

# GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

## PROGRAM INFORMATION

Sequence : Mycobacterium phage Deb65 complete sequence, 55767 bp including 10-base 3' overhang (CGGACG  
Analysis Date : 2/8/25 at 4:25:22  
Pages : 28  
Sequence Length : 55767 bp  
GC Content : 61.55%

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,

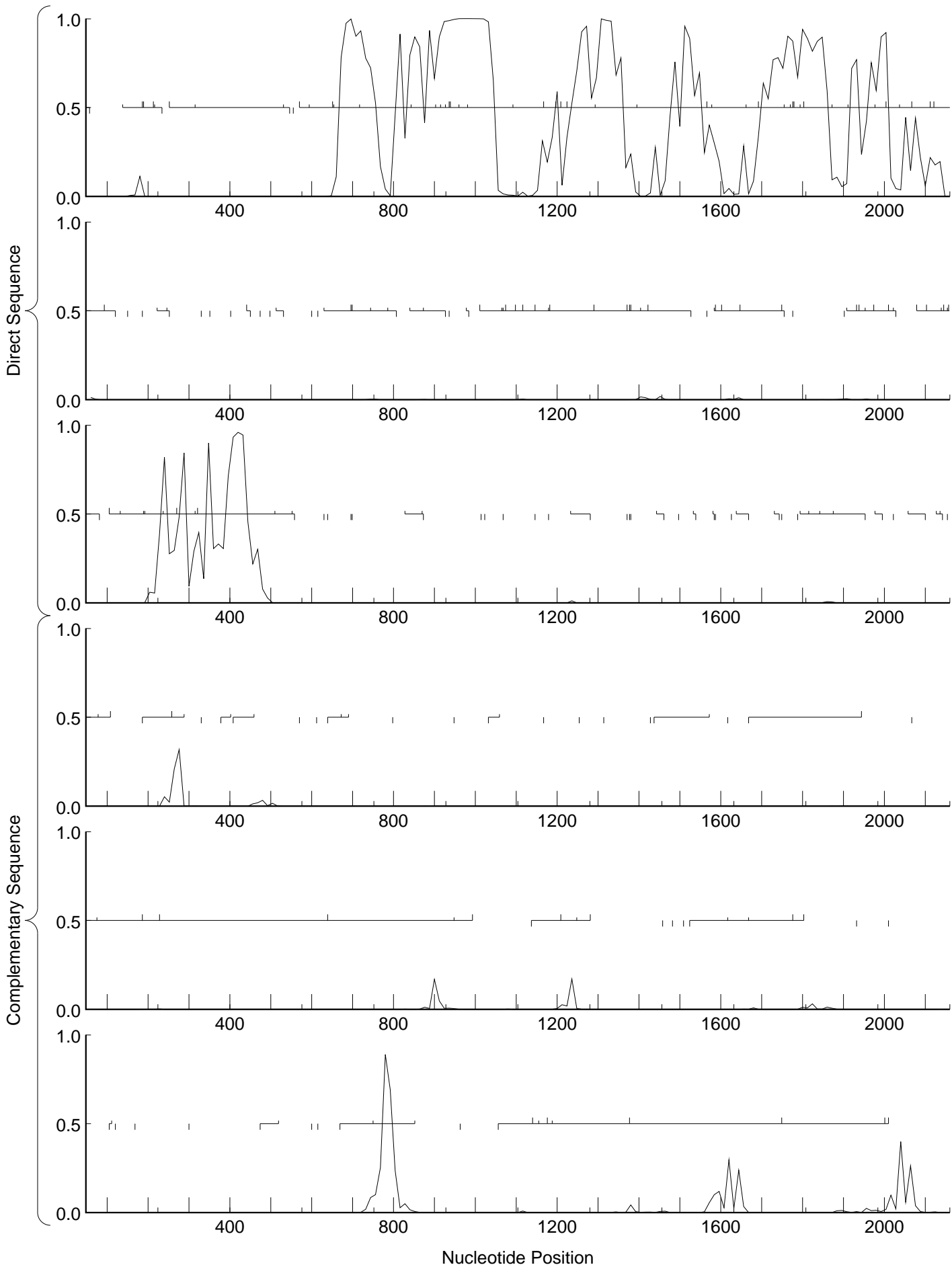
## MATRIX INFORMATION

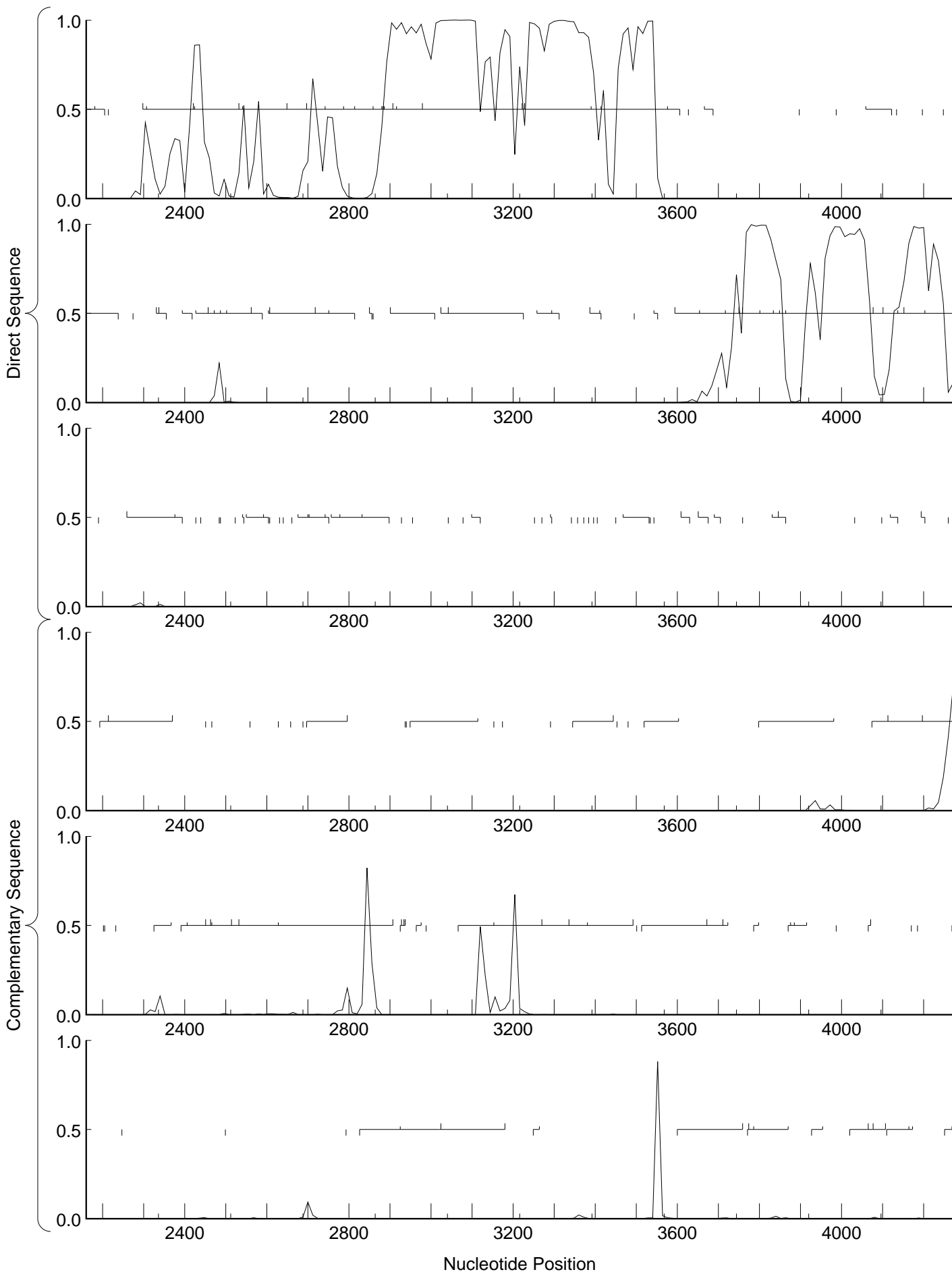
Matrix : Mycobacterium\_tuberculosis\_H37Rv  
Author : -  
Order : 4

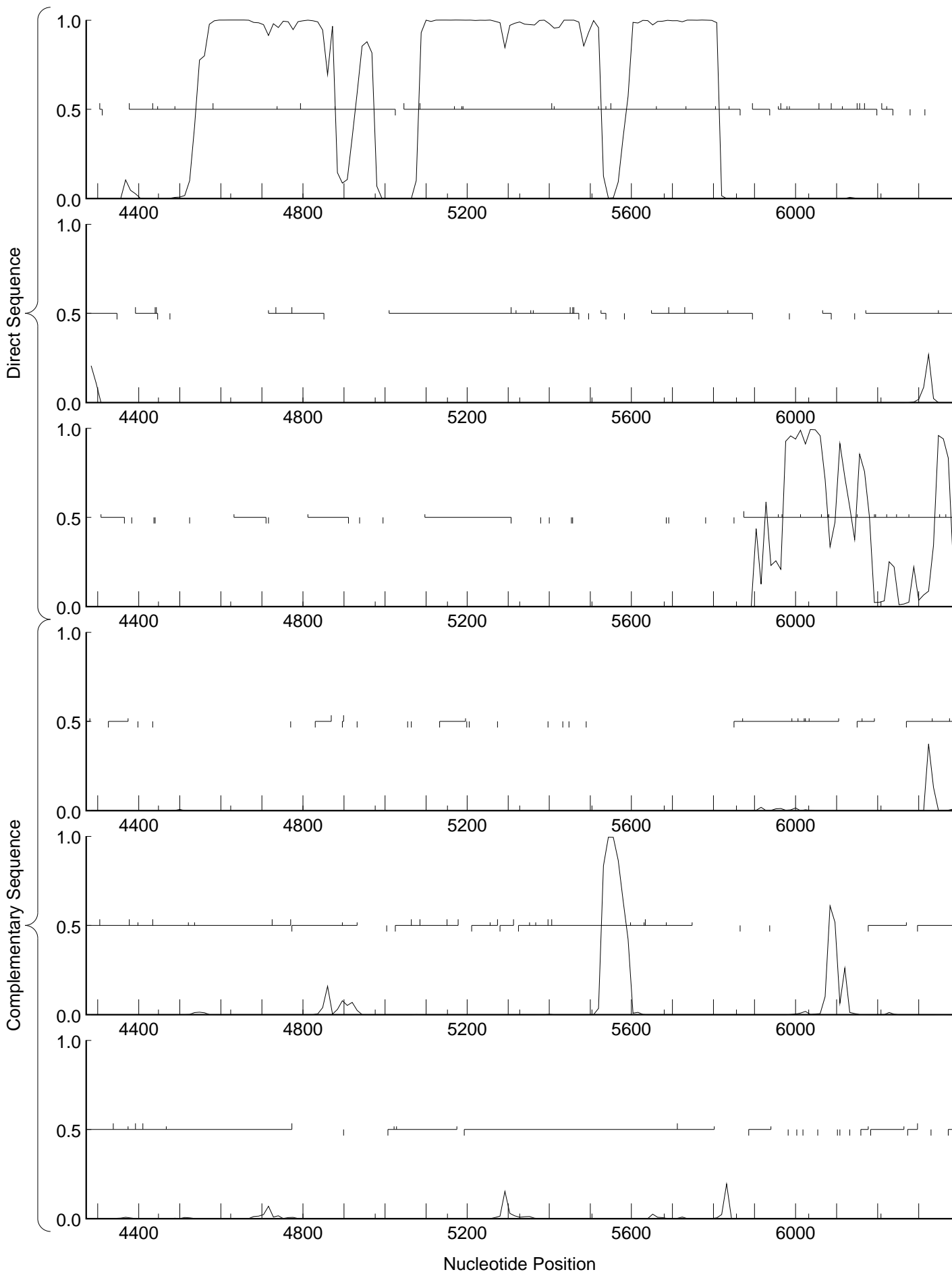
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

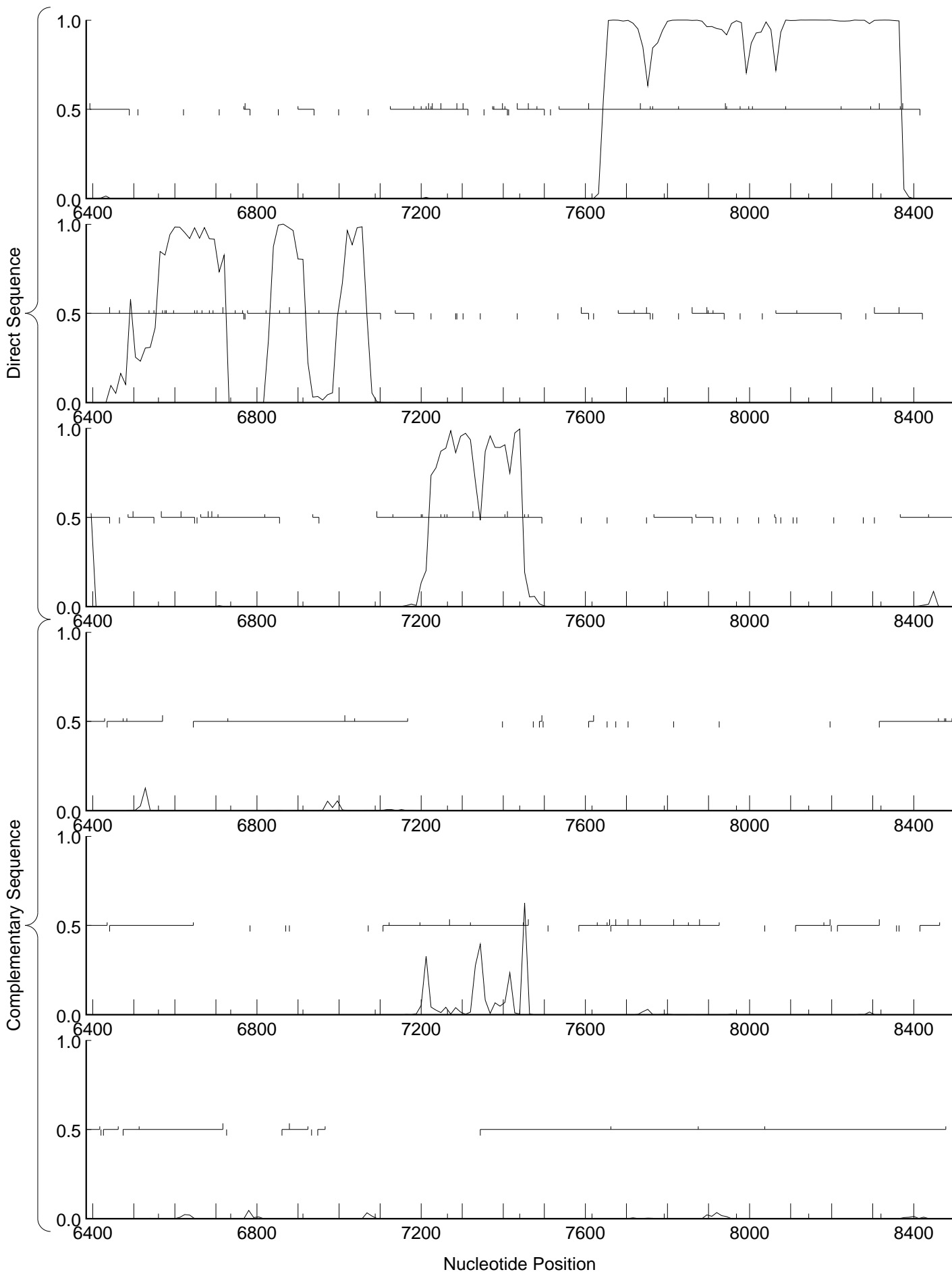
### Matrix notes & comments

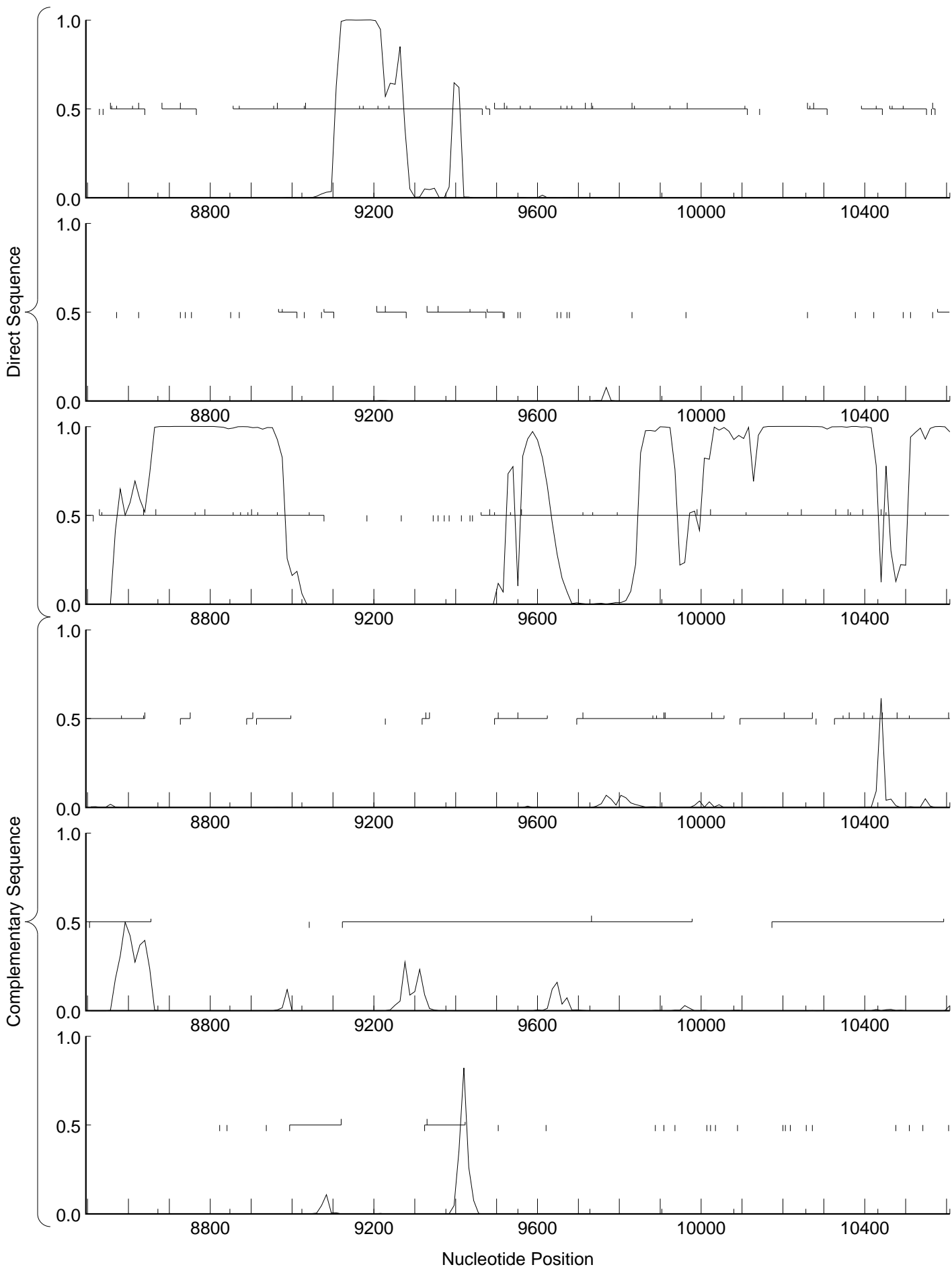
Training set derived by GeneMarkS, 4.27 September 2014  
Tue Sep 23 15:14:37 2014

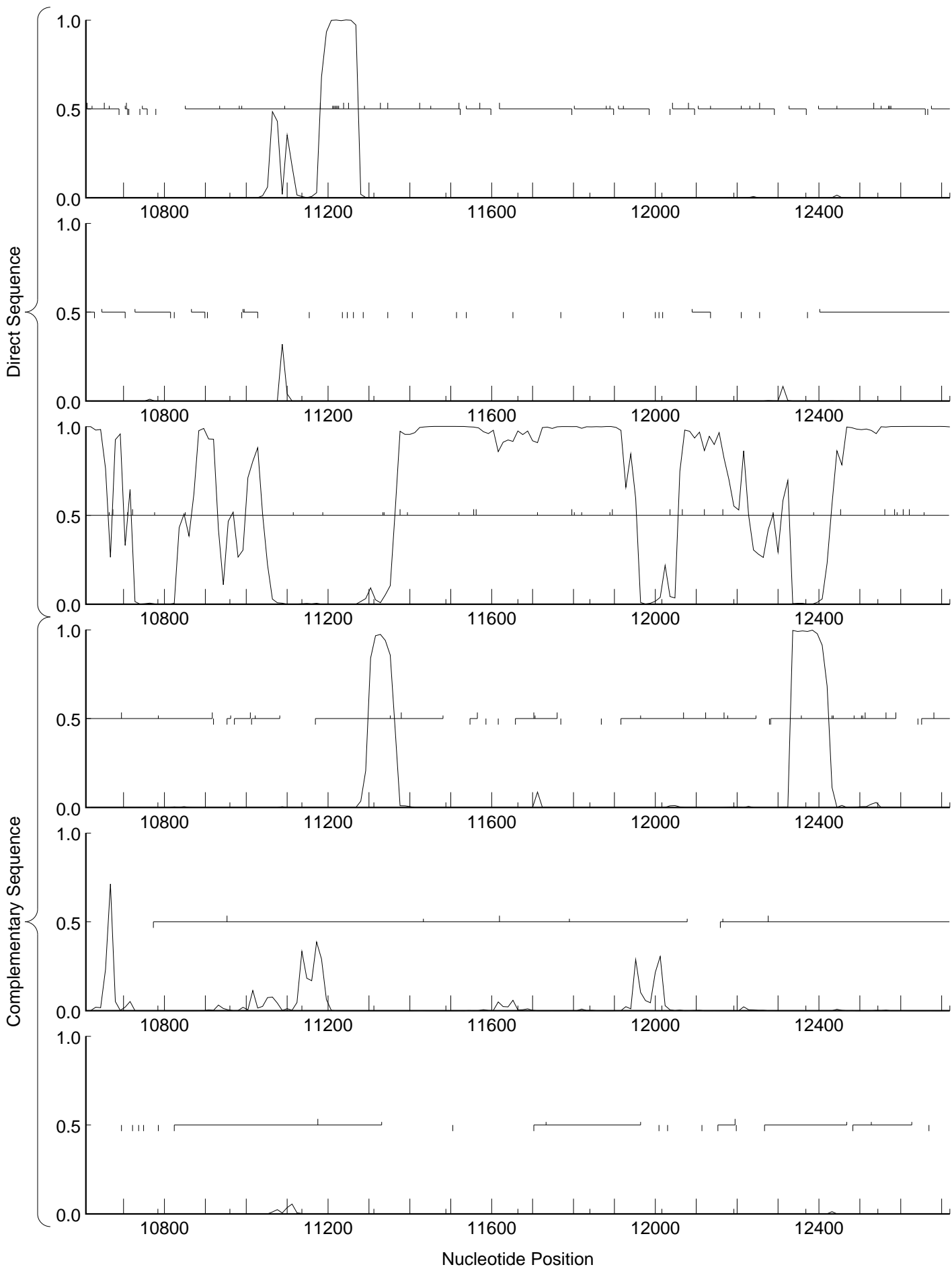


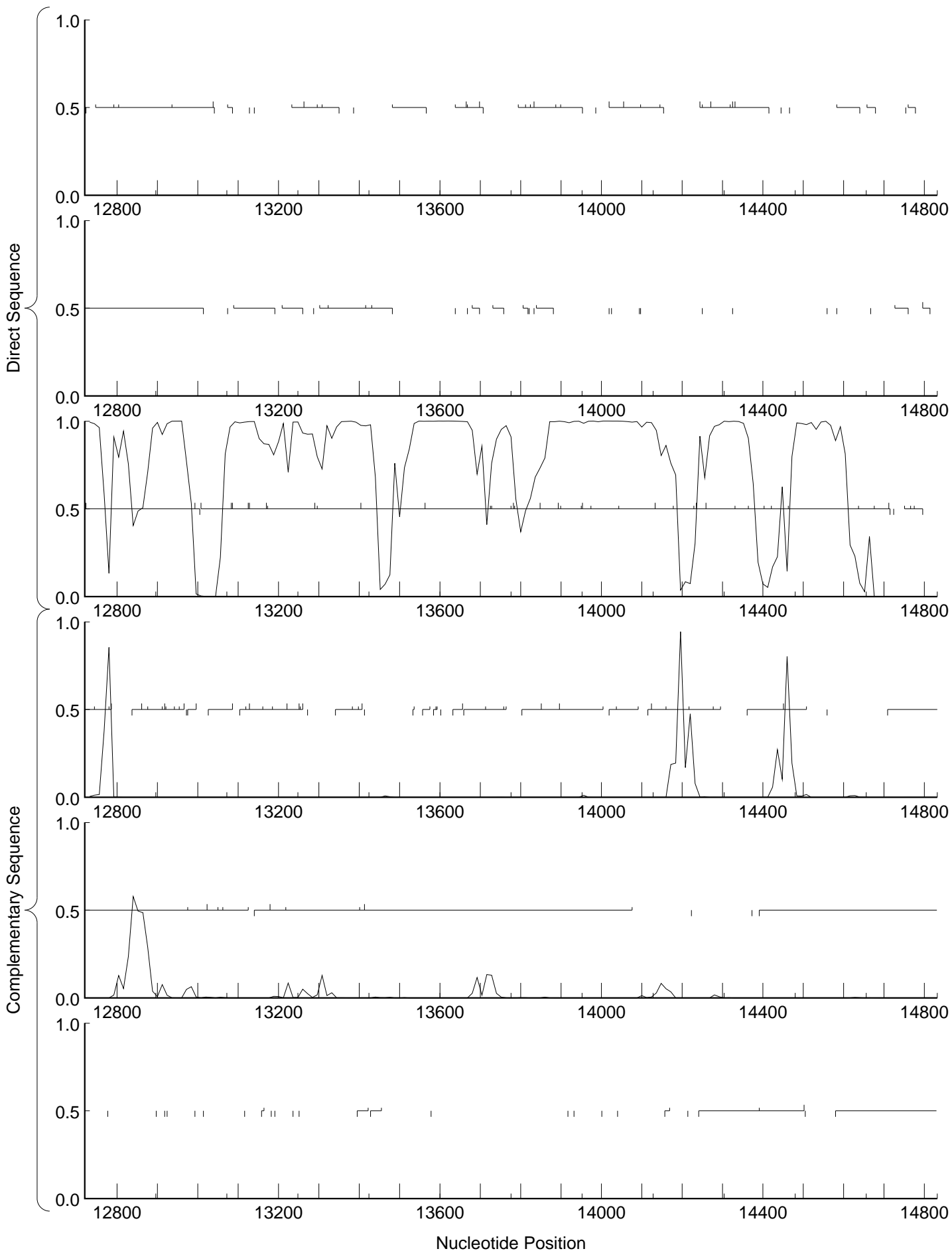




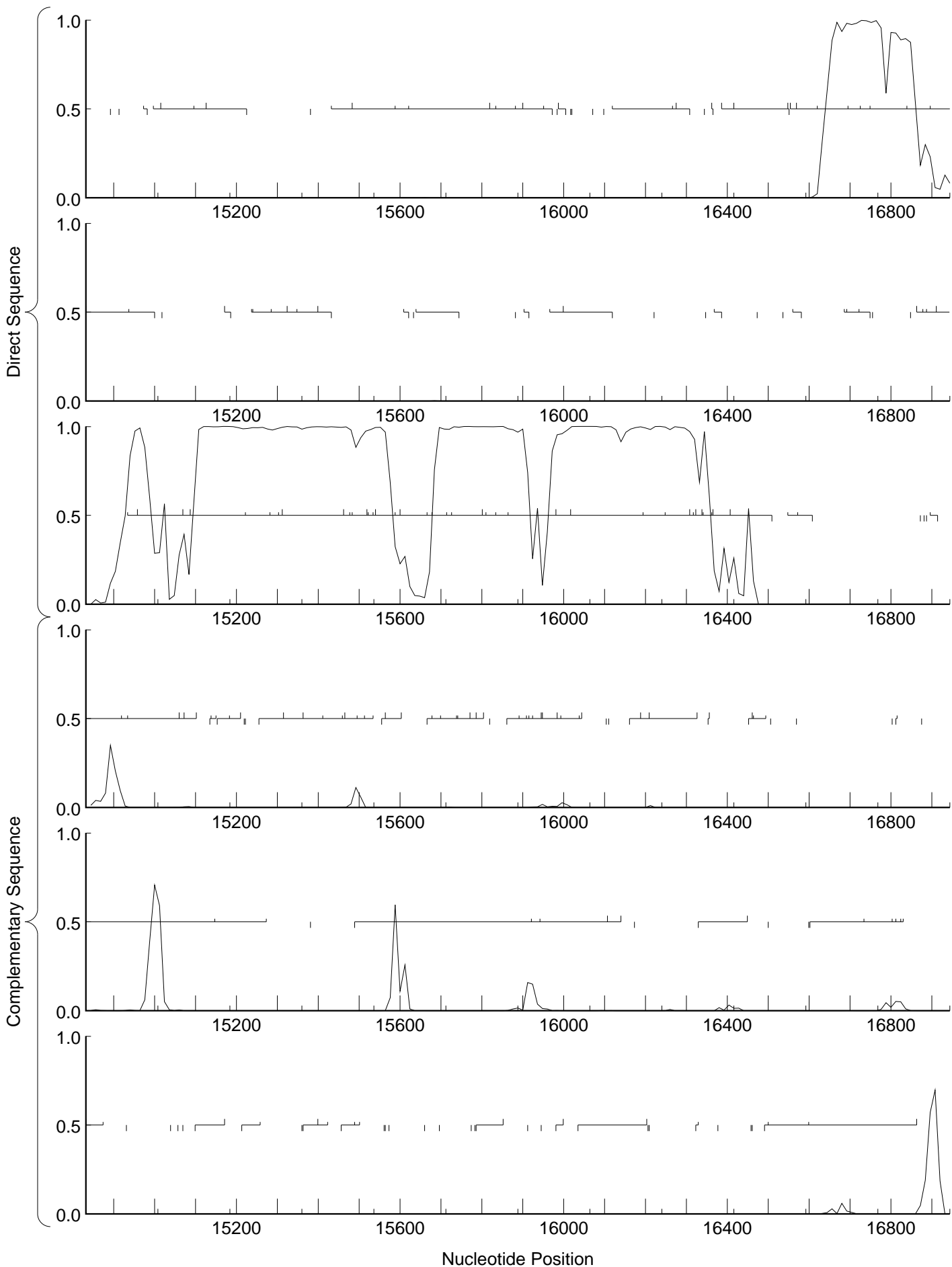


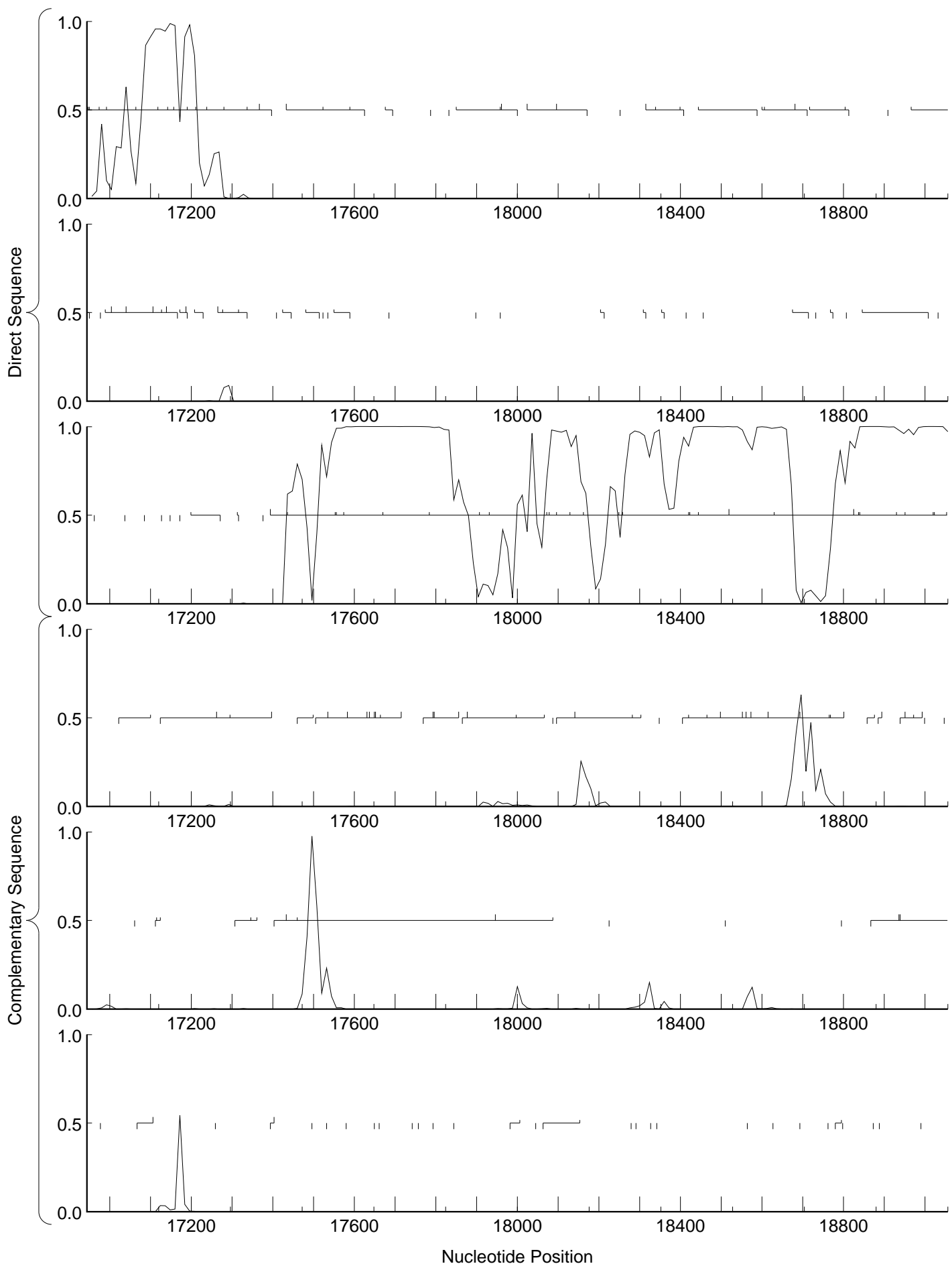


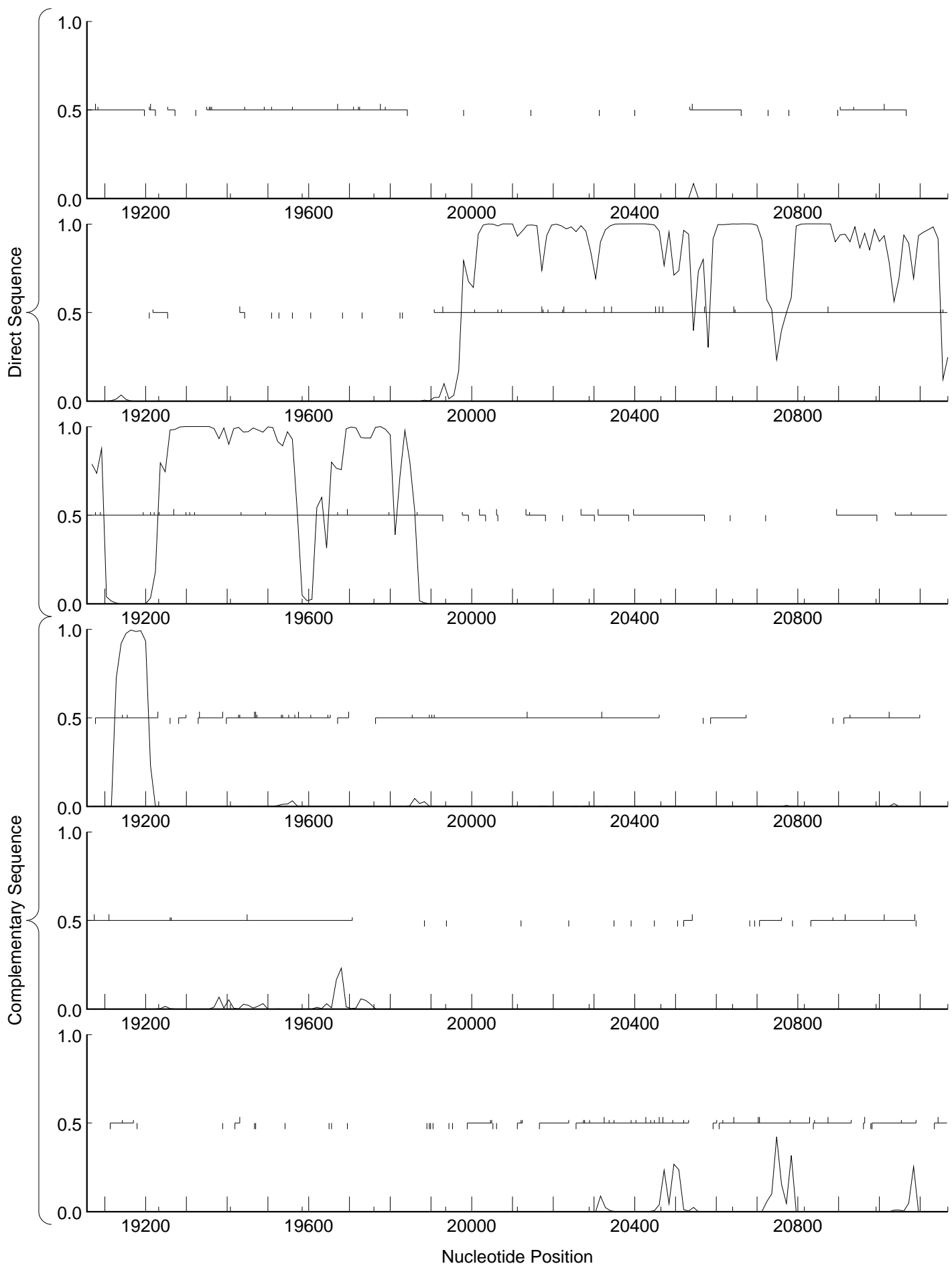


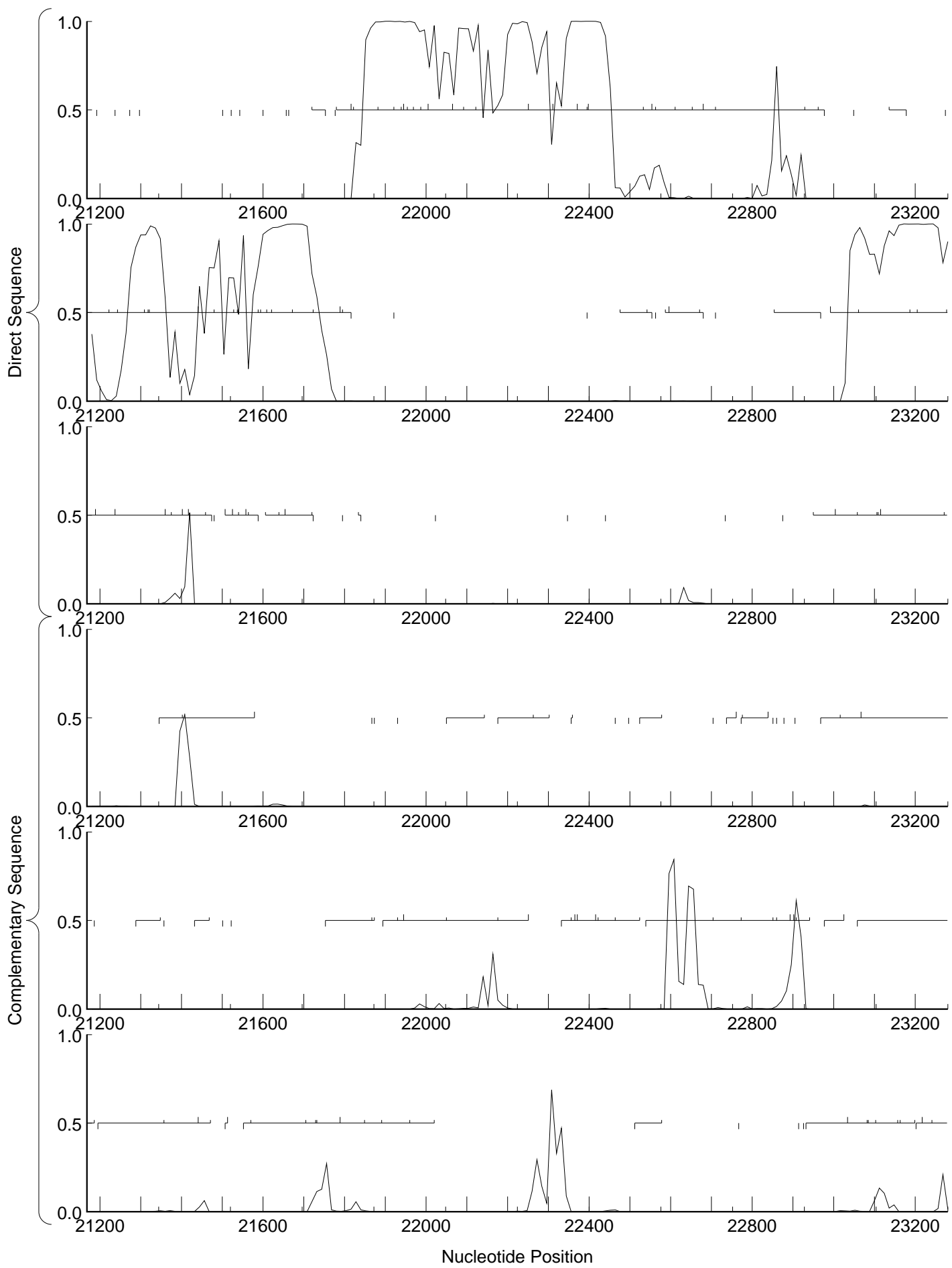




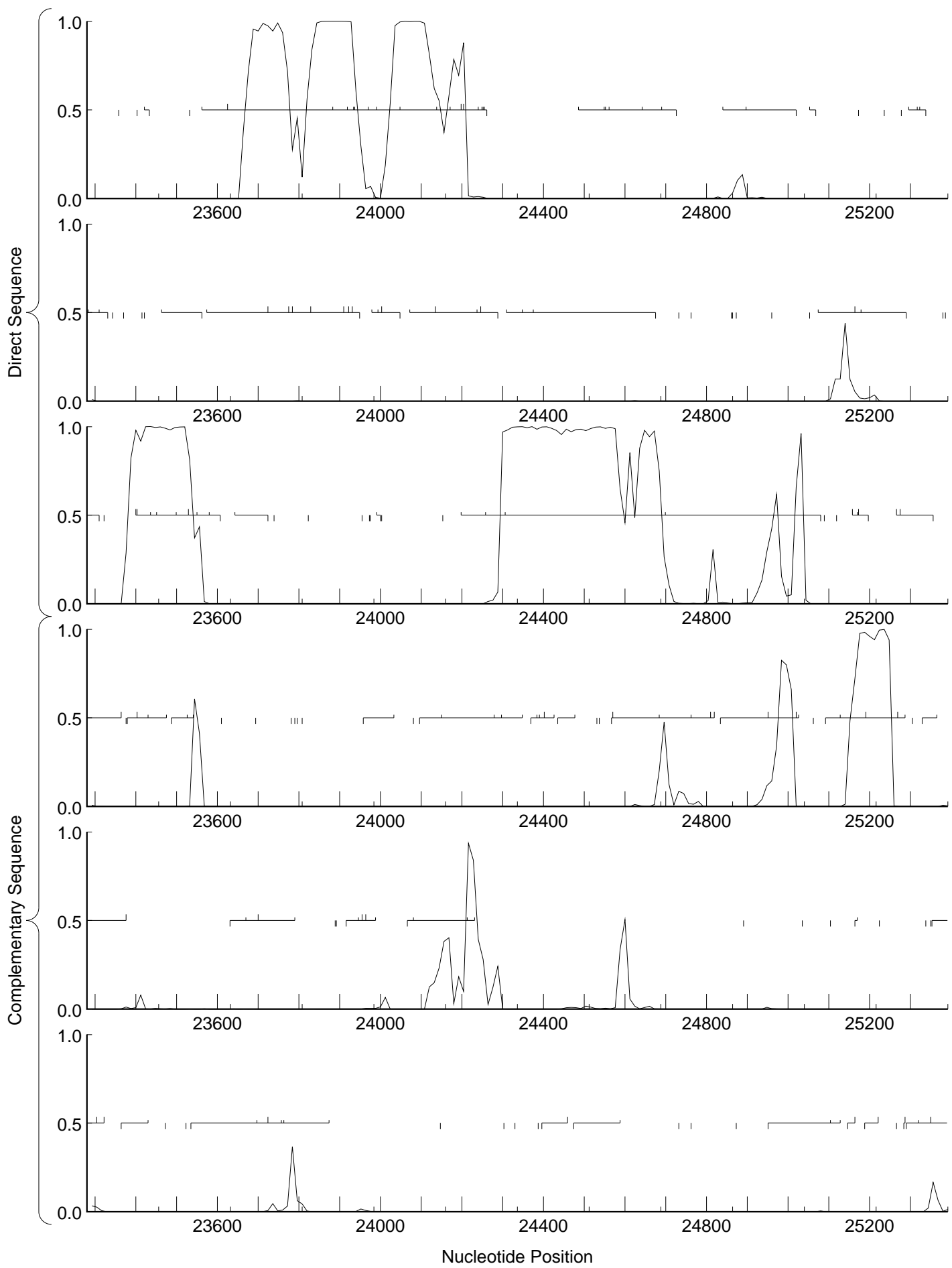




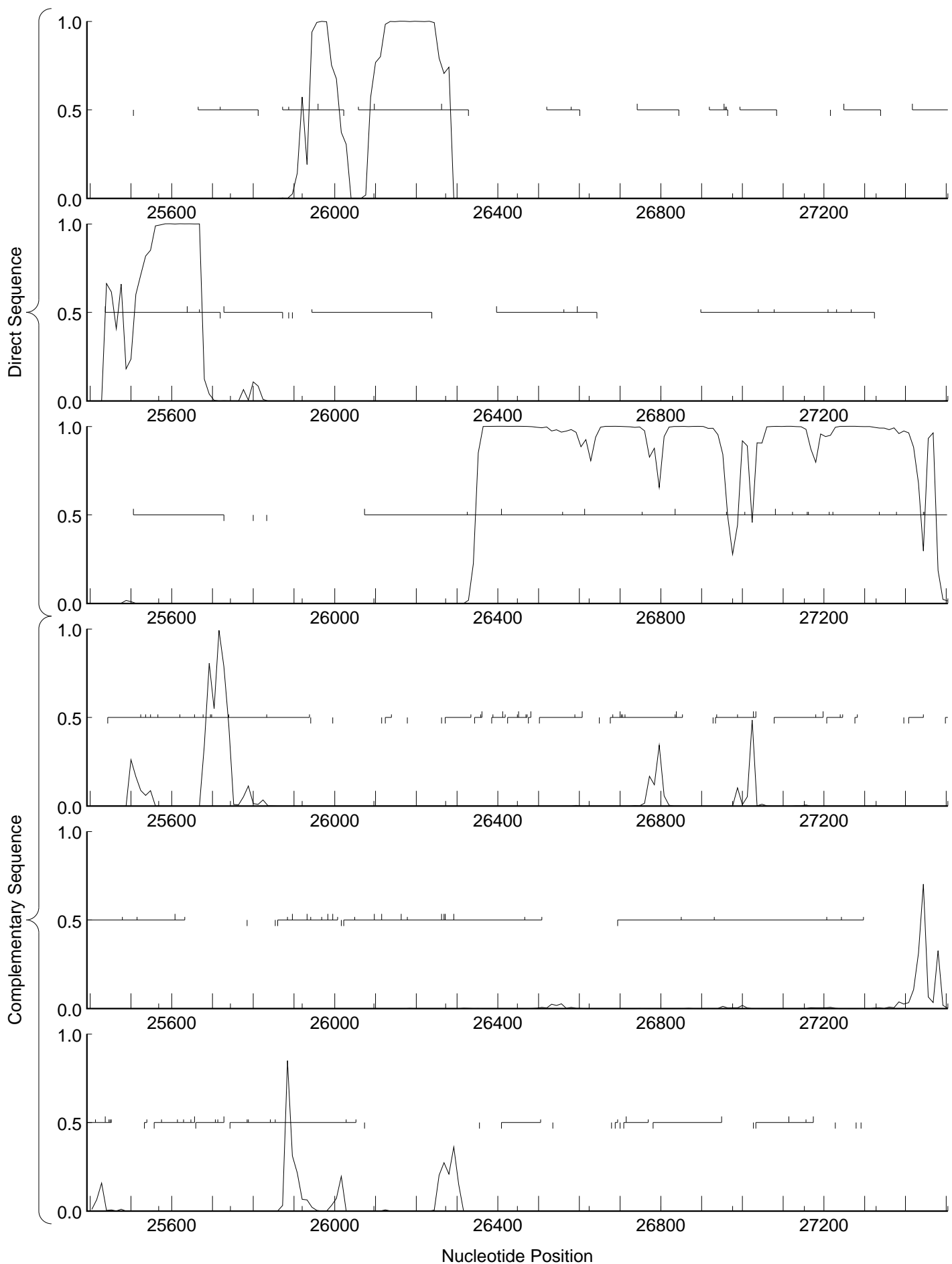


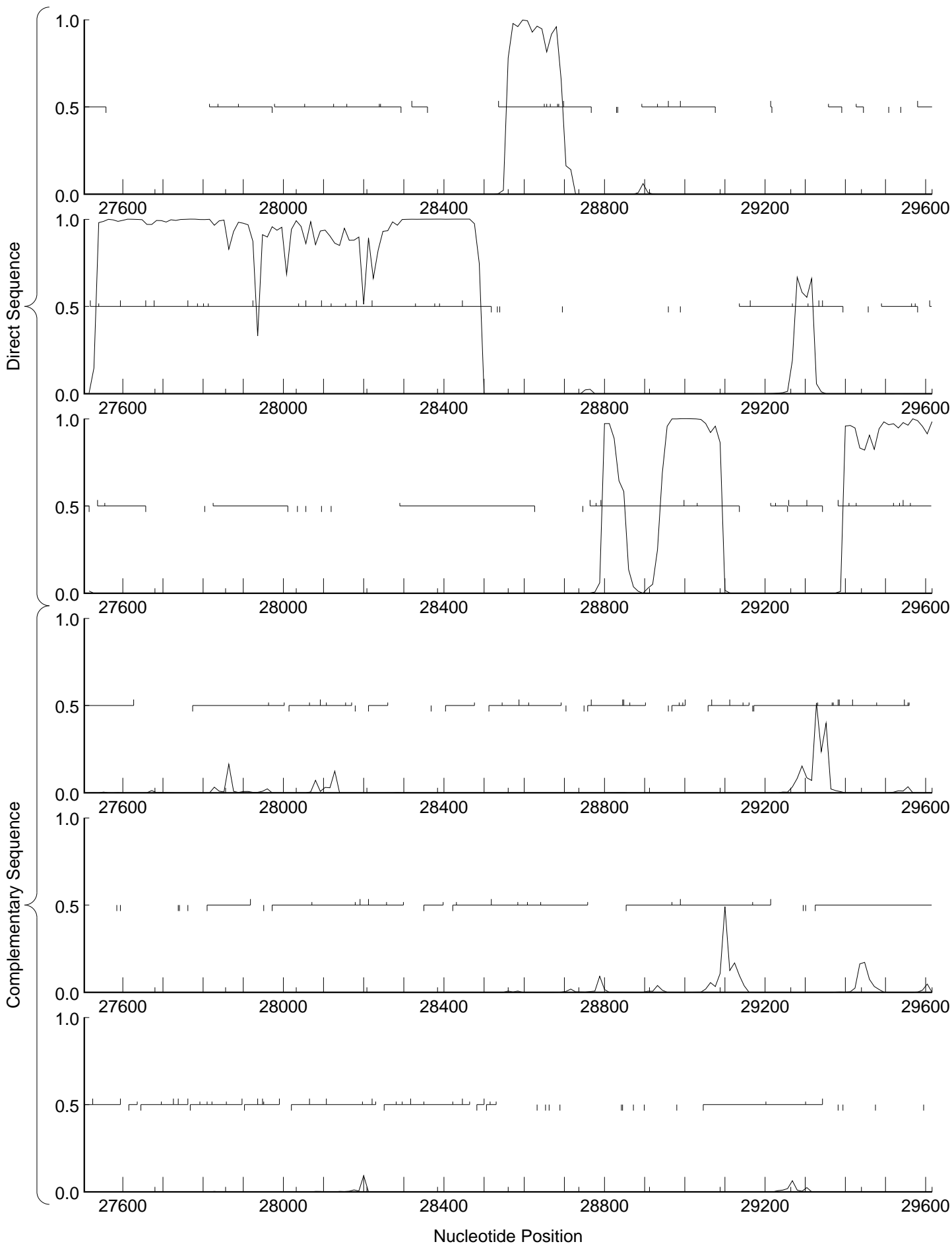


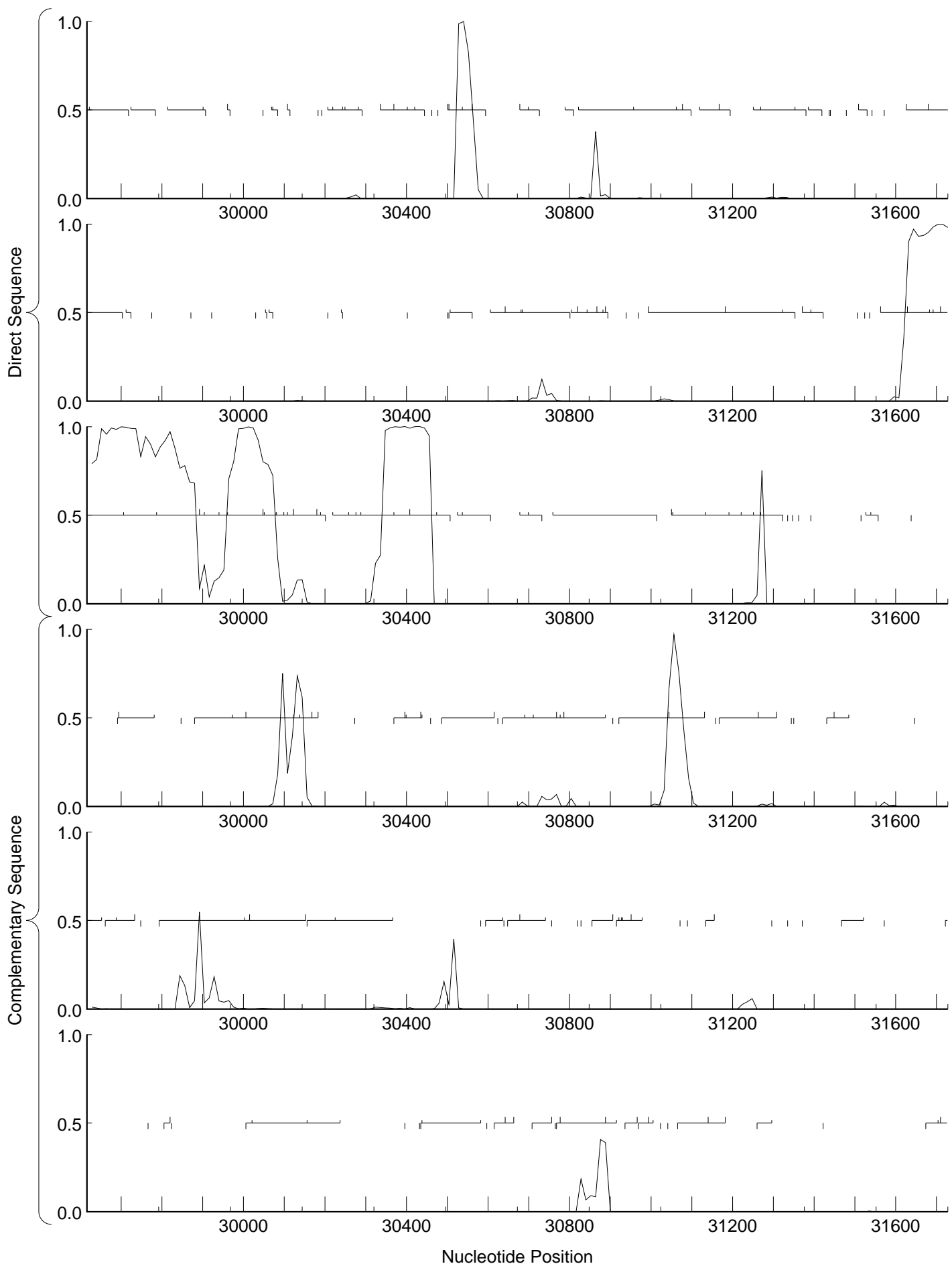
complete sequence, 55767 bp including 10-base 3' overhang (CGGACGGCGC), Cluster F1, Order 4, Window 96, Step 12, 13/28



complete sequence, 55767 bp including 10-base 3' overhang (CGGACGGCGC), Cluster F1, Order 4, Window 96, Step 12, 14/28

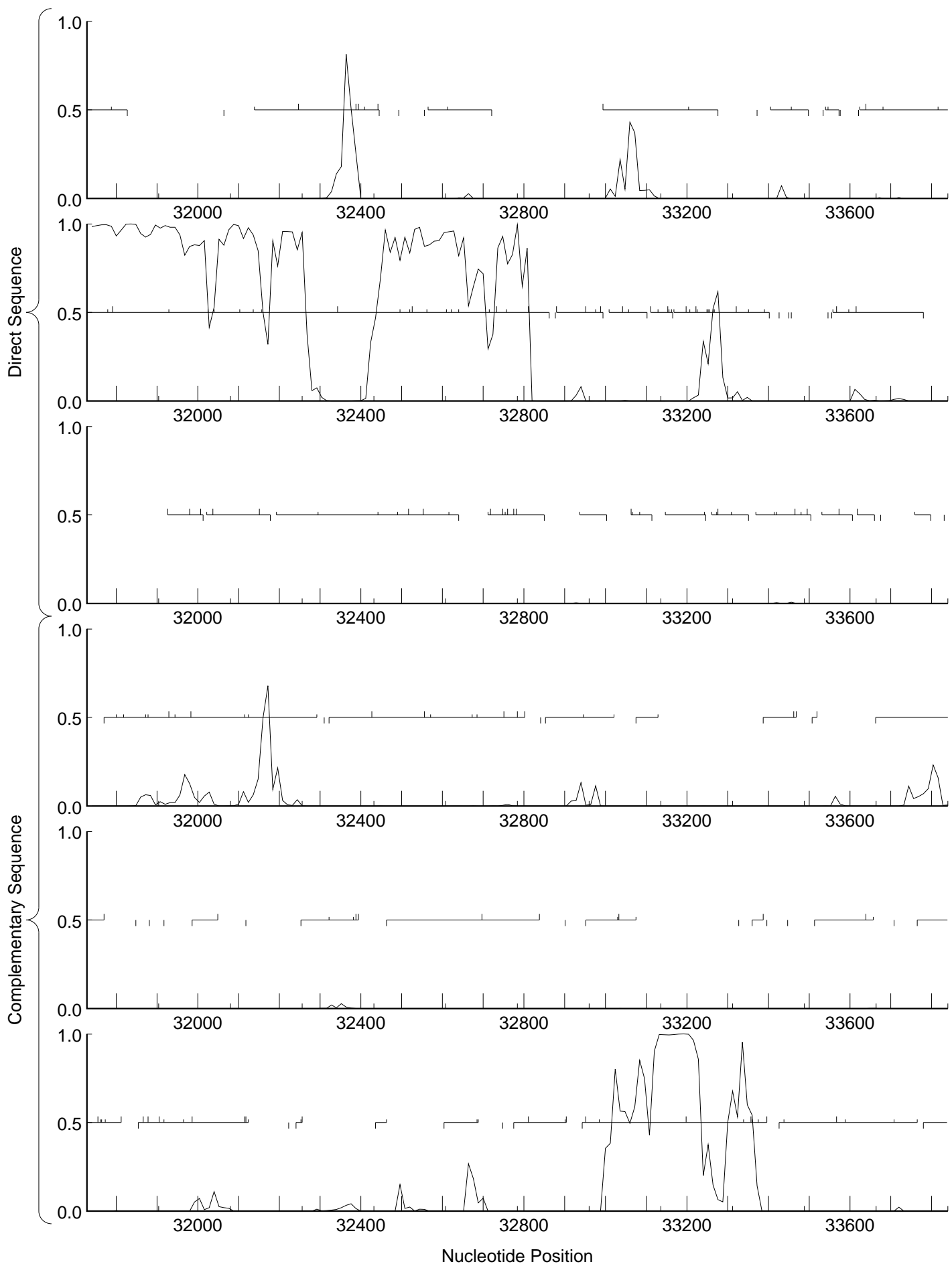




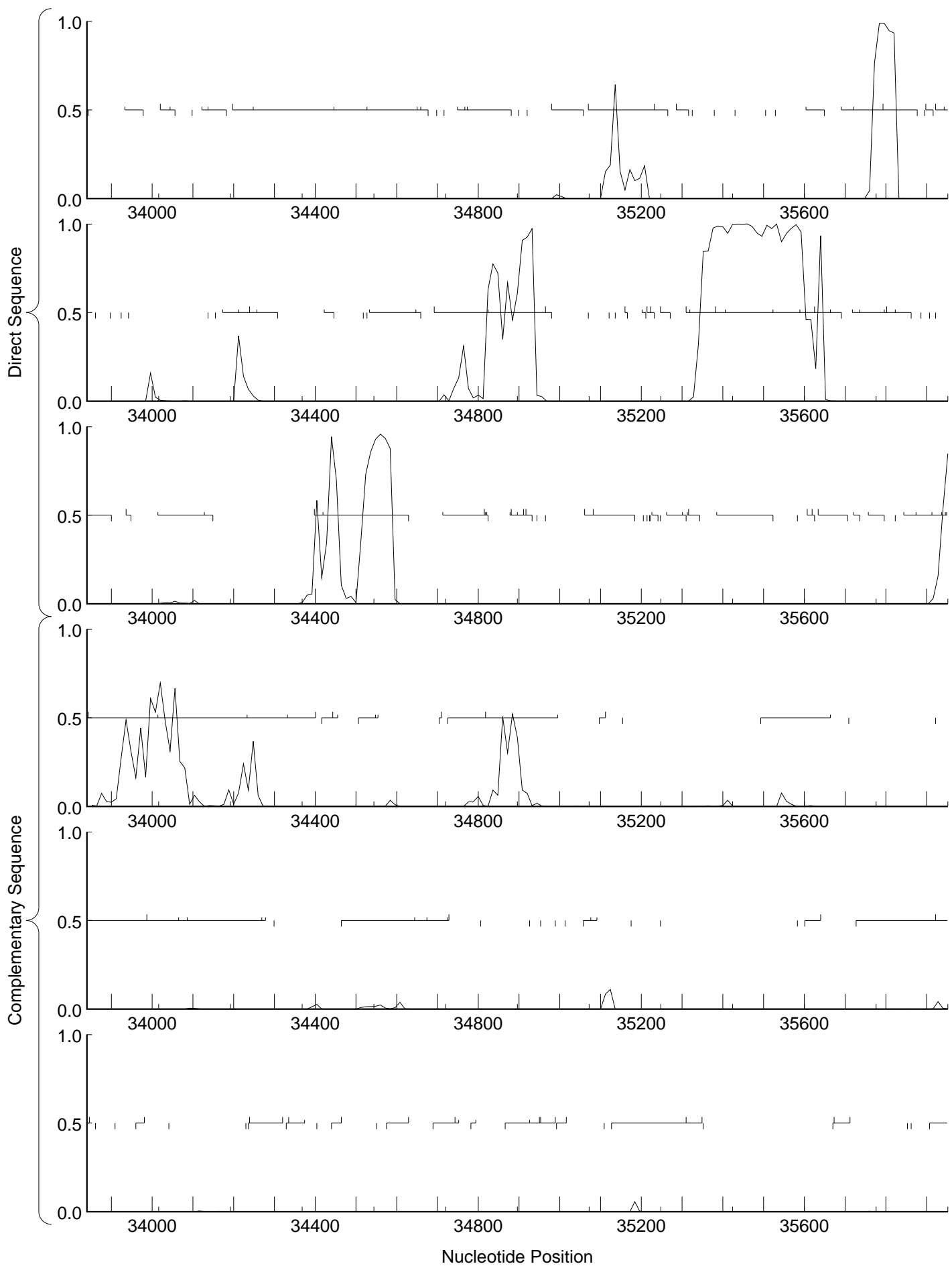




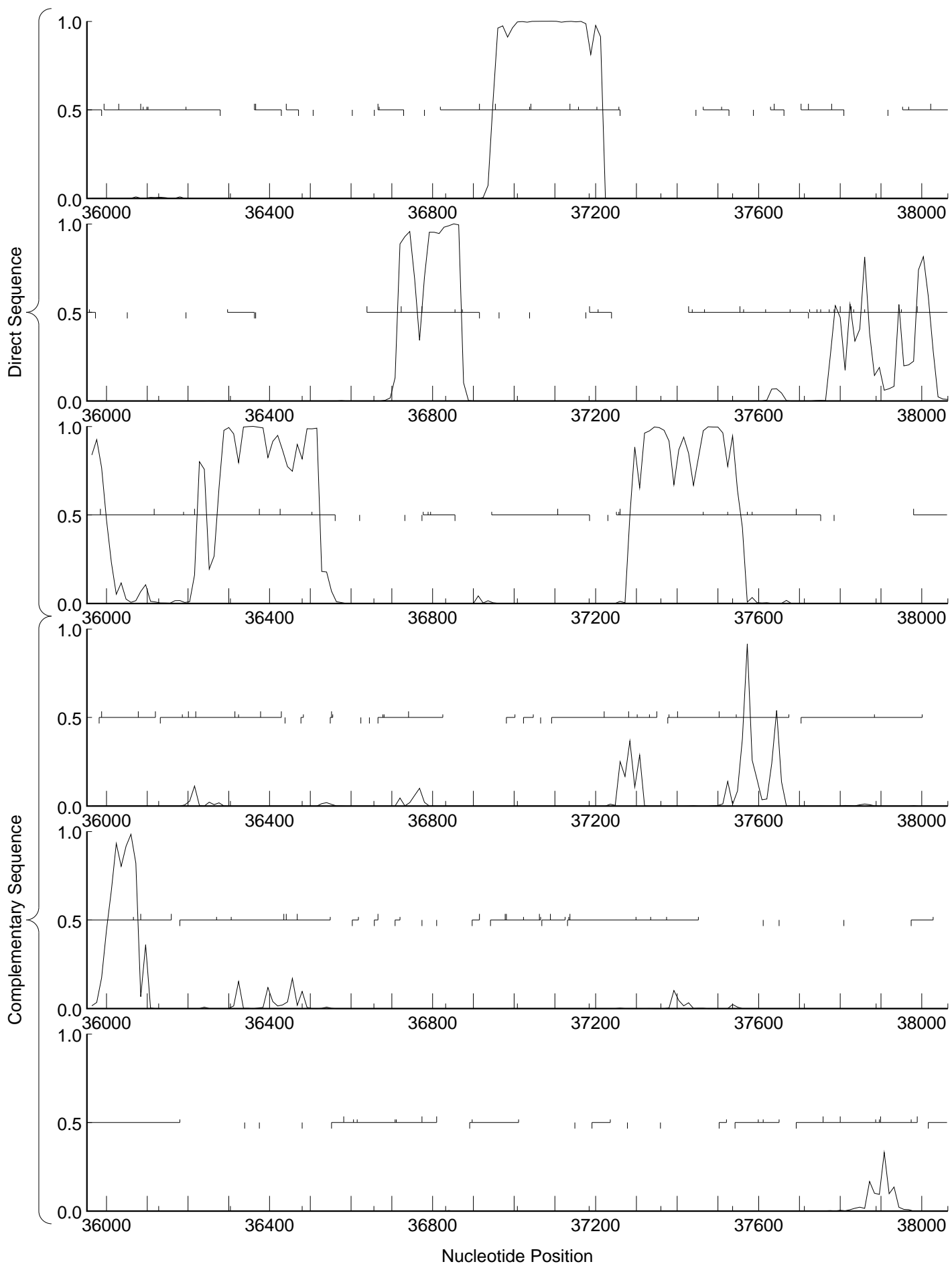
complete sequence, 55767 bp including 10-base 3' overhang (CGGACGGCGC), Cluster F1, Order 4, Window 96, Step 12, 17/28

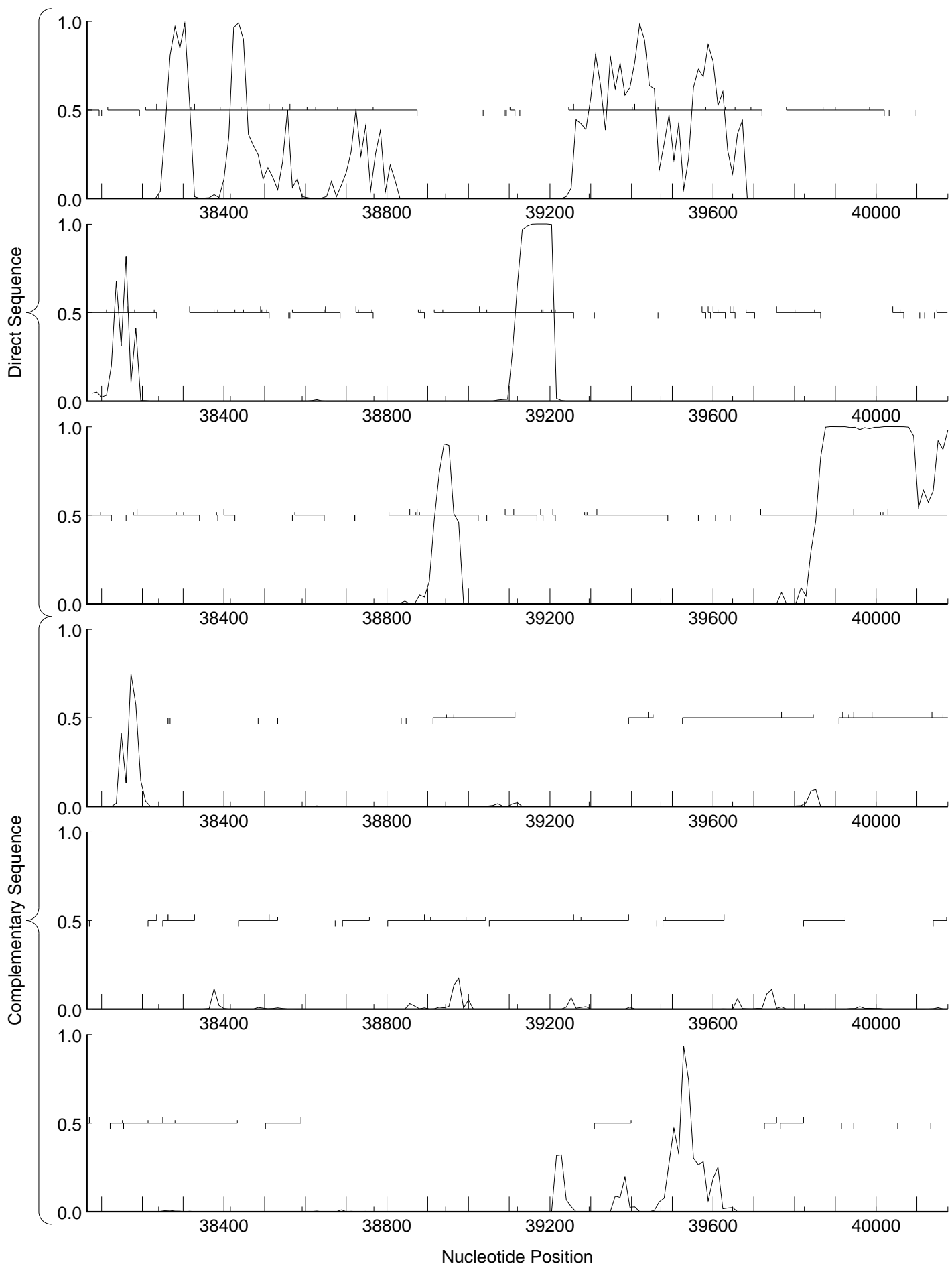


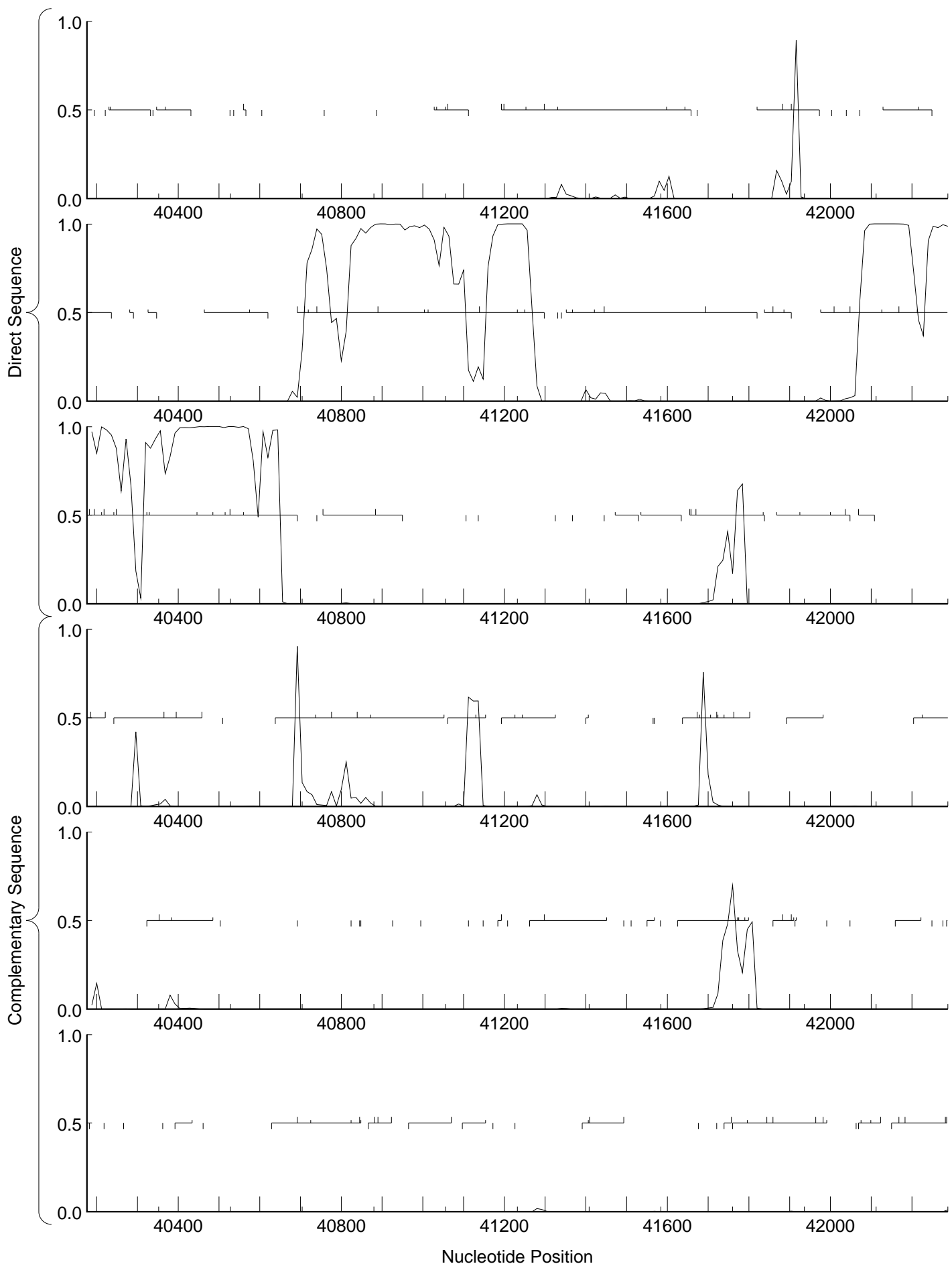
complete sequence, 55767 bp including 10-base 3' overhang (CGGACGGCGC), Cluster F1, Order 4, Window 96, Step 12, 18/28

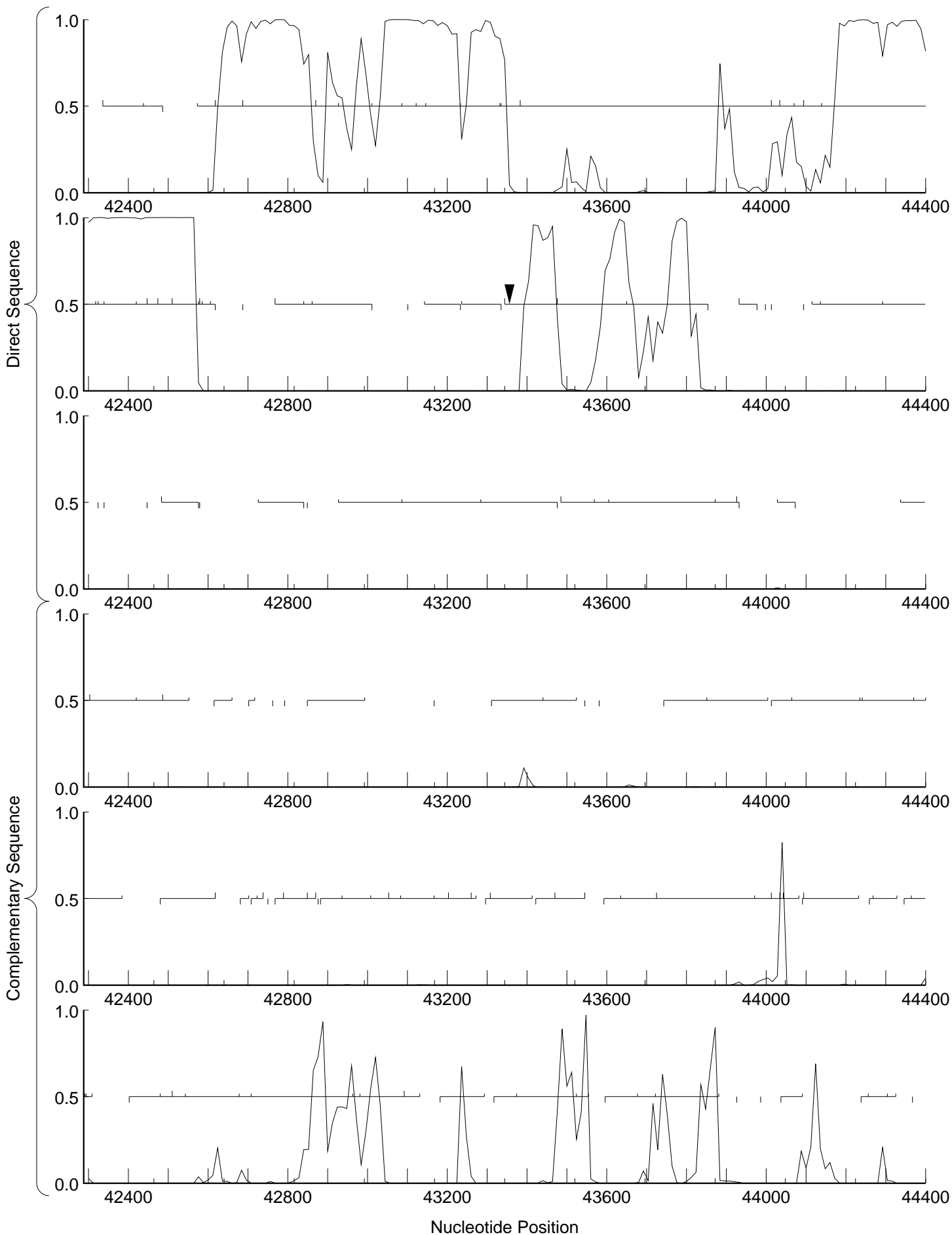


complete sequence, 55767 bp including 10-base 3' overhang (CGGACGGCGC), Cluster F1, Order 4, Window 96, Step 12, 19/28









complete sequence, 55767 bp including 10-base 3' overhang (CGGACGGCGC), Cluster F1, Order 4, Window 96, Step 12, 23/28

