

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Vetrrix complete sequence, 76520 bp including 10-base 3' overhang (TCGAT
Analysis Date : 2/5/23 at 0:07:41
Pages : 38
Sequence Length : 76520 bp
GC Content : 58.94%

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

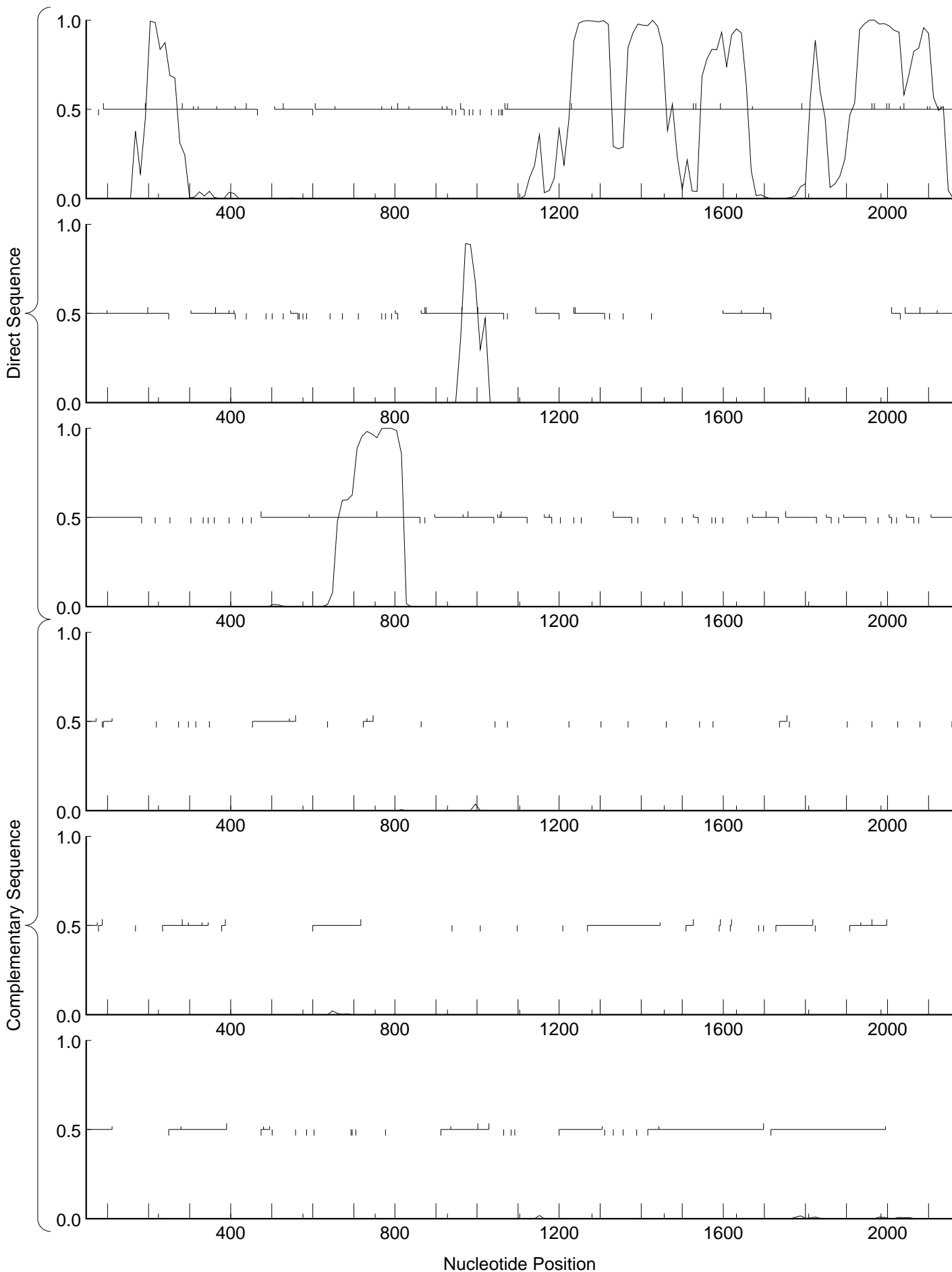
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

MATRIX INFORMATION

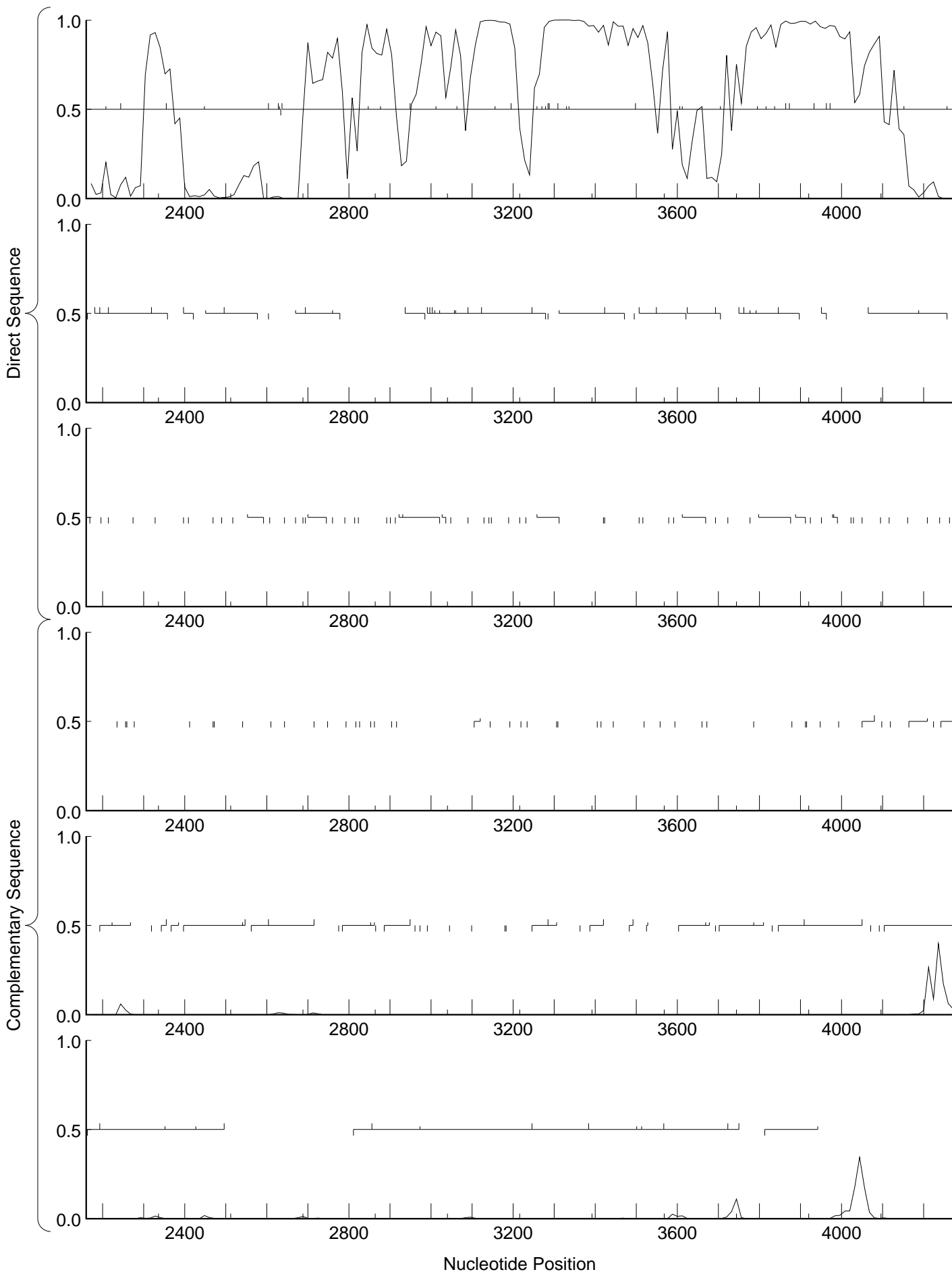
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

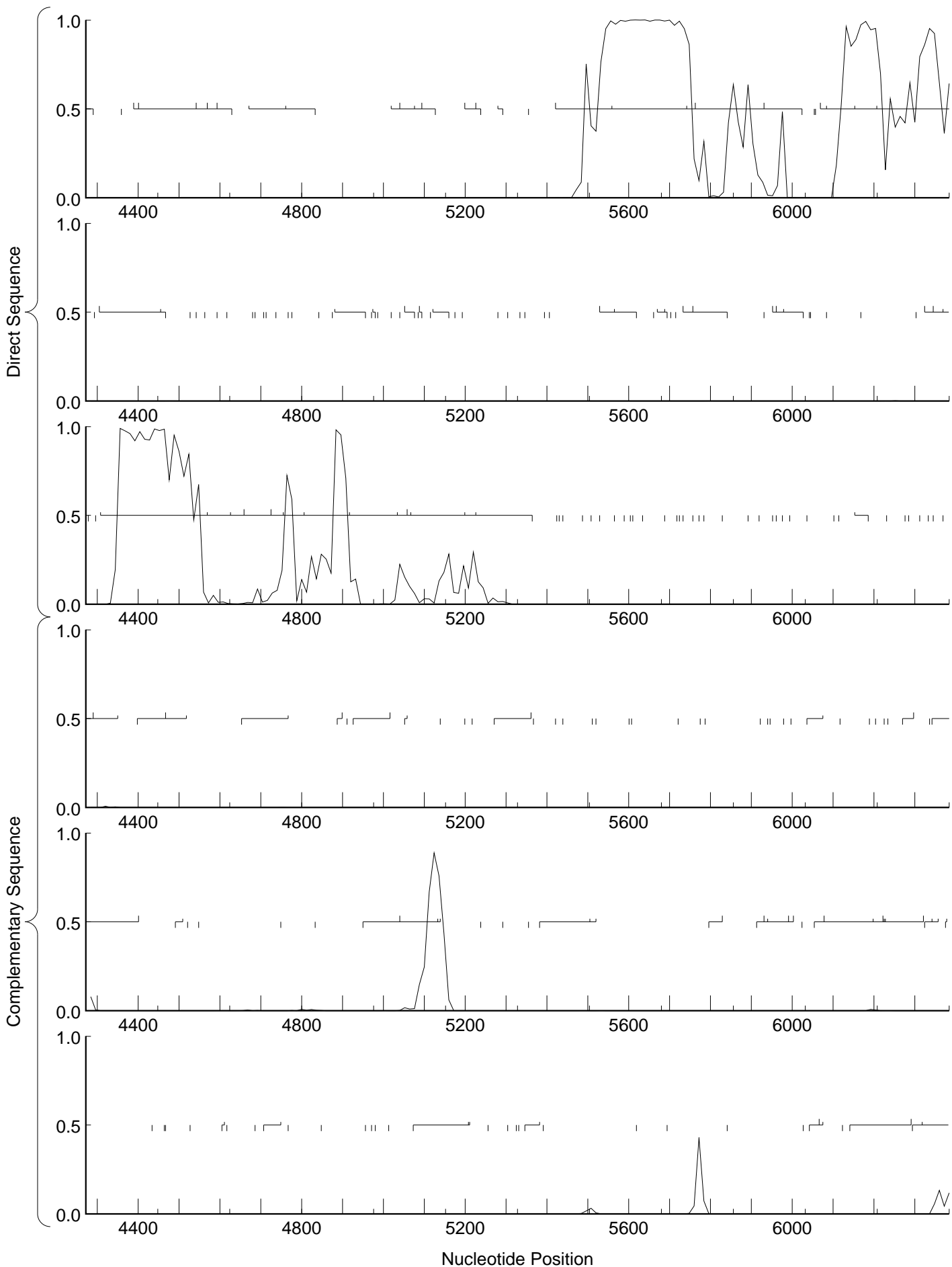
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

rix complete sequence, 76520 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 2/38

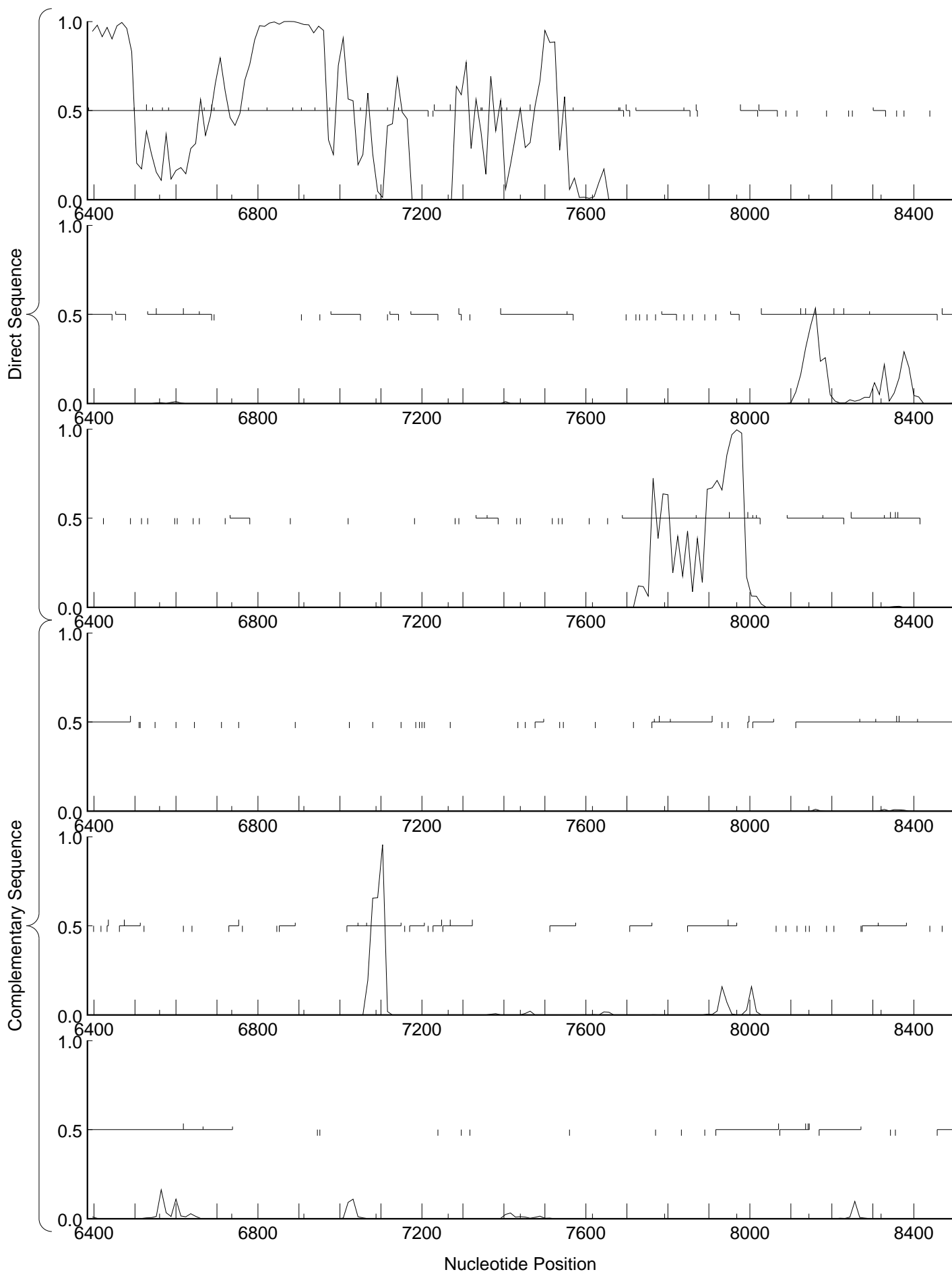


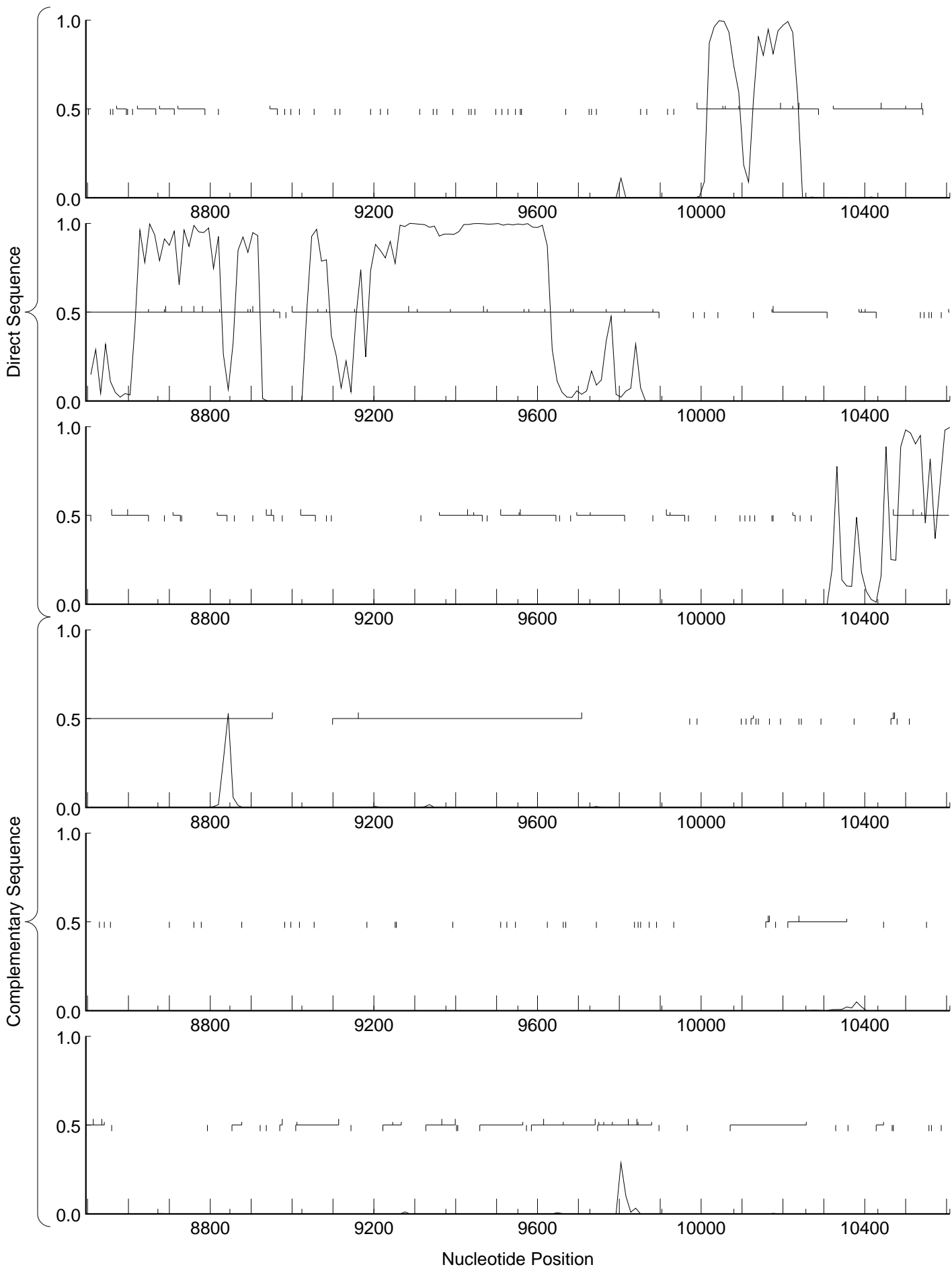
ix complete sequence, 76520 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 3/38

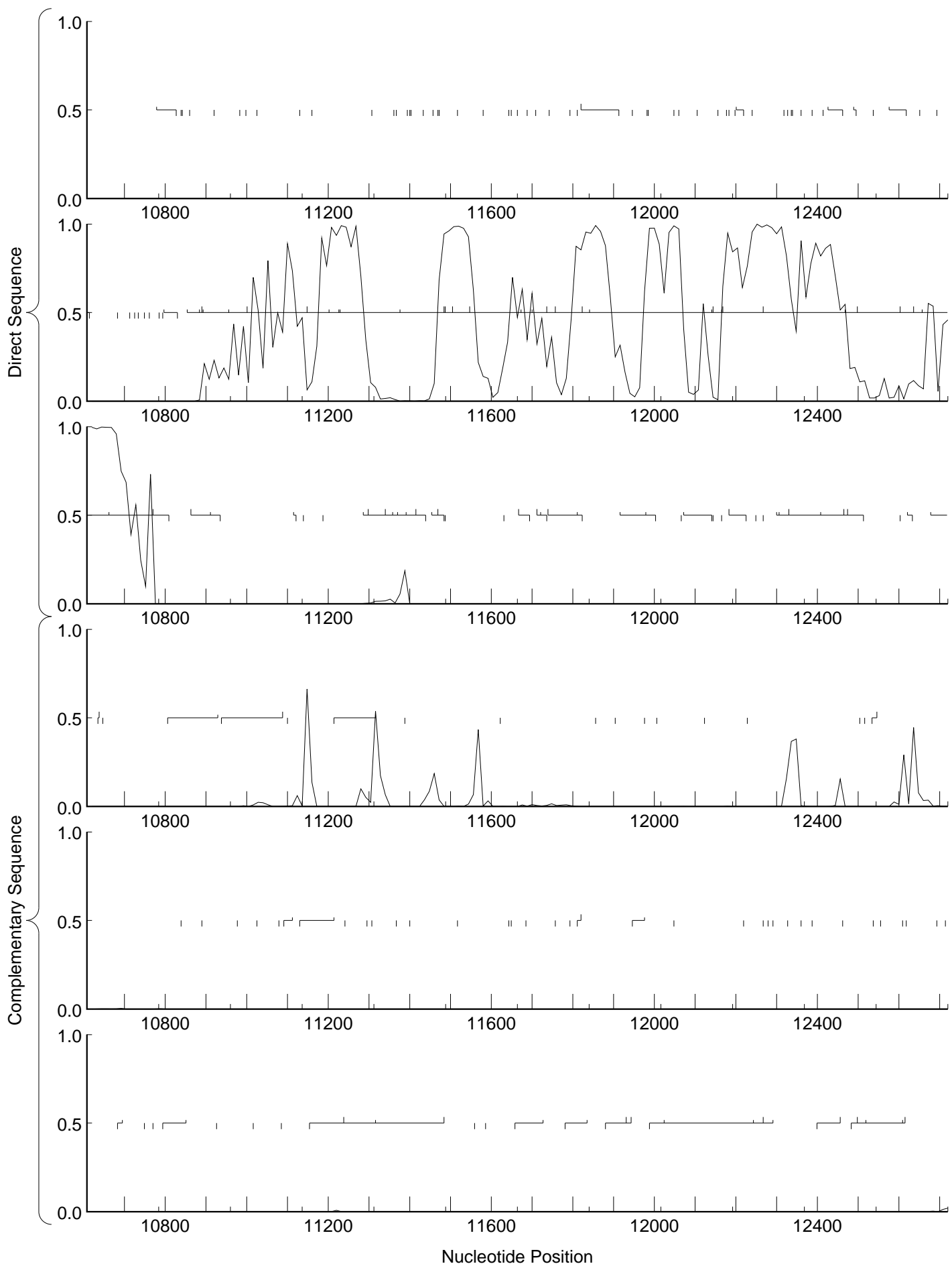


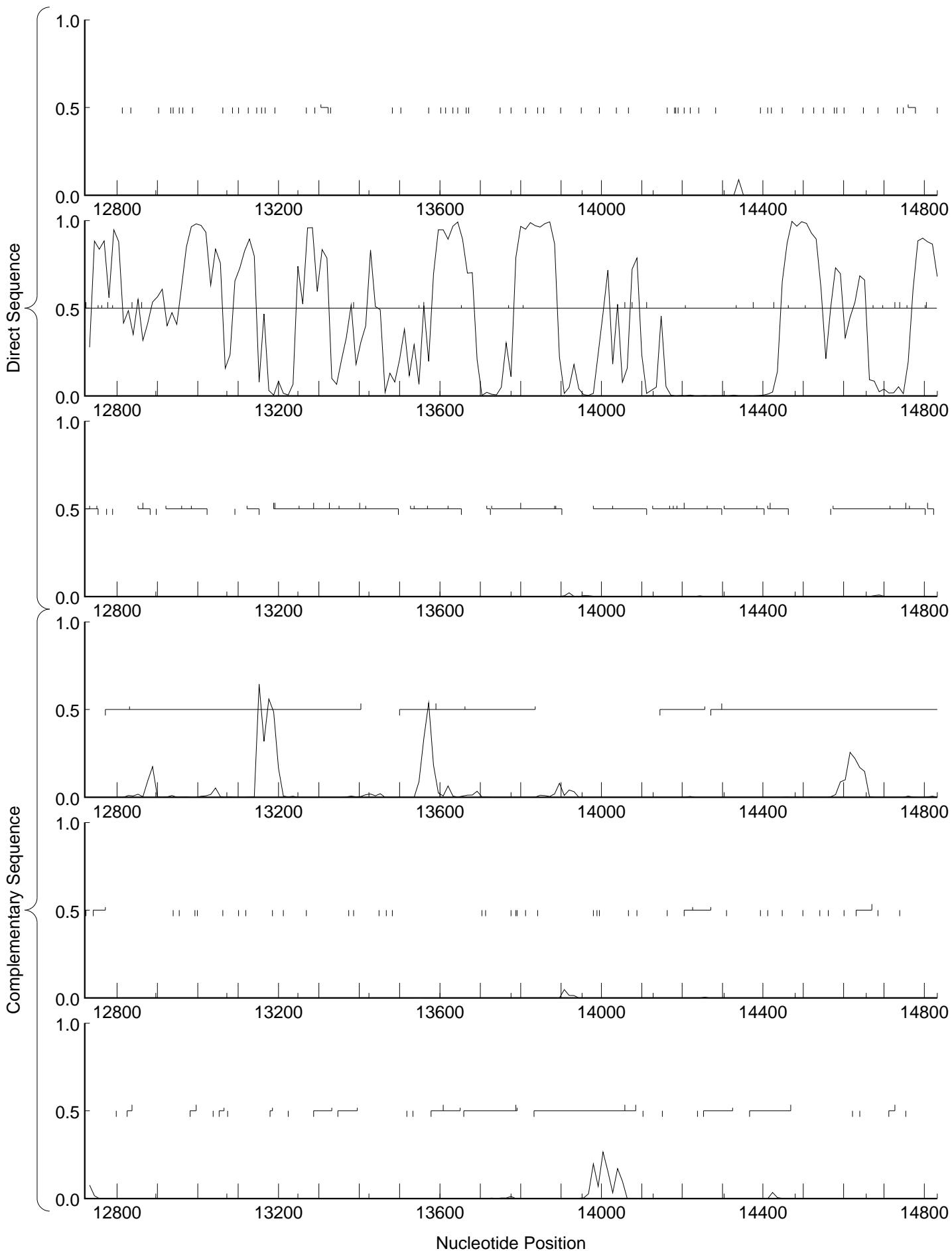


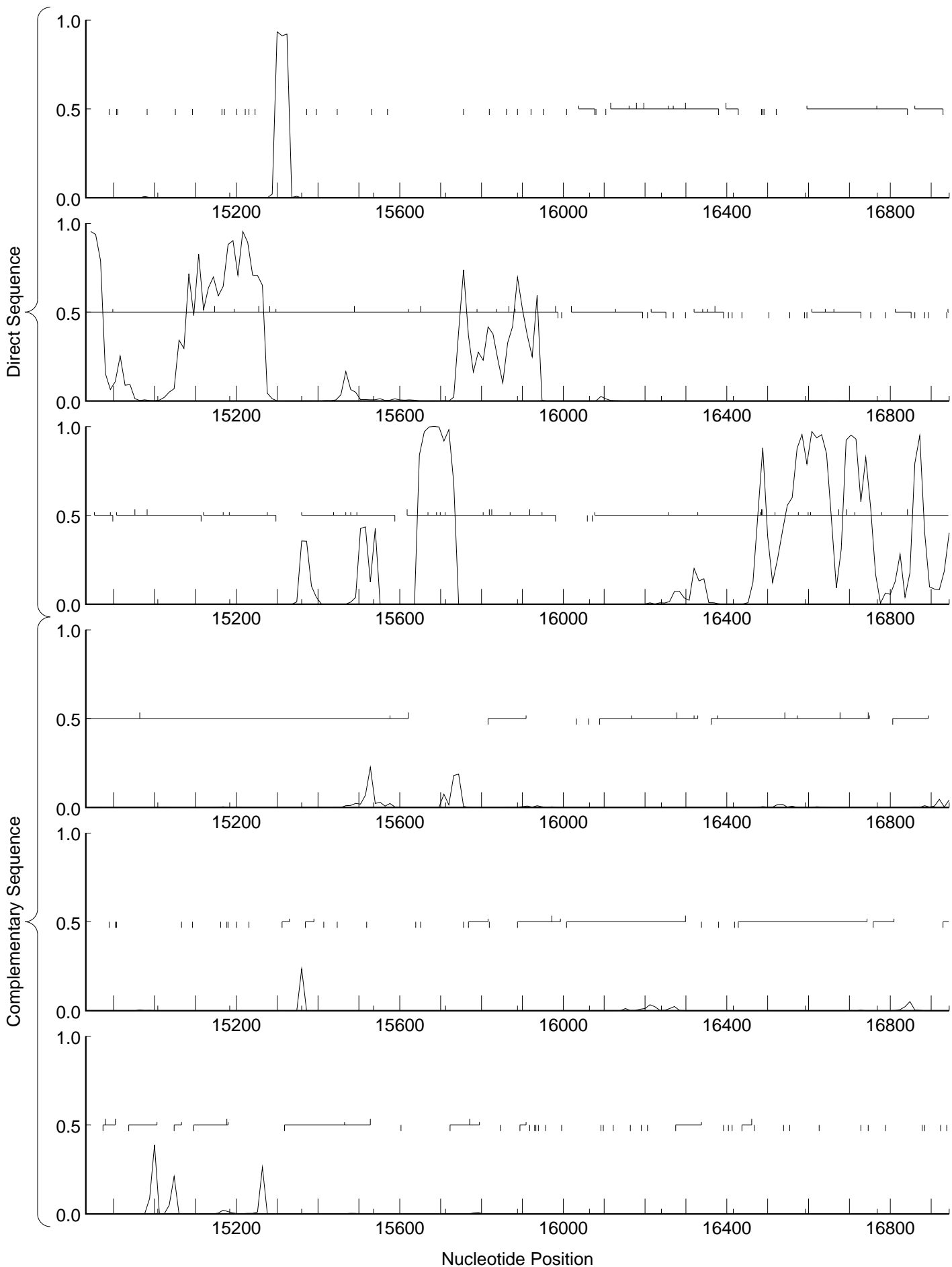
ix complete sequence, 76520 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 5/38

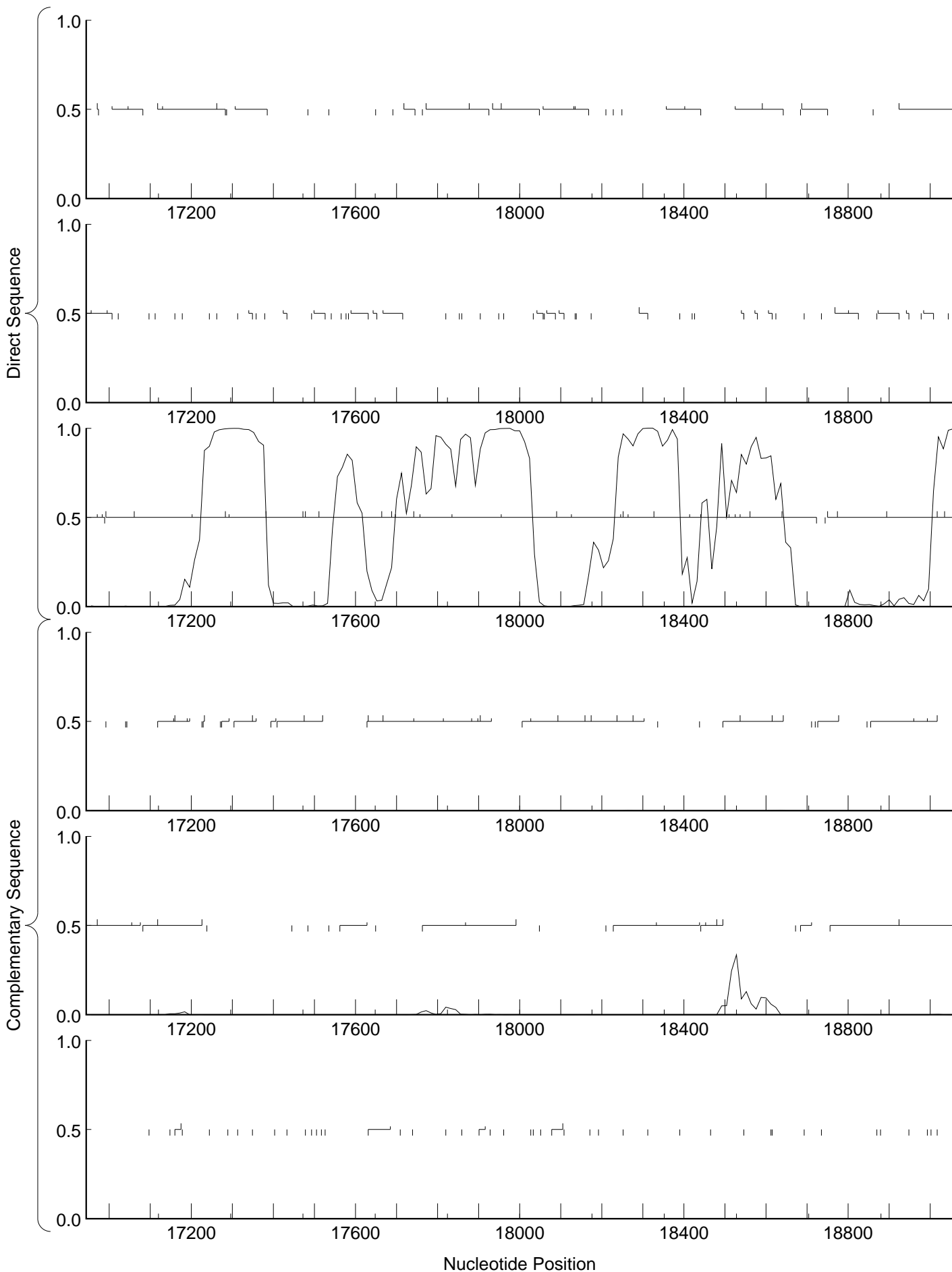


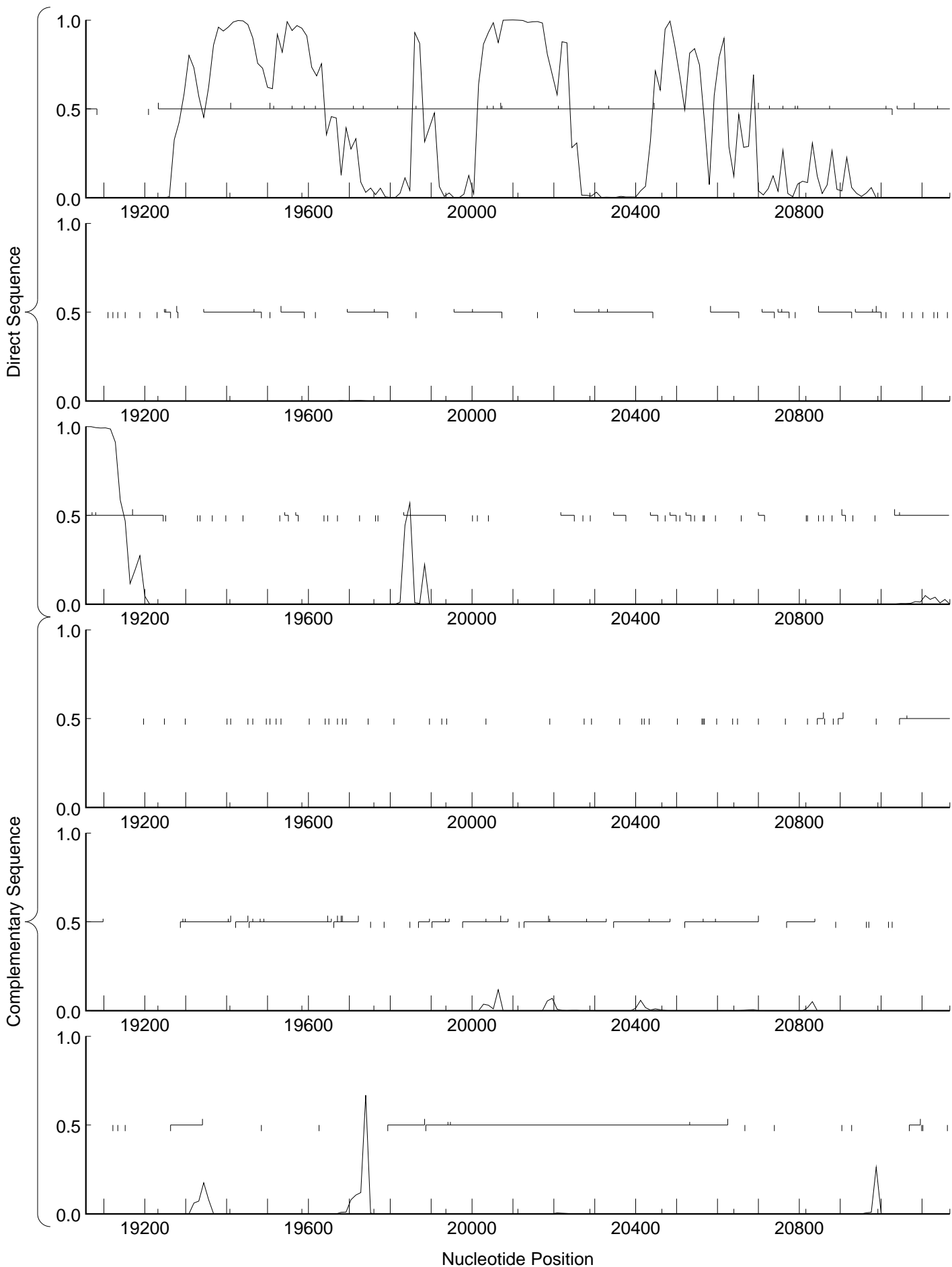


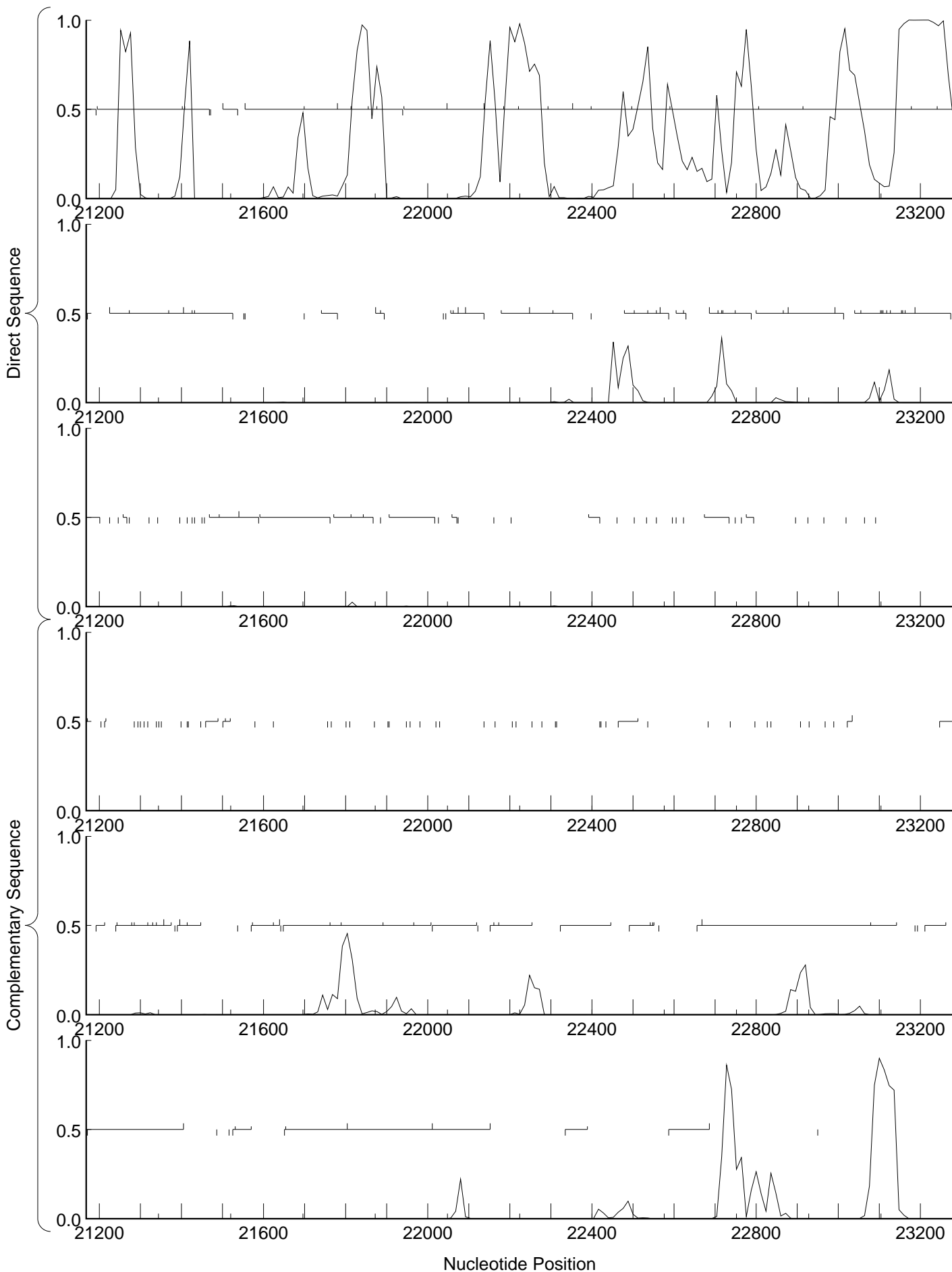


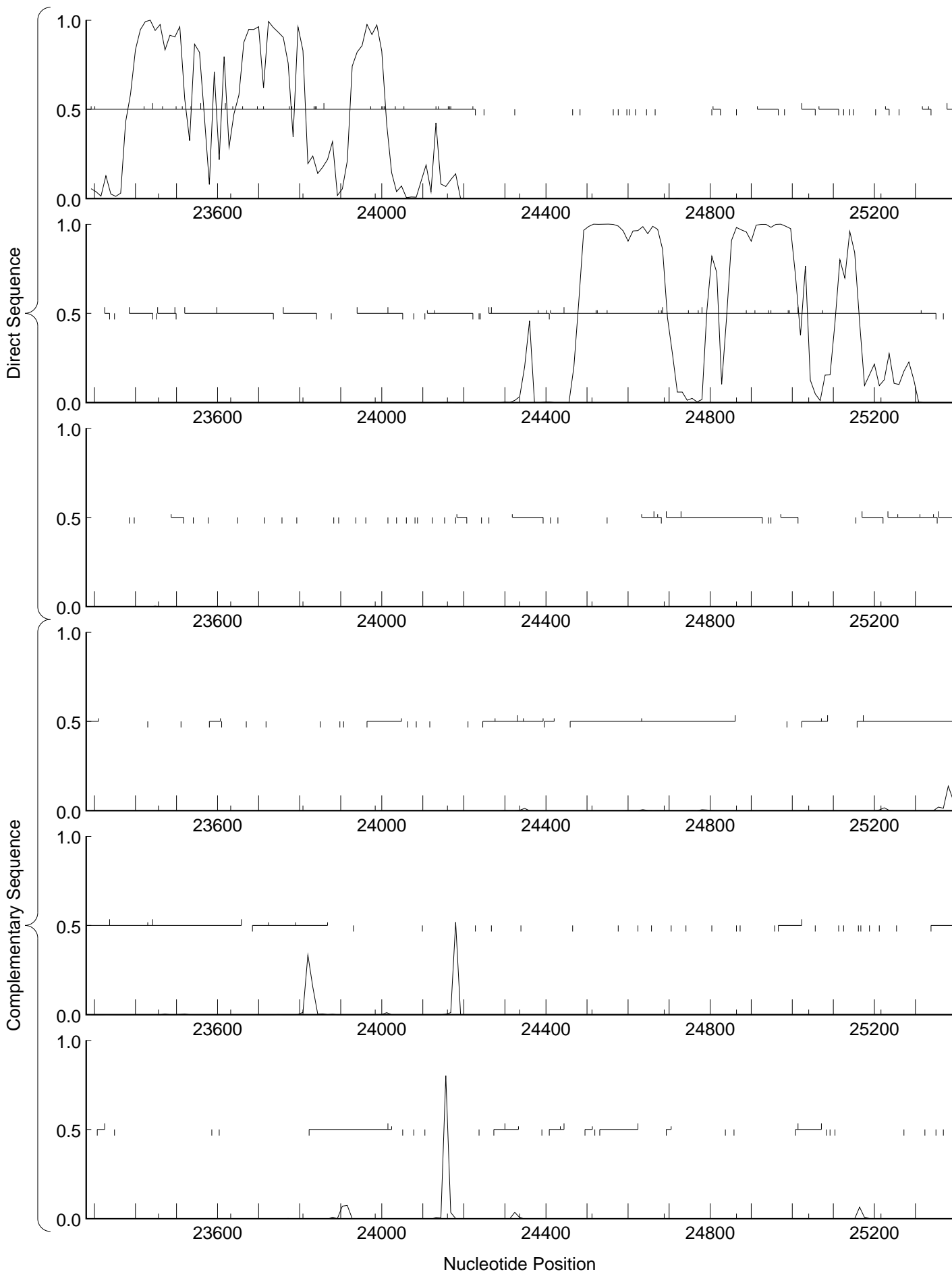


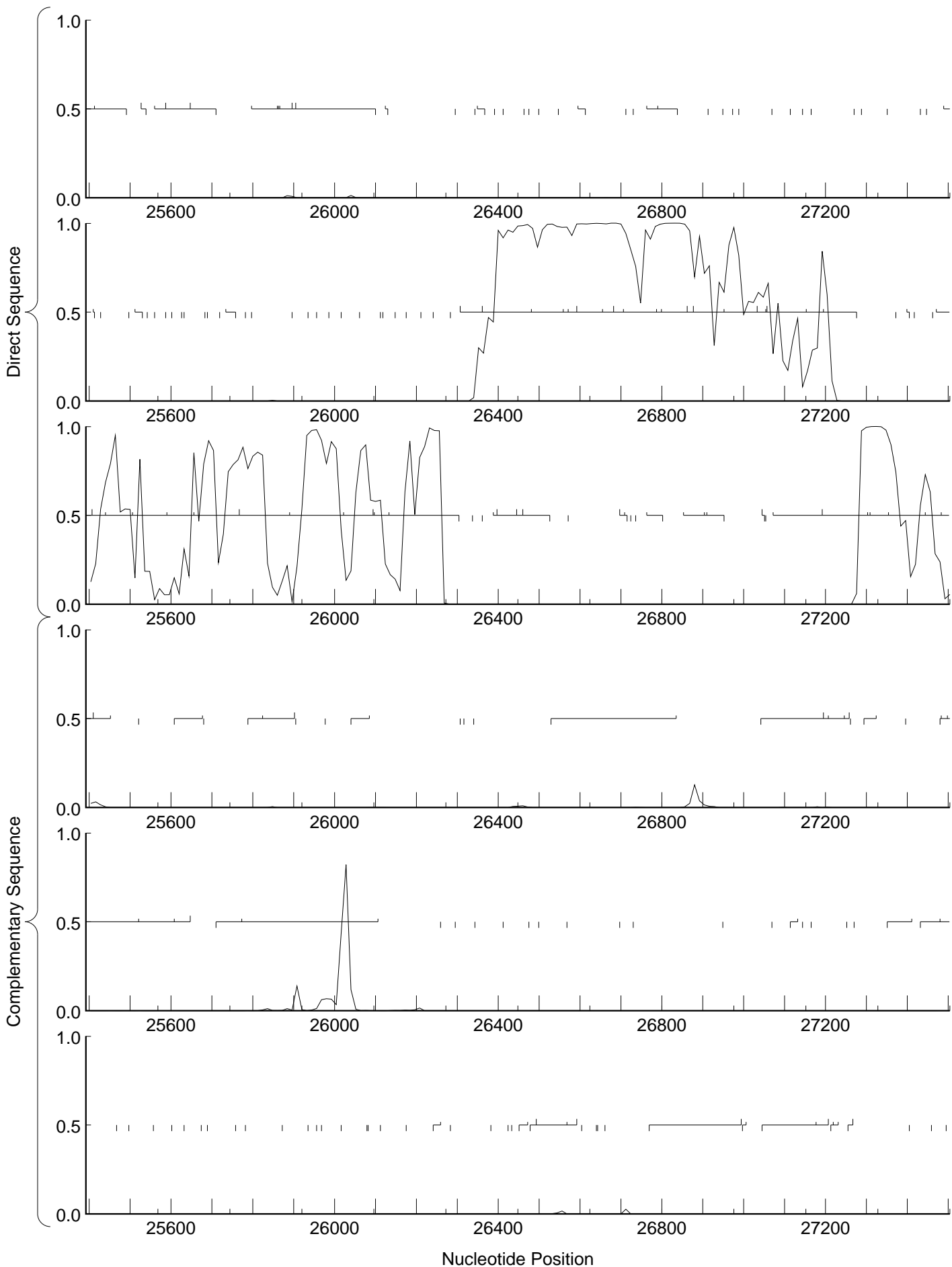


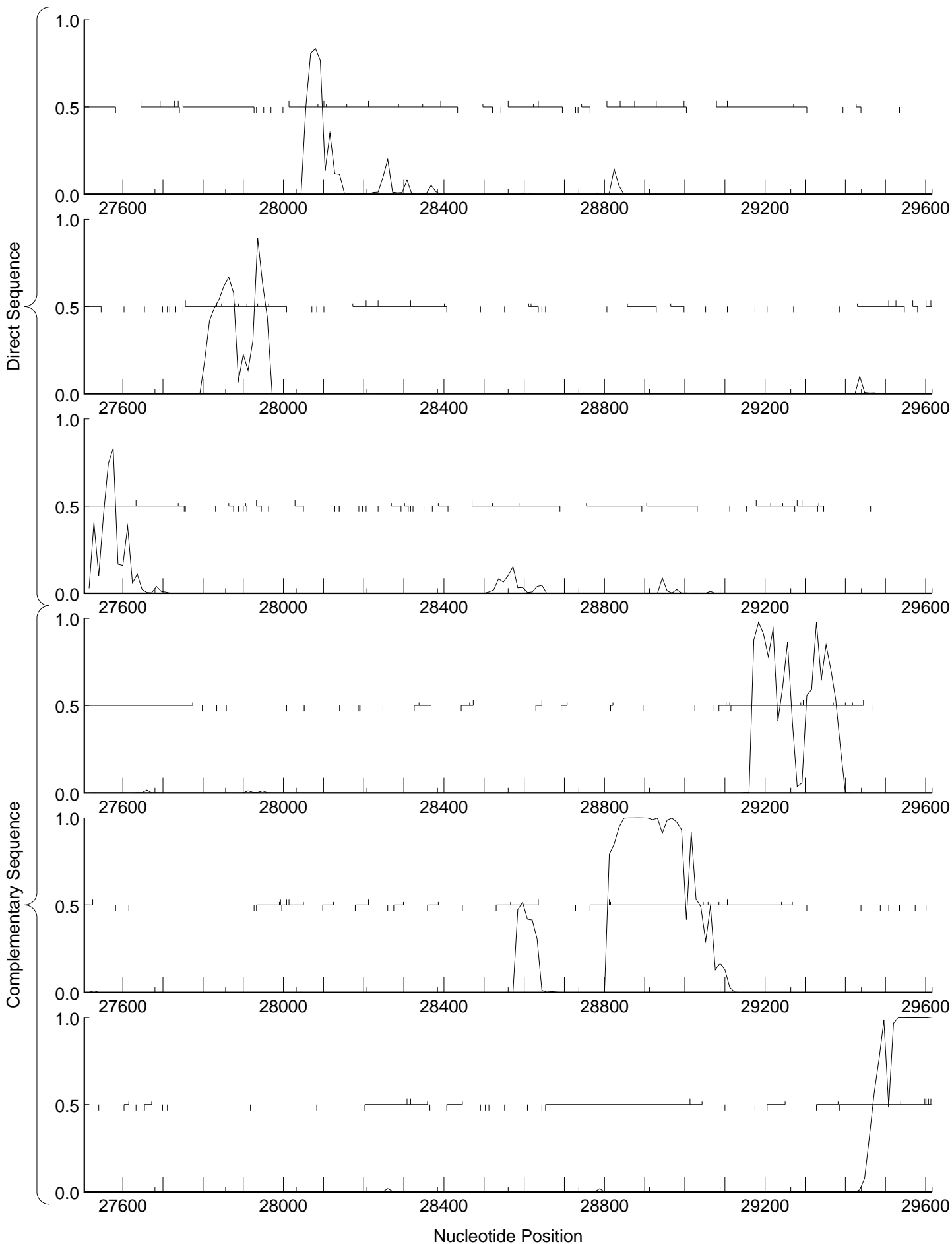


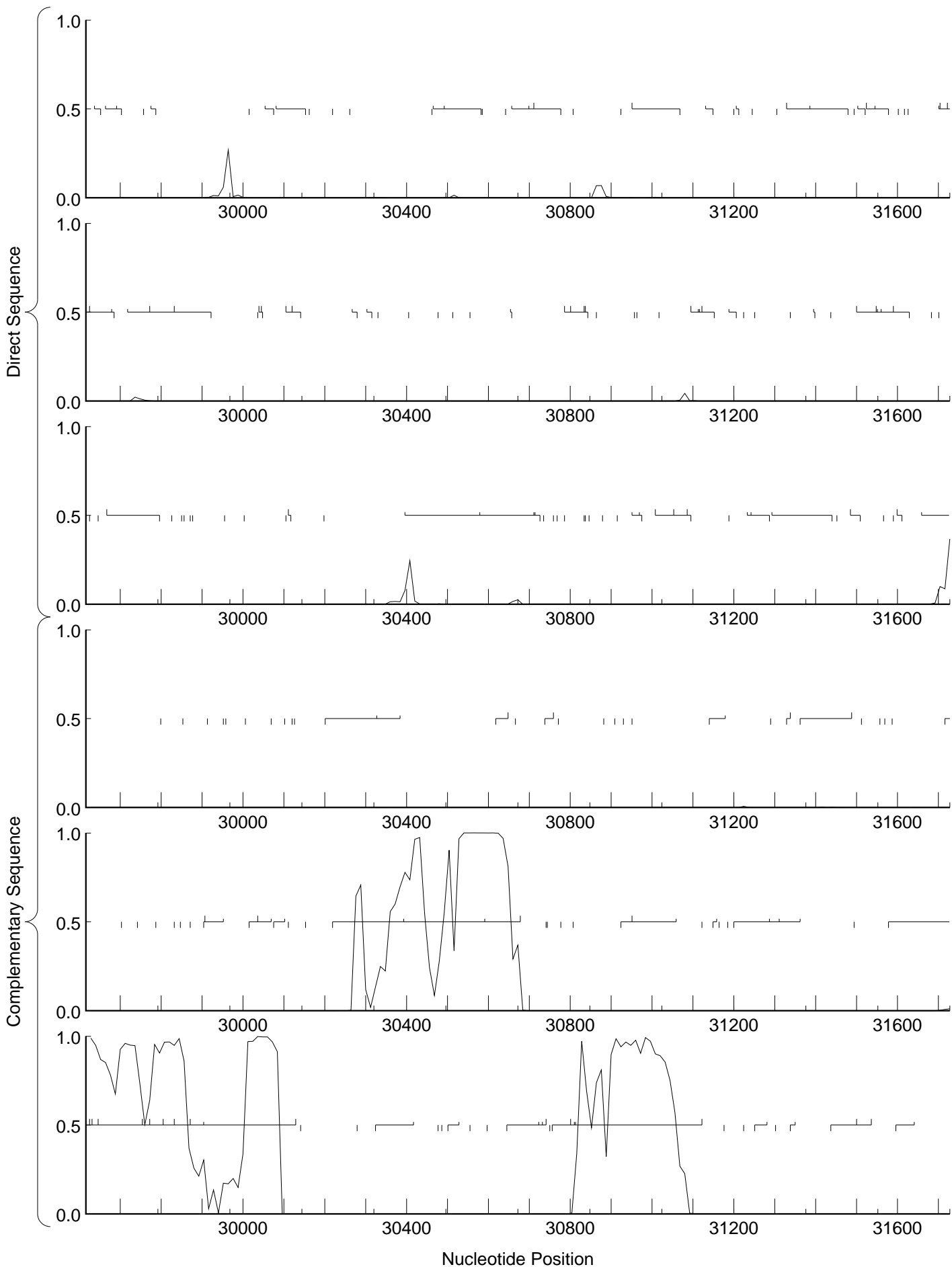


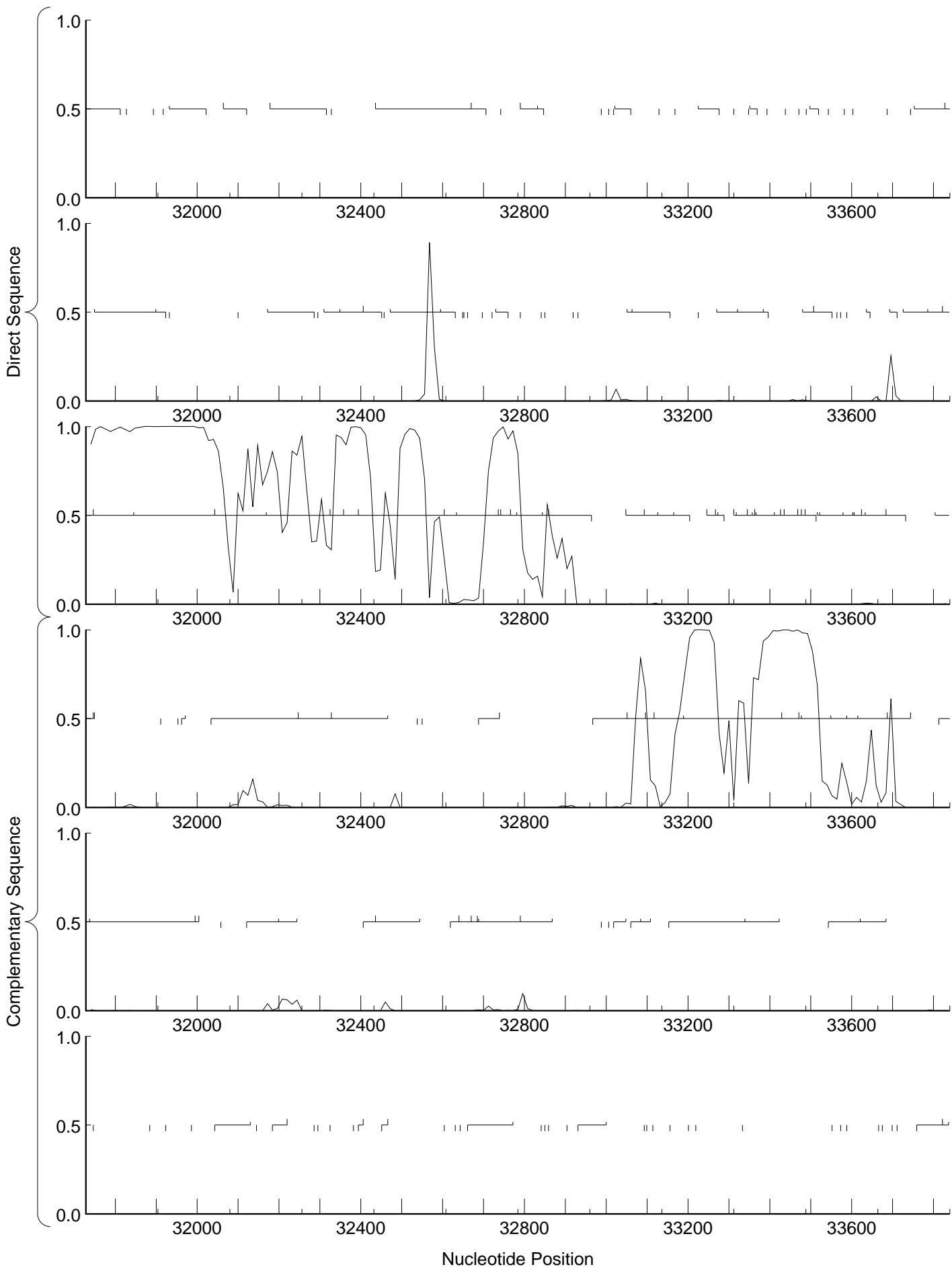


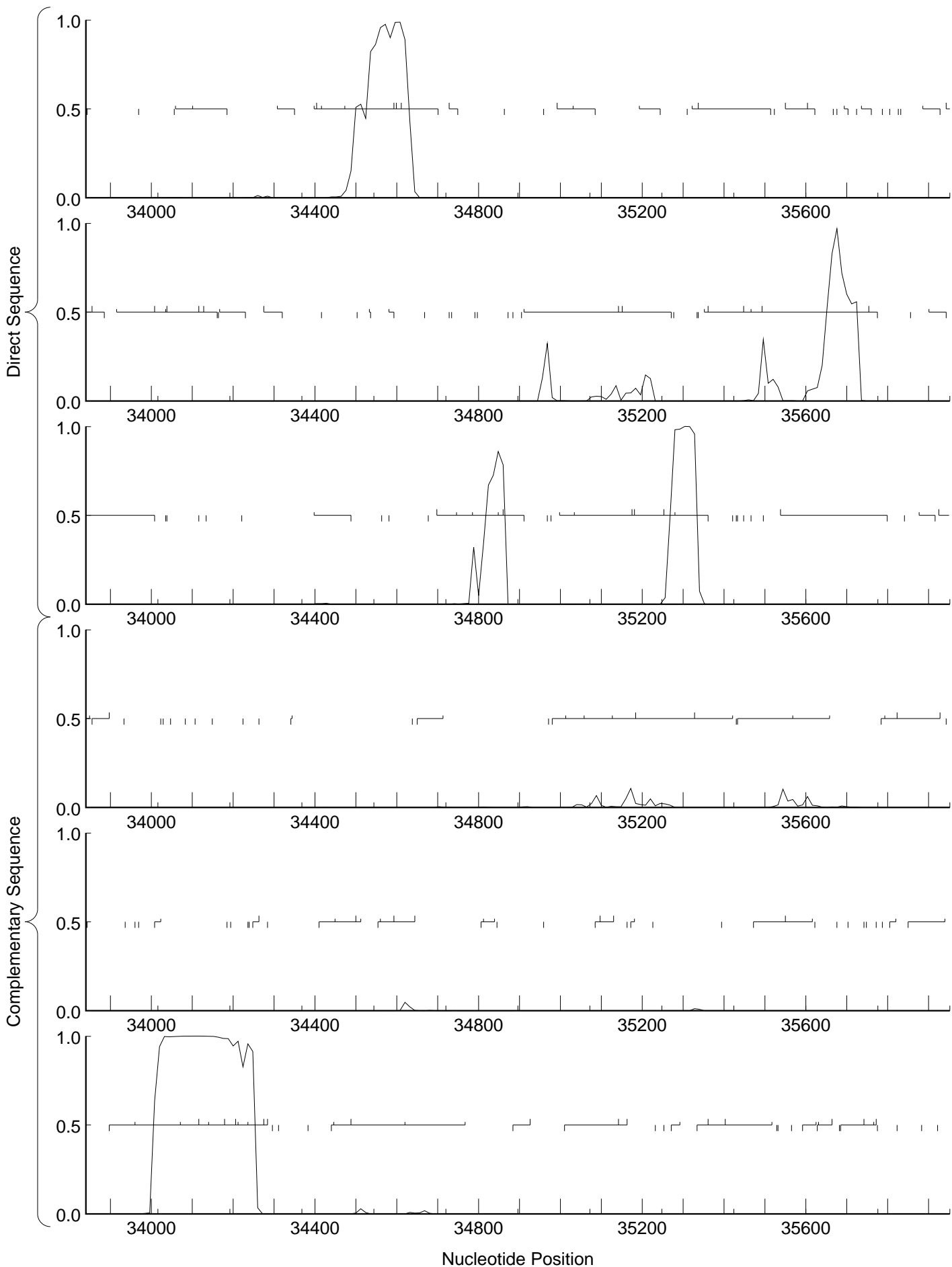


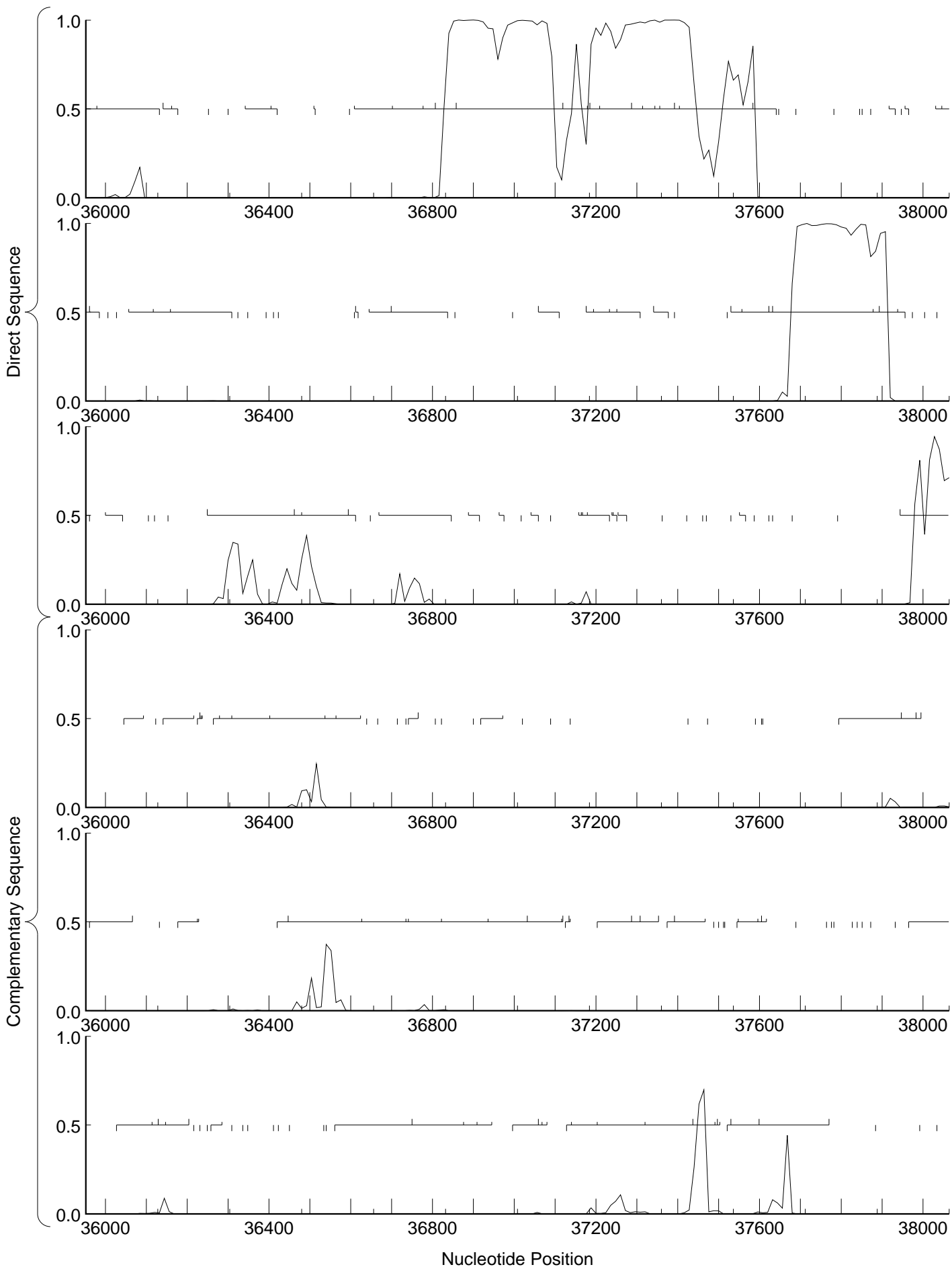


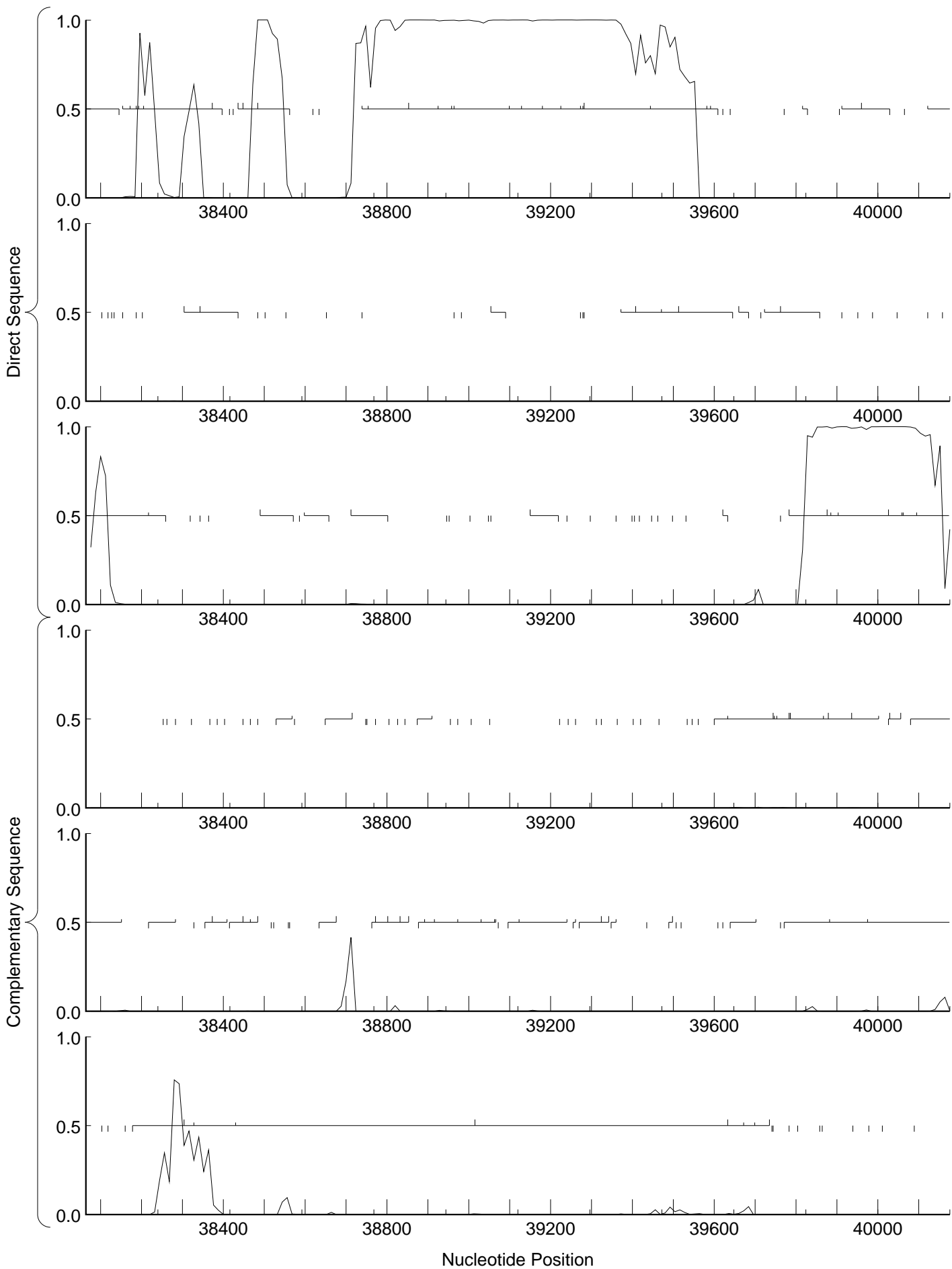


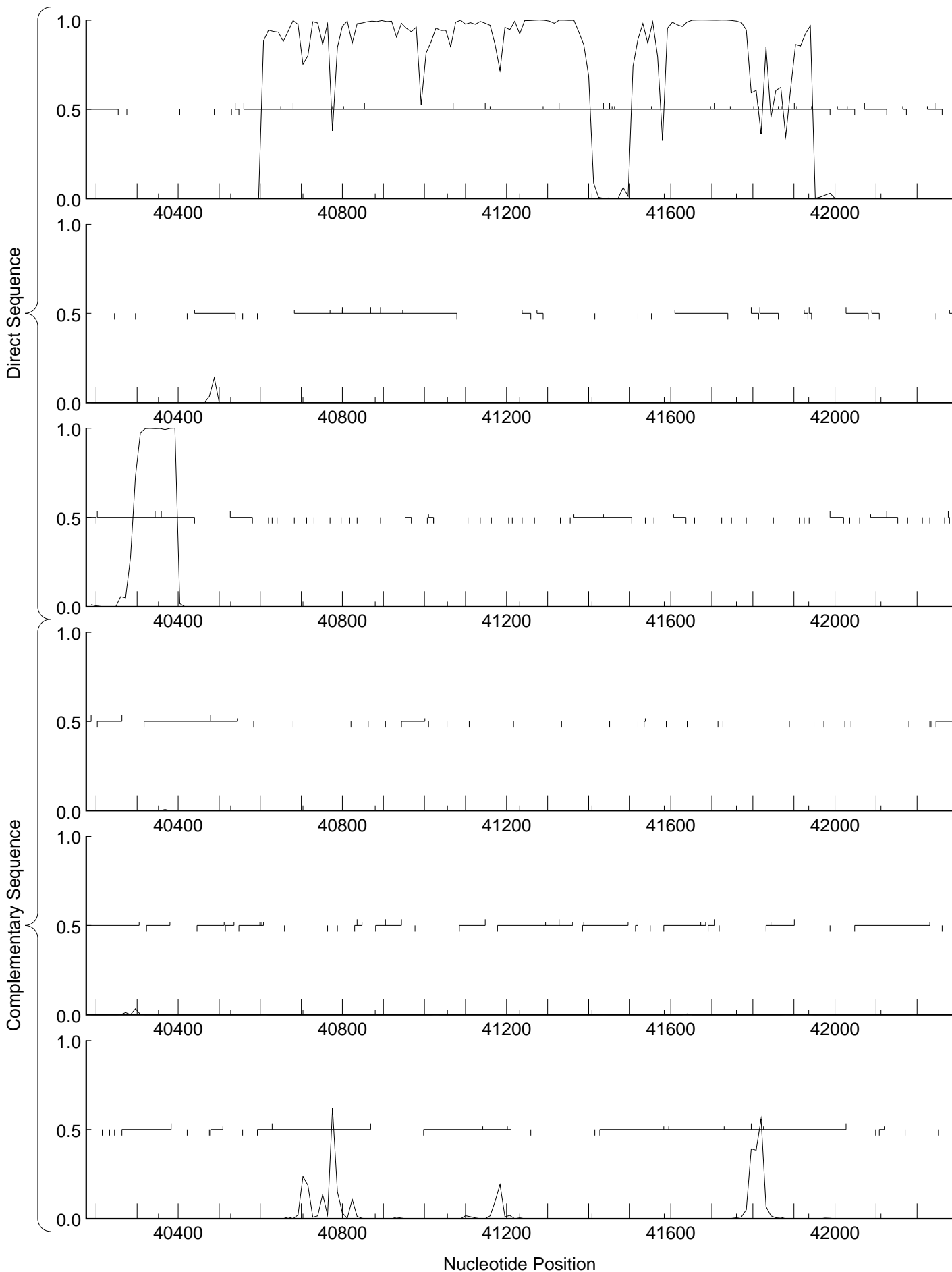


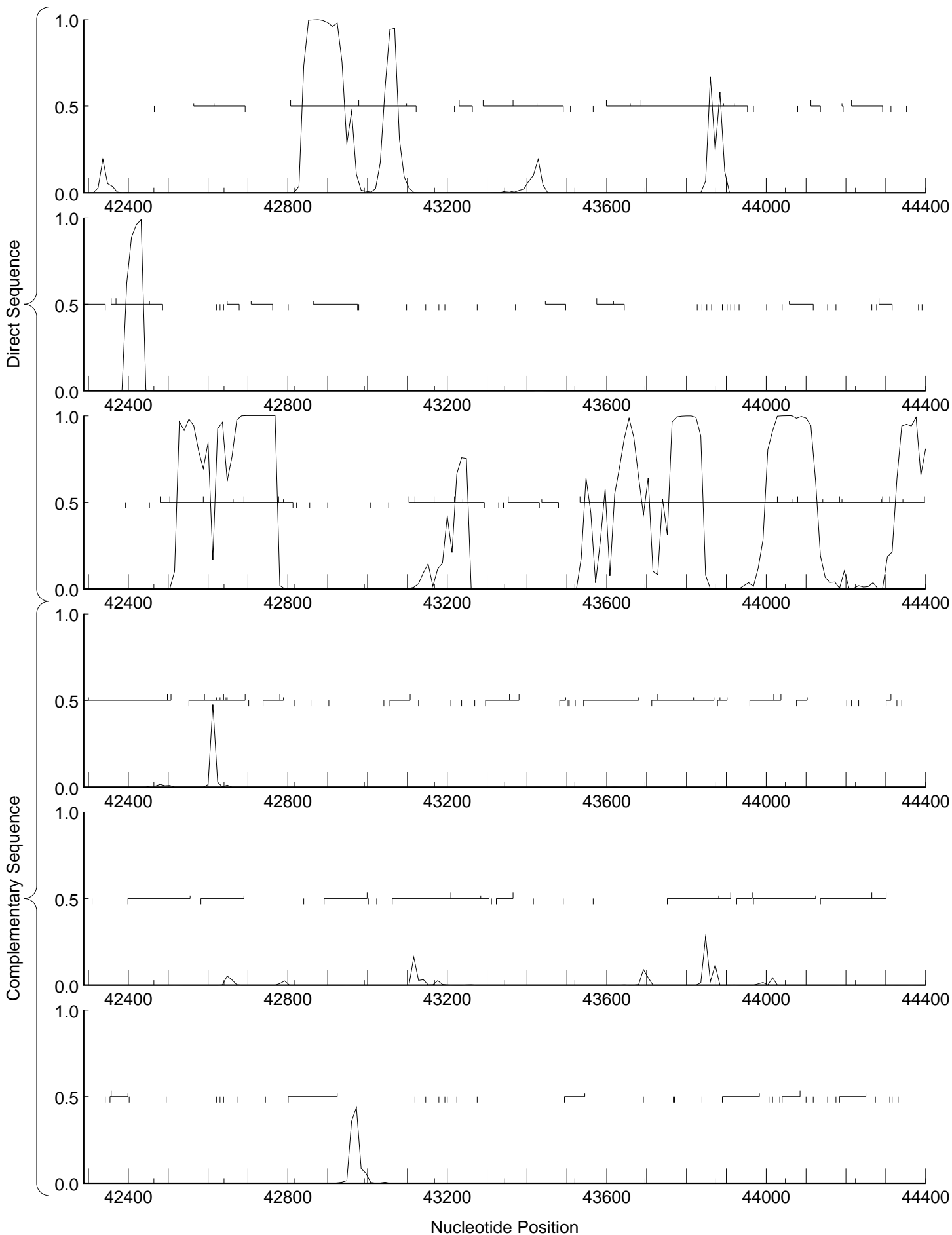


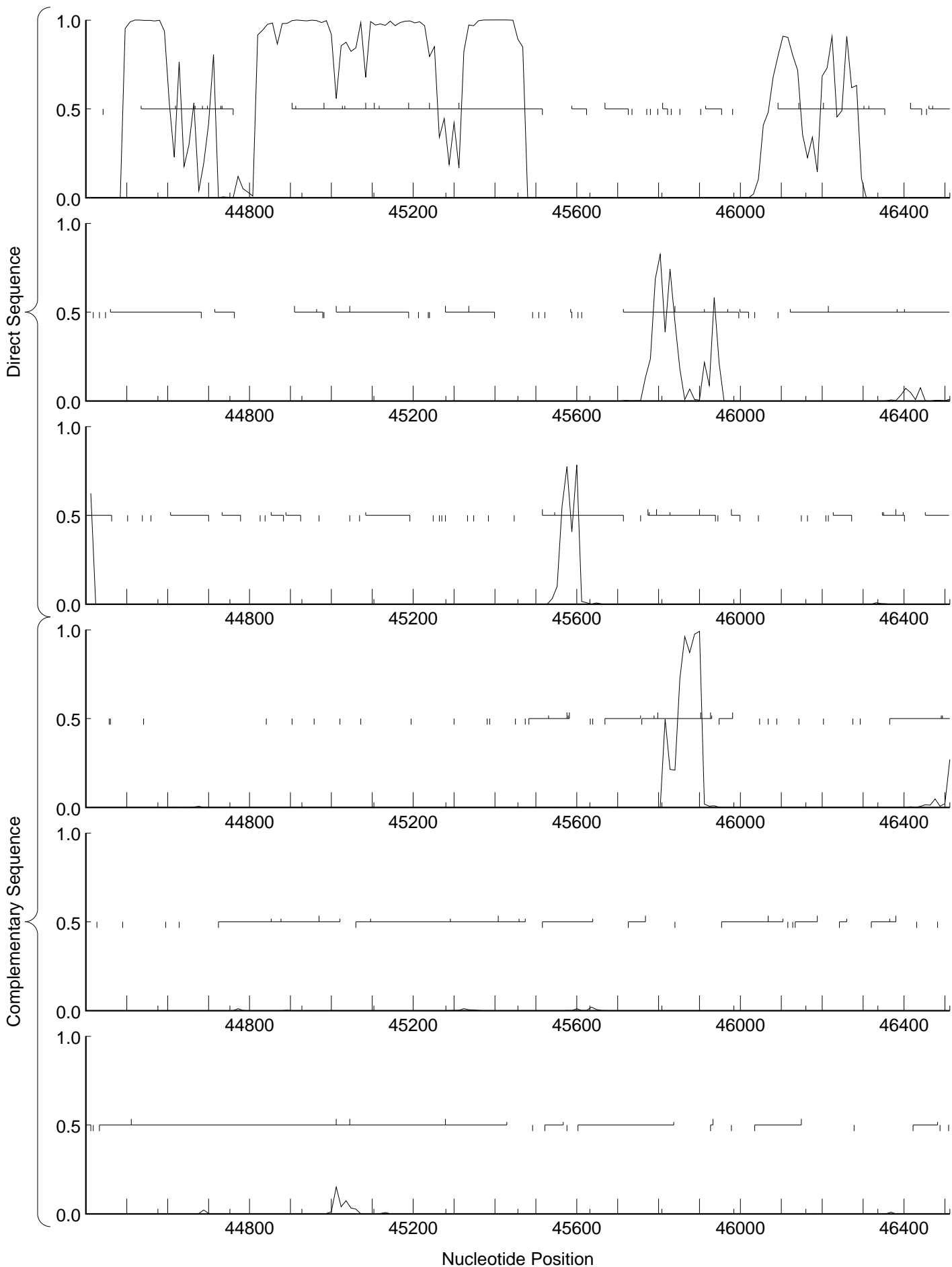


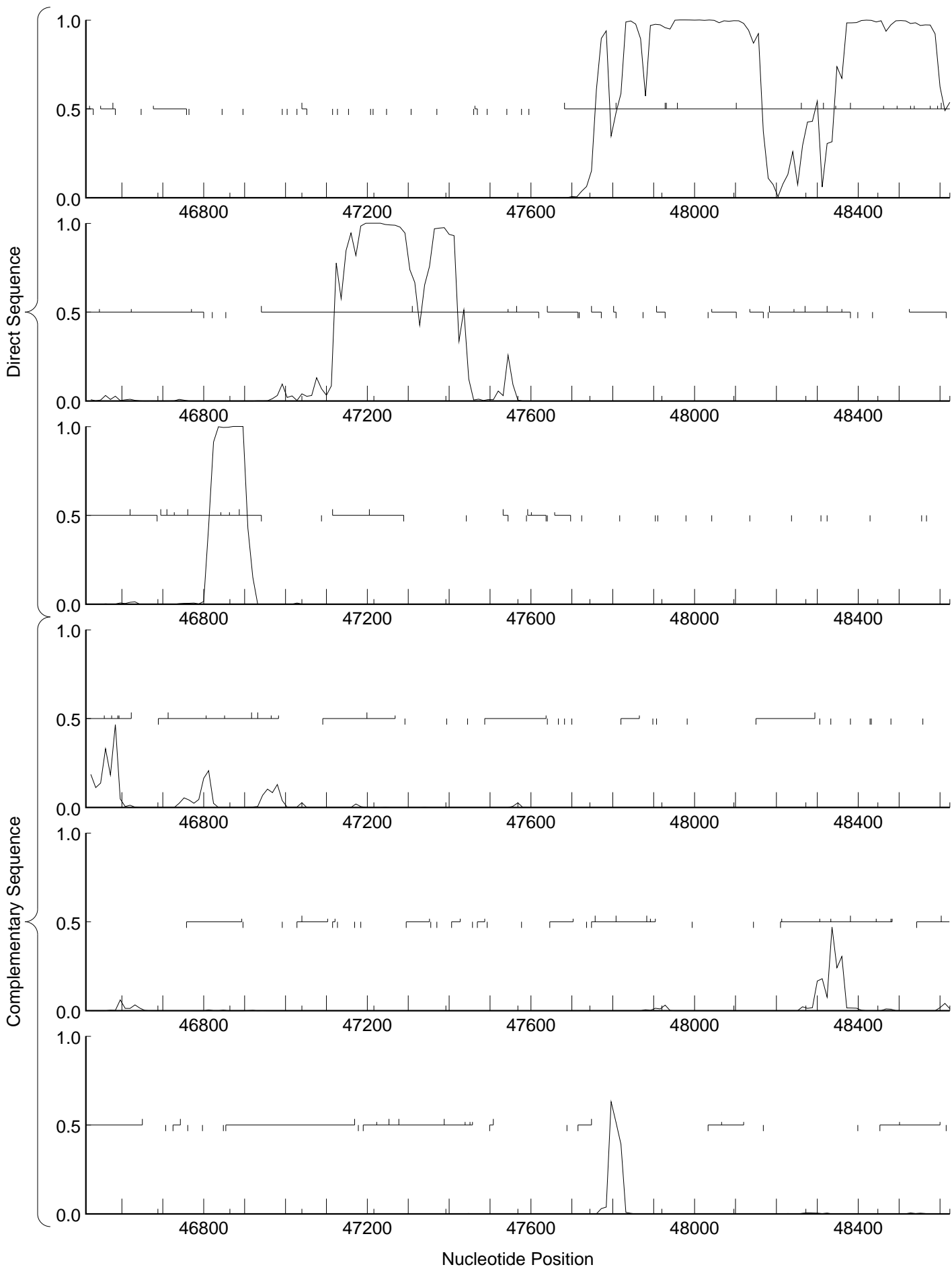


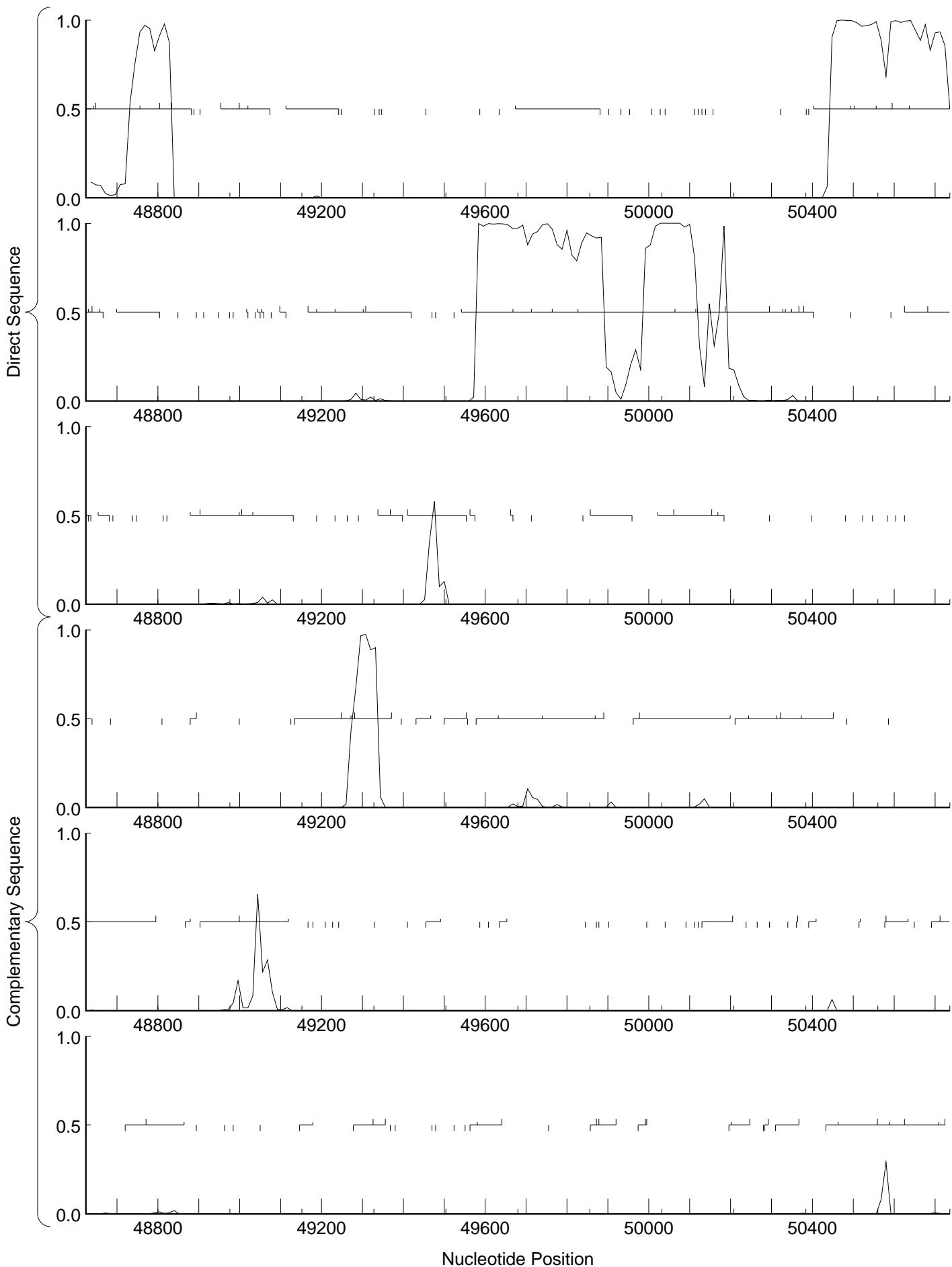


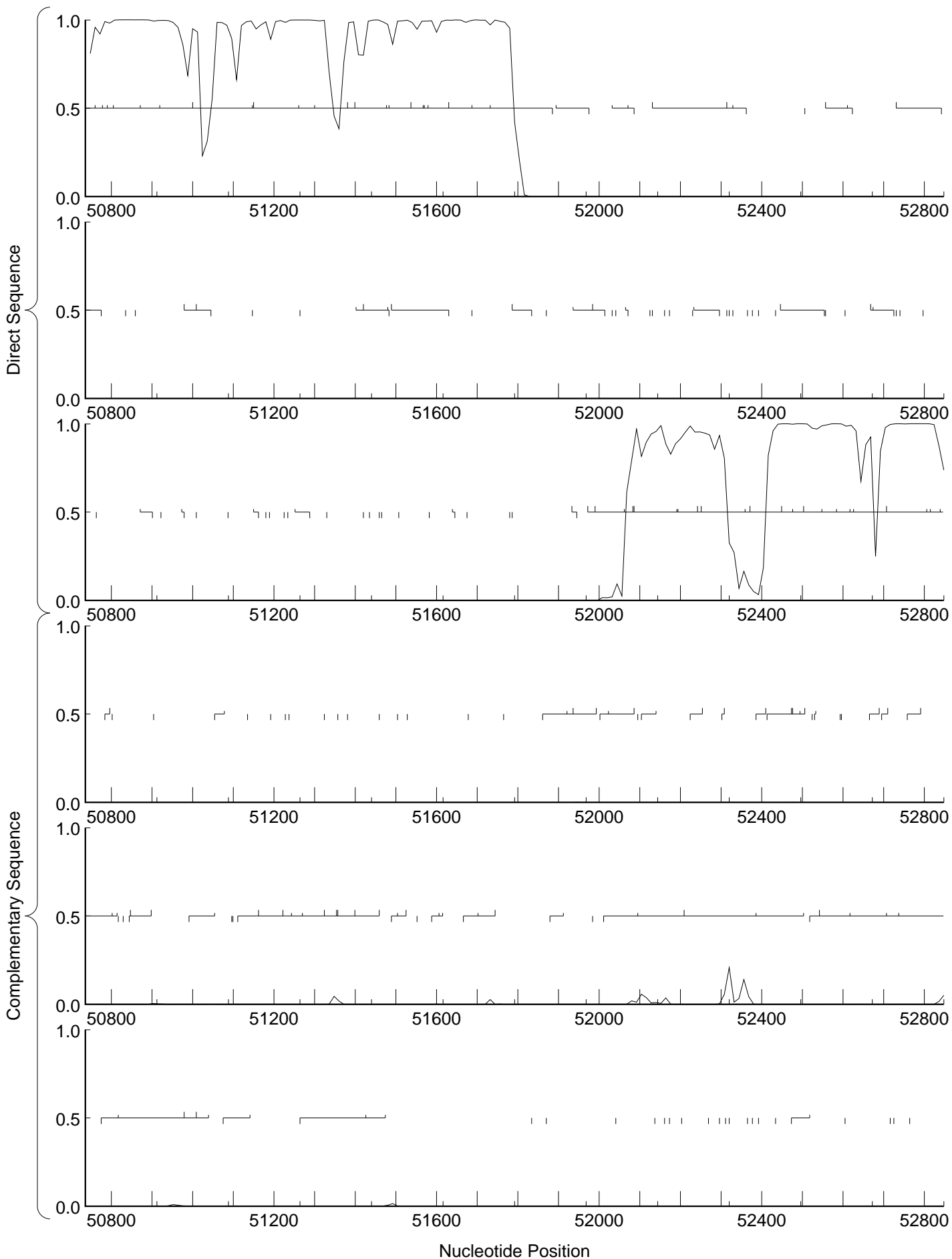


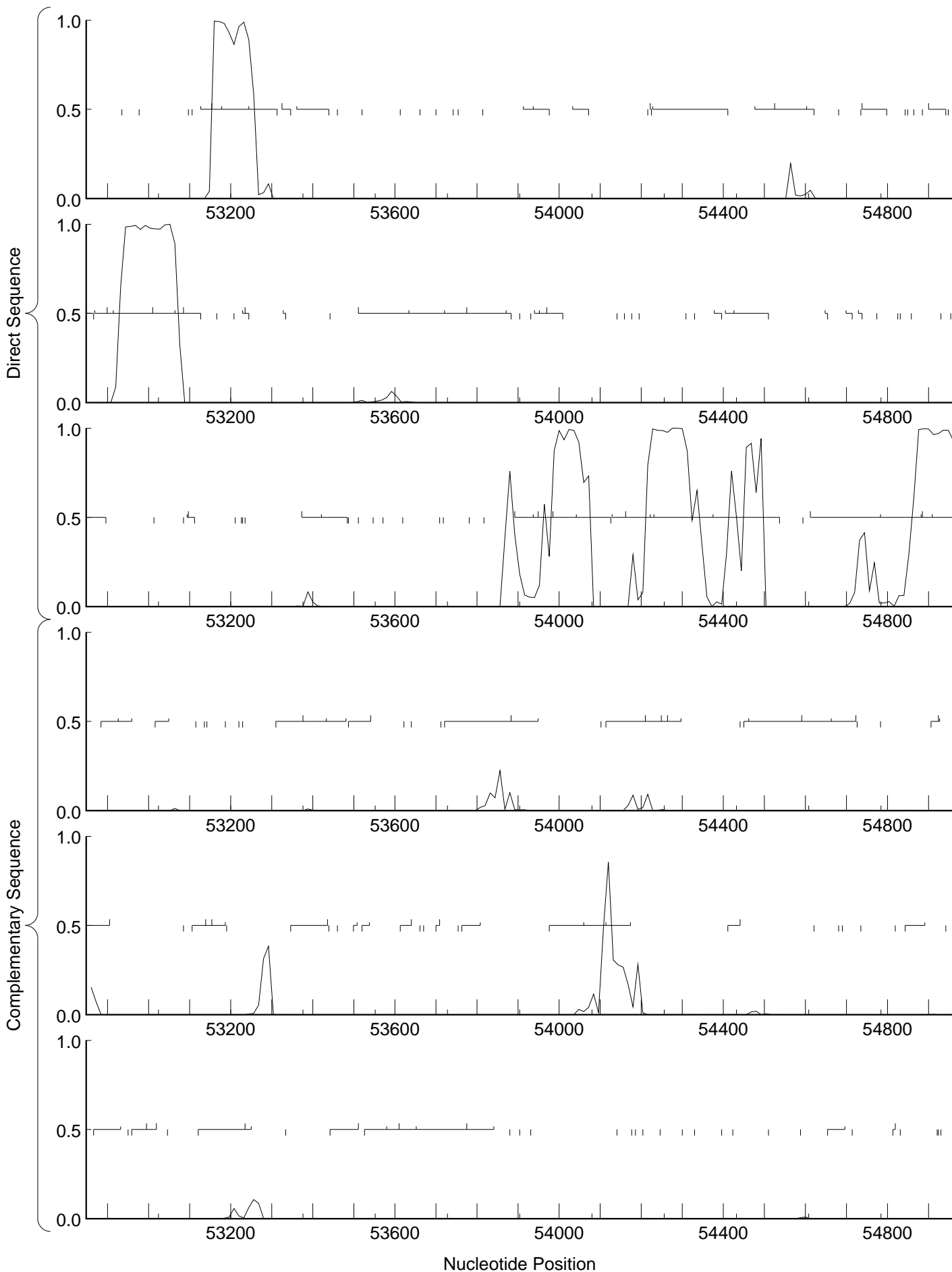


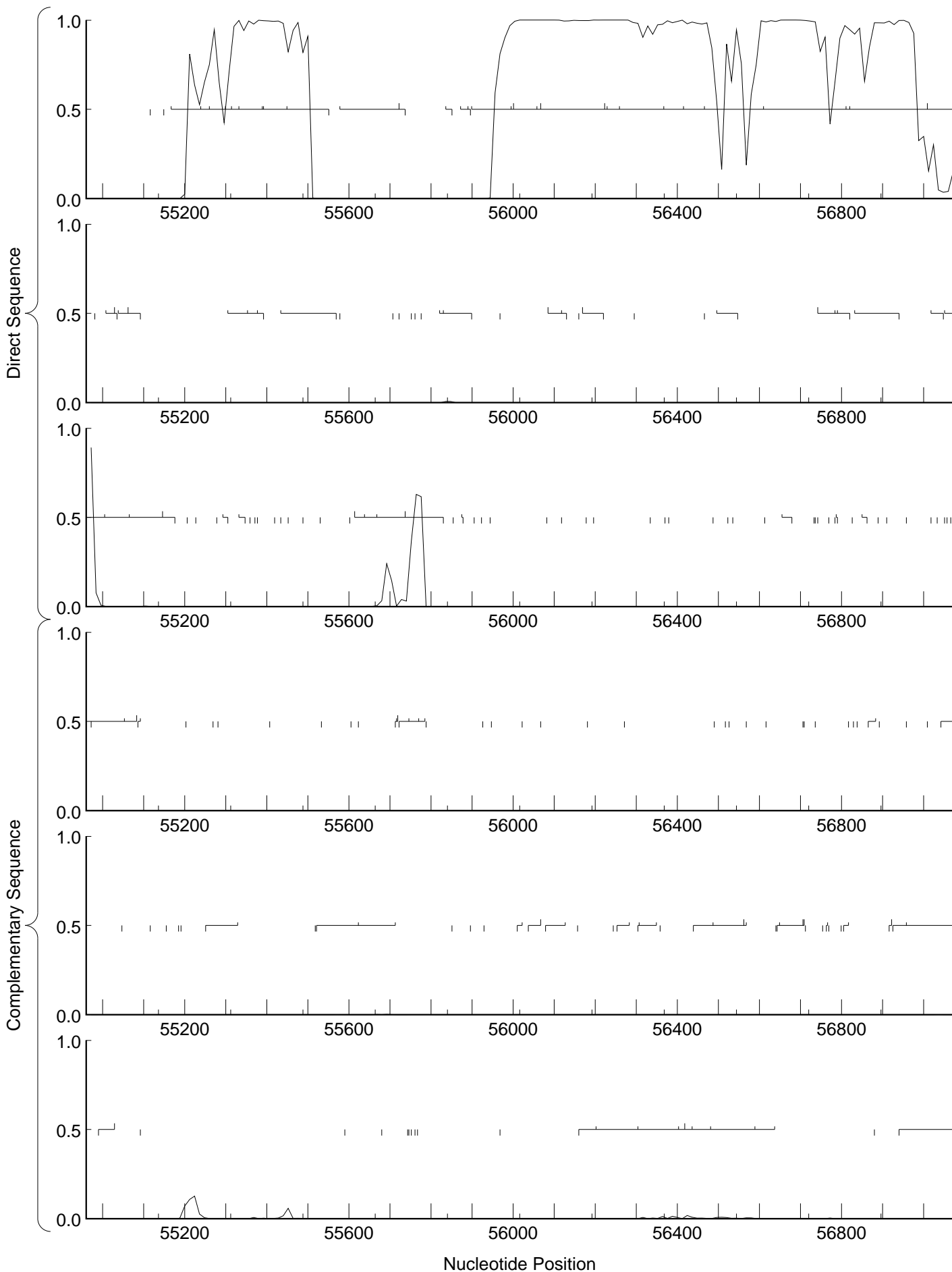


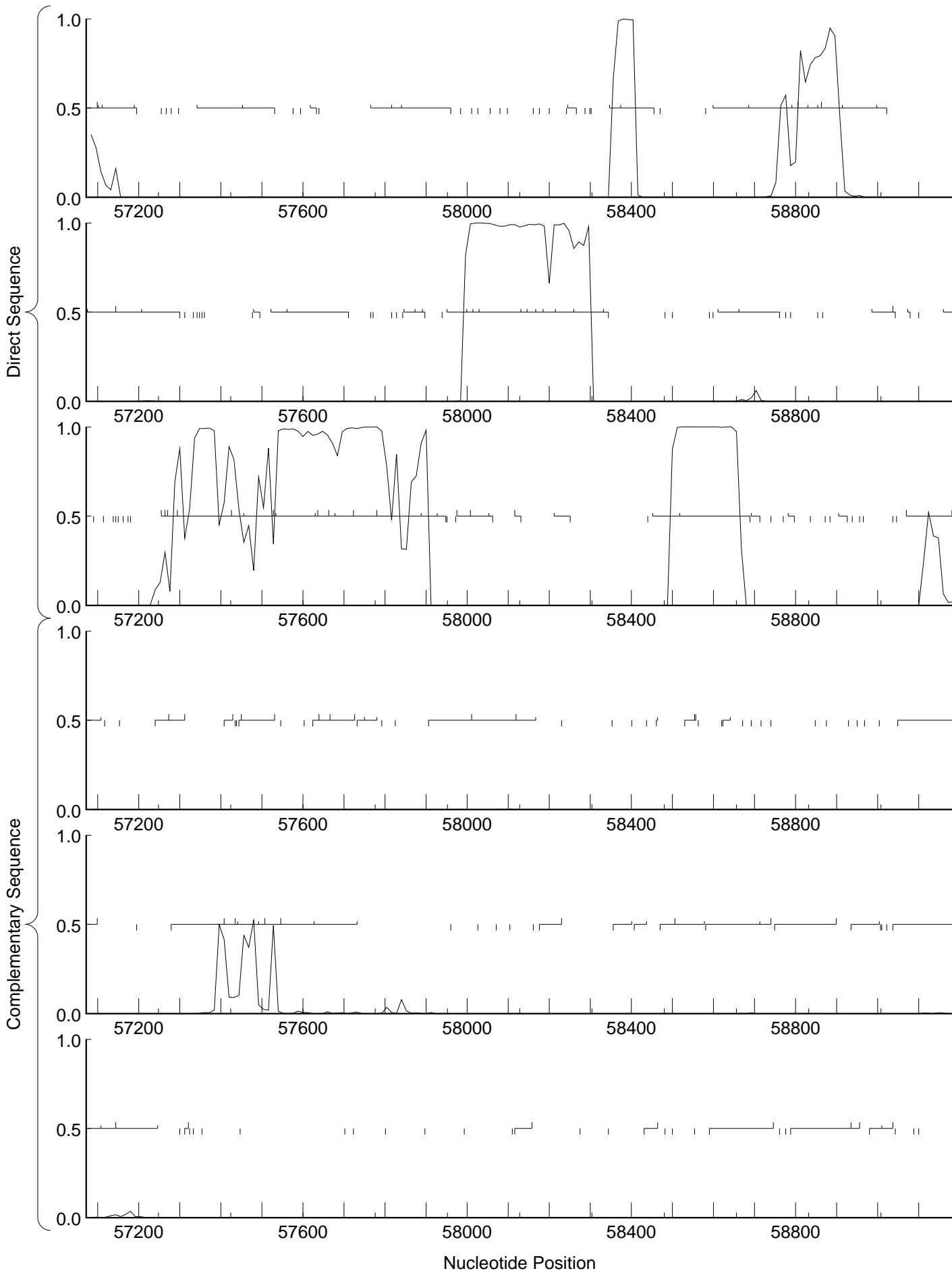












... complete sequence, 76520 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L2, Order 4, Window 96, Step 12, 30/38

