GeneMark

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

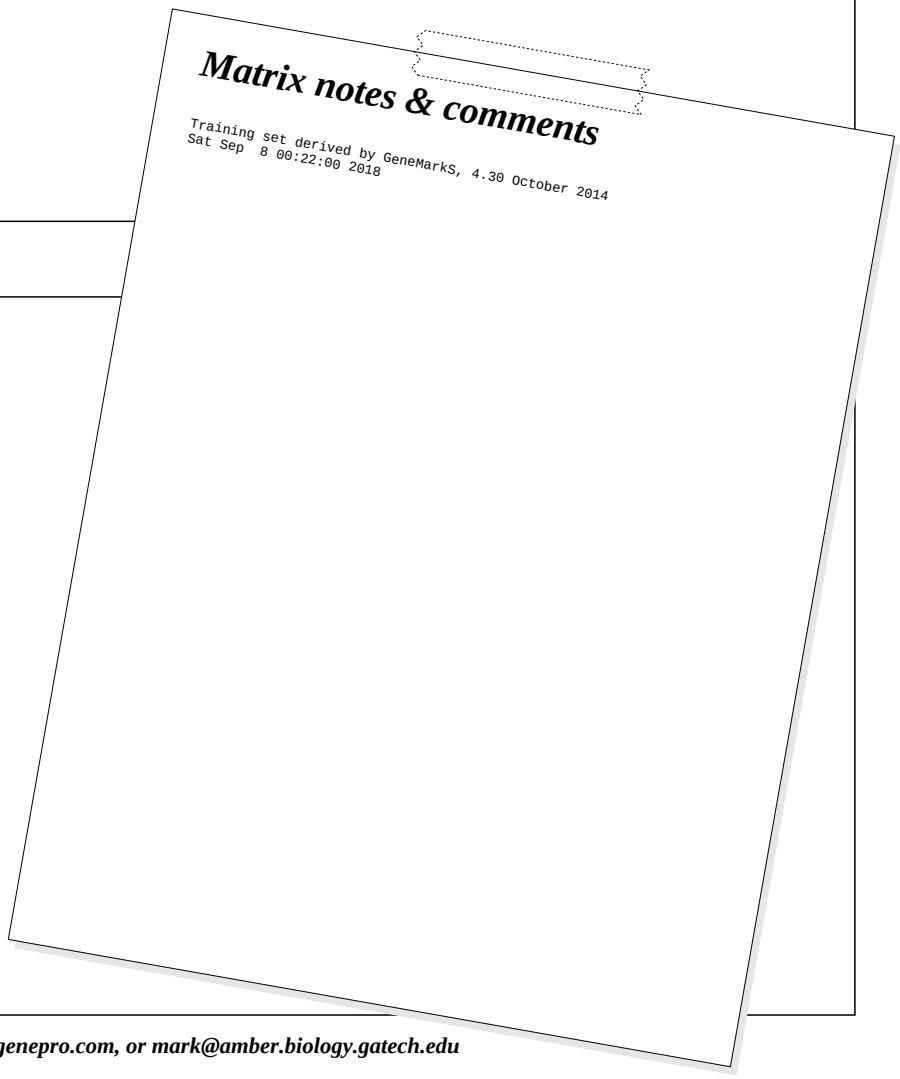
PROGRAM INFORMATION

```
Sequence         :  Arthrobacter phage Riverdale complete sequence, 43391 bp 13-base 3′ overhang (GGTAACCGTGATA
Analysis Date    :  3/29/24 at 4:35:54
Pages            :  22
Sequence Length  :  43391 bp
GC Content       :   60.99%

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

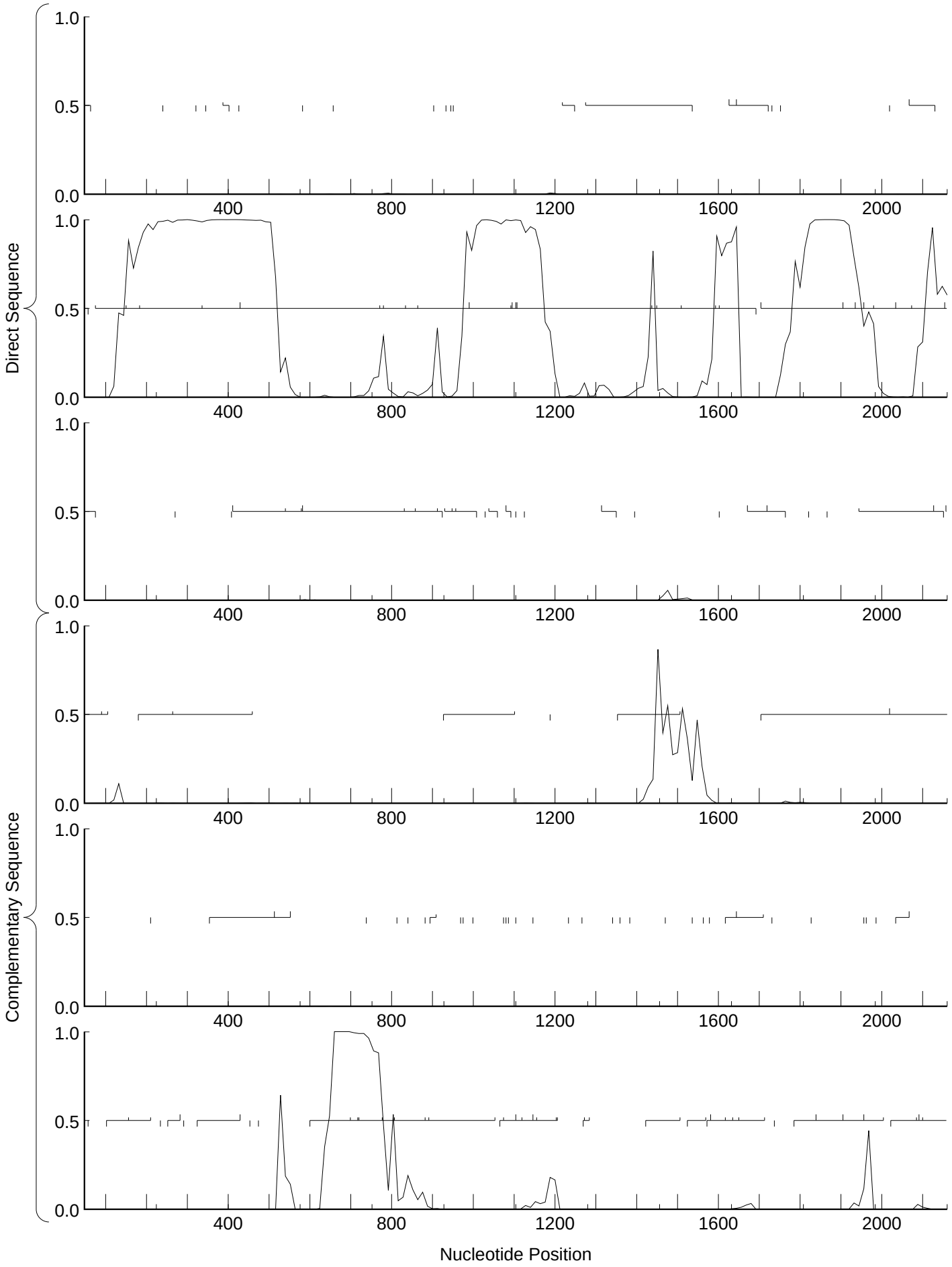
```
Matrix     :  -
Author     :  -

Order      :  4

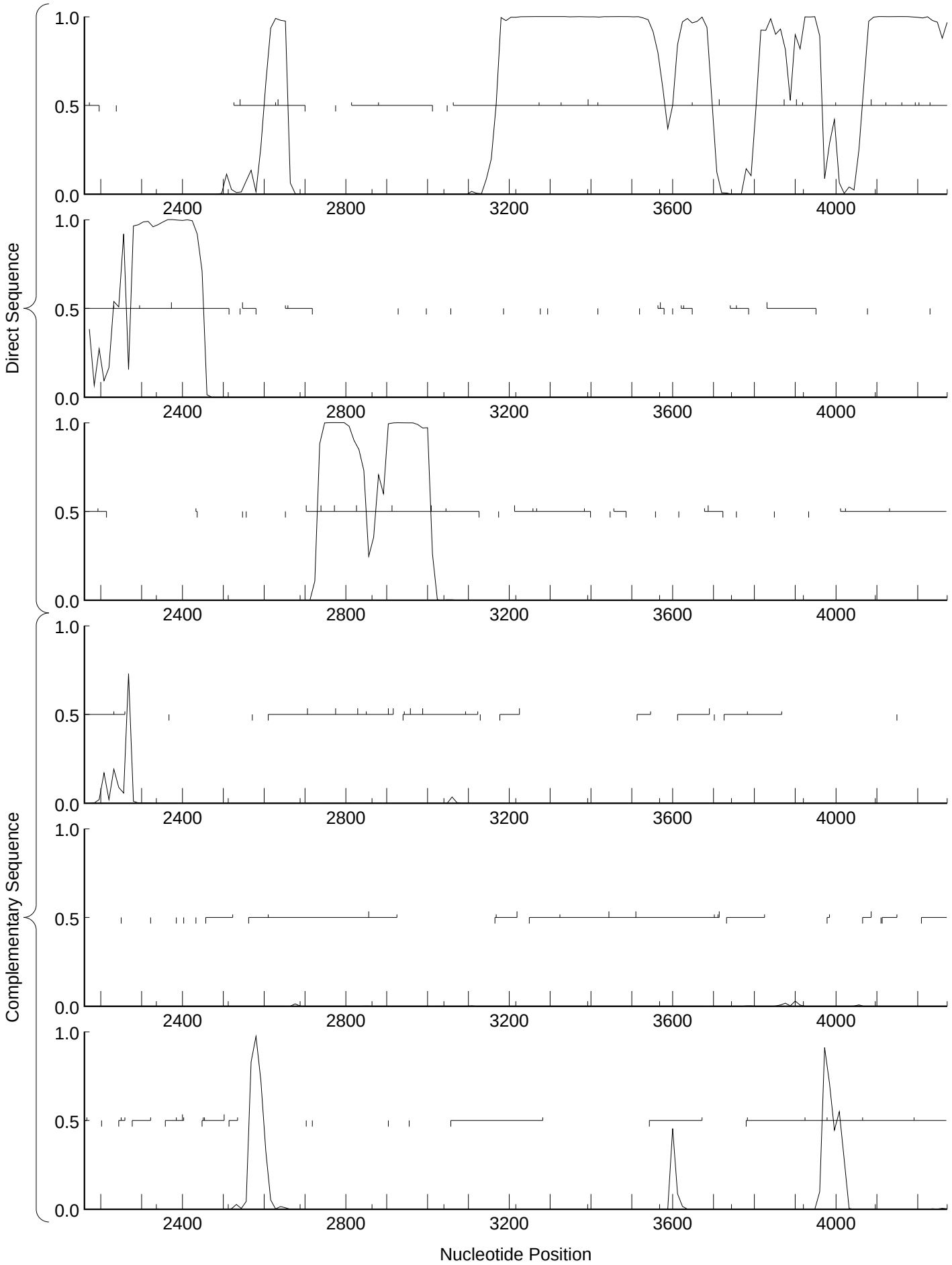

  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```

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

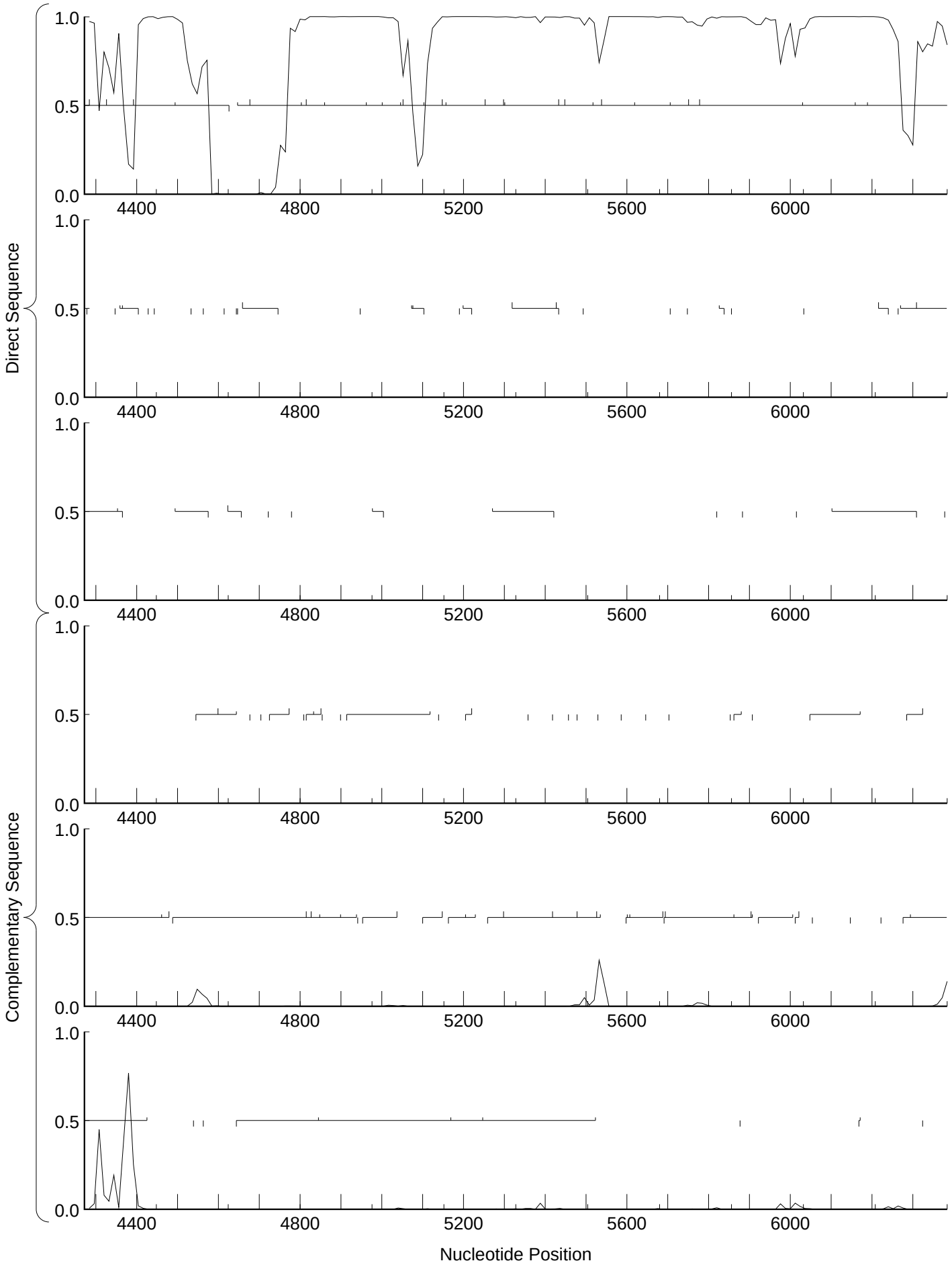
erdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 2/22



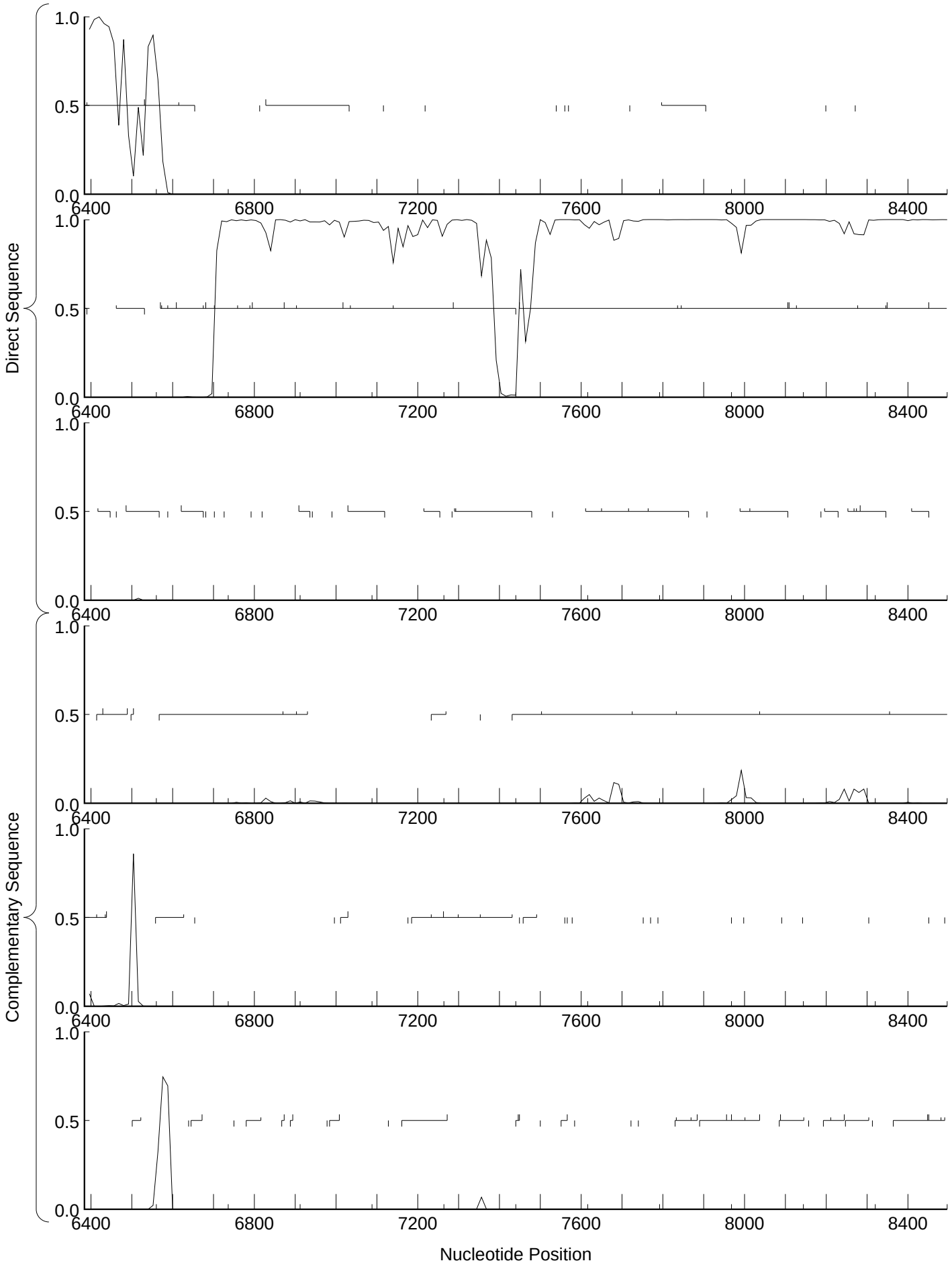
erdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 3/22



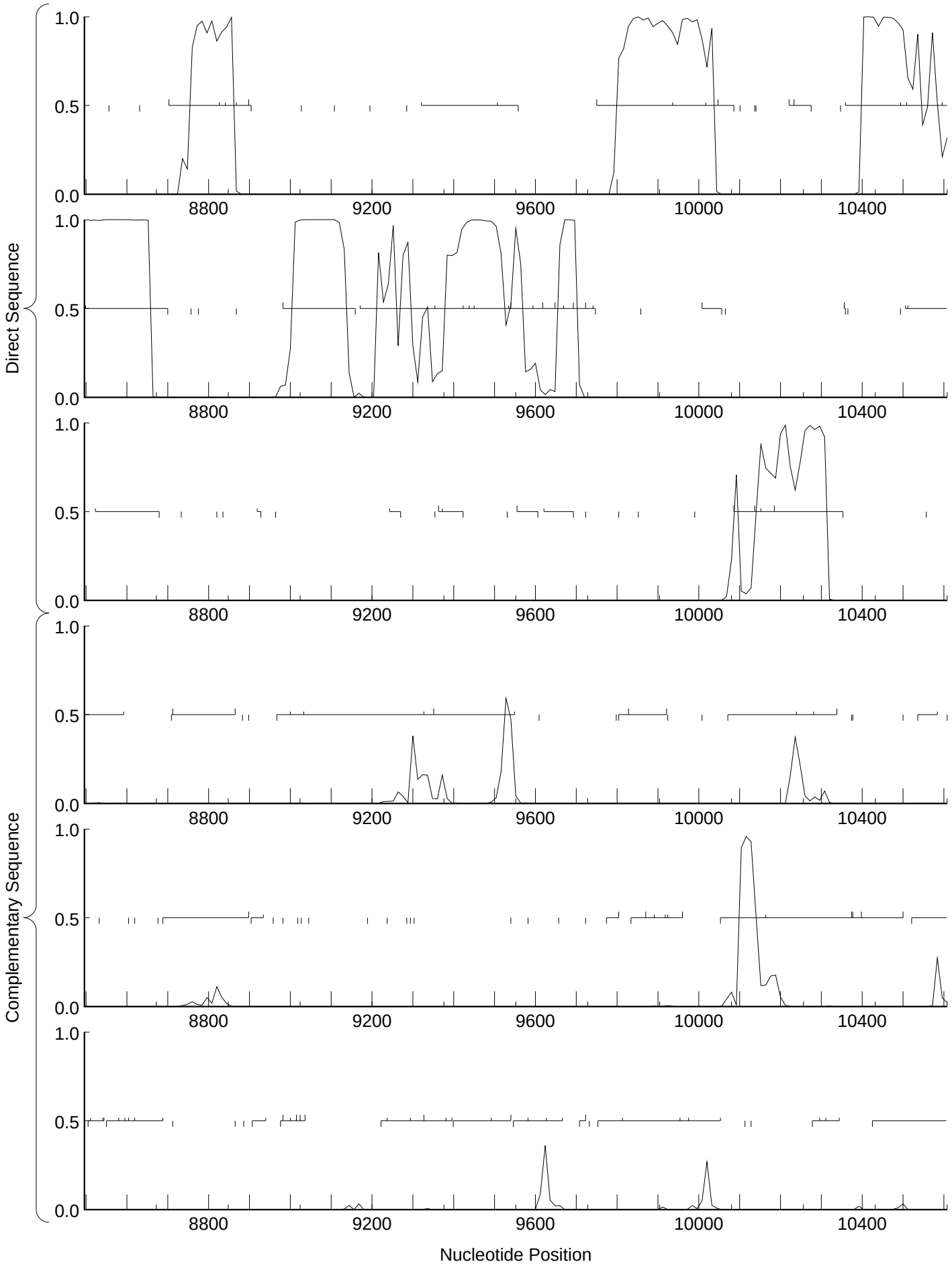
erdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 4/22



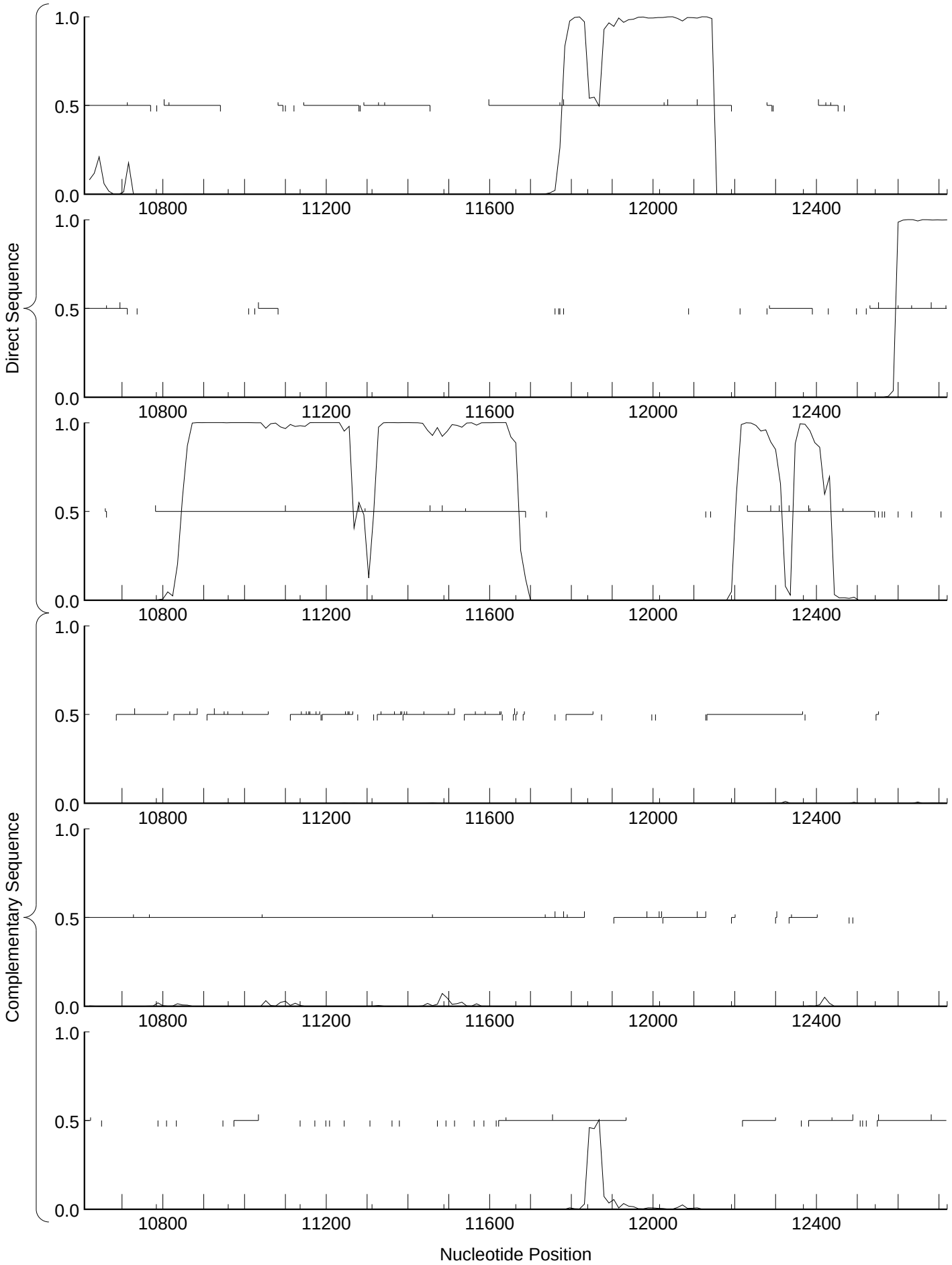
erdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 5/22



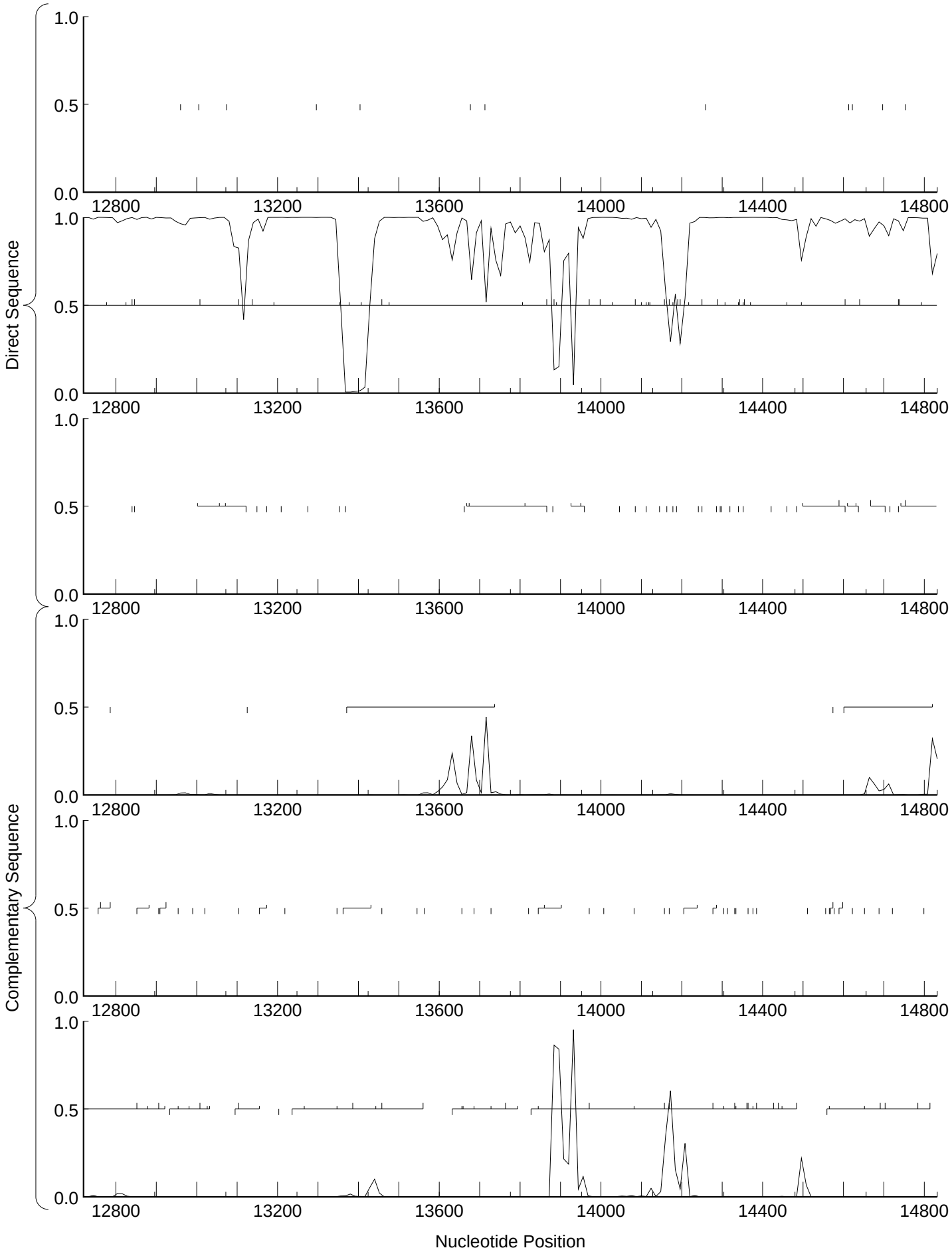
erdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 6/22



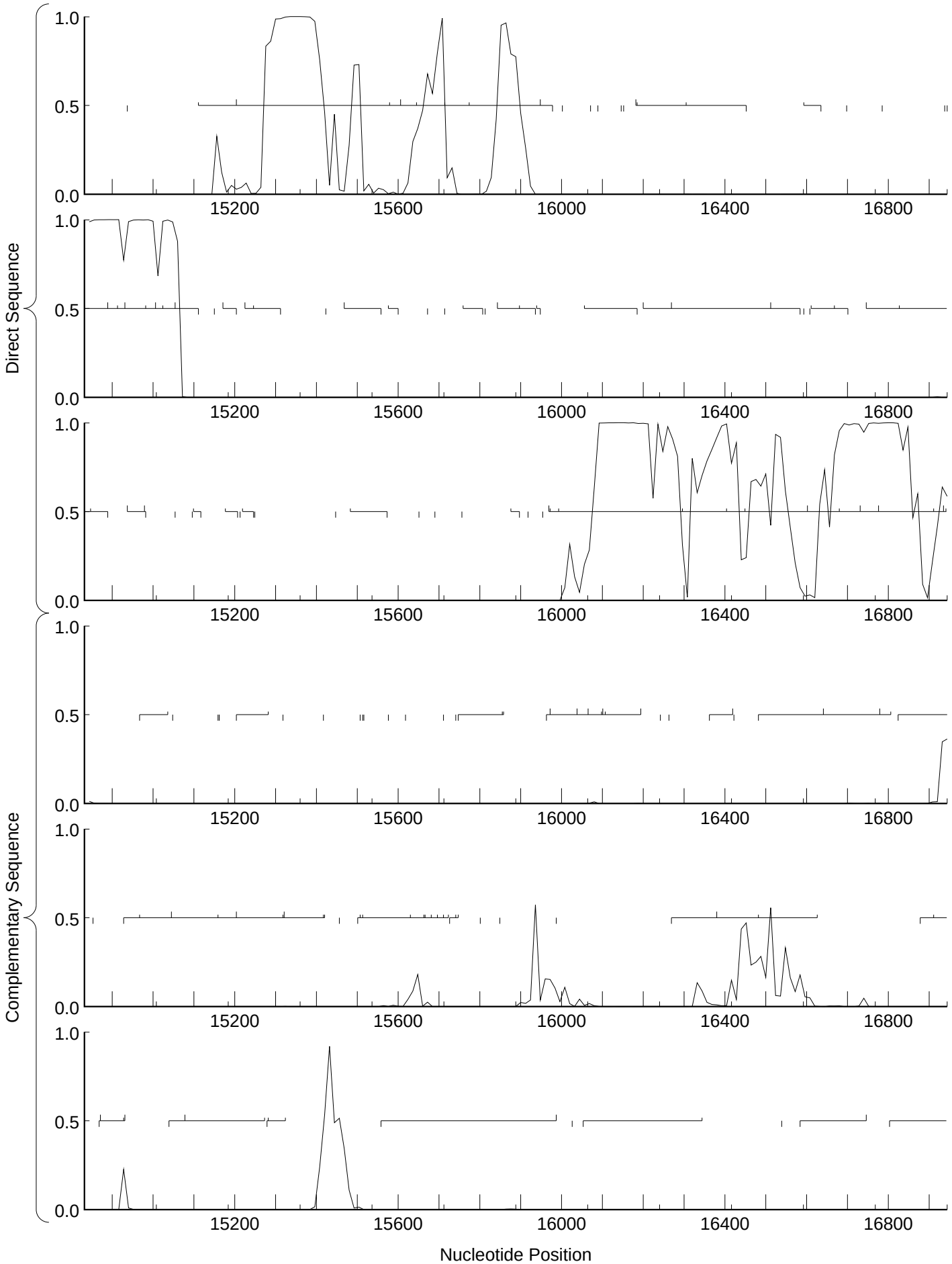
erdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 7/22



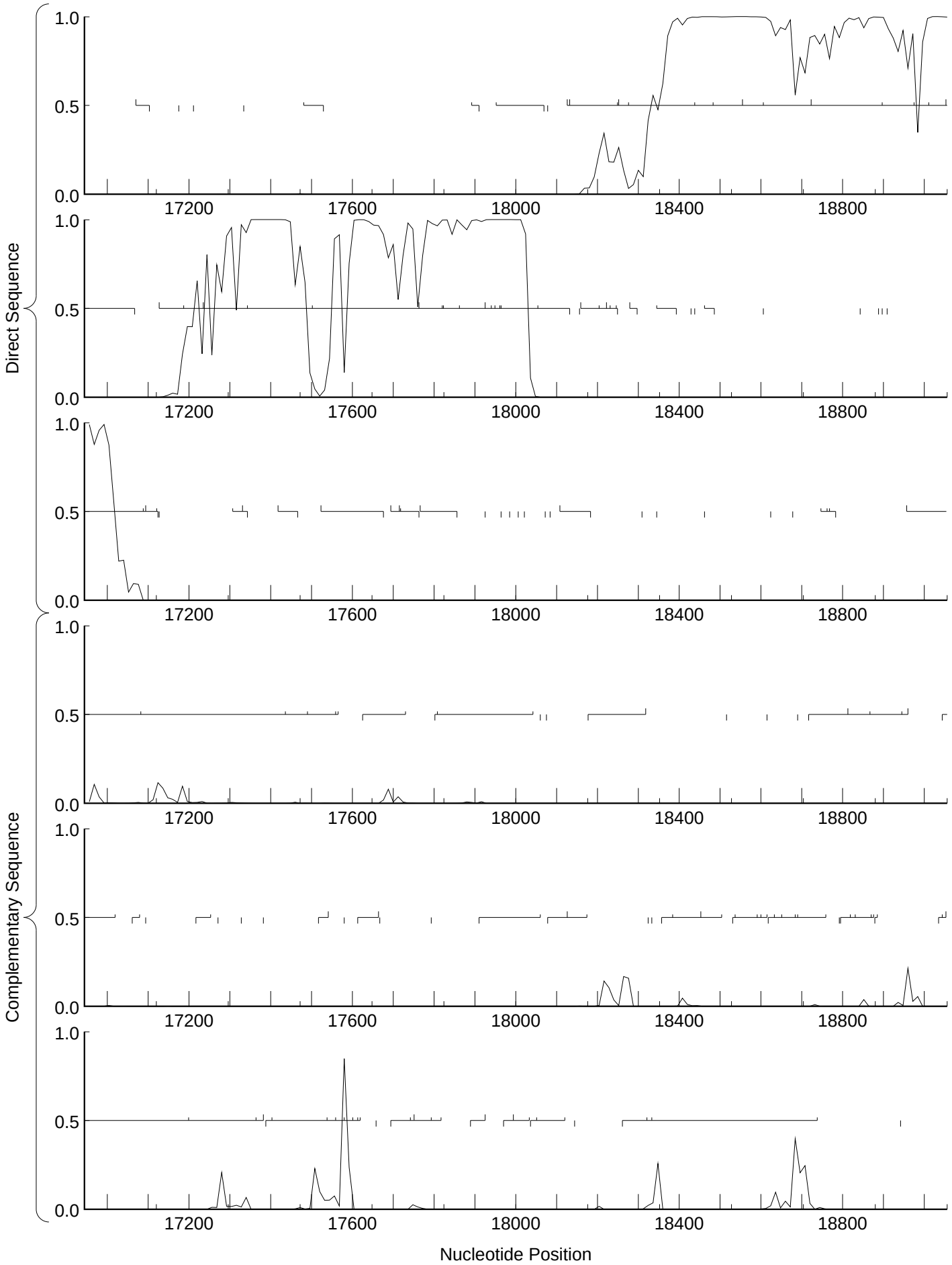
erdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 8/22



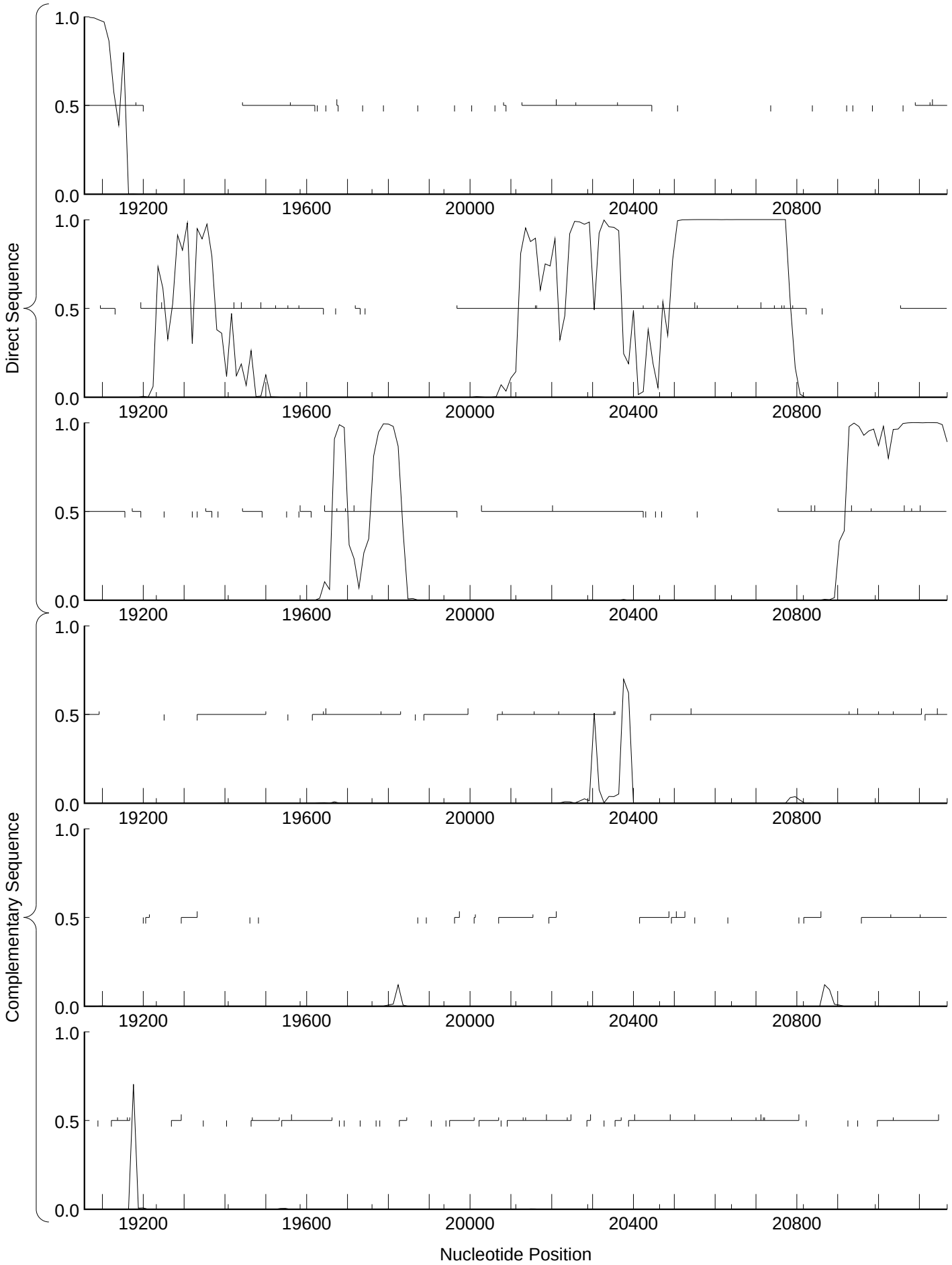
erdale complete sequence, 43391 bp 13-base 3' overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 9/22



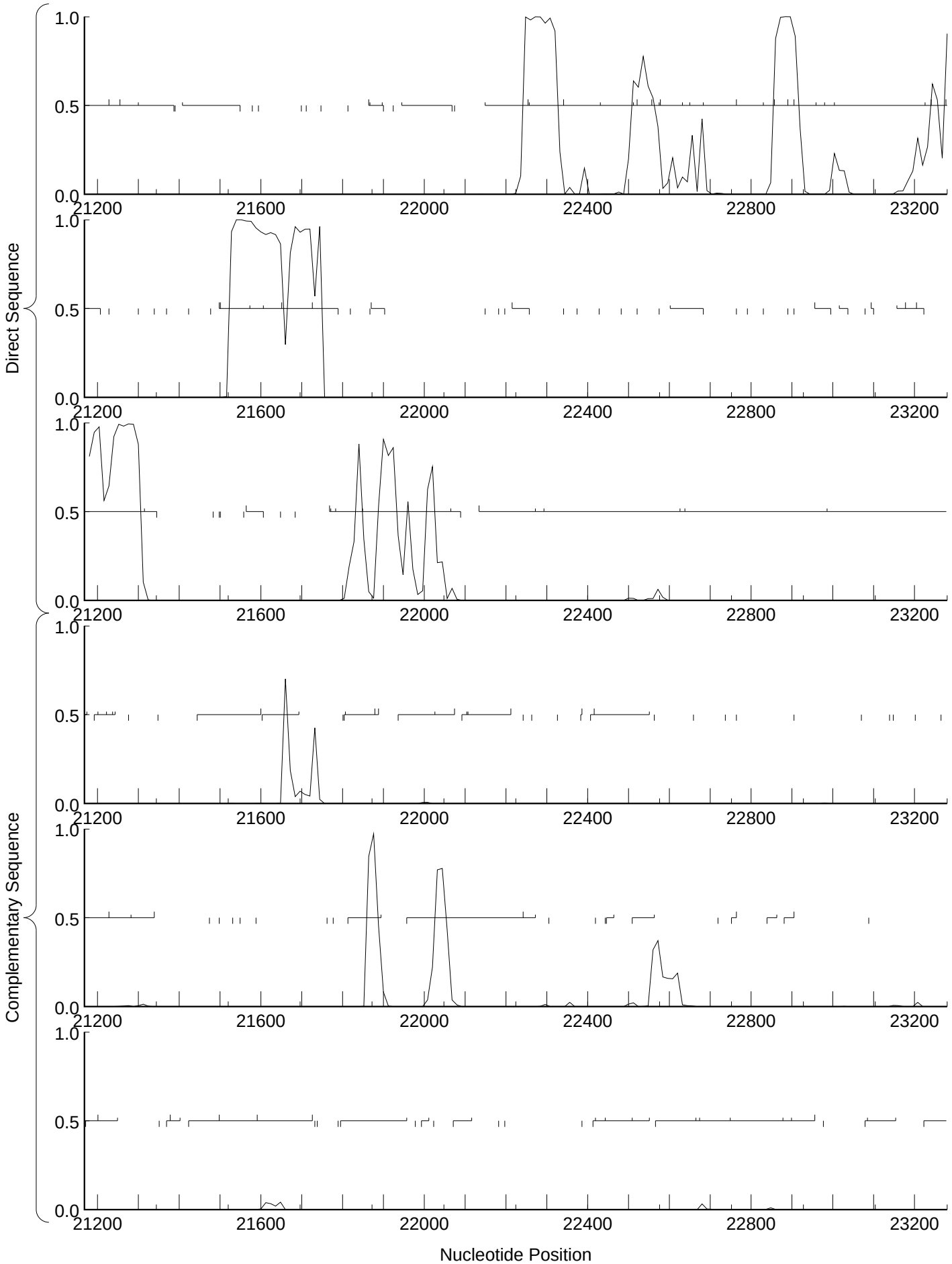
rdale complete sequence, 43391 bp 13-base 3' overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 10/22



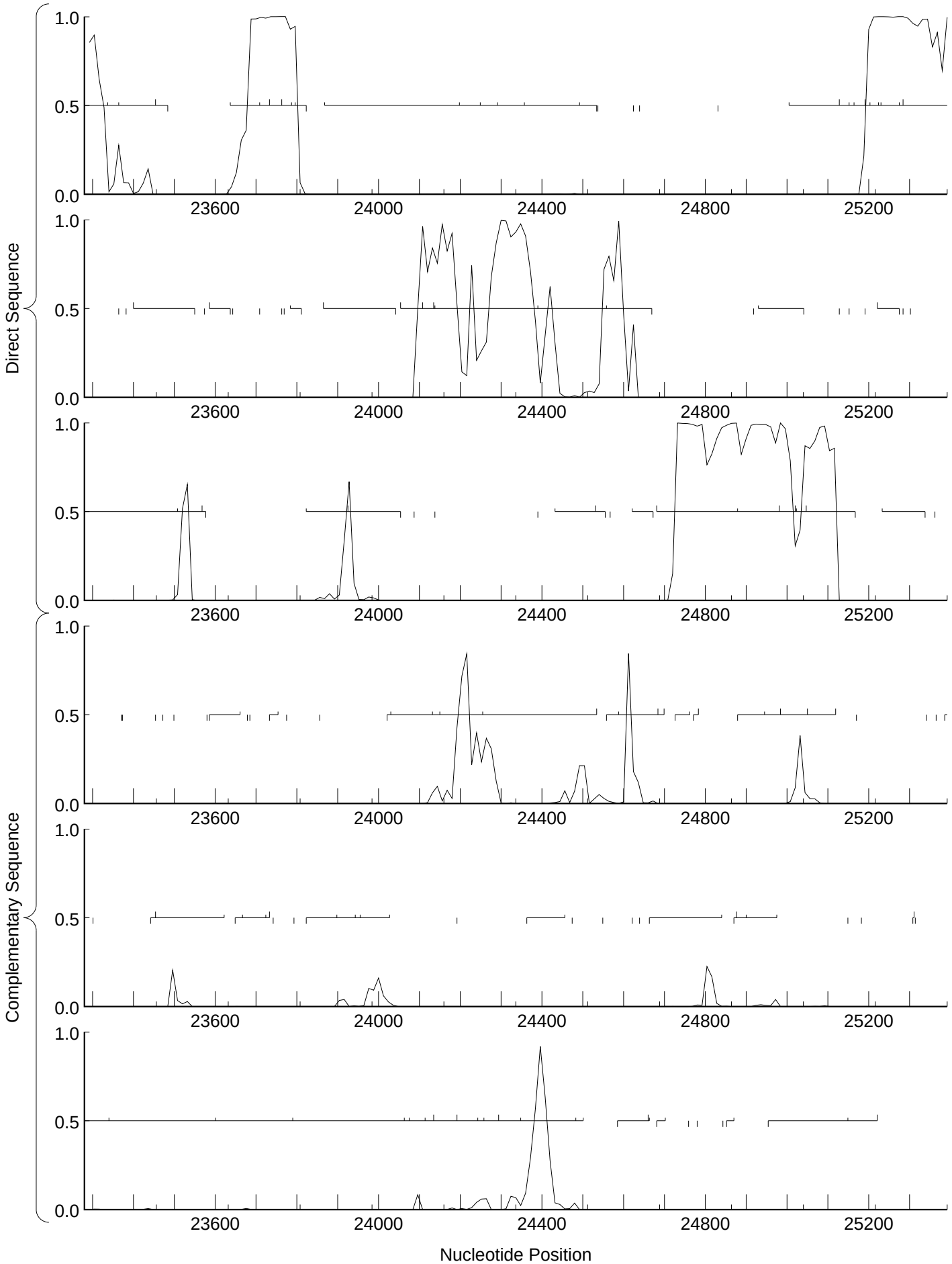
rdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 11/22



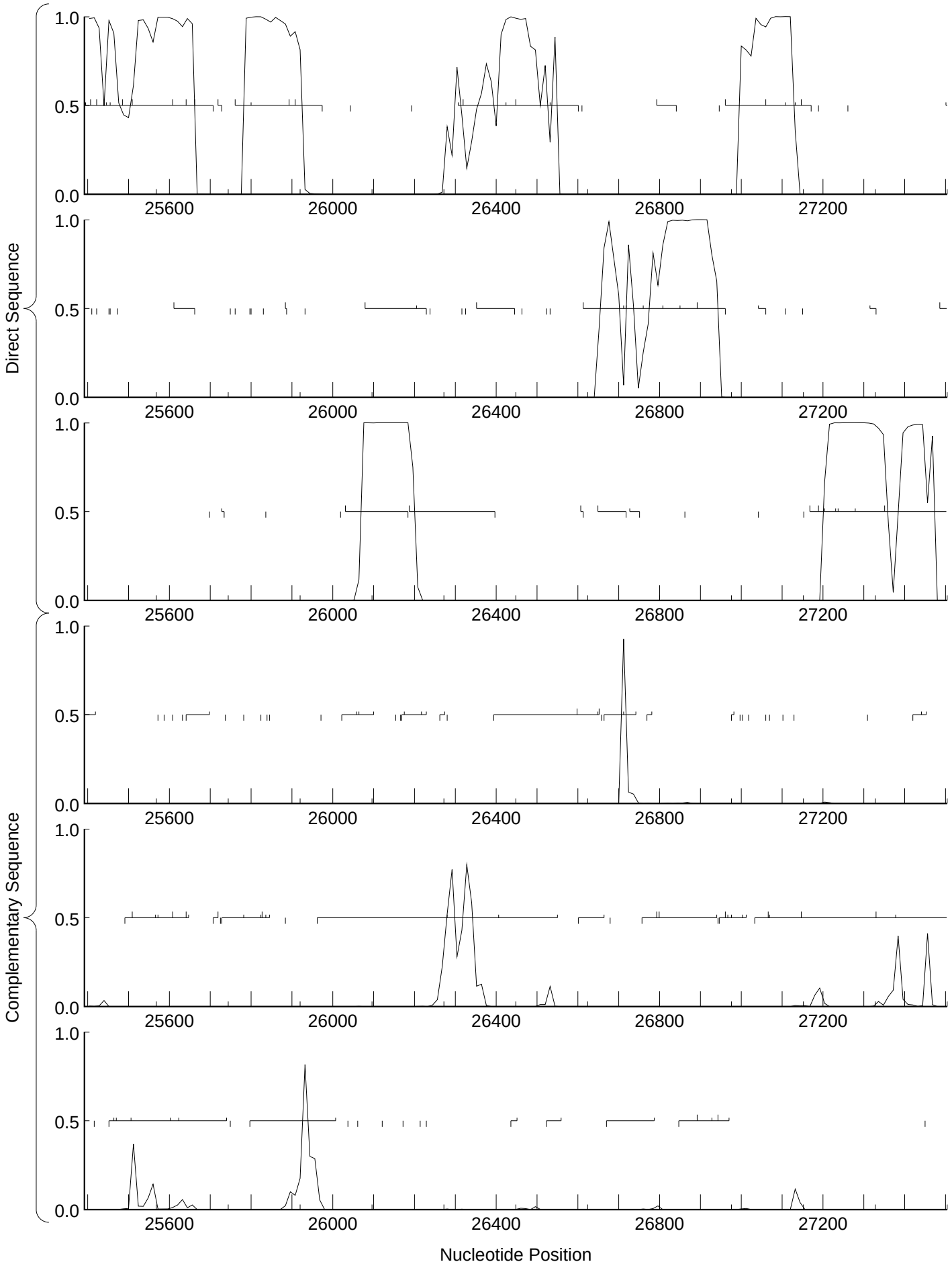
rdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 12/22



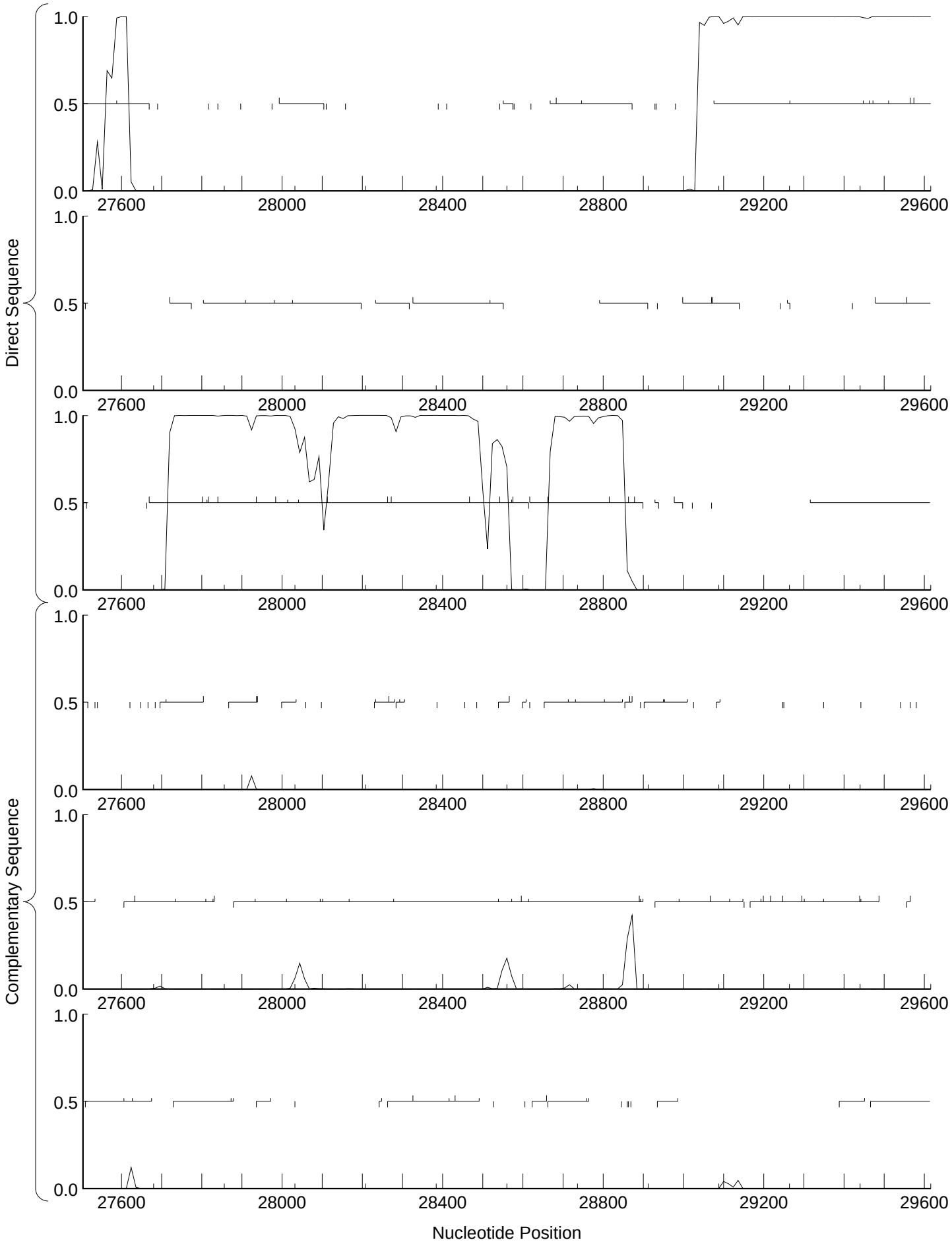
rdale complete sequence, 43391 bp 13-base 3' overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 13/22



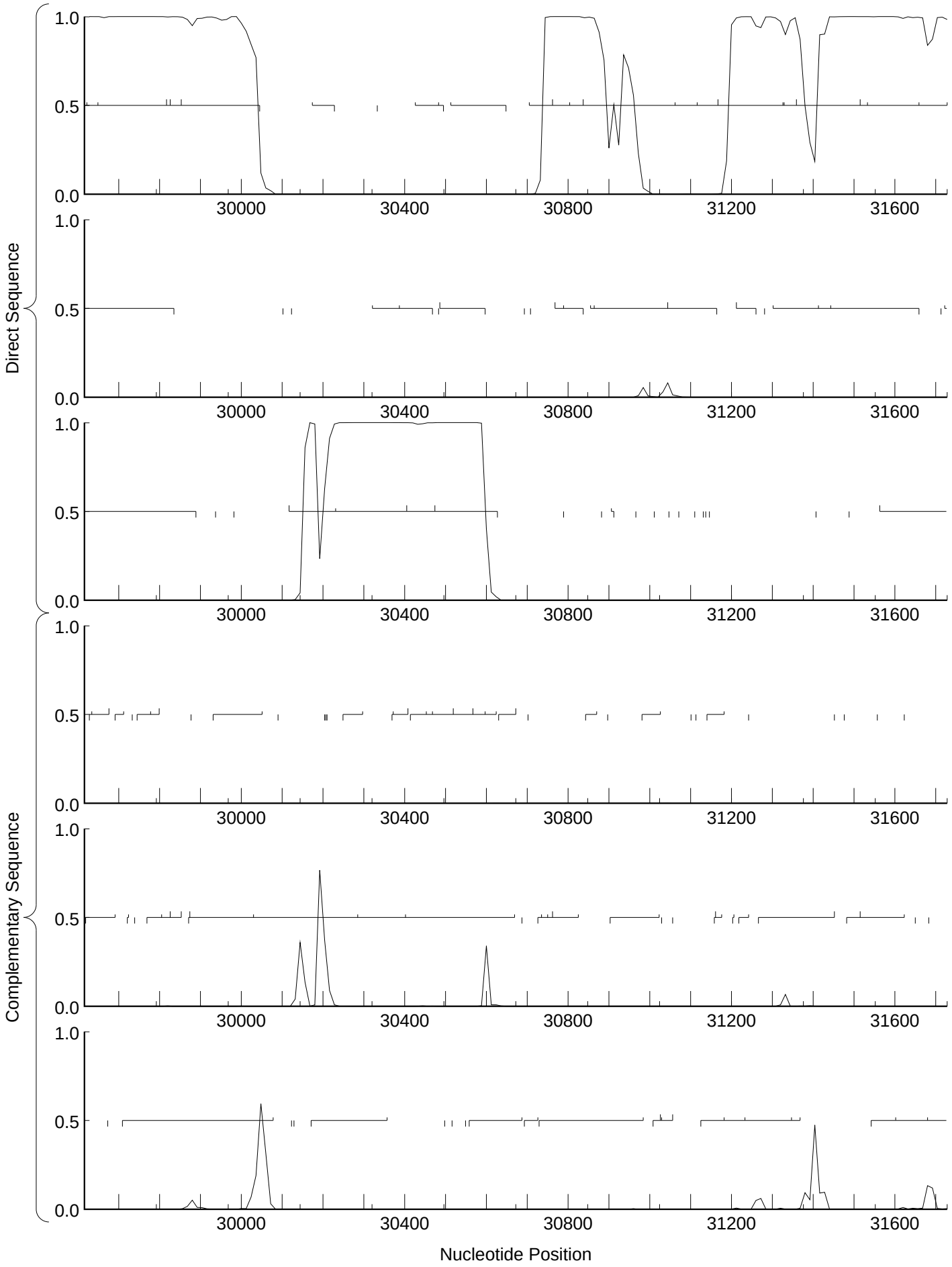
rdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 14/22



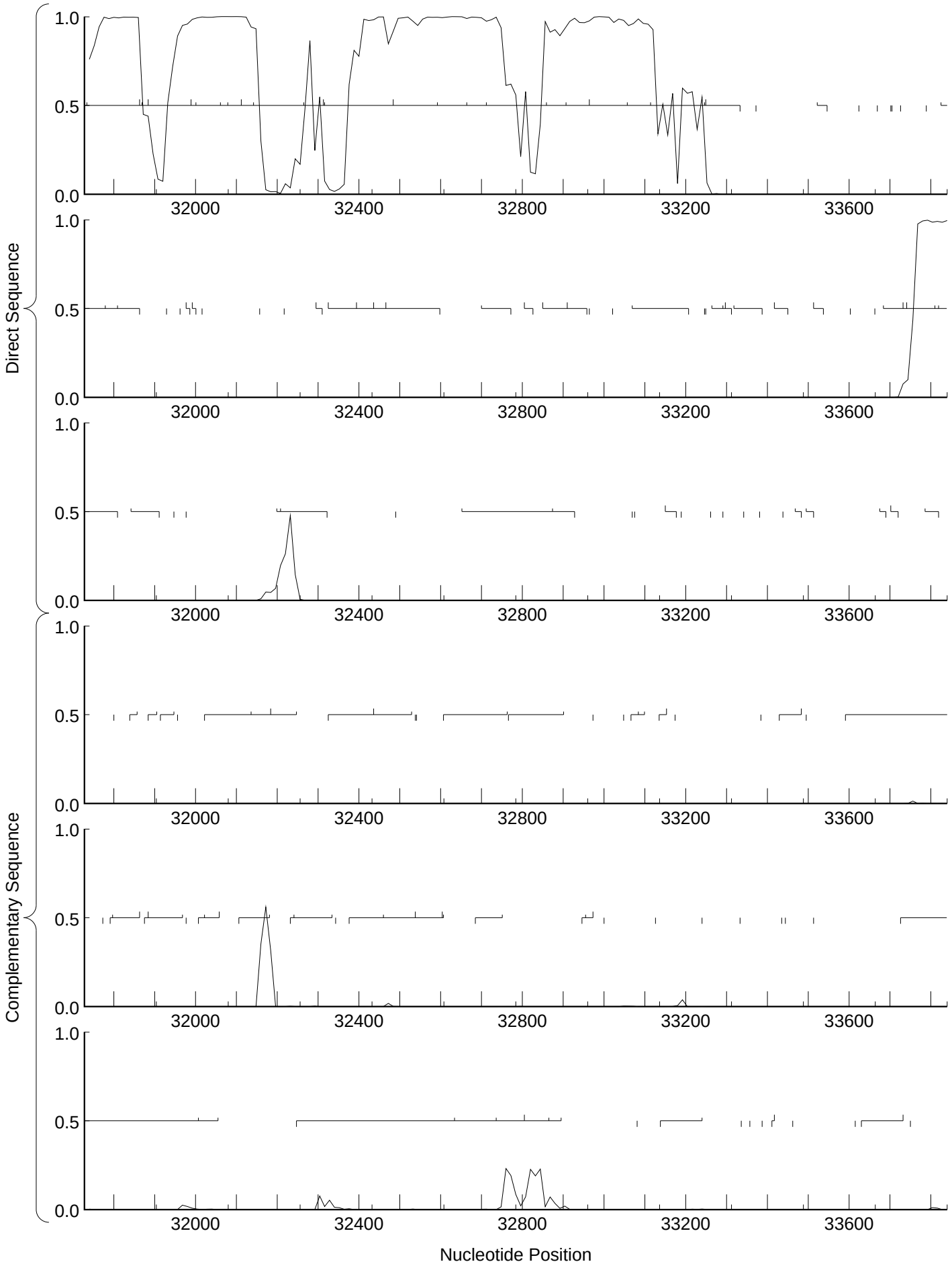
rdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 15/22



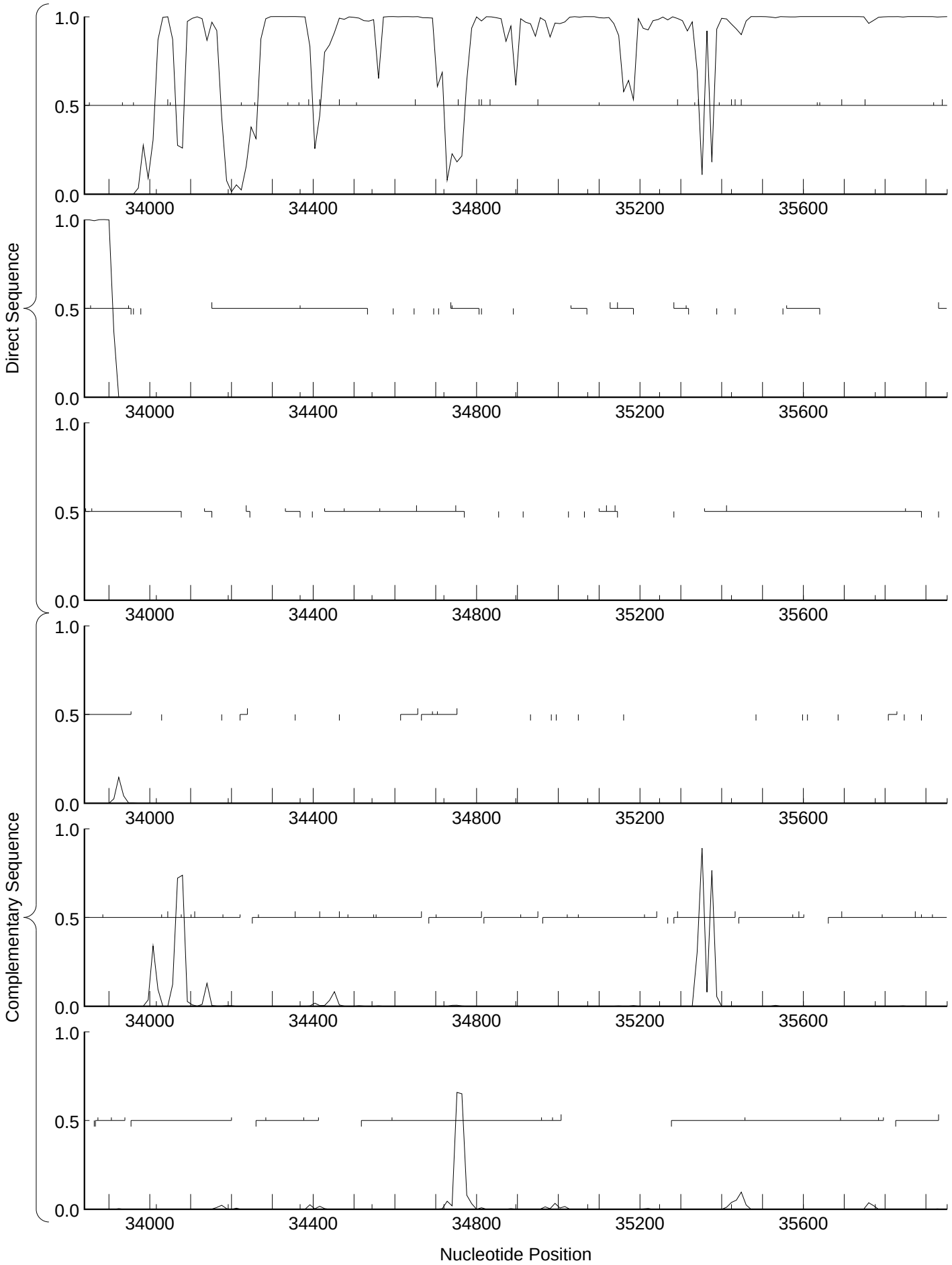
rdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 16/22



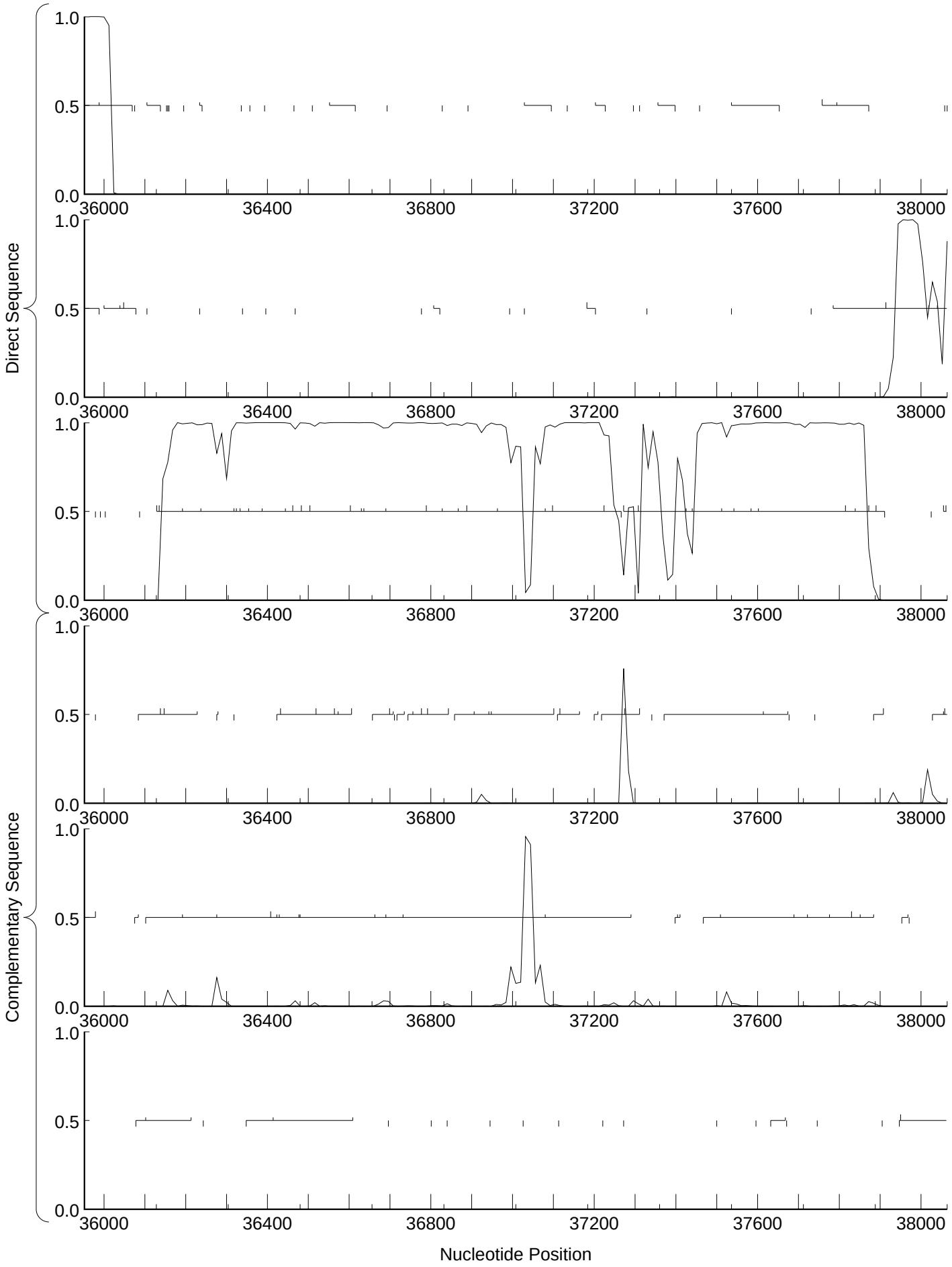
rdale complete sequence, 43391 bp 13-base 3' overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 17/22



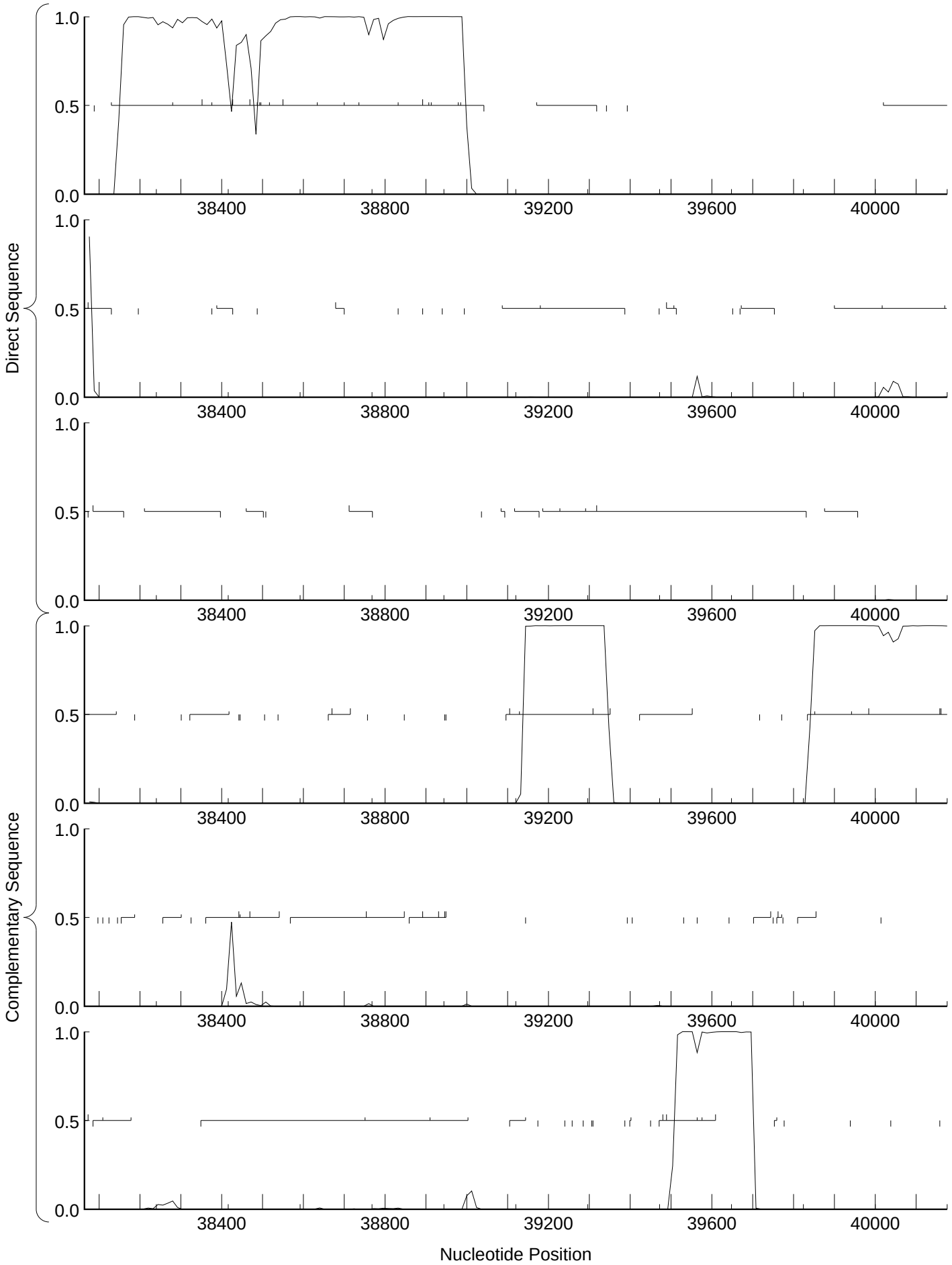
rdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 18/22



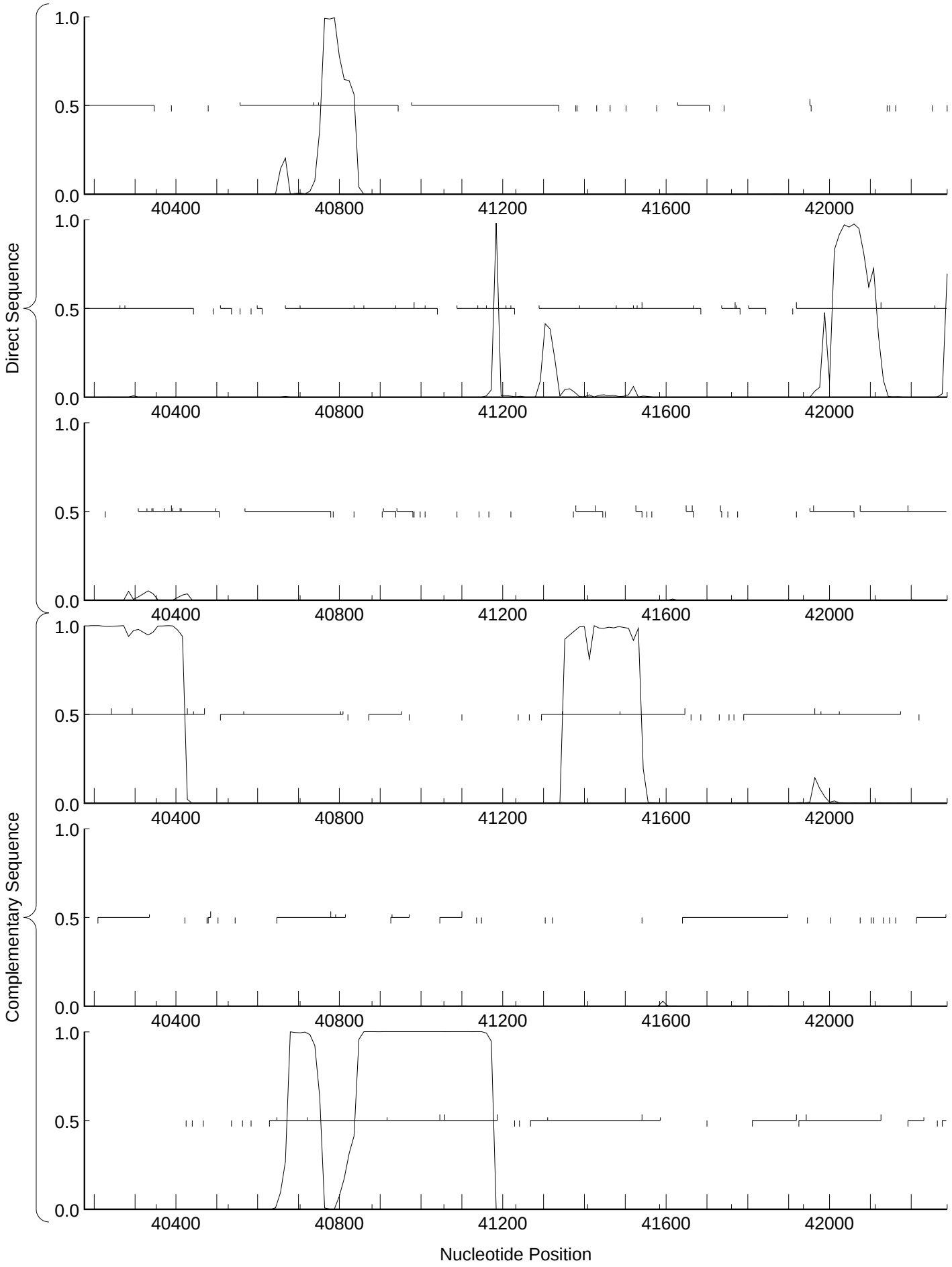
rdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 19/22



rdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 20/22



rdale complete sequence, 43391 bp 13-base 3' overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 21/22



rdale complete sequence, 43391 bp 13-base 3’ overhang (GGTAACCGTGATA), Cluster AK, Order 4, Window 96, Step 12, 22/22

