

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Rose5 complete sequence, 73949 bp including 10-base 3' overhang (TCGAAC
Analysis Date : 9/19/21 at 16:32:56
Pages : 37
Sequence Length : 73949 bp
GC Content : 58.72%

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,

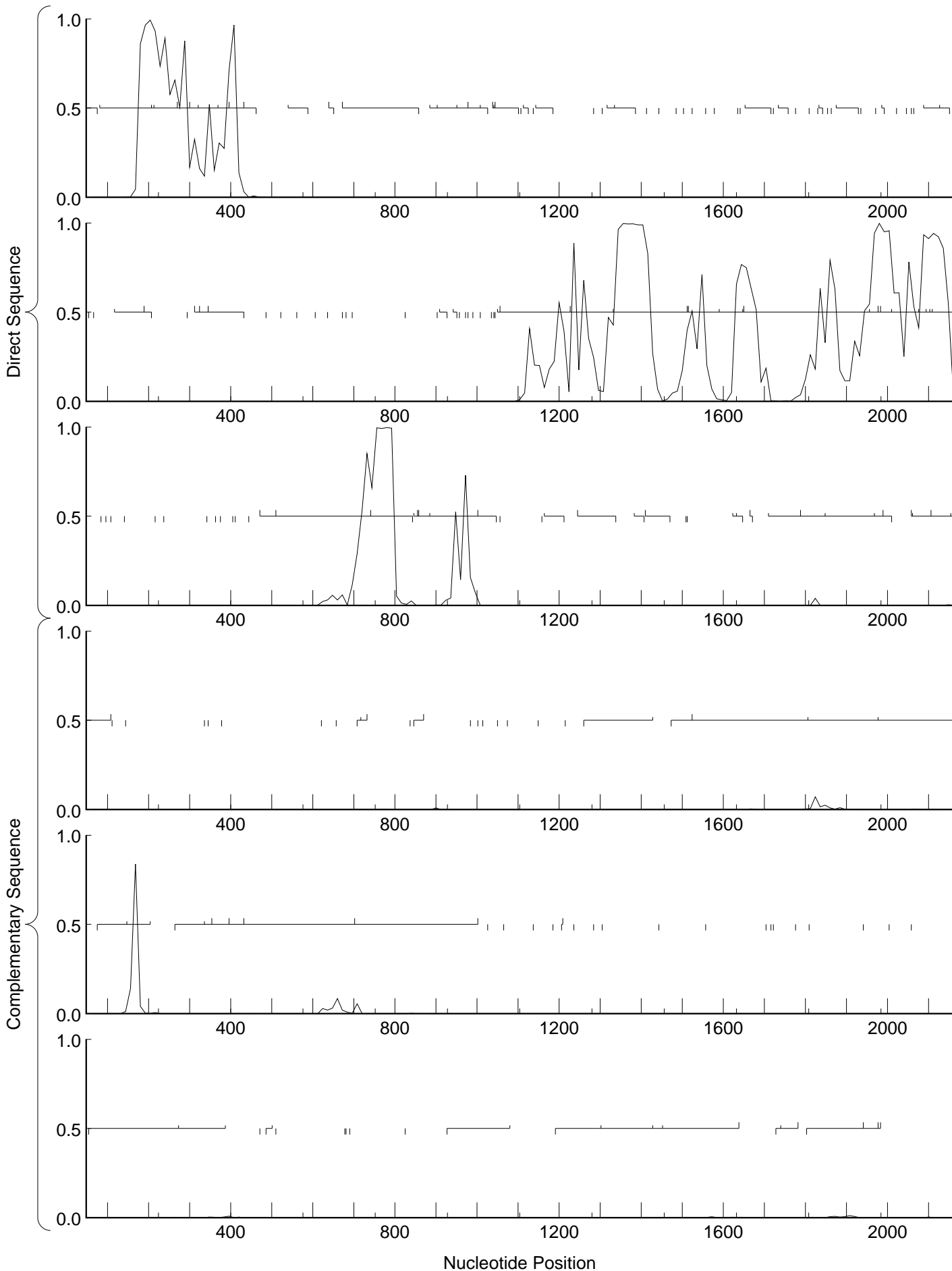
Matrix notes & comments

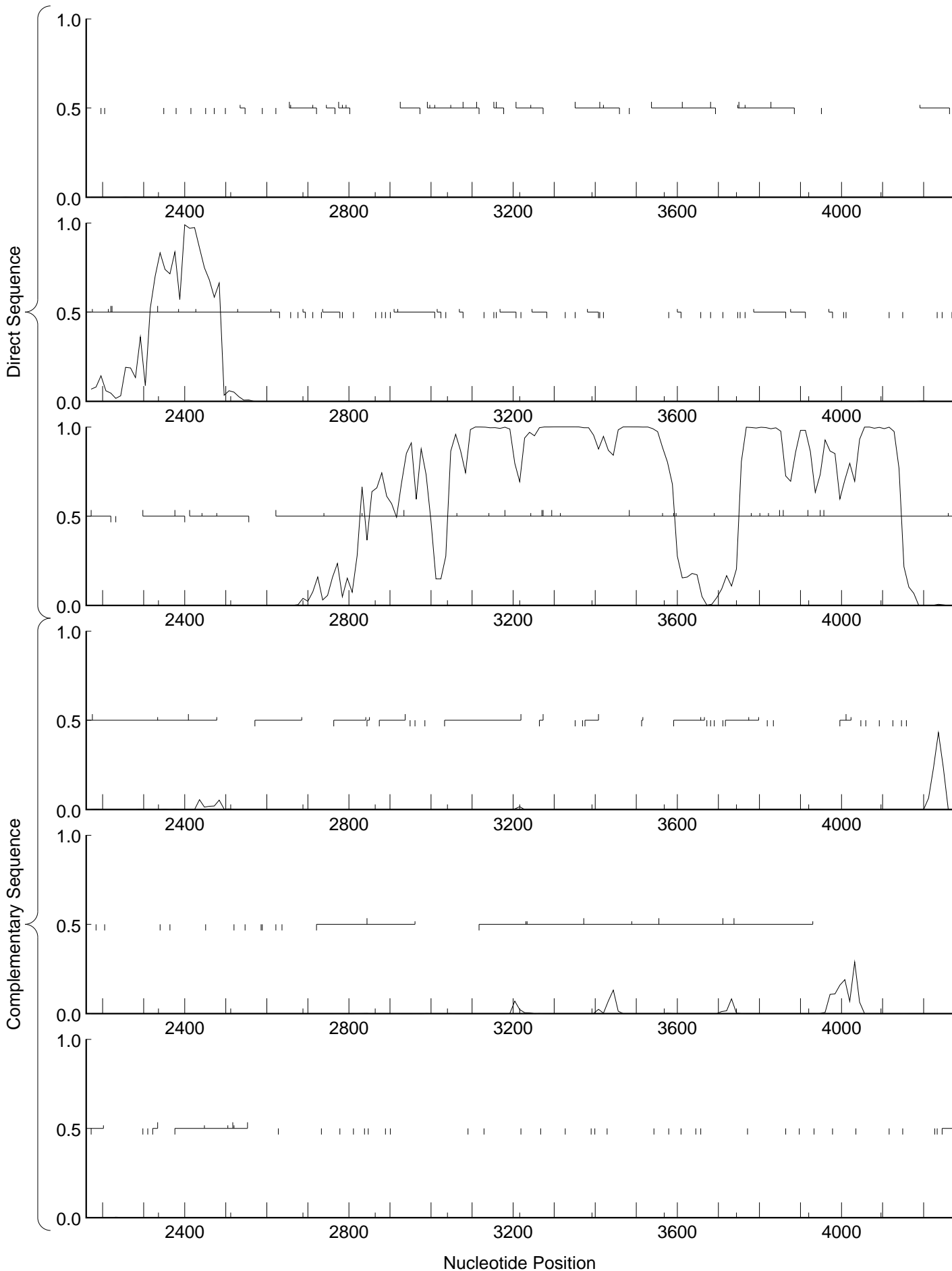
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

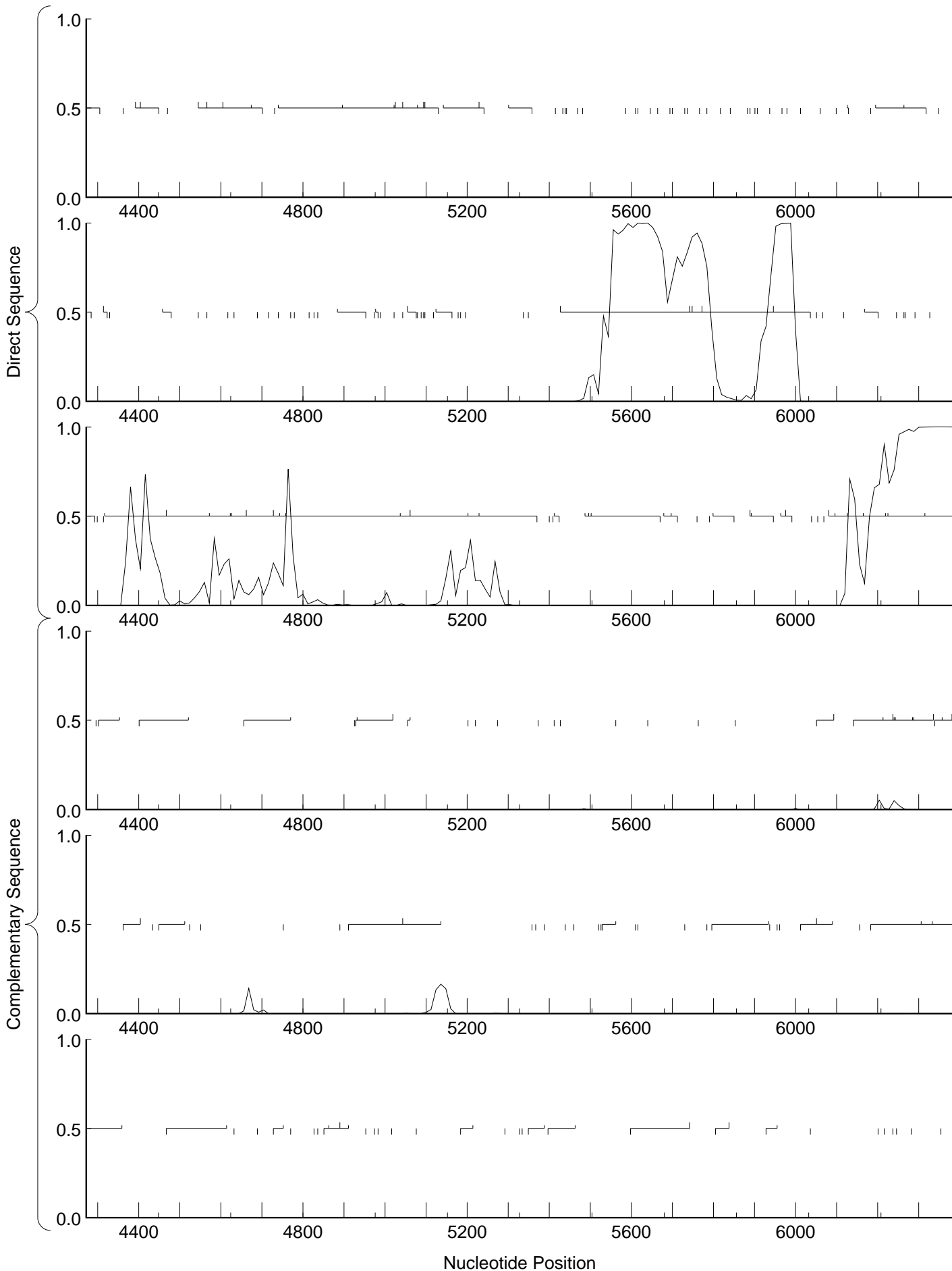
MATRIX INFORMATION

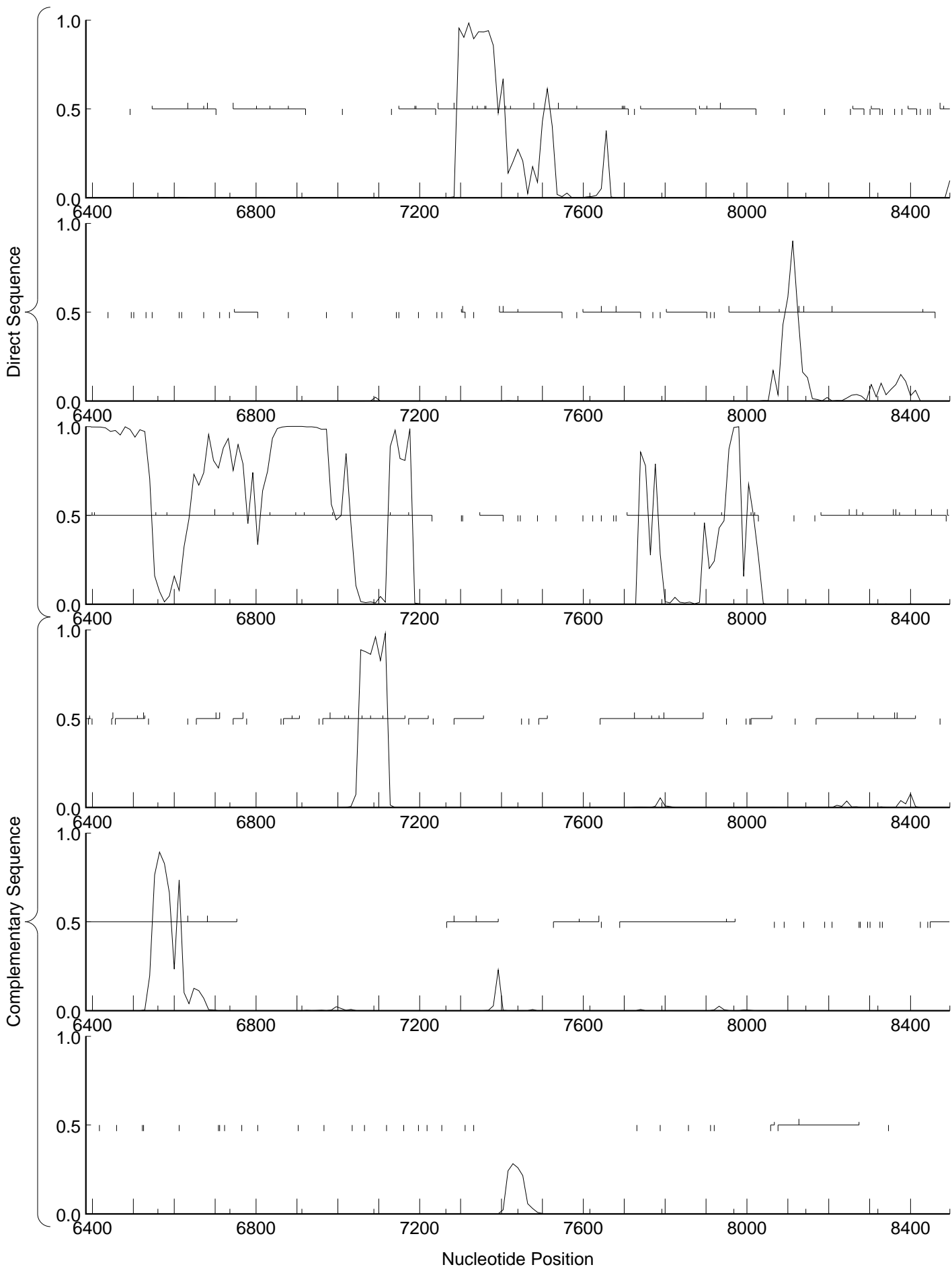
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

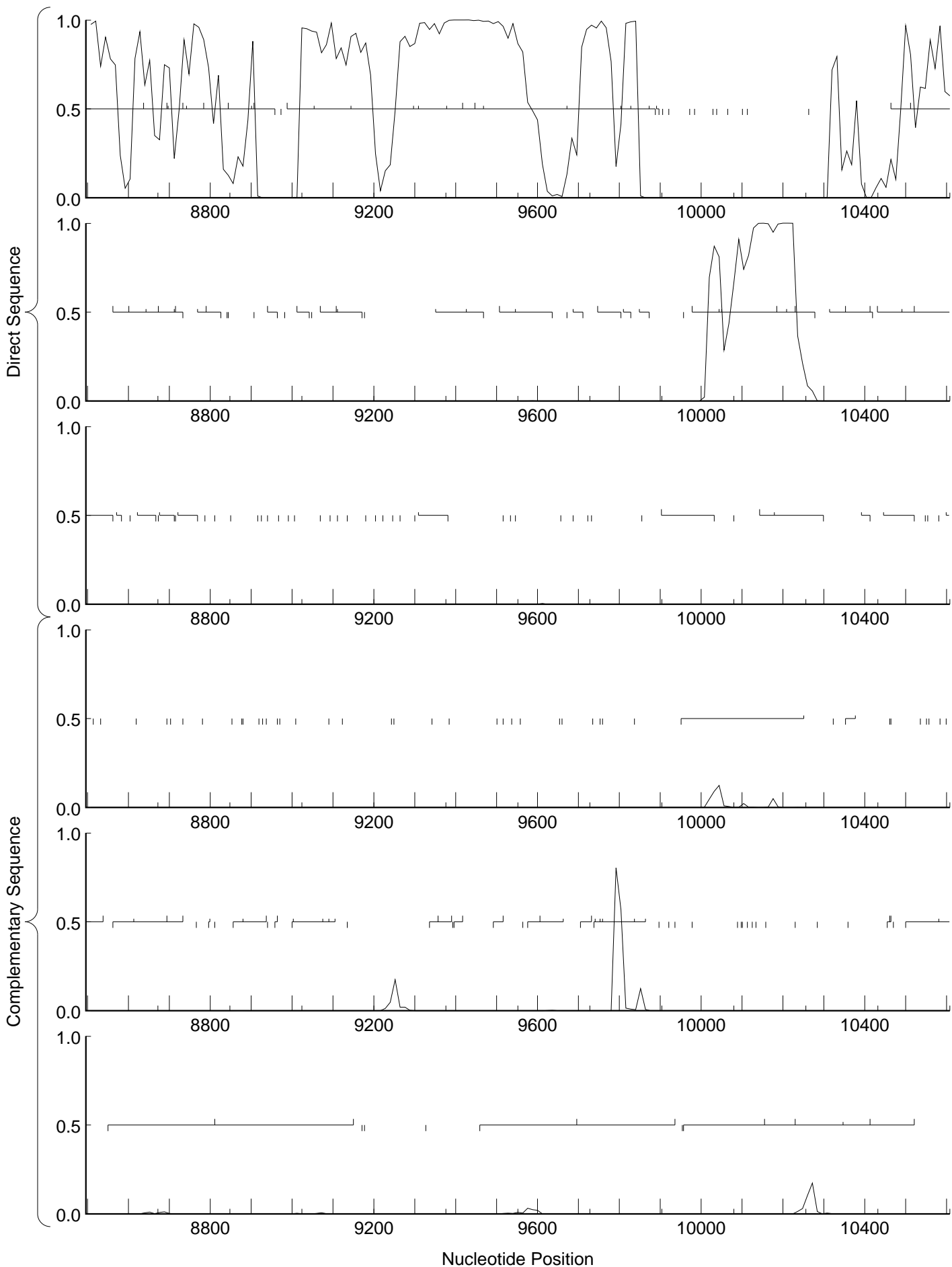
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

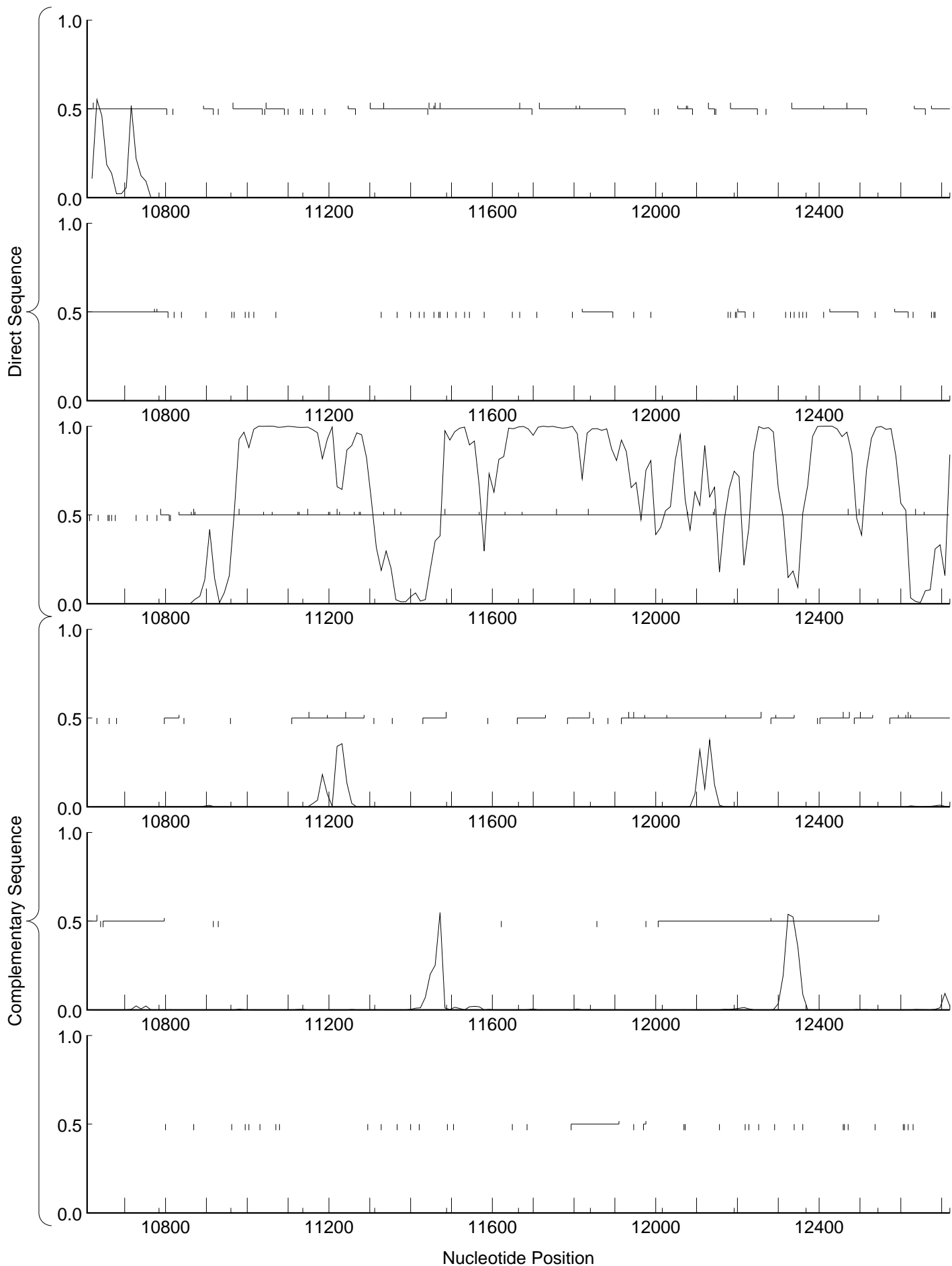


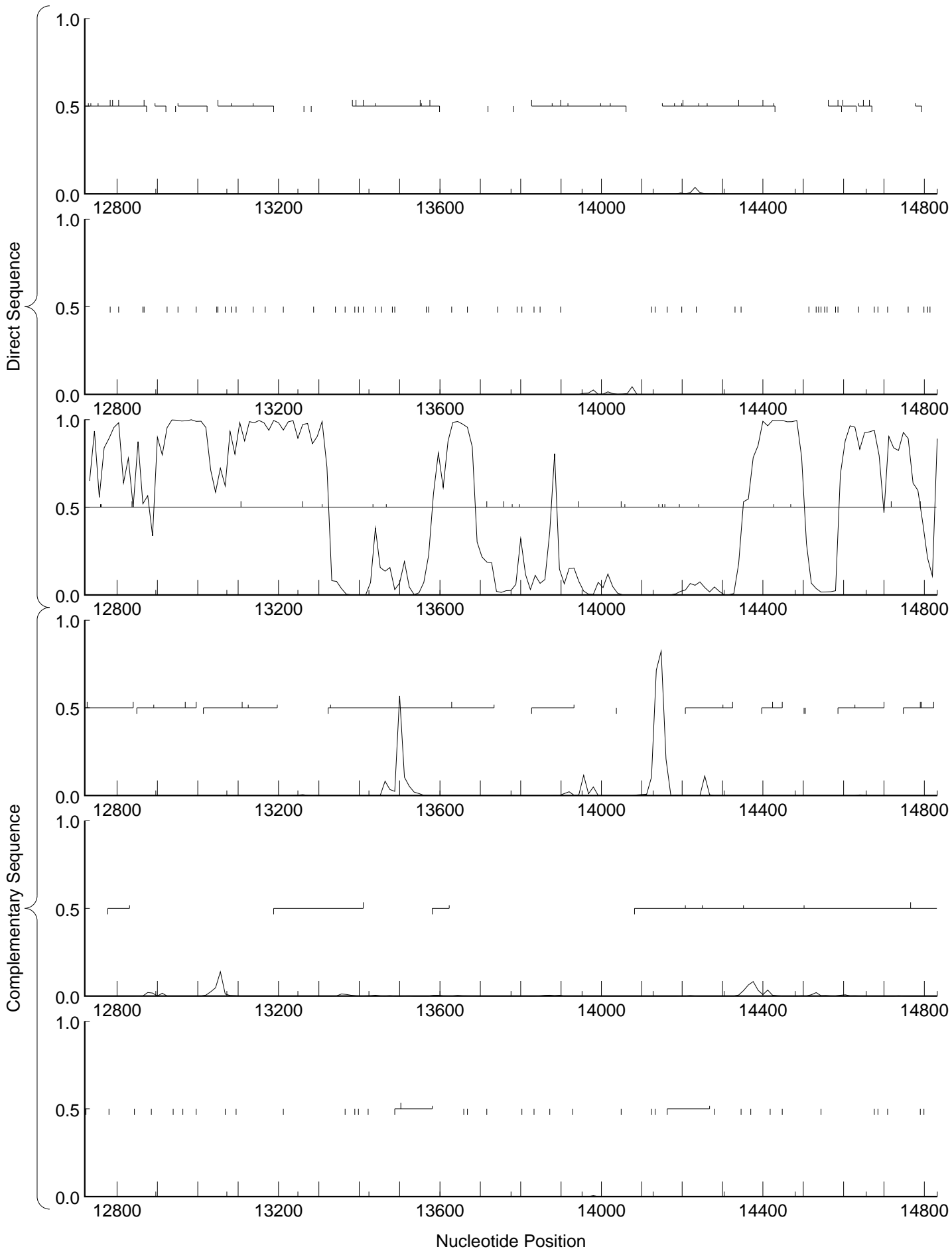


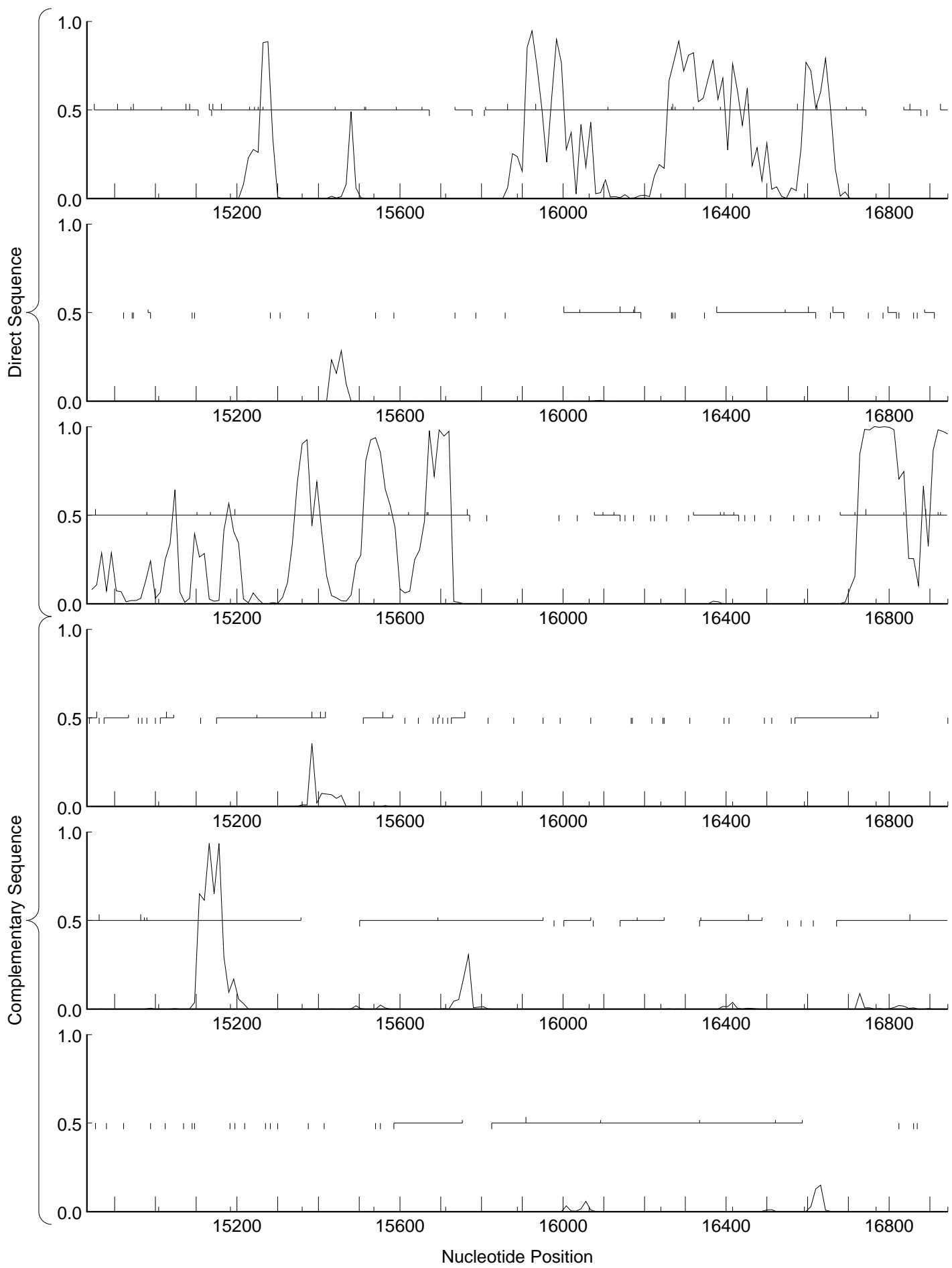


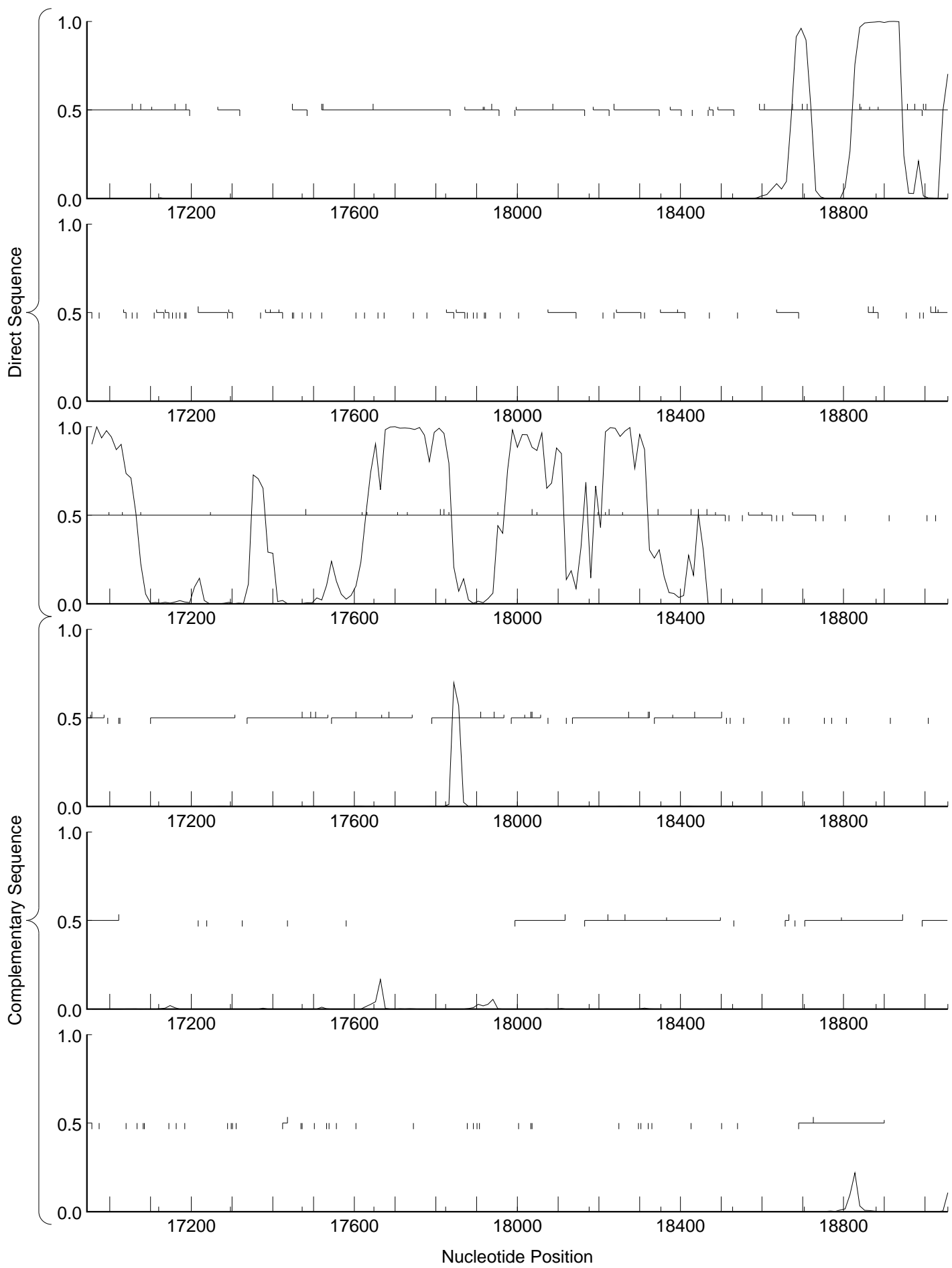




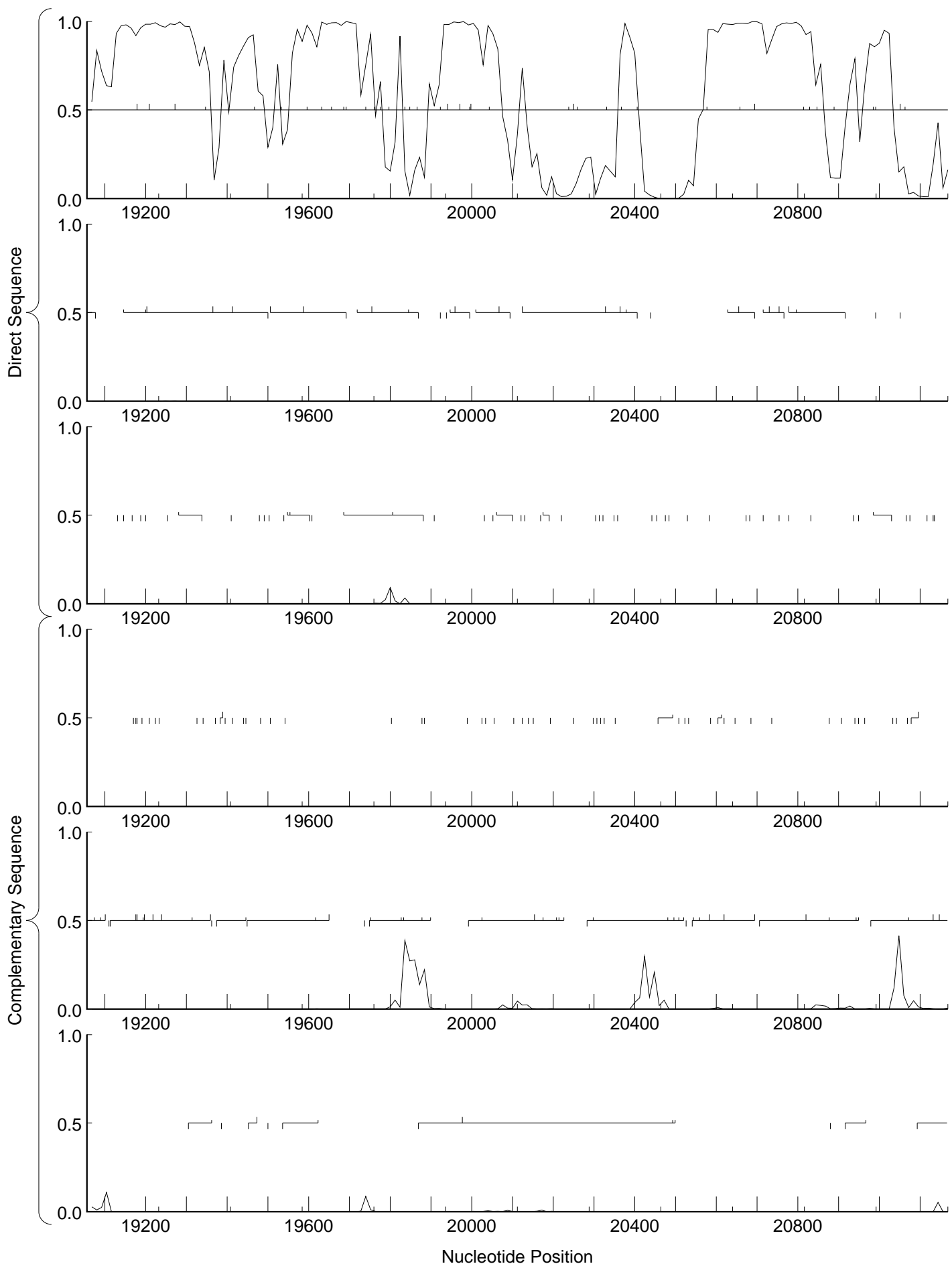


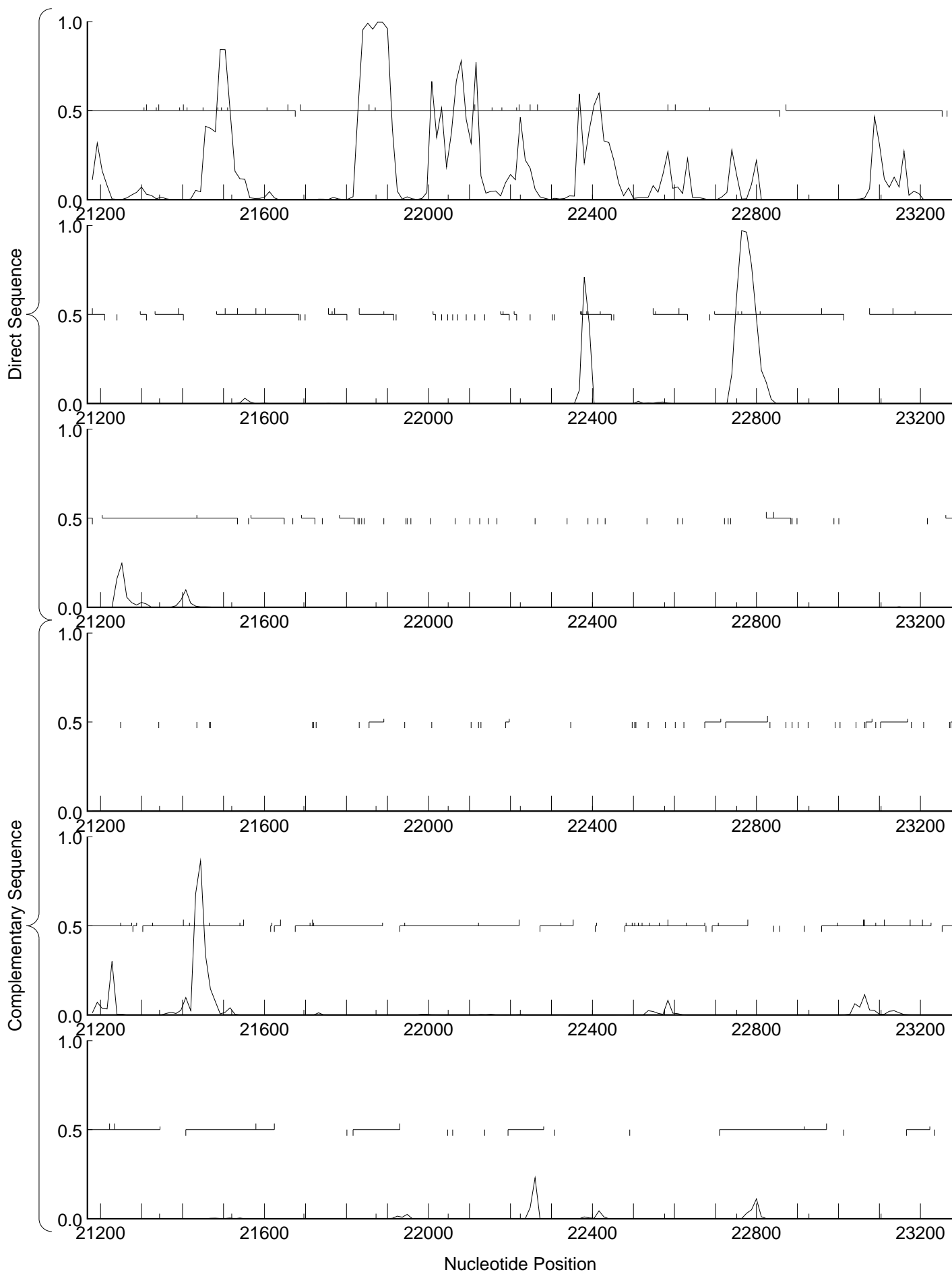




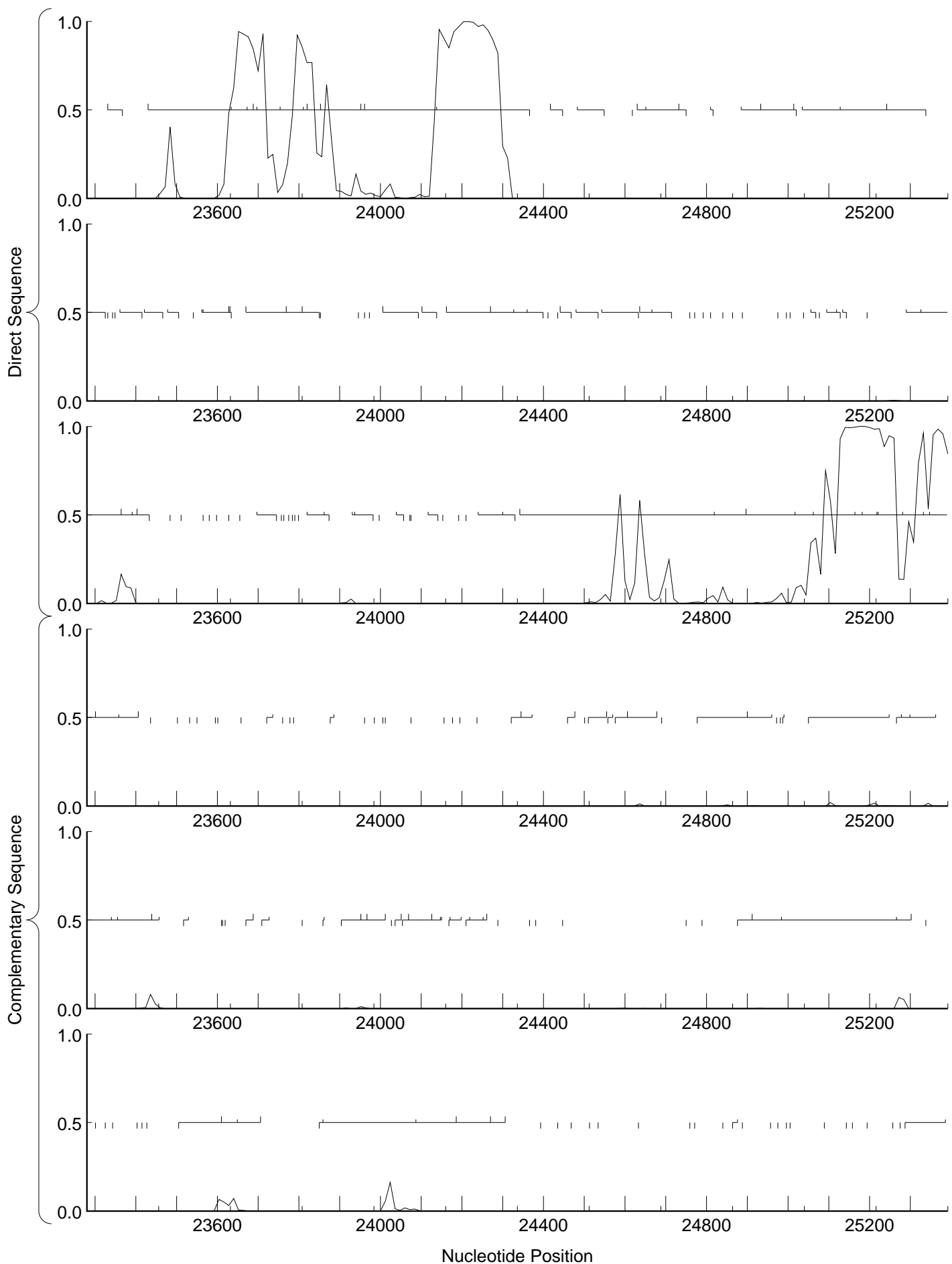


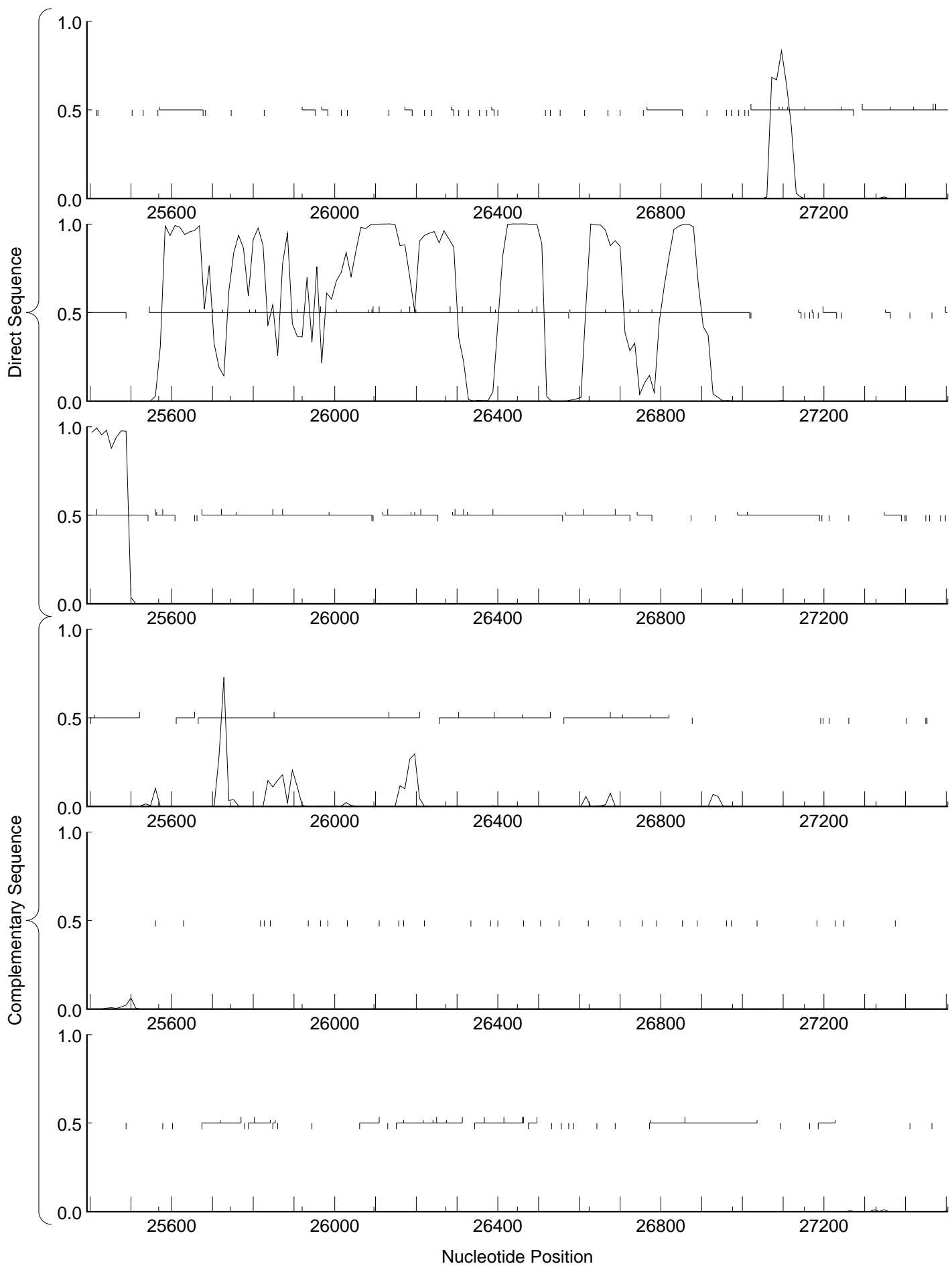
complete sequence, 73949 bp including 10-base 3' overhang (TCGAACGGCC), Cluster L1, Order 4, Window 96, Step 12, 11/37

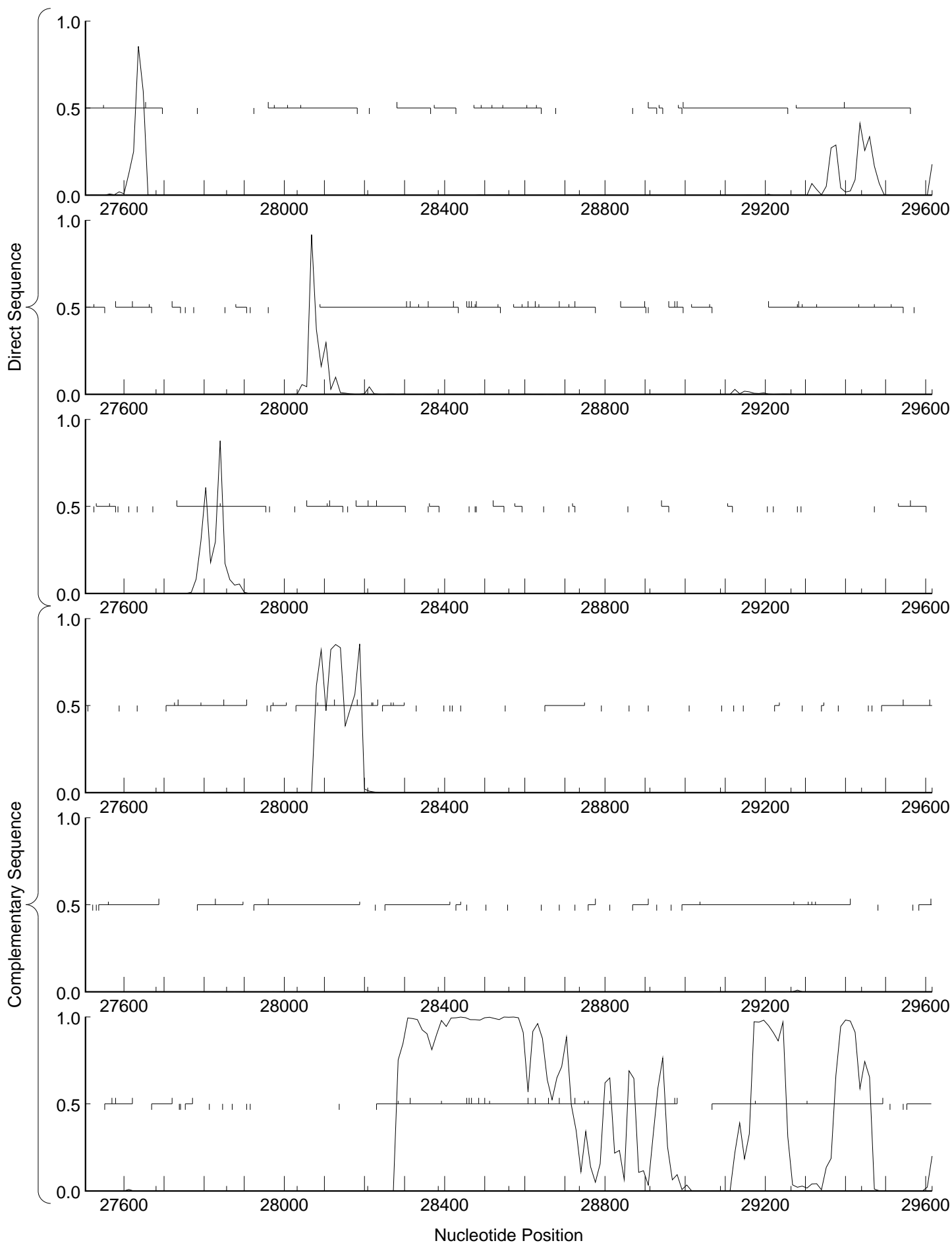


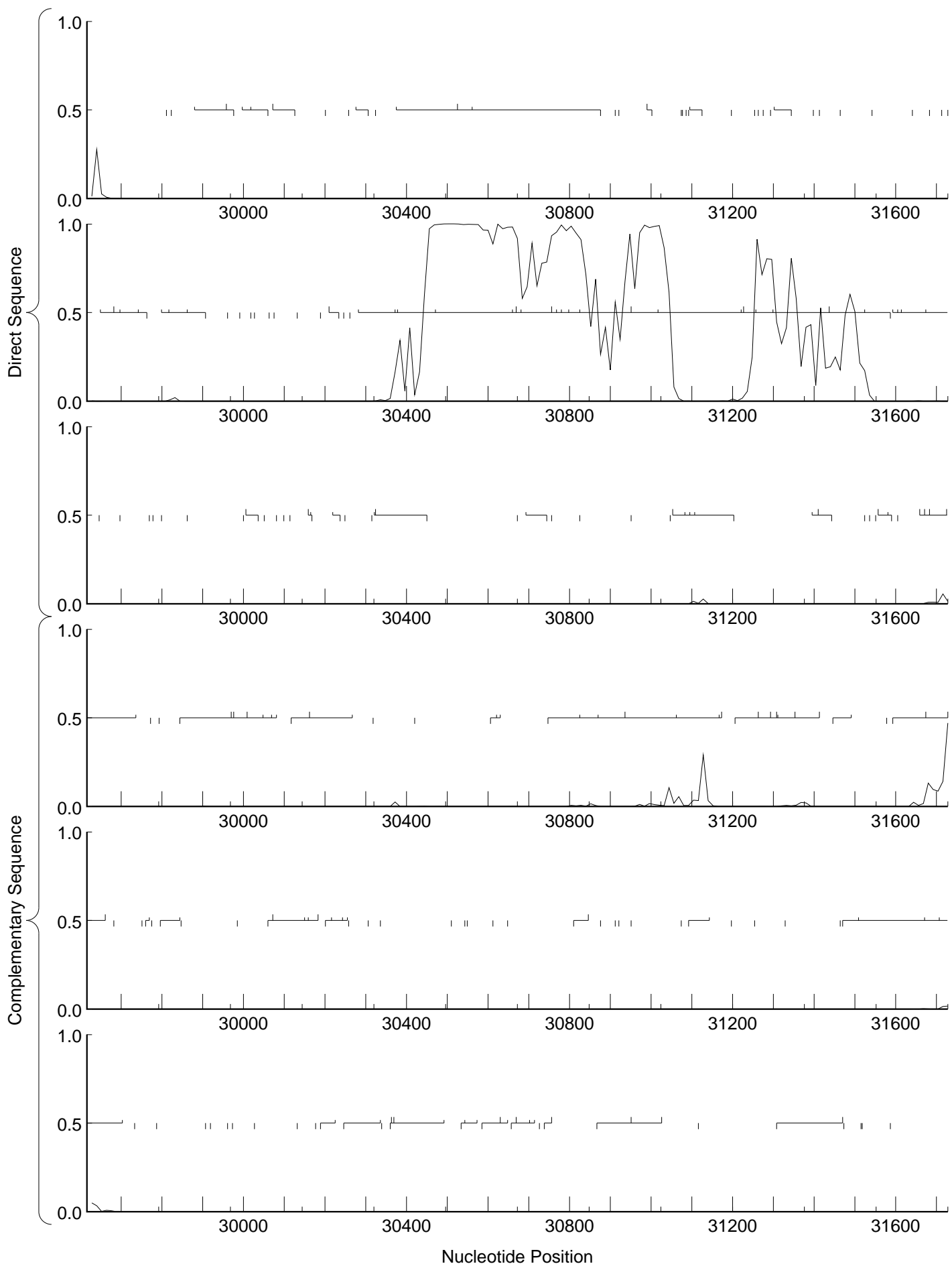


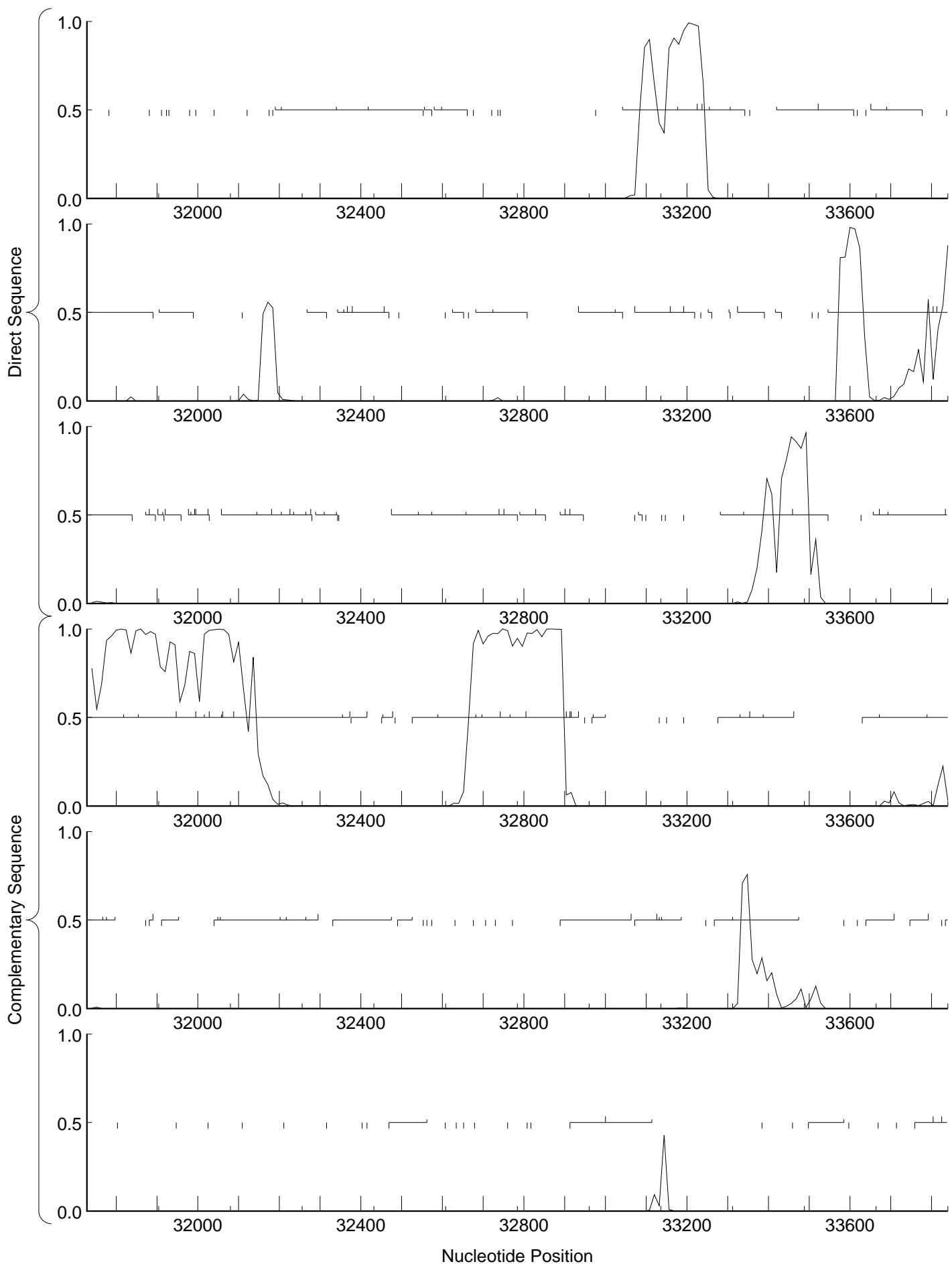
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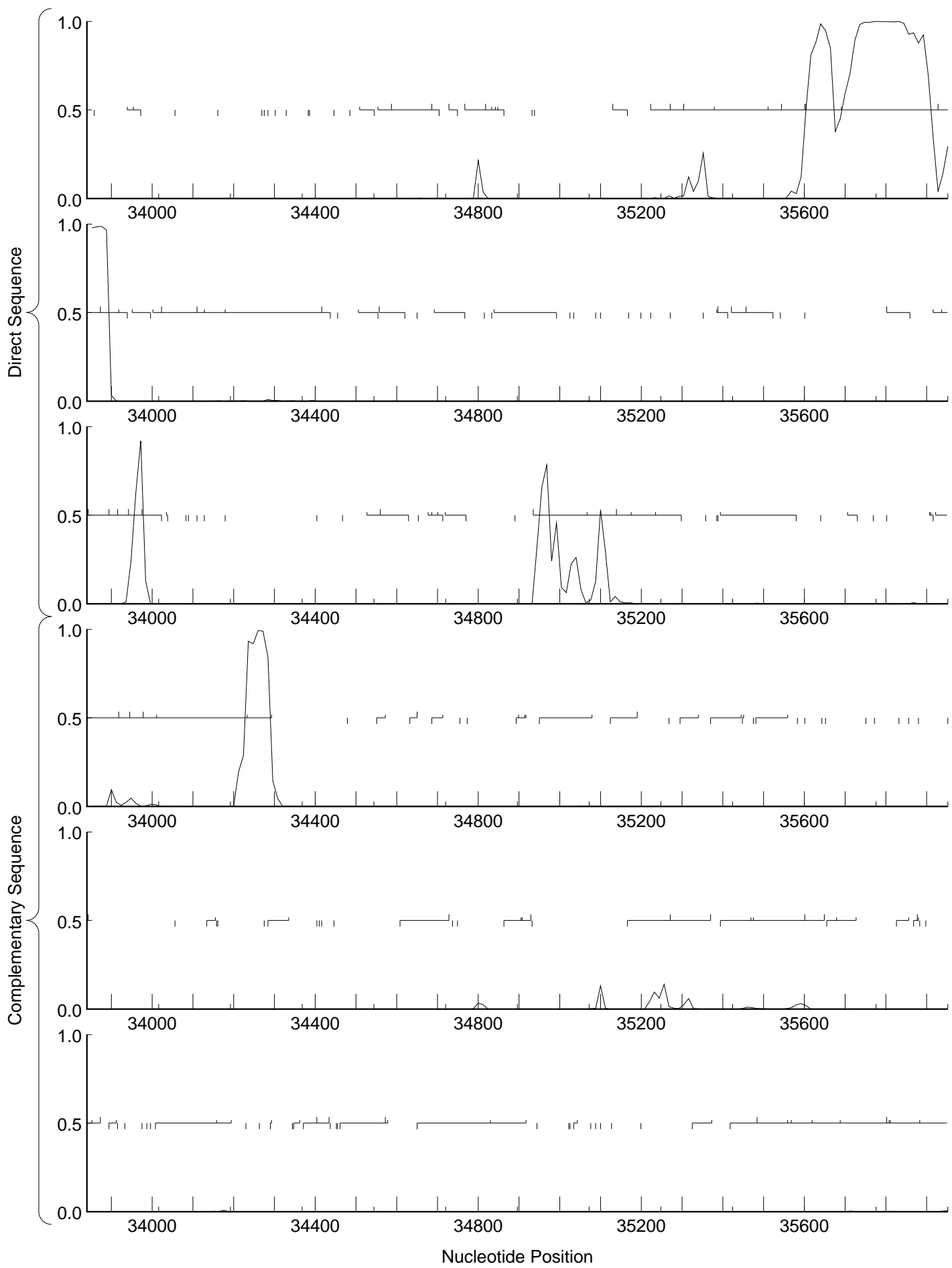




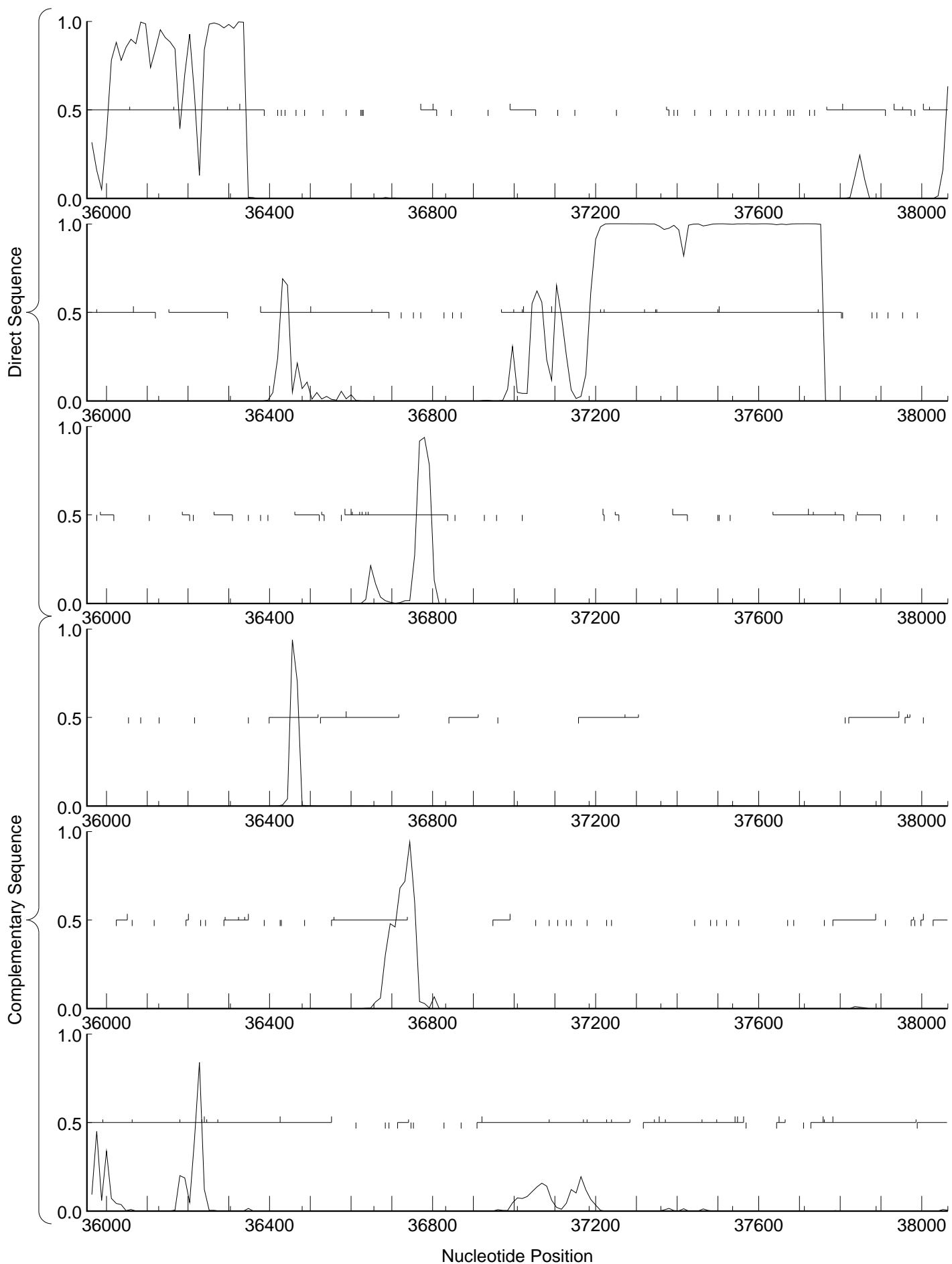


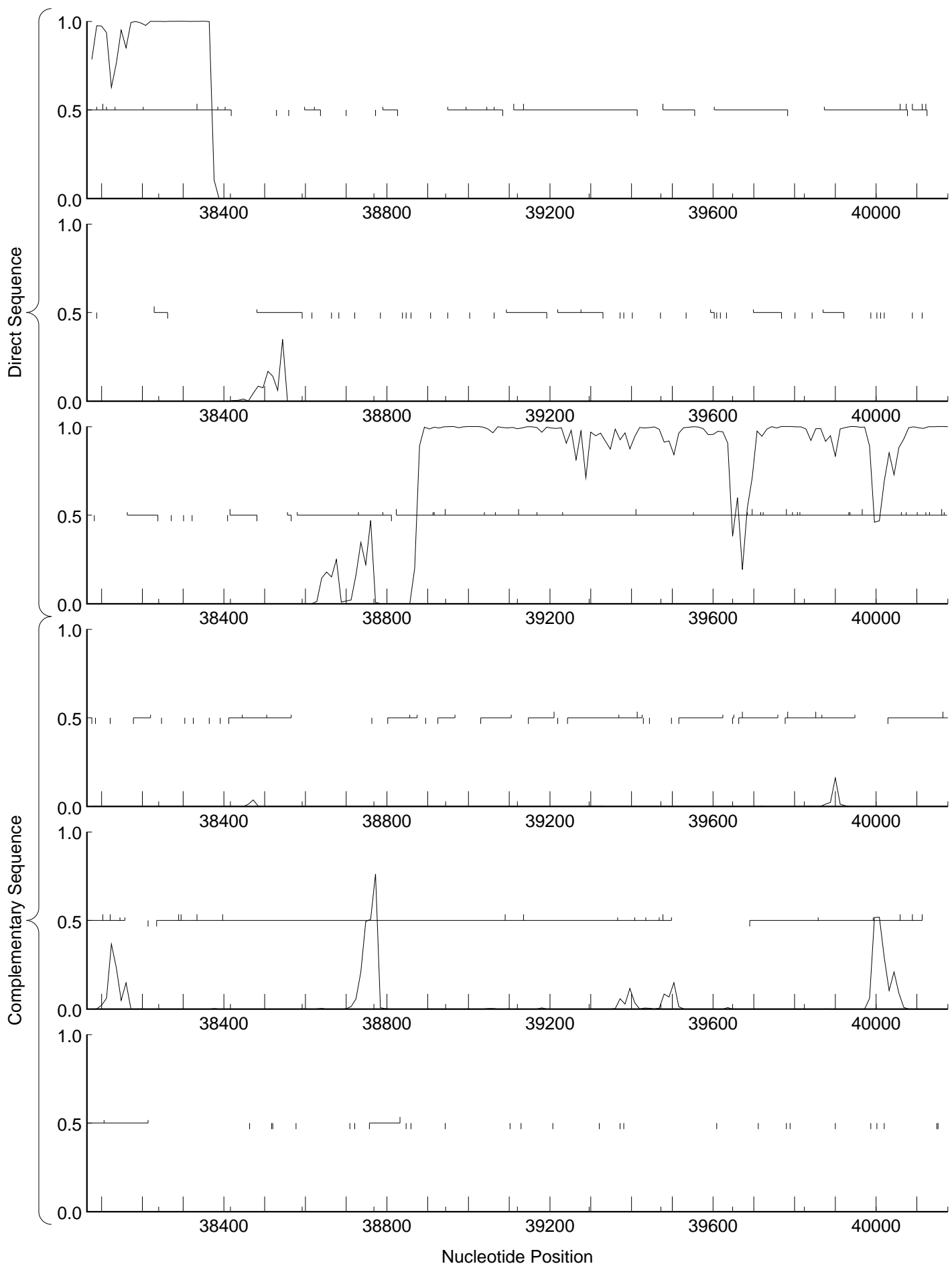




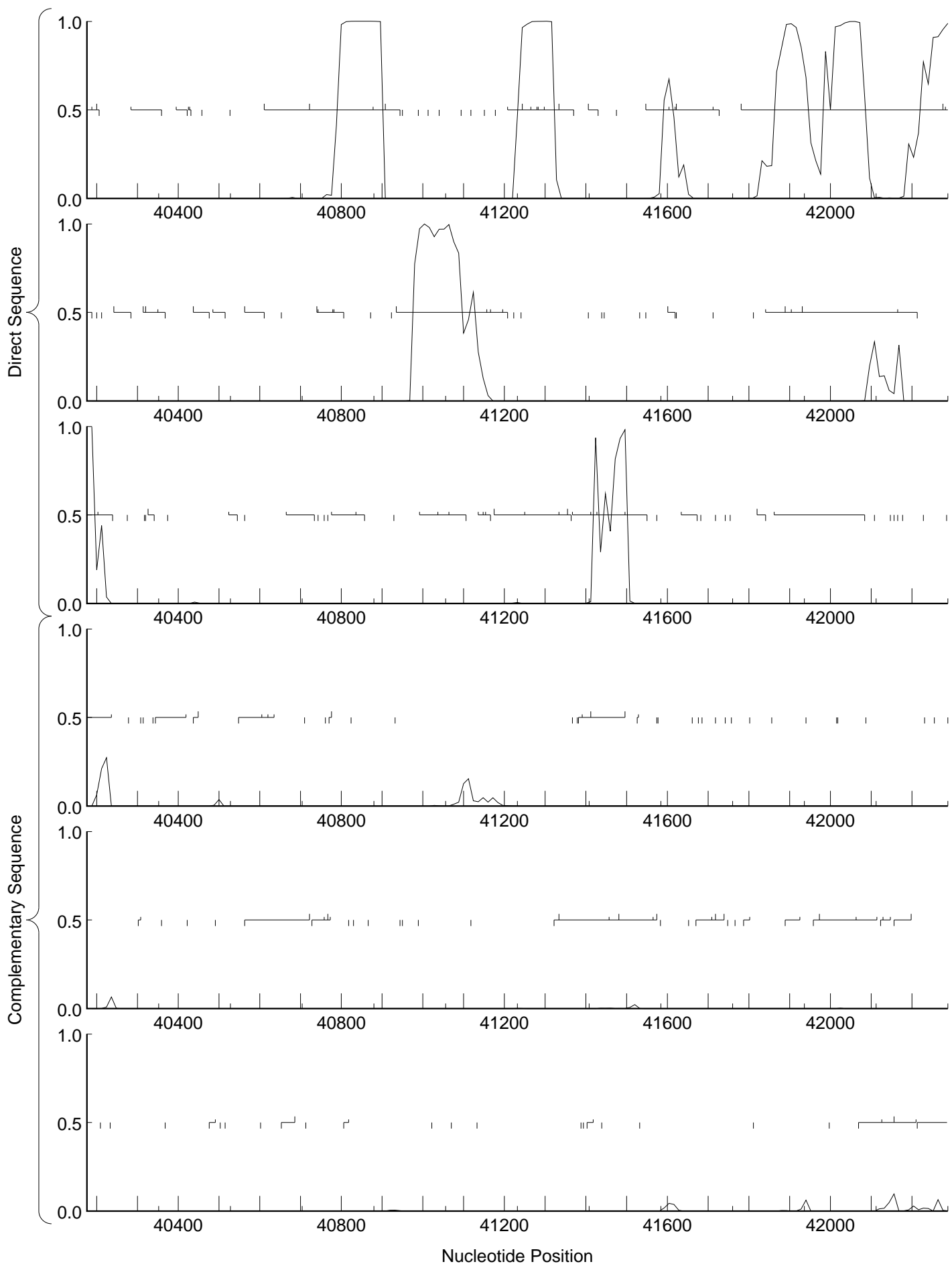


complete sequence, 73949 bp including 10-base 3' overhang (TCGAACGGCC), Cluster L1, Order 4, Window 96, Step 12, 19/37

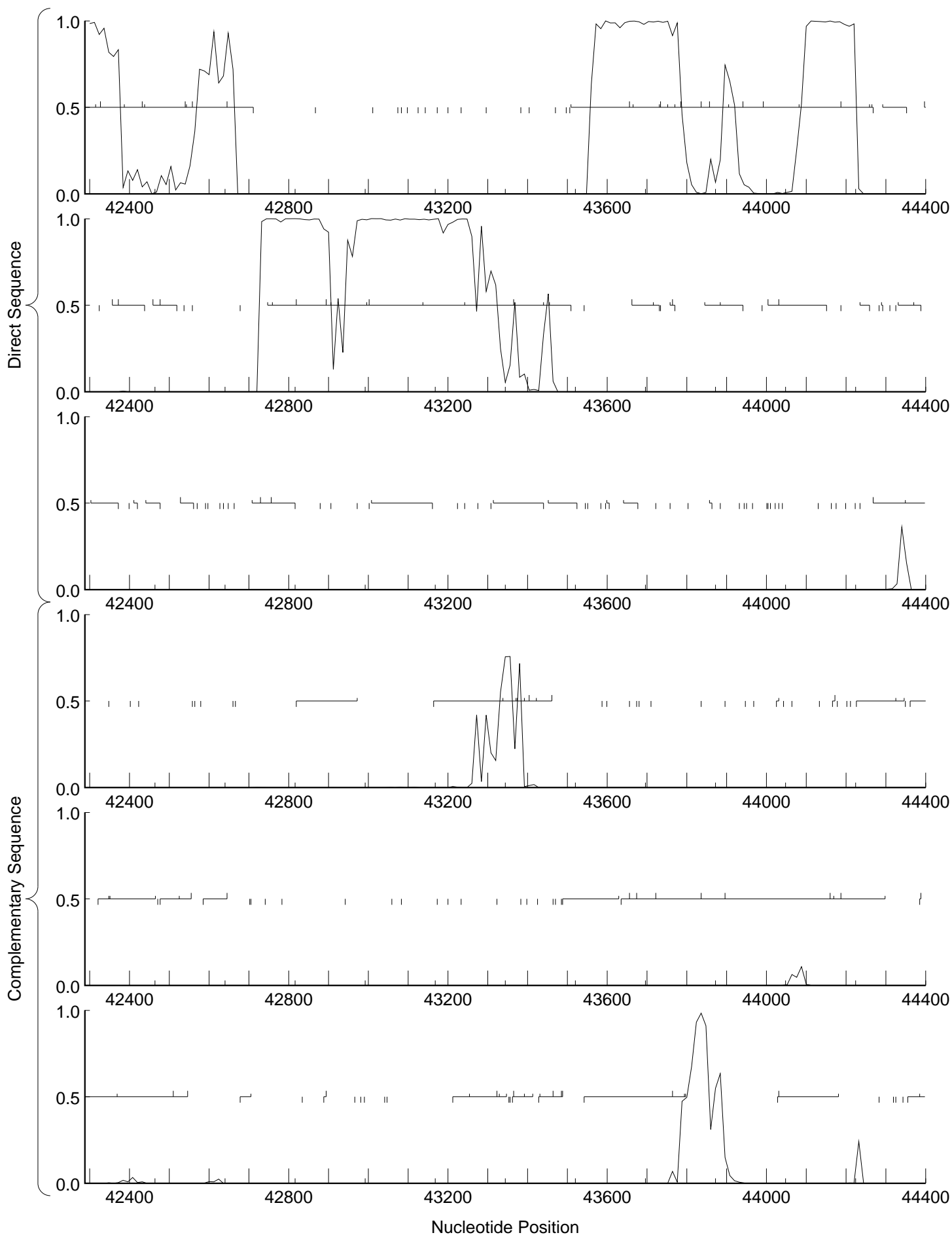


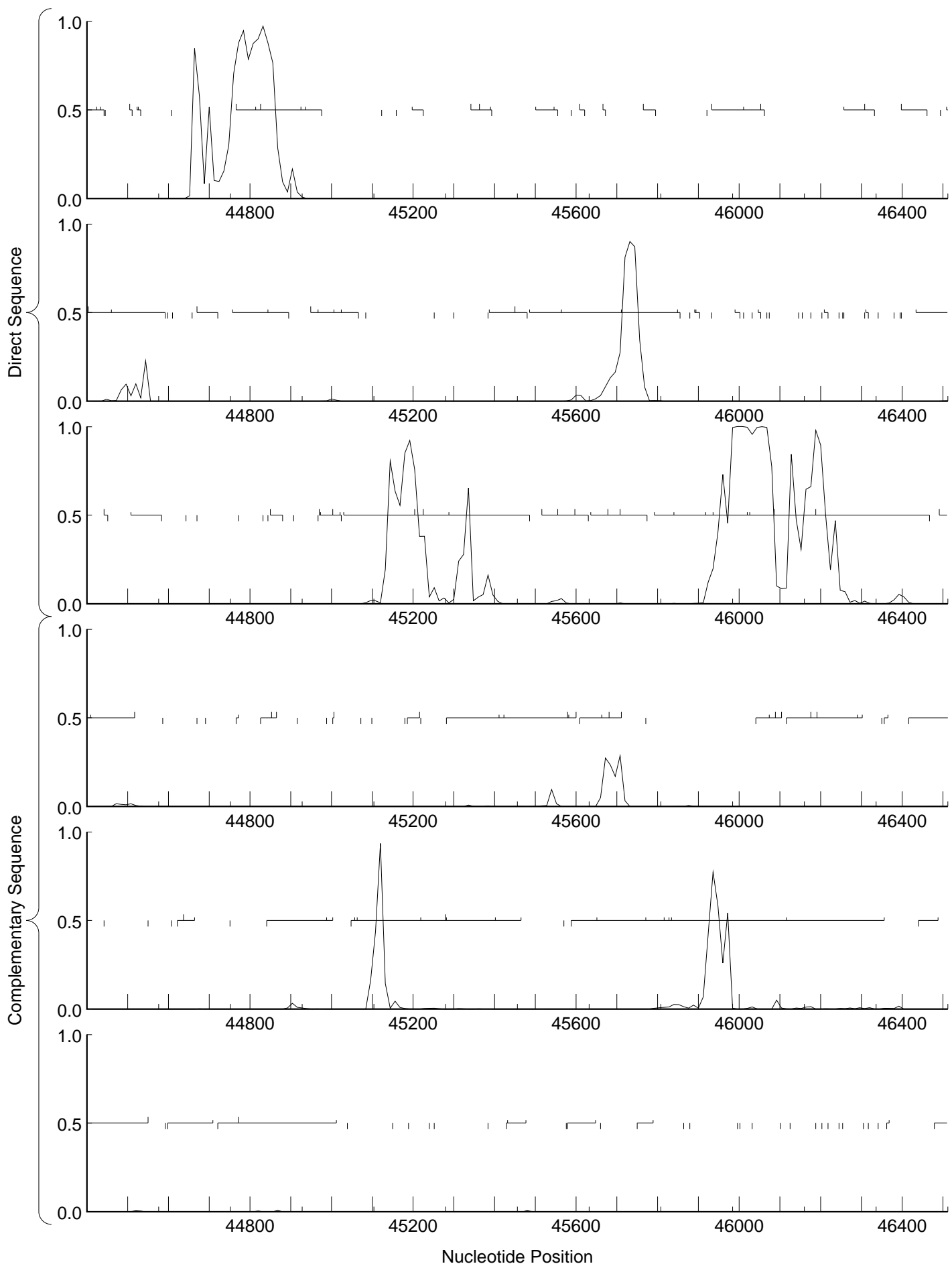


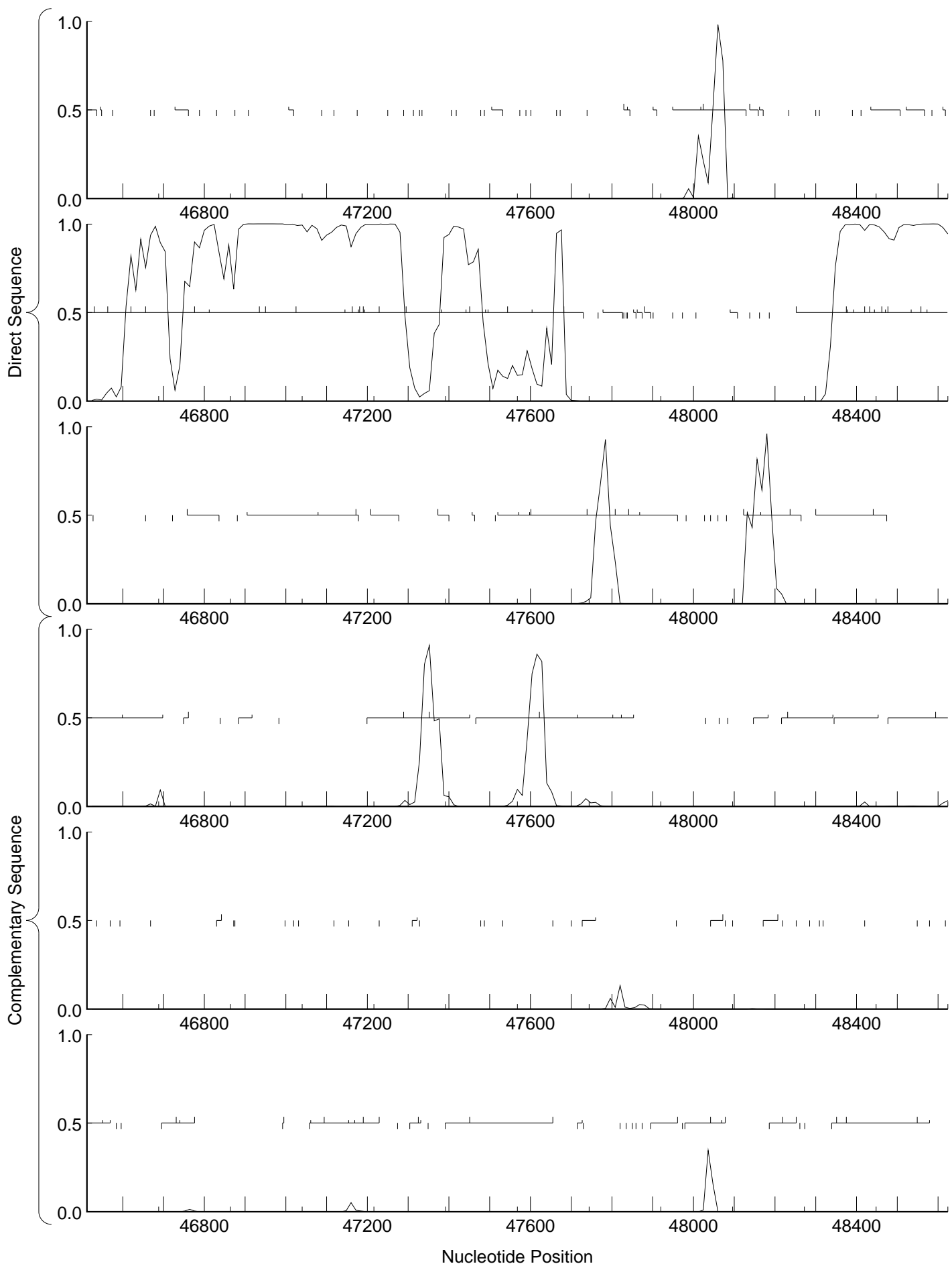
complete sequence, 73949 bp including 10-base 3' overhang (TCGAACGGCC), Cluster L1, Order 4, Window 96, Step 12, 21/37

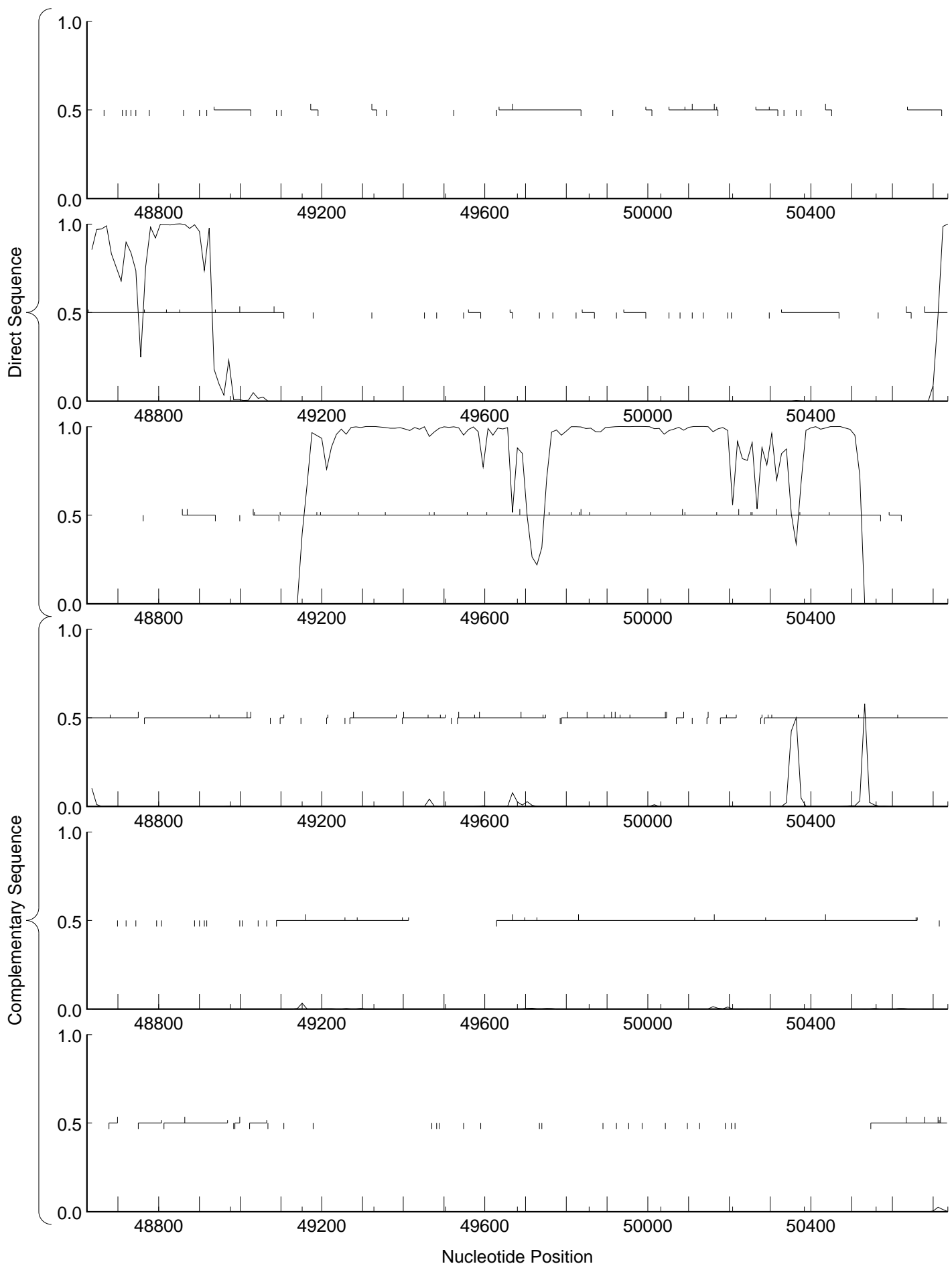


complete sequence, 73949 bp including 10-base 3' overhang (TCGAACGGCC), Cluster L1, Order 4, Window 96, Step 12, 22/37

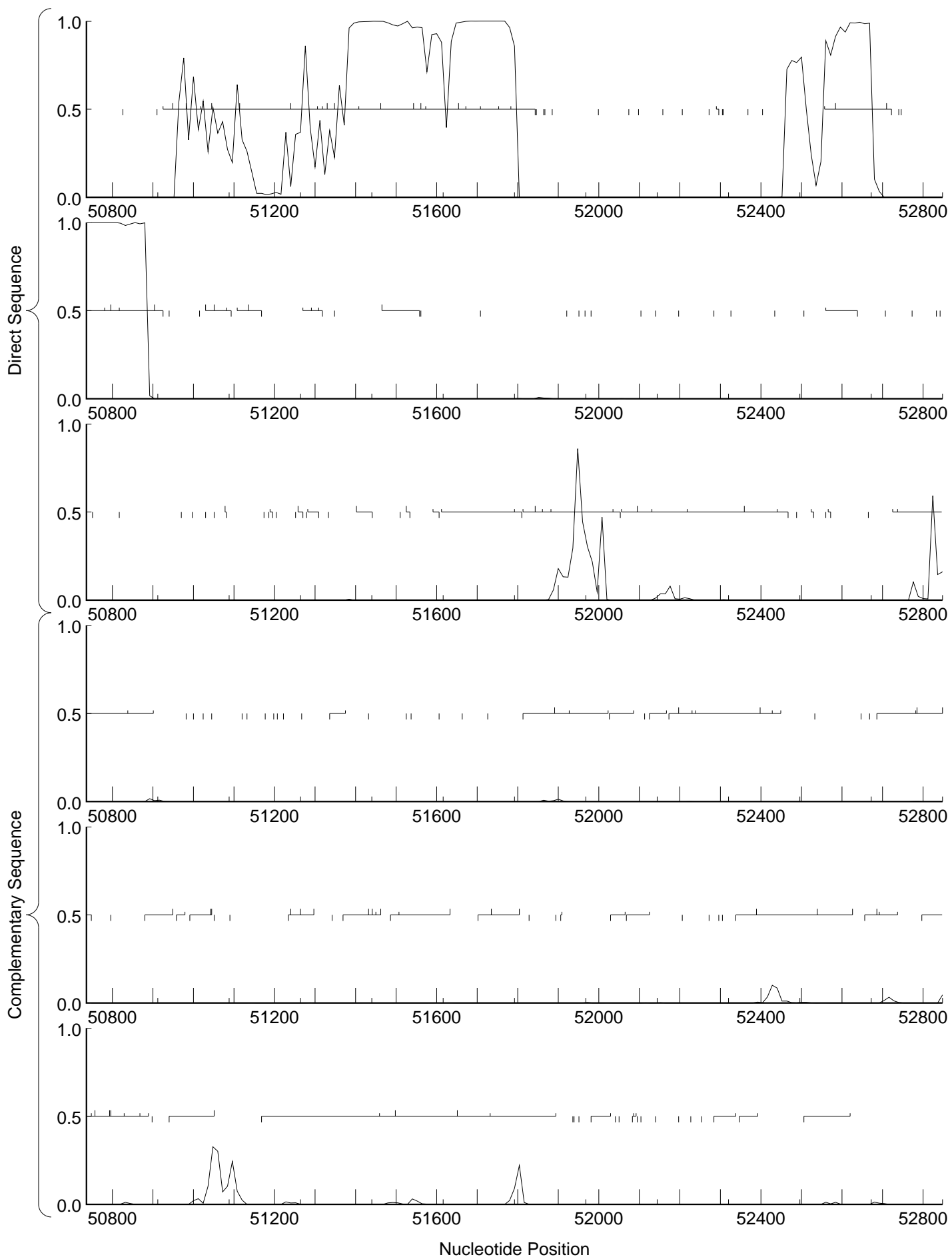


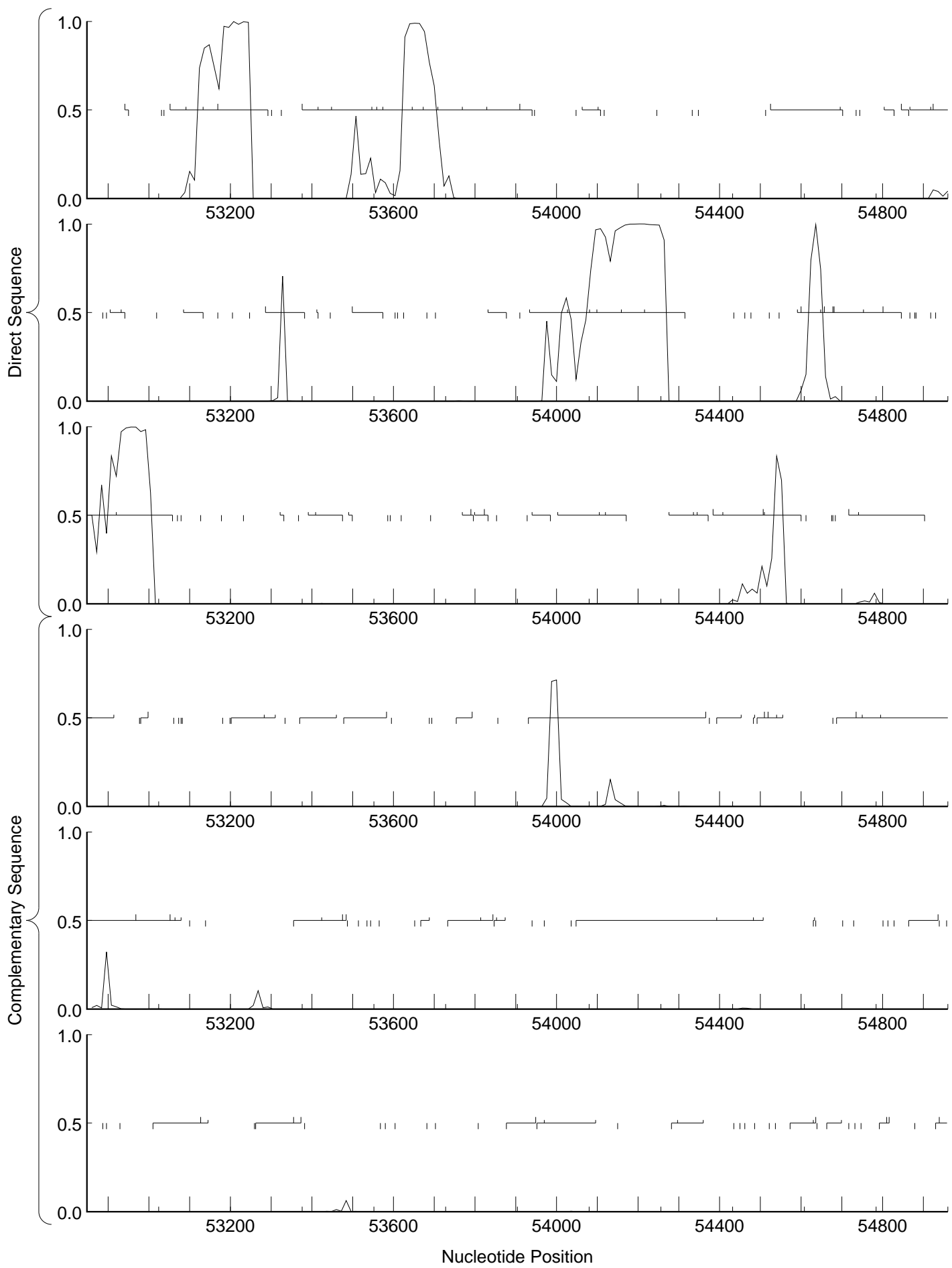


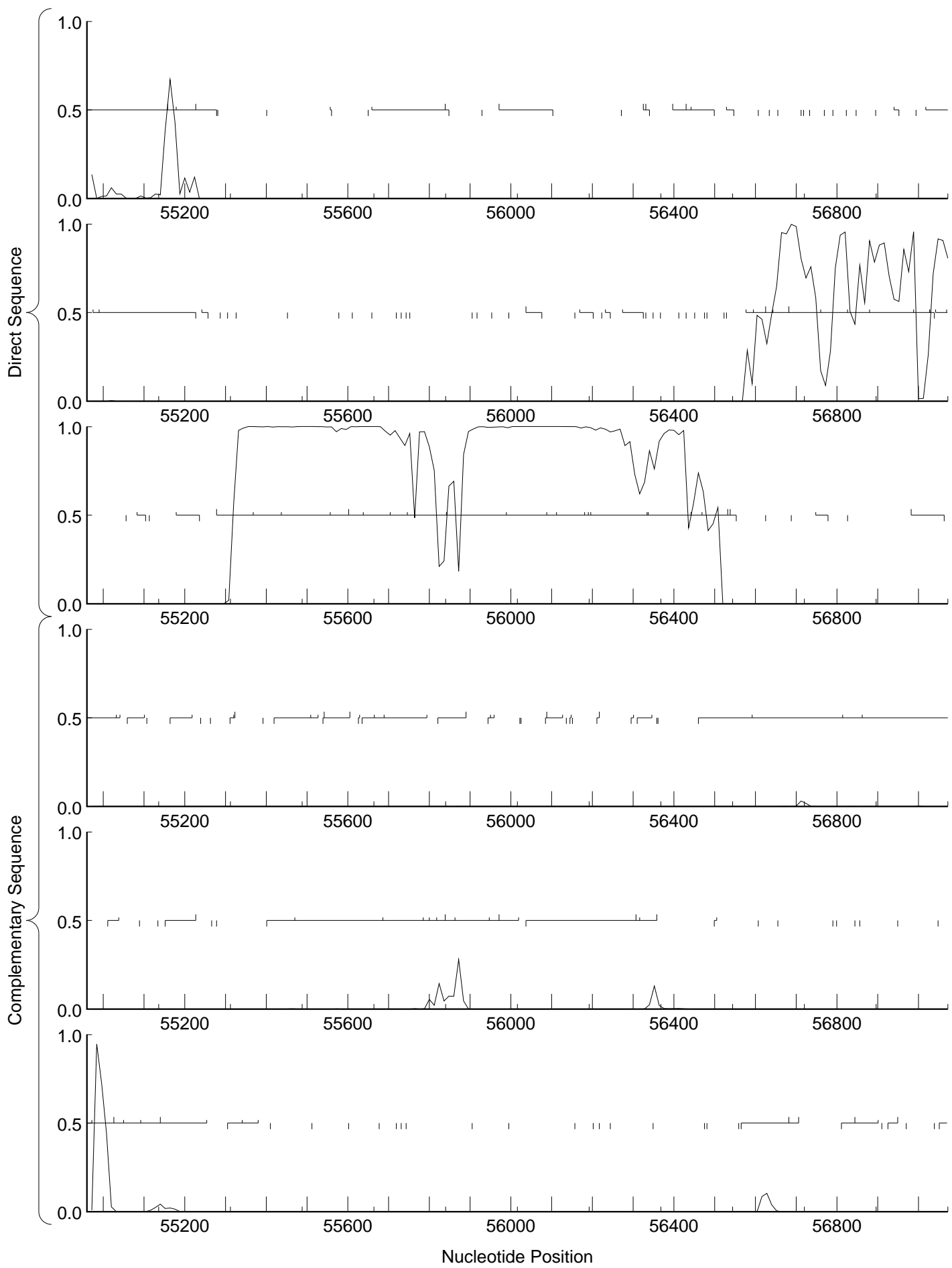


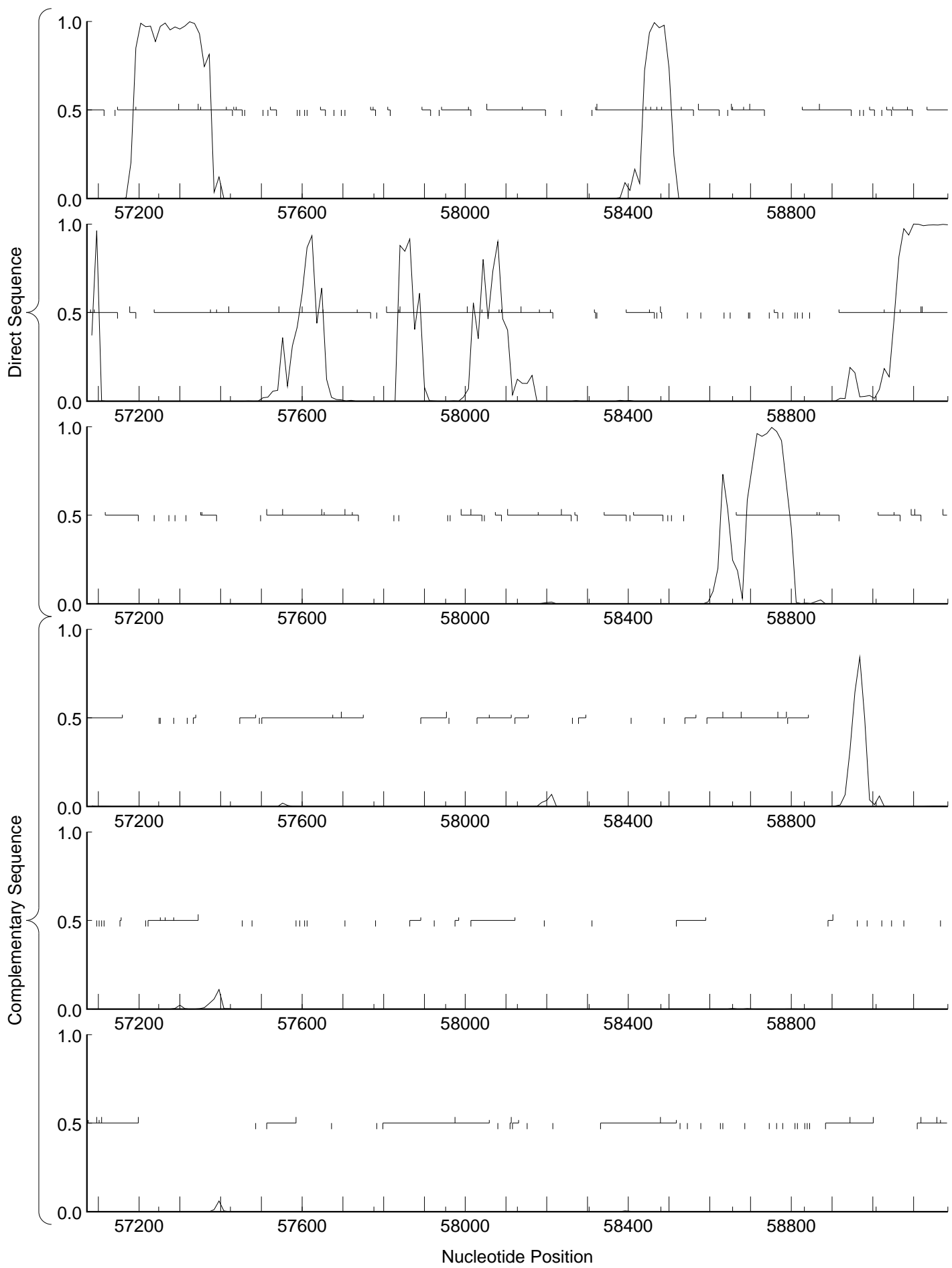


complete sequence, 73949 bp including 10-base 3' overhang (TCGAACGGCC), Cluster L1, Order 4, Window 96, Step 12, 26/37

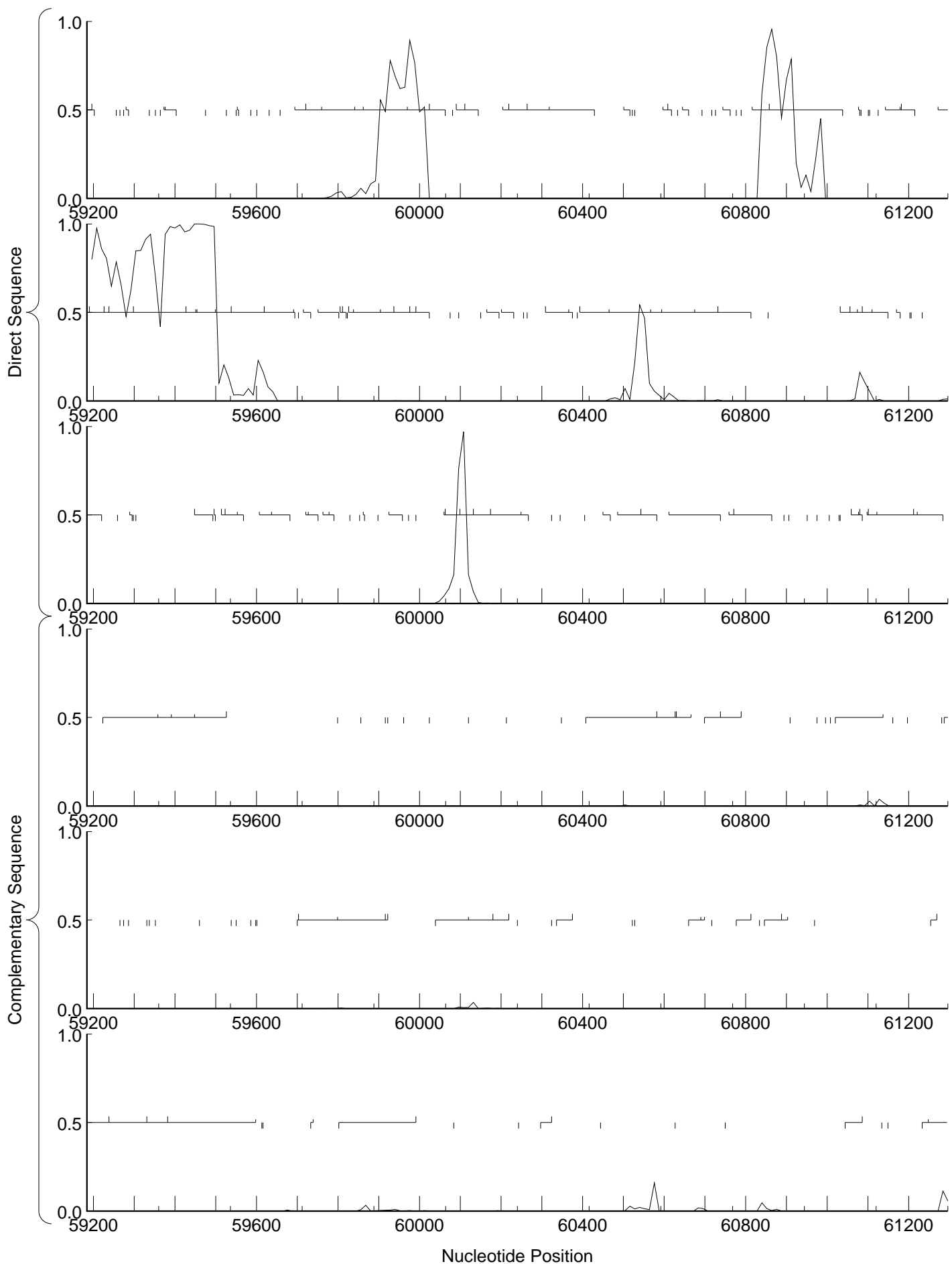


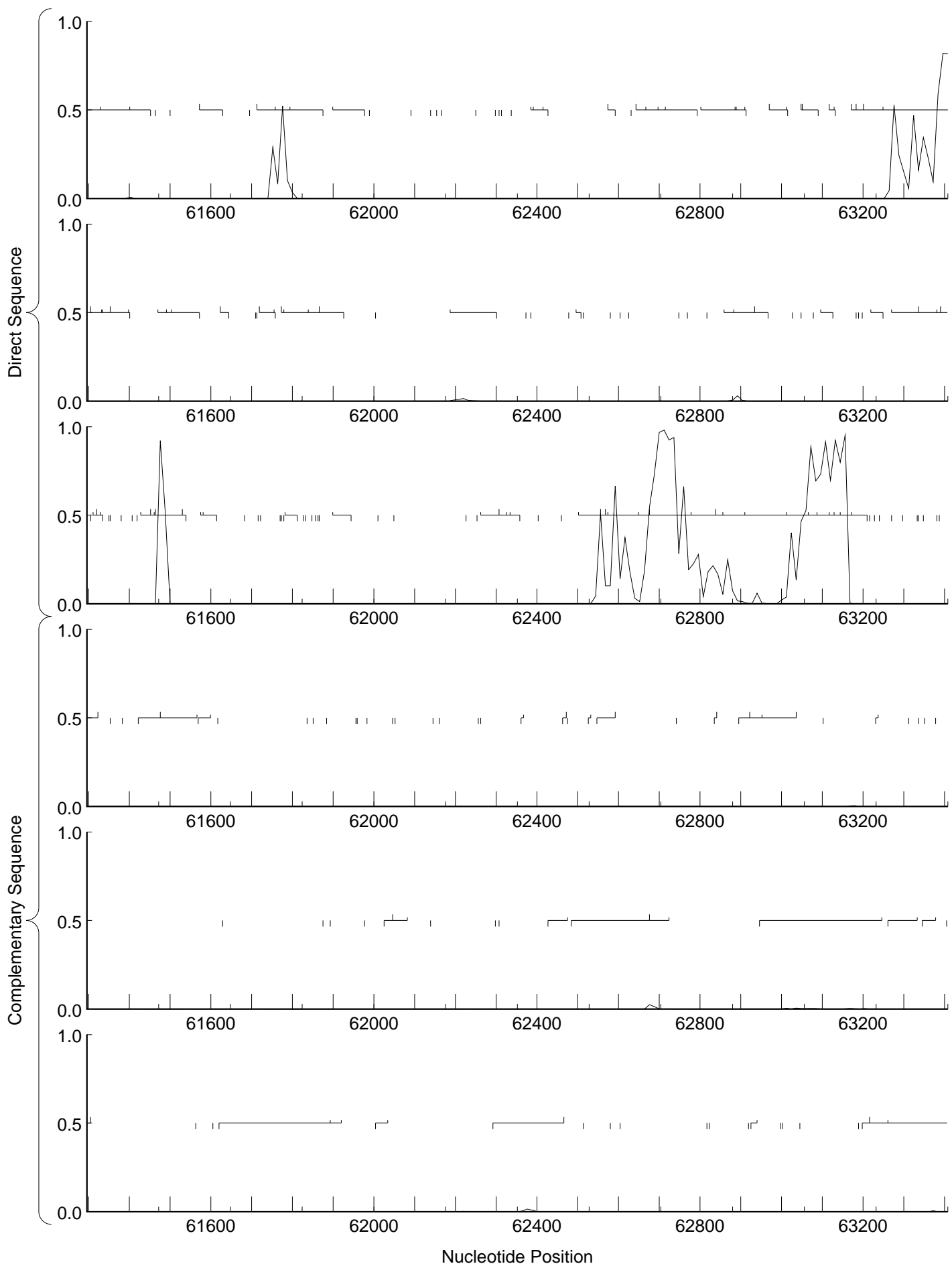


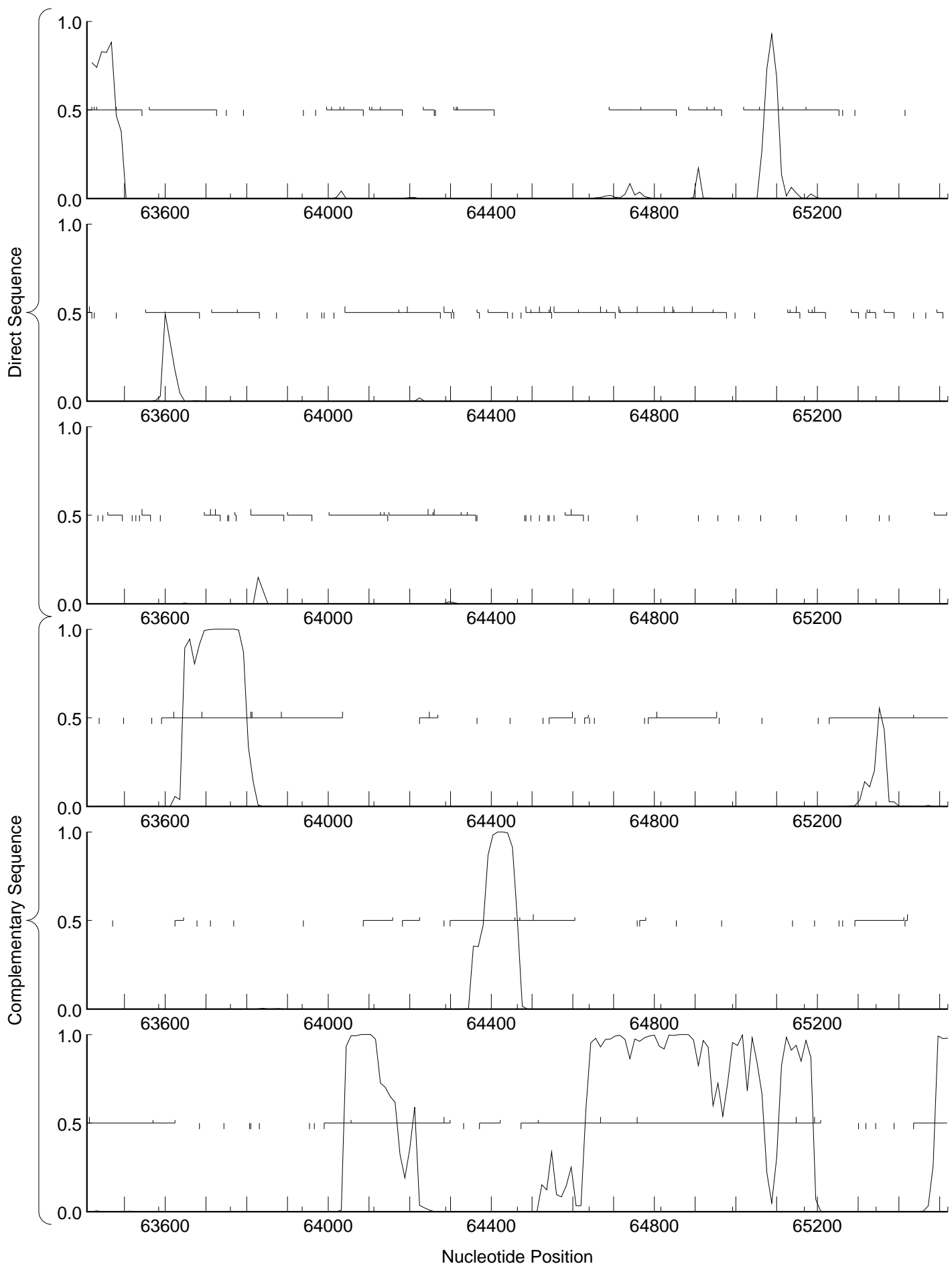


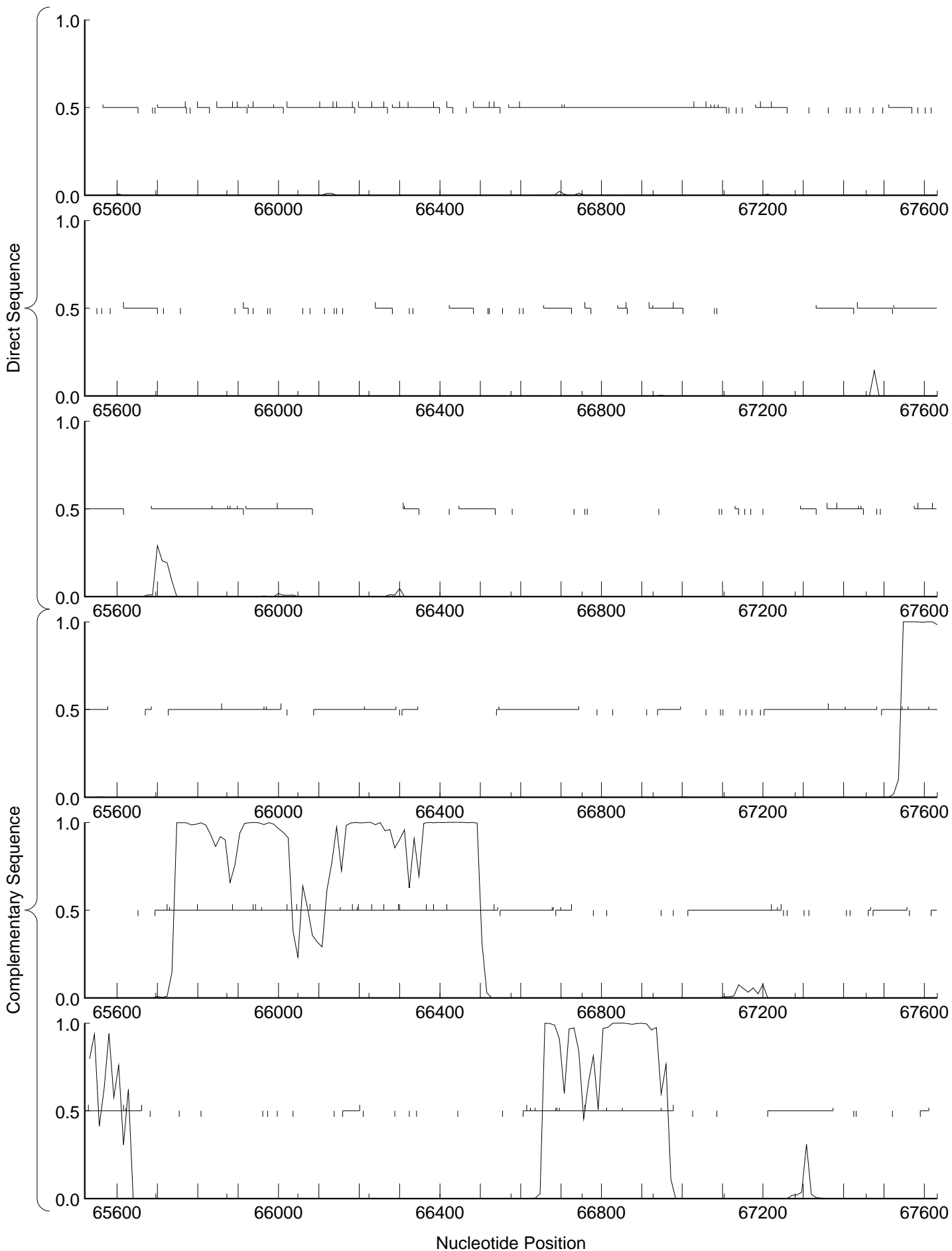


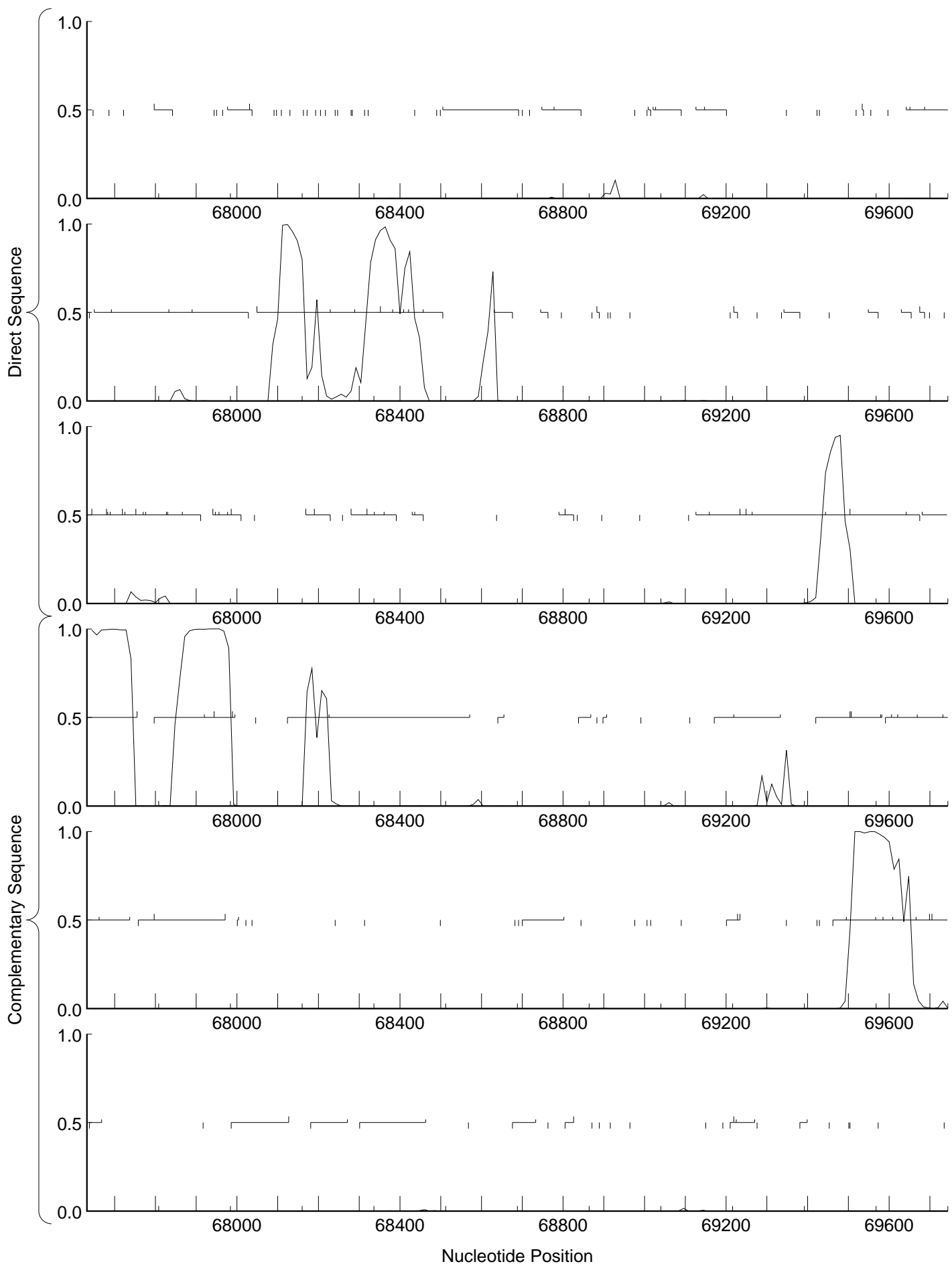
complete sequence, 73949 bp including 10-base 3' overhang (TCGAACGGCC), Cluster L1, Order 4, Window 96, Step 12, 30/37











complete sequence, 73949 bp including 10-base 3' overhang (TCGAACGGCC), Cluster L1, Order 4, Window 96, Step 12, 35/37

