

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Bryler complete sequence, 57666 bp including 10-base 3' overhang (CTCGG
Analysis Date : 8/24/19 at 18:29:55
Pages : 29
Sequence Length : 57666 bp
GC Content : 66.10%

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,

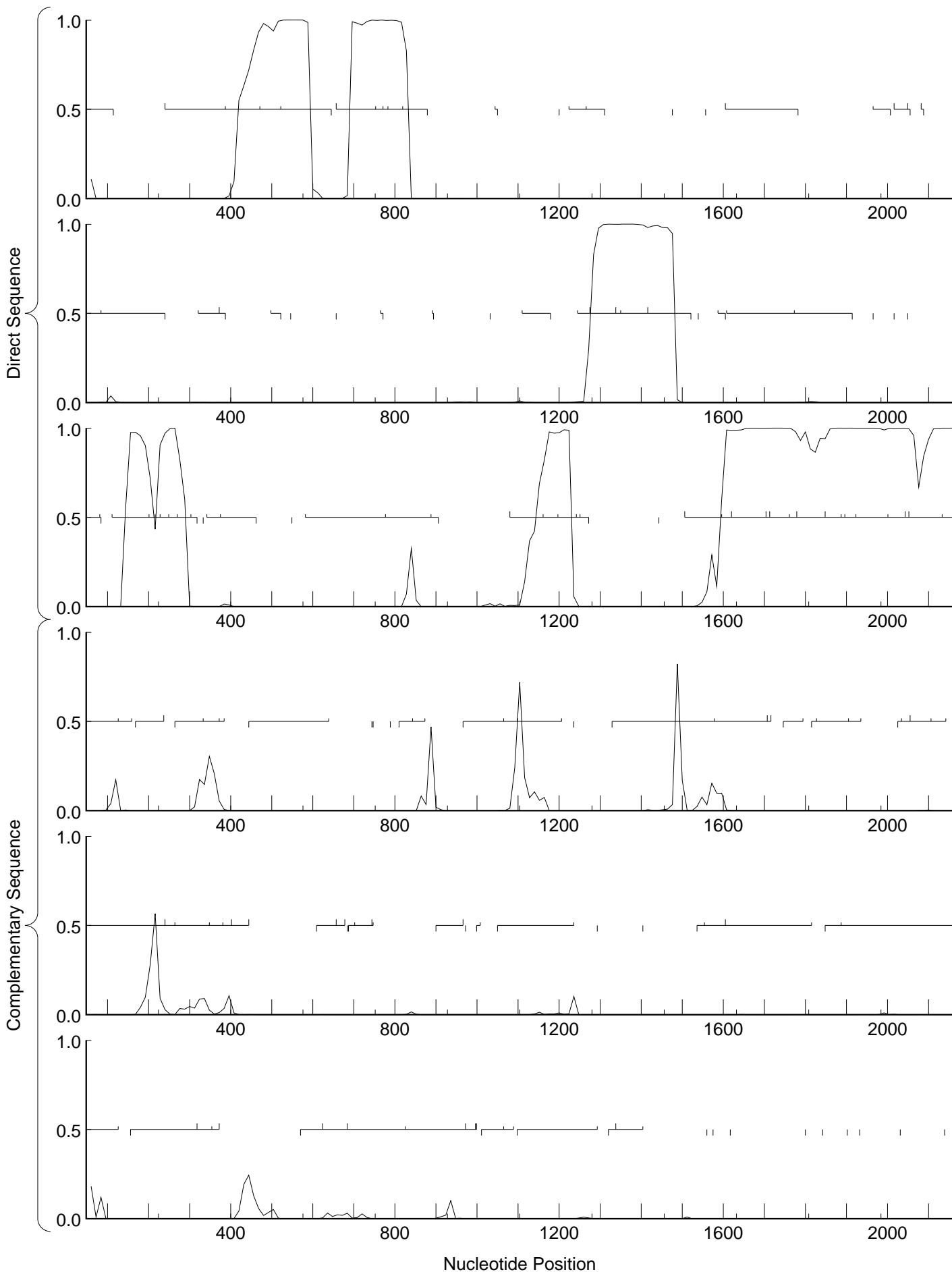
Matrix notes & comments

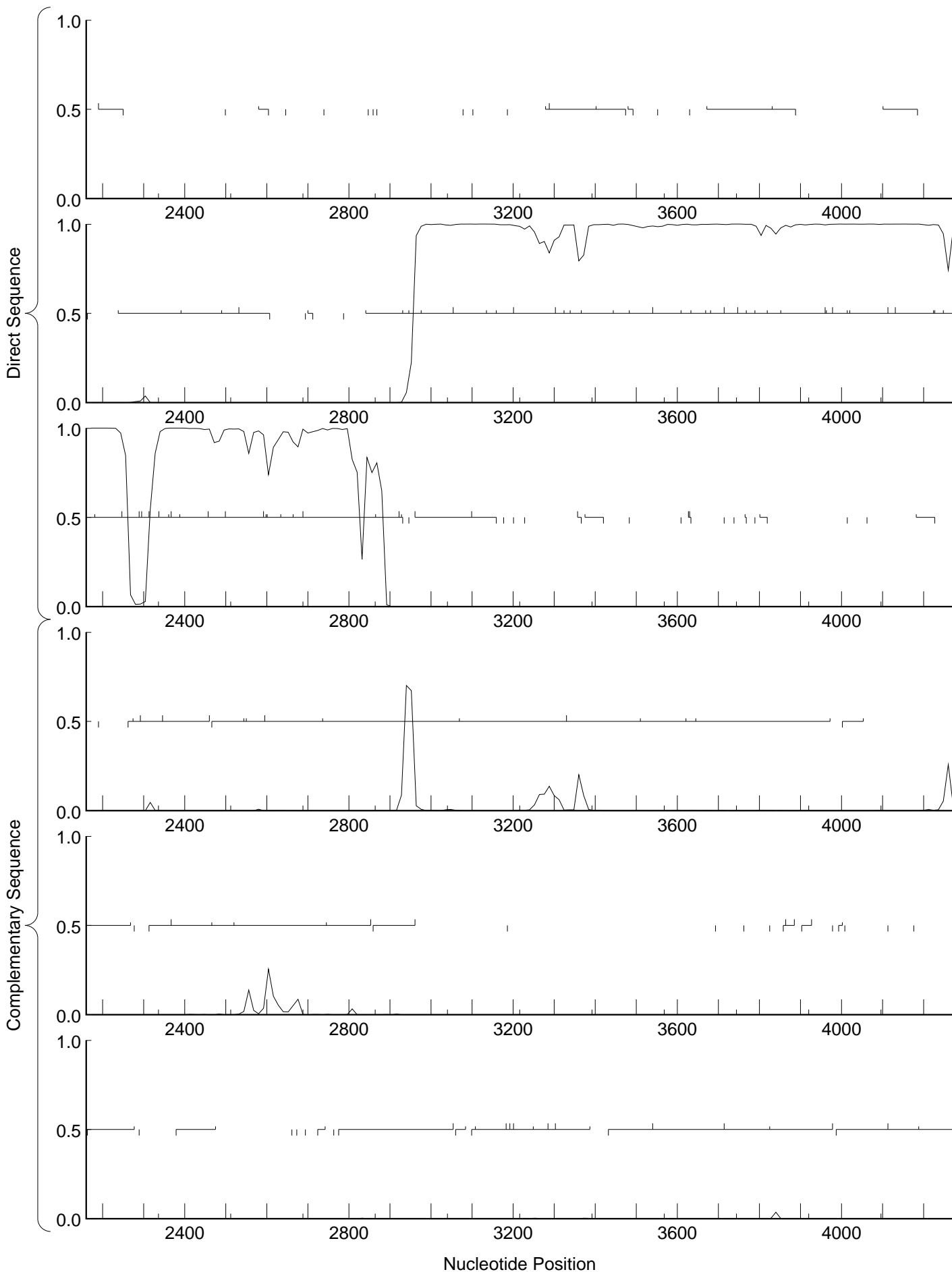
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

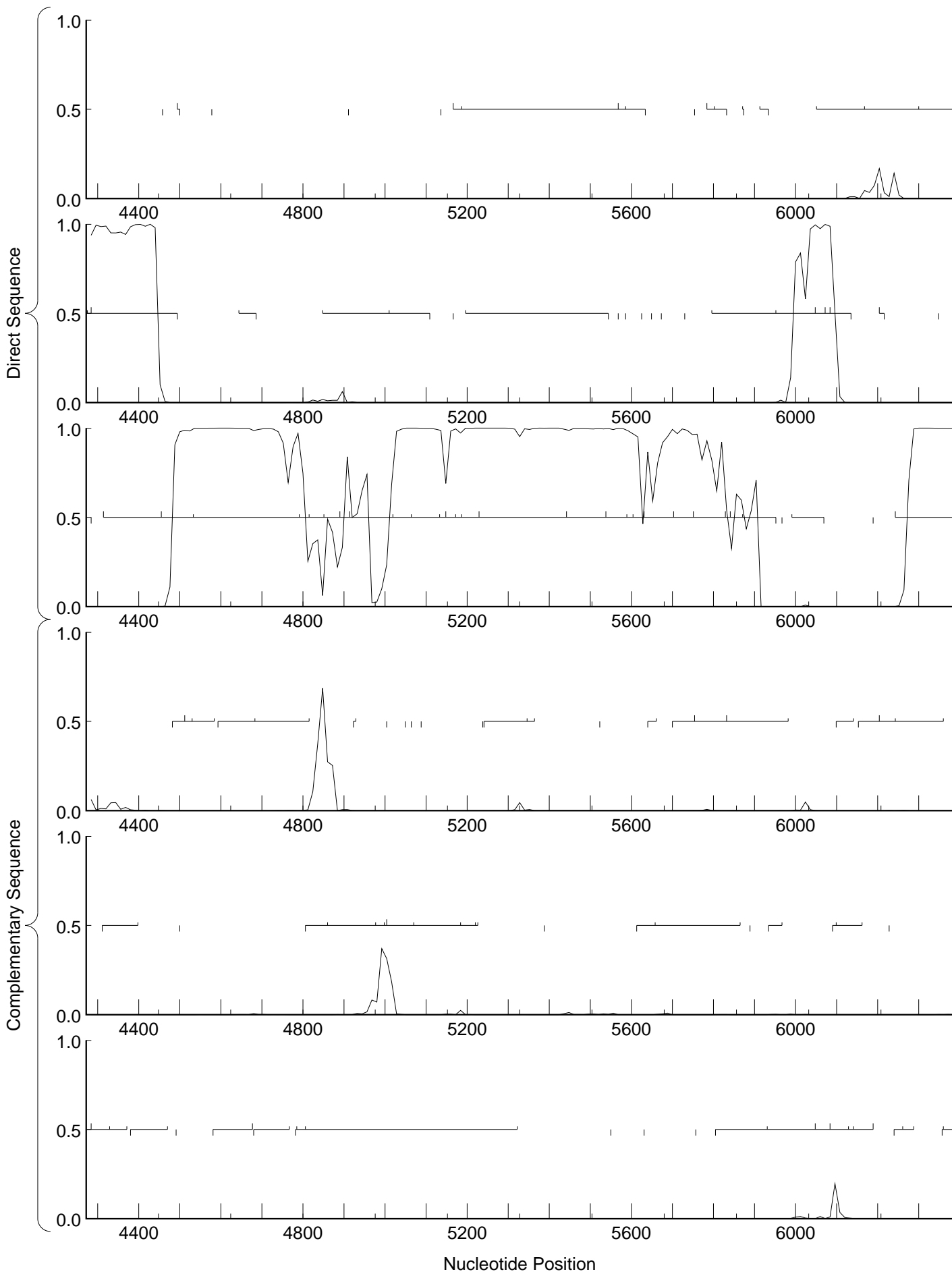
MATRIX INFORMATION

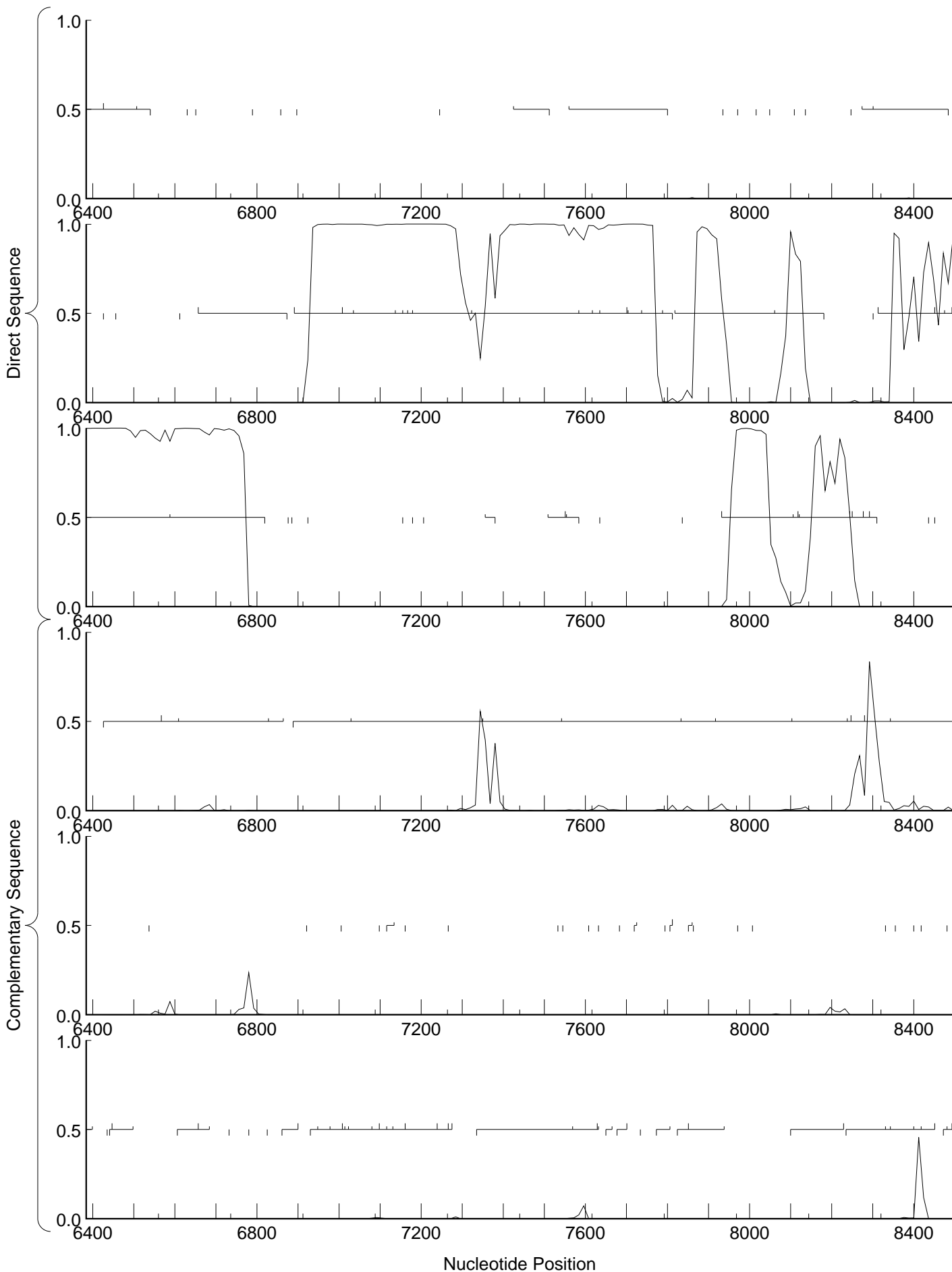
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

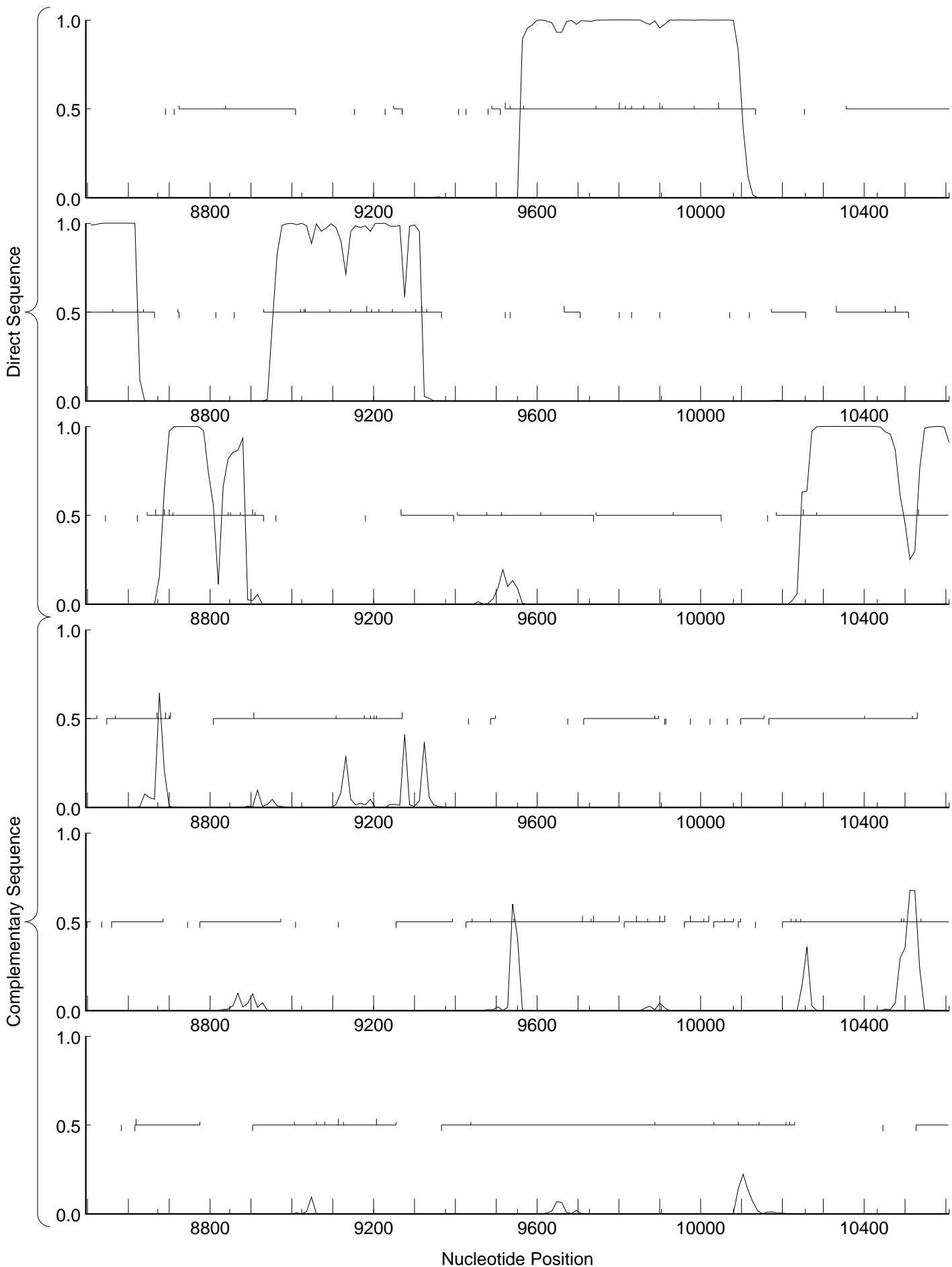
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

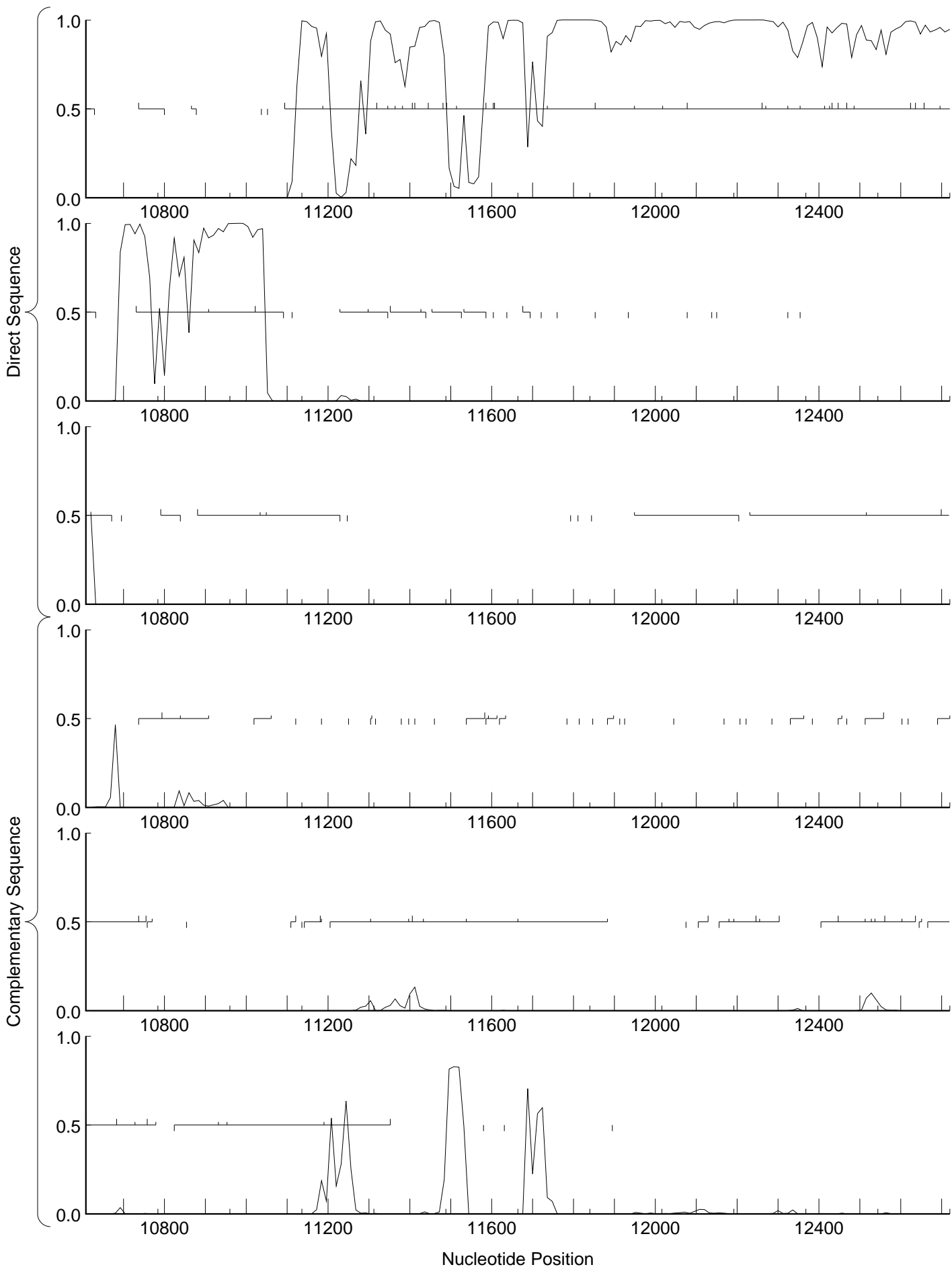


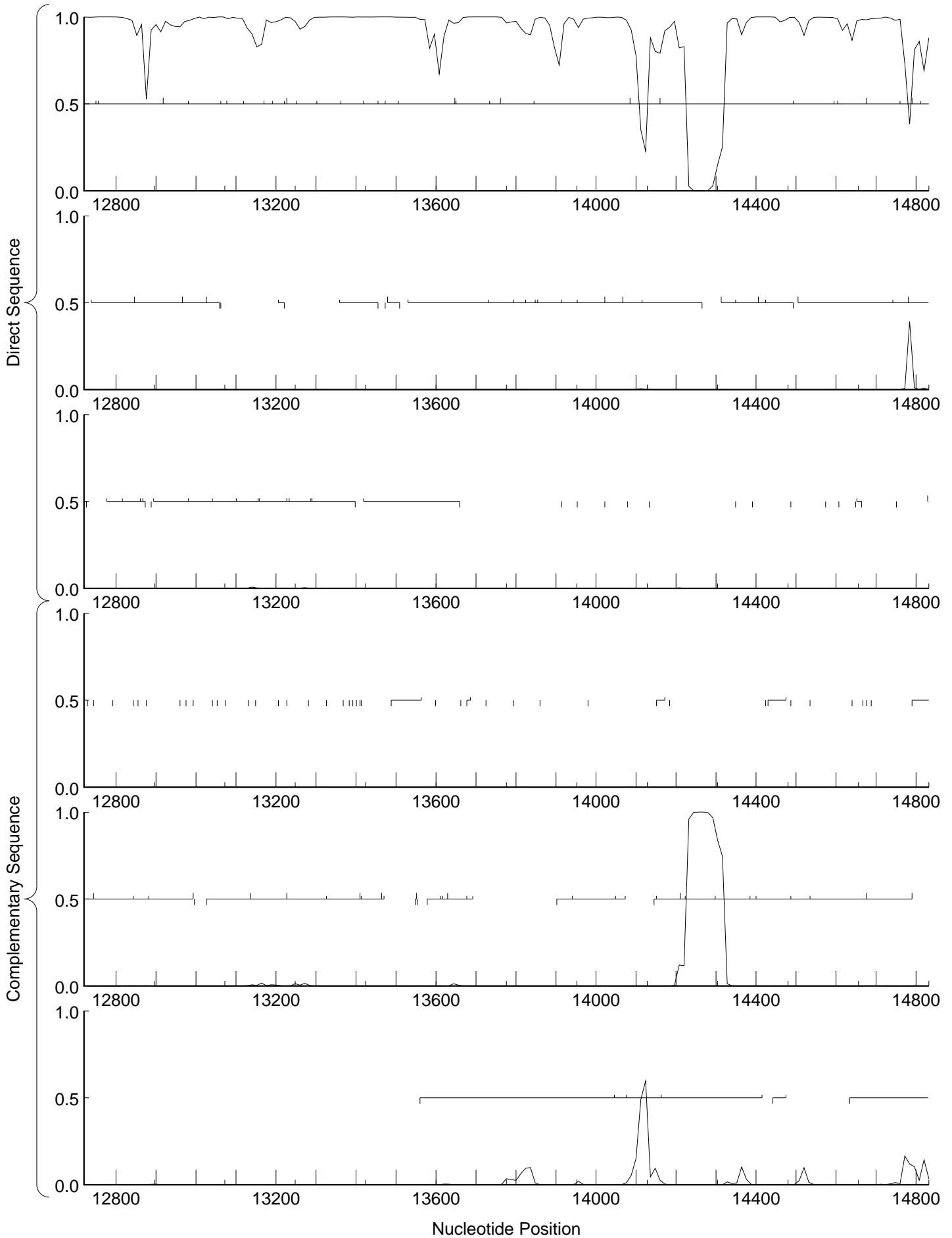


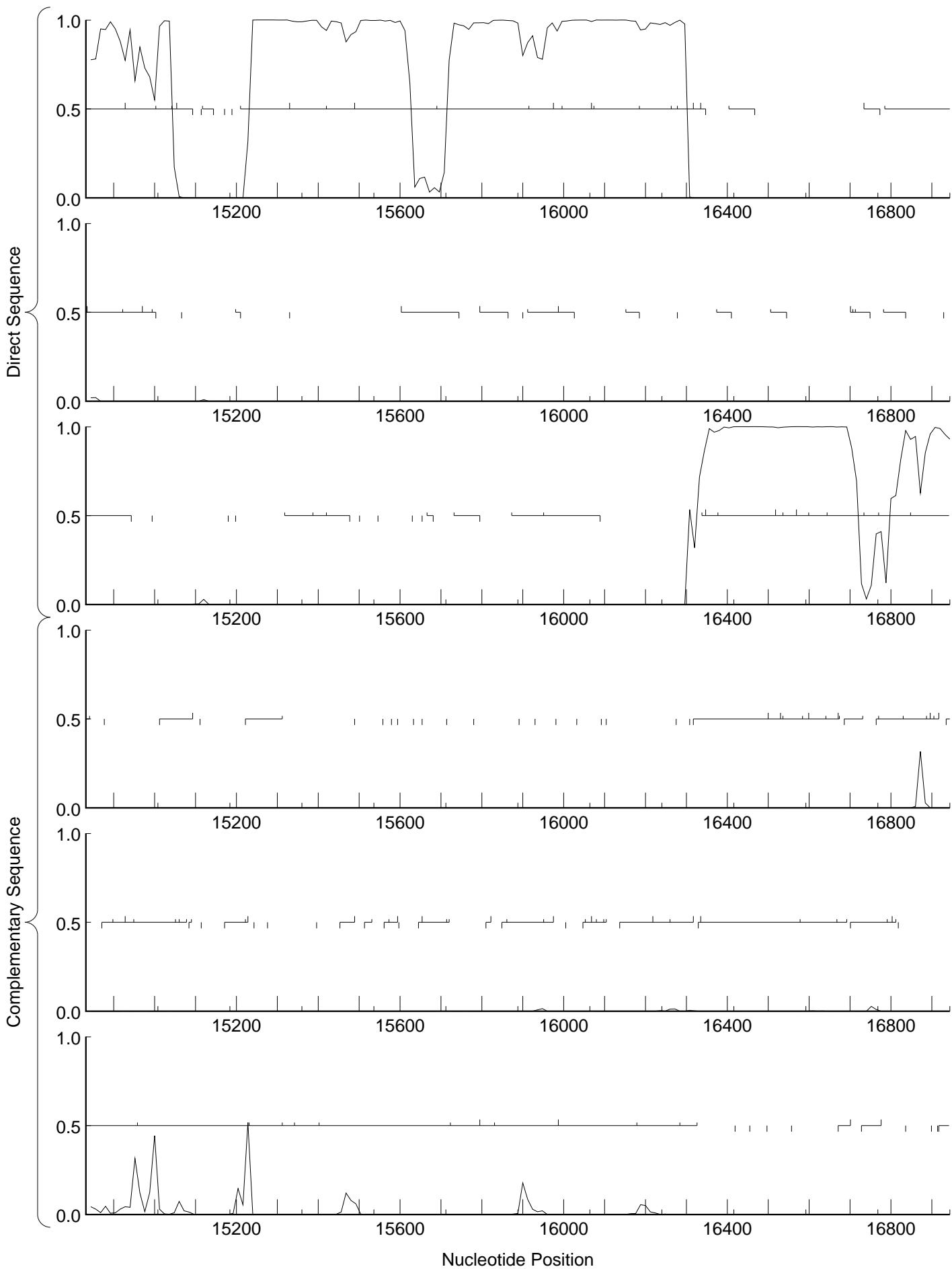


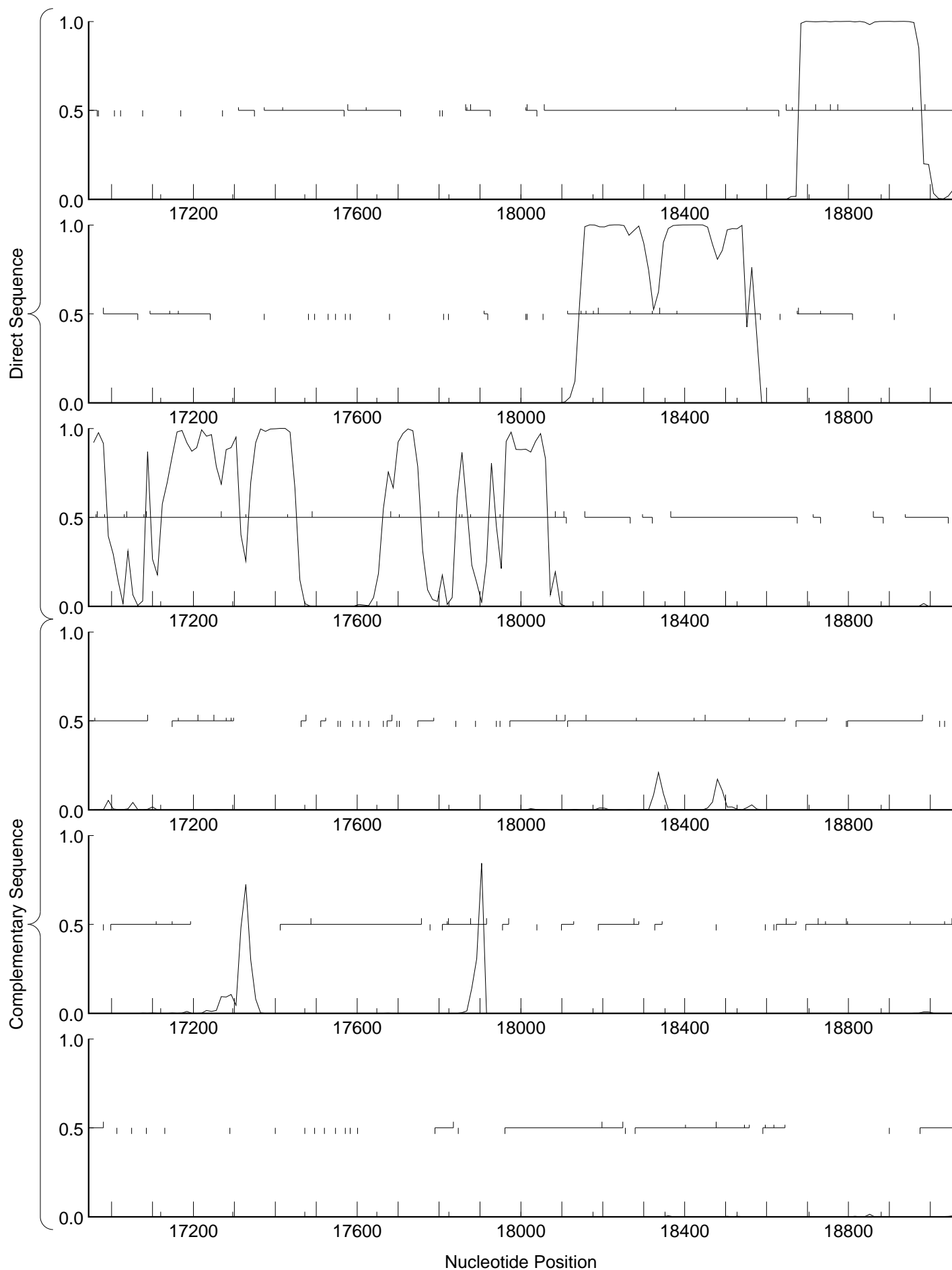


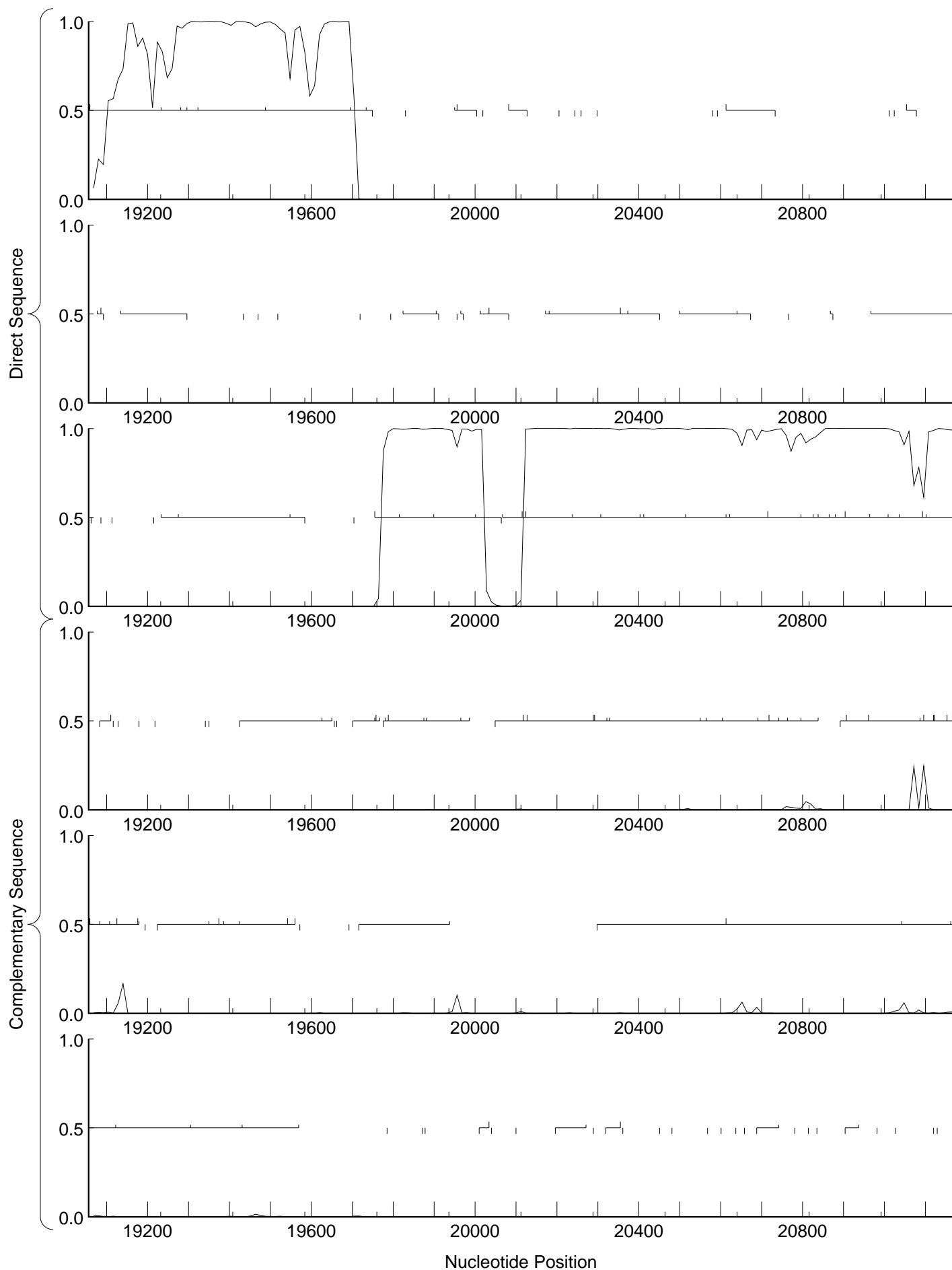


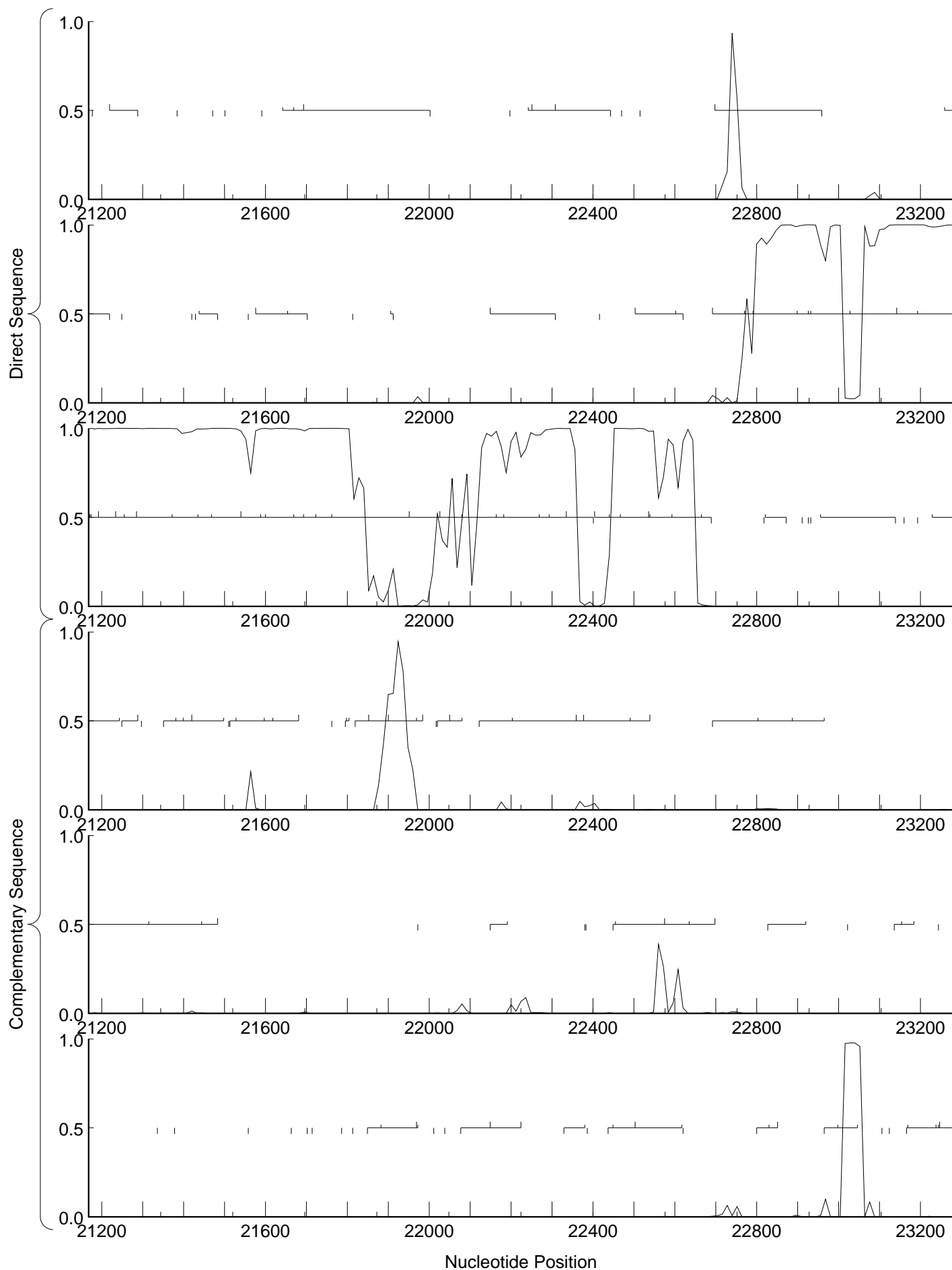


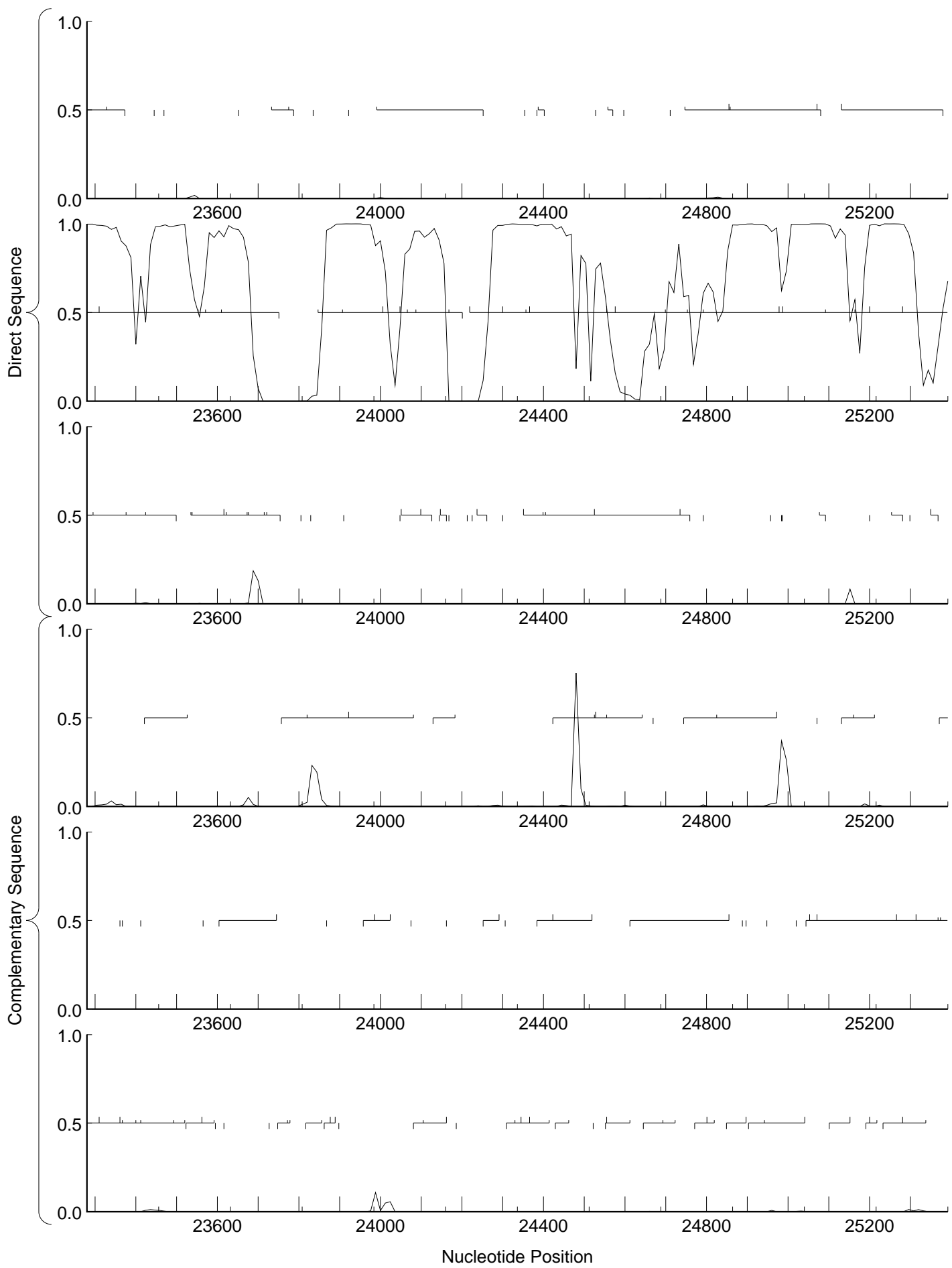


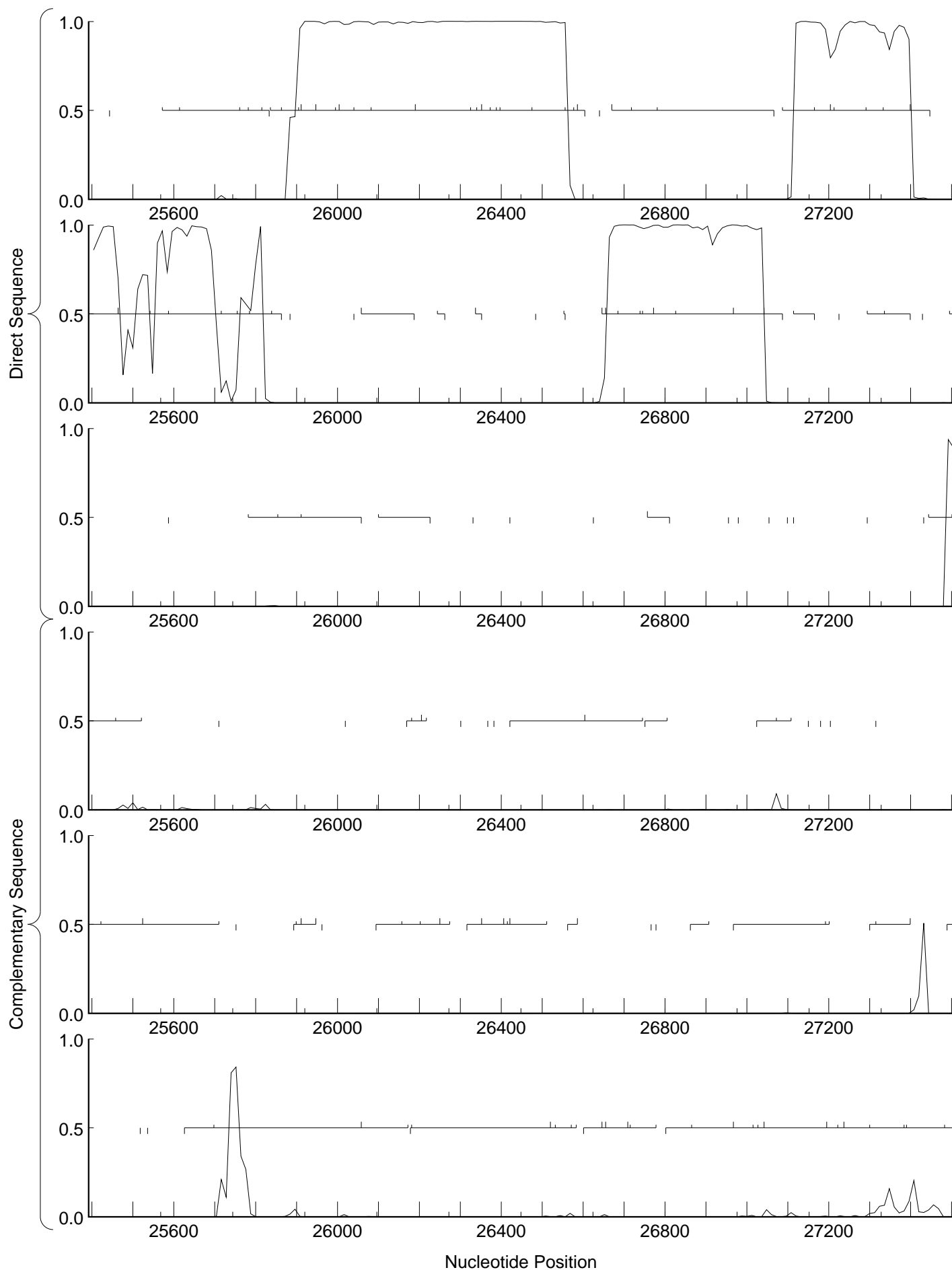


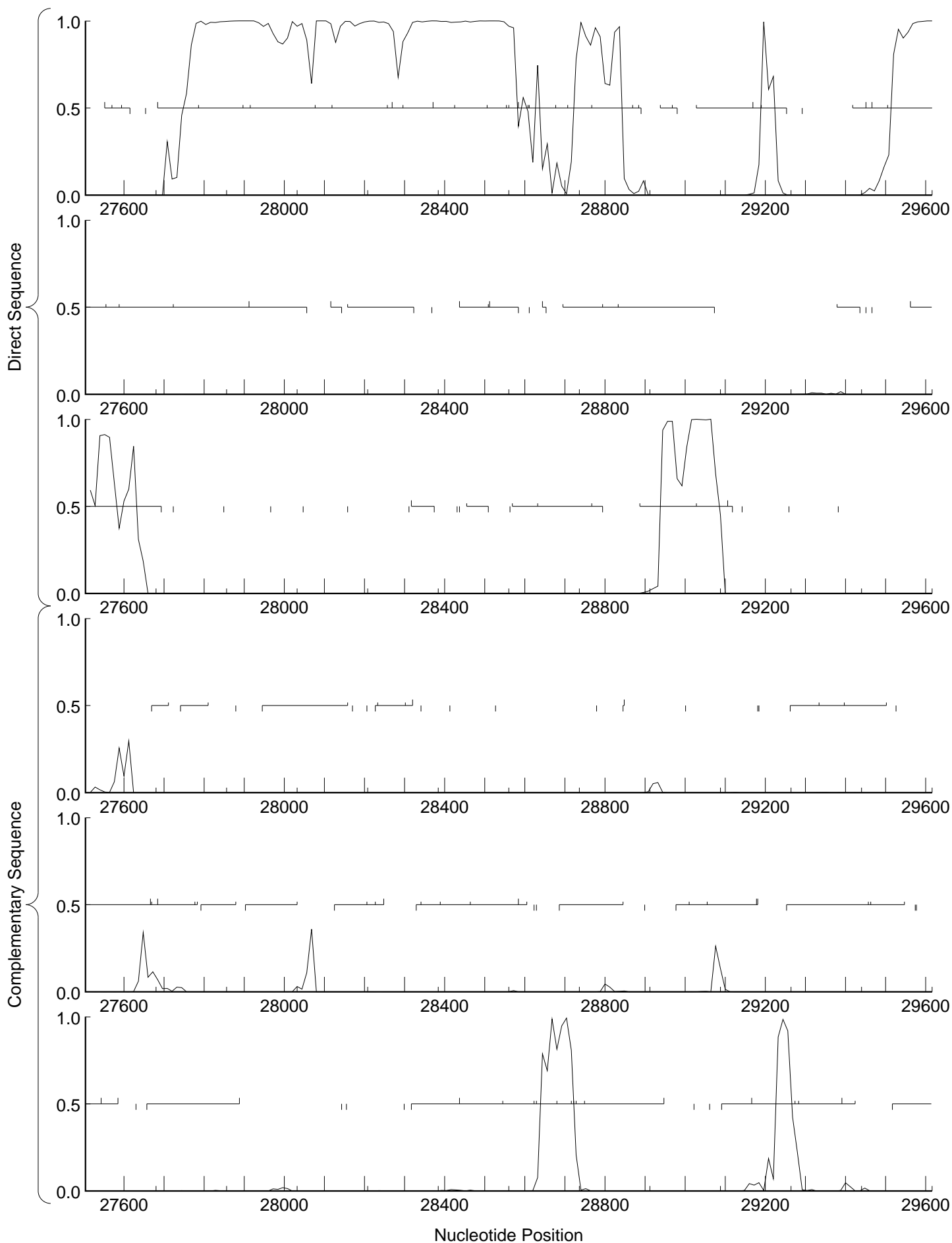


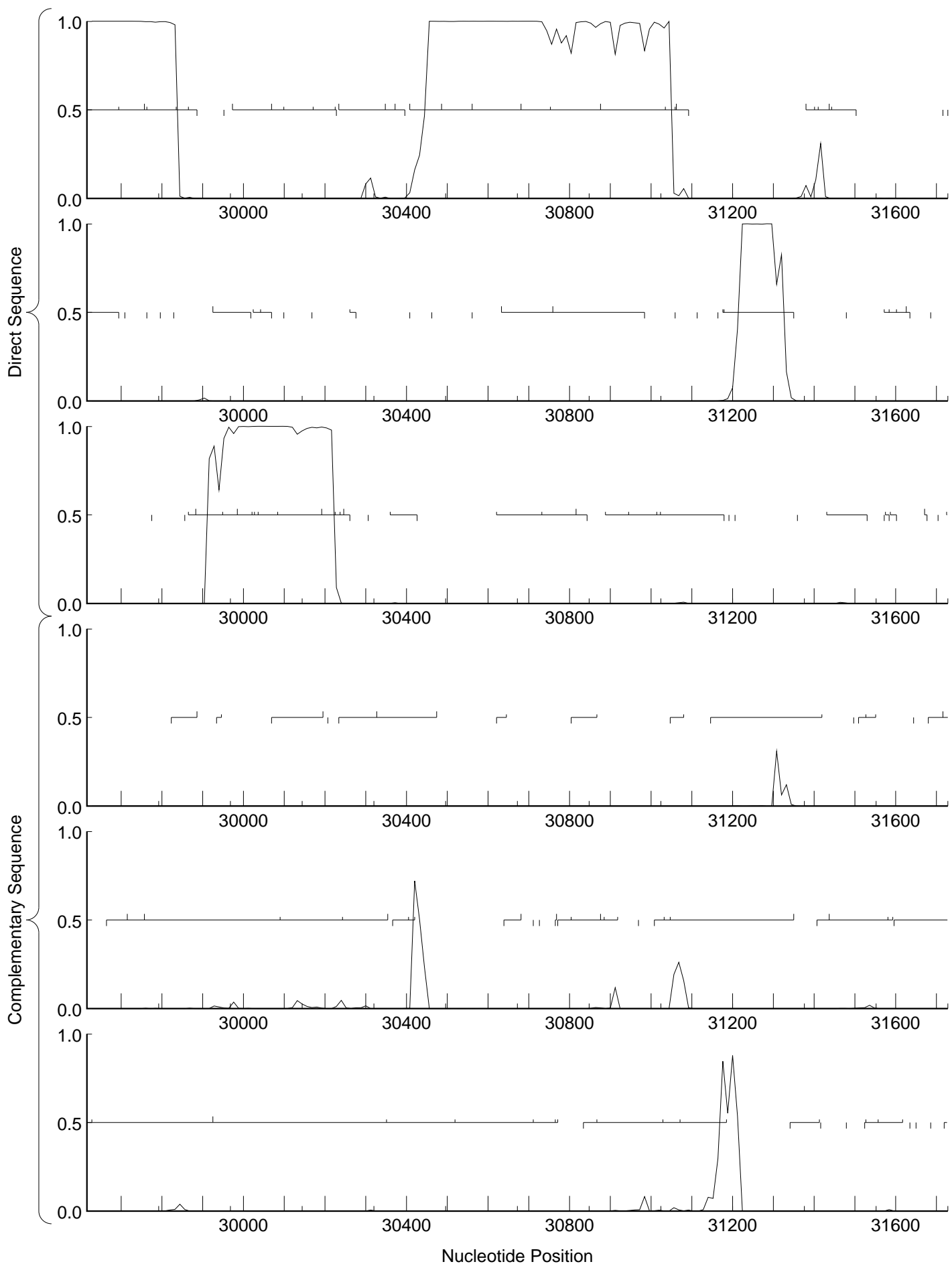


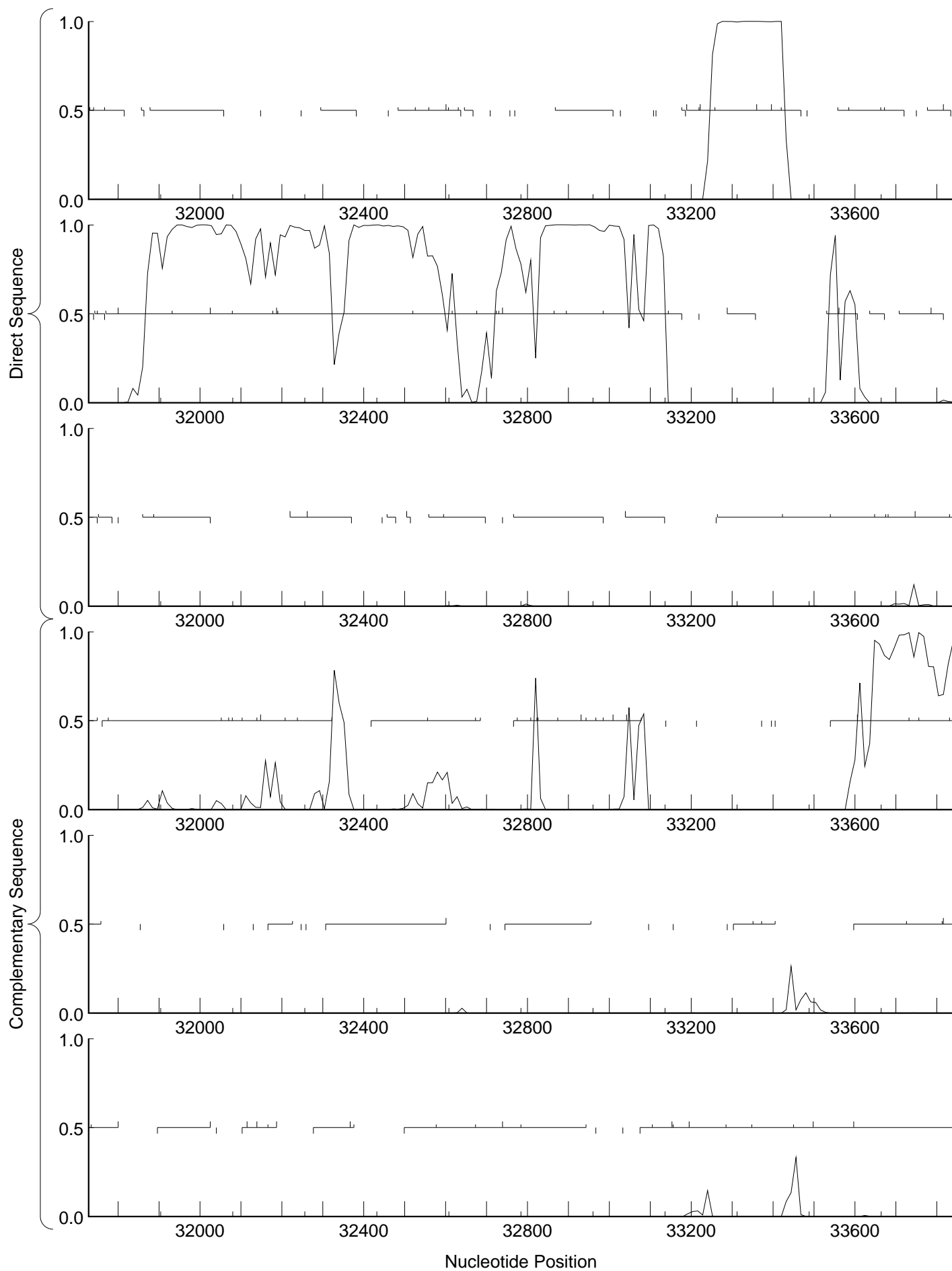


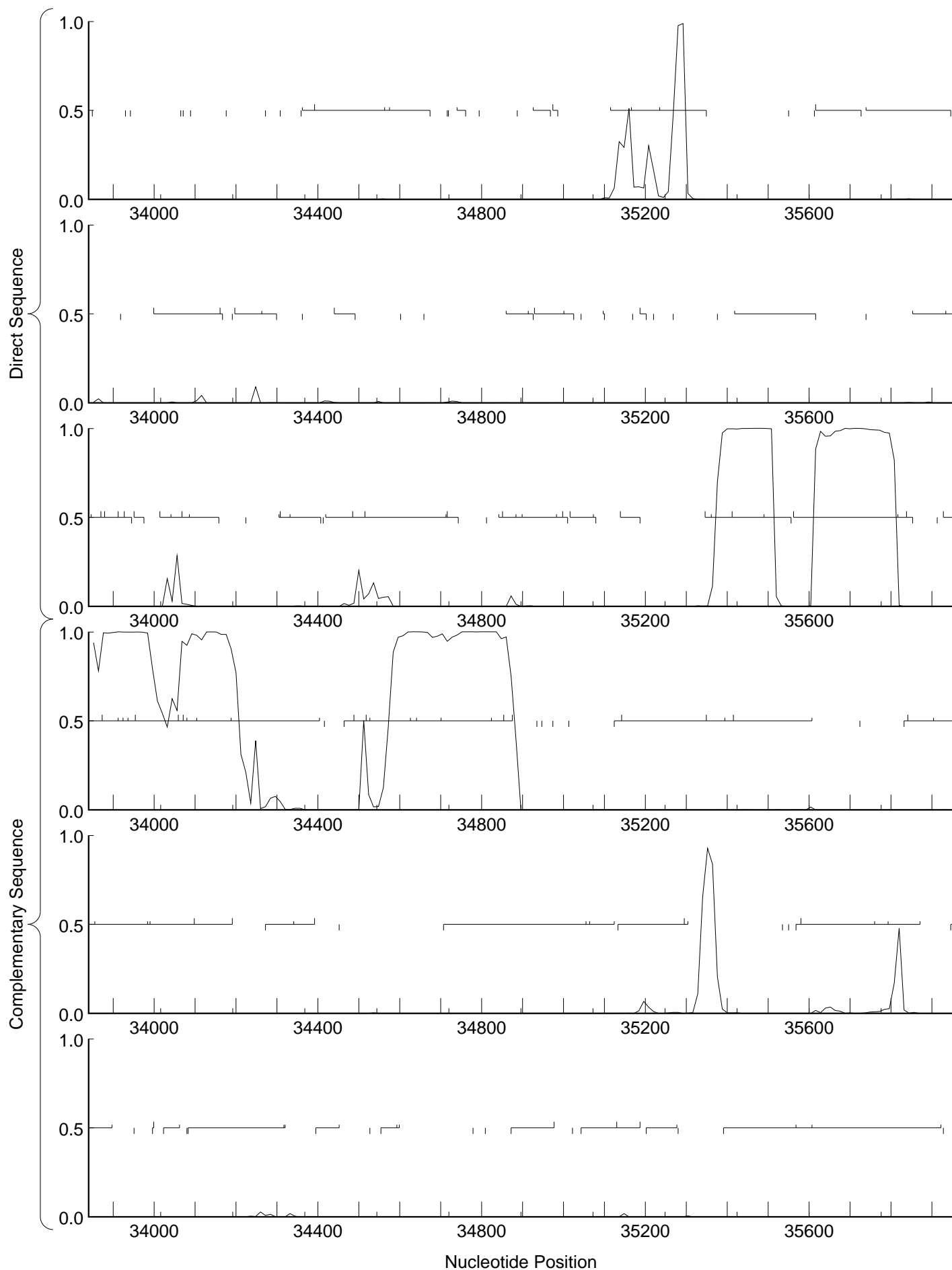












complete sequence, 57666 bp including 10-base 3' overhang (CTCGGGGCAT), Cluster K6, Order 4, Window 96, Step 12, 19/29

