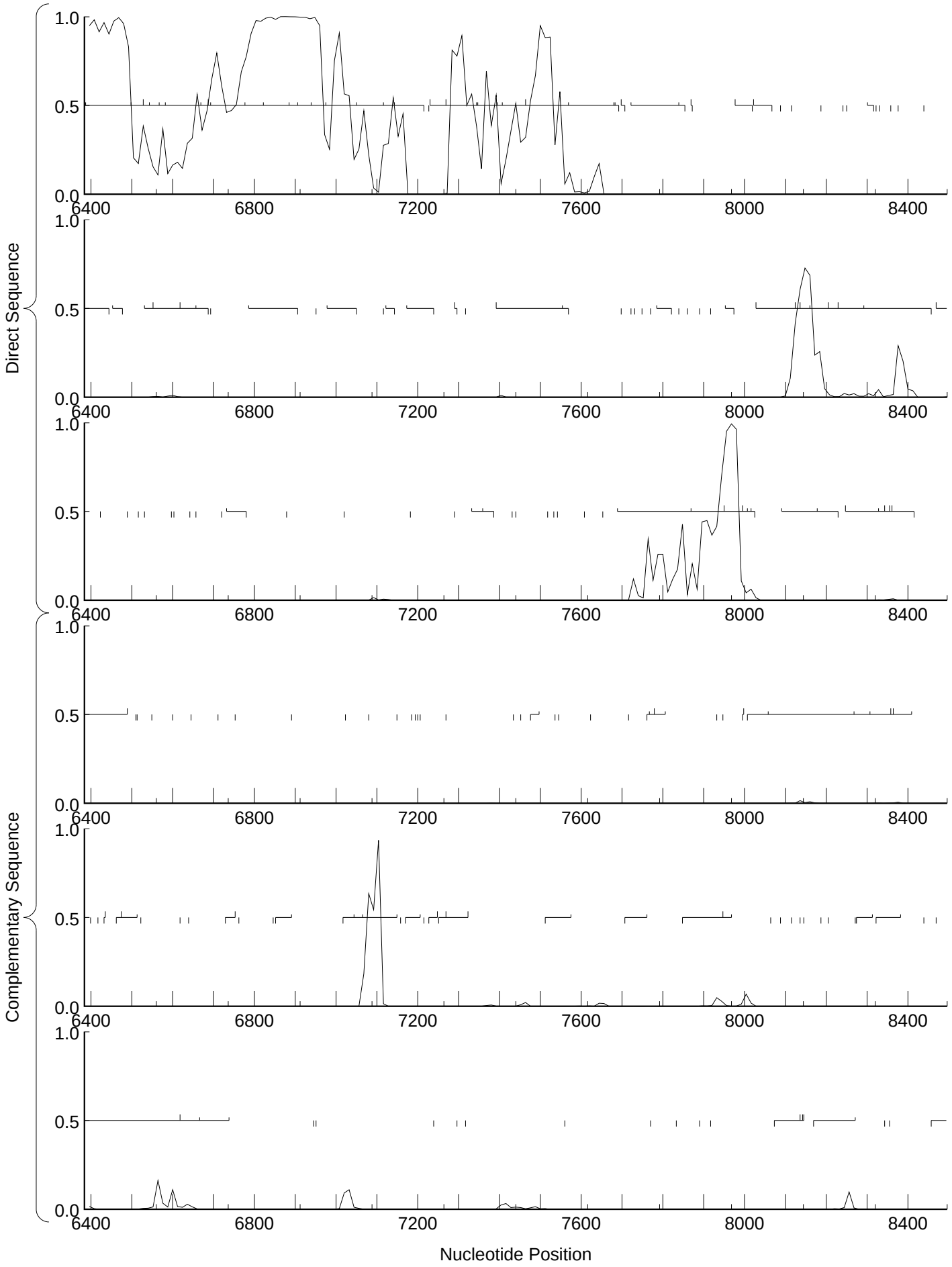
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

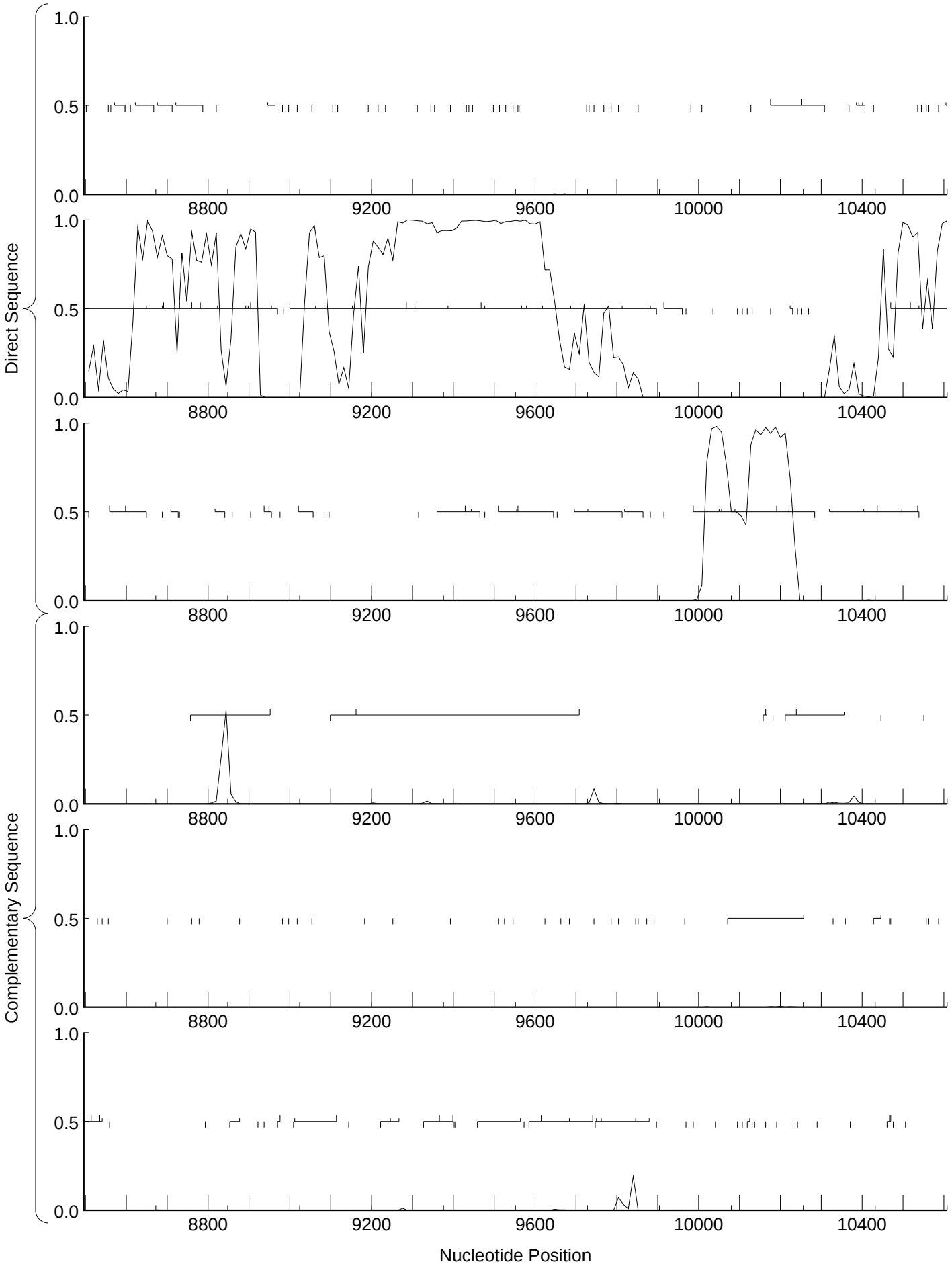
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

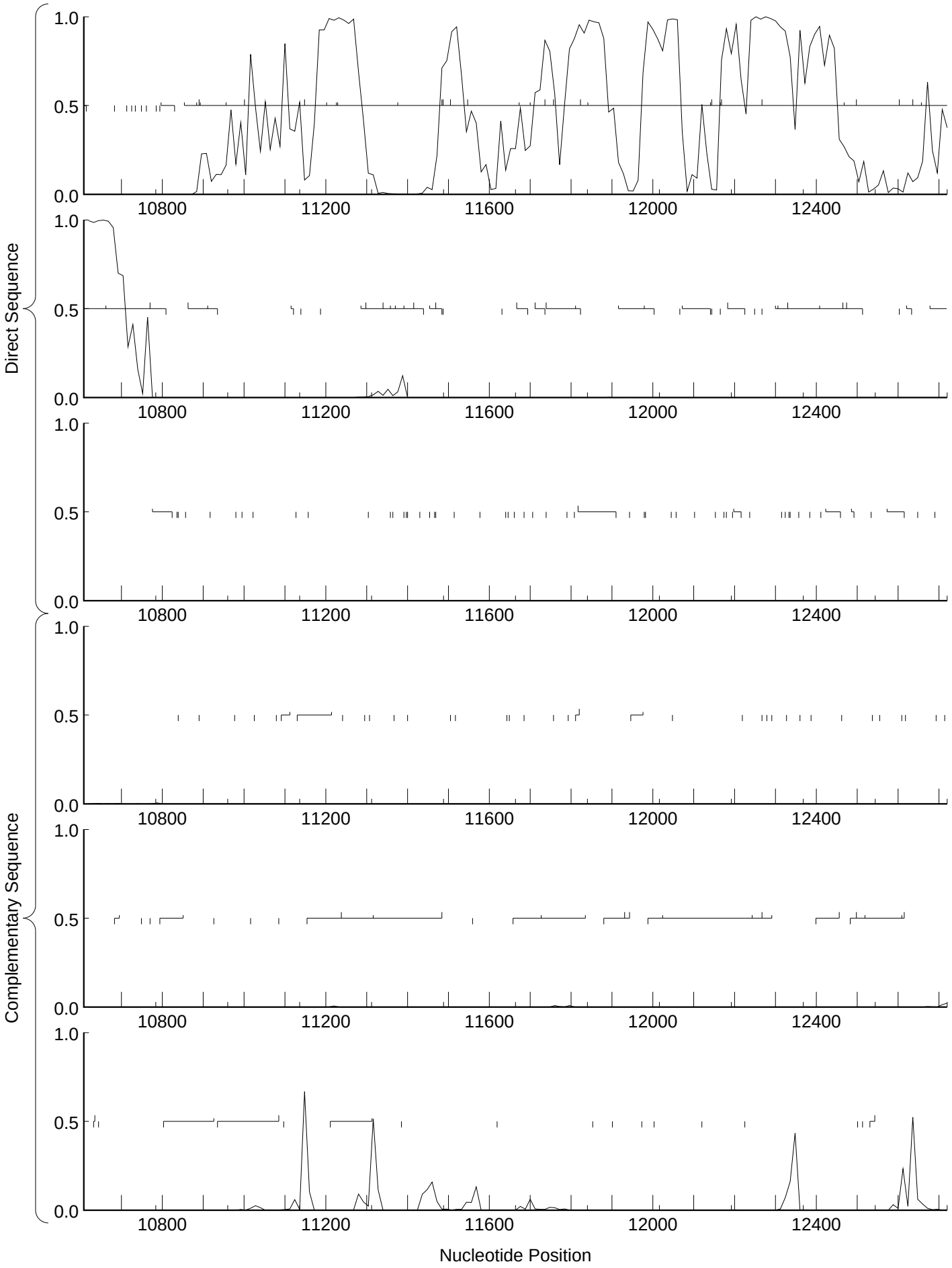

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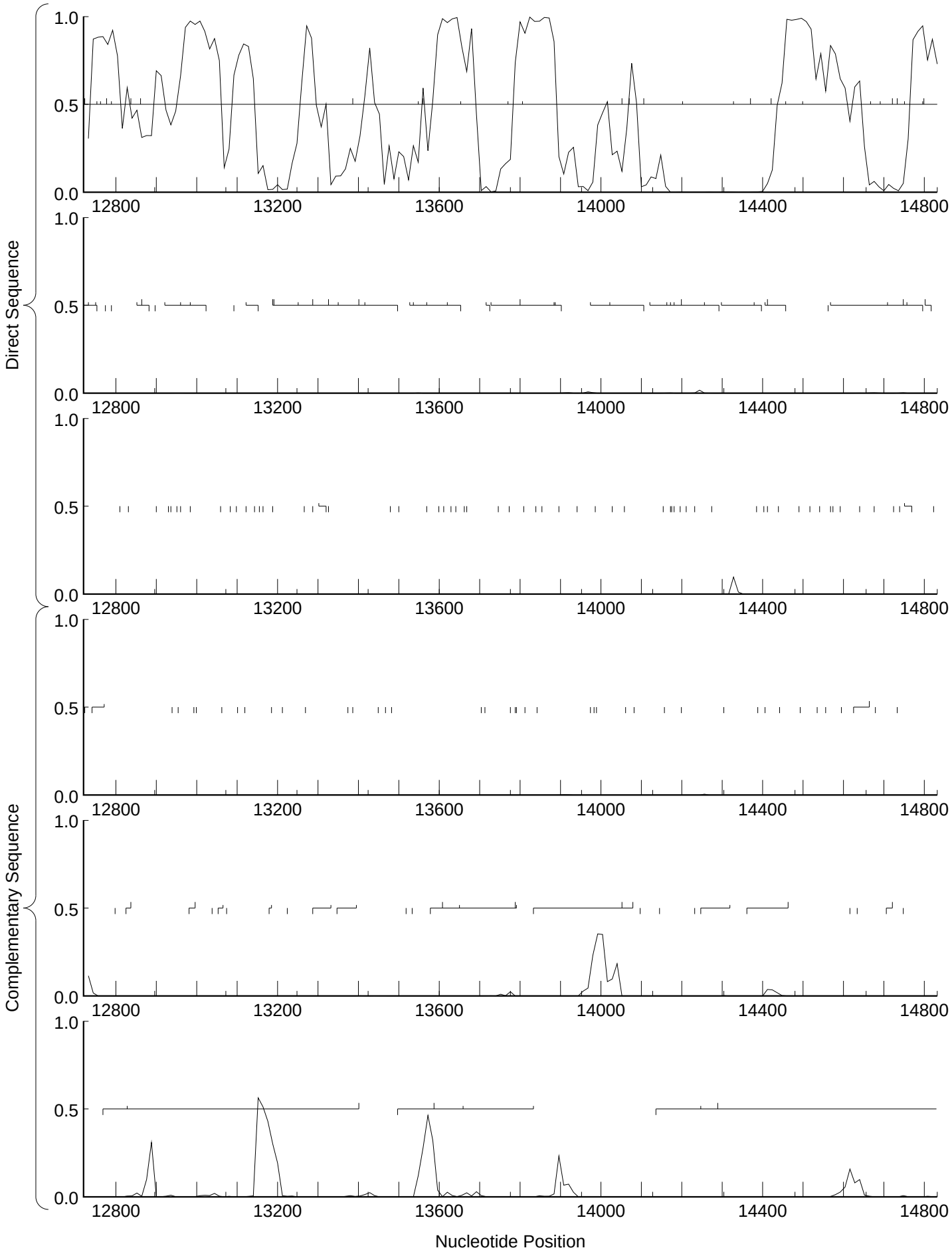

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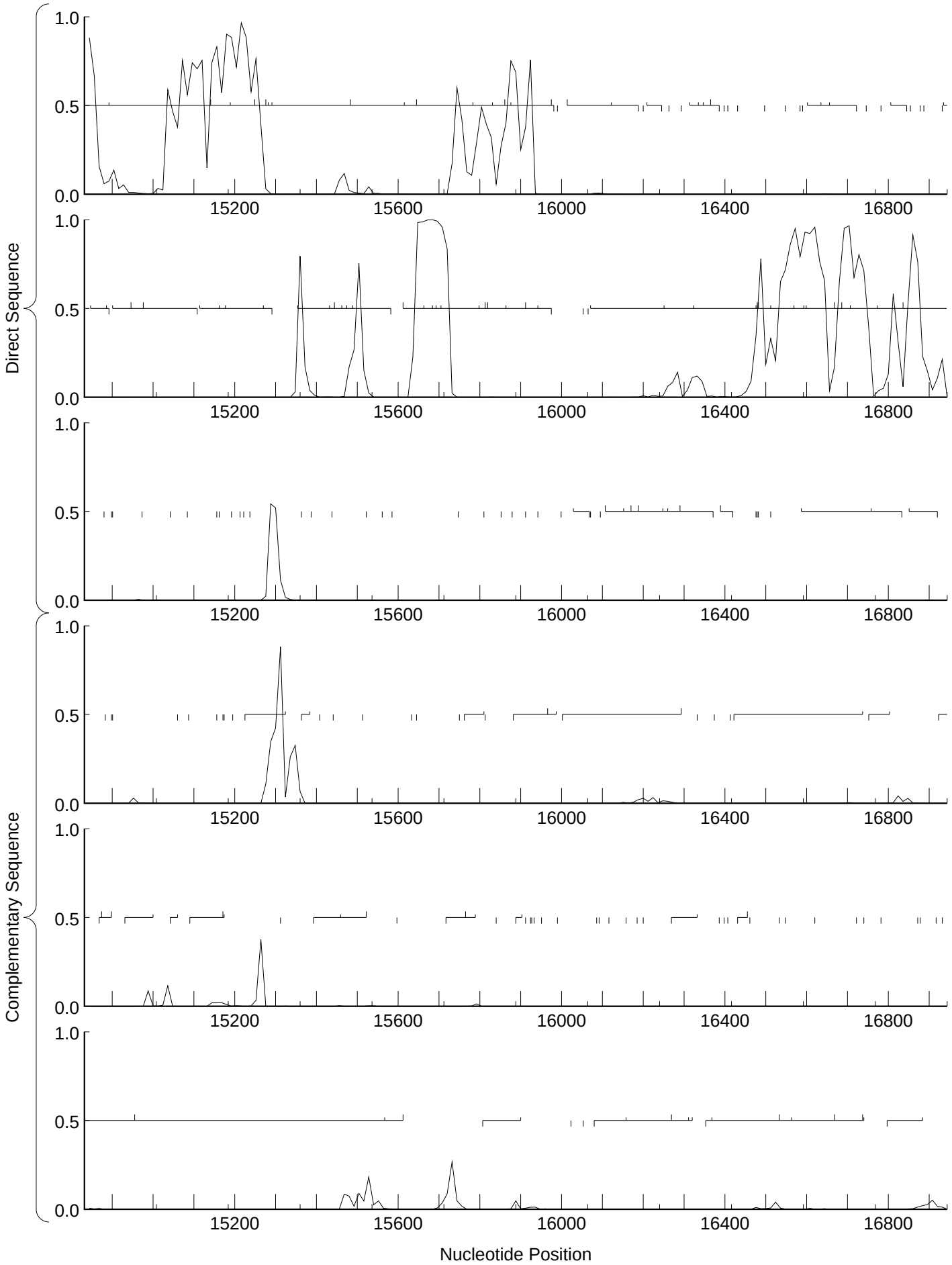

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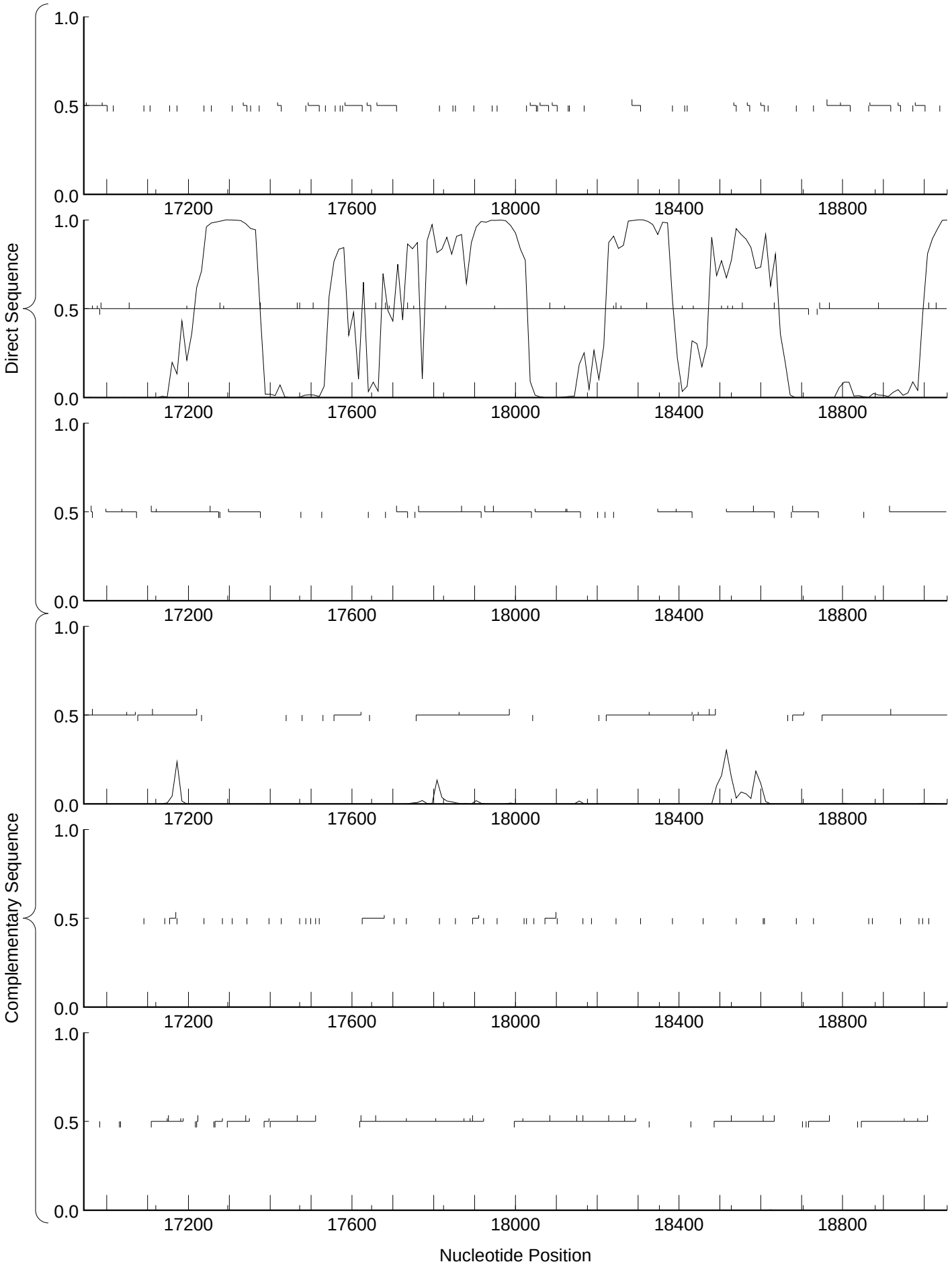

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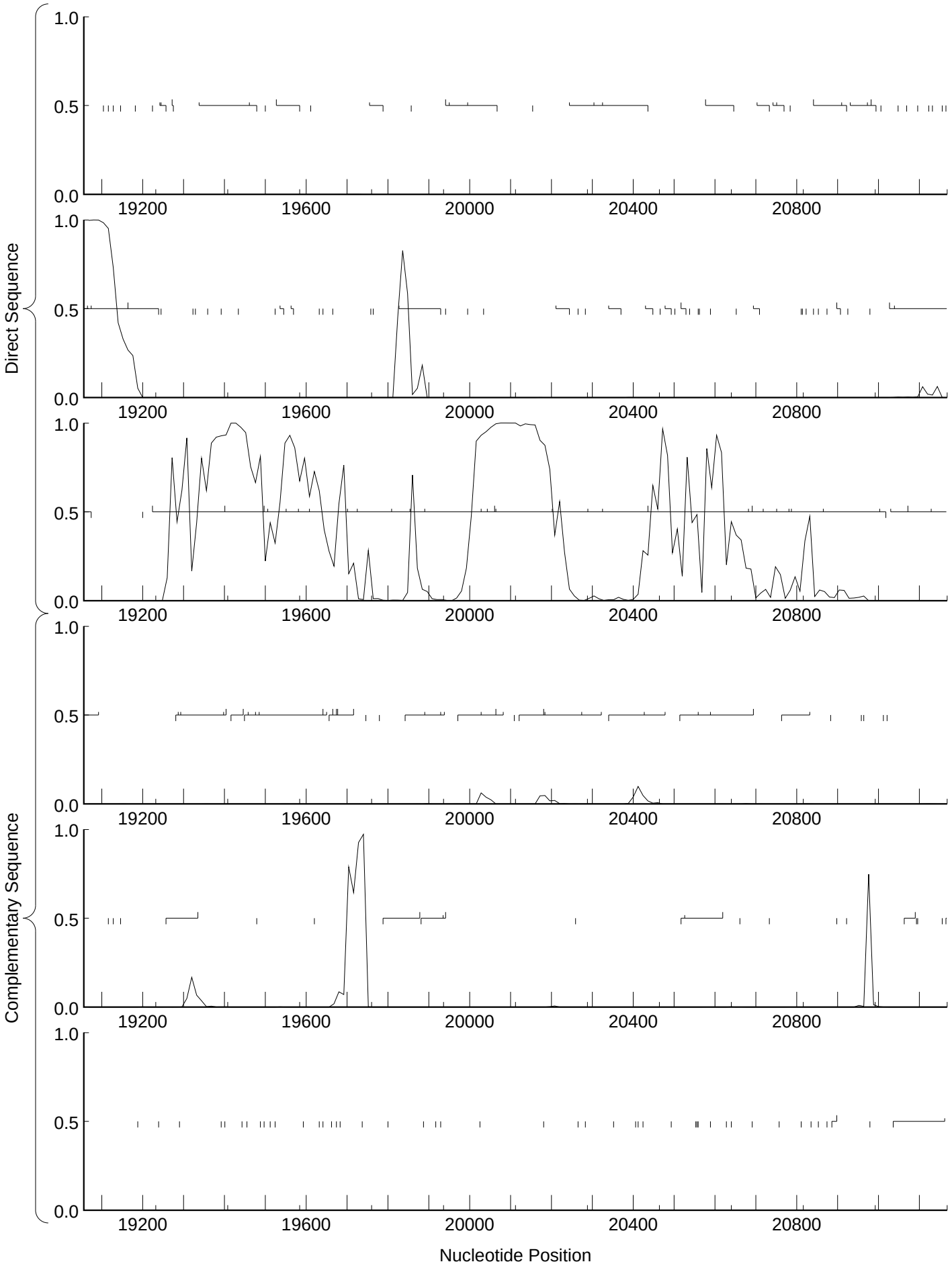

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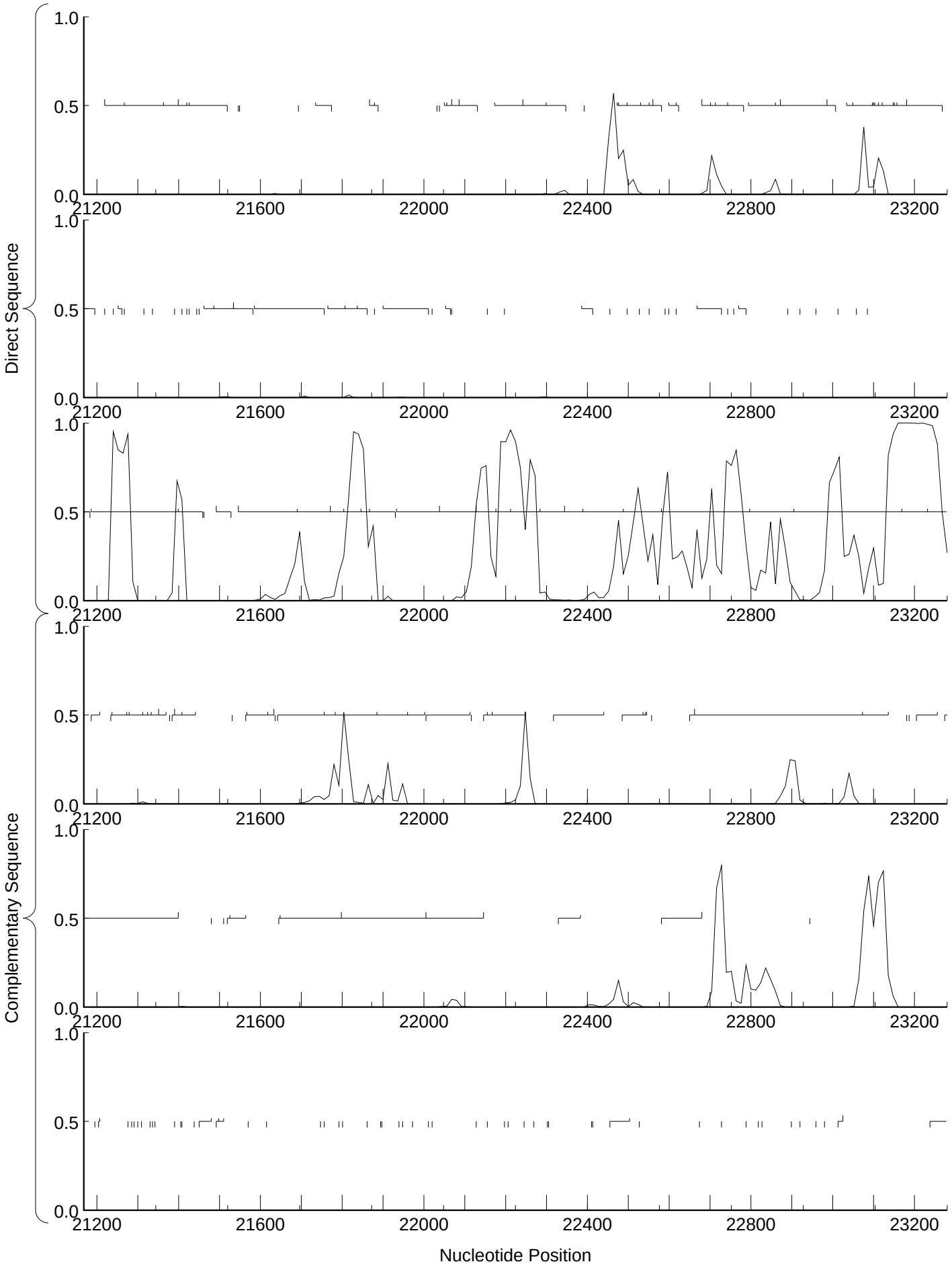
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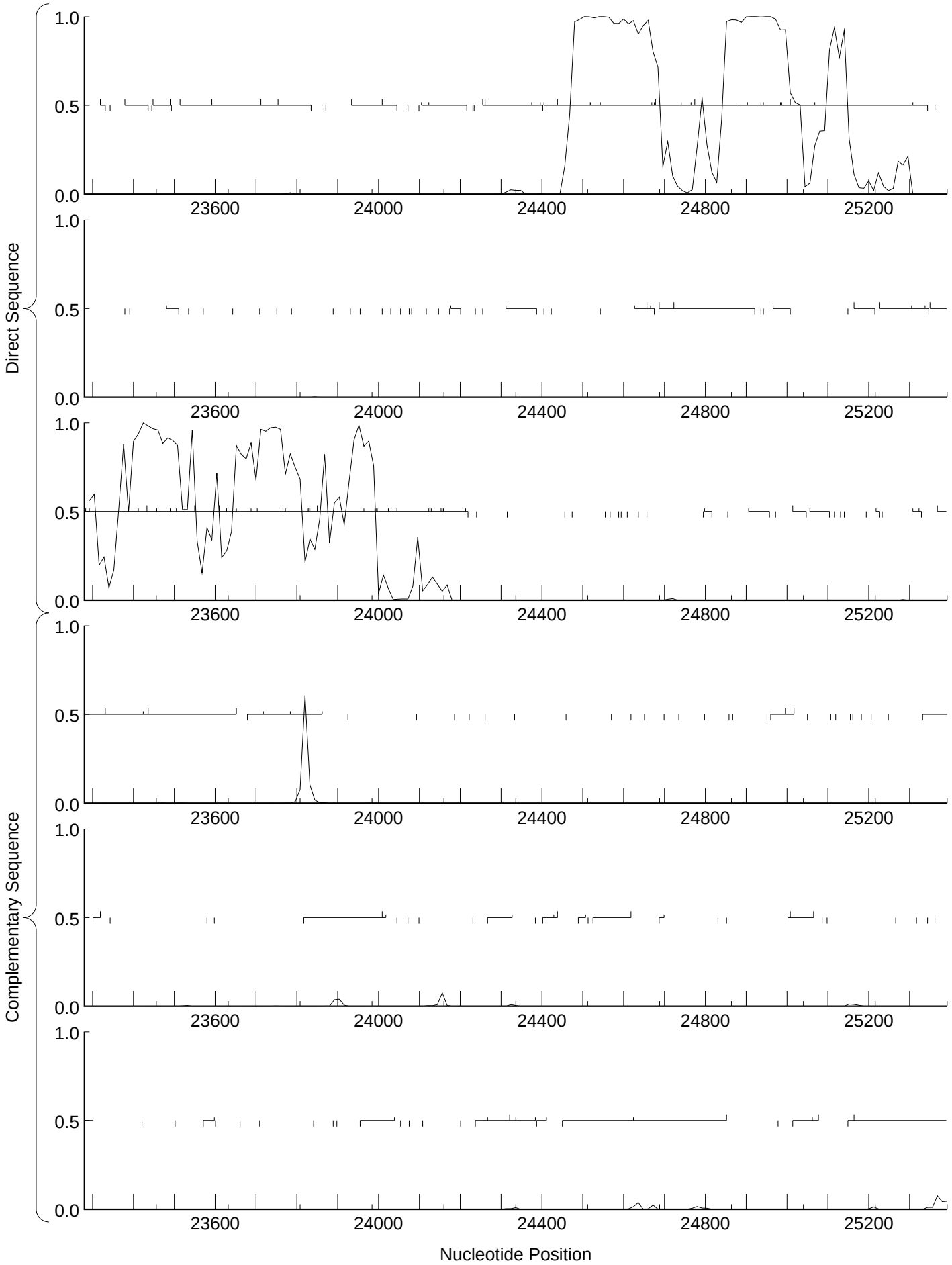
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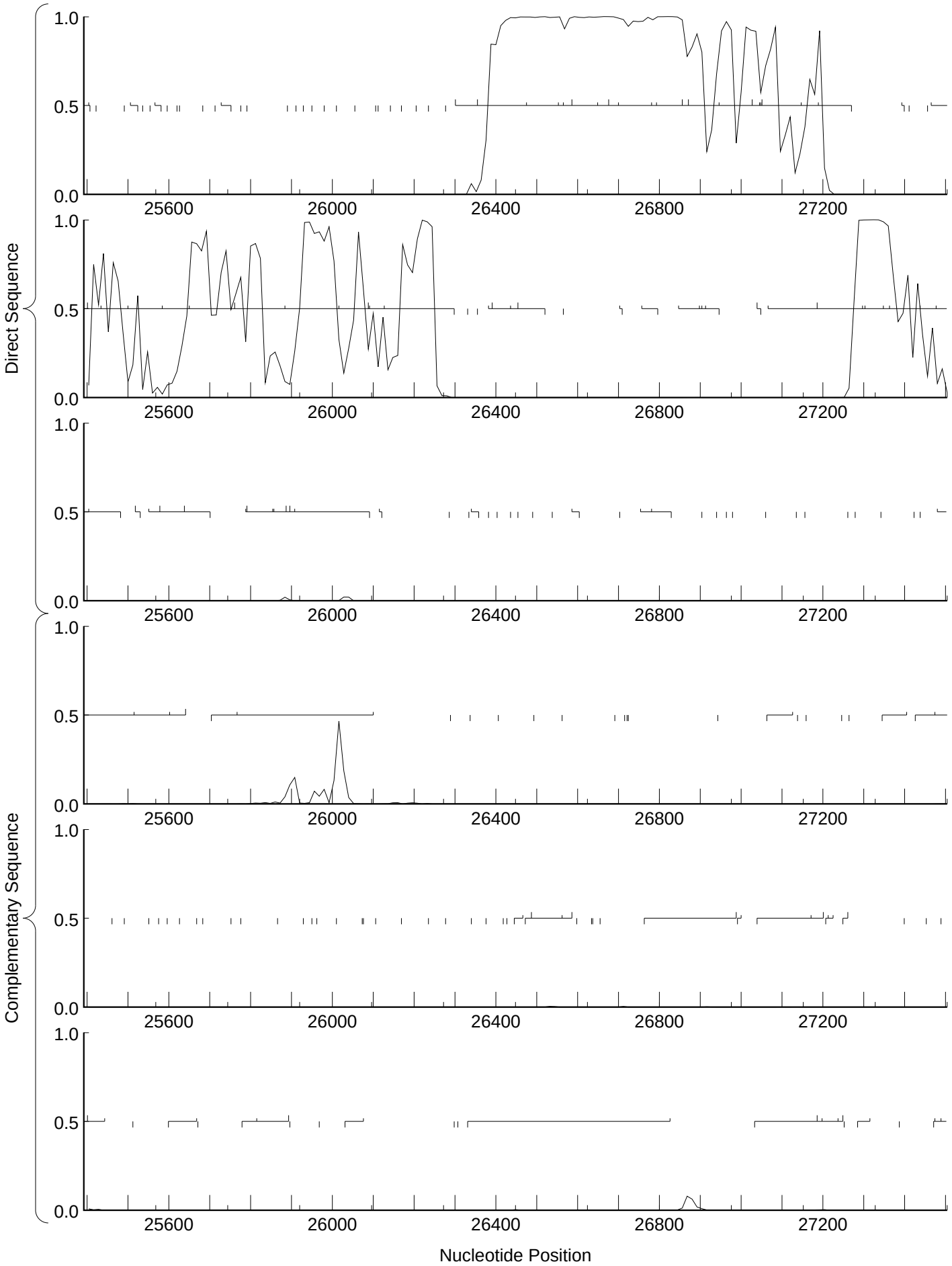
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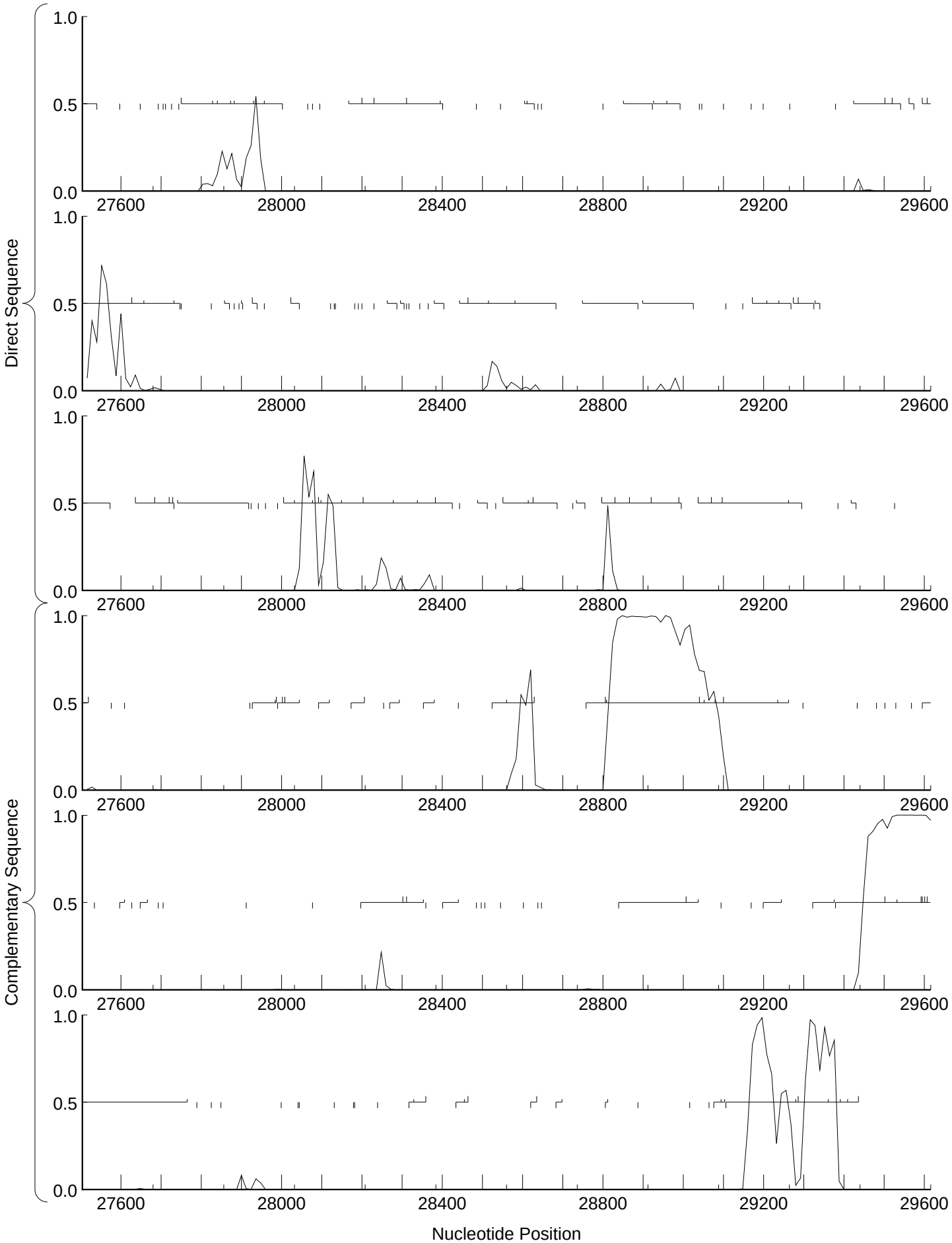
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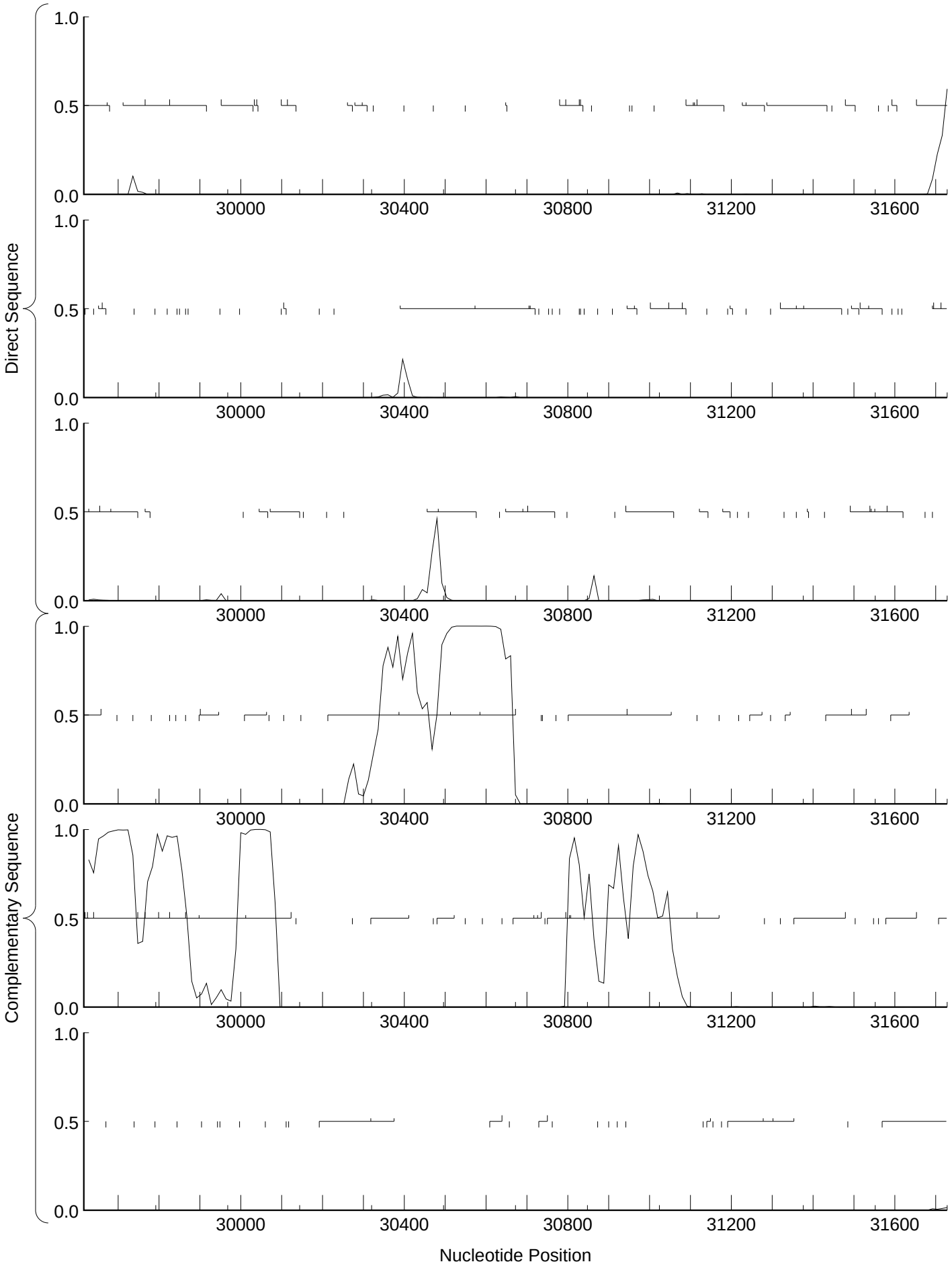
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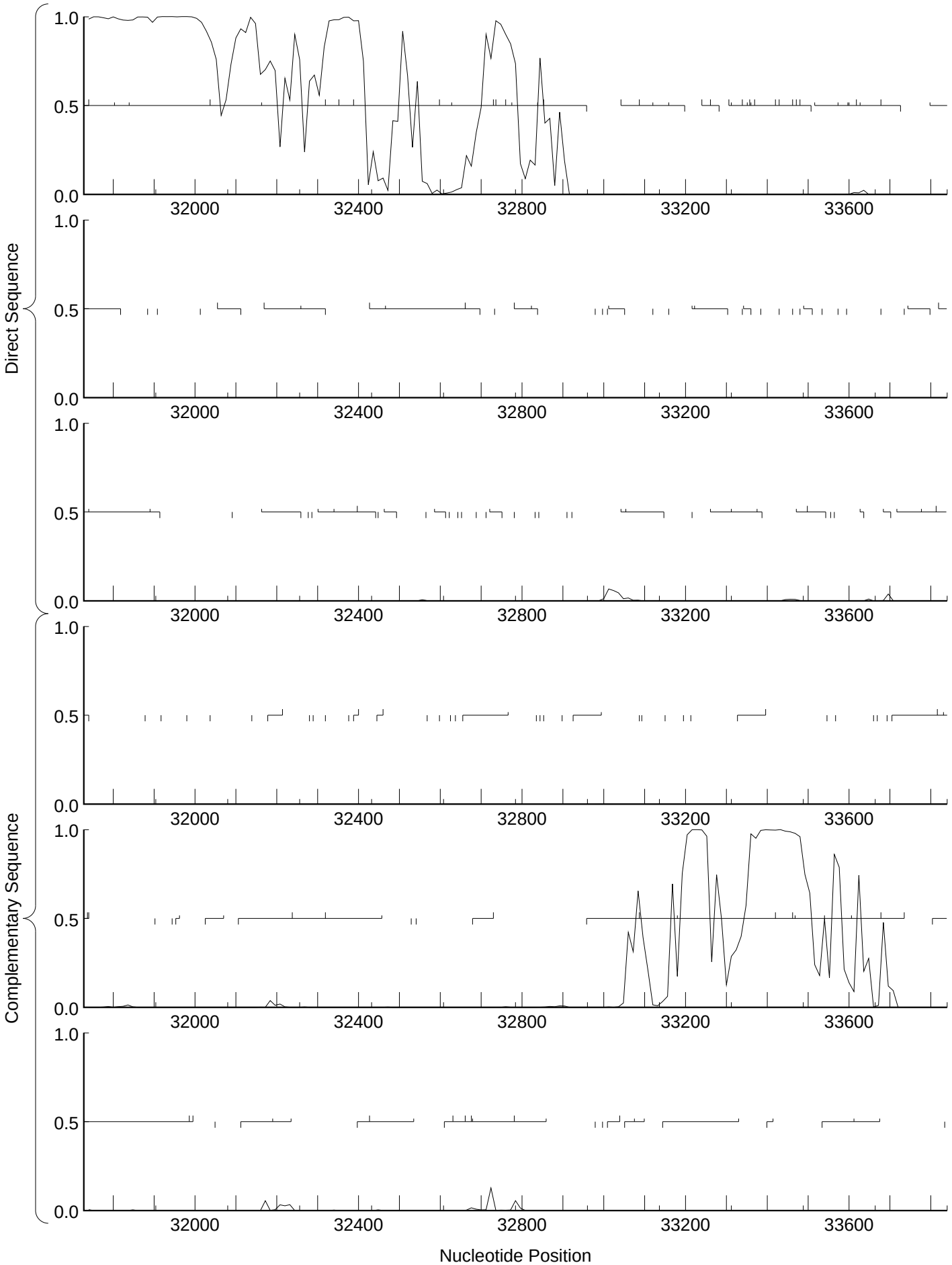
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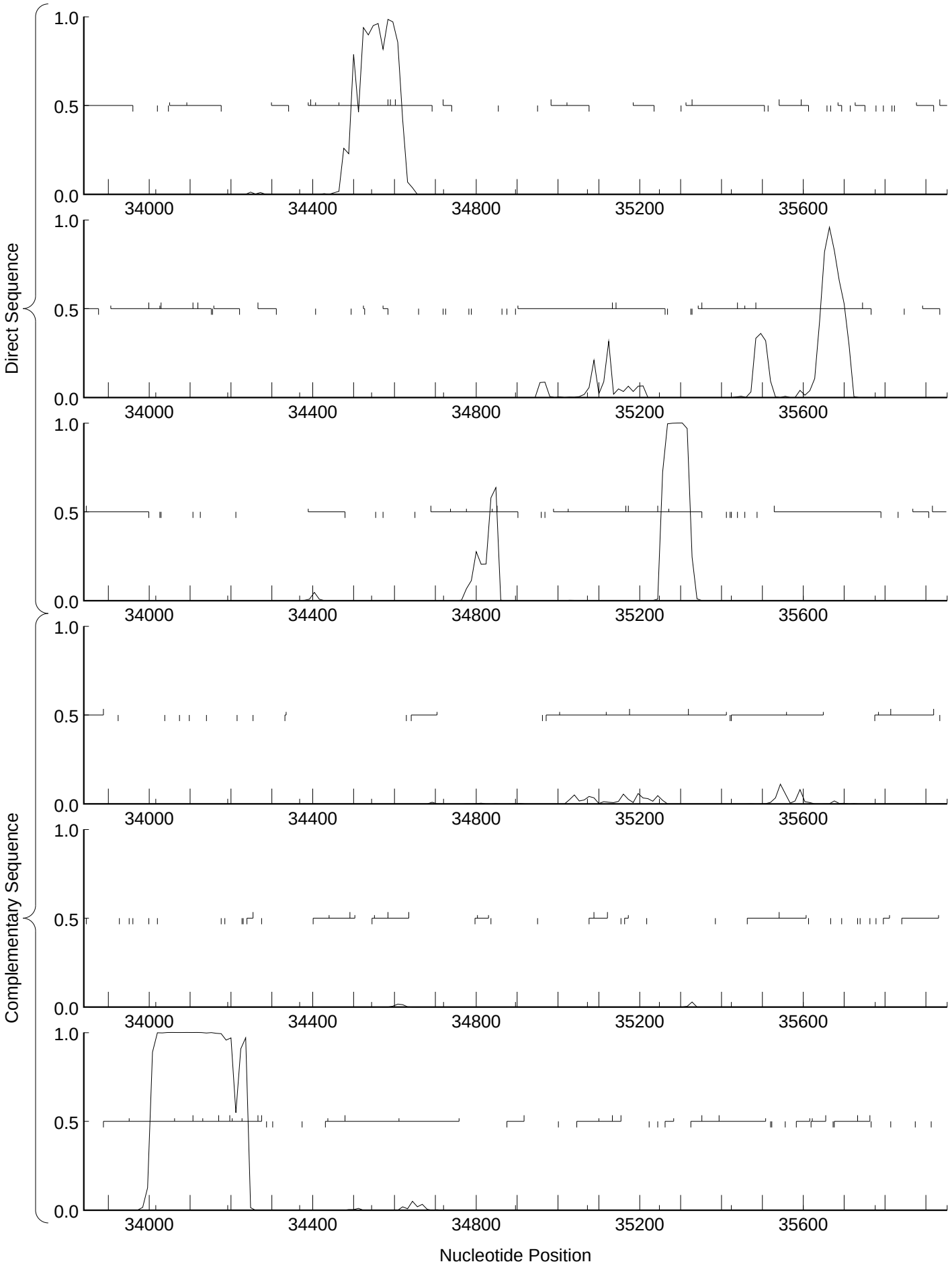
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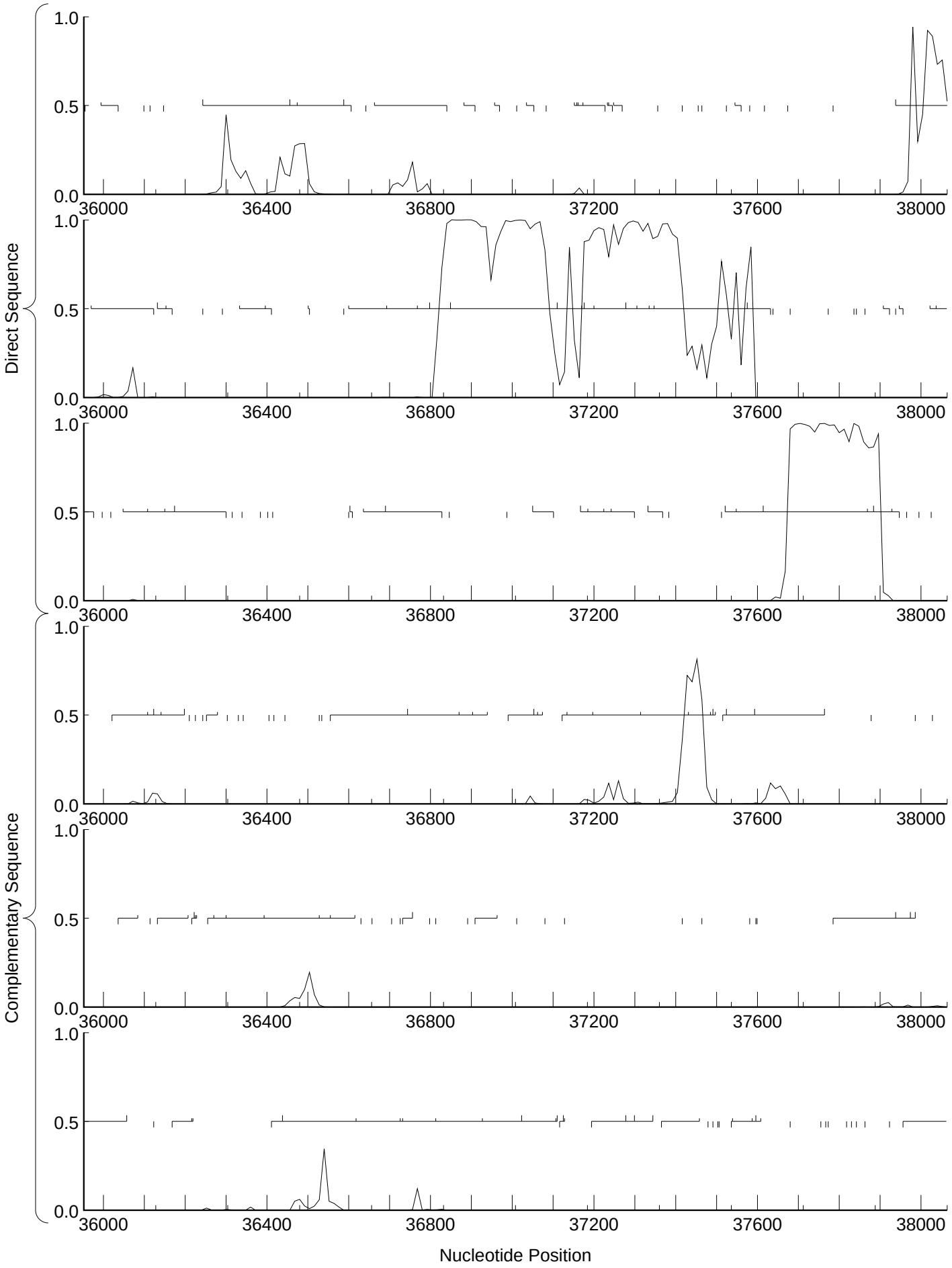
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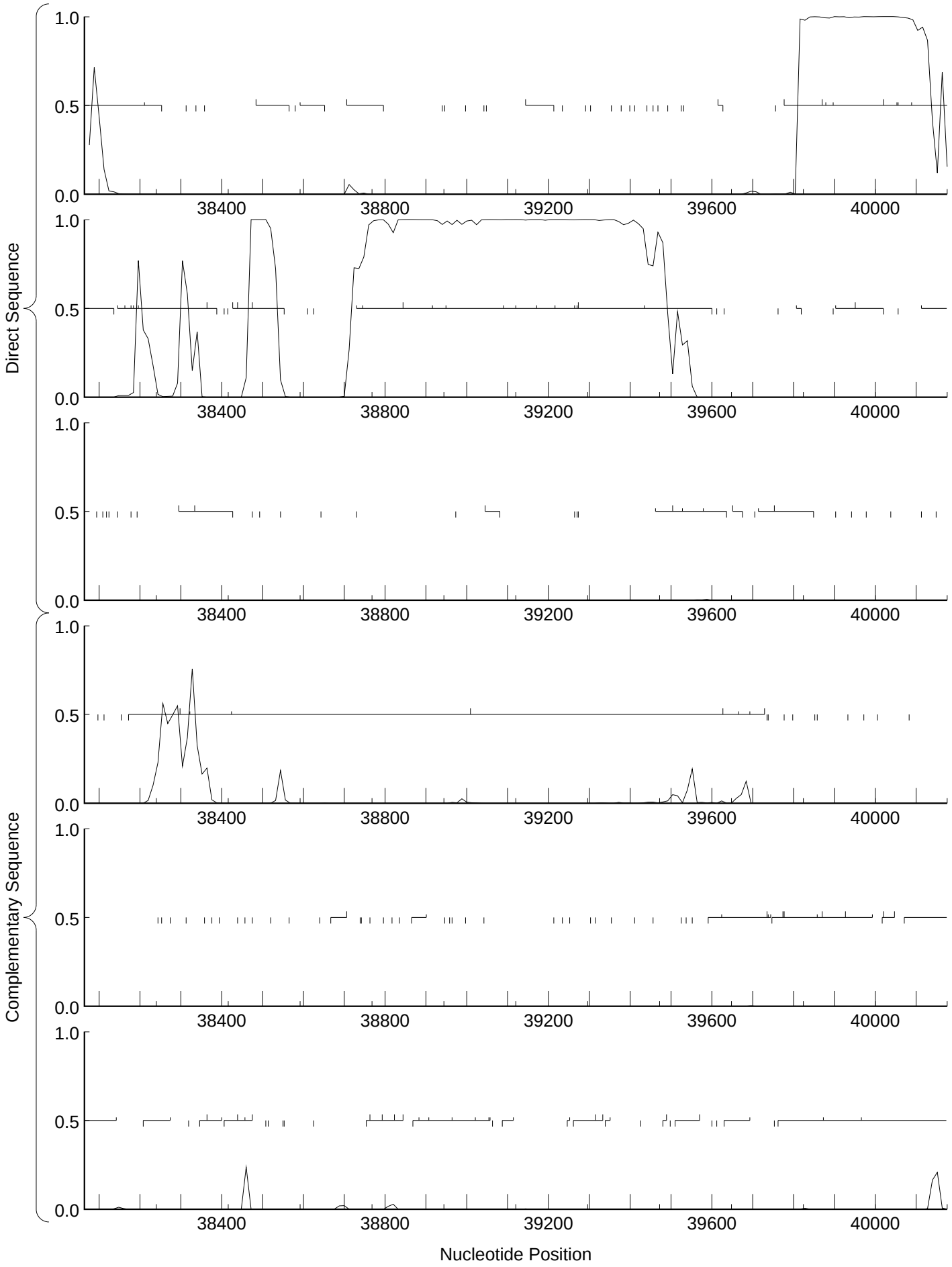
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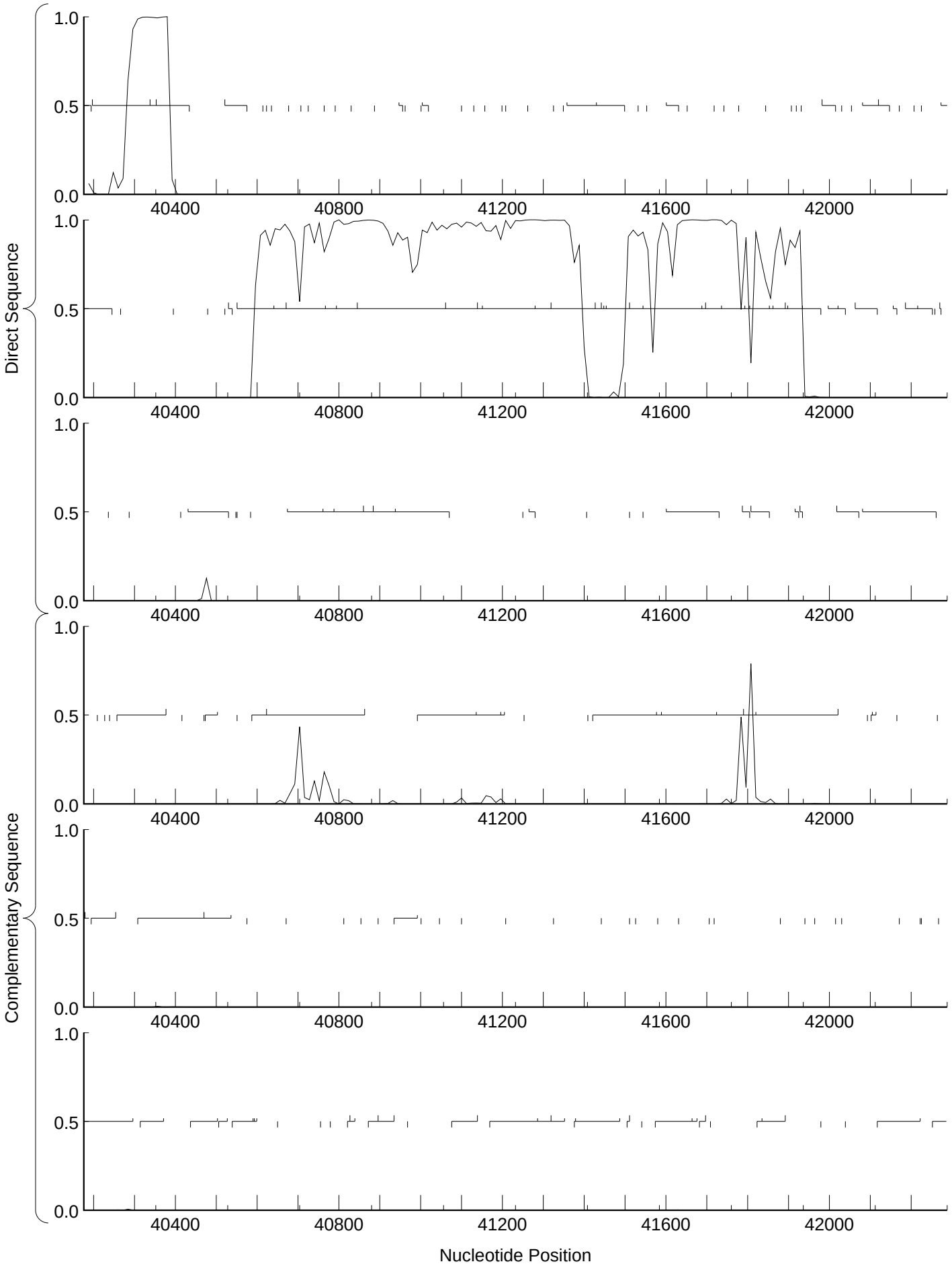
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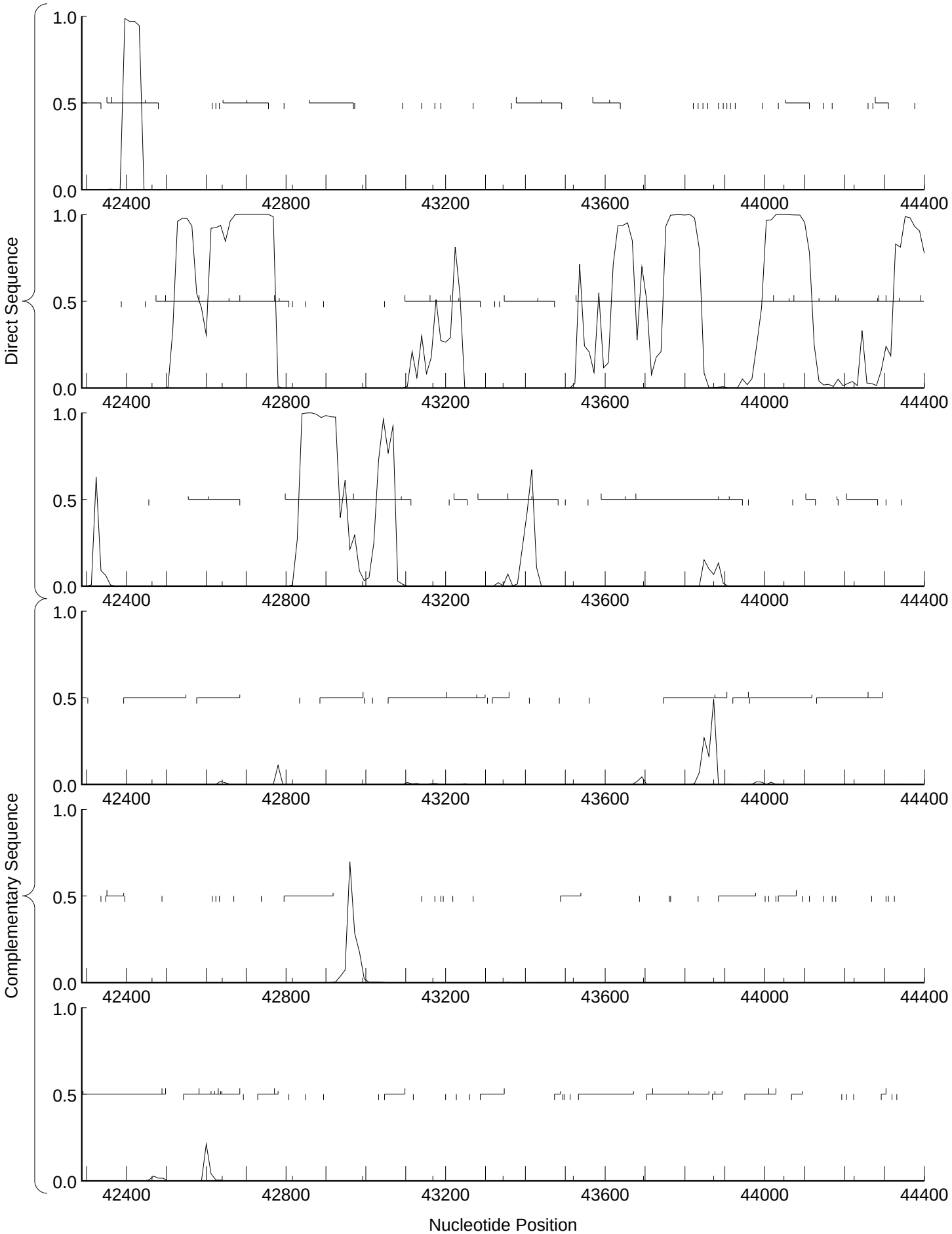
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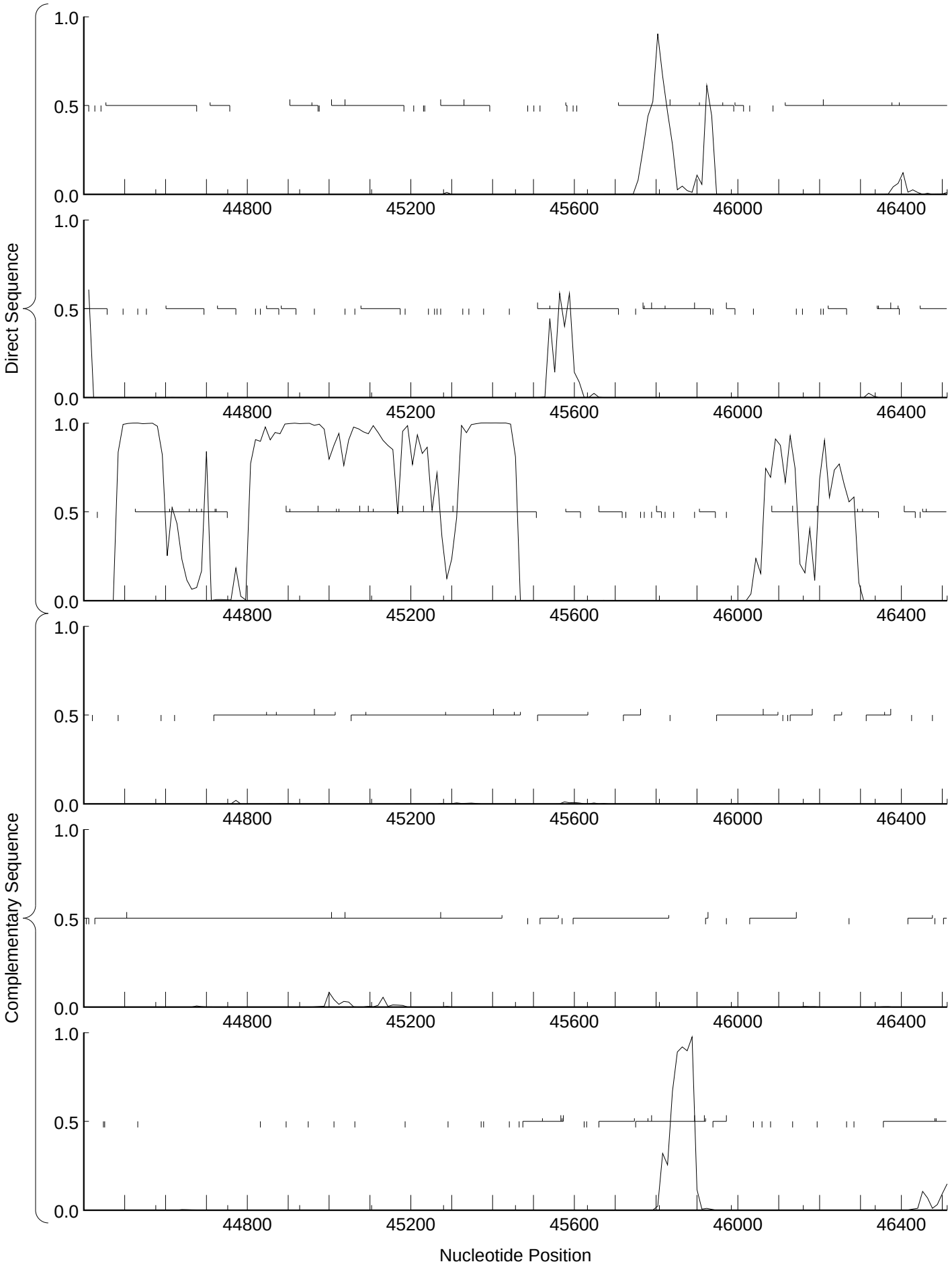
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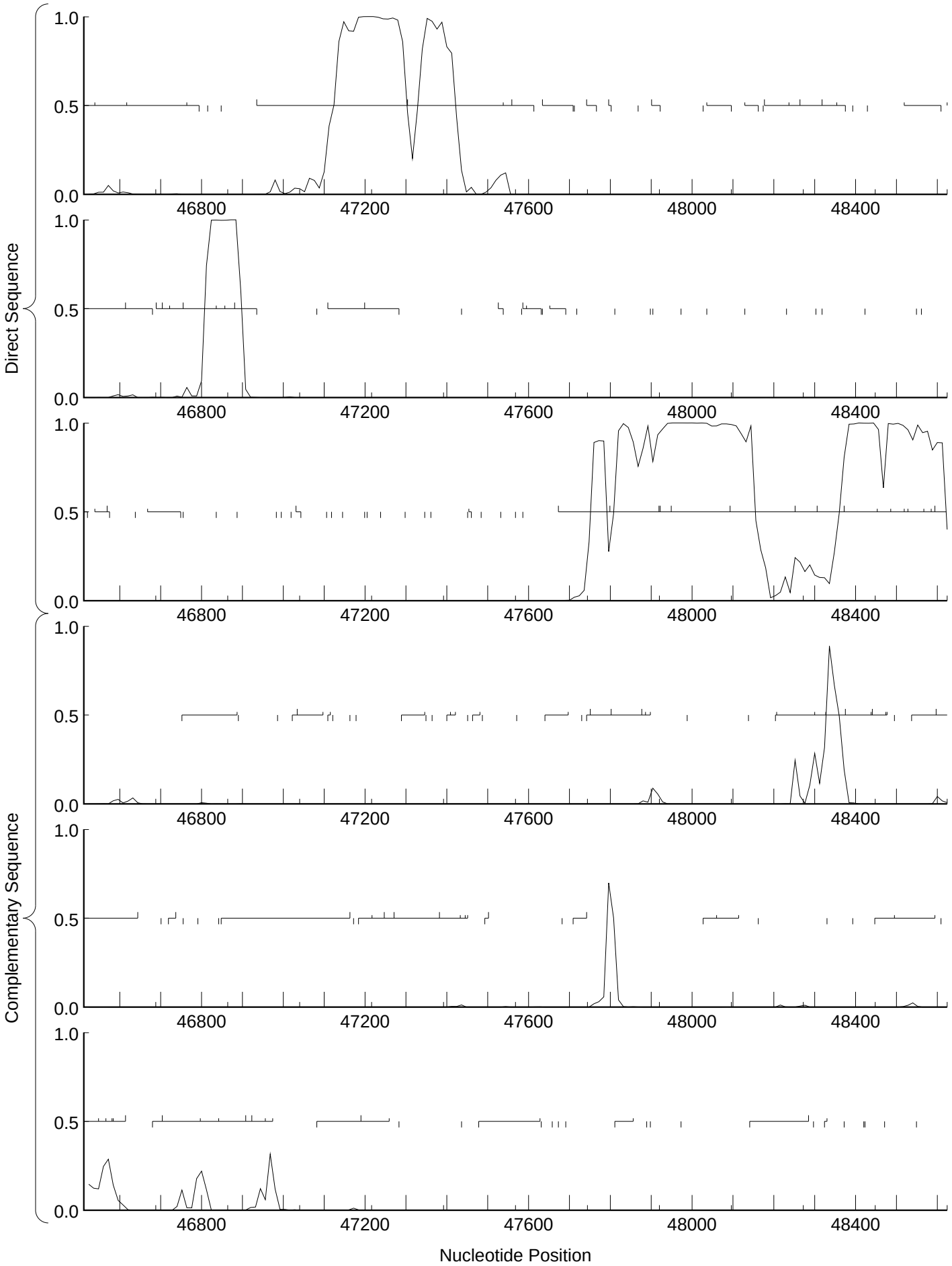
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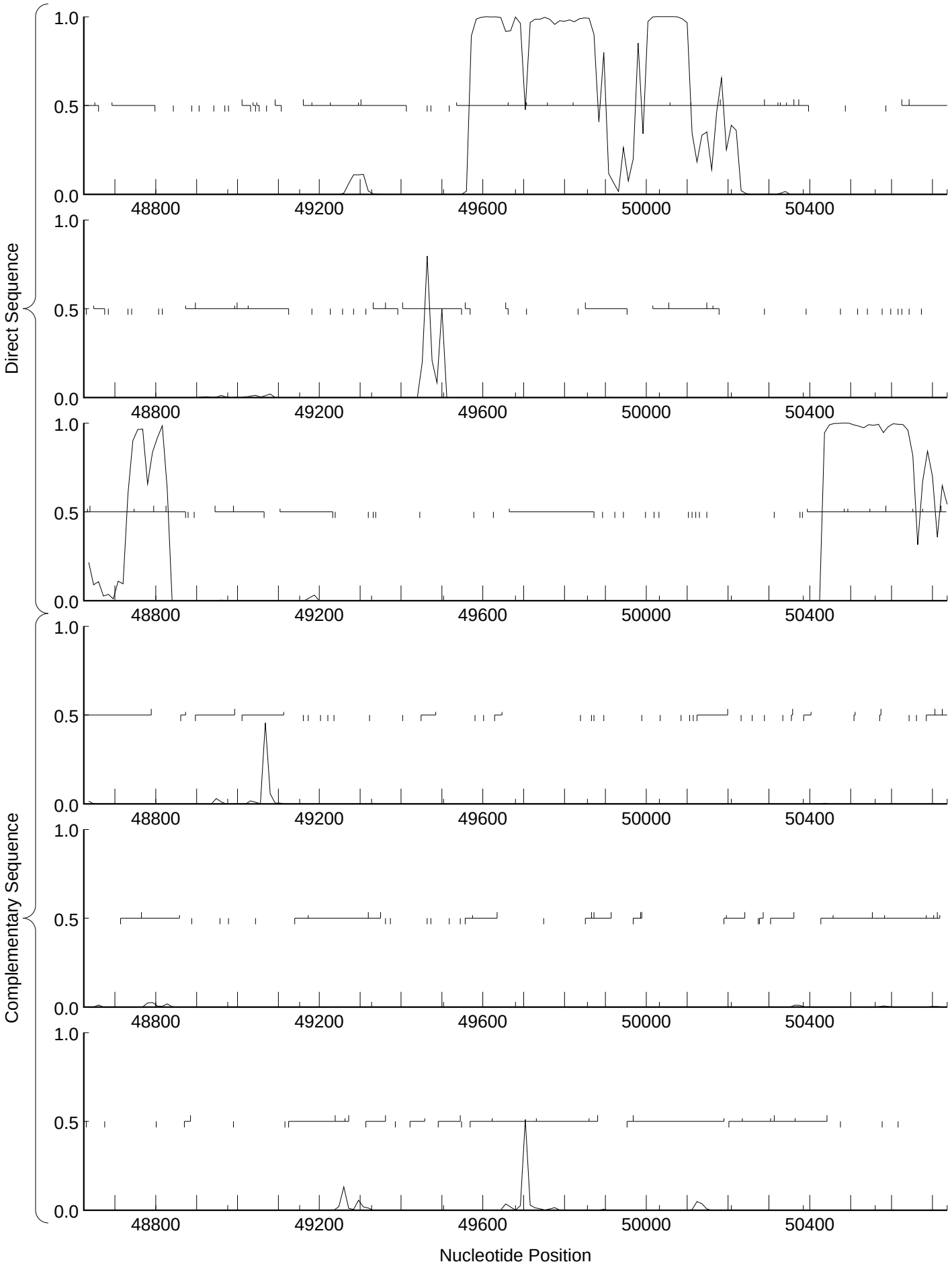
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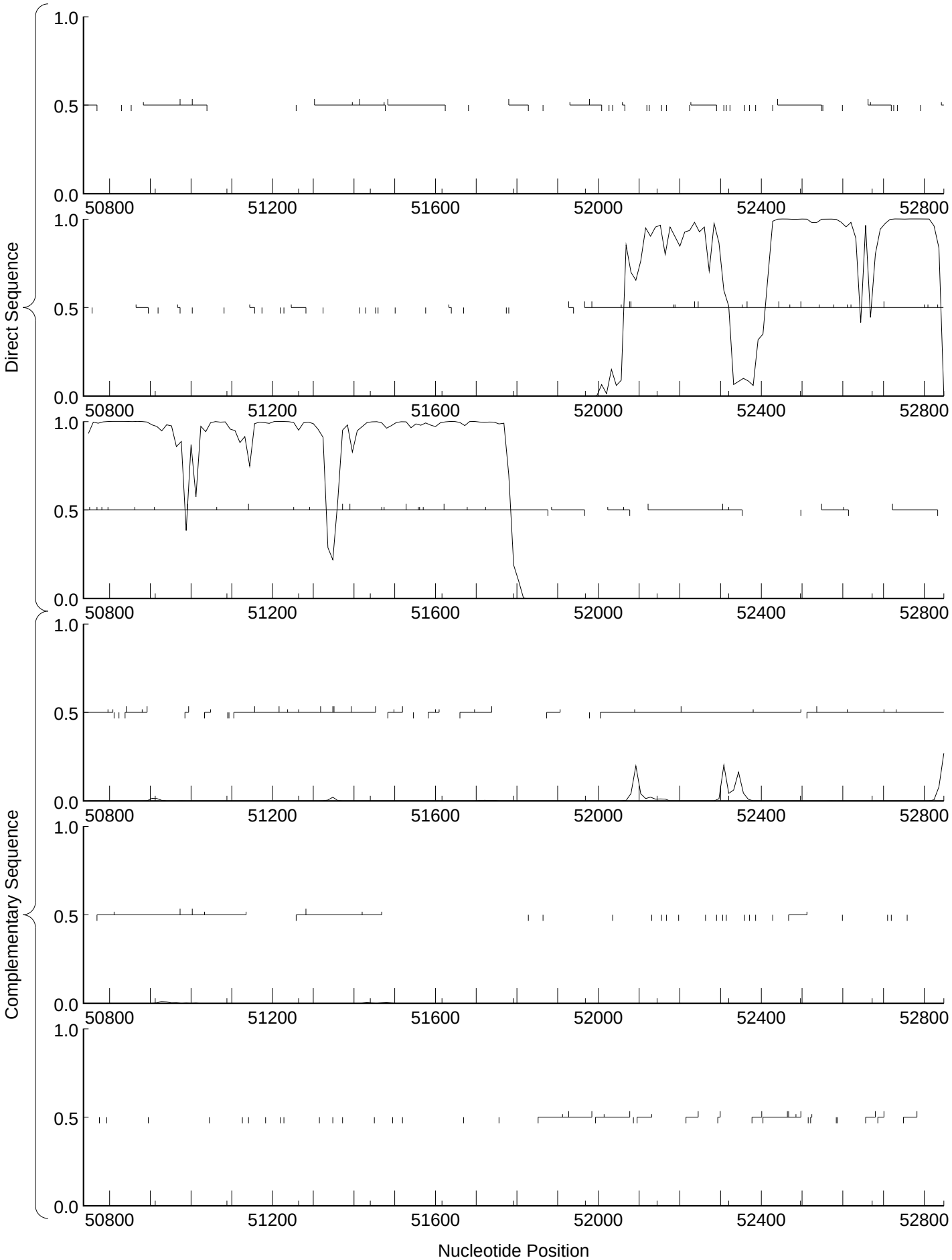
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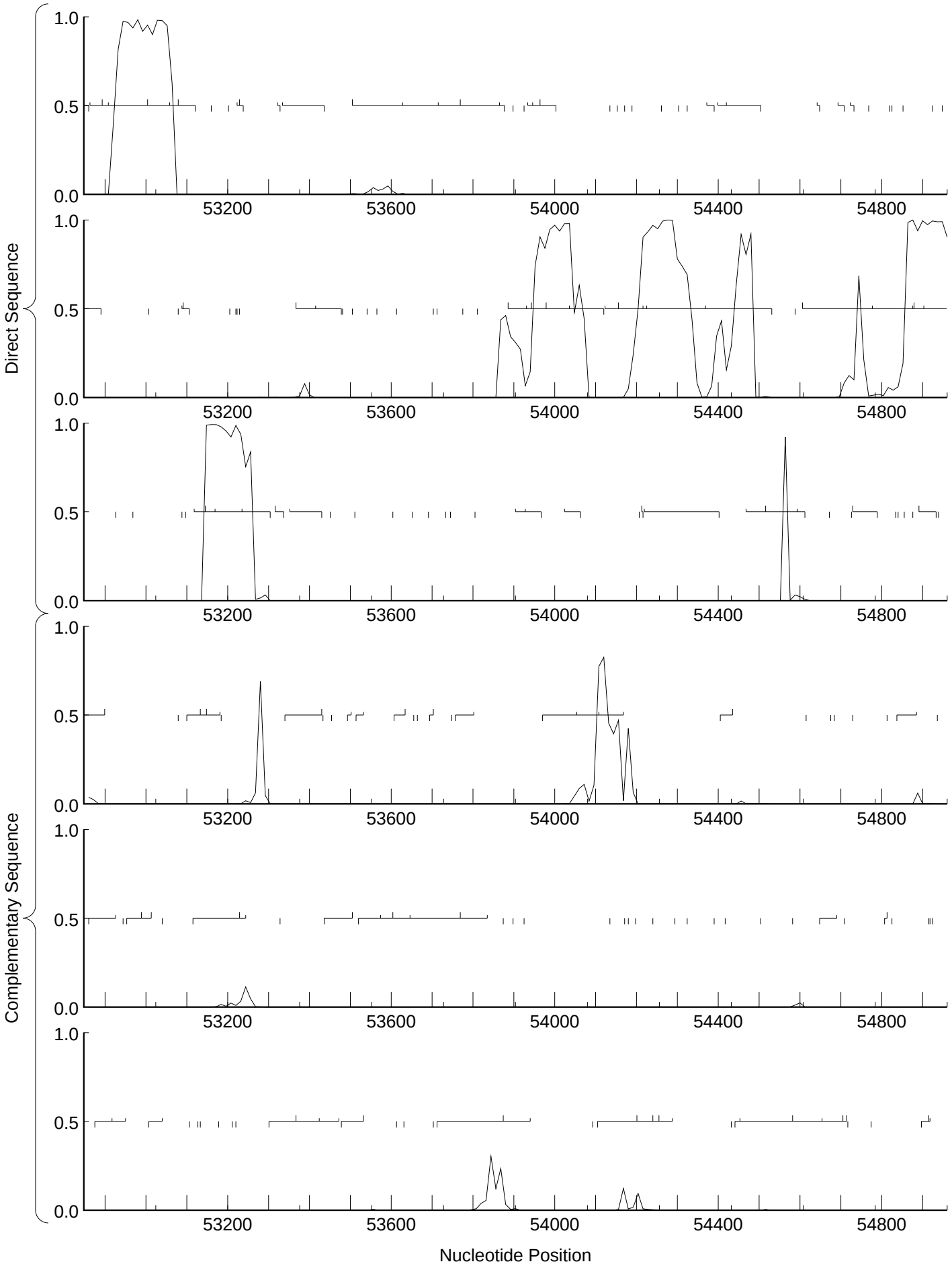
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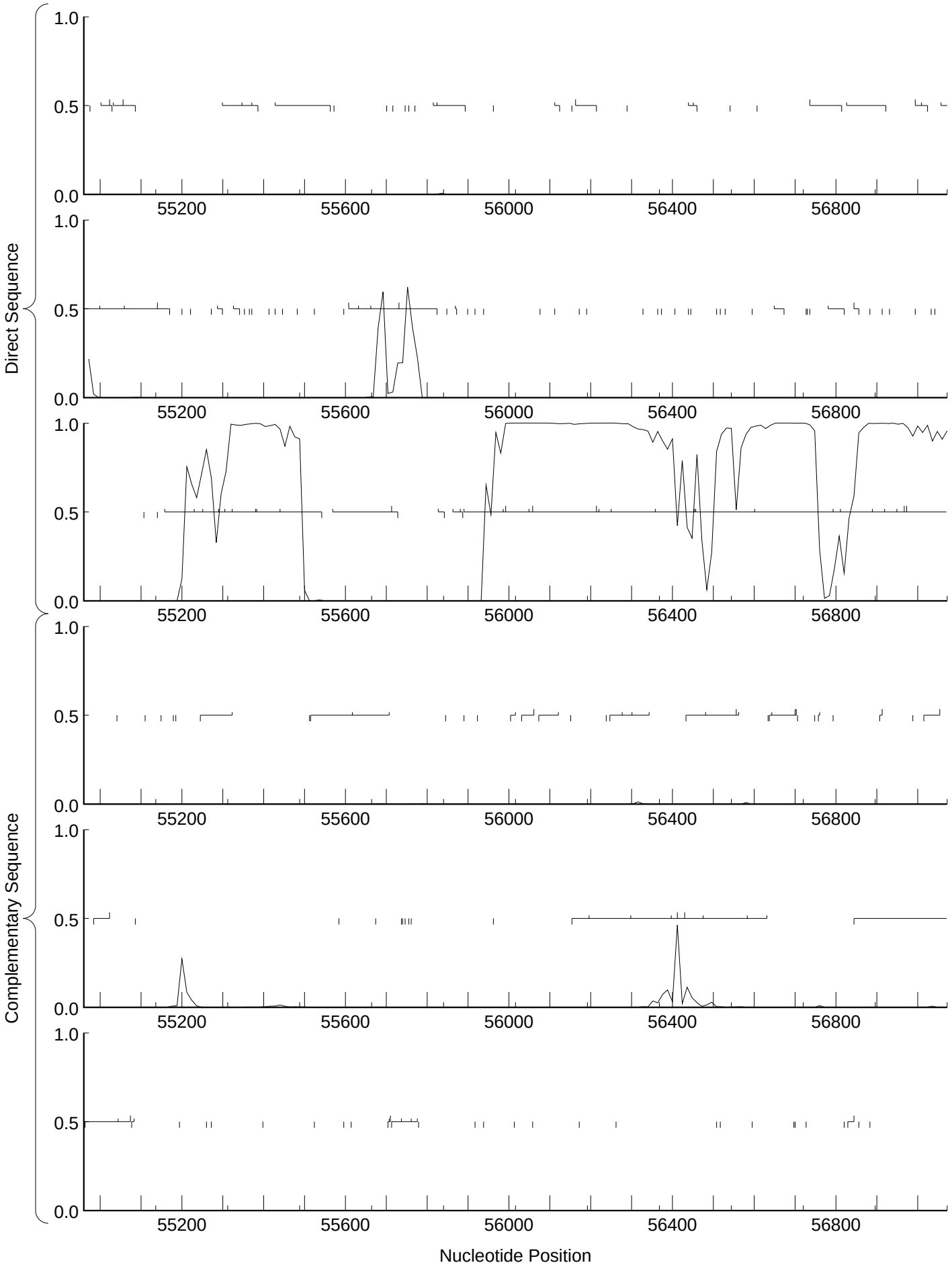


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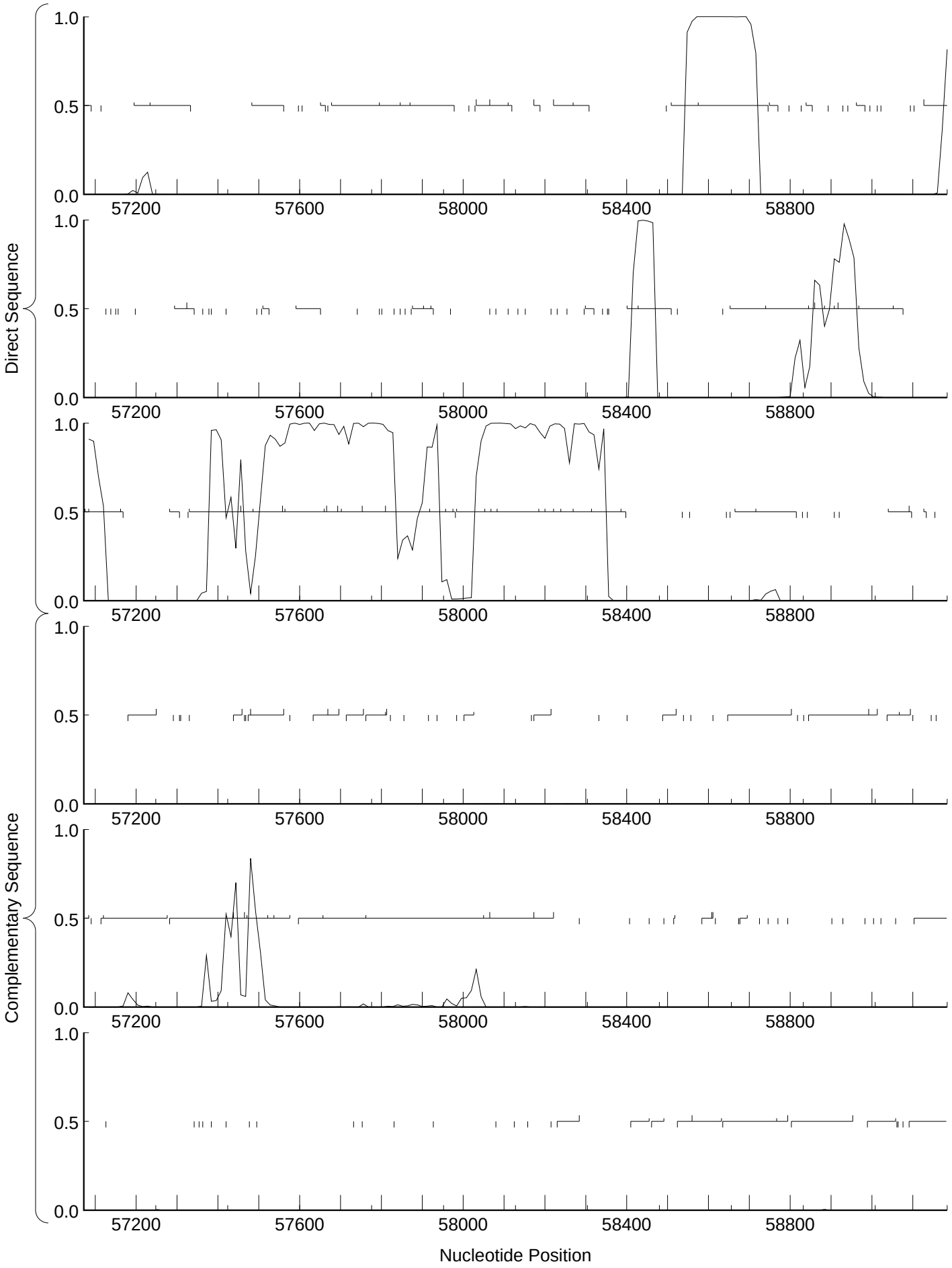


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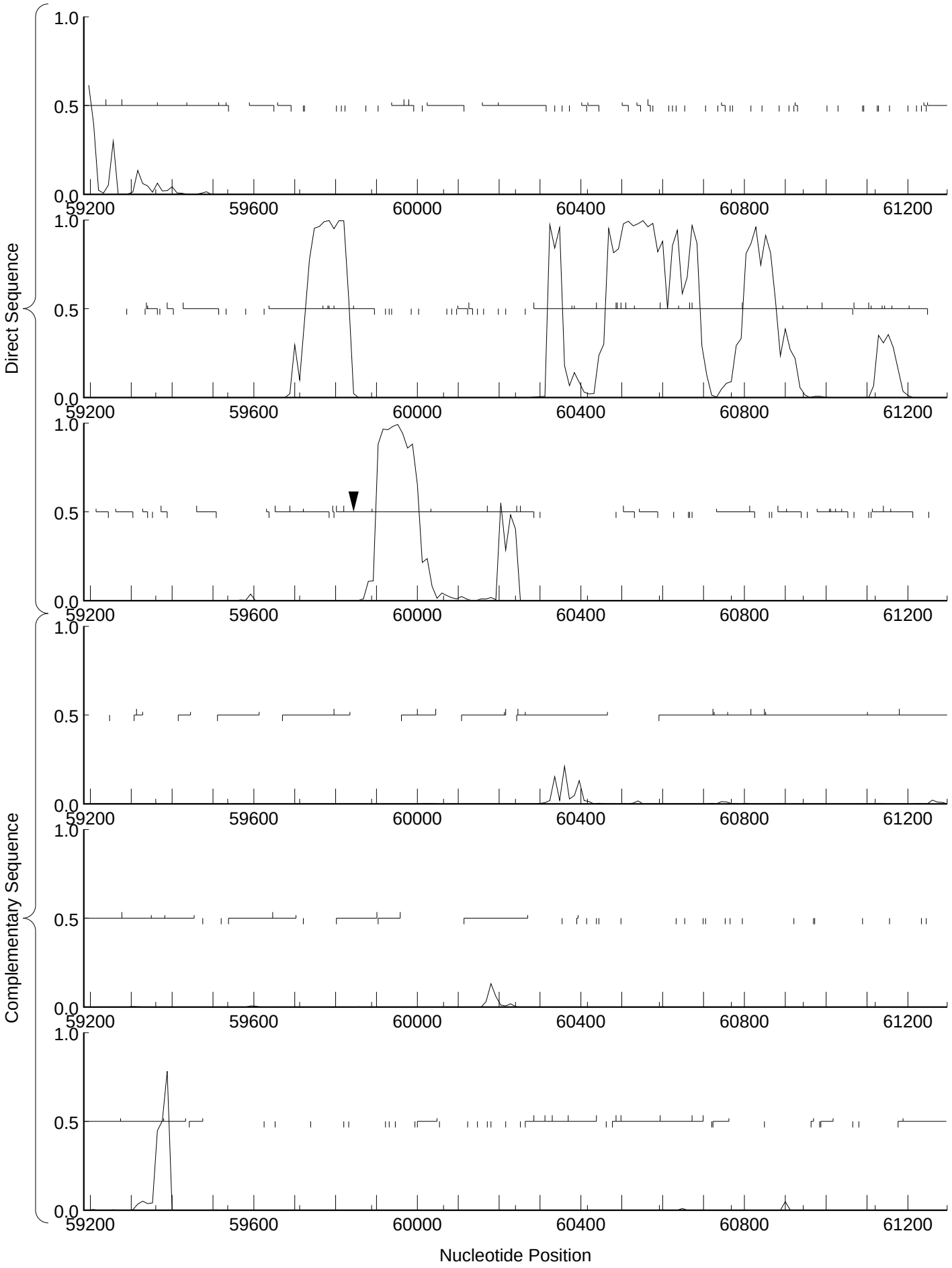



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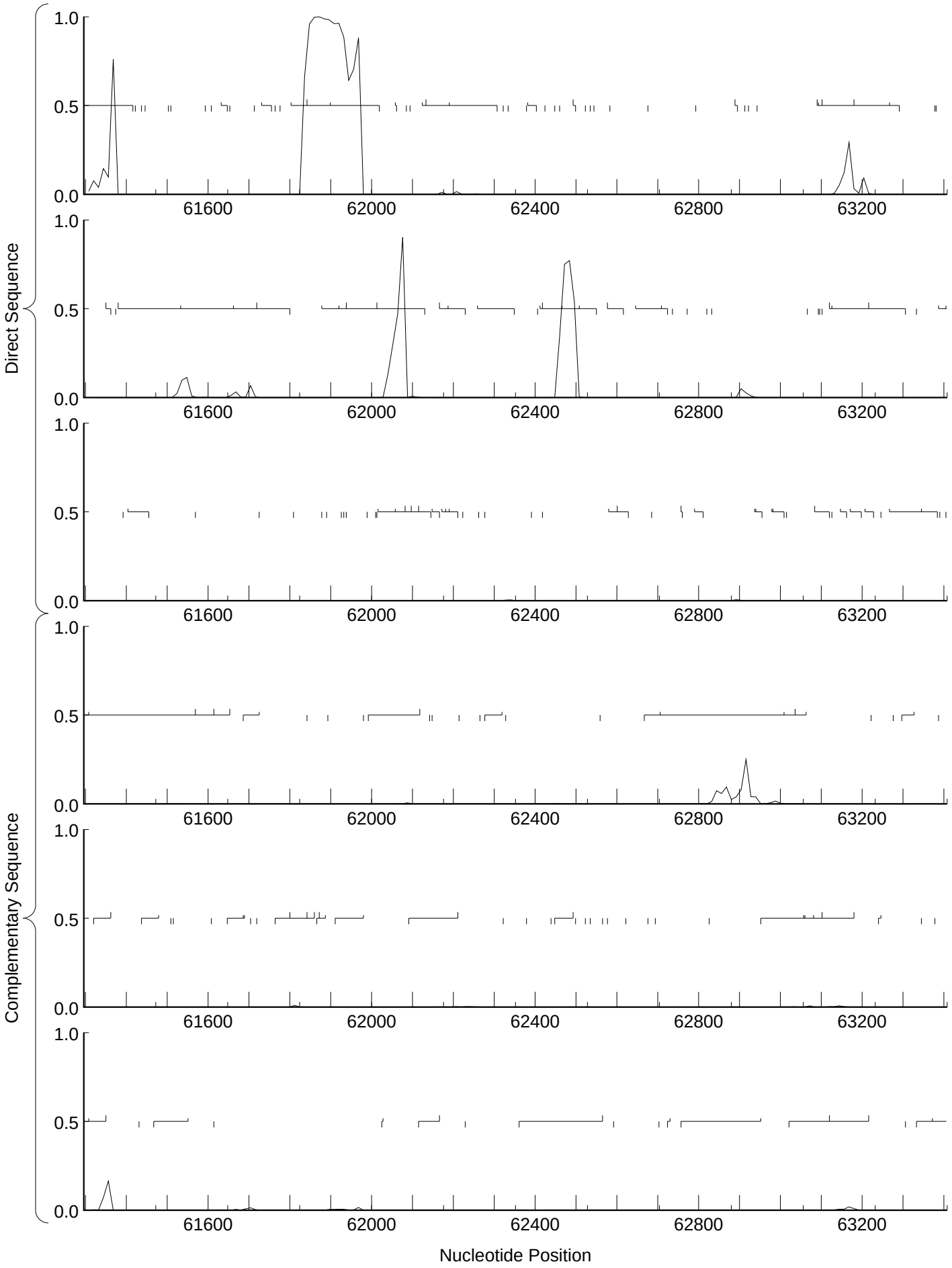
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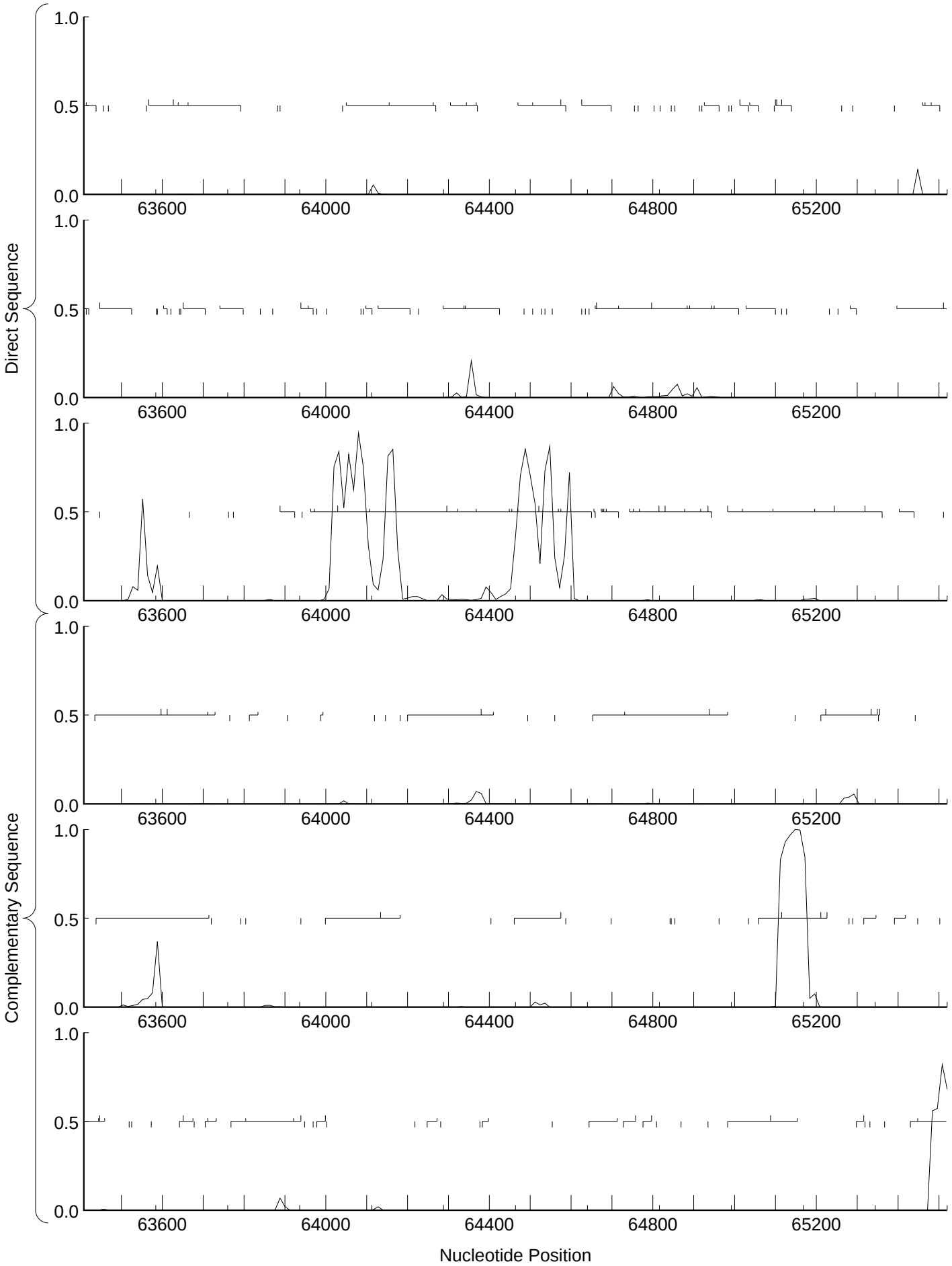
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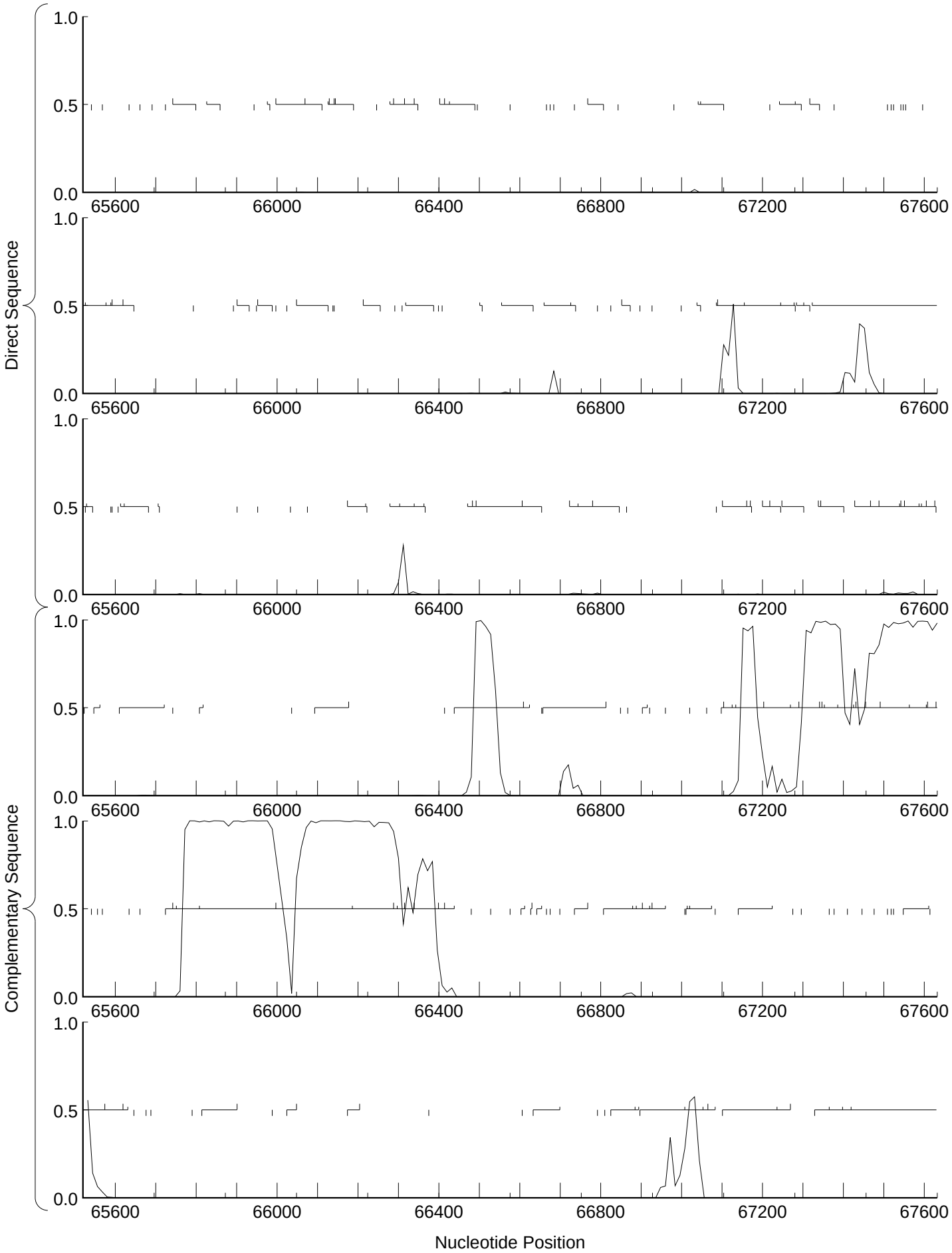
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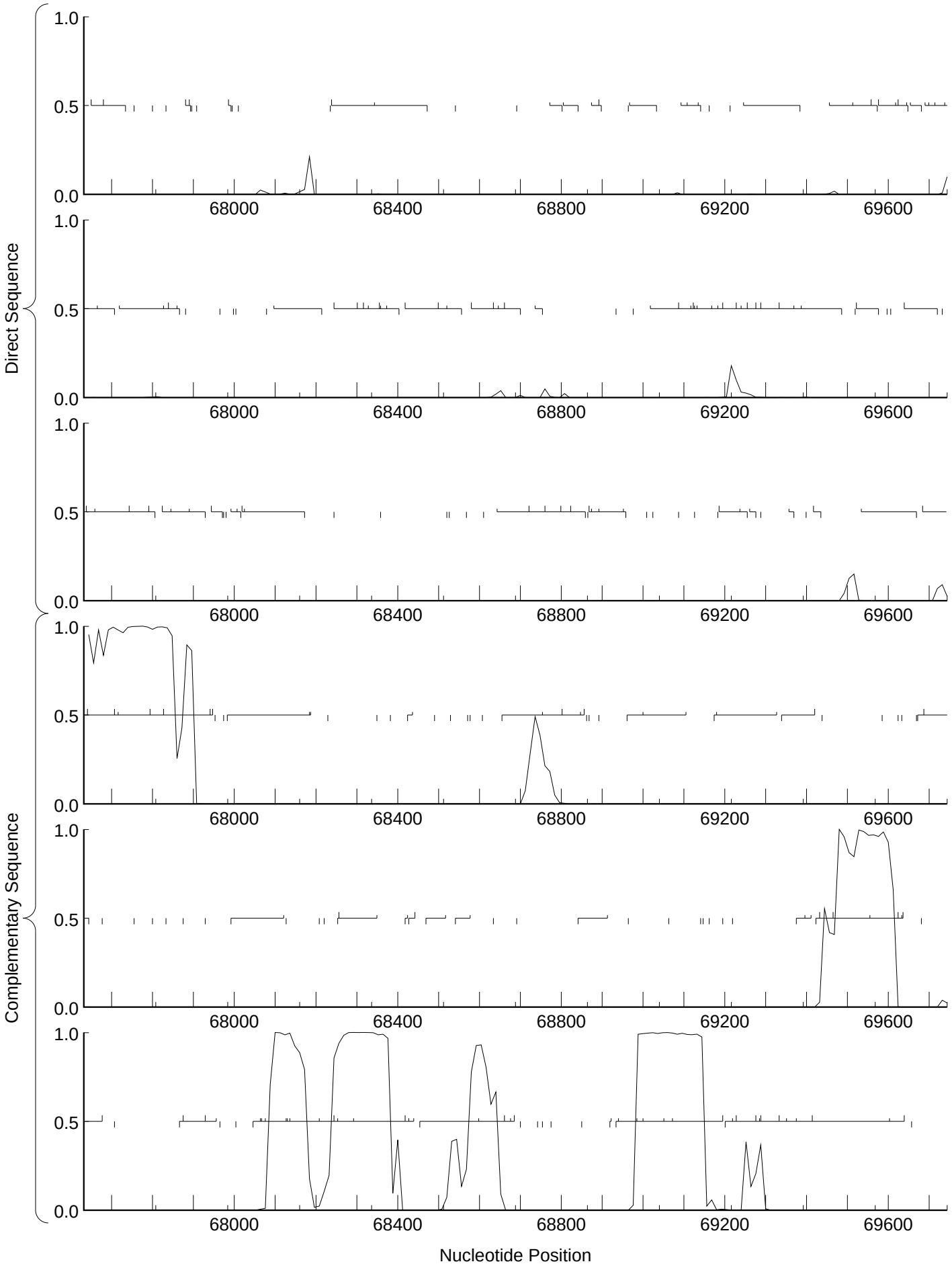
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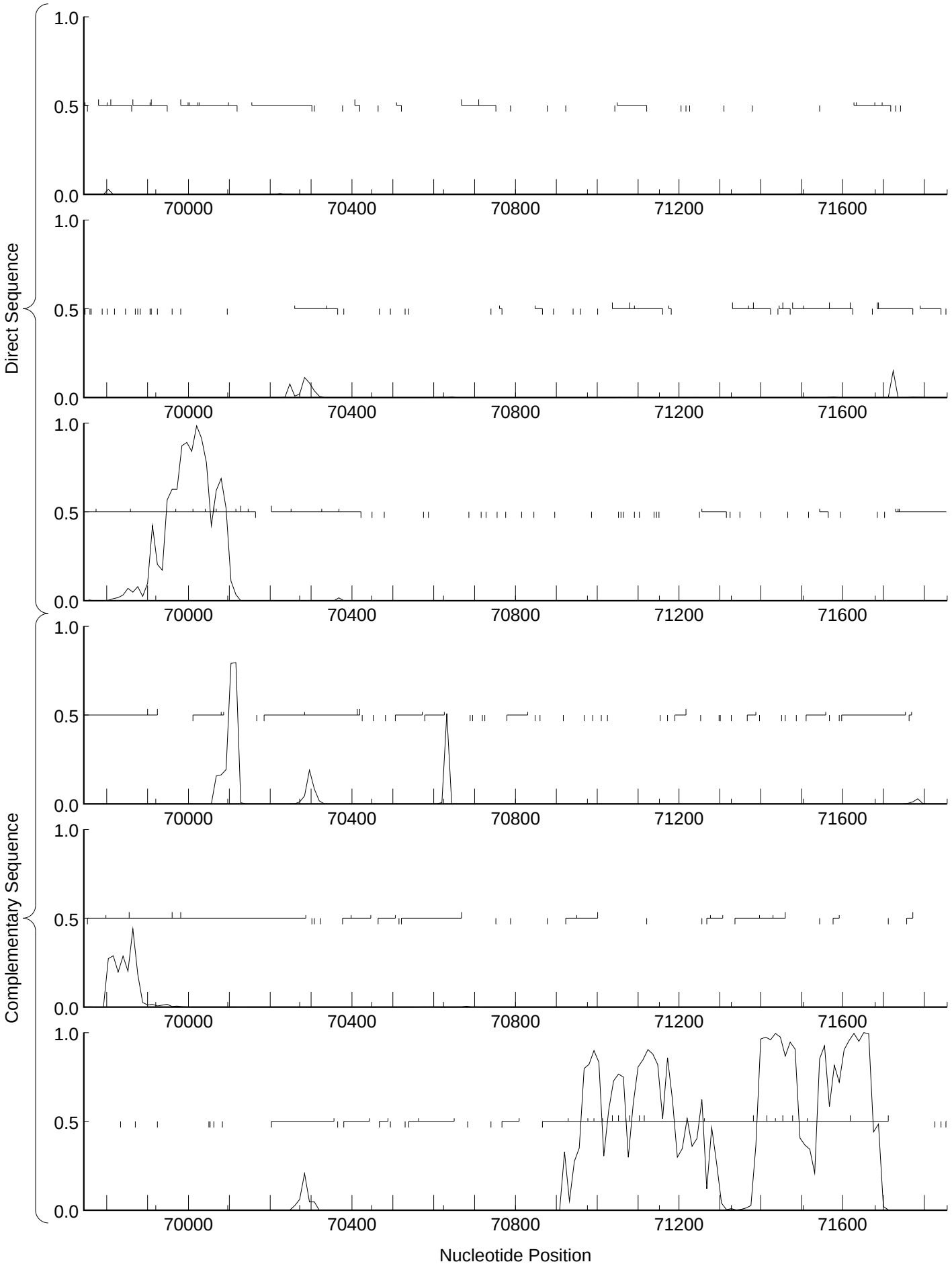
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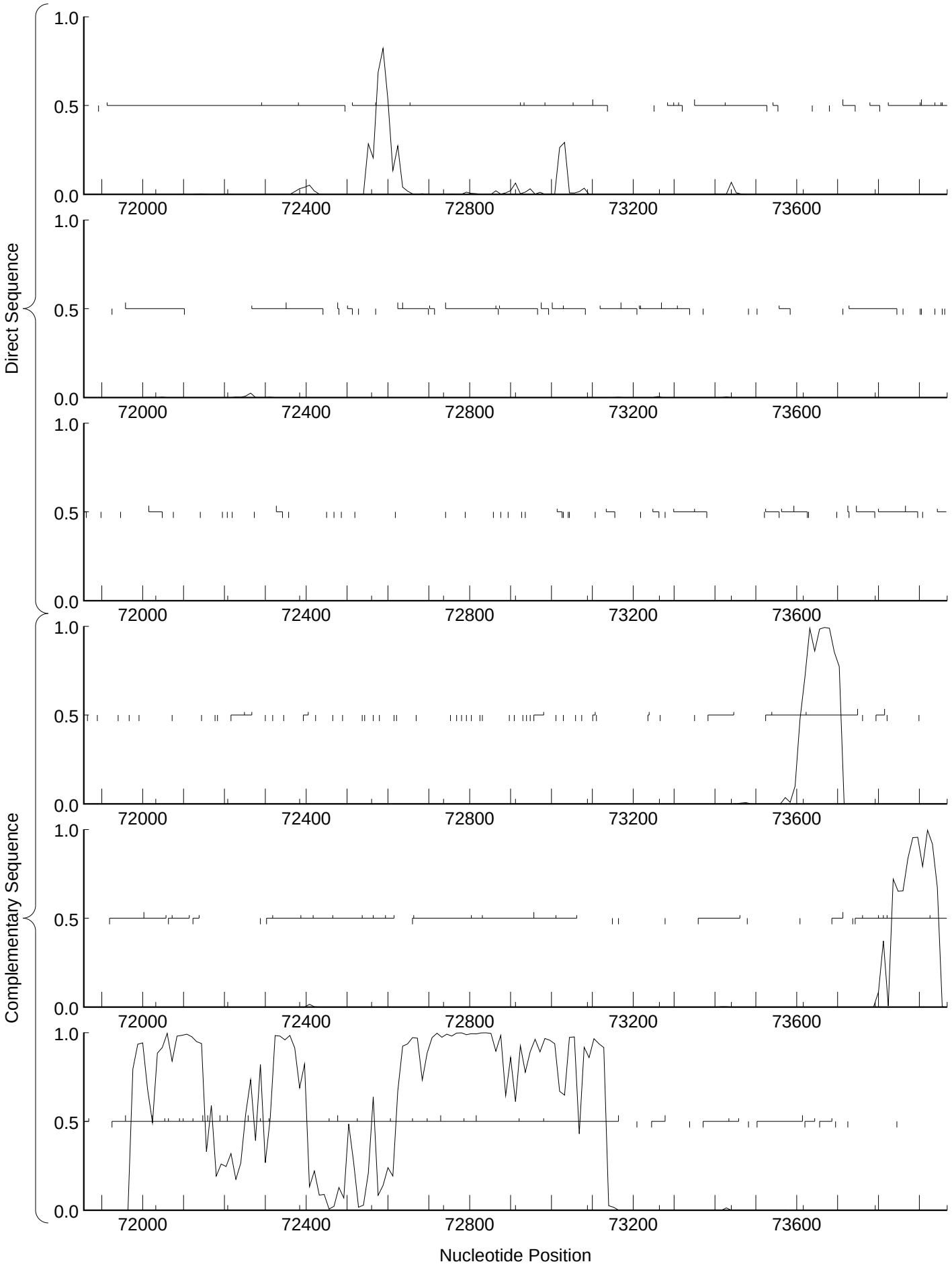
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e complete sequence, 75440 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 35/37



e complete sequence, 75440 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 36/37



e complete sequence, 75440 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 37/37

