GeneMark

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

PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Eponine complete sequence, 58678 bp including 11-base 3’ overhang (CTCG
Analysis Date    :  3/29/24 at 3:56:39
Pages            :  29
Sequence Length  :  58678 bp
GC Content       :   67.42%

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

PS-Version       :  1.2

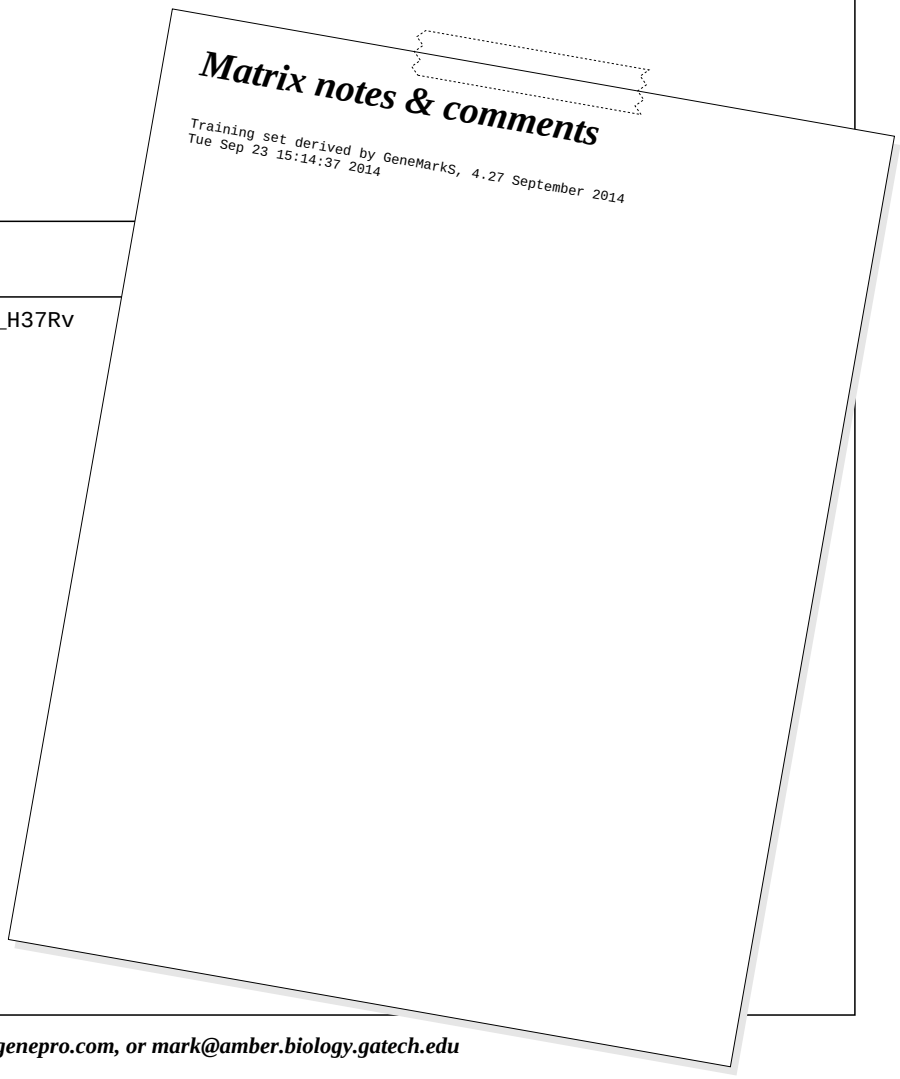
GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
```

MATRIX INFORMATION

```
Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

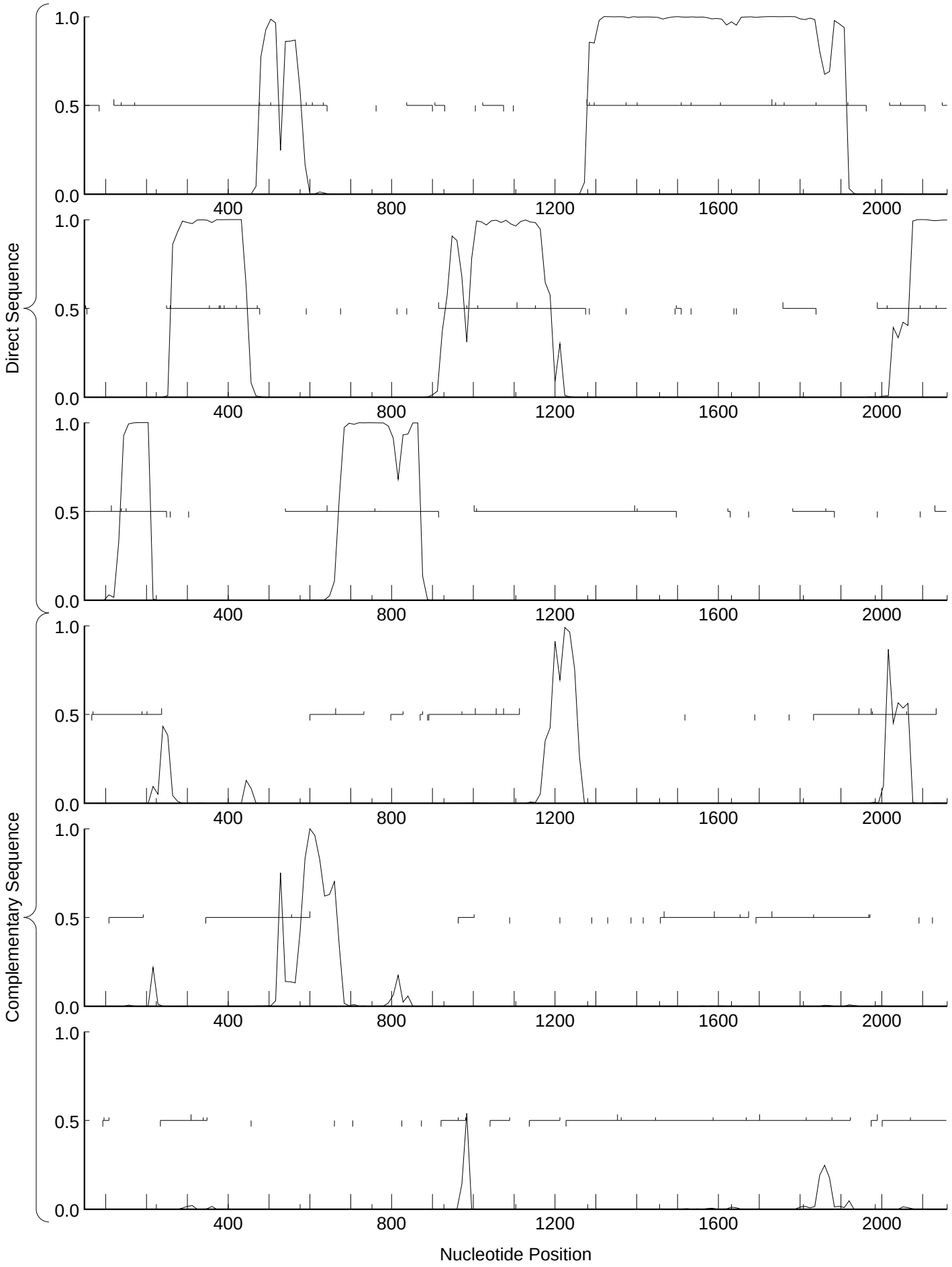
Order      :  4


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

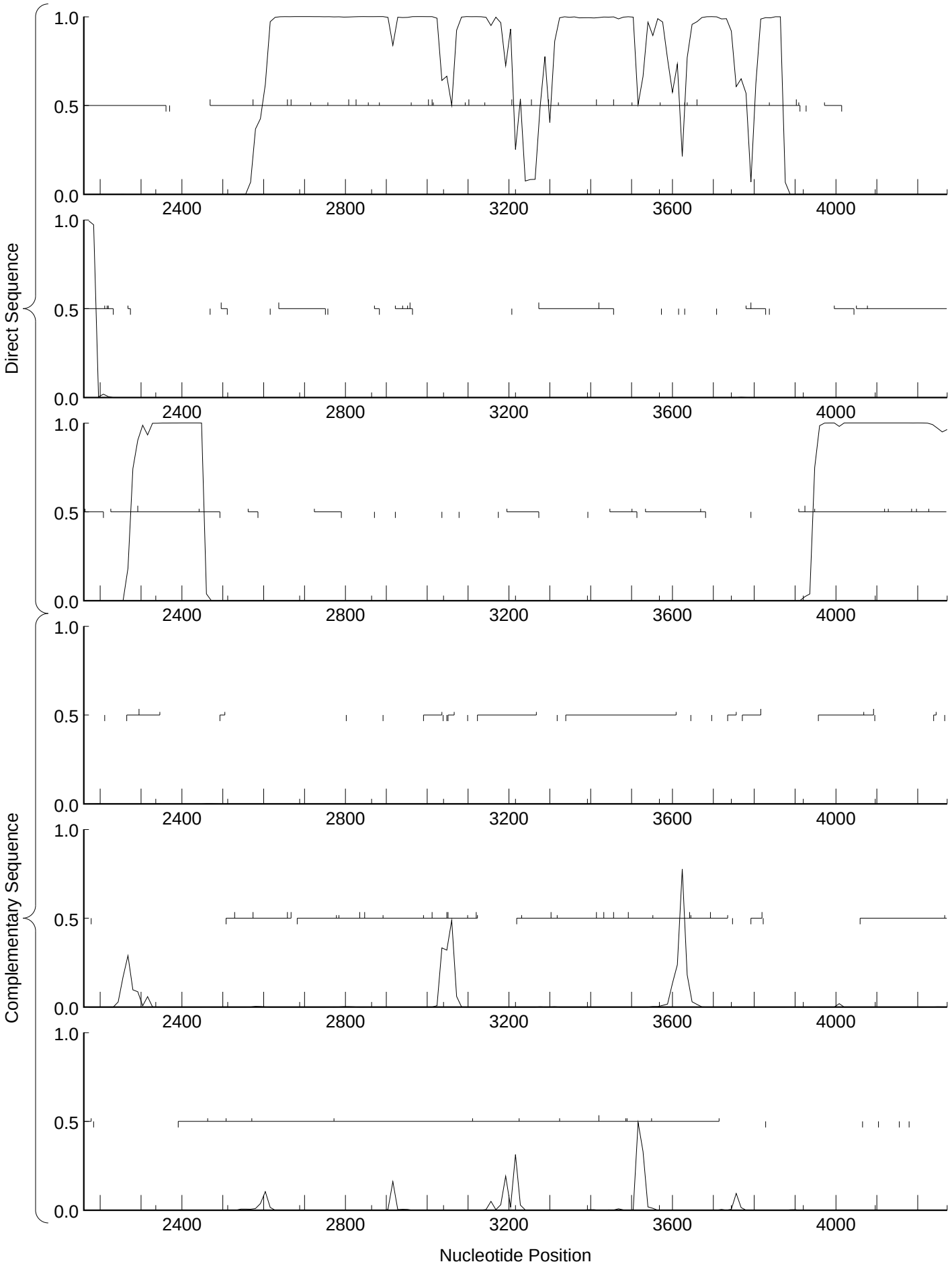


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

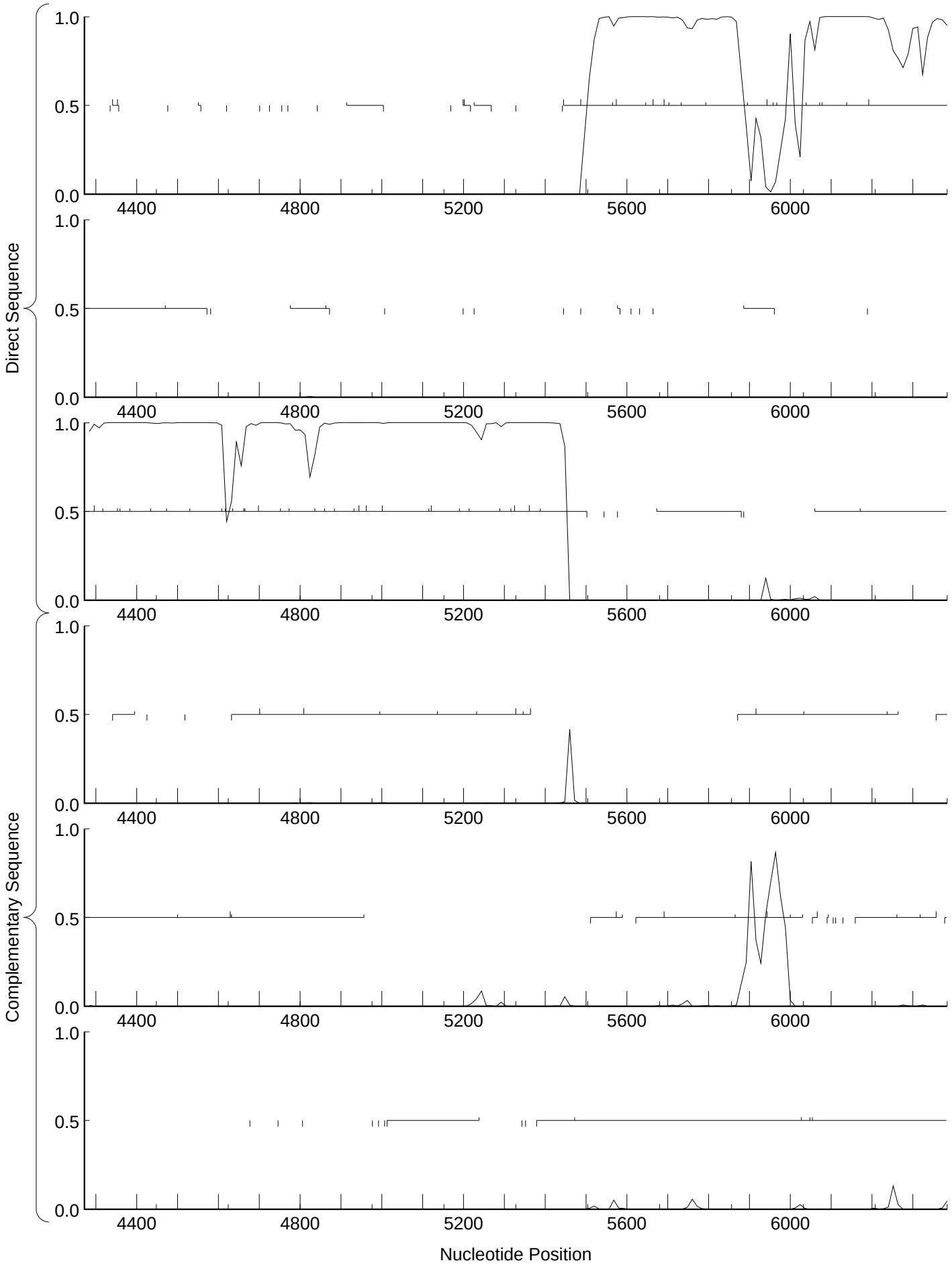
complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 2/29



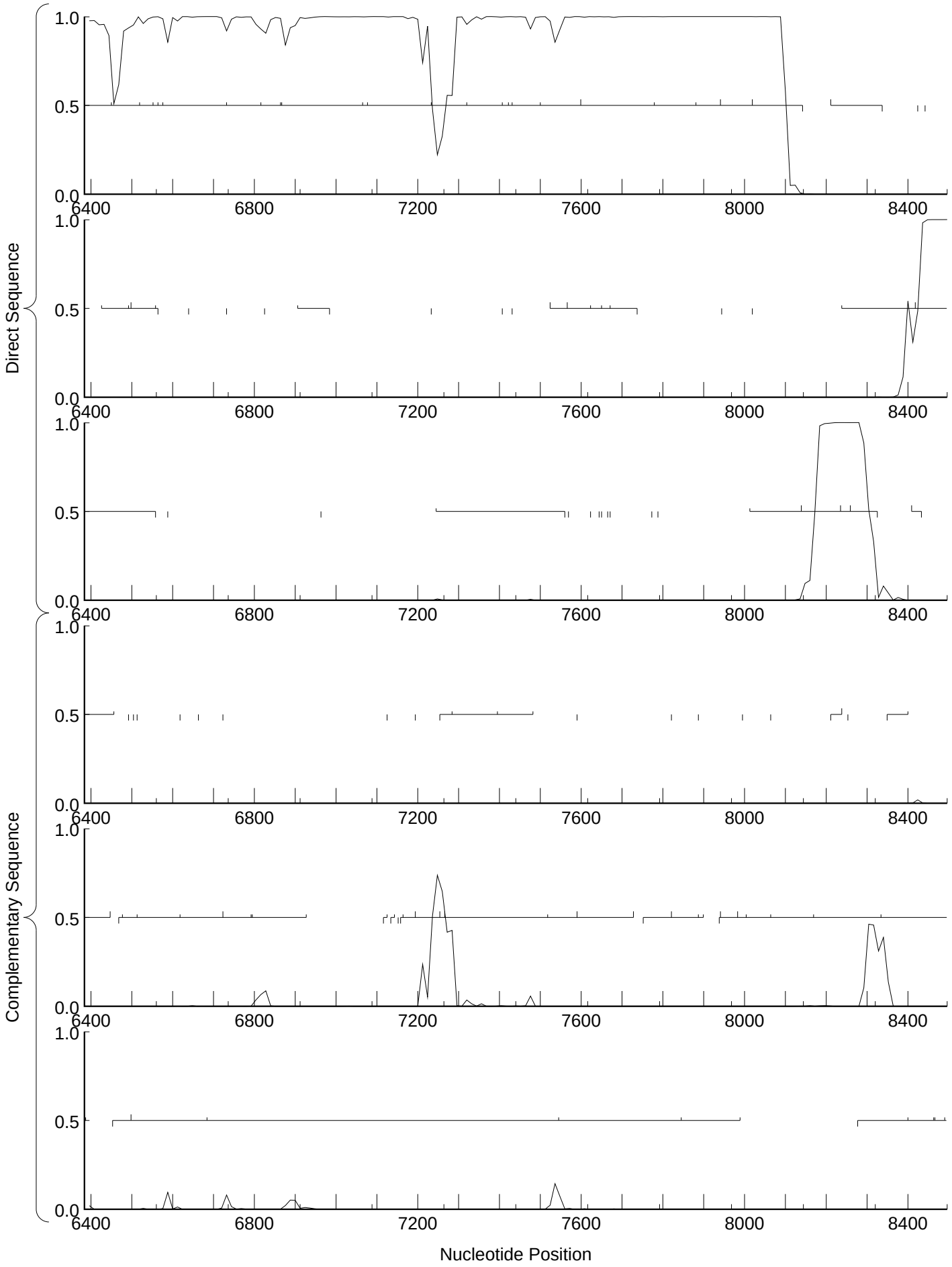
complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 3/29



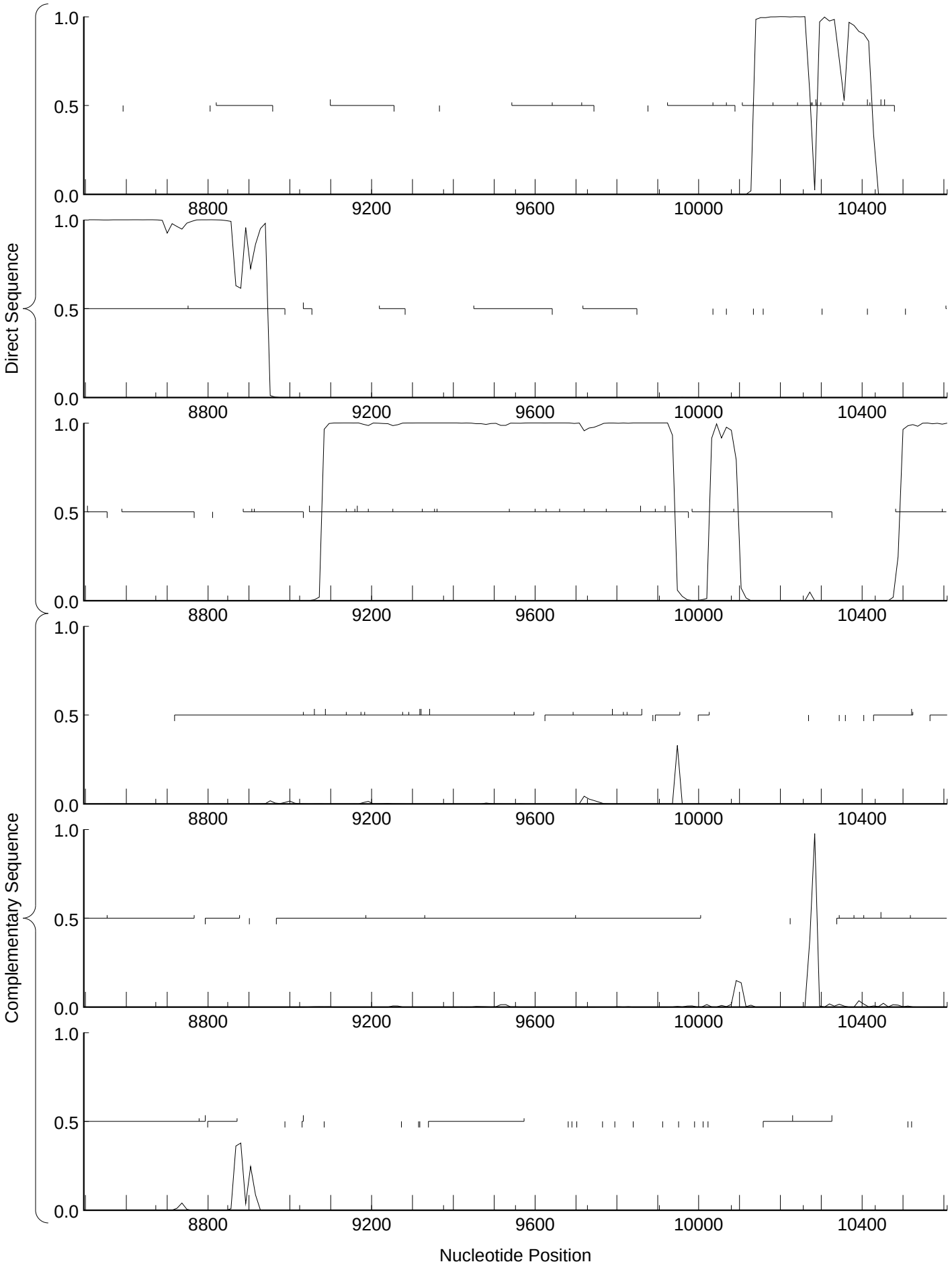
complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 4/29



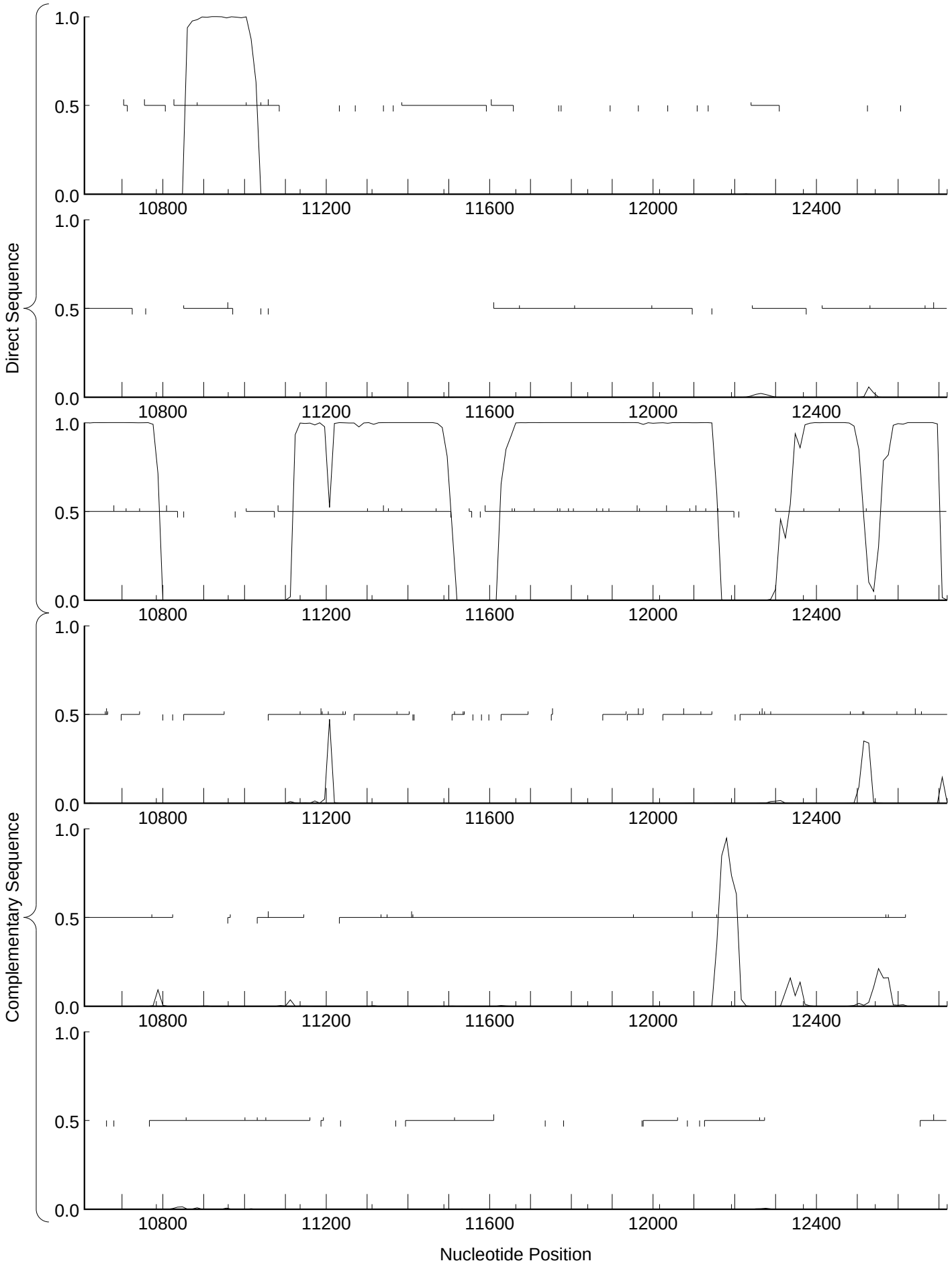
complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 5/29



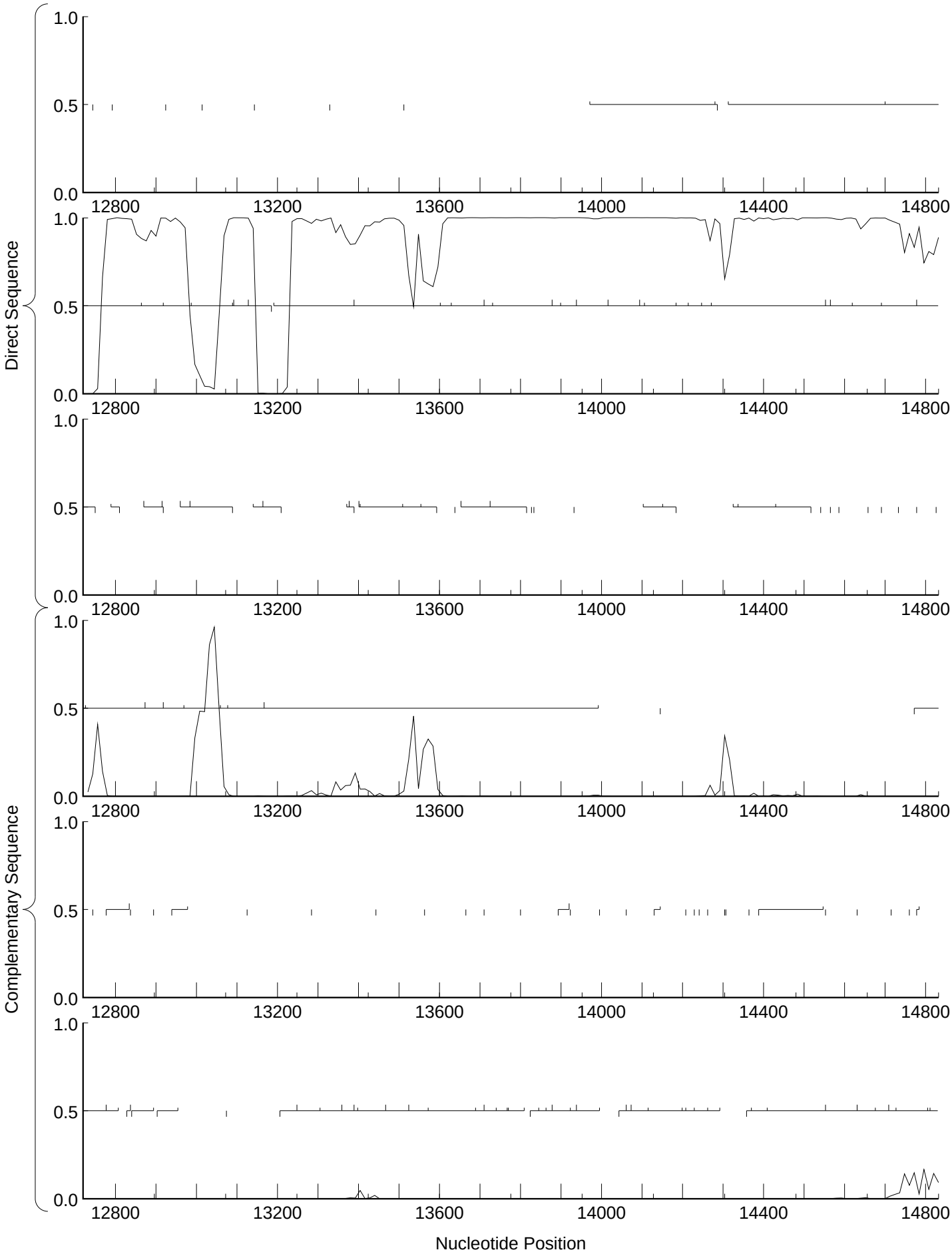
complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 6/29



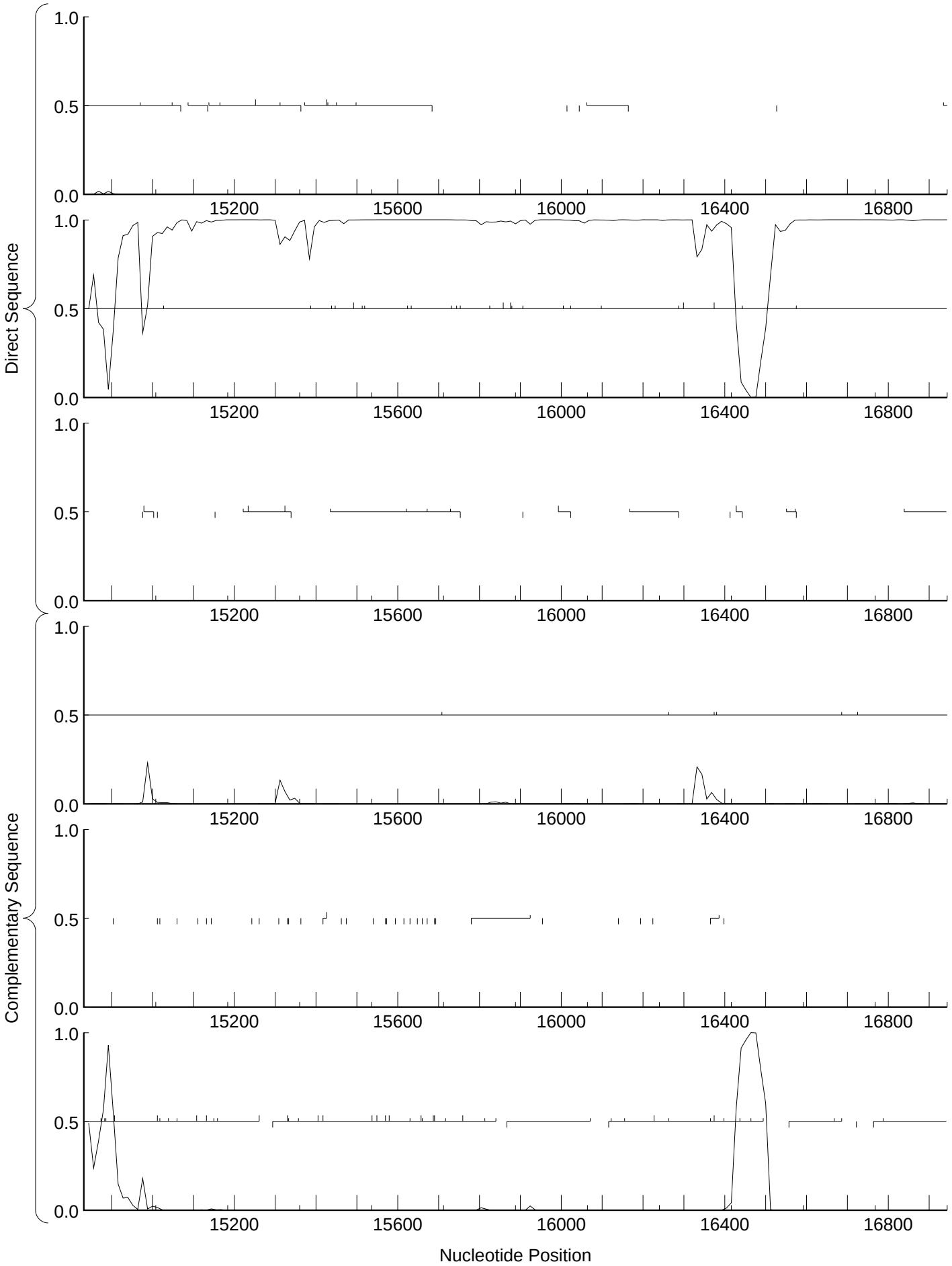
complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 7/29



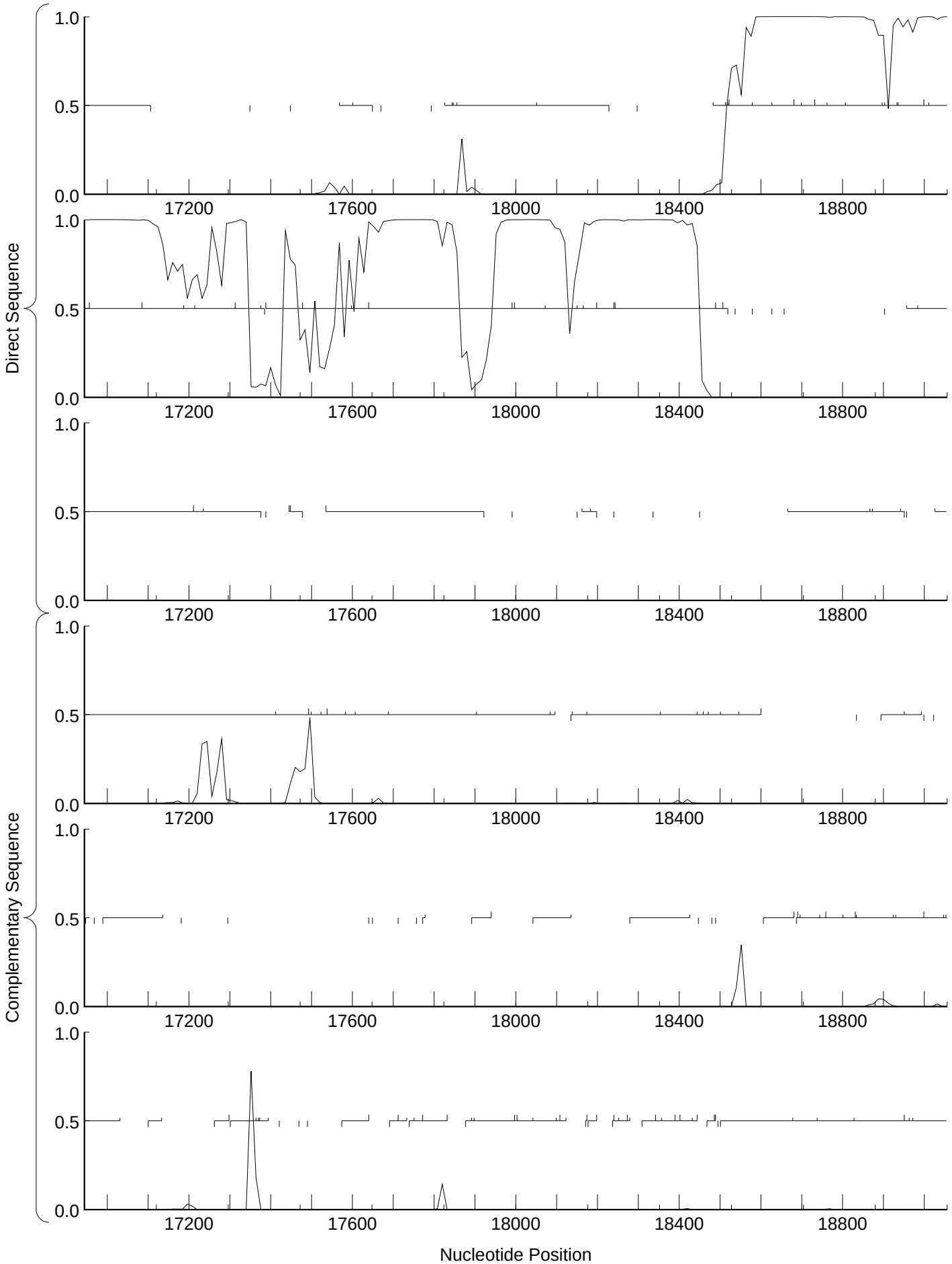
complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 8/29



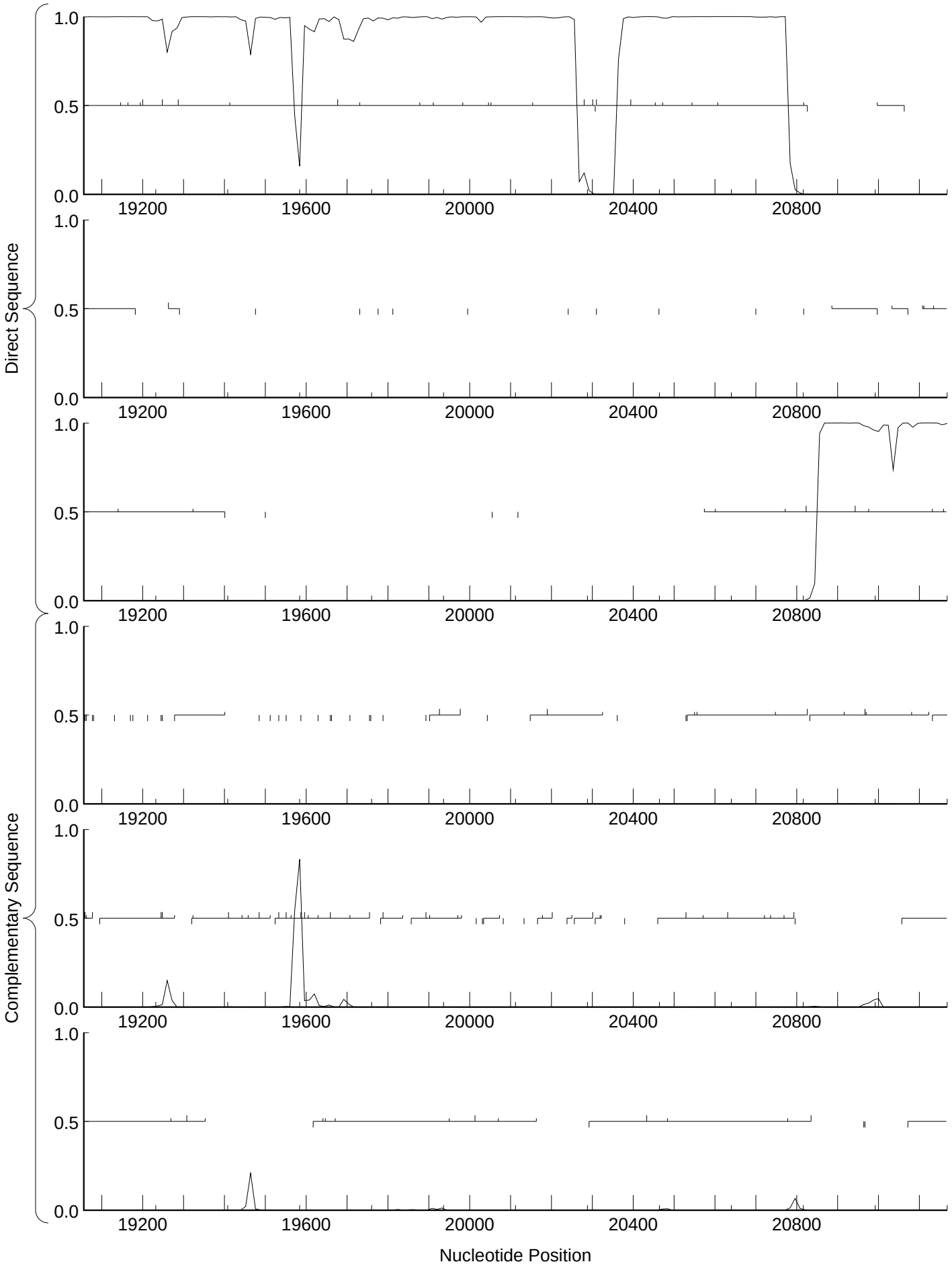
complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 9/29



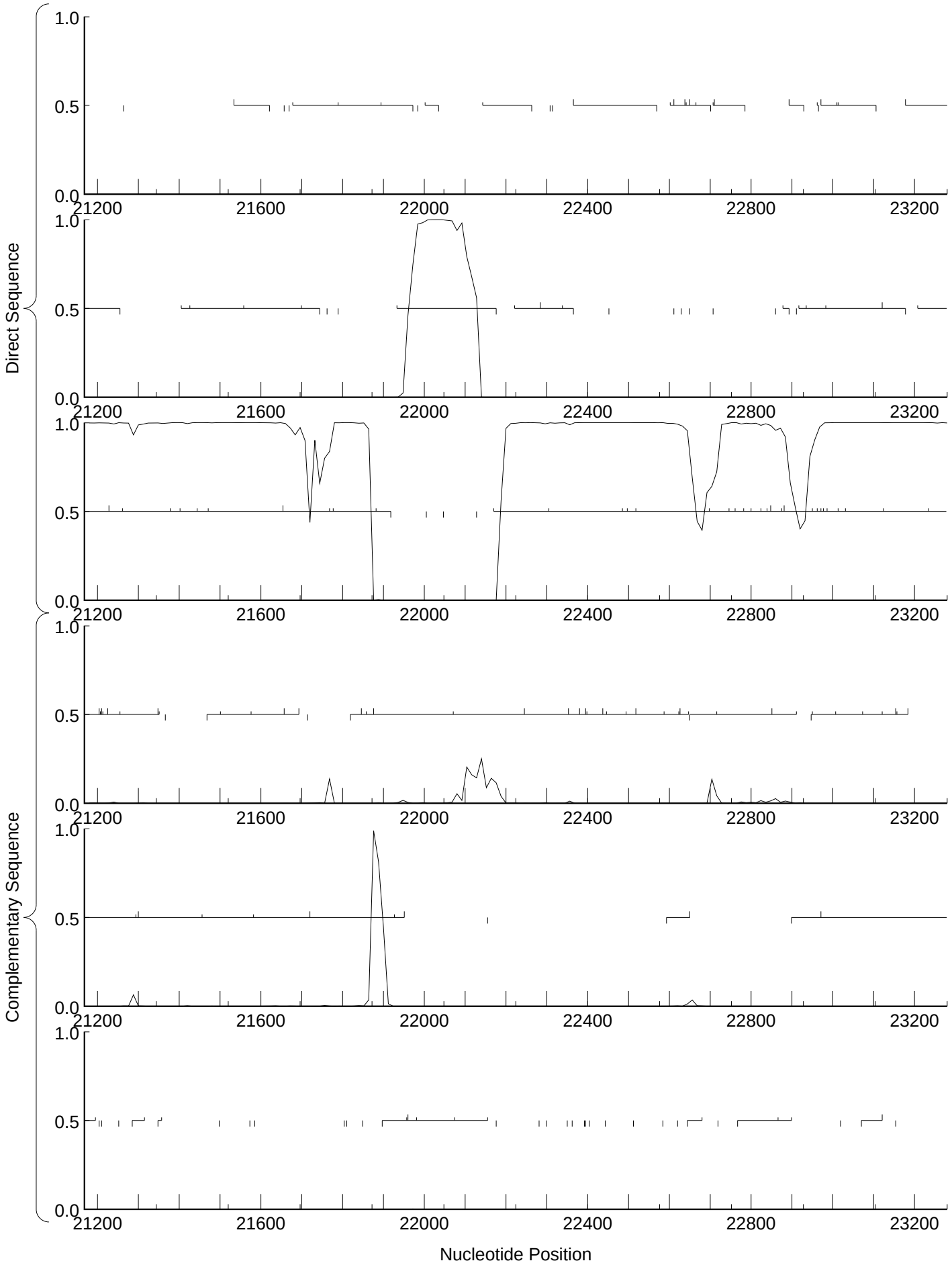
complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 10/29



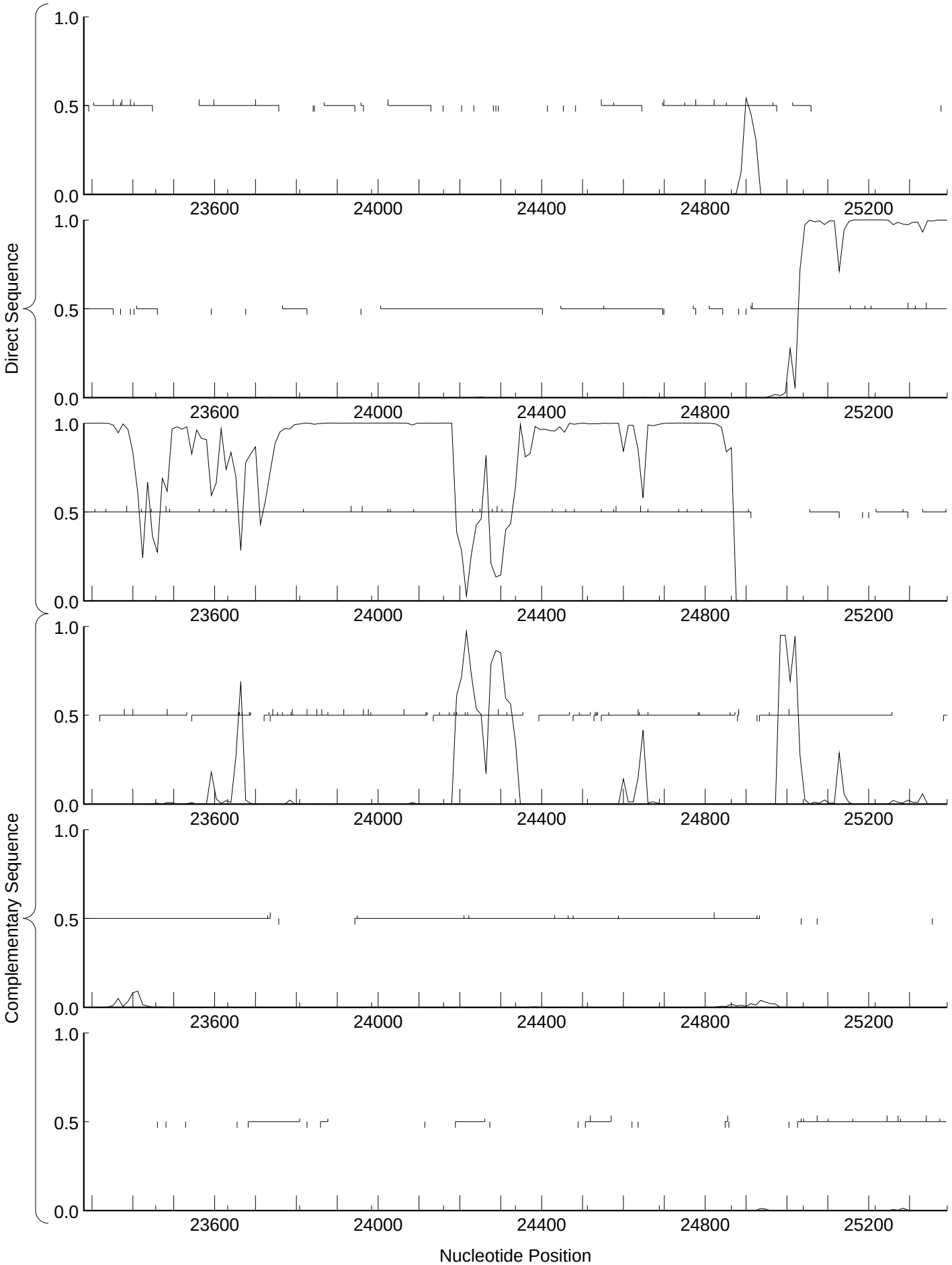
complete sequence, 58678 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 11/29

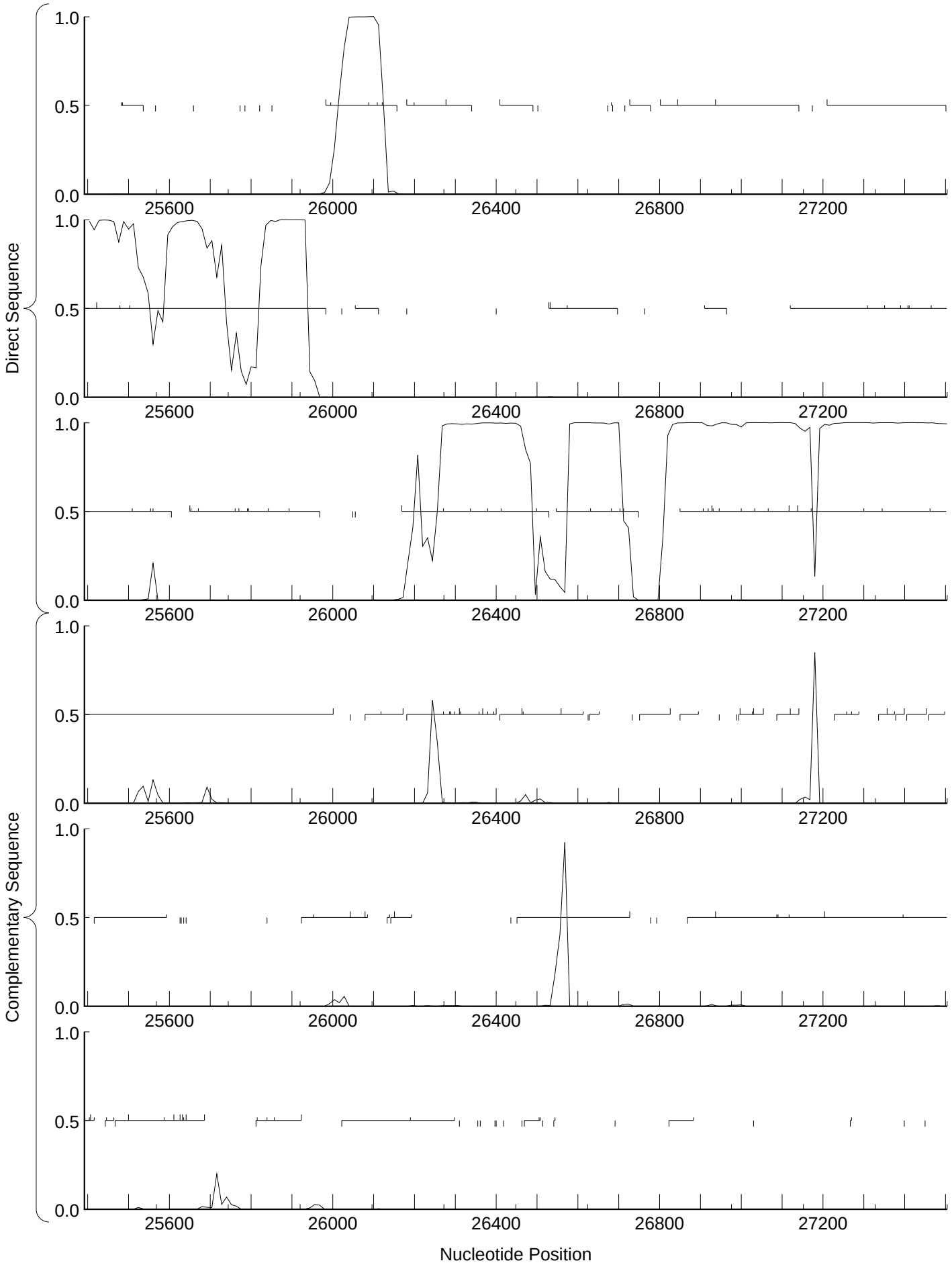


complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 12/29

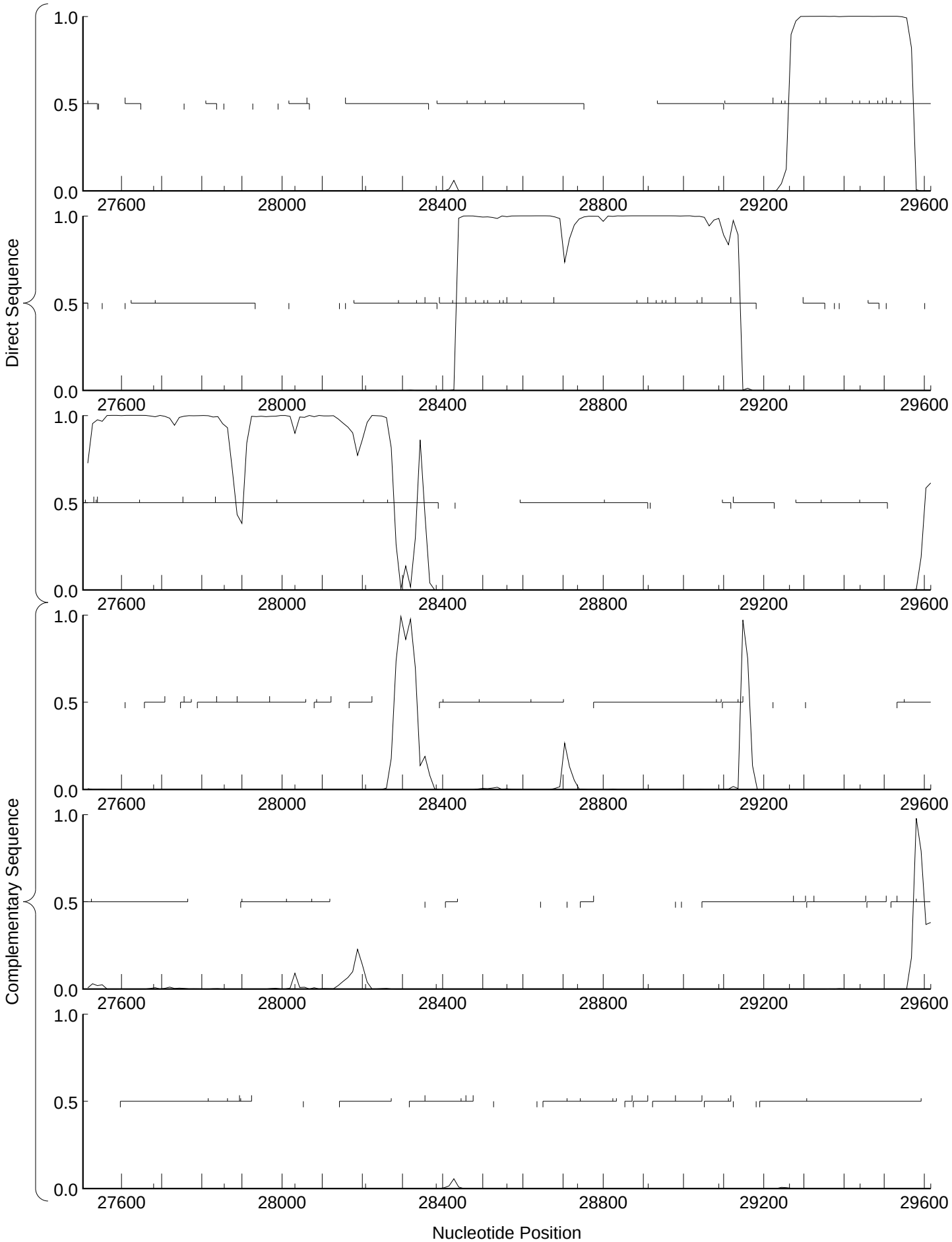


complete sequence, 58678 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 13/29

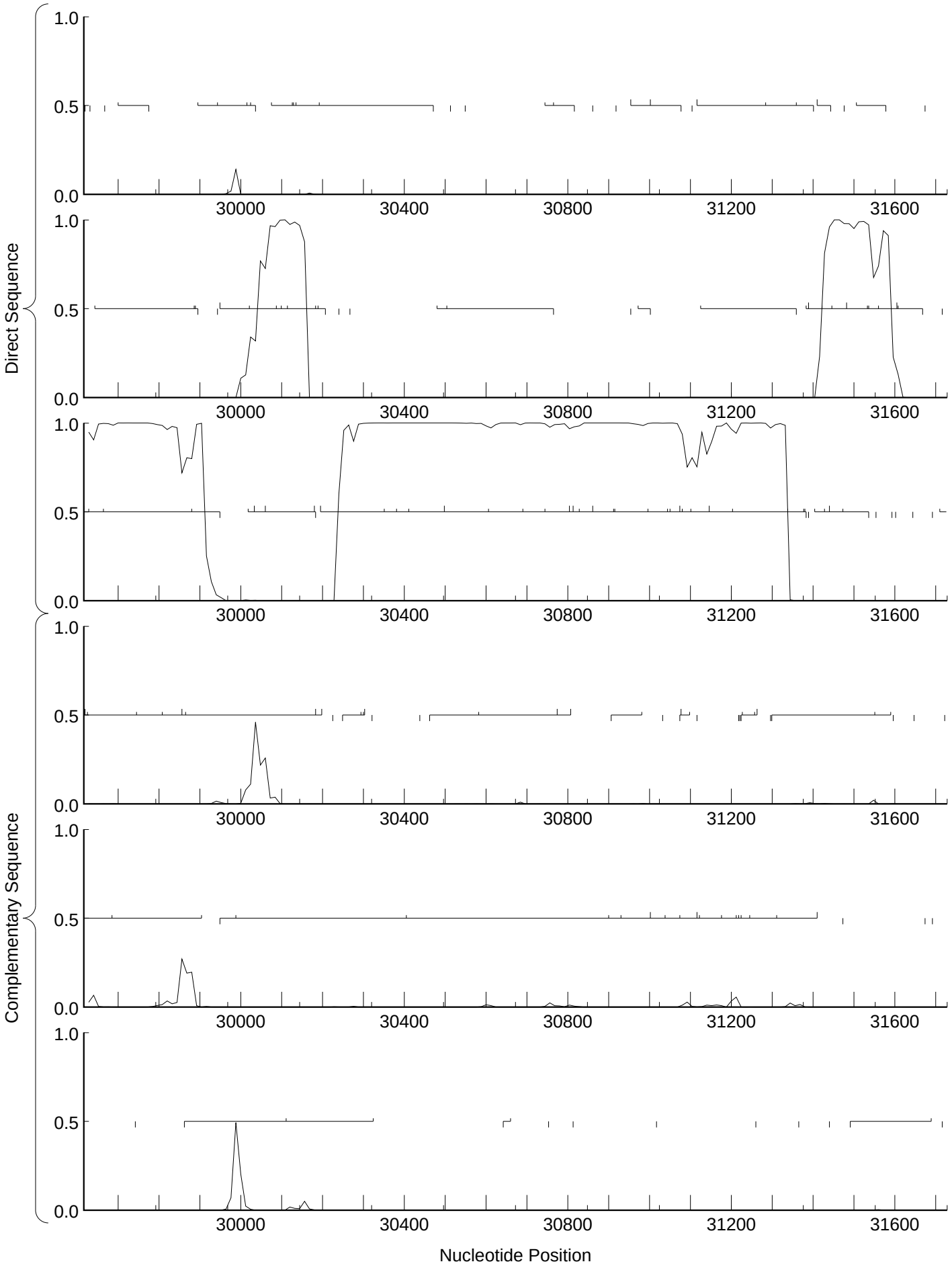


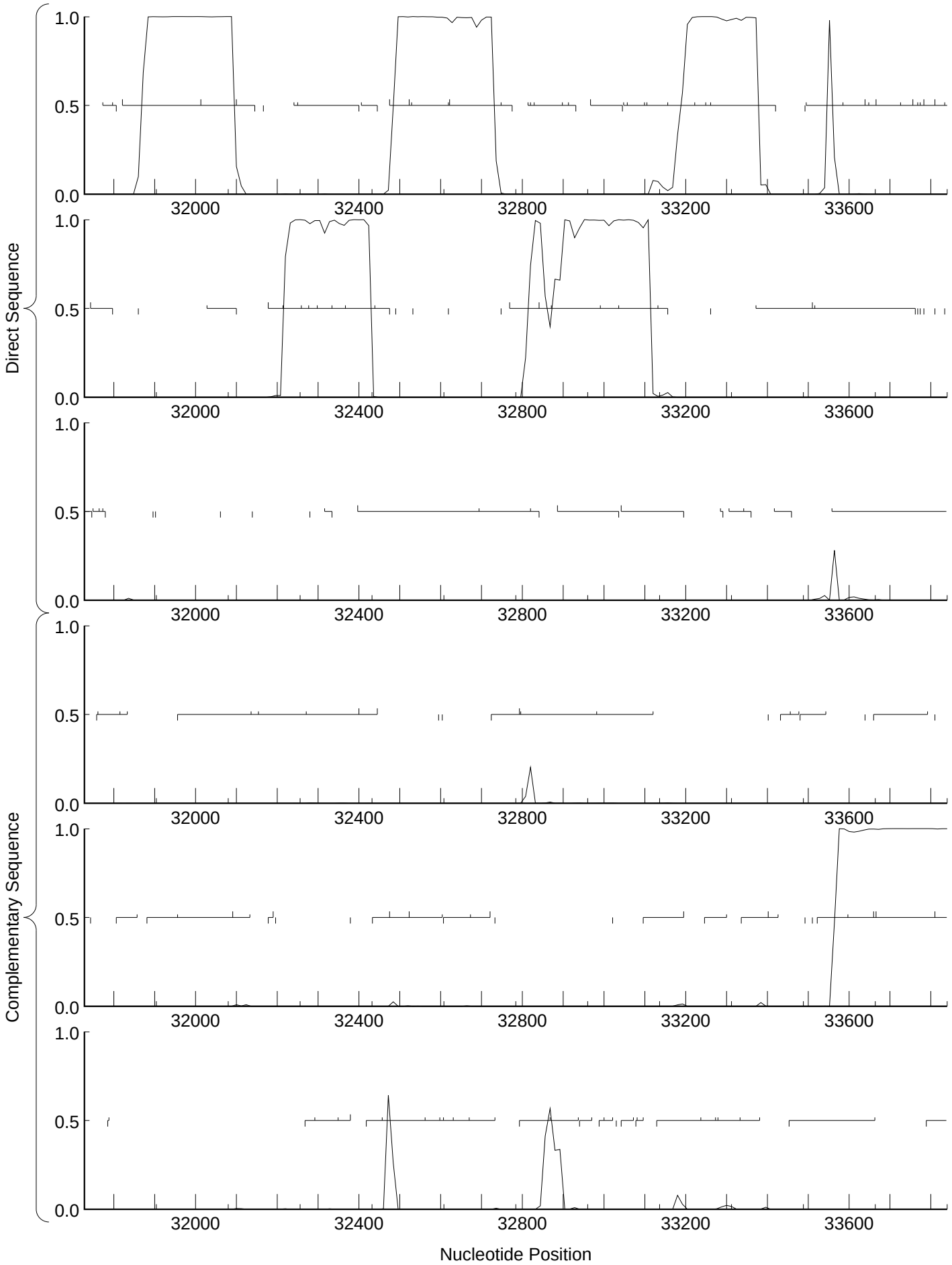

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
25600
26000
26400
26800
27200
Nucleotide Position

complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 15/29

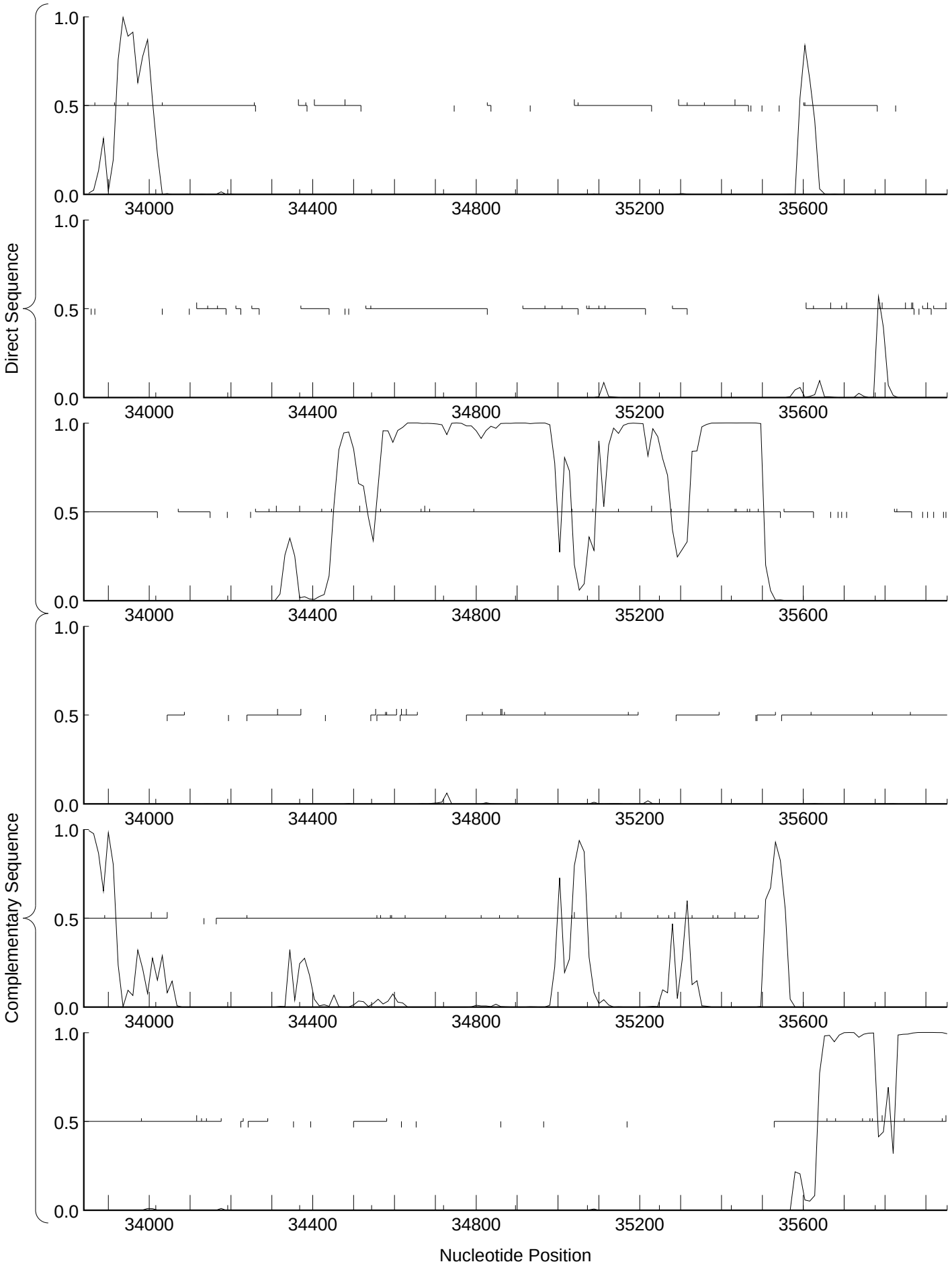


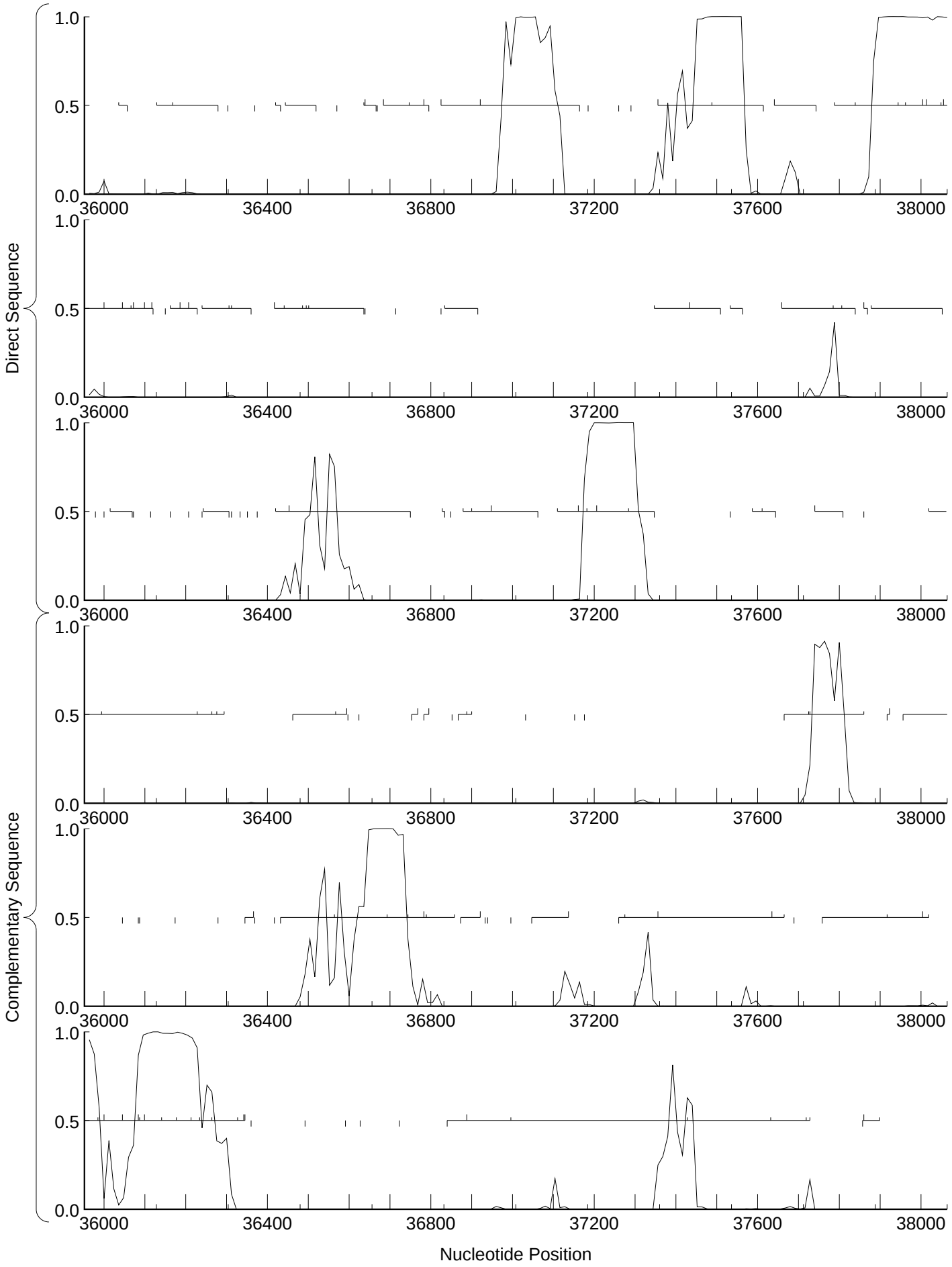
complete sequence, 58678 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 16/29

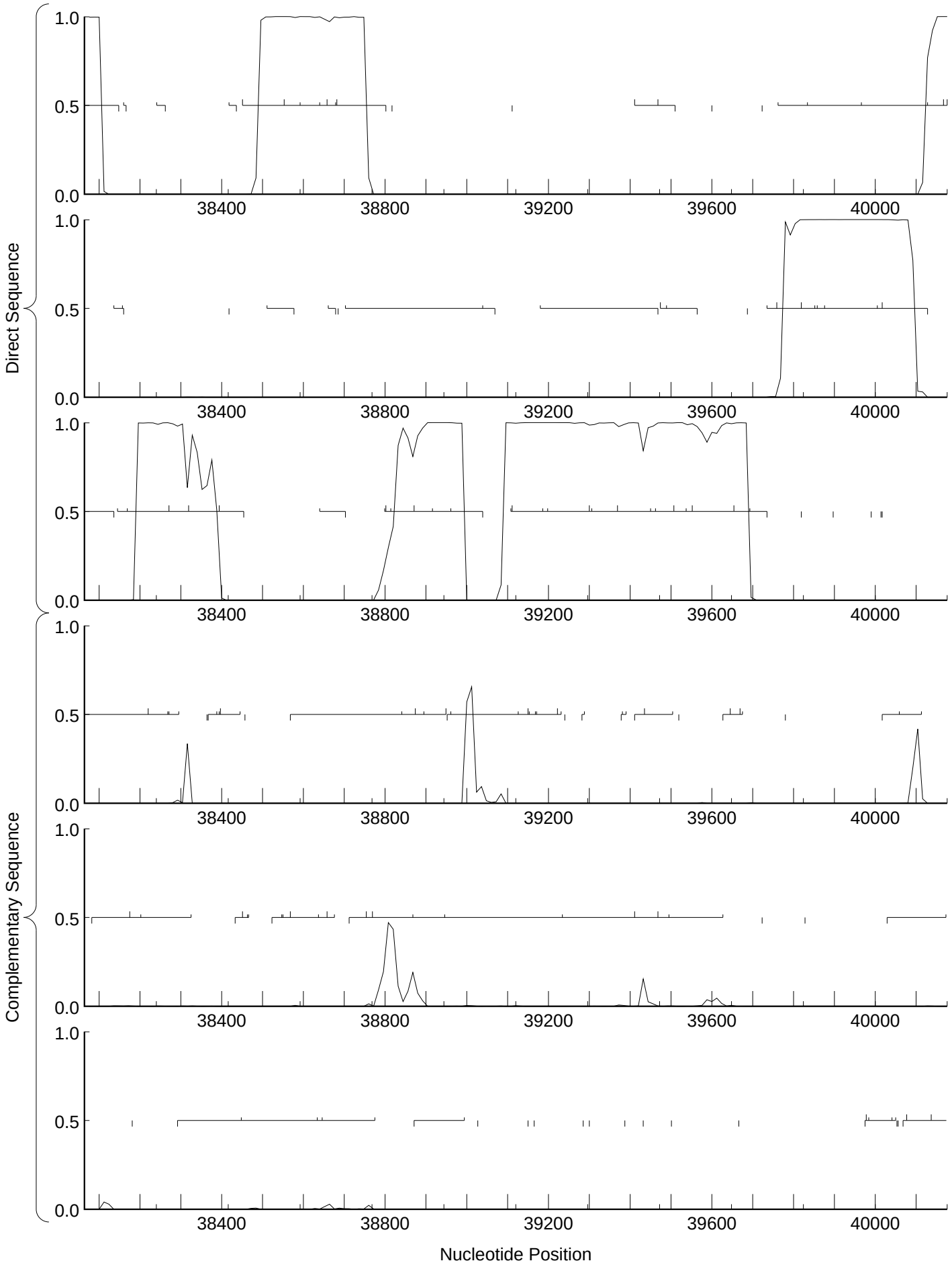


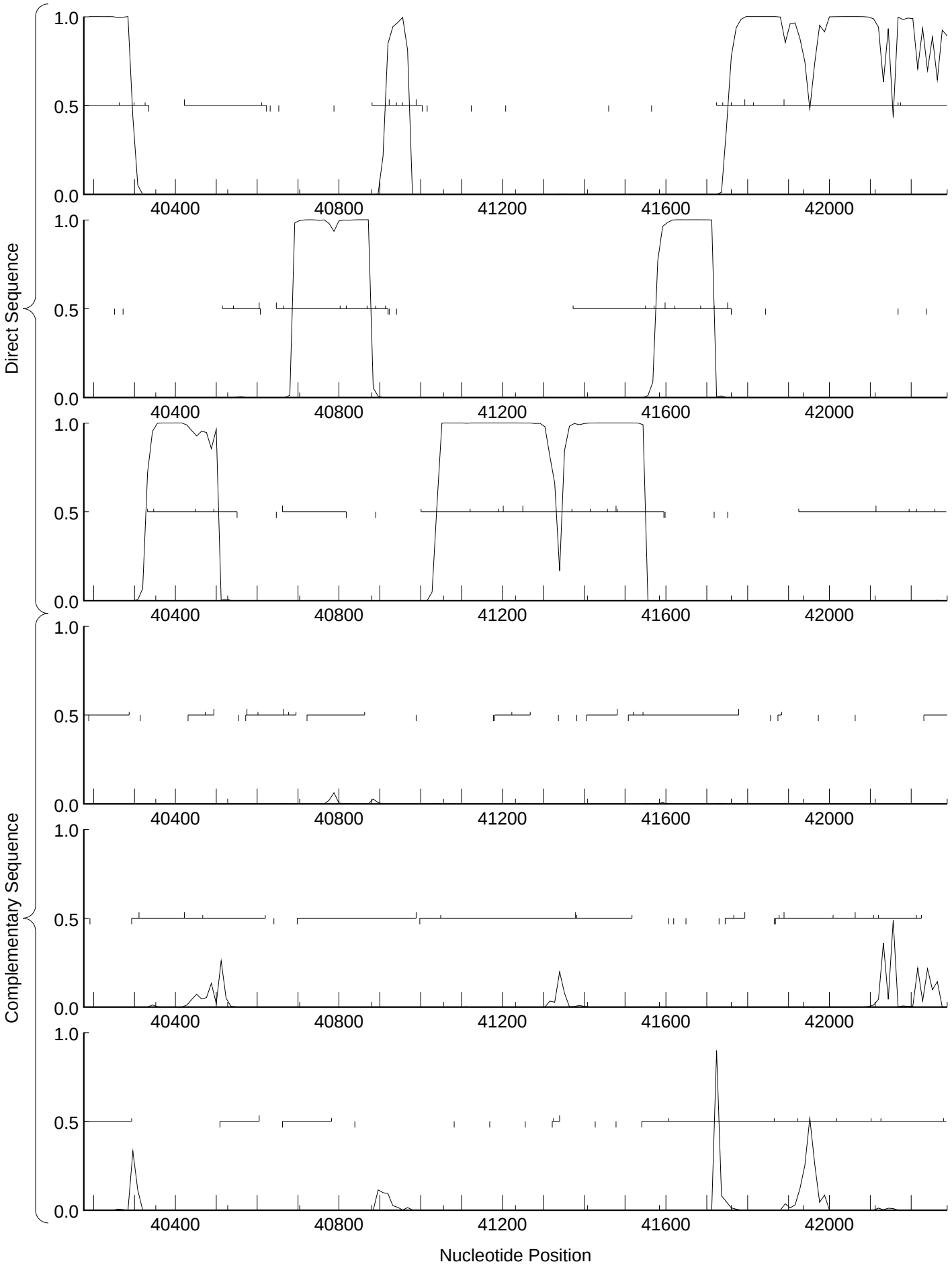

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
32000
32400
32800
33200
33600
Nucleotide Position

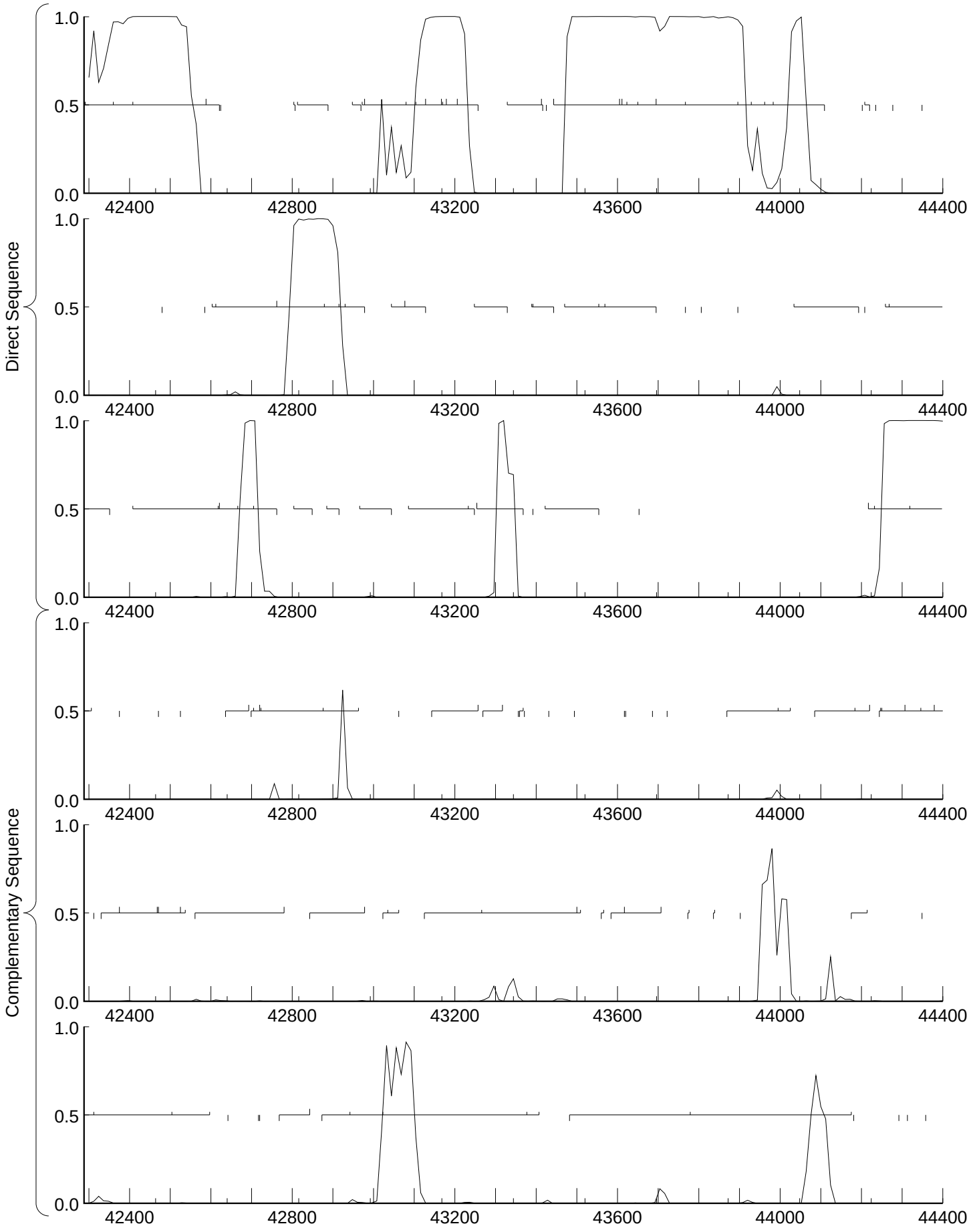
complete sequence, 58678 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 18/29



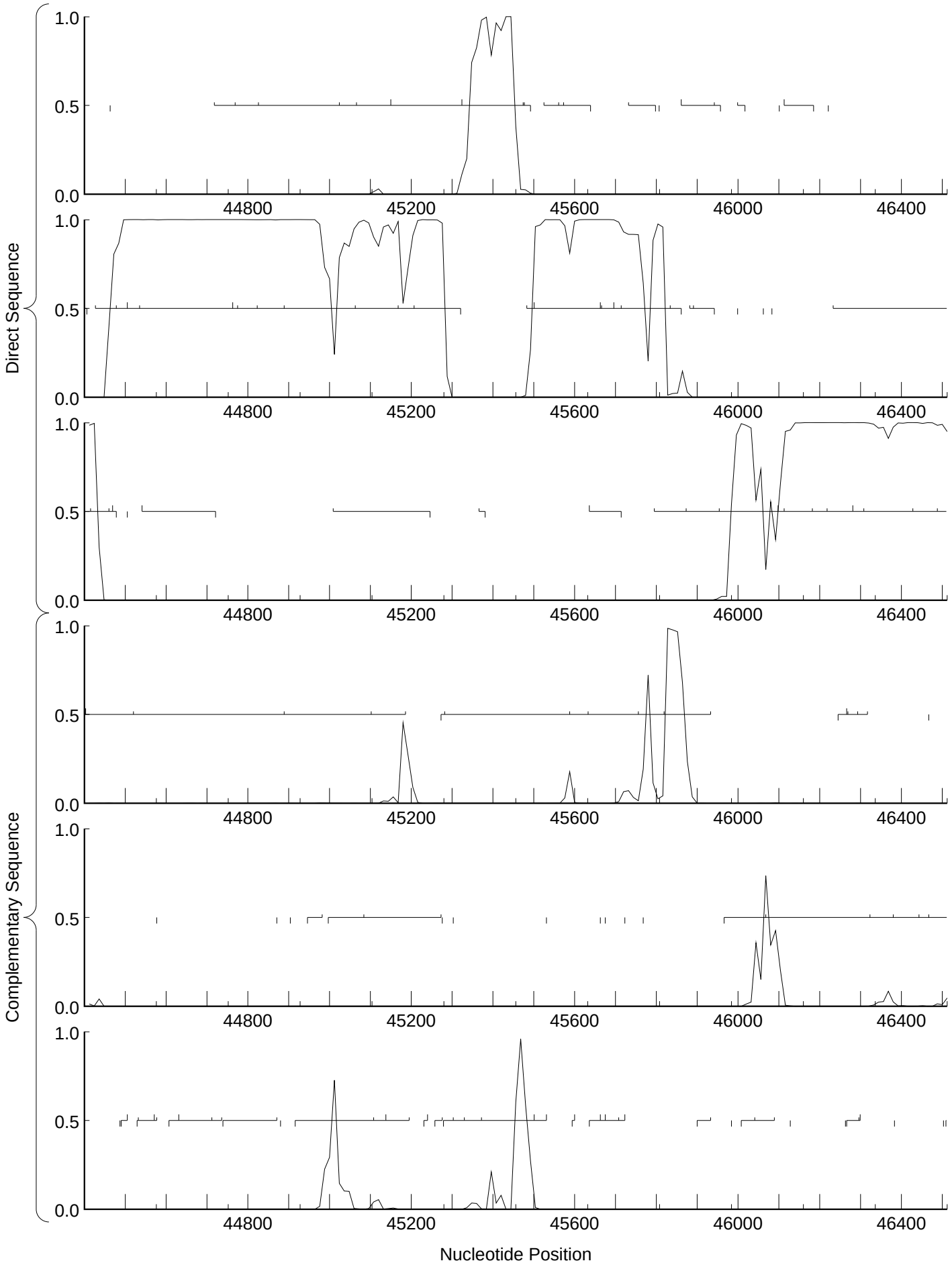

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
36000
36400
36800
37200
37600
38000
Nucleotide Position

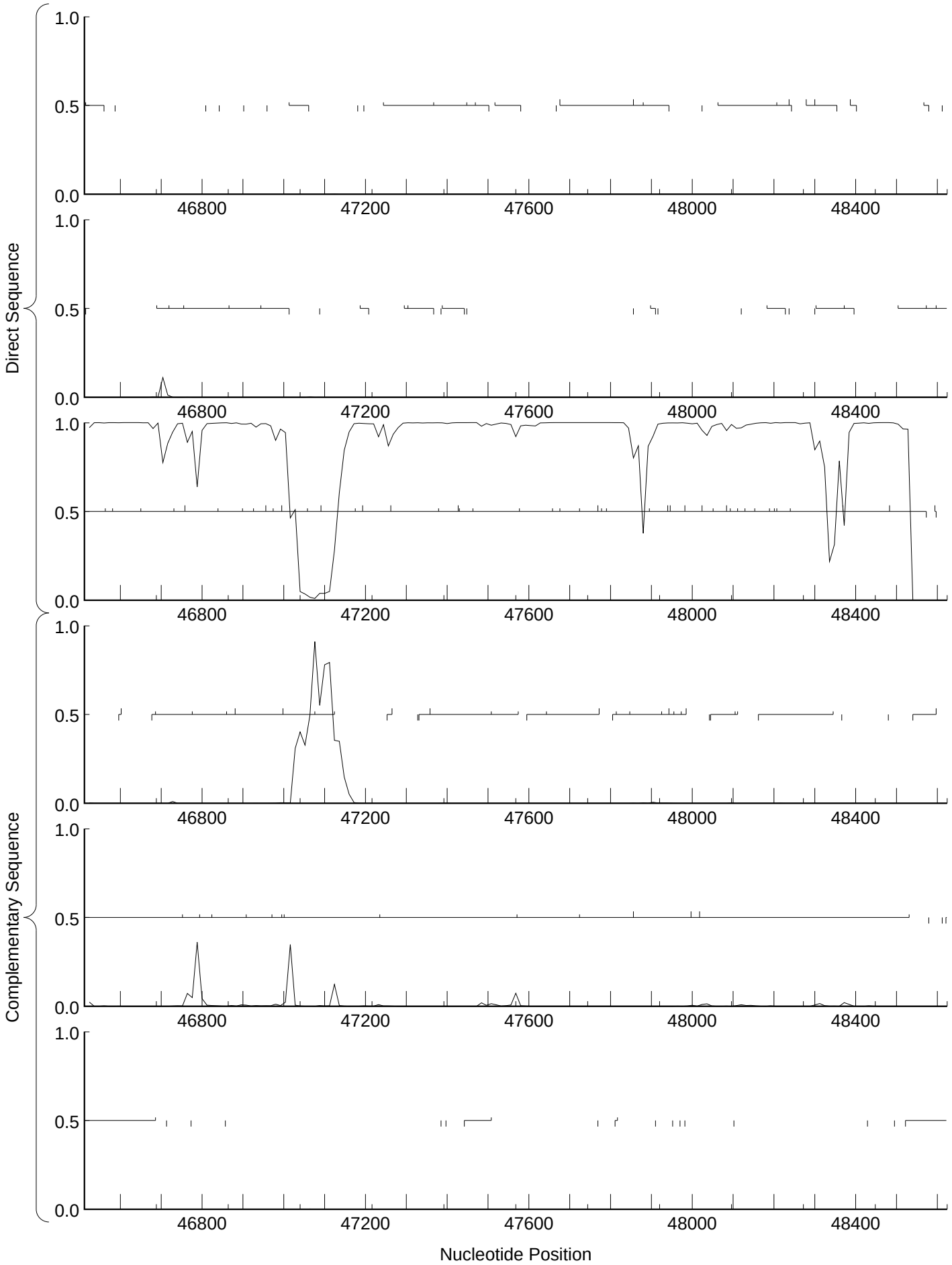

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

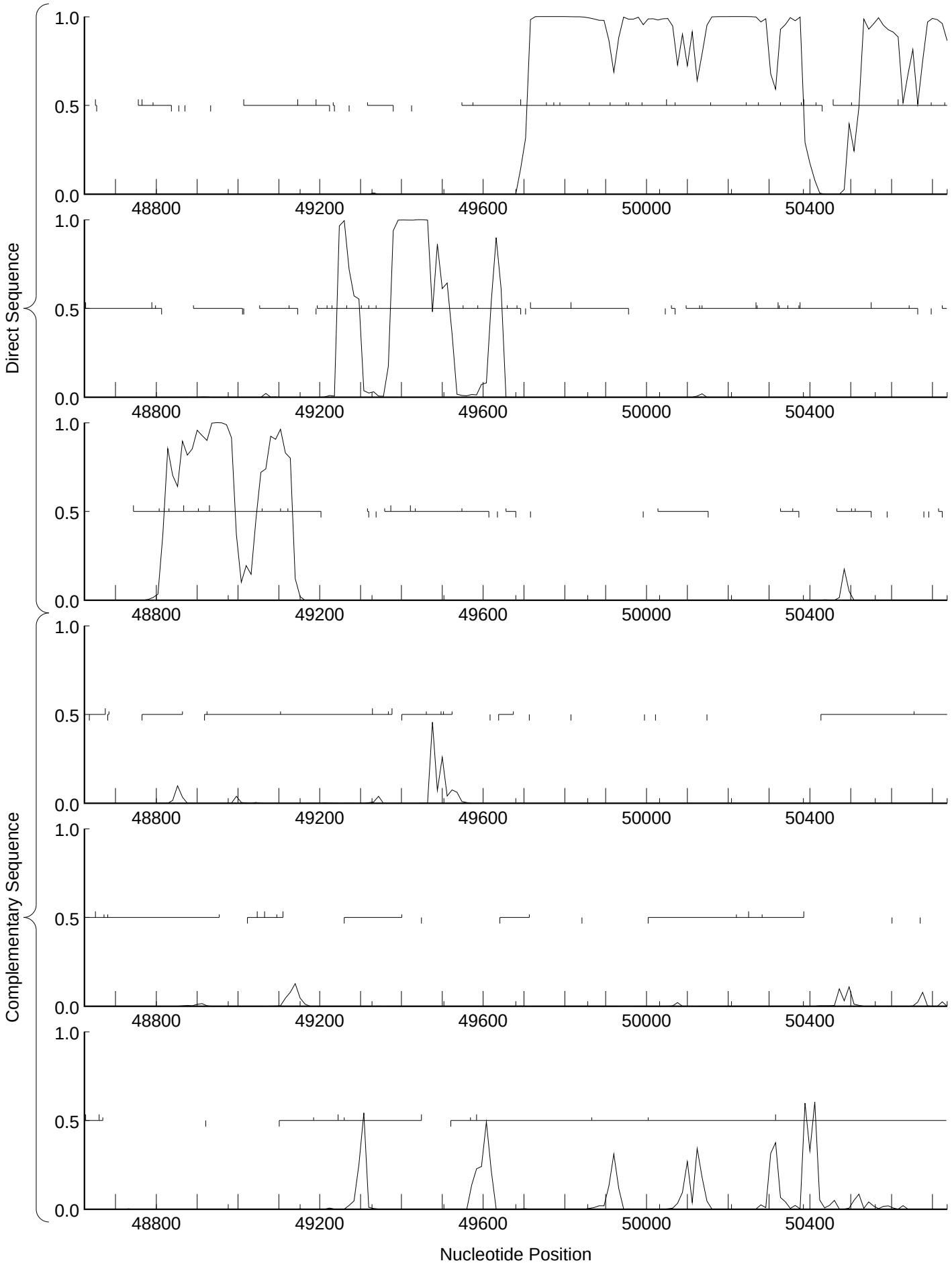

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
40400
40800
41200
41600
42000
Nucleotide Position

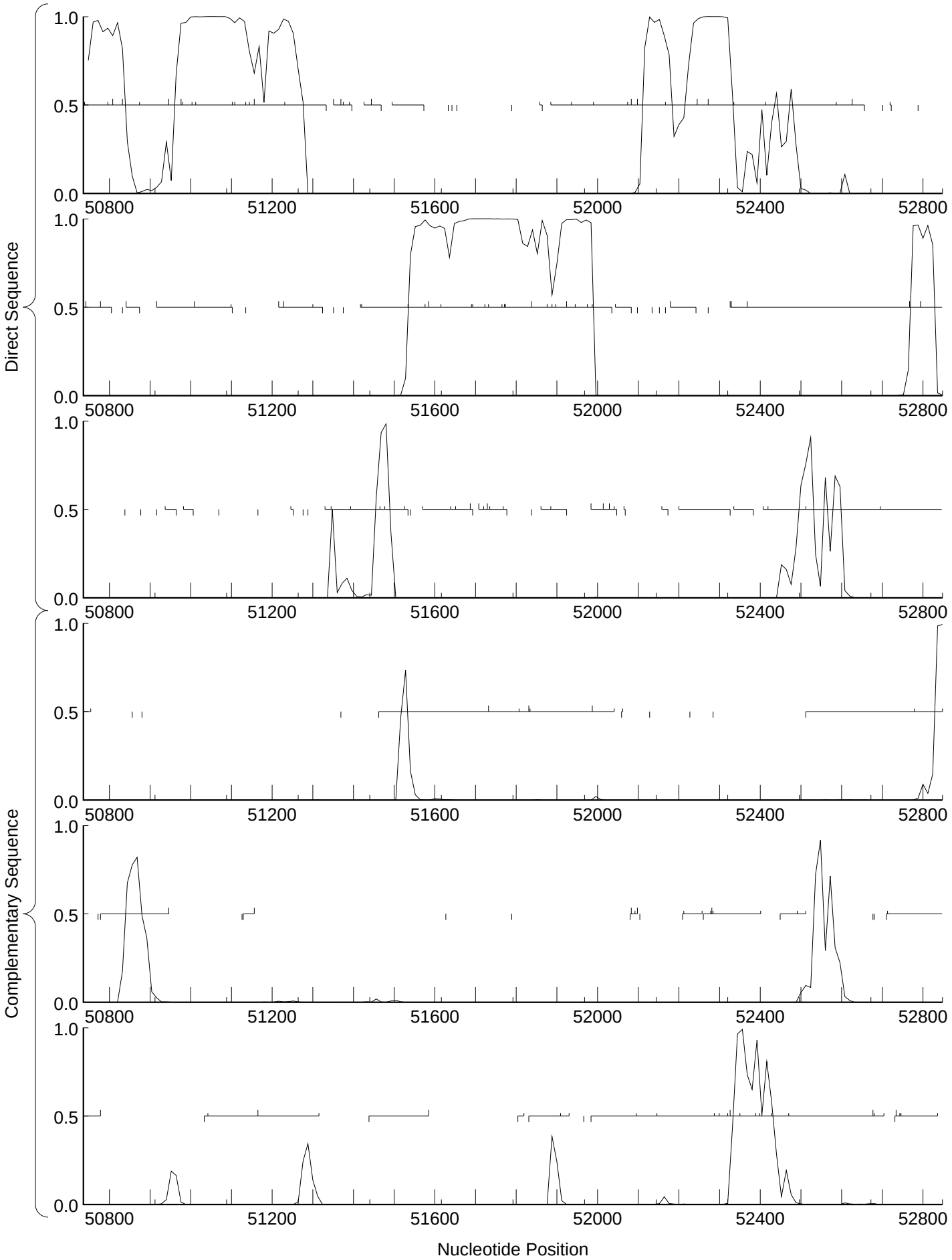

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
42400
42800
43200
43600
44000
44400
Nucleotide Position

complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 23/29

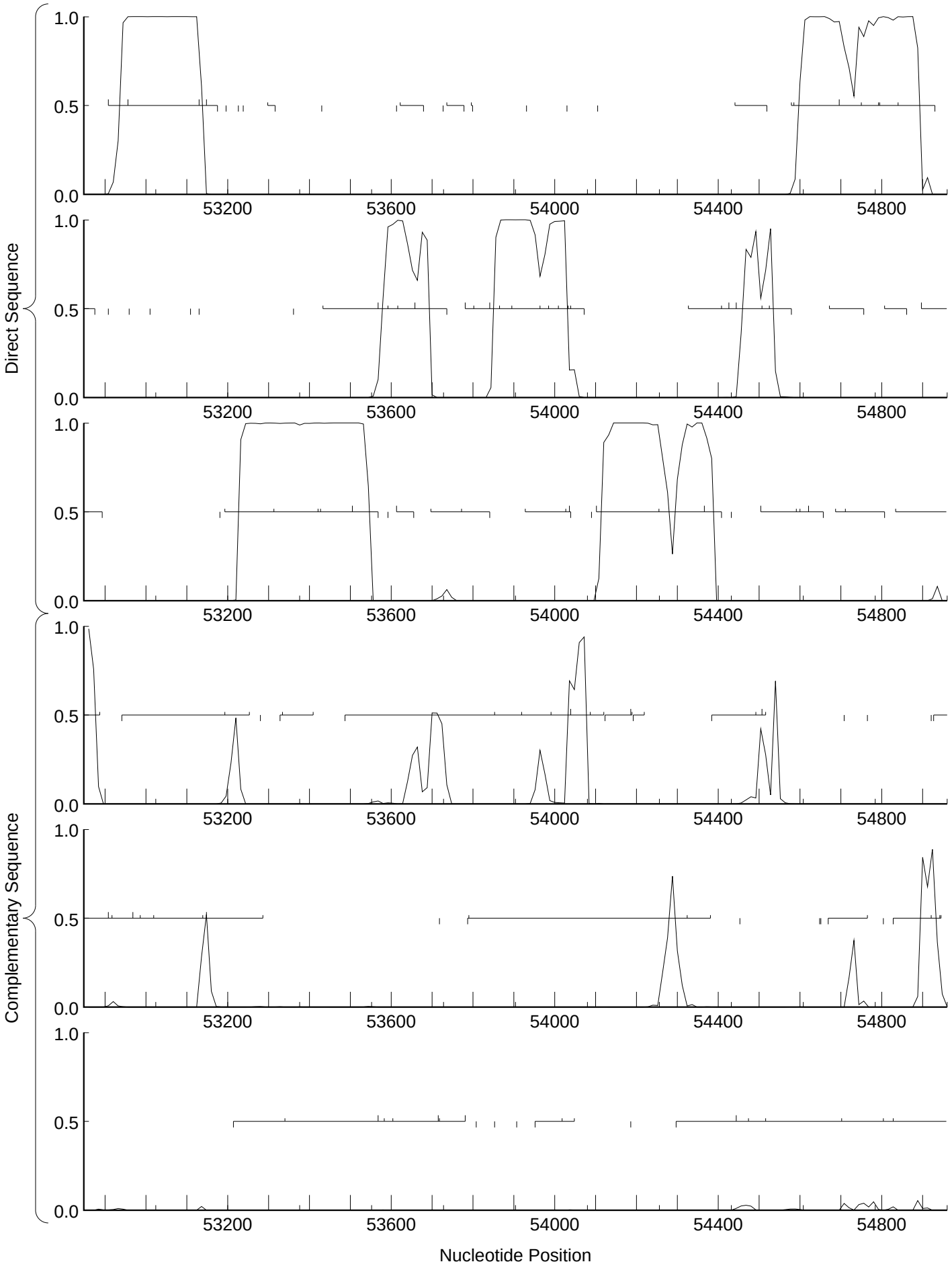



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
46800
47200
47600
48000
48400
Nucleotide Position

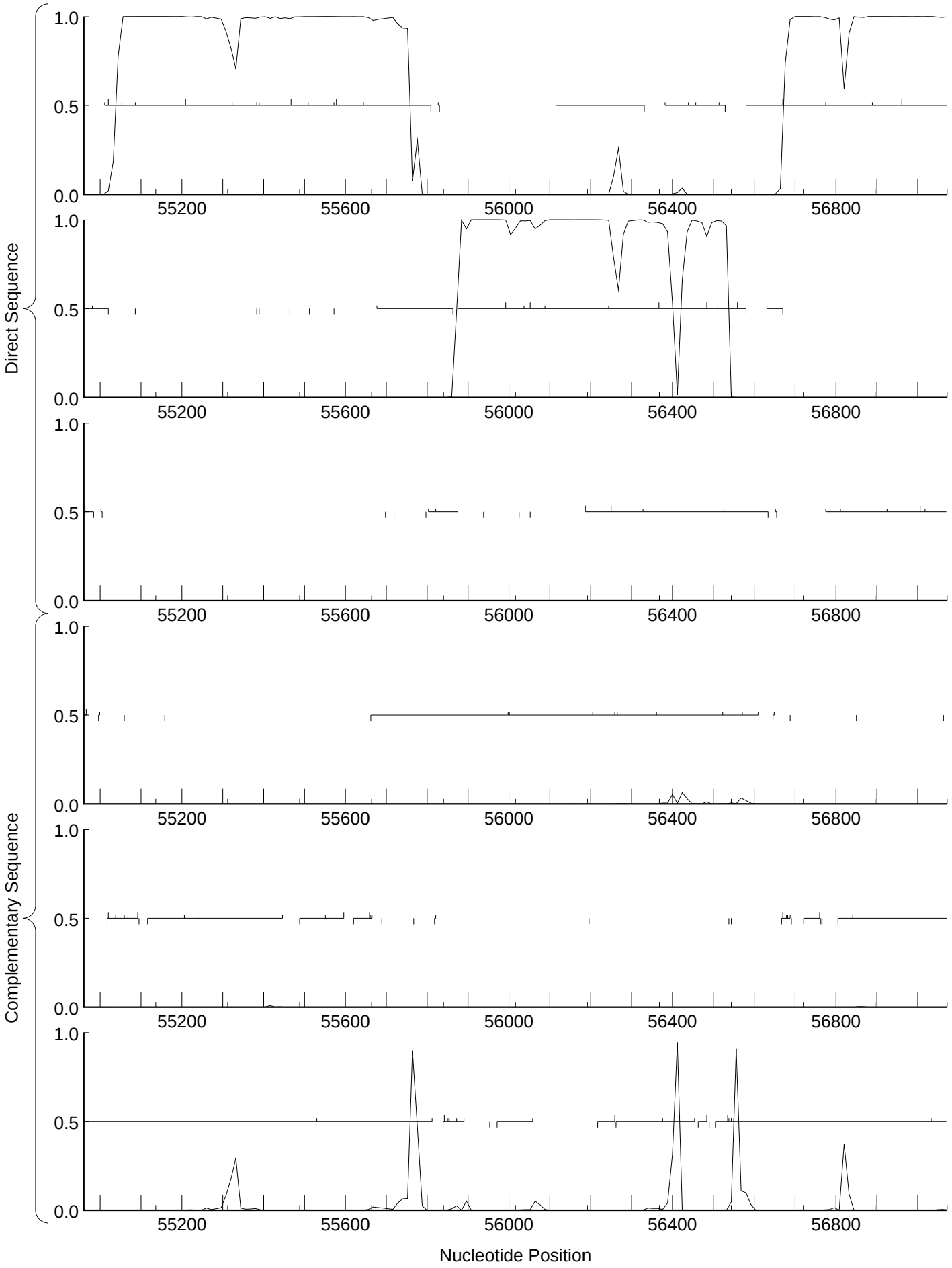

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
48800
49200
49600
50000
50400
Nucleotide Position


Direct Sequence
Complementary Sequence
1.0
0.5
0.0
50800
51200
51600
52000
52400
52800
Nucleotide Position

complete sequence, 58678 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 27/29



complete sequence, 58678 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 28/29



complete sequence, 58678 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 29/29

