

# GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

## PROGRAM INFORMATION

Sequence : Mycobacterium phage Tiri complete sequence, 59449 bp including 11-base 3' overhang (CTCGTAG  
Analysis Date : 9/26/21 at 21:22:05  
Pages : 30  
Sequence Length : 59449 bp  
GC Content : 66.64%

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,

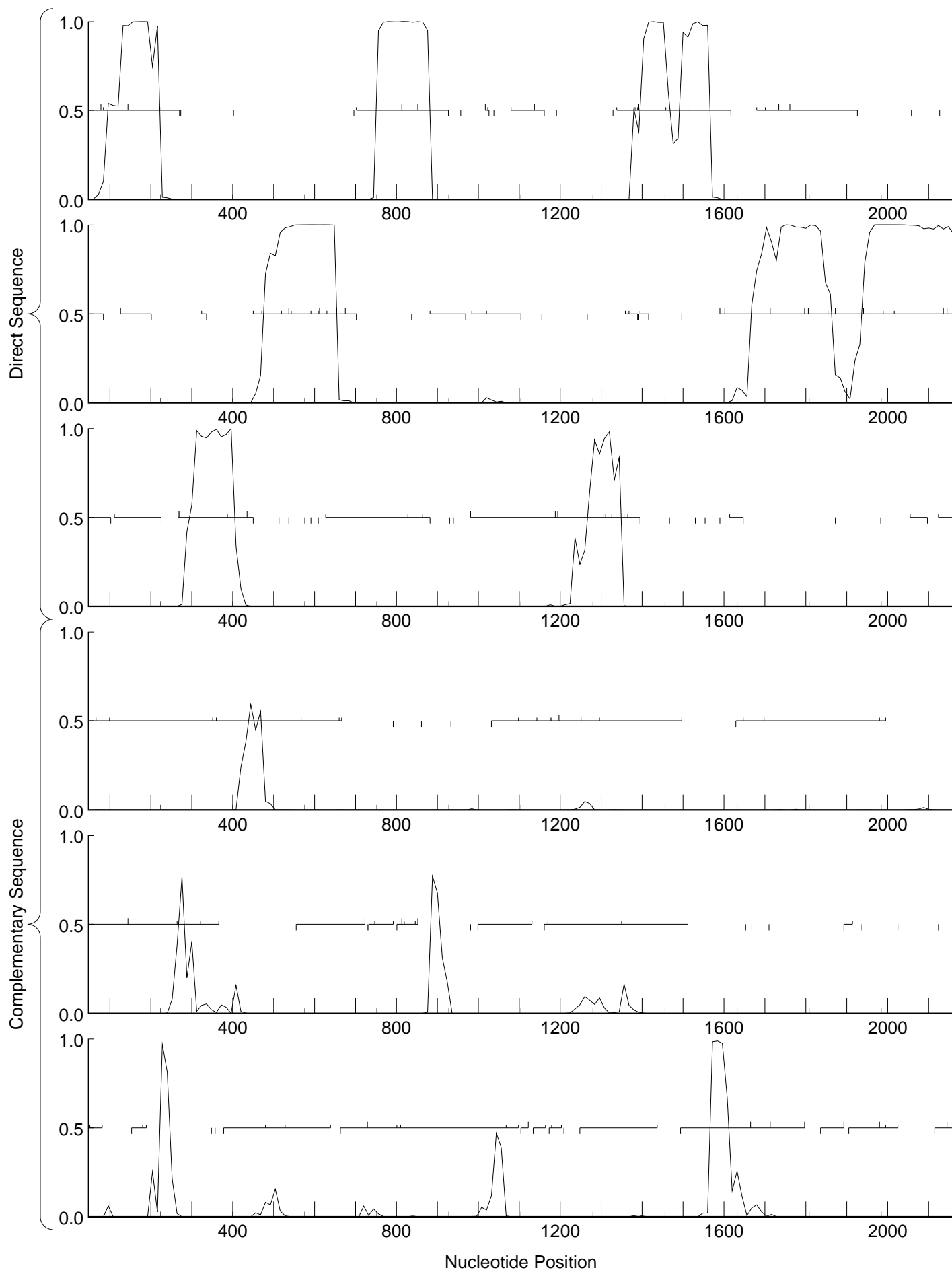
## Matrix notes & comments

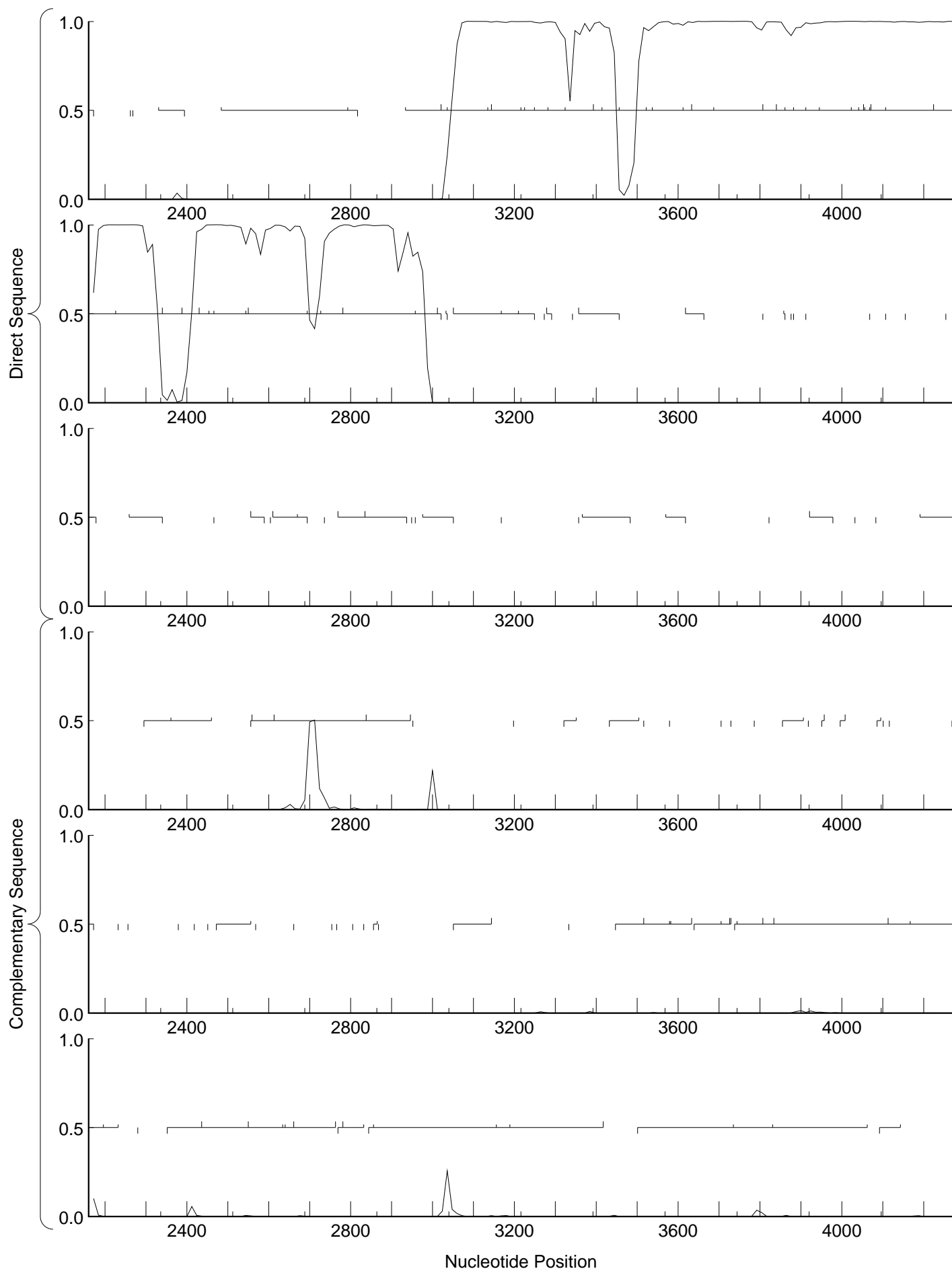
Training set derived by GeneMarkS, 4.27 September 2014  
Tue Sep 23 15:14:37 2014

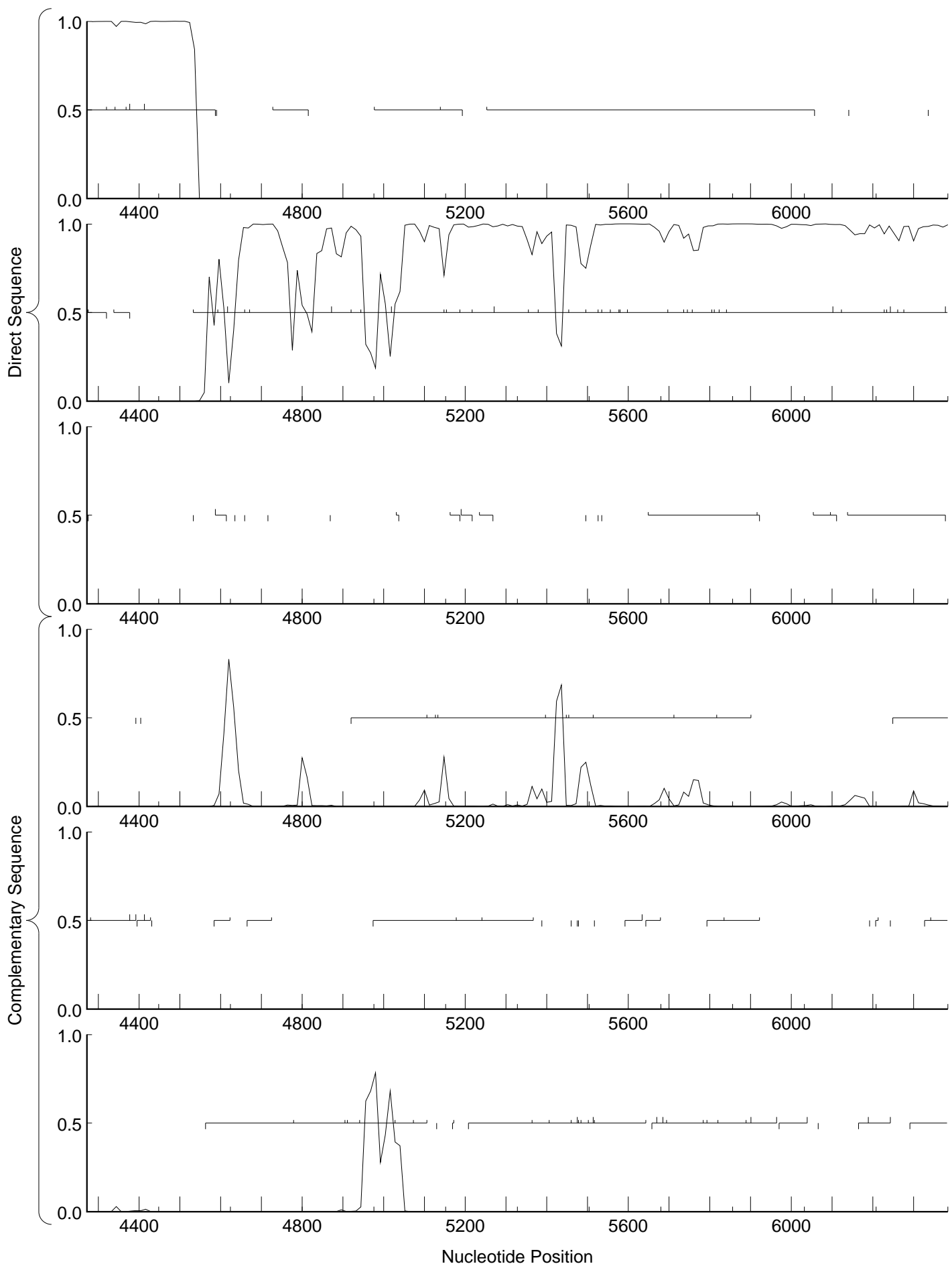
## MATRIX INFORMATION

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

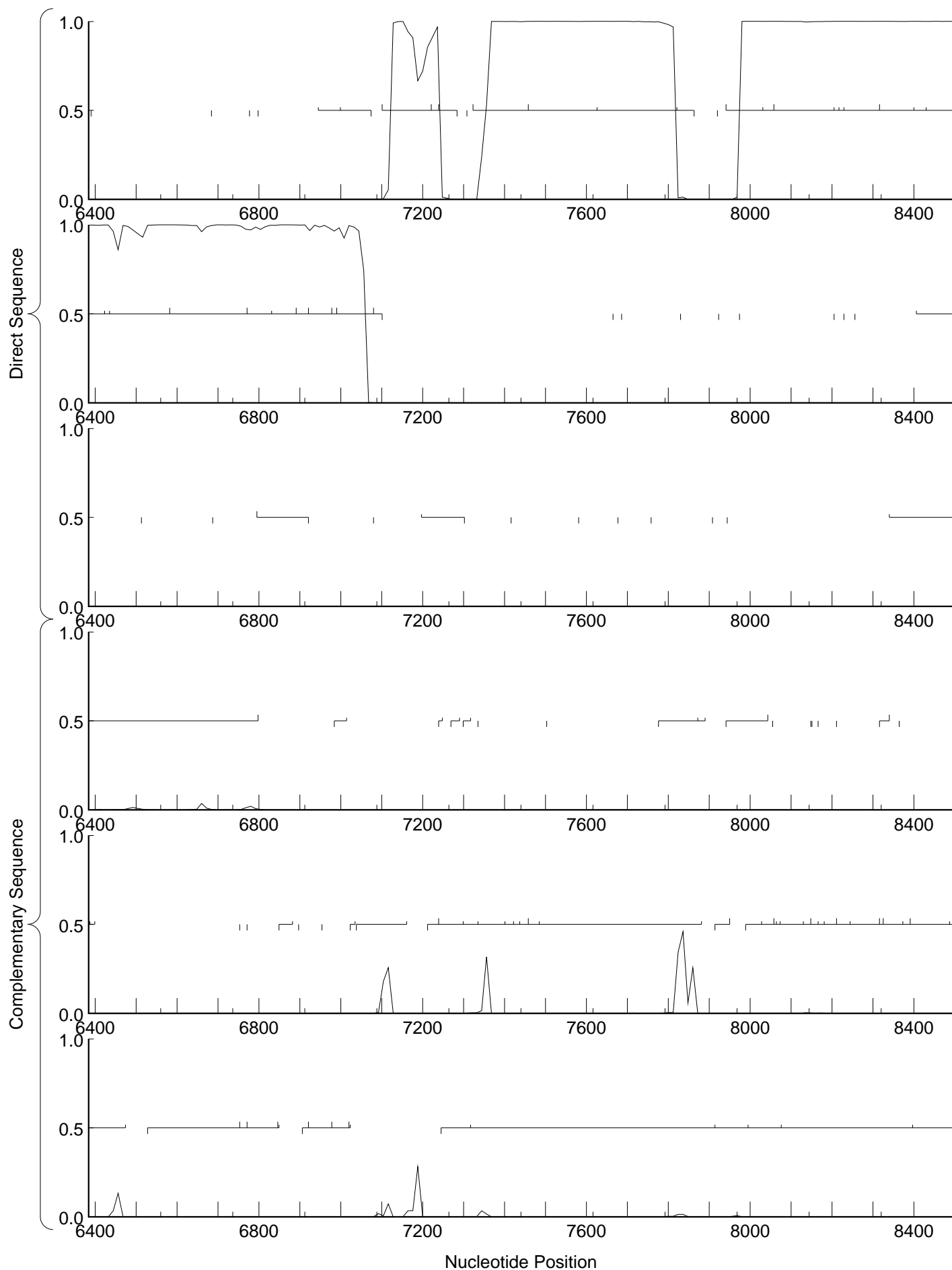
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

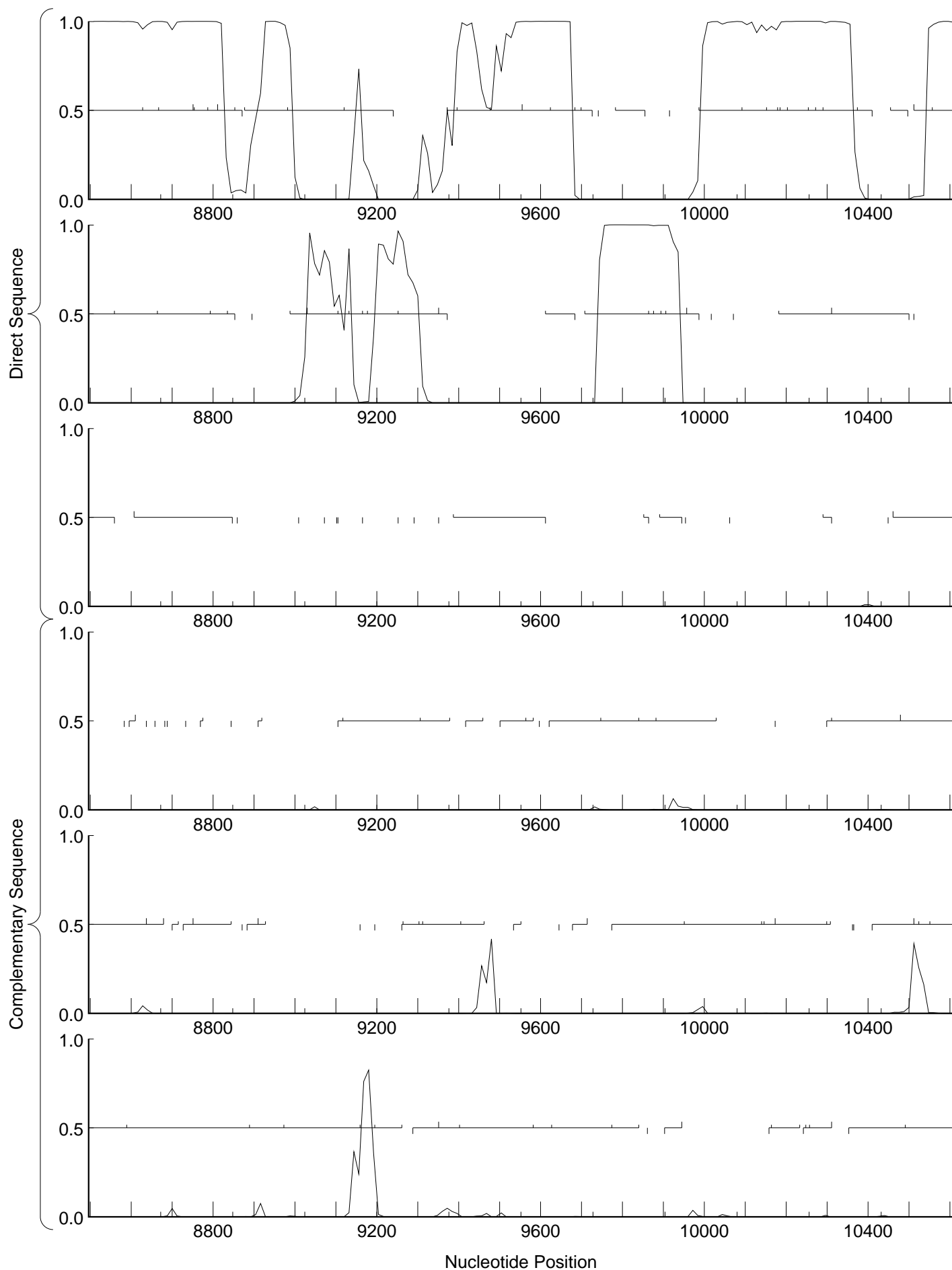




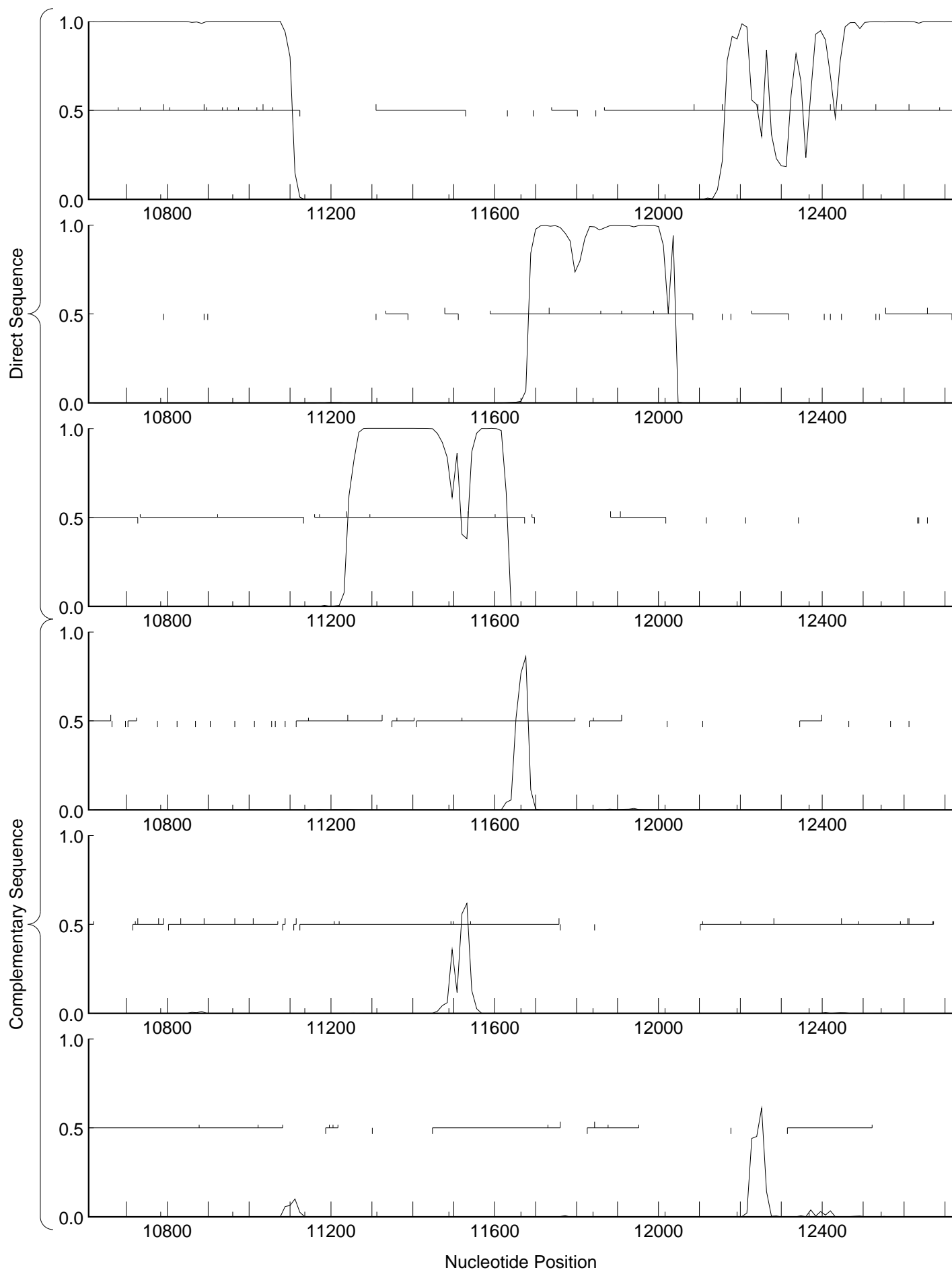


complete sequence, 59449 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 5/30





complete sequence, 59449 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 7/30



complete sequence, 59449 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 8/30

