GeneMark

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

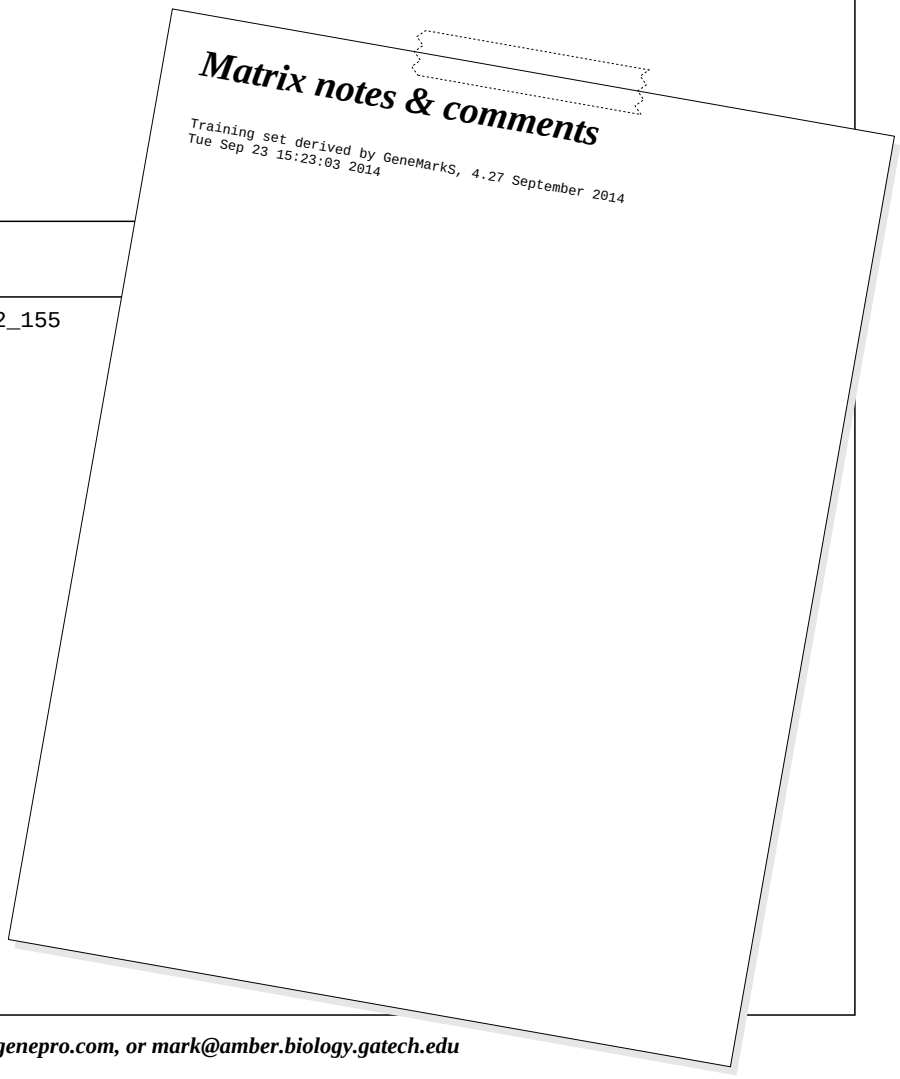
PROGRAM INFORMATION

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Analysis Date    :  3/28/24 at 5:03:48
Pages            :  34
Sequence Length  :  68568 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,
                    List ORFs,
                    List regions and/or splice sites,
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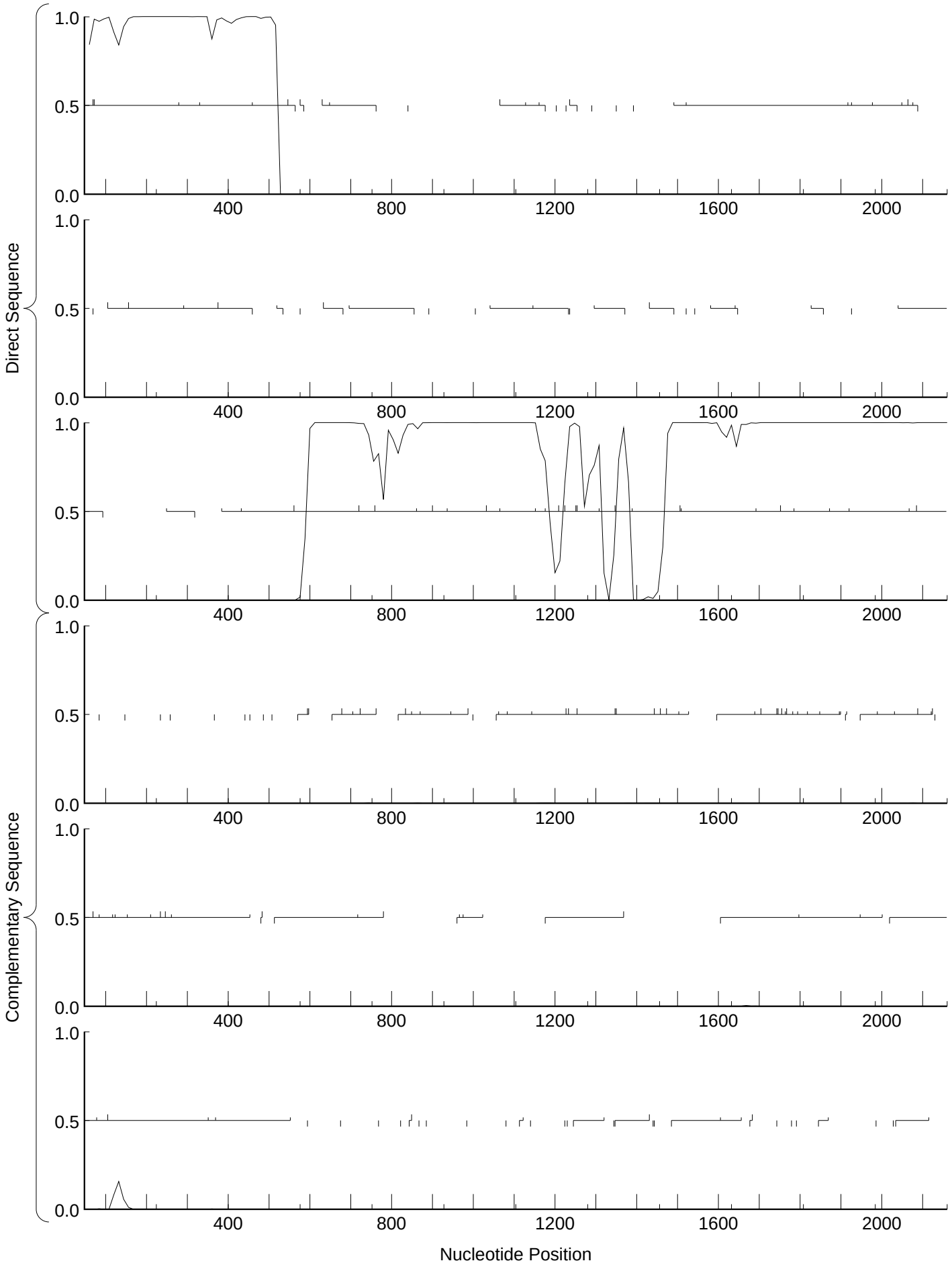
MATRIX INFORMATION

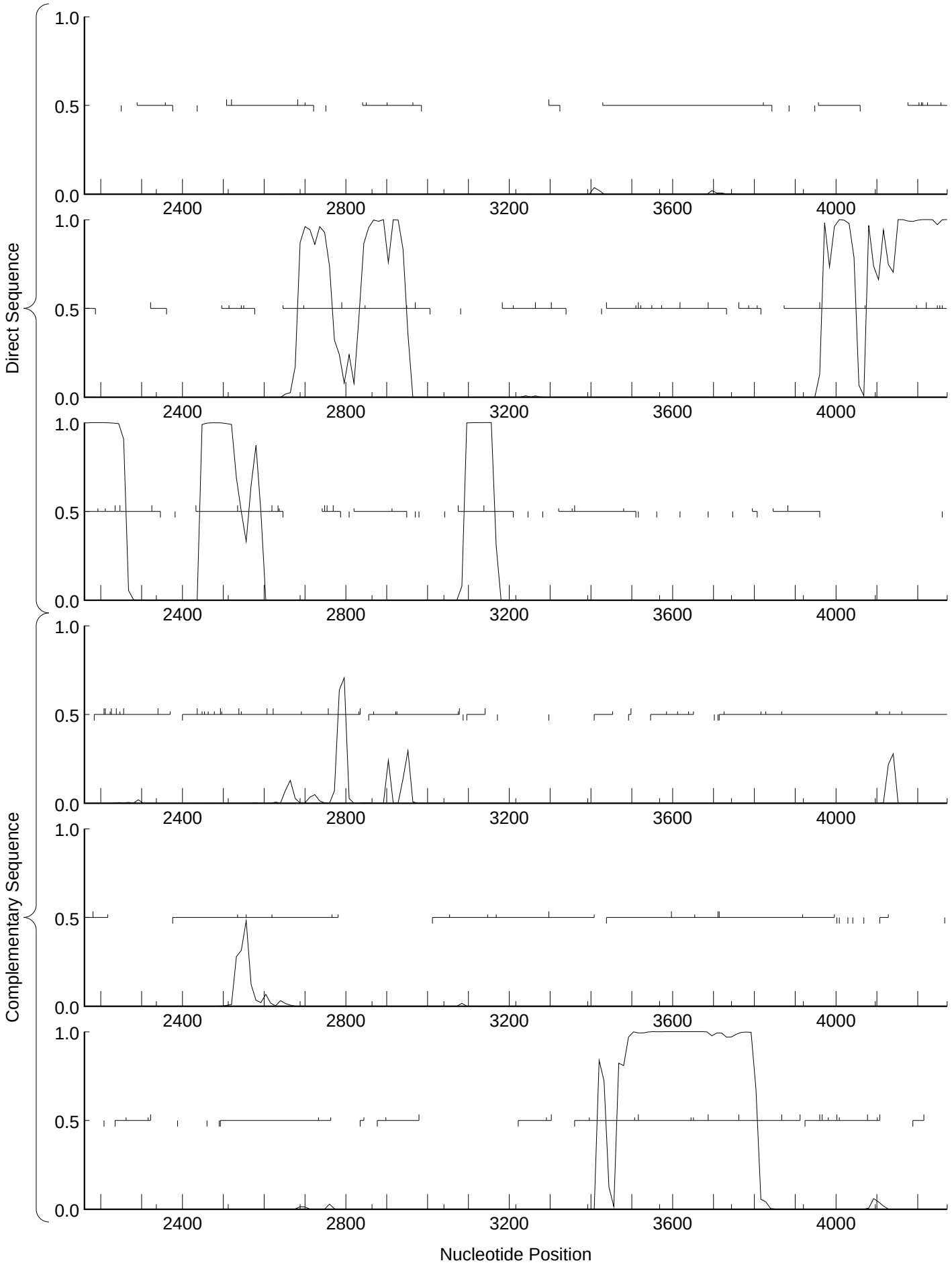
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Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

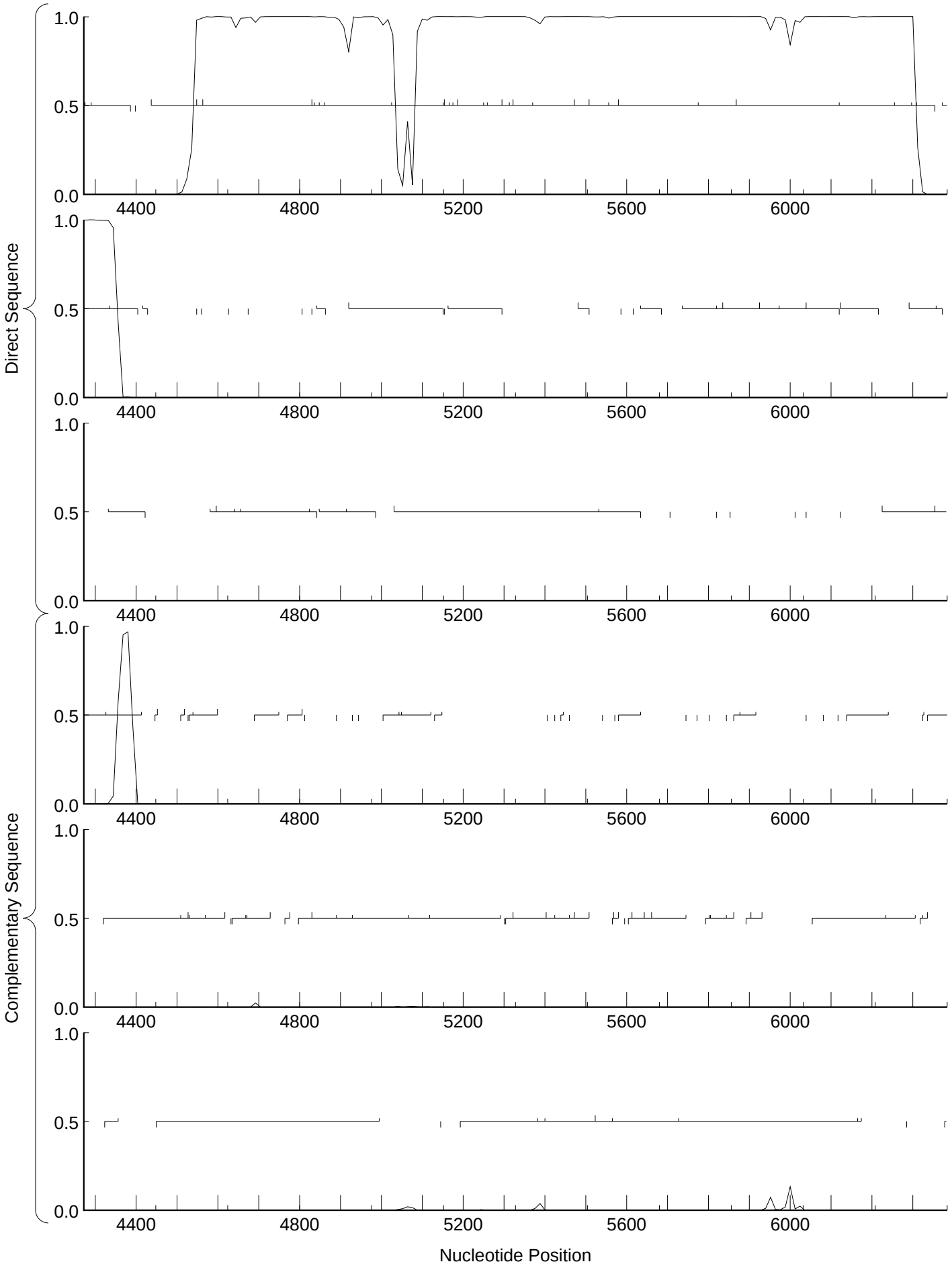
Order      :  4
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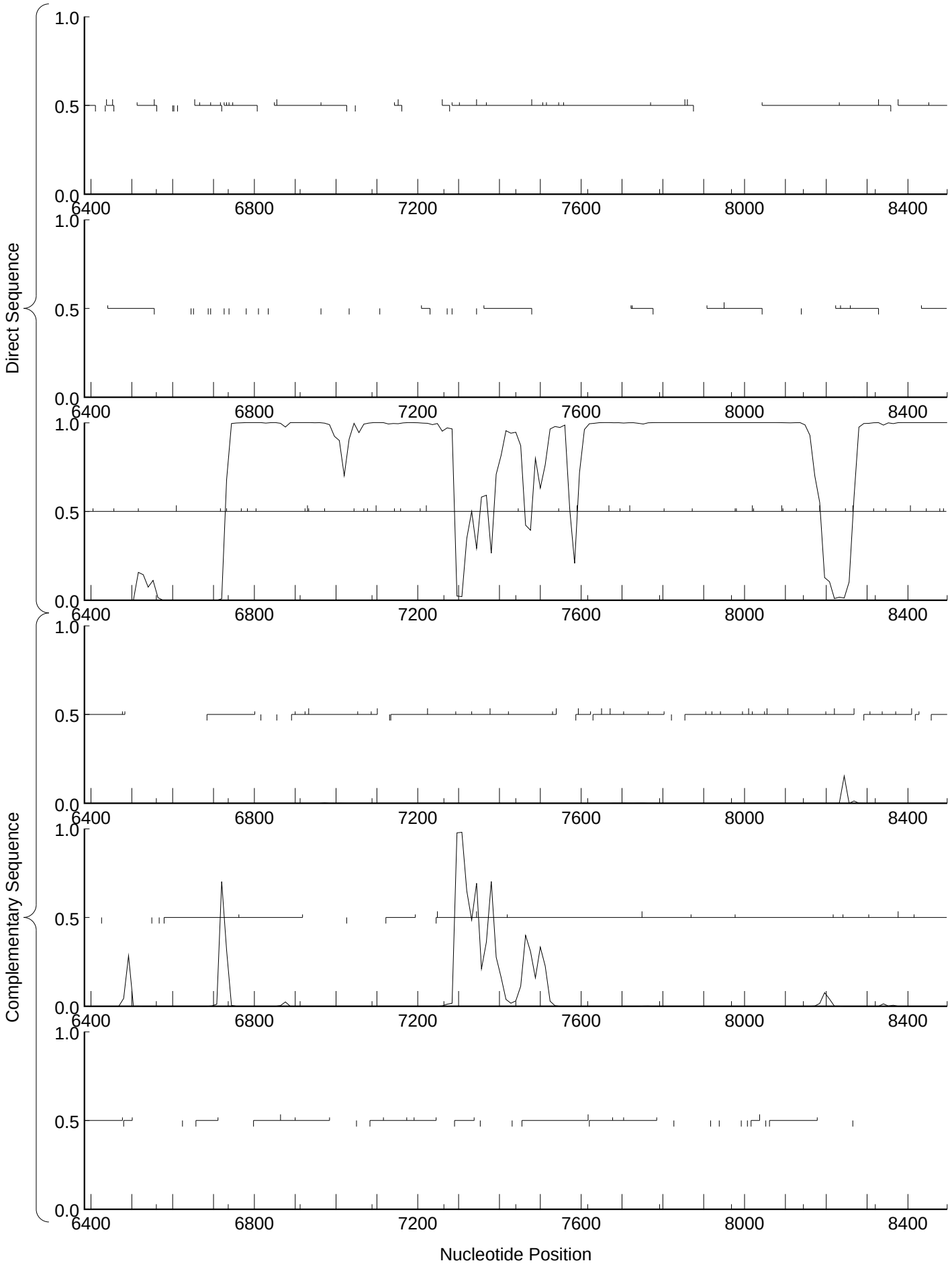

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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

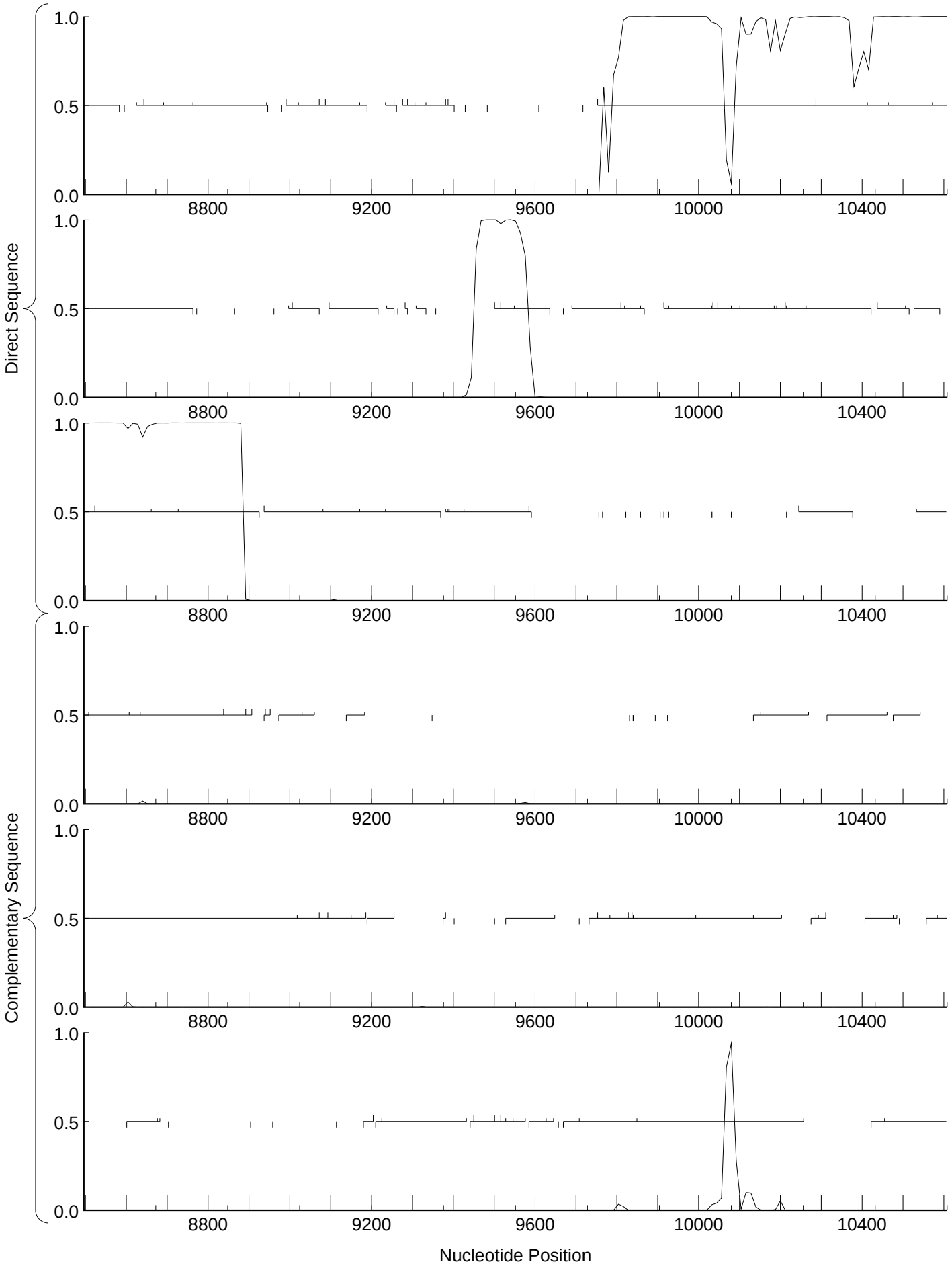

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

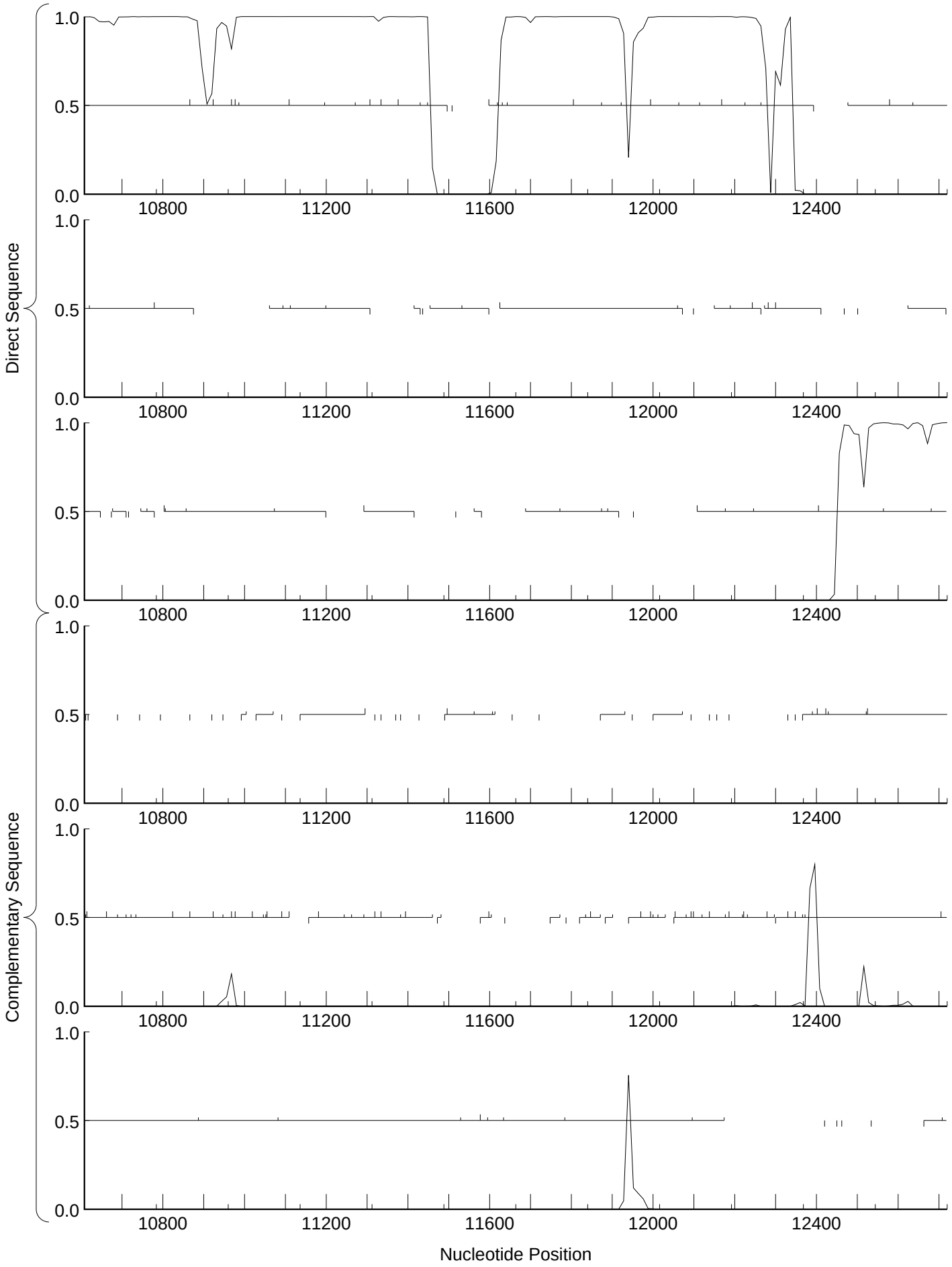

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1200
1600
2000
Nucleotide Position

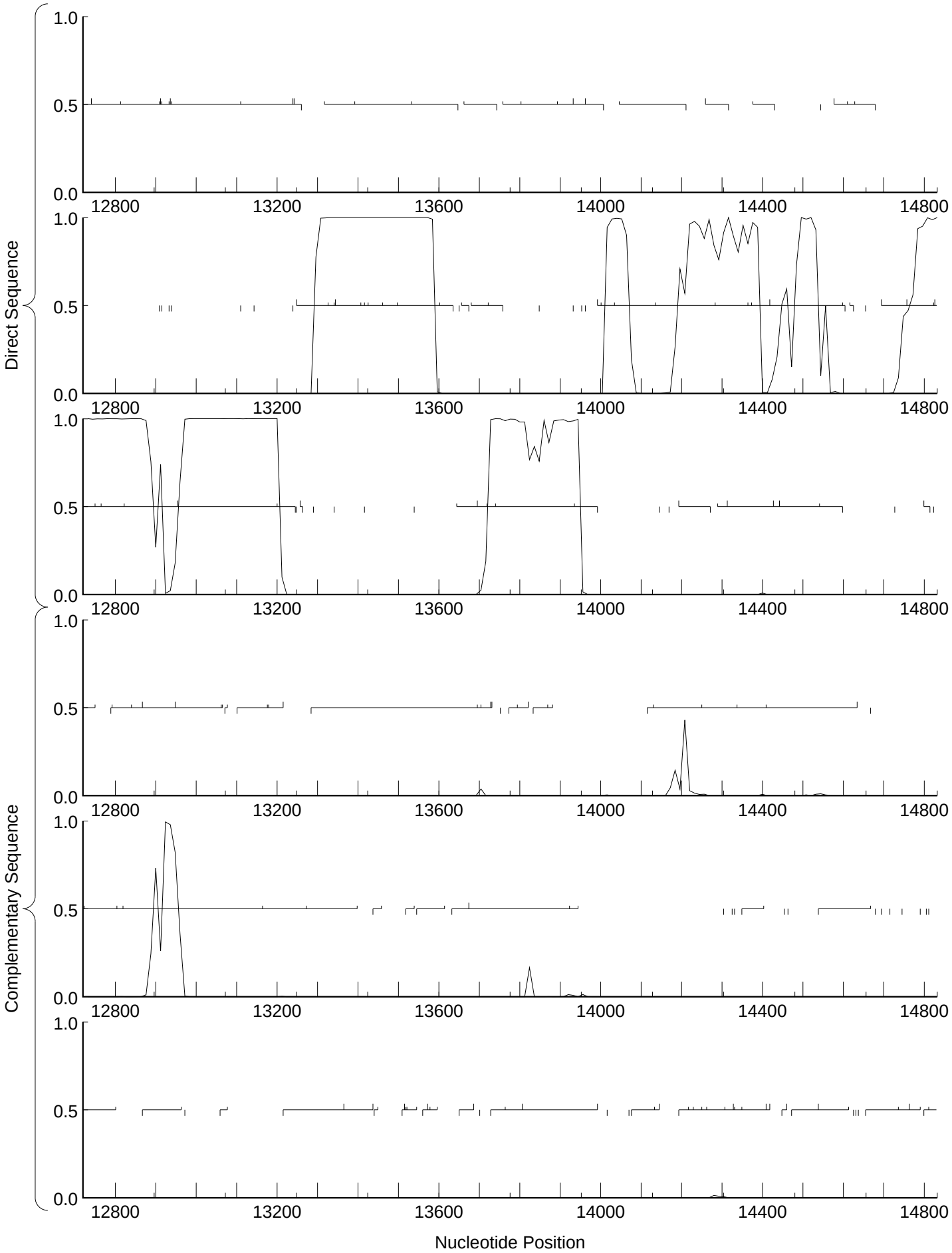

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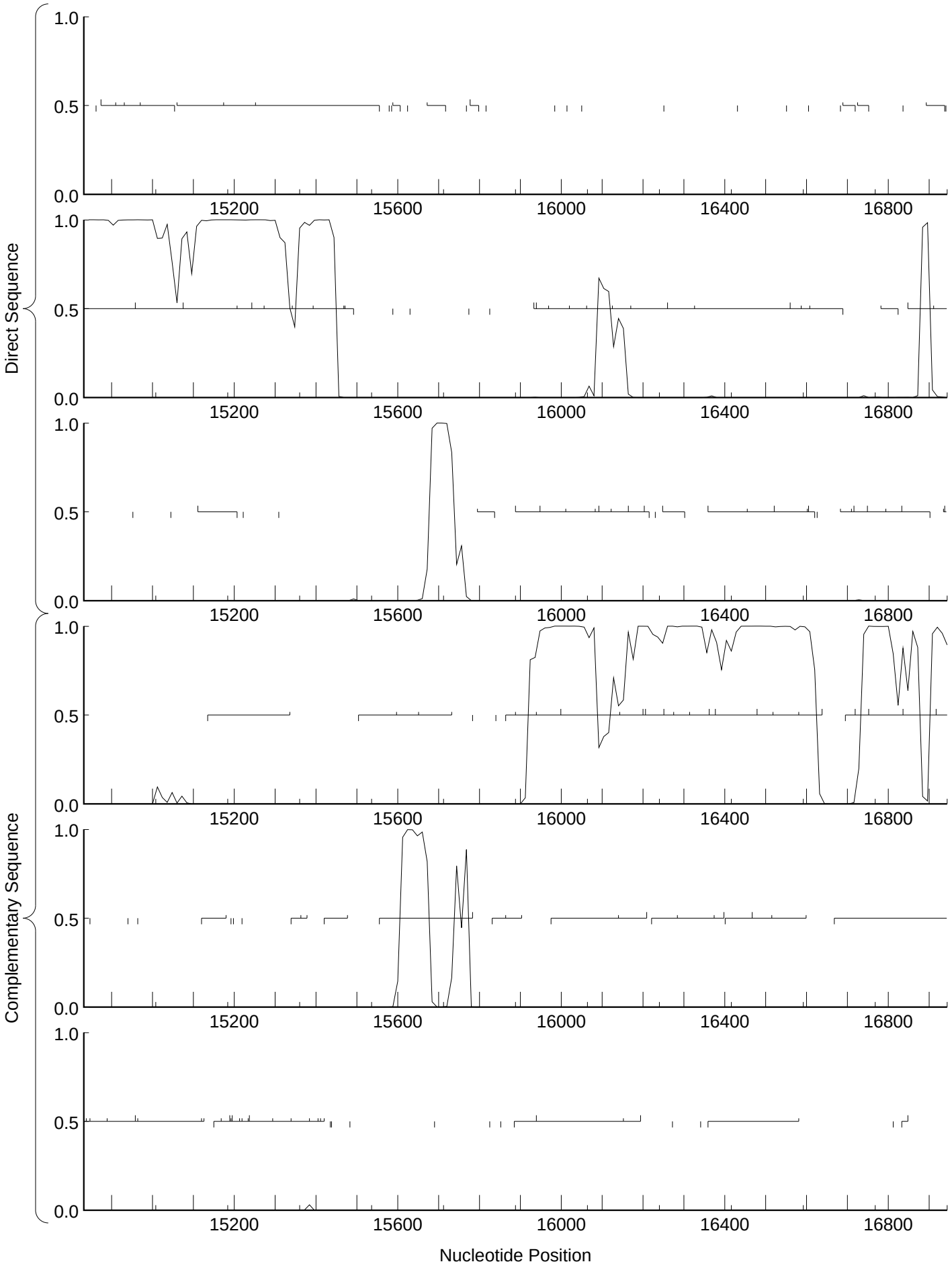

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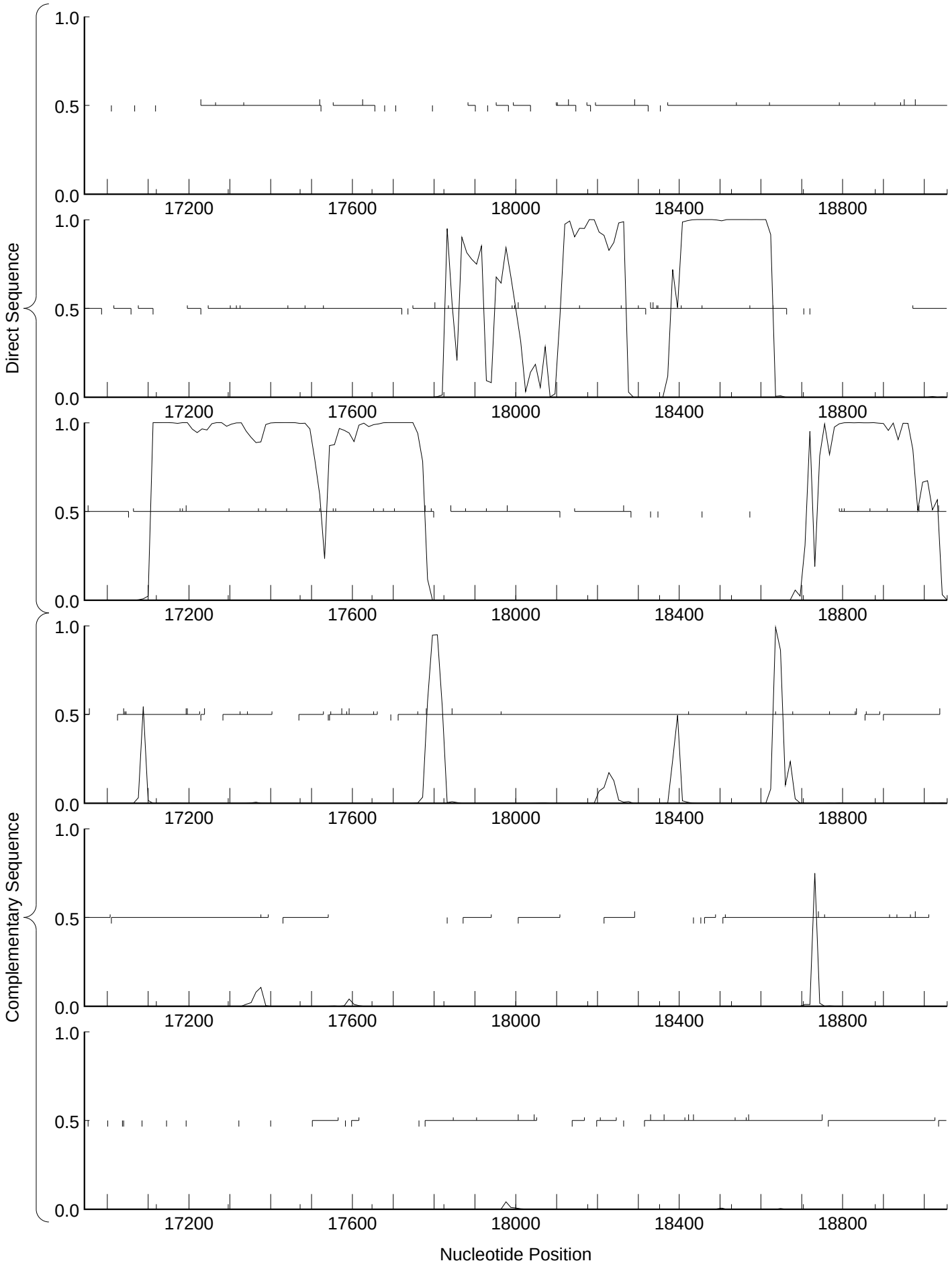

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Nucleotide Position

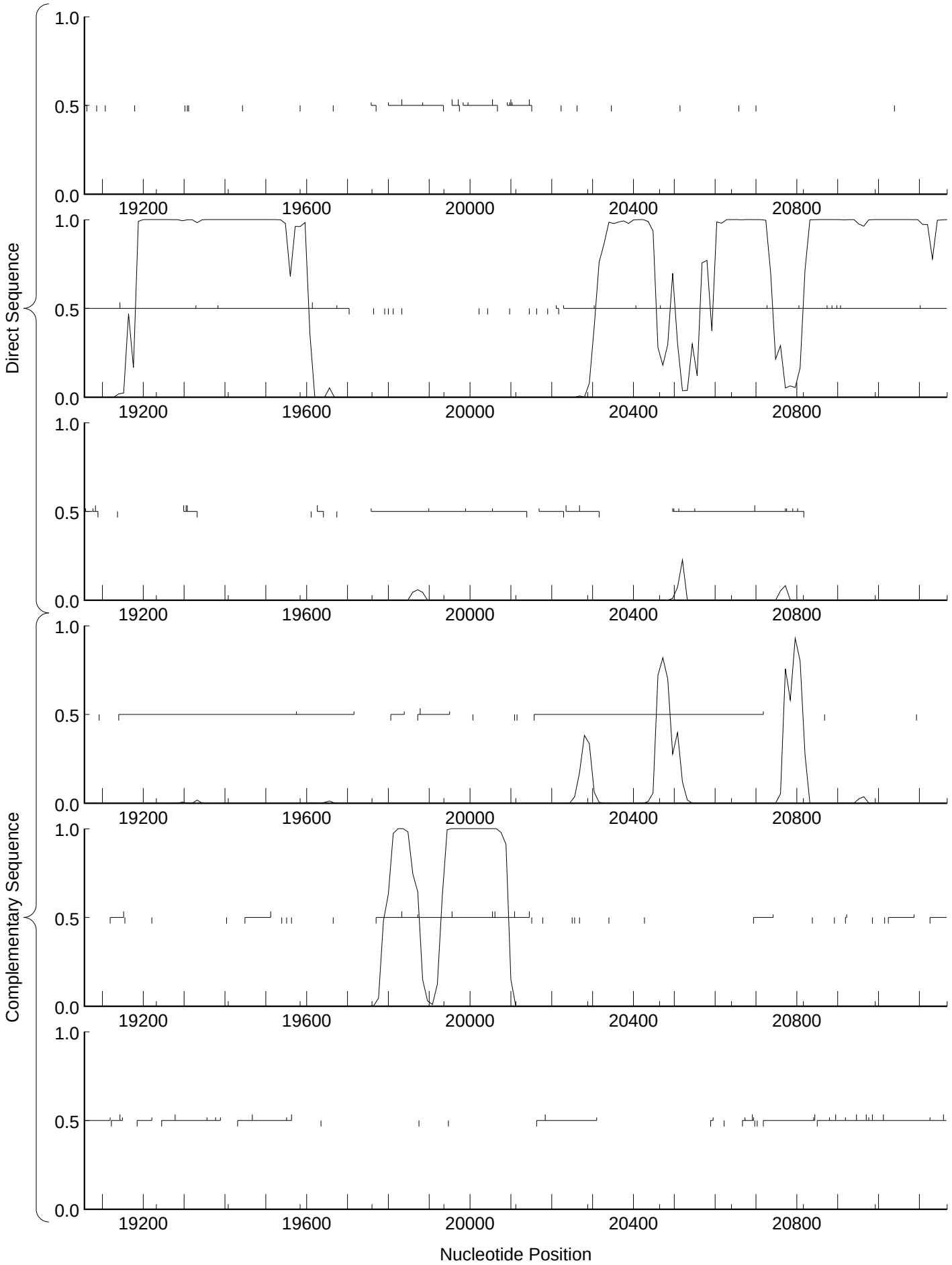

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11200
11600
12000
12400
Nucleotide Position


Direct Sequence
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13600
14000
14400
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Nucleotide Position

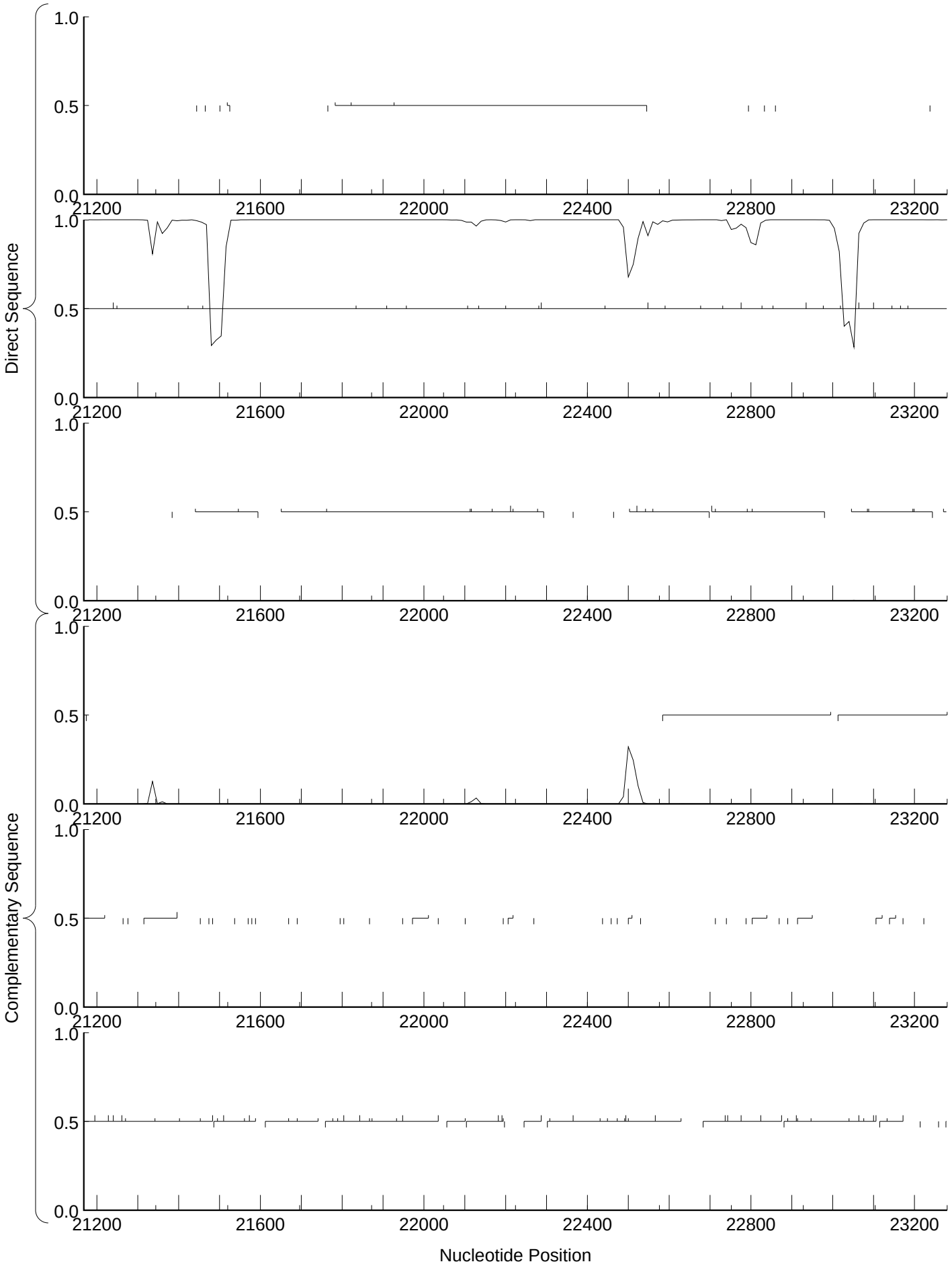

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16000
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Direct Sequence
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18000
18400
18800
Nucleotide Position

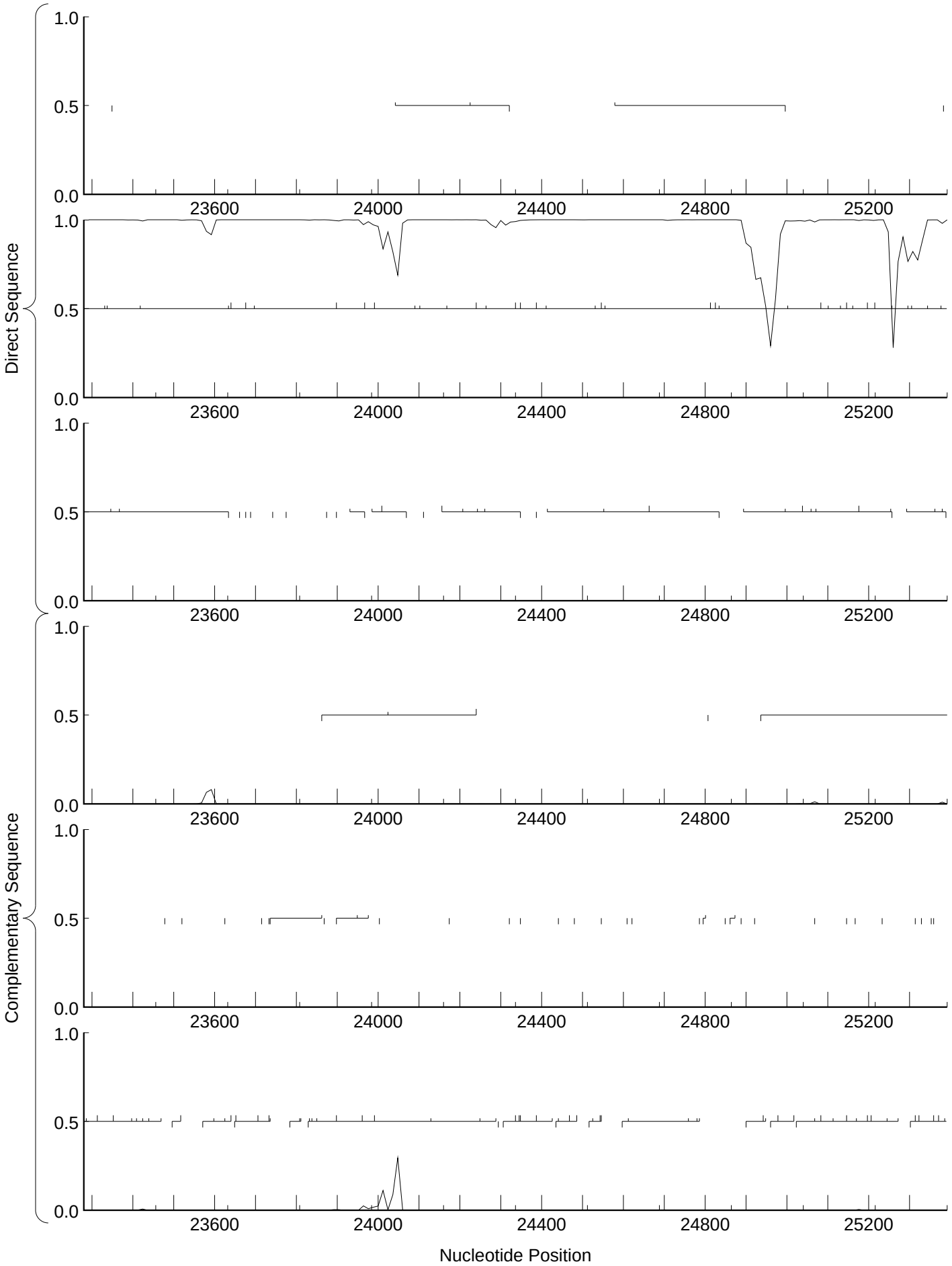
acterium phage Spartan300 Complete Sequence, 68568 bp, Circularly Permuted, Cluster B1, Order 4, Window 96, Step 12, 11/34



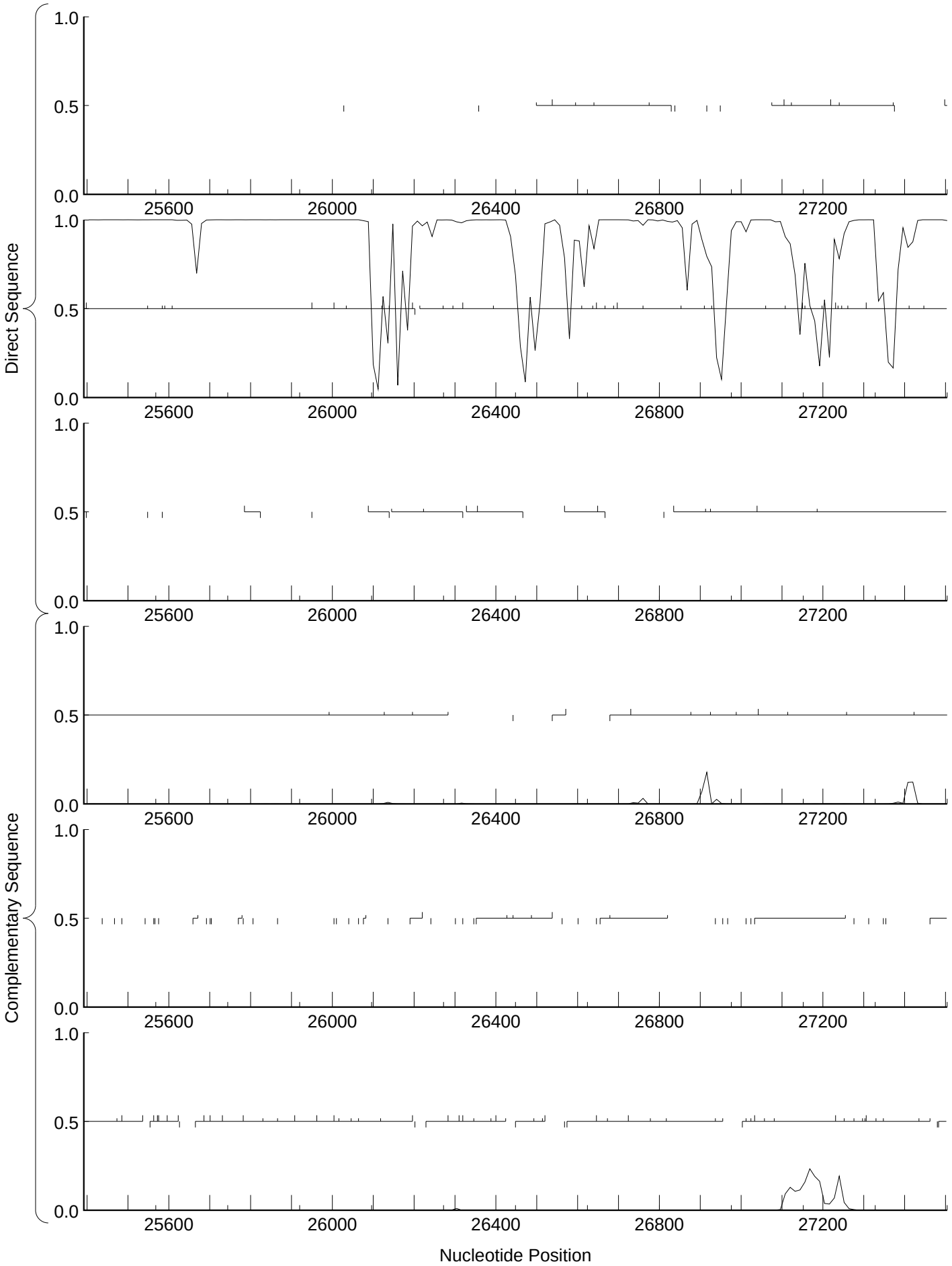
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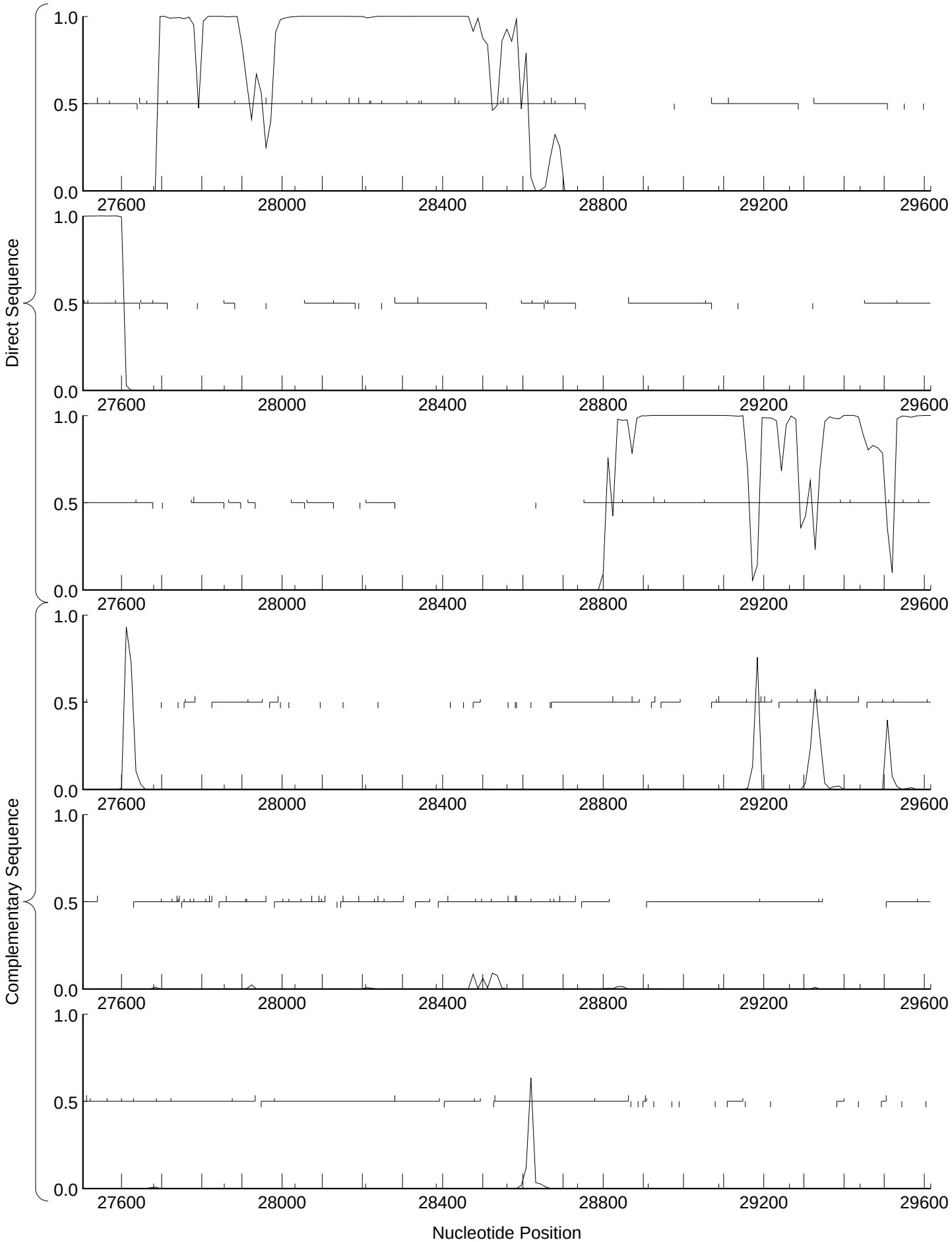
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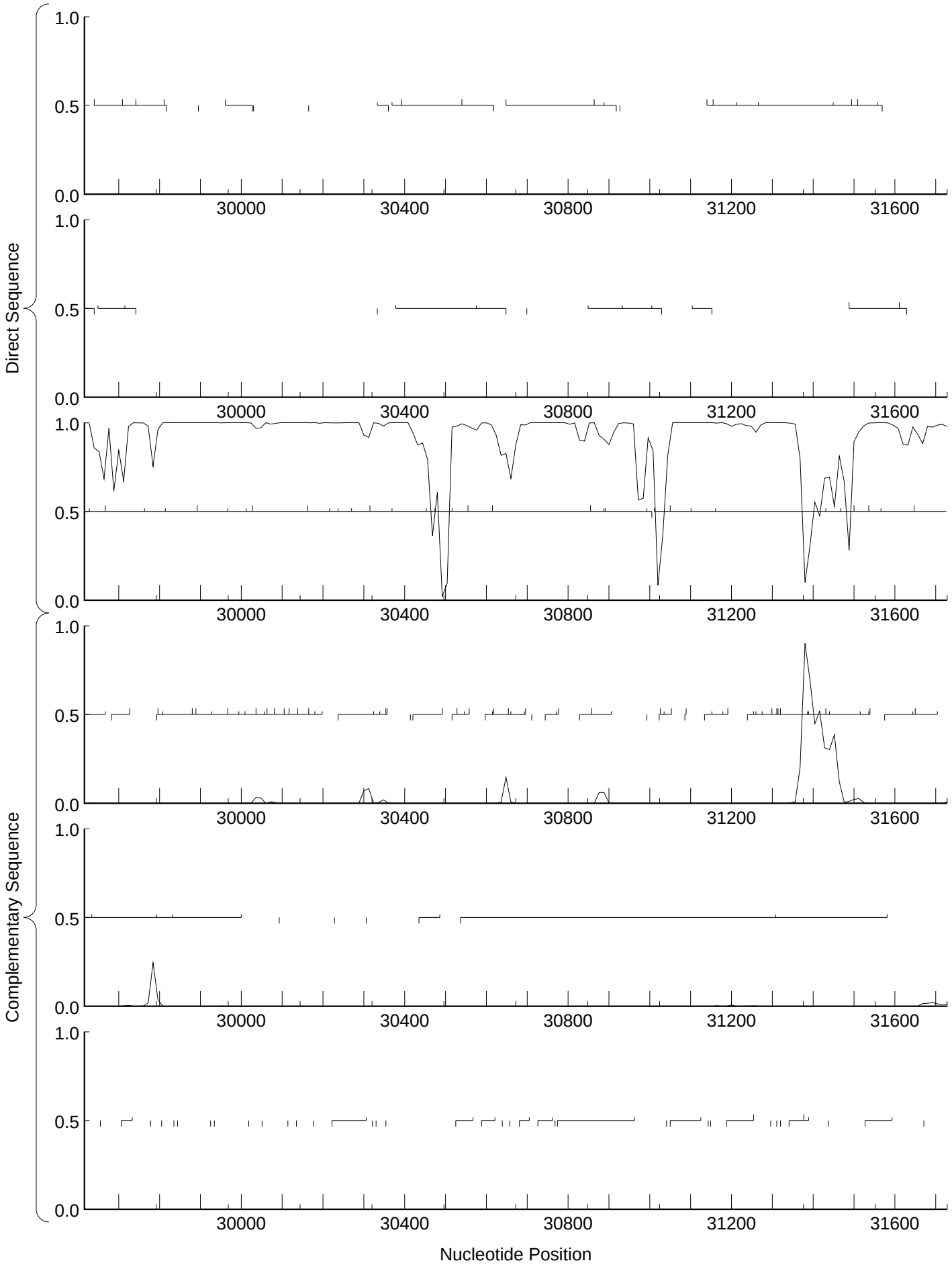
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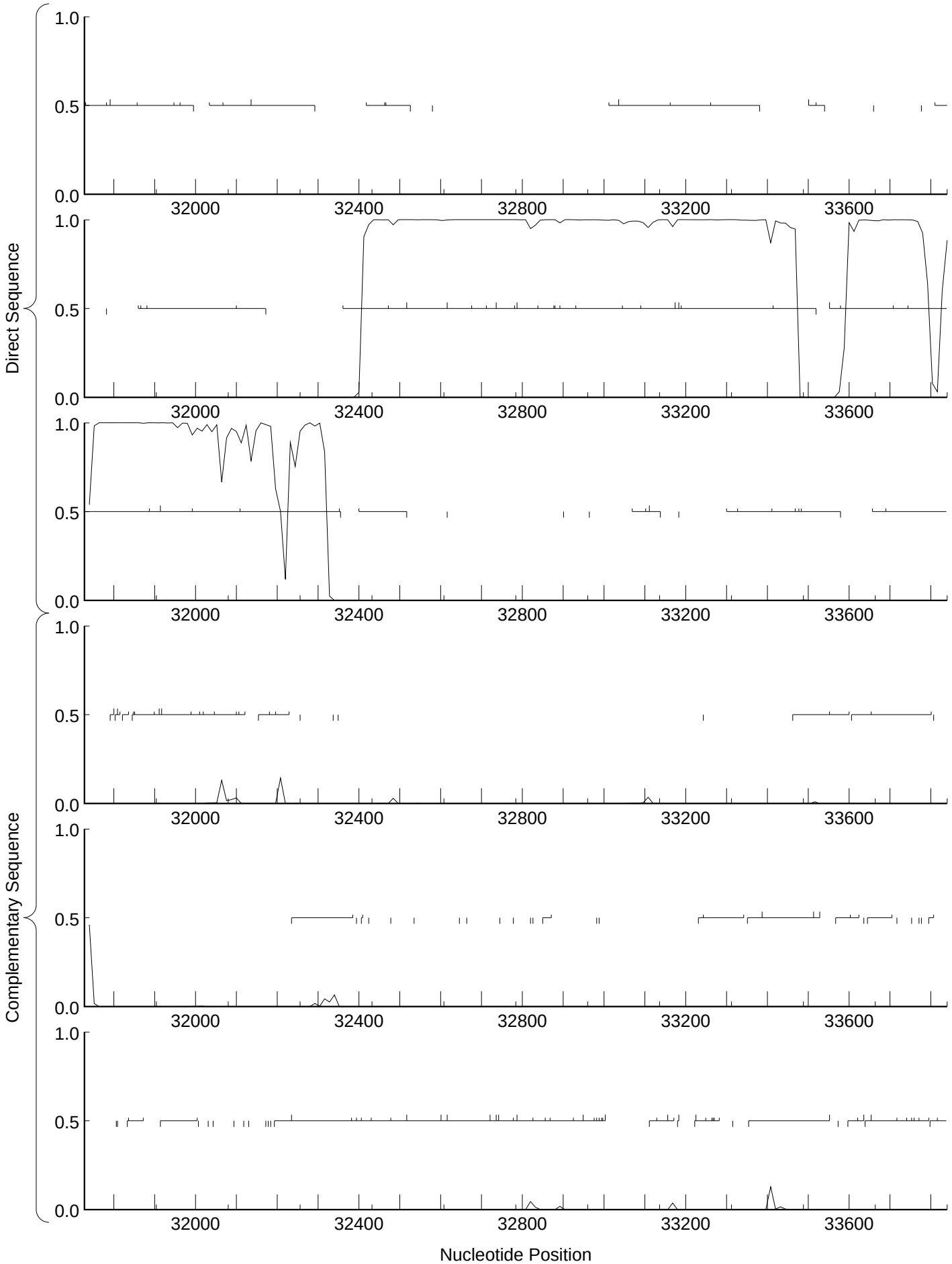
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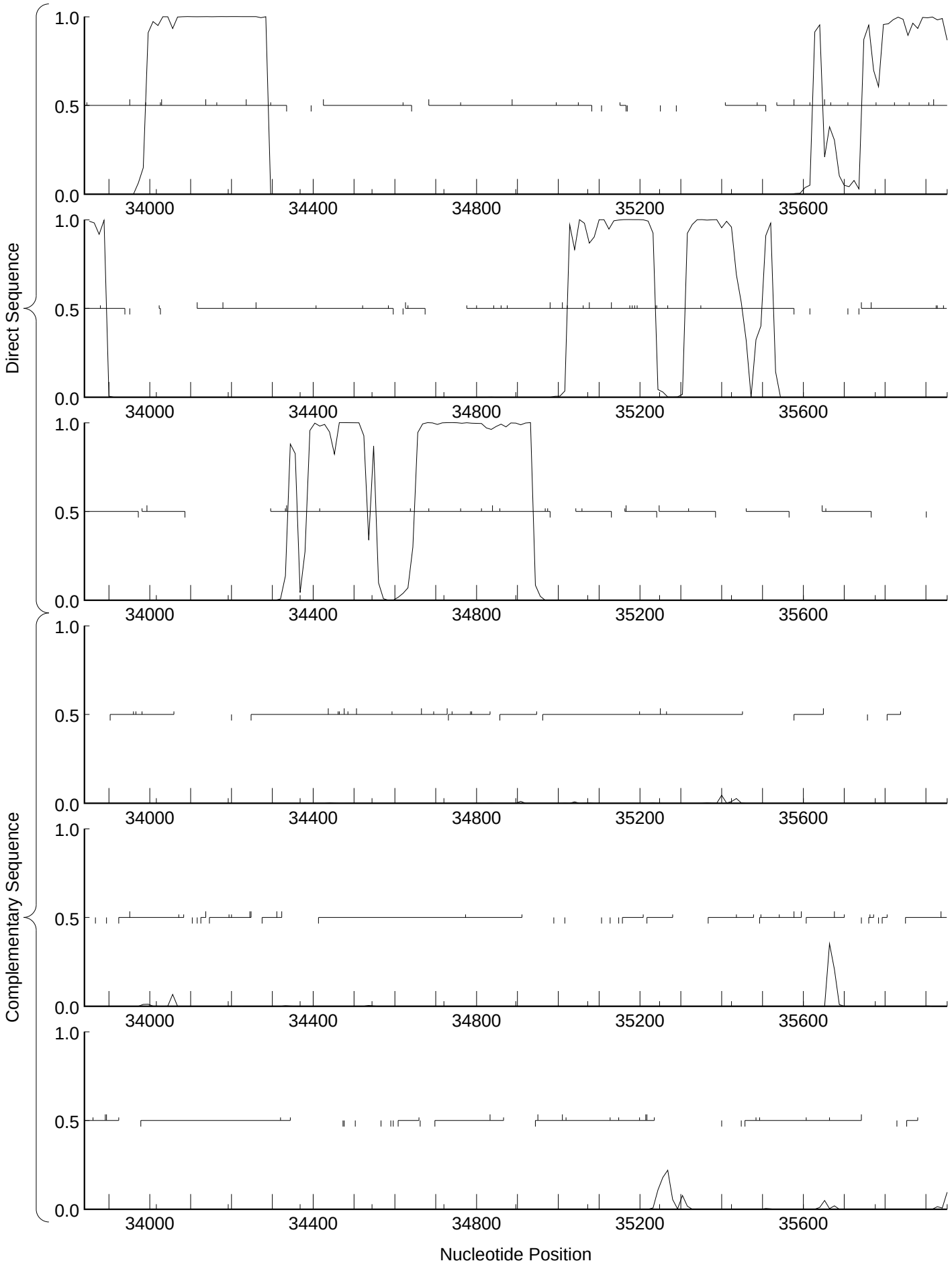
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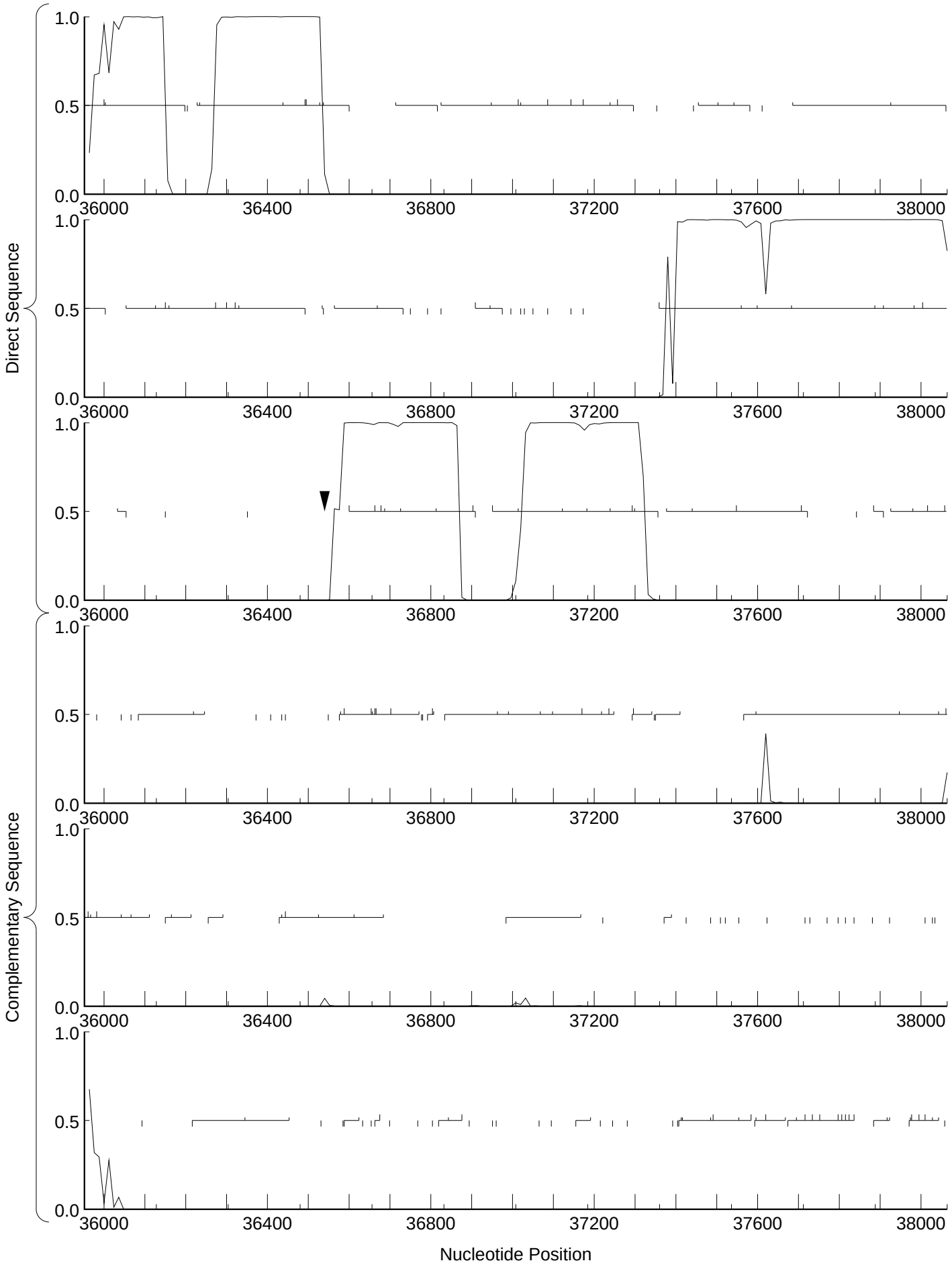
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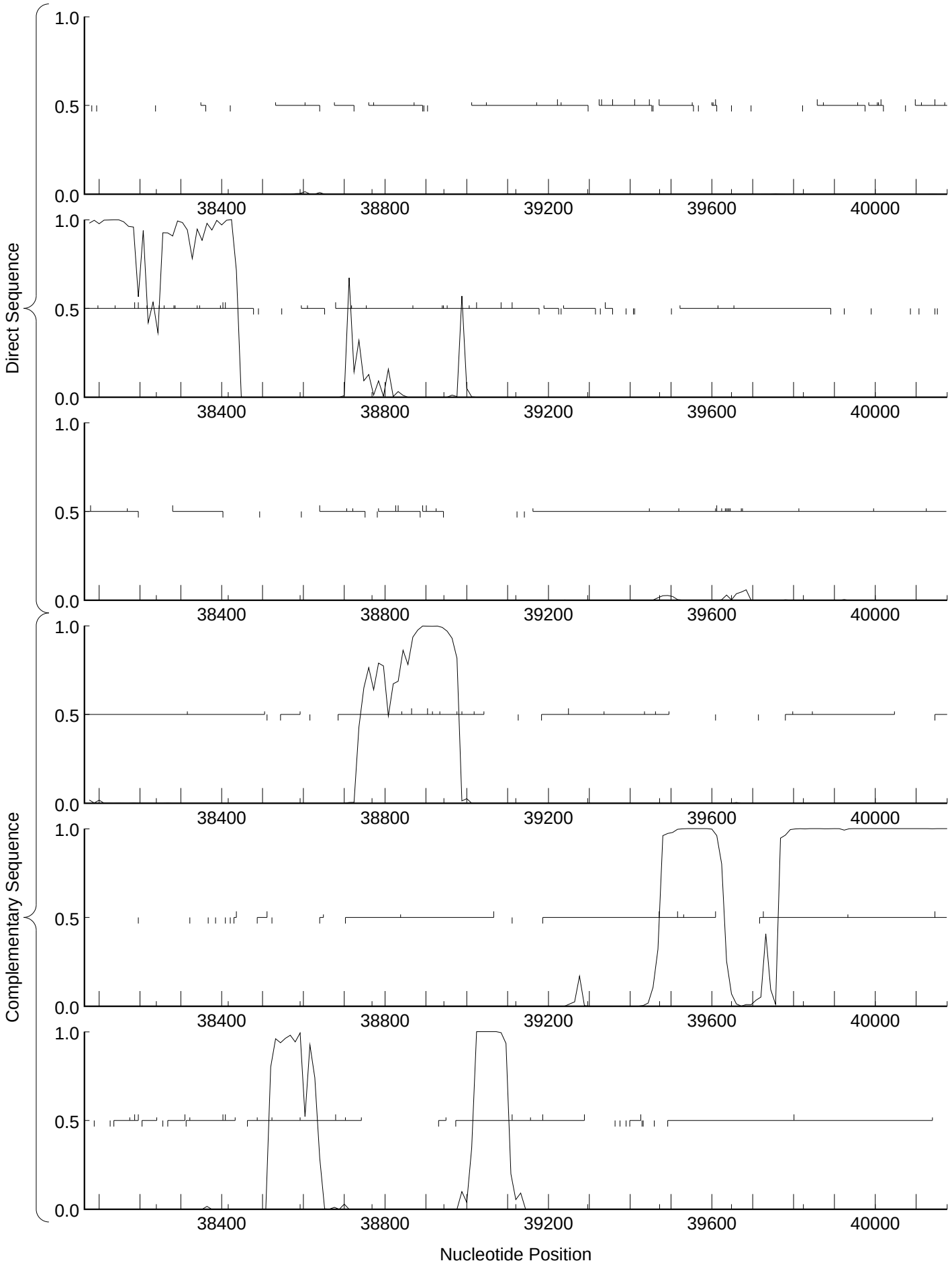


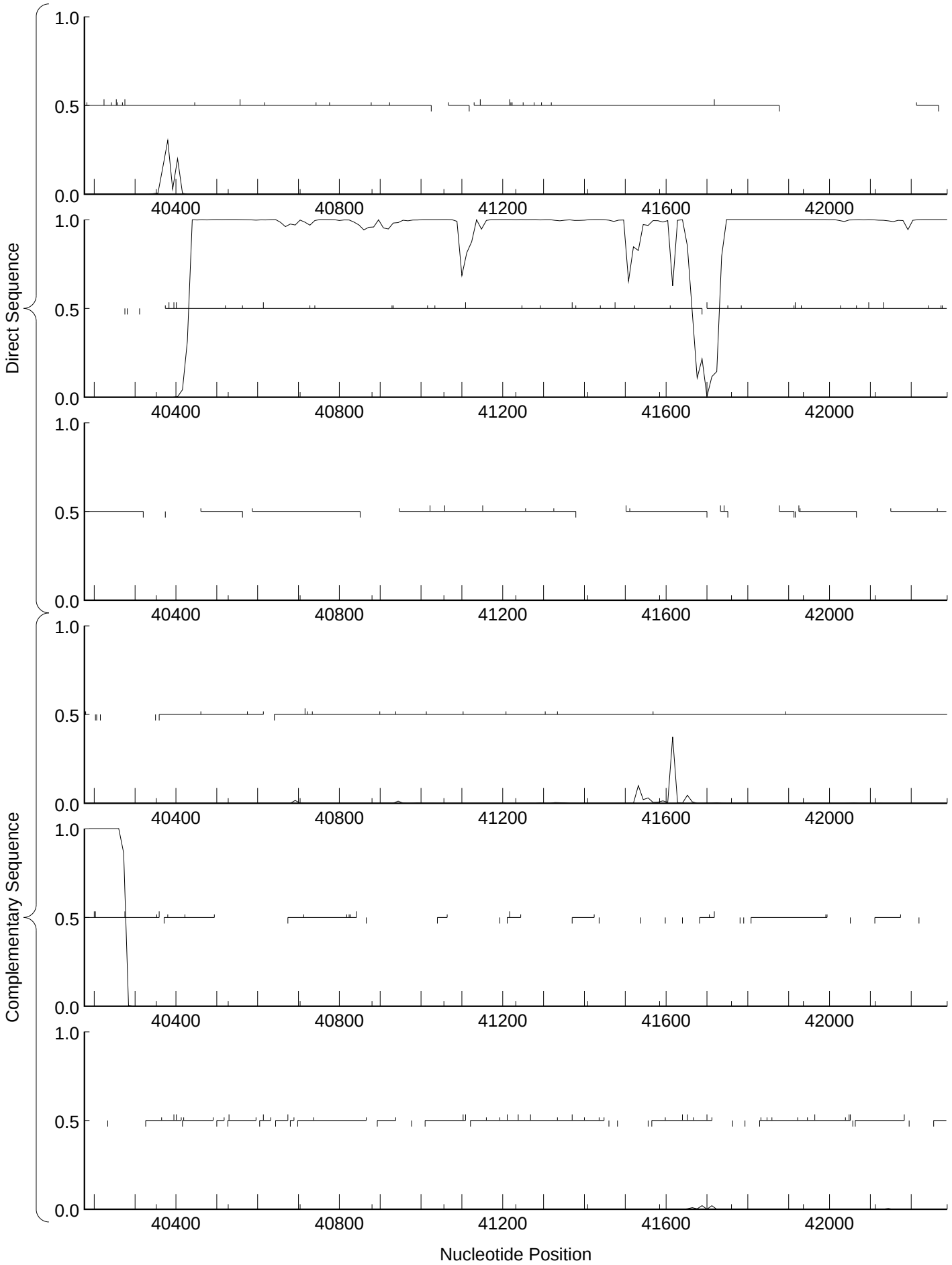
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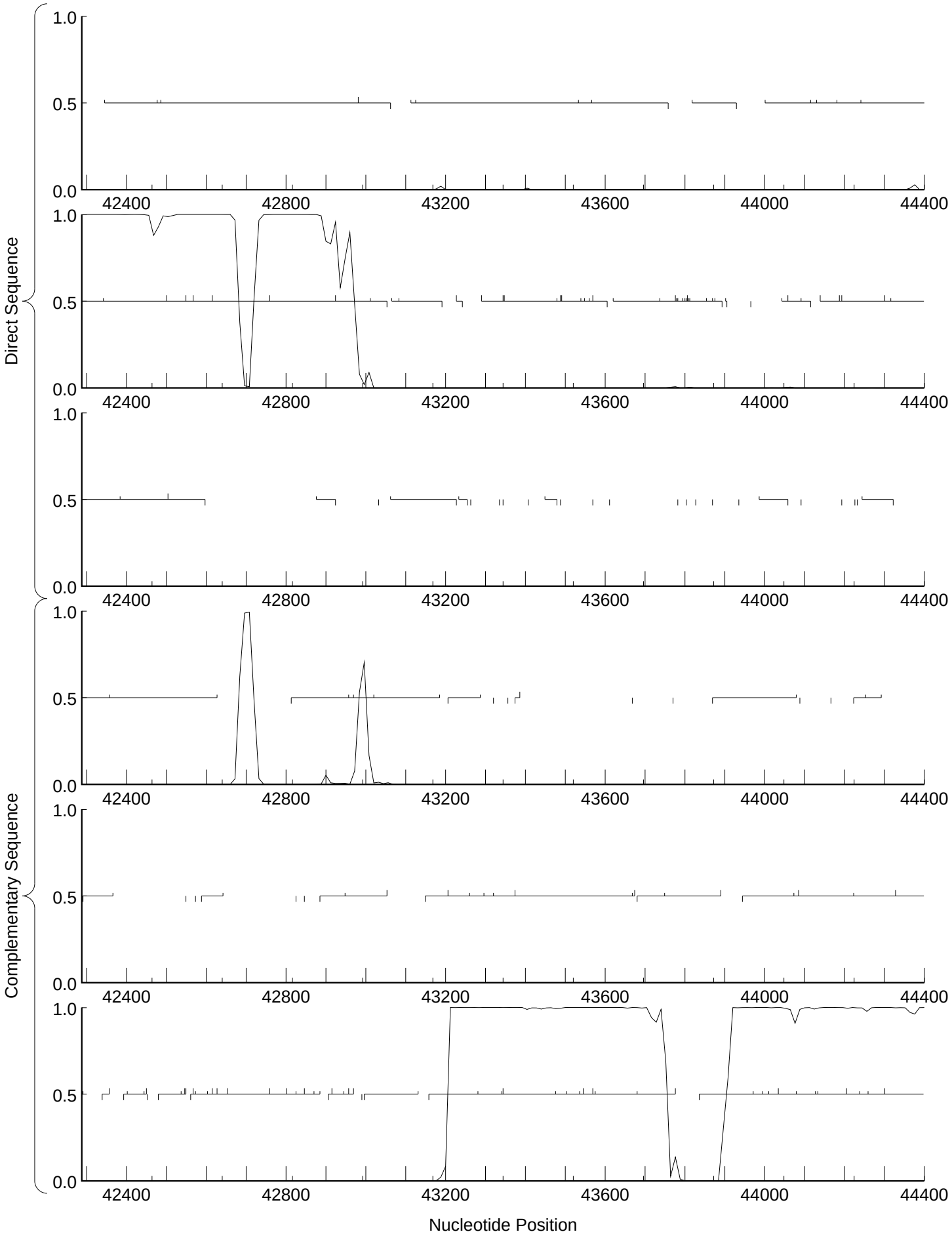
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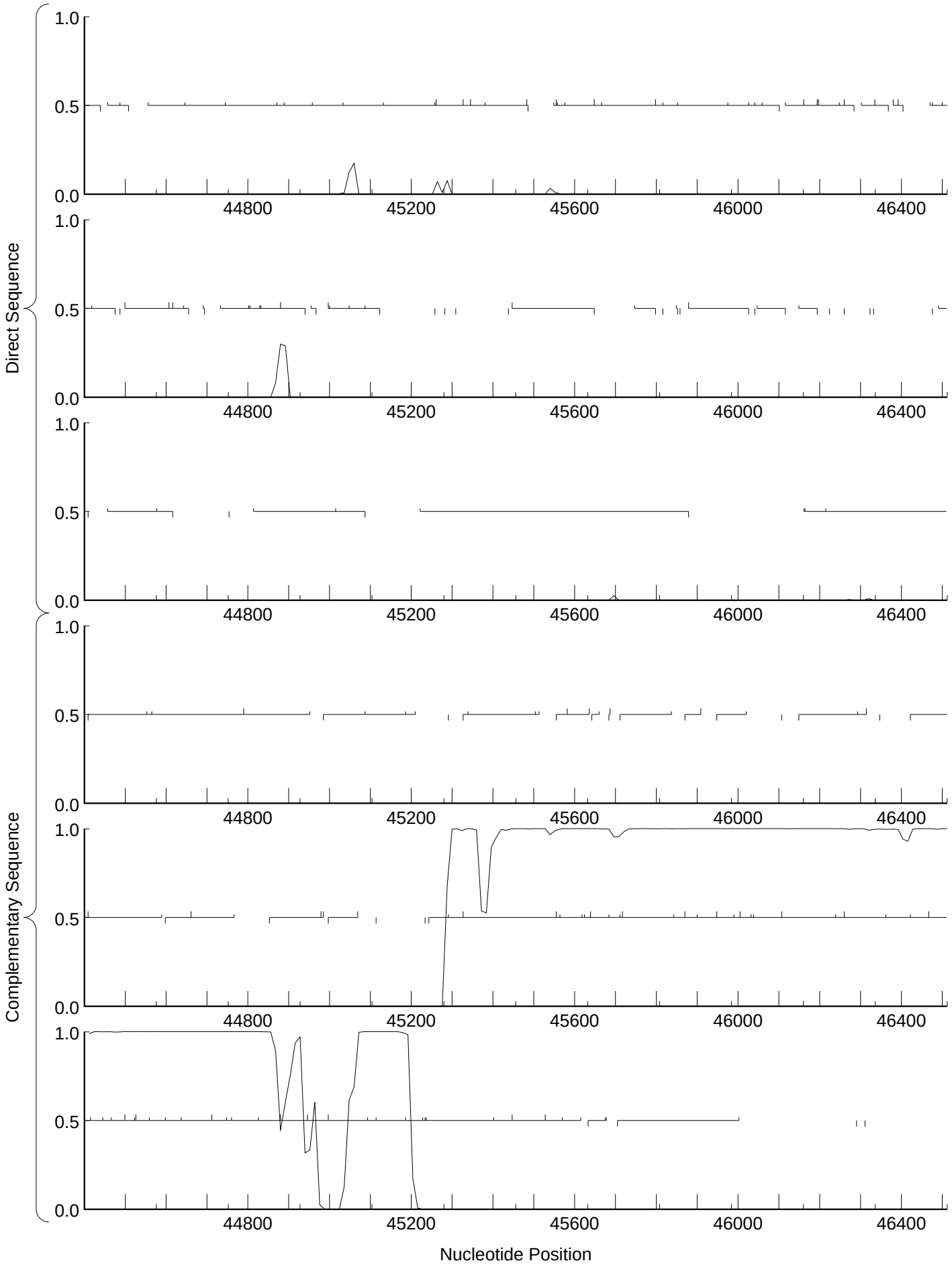

Direct Sequence
Complementary Sequence
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0.5
0.0
38400
38800
39200
39600
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Nucleotide Position


Direct Sequence
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40800
41200
41600
42000
Nucleotide Position

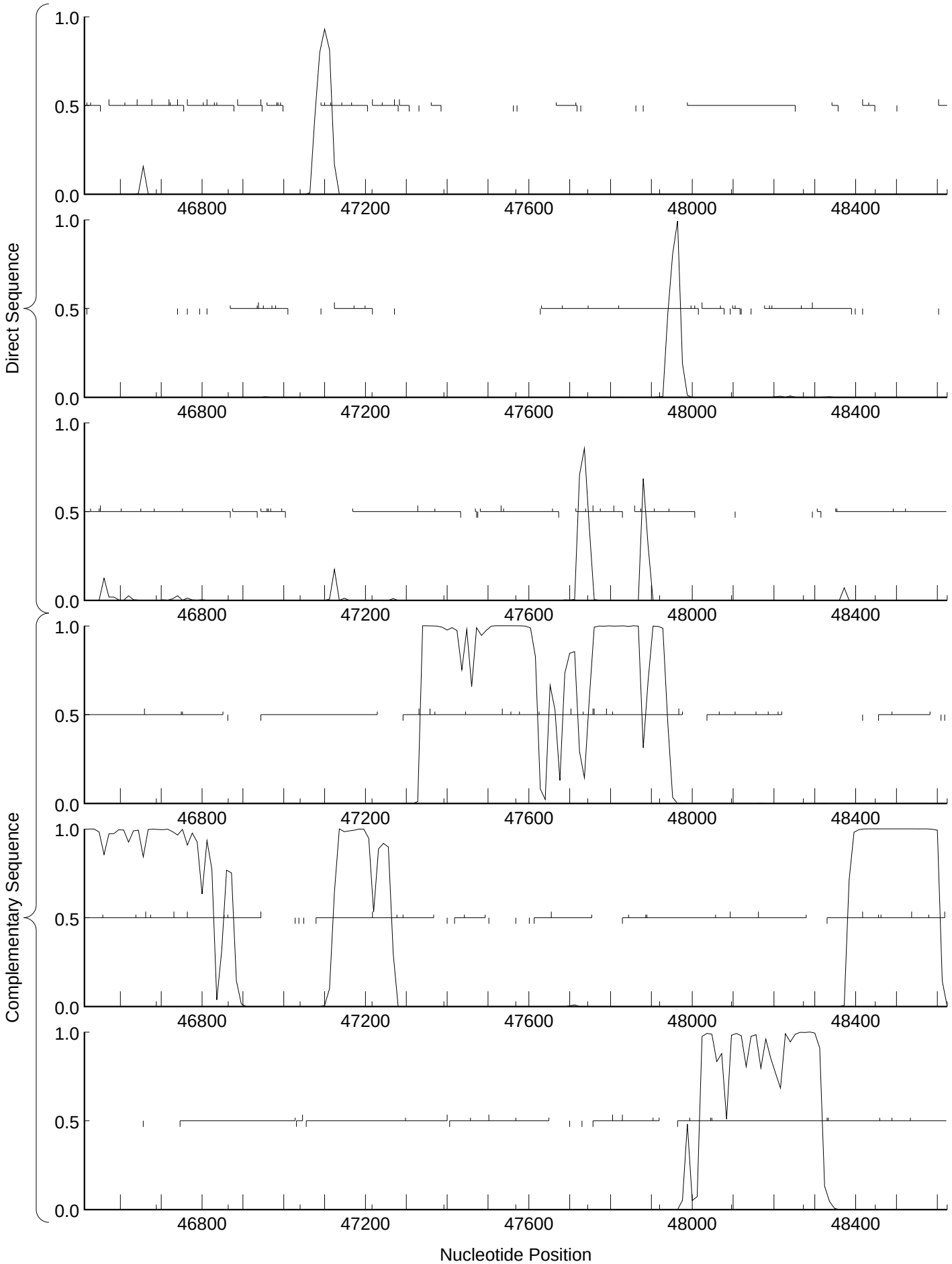
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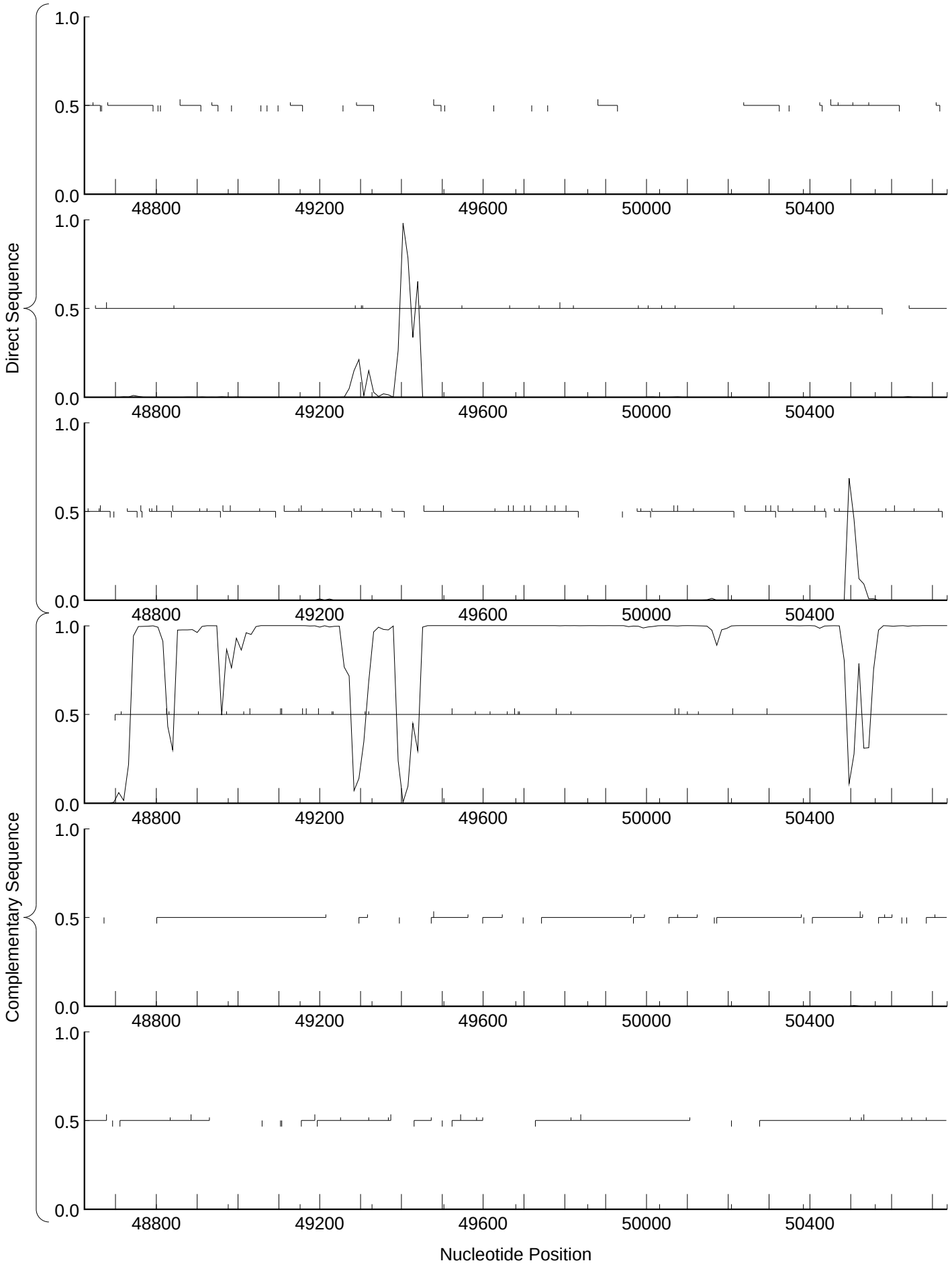
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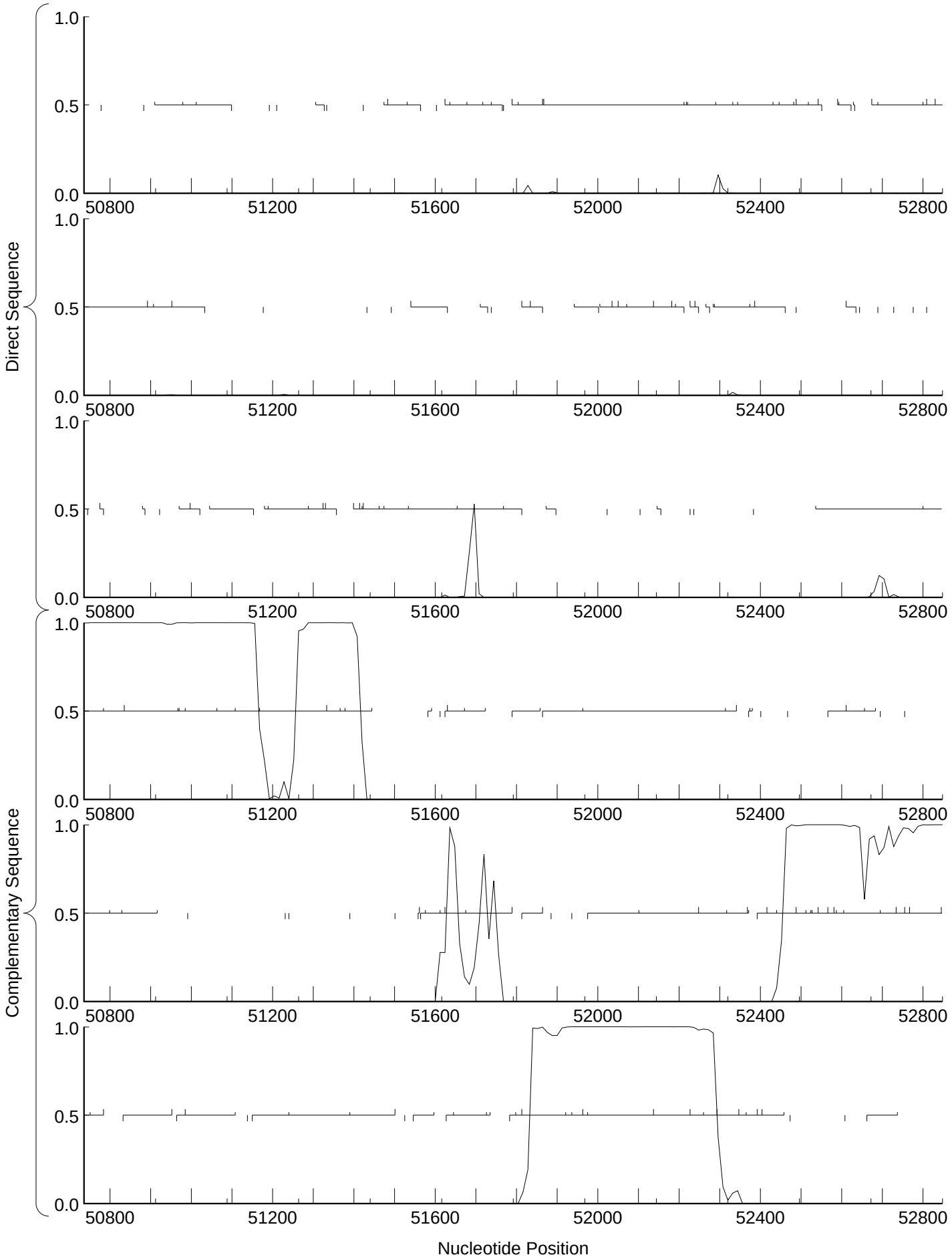


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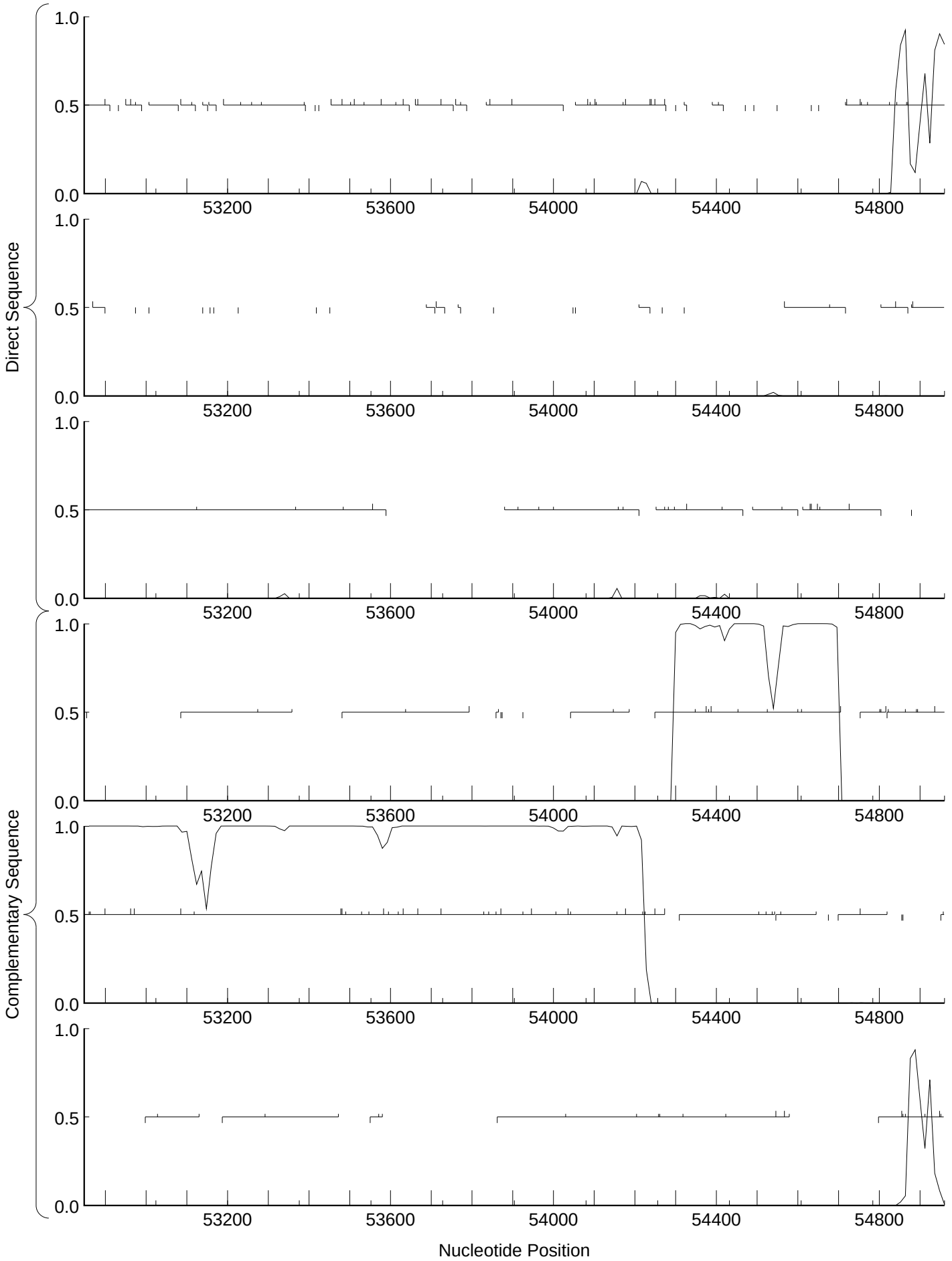


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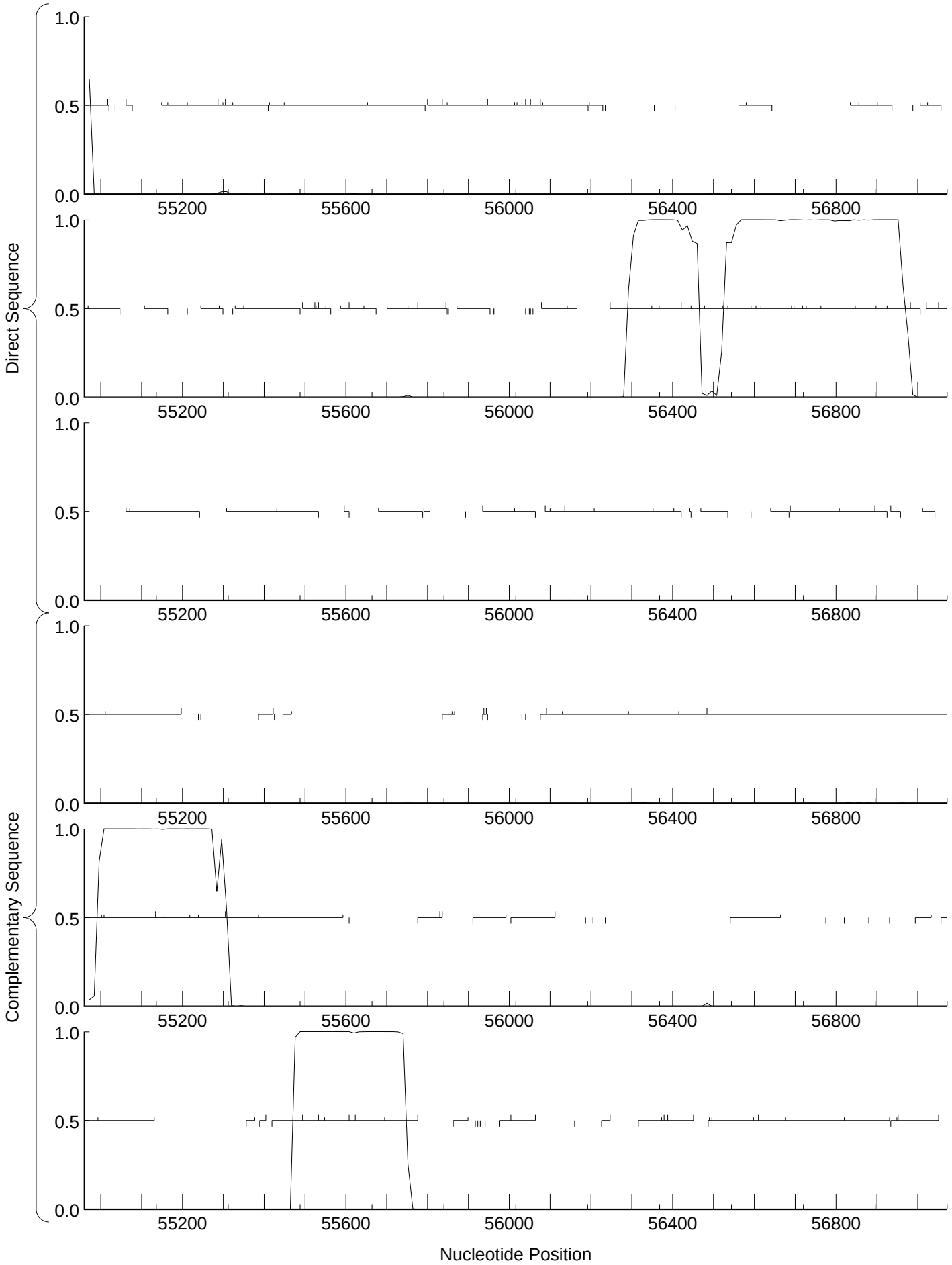


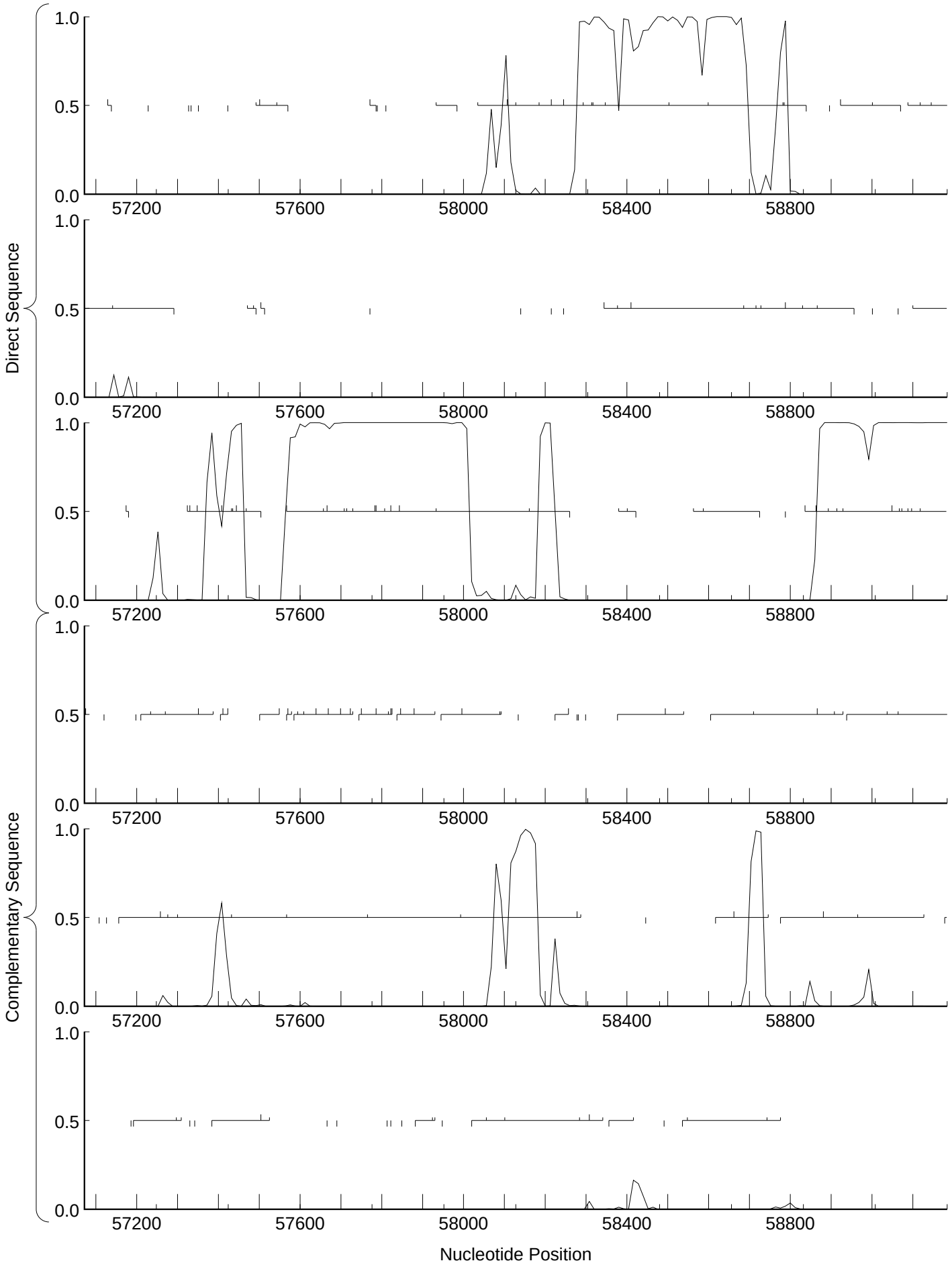

Direct Sequence
Complementary Sequence
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0.5
0.0
50800
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52000
52400
52800
Nucleotide Position

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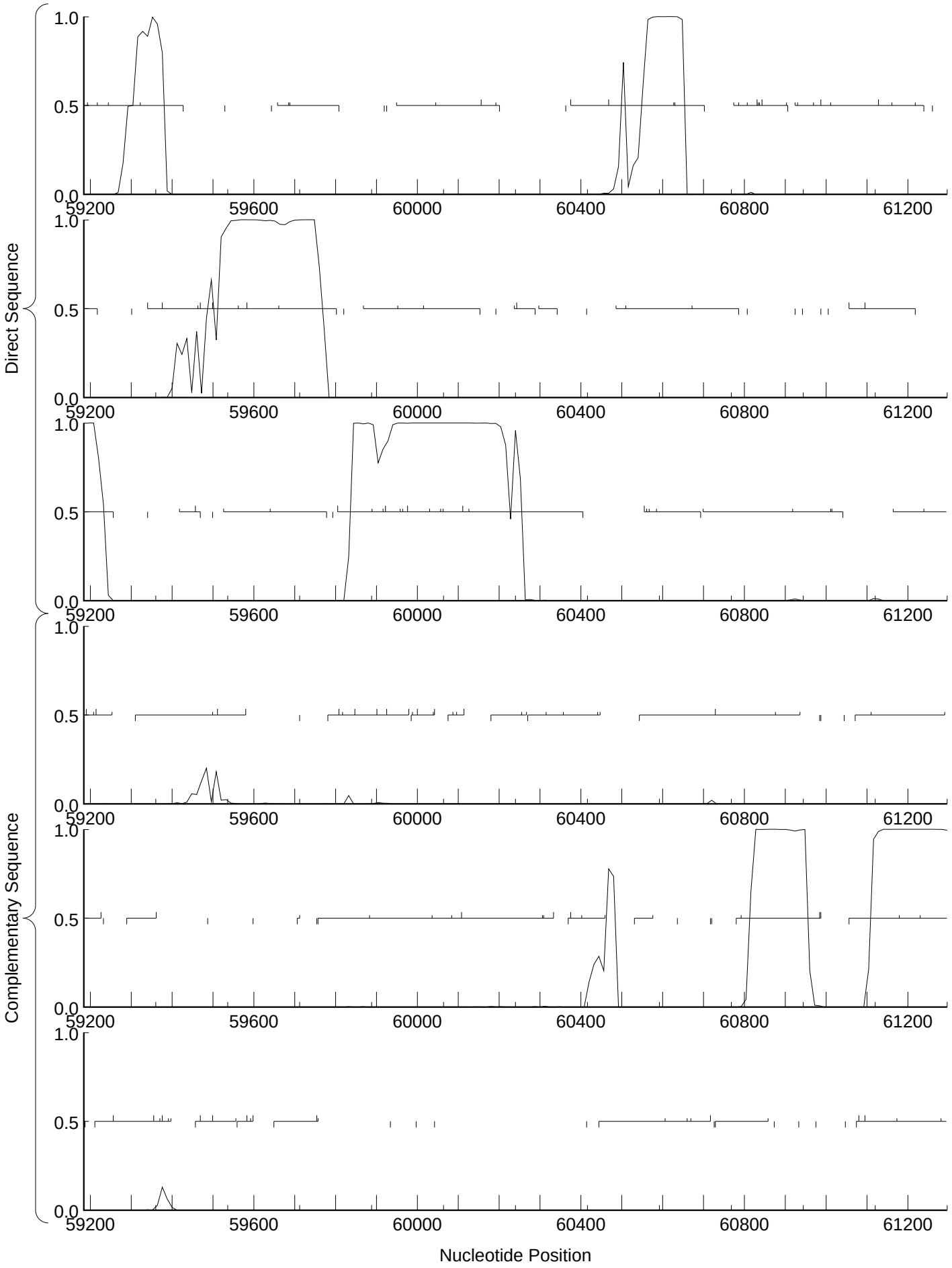


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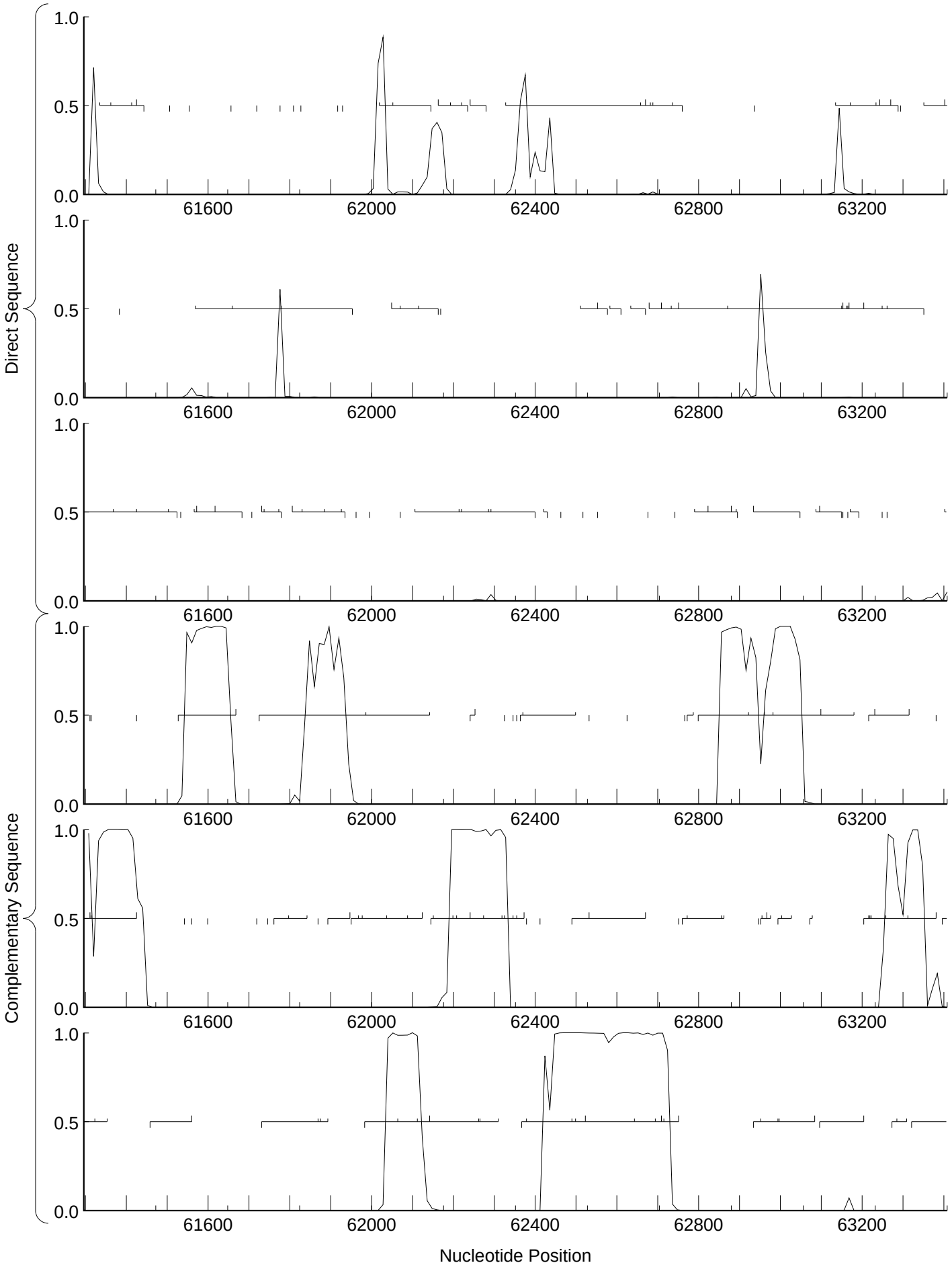



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Complementary Sequence
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Nucleotide Position

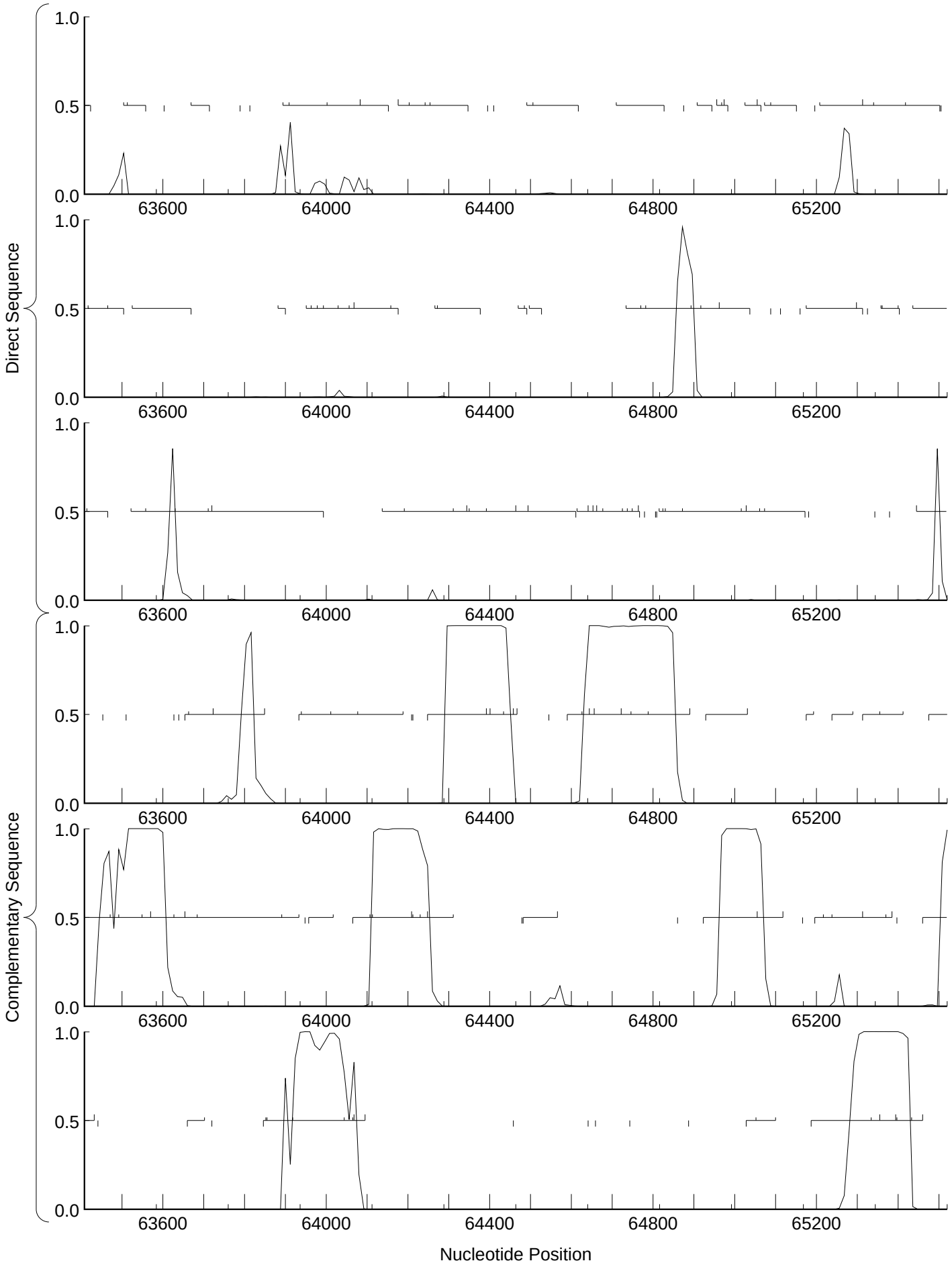
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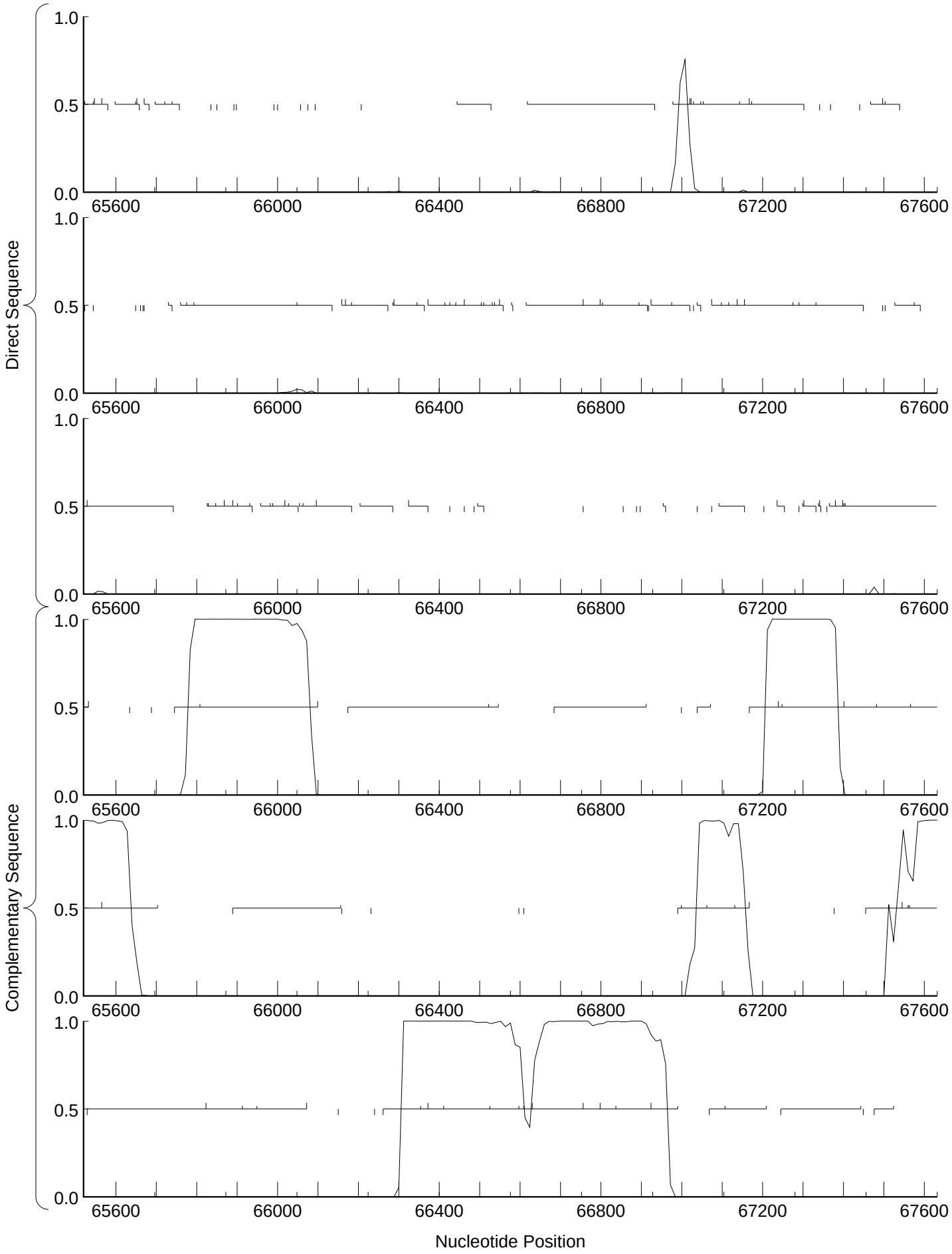
acterium phage Spartan300 Complete Sequence, 68568 bp, Circularly Permuted, Cluster B1, Order 4, Window 96, Step 12, 31/34



acterium phage Spartan300 Complete Sequence, 68568 bp, Circularly Permuted, Cluster B1, Order 4, Window 96, Step 12, 32/34



acterium phage Spartan300 Complete Sequence, 68568 bp, Circularly Permuted, Cluster B1, Order 4, Window 96, Step 12, 33/34



acterium phage Spartan300 Complete Sequence, 68568 bp, Circularly Permuted, Cluster B1, Order 4, Window 96, Step 12, 34/34

