

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Gordonia phage FroggyToad complete sequence, 46692 bp including 11-base 3' overhang (TACCAG  
Analysis Date : 3/25/25 at 15:07:27  
Pages : 24  
Sequence Length : 46692 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,

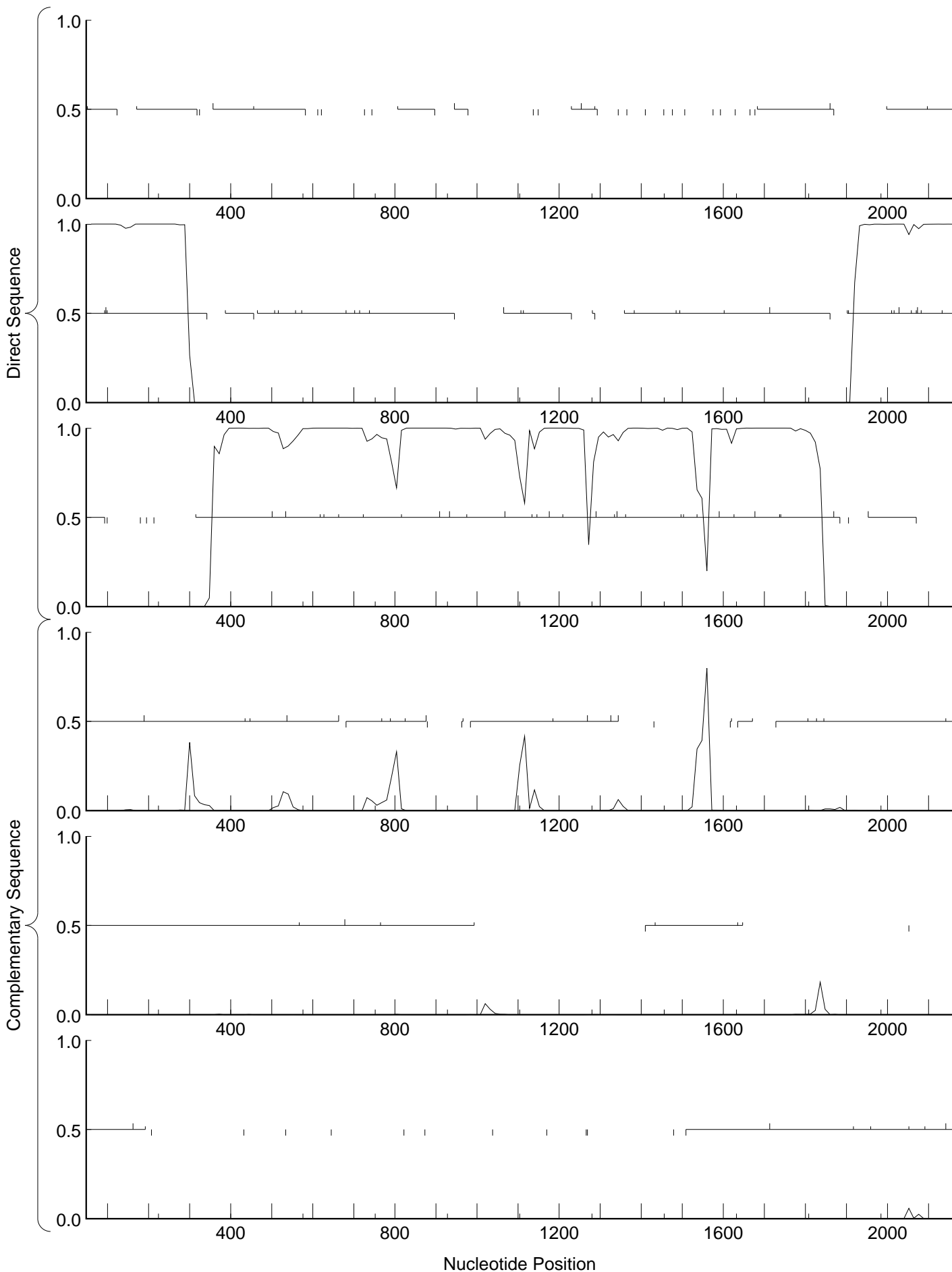
## MATRIX INFORMATION

Matrix : -  
Author : -  
Order : 4

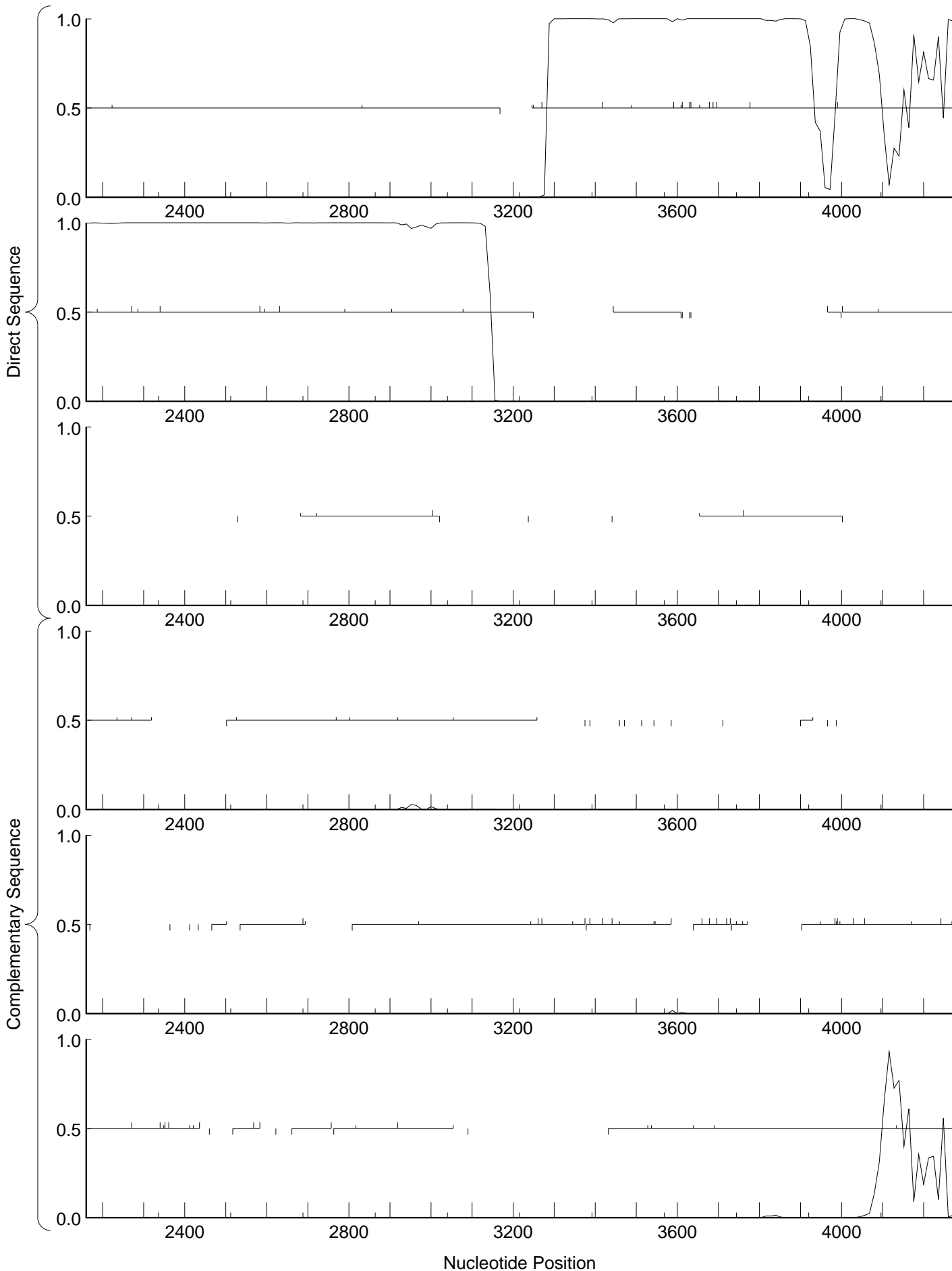
Send questions / comments to:  
Dr. M. Borodovsky  
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Atlanta, GA 30332-0230

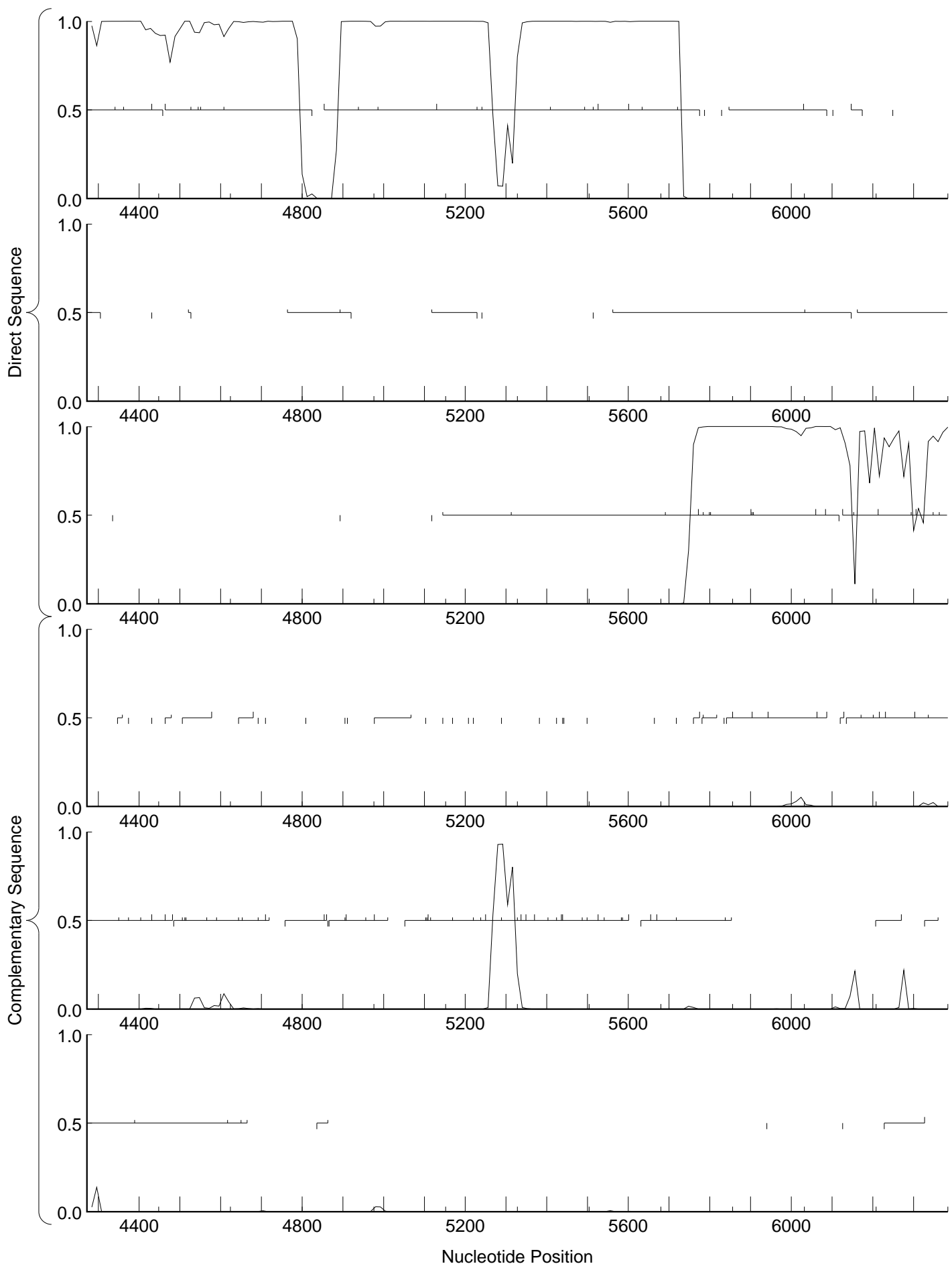
### Matrix notes & comments

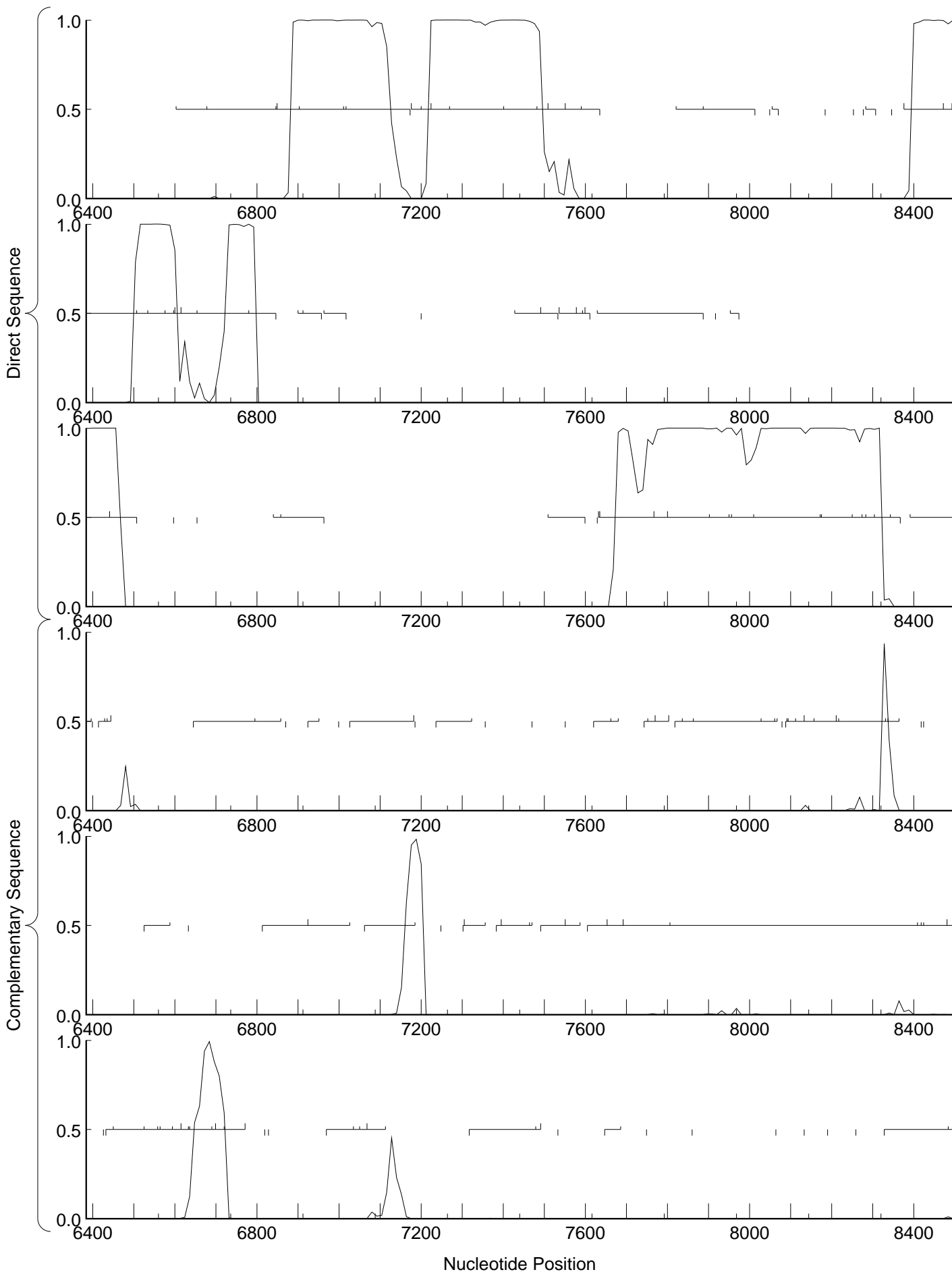
Training set derived by GeneMarkS, 4.30 October 2014  
Sat Sep 8 00:36:14 2018

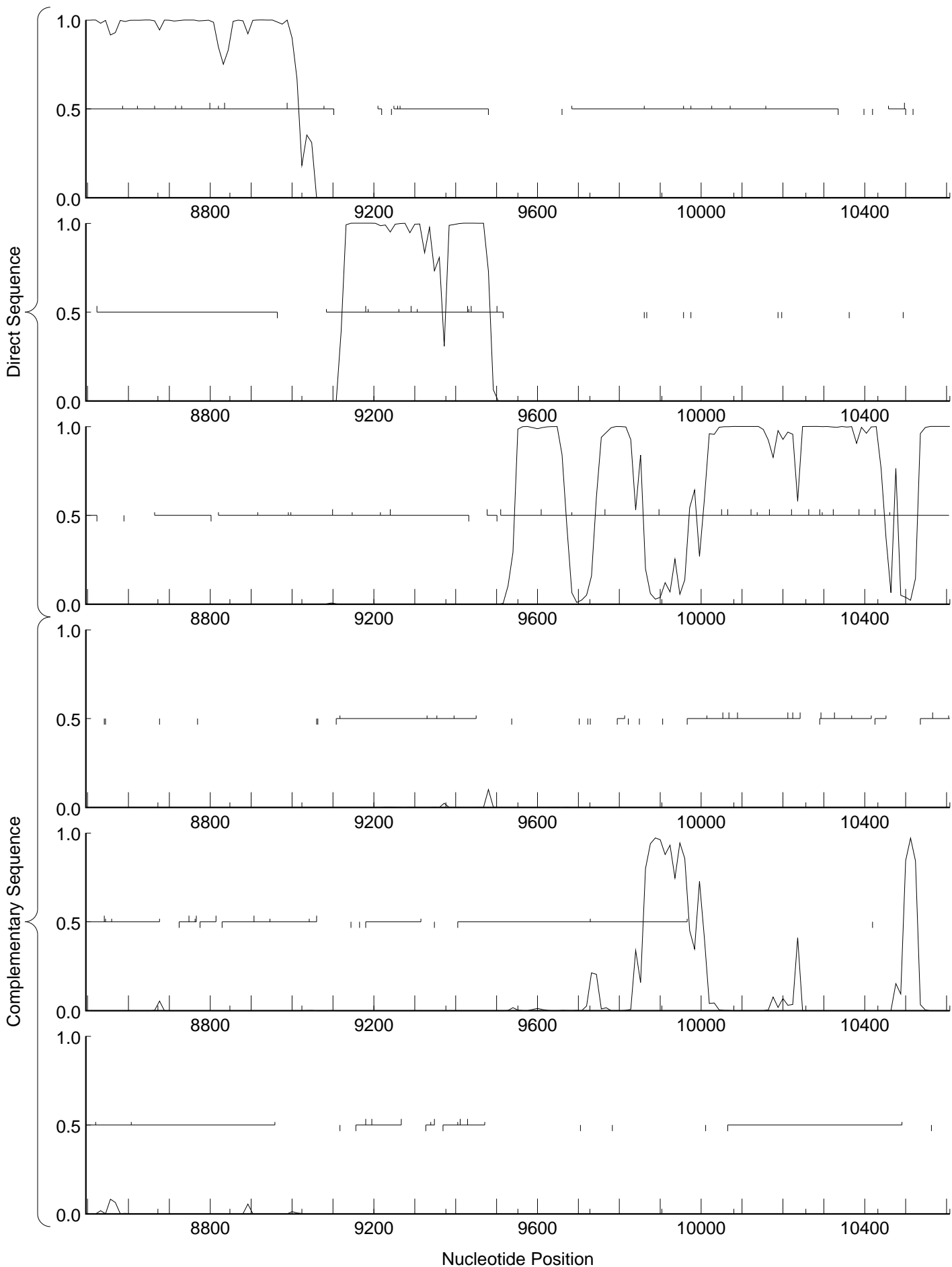


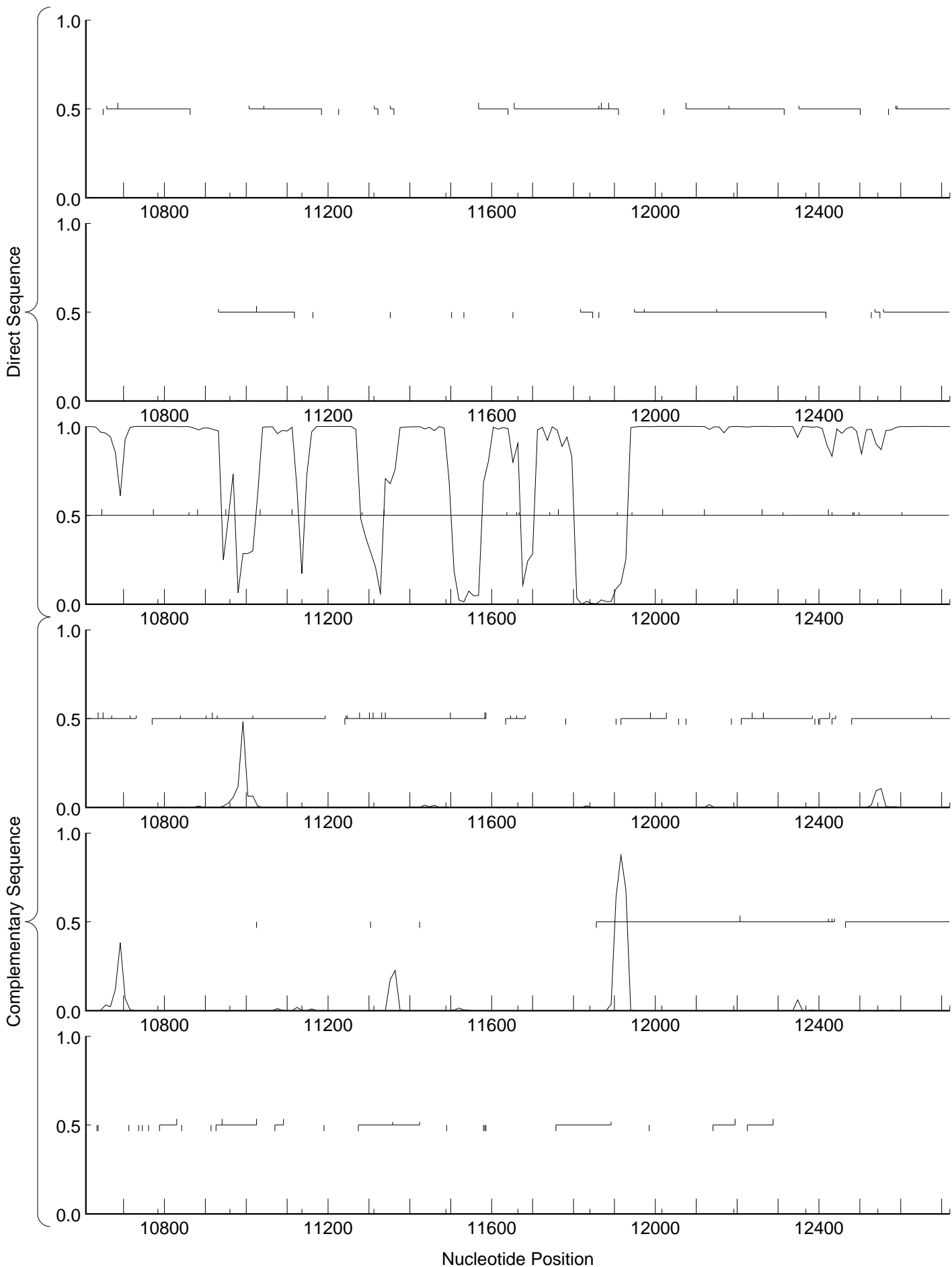
complete sequence, 46692 bp including 11-base 3' overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 3/24

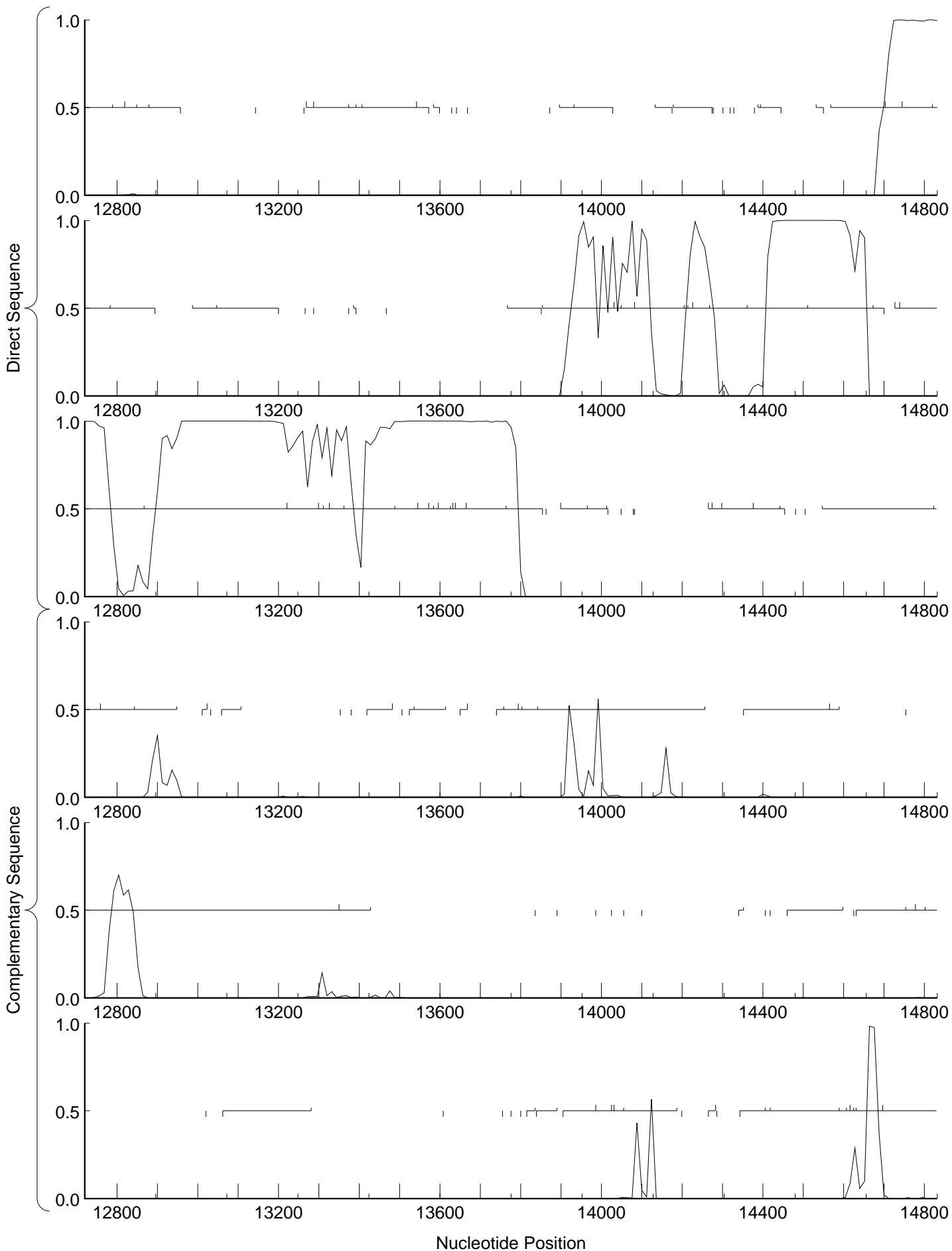




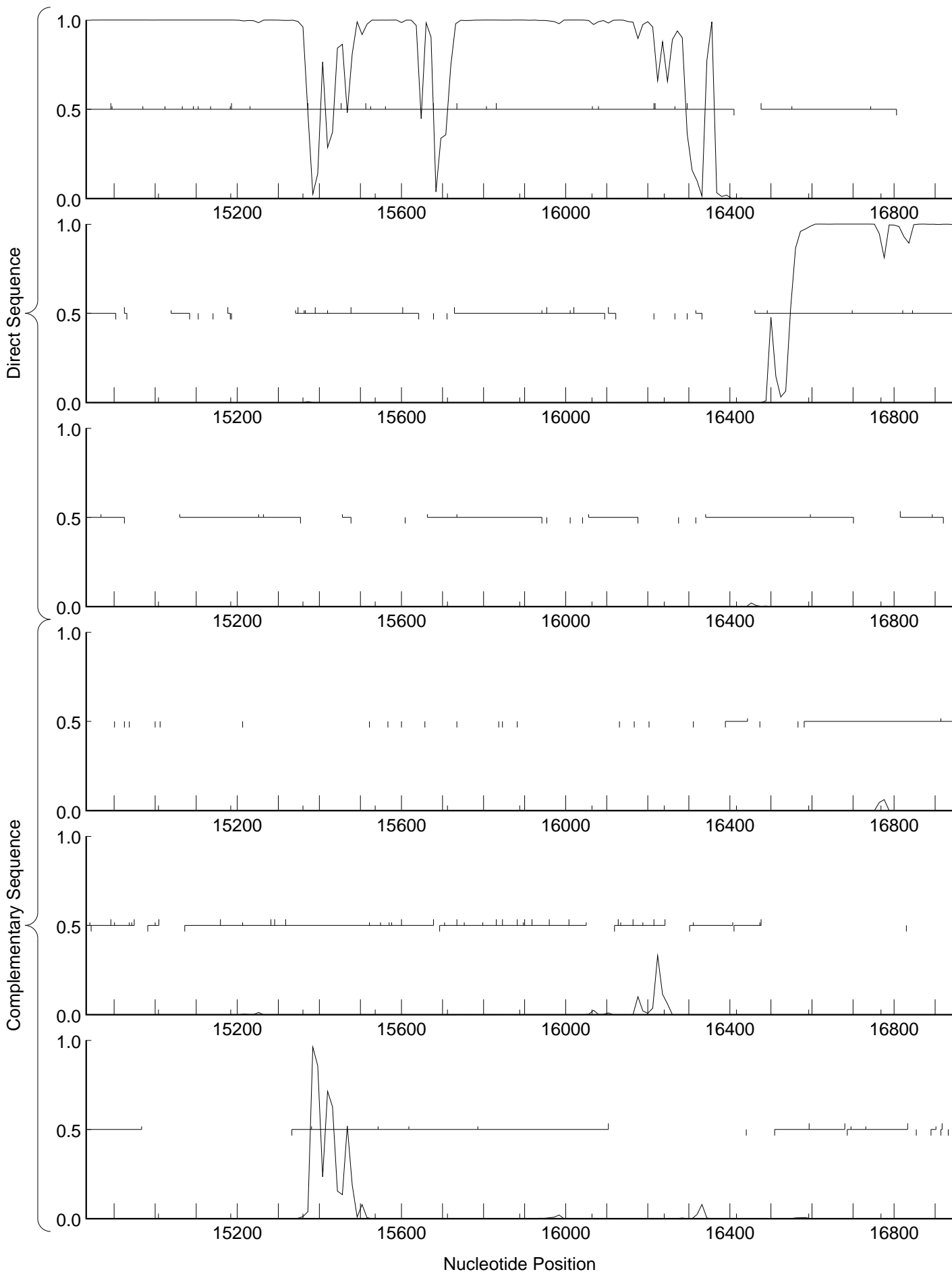


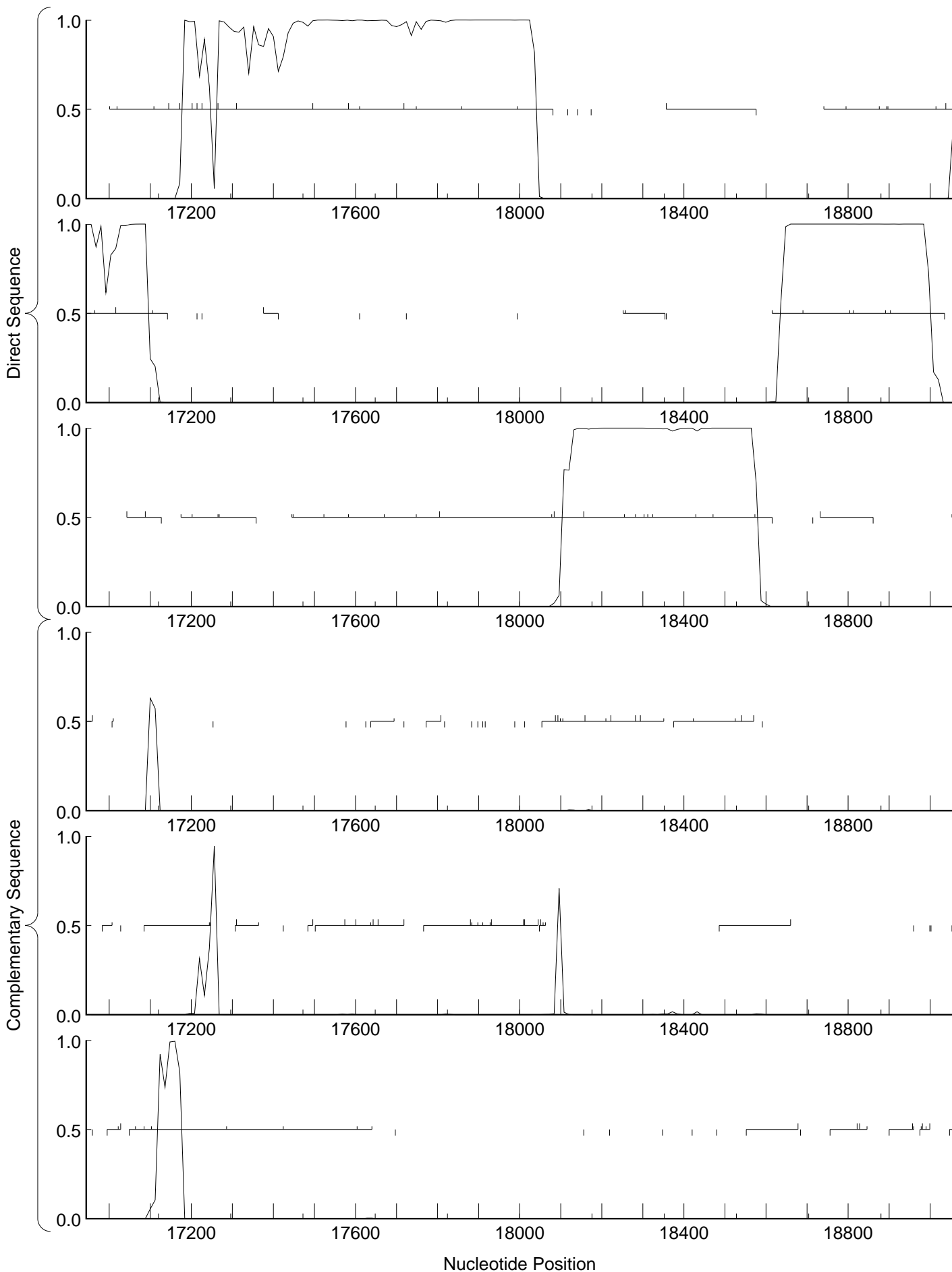


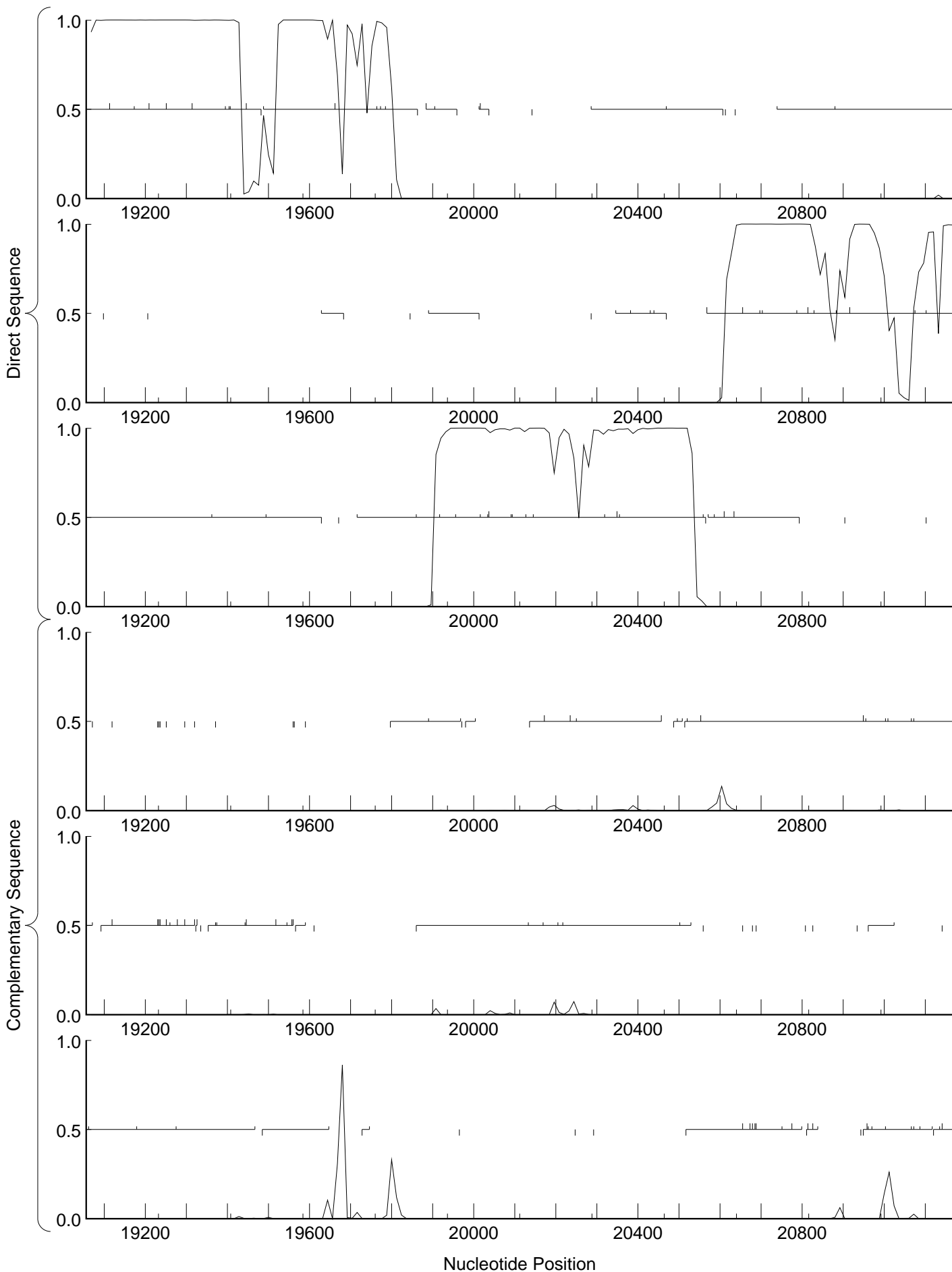


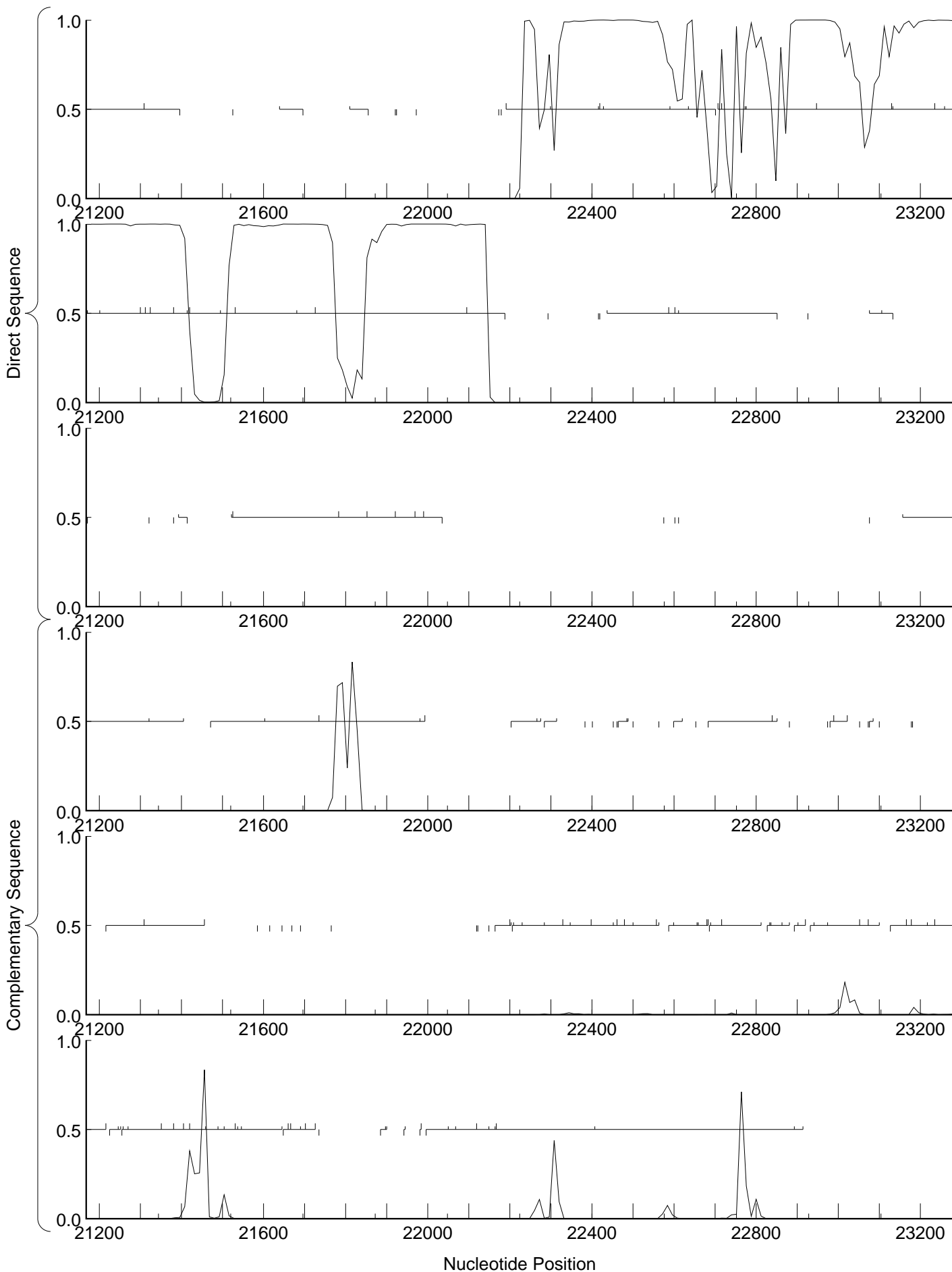


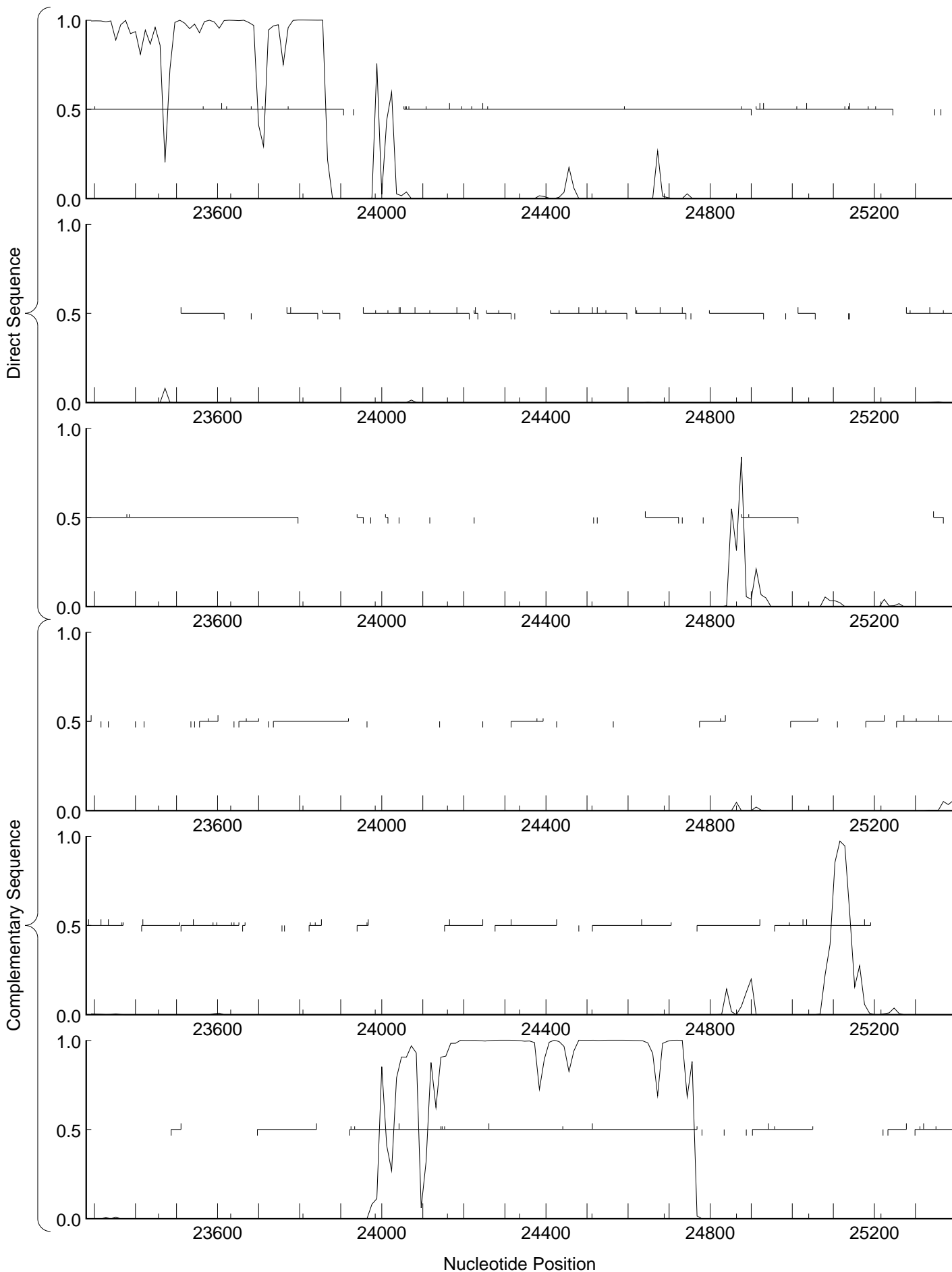


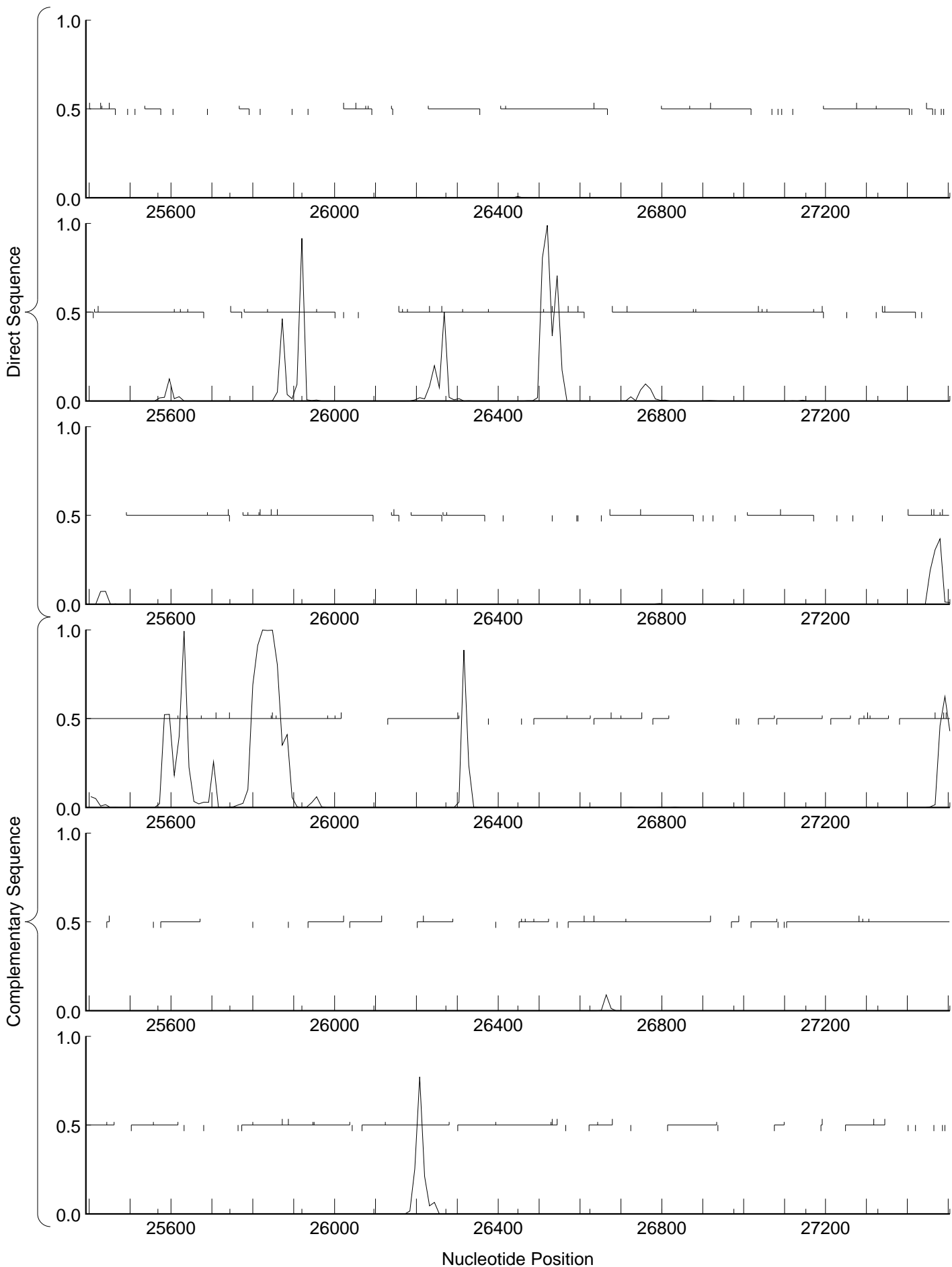


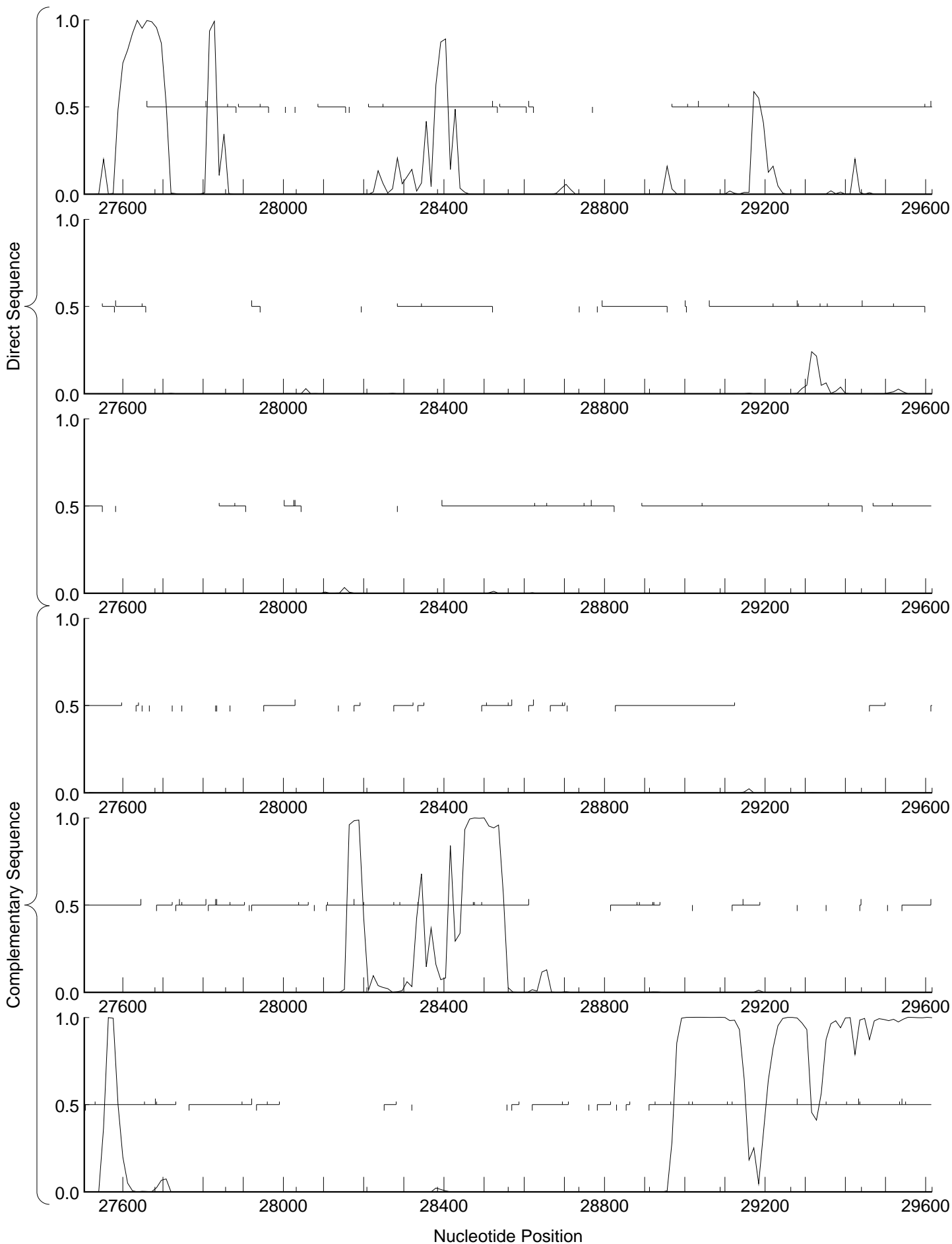


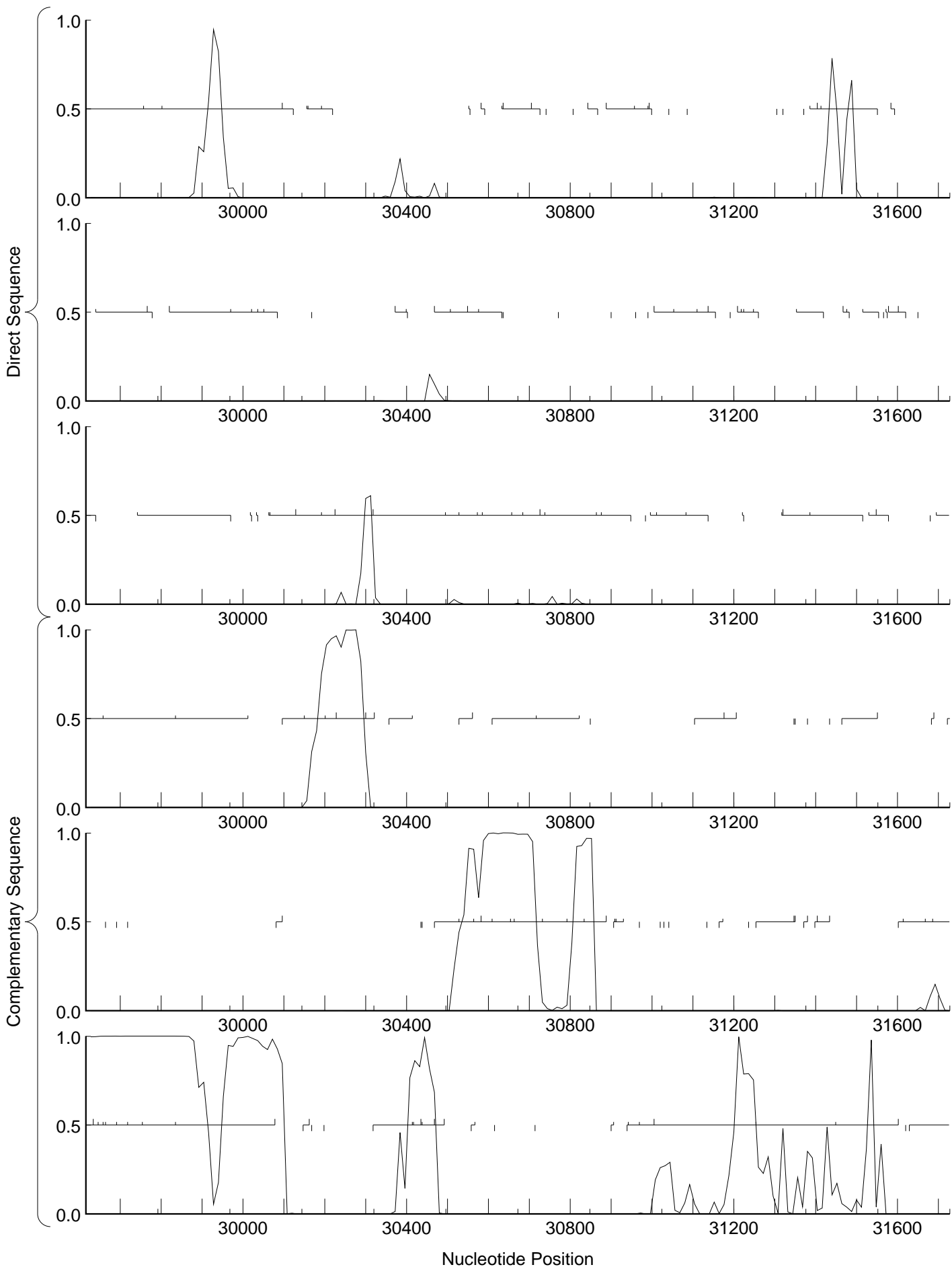




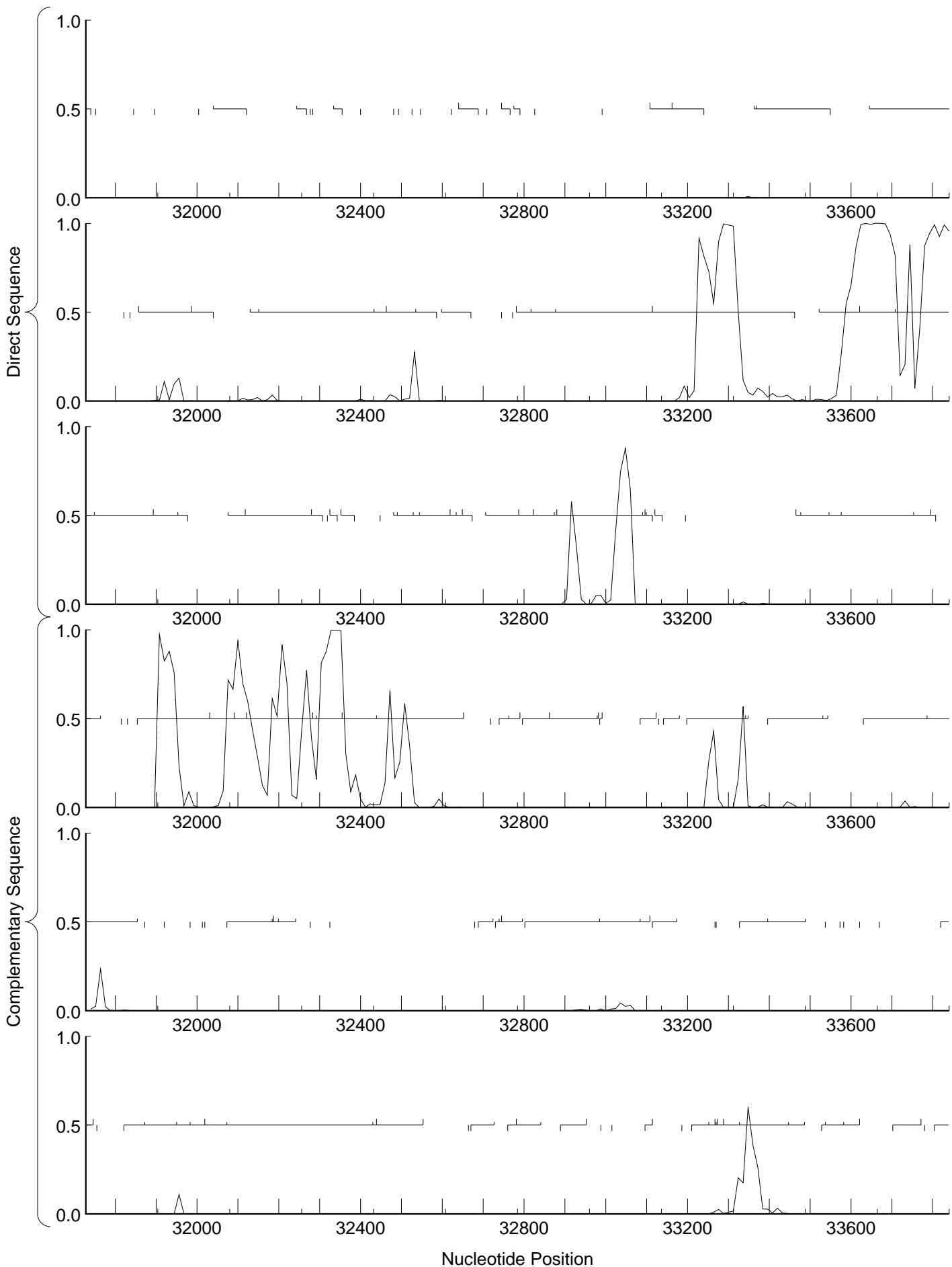


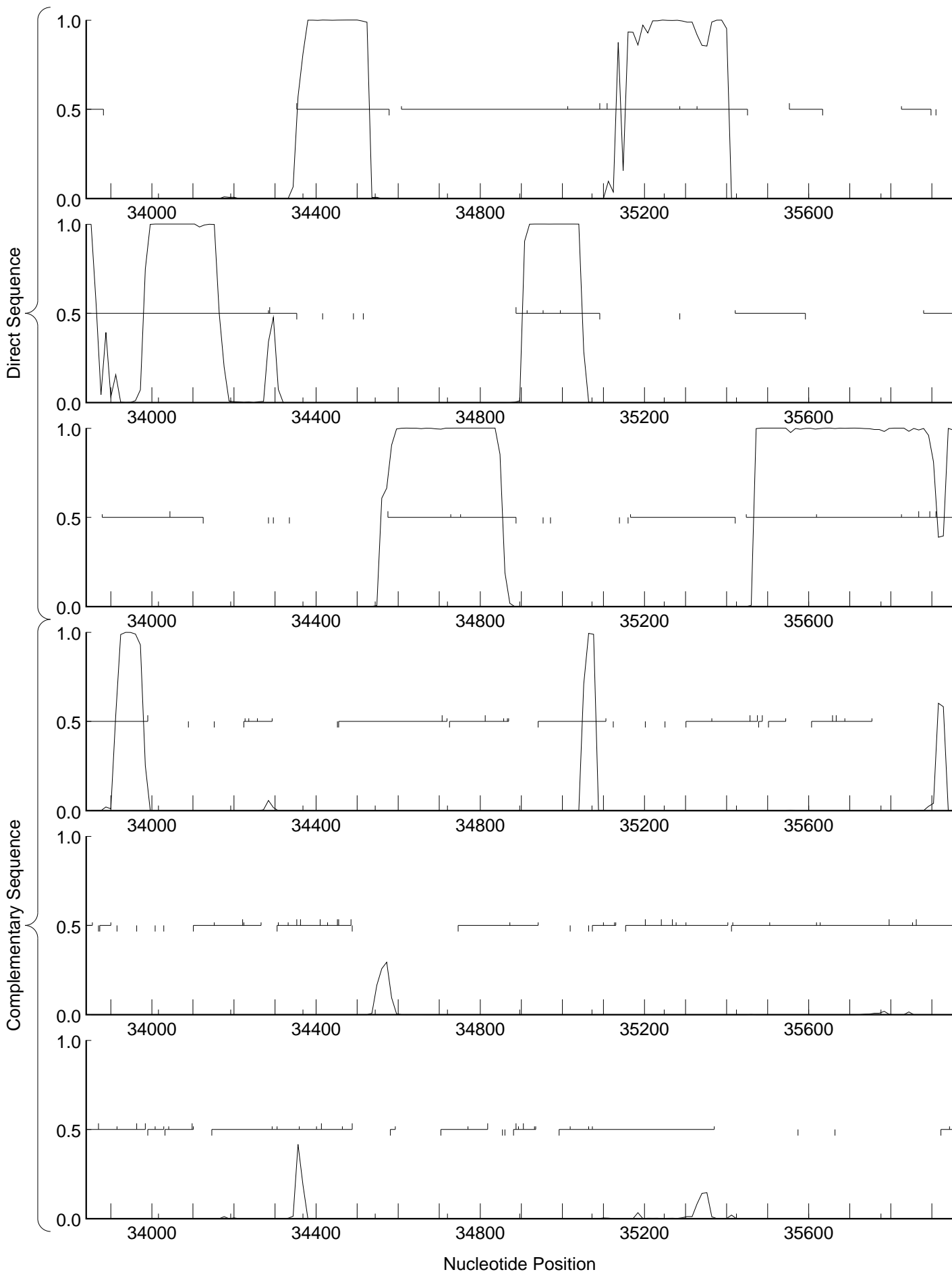


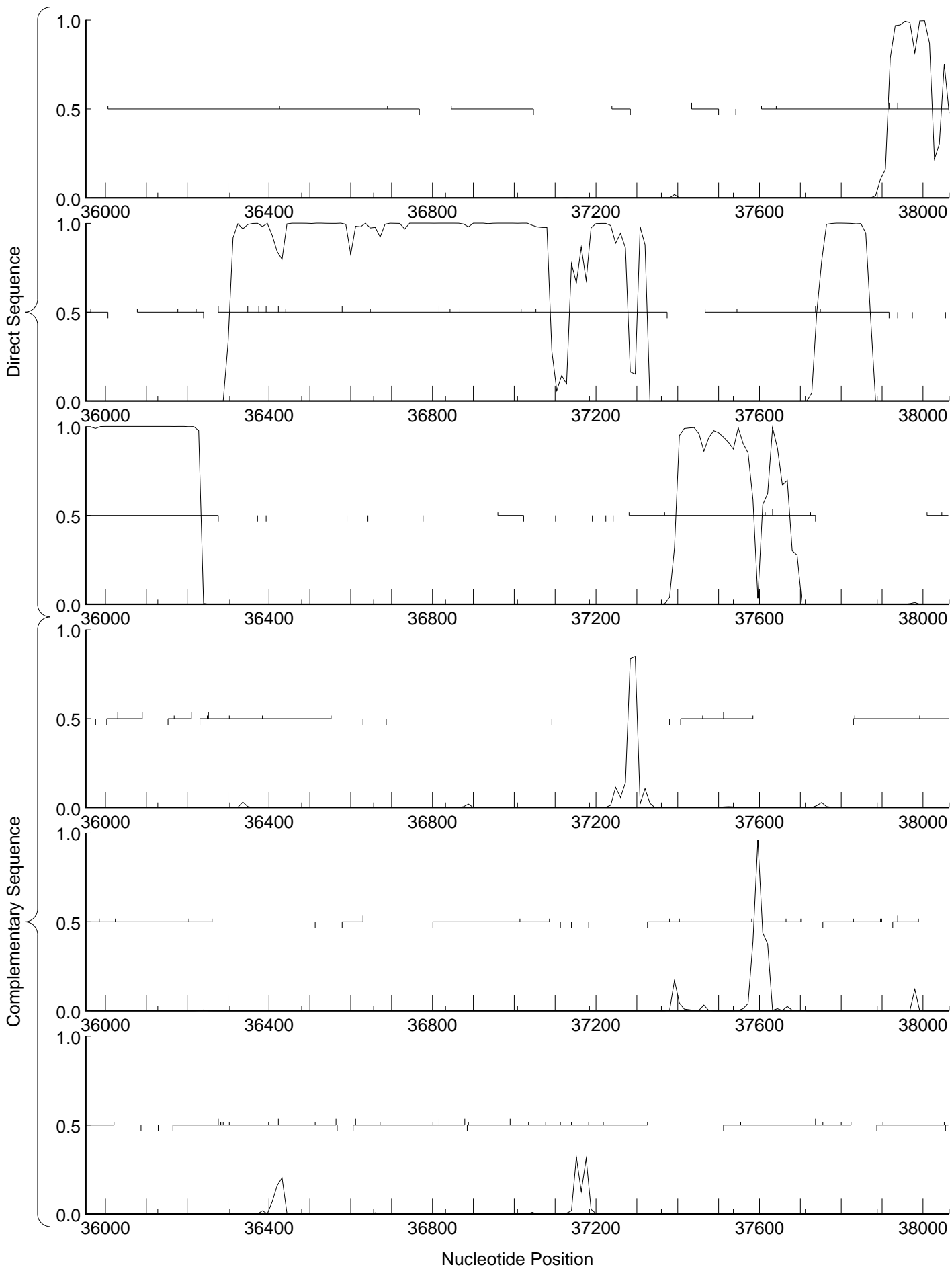


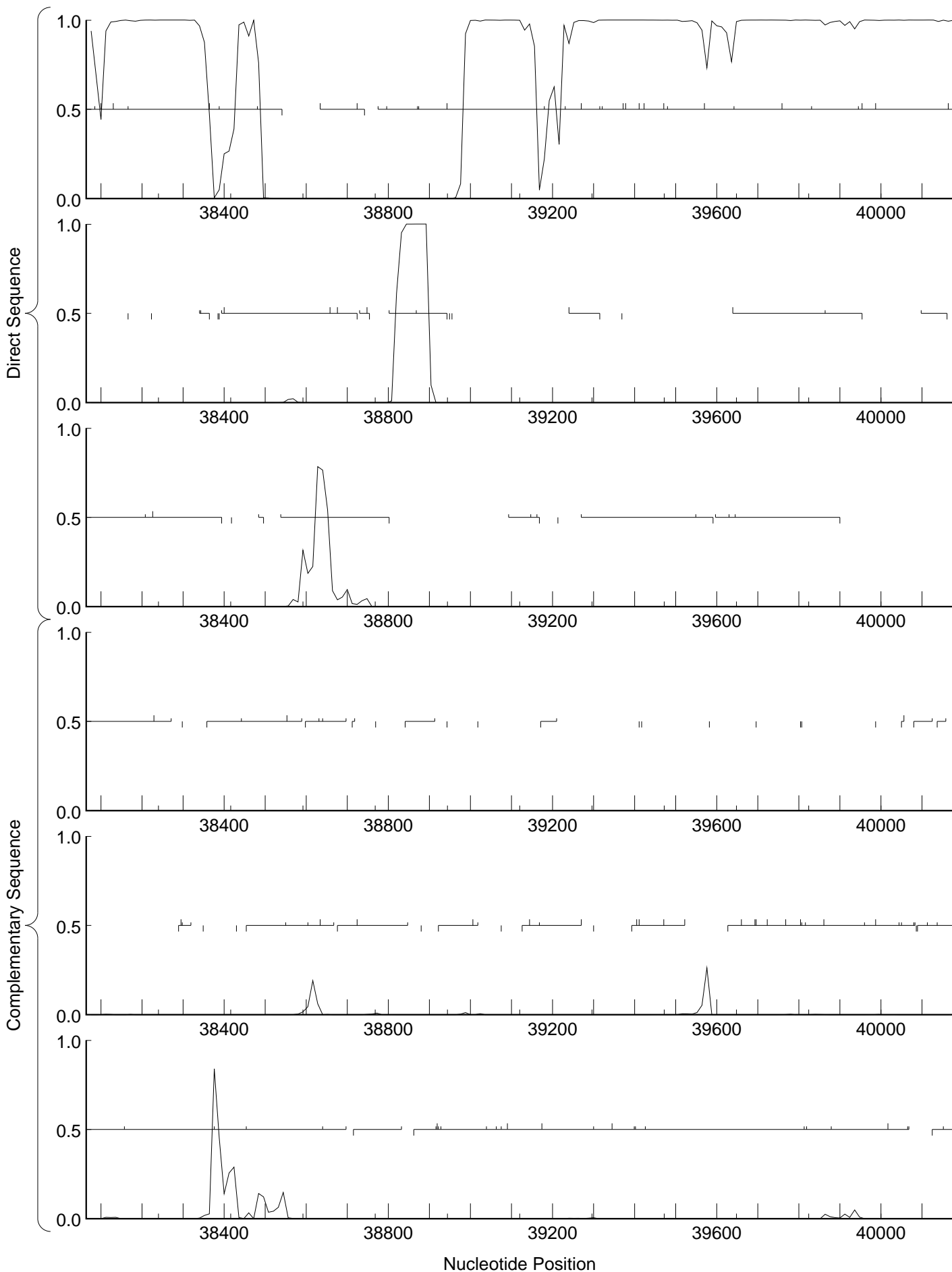


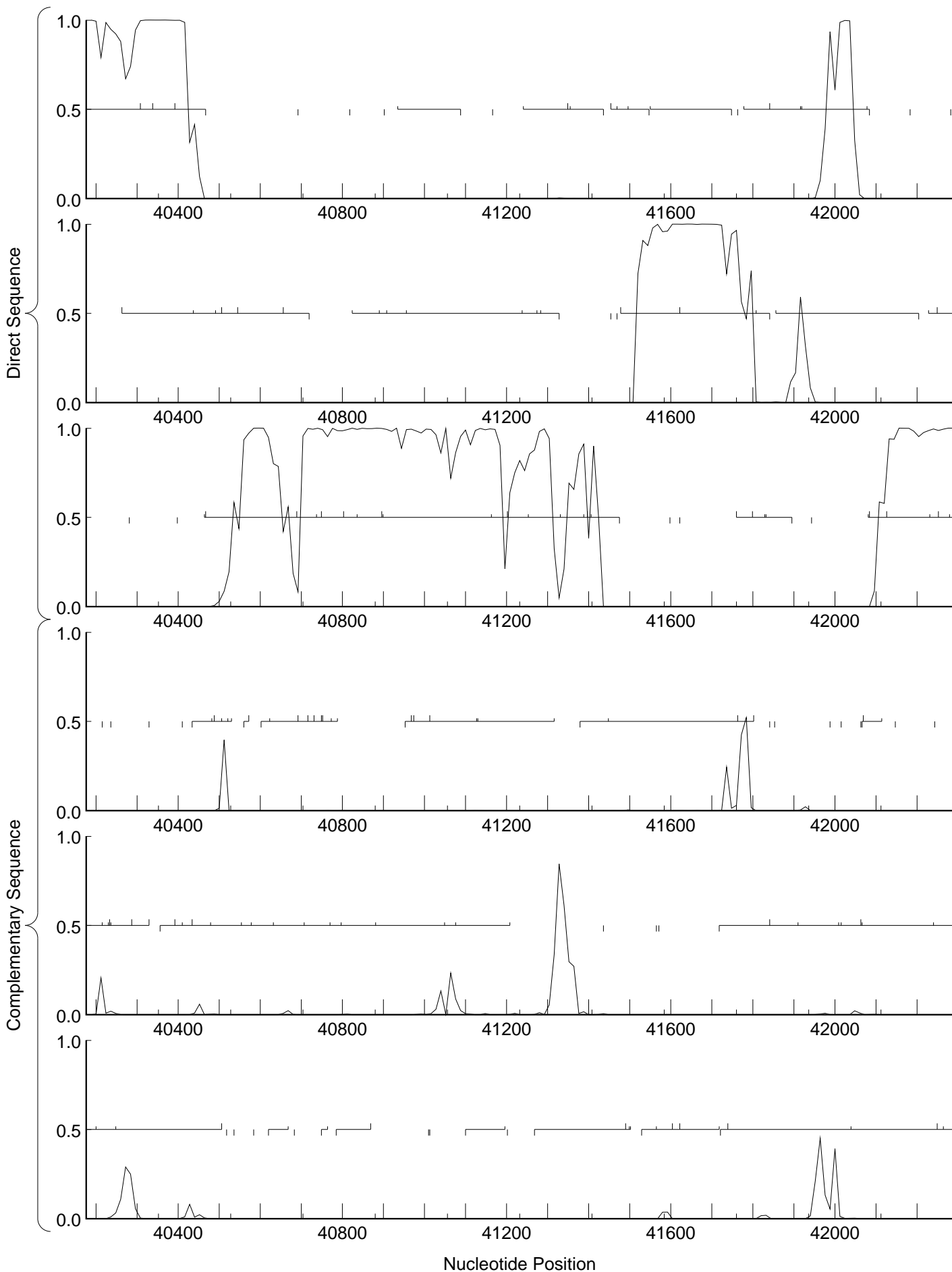


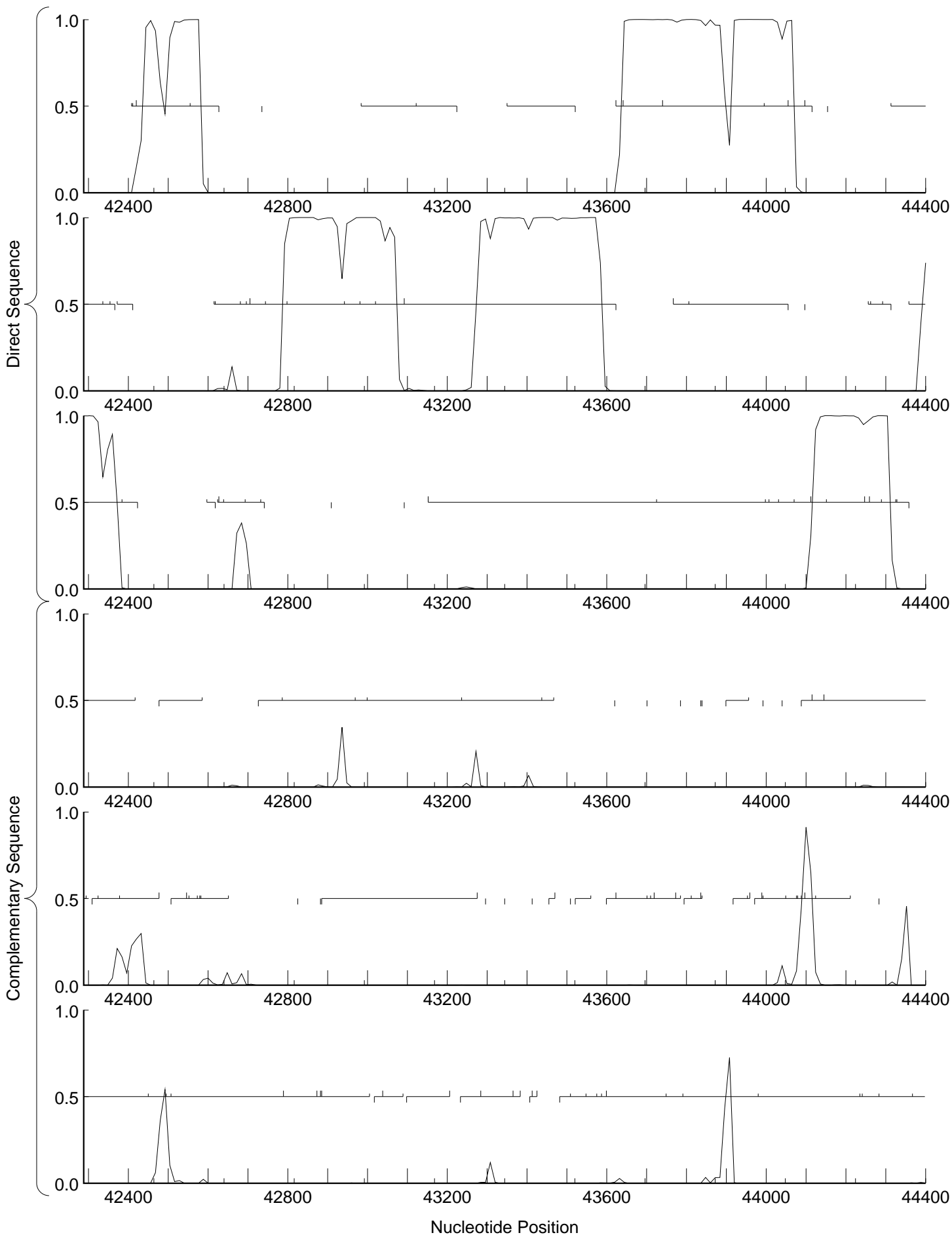












Complete sequence, 46692 bp including 11-base 3' overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 23/24

