

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Ellson complete sequence, 75945 bp including 10-base 3' overhang (TCGAT
Analysis Date : 4/24/25 at 5:49:00
Pages : 37
Sequence Length : 75945 bp
GC Content : 59.35%

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,

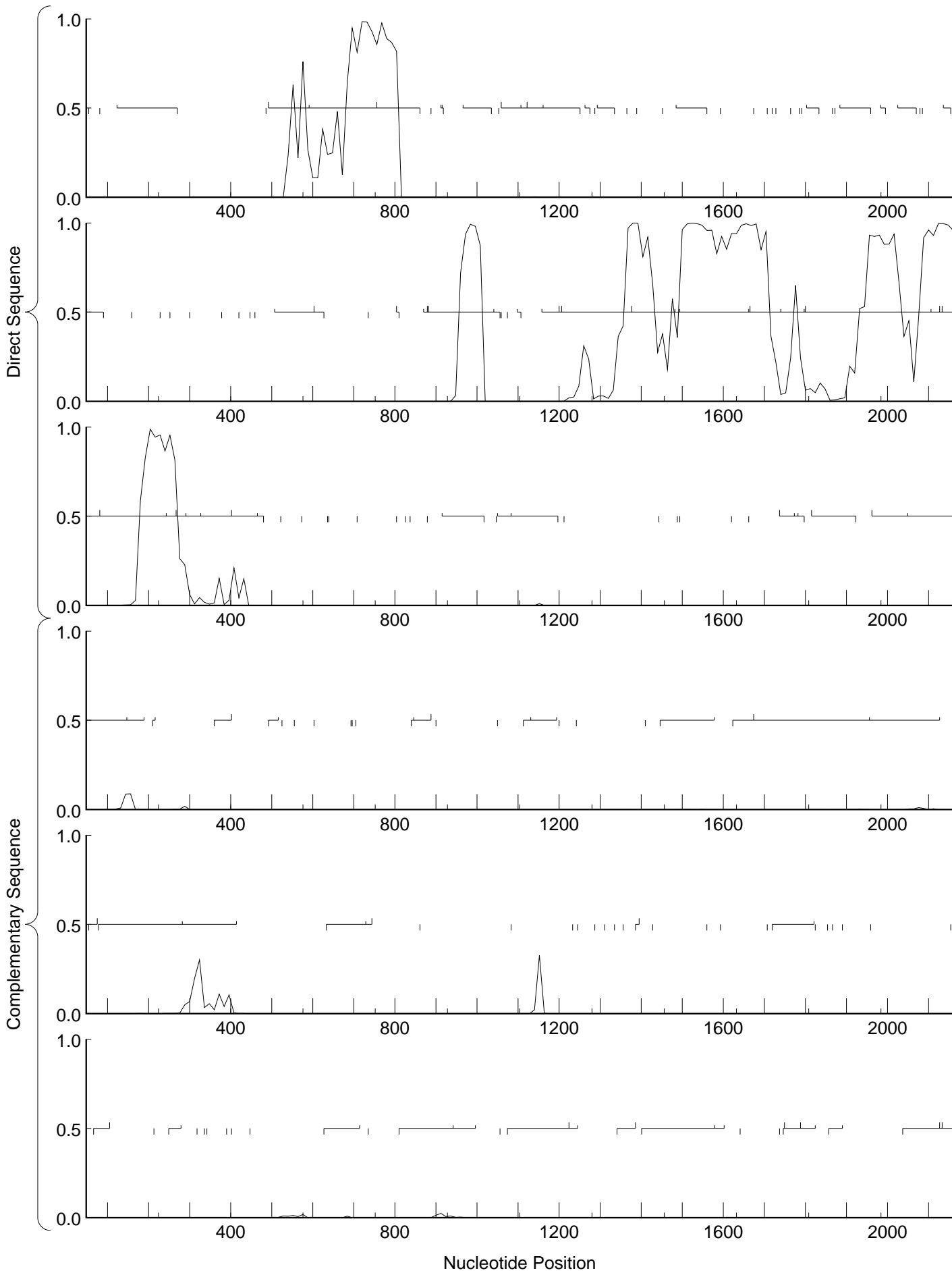
Matrix notes & comments

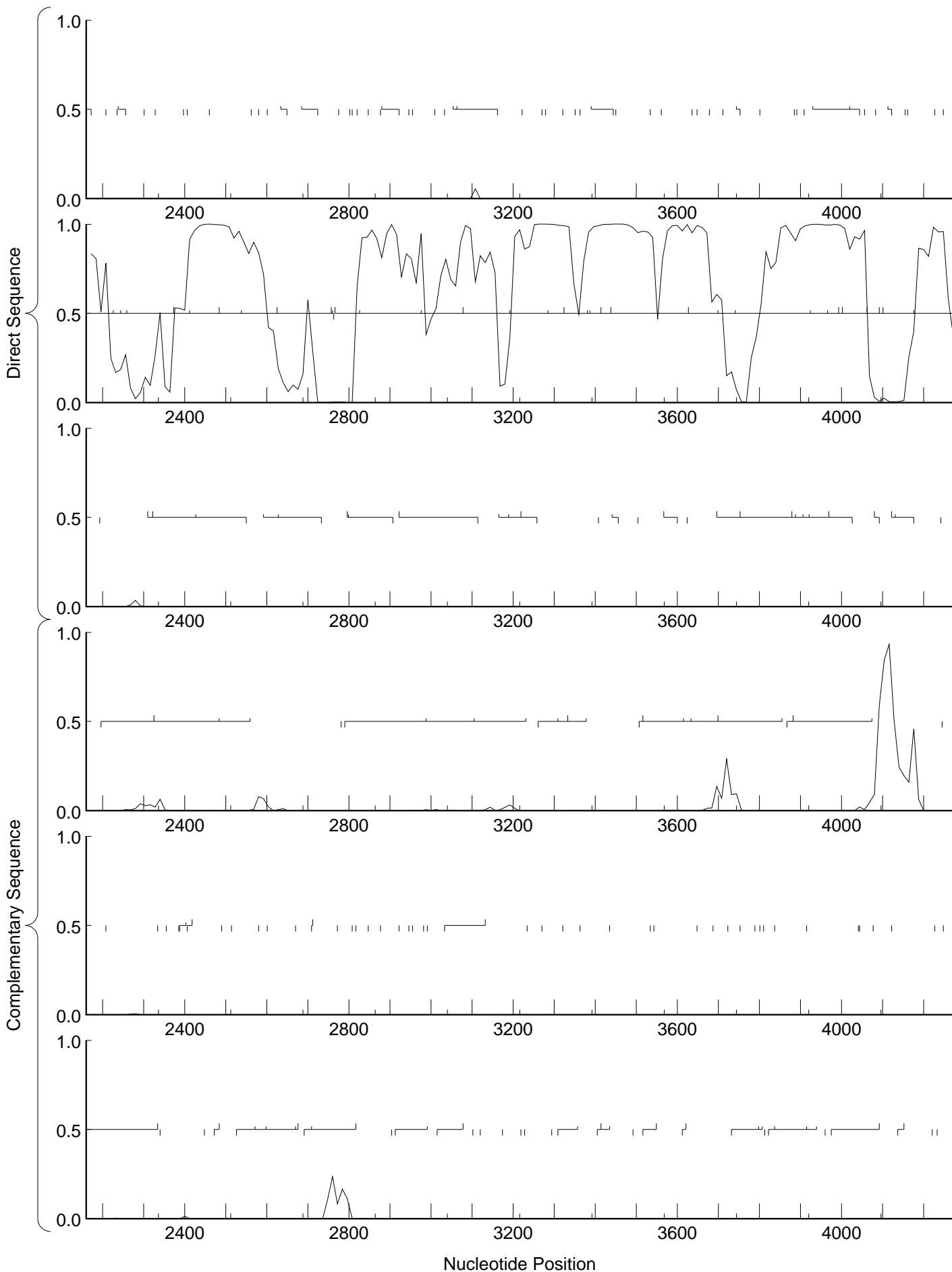
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

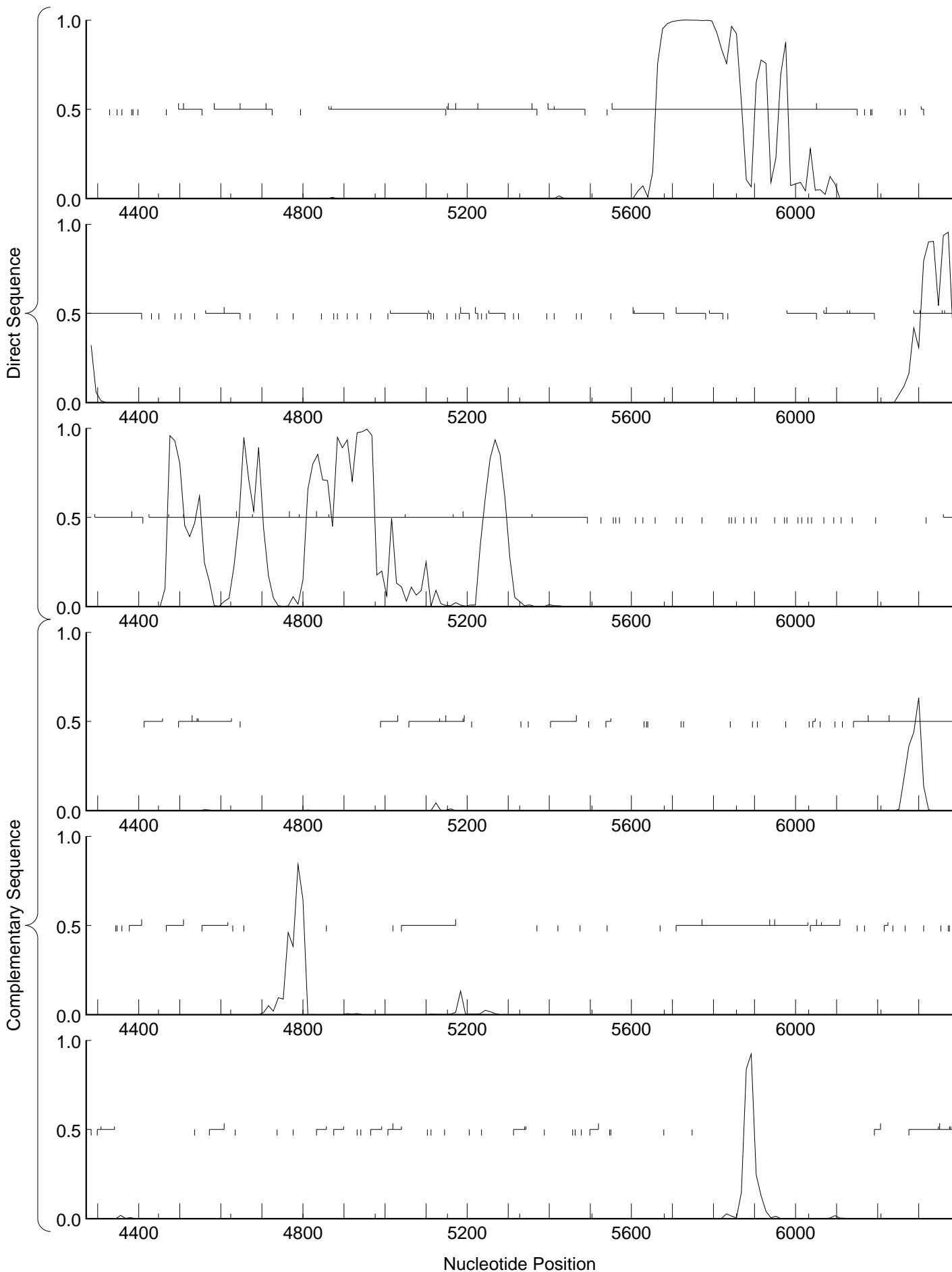
MATRIX INFORMATION

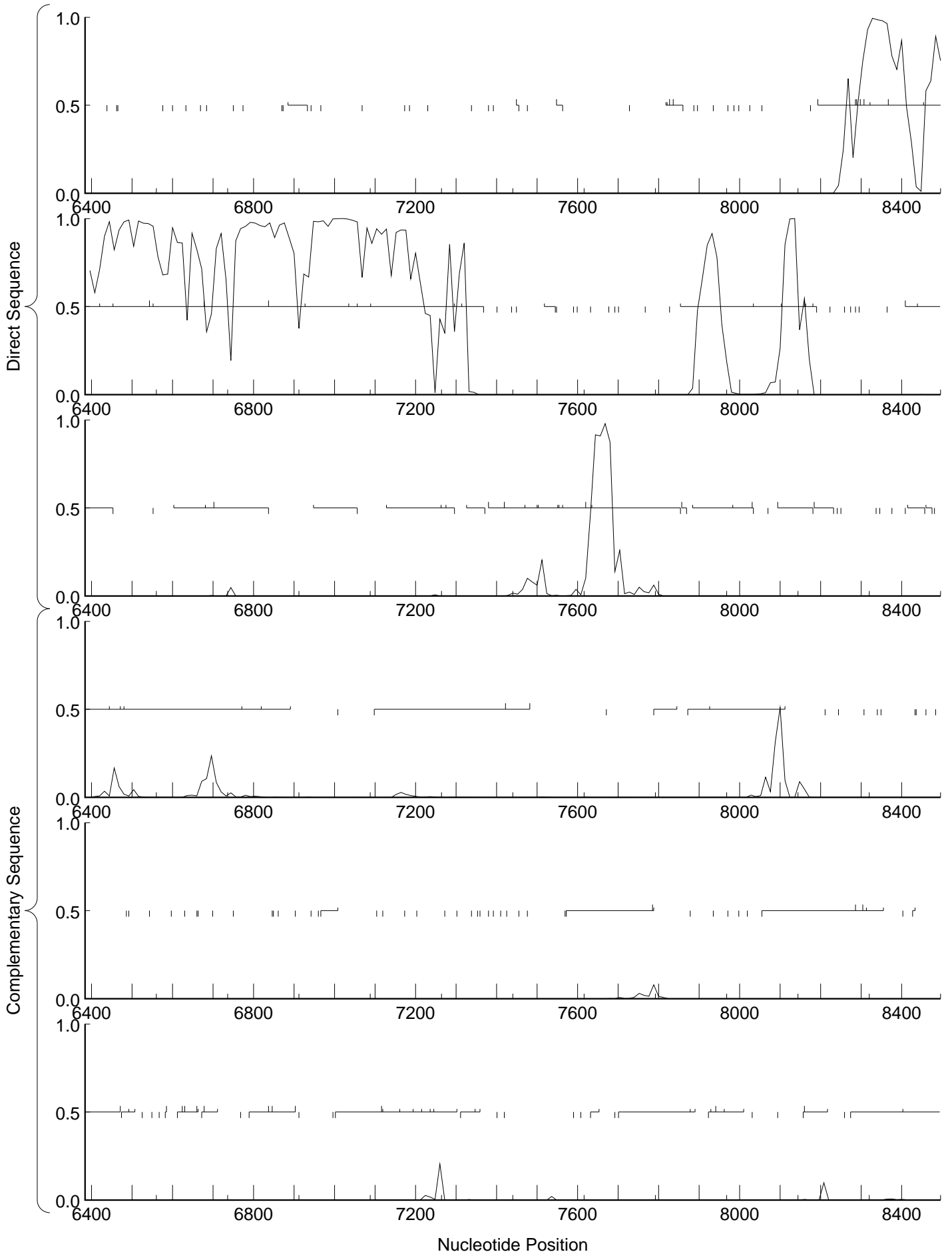
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

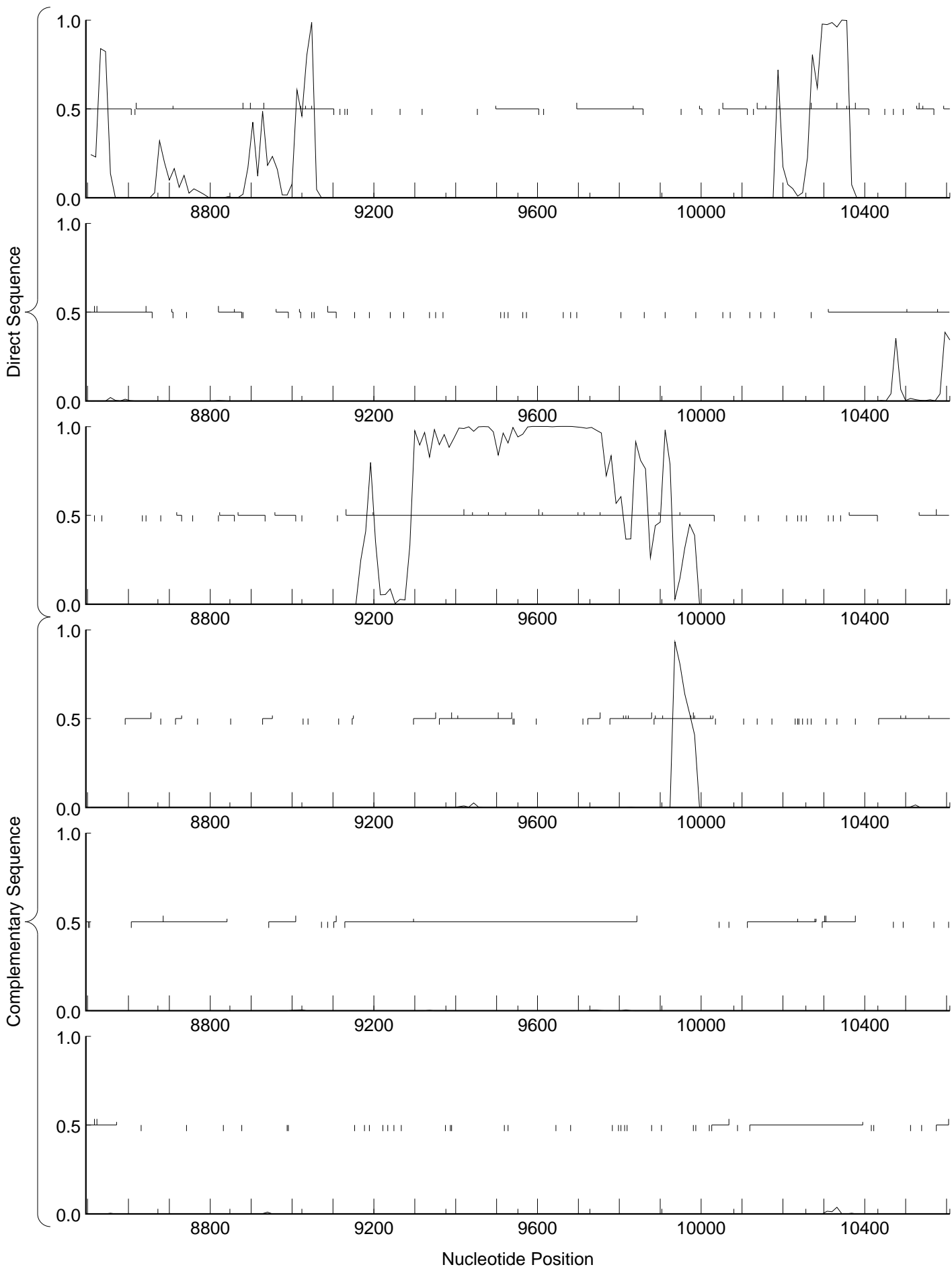
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

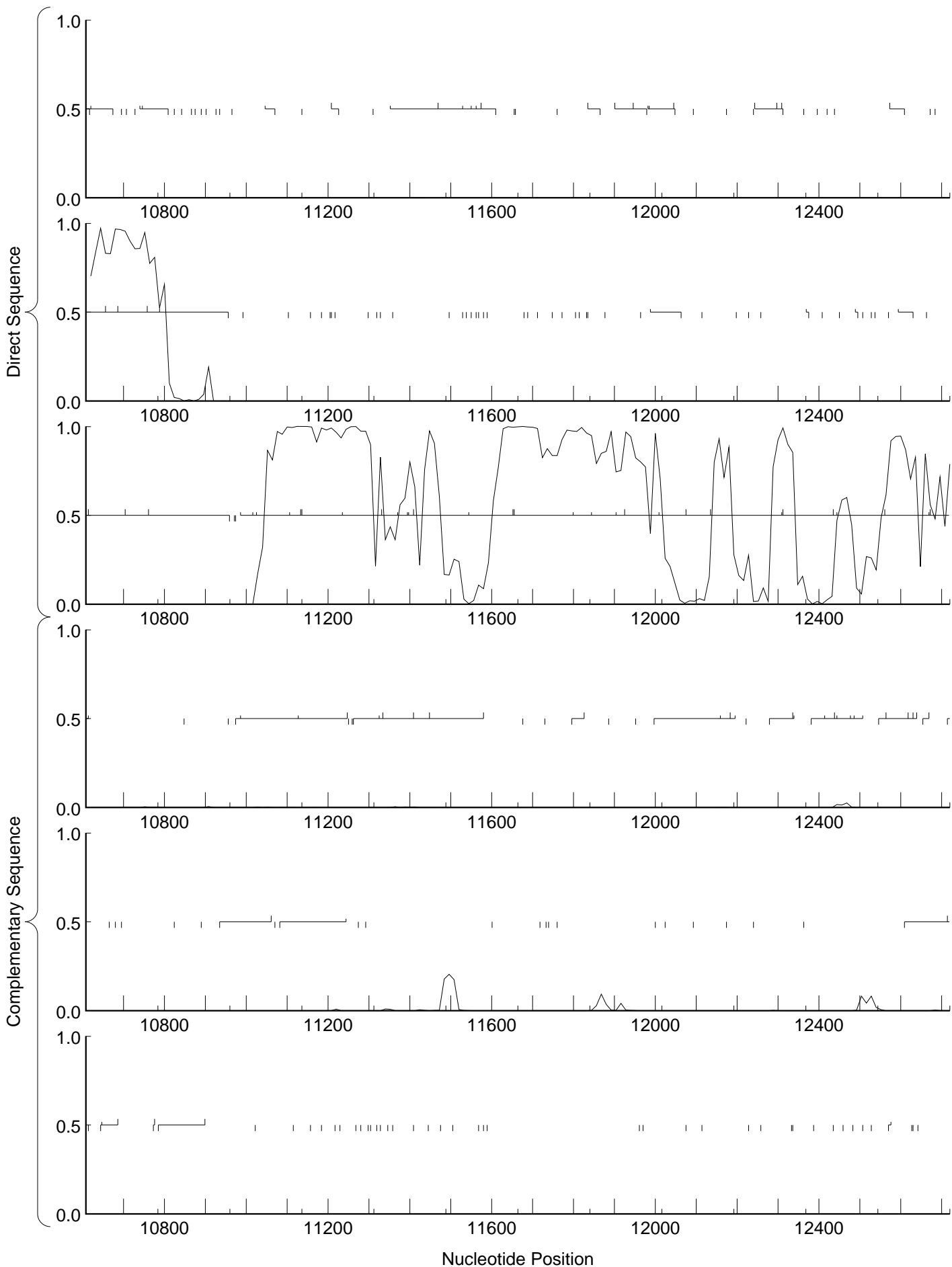


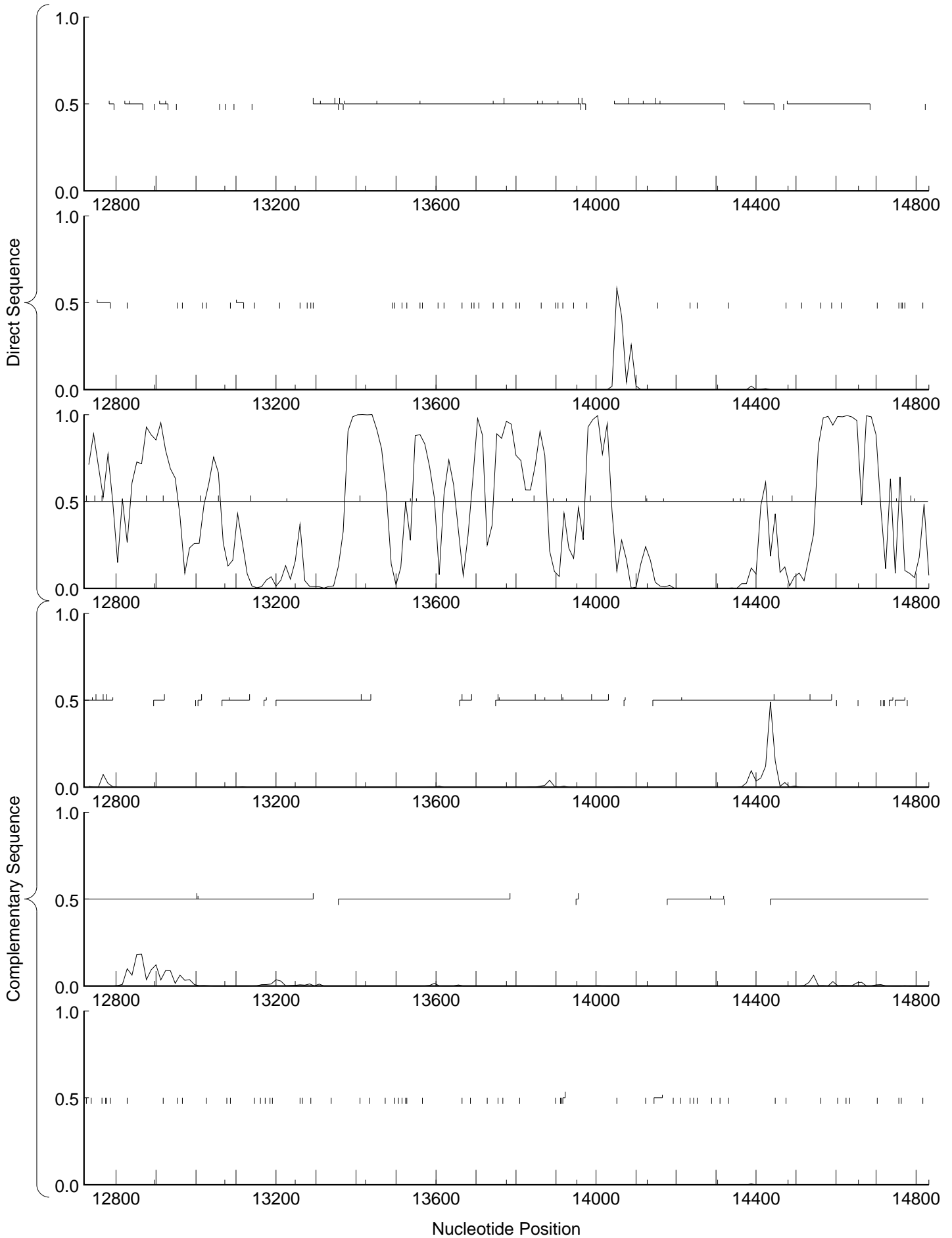


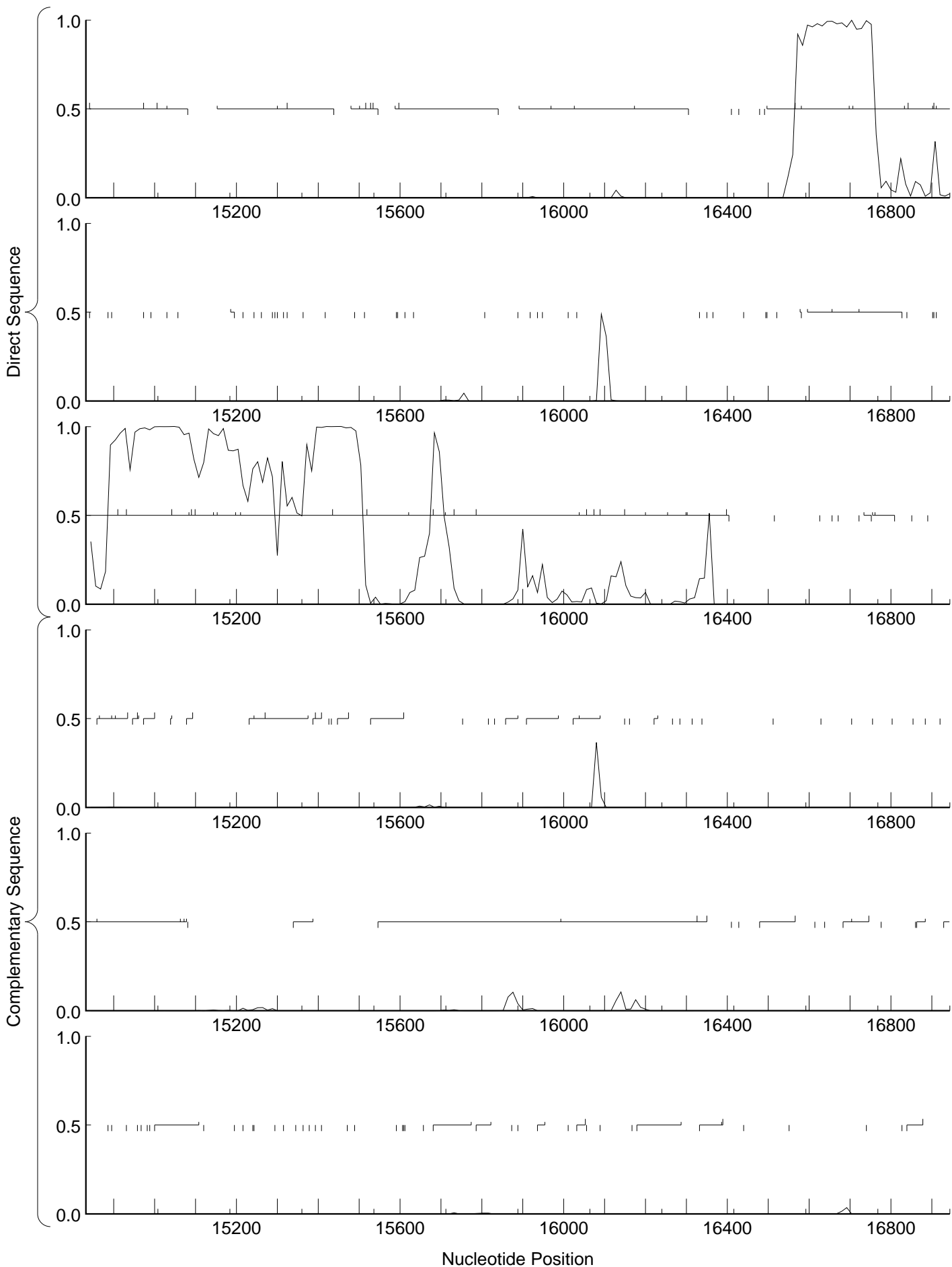


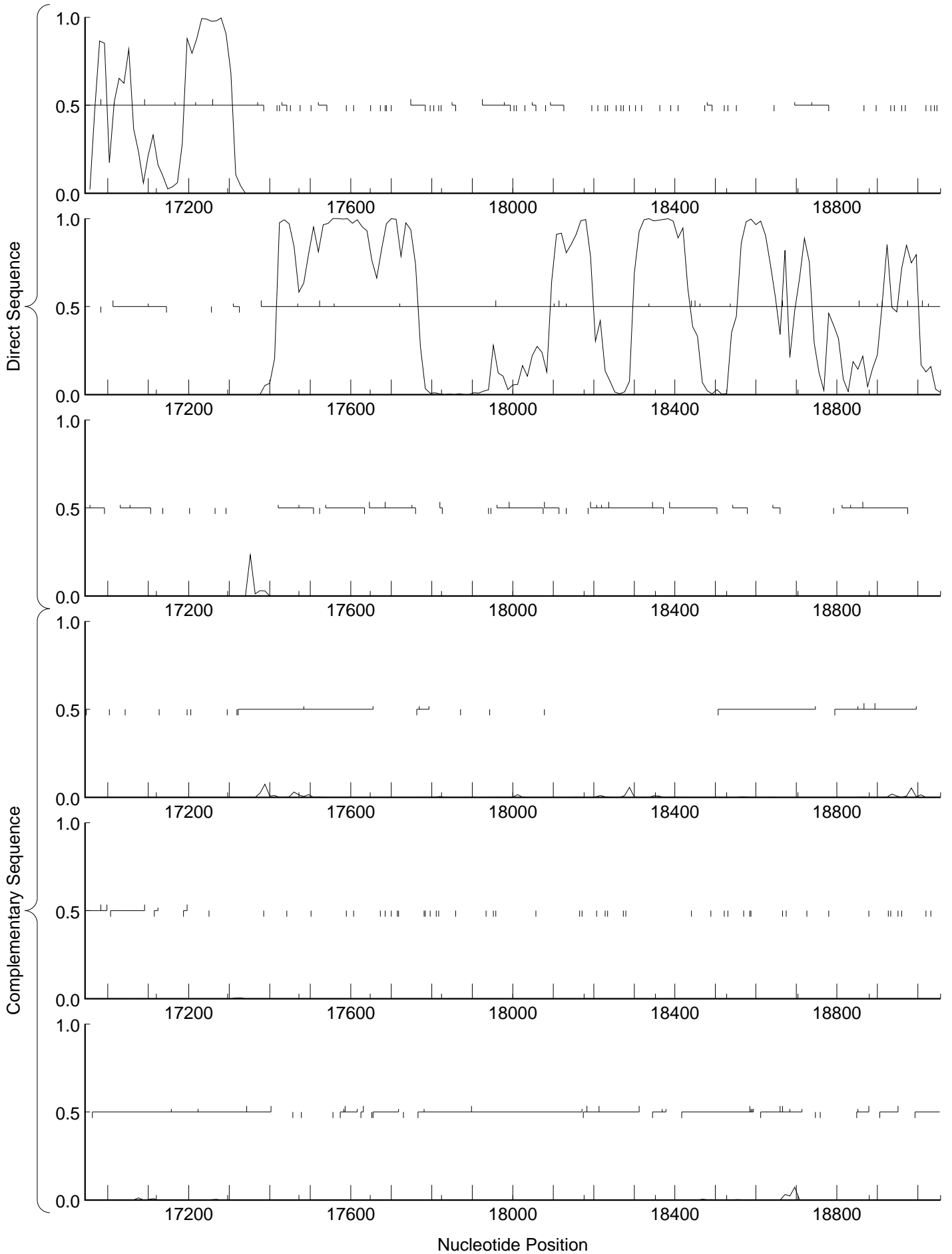


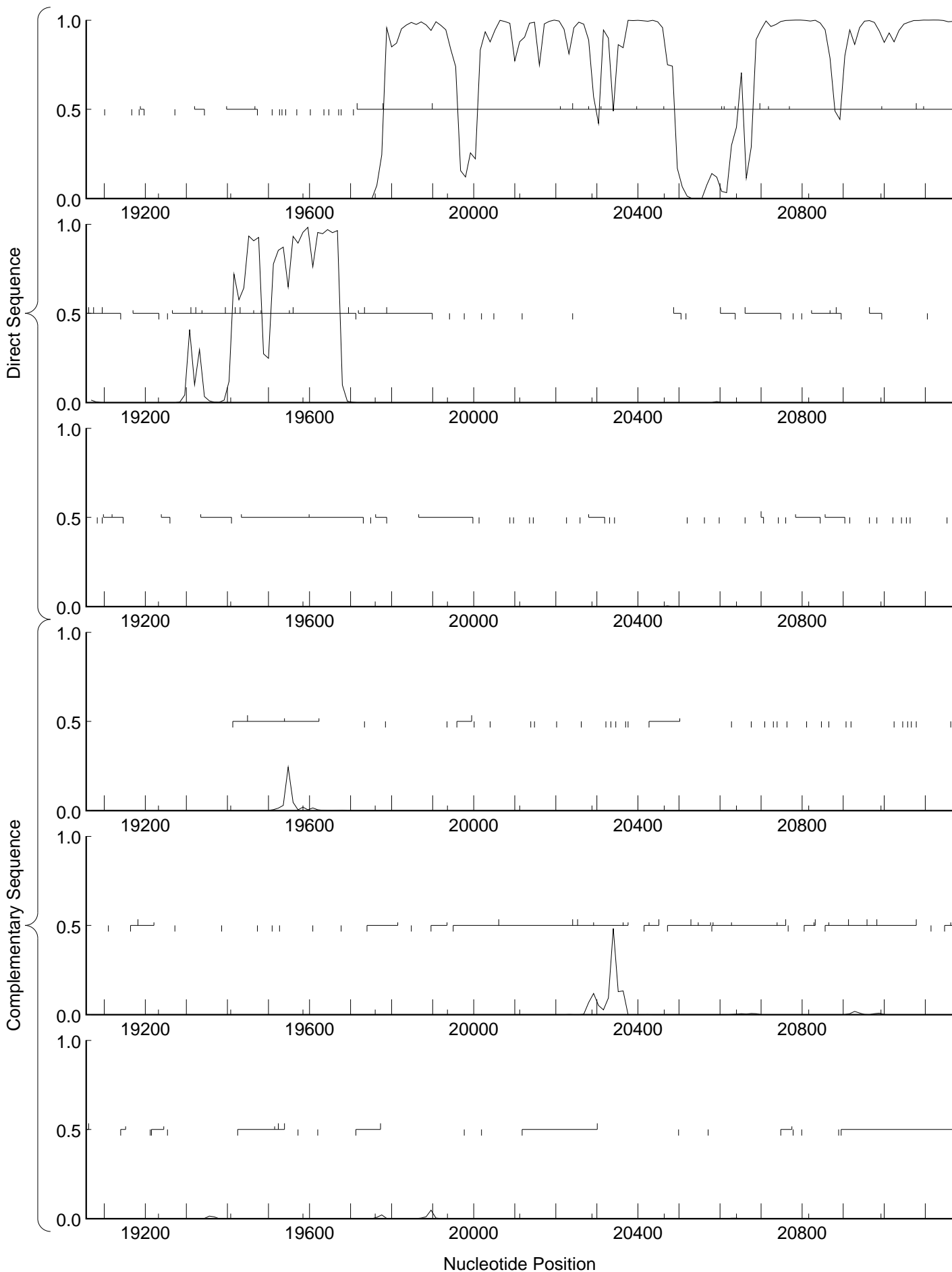


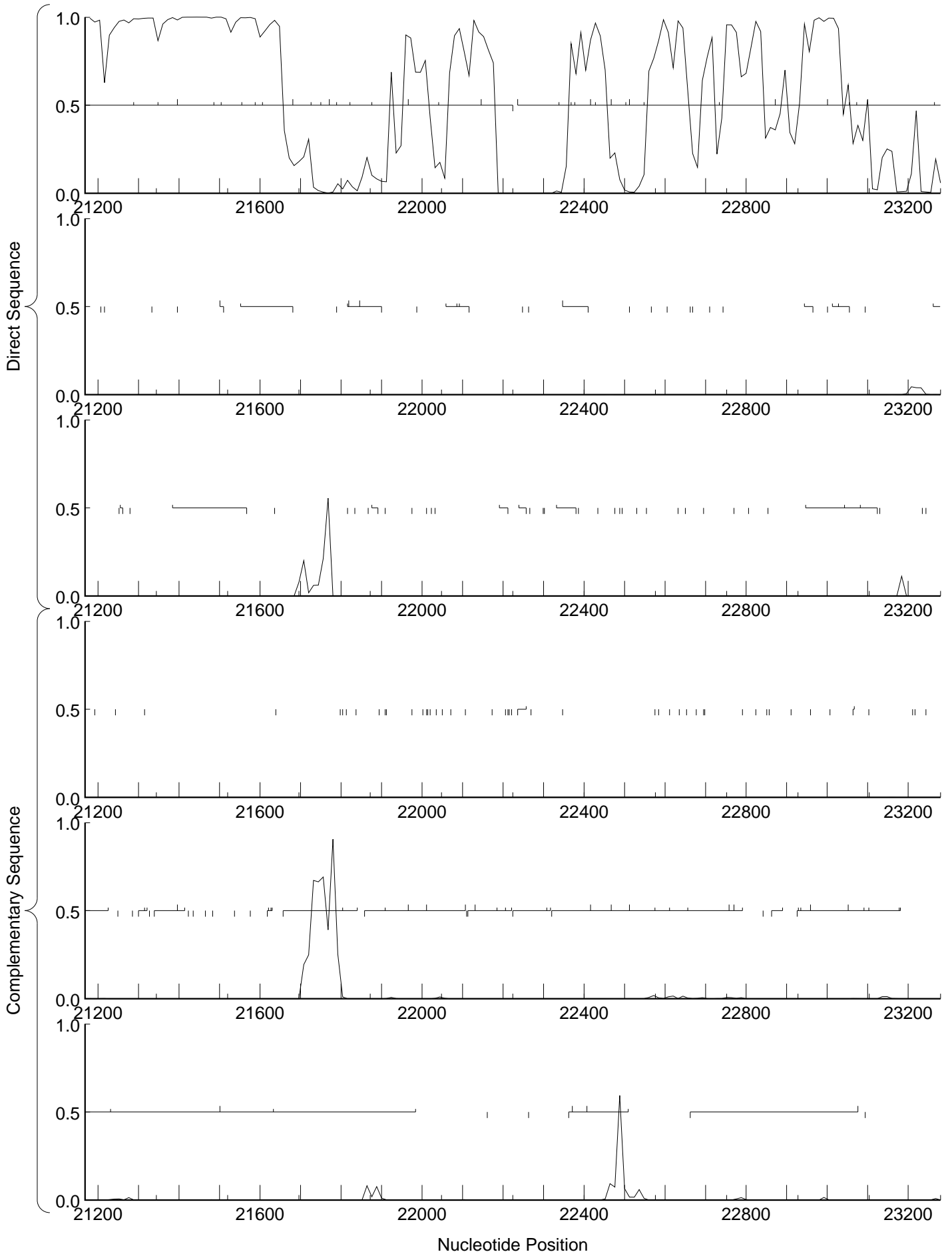


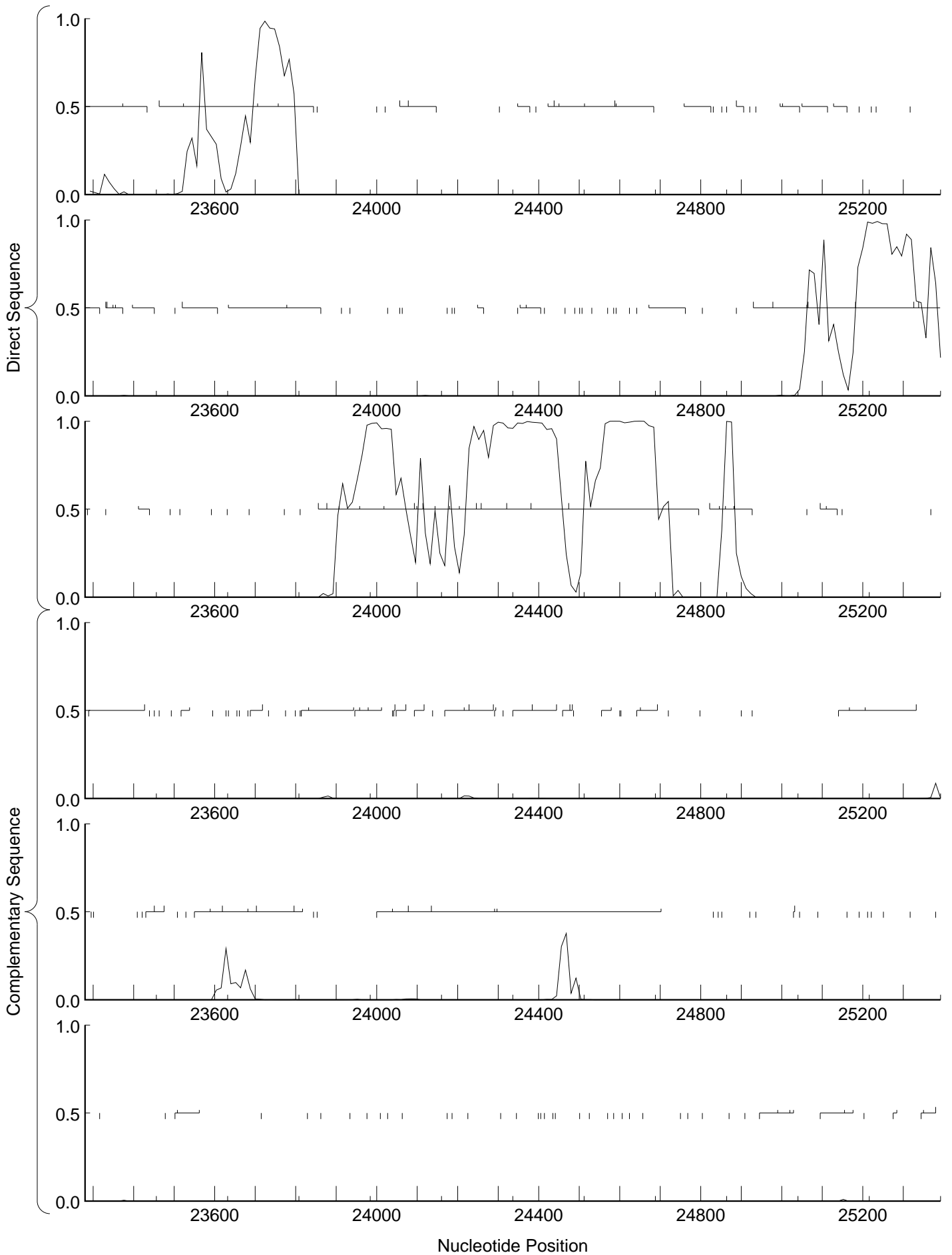




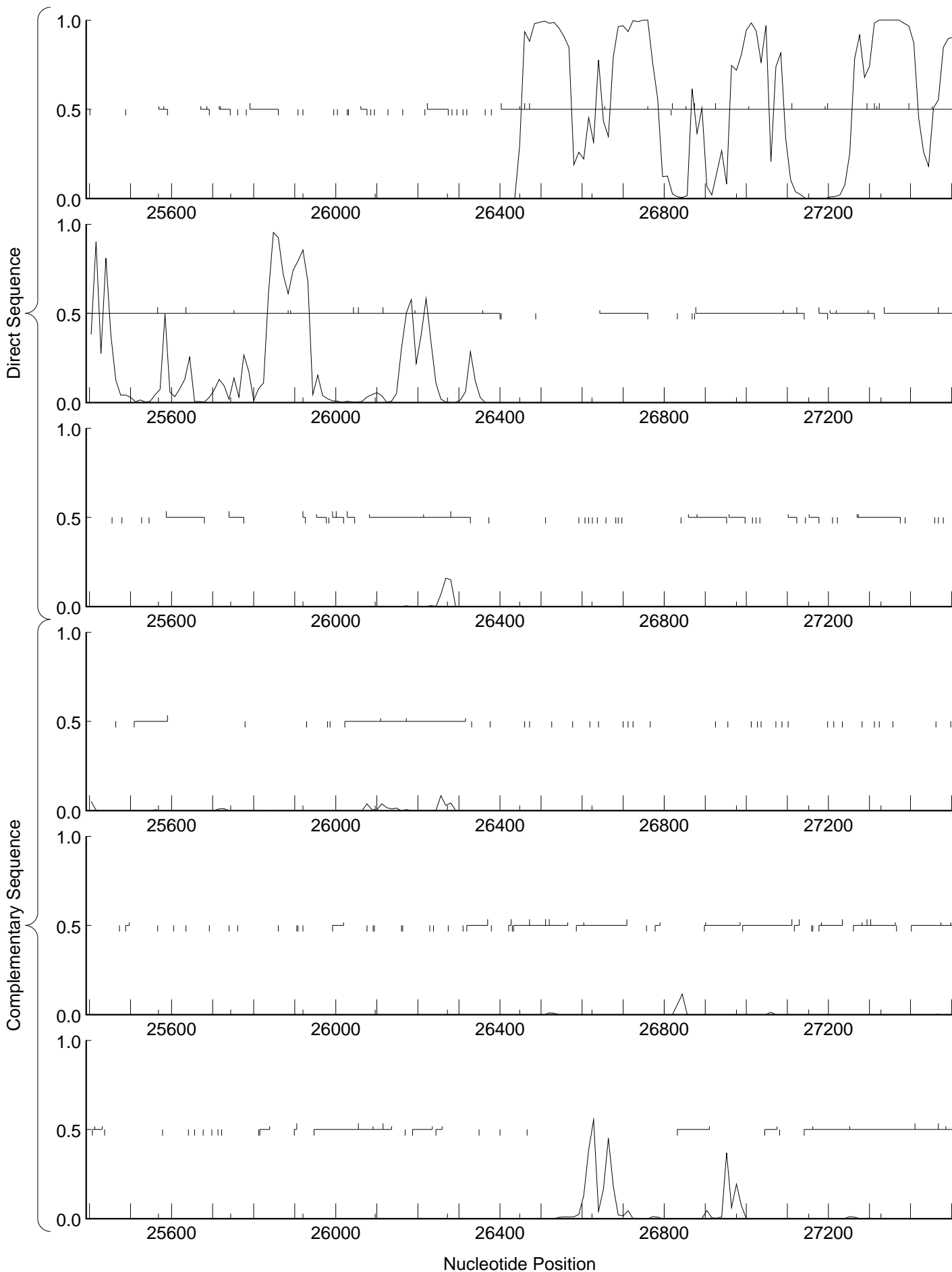


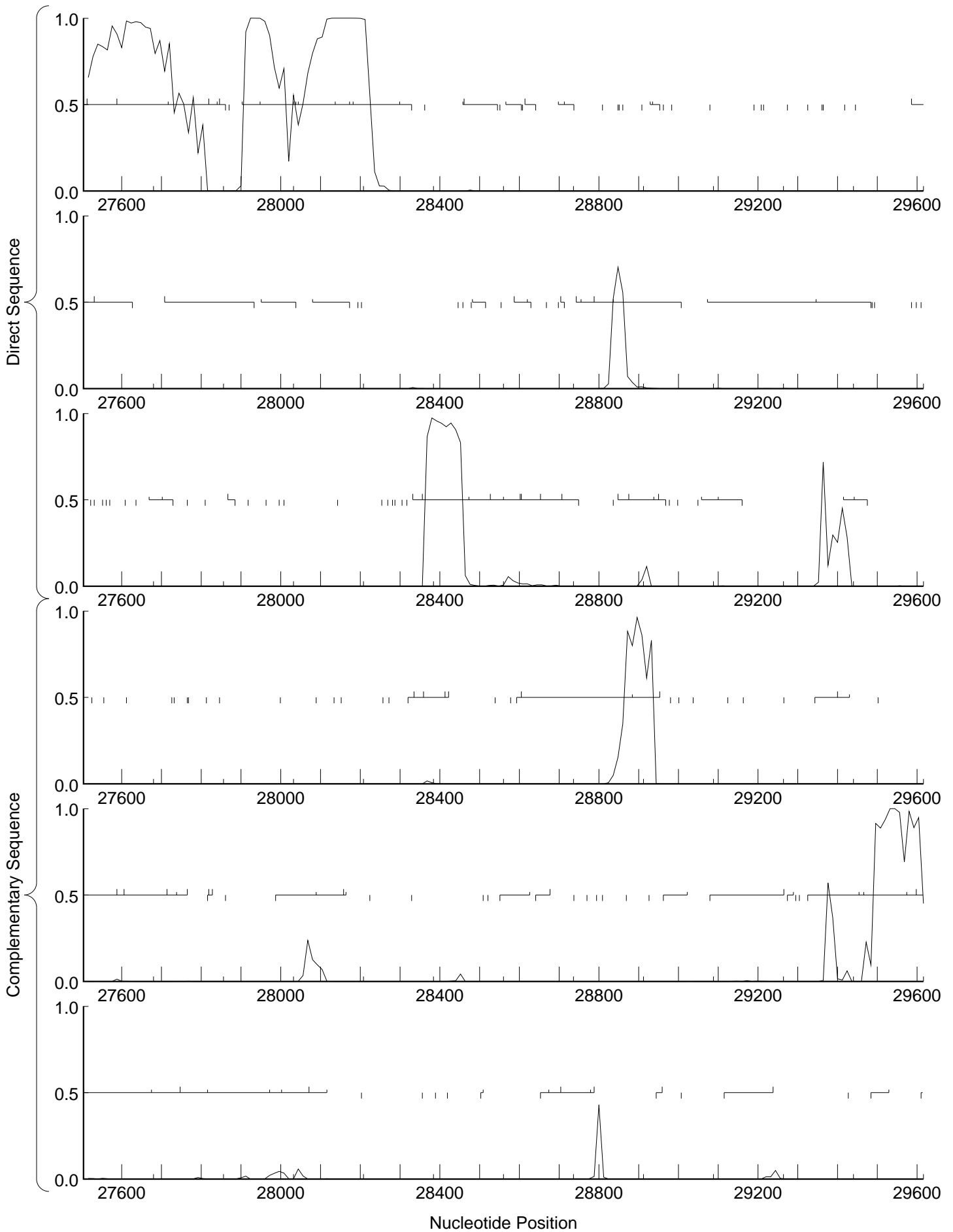


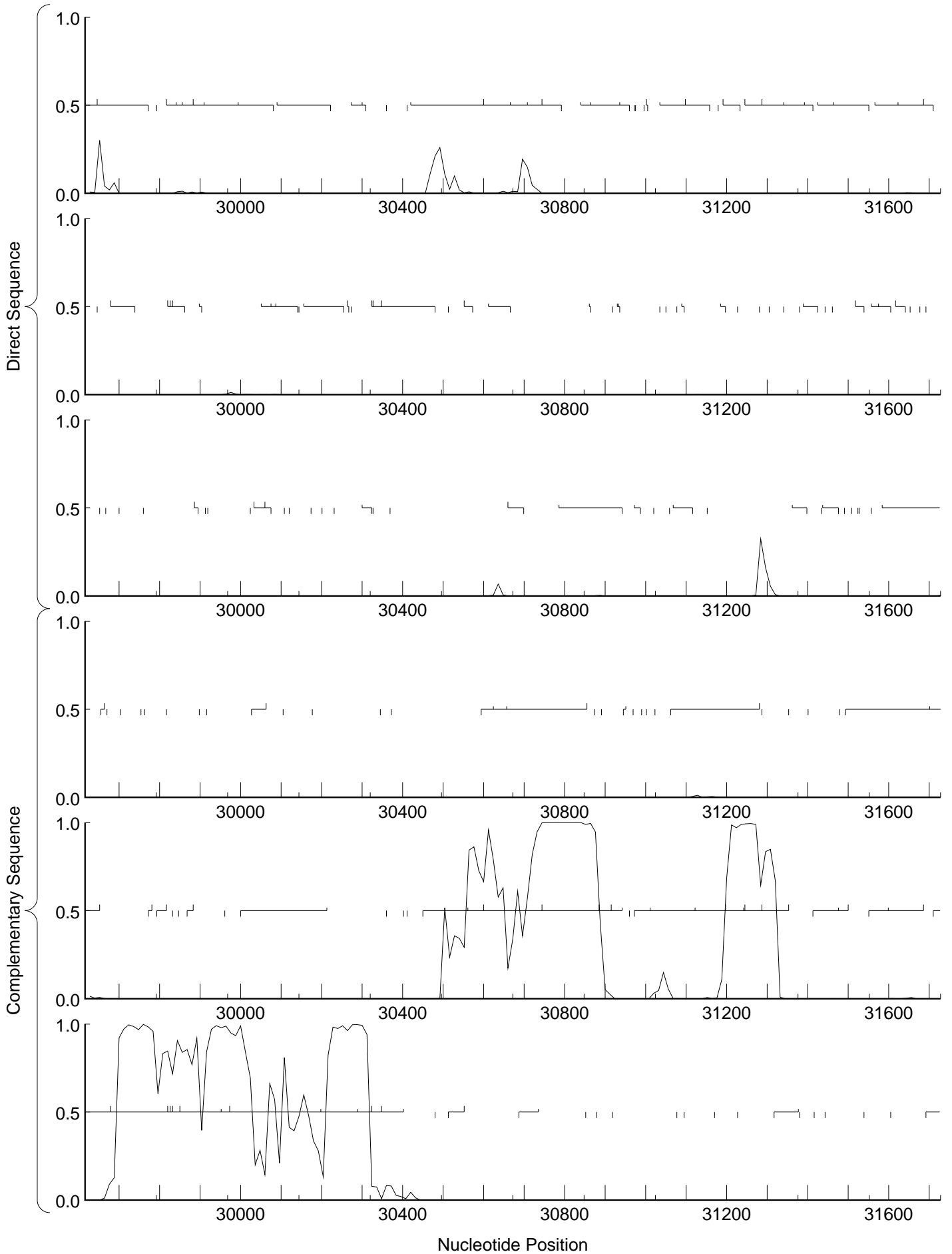


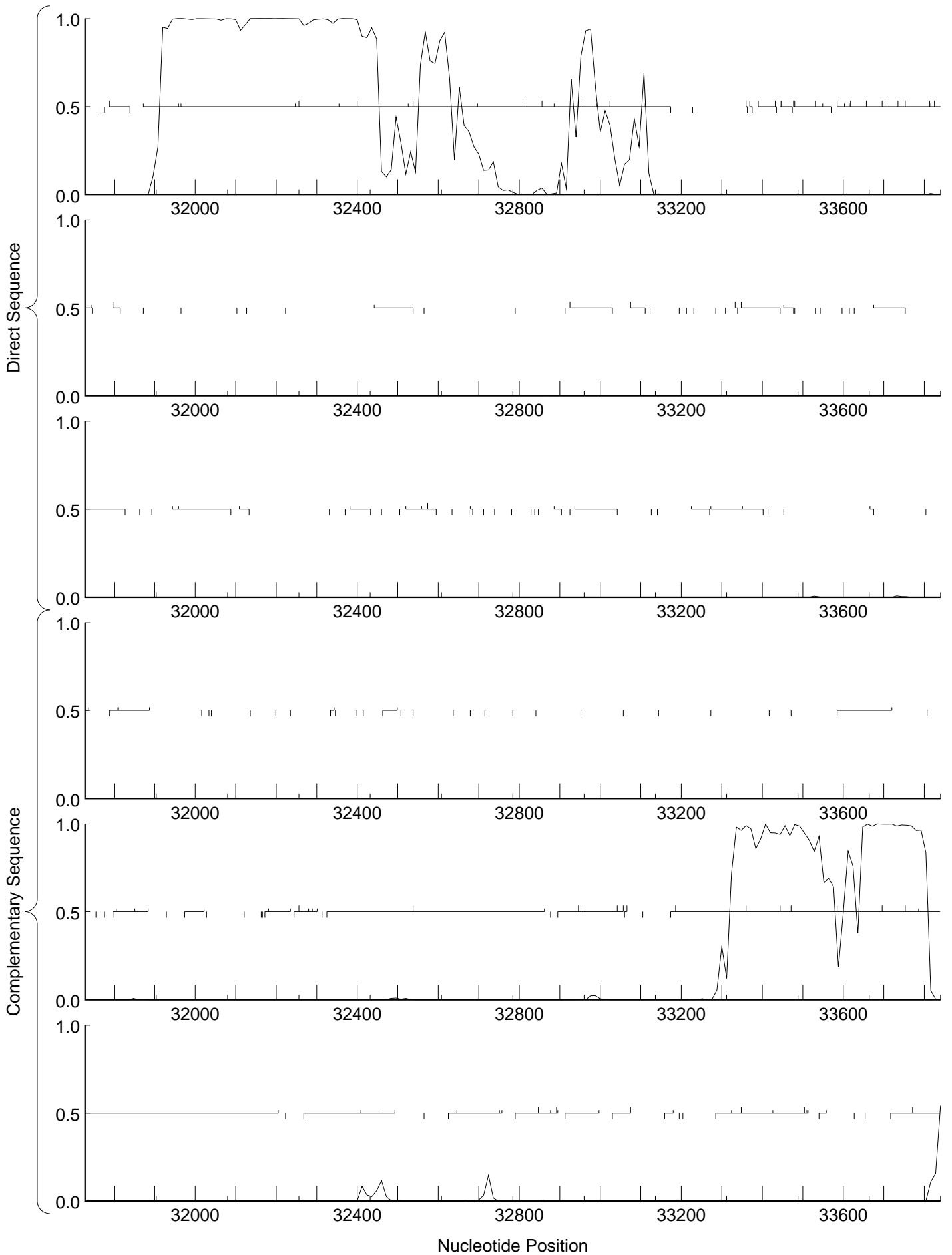


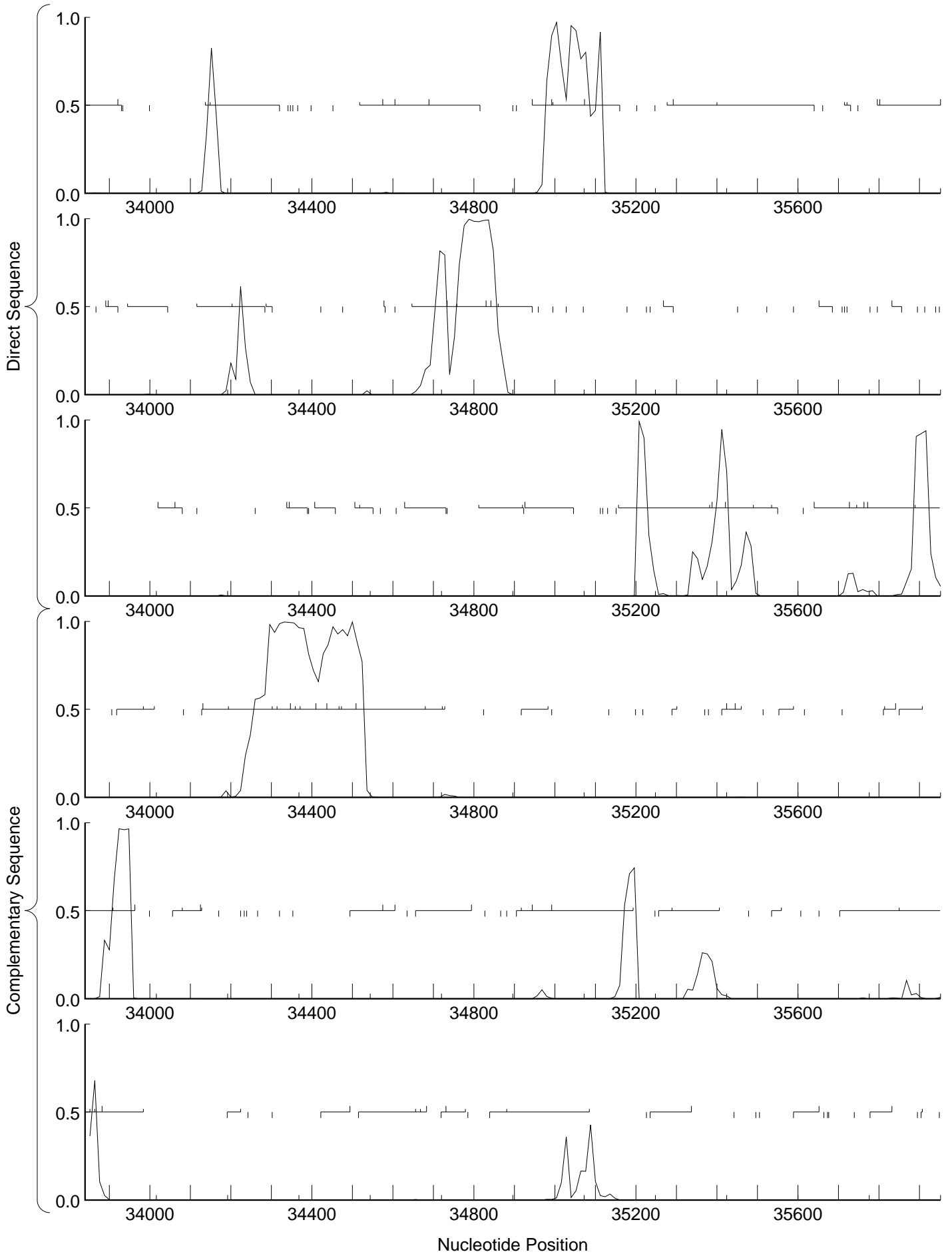
Complete sequence, 75945 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L3, Order 4, Window 96, Step 12, 14/37











Complete sequence, 75945 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L3, Order 4, Window 96, Step 12, 19/37

