GeneMark

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

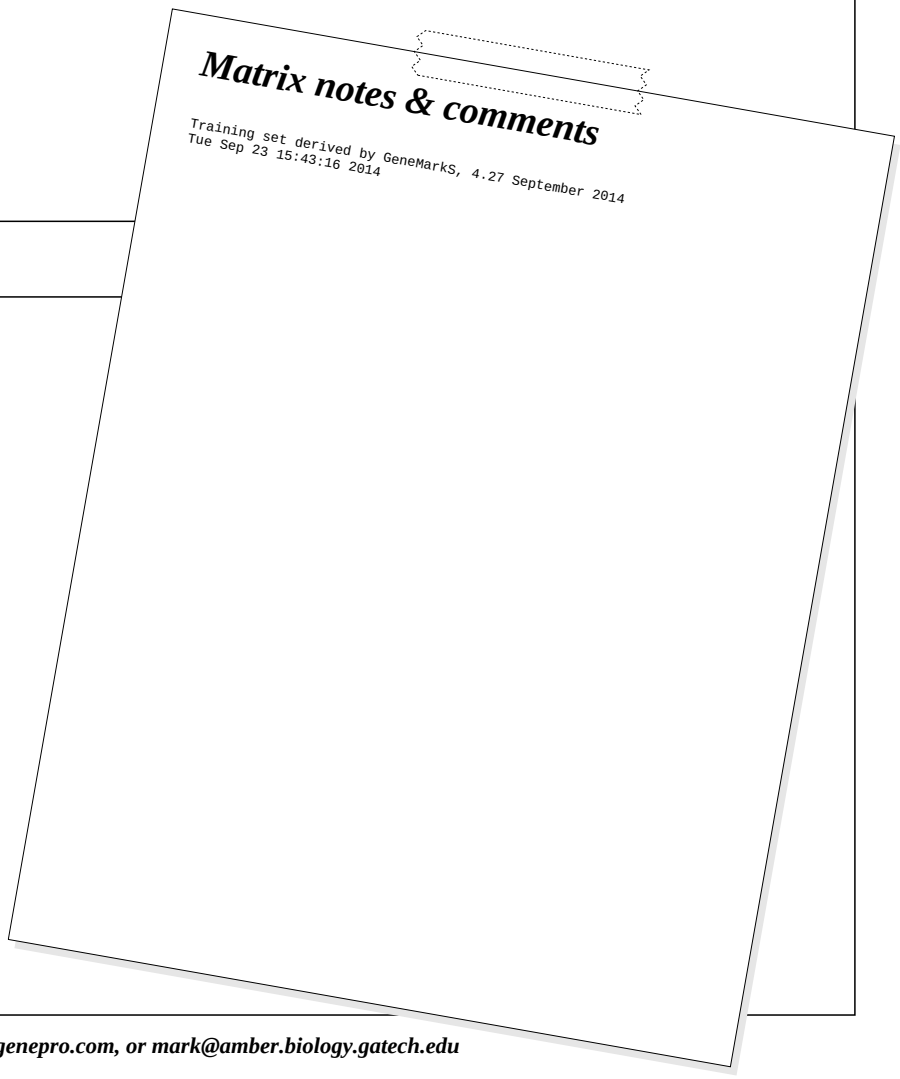
PROGRAM INFORMATION

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Sequence         :  Streptomyces phage Janus complete sequence, 50632 bp including 11-base 3' overhang (CGCCGTG
Analysis Date    :  3/28/24 at 22:14:23
Pages            :  25
Sequence Length  :  50632 bp
GC Content       :   67.05%

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

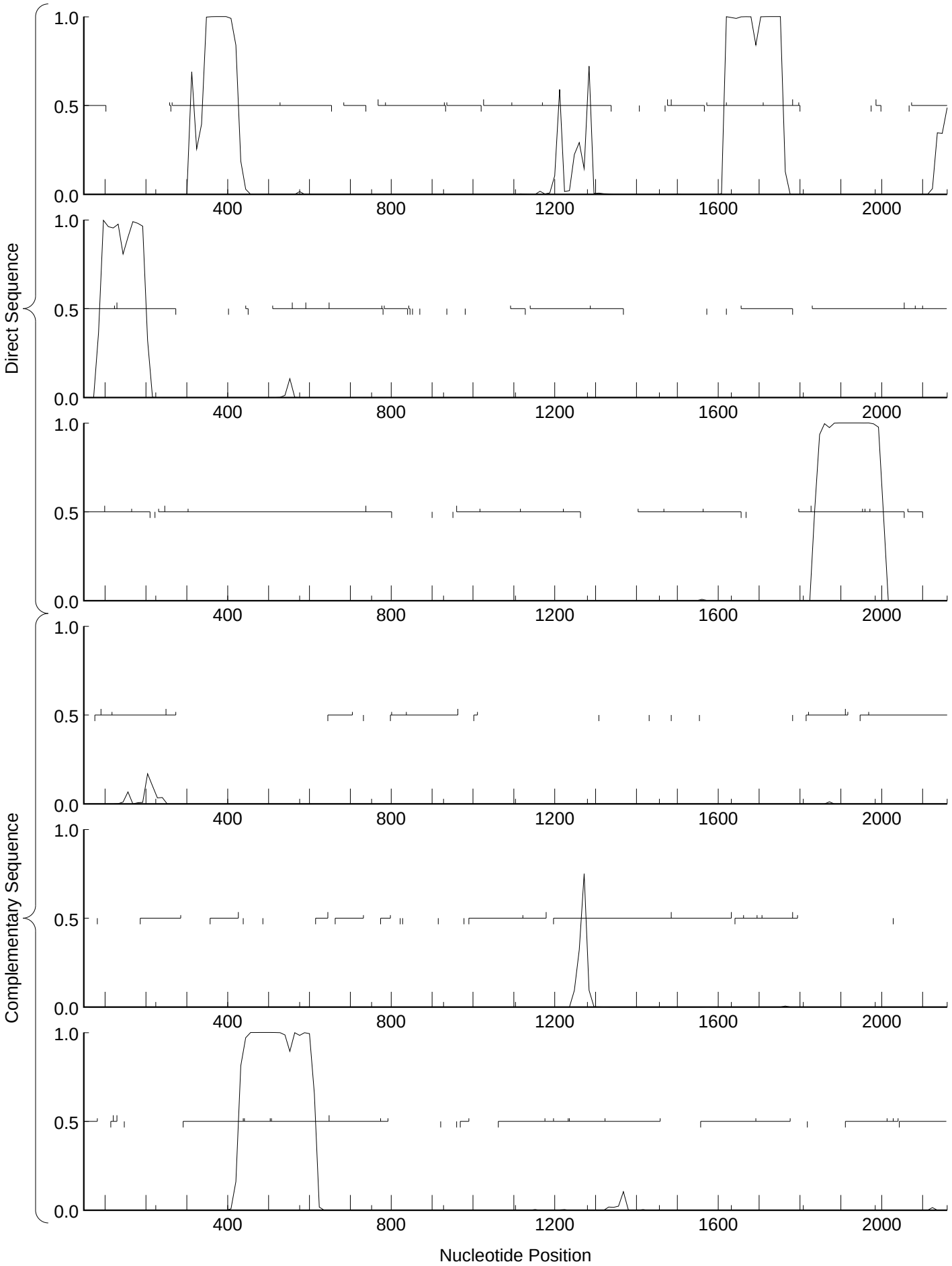
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Matrix     :  Streptomyces_scabiei_87_22
Author     :  -

Order      :  4
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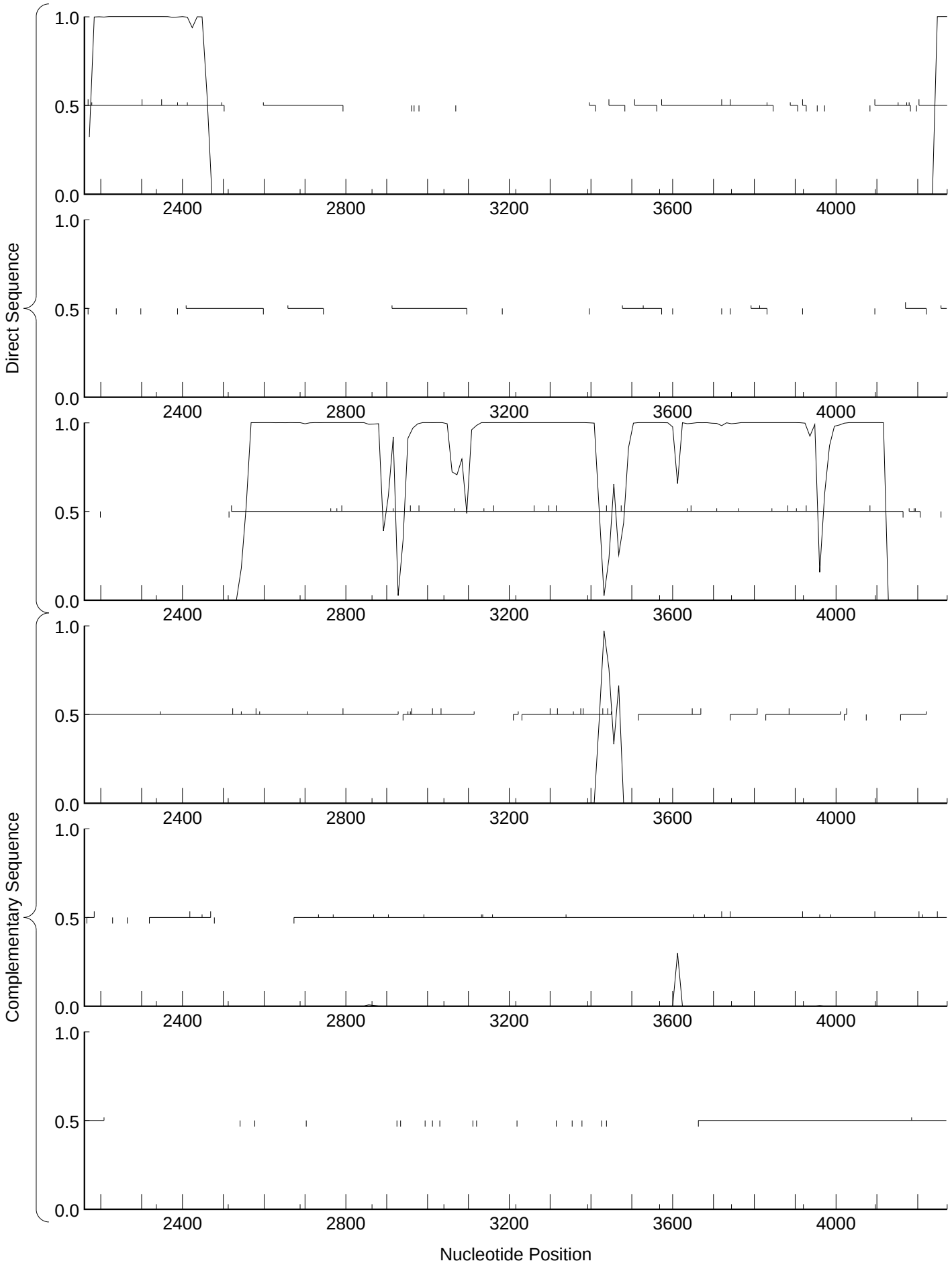

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

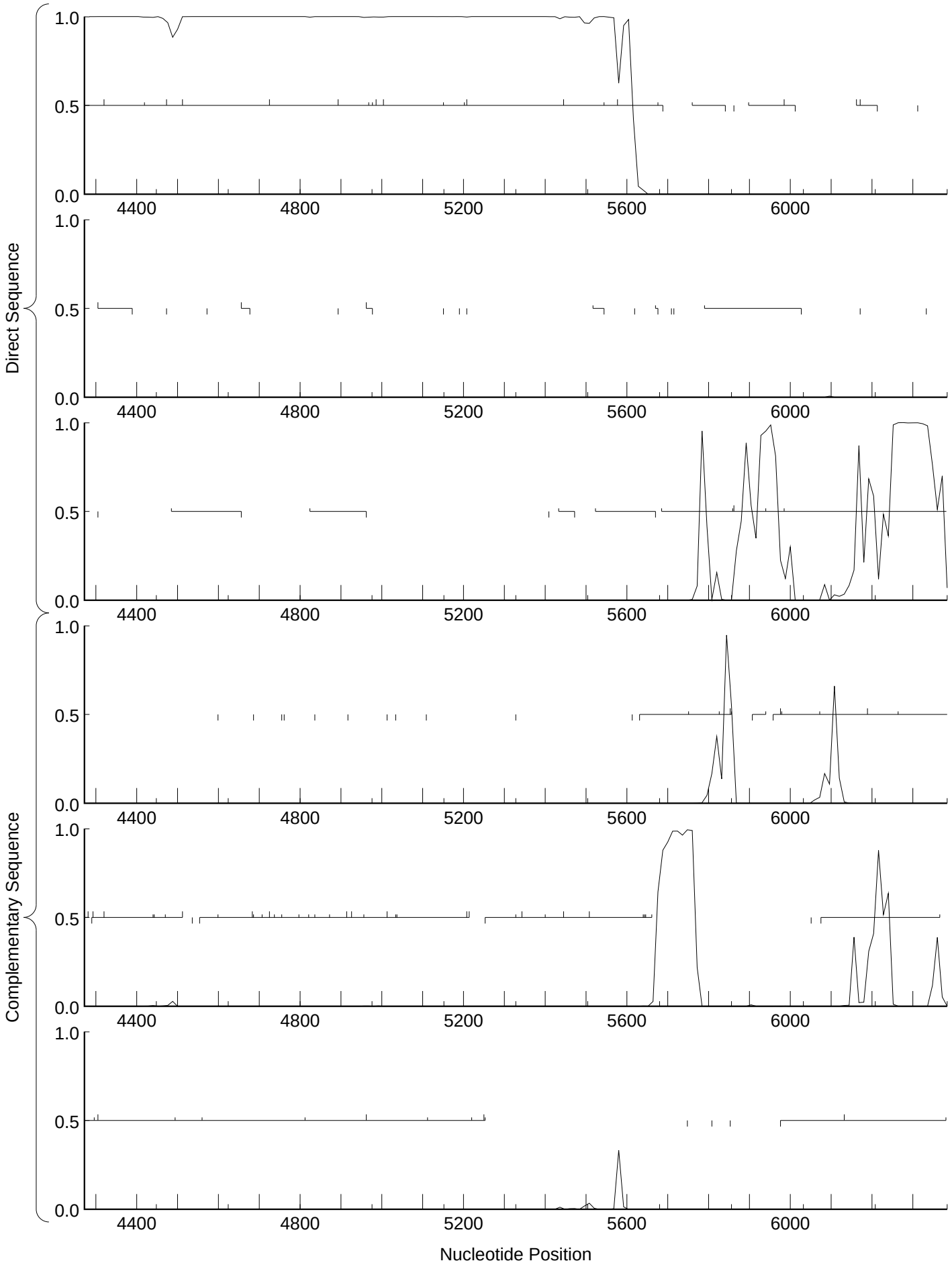
complete sequence, 50632 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 2/25



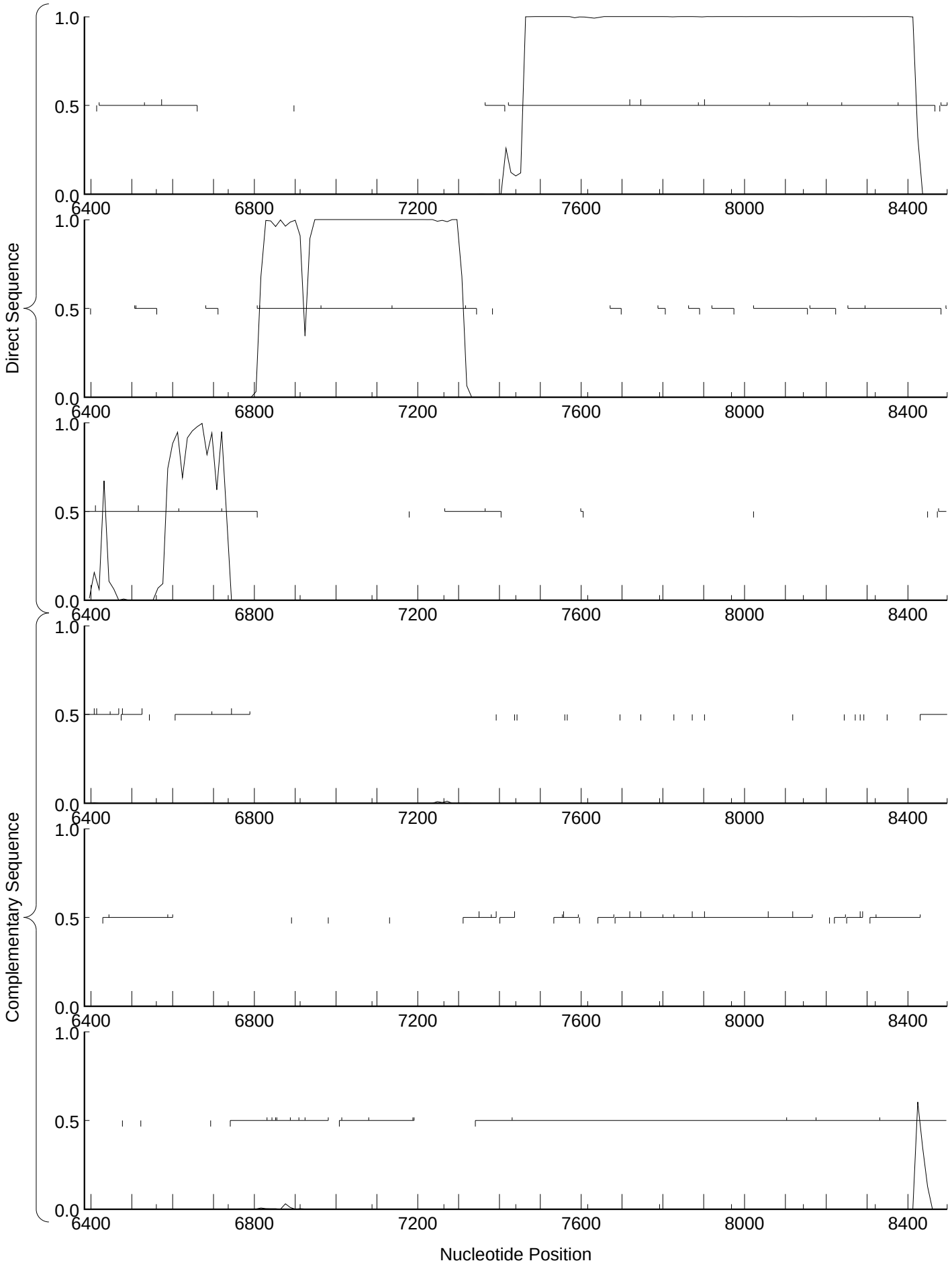
complete sequence, 50632 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 3/25



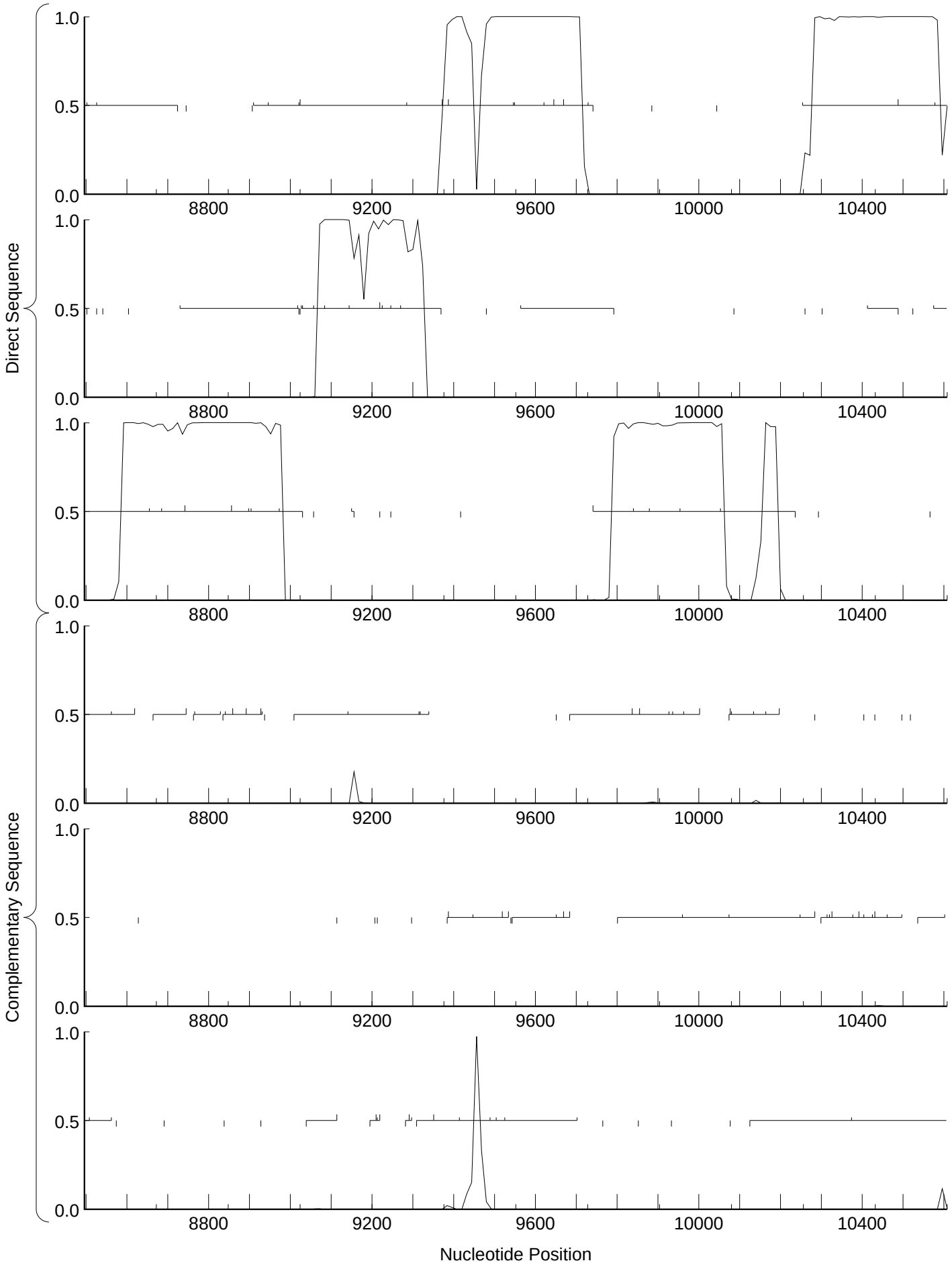
complete sequence, 50632 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 4/25



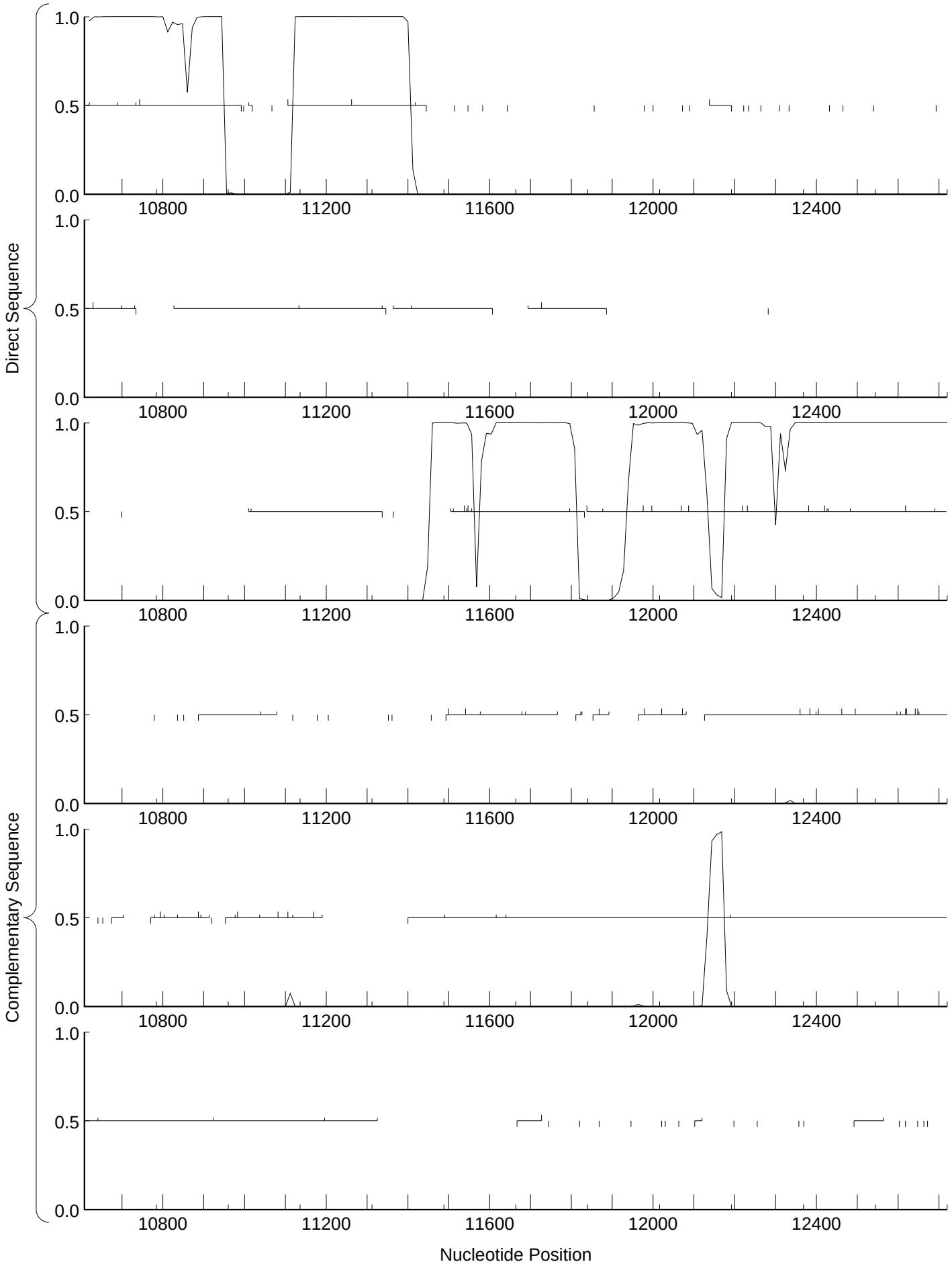
complete sequence, 50632 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 5/25



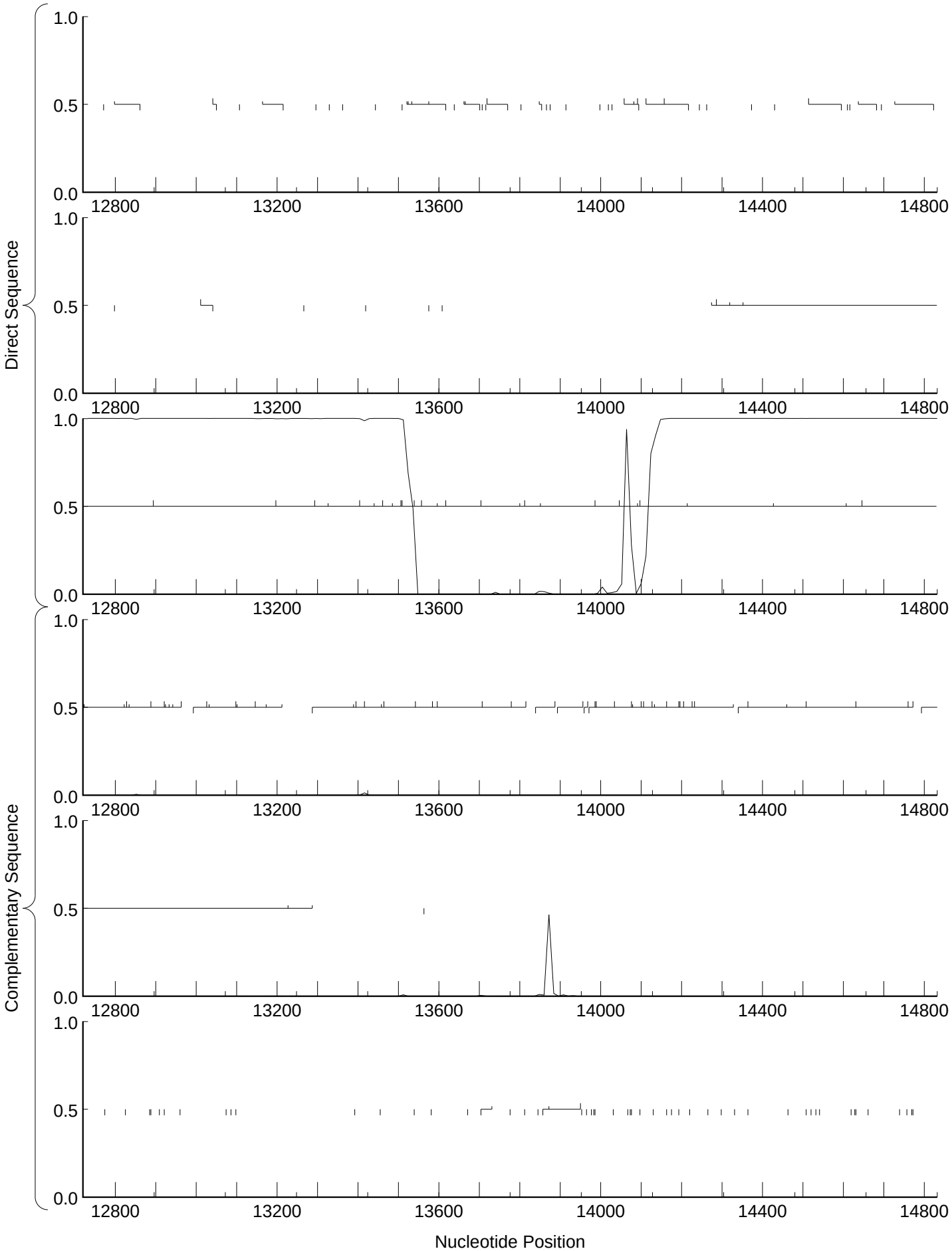
complete sequence, 50632 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 6/25



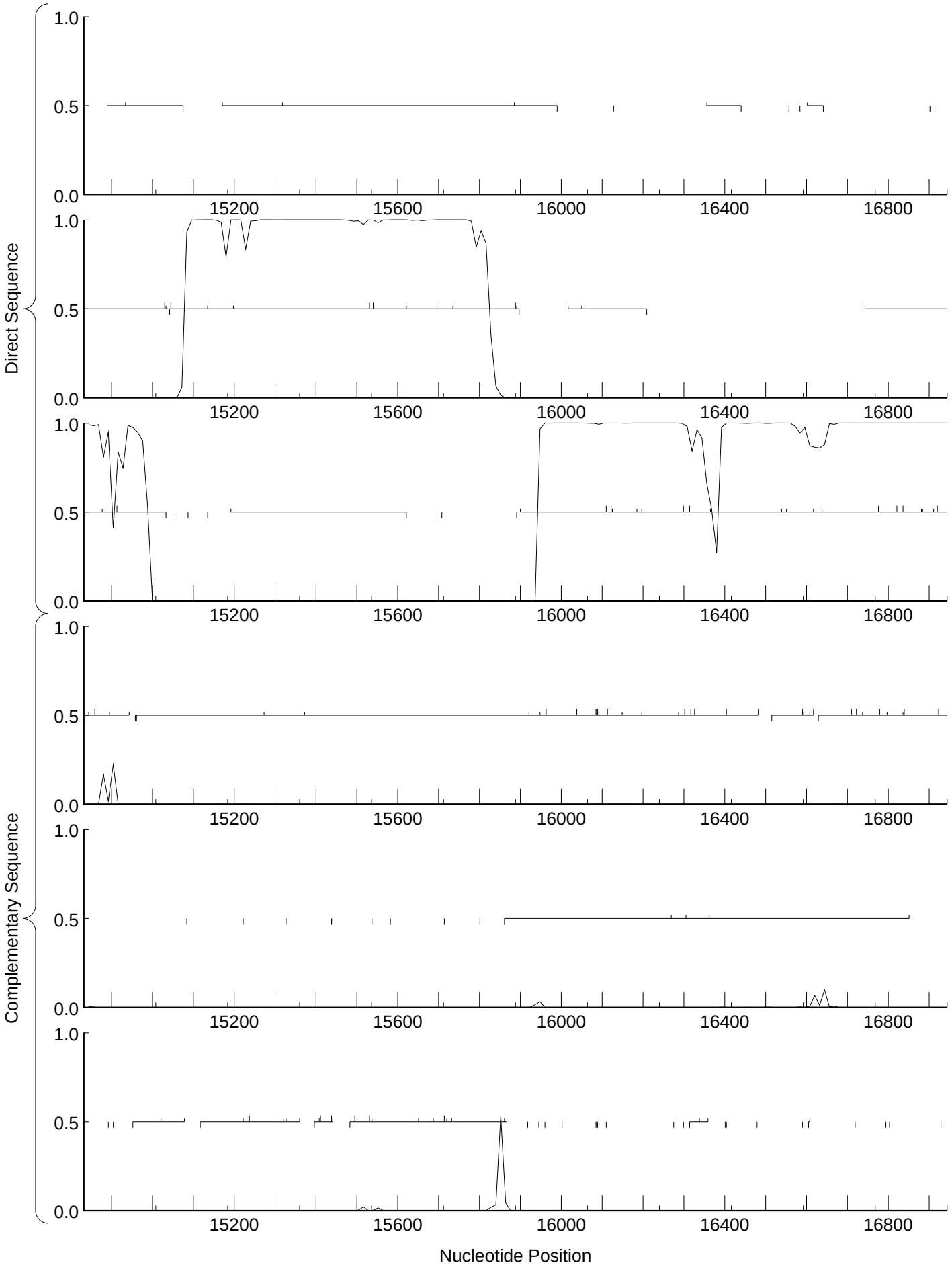
complete sequence, 50632 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 7/25



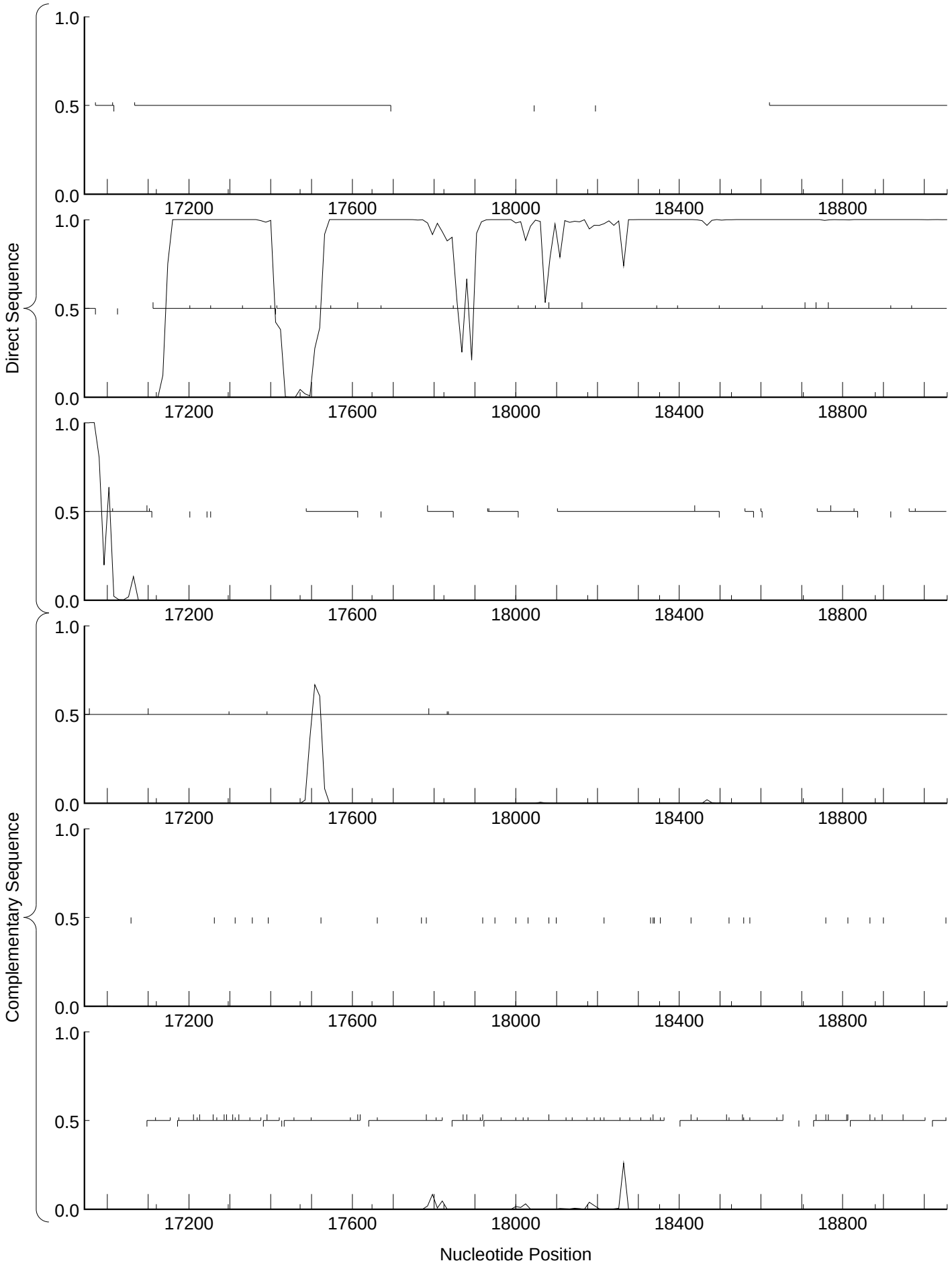
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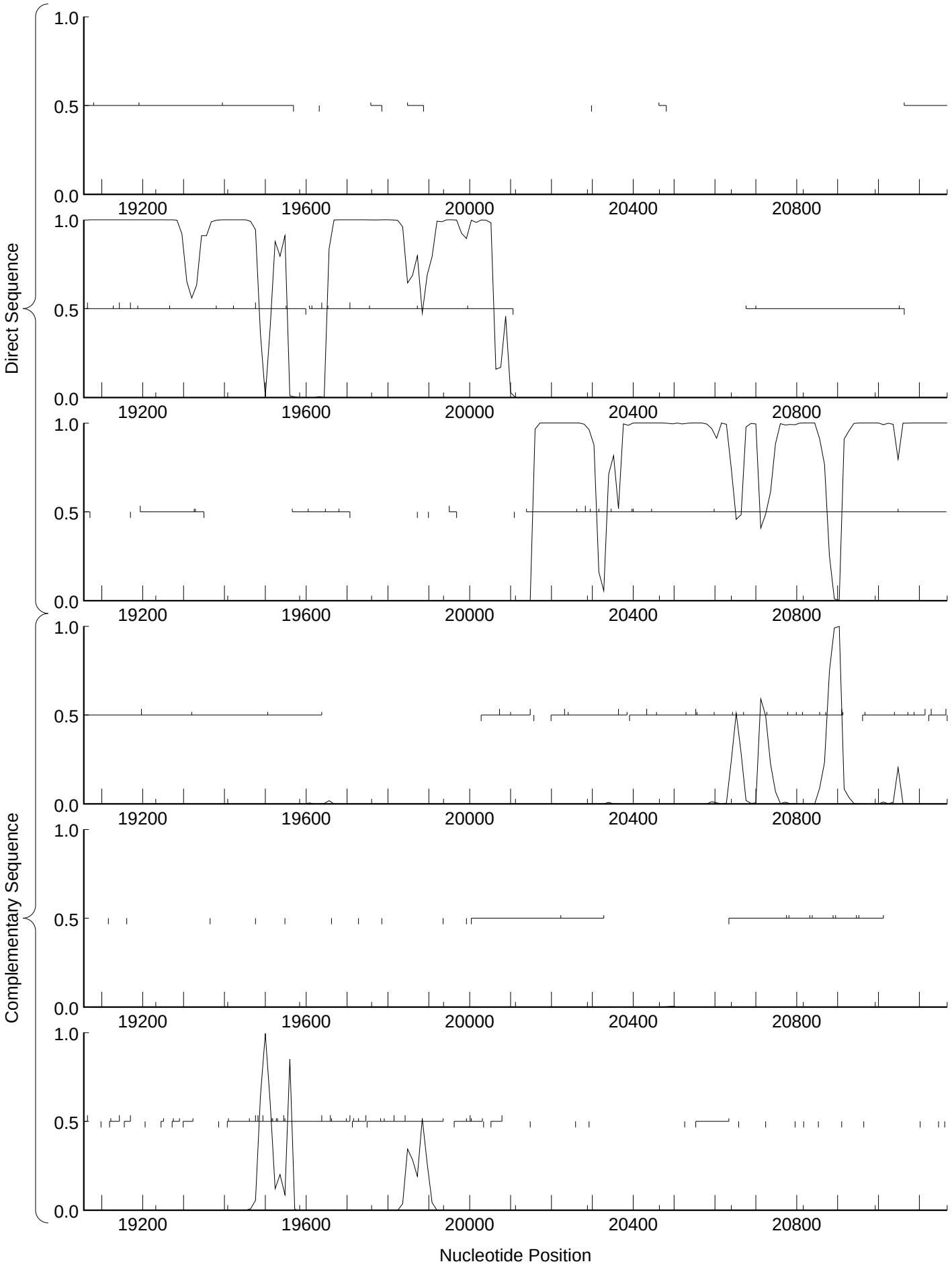
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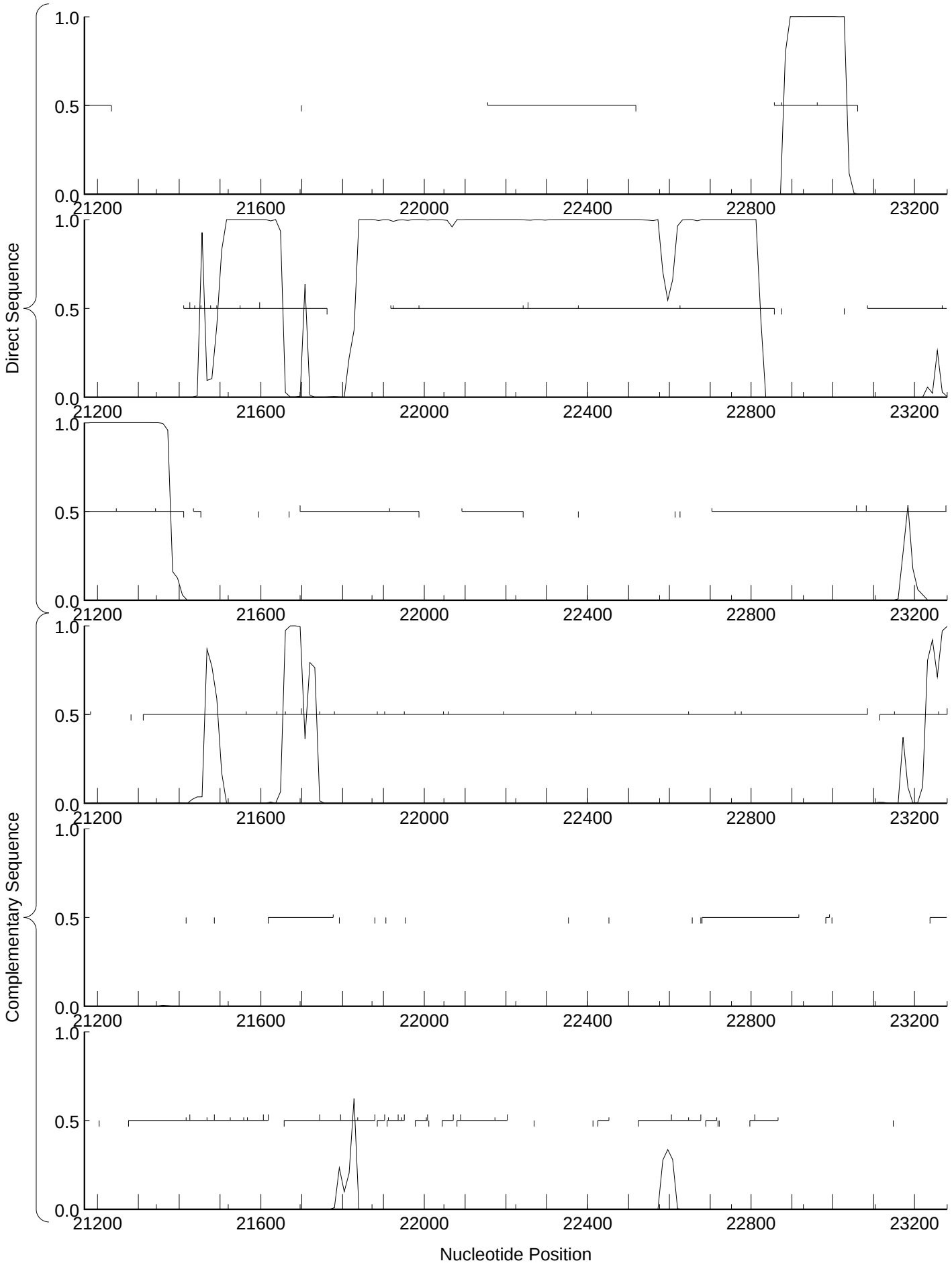
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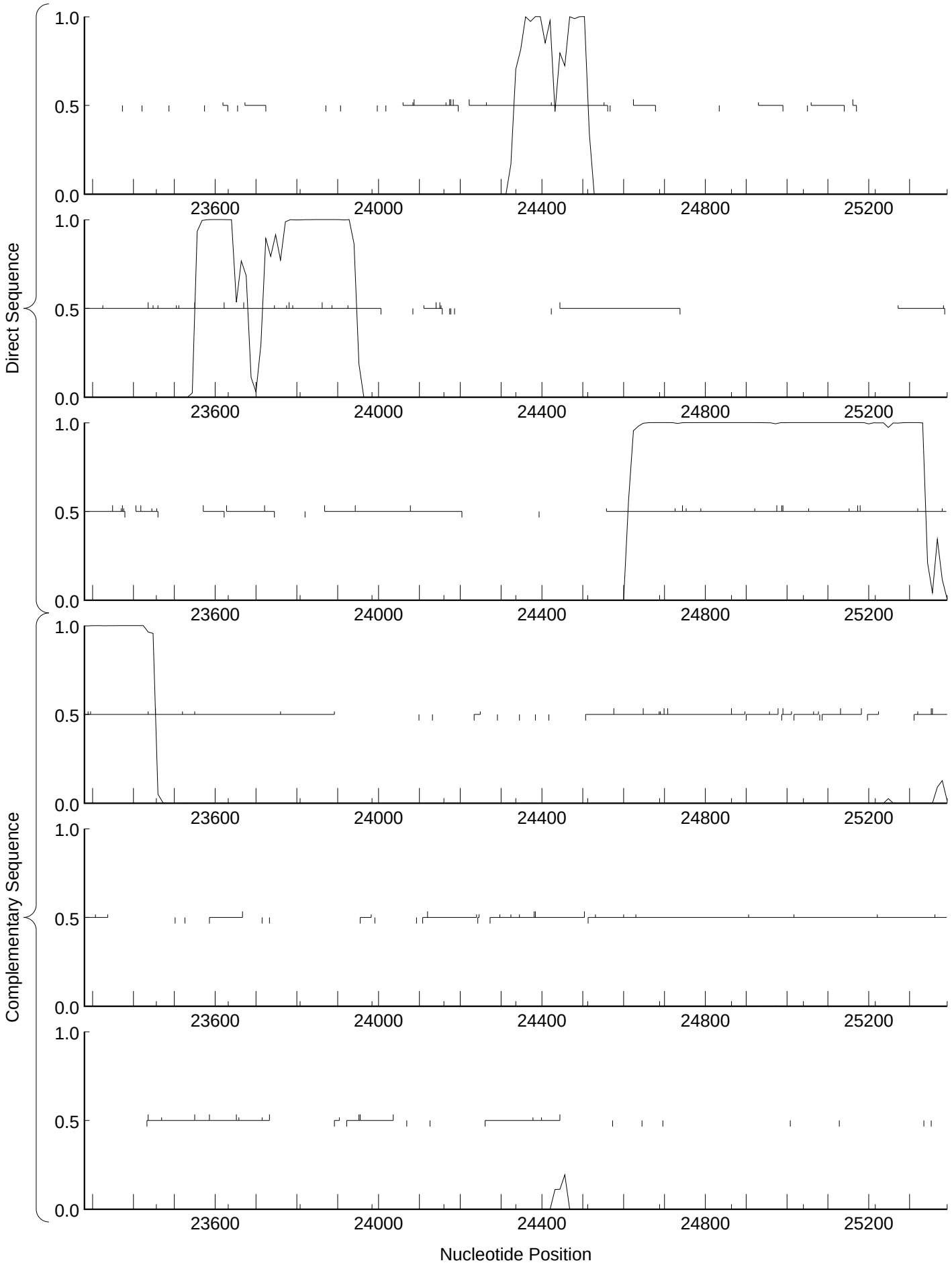
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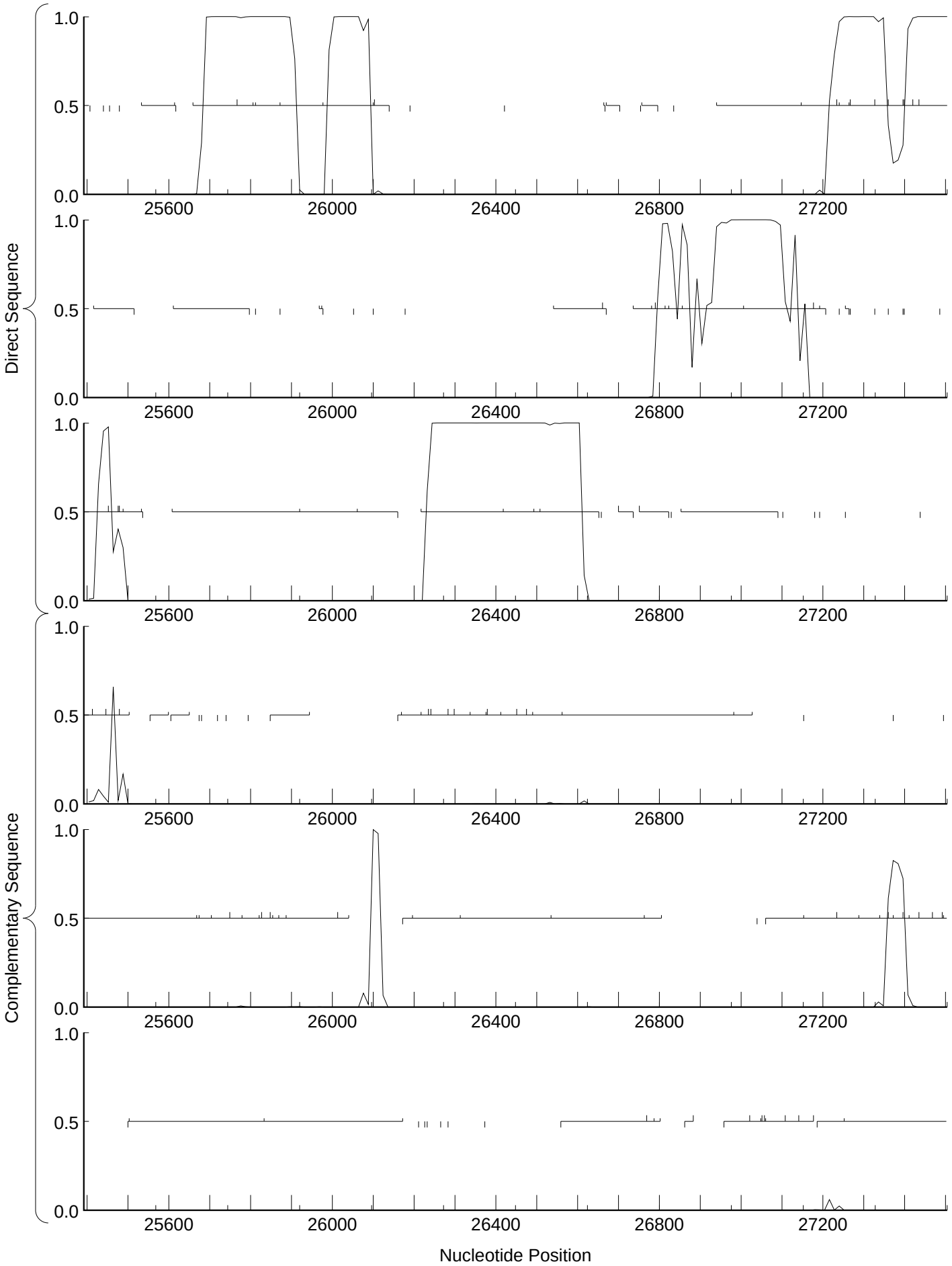


omplete sequence, 50632 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 12/25

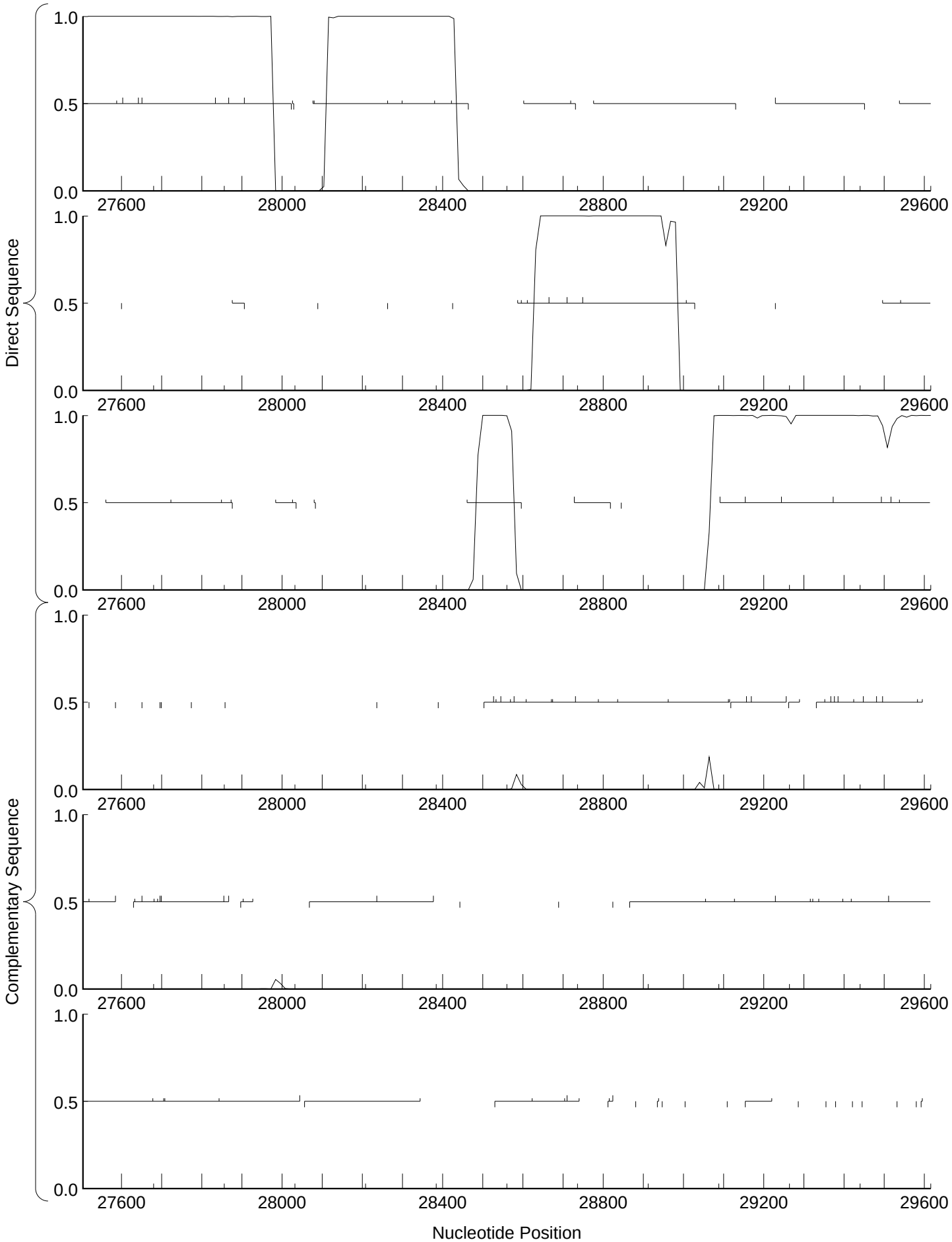


omplete sequence, 50632 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 13/25

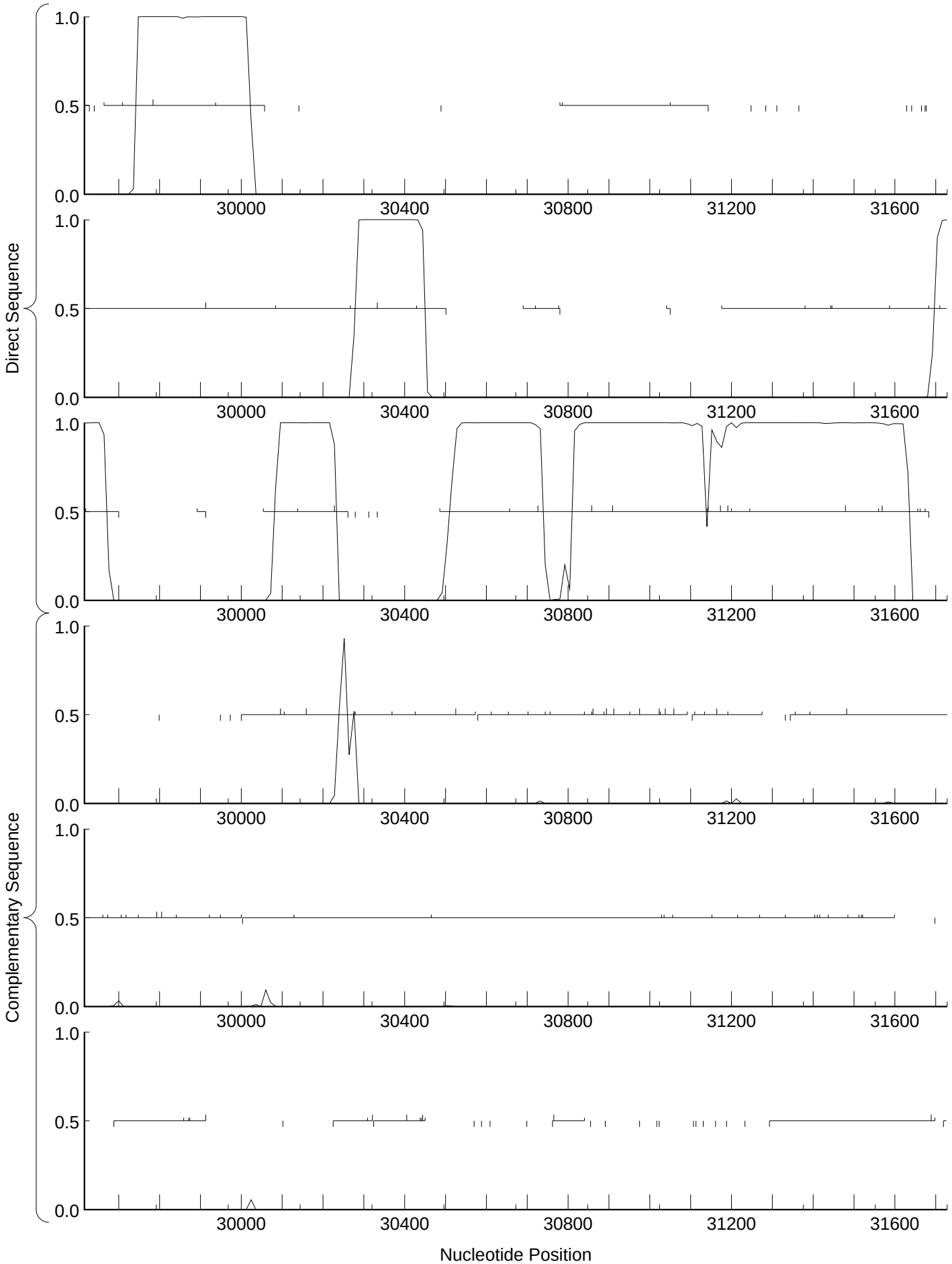


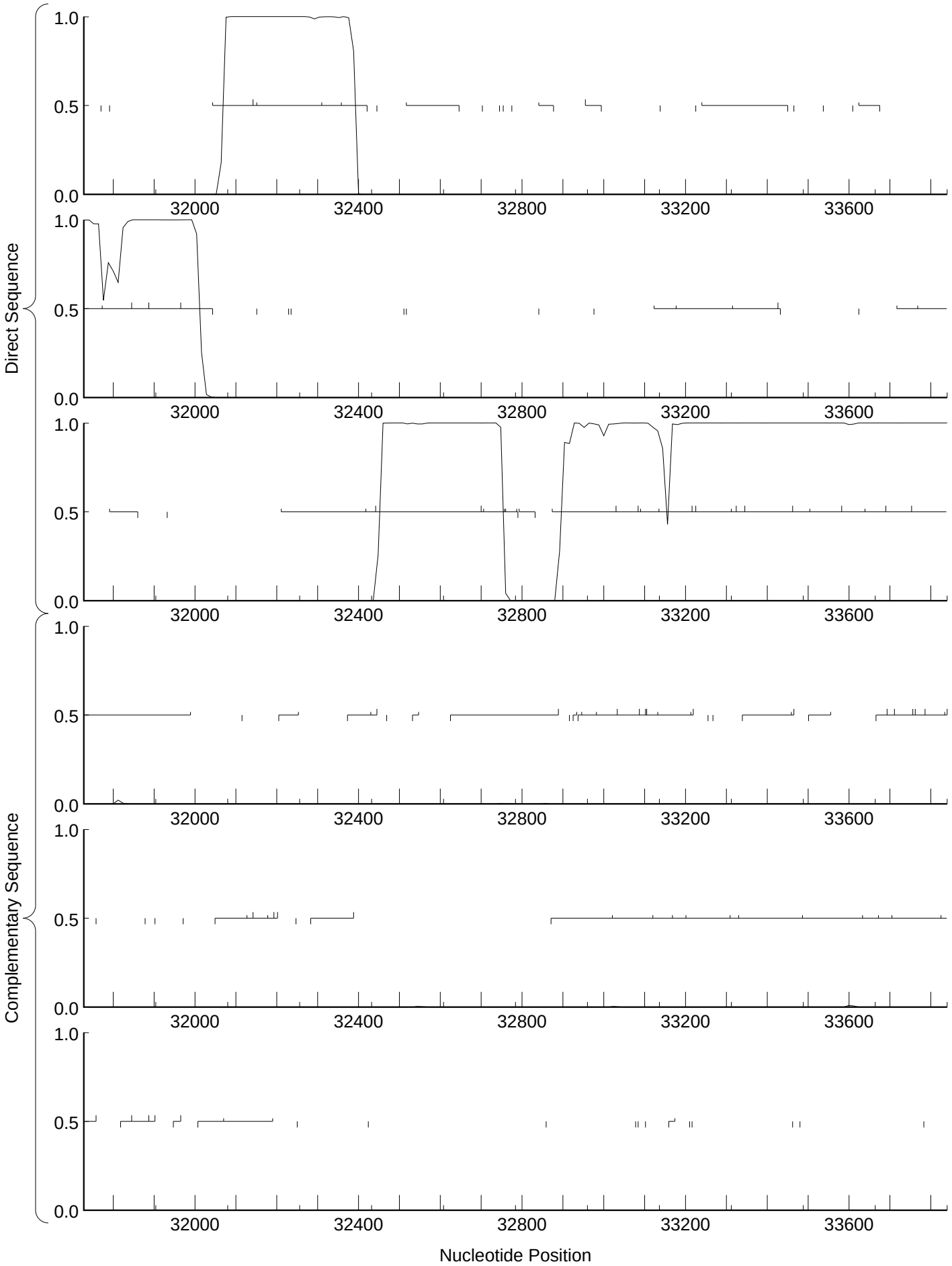

Direct Sequence
Complementary Sequence
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0.5
0.0
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26000
26400
26800
27200
Nucleotide Position

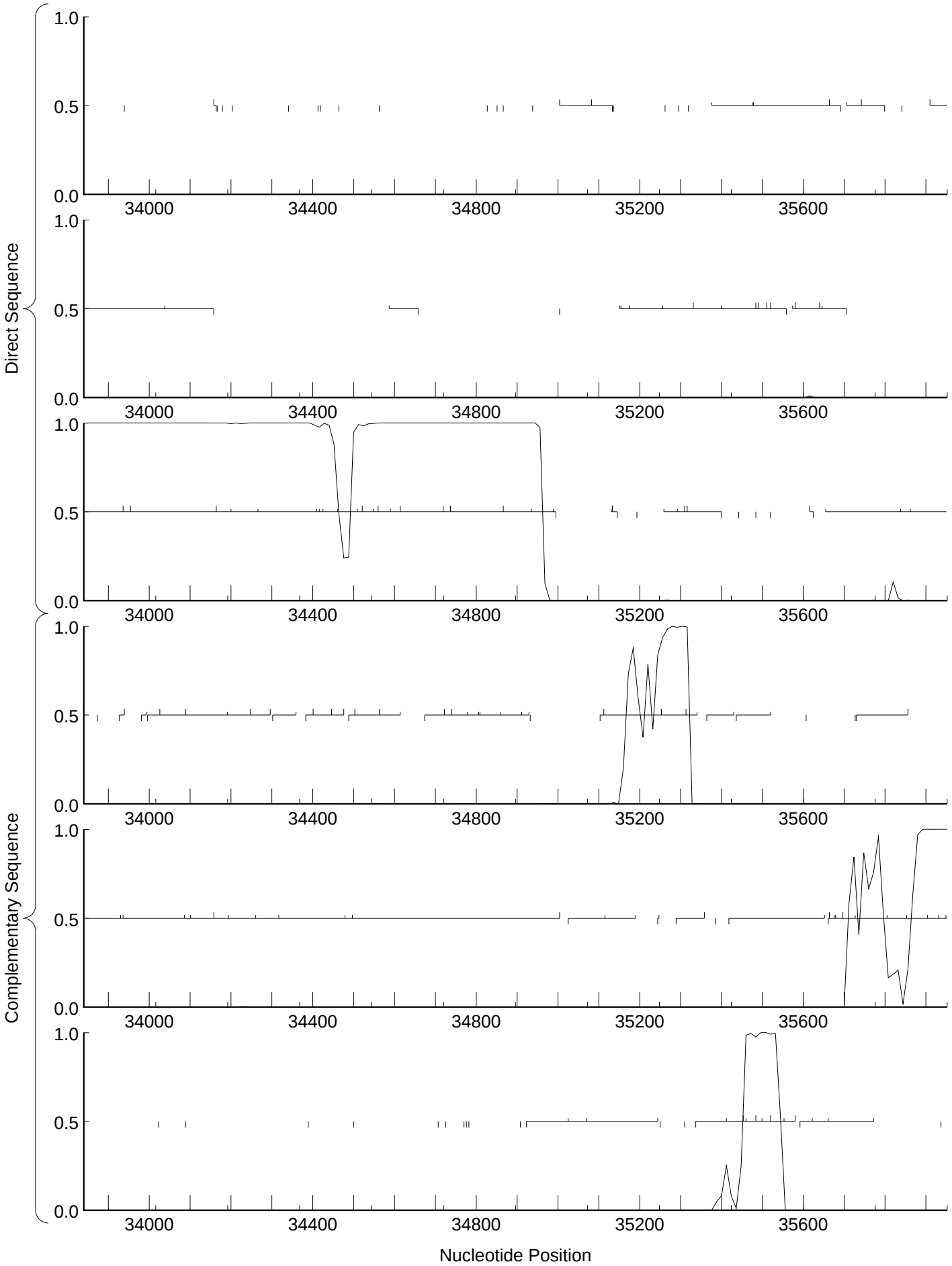
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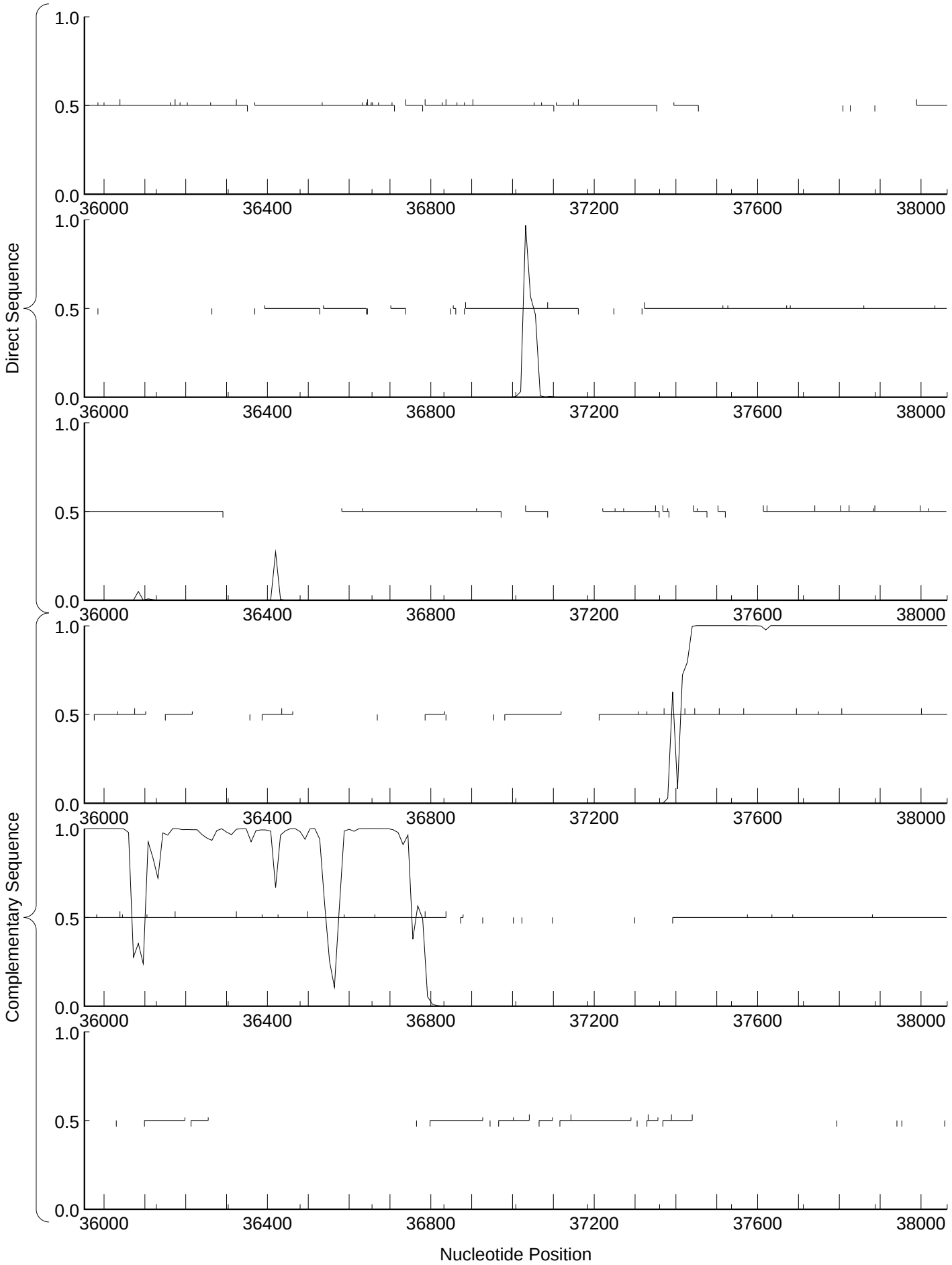


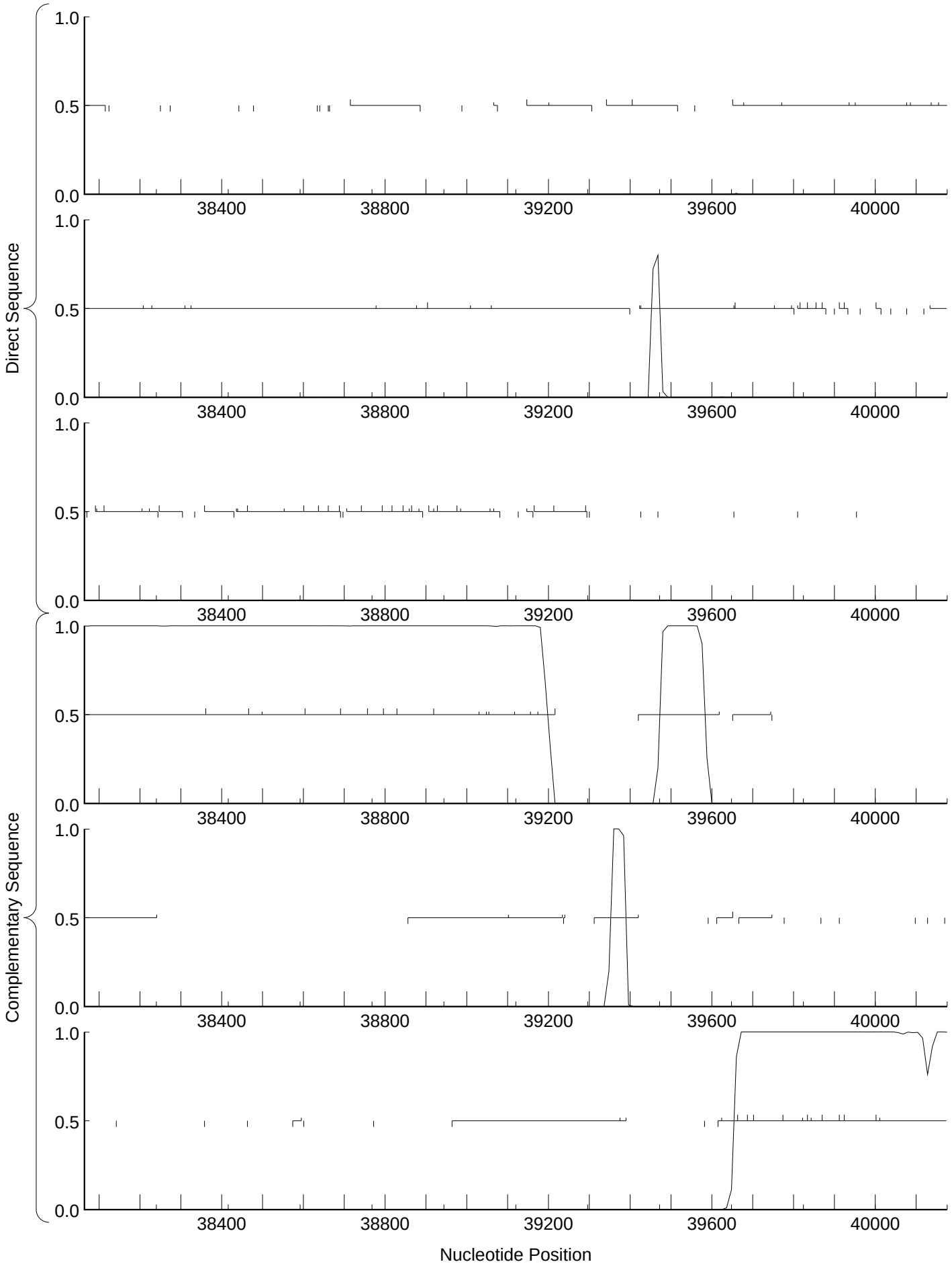
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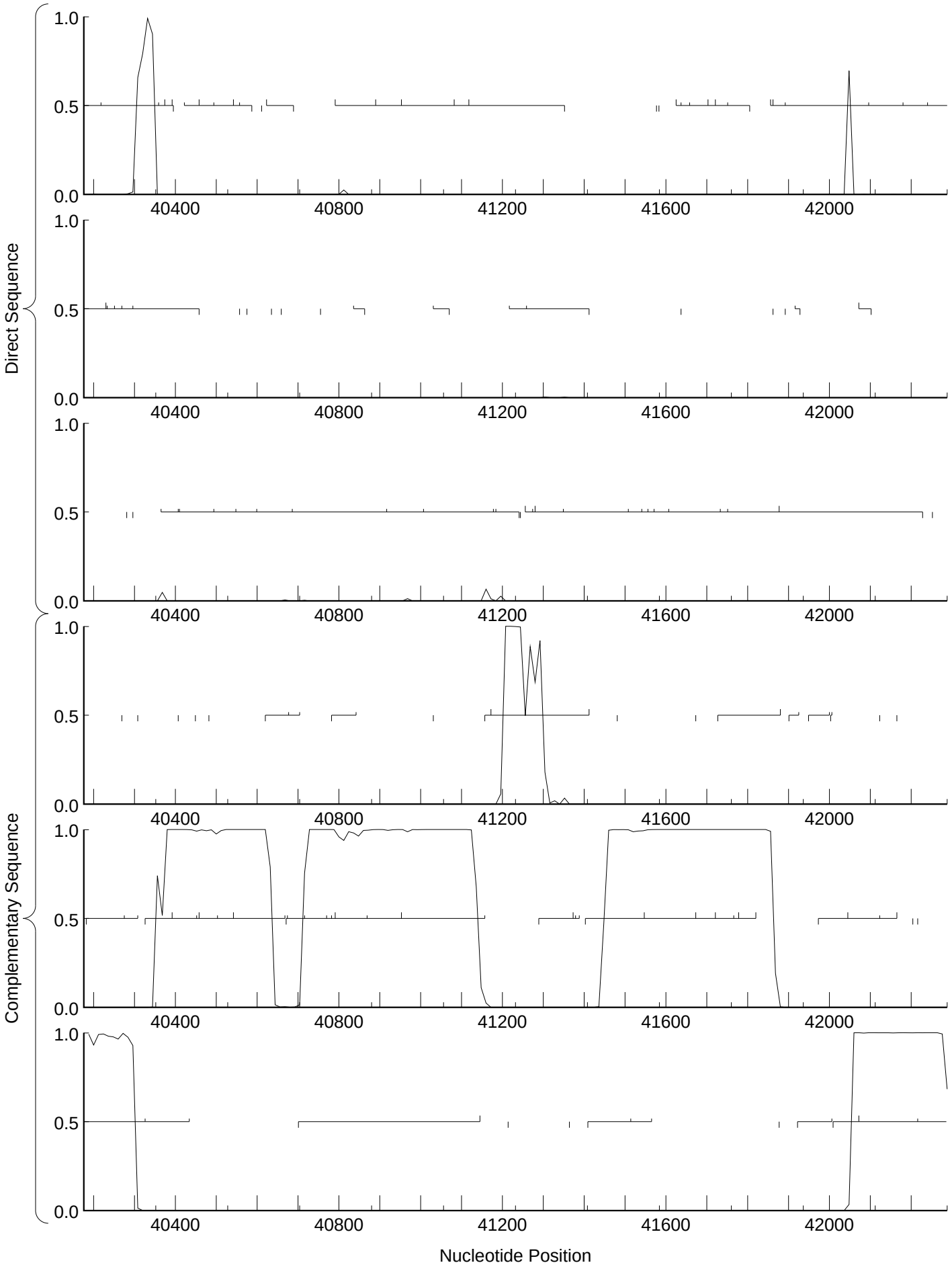

Direct Sequence
Complementary Sequence
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0.5
0.0
32000
32400
32800
33200
33600
Nucleotide Position

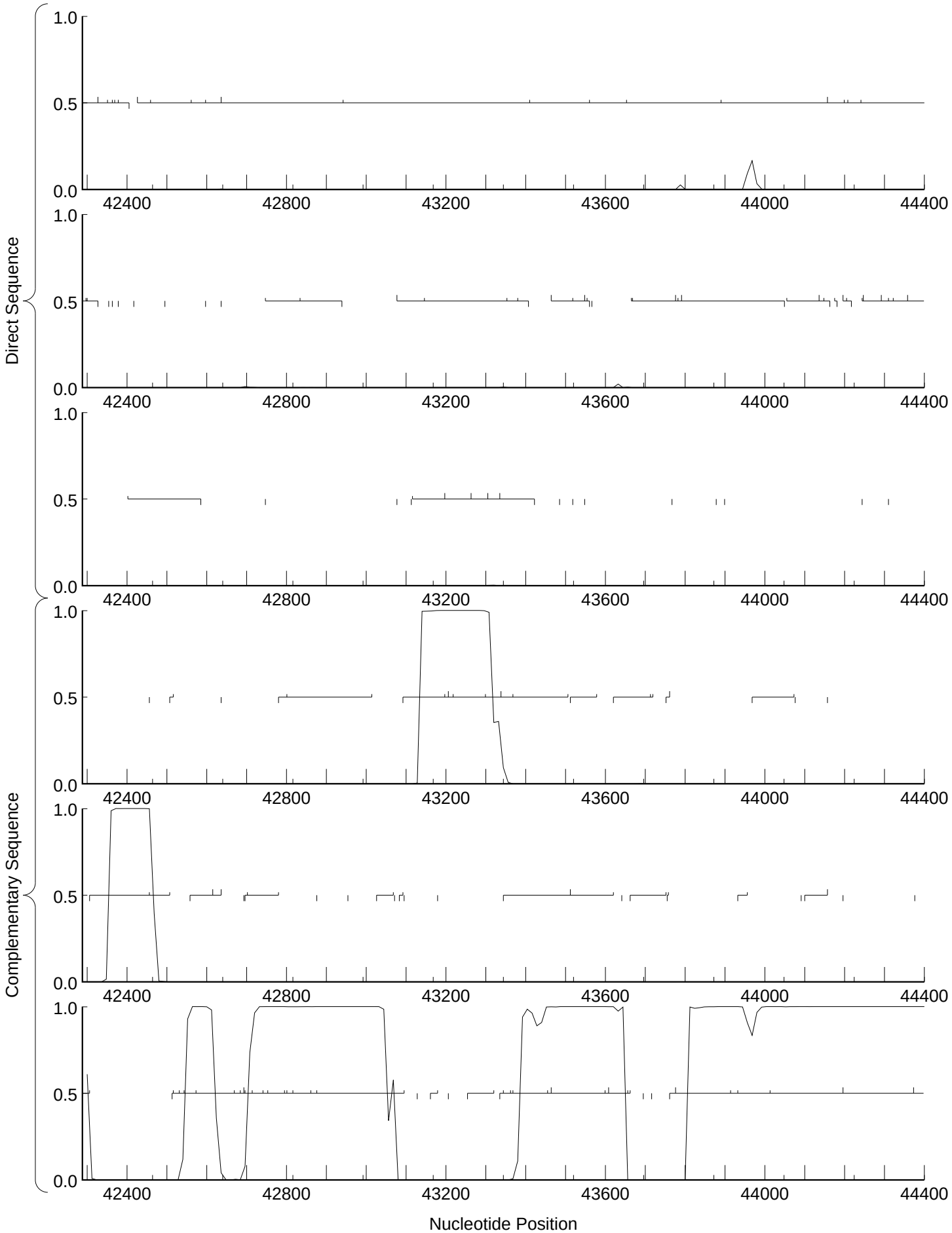

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position


Direct Sequence
Complementary Sequence
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0.5
0.0
36000
36400
36800
37200
37600
38000
Nucleotide Position

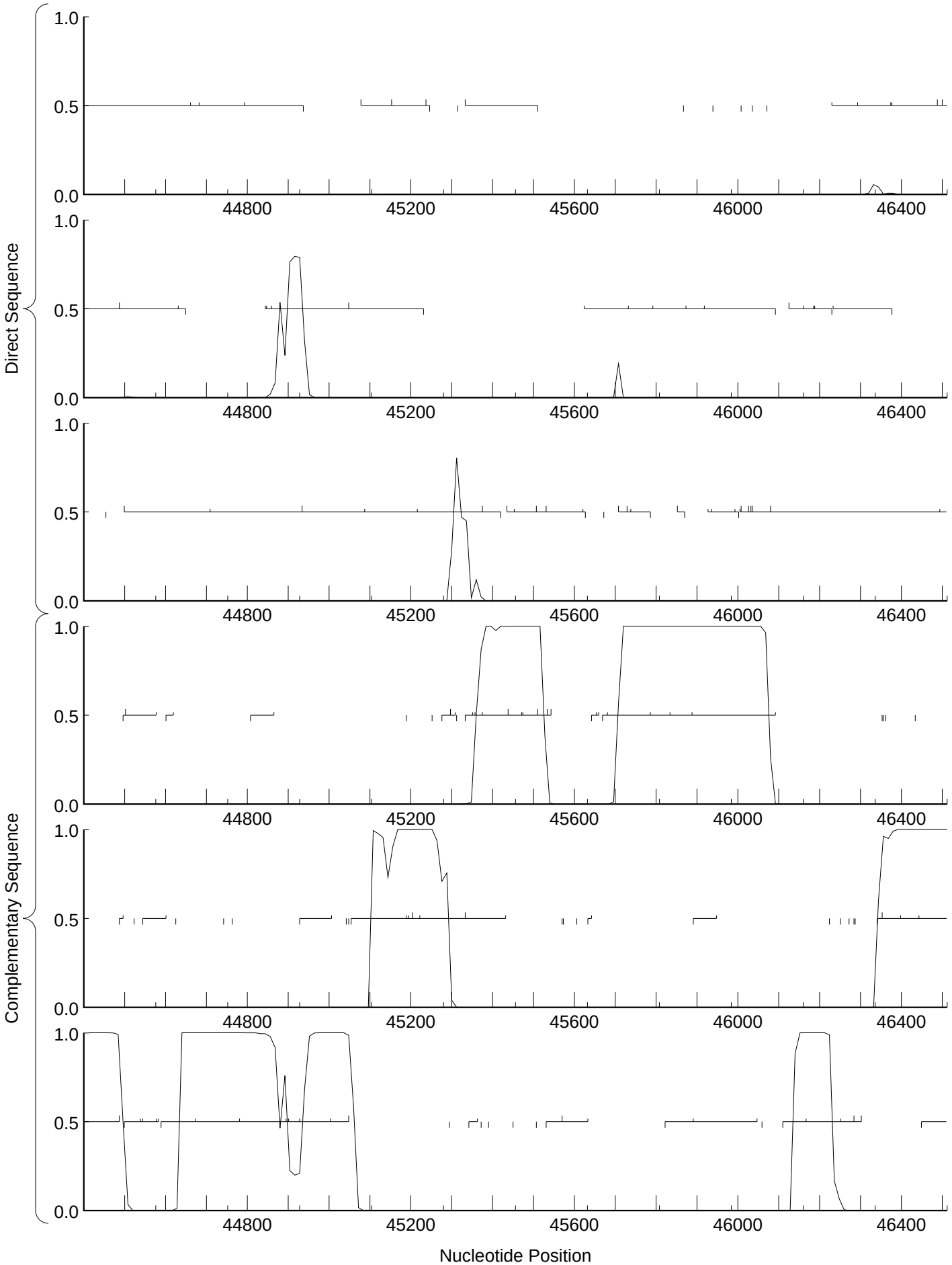

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

omplete sequence, 50632 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 21/25

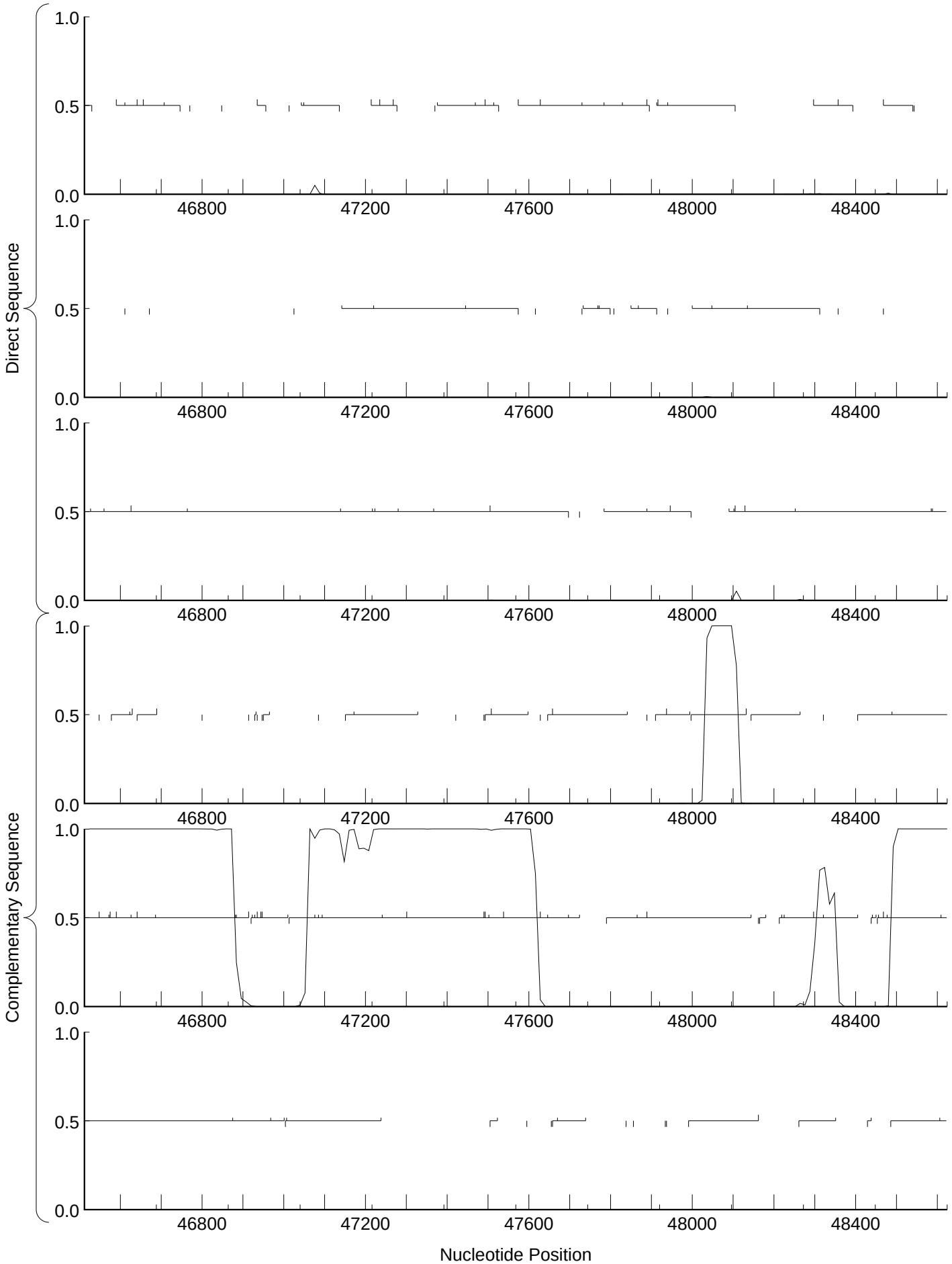



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
42400
42800
43200
43600
44000
44400
Nucleotide Position

omplete sequence, 50632 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 23/25



omplete sequence, 50632 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 24/25



omplete sequence, 50632 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 25/25

