GeneMark

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

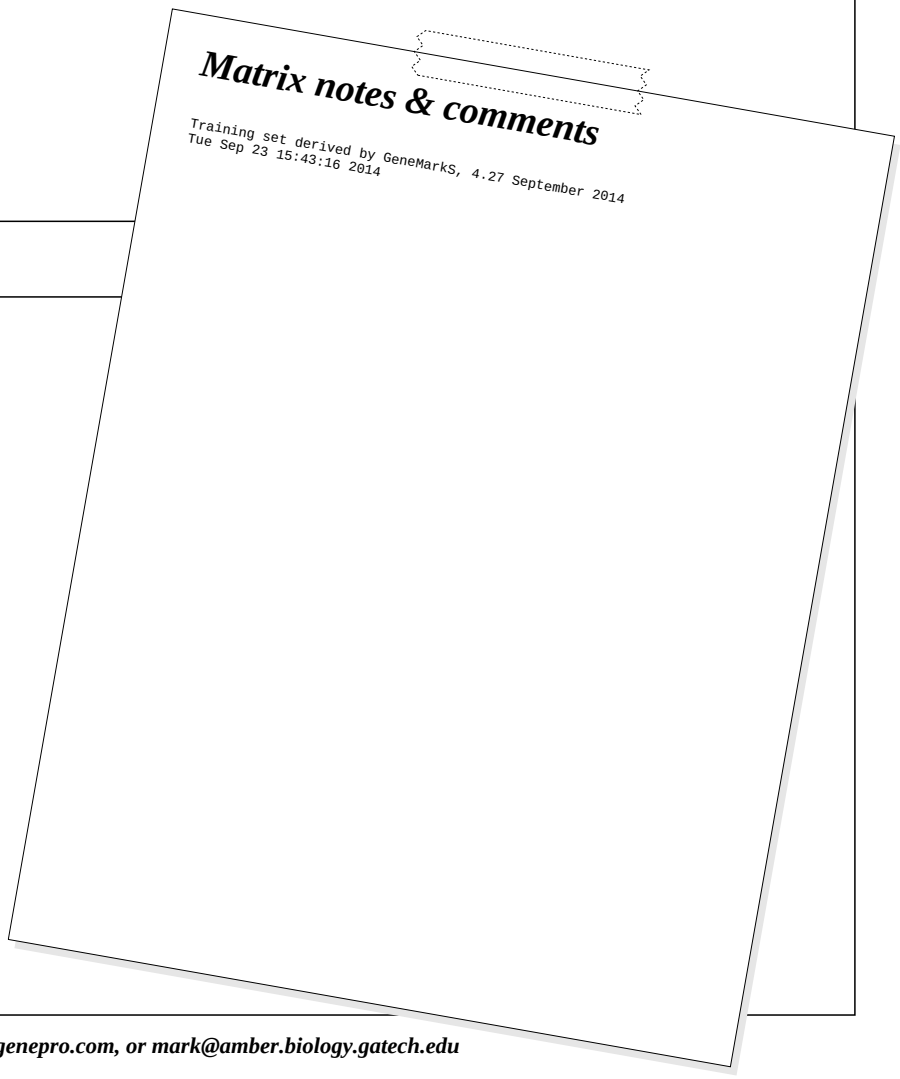
PROGRAM INFORMATION

```
Sequence         :  Streptomyces phage Pablito complete sequence, 49492 bp including 11-base 3' overhang (CGCCG
Analysis Date    :  3/28/24 at 17:38:22
Pages            :  25
Sequence Length  :  49492 bp
GC Content       :   66.41%

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

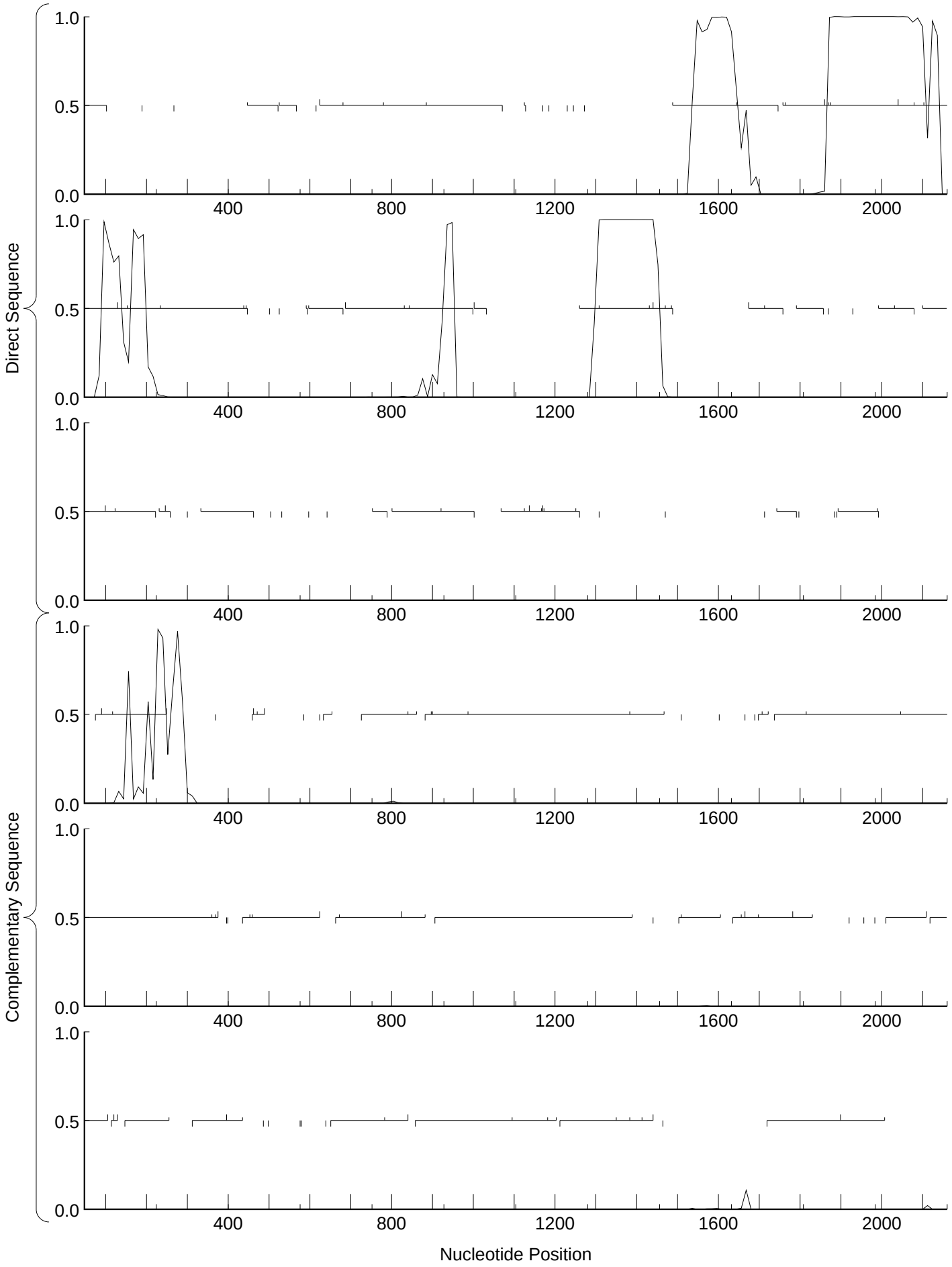
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Matrix     :  Streptomyces_scabiei_87_22
Author     :  -

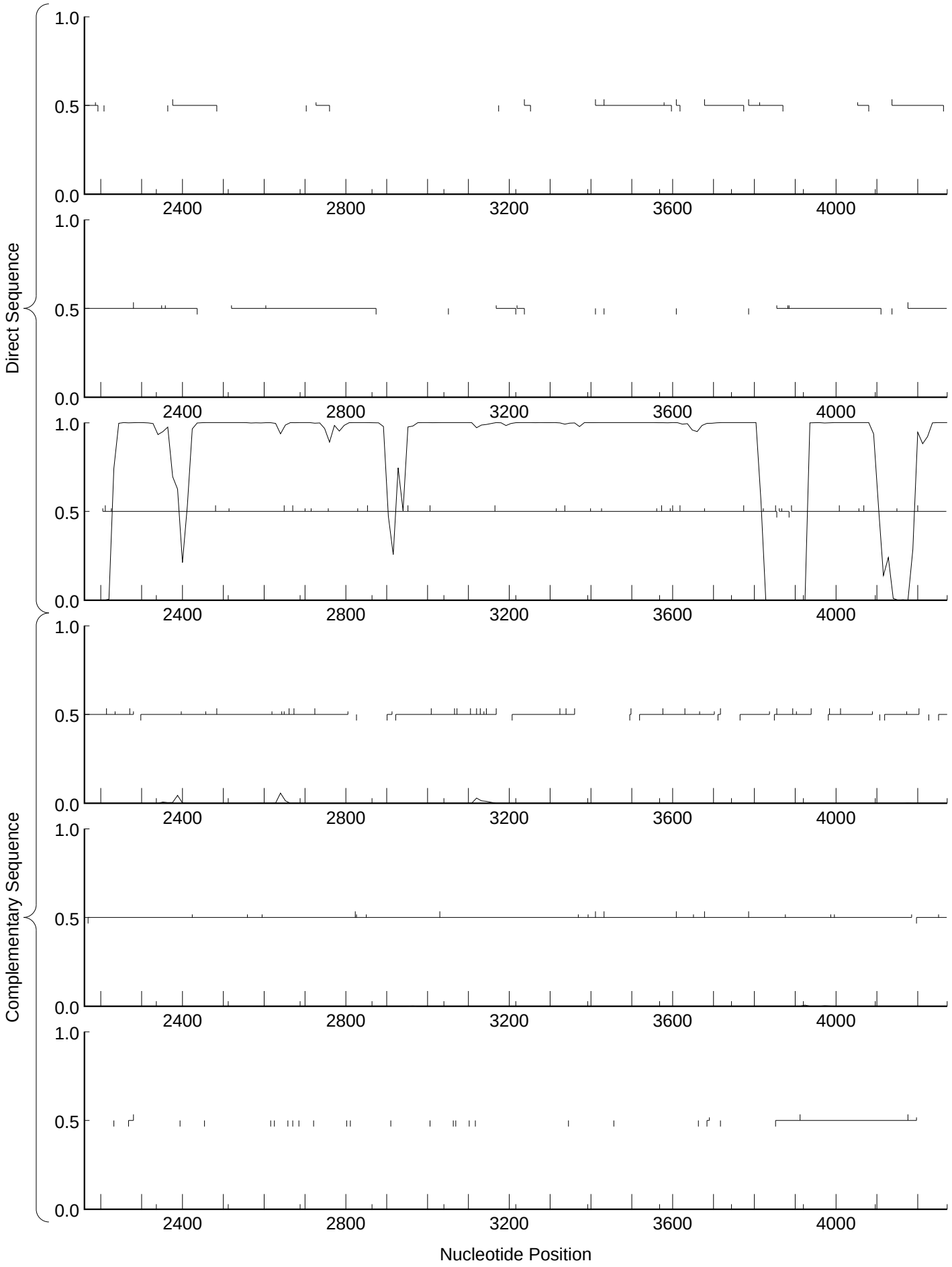
Order      :  4
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  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

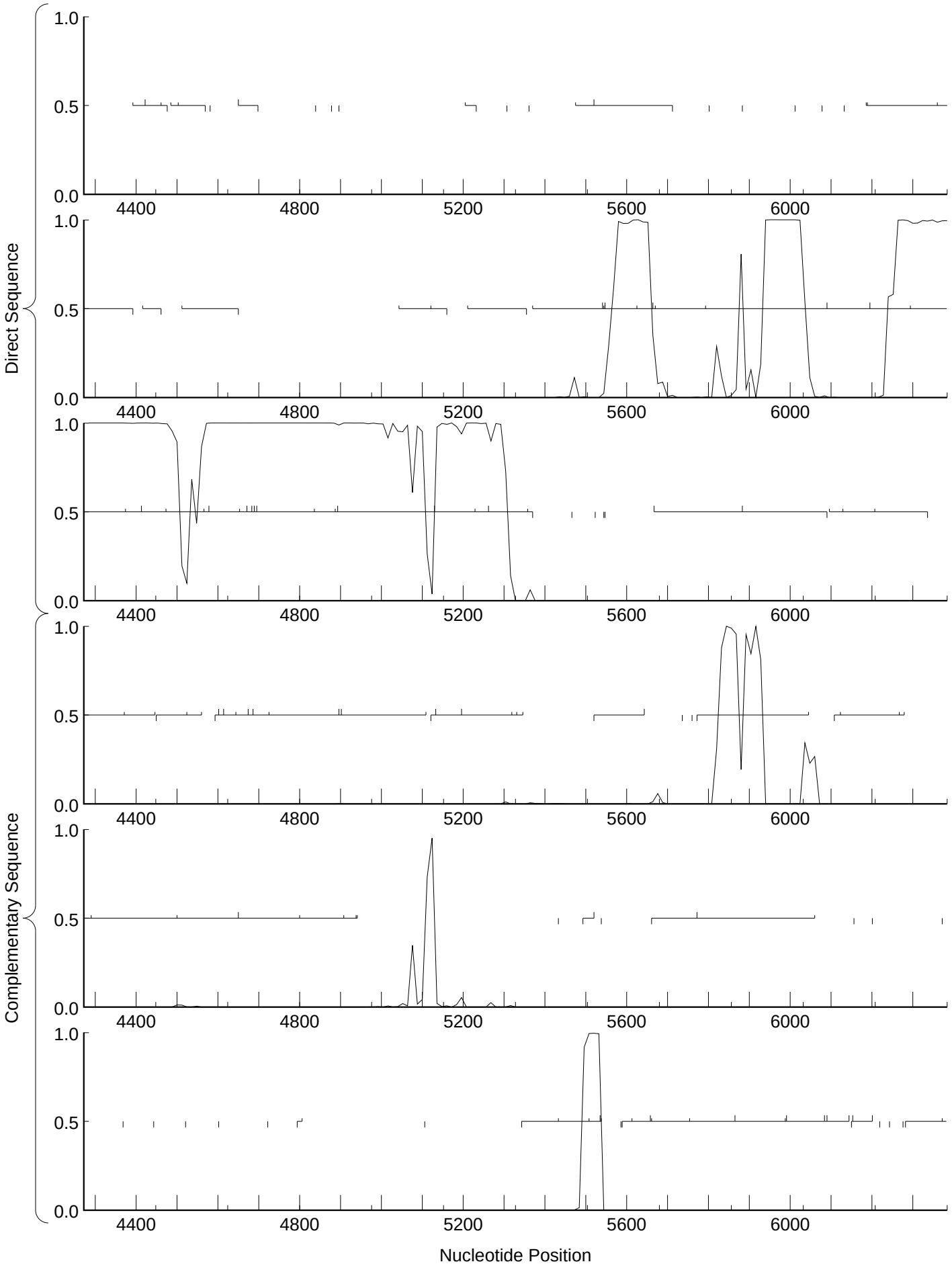

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

complete sequence, 49492 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 2/25

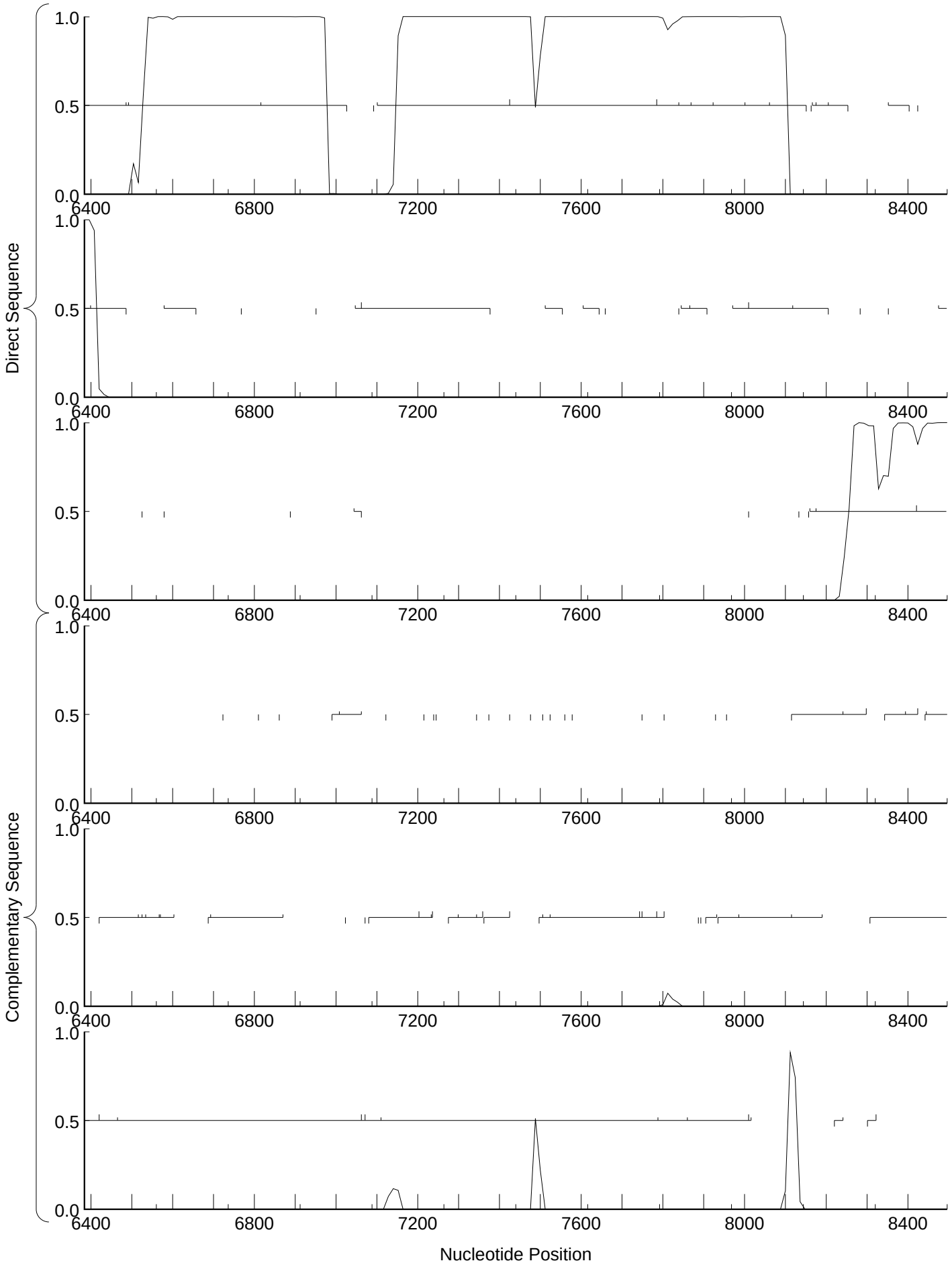



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
2400
2800
3200
3600
4000
Nucleotide Position

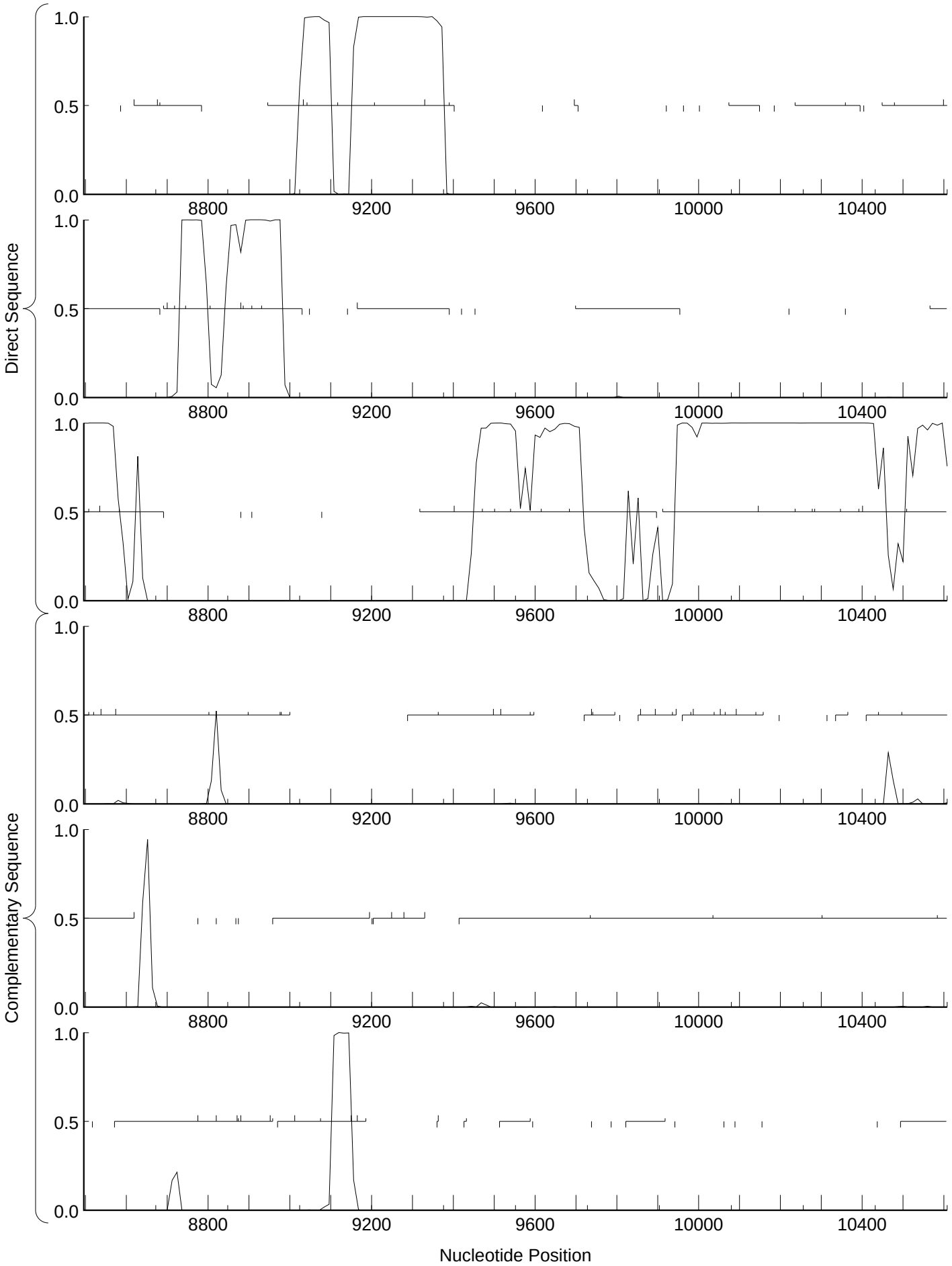
complete sequence, 49492 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 4/25



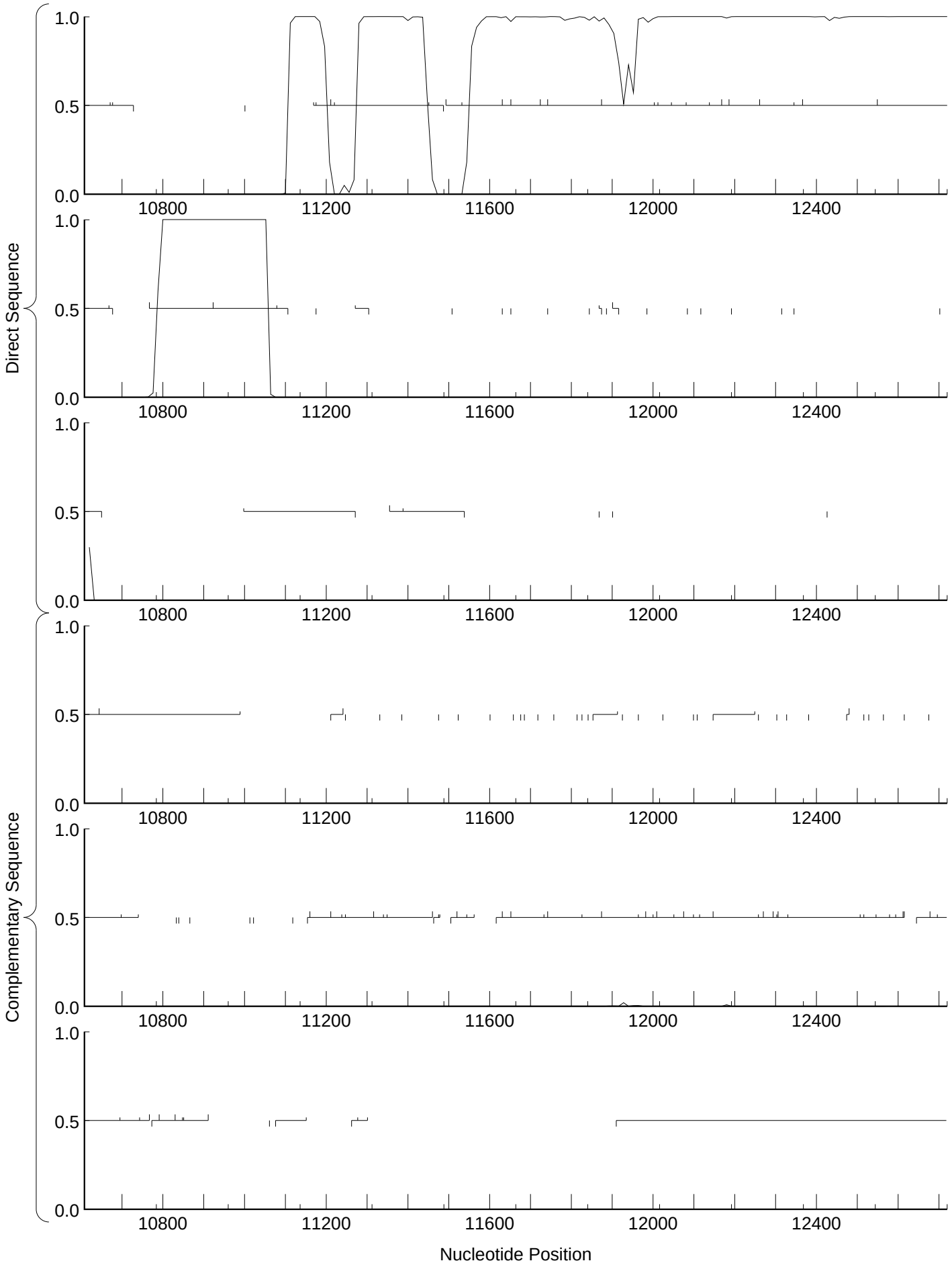
complete sequence, 49492 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 5/25



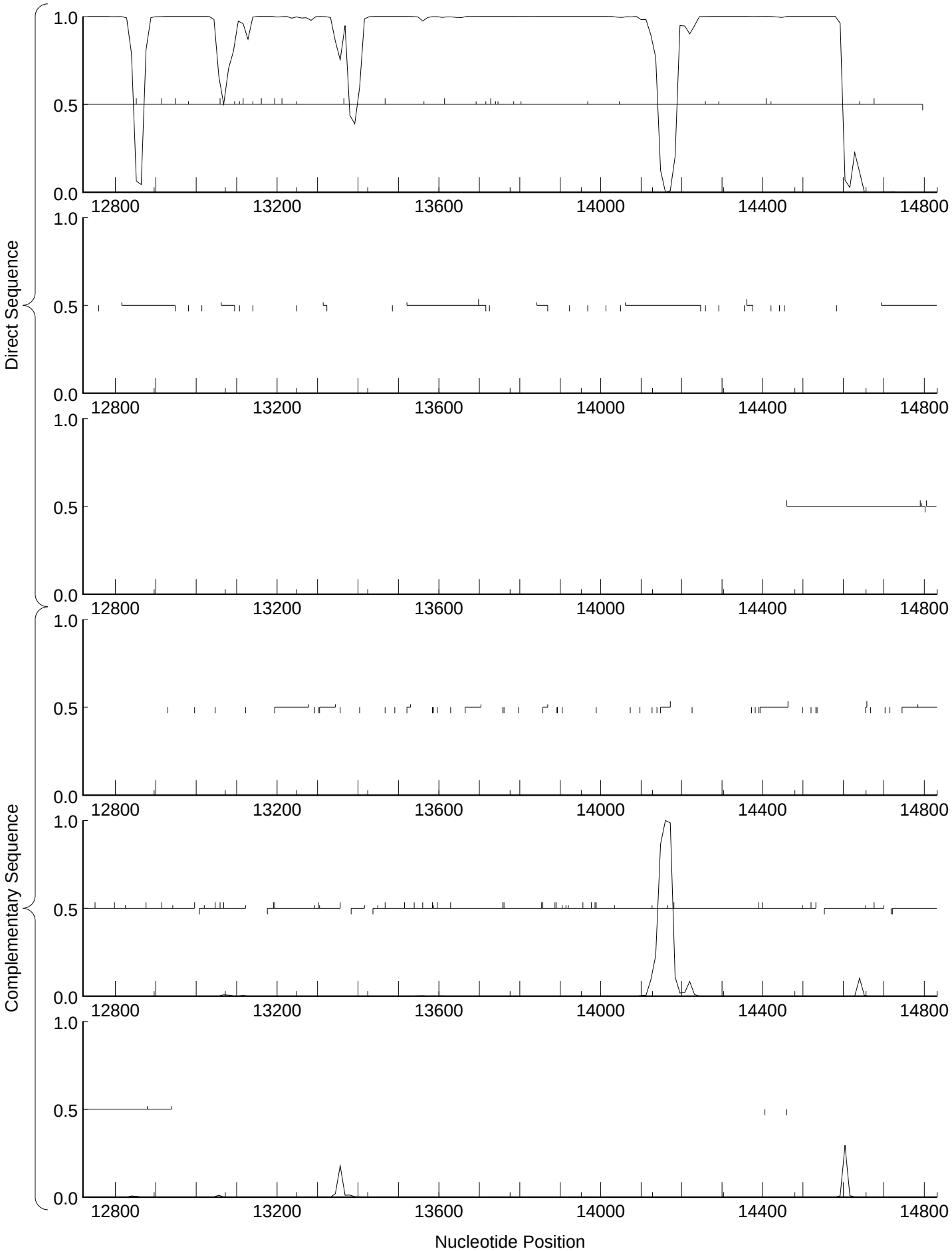
complete sequence, 49492 bp including 11-base 3' overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 6/25



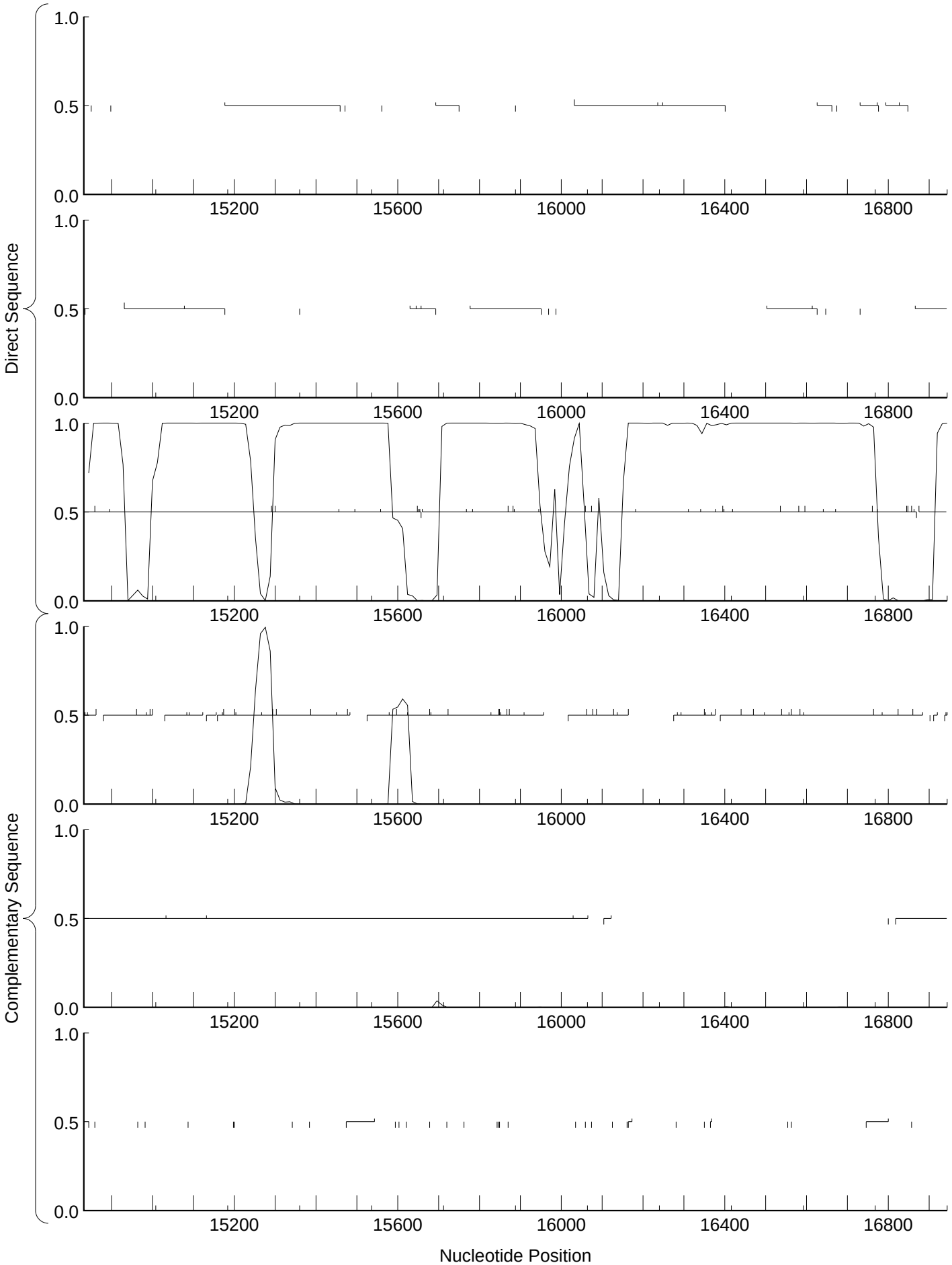
complete sequence, 49492 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 7/25

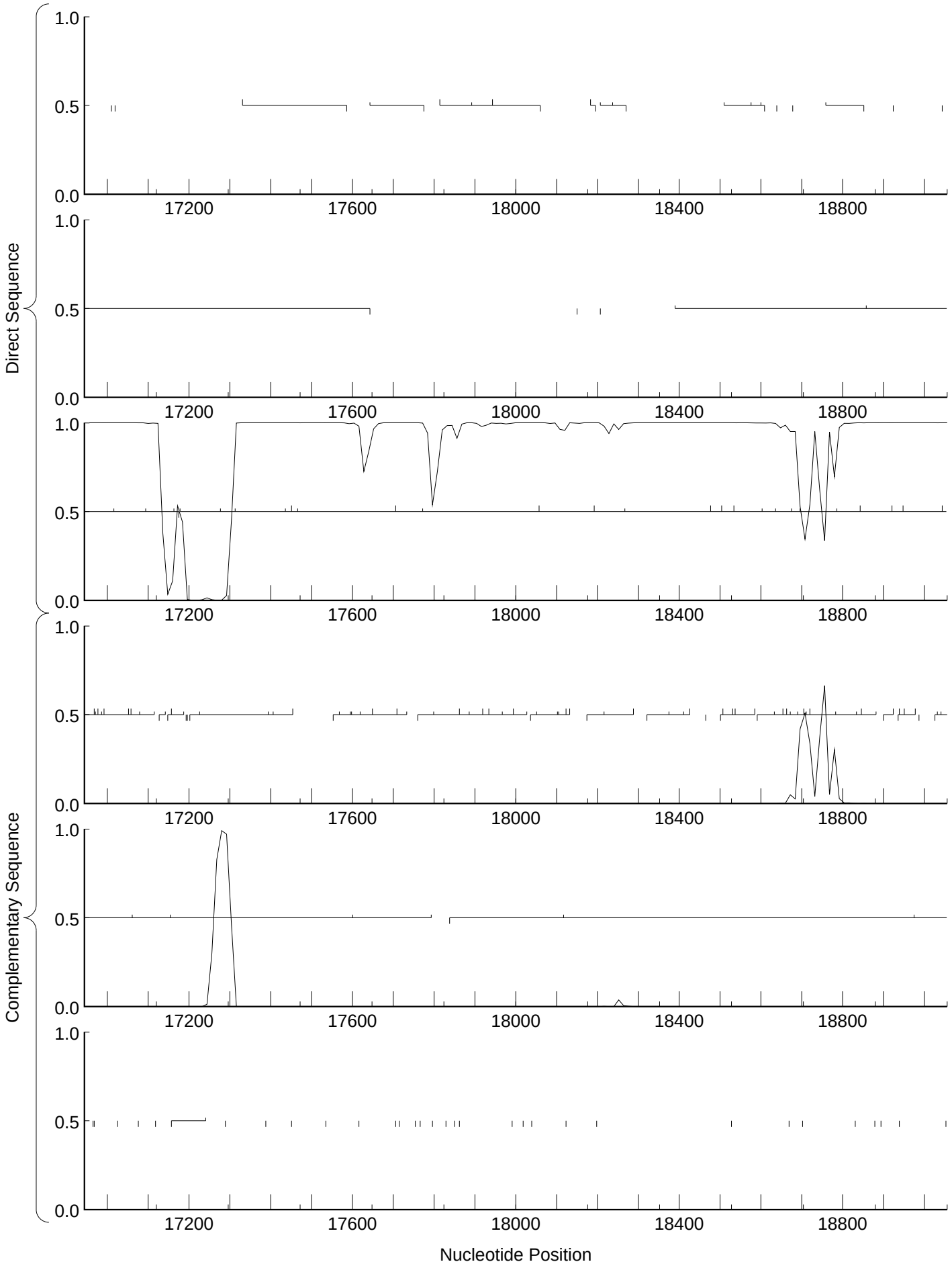


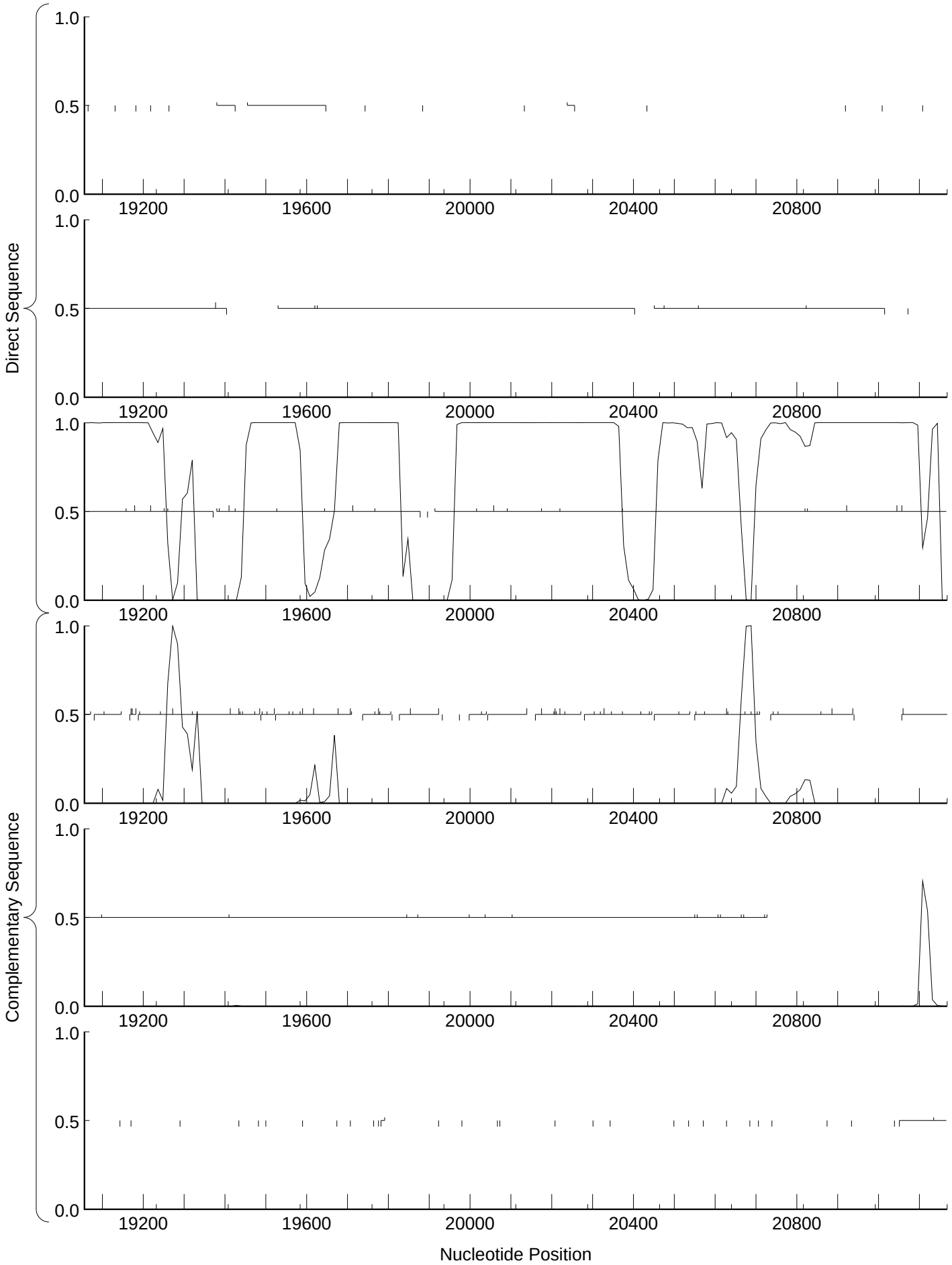
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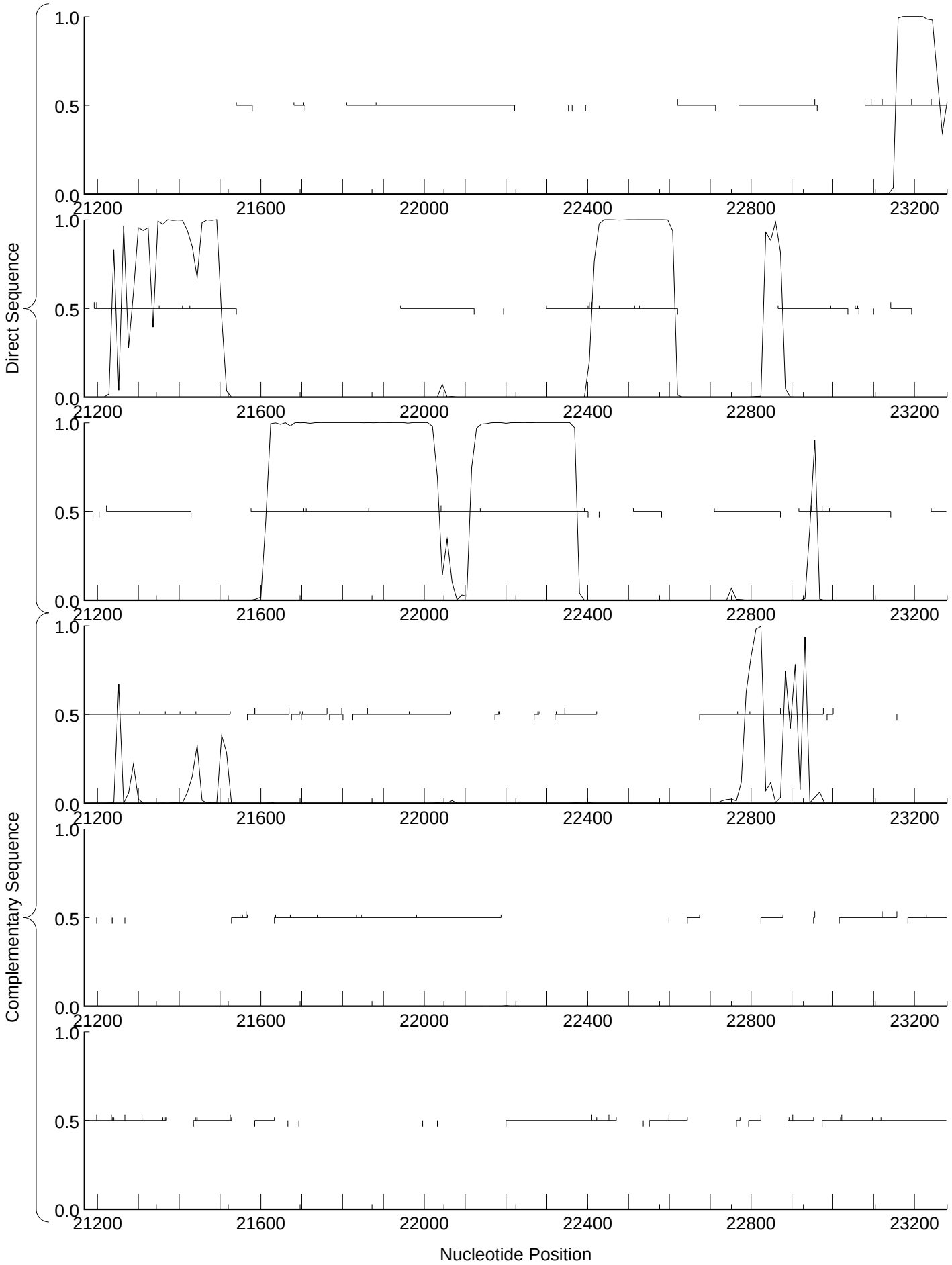
complete sequence, 49492 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 9/25



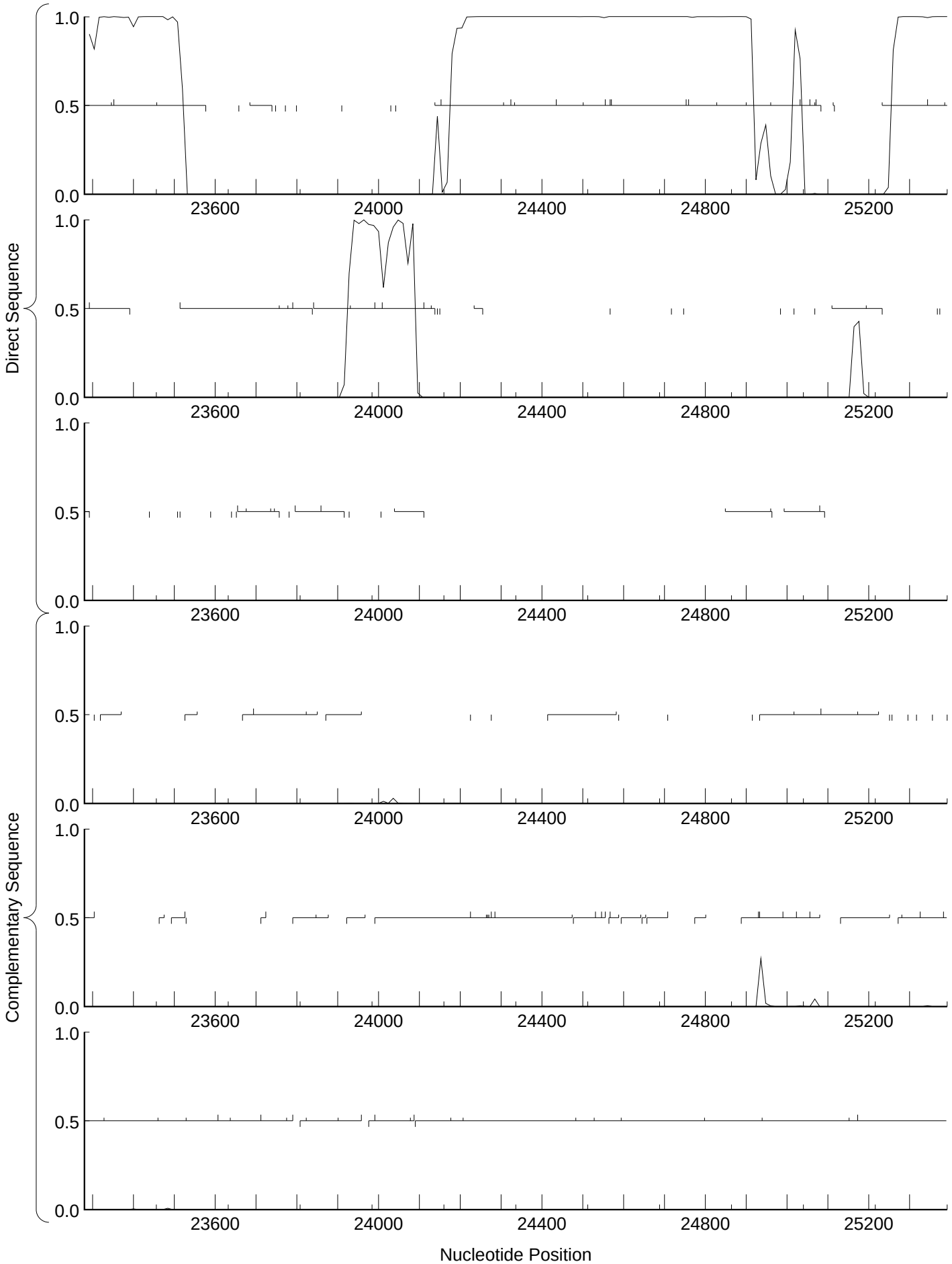

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
17200
17600
18000
18400
18800
Nucleotide Position


Direct Sequence
Complementary Sequence
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0.5
0.0
19200
19600
20000
20400
20800
Nucleotide Position

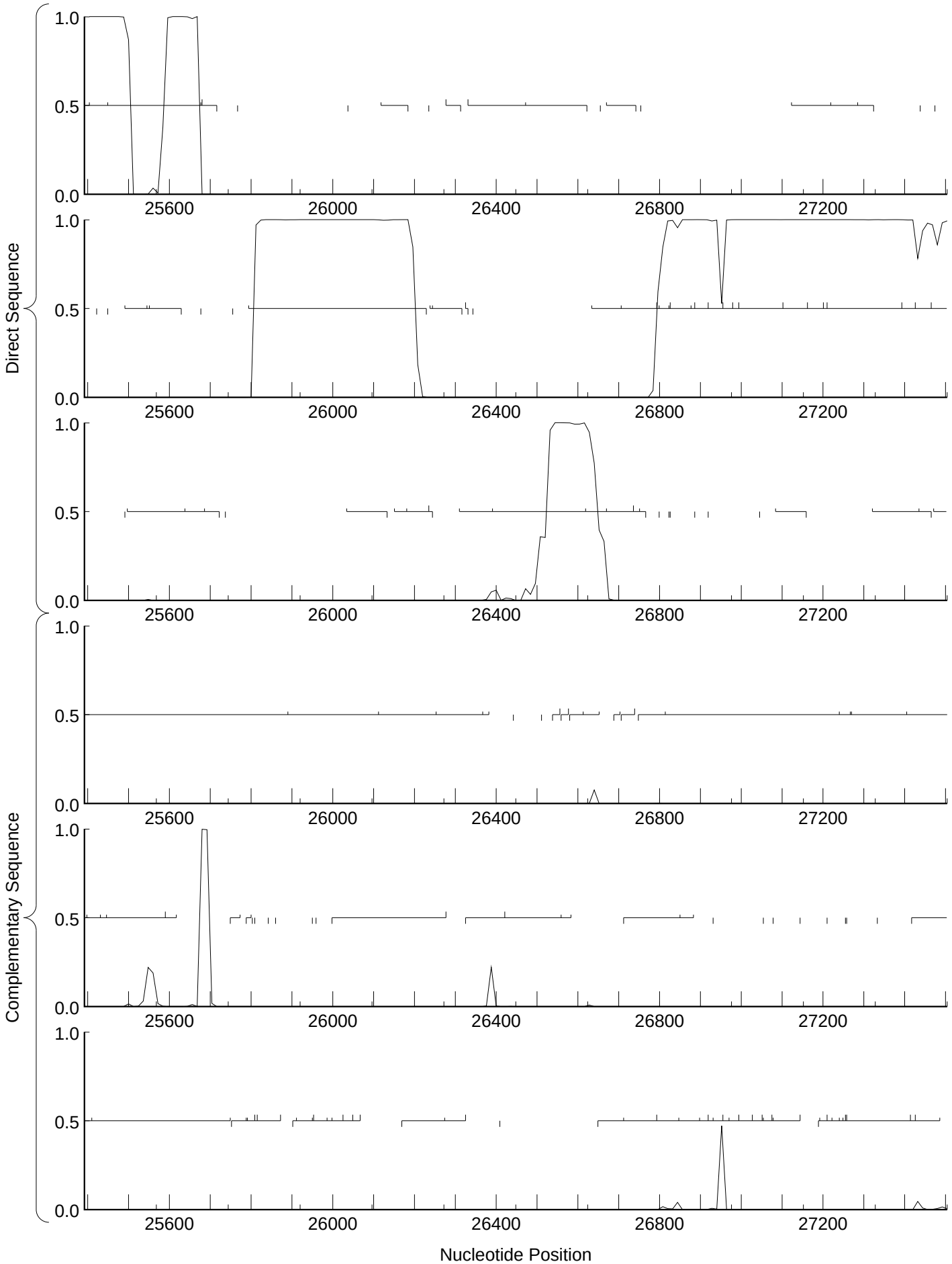
omplete sequence, 49492 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 12/25

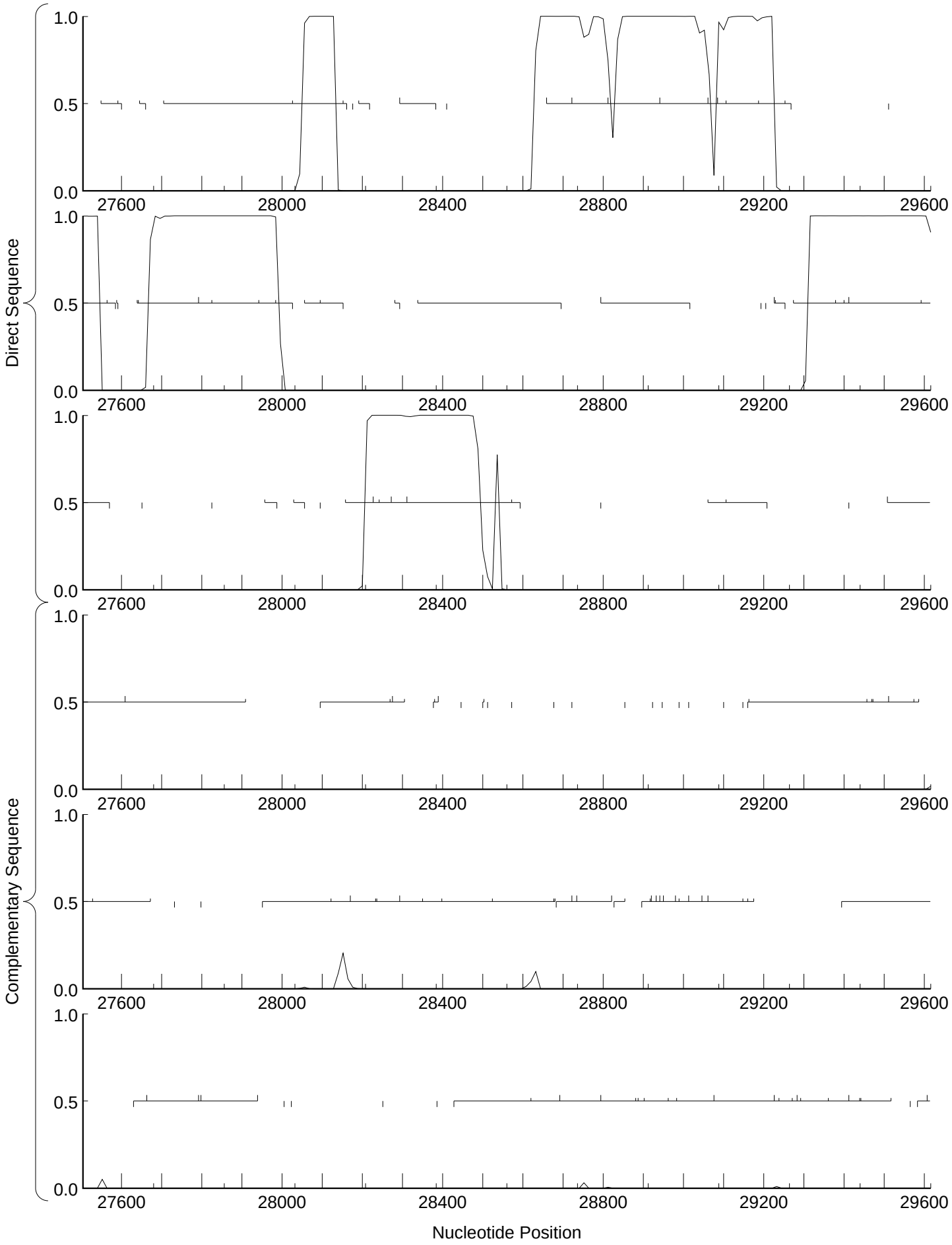


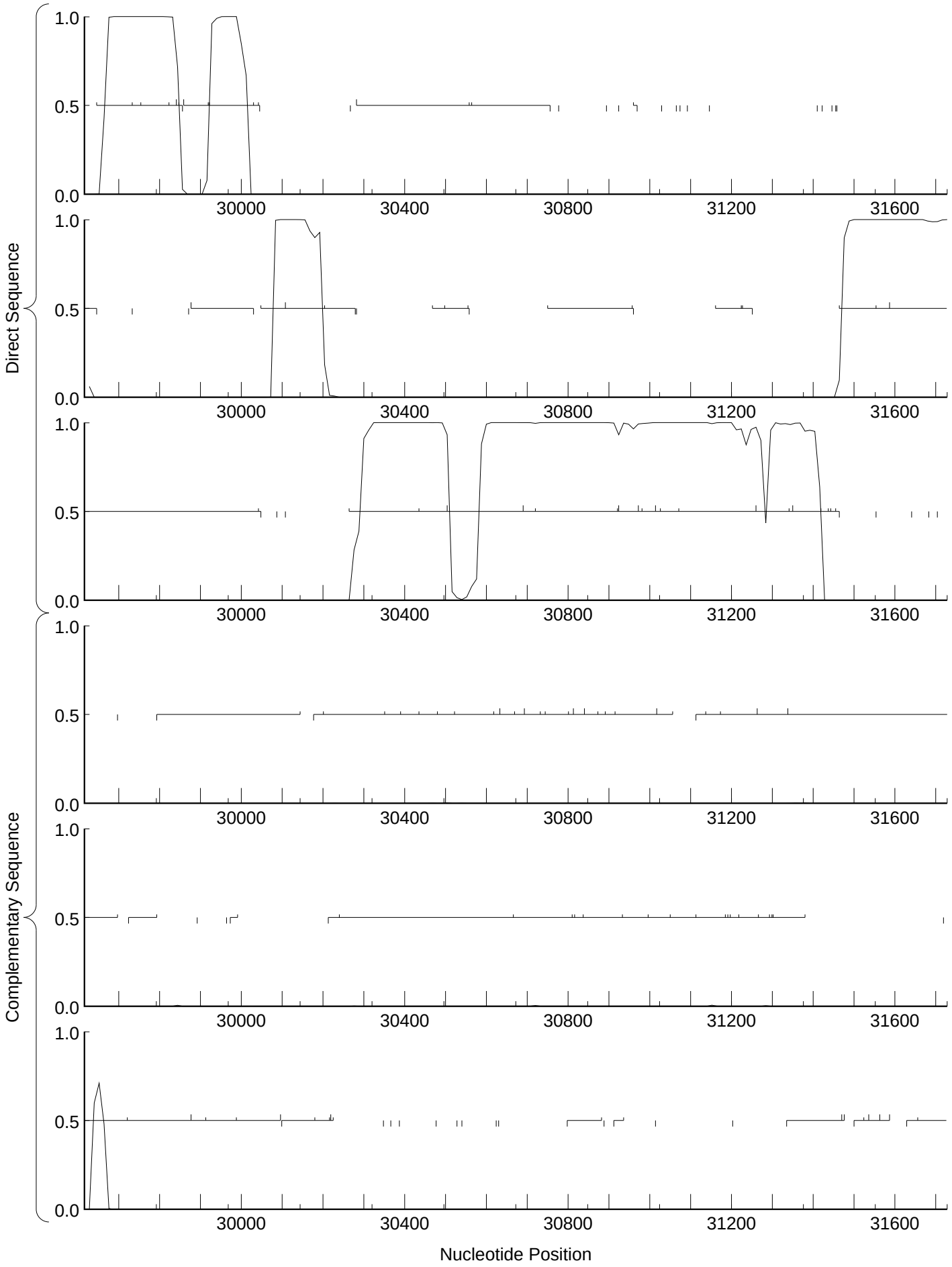
omplete sequence, 49492 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 13/25



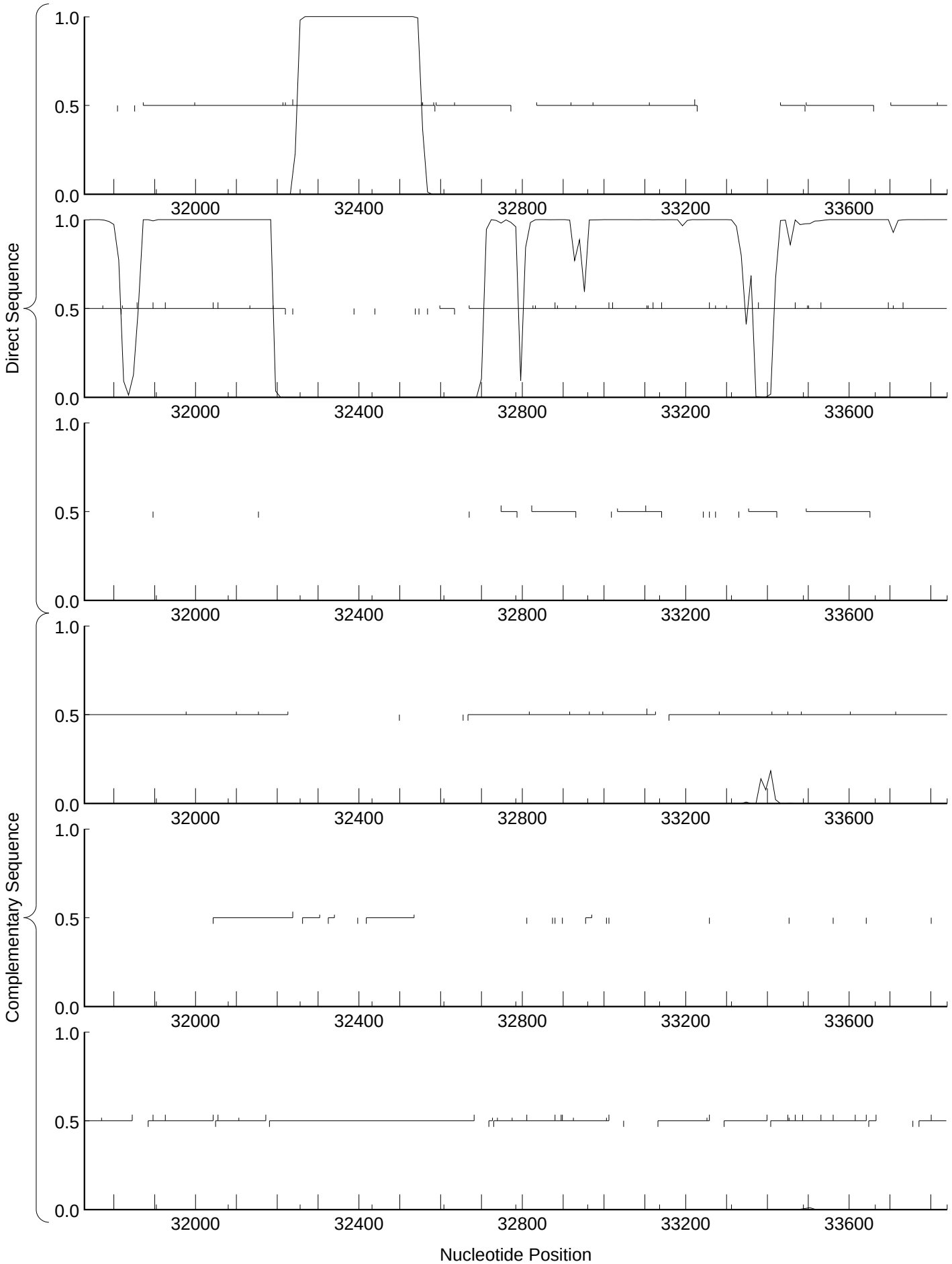
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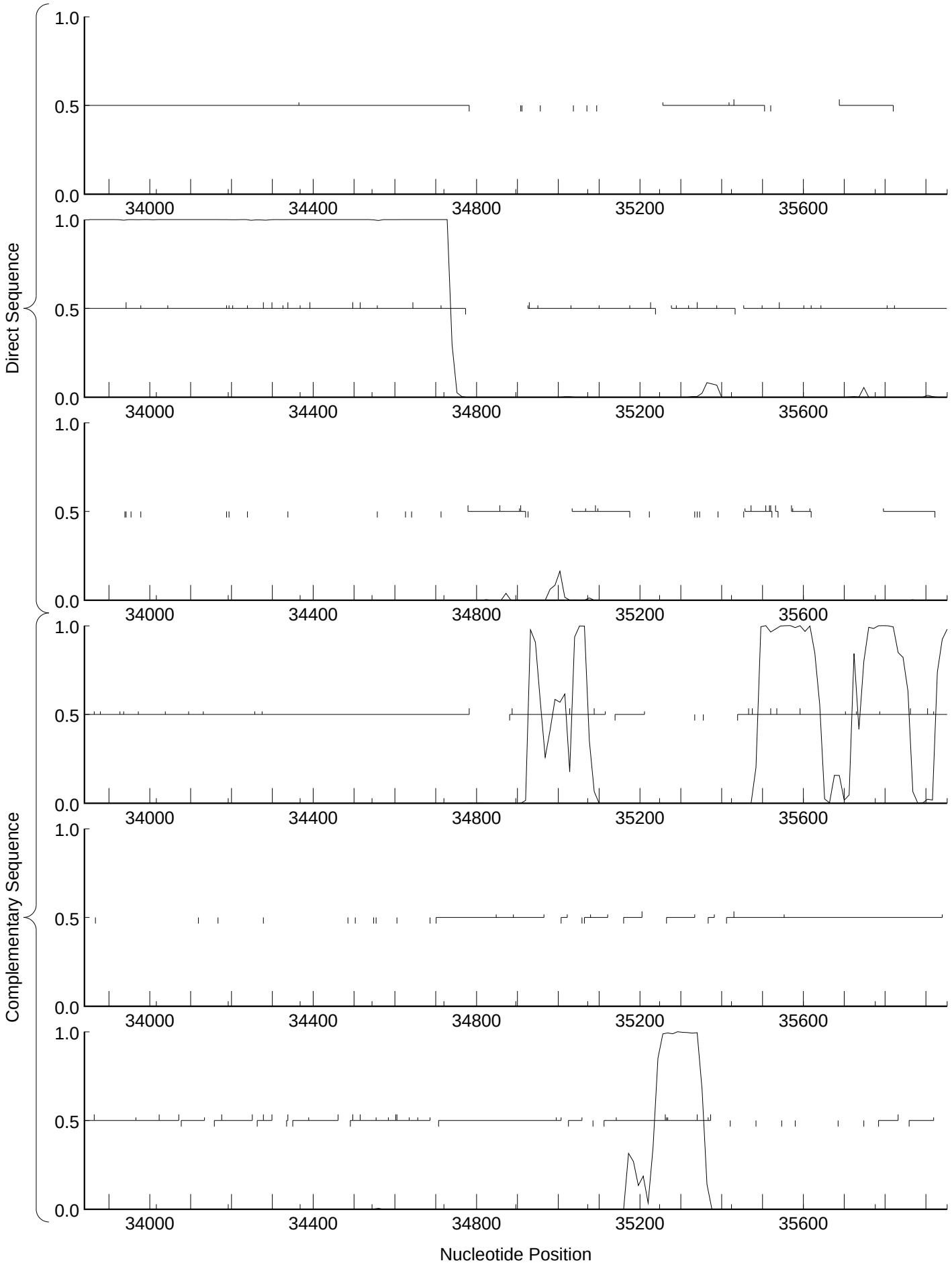



Direct Sequence
Complementary Sequence
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0.5
0.0
27600
28000
28400
28800
29200
29600
Nucleotide Position

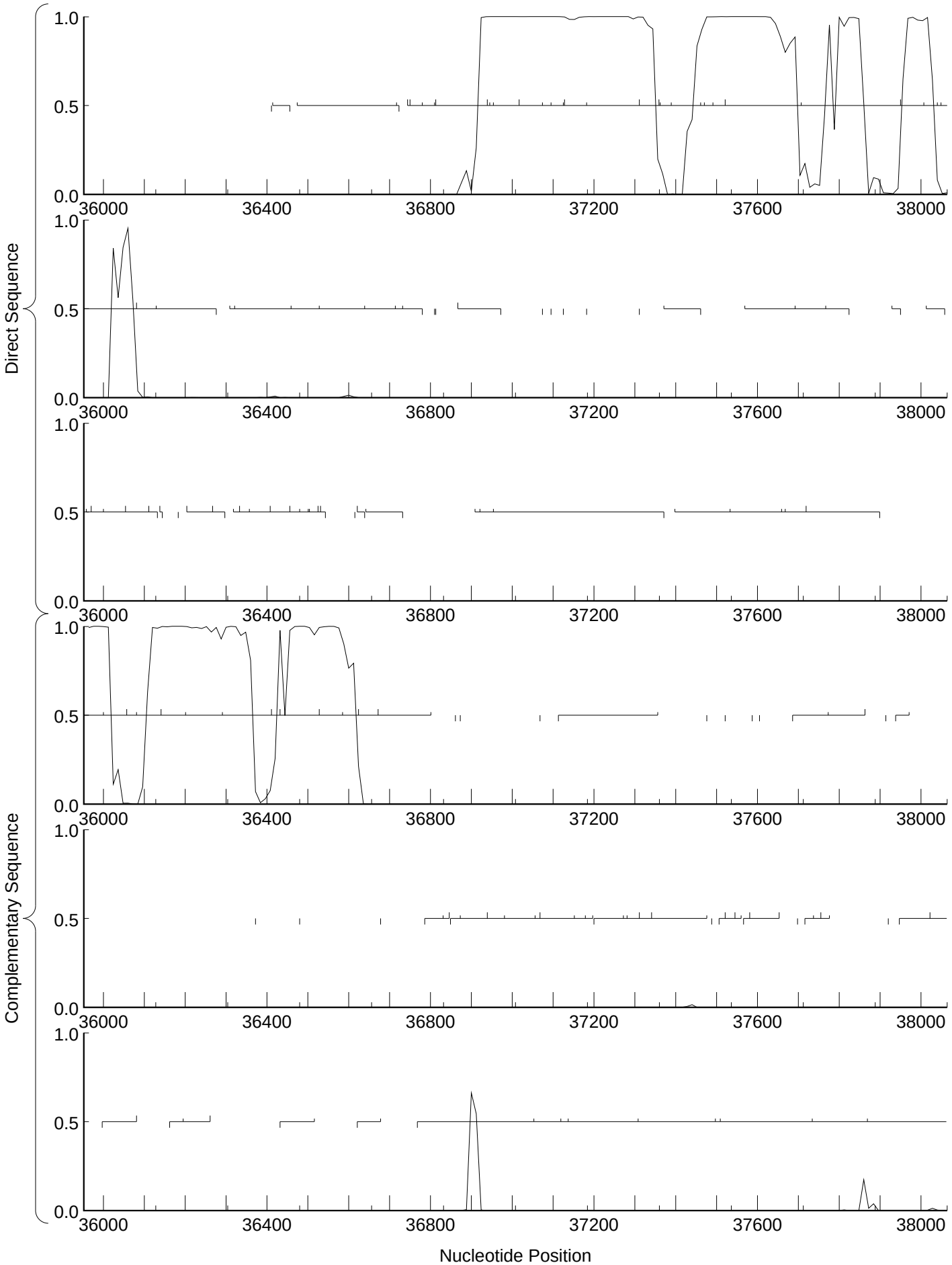

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Complementary Sequence
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Nucleotide Position

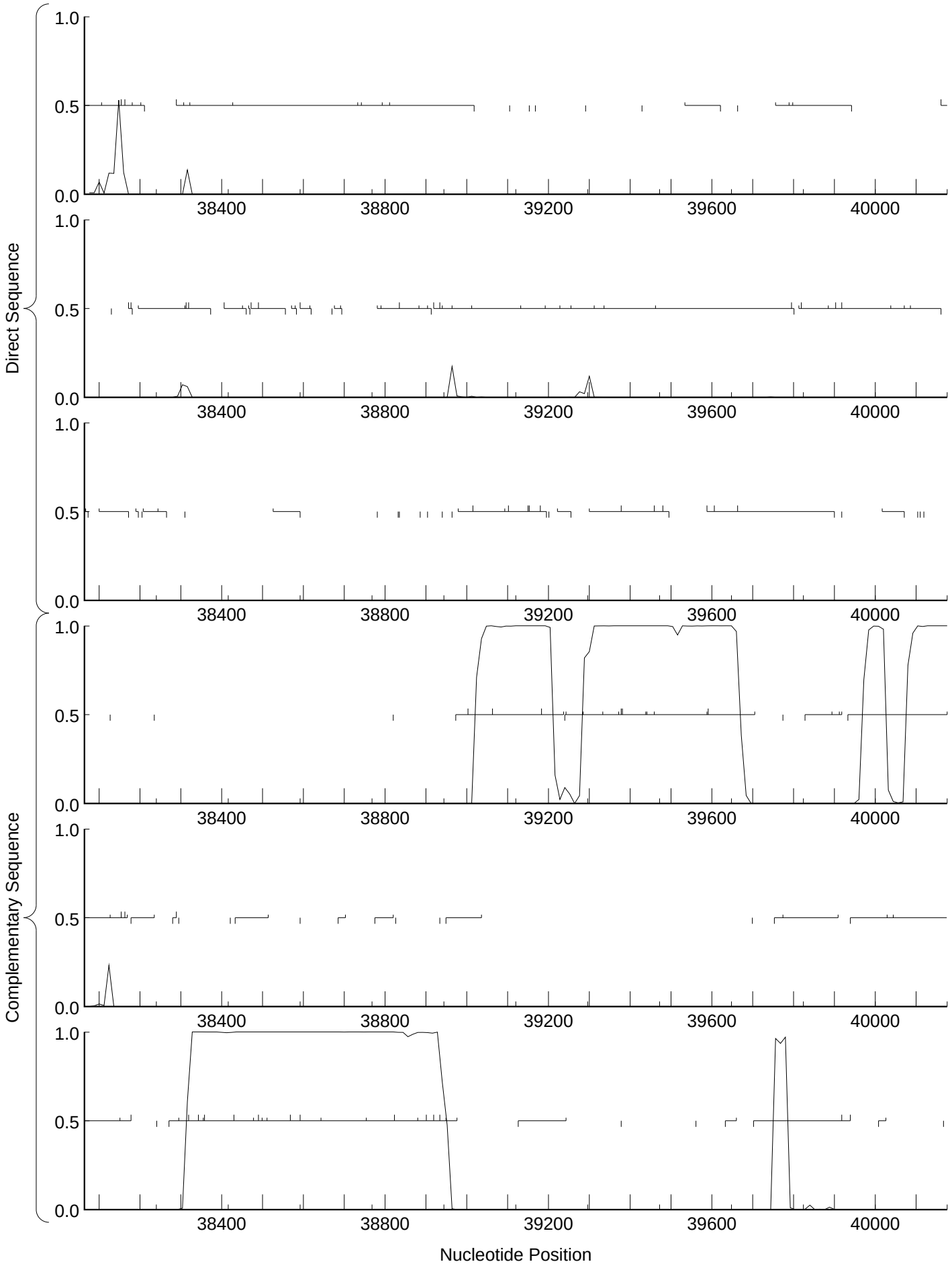
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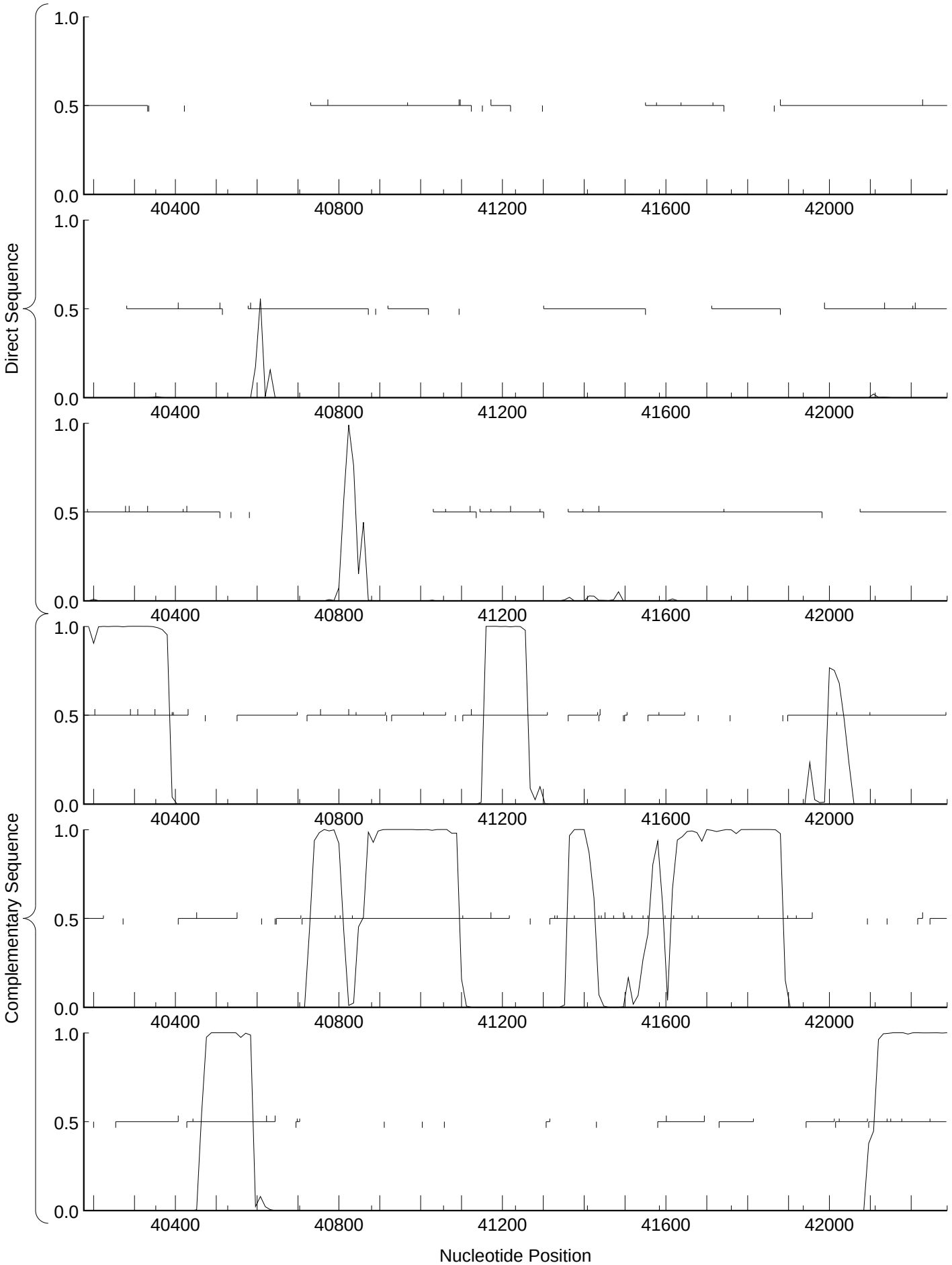

Direct Sequence
Complementary Sequence
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0.0
34000
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34800
35200
35600
Nucleotide Position

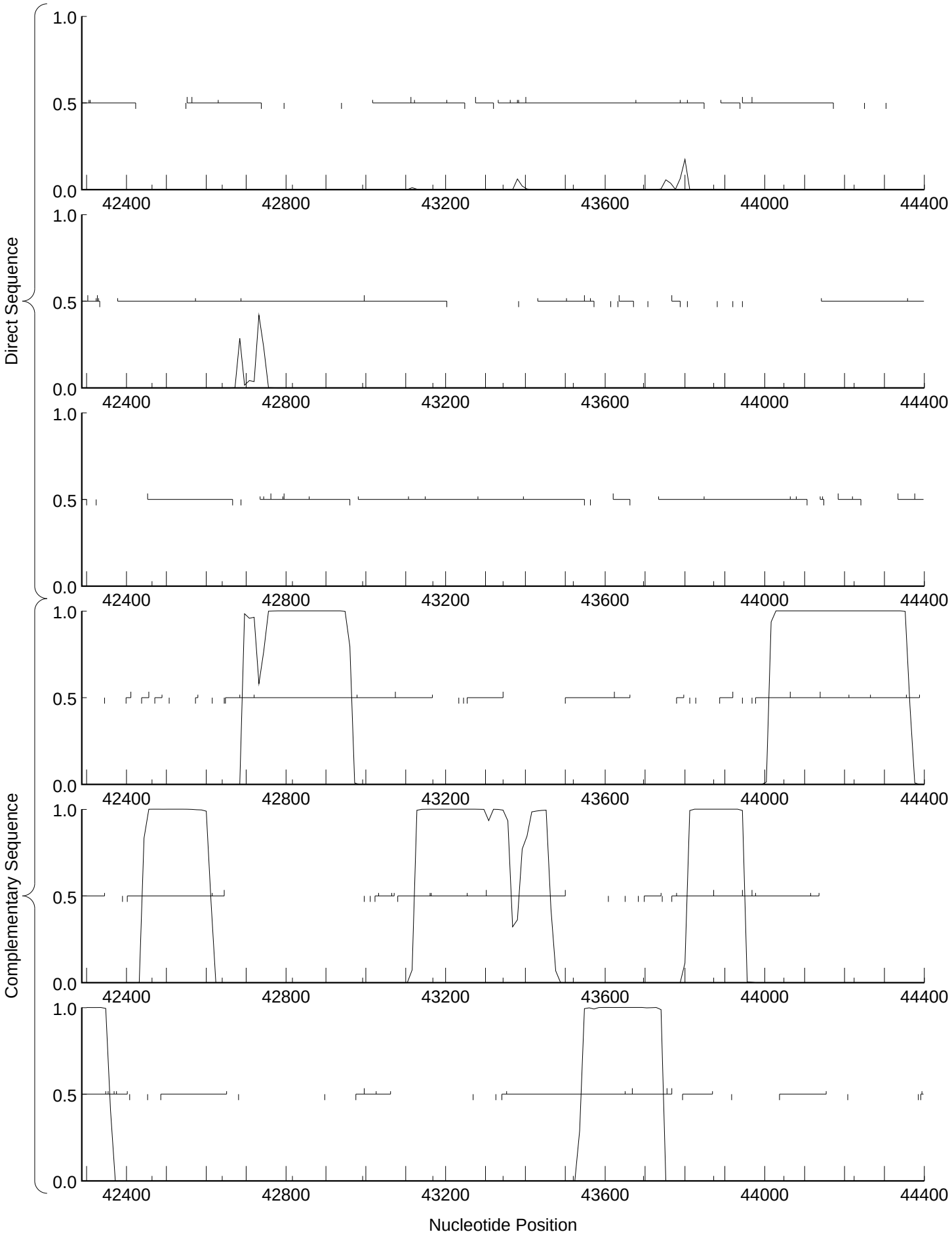
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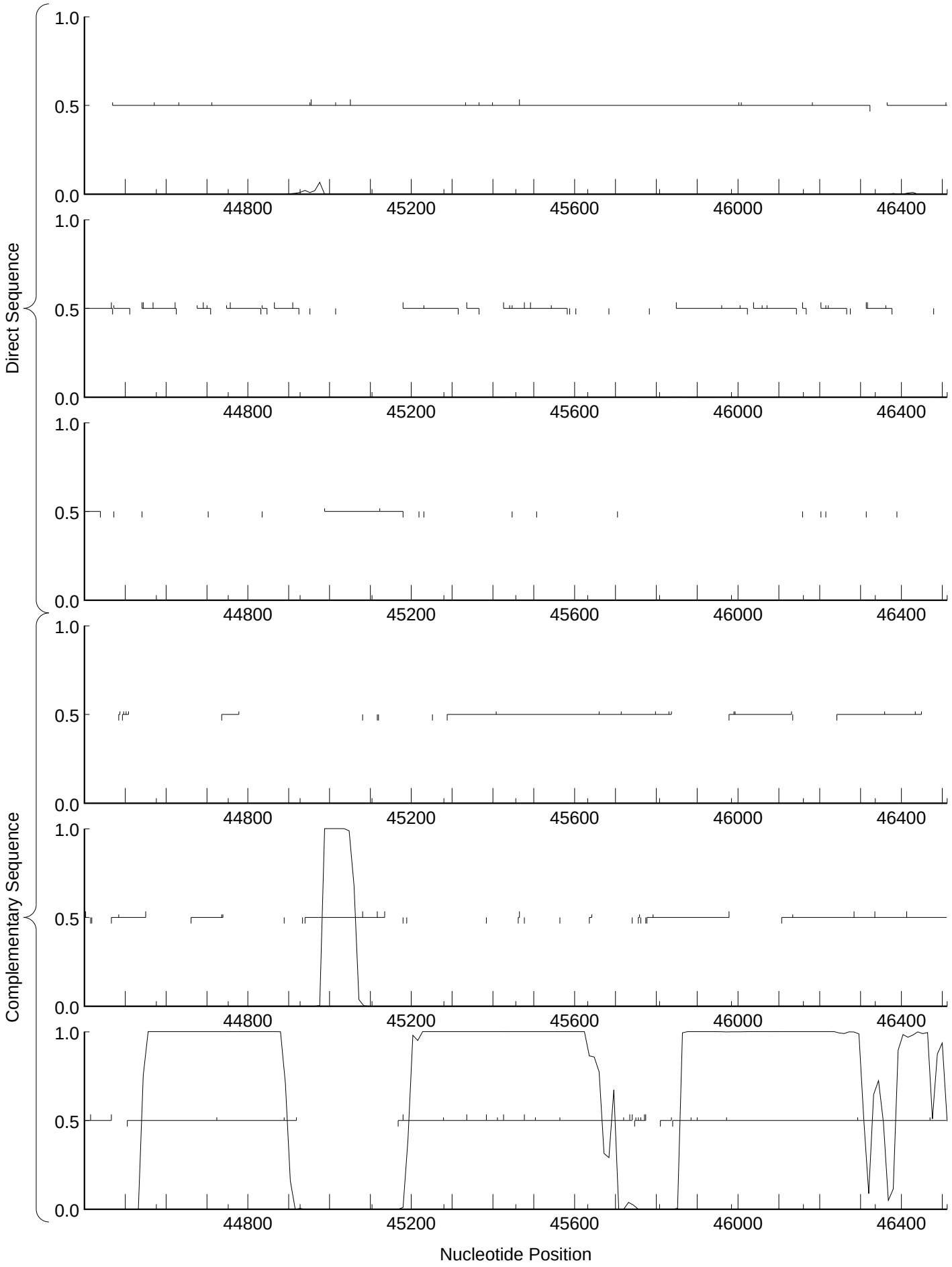

Direct Sequence
Complementary Sequence
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0.0
38400
38800
39200
39600
40000
Nucleotide Position

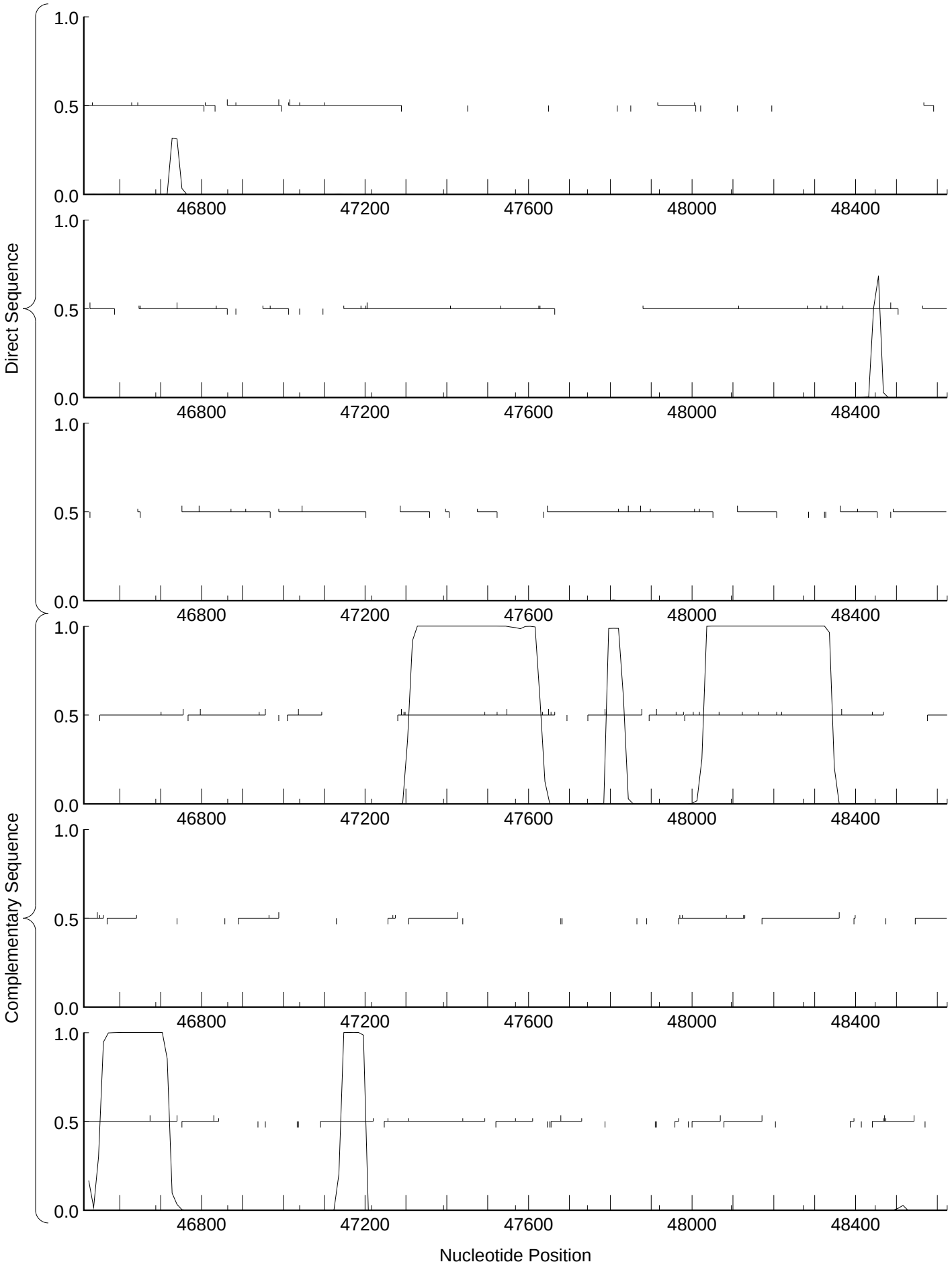
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Direct Sequence
Complementary Sequence
1.0
0.5
0.0
42400
42800
43200
43600
44000
44400
Nucleotide Position

omplete sequence, 49492 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 23/25




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
46800
47200
47600
48000
48400
Nucleotide Position

omplete sequence, 49492 bp including 11-base 3’ overhang (CGCCGTGTCTT), Cluster BD2, Order 4, Window 96, Step 12, 25/25

