GeneMark

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

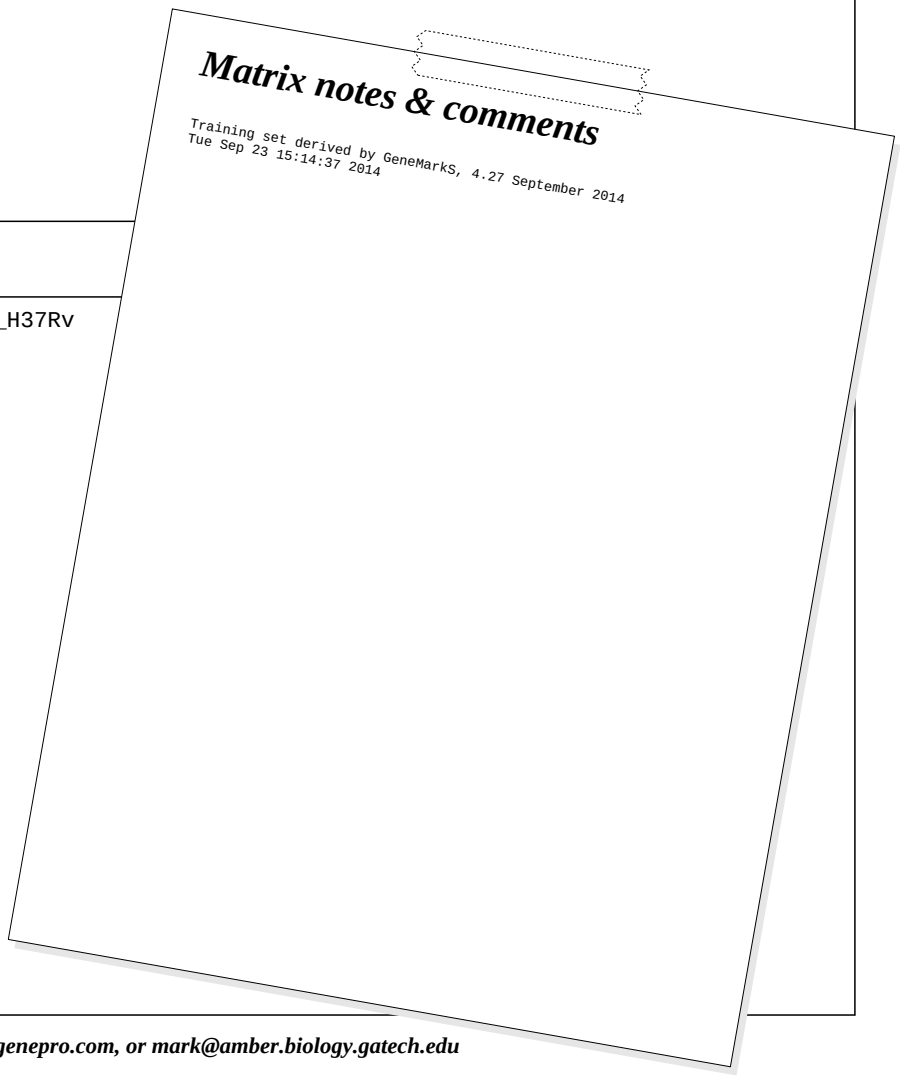
PROGRAM INFORMATION

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Sequence         :  Mycobacterium phage Phaedrus complete genome
Analysis Date    :  3/29/24 at 1:42:28
Pages            :  34
Sequence Length  :  68090 bp
GC Content       :   67.57%

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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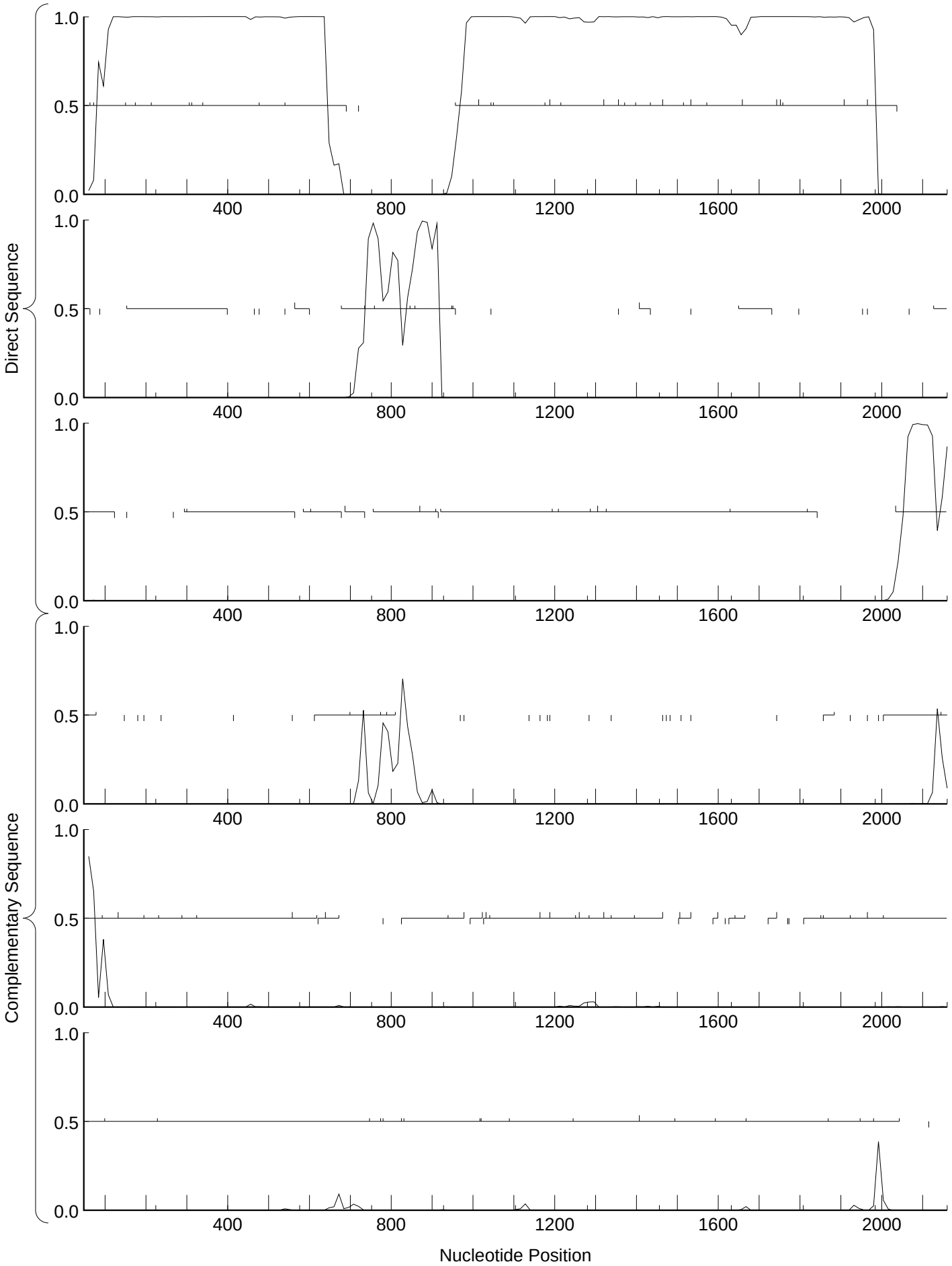
MATRIX INFORMATION

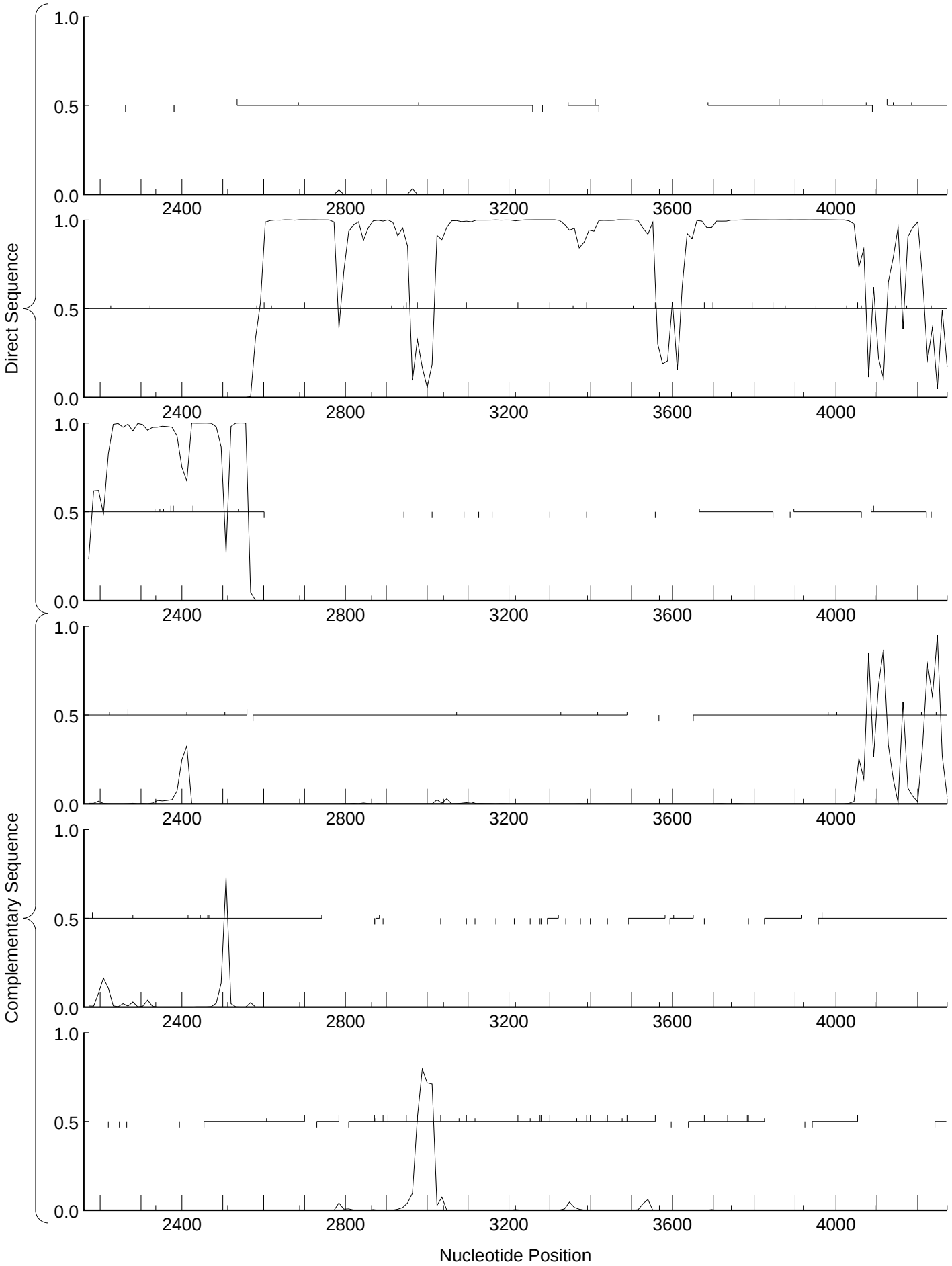
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Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

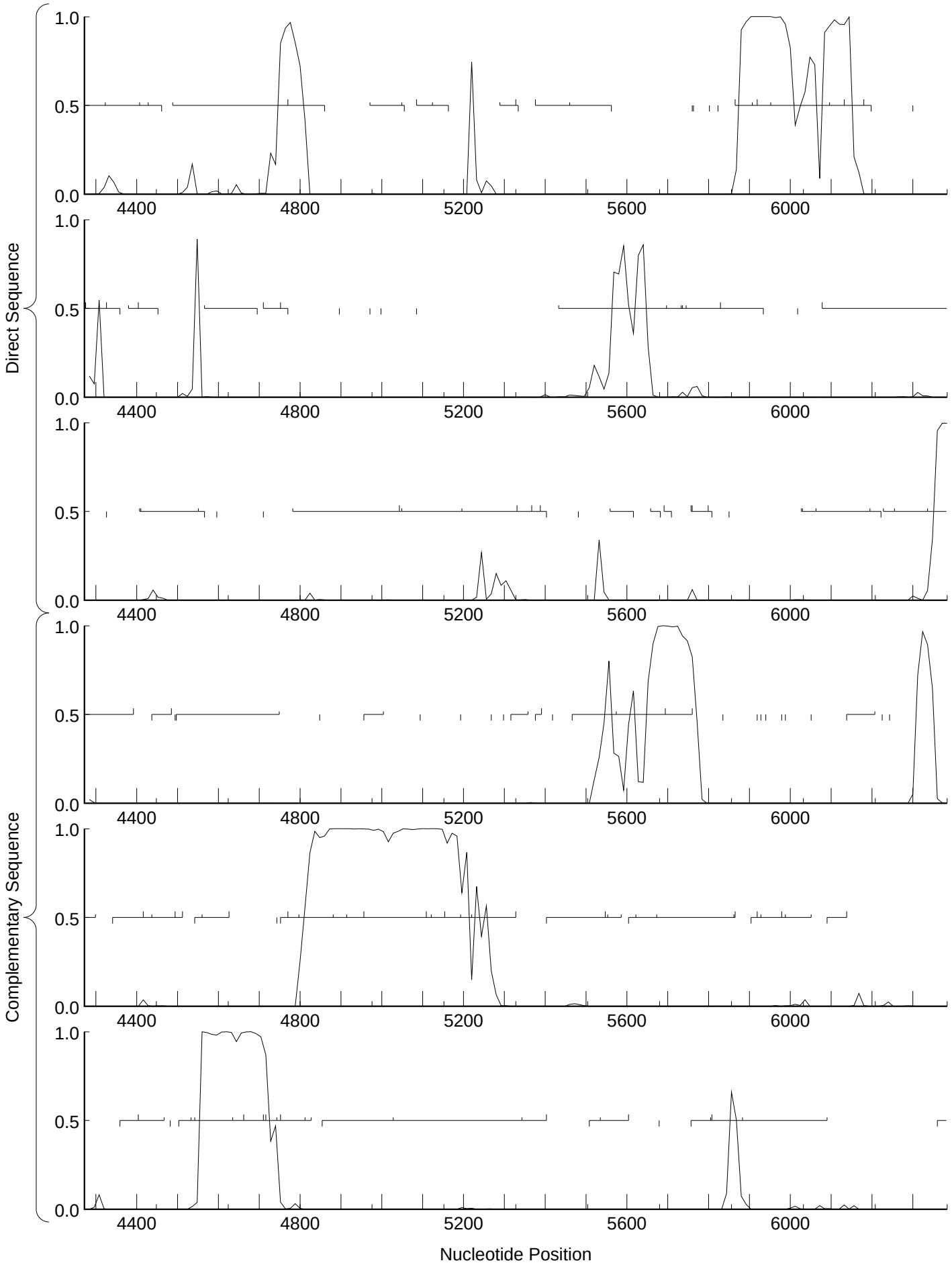
Order      :  4
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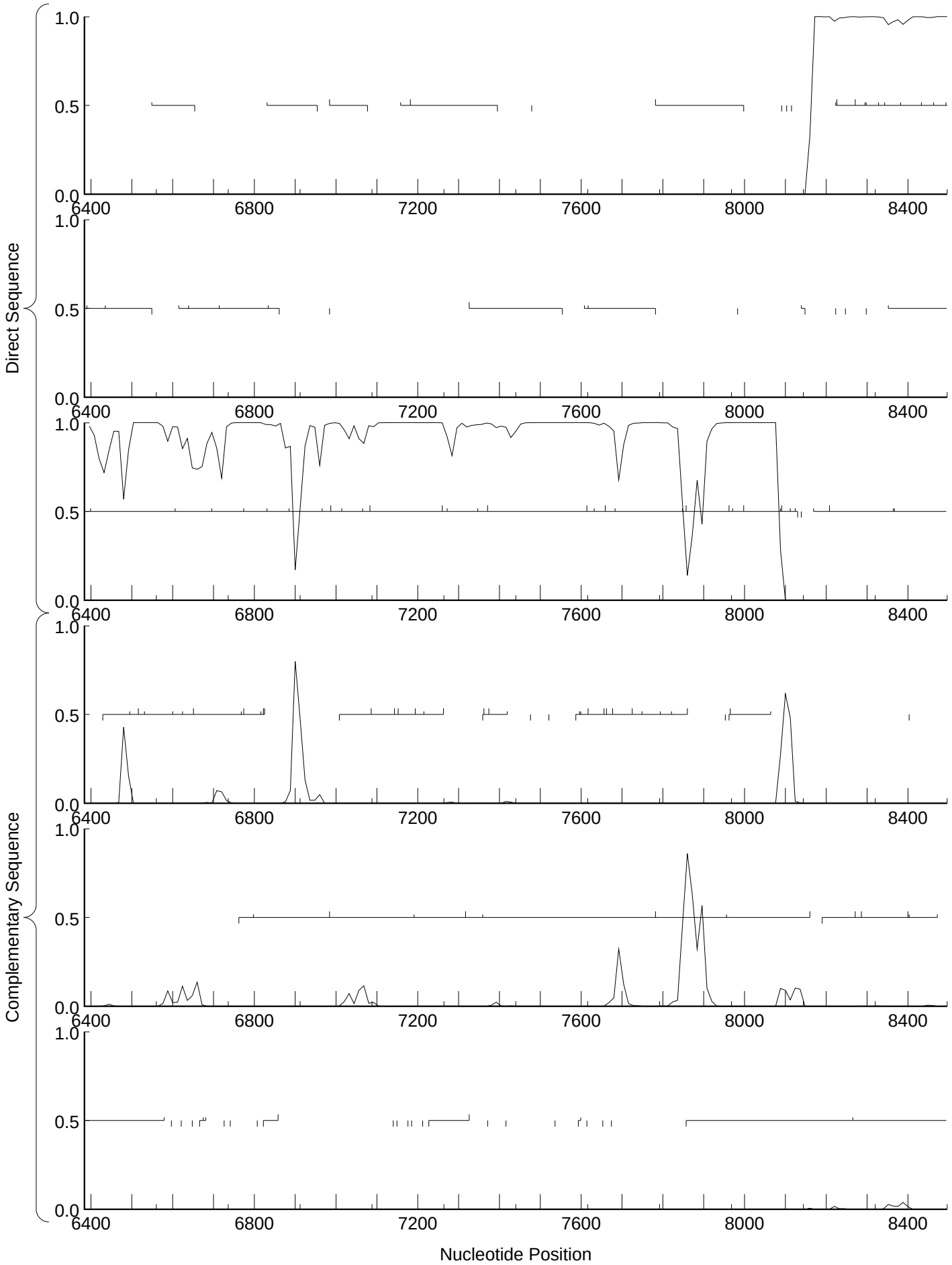

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

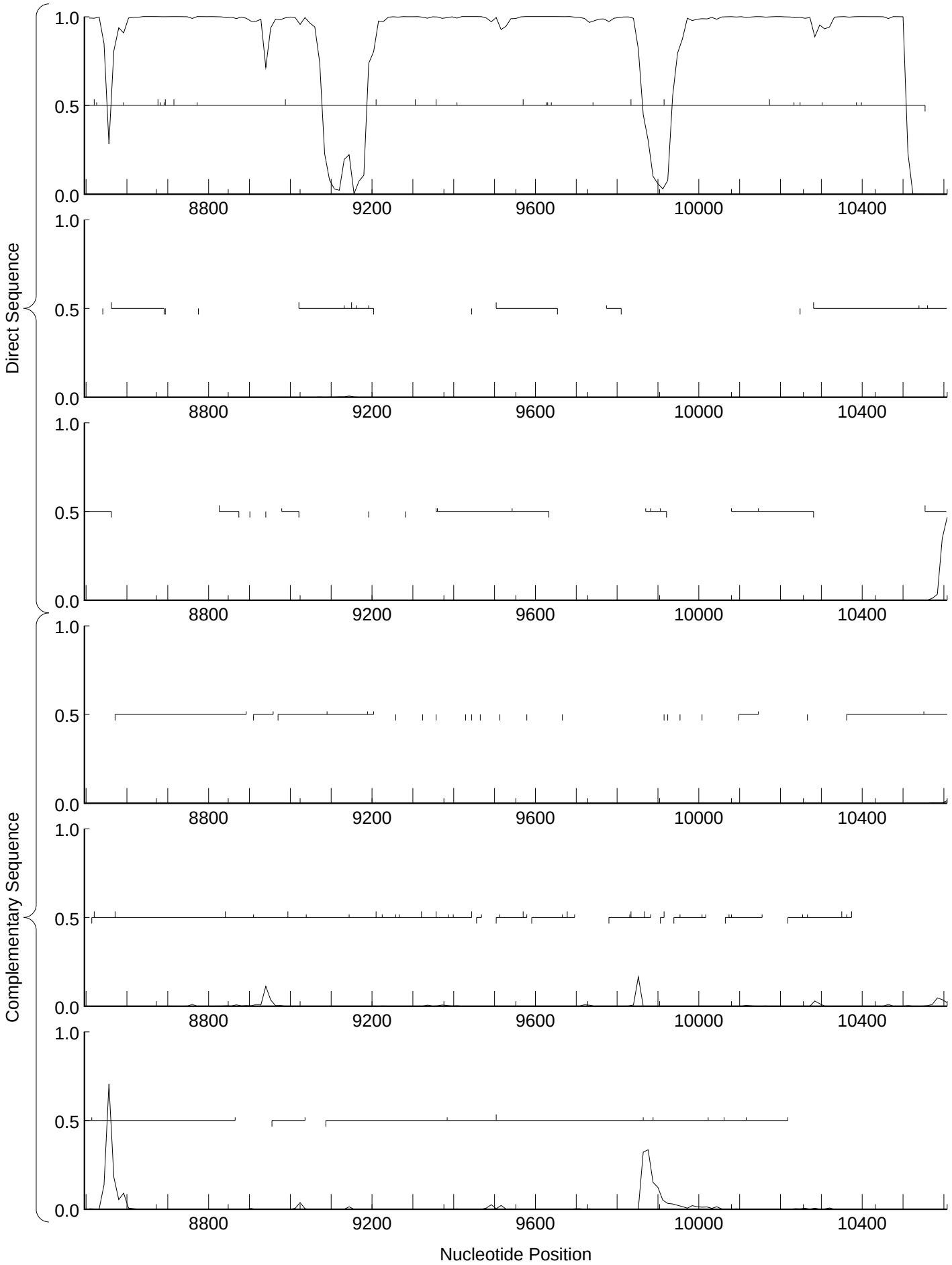

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

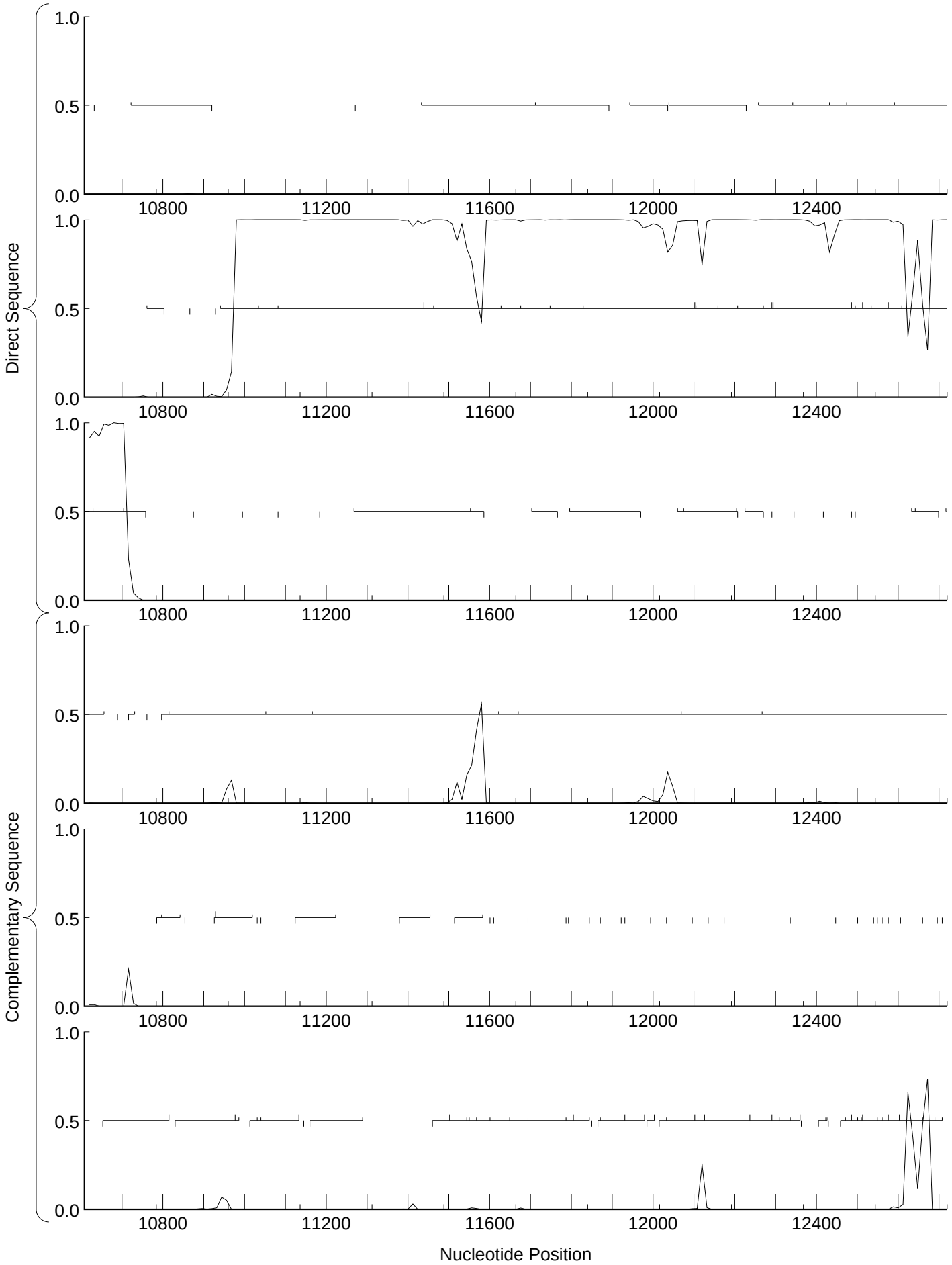

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Complementary Sequence
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0.5
0.0
400
800
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1600
2000
Nucleotide Position


Direct Sequence
Complementary Sequence
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Nucleotide Position

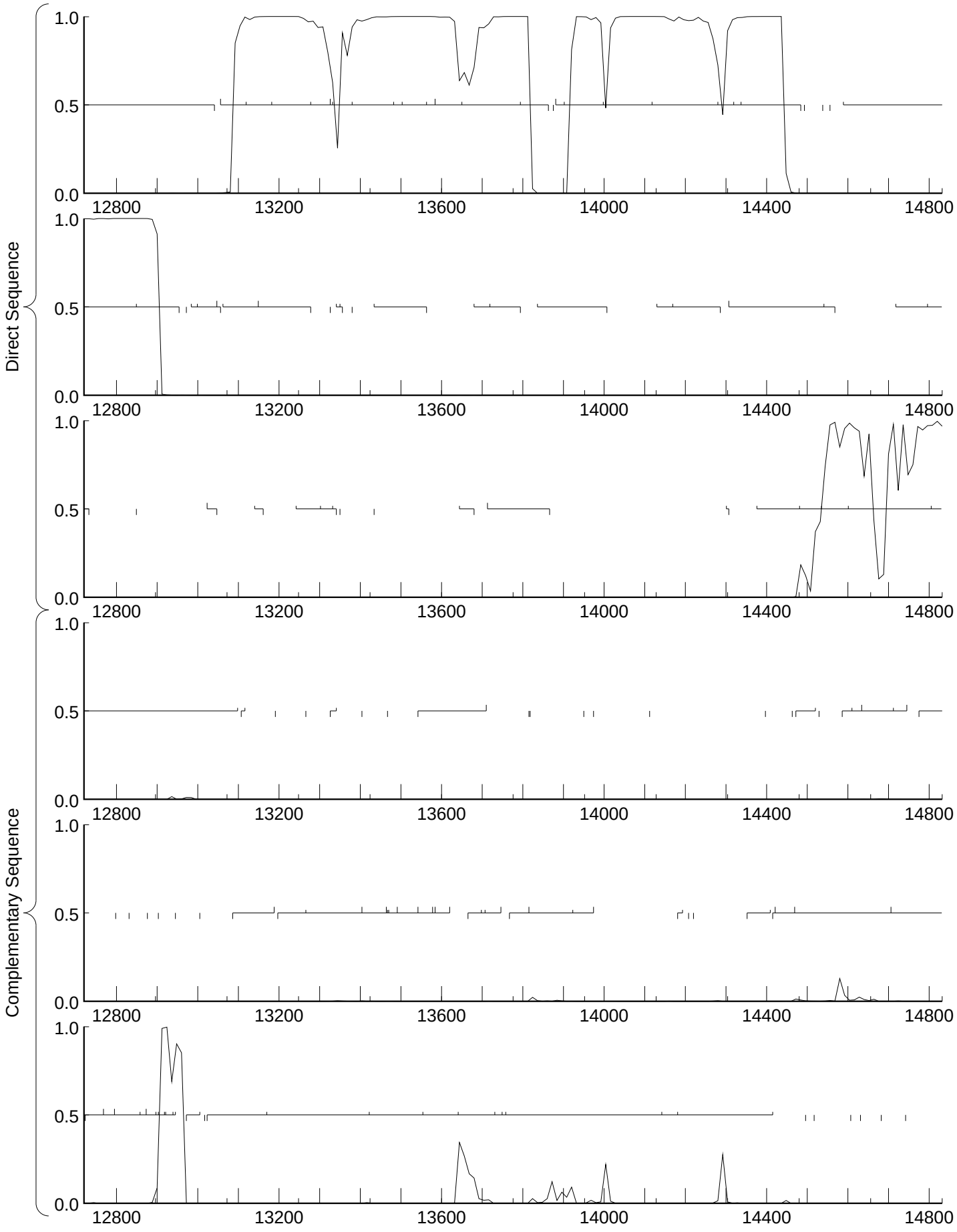

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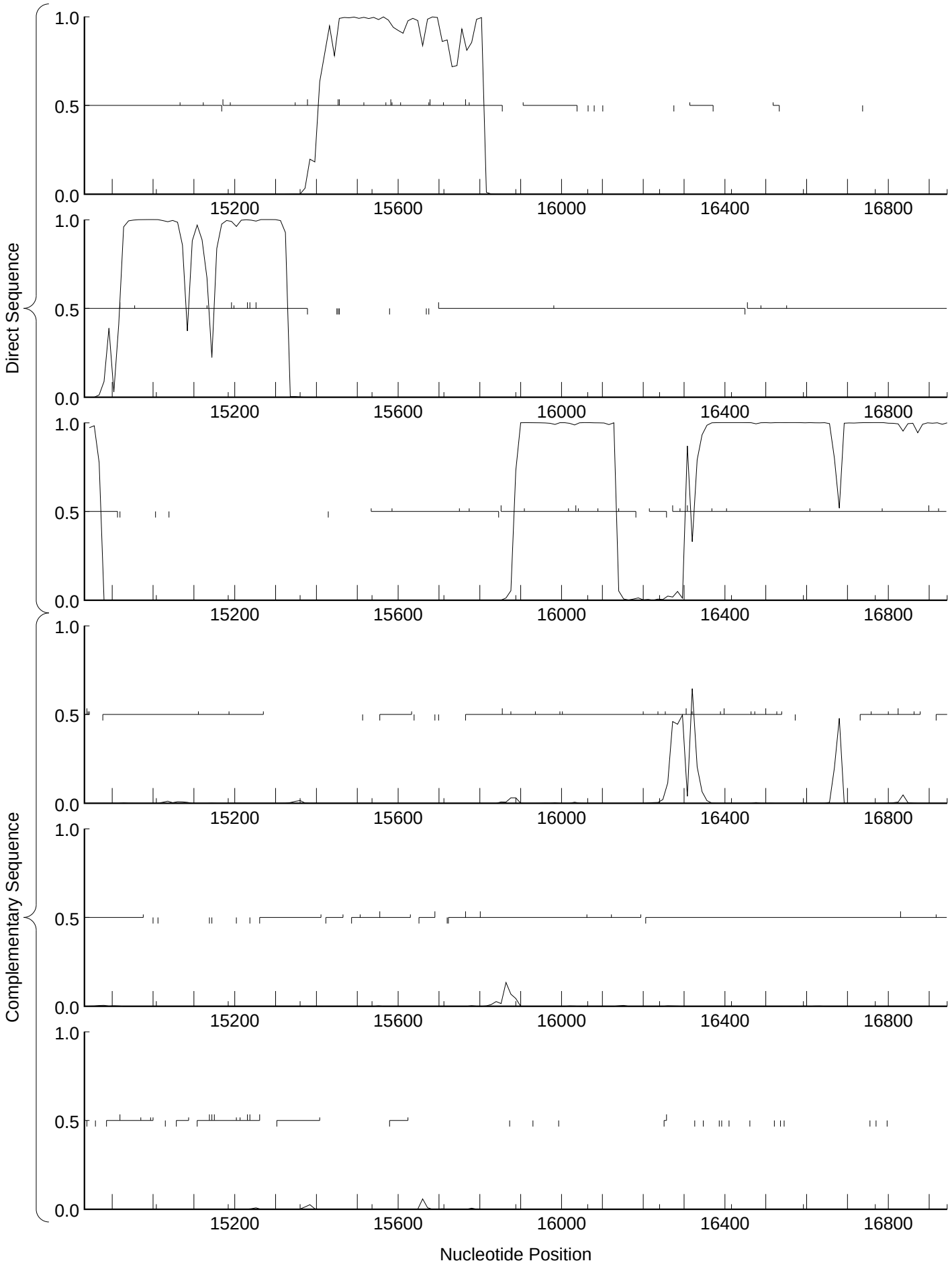

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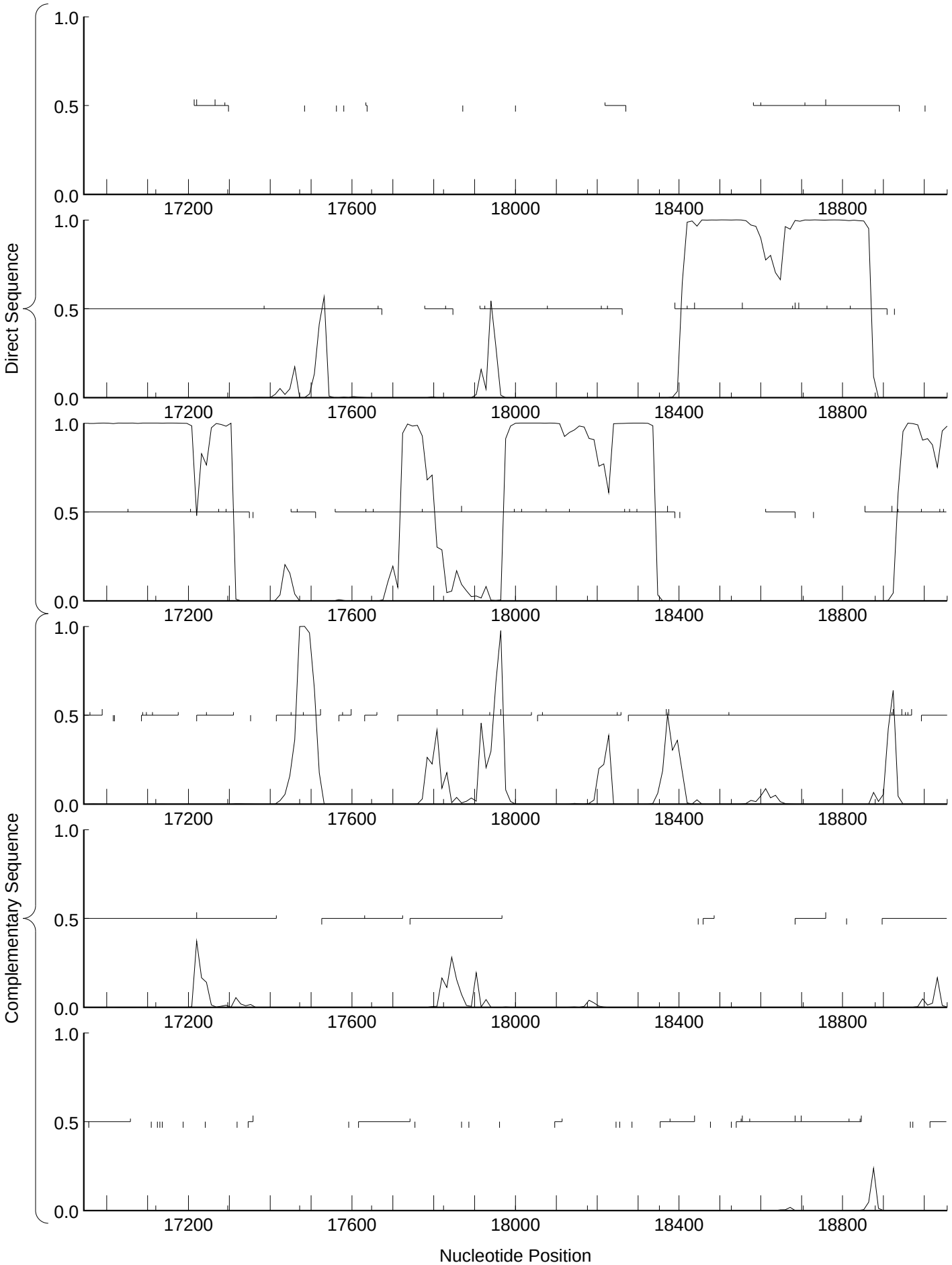

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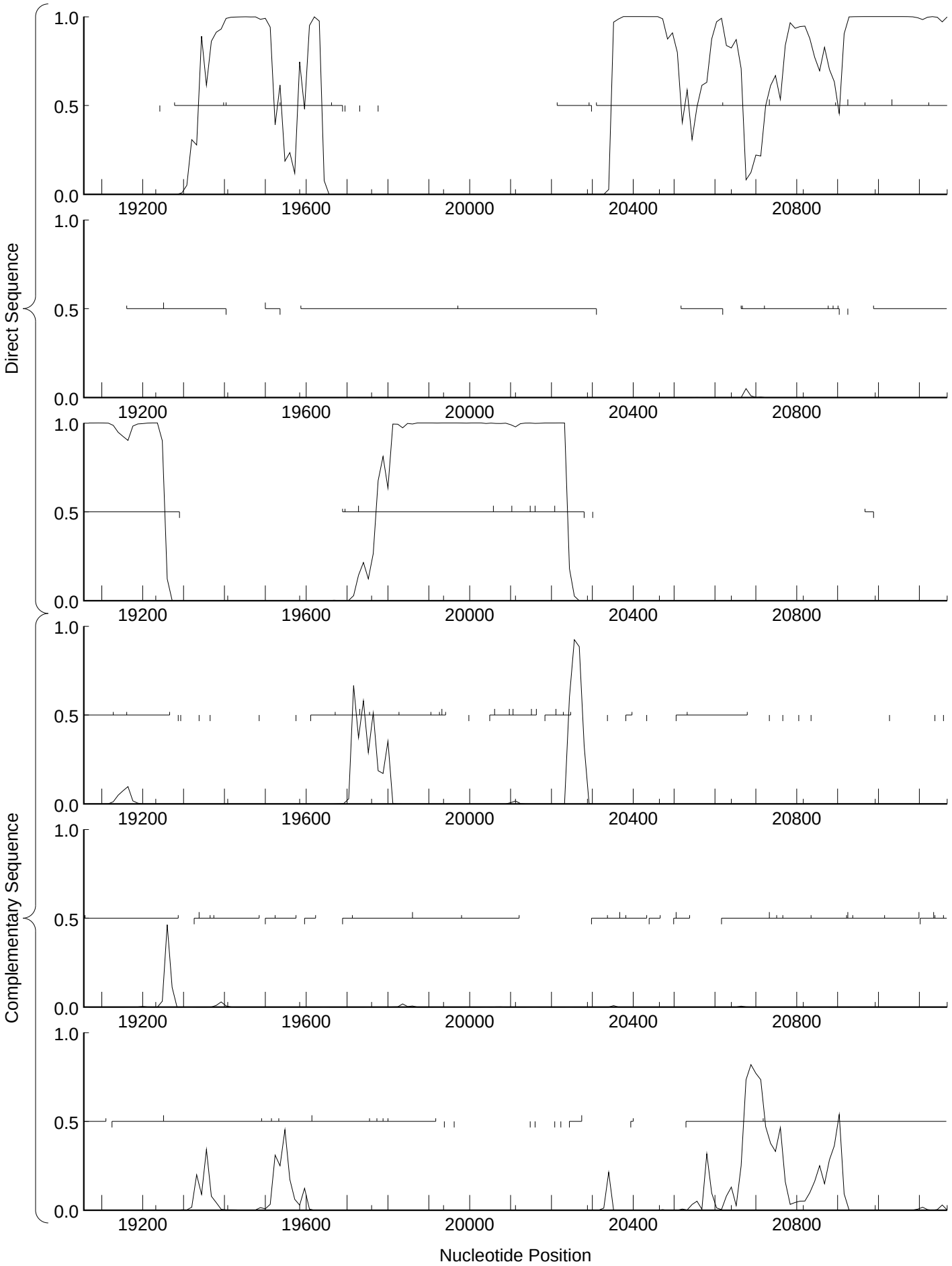

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Nucleotide Position

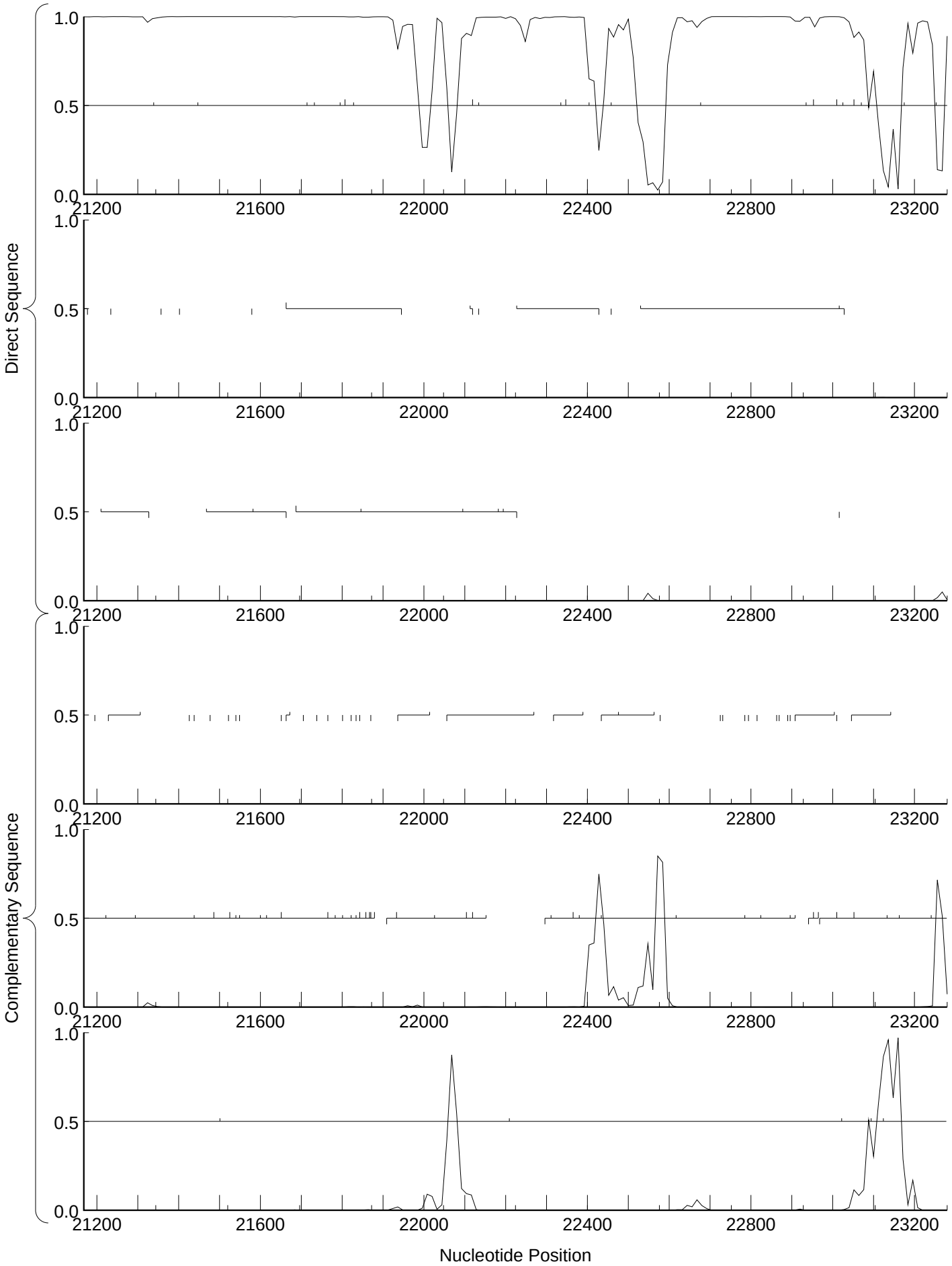
Mycobacterium phage Phaedrus complete genome, Order 4, Window 96, Step 12, 8/34

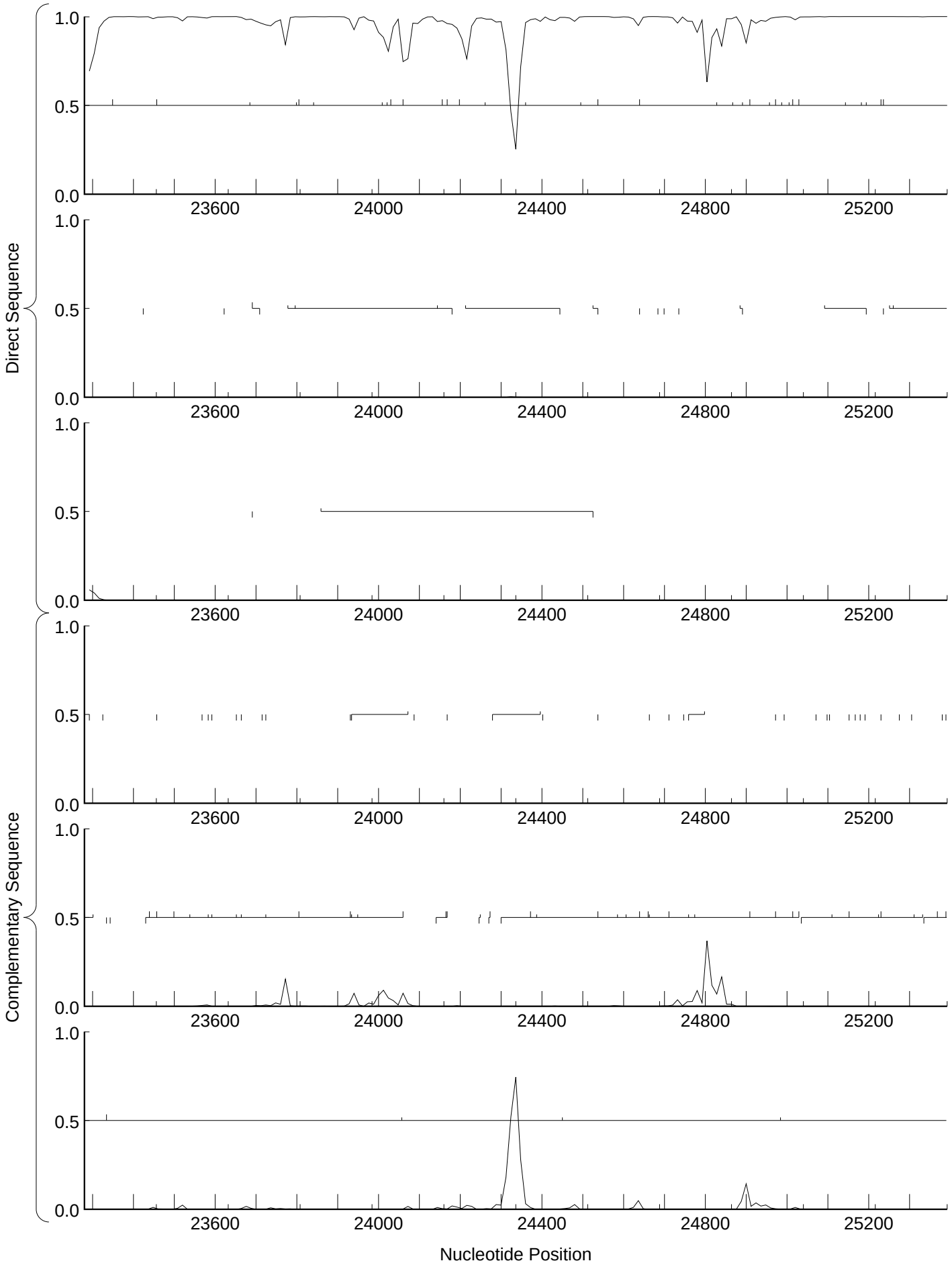


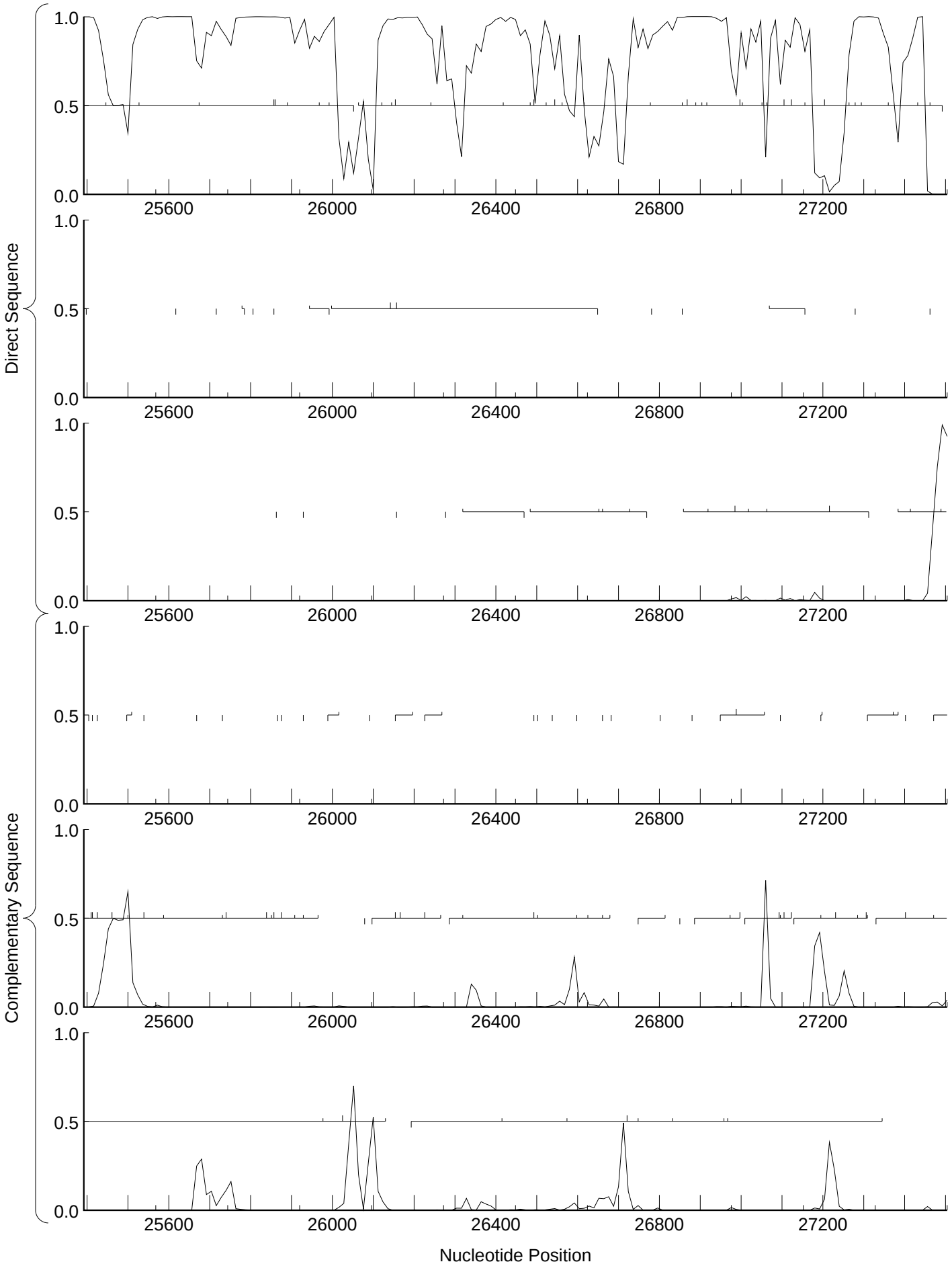

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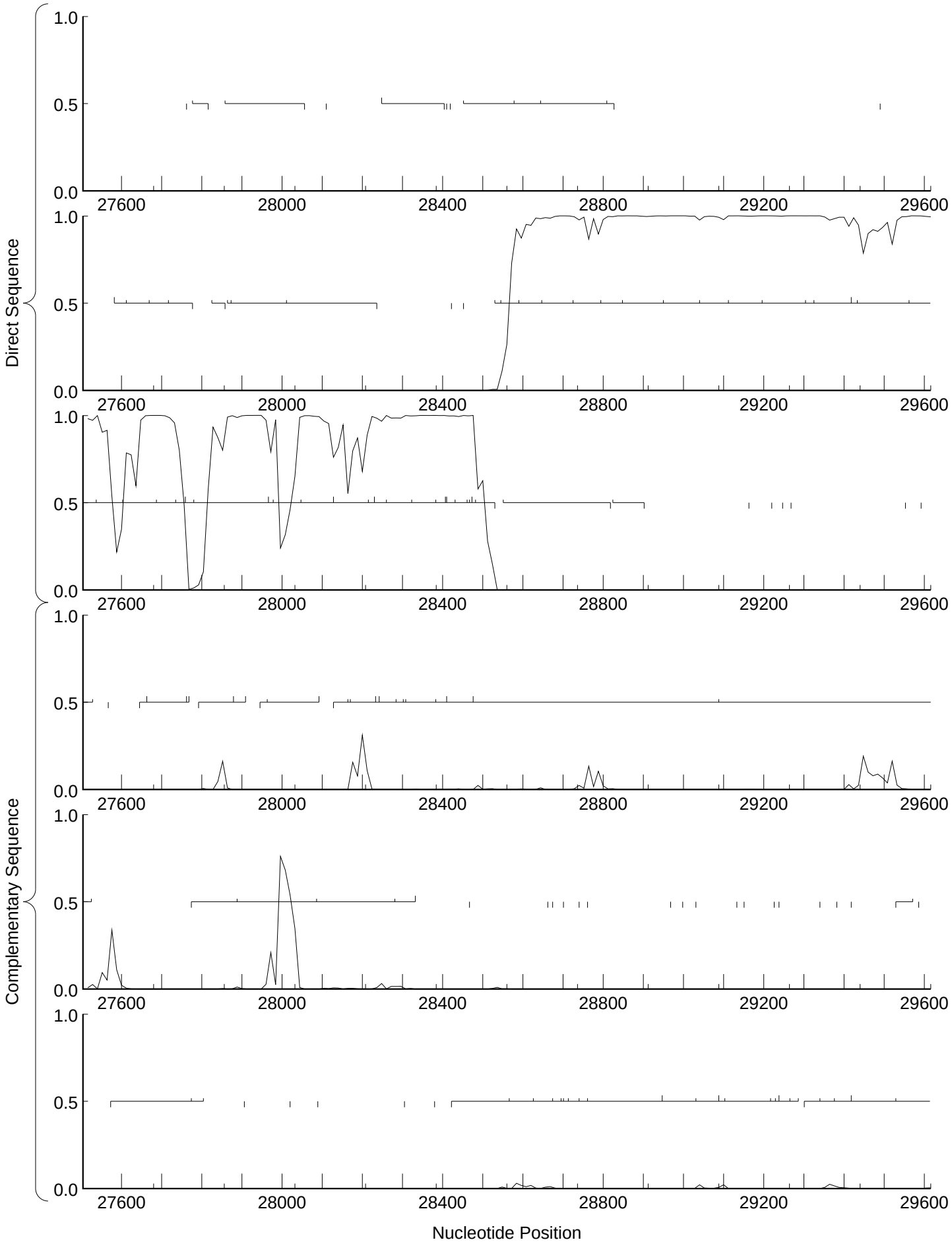

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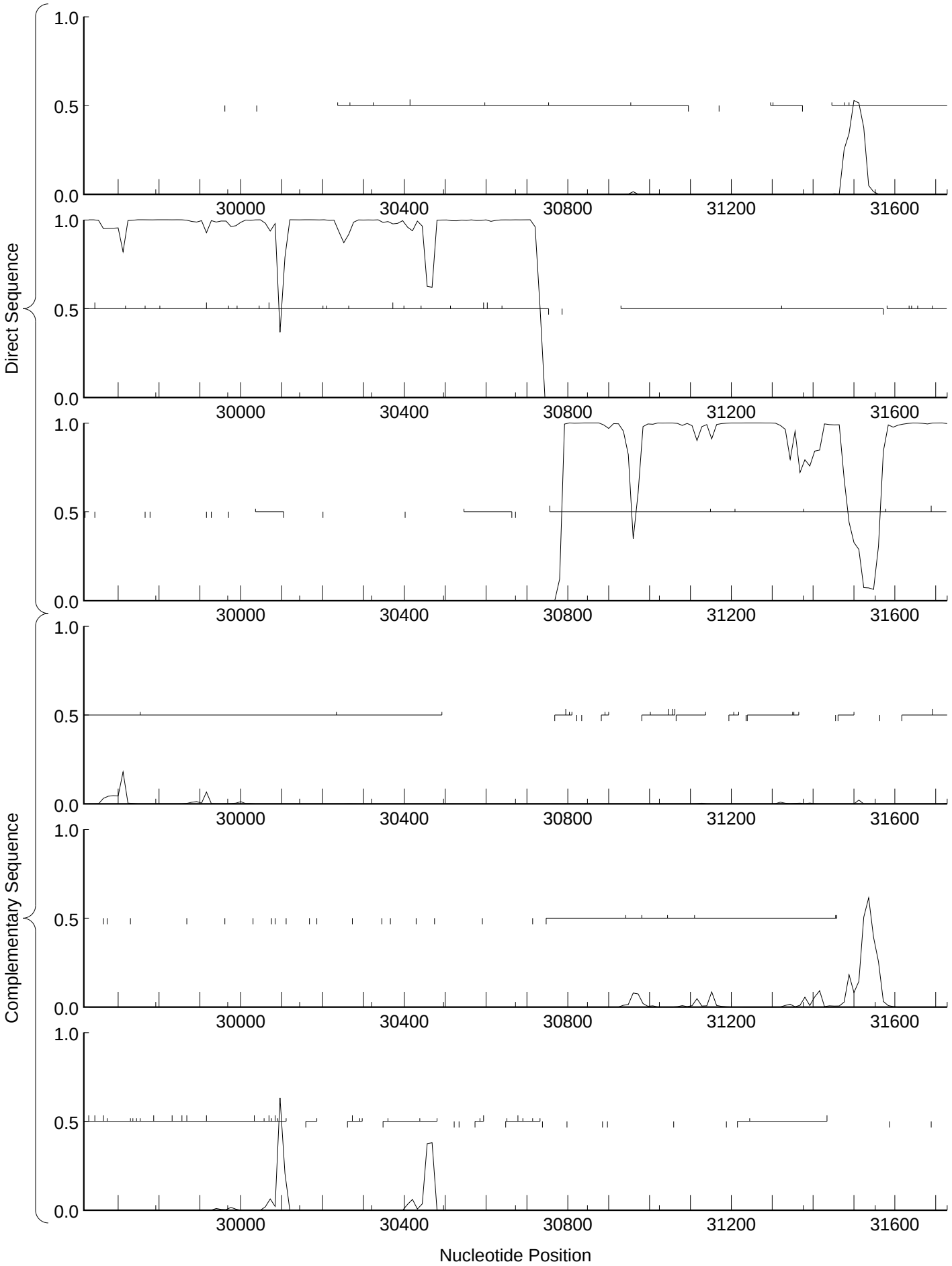

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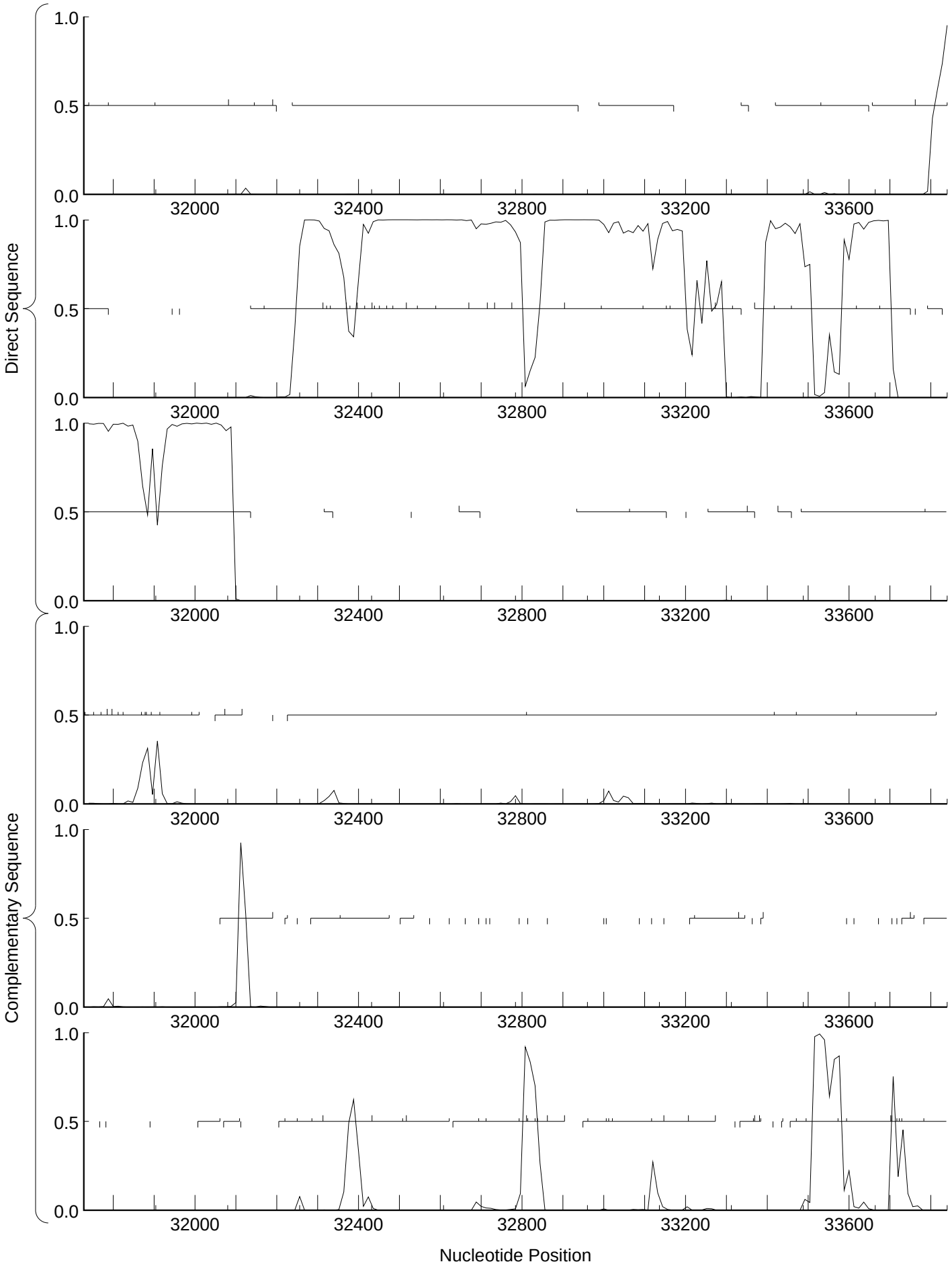

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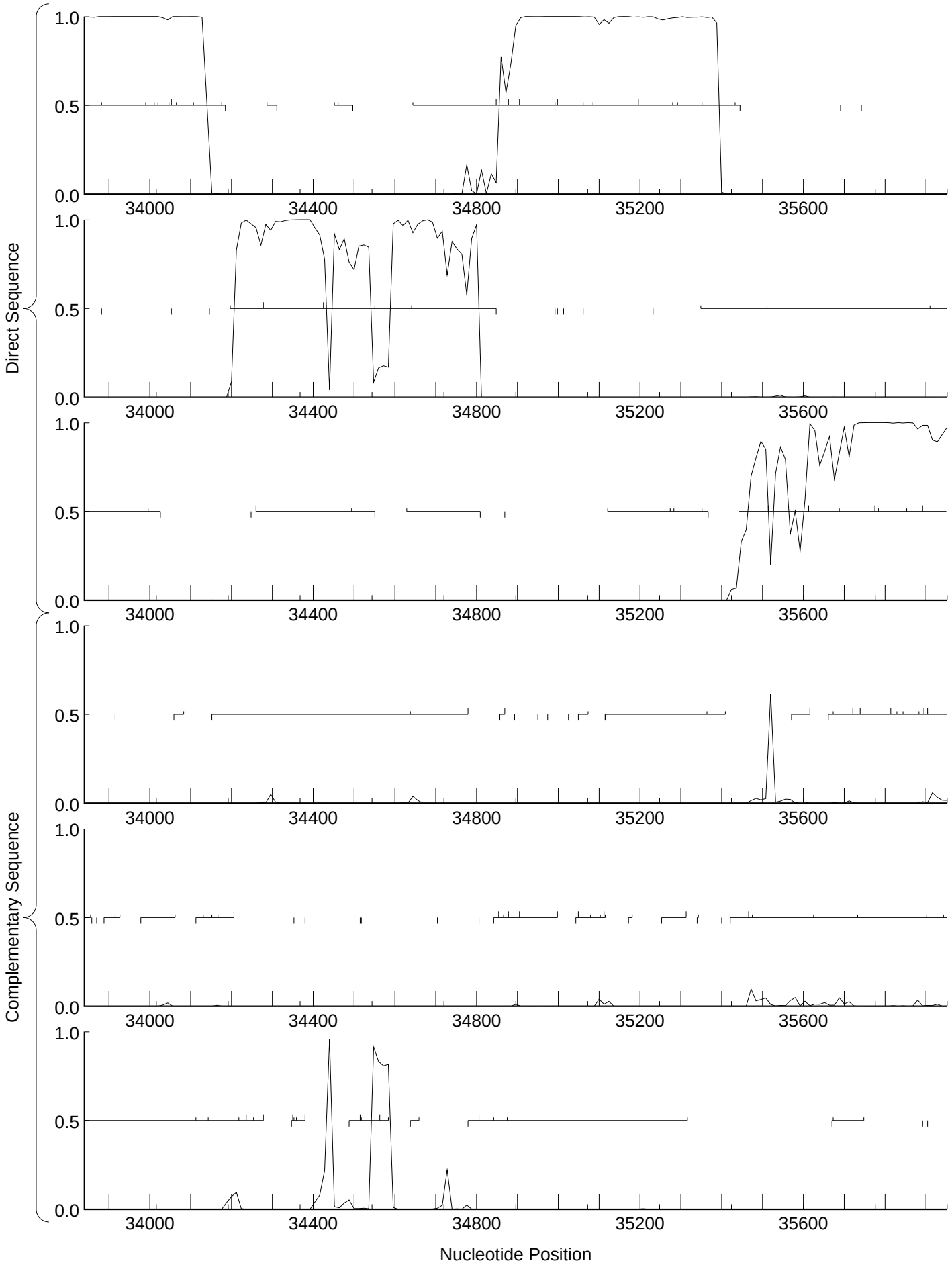

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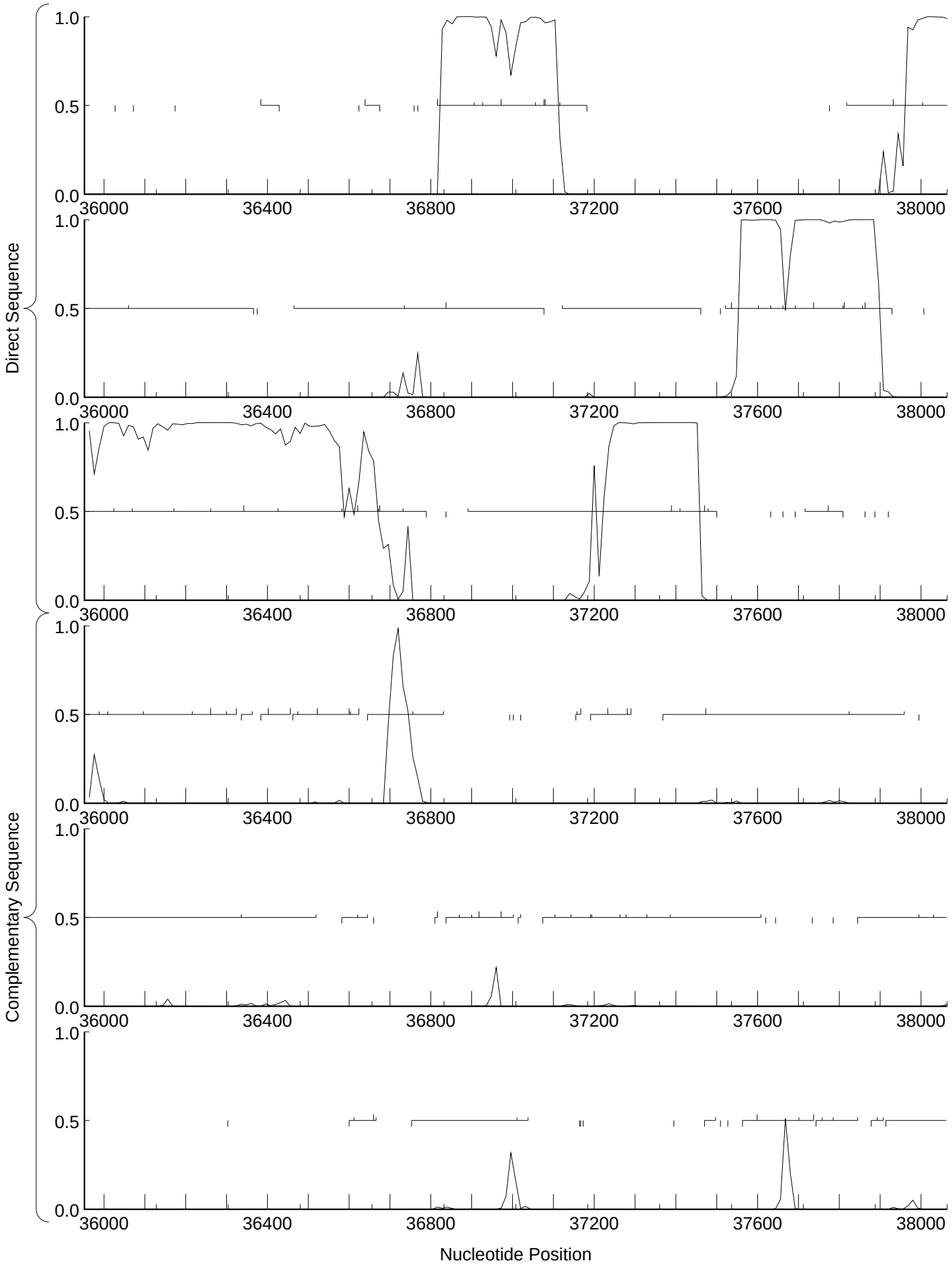

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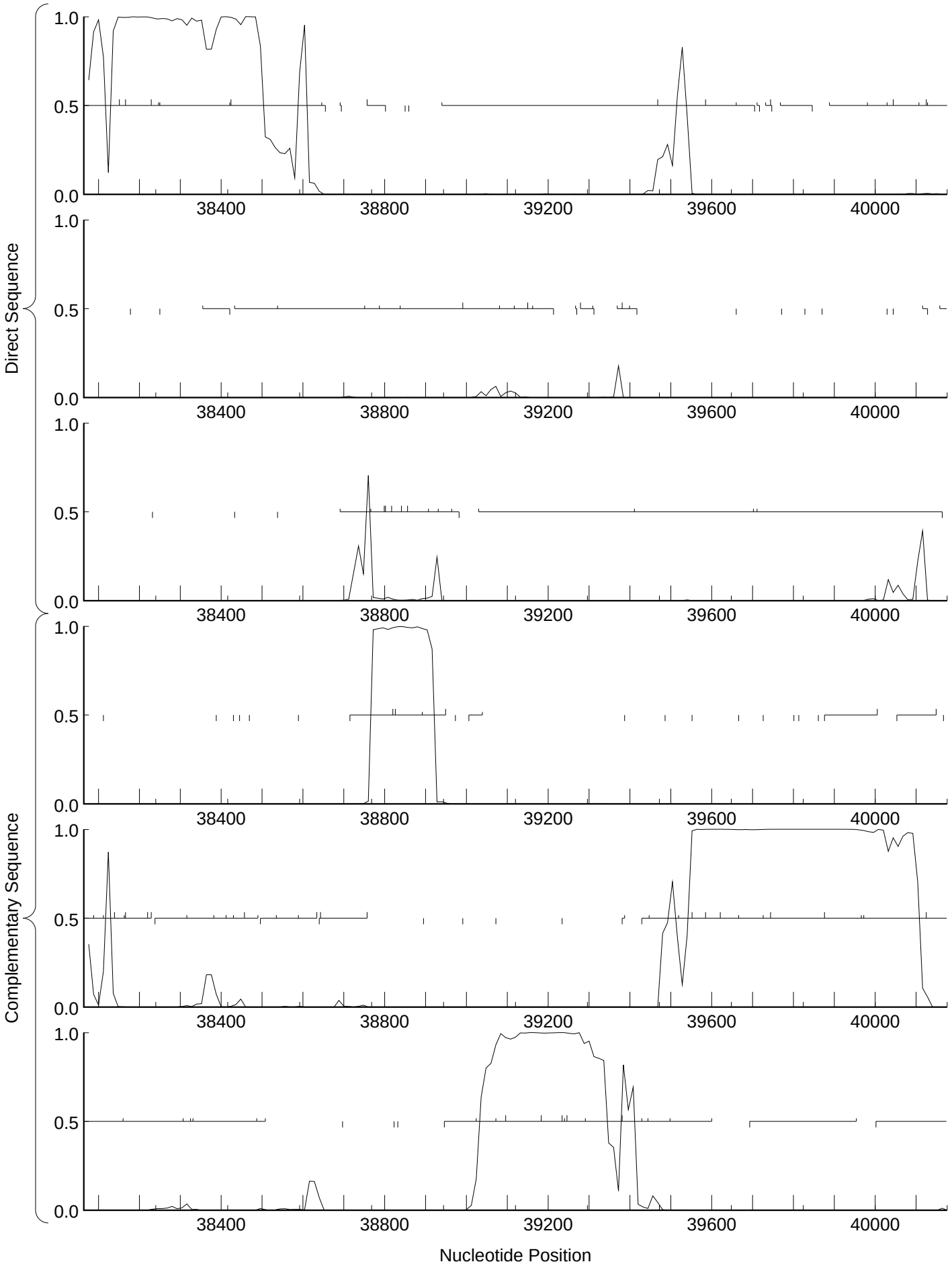

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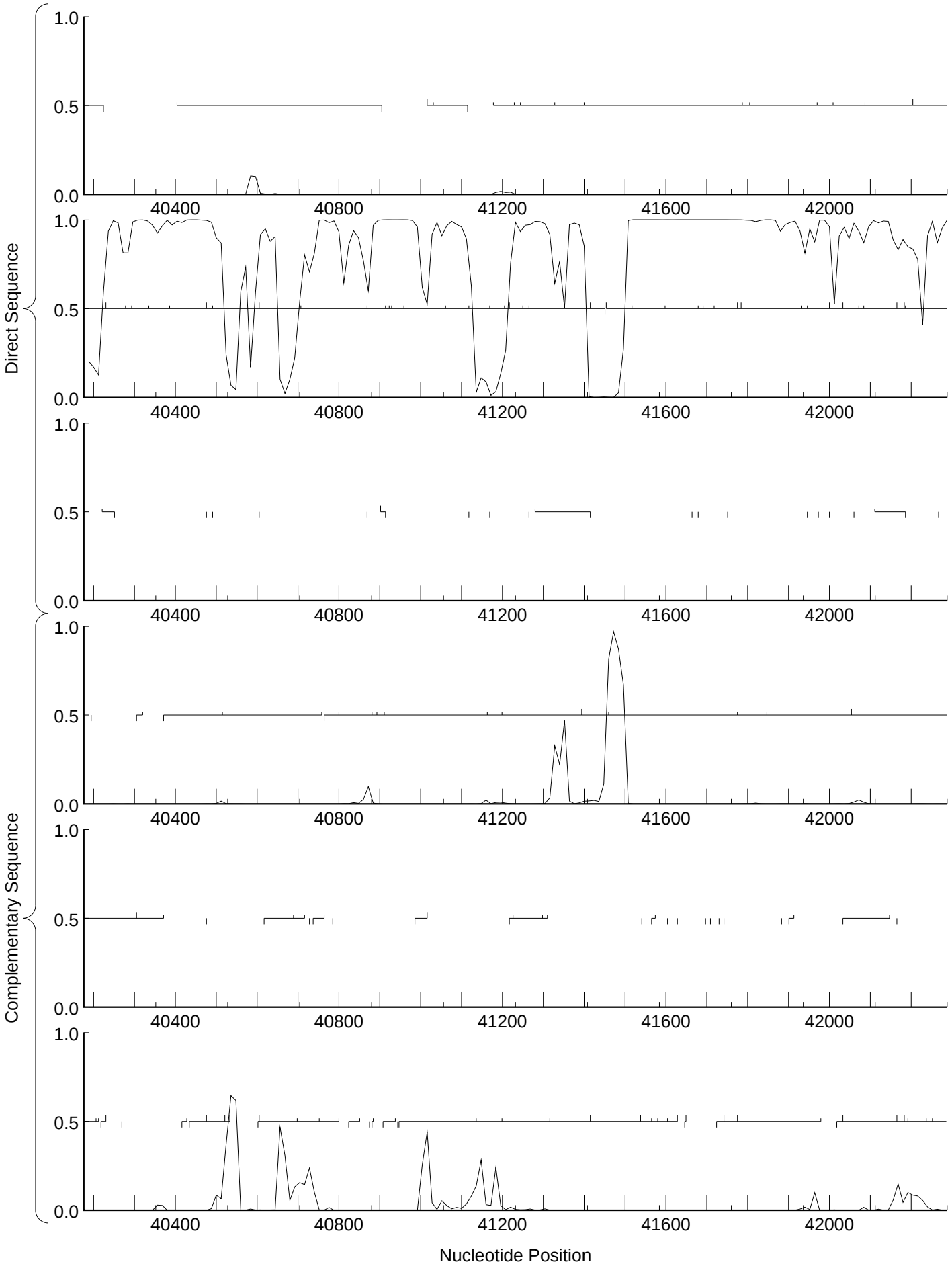

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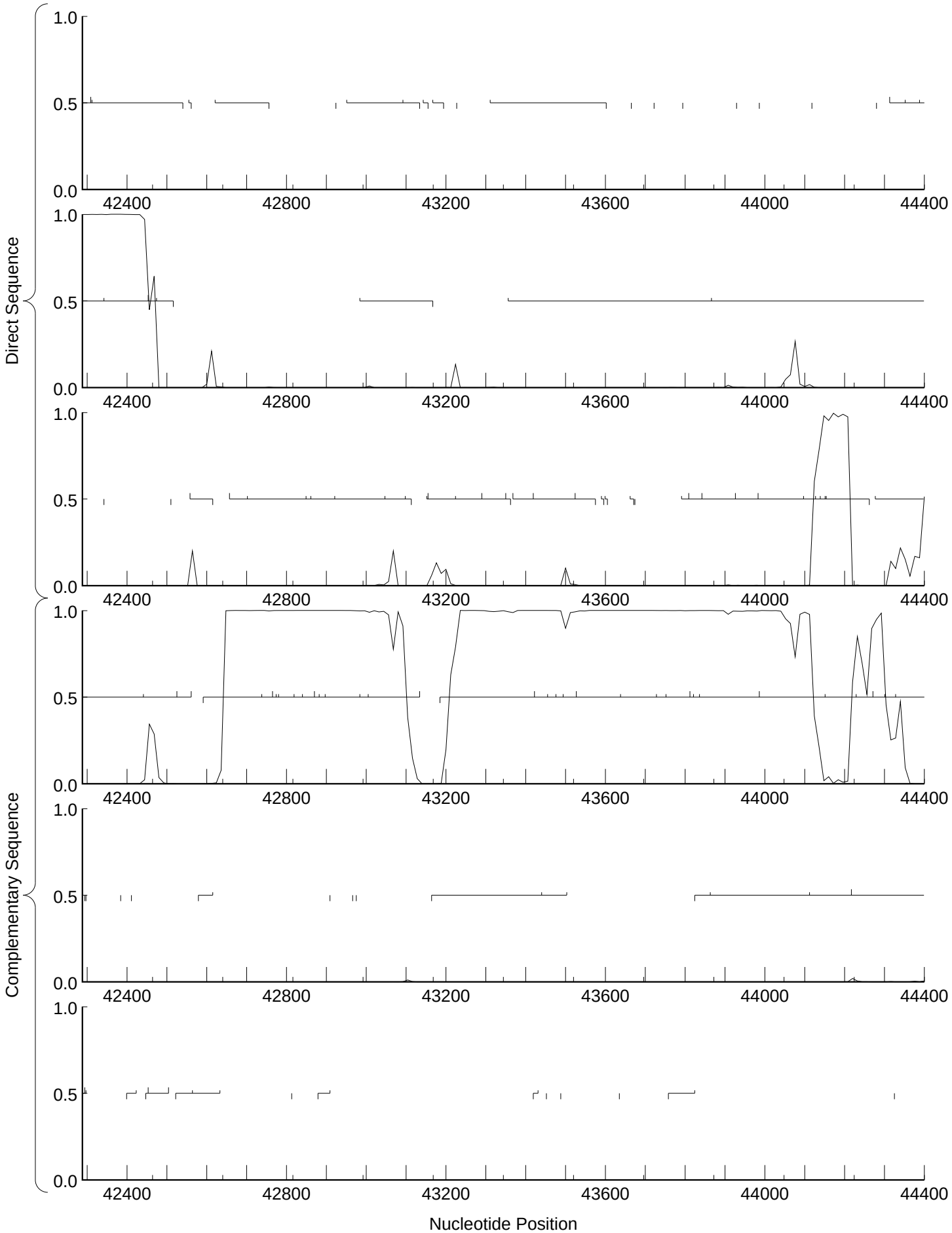

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Nucleotide Position

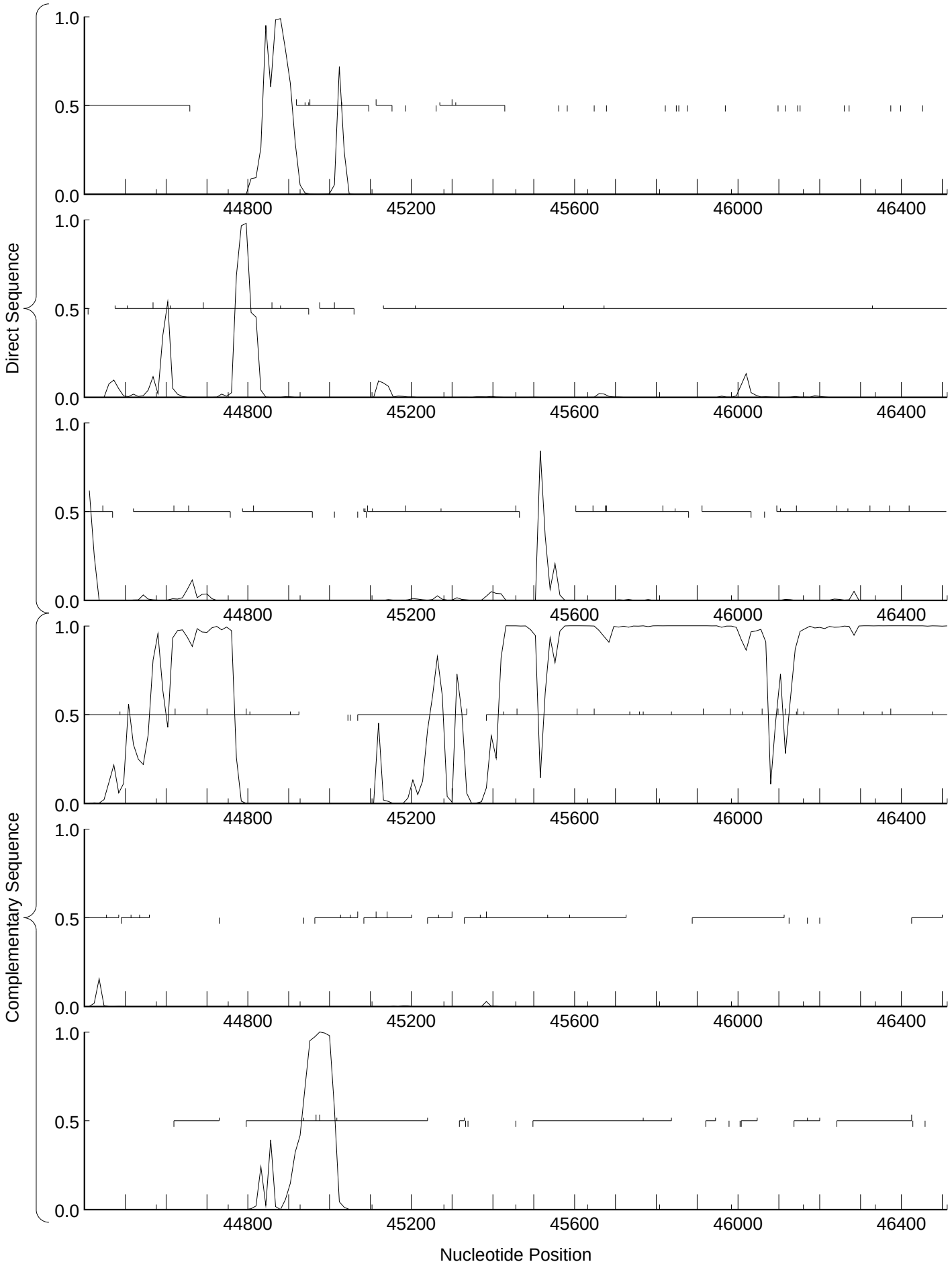

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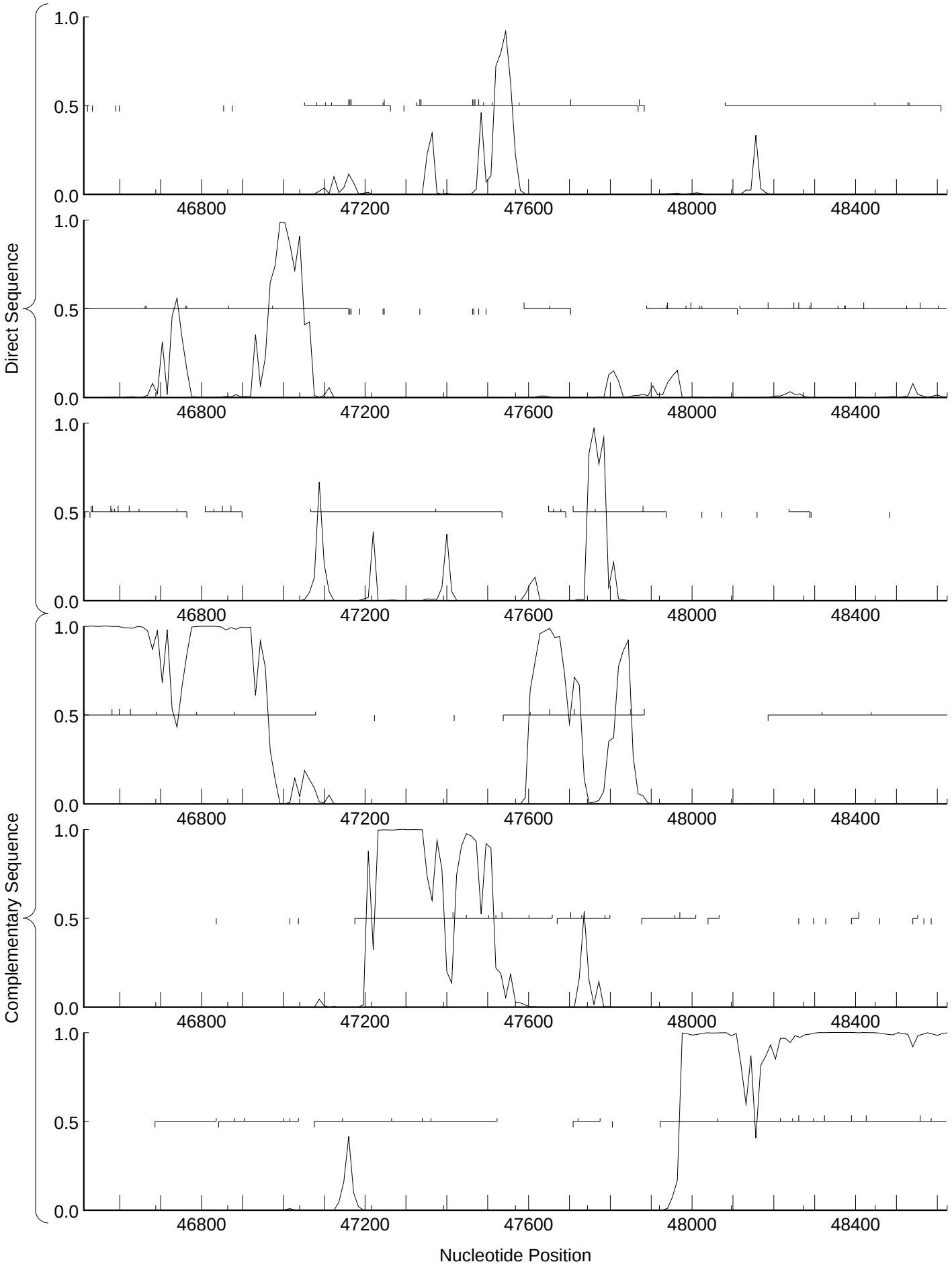

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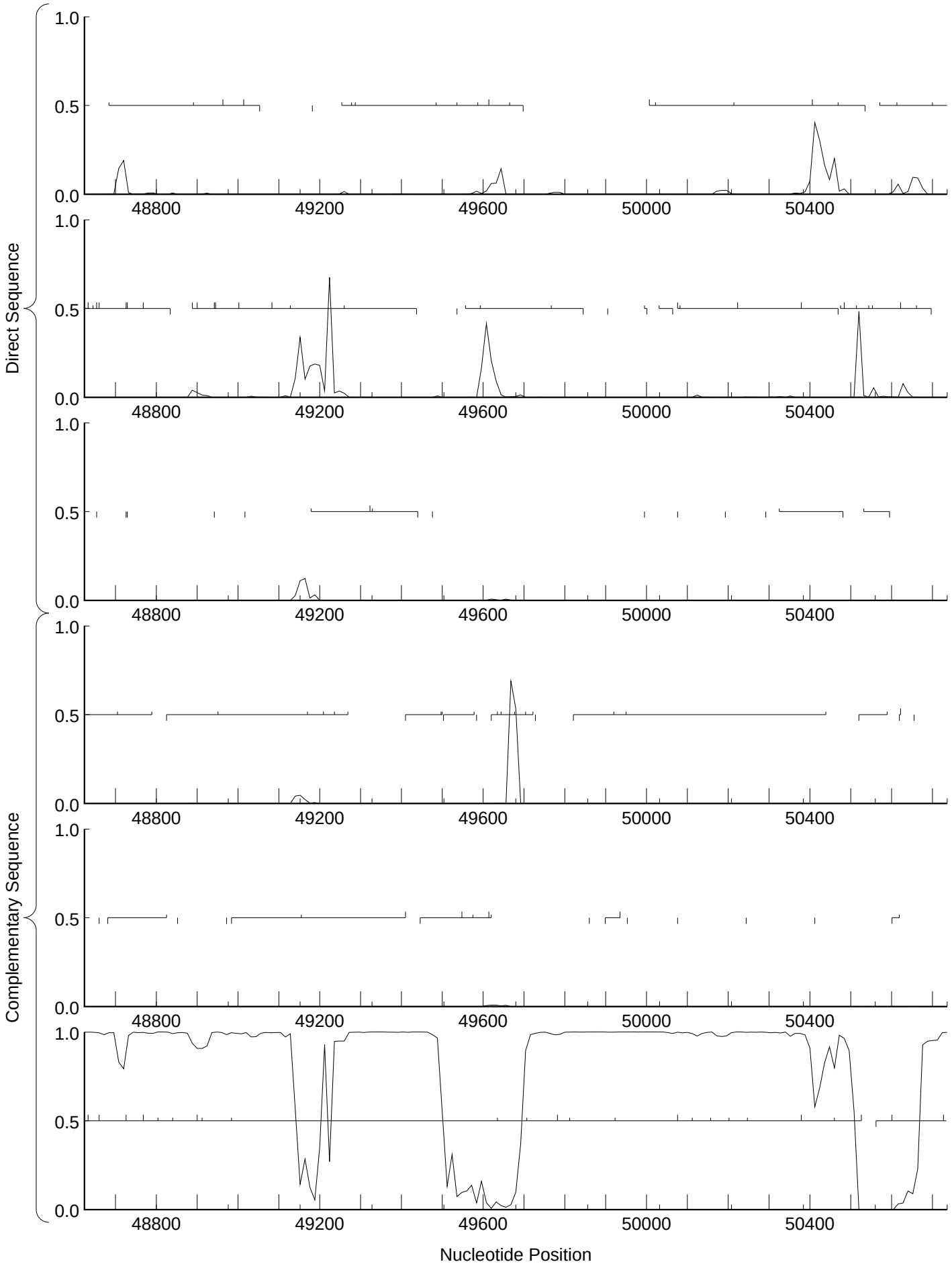

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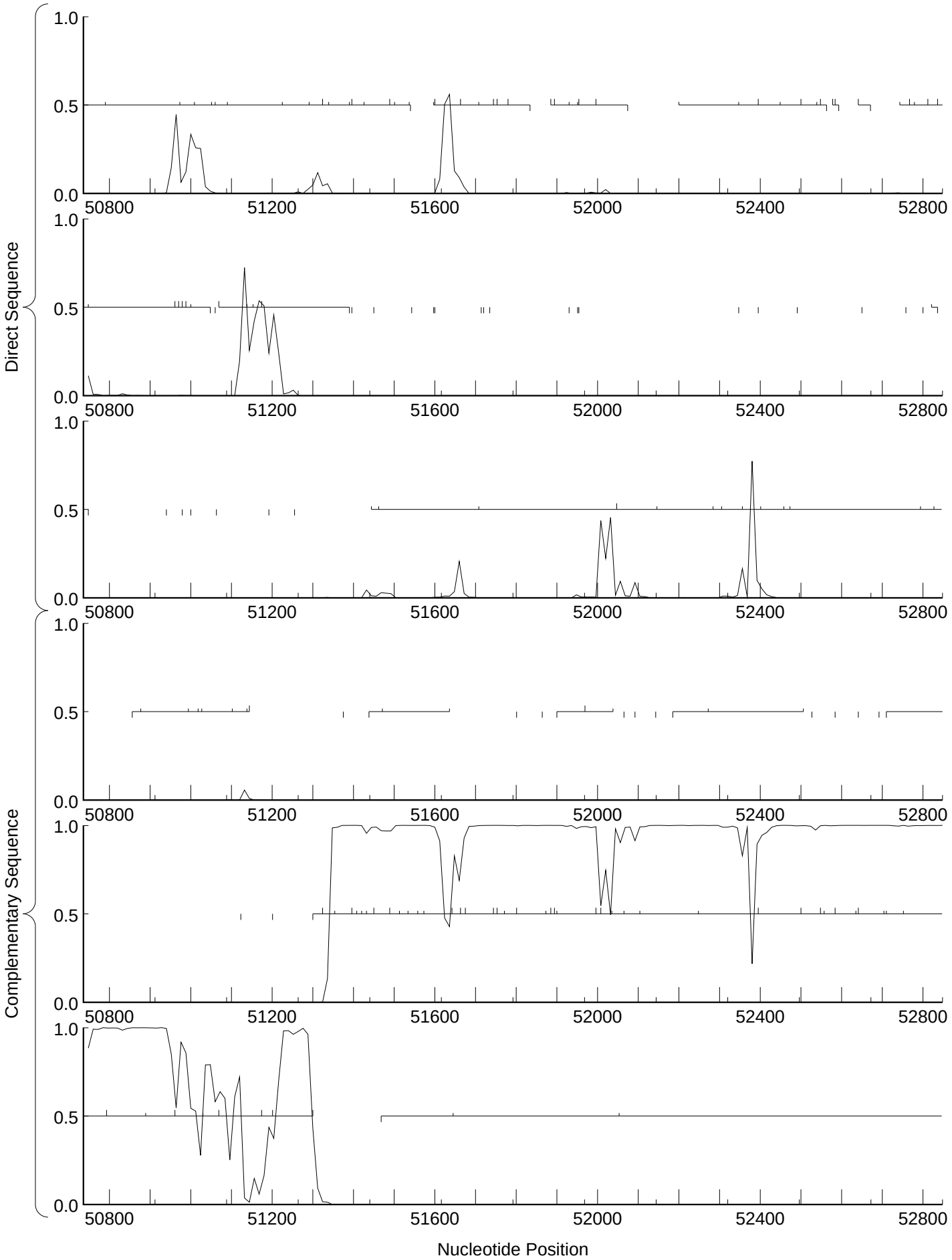

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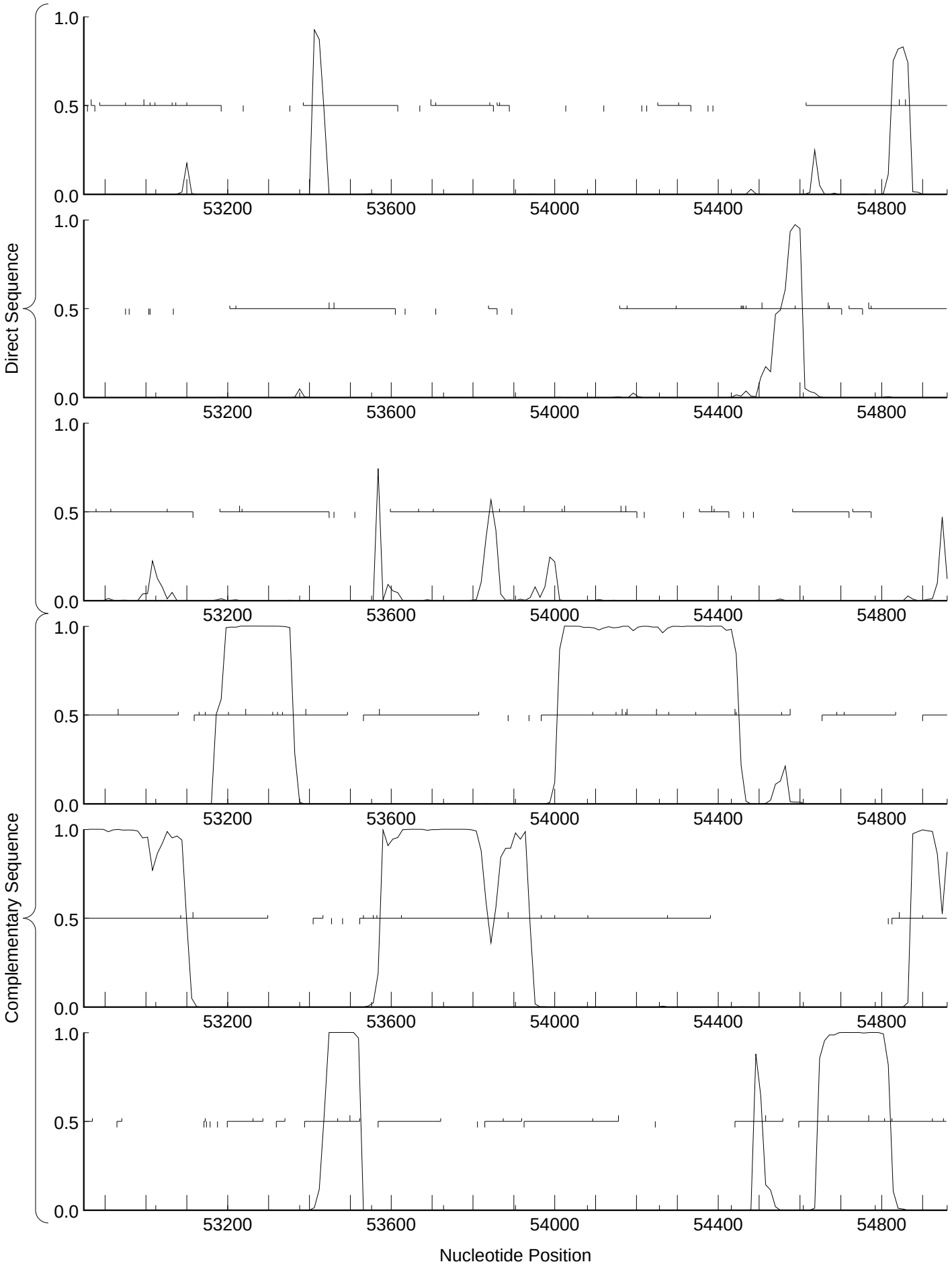

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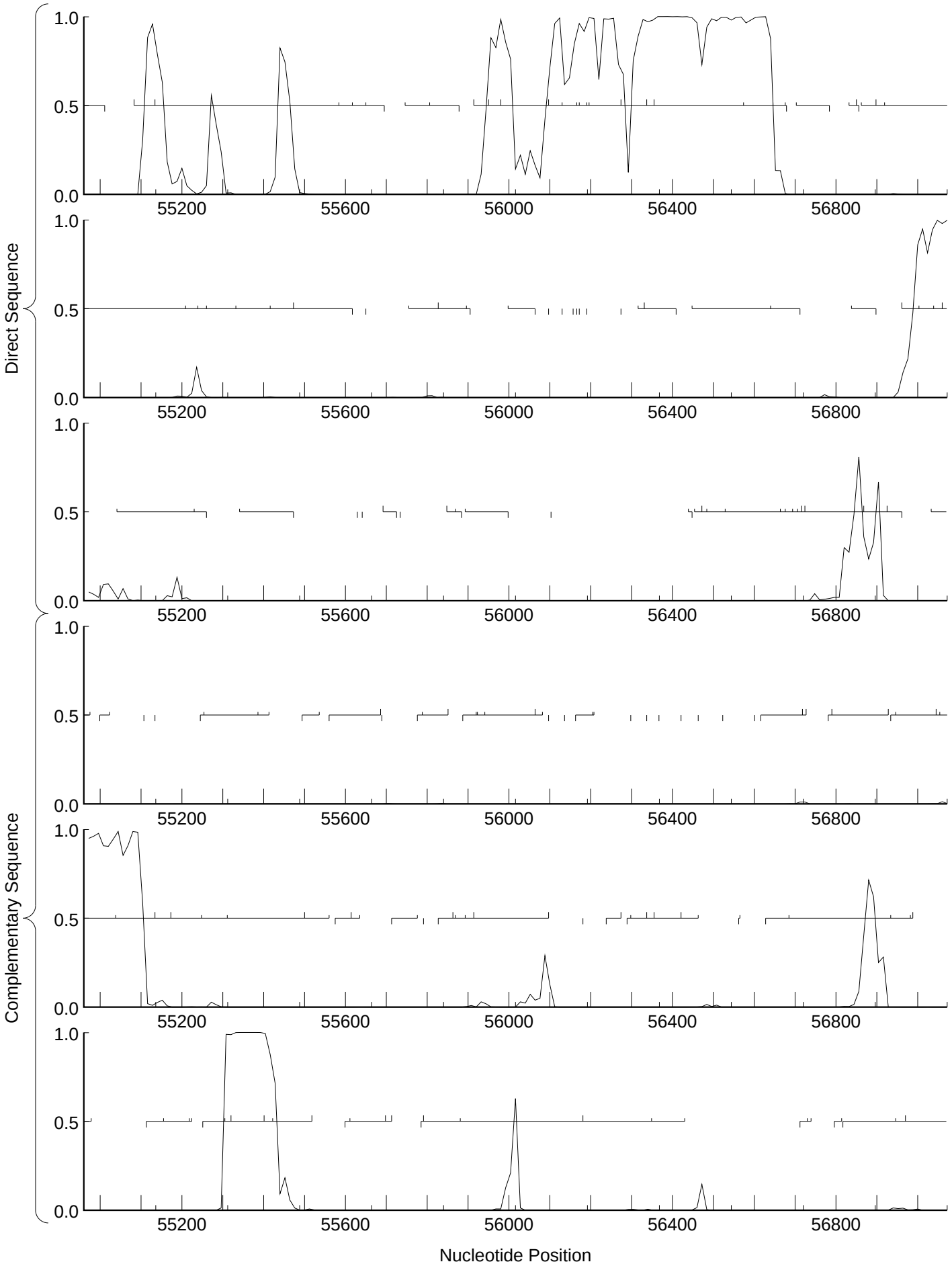

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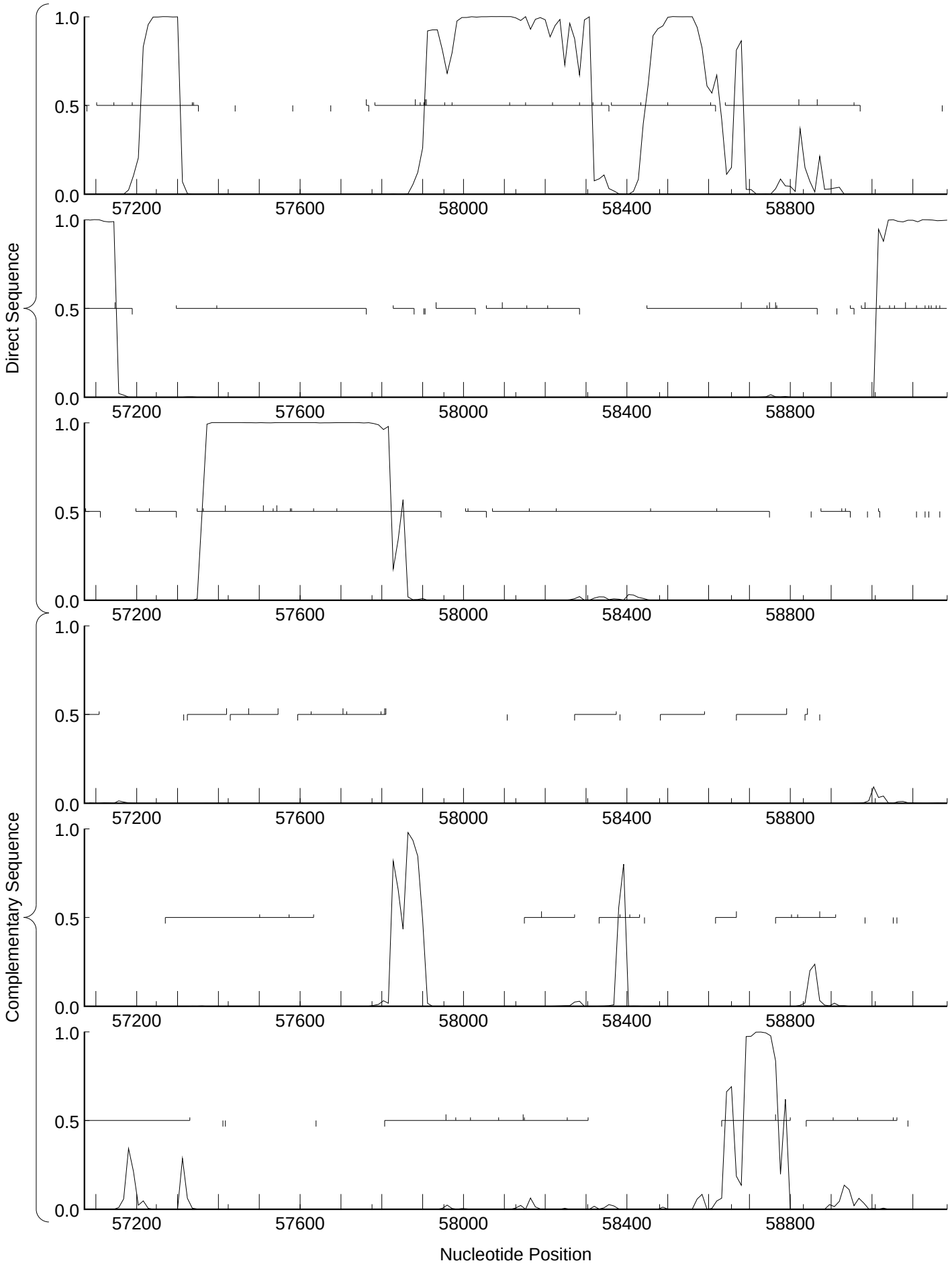

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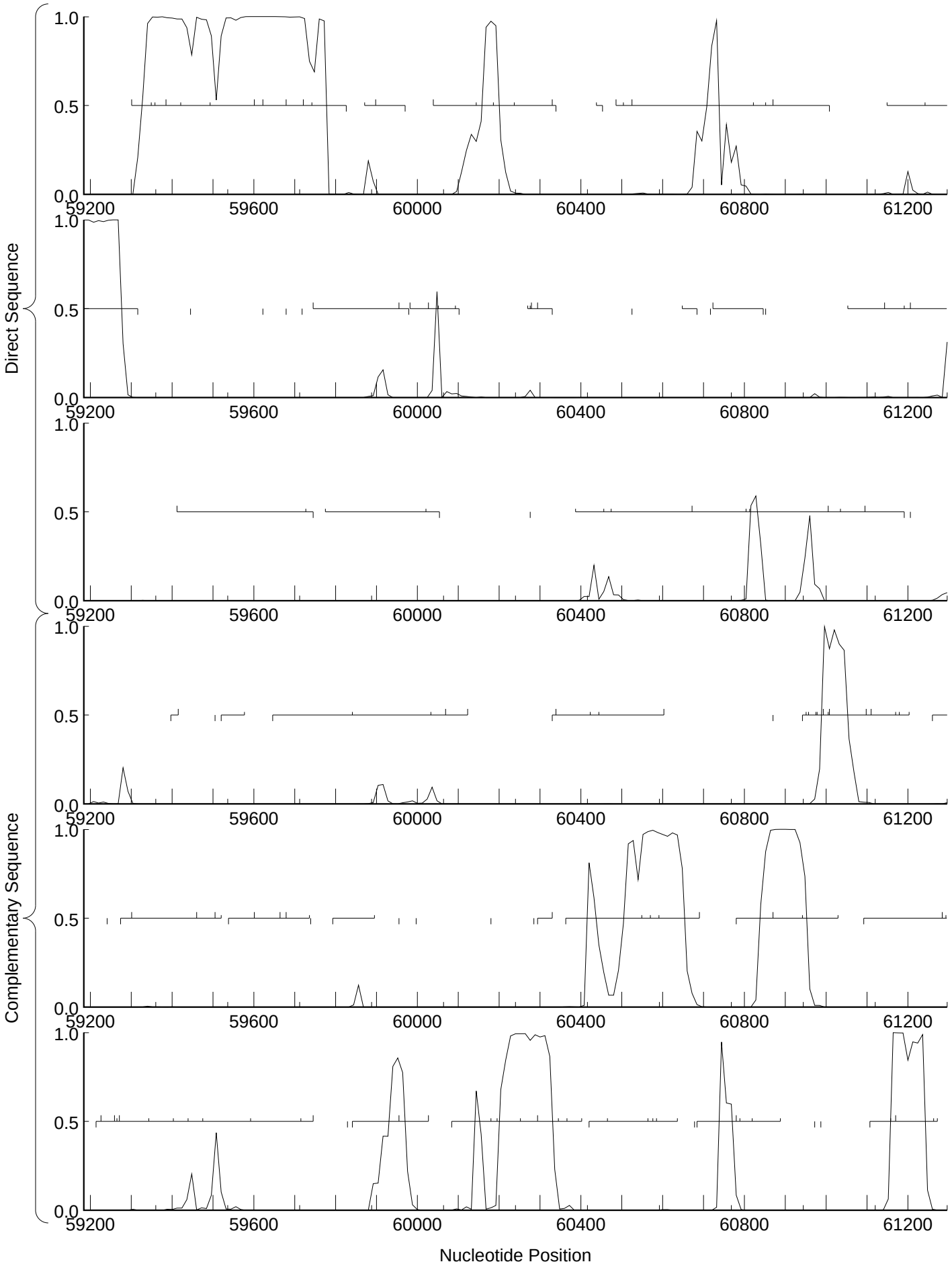

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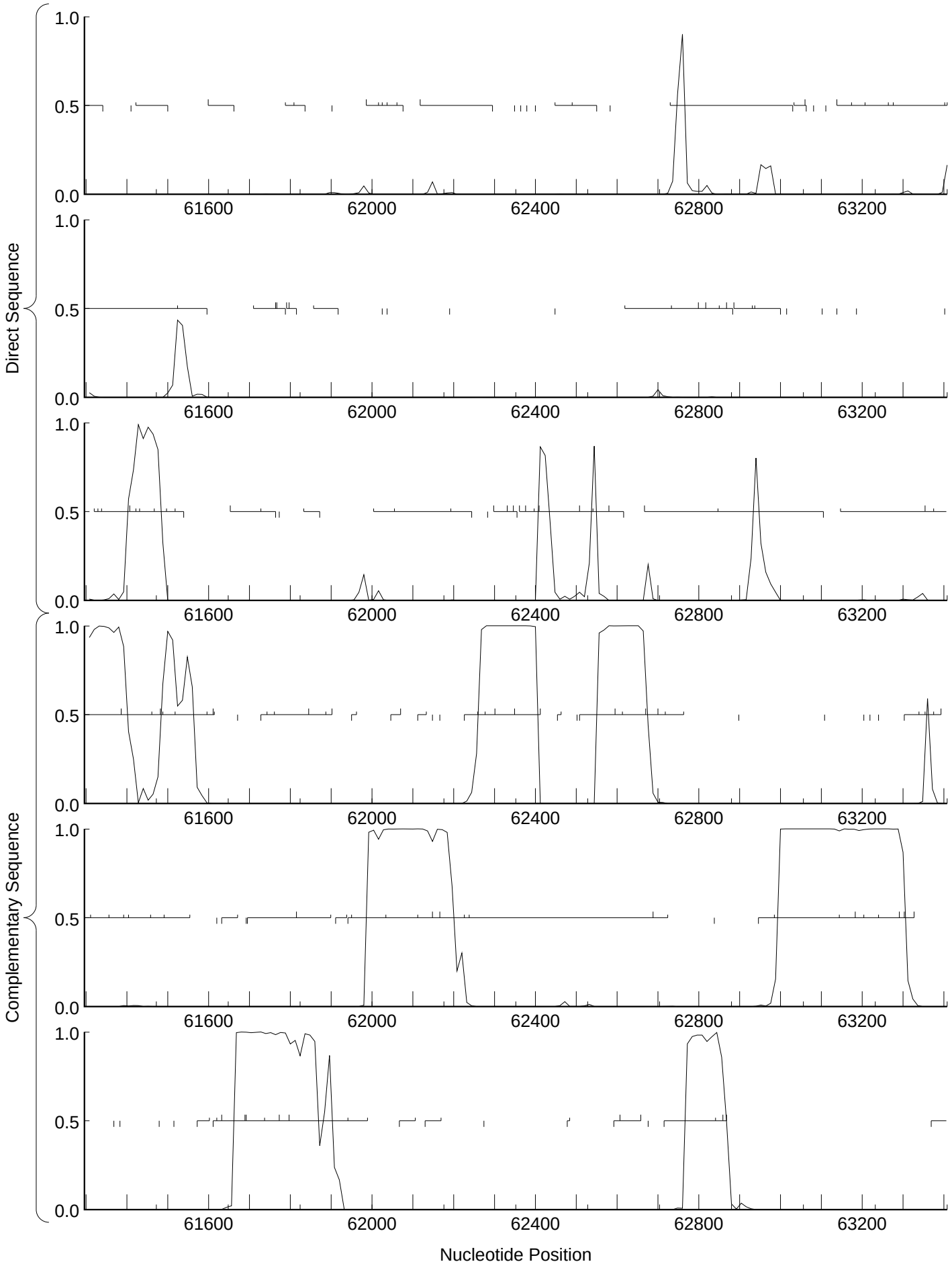

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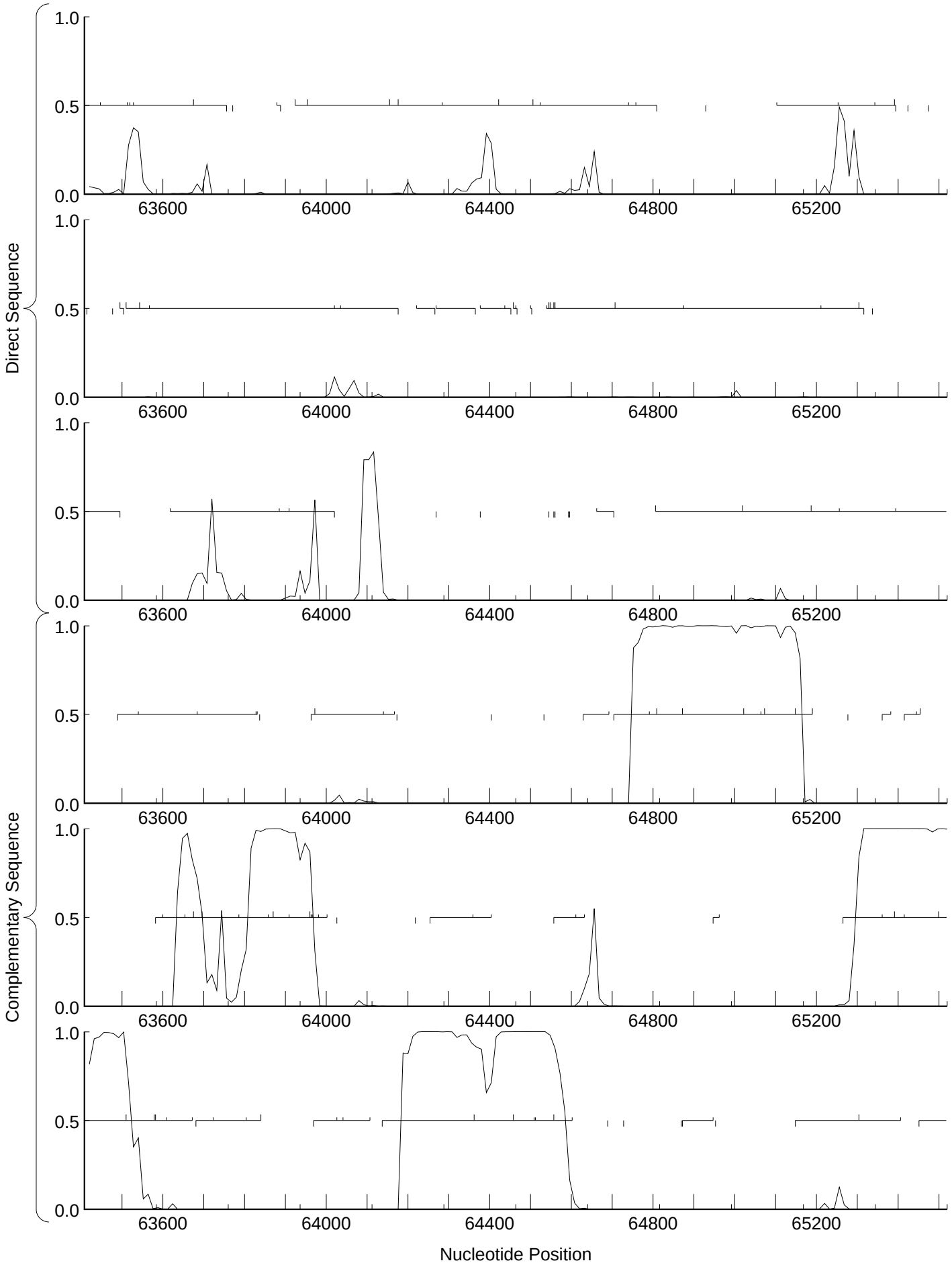

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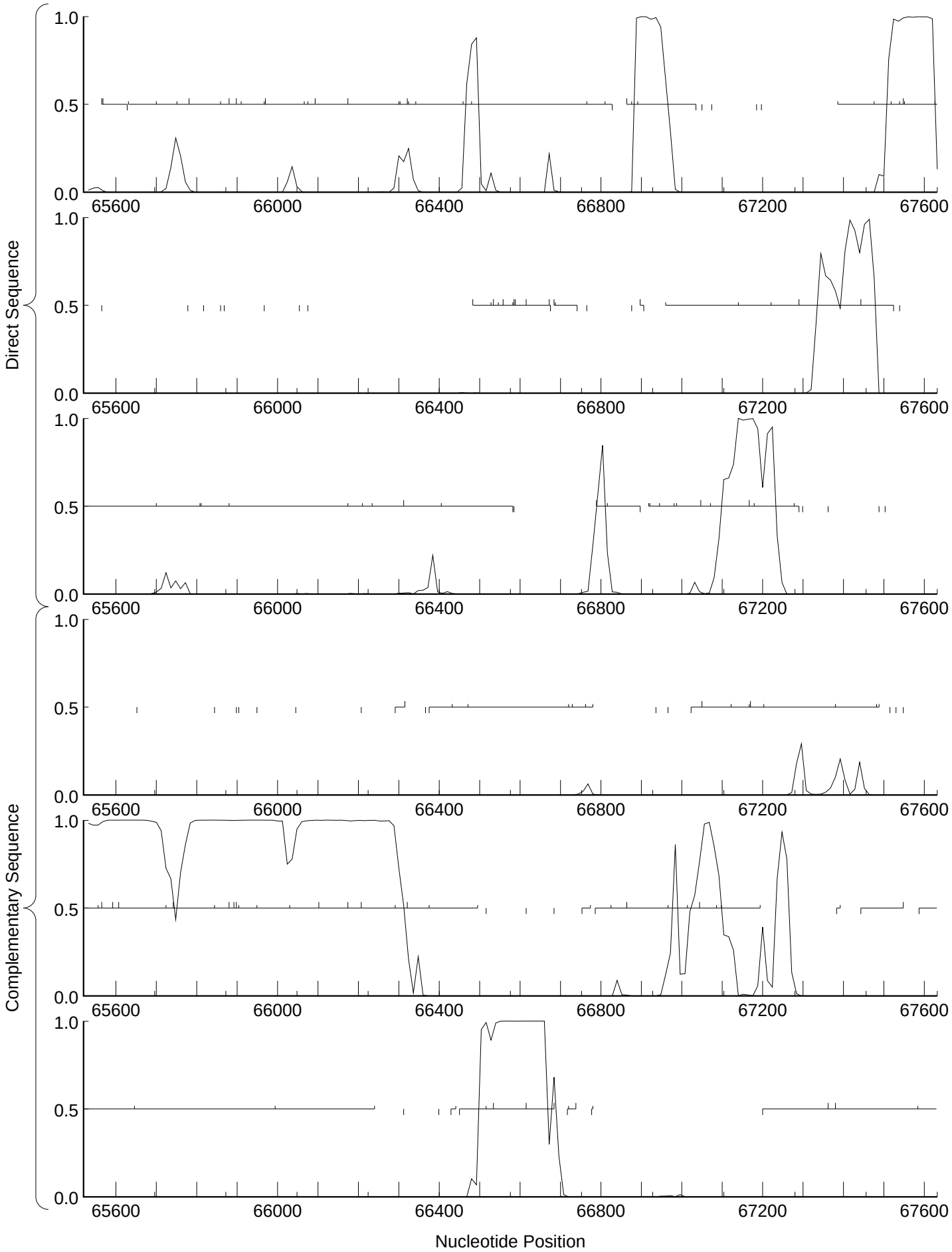

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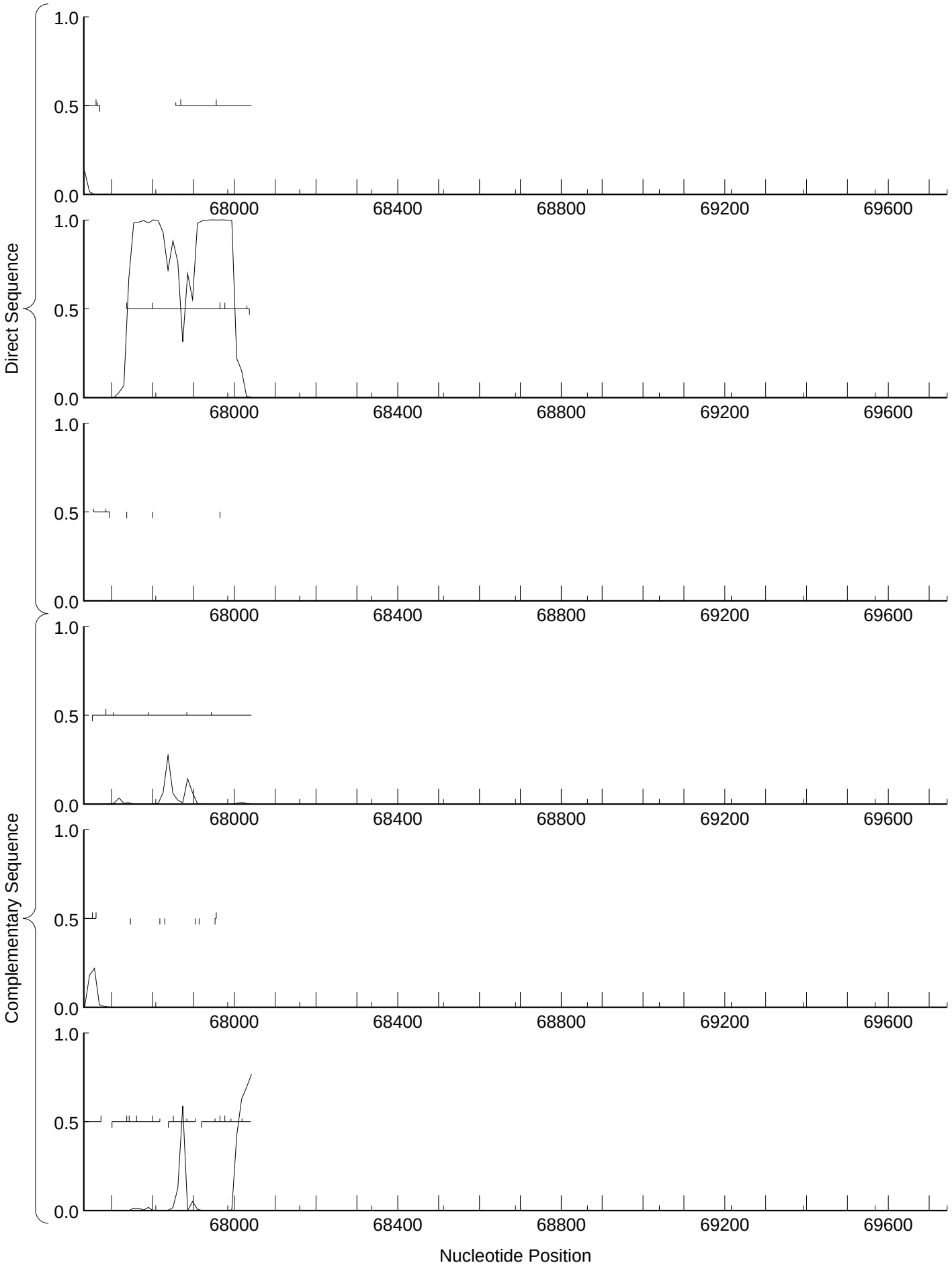

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Direct Sequence
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Direct Sequence
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63200
Nucleotide Position


Direct Sequence
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Direct Sequence
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Direct Sequence
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