# GeneMark

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

## PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Myrale complete sequence, 76419 bp including 9-base 3' overhang (CGCTTG
Analysis Date    :  3/29/24 at 3:35:18
Pages            :  38
Sequence Length  :  76419 bp
GC Content       :   62.83%

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

PS-Version       :  1.2

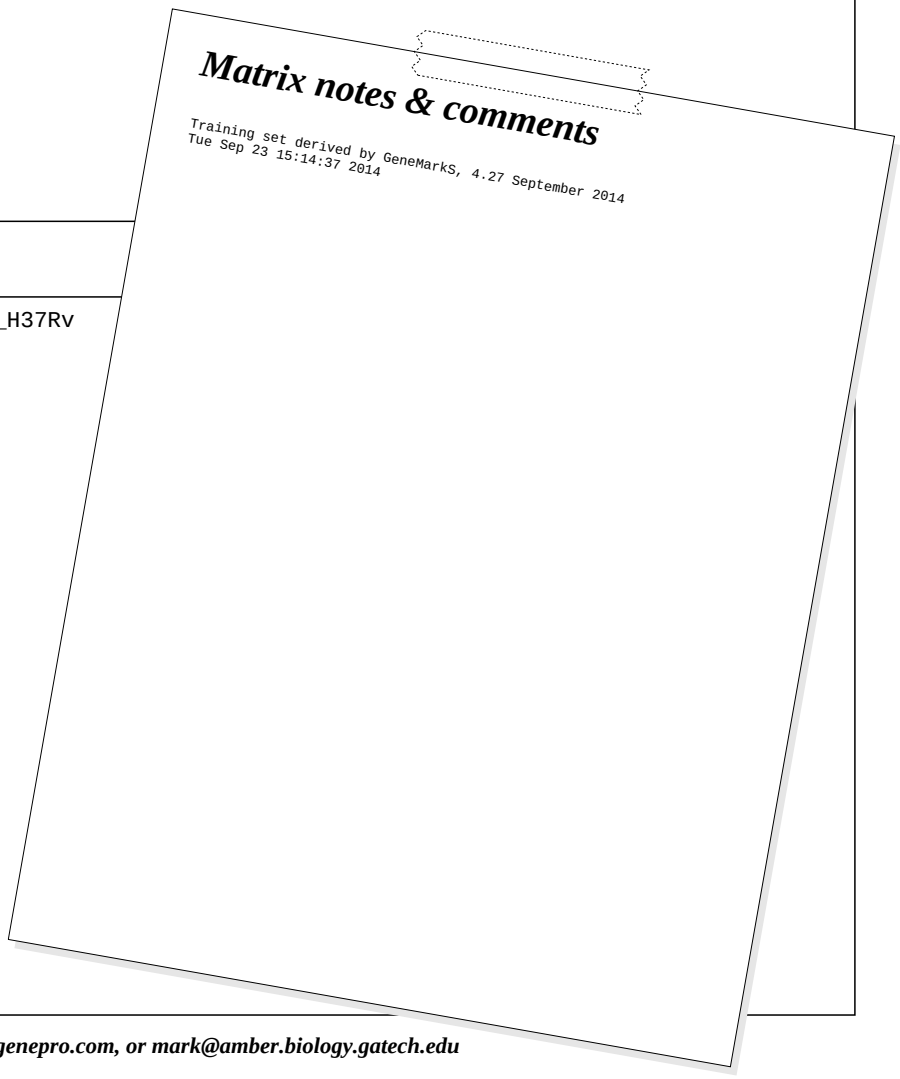
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                    Mark ORFs / splice sites,
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## MATRIX INFORMATION

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Author     :  -

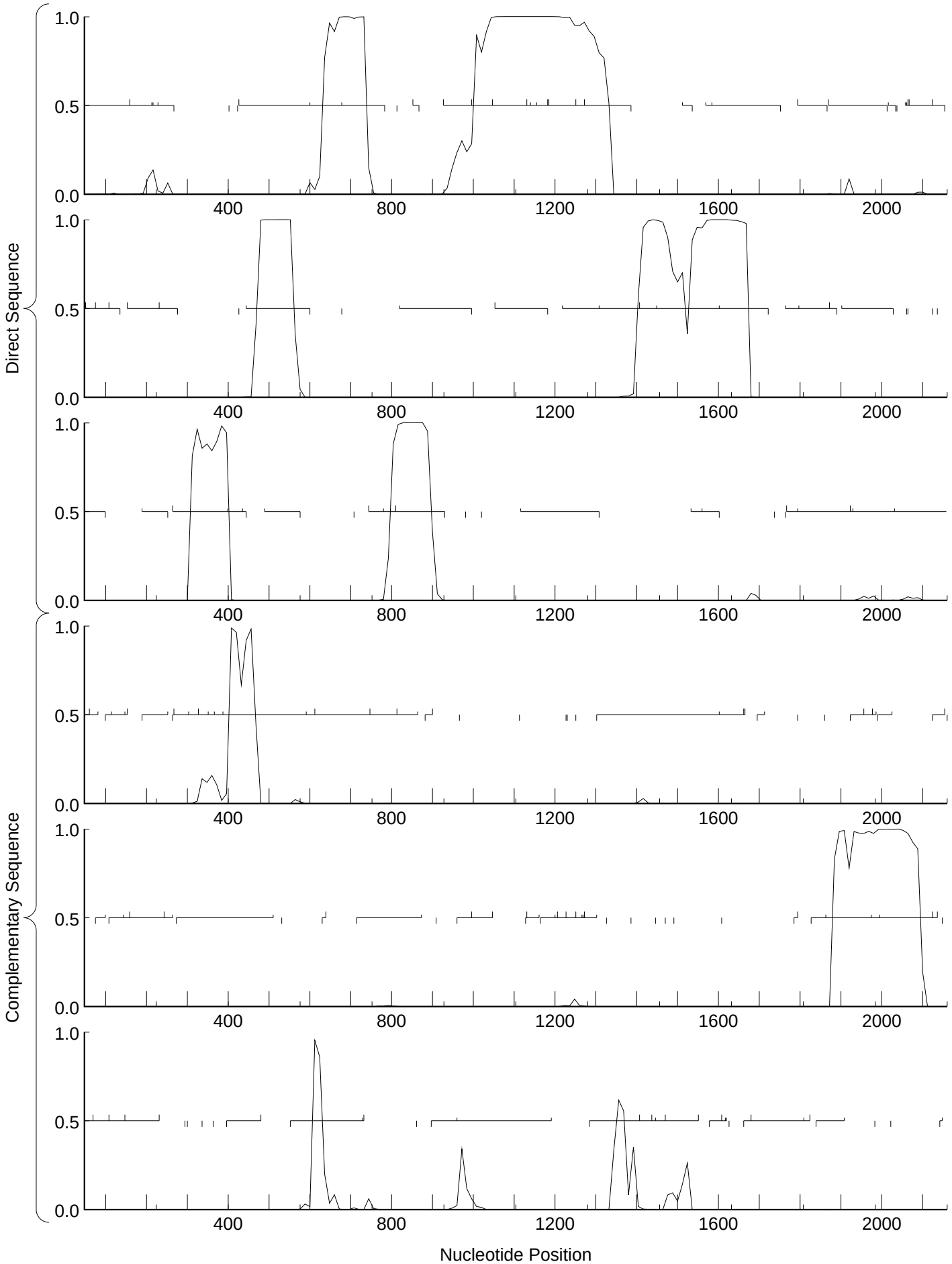
Order      :  4


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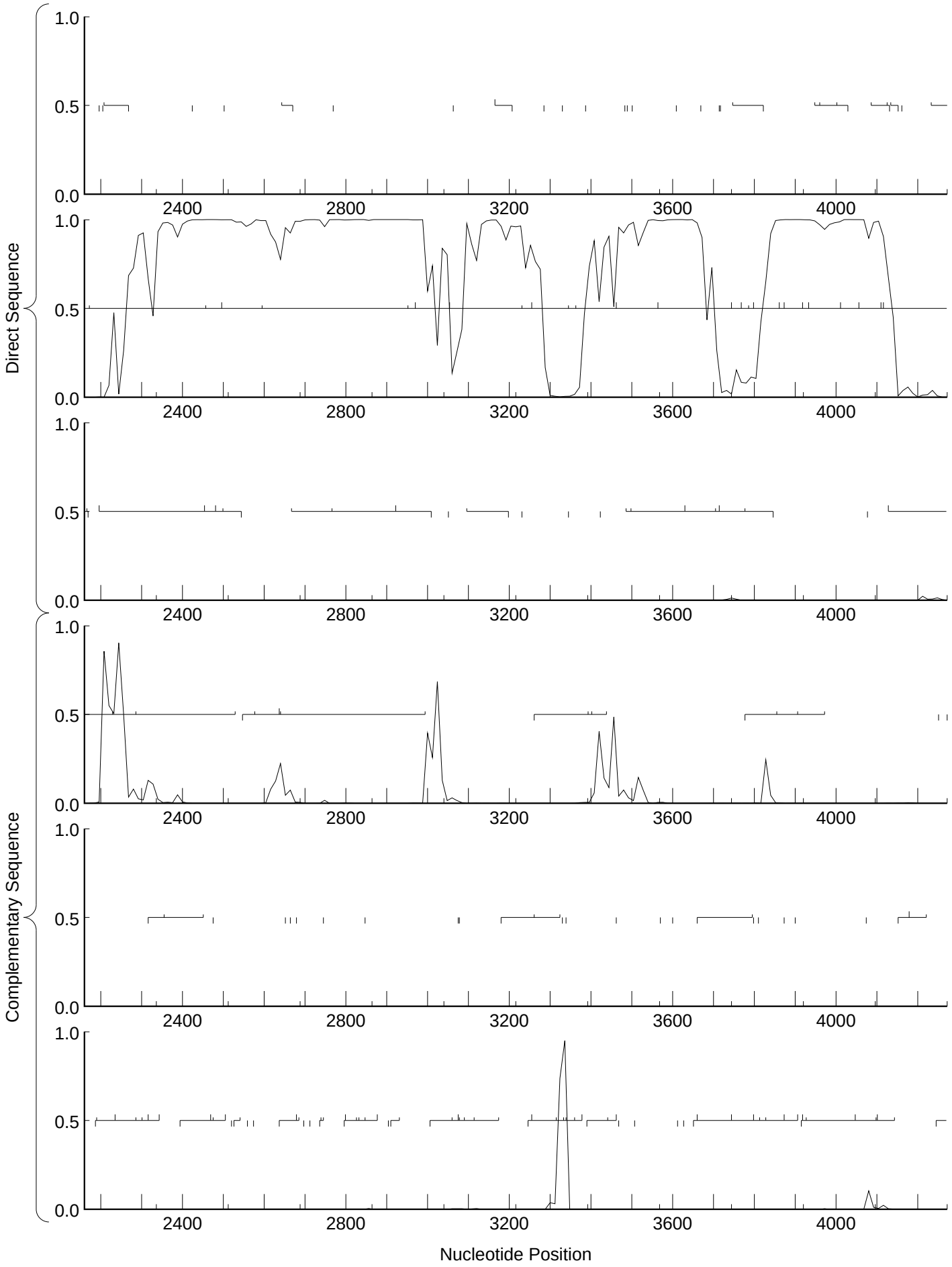



*For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu*

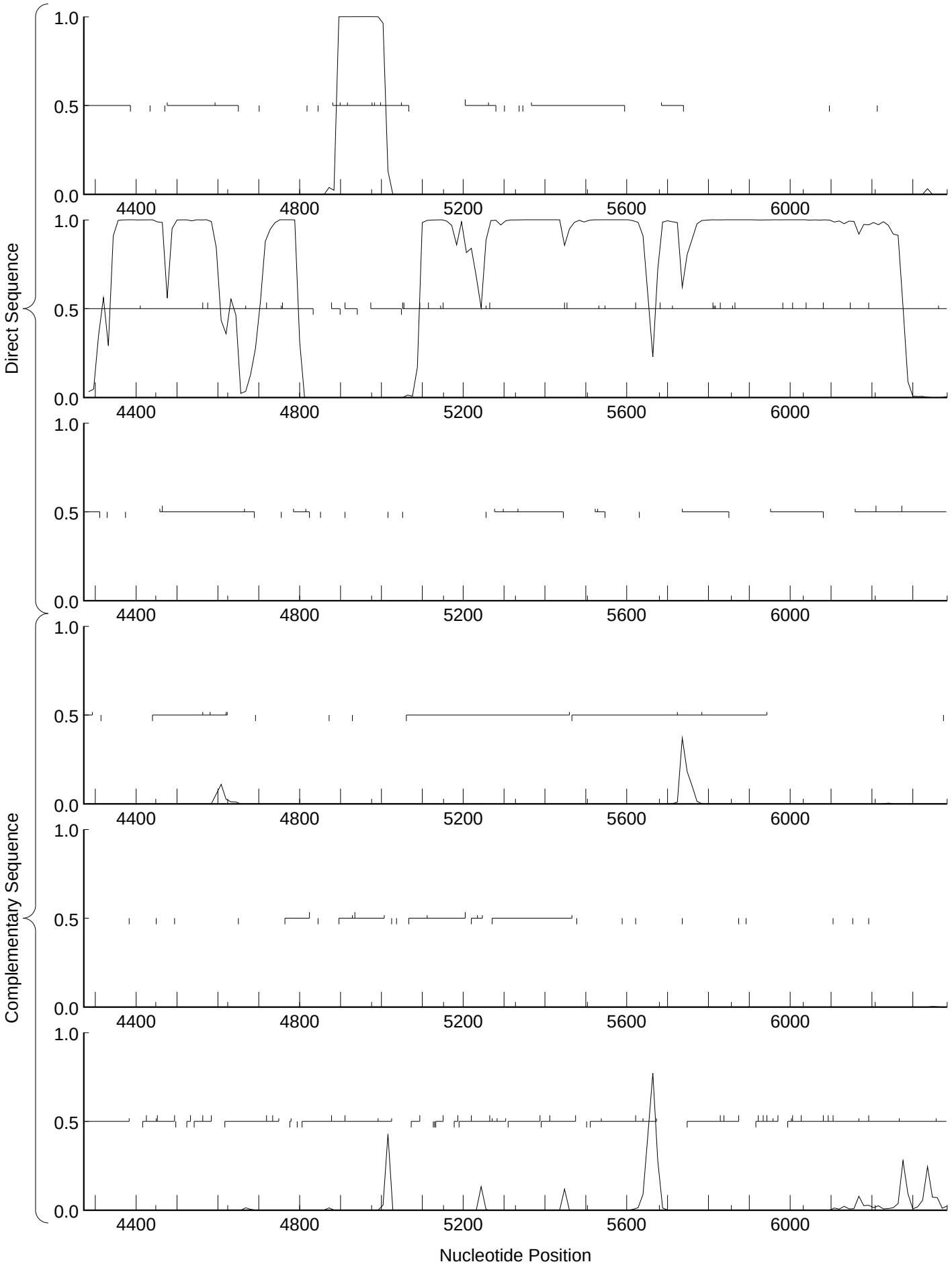
Myrale complete sequence, 76419 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 2/38



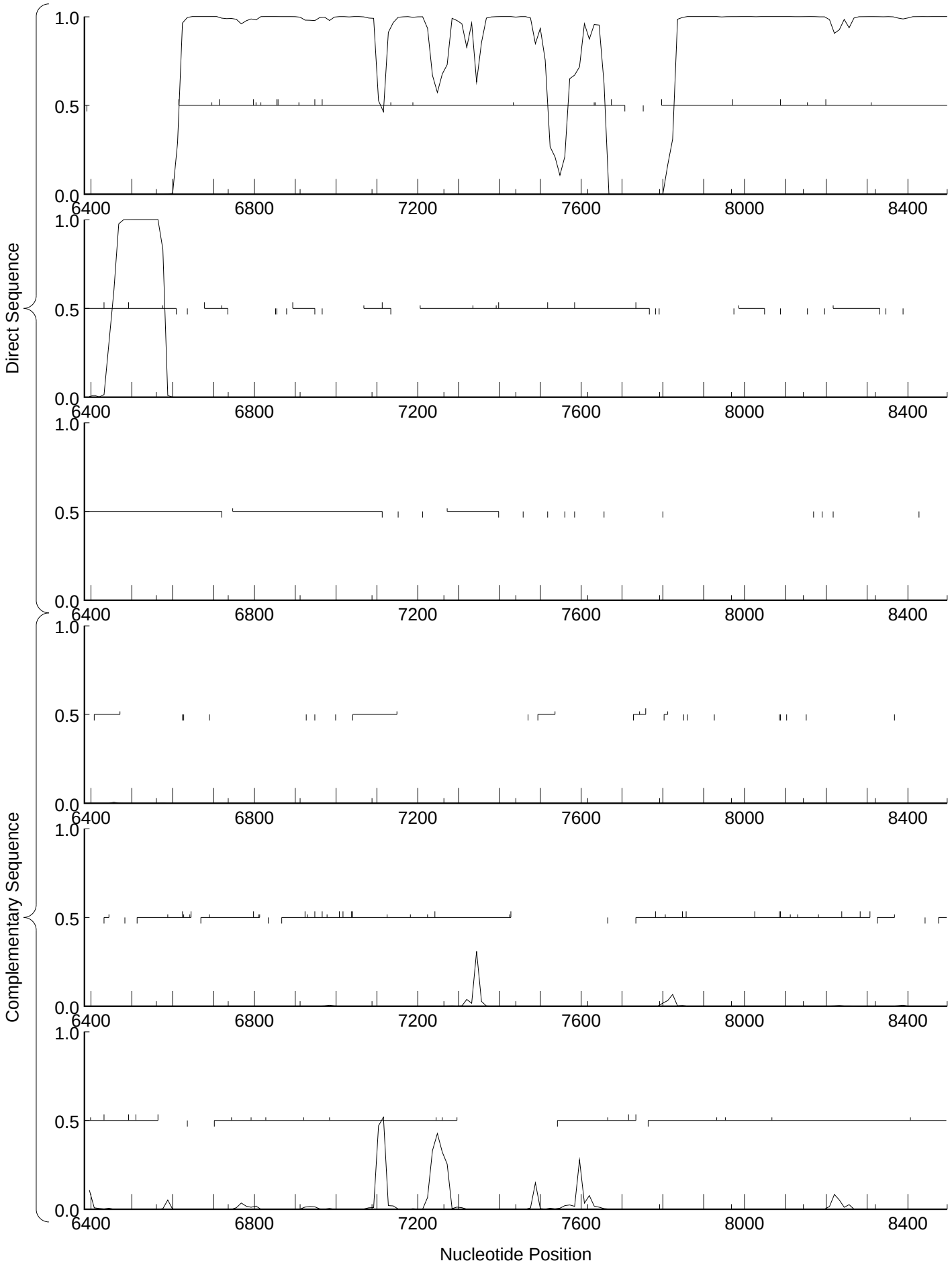
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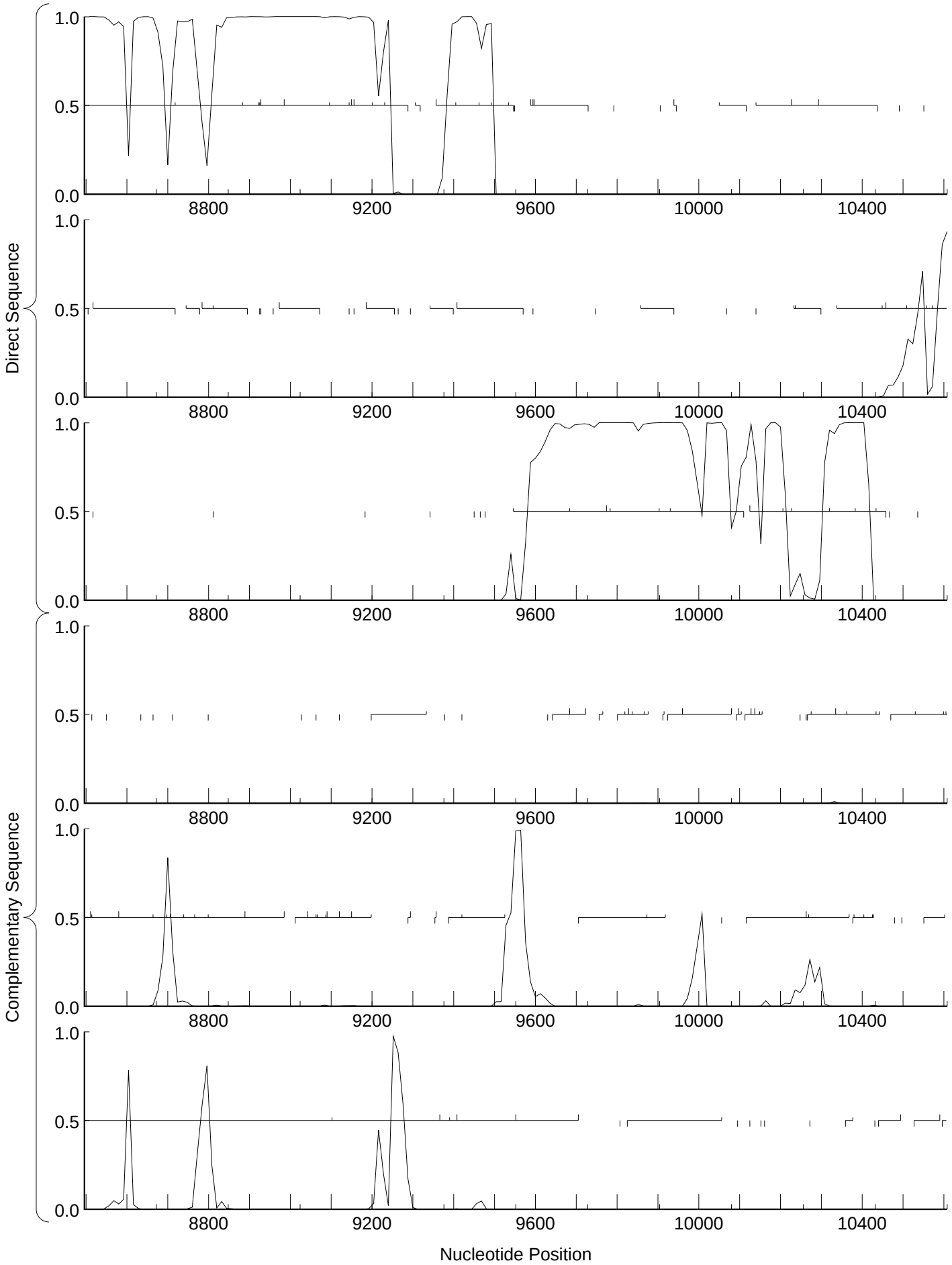
Myrale complete sequence, 76419 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 4/38



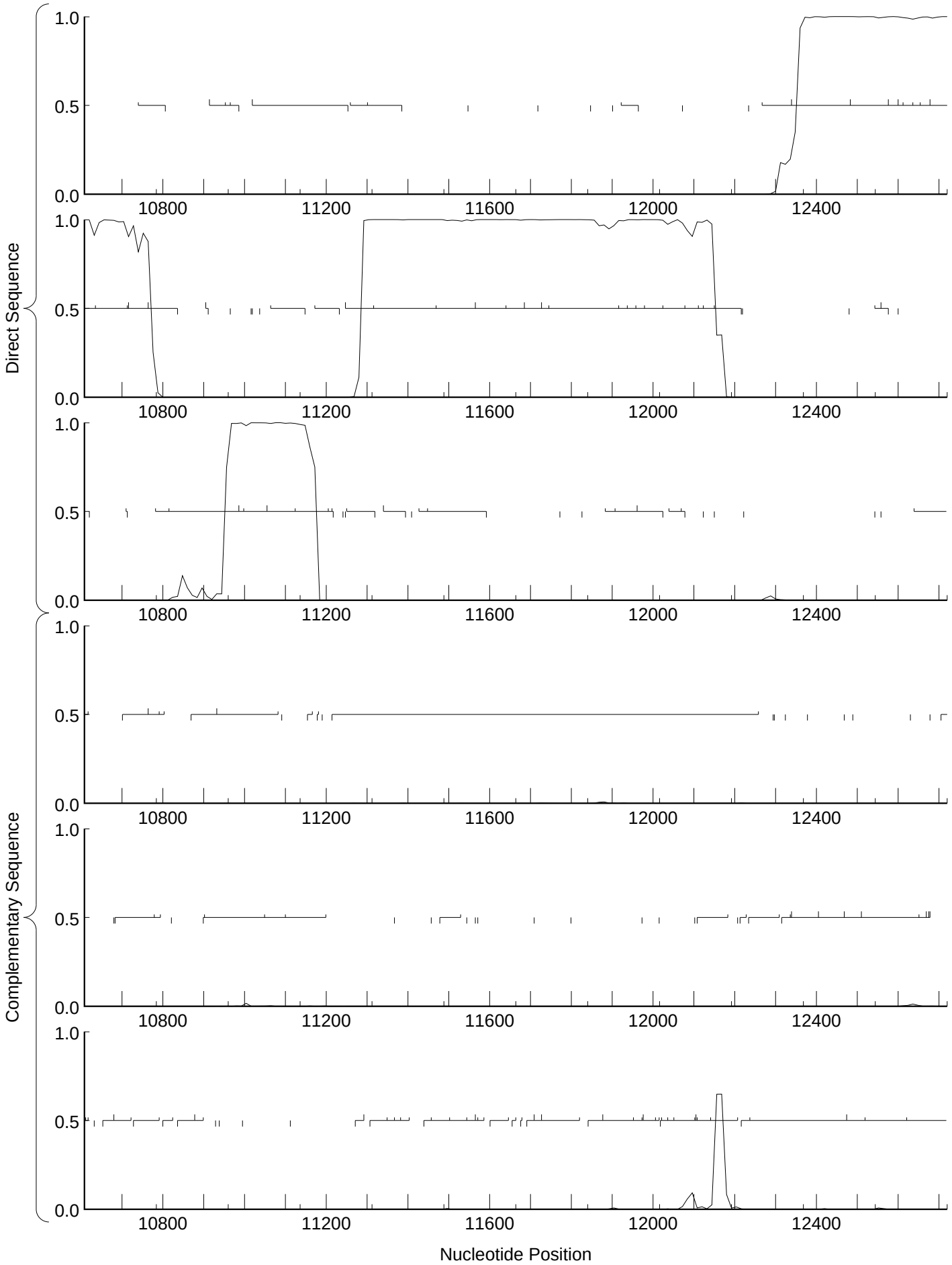
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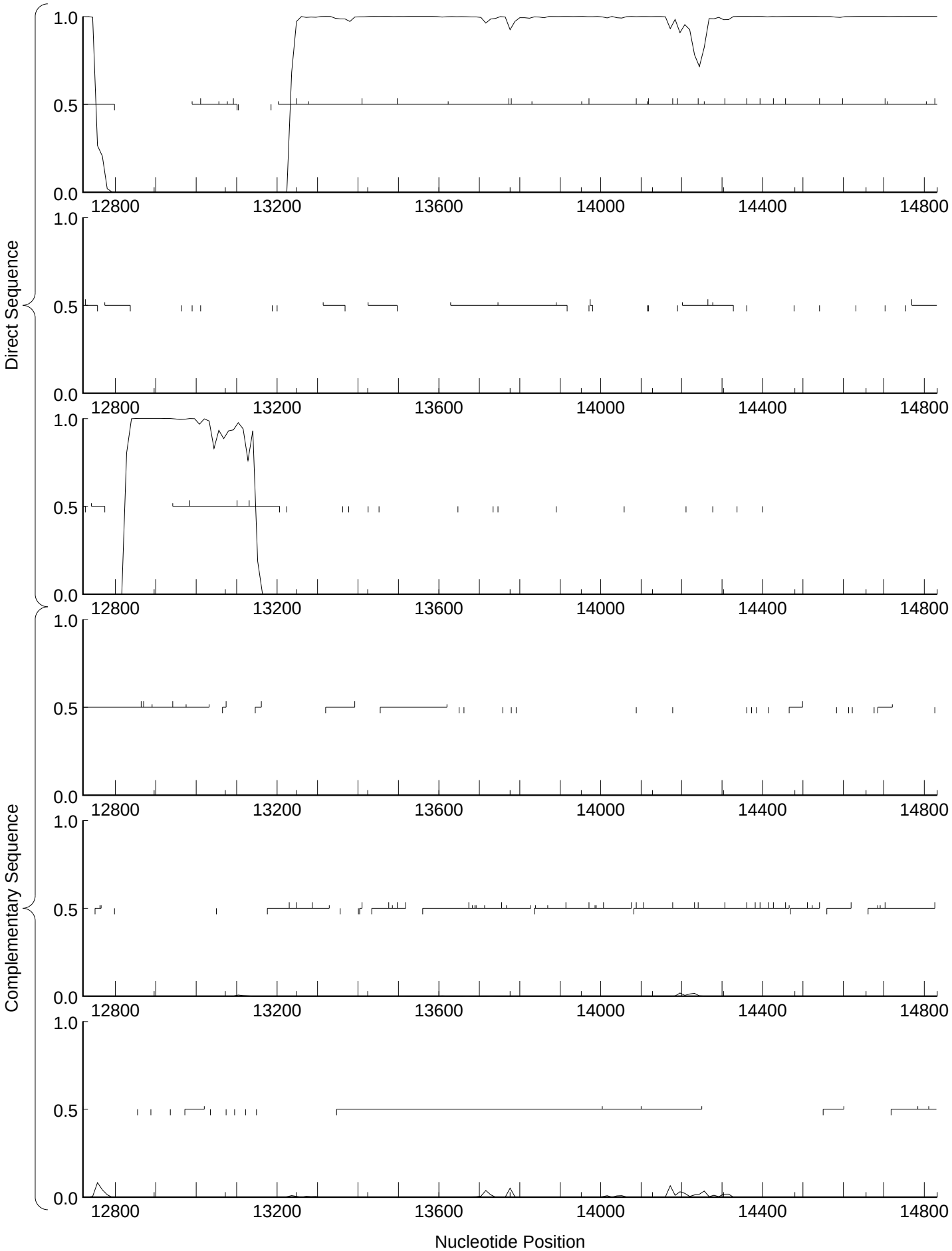
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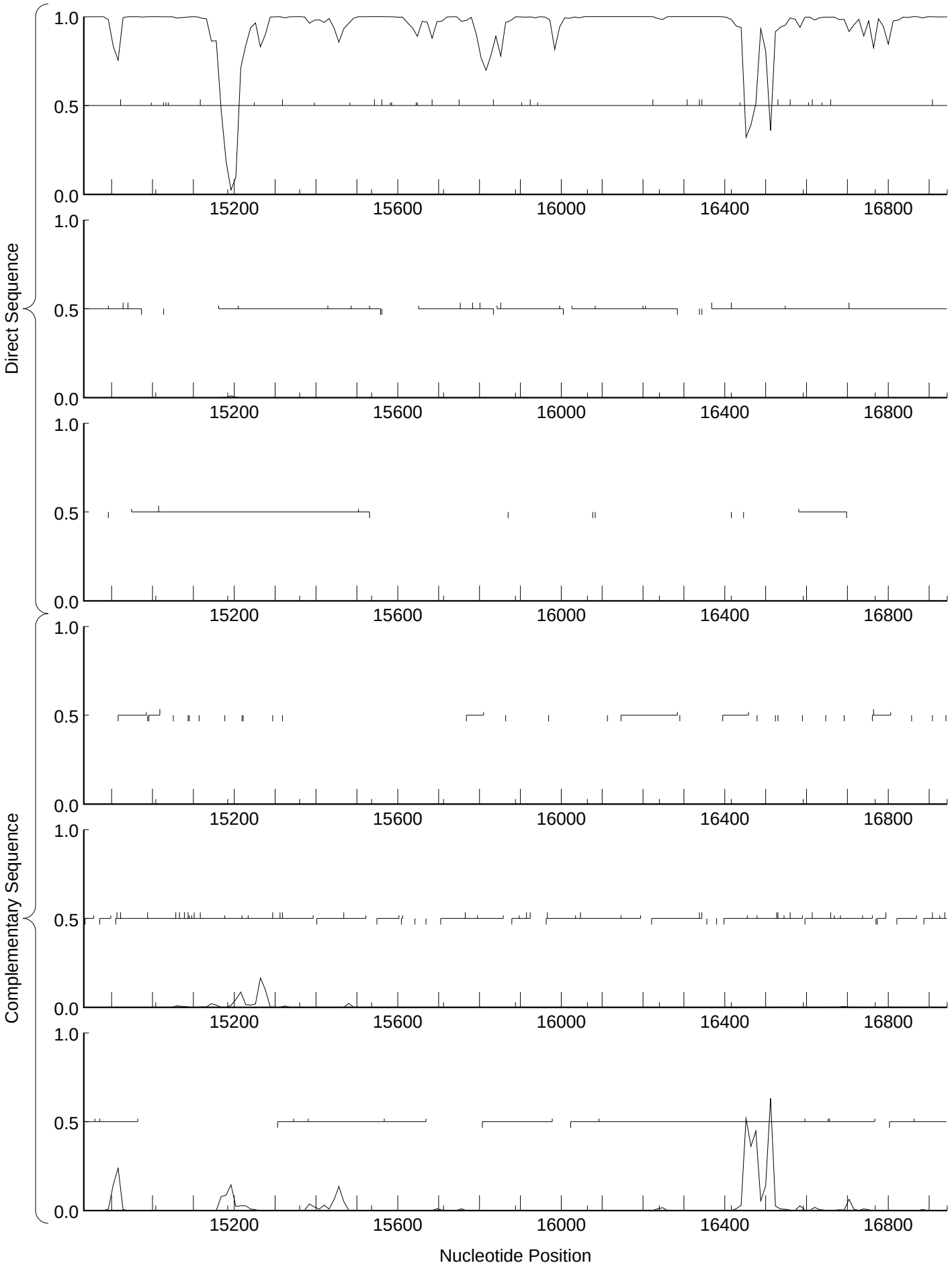
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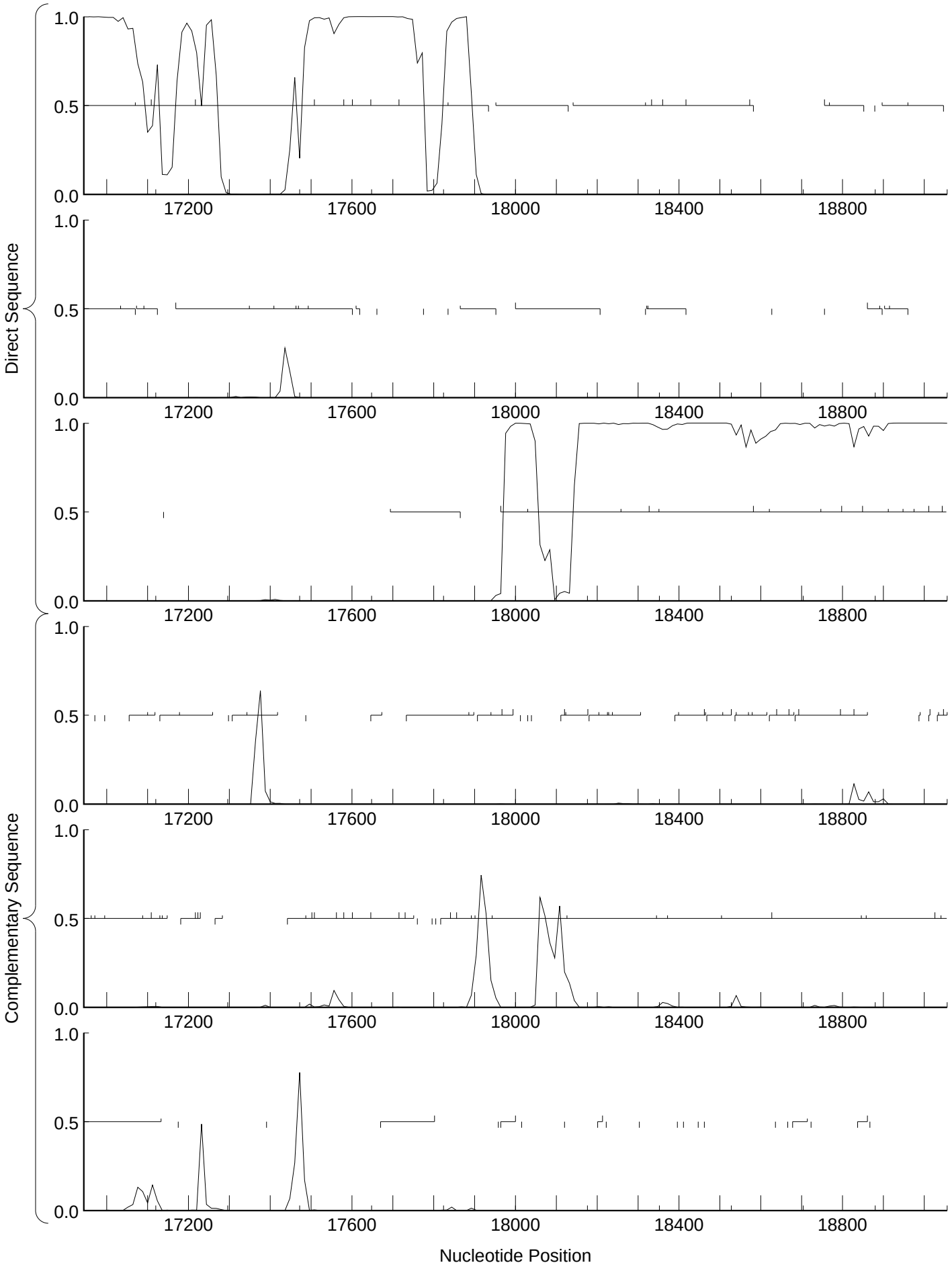
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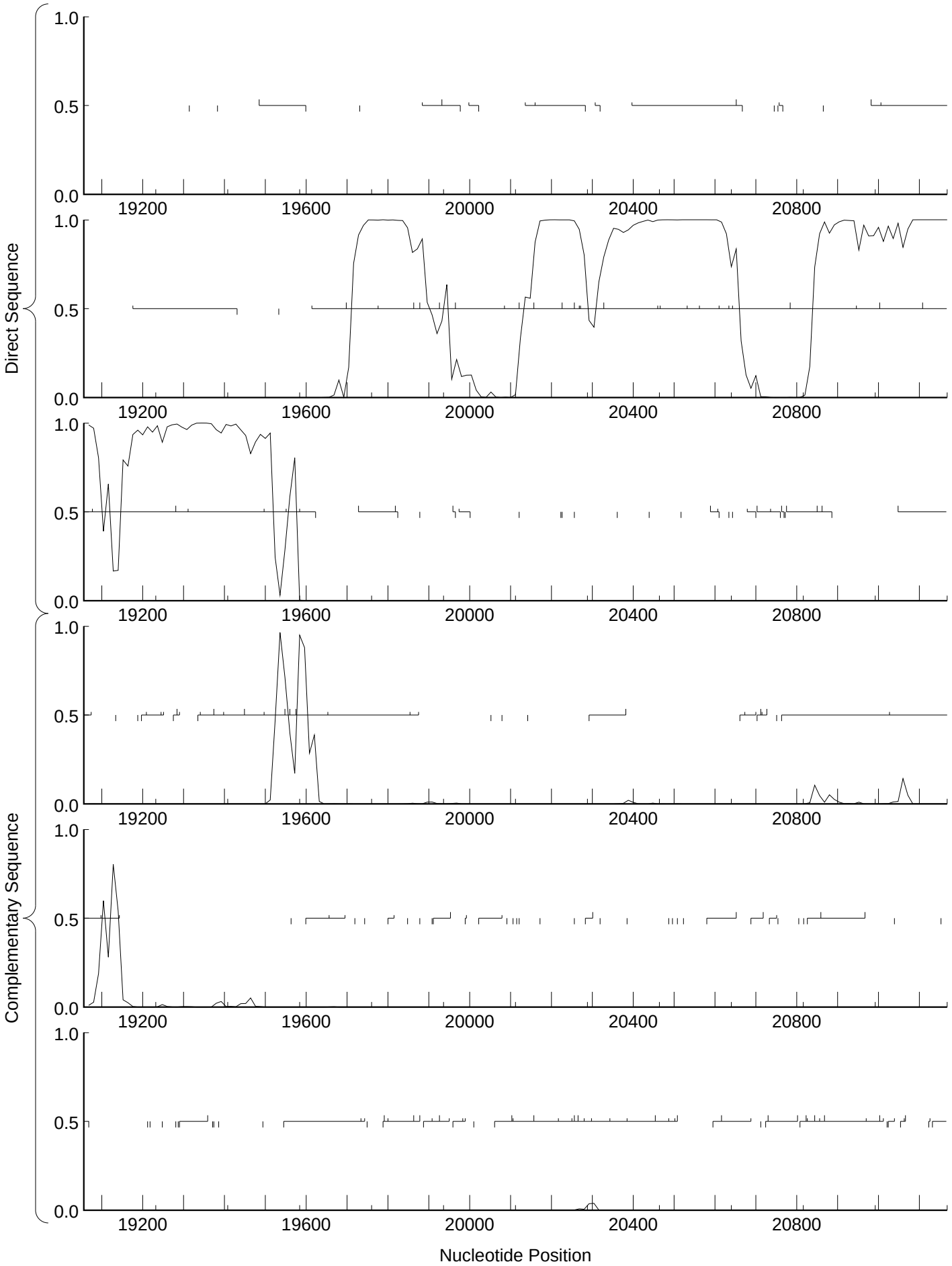
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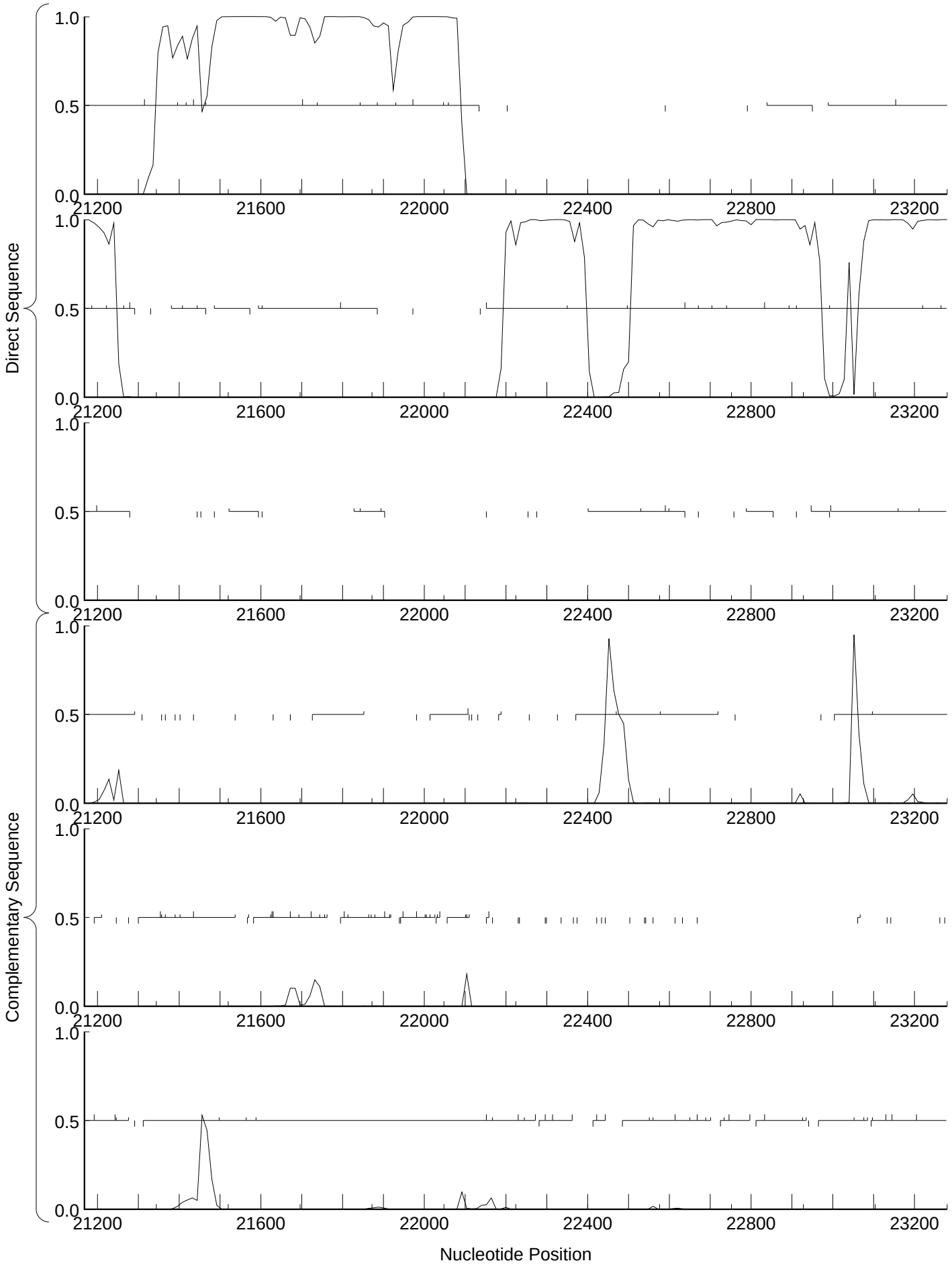
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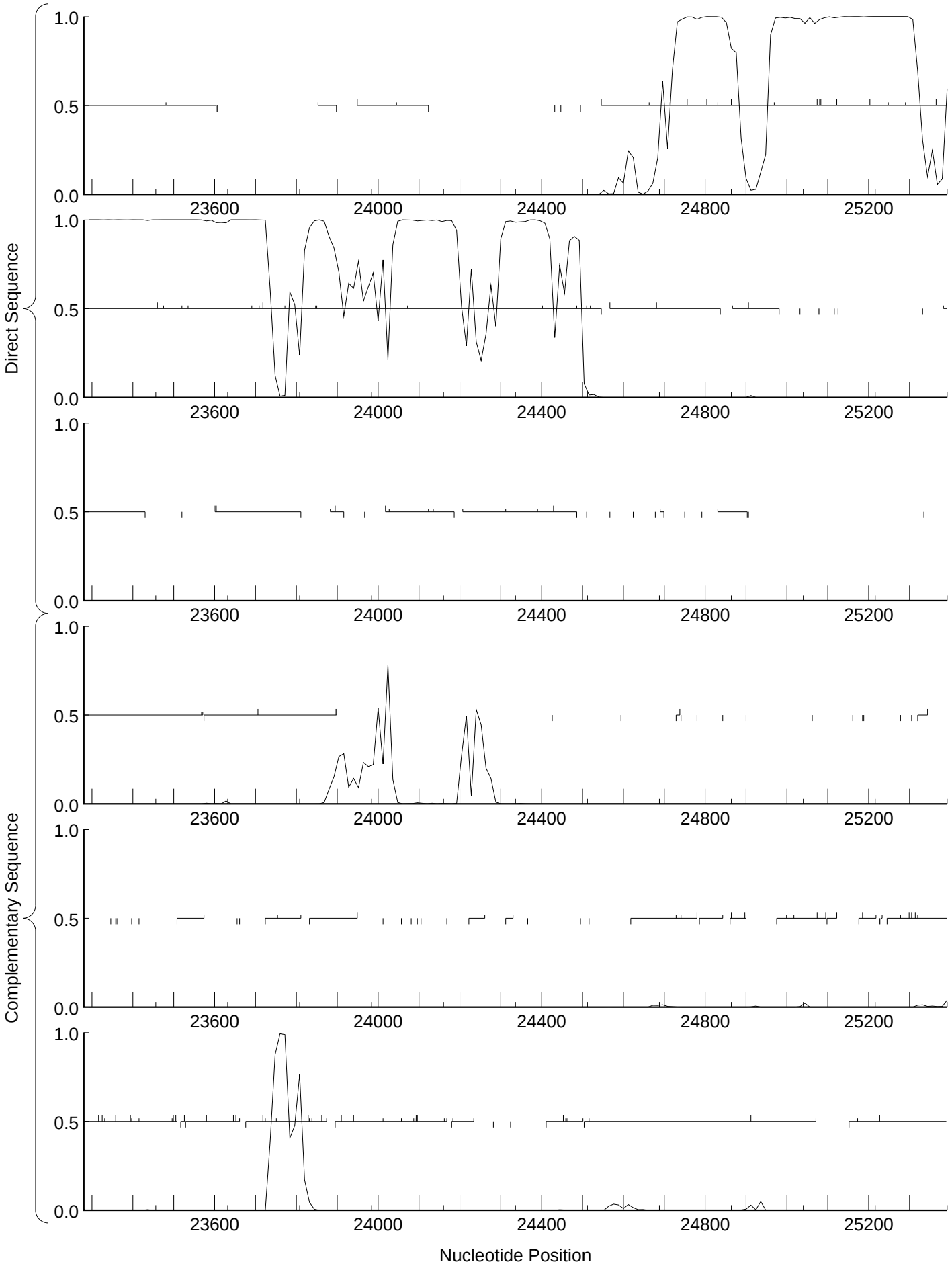
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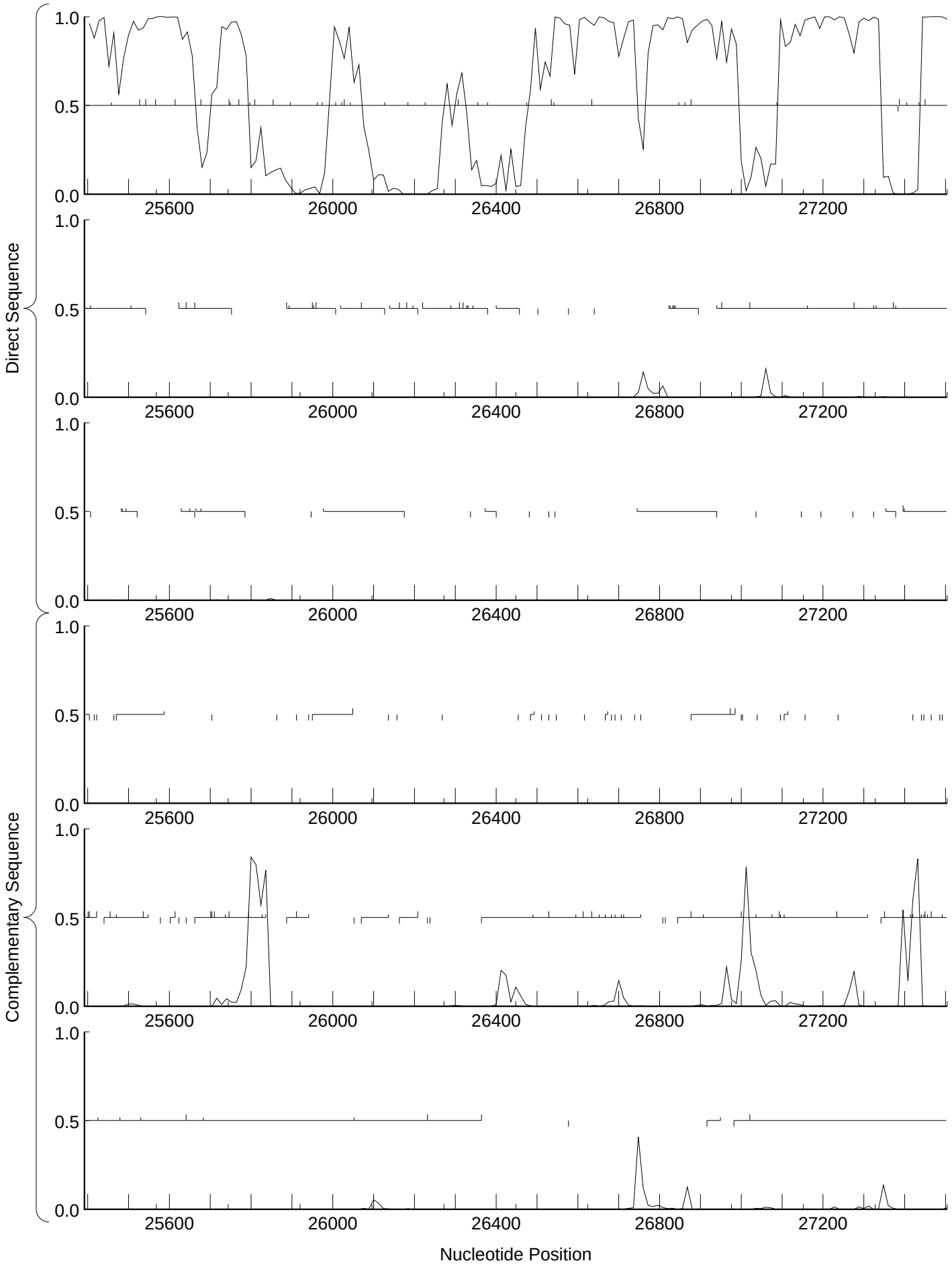
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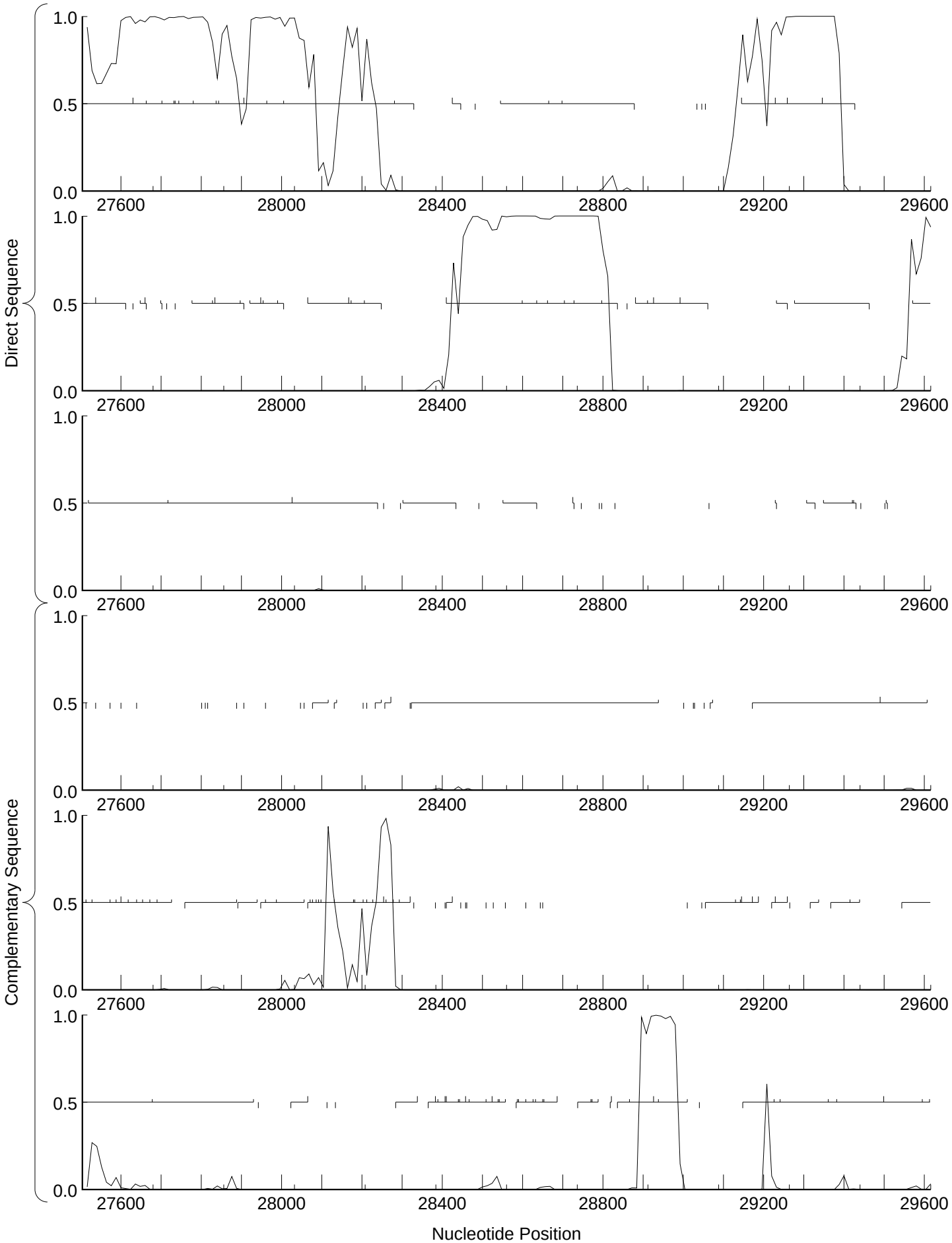
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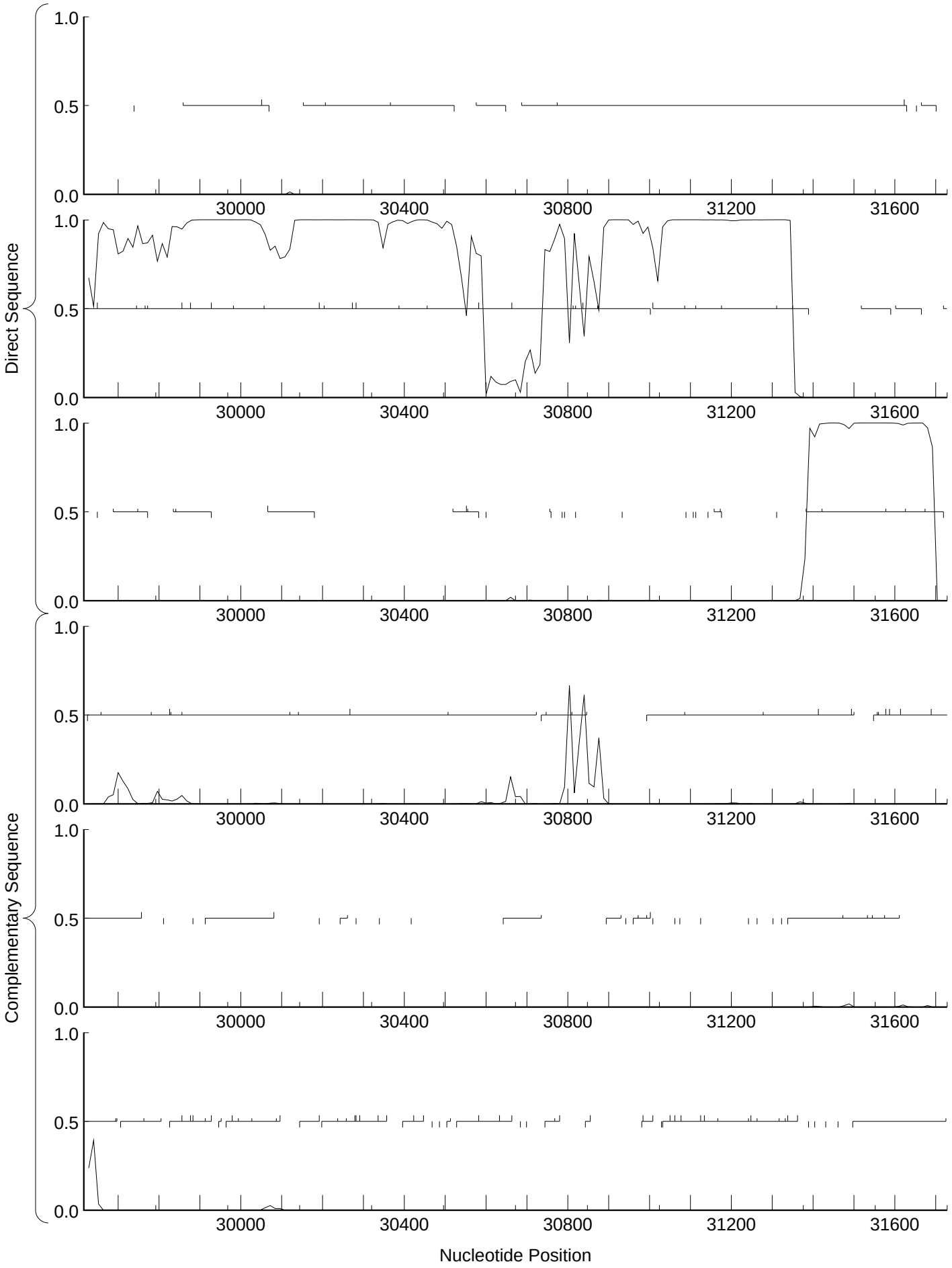
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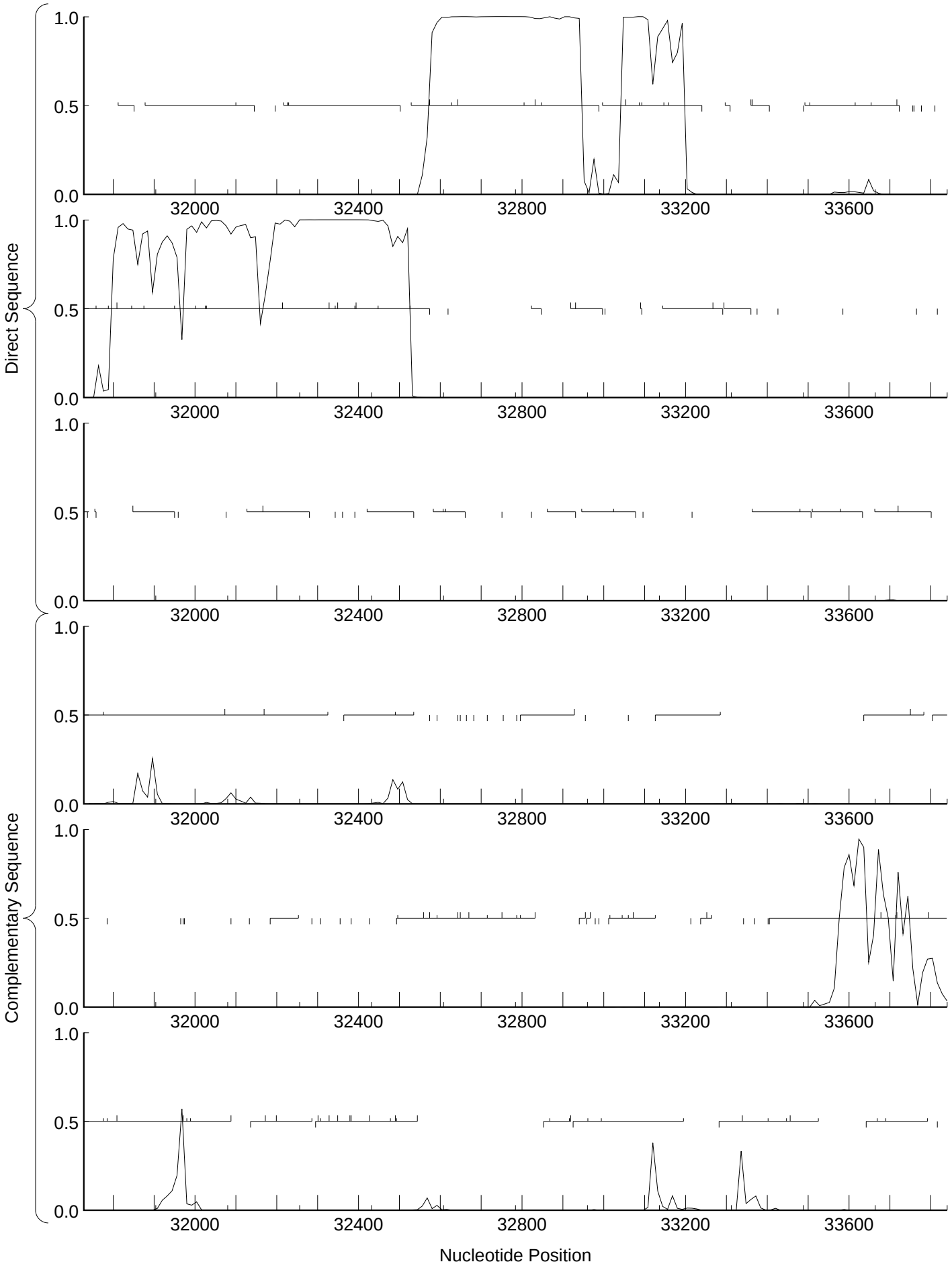
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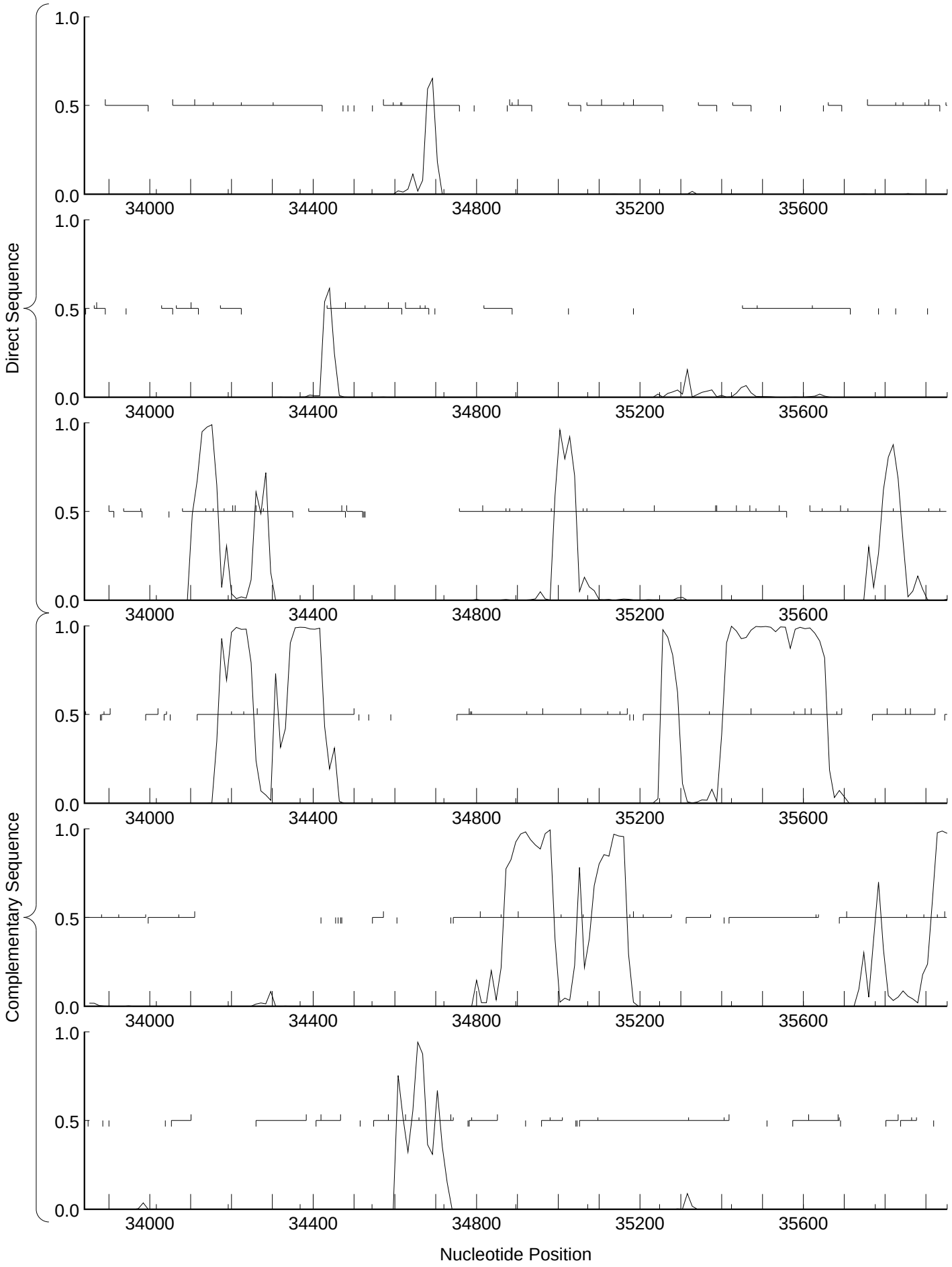
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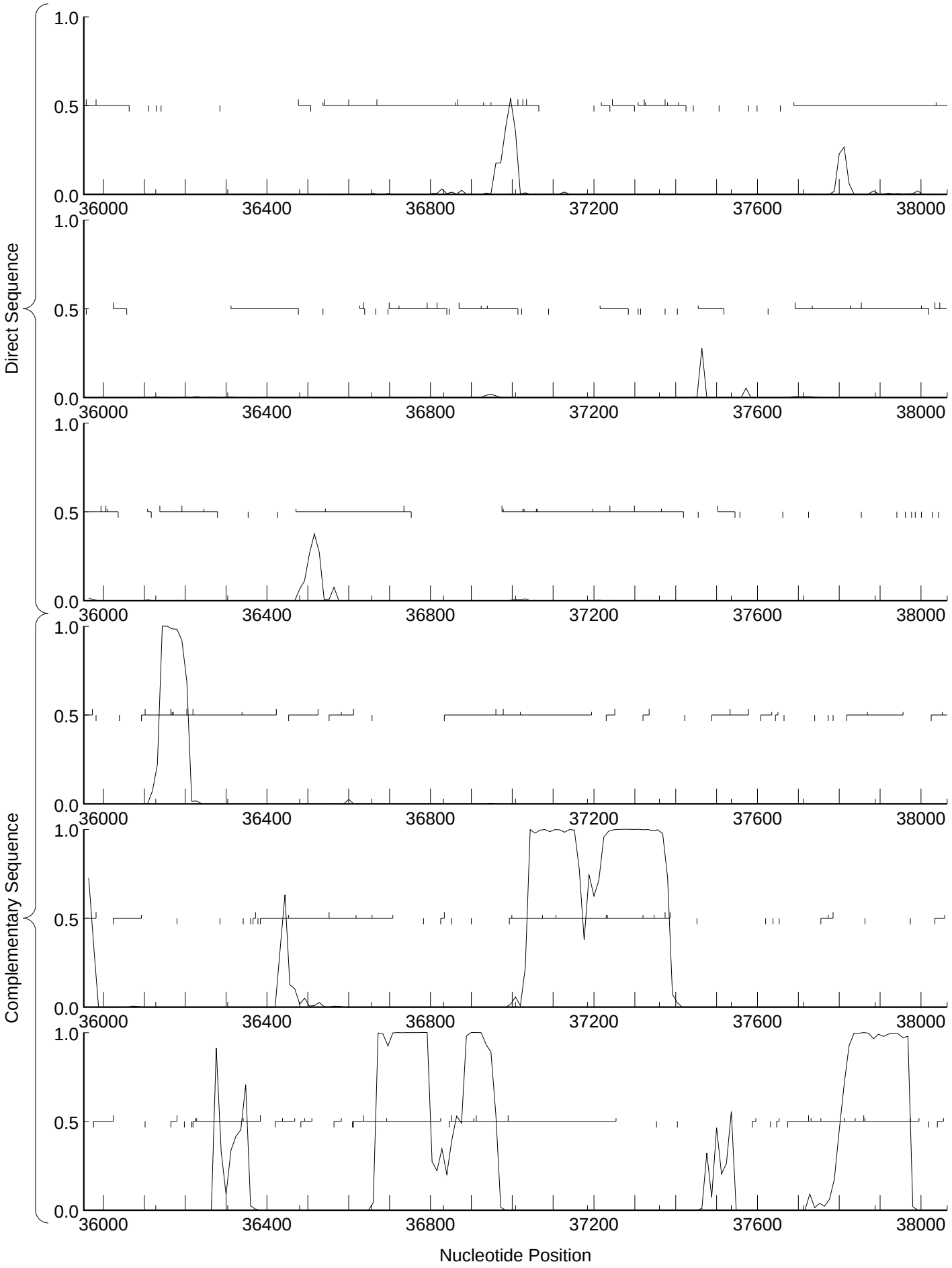
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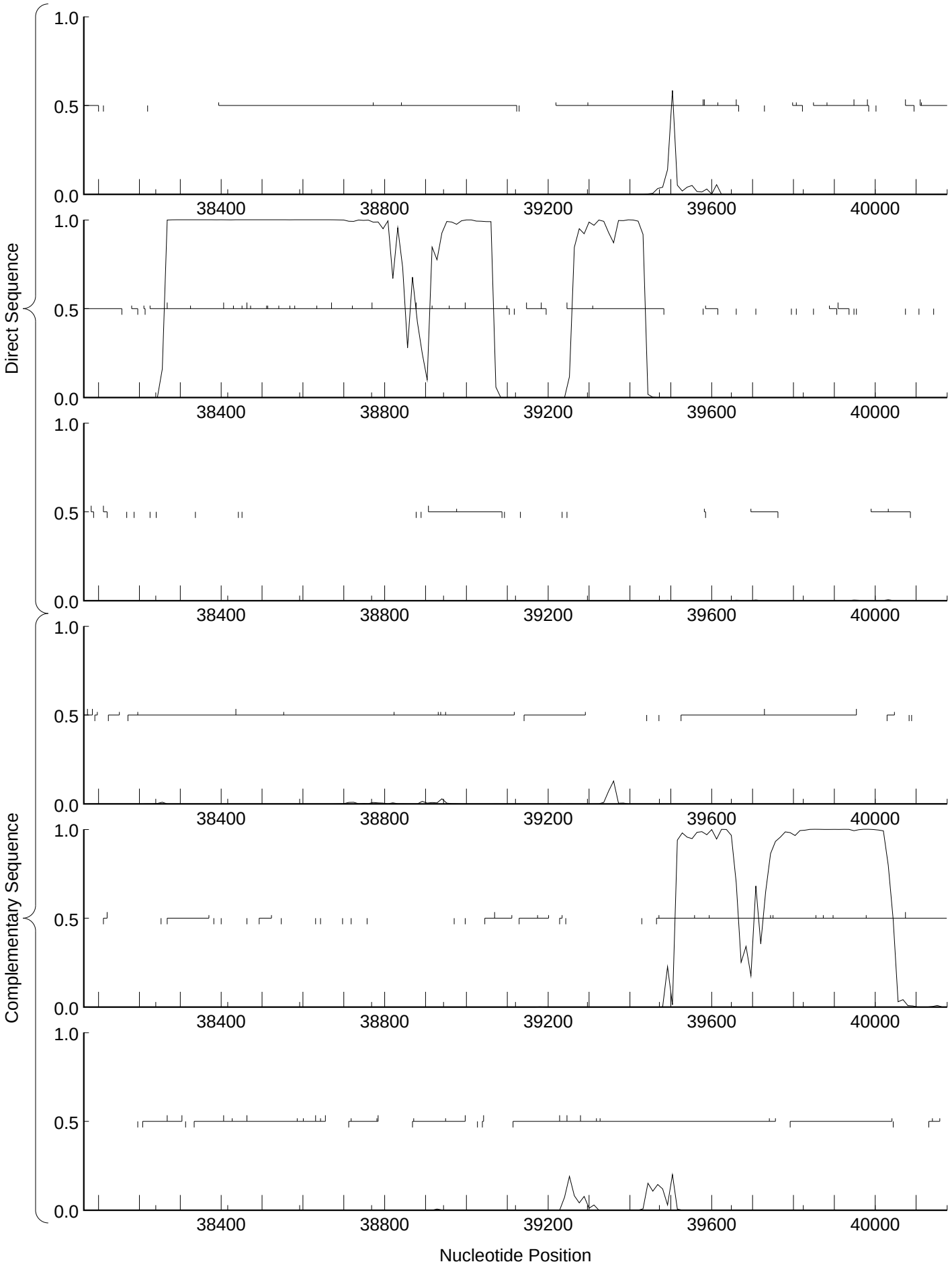
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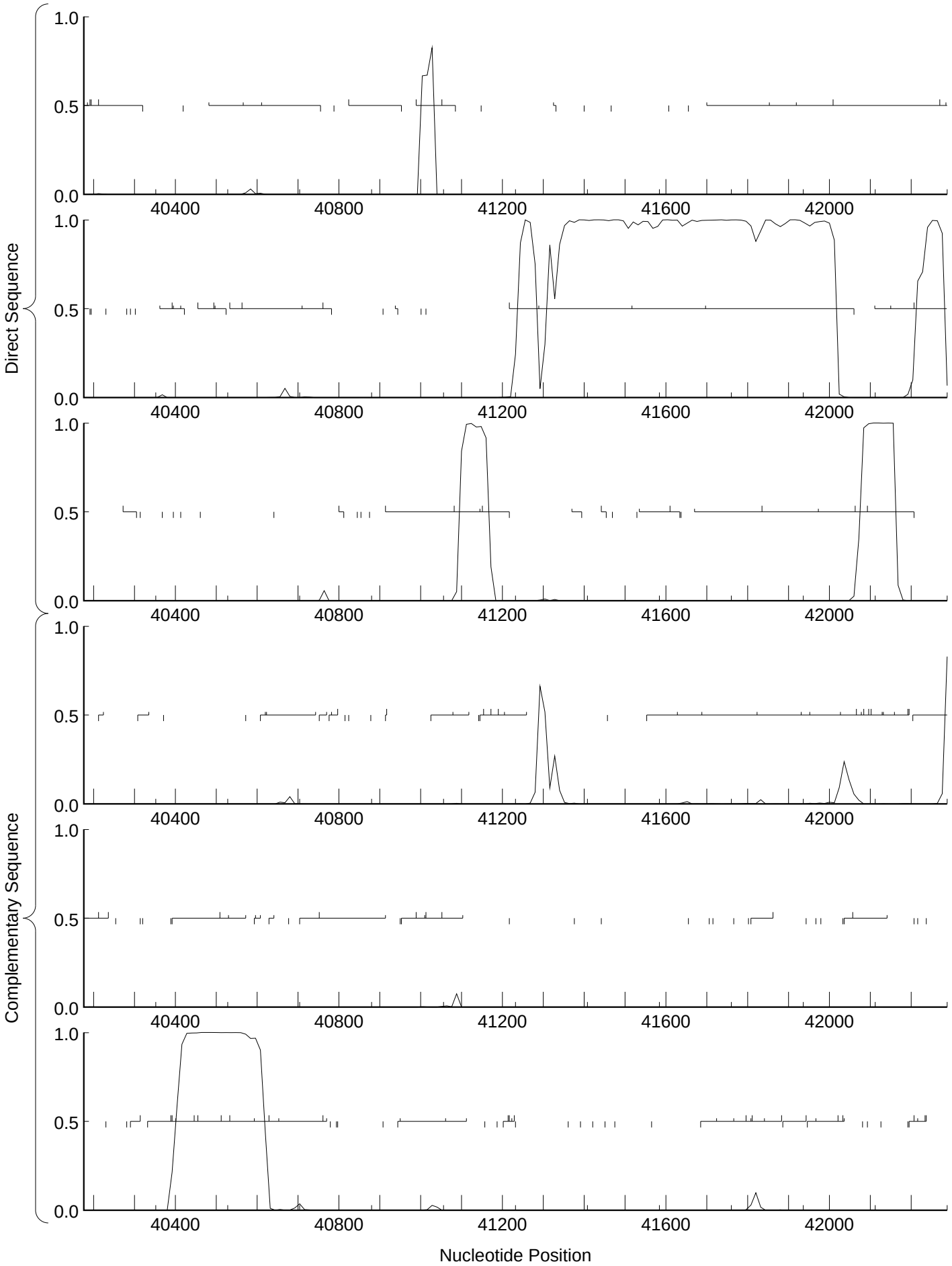
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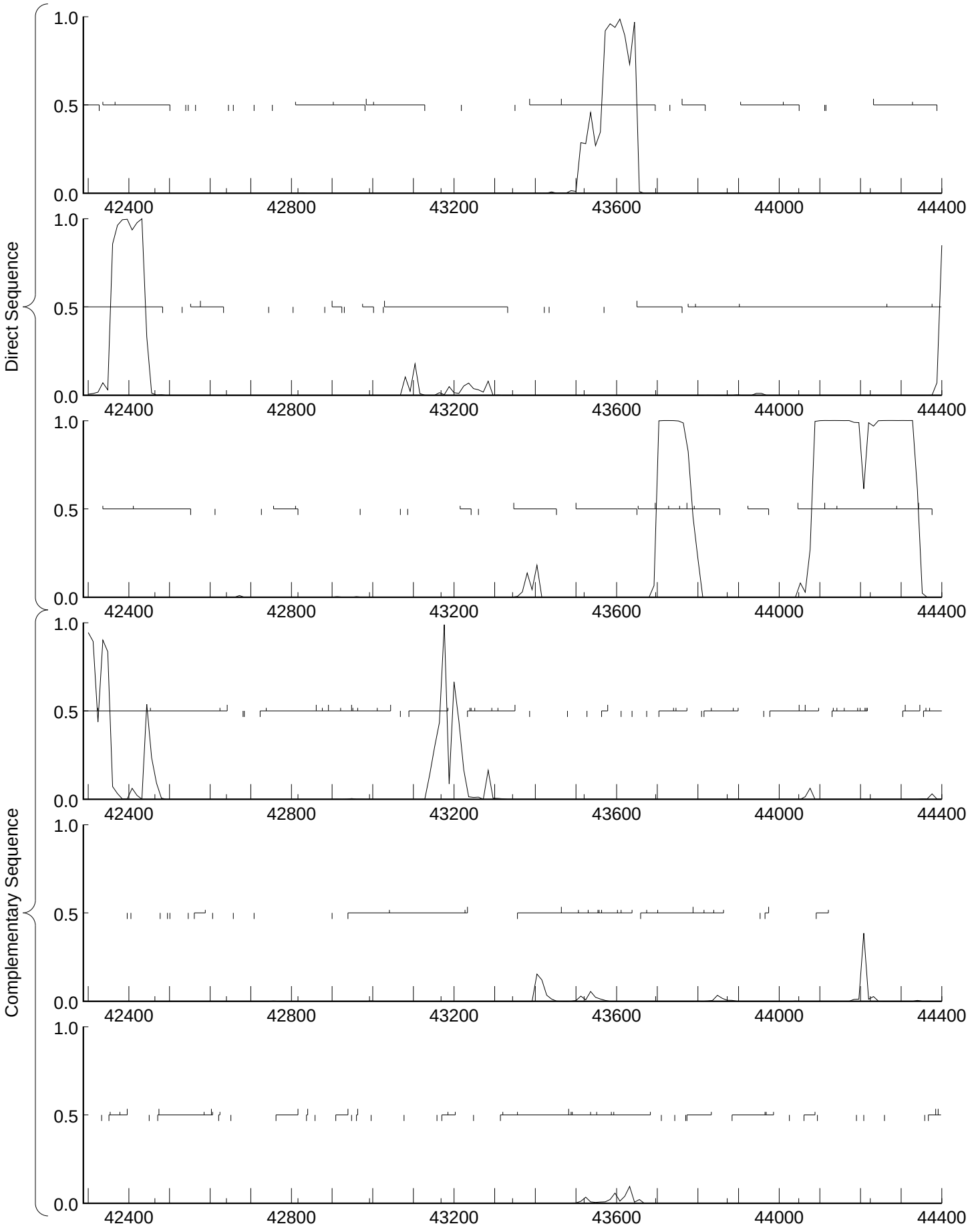


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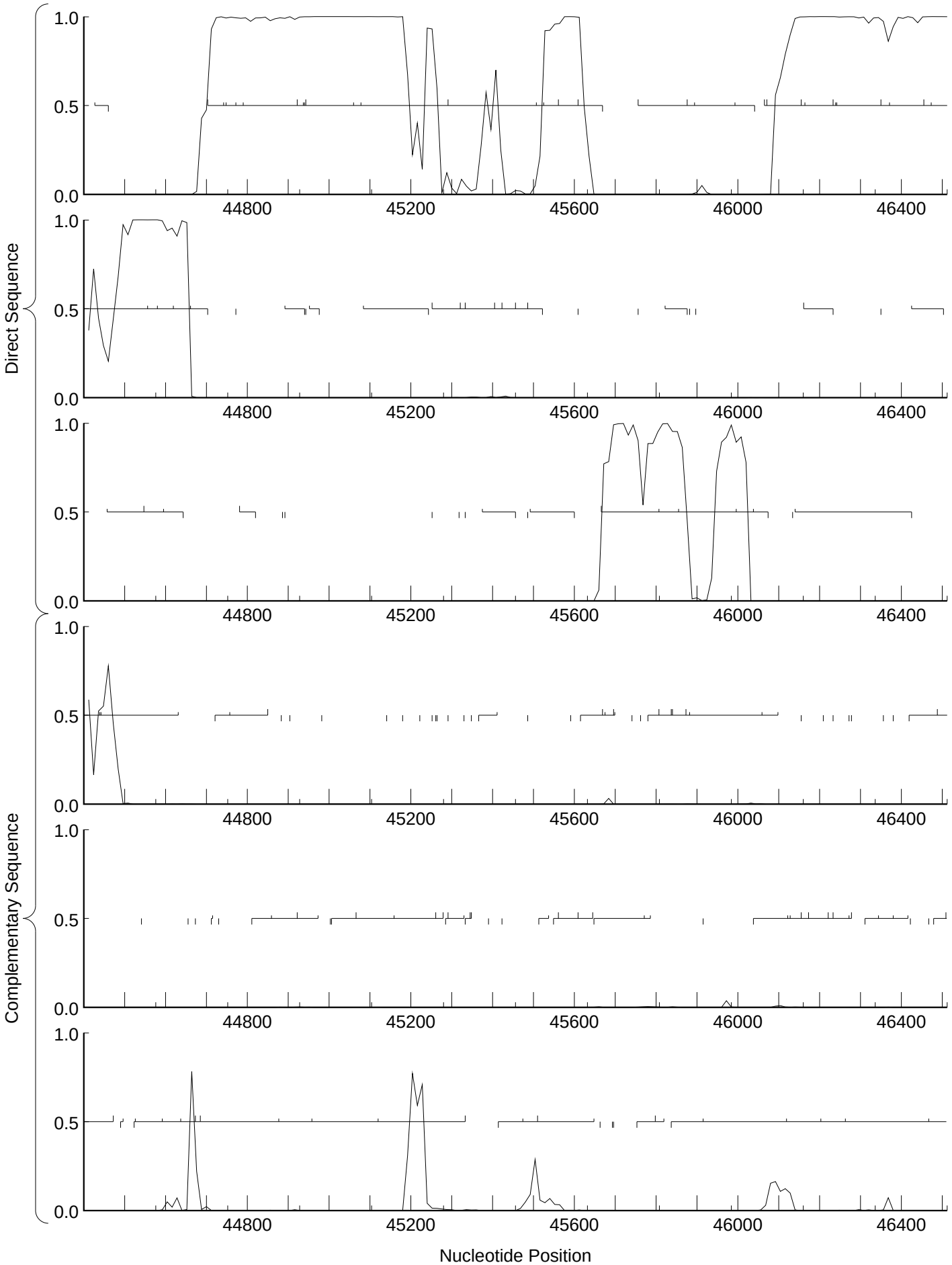


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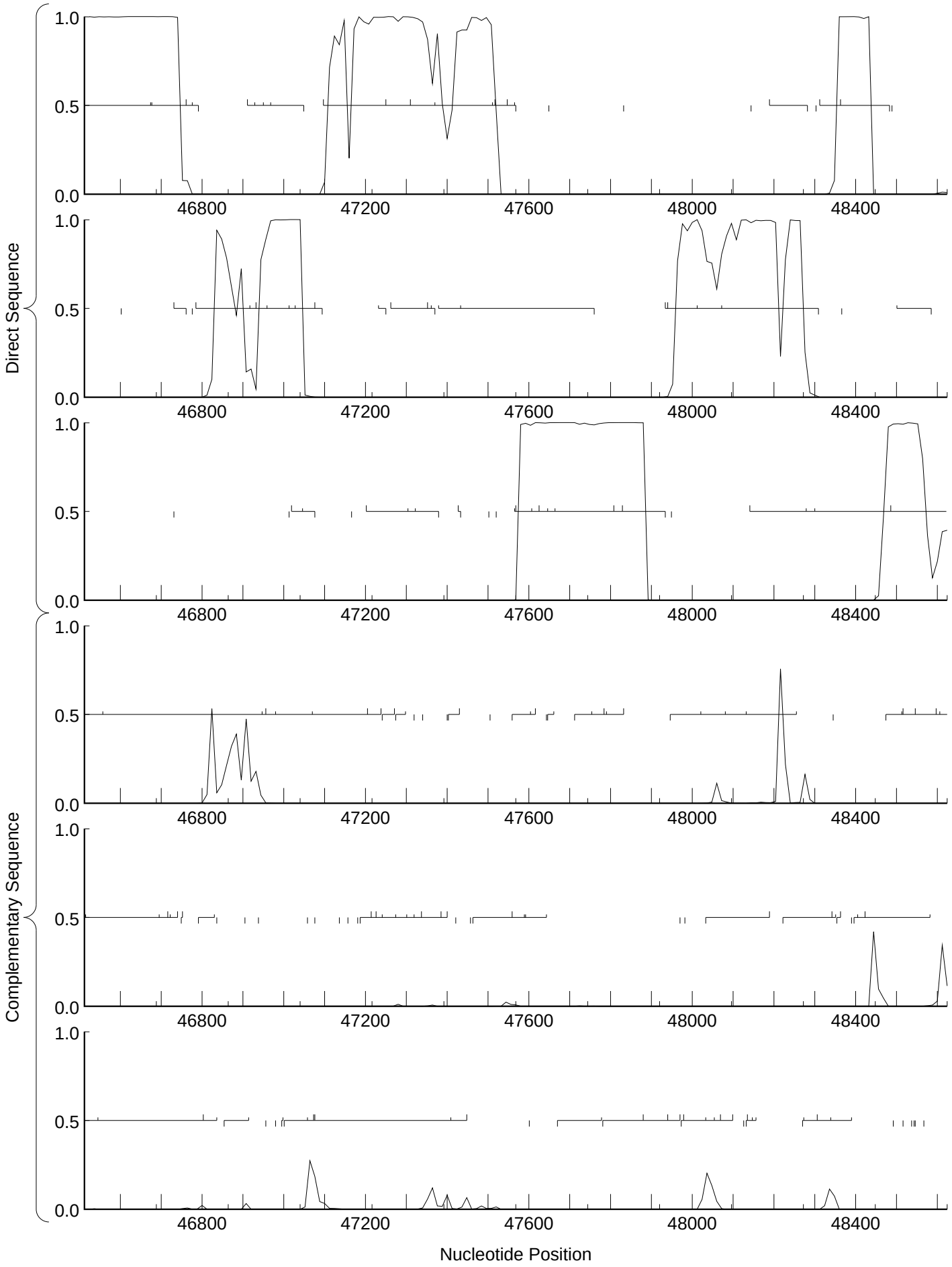



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44000
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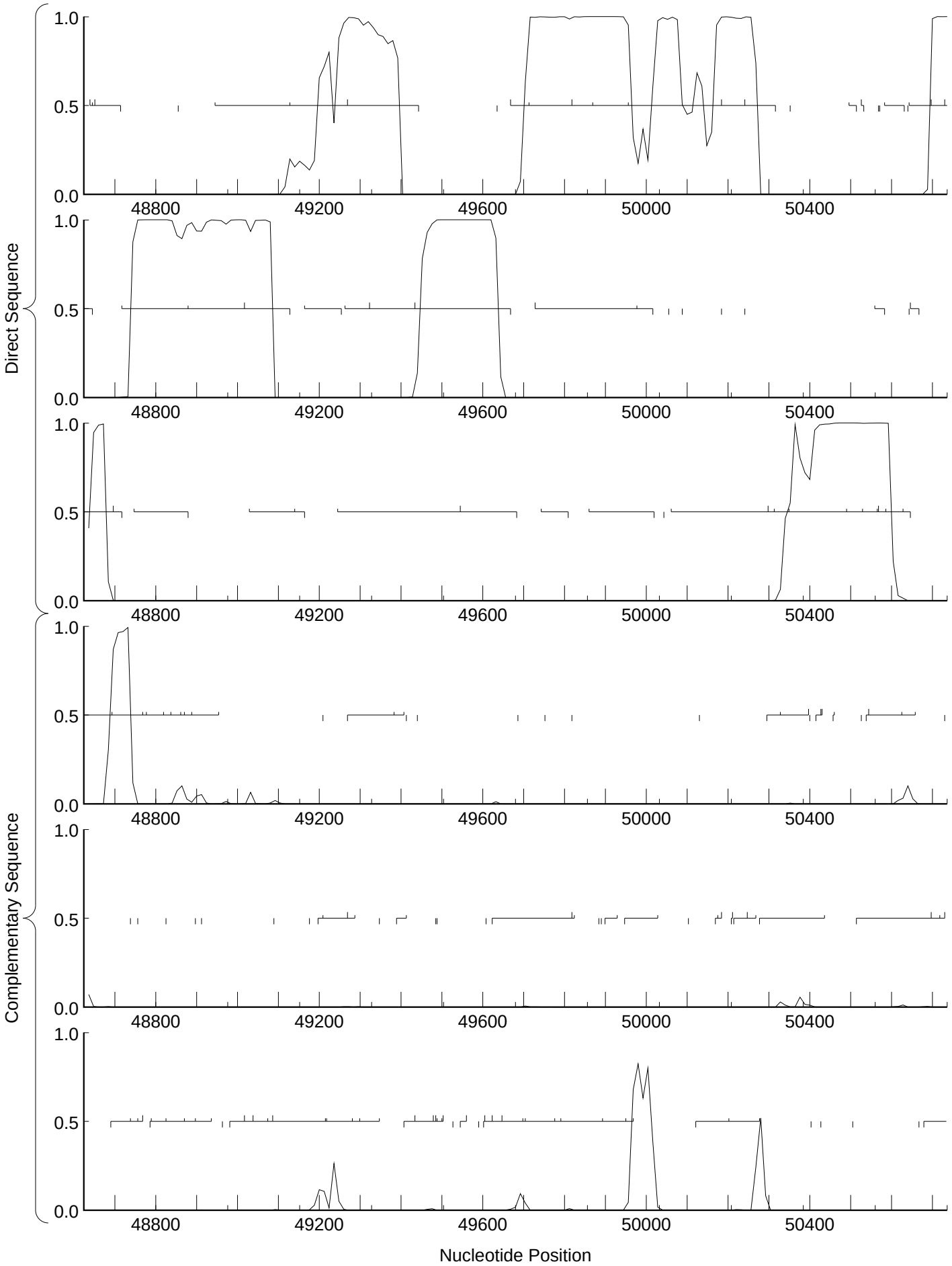
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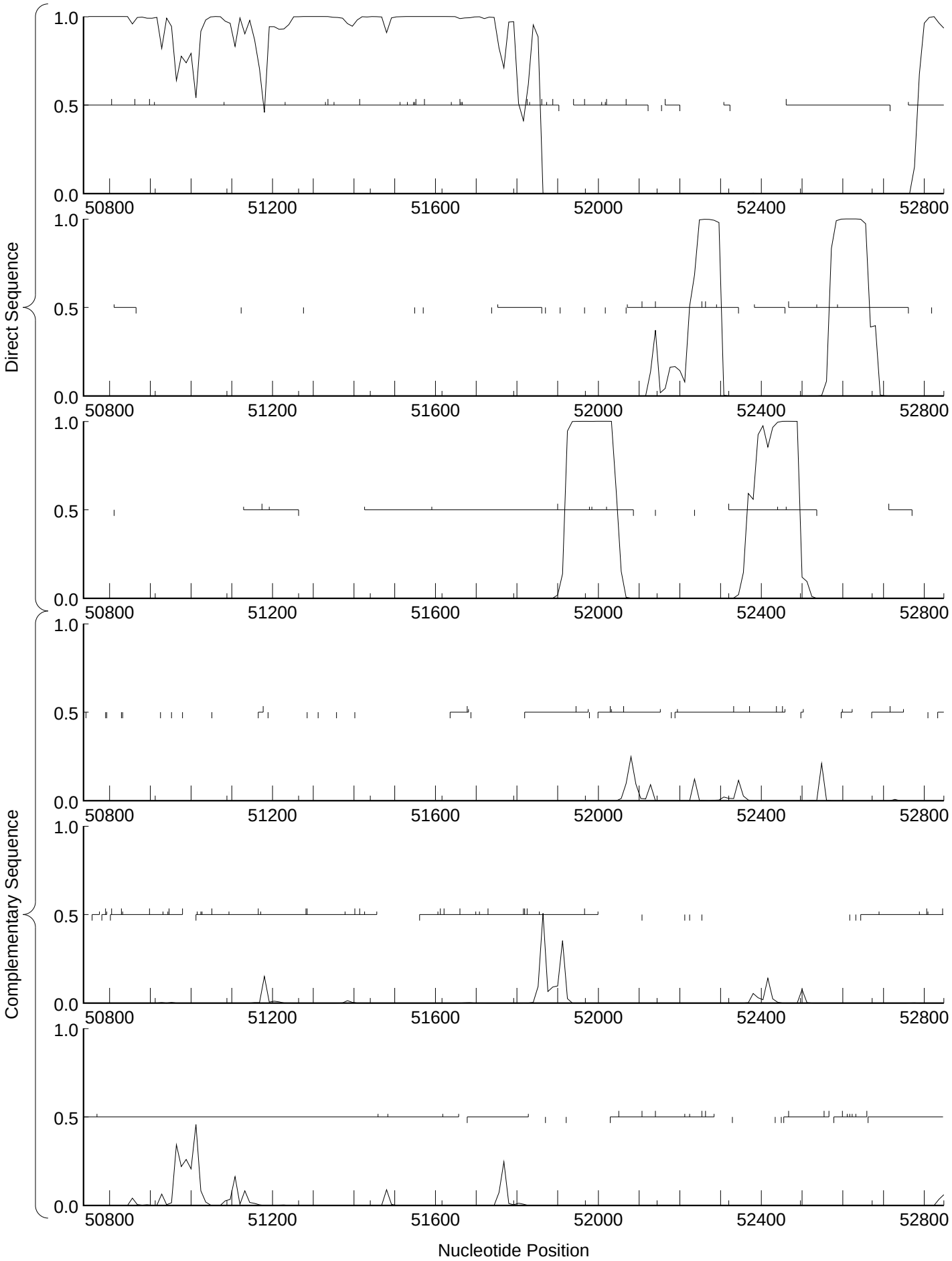
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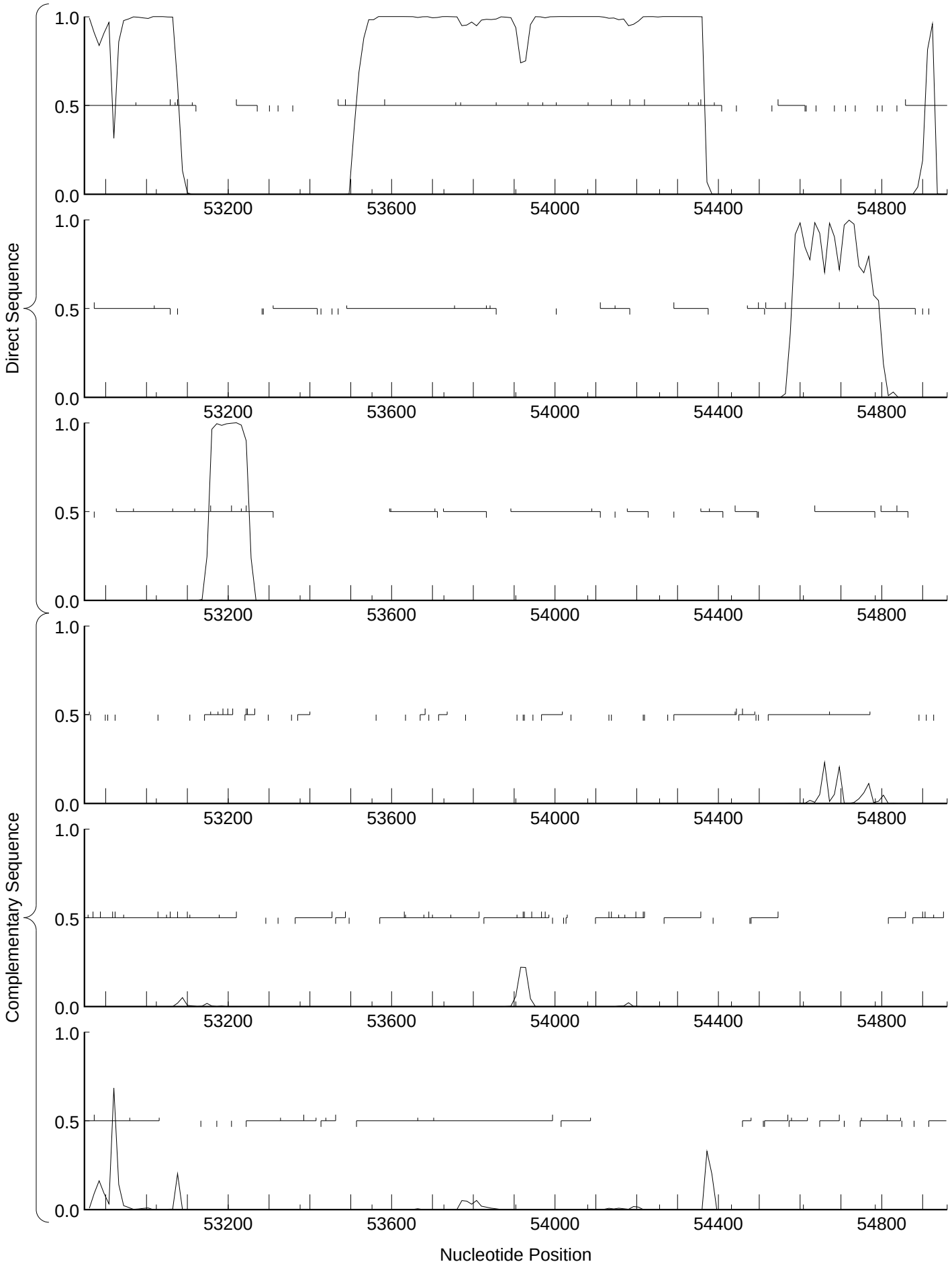
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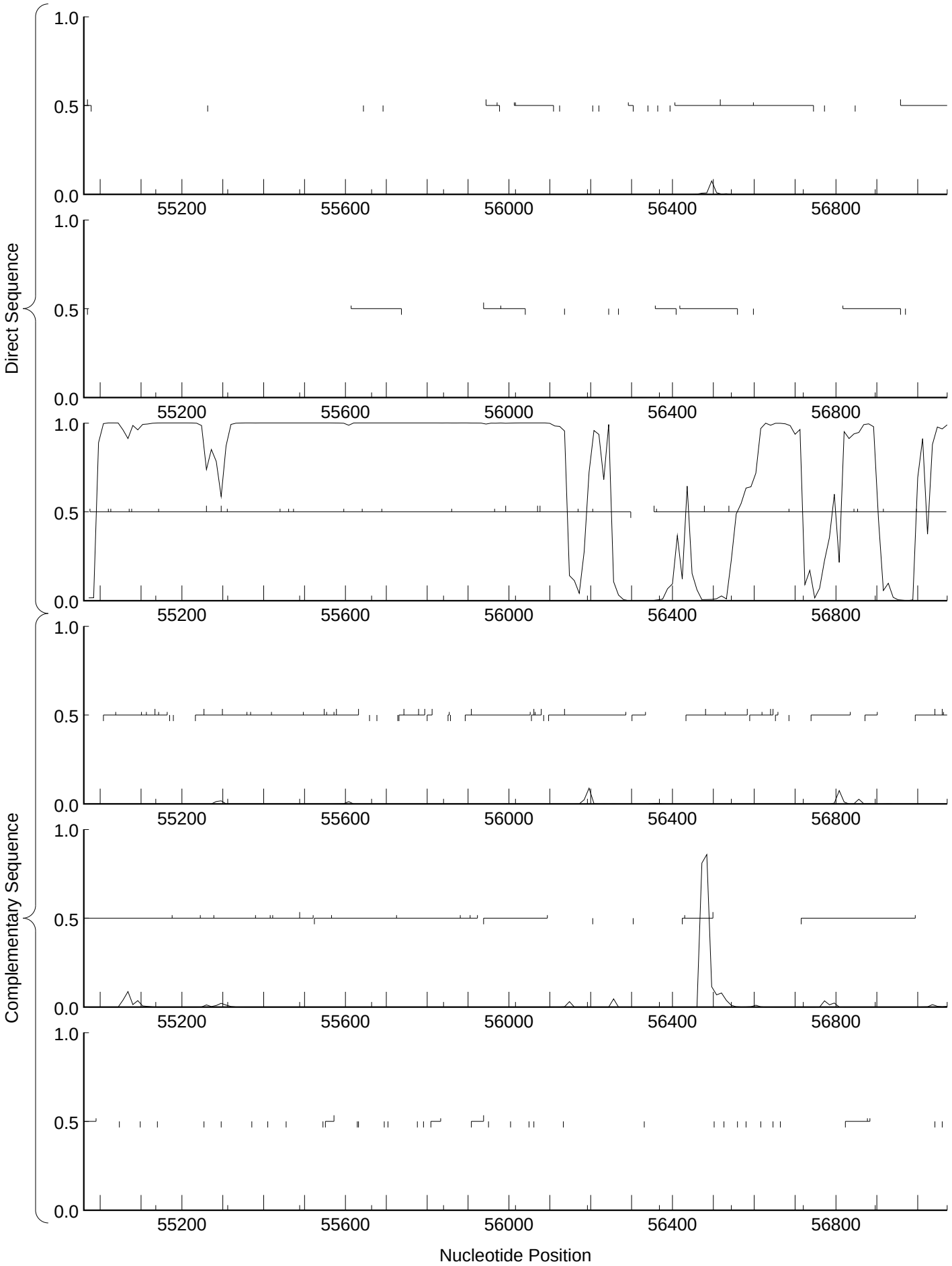
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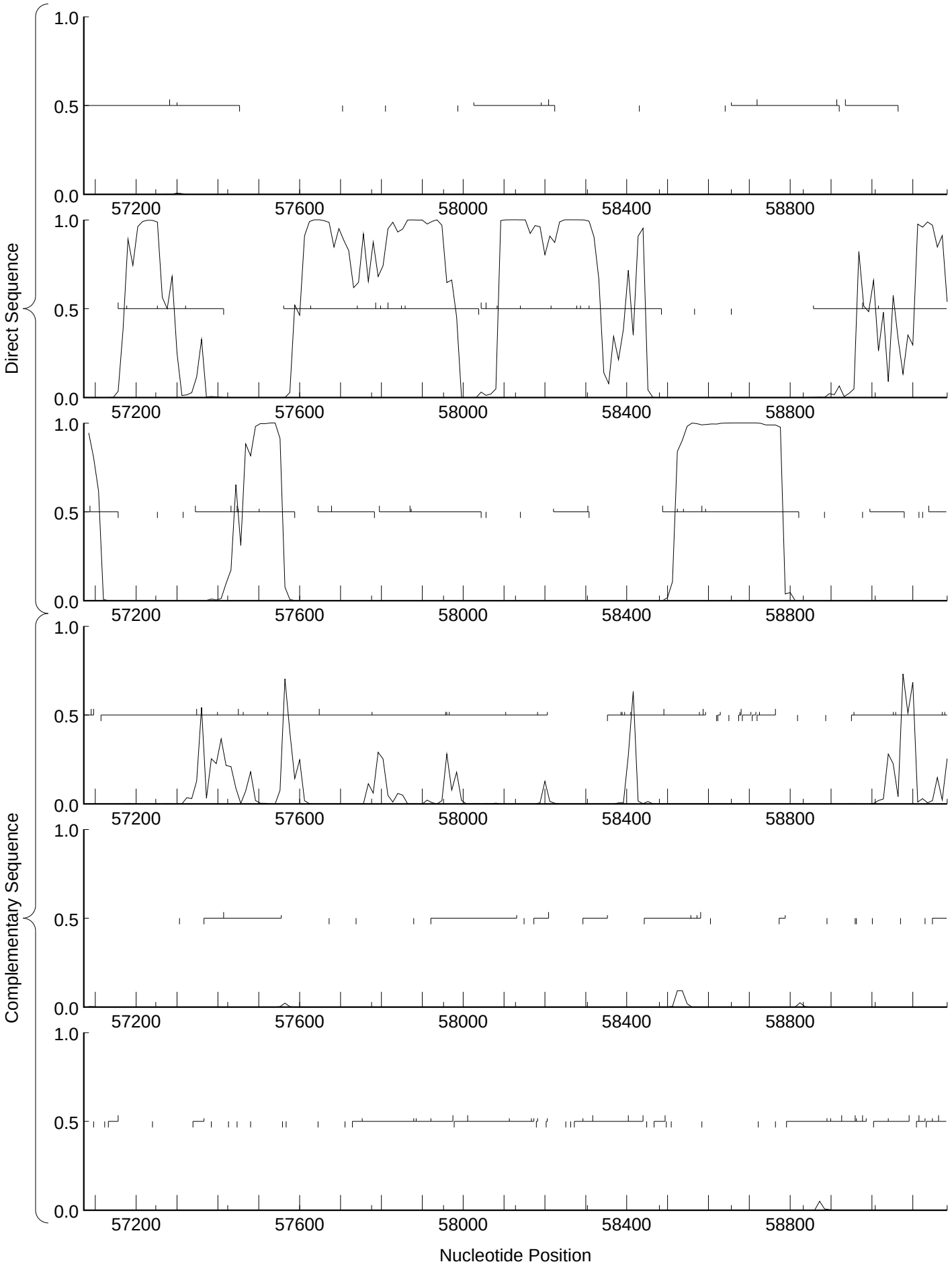
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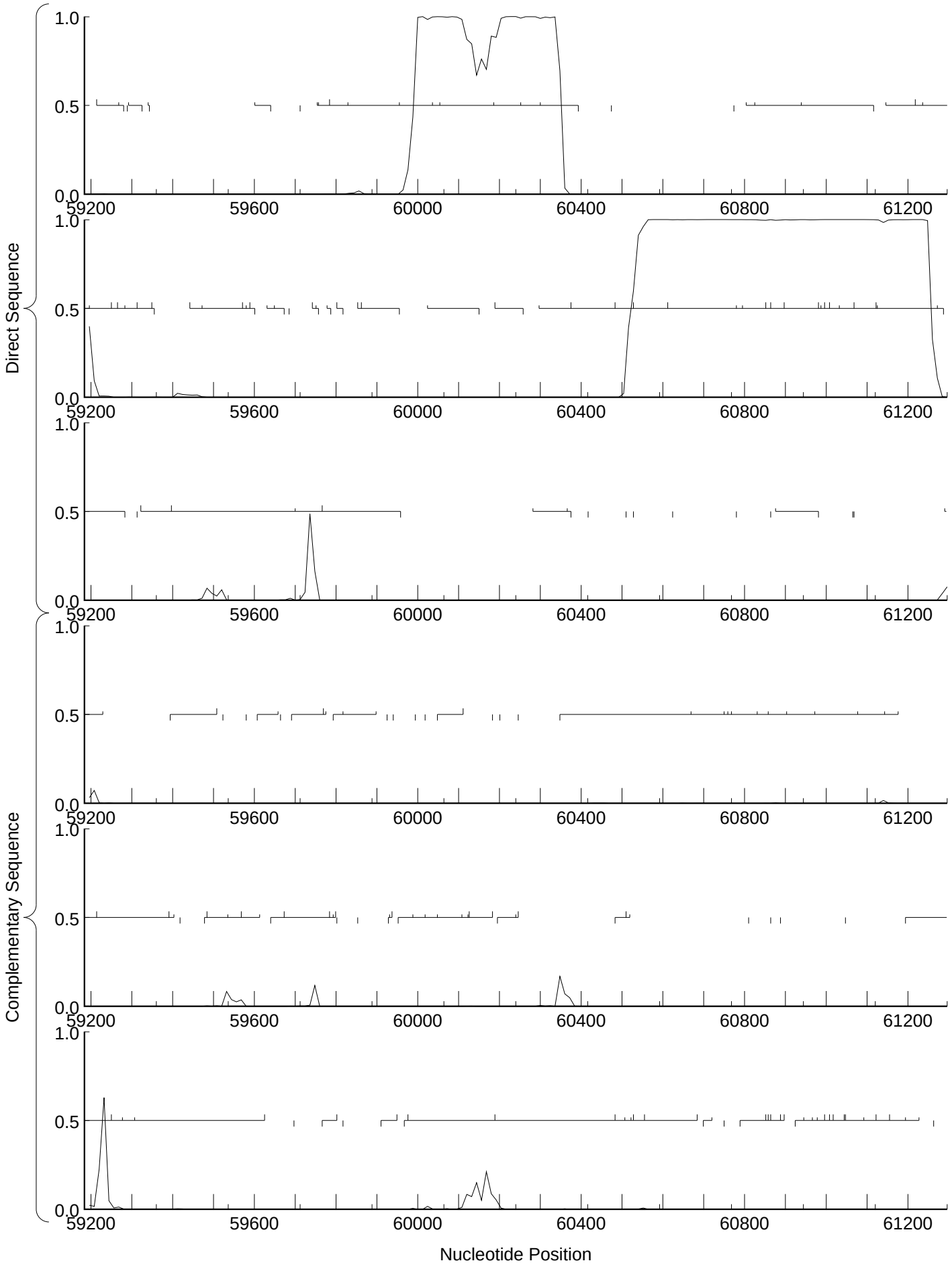
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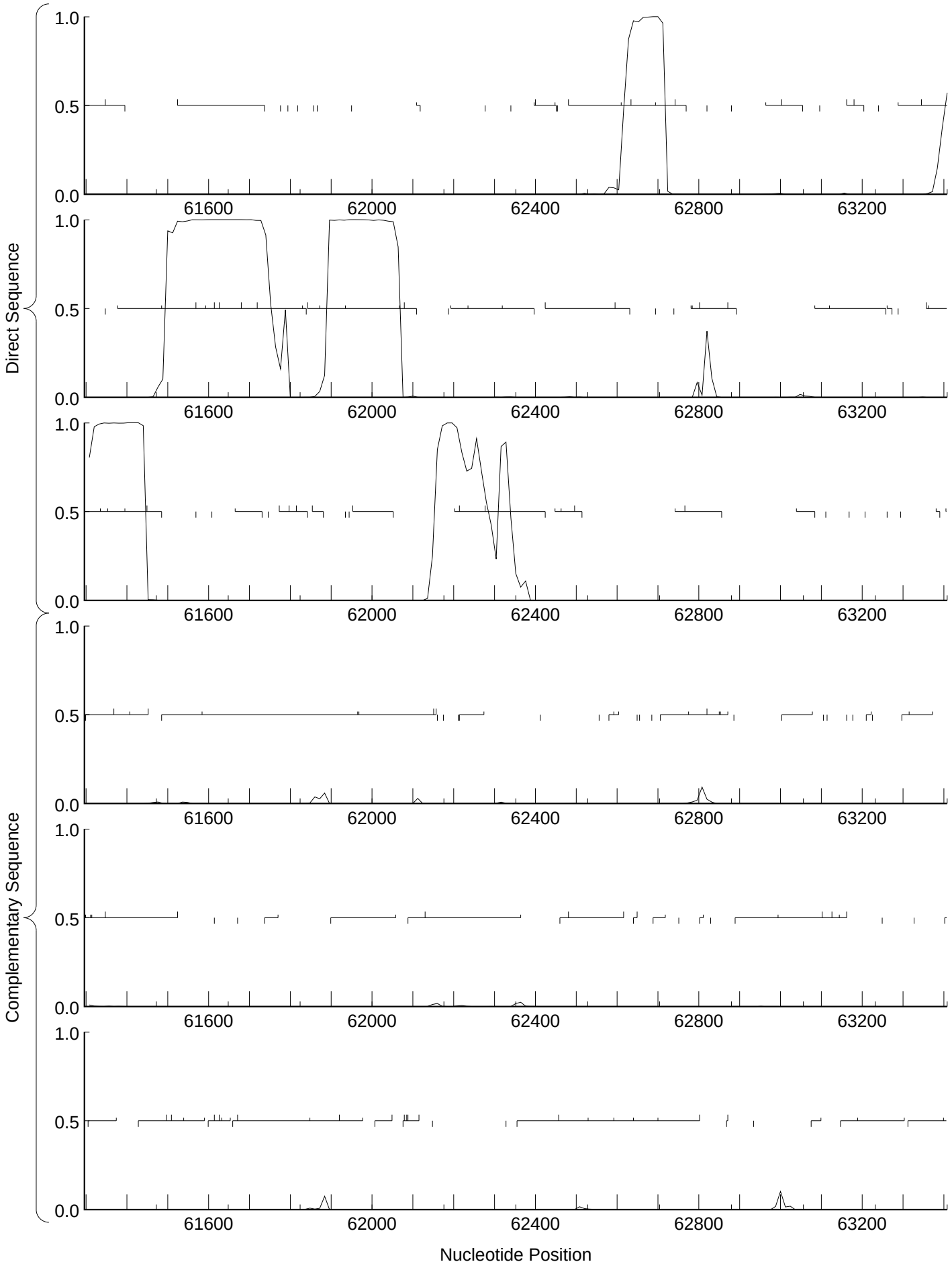
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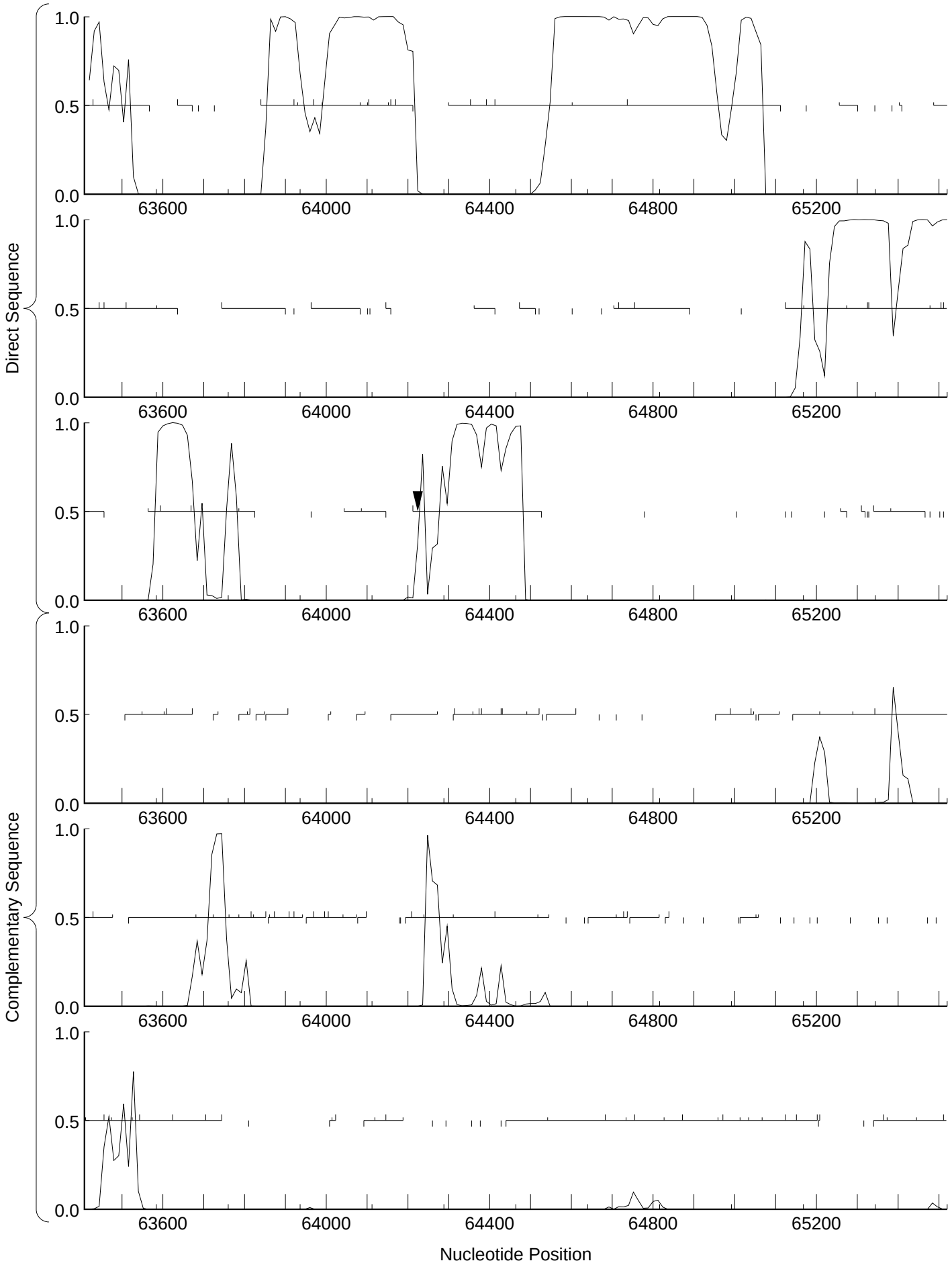
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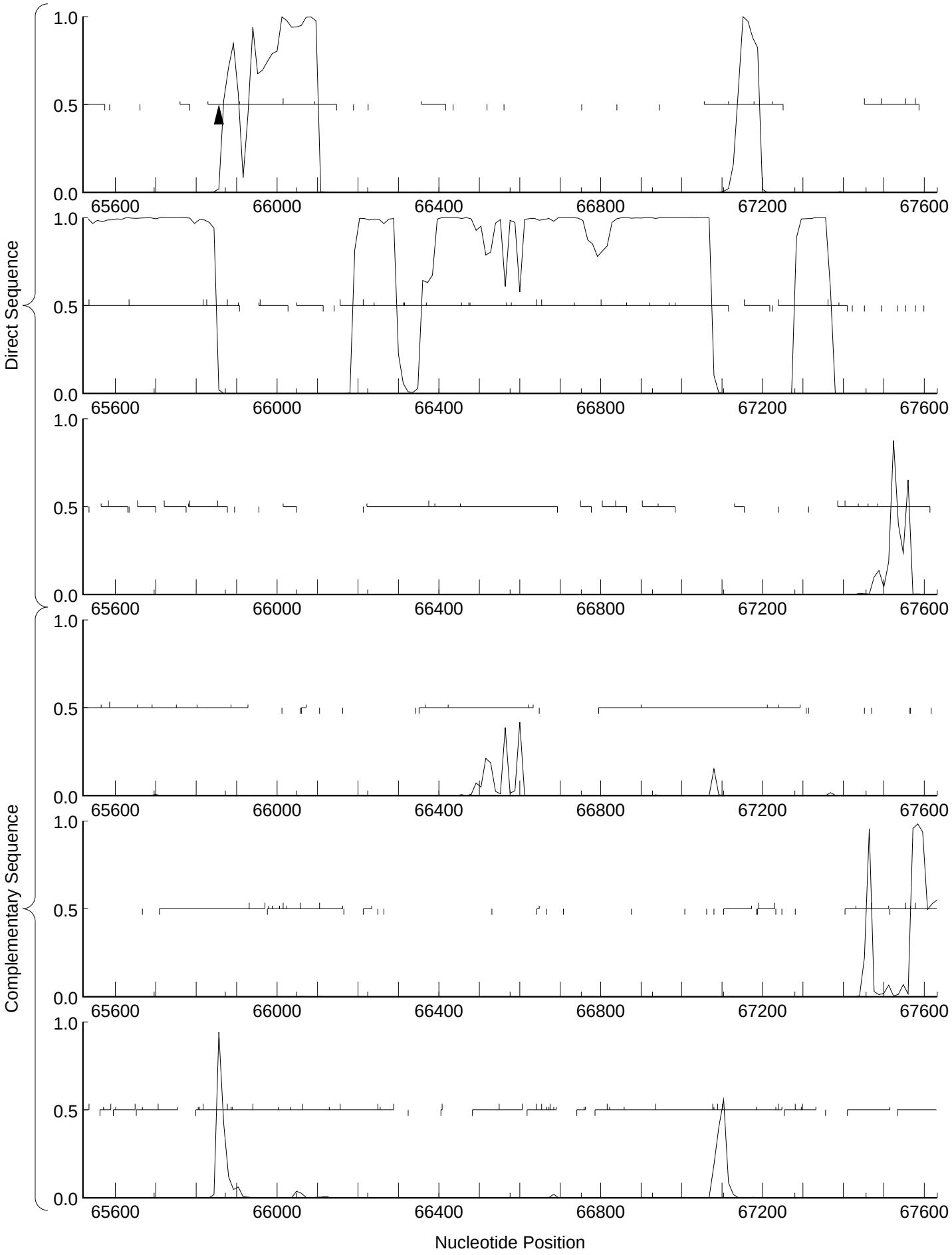
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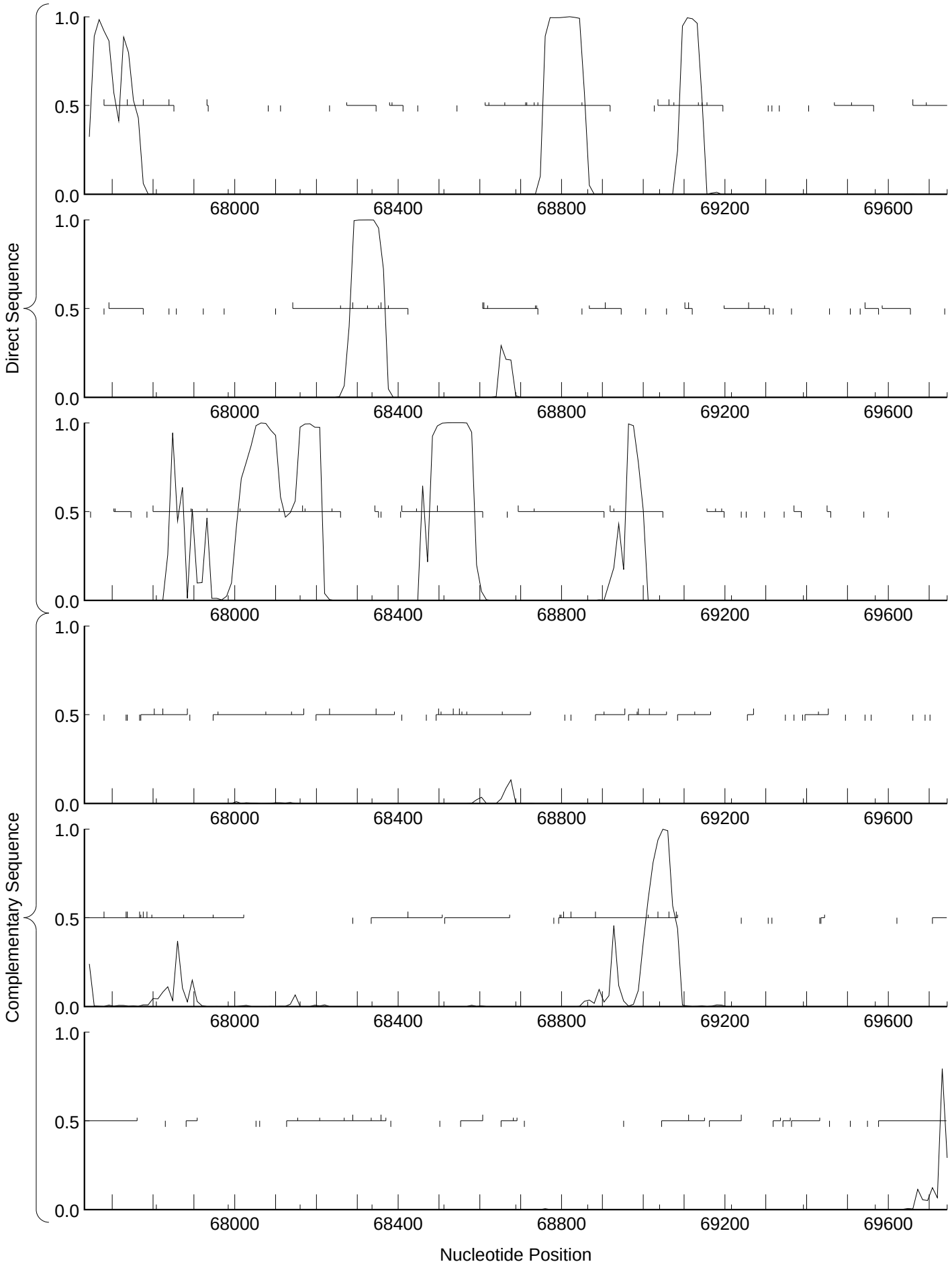
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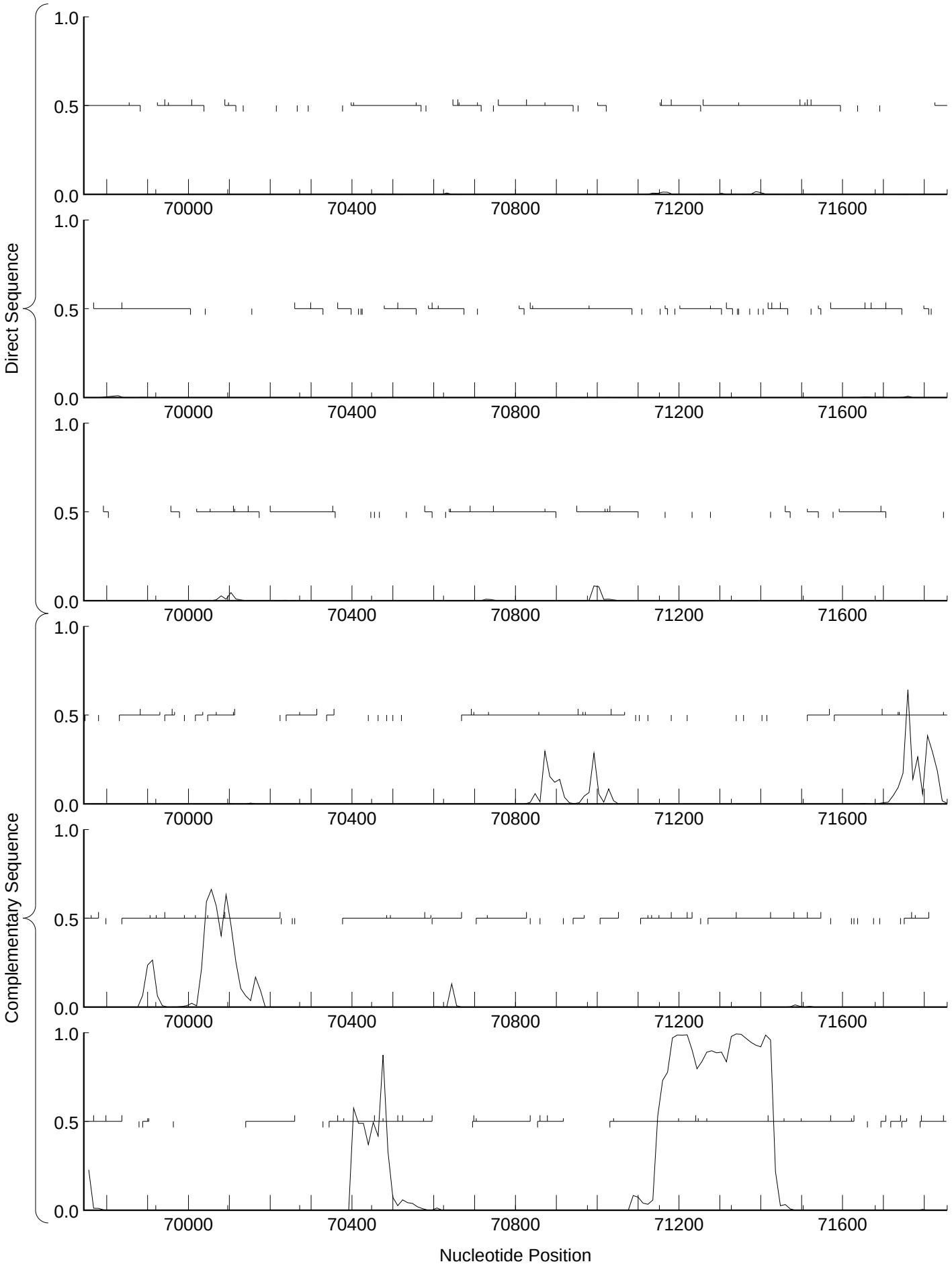
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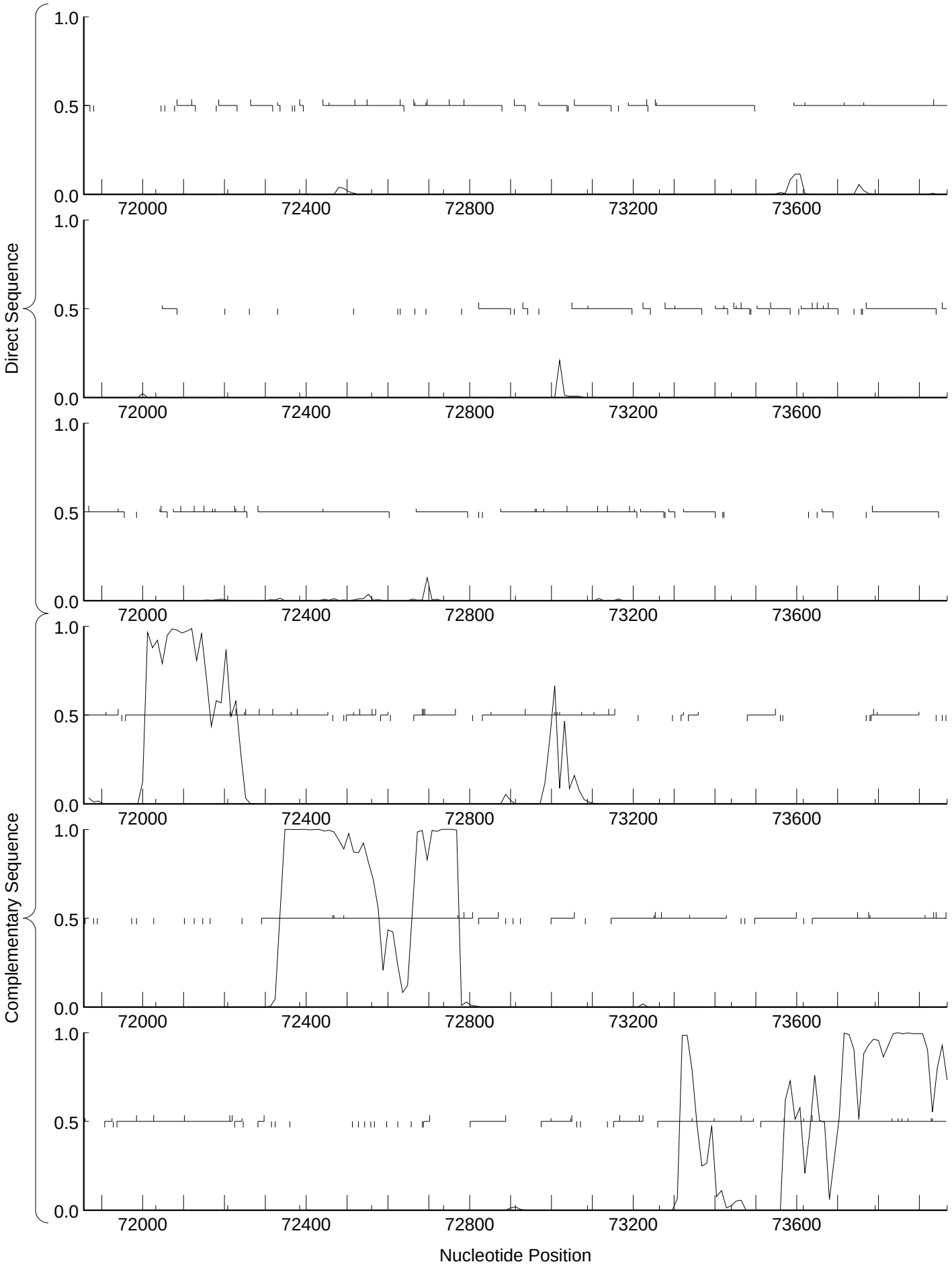
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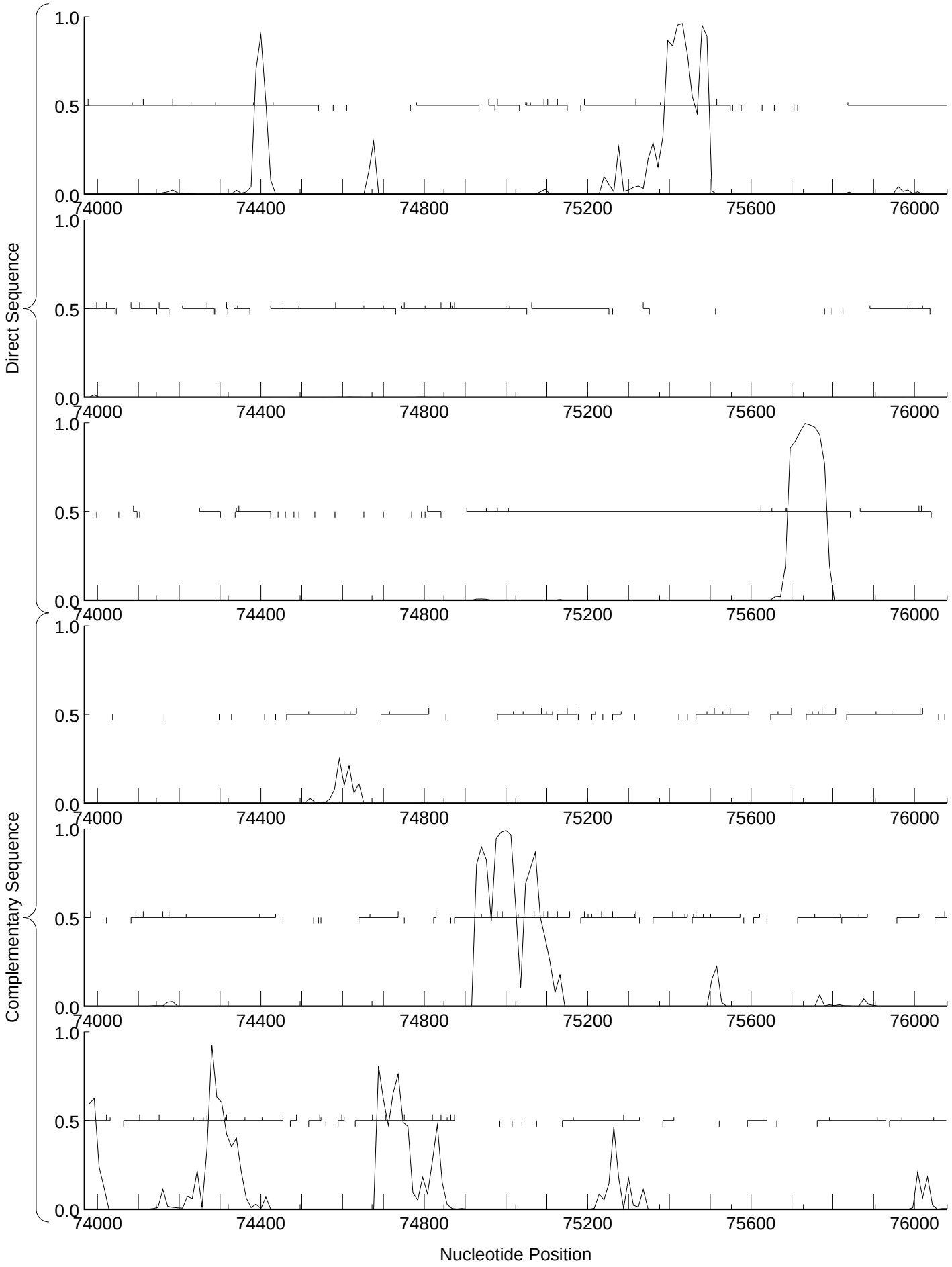
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yrale complete sequence, 76419 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 37/38



yrale complete sequence, 76419 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 38/38

