

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Yeet412 complete sequence, 46684 bp including 11-base 3' overhang (TACCAGGGG
Analysis Date : 2/27/21 at 7:39:24
Pages : 24
Sequence Length : 46684 bp
GC Content : 66.77%

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,

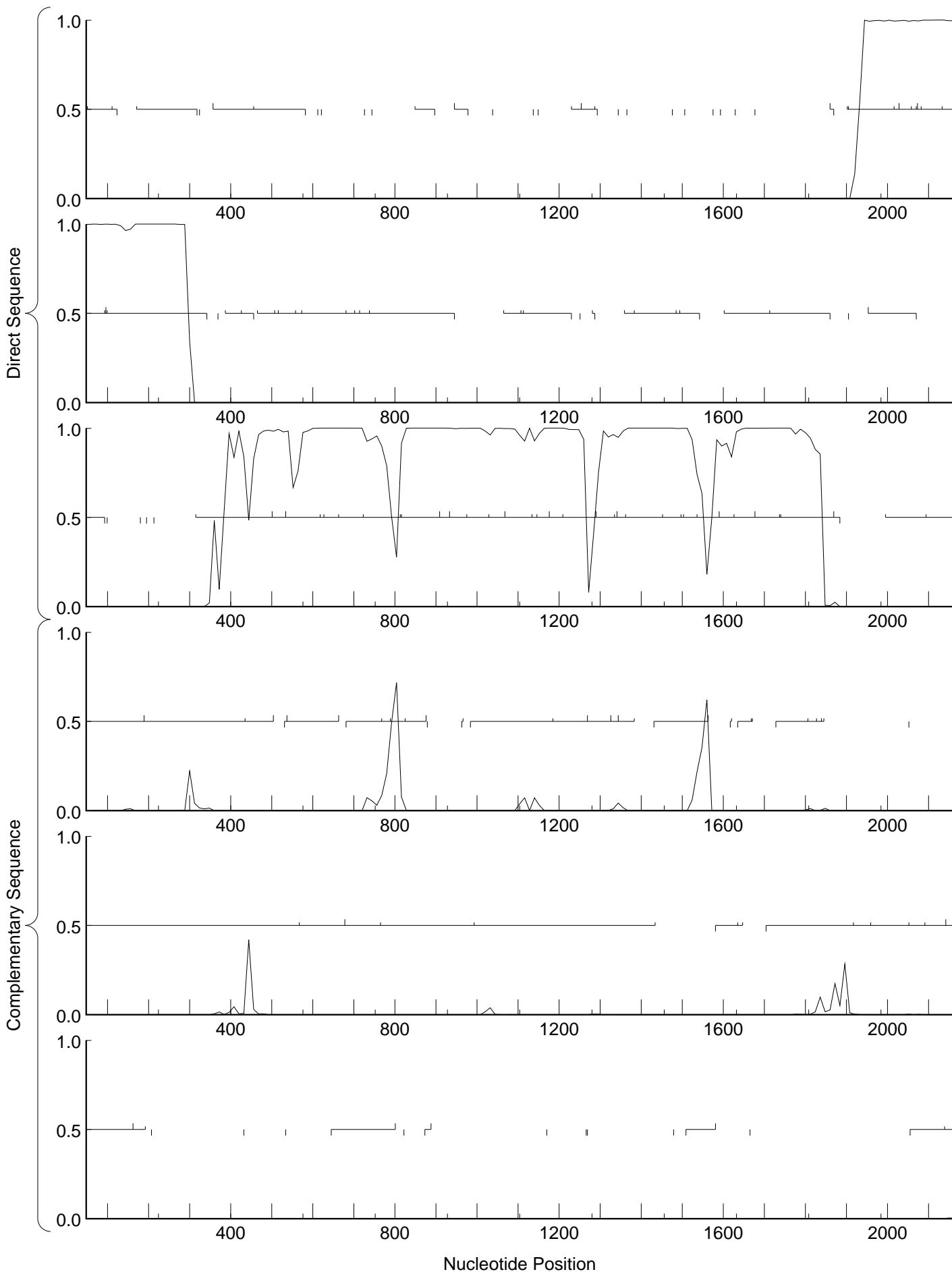
MATRIX INFORMATION

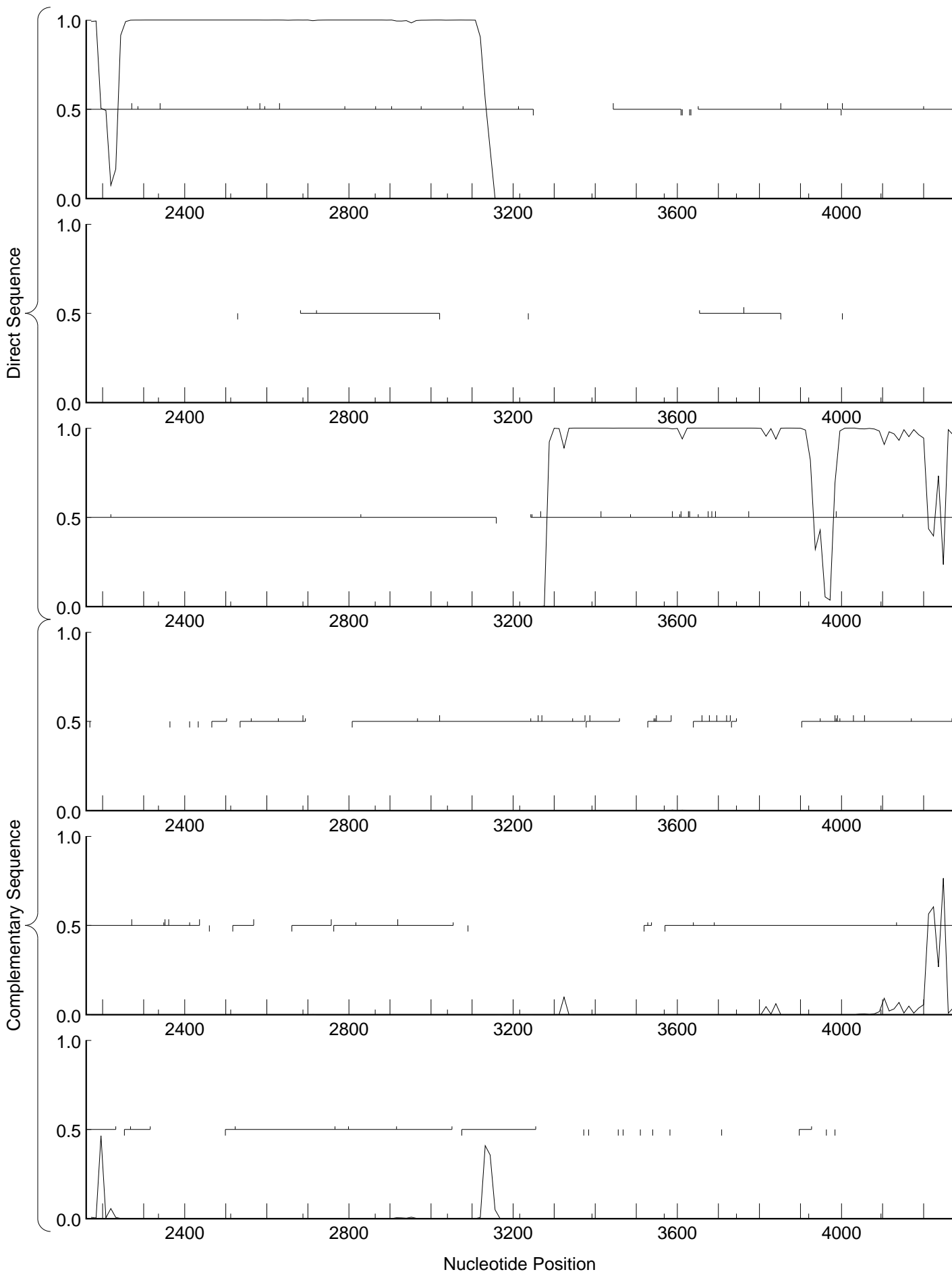
Matrix : -
Author : -
Order : 4

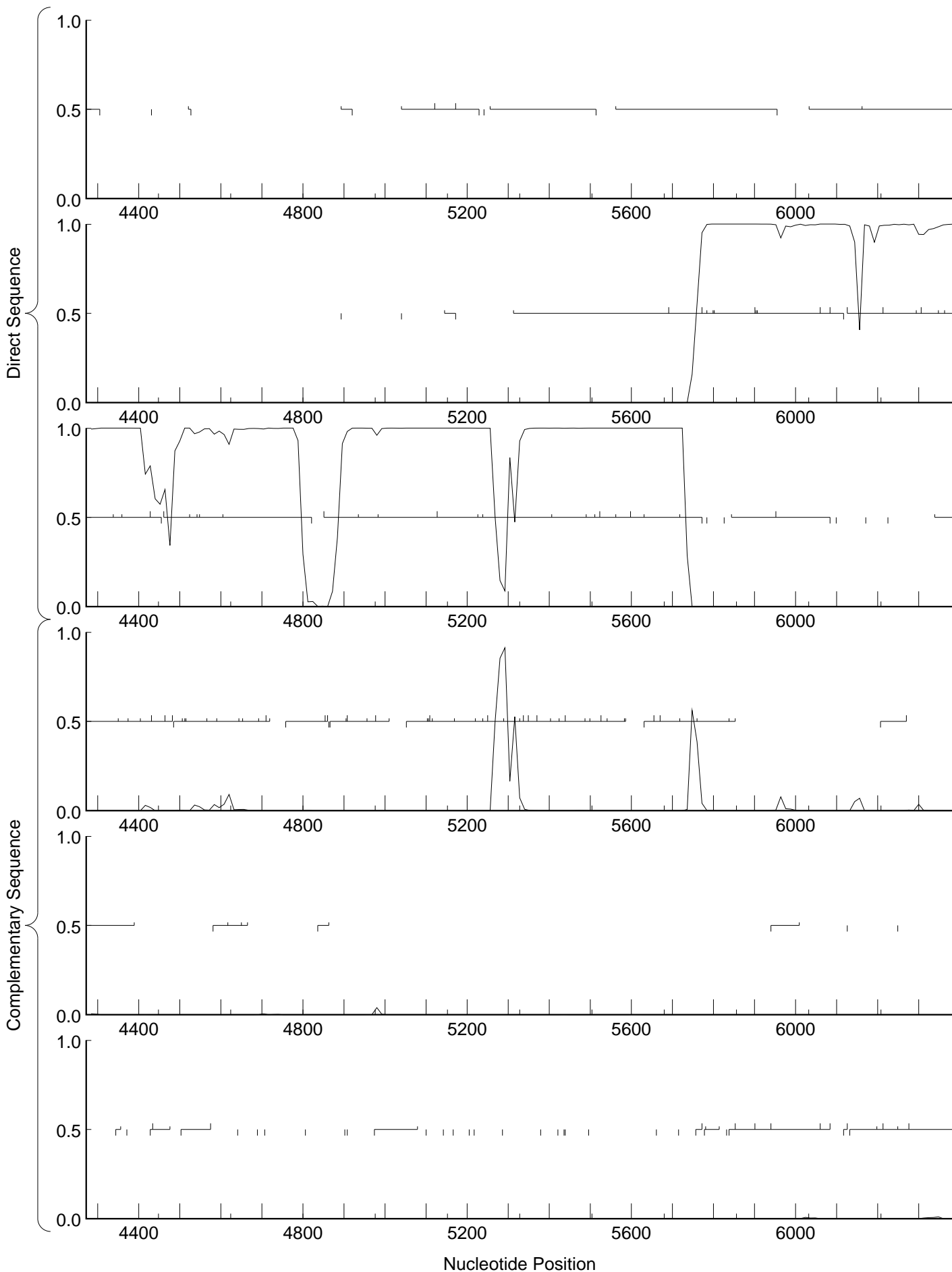
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

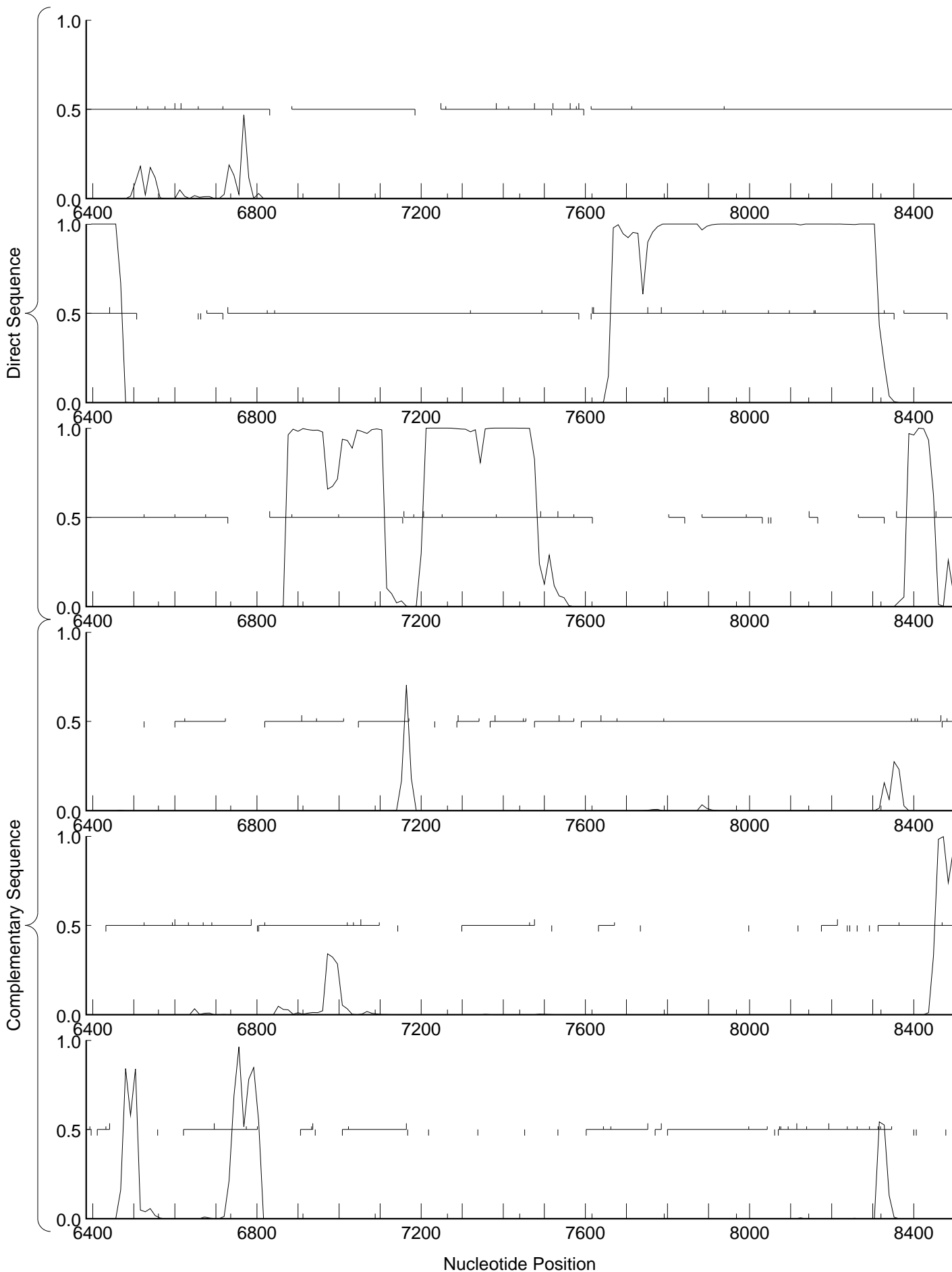
Matrix notes & comments

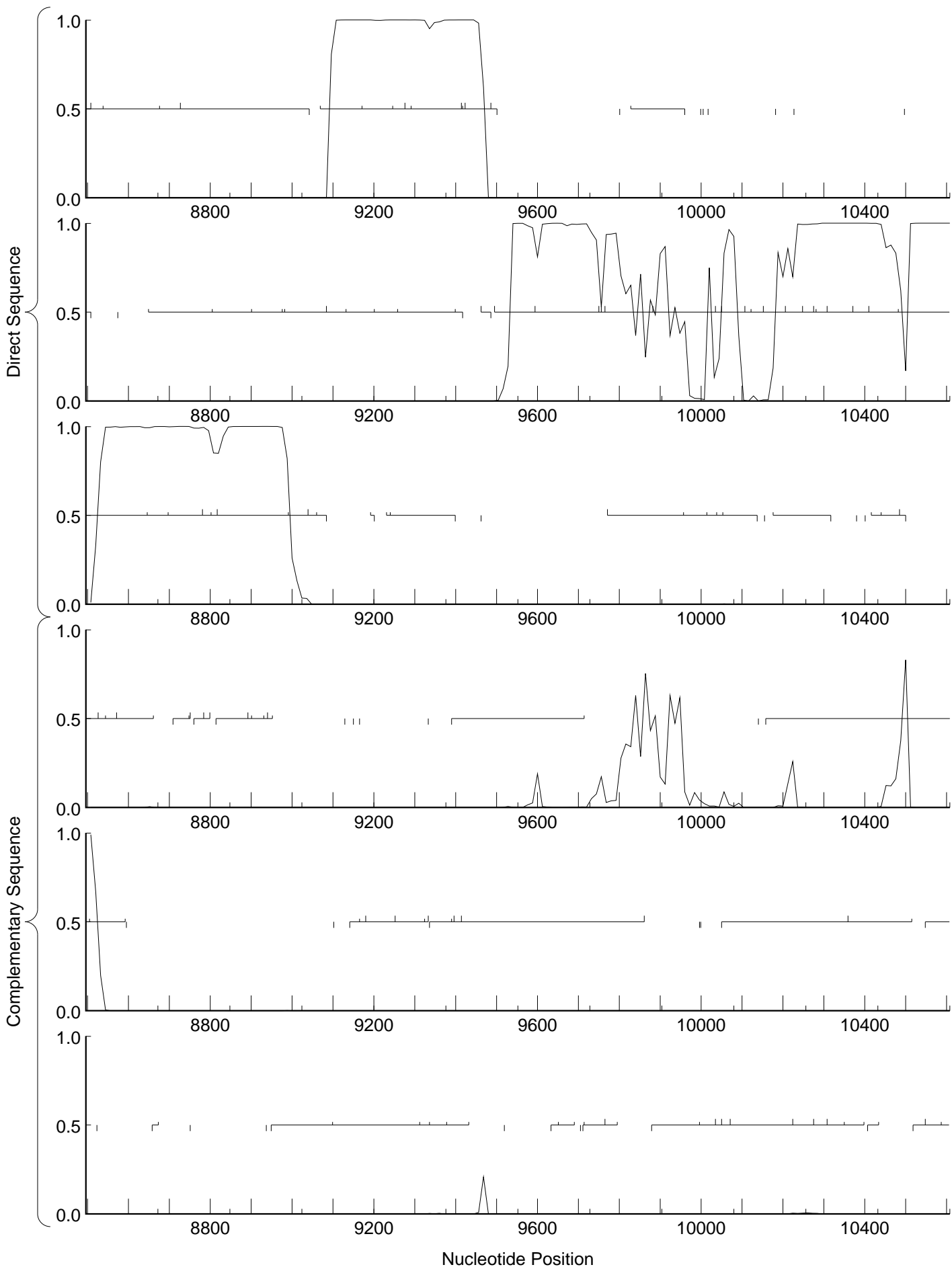
Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:36:14 2018

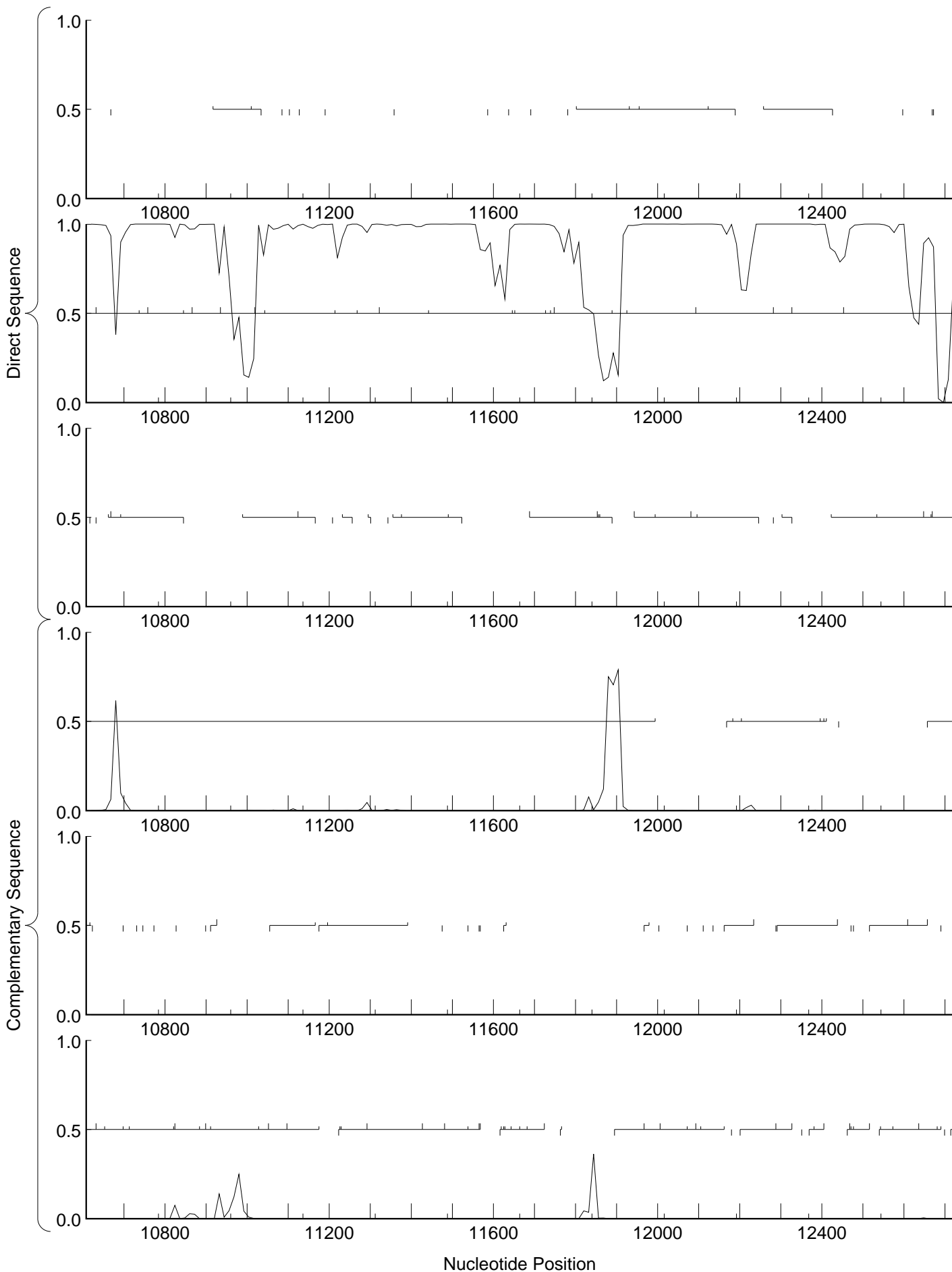


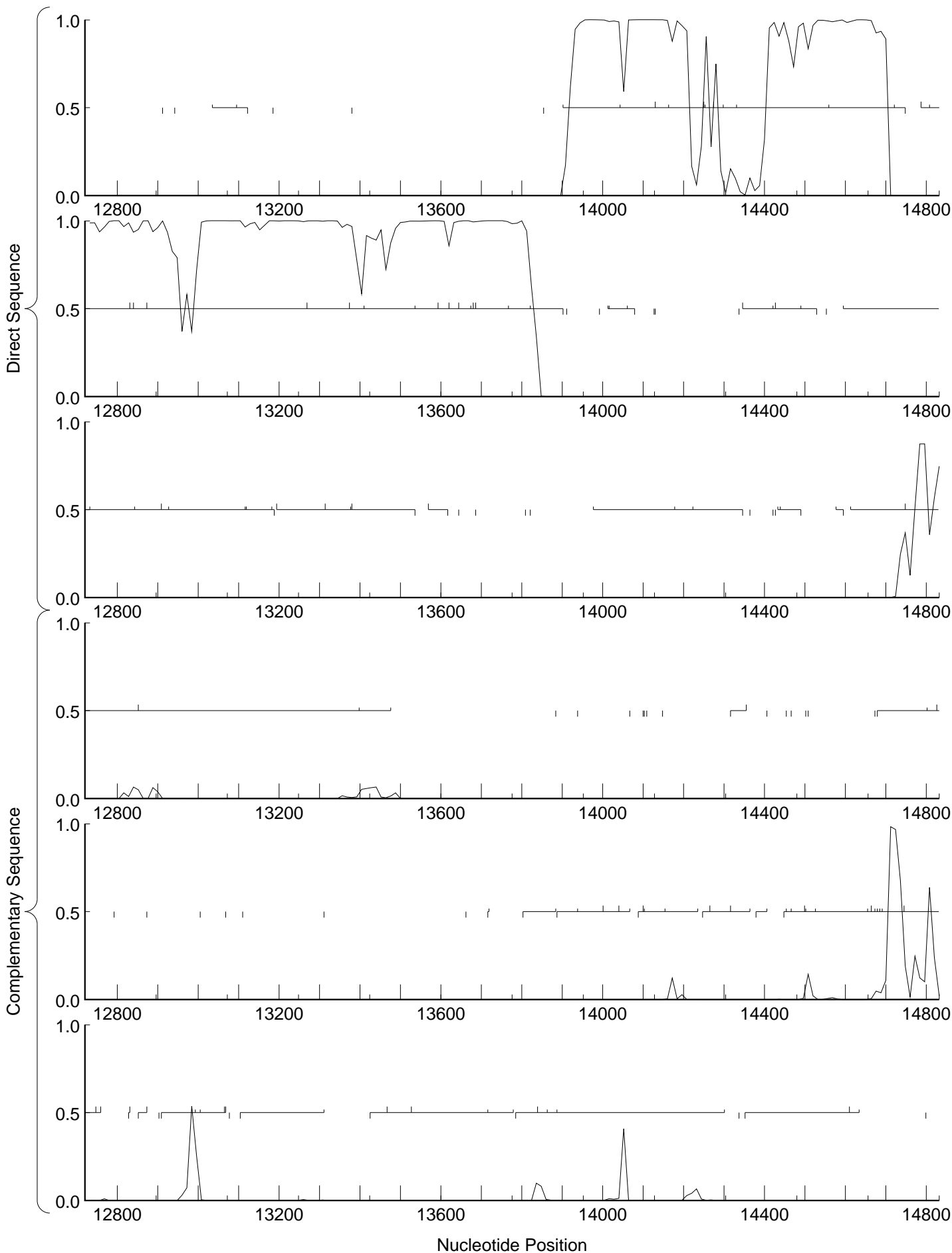


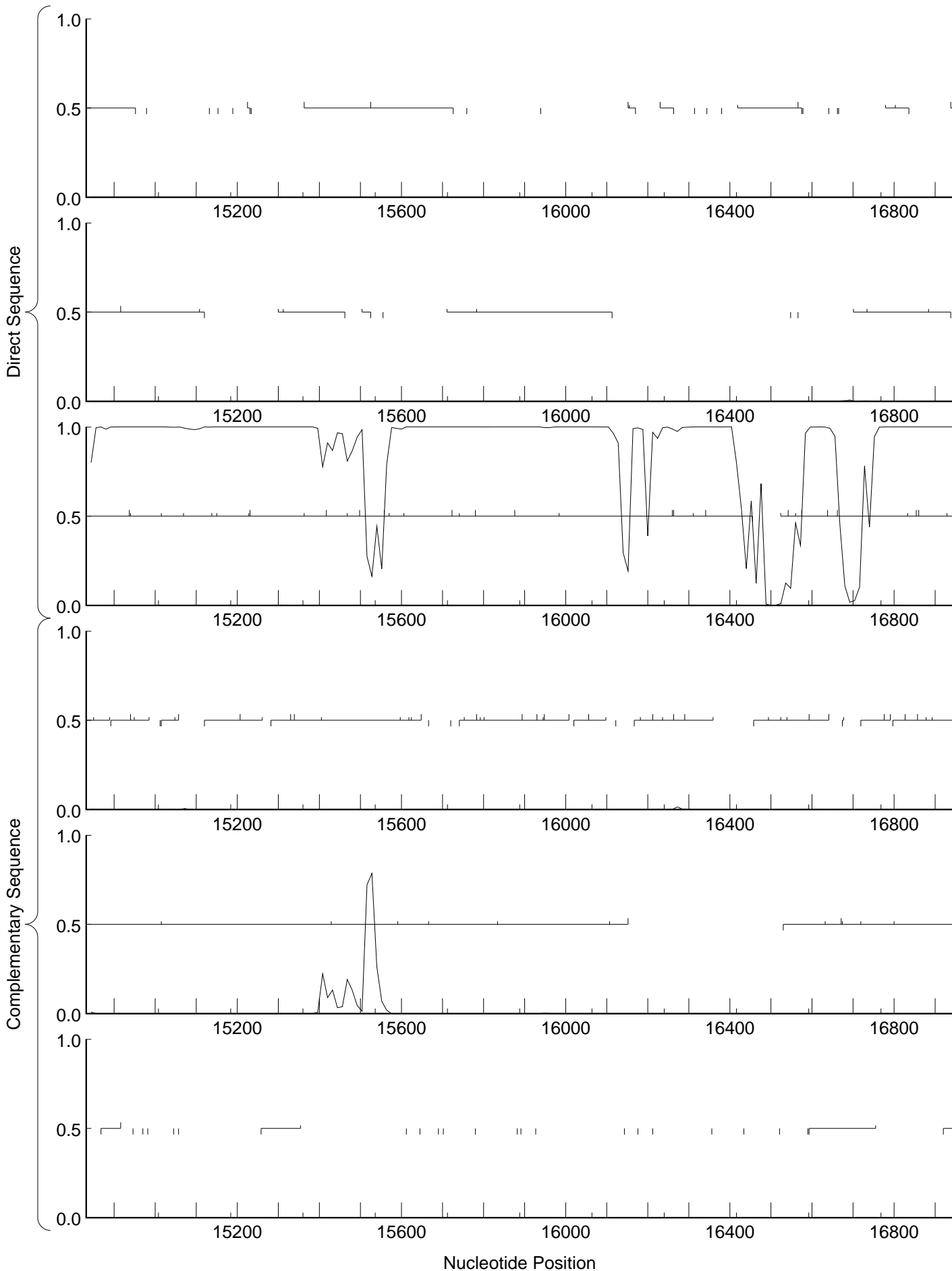


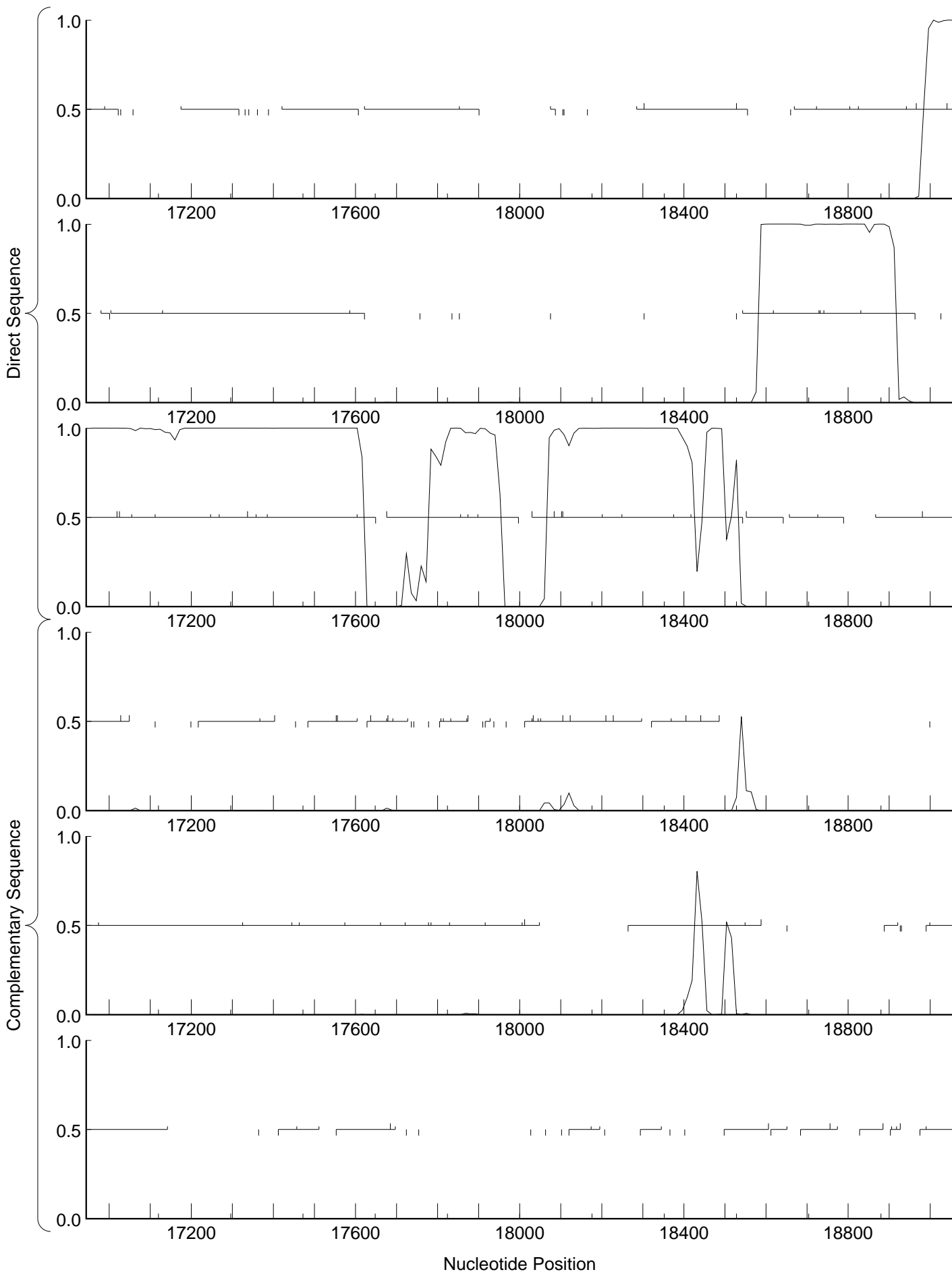


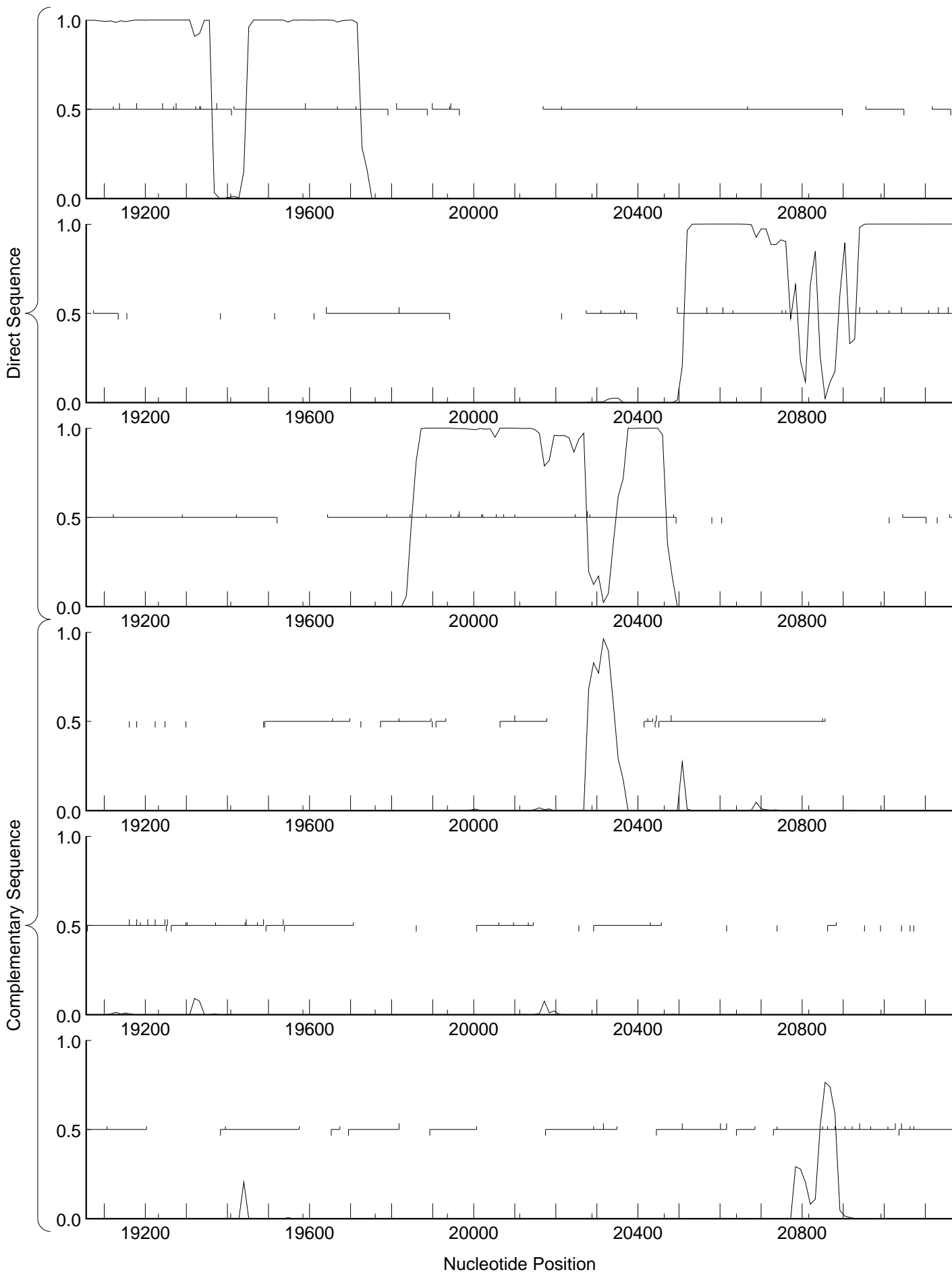


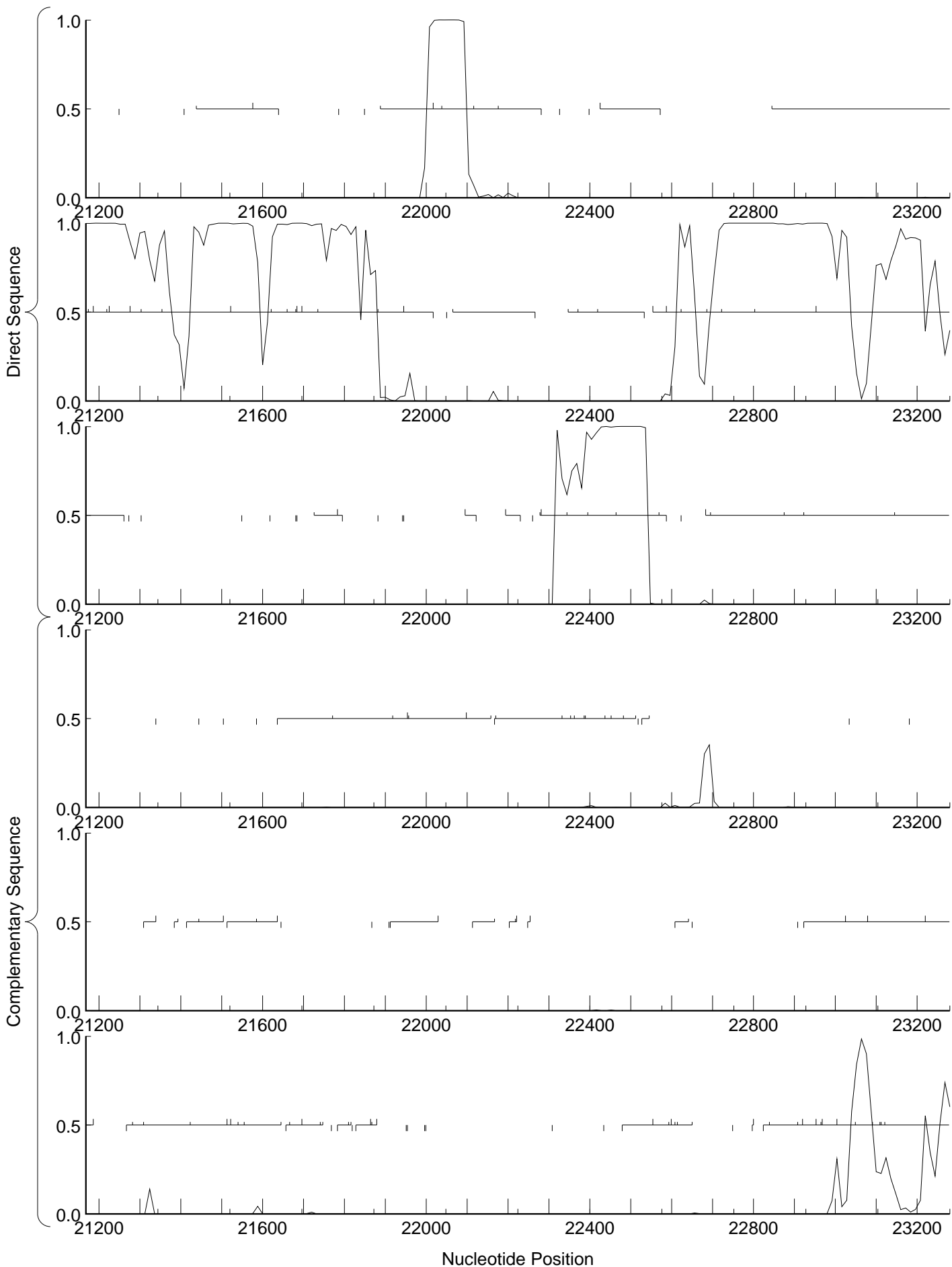




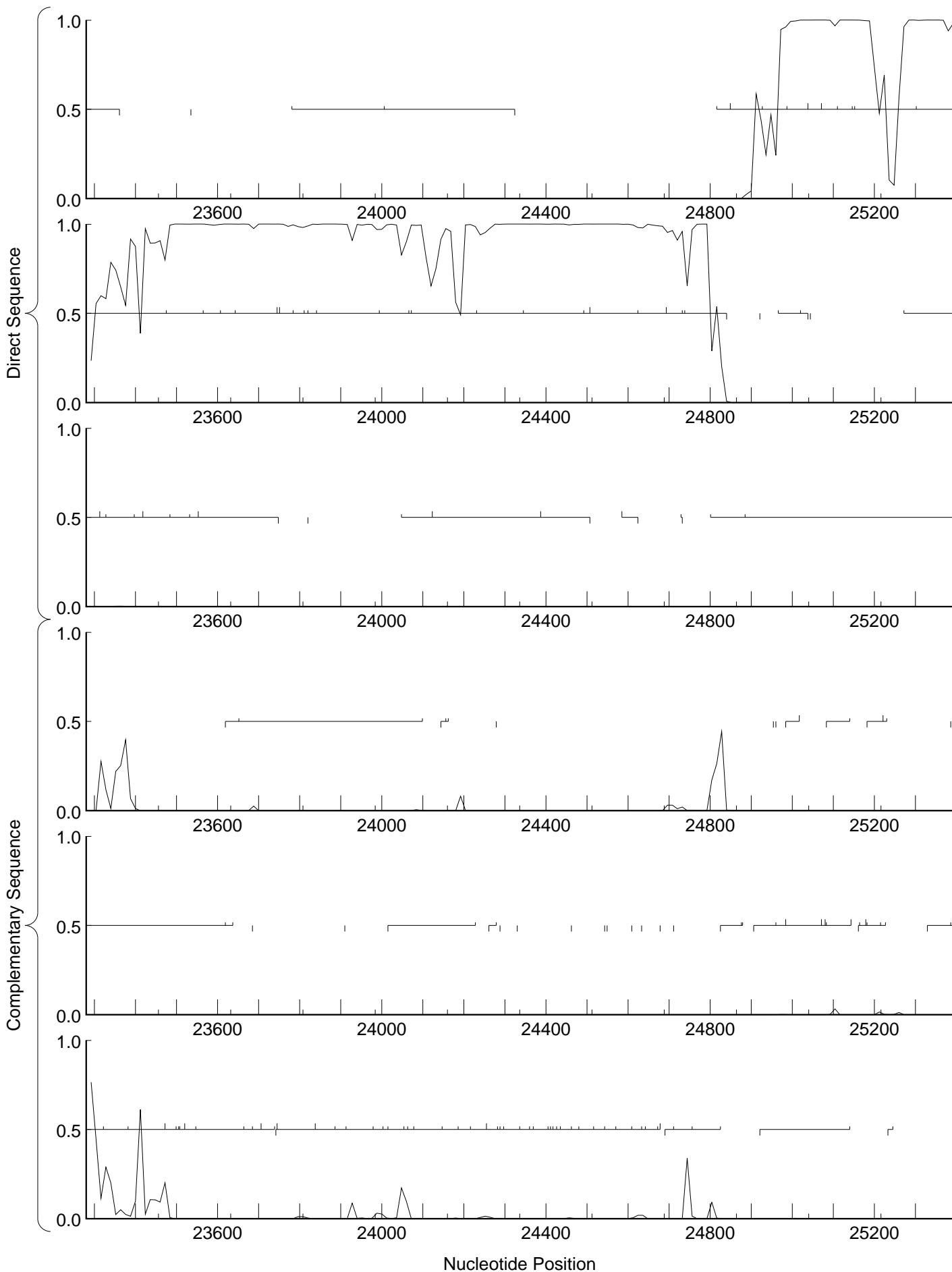


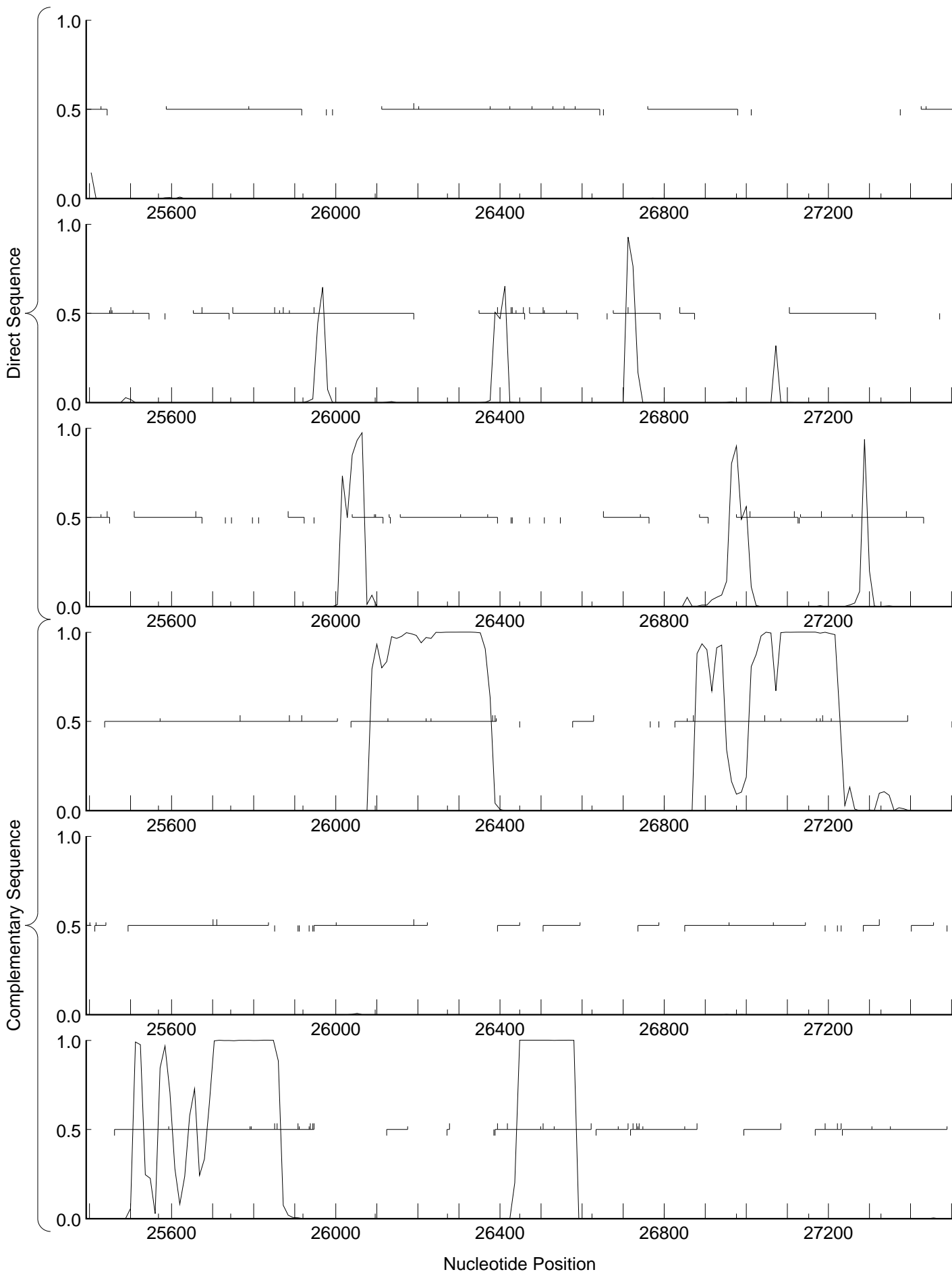


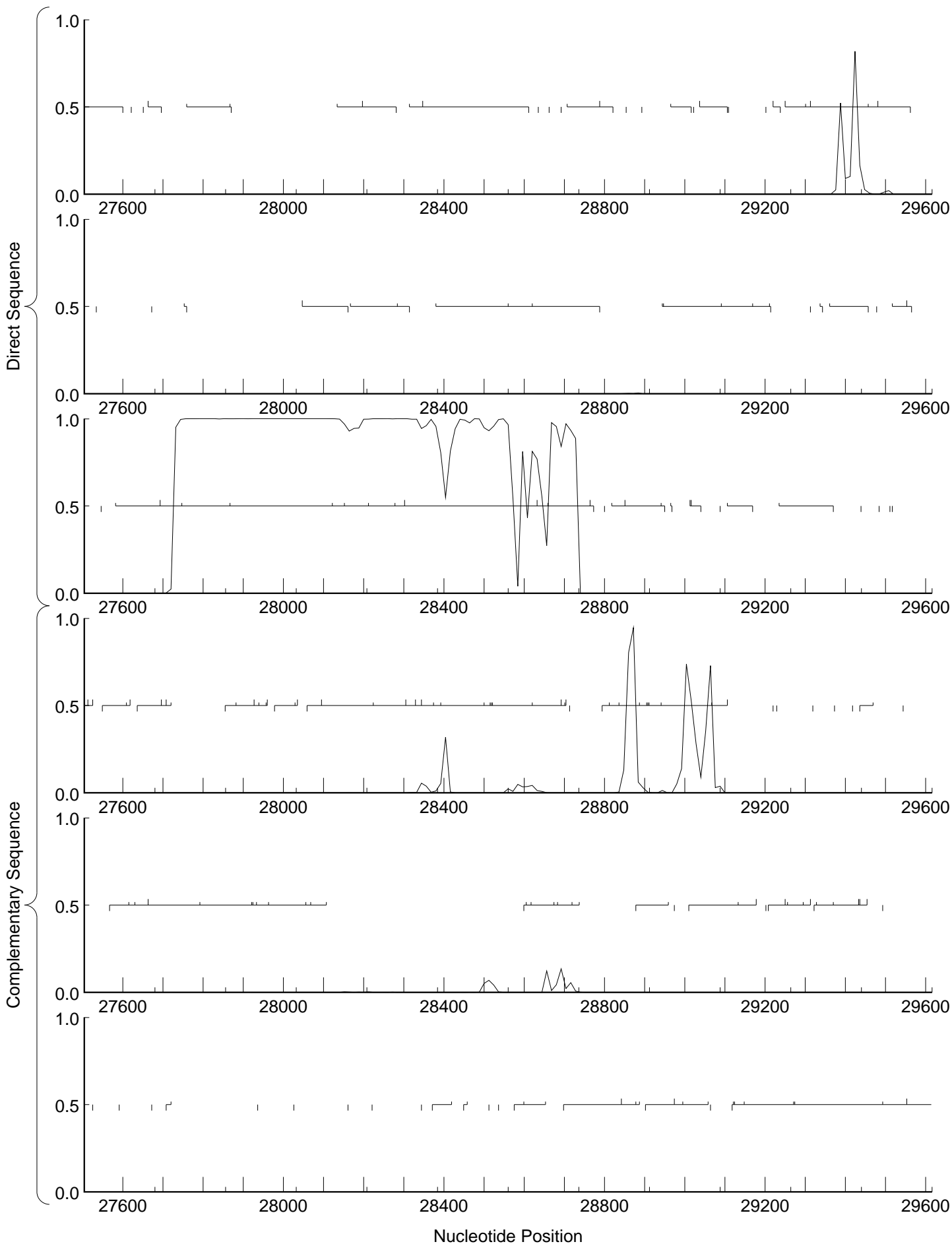


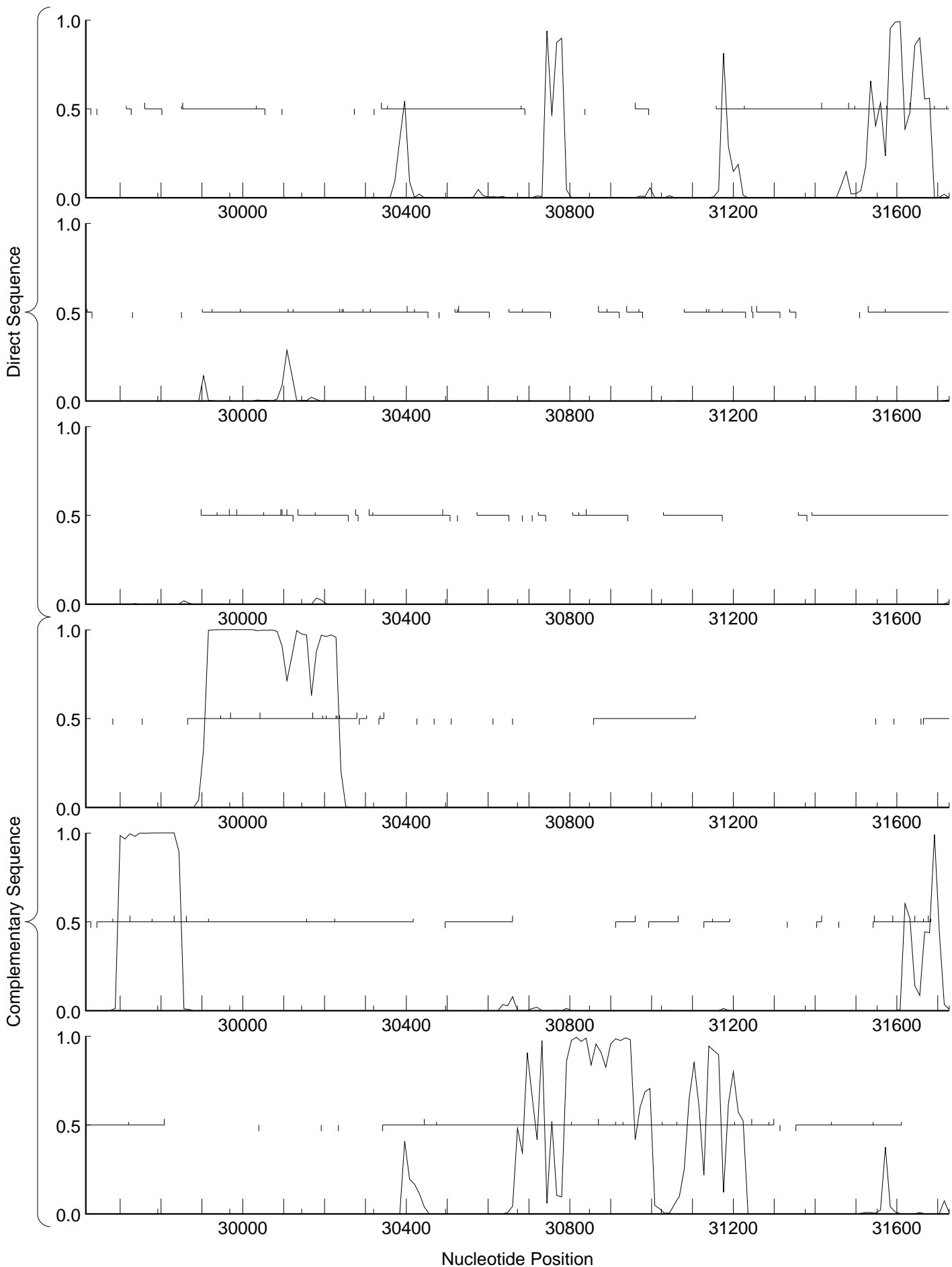


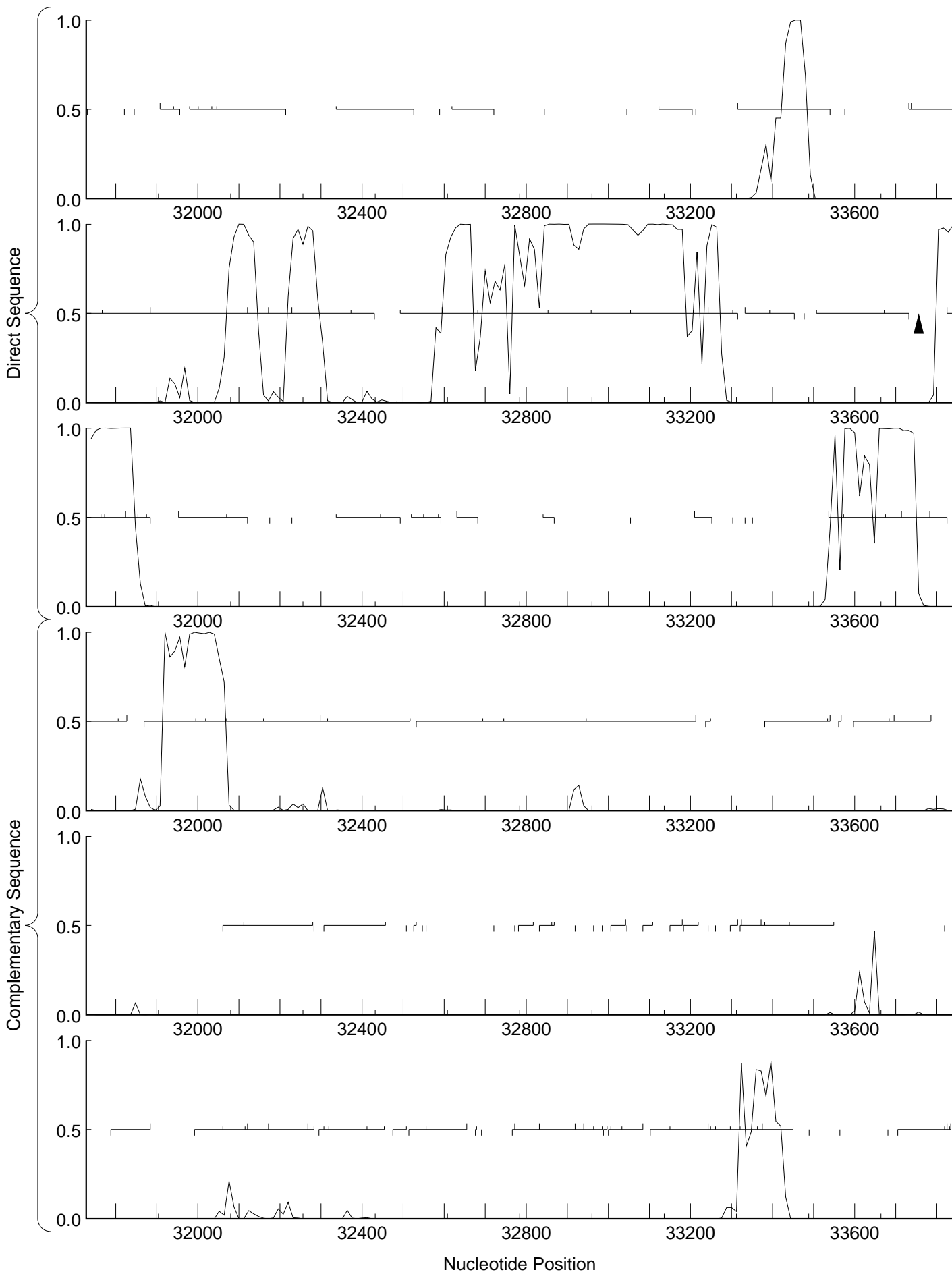
complete sequence, 46684 bp including 11-base 3' overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 13/24

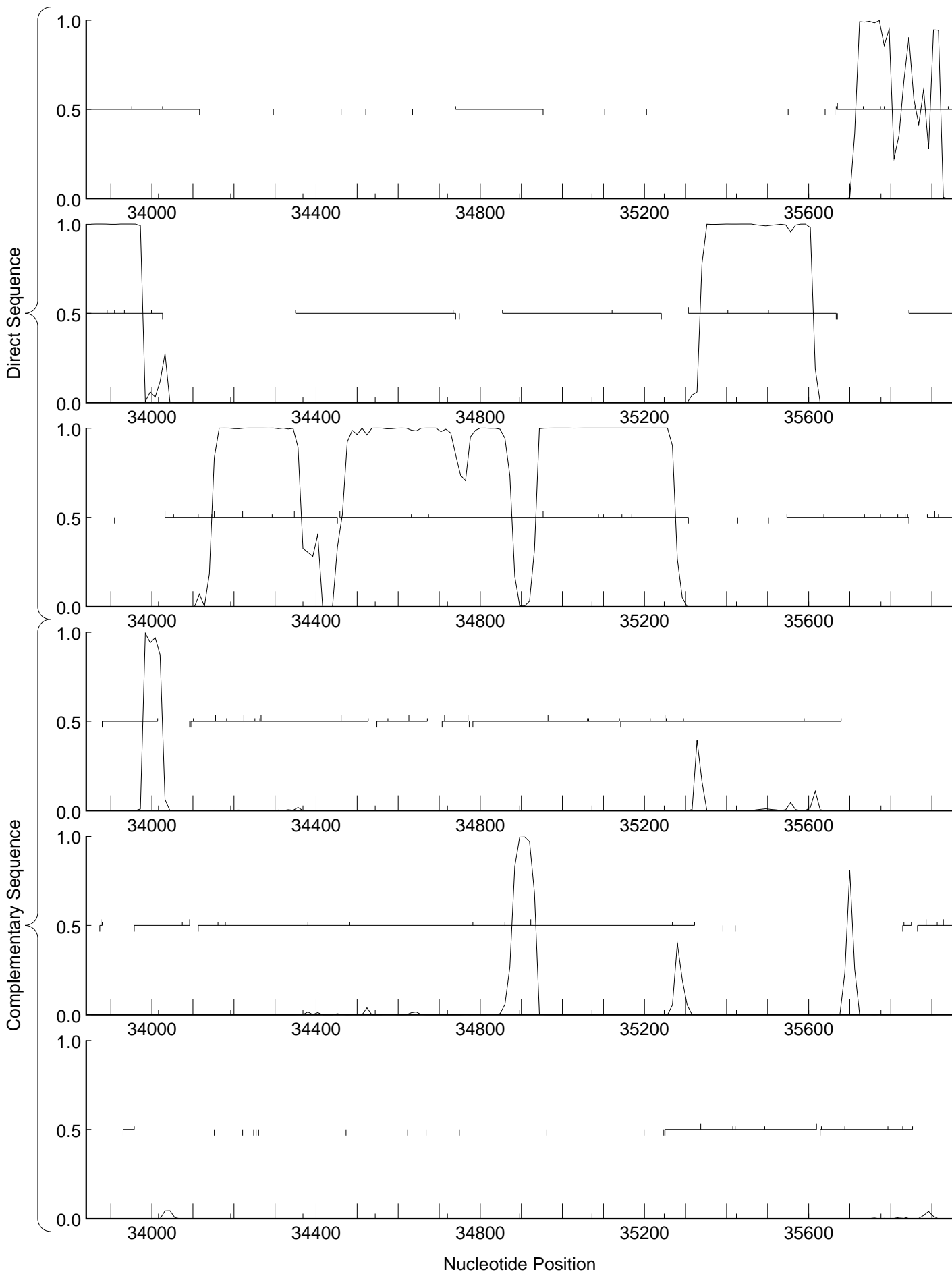


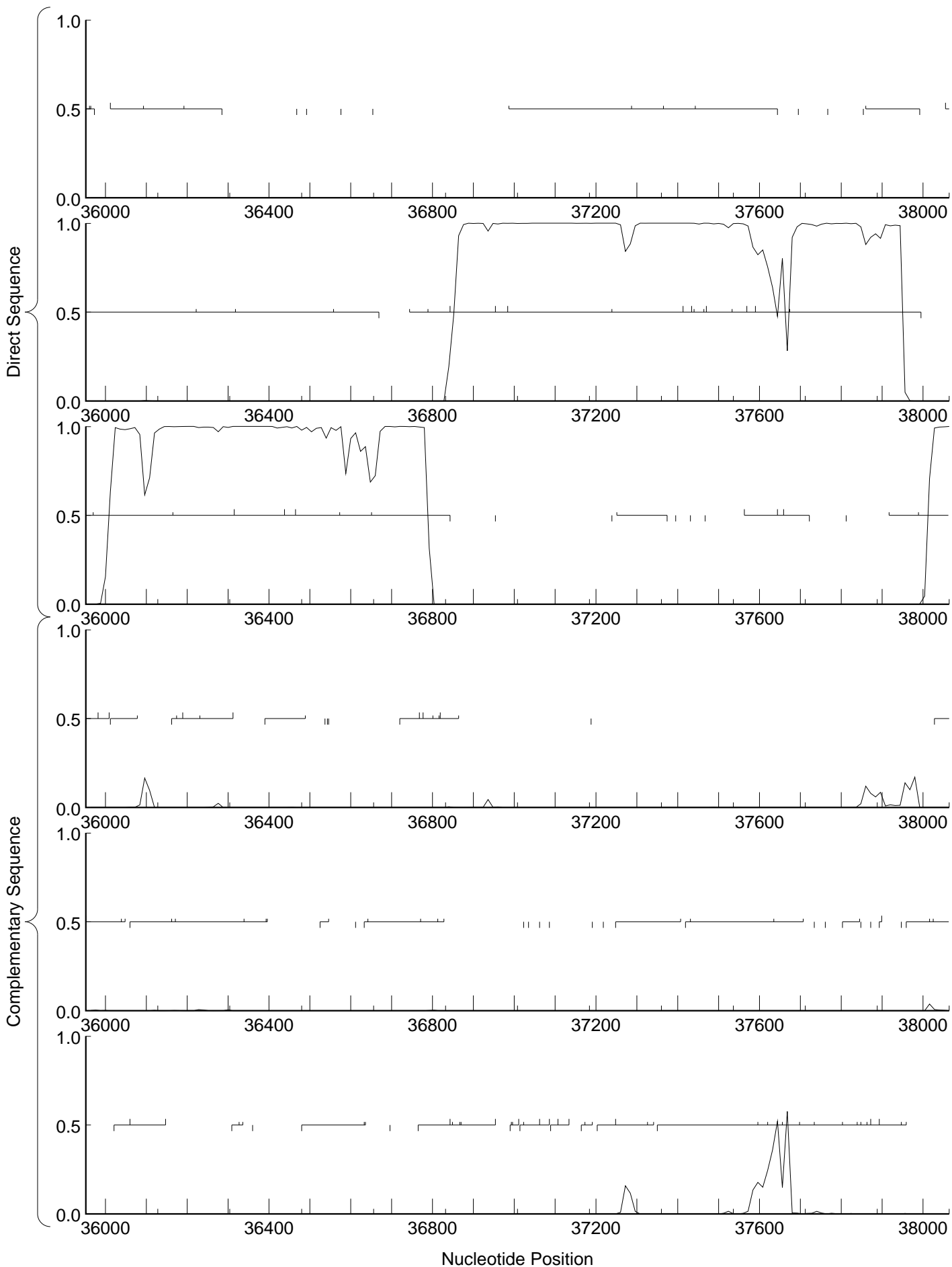


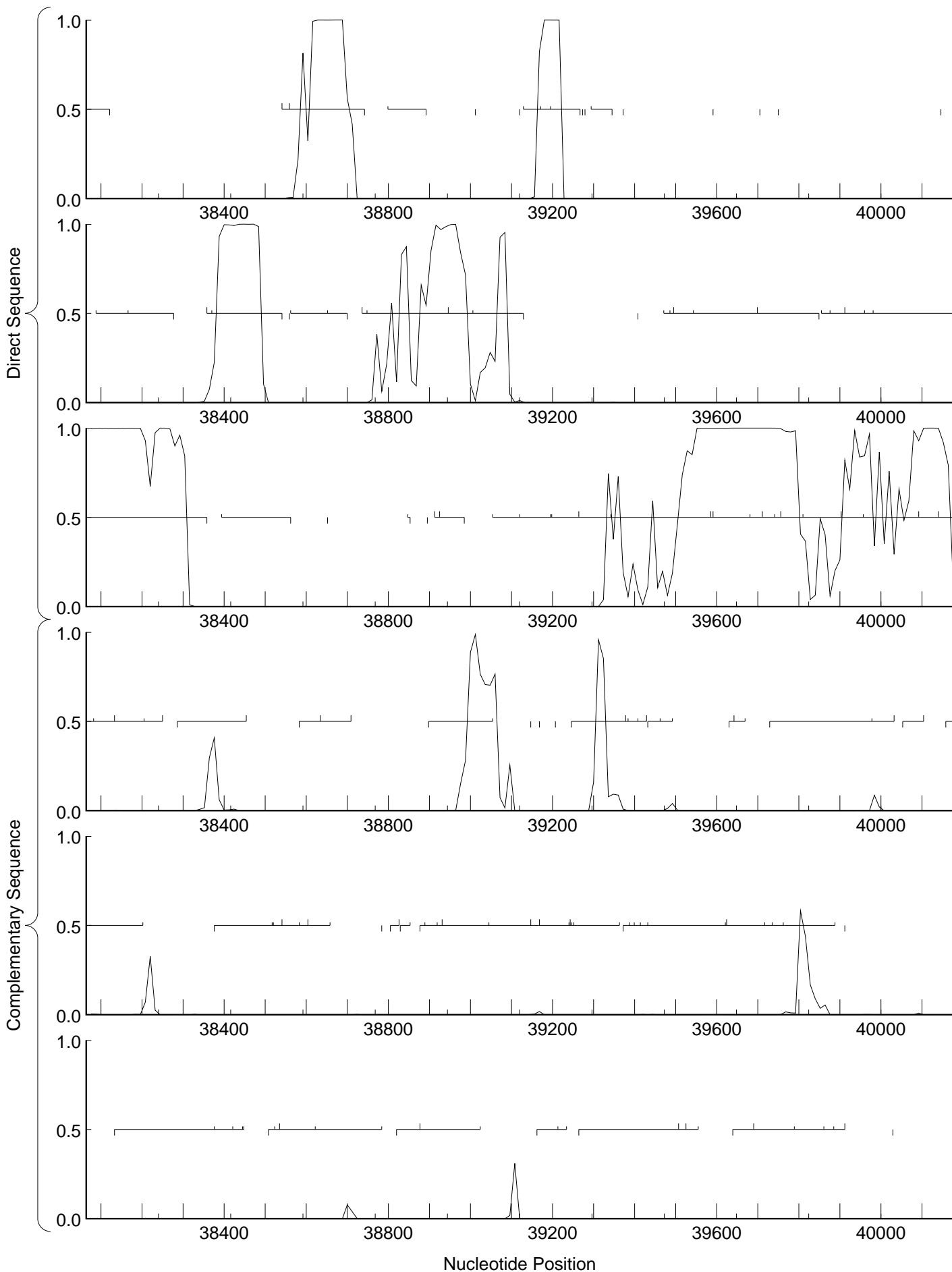


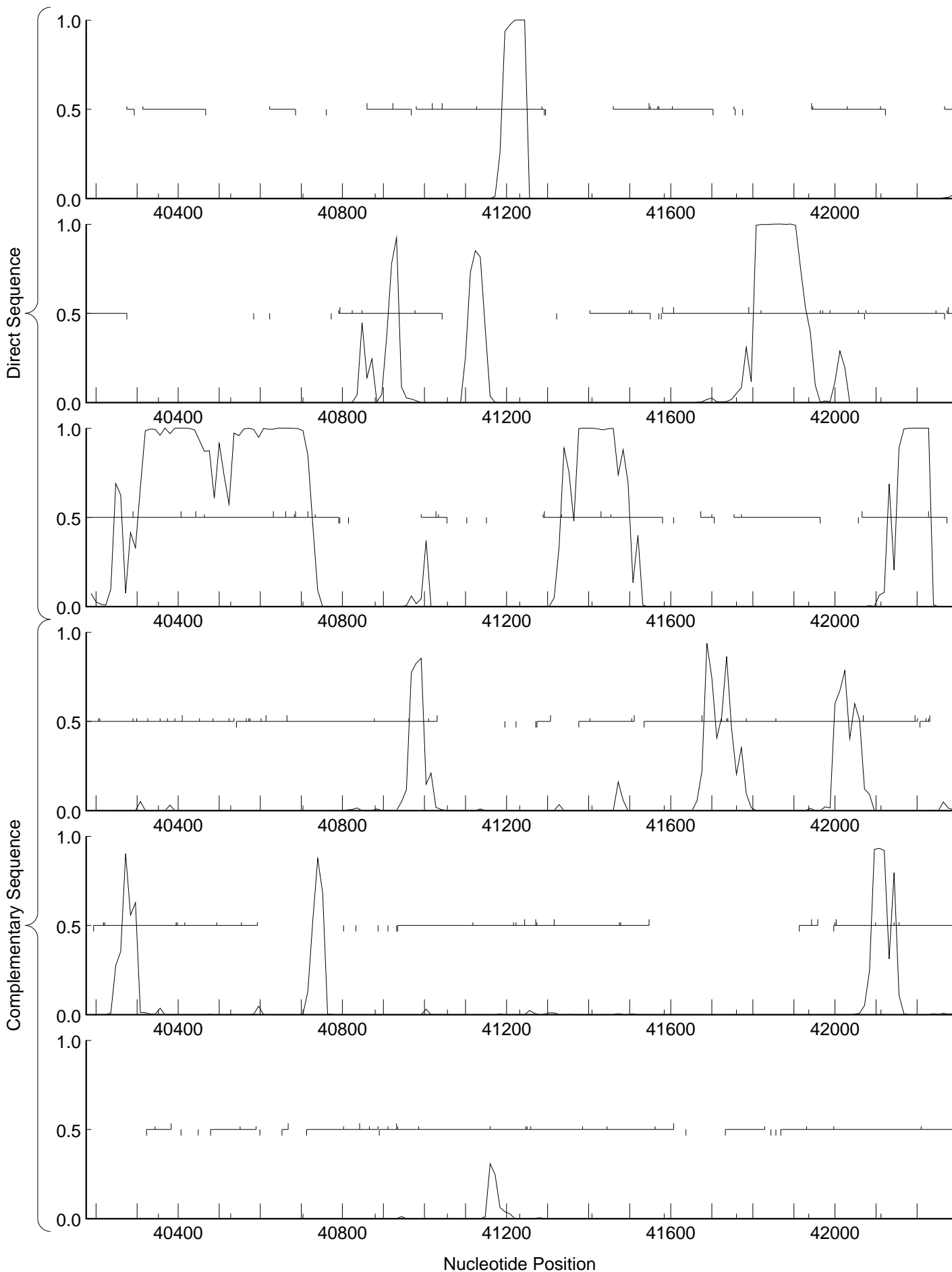


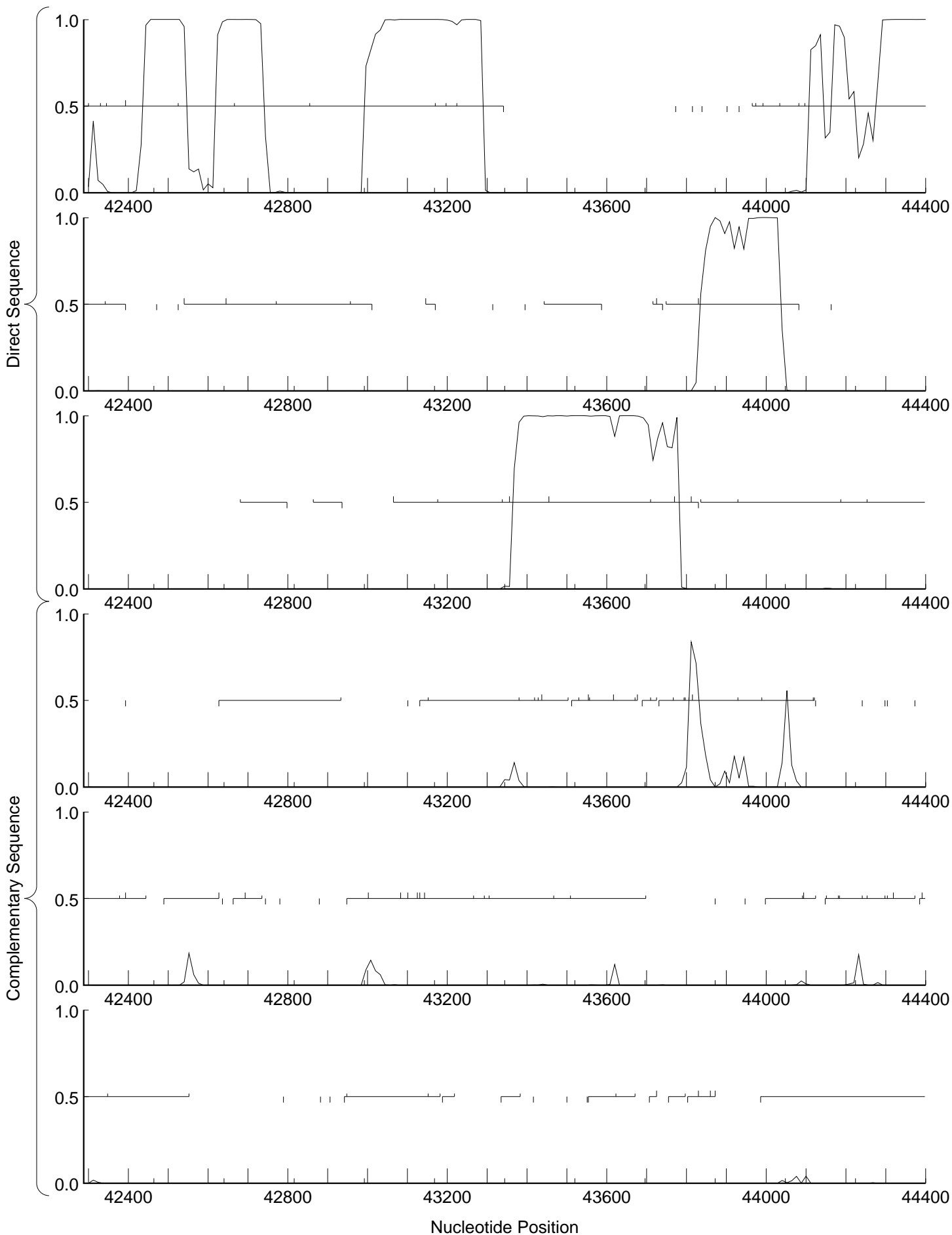












Complete sequence, 46684 bp including 11-base 3' overhang (TACCAGGGGGA), Cluster CZ2, Order 4, Window 96, Step 12, 23/24

