

# GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

## PROGRAM INFORMATION

Sequence : Gordonia phage Clark complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA  
Analysis Date : 3/17/18 at 16:23:01  
Pages : 24  
Sequence Length : 47716 bp  
GC Content : 66.53%

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 notes & comments

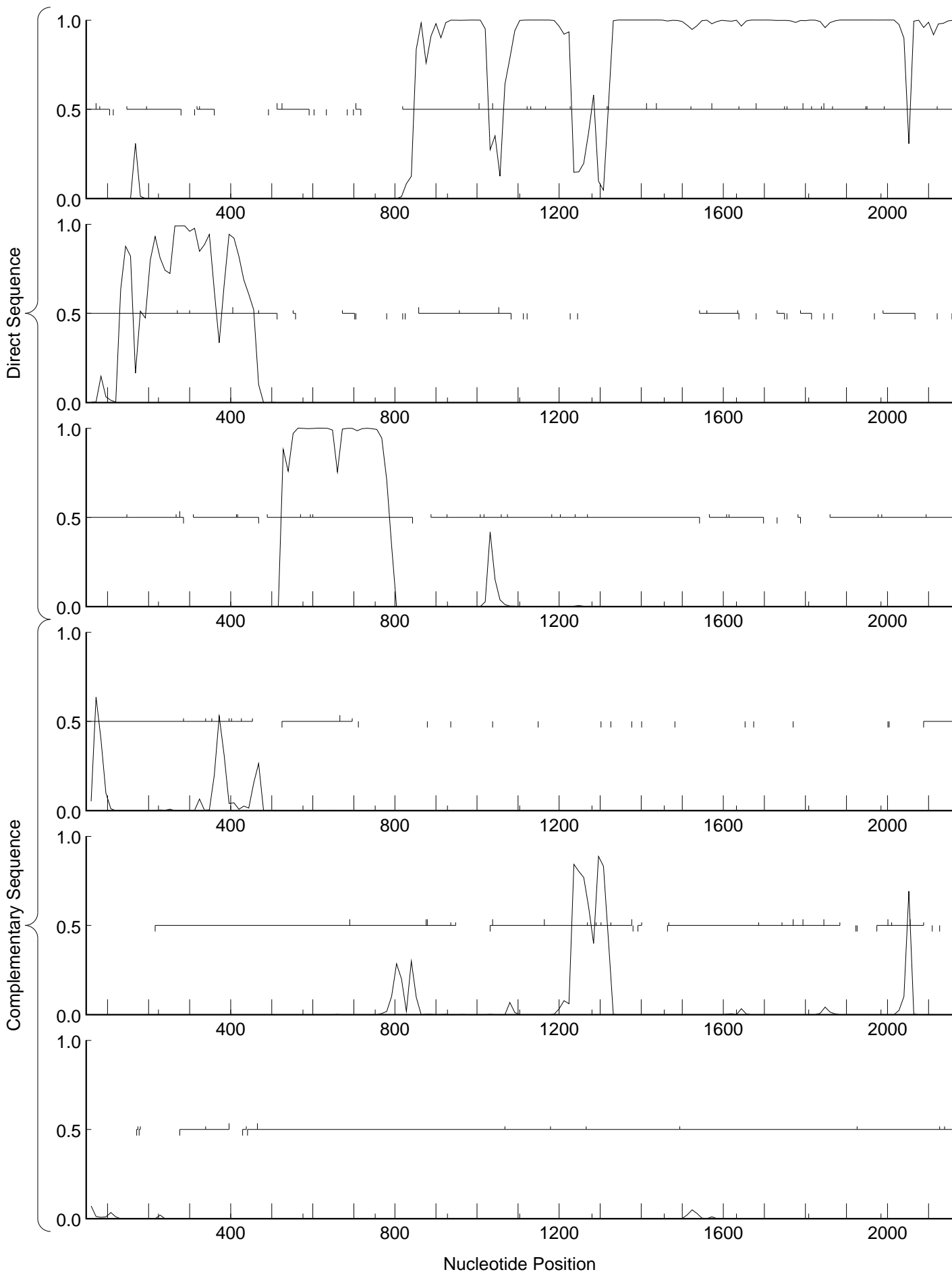
Training set derived by GeneMarkS, 4.27 September 2014  
Tue Sep 23 15:14:37 2014

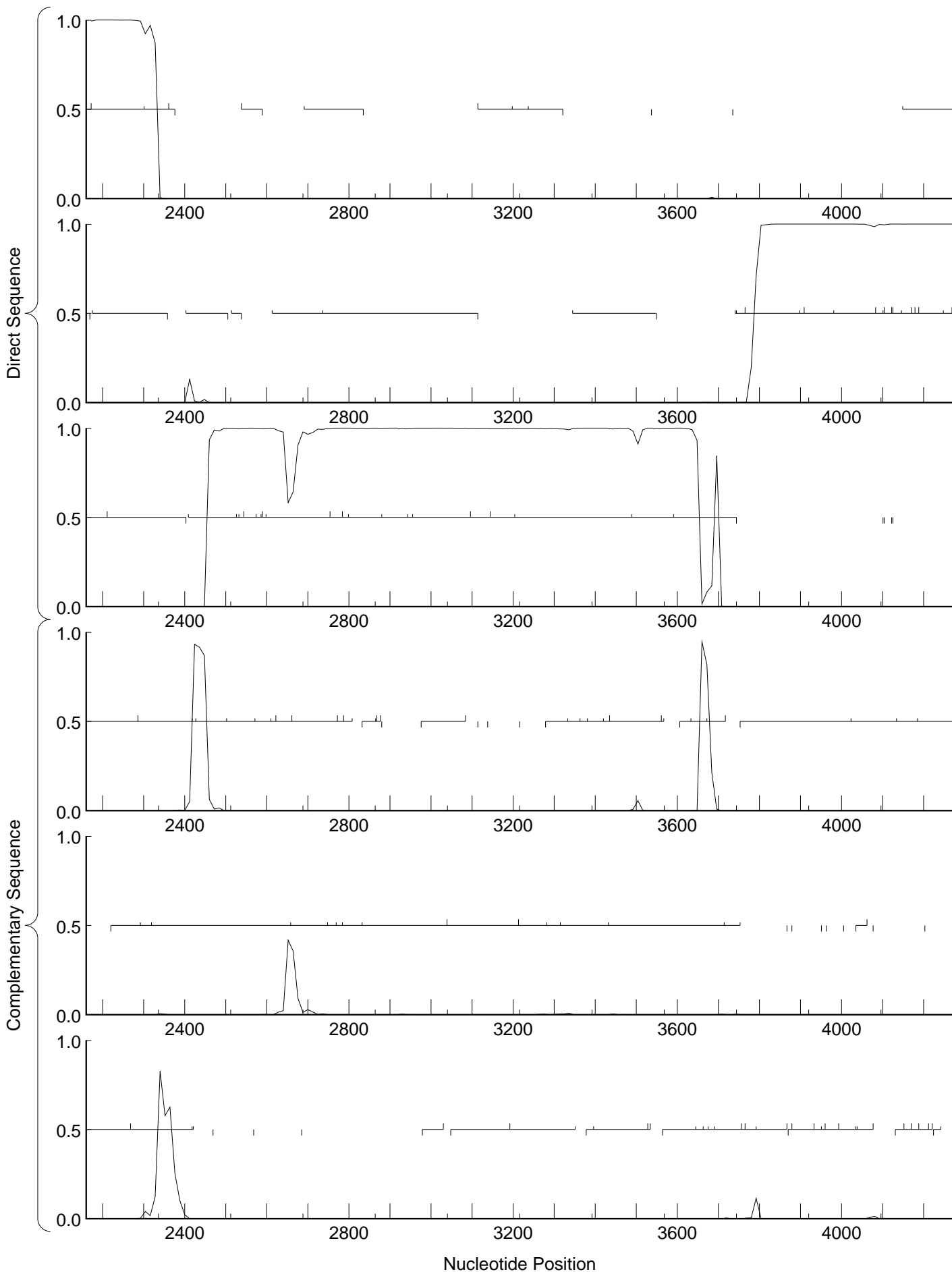
## MATRIX INFORMATION

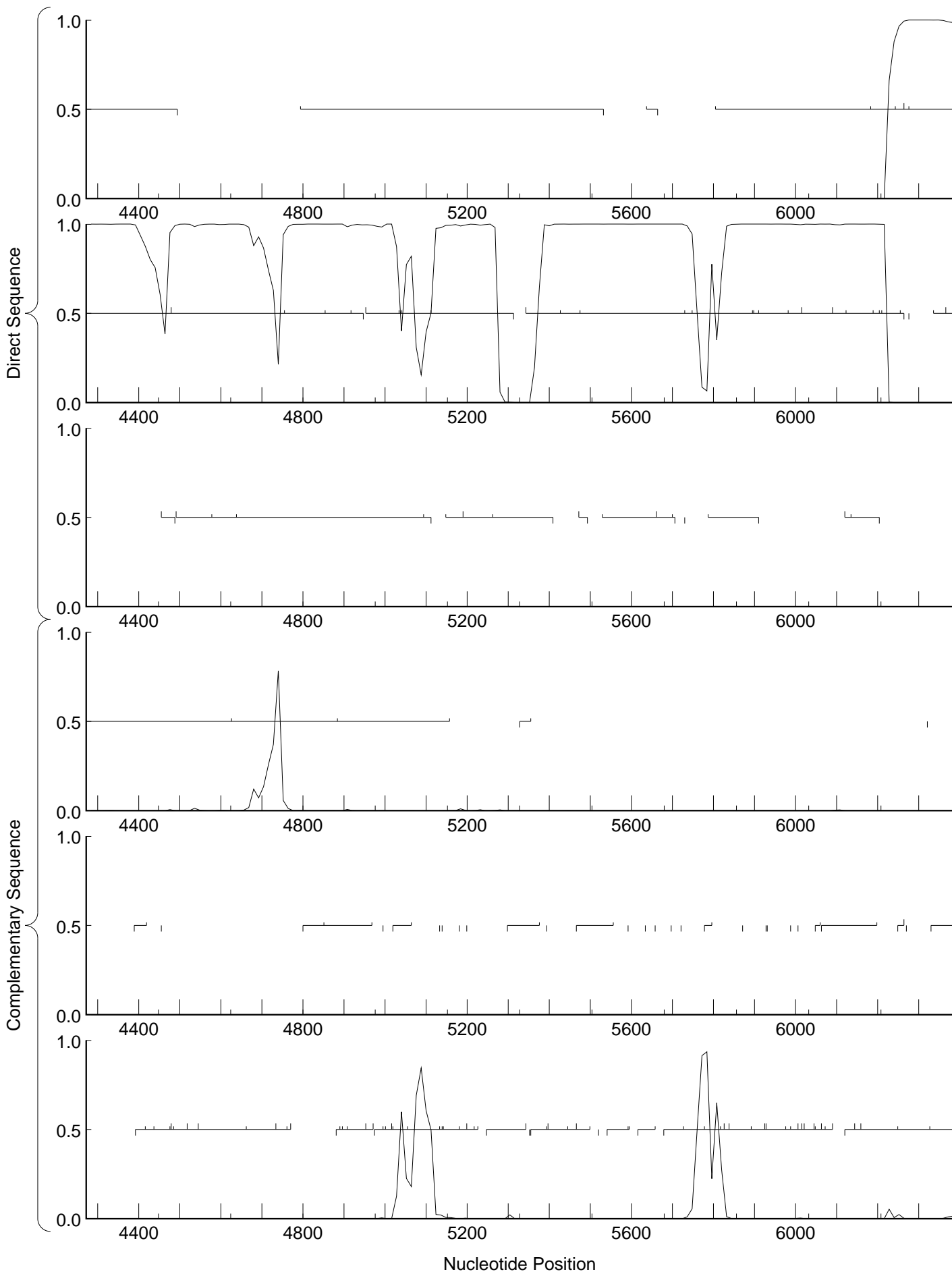
Matrix : Mycobacterium\_tuberculosis\_H37Rv  
Author : -  
Order : 4

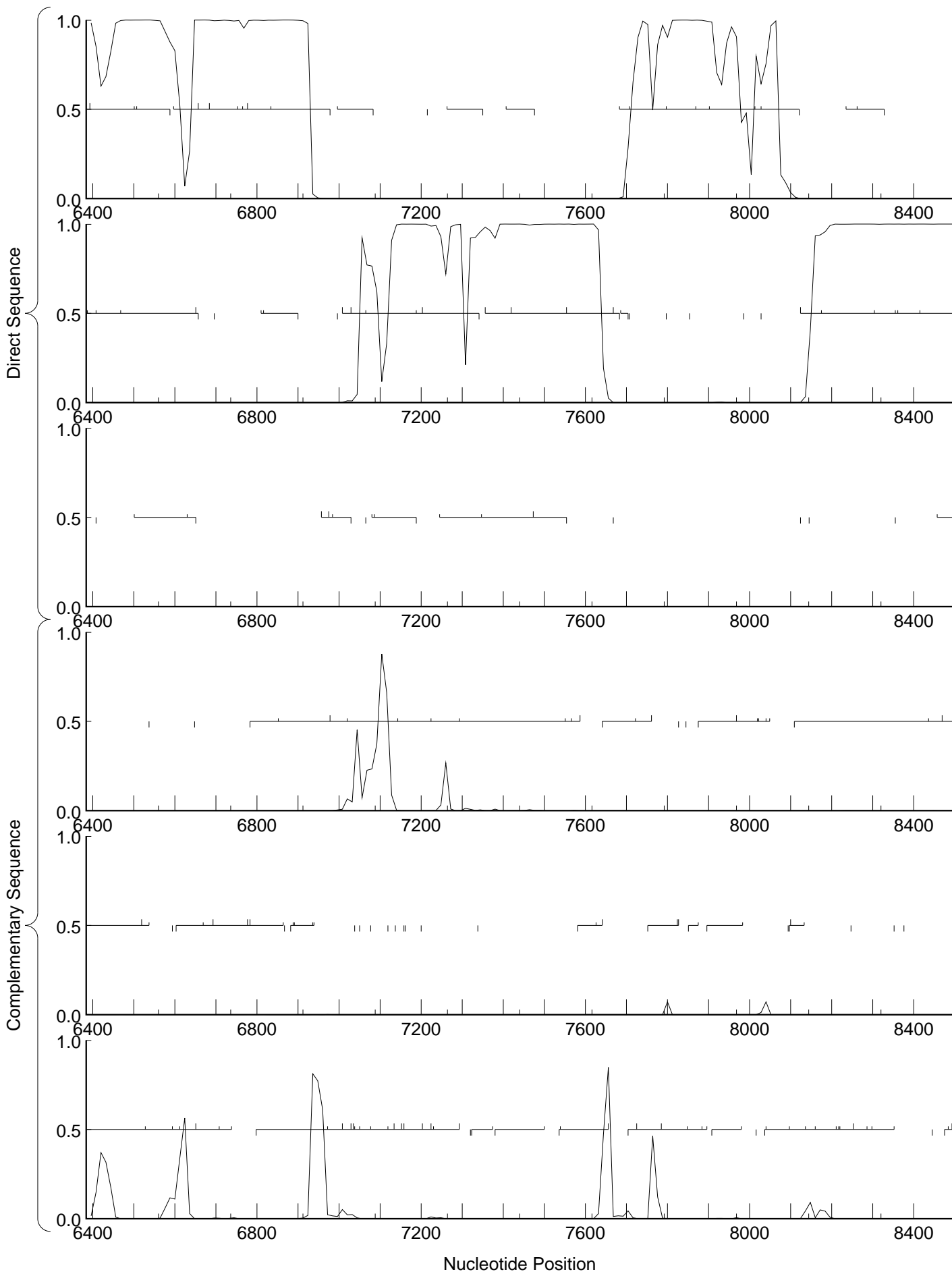
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

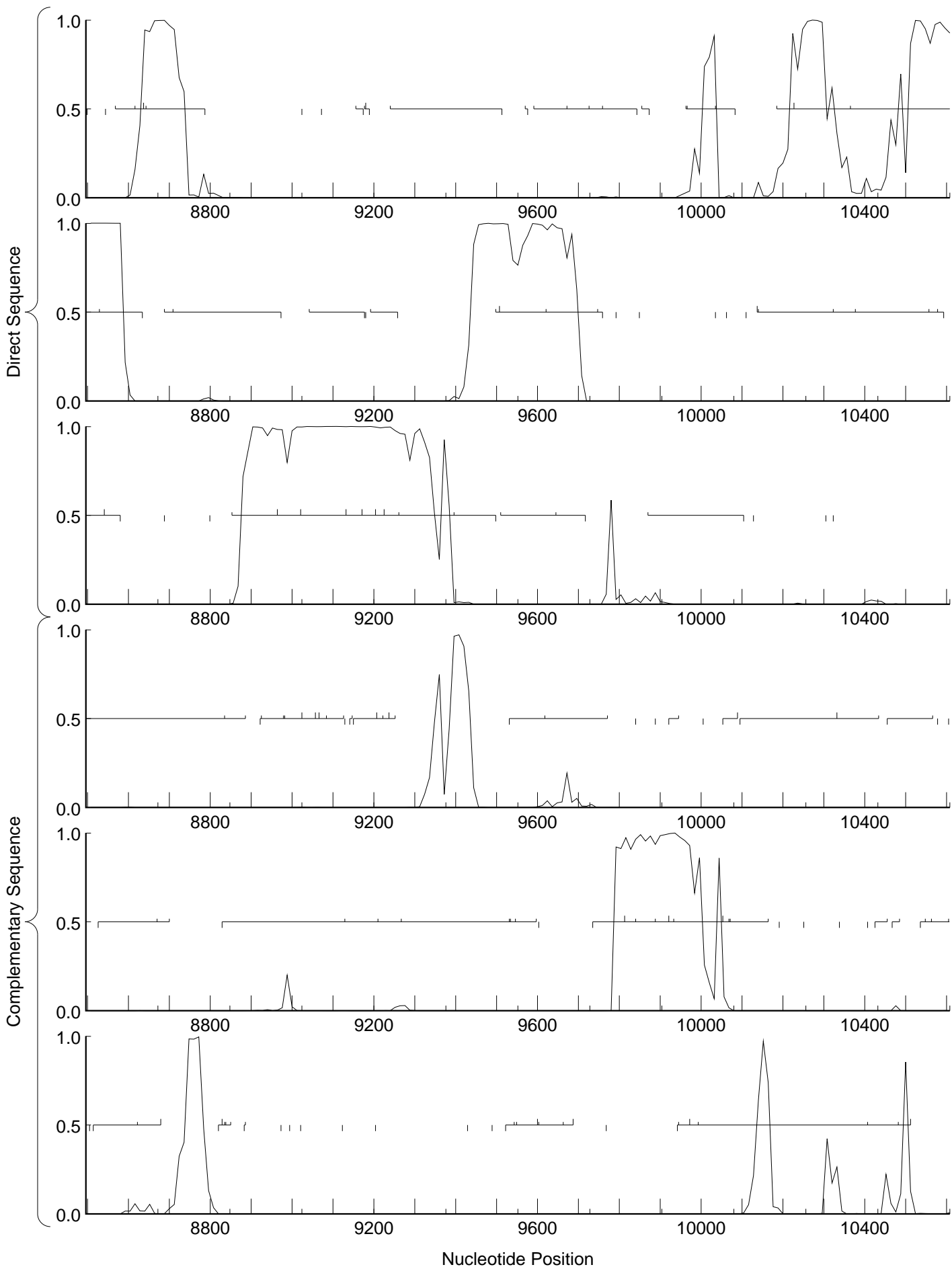
For help or information, send electronic mail to: [custserv@genepro.com](mailto:custserv@genepro.com), or [mark@amber.biology.gatech.edu](mailto:mark@amber.biology.gatech.edu)



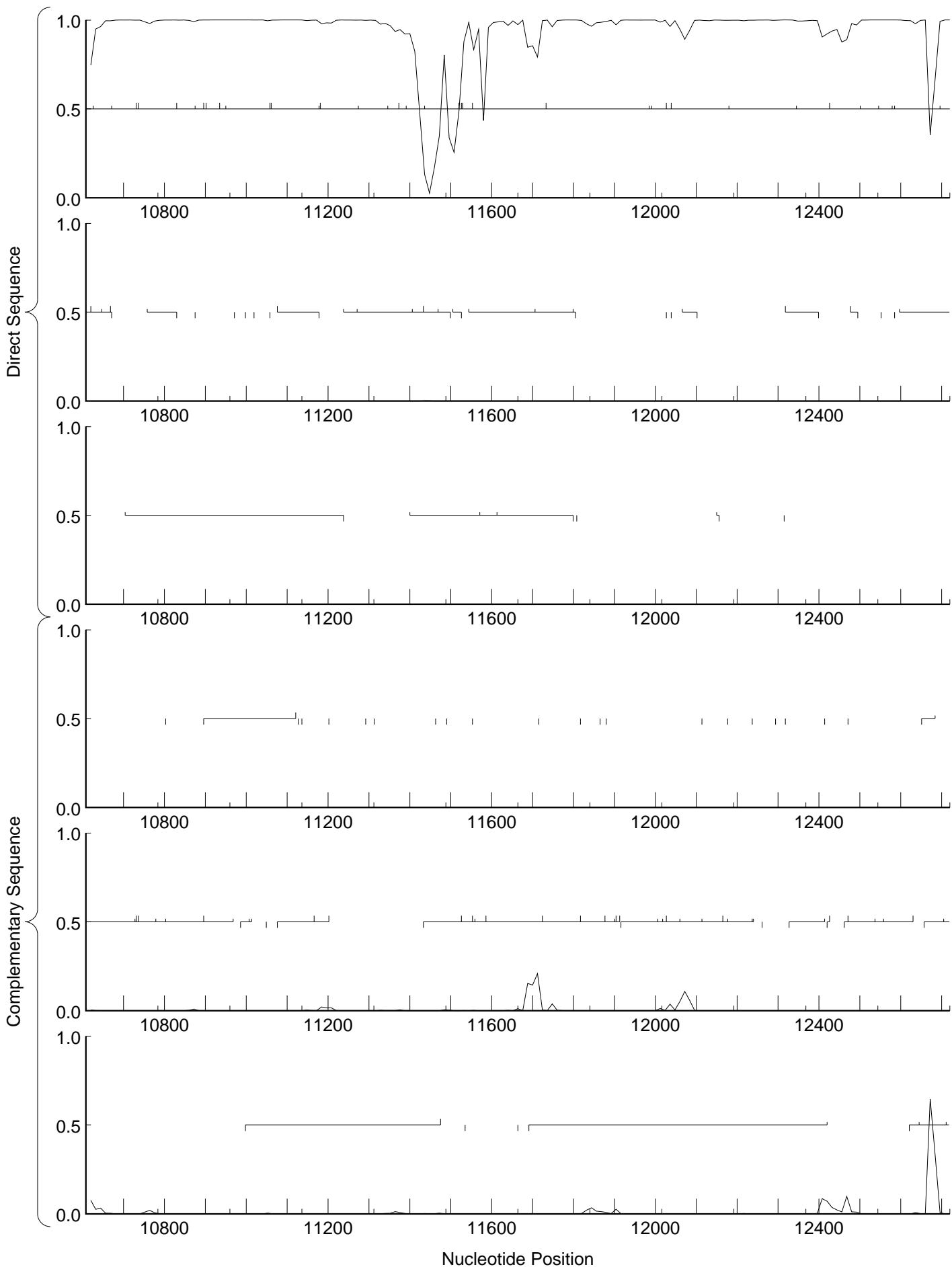




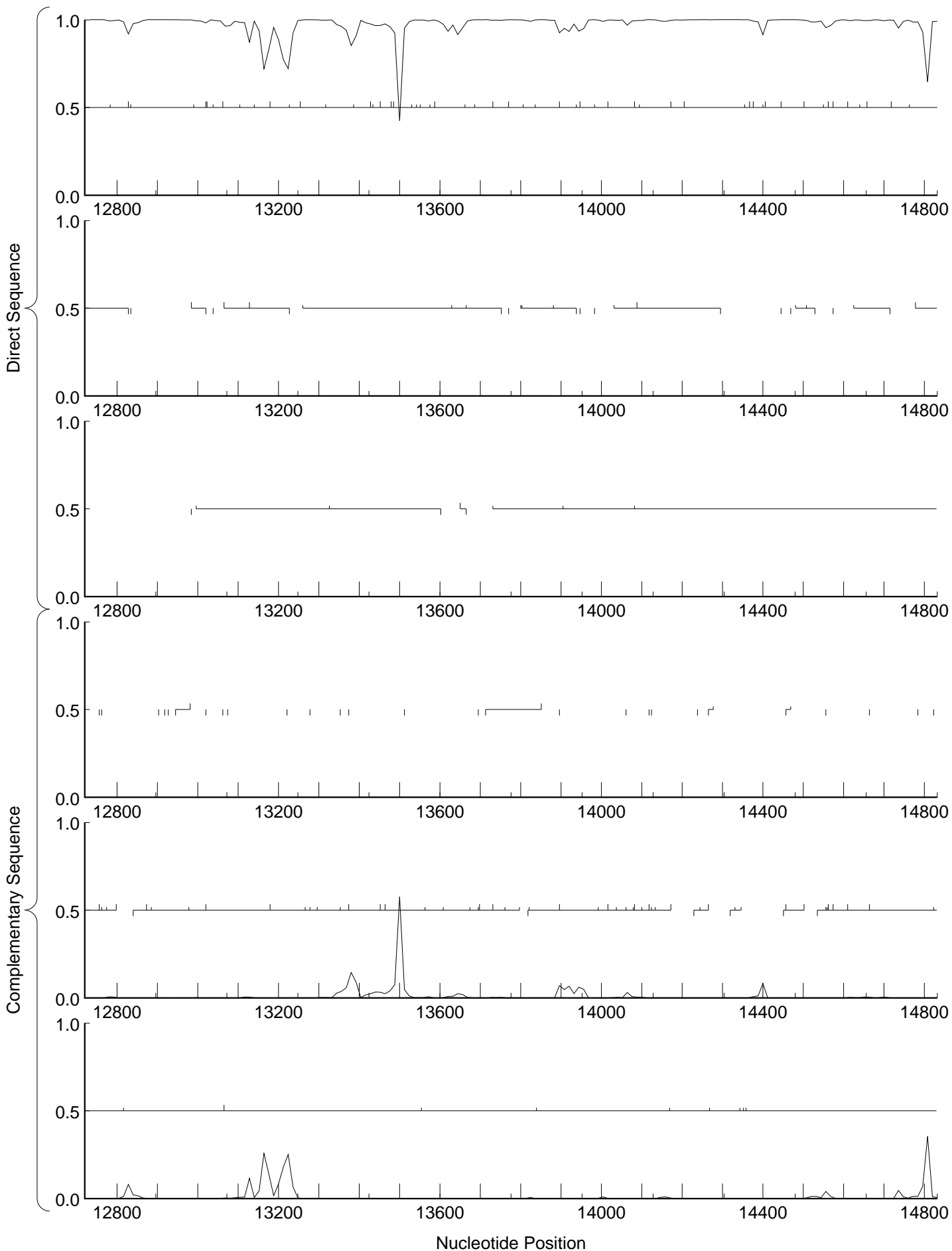




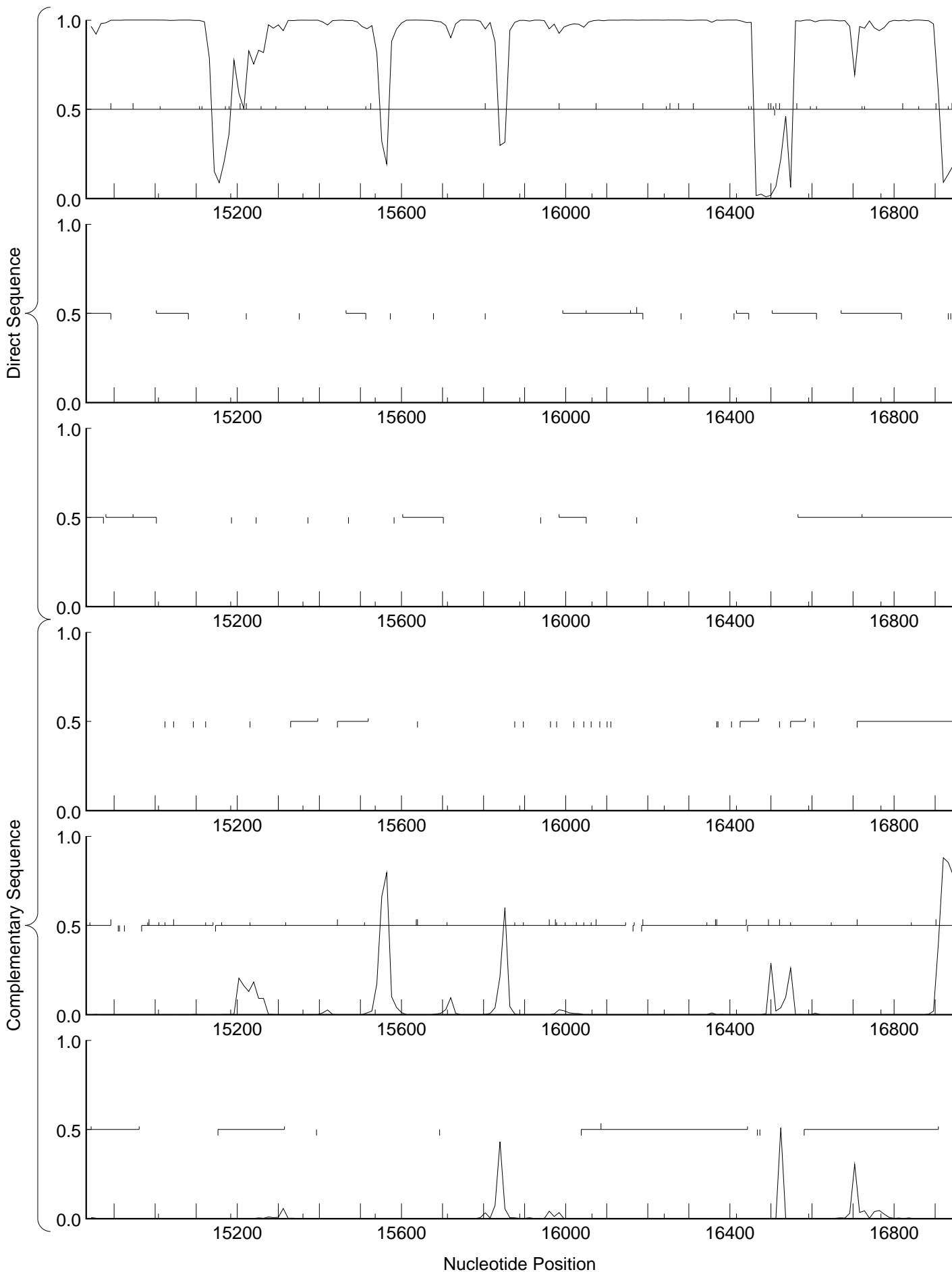
complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 4, Window 96, Step 12, 7/24

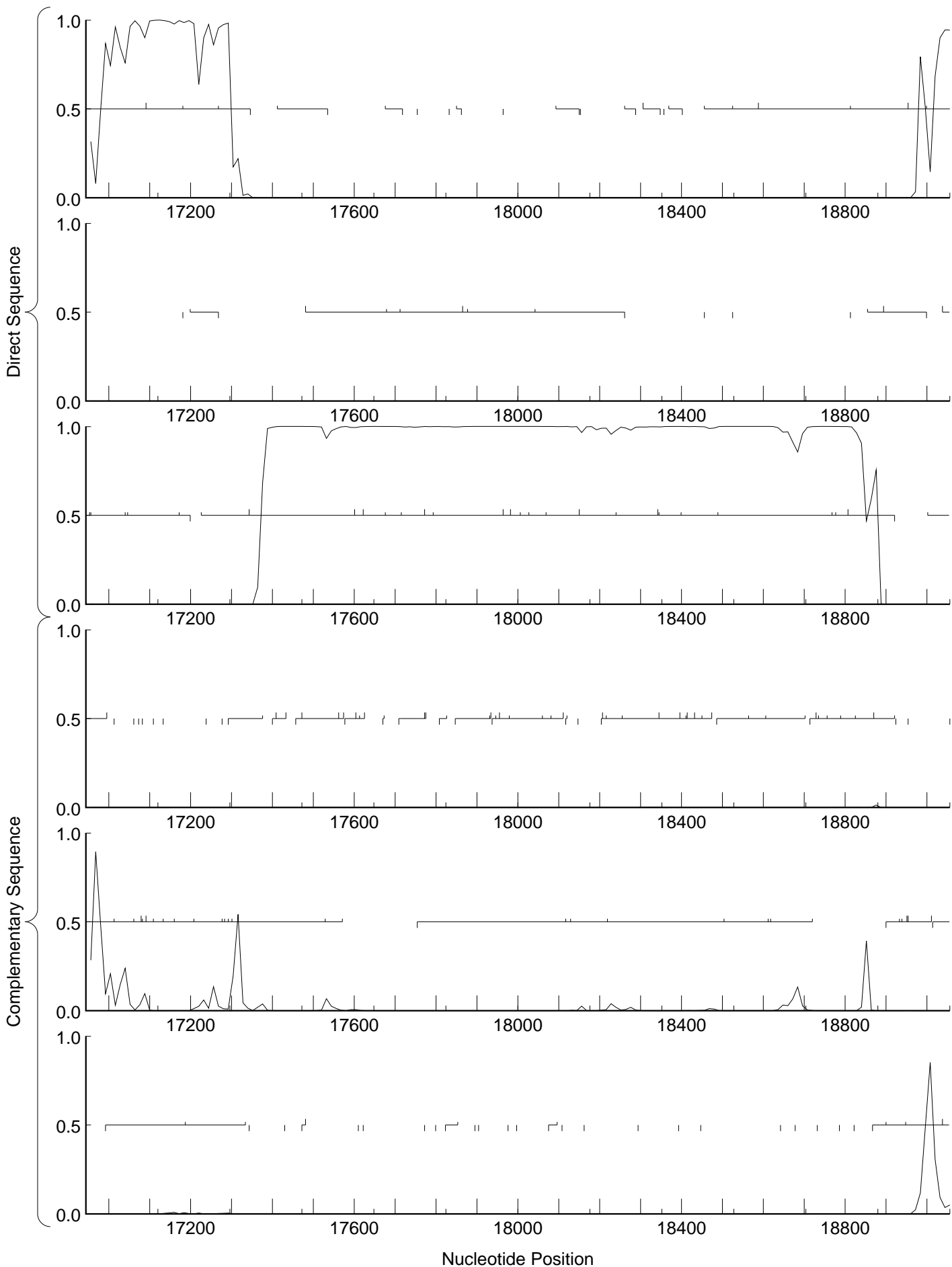


complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 4, Window 96, Step 12, 8/24

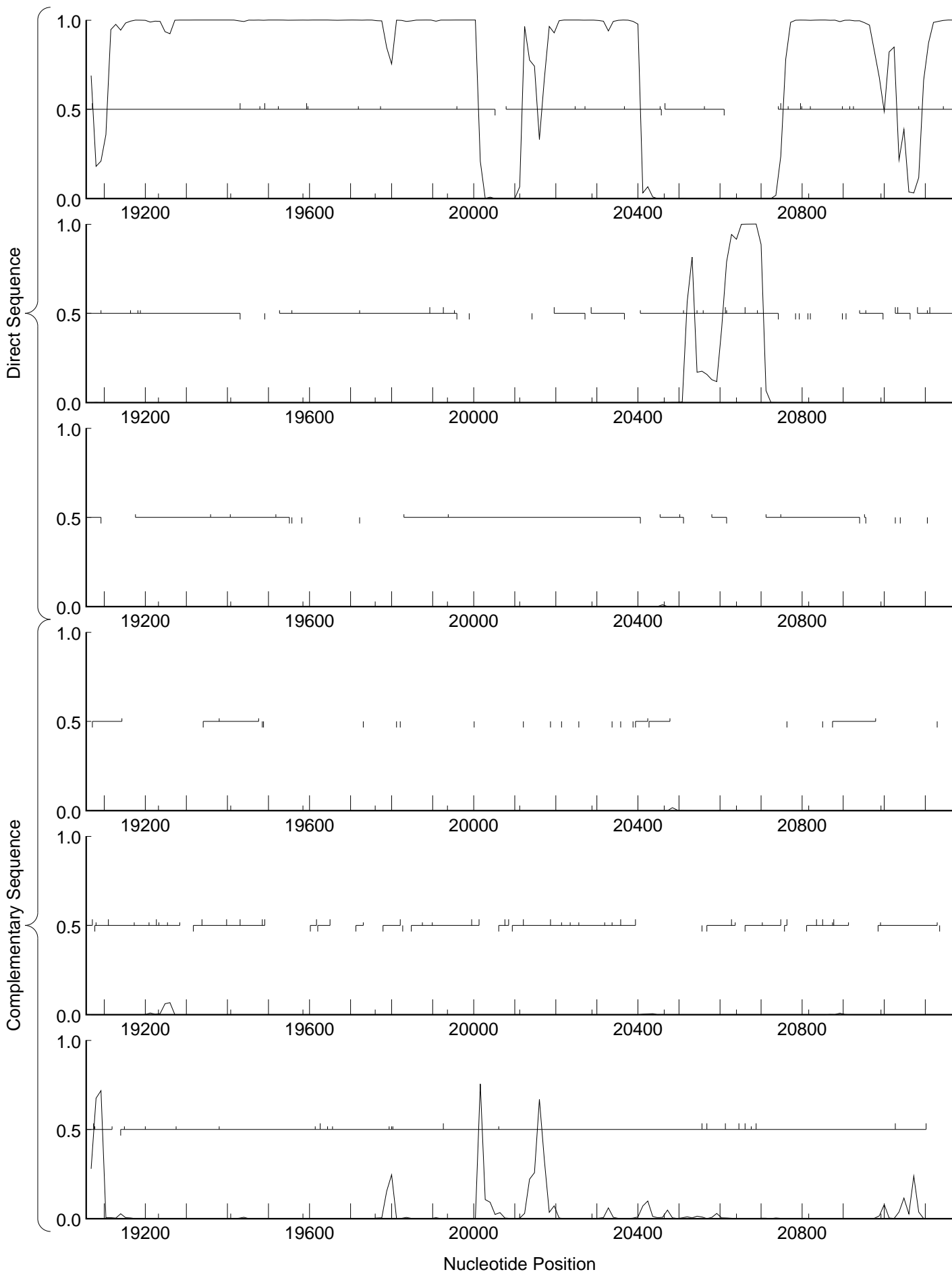


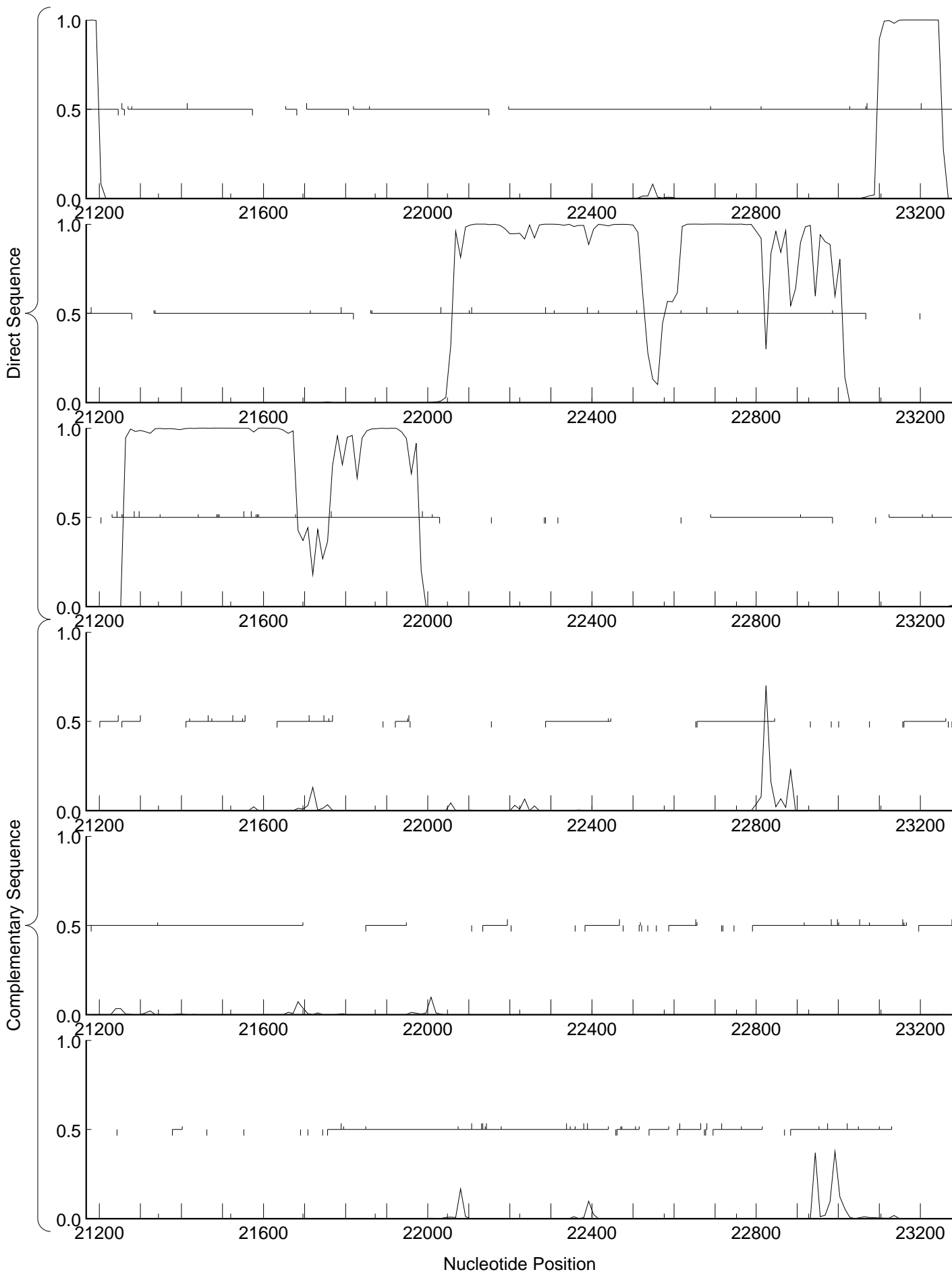




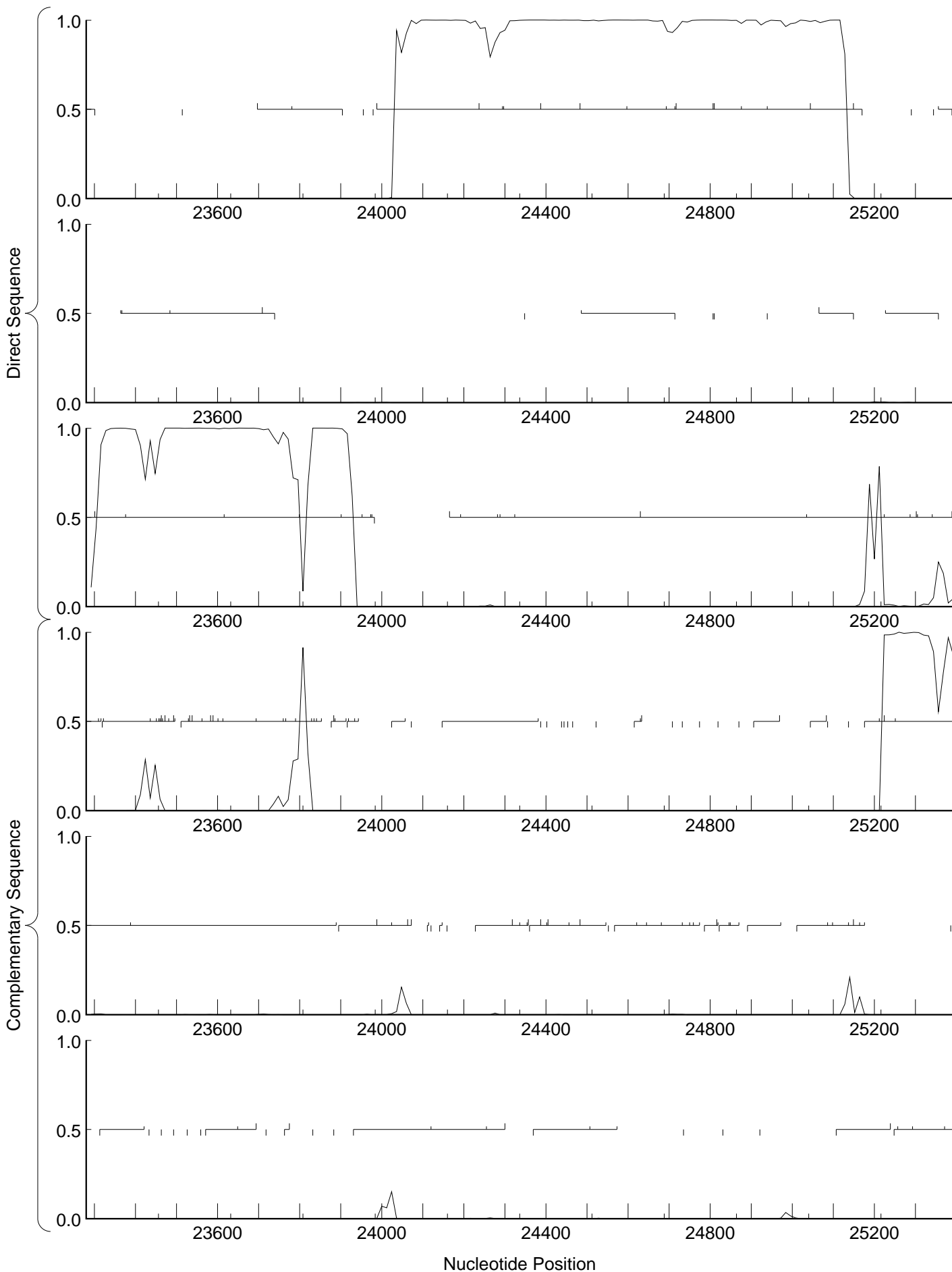


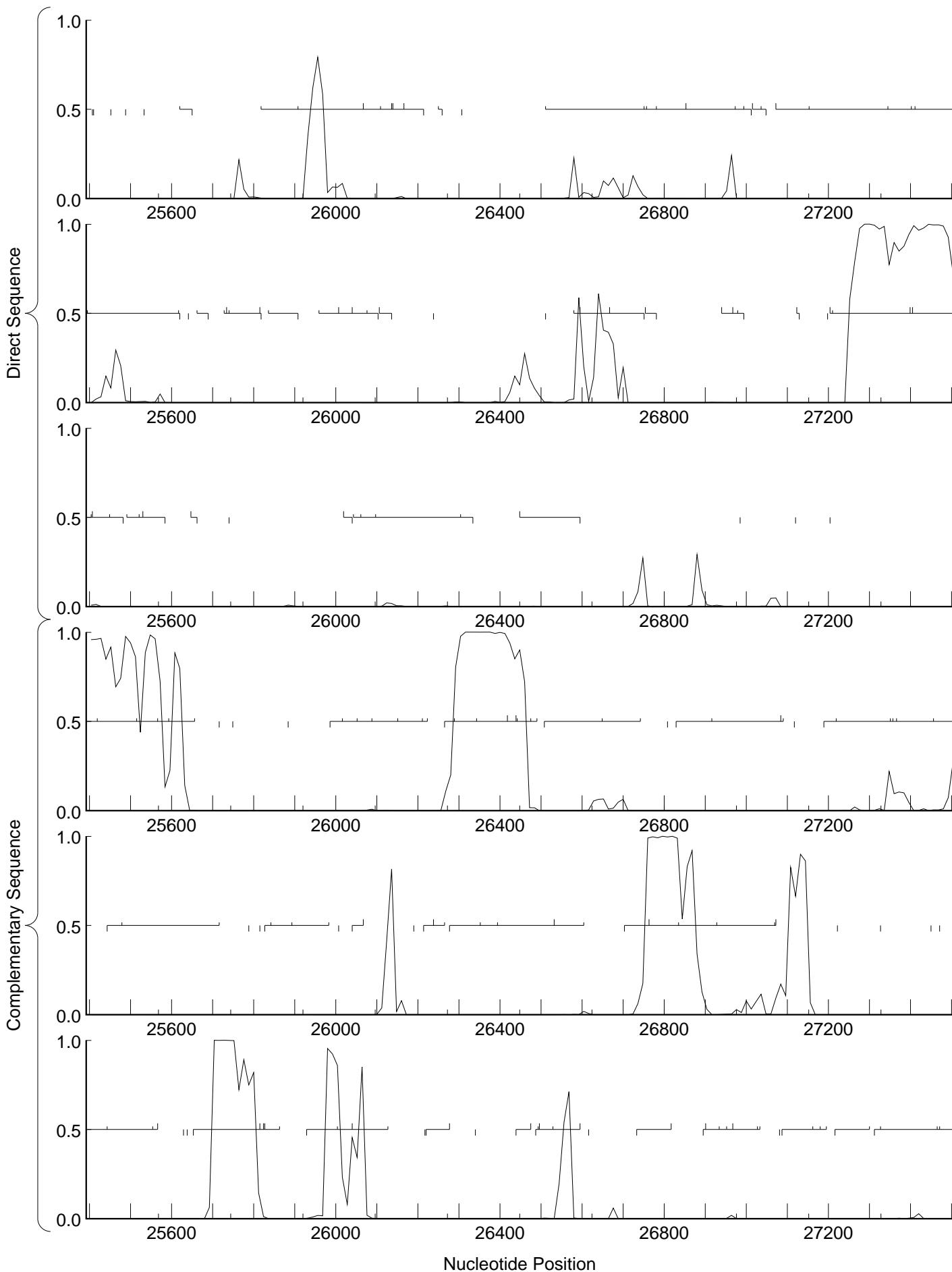
complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 4, Window 96, Step 12, 11/24



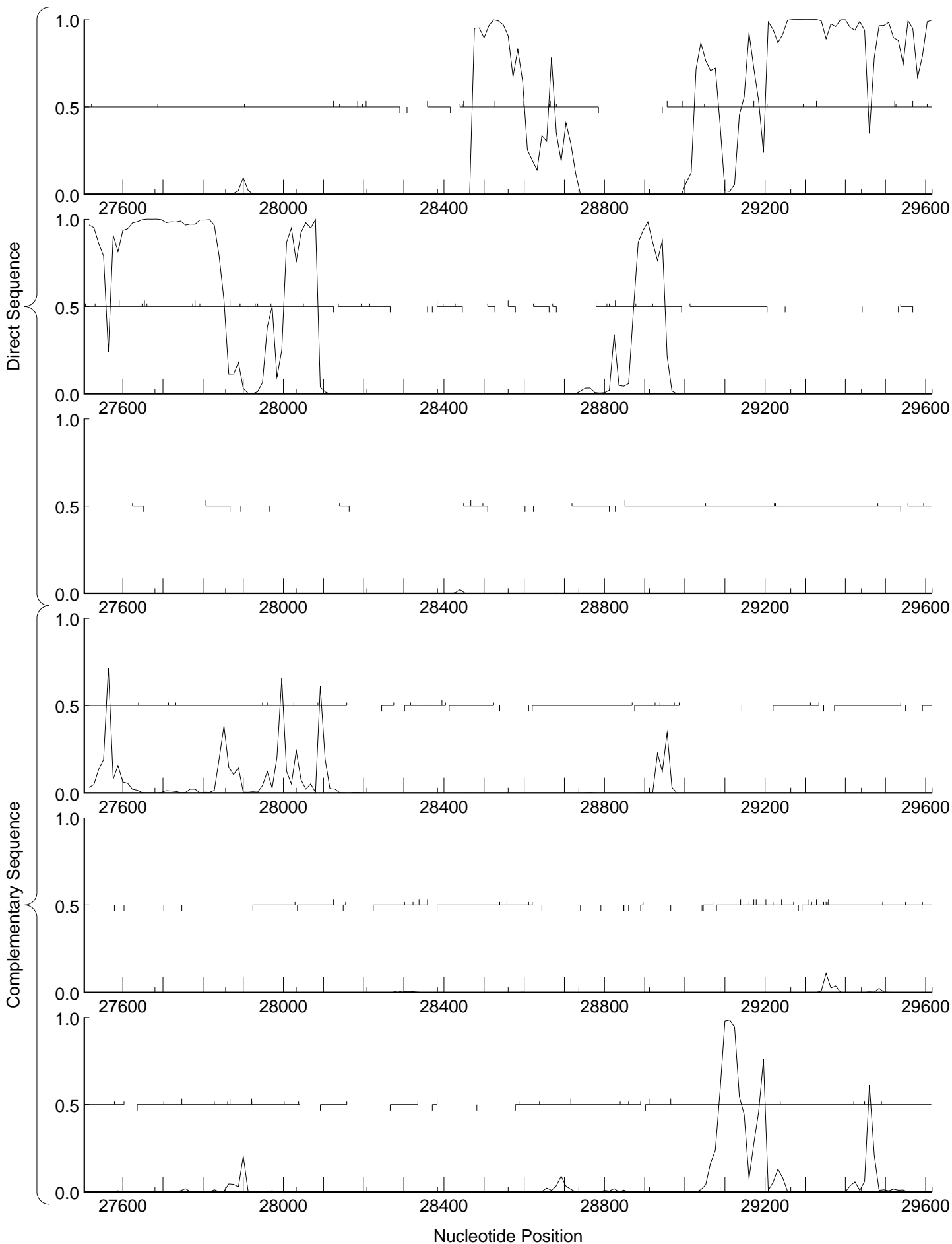


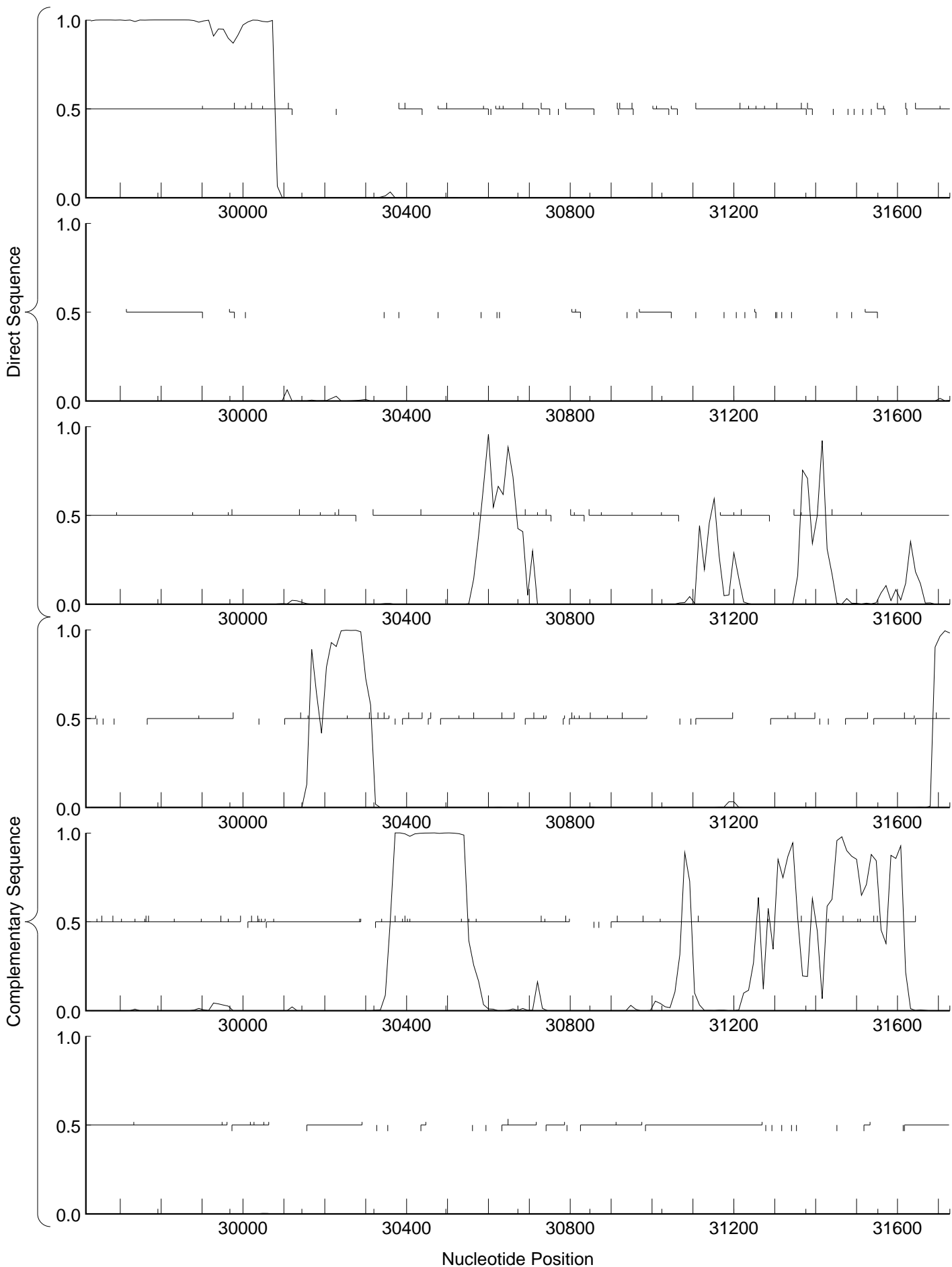
Complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 4, Window 96, Step 12, 13/24





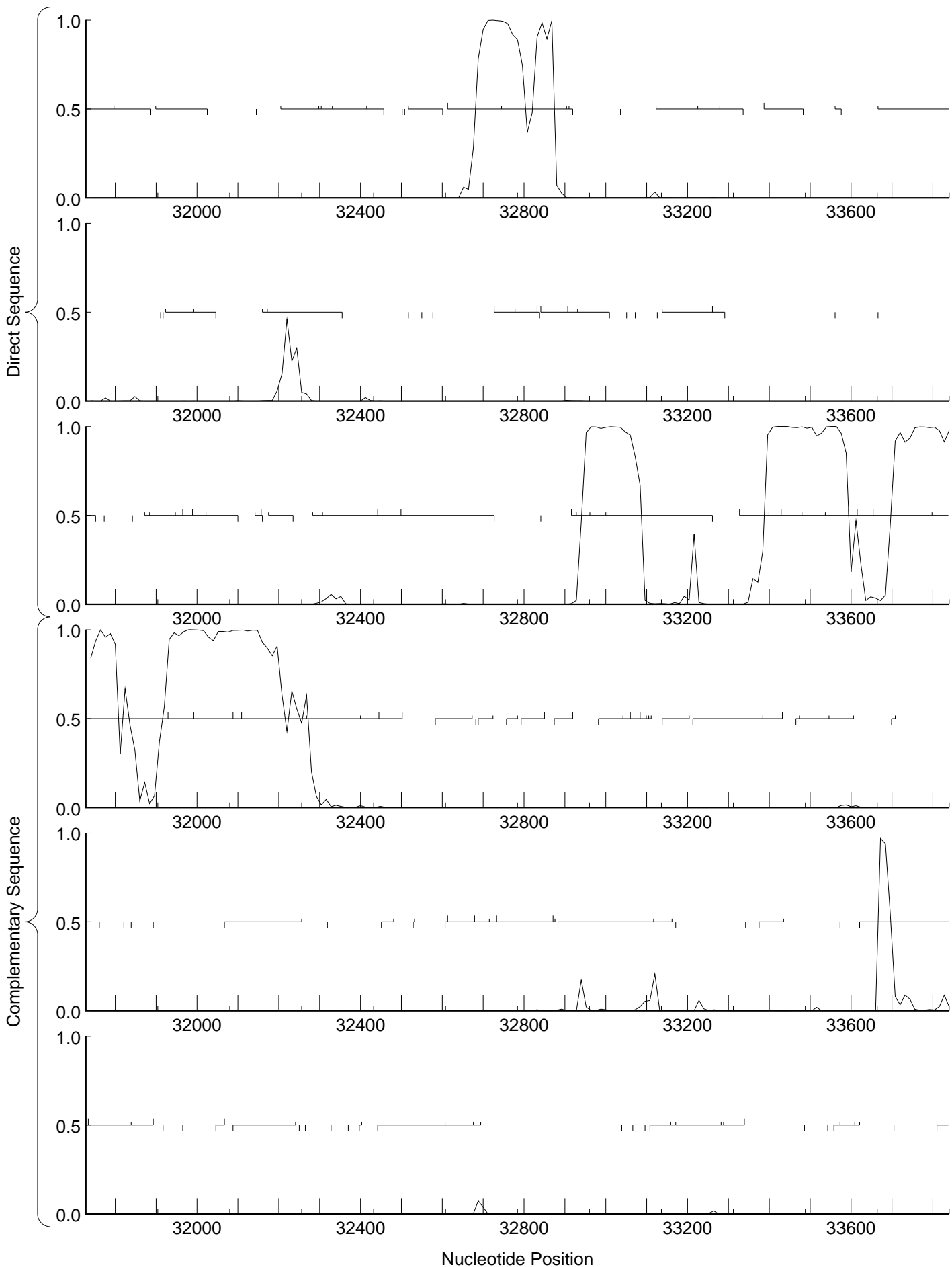
complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 4, Window 96, Step 12, 15/24

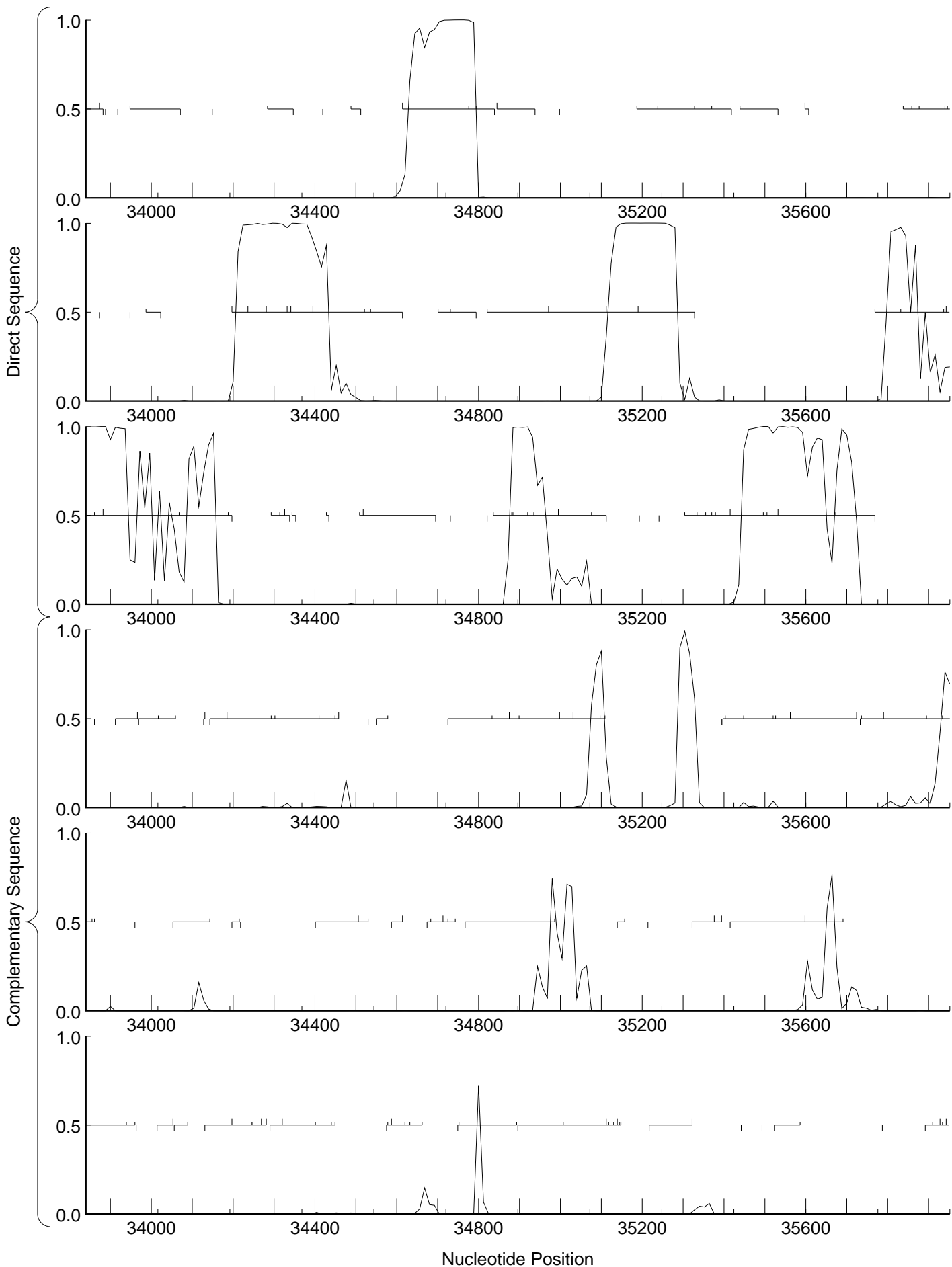


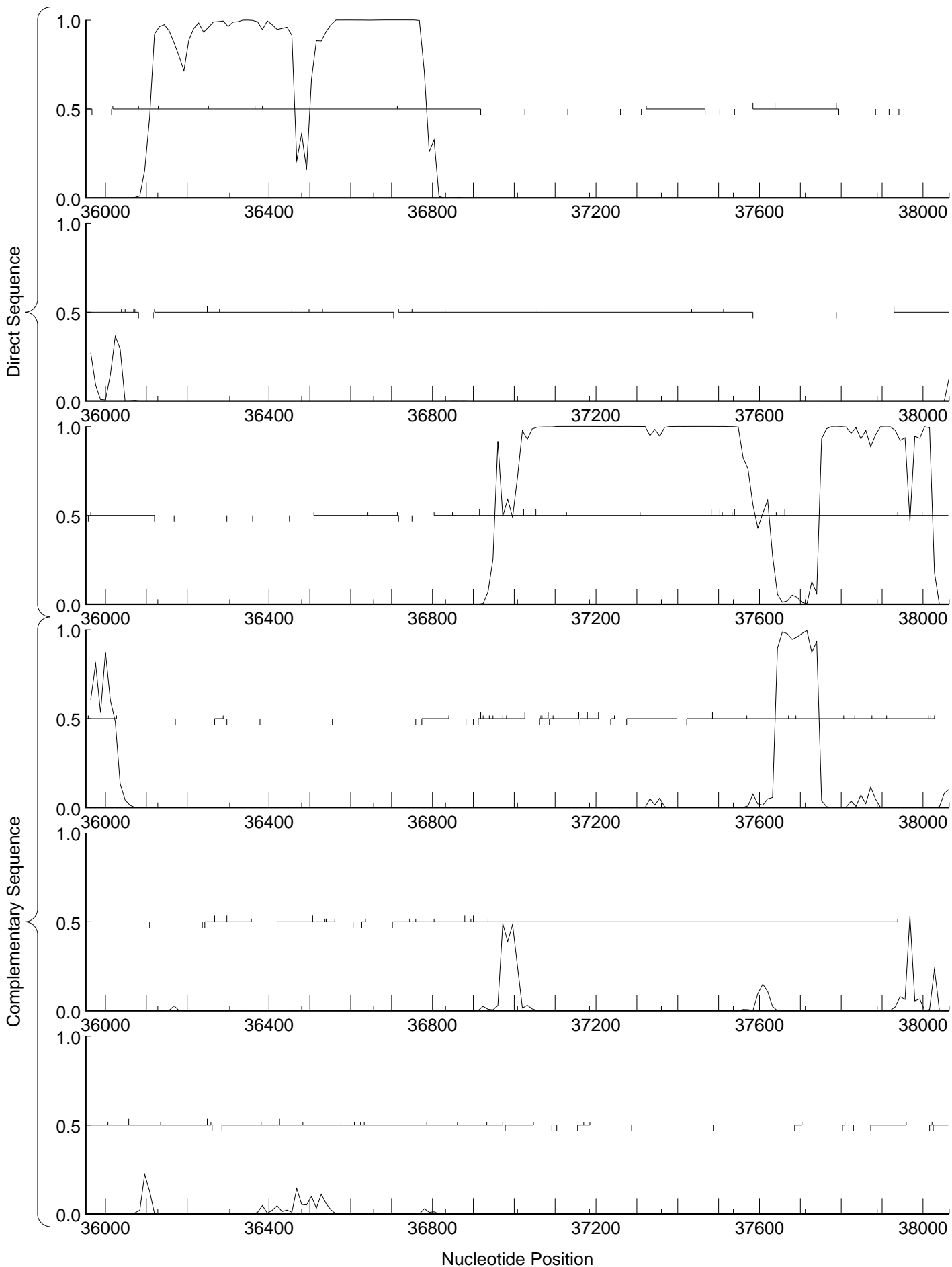


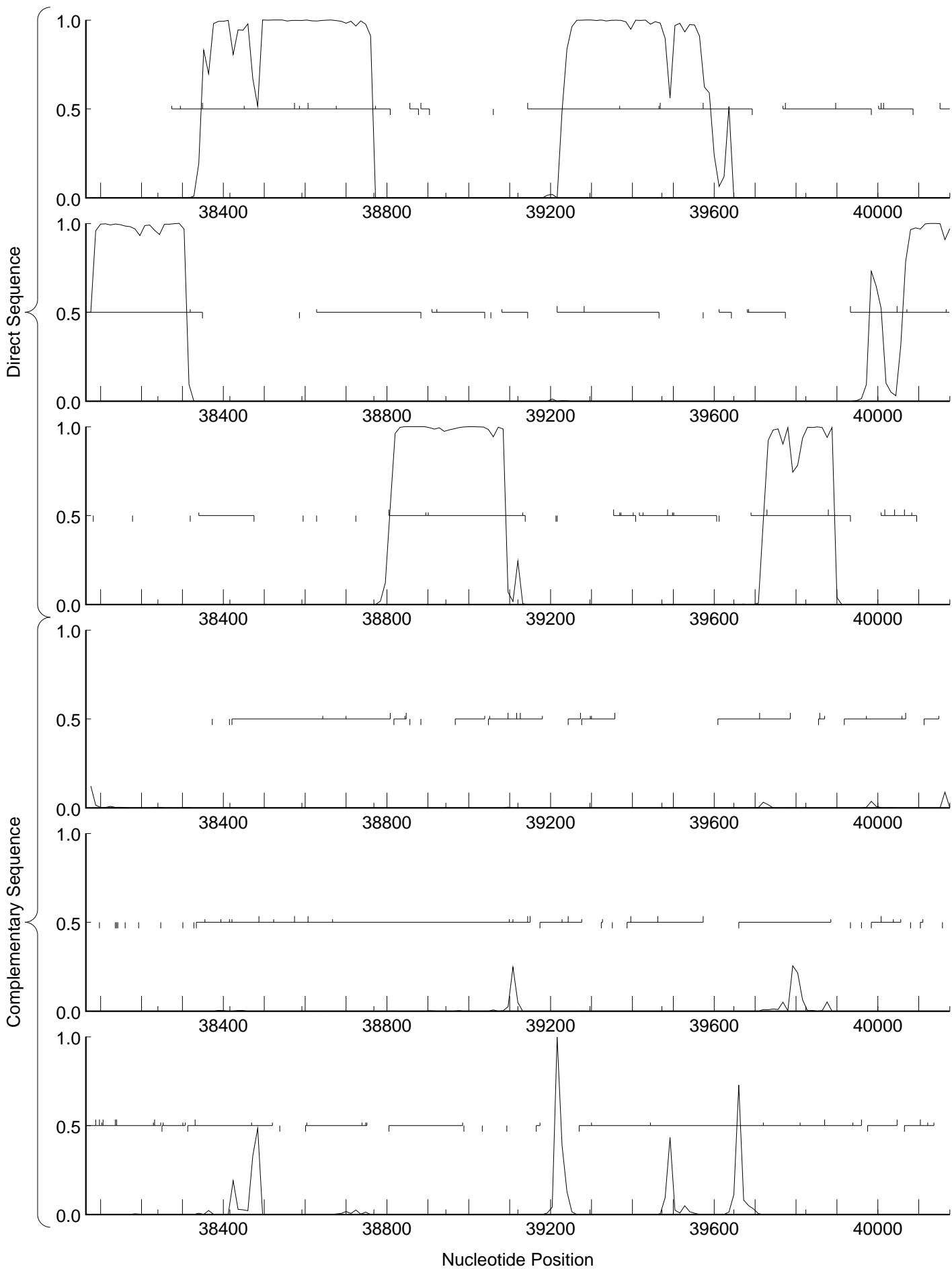


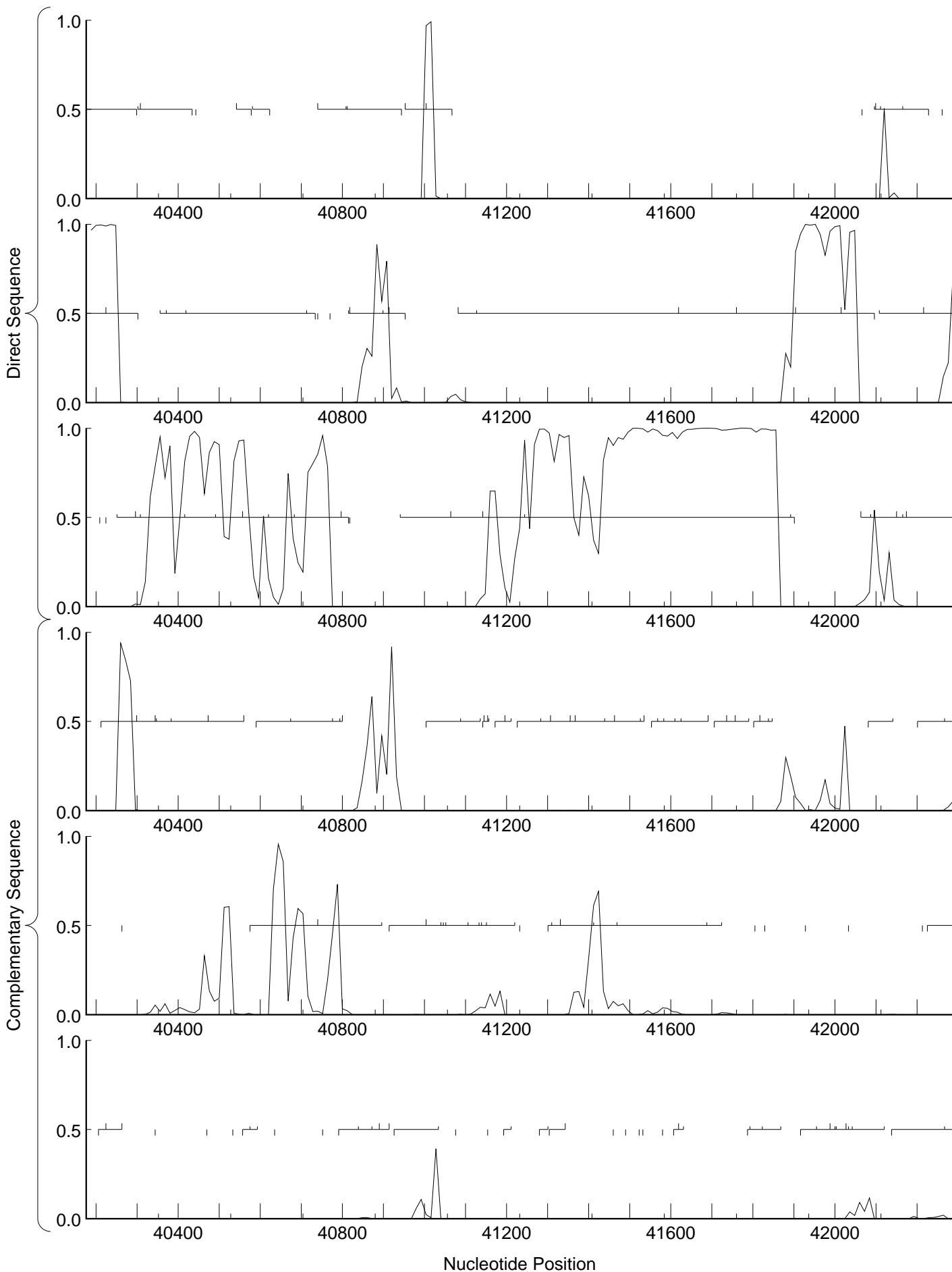
complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 4, Window 96, Step 12, 17/24

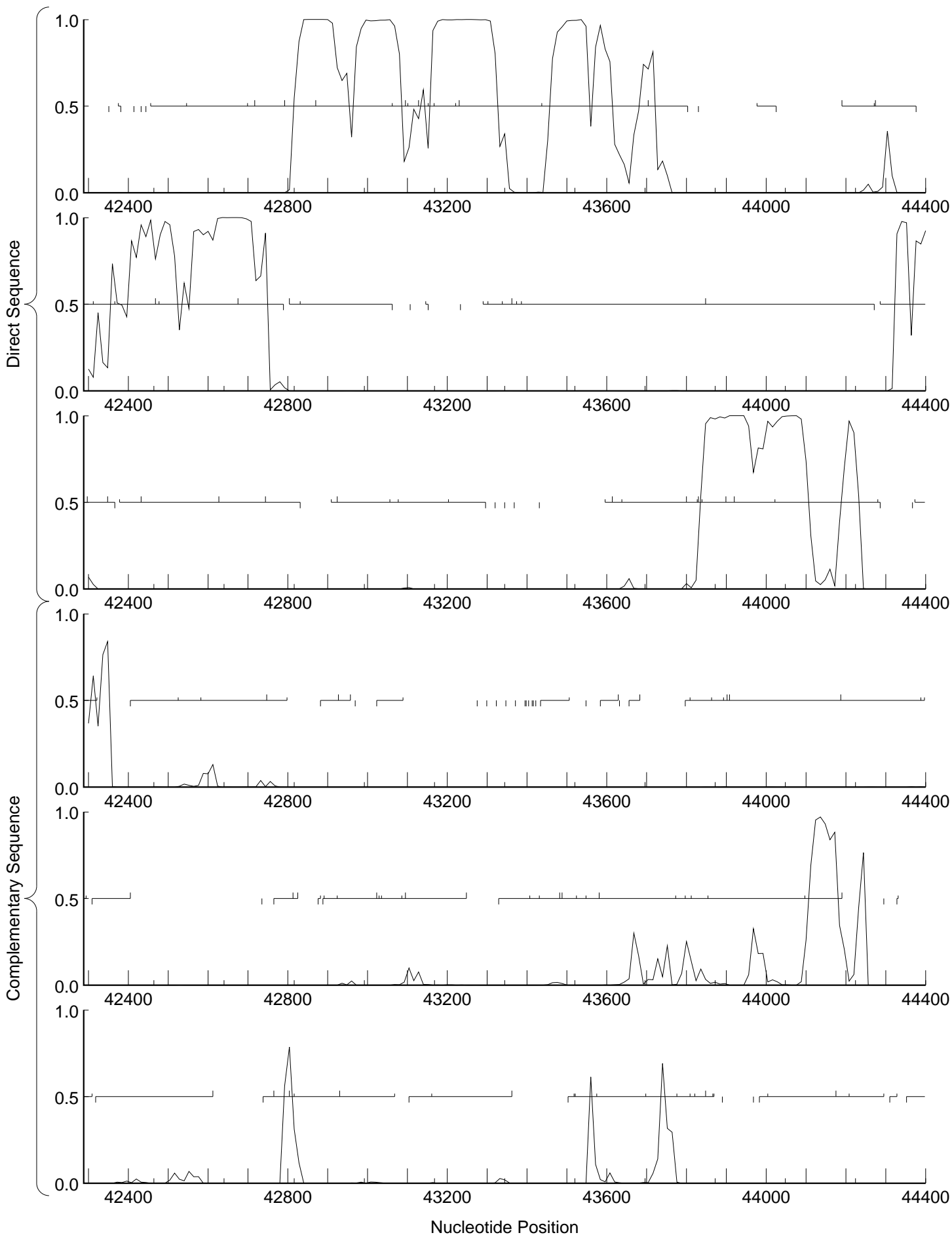












Complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 4, Window 96, Step 12, 23/24

