

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Arthrobacter phage Amelia complete sequence, 38128 bp including 12-base 3' overhang (GAGTTG  
Analysis Date : 10/6/22 at 13:55:52  
Pages : 20  
Sequence Length : 38128 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,

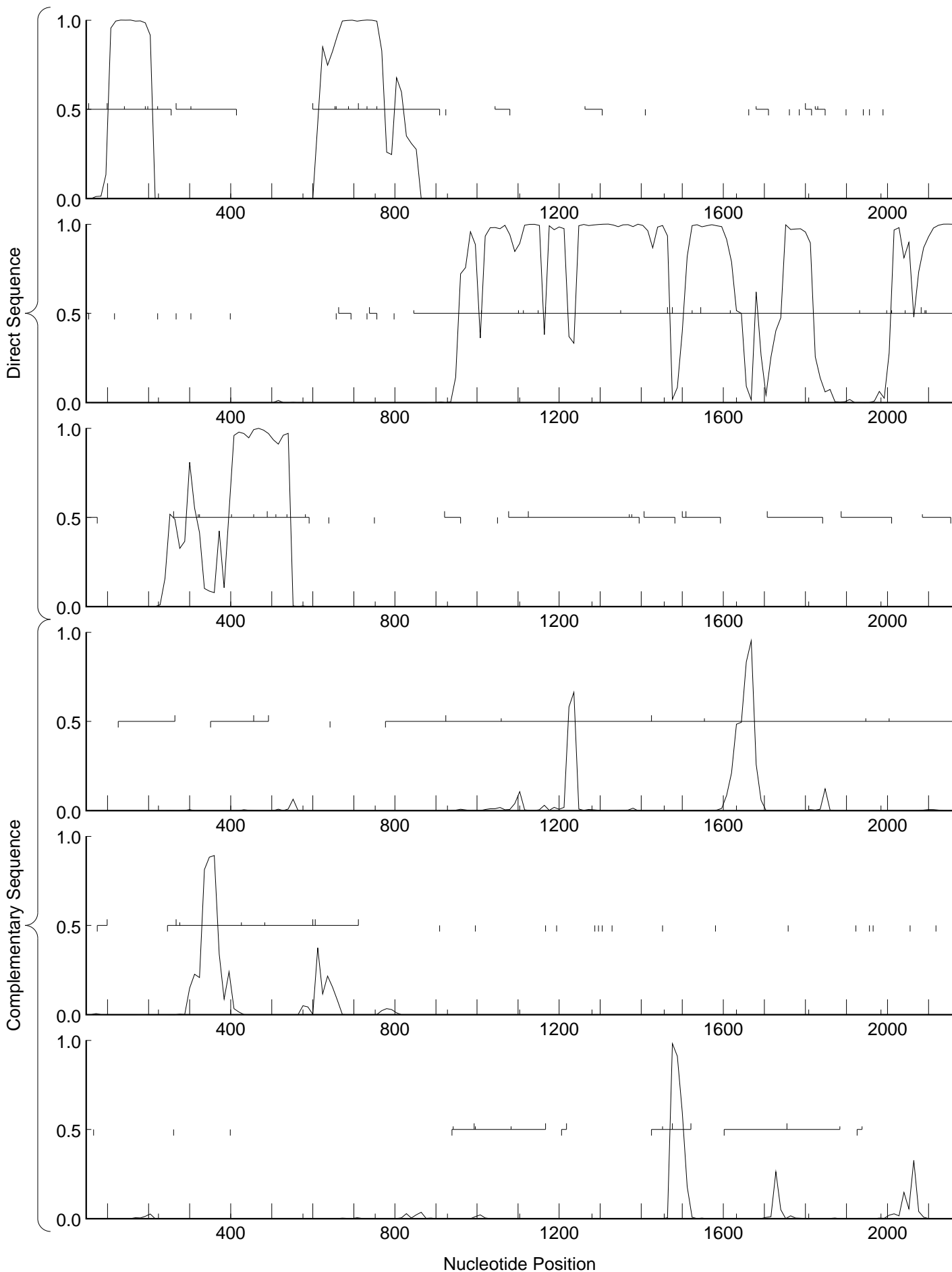
## Matrix notes & comments

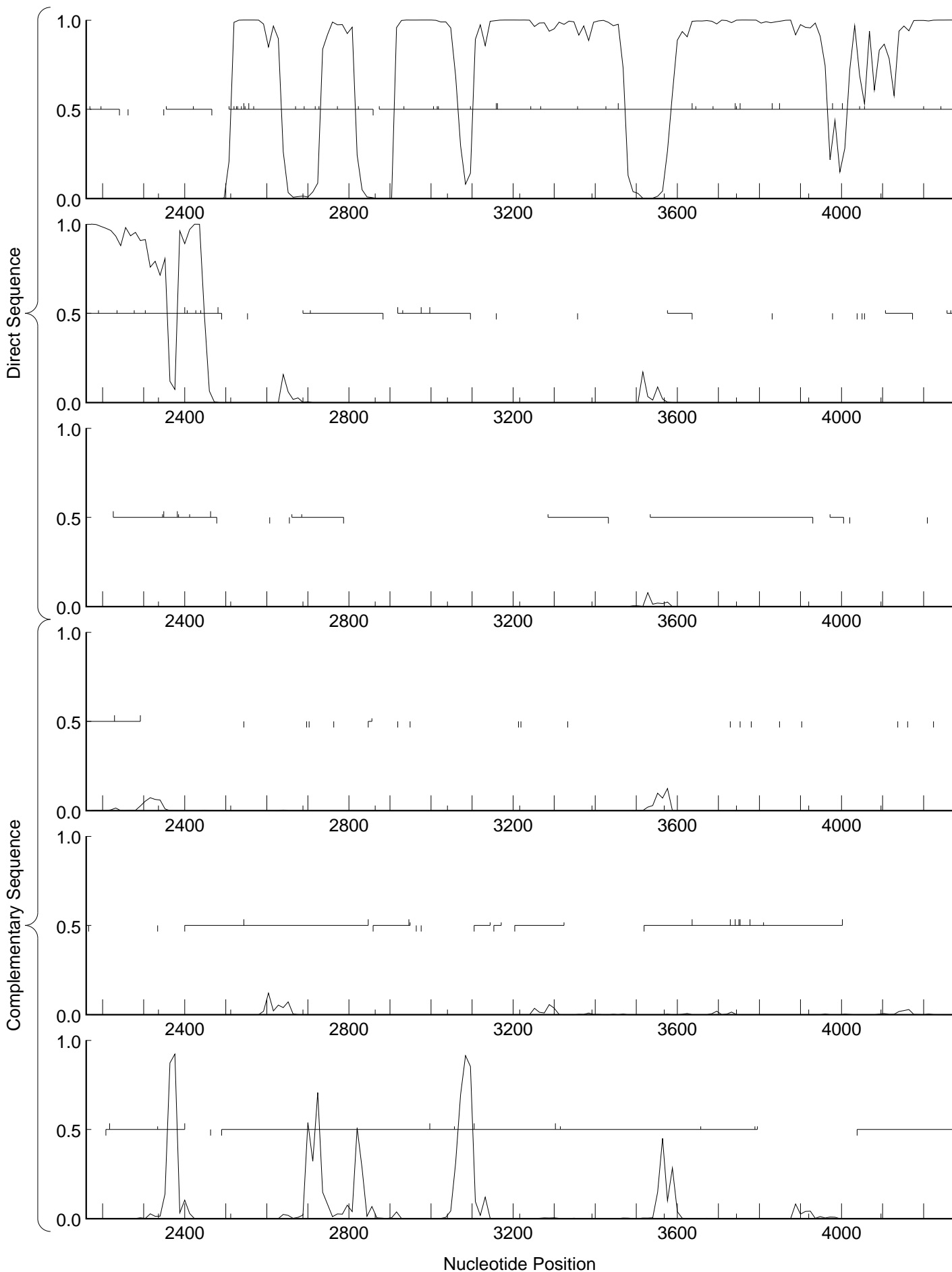
Training set derived by GeneMarkS, 4.30 October 2014  
Sat Sep 8 00:22:00 2018

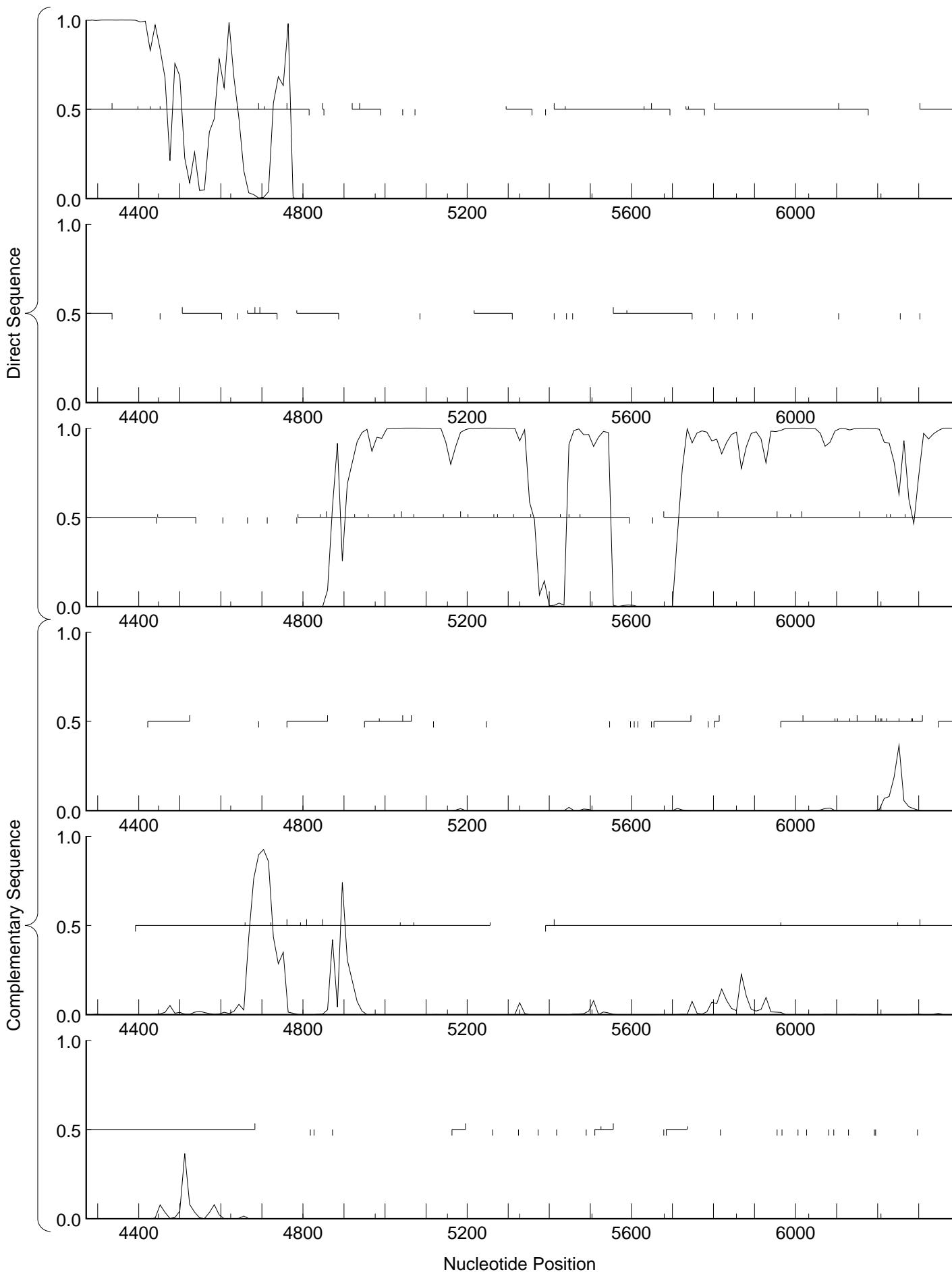
## MATRIX INFORMATION

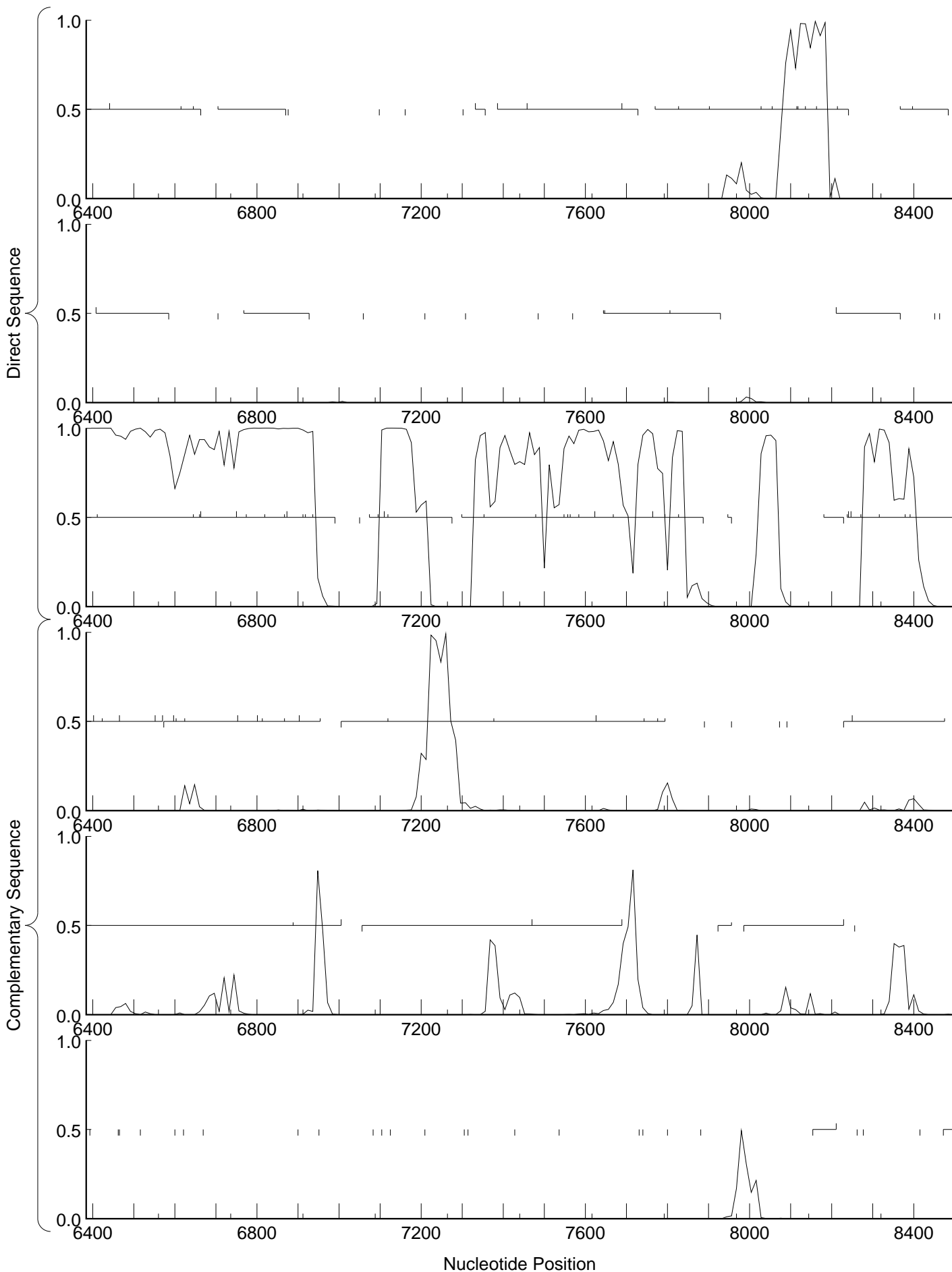
Matrix : -  
Author : -  
Order : 4

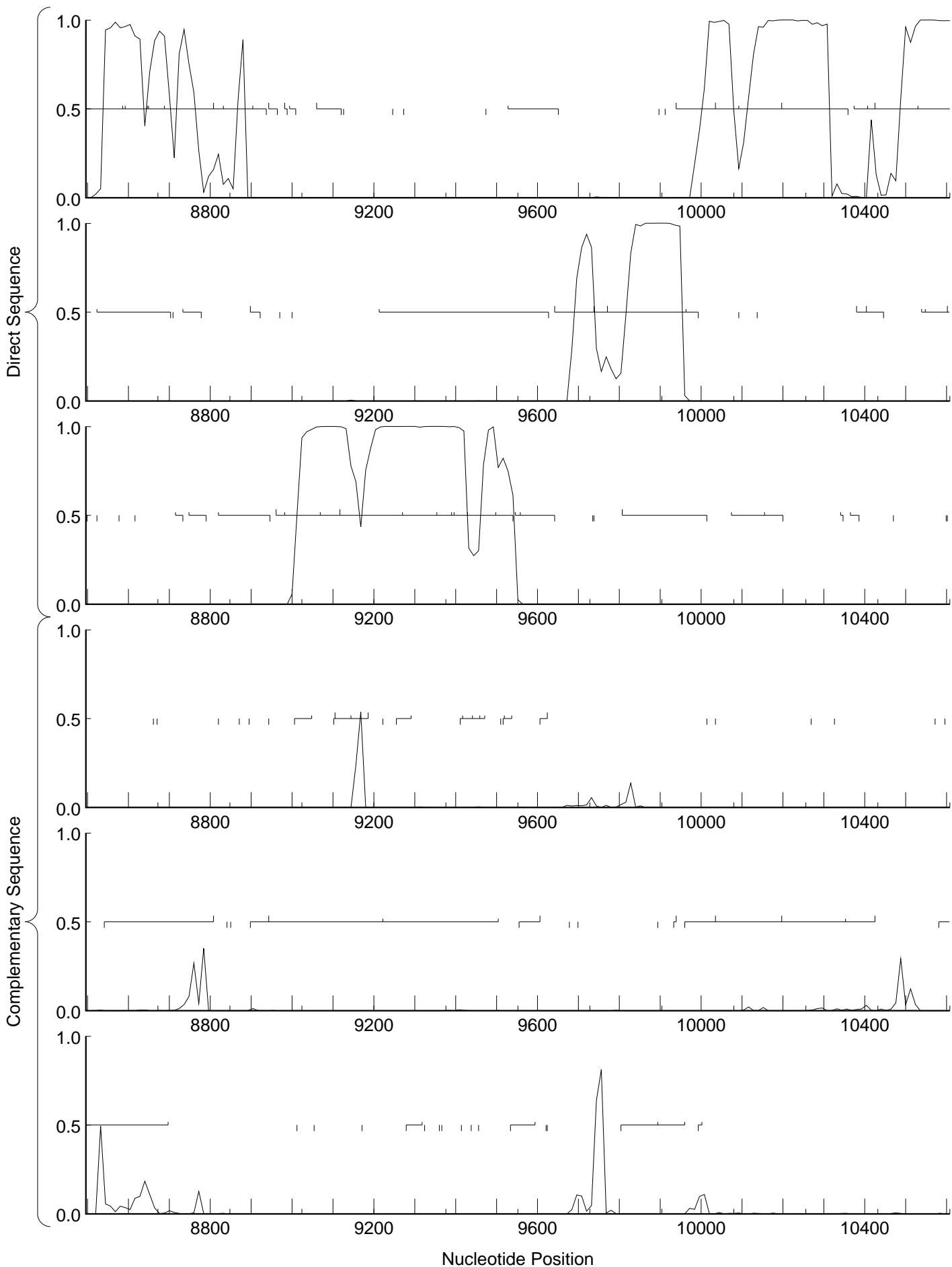
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

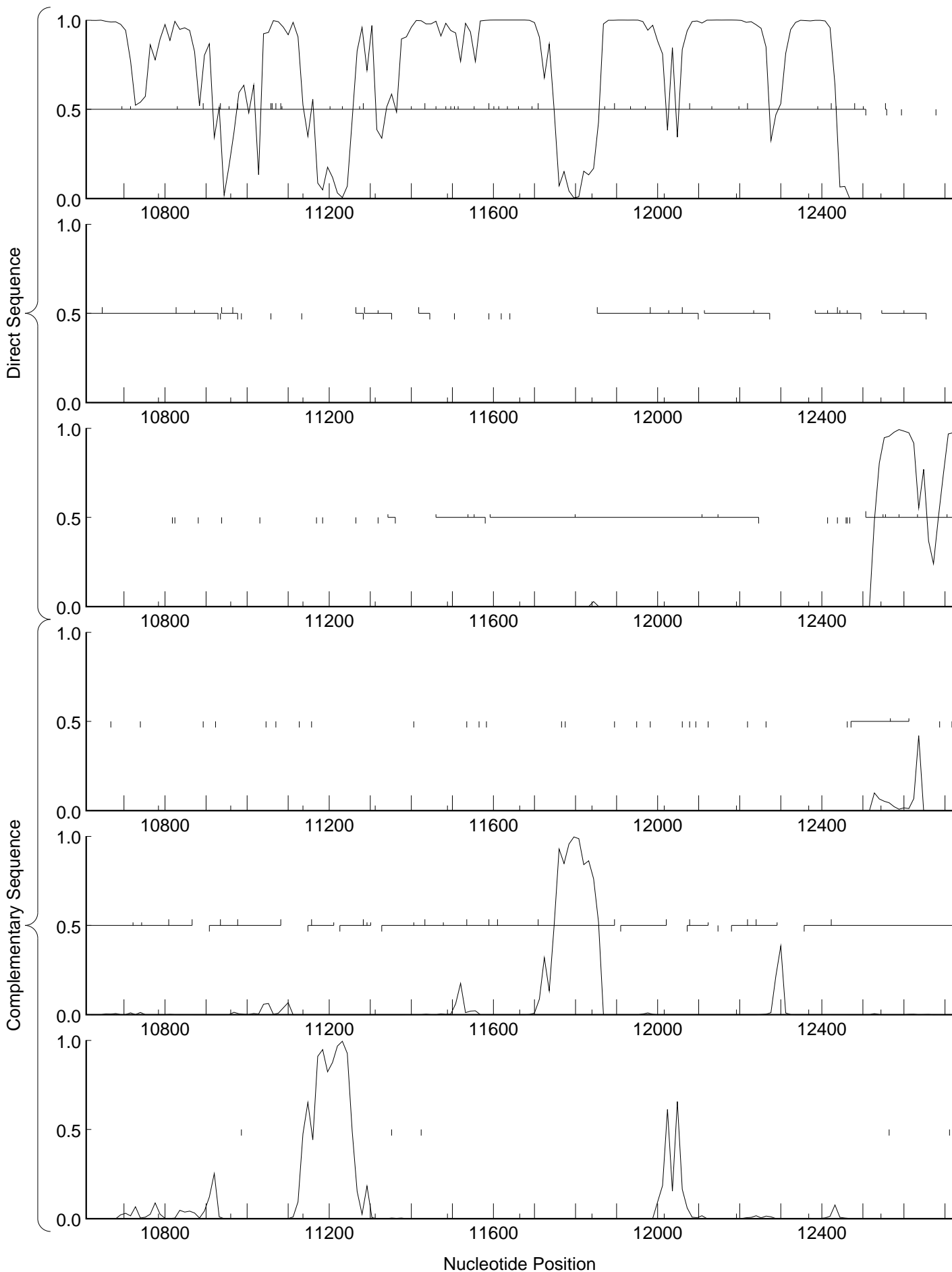


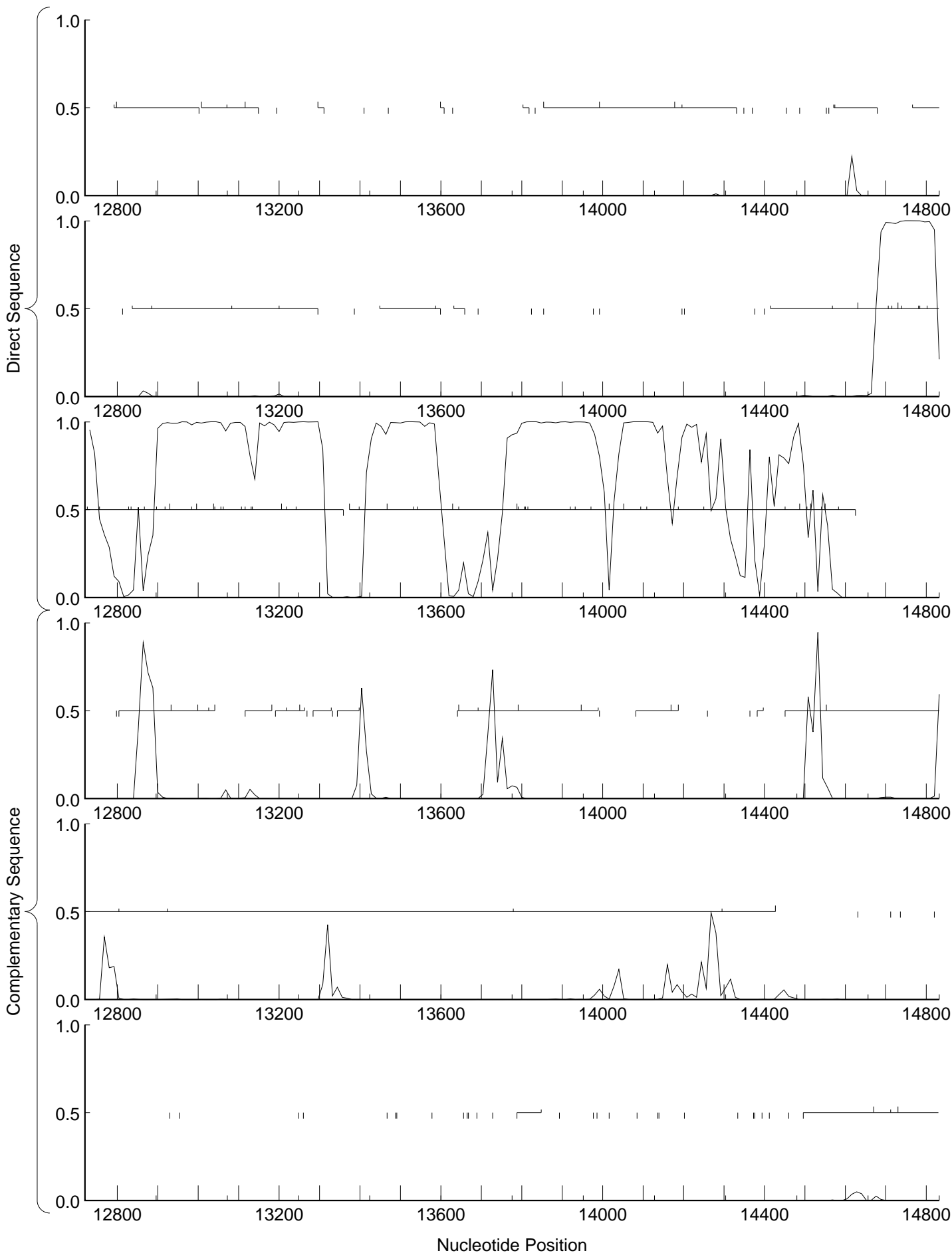




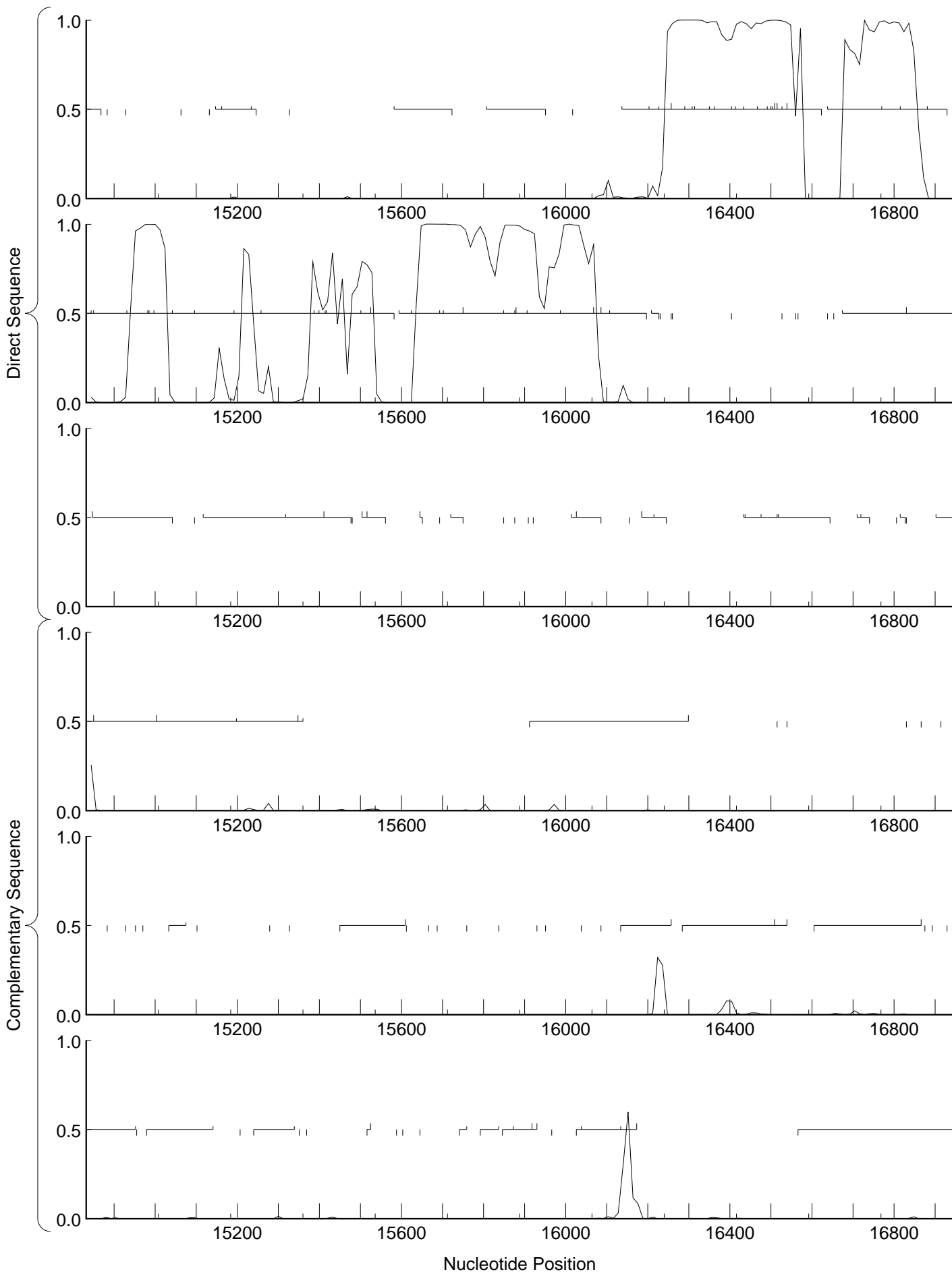


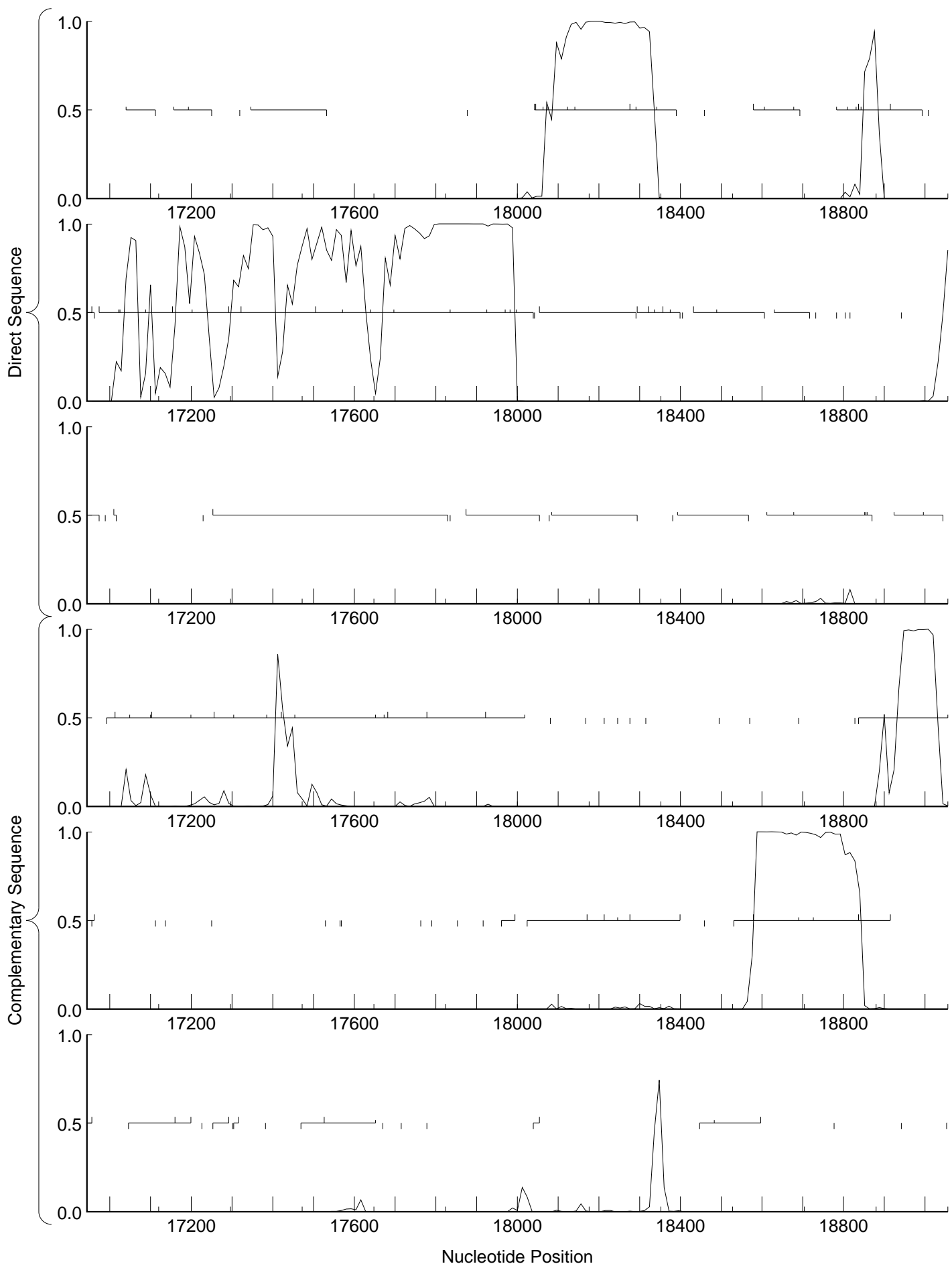


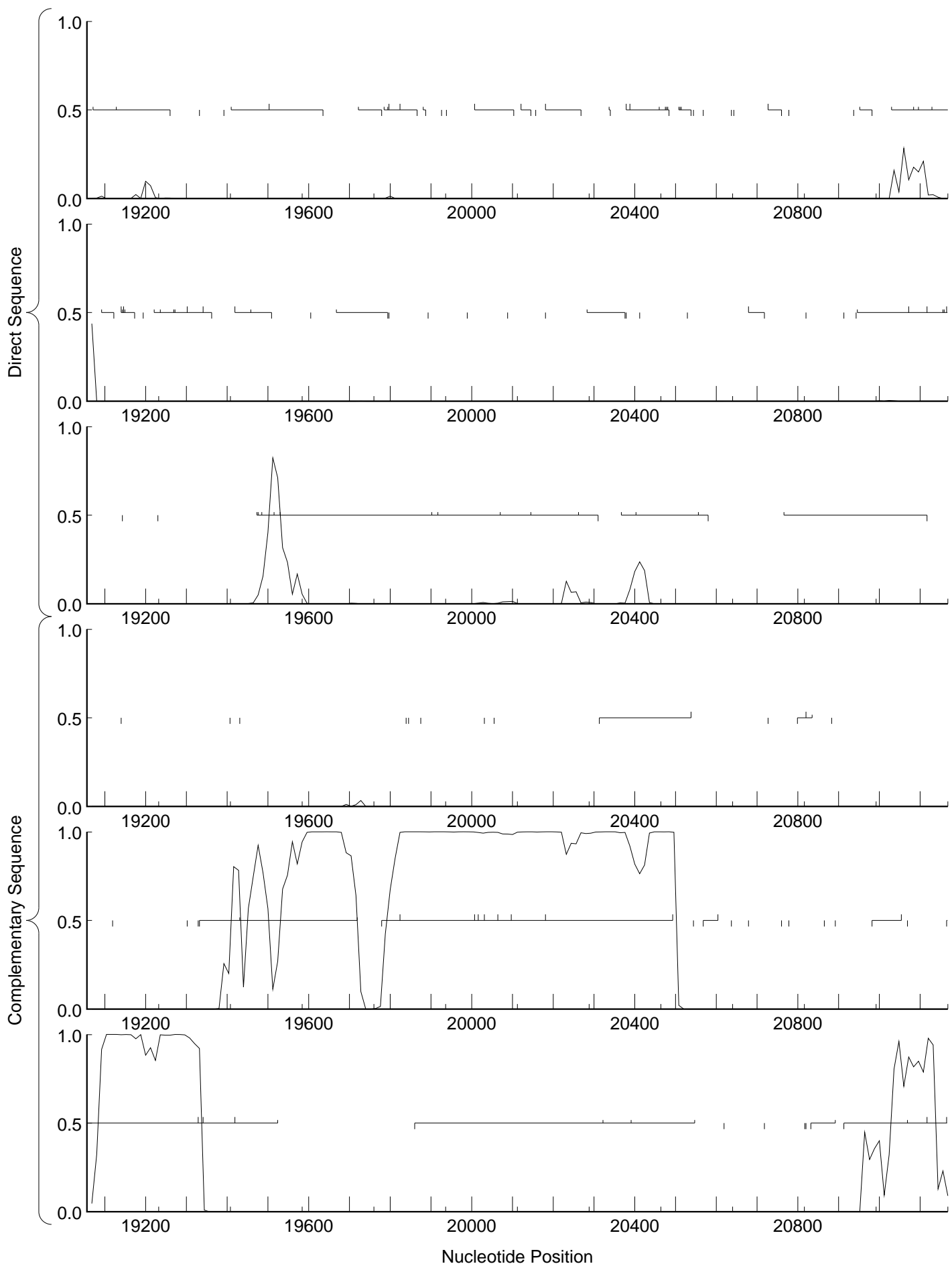


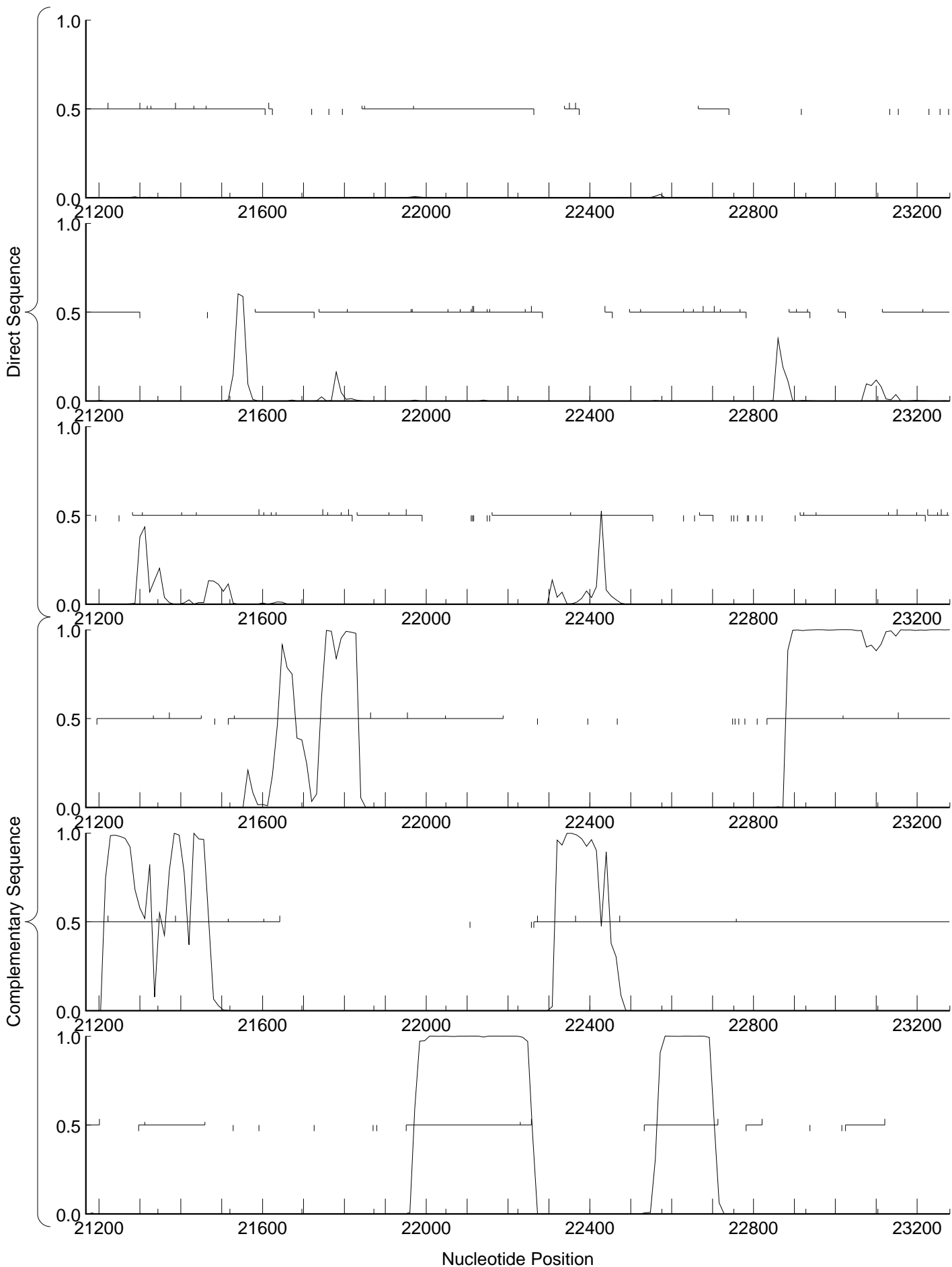


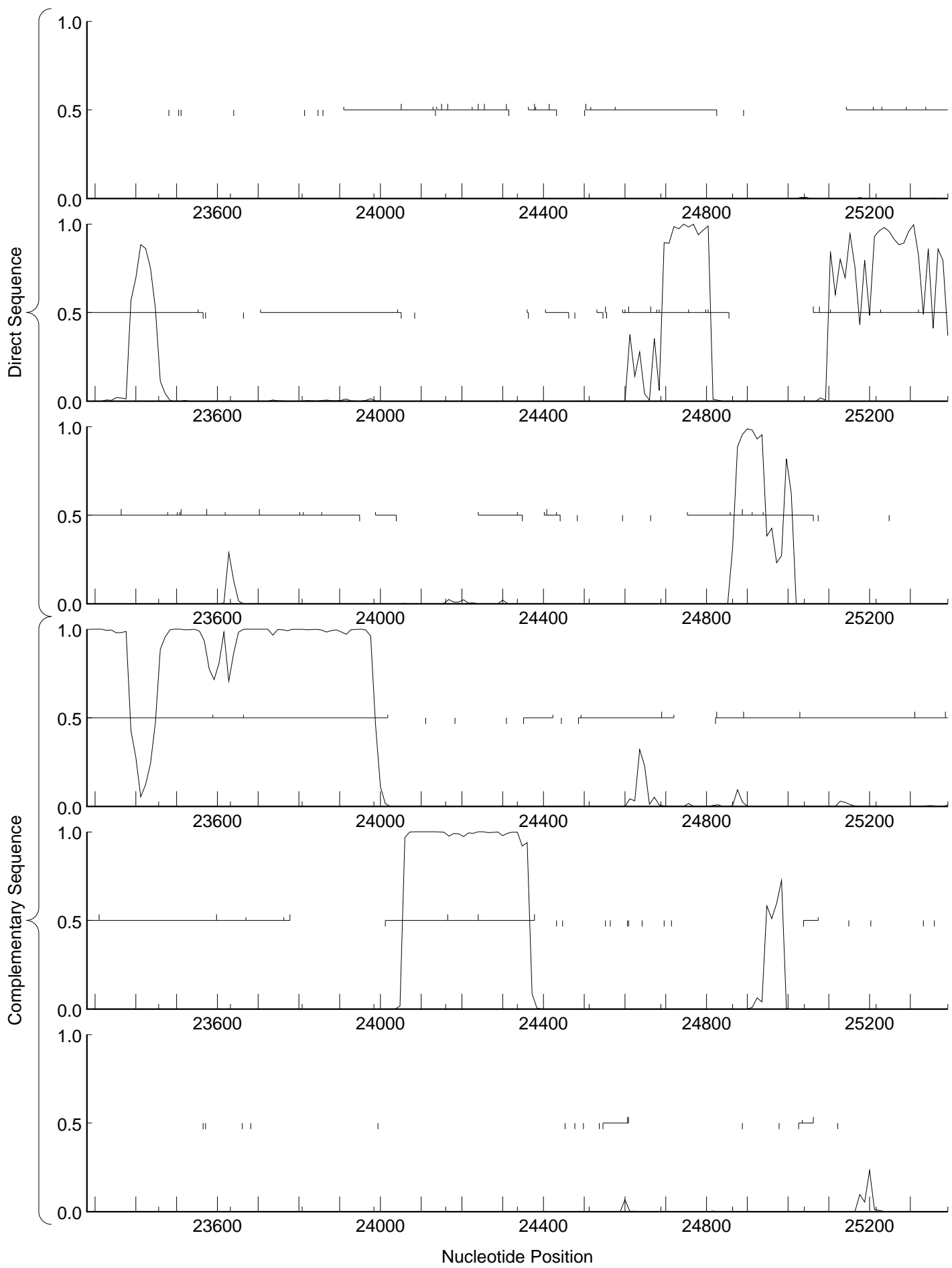




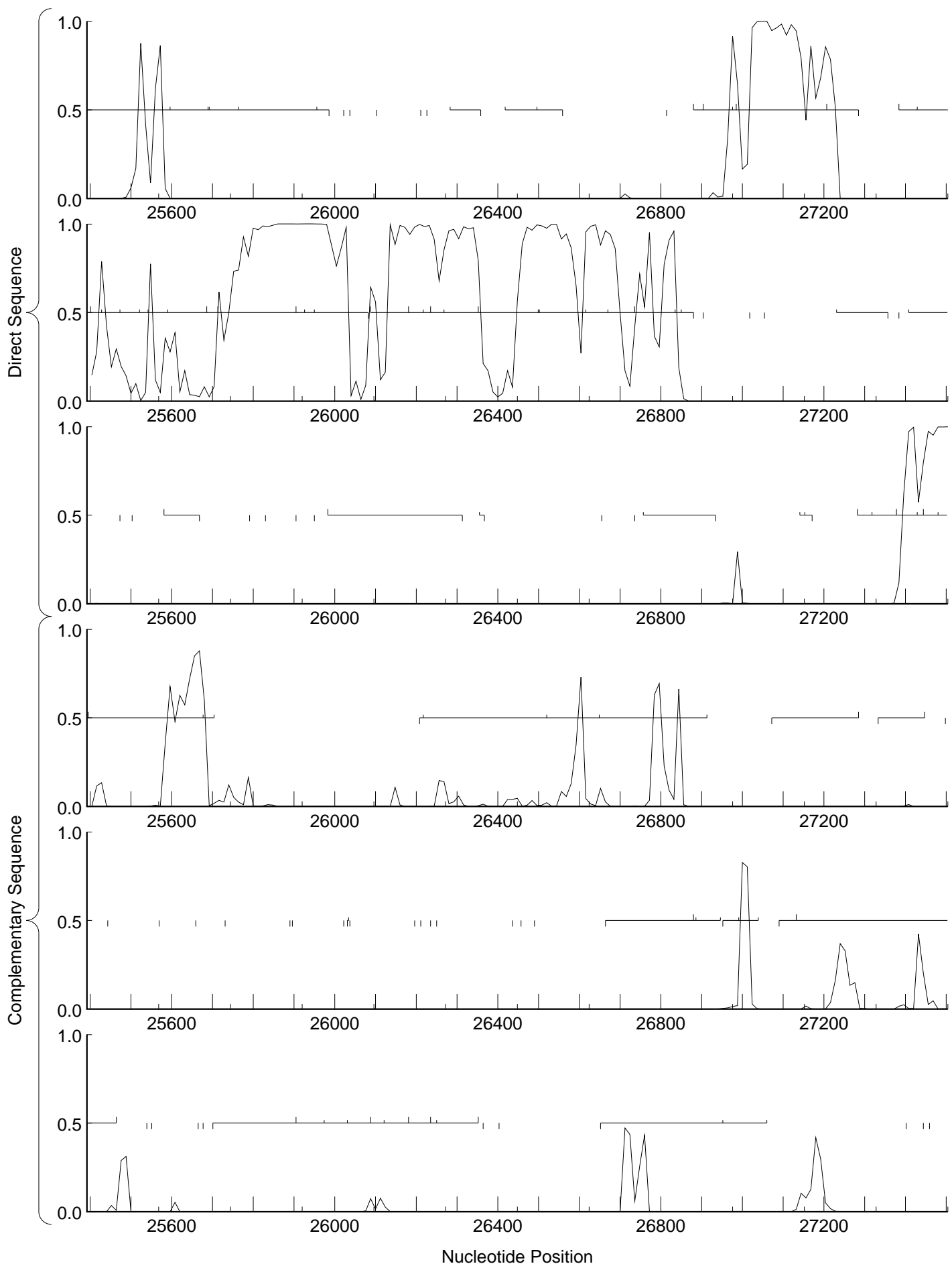




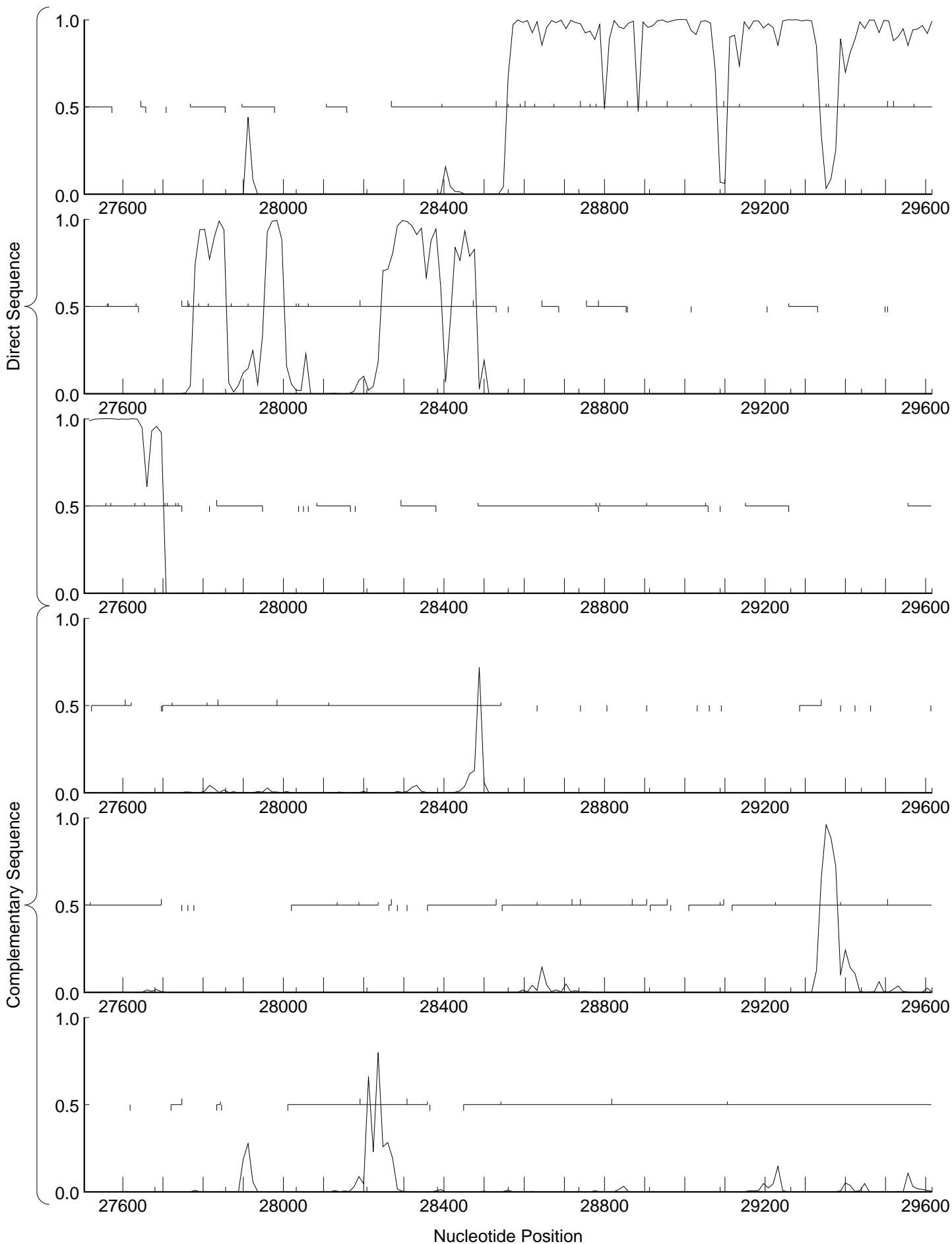




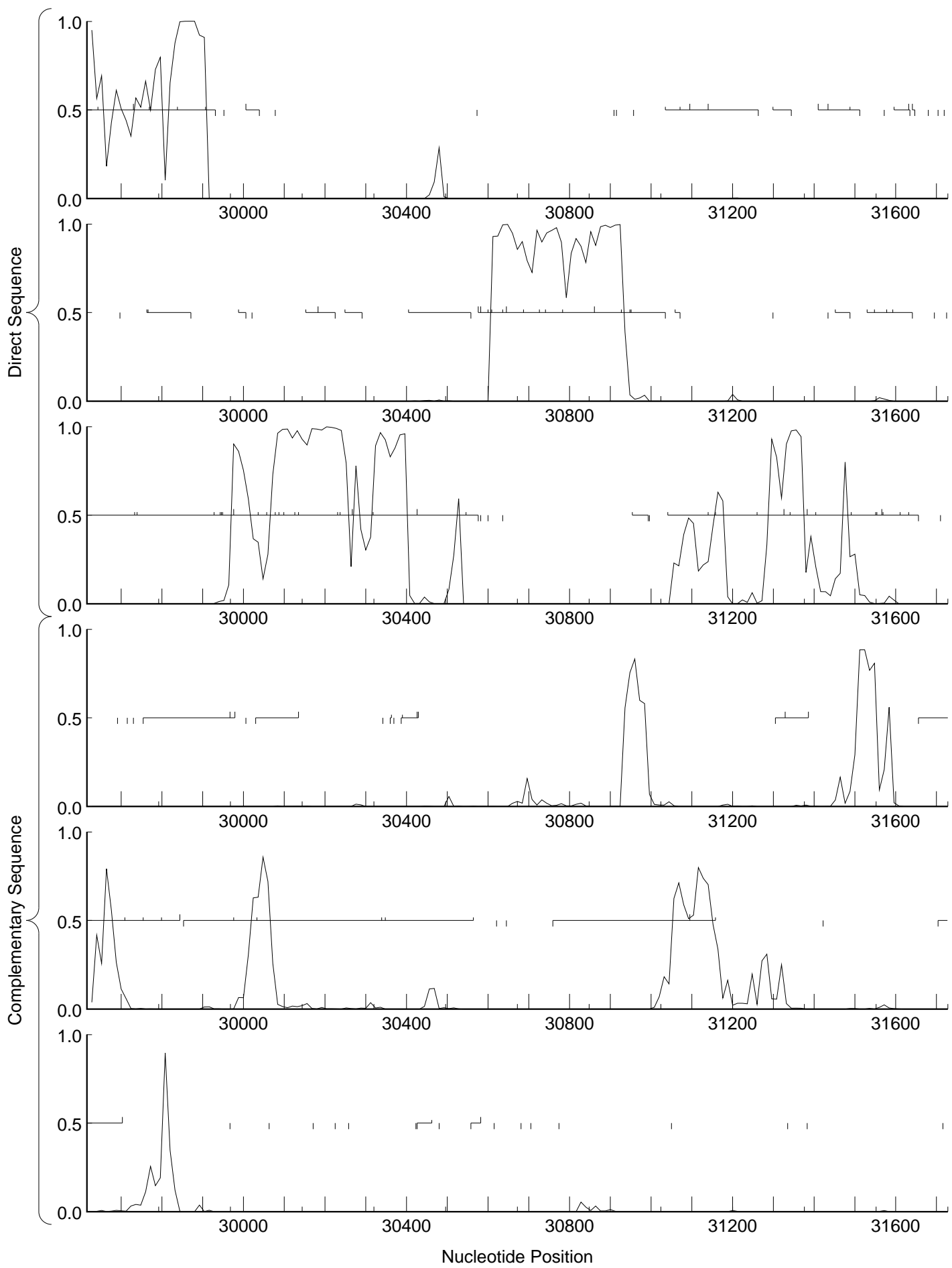
plete sequence, 38128 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 14/20



plete sequence, 38128 bp including 12-base 3' overhang (GAGTTGCCGCA), Cluster AS2, Order 4, Window 96, Step 12, 15/20



plete sequence, 38128 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 16/20





plete sequence, 38128 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 17/20

