GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Gordonia phage SweatNTears complete sequence, 45917 bp including 10-base 3' overhang (CGGTA
Analysis Date    :  3/29/24 at 2:20:53
Pages            :  23
Sequence Length  :  45917 bp
GC Content       :   59.93%

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,
```

MATRIX INFORMATION

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Matrix     :  -
Author     :  -

Order      :  4

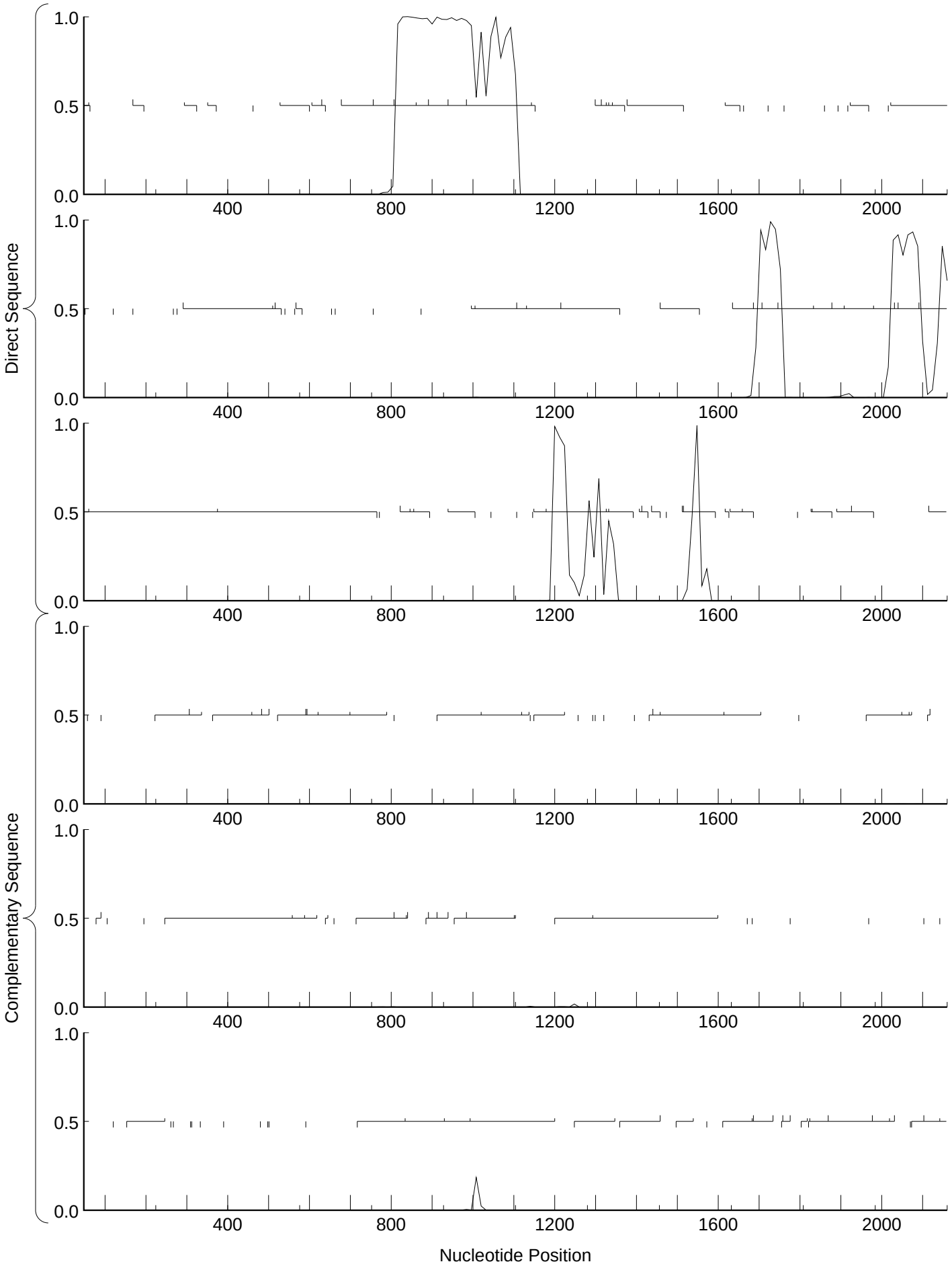

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

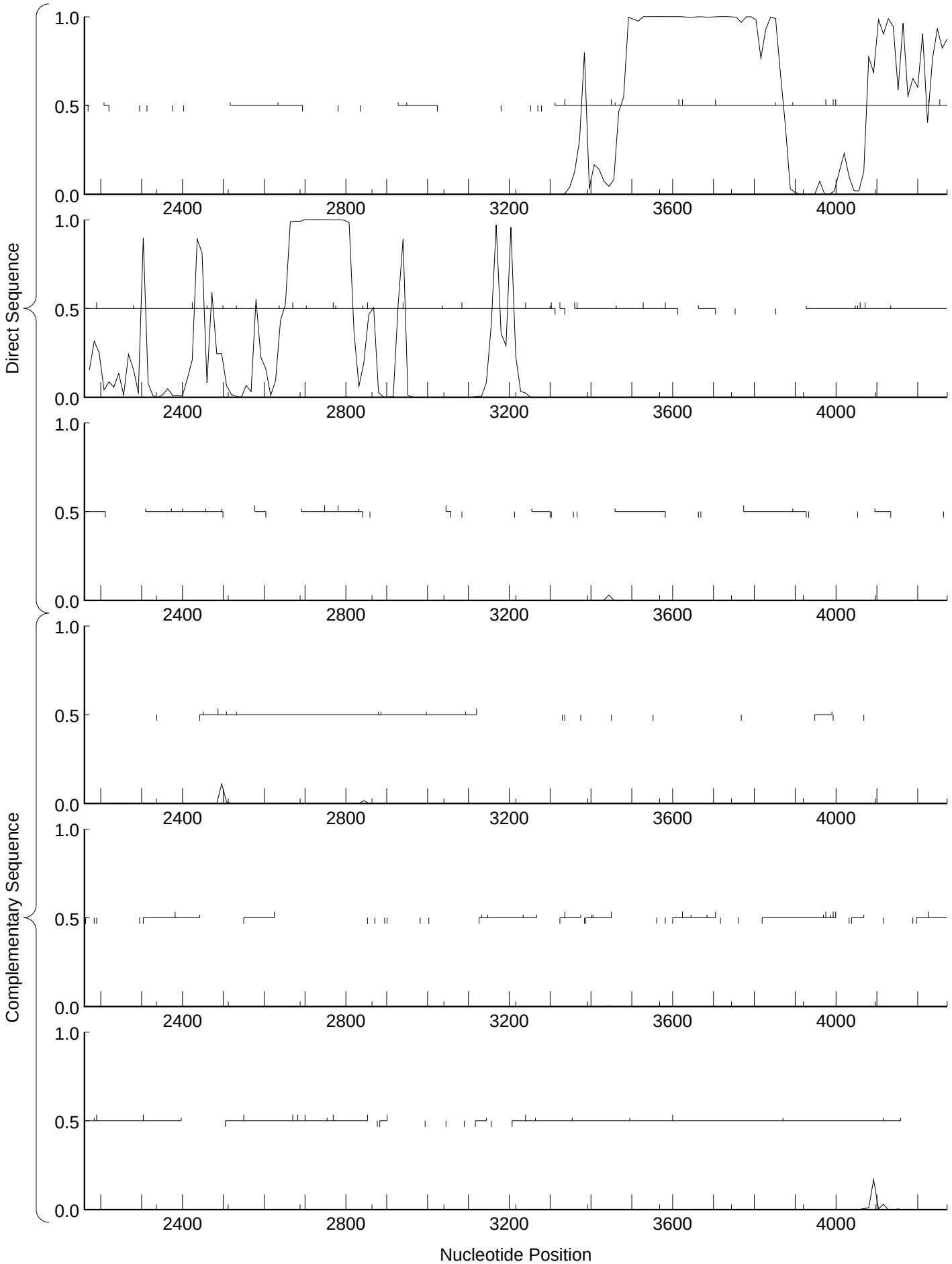
Matrix notes & comments

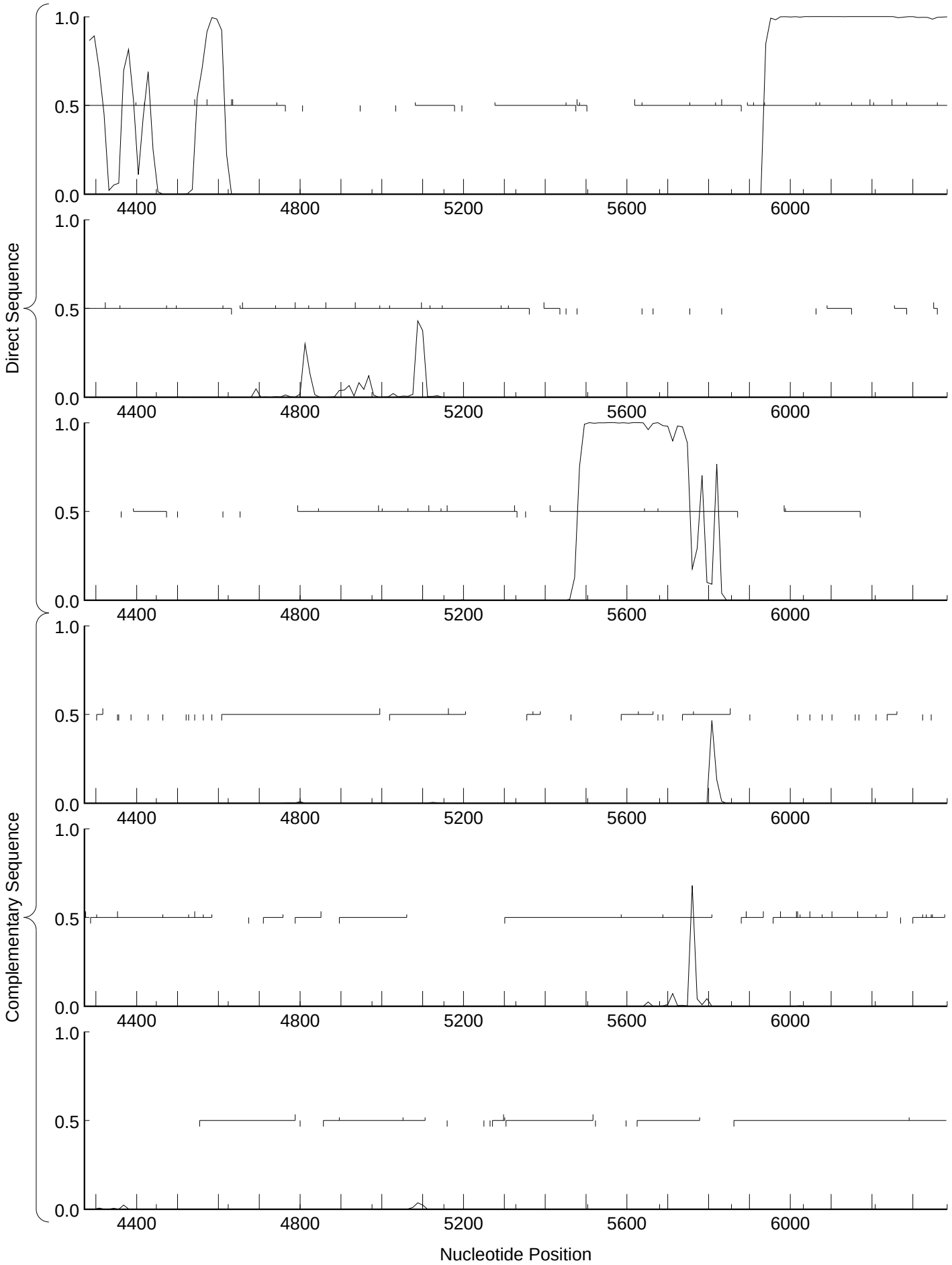
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Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep  8 00:36:14 2018
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For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

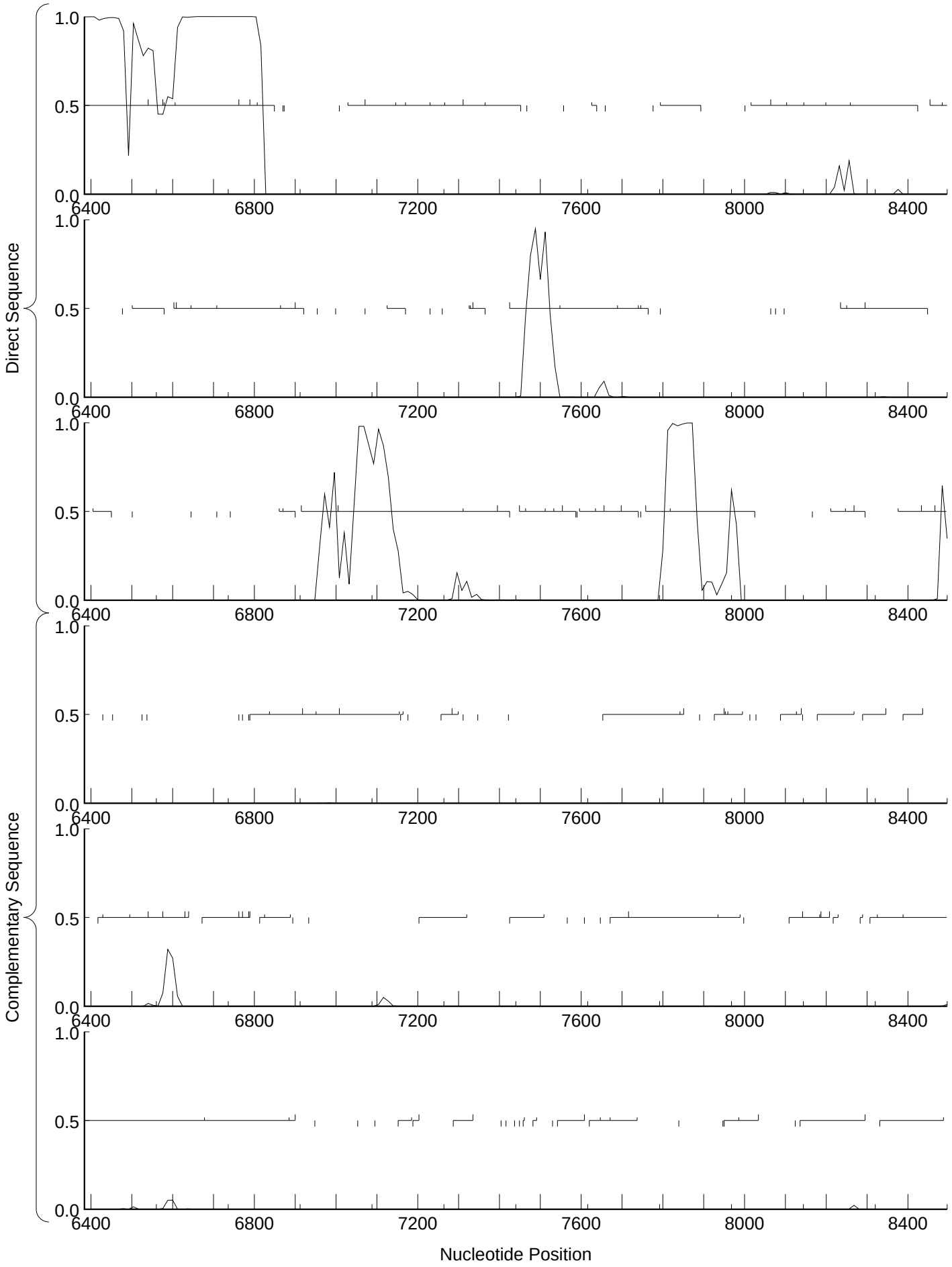
s complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 2/23

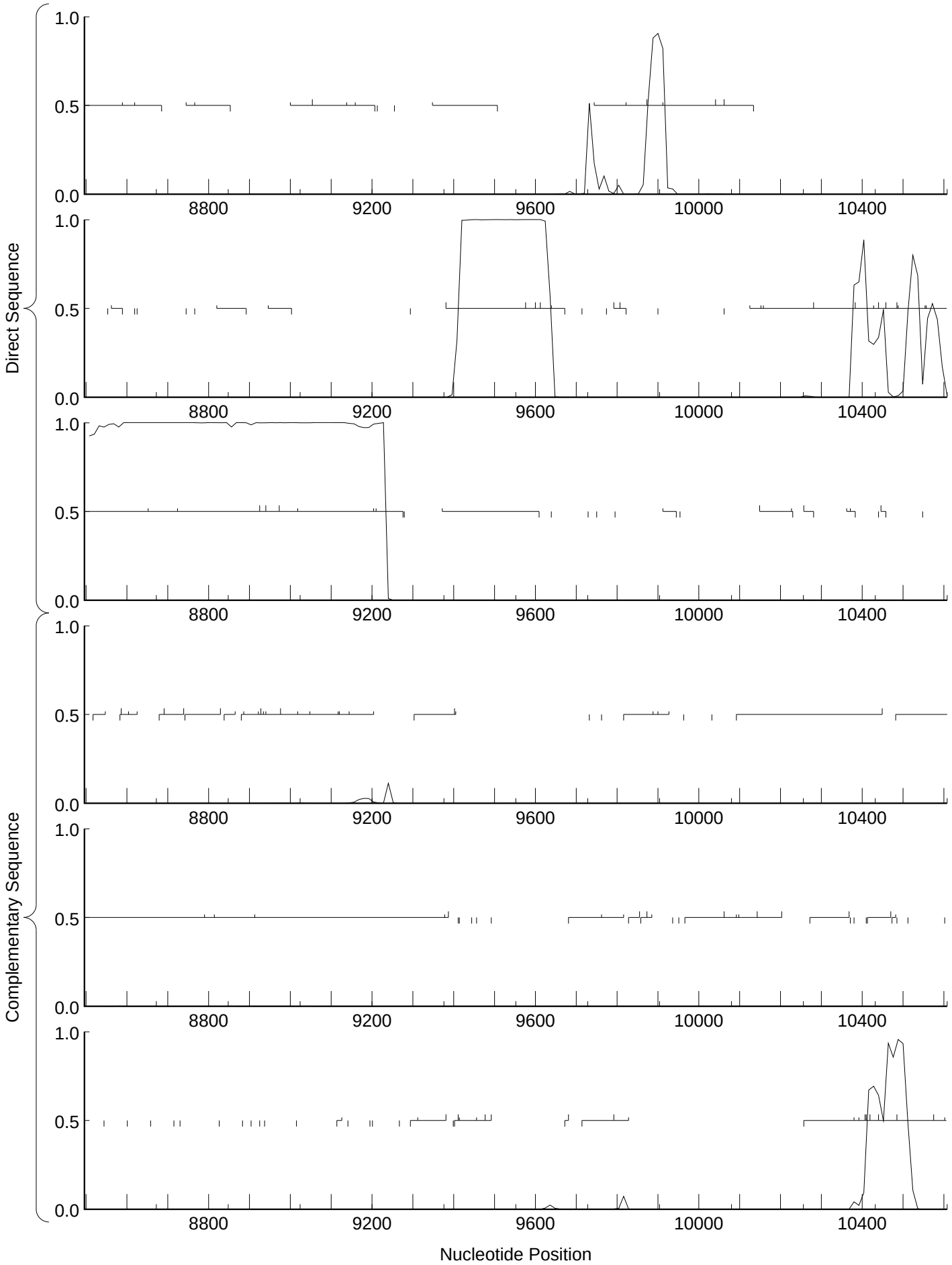



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
2400
2800
3200
3600
4000
Nucleotide Position

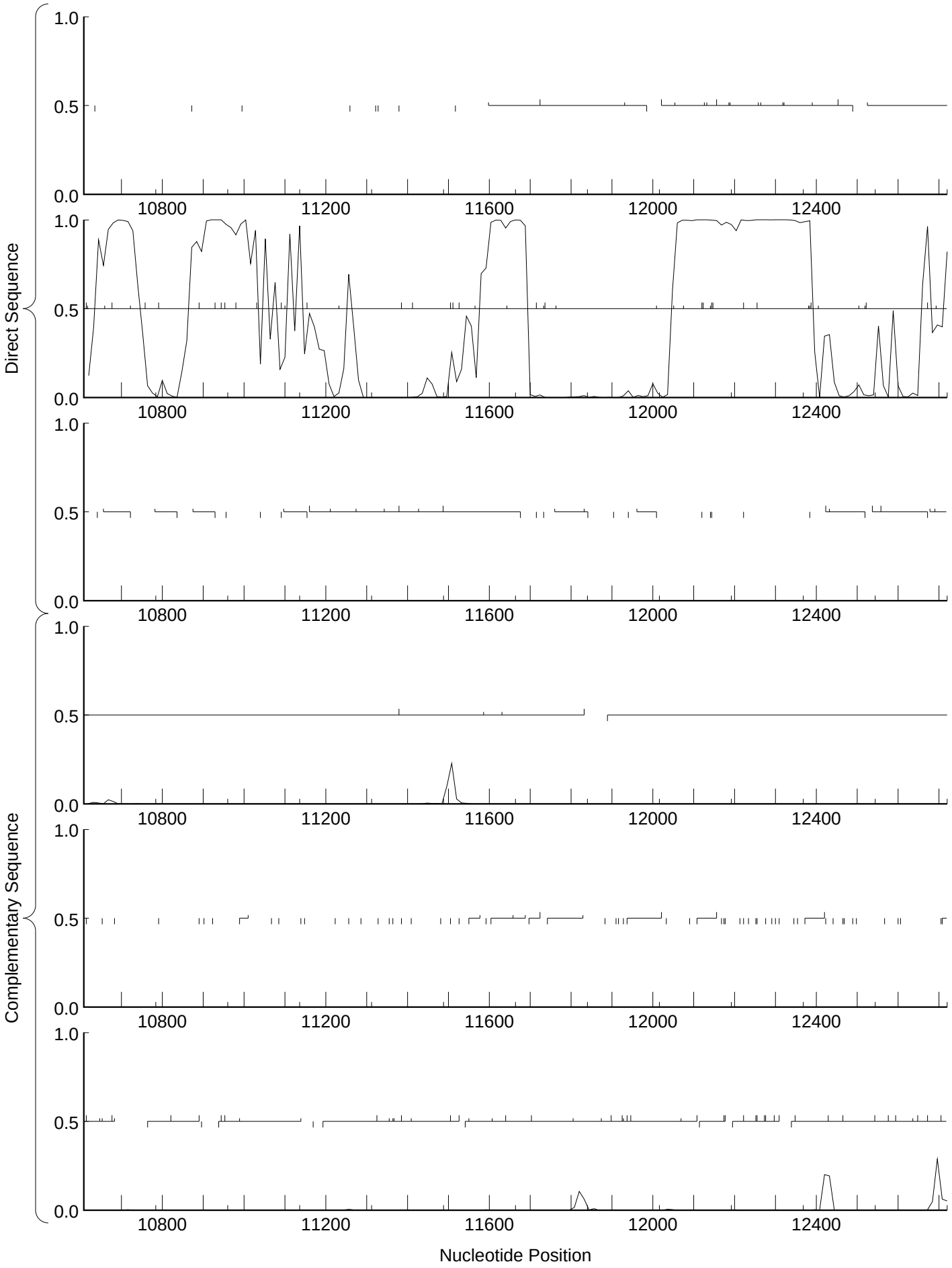

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
4400
4800
5200
5600
6000
Nucleotide Position

s complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 5/23

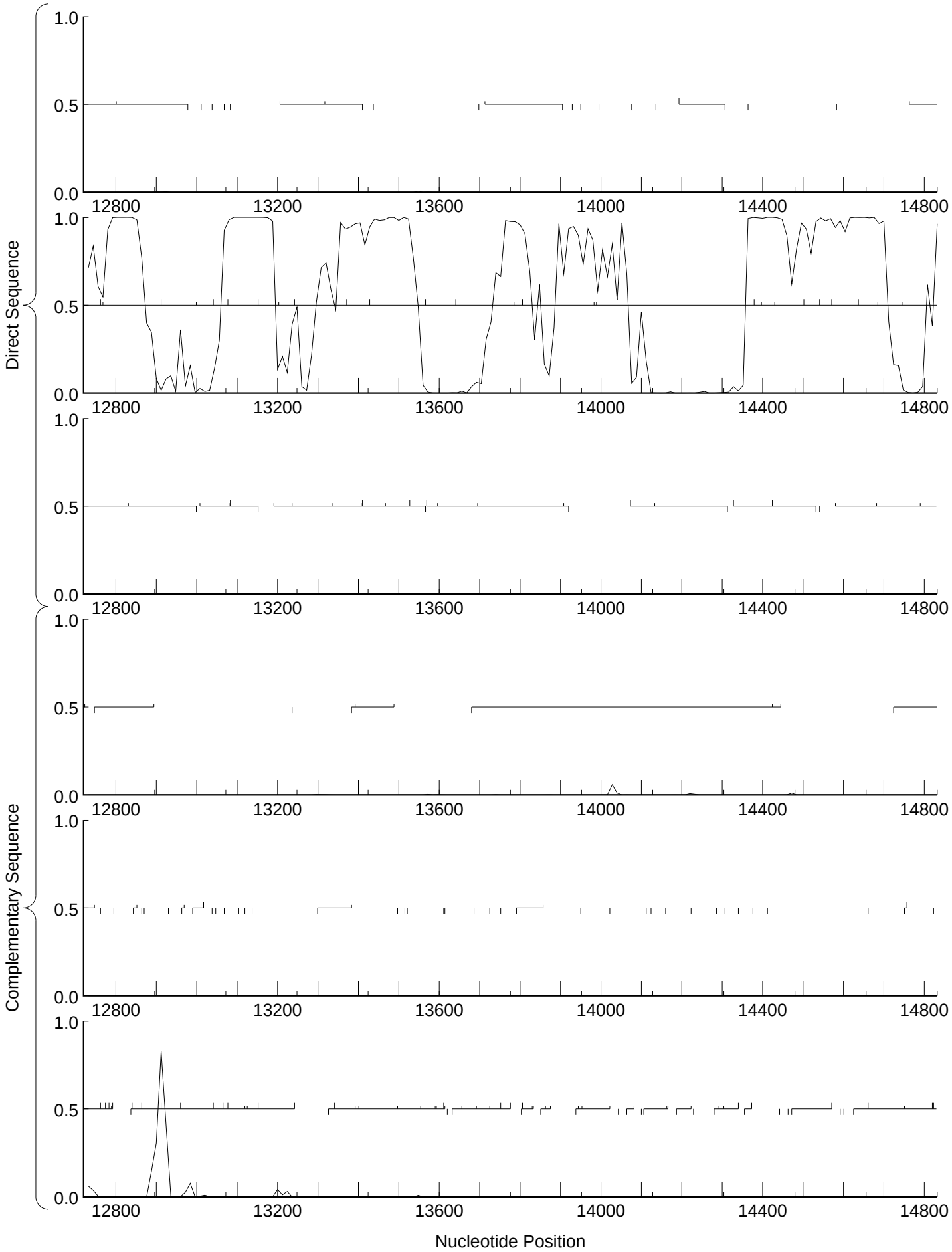



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
8800
9200
9600
10000
10400
Nucleotide Position

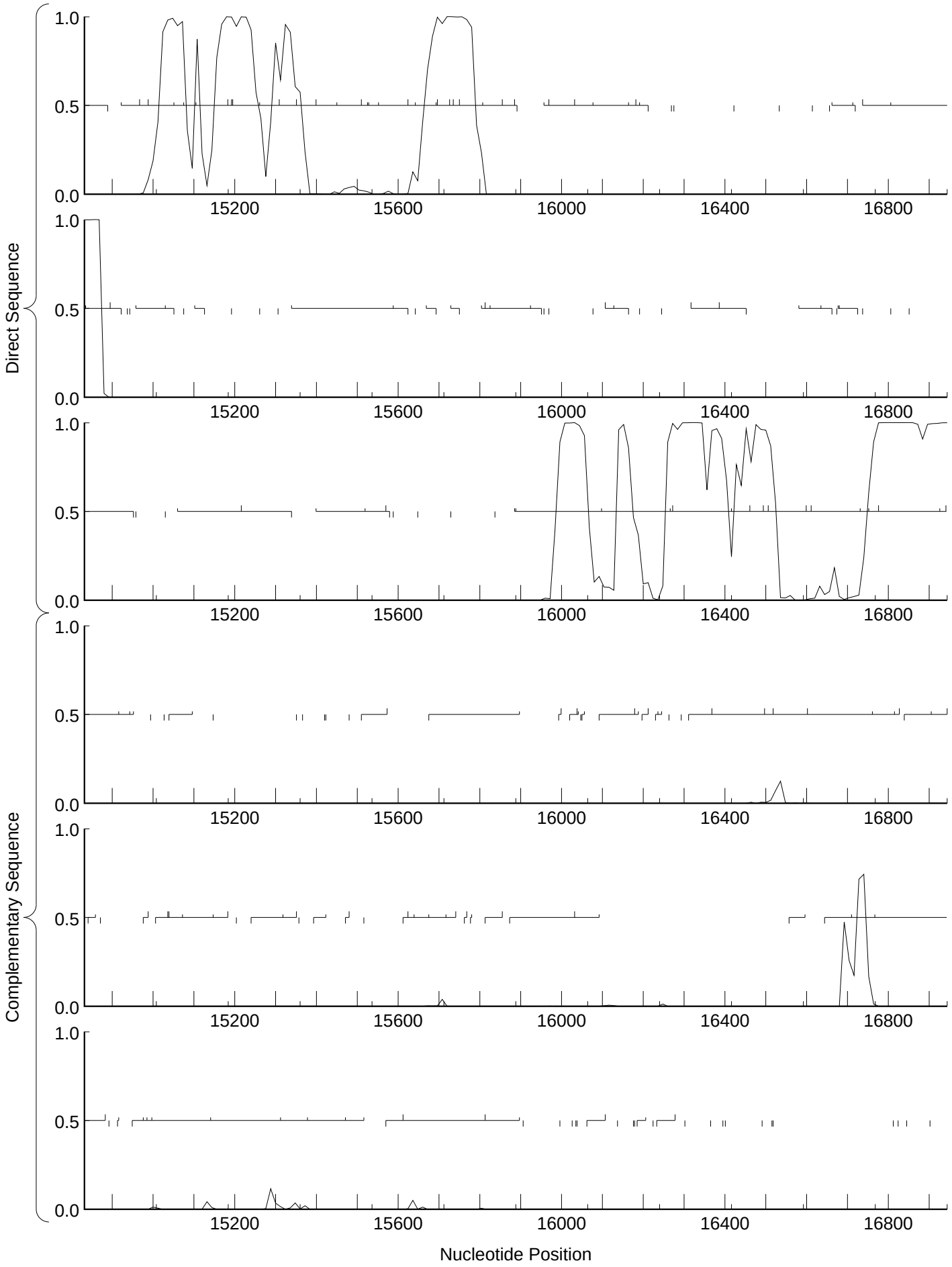
s complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 7/23



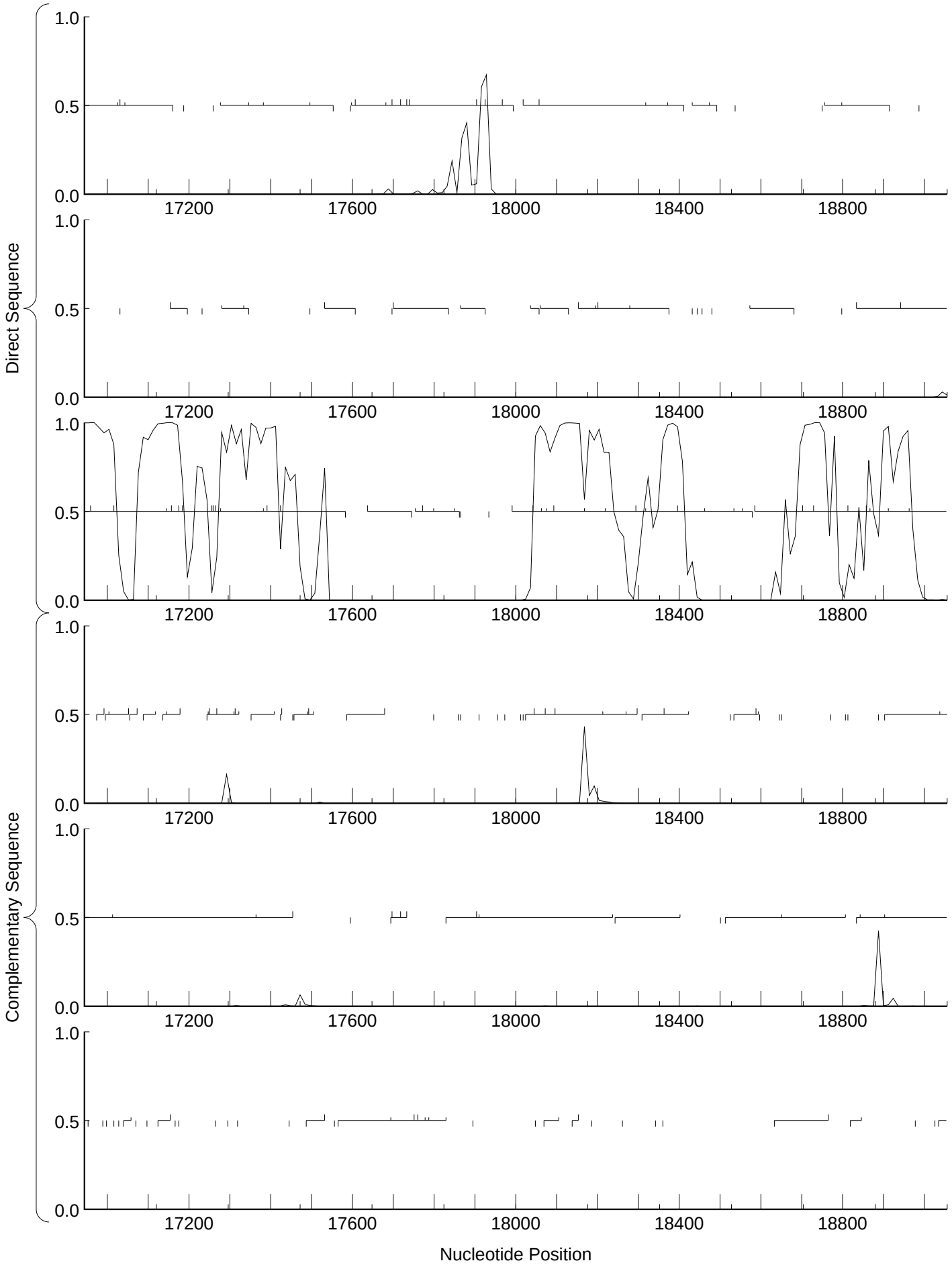
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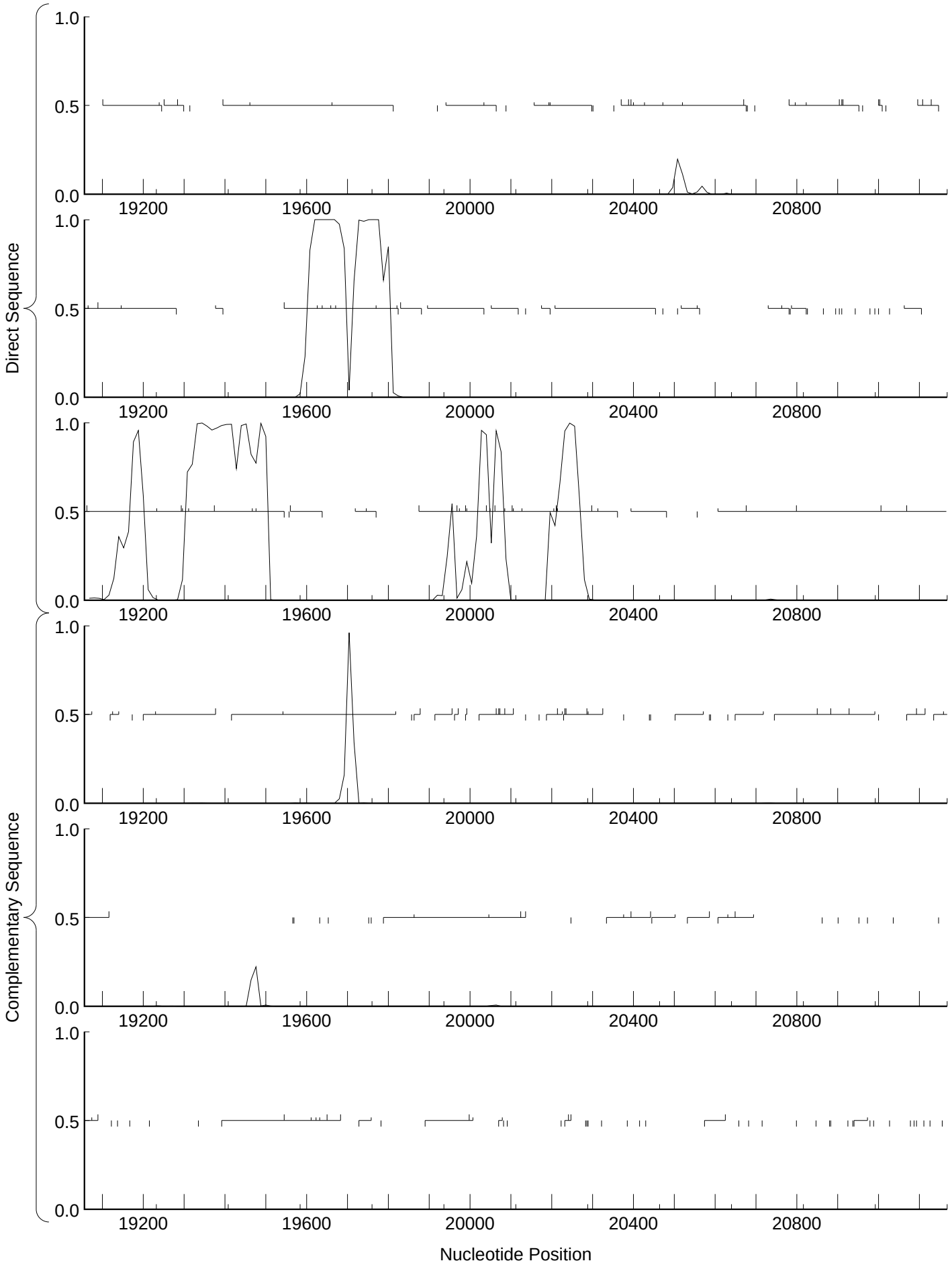
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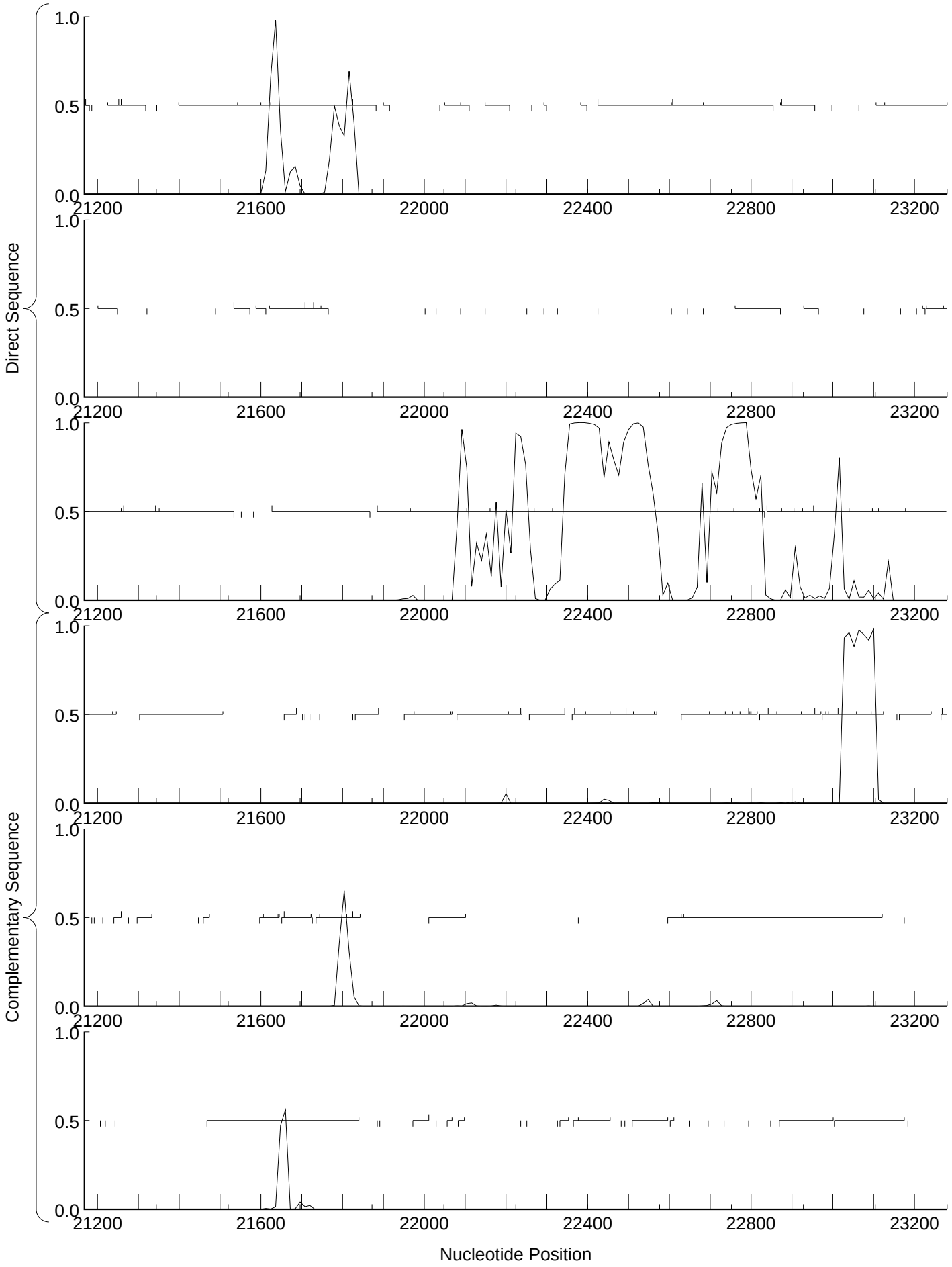
complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 10/23



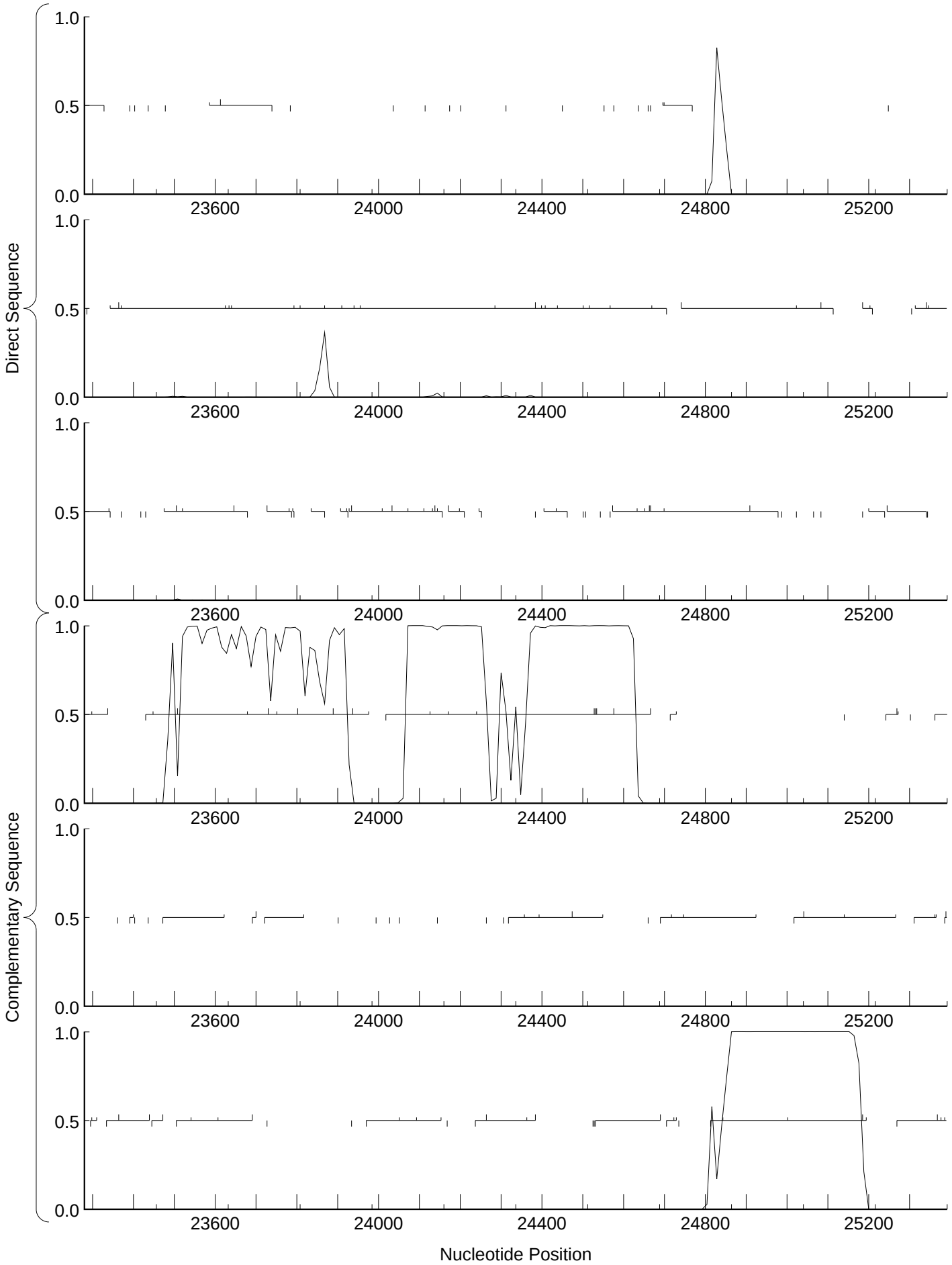
complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 11/23



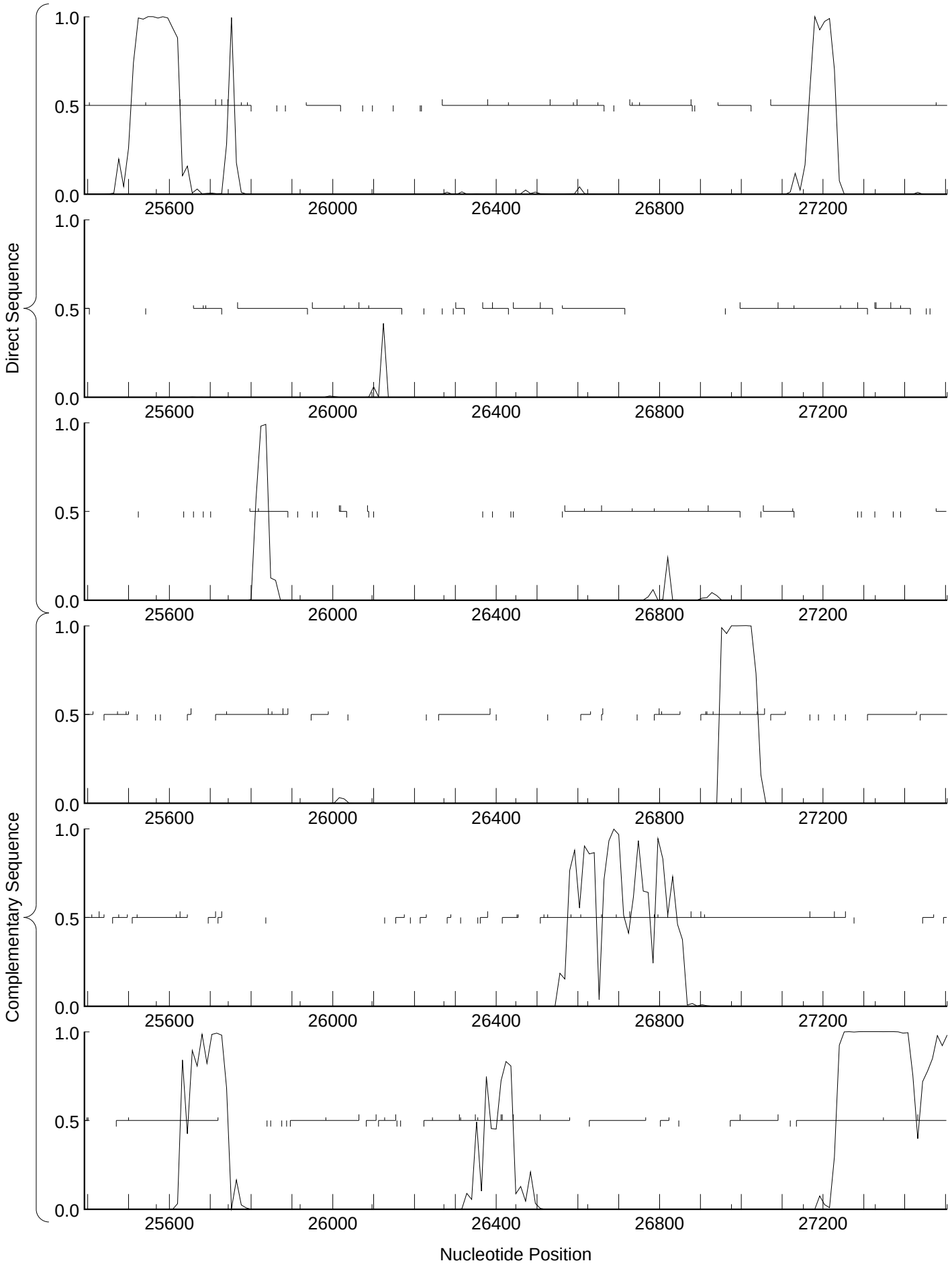
complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 12/23



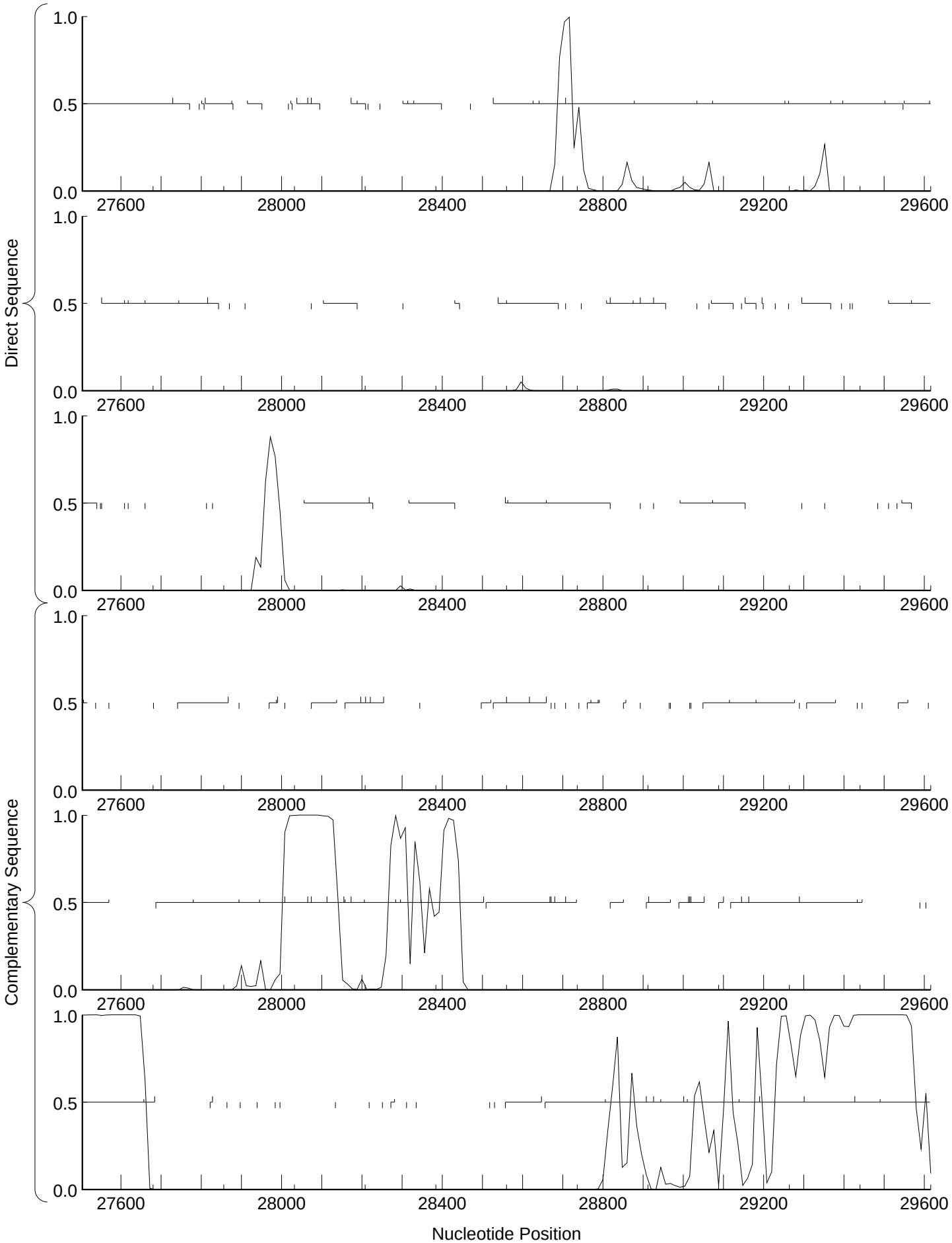
complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 13/23



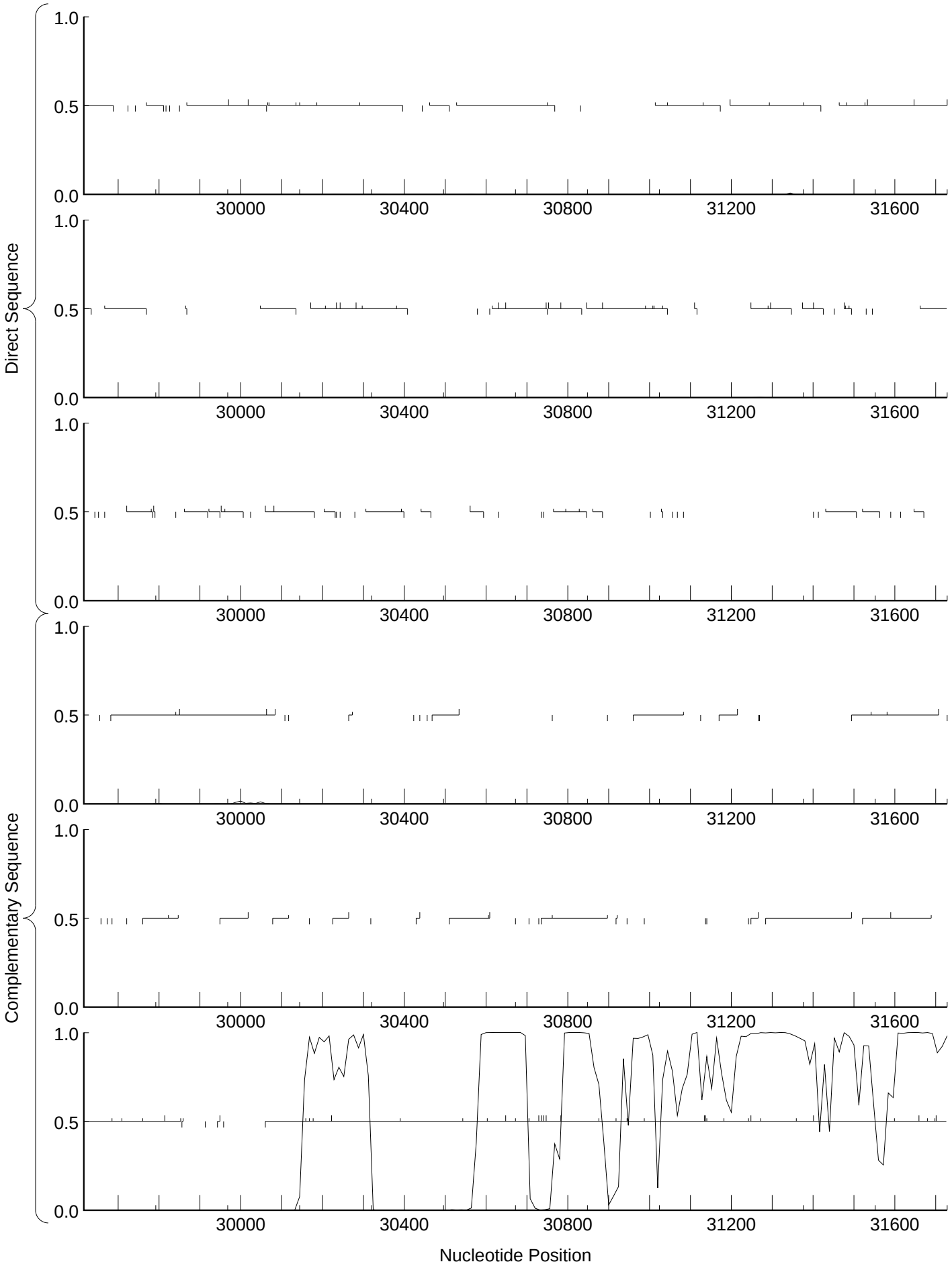
complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 14/23



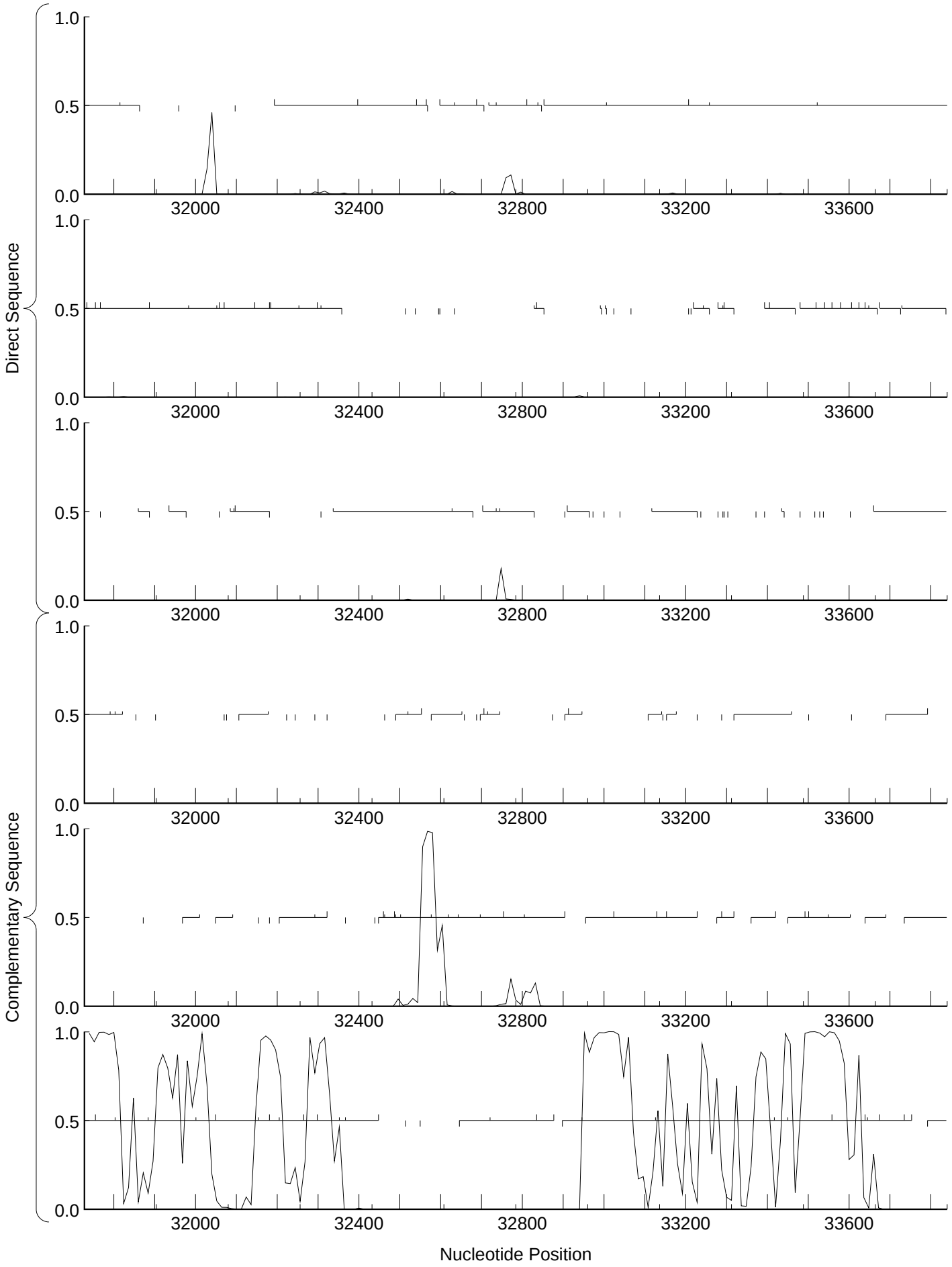
complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 15/23



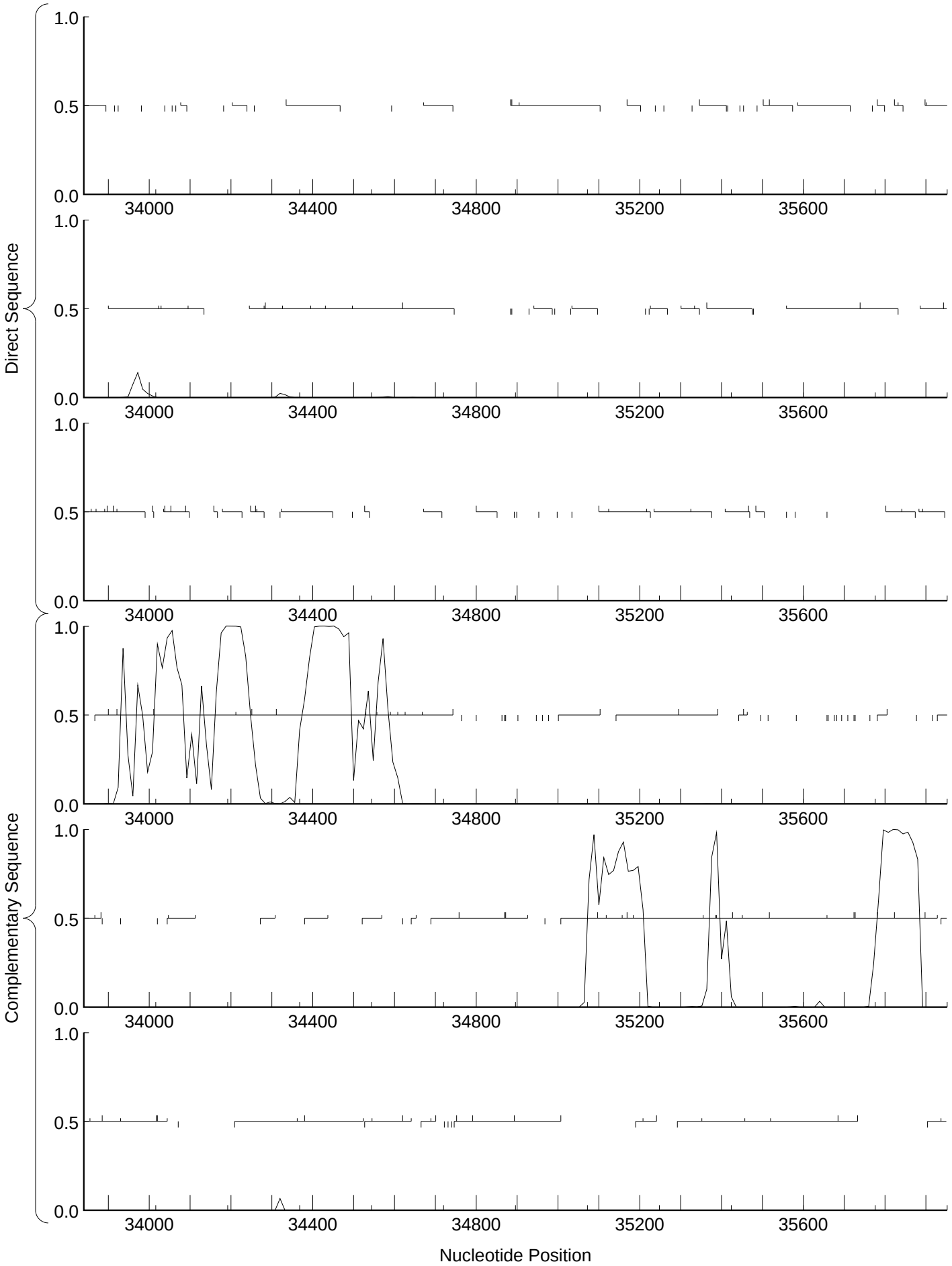
complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 16/23



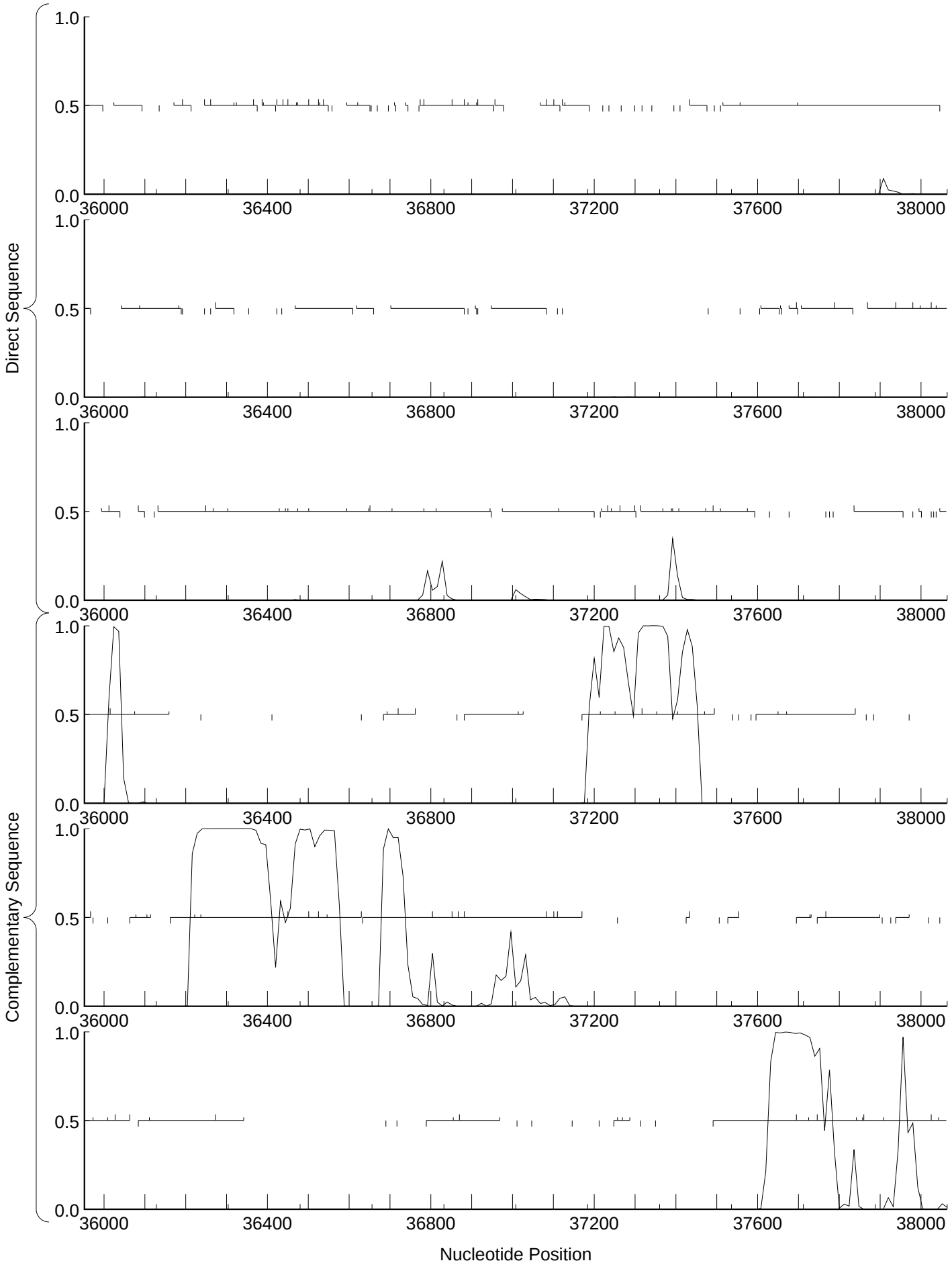
complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 17/23



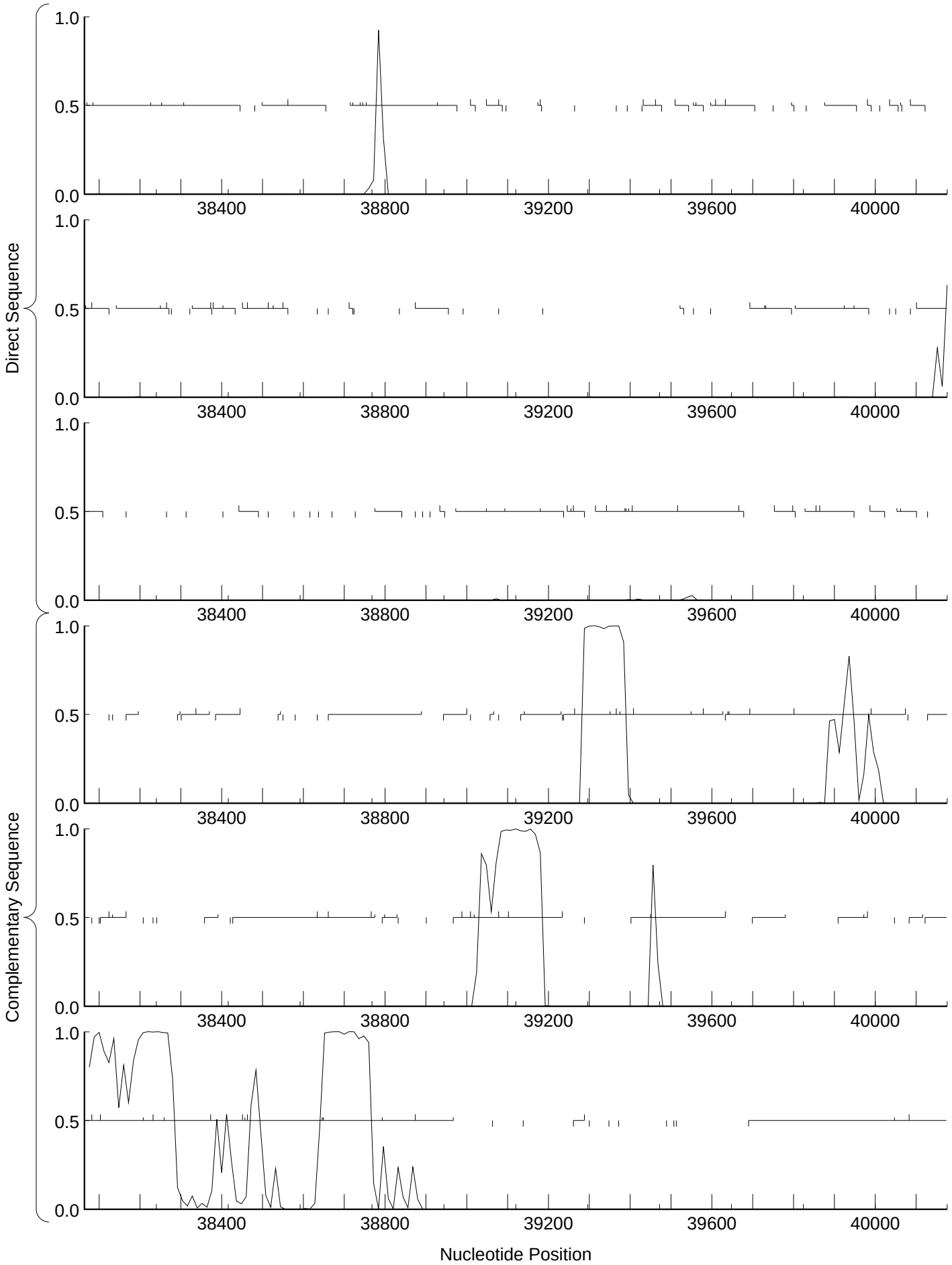
complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 18/23



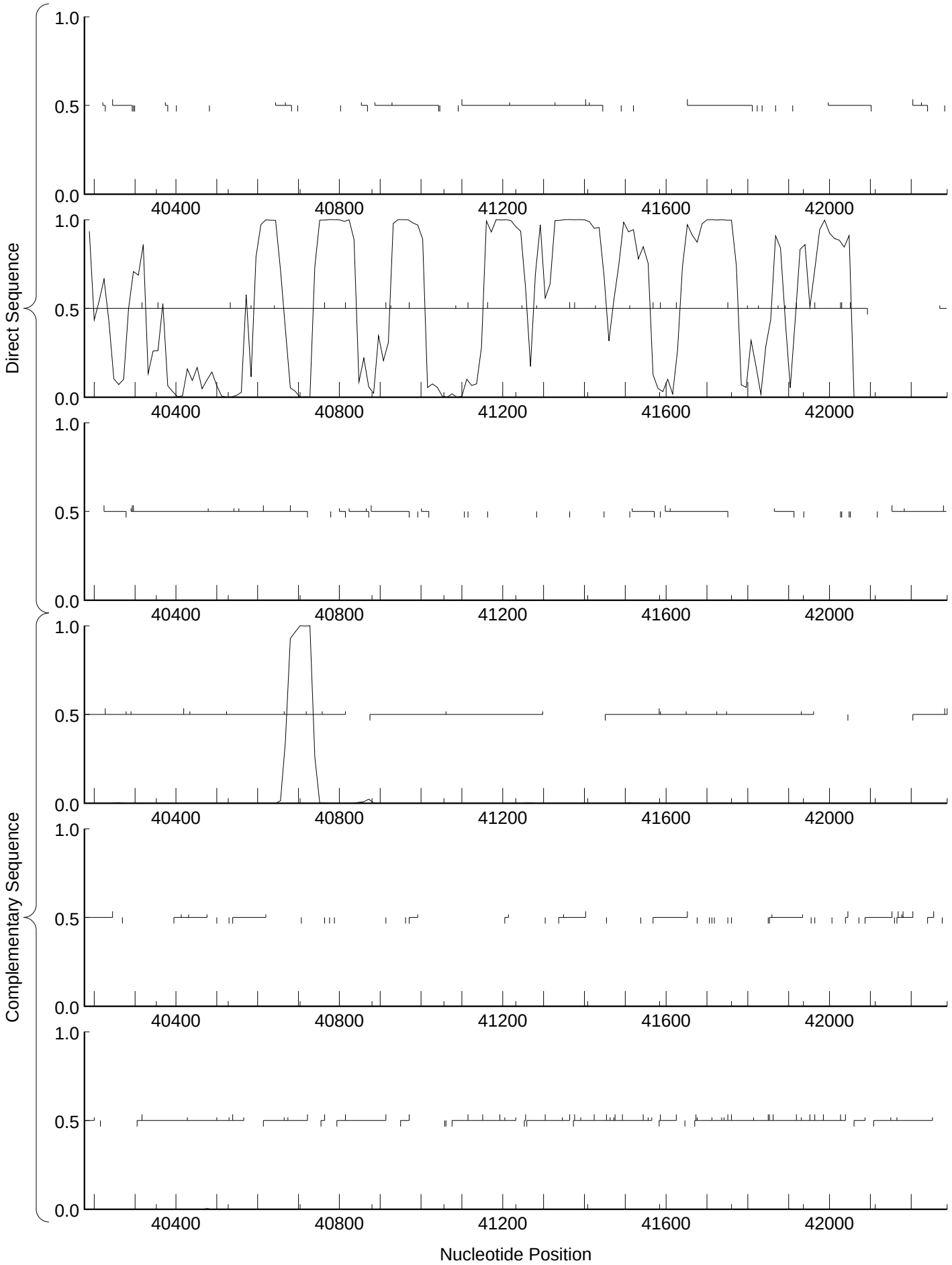
complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 19/23



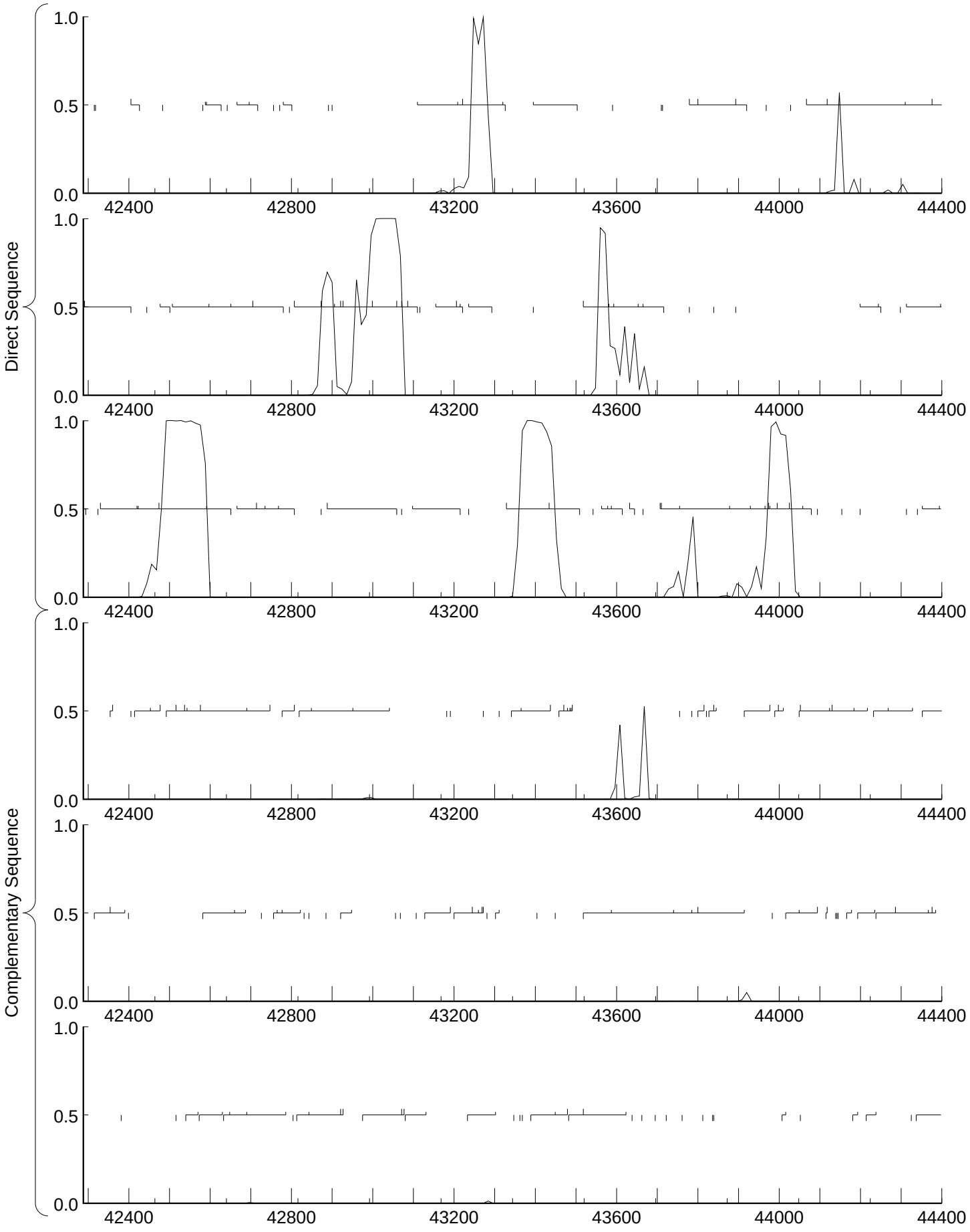
complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 20/23



complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 21/23



complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 22/23



complete sequence, 45917 bp including 10-base 3’ overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 23/23

