# GeneMark

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

## PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage WunderPhul complete sequence, 48724 bp including 10-base 3’ overhang (C
Analysis Date    :  3/28/24 at 18:23:12
Pages            :  25
Sequence Length  :  48724 bp
GC Content       :   61.54%

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

PS-Version       :  1.2

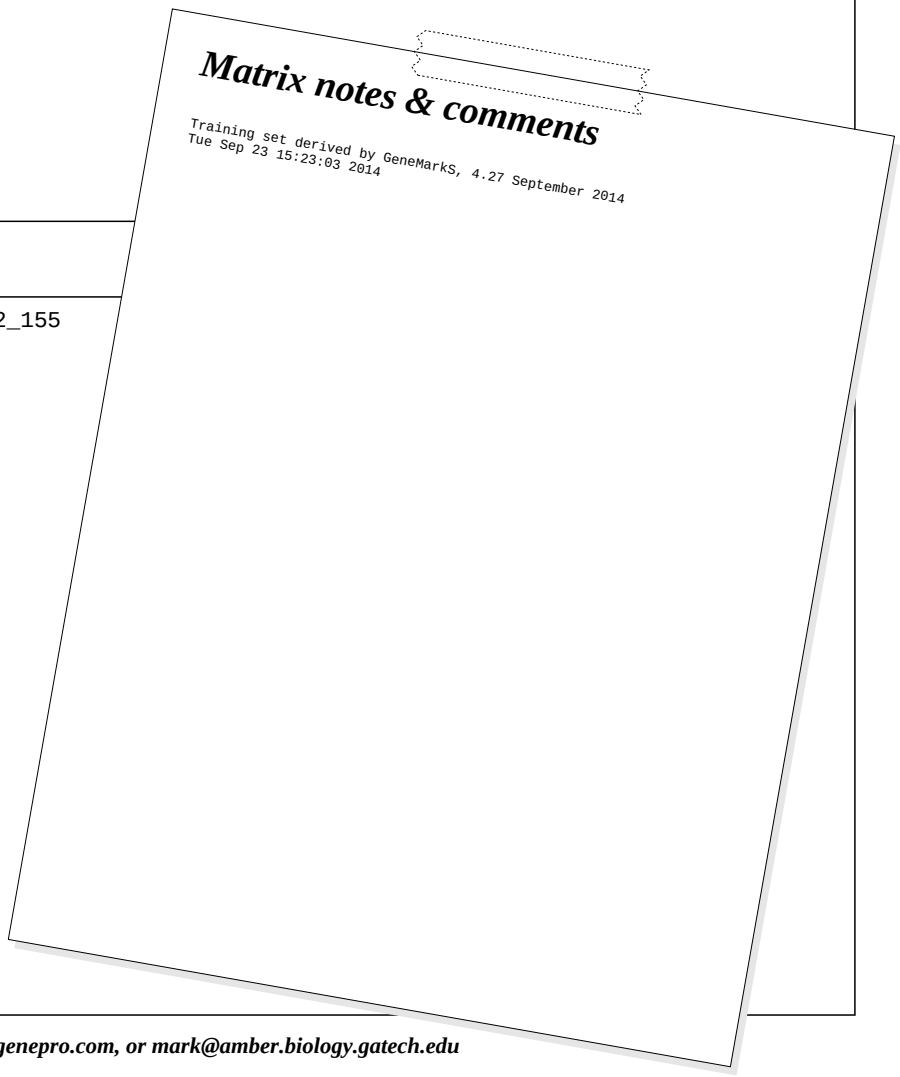
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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## MATRIX INFORMATION

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Author     :  -

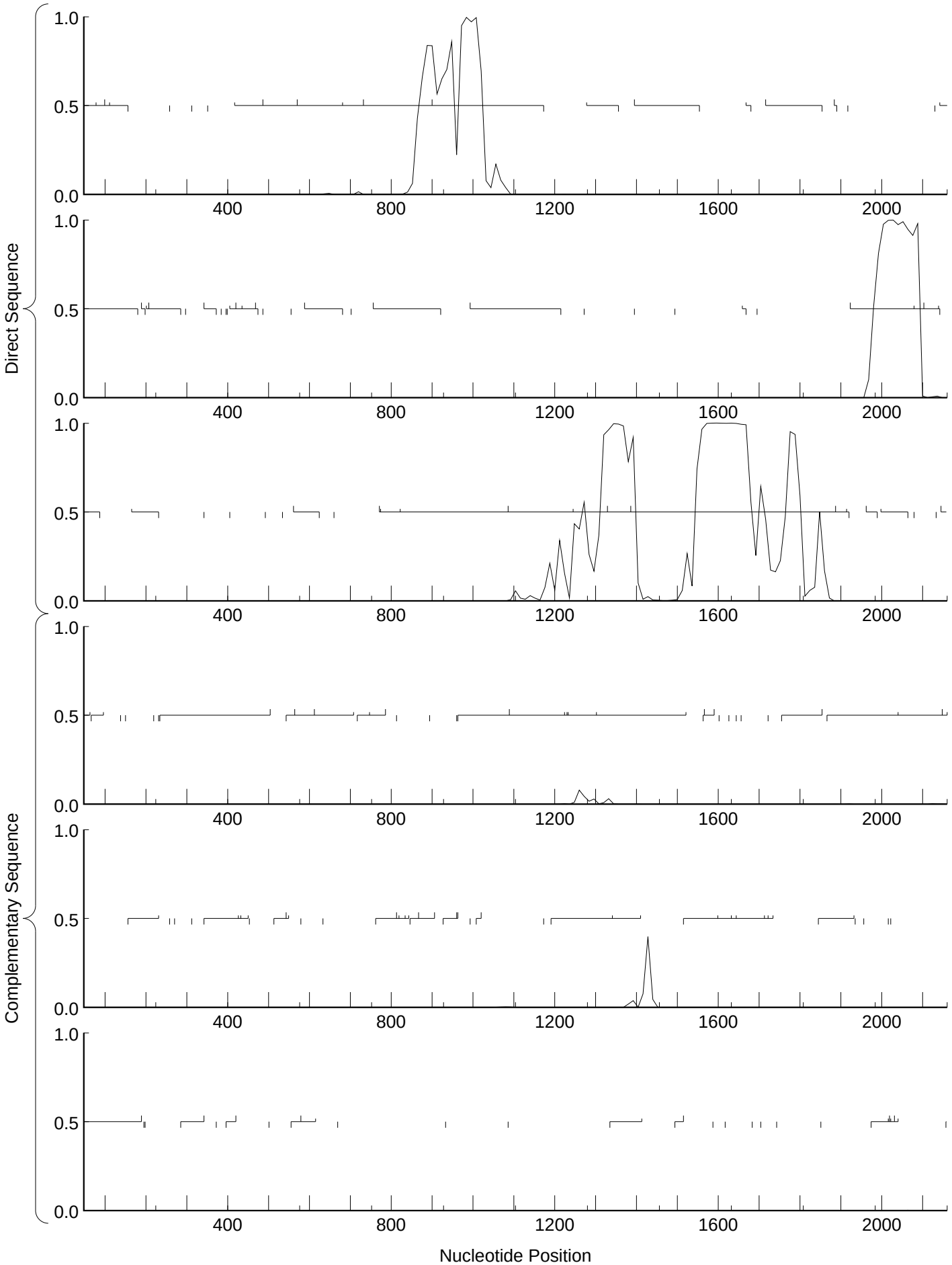
Order      :  4


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

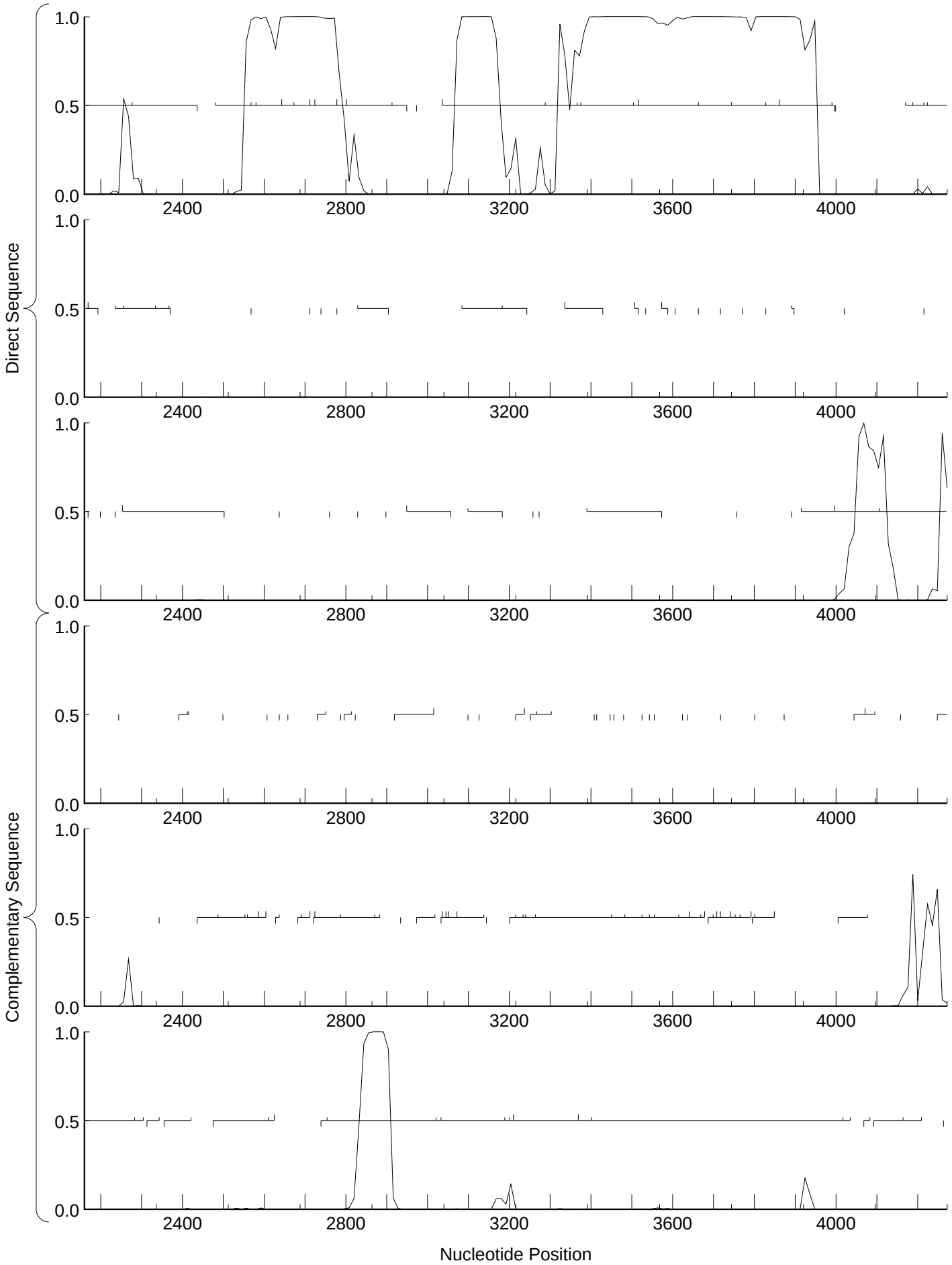



*For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu*

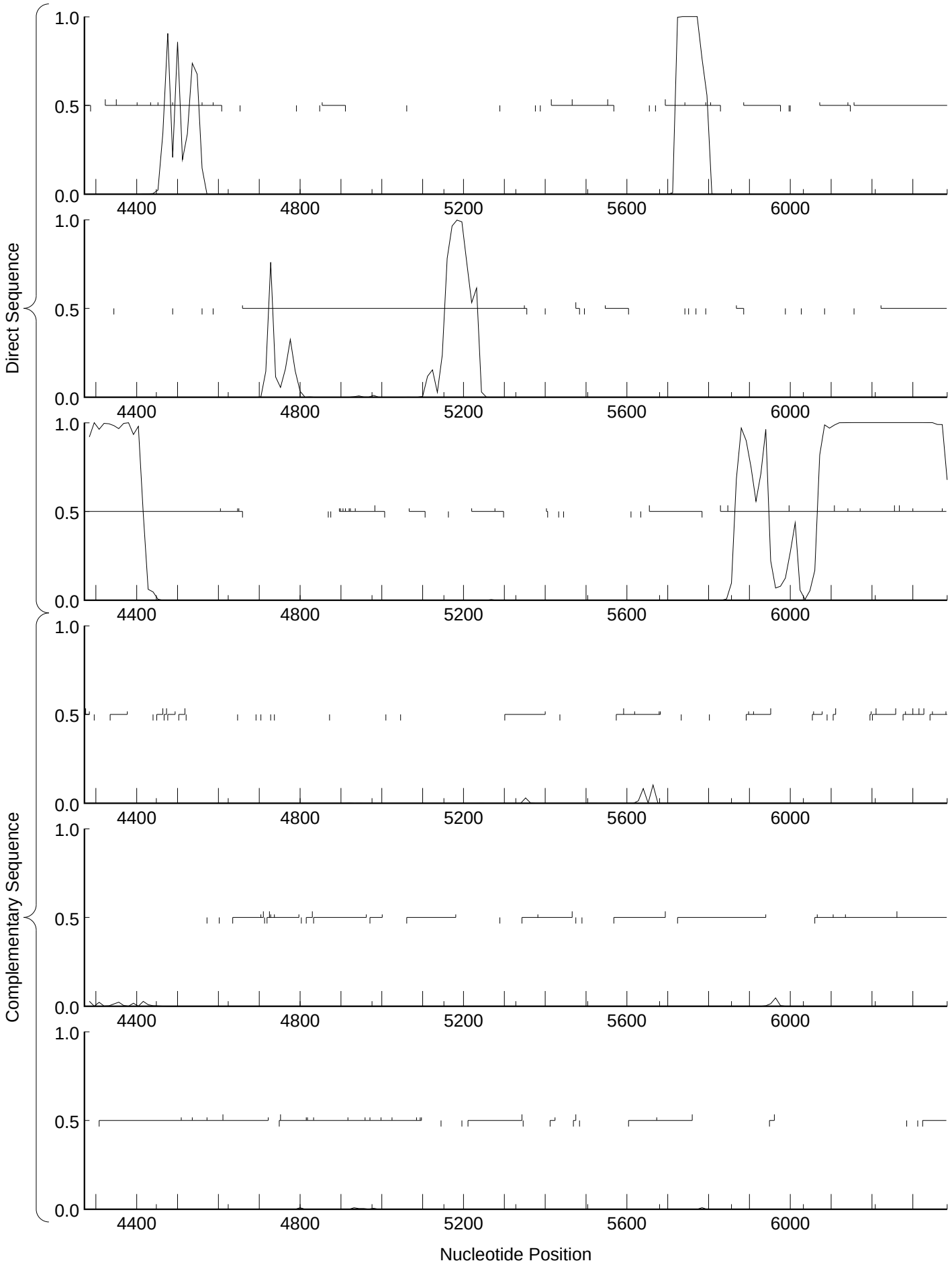
ul complete sequence, 48724 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 2/25



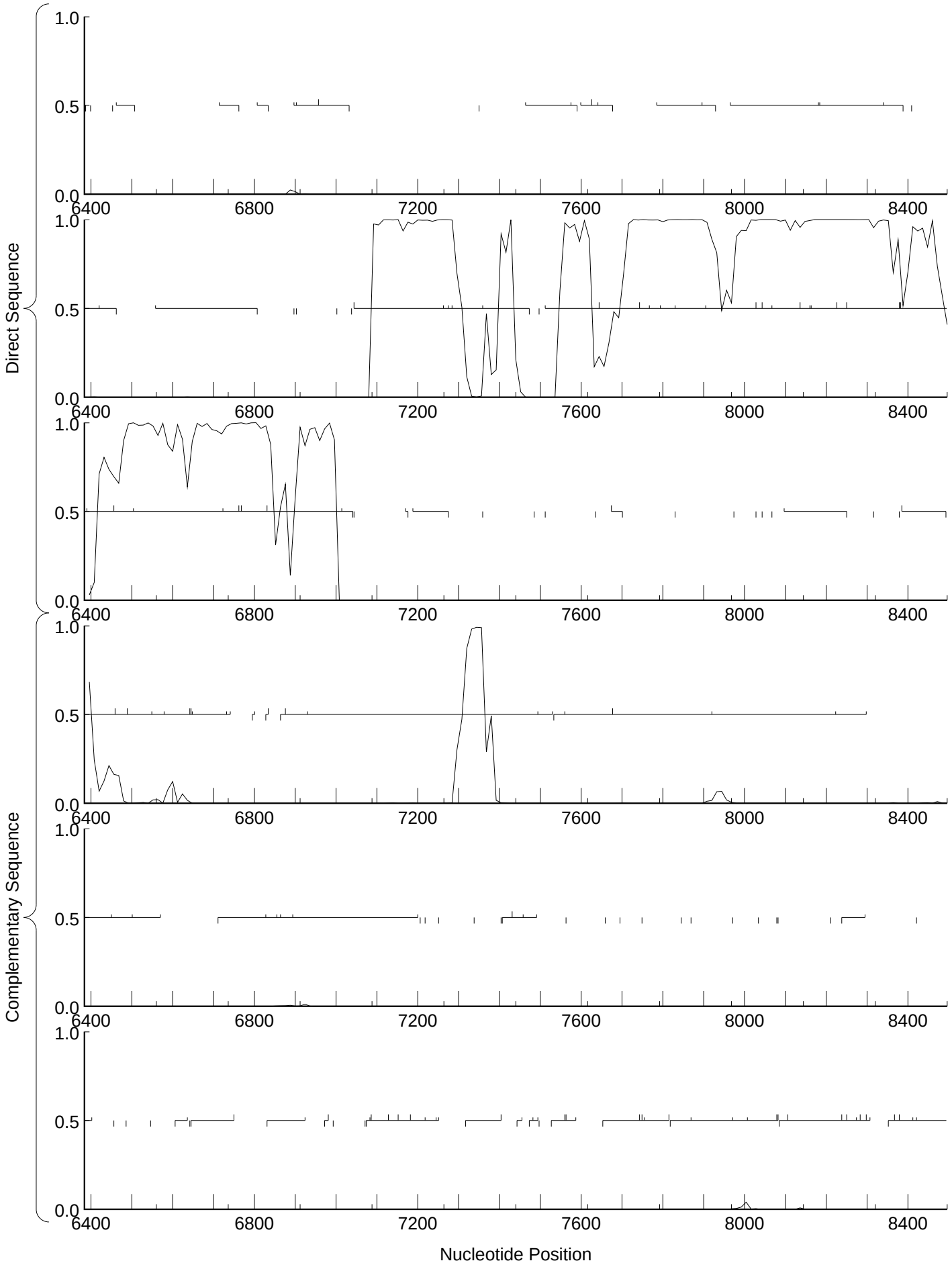
ul complete sequence, 48724 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 3/25



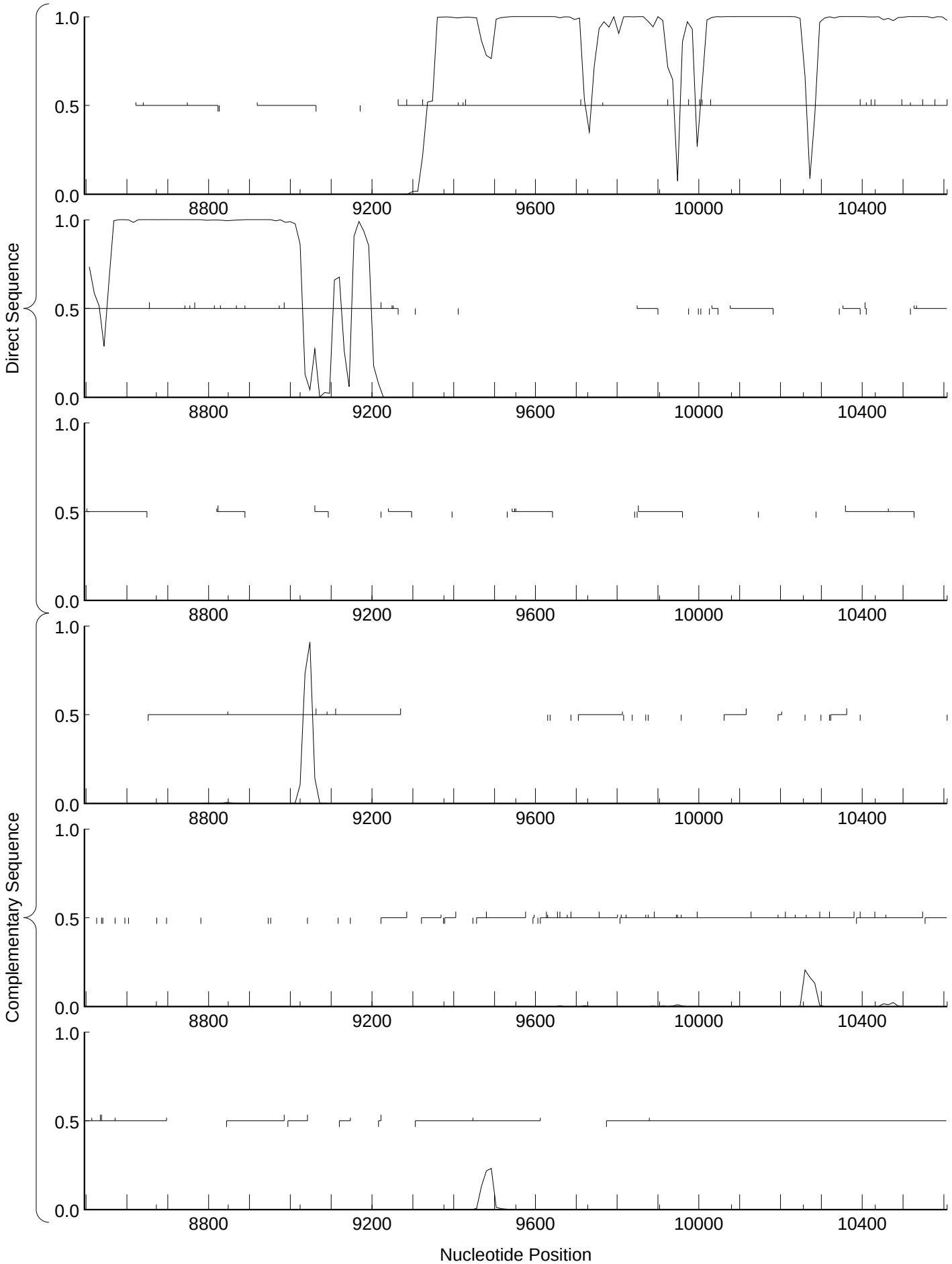
ul complete sequence, 48724 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 4/25



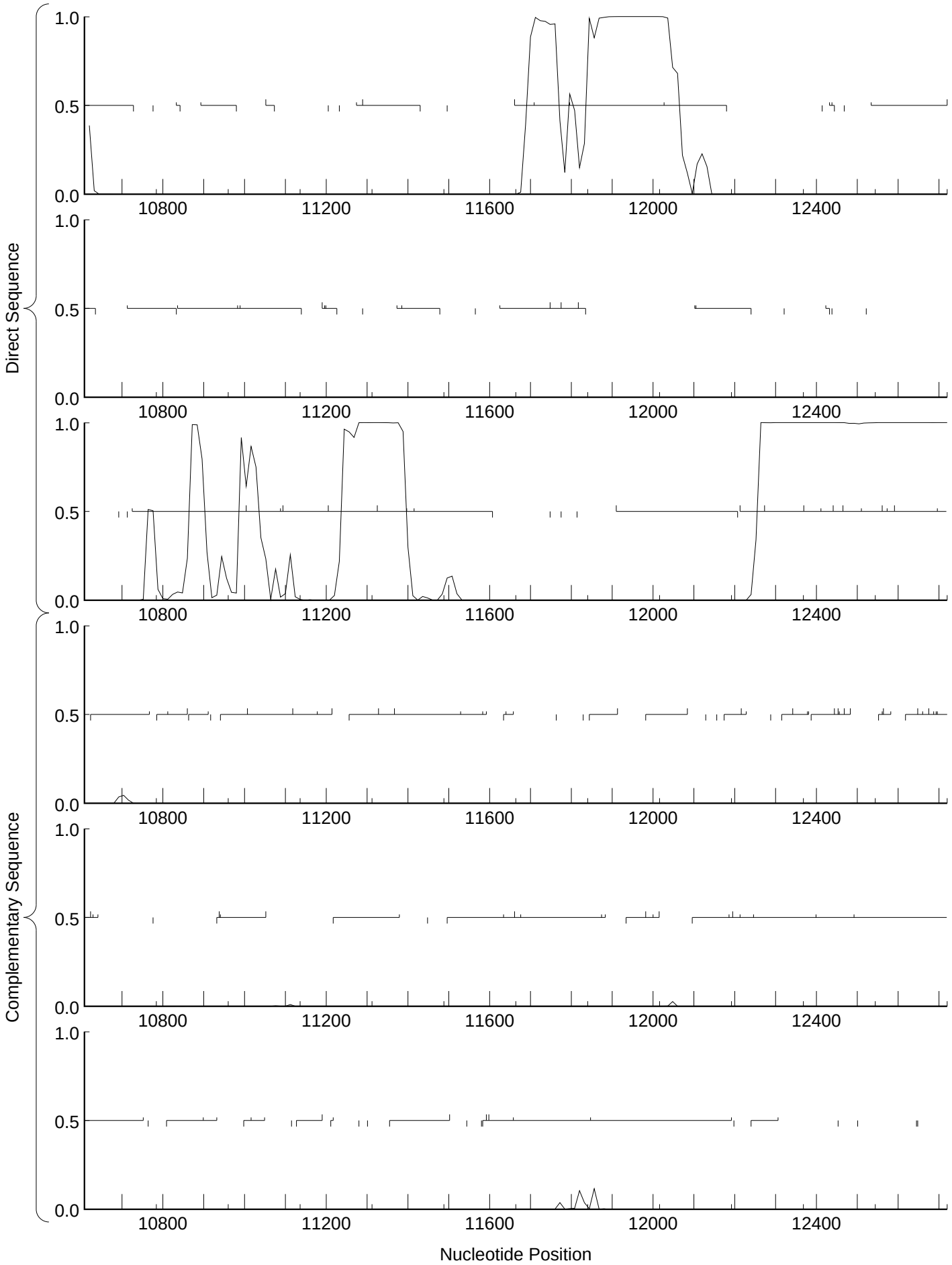
ul complete sequence, 48724 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 5/25



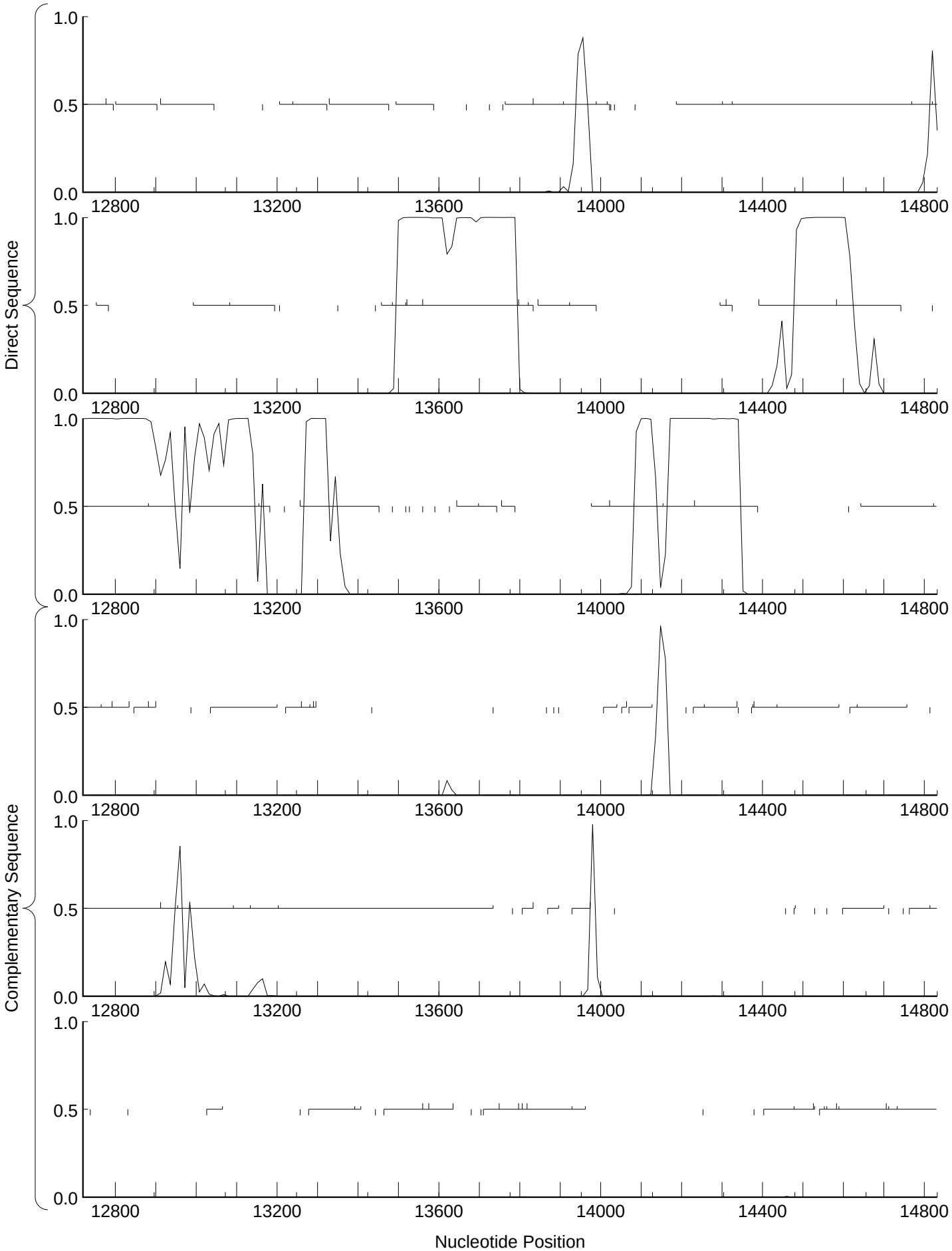
ul complete sequence, 48724 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 6/25



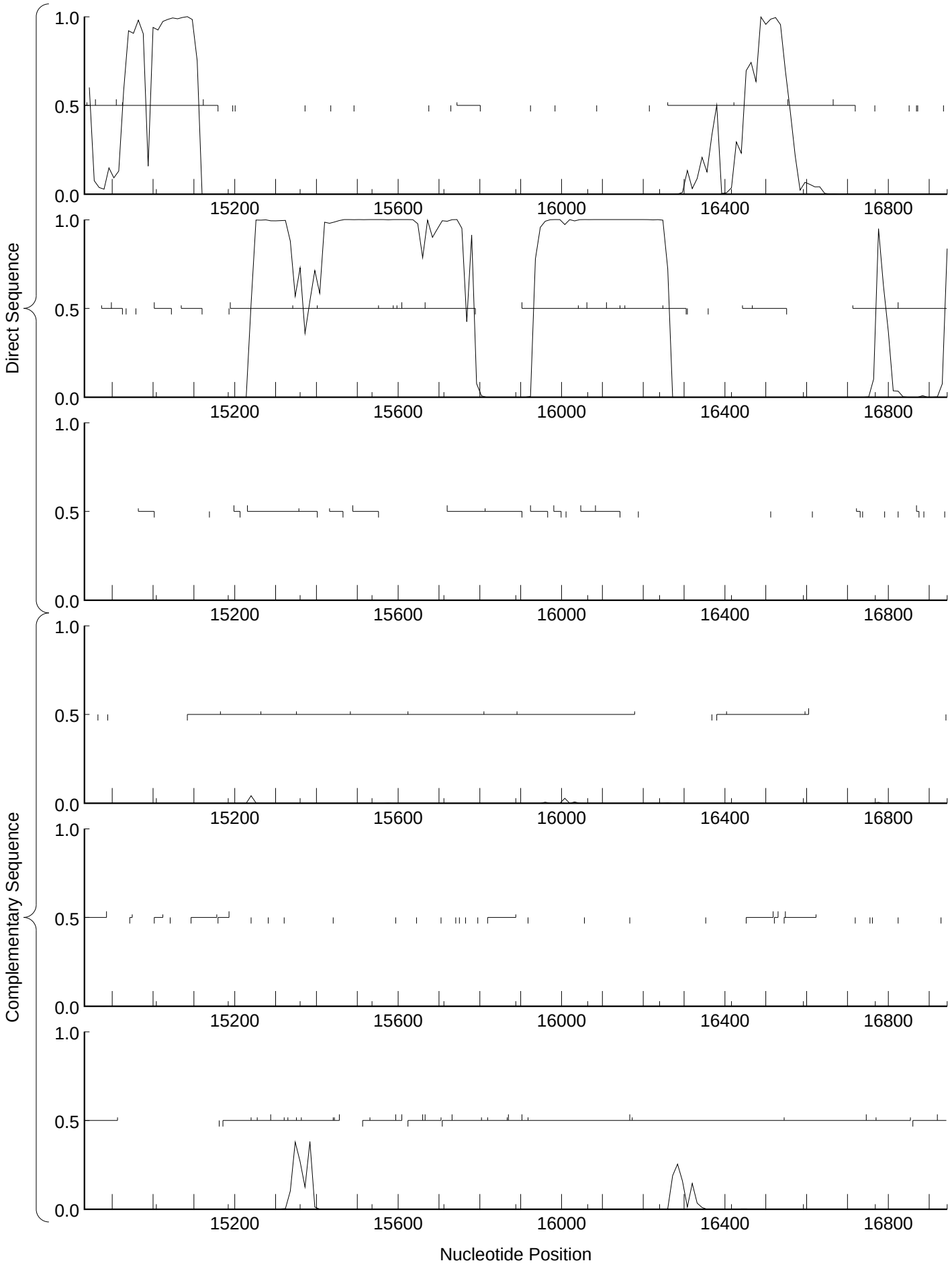
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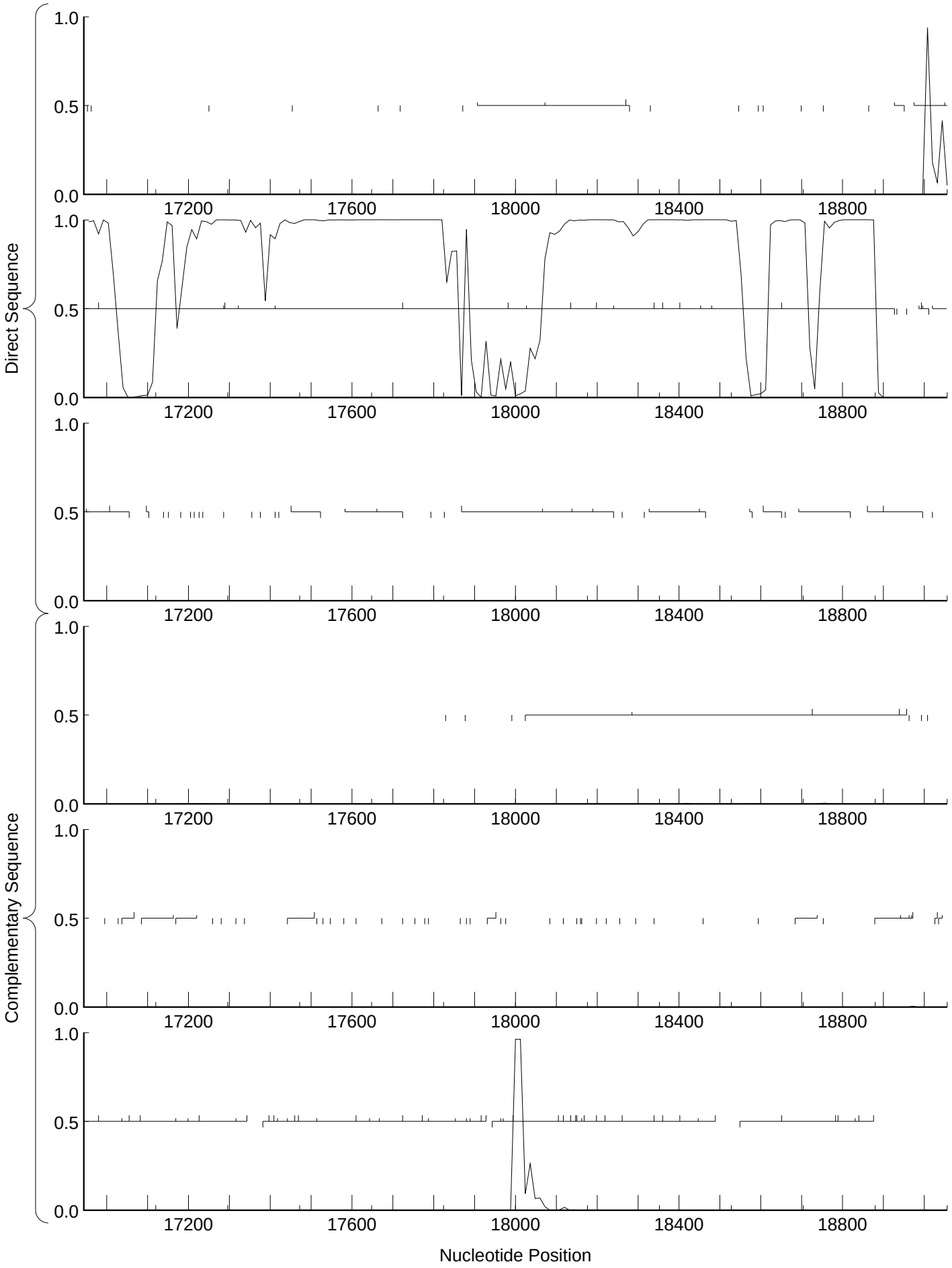
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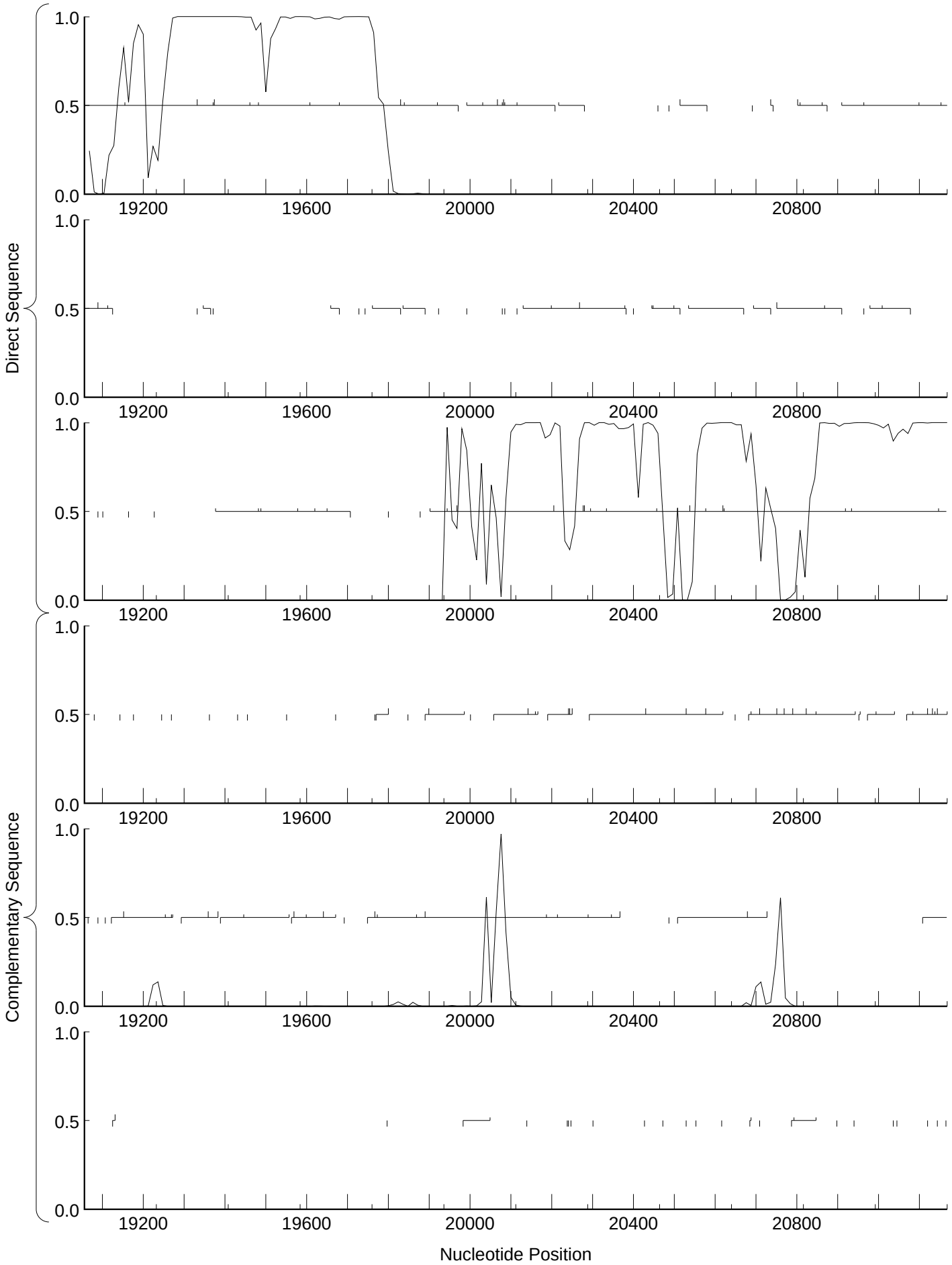
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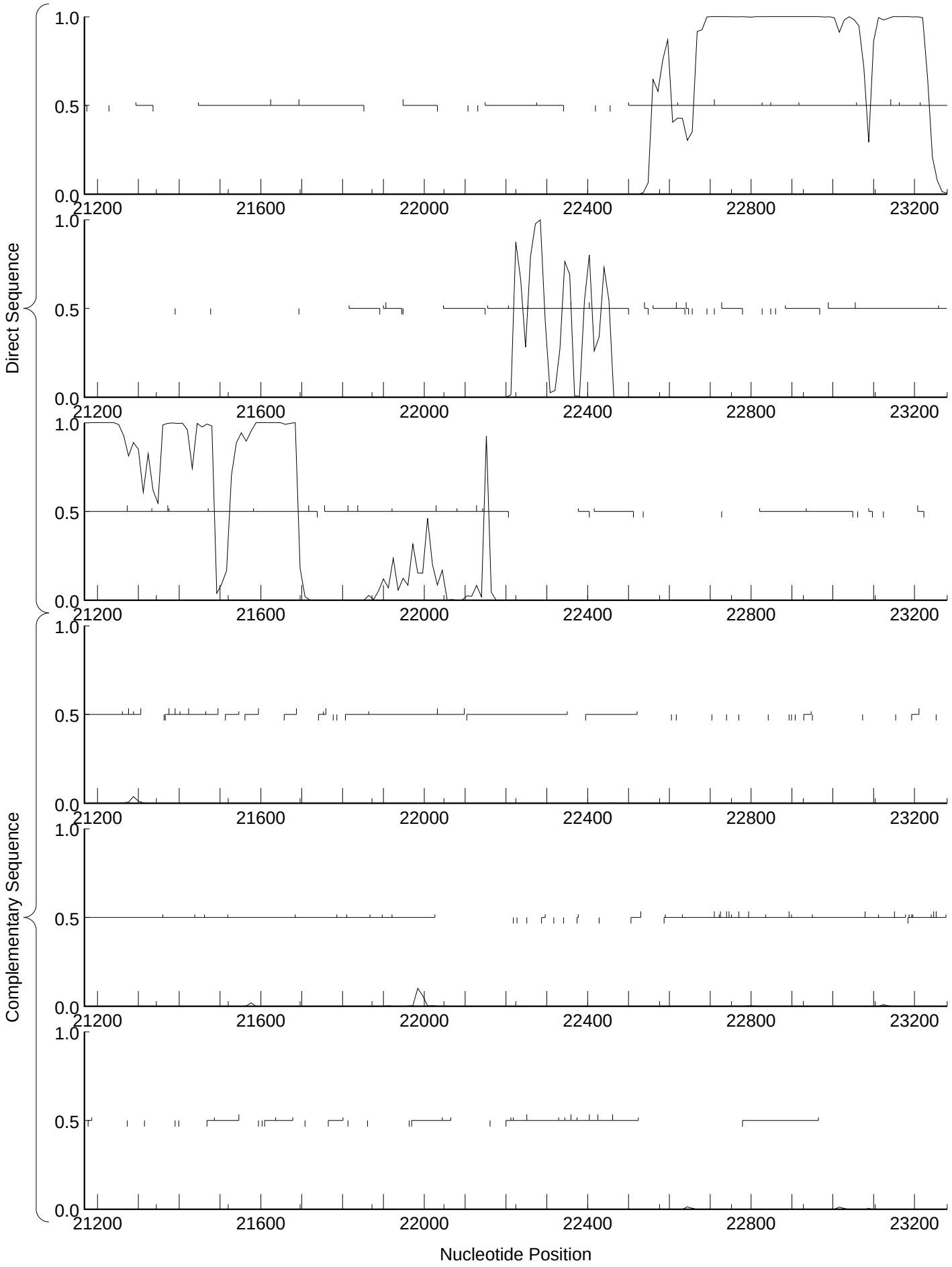
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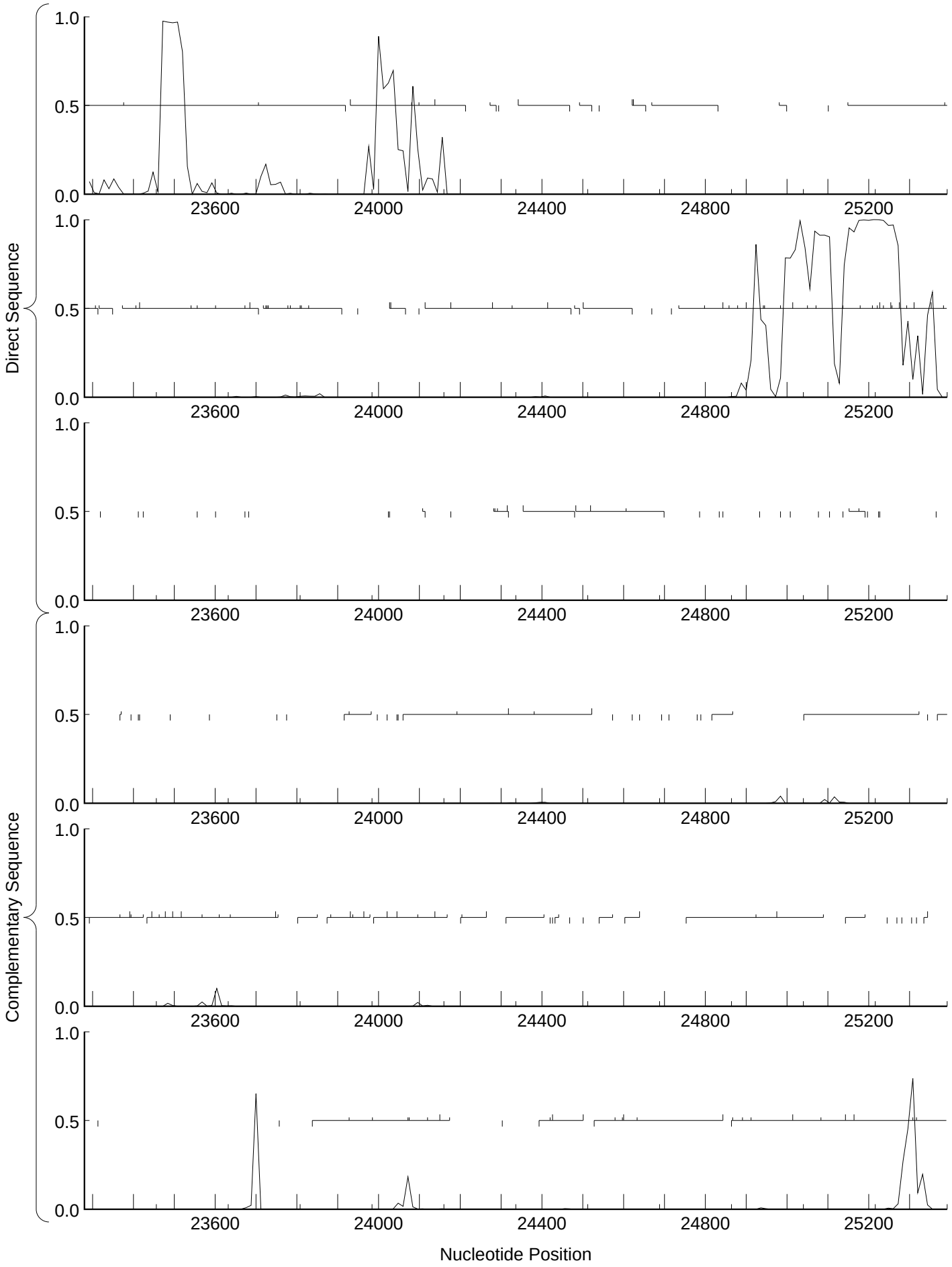
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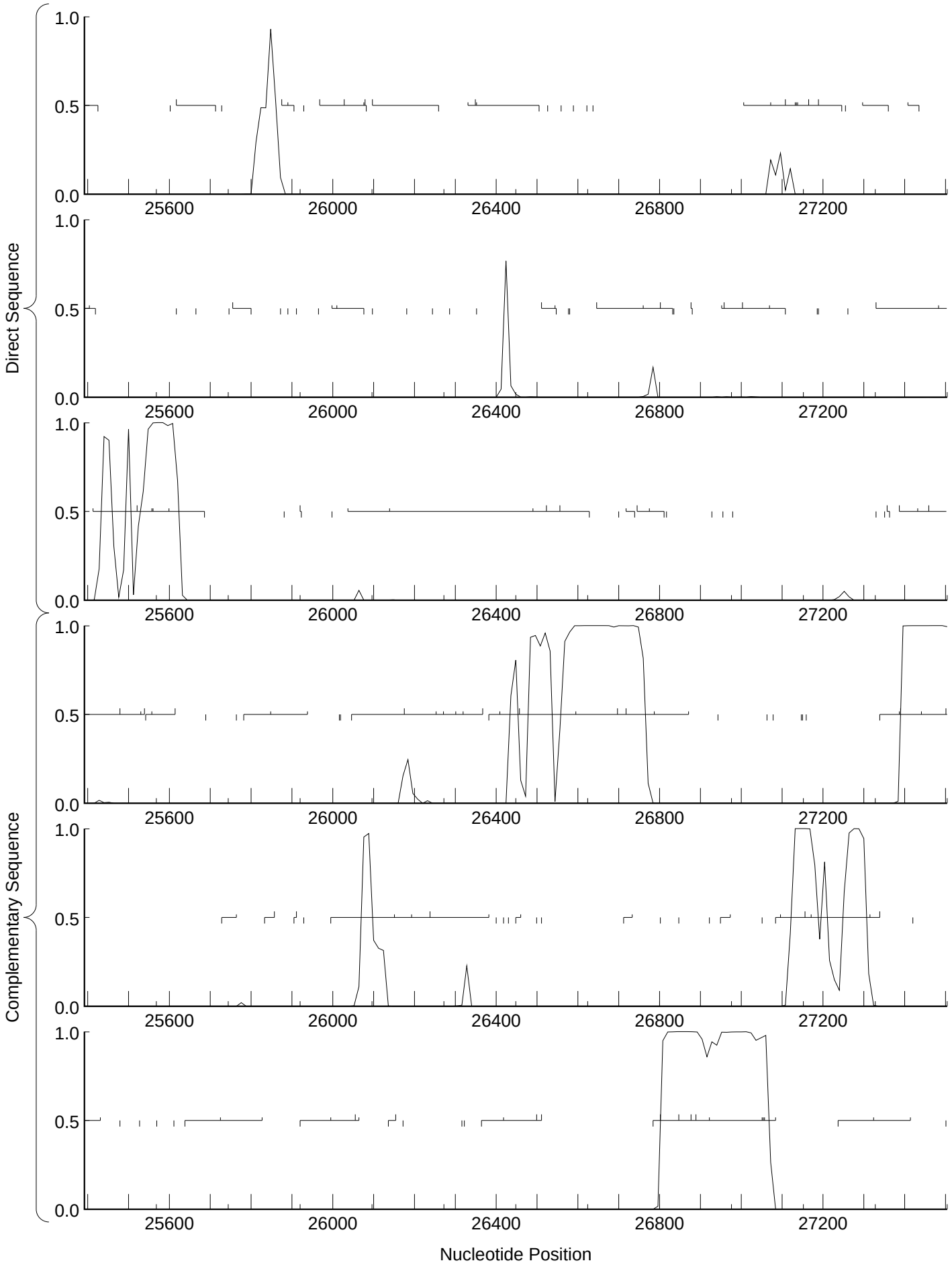
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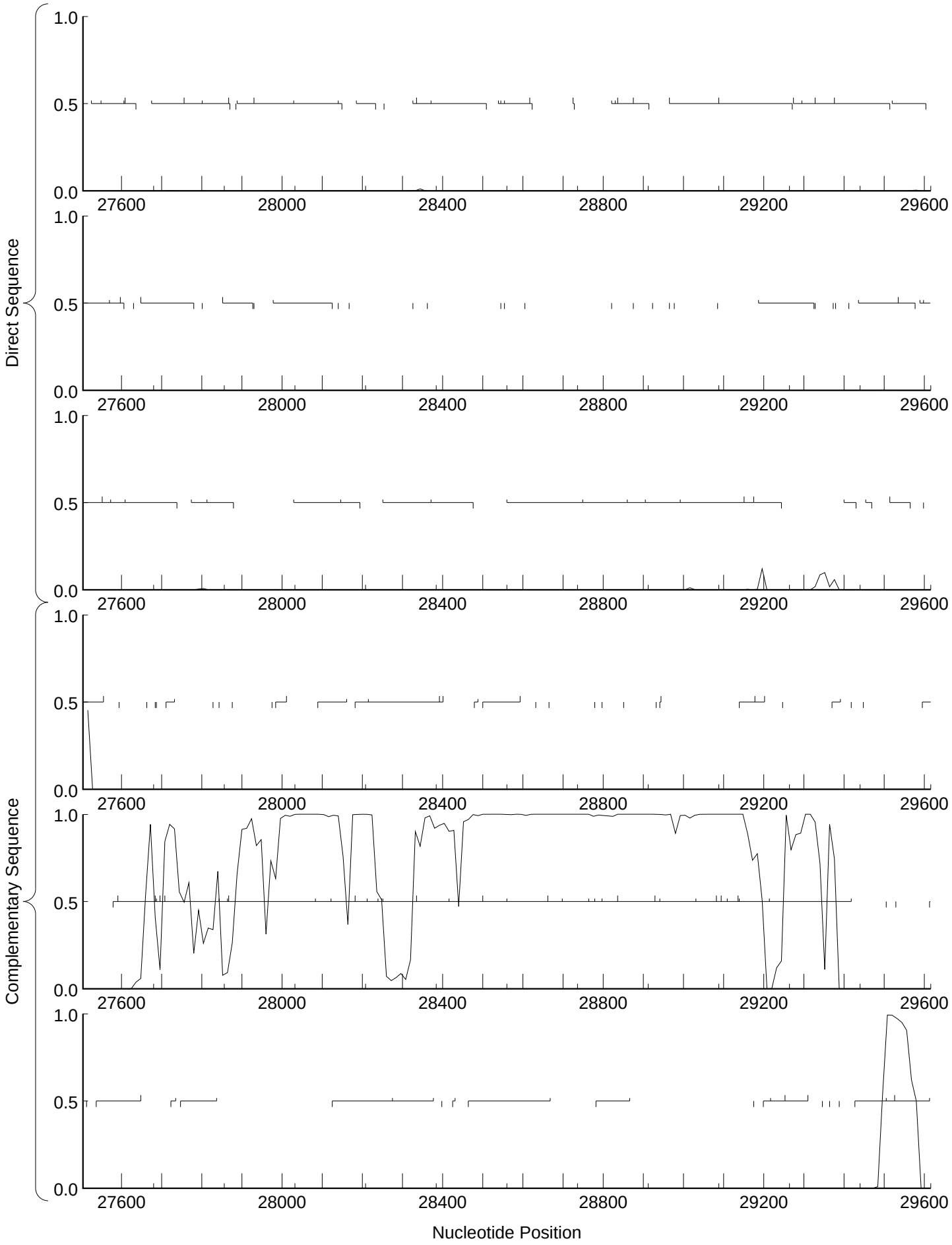
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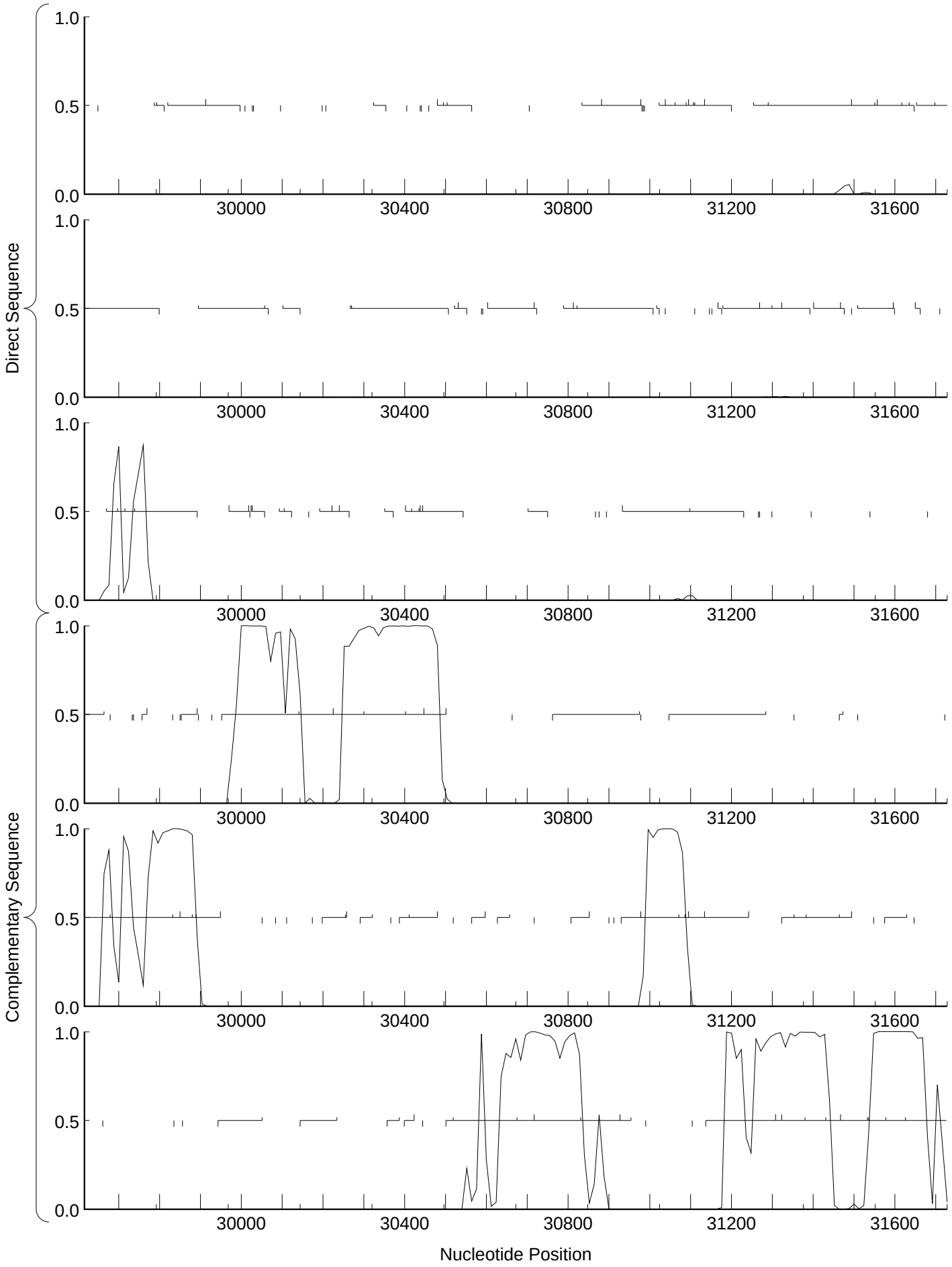
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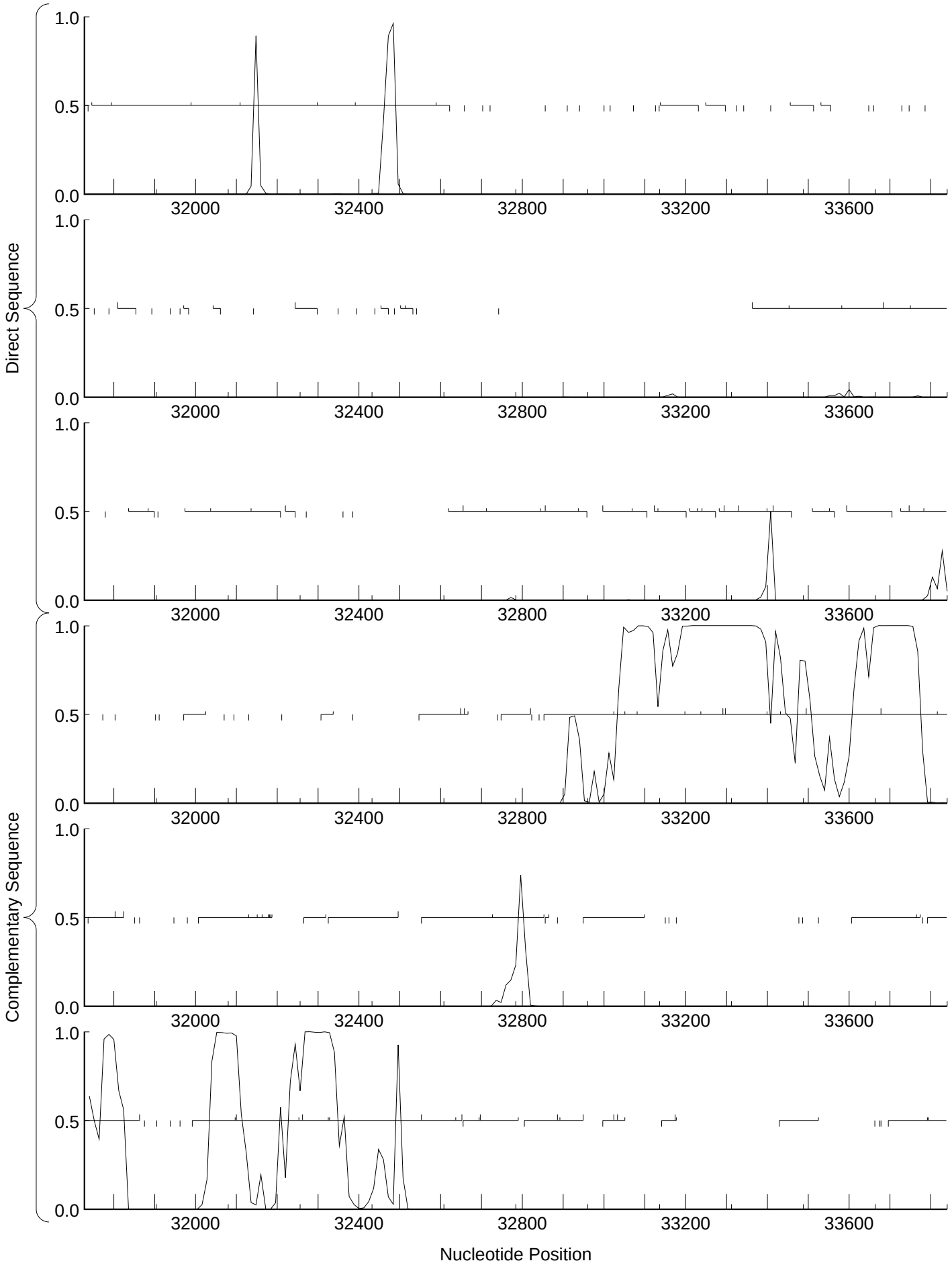
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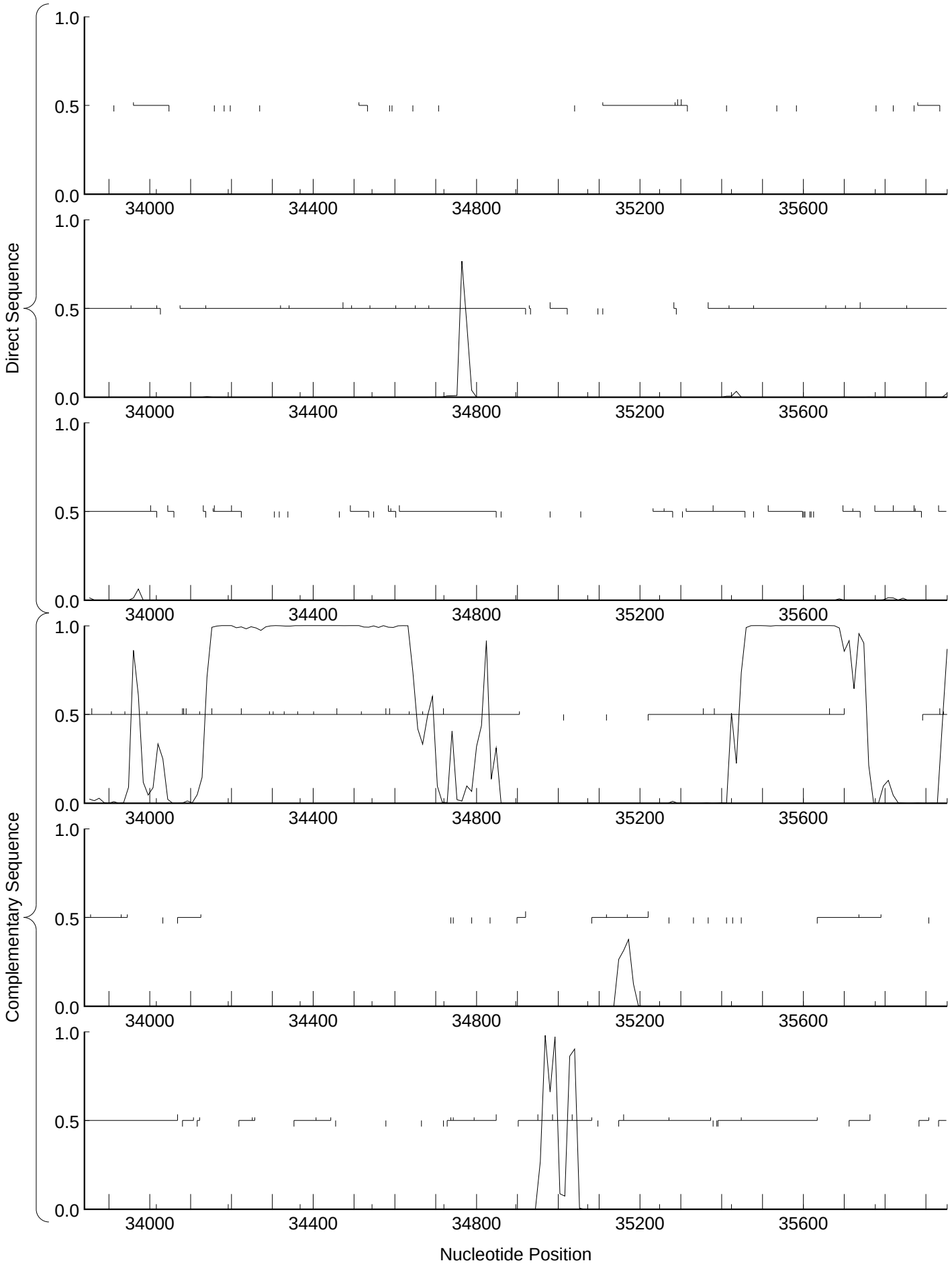
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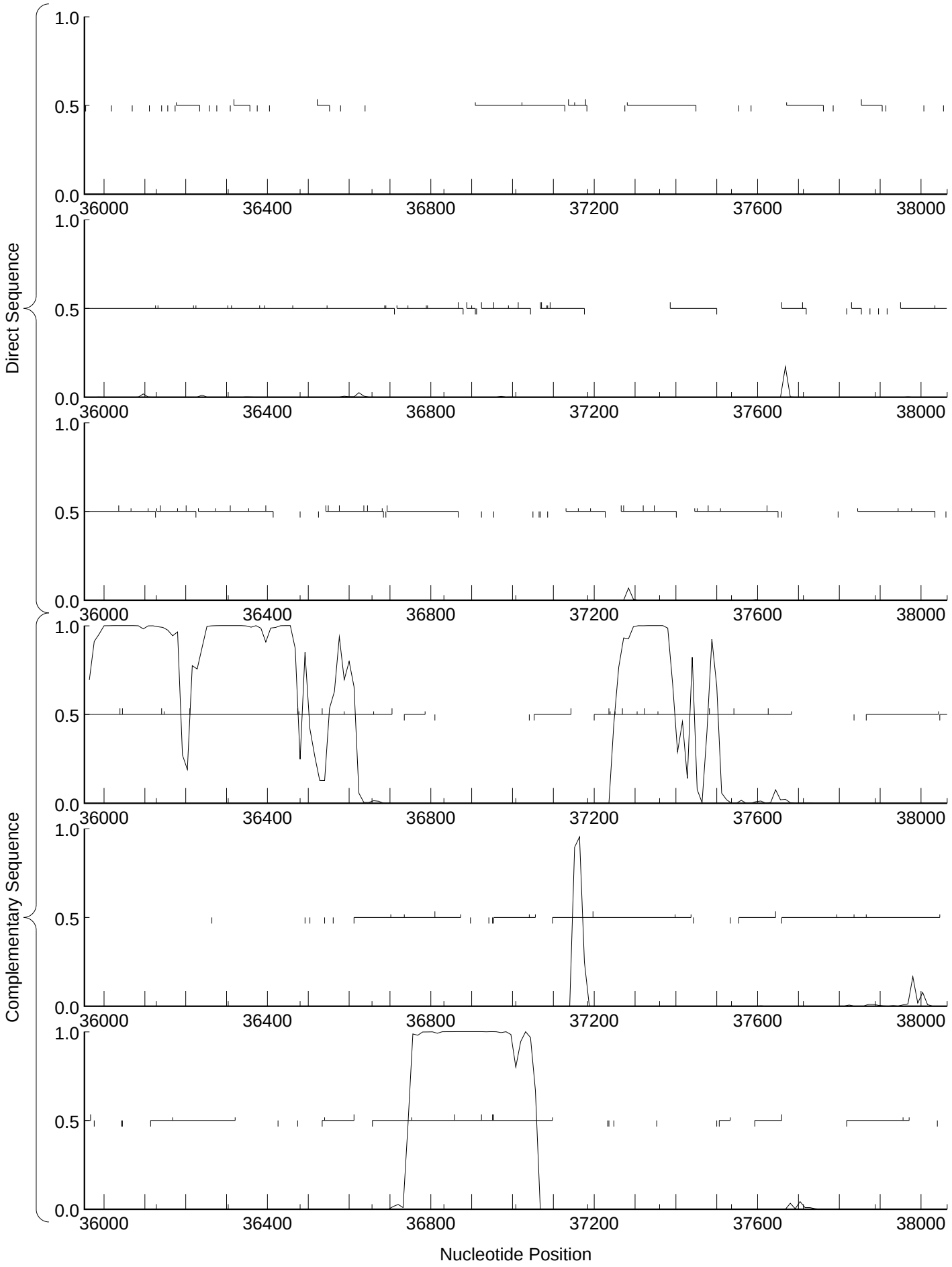
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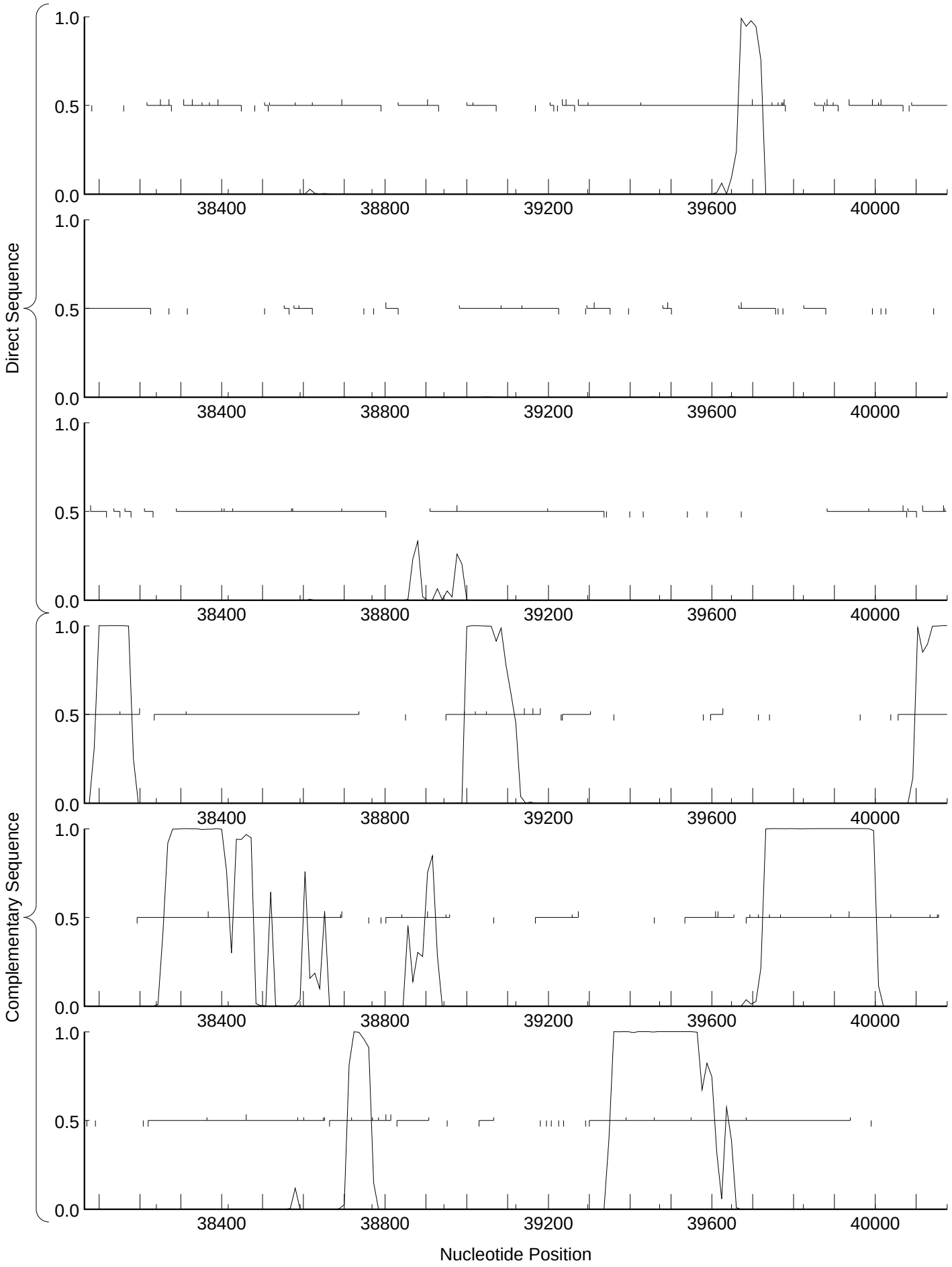
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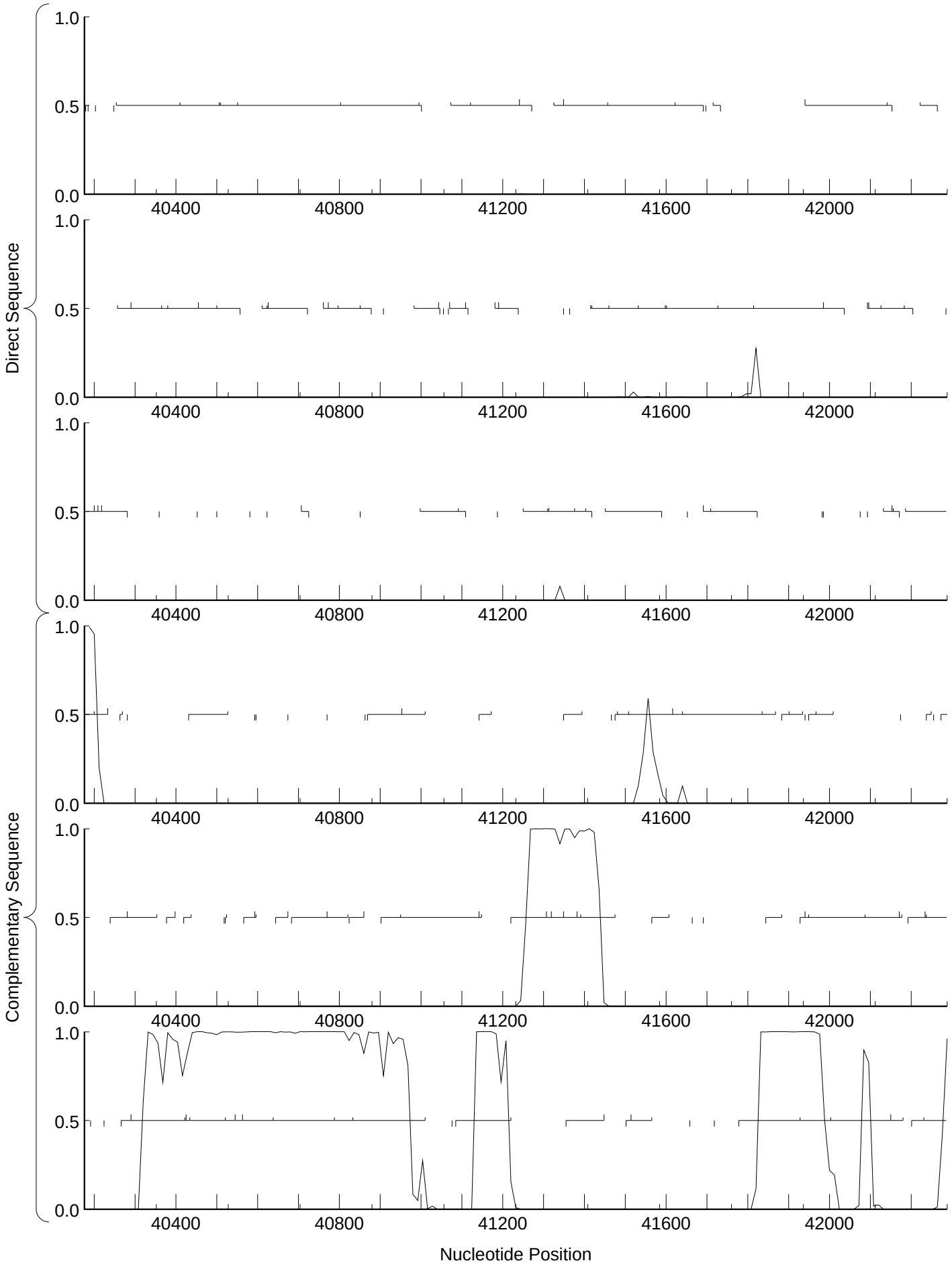
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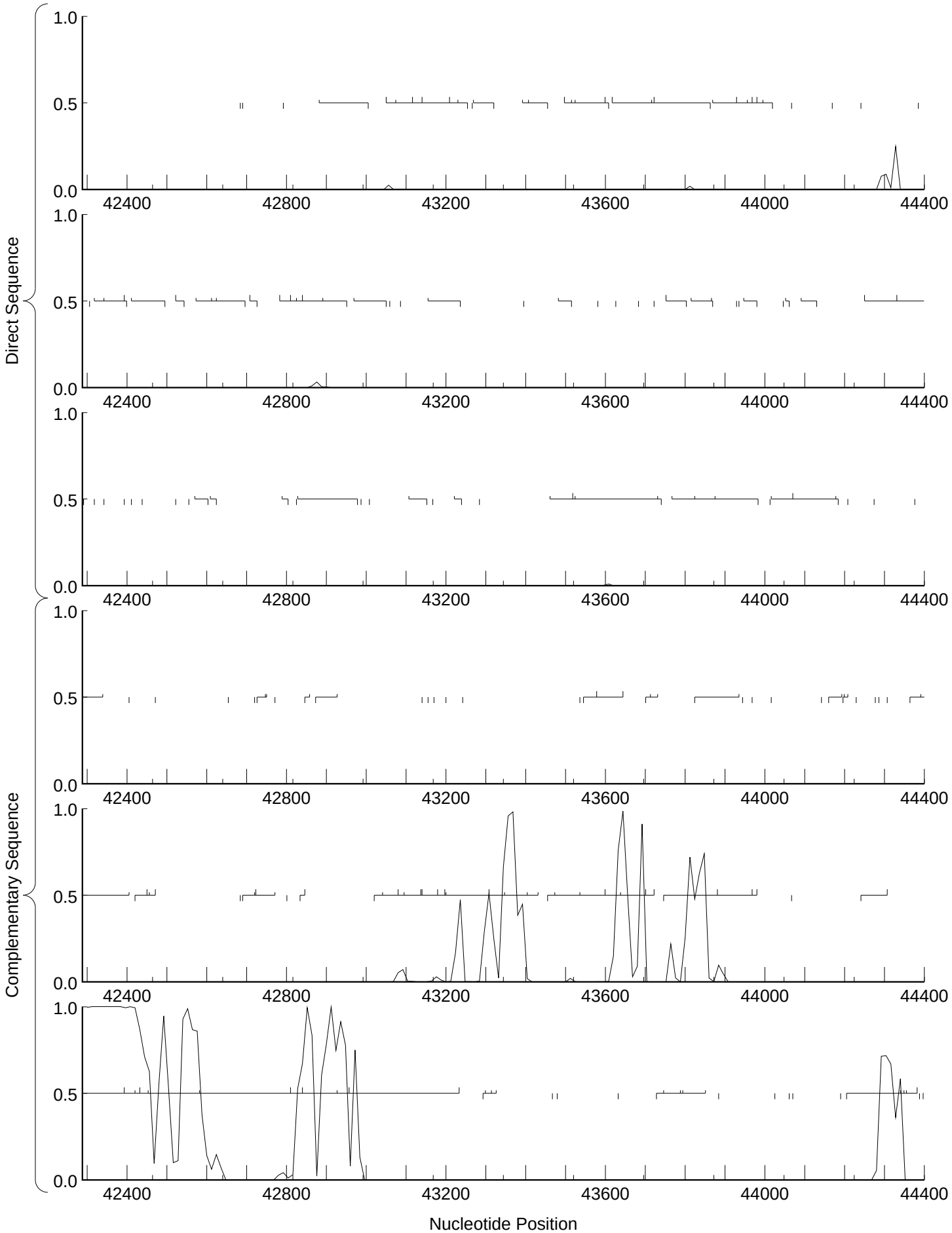
complete sequence, 48724 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 20/25



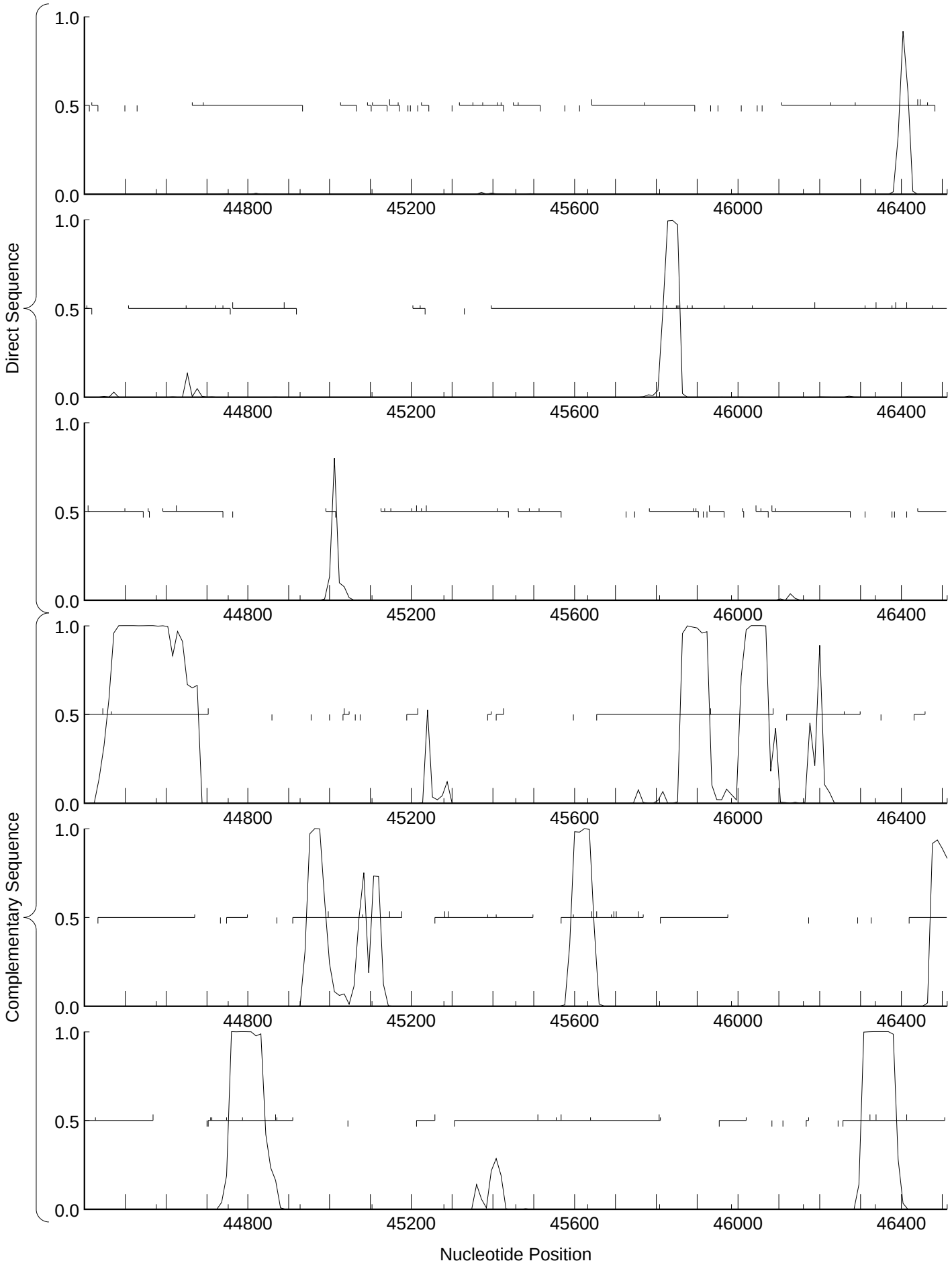
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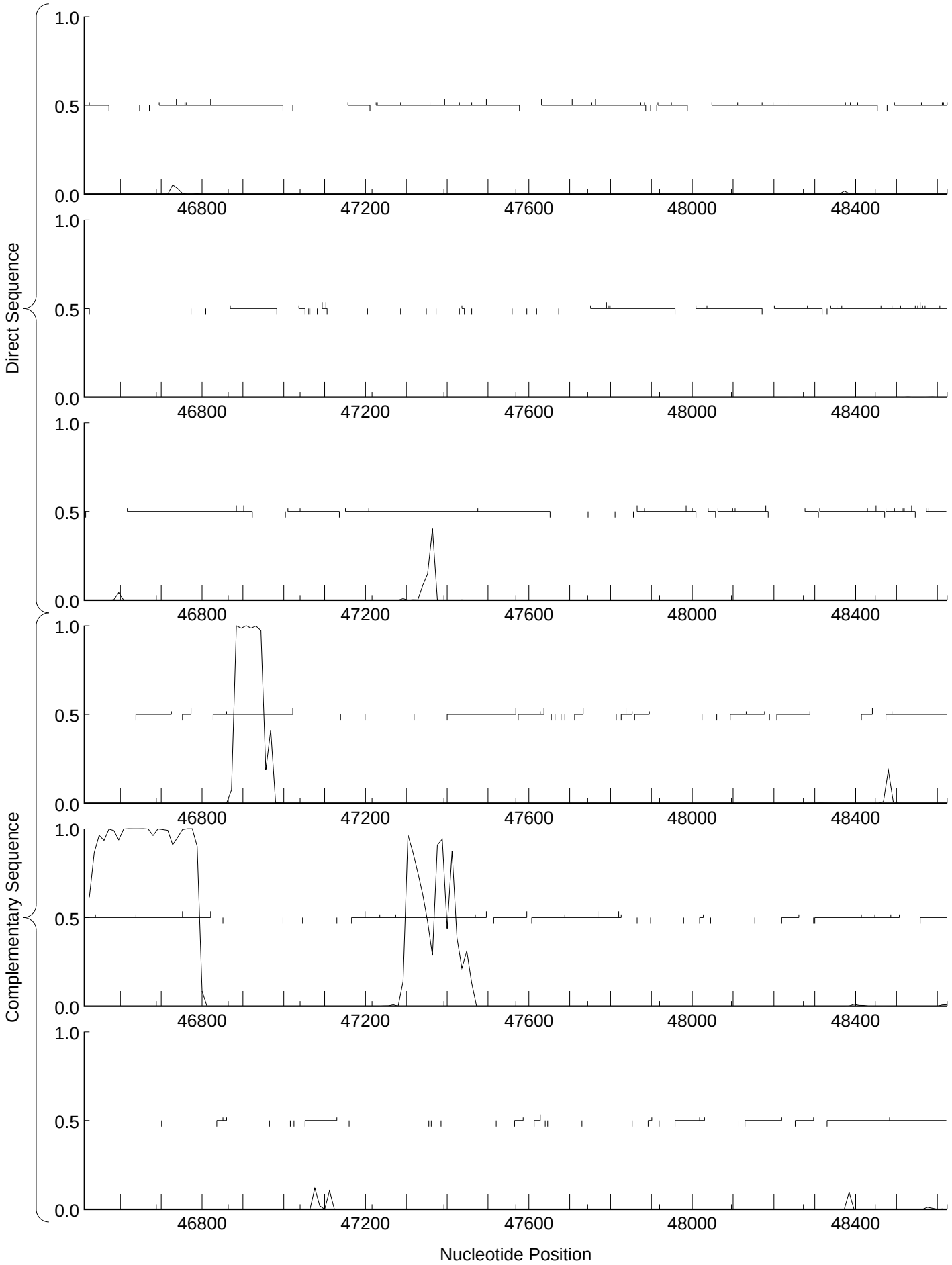
complete sequence, 48724 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 22/25



complete sequence, 48724 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 23/25



complete sequence, 48724 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 24/25



complete sequence, 48724 bp including 10-base 3’ overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 25/25

