

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Mycobacterium phage Soap141 complete sequence, 76871 bp including 10-base 3' overhang (TCGA)
Analysis Date : 3/23/25 at 12:59:08
Pages : 38
Sequence Length : 76871 bp
GC Content : 58.96%

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,

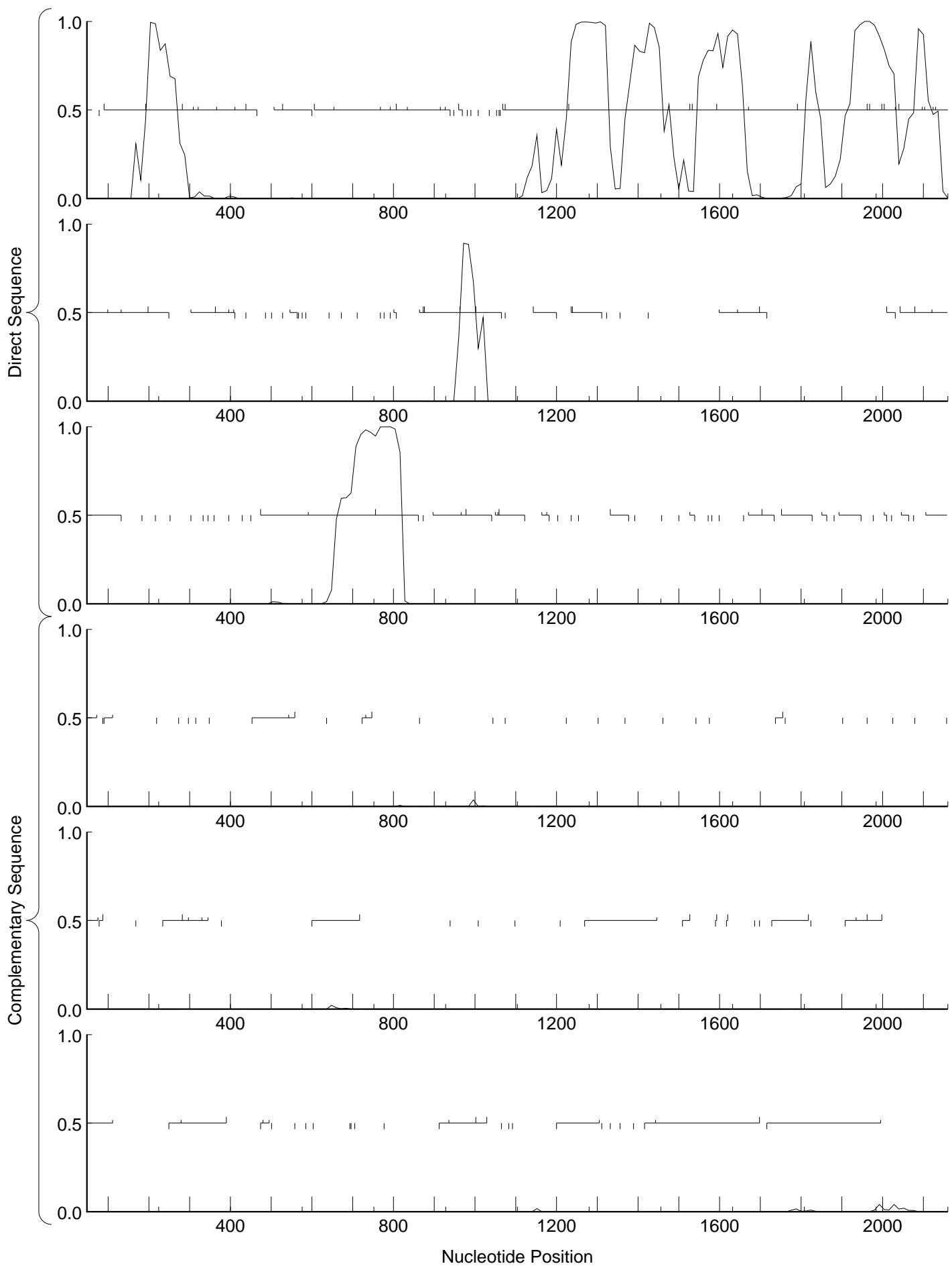
MATRIX INFORMATION

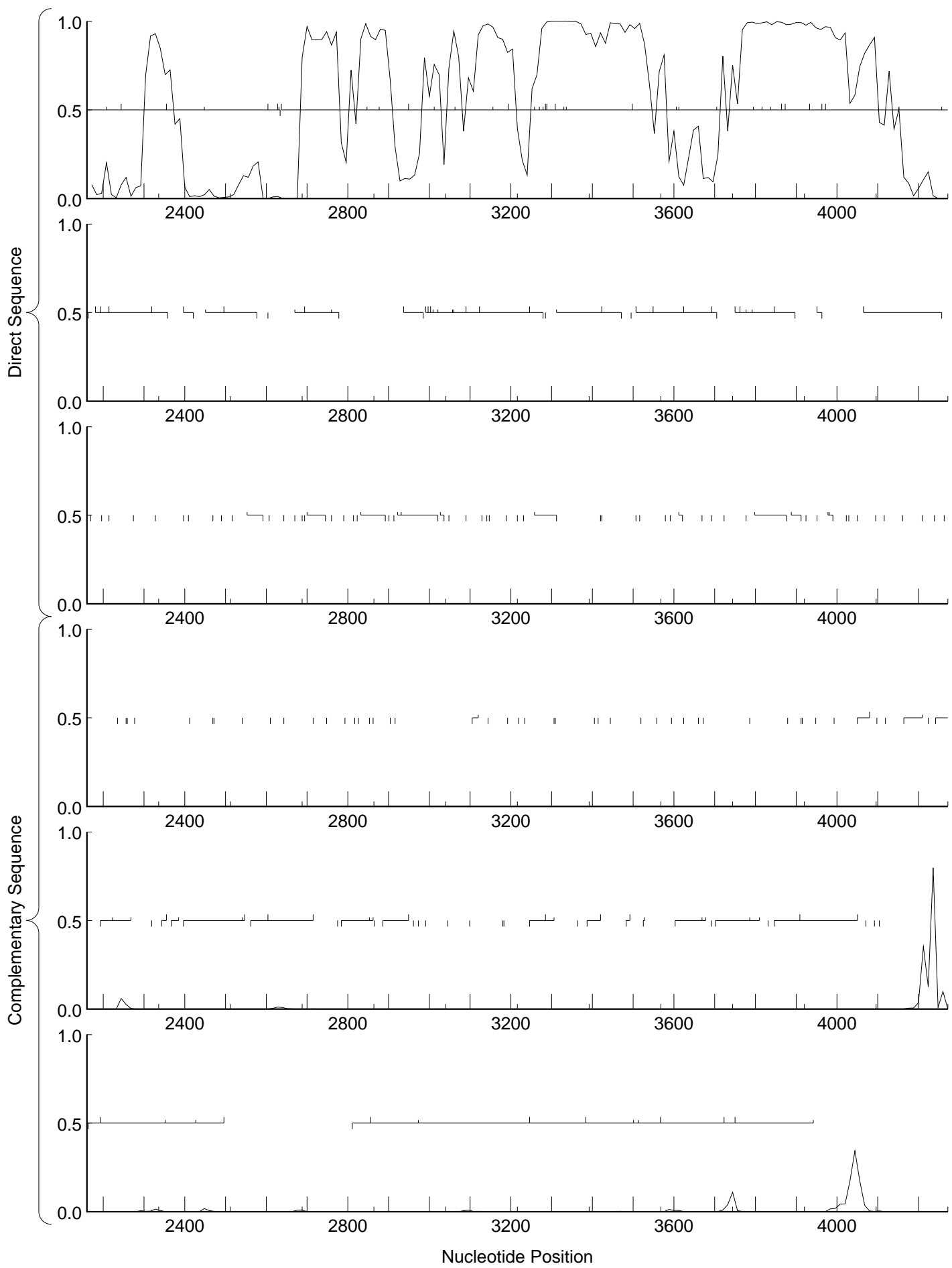
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

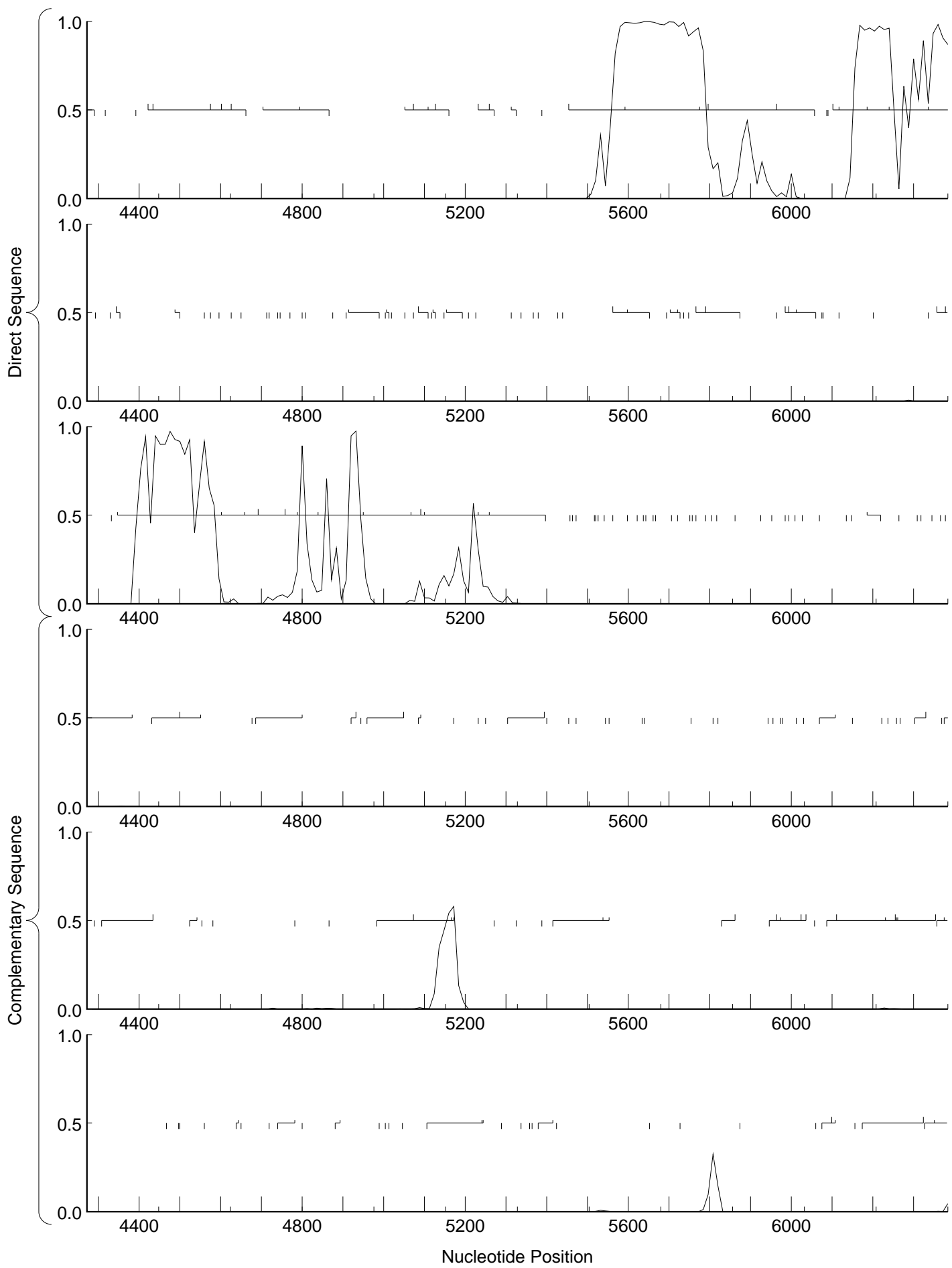
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

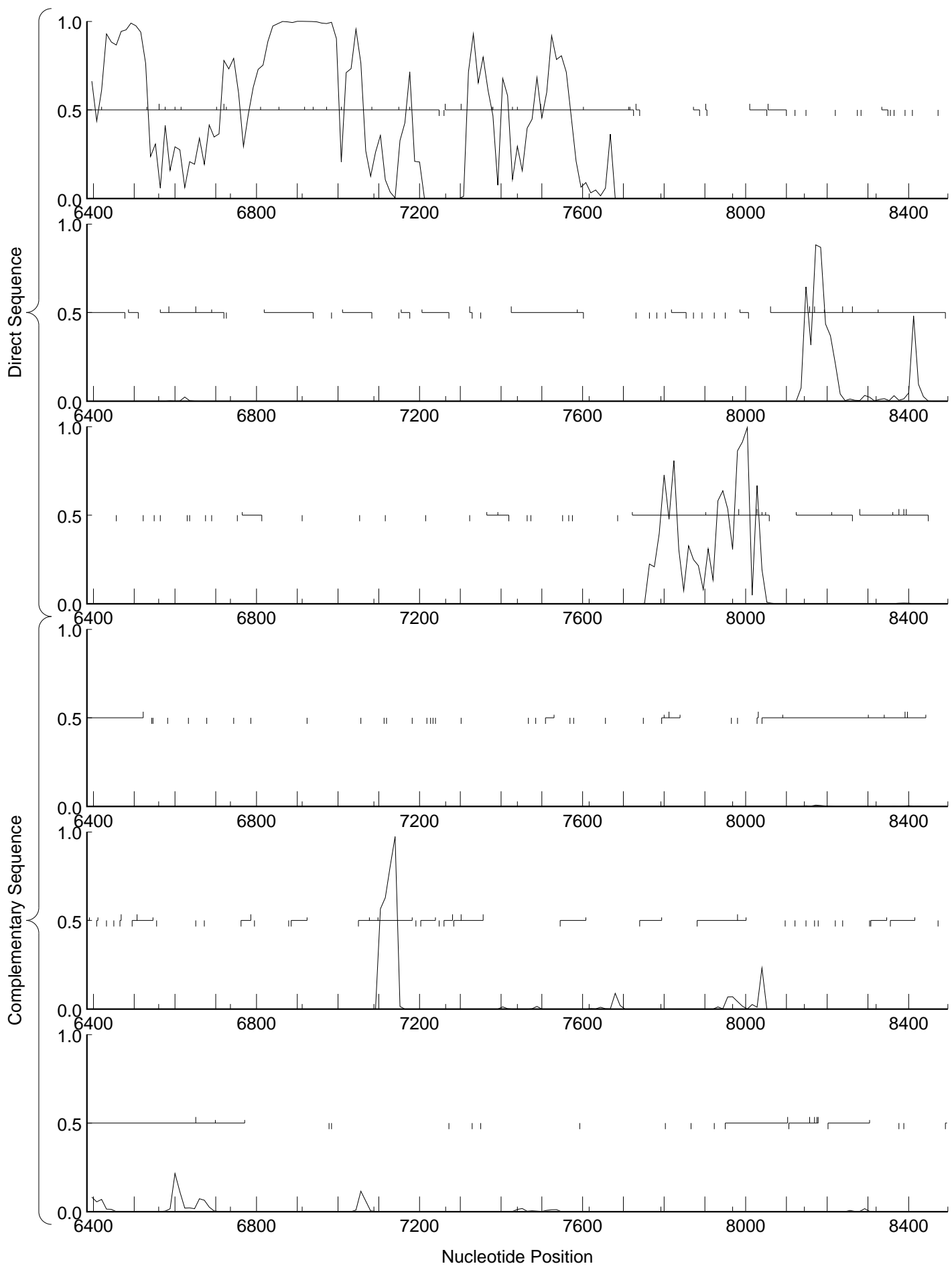
Matrix notes & comments

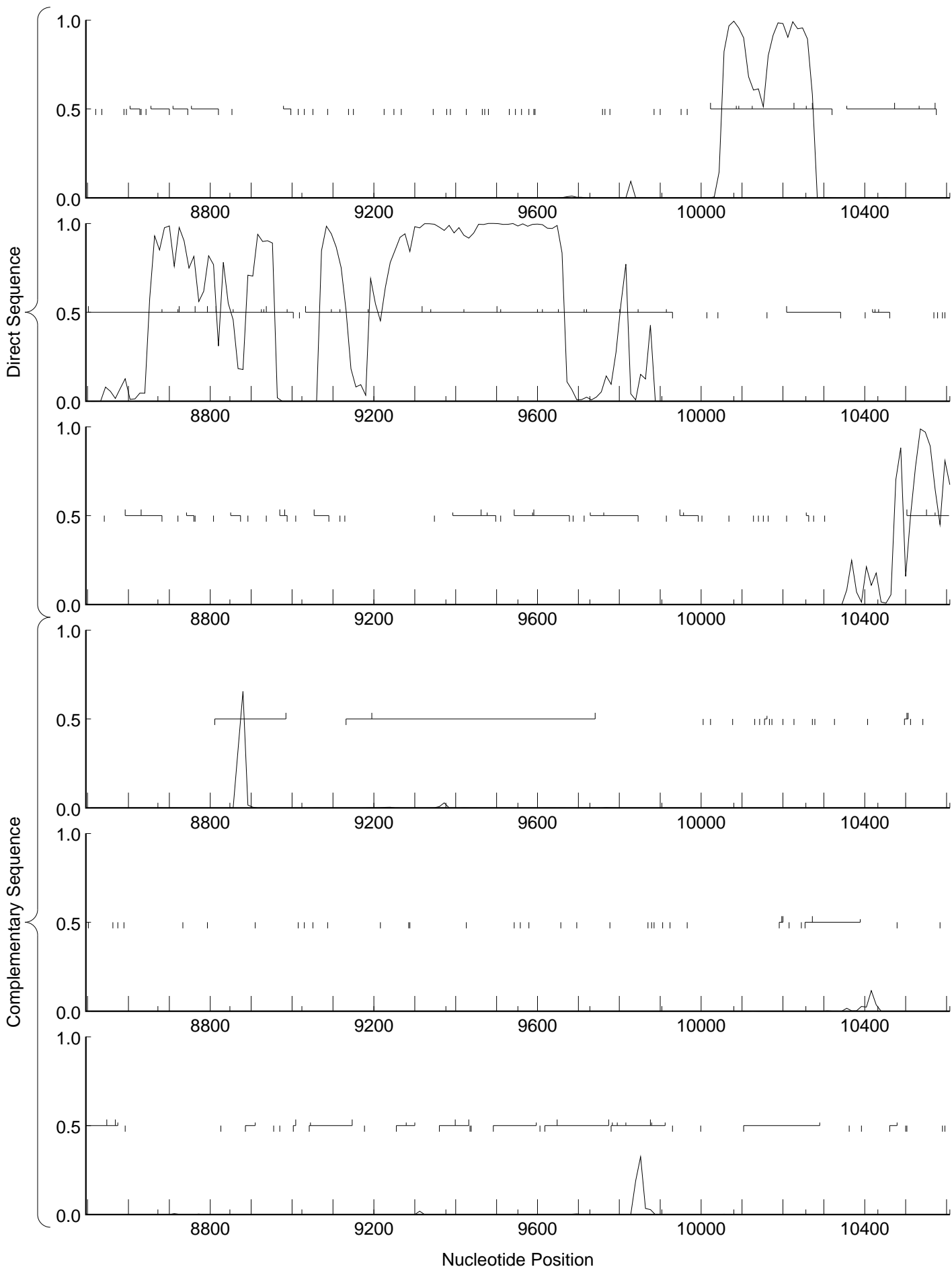
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

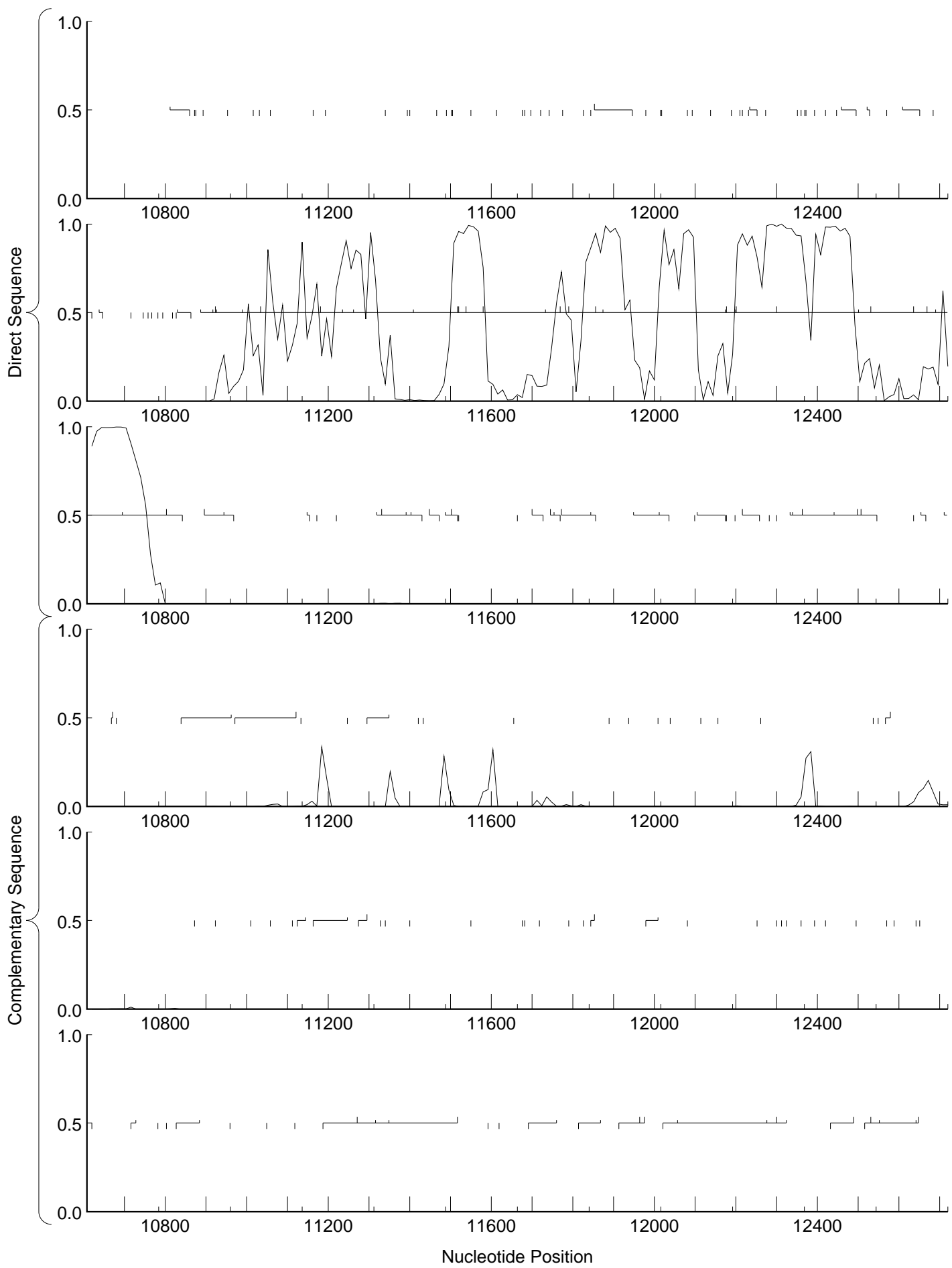


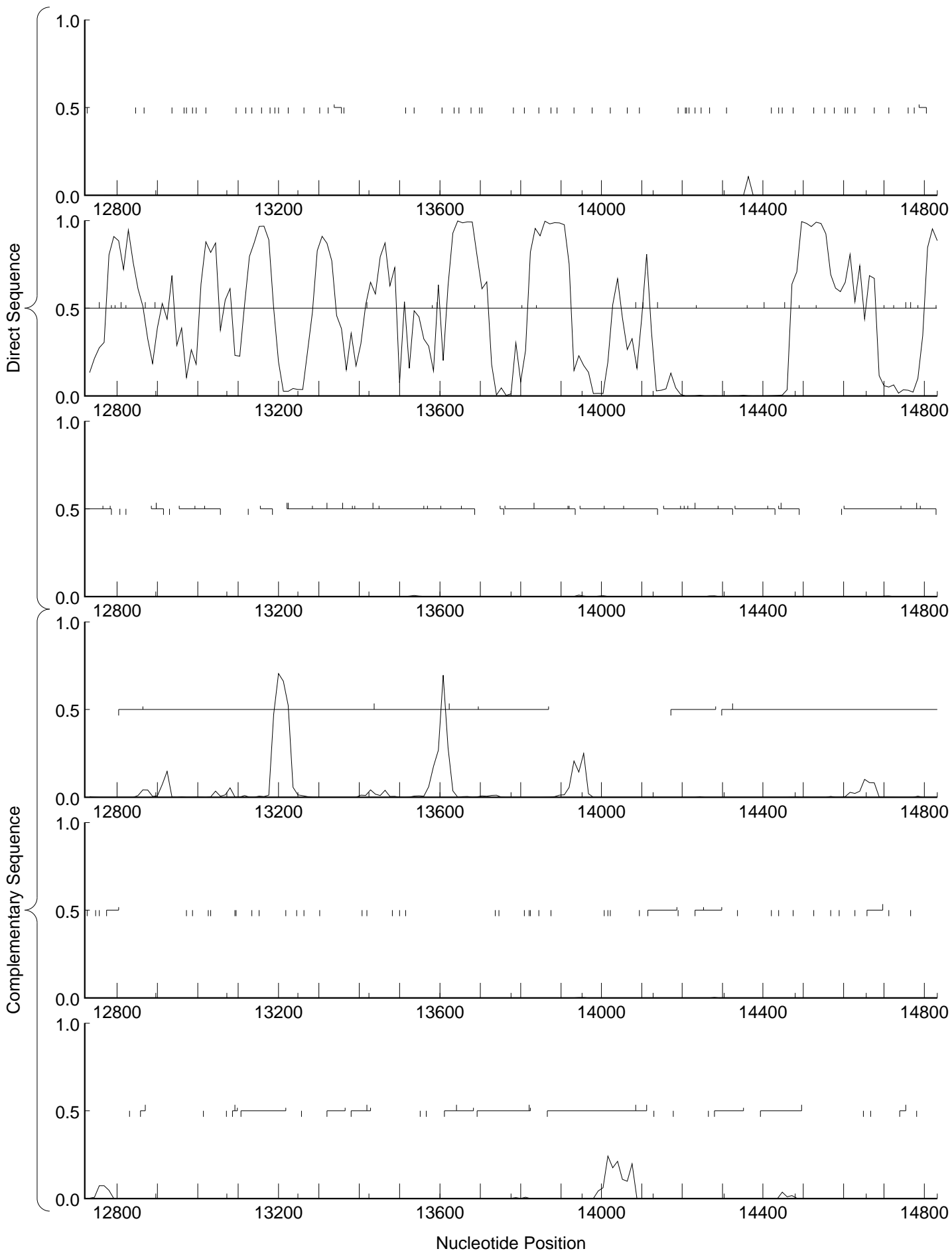


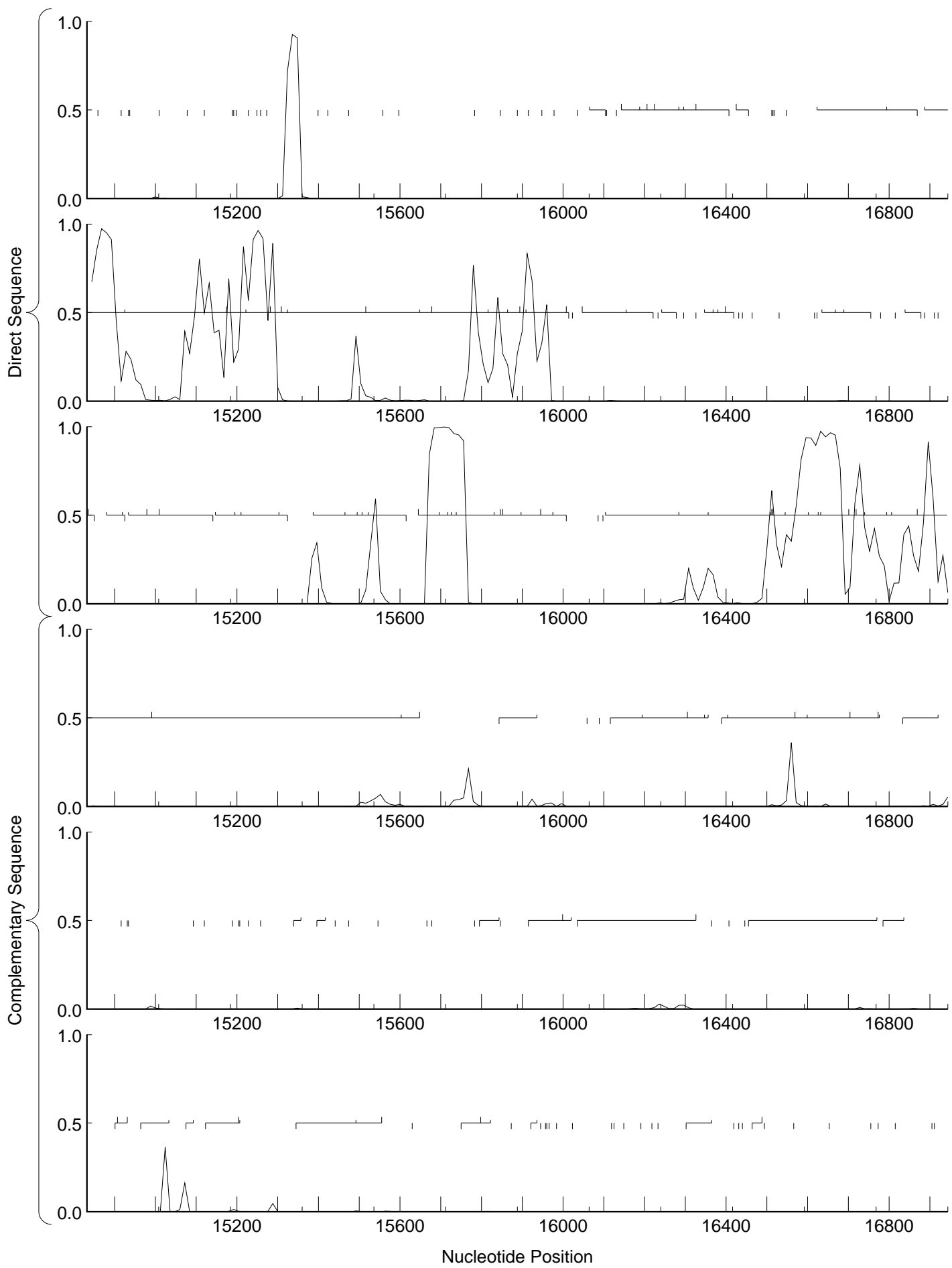


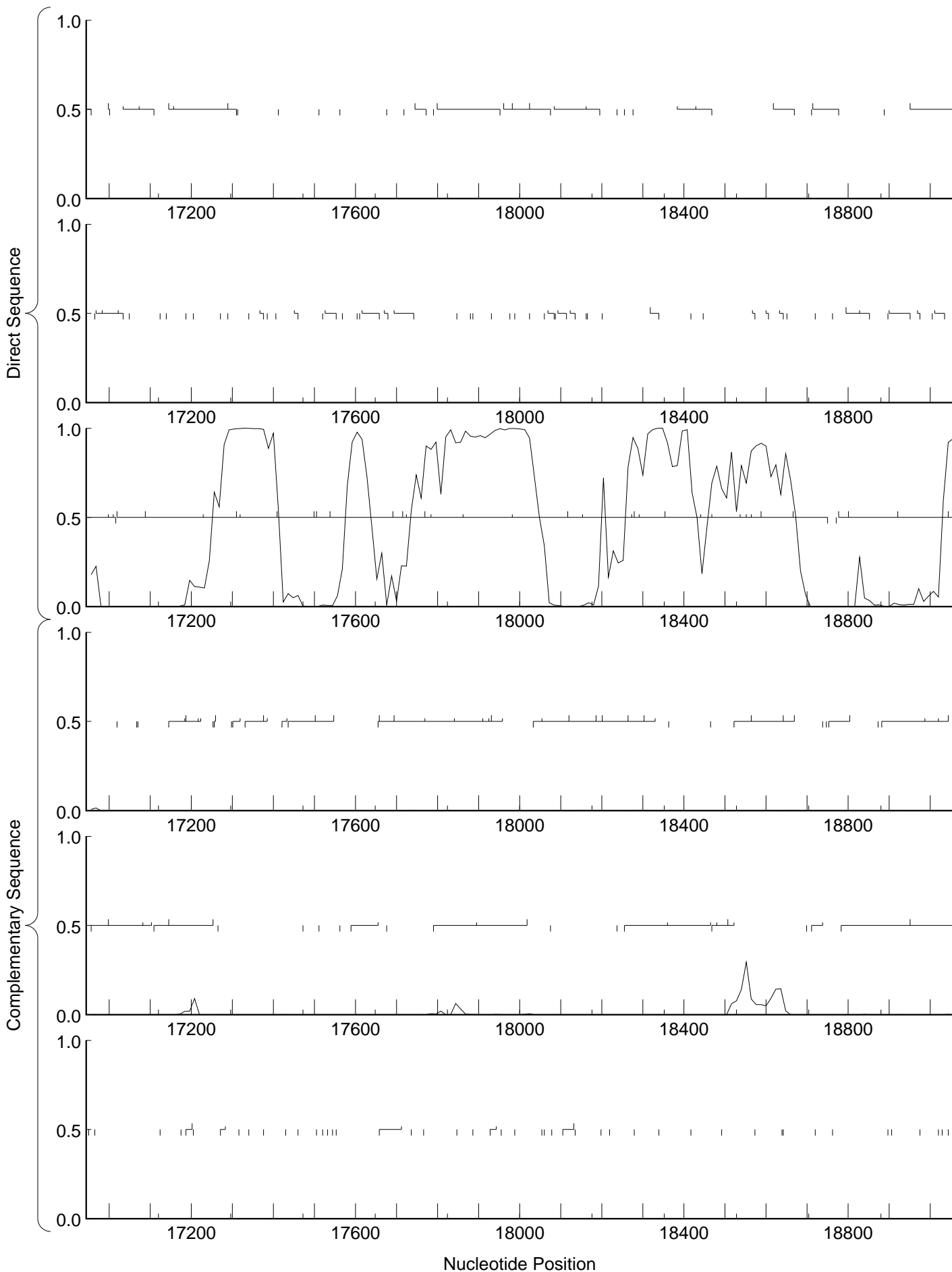


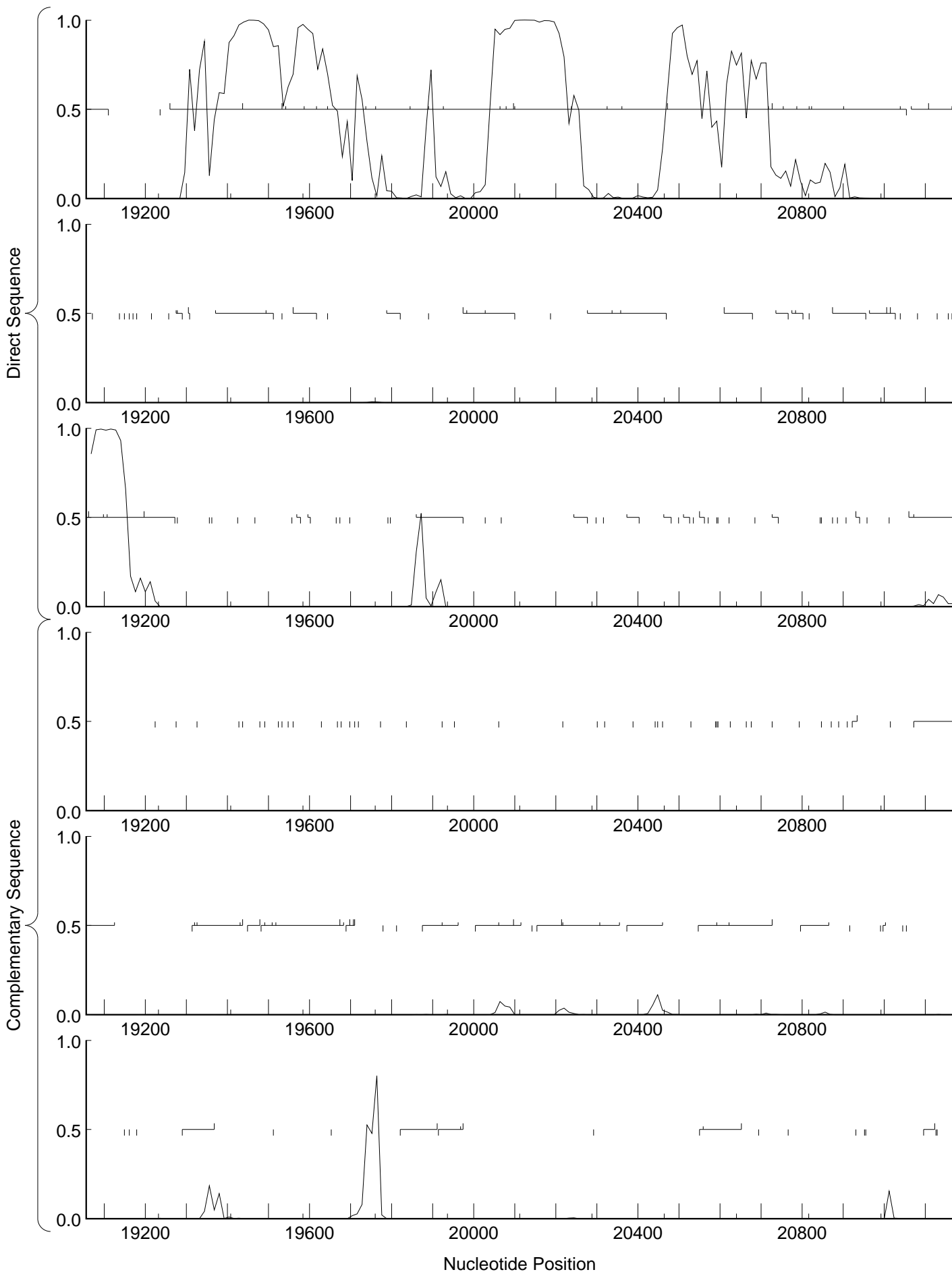




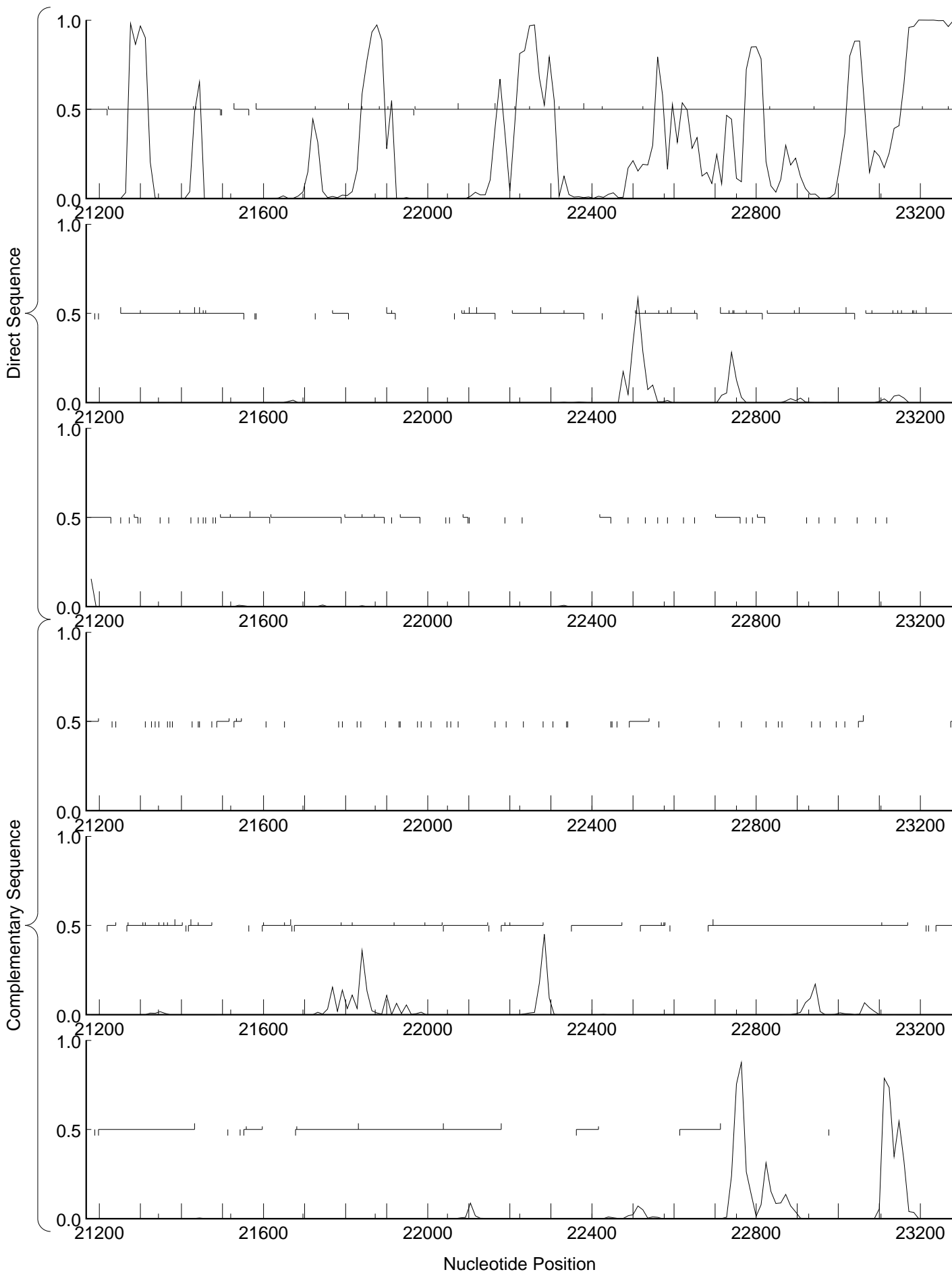


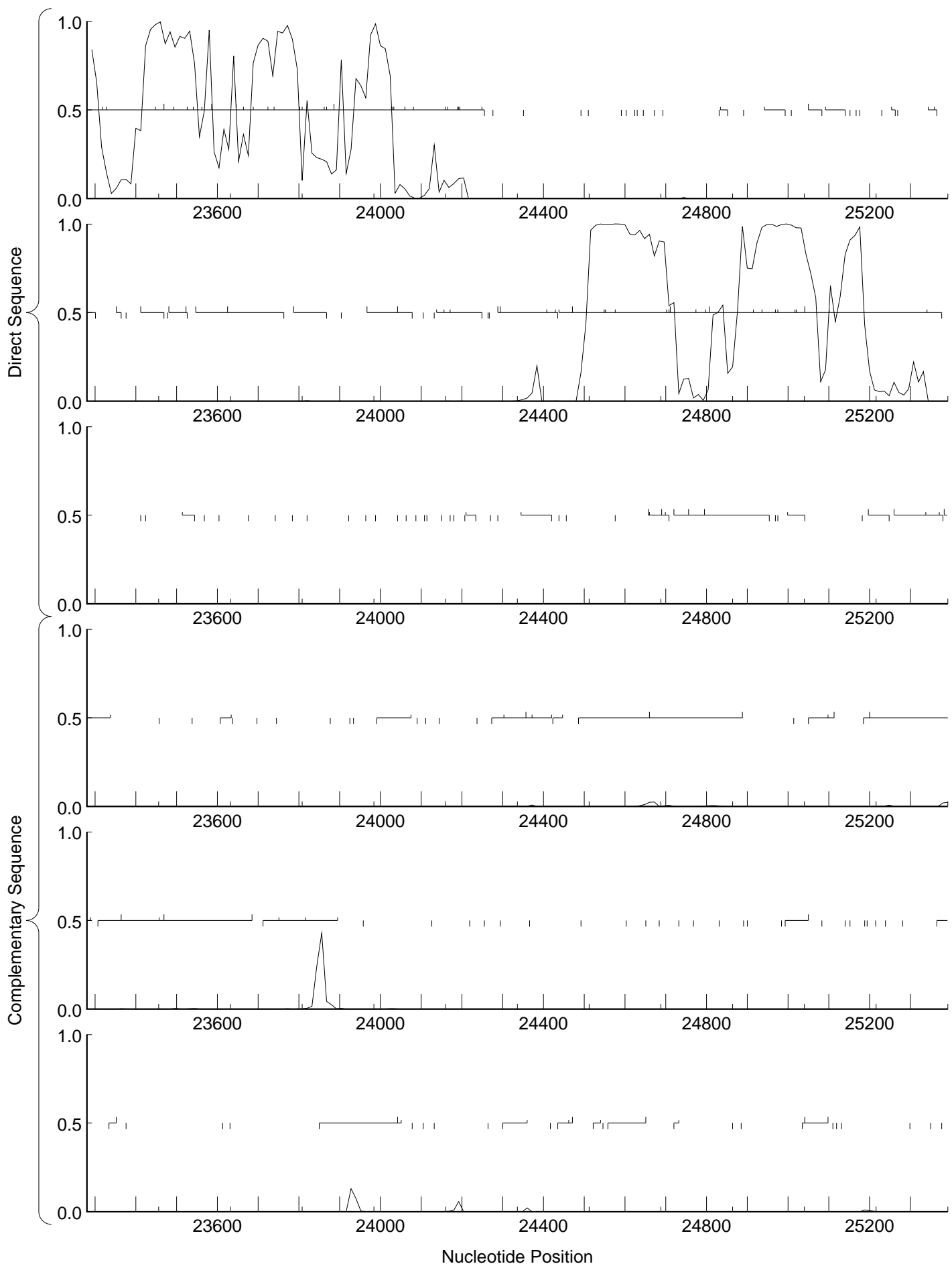


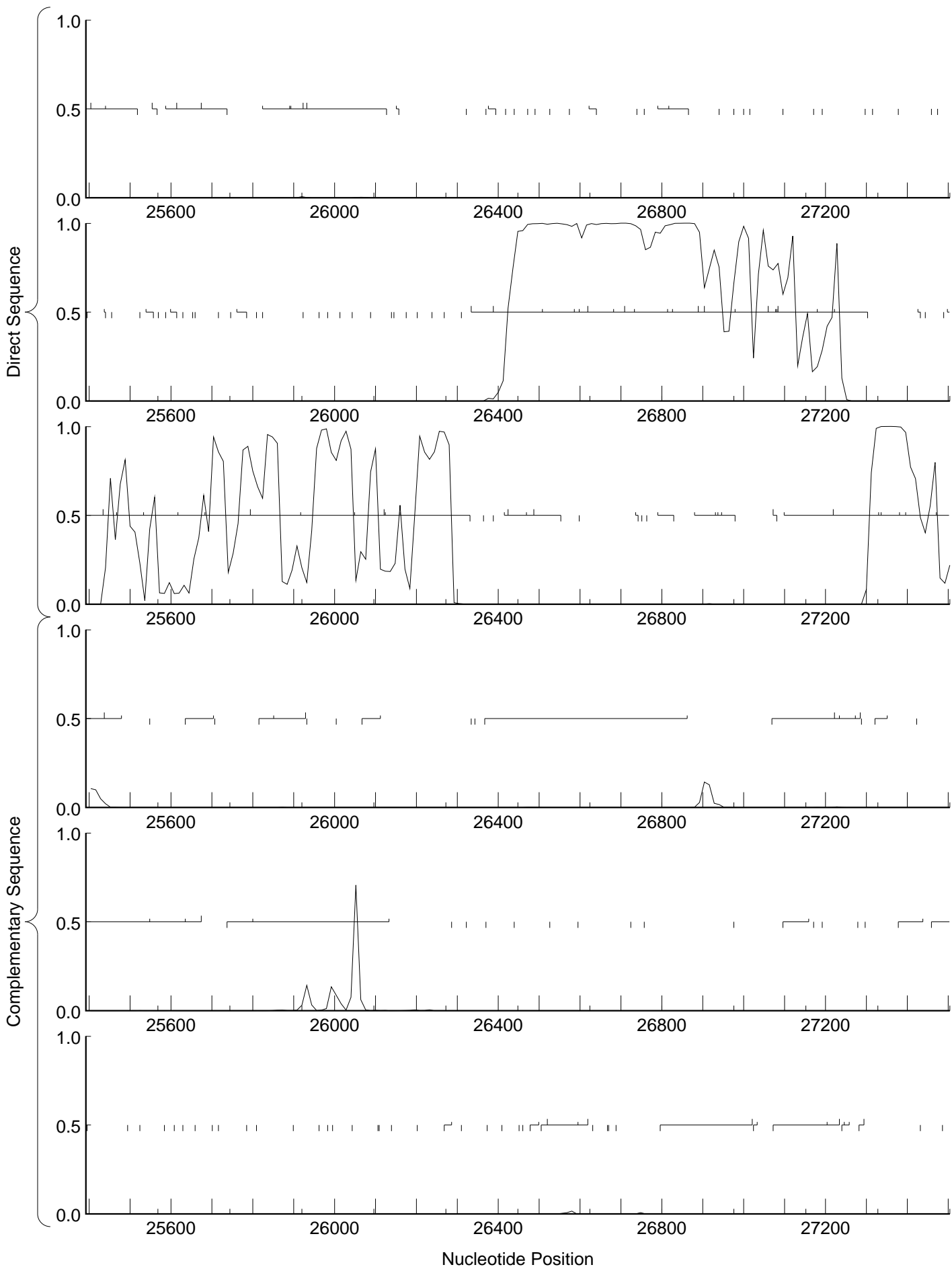


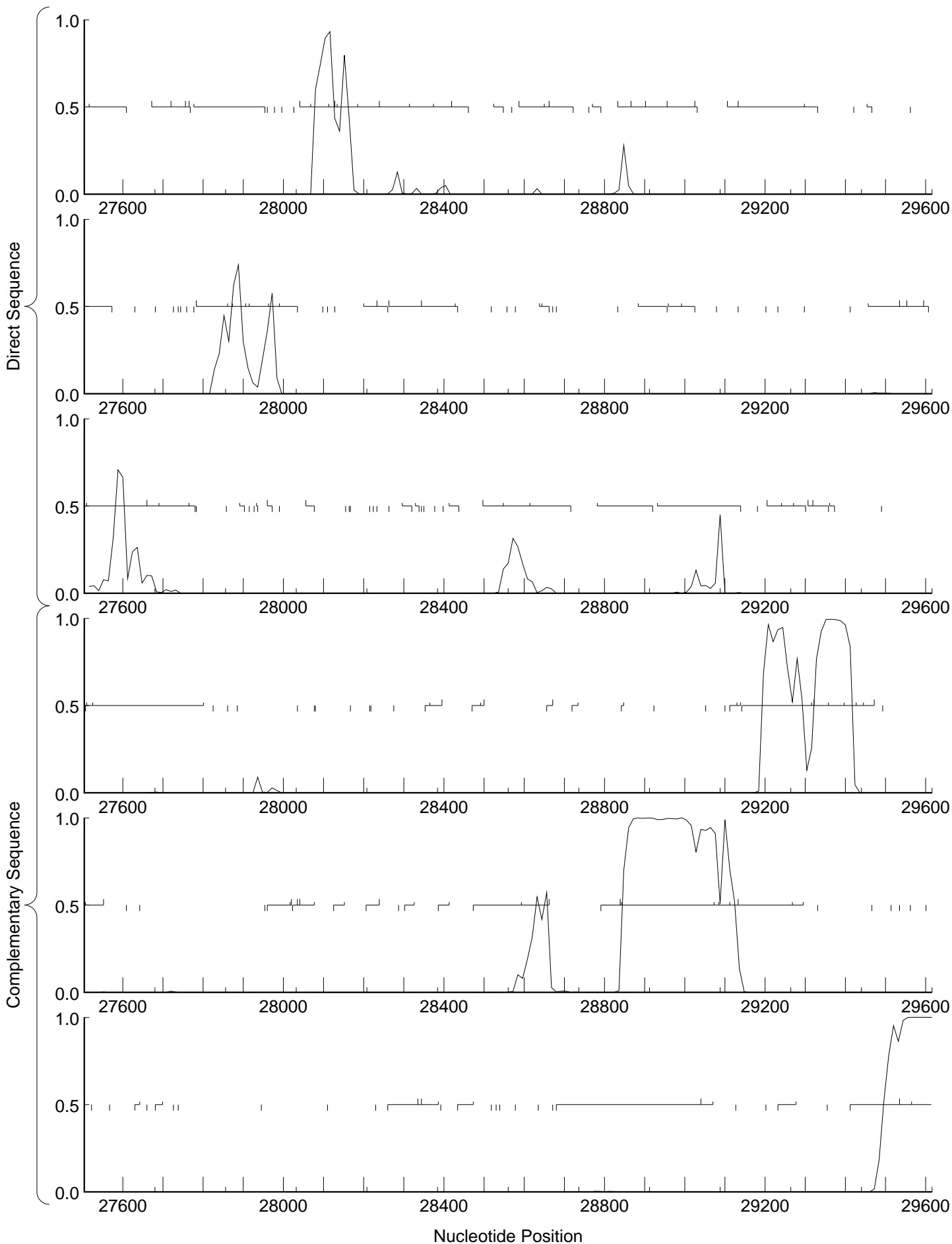


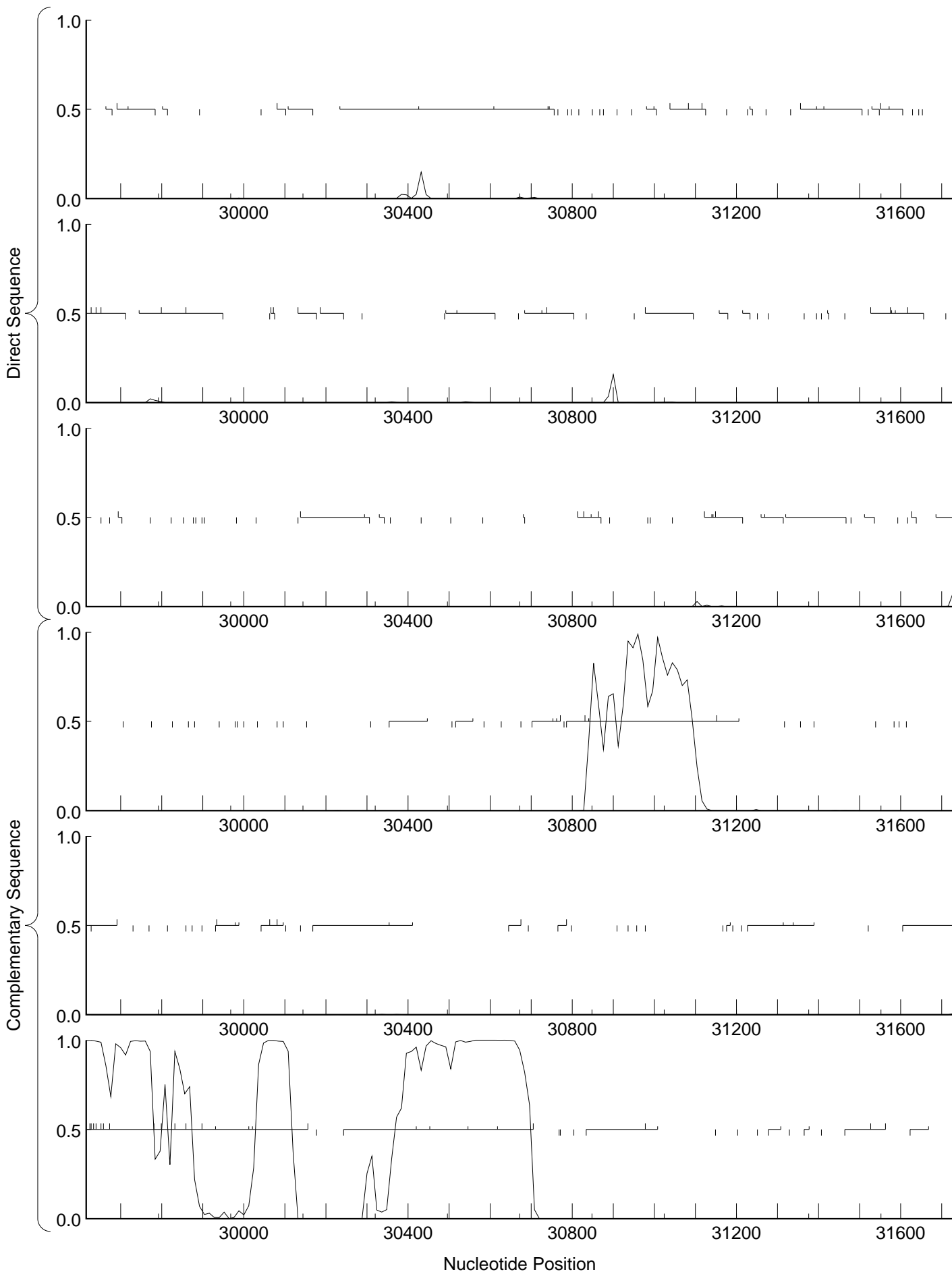
Complete sequence, 76871 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 12/38

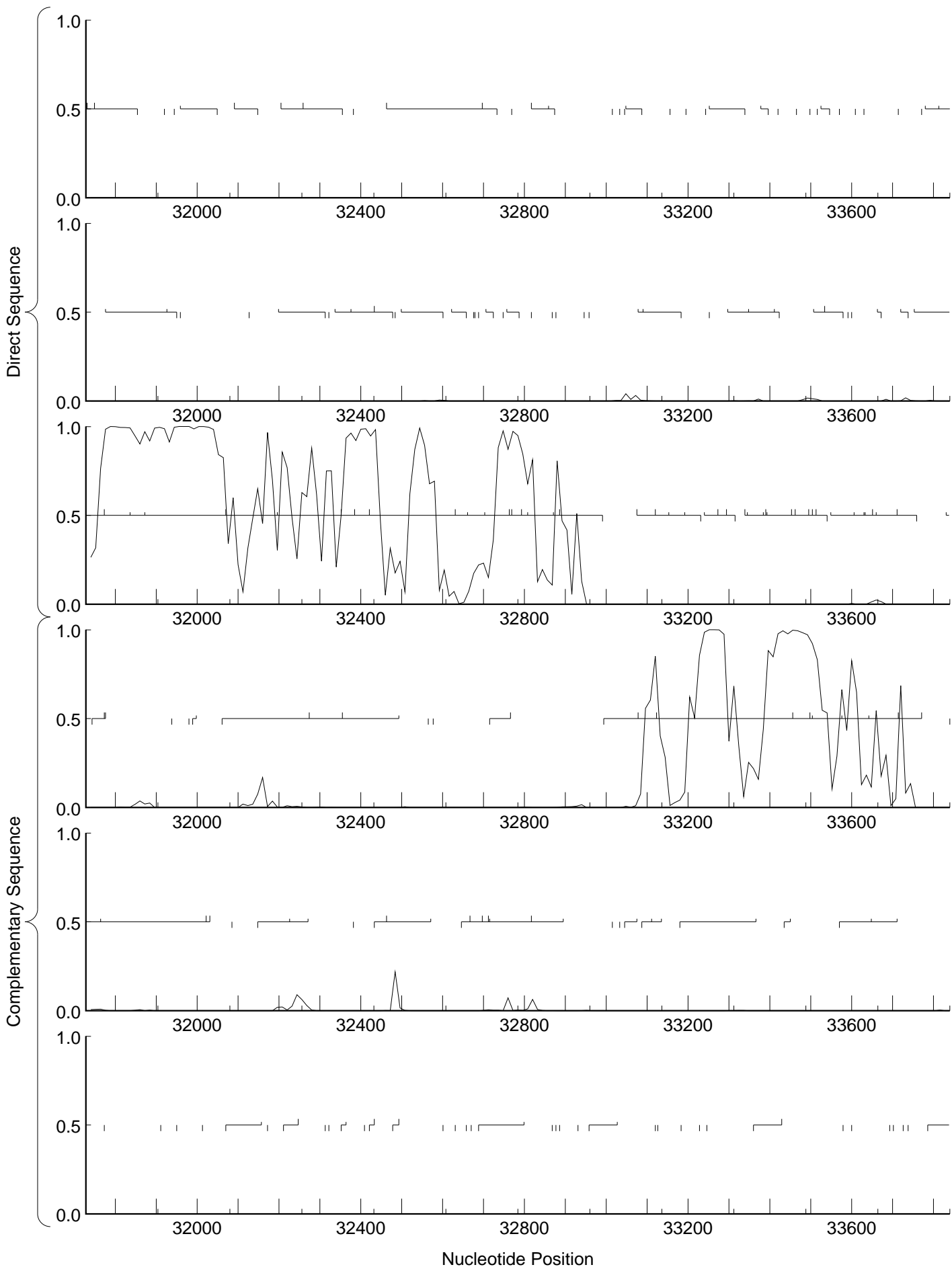


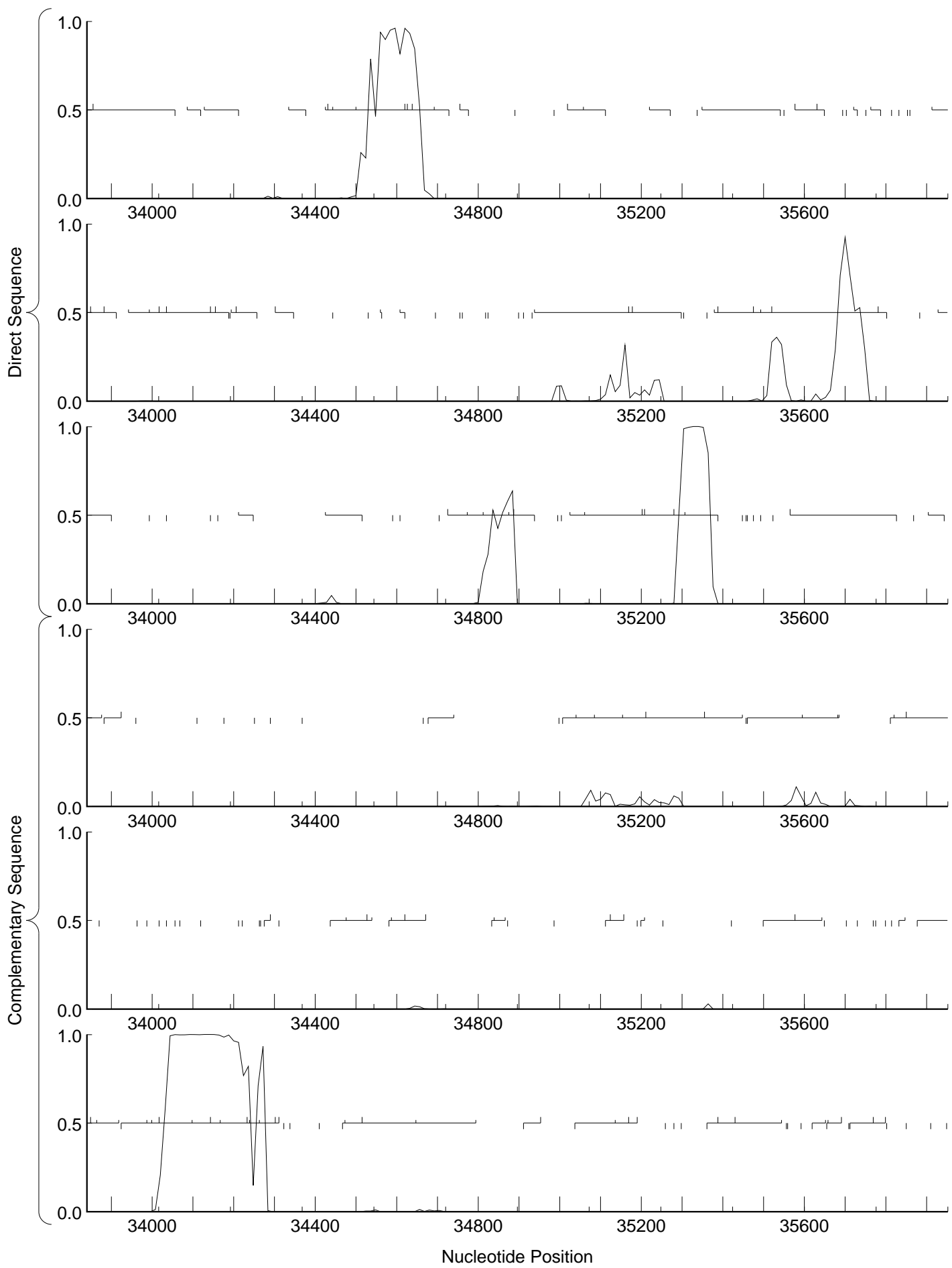


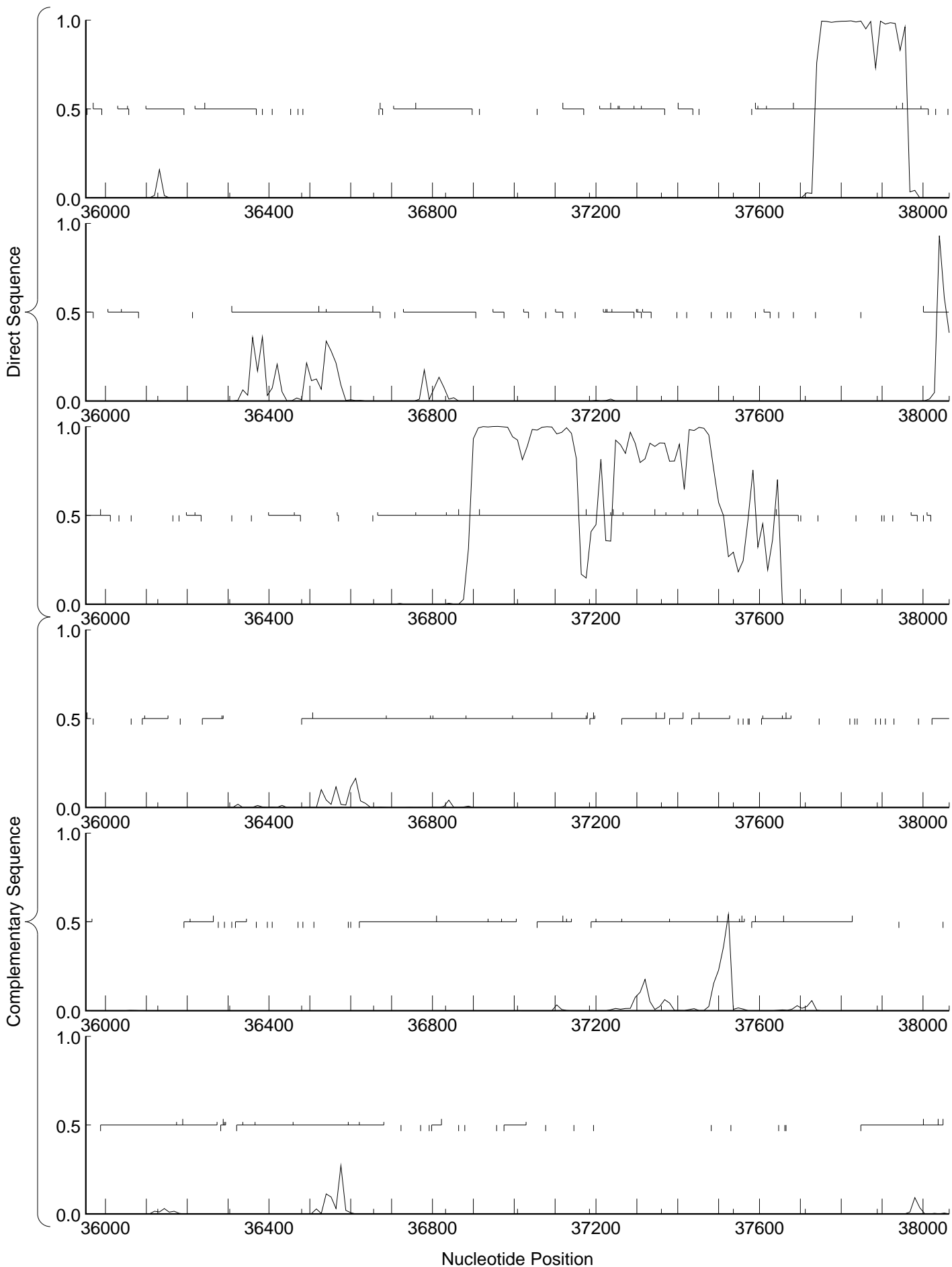


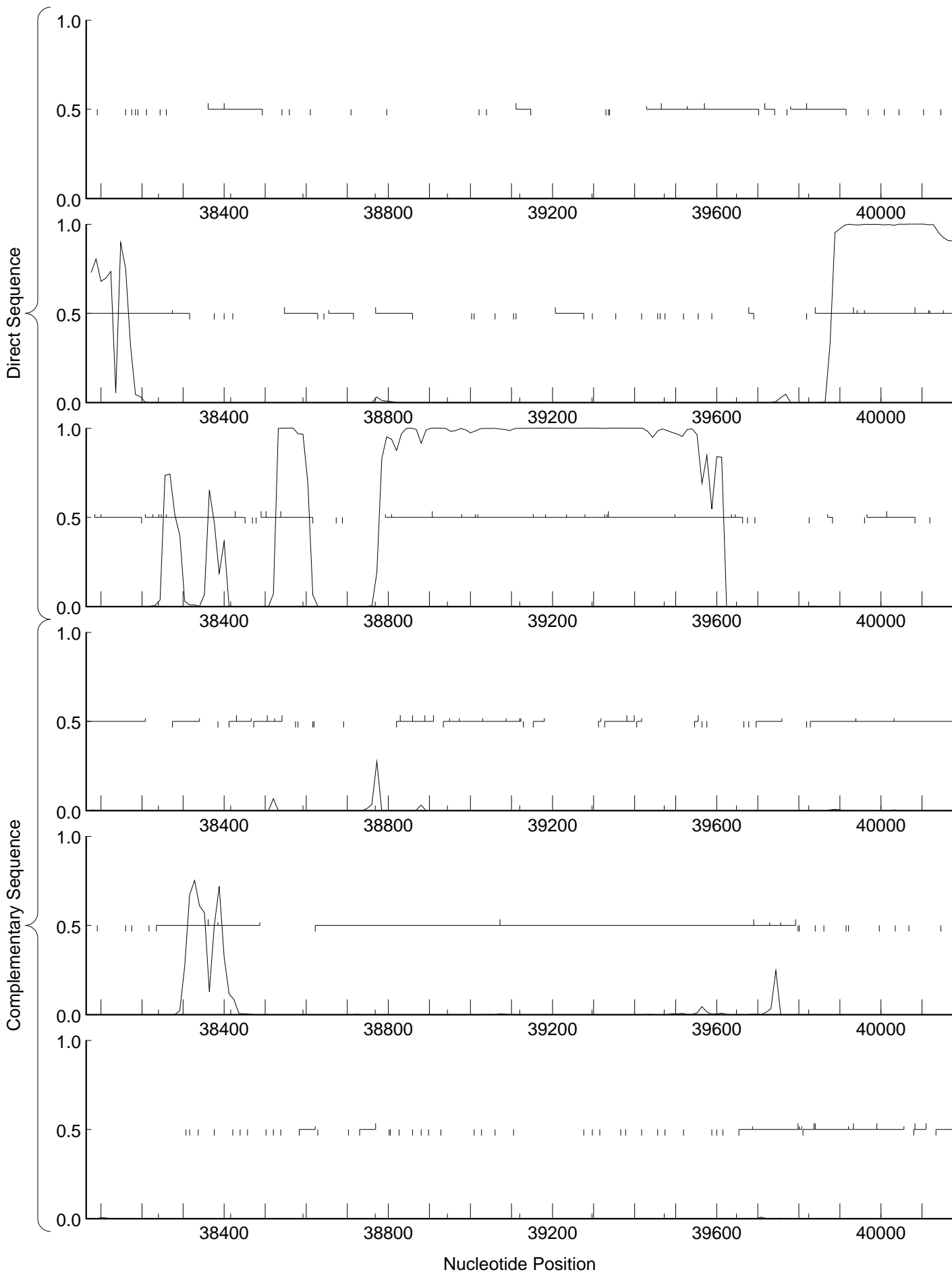


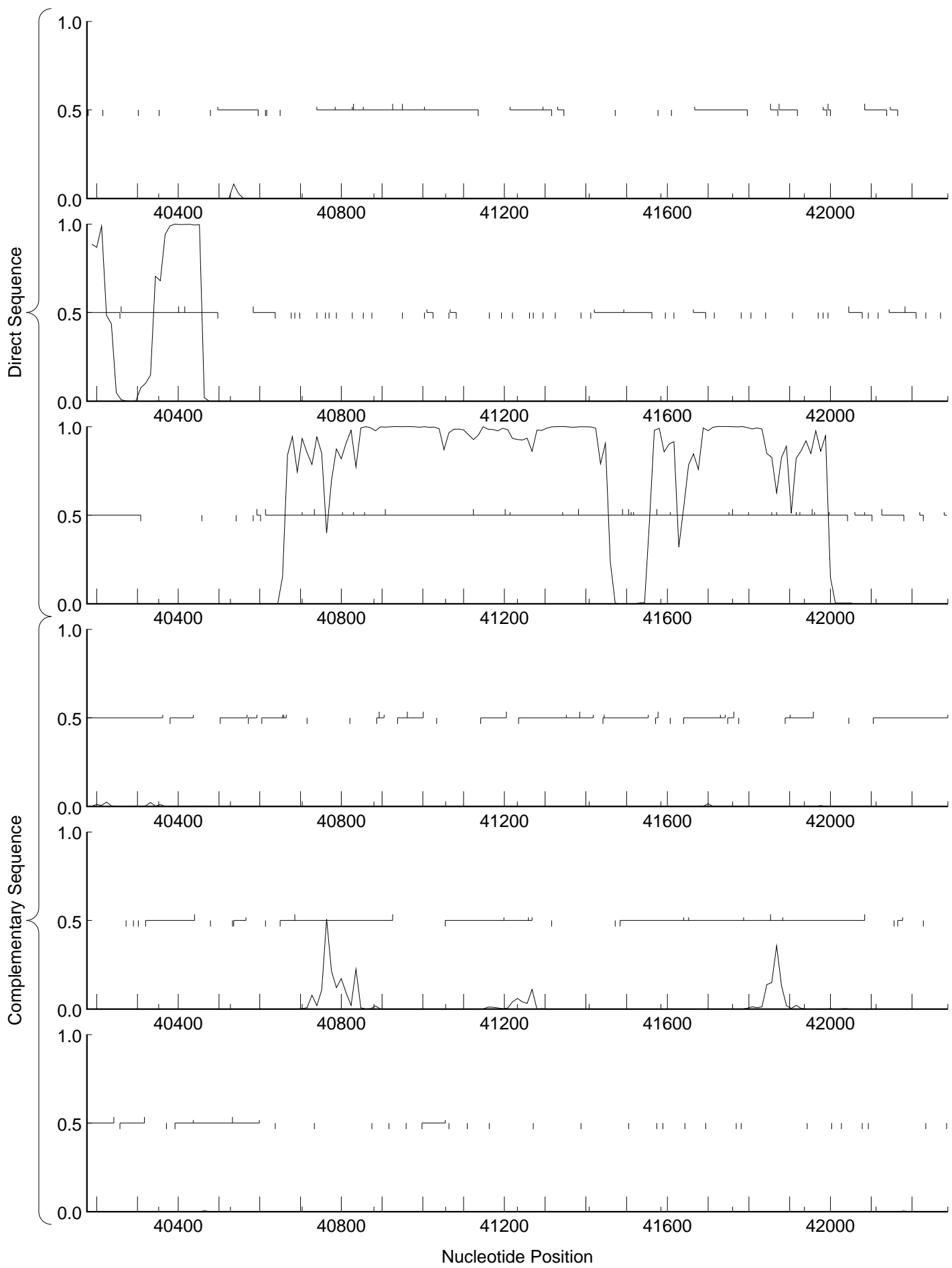


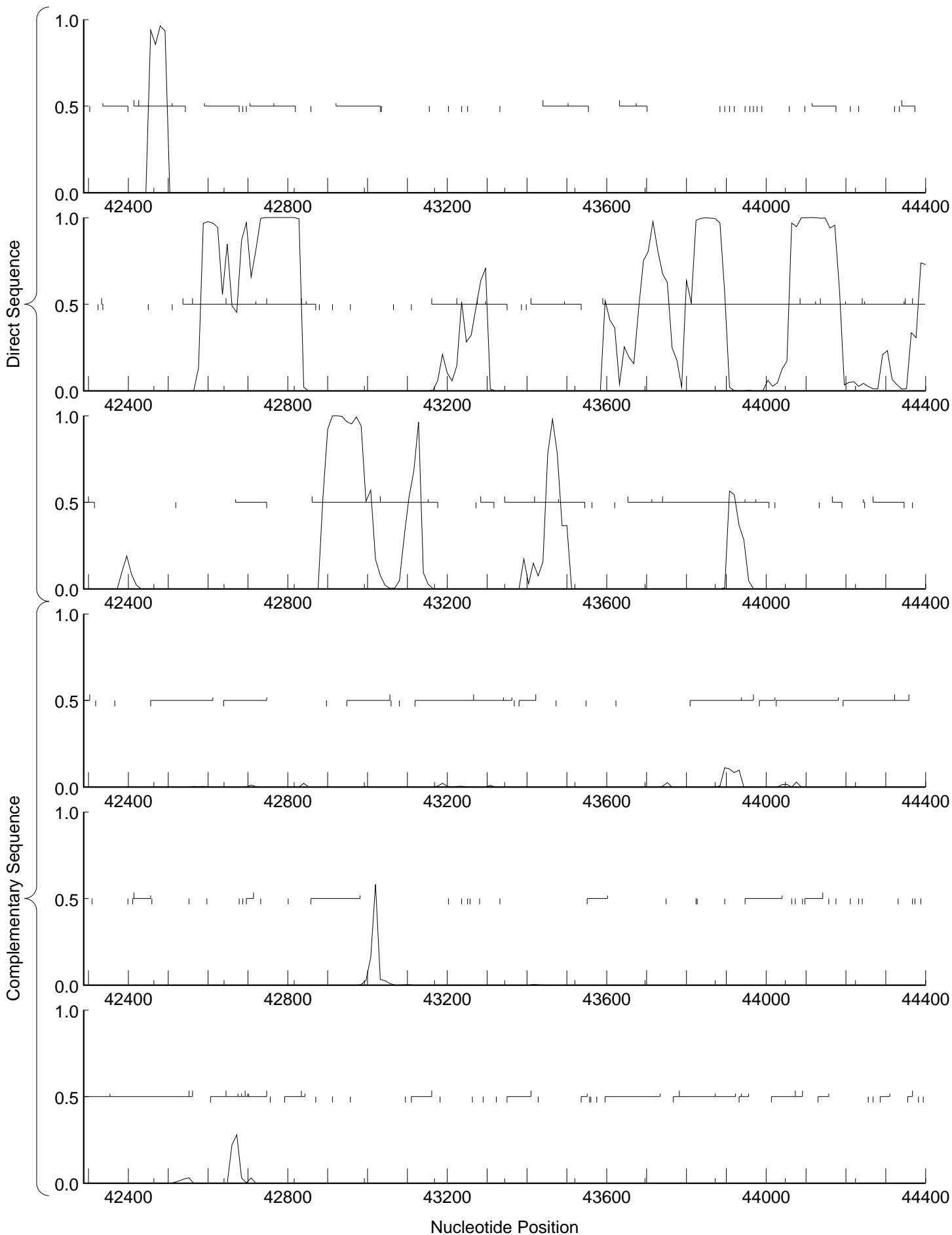


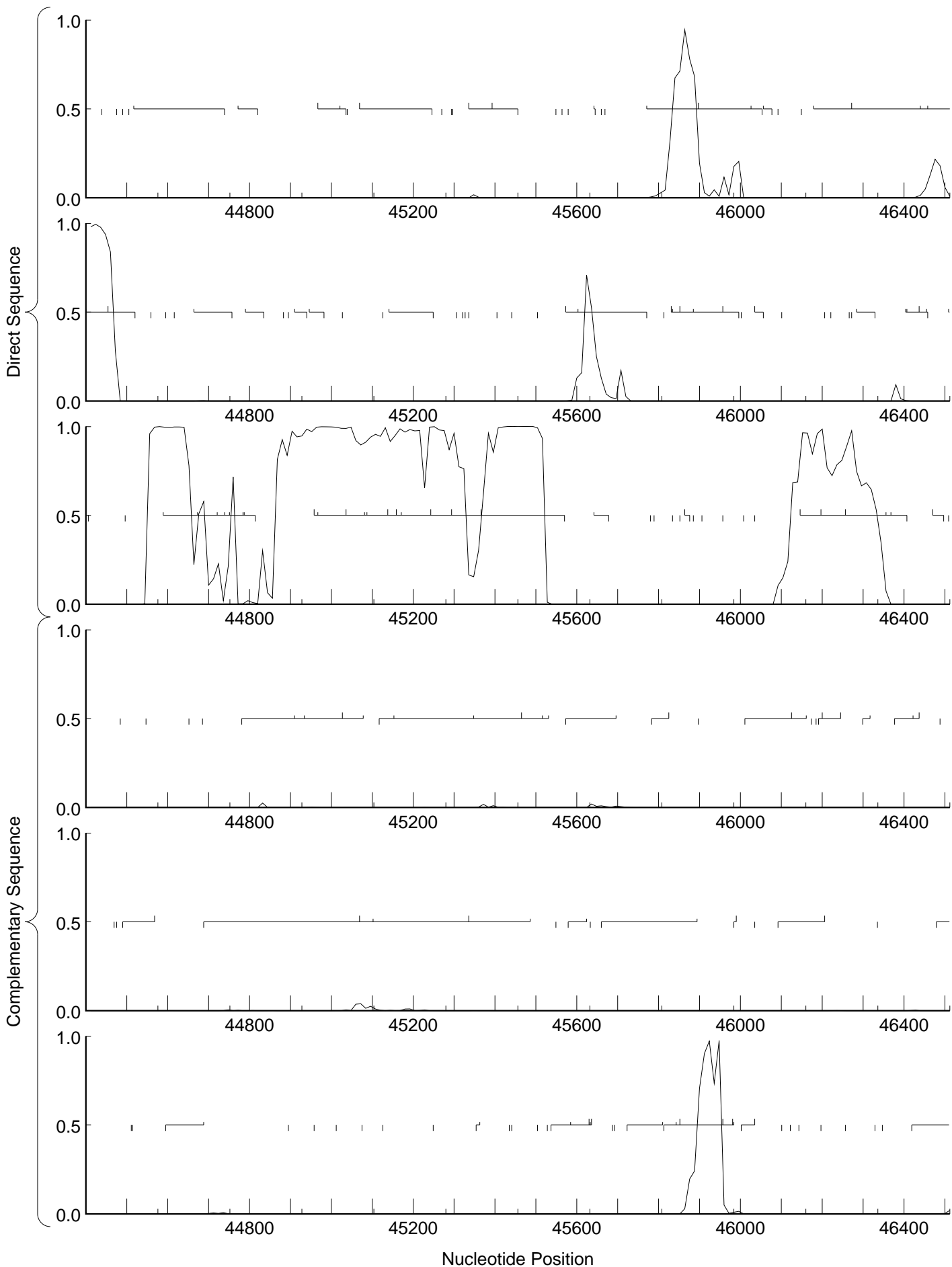


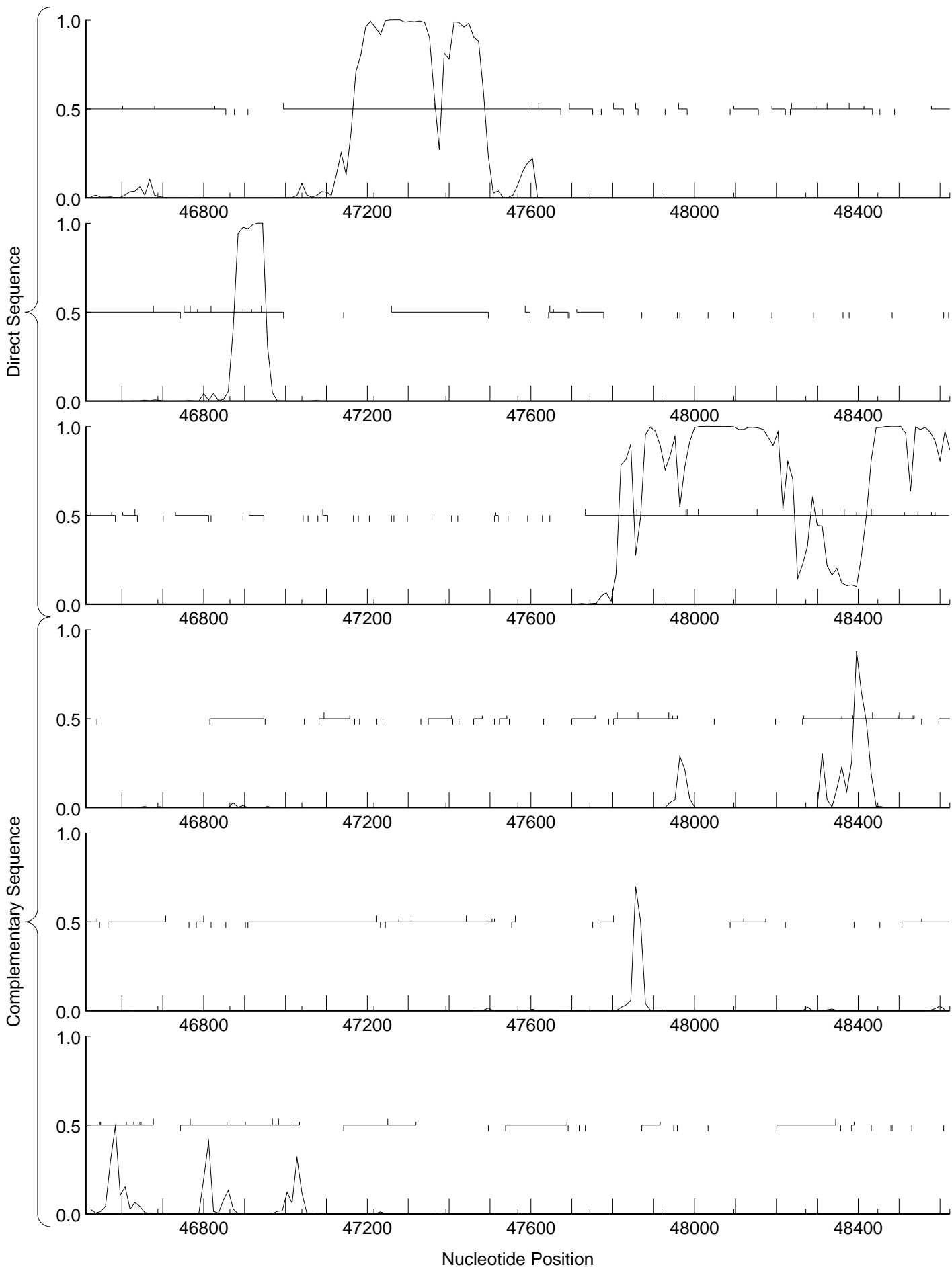


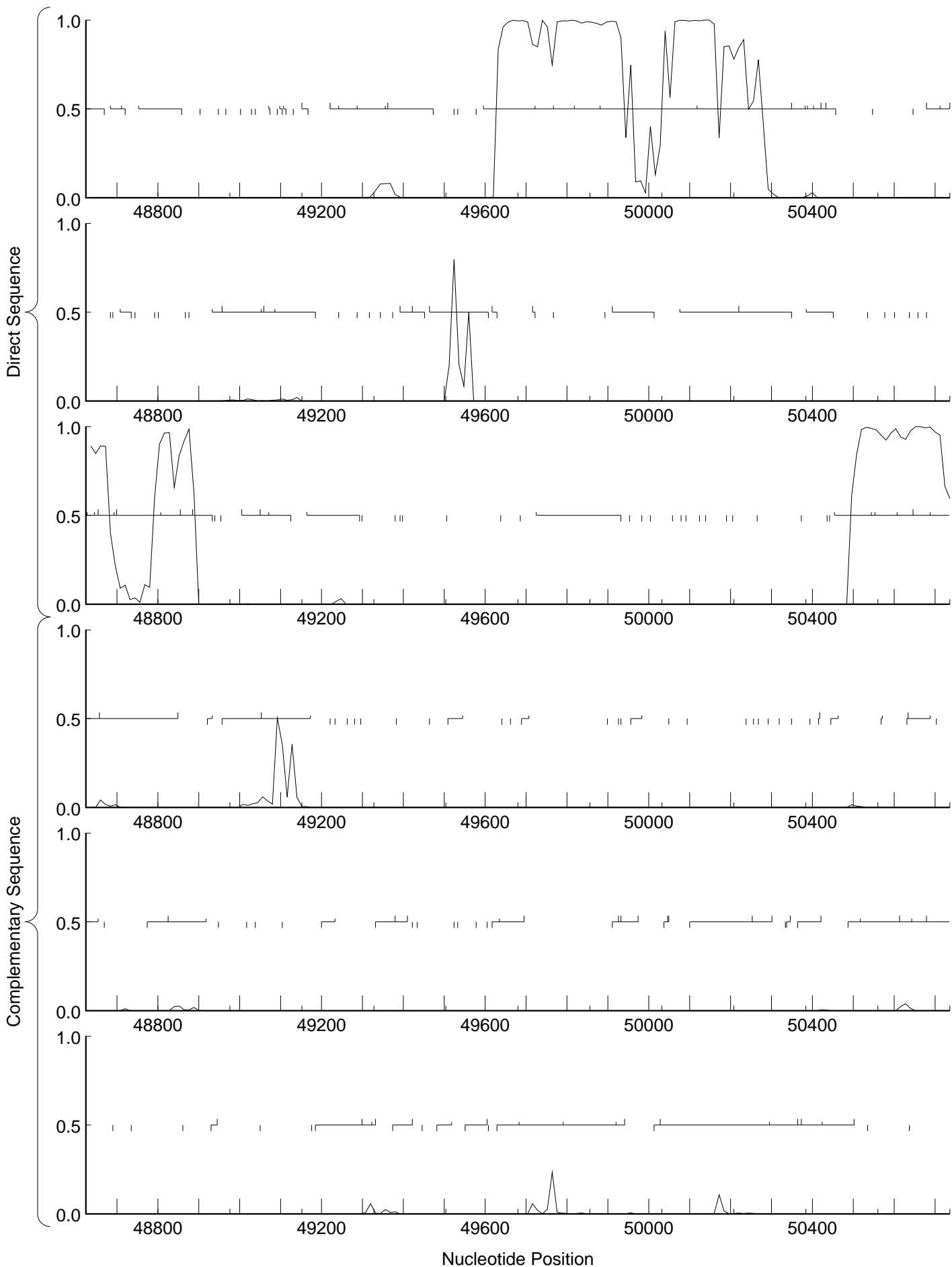


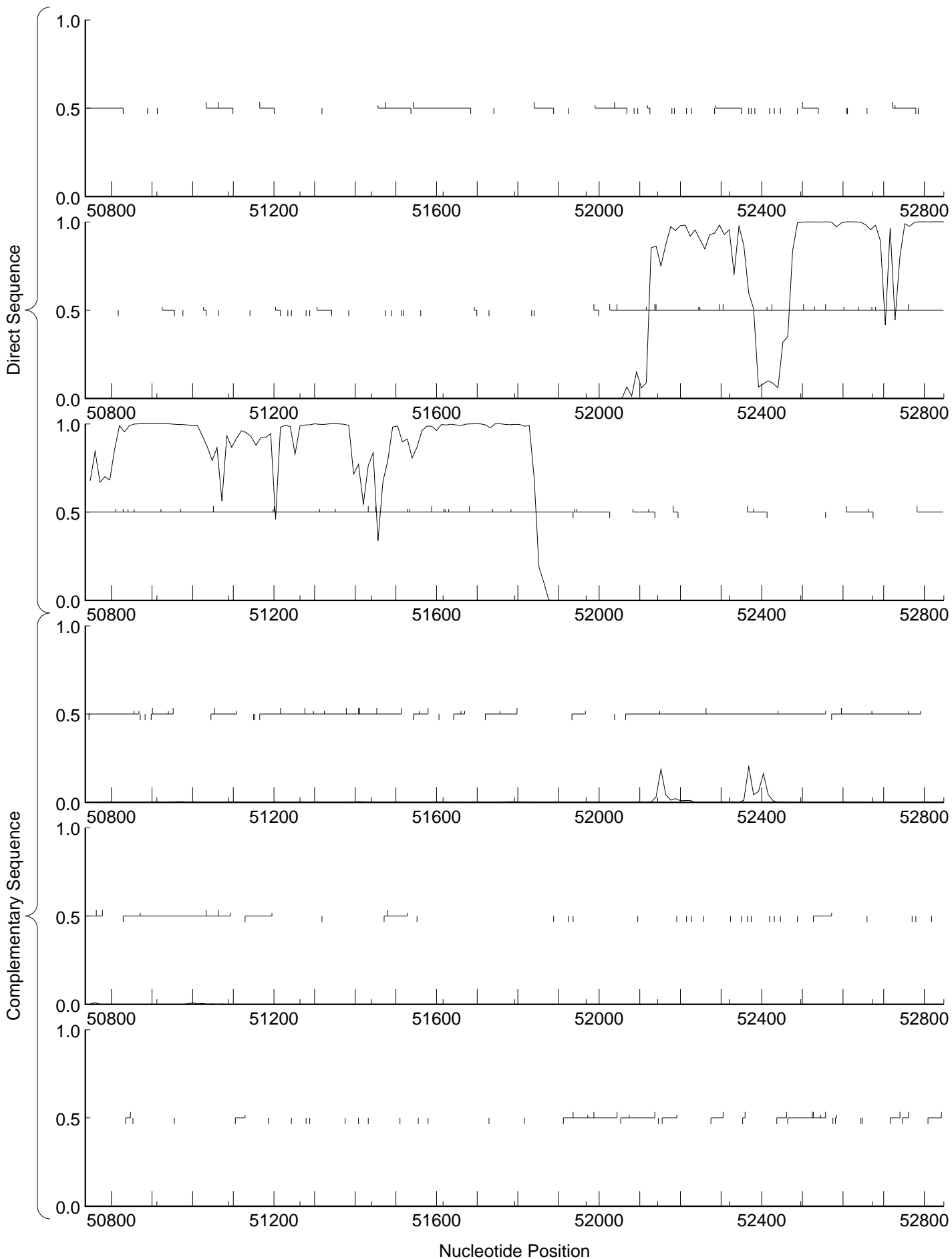


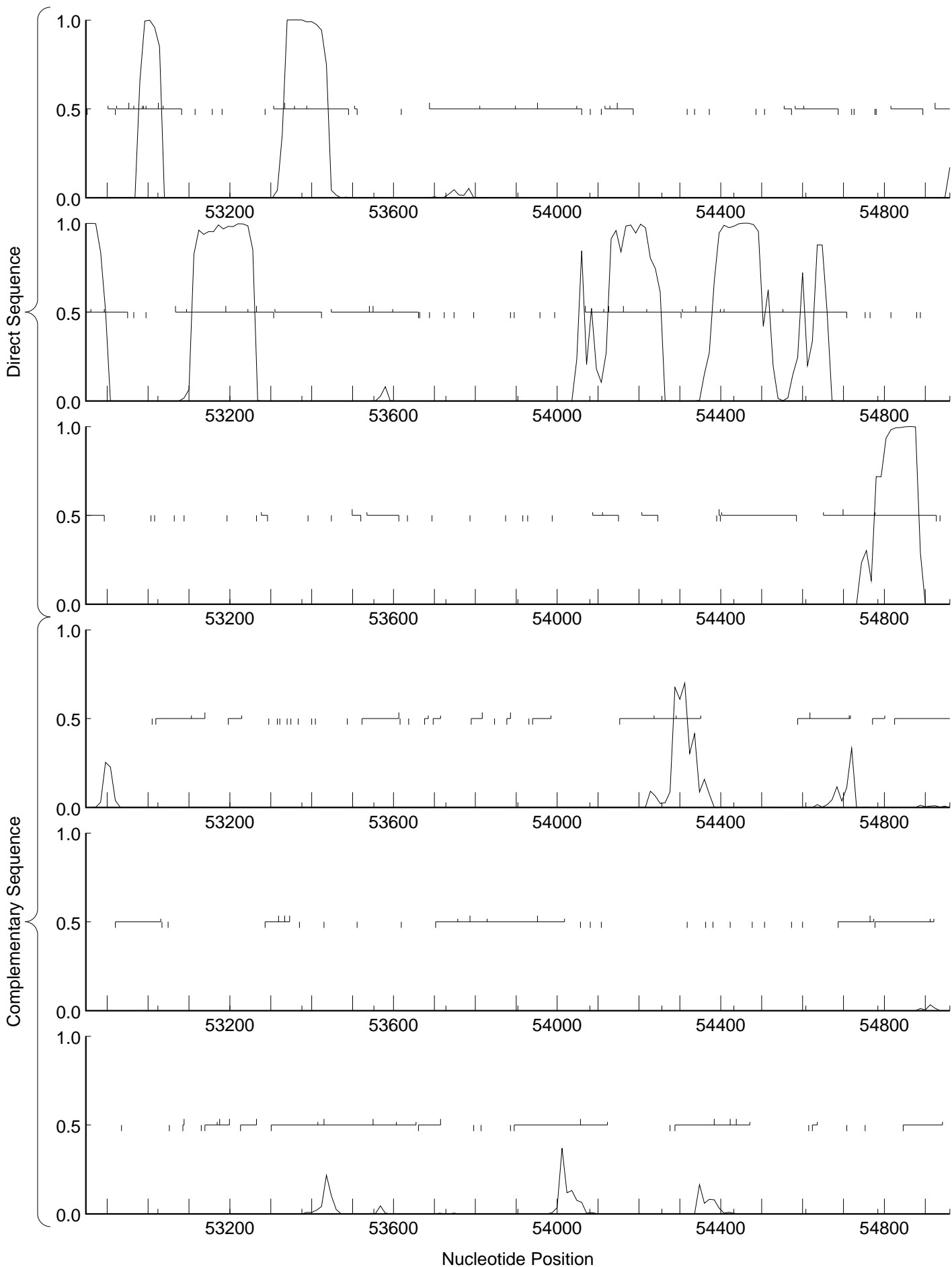


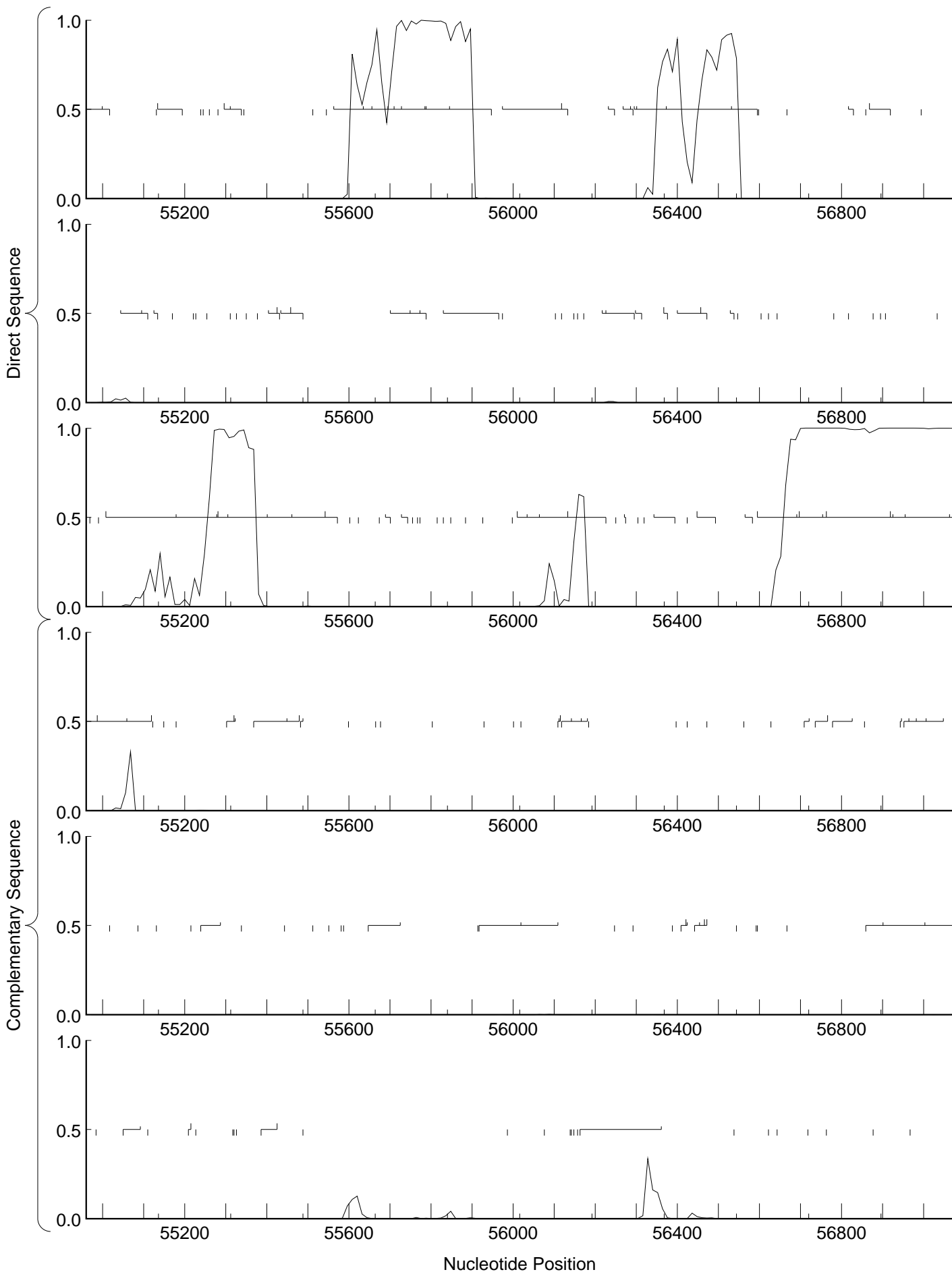


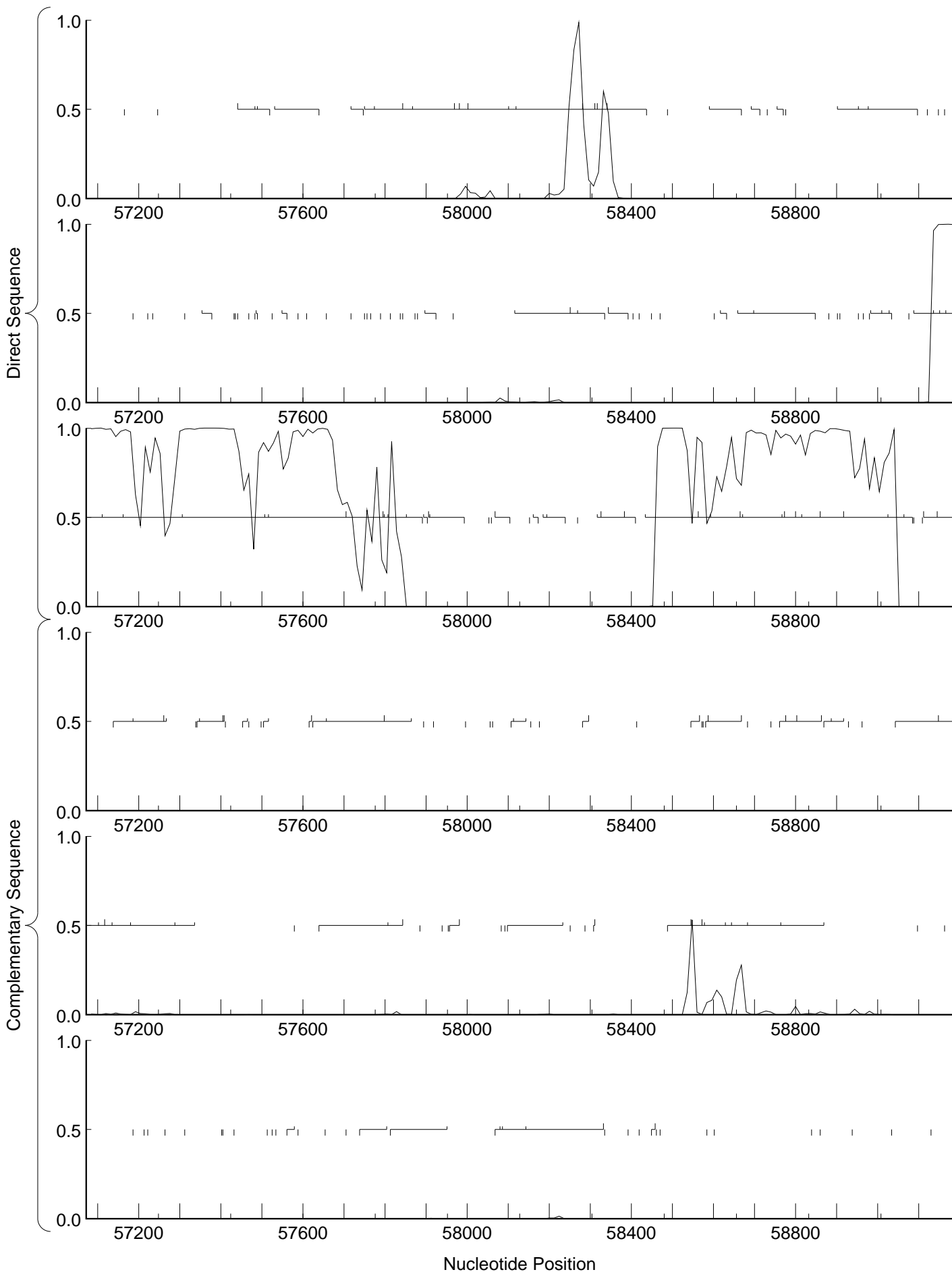




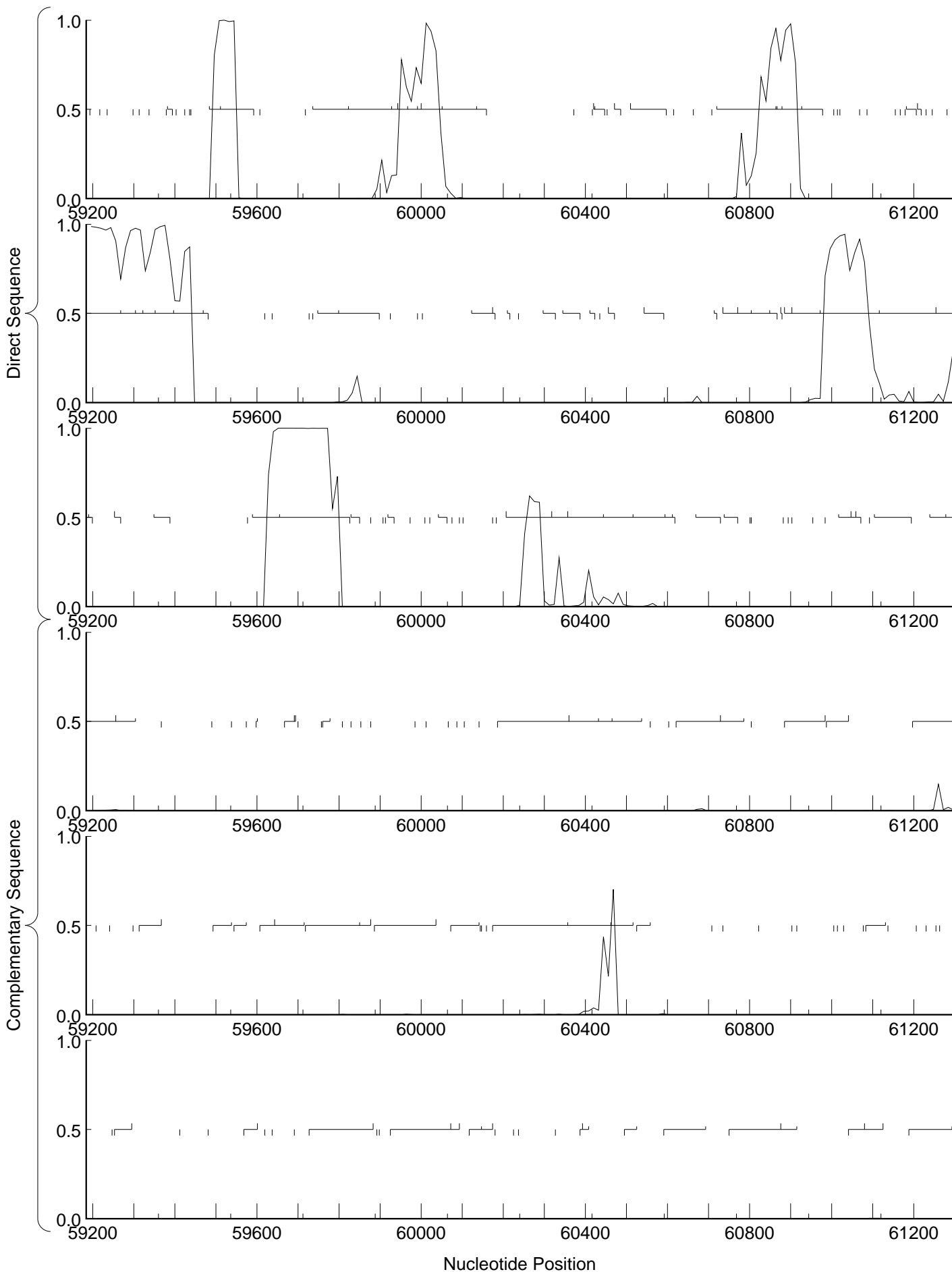


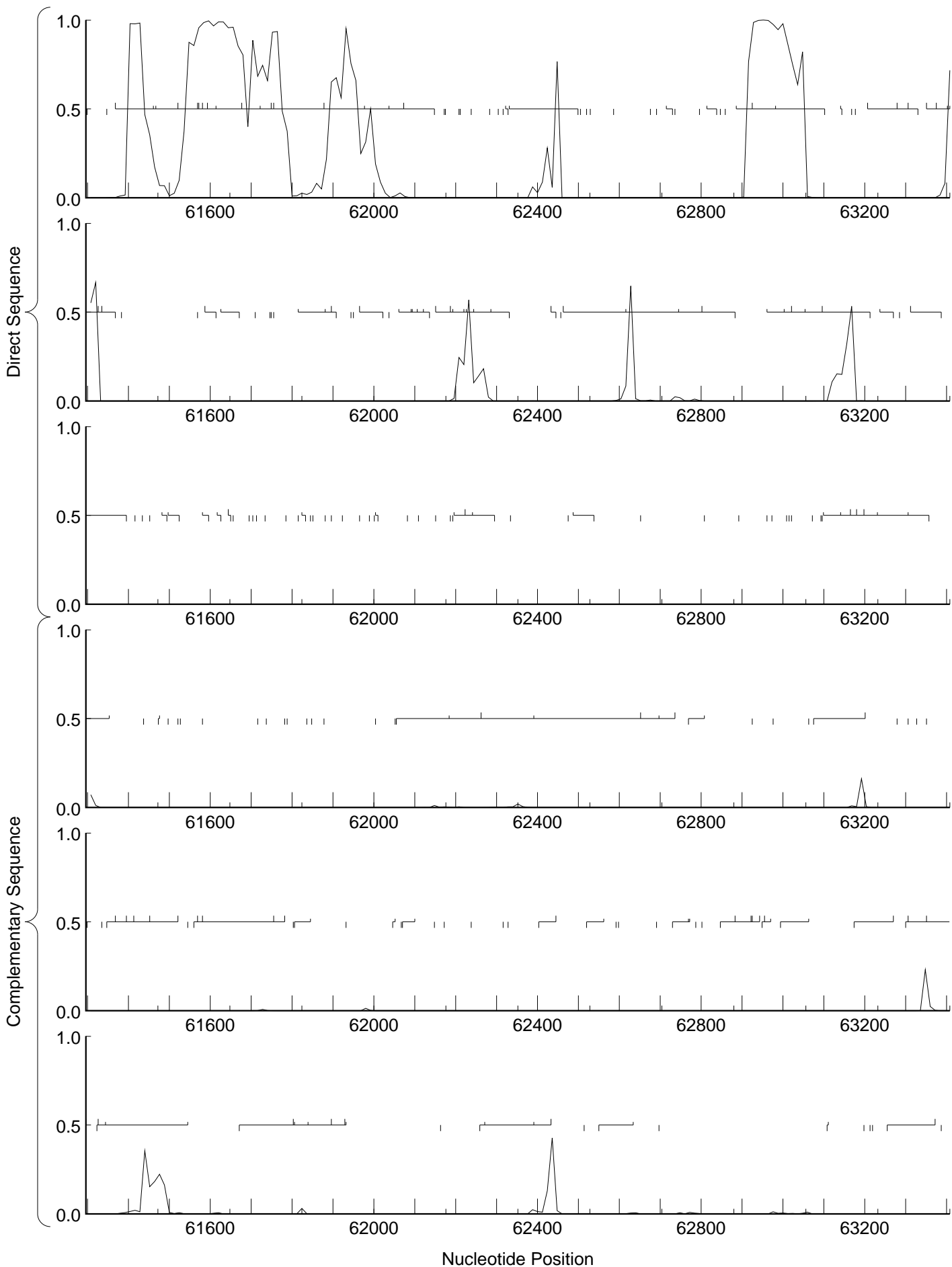


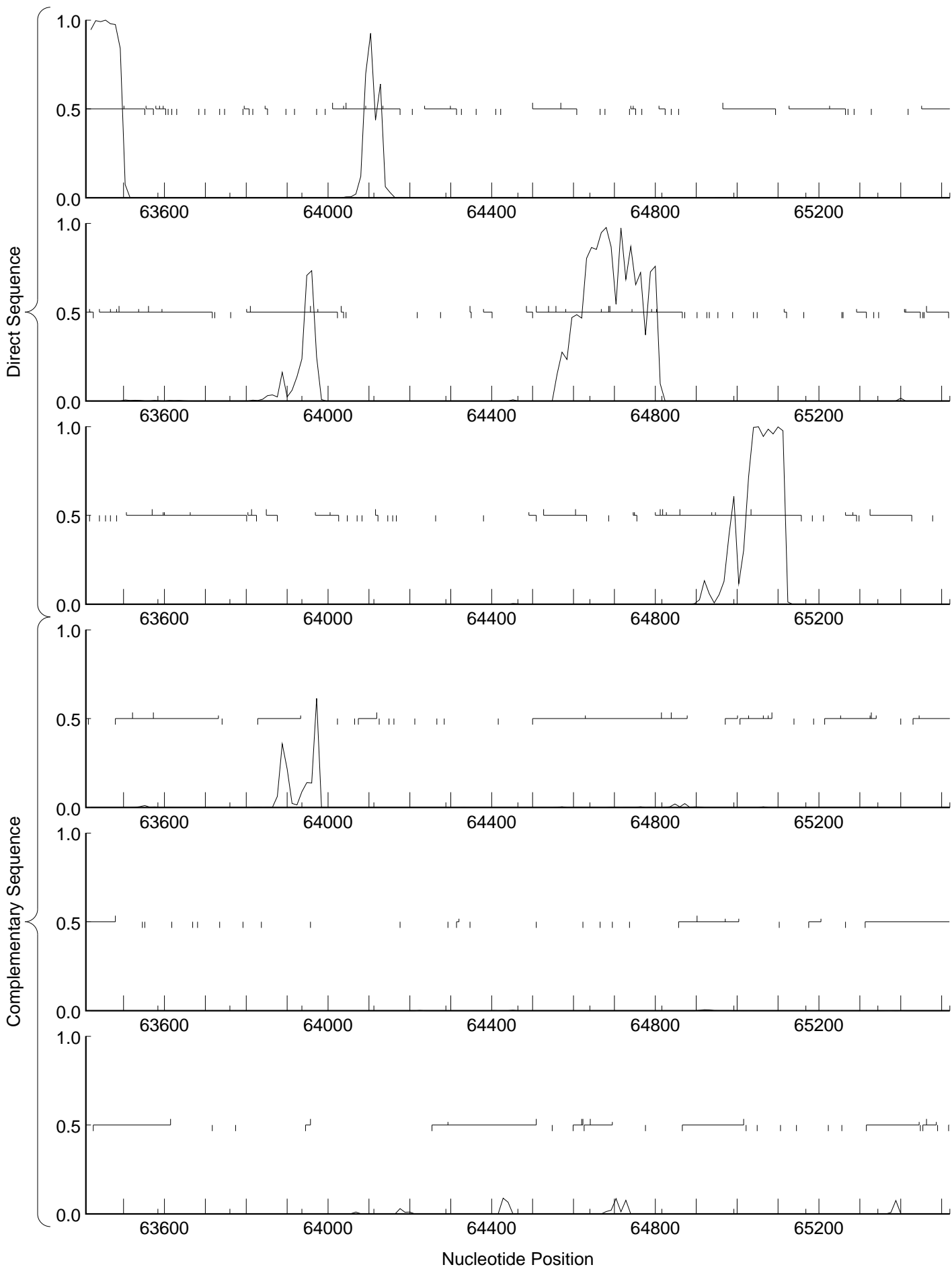


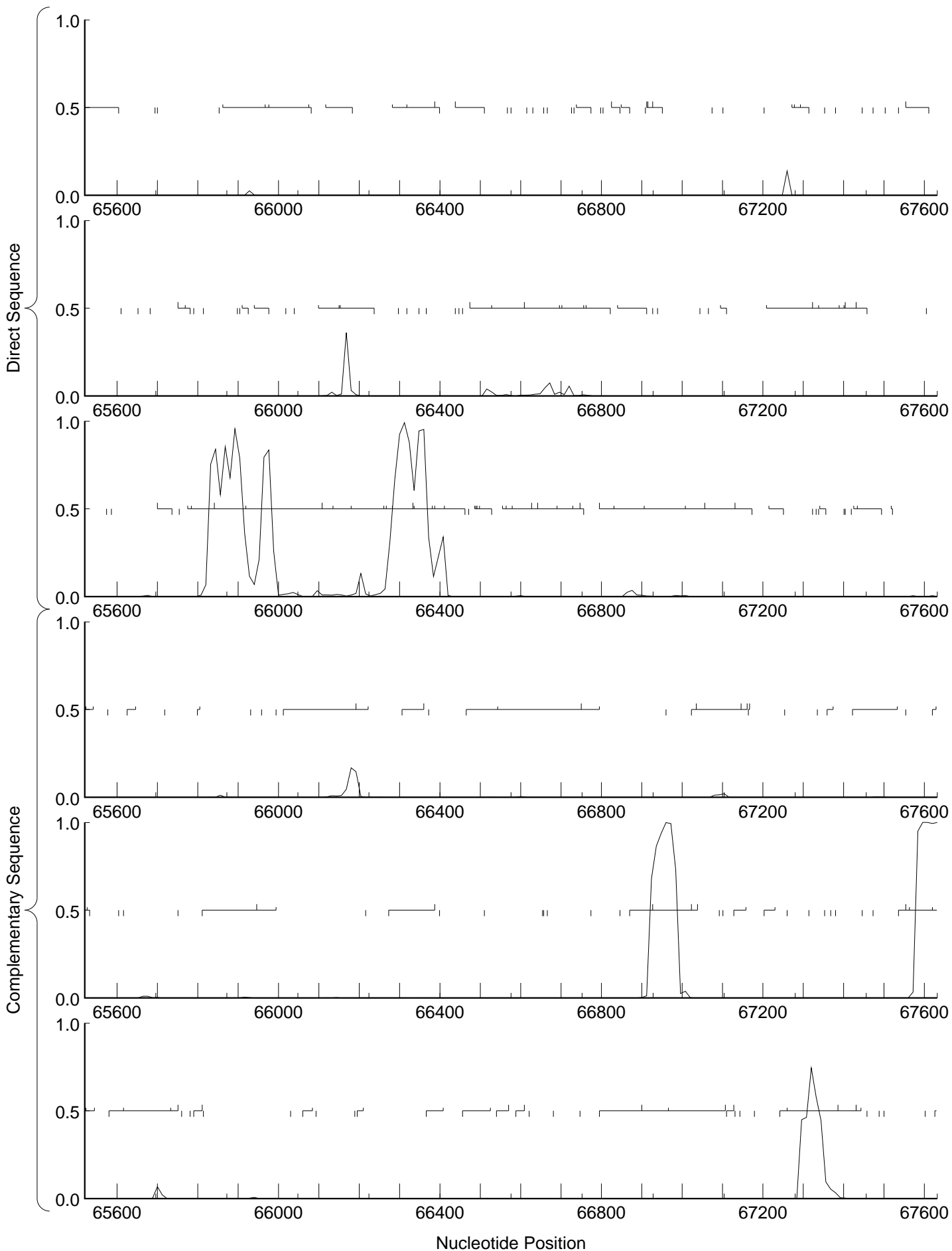


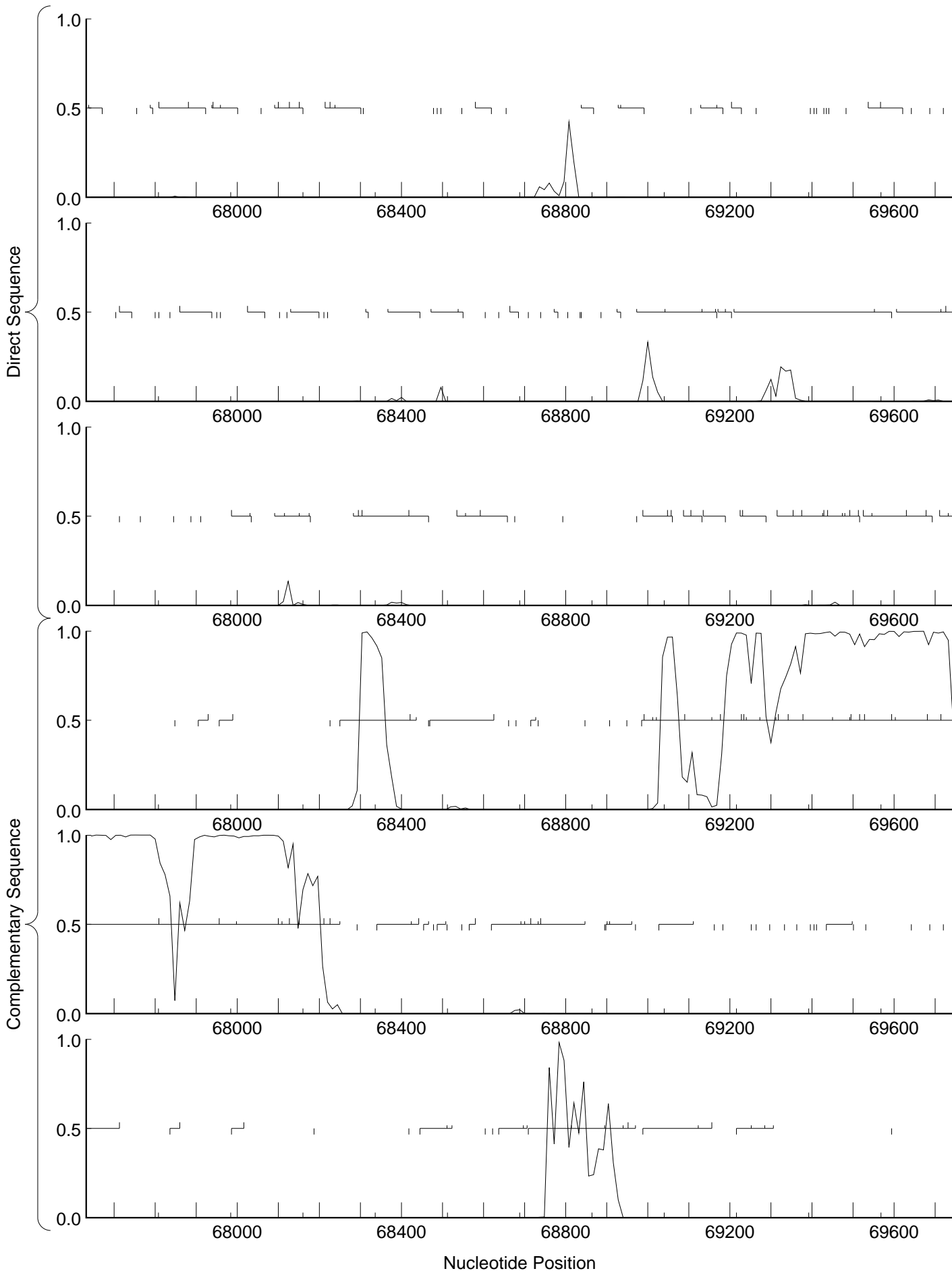
complete sequence, 76871 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 30/38











complete sequence, 76871 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 35/38

