

# GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

## PROGRAM INFORMATION

Sequence : Mycobacterium phage Jamie19 complete sequence, 41278 bp including 13-base 3' overhang (CCCG)  
Analysis Date : 10/14/19 at 2:22:51  
Pages : 21  
Sequence Length : 41278 bp  
GC Content : 66.24%

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,

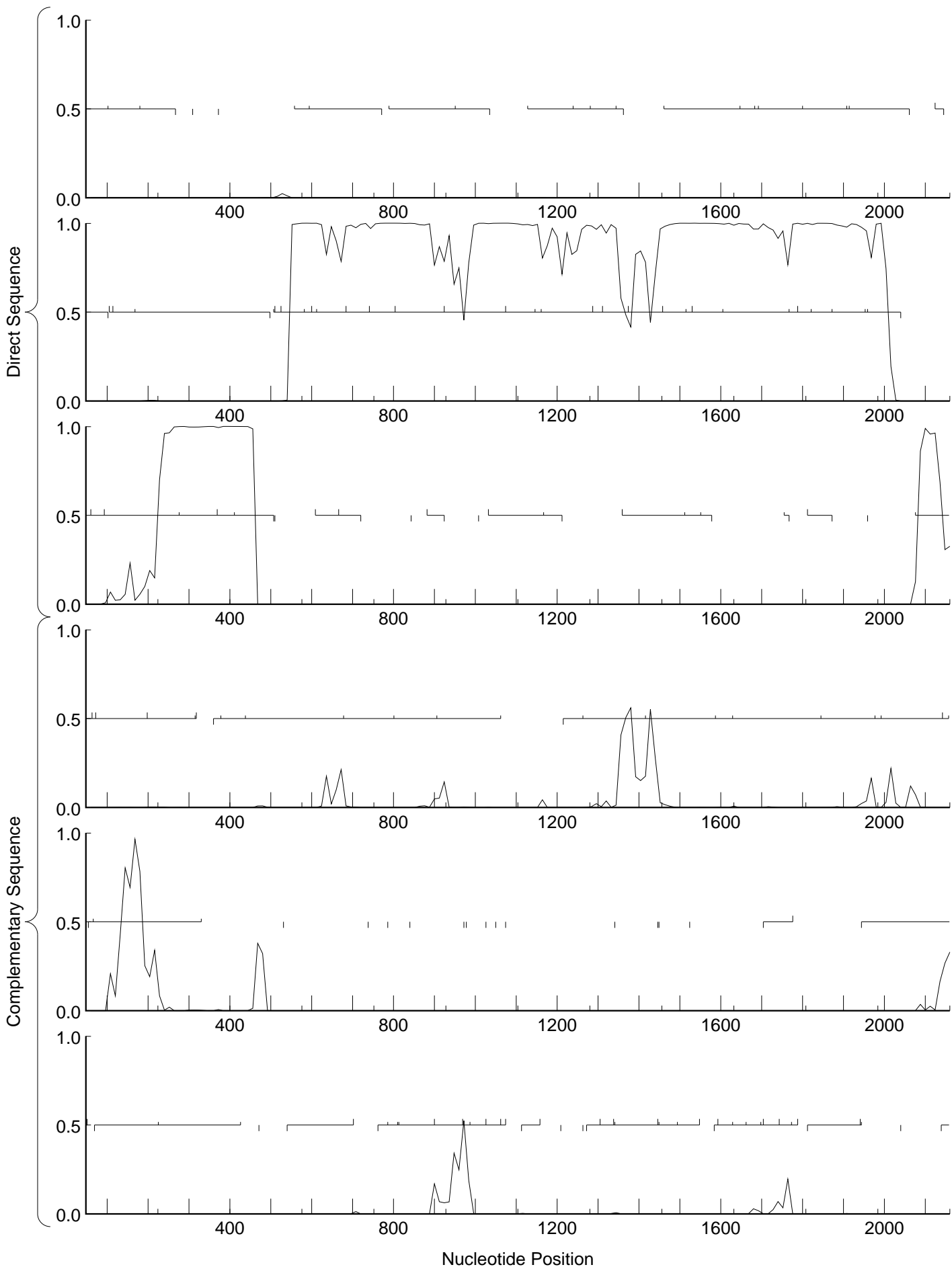
## Matrix notes & comments

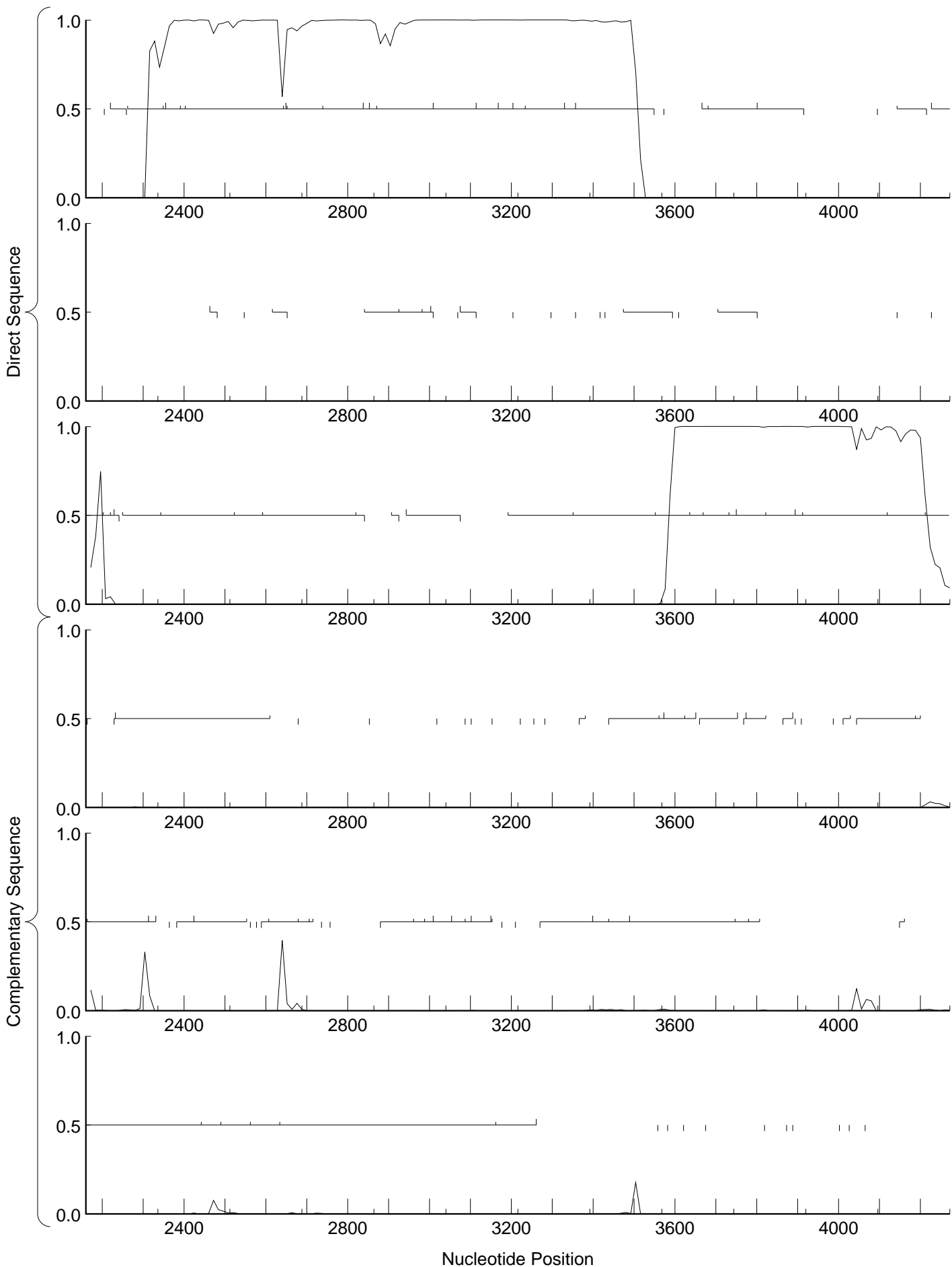
Training set derived by GeneMarkS, 4.27 September 2014  
Tue Sep 23 15:14:37 2014

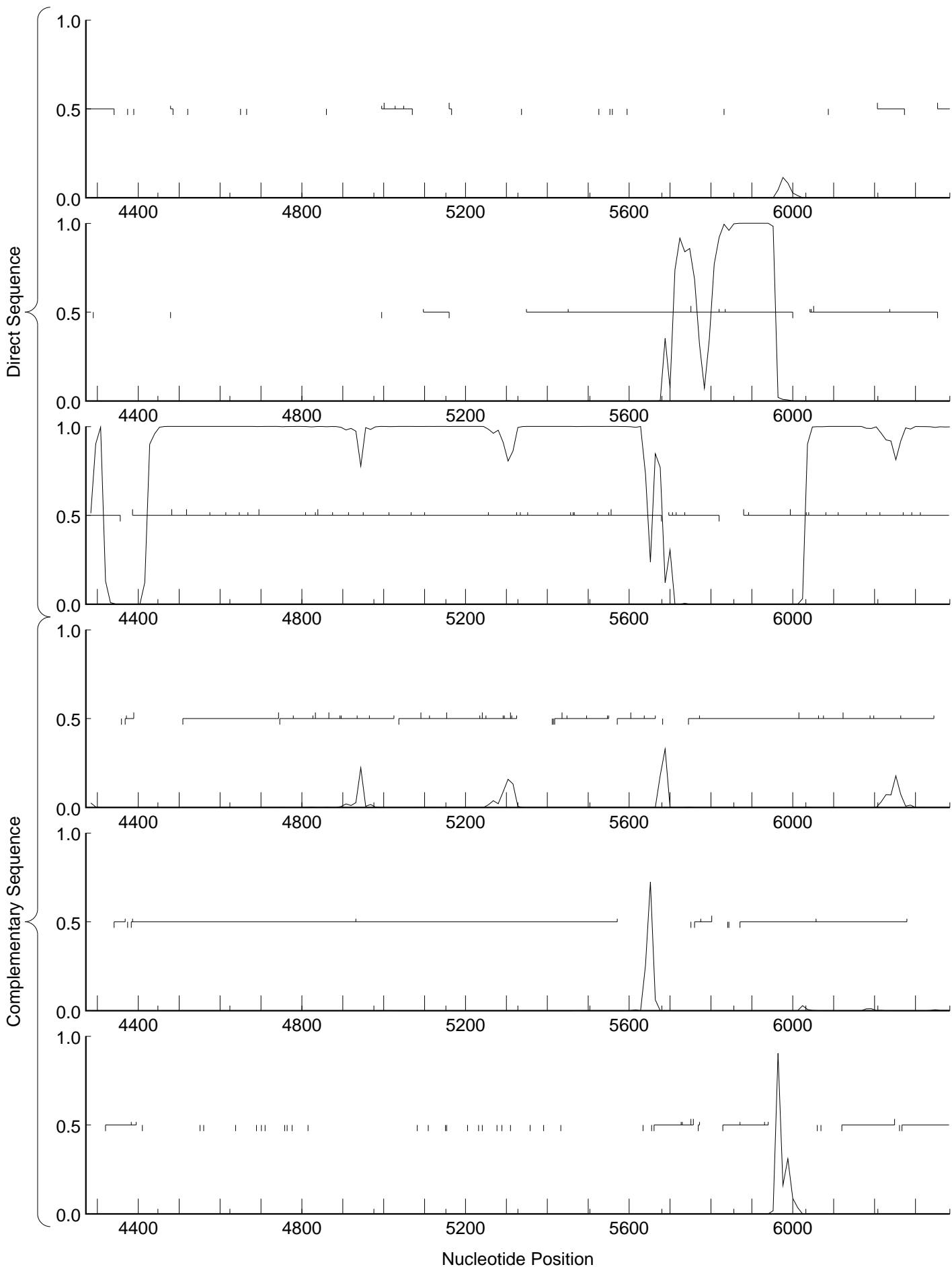
## MATRIX INFORMATION

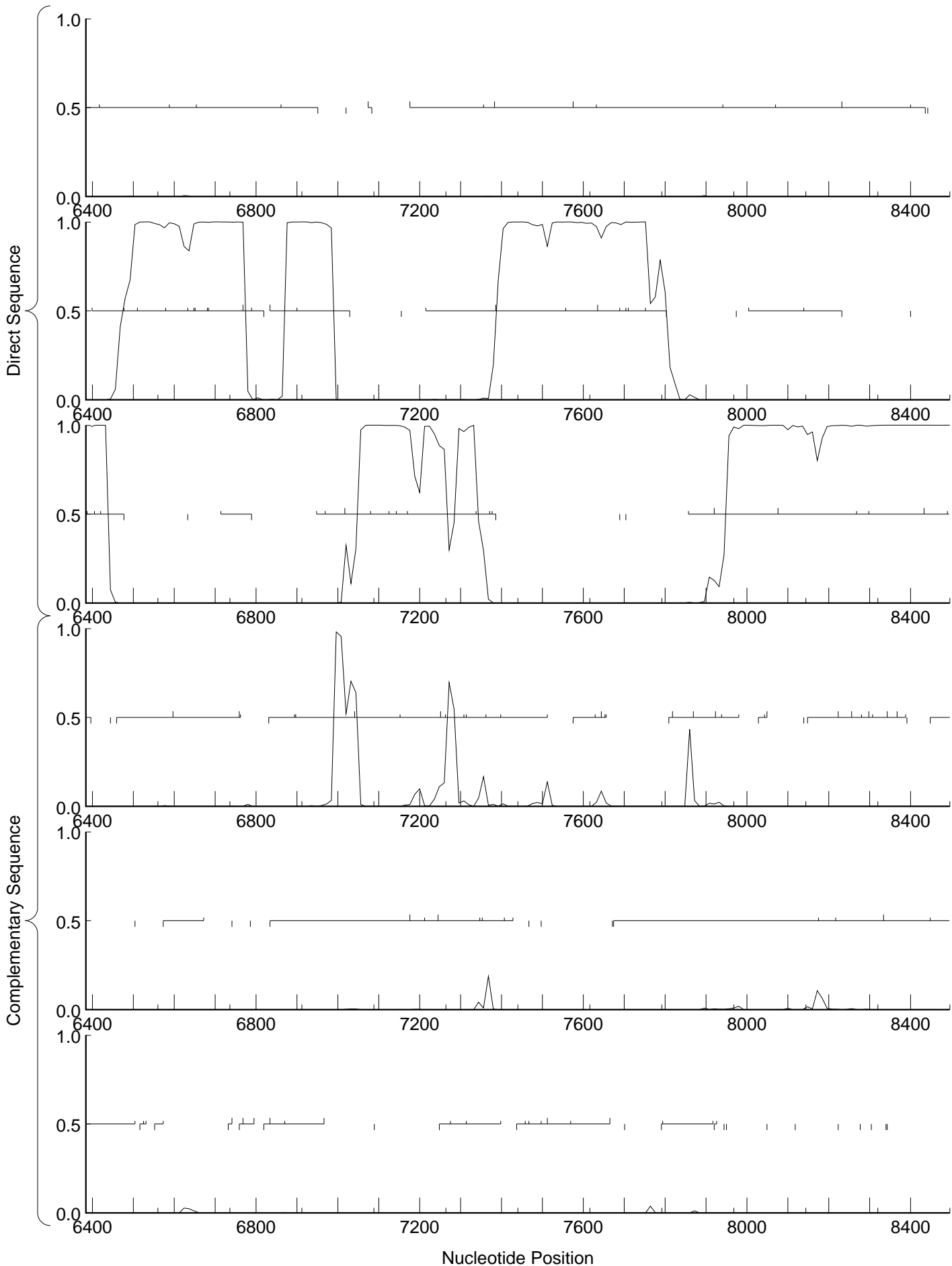
Matrix : Mycobacterium\_tuberculosis\_H37Rv  
Author : -  
Order : 4

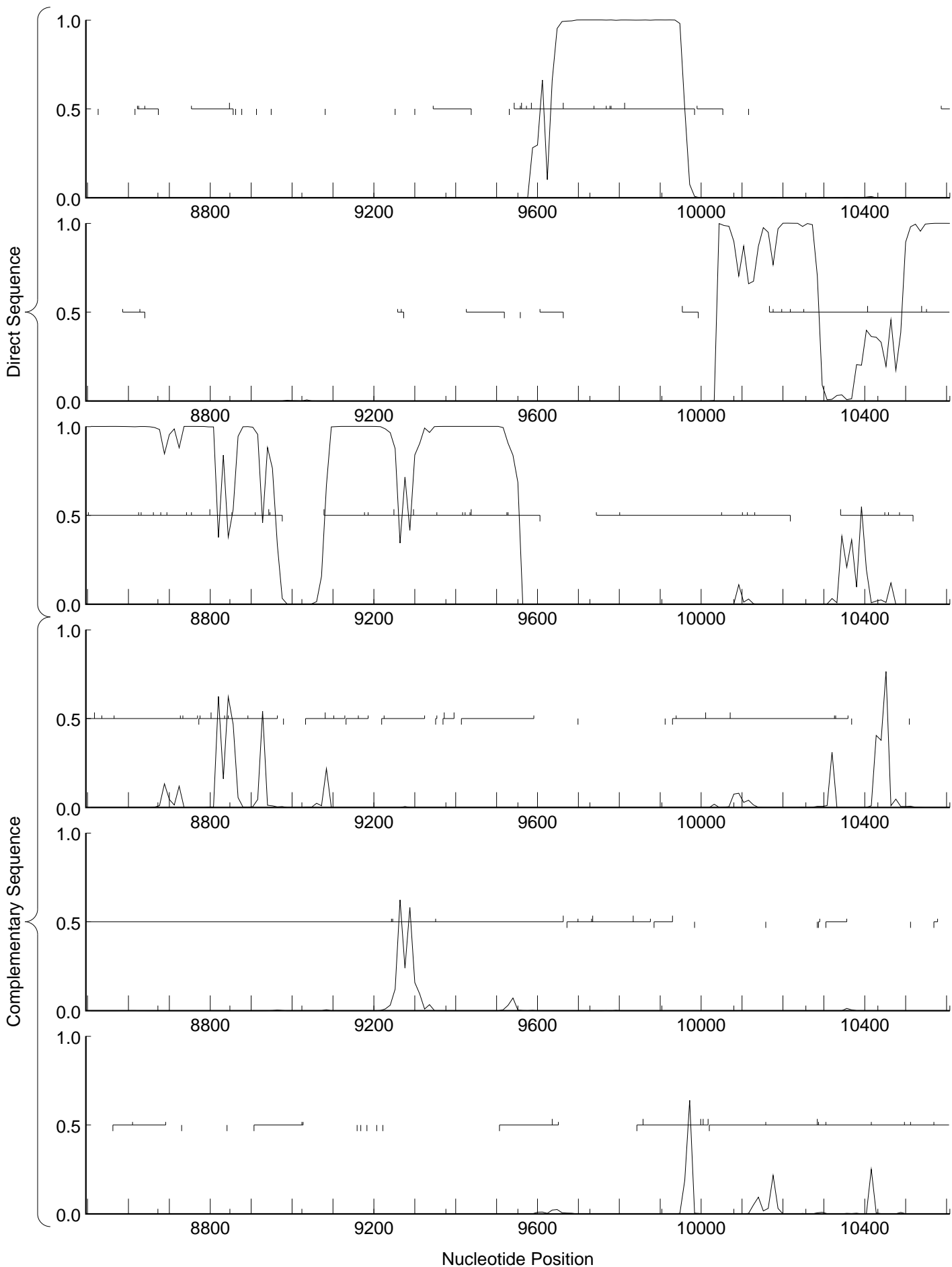
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

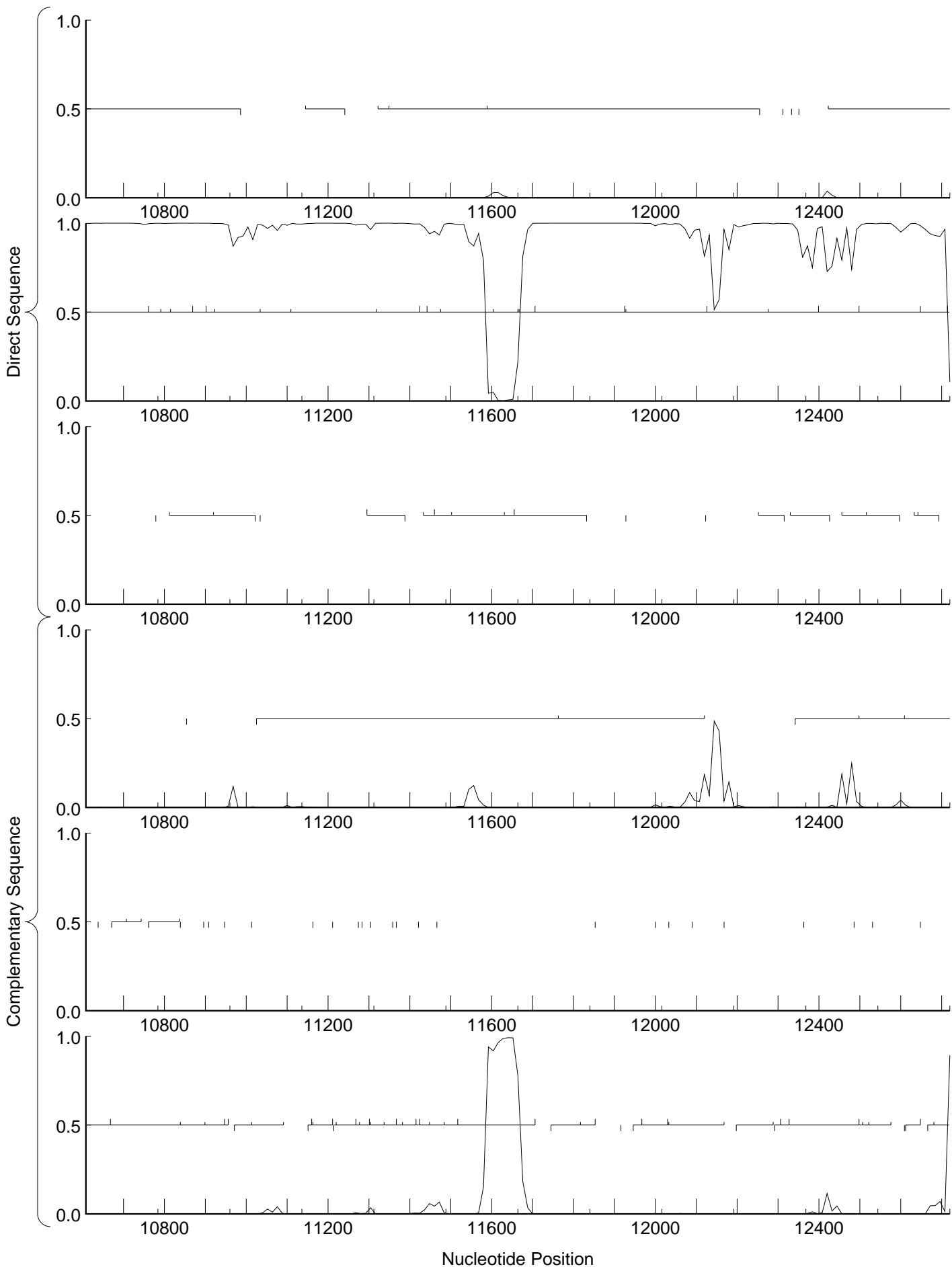


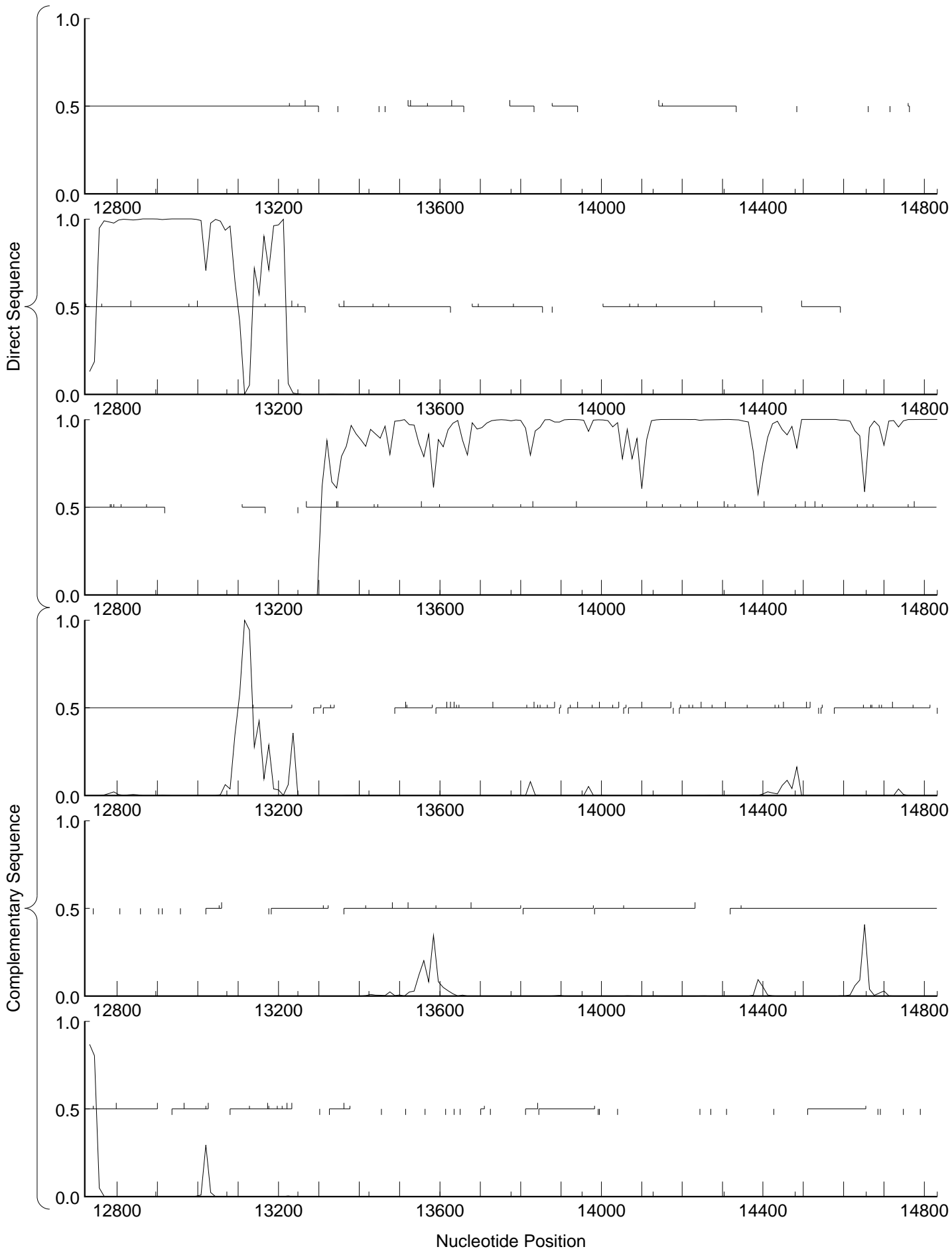




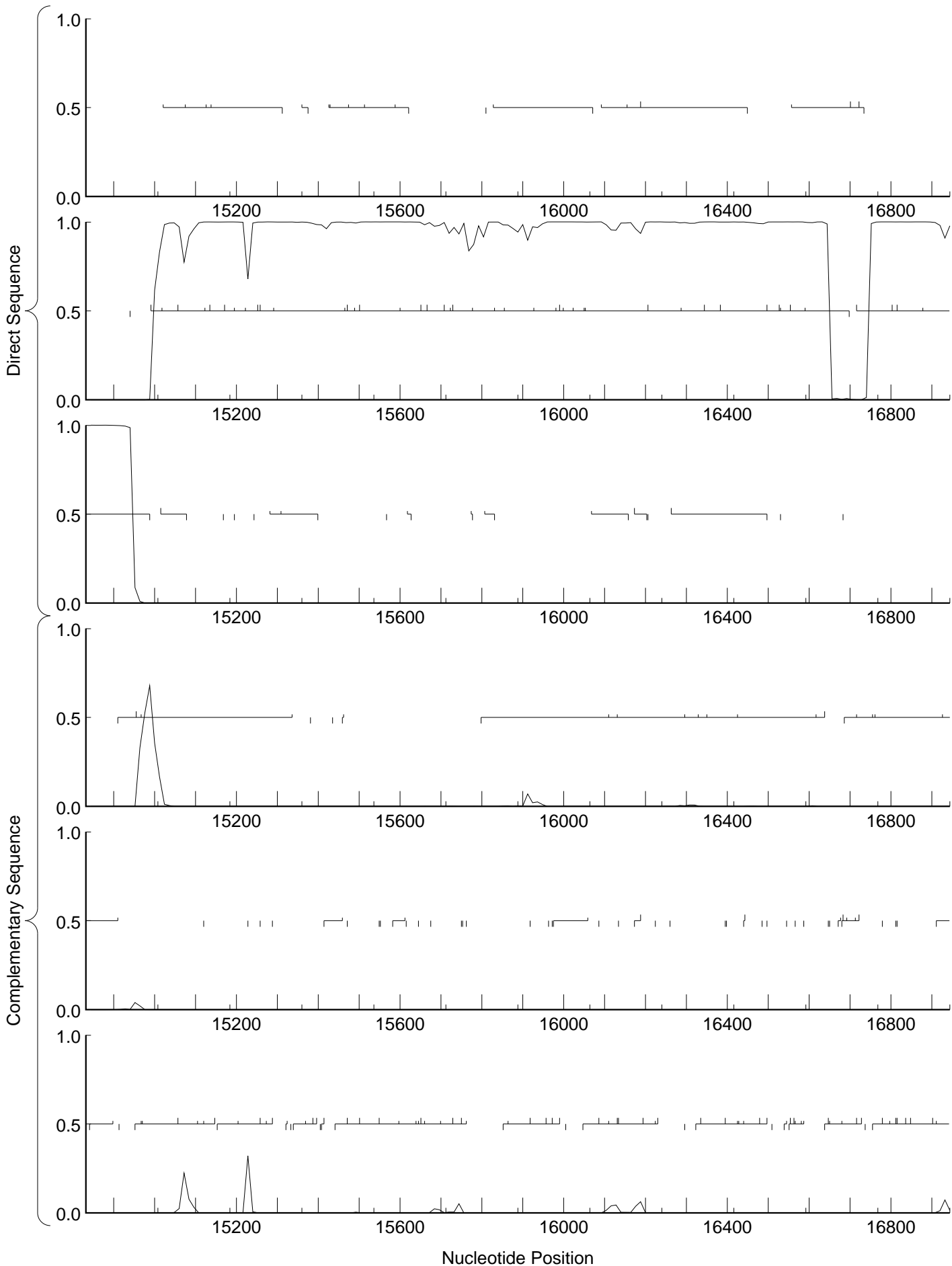


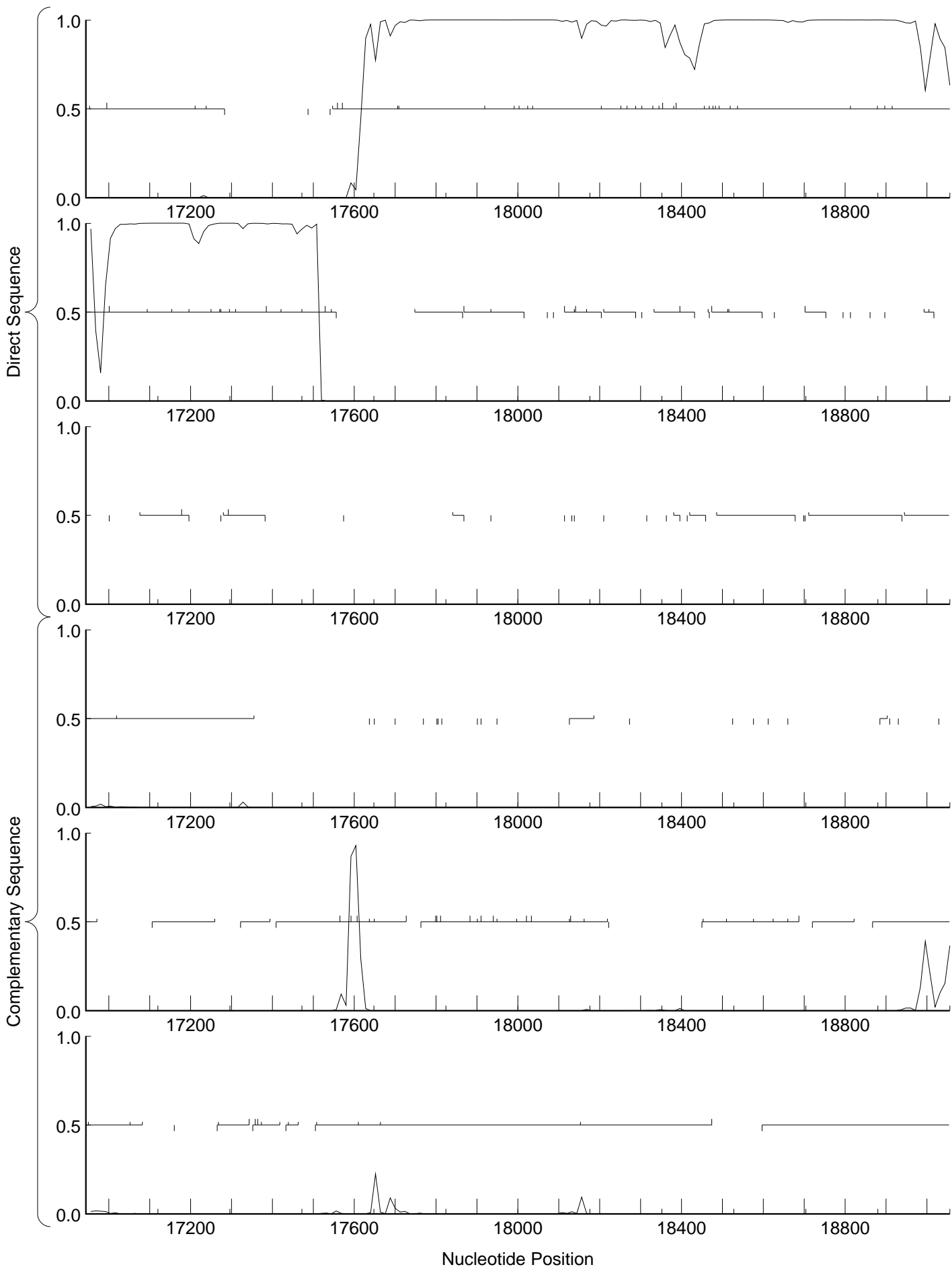




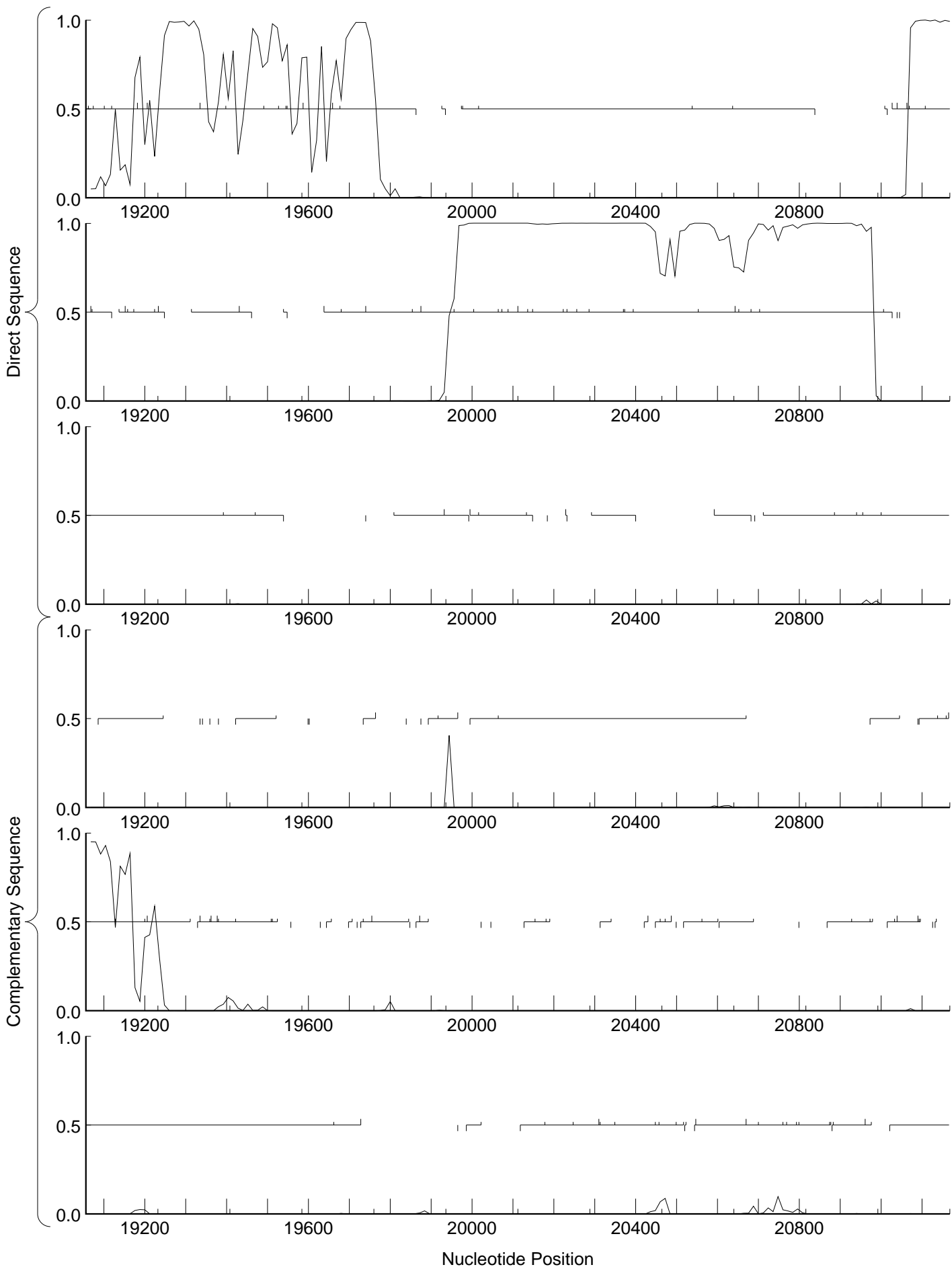


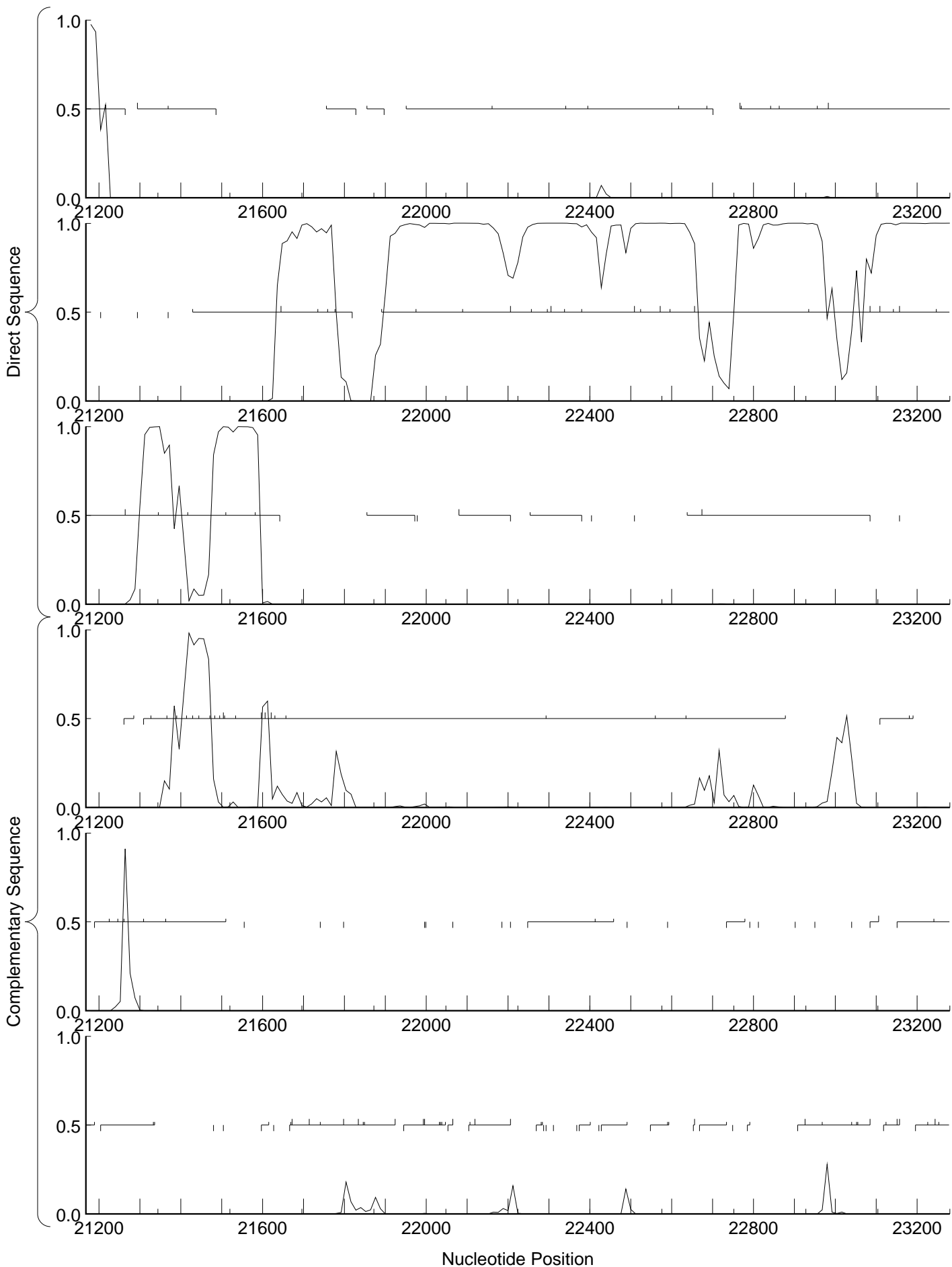


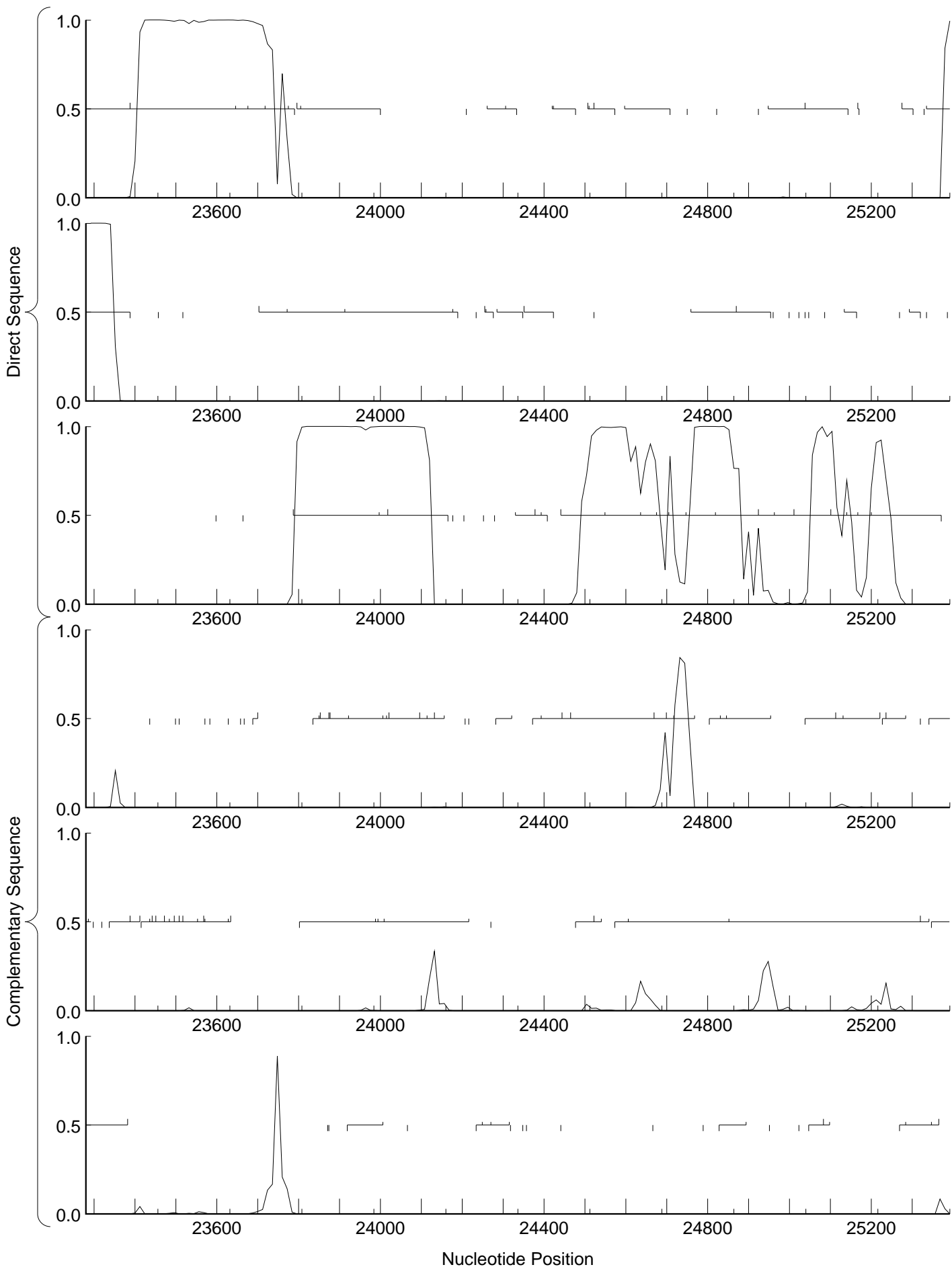


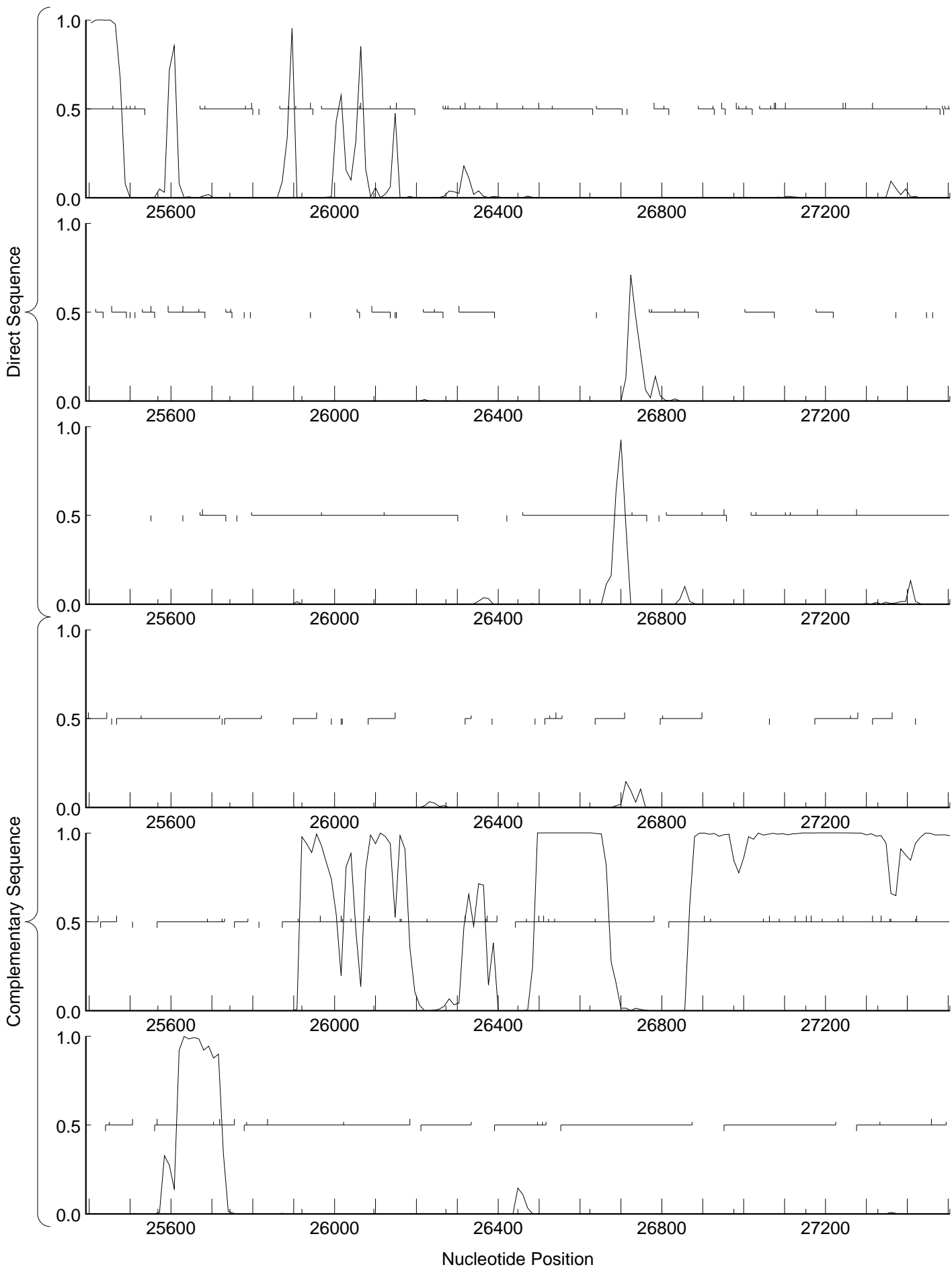


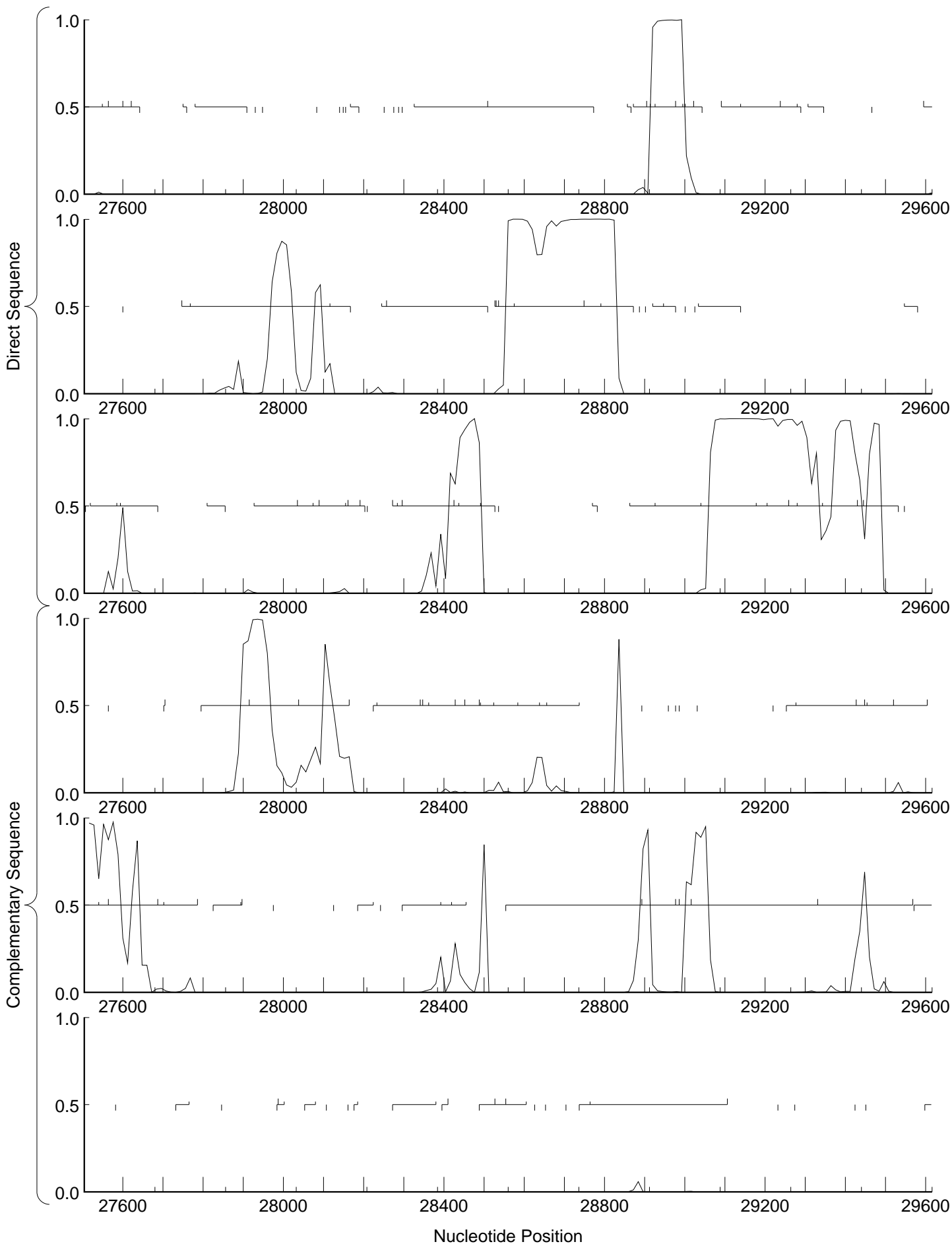
Complete sequence, 41278 bp including 13-base 3' overhang (CCCGCCGCCTTGG), Cluster N, Order 4, Window 96, Step 12, 11/21

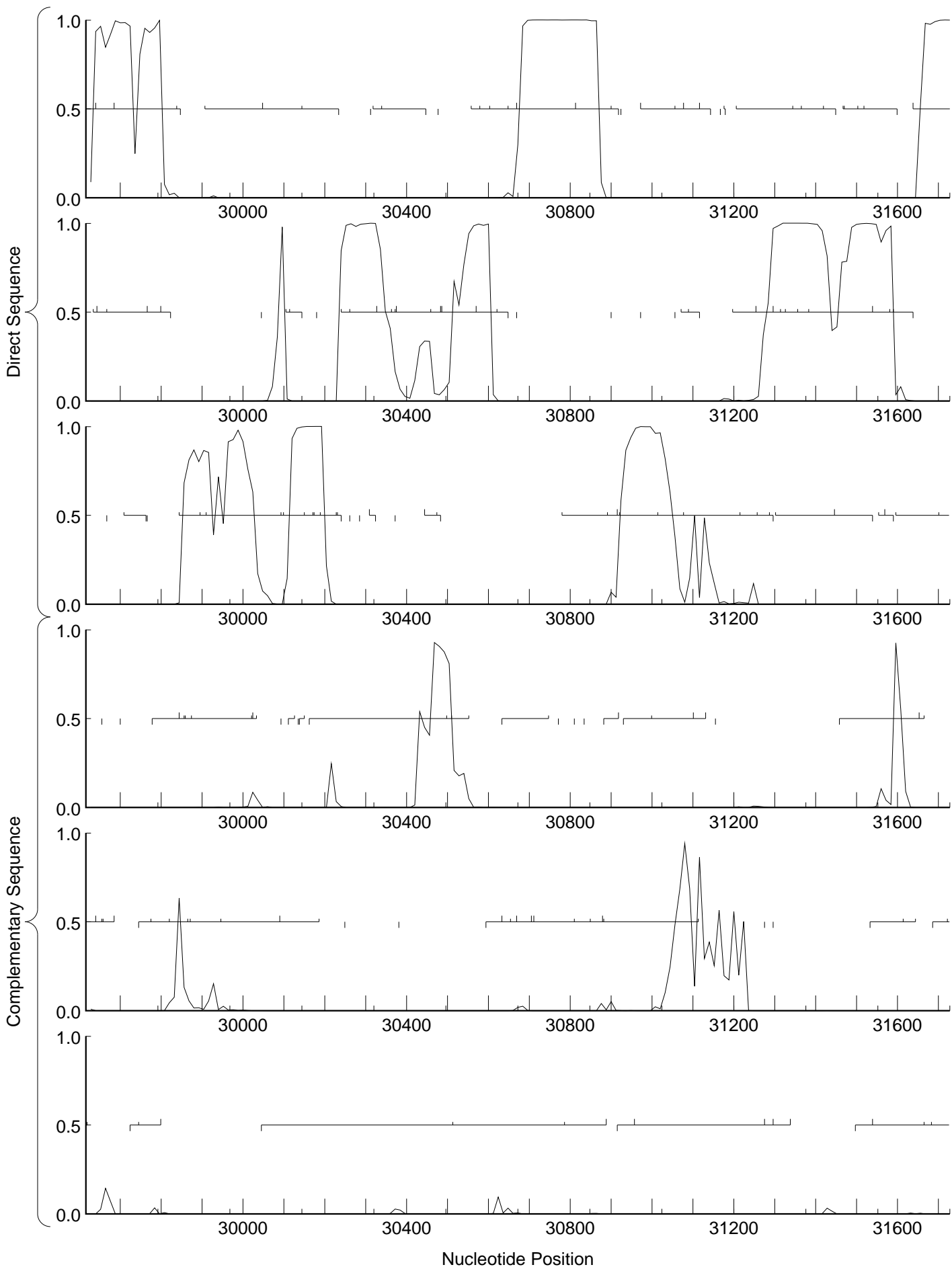














complete sequence, 41278 bp including 13-base 3' overhang (CCCGCCGCCTTGG), Cluster N, Order 4, Window 96, Step 12, 17/21

