

# GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

## PROGRAM INFORMATION

Sequence : Mycobacterium phage DrSeegs complete sequence, 76652 bp including 10-base 3' overhang (TCGA)  
Analysis Date : 2/3/23 at 5:40:05  
Pages : 38  
Sequence Length : 76652 bp  
GC Content : 58.93%

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,

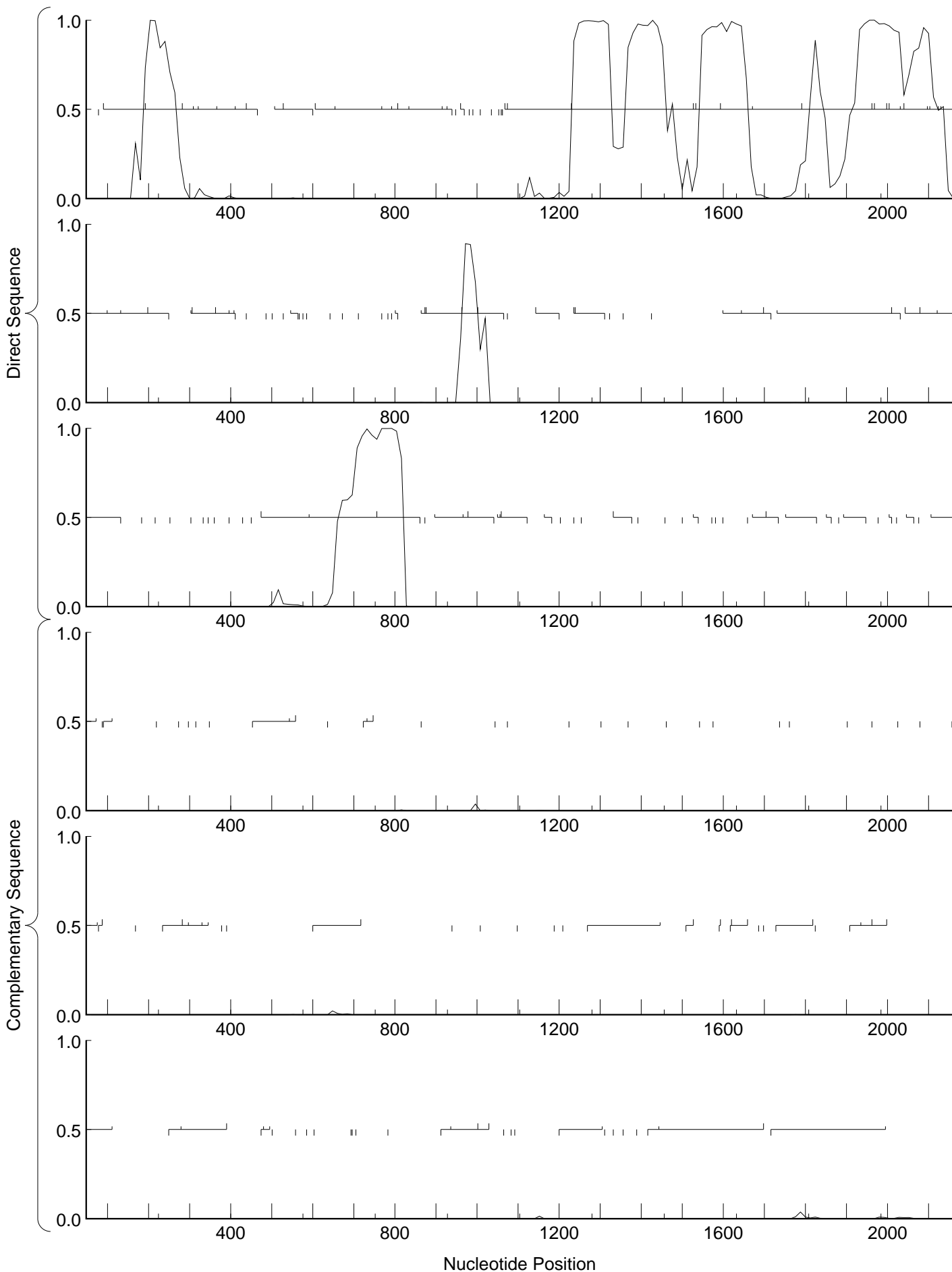
## MATRIX INFORMATION

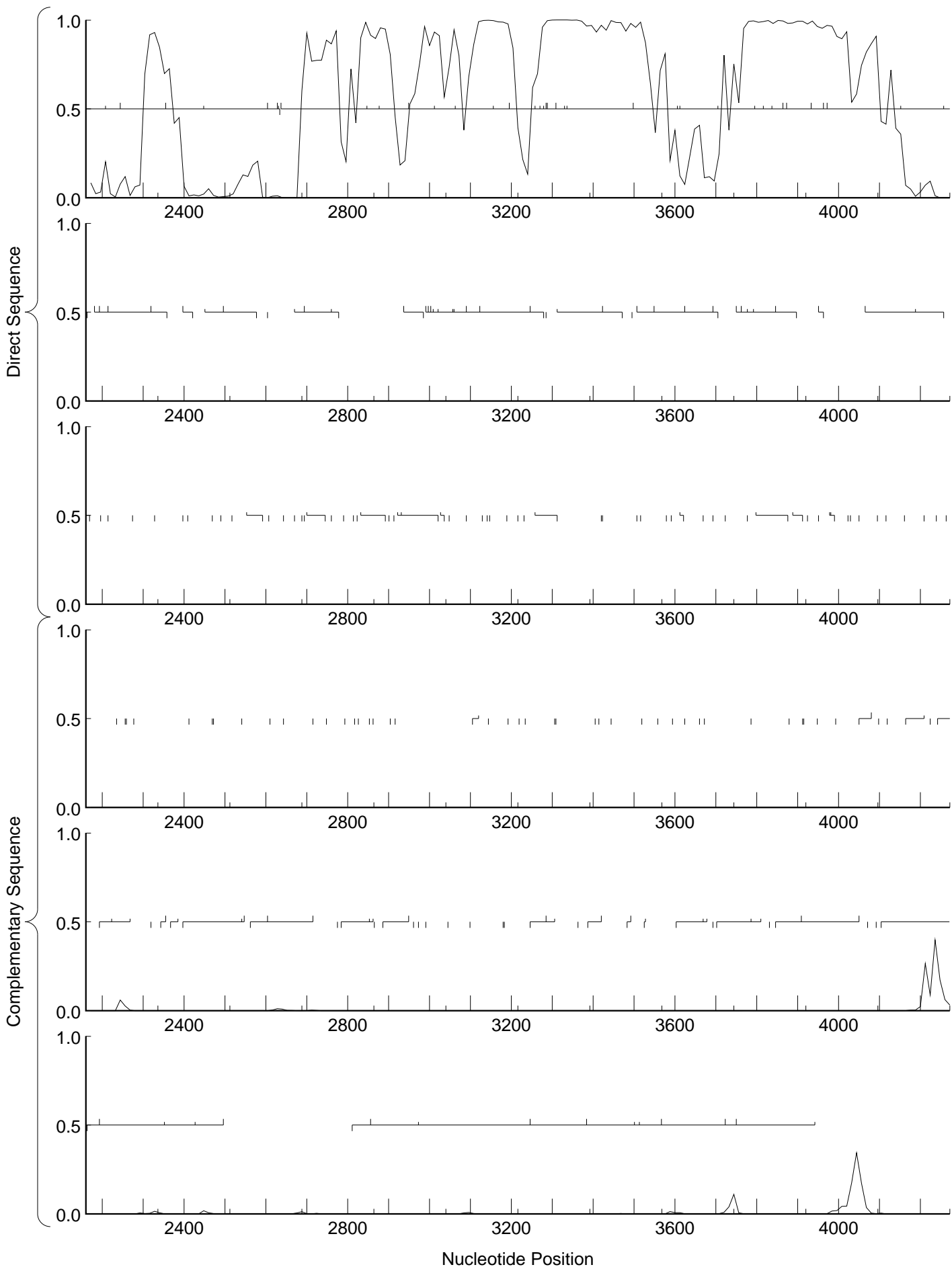
Matrix : Mycobacterium\_tuberculosis\_H37Rv  
Author : -  
Order : 4

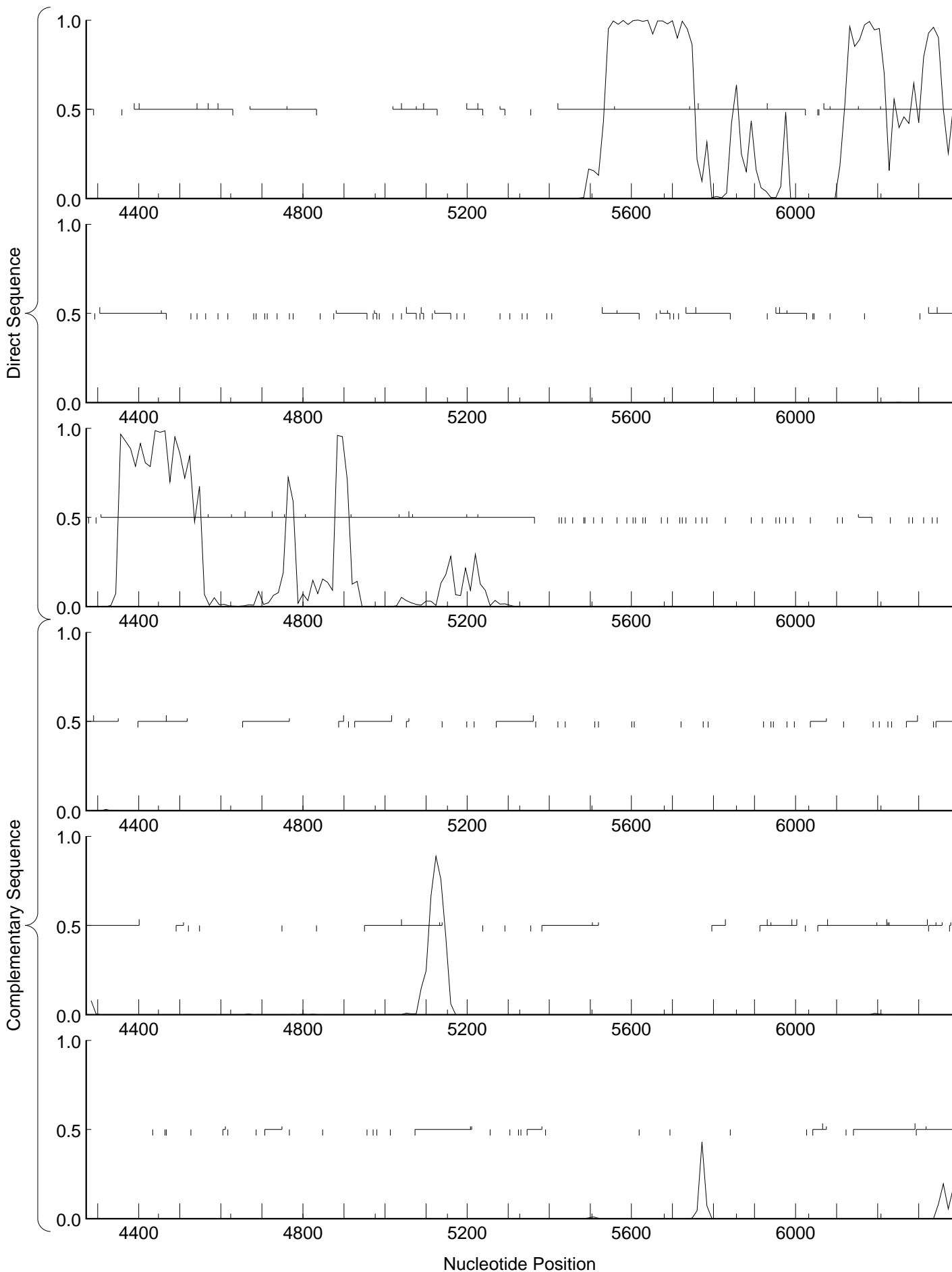
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

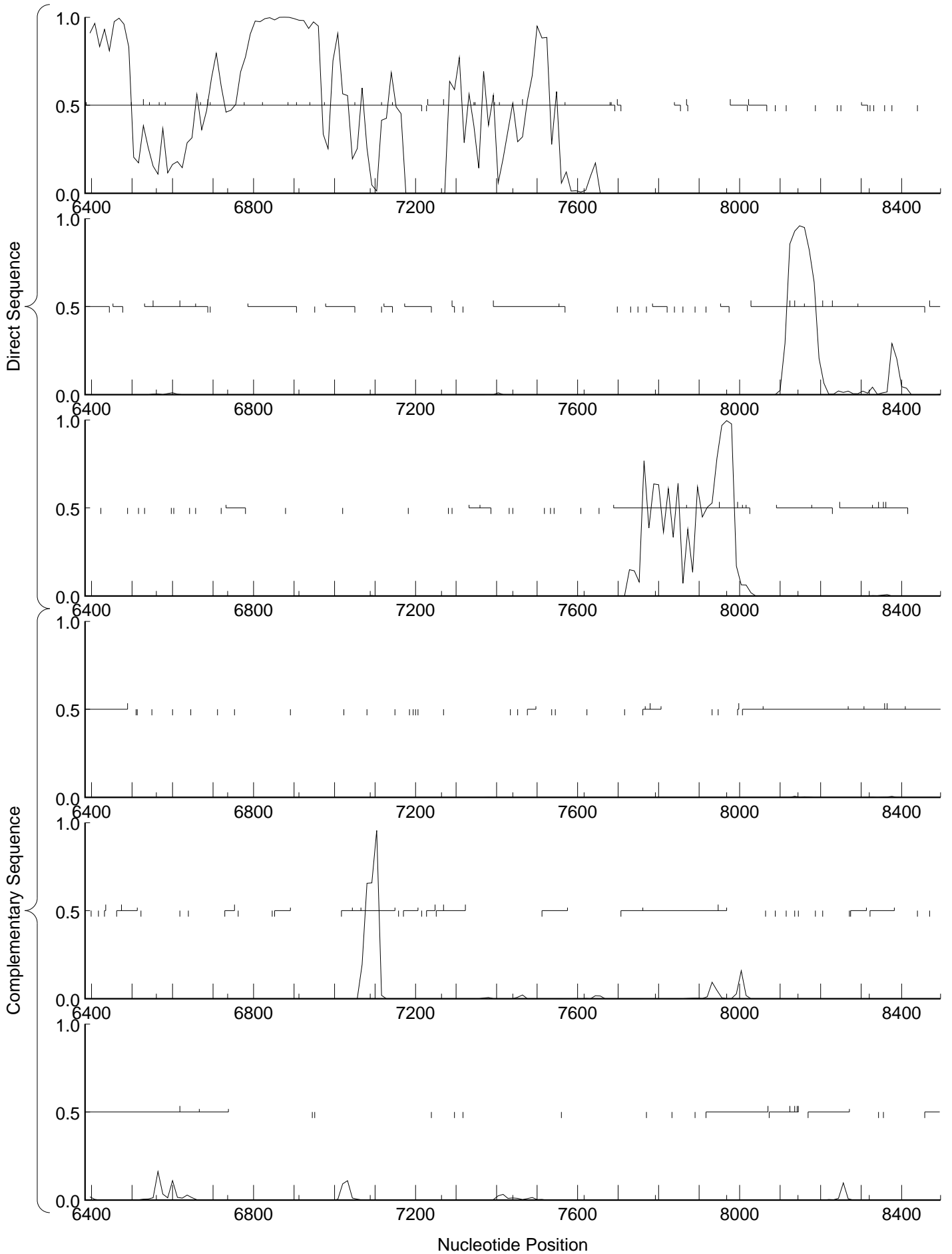
### Matrix notes & comments

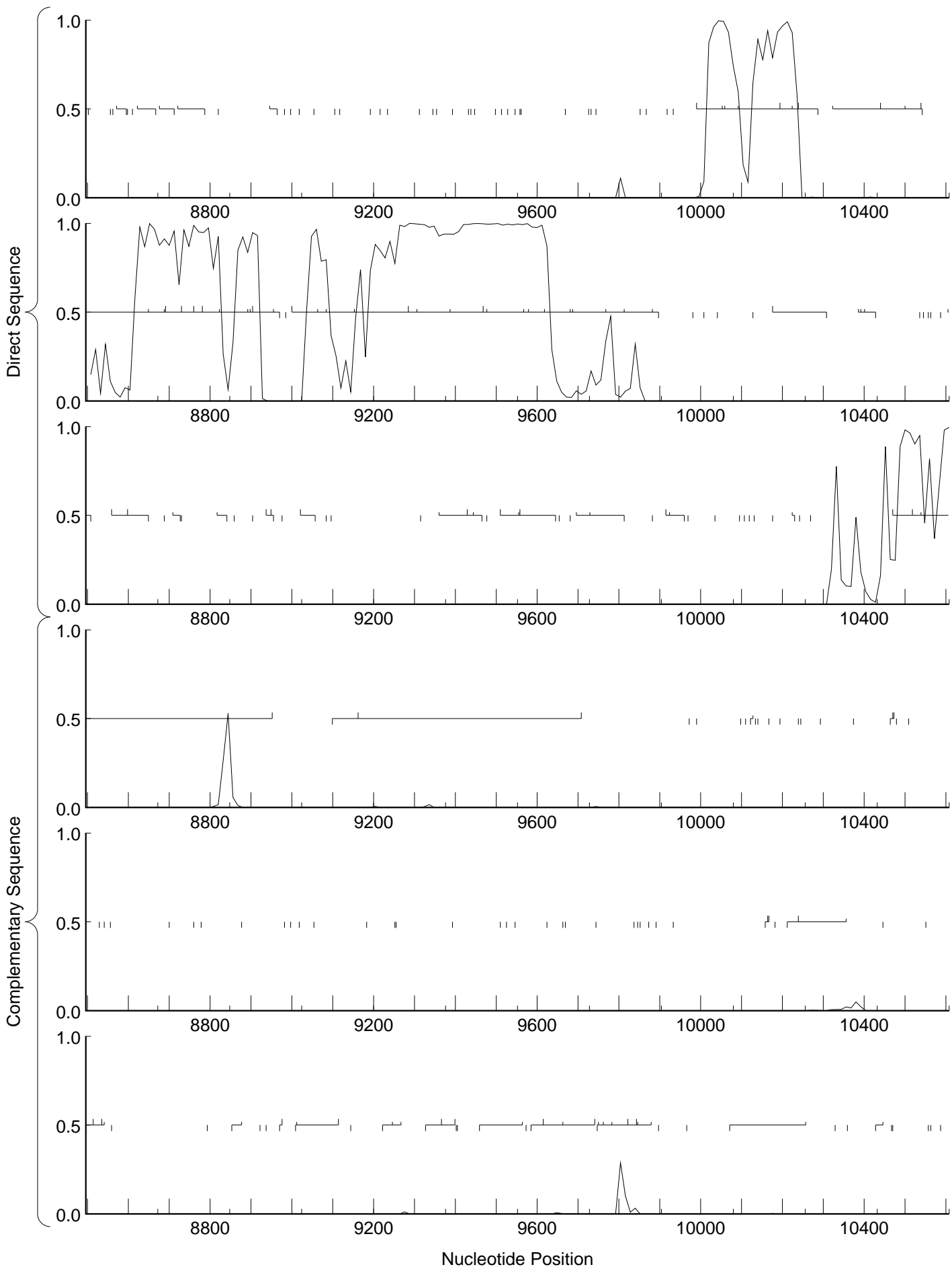
Training set derived by GeneMarkS, 4.27 September 2014  
Tue Sep 23 15:14:37 2014

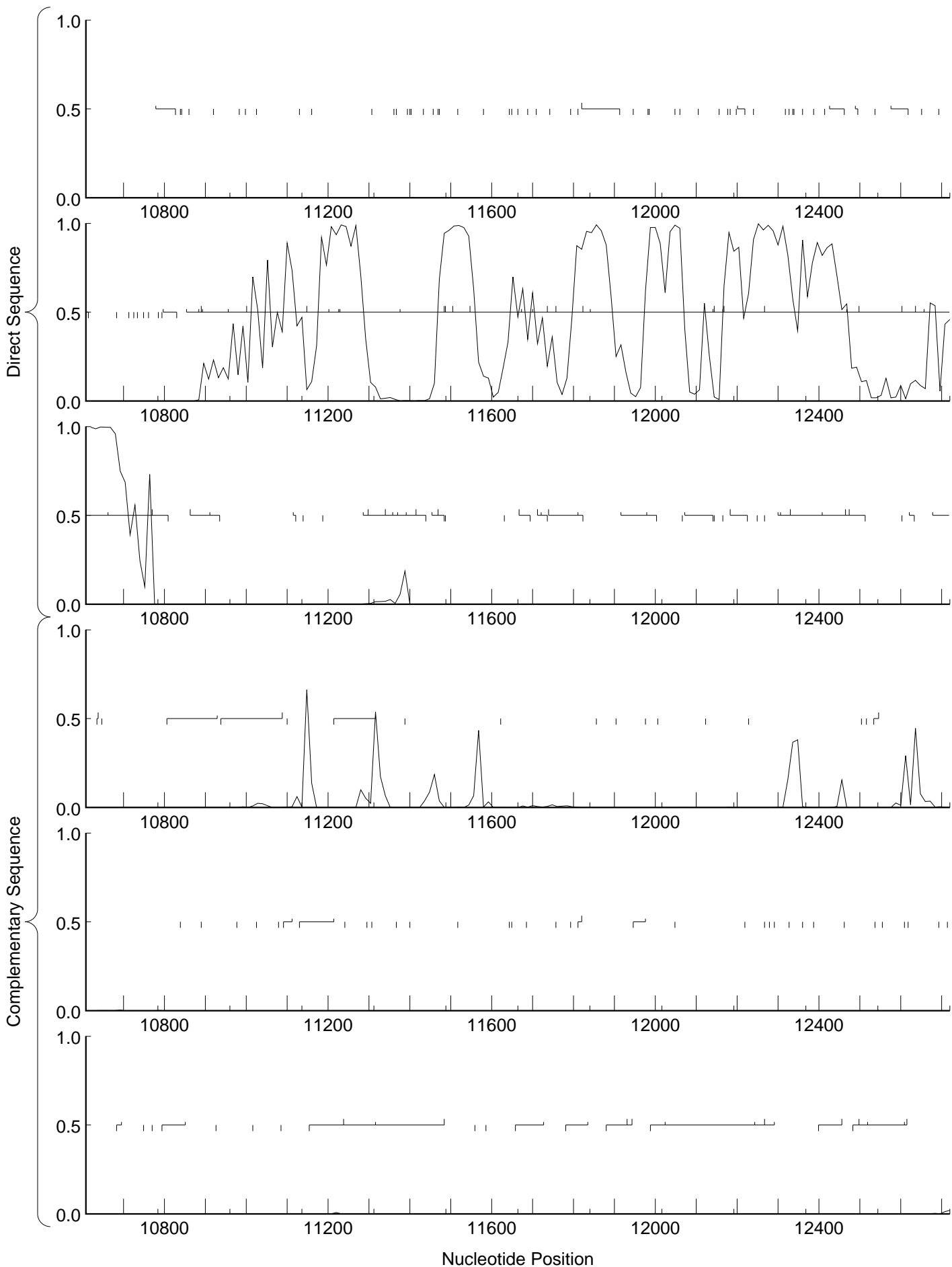


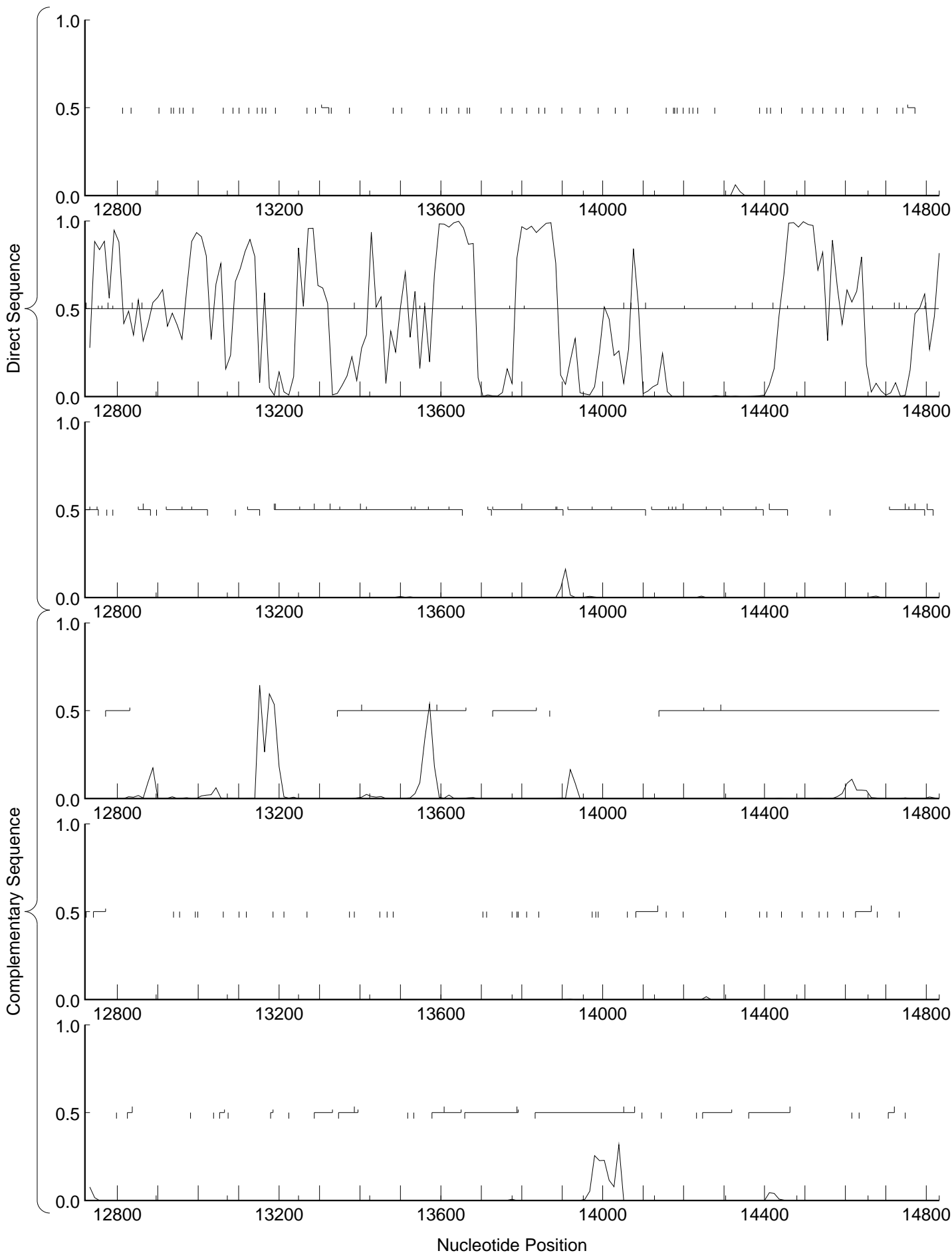




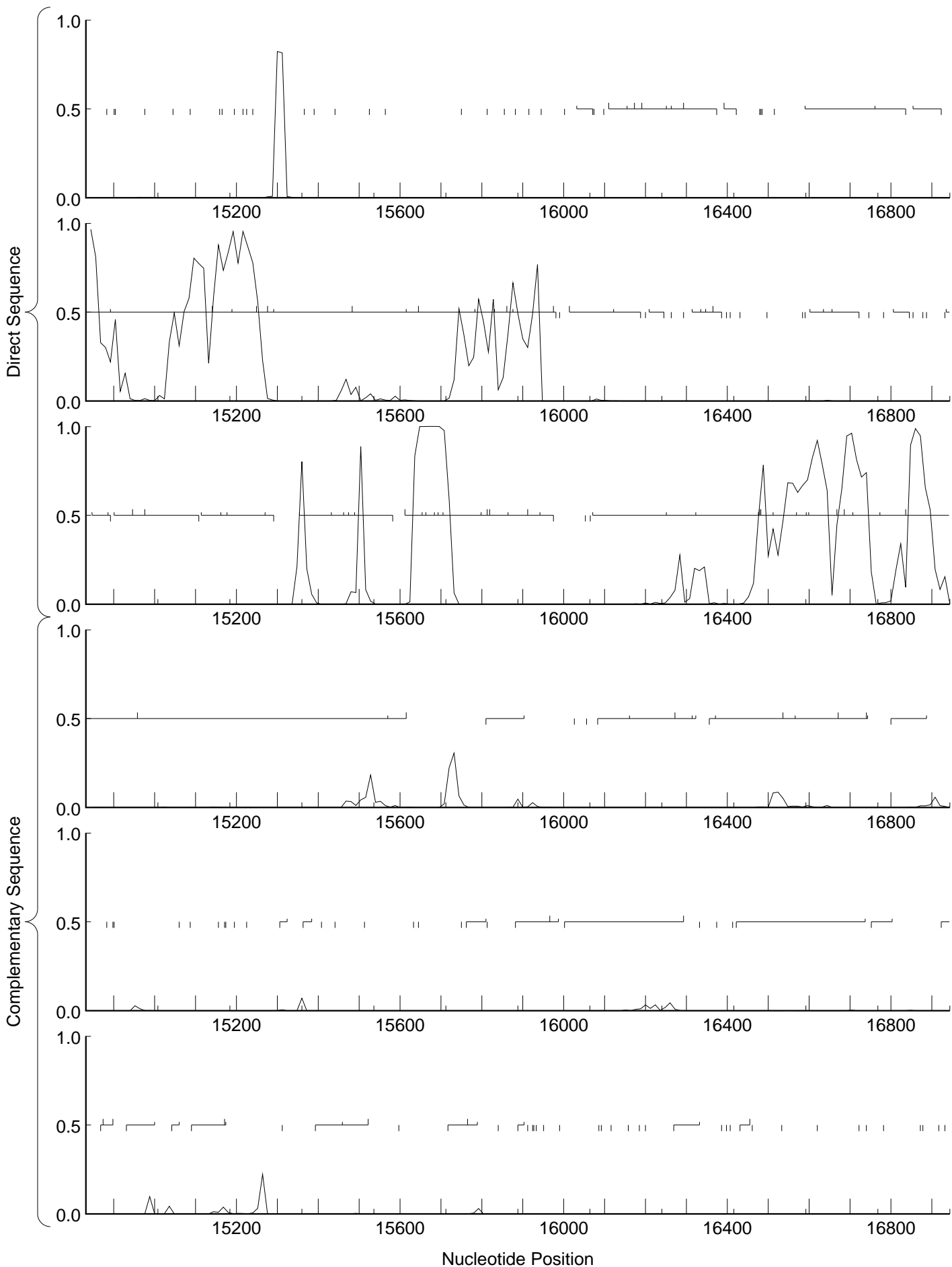


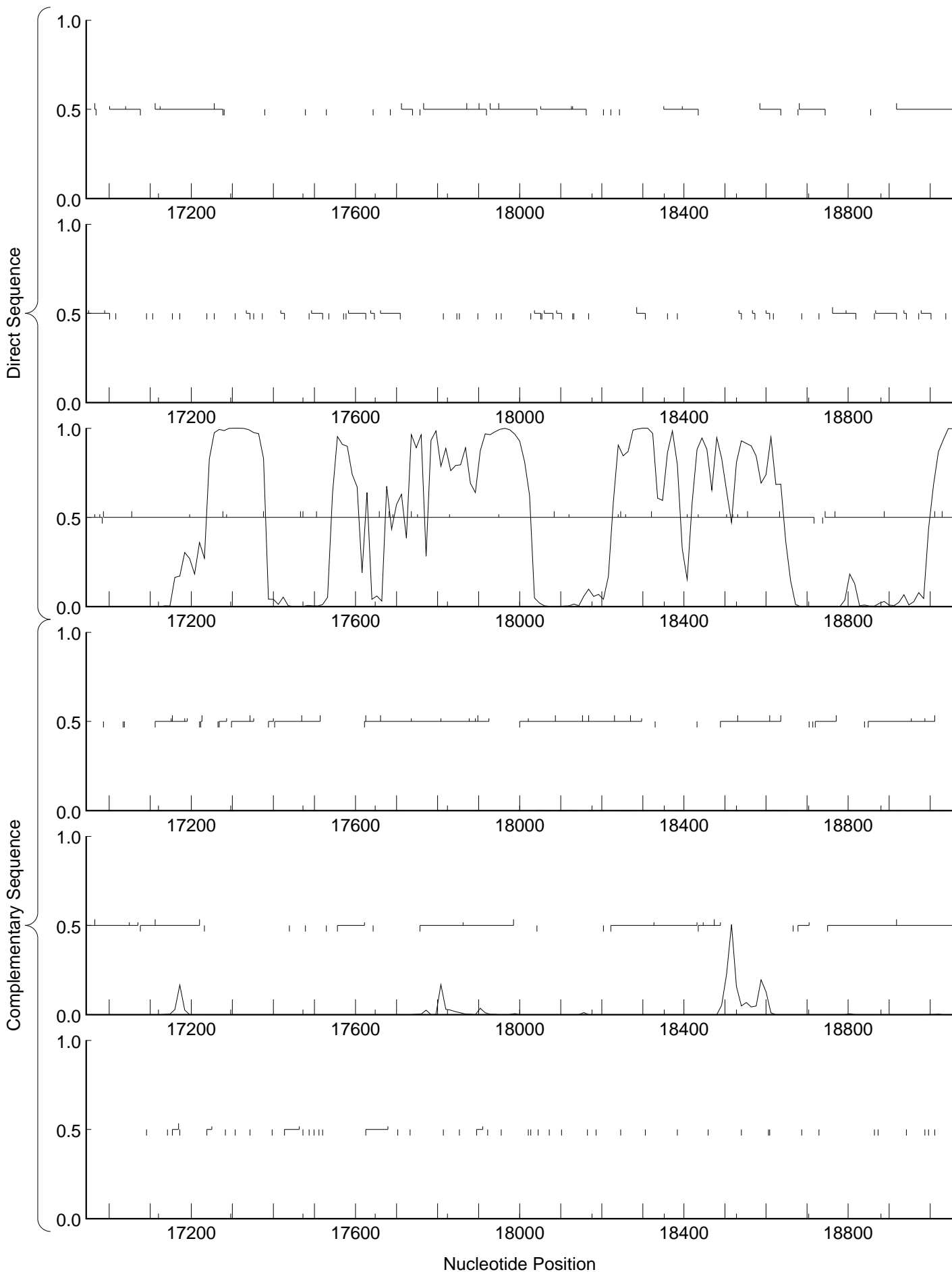












s complete sequence, 76652 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 11/38

