

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Microbacterium phage Barnstormer complete sequence, 46860 bp including 11-base 3' overhang  
Analysis Date : 8/19/22 at 13:05:36  
Pages : 24  
Sequence Length : 46860 bp  
GC Content : 69.53%

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,

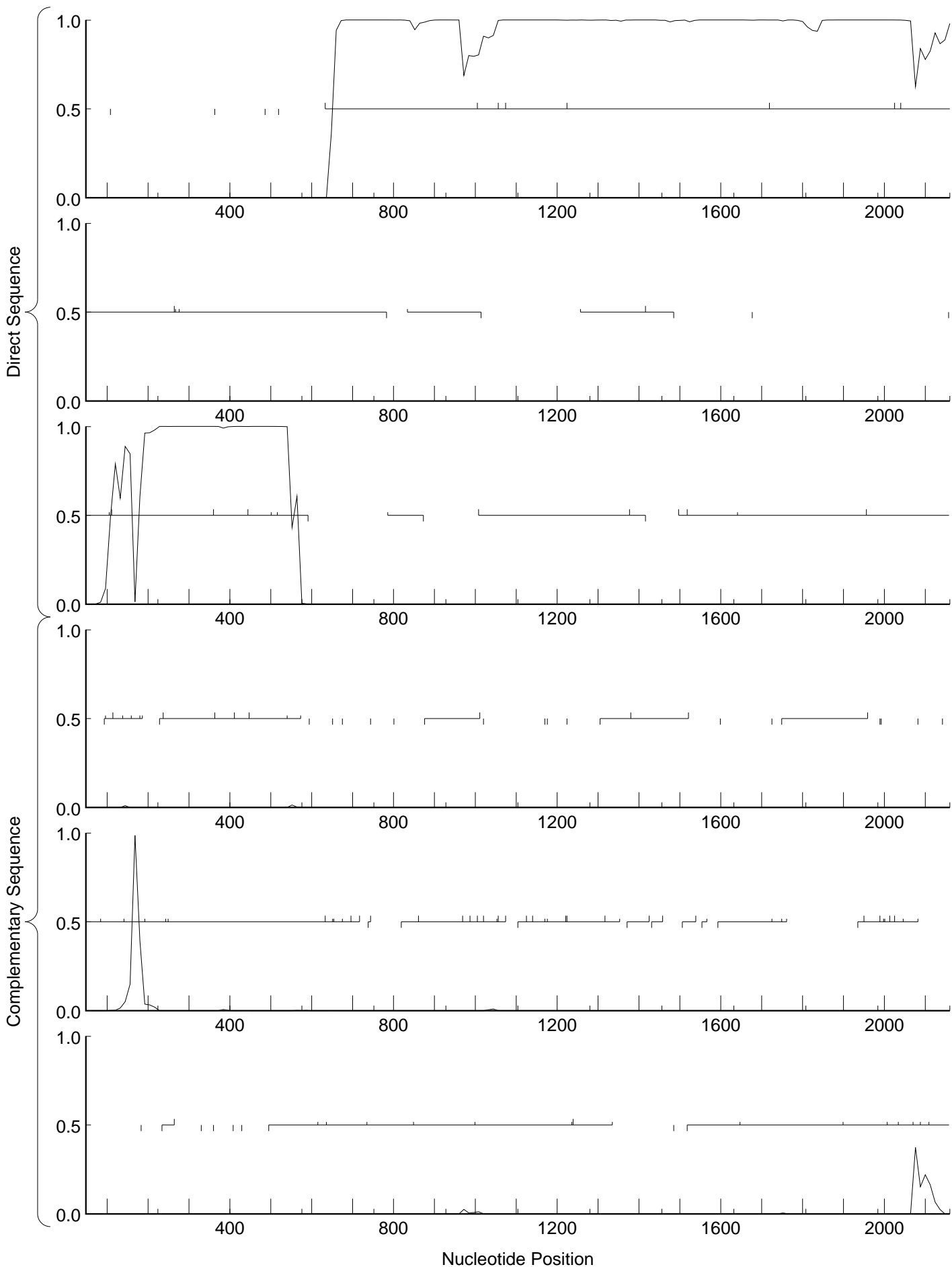
## MATRIX INFORMATION

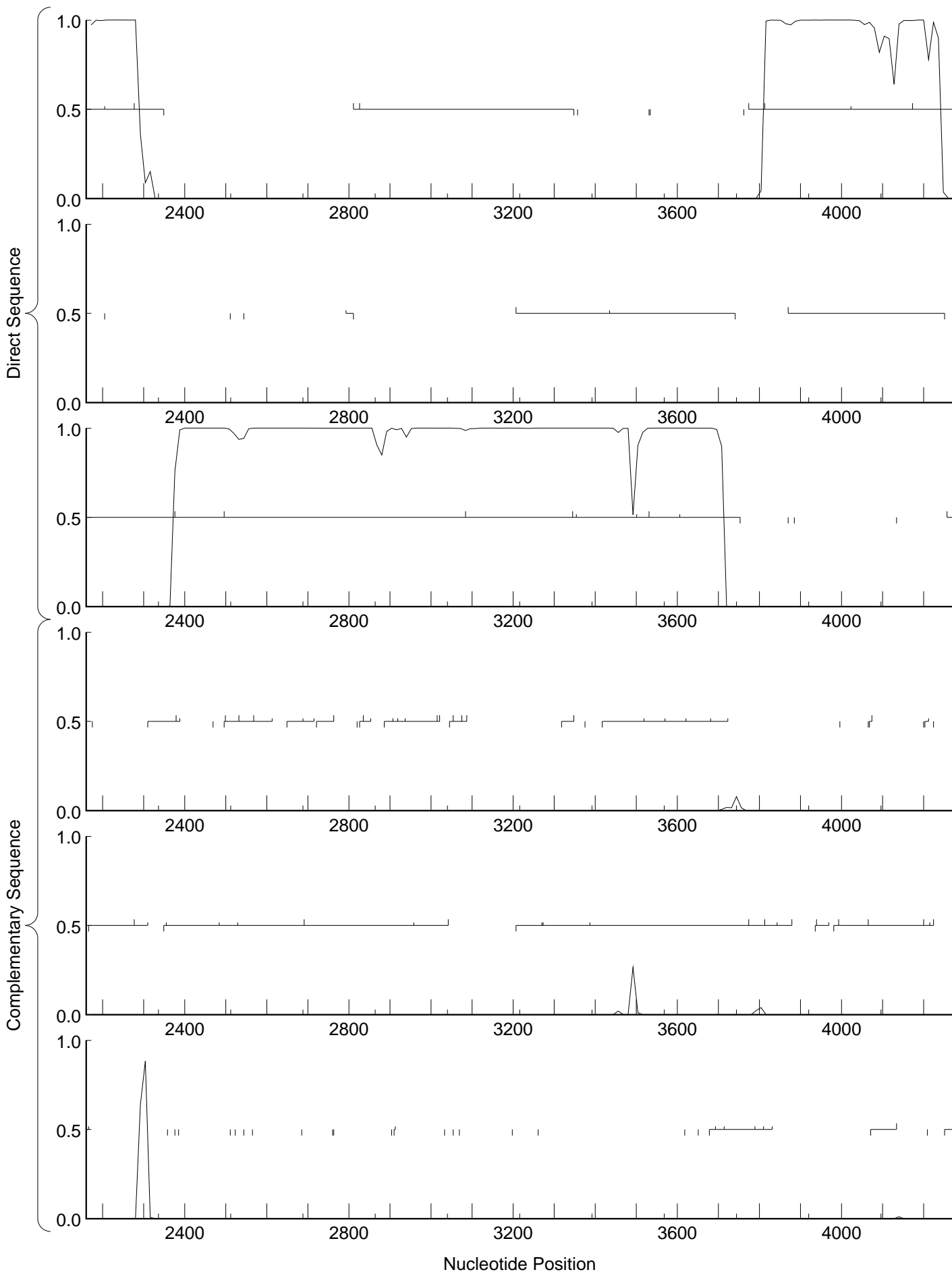
Matrix : -  
Author : -  
Order : 4

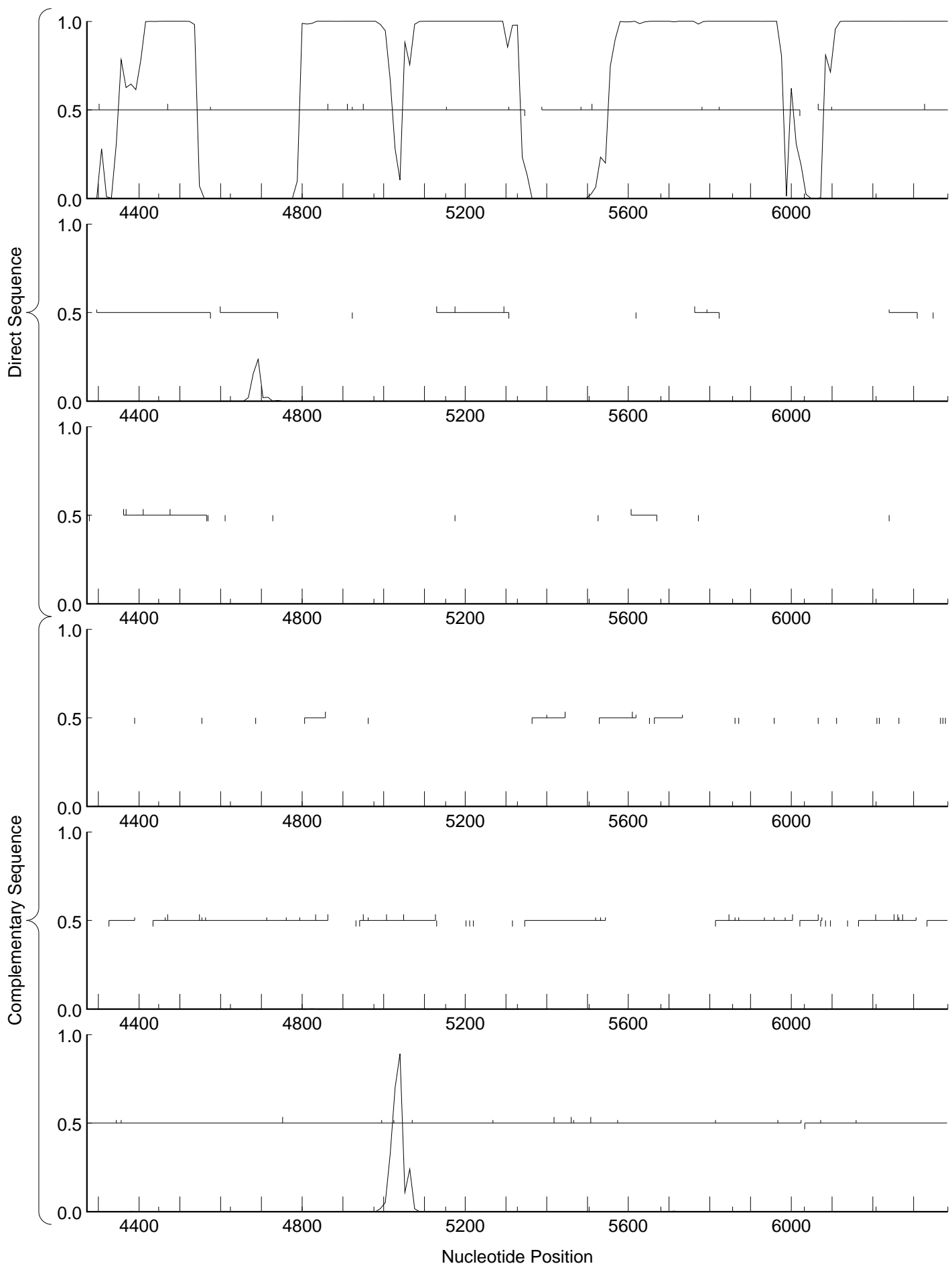
Send questions / comments to:  
Dr. M. Borodovsky  
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Atlanta, GA 30332-0230

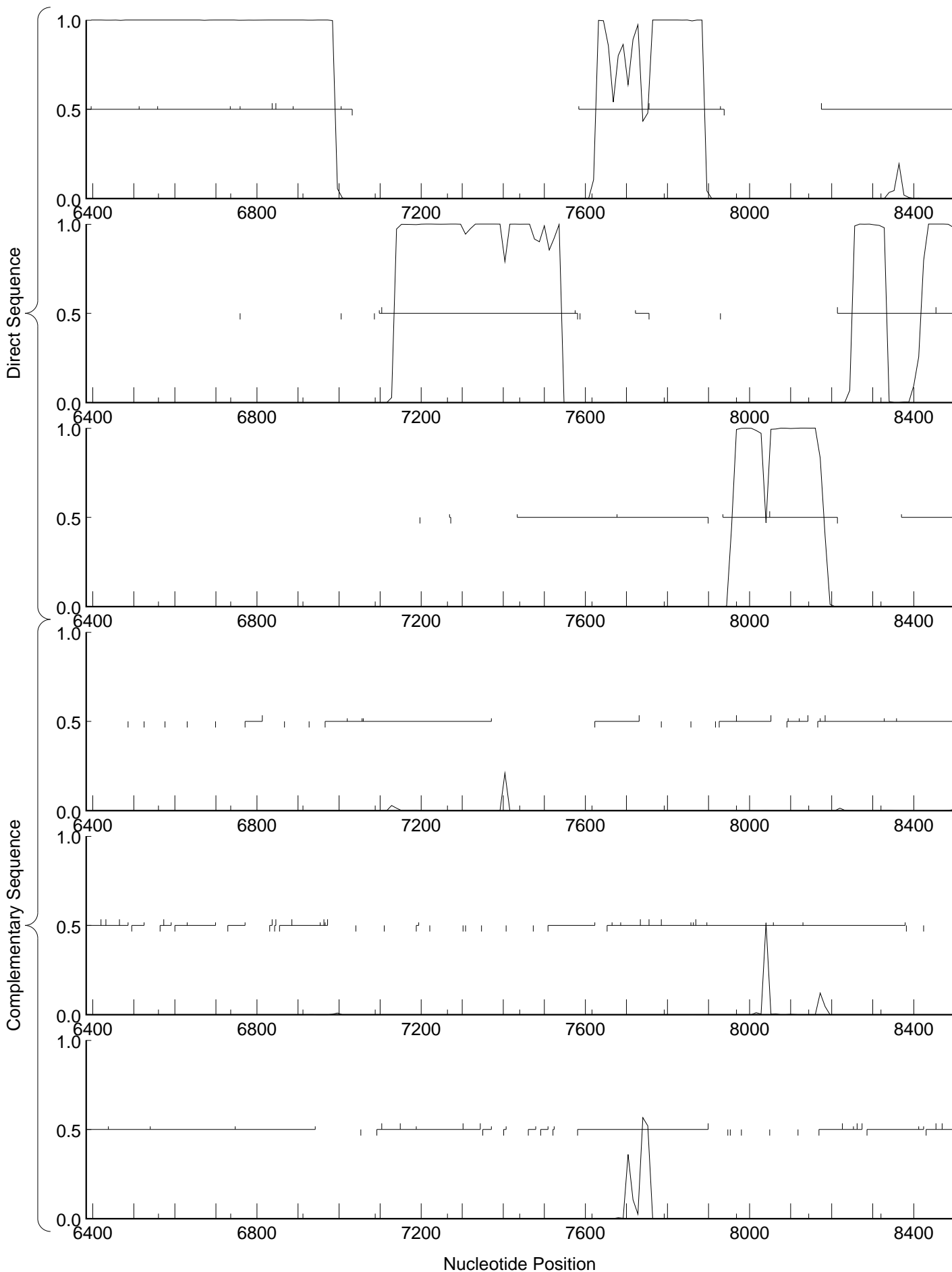
### Matrix notes & comments

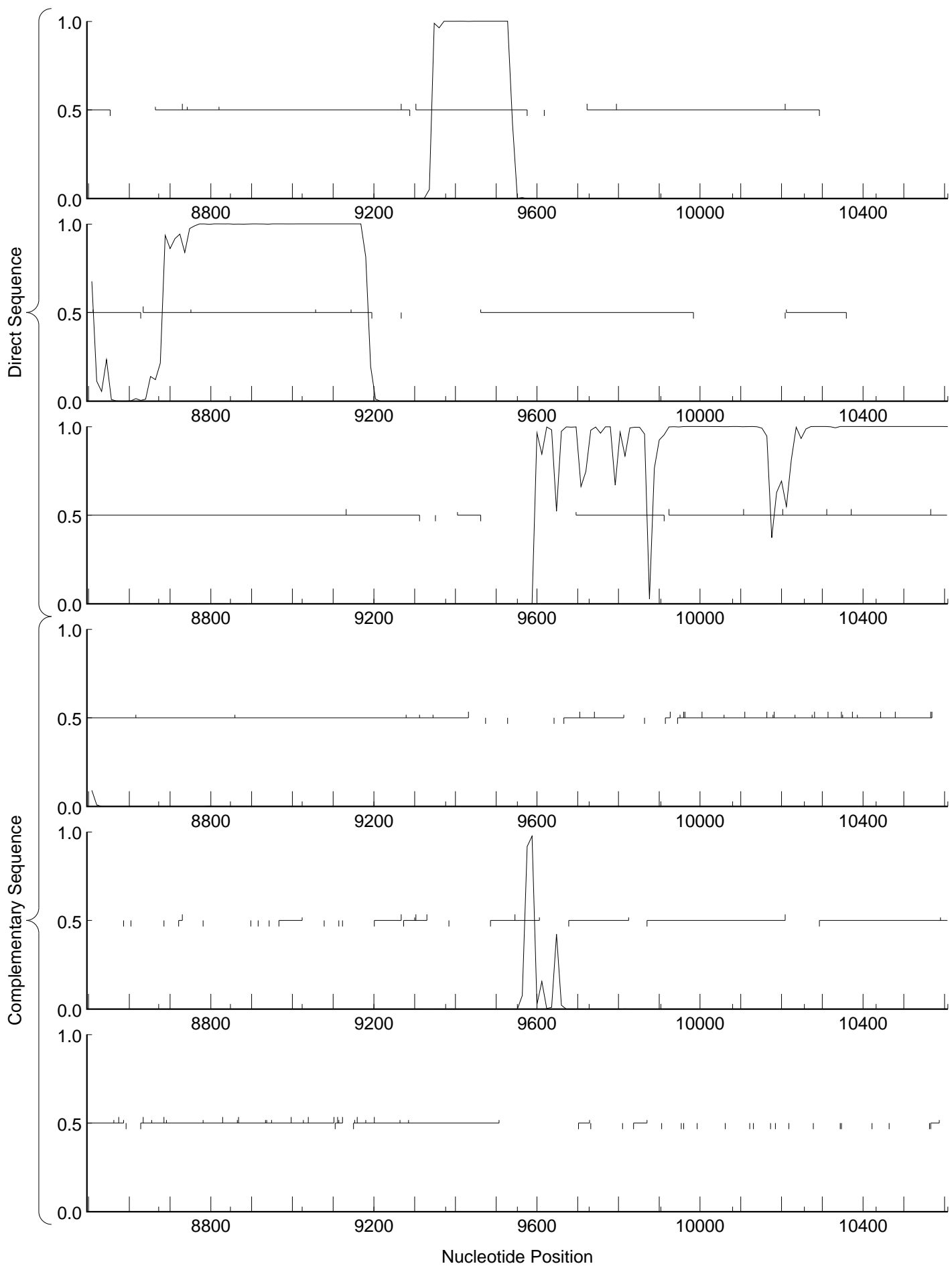
Training set derived by GeneMarkS, 4.30 October 2014  
Fri Sep 7 18:42:59 2018

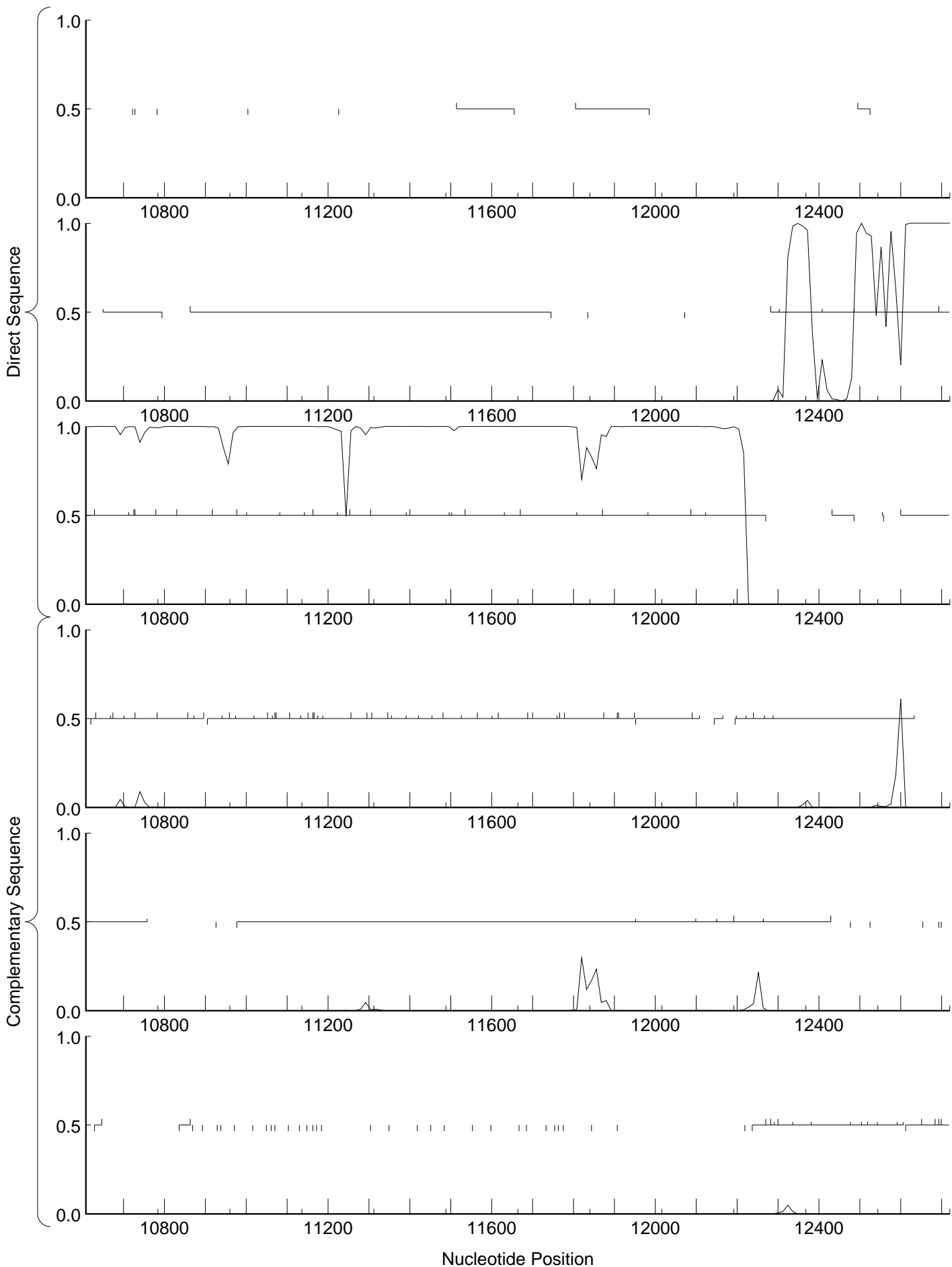




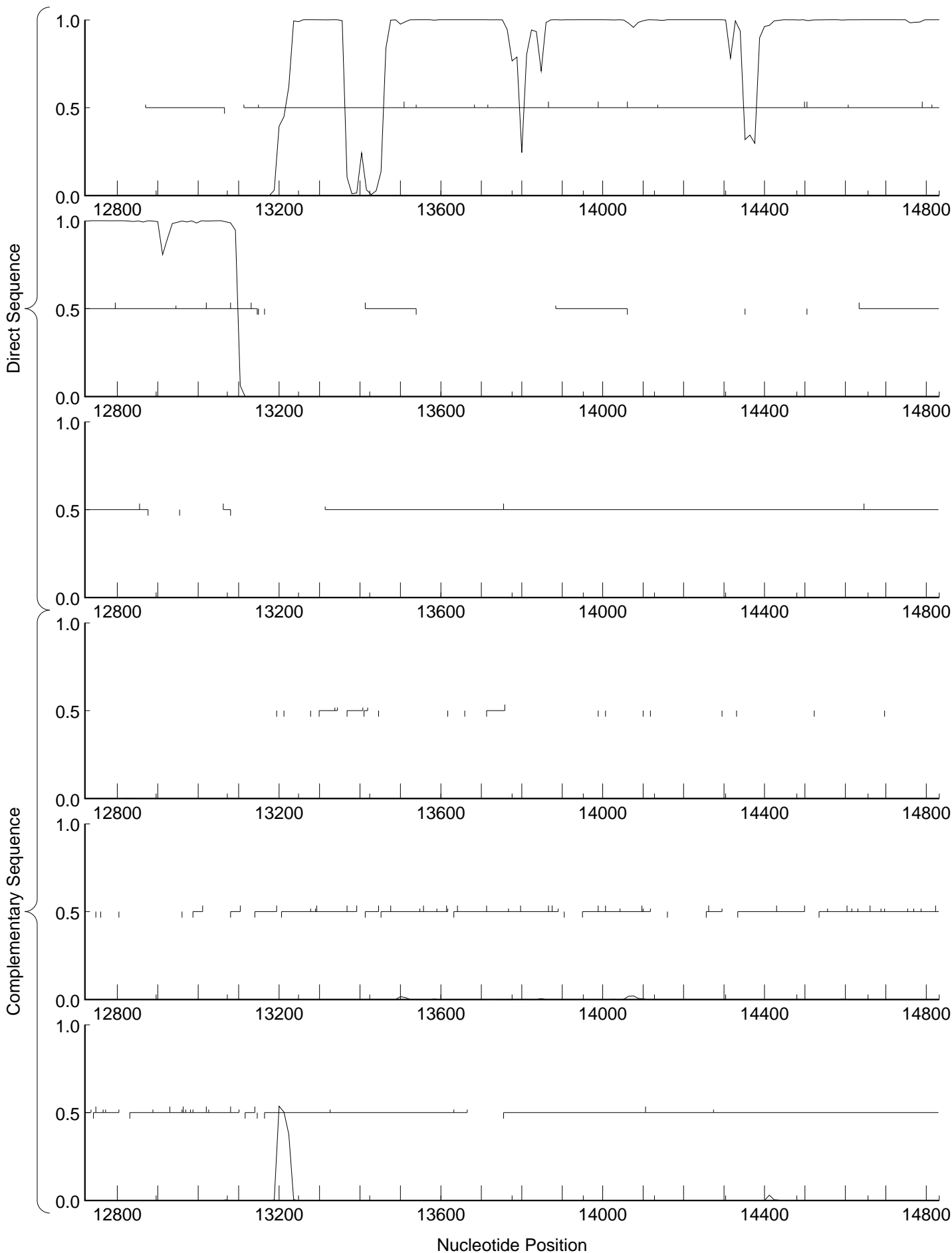




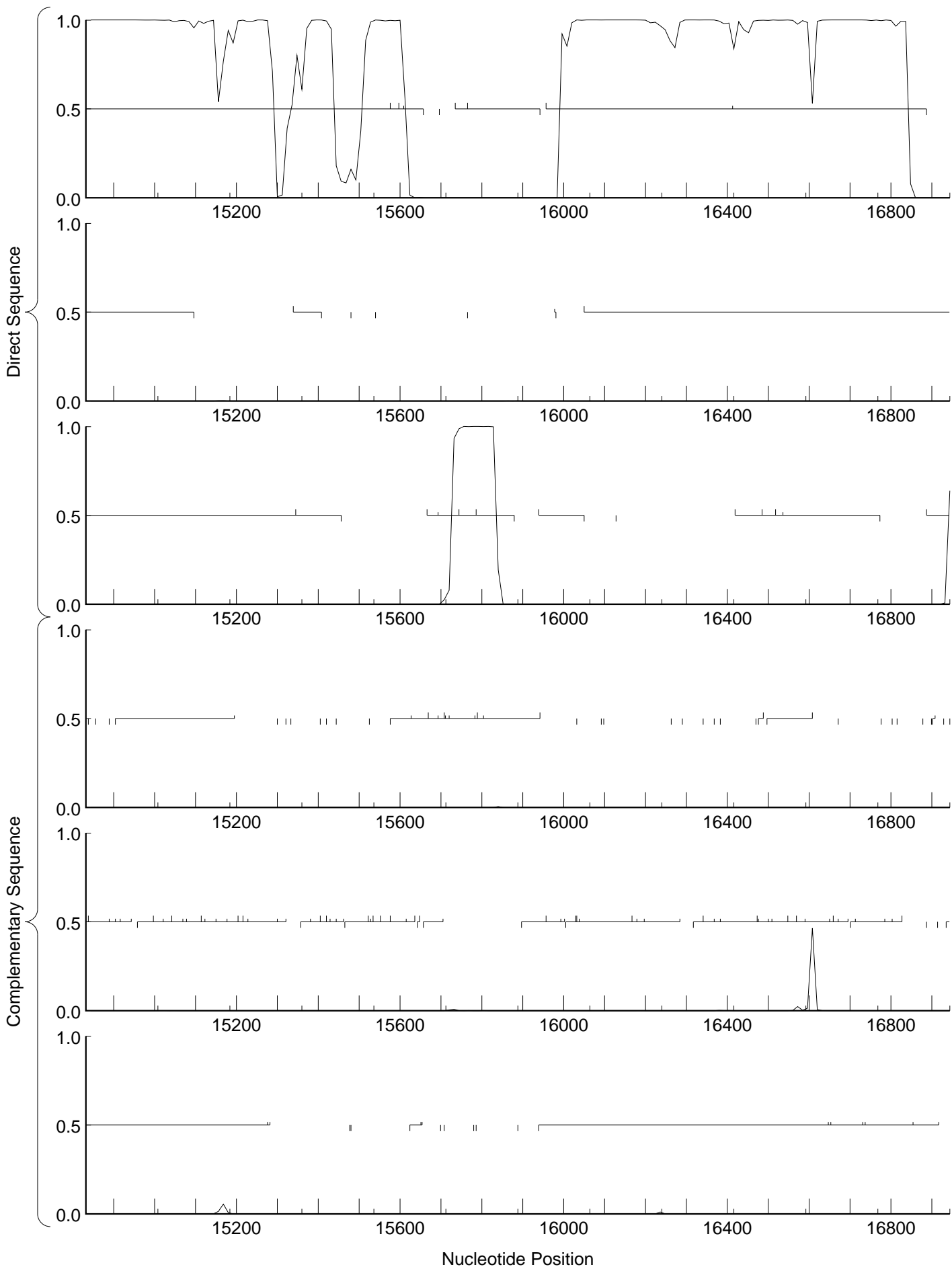


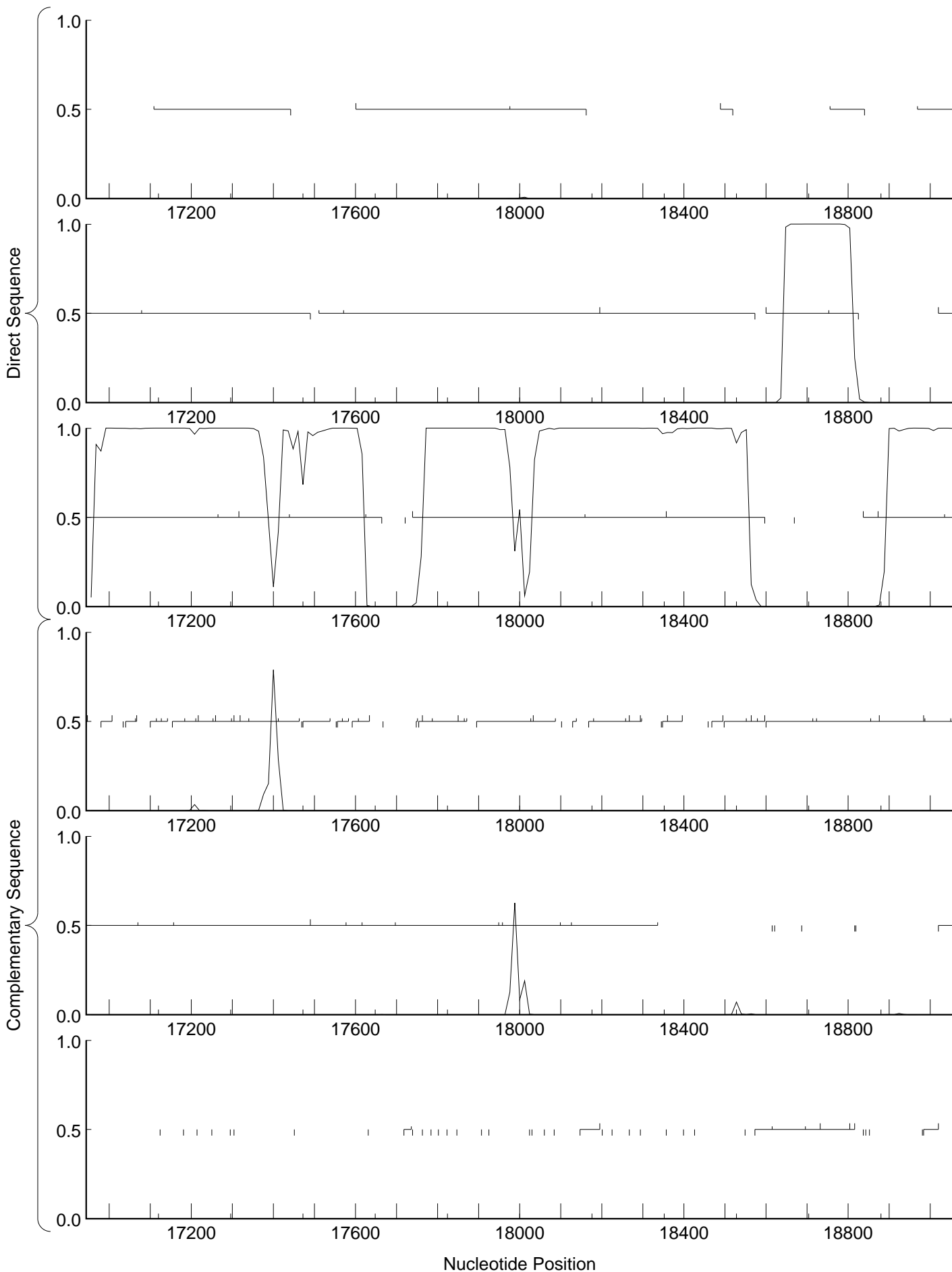


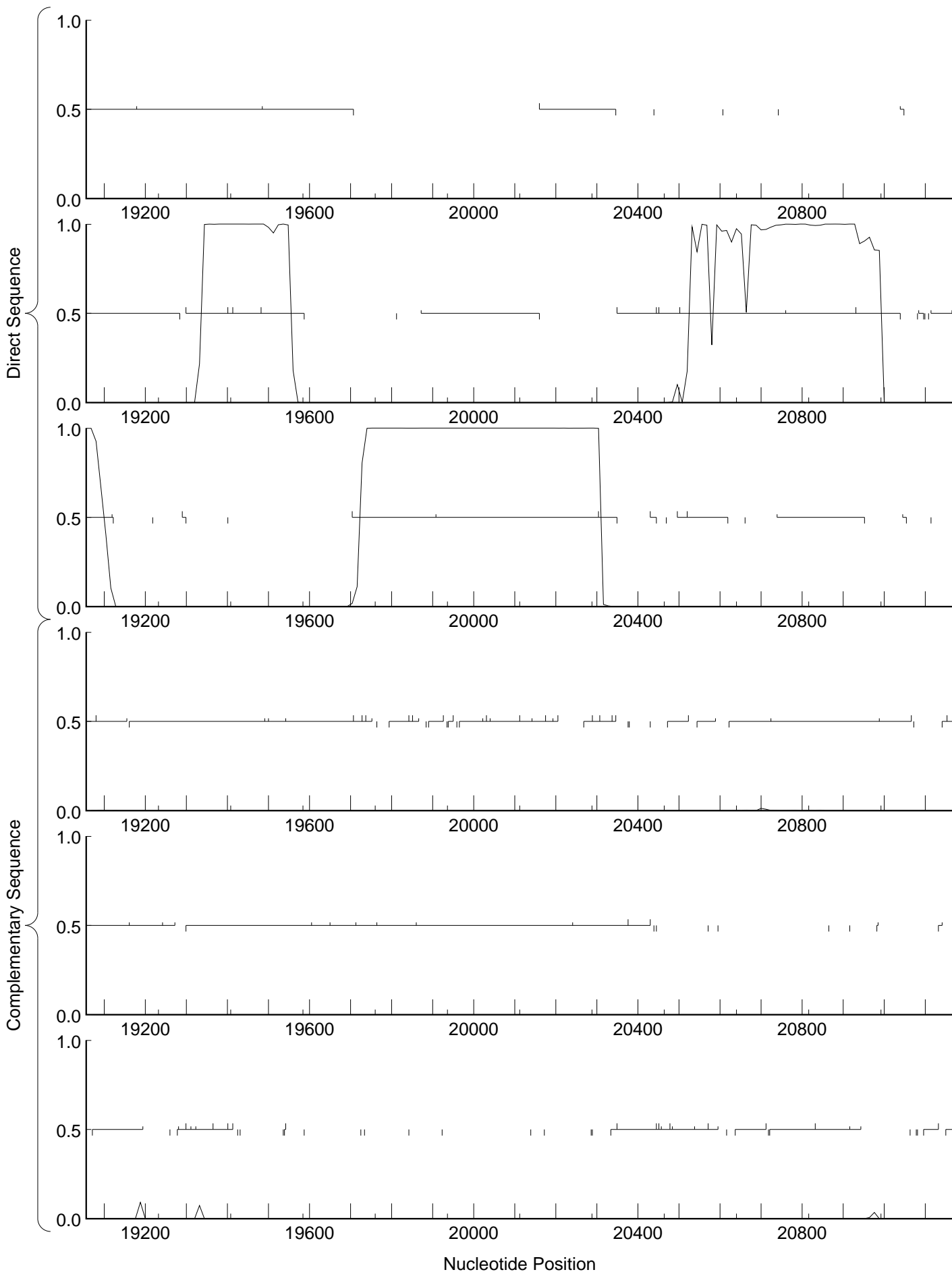
complete sequence, 46860 bp including 11-base 3' overhang (CGAGGAGGCAT), Cluster EH, Order 4, Window 96, Step 12, 8/24

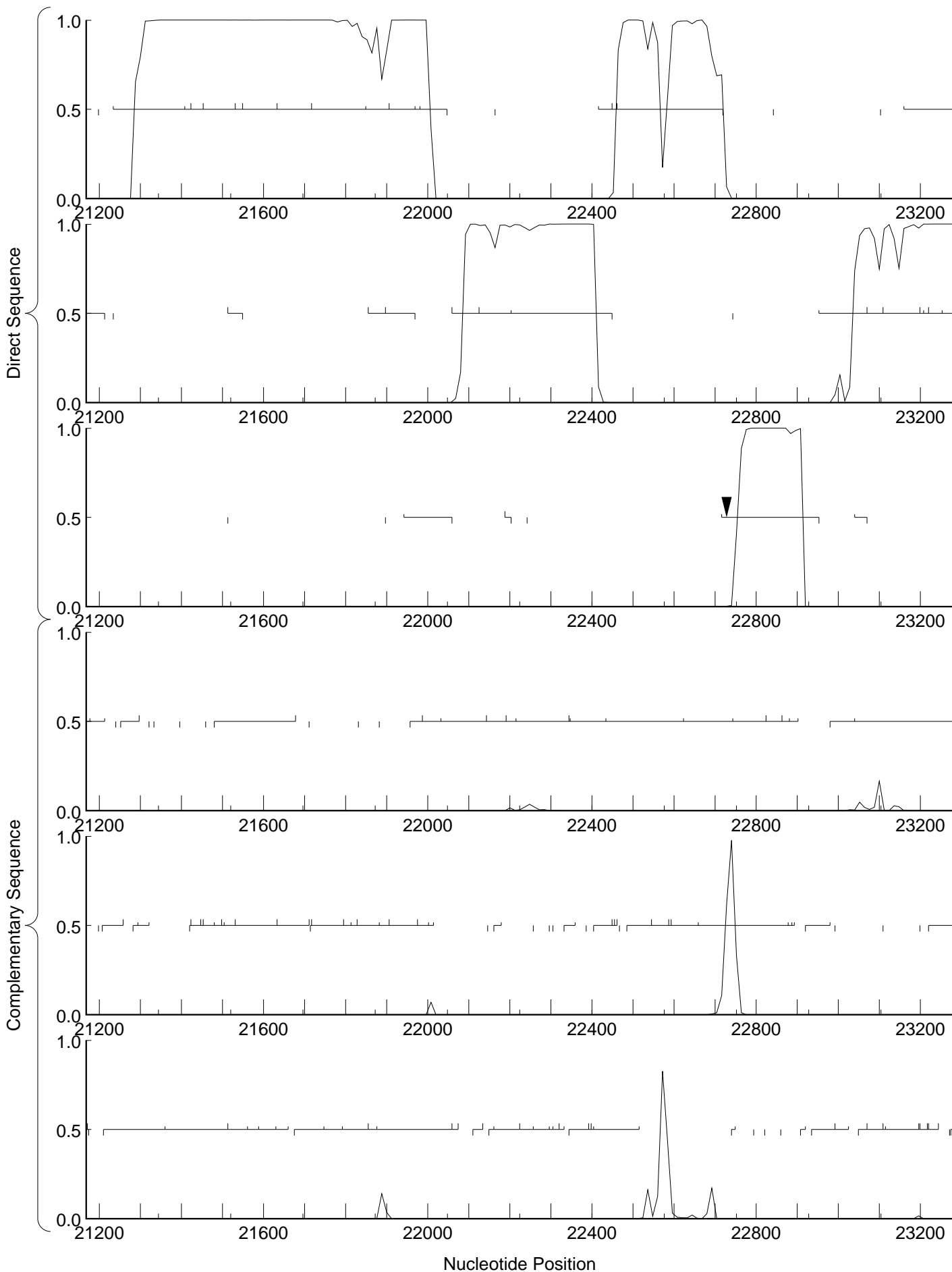


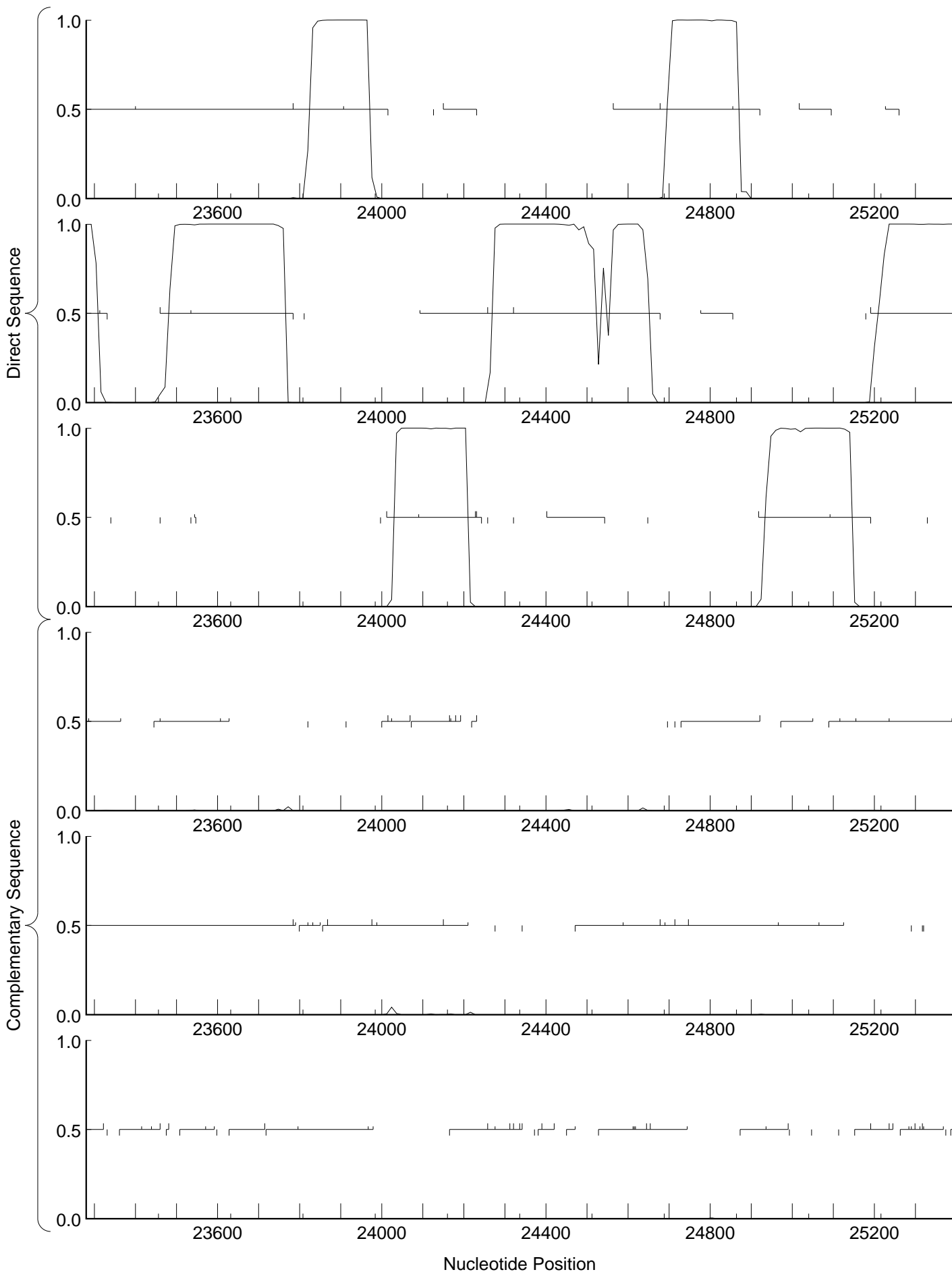


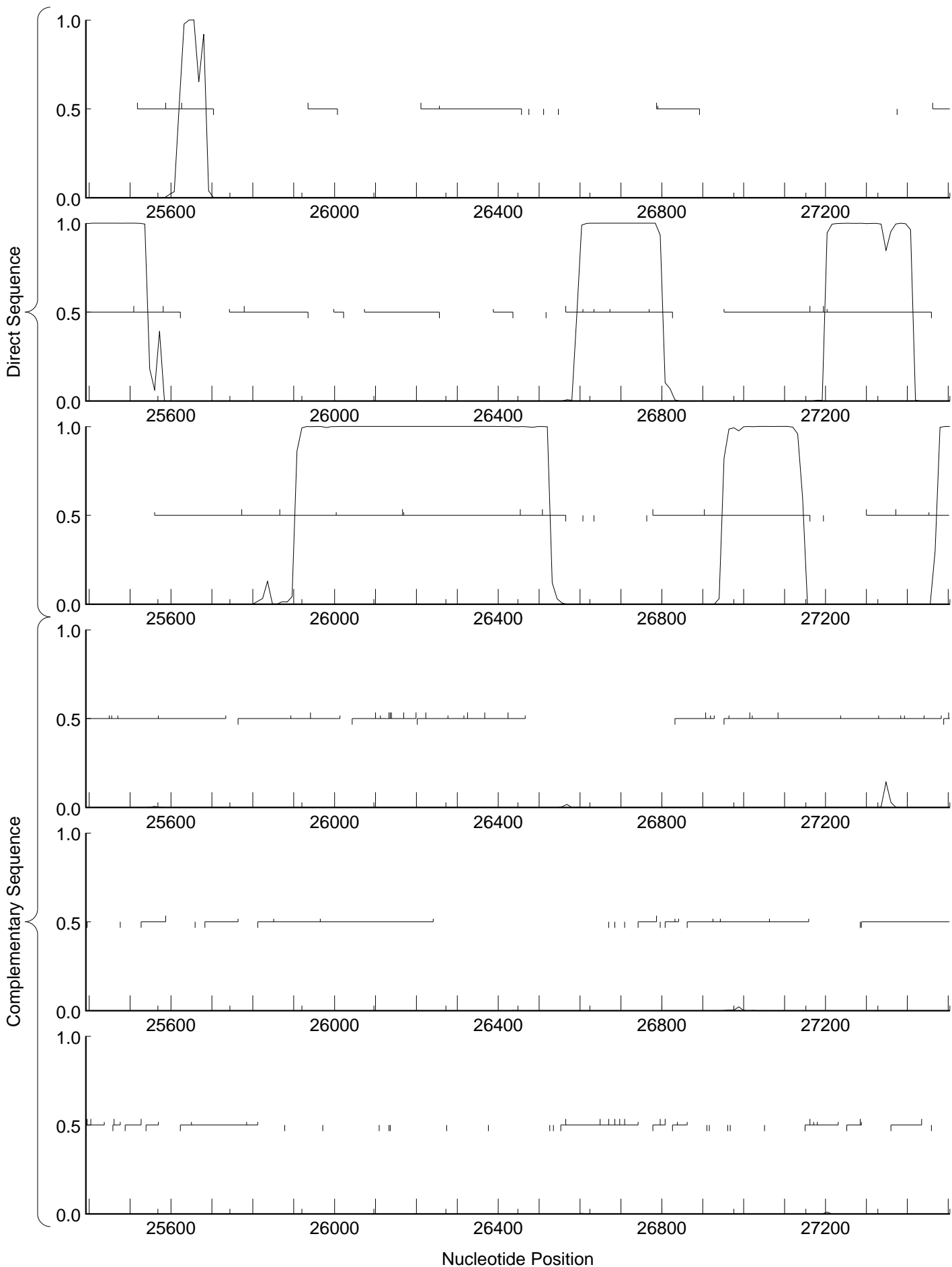












complete sequence, 46860 bp including 11-base 3' overhang (CGAGGAGGCAT), Cluster EH, Order 4, Window 96, Step 12, 15/24

