GeneMark

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

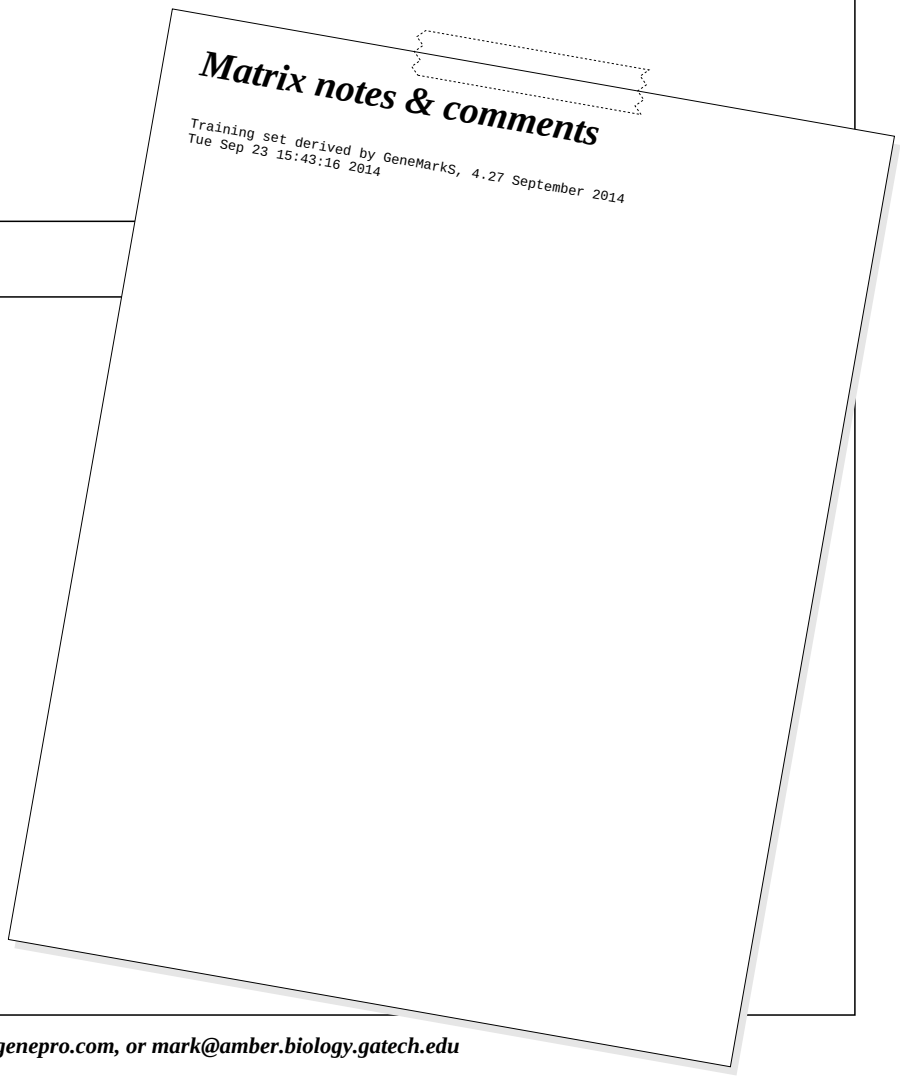
PROGRAM INFORMATION

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Analysis Date    :  3/29/24 at 3:09:45
Pages            :  34
Sequence Length  :  69299 bp
GC Content       :   64.24%

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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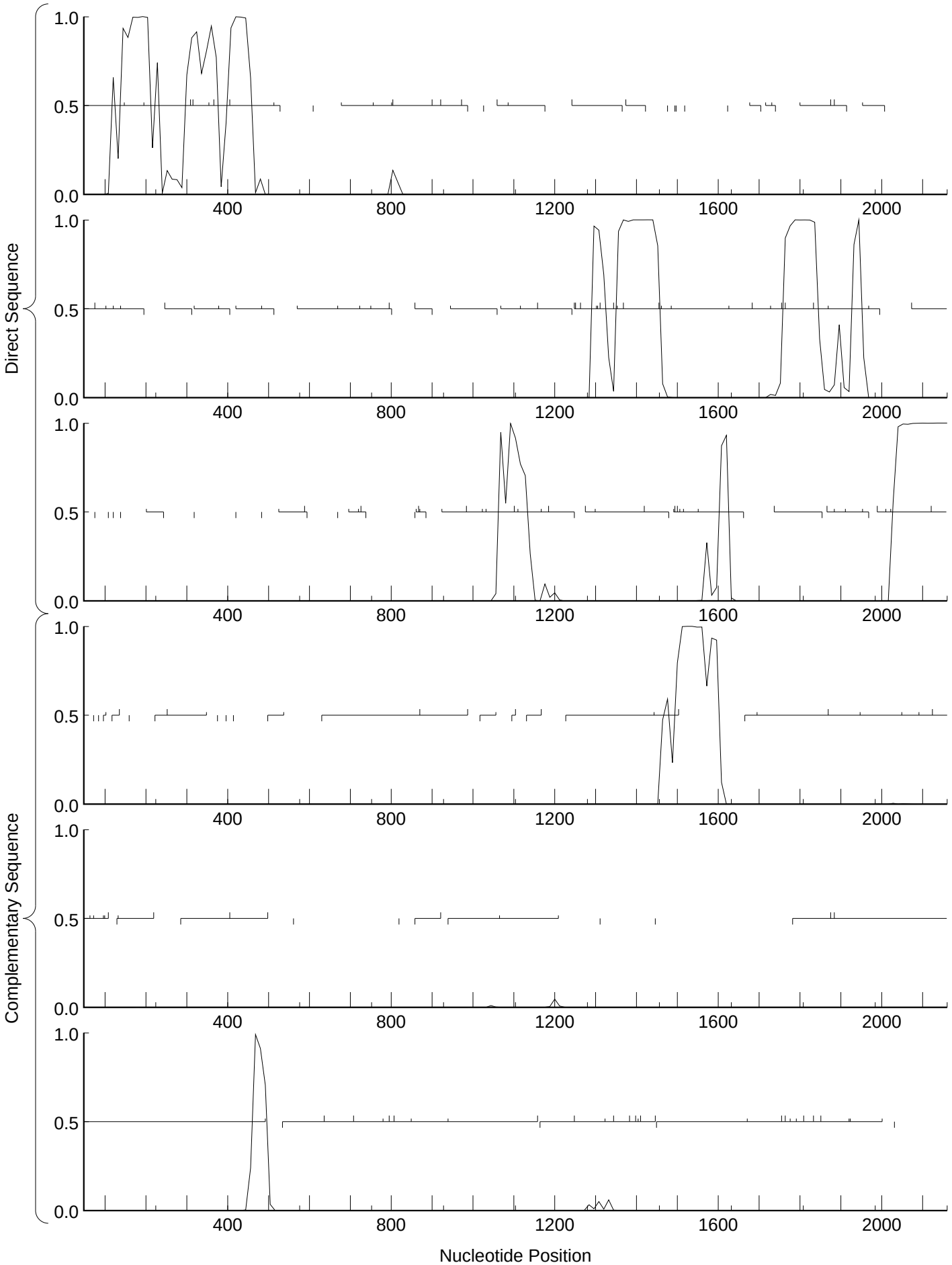
MATRIX INFORMATION

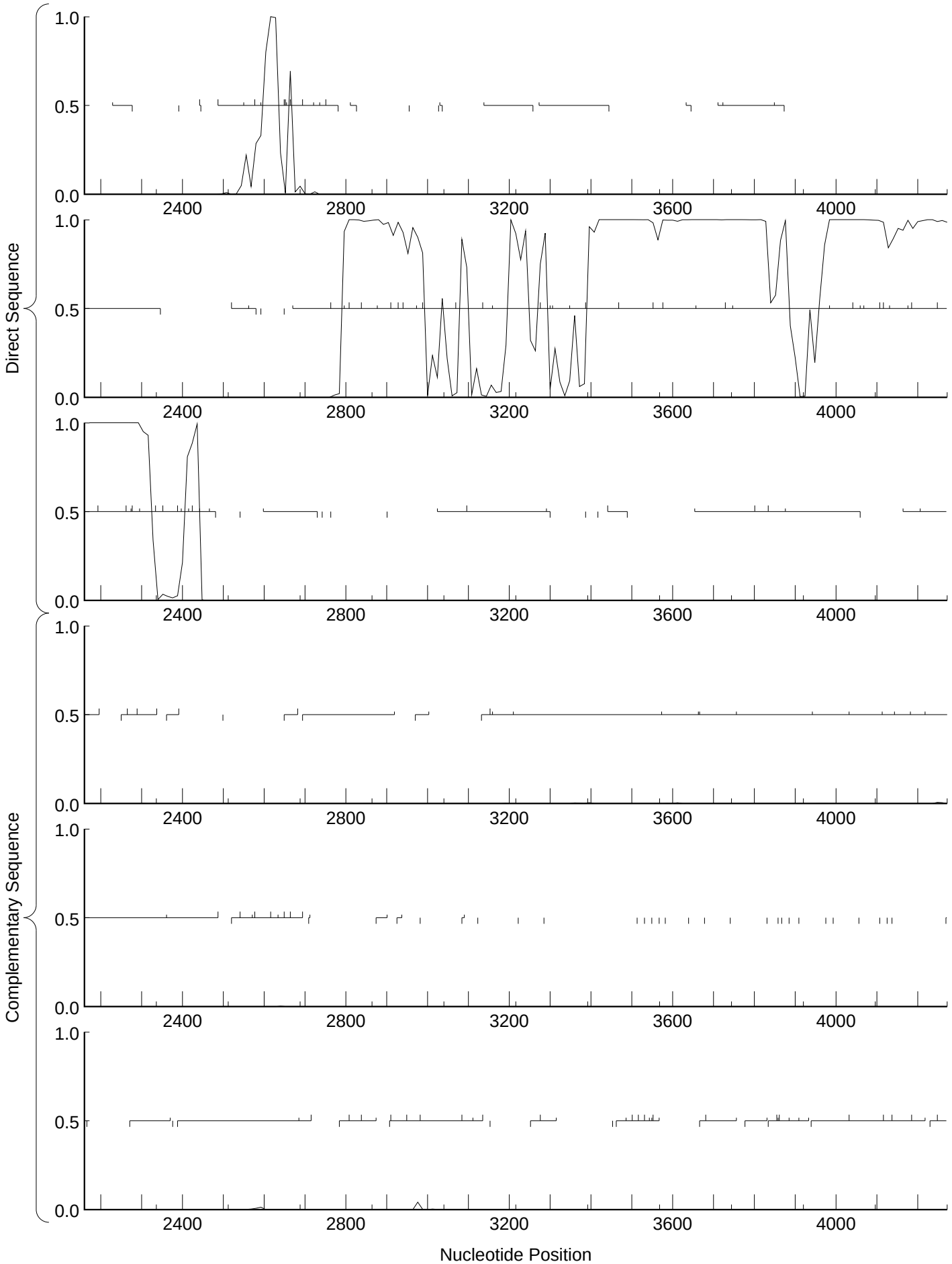
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Matrix     :  Streptomyces_scabiei_87_22
Author     :  -

Order      :  4
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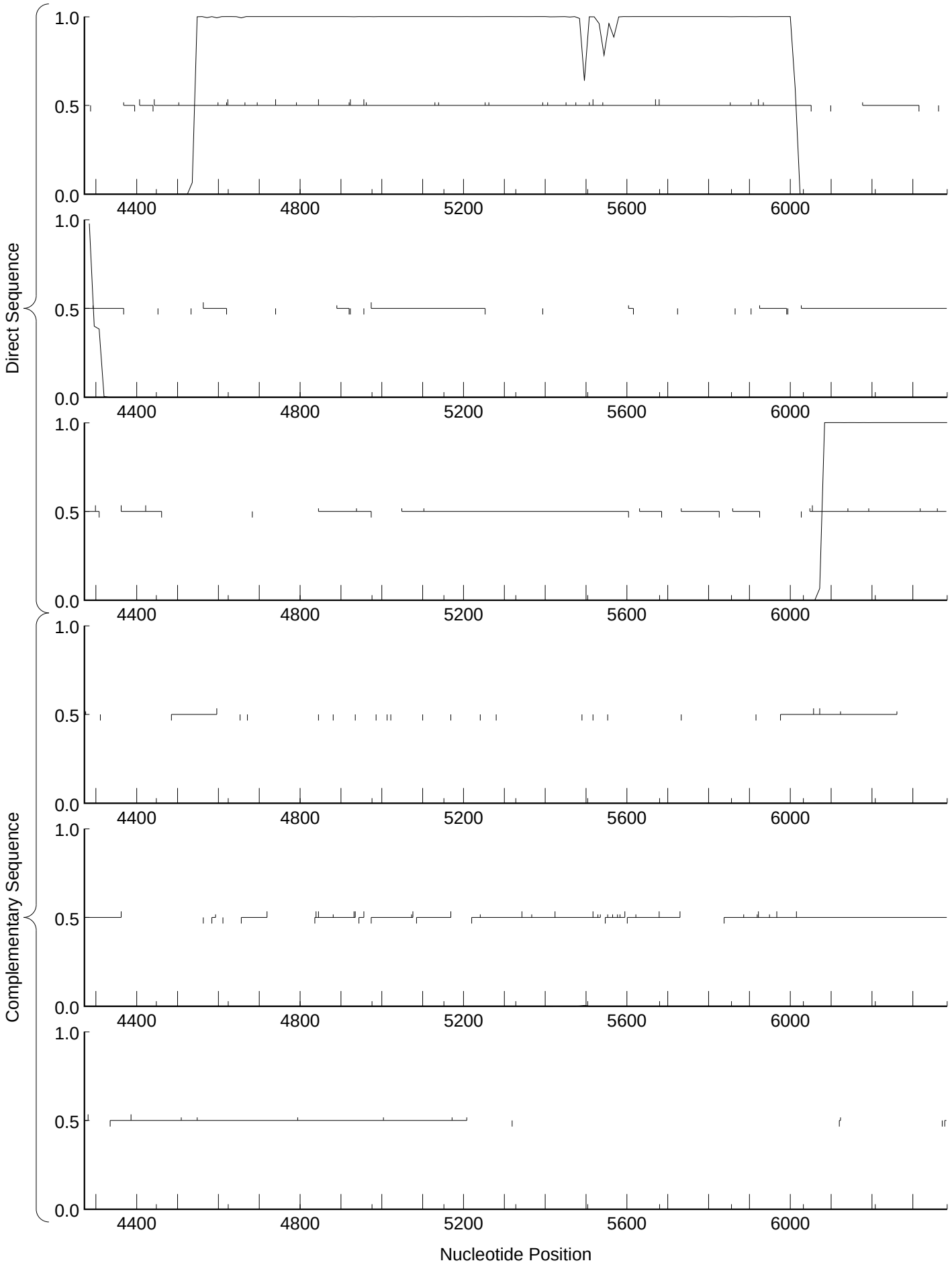

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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
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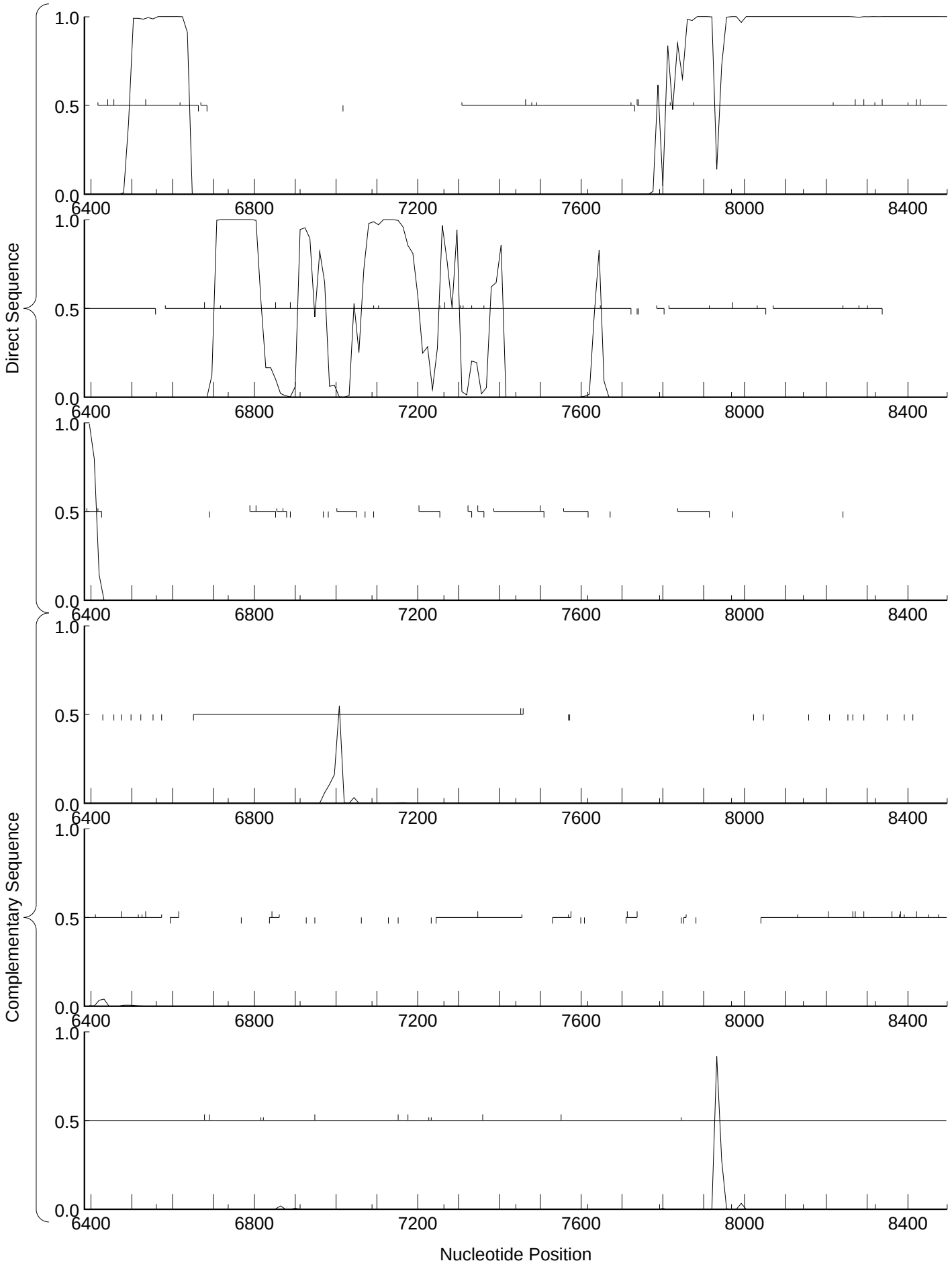

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

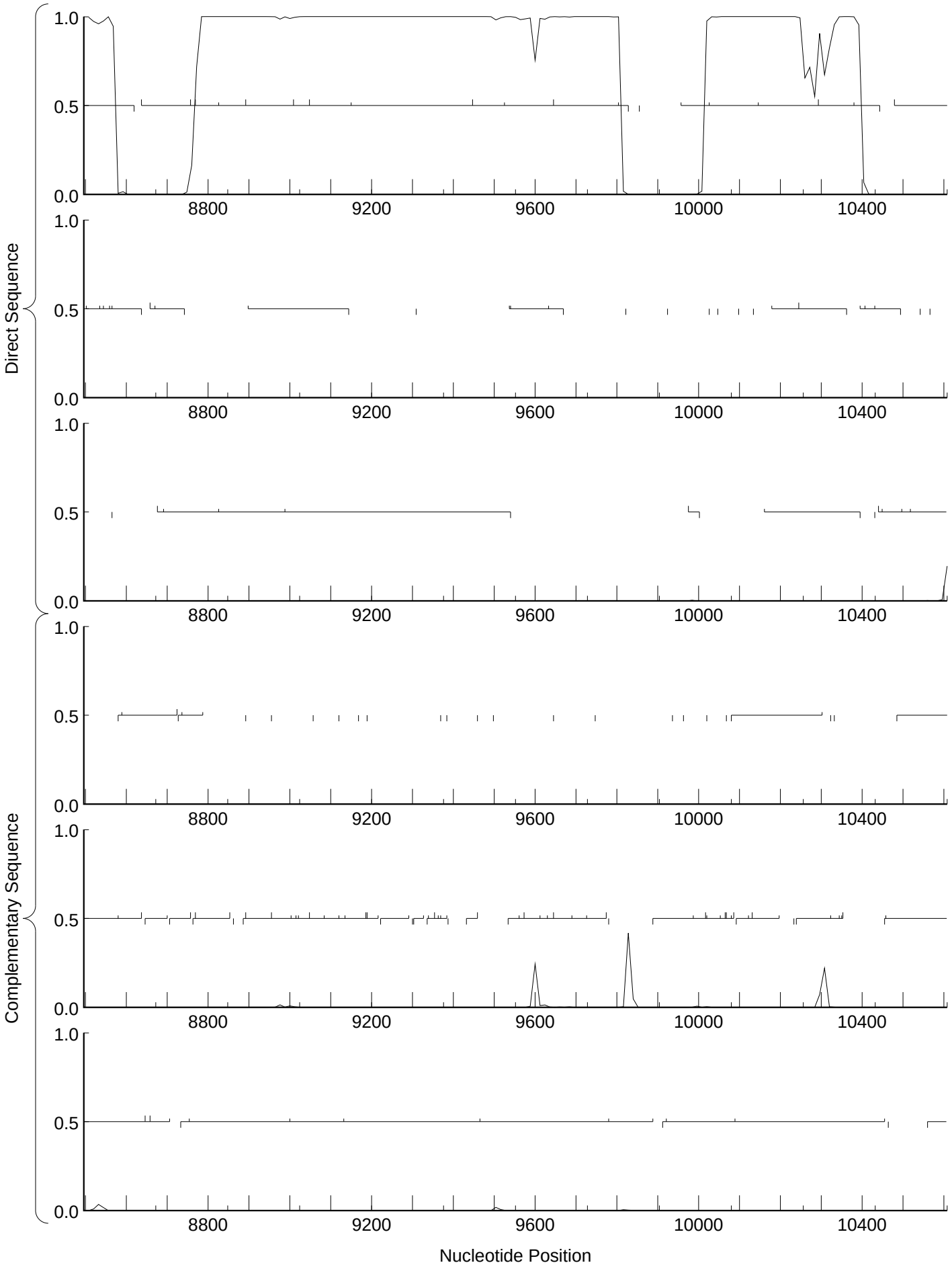

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
400
800
1200
1600
2000
Nucleotide Position

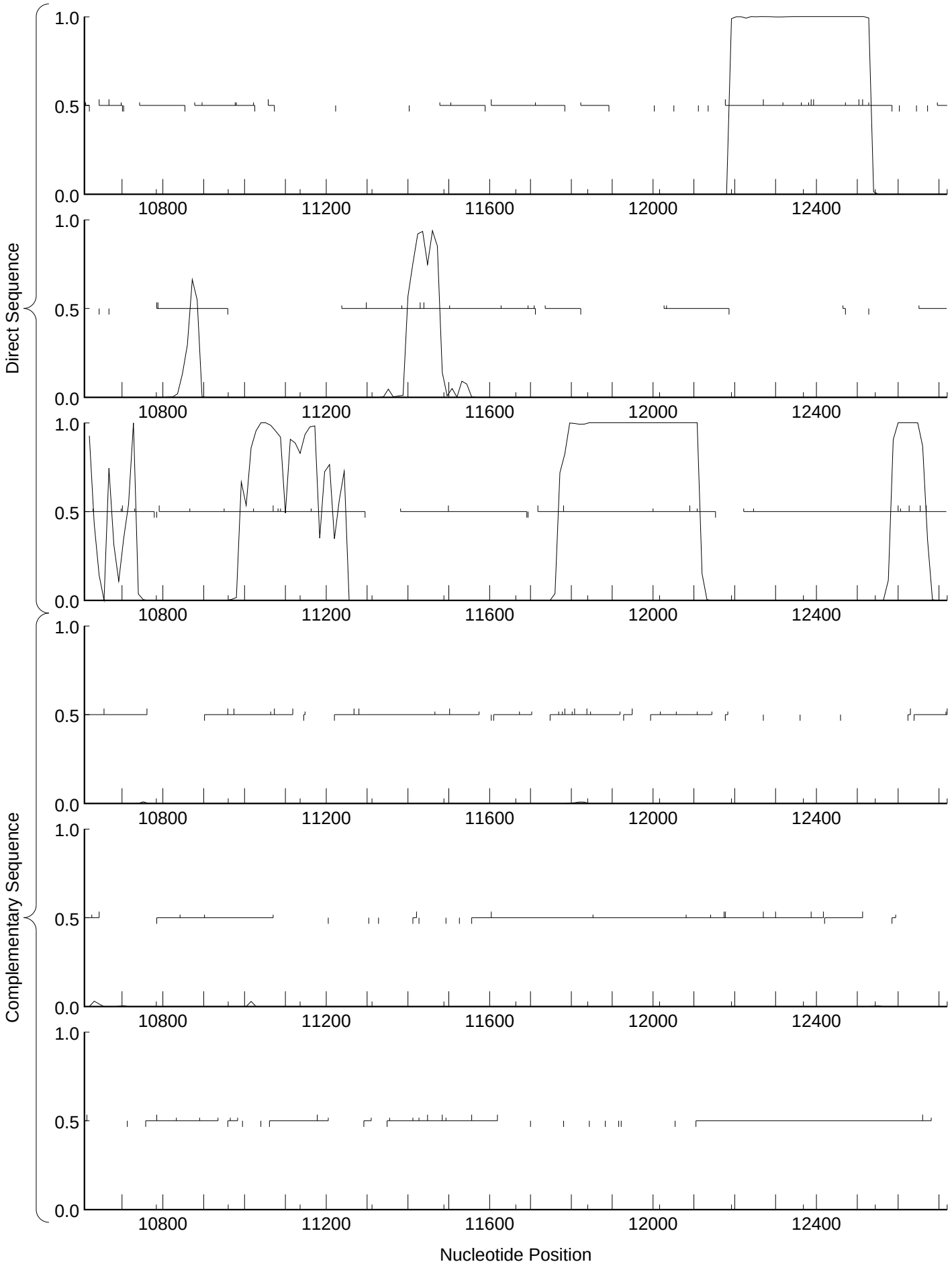

Direct Sequence
Complementary Sequence
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3200
3600
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Nucleotide Position

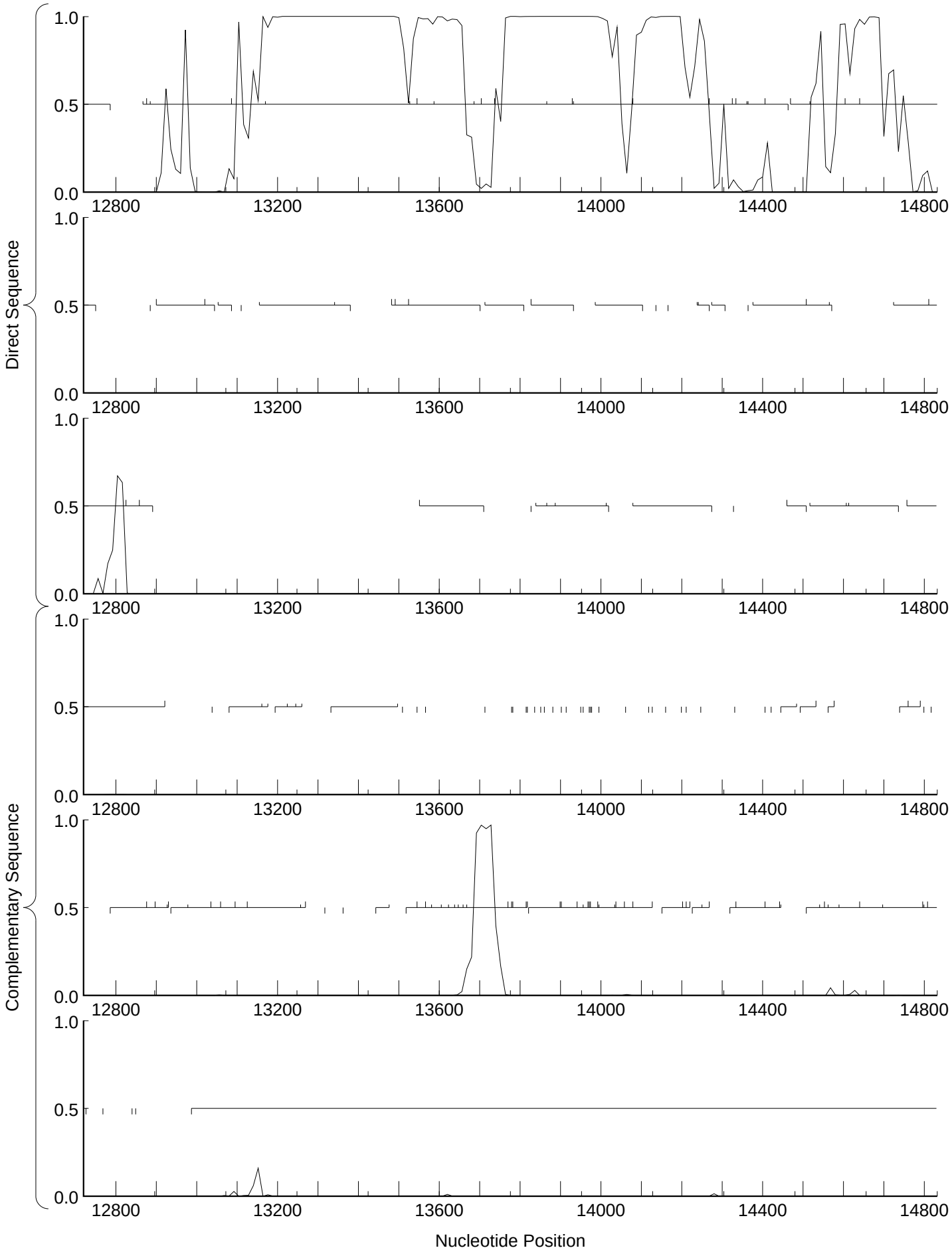
ptomyces phage ShakeNBake complete sequence, 69299 bp, circularly permuted, Cluster BN, Order 4, Window 96, Step 12, 4/34



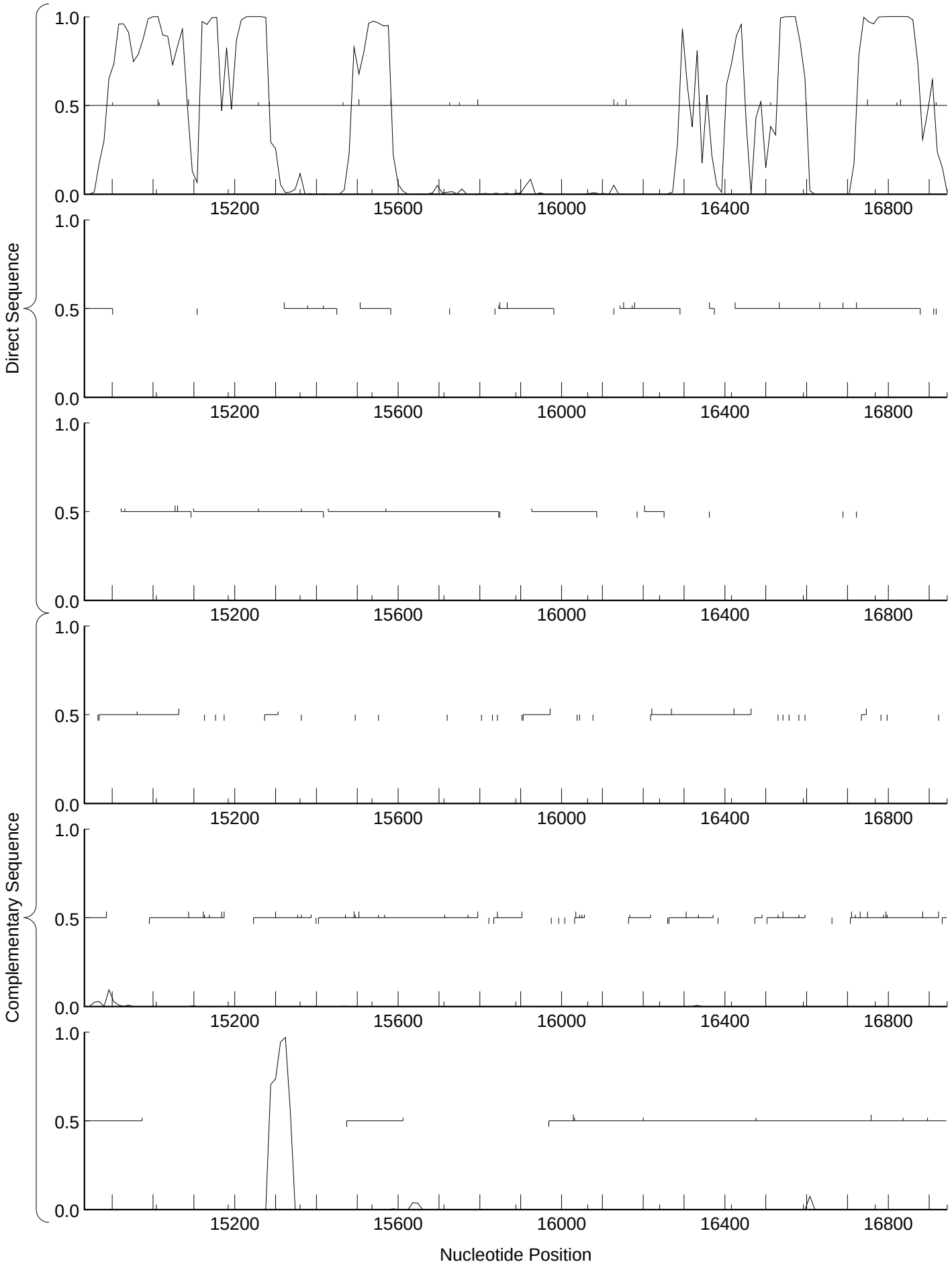

Direct Sequence
Complementary Sequence
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0.5
0.0
6400
6800
7200
7600
8000
8400
Nucleotide Position


Direct Sequence
Complementary Sequence
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0.5
0.0
8800
9200
9600
10000
10400
Nucleotide Position

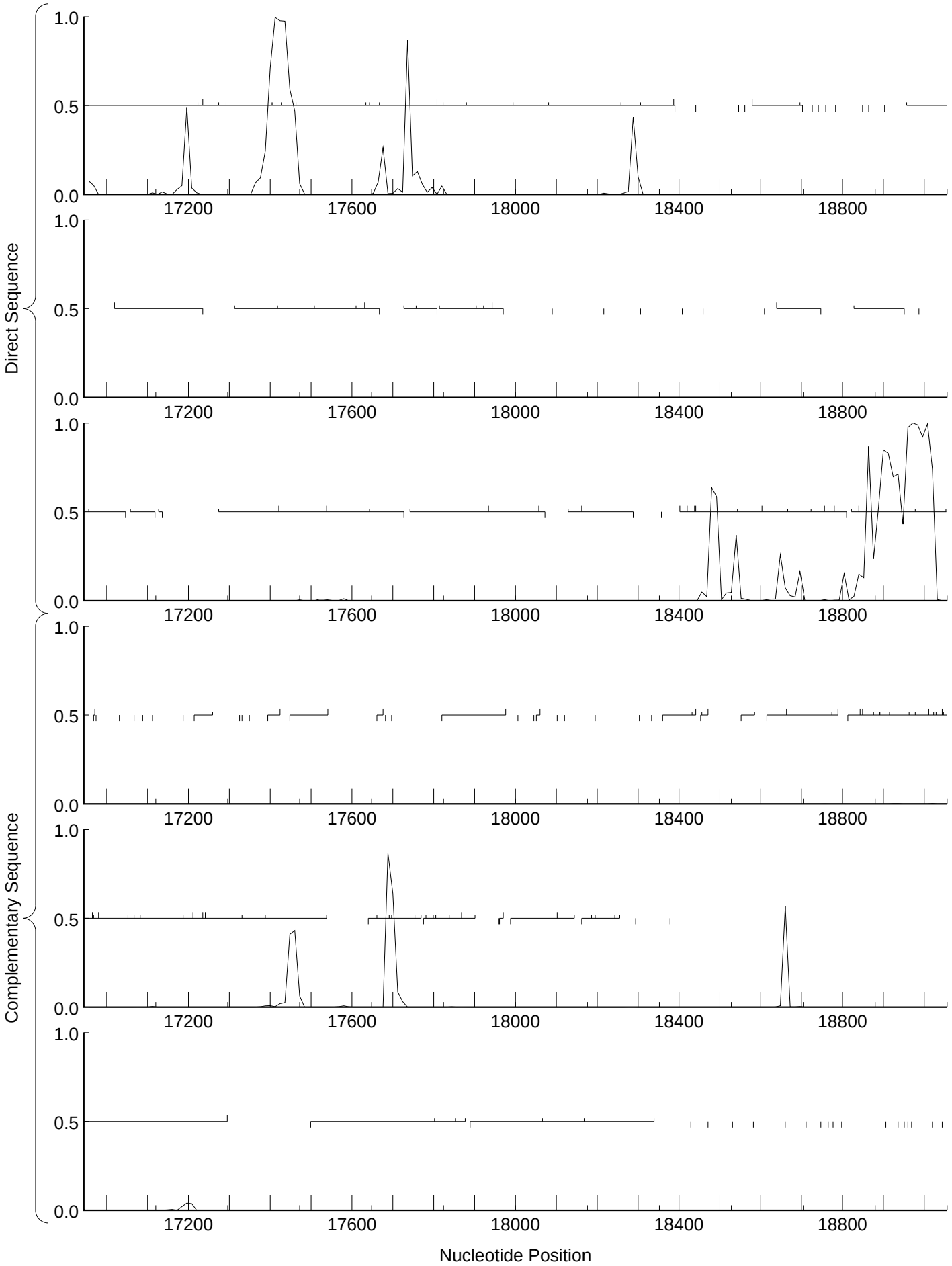

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
10800
11200
11600
12000
12400
Nucleotide Position


Direct Sequence
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12800
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14400
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Nucleotide Position

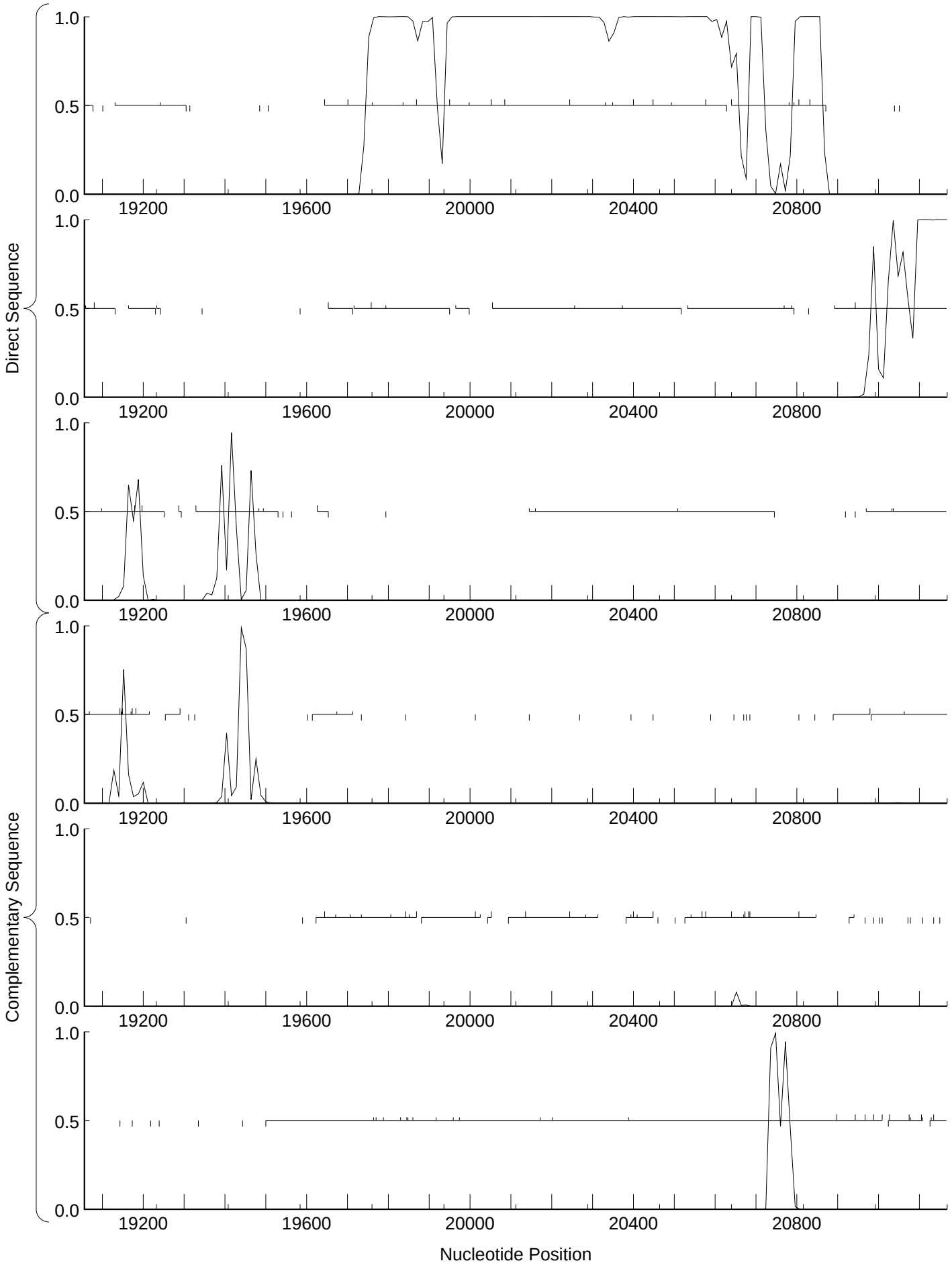
ptomyces phage ShakeNBake complete sequence, 69299 bp, circularly permuted, Cluster BN, Order 4, Window 96, Step 12, 9/34



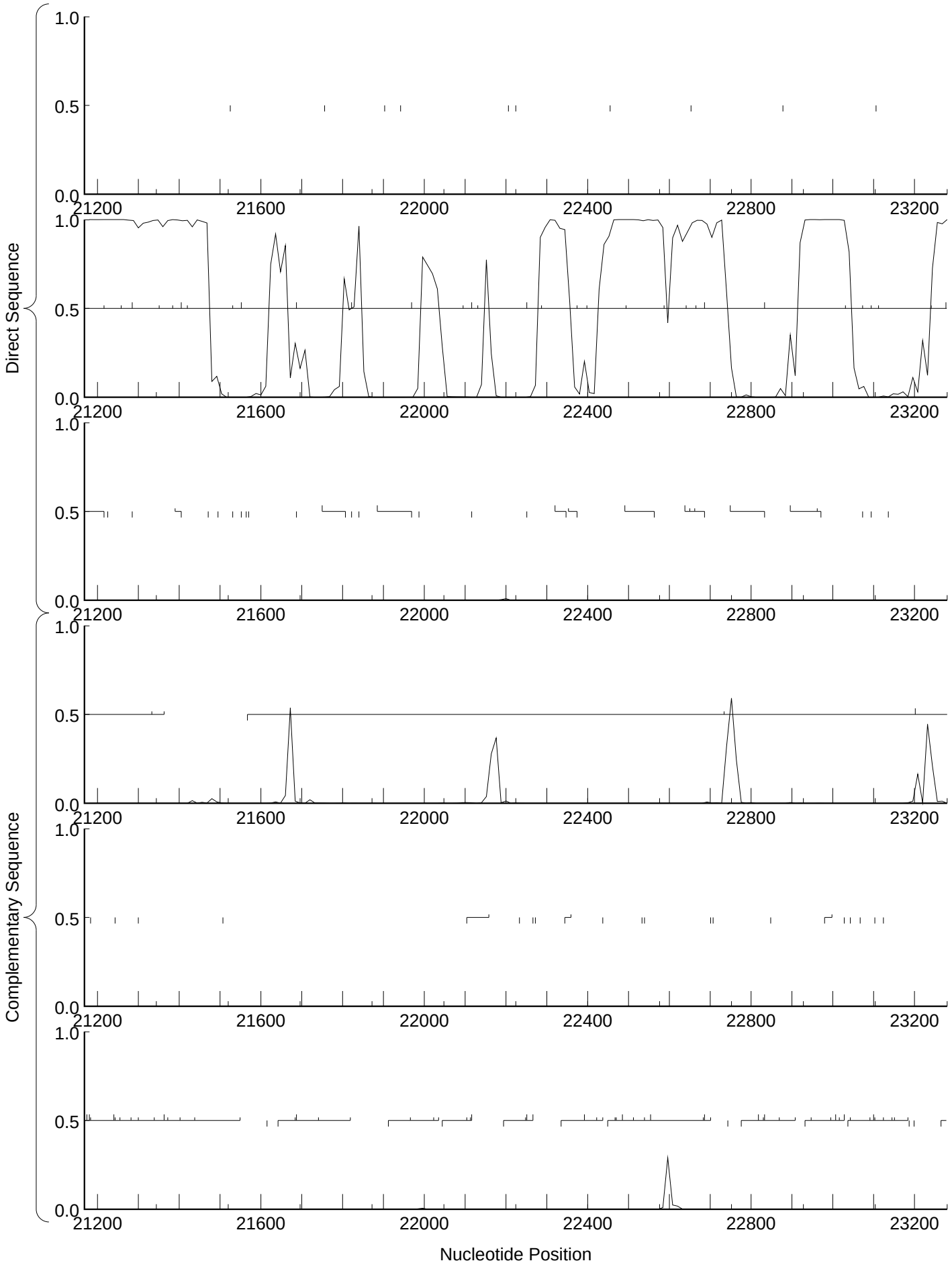
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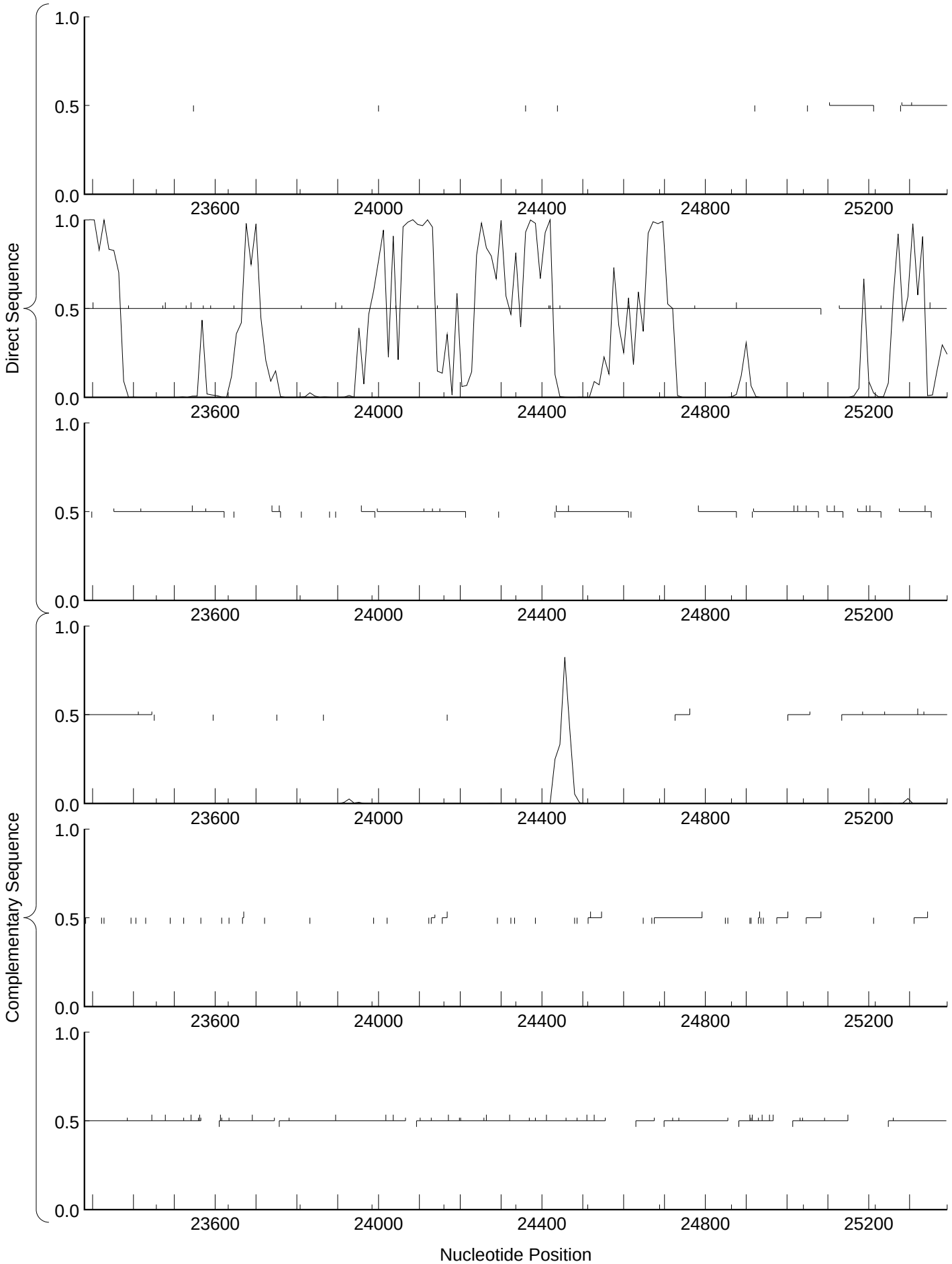
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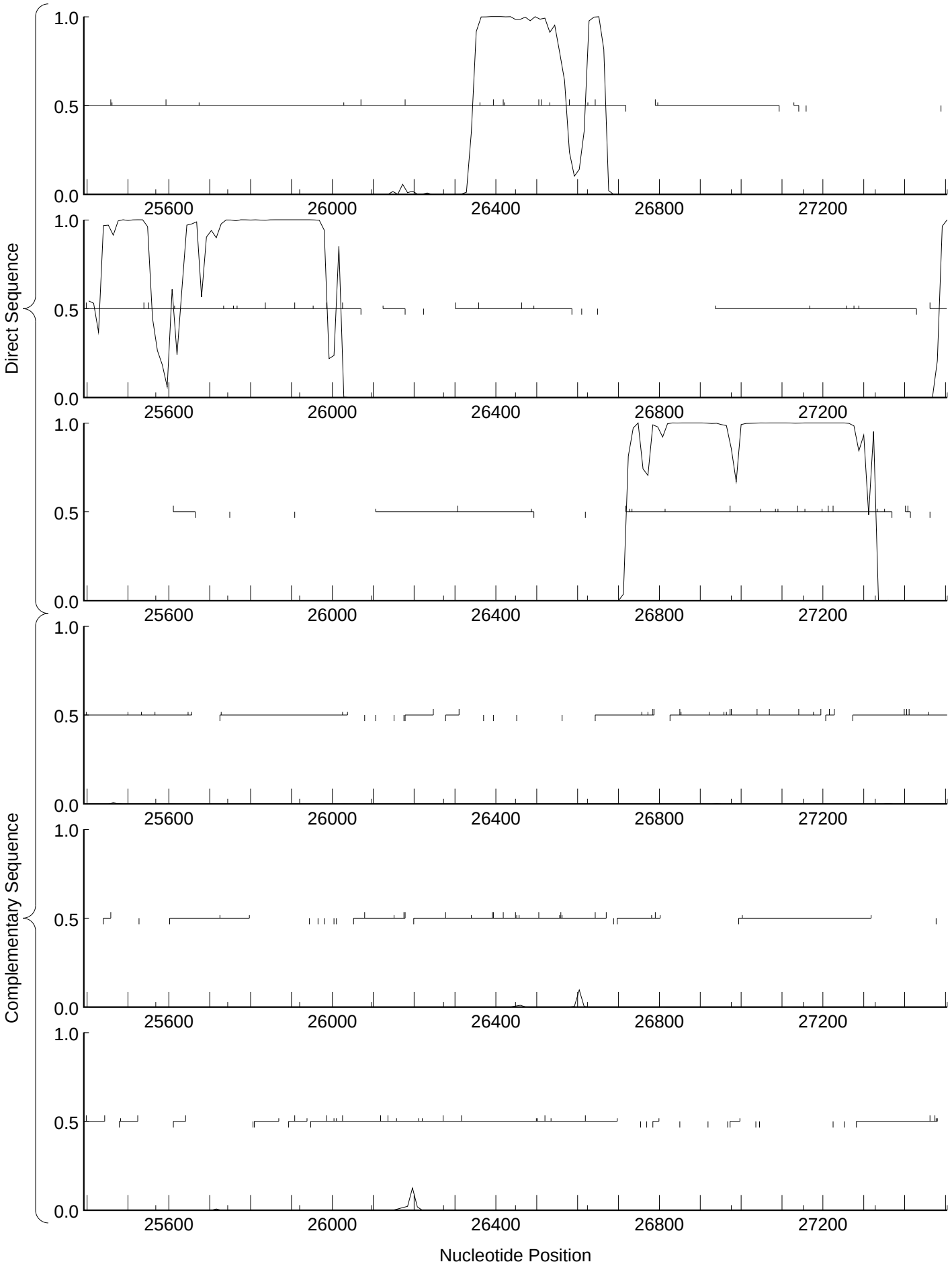
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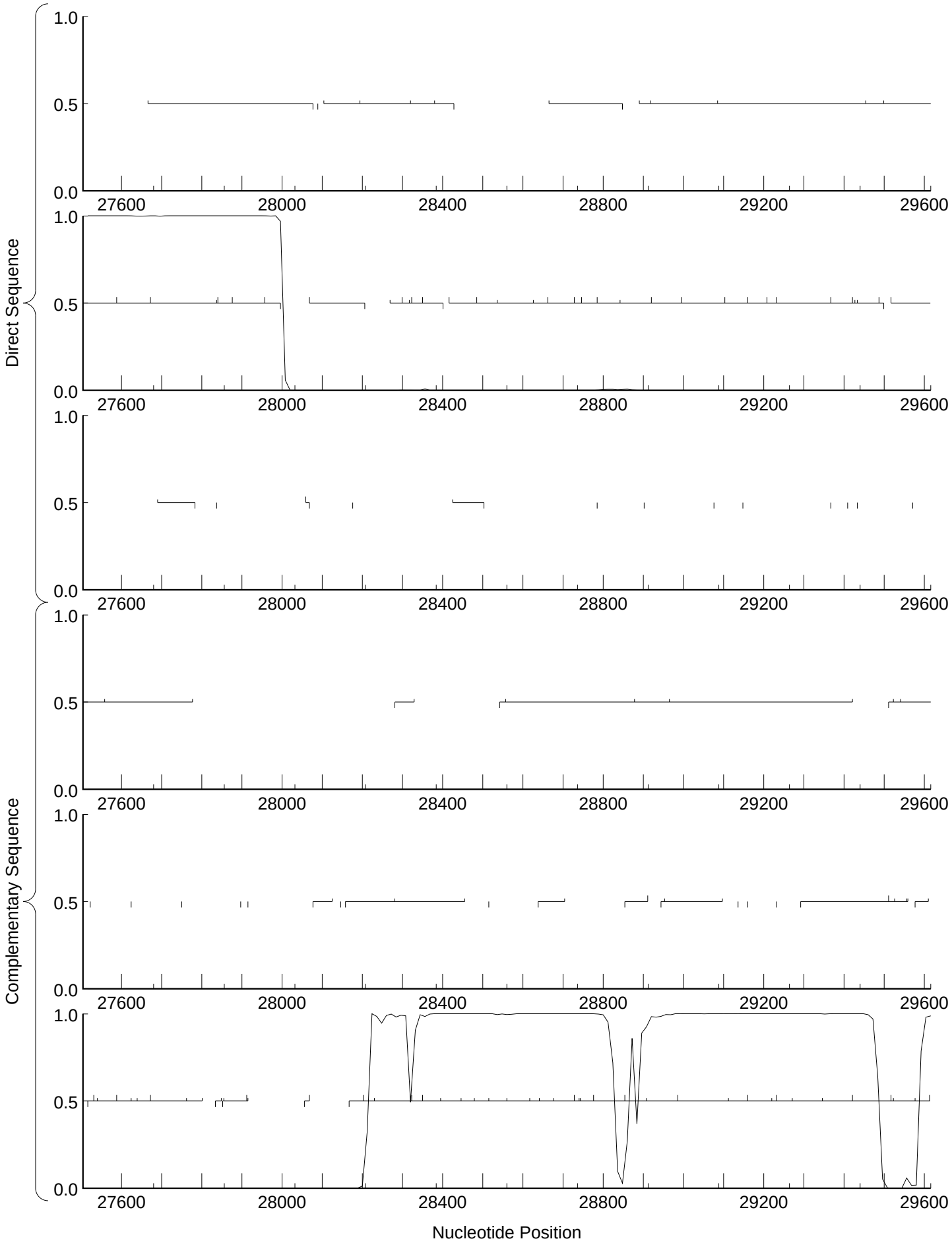
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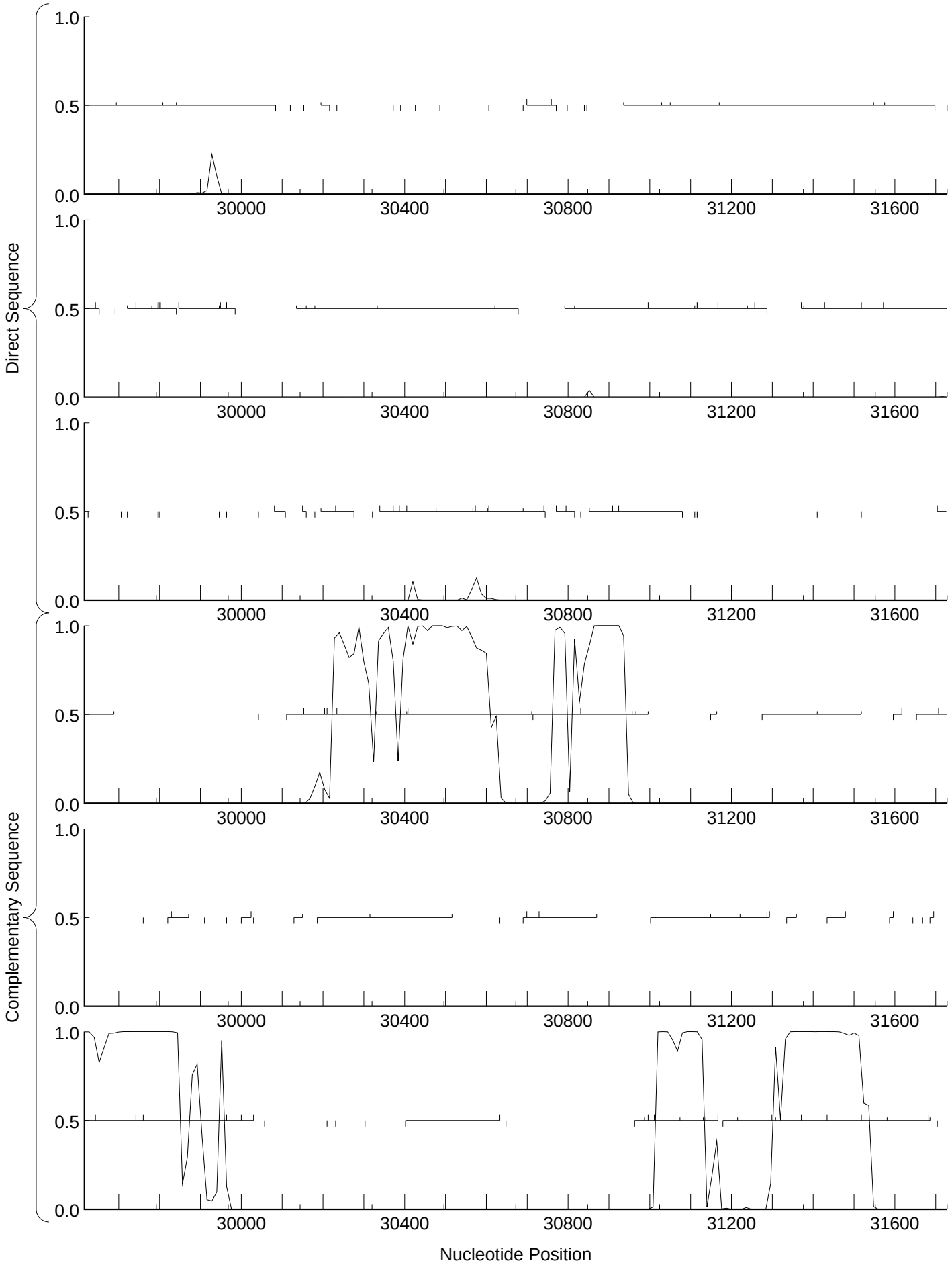
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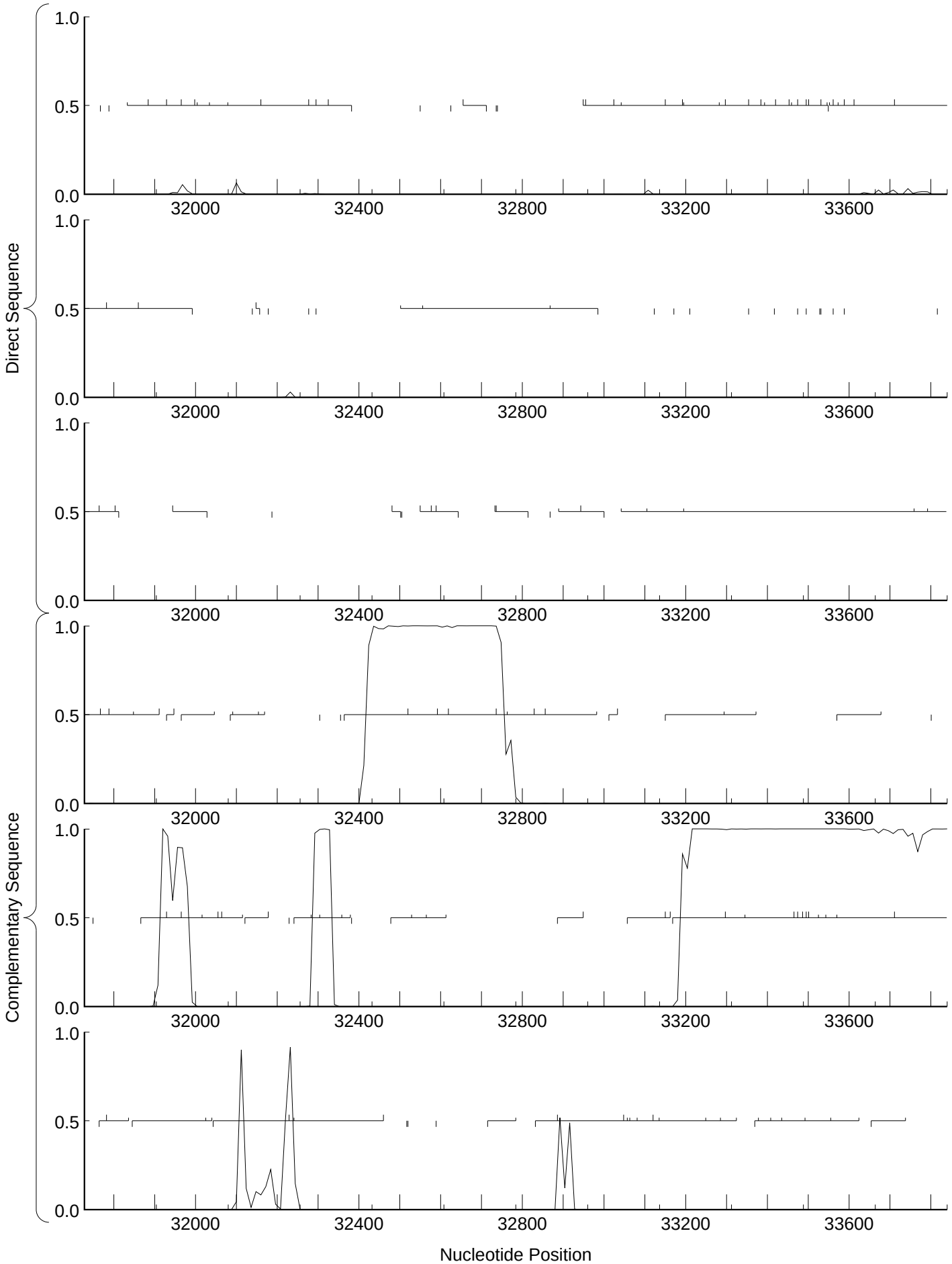
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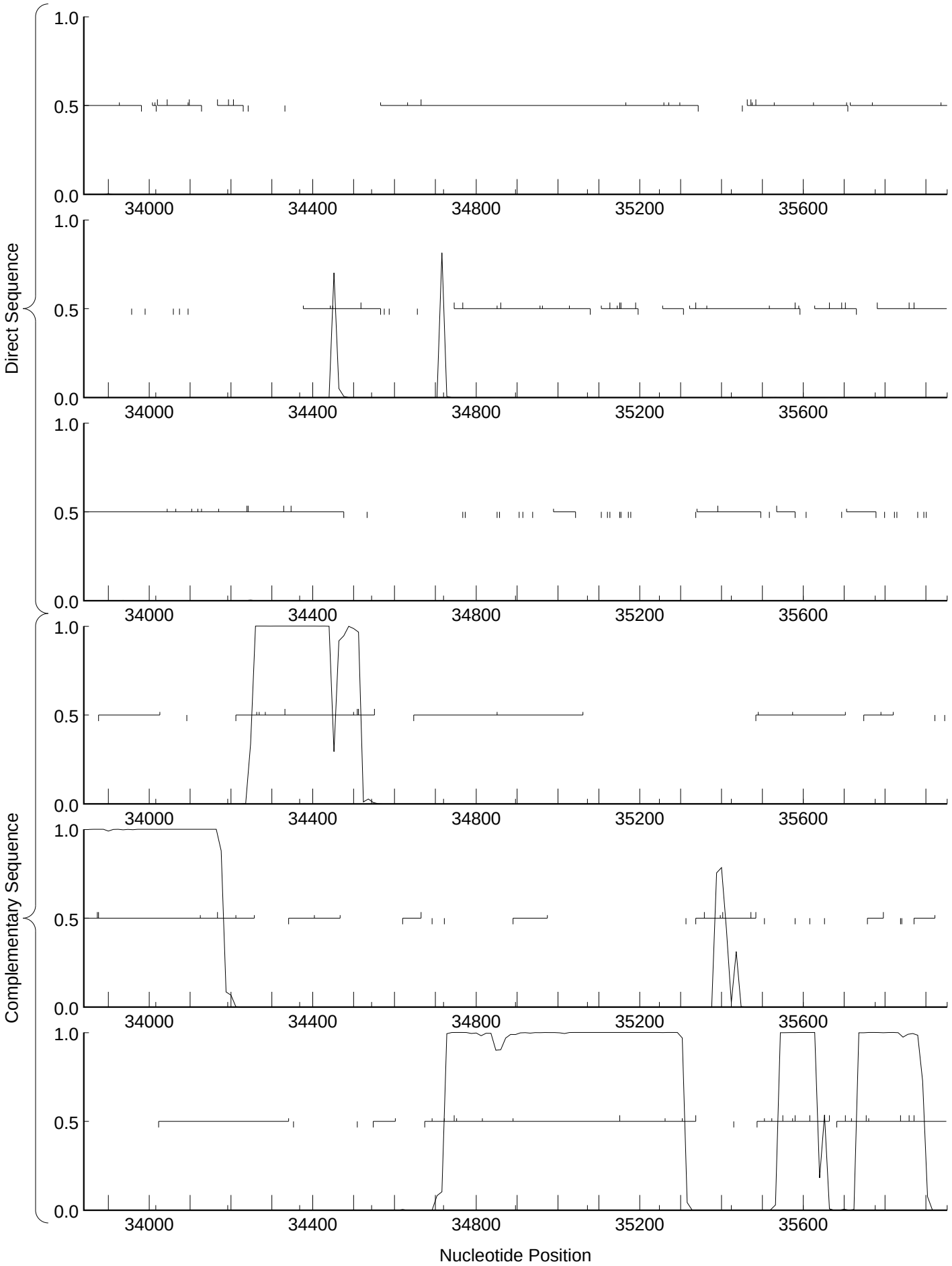
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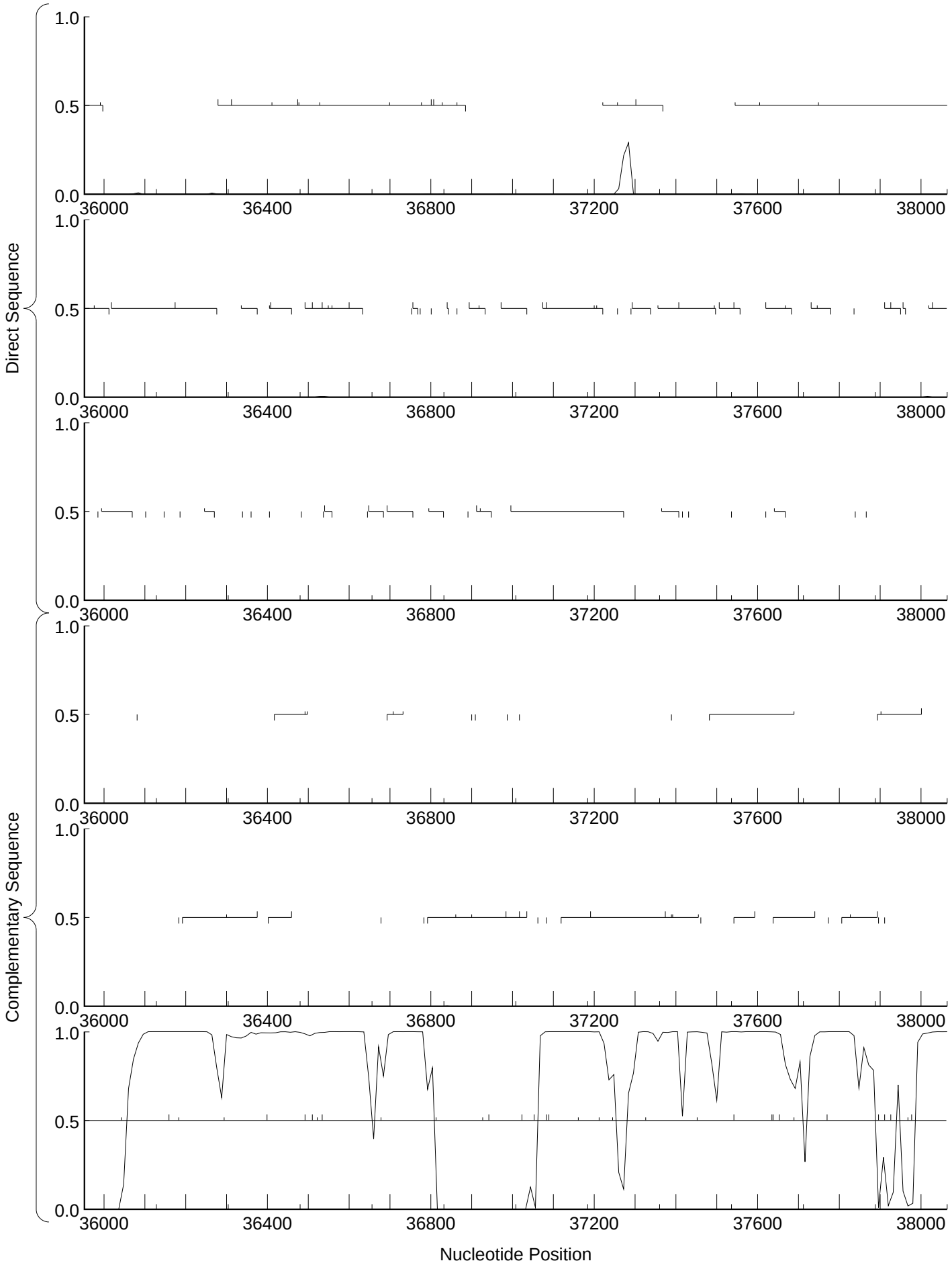
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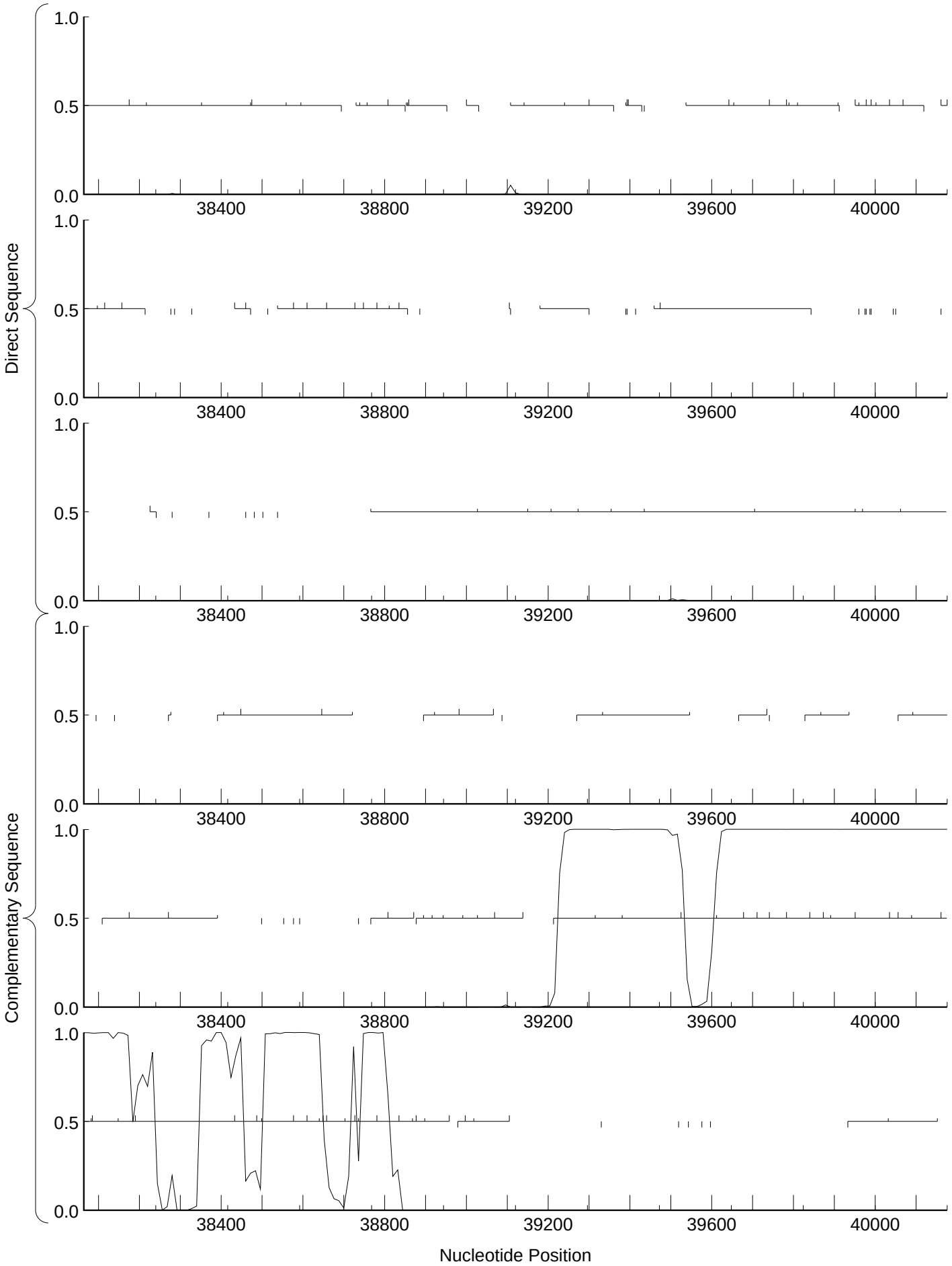
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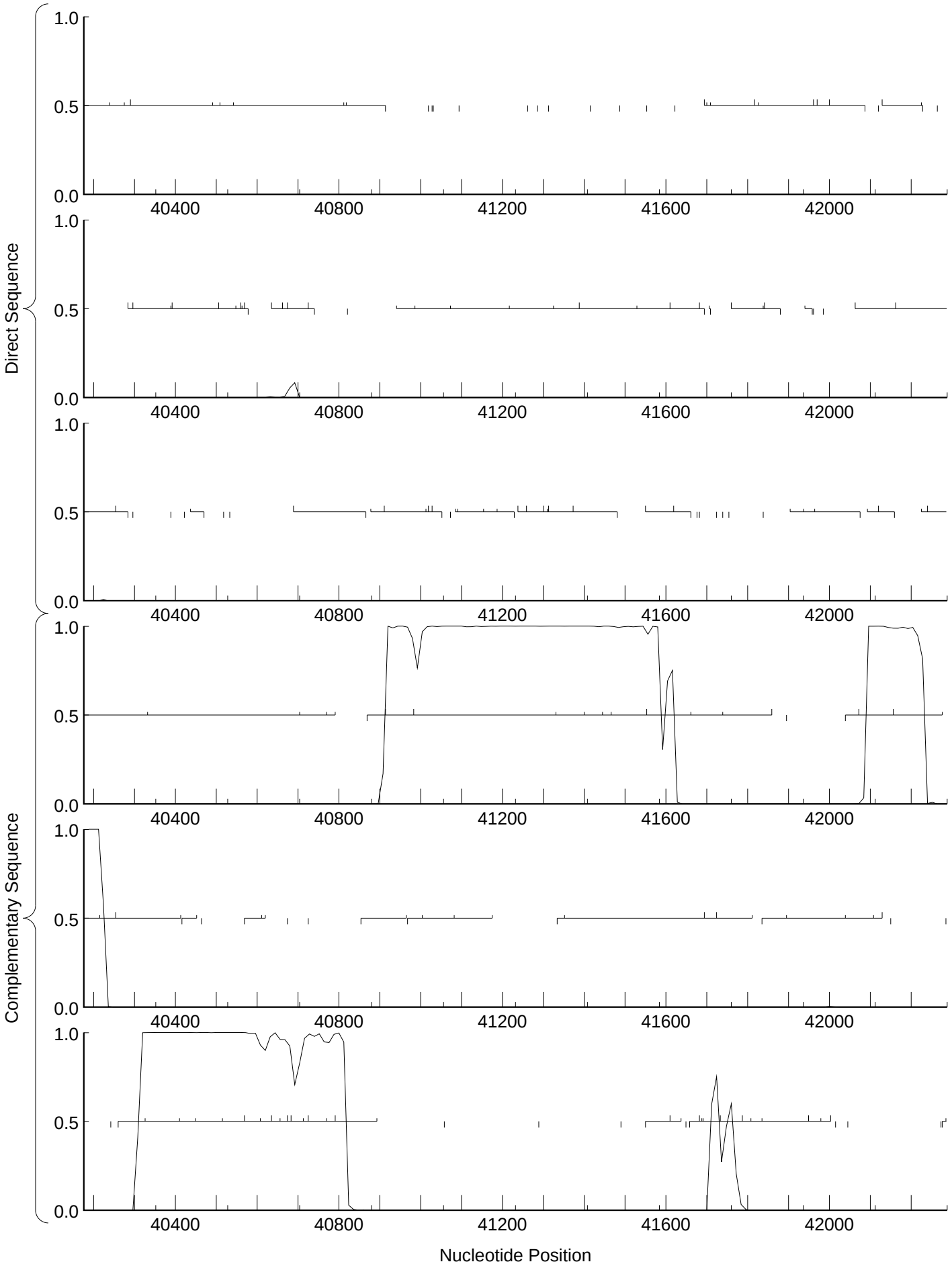
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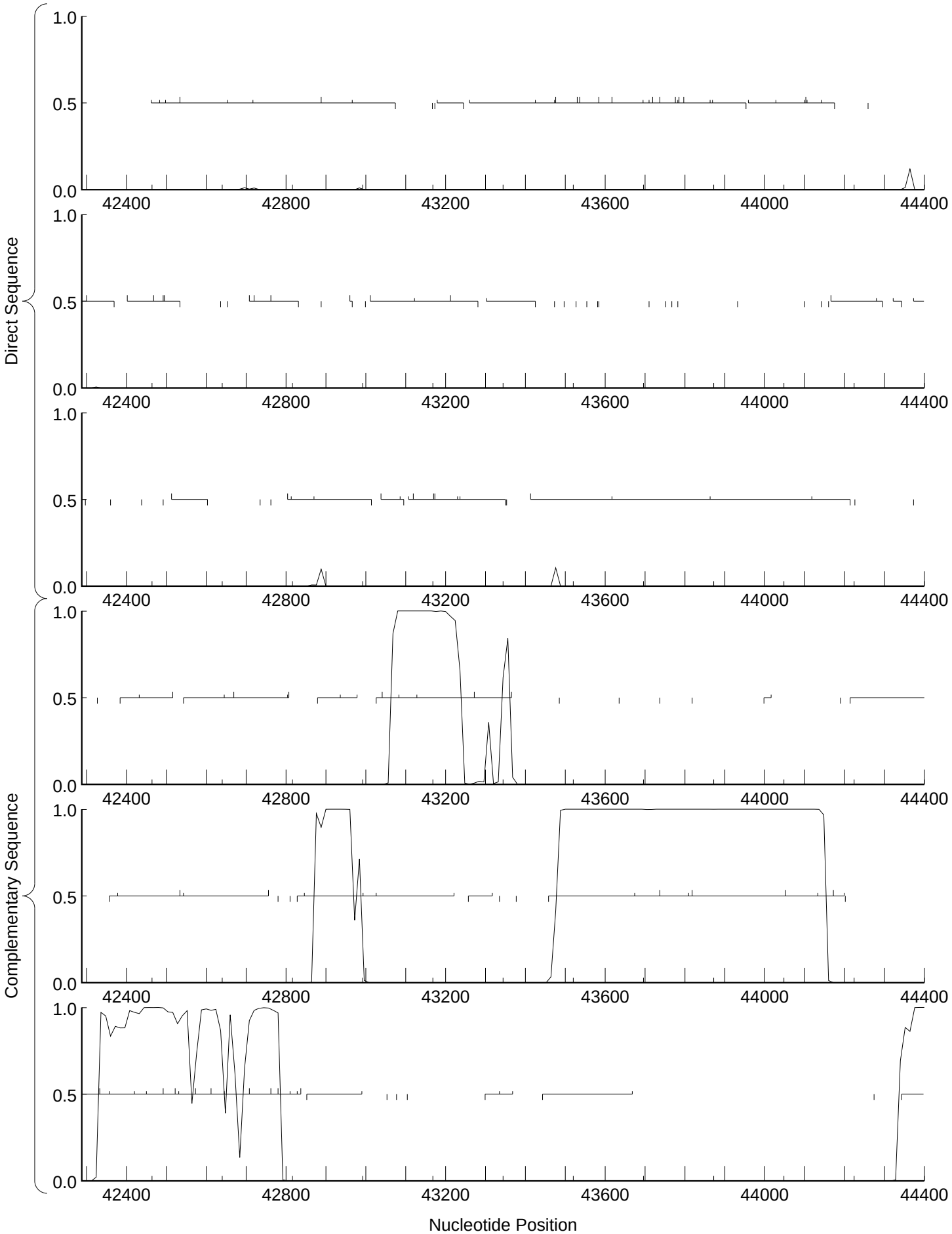
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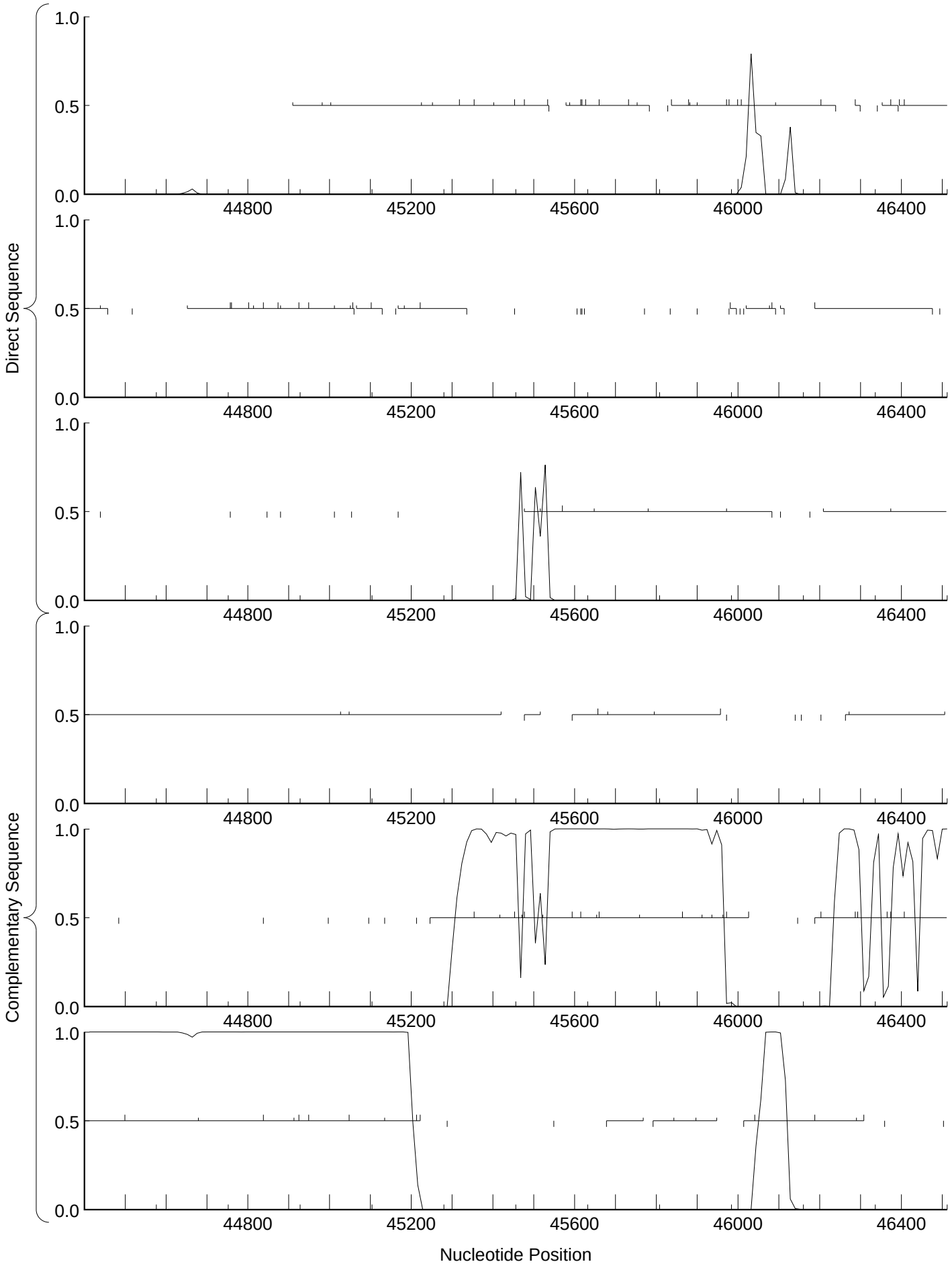
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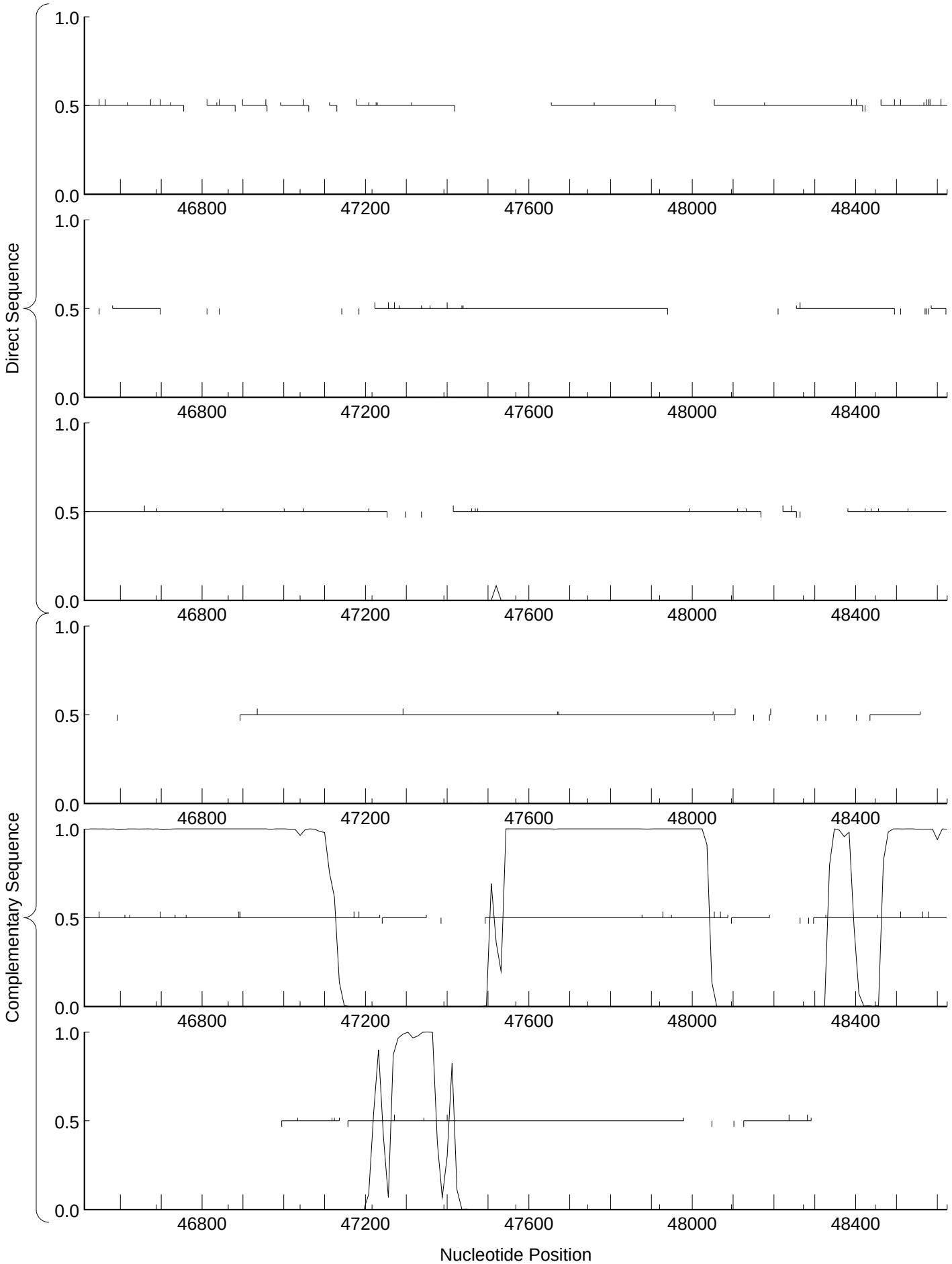
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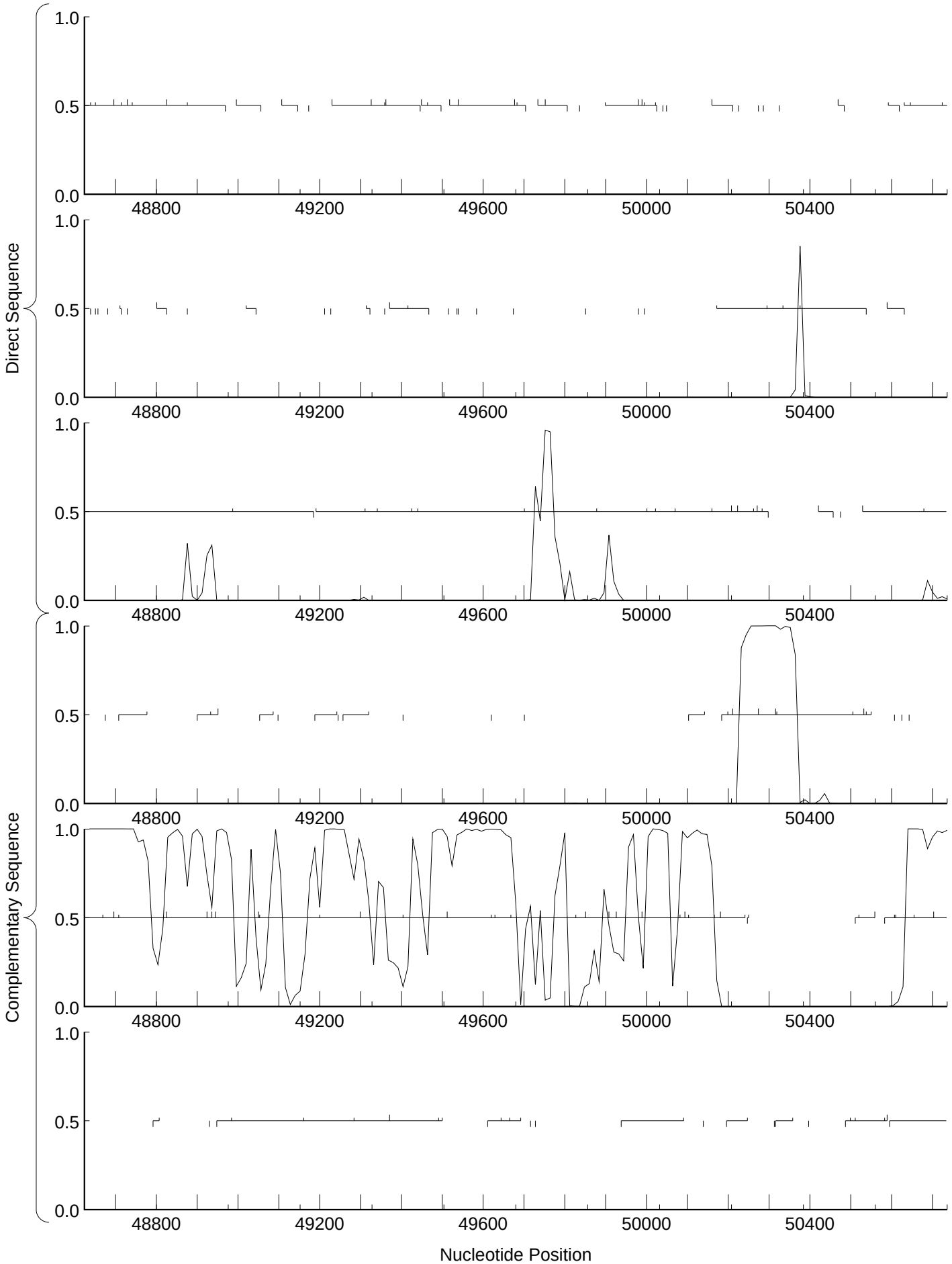
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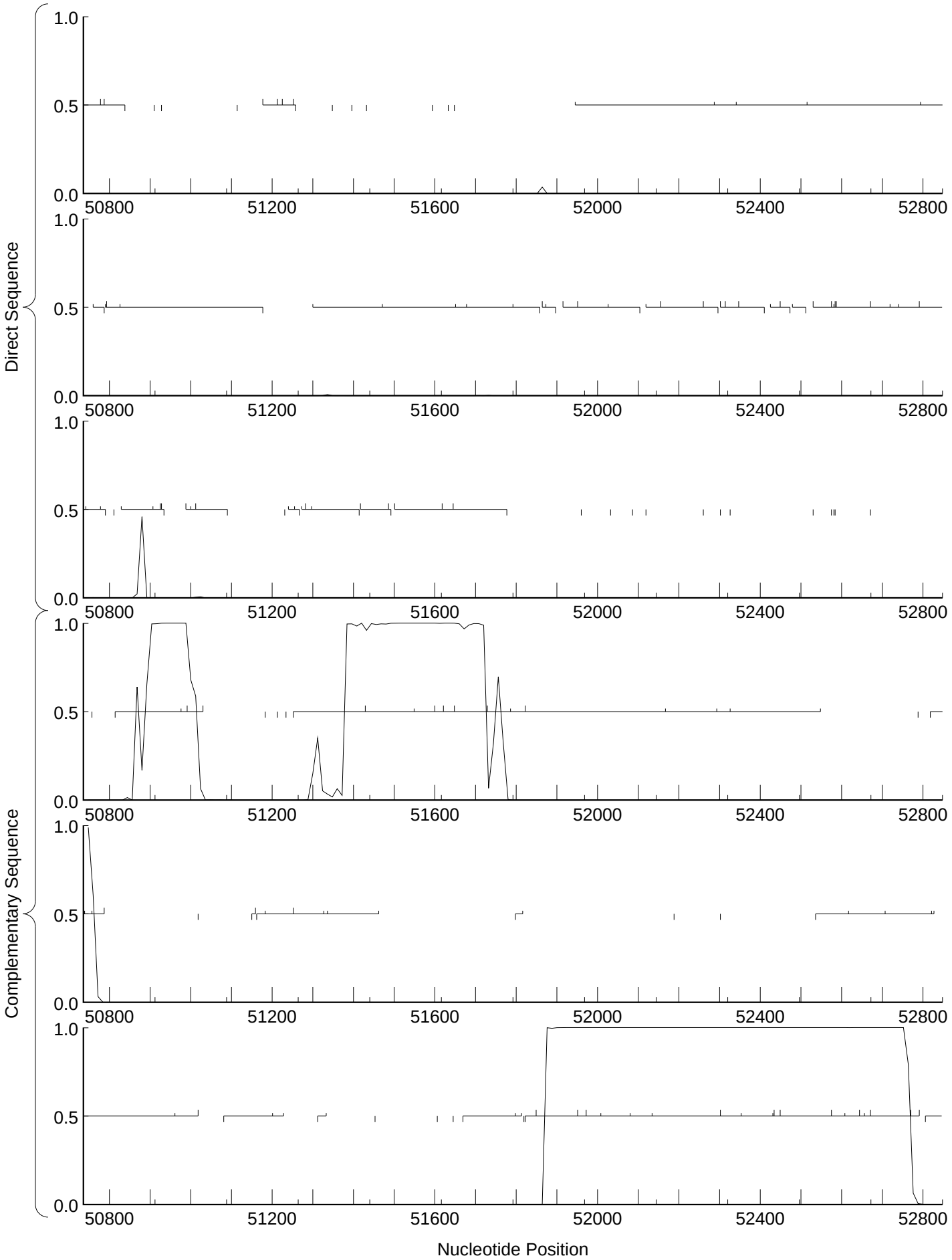
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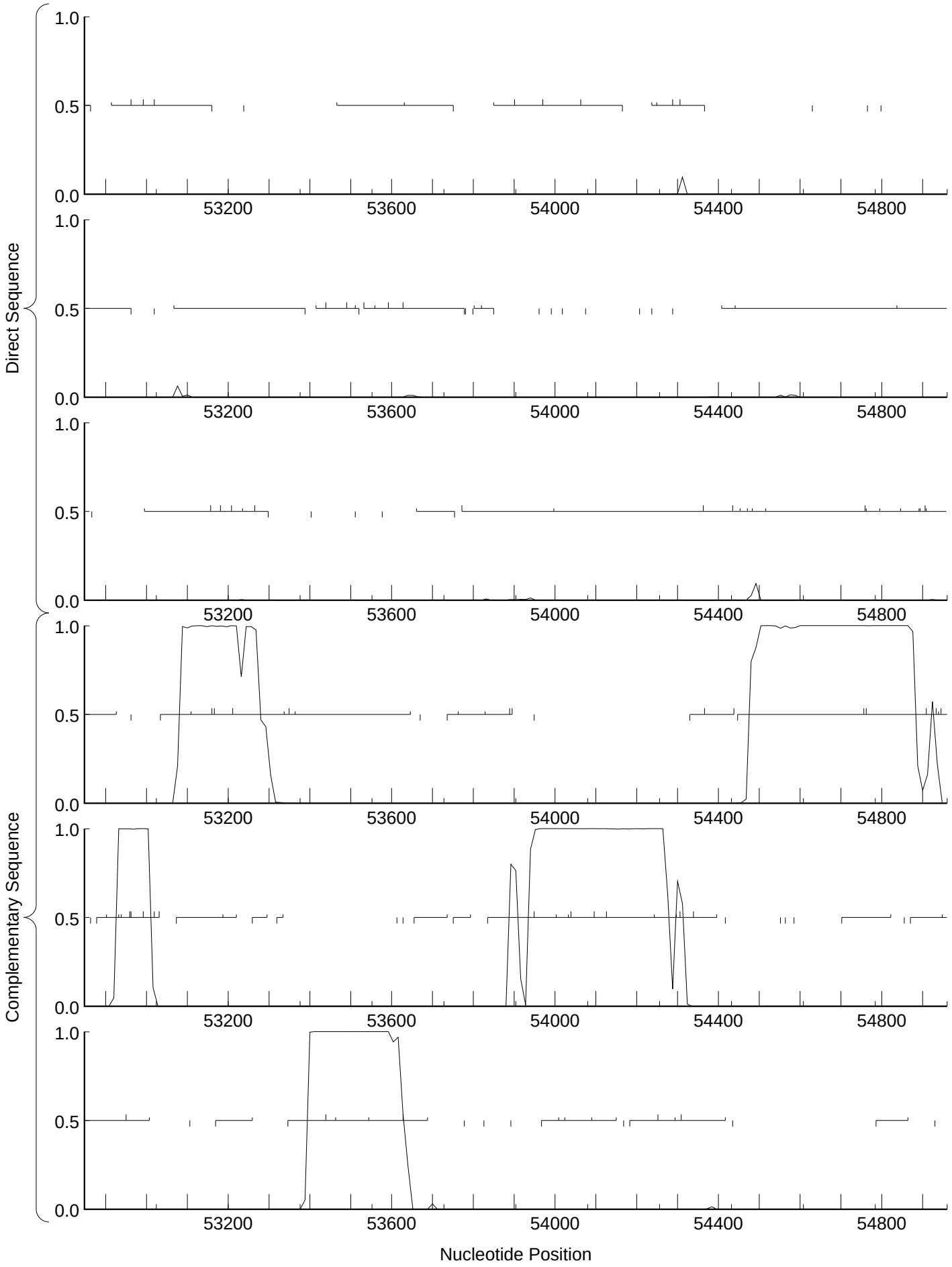
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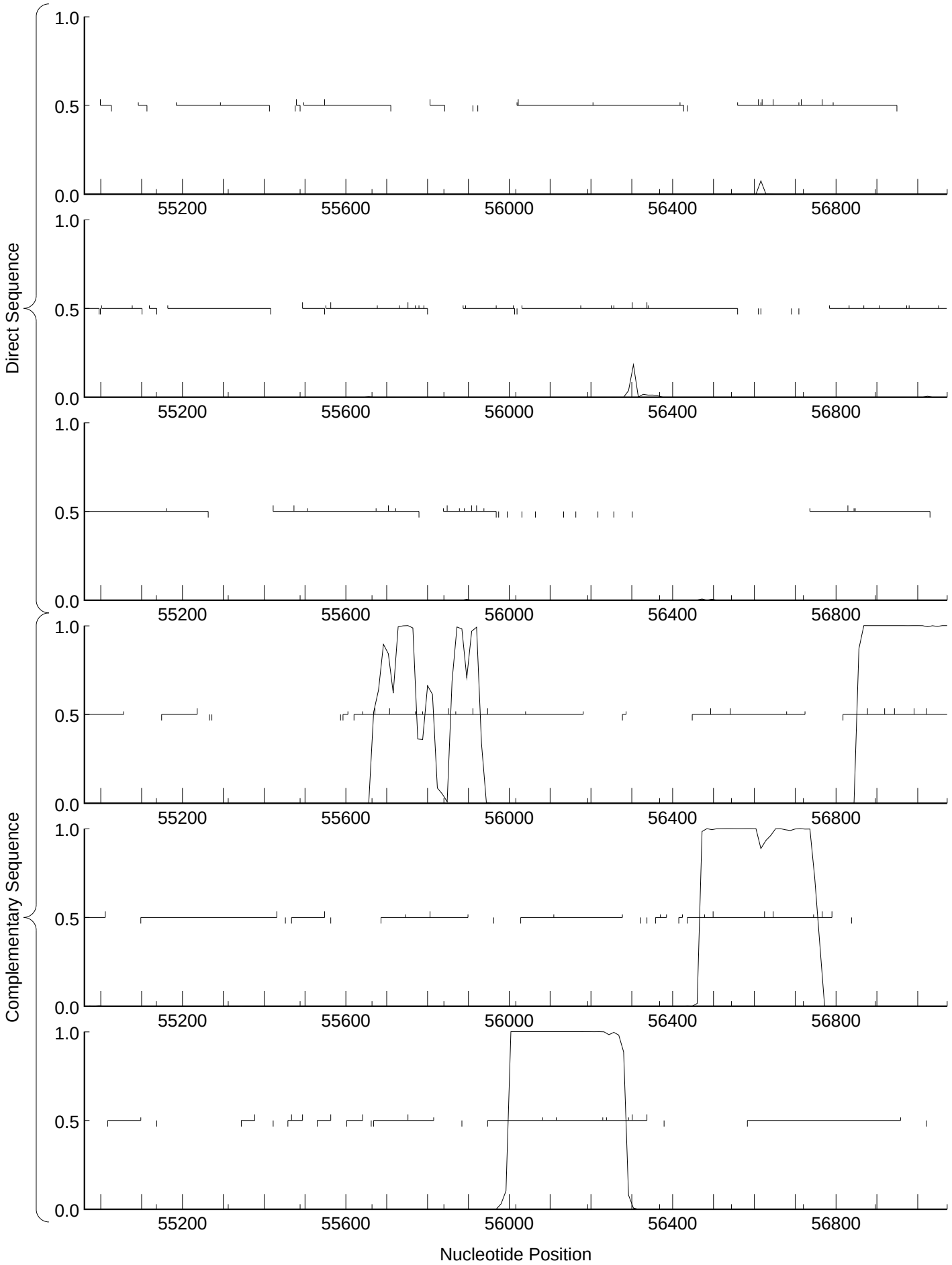
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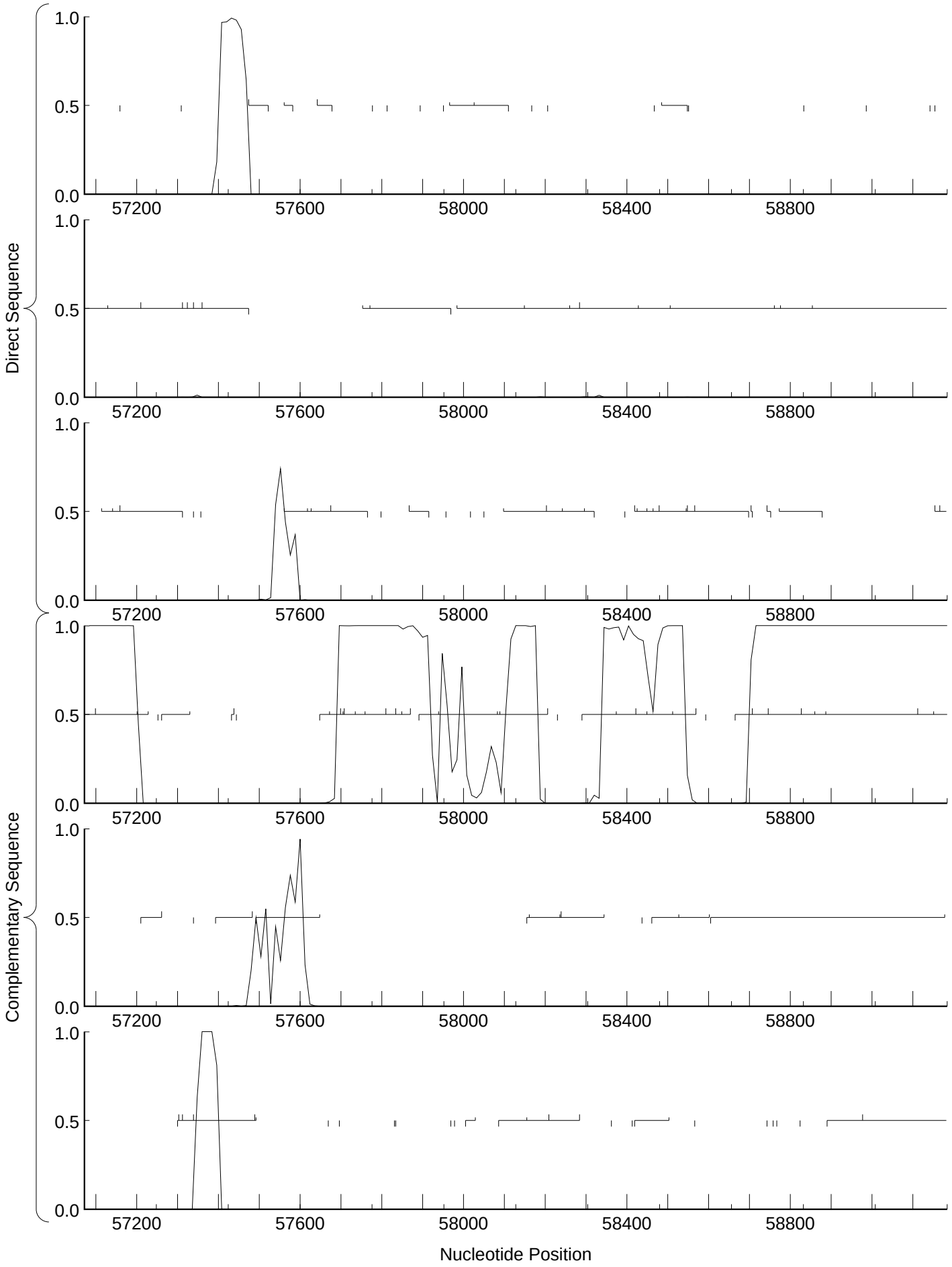
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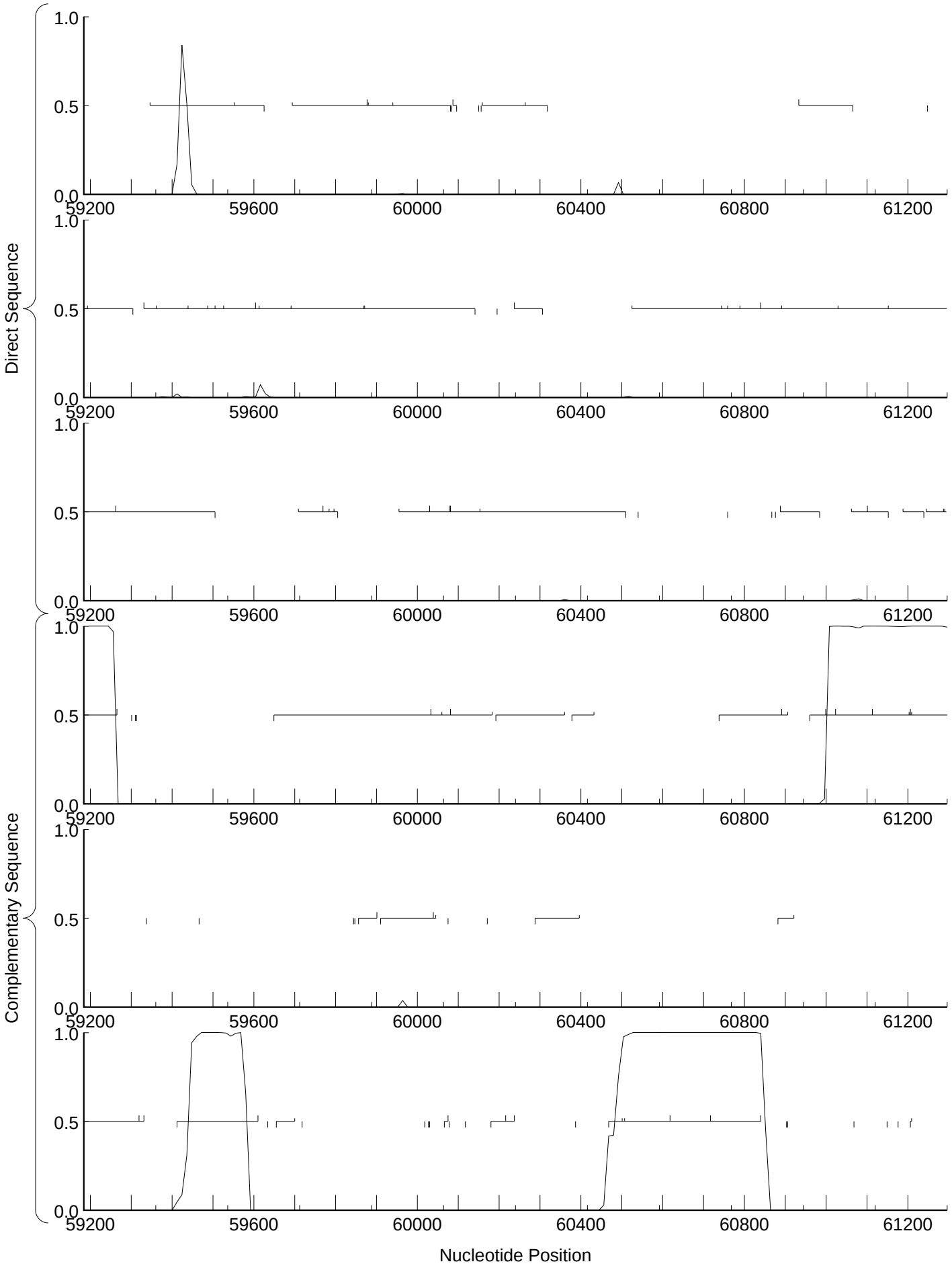
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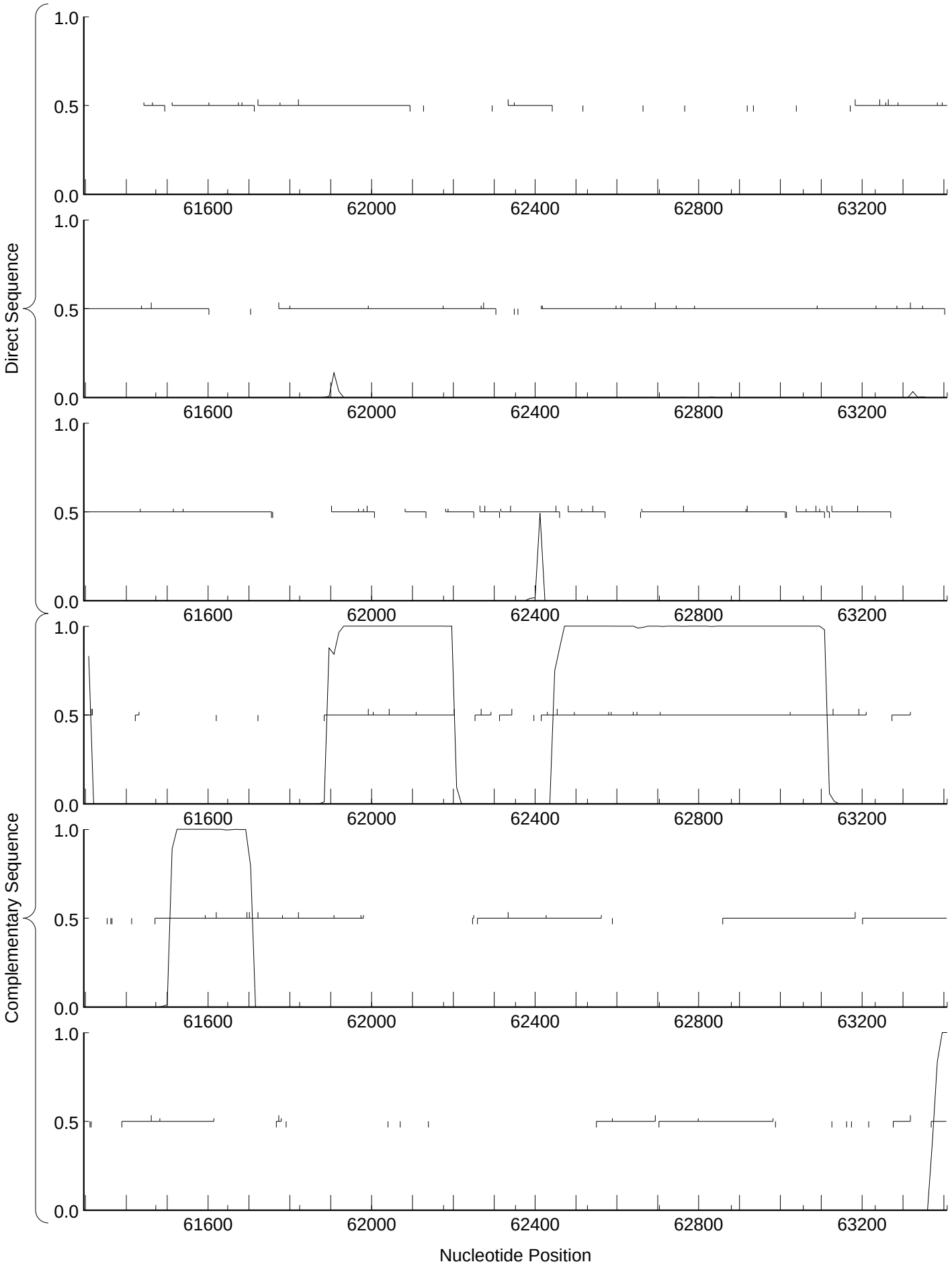
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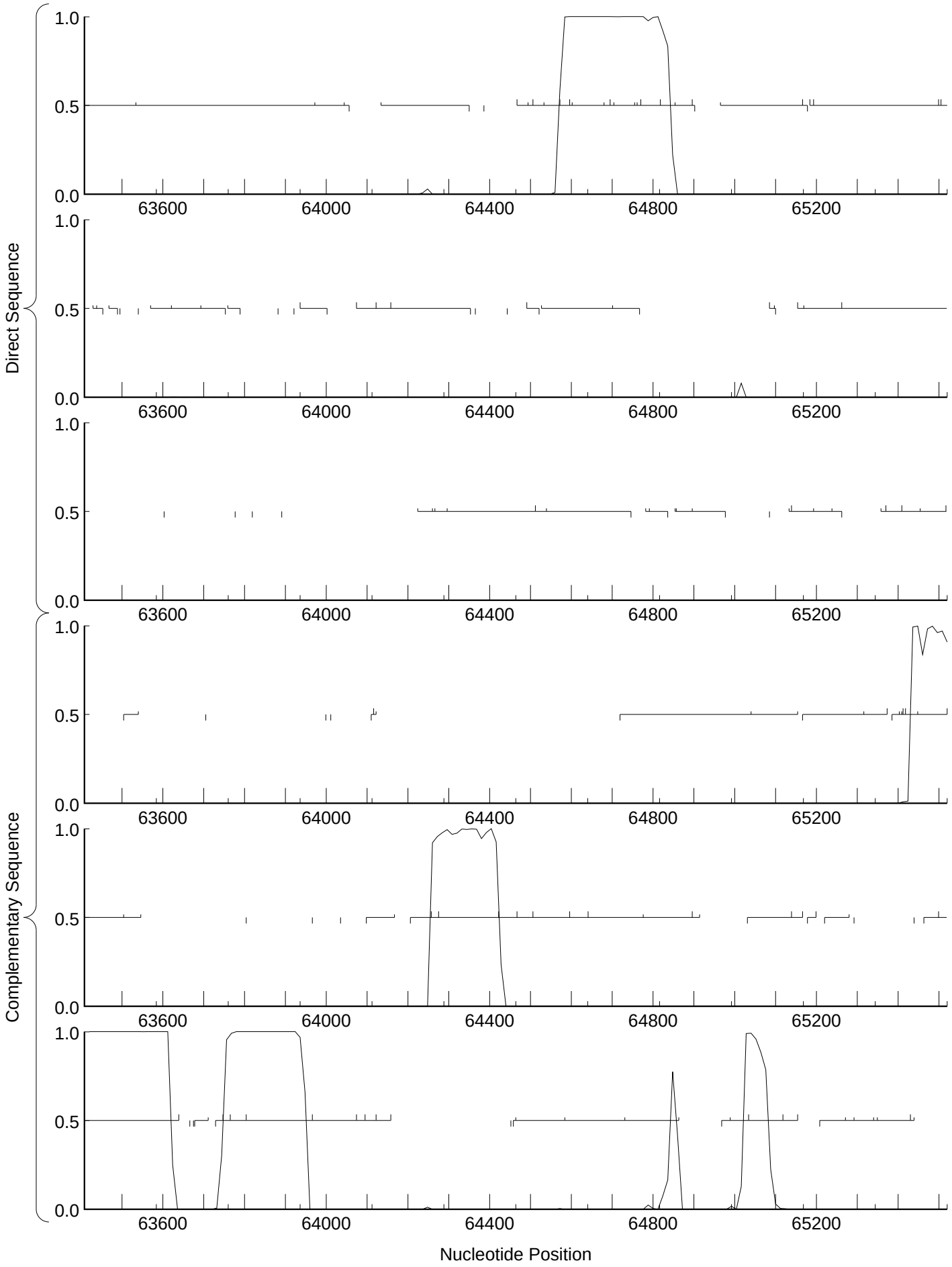
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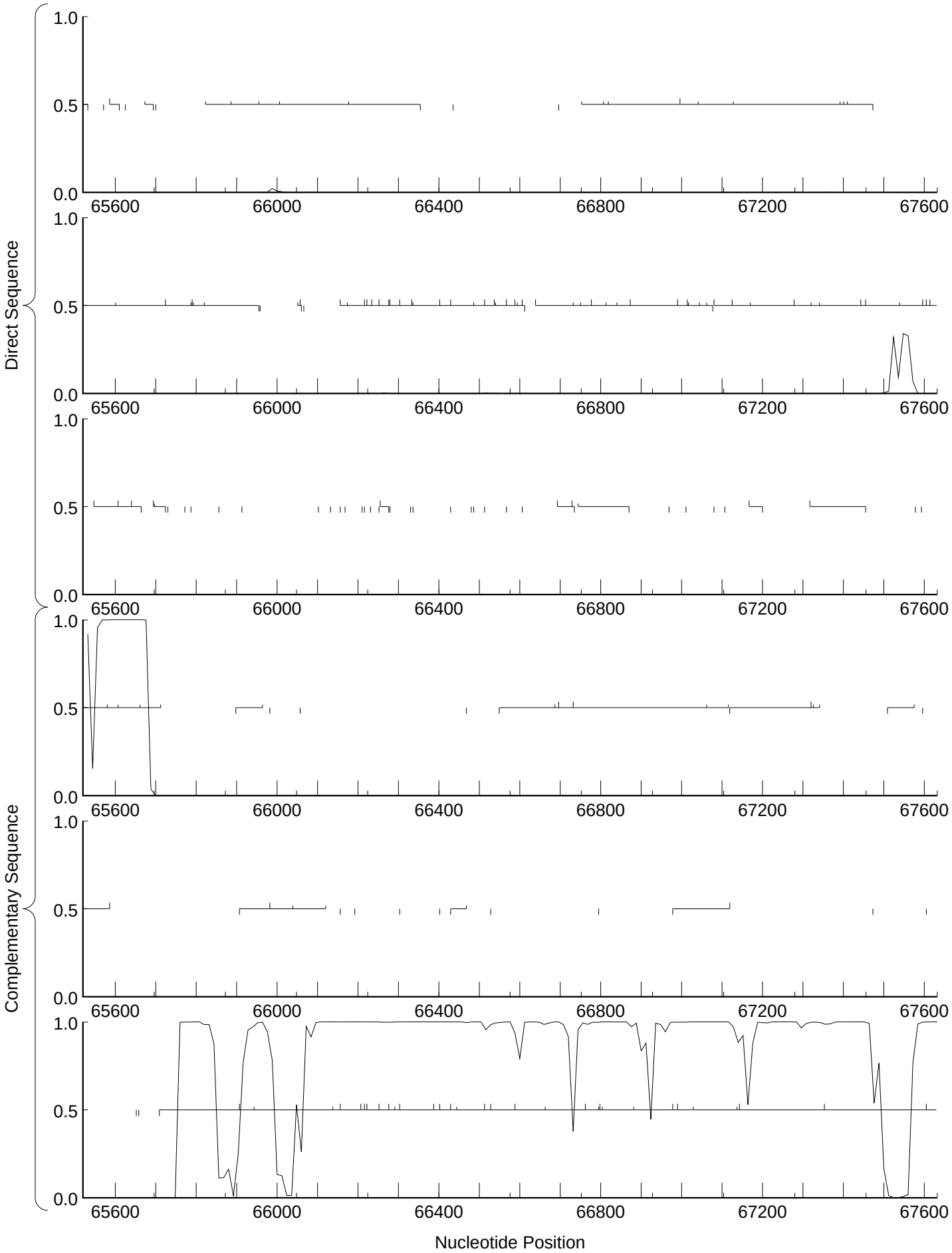
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omyces phage ShakeNBake complete sequence, 69299 bp, circularly permuted, Cluster BN, Order 4, Window 96, Step 12, 32/34



omyces phage ShakeNBake complete sequence, 69299 bp, circularly permuted, Cluster BN, Order 4, Window 96, Step 12, 33/34



omyces phage ShakeNBake complete sequence, 69299 bp, circularly permuted, Cluster BN, Order 4, Window 96, Step 12, 34/34

