

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage DyoEdafos complete sequence, 77566 bp including 10-base 3' overhang (TC
Analysis Date : 10/23/19 at 8:23:40
Pages : 38
Sequence Length : 77566 bp
GC Content : 58.54%

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,

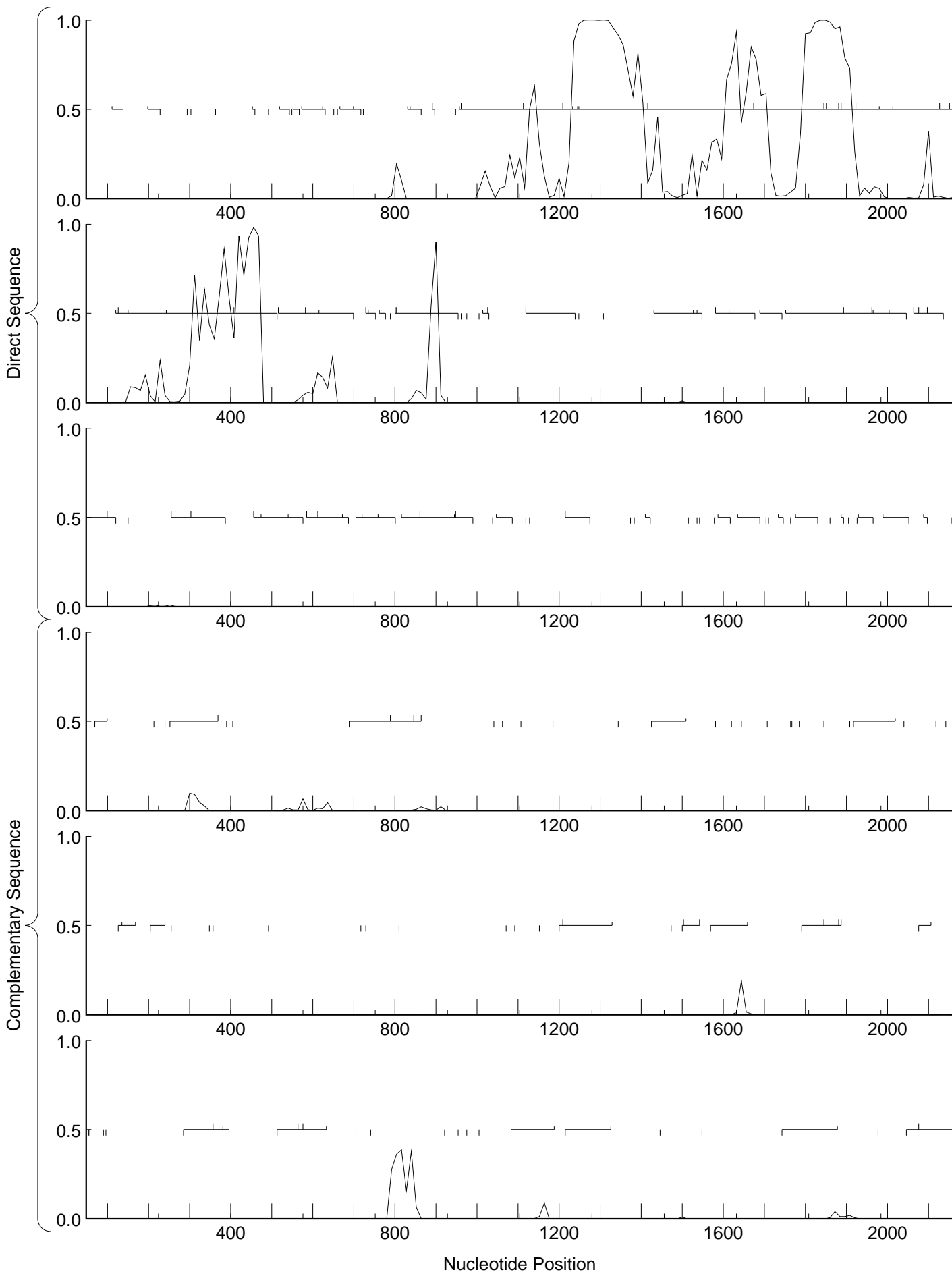
Matrix notes & comments

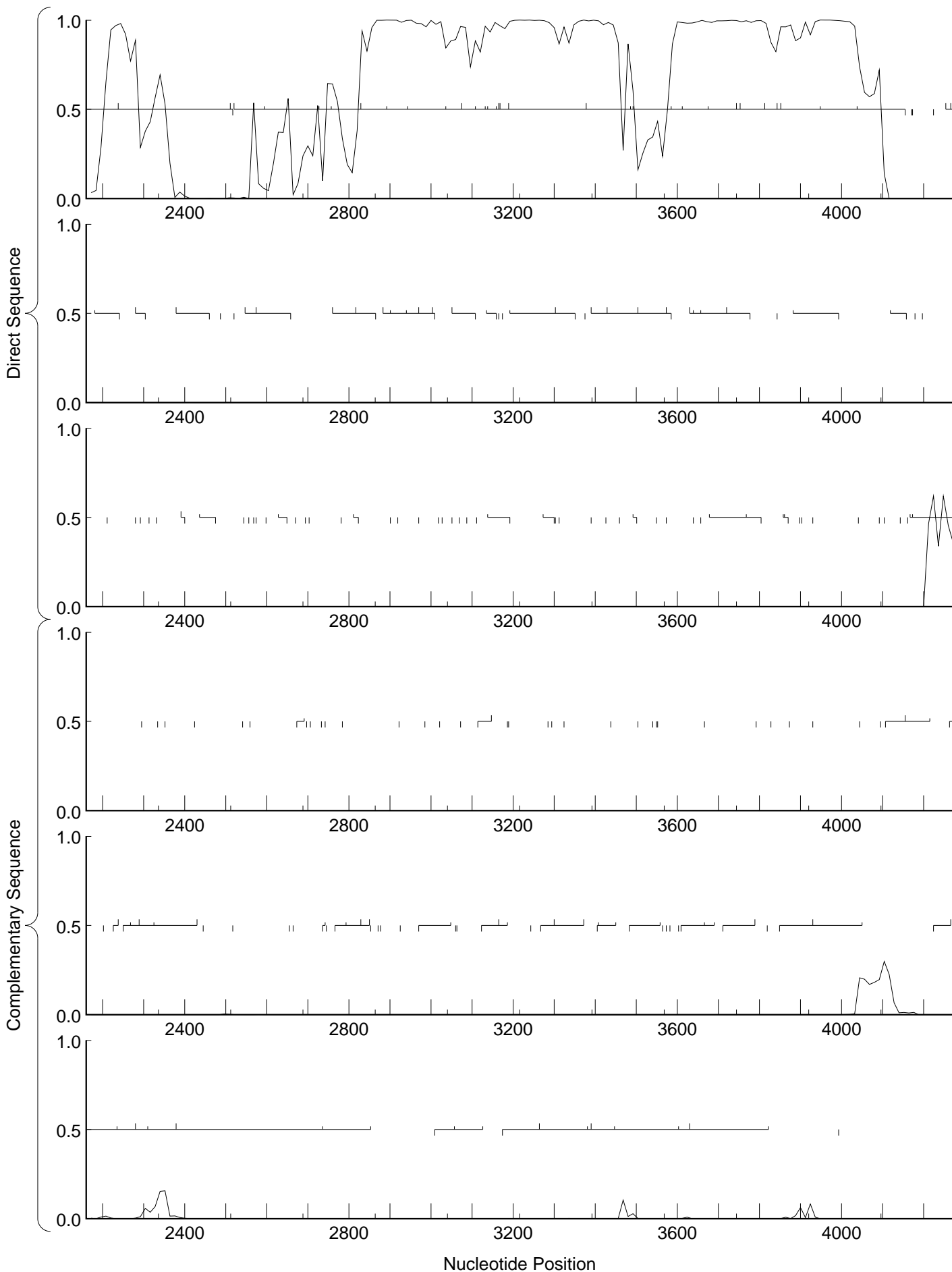
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

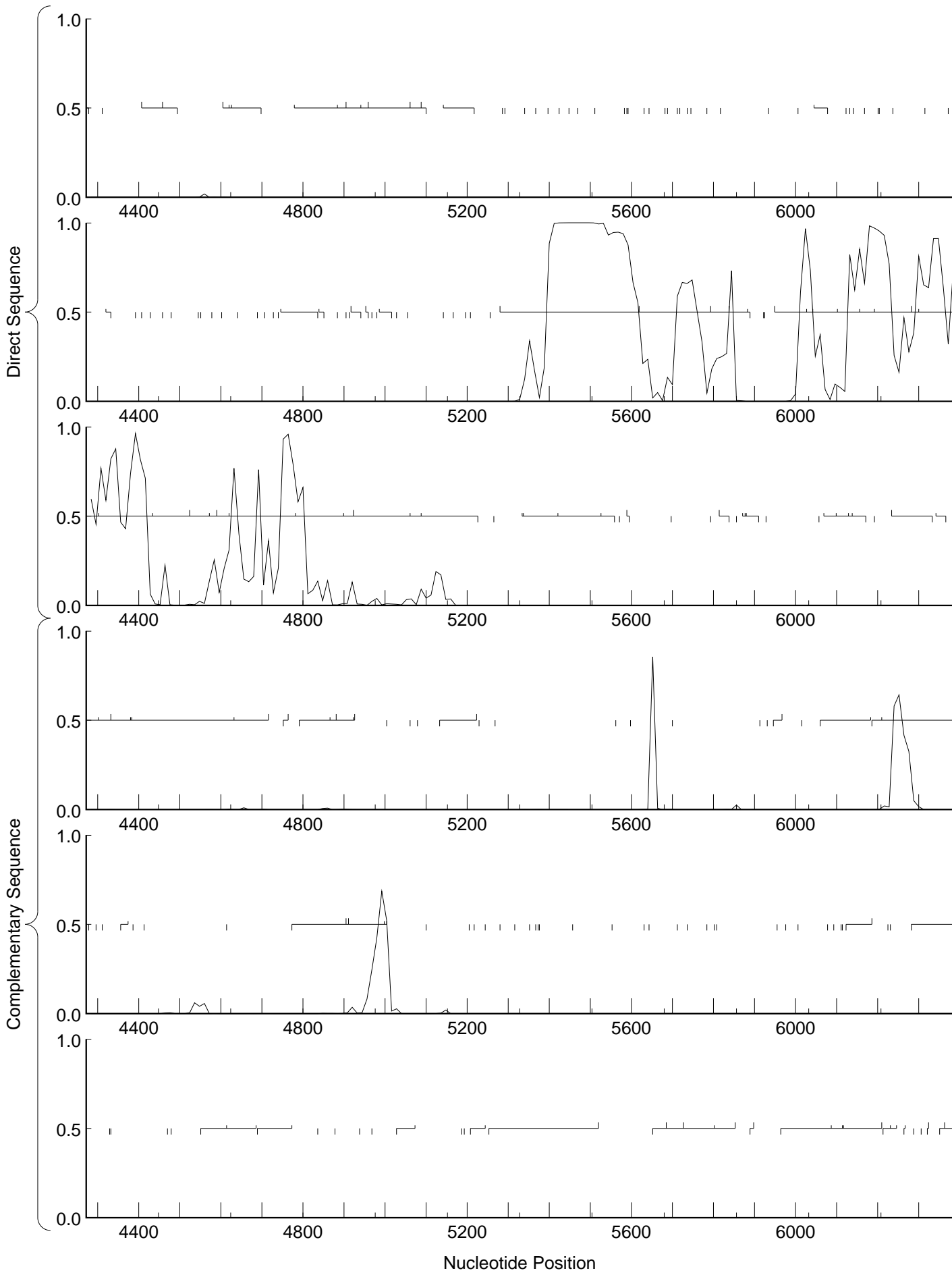
MATRIX INFORMATION

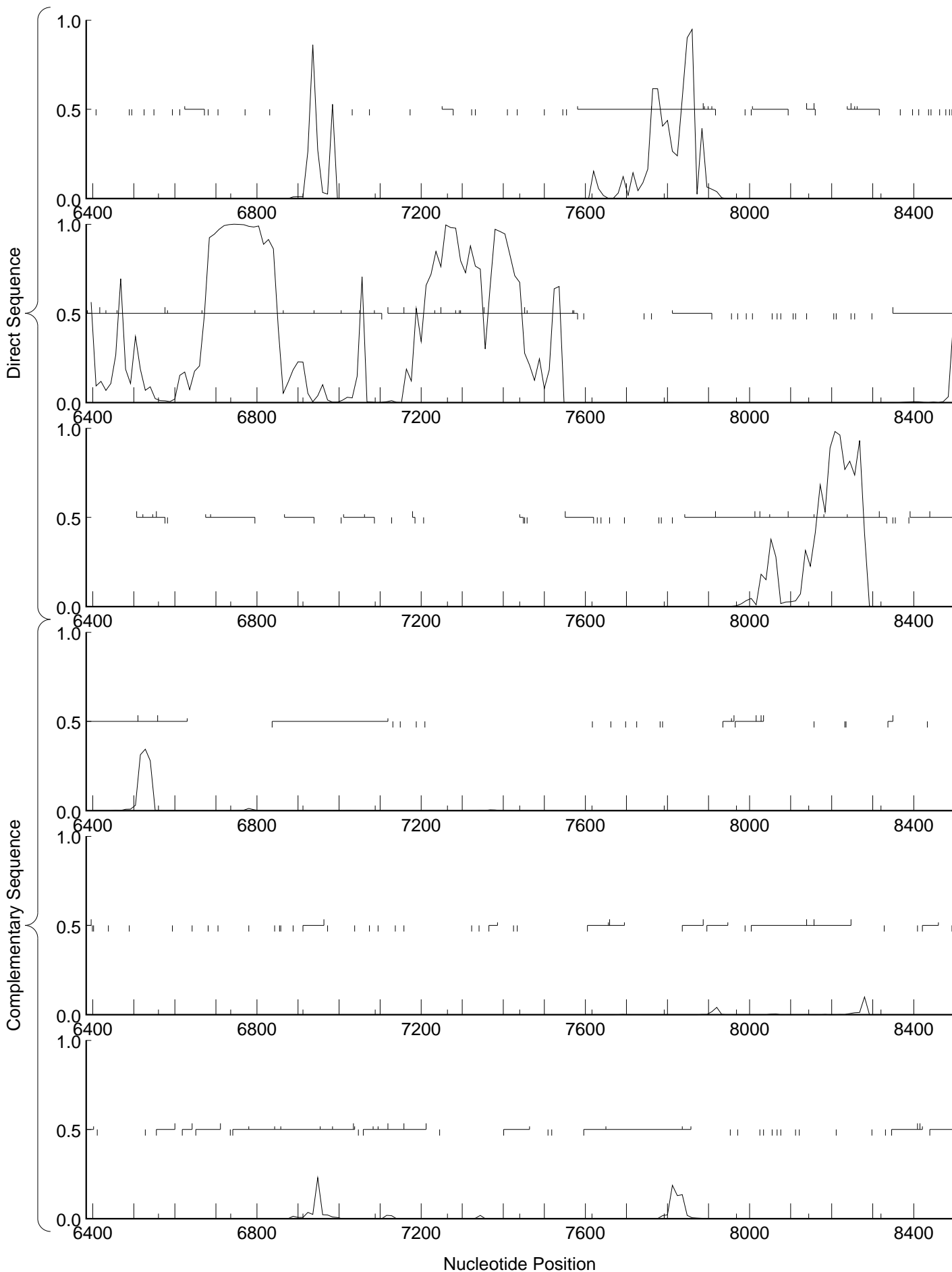
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

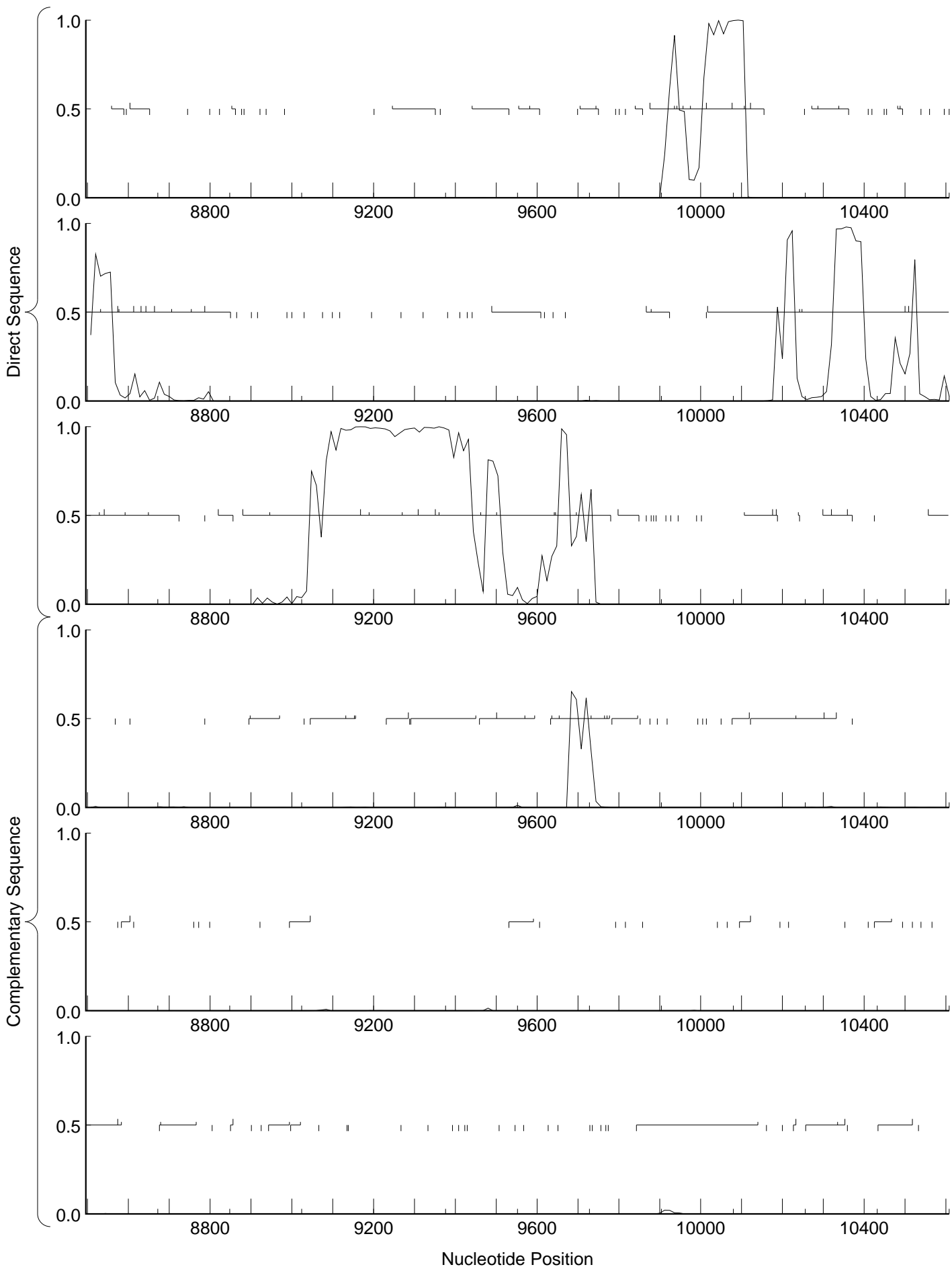
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

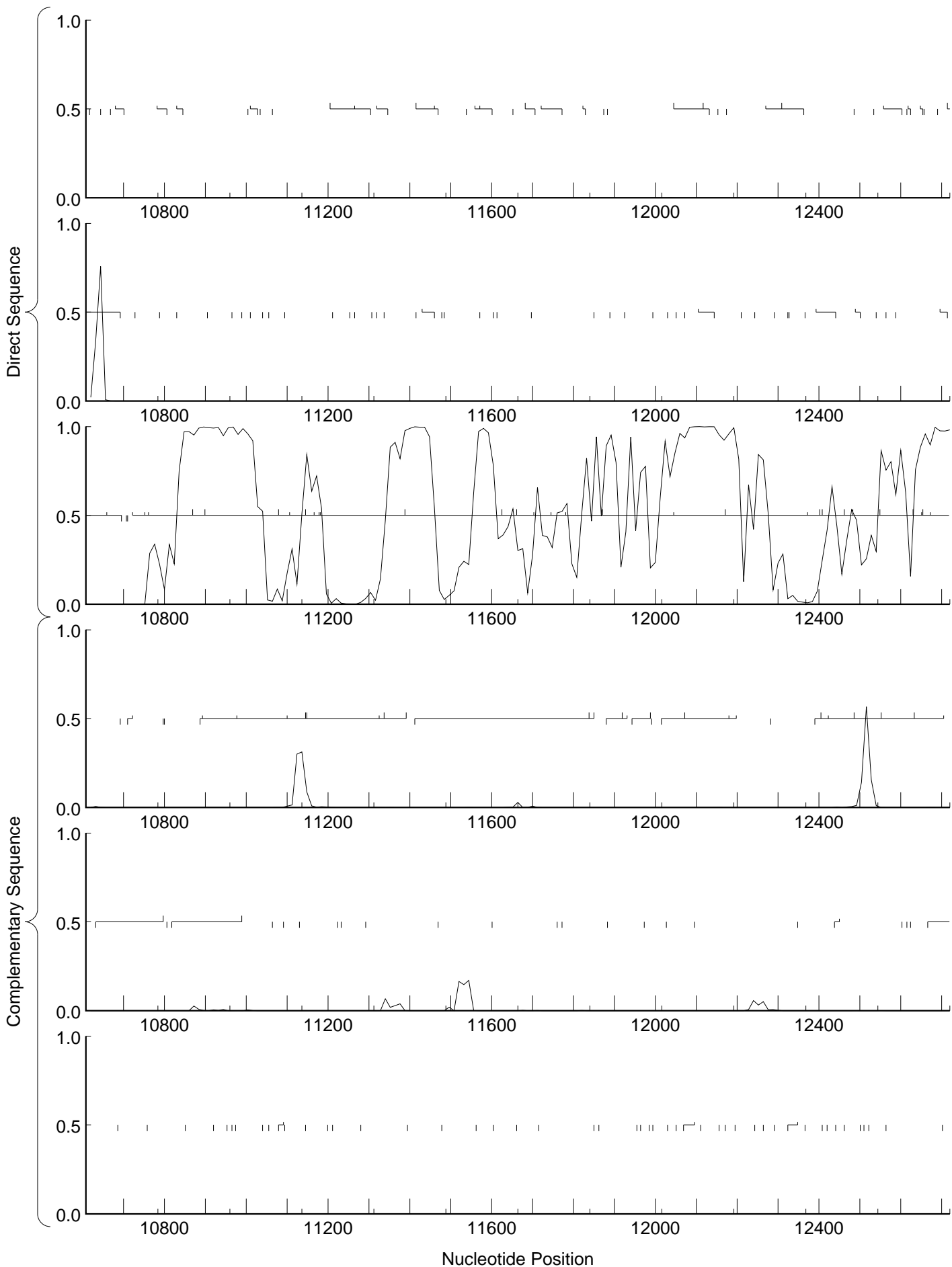


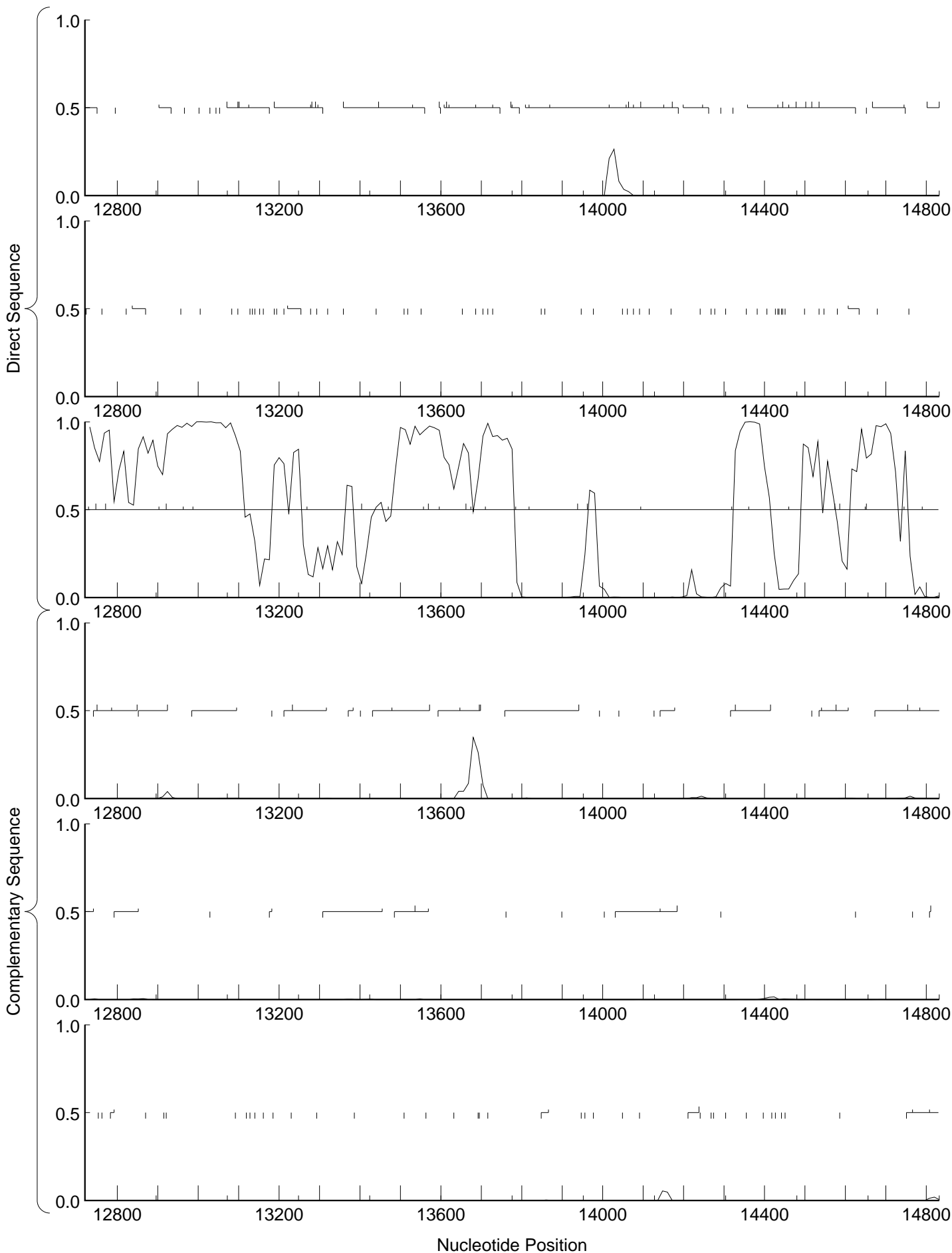


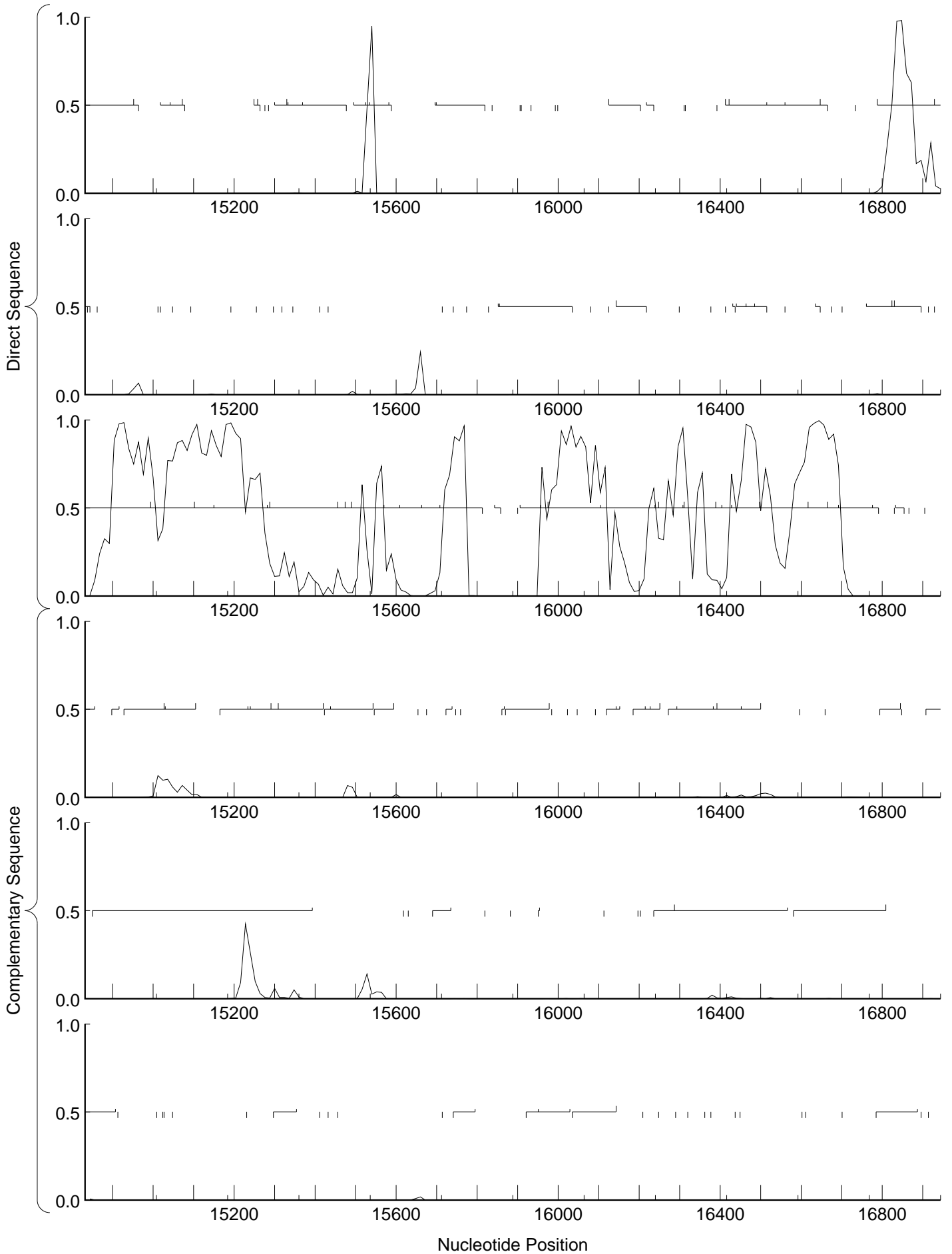


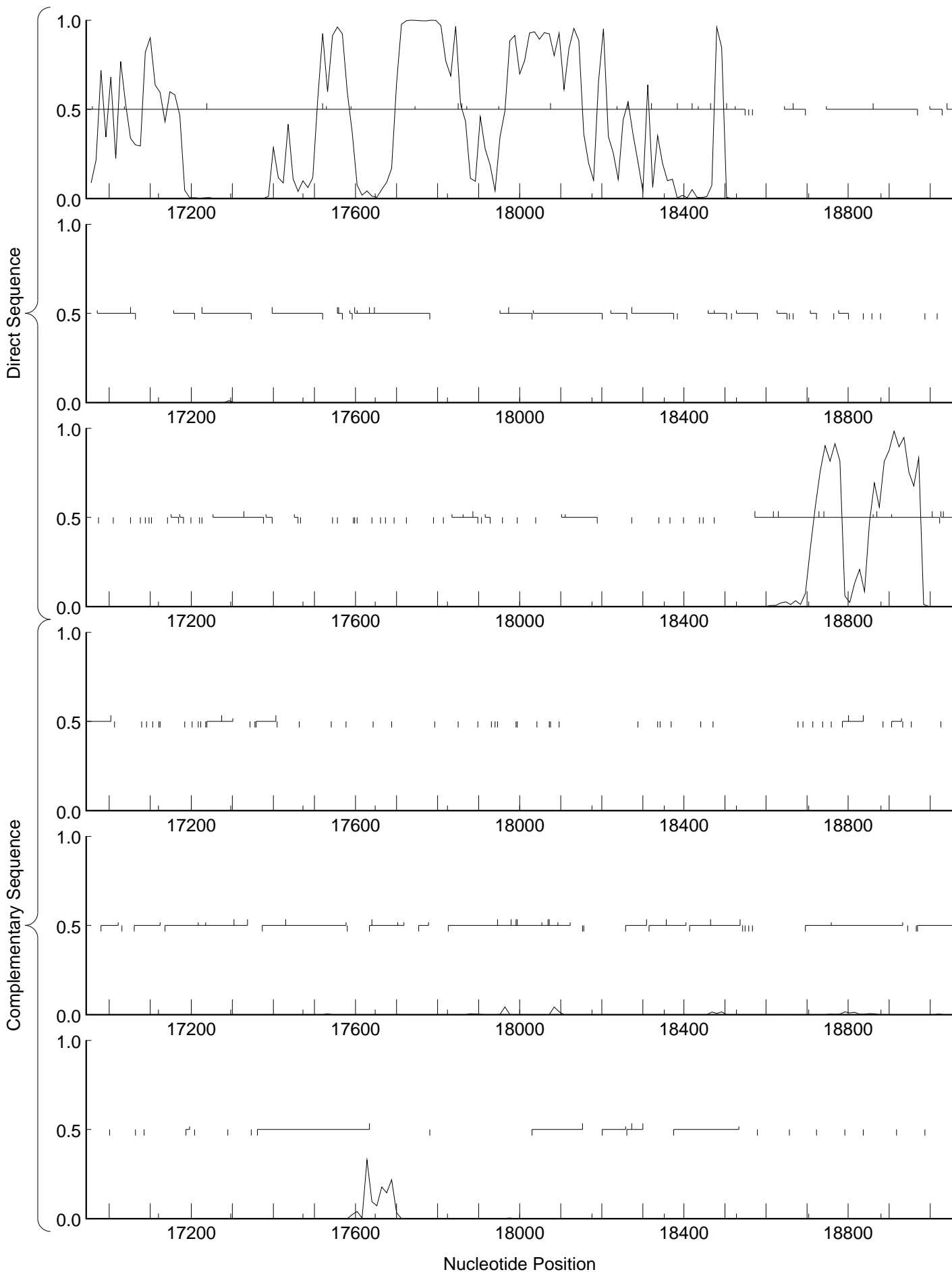




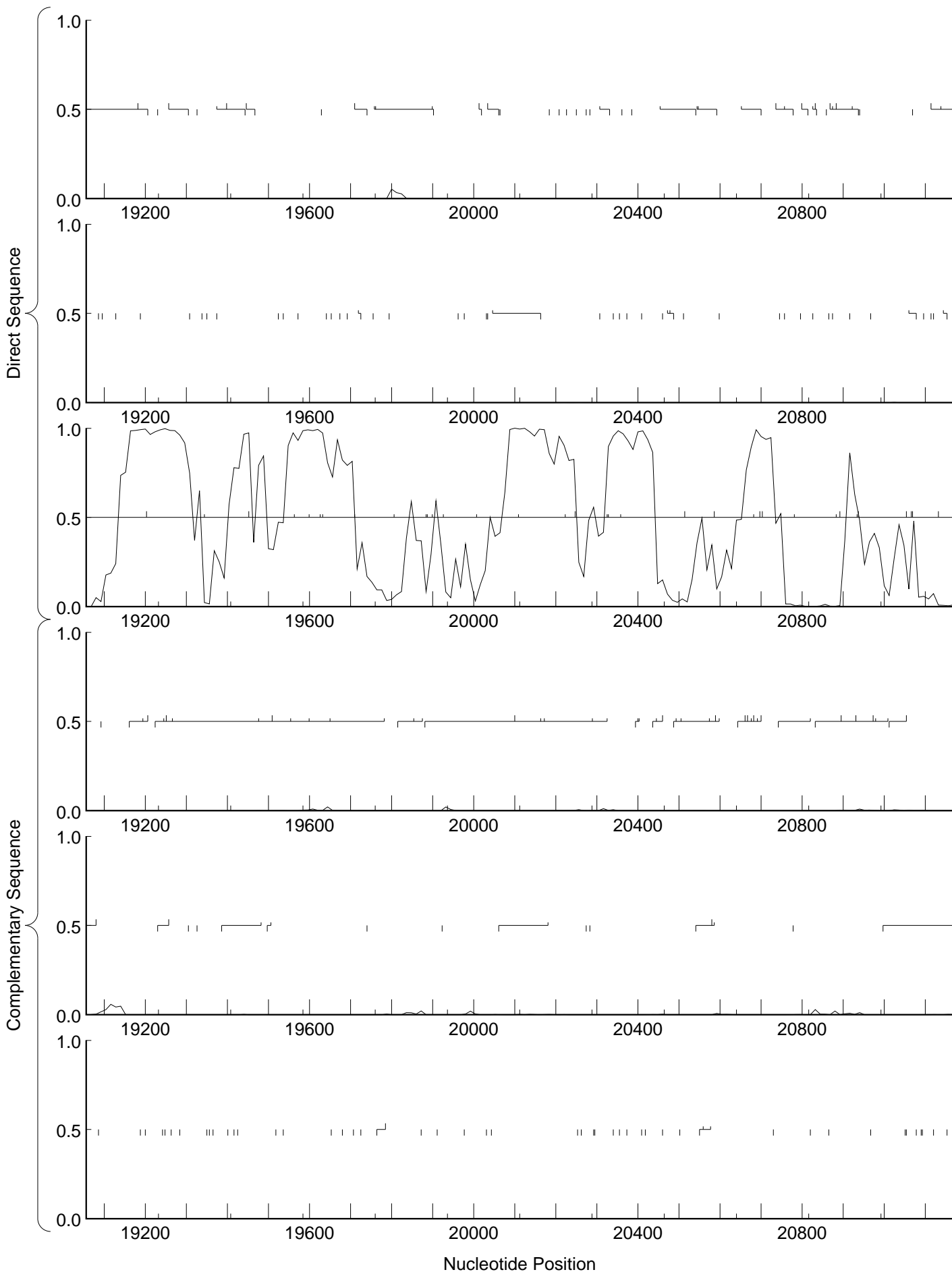


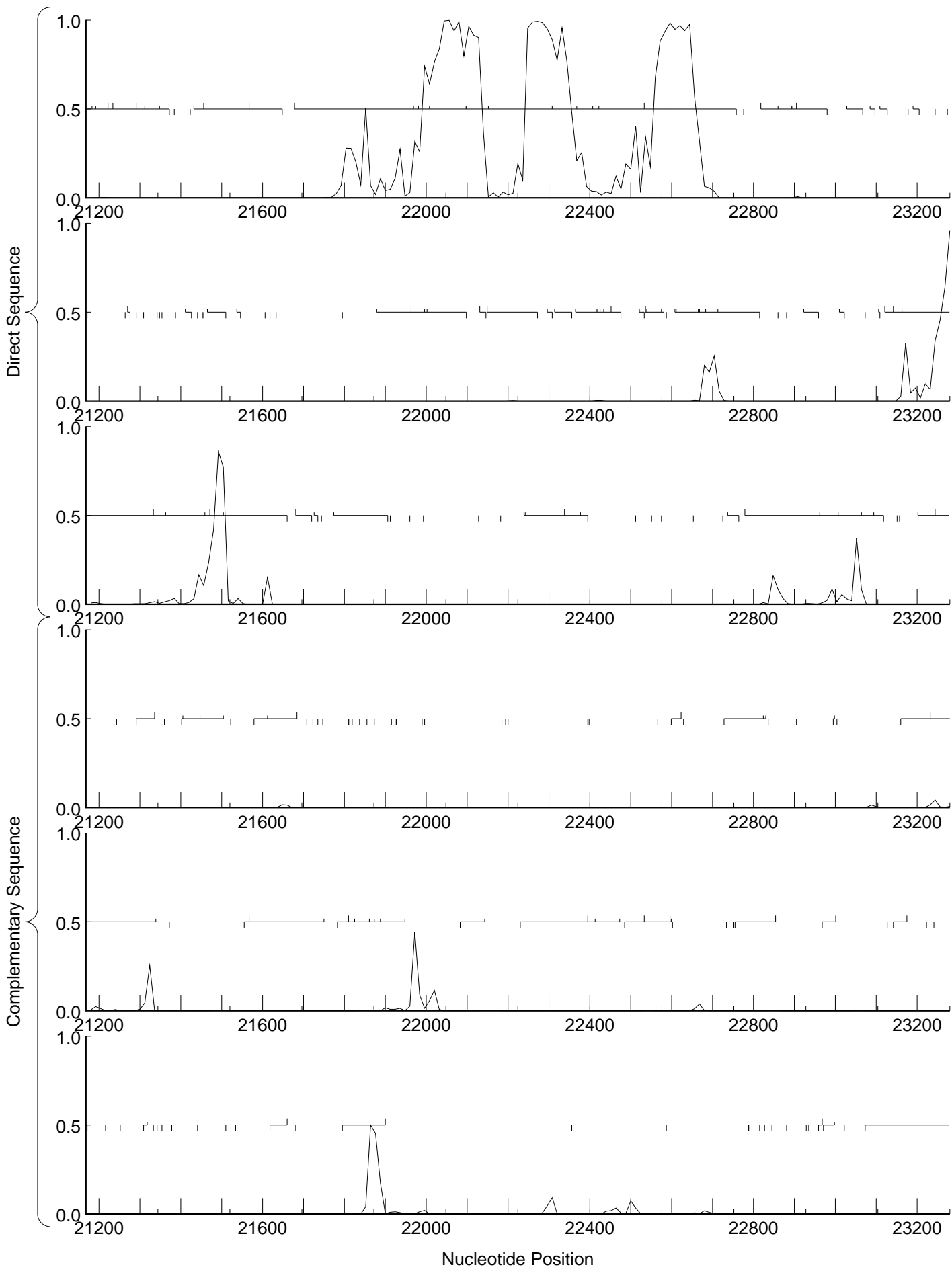


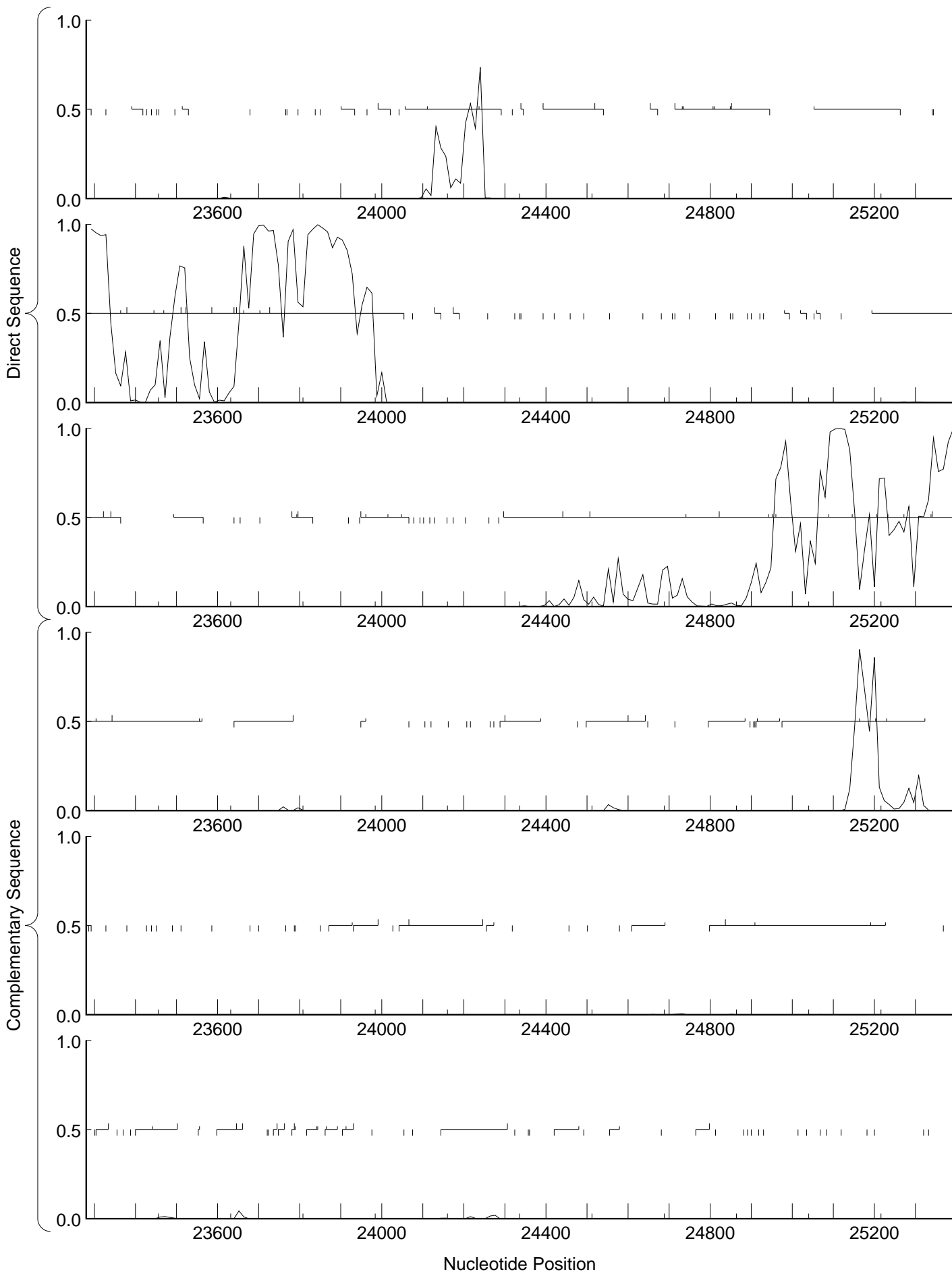


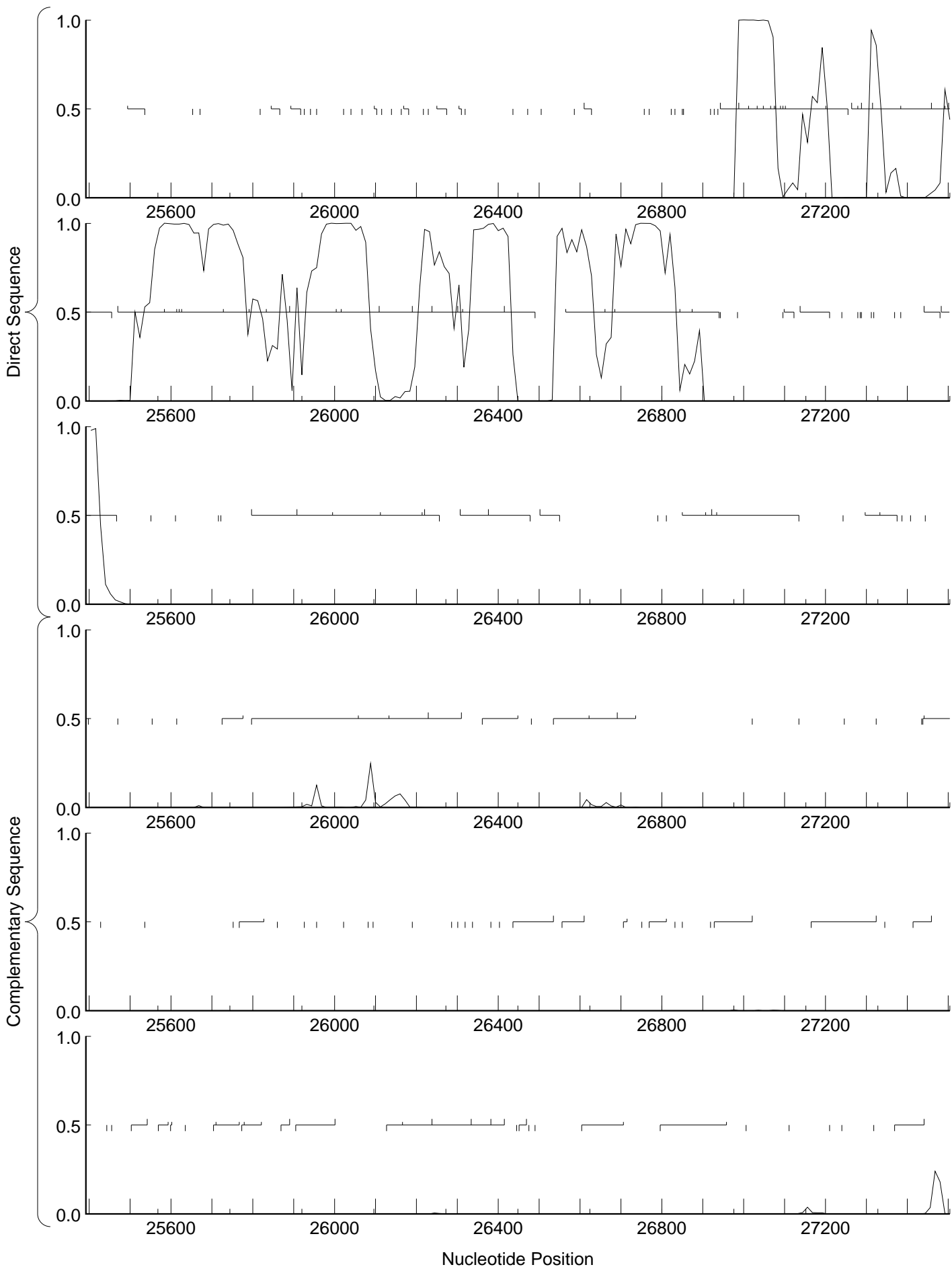


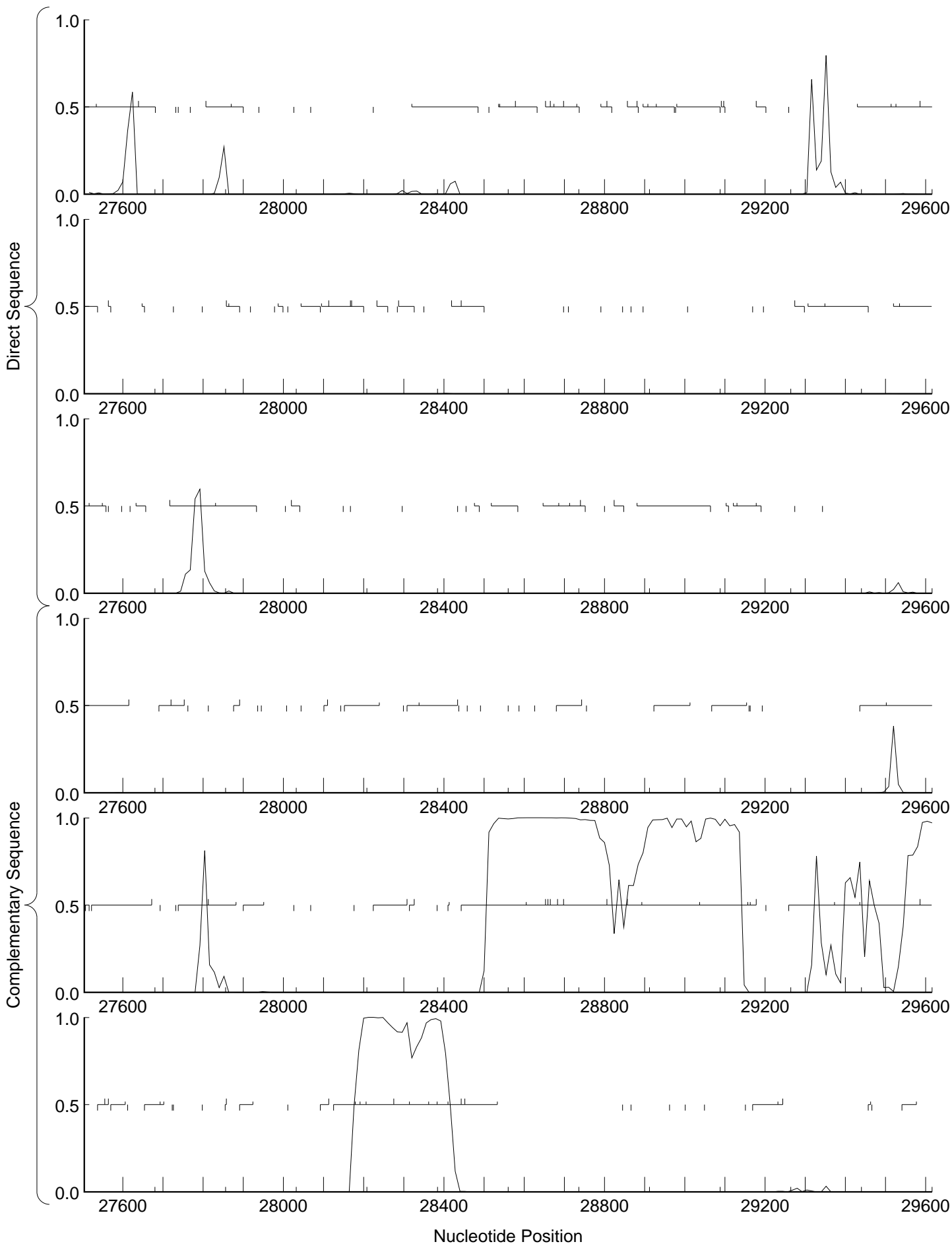
s complete sequence, 77566 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L4, Order 4, Window 96, Step 12, 11/38

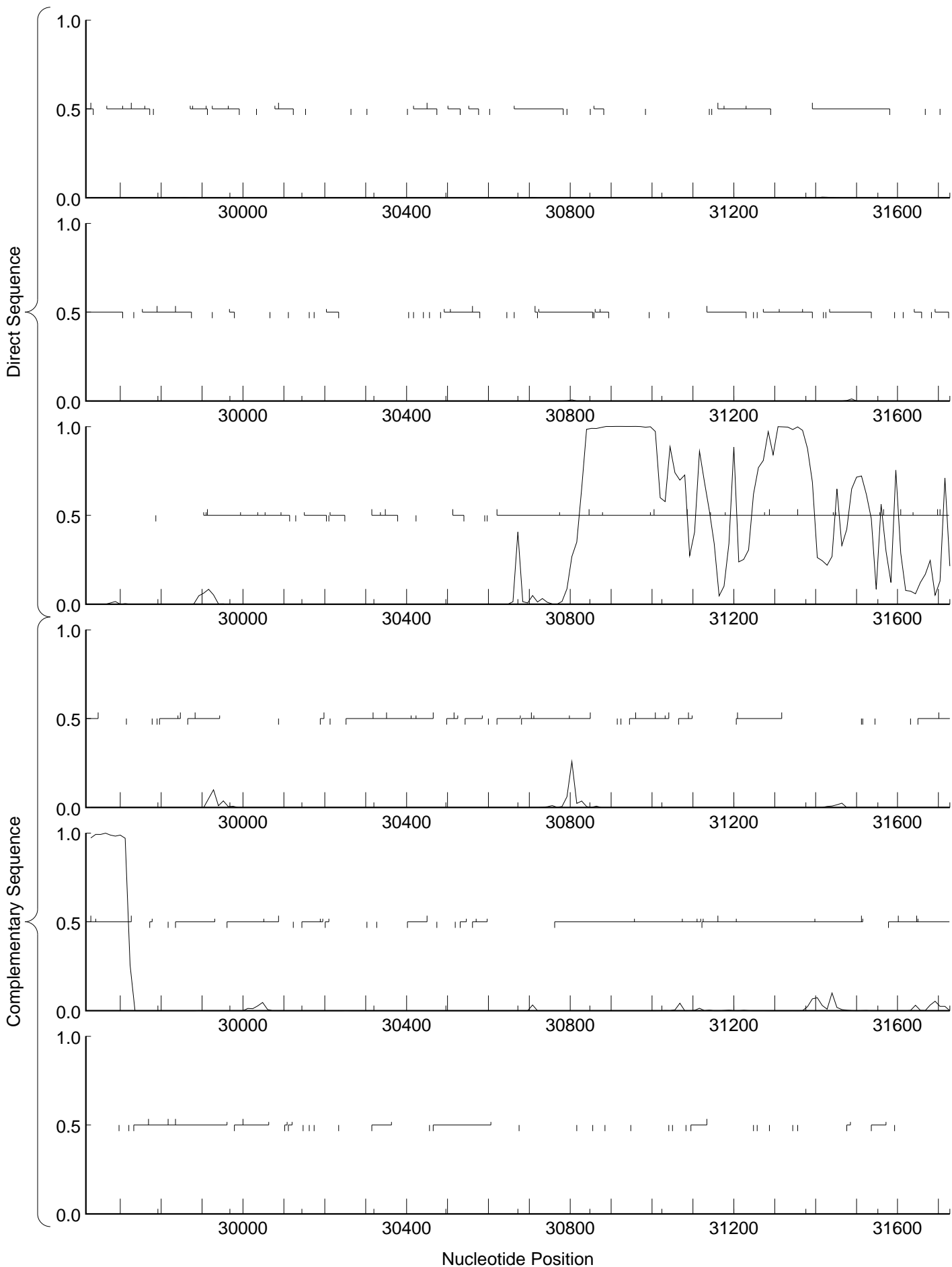


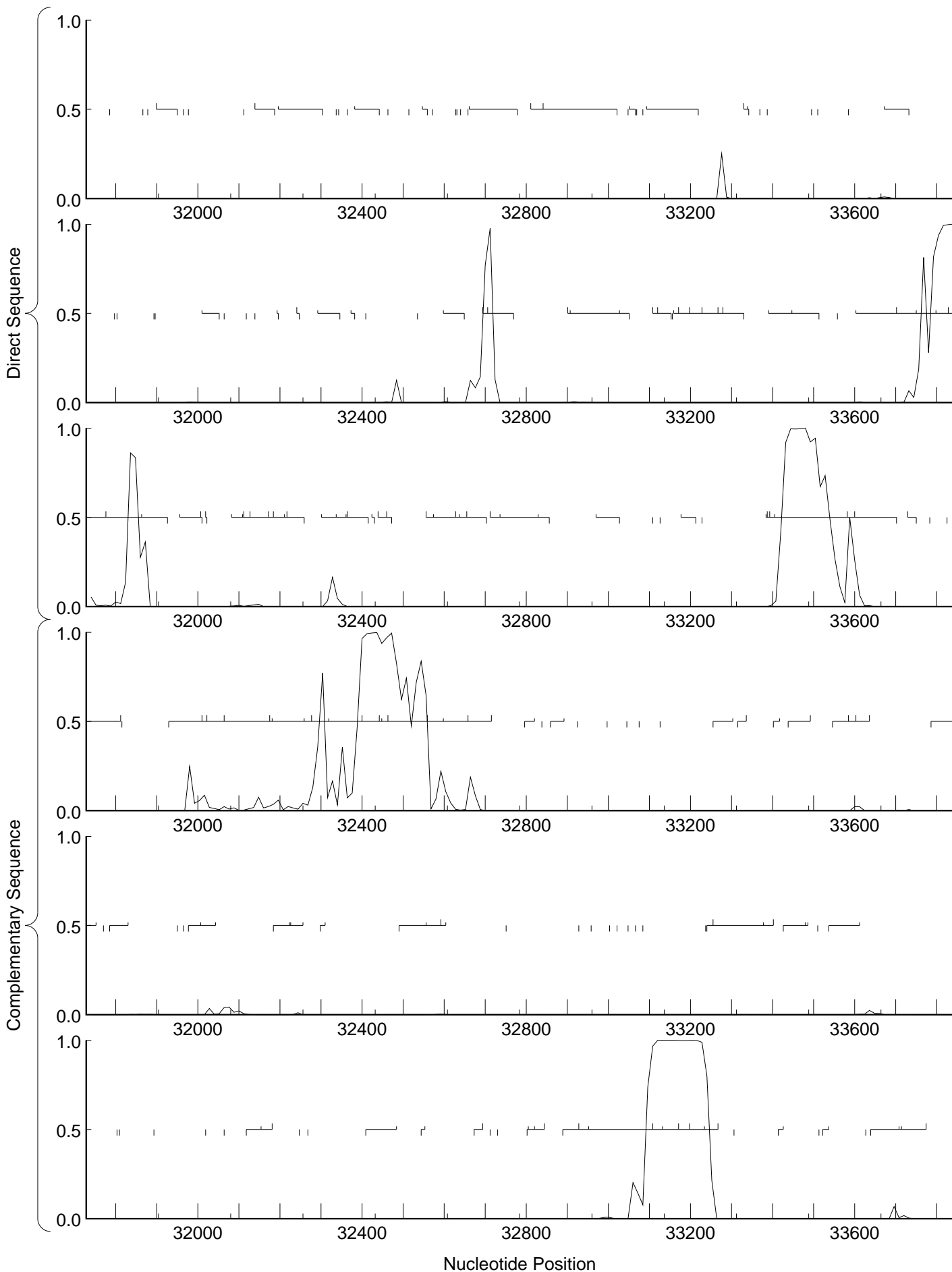


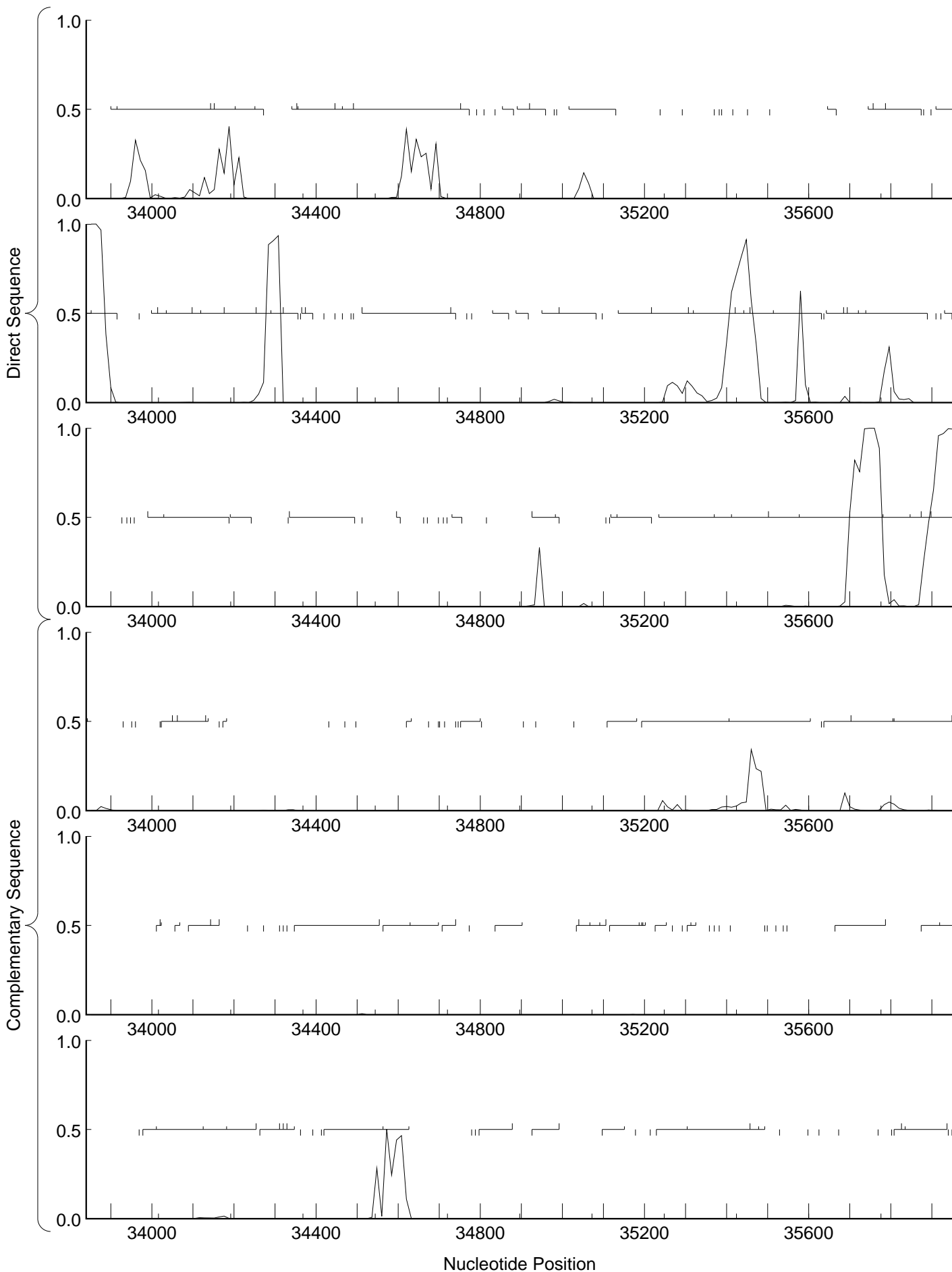


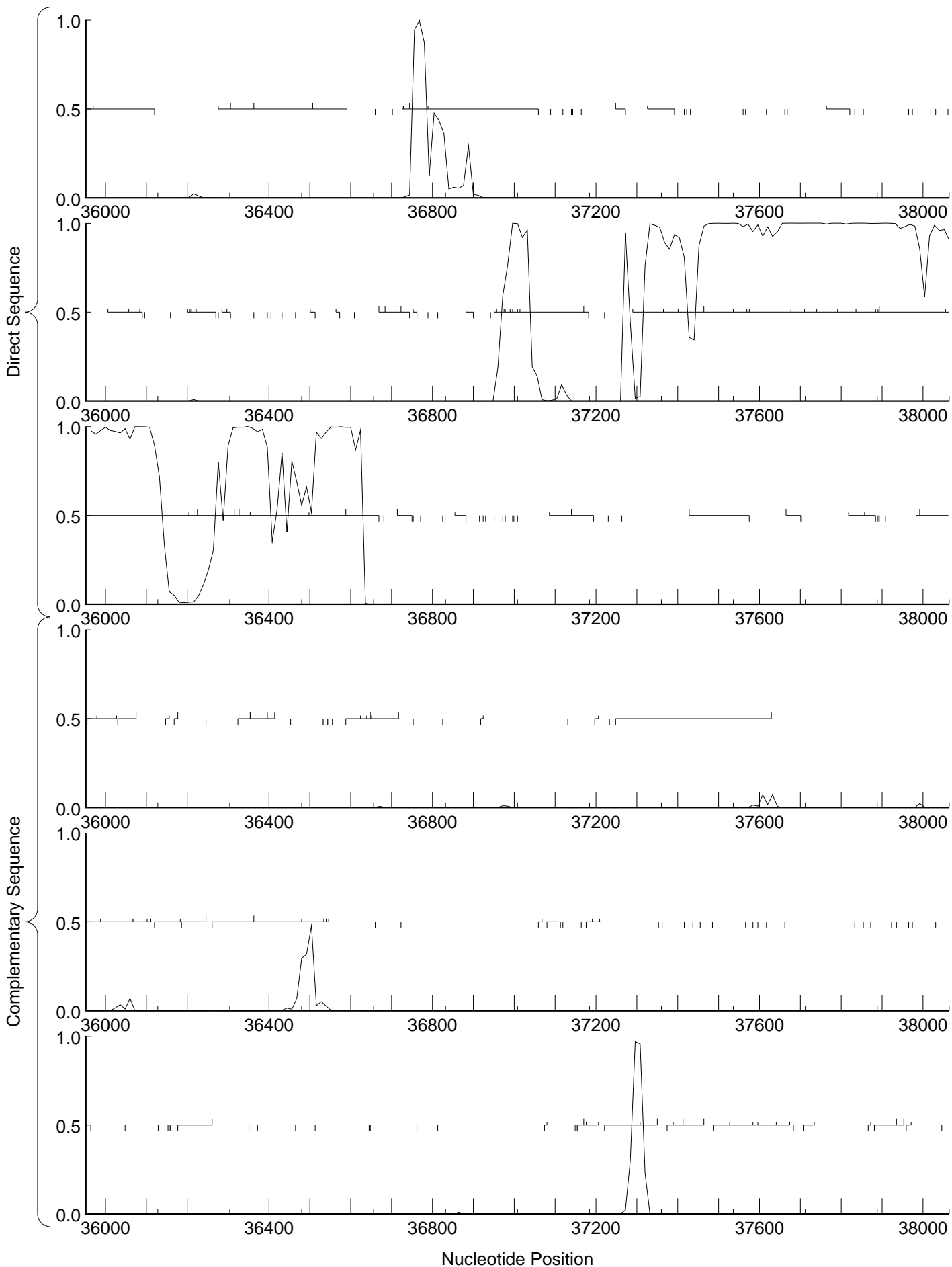




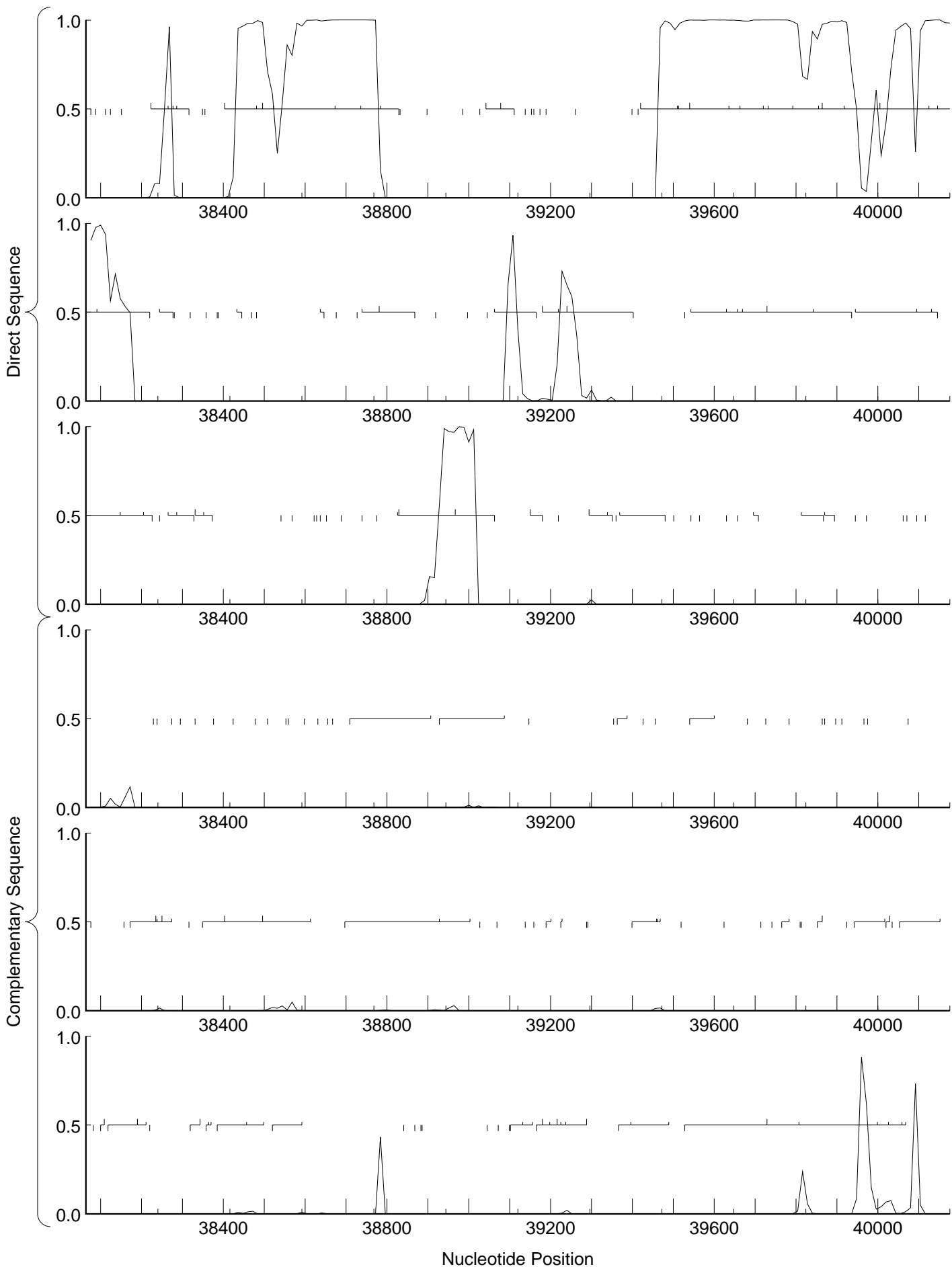


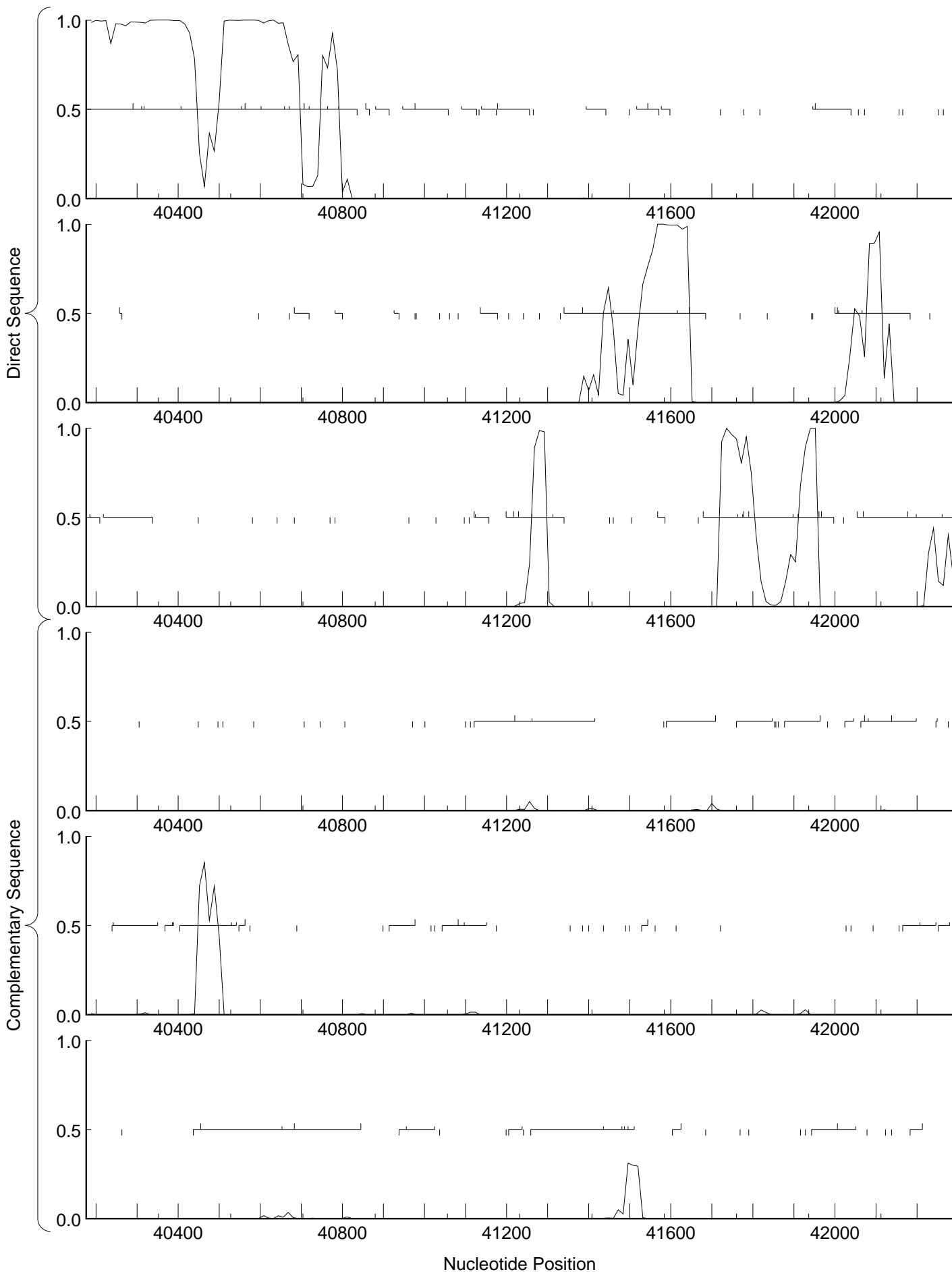


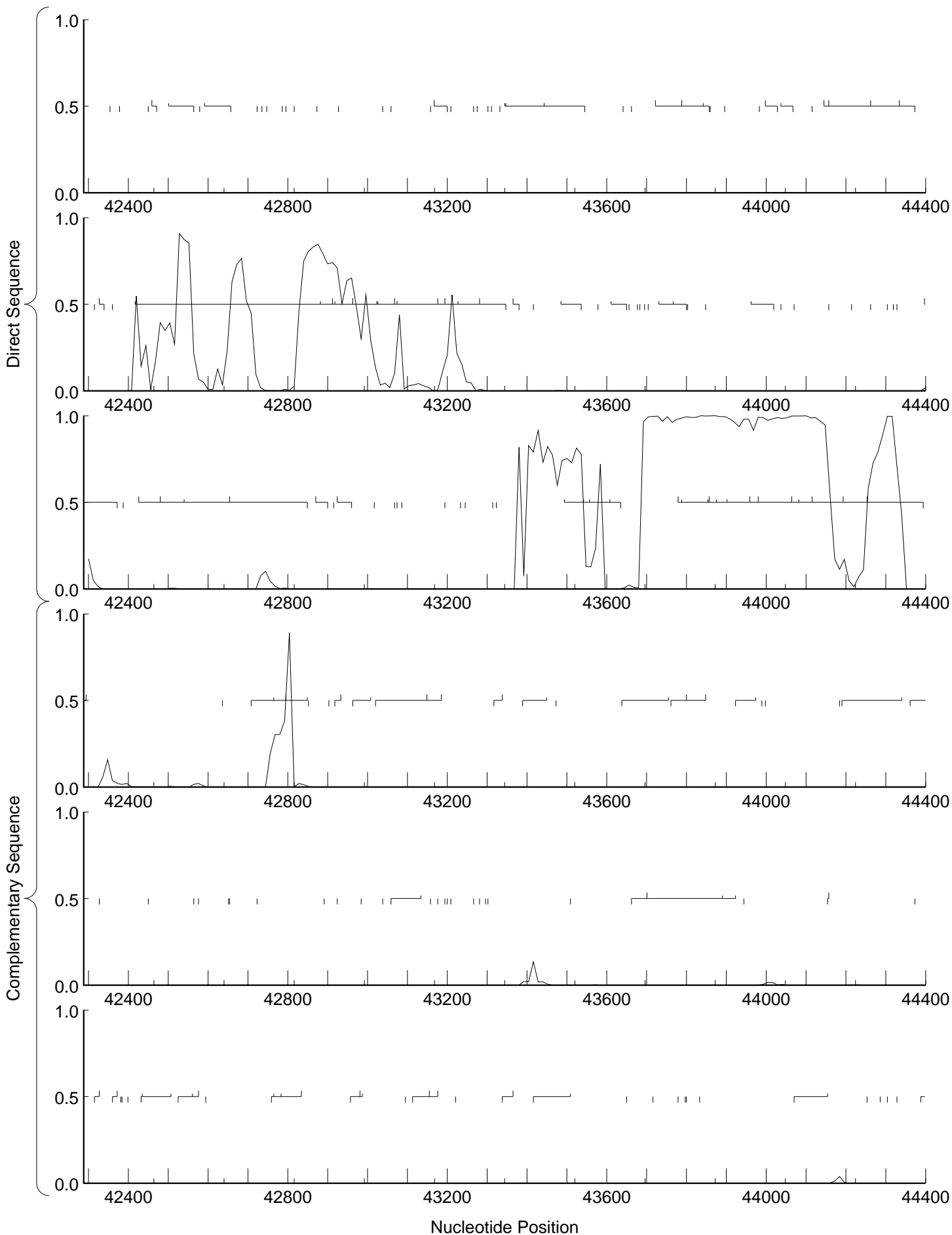


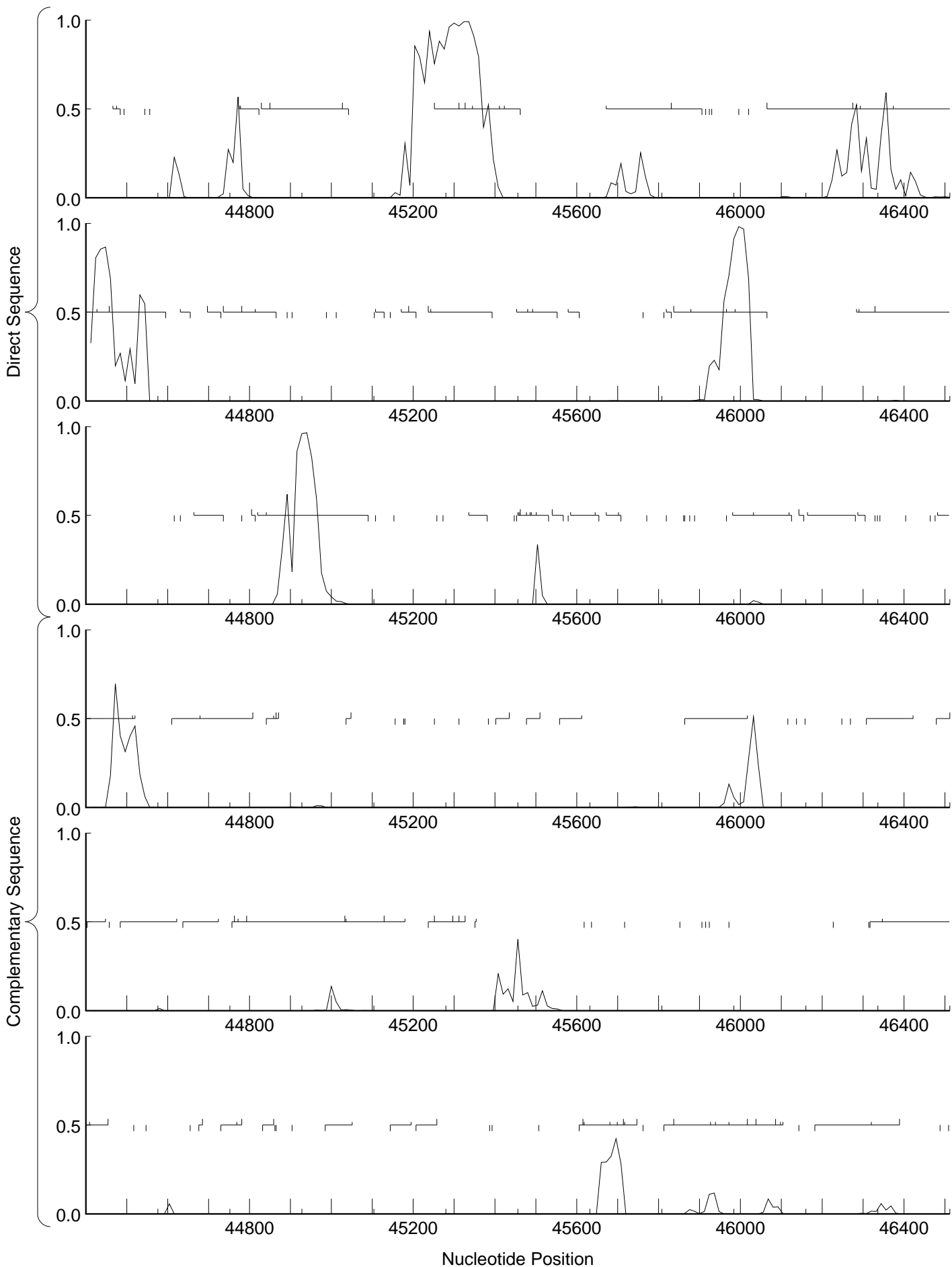


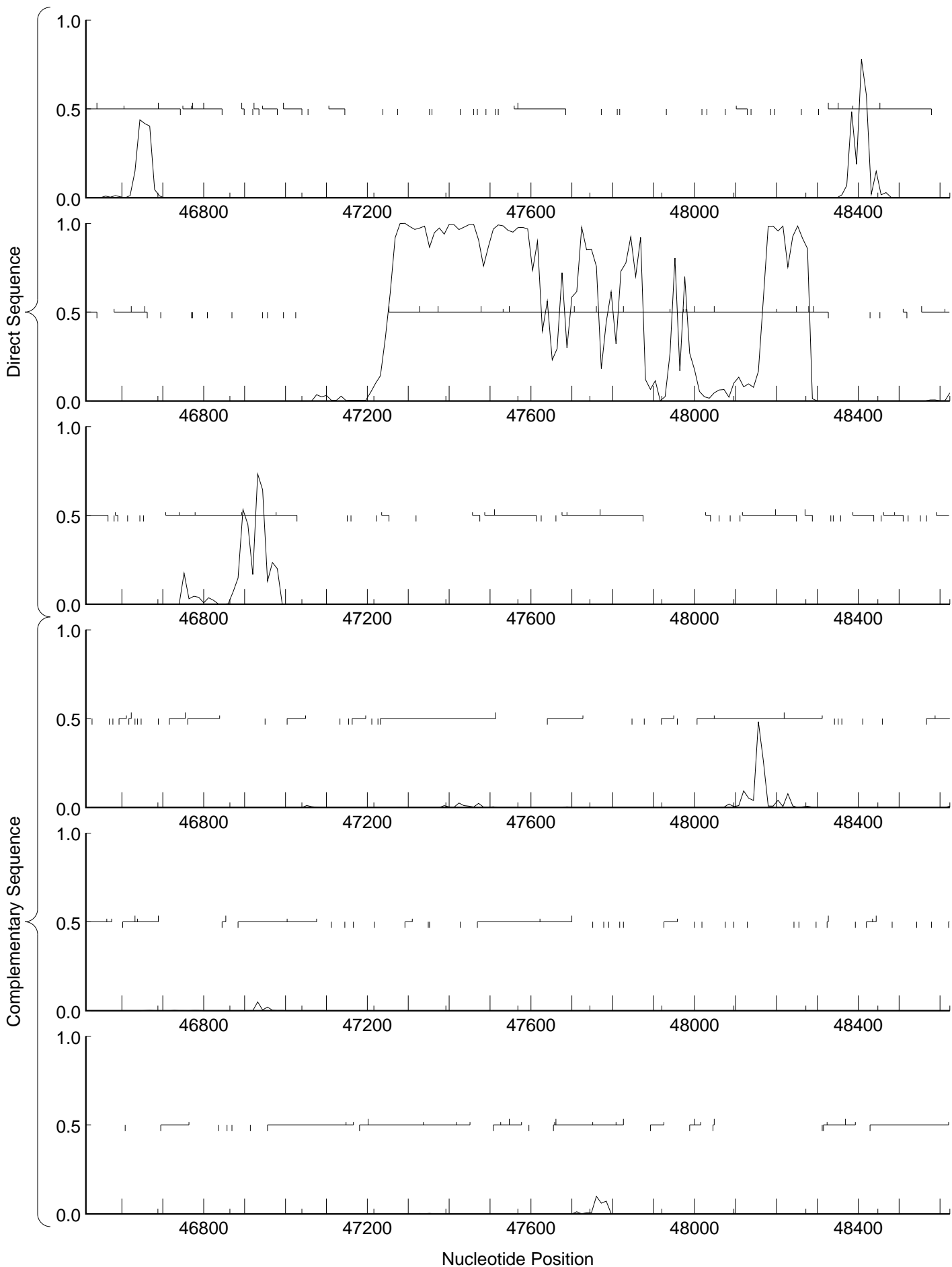
s complete sequence, 77566 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L4, Order 4, Window 96, Step 12, 20/38

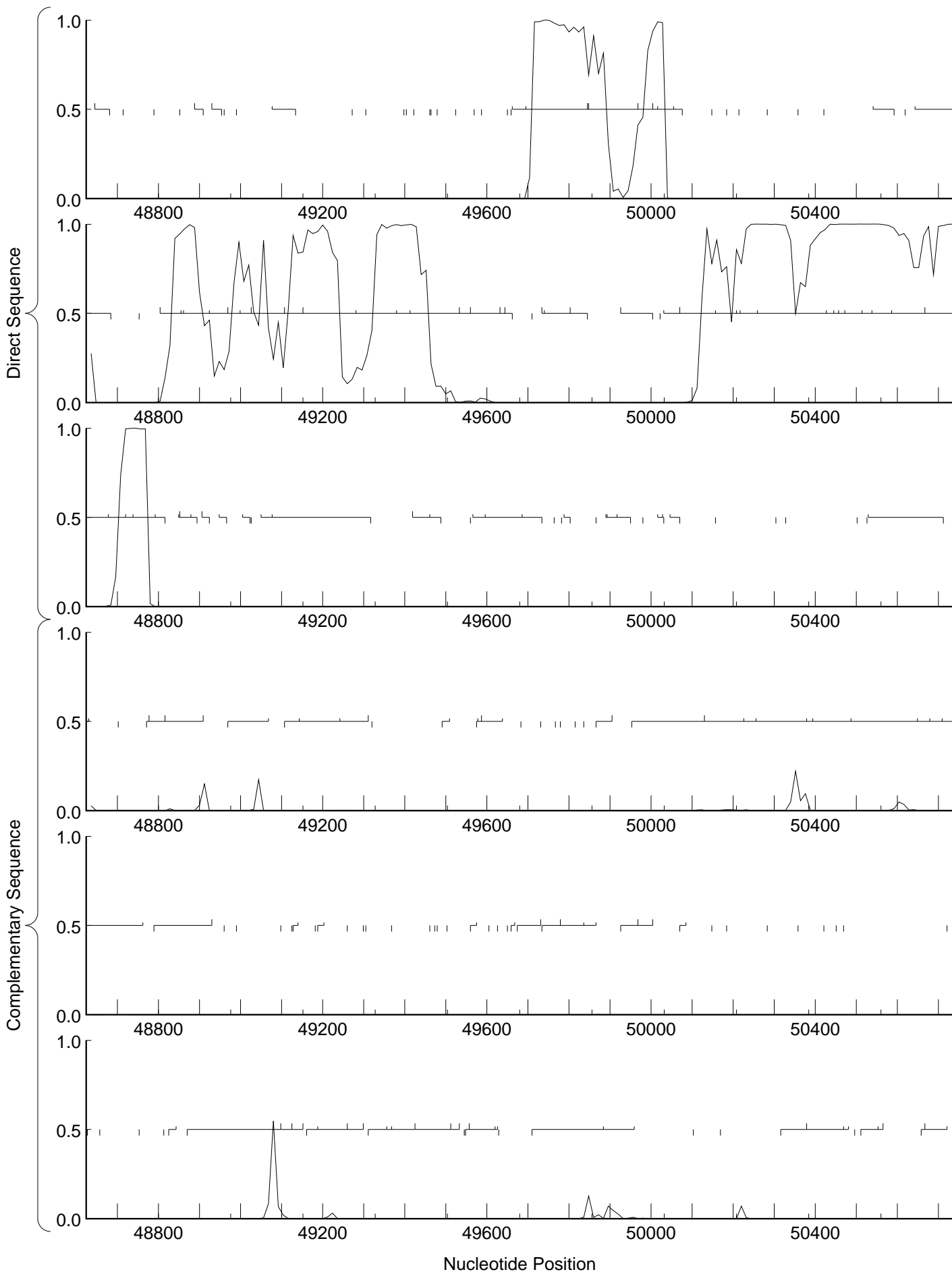


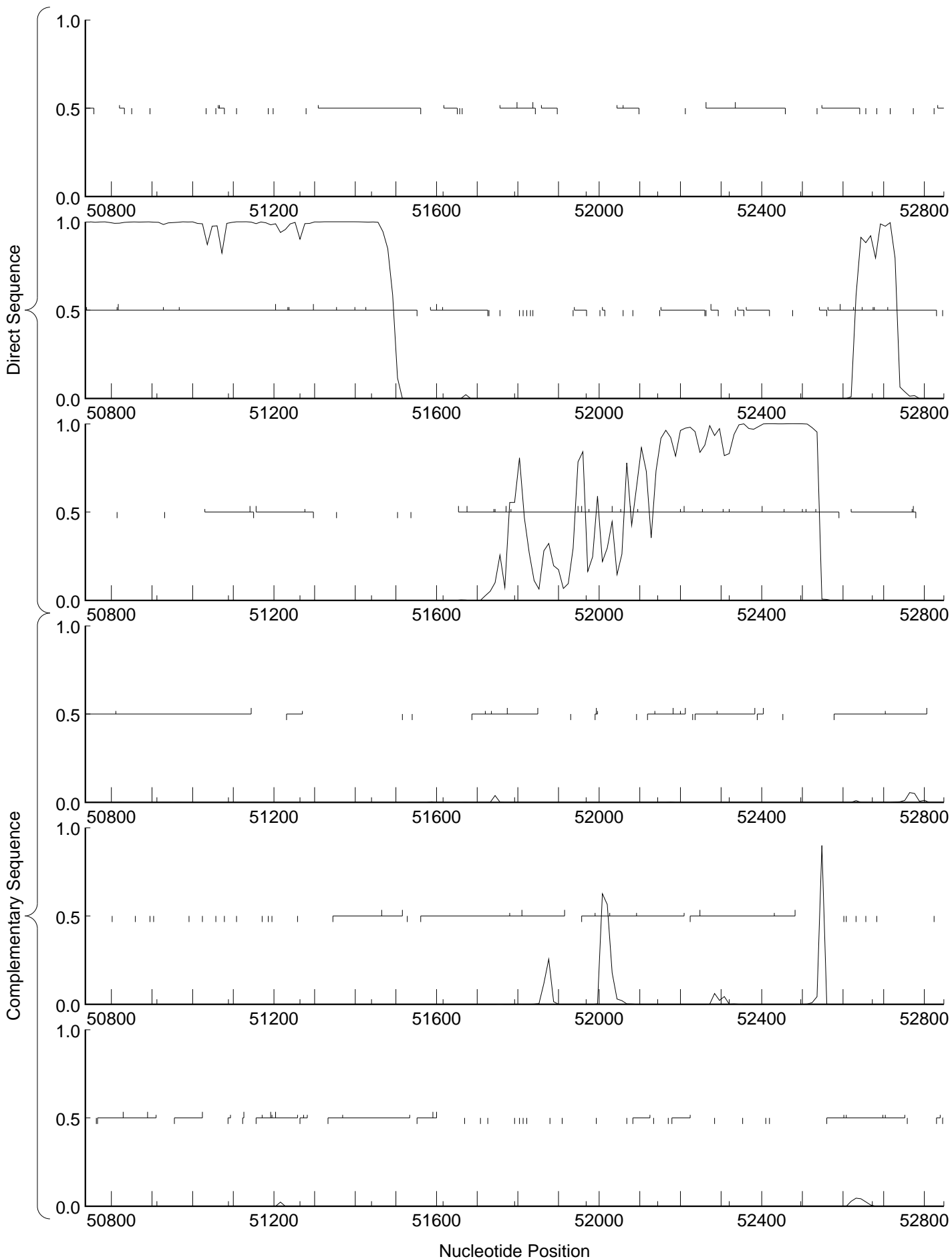


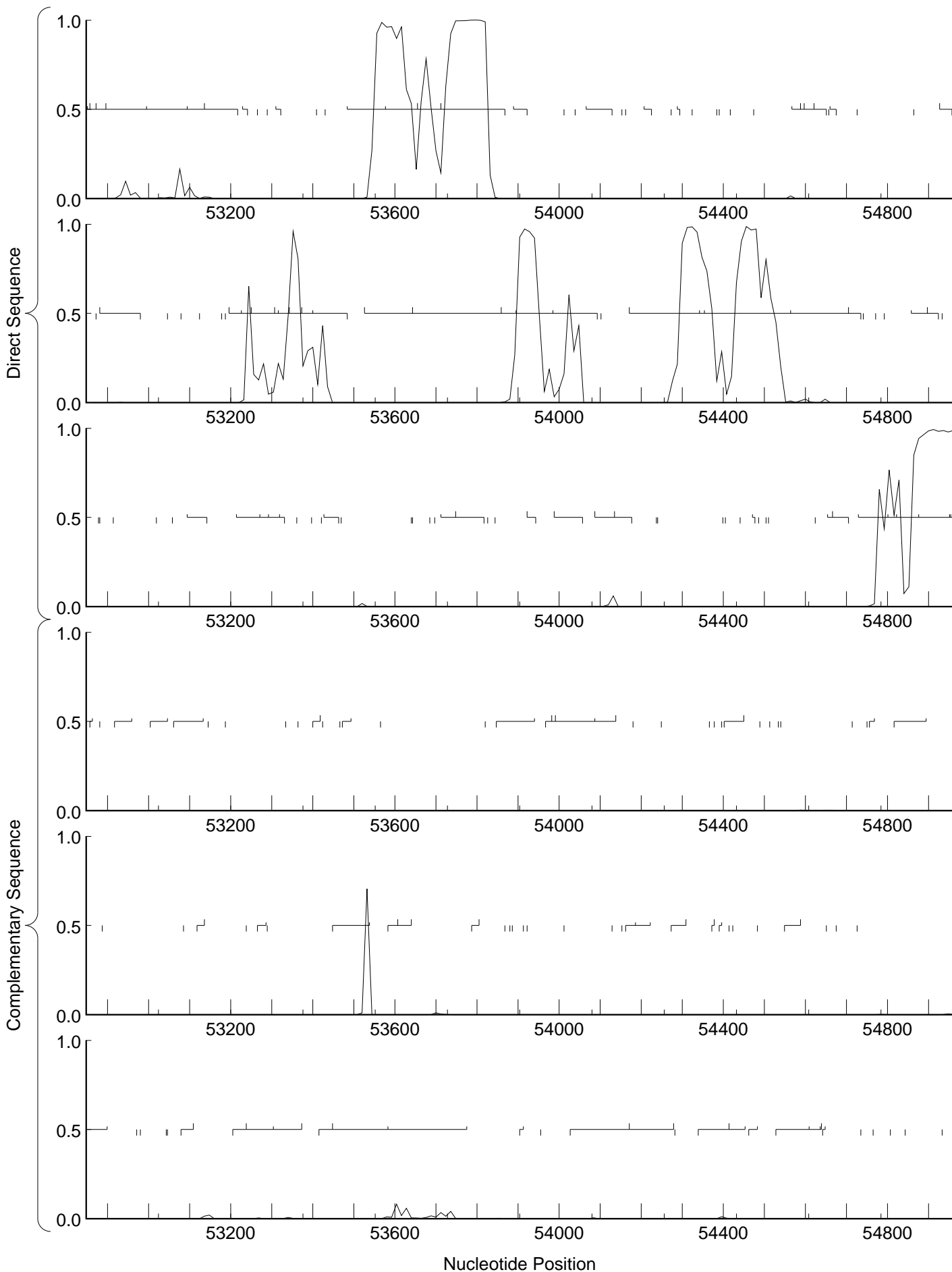


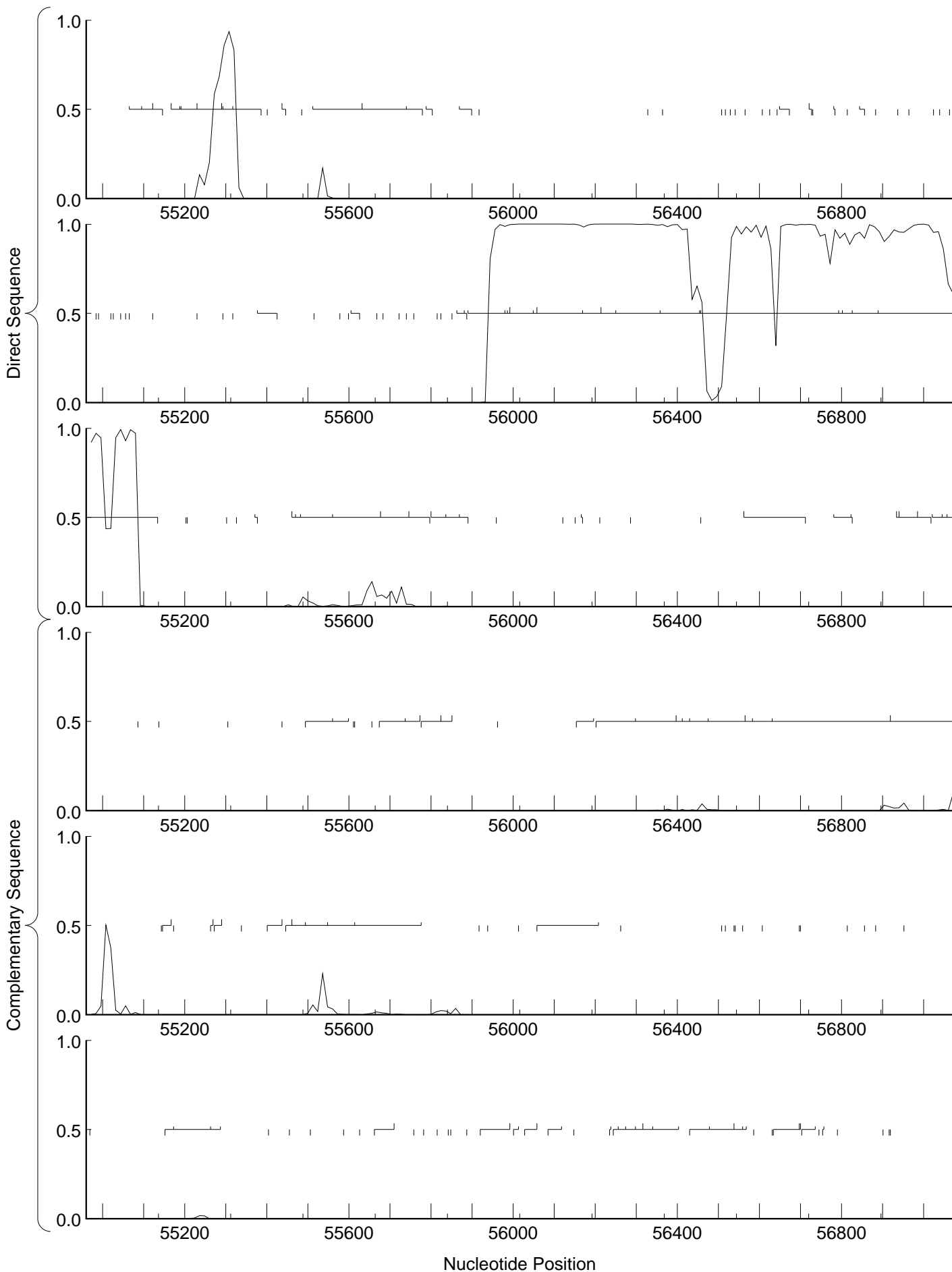


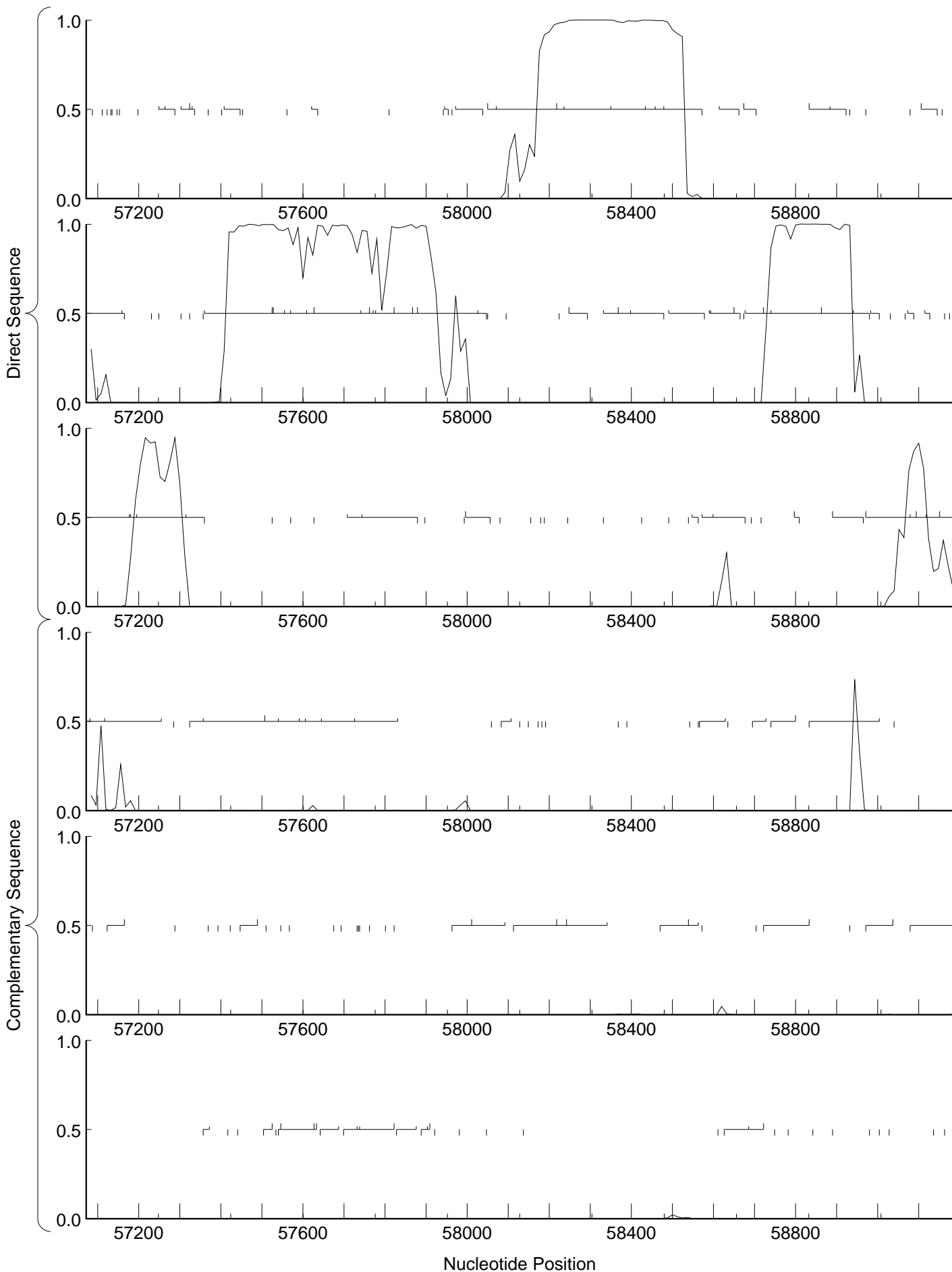




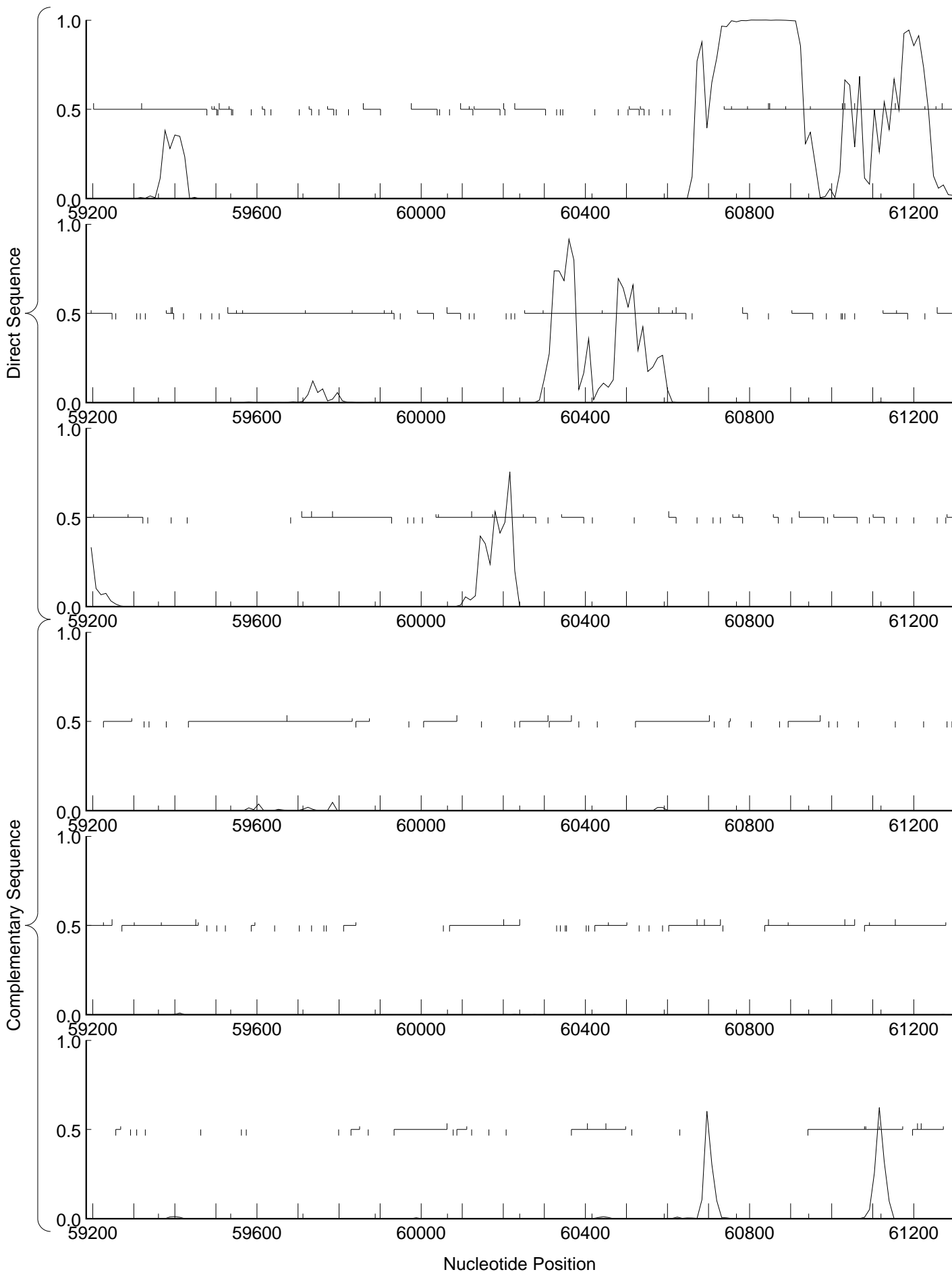


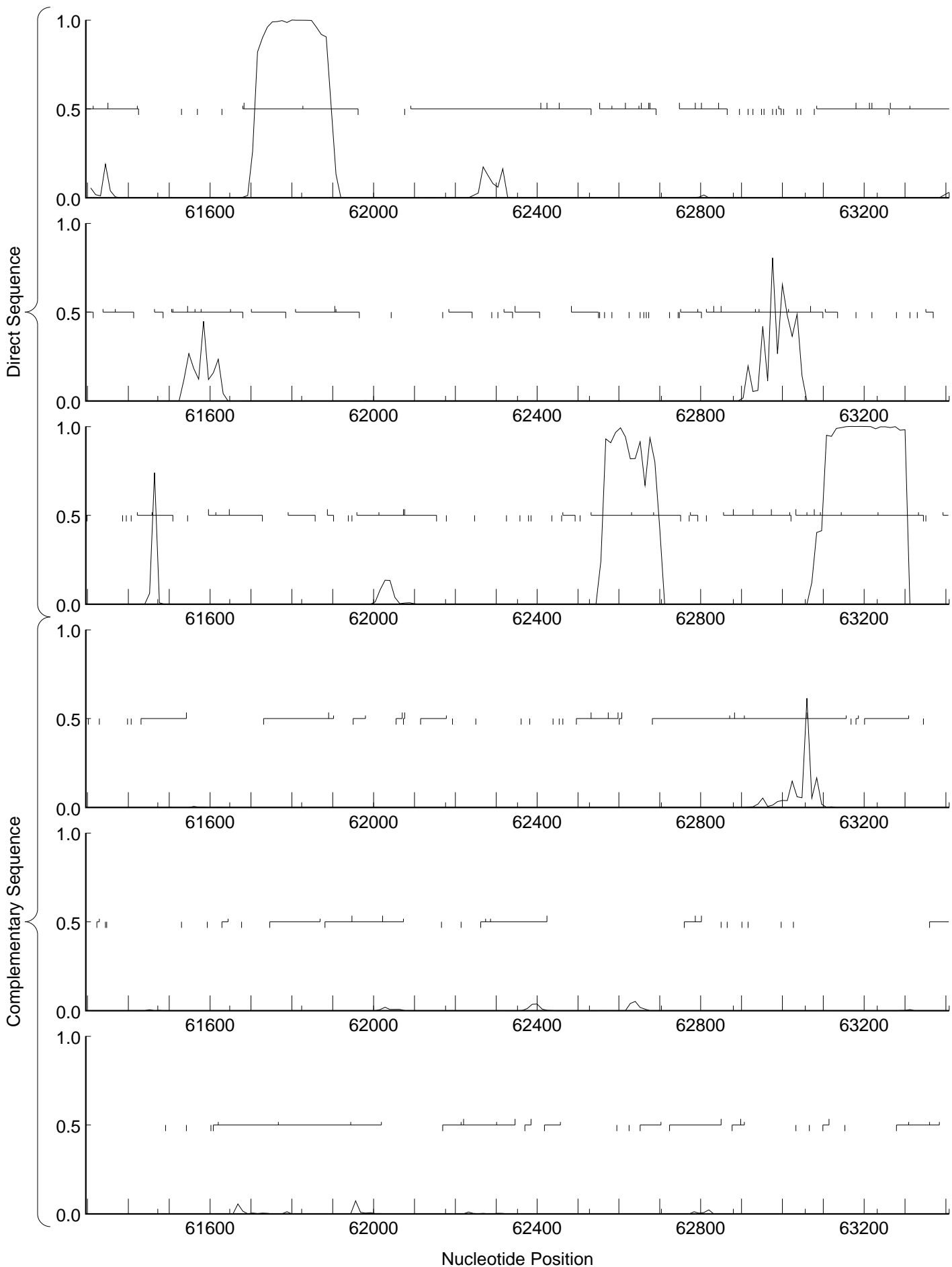


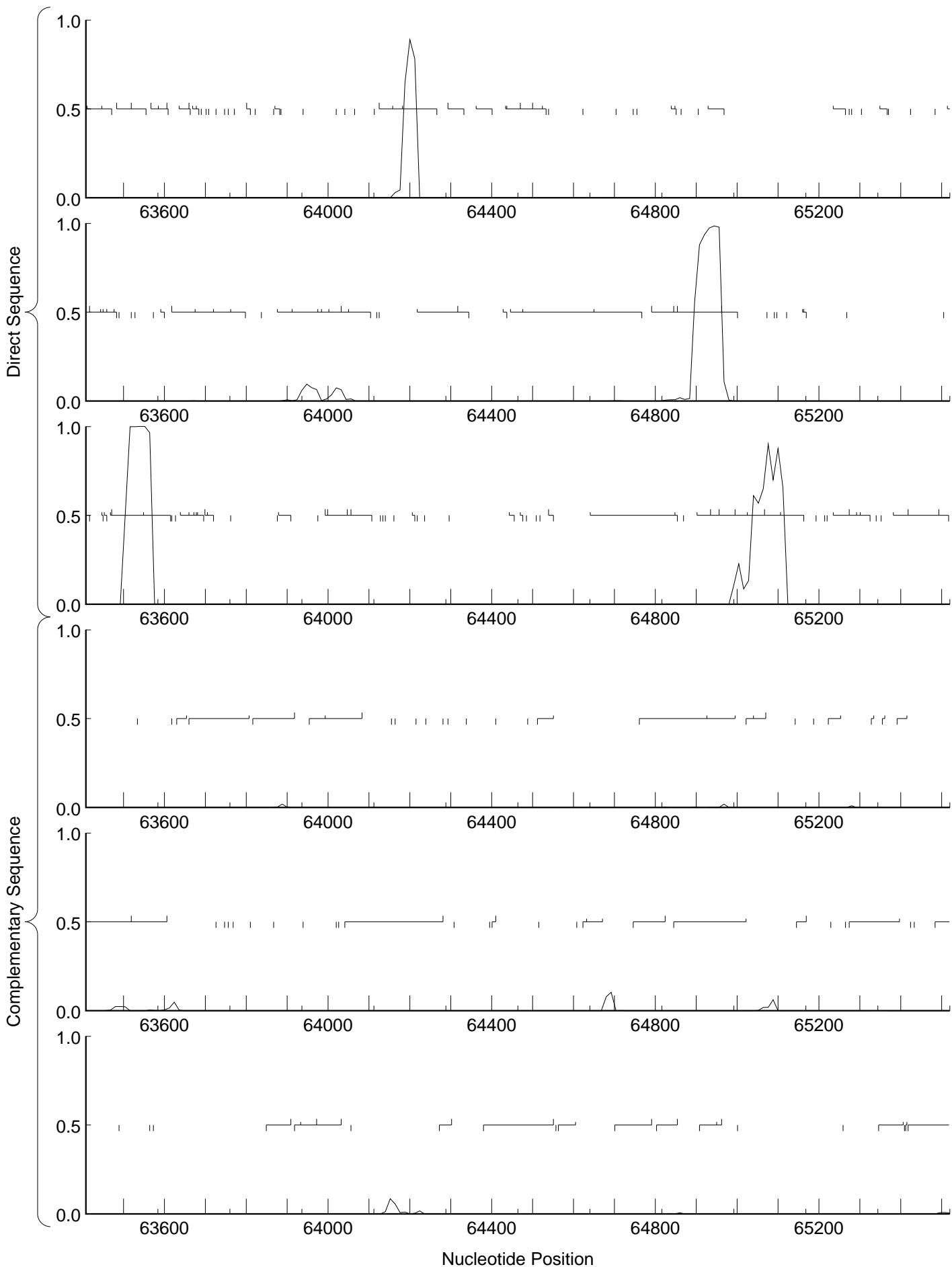


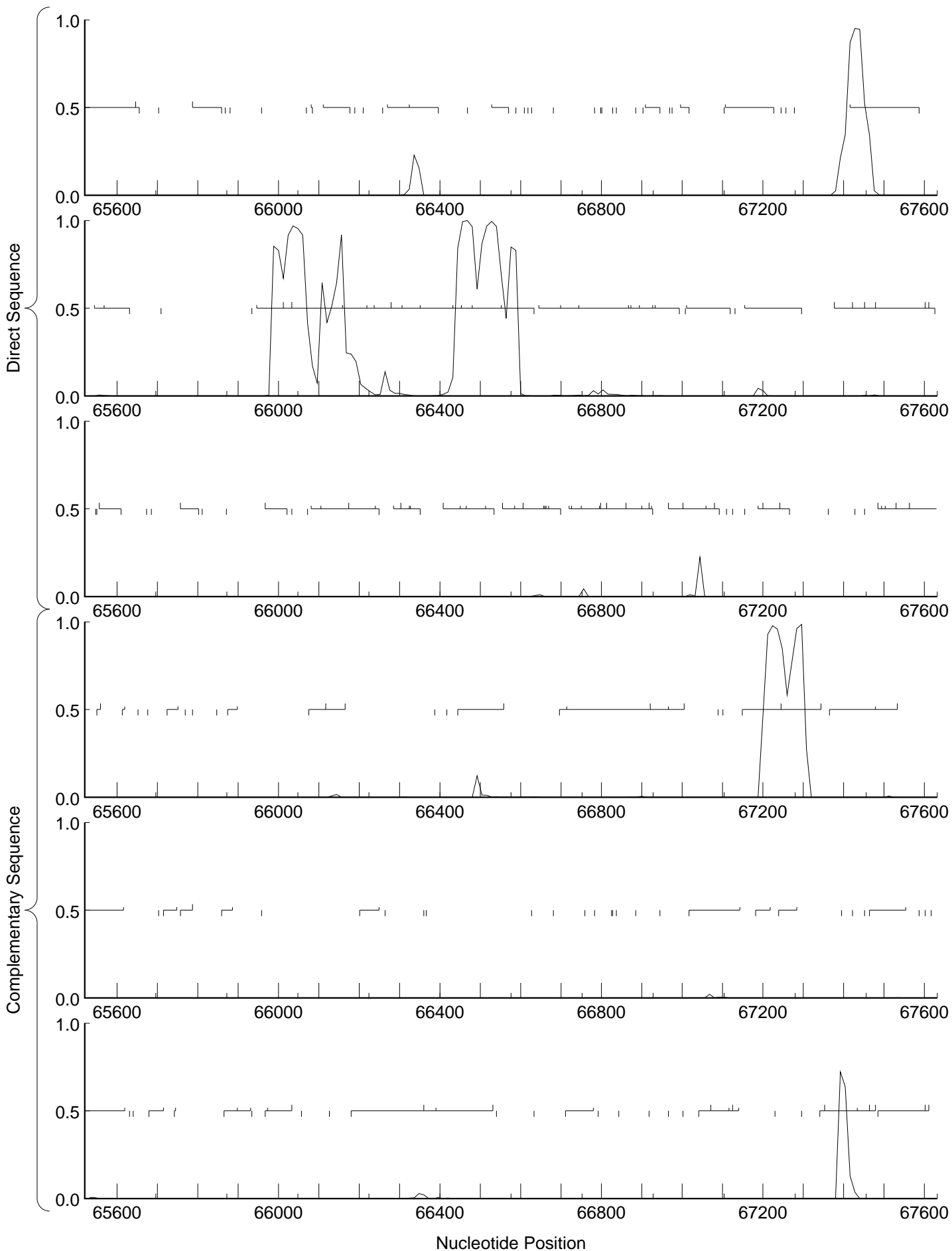


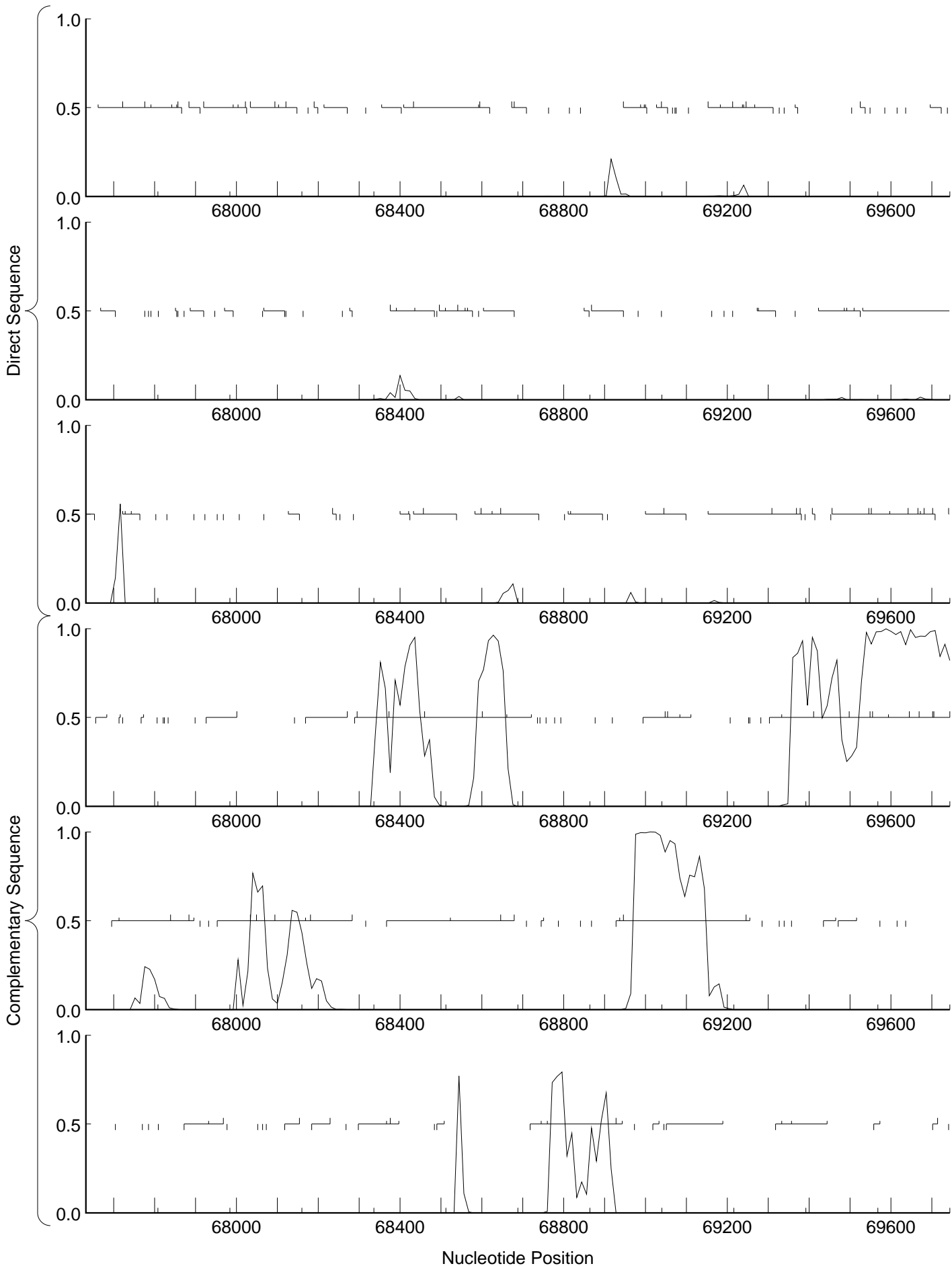
s complete sequence, 77566 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L4, Order 4, Window 96, Step 12, 30/38

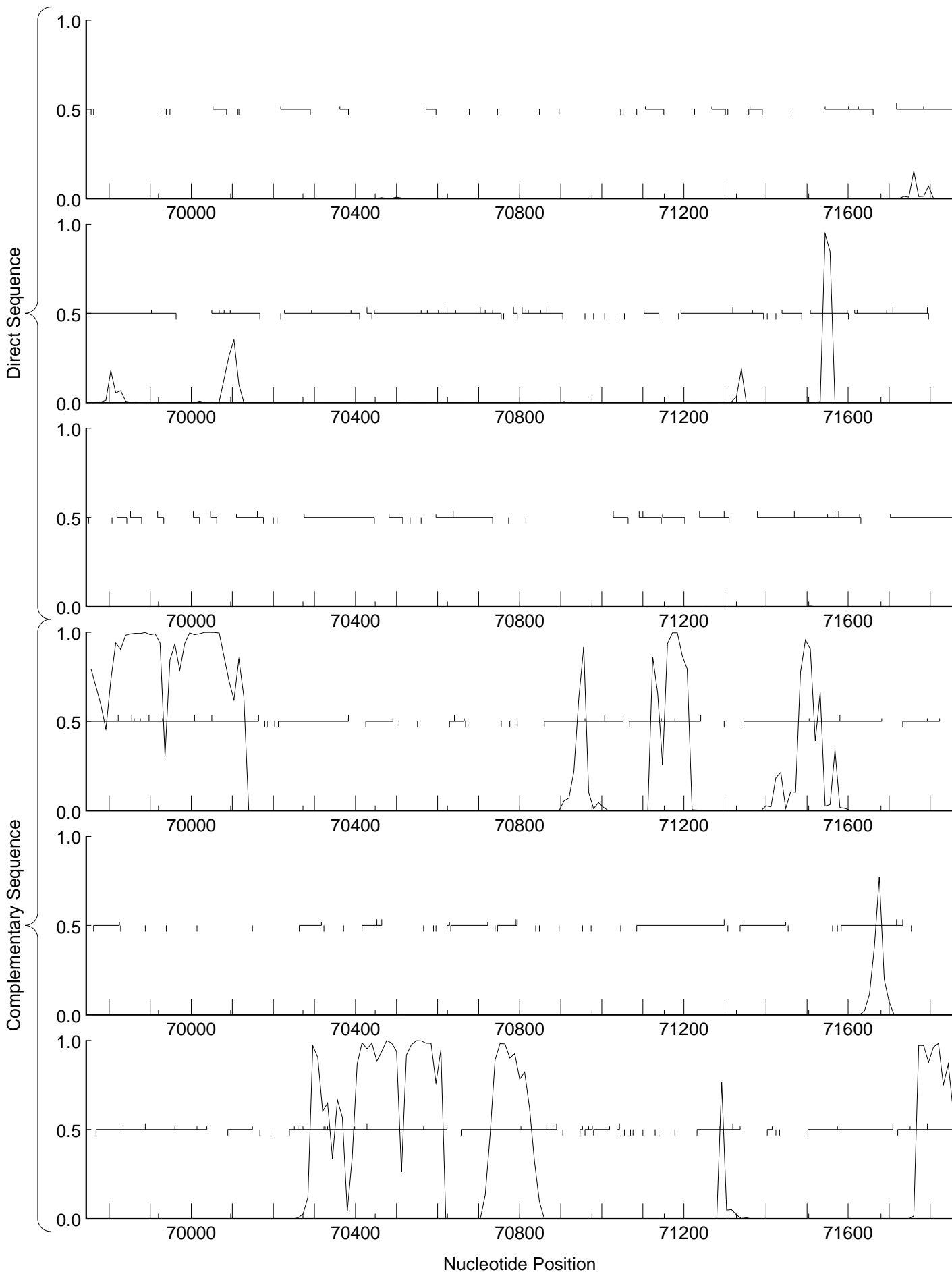


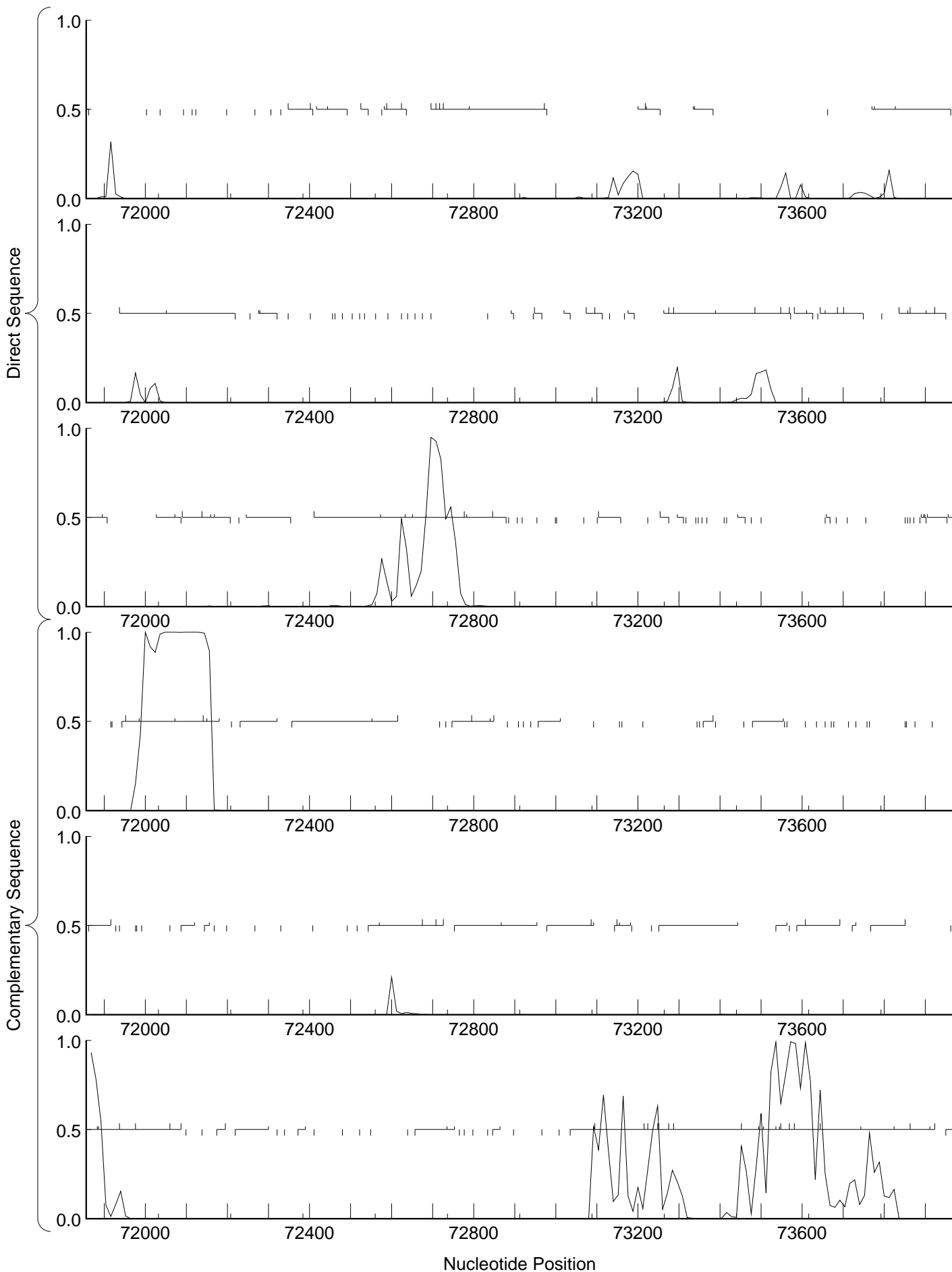












s complete sequence, 77566 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L4, Order 4, Window 96, Step 12, 37/38

