GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Hammer
Analysis Date    :  3/28/24 at 20:45:56
Pages            :  26
Sequence Length  :  51889 bp
GC Content       :   61.33%

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

PS-Version       :  1.2

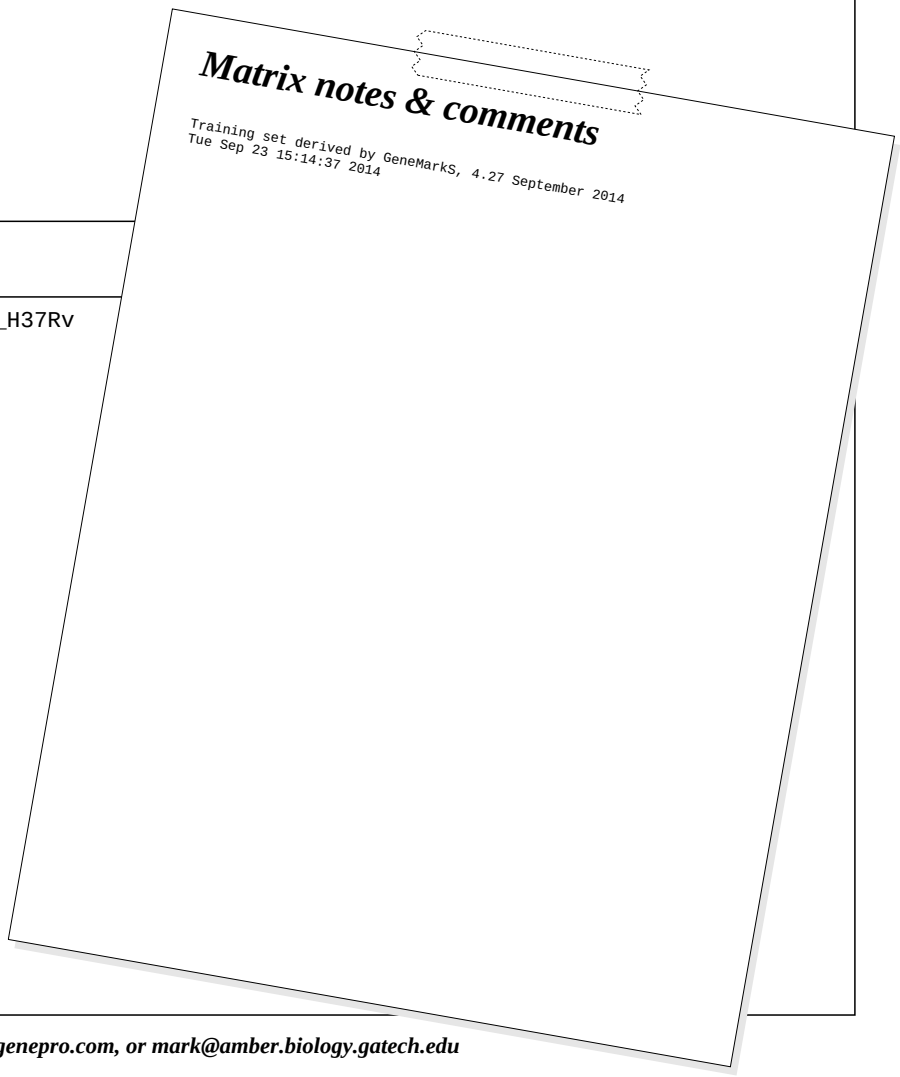
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
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MATRIX INFORMATION

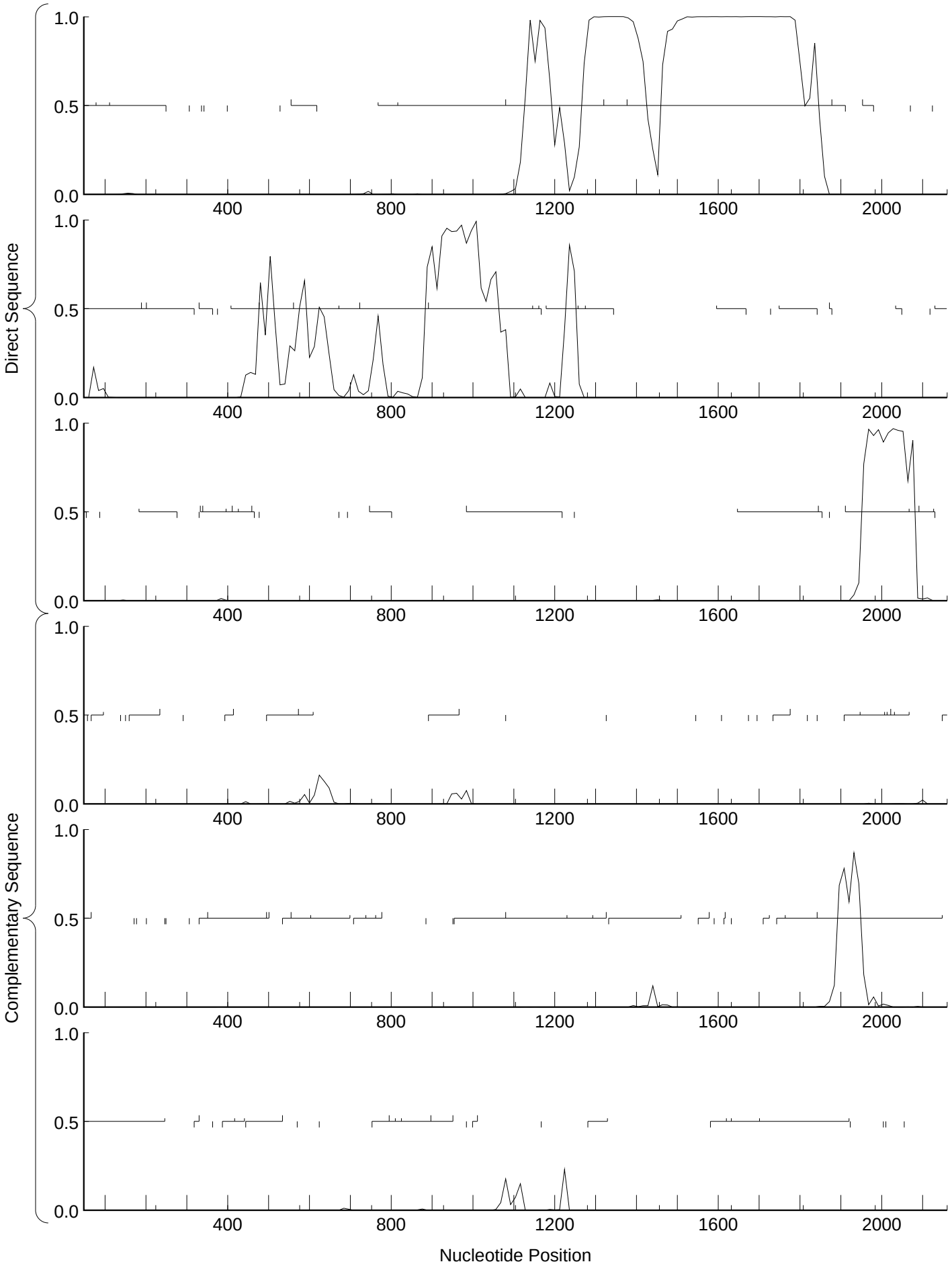
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Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

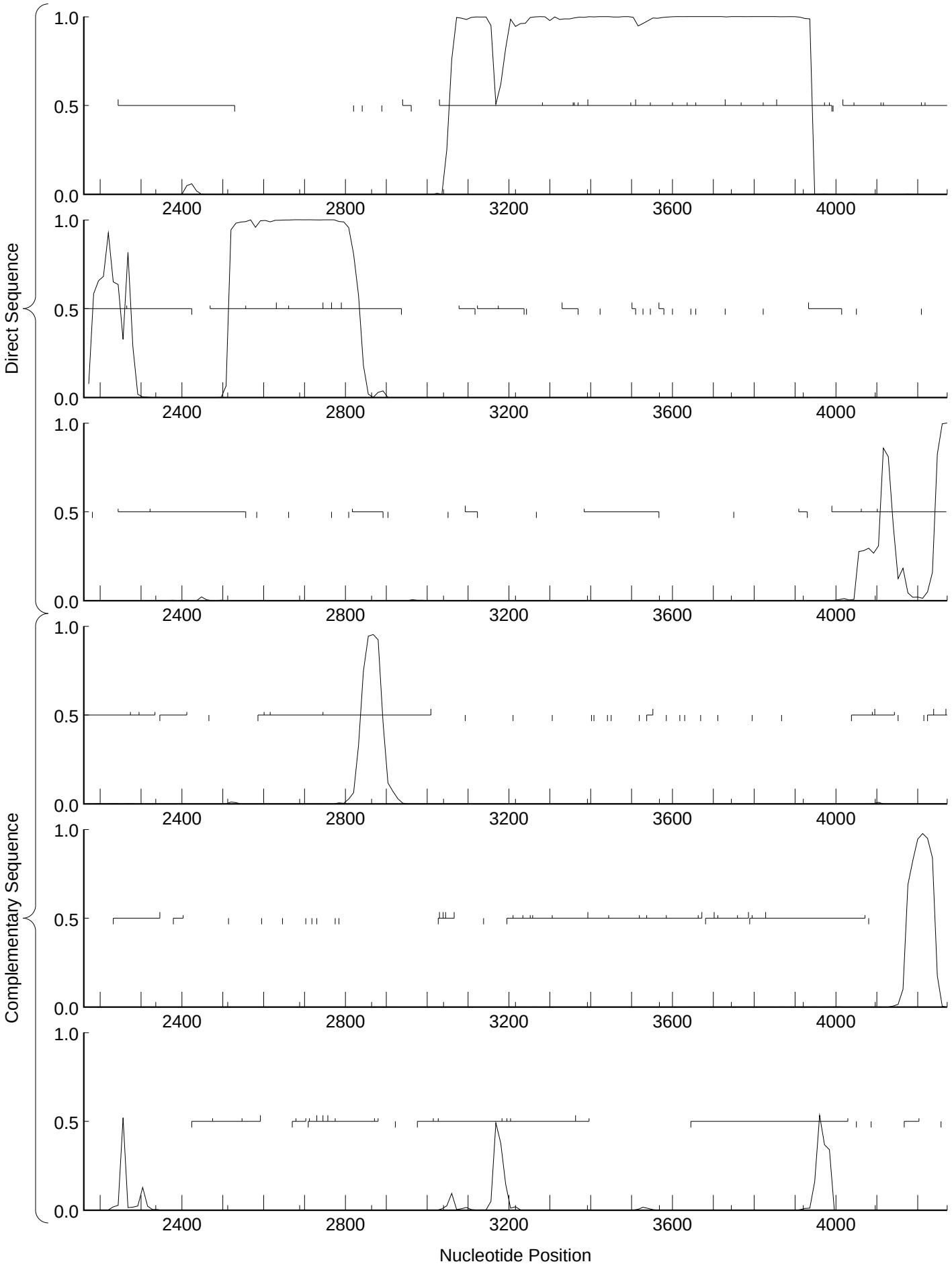
Order      :  4

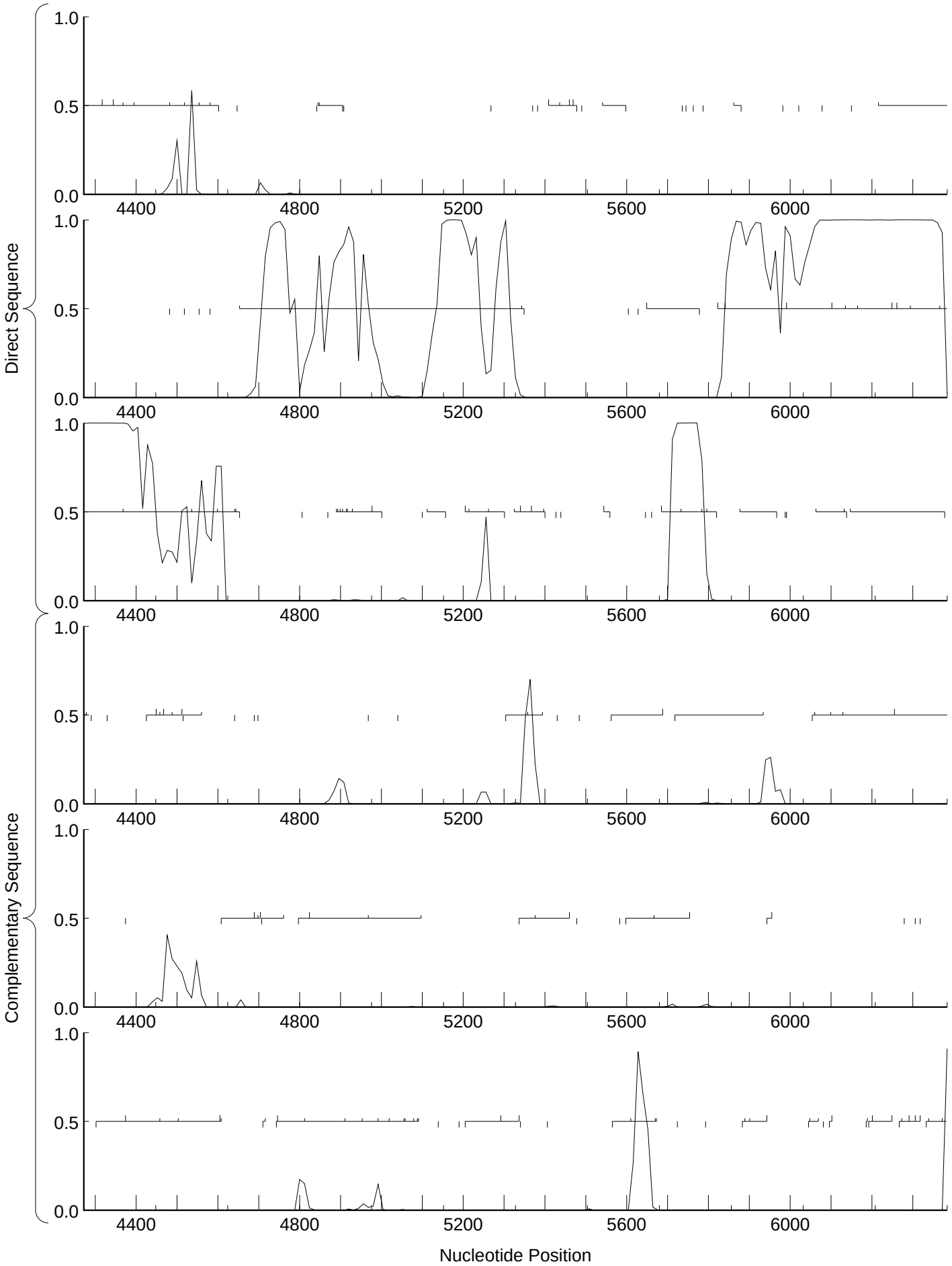

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

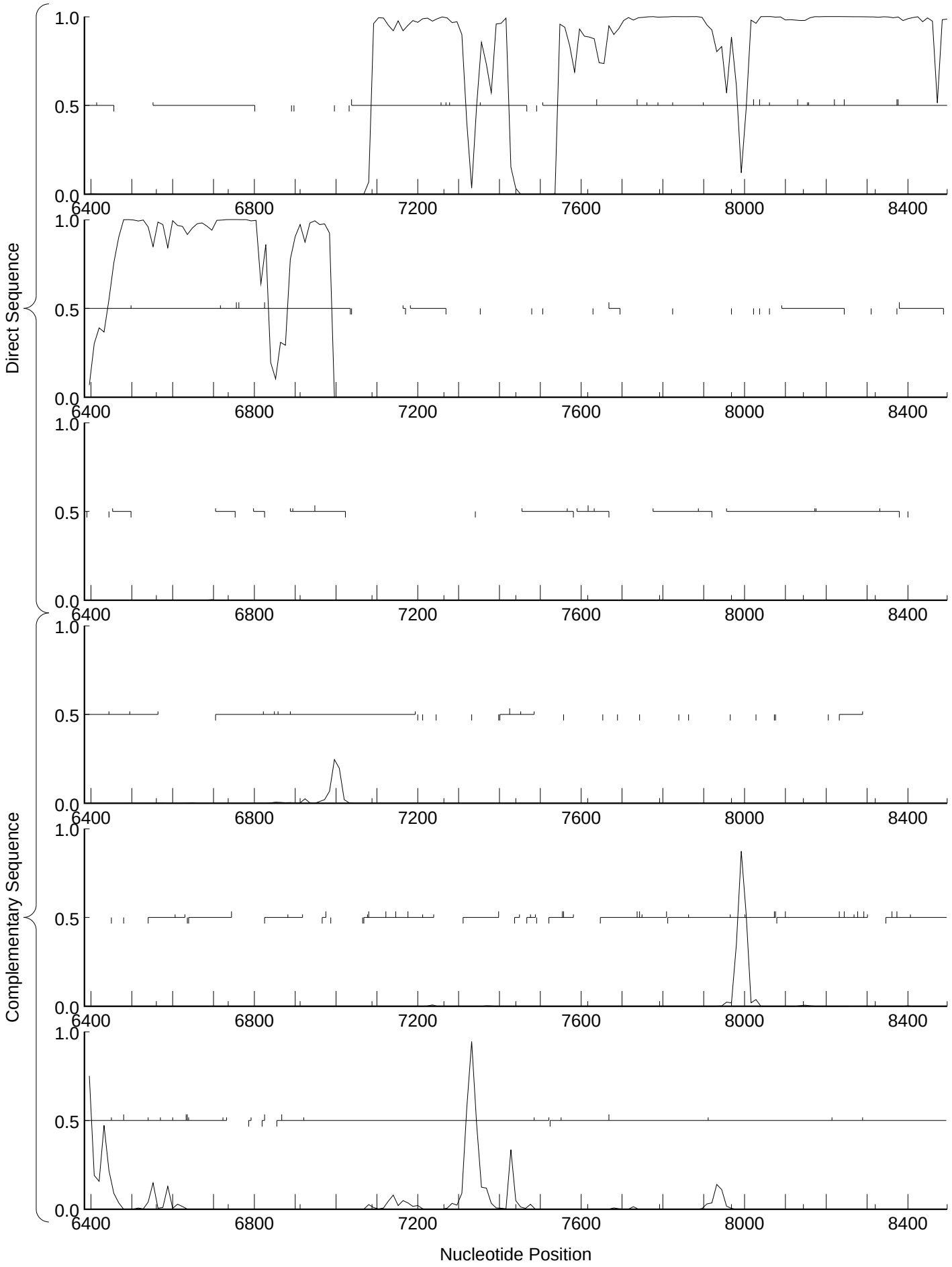


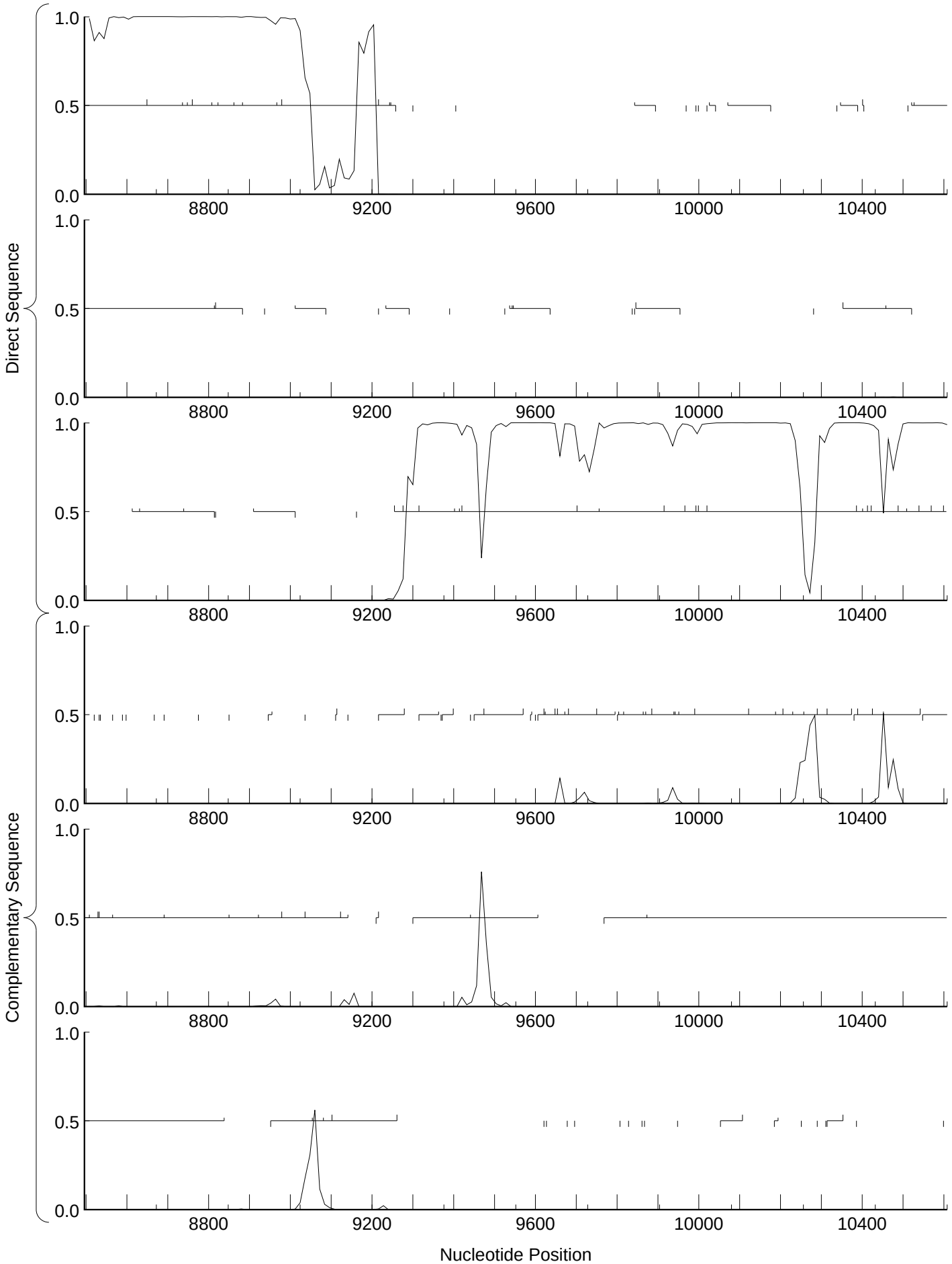
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

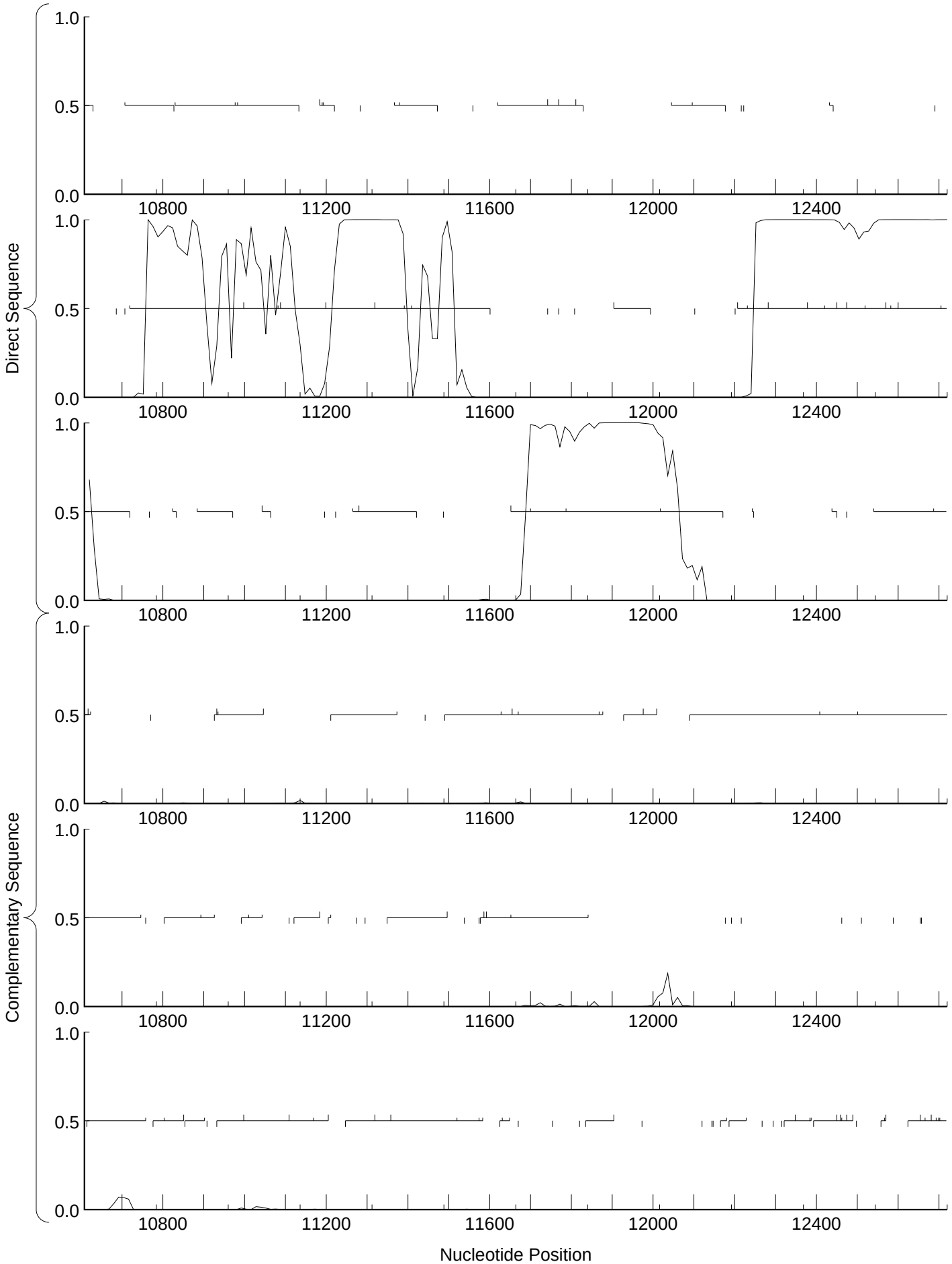

Direct Sequence
Complementary Sequence
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800
1200
1600
2000
Nucleotide Position

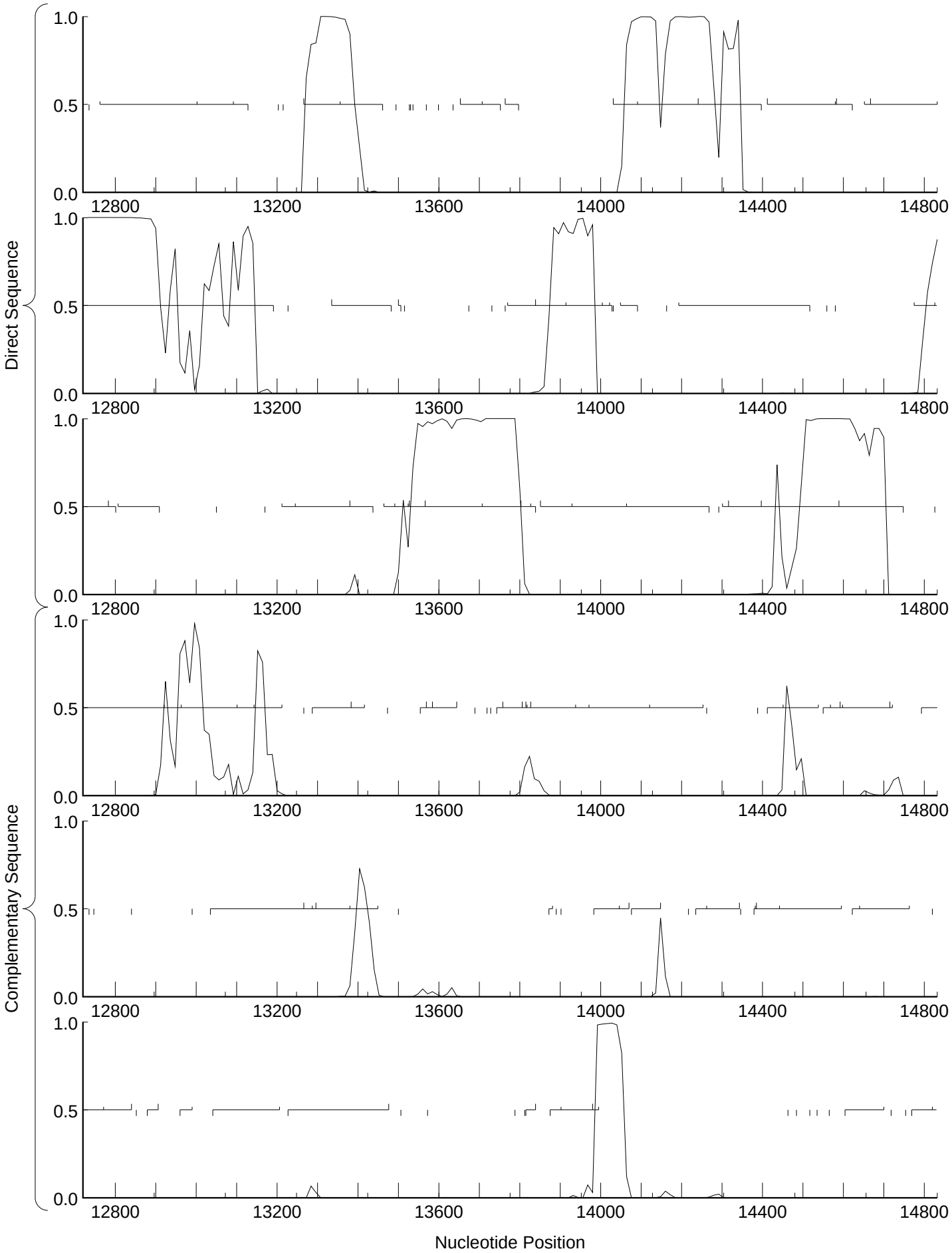

Direct Sequence
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Nucleotide Position

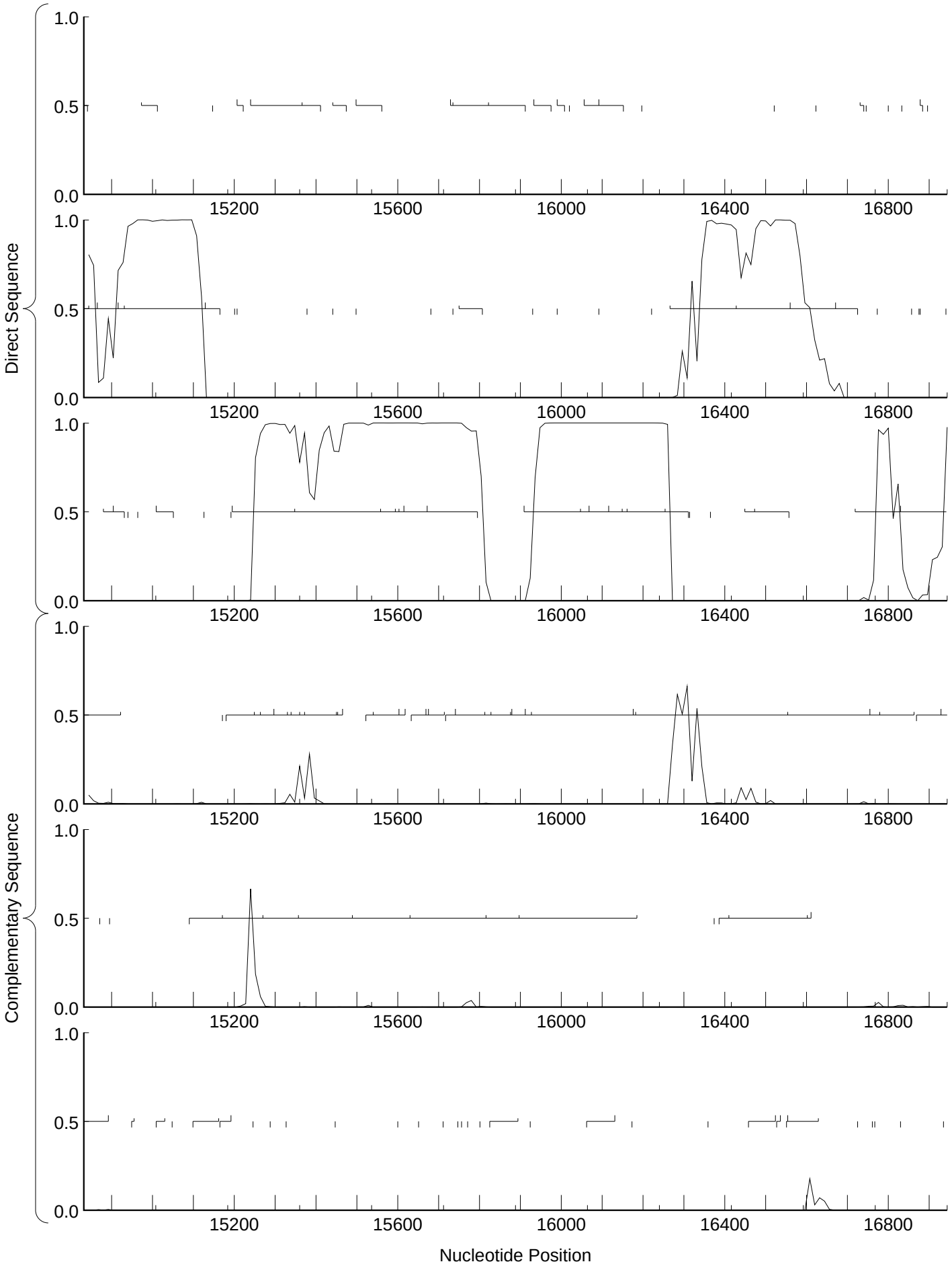

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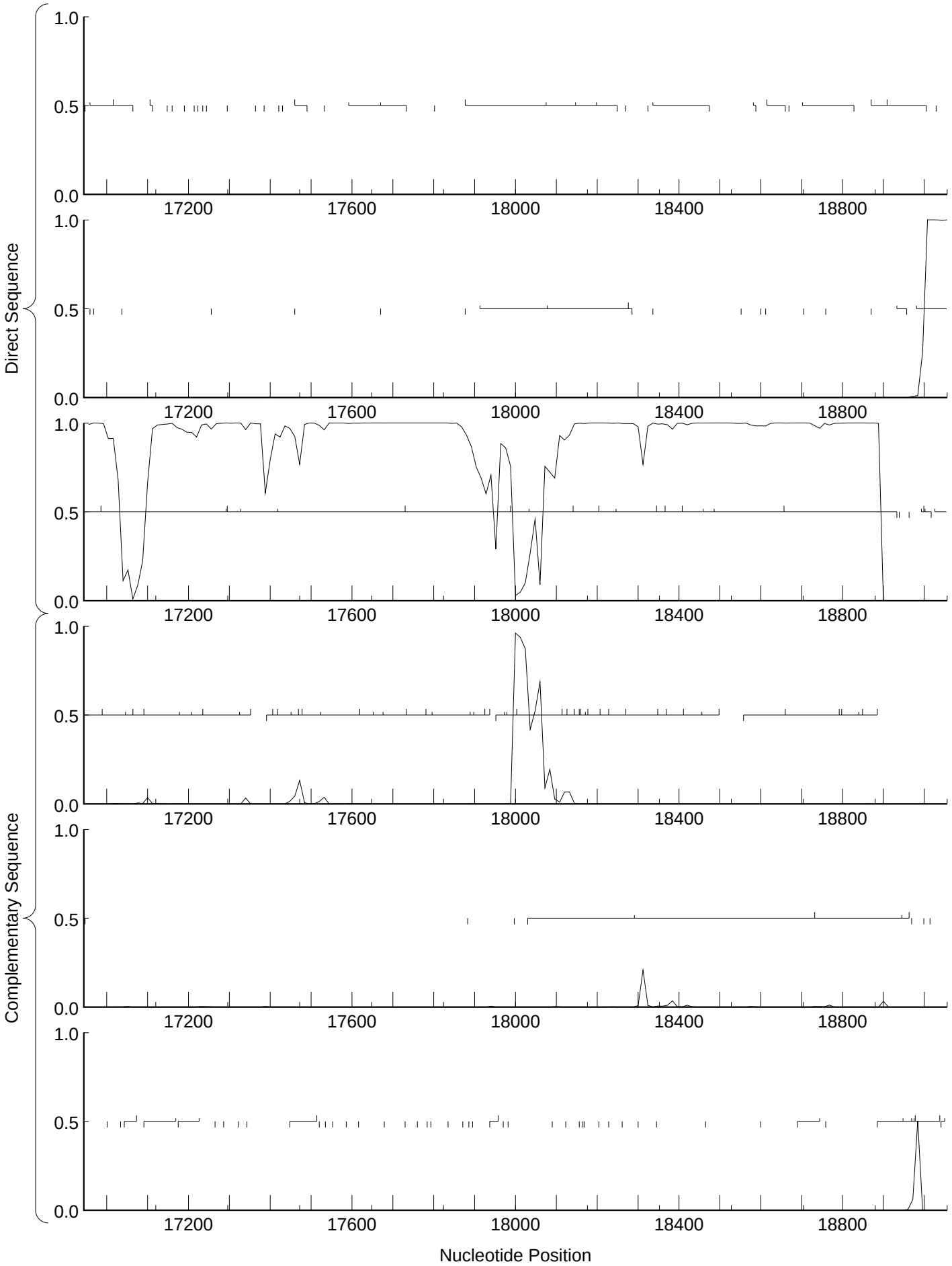

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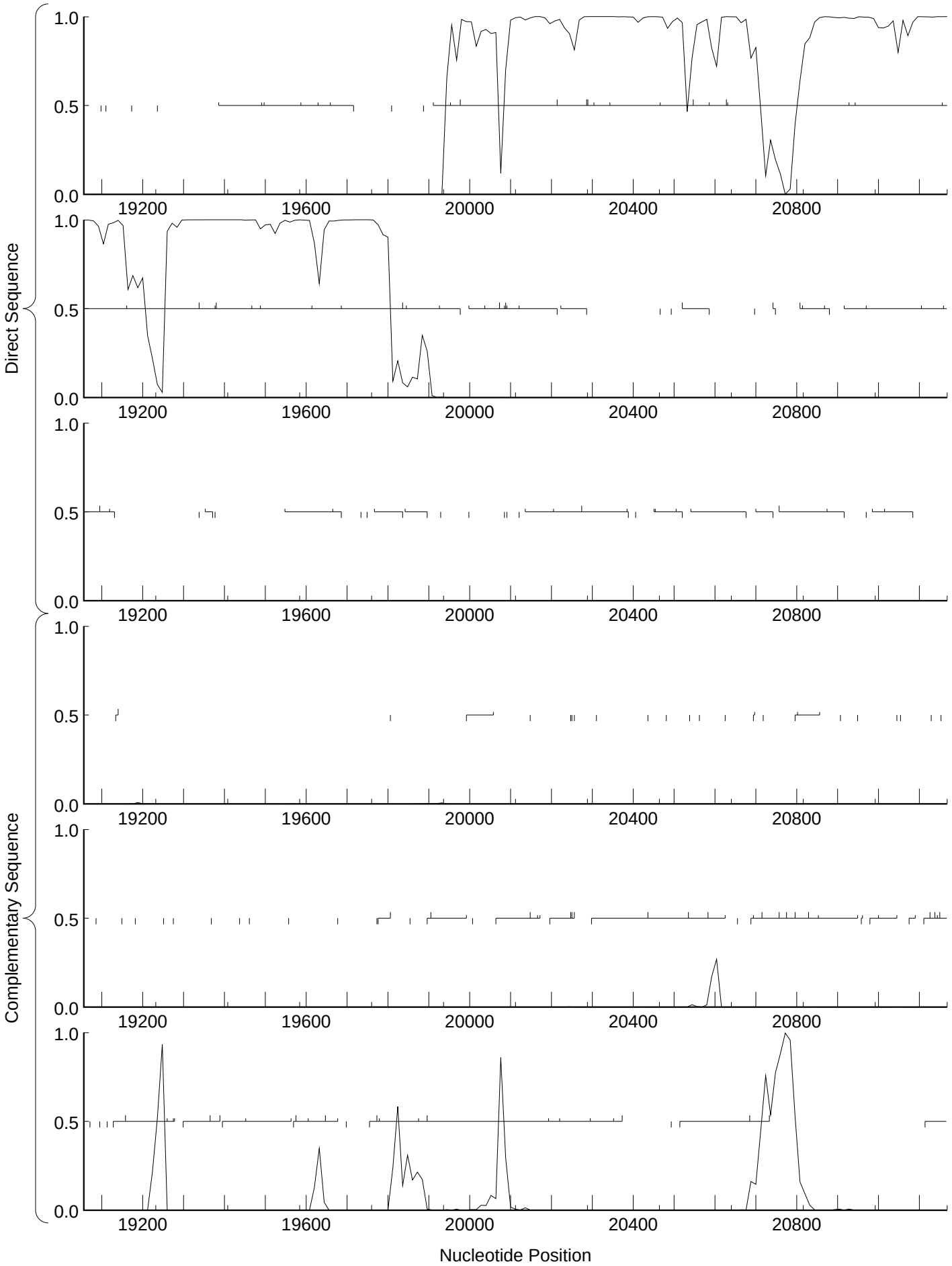

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Nucleotide Position

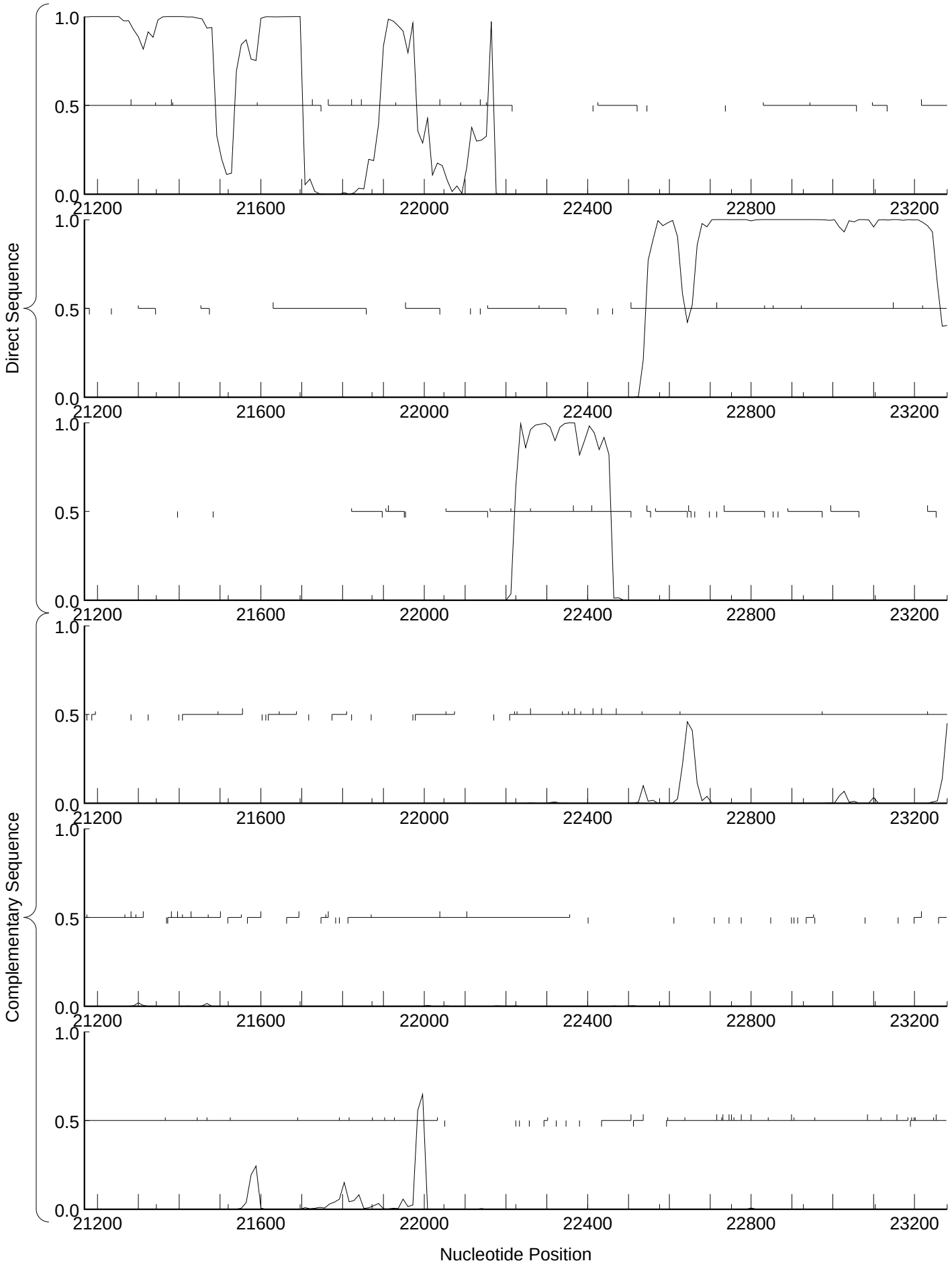

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Nucleotide Position

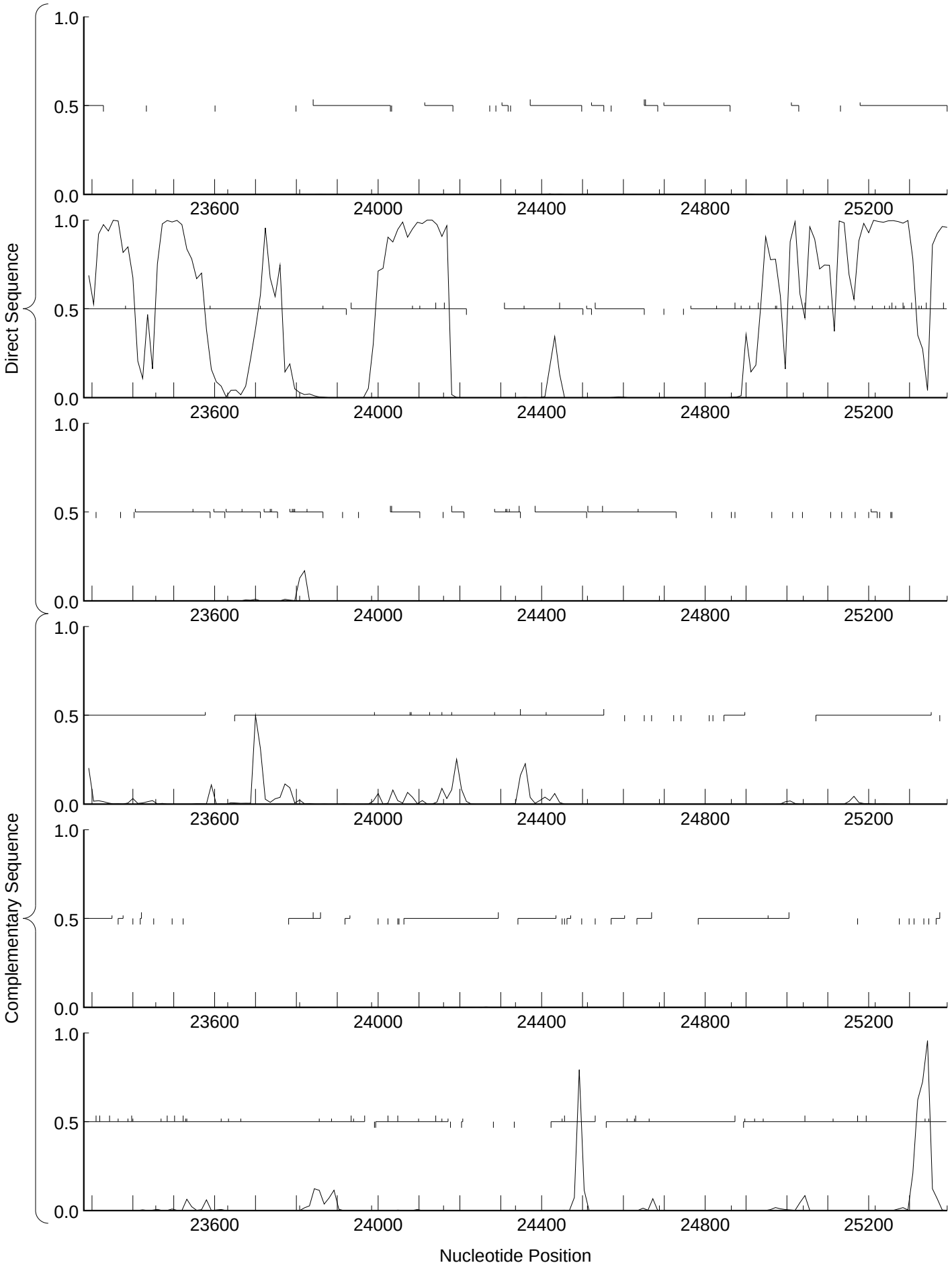

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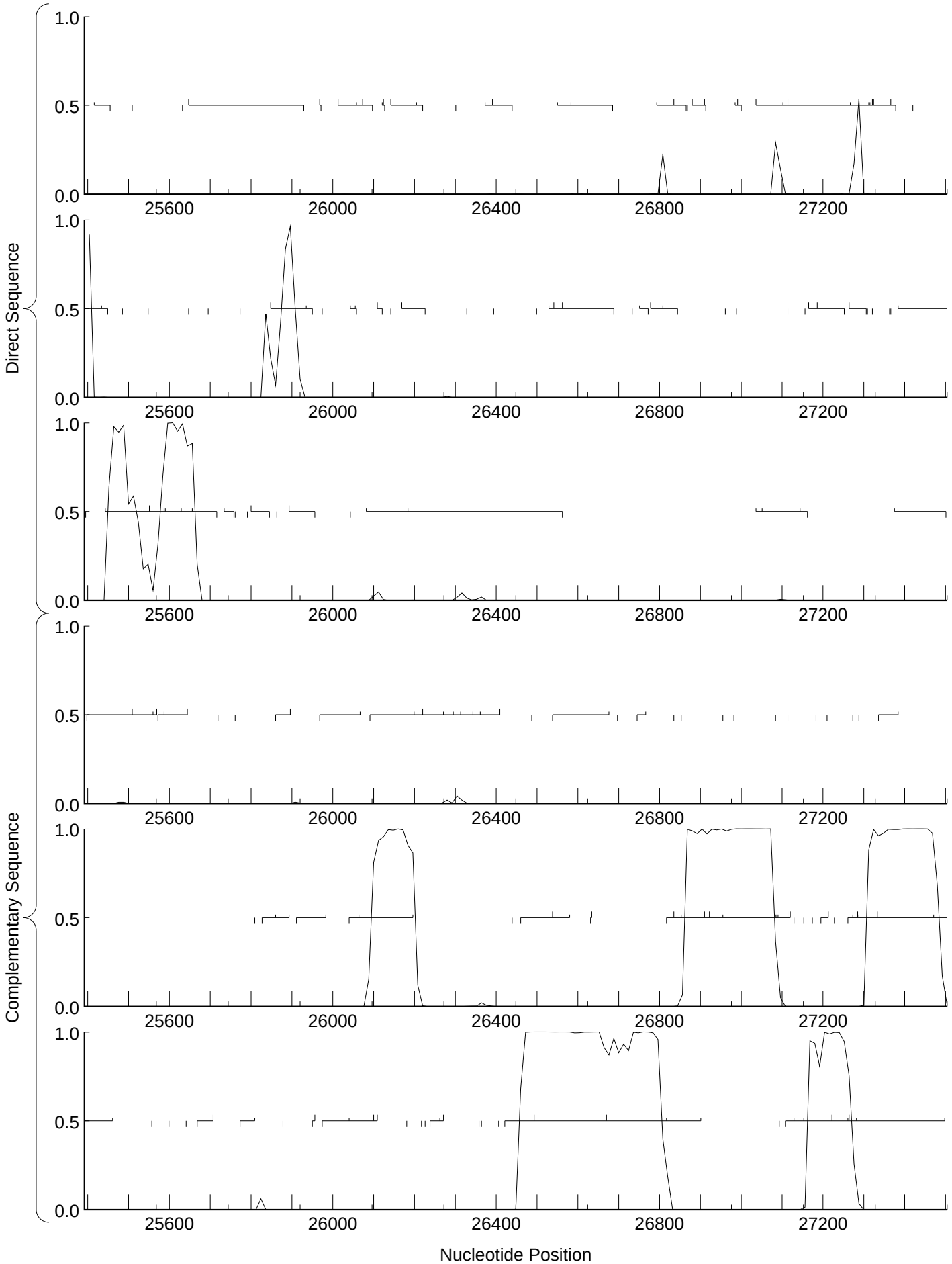

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Nucleotide Position

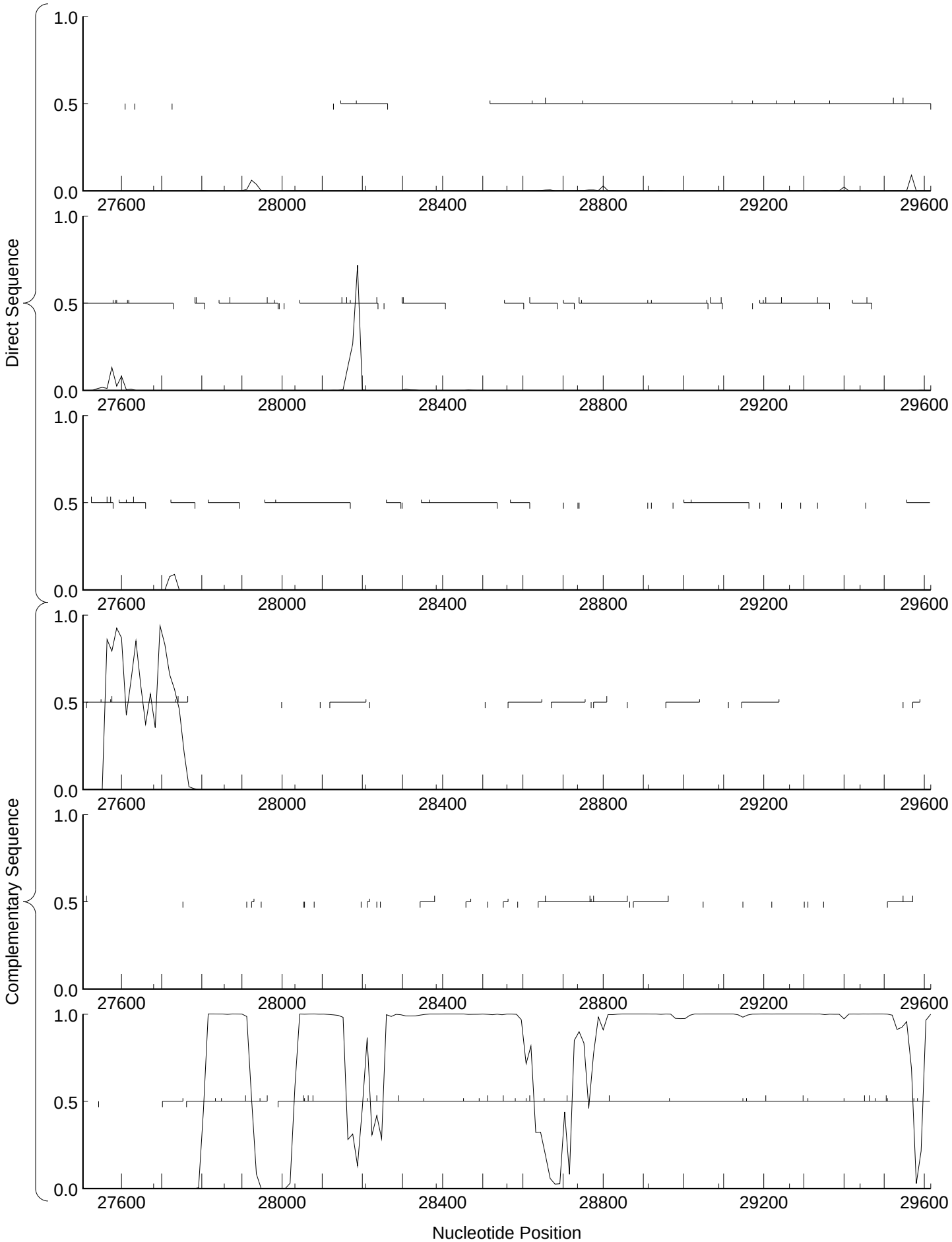

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Nucleotide Position

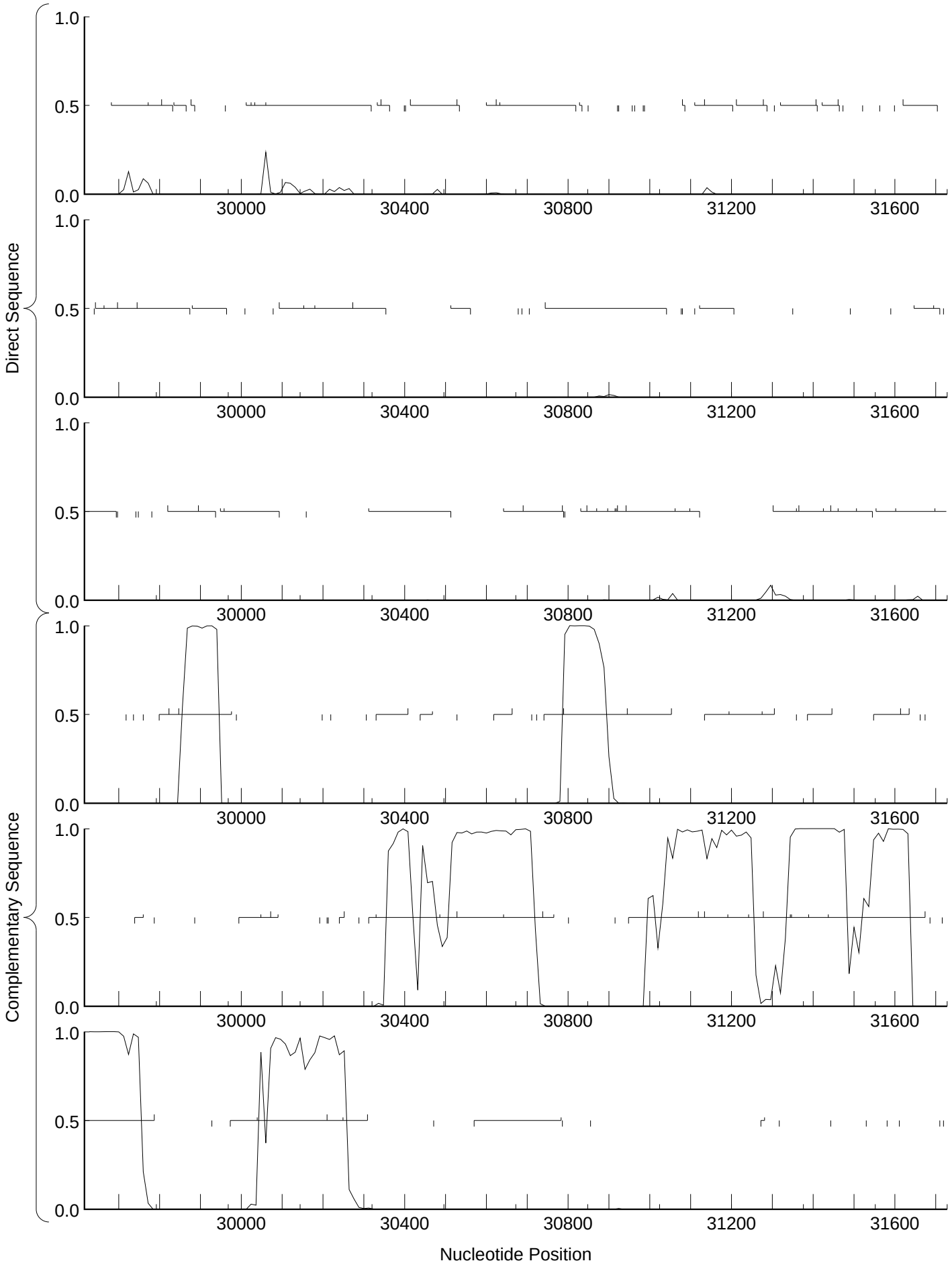

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Nucleotide Position

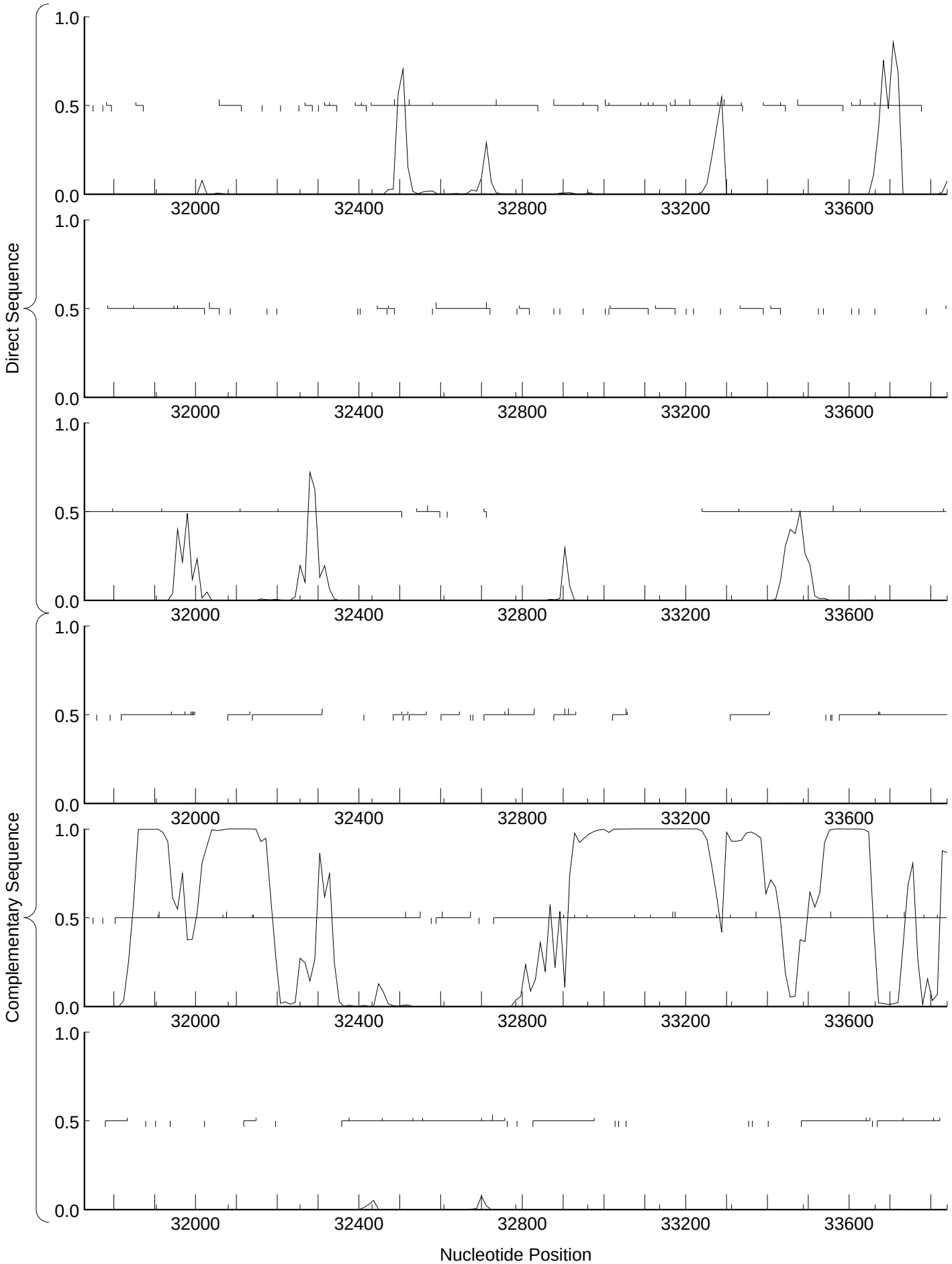

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Complementary Sequence
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Nucleotide Position

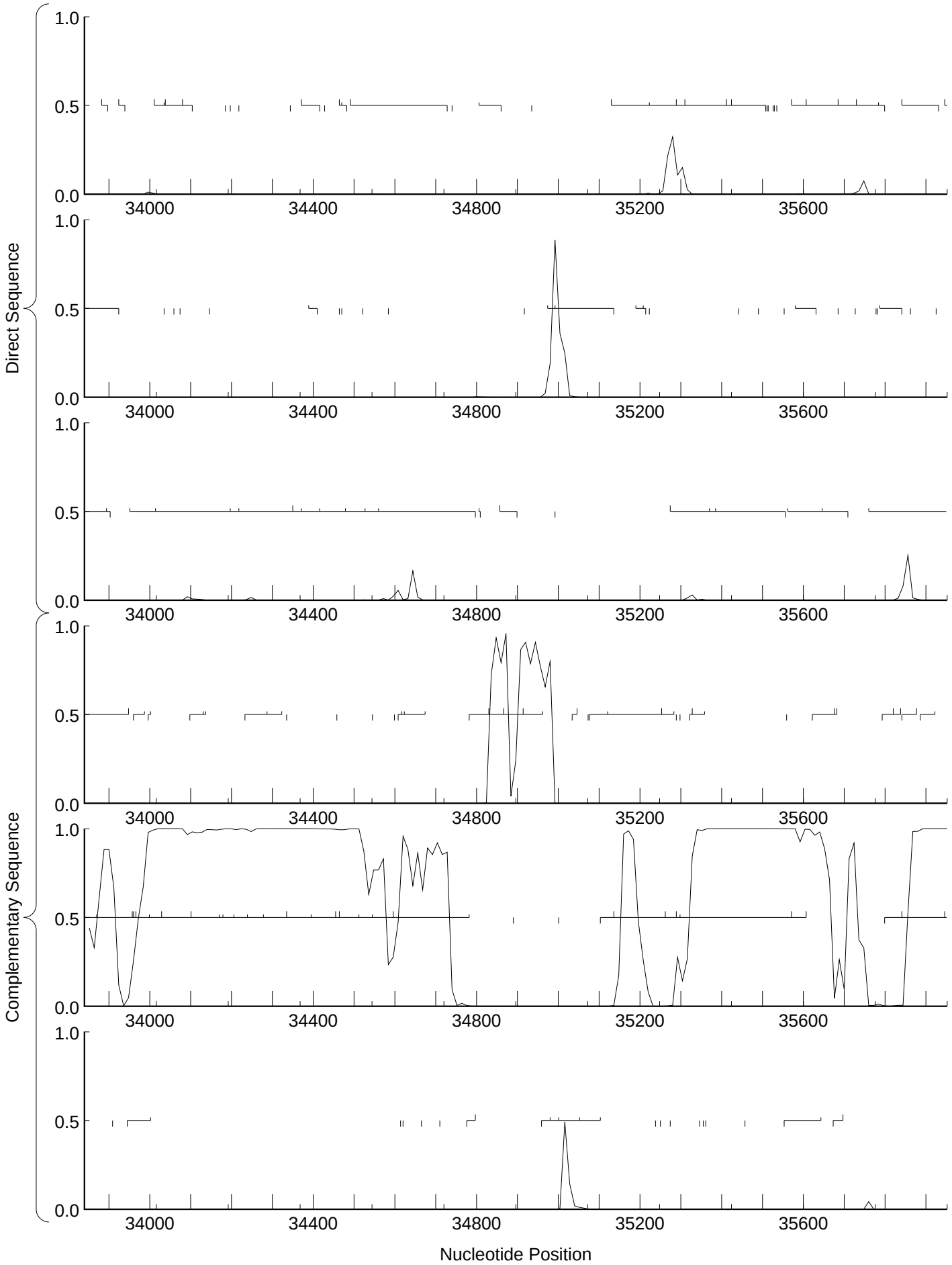

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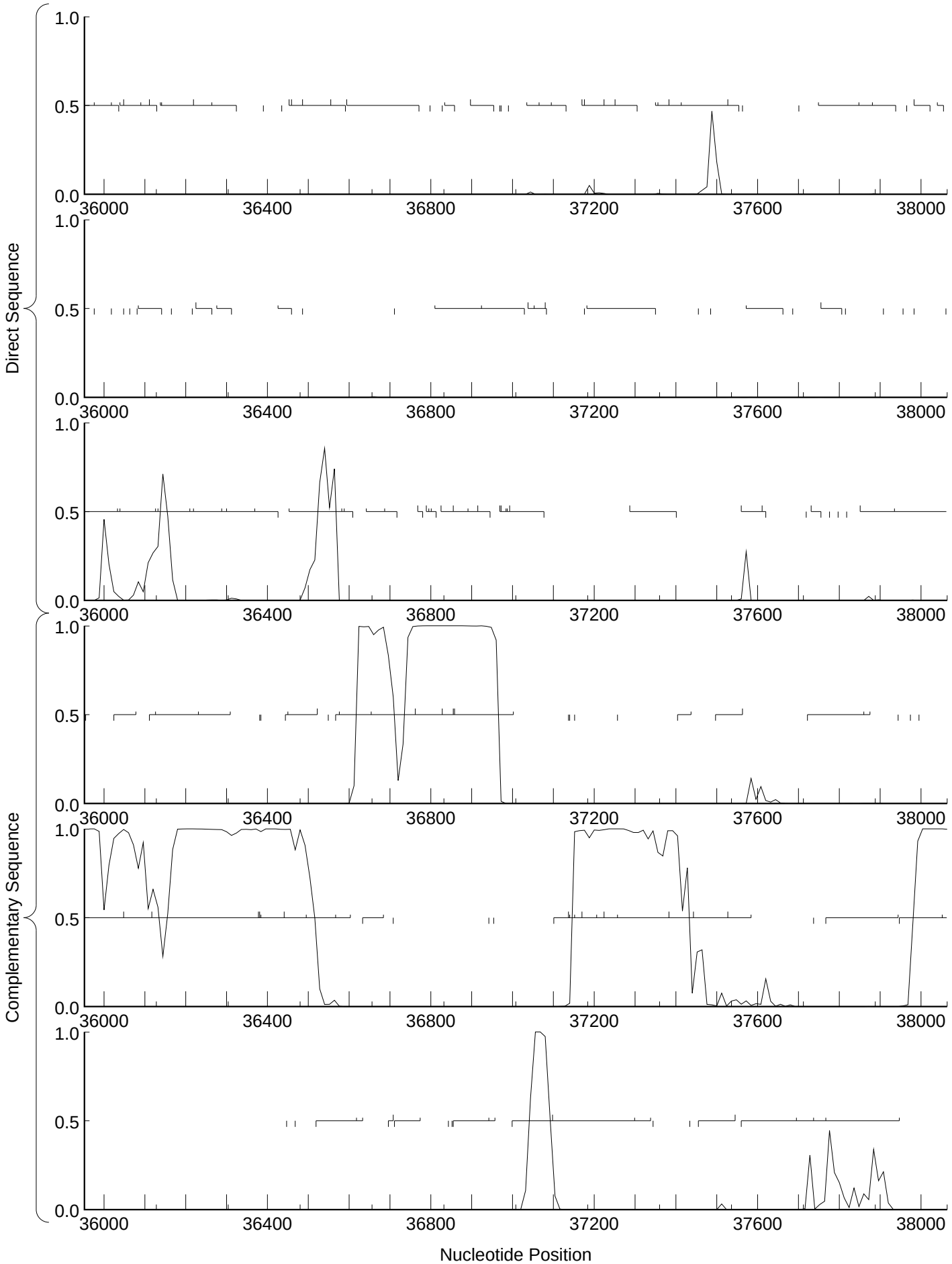

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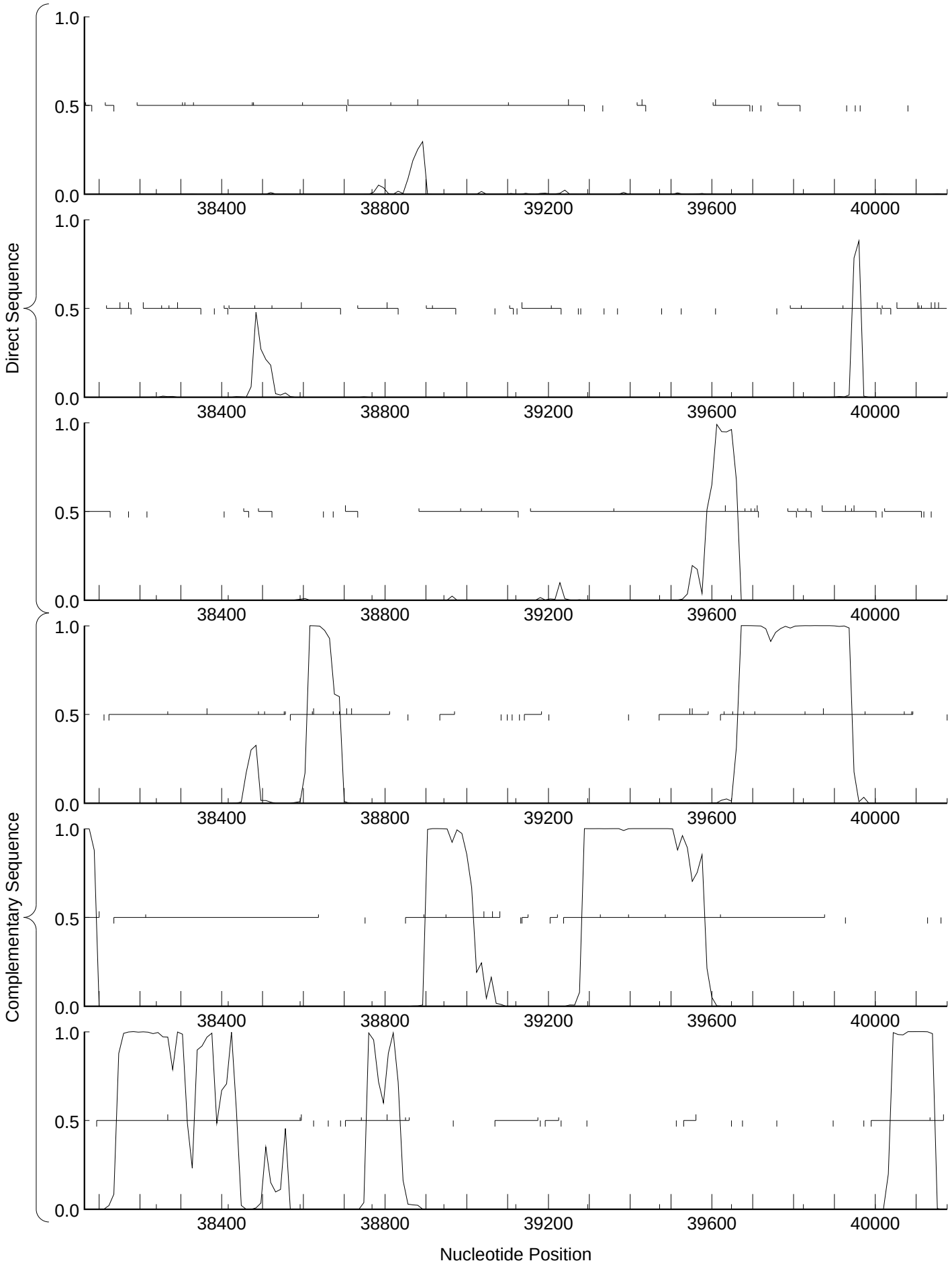

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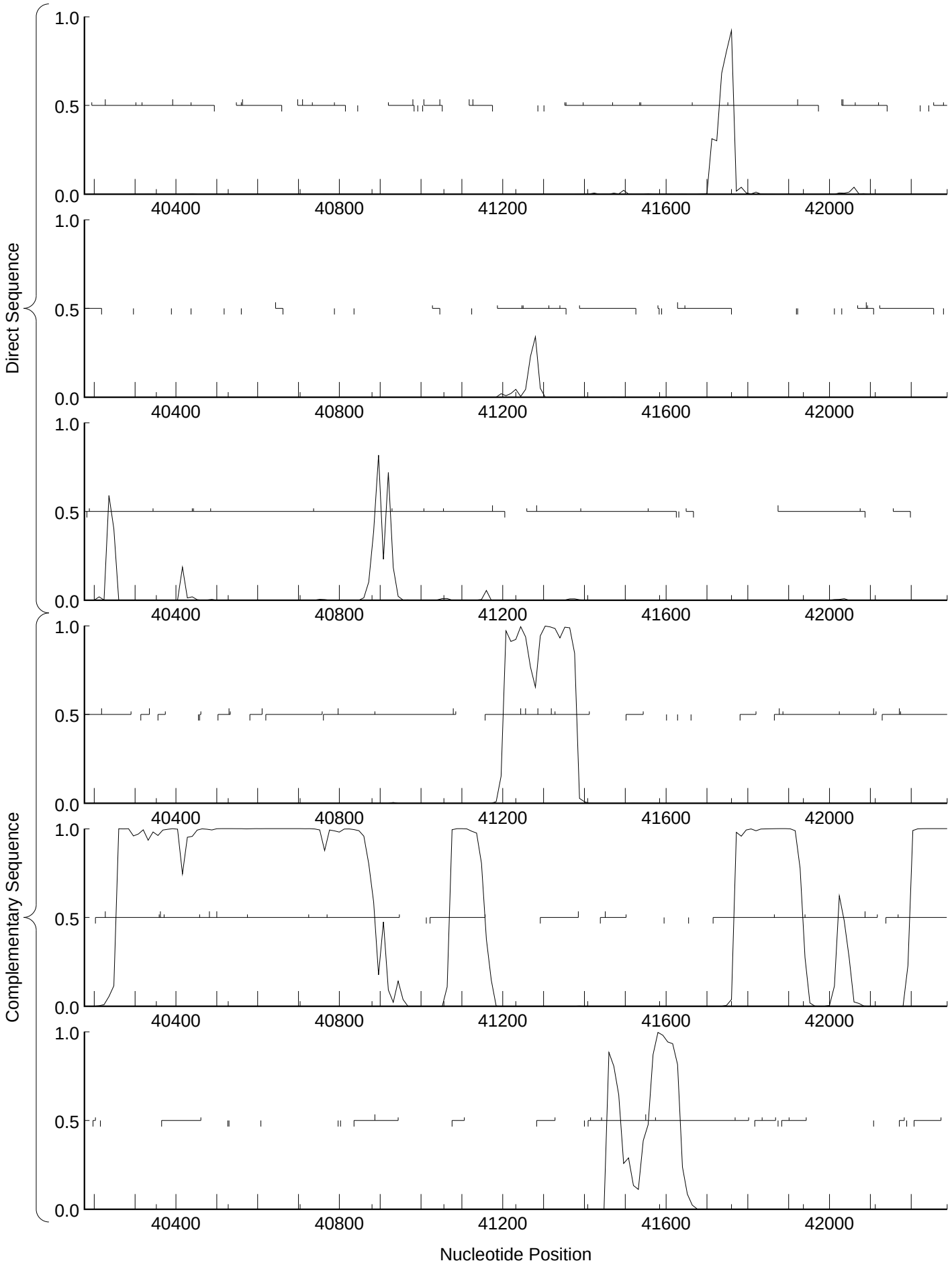

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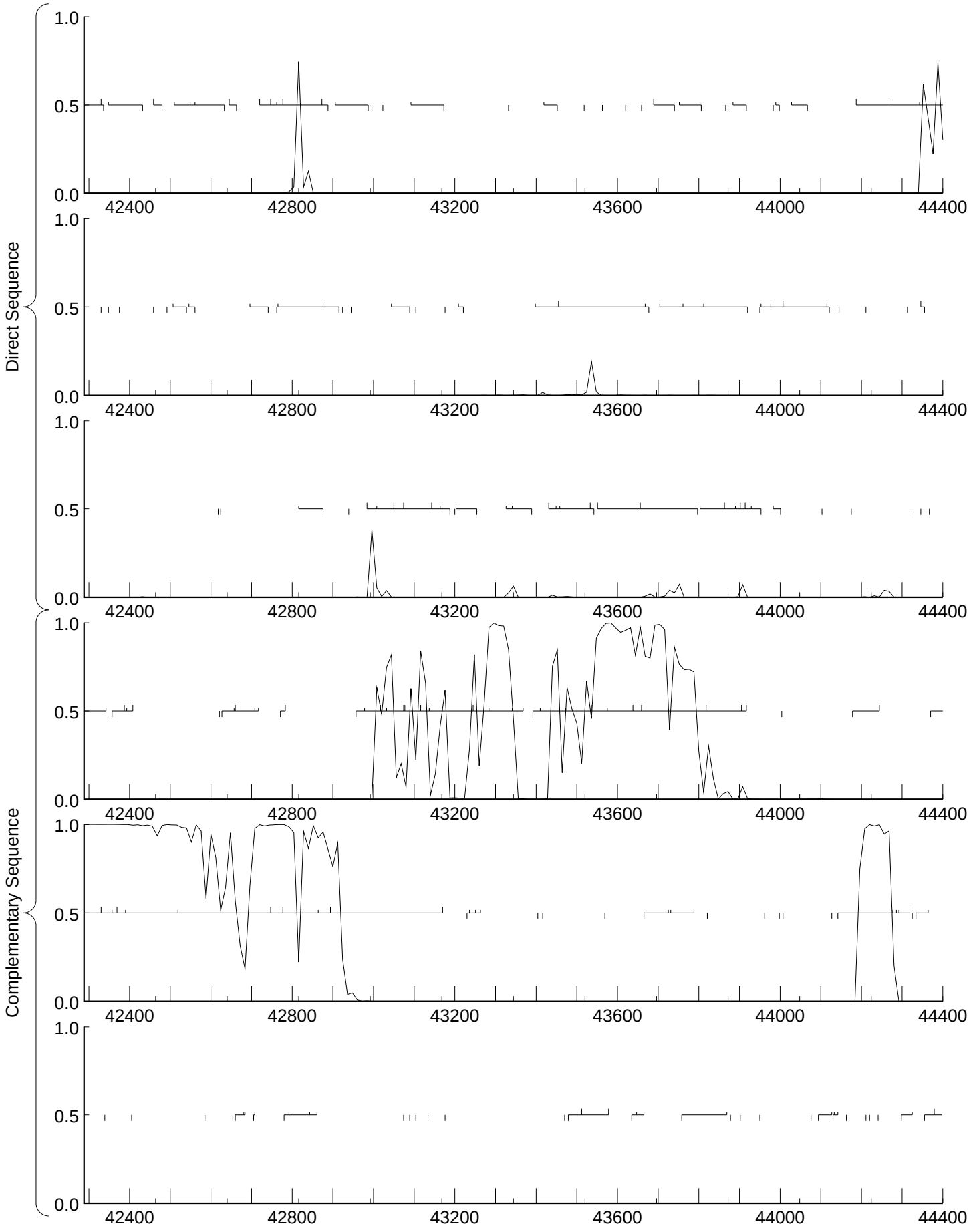

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Nucleotide Position

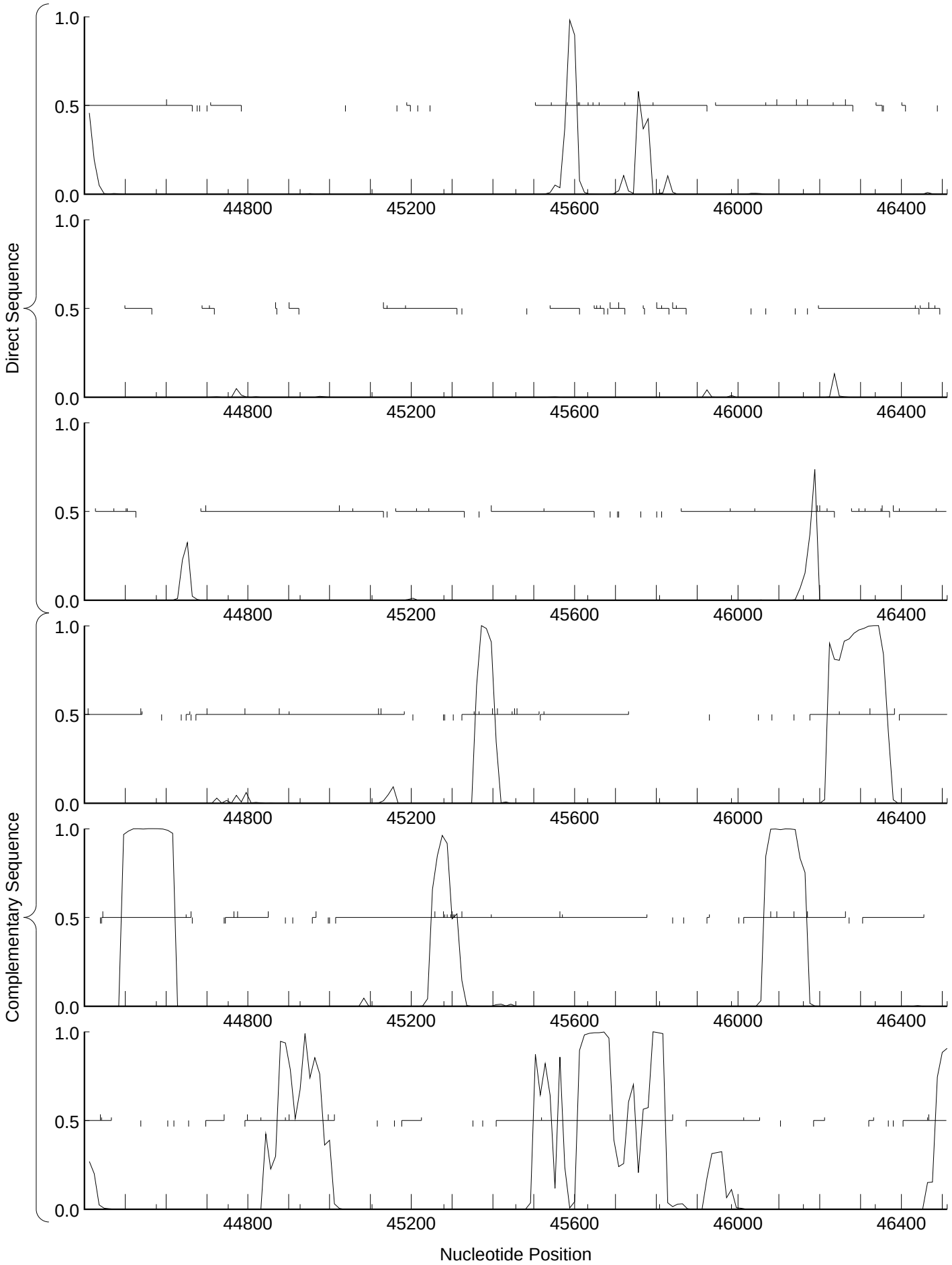

Direct Sequence
Complementary Sequence
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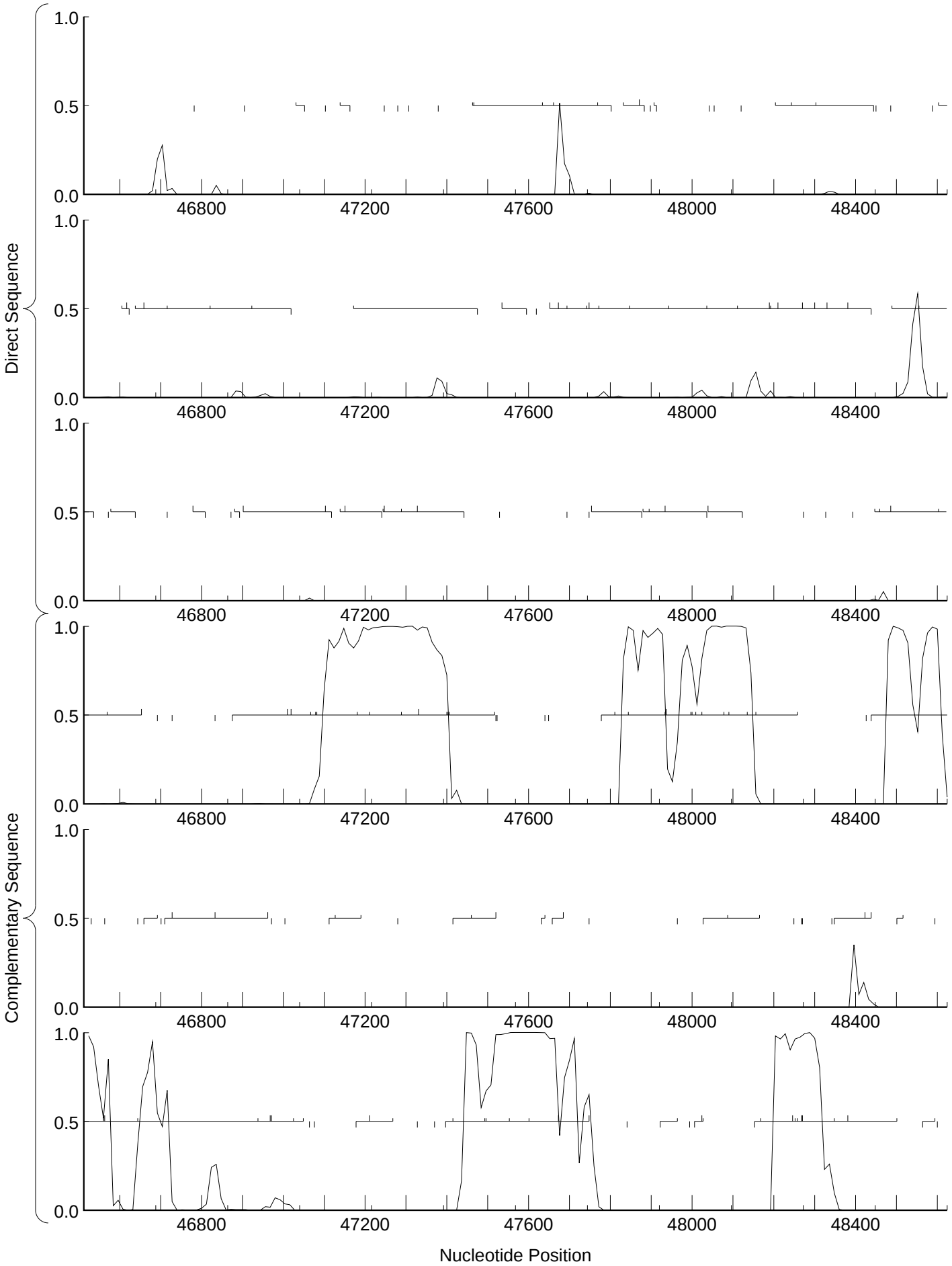

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Nucleotide Position

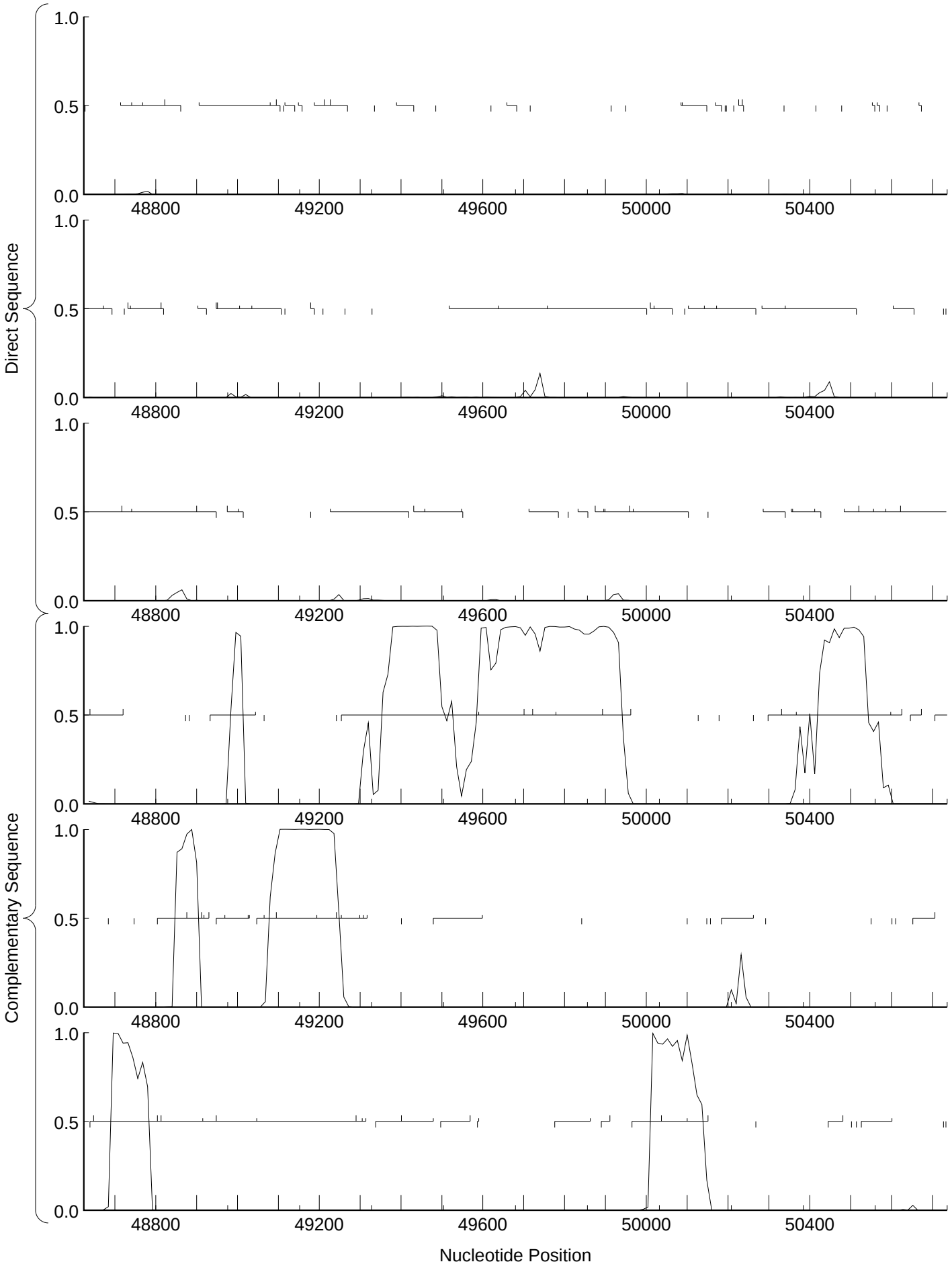

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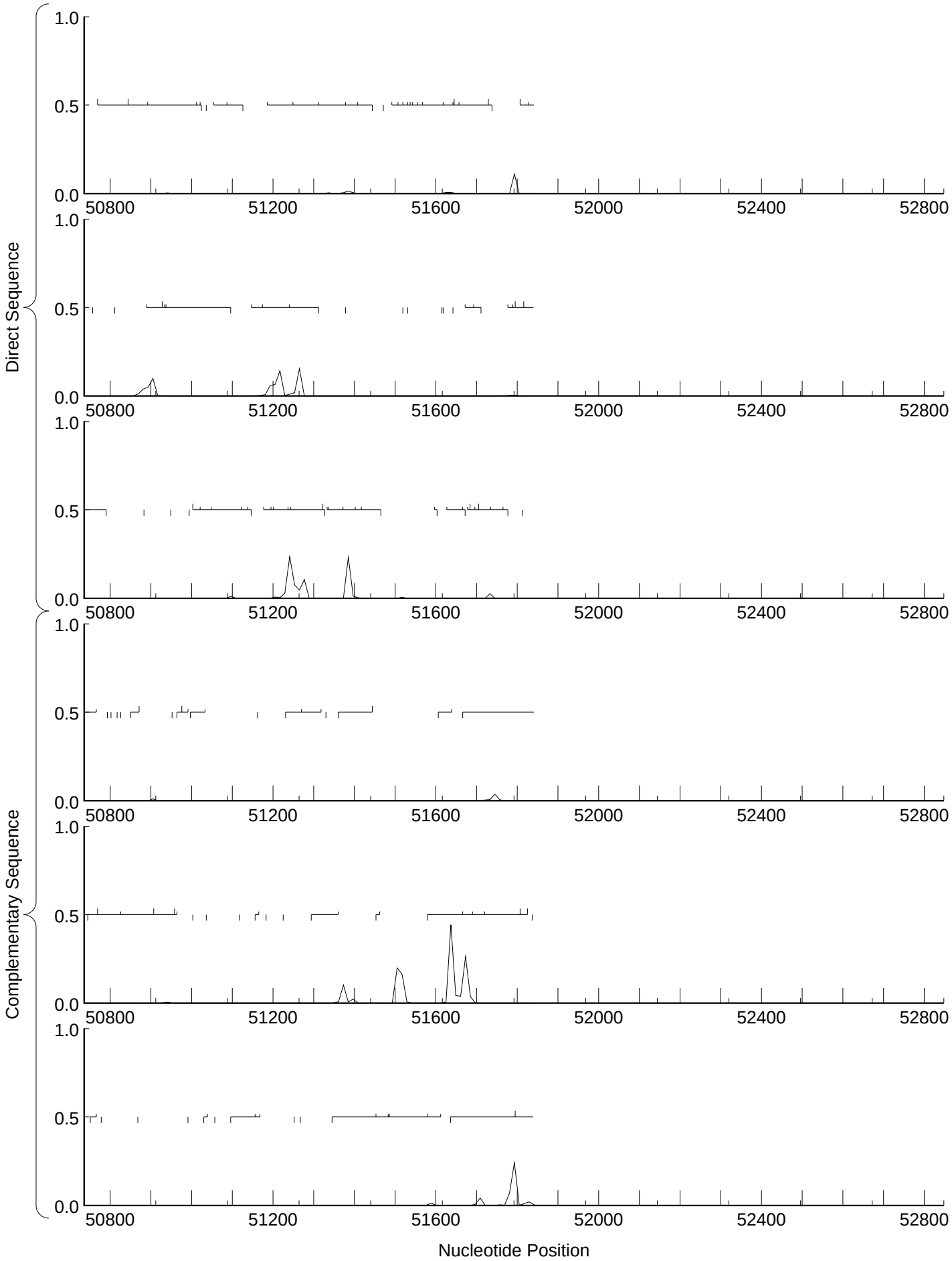

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Direct Sequence
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44000
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Nucleotide Position


Direct Sequence
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Direct Sequence
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