

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Arthrobacter phage Kepler complete sequence, 38449 bp including 12-base 3' overhang (GAGTTG)  
Analysis Date : 2/23/18 at 17:11:45  
Pages : 20  
Sequence Length : 38449 bp  
GC Content : 66.74%

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,

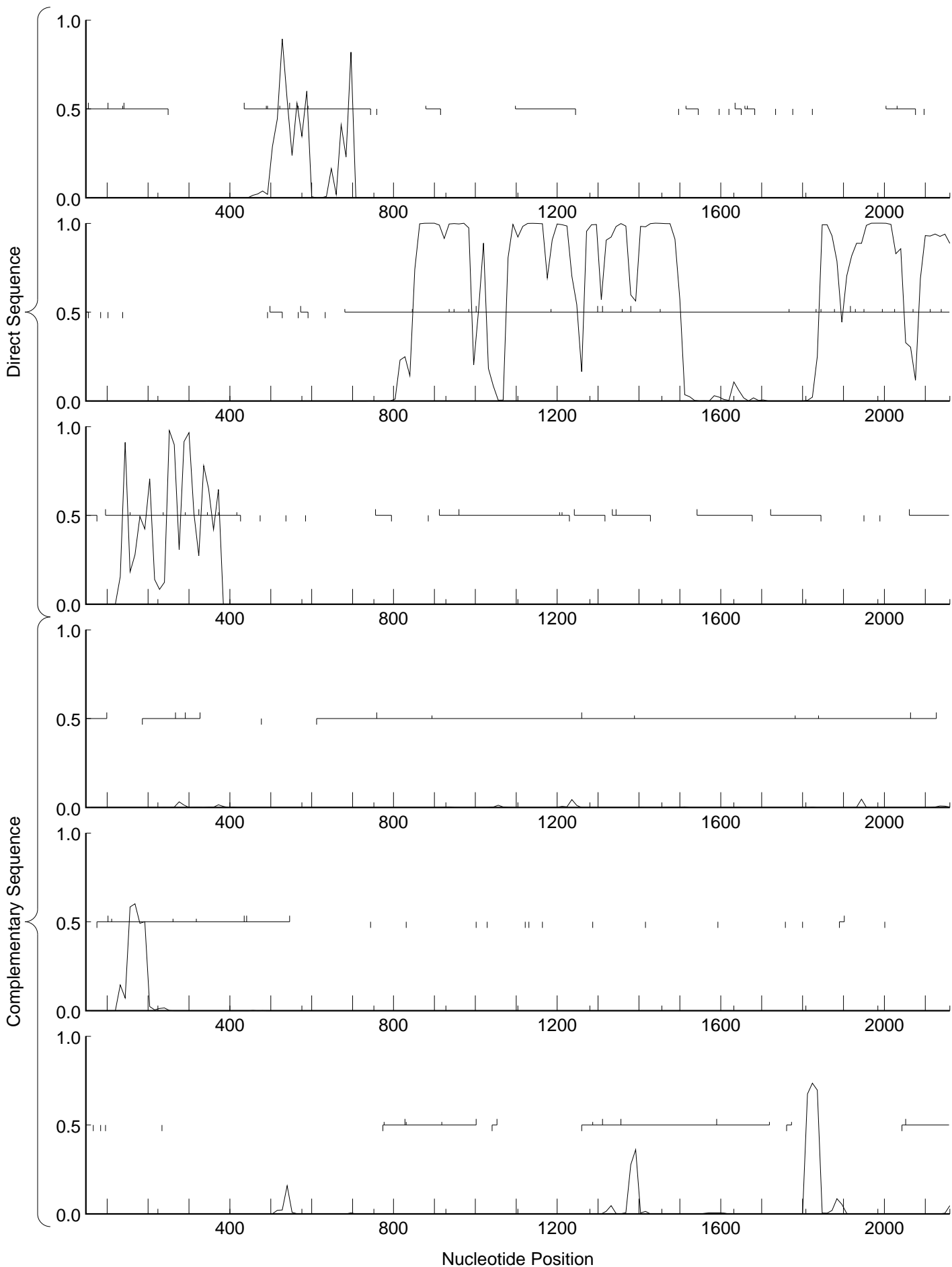
## Matrix notes & comments

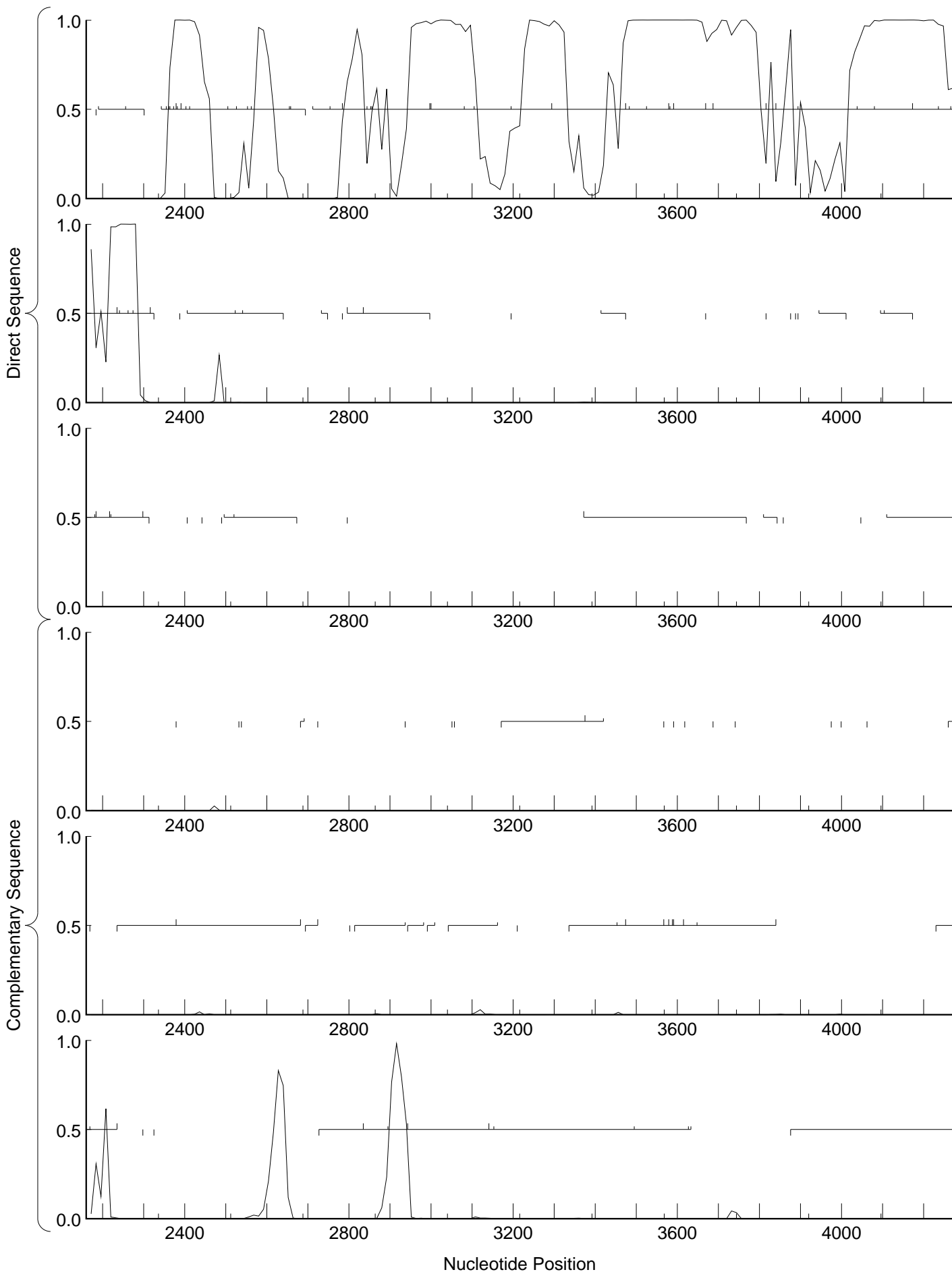
Training set derived by GeneMarkS, 4.27 September 2014  
Tue Sep 23 15:23:03 2014

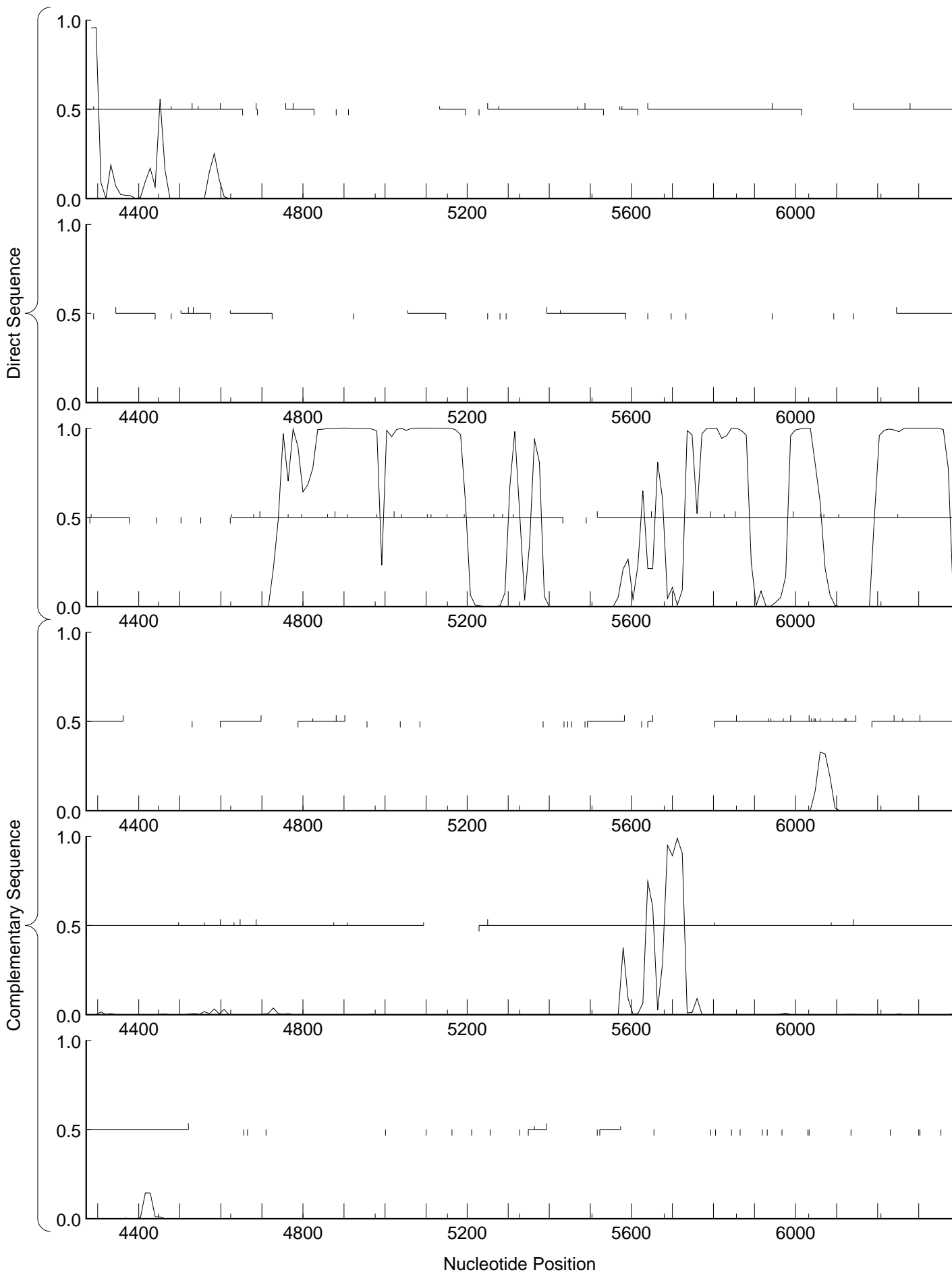
## MATRIX INFORMATION

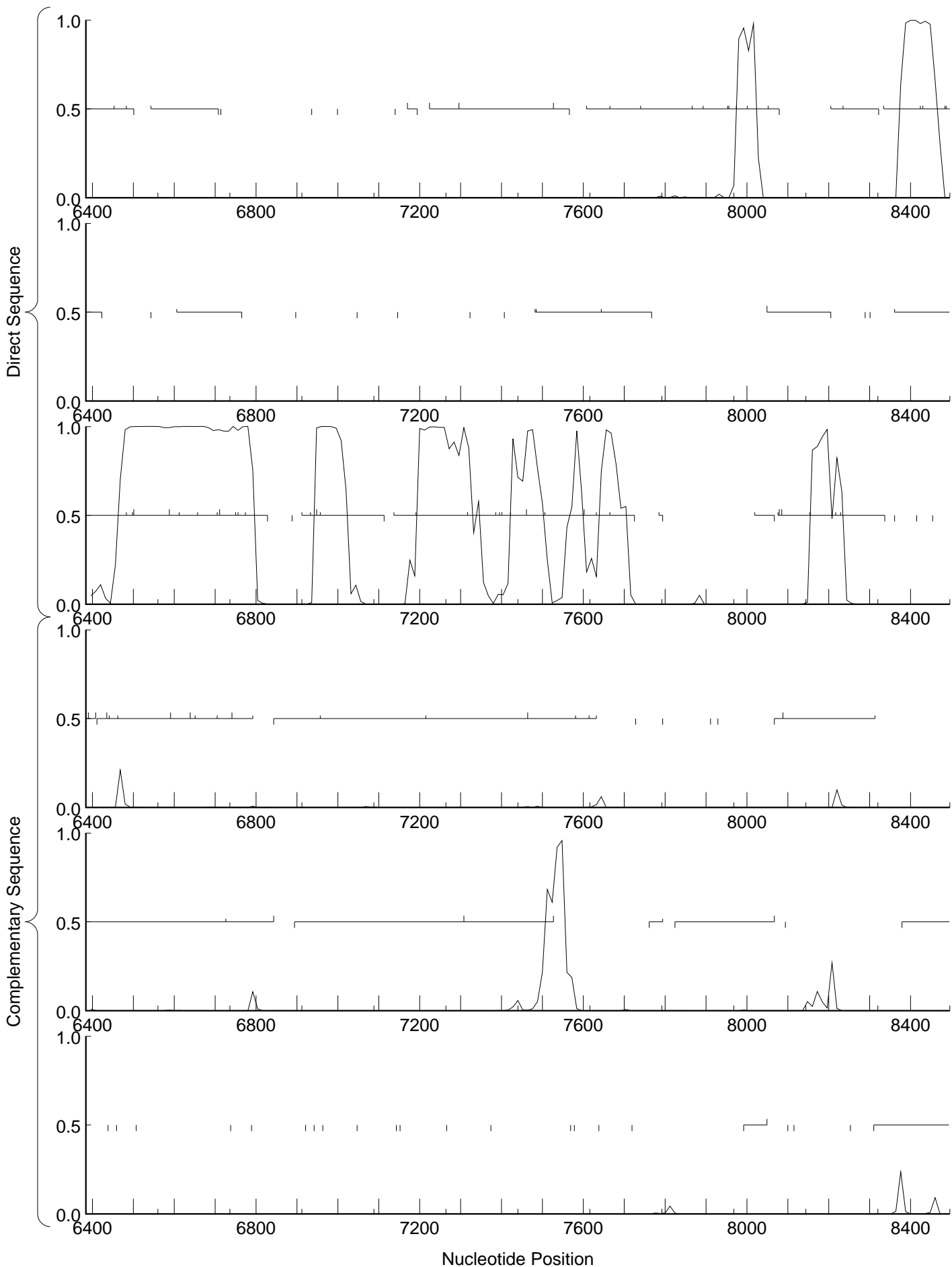
Matrix : Mycobacterium\_smegmatis\_MC2\_155  
Author : -  
Order : 4

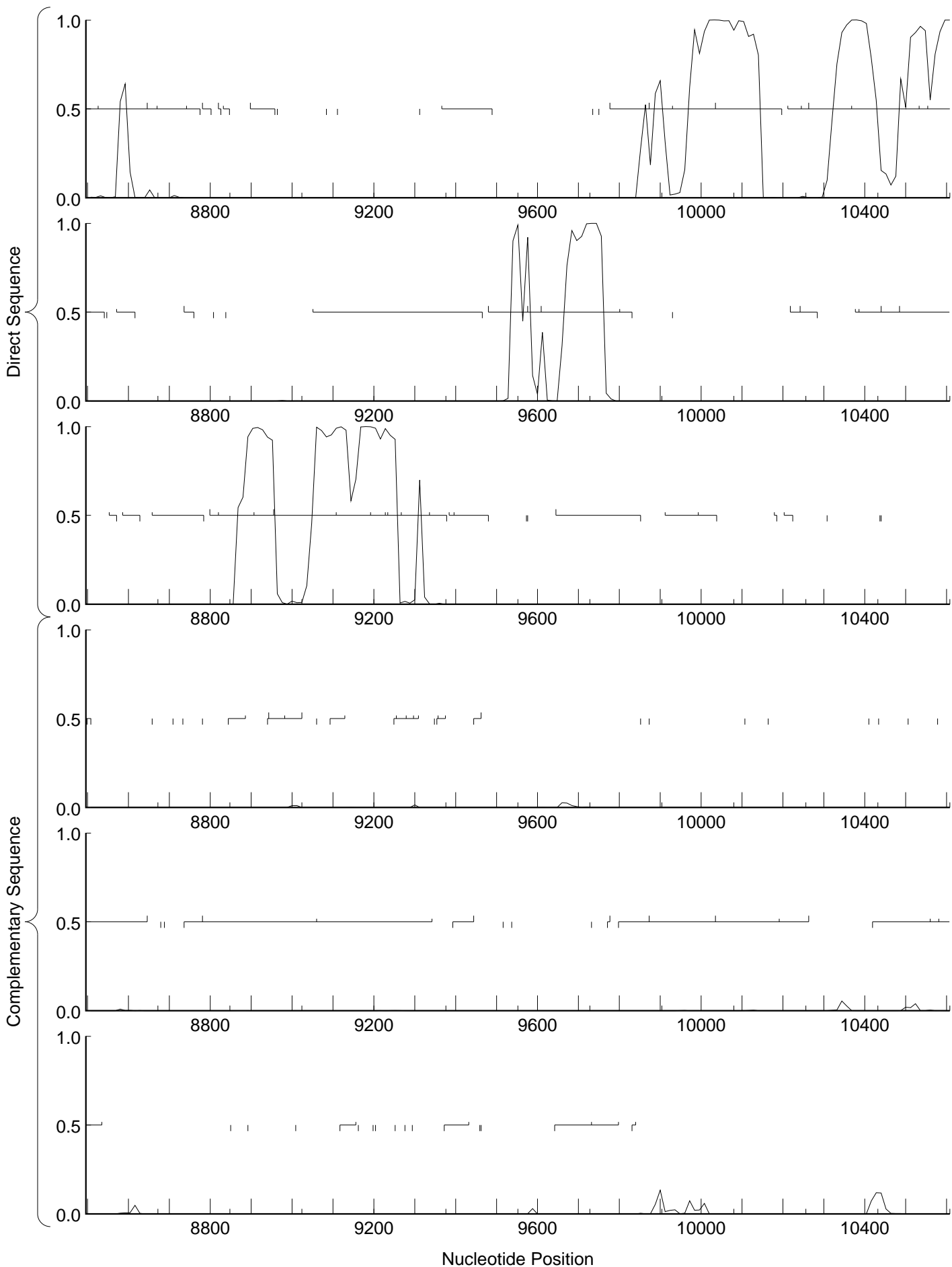
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

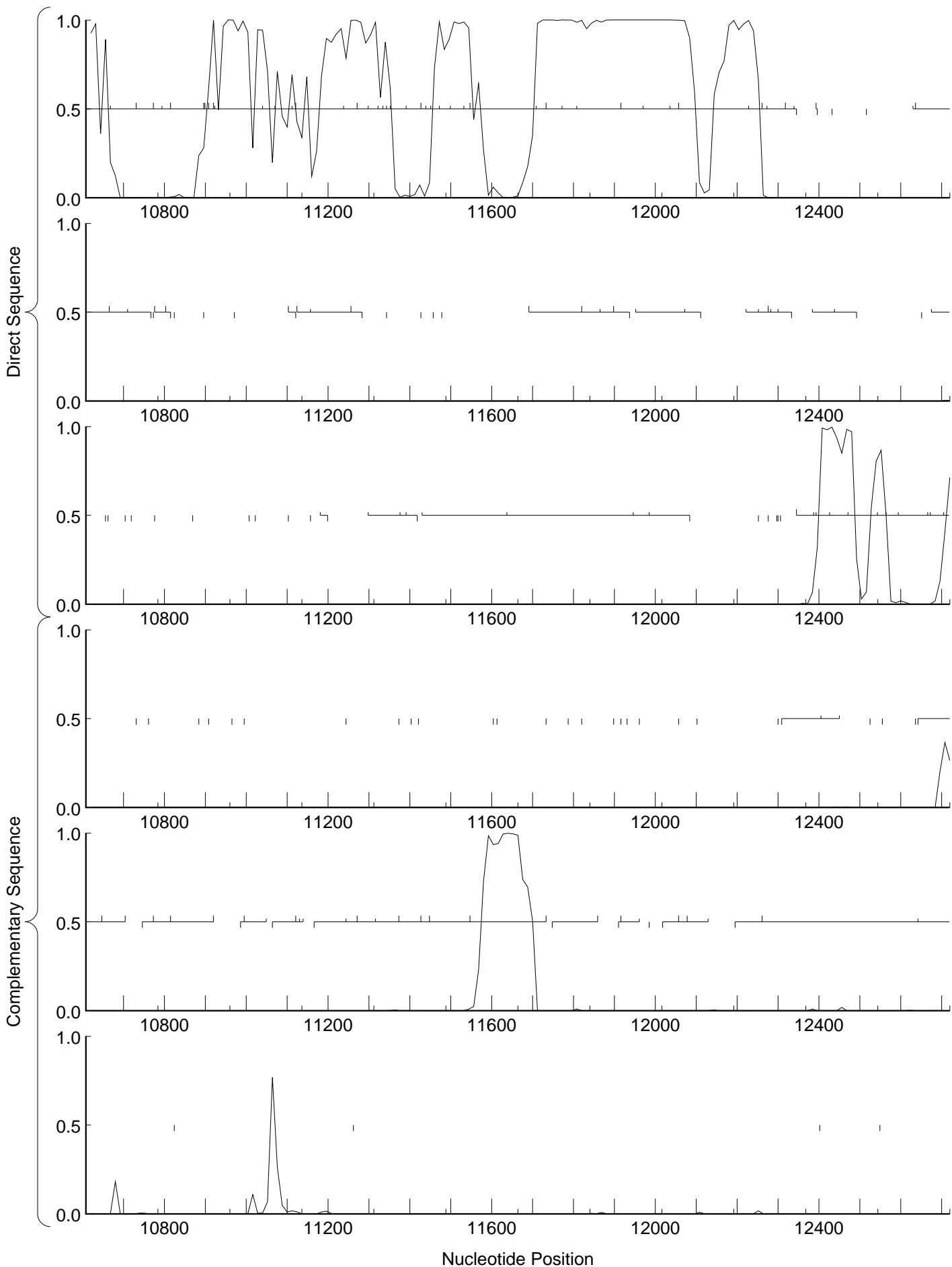


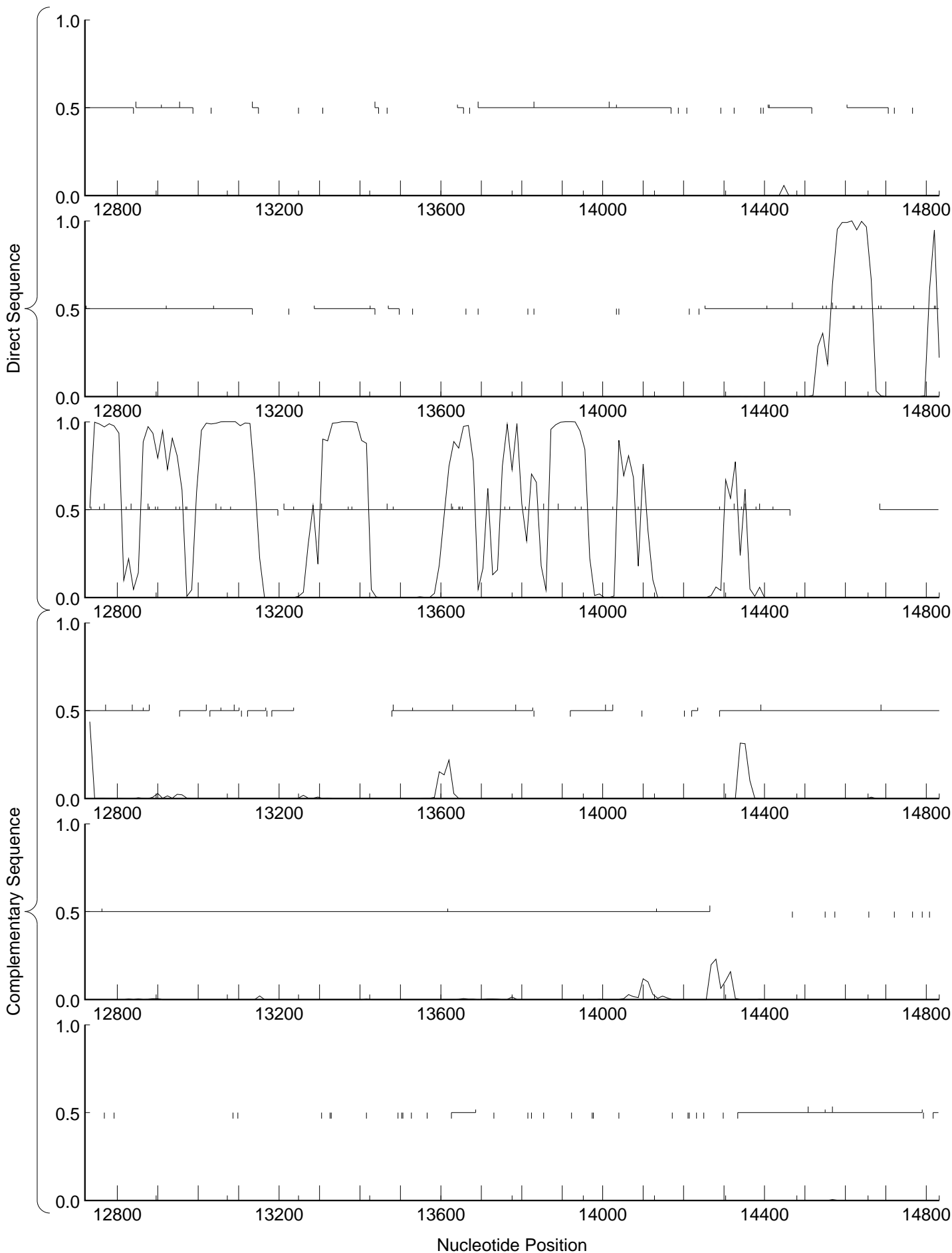




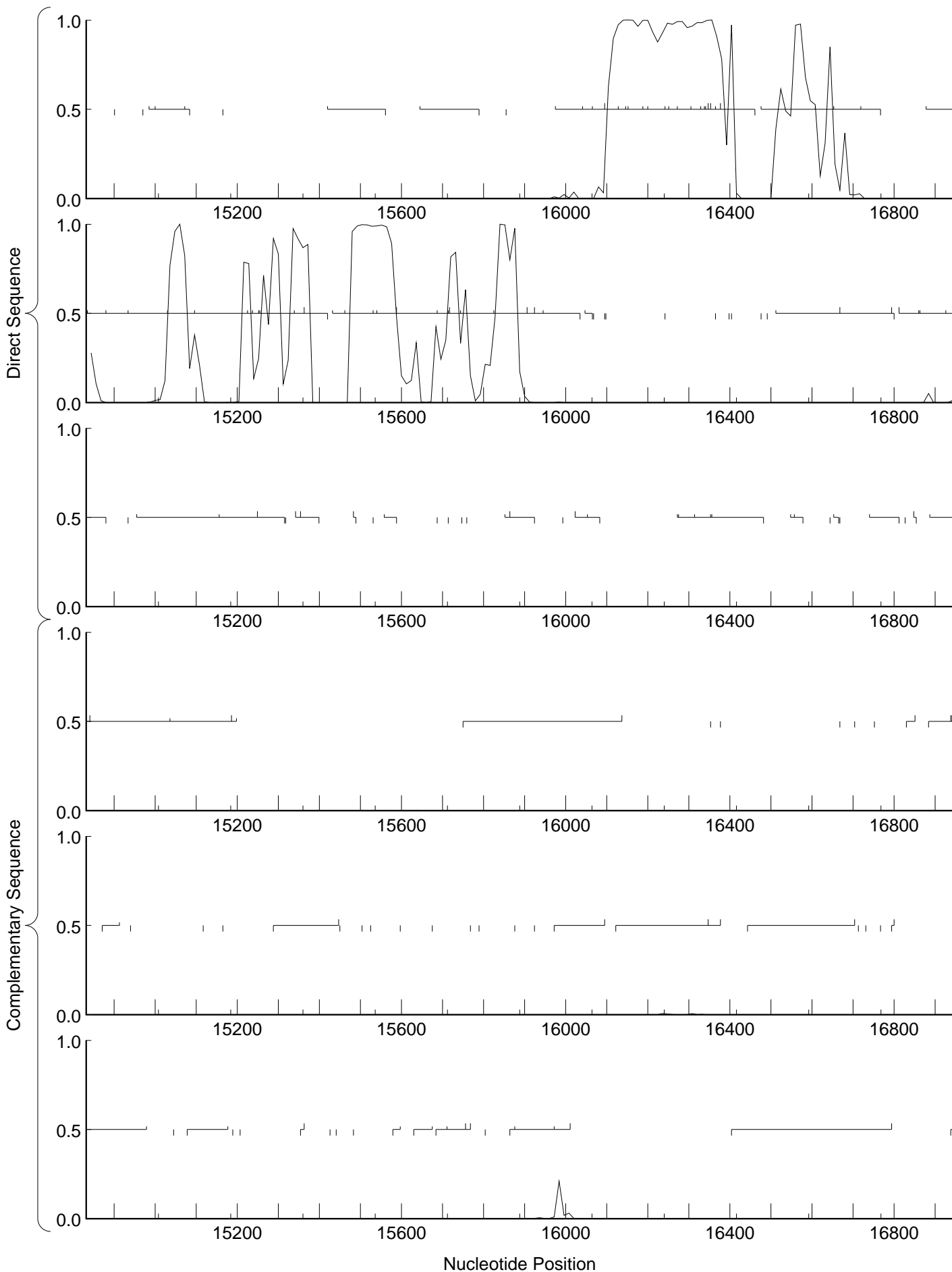


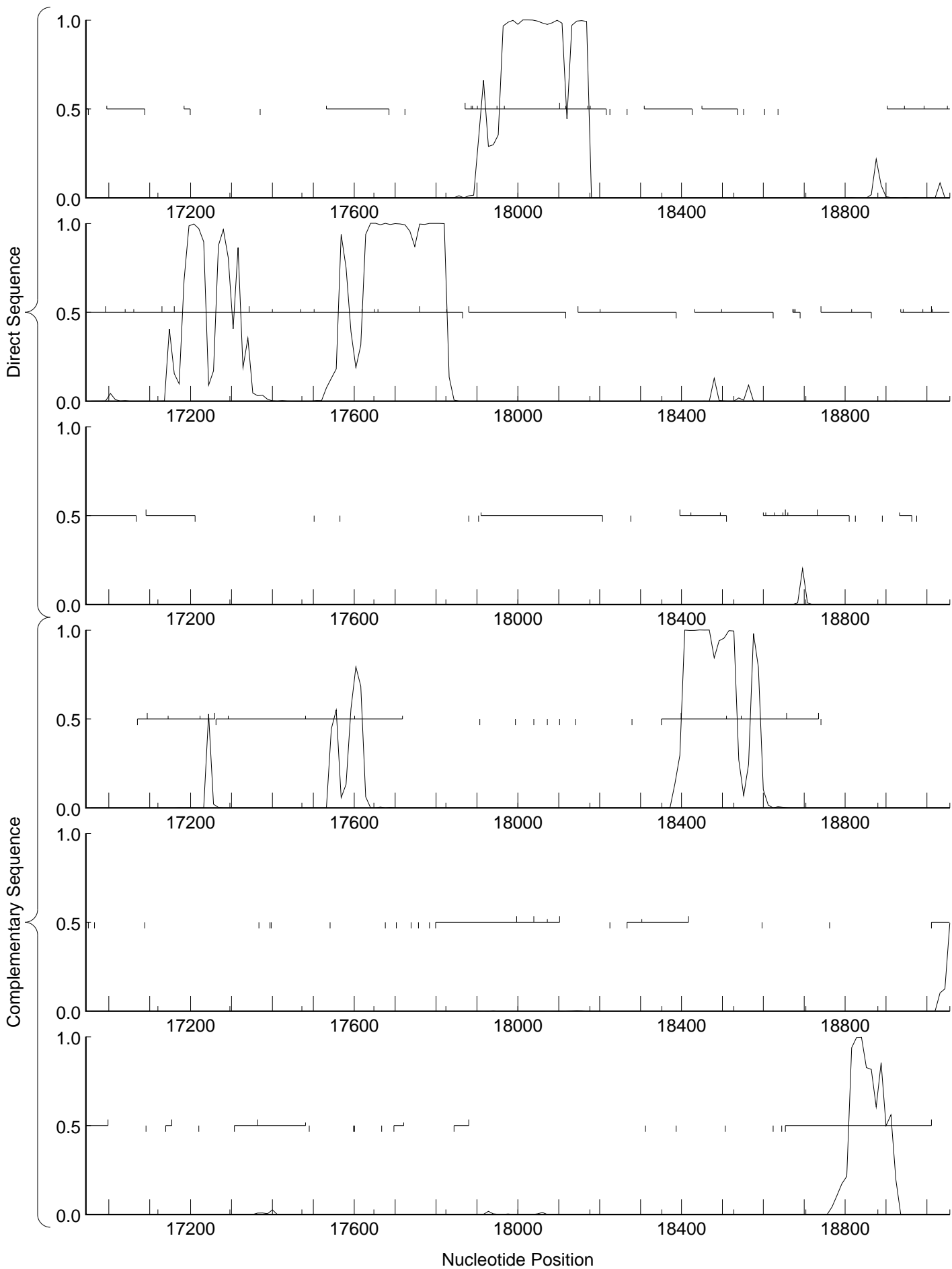


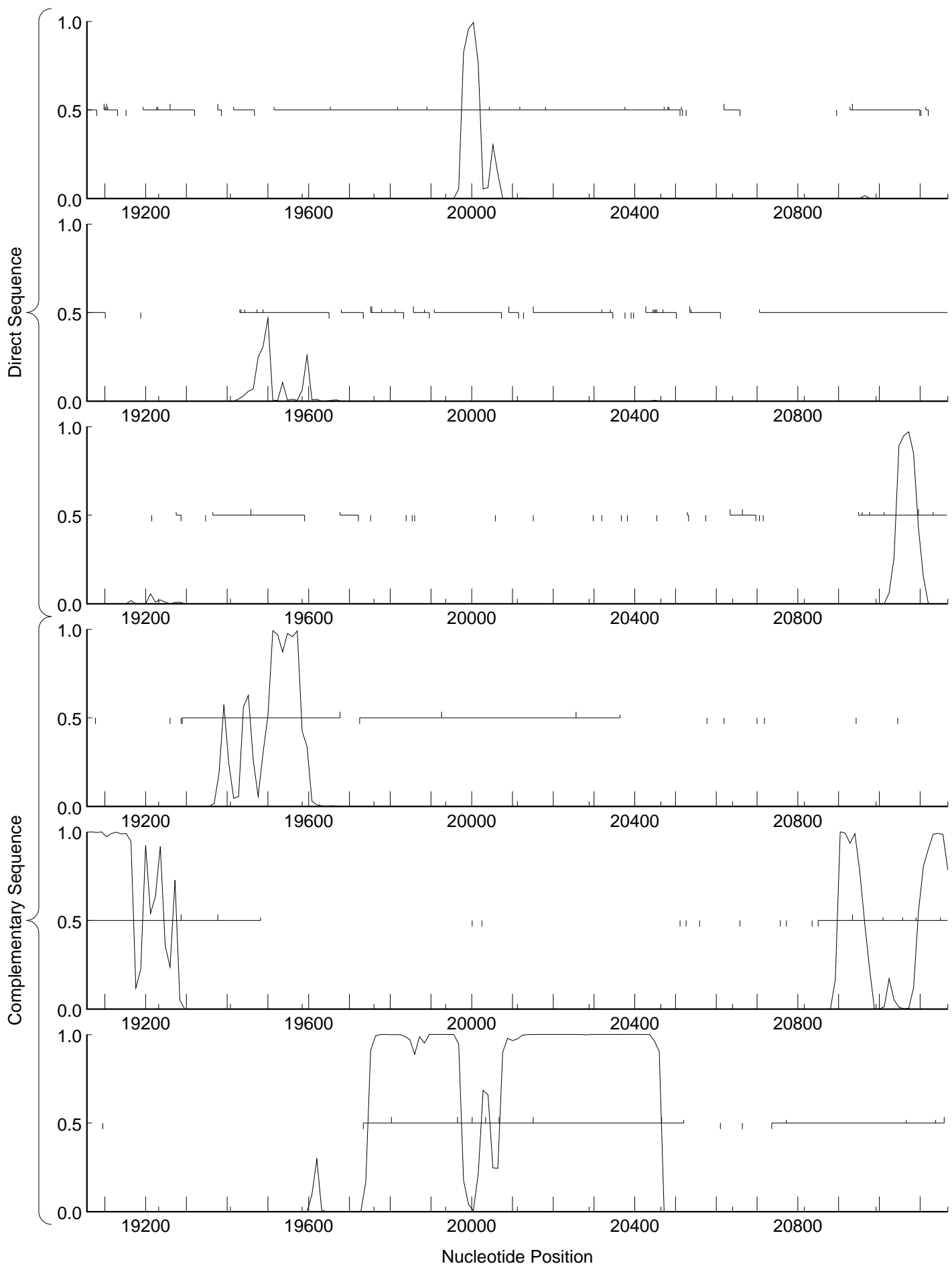


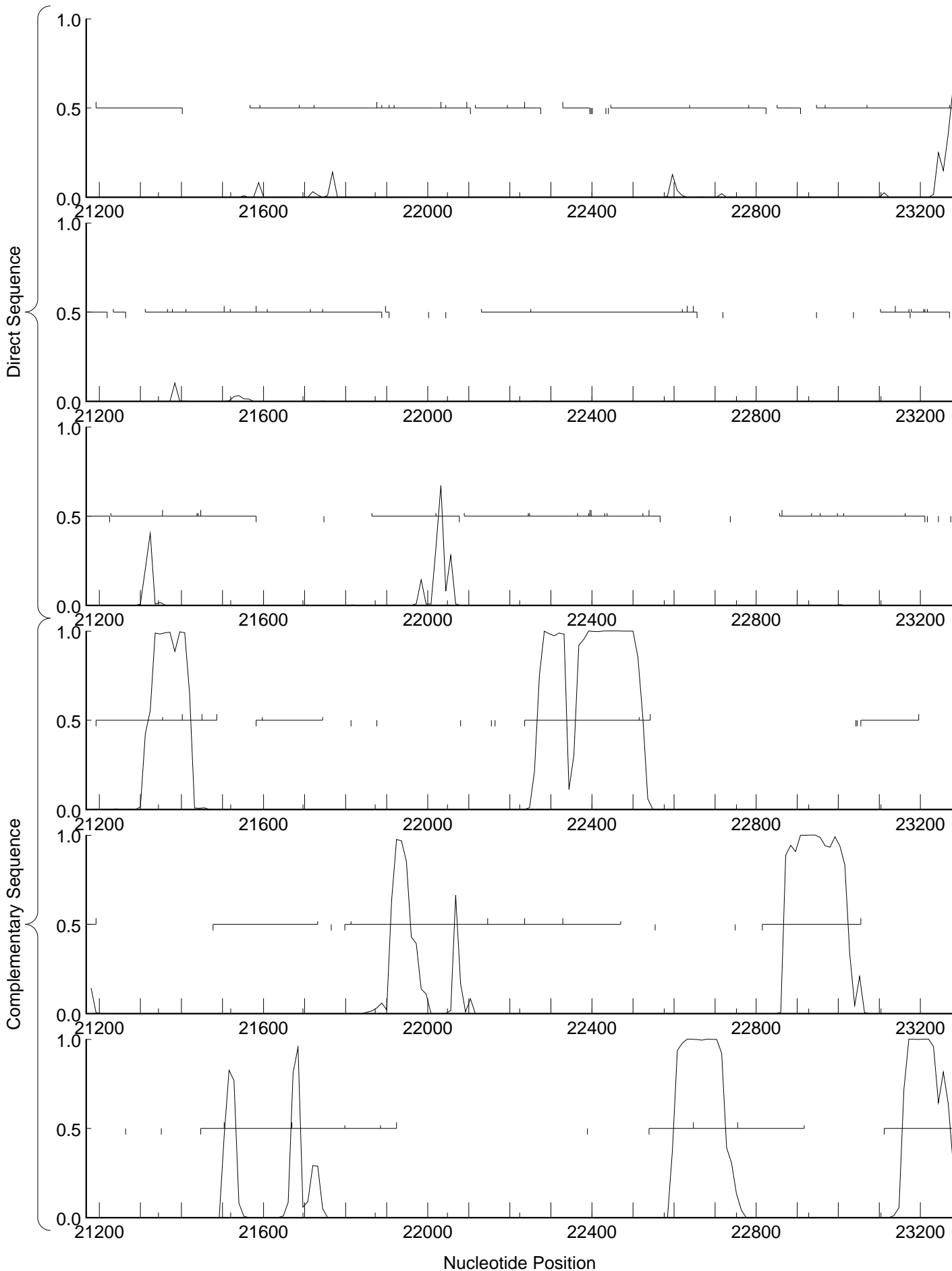




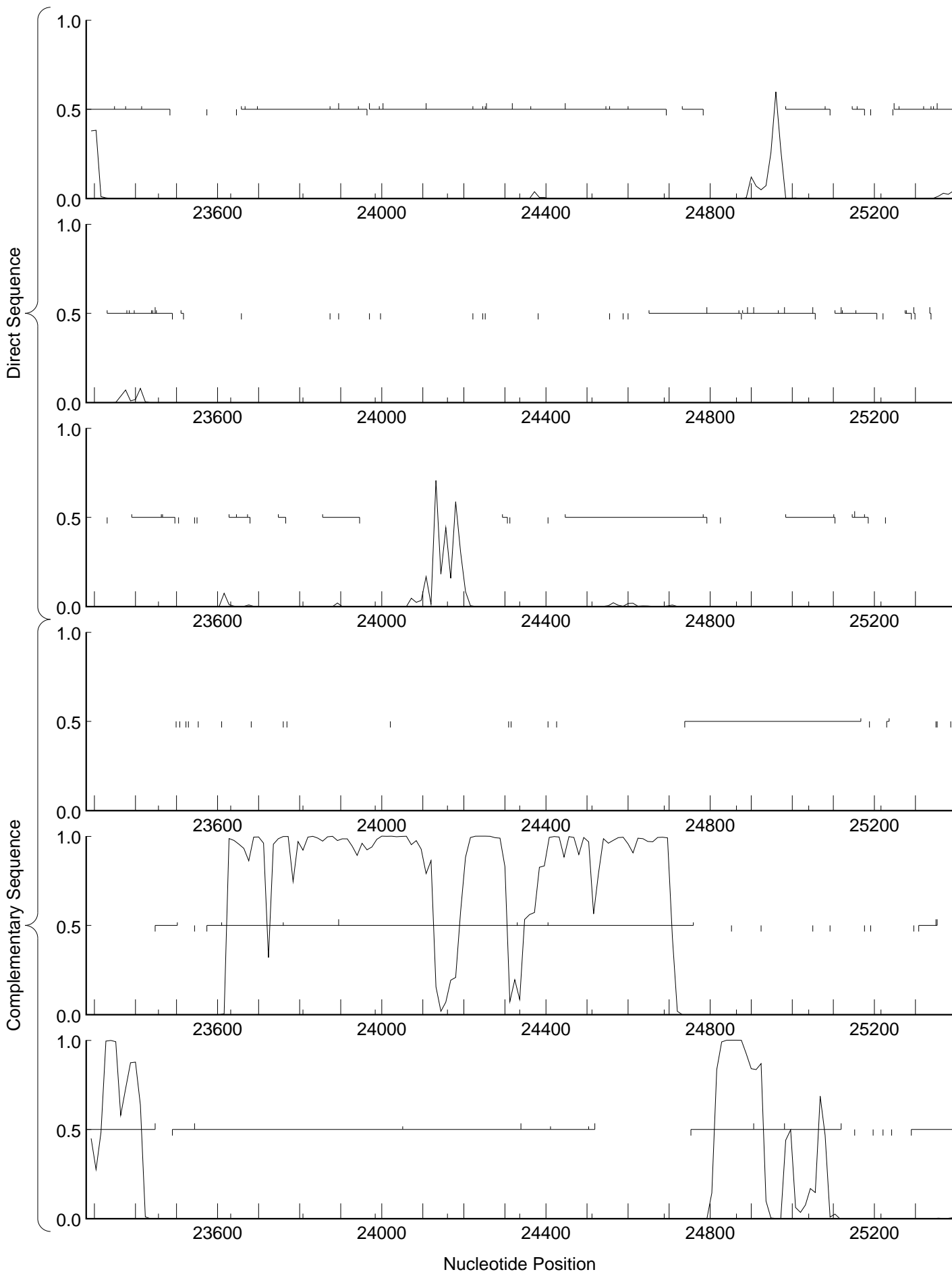


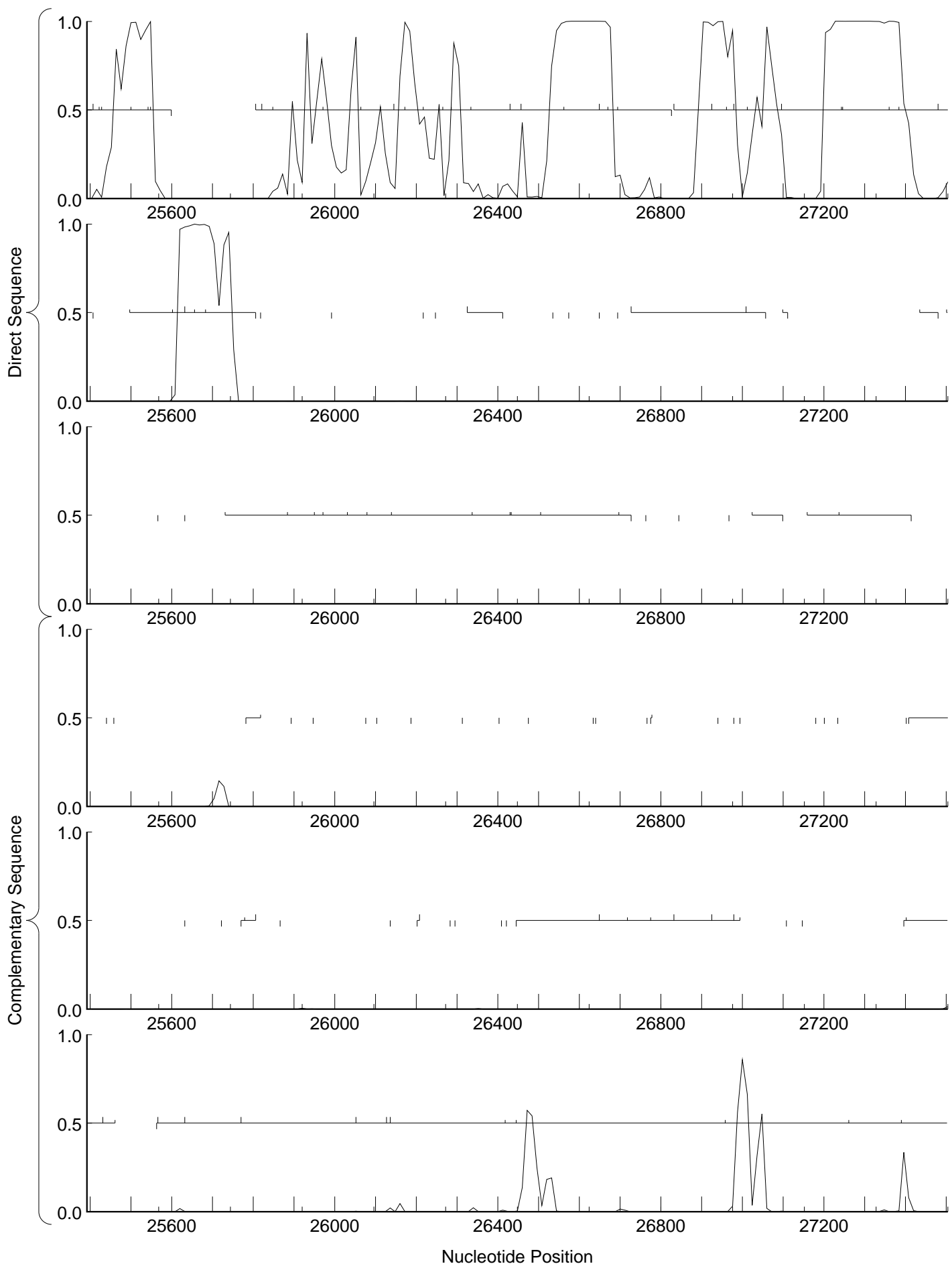


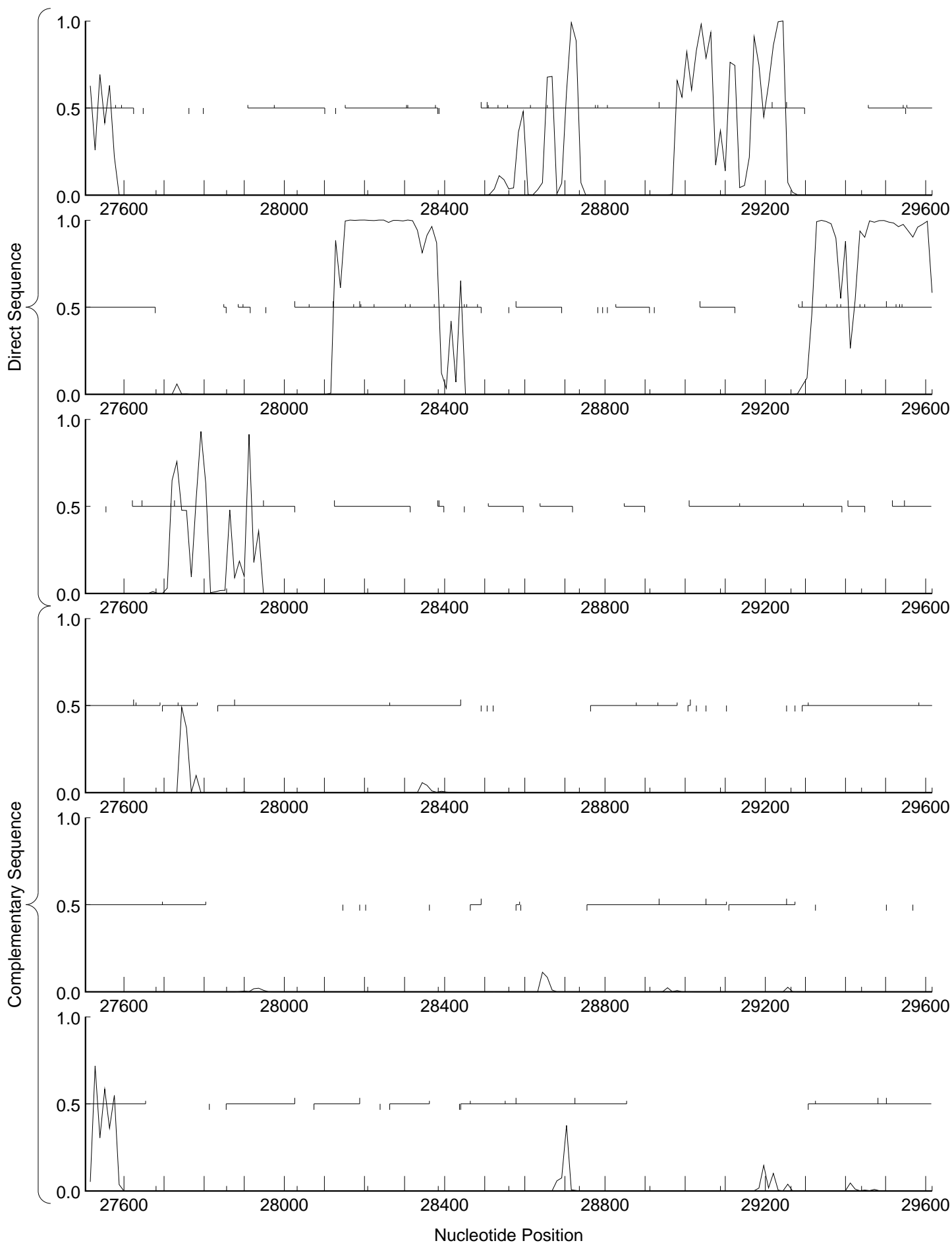




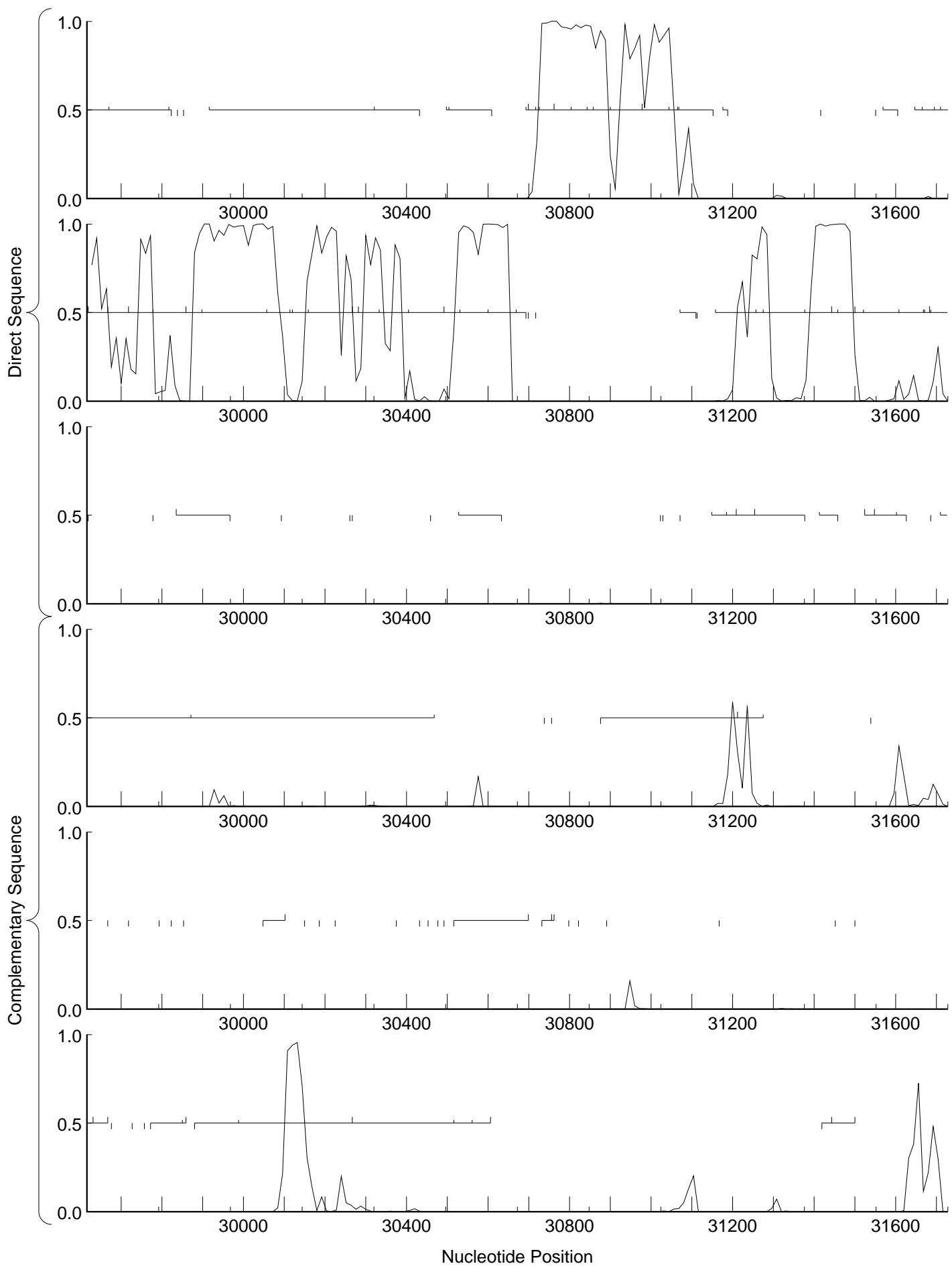
plete sequence, 38449 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 13/20







plete sequence, 38449 bp including 12-base 3' overhang (GAGTTGCCGCA), Cluster AS2, Order 4, Window 96, Step 12, 16/20





plete sequence, 38449 bp including 12-base 3' overhang (GAGTTGCCGCA), Cluster AS2, Order 4, Window 96, Step 12, 17/20

