GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage CharlieGBrown complete sequence, 68604 bp, circularly permuted, Cluster
Analysis Date    :  3/29/24 at 9:17:53
Pages            :  34
Sequence Length  :  68604 bp
GC Content       :   66.39%

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

PS-Version       :  1.2

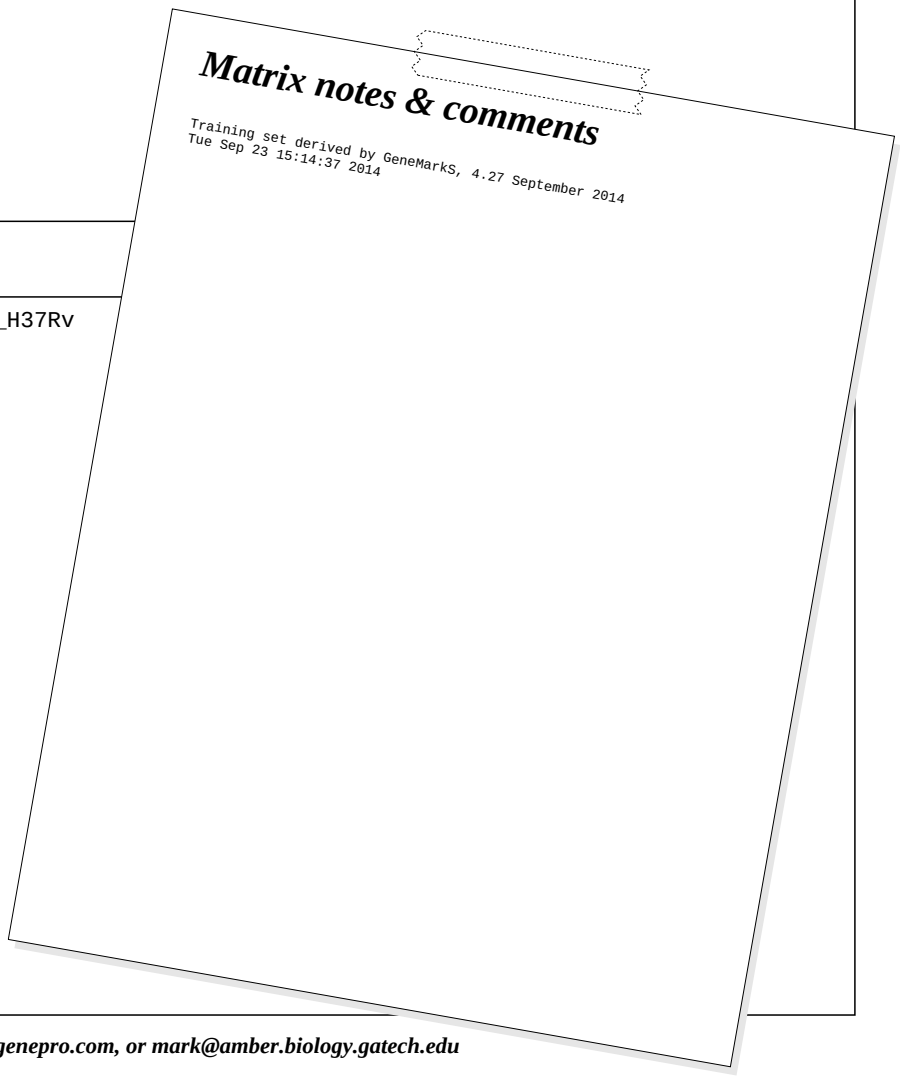
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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MATRIX INFORMATION

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Author     :  -

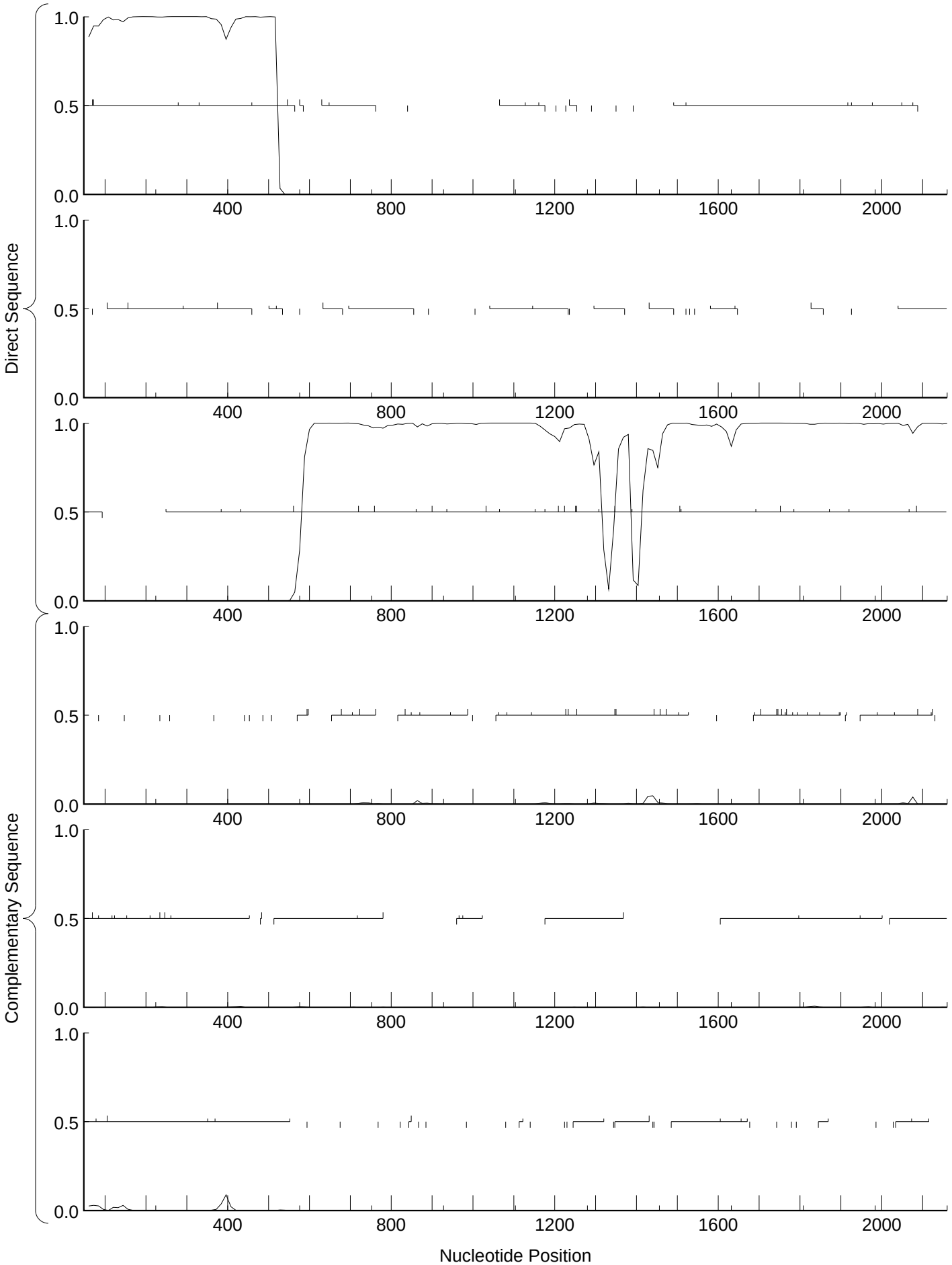
Order      :  4


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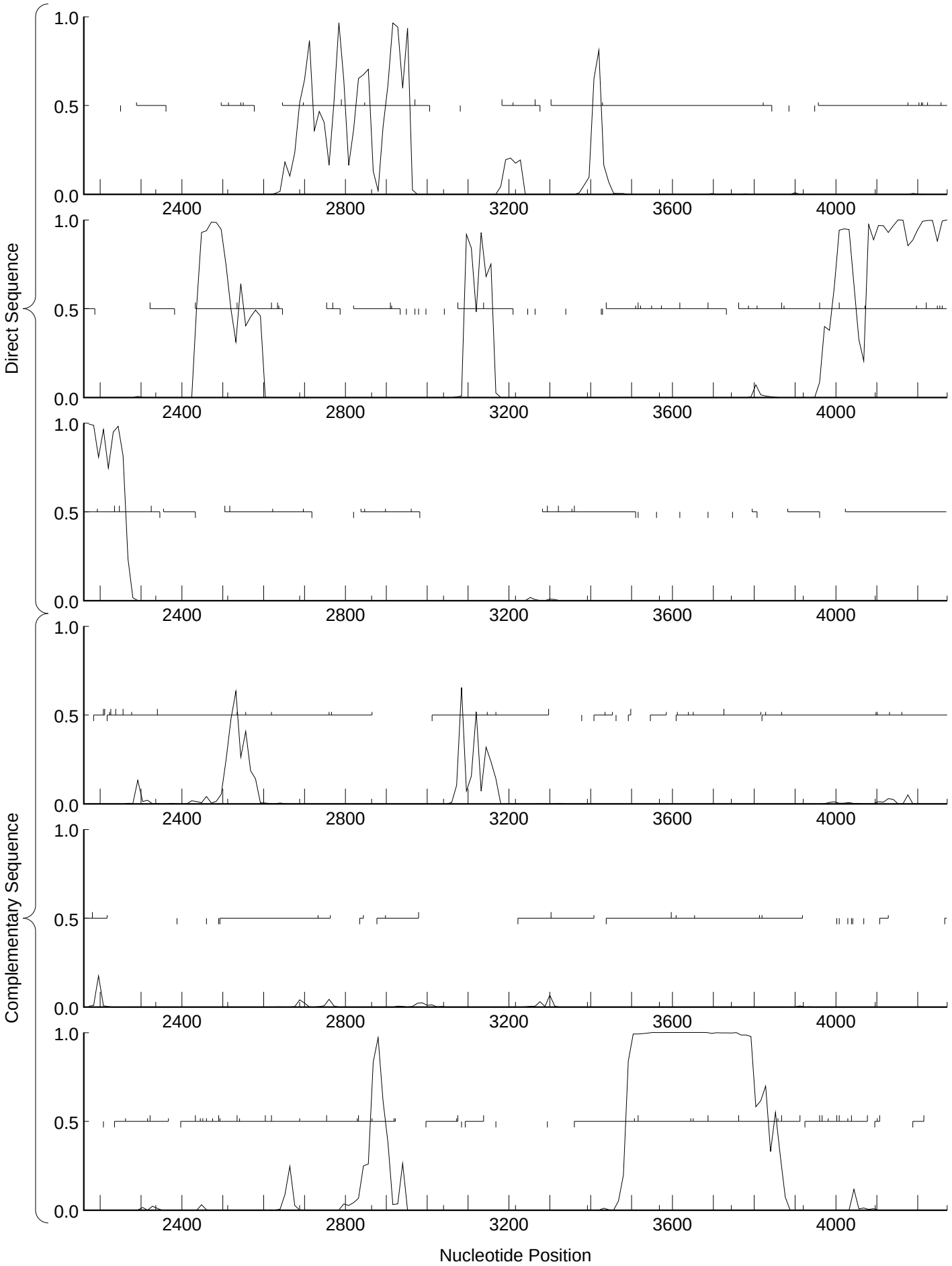


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

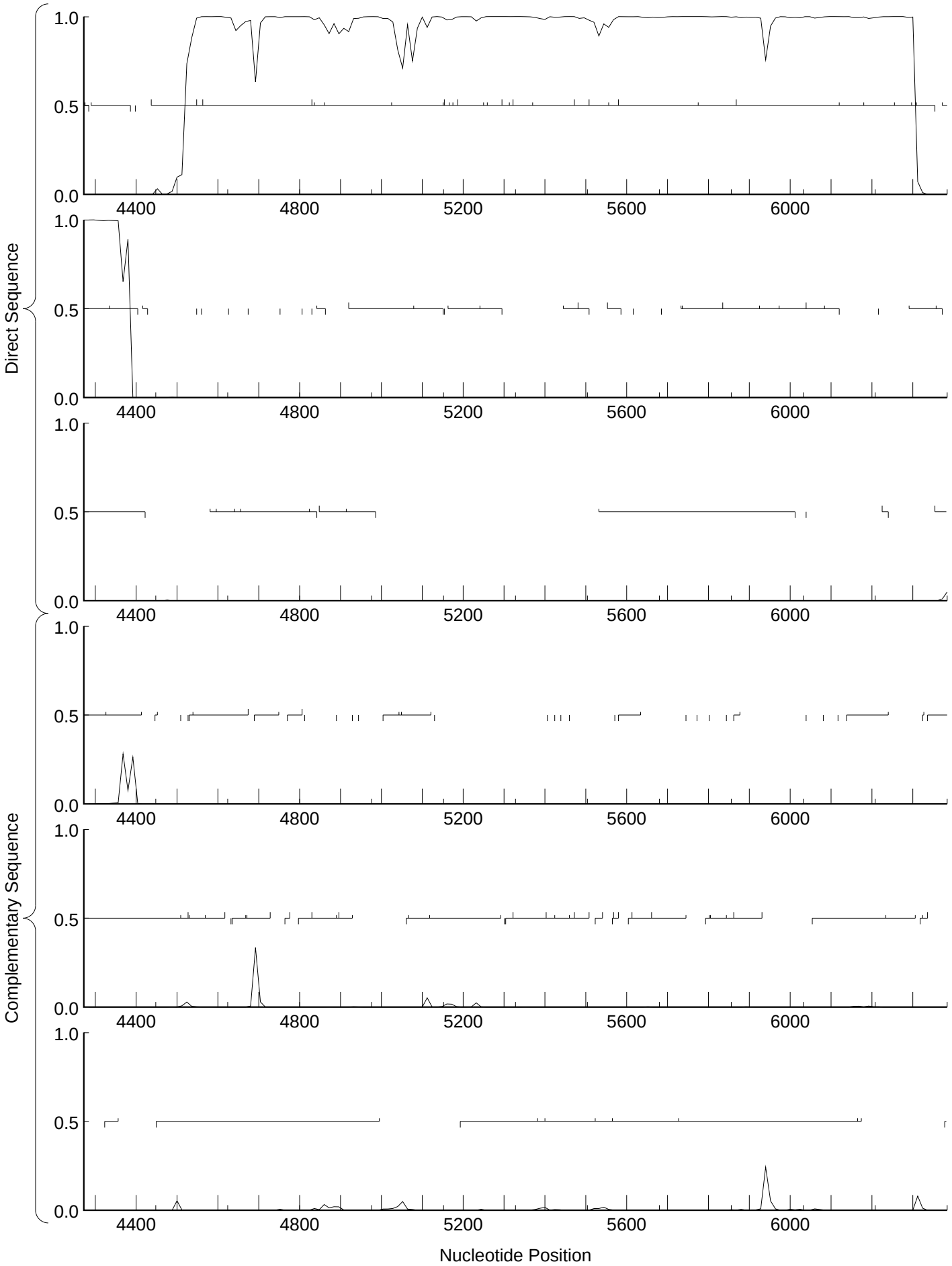
cterium phage CharlieGBrown complete sequence, 68604 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 2/34



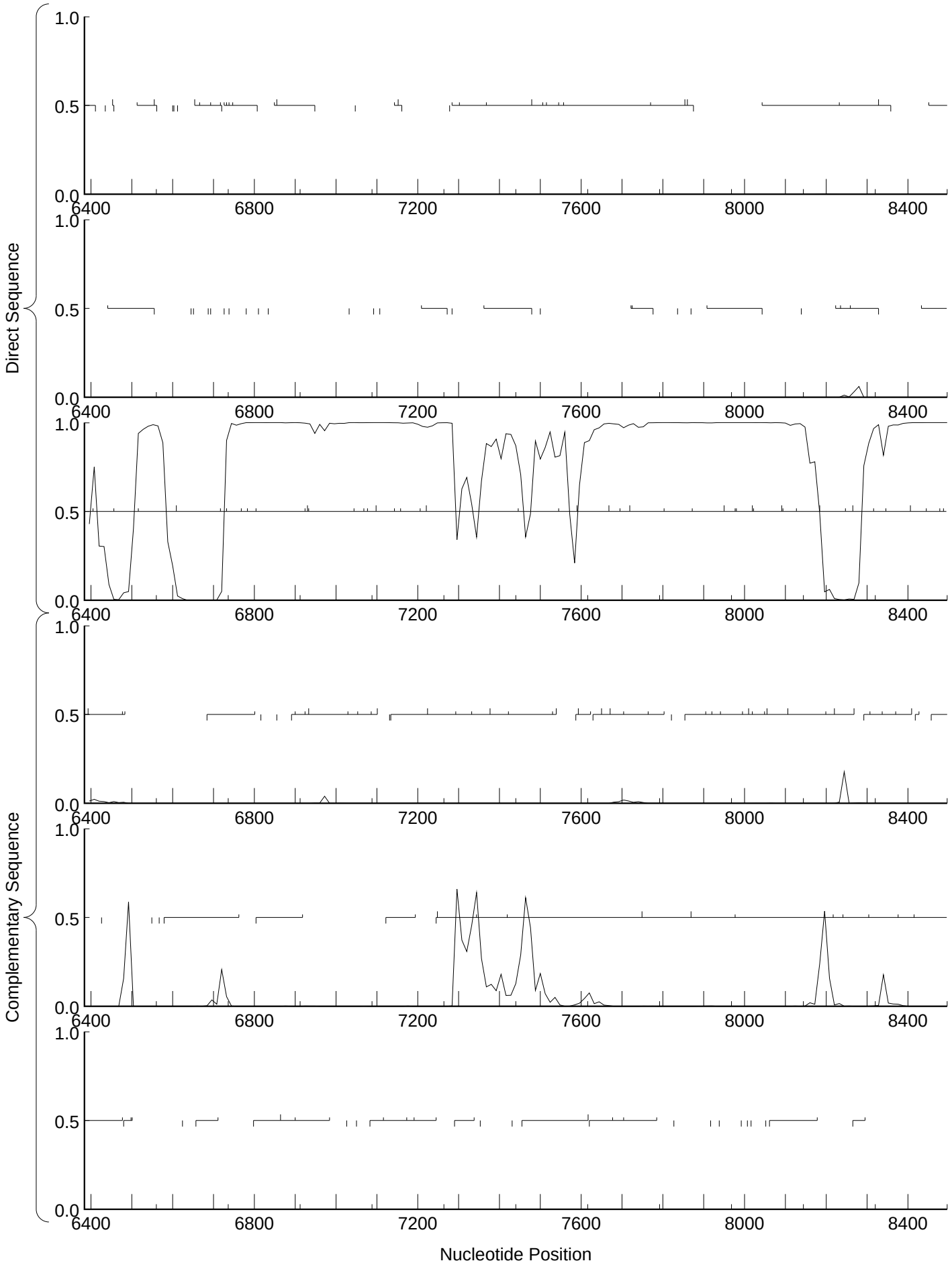
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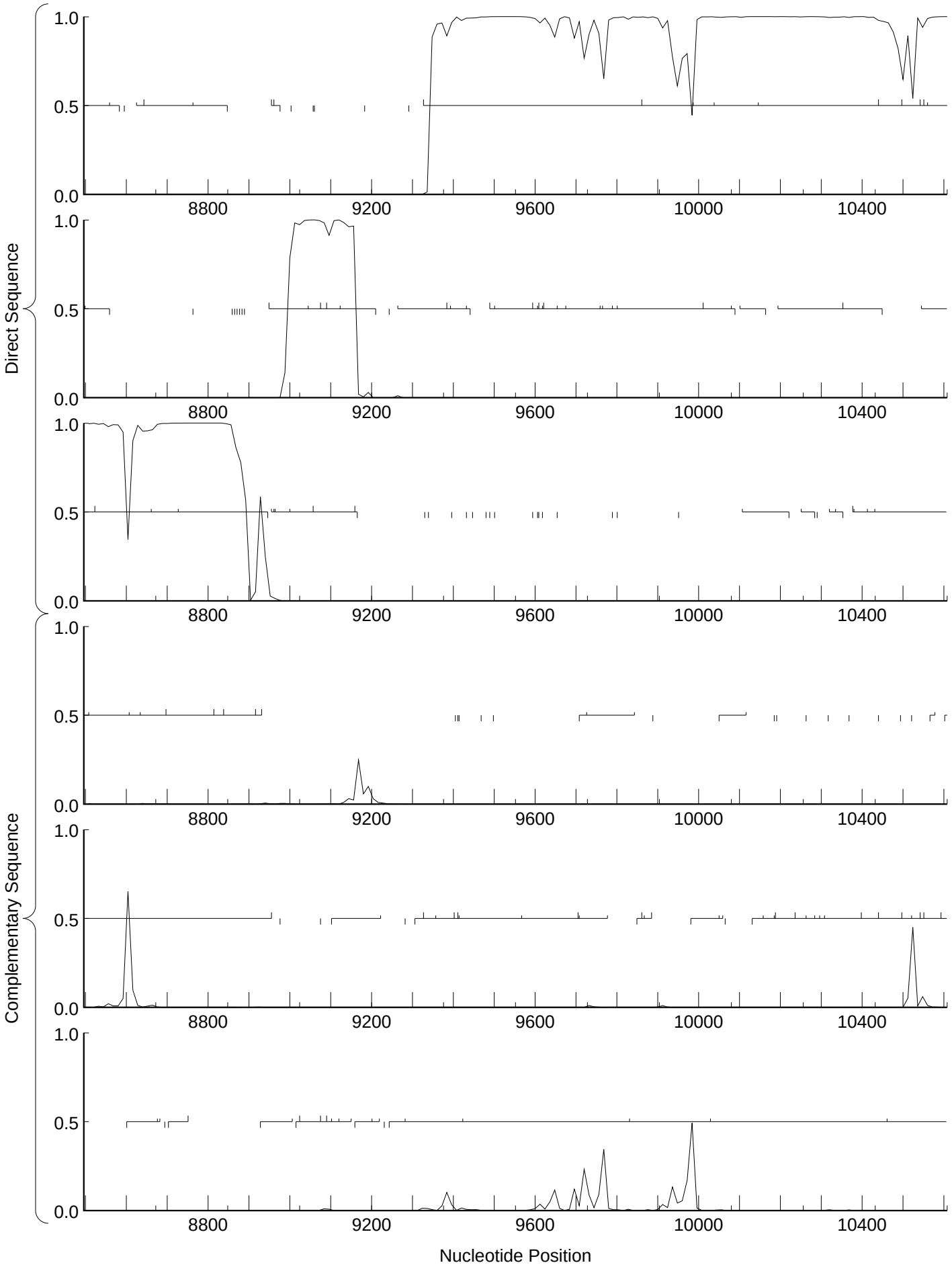
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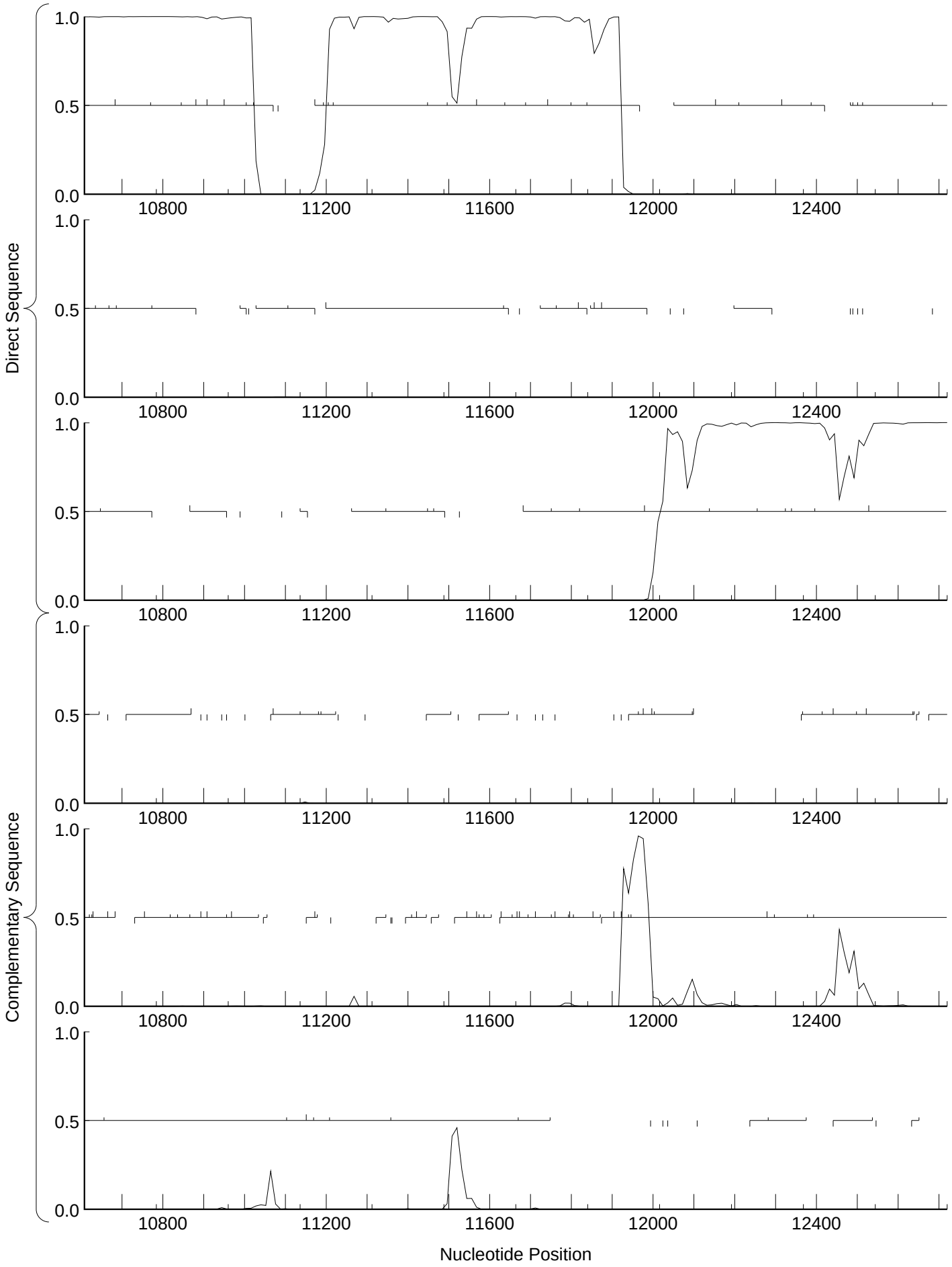
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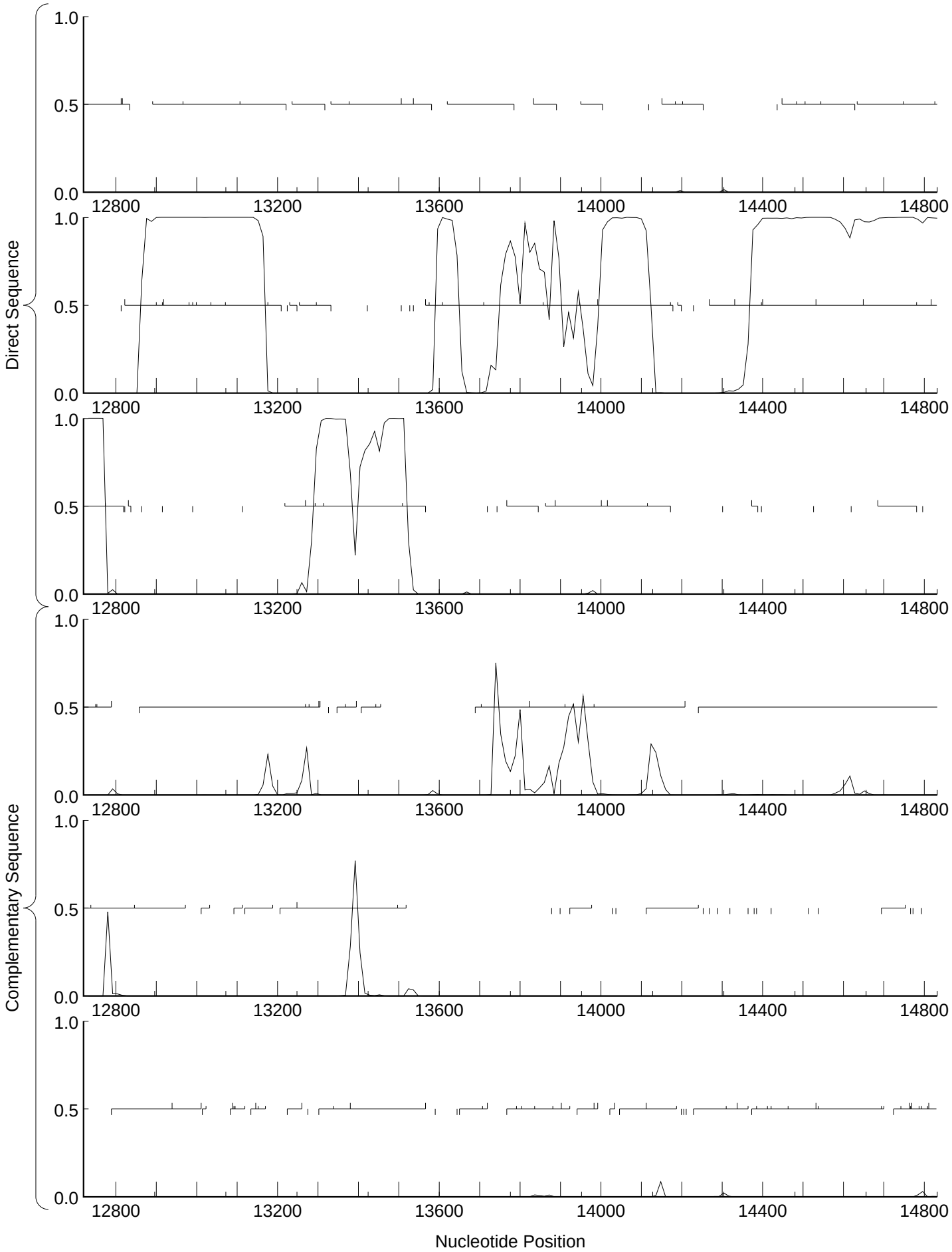
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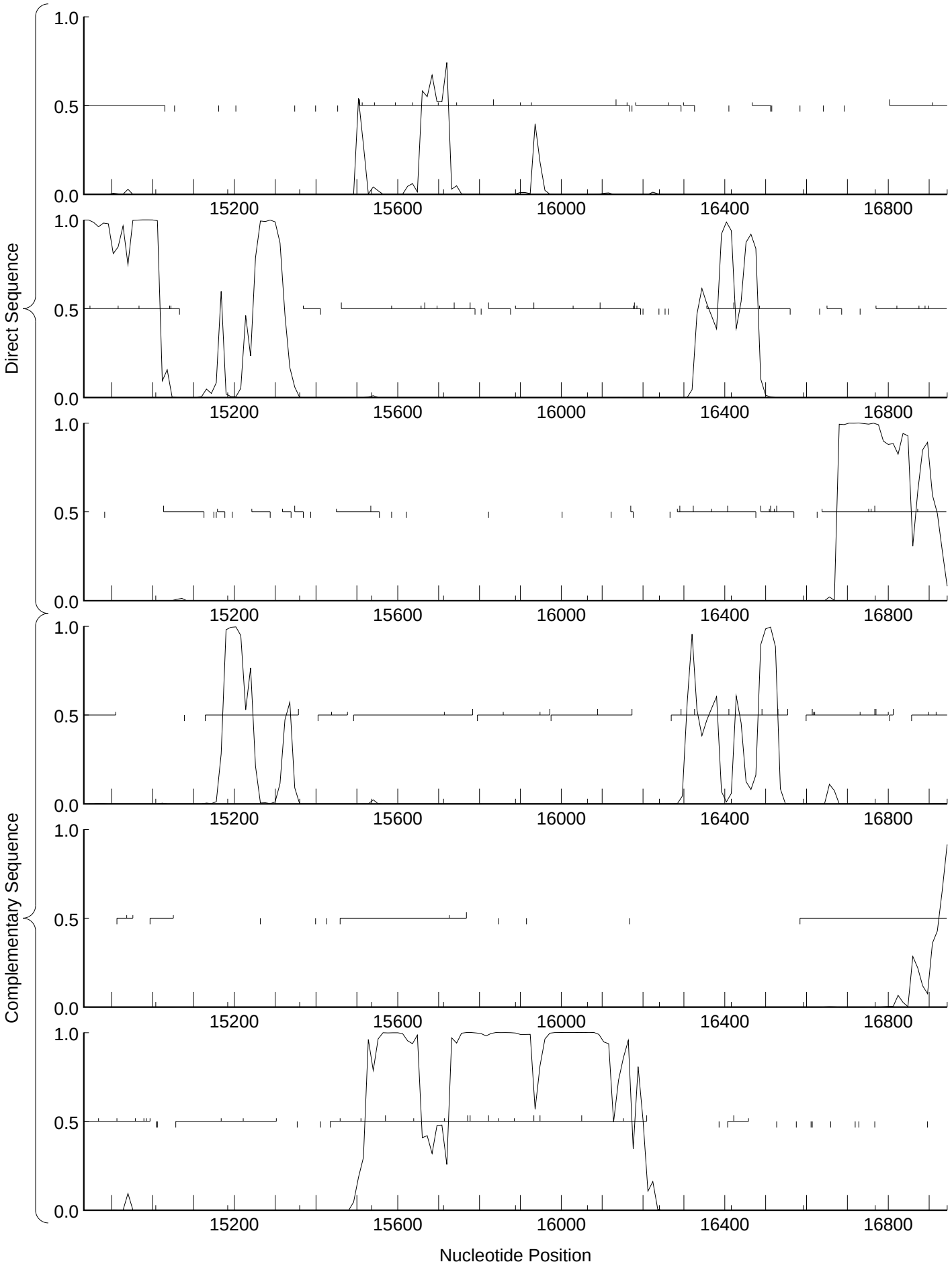
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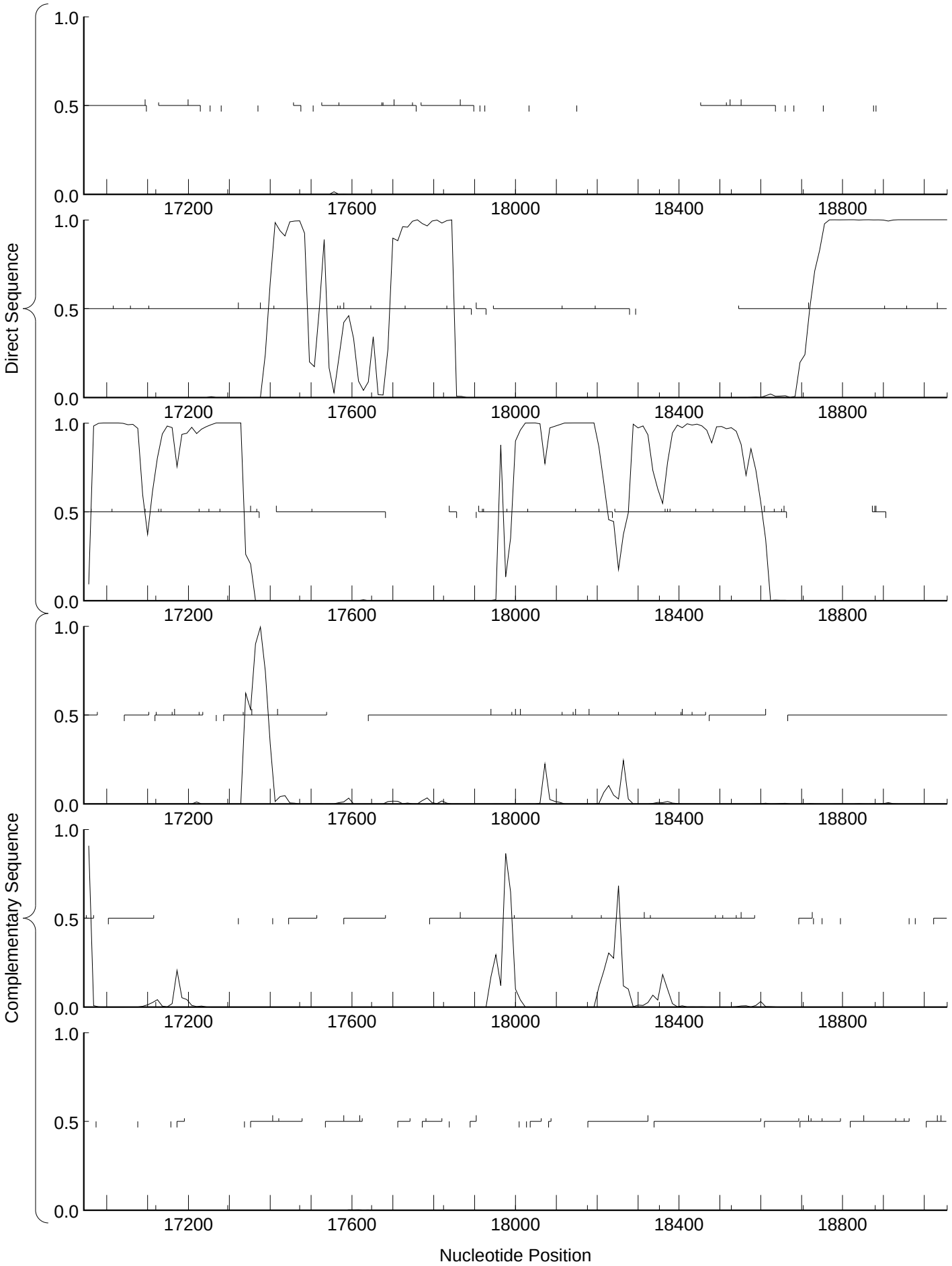
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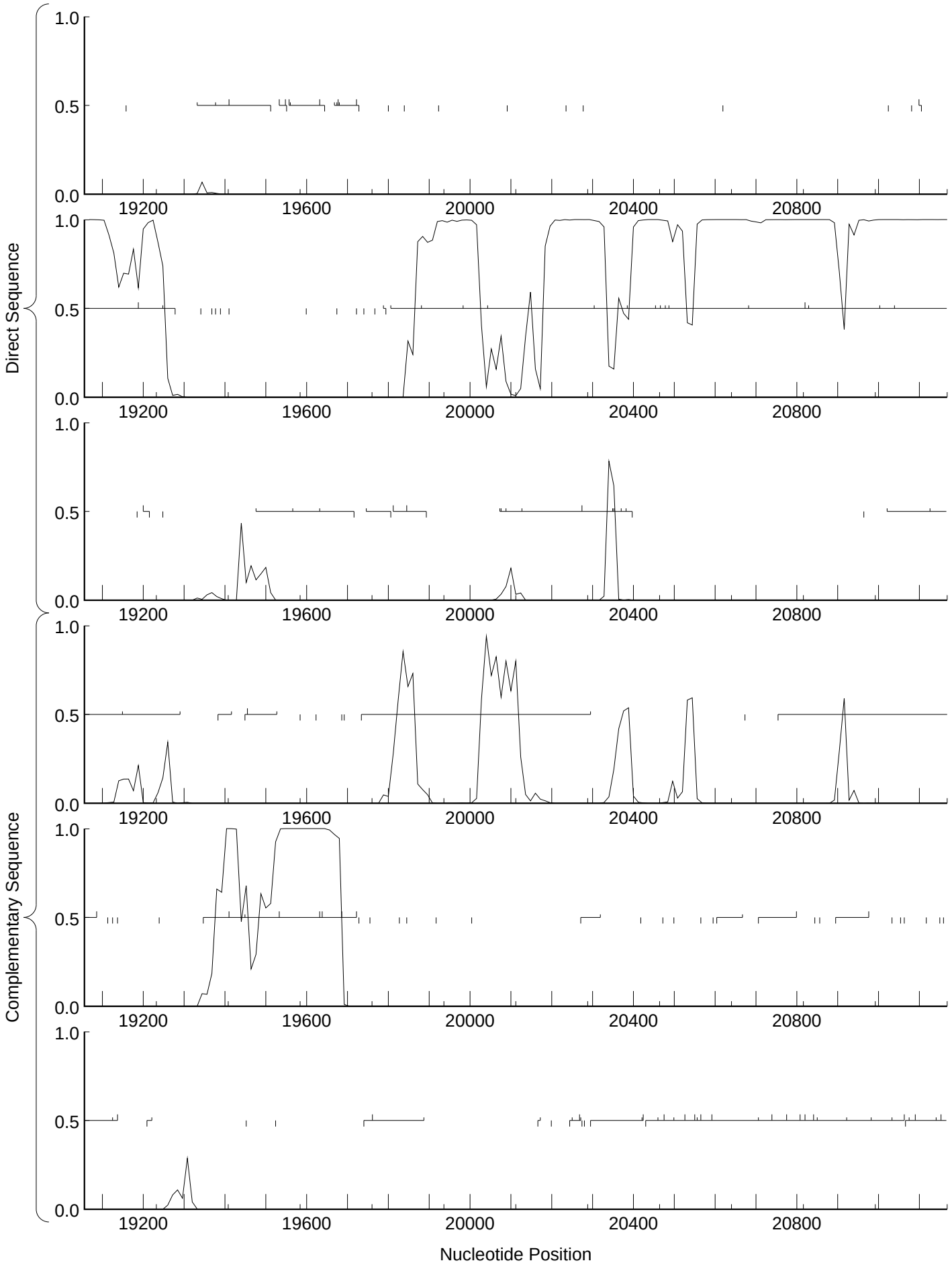
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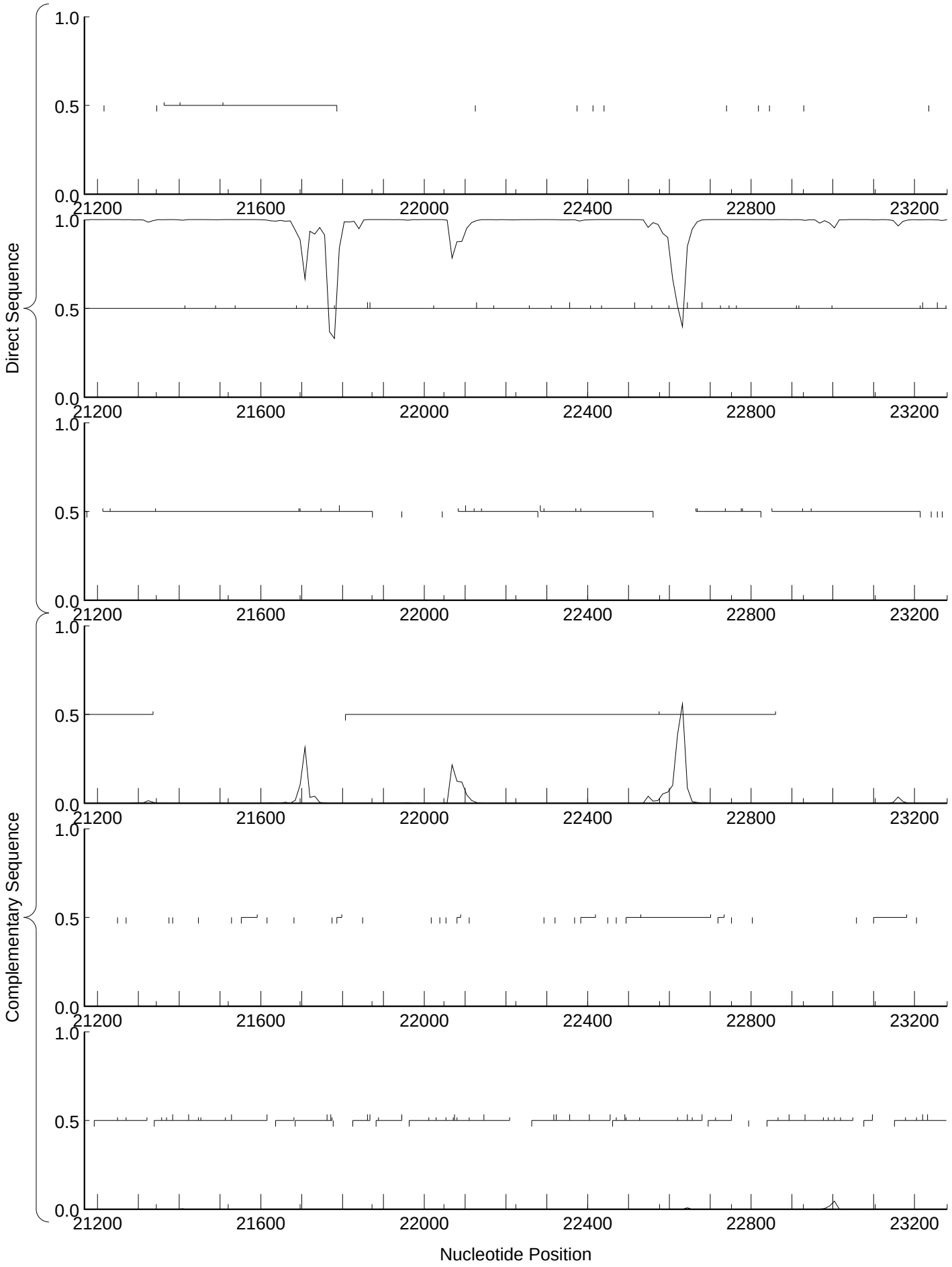
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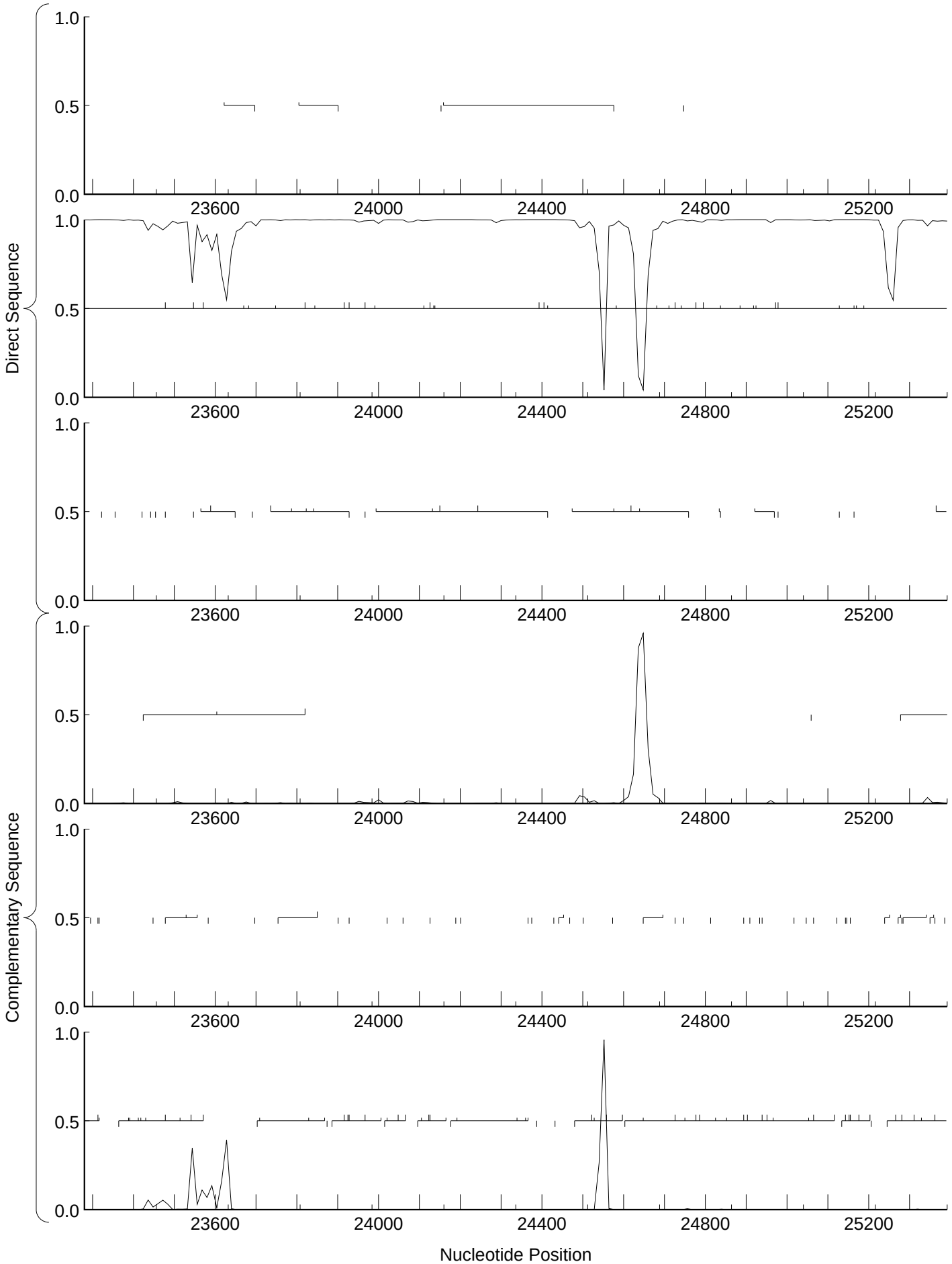
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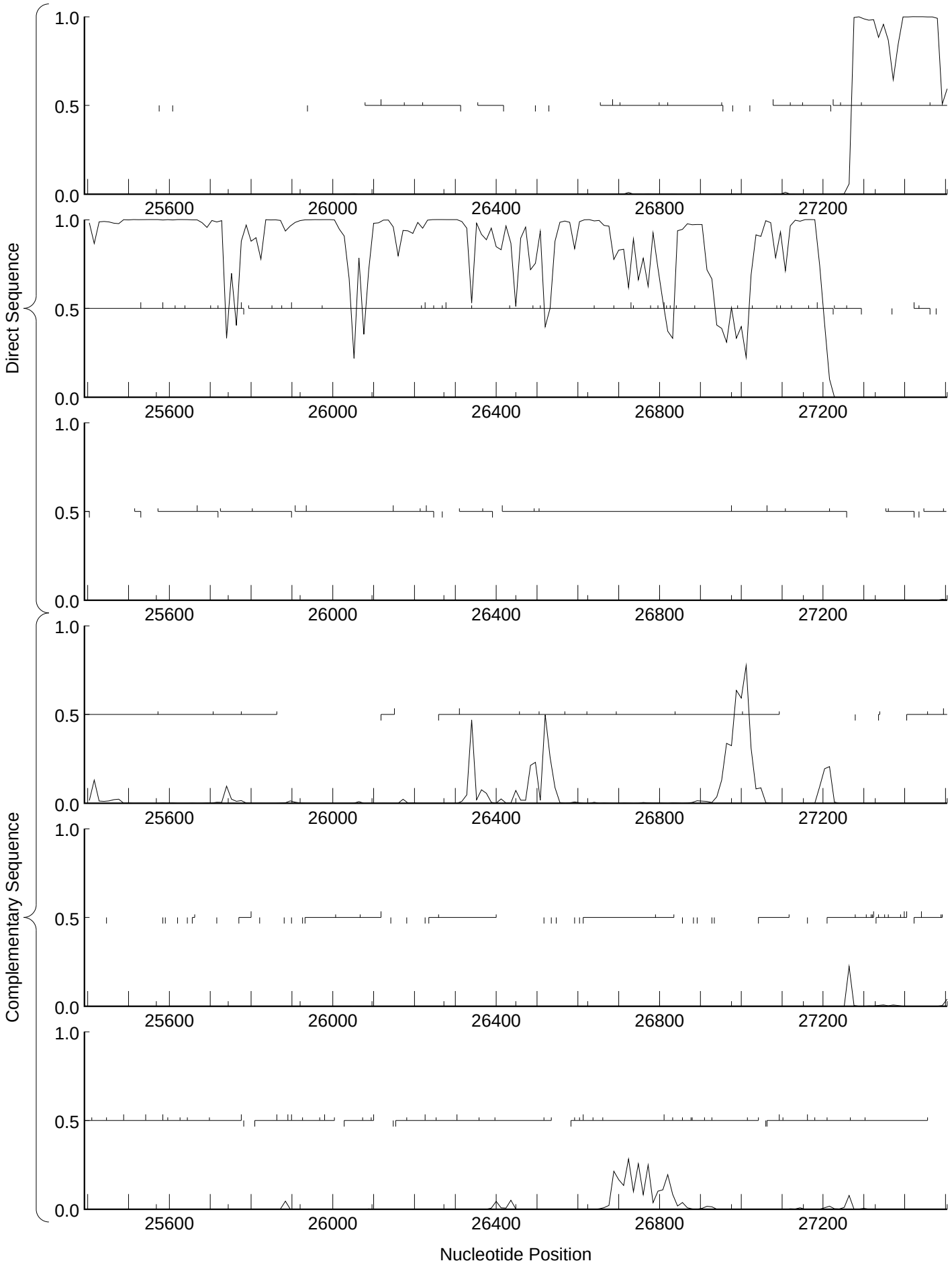
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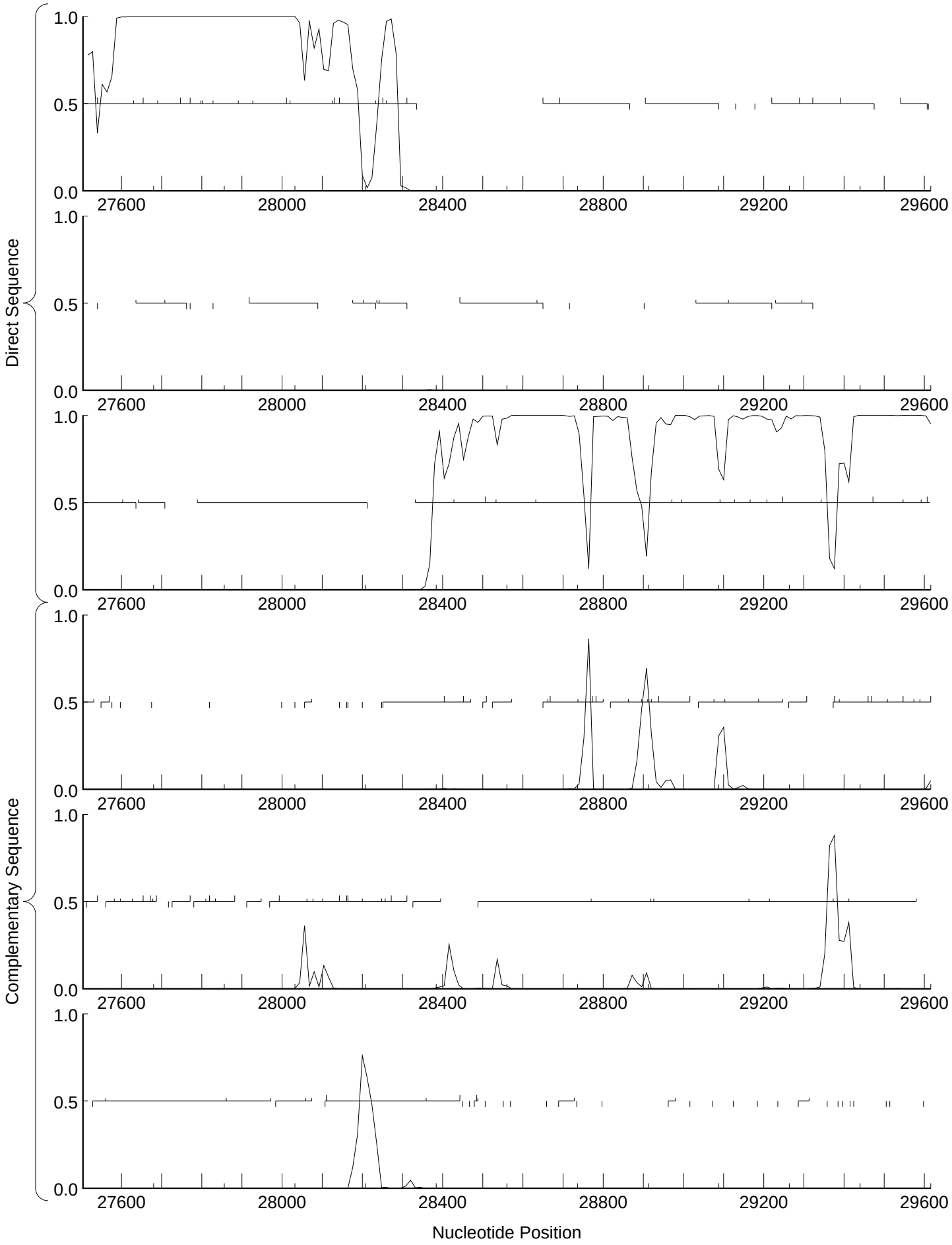
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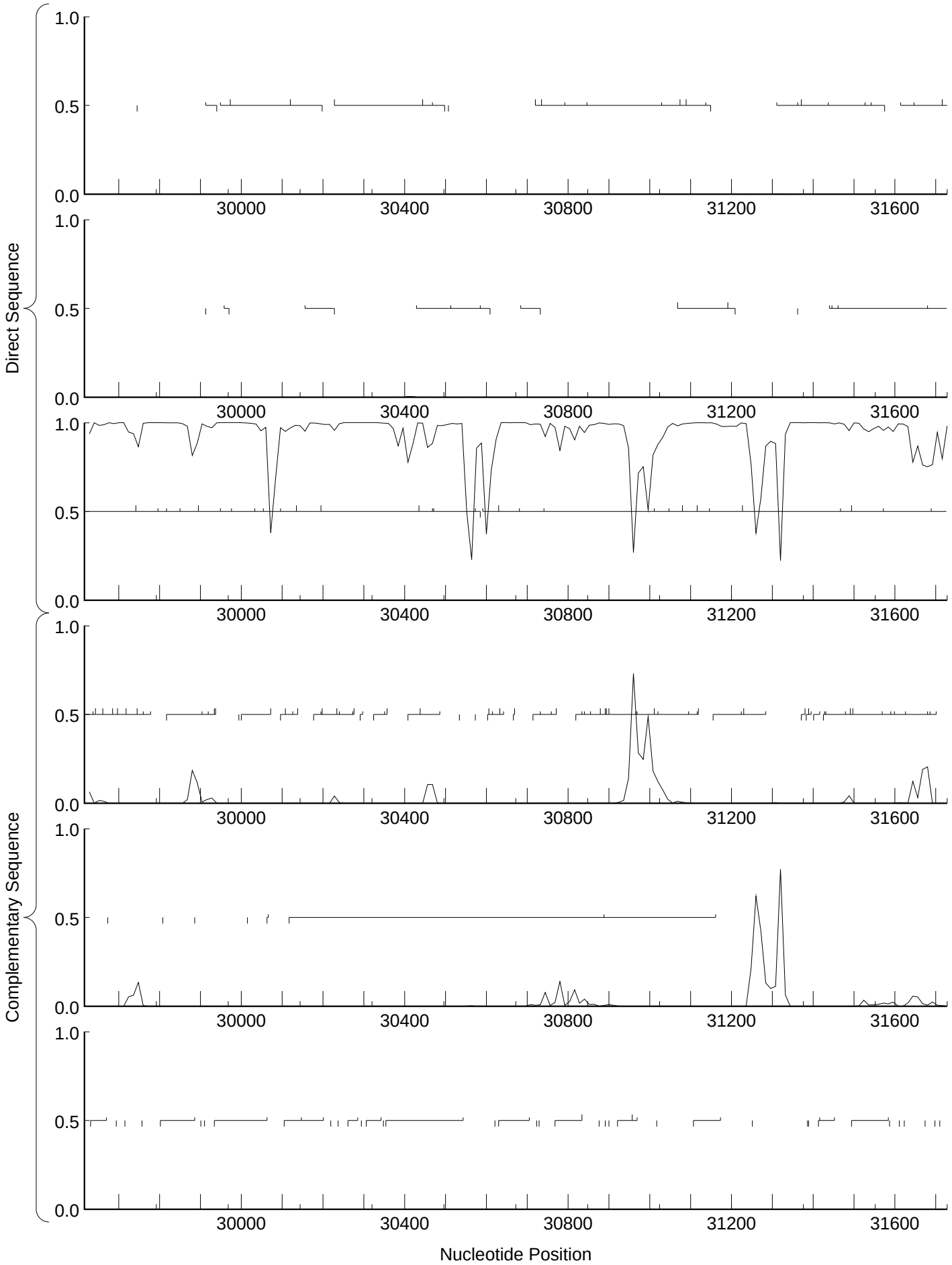
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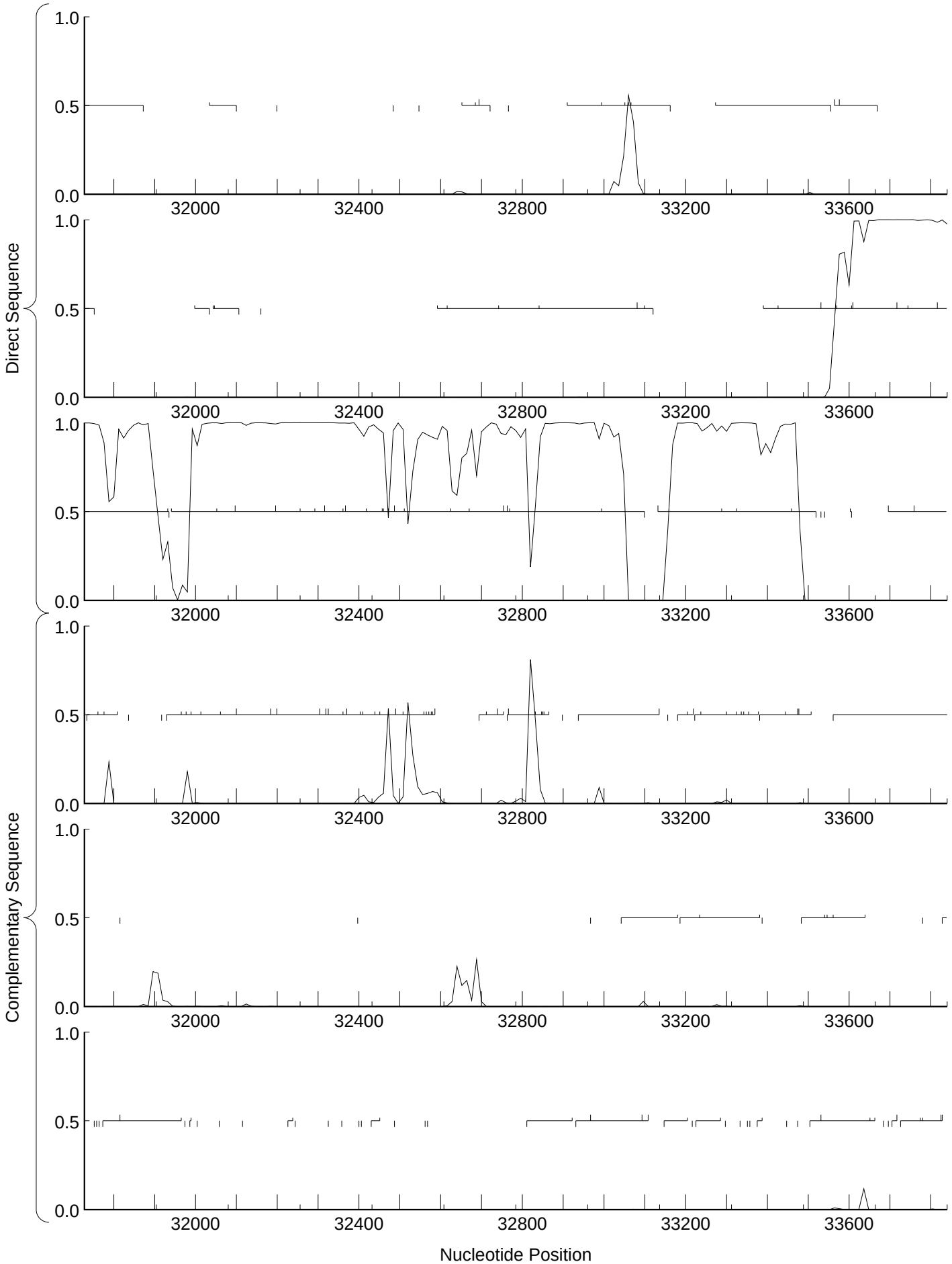
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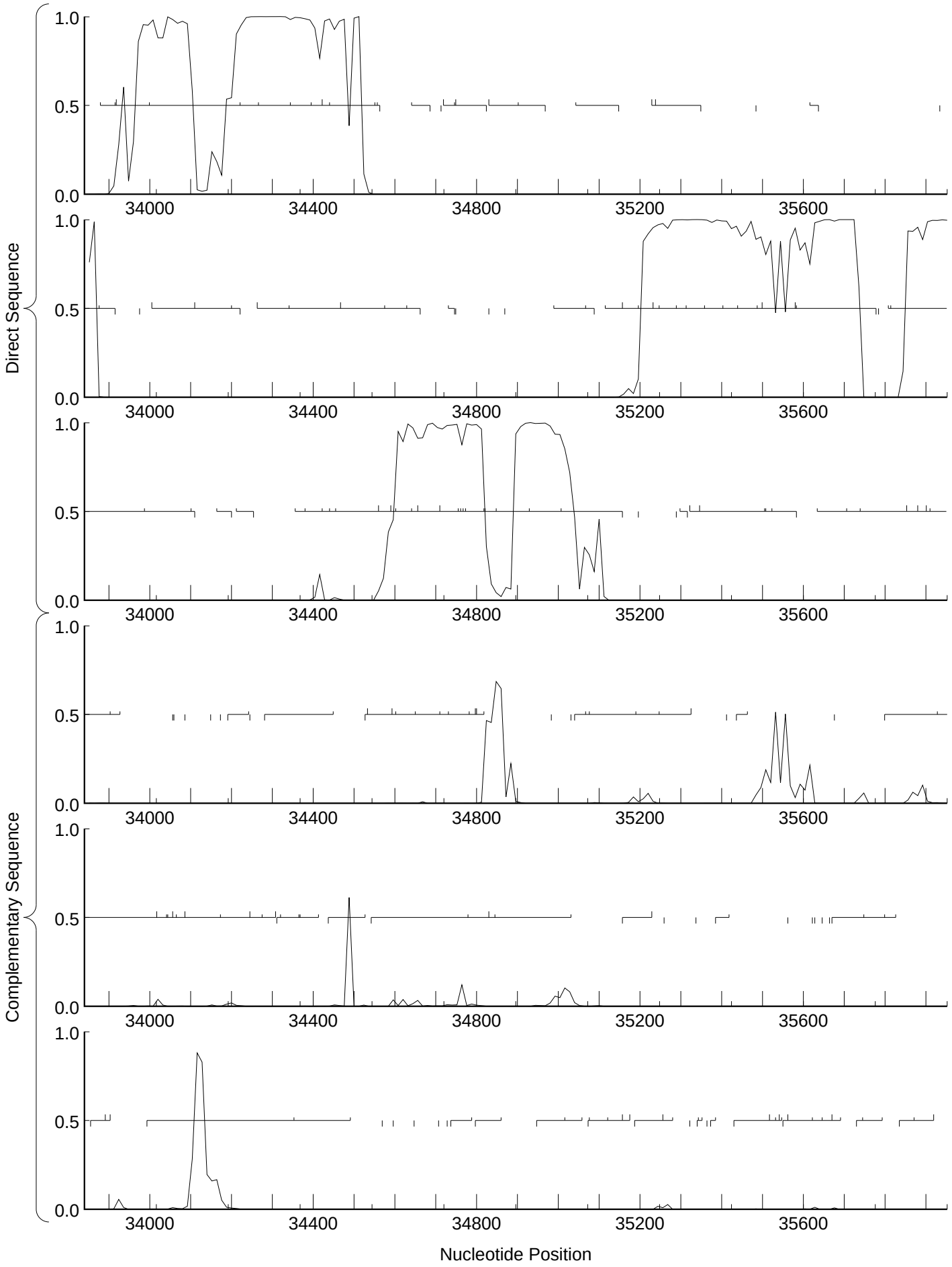
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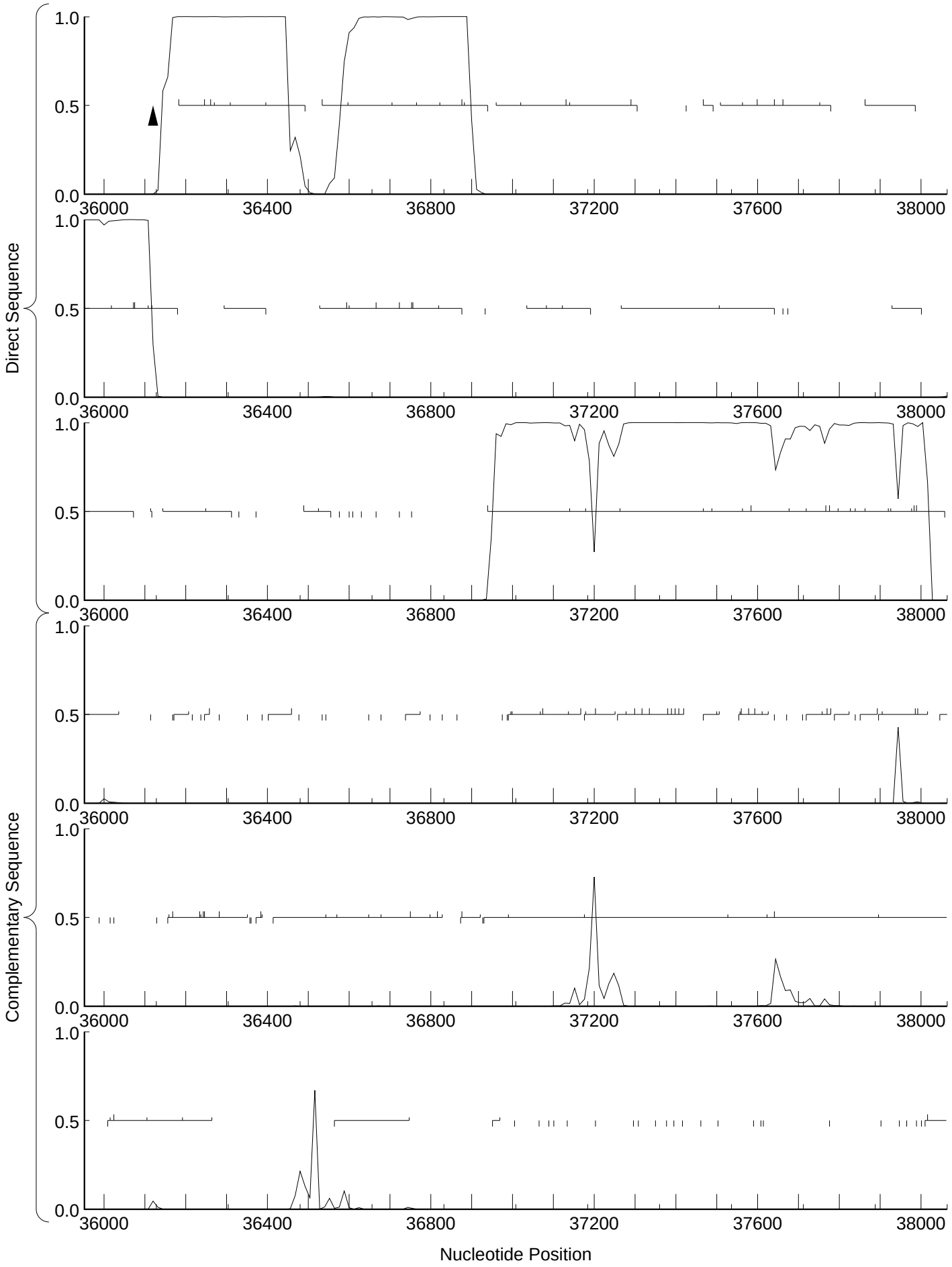
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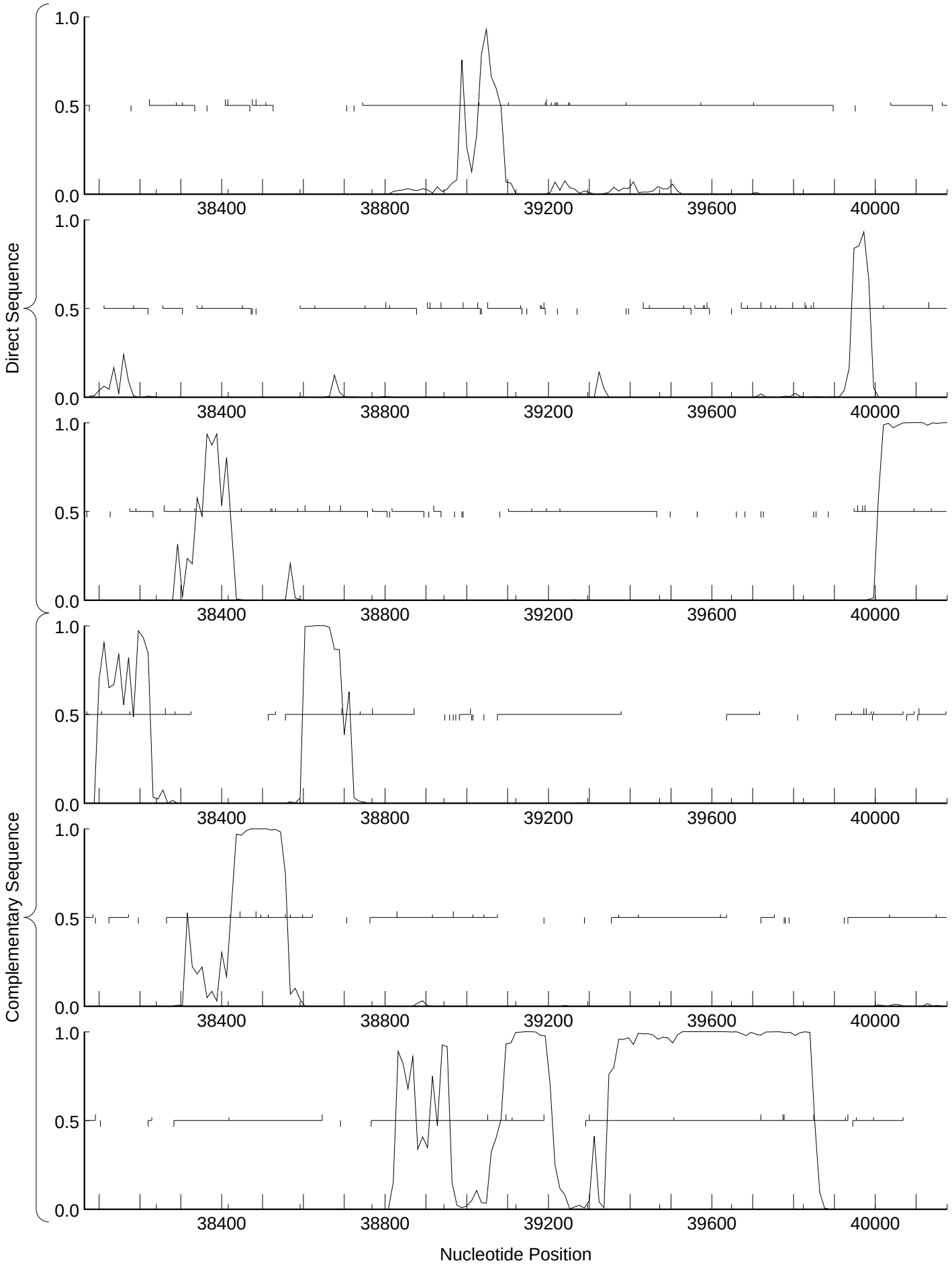


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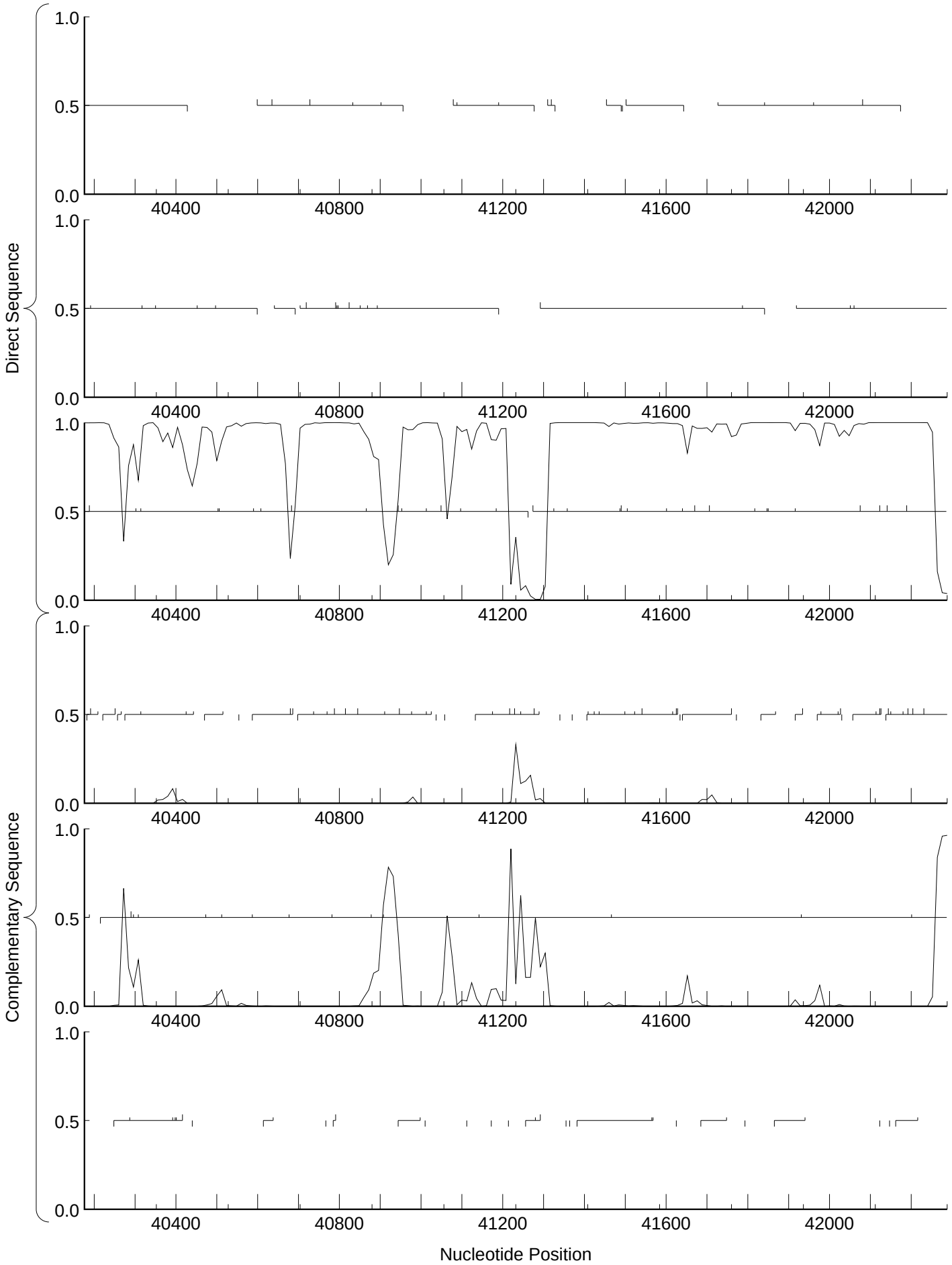


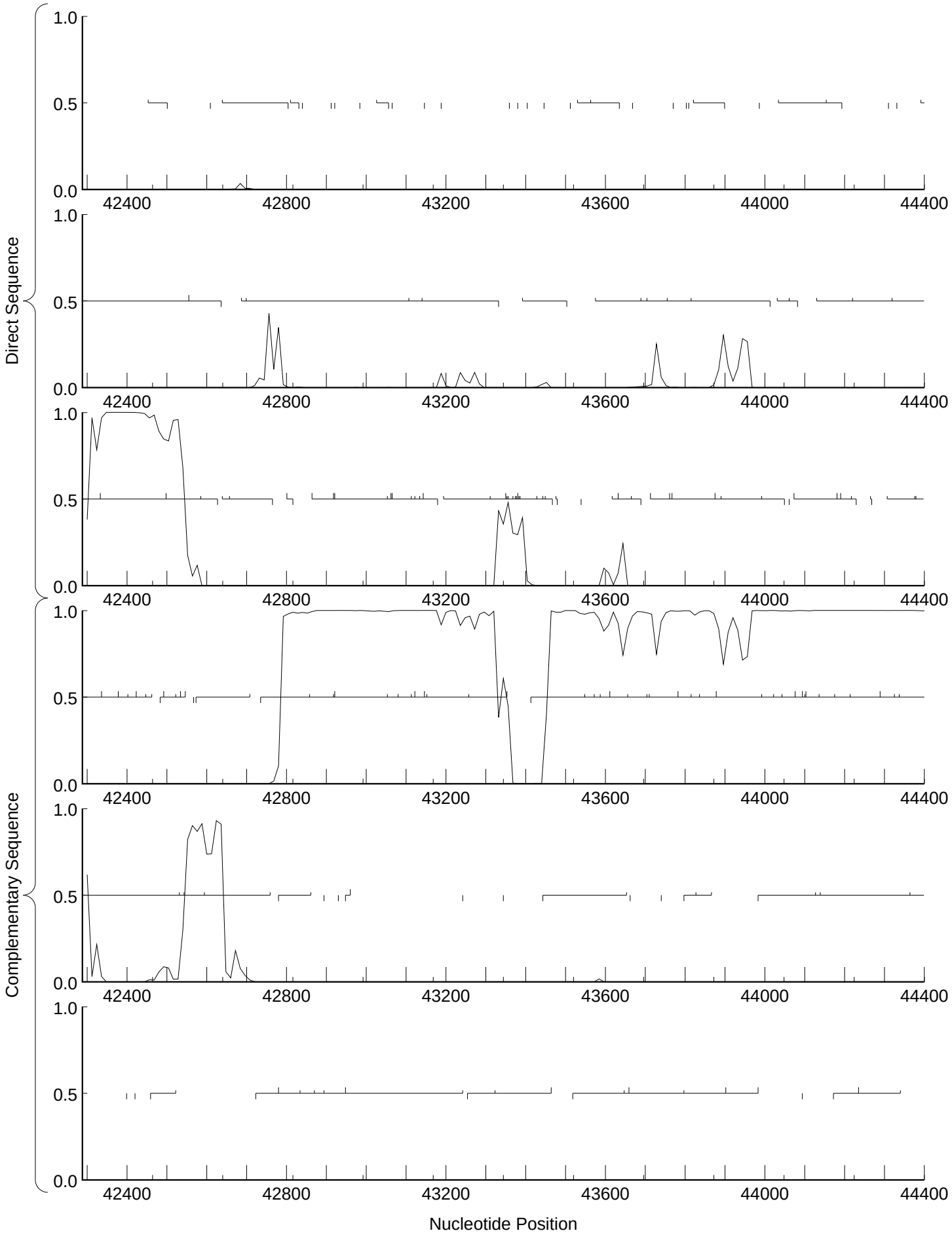
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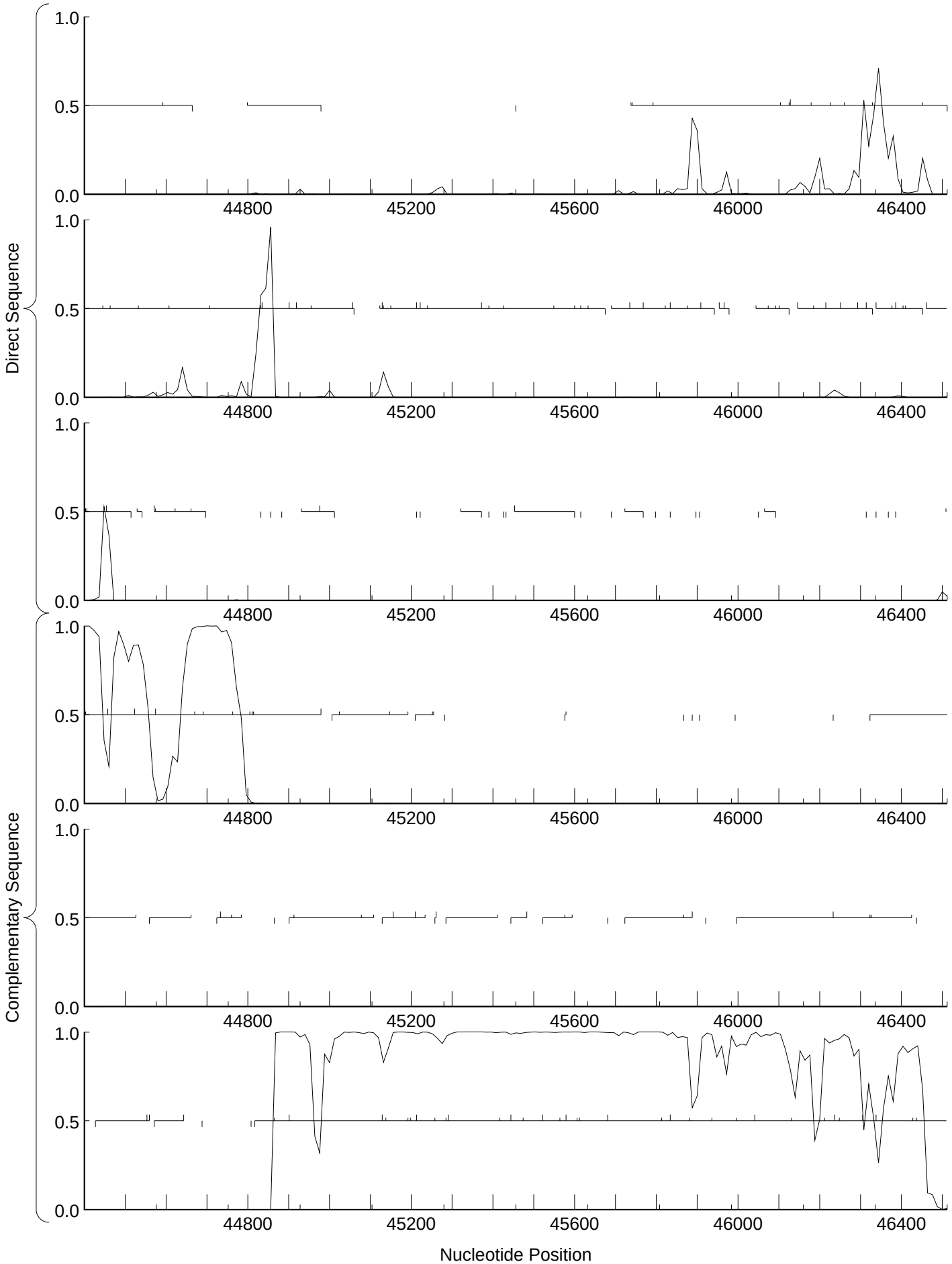



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

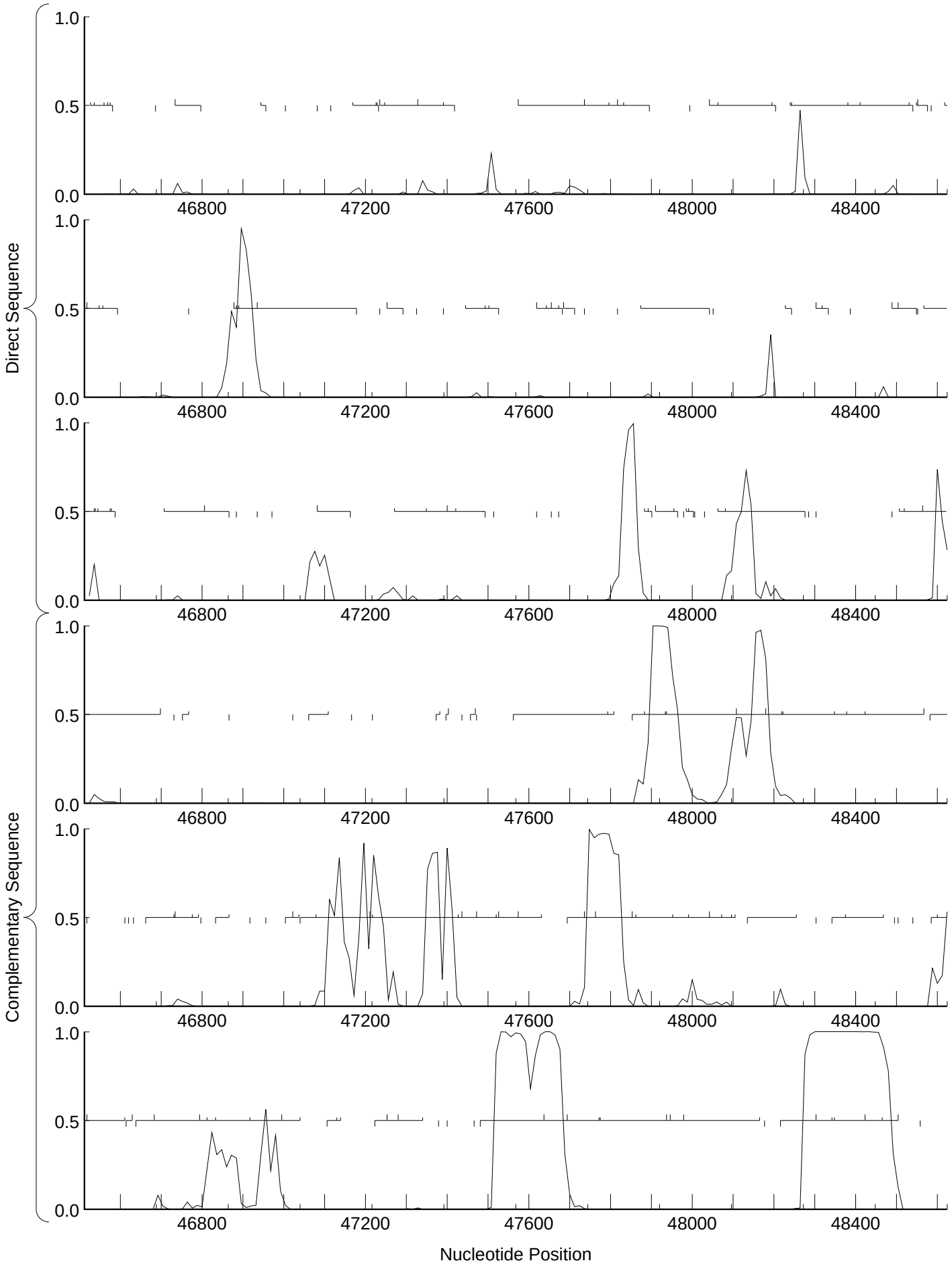
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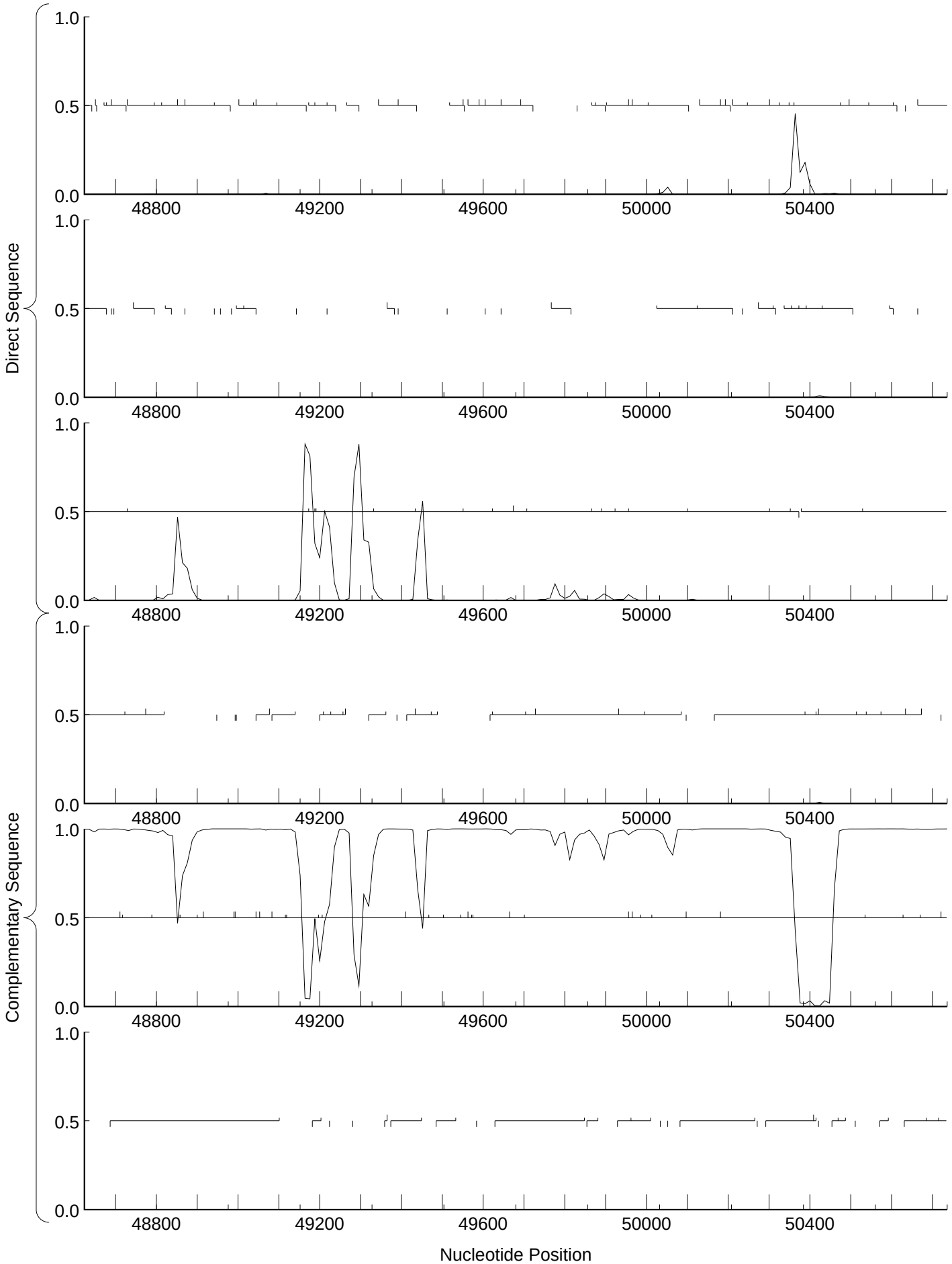

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
42400
42800
43200
43600
44000
44400
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
44800
45200
45600
46000
46400
Nucleotide Position

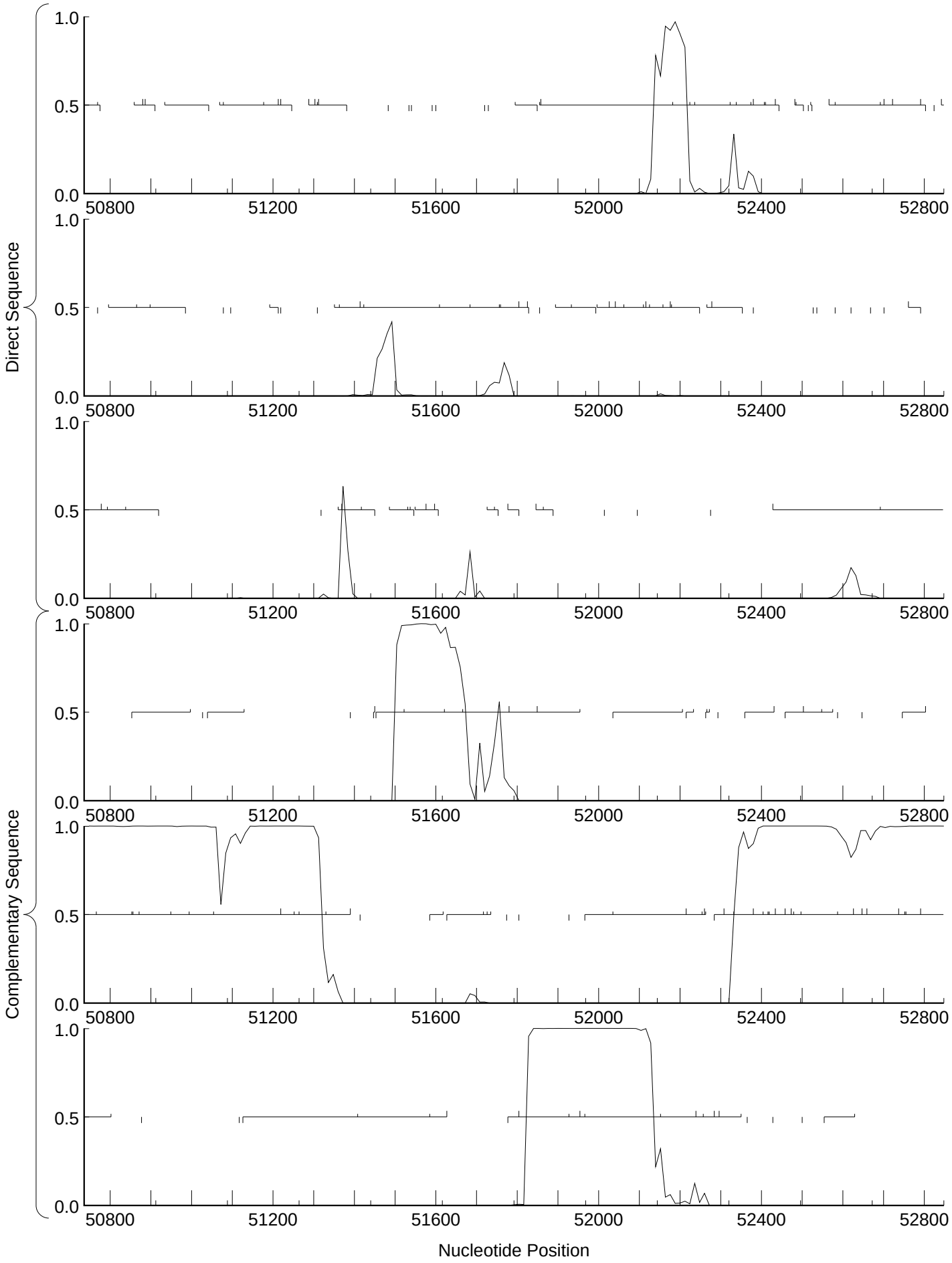
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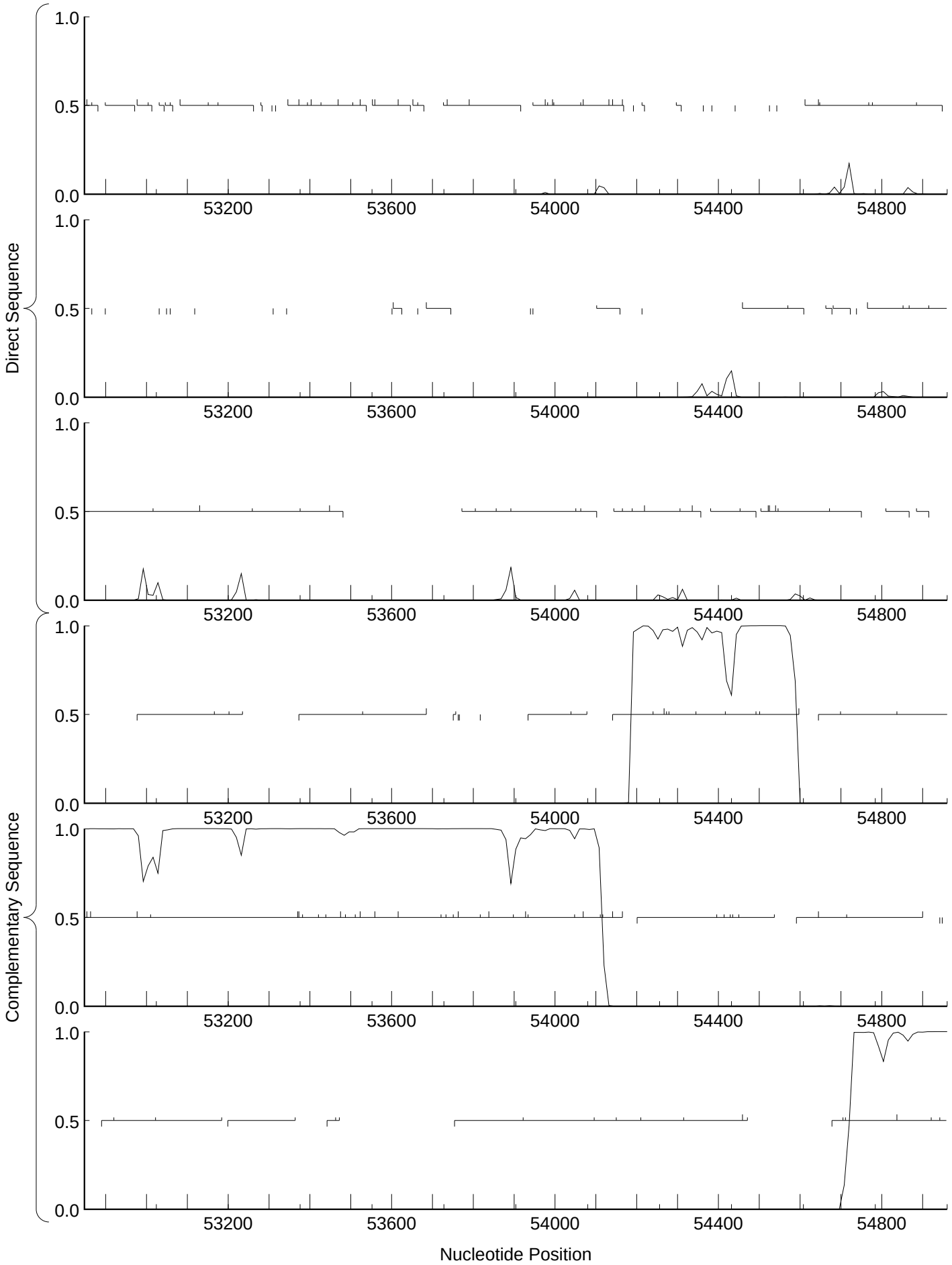
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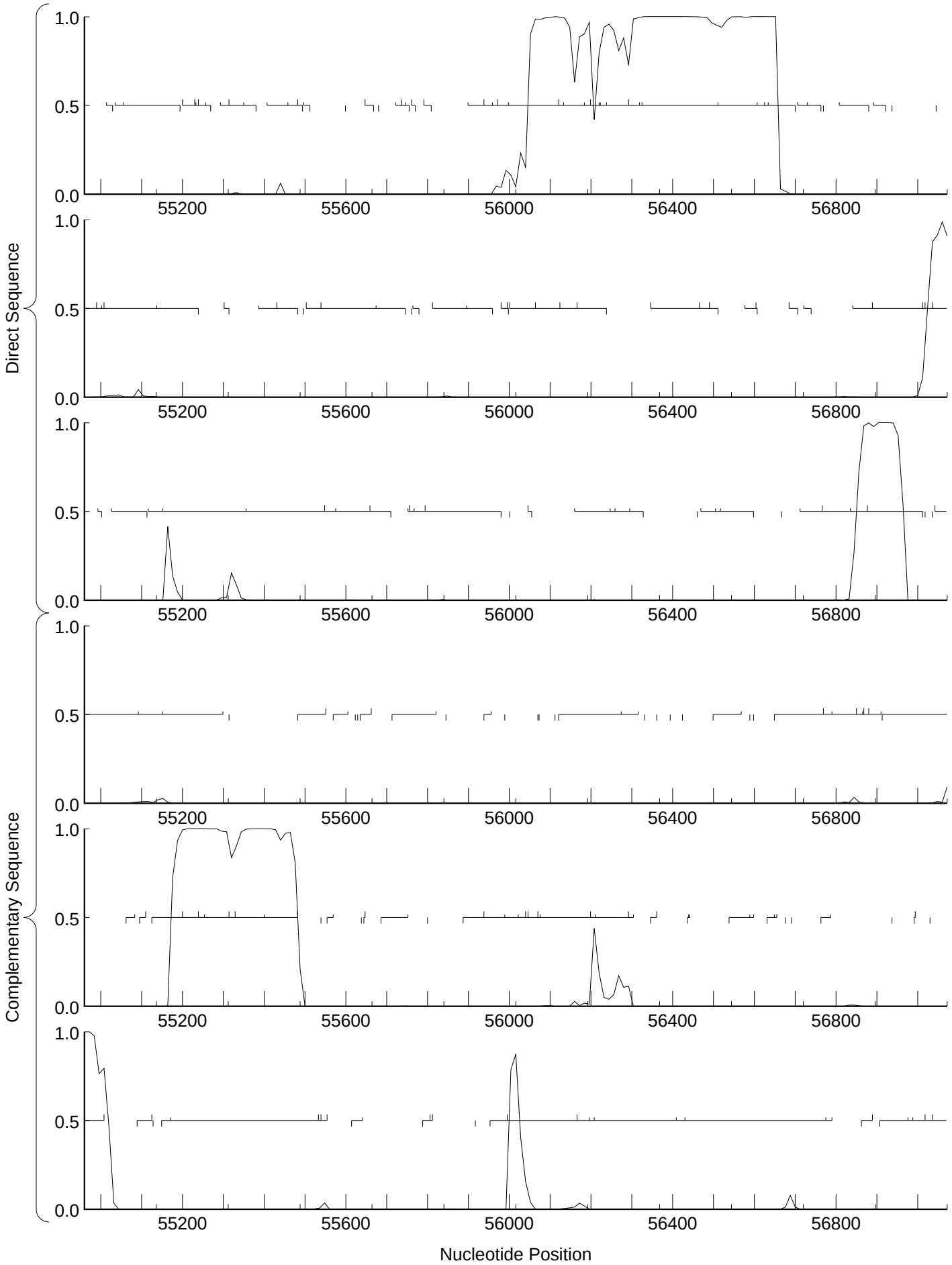
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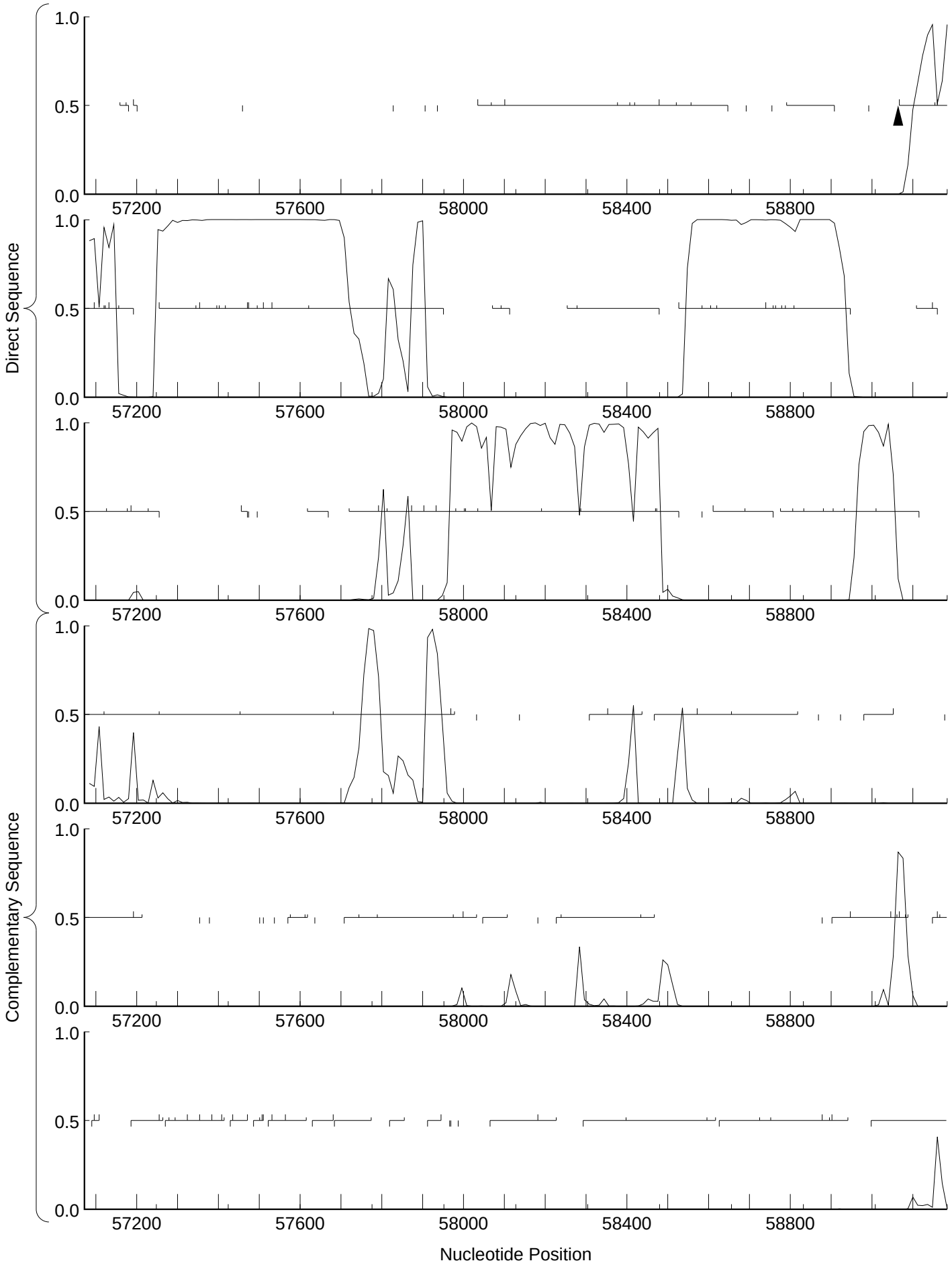
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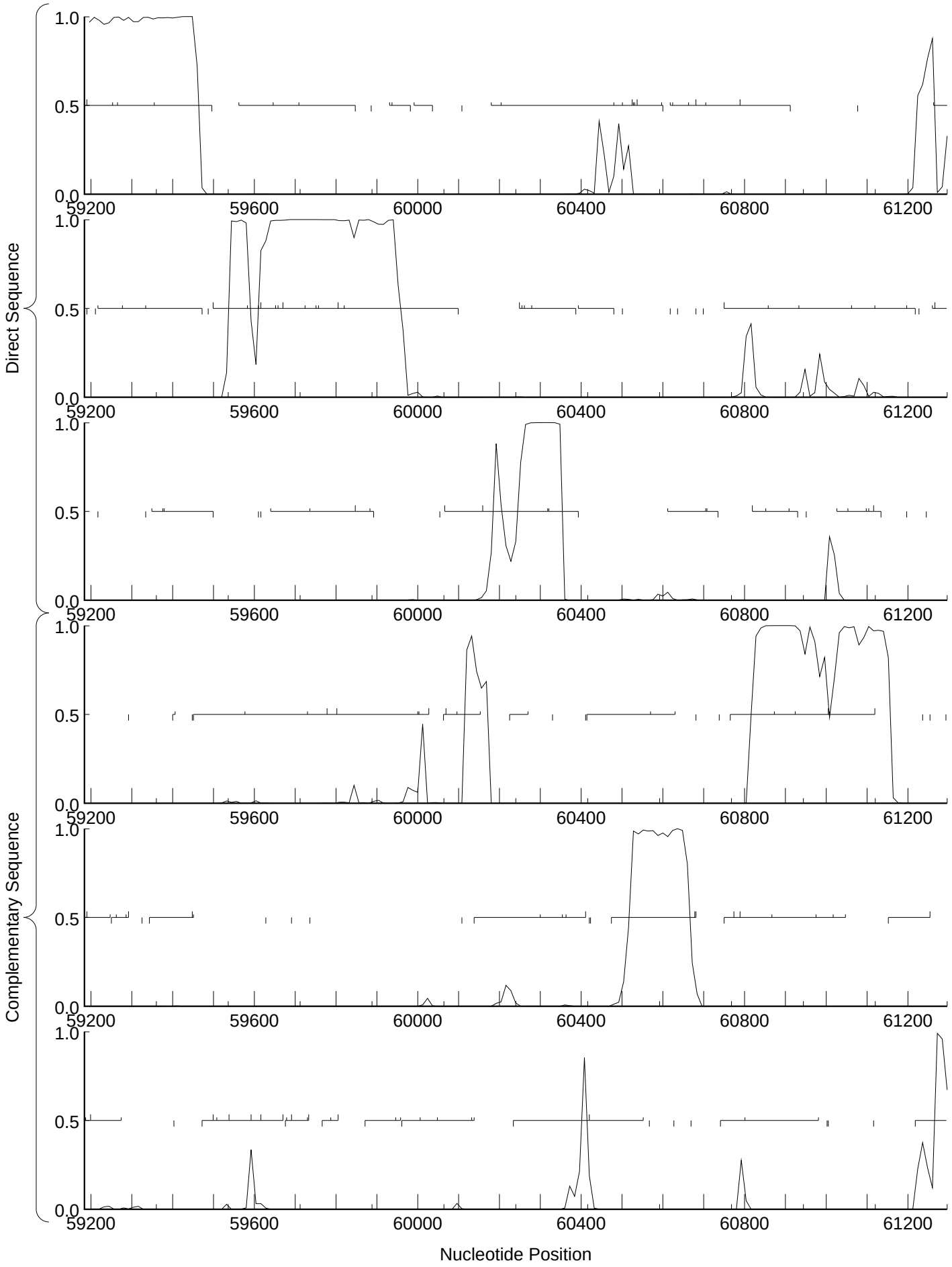
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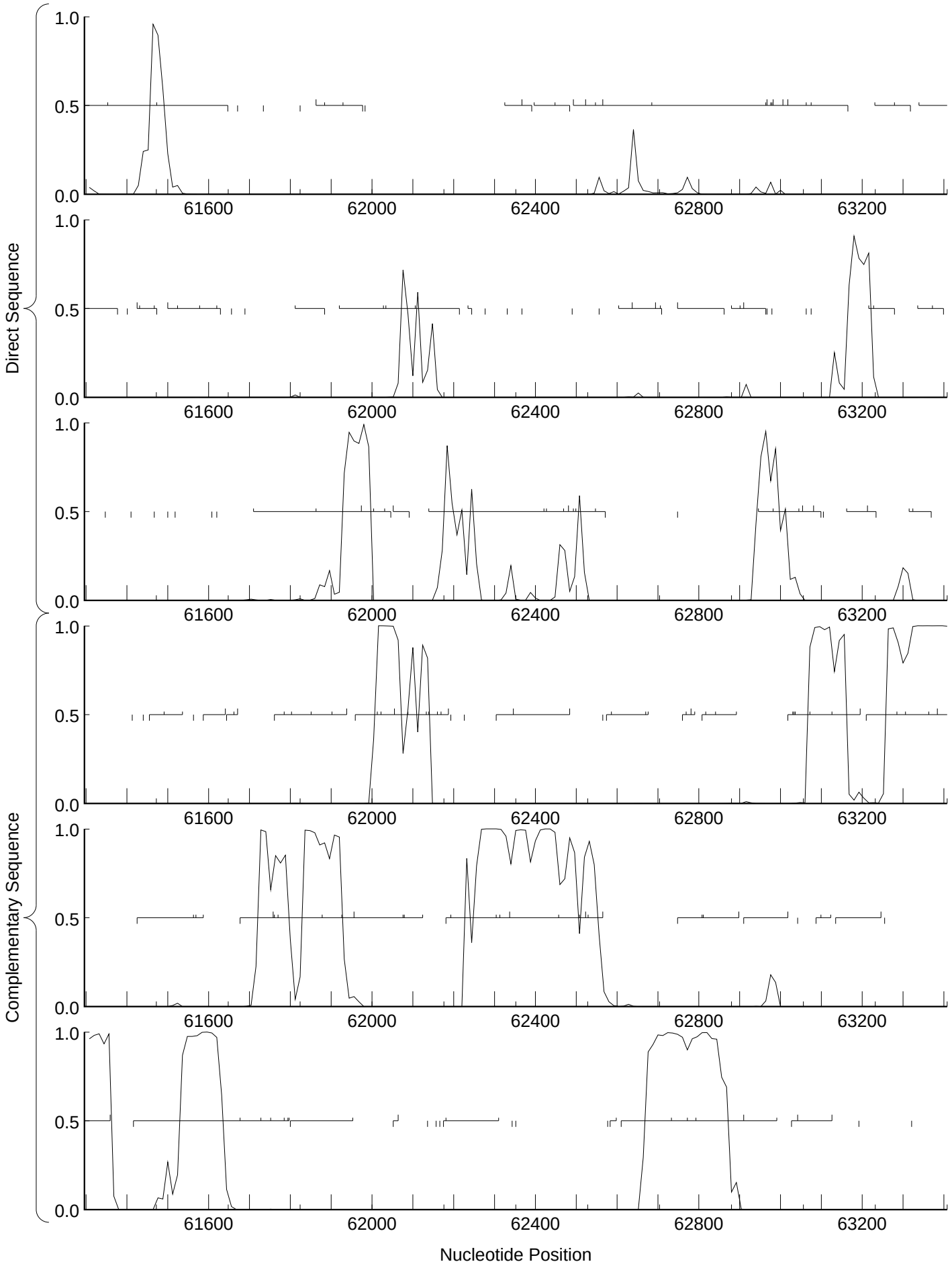
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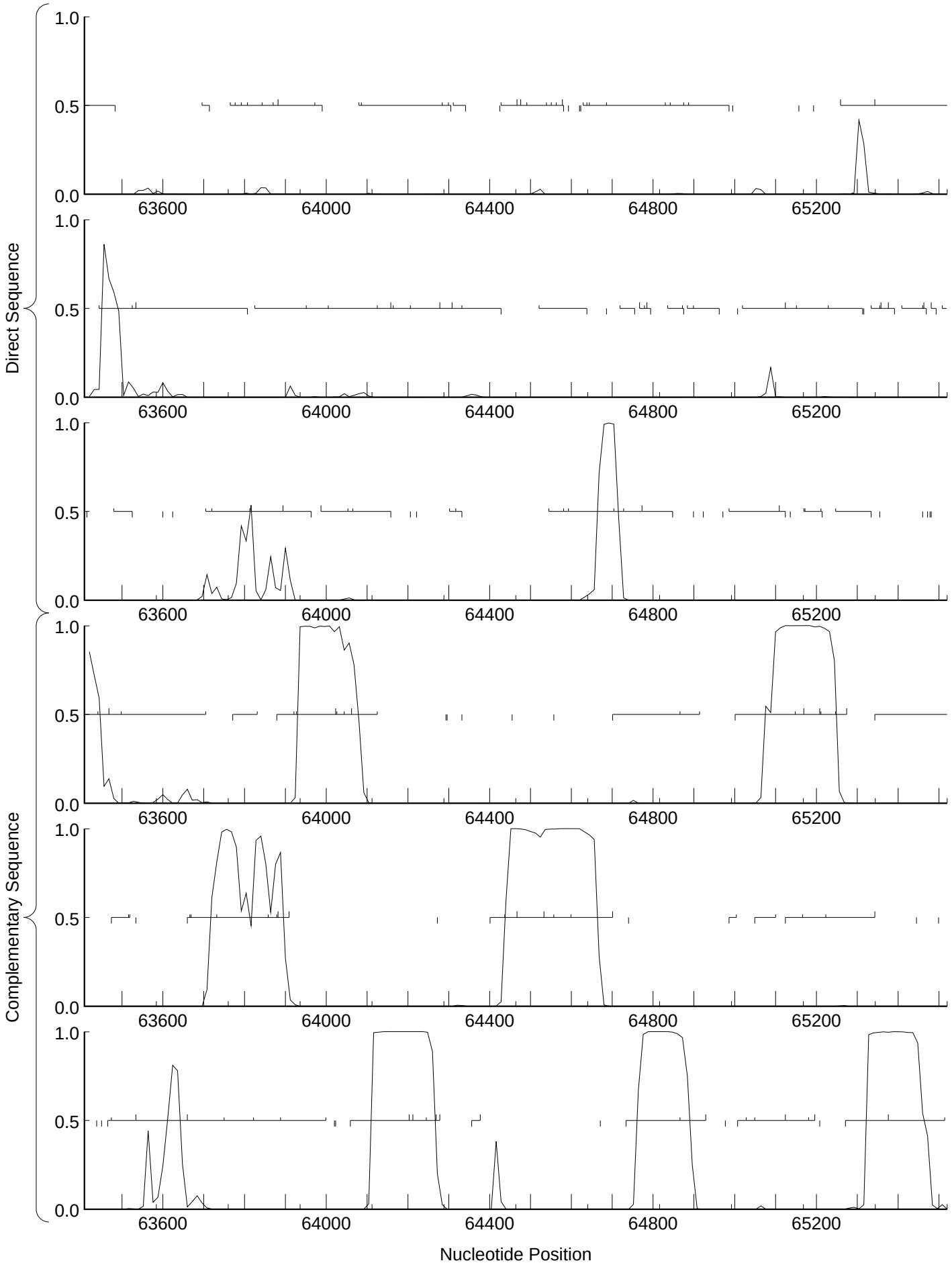


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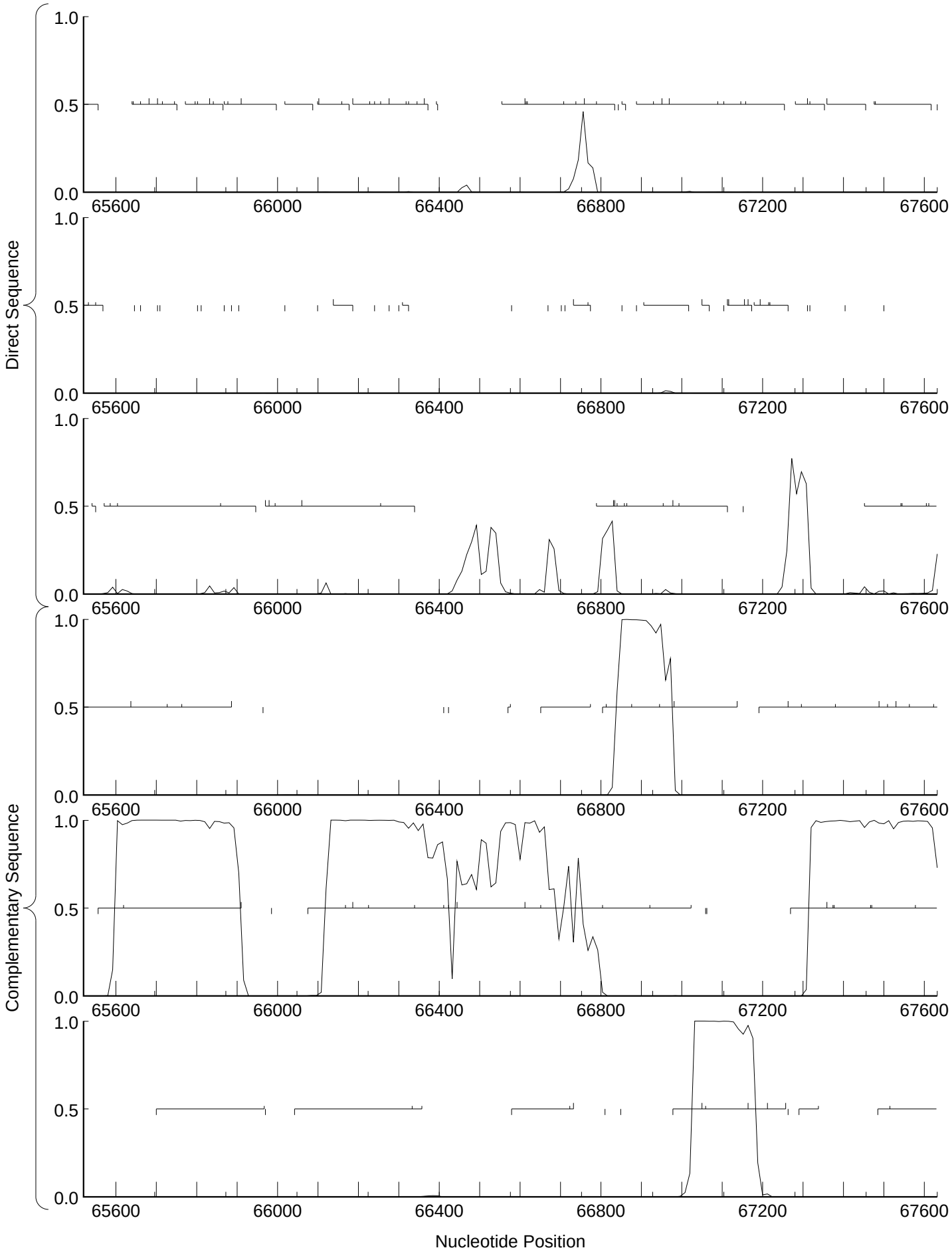


terium phage CharlieGBrown complete sequence, 68604 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 31/34




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
63600
64000
64400
64800
65200
Nucleotide Position

terium phage CharlieGBrown complete sequence, 68604 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 33/34



terium phage CharlieGBrown complete sequence, 68604 bp, circularly permuted, Cluster B1, Order 4, Window 96, Step 12, 34/34

