# GeneMark

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

## PROGRAM INFORMATION

```
Sequence         :  Streptomyces phage Rainydai complete sequence, 57623 bp including 9-base 3' overhang (CGCCC
Analysis Date    :  3/28/24 at 4:44:19
Pages            :  29
Sequence Length  :  57623 bp
GC Content       :   58.08%

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

PS-Version       :  1.2

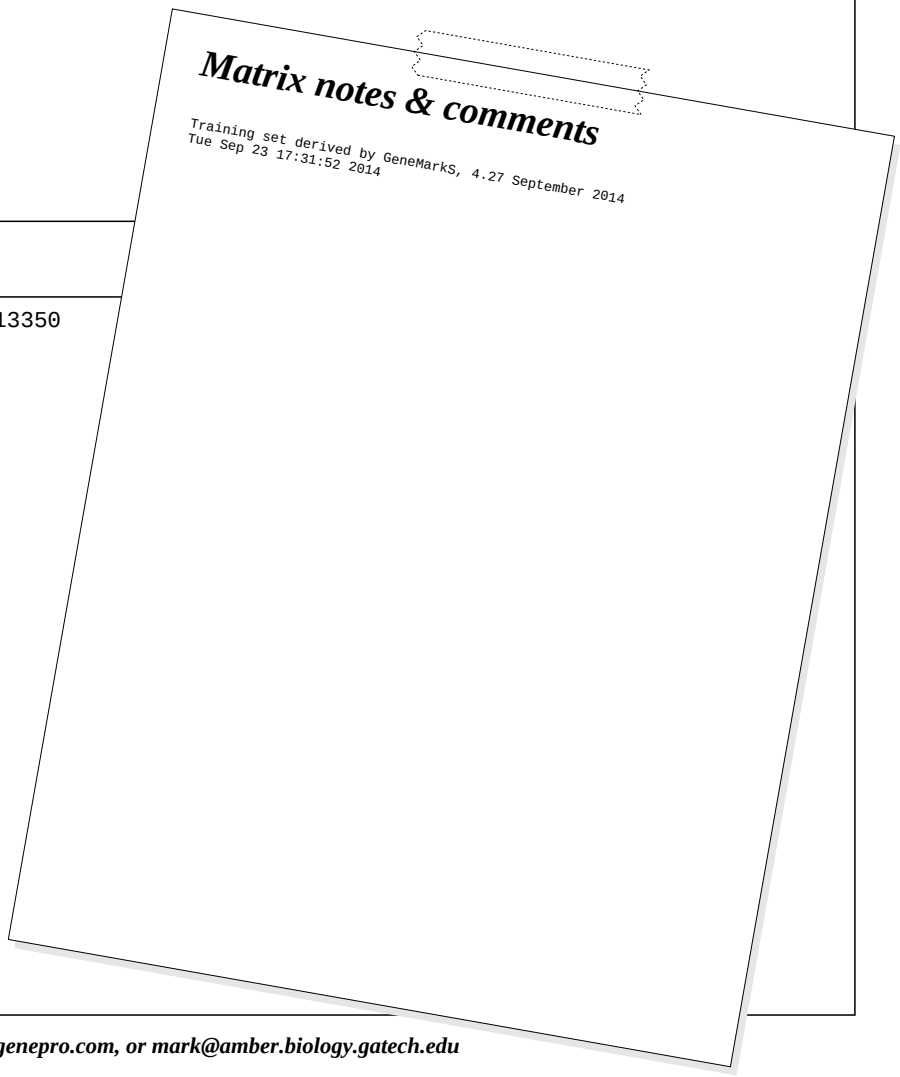
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
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## MATRIX INFORMATION

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Author     :  -

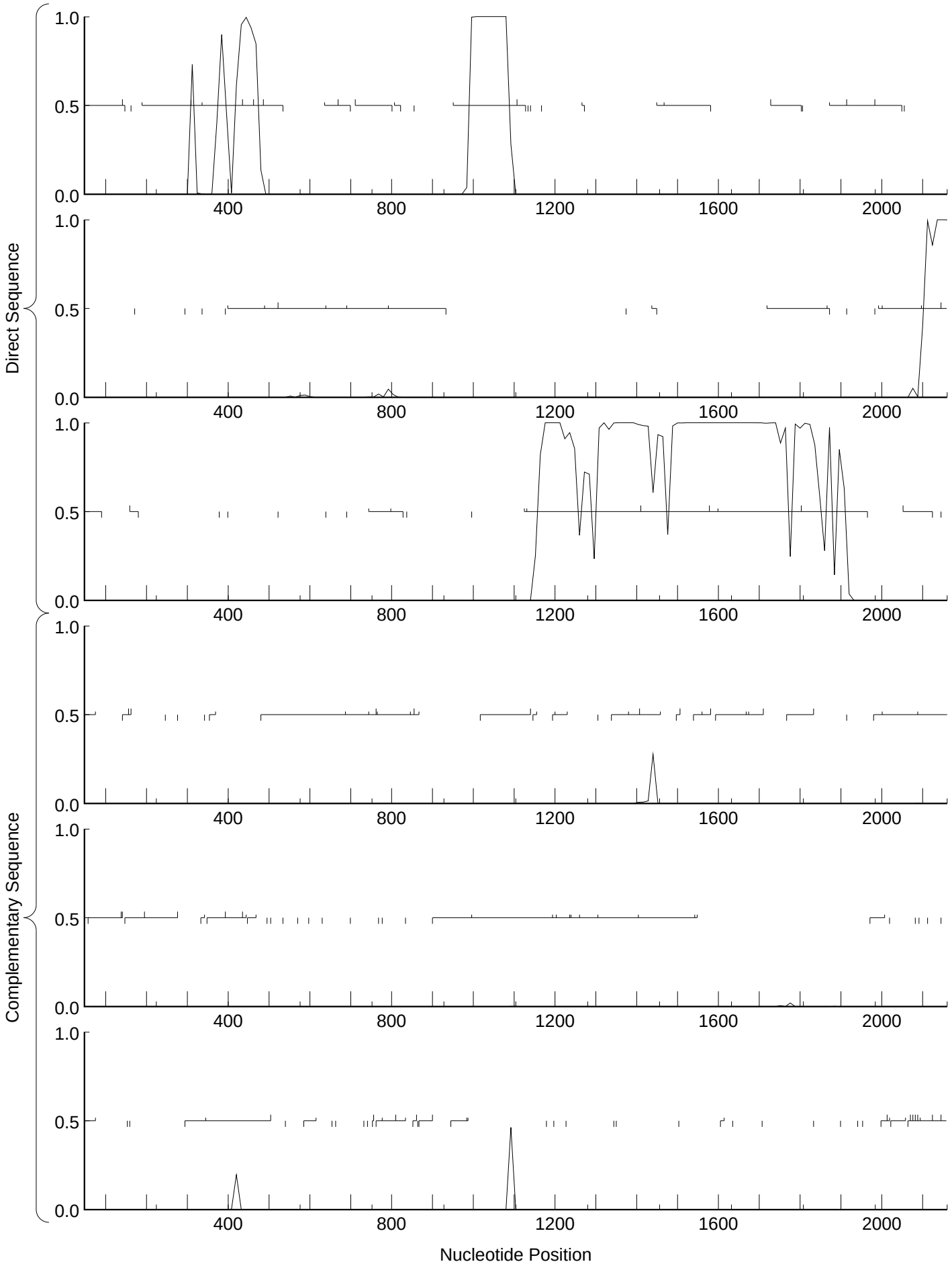
Order      :  4


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

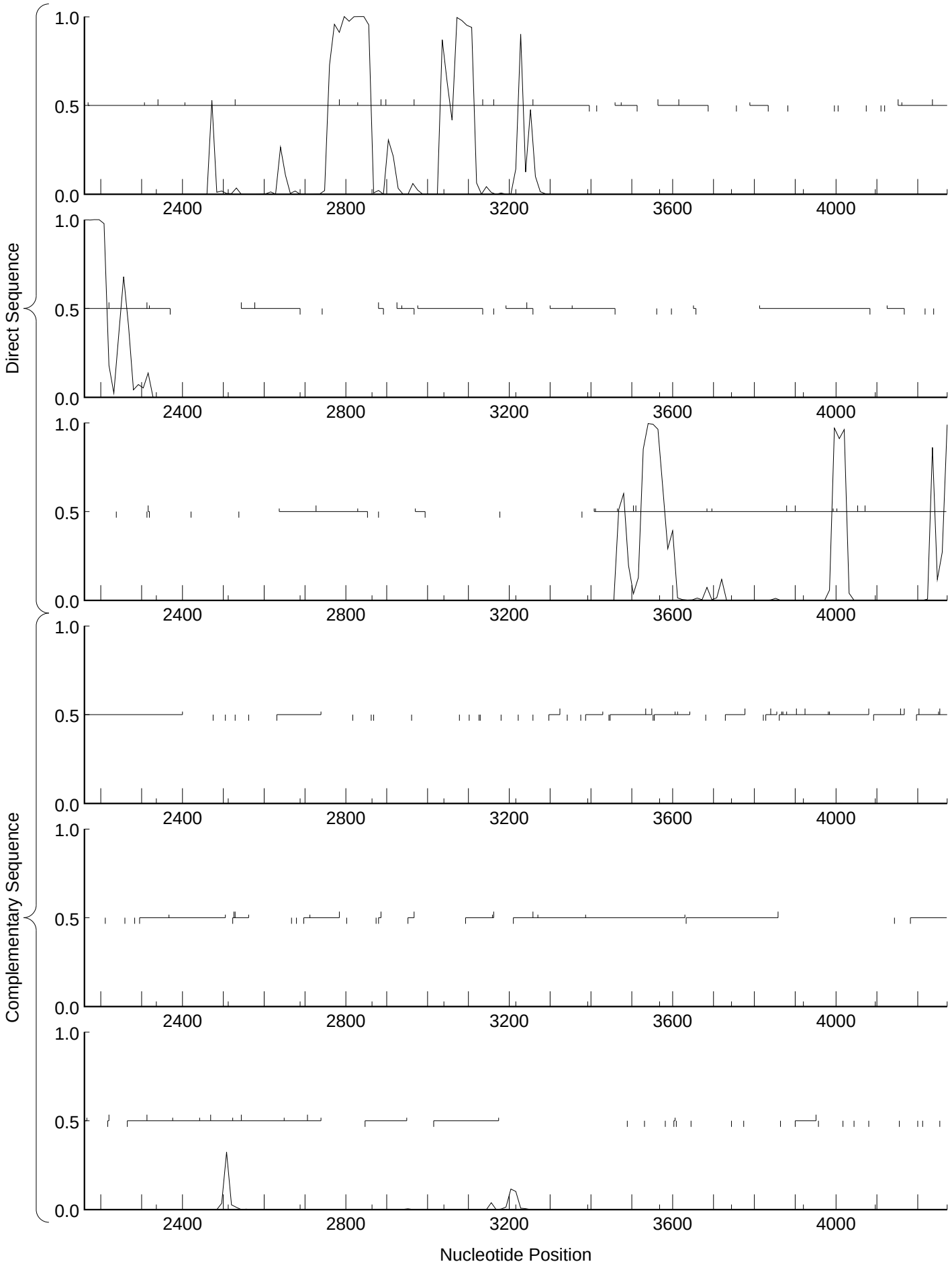



*For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu*

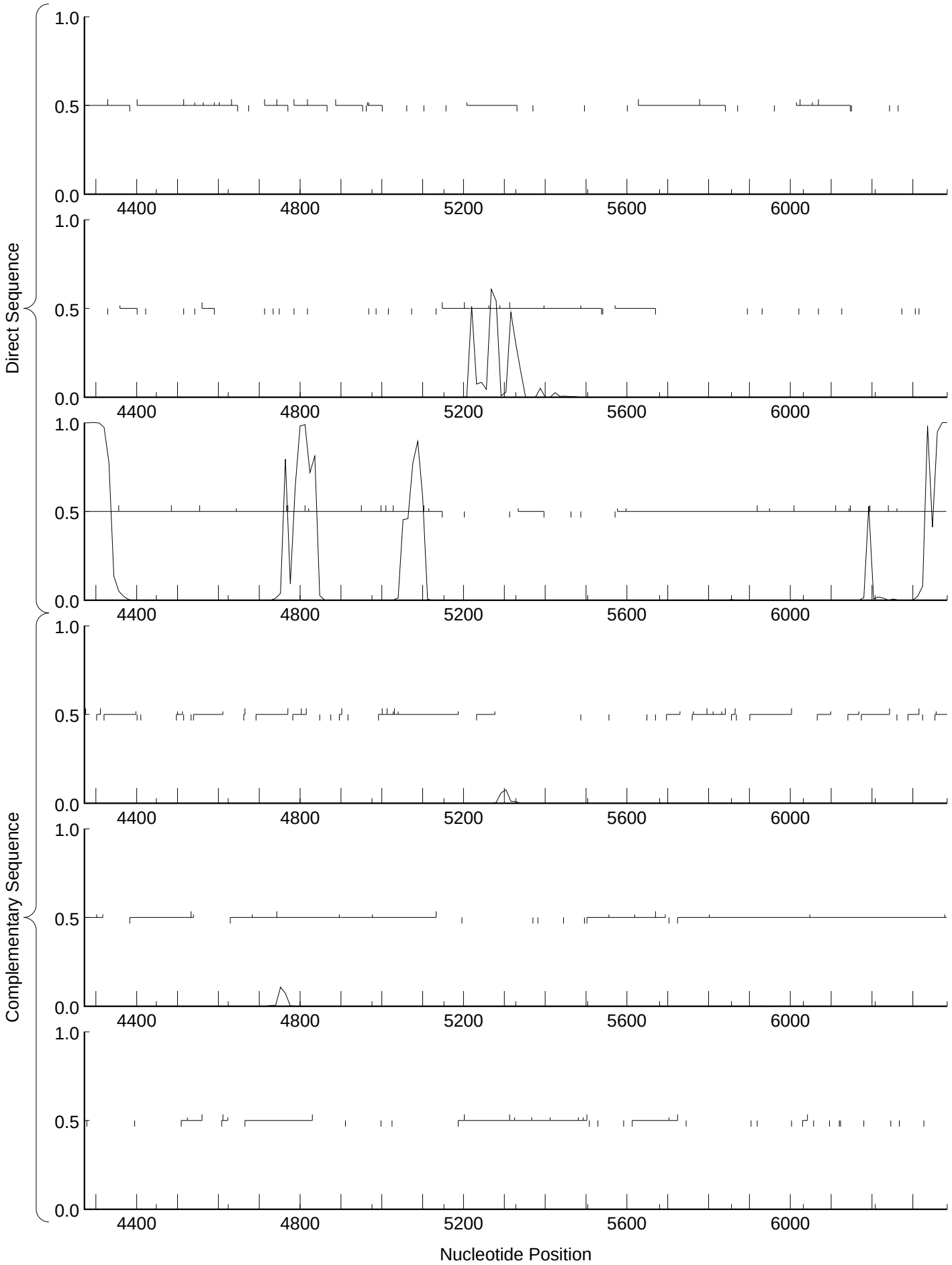
ydai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 2/29



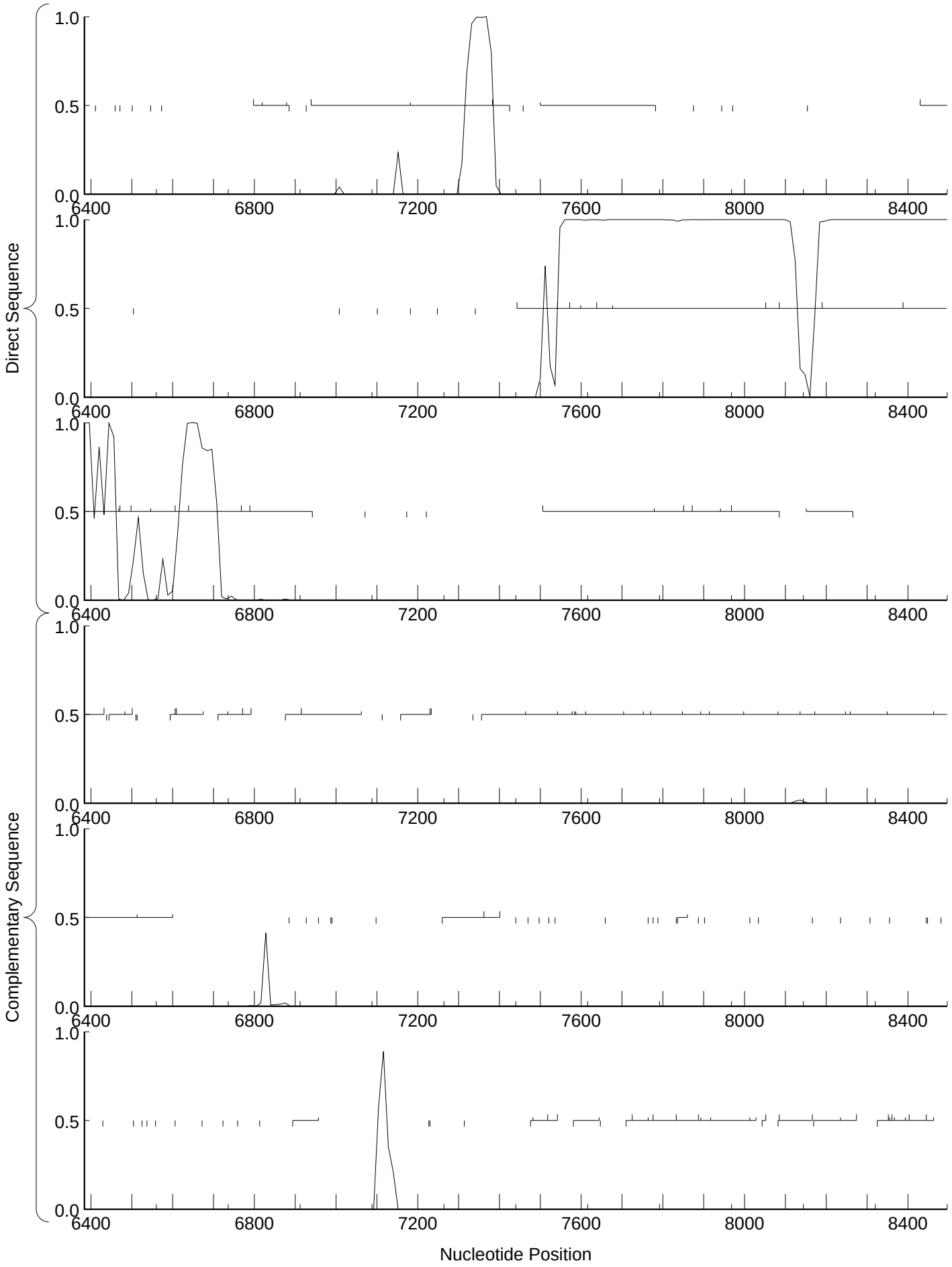
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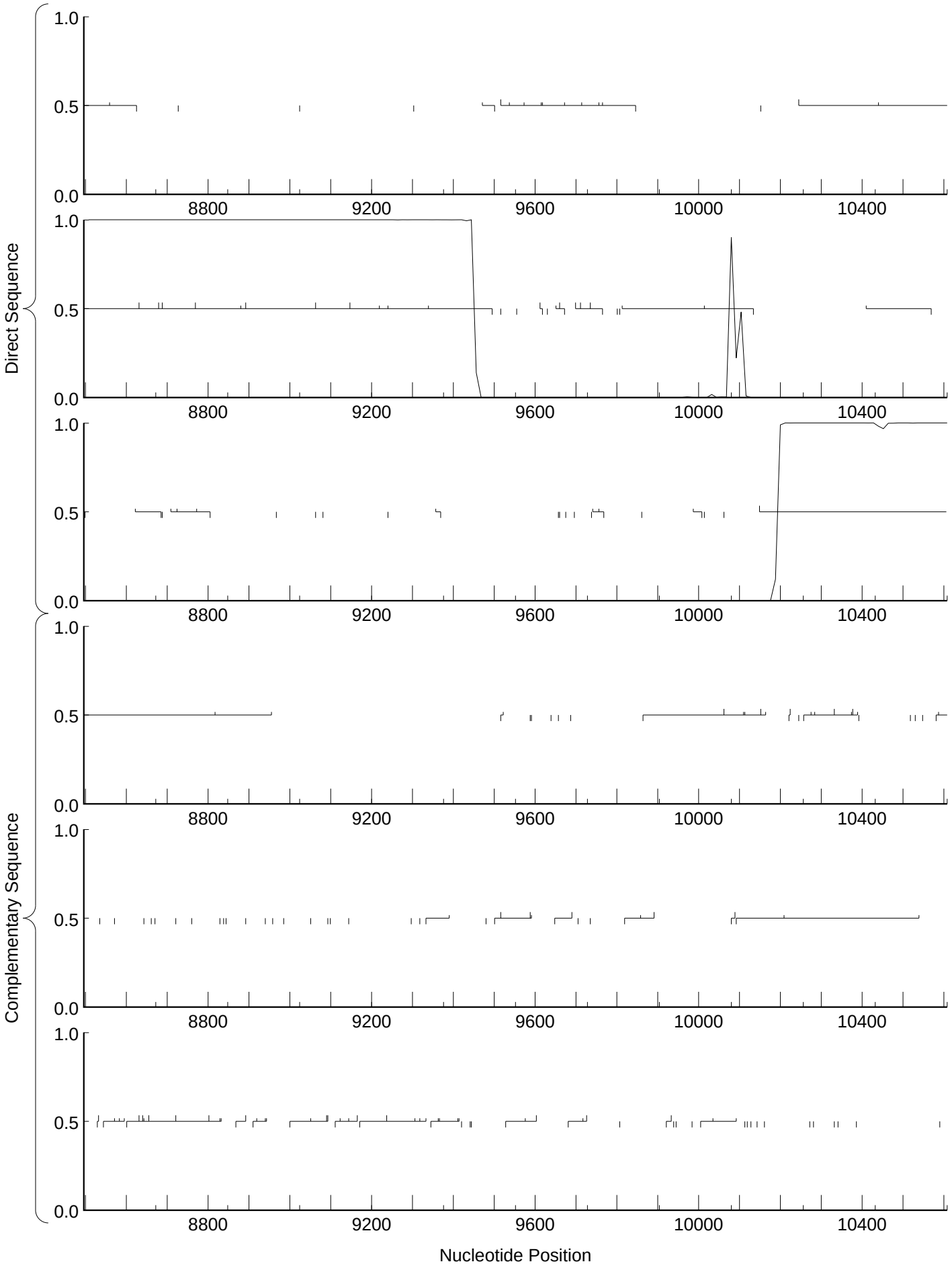
ydai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 4/29



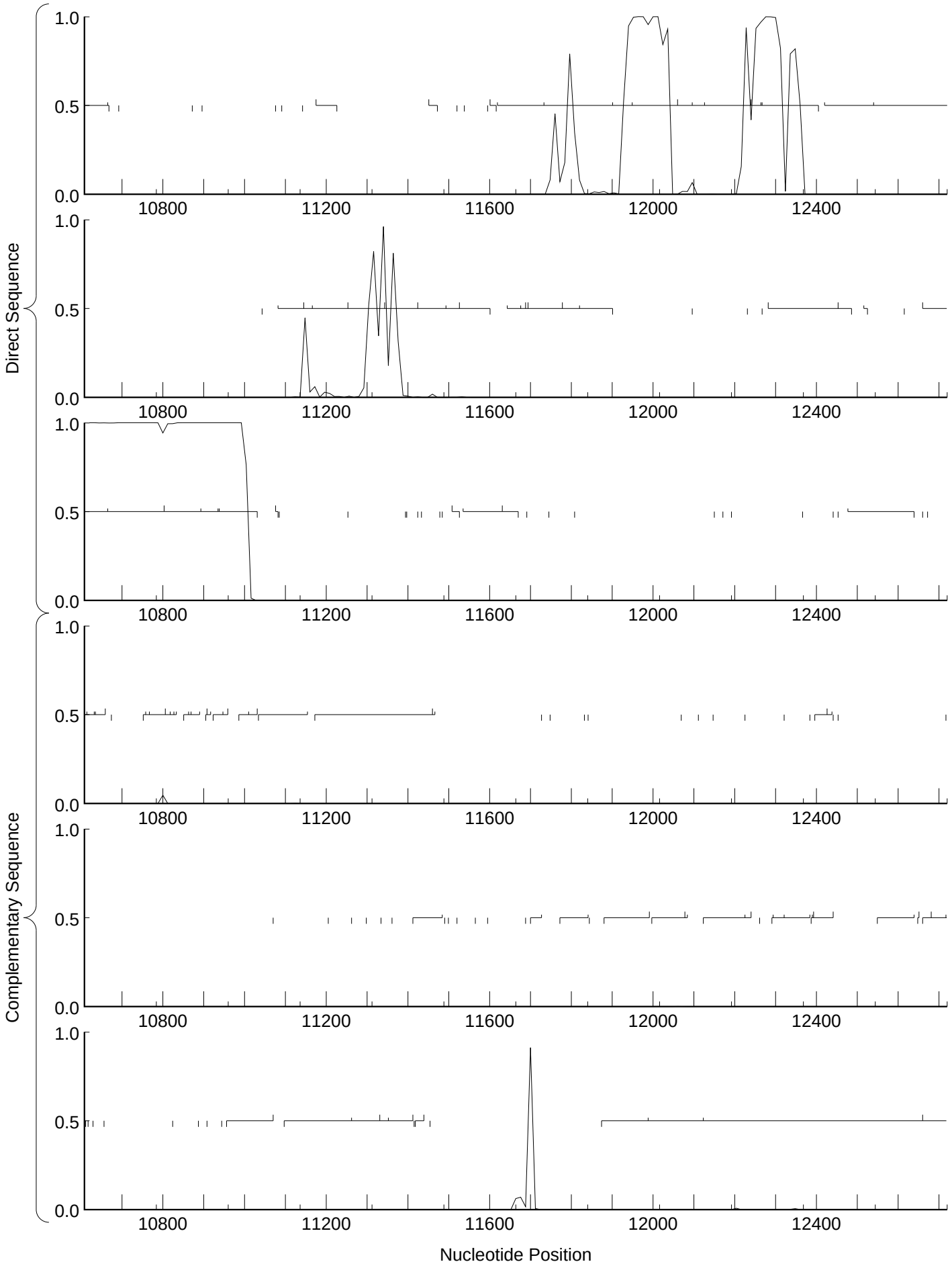
ydai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 5/29



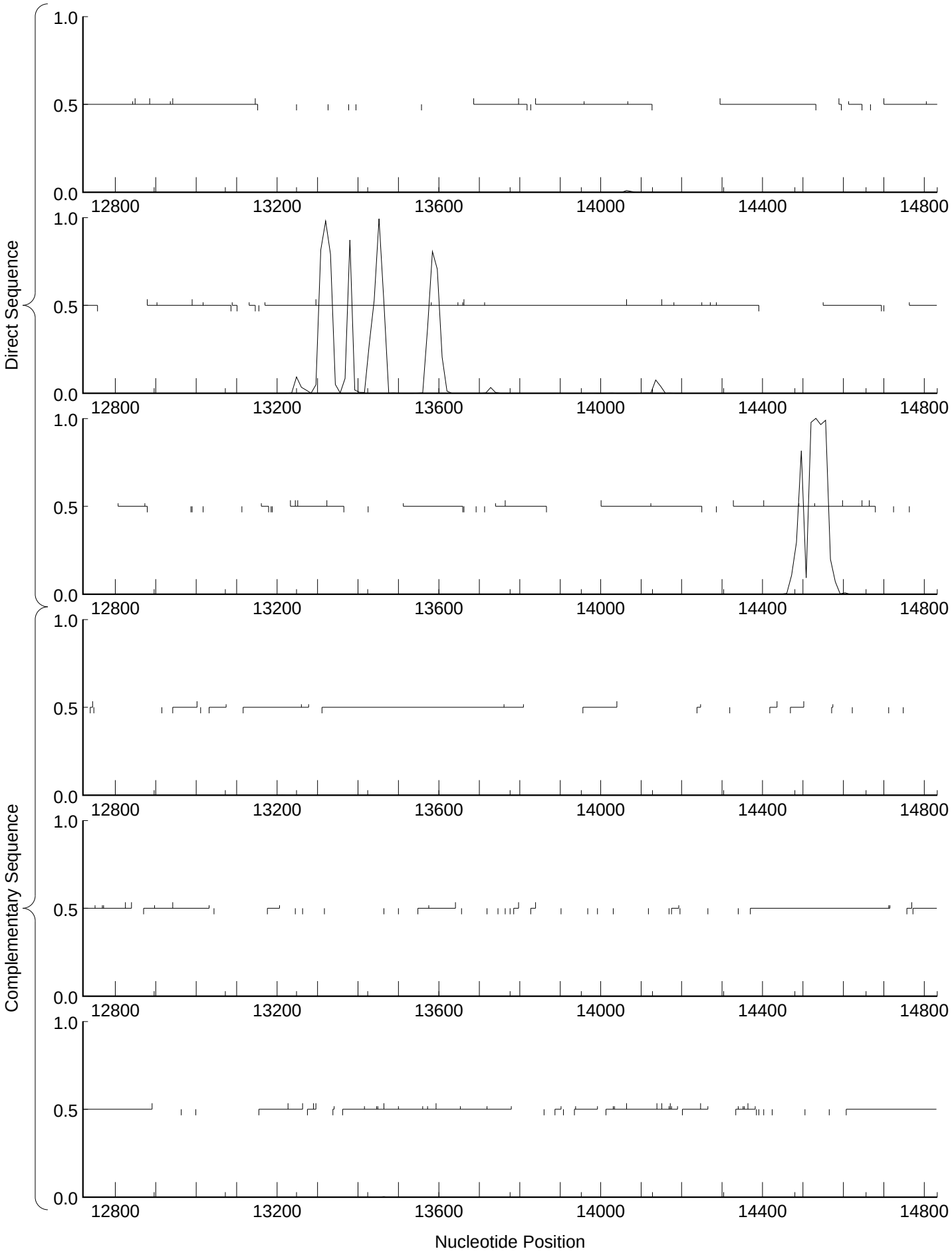
ydai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 6/29



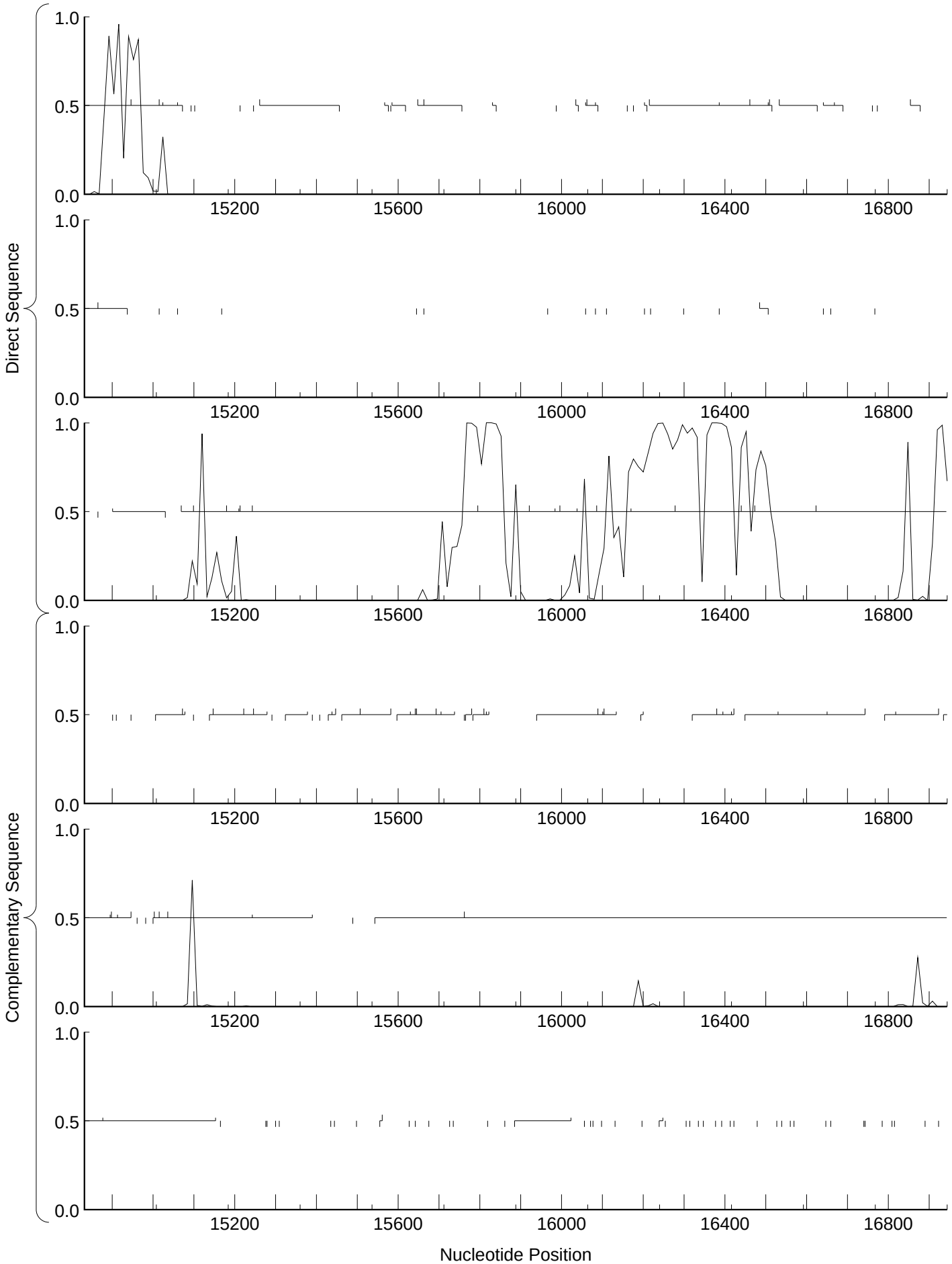
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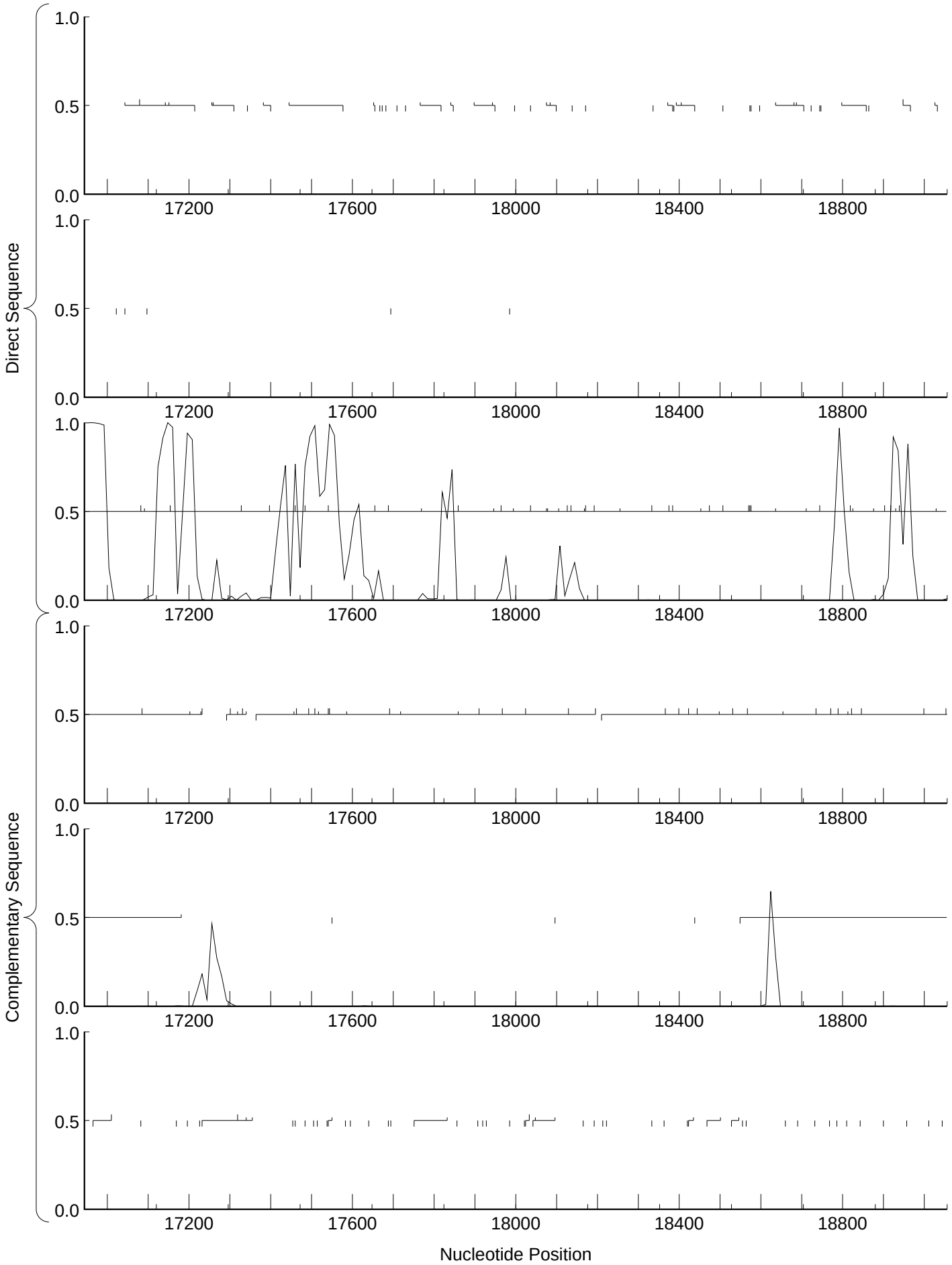
ydai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 8/29



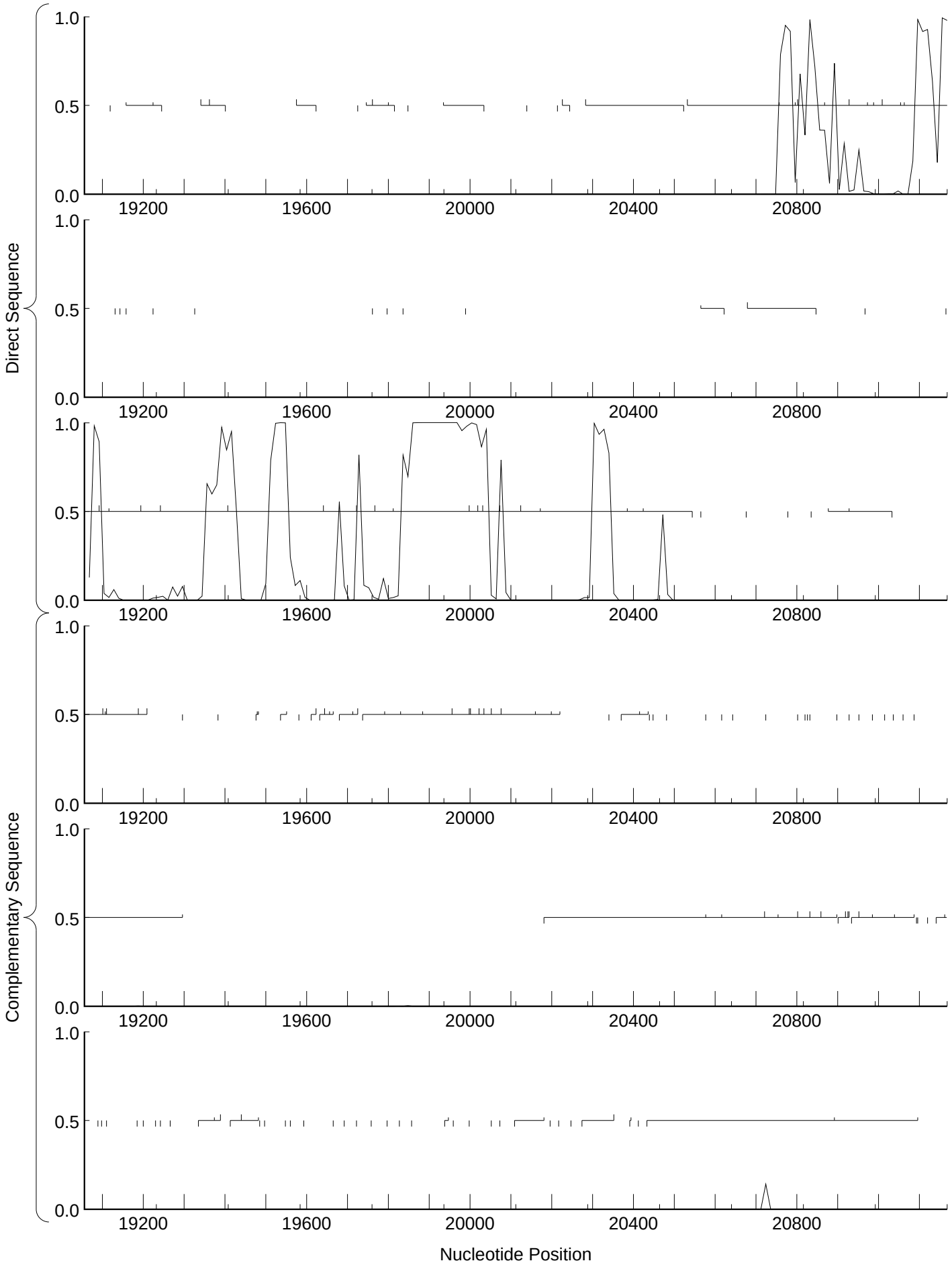
ydai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 9/29



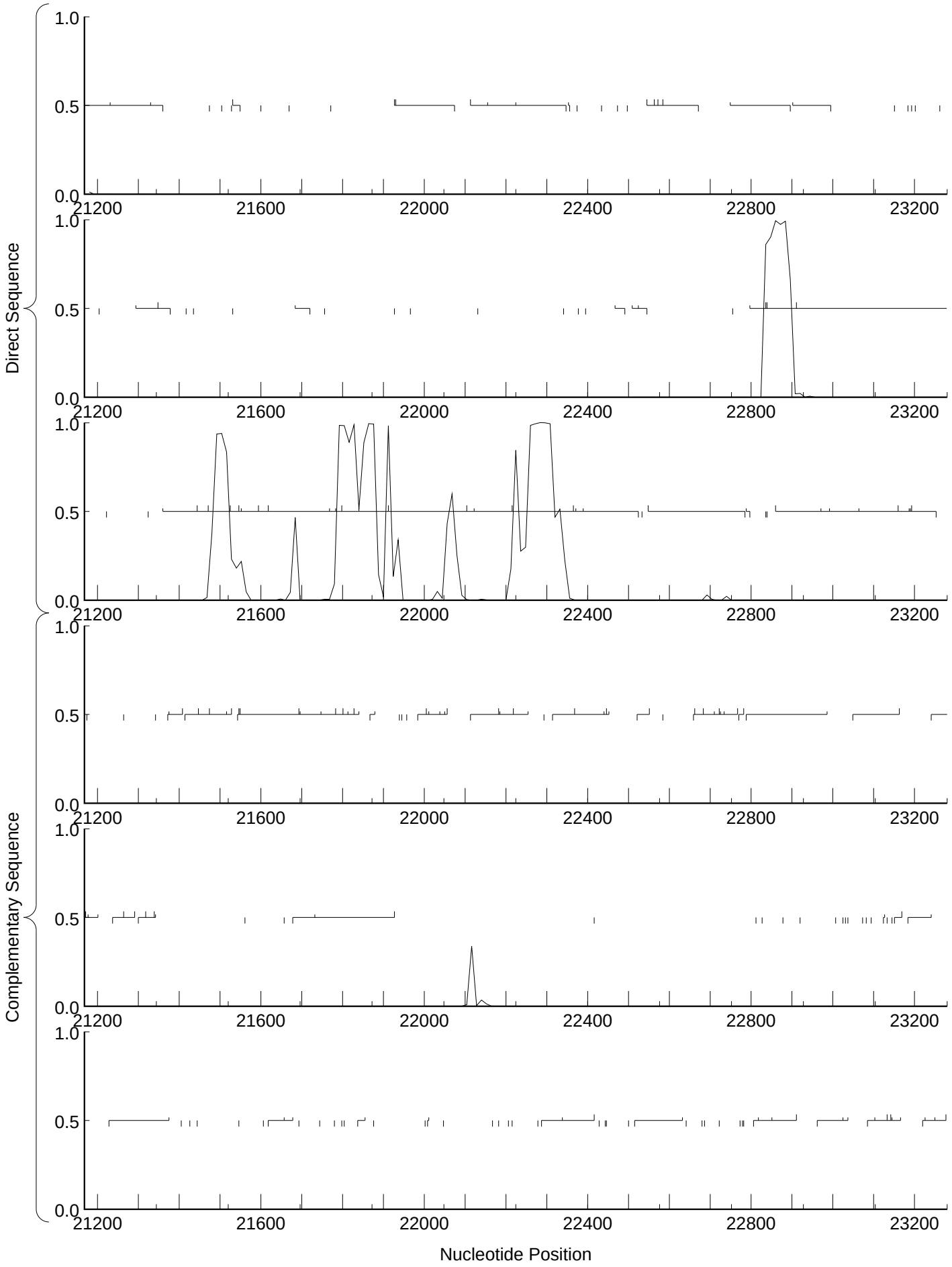
lai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 10/29



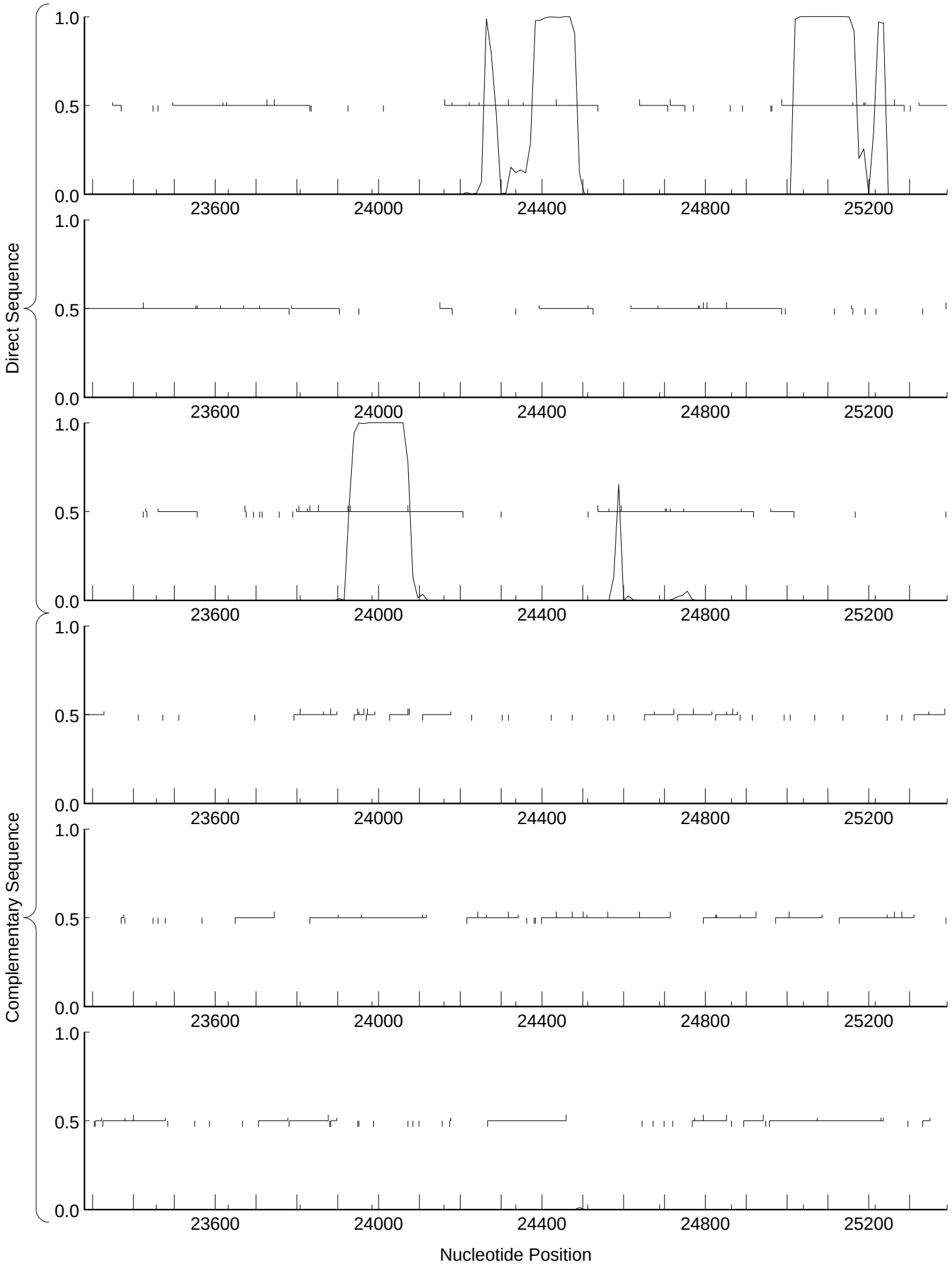
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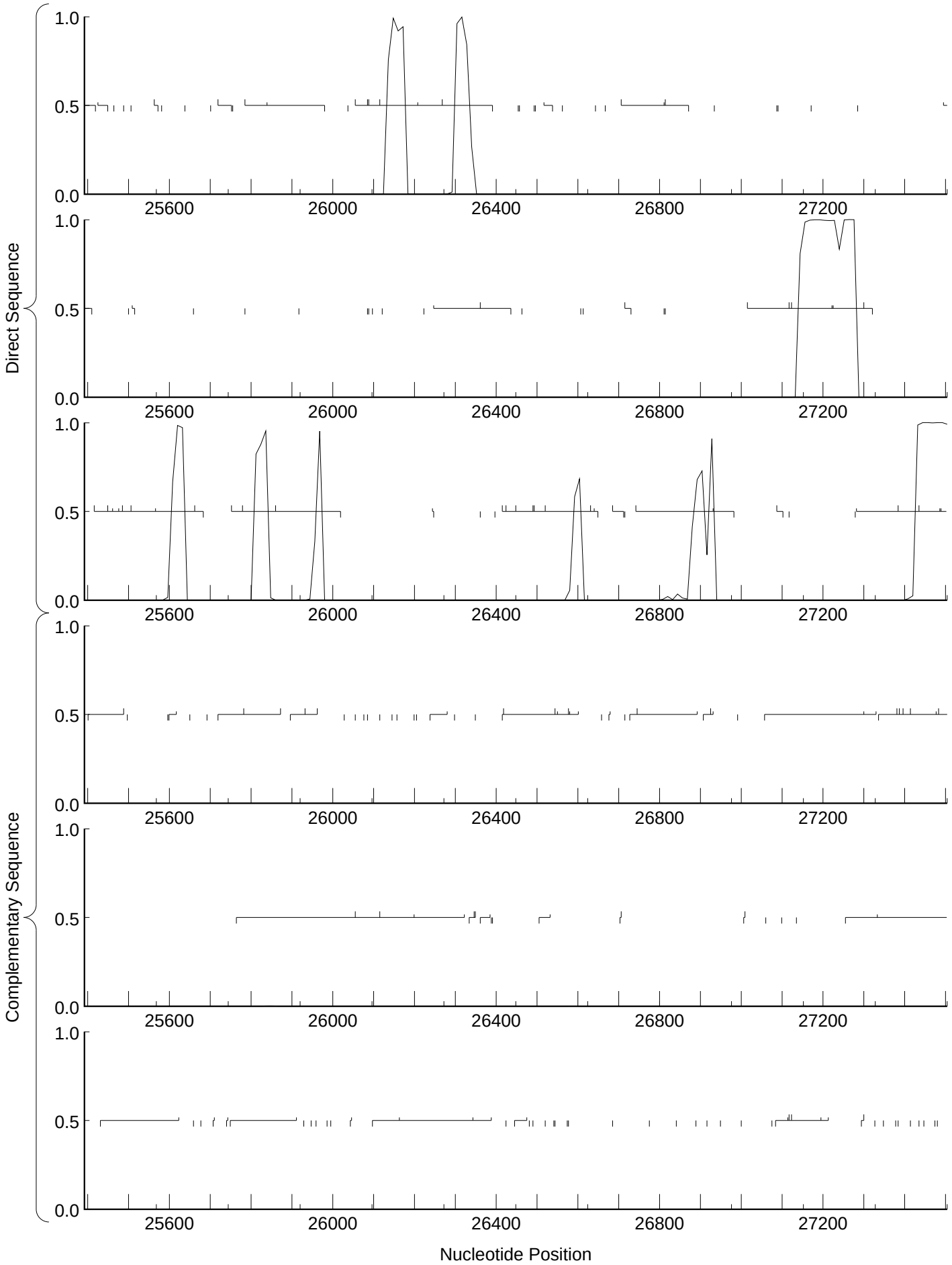
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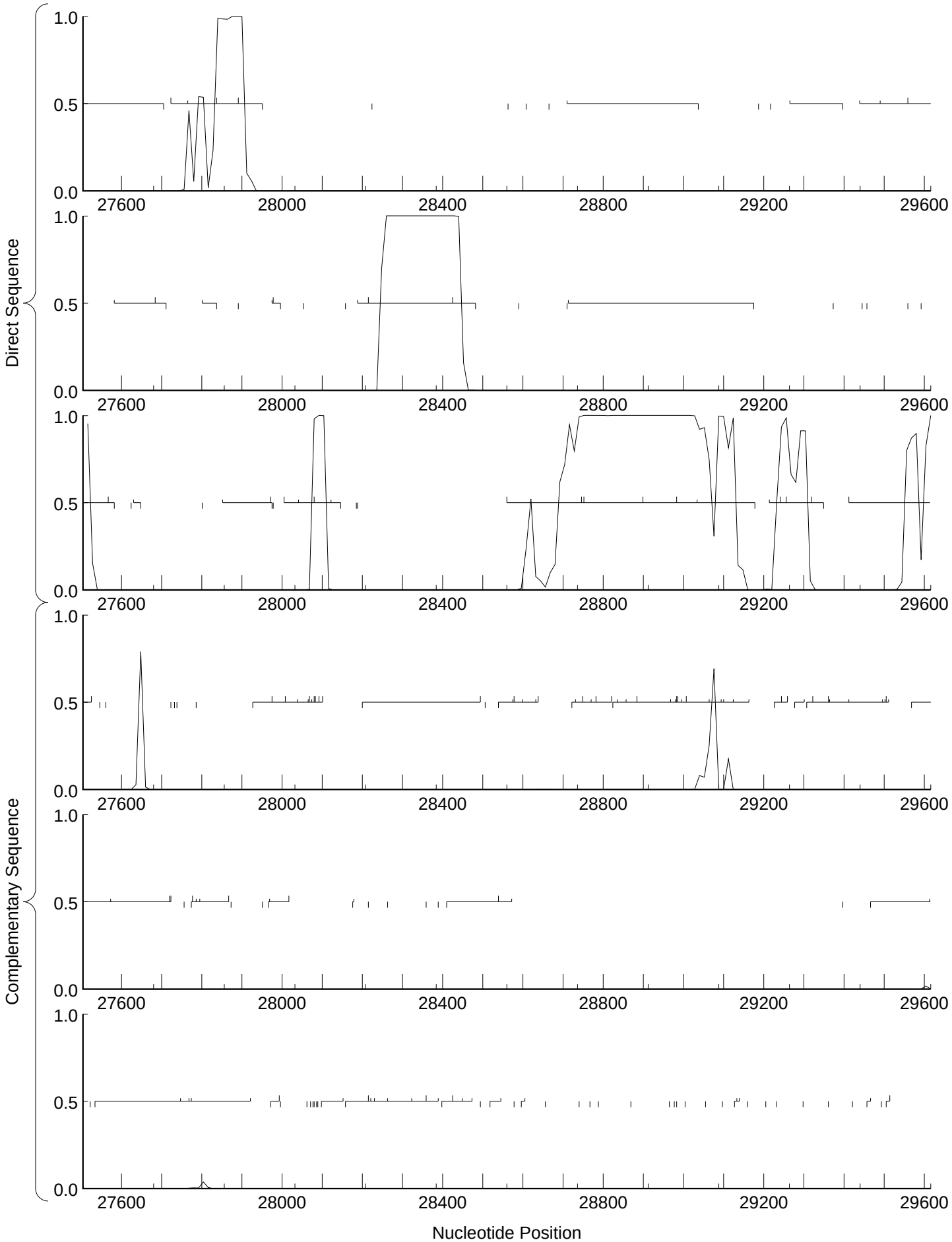
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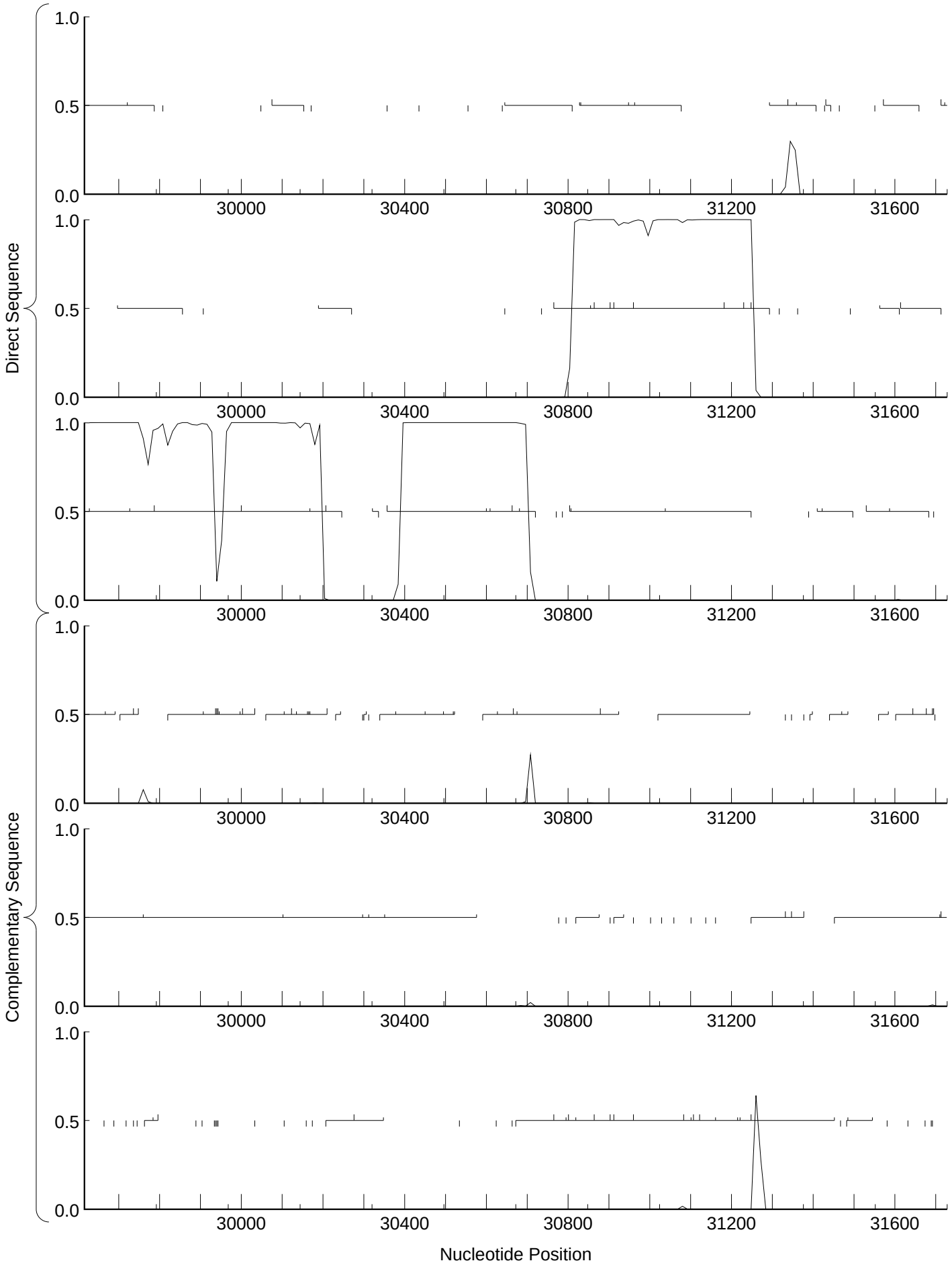


lai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 14/29

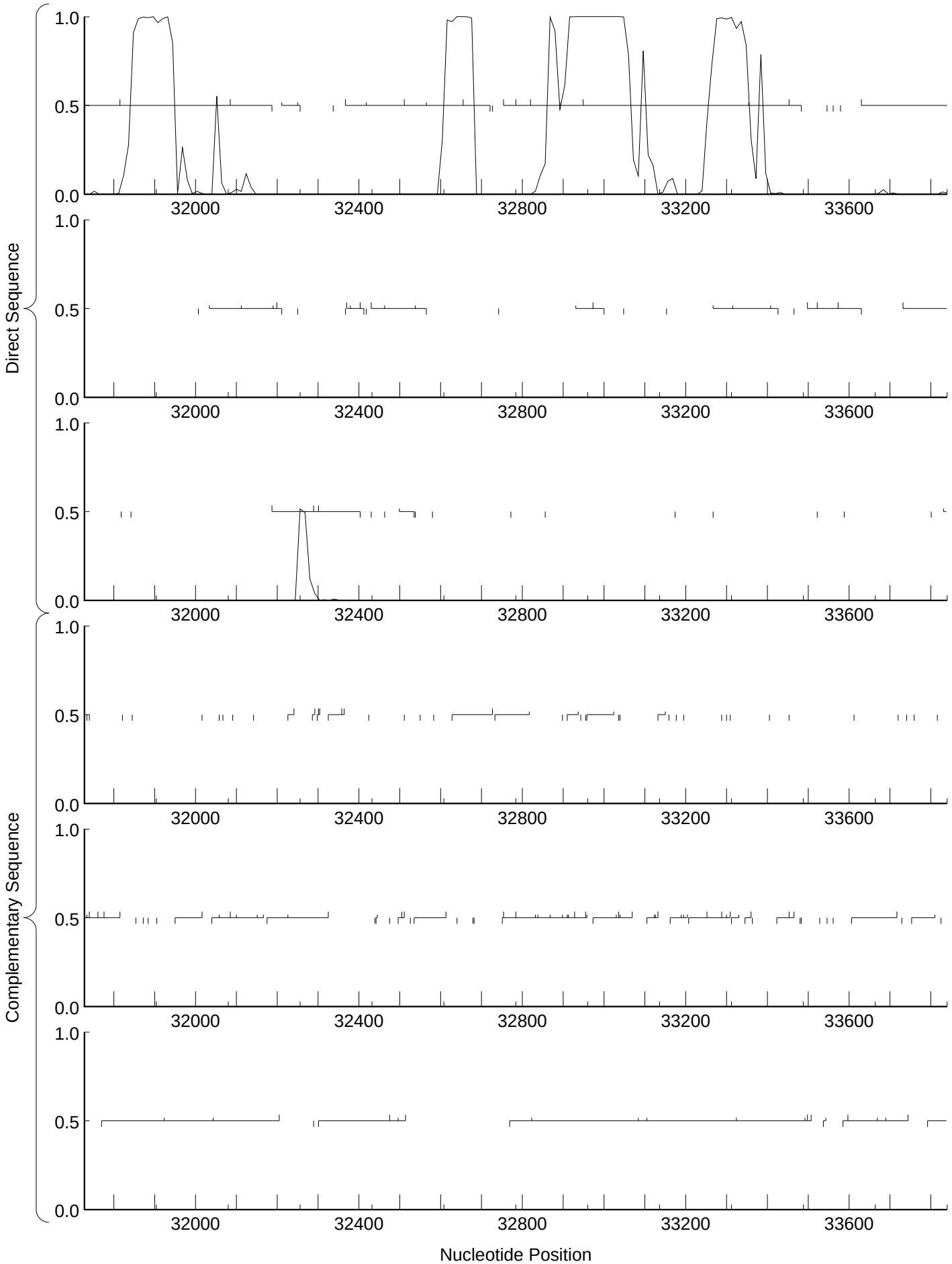


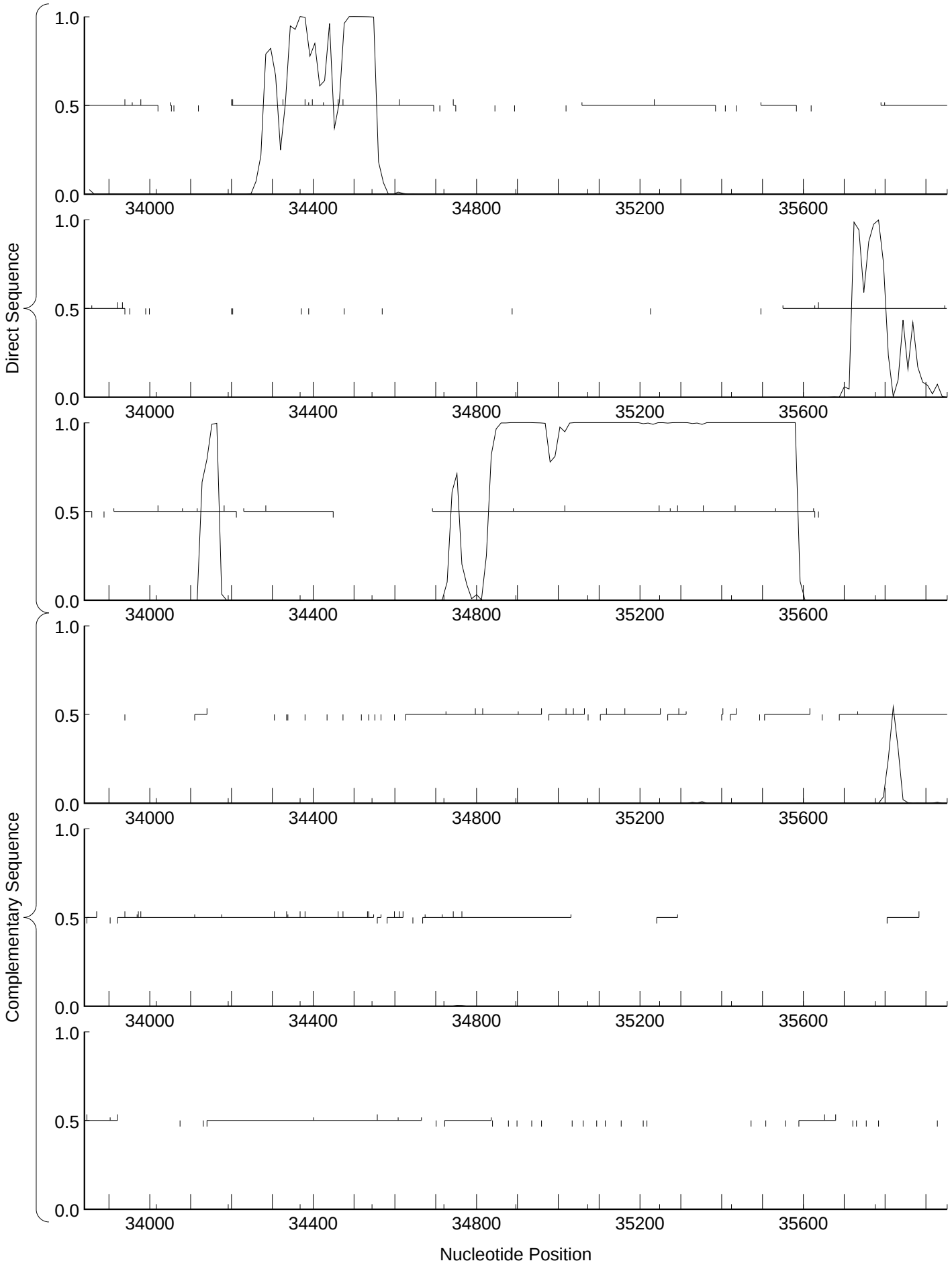
lai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 15/29



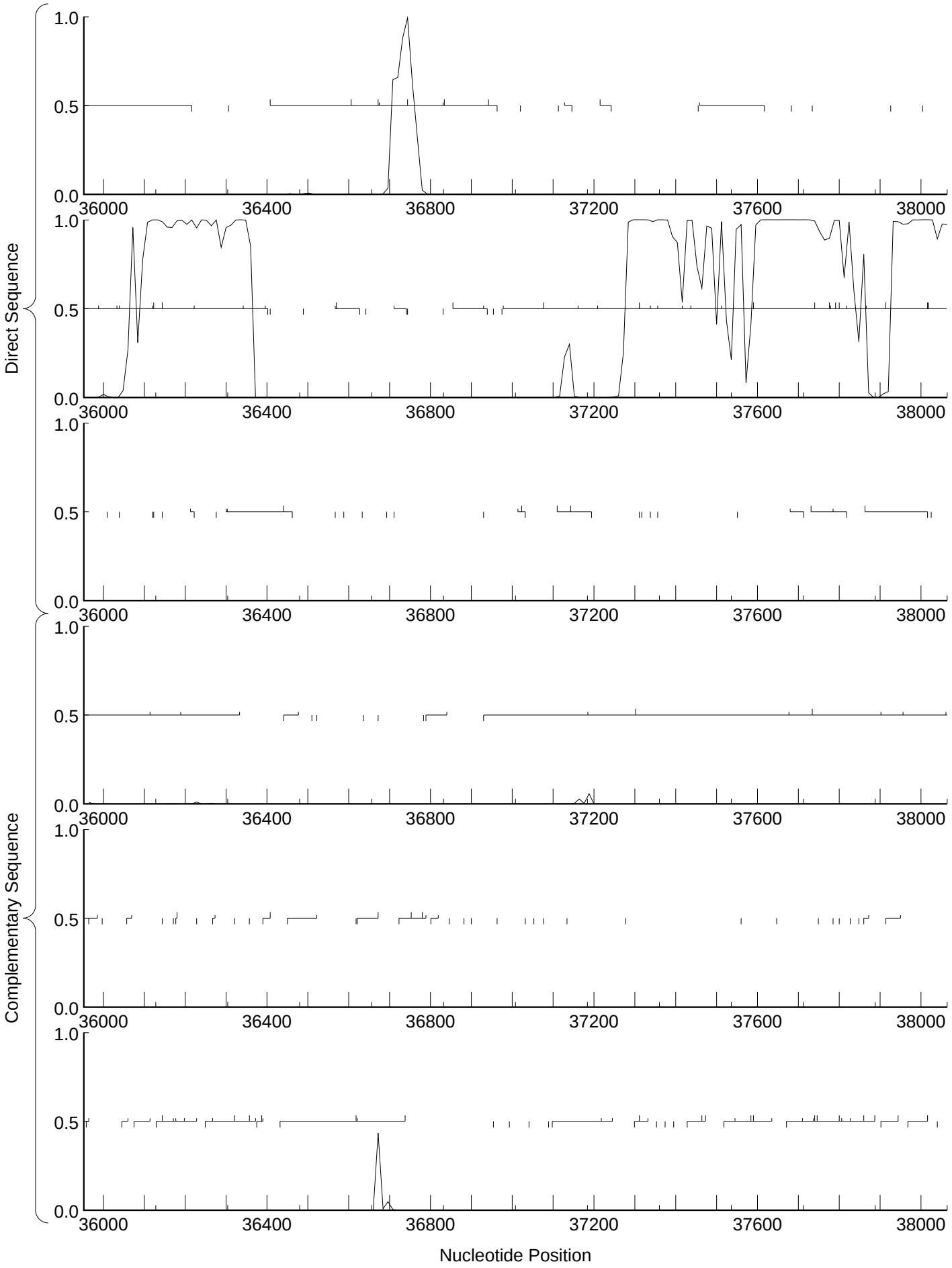

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
30000
30400
30800
31200
31600
Nucleotide Position

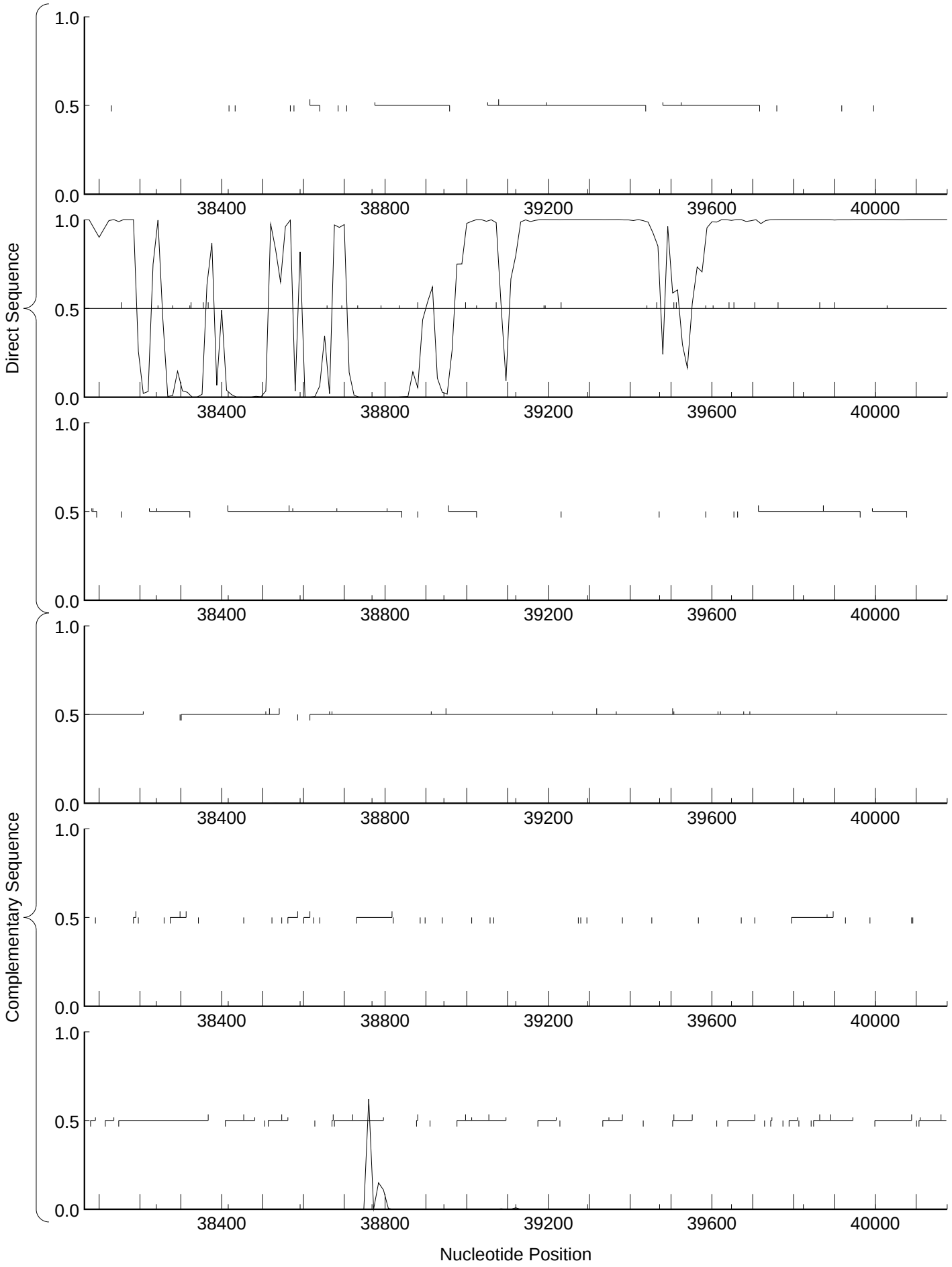
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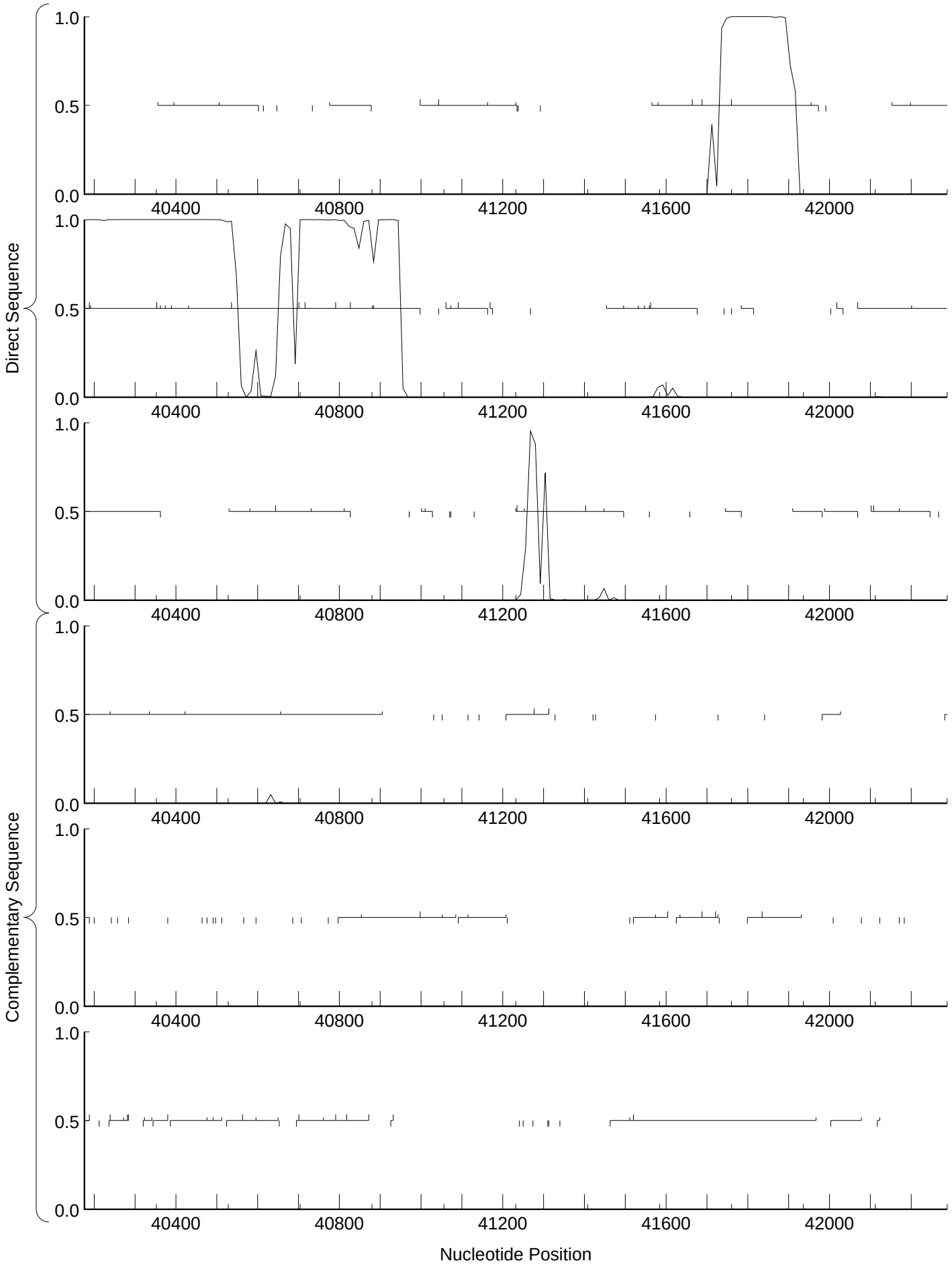


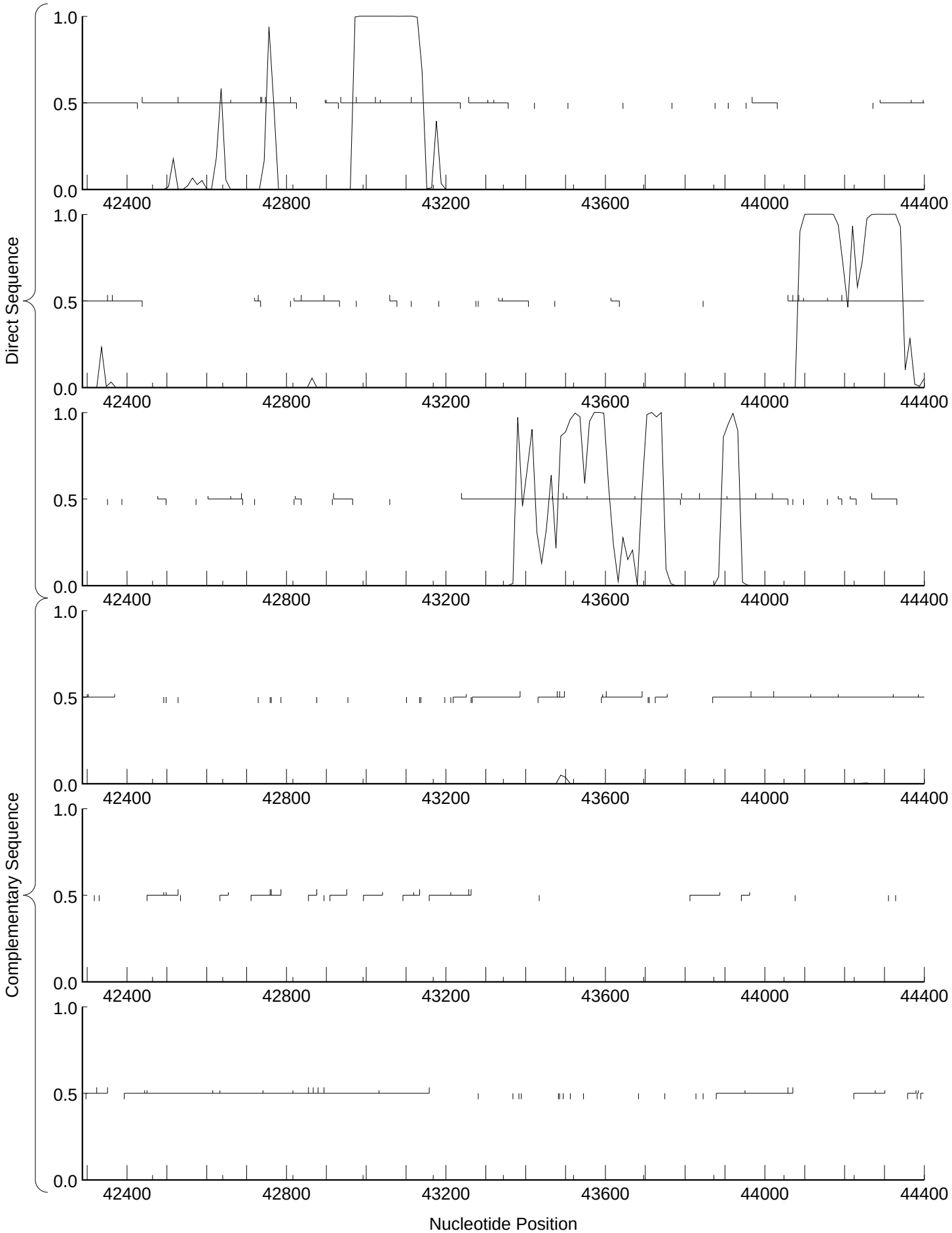

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

lai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 19/29

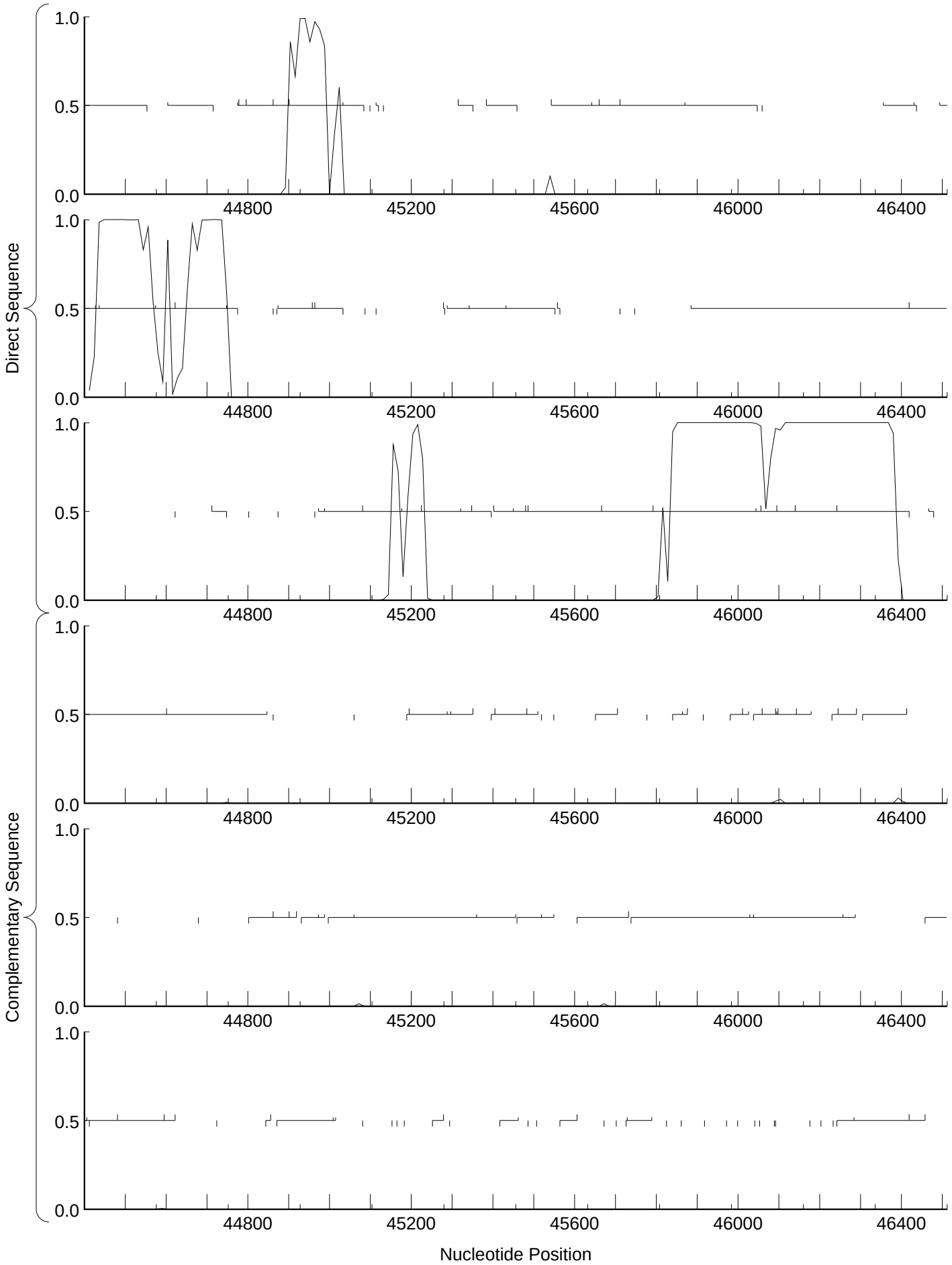



Direct Sequence
Complementary Sequence
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0.5
0.0
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38800
39200
39600
40000
Nucleotide Position

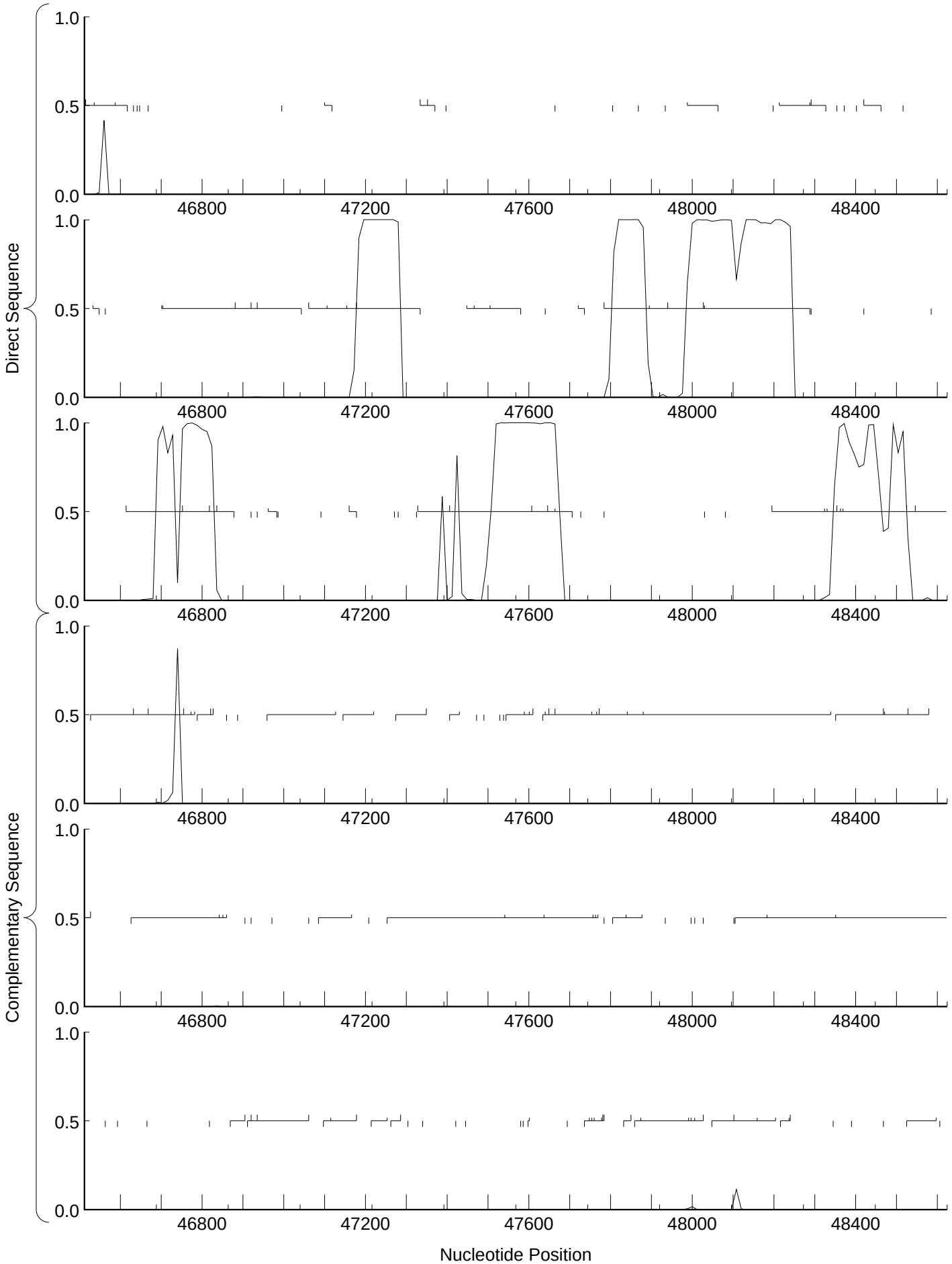

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
40400
40800
41200
41600
42000
Nucleotide Position


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

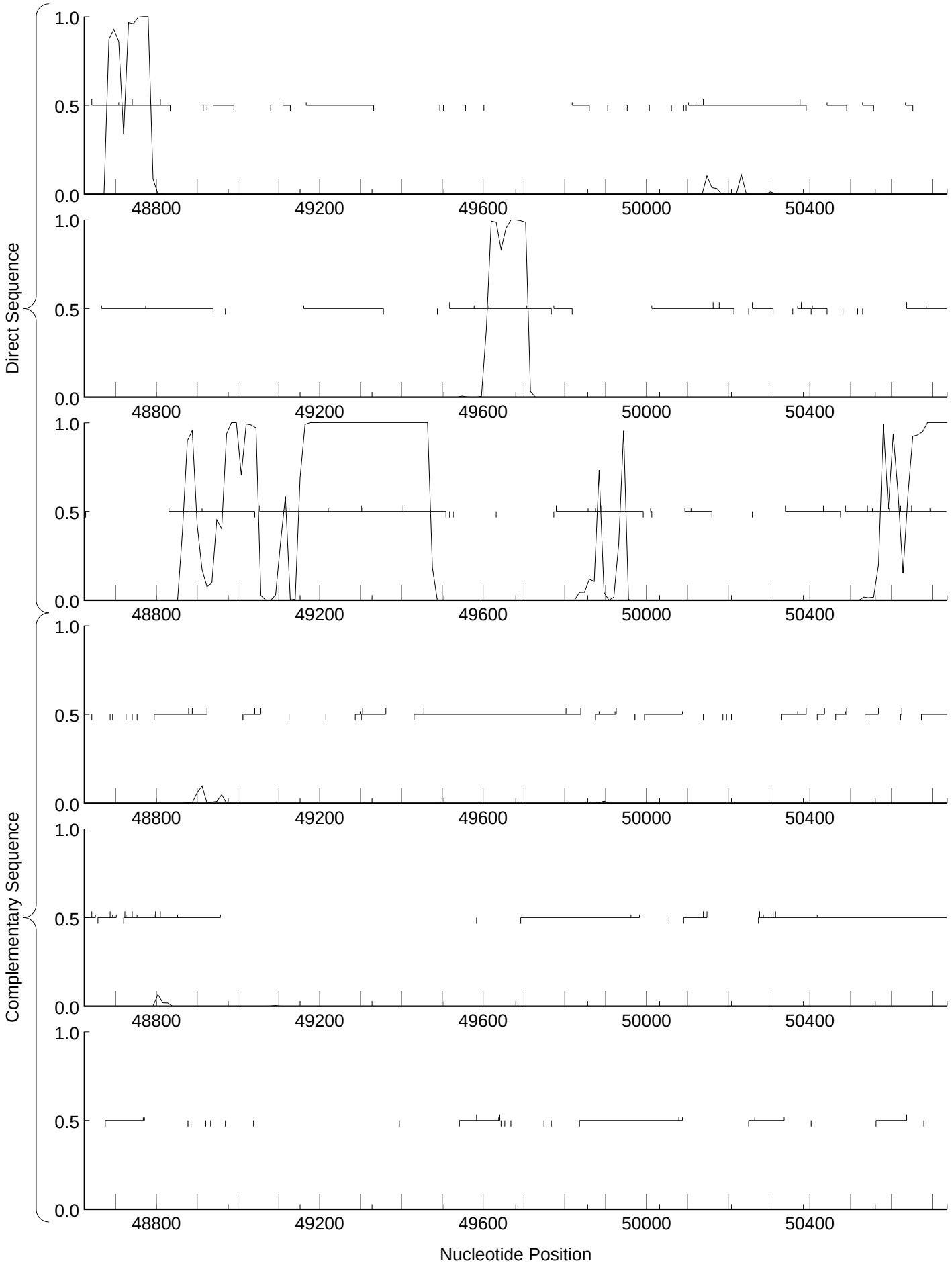
lai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 23/29



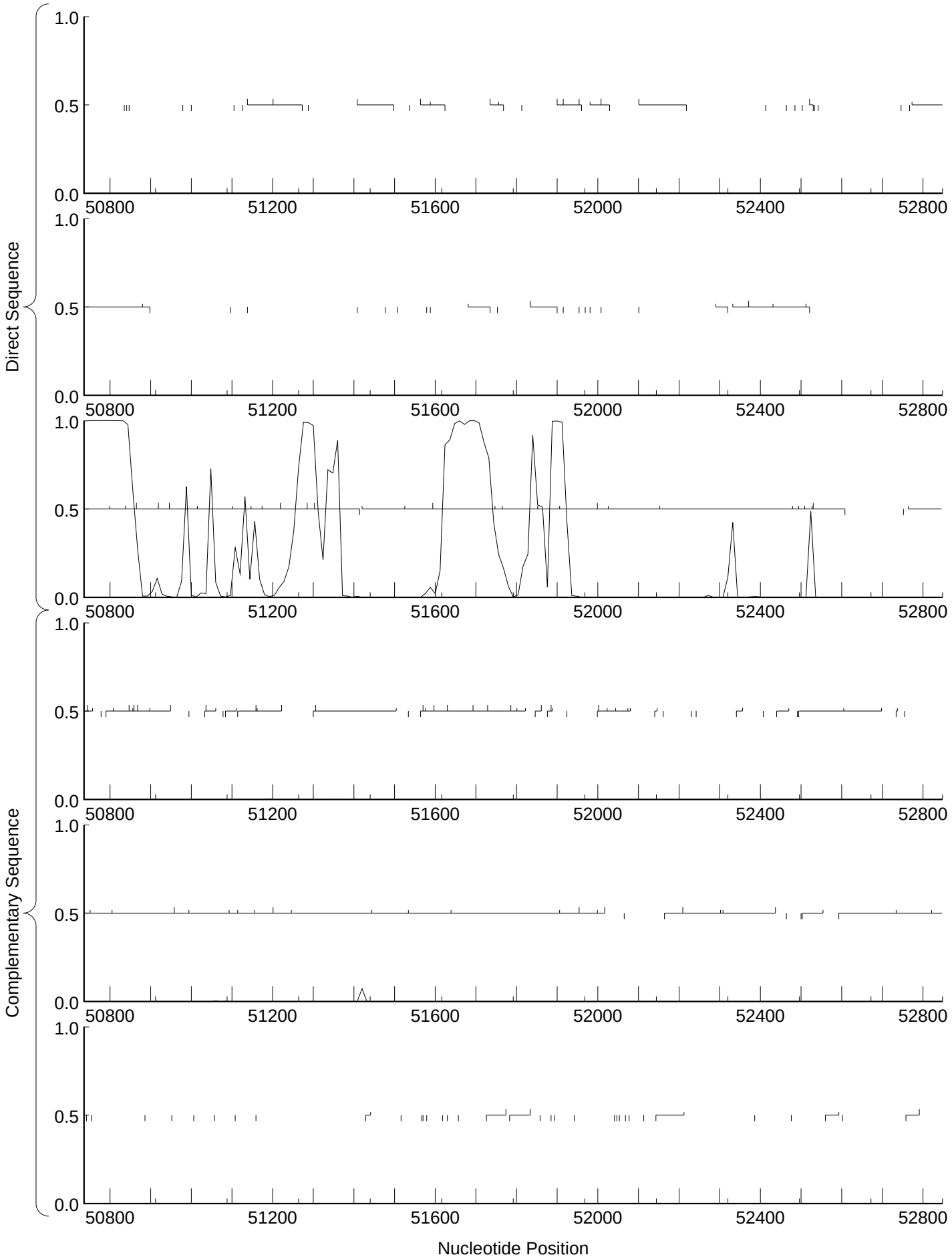
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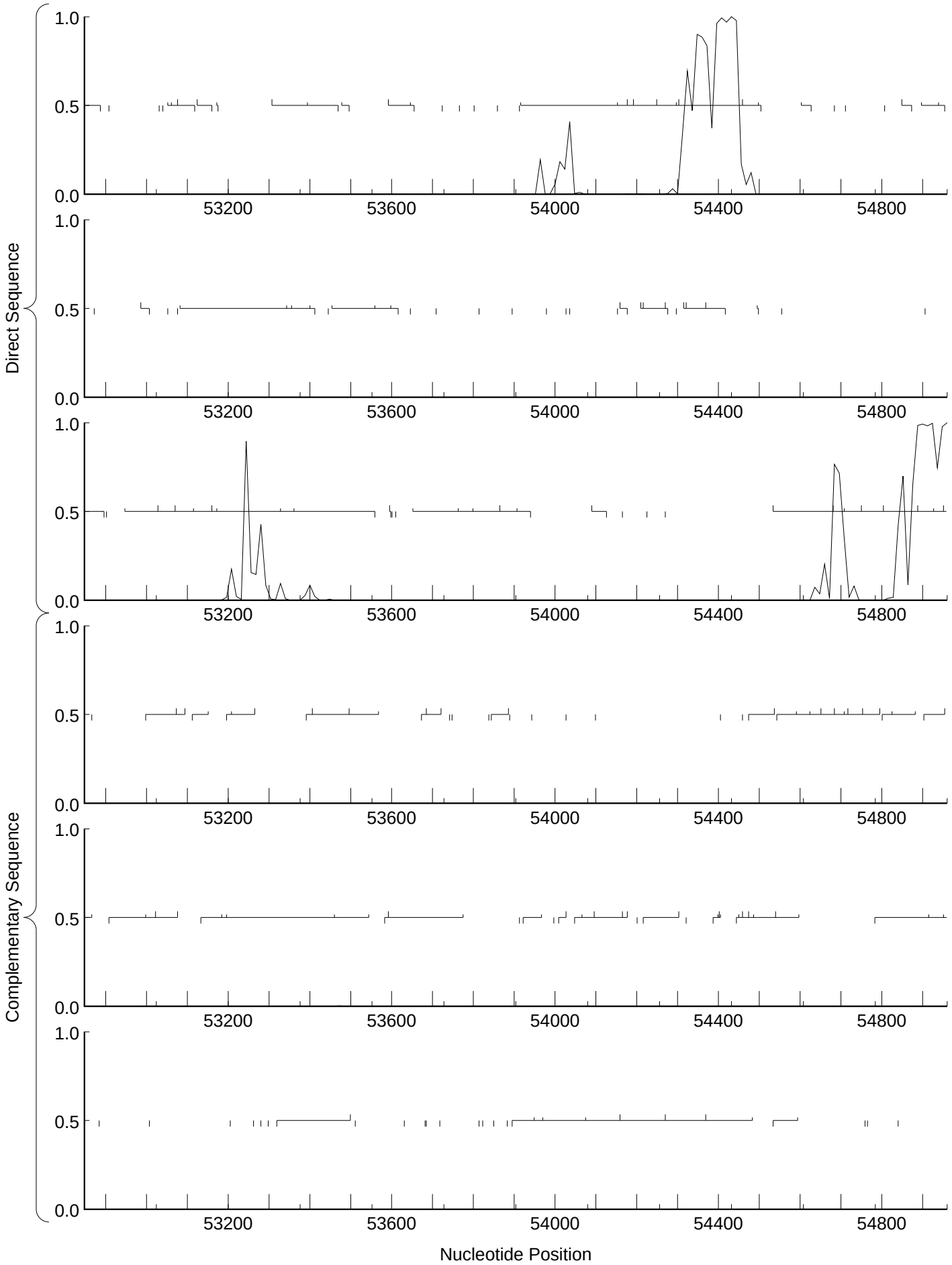


lai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 25/29

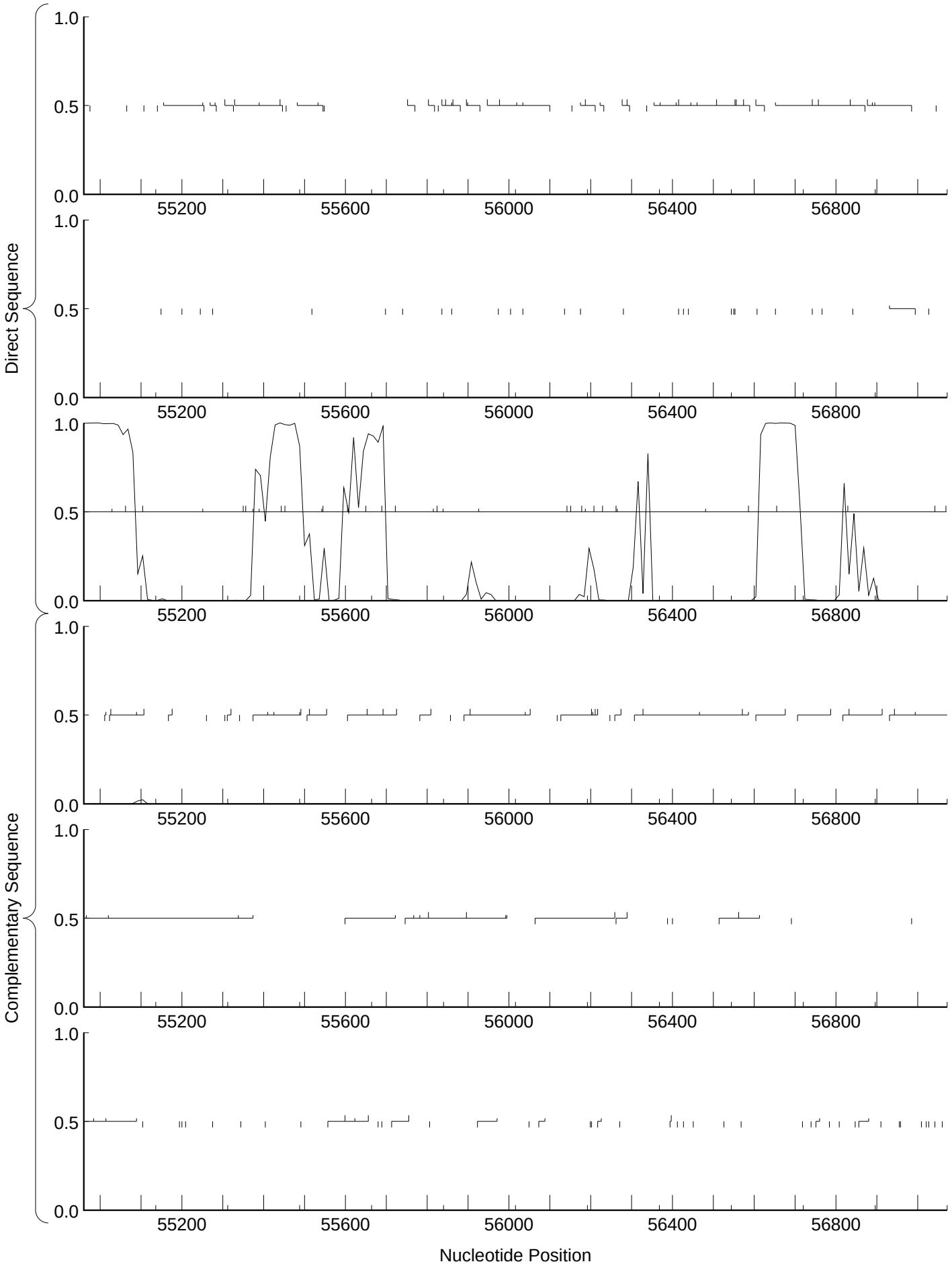


lai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 26/29




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
53200
53600
54000
54400
54800
Nucleotide Position

lai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 28/29



lai complete sequence, 57623 bp including 9-base 3’ overhang (CGCCCGCCT), Cluster BI4, Order 4, Window 96, Step 12, 29/29

