GeneMark

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

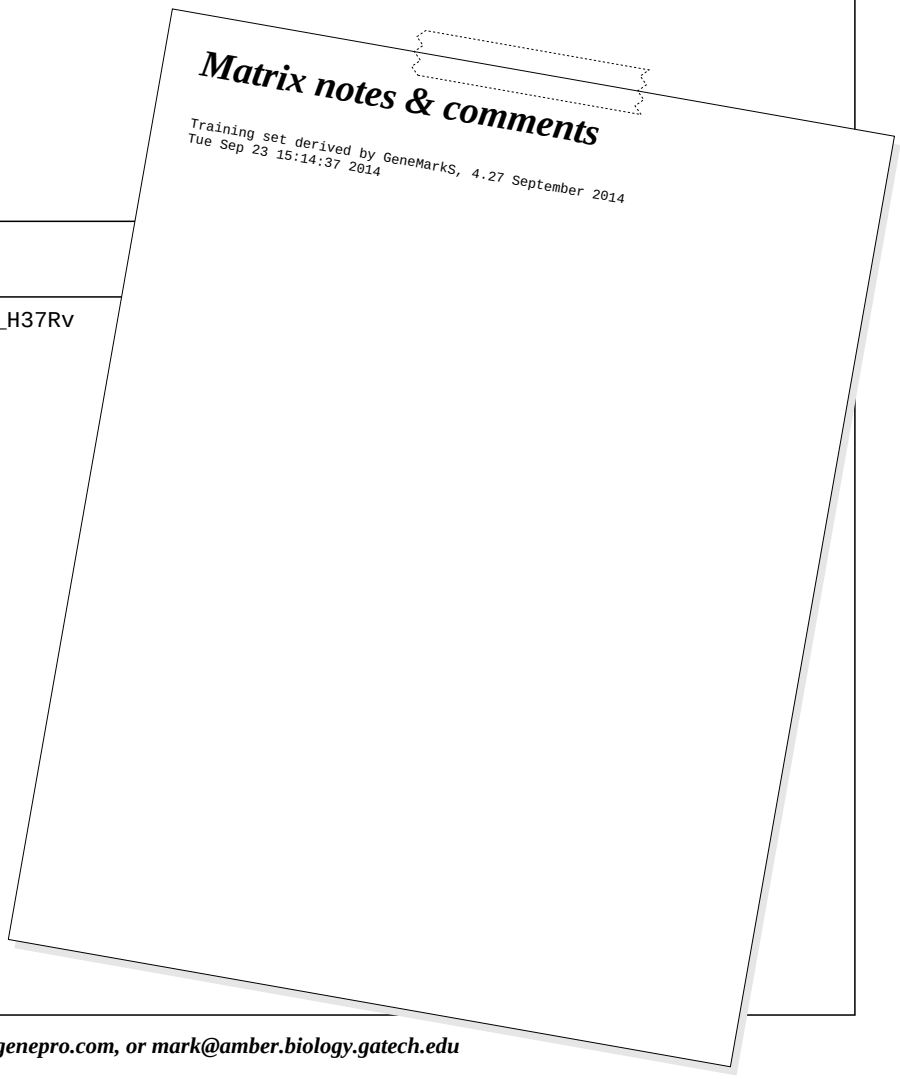
PROGRAM INFORMATION

```
Sequence         :  Mycobacterium phage Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTG
Analysis Date    :  3/28/24 at 11:31:40
Pages            :  37
Sequence Length  :  75360 bp
GC Content       :   63.01%

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

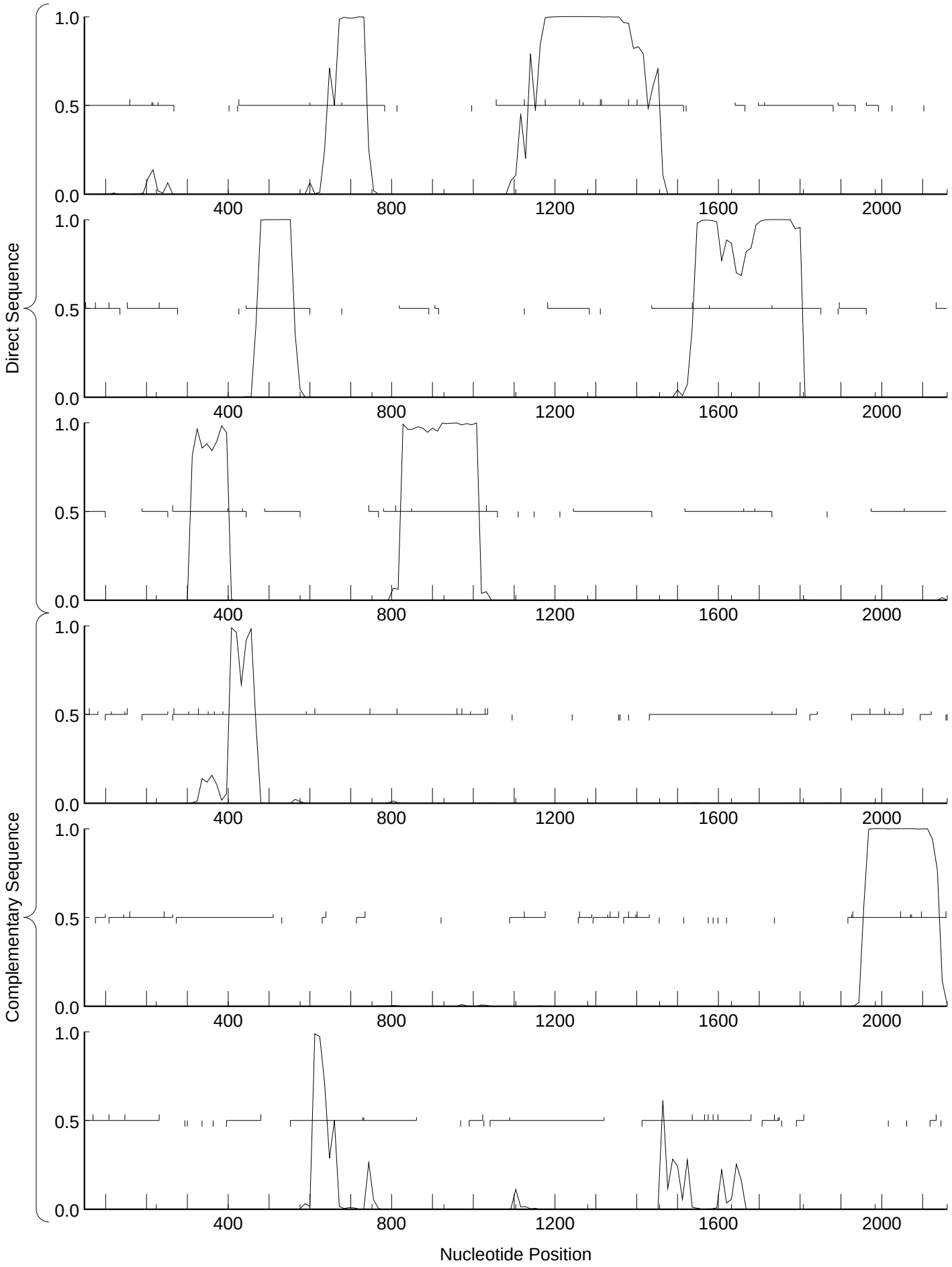
```
Matrix     :  Mycobacterium_tuberculosis_H37Rv
Author     :  -

Order      :  4

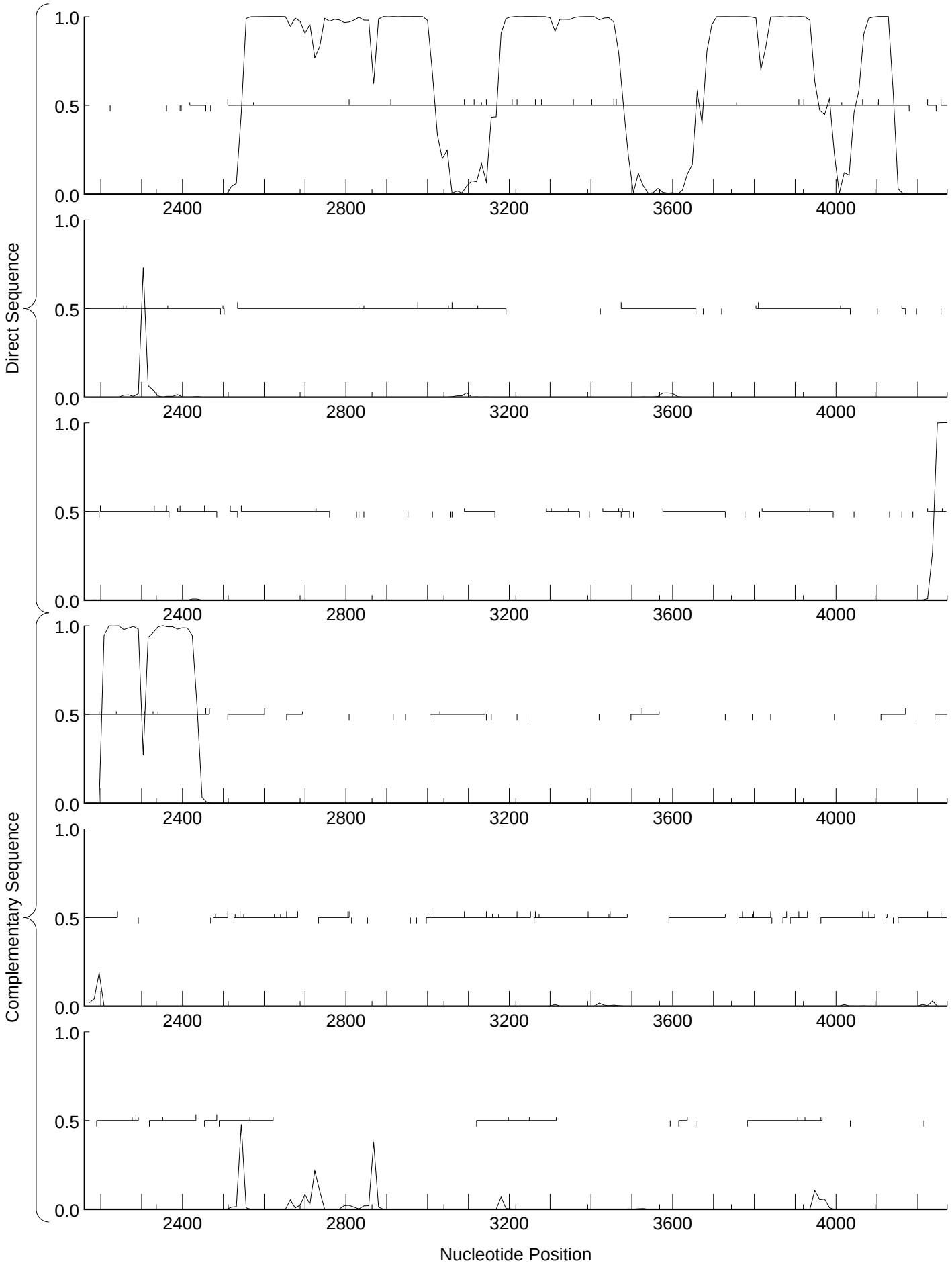

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

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

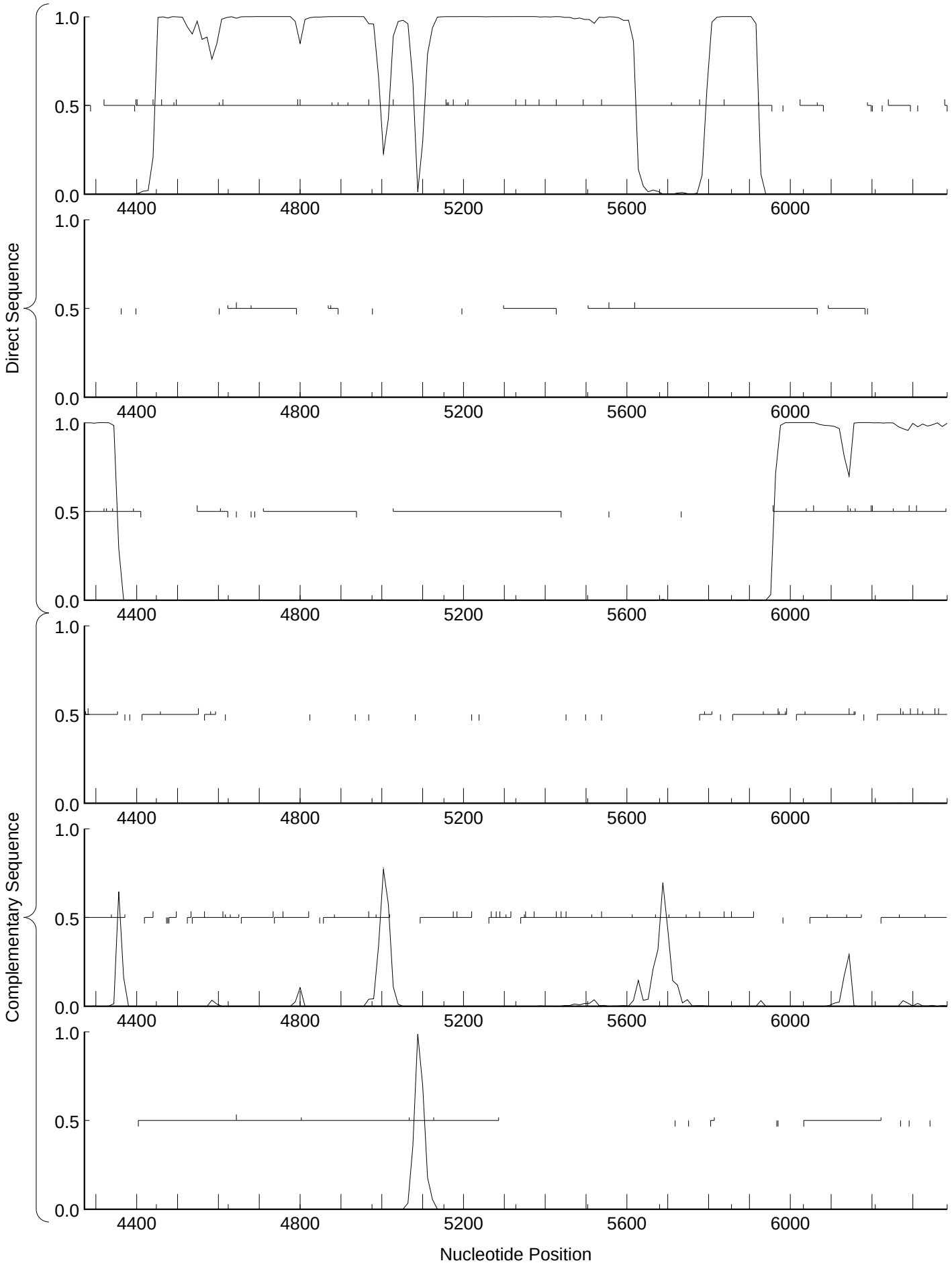
e Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 2/37



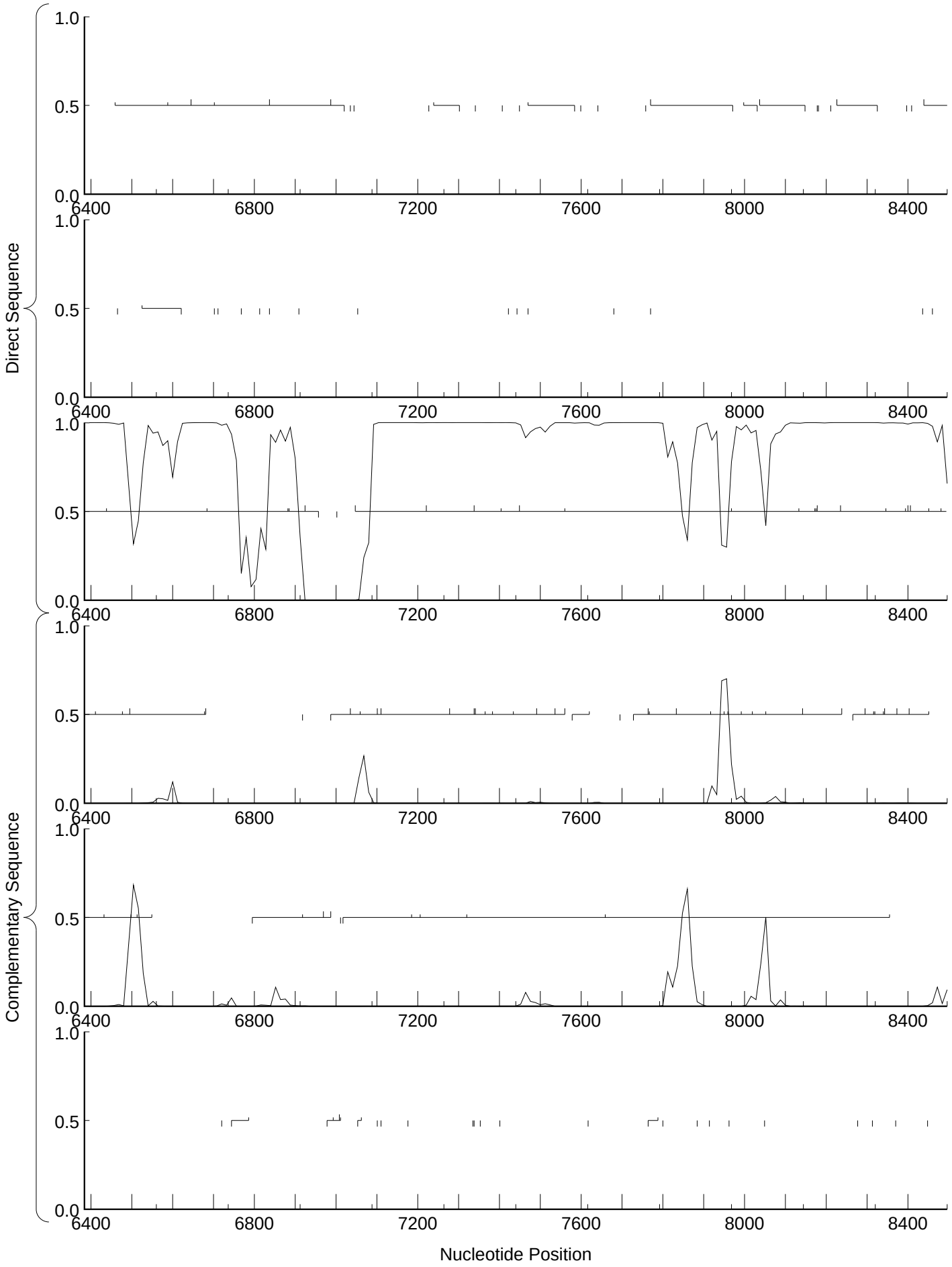
e Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 3/37



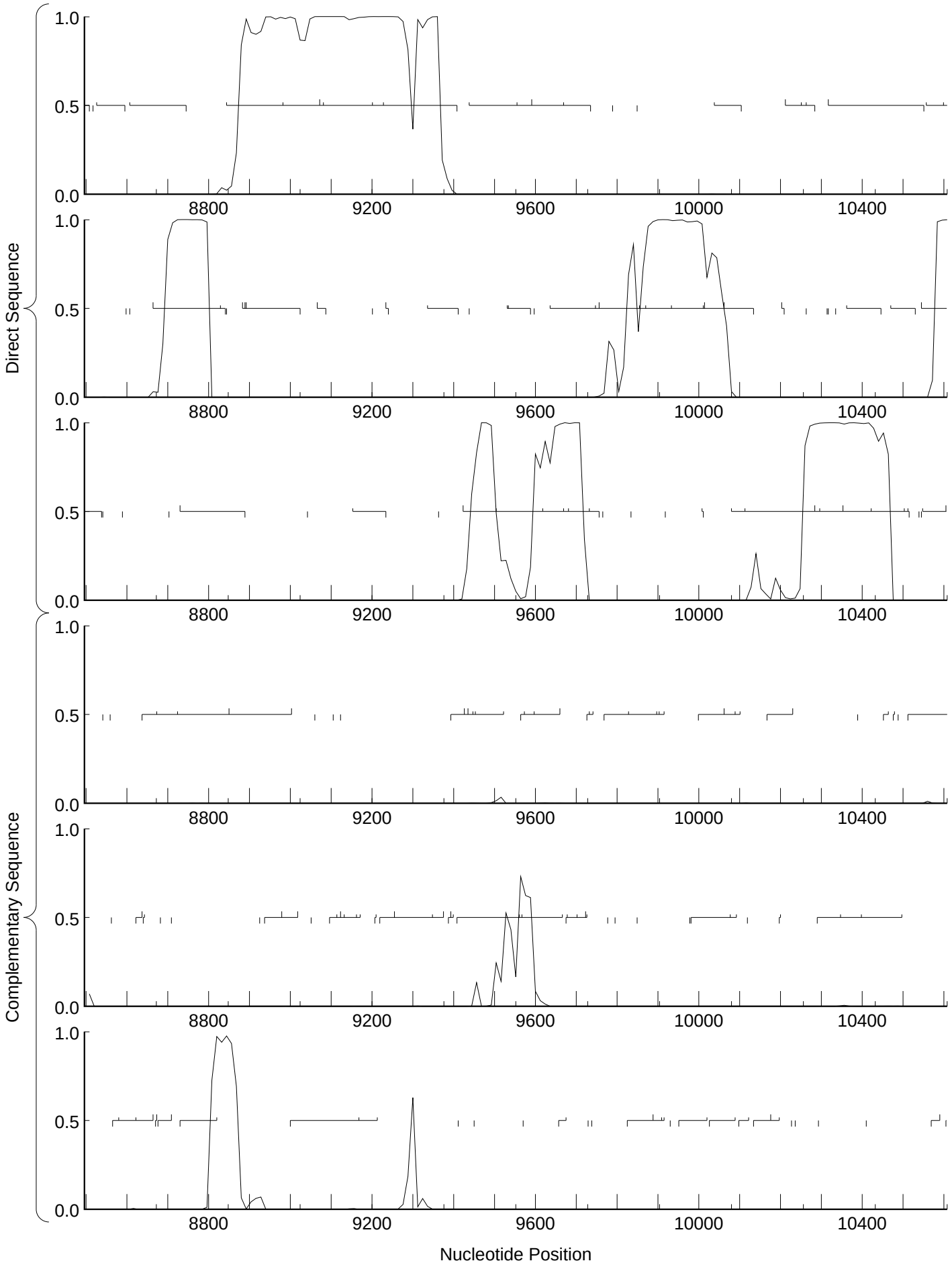
e Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 4/37



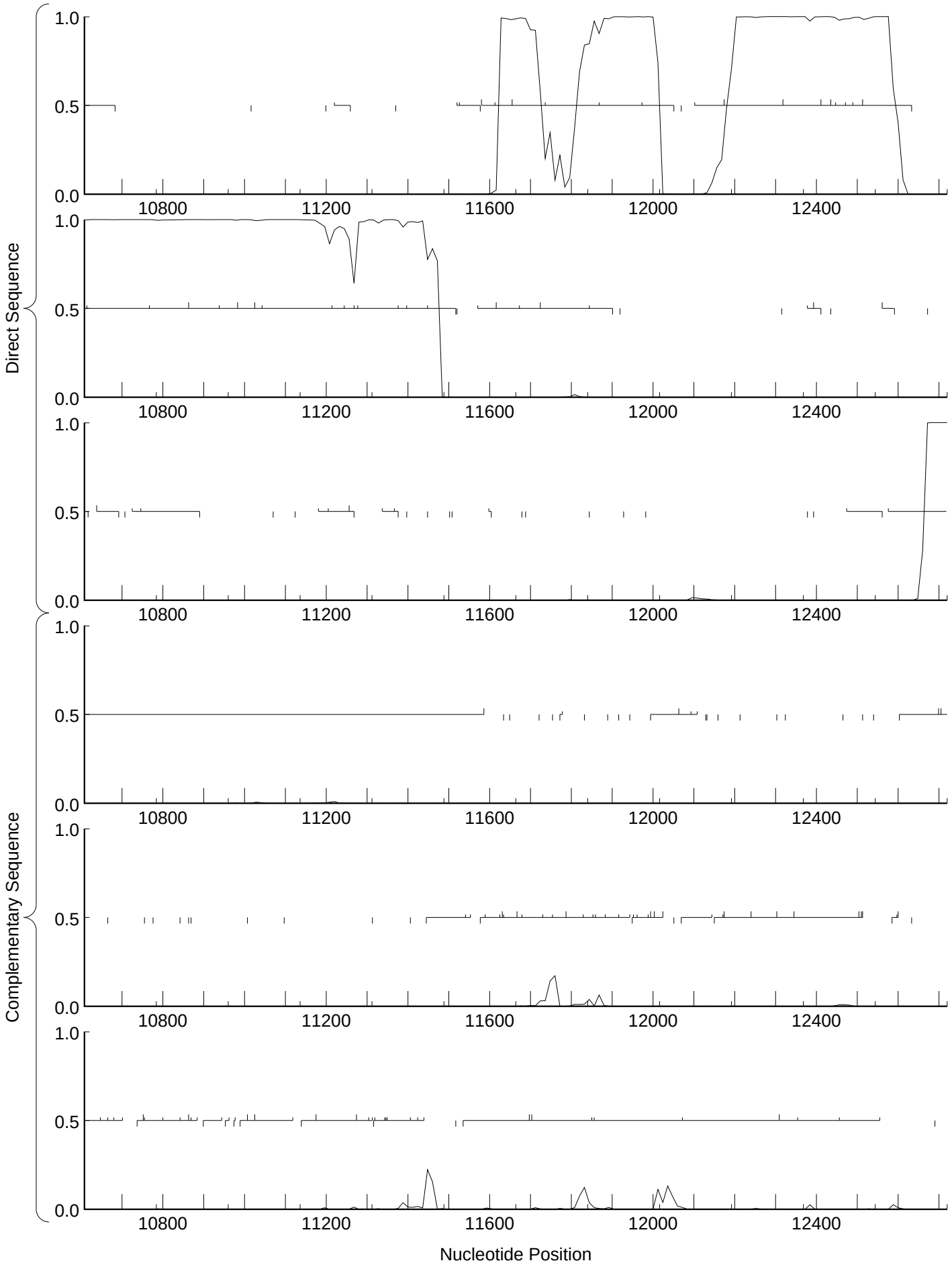
e Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 5/37



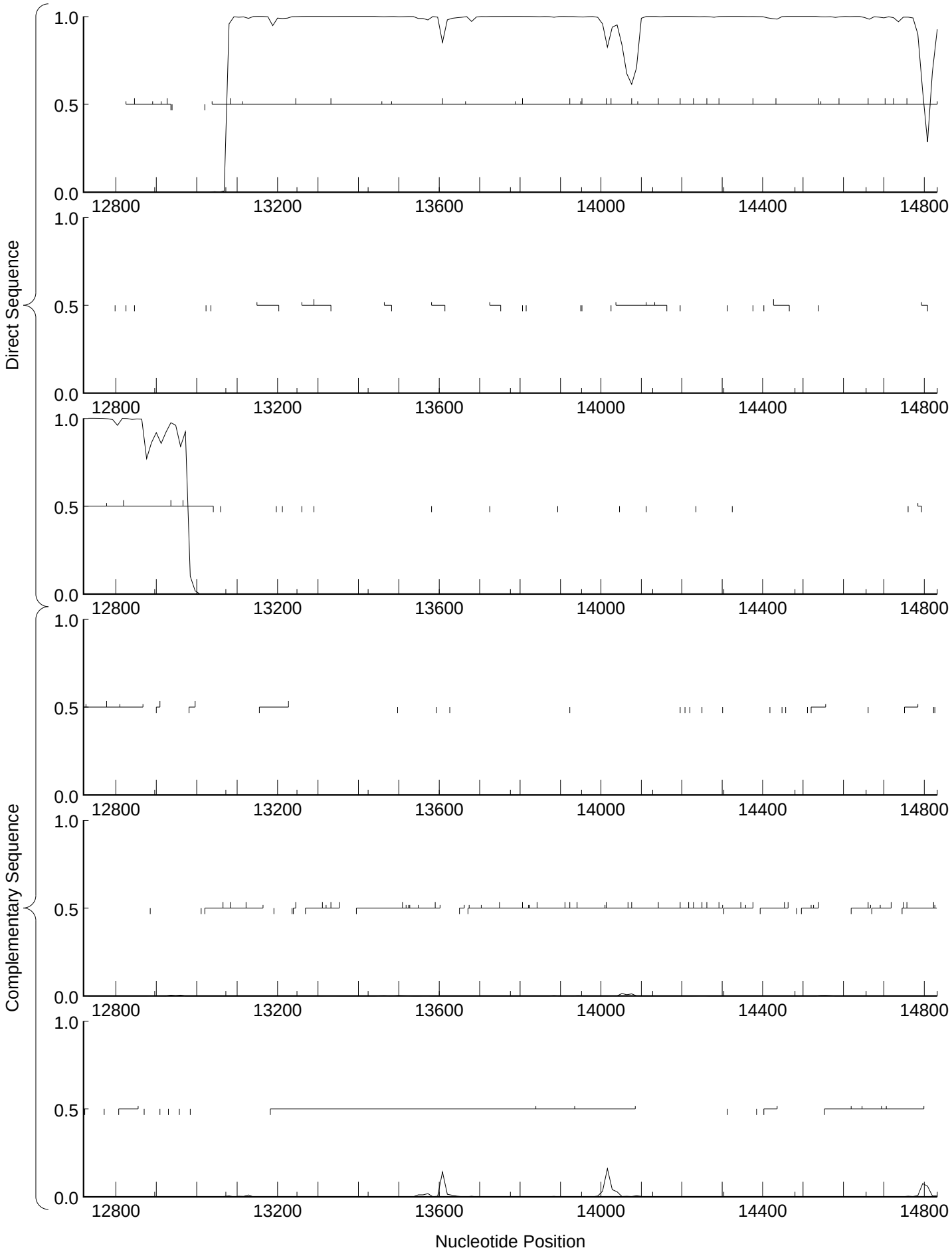
e Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 6/37



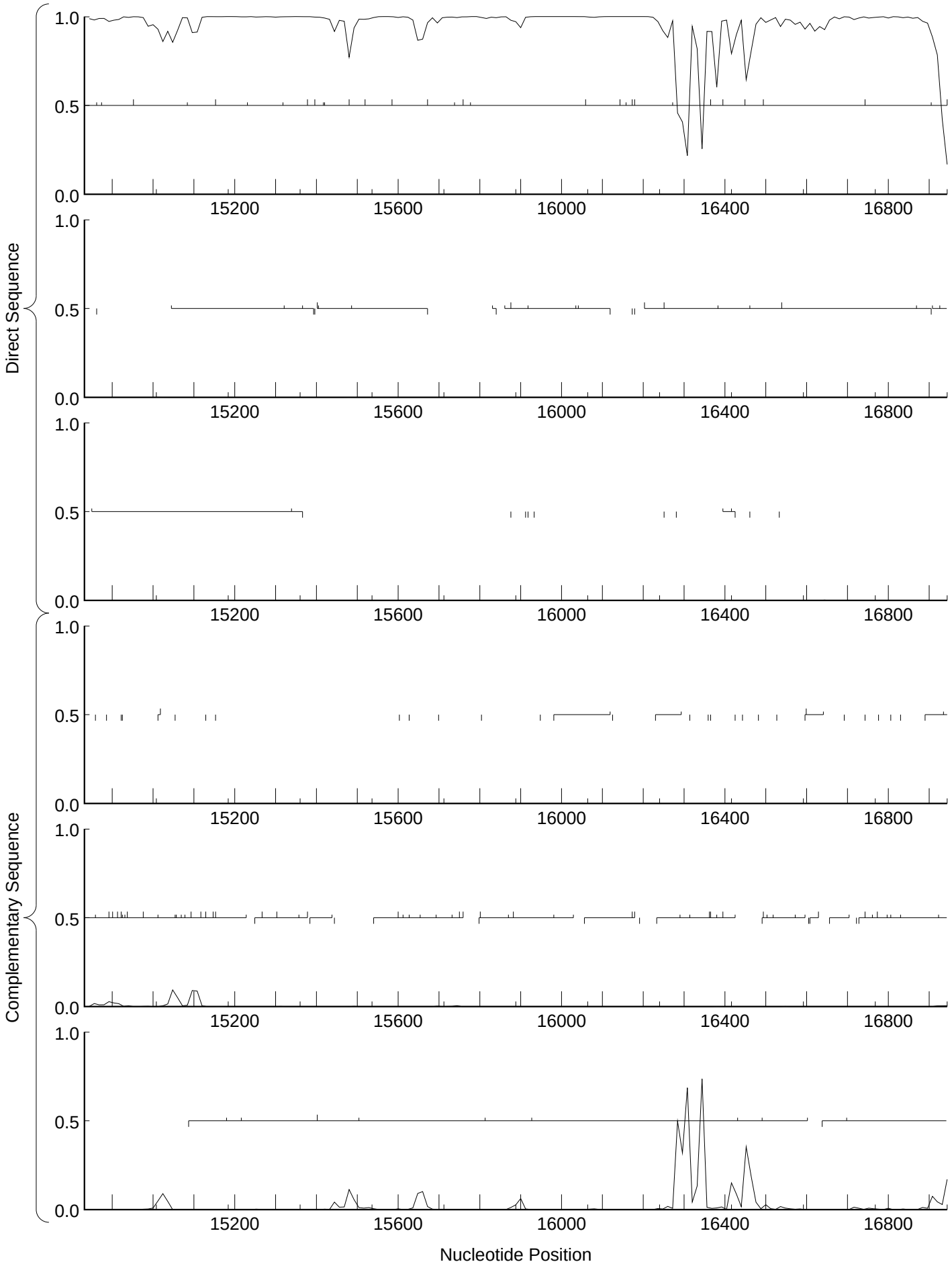
e Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 7/37



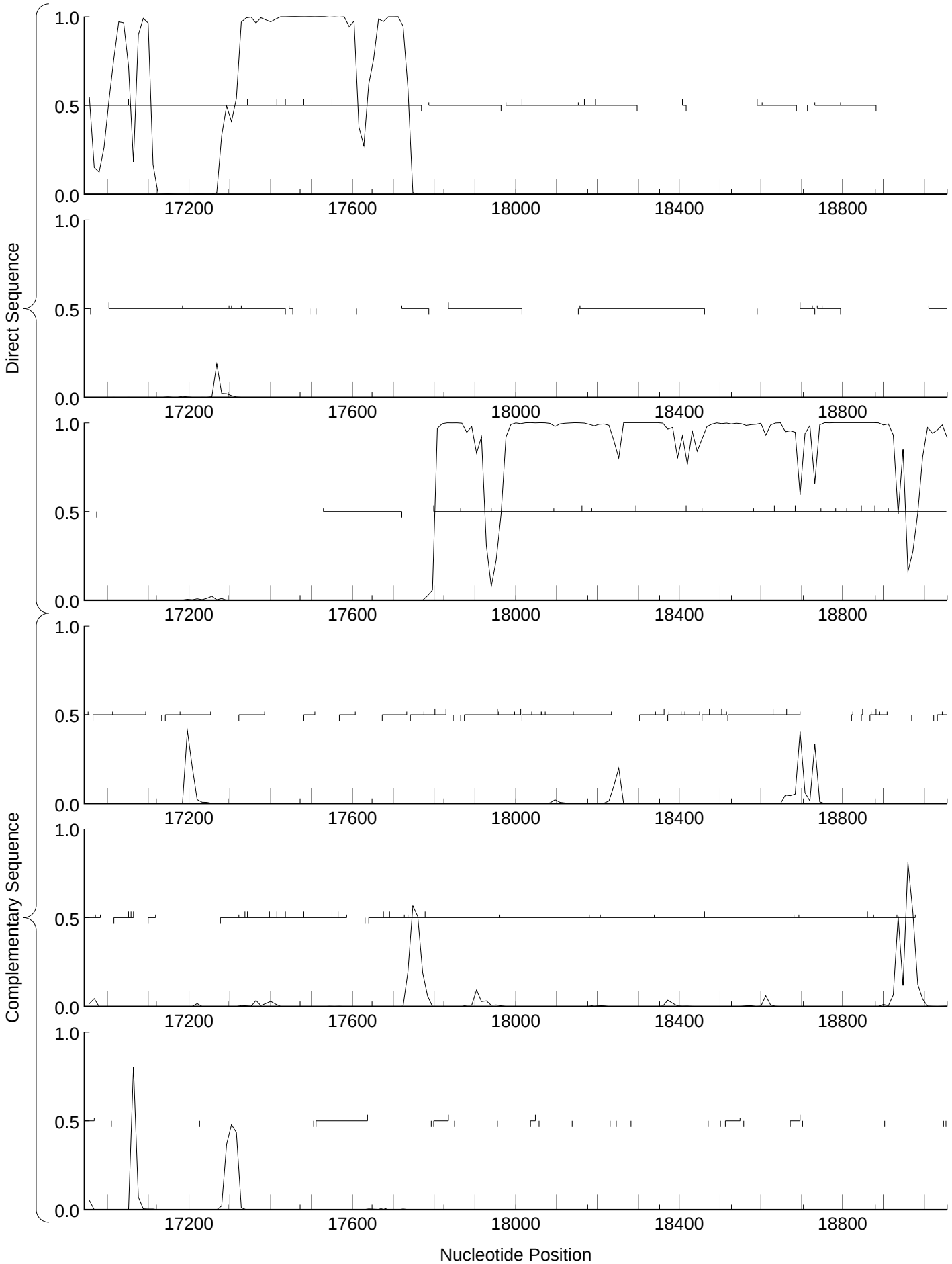
e Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 8/37



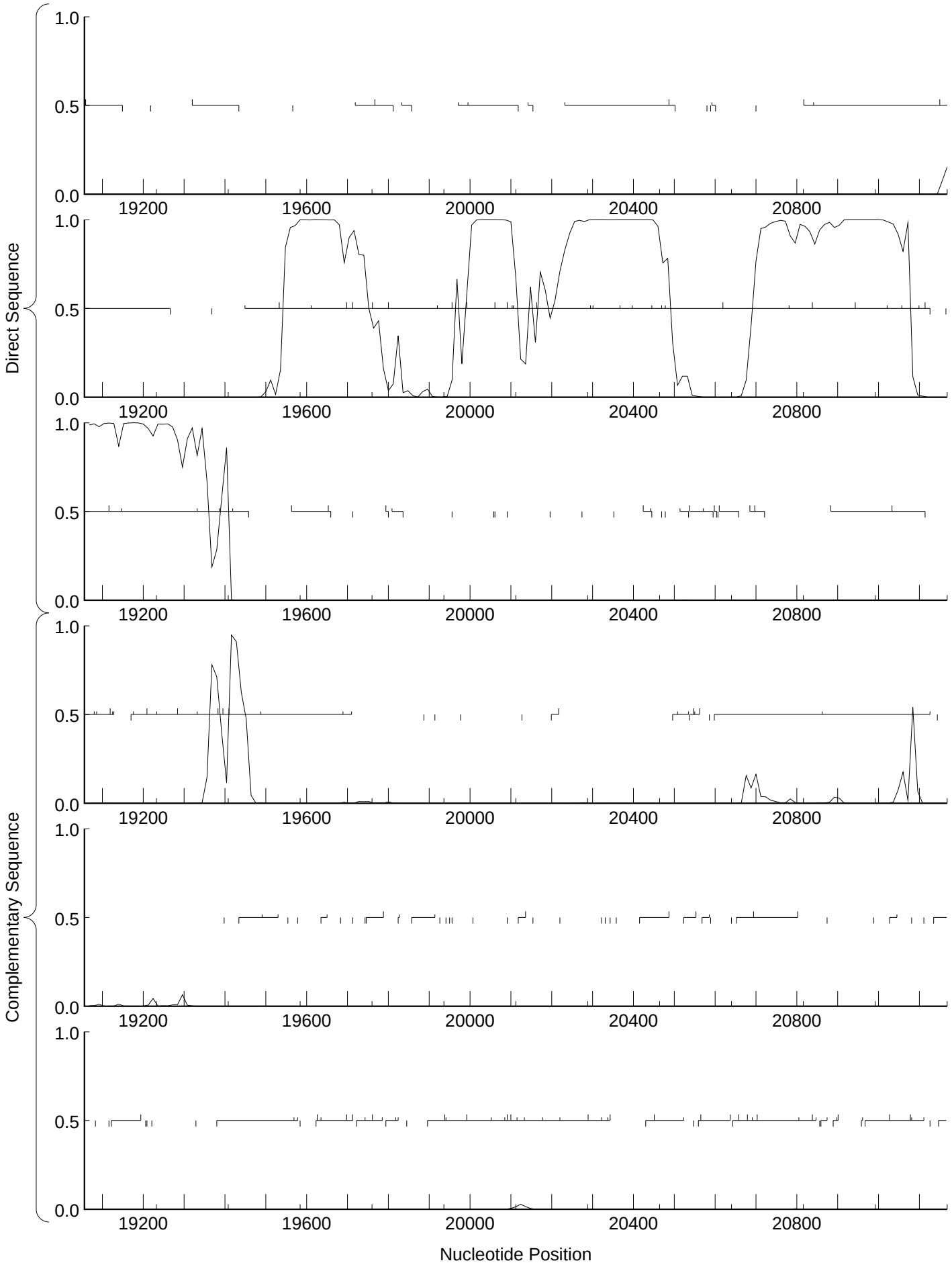
e Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 9/37



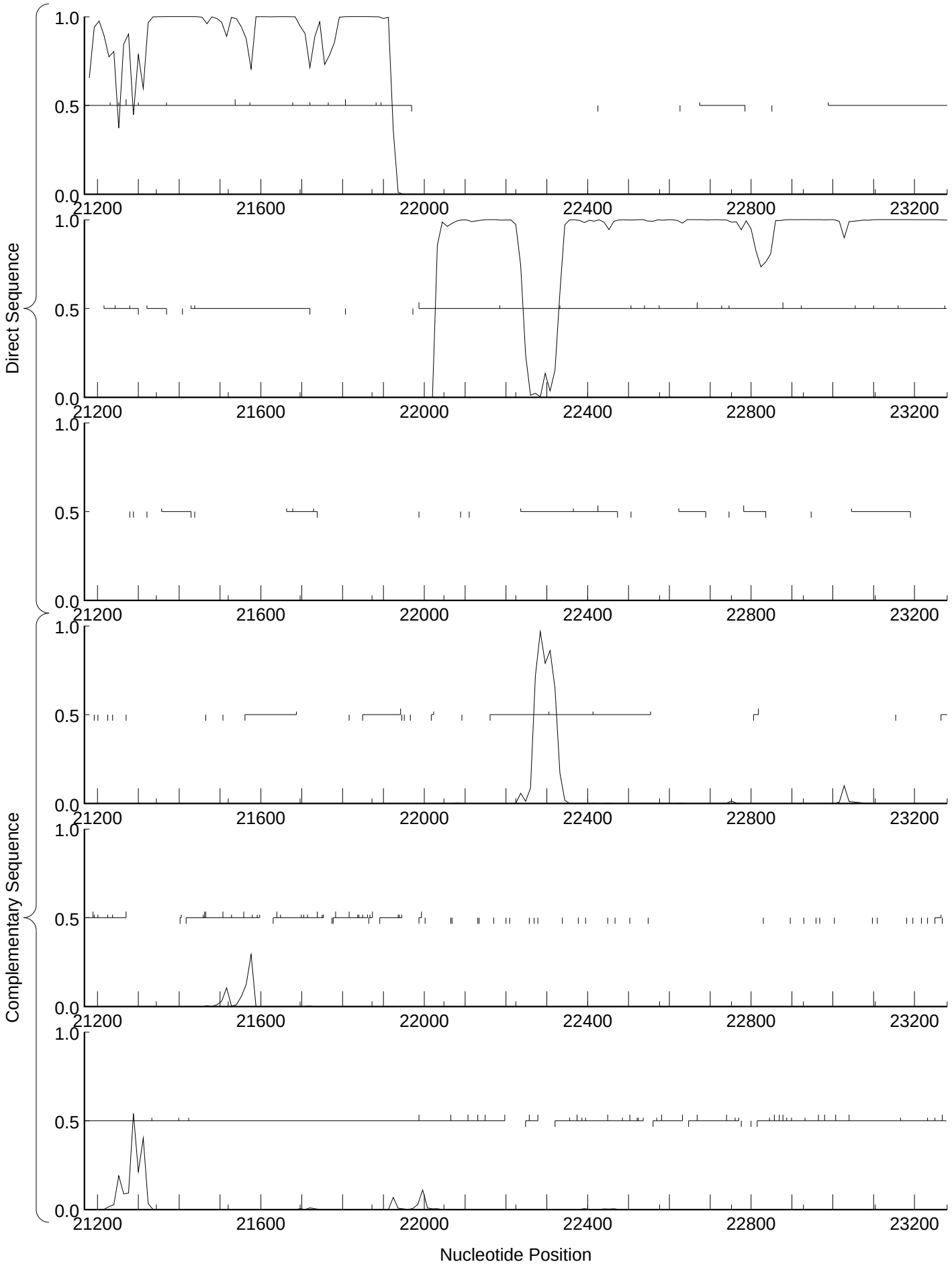
Lilizi complete sequence, 75360 bp including 9-base 3' overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 10/37



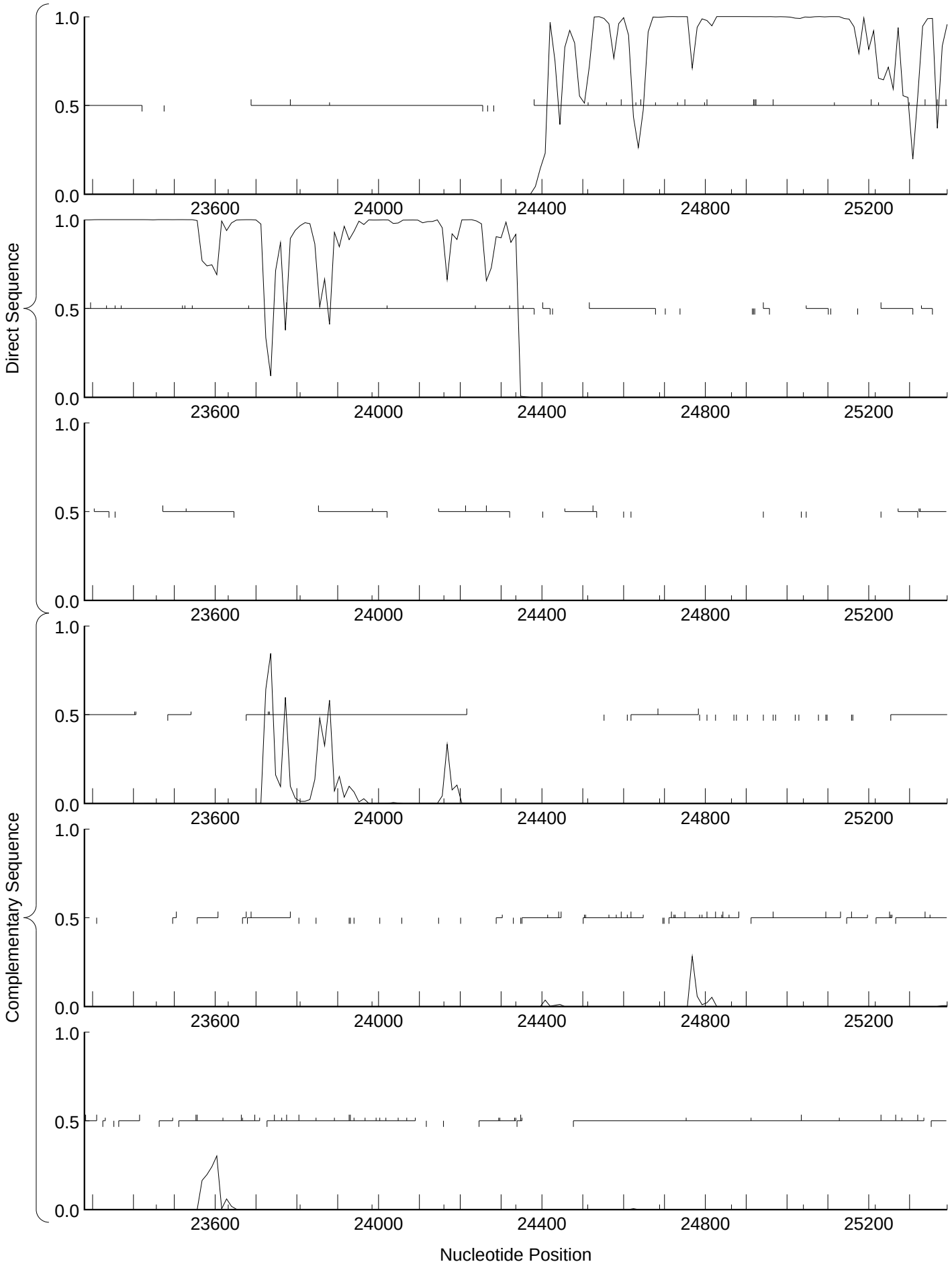
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 11/37



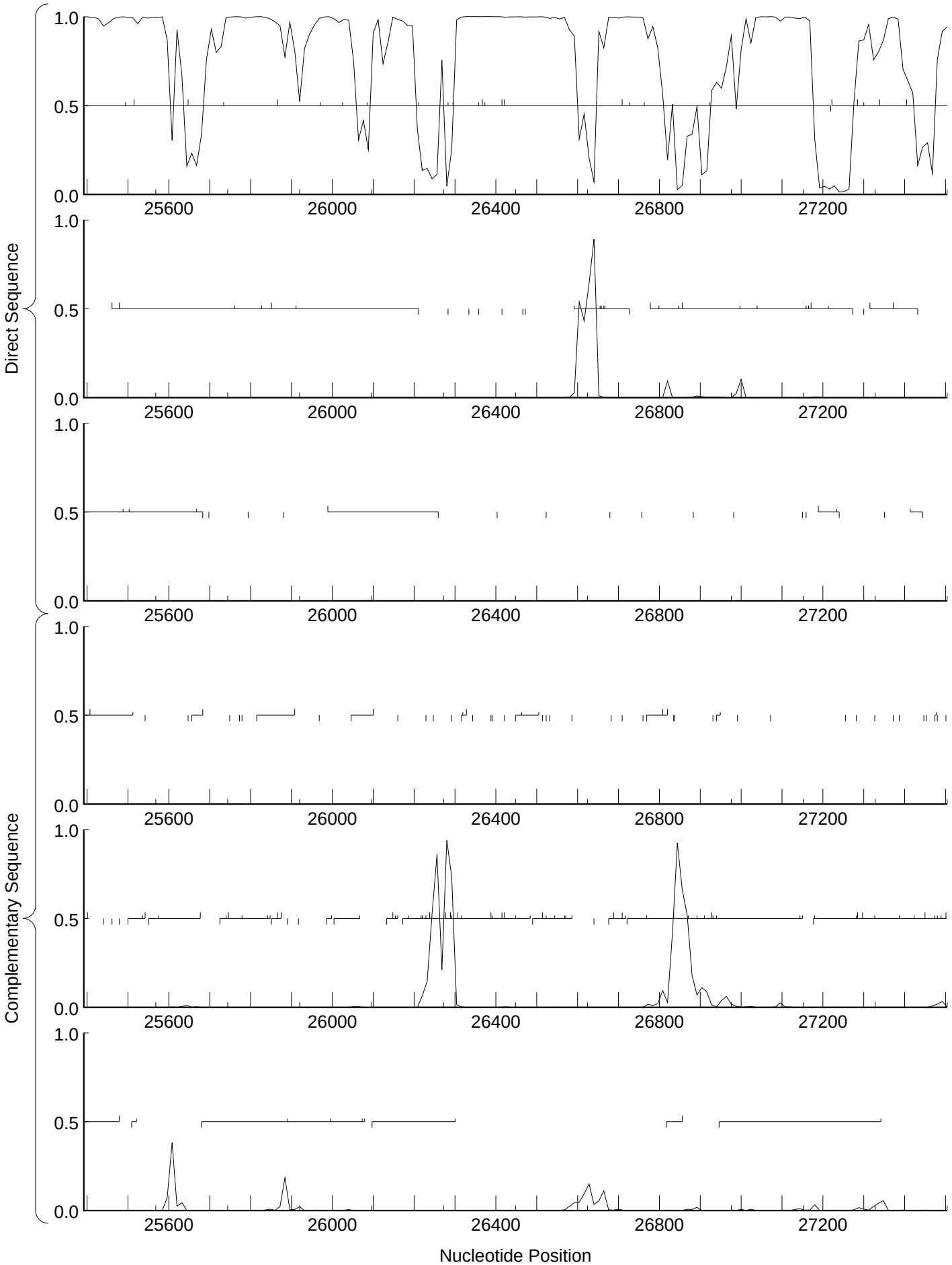
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 12/37



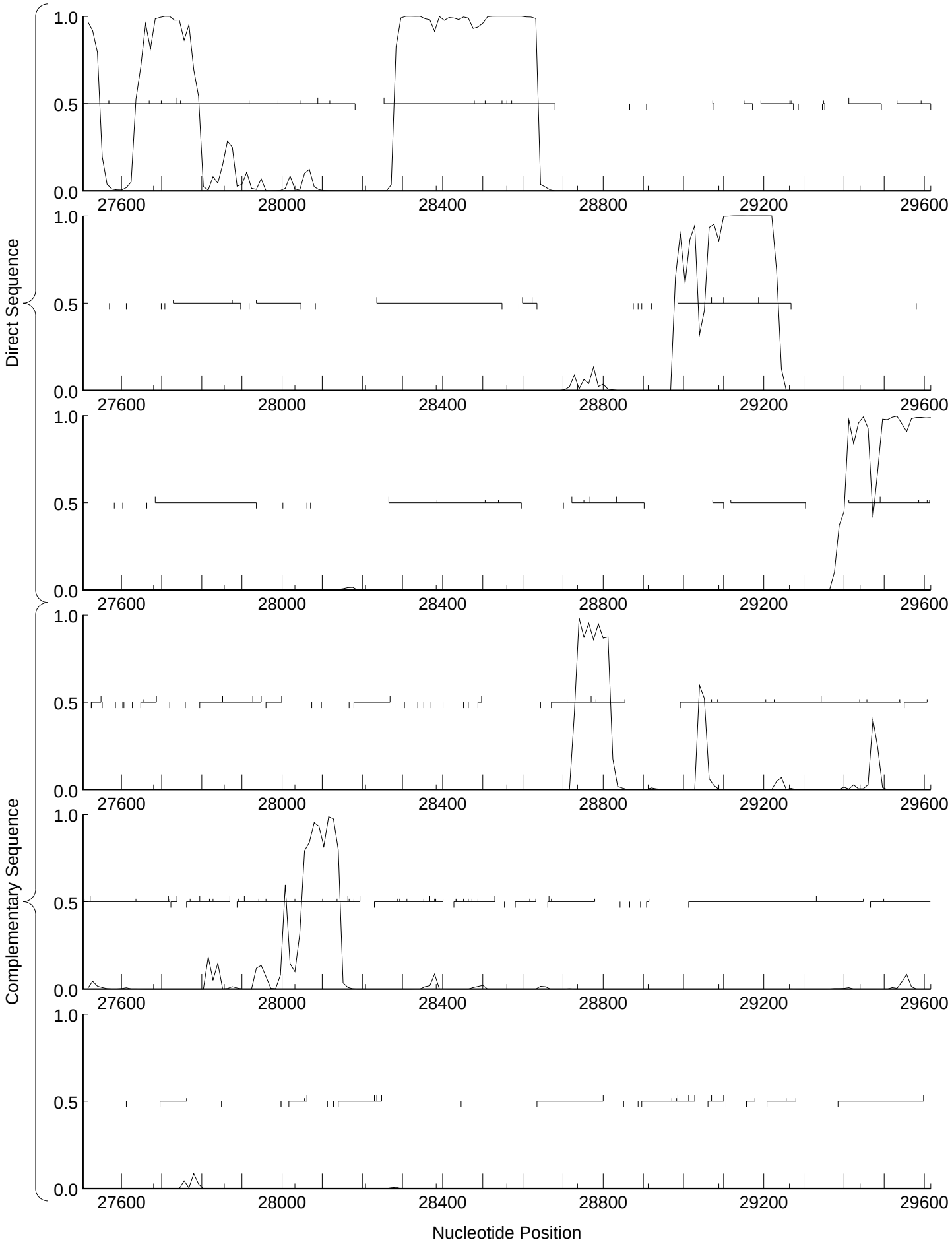
Lilizi complete sequence, 75360 bp including 9-base 3' overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 13/37



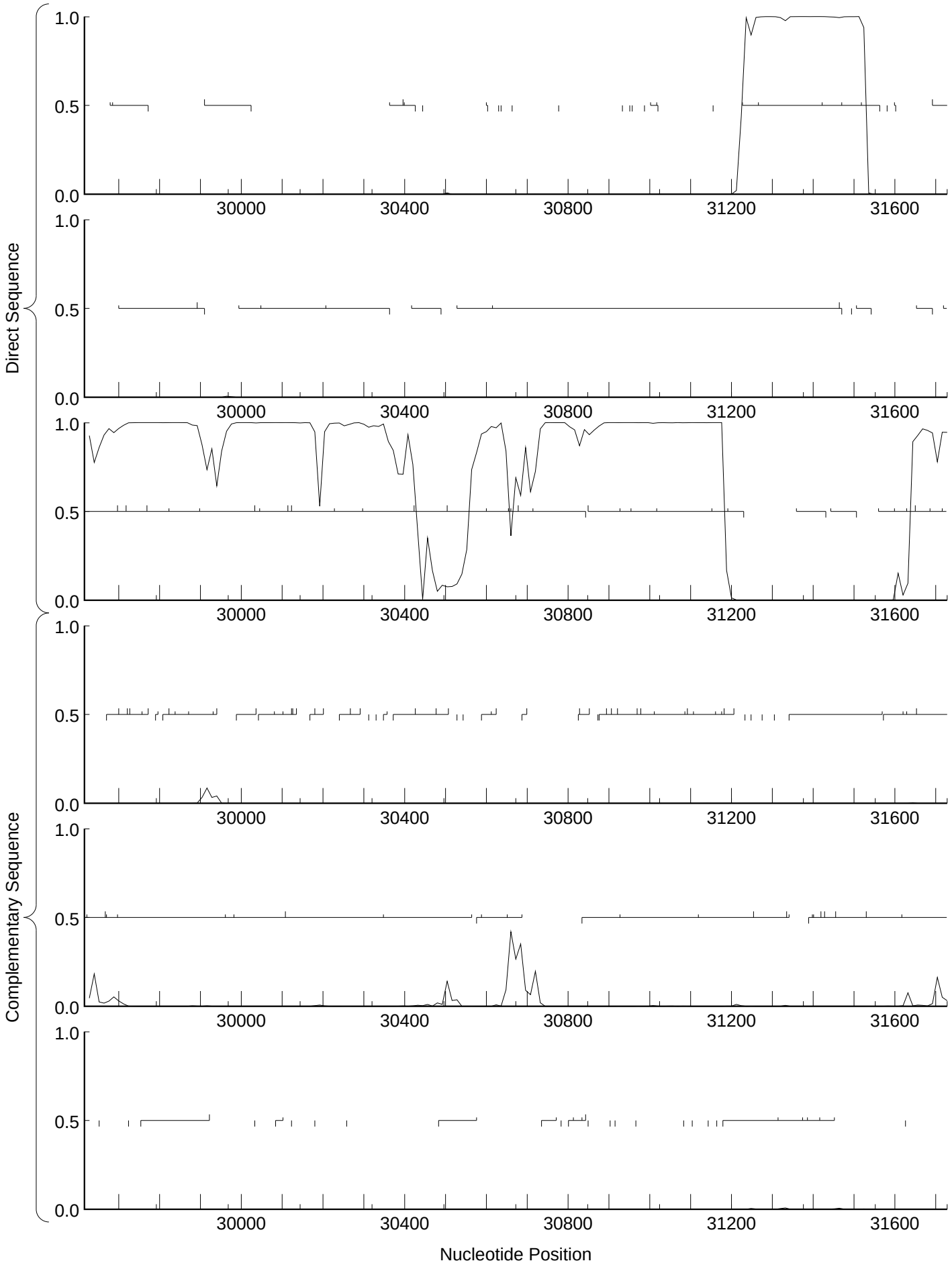
Lilizi complete sequence, 75360 bp including 9-base 3' overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 14/37



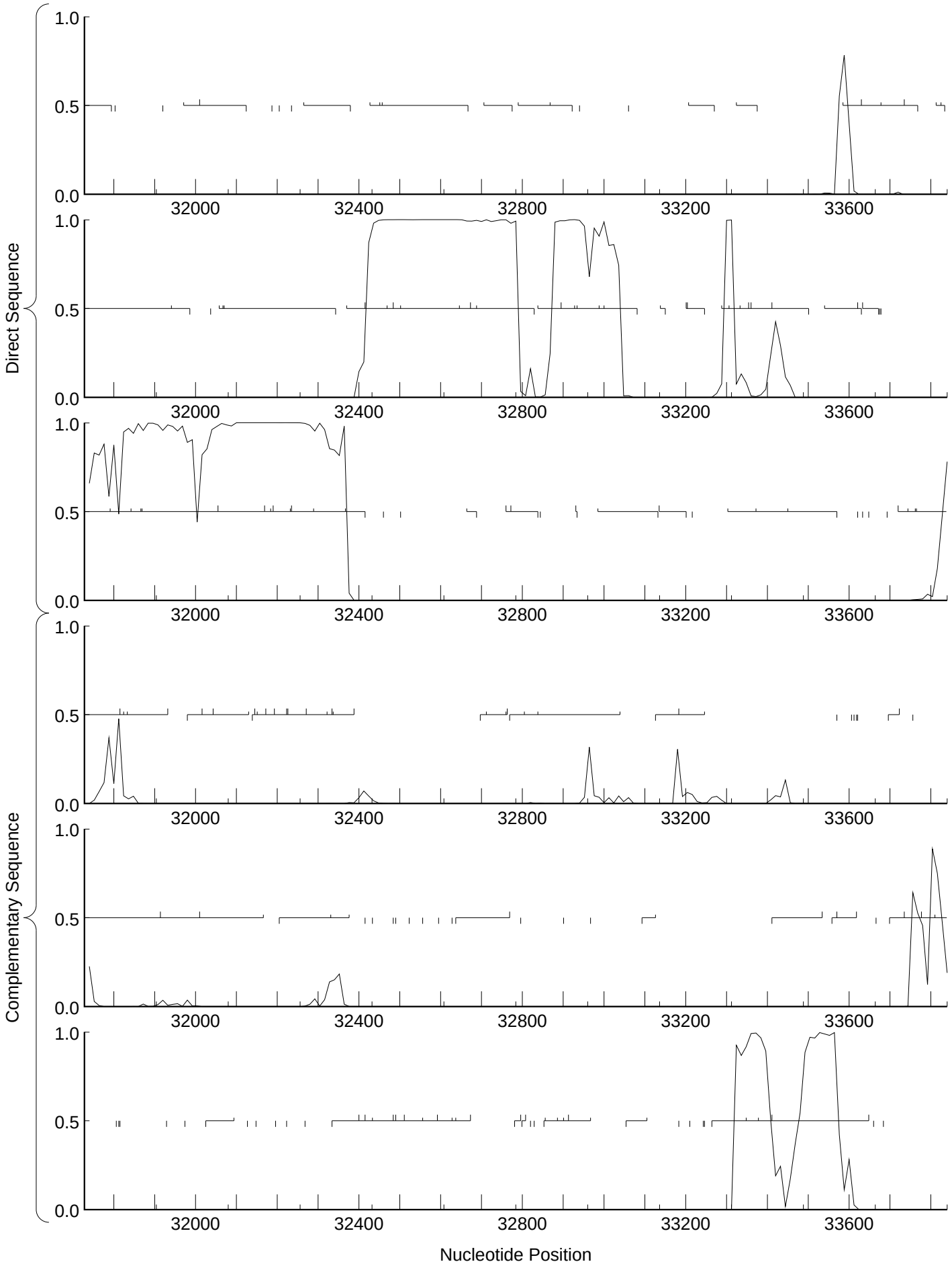
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 15/37



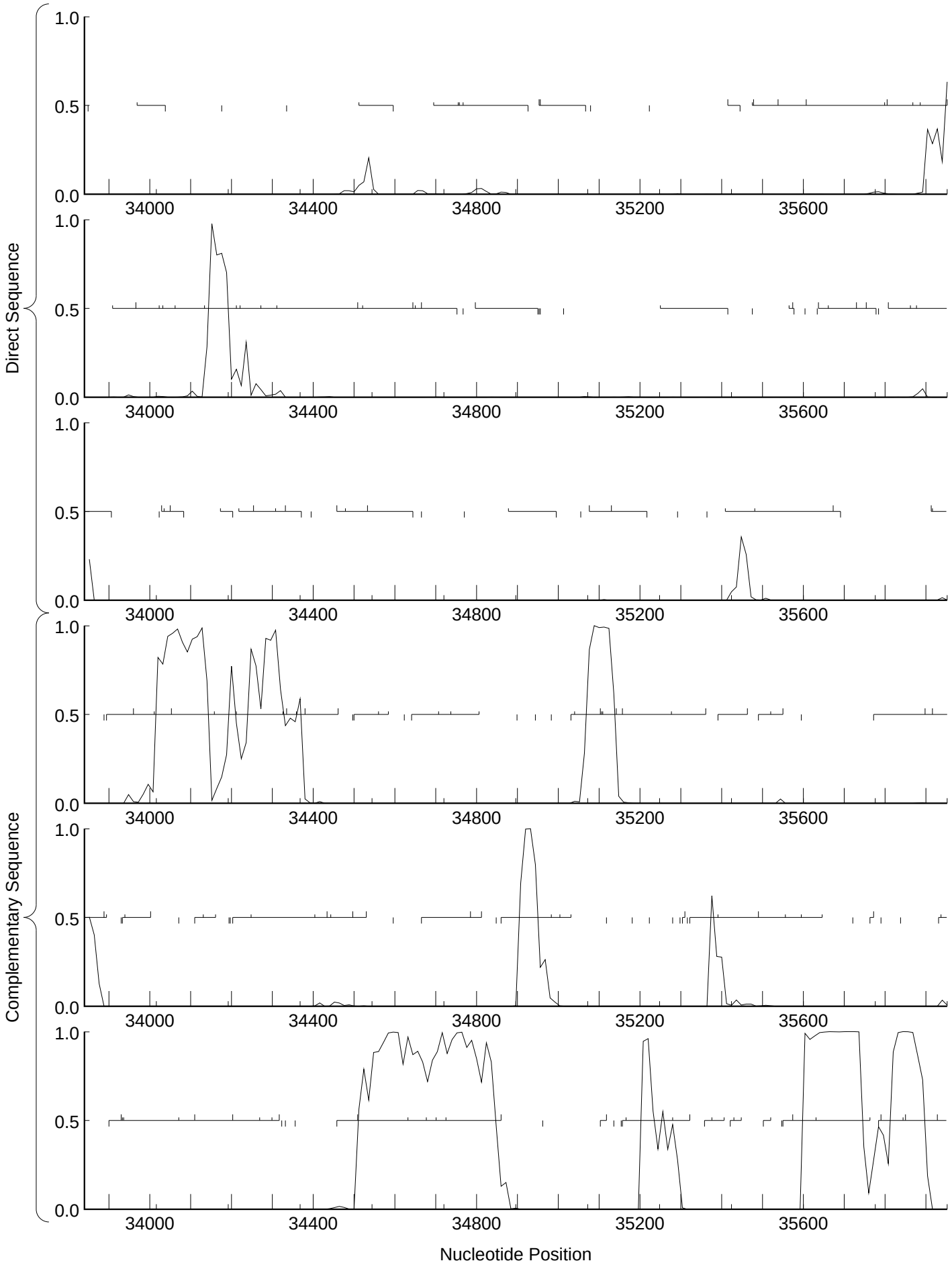
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 16/37



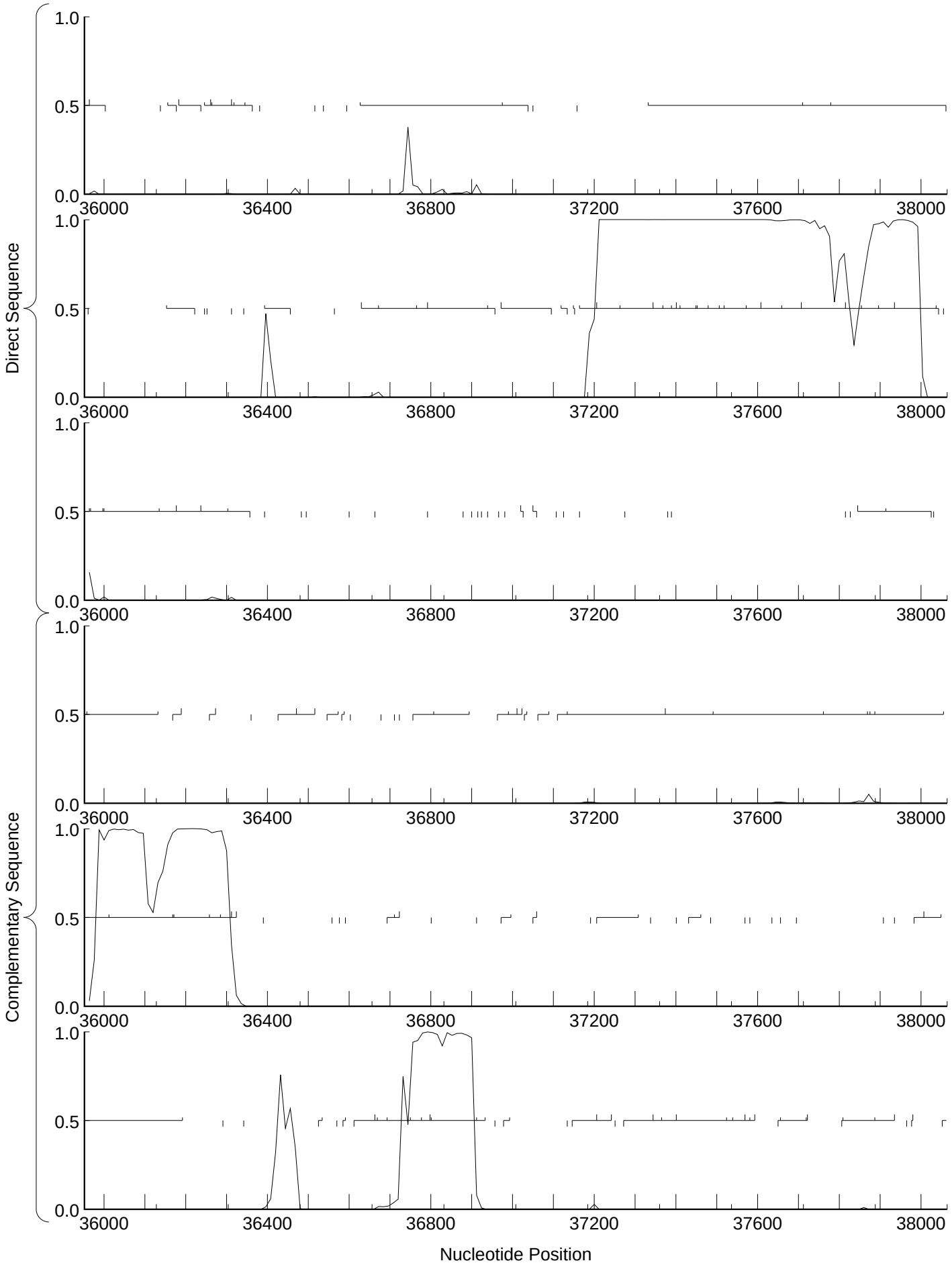
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 17/37



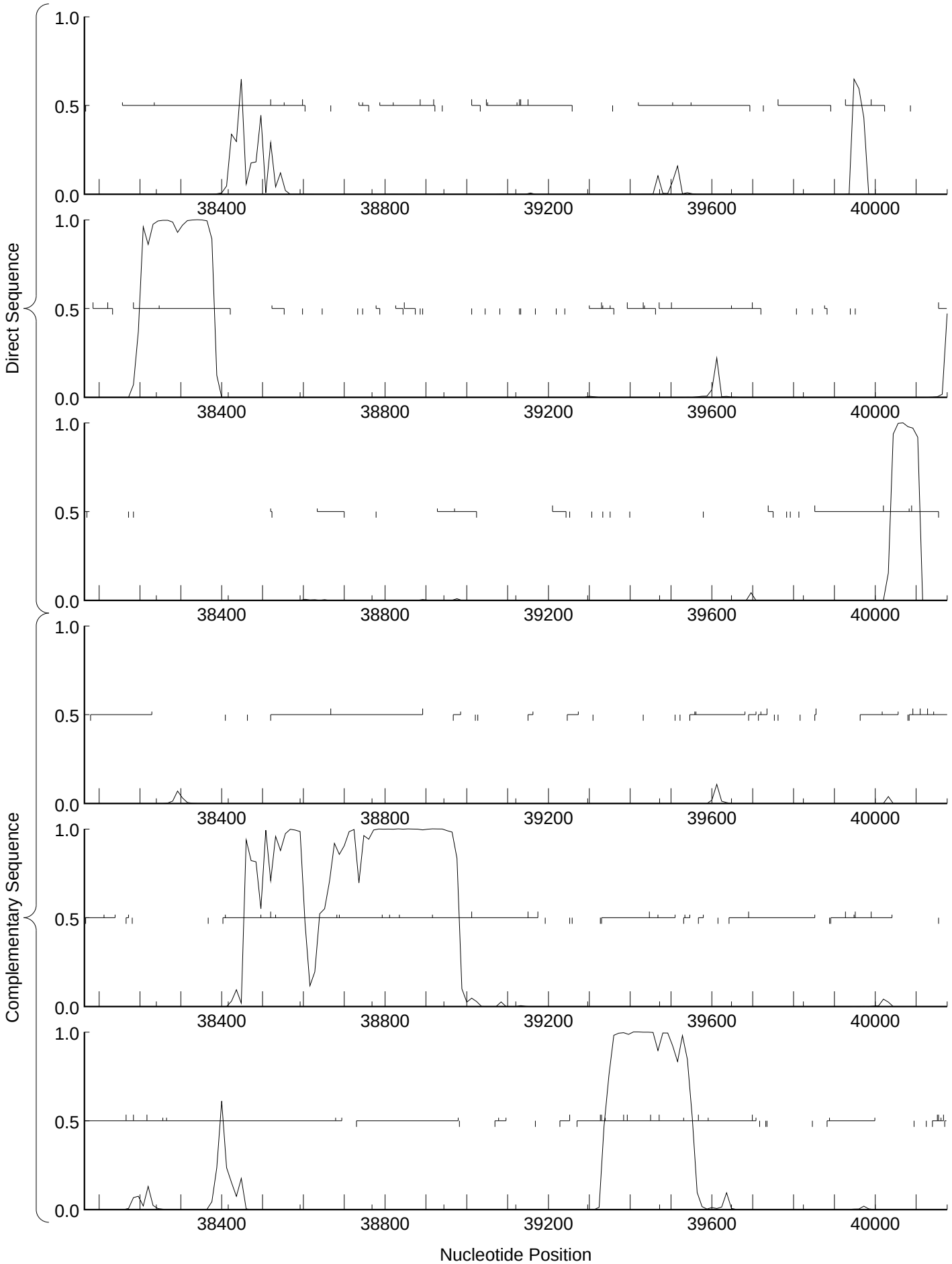
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 18/37



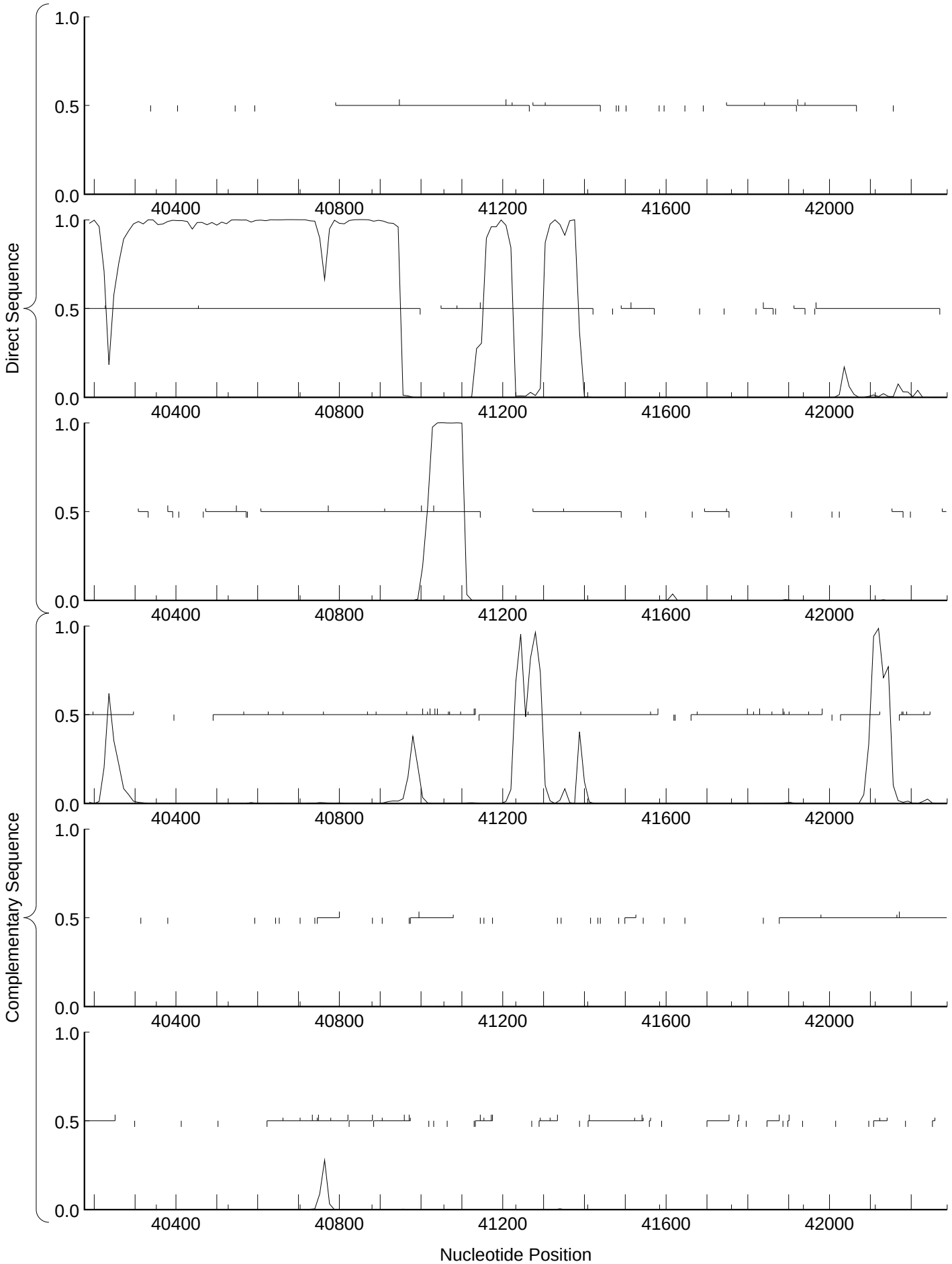
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 19/37



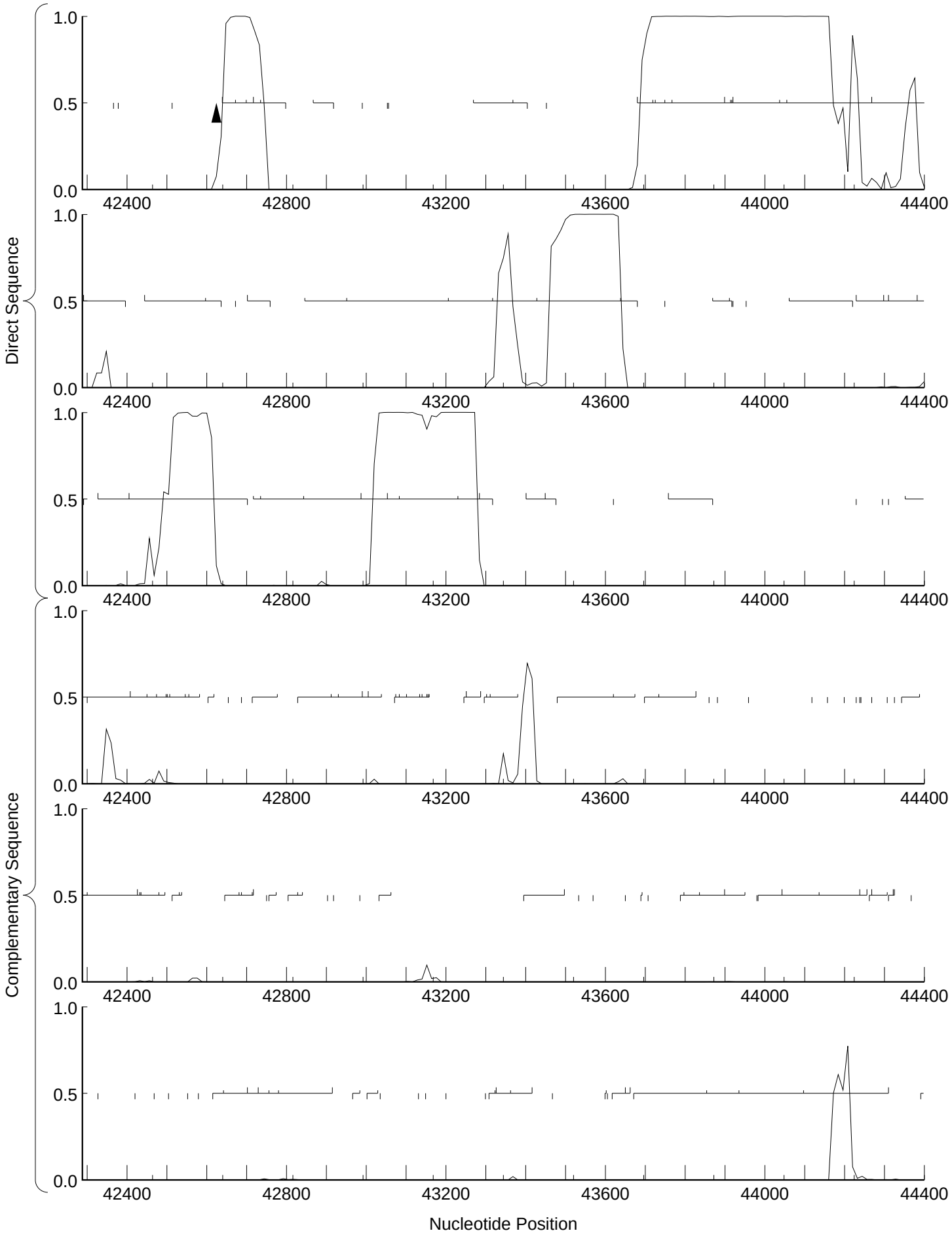
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 20/37



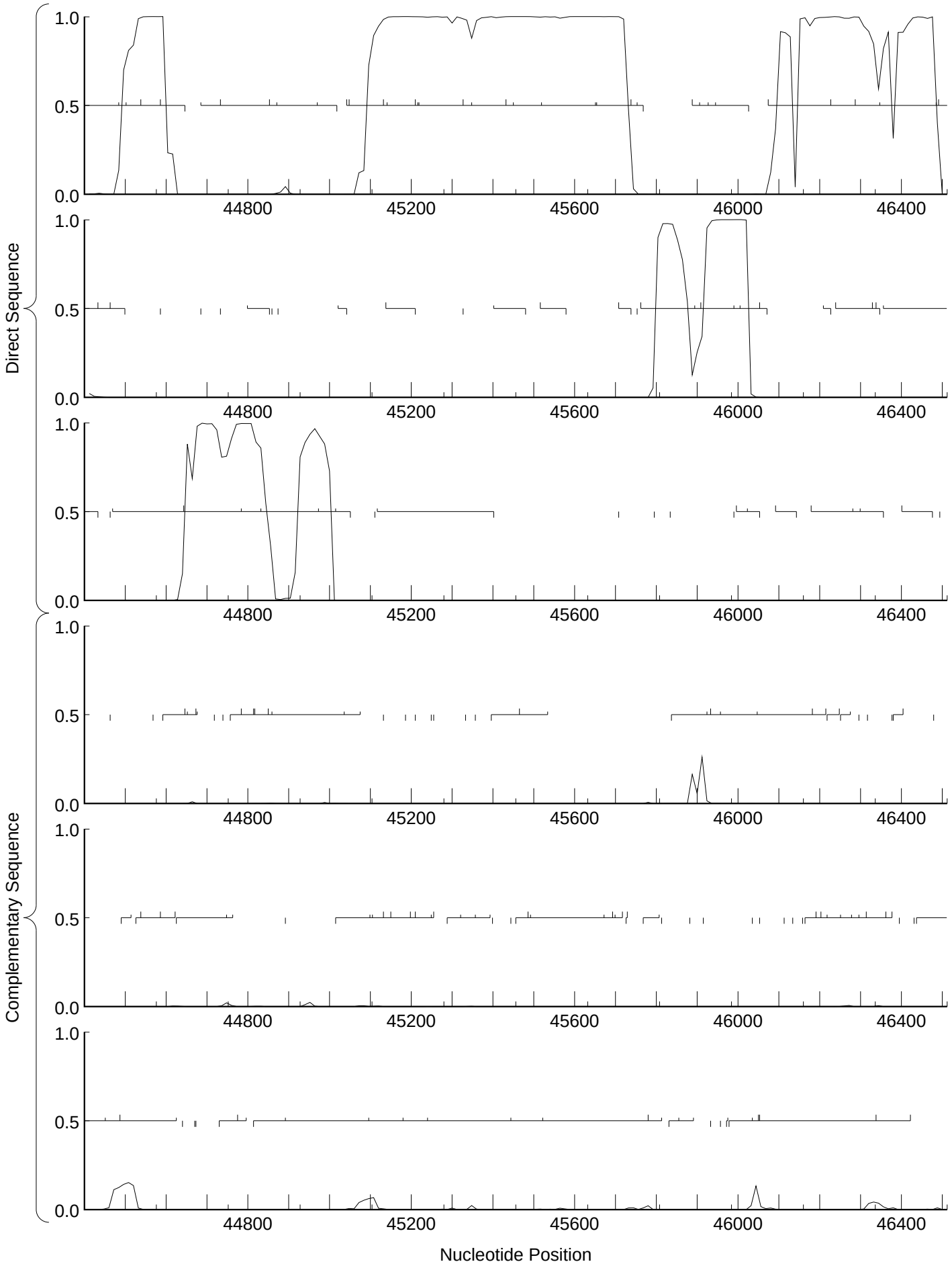
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 21/37



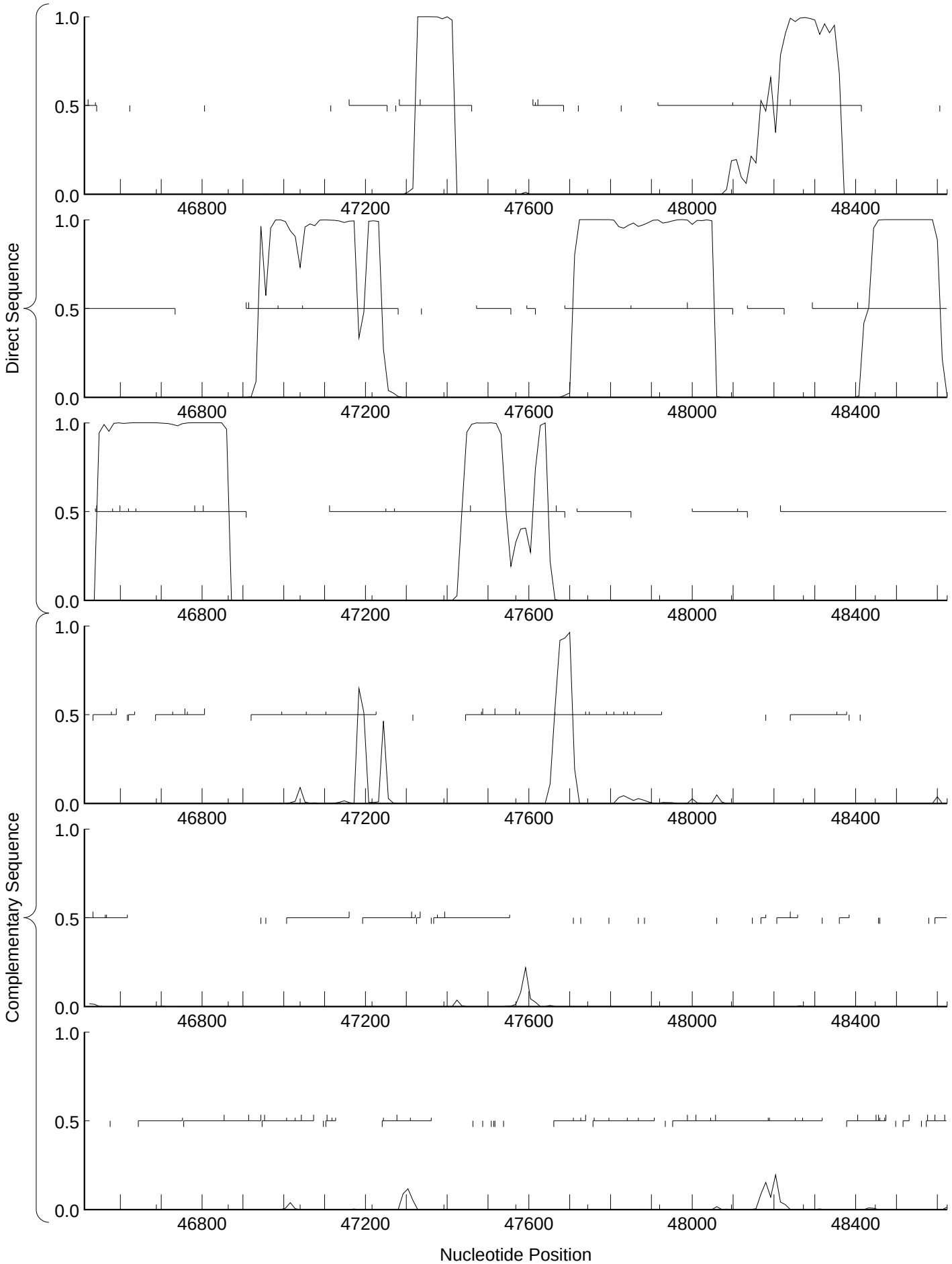
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 22/37



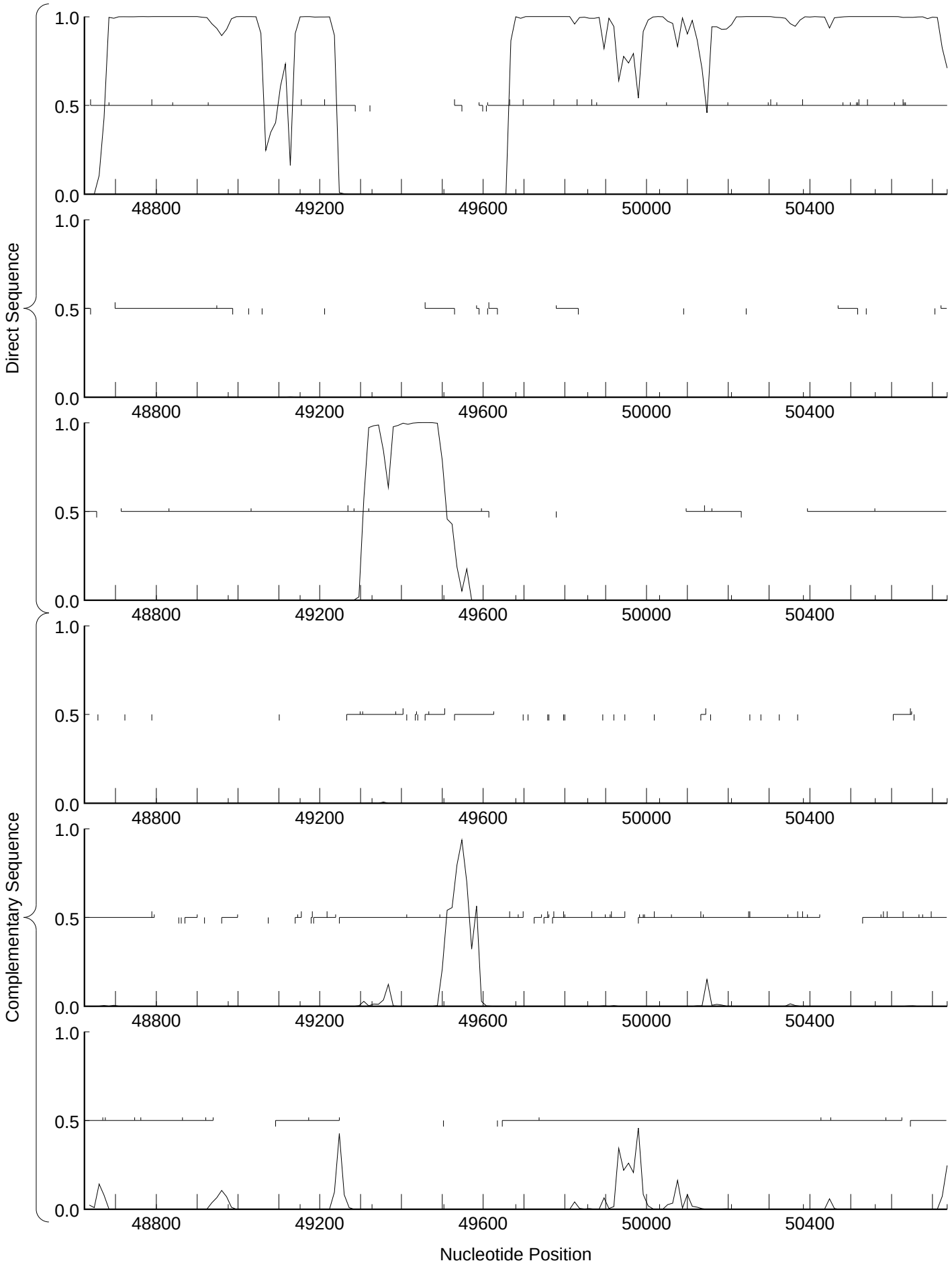
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 23/37



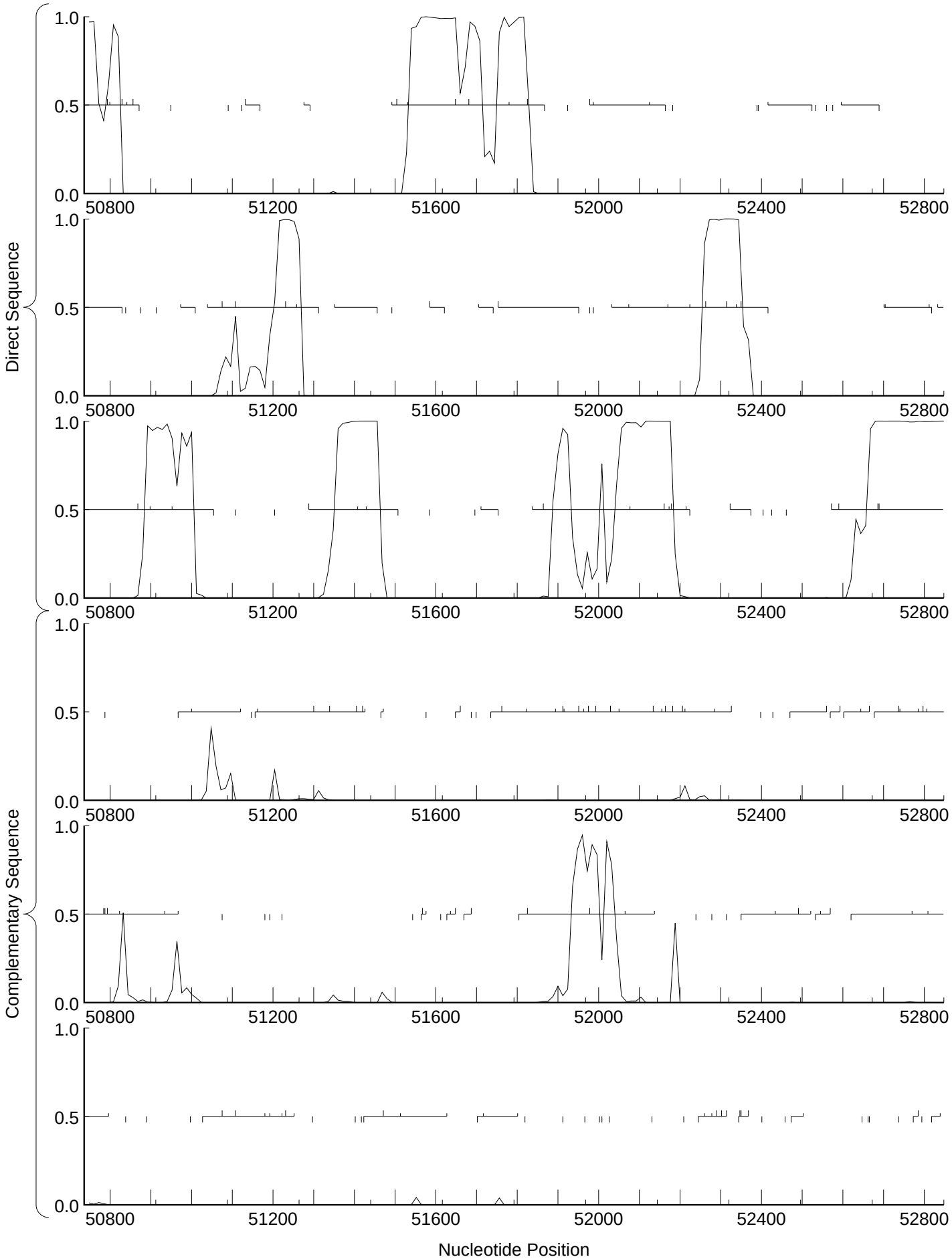
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 24/37



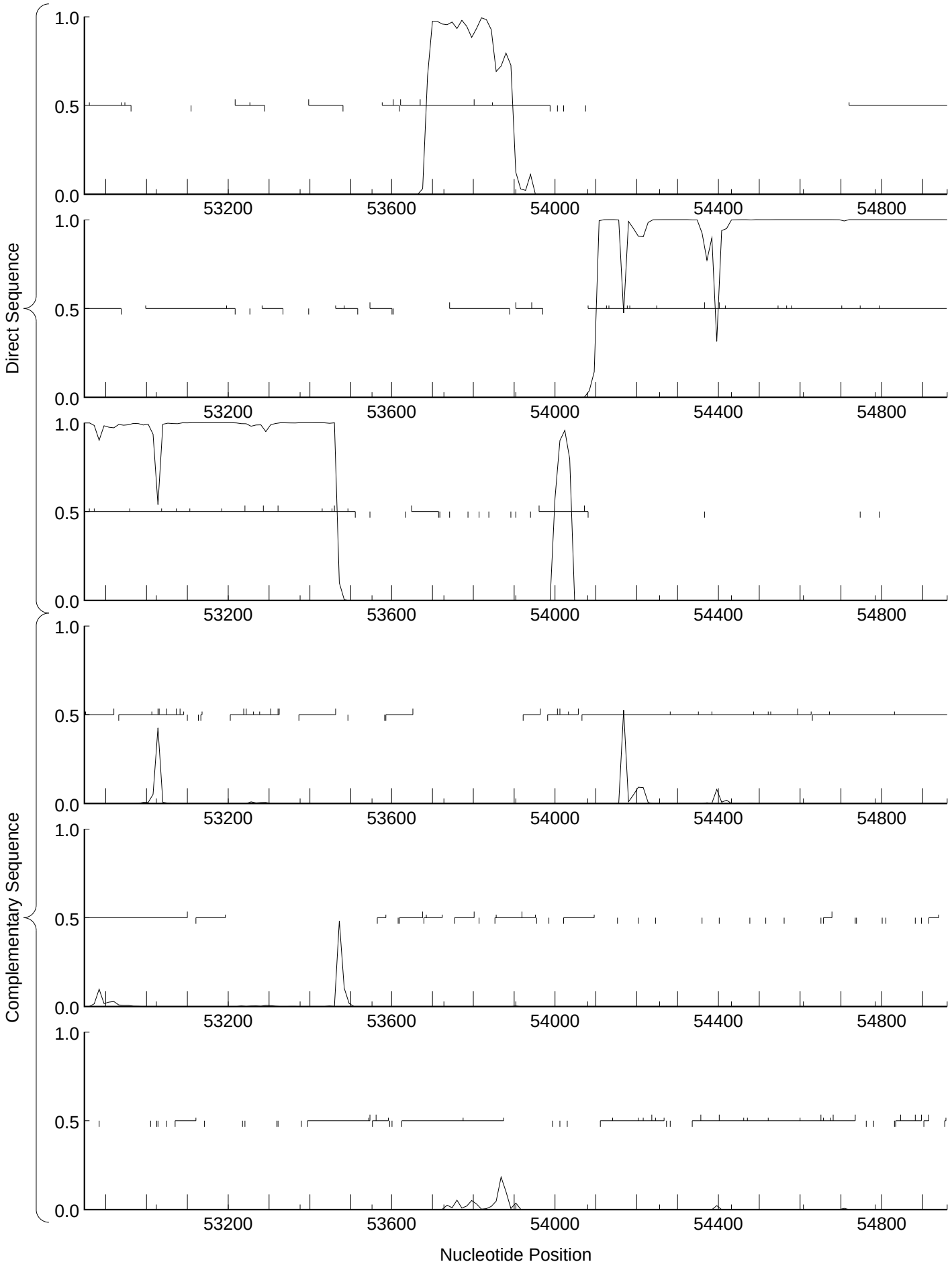
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 25/37



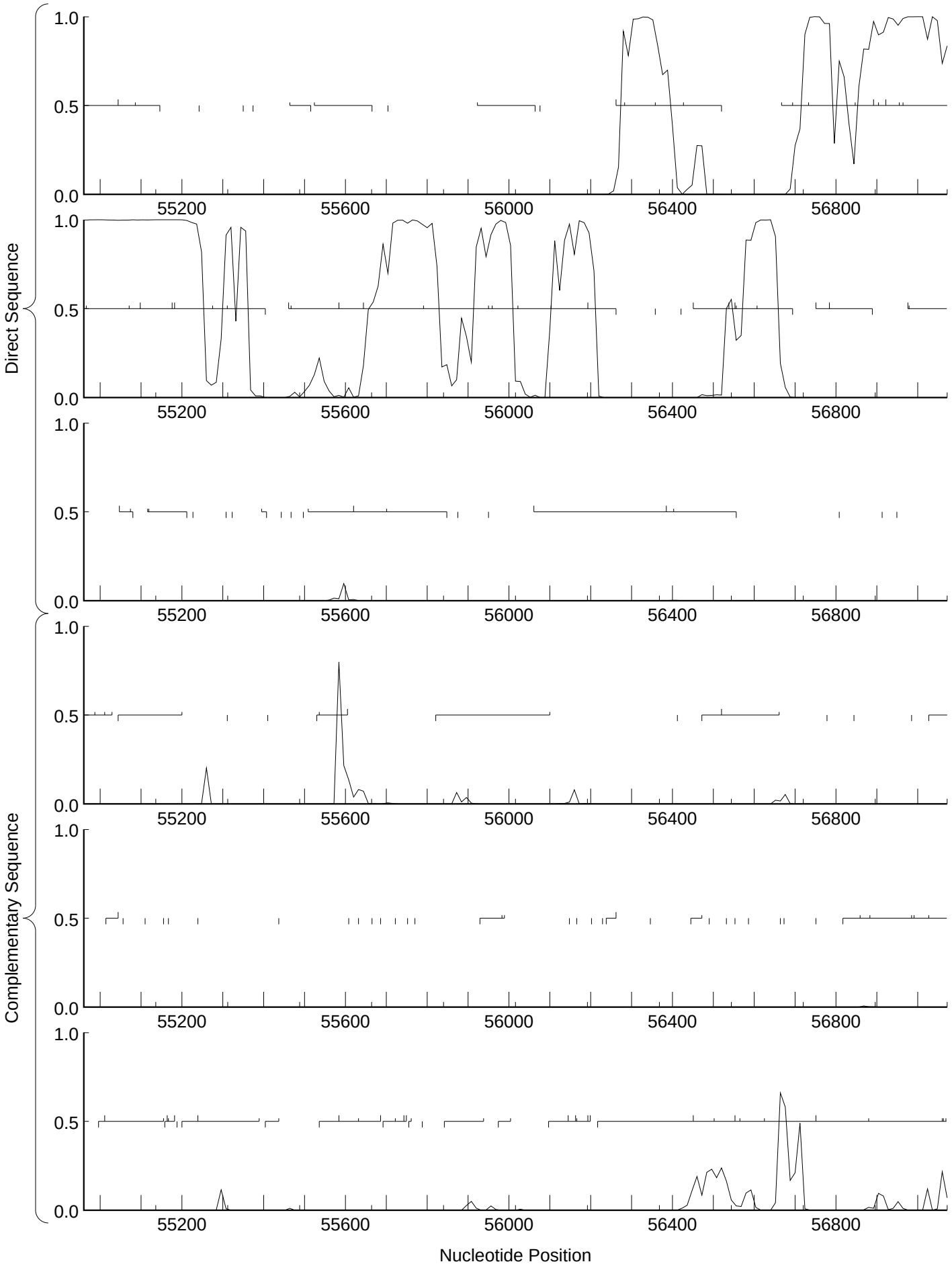
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 26/37



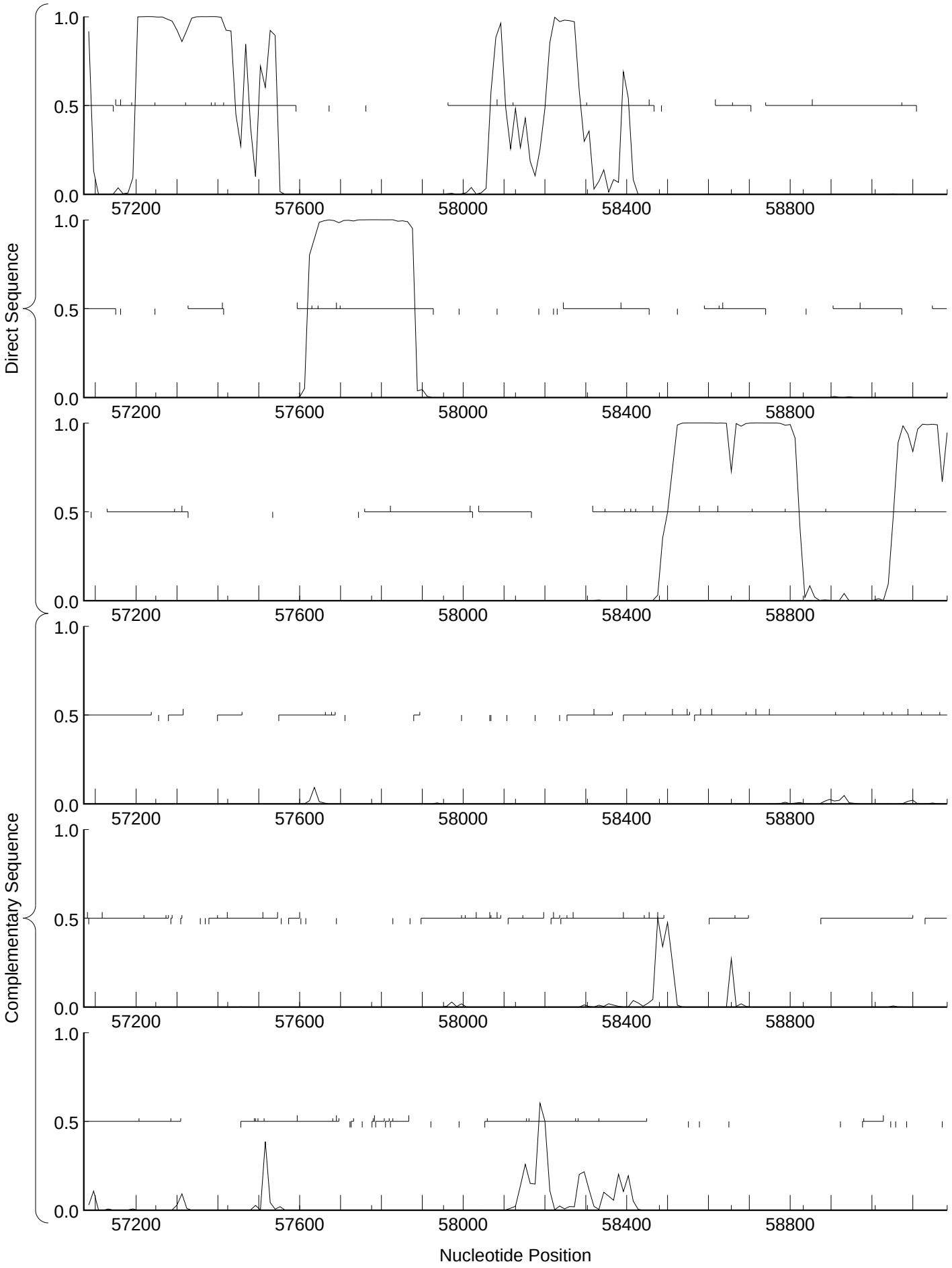
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 27/37



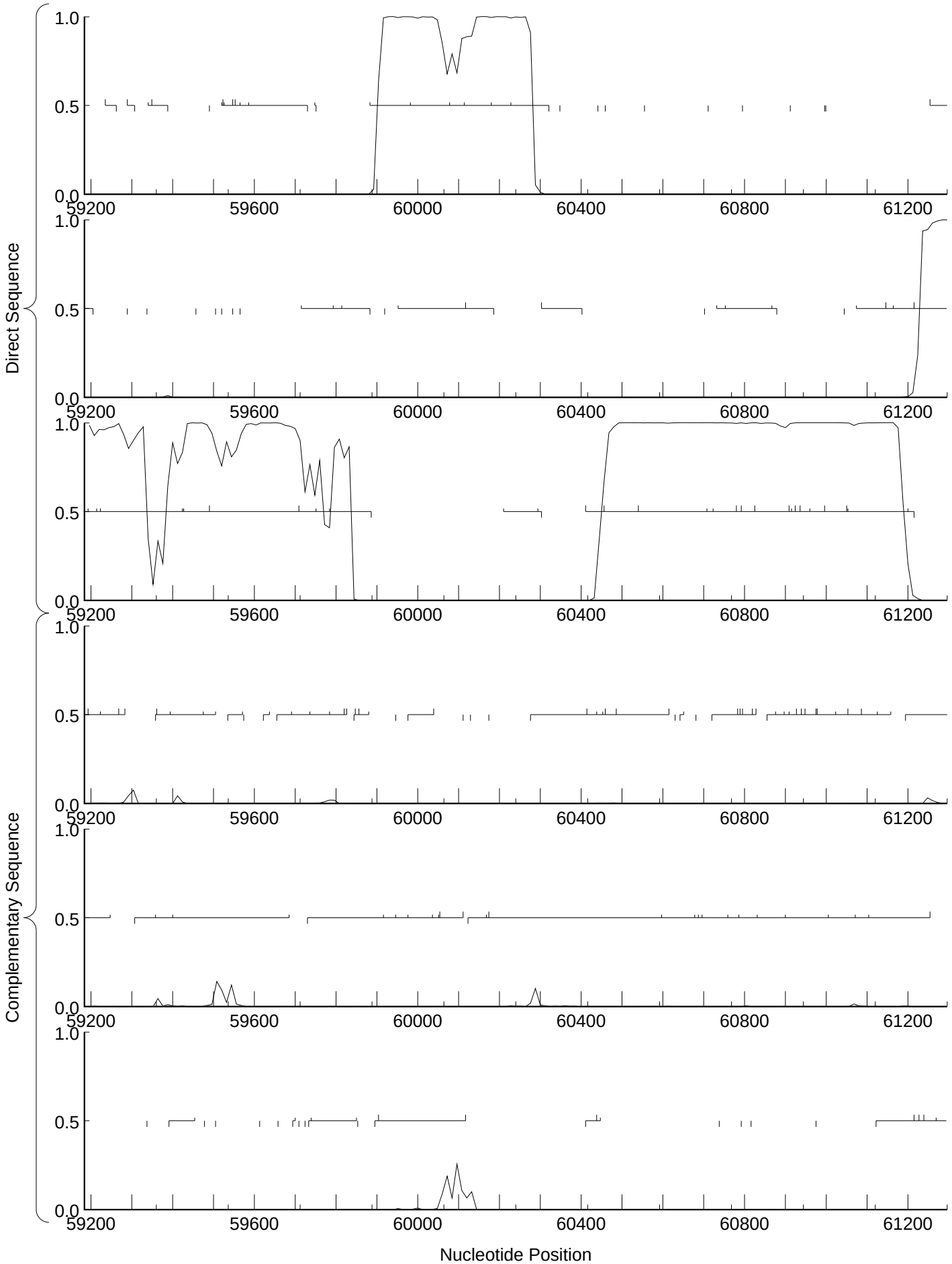
Lilizi complete sequence, 75360 bp including 9-base 3' overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 28/37



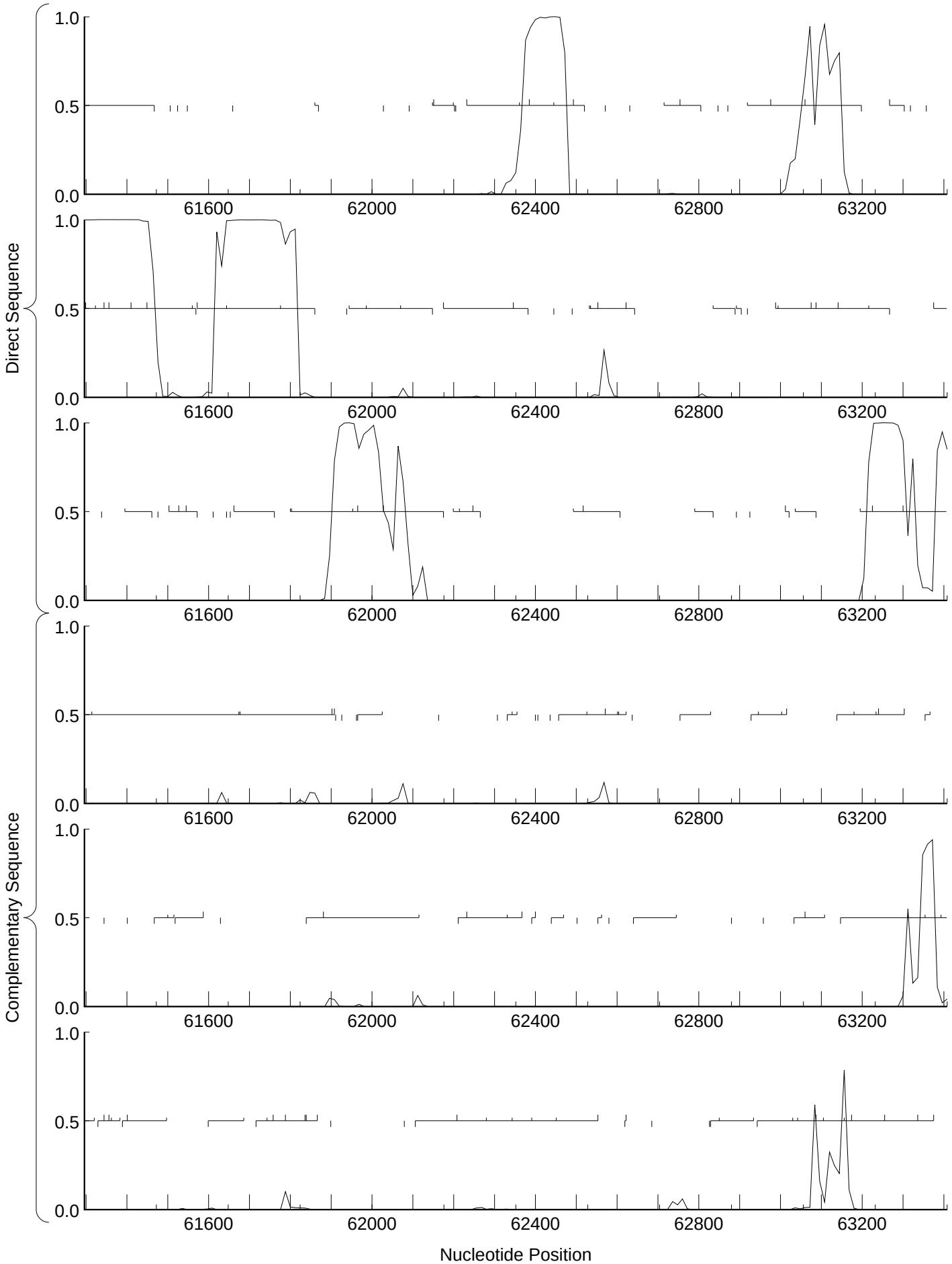
Lilizi complete sequence, 75360 bp including 9-base 3' overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 29/37



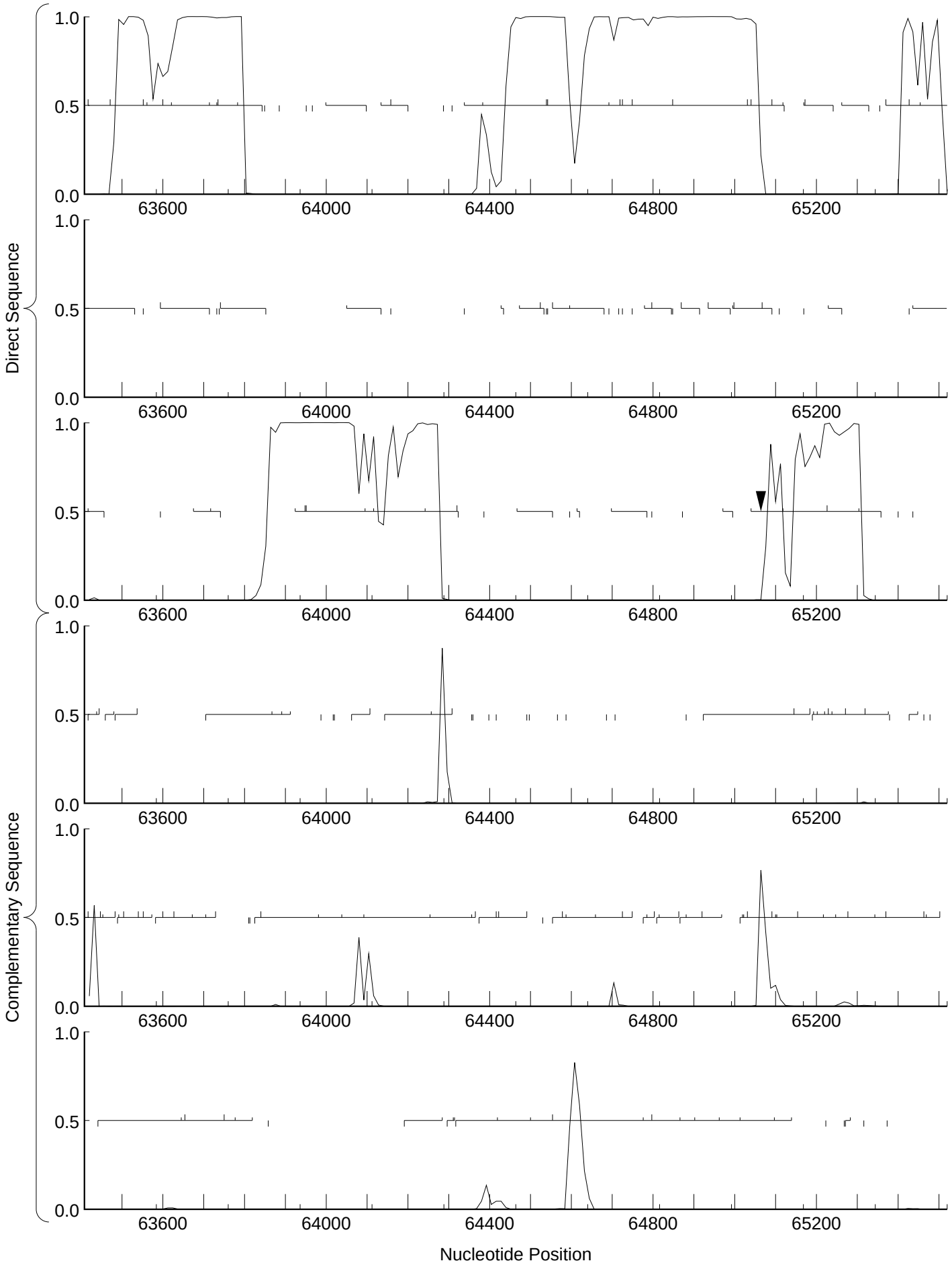
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 30/37



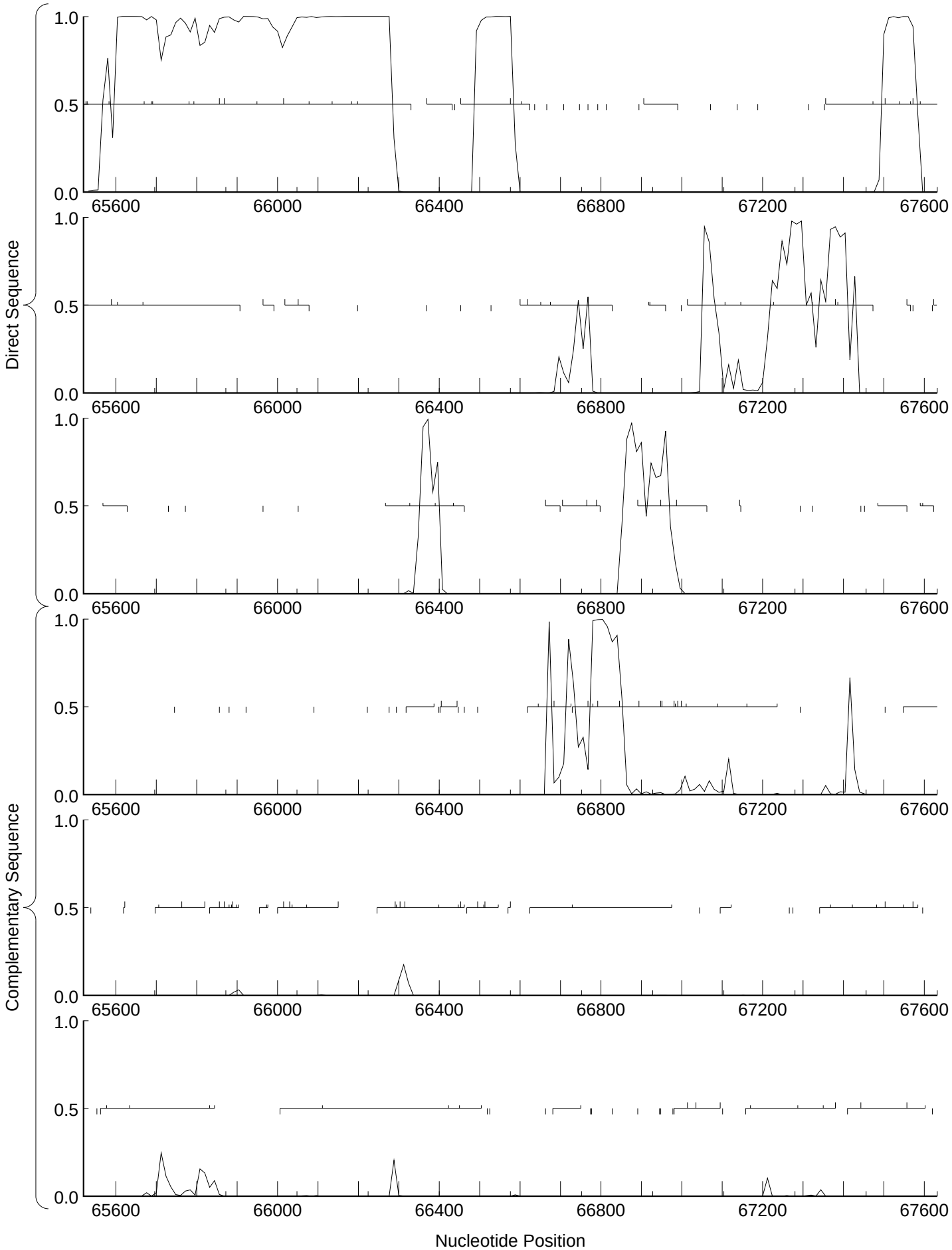
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 31/37



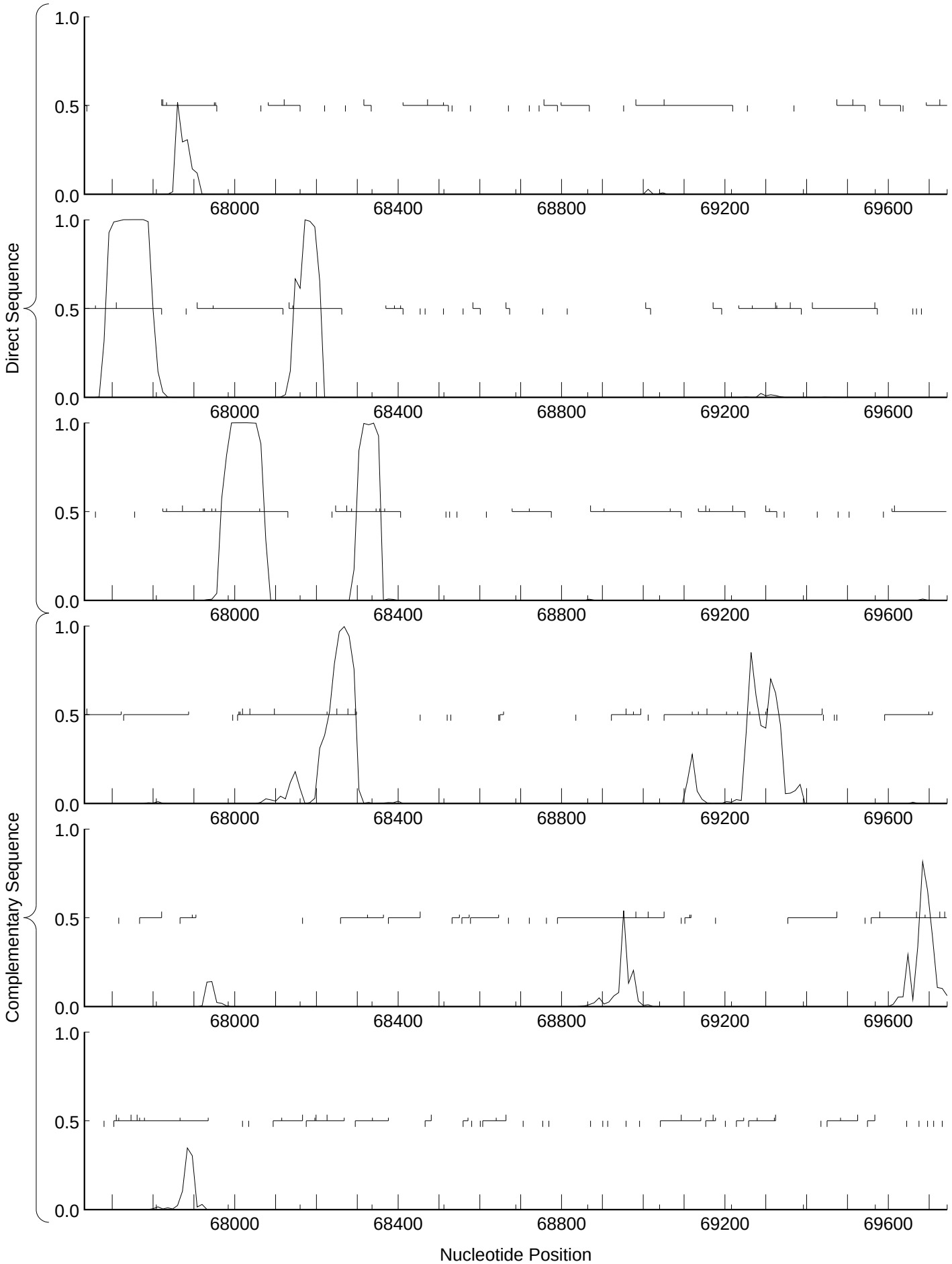
Lilizi complete sequence, 75360 bp including 9-base 3' overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 32/37



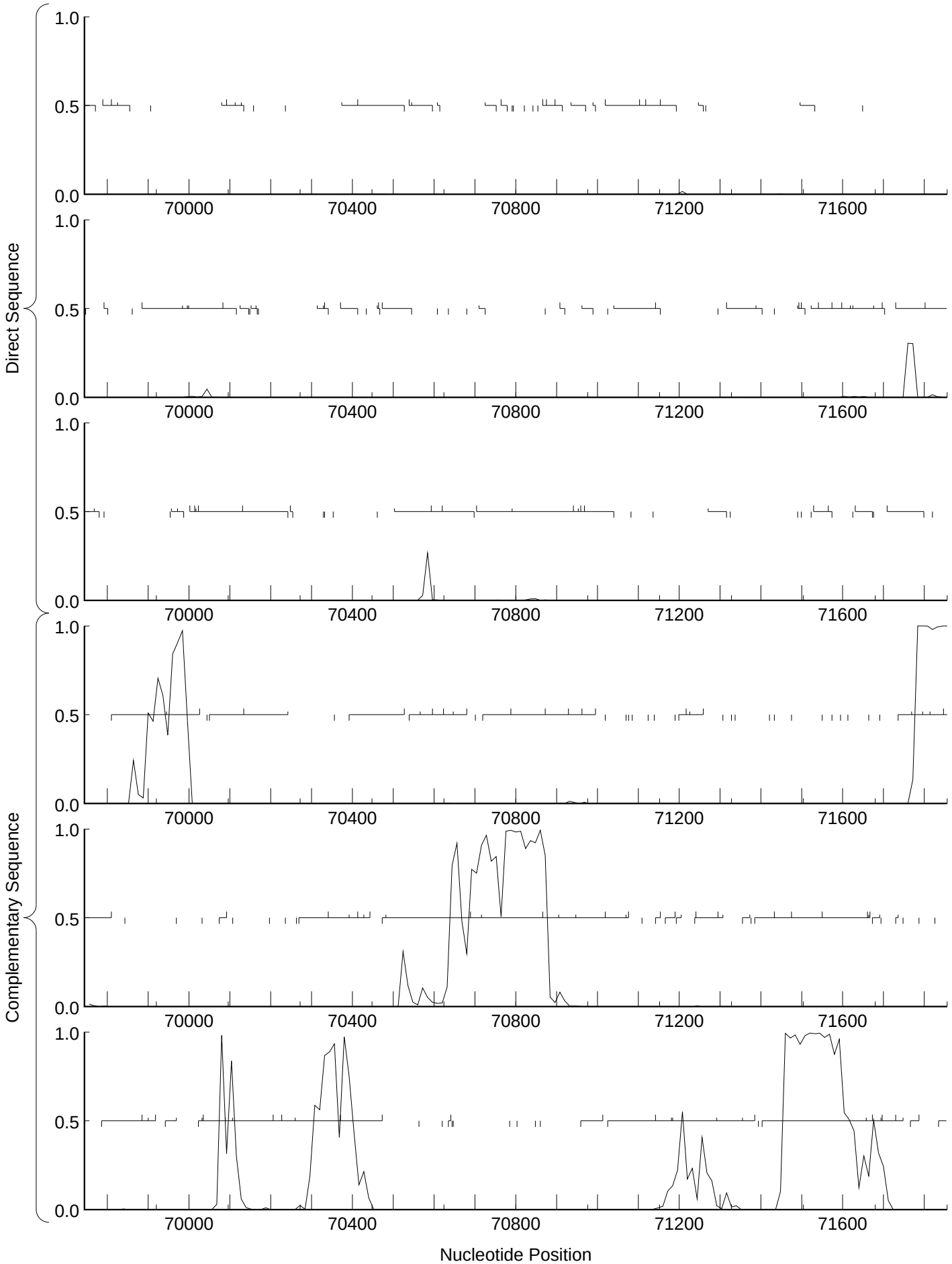
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 33/37



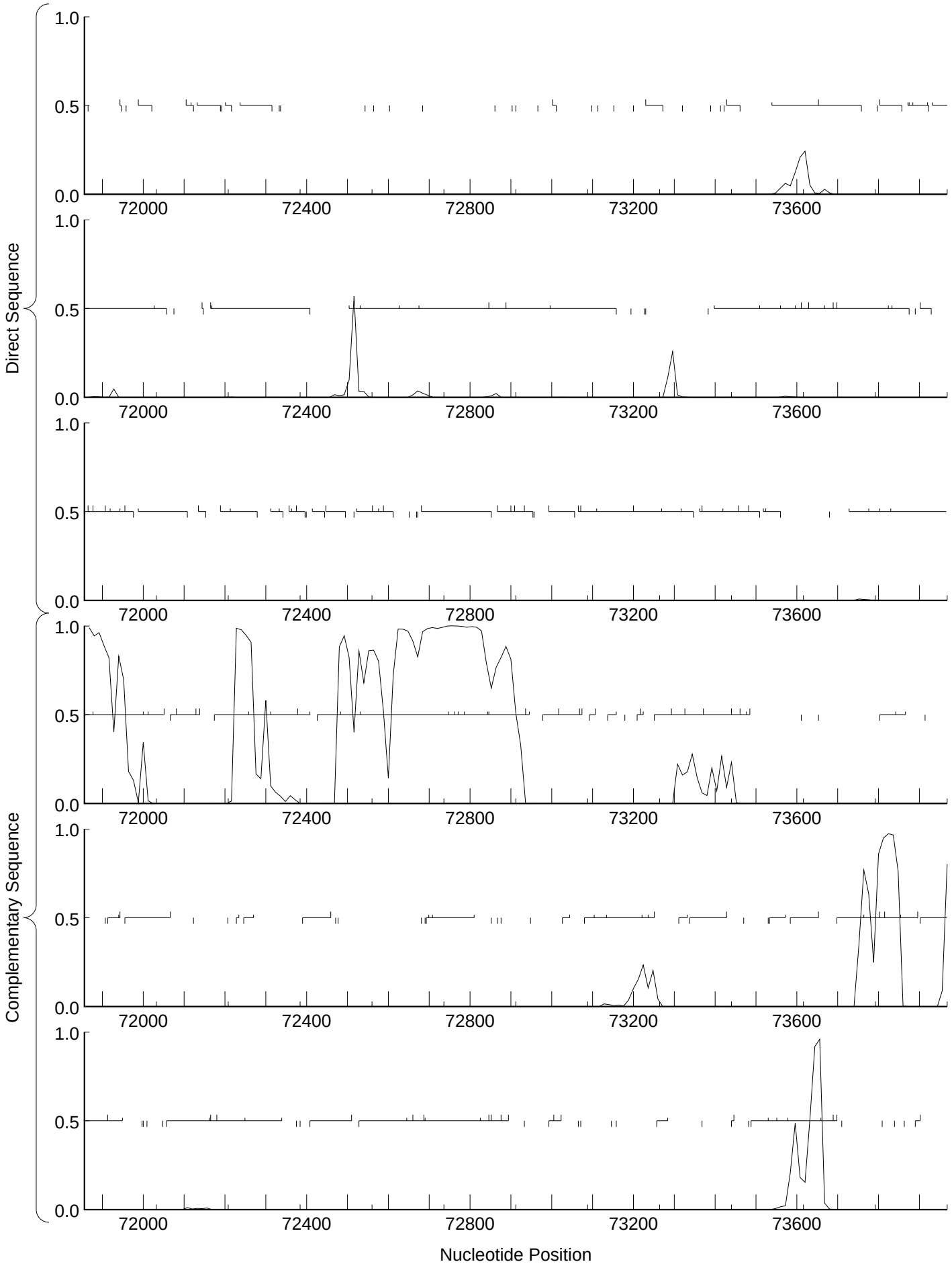
Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 34/37



Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 35/37



Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 36/37



Lilizi complete sequence, 75360 bp including 9-base 3’ overhang (CGCTTGTCA), Cluster E, Order 4, Window 96, Step 12, 37/37

