

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Gordonia phage LordFarquaad complete sequence, 47863 bp including 11-base 3' overhang (TACC
Analysis Date : 3/25/25 at 14:13:46
Pages : 24
Sequence Length : 47863 bp
GC Content : 66.83%

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,

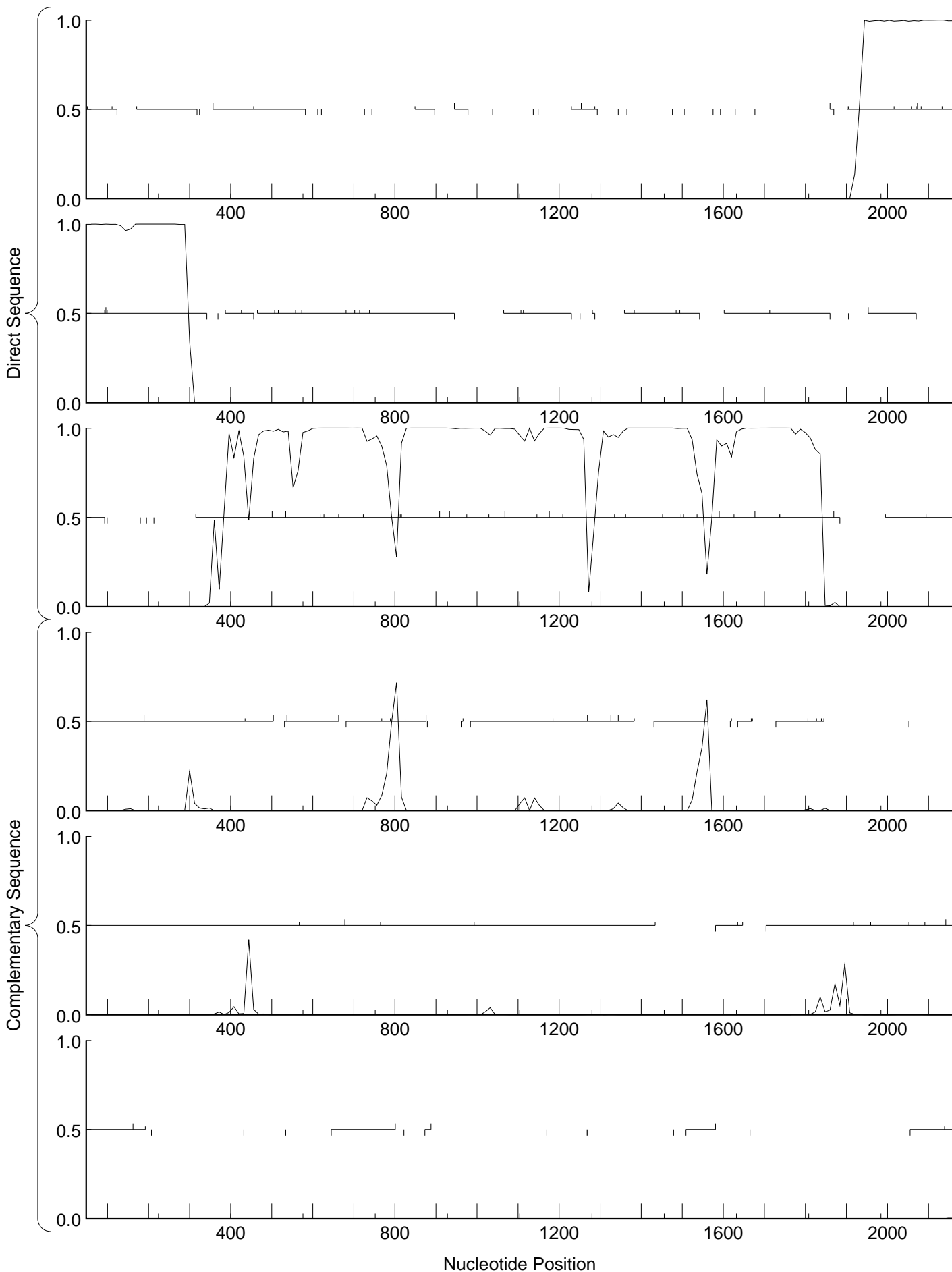
MATRIX INFORMATION

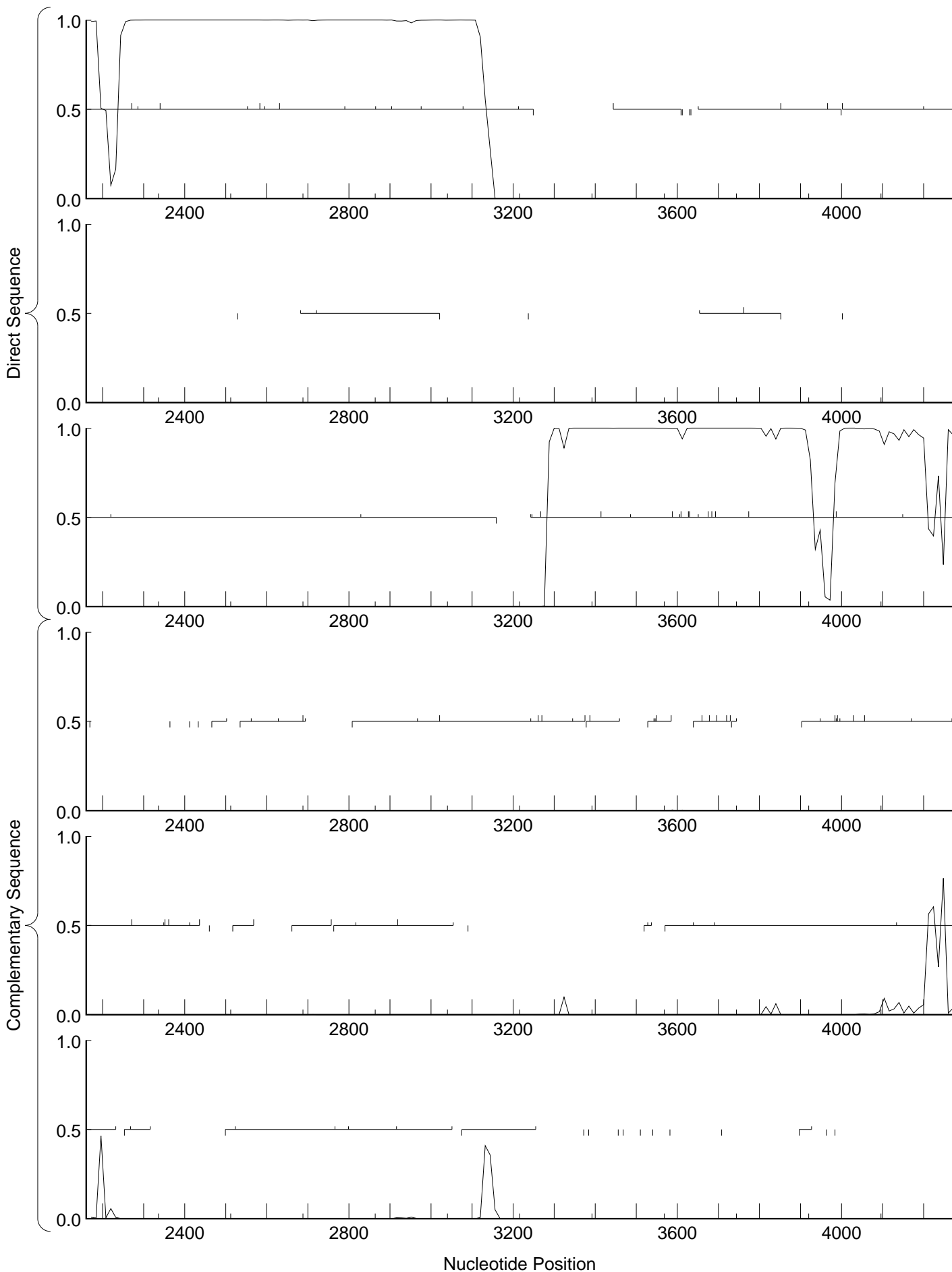
Matrix : -
Author : -
Order : 4

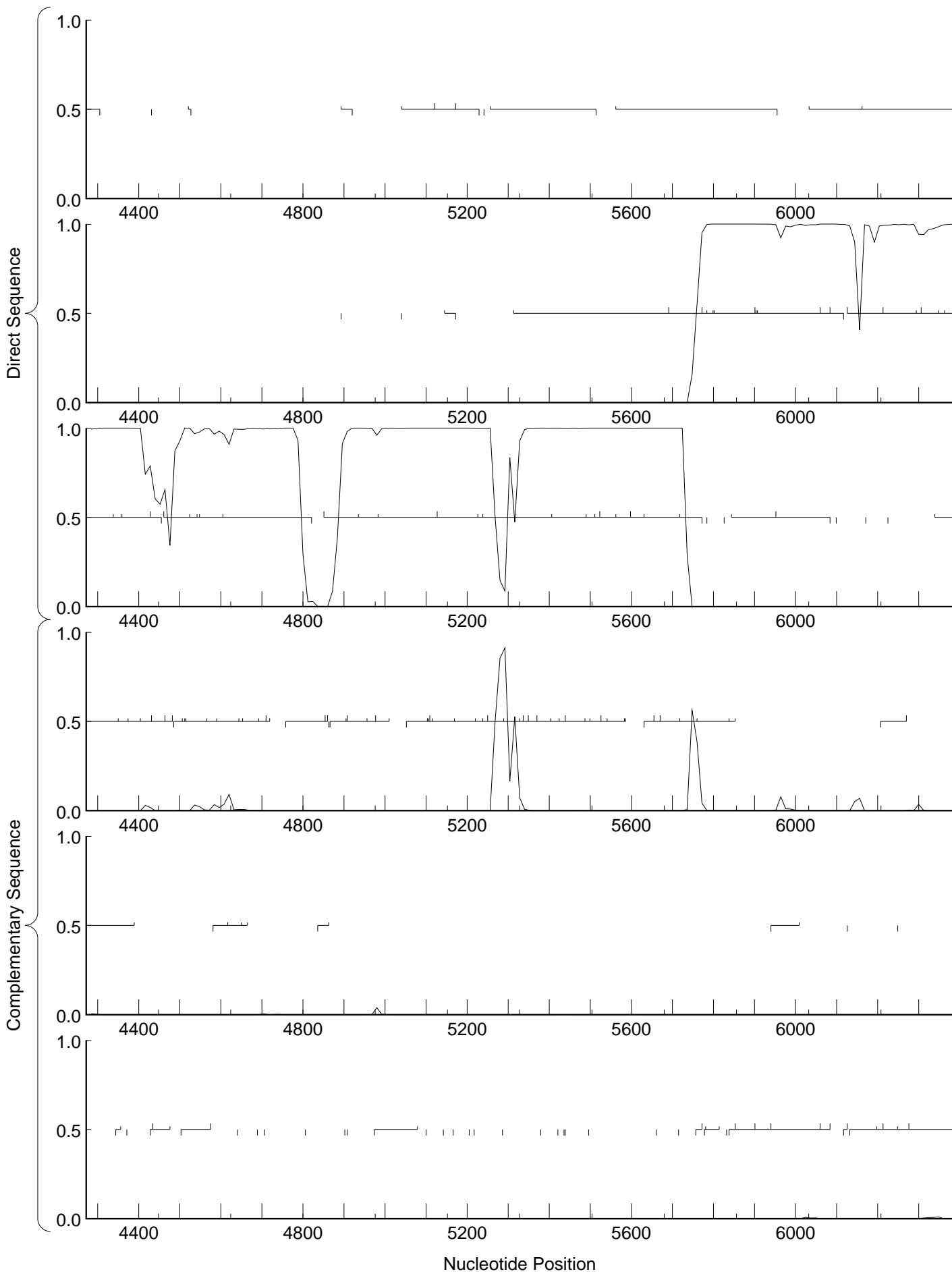
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

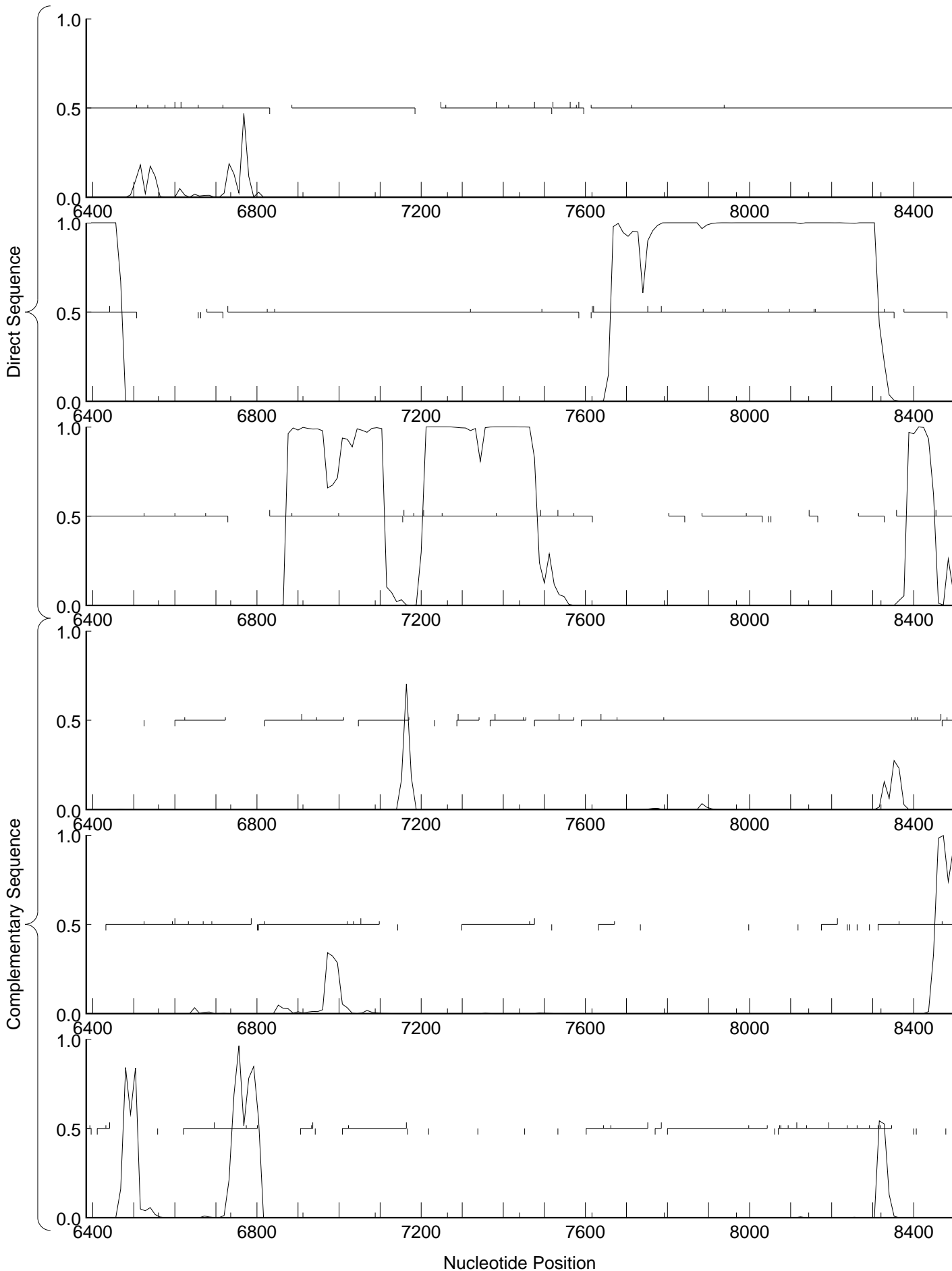
Matrix notes & comments

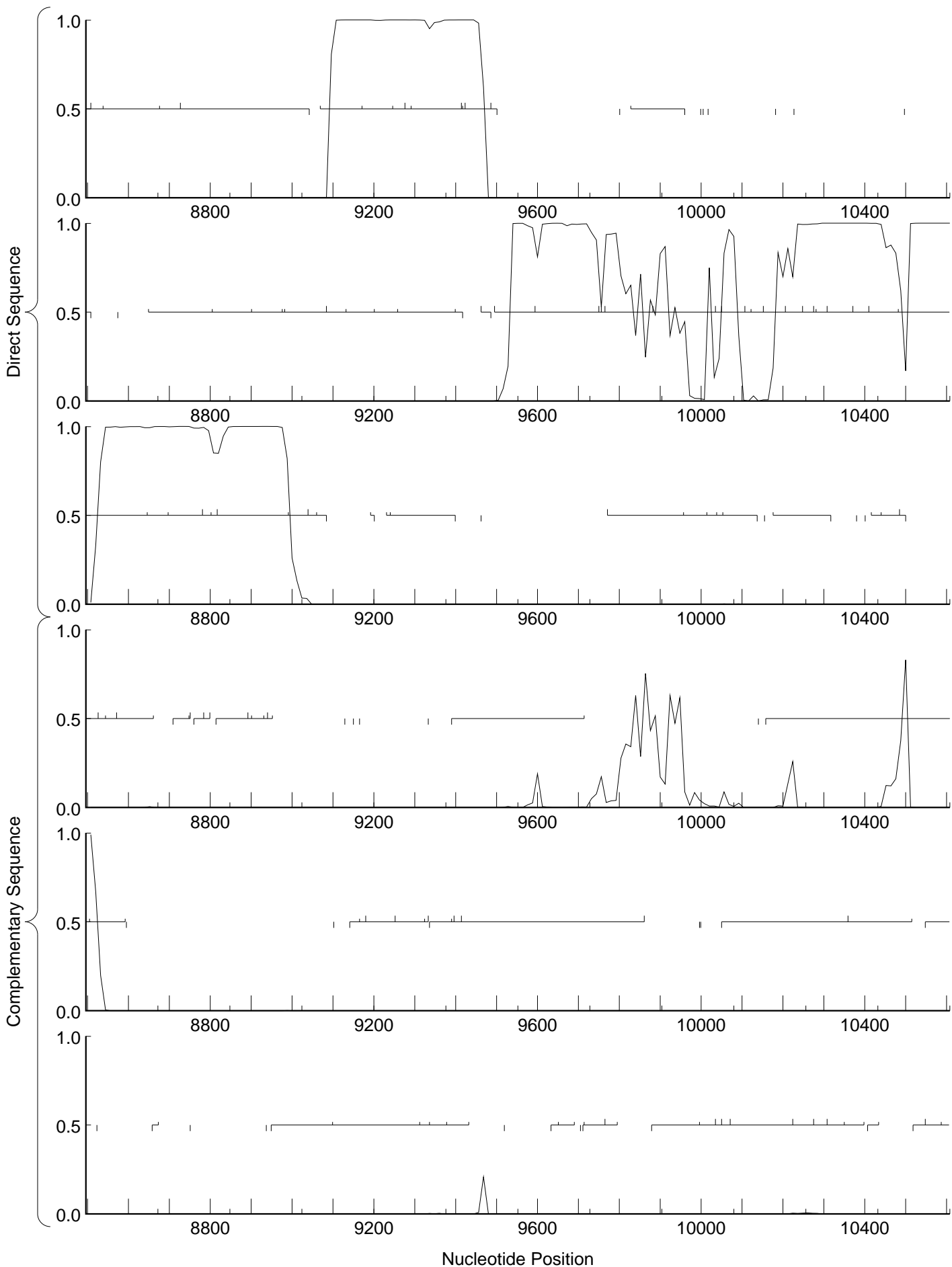
Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:36:14 2018

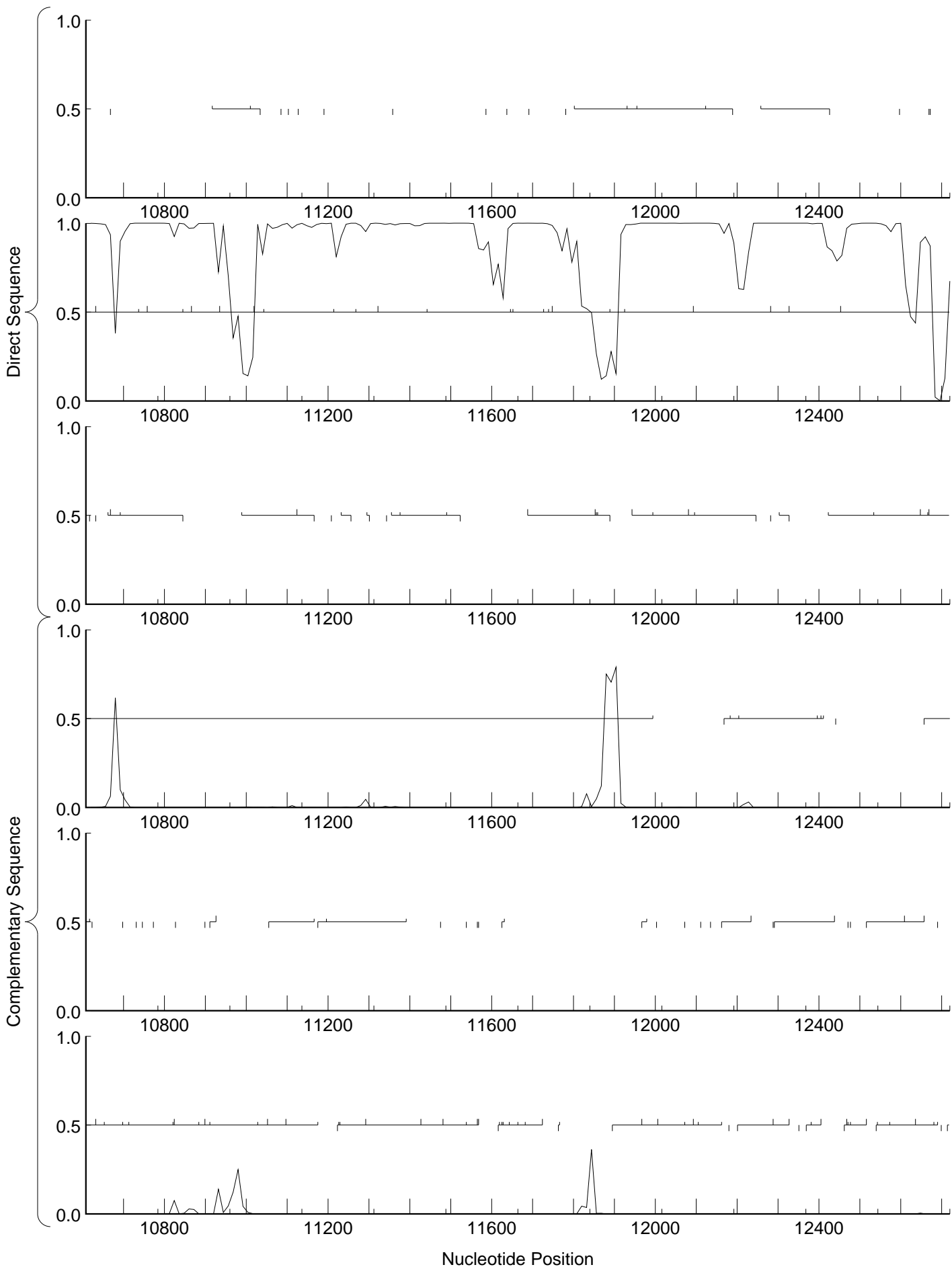


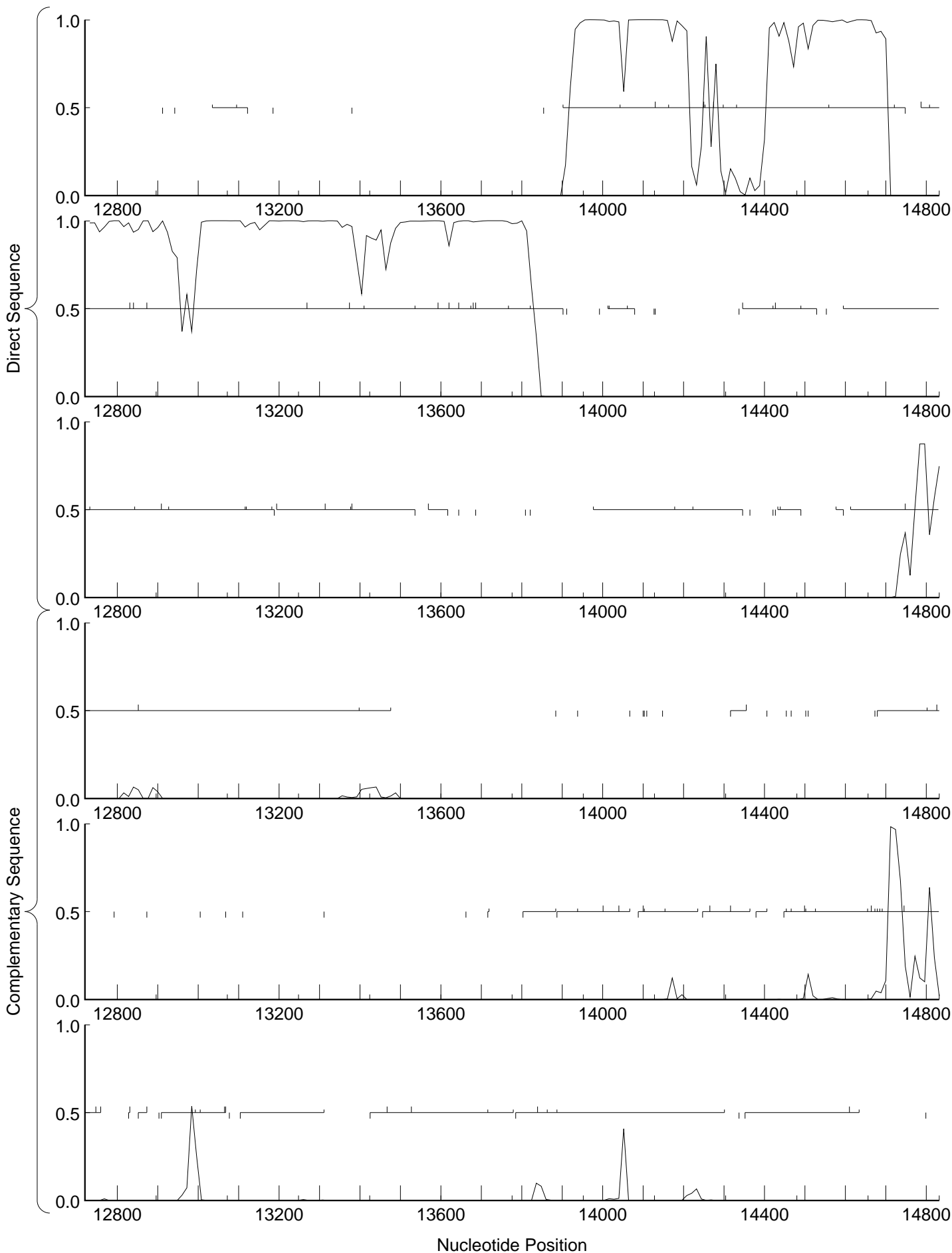


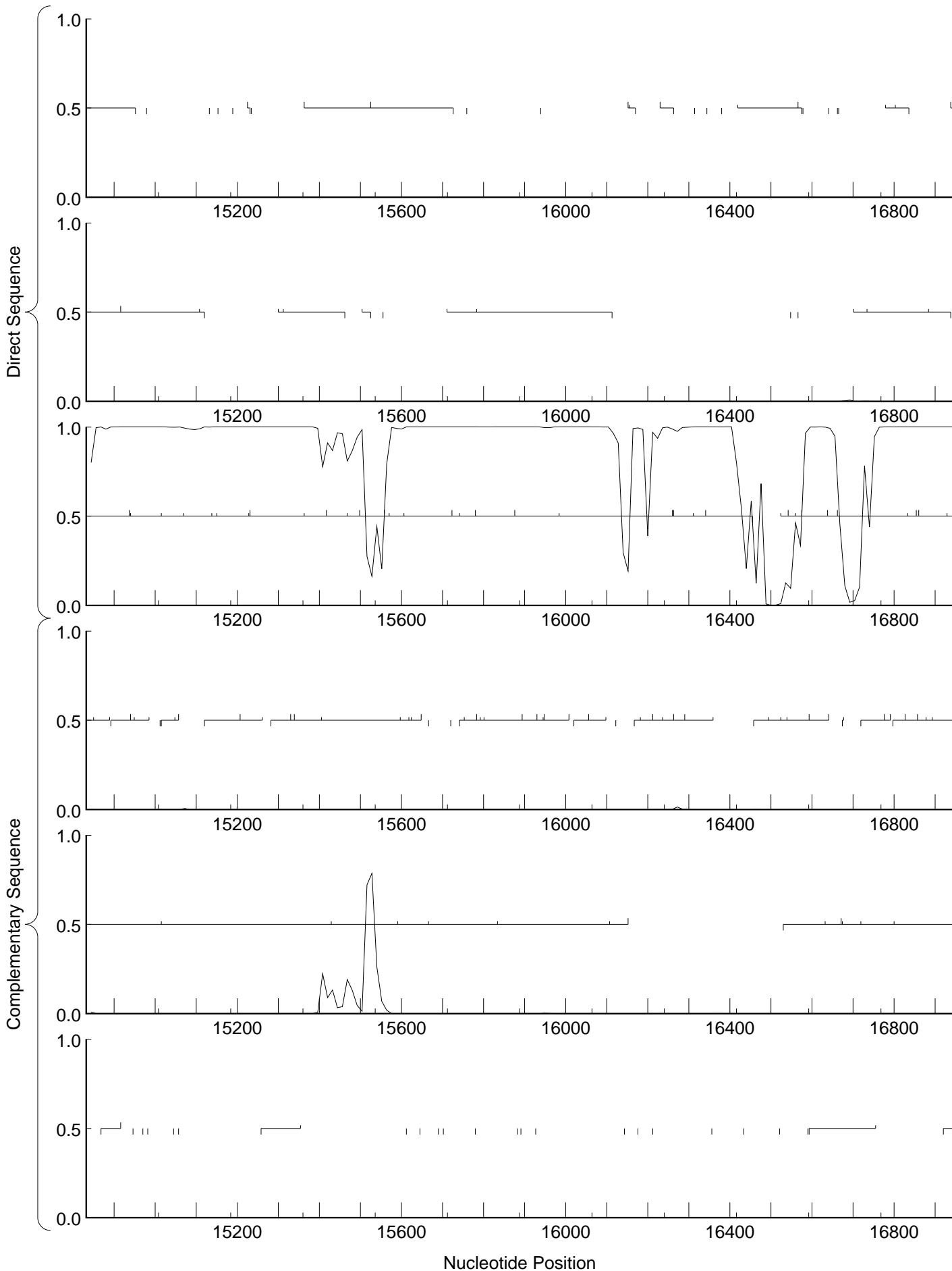


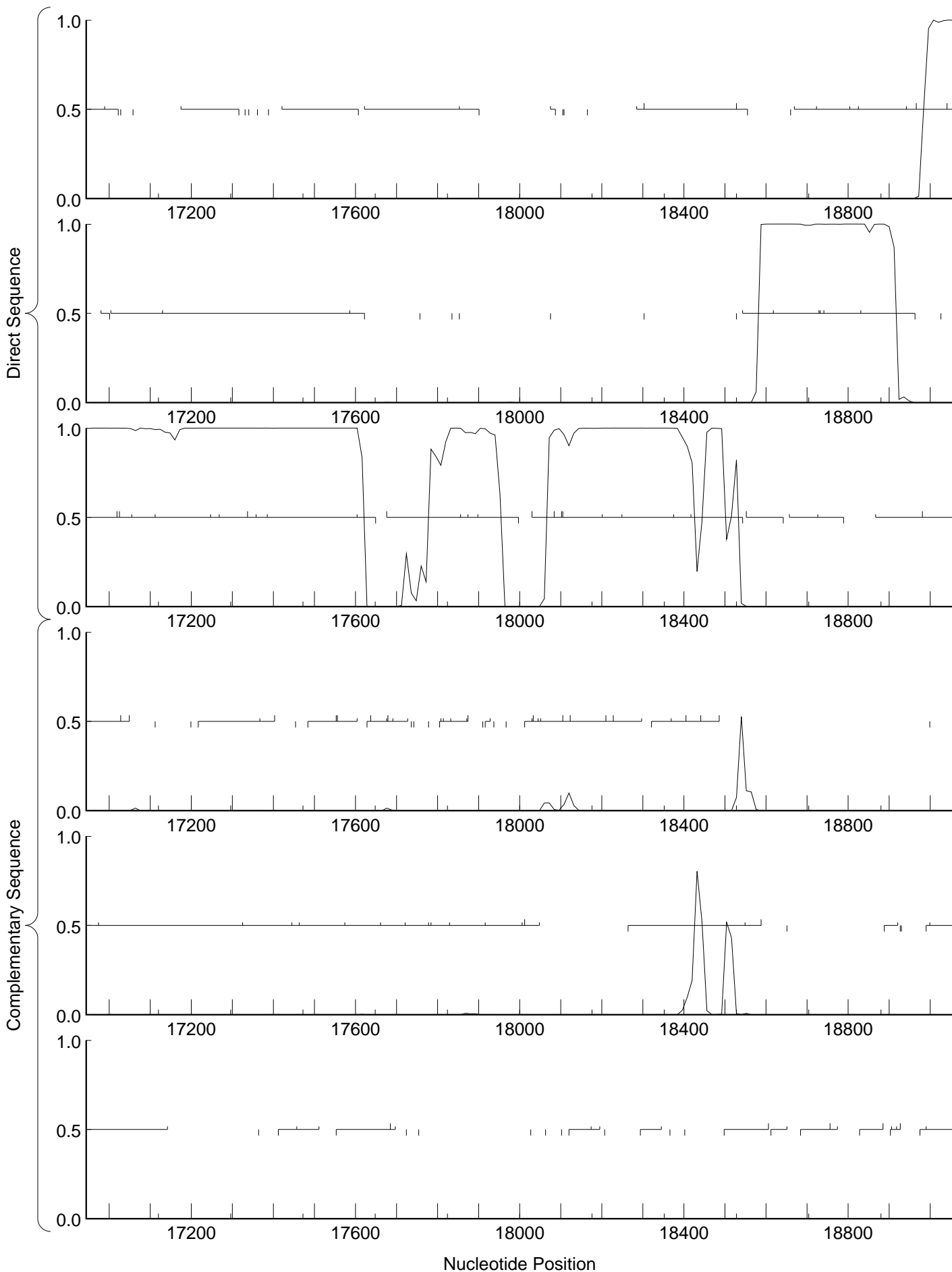


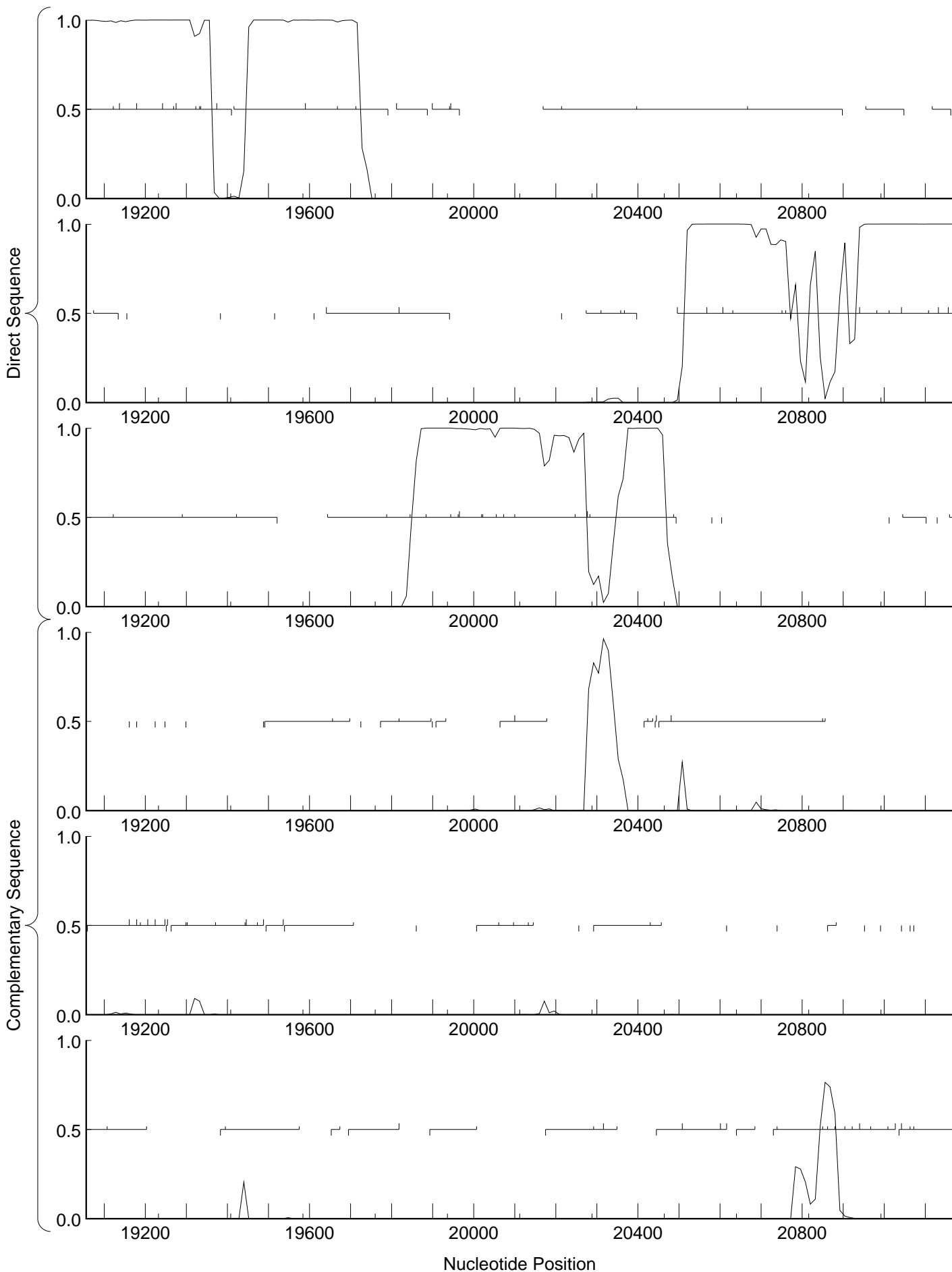


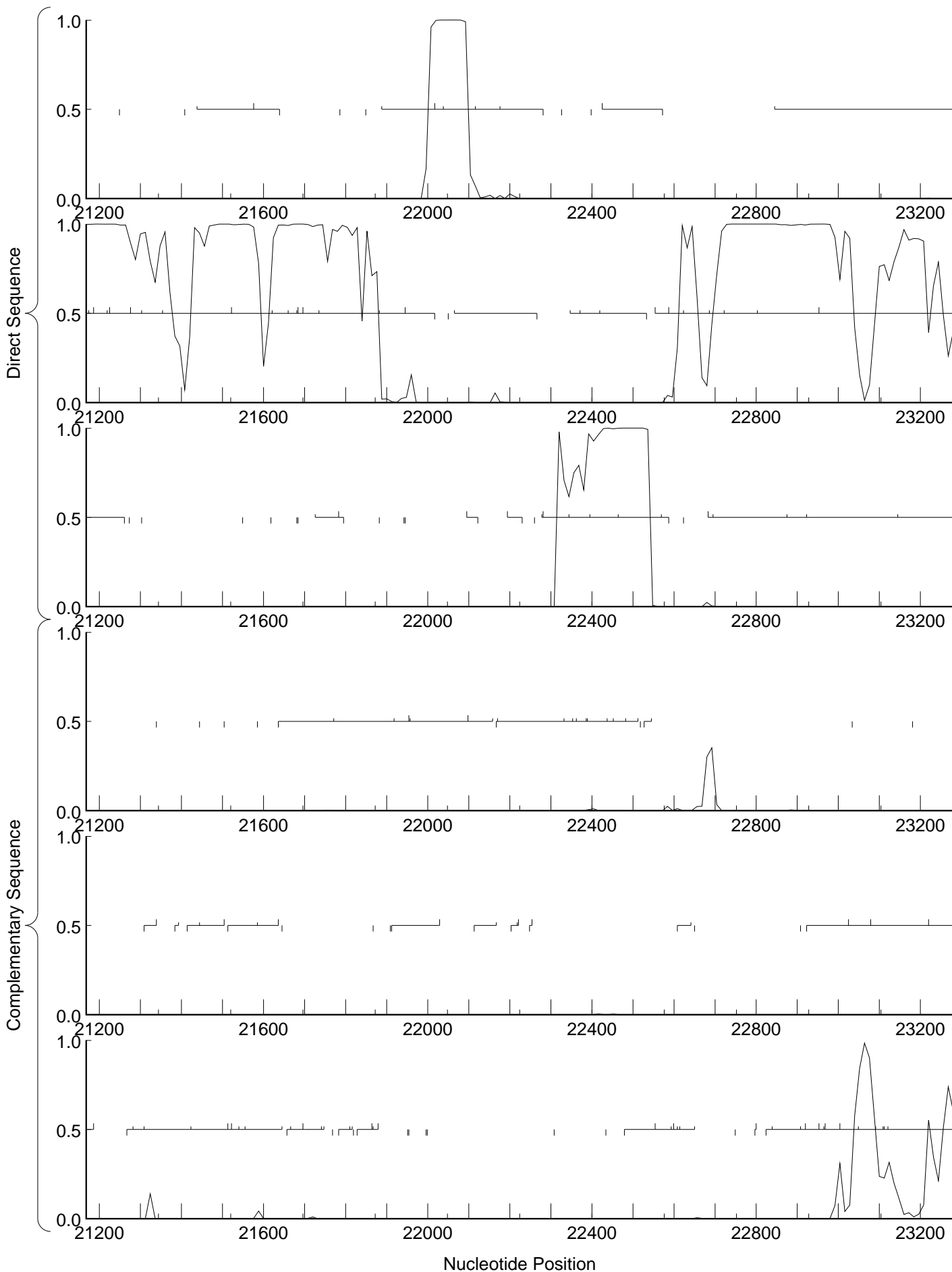












complete sequence, 47863 bp including 11-base 3' overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 13/24

