

# GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

## PROGRAM INFORMATION

Sequence : Mycobacterium phage Moostard complete sequence, 69480 bp including 10-base 3' overhang (TCG  
Analysis Date : 2/1/23 at 20:07:58  
Pages : 34  
Sequence Length : 69480 bp  
GC Content : 59.13%

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,

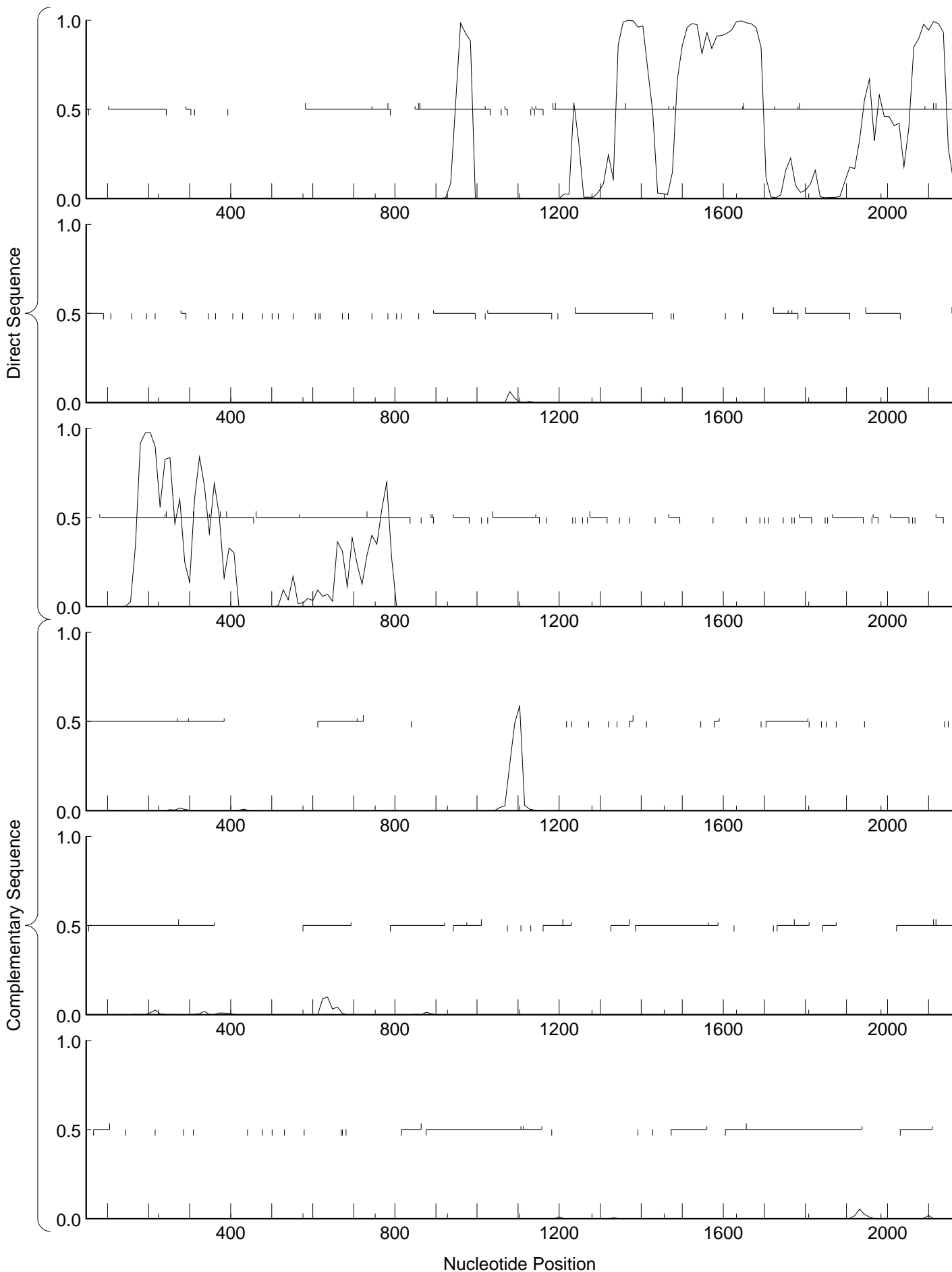
## MATRIX INFORMATION

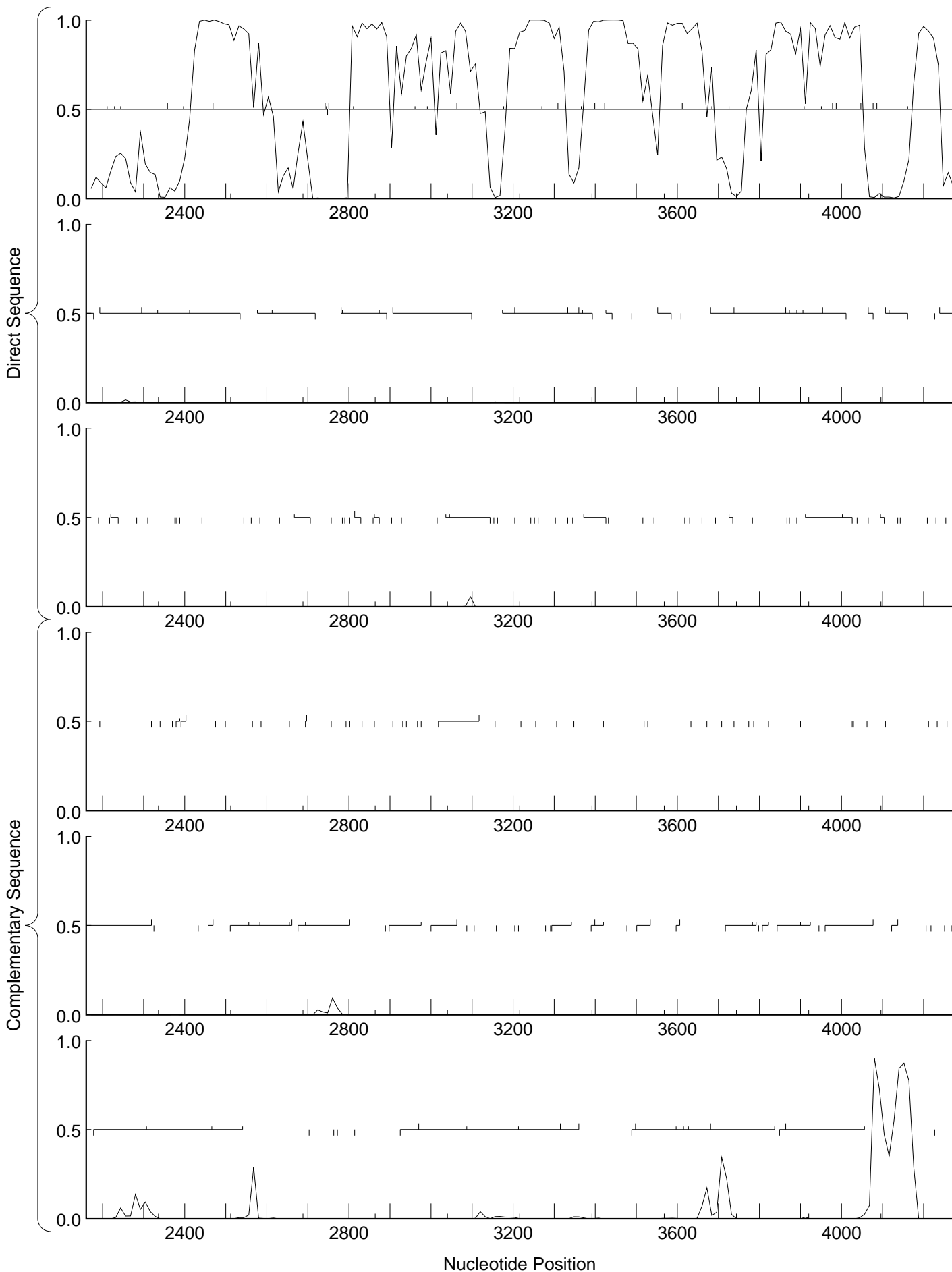
Matrix : Mycobacterium\_tuberculosis\_H37Rv  
Author : -  
Order : 4

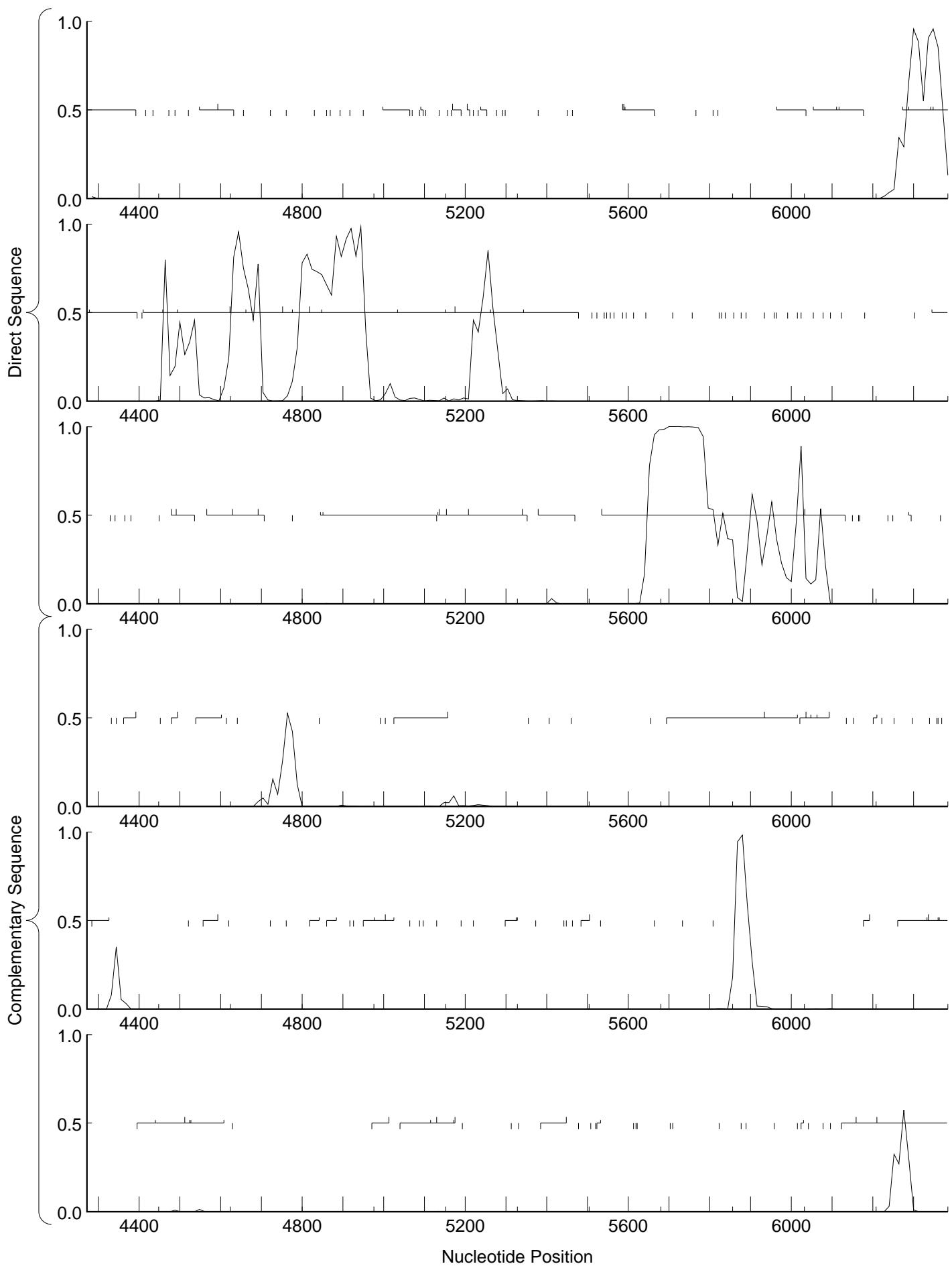
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

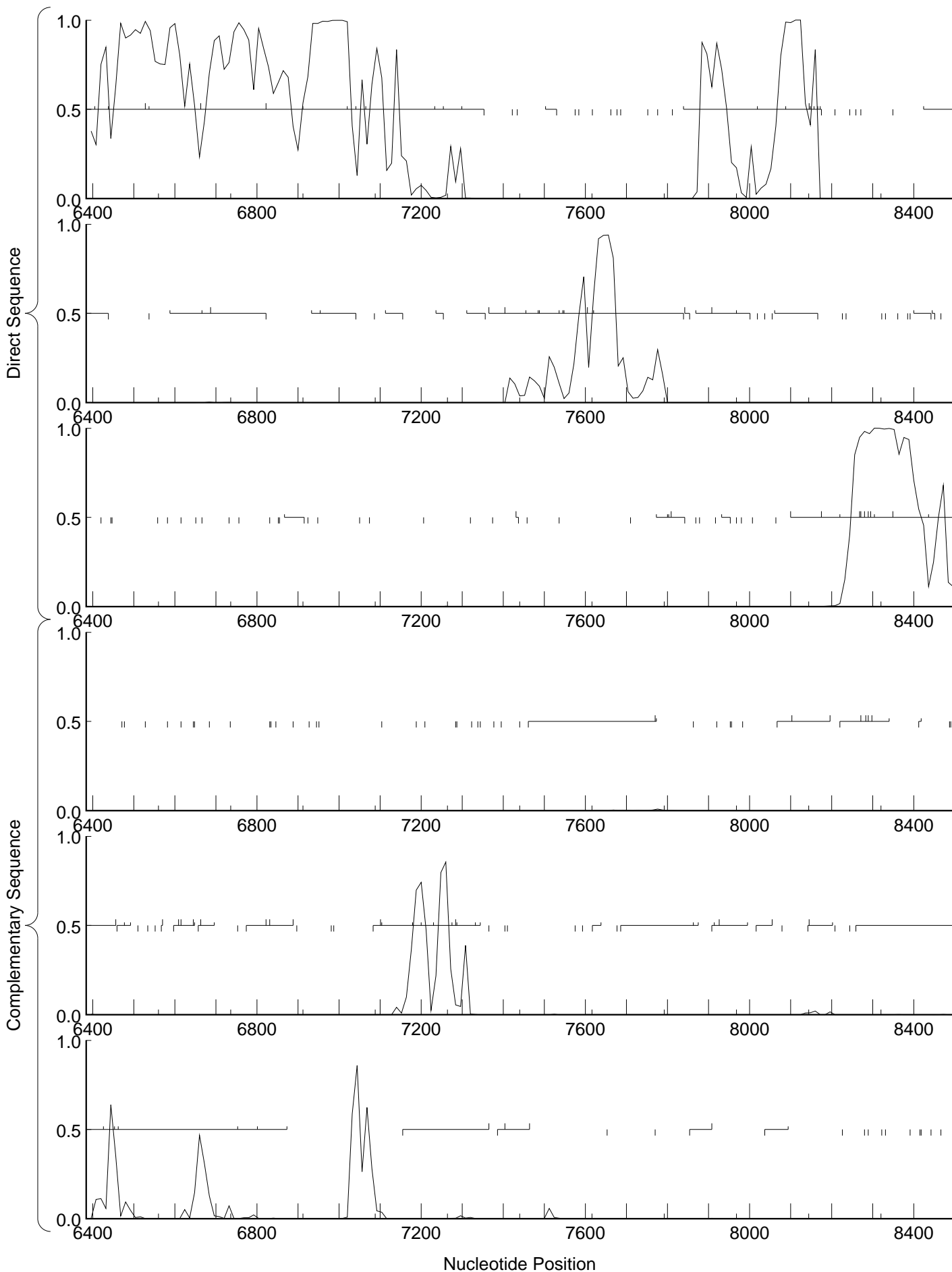
### Matrix notes & comments

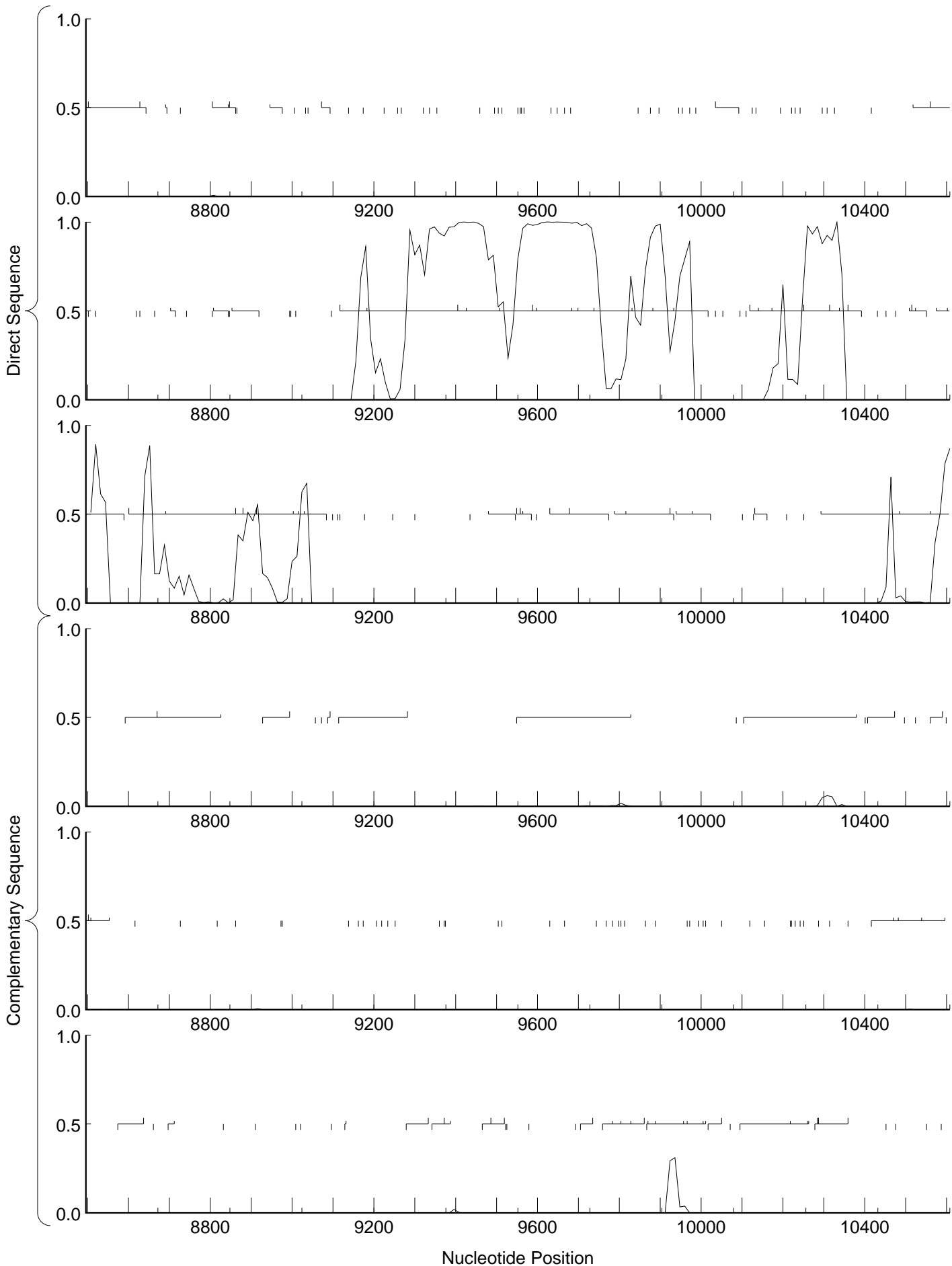
Training set derived by GeneMarkS, 4.27 September 2014  
Tue Sep 23 15:14:37 2014



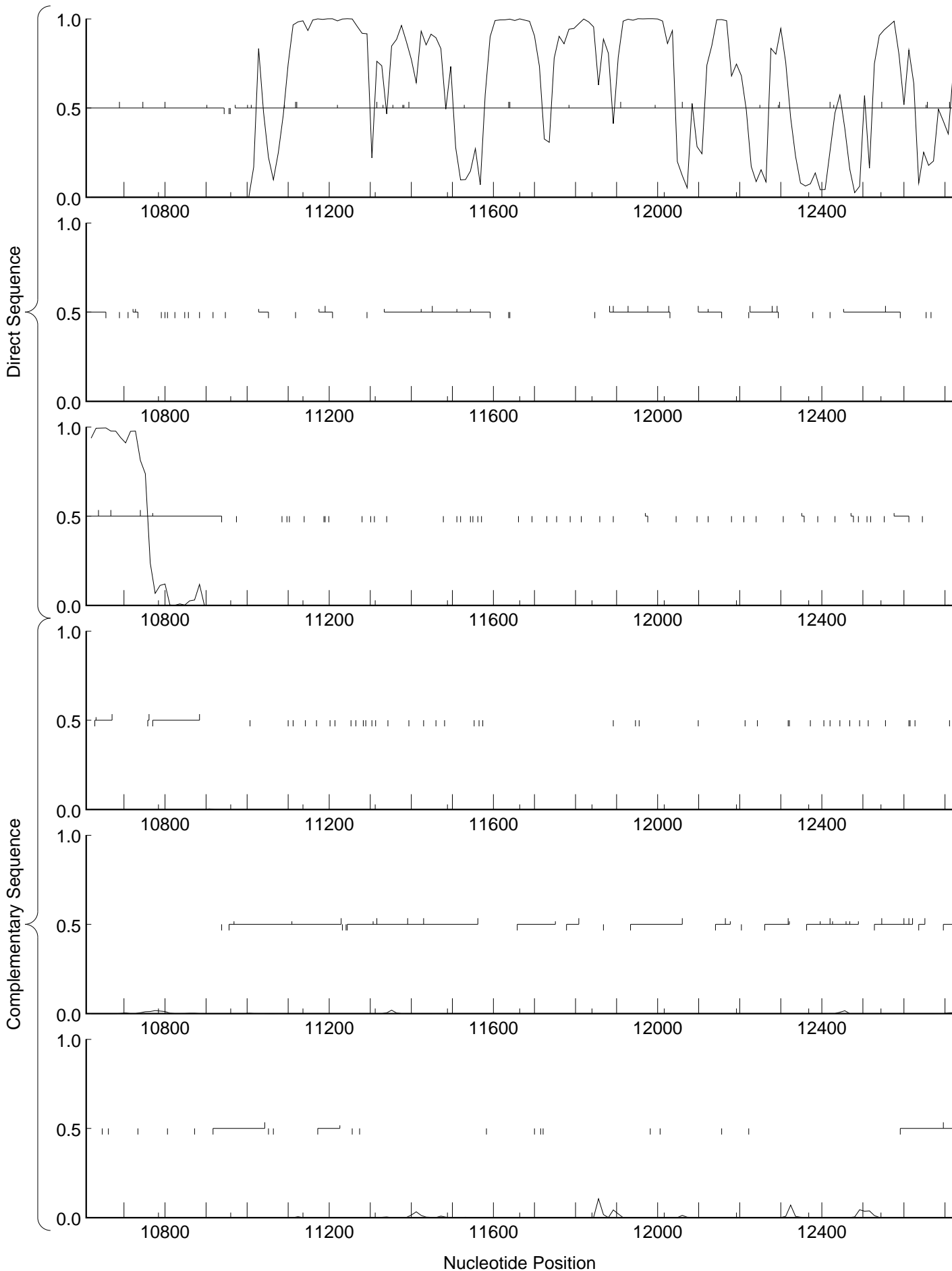


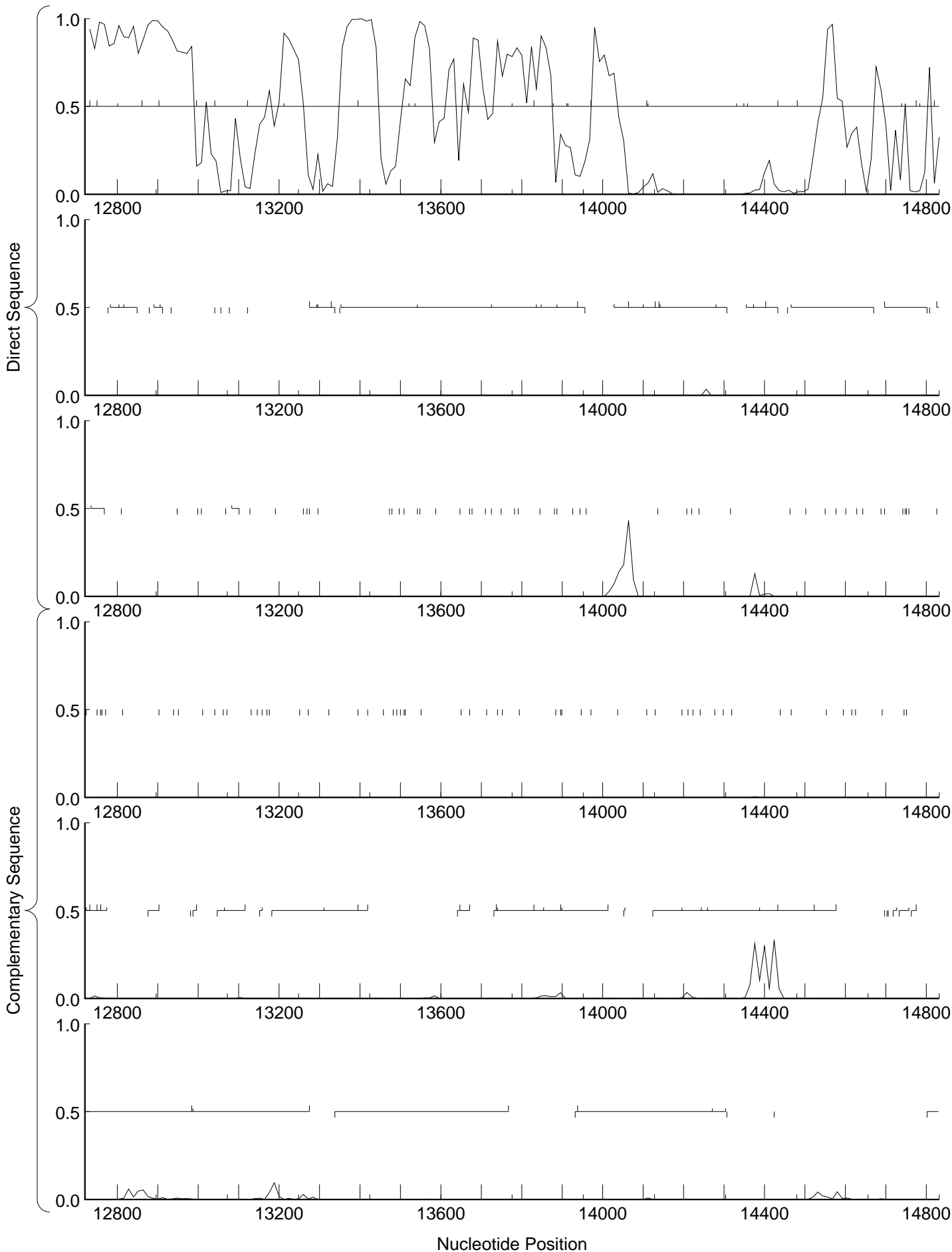




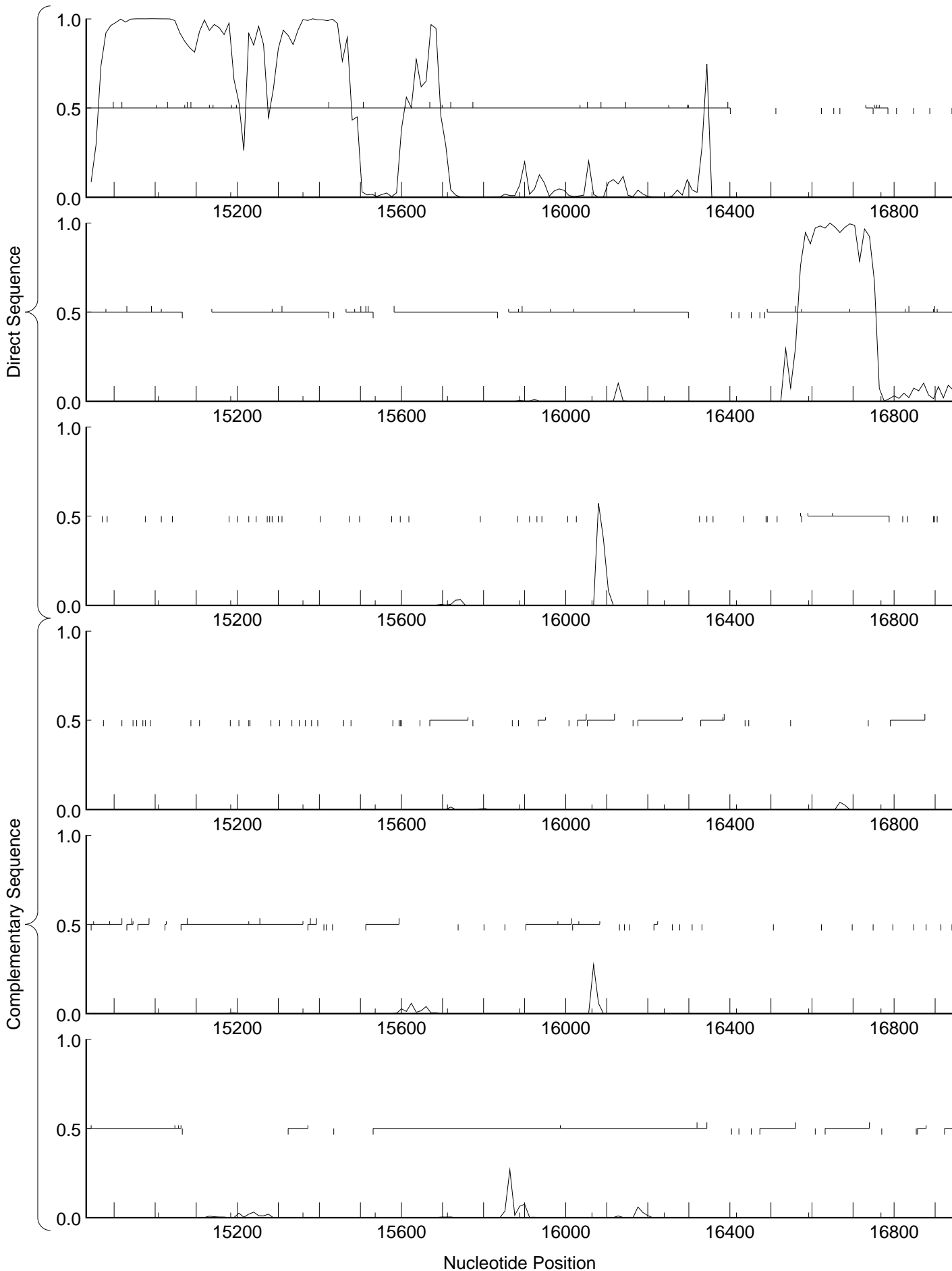


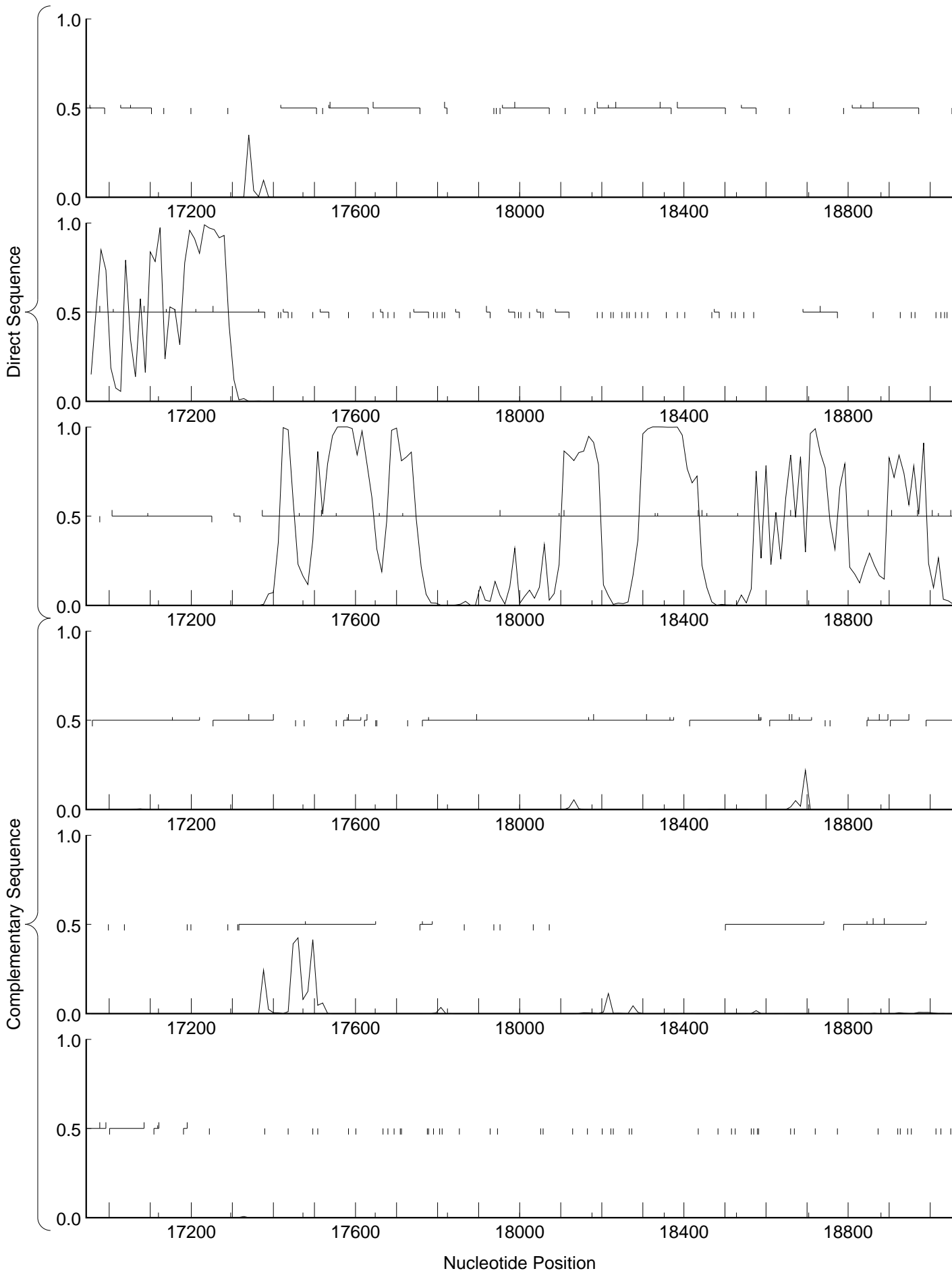
rd complete sequence, 69480 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L3, Order 4, Window 96, Step 12, 7/34

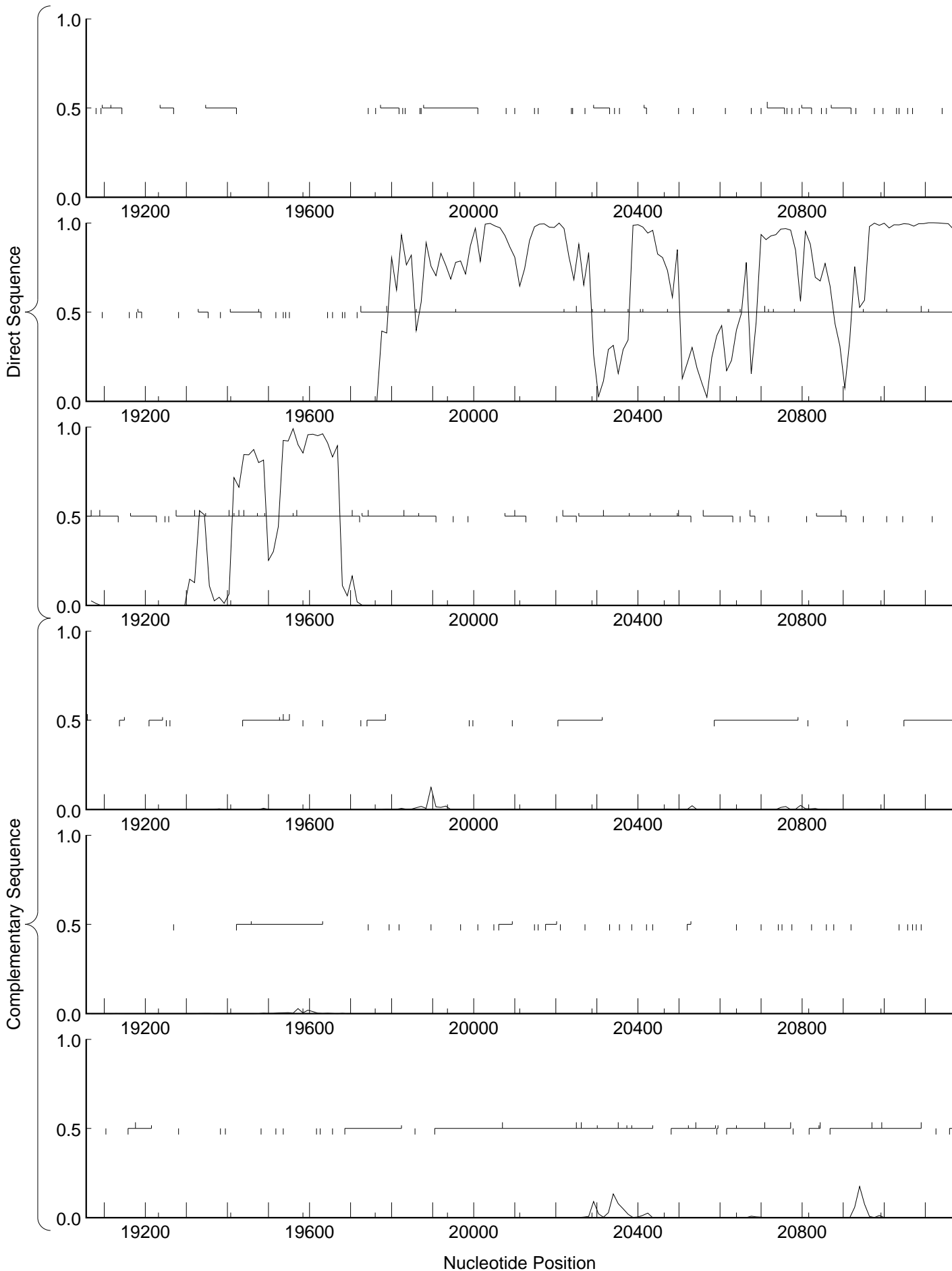


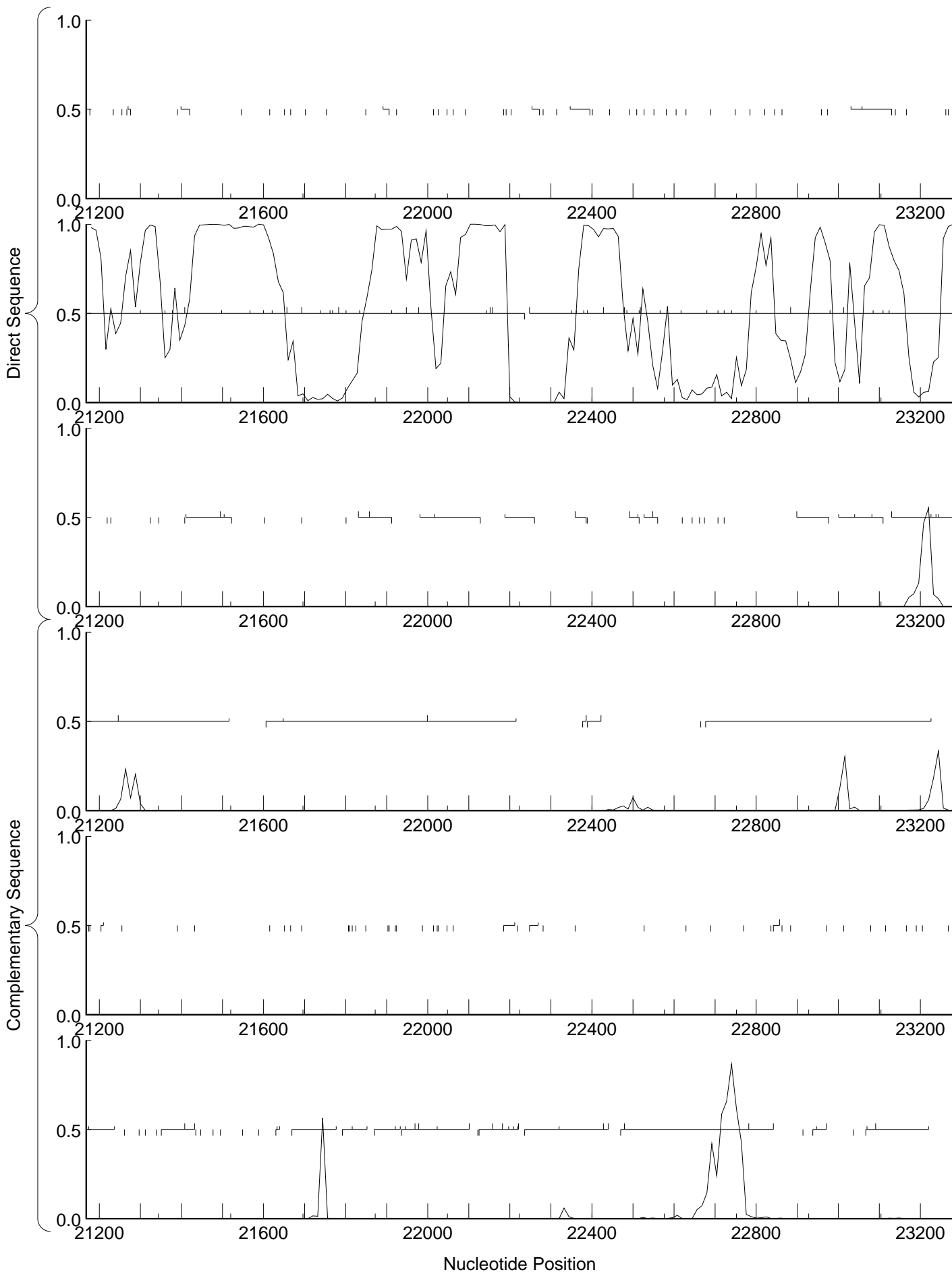


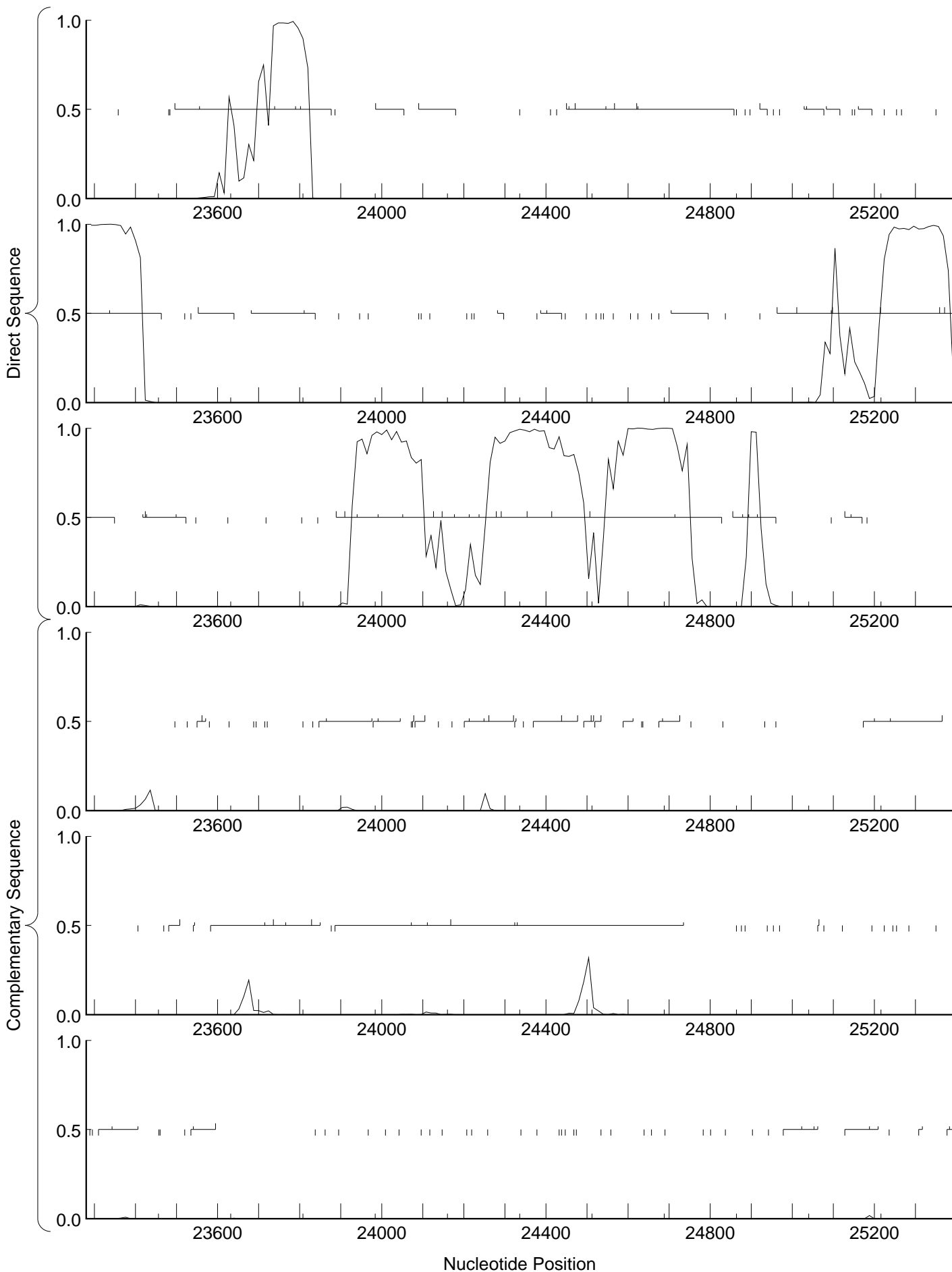


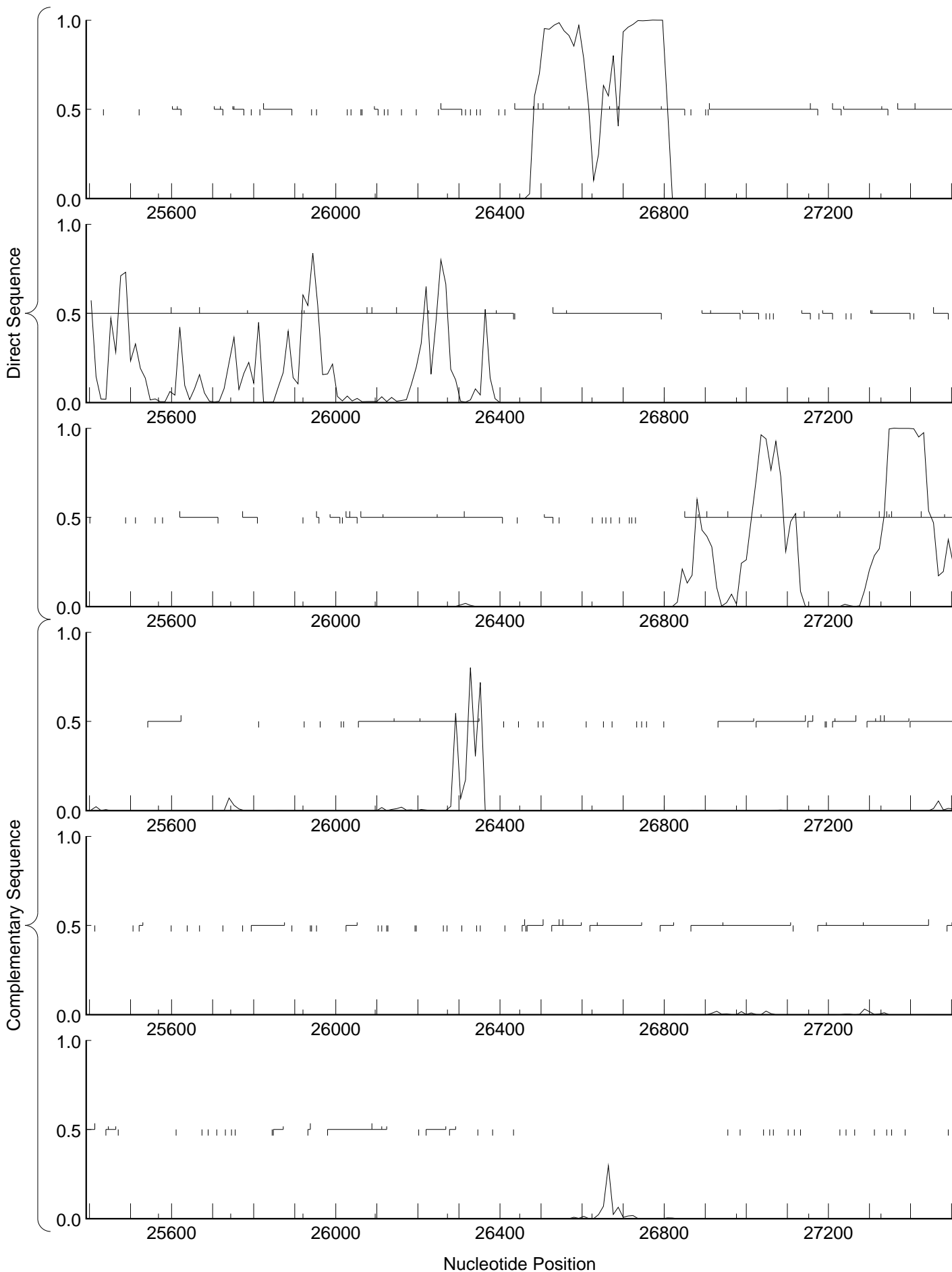


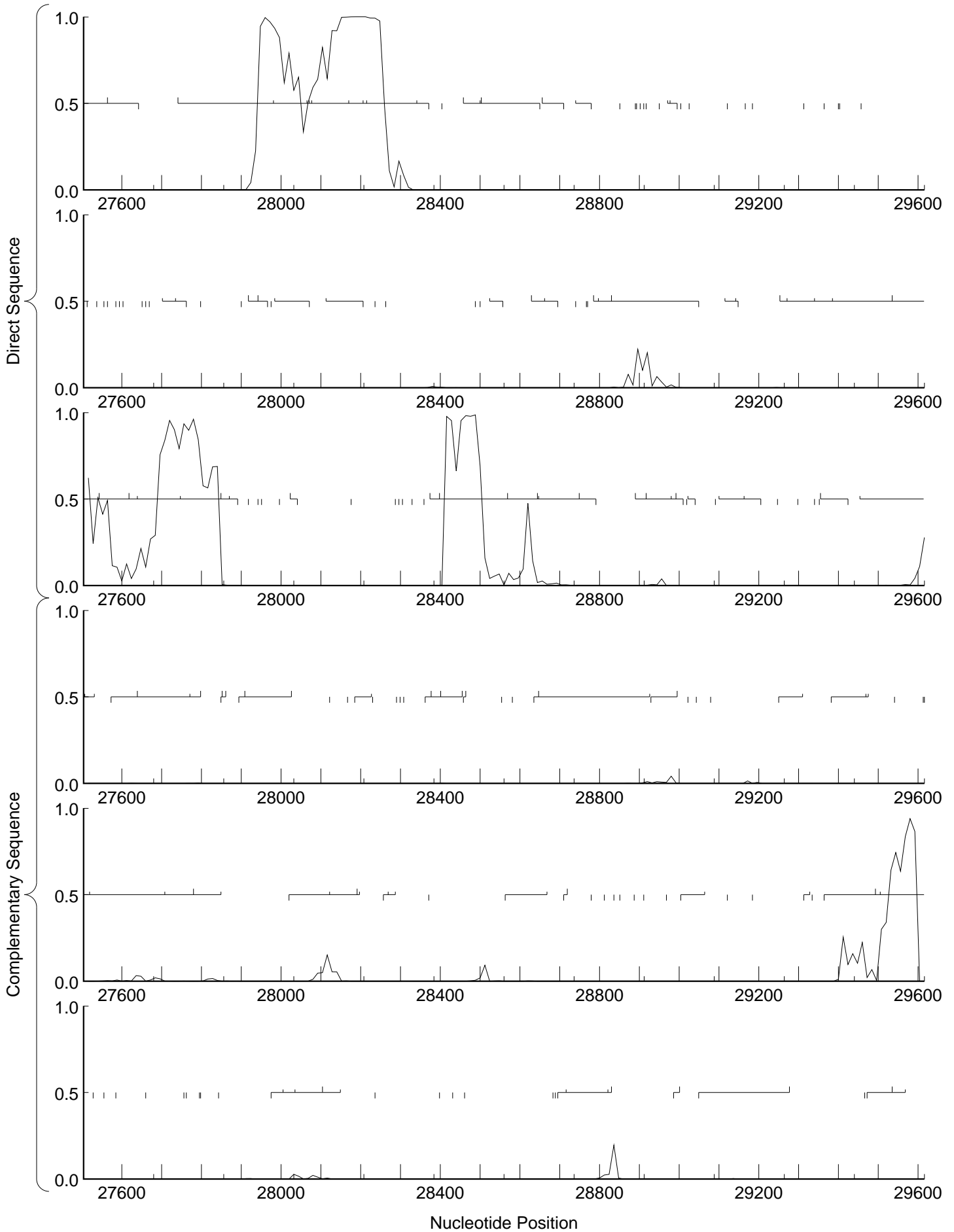


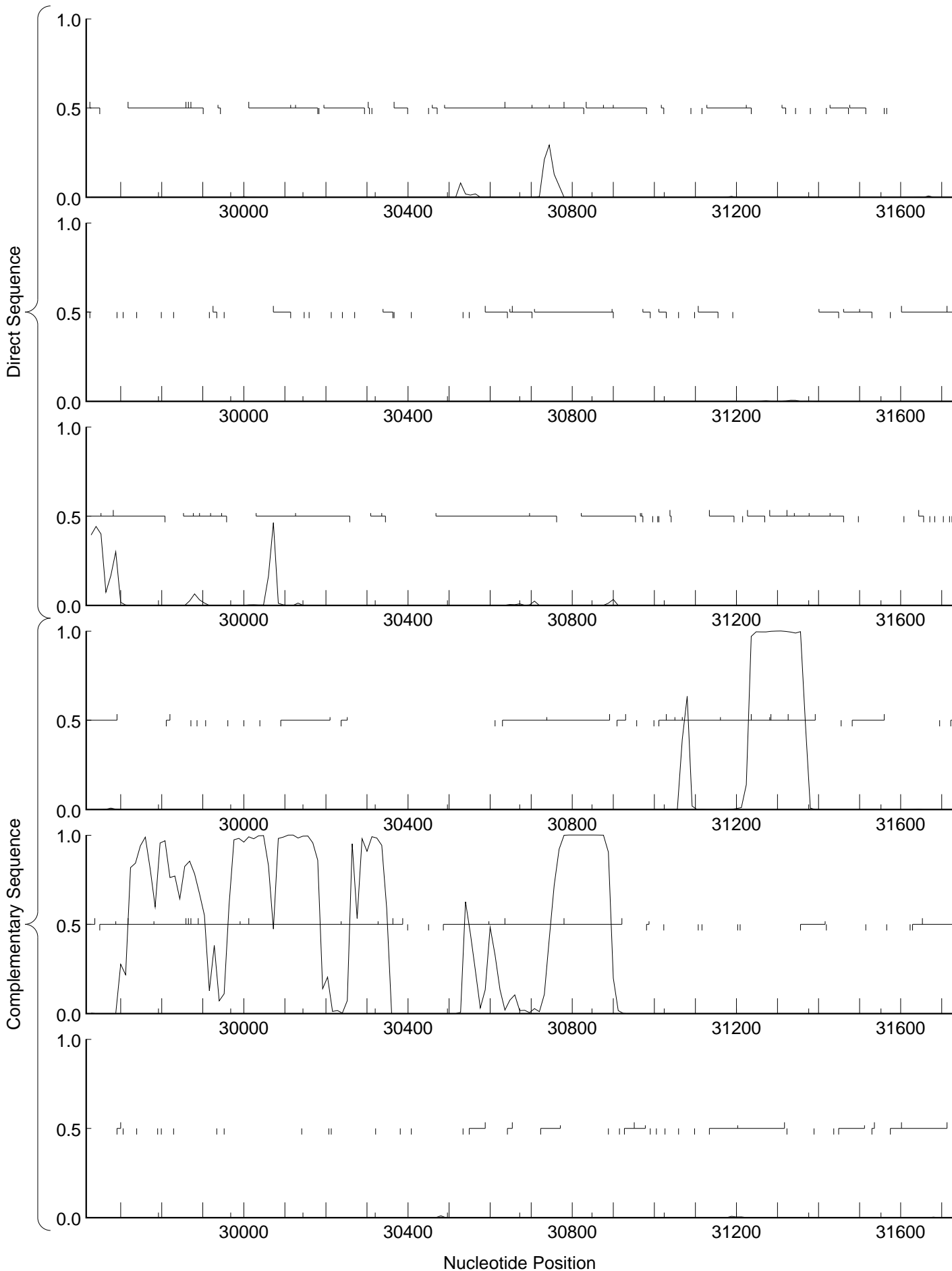














Complete sequence, 69480 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L3, Order 4, Window 96, Step 12, 17/34

