GeneMark

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

PROGRAM INFORMATION

```
Sequence          :  Mycobacterium phage KingVeveve complete sequence, 68043 bp, circularly permuted, Cluster B1
Analysis Date     :  3/28/24 at 9:14:20
Pages             :  34
Sequence Length   :  68043 bp
GC Content        :   66.50%

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

PS-Version        :  1.2

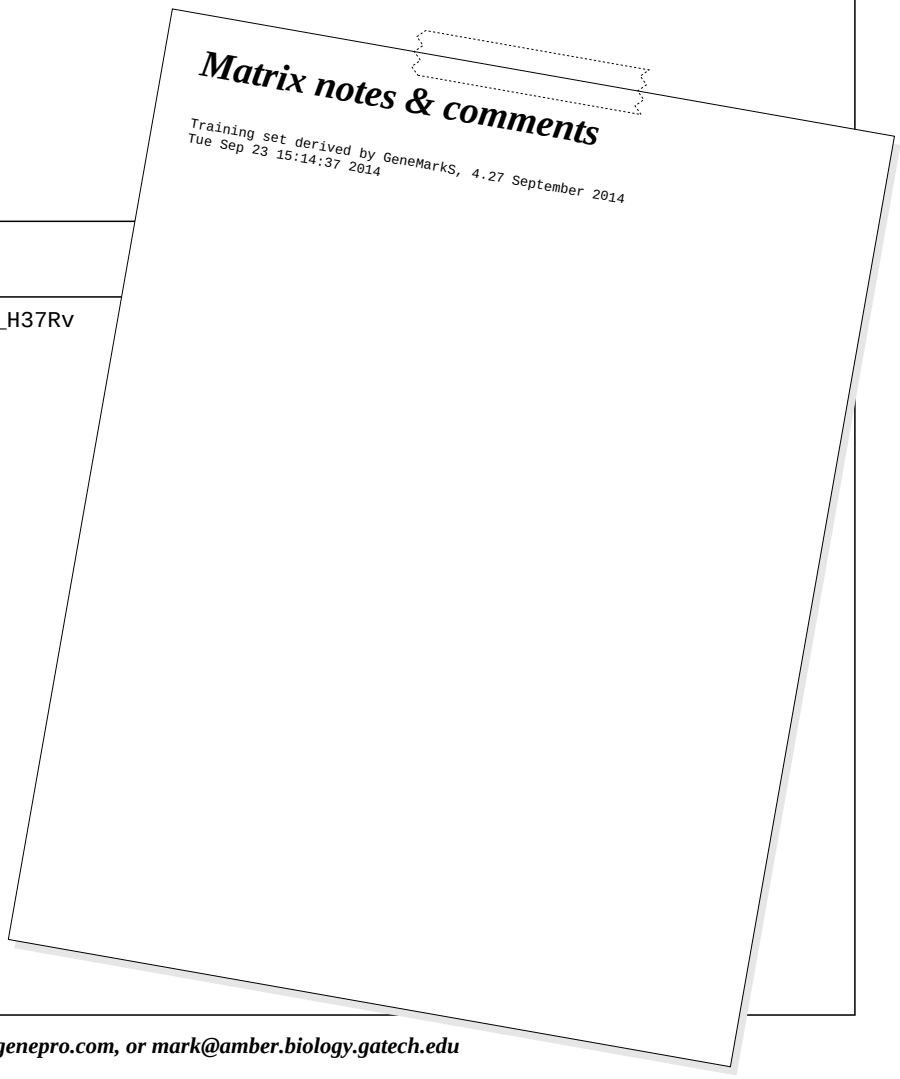
GeneMark Options  :  PostScript graph,
                     Mark ORFs / splice sites,
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MATRIX INFORMATION

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Matrix      :  Mycobacterium_tuberculosis_H37Rv
Author      :  -

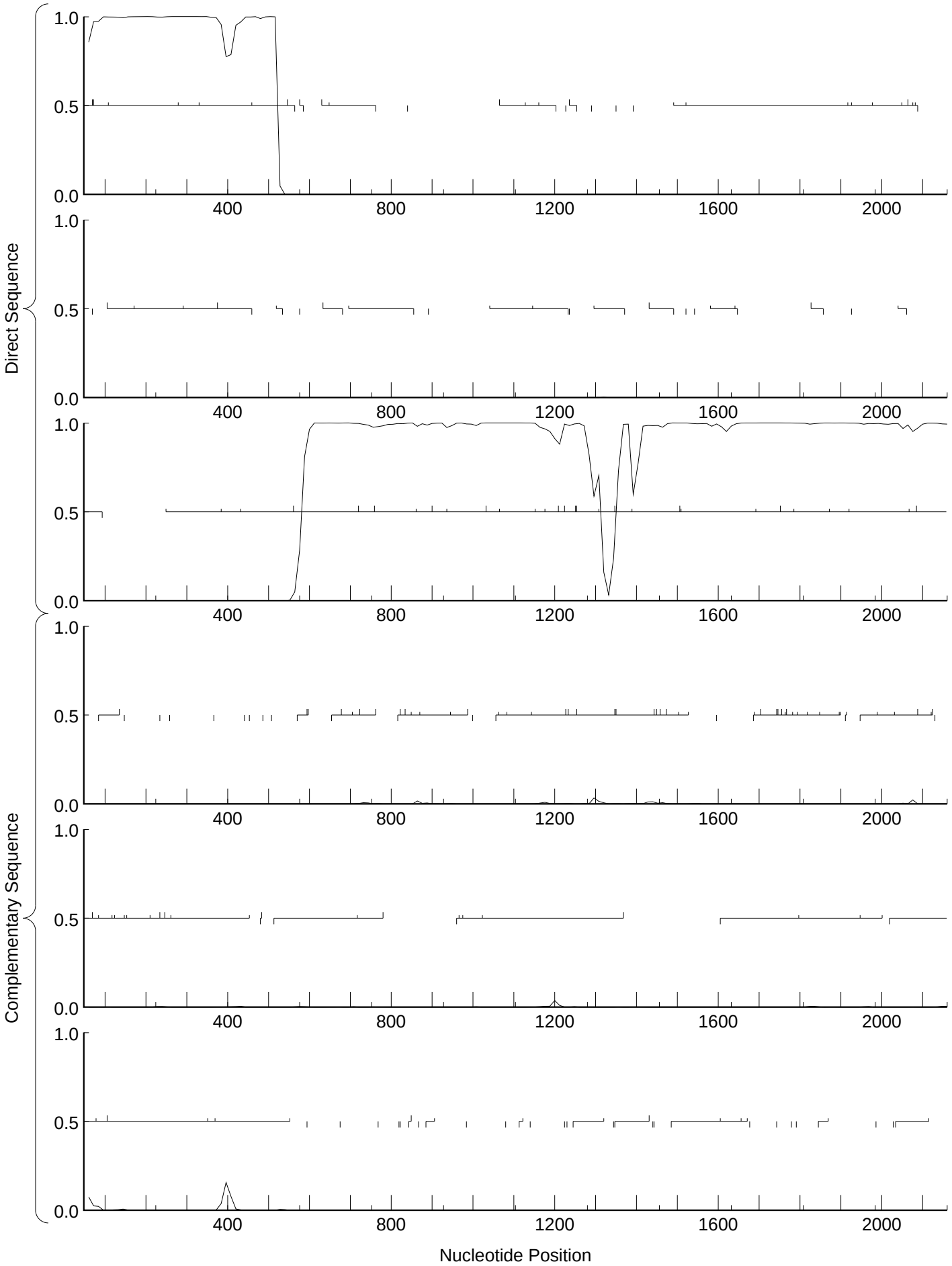
Order       :  4


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

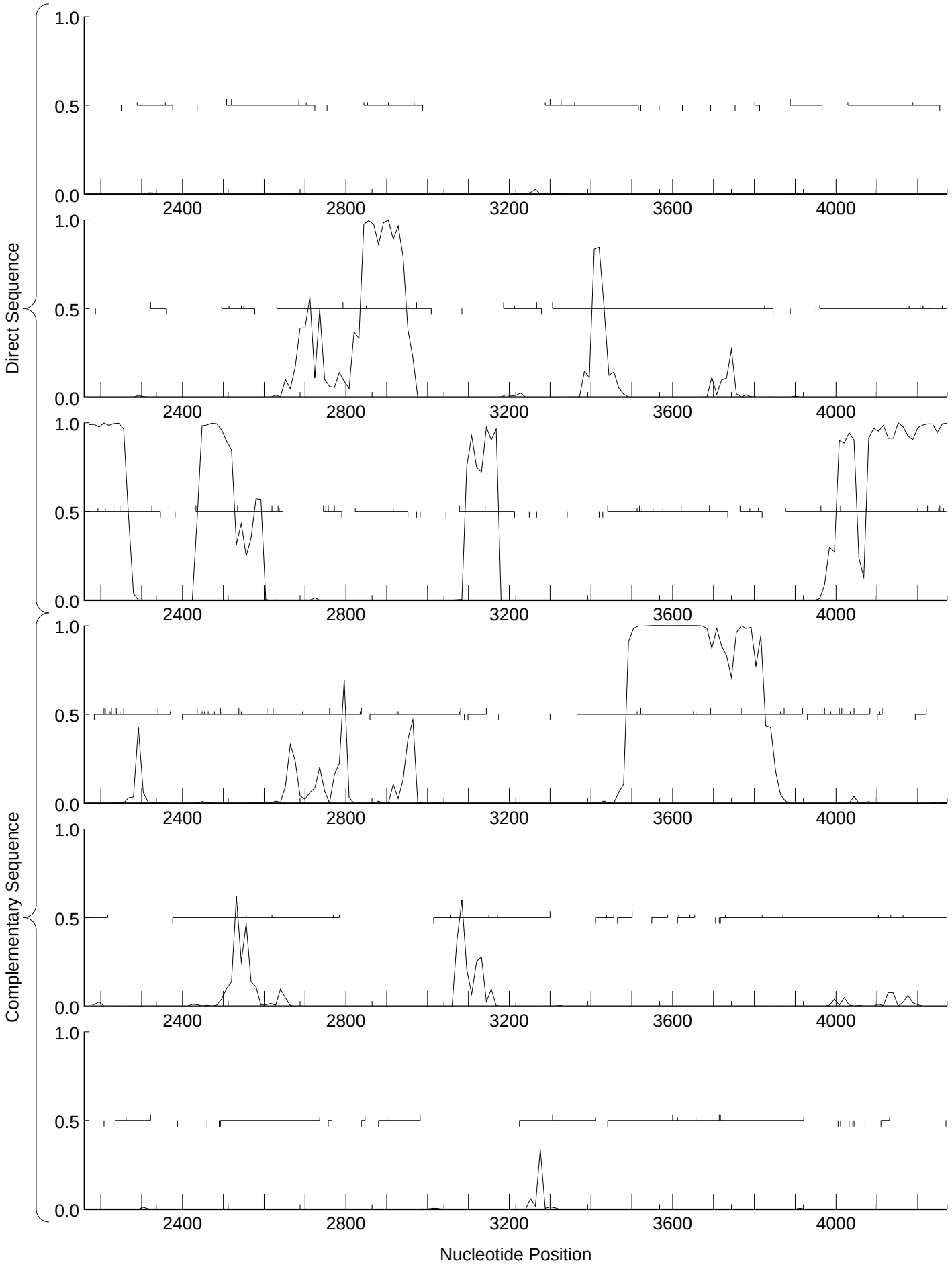


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

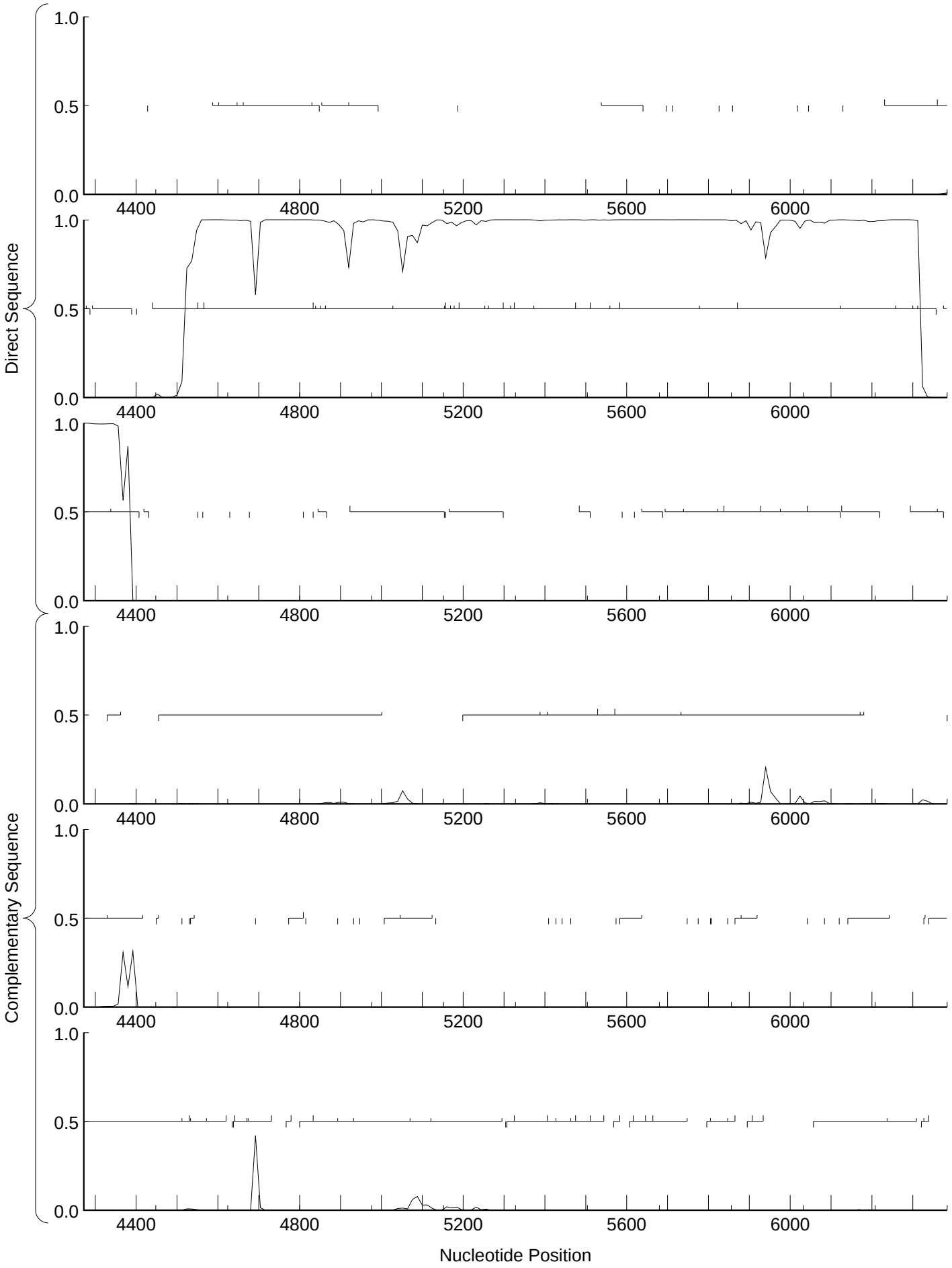
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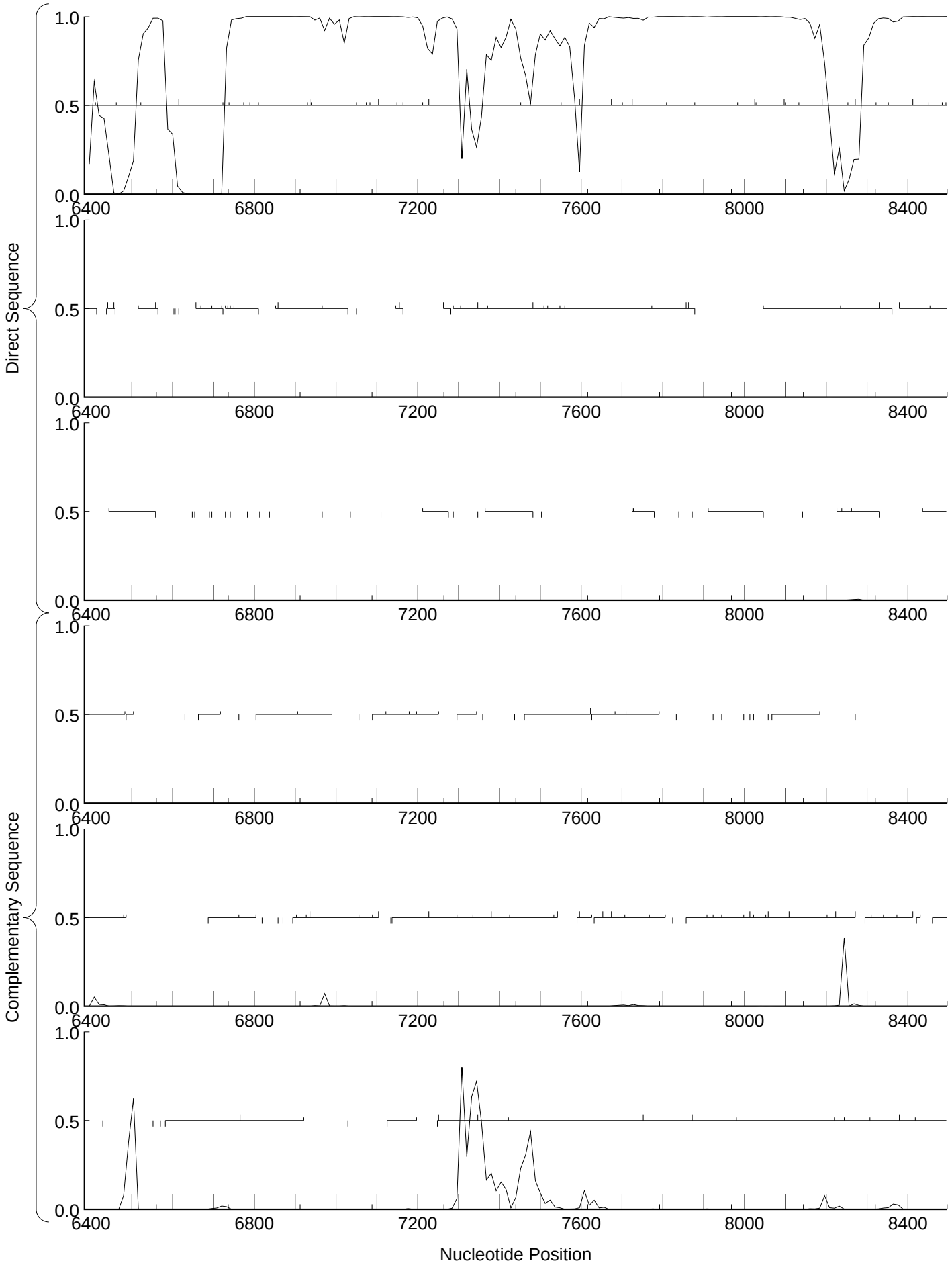
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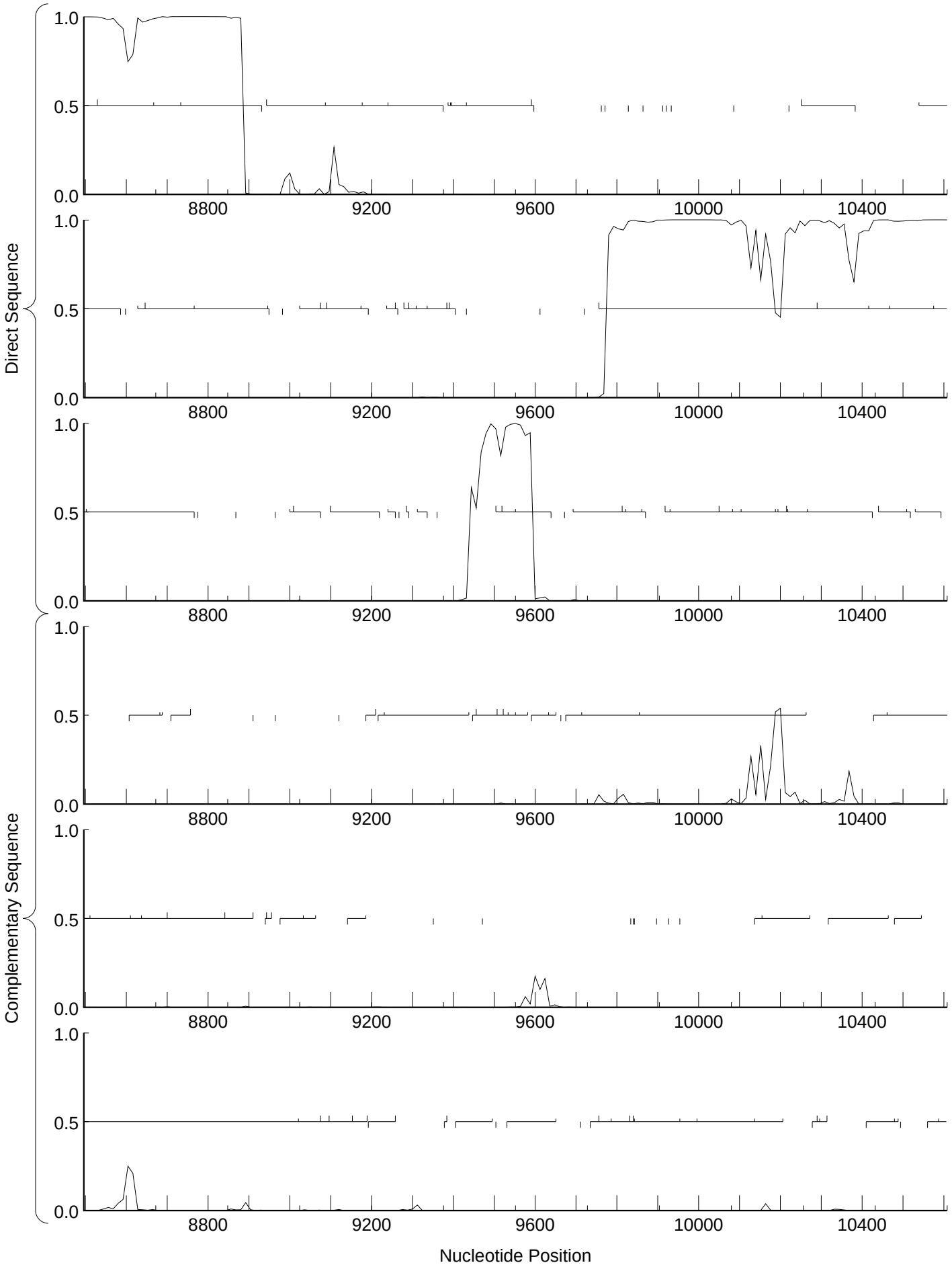
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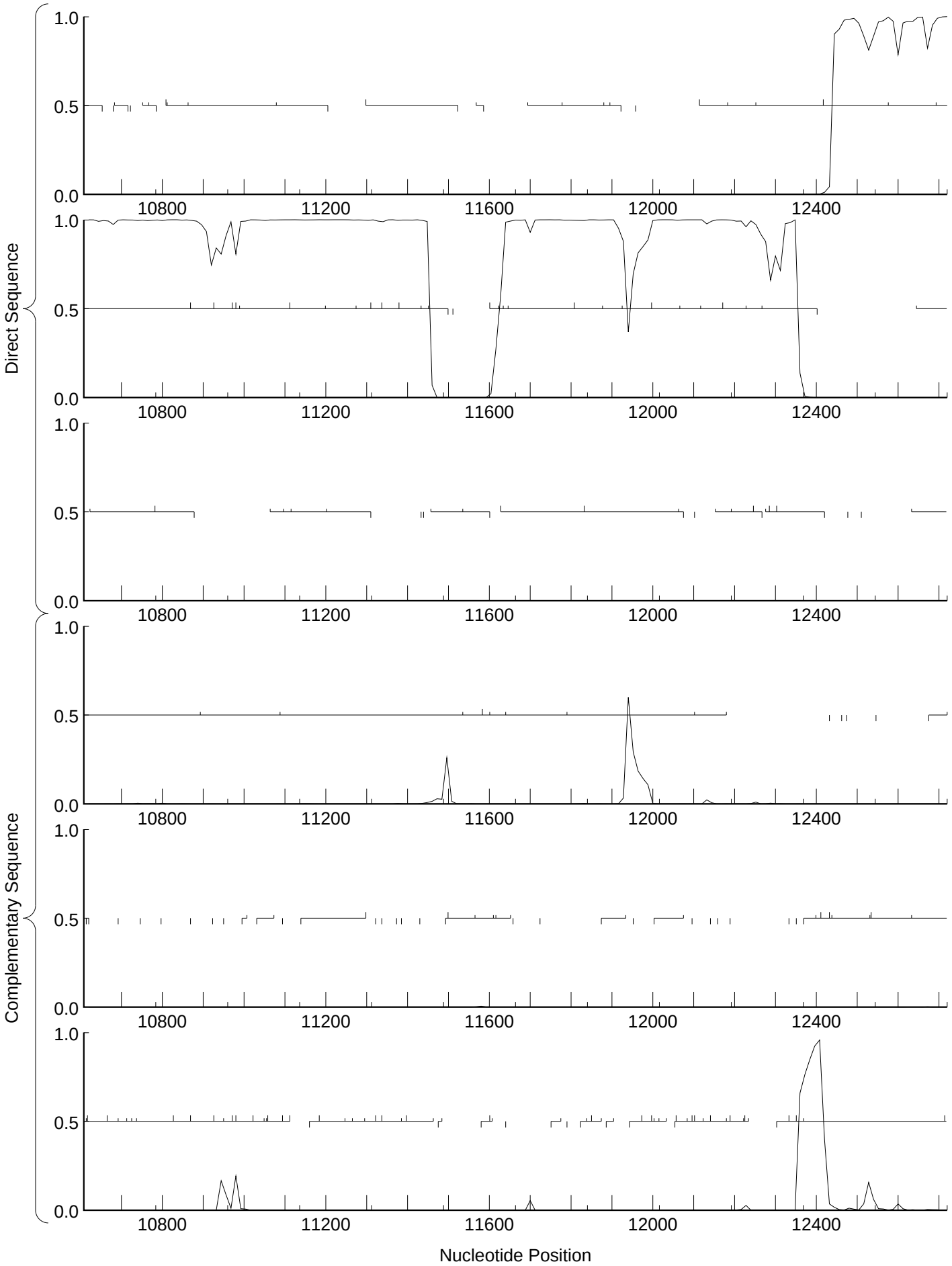
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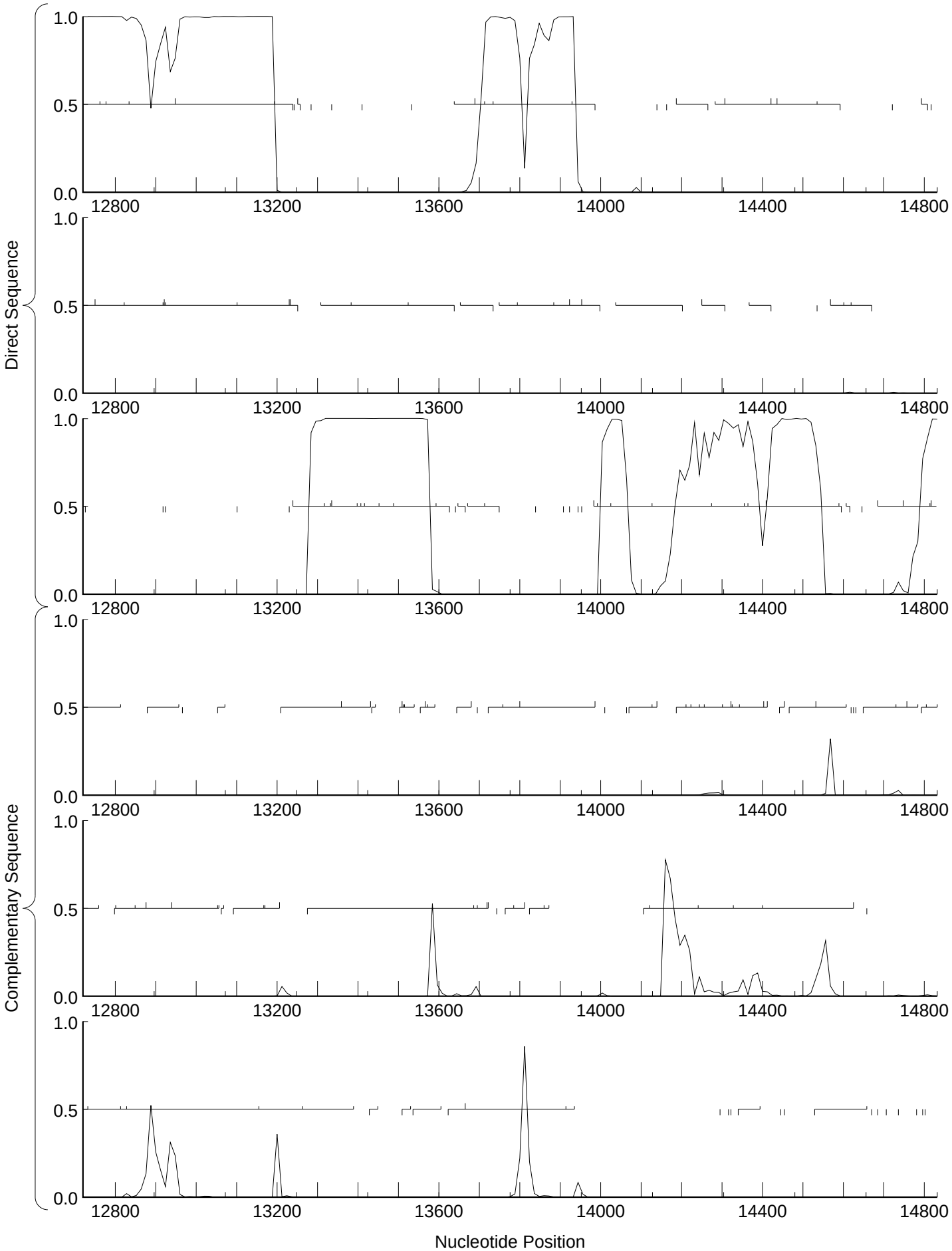
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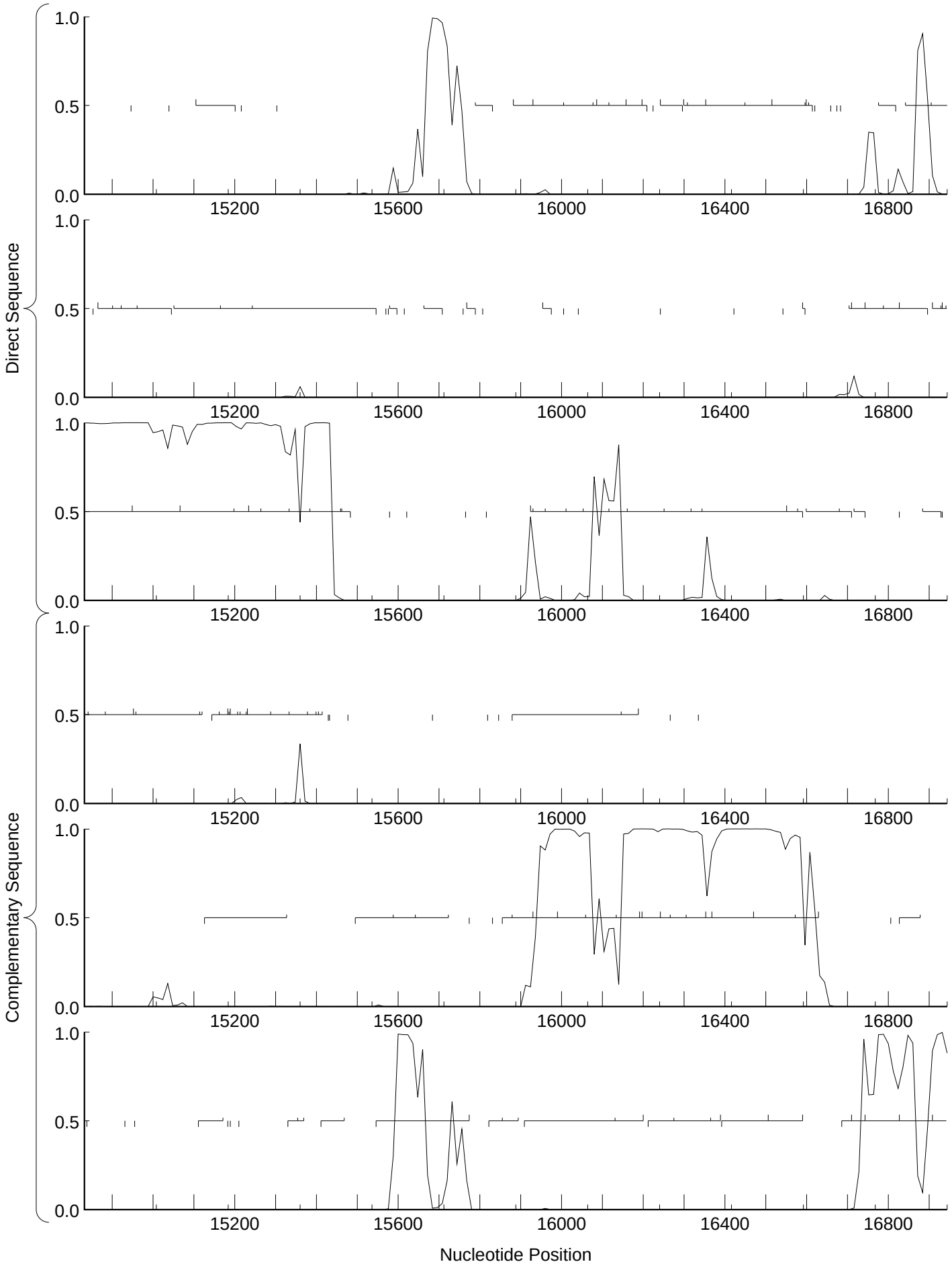
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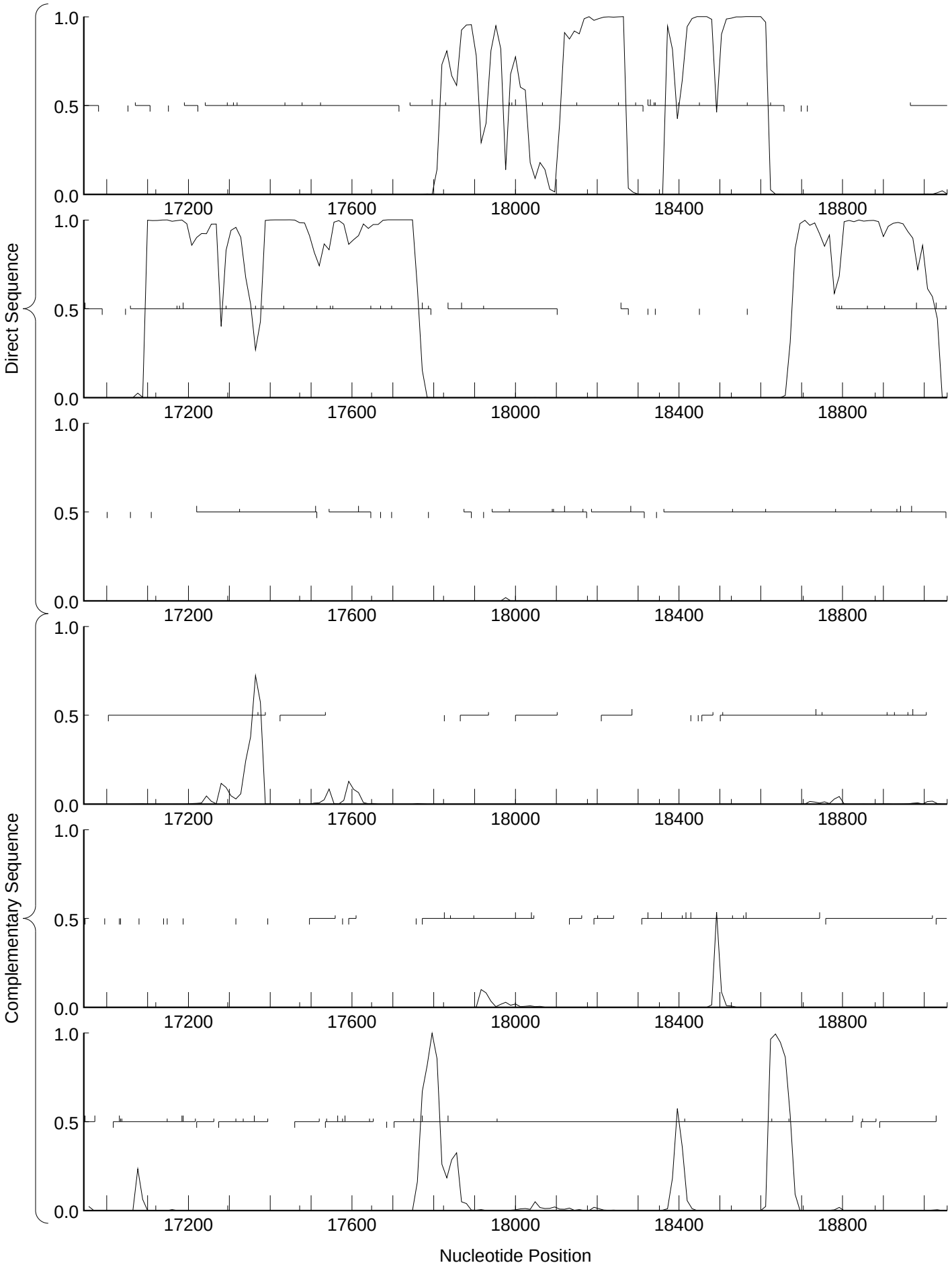


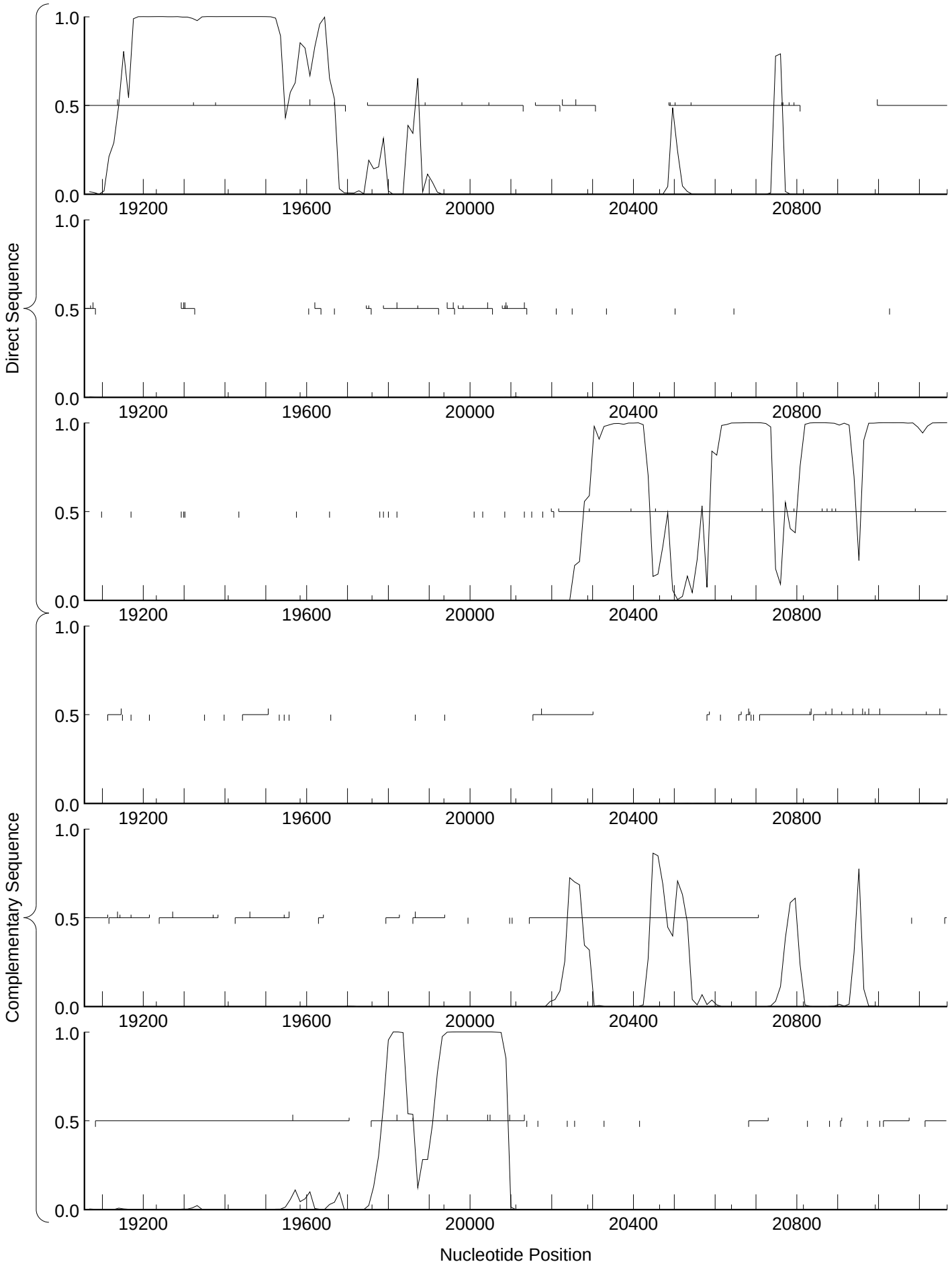
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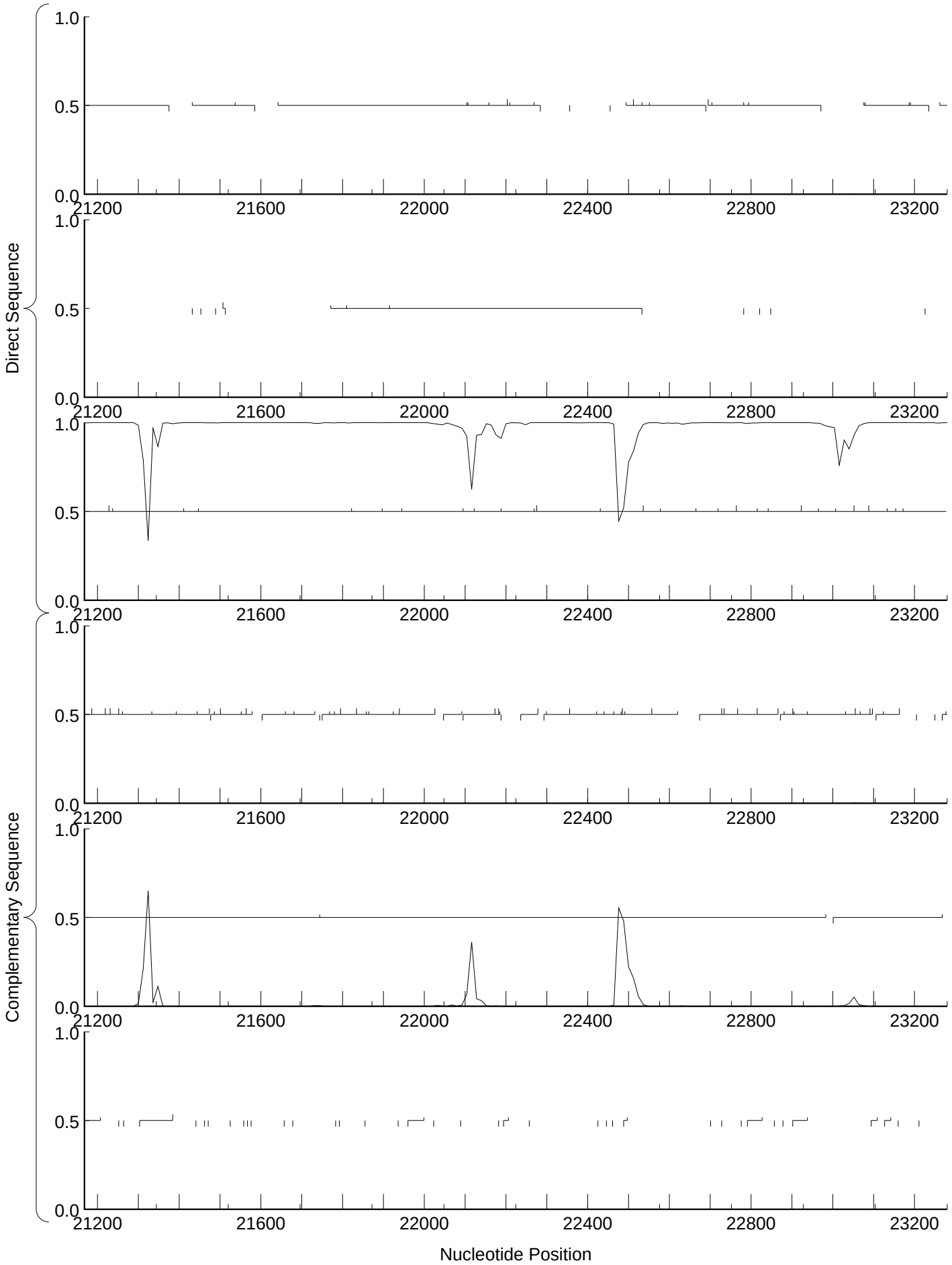
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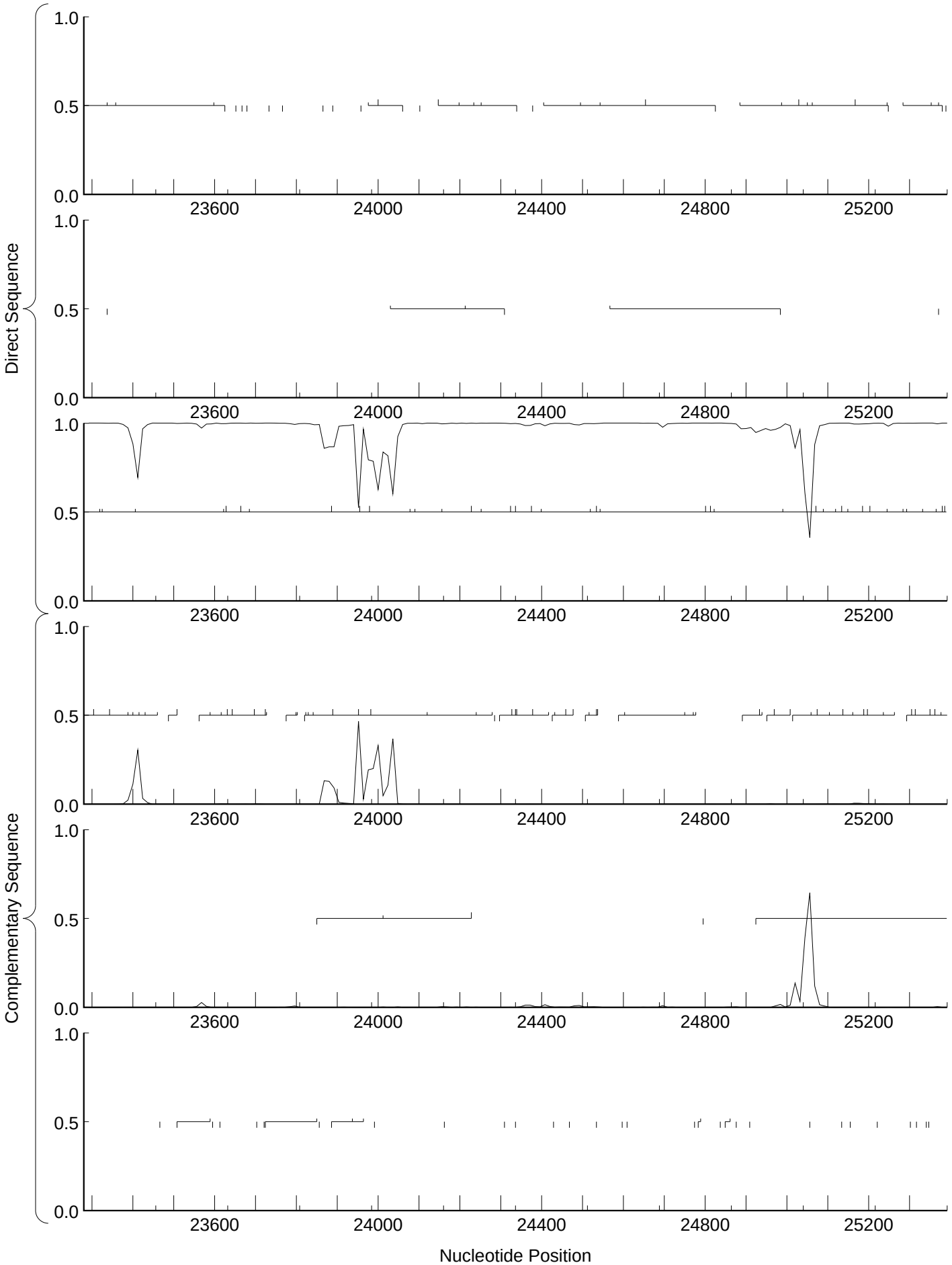


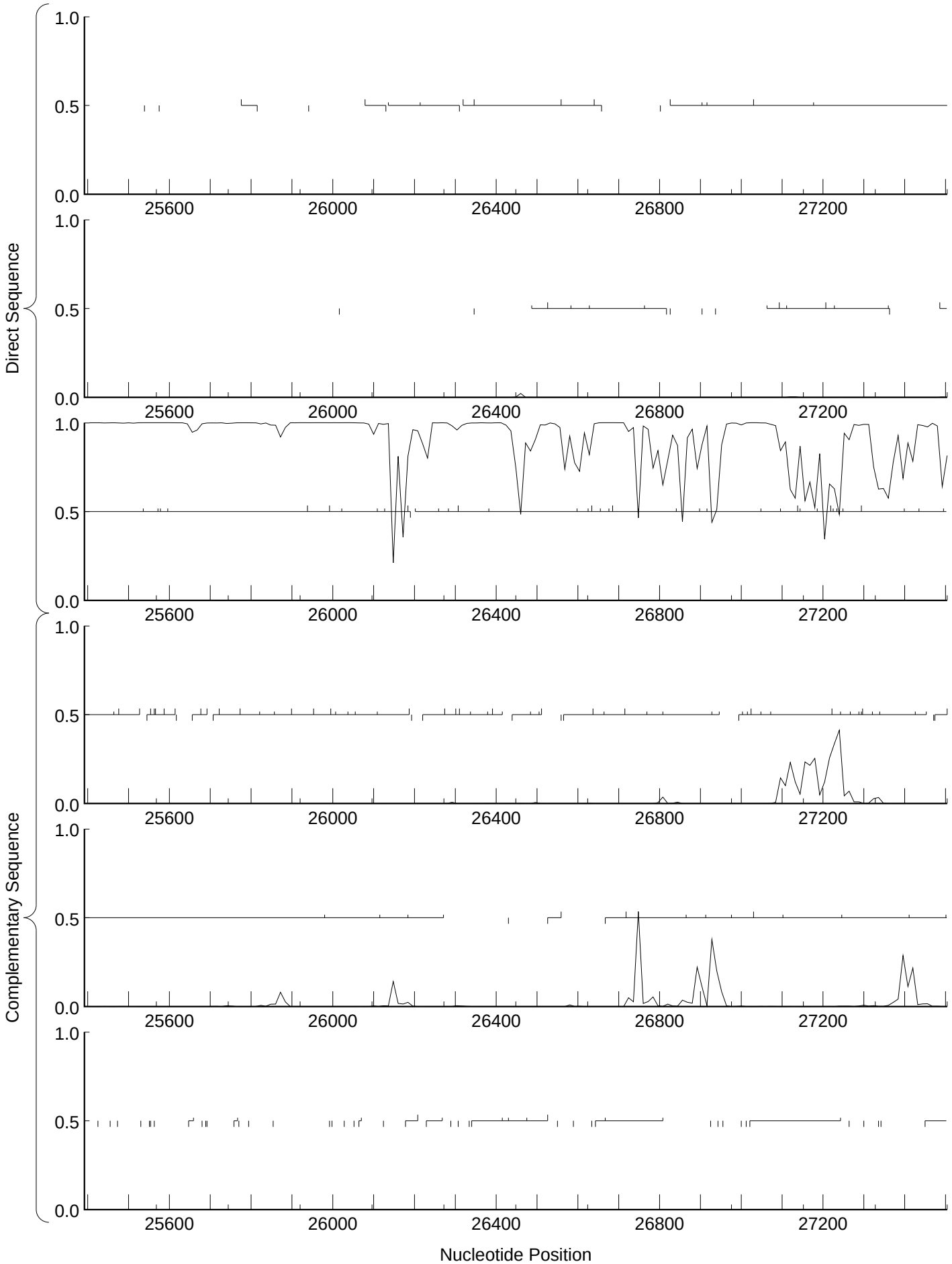

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Complementary Sequence
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0.0
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17600
18000
18400
18800
Nucleotide Position

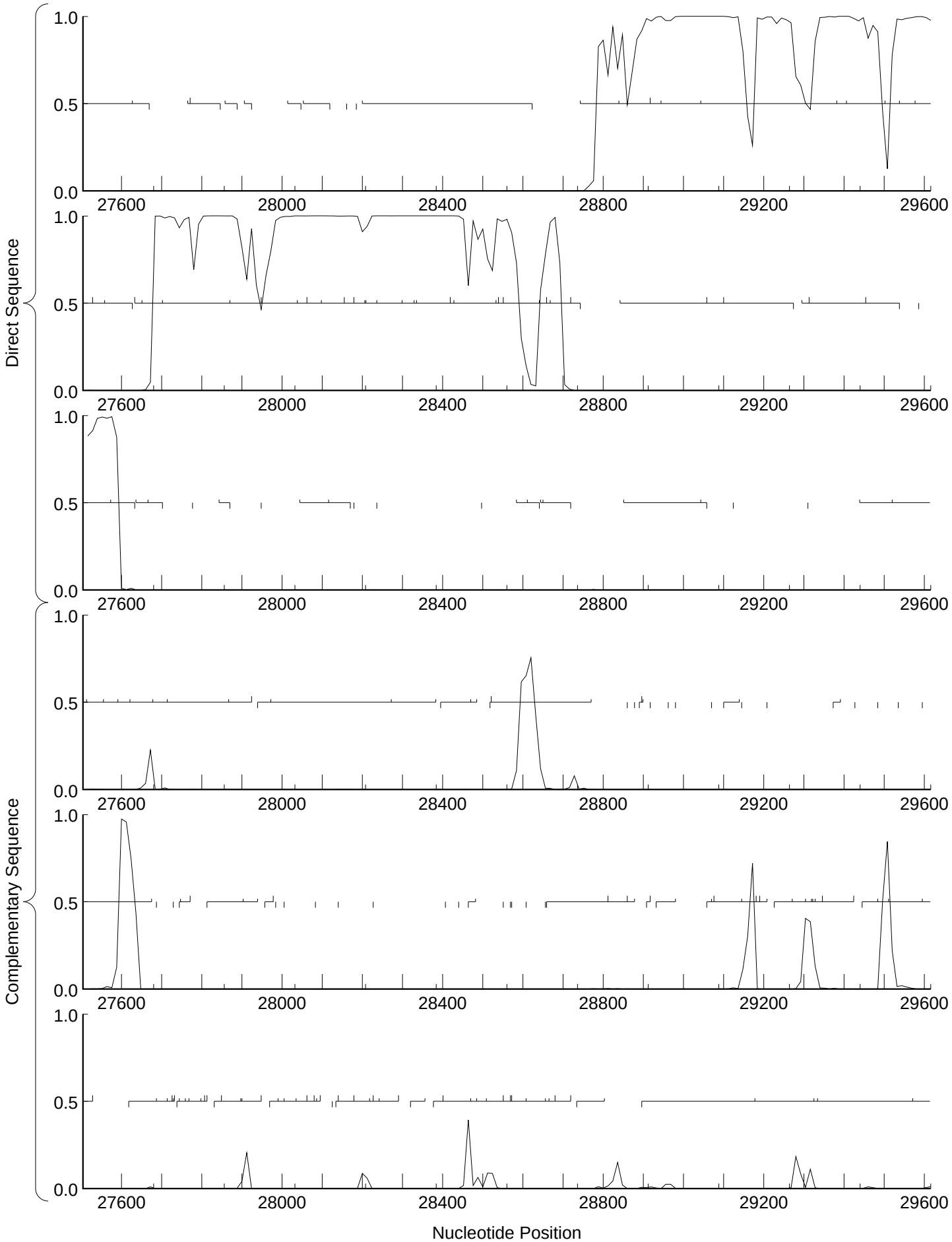

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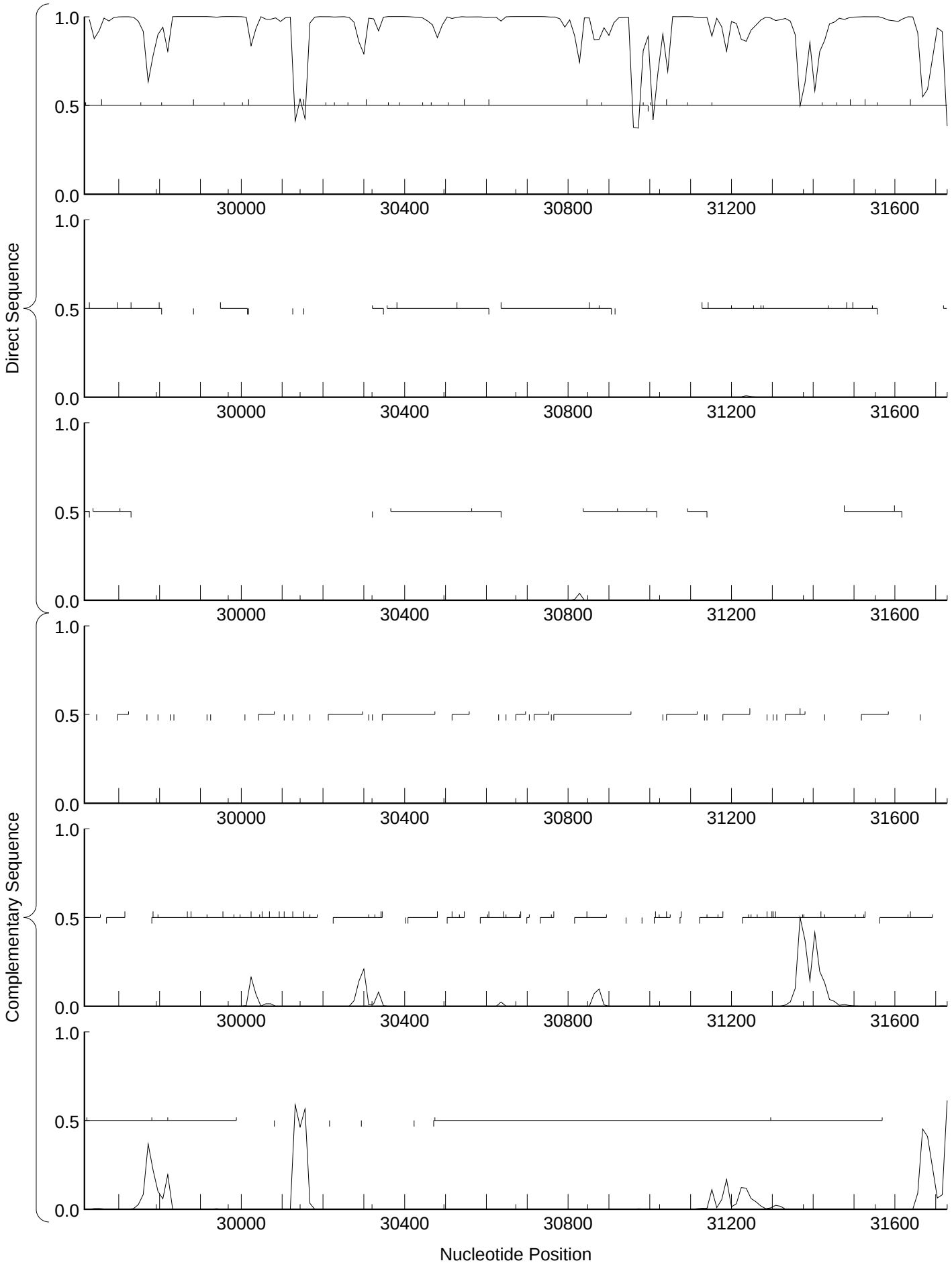
bacterium phage KingVeveve complete sequence, 68043 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 12/34



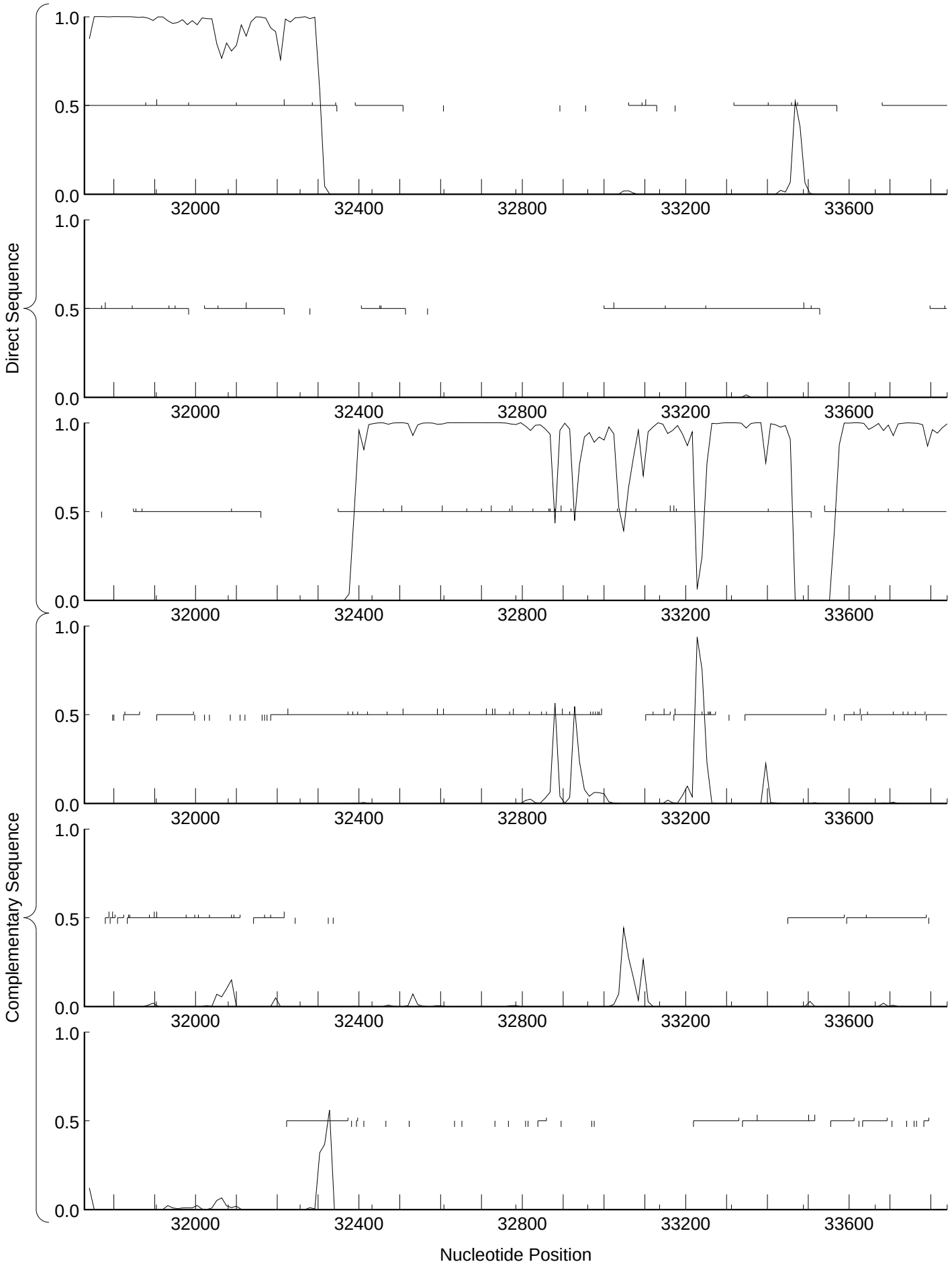

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Nucleotide Position

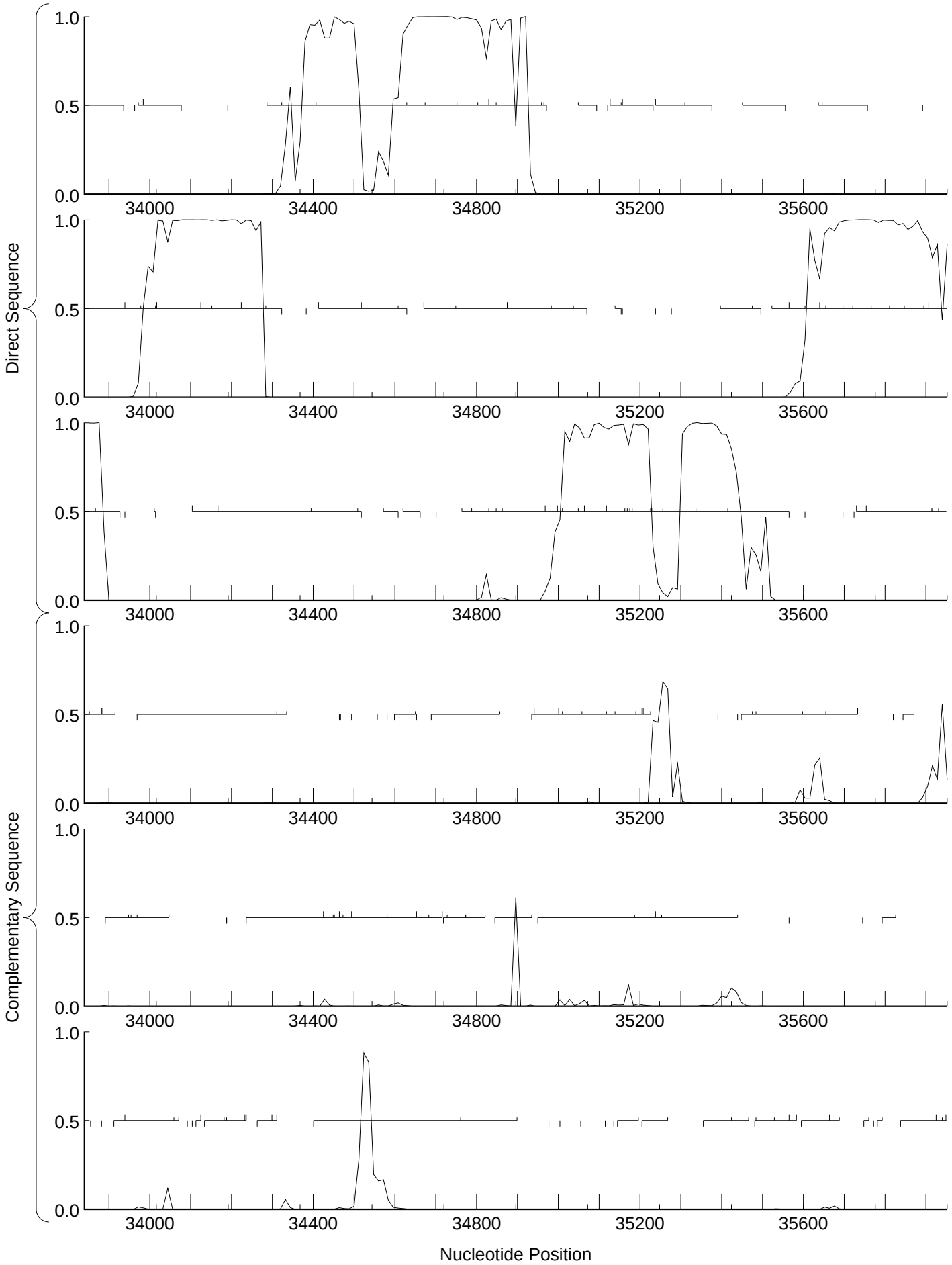

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Nucleotide Position

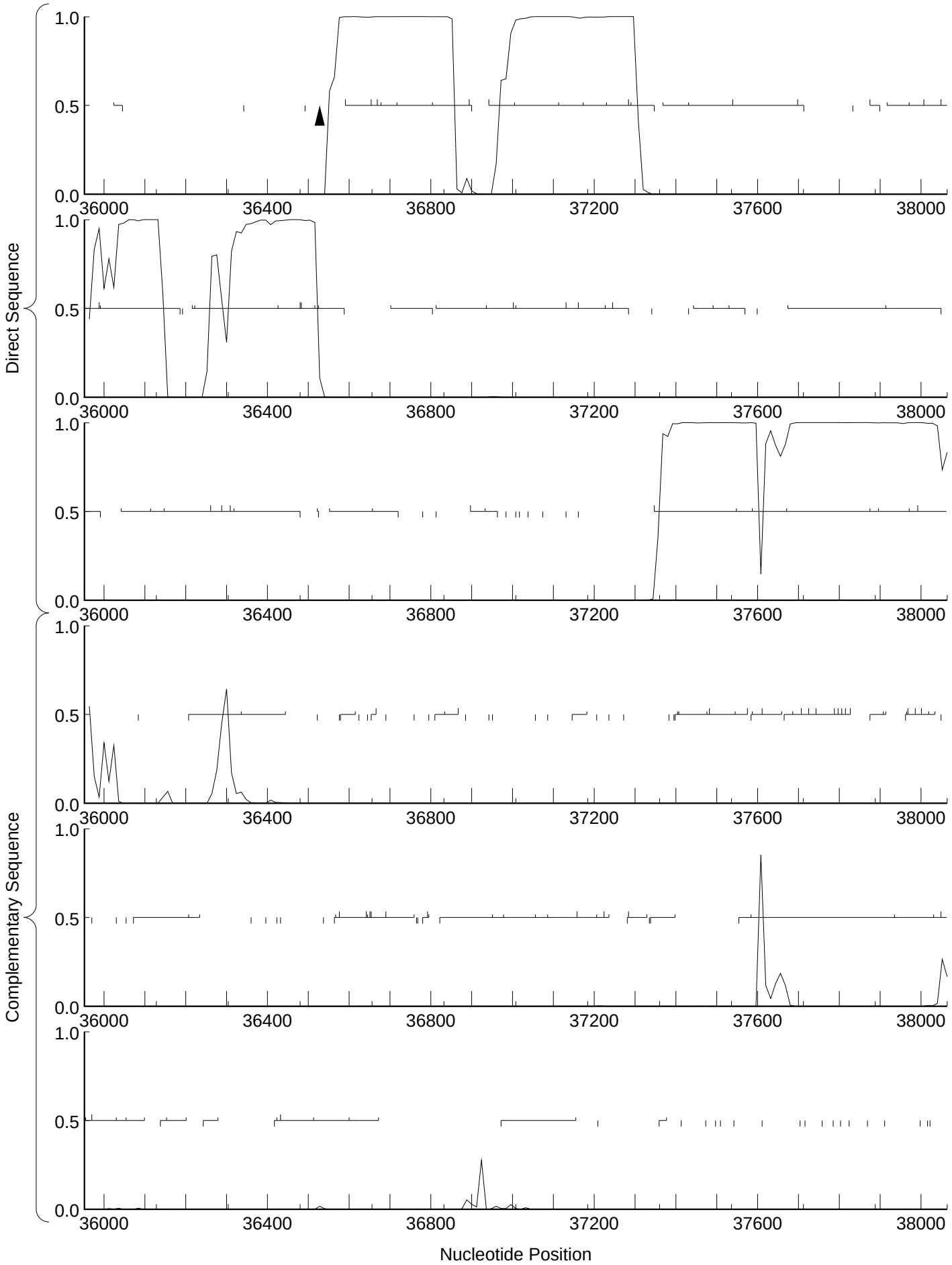

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Nucleotide Position

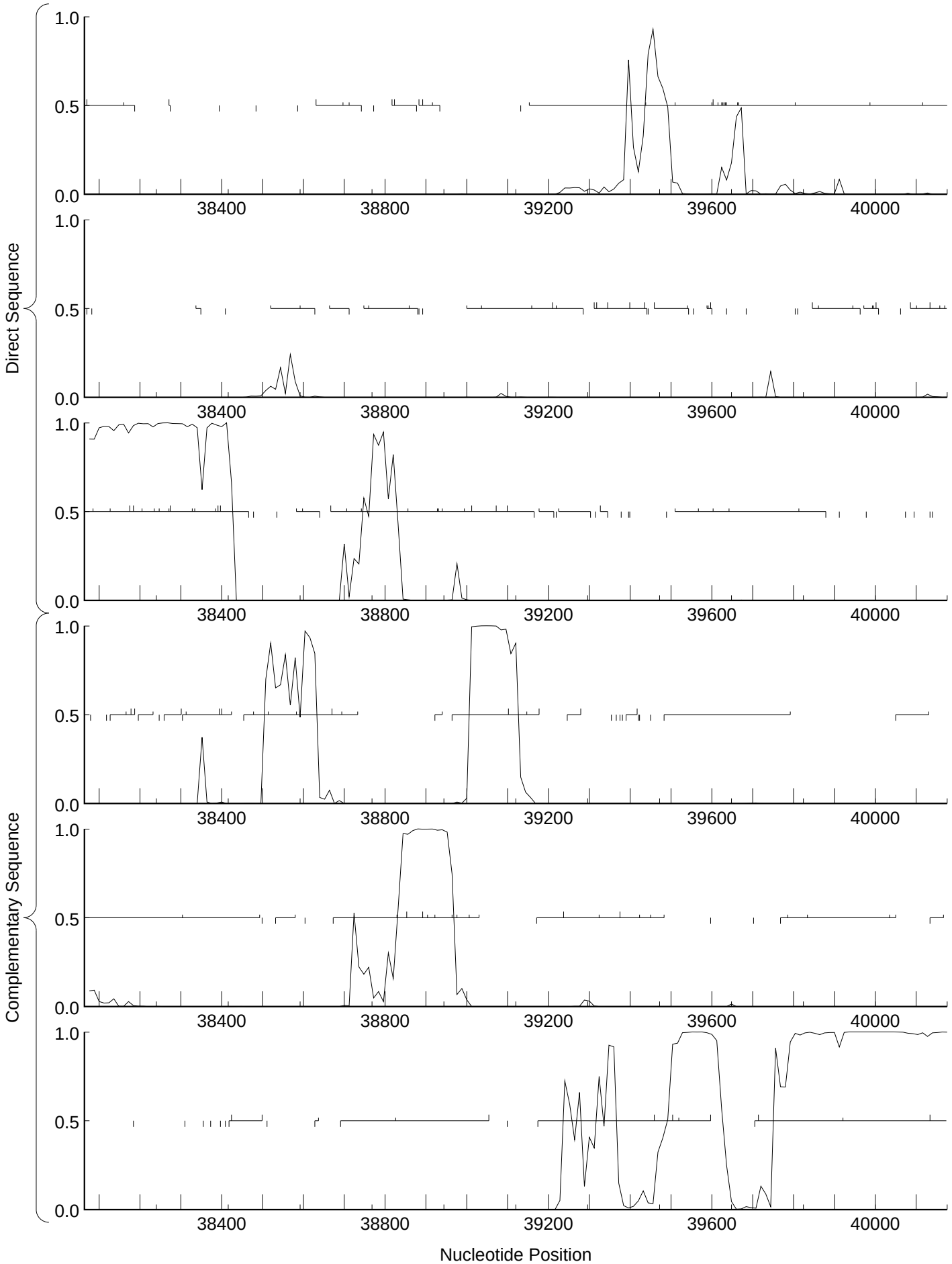

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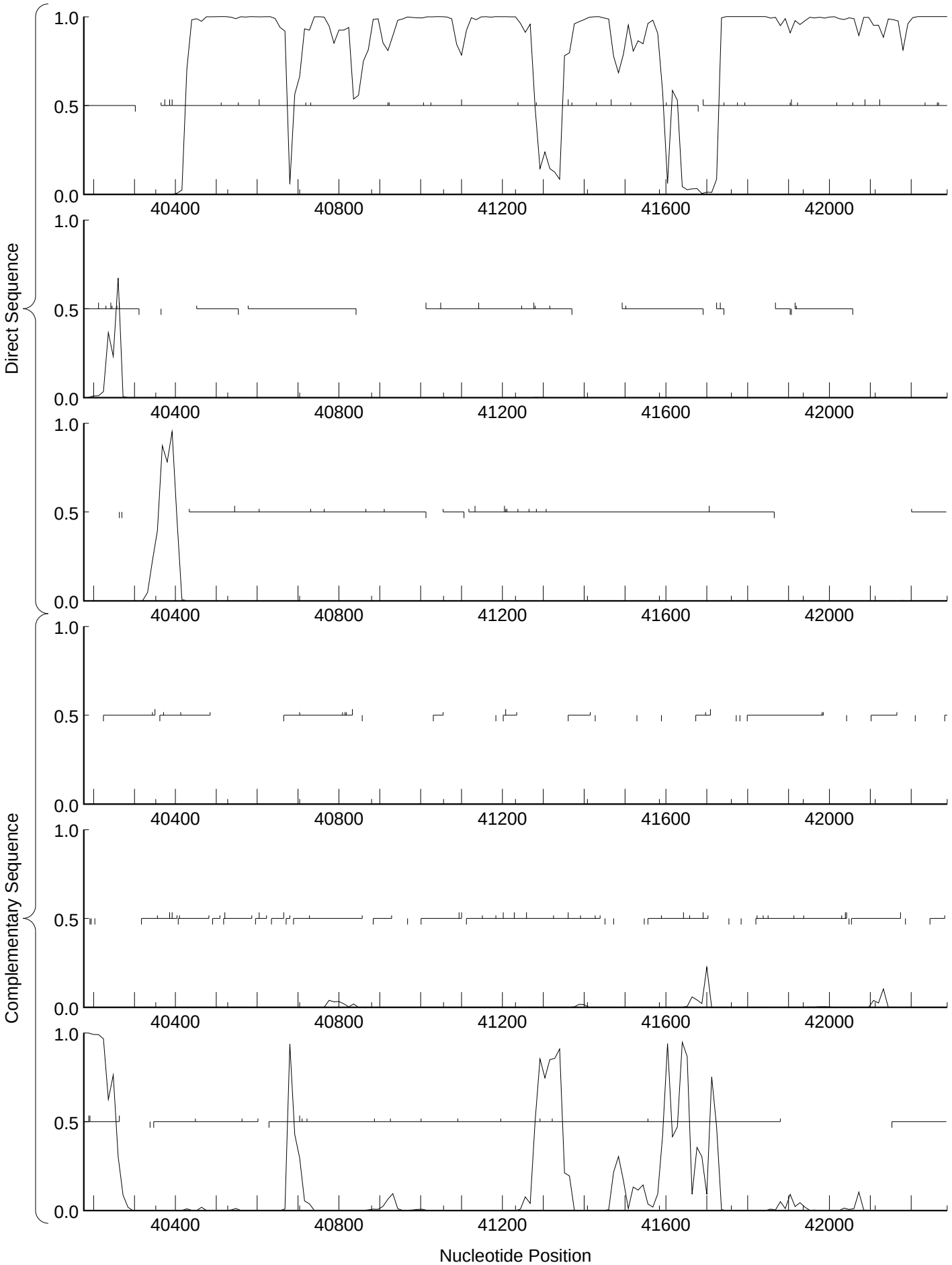
bacterium phage KingVeveve complete sequence, 68043 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 17/34

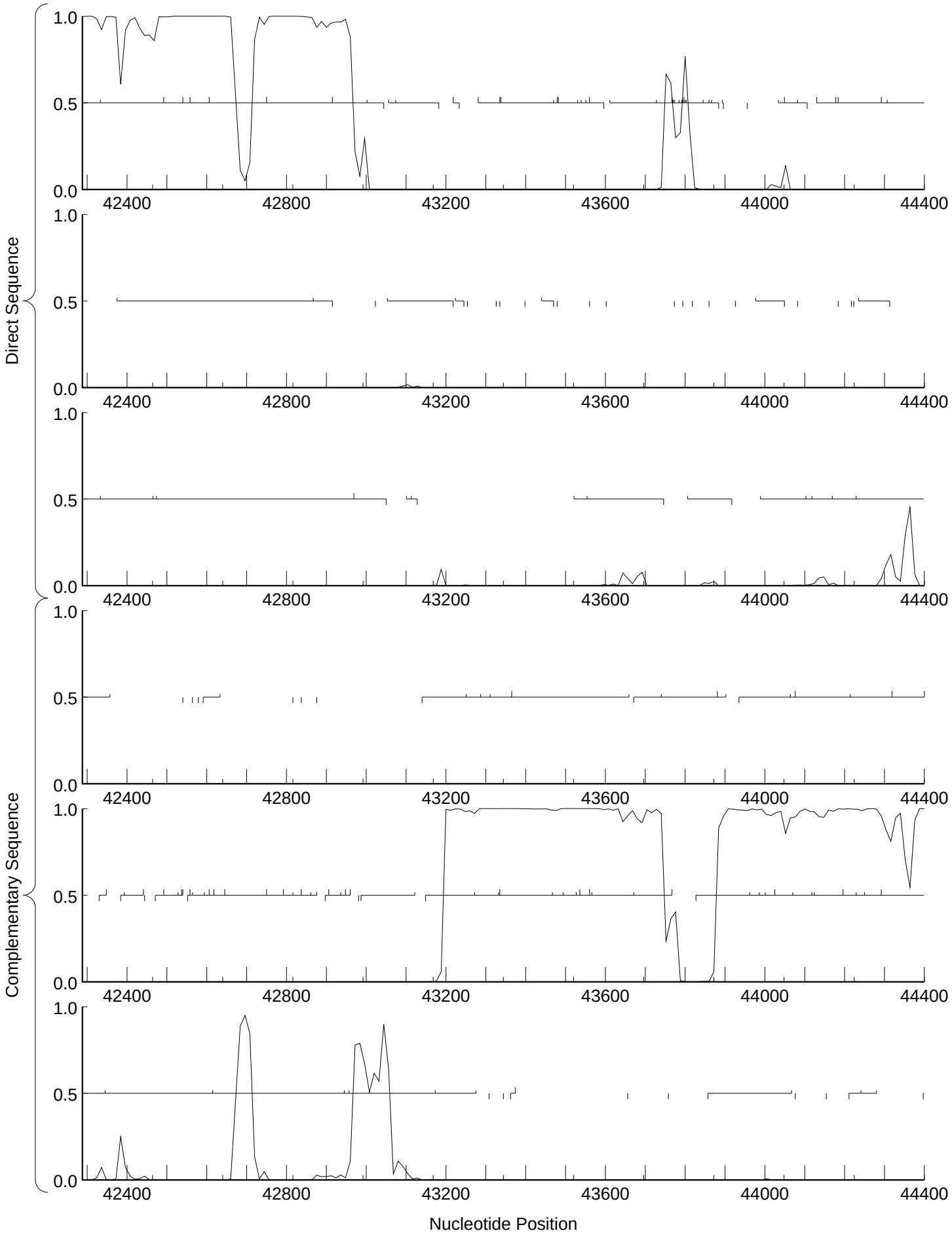


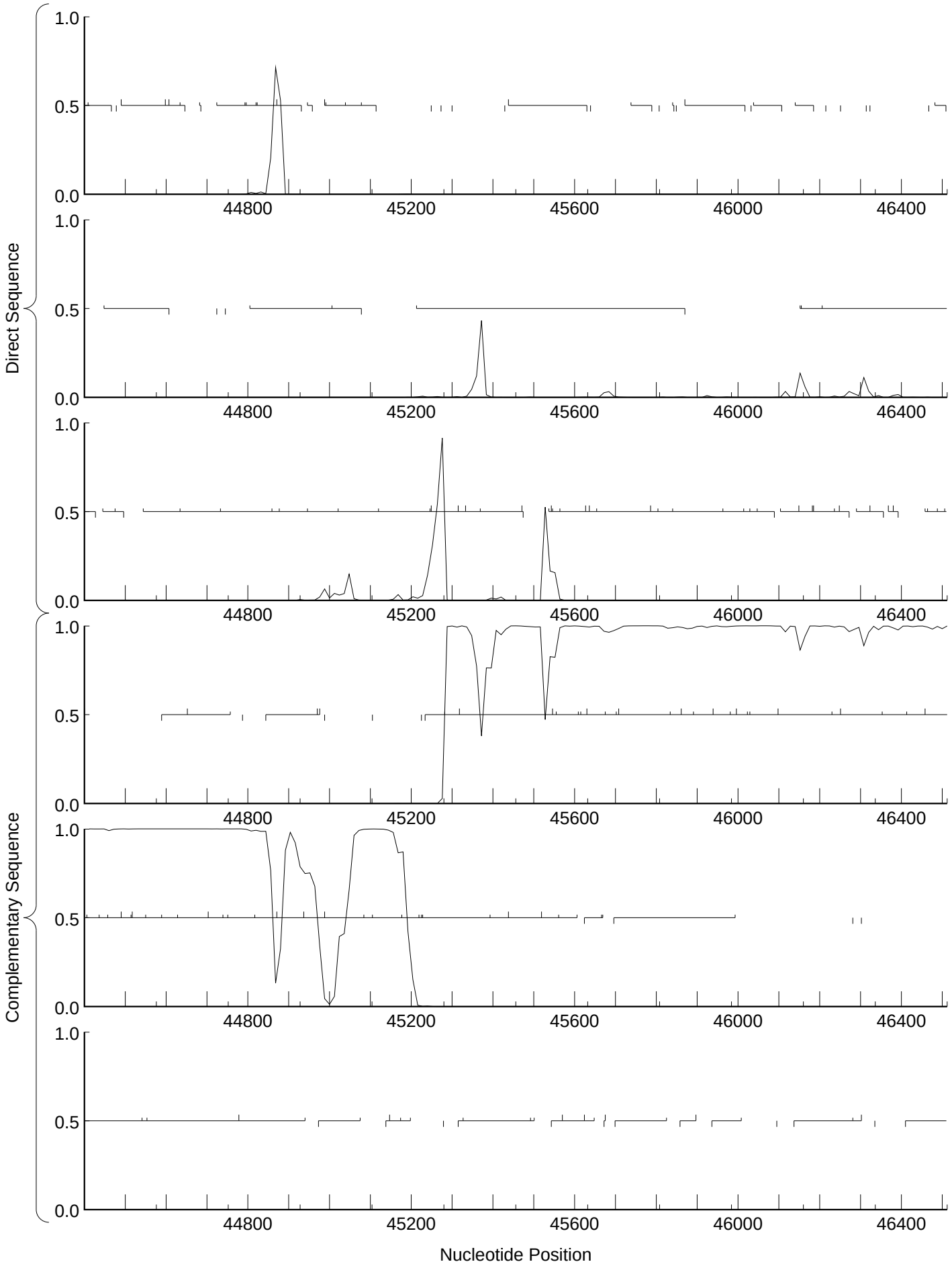

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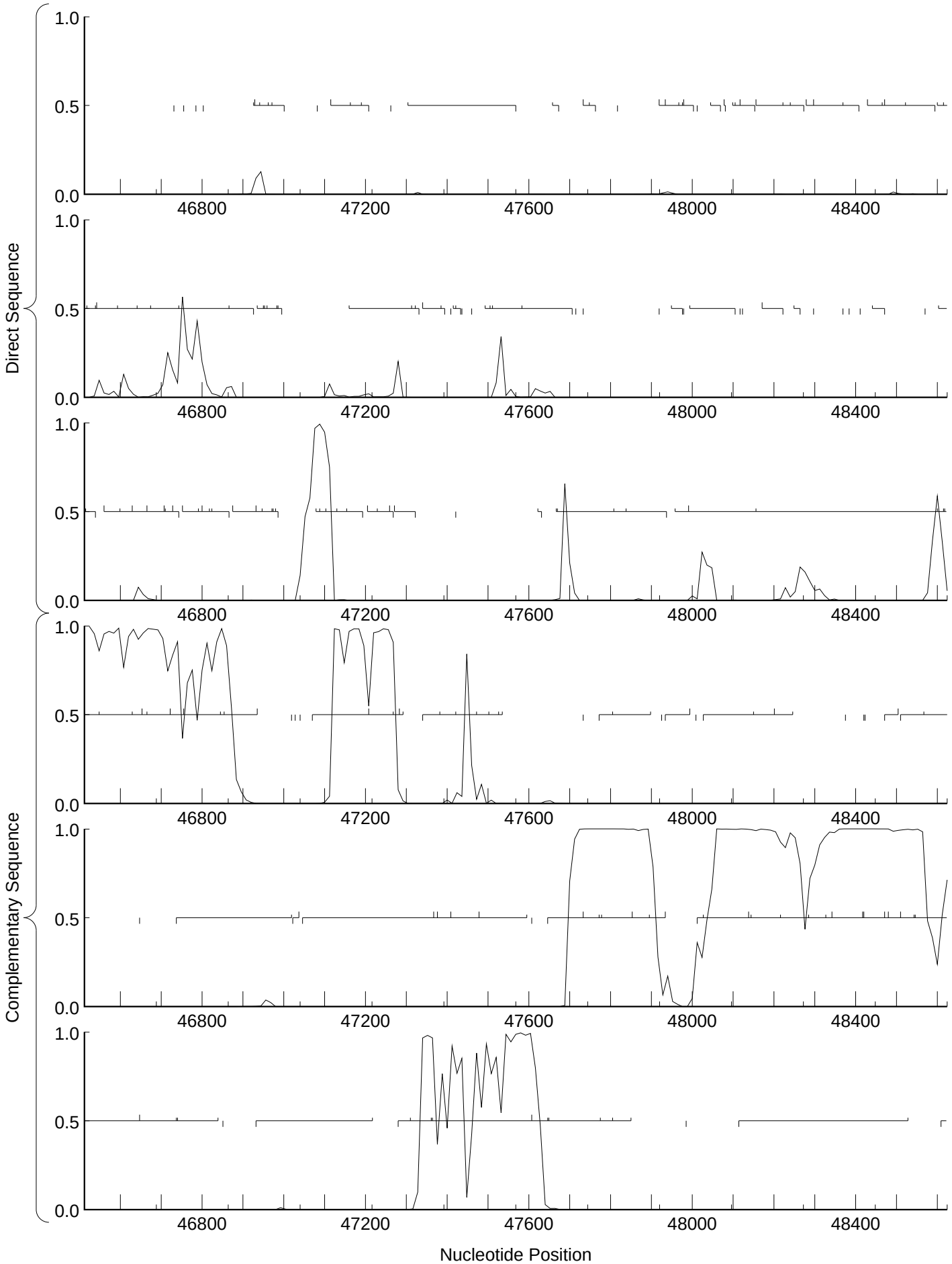

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Nucleotide Position

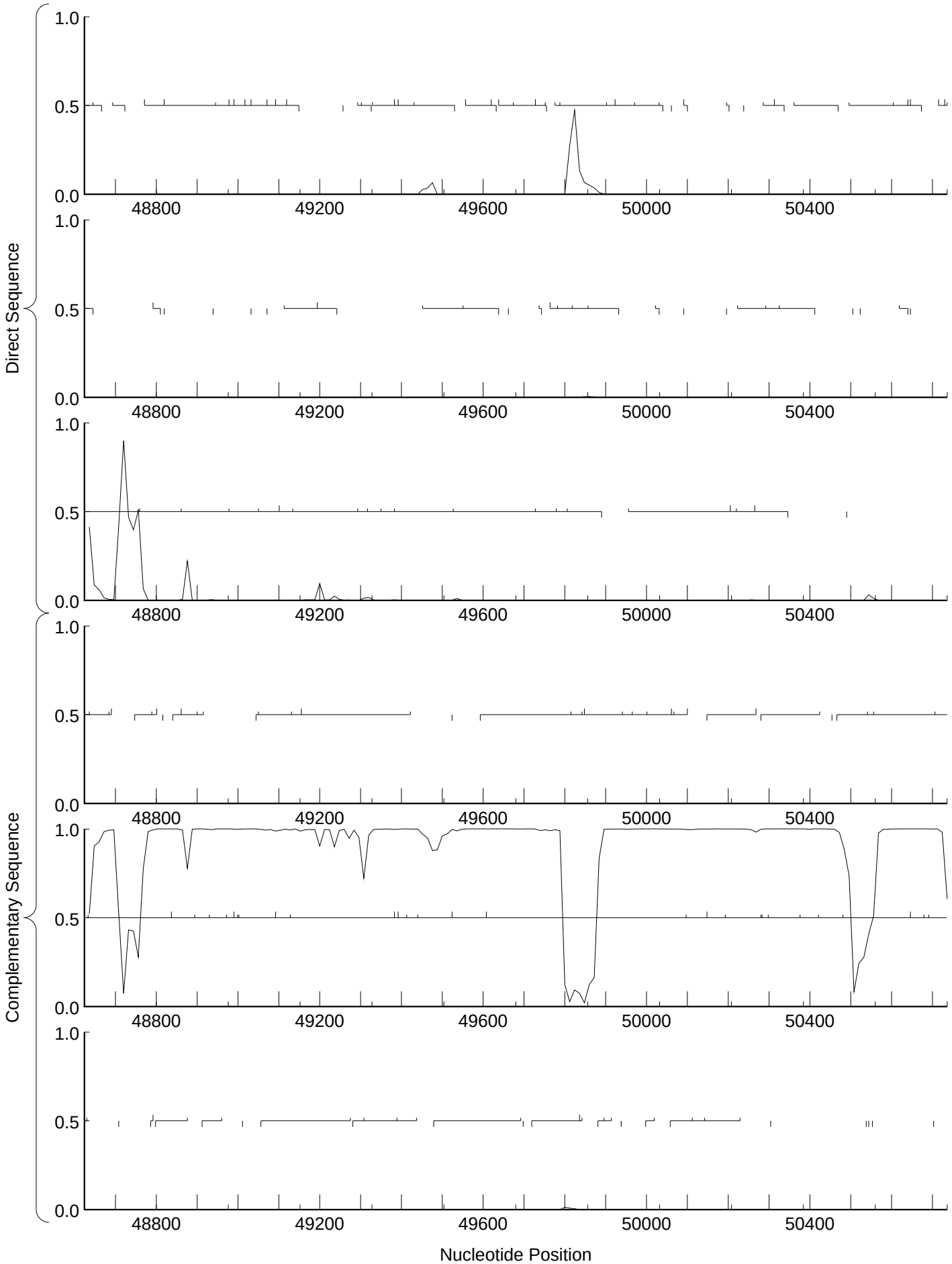

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Nucleotide Position

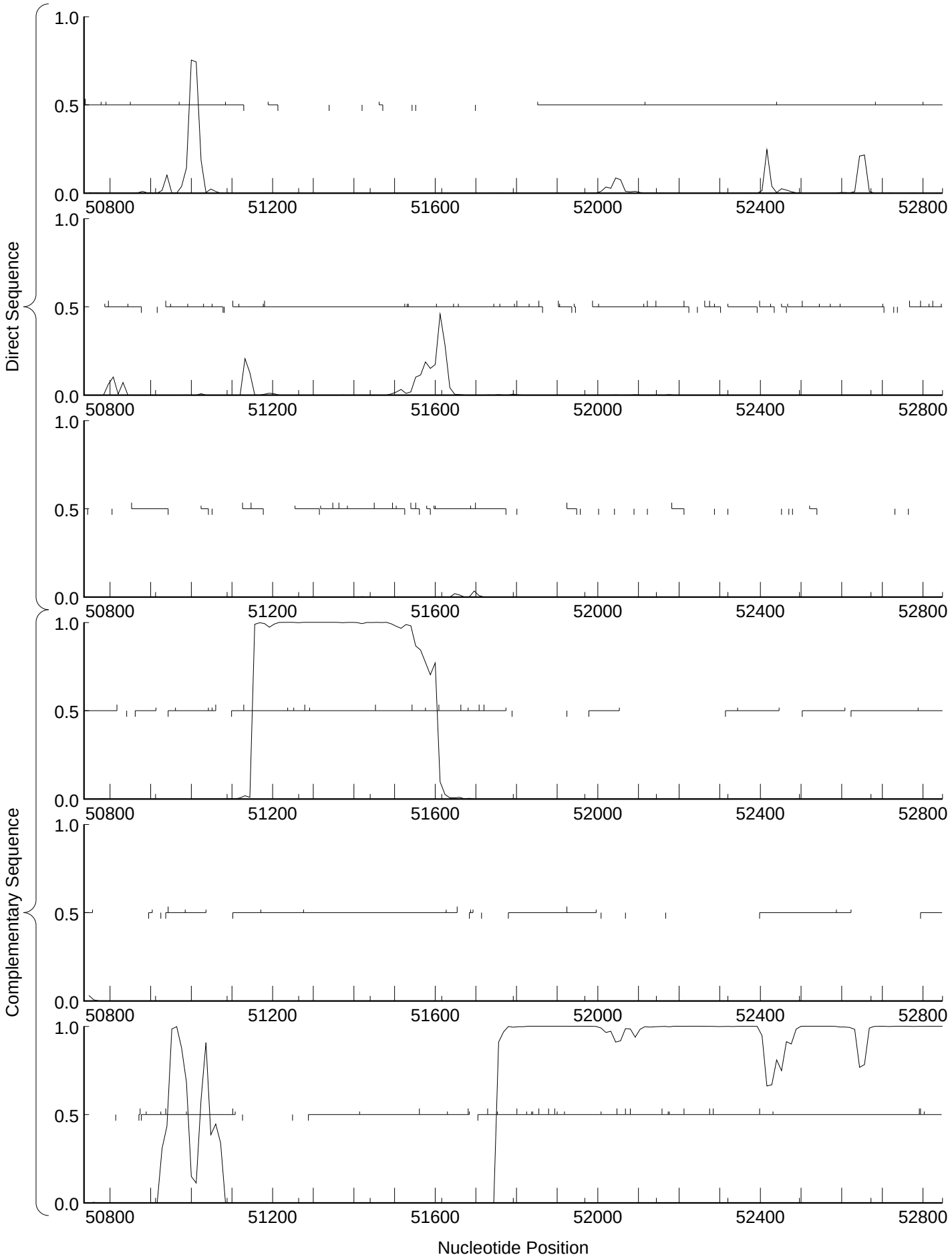

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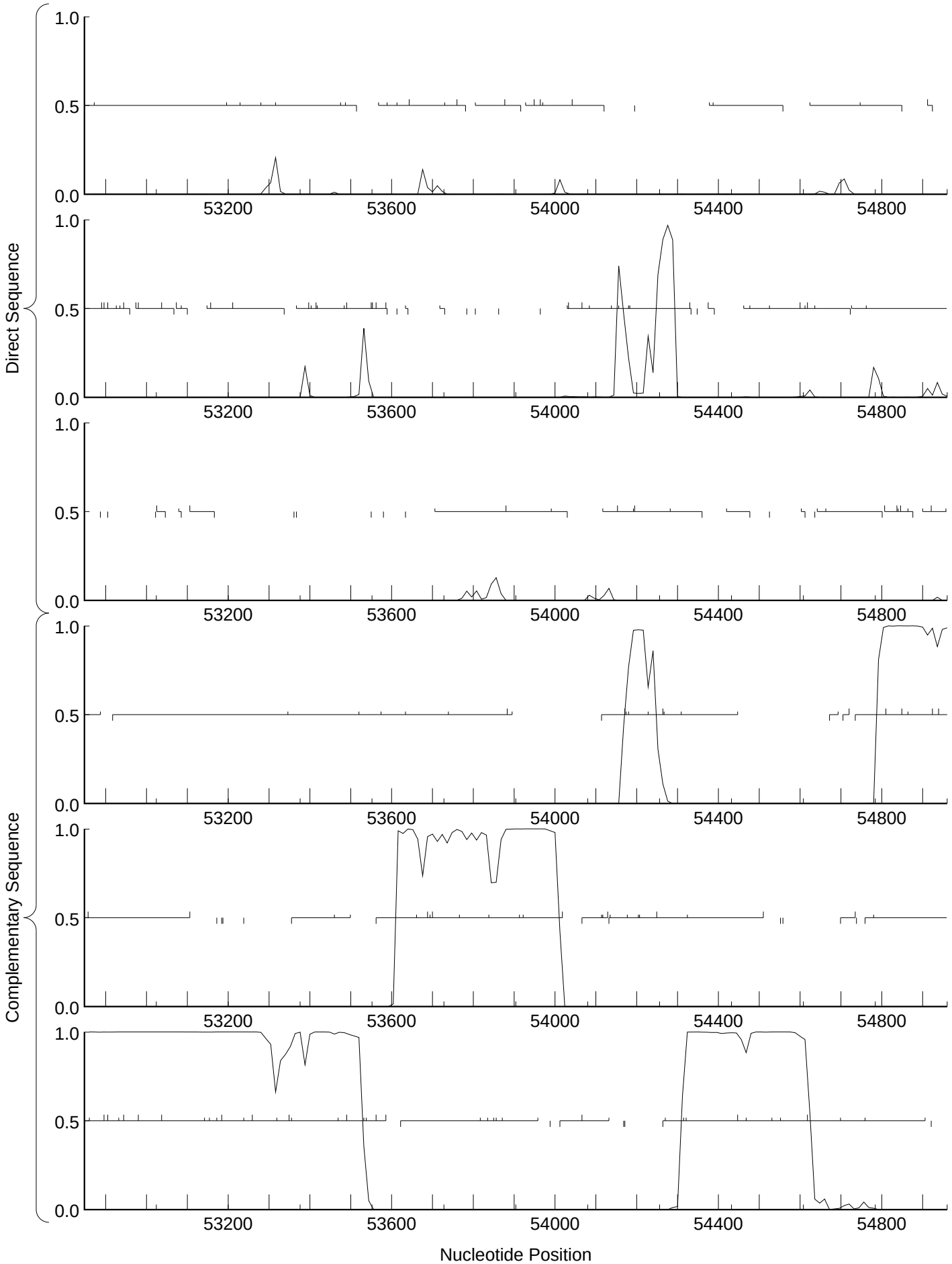

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Nucleotide Position

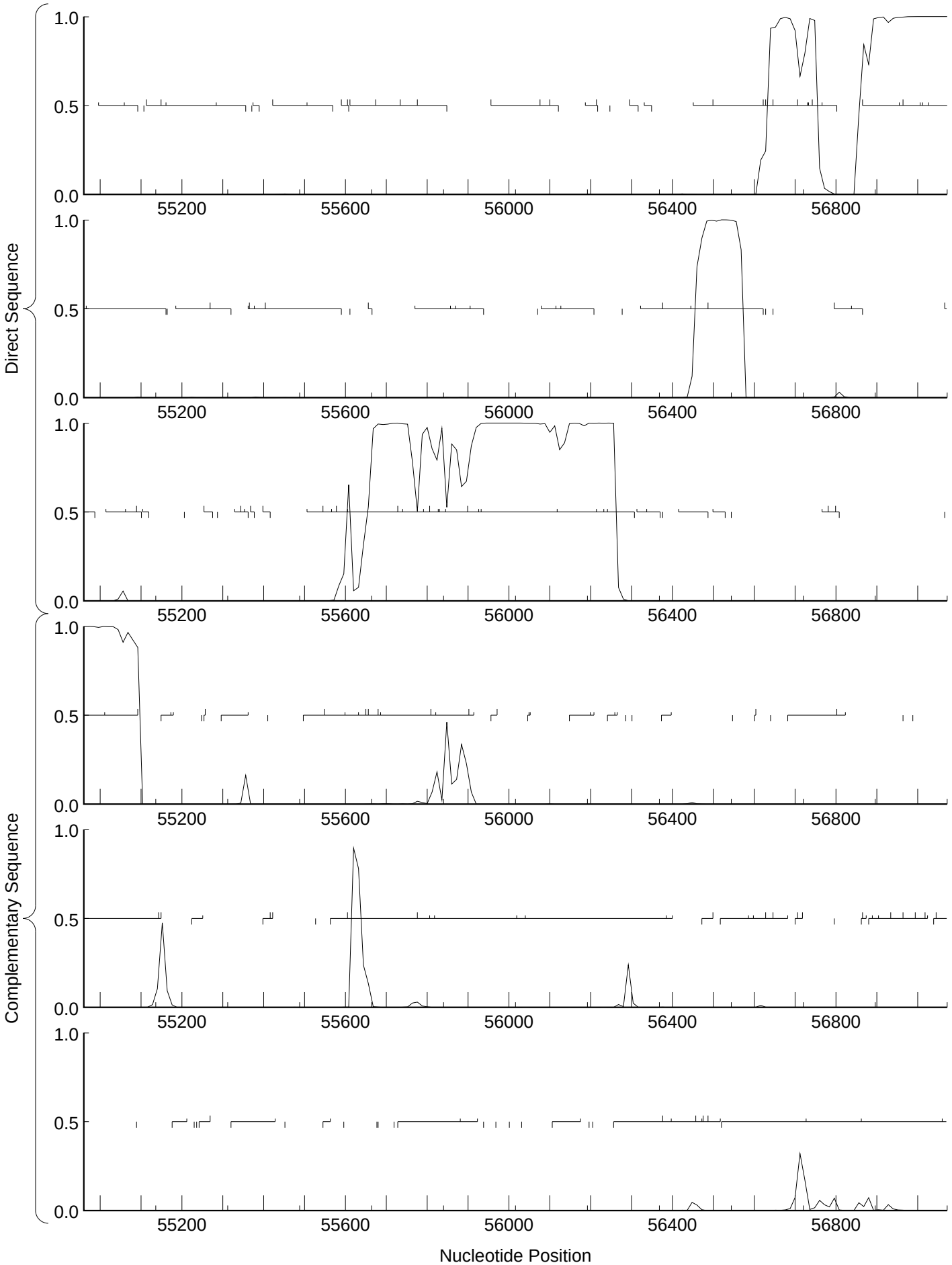

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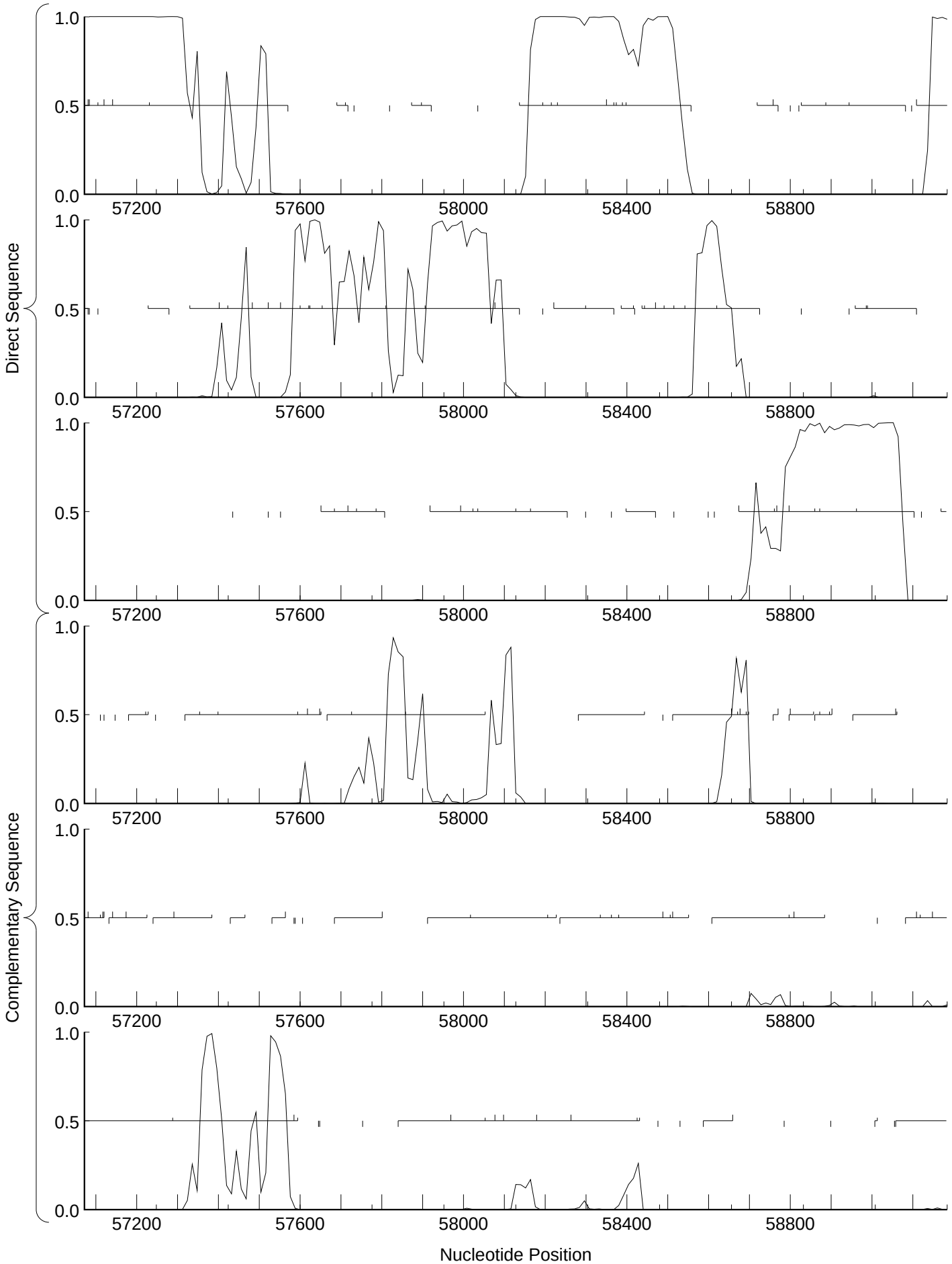

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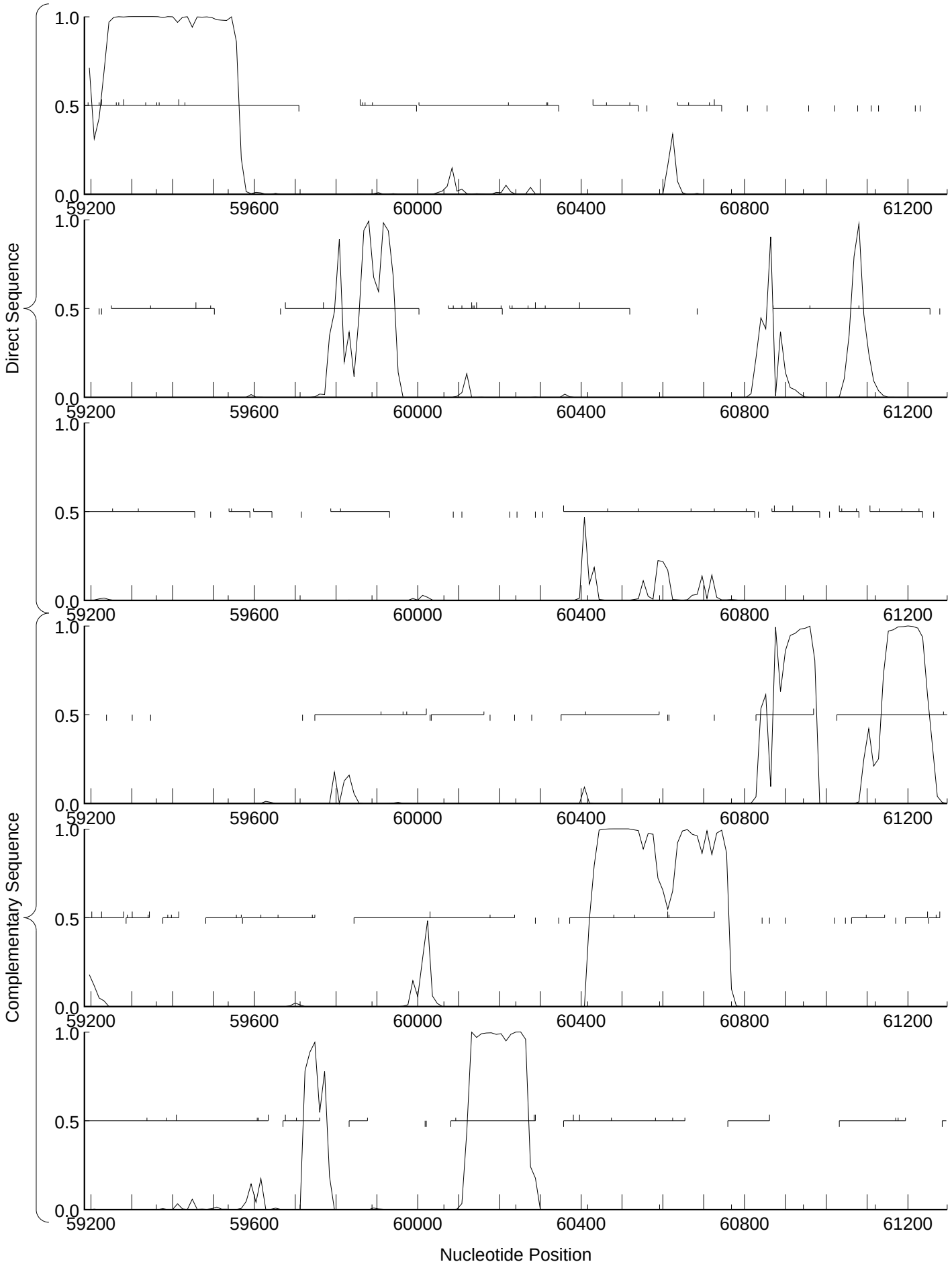

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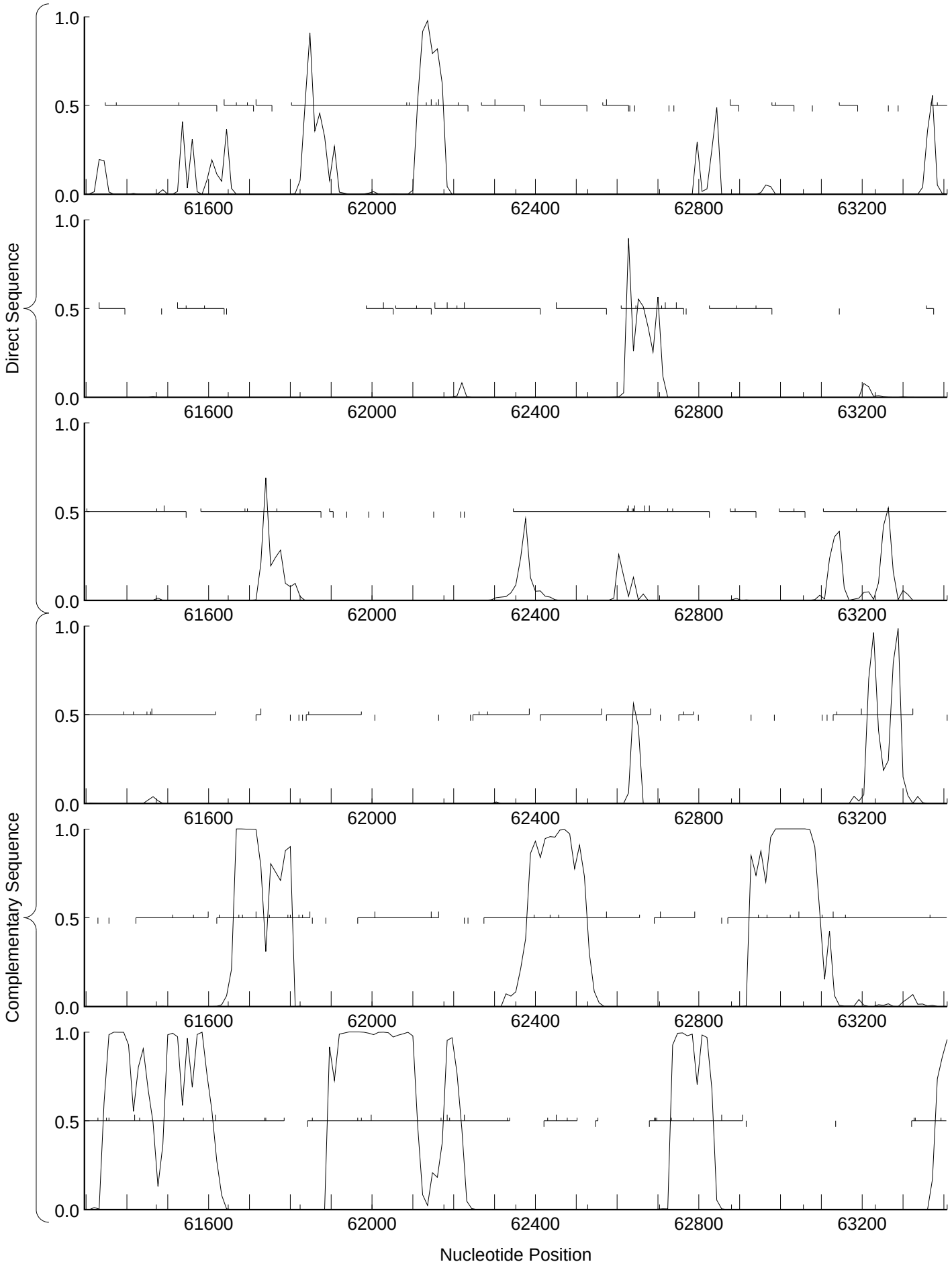

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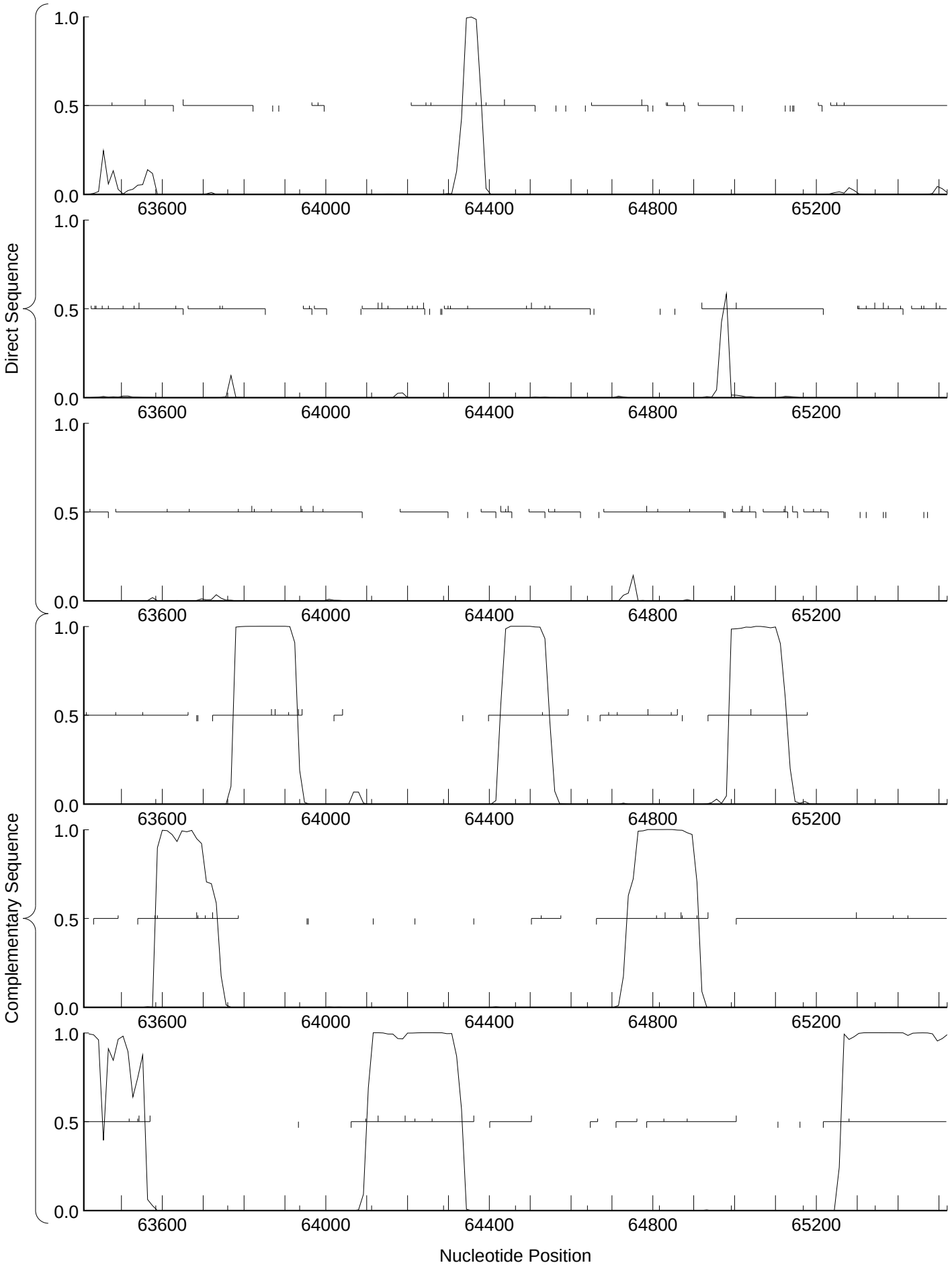

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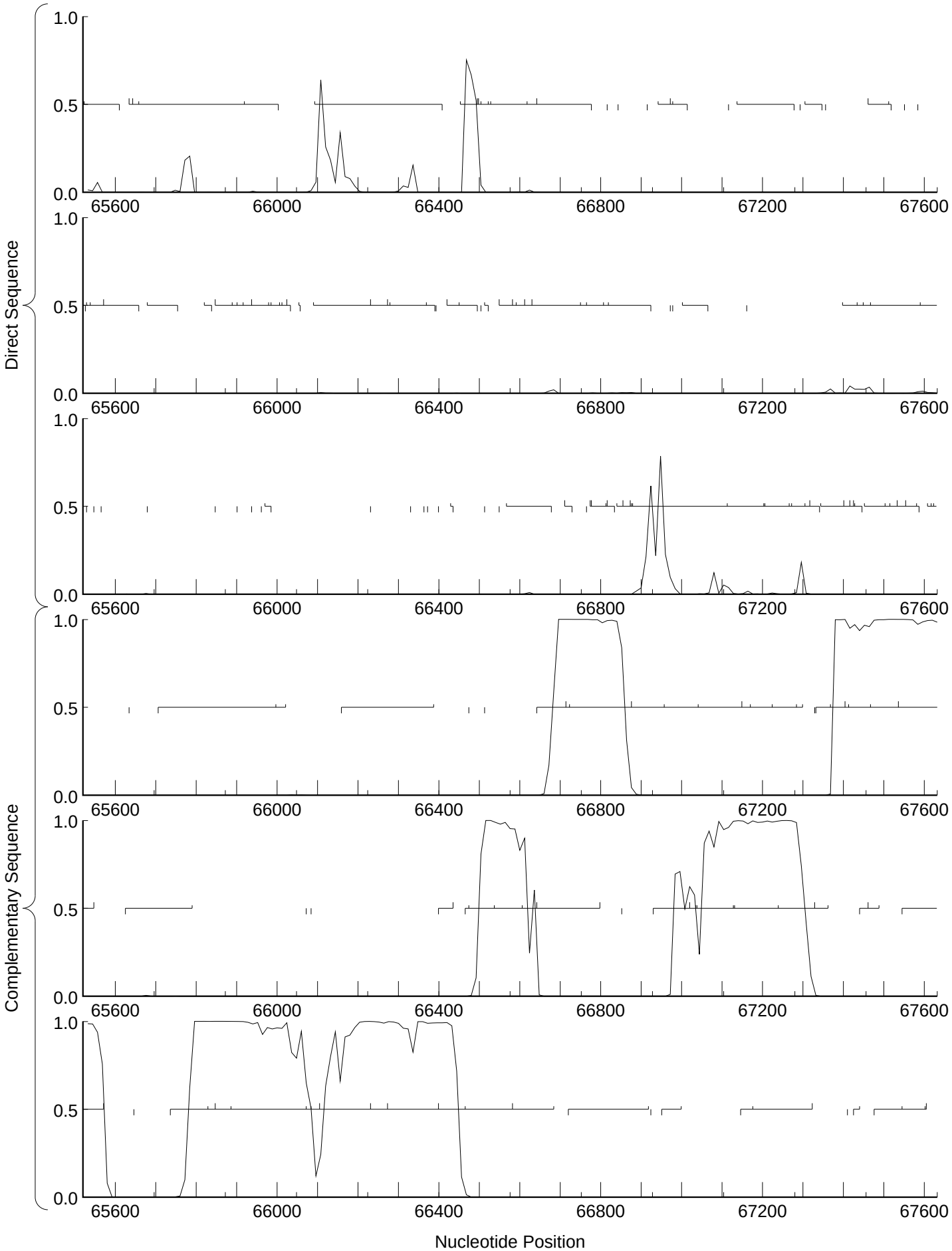

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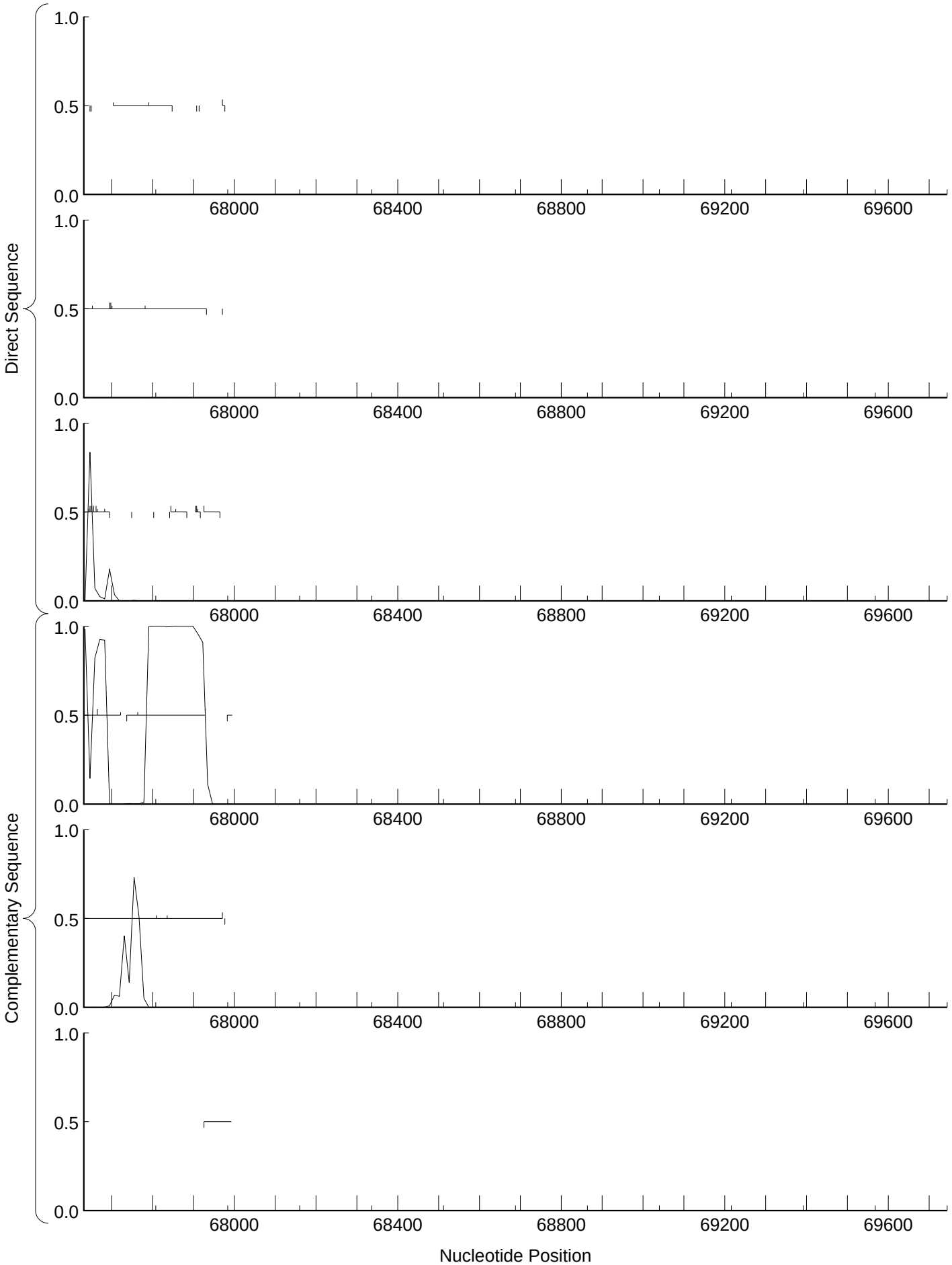

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Nucleotide Position


Direct Sequence
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60800
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Nucleotide Position


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63200
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Direct Sequence
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64800
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Nucleotide Position


Direct Sequence
Complementary Sequence
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66000
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67200
67600
Nucleotide Position


Direct Sequence
Complementary Sequence
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