GeneMark

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

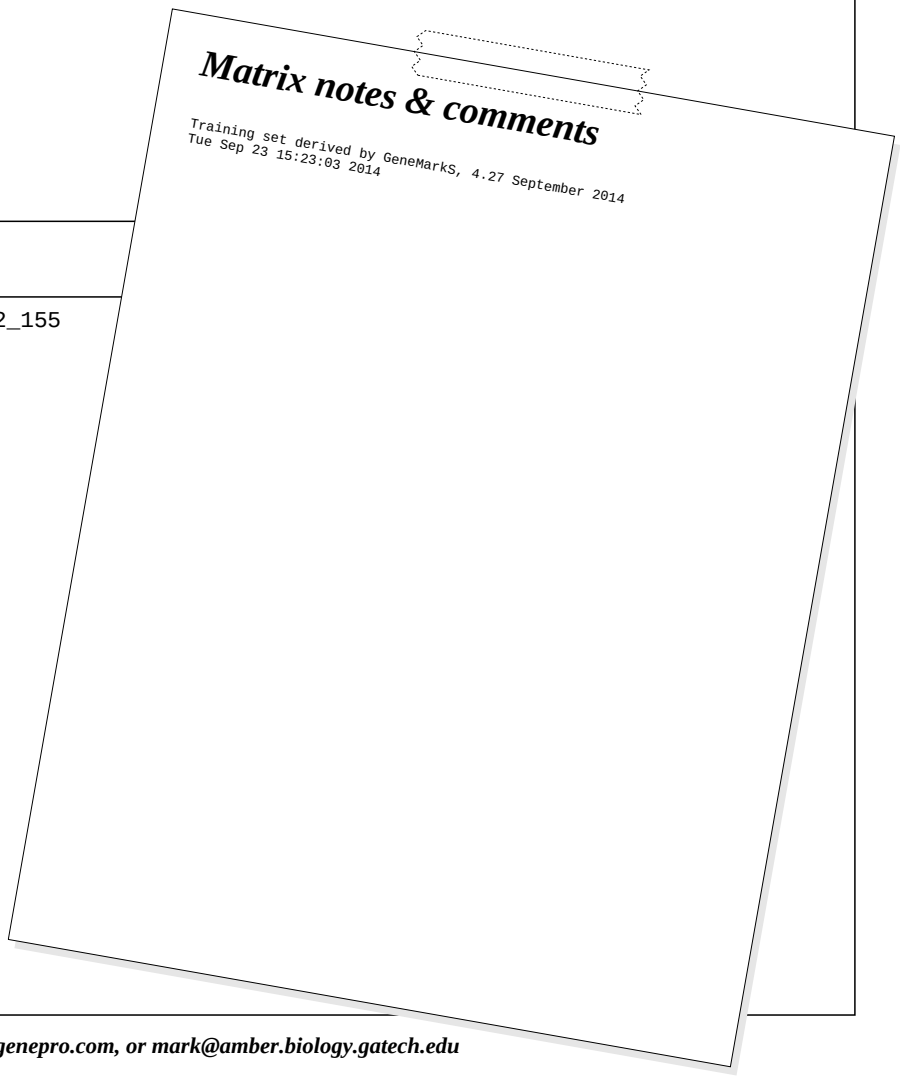
PROGRAM INFORMATION

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Pages            :  75
Sequence Length  :  155943 bp
GC Content       :   64.75%

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,
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MATRIX INFORMATION

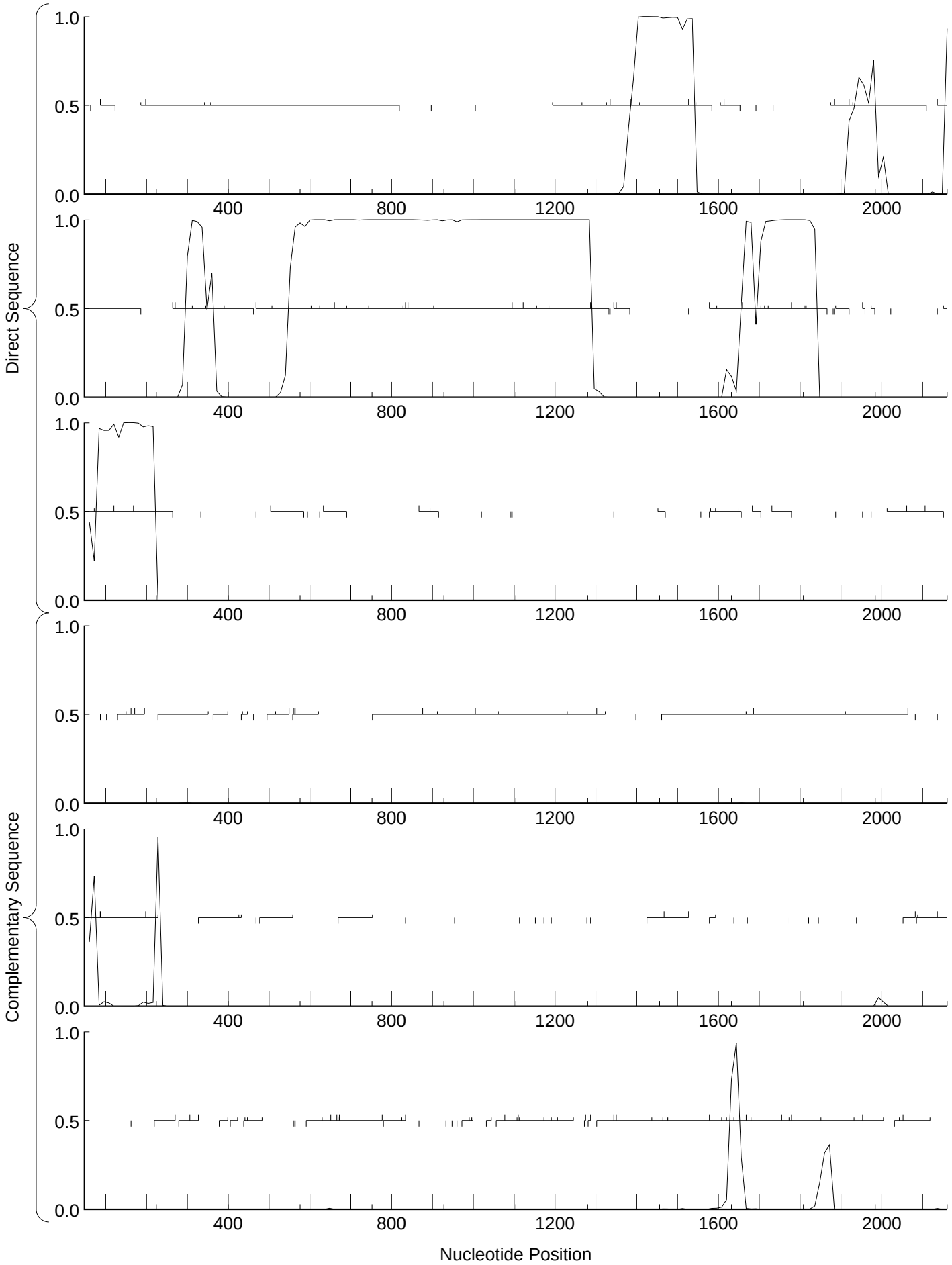
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Author     :  -

Order      :  4
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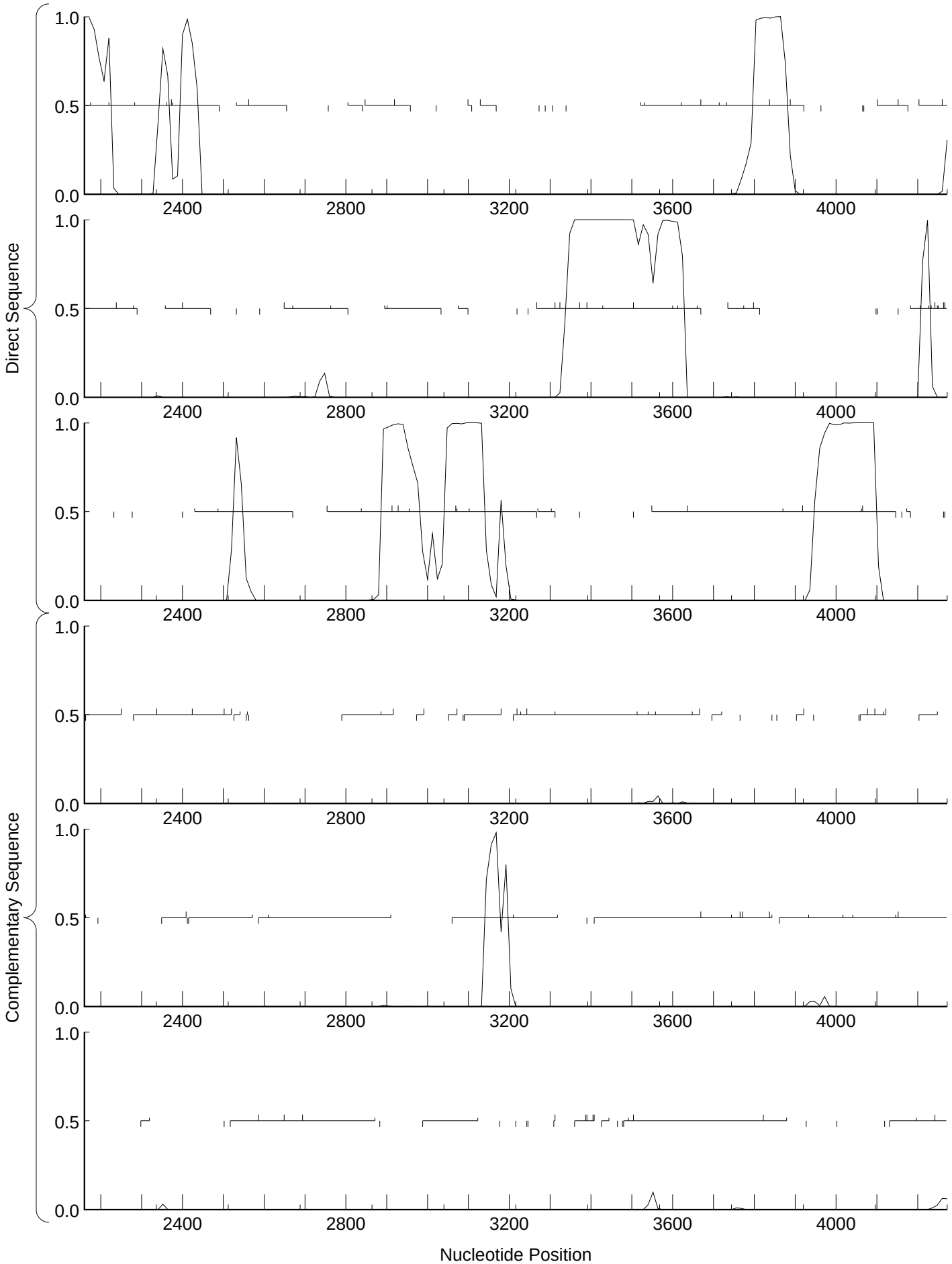

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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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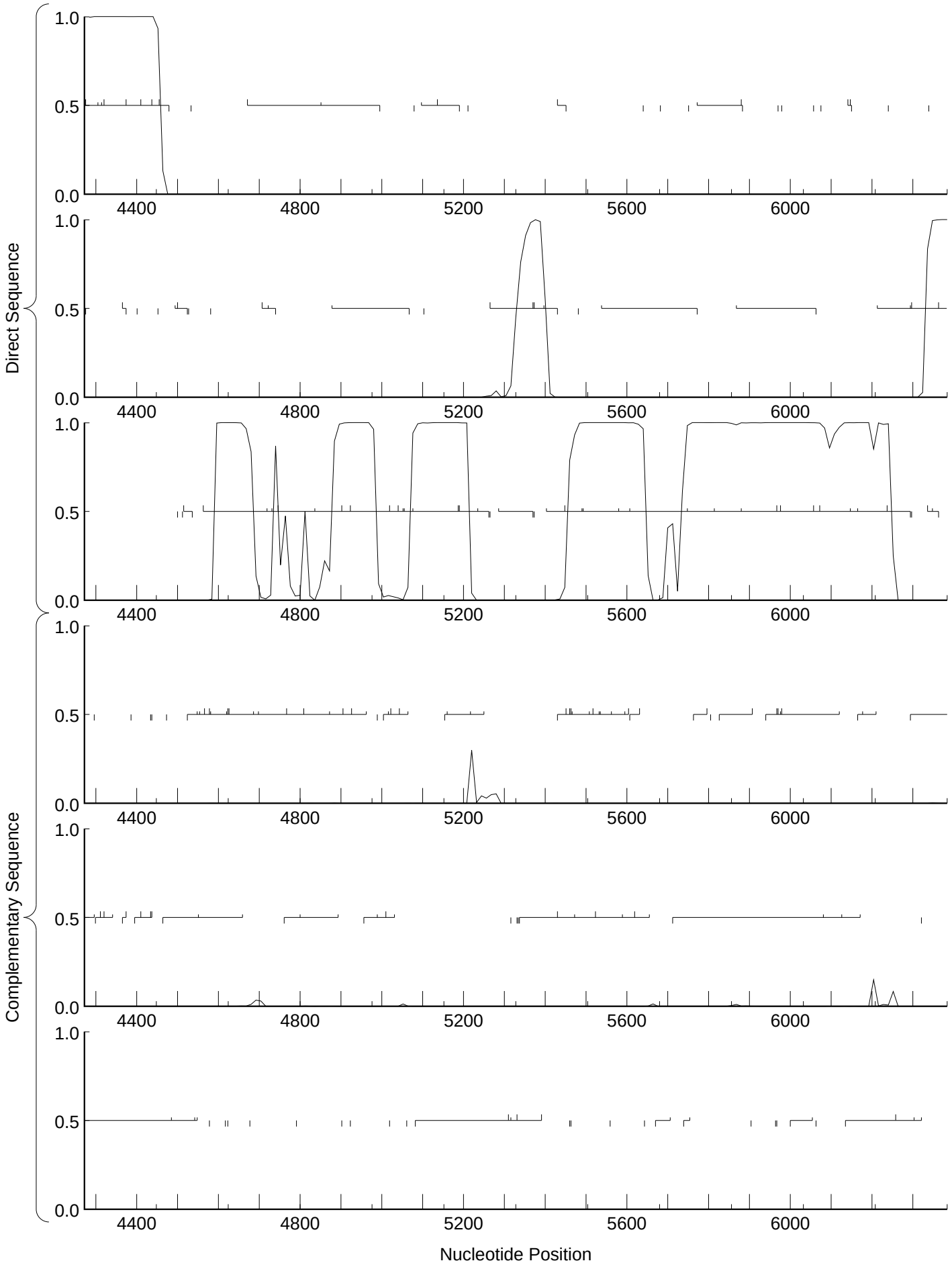

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

terium phage NuevoMundo Complete Sequence, 155943 bp, Circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 2/75

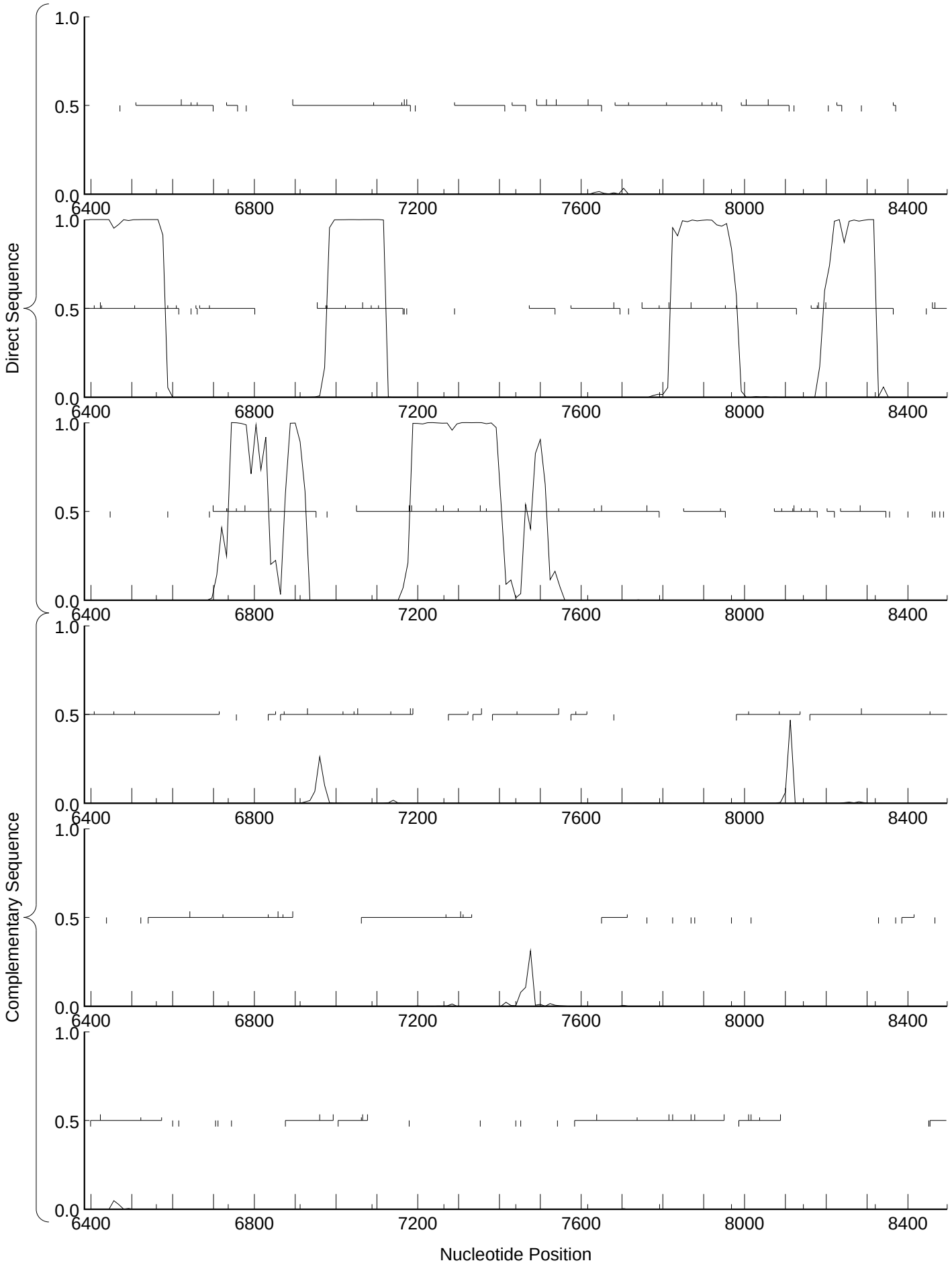


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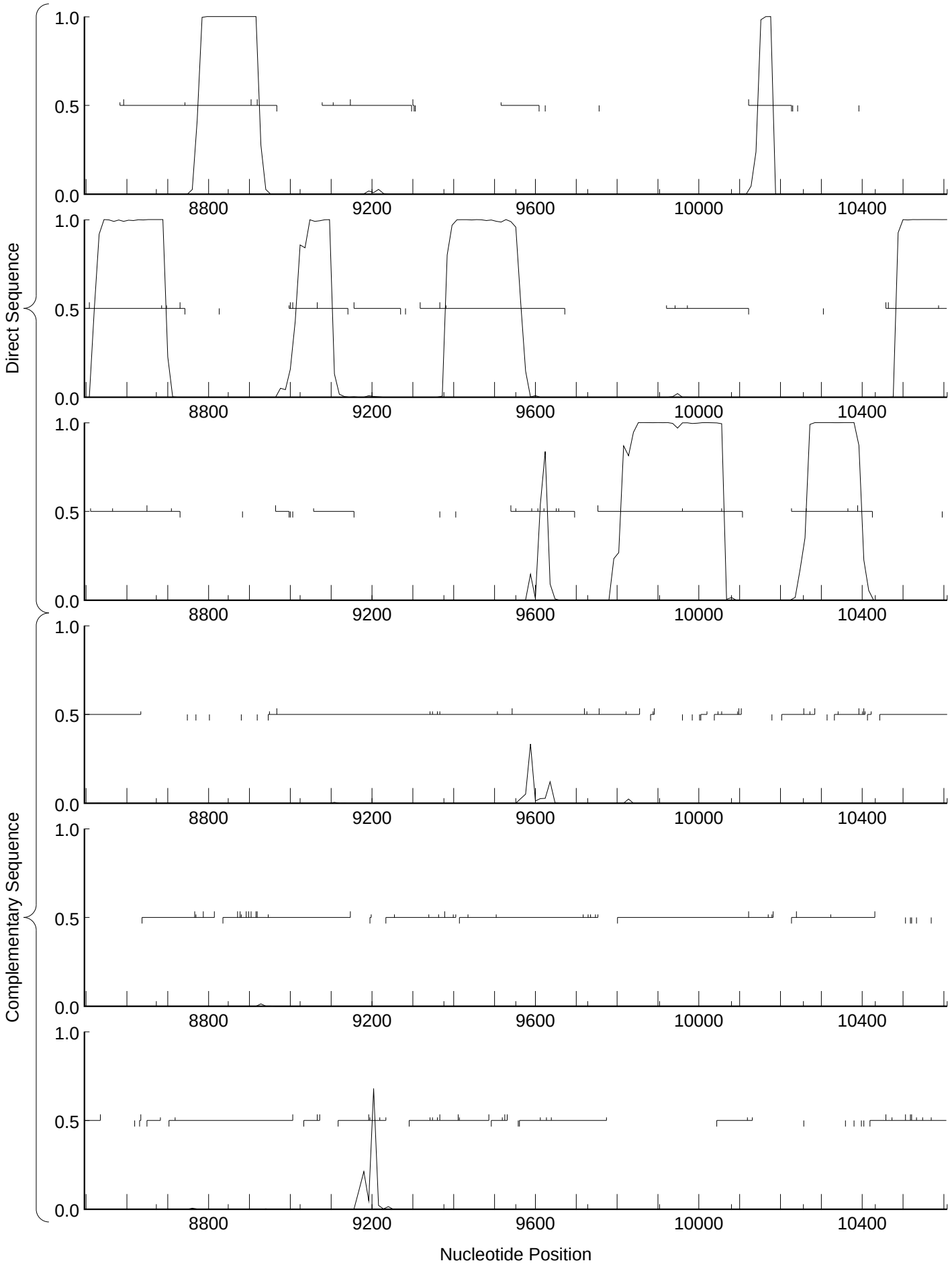



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Complementary Sequence
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5600
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Nucleotide Position

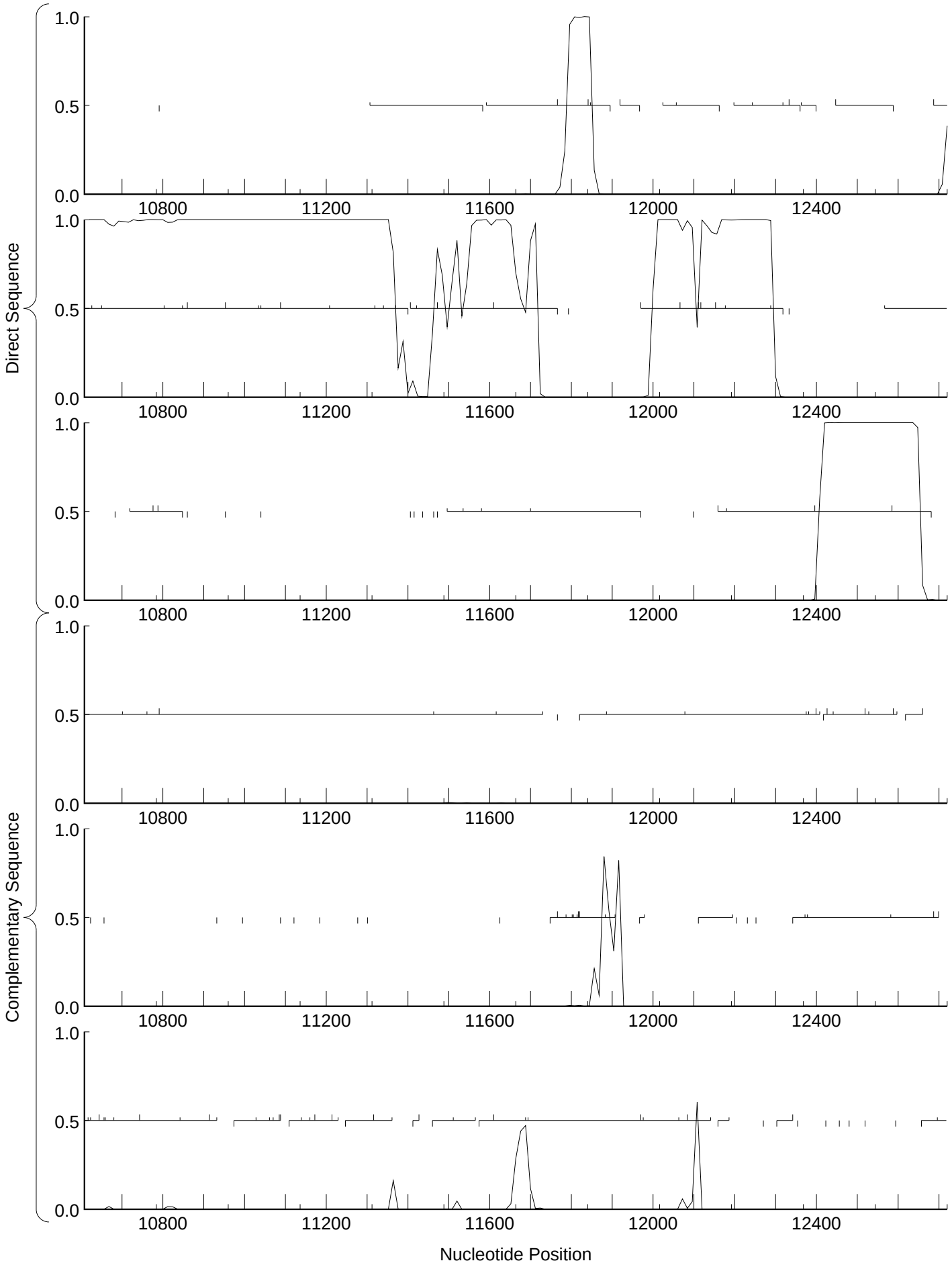
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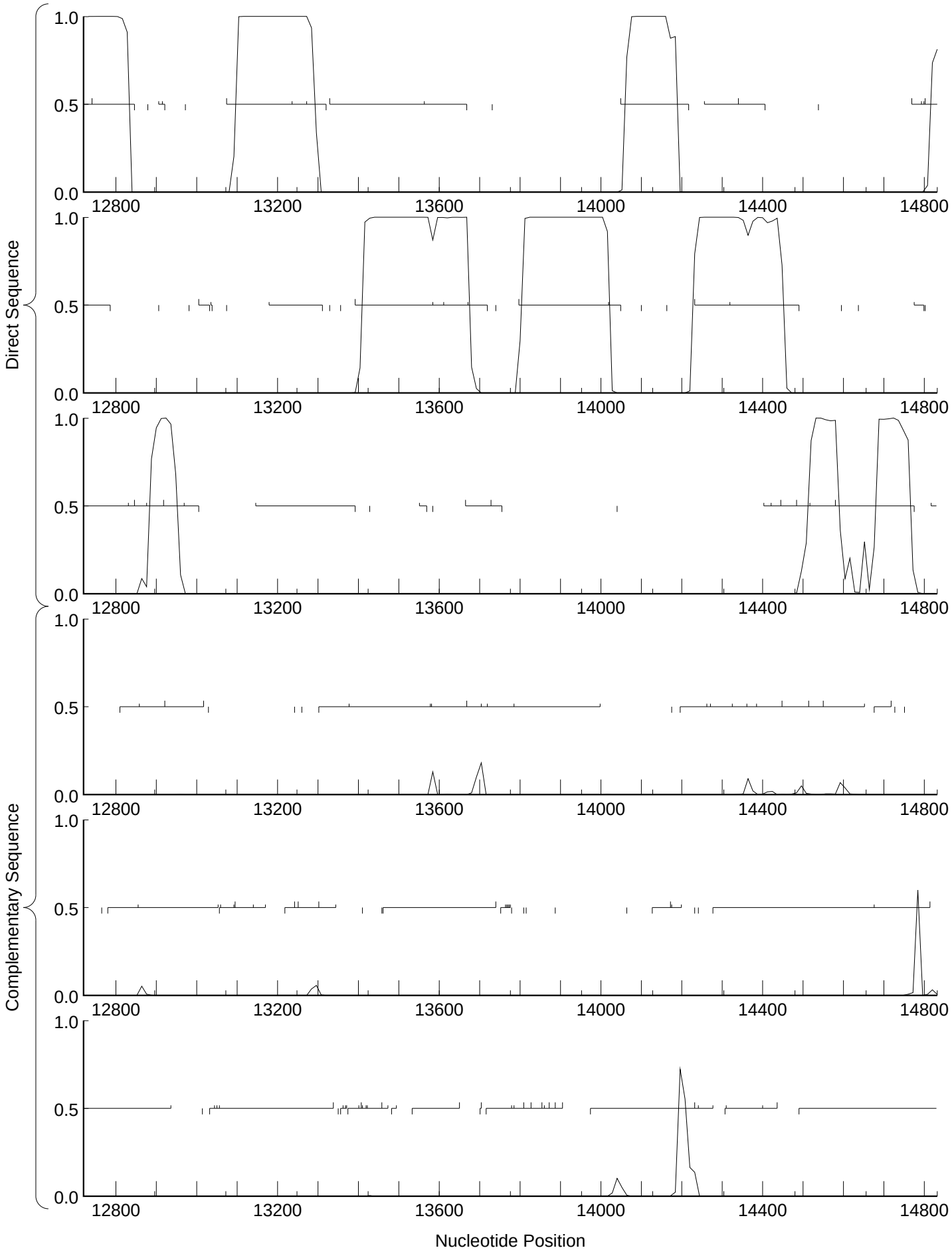
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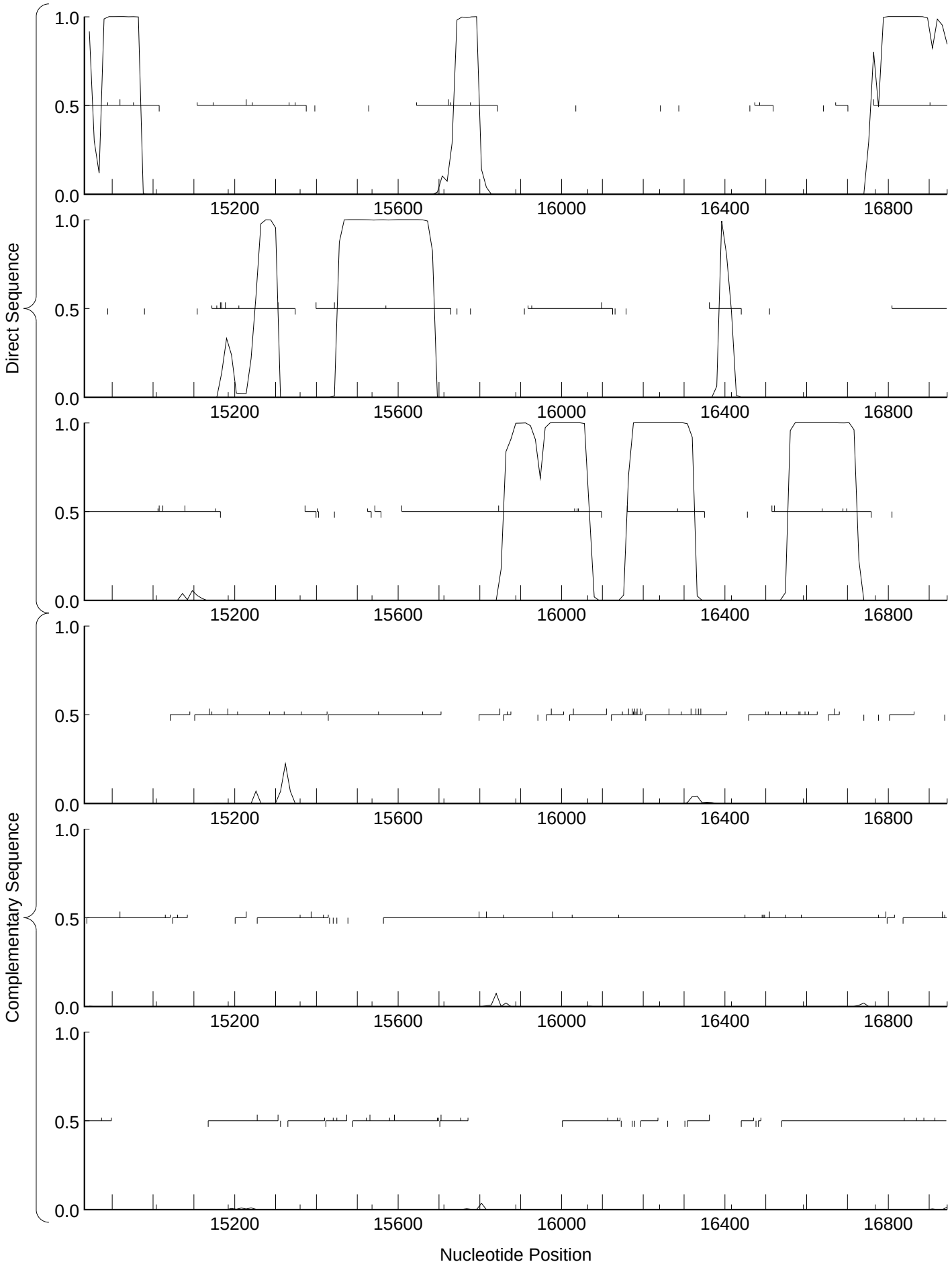
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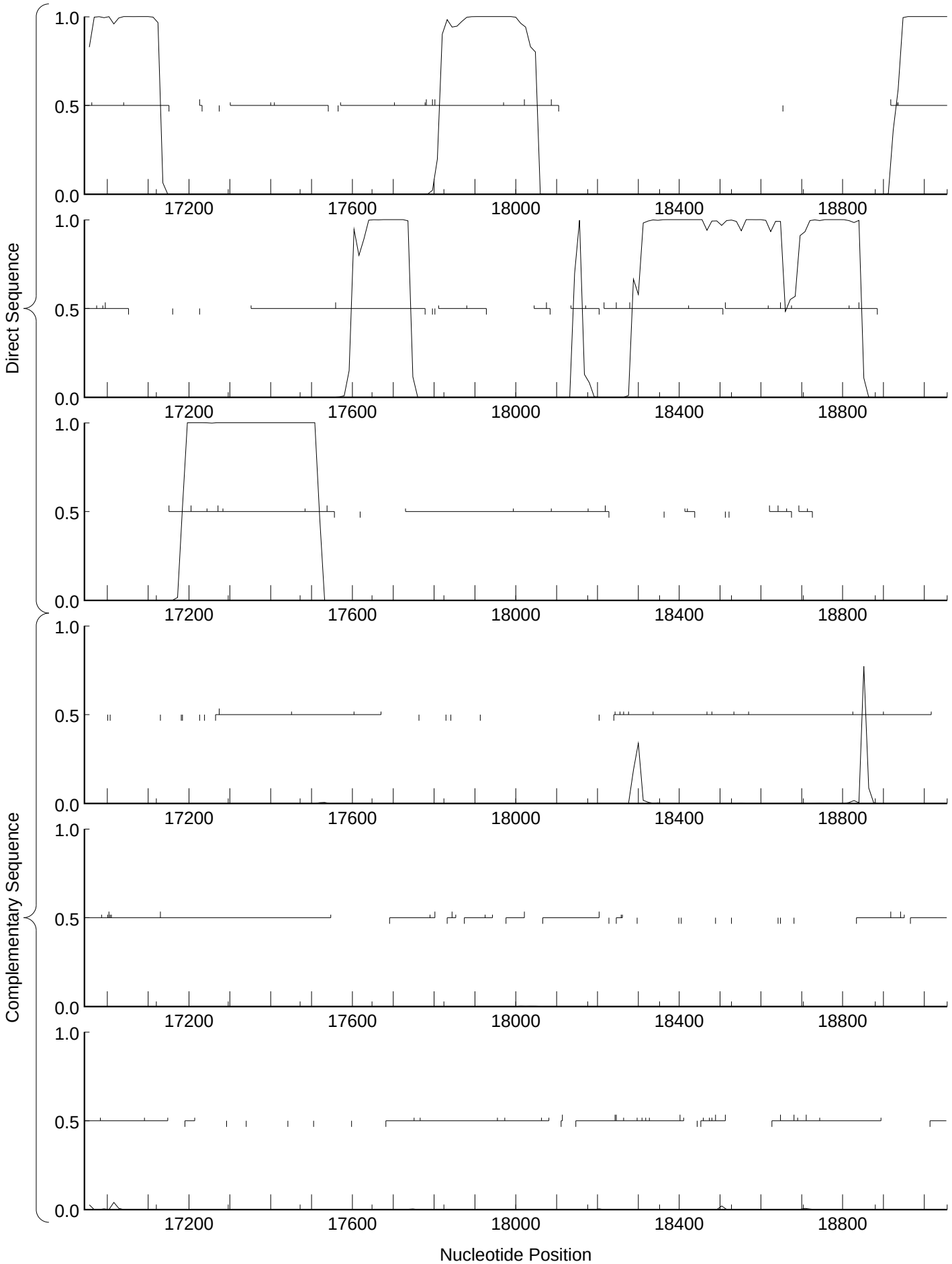
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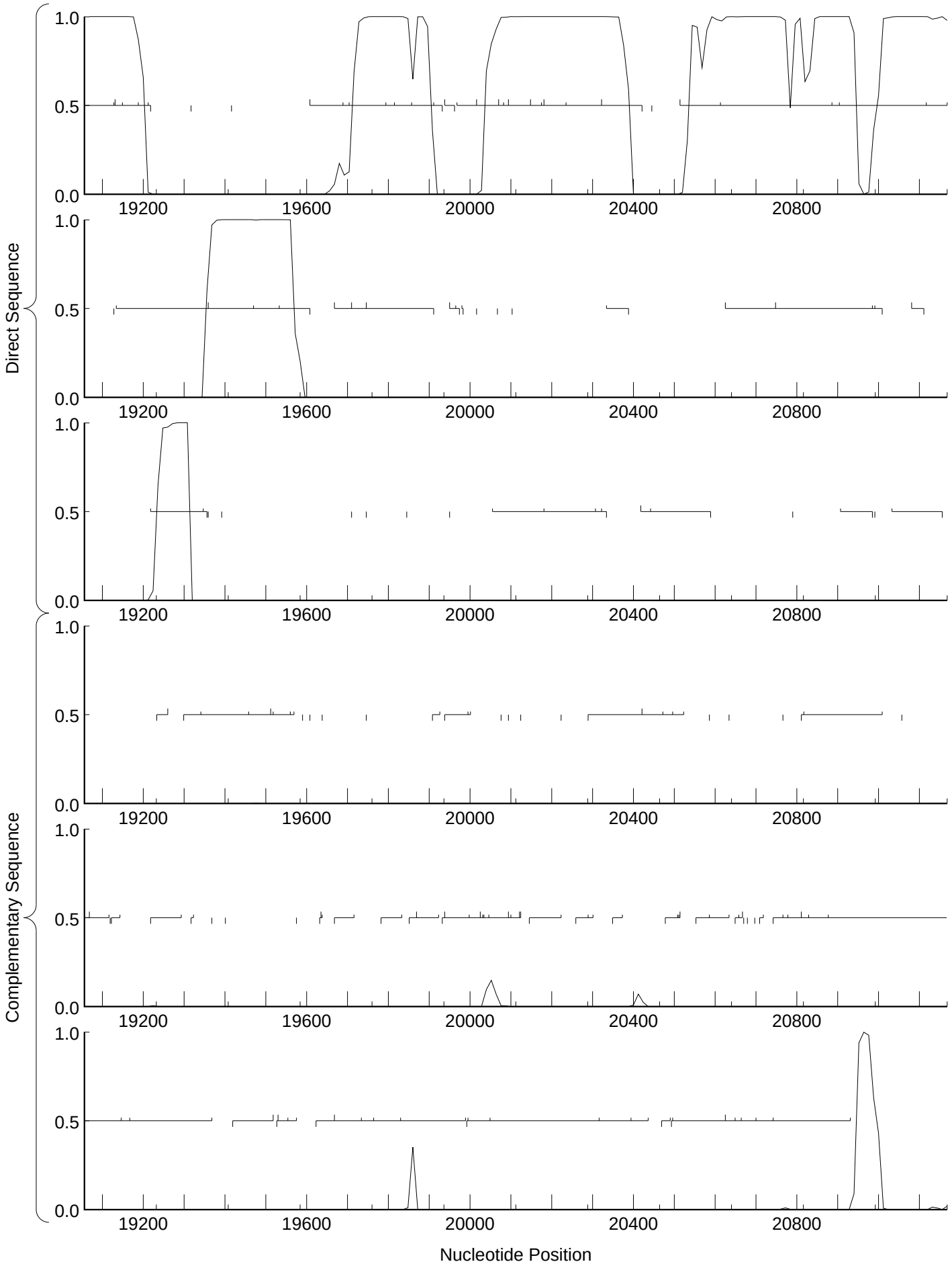
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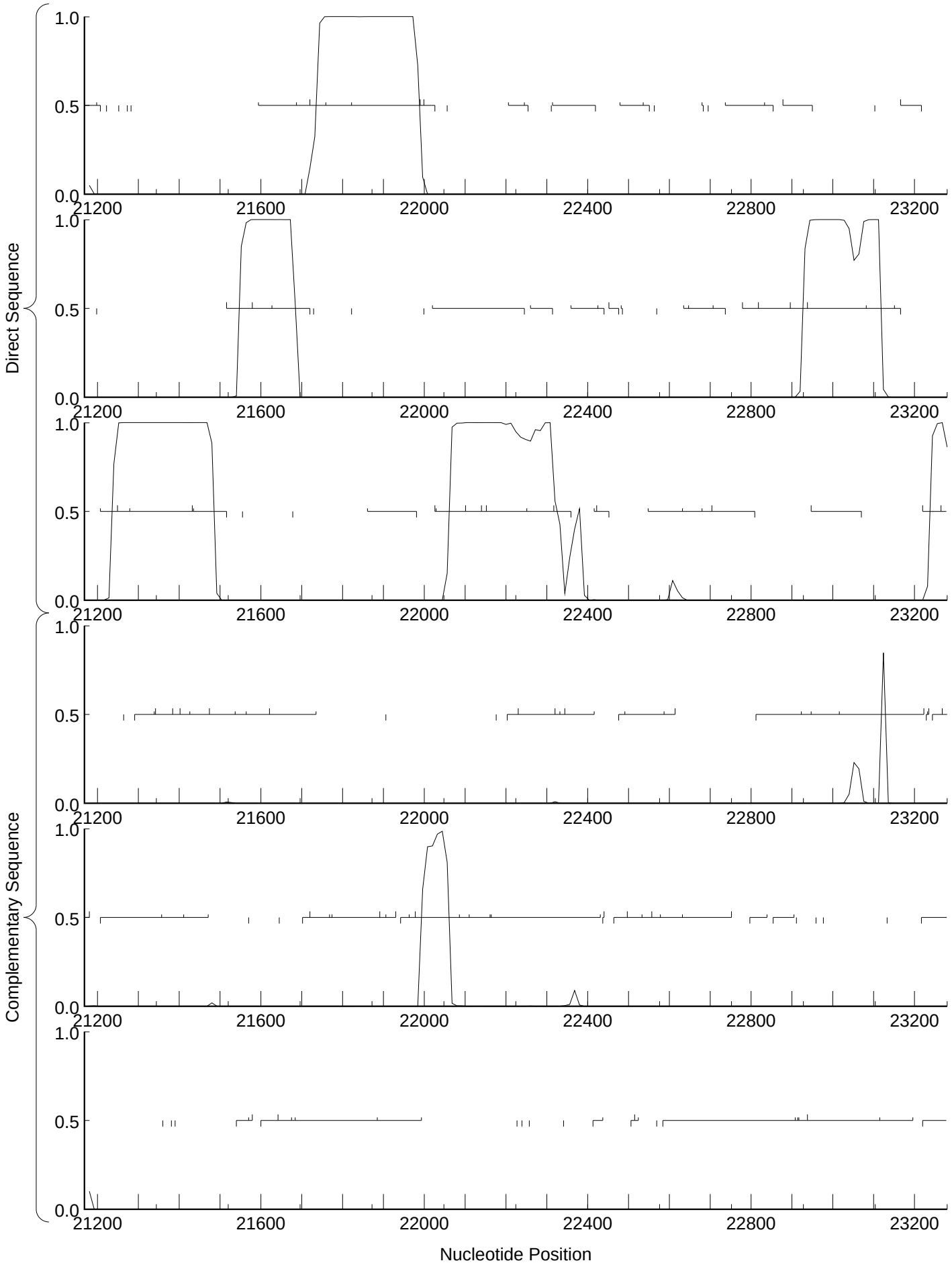


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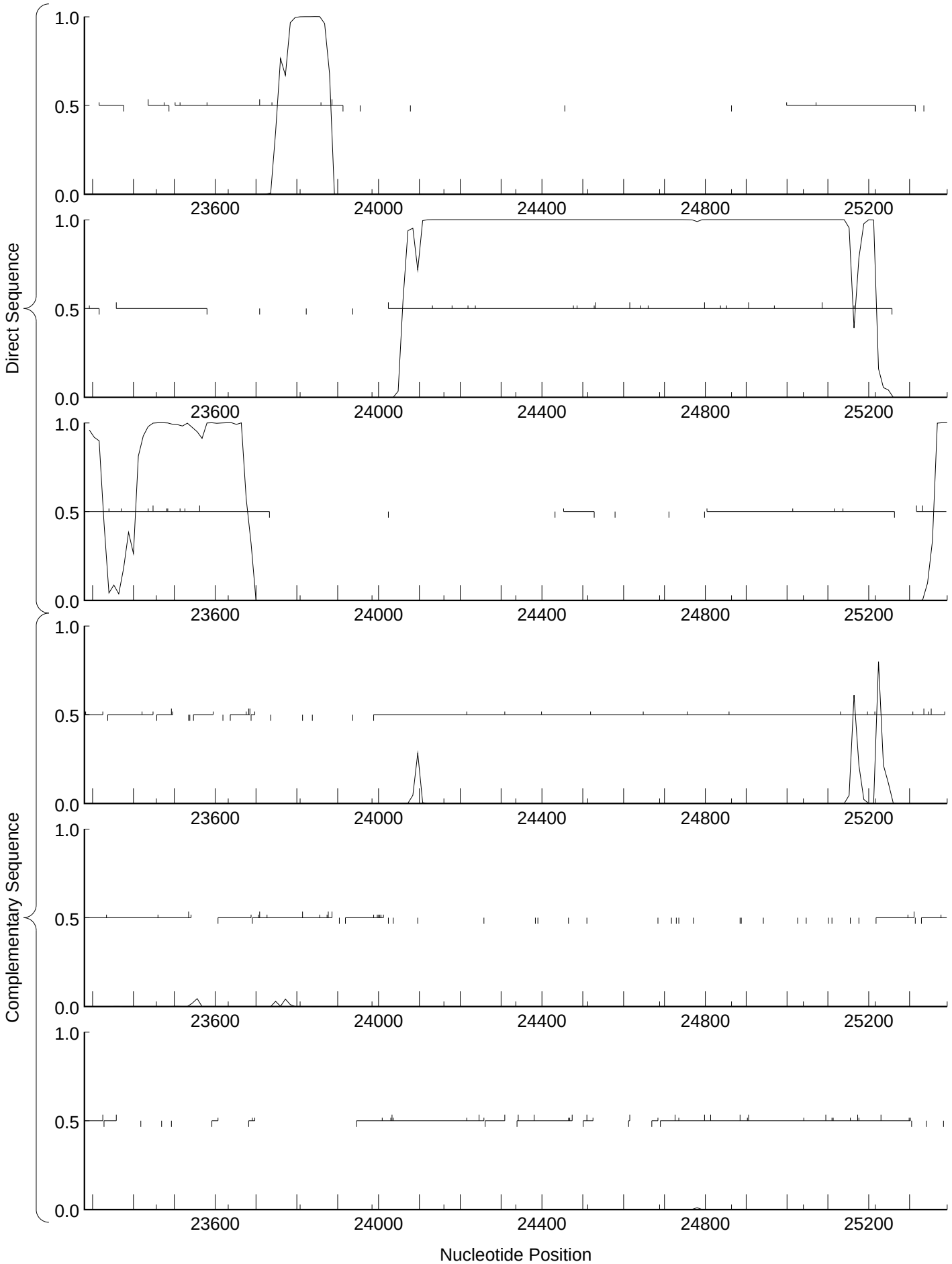


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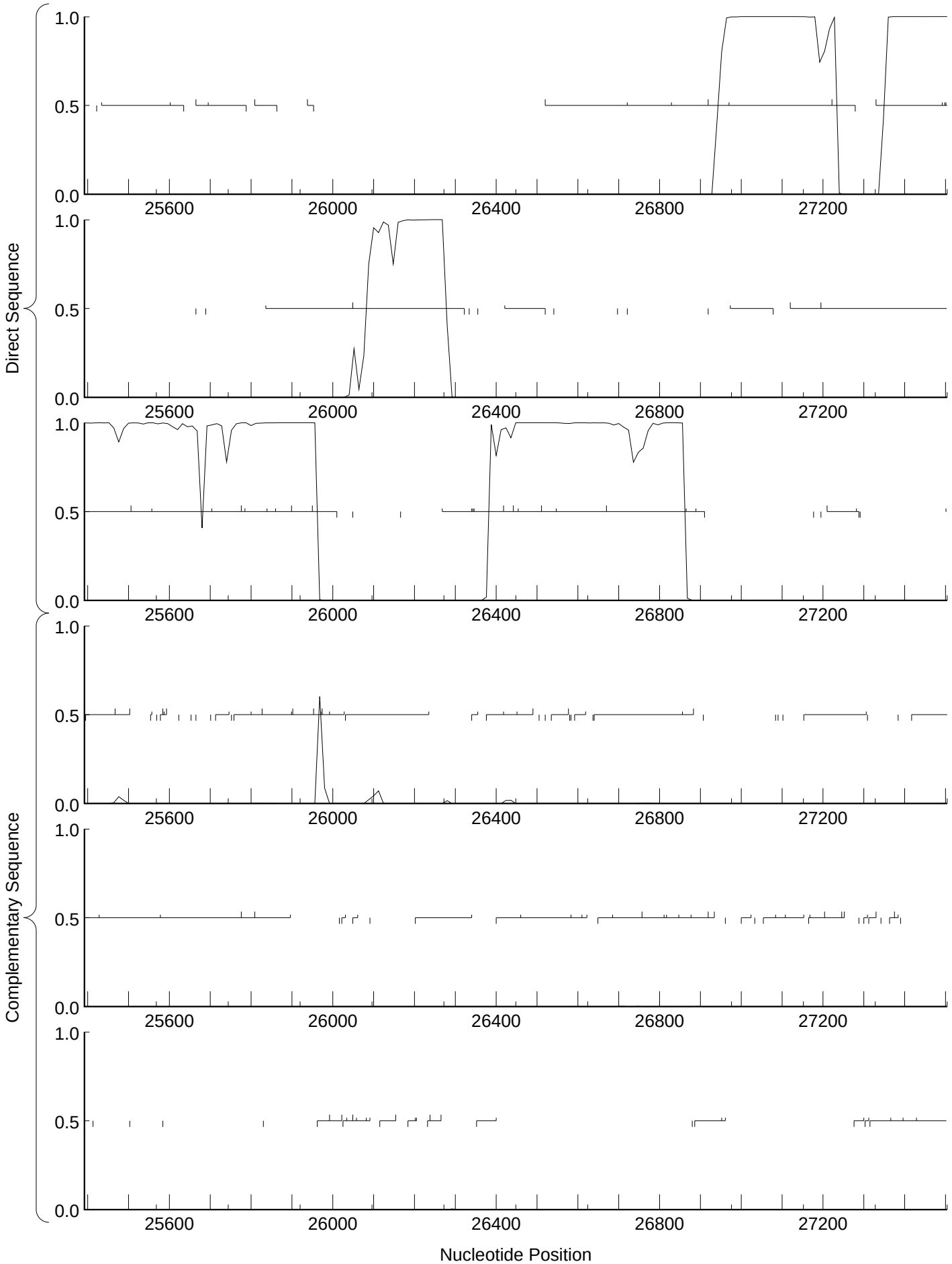



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Complementary Sequence
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22000
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22800
23200
Nucleotide Position

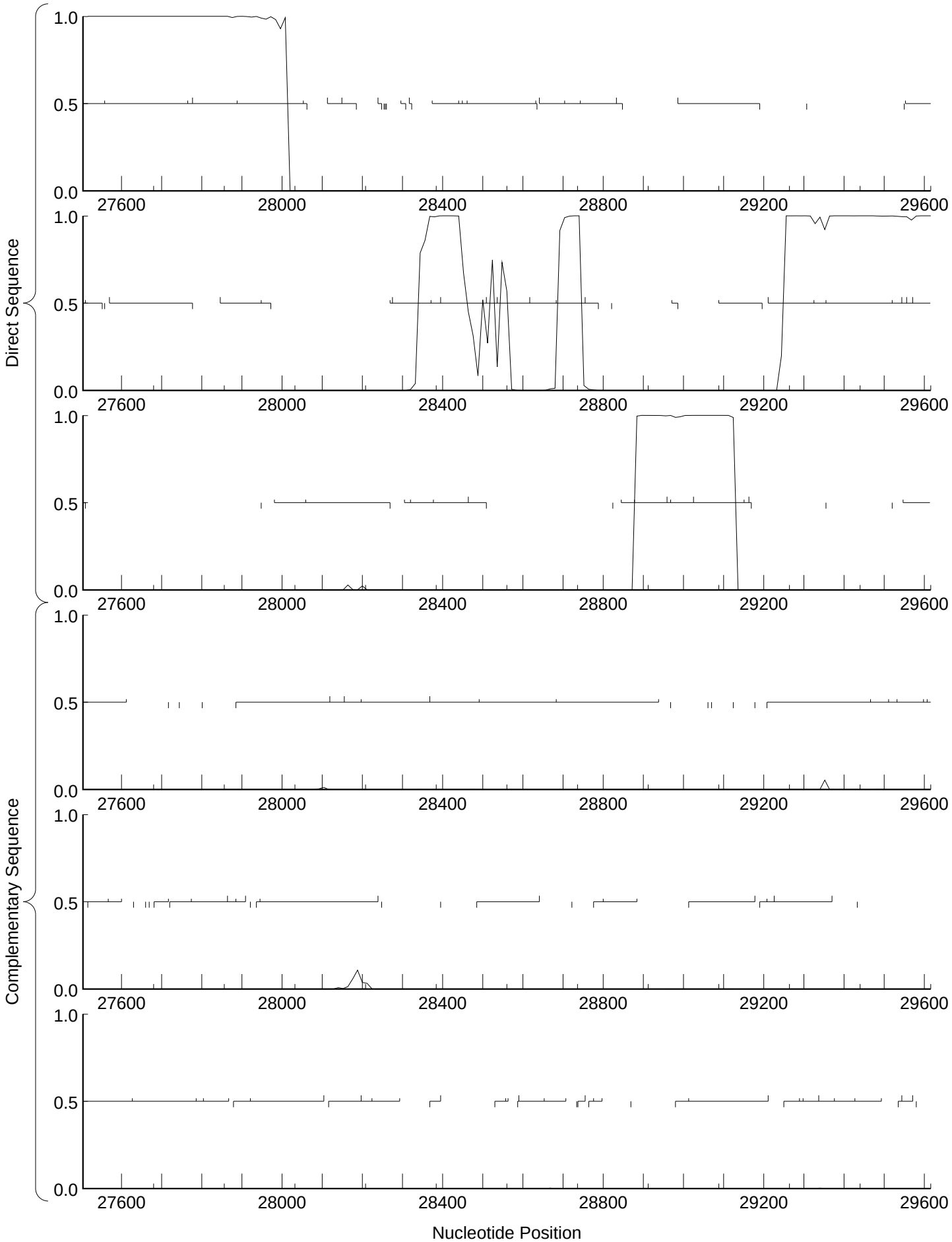
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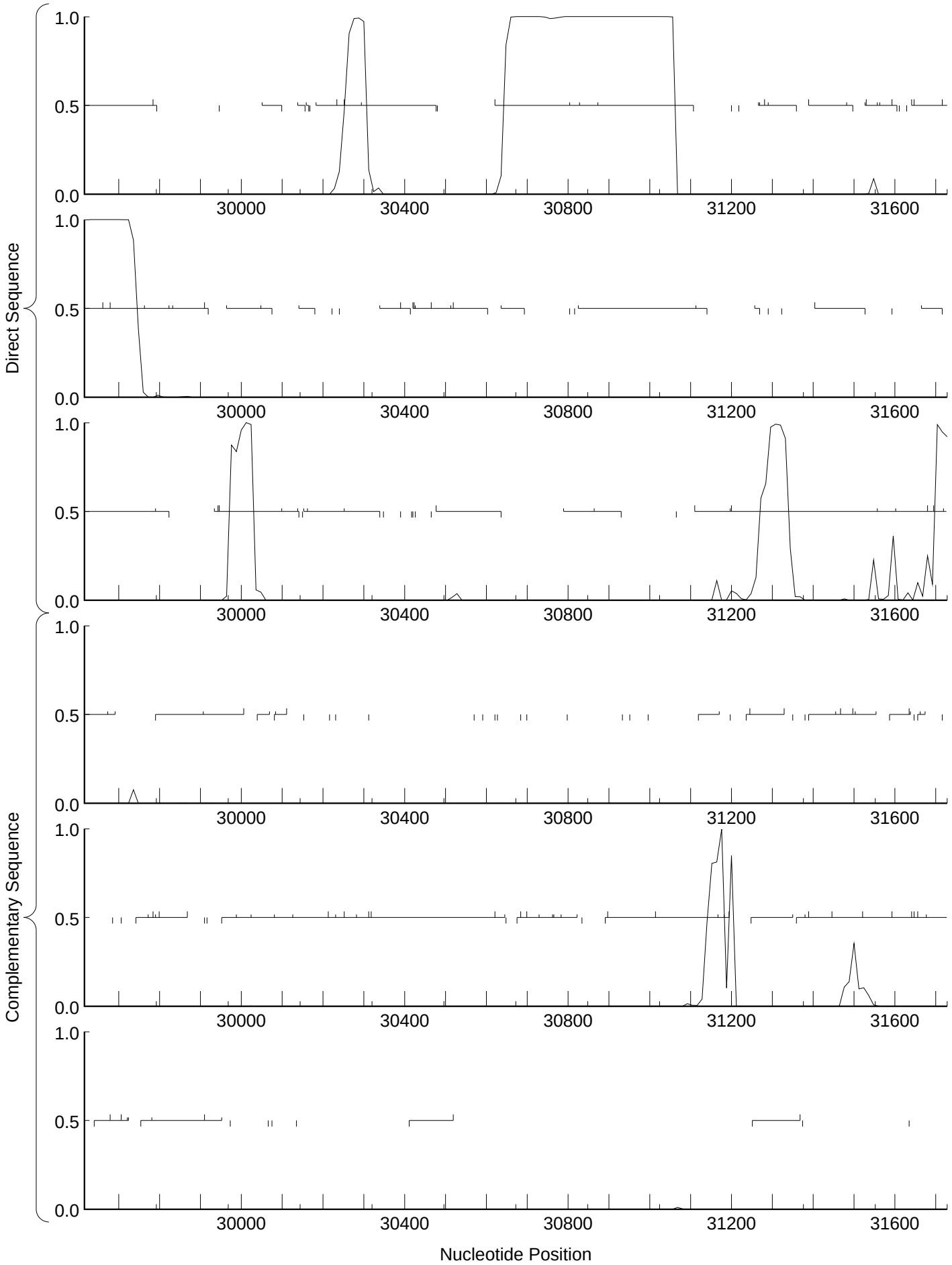
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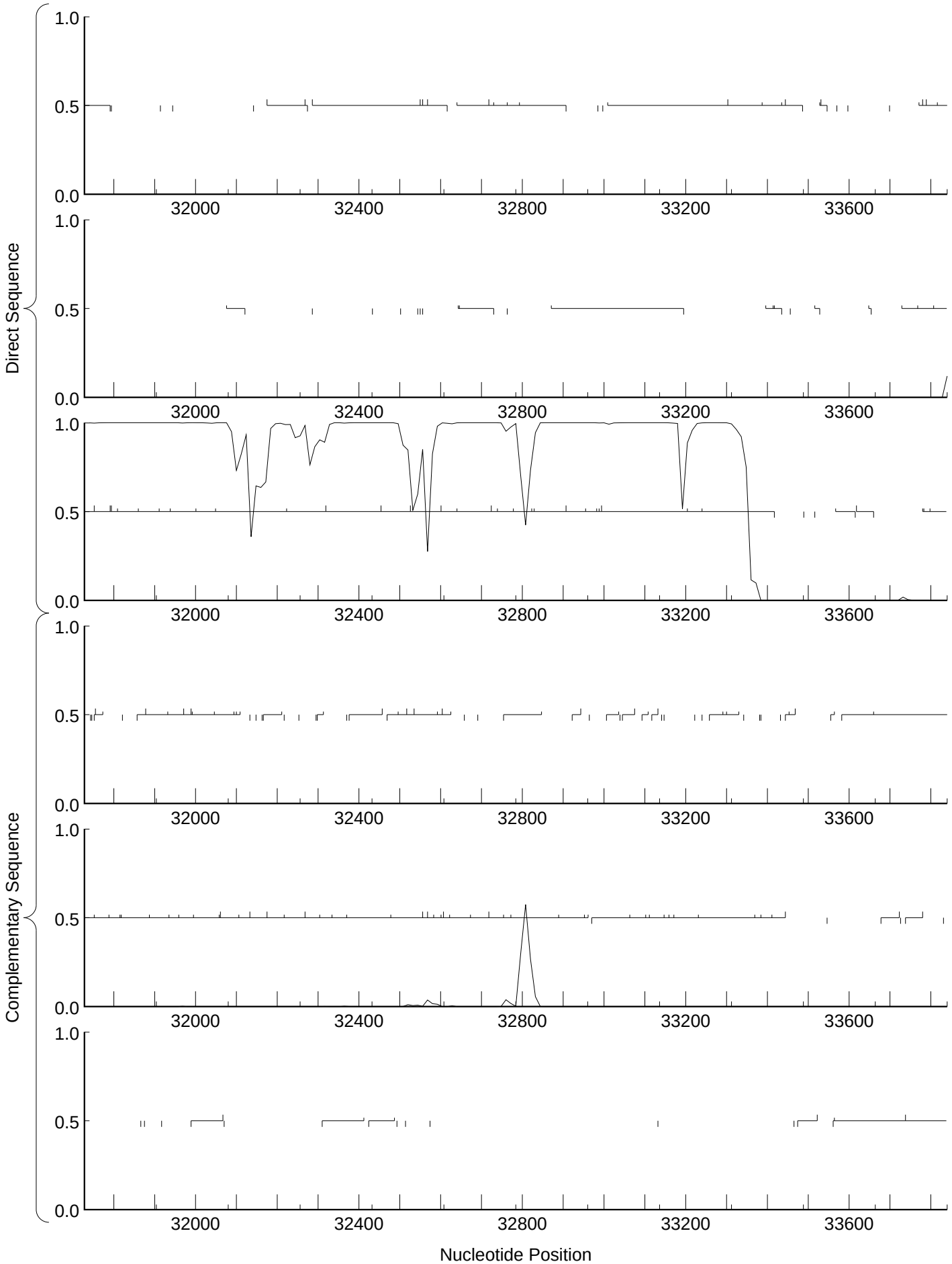
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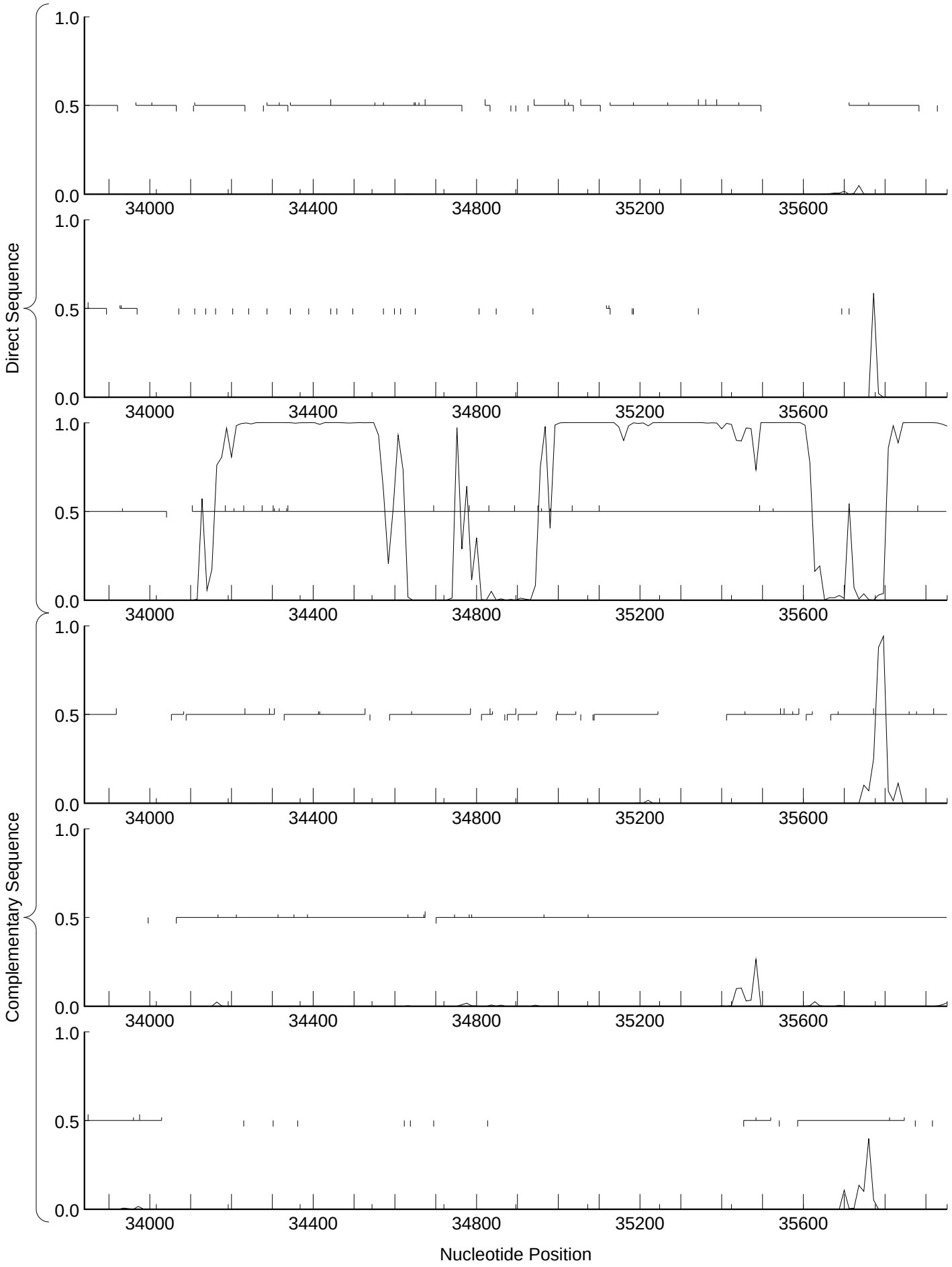
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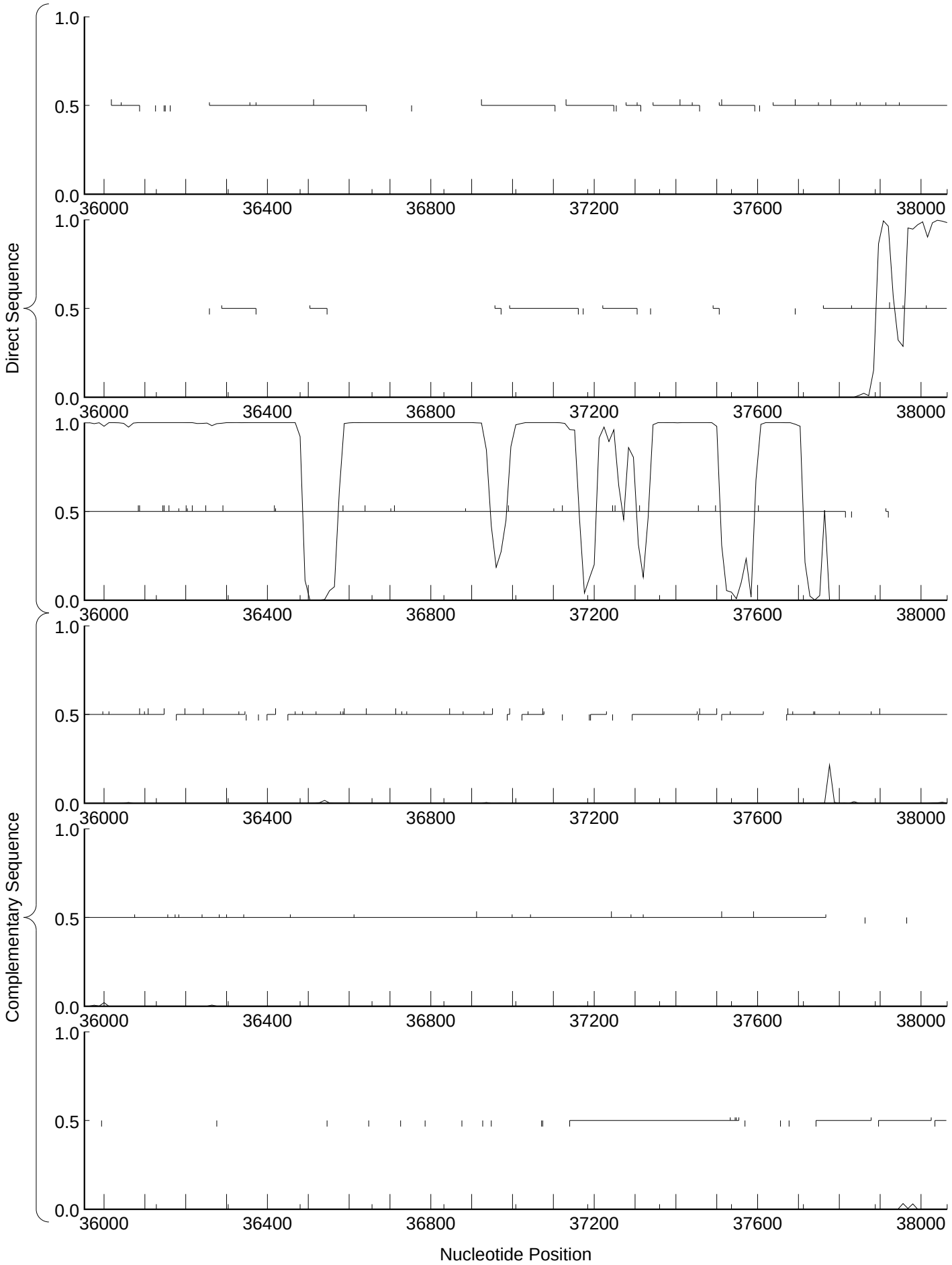
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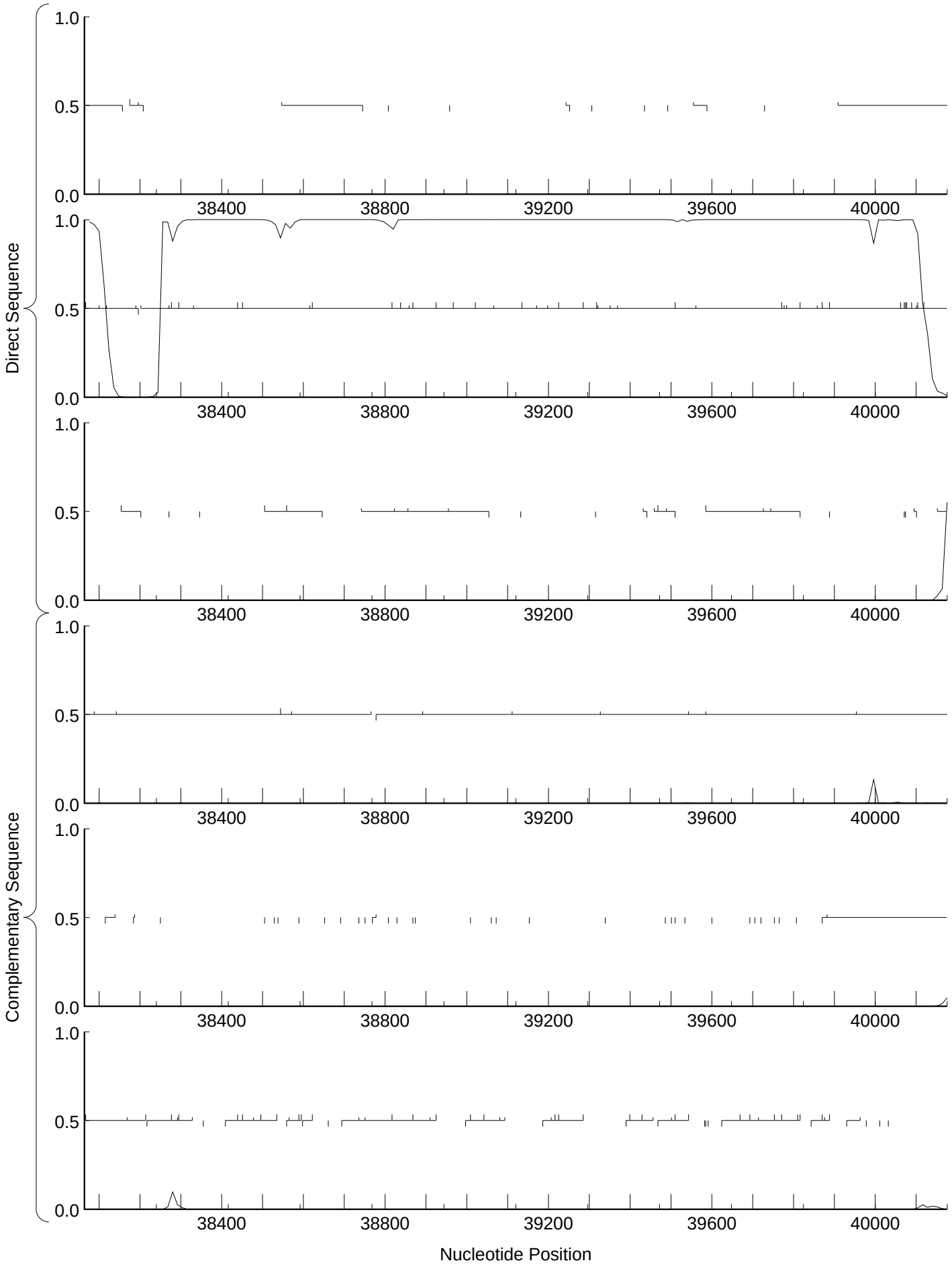


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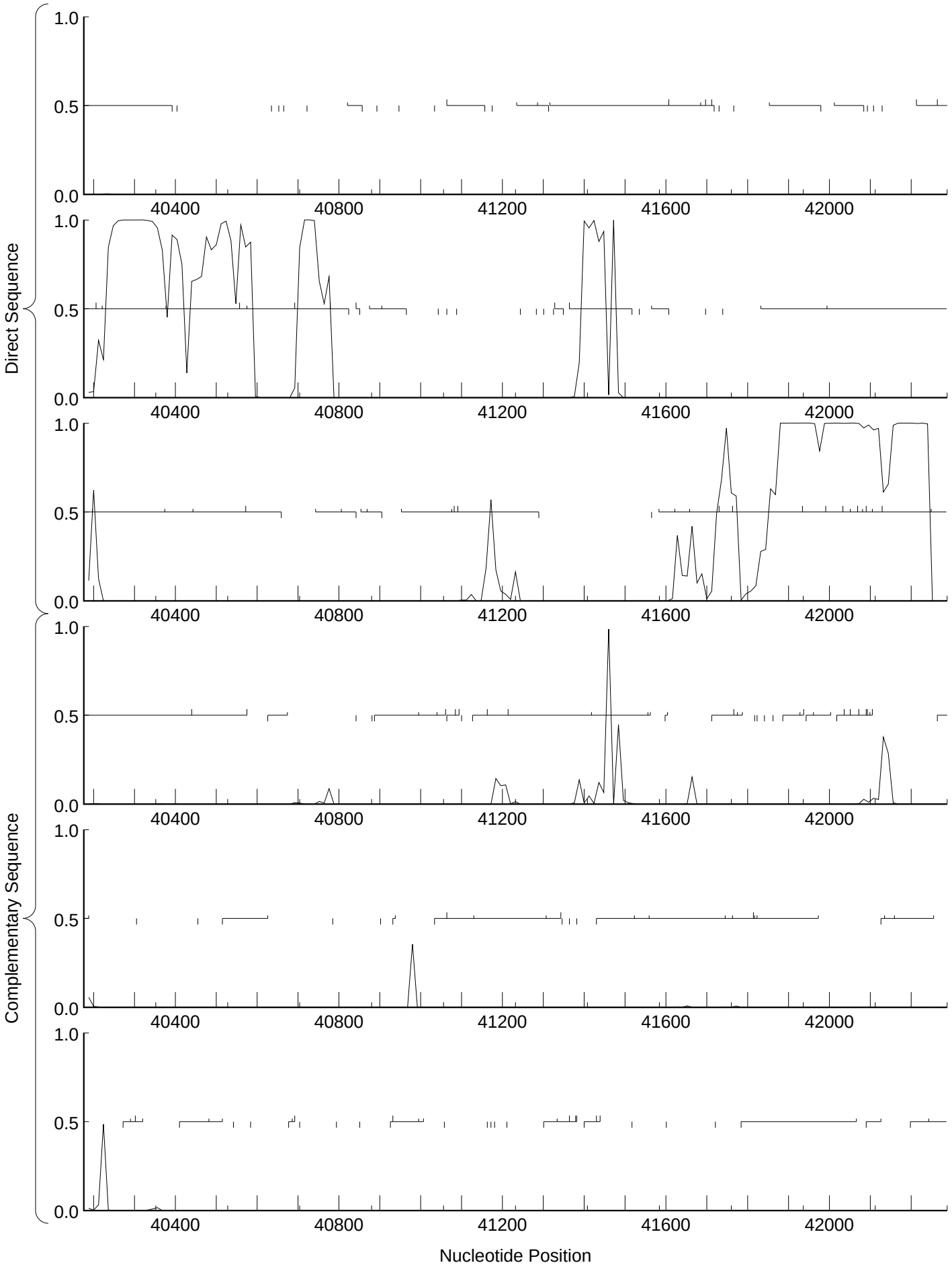


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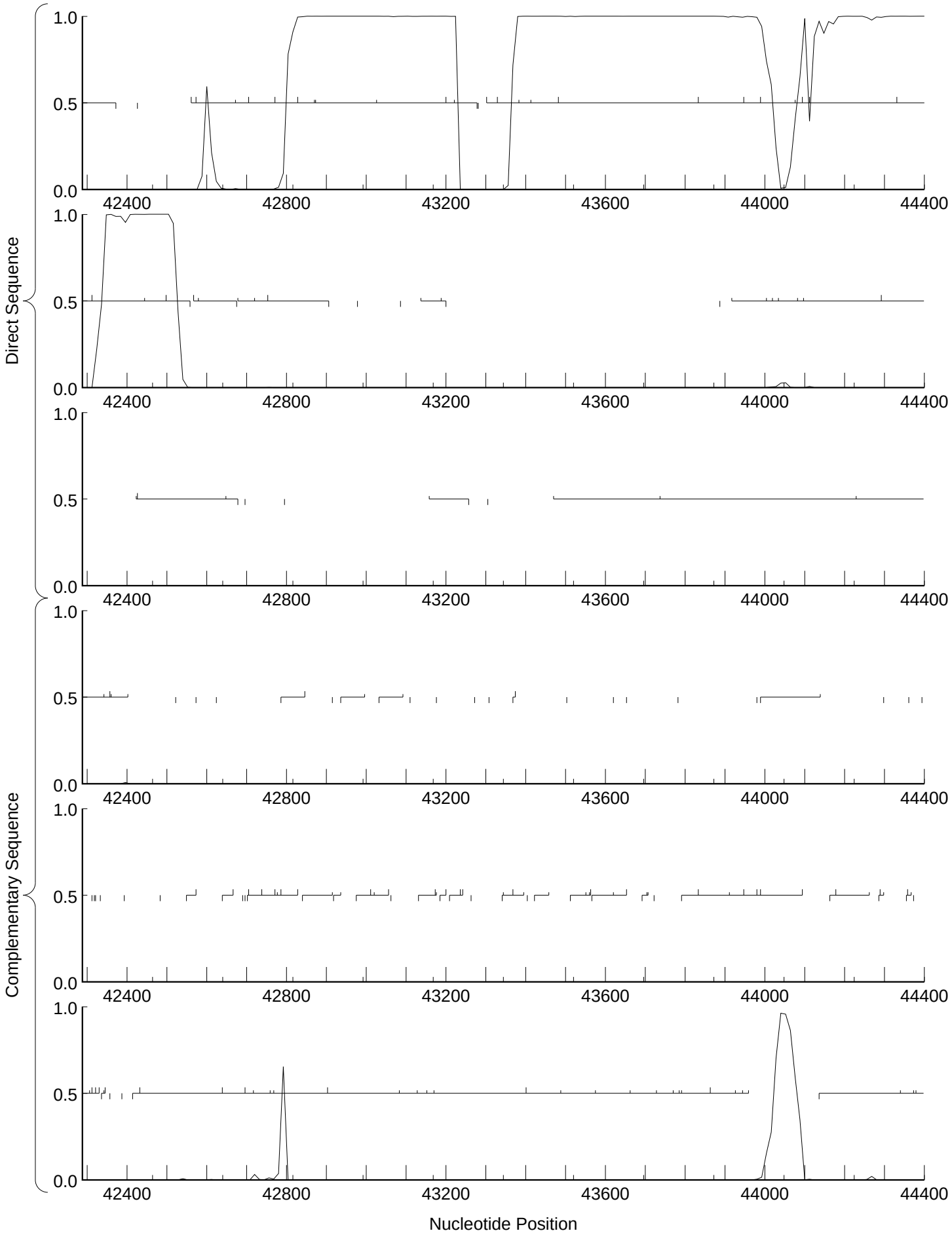


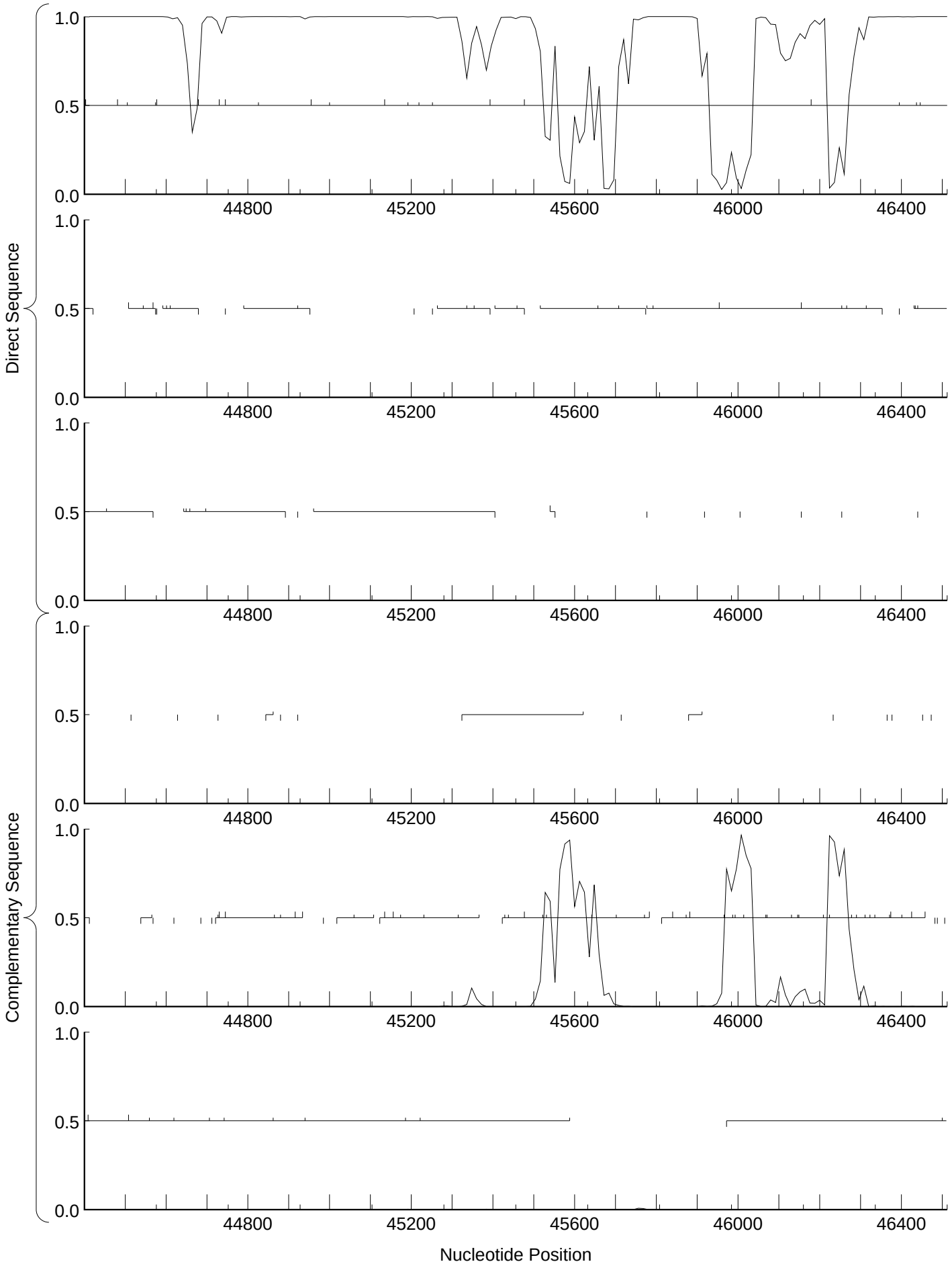

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

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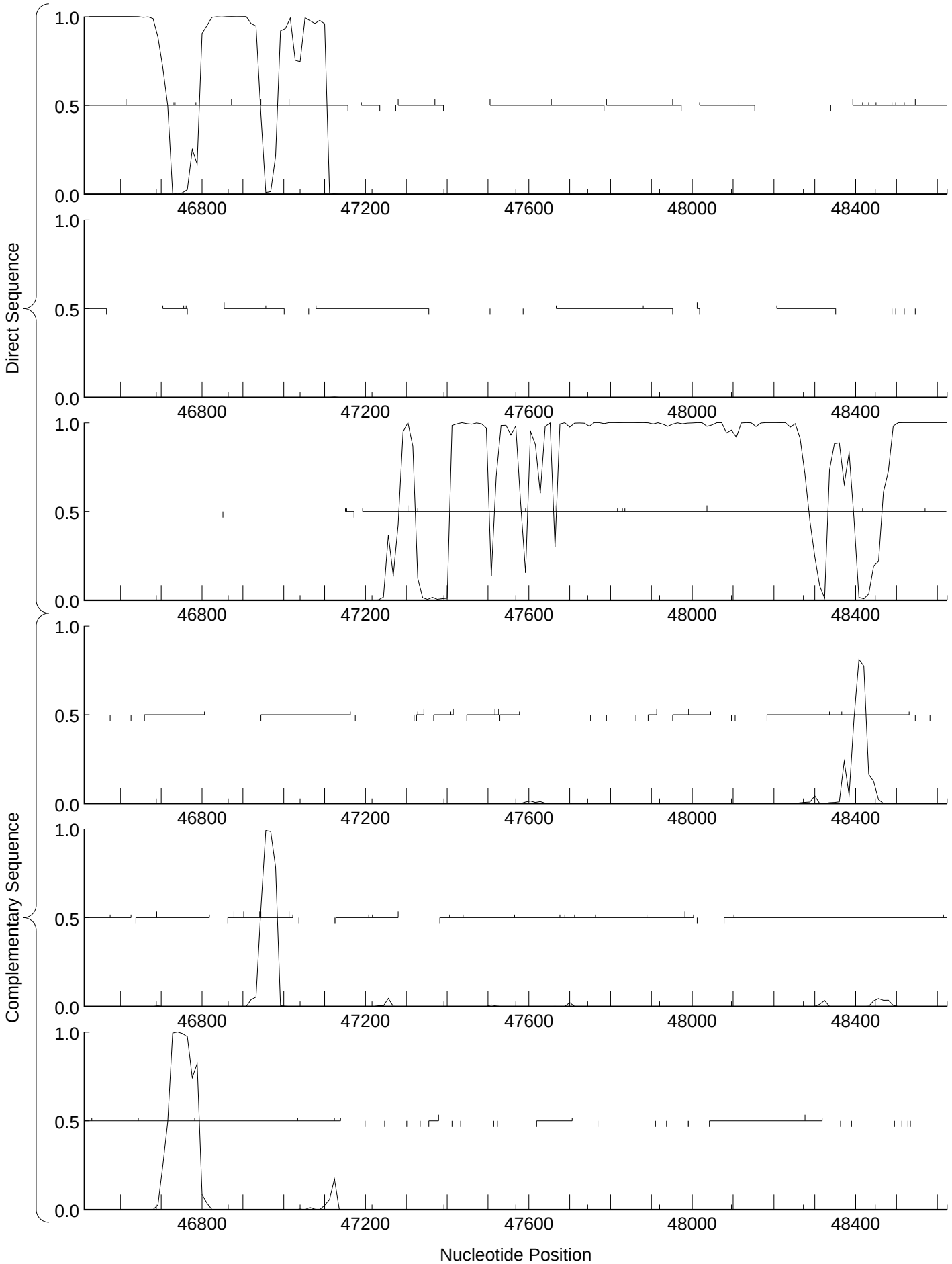


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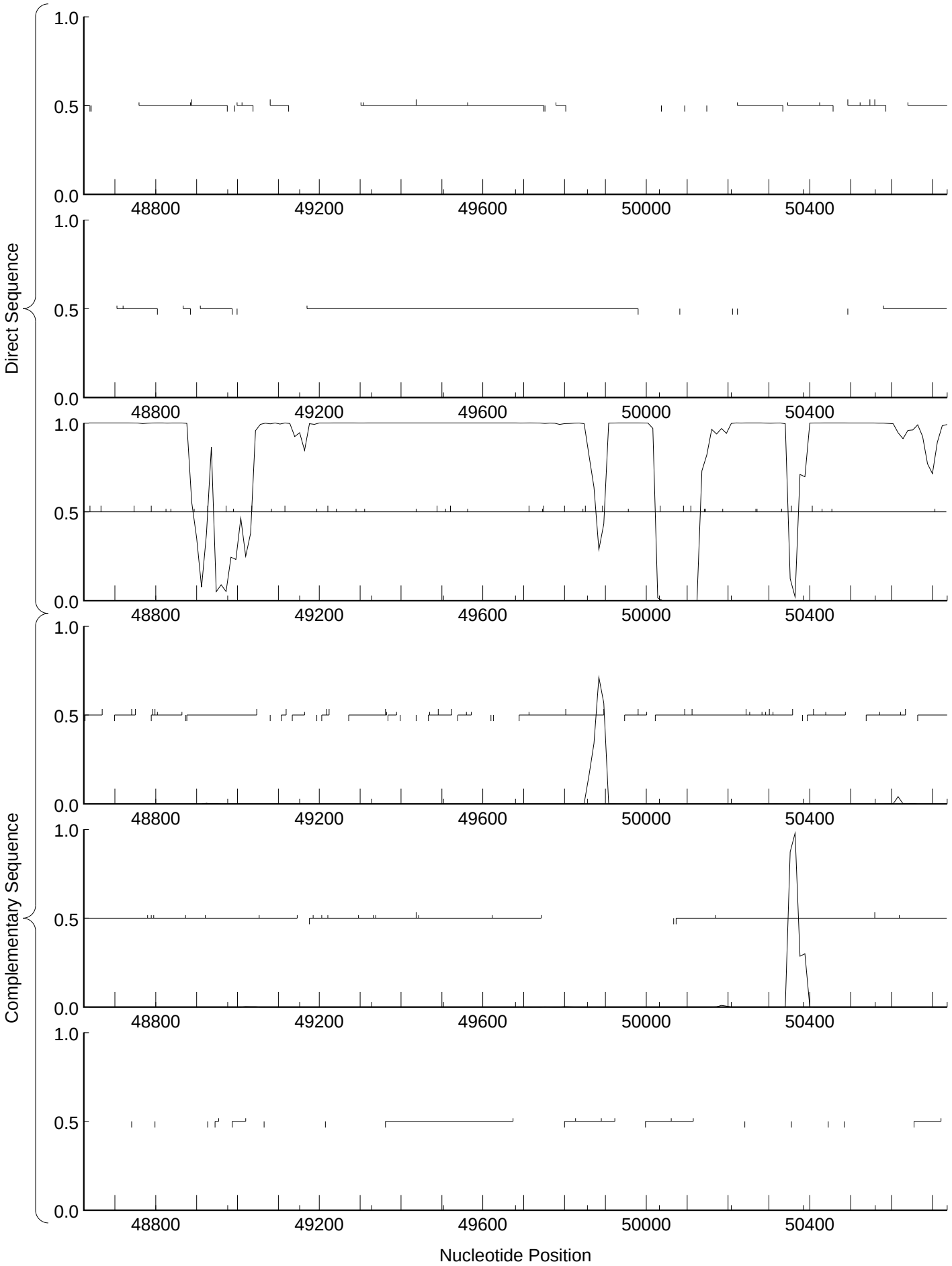



Direct Sequence
Complementary Sequence
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0.0
44800
45200
45600
46000
46400
Nucleotide Position

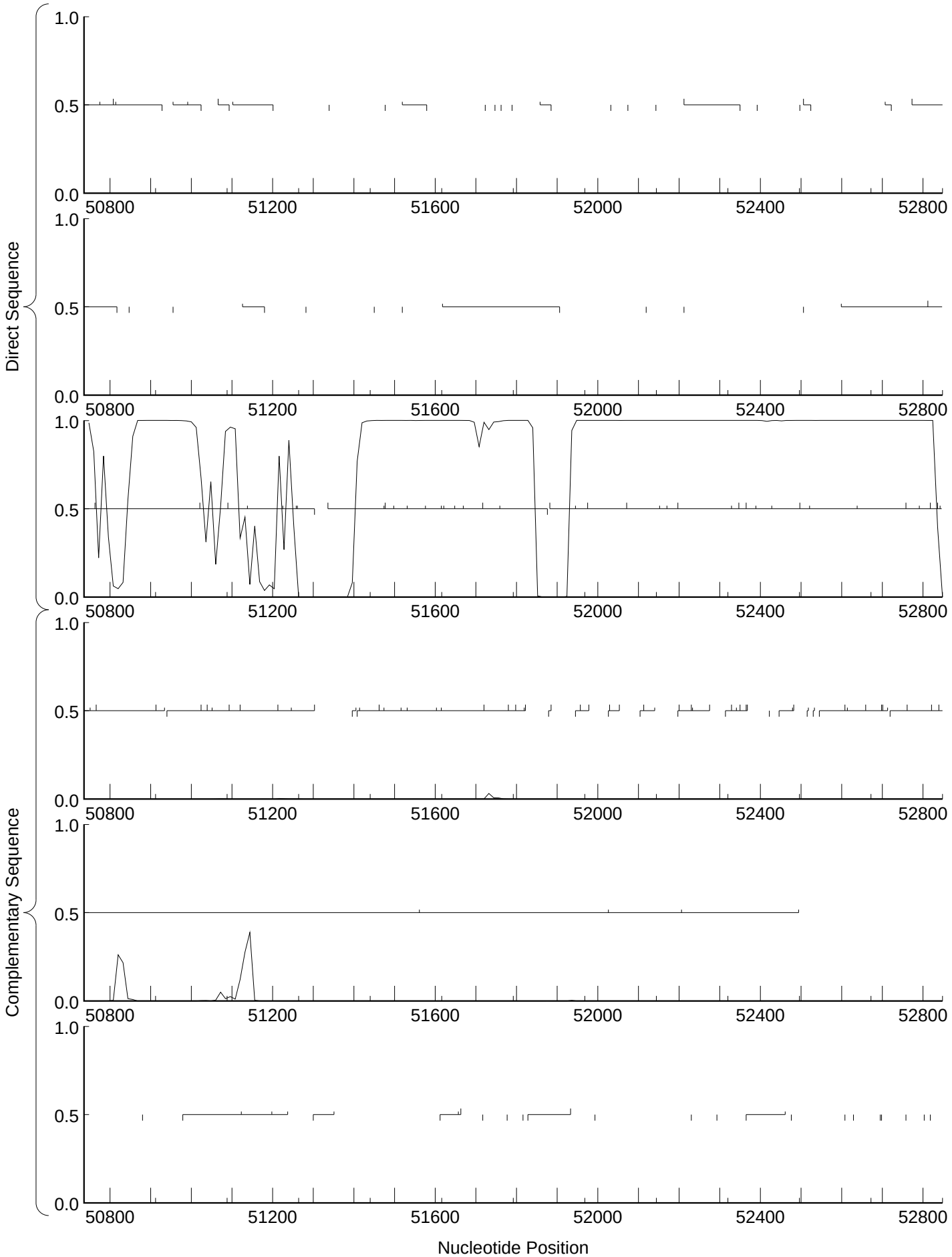
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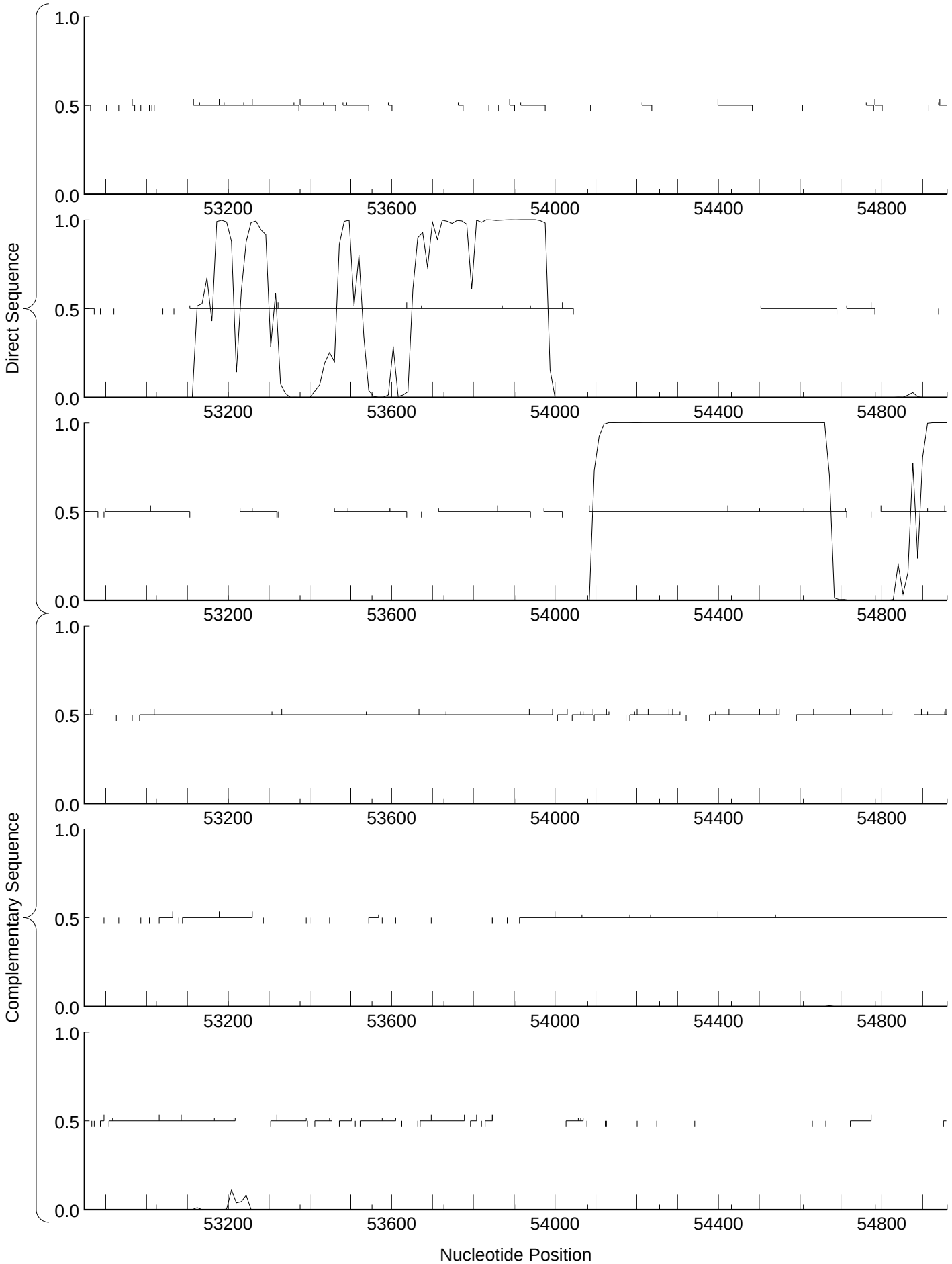
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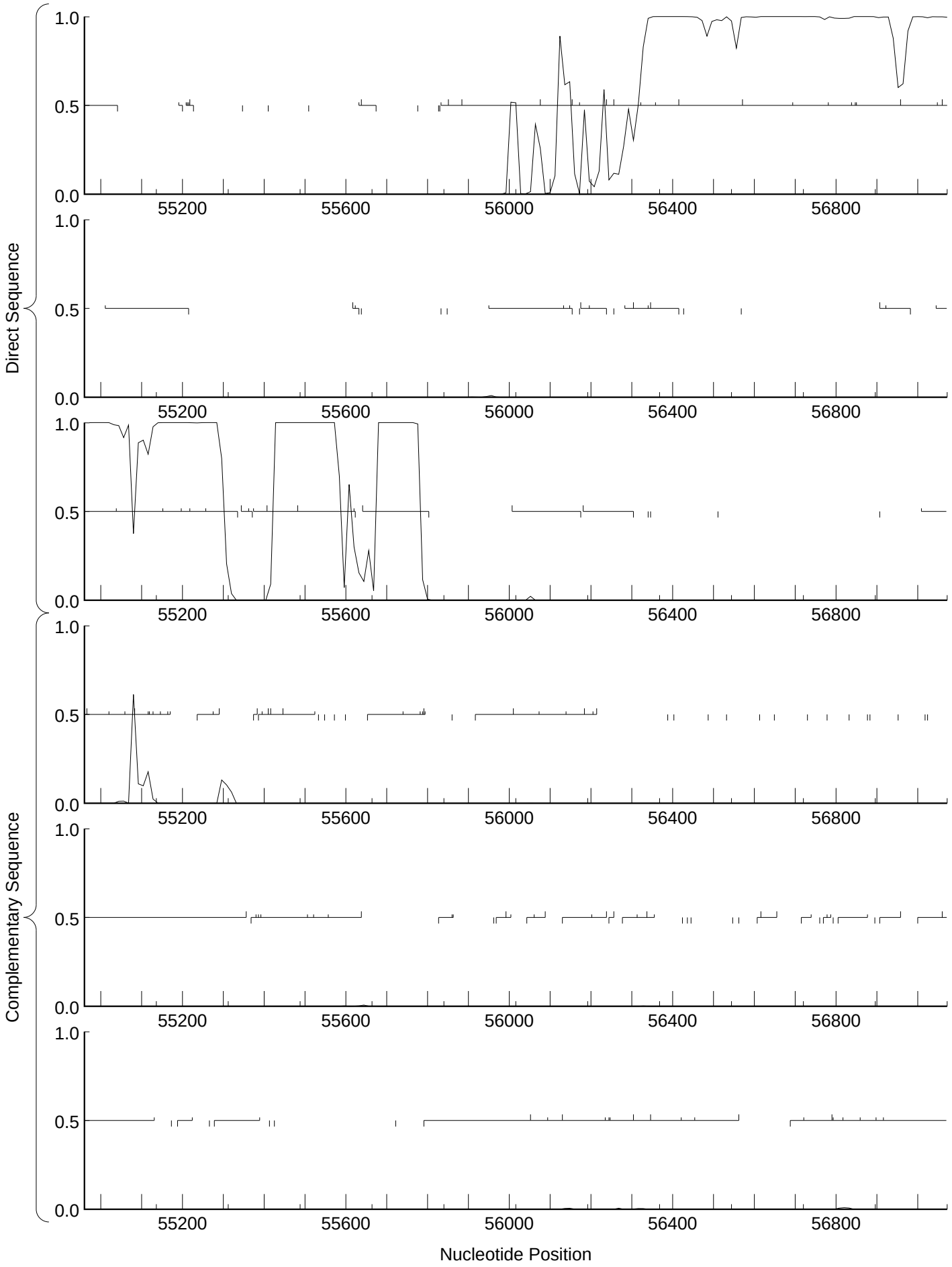


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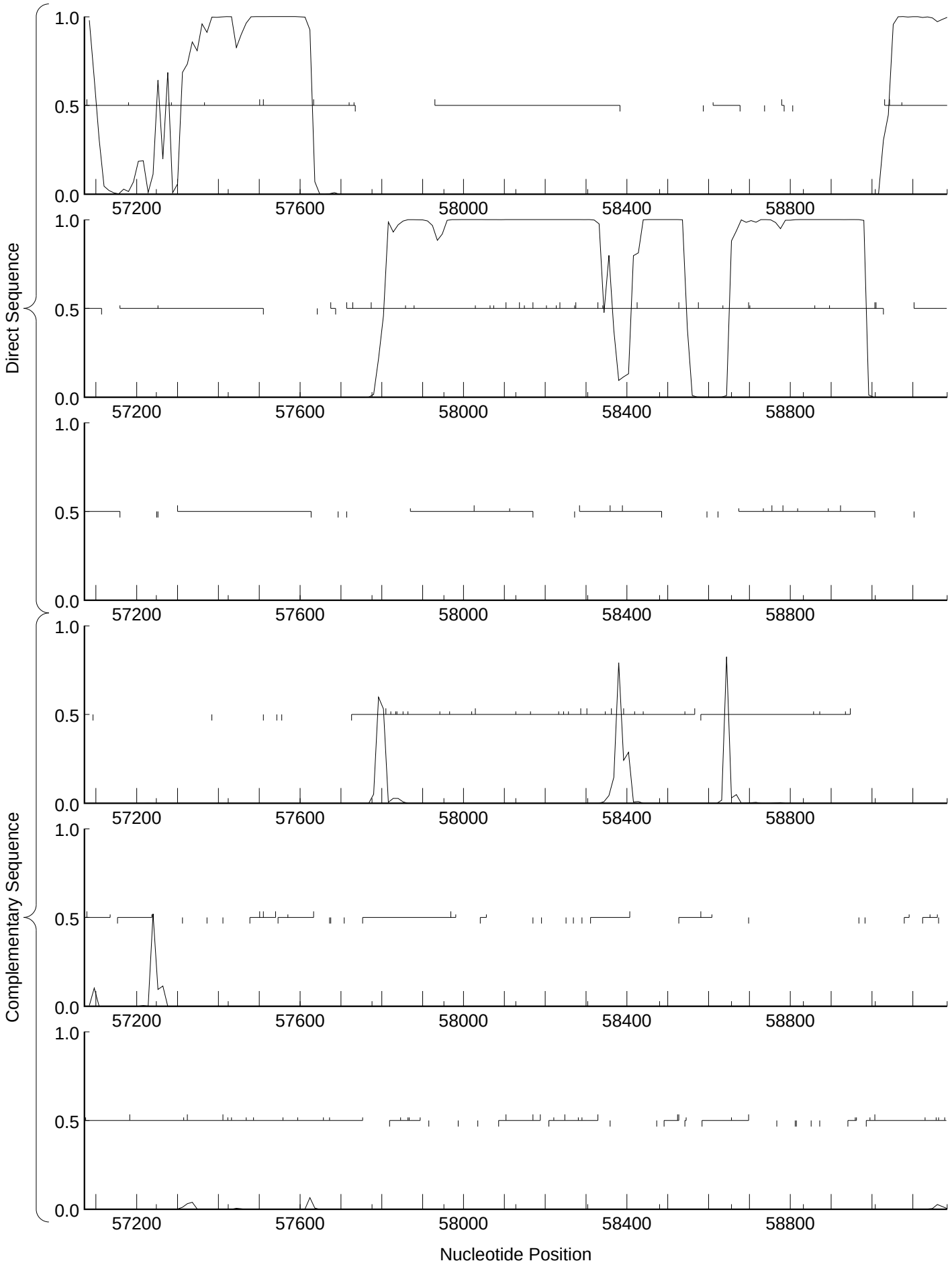


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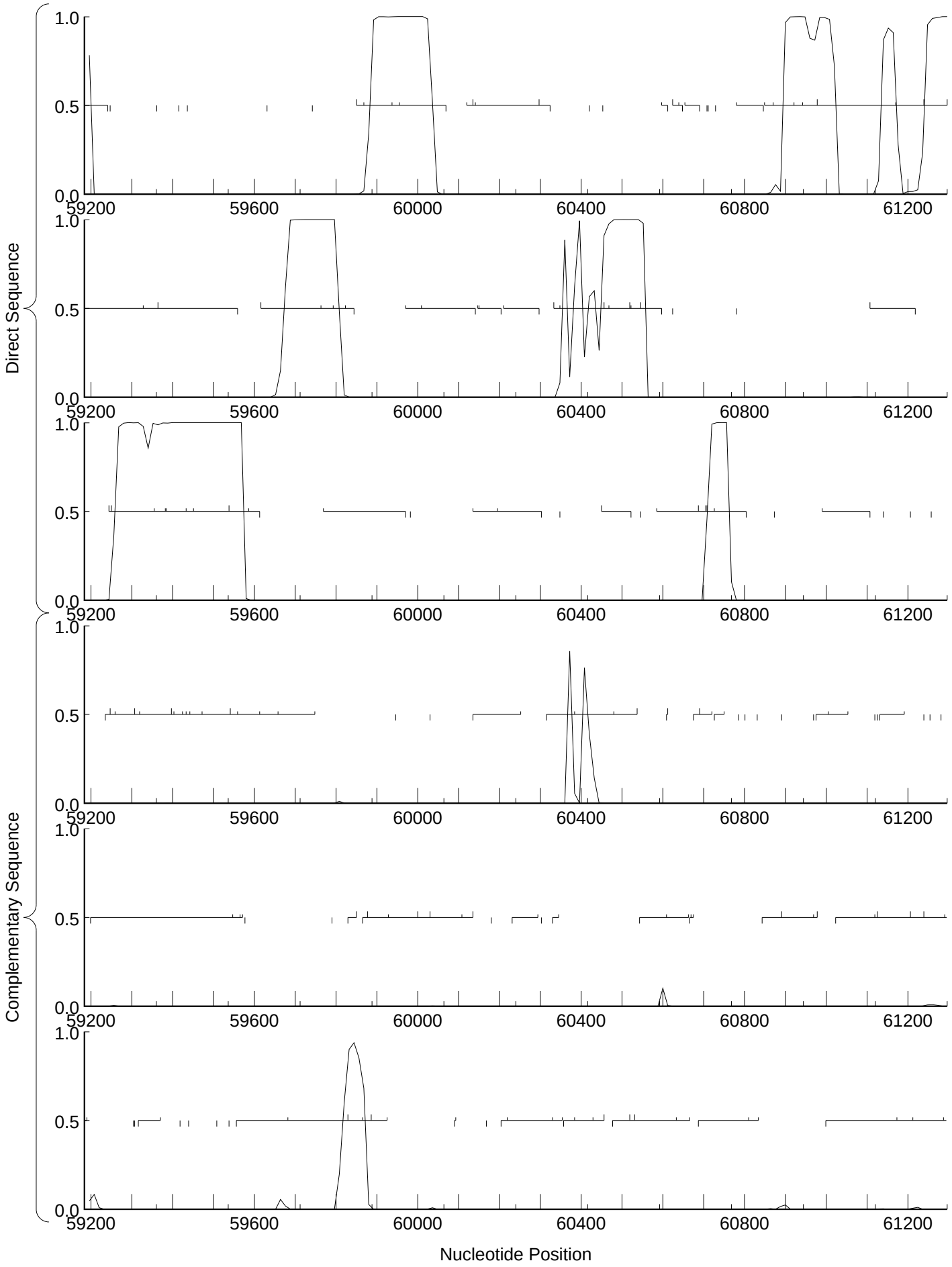



Direct Sequence
Complementary Sequence
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0.0
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55600
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56400
56800
Nucleotide Position

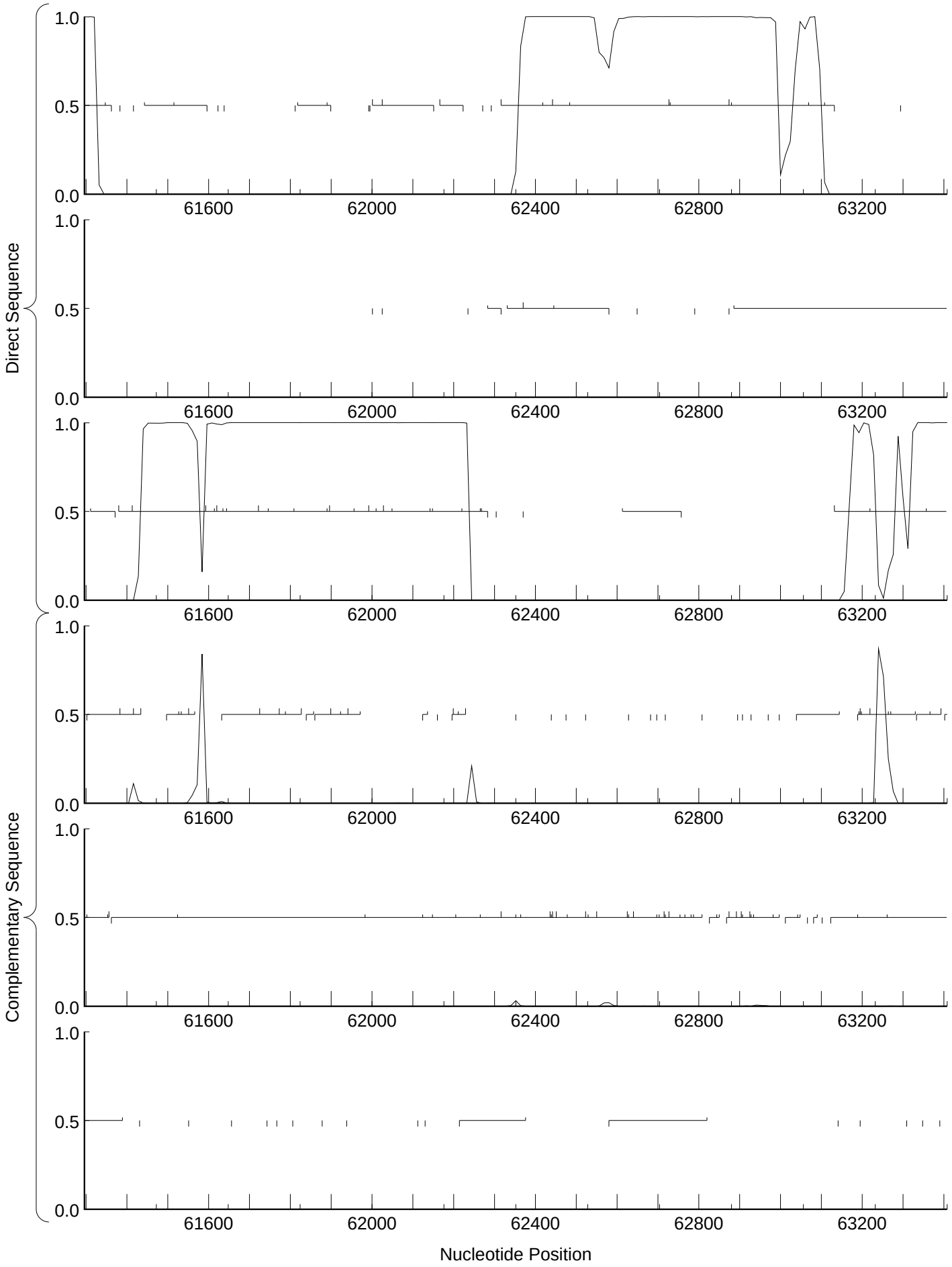
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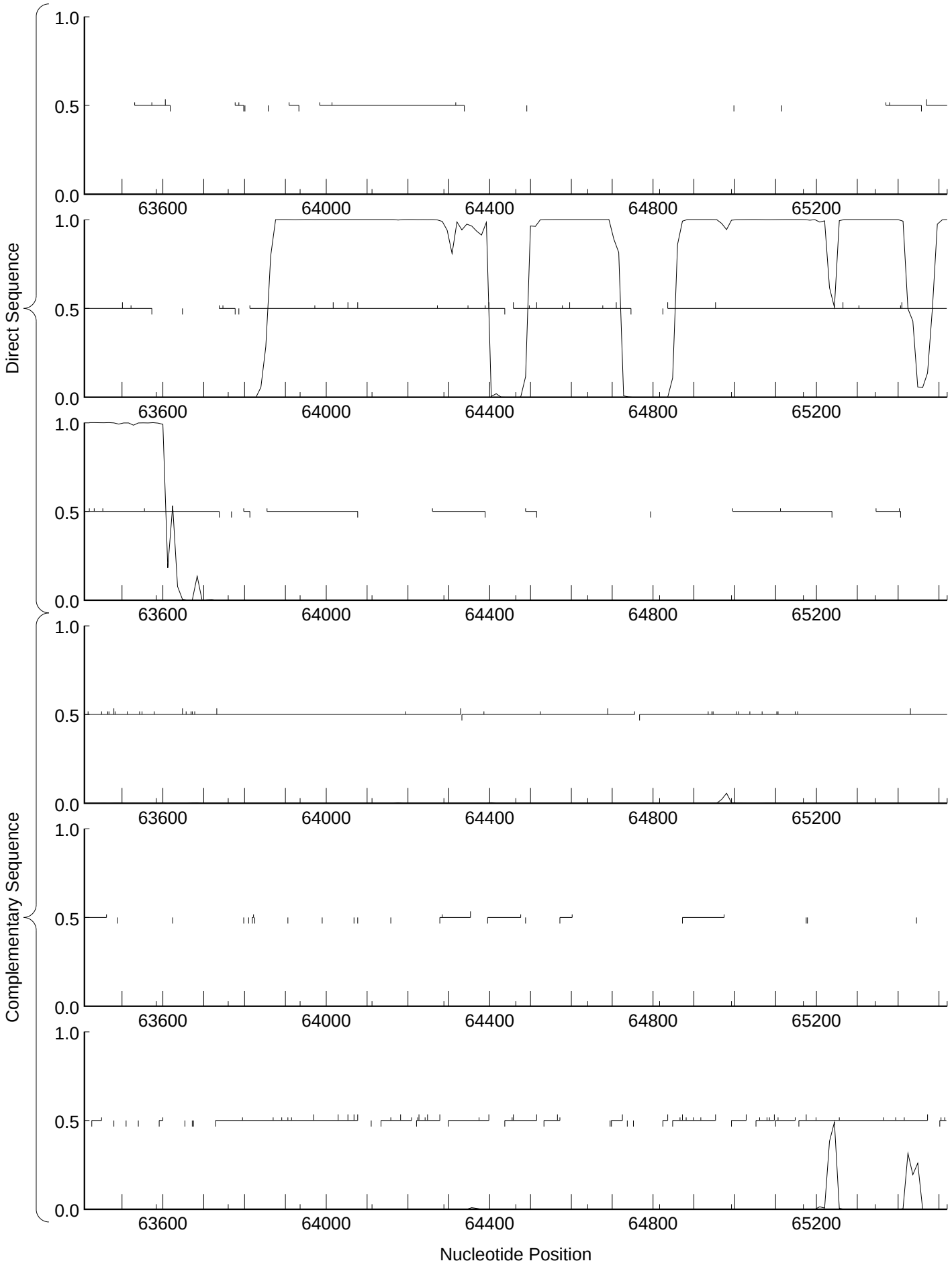
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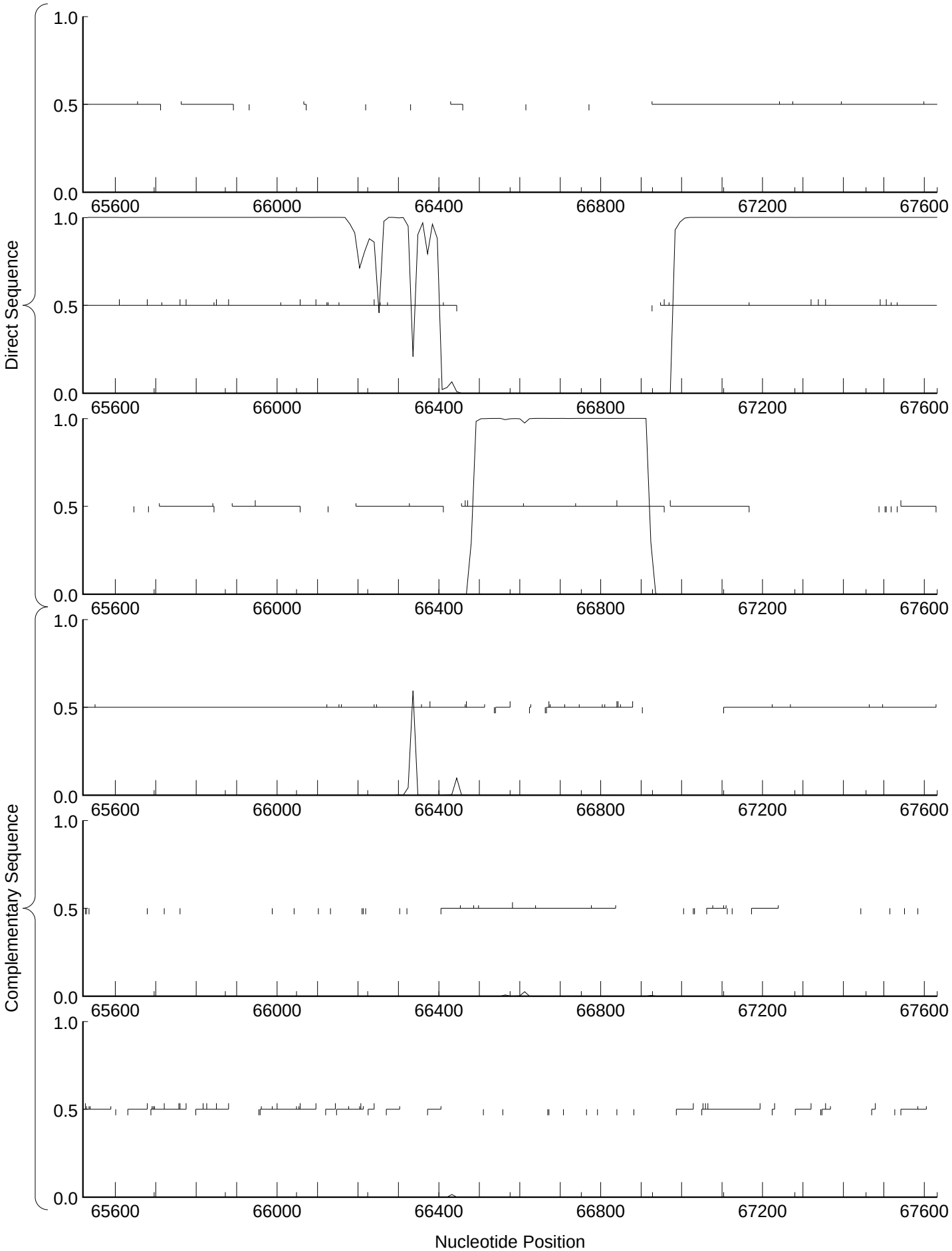
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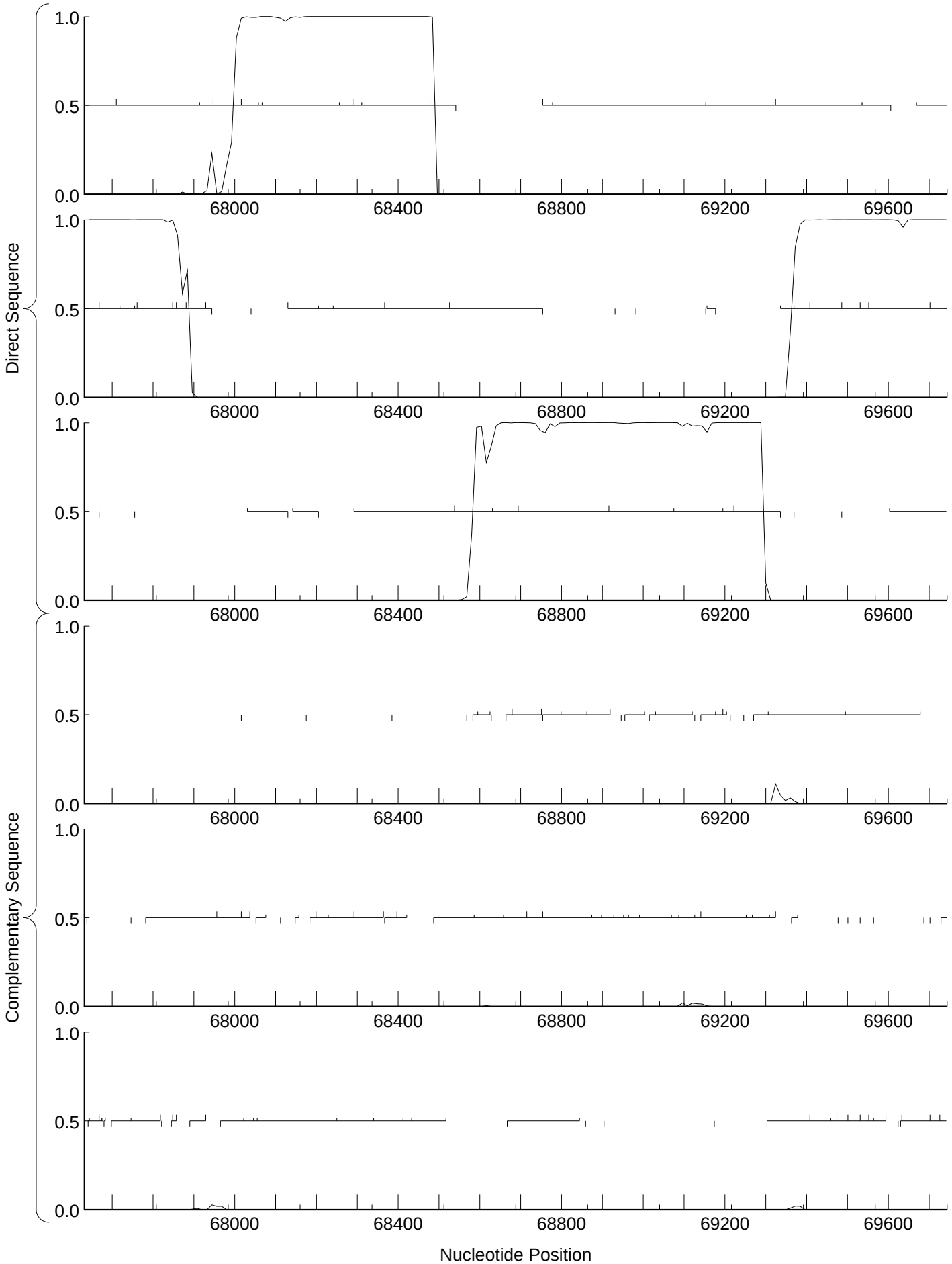
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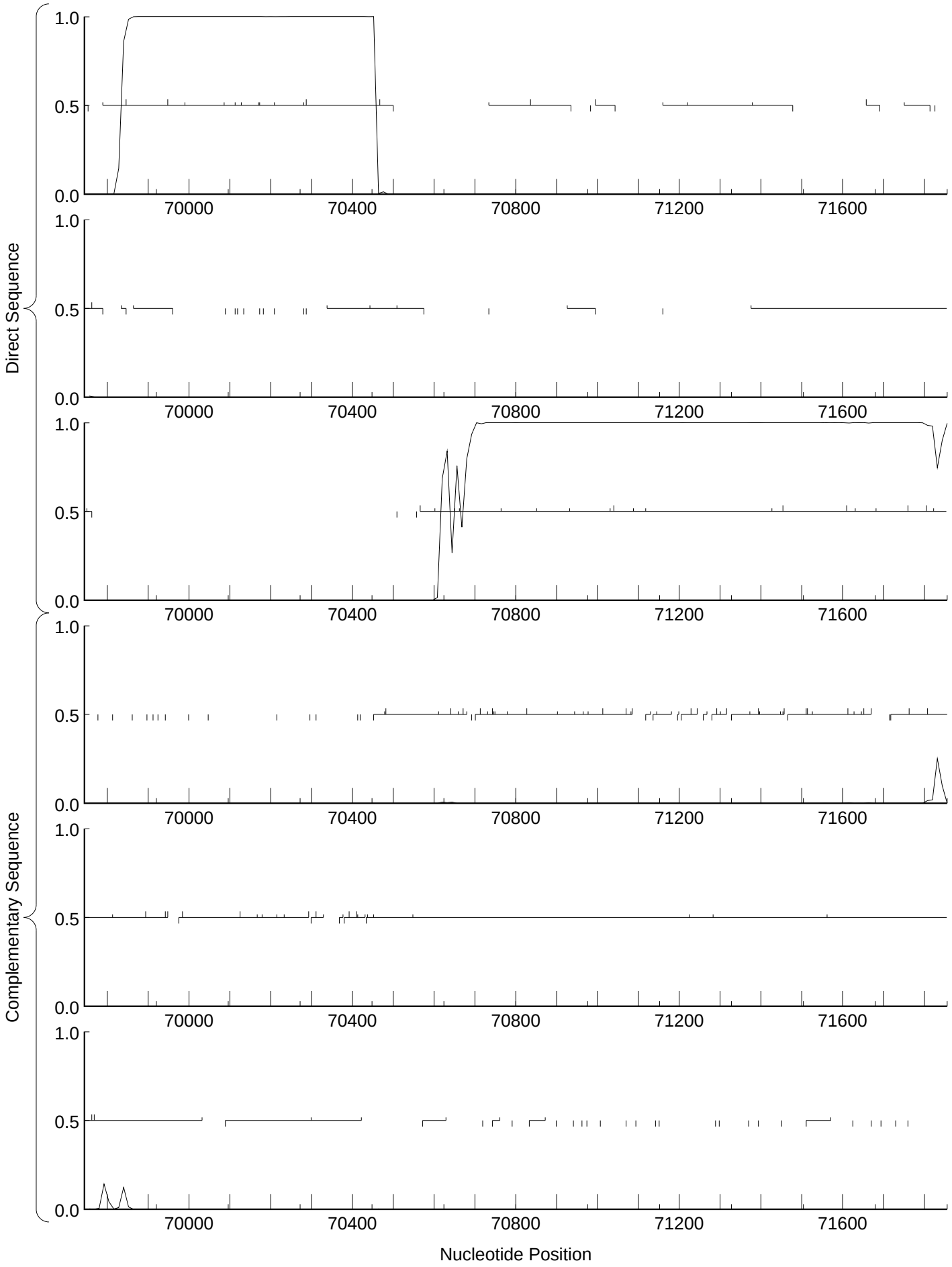


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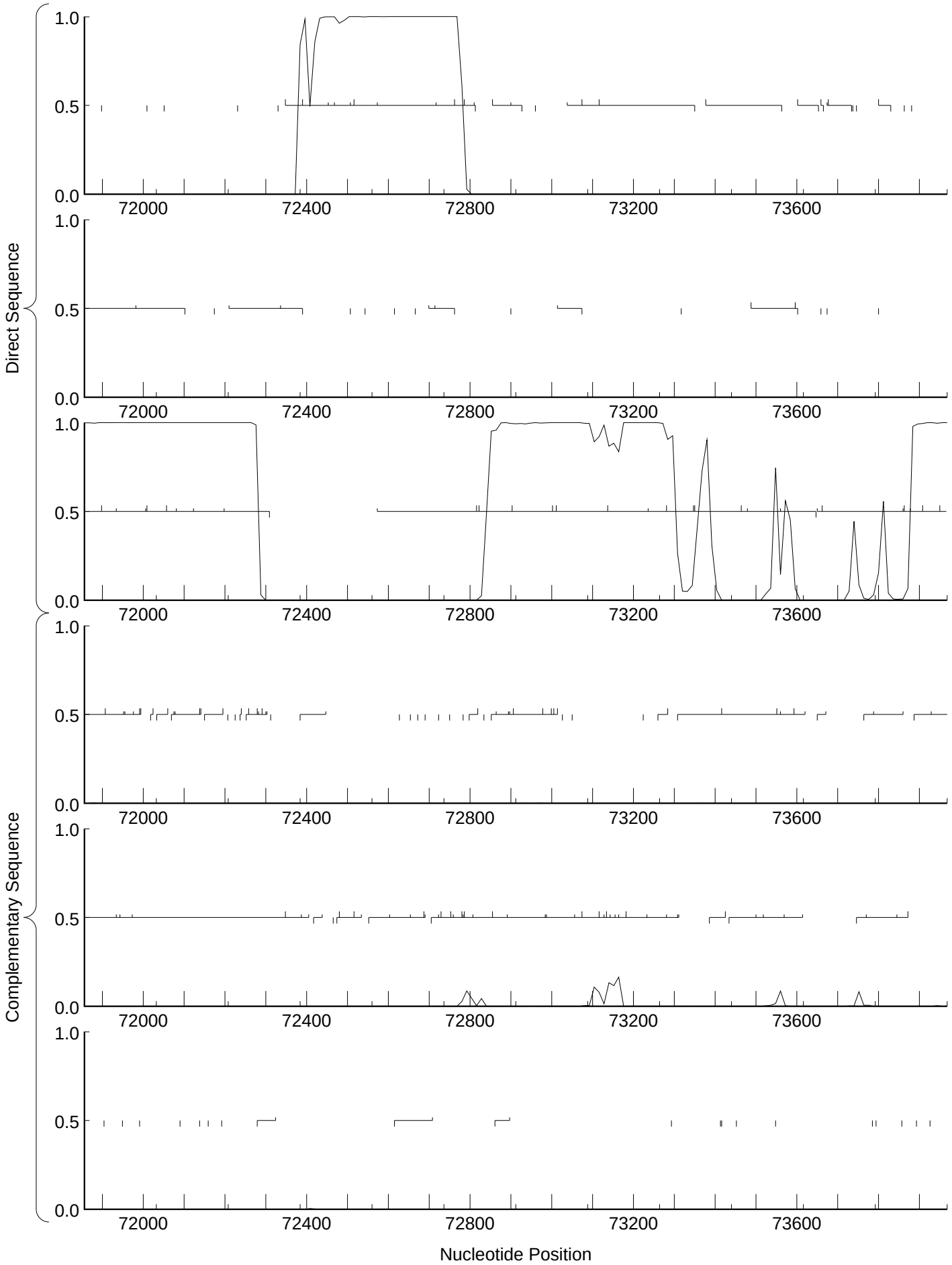


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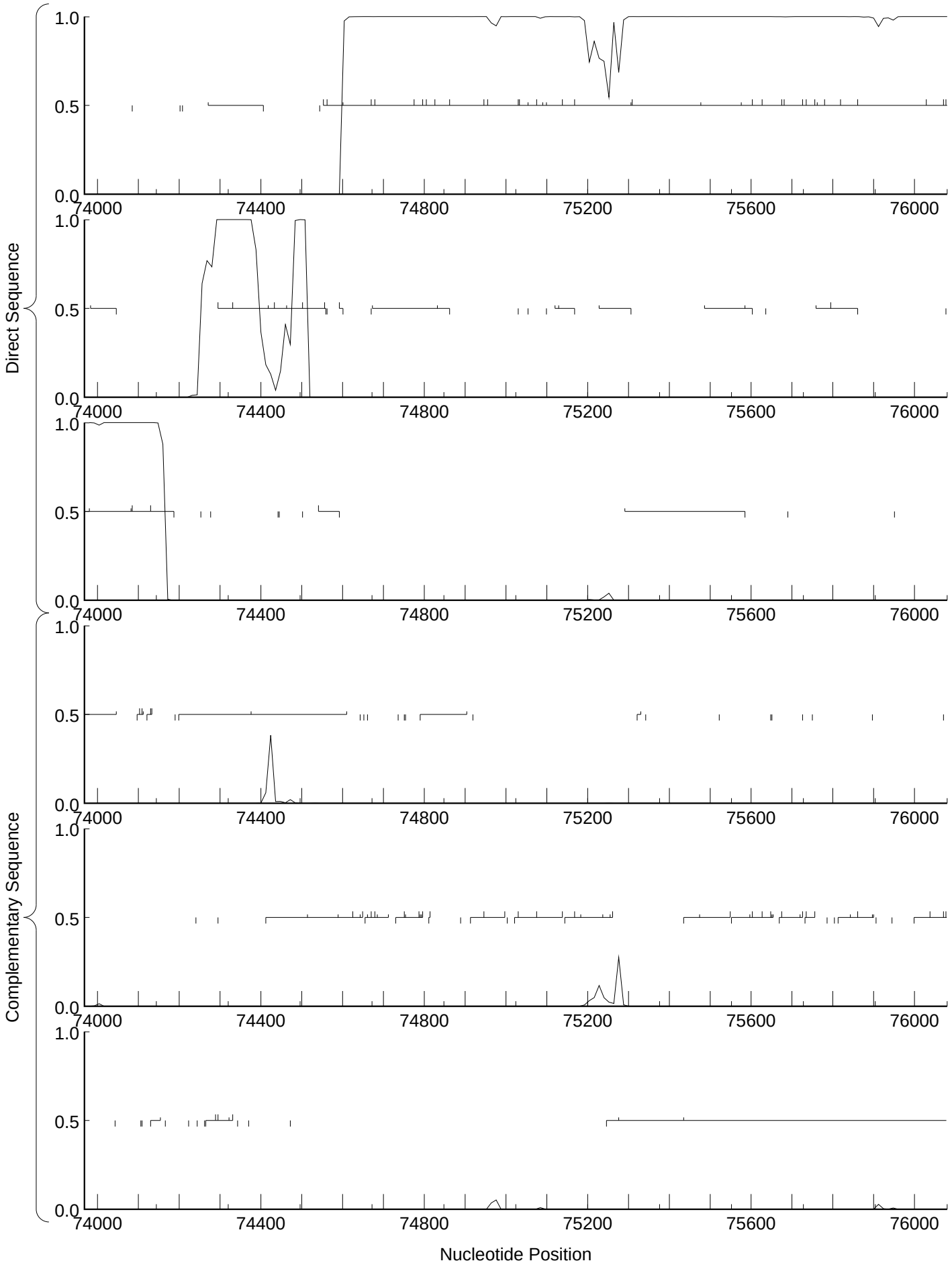



Direct Sequence
Complementary Sequence
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0.5
0.0
70000
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70800
71200
71600
Nucleotide Position

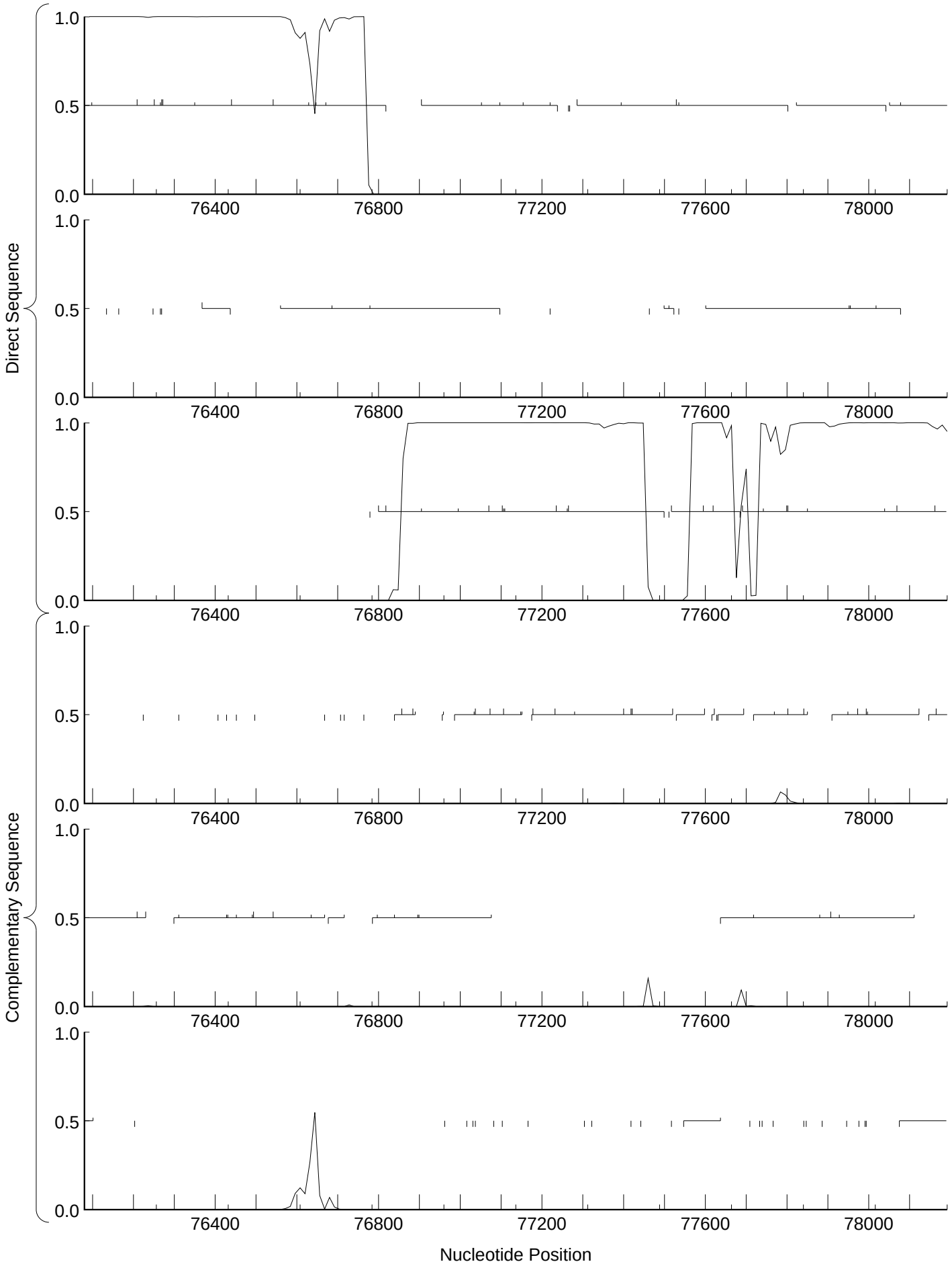
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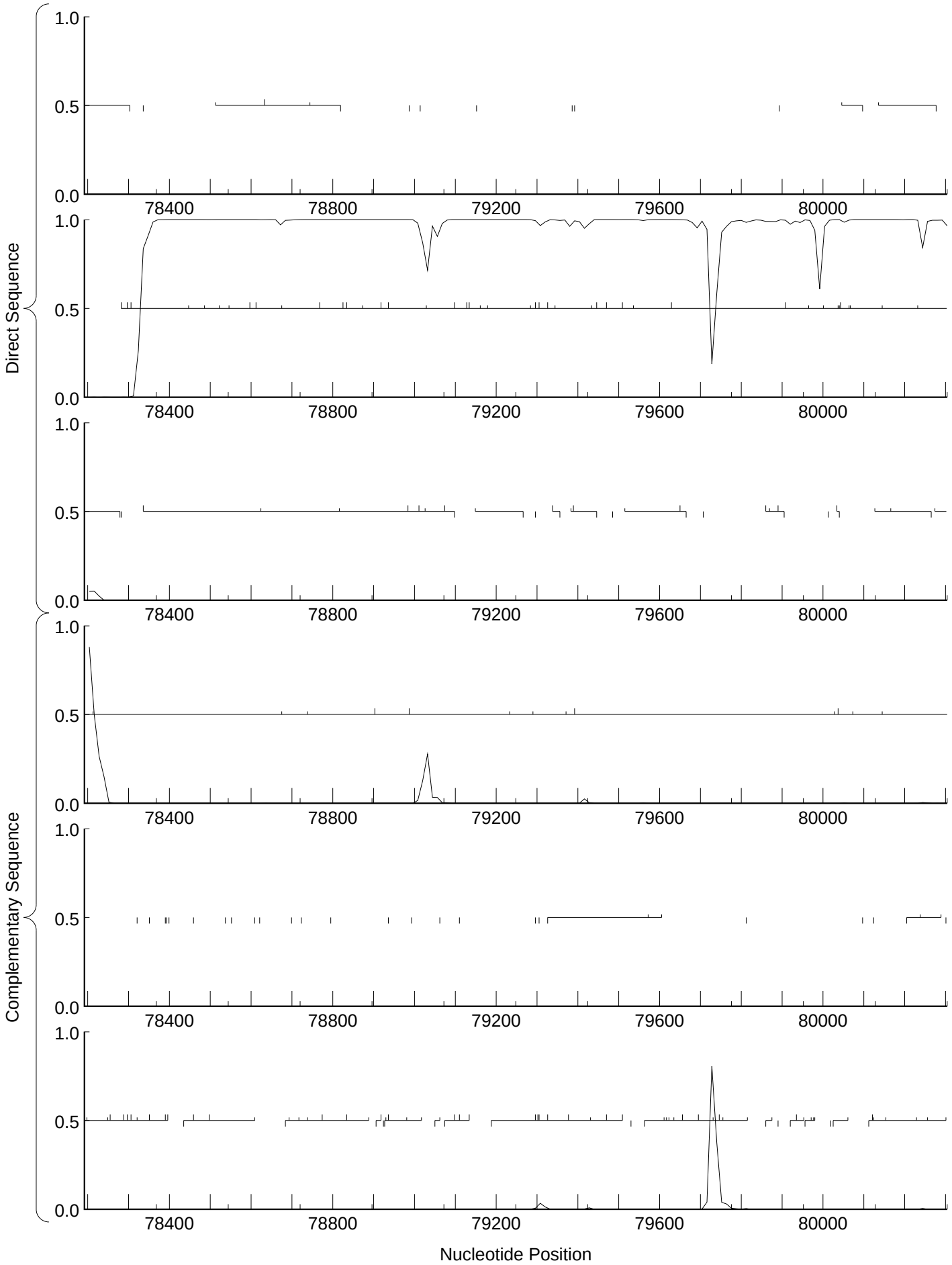
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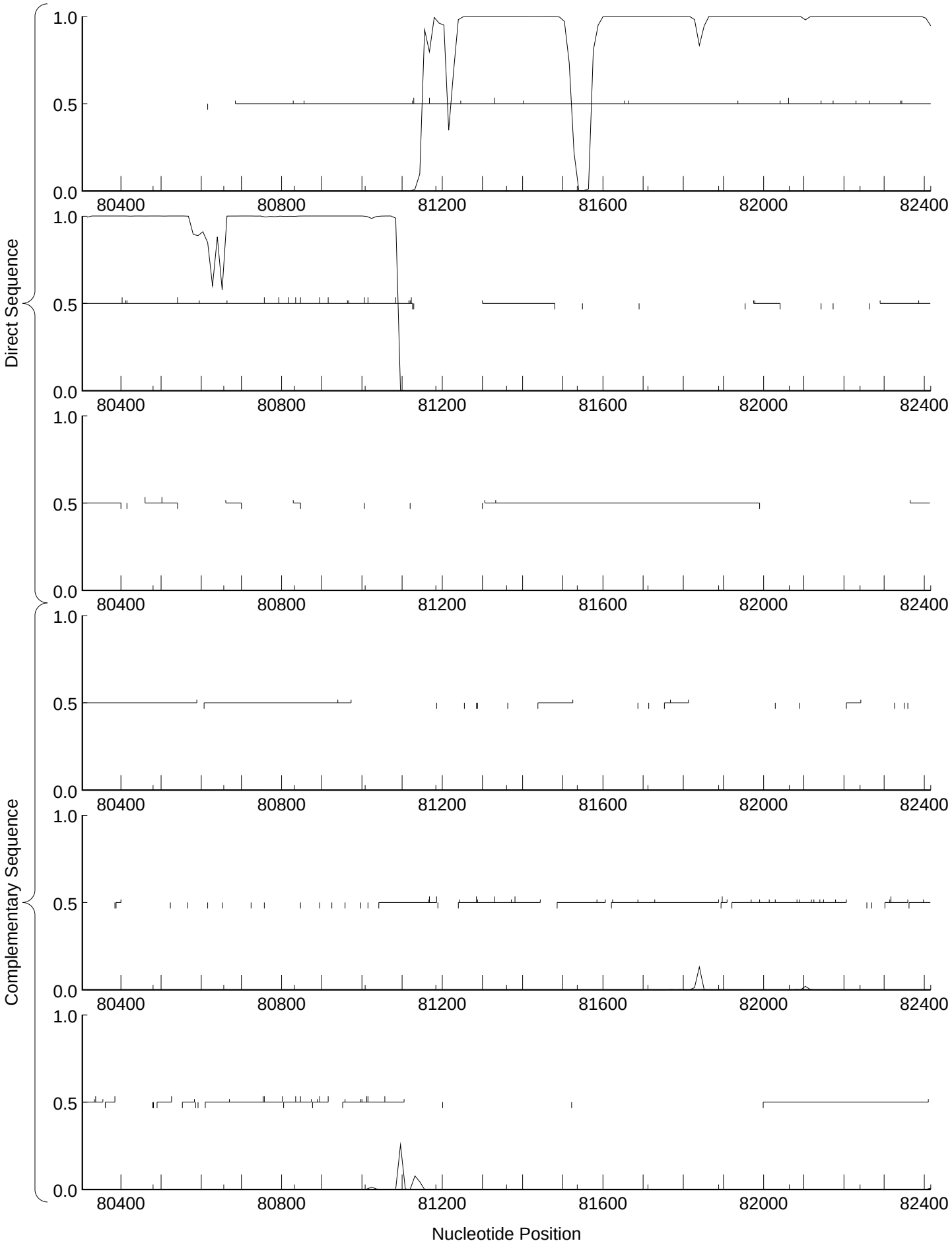
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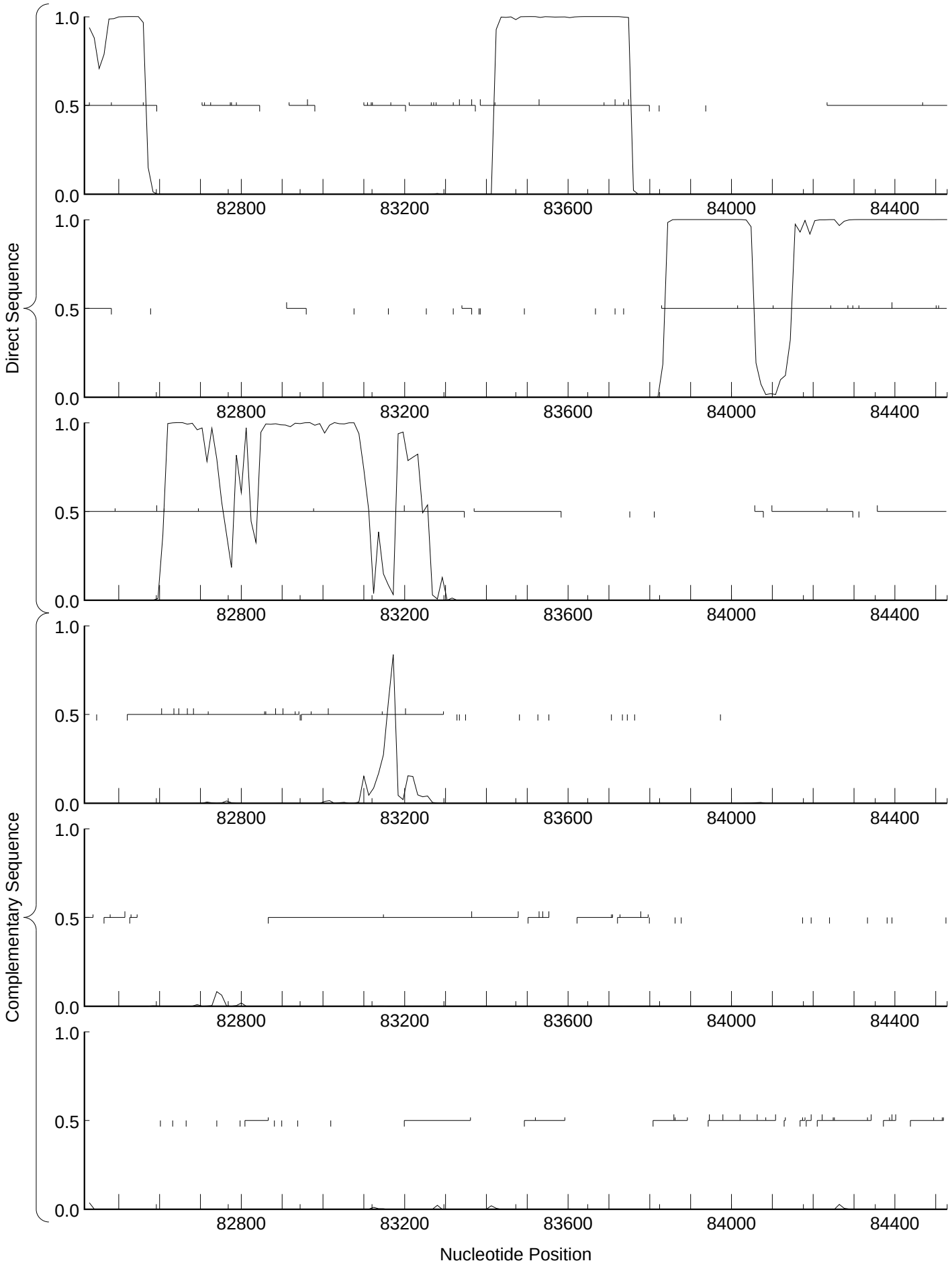
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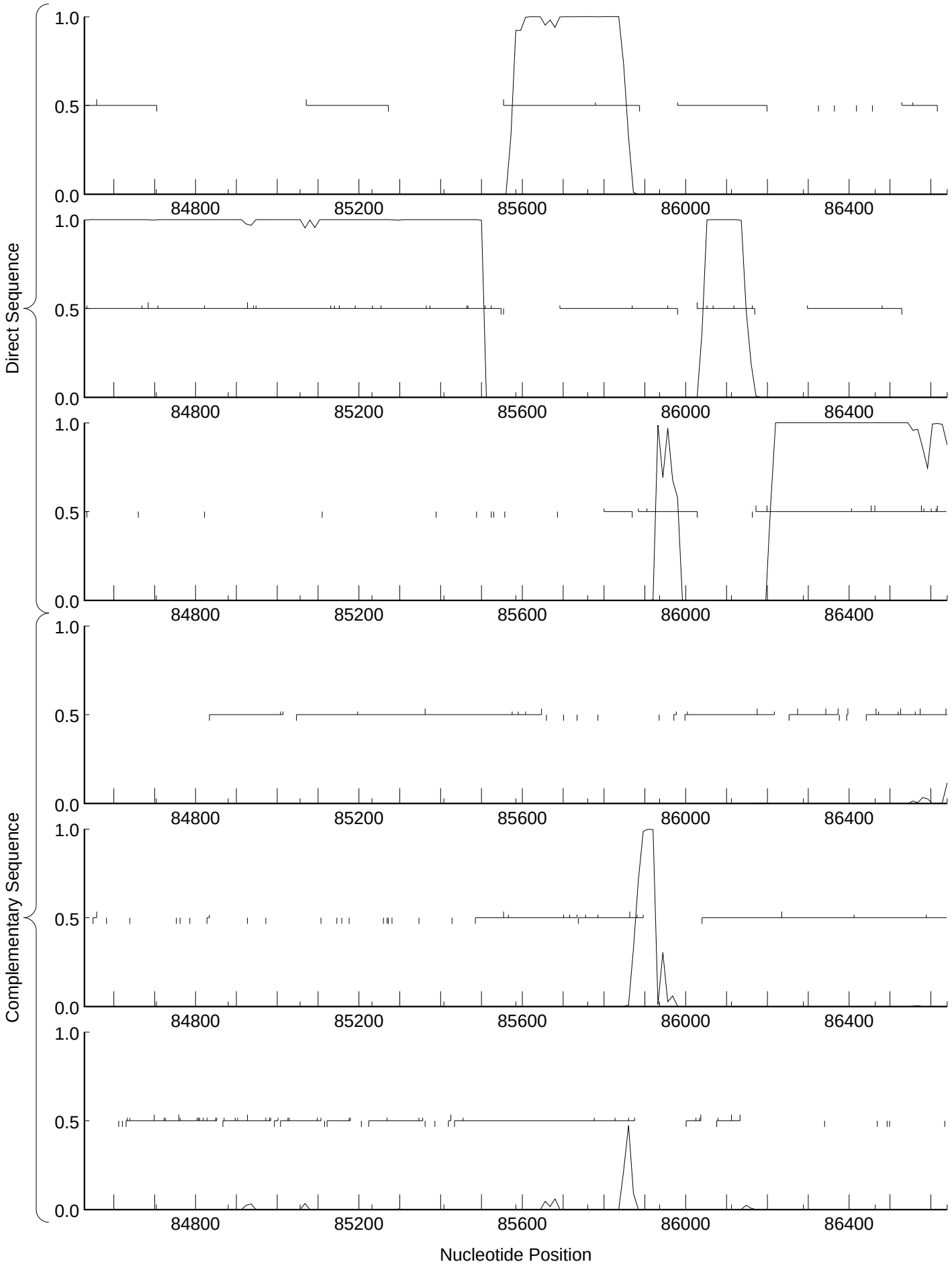
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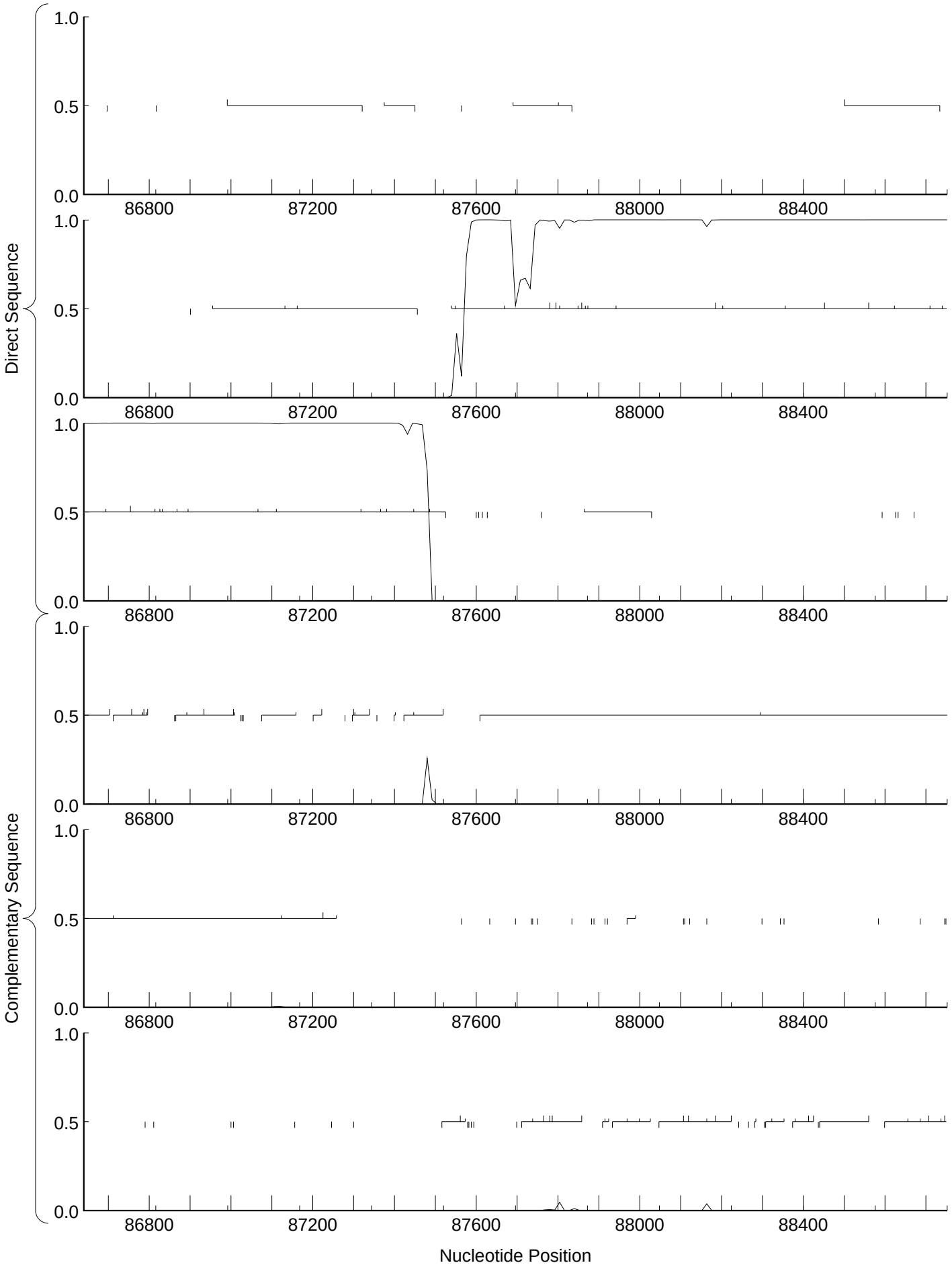
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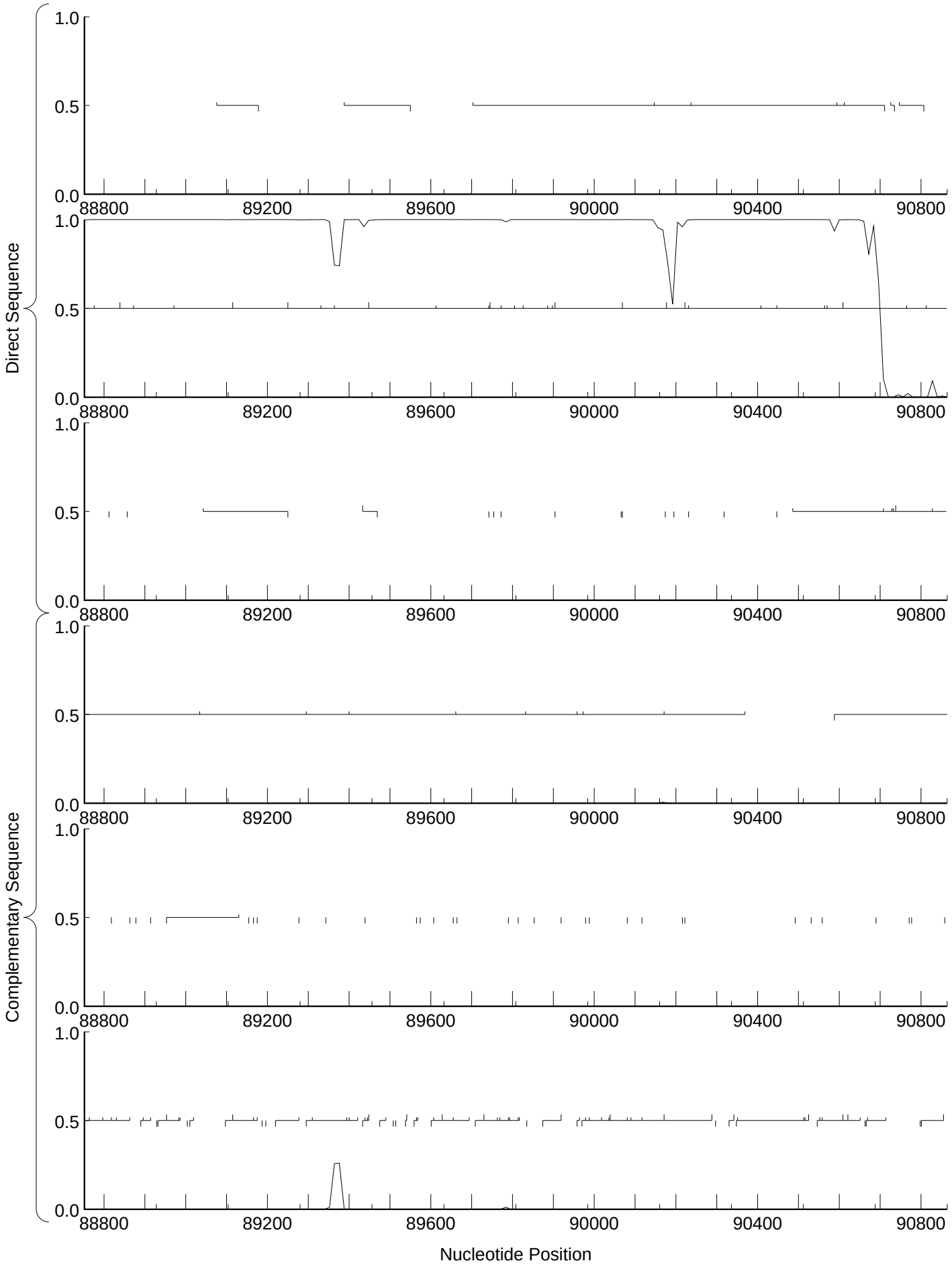
erium phage NuevoMundo Complete Sequence, 155943 bp, Circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 42/75



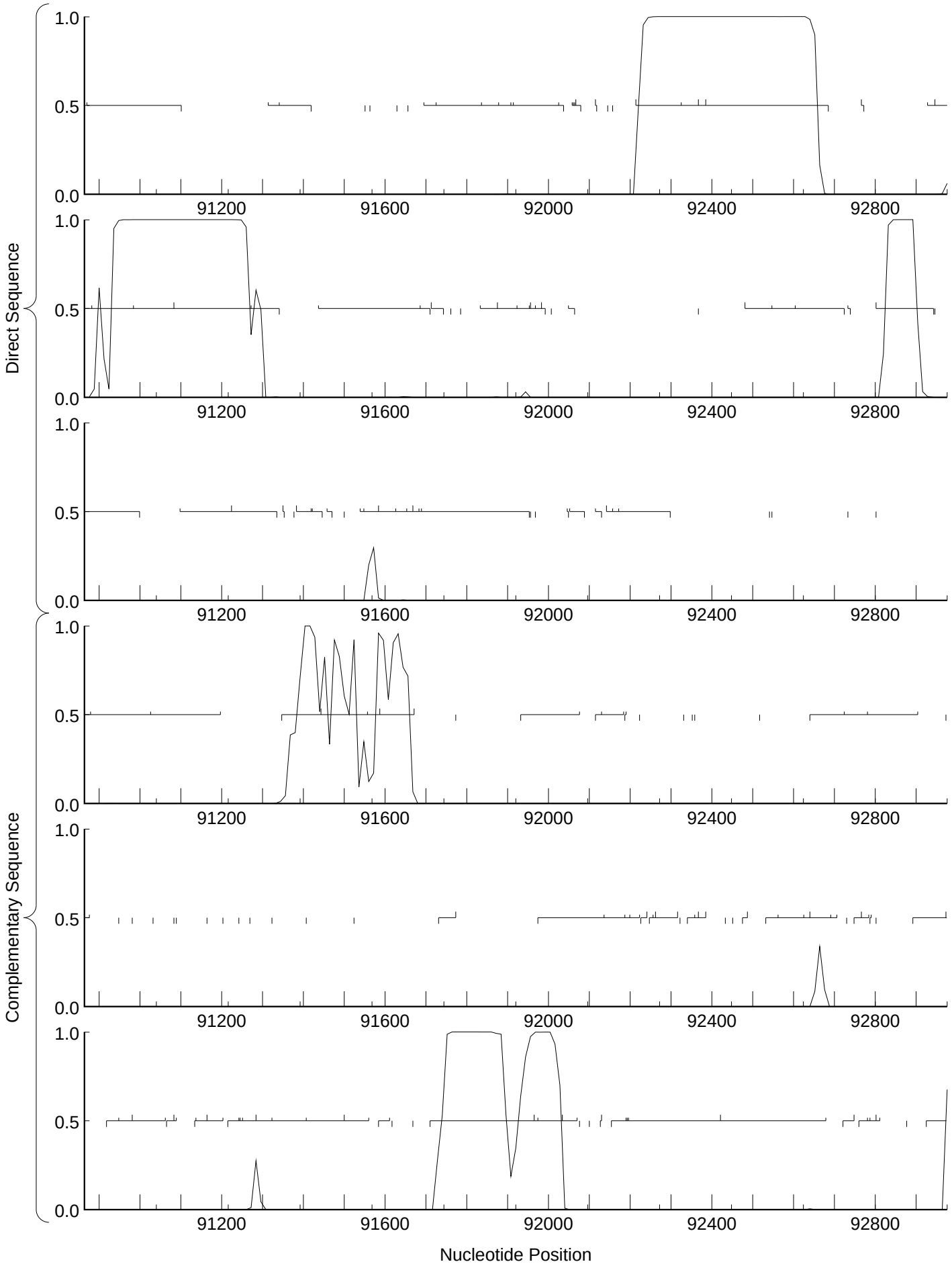
erium phage NuevoMundo Complete Sequence, 155943 bp, Circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 43/75



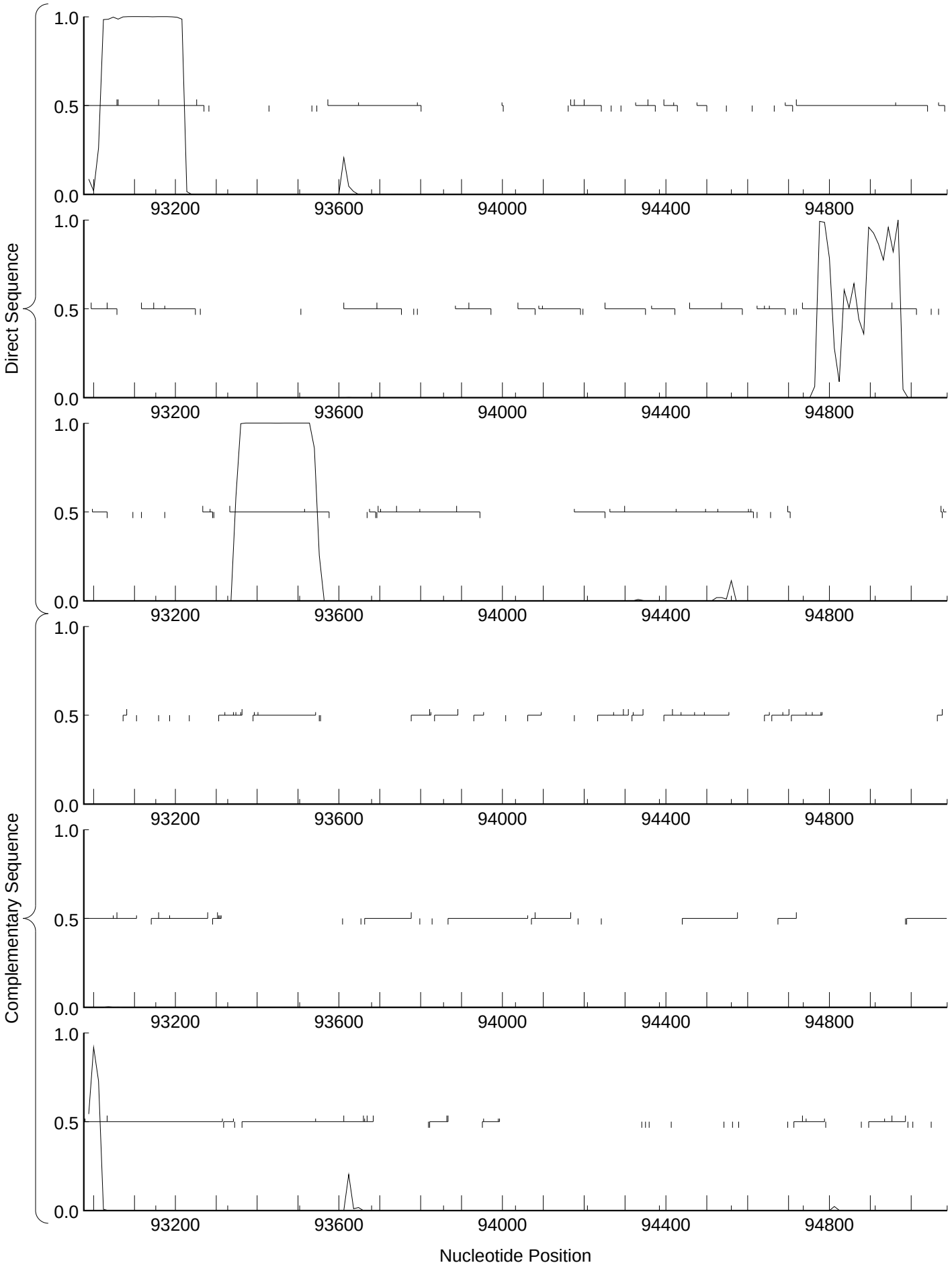
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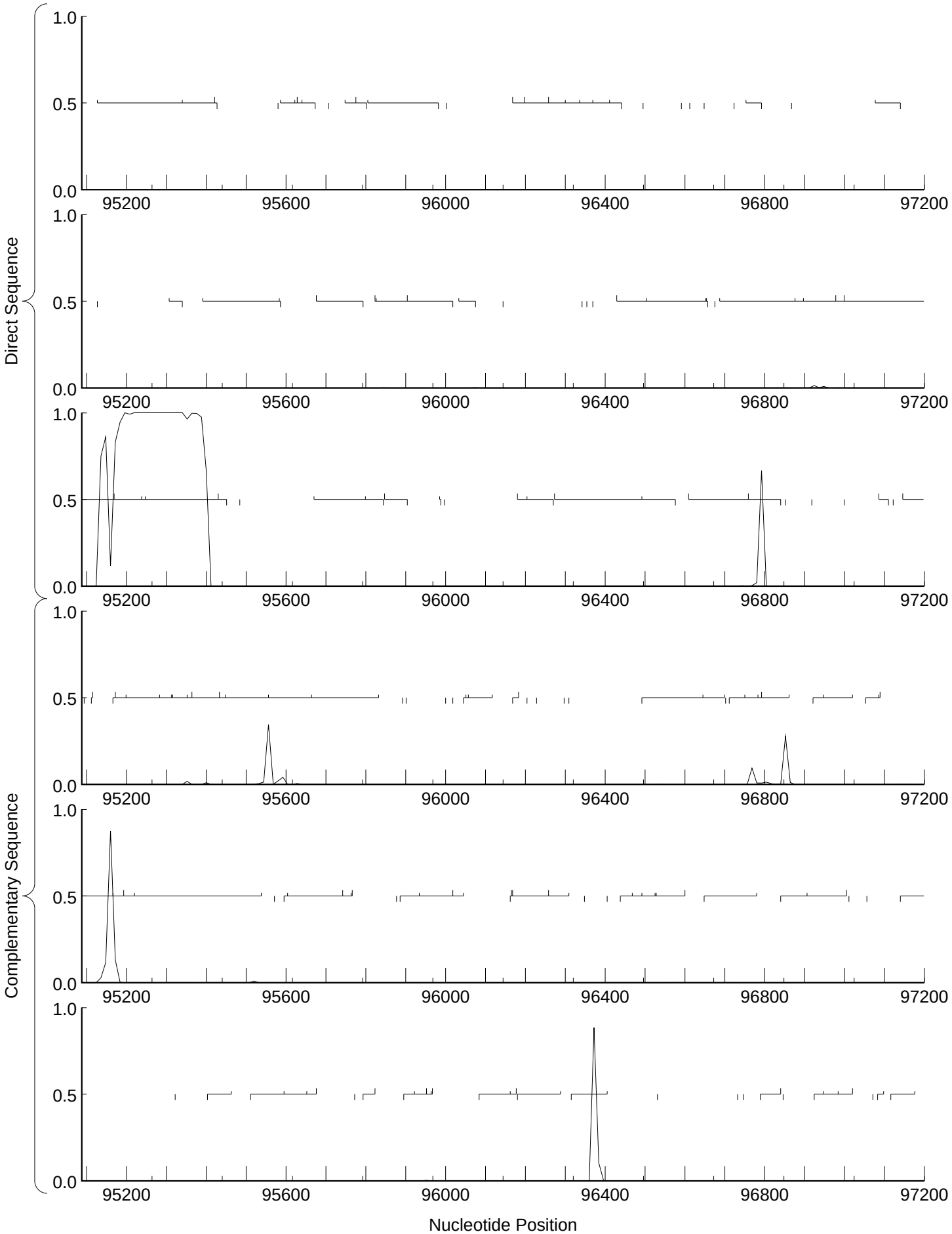
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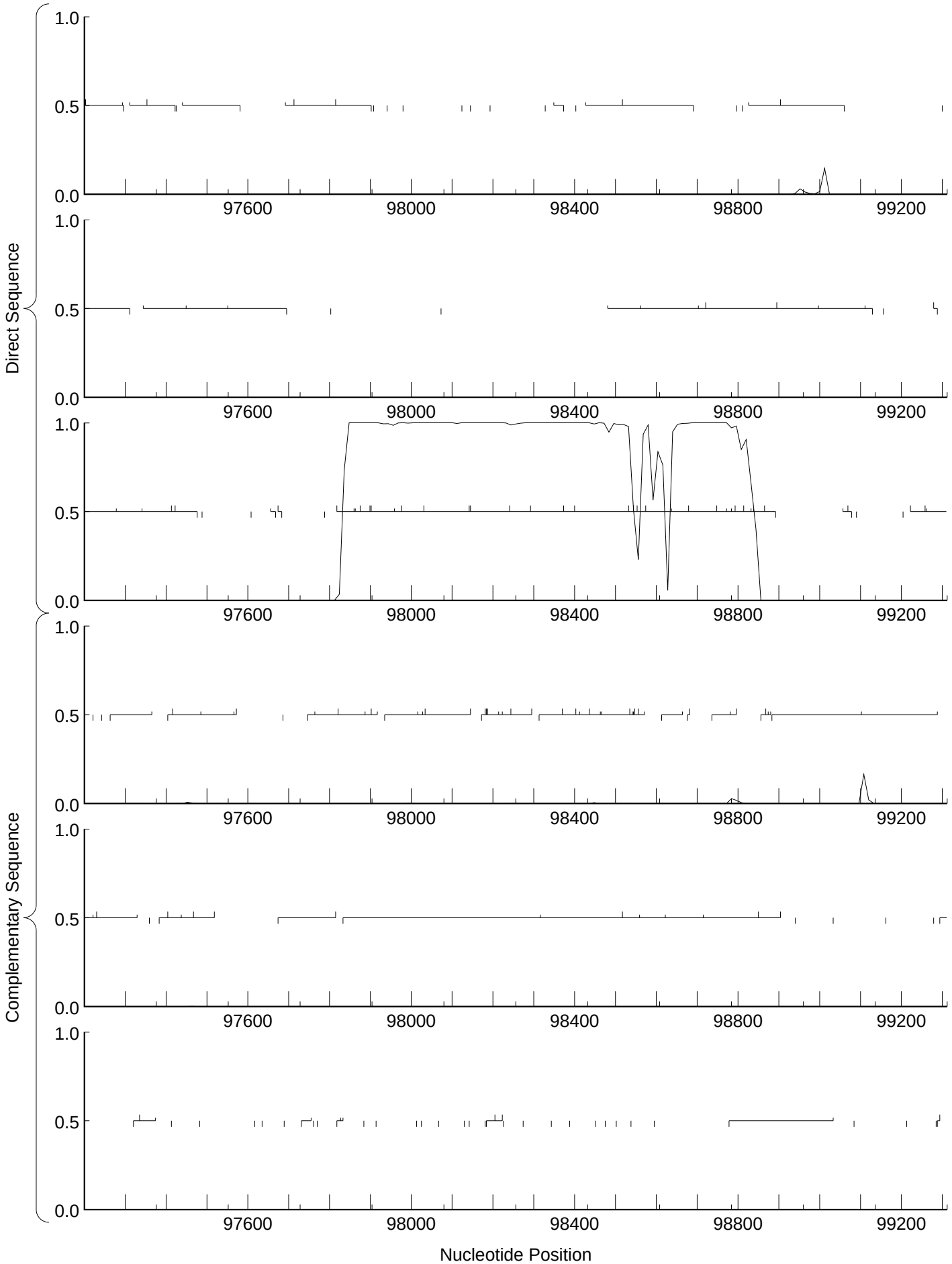
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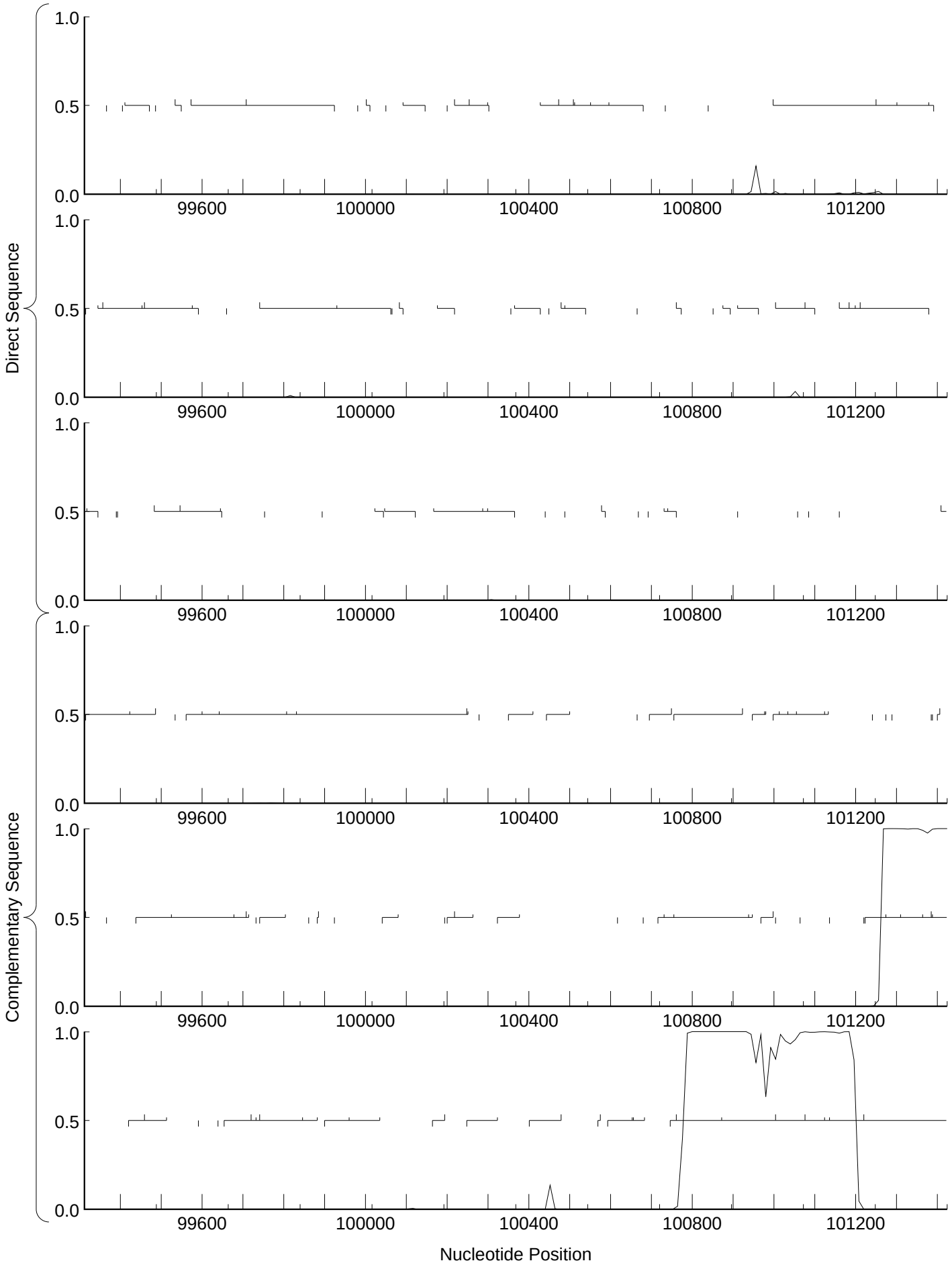


erium phage NuevoMundo Complete Sequence, 155943 bp, Circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 47/75

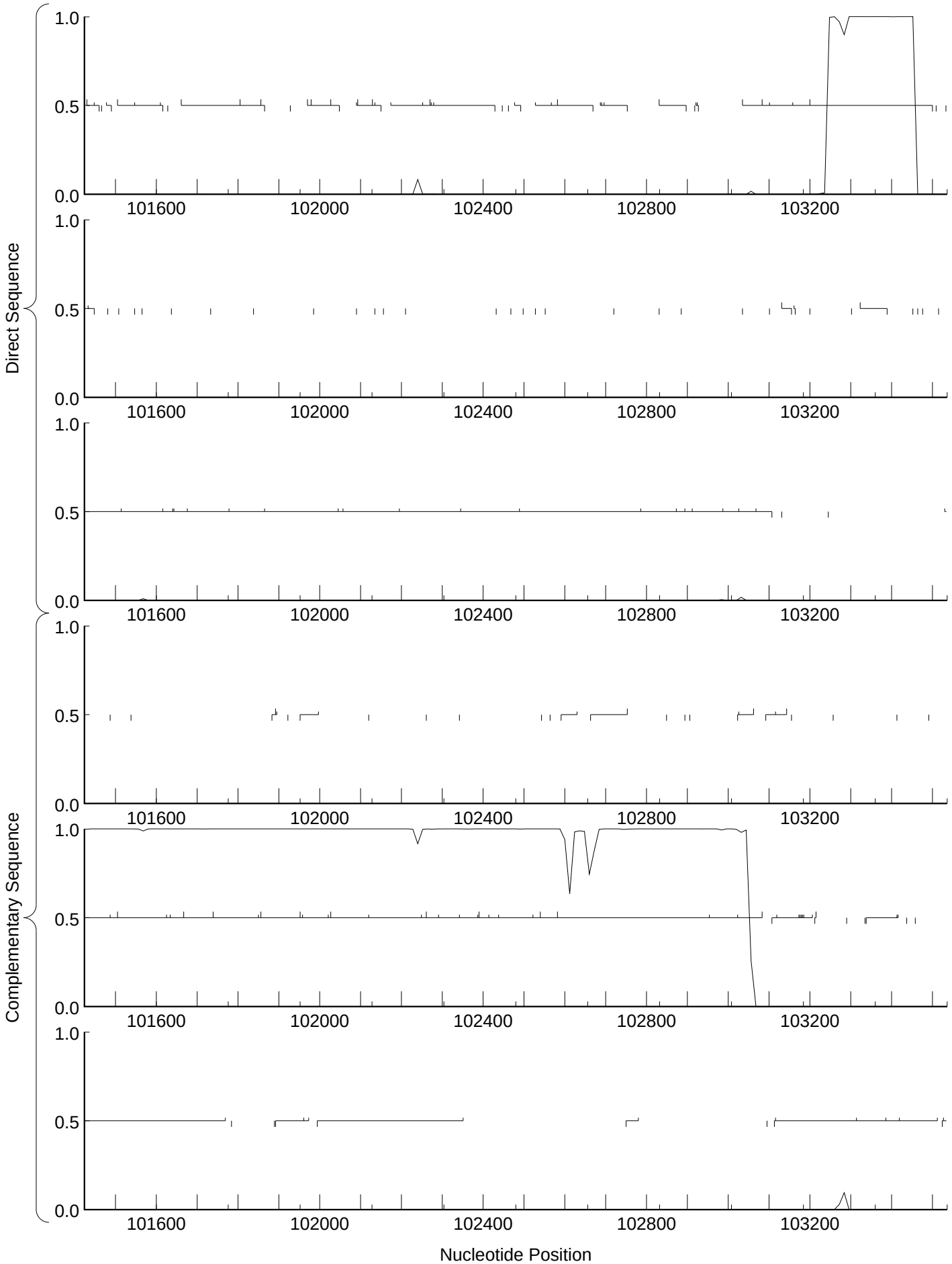


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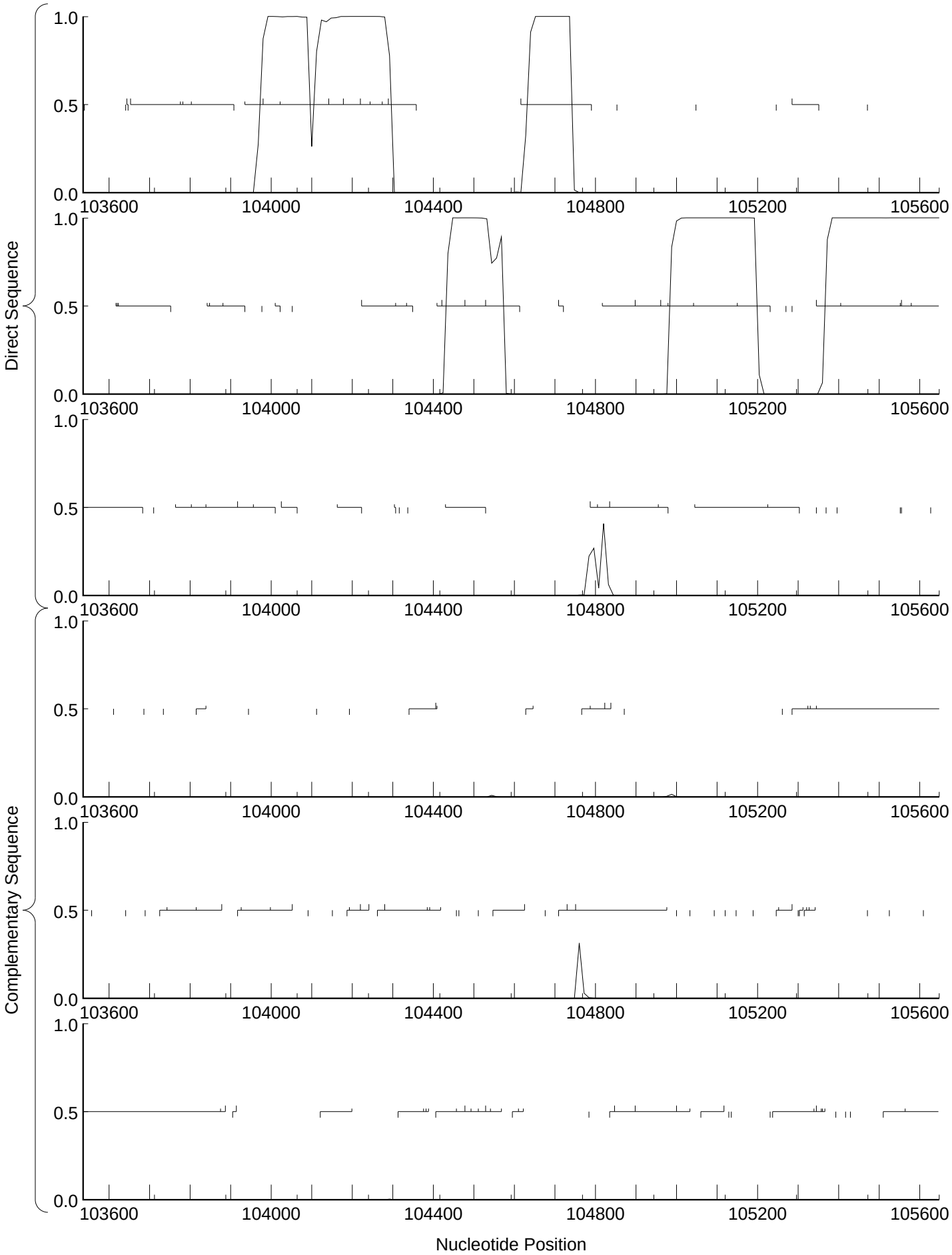



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
99600
100000
100400
100800
101200
Nucleotide Position

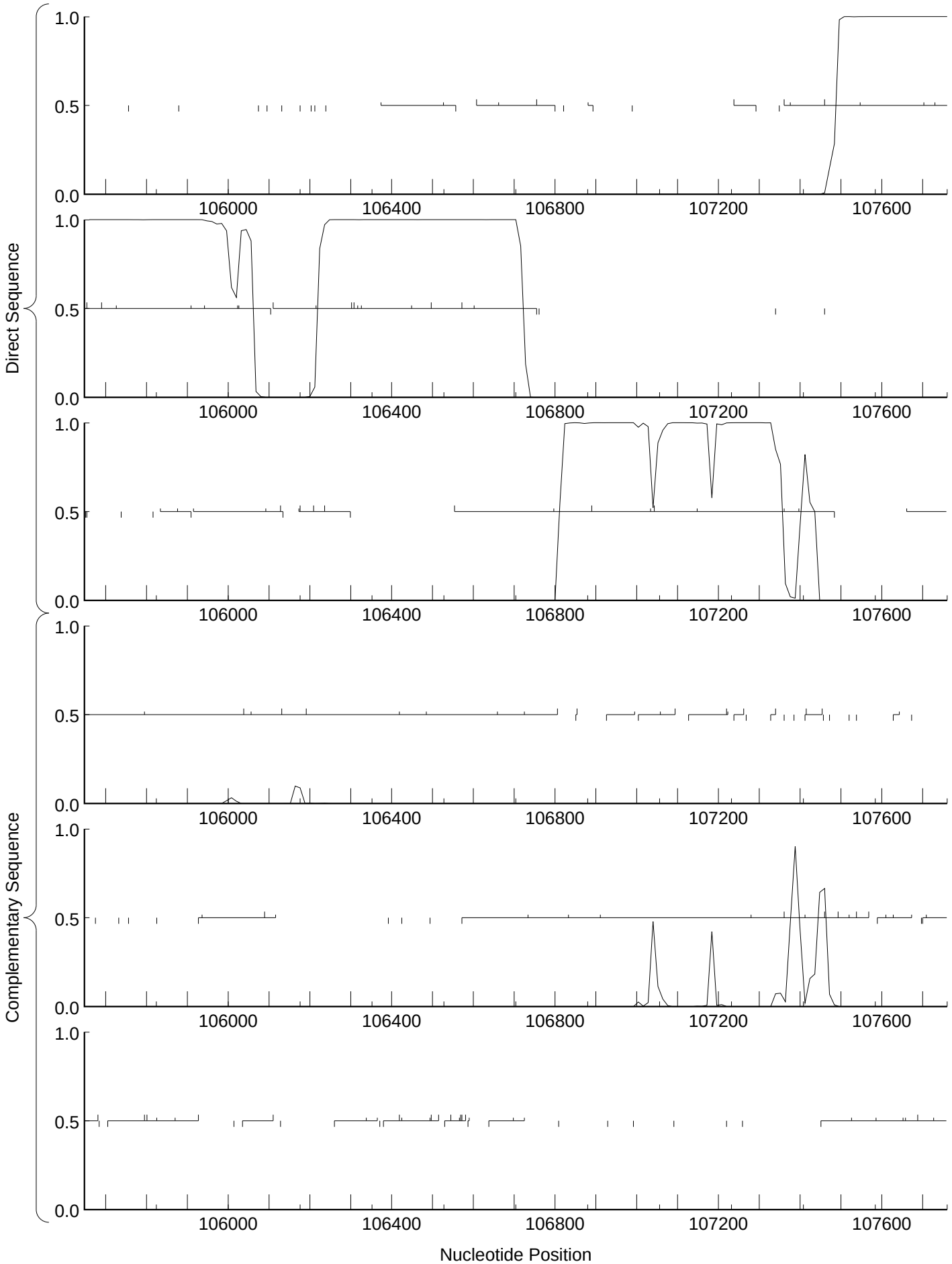
erium phage NuevoMundo Complete Sequence, 155943 bp, Circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 50/75



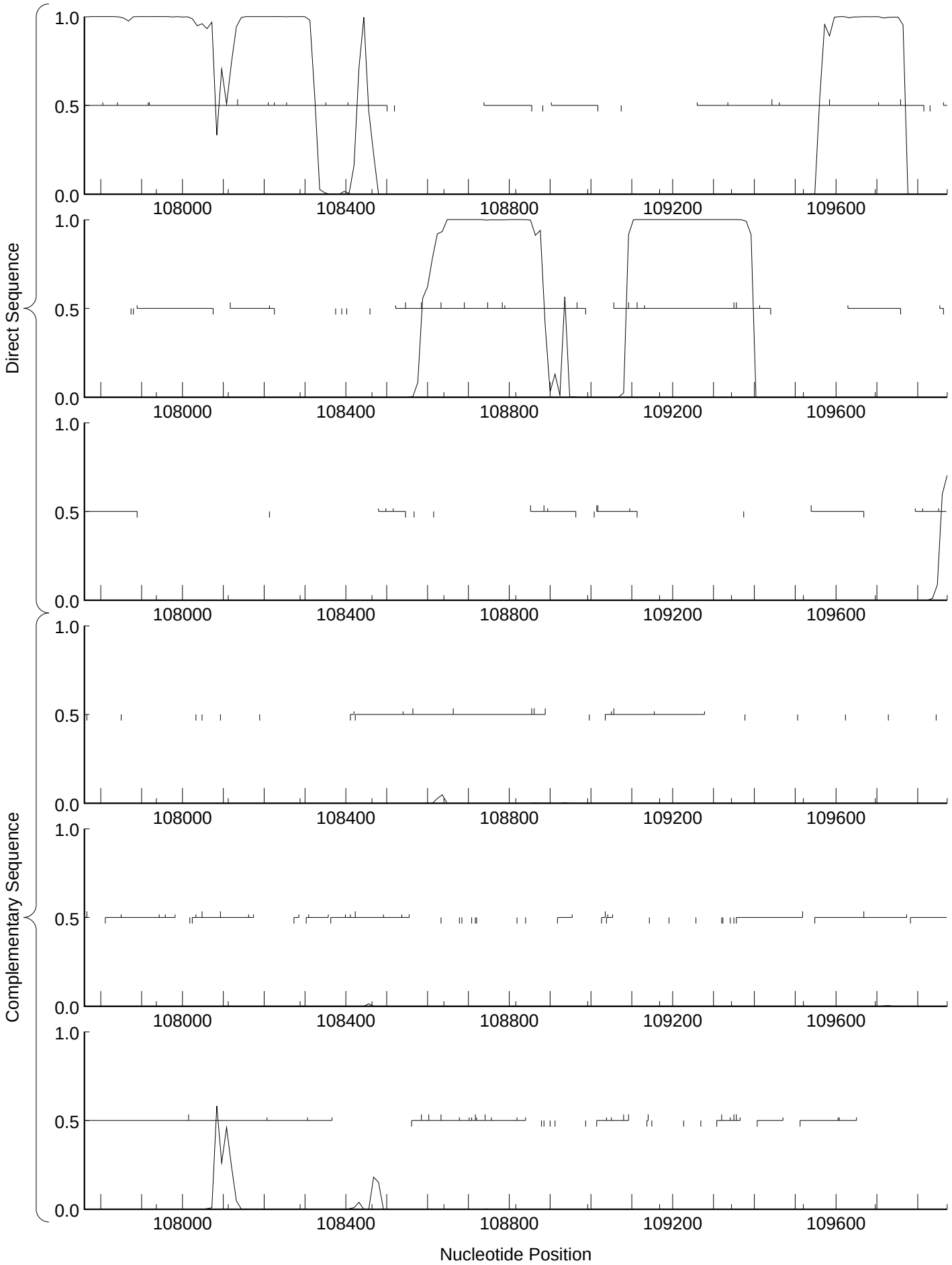
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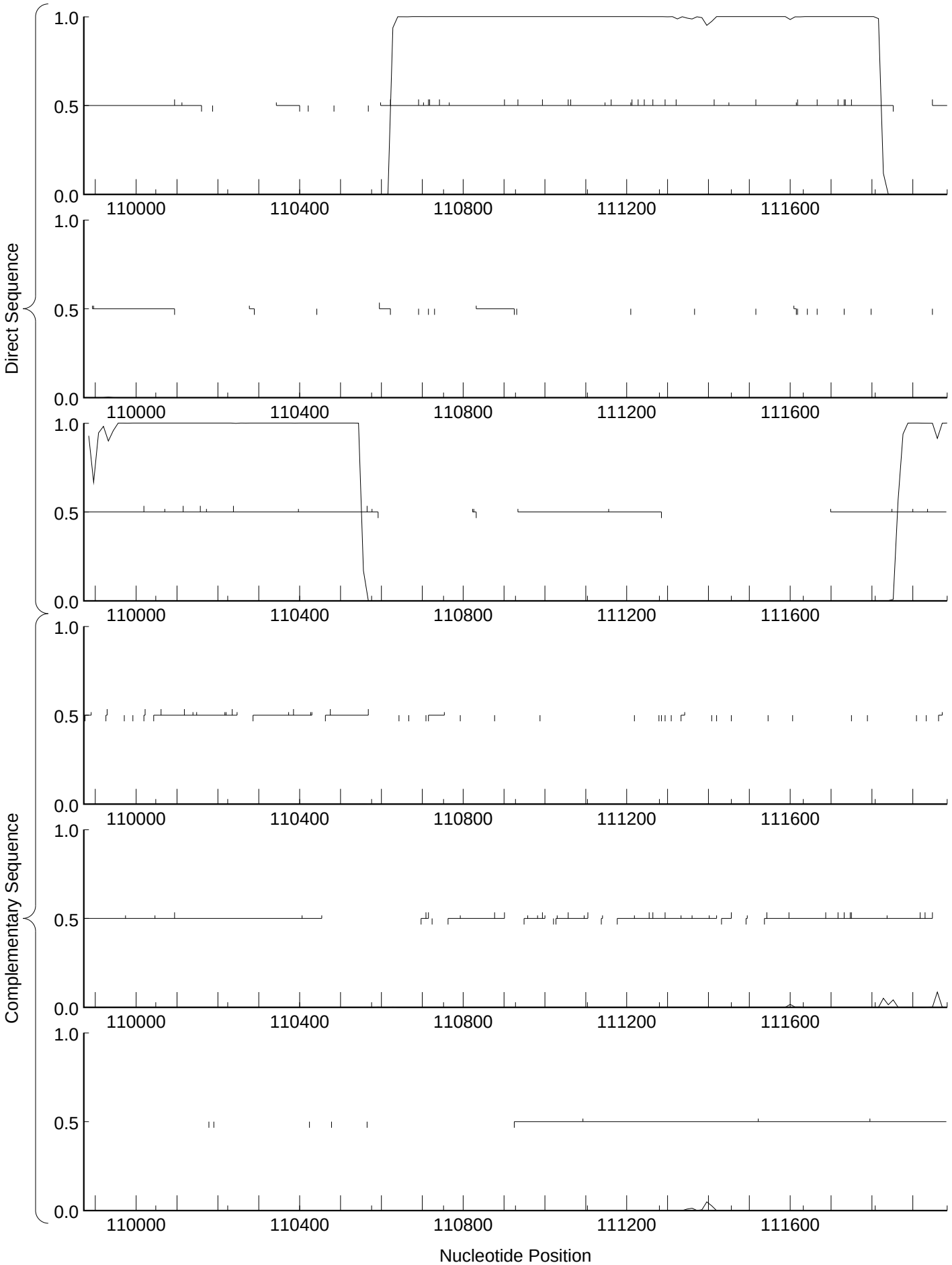
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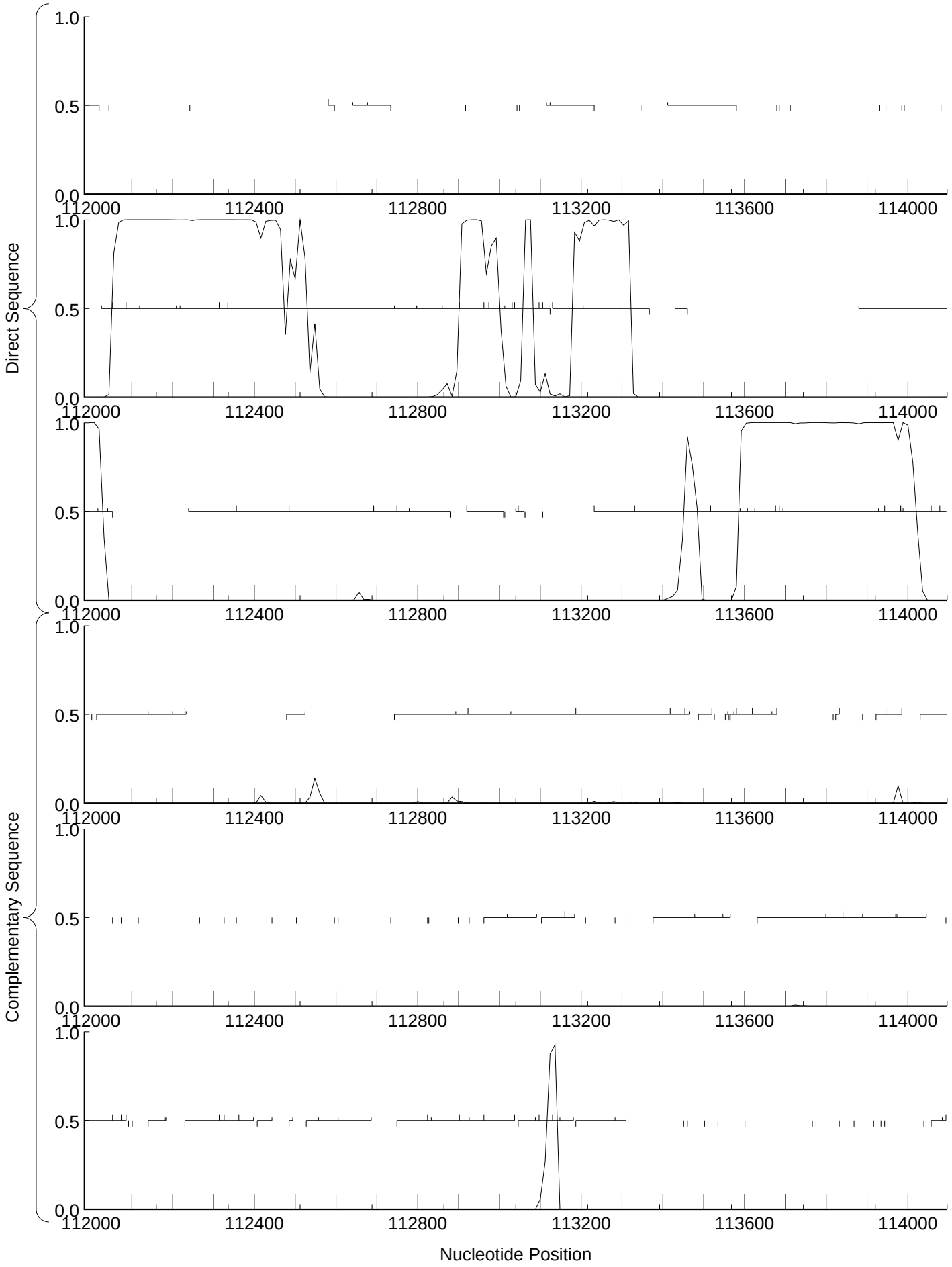
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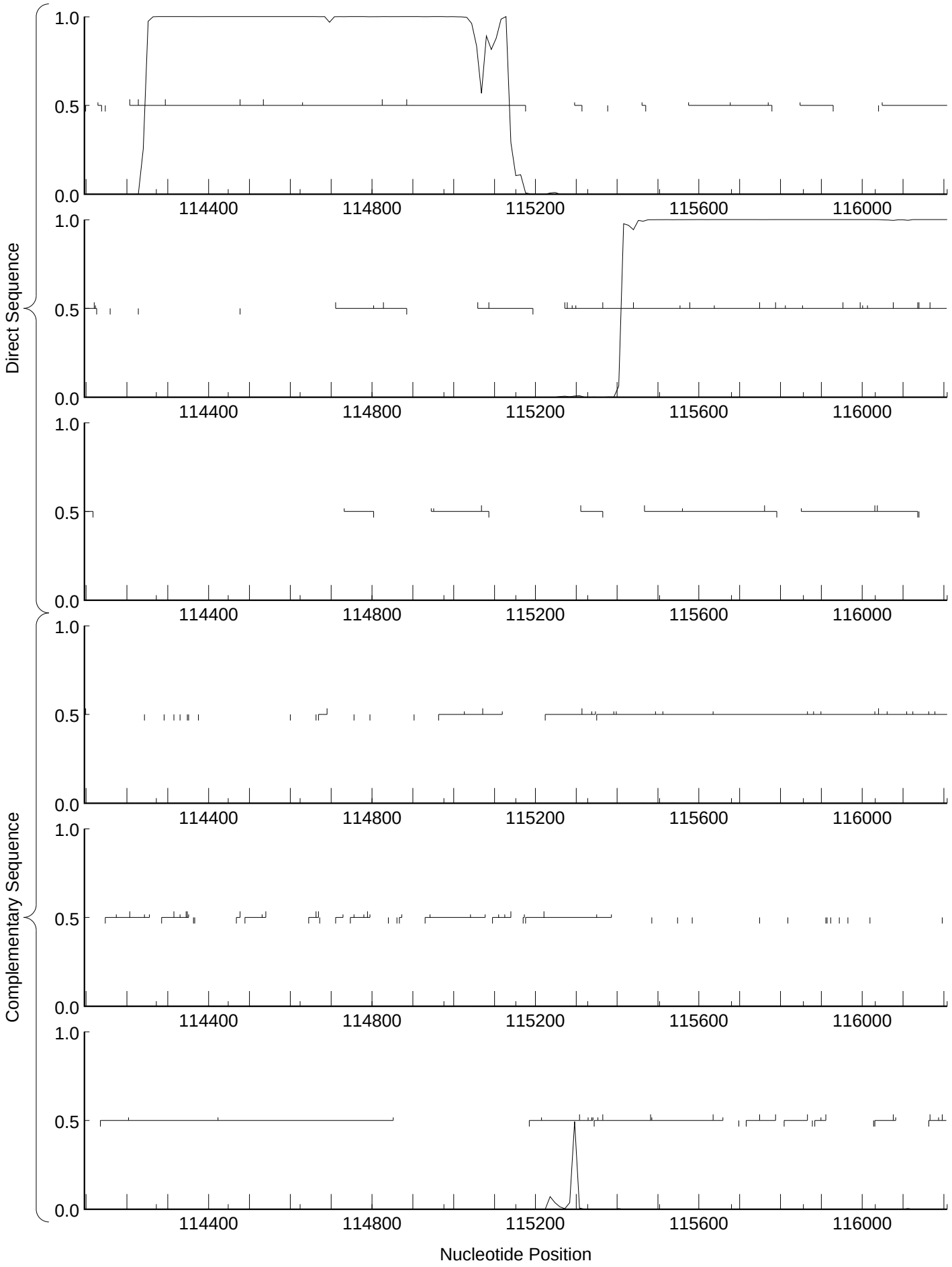


erium phage NuevoMundo Complete Sequence, 155943 bp, Circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 54/75

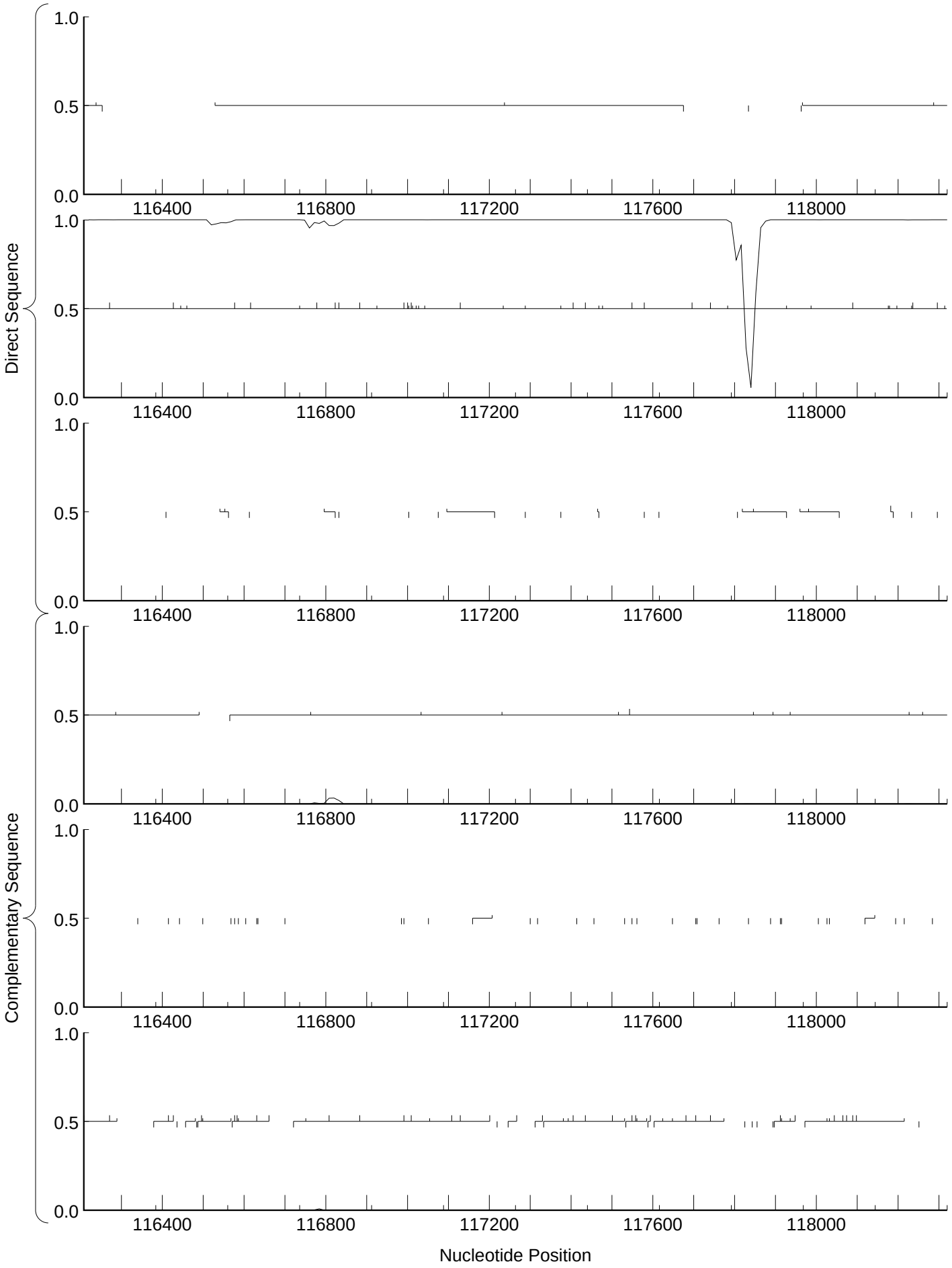


erium phage NuevoMundo Complete Sequence, 155943 bp, Circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 55/75

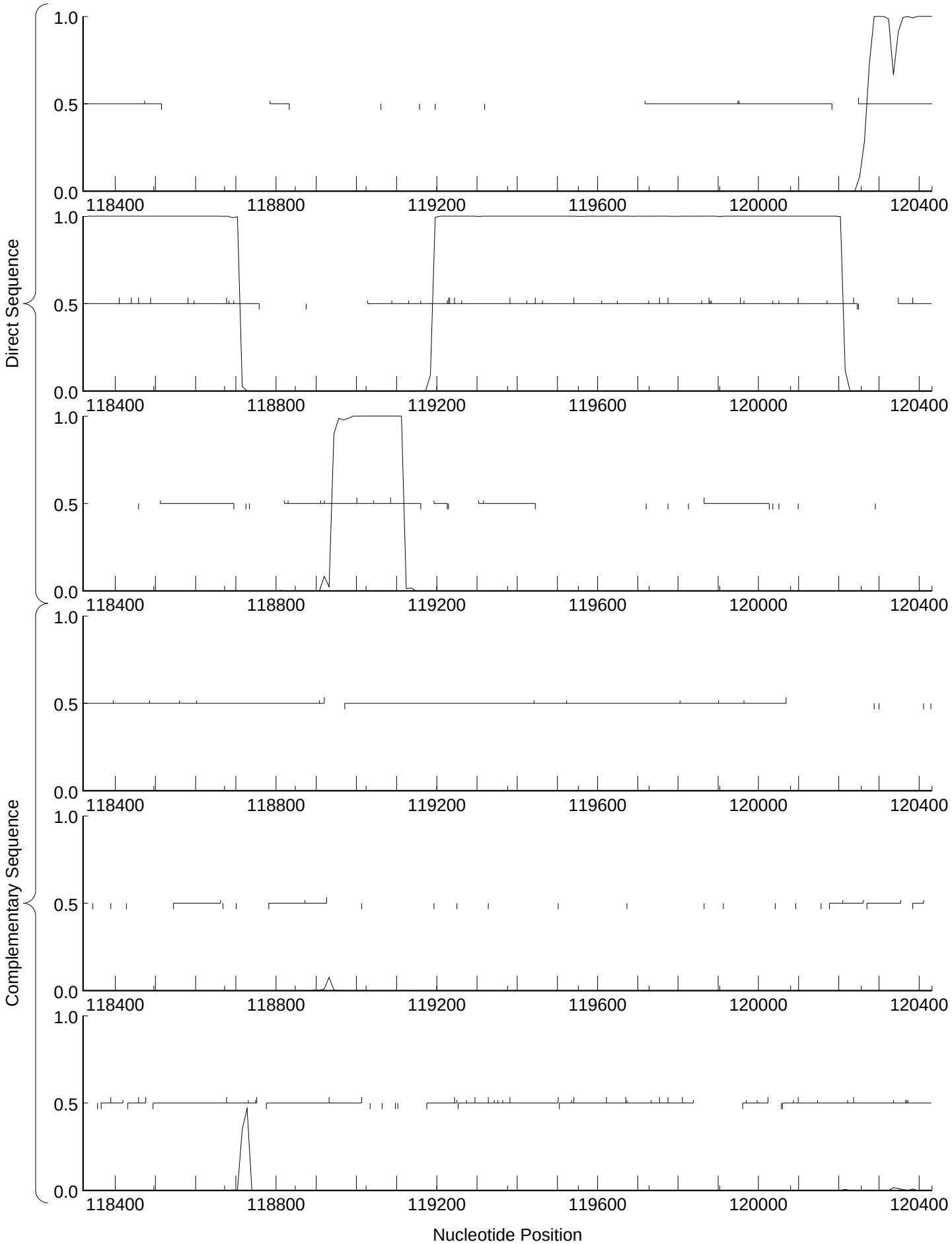



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
114400
114800
115200
115600
116000
Nucleotide Position

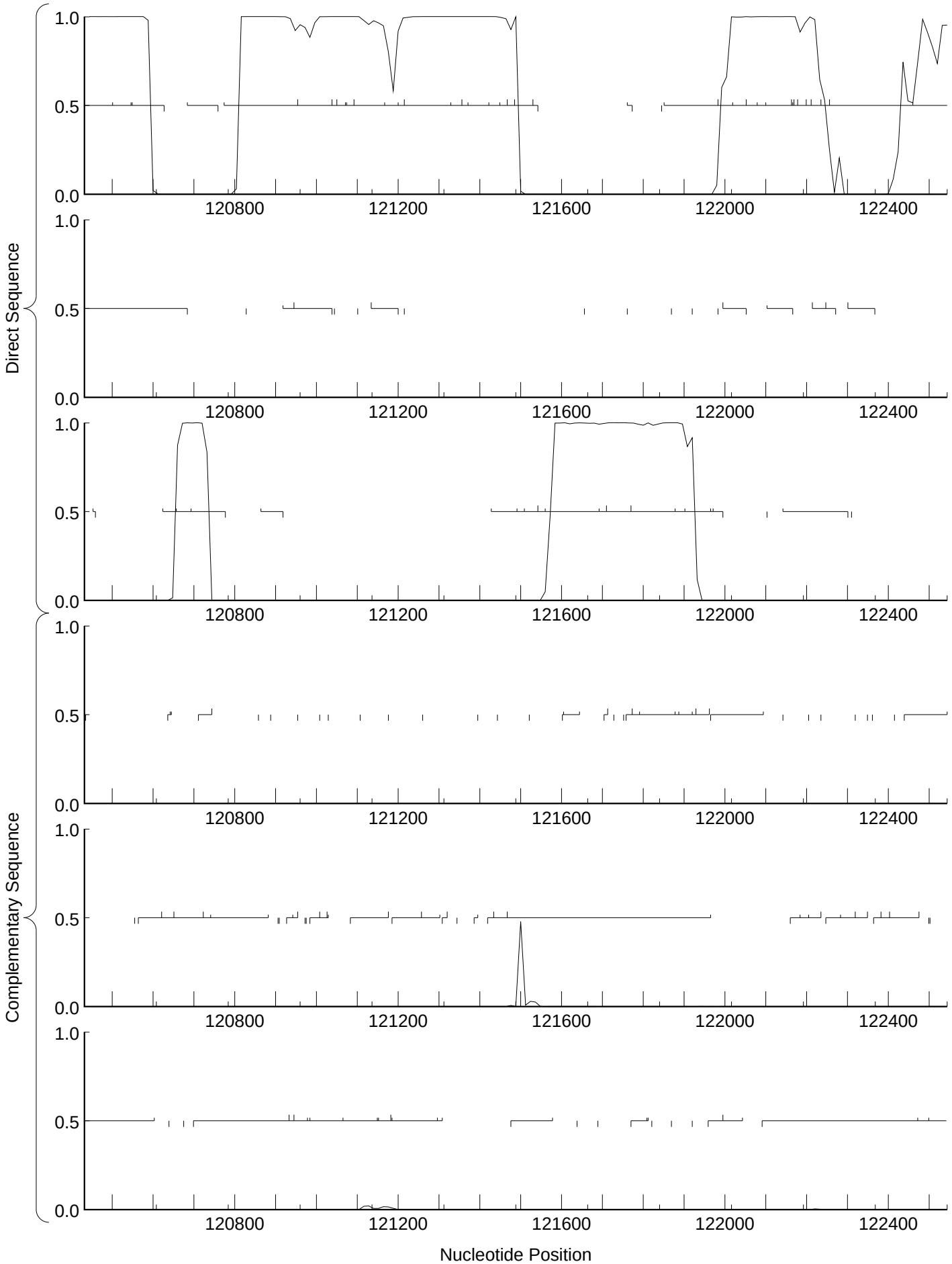
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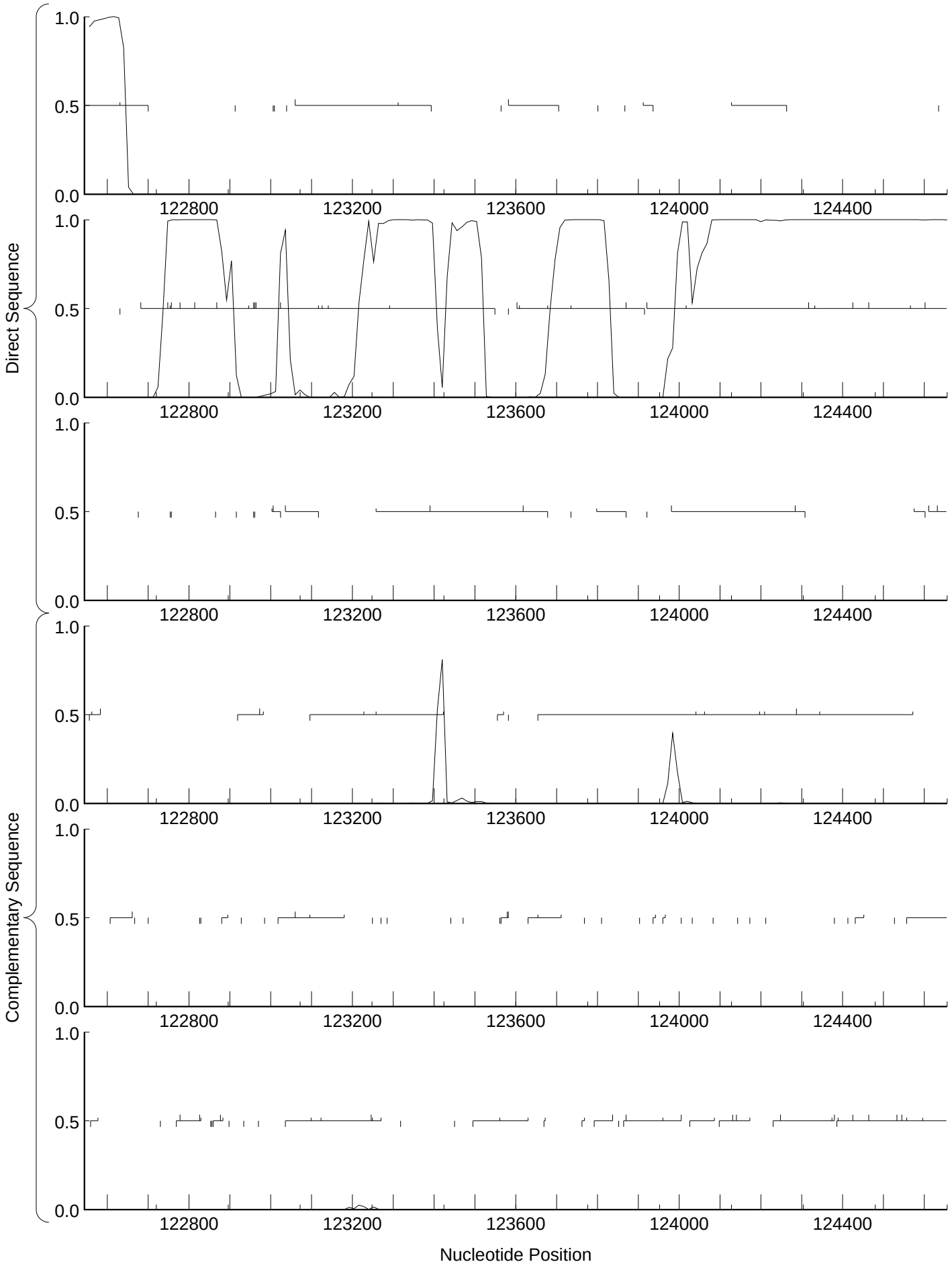
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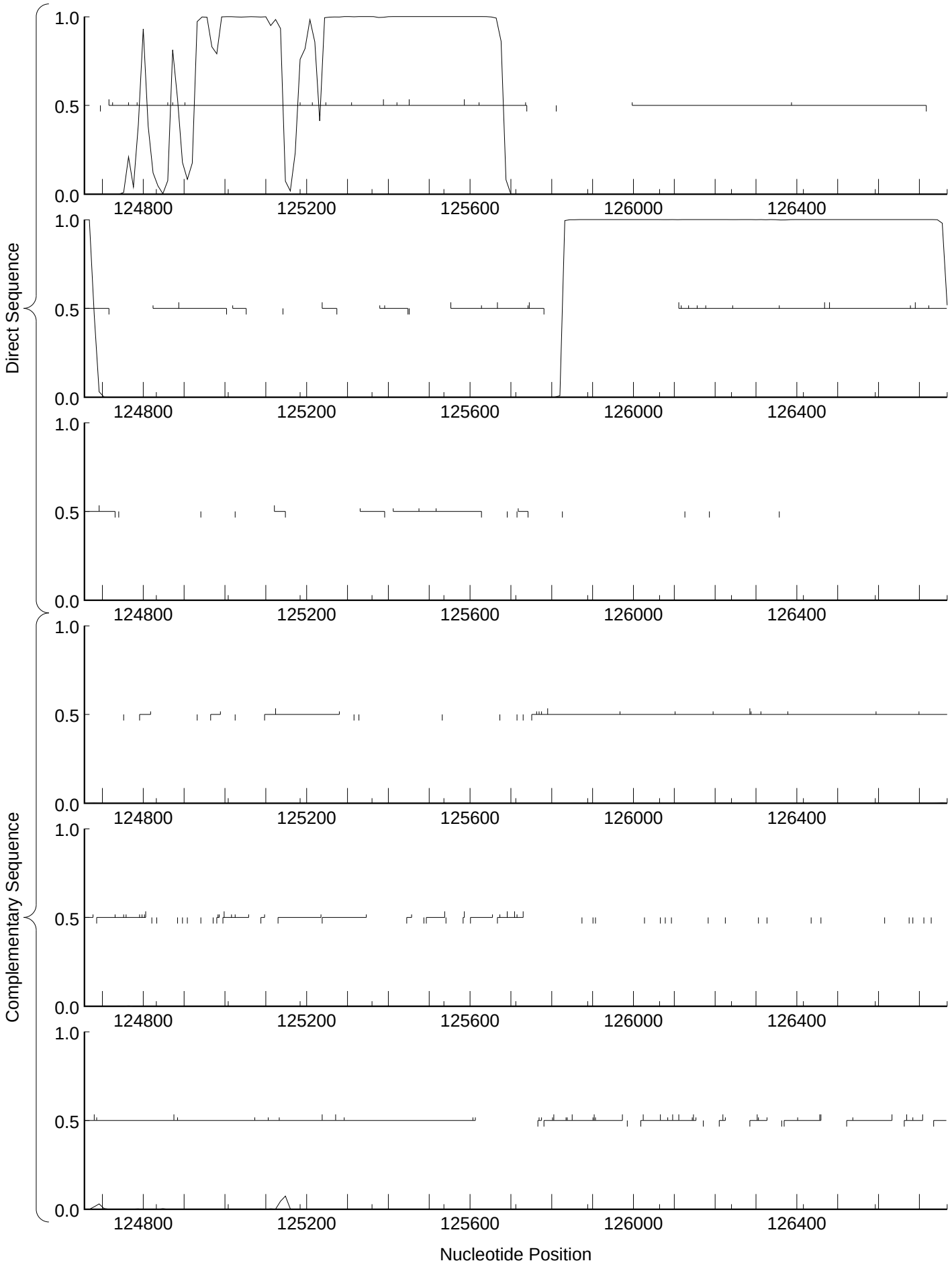


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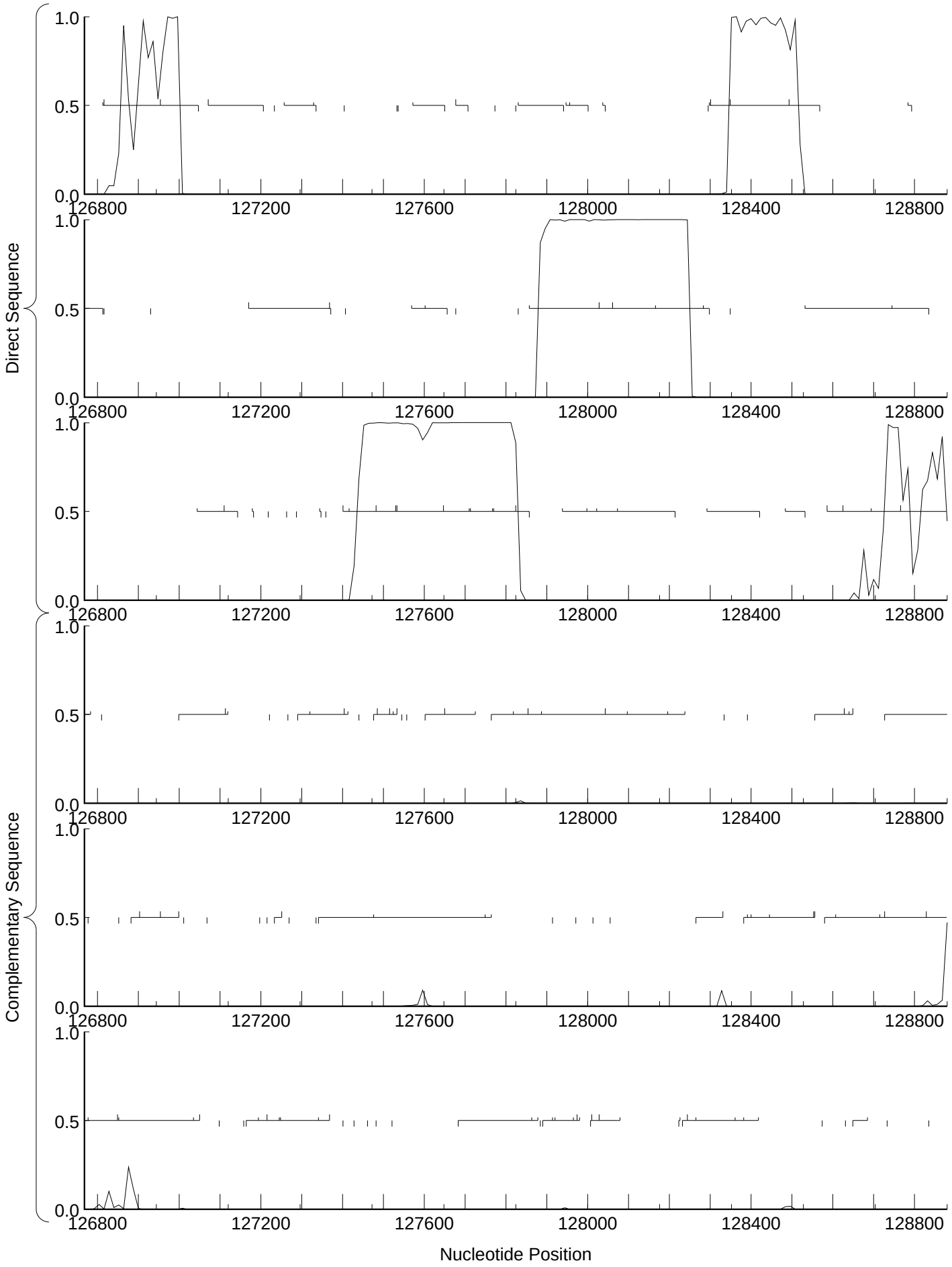


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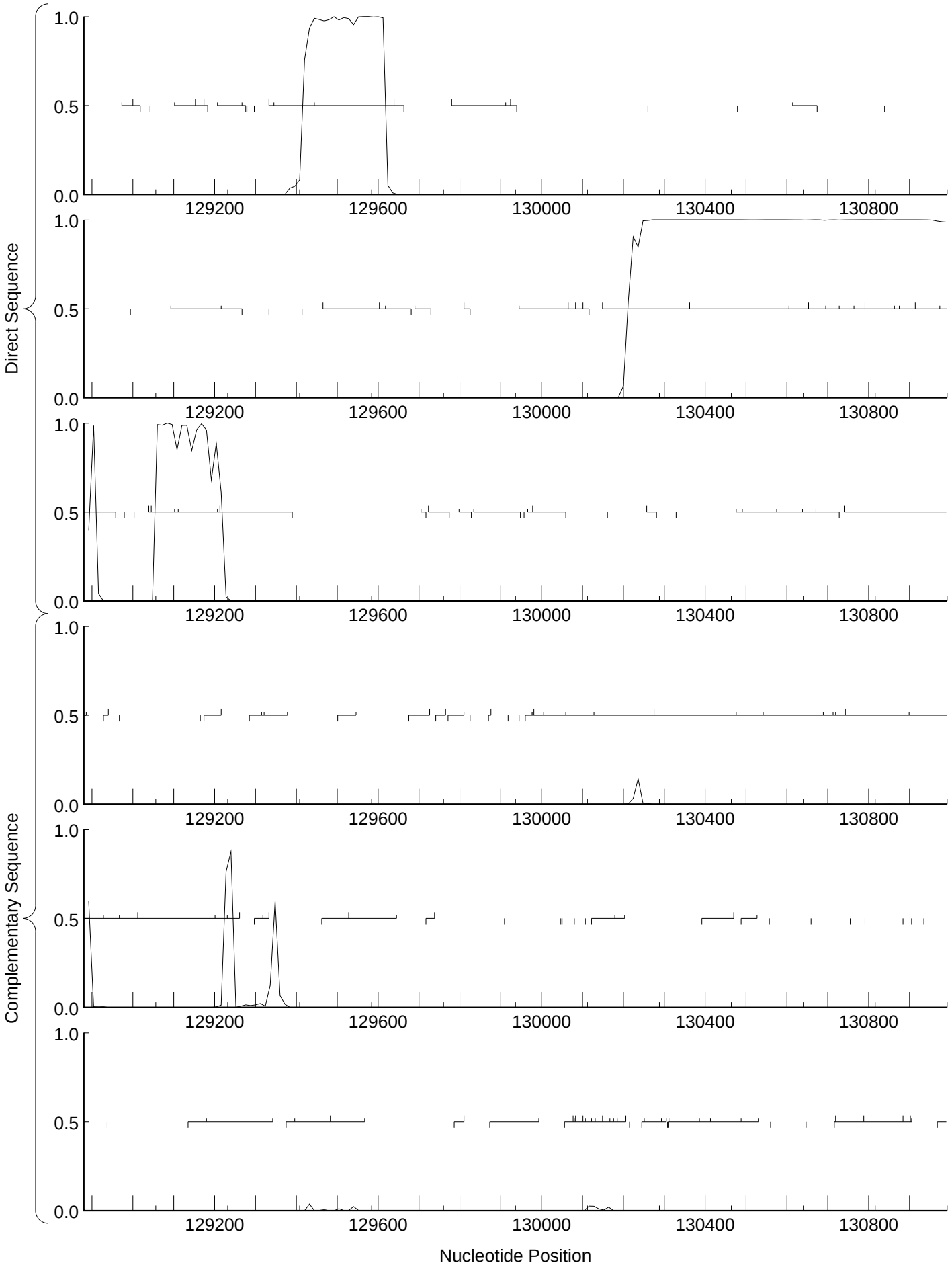



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
124800
125200
125600
126000
126400
Nucleotide Position

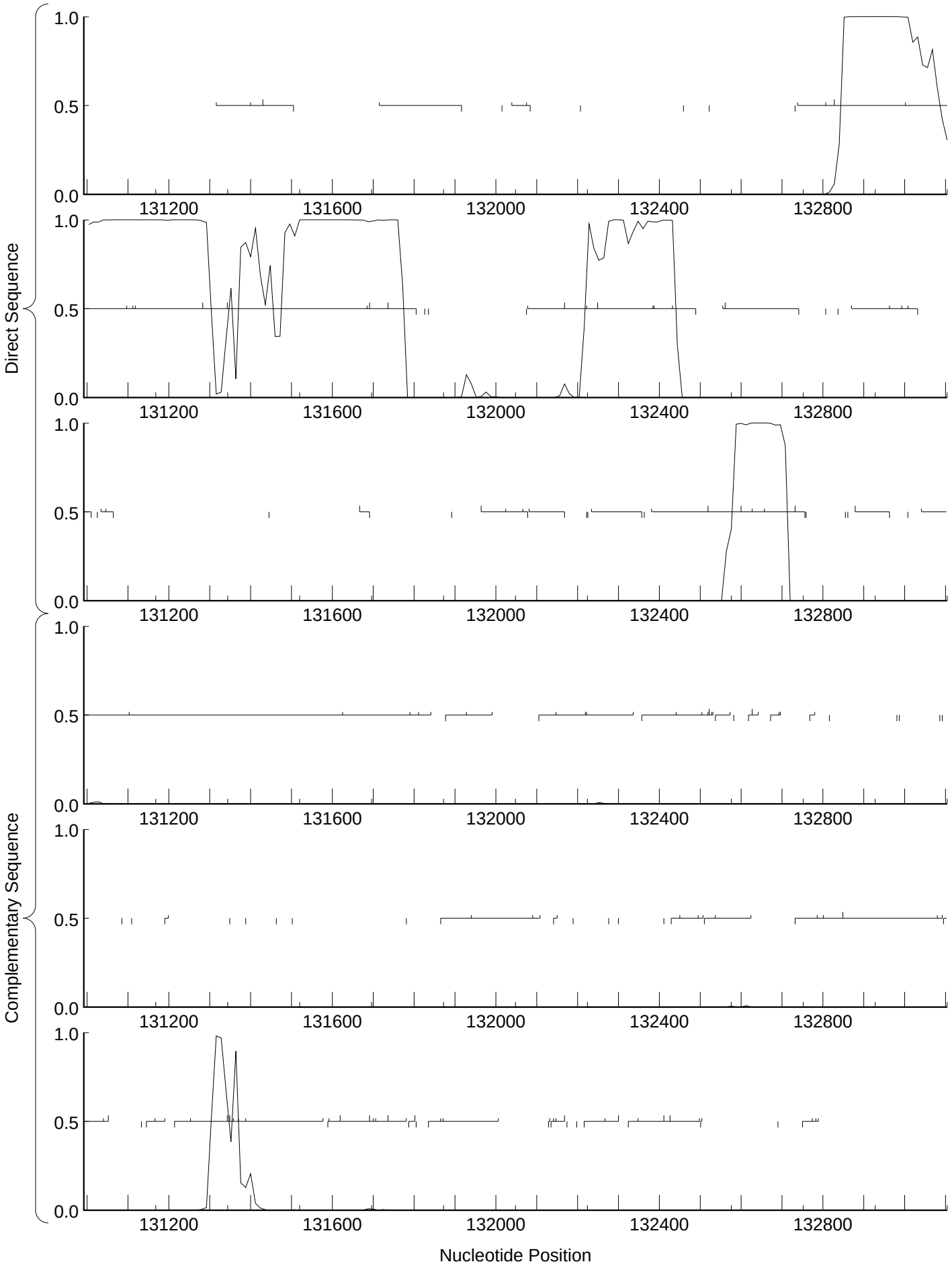
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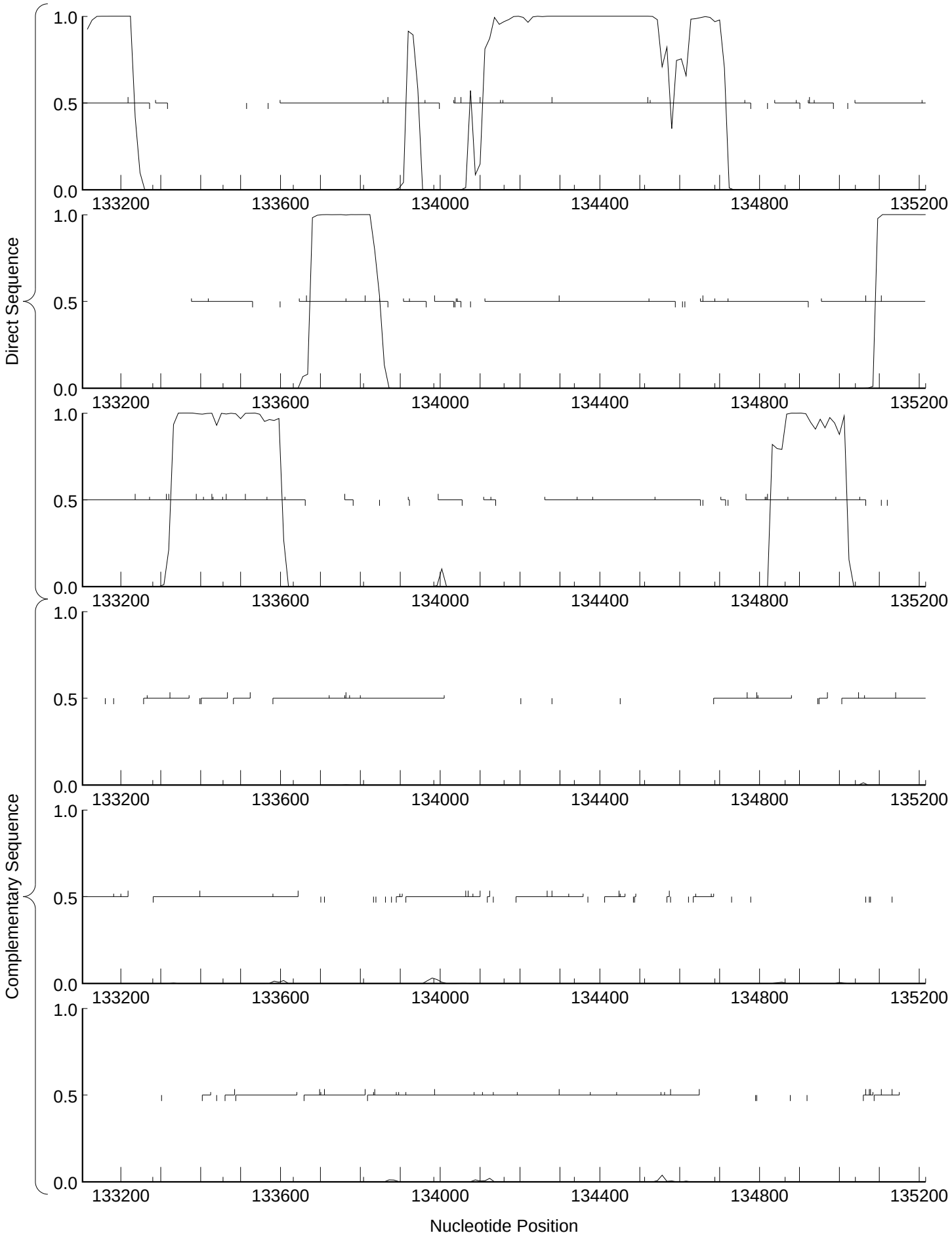
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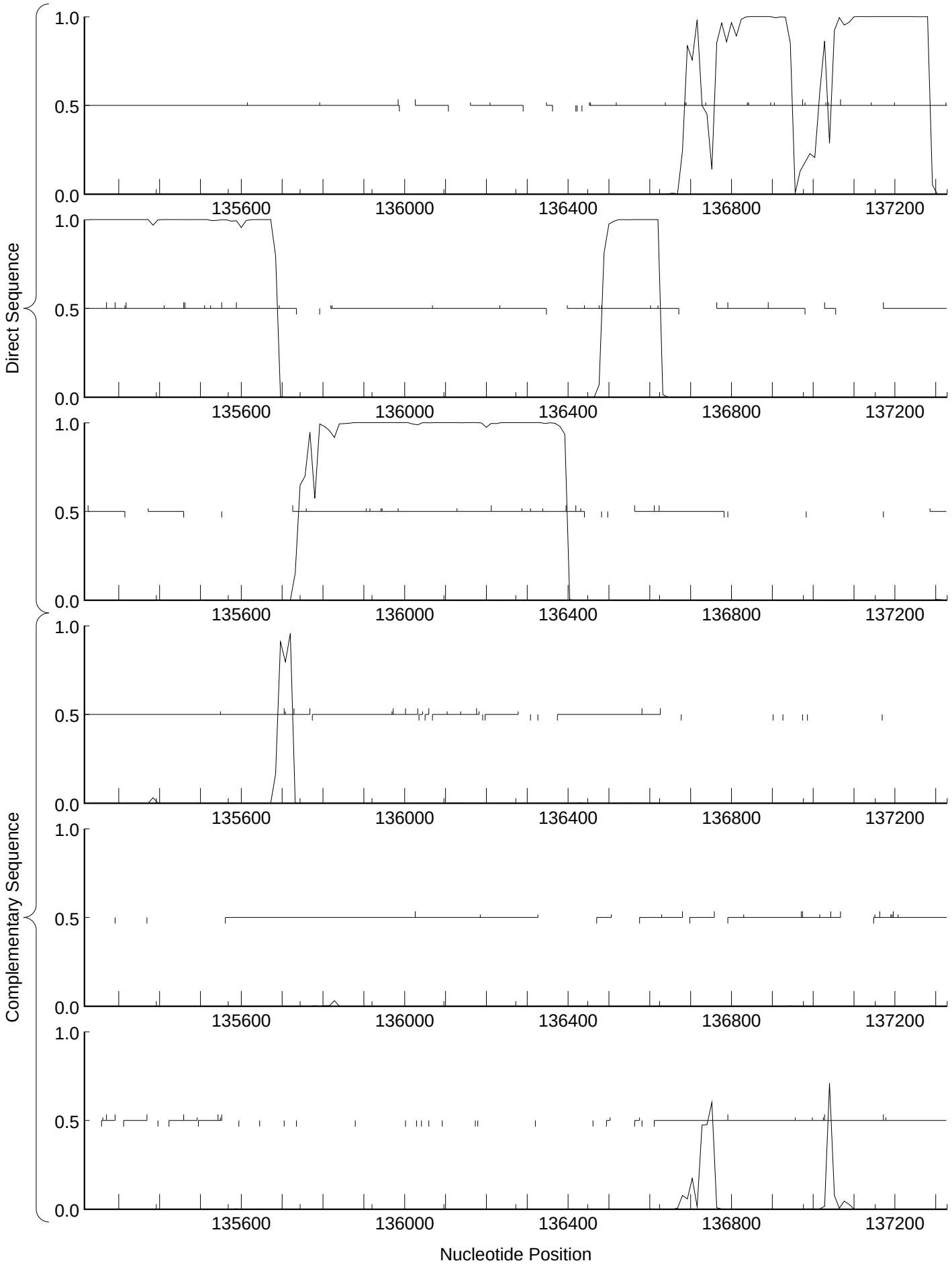


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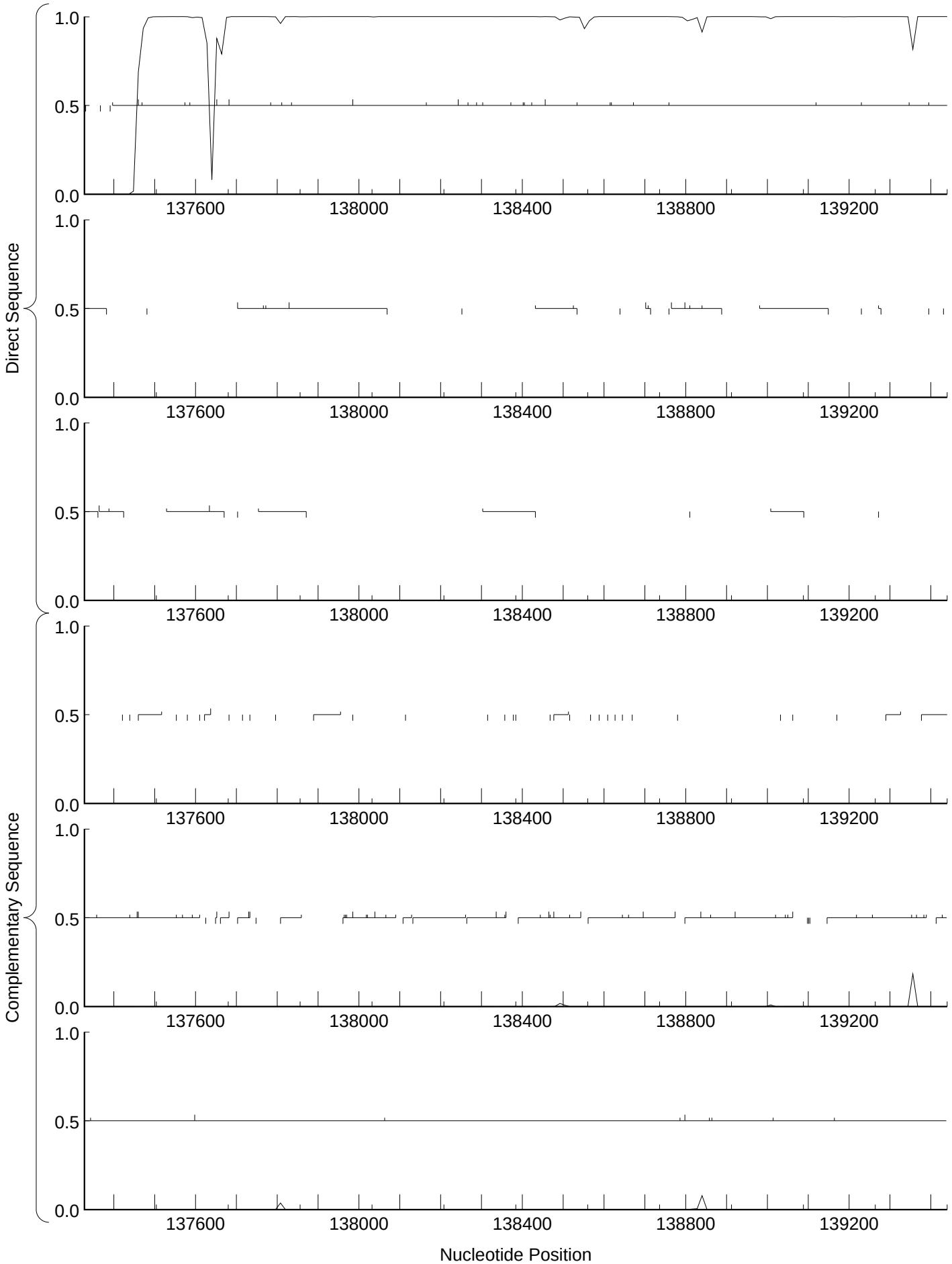


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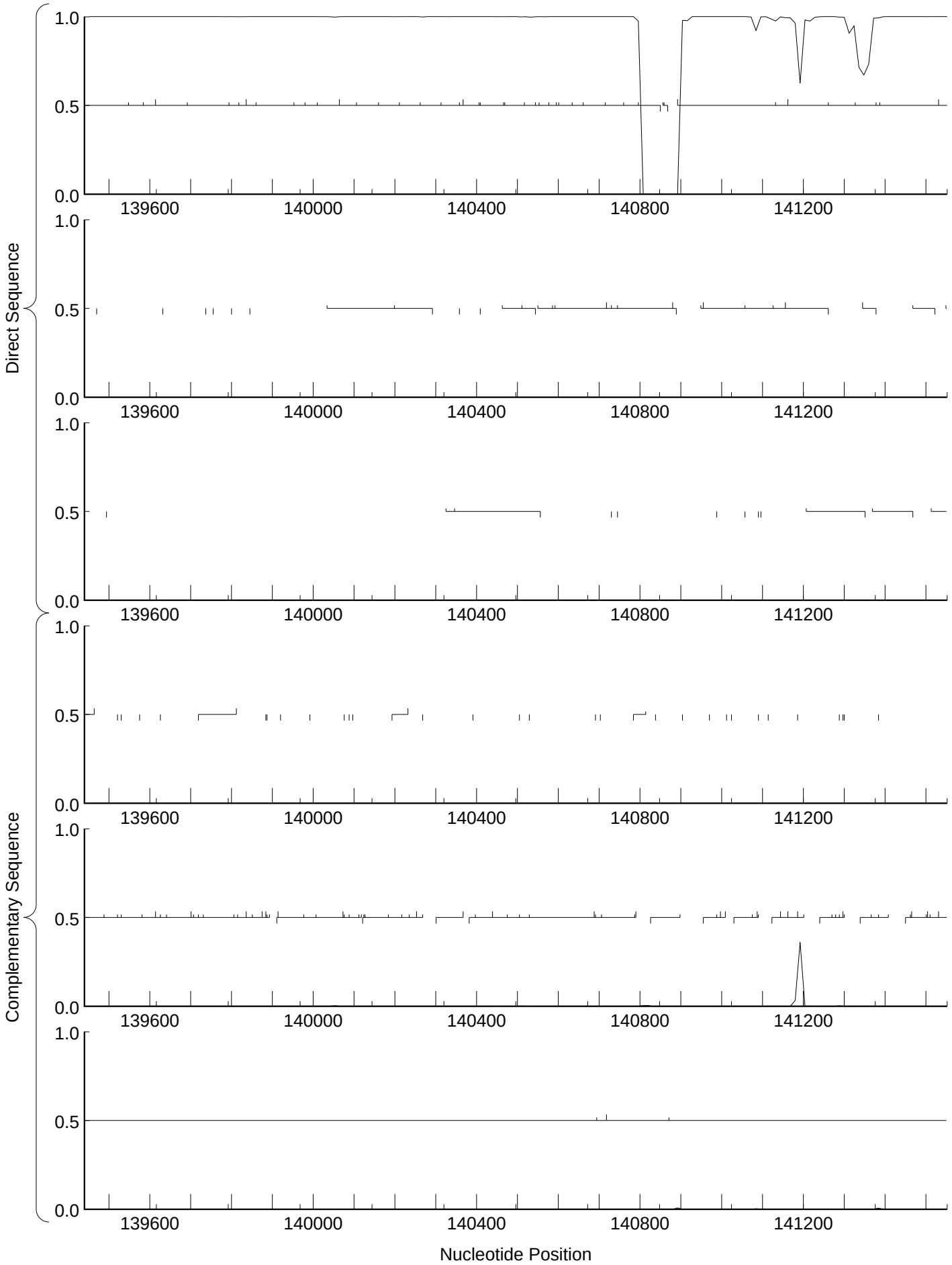



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
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136000
136400
136800
137200
Nucleotide Position

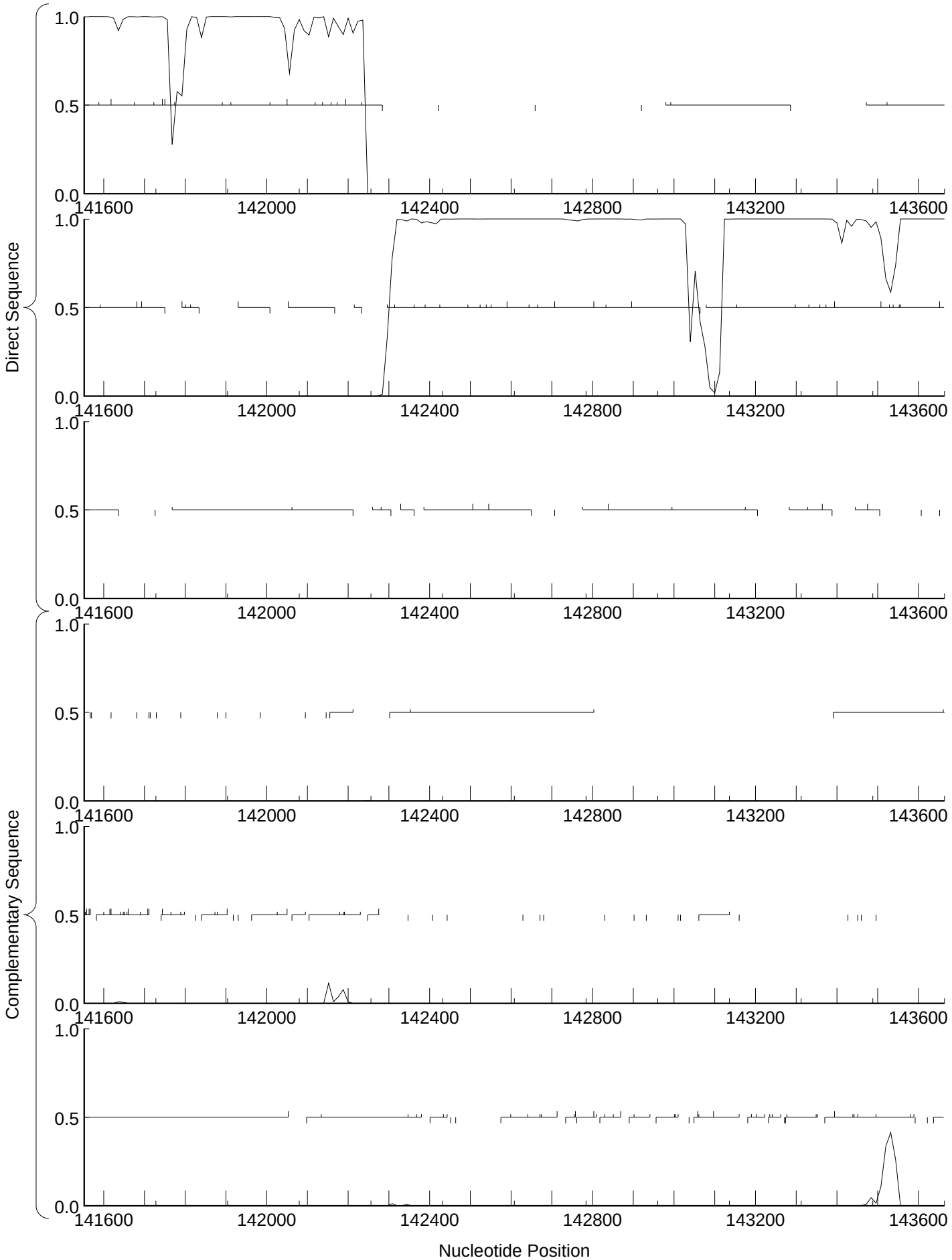
erium phage NuevoMundo Complete Sequence, 155943 bp, Circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 67/75



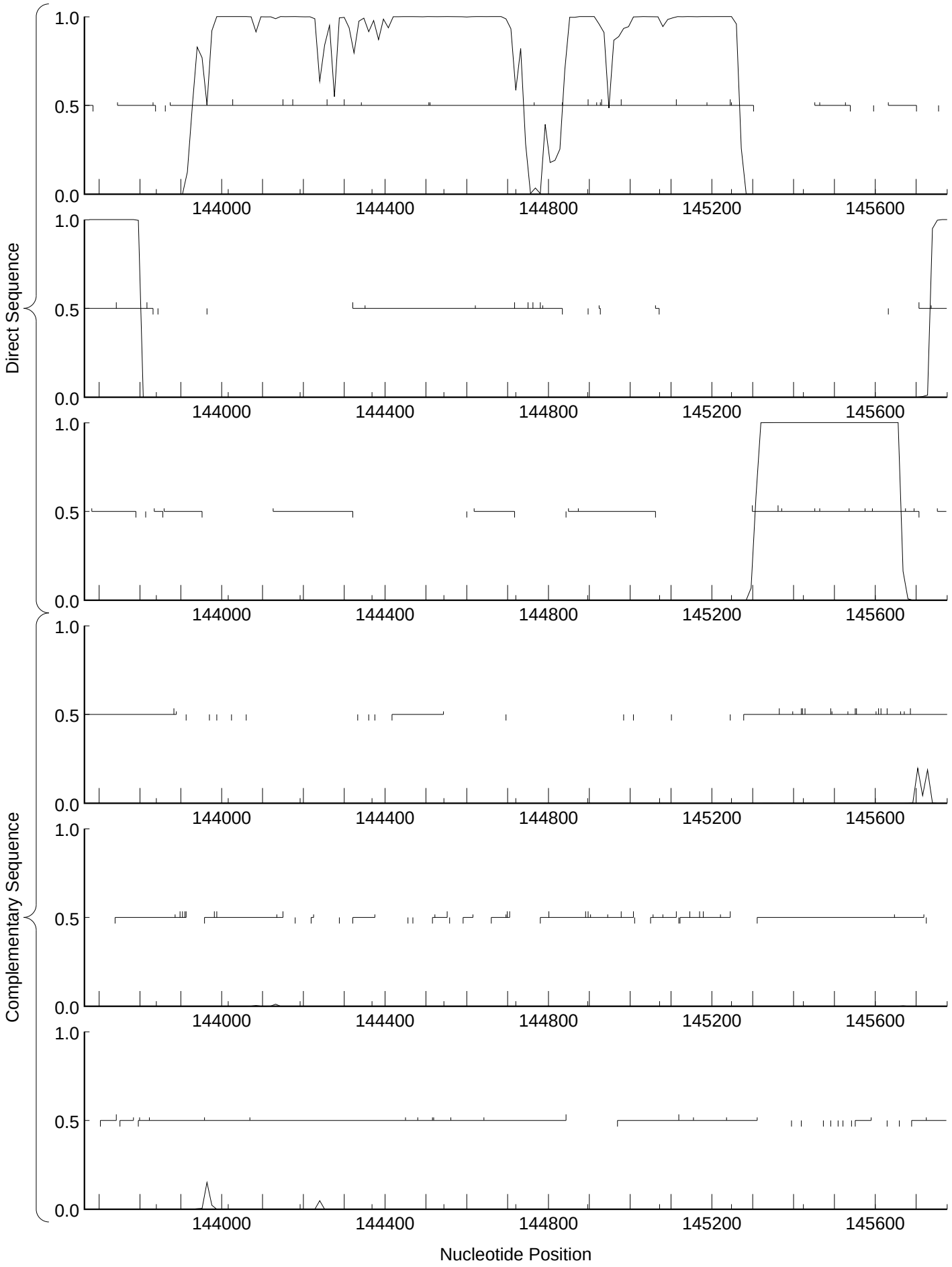
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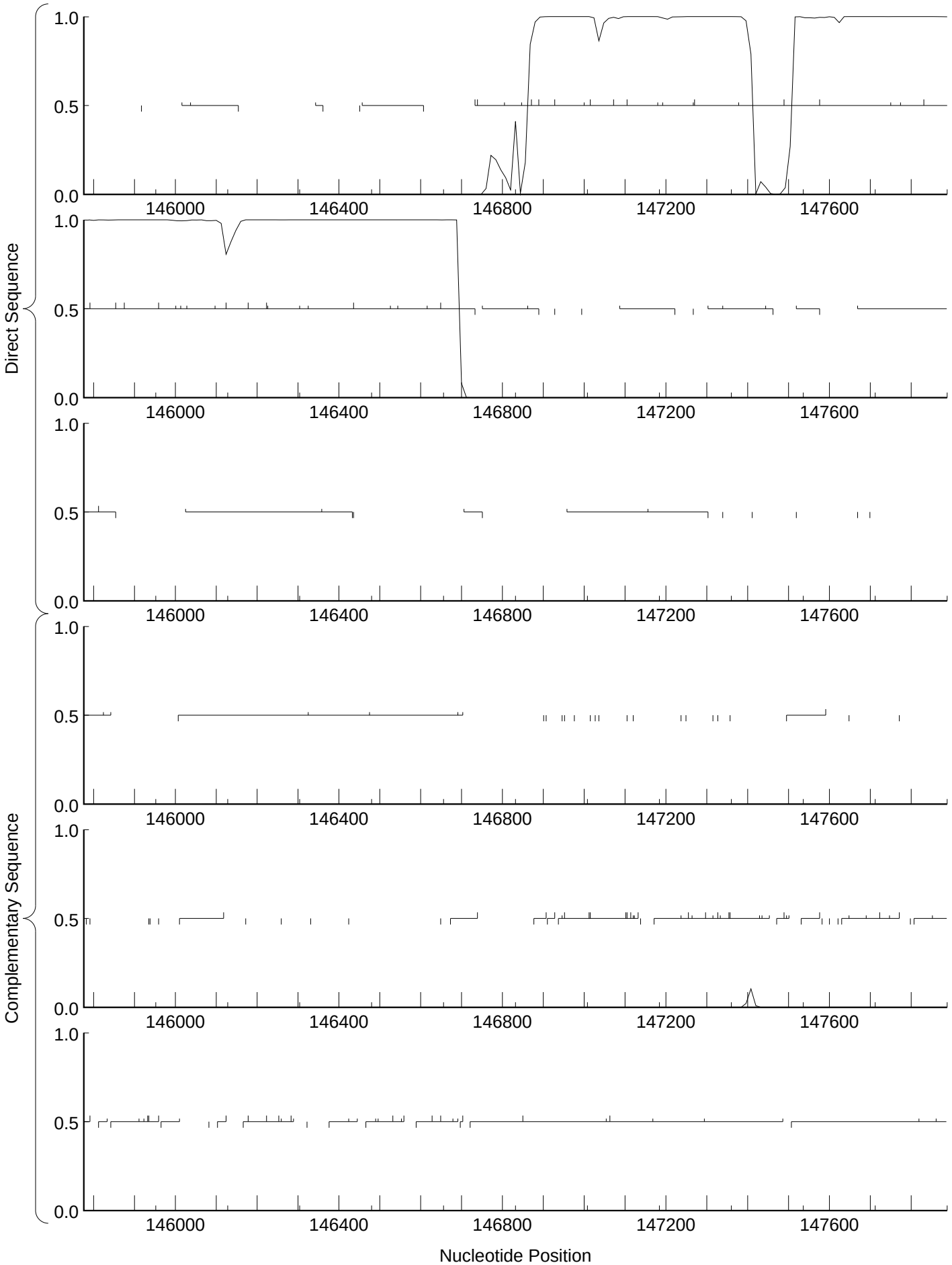
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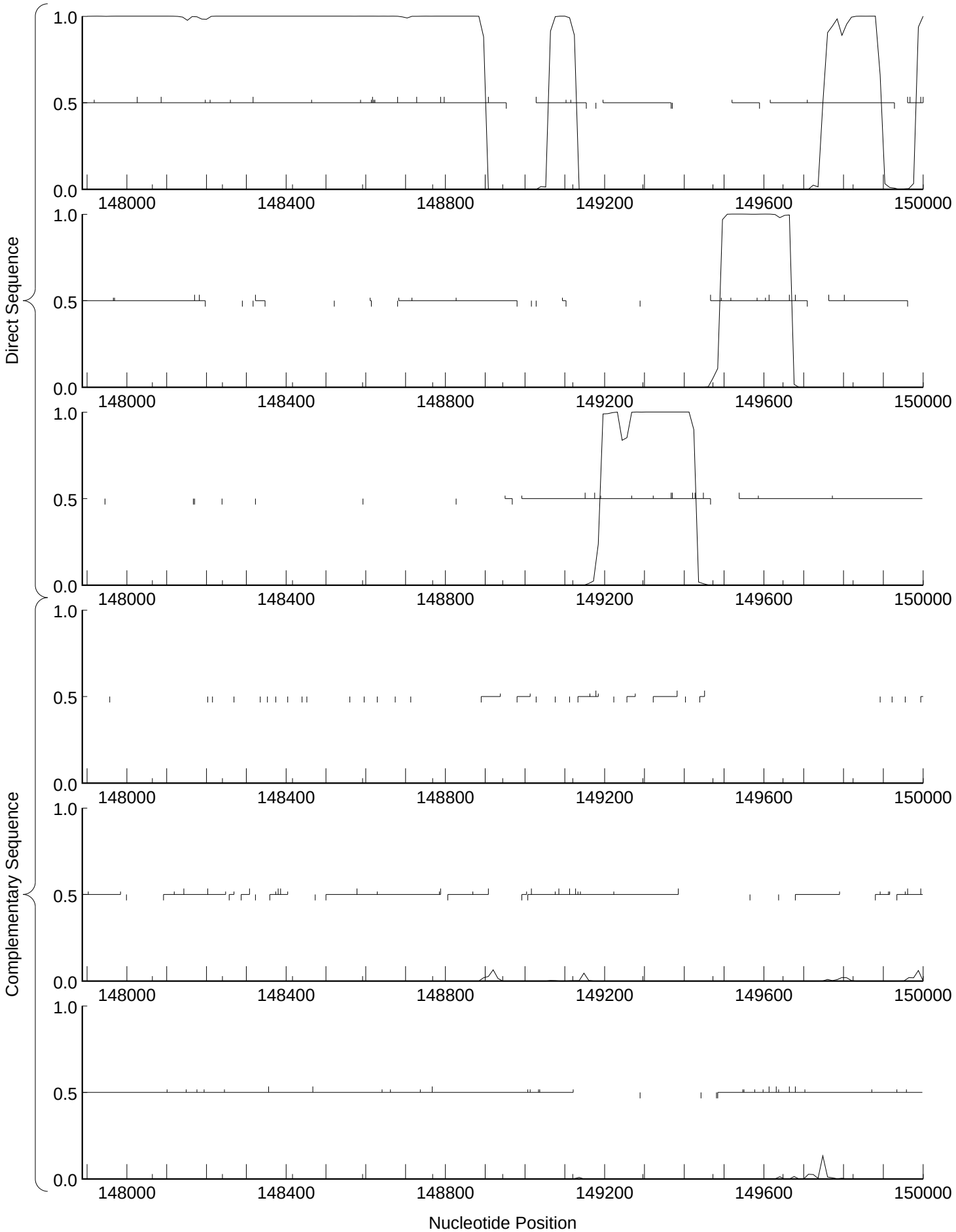
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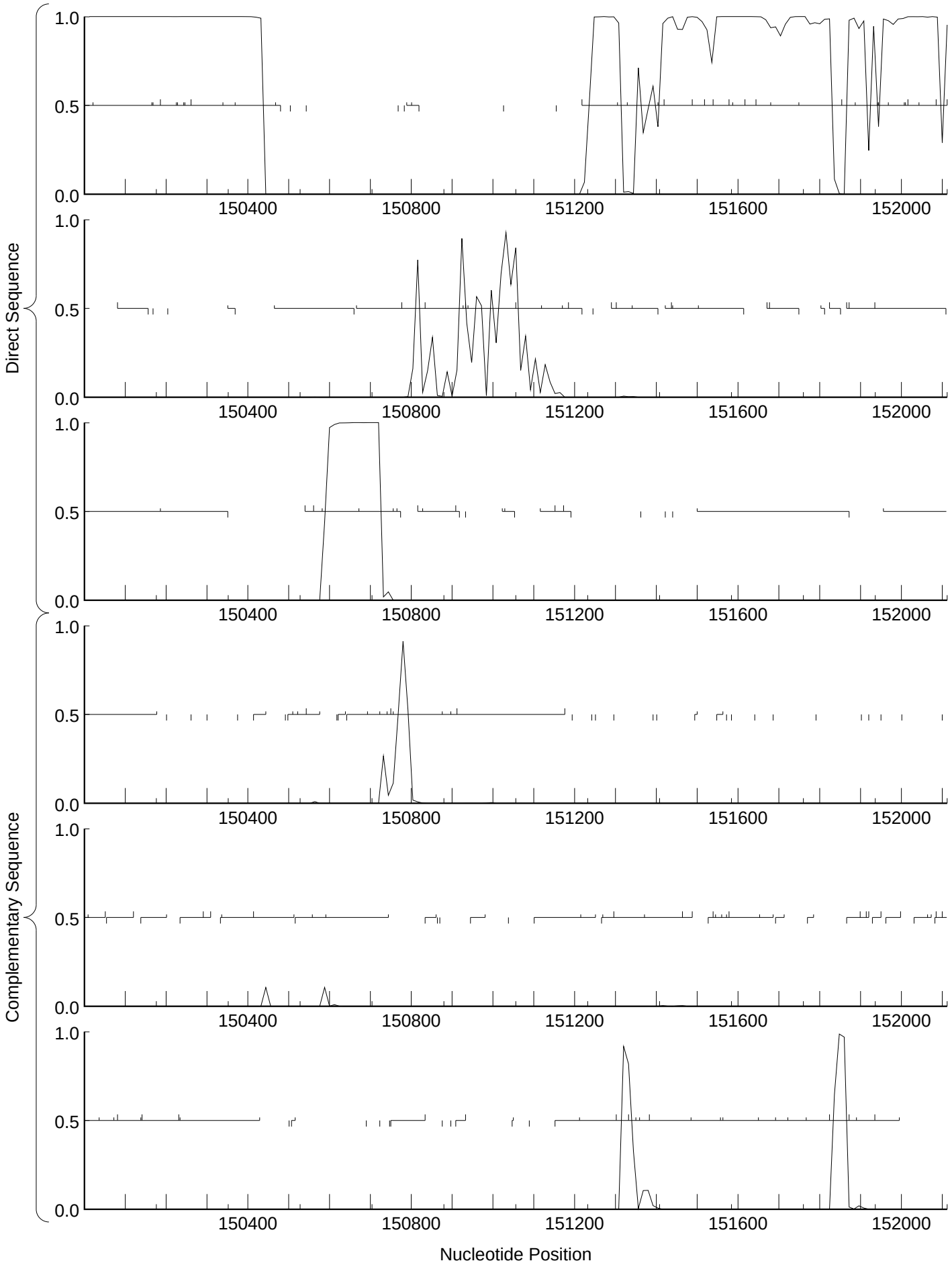
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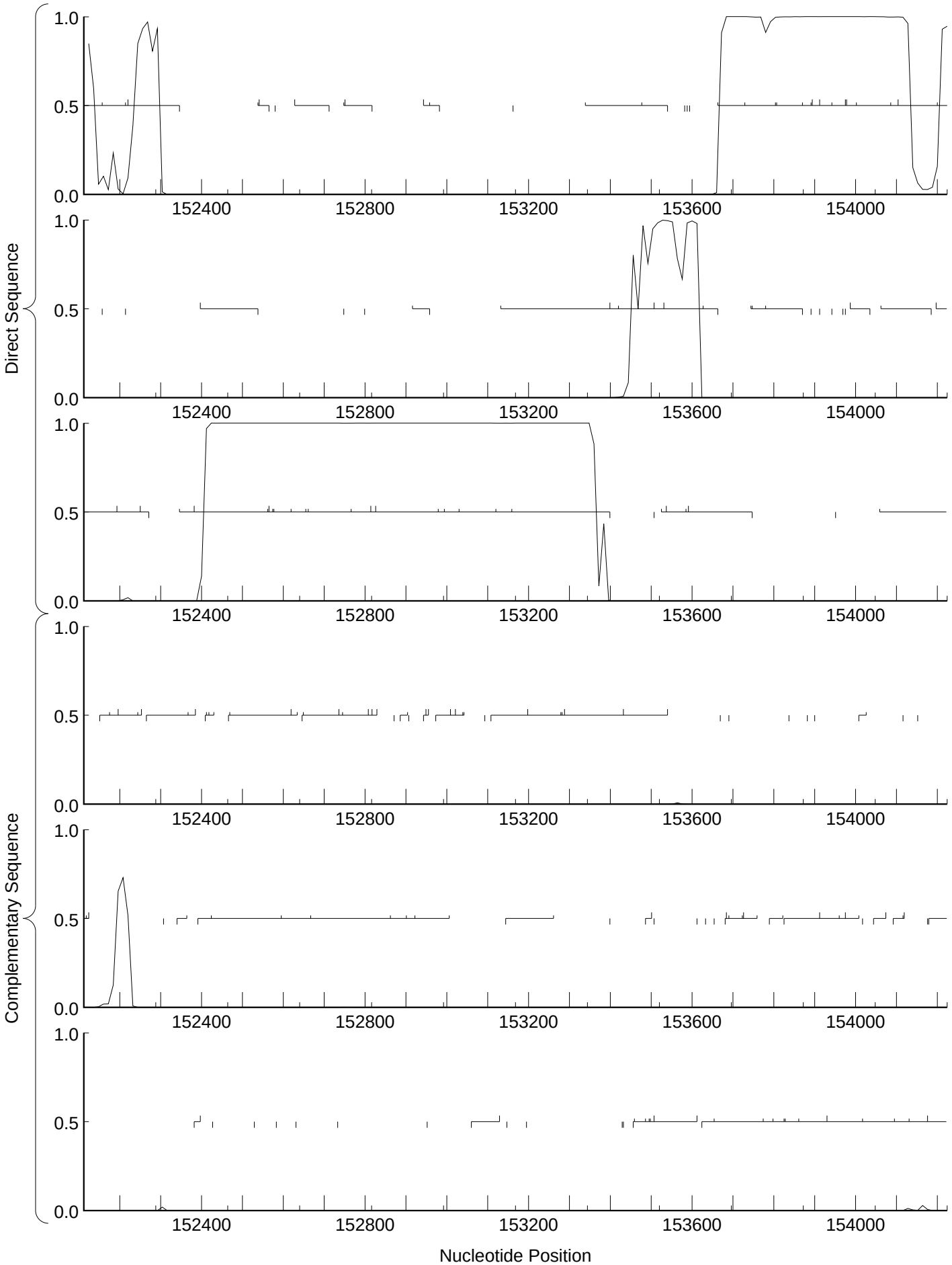
erium phage NuevoMundo Complete Sequence, 155943 bp, Circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 72/75



erium phage NuevoMundo Complete Sequence, 155943 bp, Circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 73/75



erium phage NuevoMundo Complete Sequence, 155943 bp, Circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 74/75



erium phage NuevoMundo Complete Sequence, 155943 bp, Circularly permuted, Cluster C1, Order 4, Window 96, Step 12, 75/75

