GeneMark

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

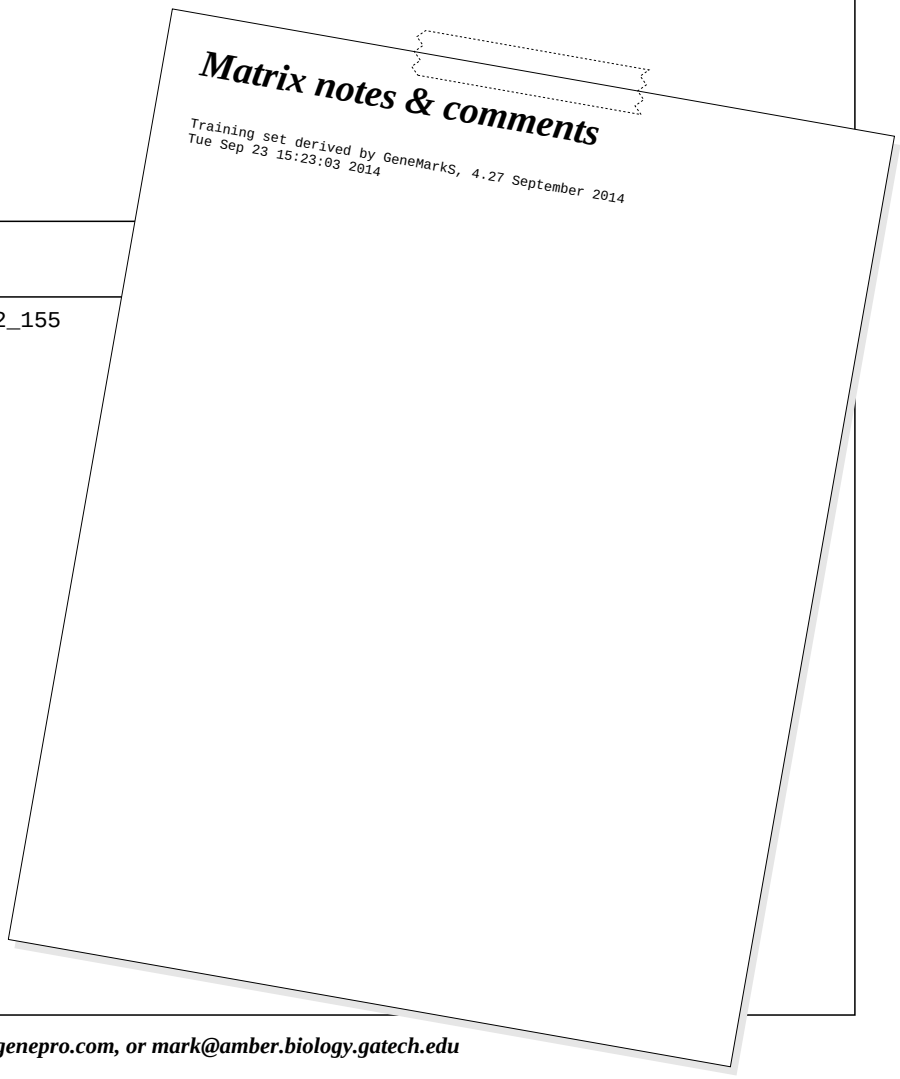
PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Slarp complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCG
Analysis Date    :  3/28/24 at 21:16:30
Pages            :  29
Sequence Length  :  57256 bp
GC Content       :   67.96%

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

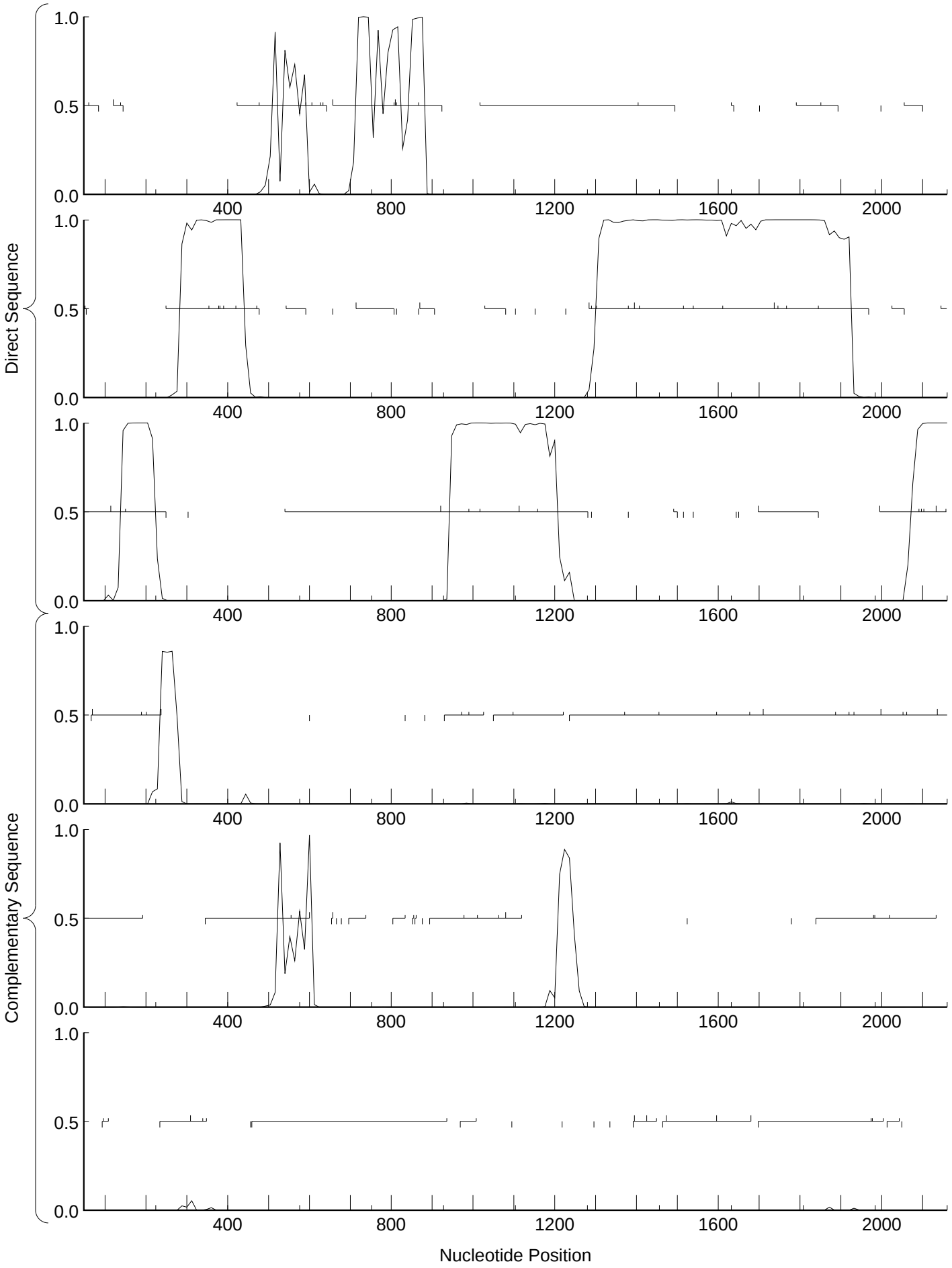
```
Matrix     :  Mycobacterium_smegmatis_MC2_155
Author     :  -

Order      :  4
```

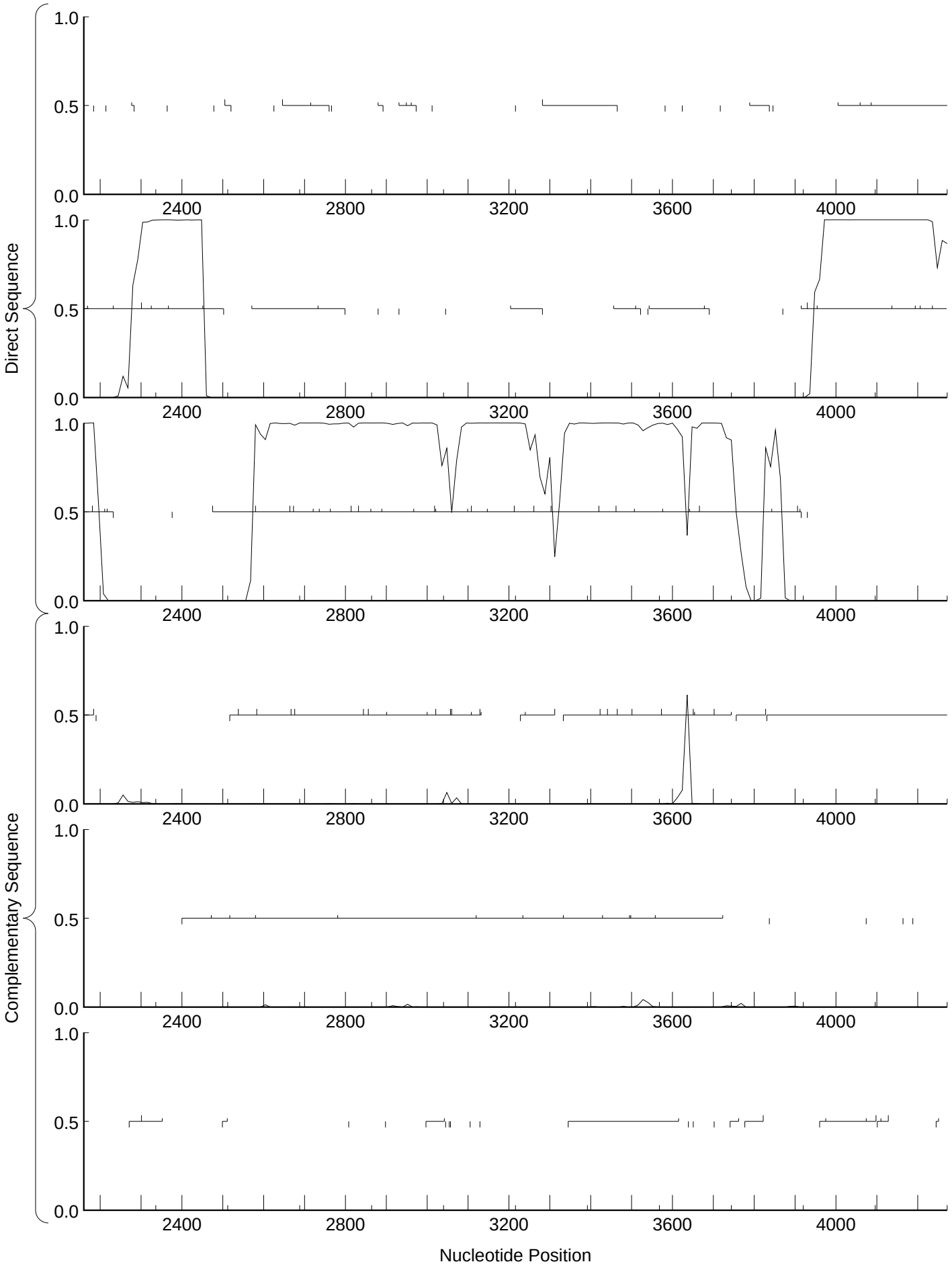

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


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

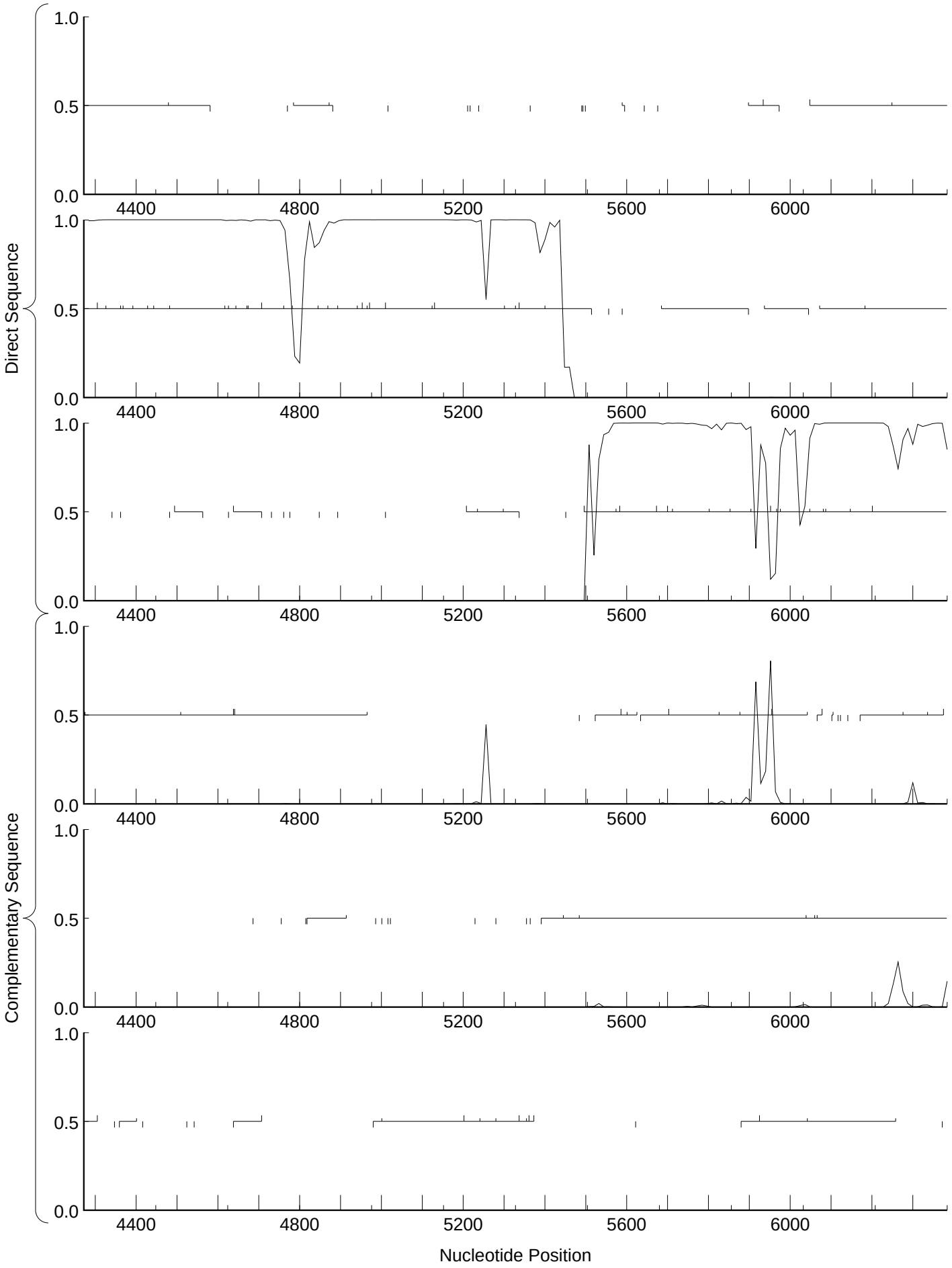
complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 2/29



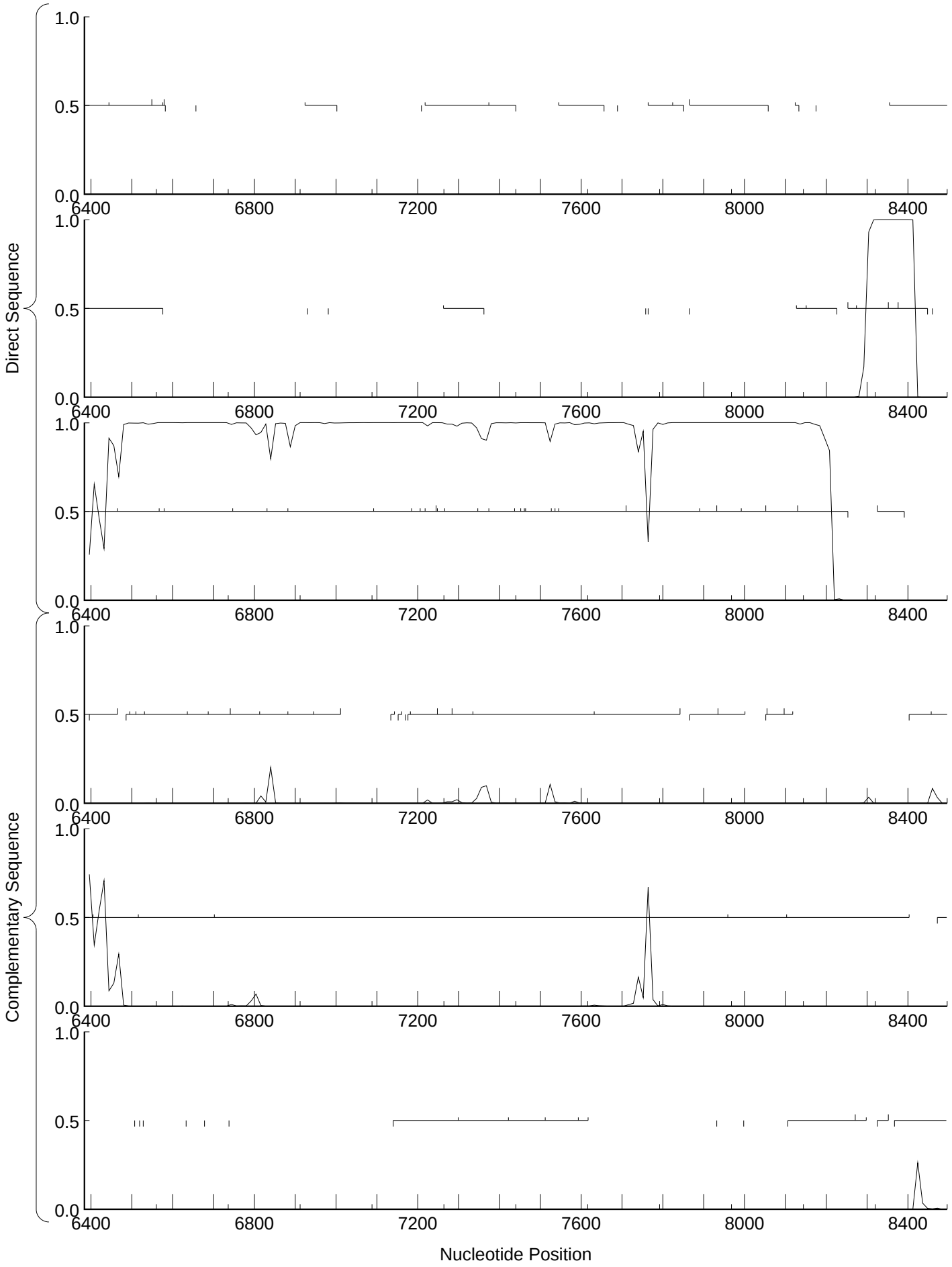
complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 3/29



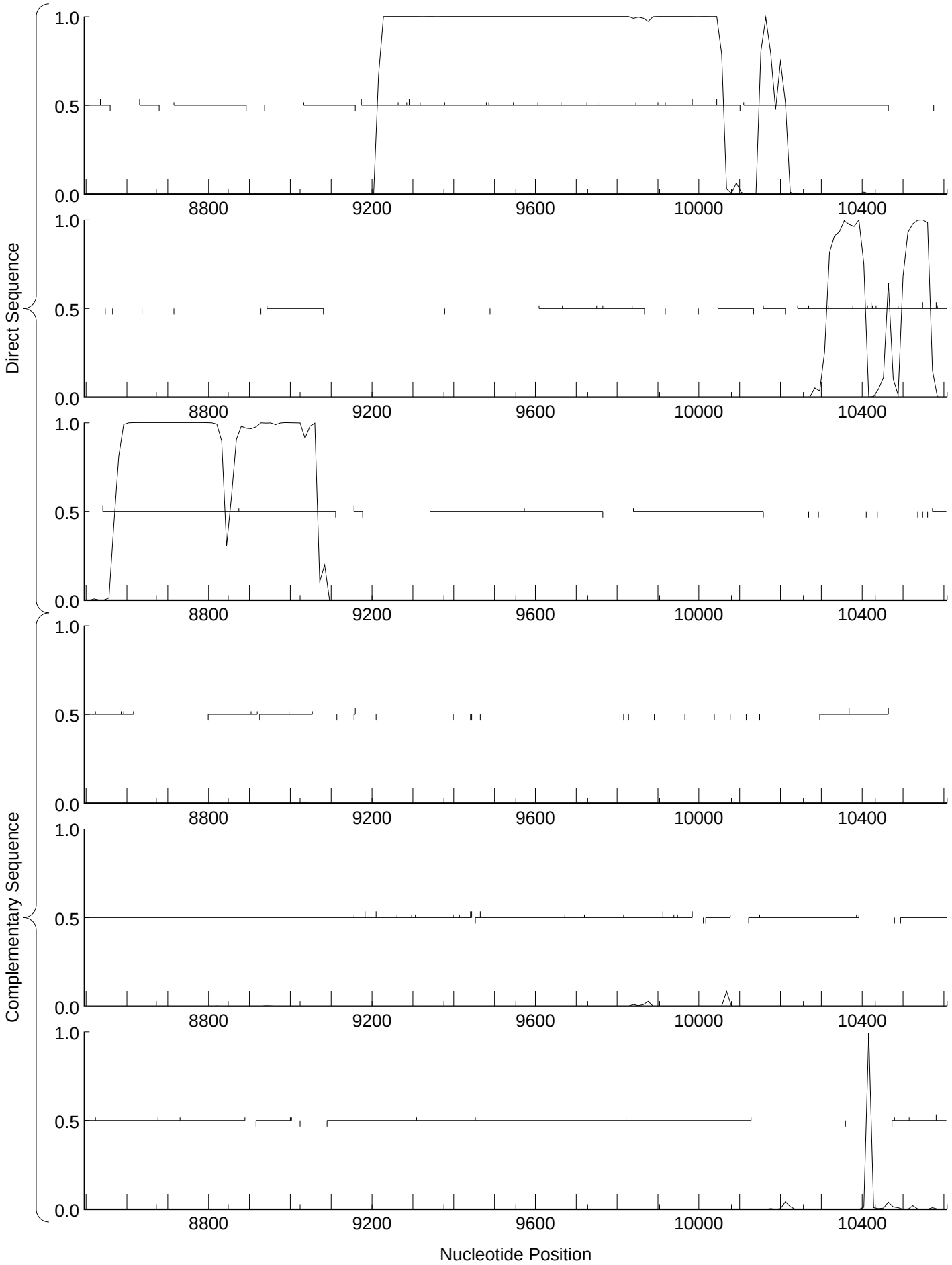
complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 4/29



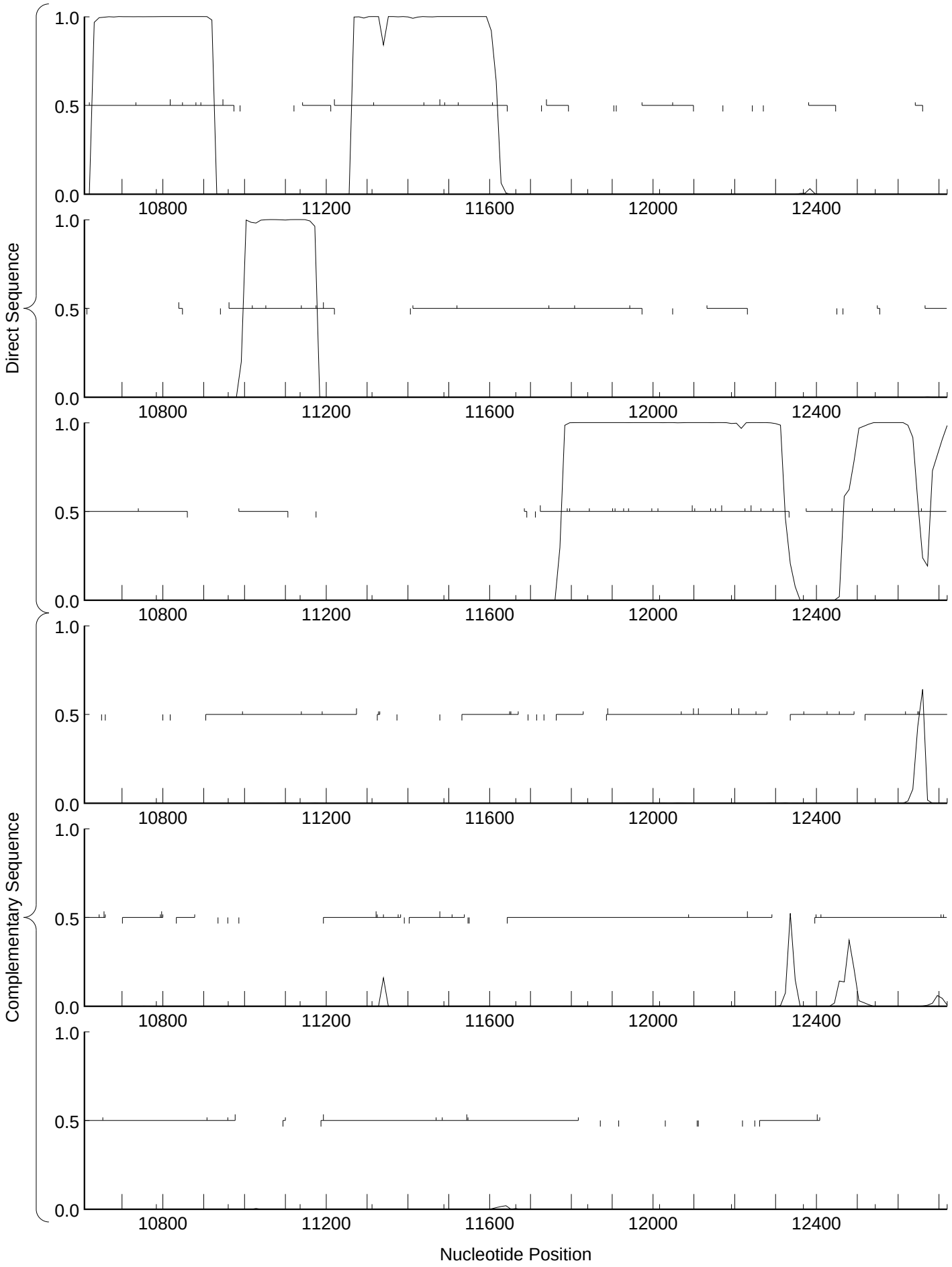
complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 5/29



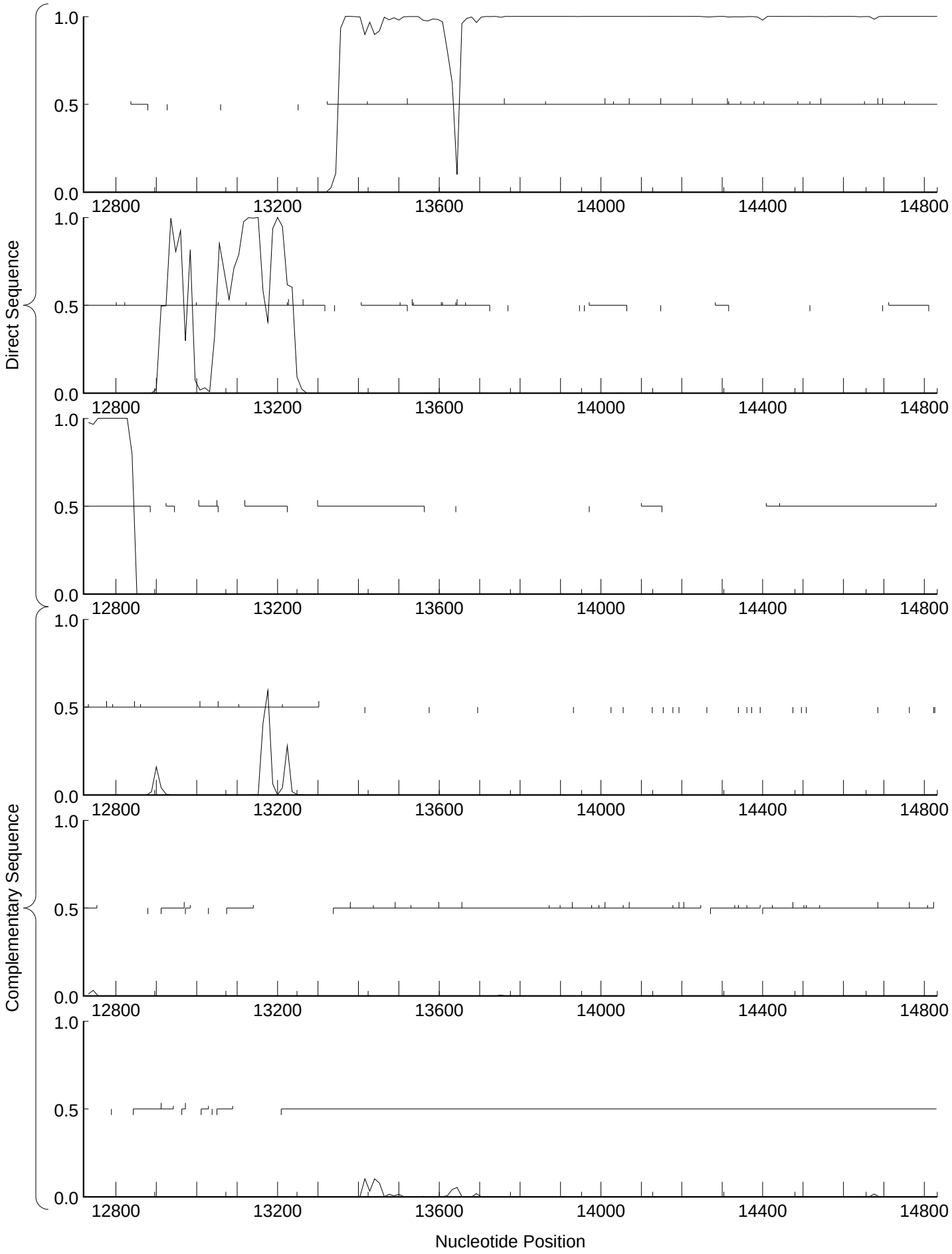
complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 6/29



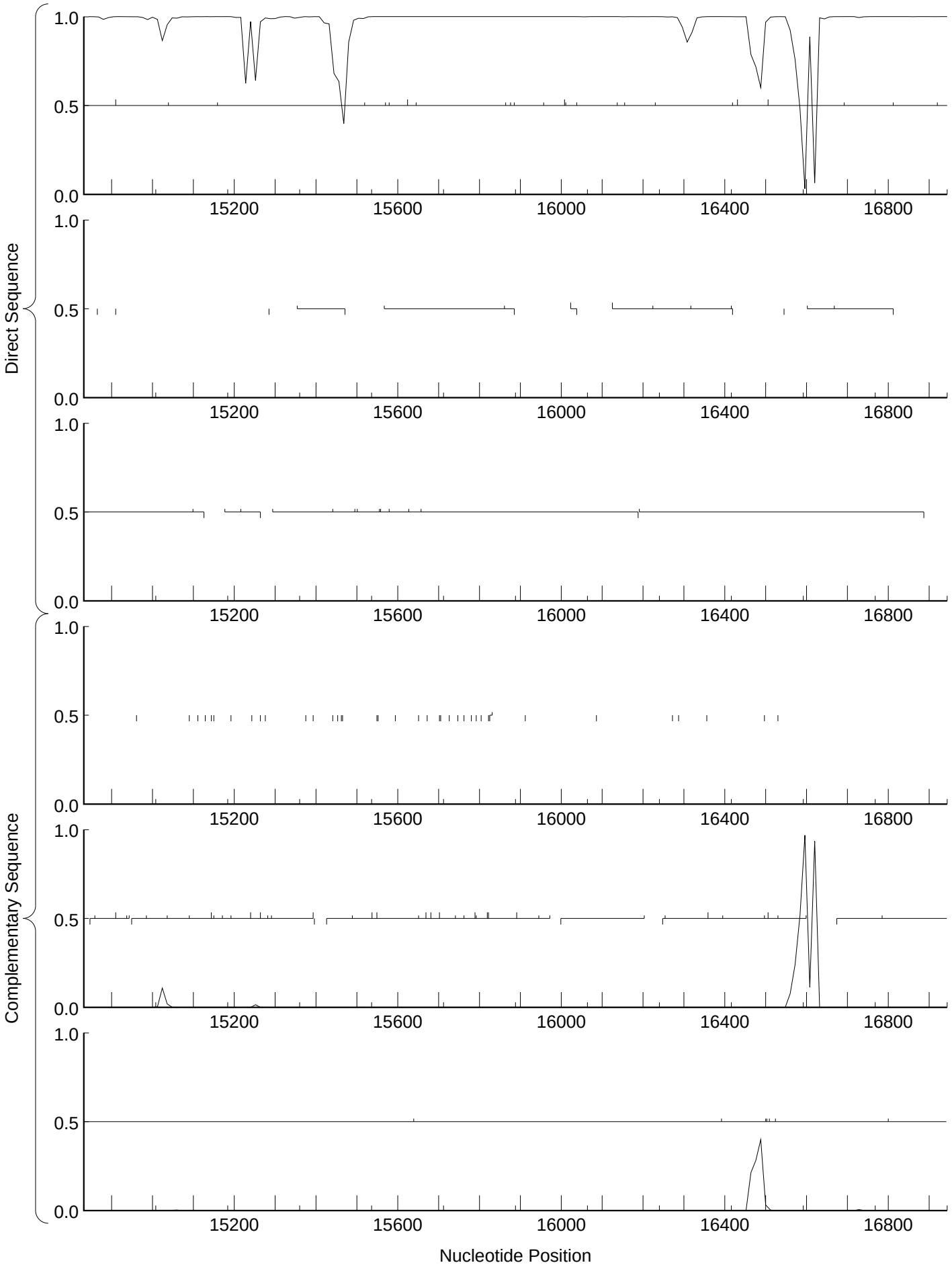
complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 7/29



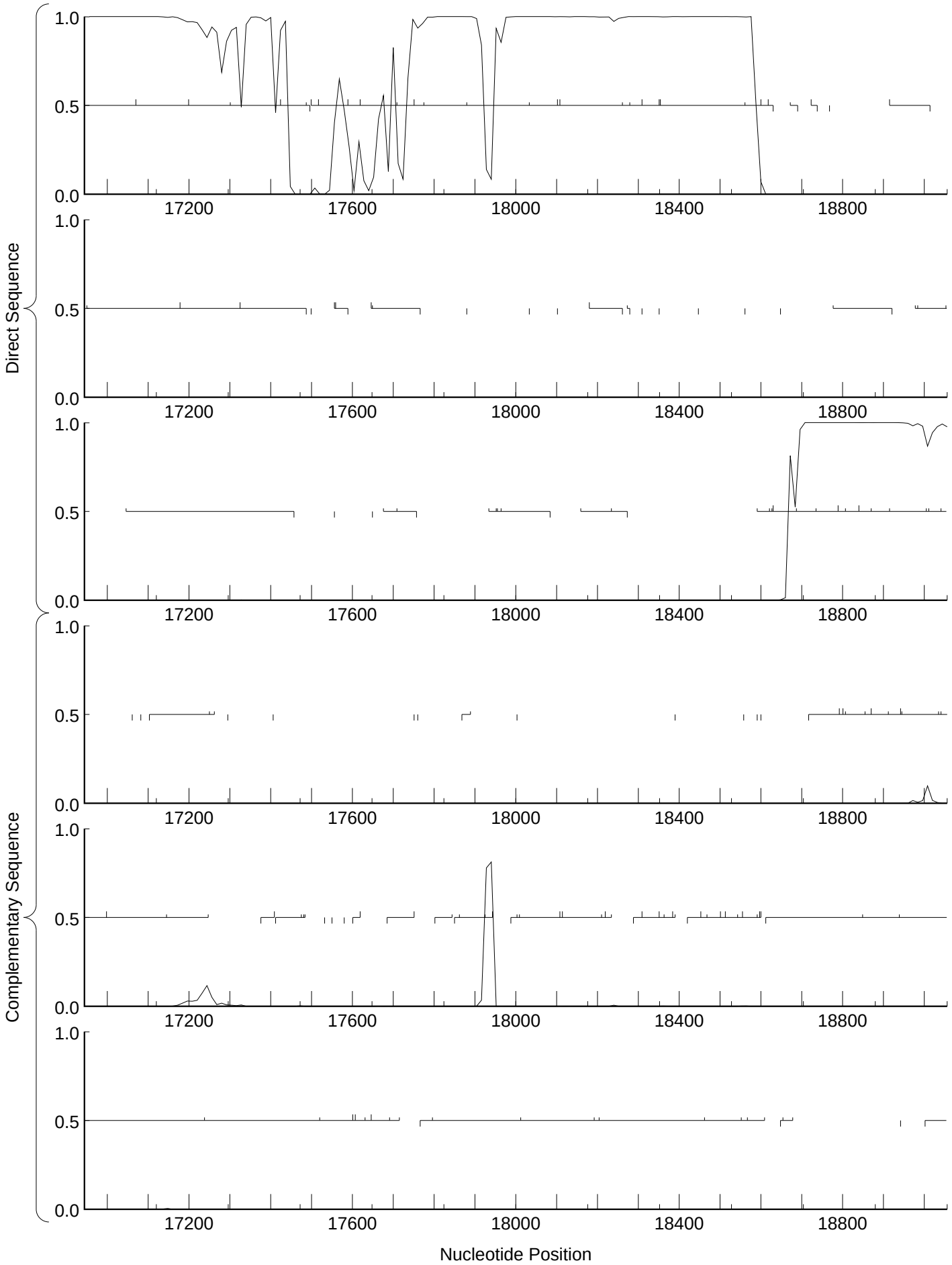
complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 8/29



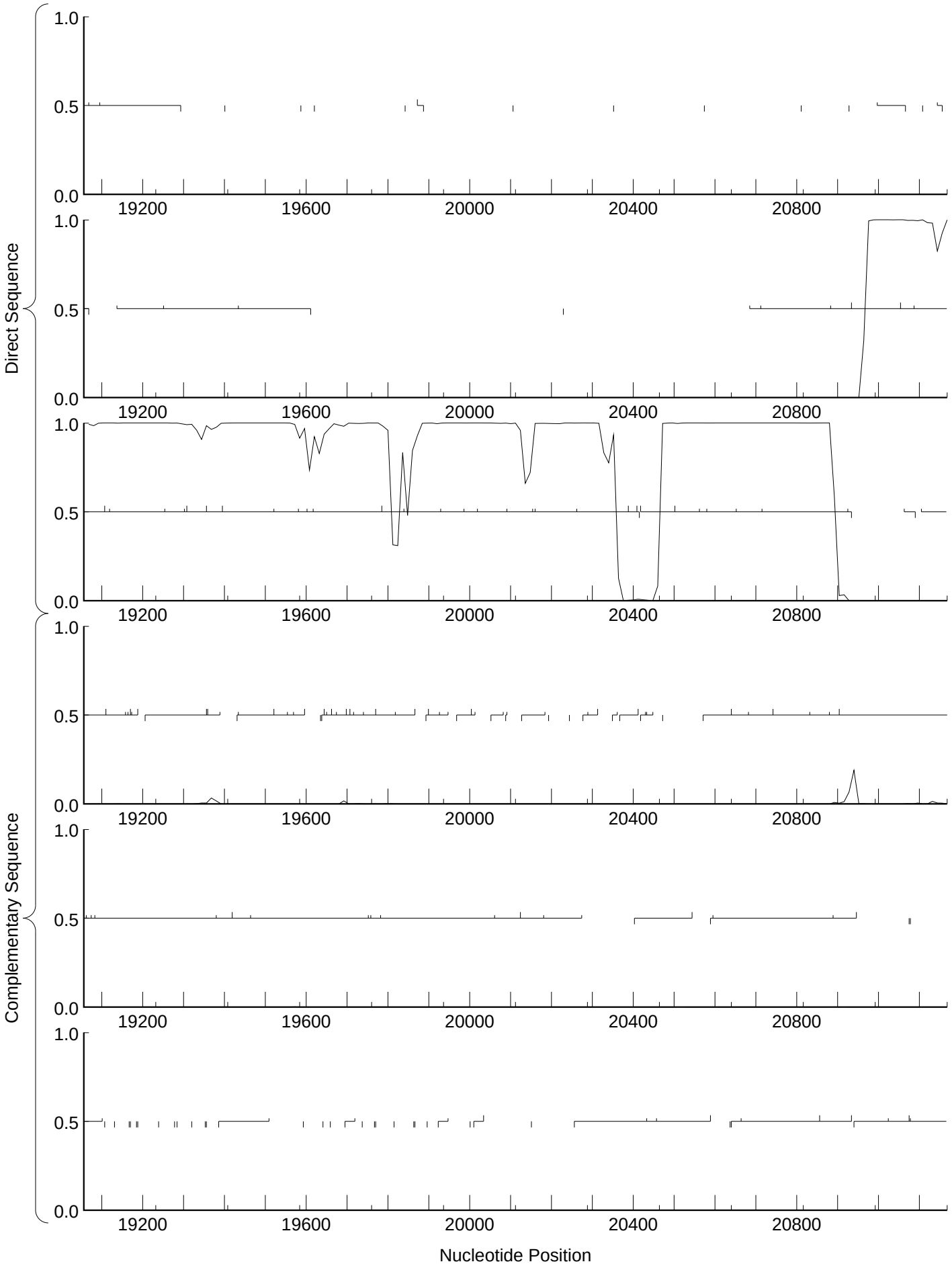
complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 9/29

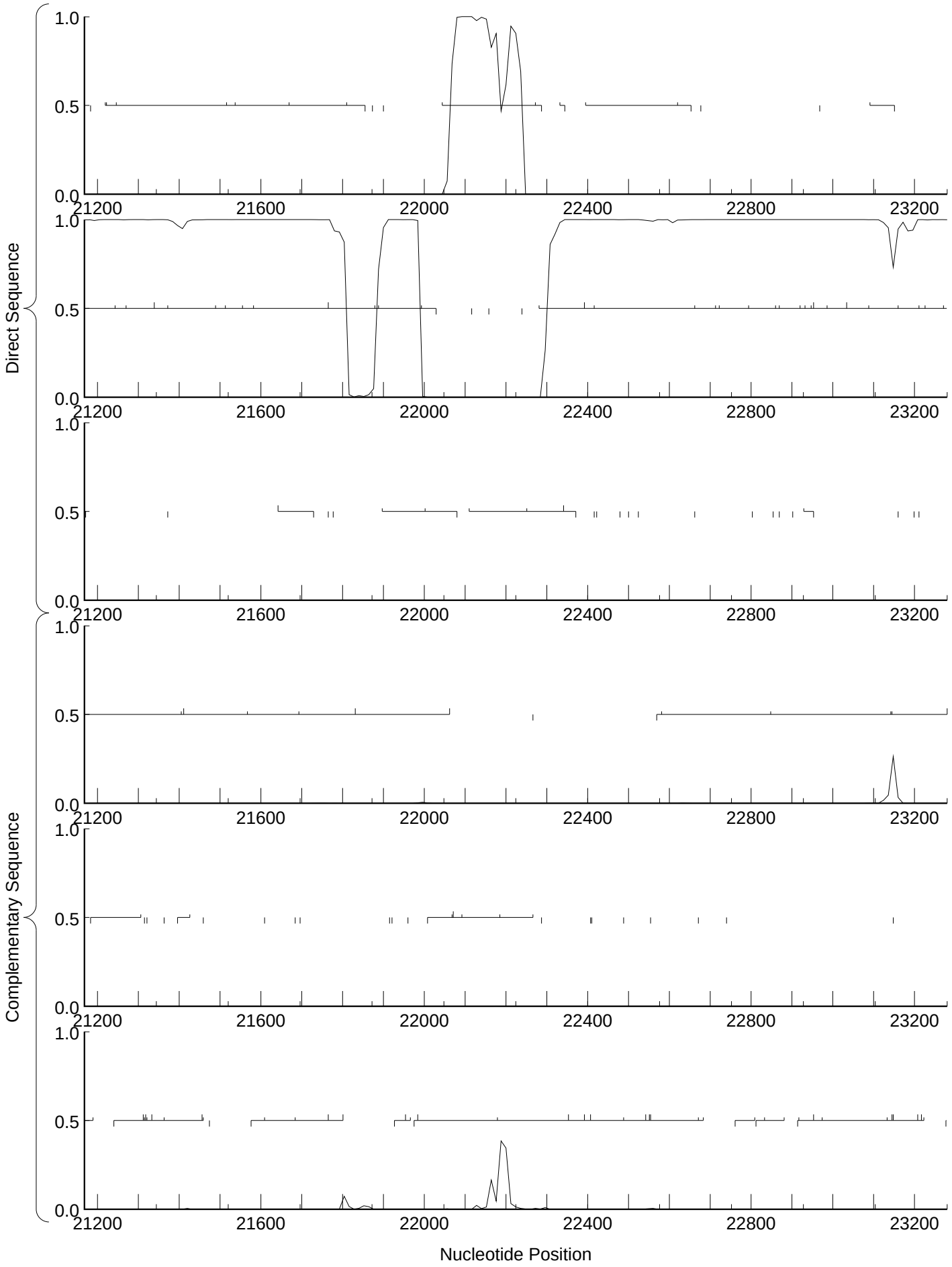


complete sequence, 57256 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 10/29

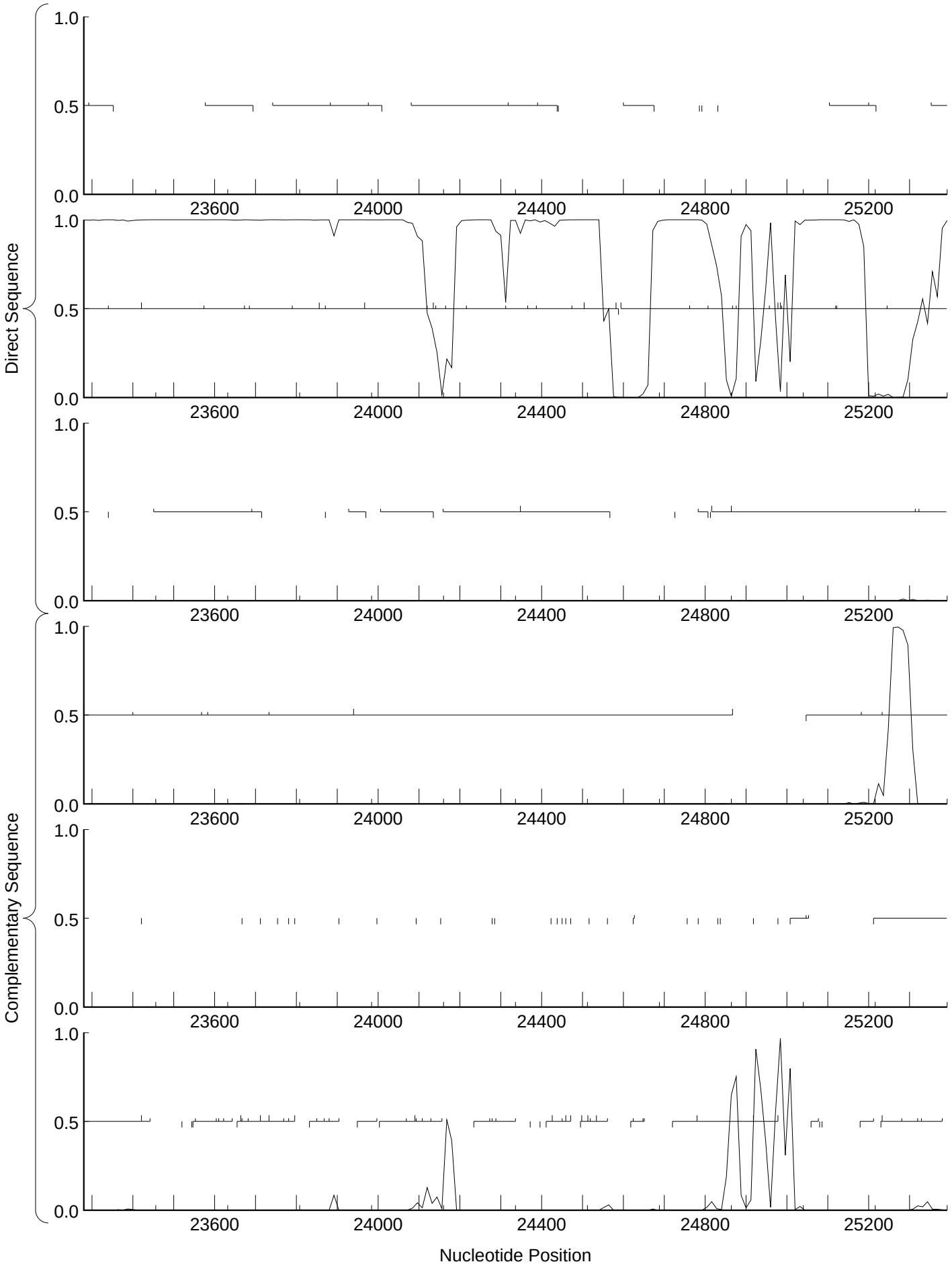


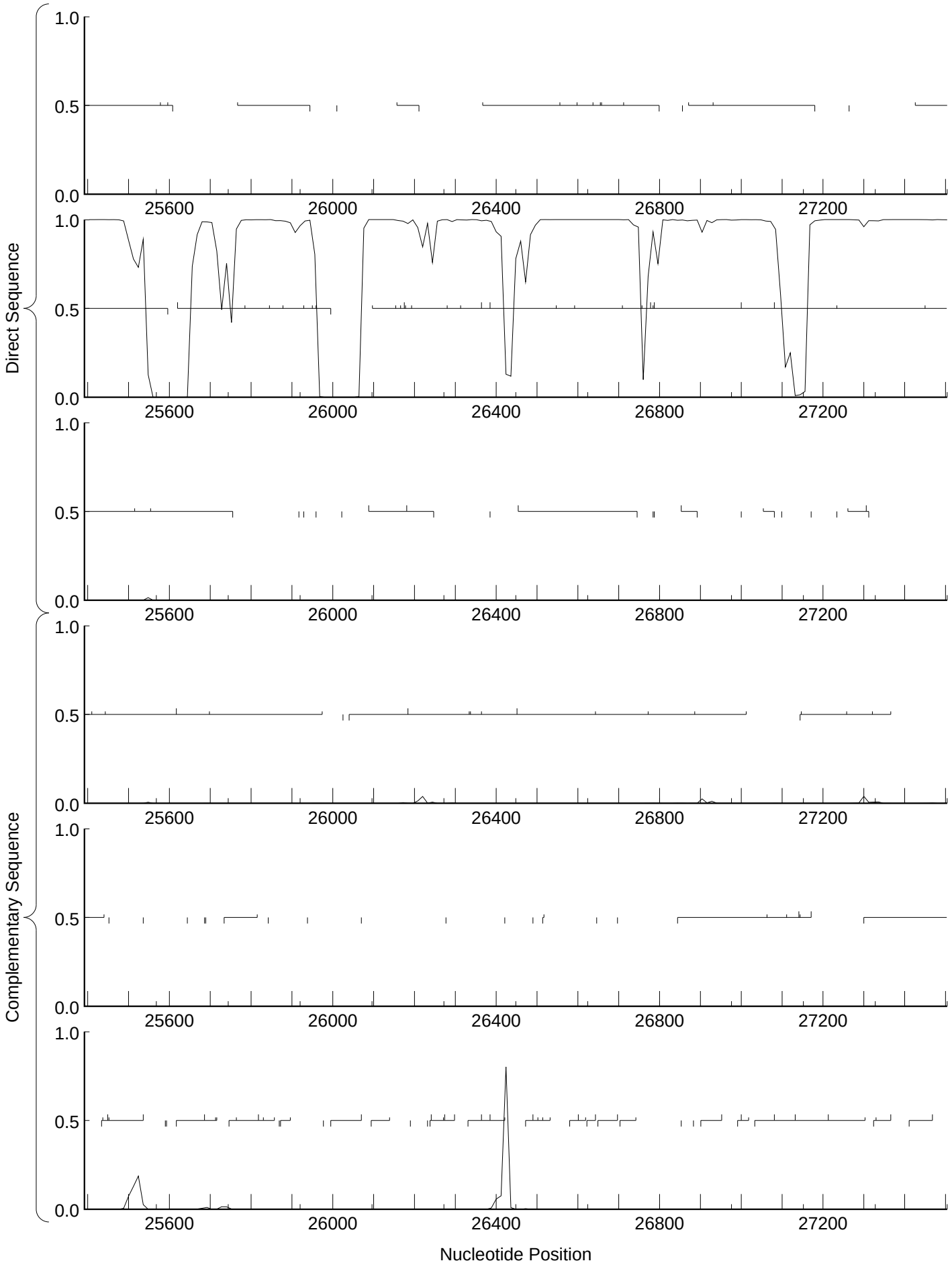
complete sequence, 57256 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 11/29



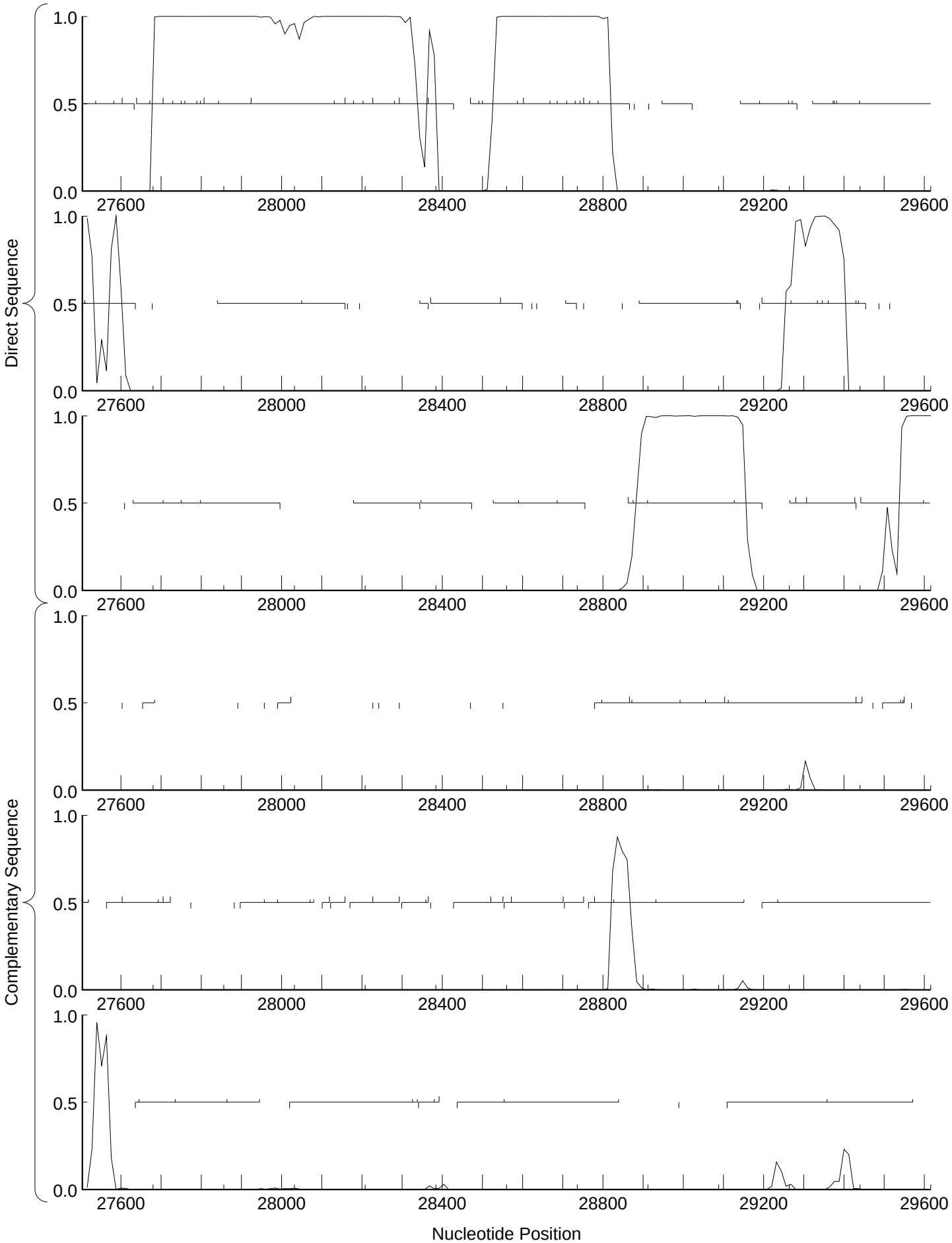

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
21200
21600
22000
22400
22800
23200
Nucleotide Position

complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 13/29

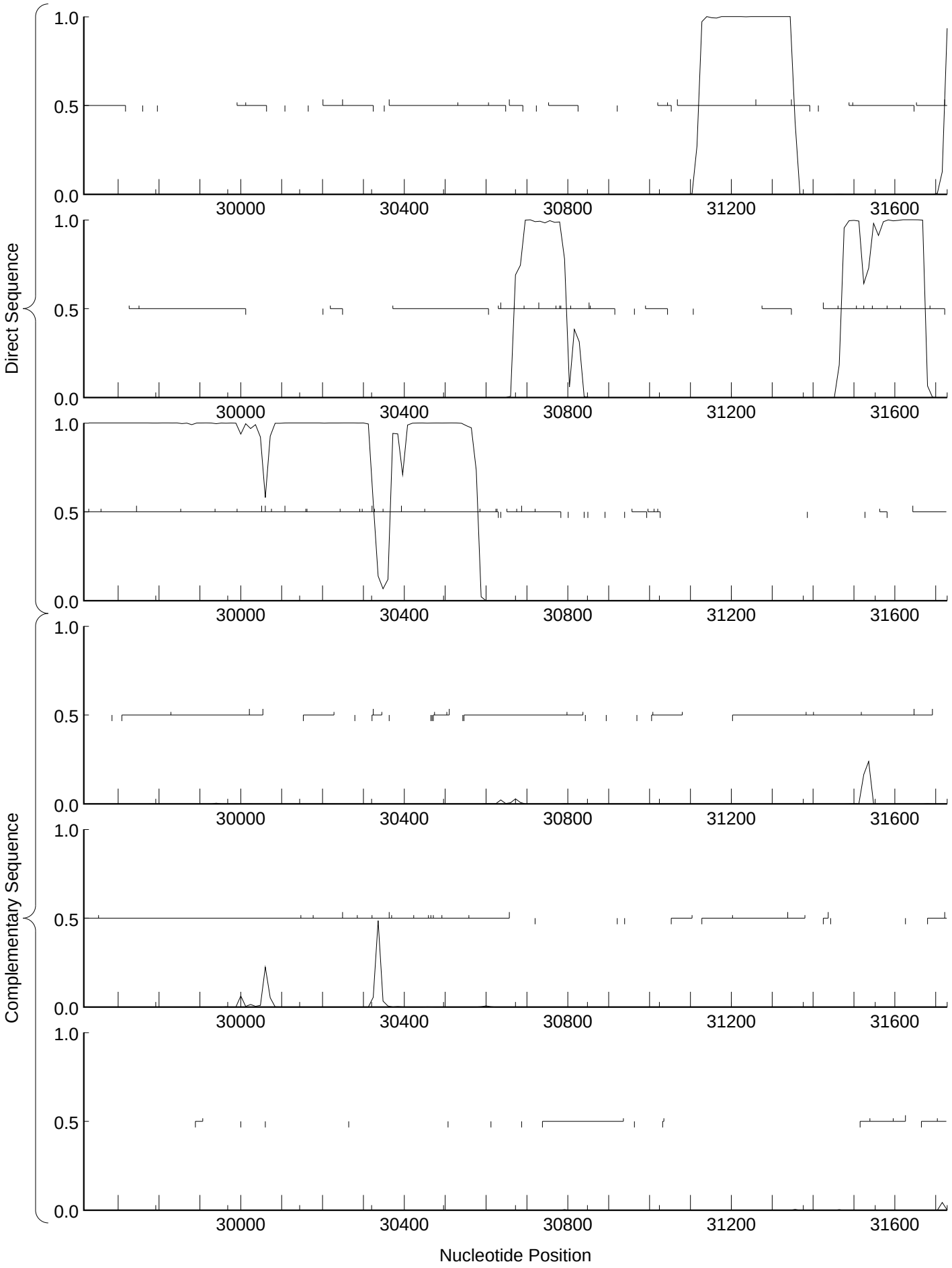


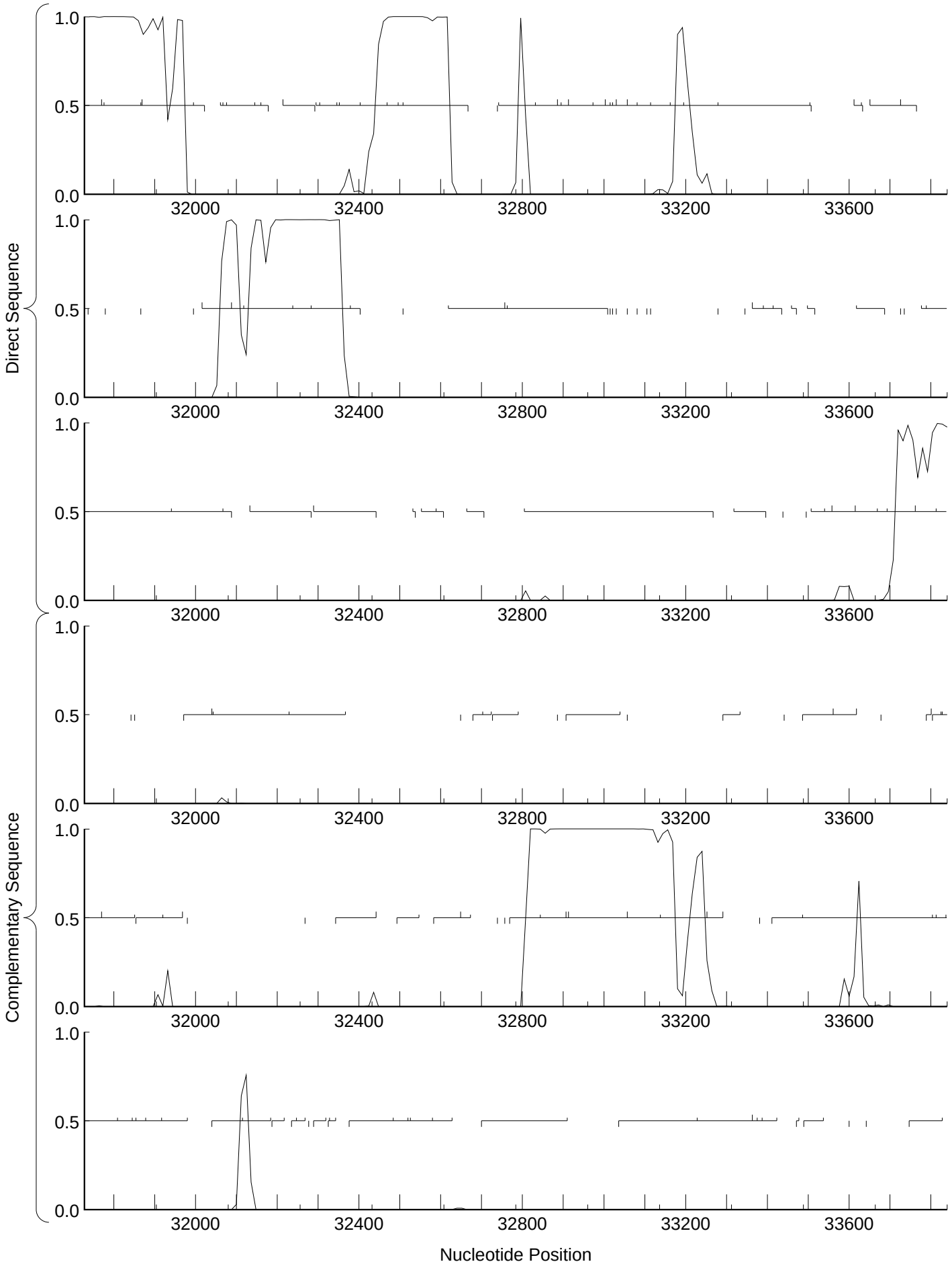

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
25600
26000
26400
26800
27200
Nucleotide Position

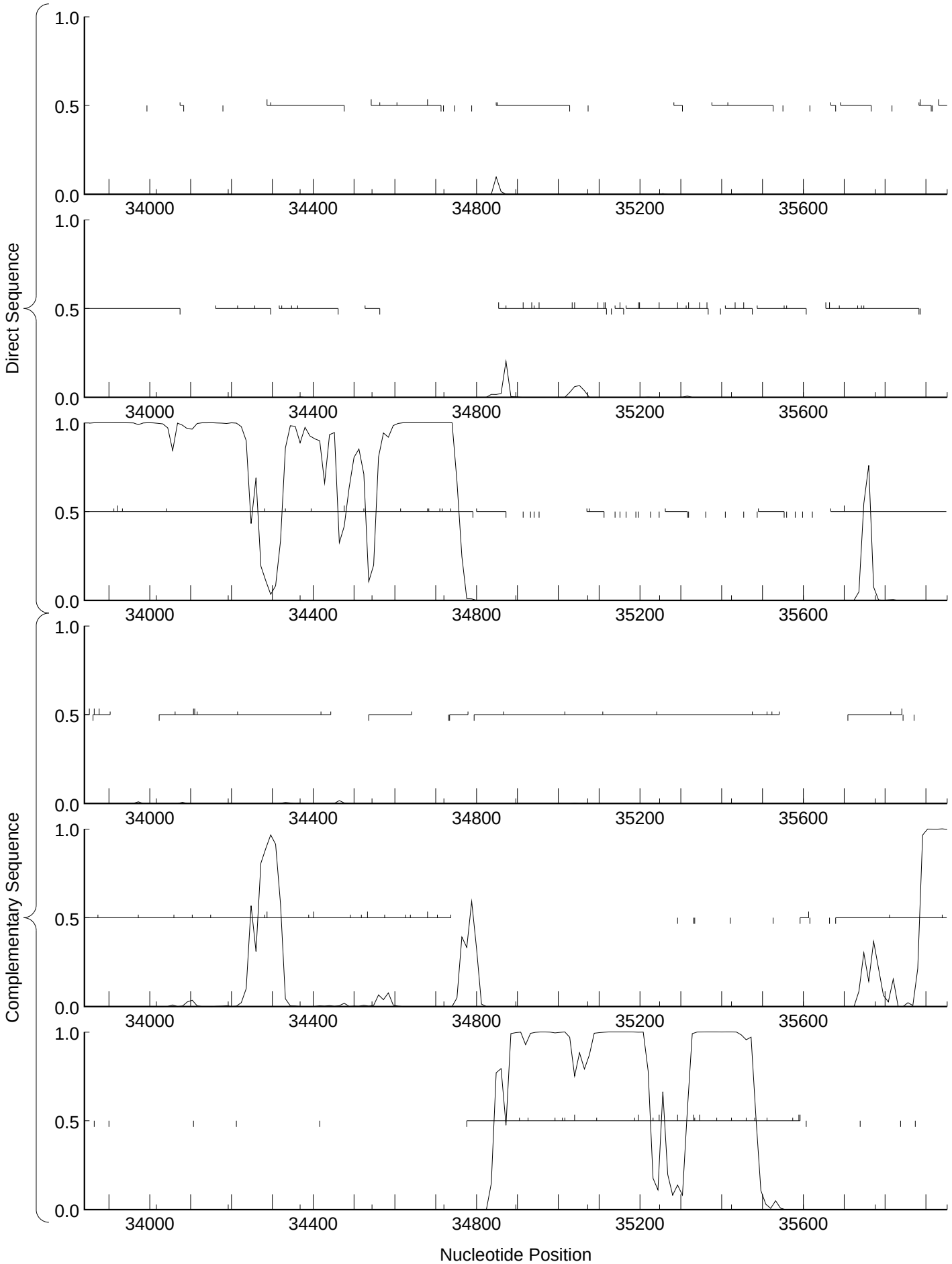
complete sequence, 57256 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 15/29

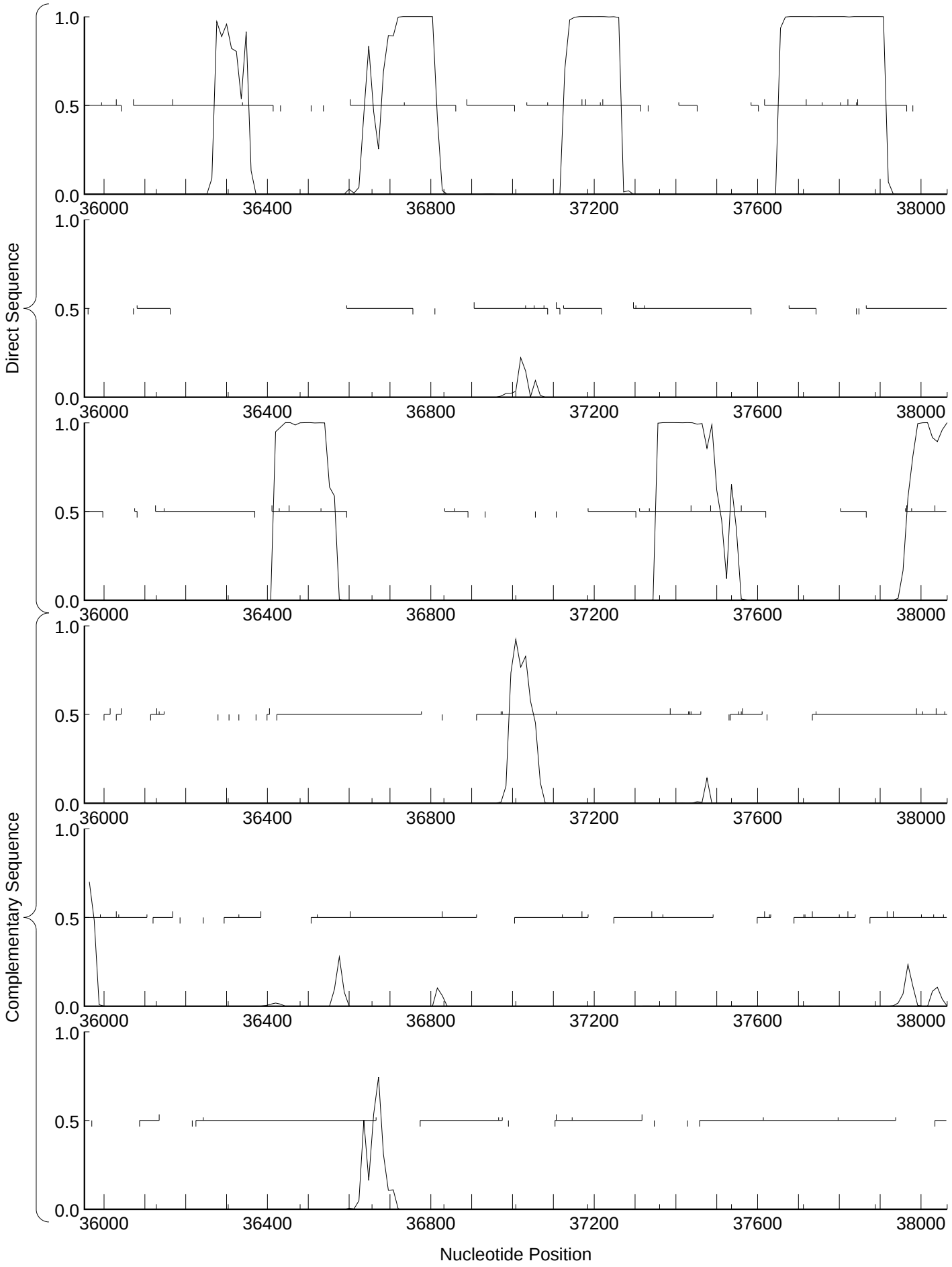


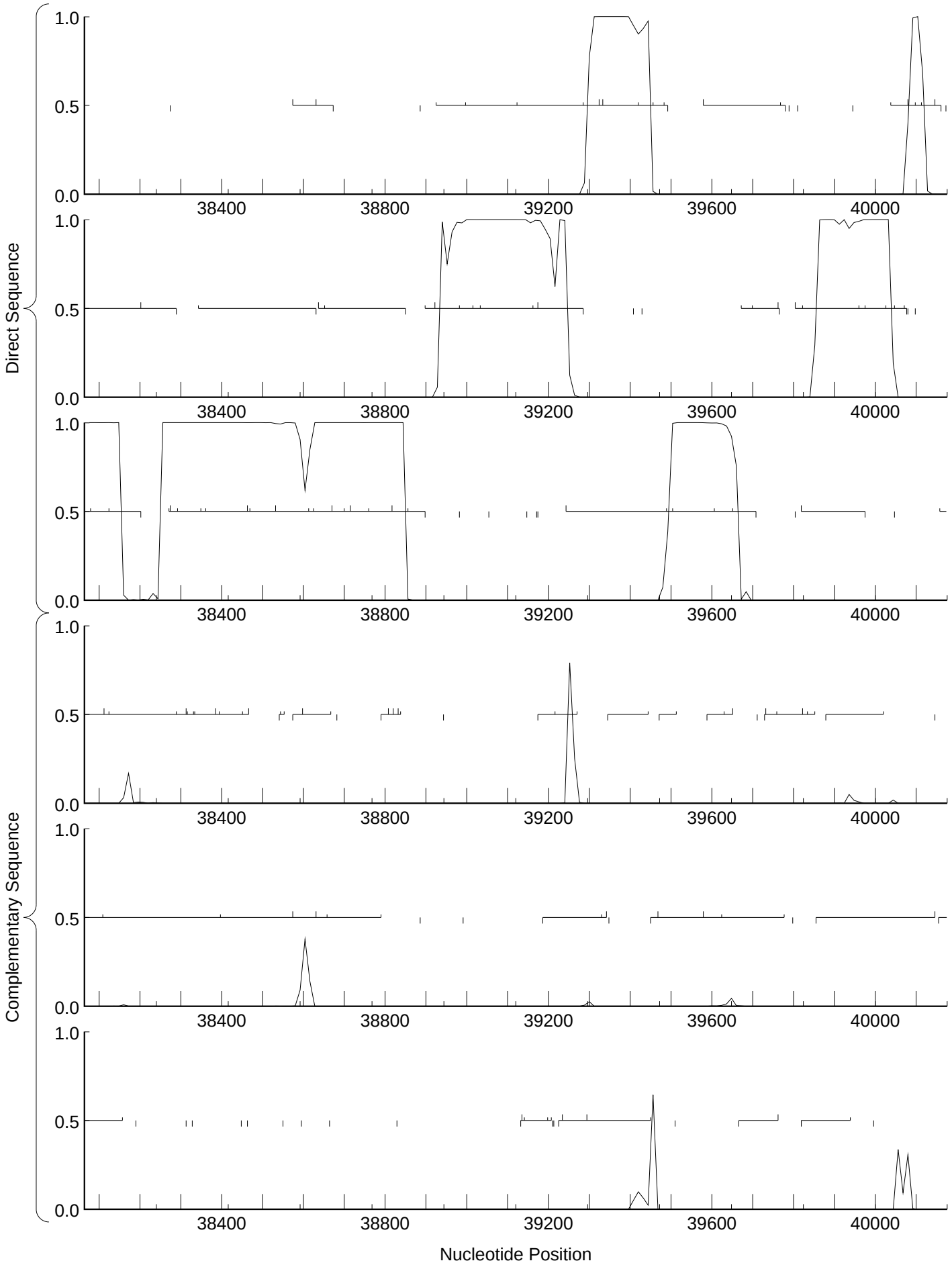
complete sequence, 57256 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 16/29



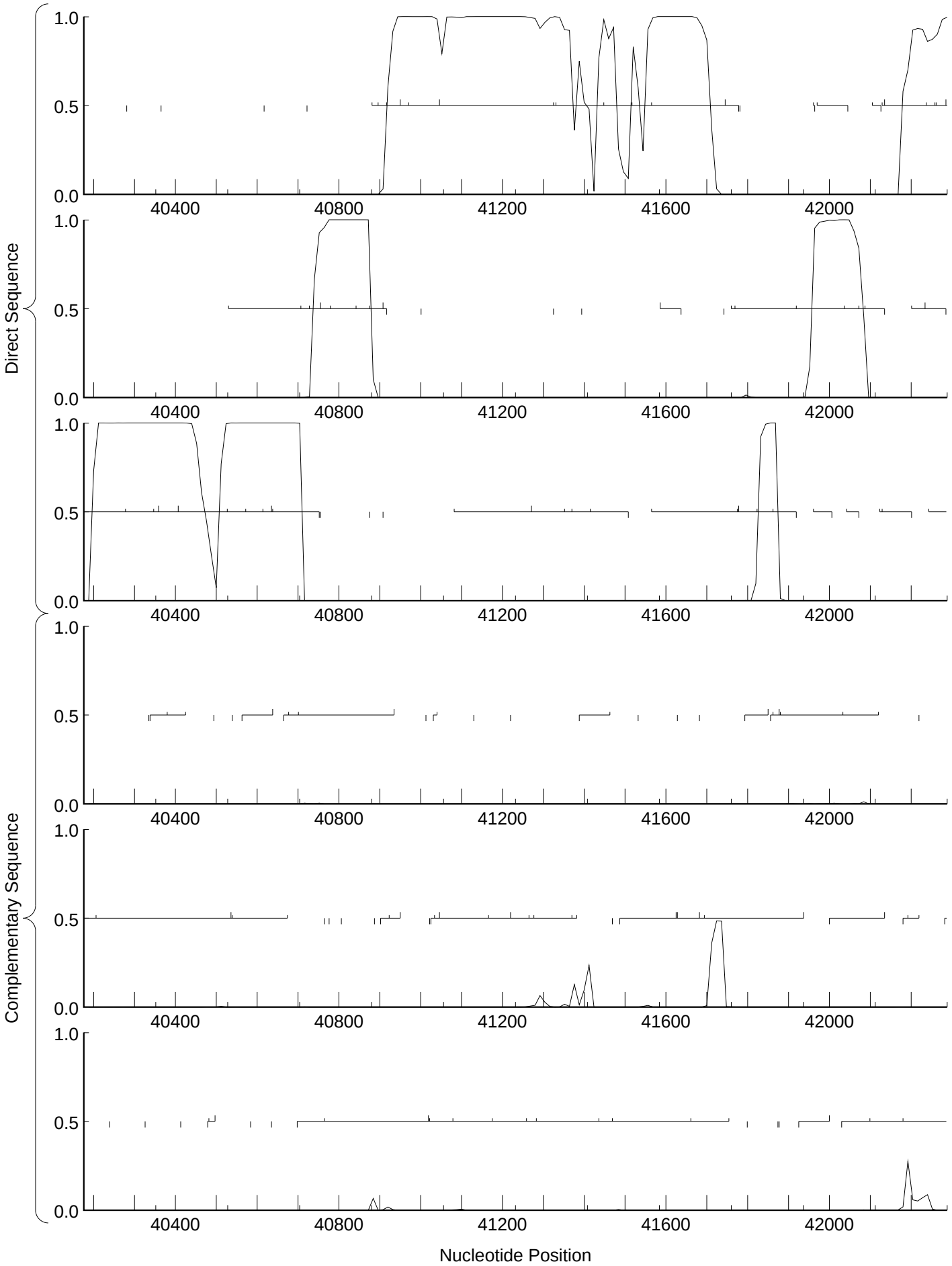

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
32000
32400
32800
33200
33600
Nucleotide Position

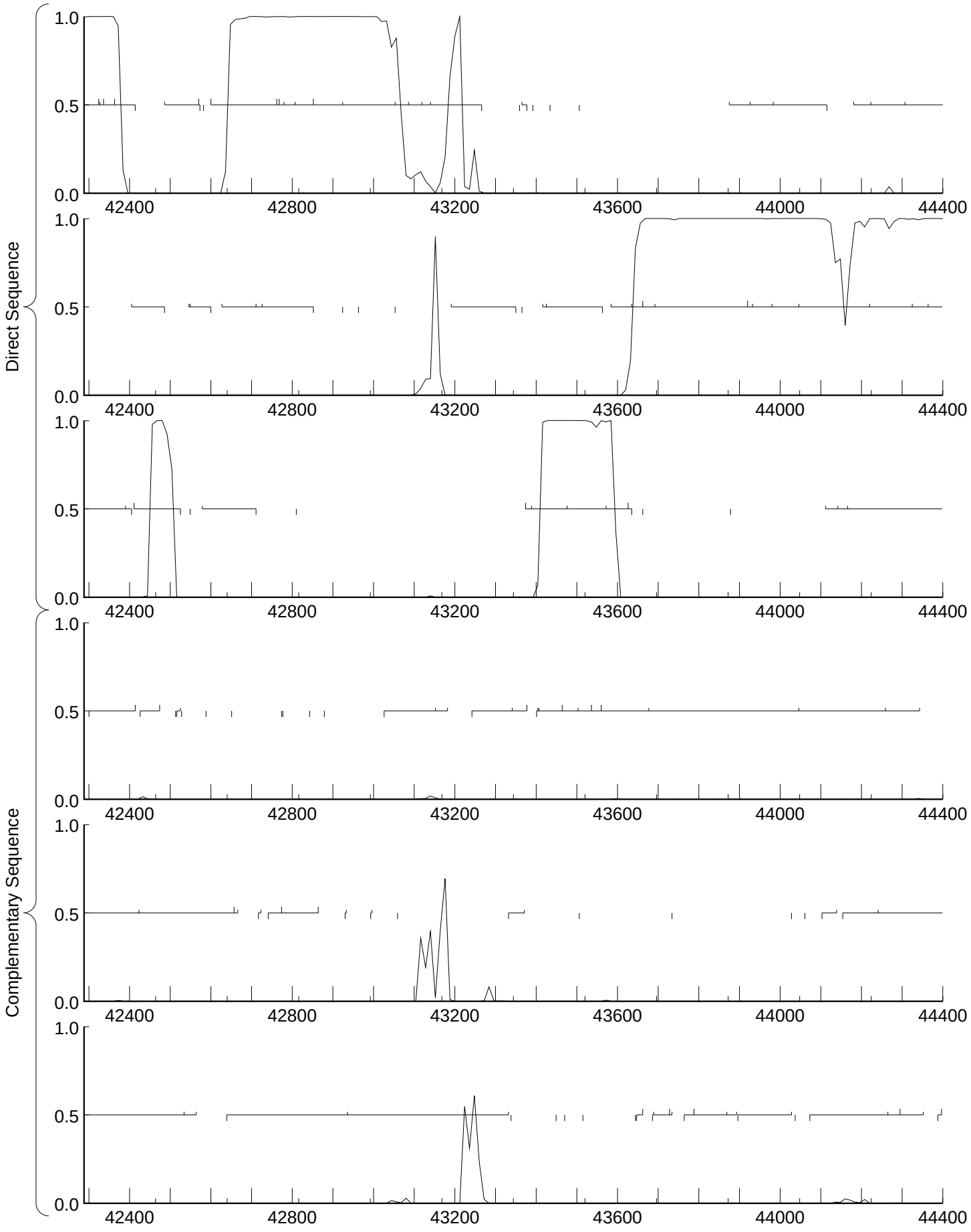

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position


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

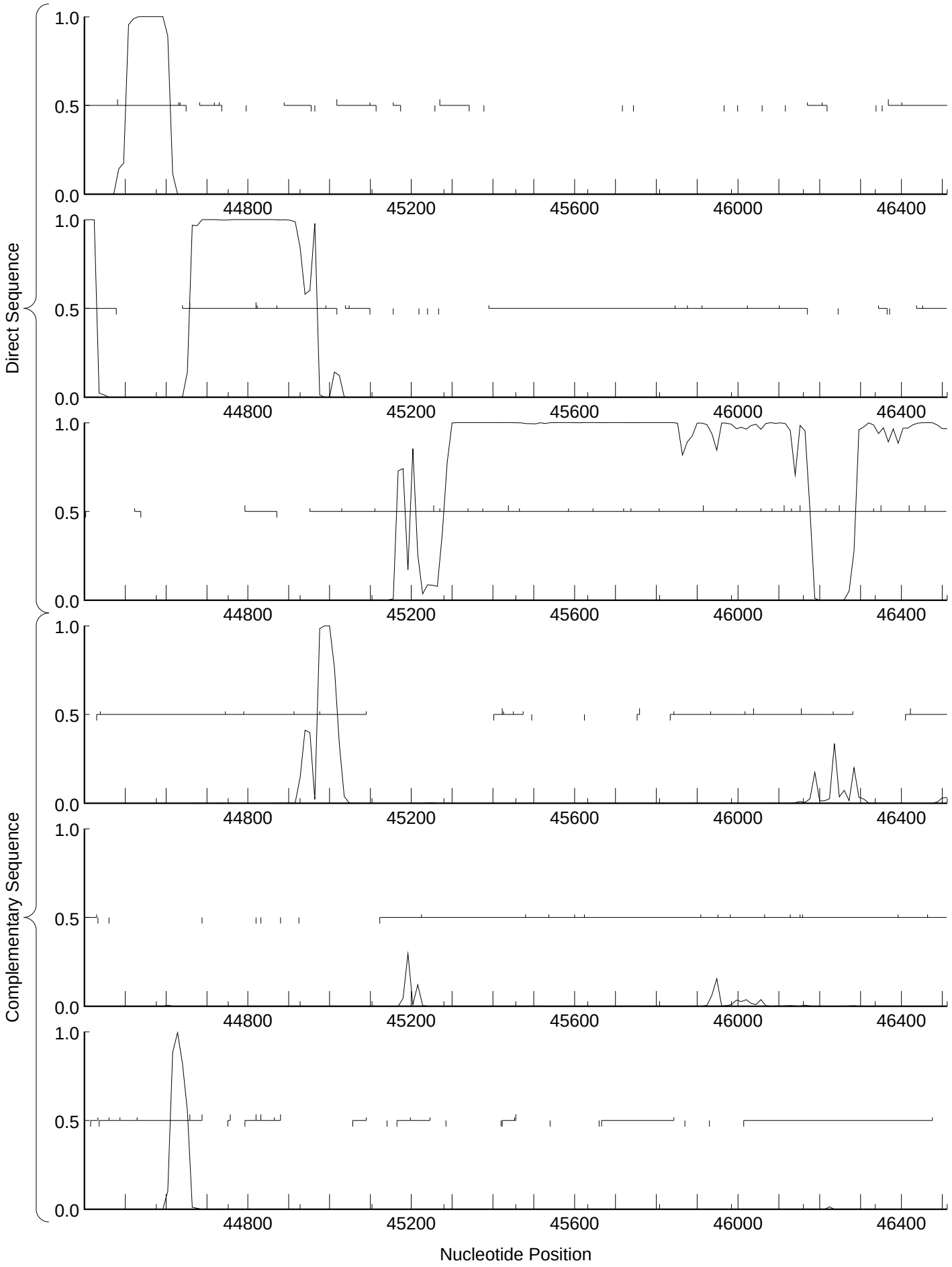

Direct Sequence
Complementary Sequence
1.0
0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 21/29

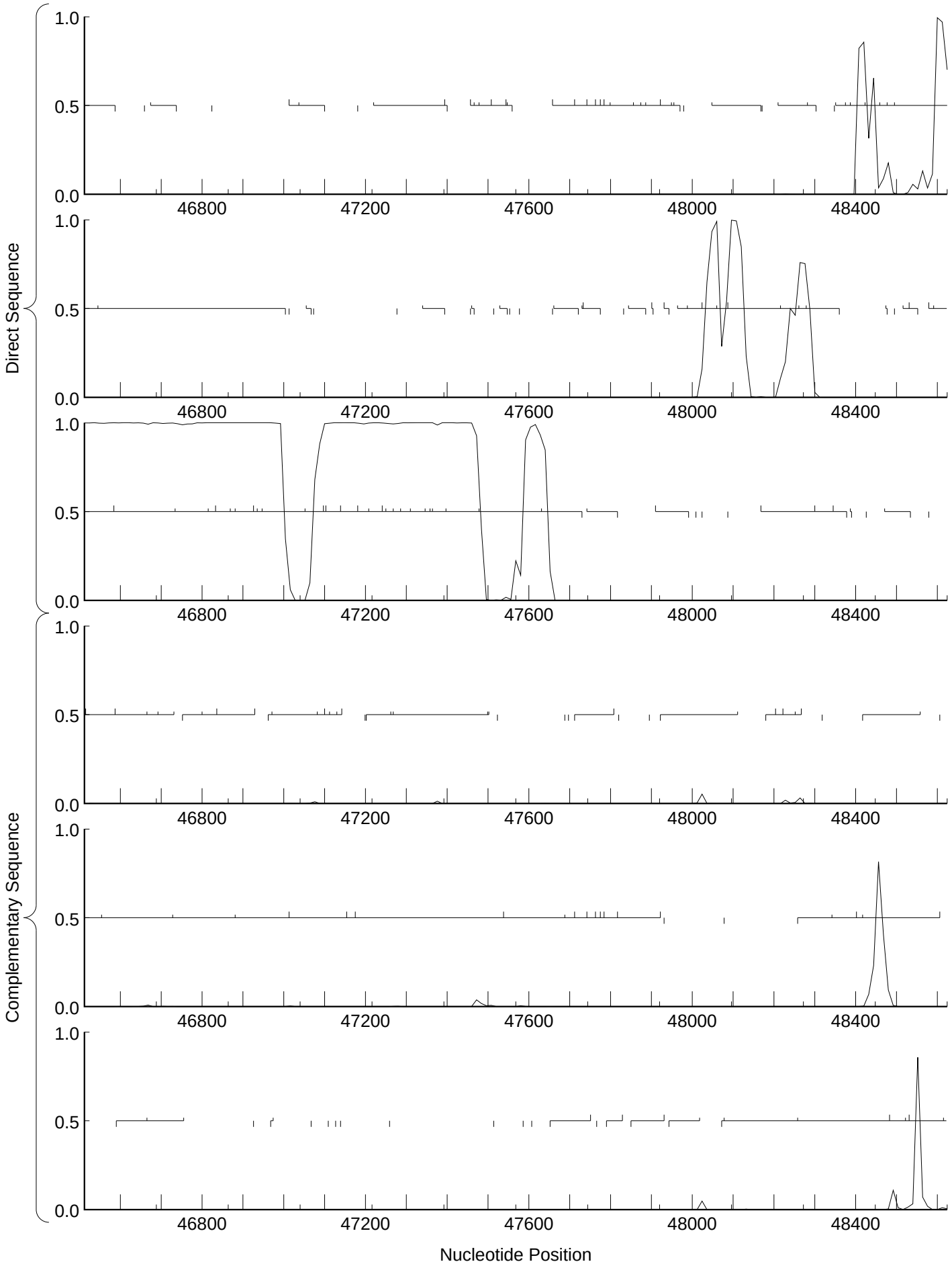



Direct Sequence
Complementary Sequence
1.0
0.5
0.0
42400
42800
43200
43600
44000
44400
Nucleotide Position

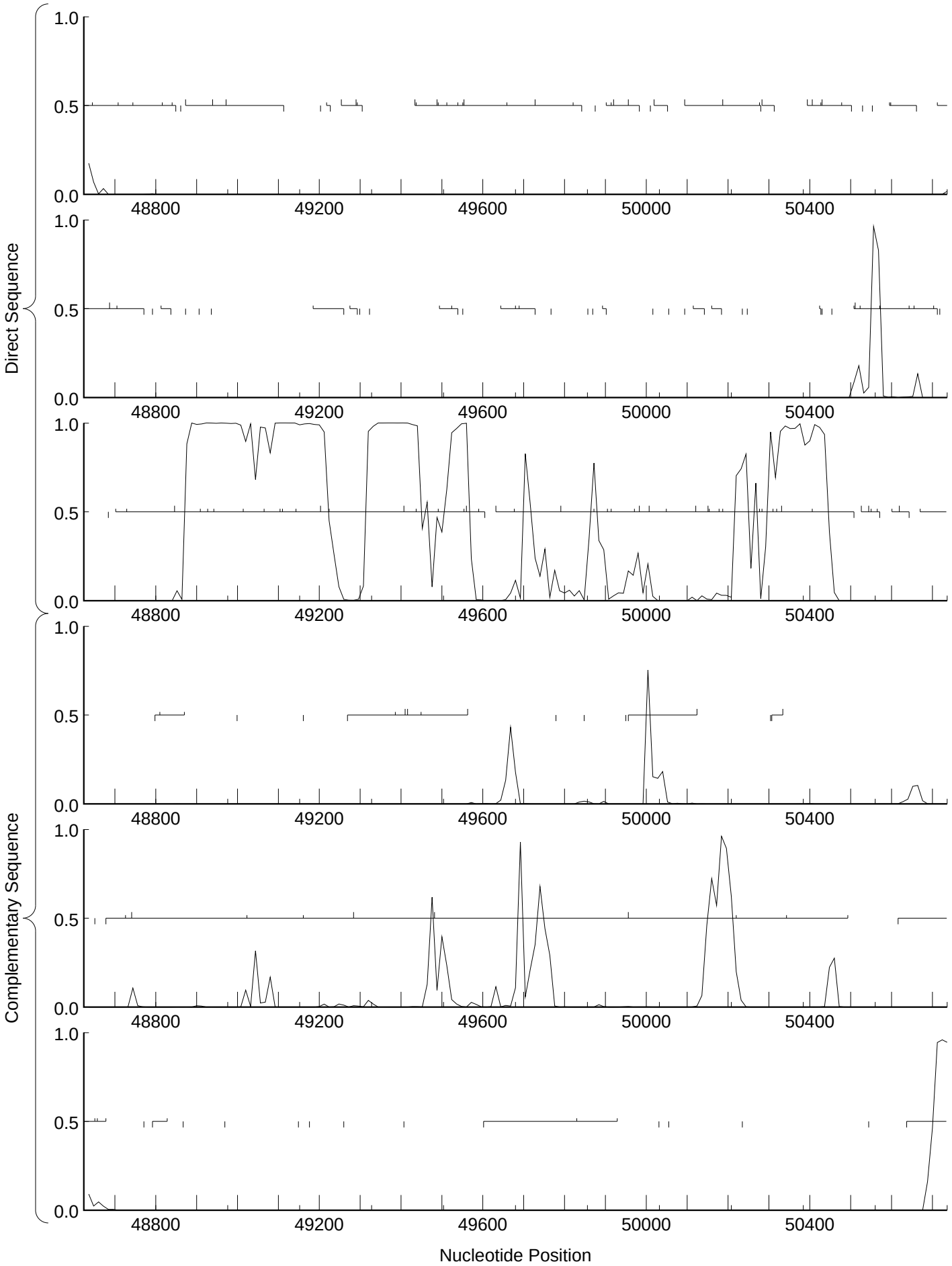
complete sequence, 57256 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 23/29

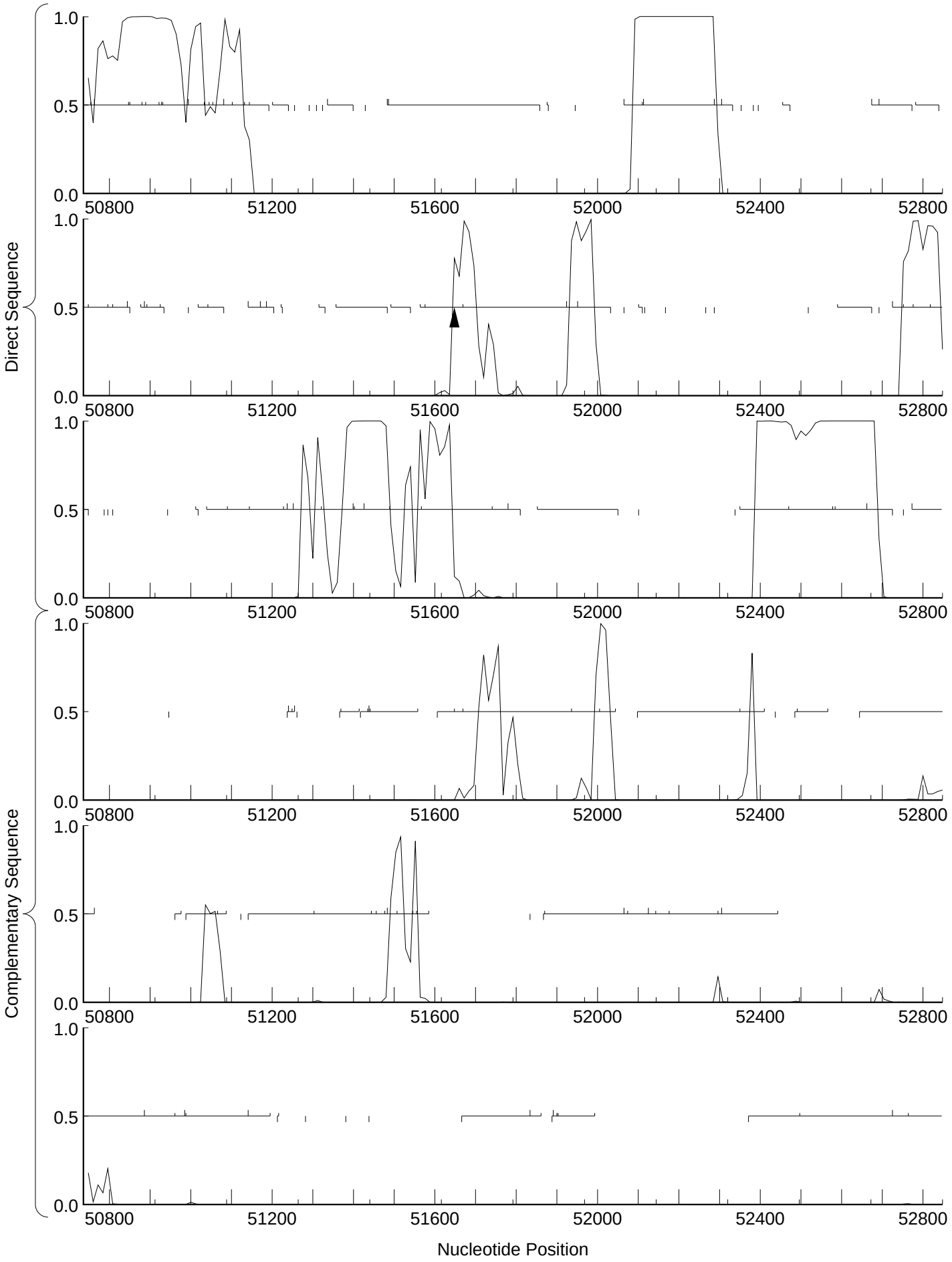


complete sequence, 57256 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 24/29

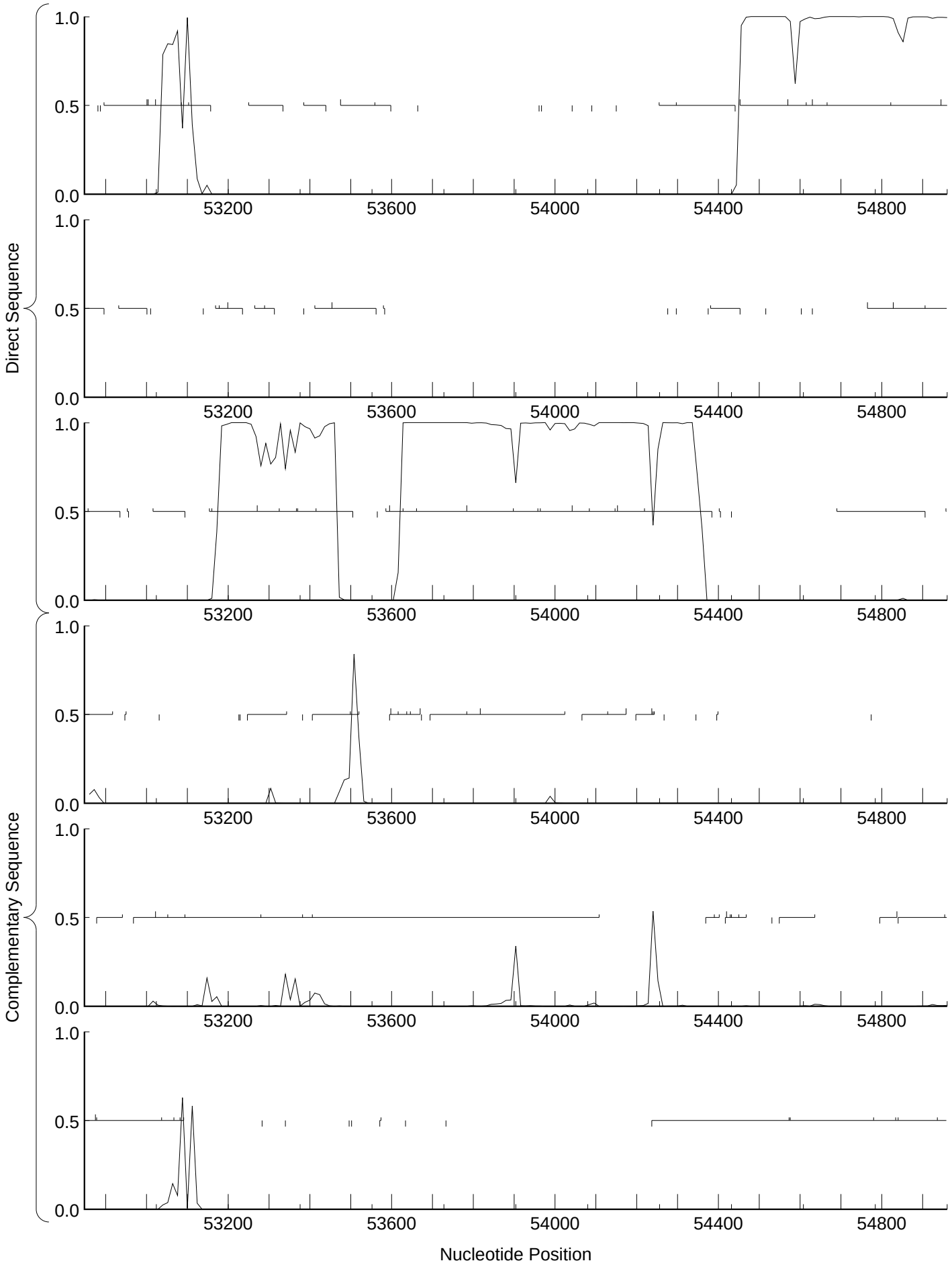


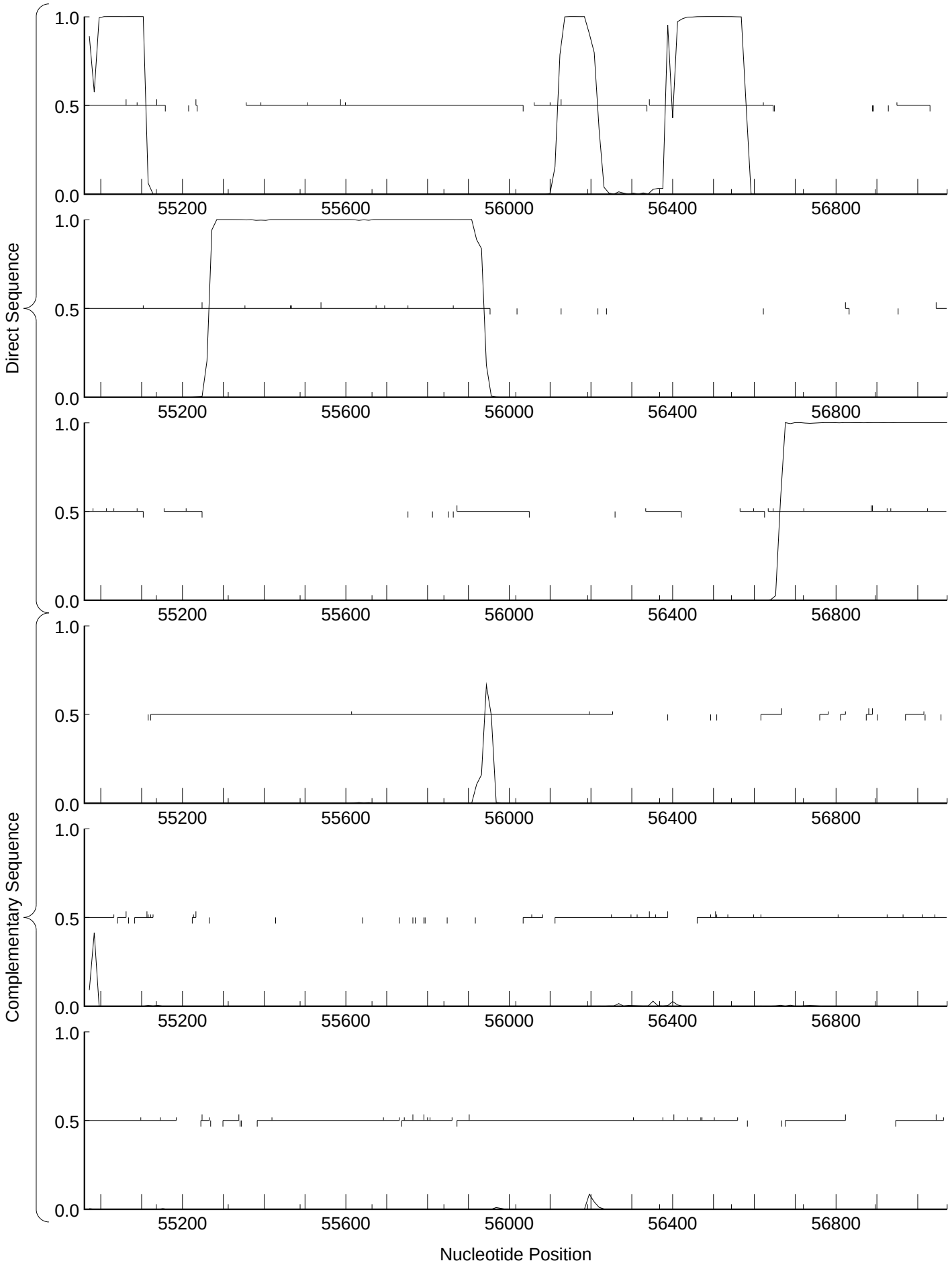
complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 25/29




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

complete sequence, 57256 bp including 11-base 3' overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 27/29




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
55200
55600
56000
56400
56800
Nucleotide Position

complete sequence, 57256 bp including 11-base 3’ overhang (CTCGCGGCCAT), Cluster K4, Order 4, Window 96, Step 12, 29/29

