

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium Phage Mangethe complete sequence, 47,612 bp including 12-base 3' overhang (CC
Analysis Date : 9/27/21 at 6:23:58
Pages : 24
Sequence Length : 47612 bp
GC Content : 67.37%

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

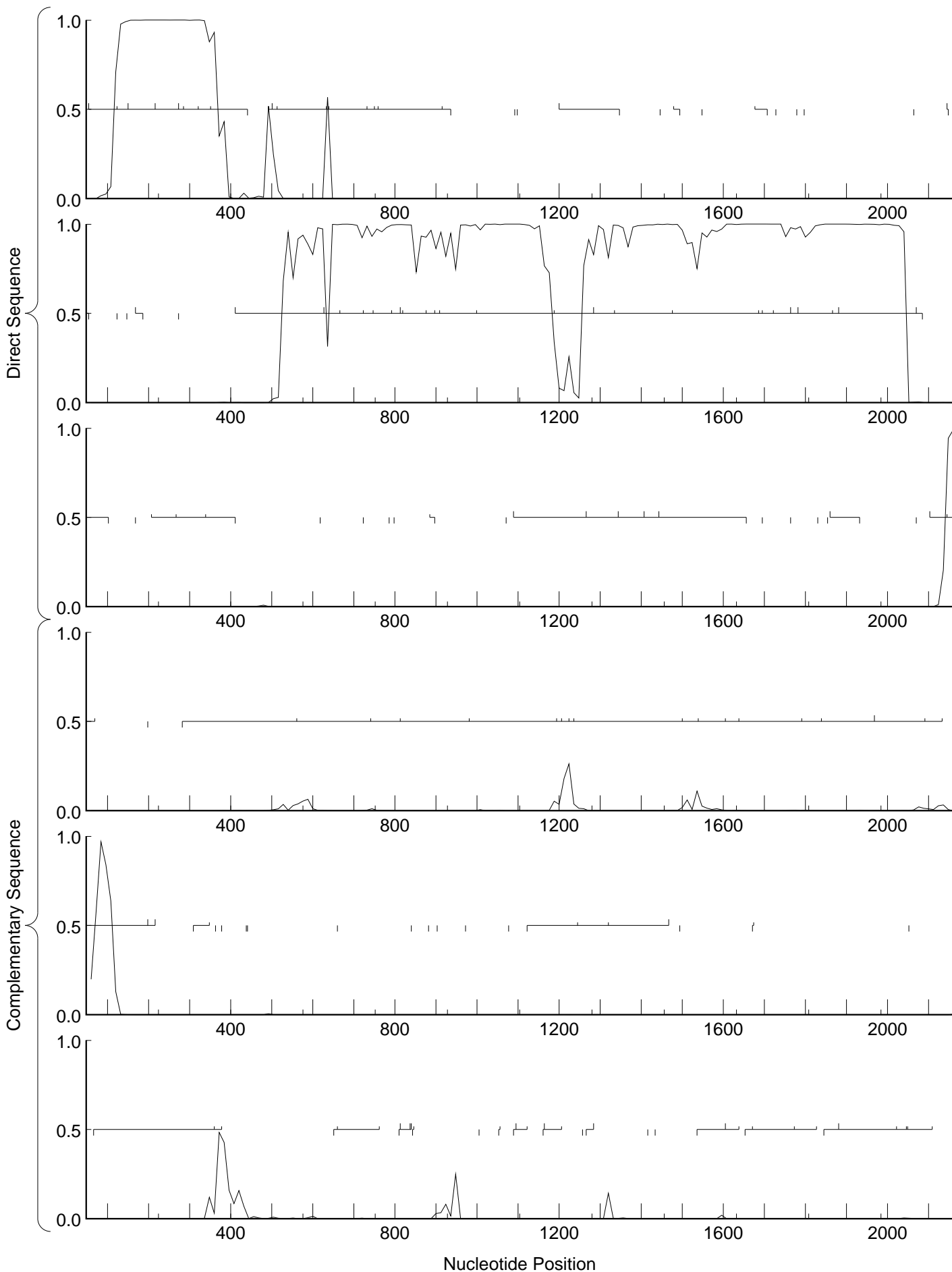
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

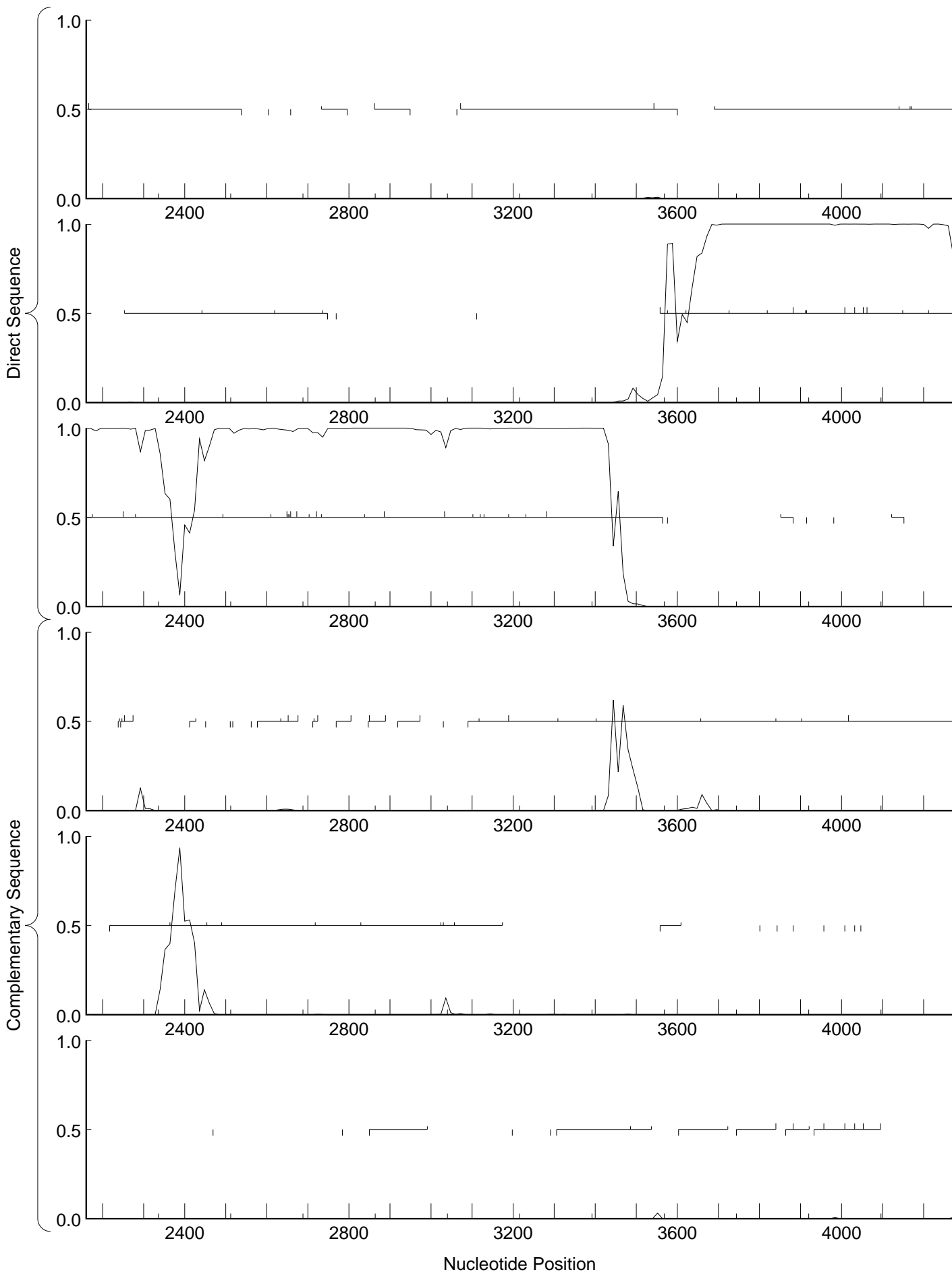
MATRIX INFORMATION

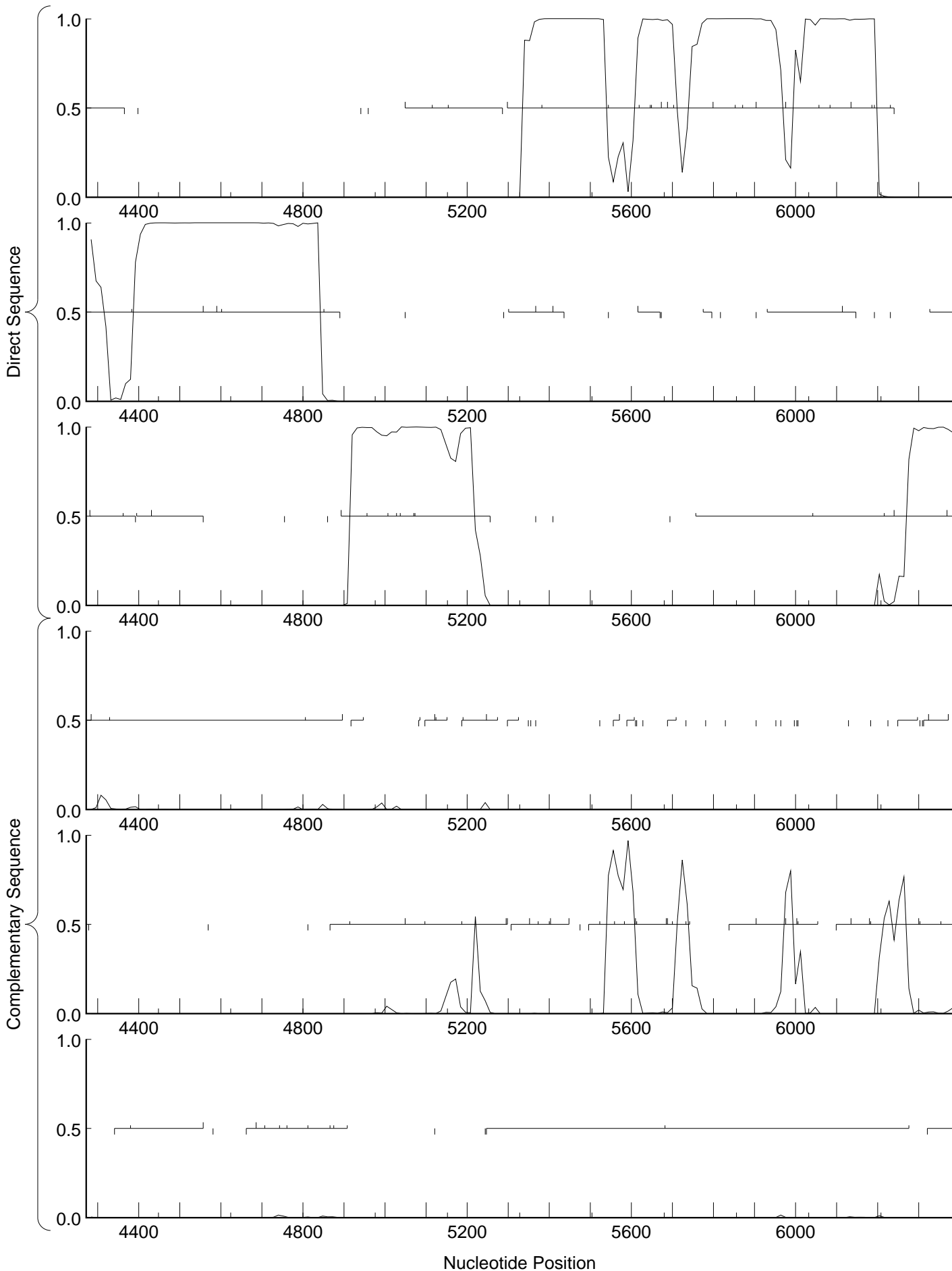
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

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

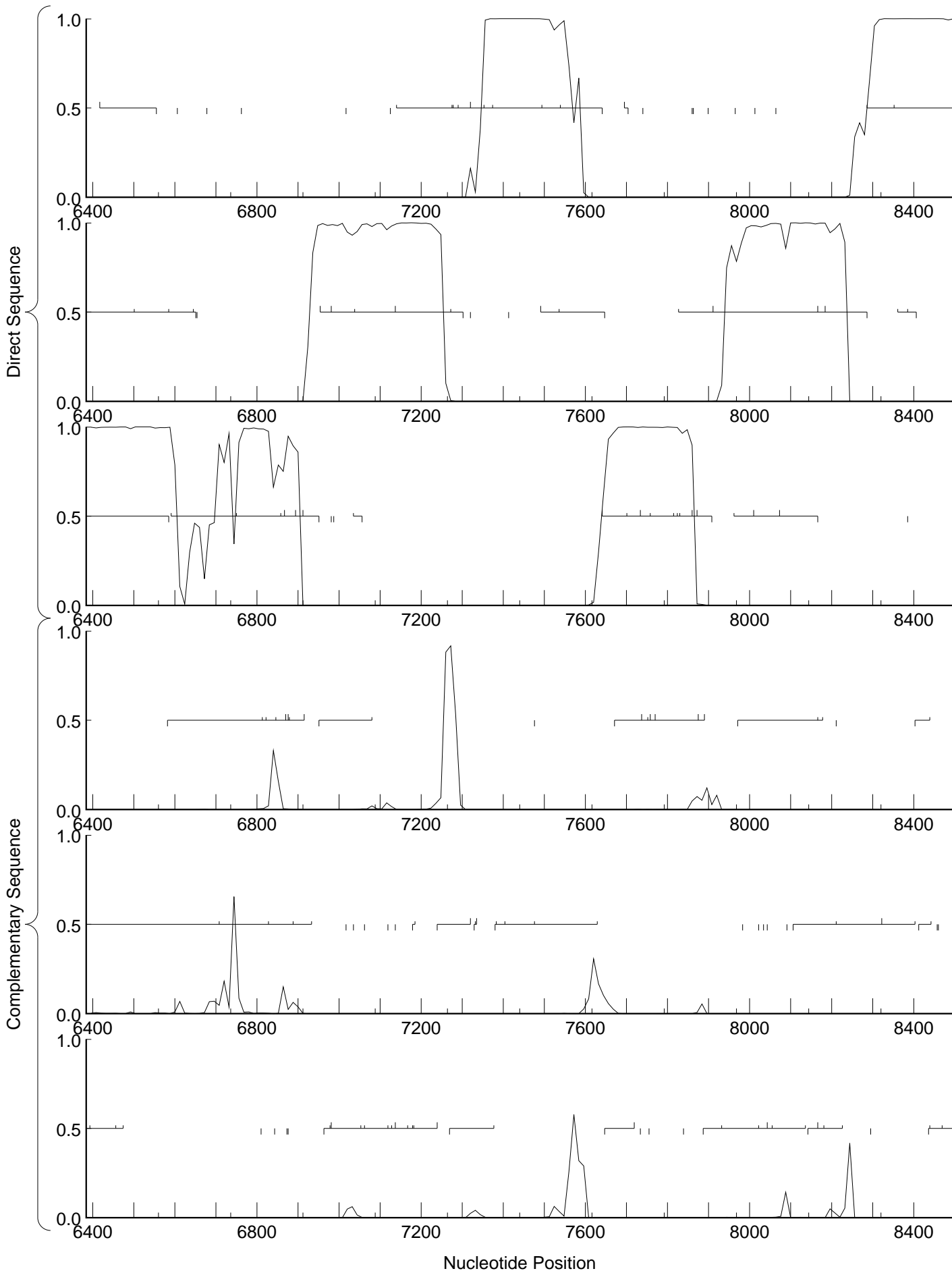
For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

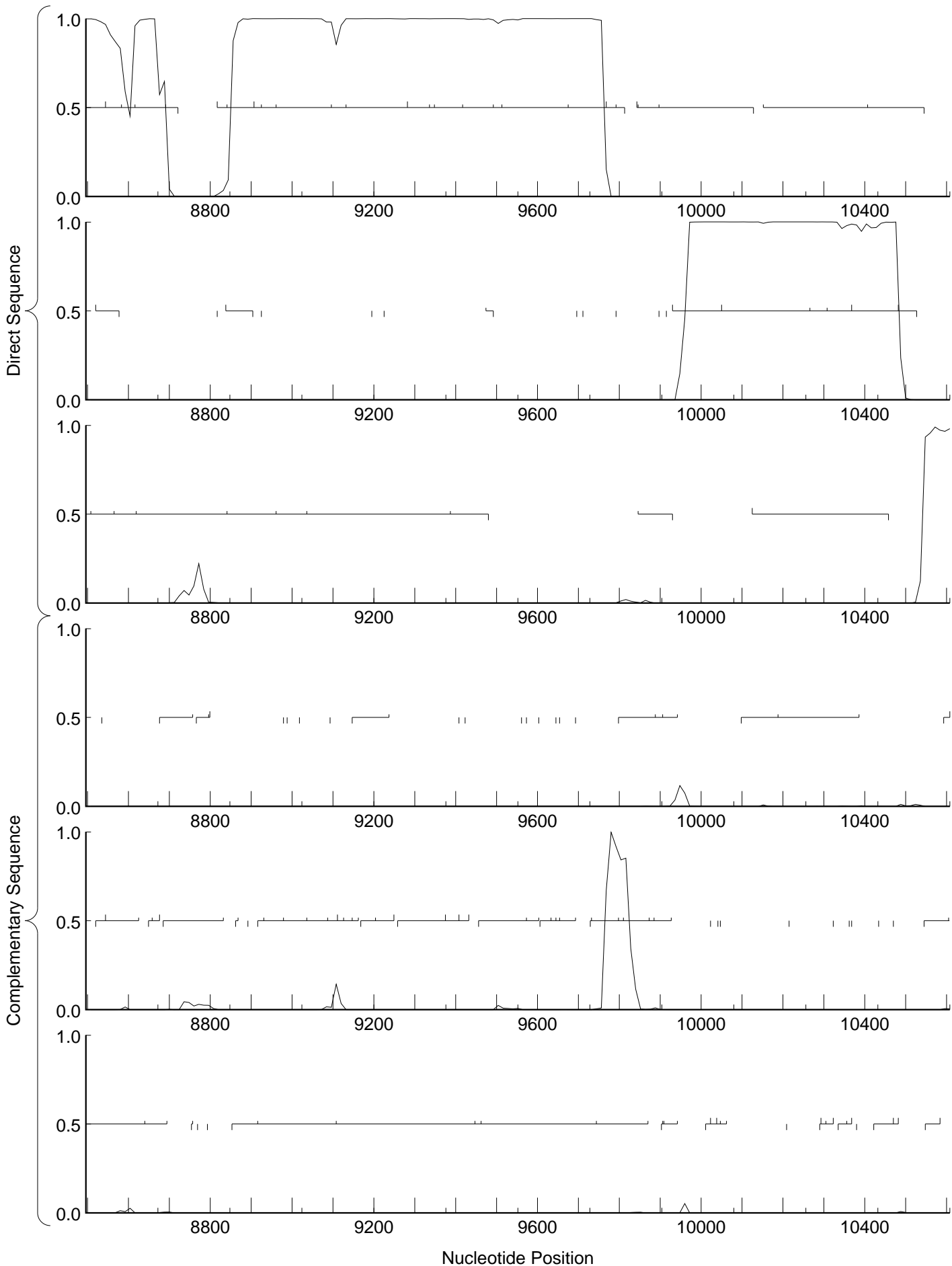




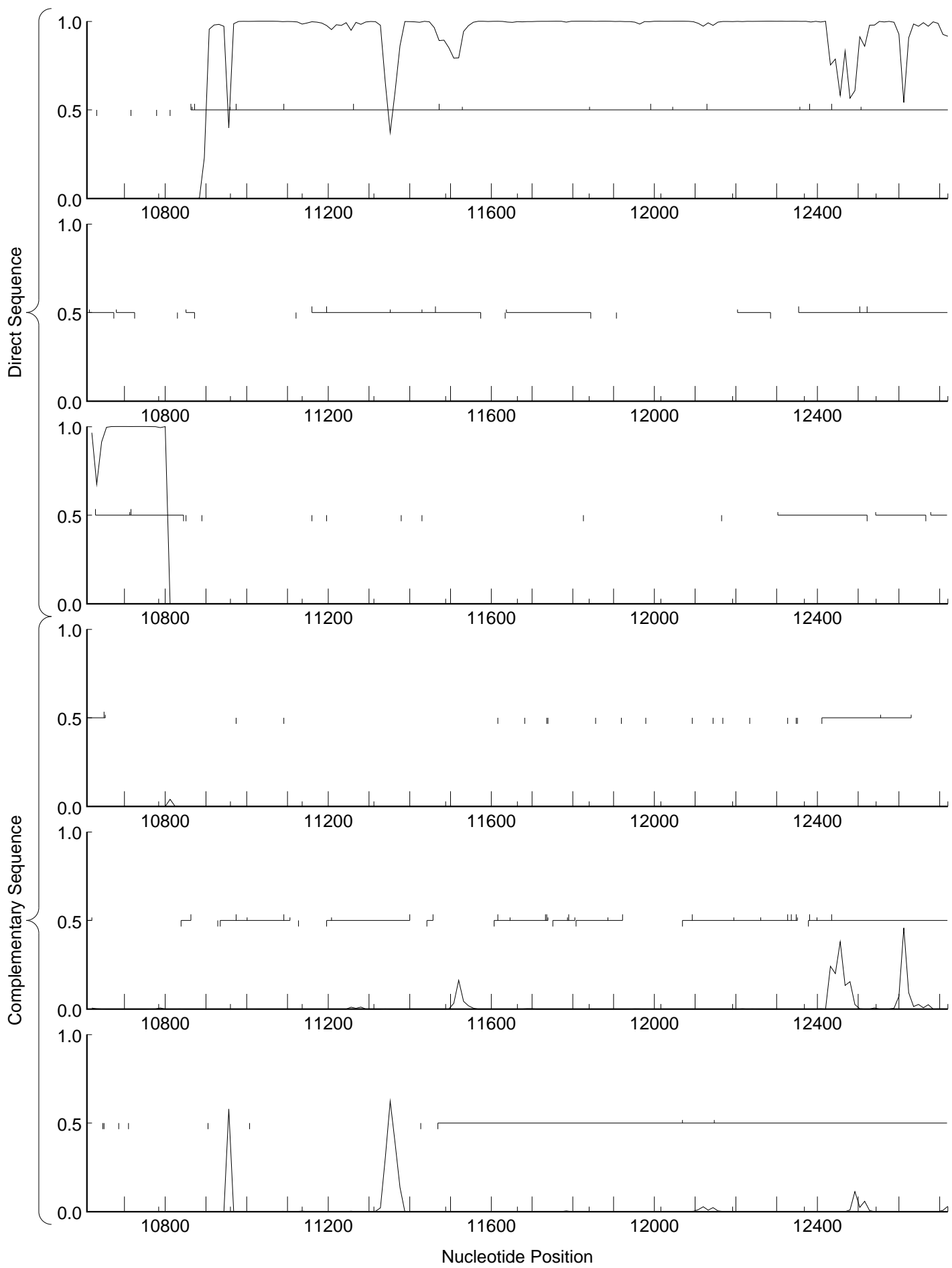


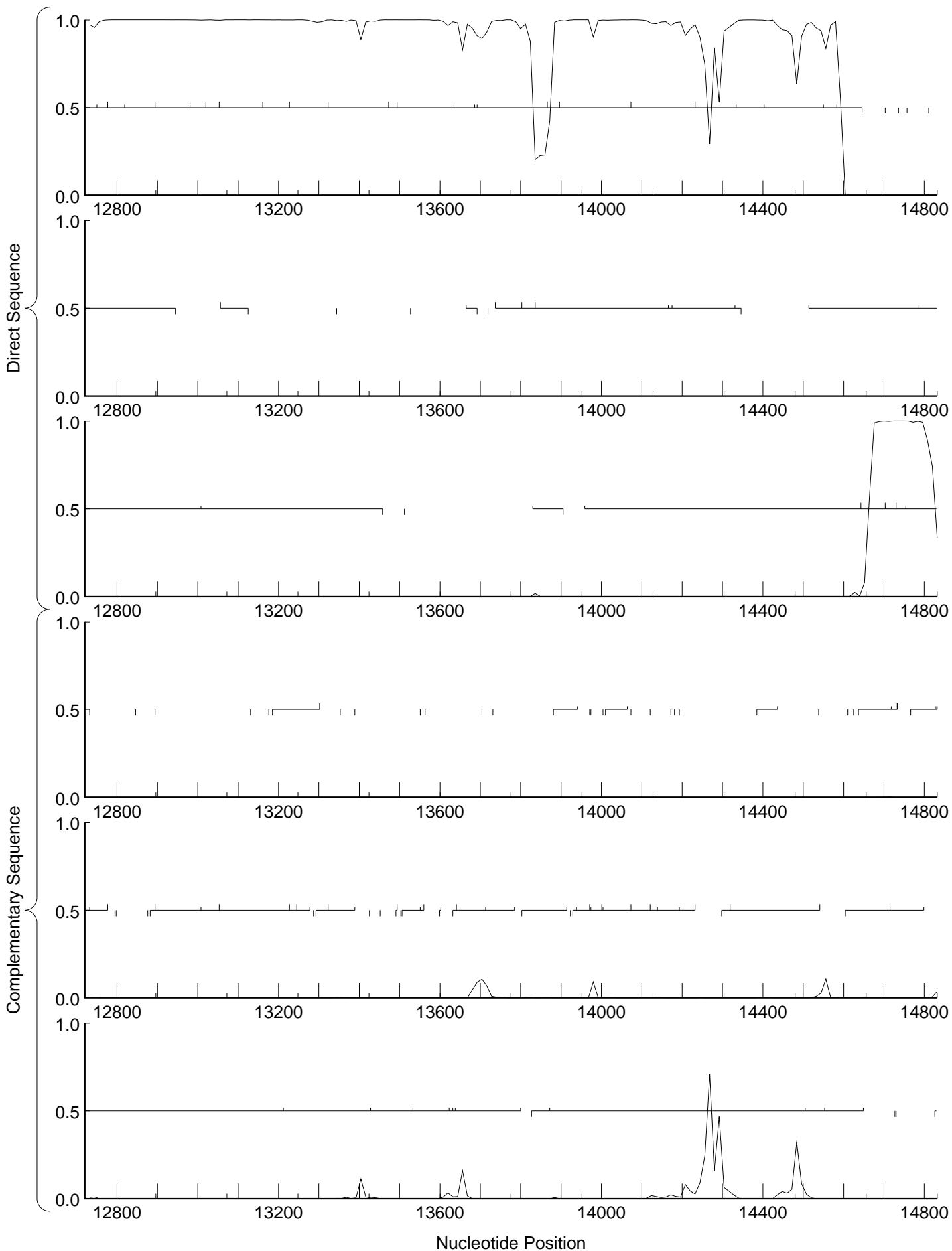
complete sequence, 47,612 bp including 12-base 3' overhang (CCCGCCCCCGA), Cluster P, Order 4, Window 96, Step 12, 5/24

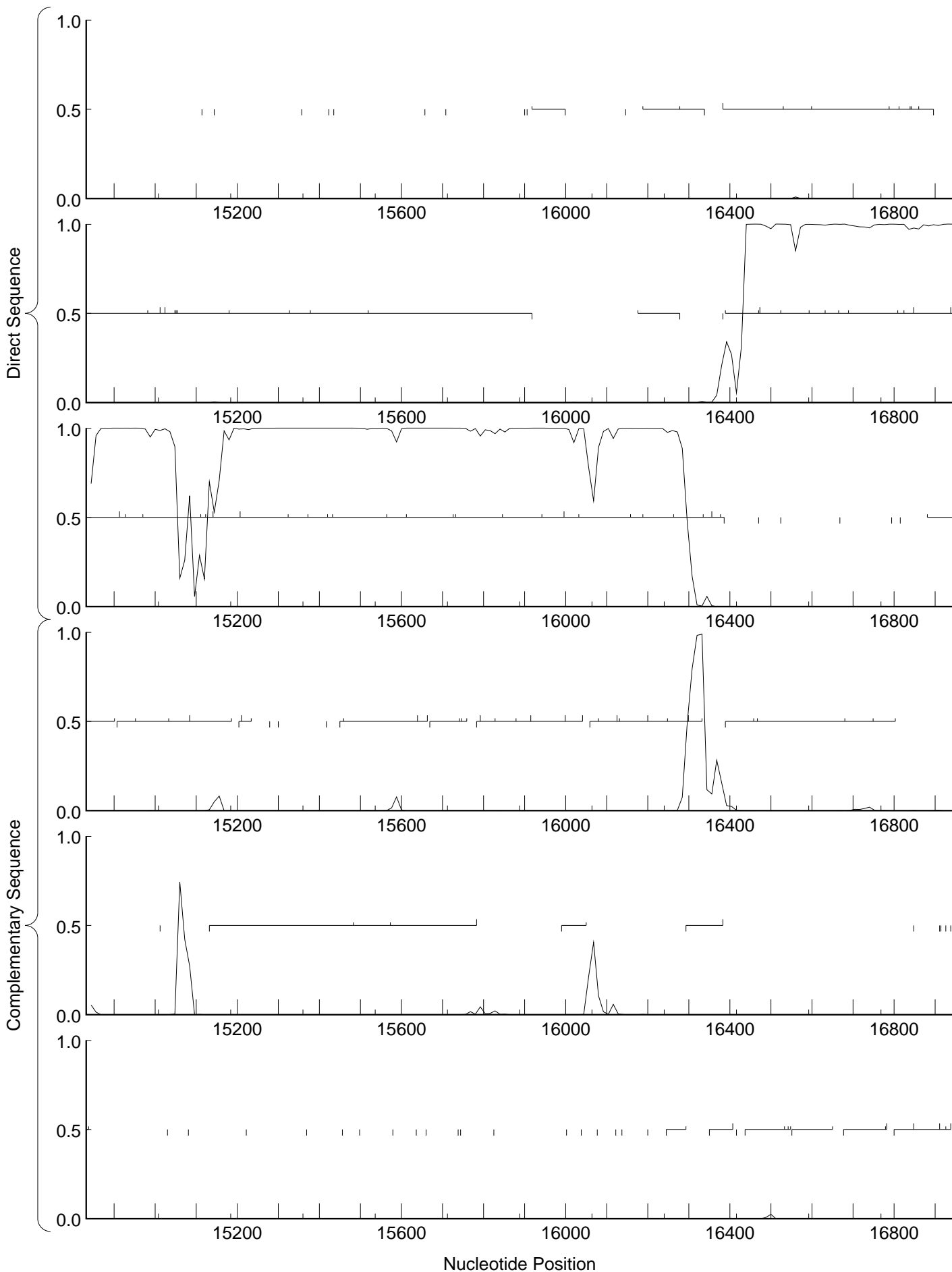


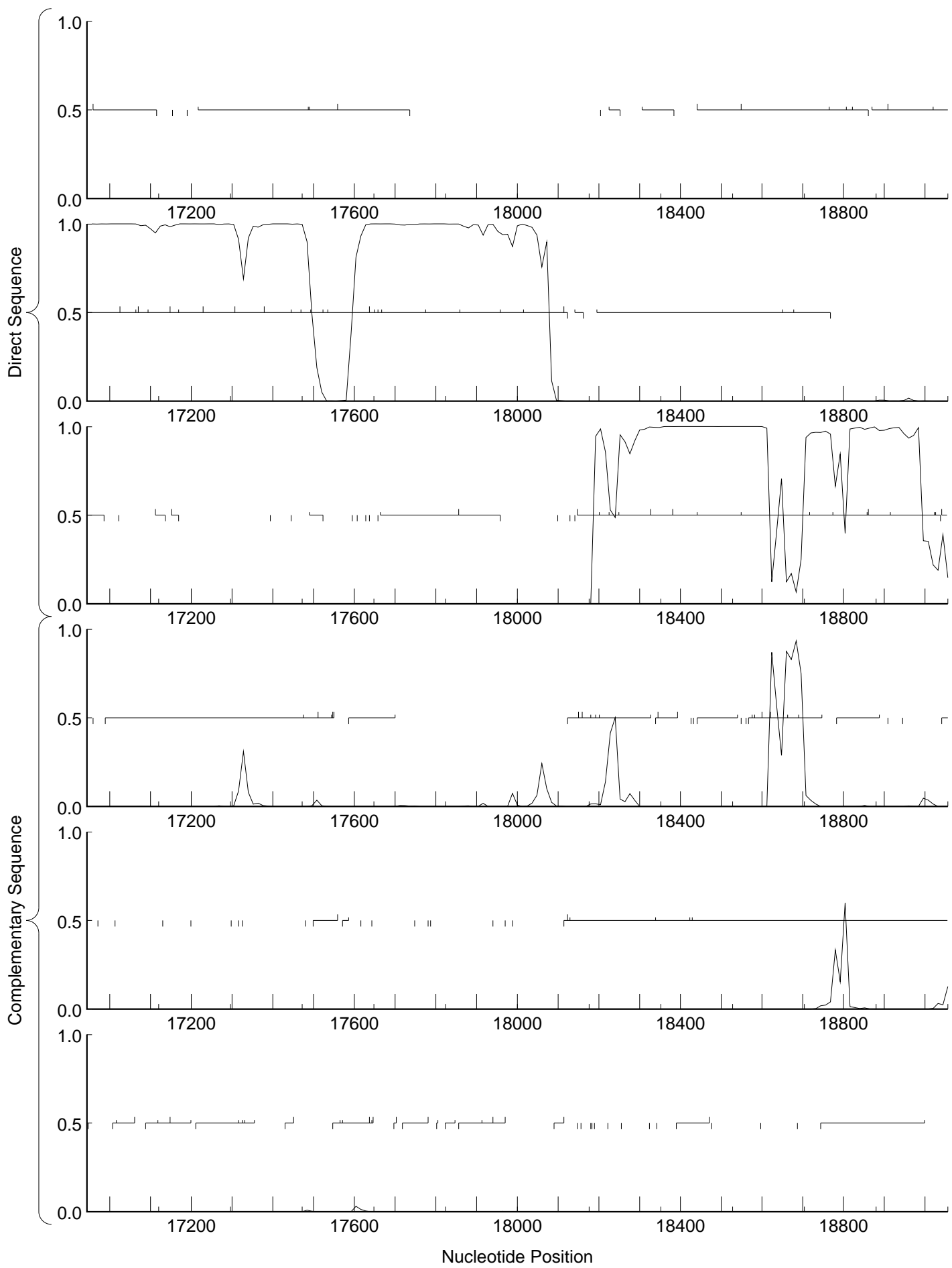


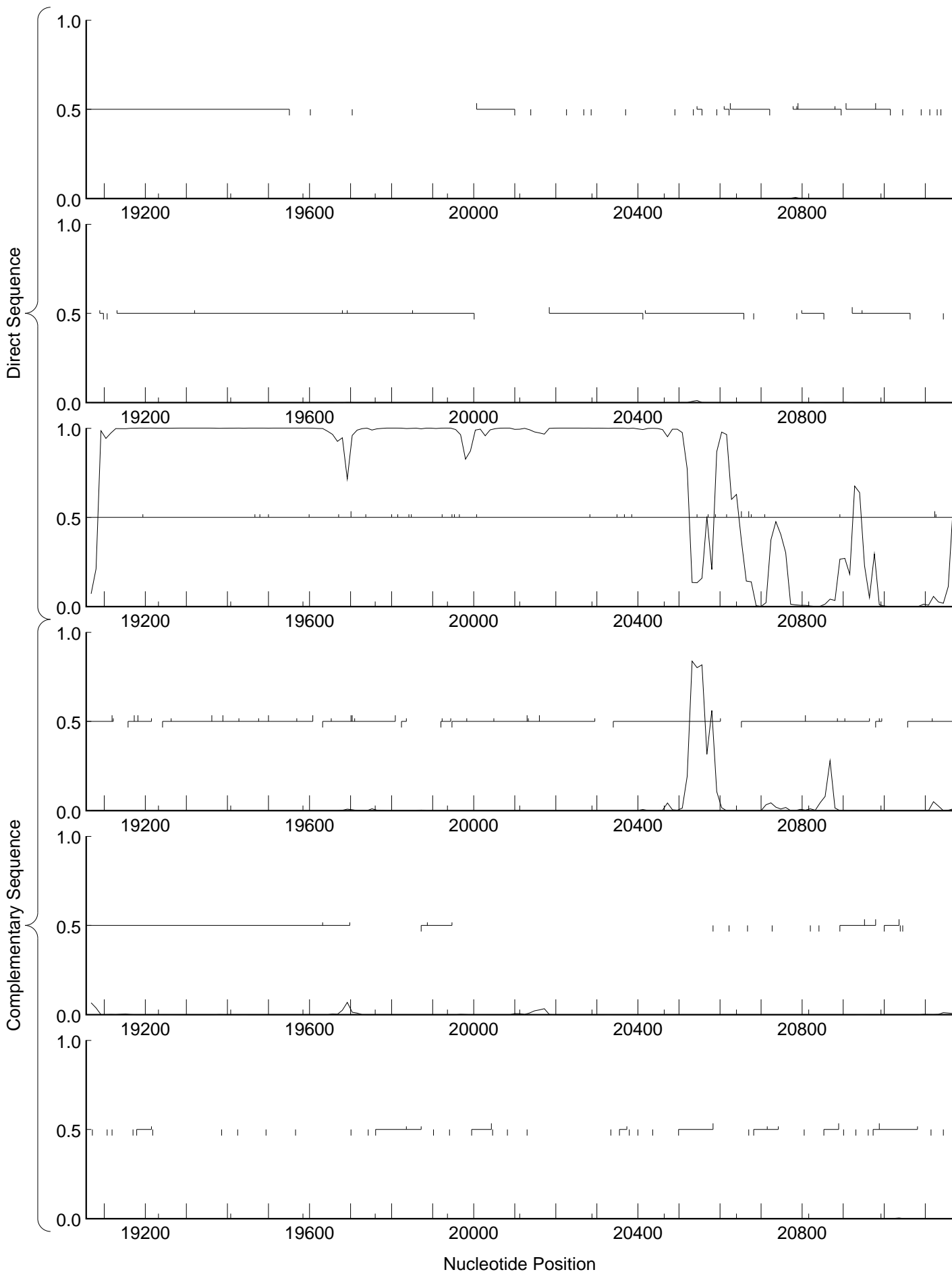
complete sequence, 47,612 bp including 12-base 3' overhang (CCCGCCCCCGA), Cluster P, Order 4, Window 96, Step 12, 7/24

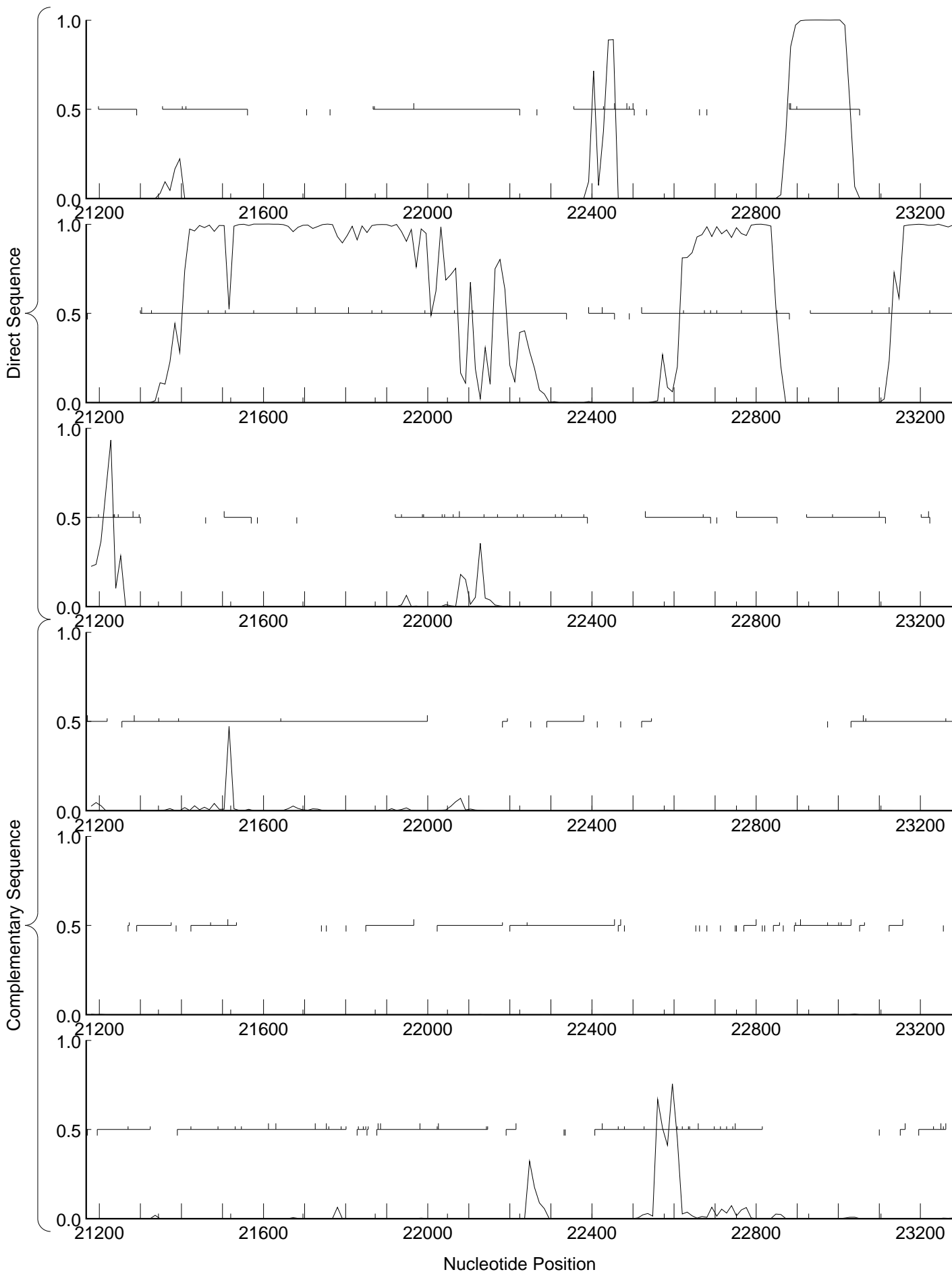




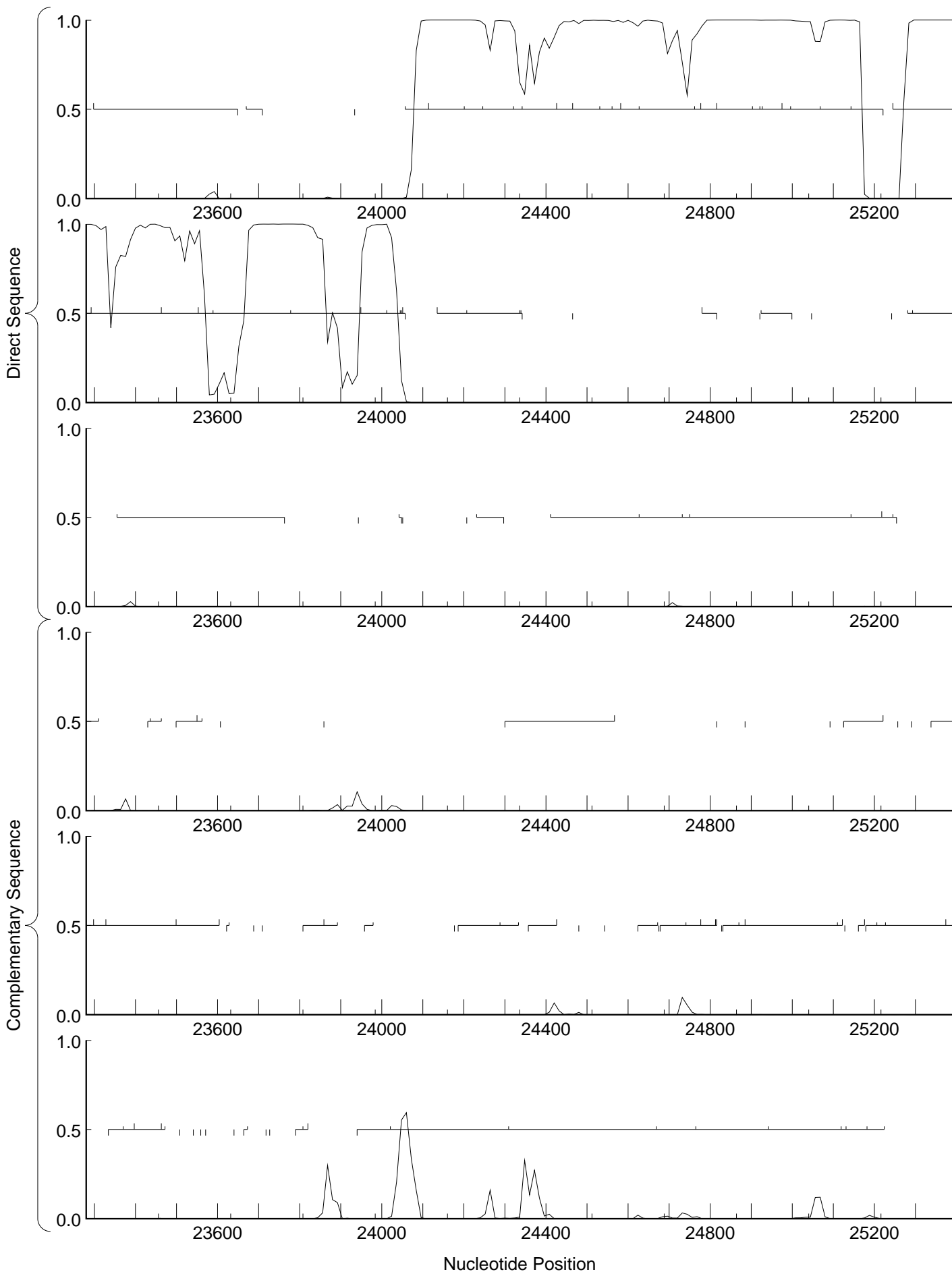


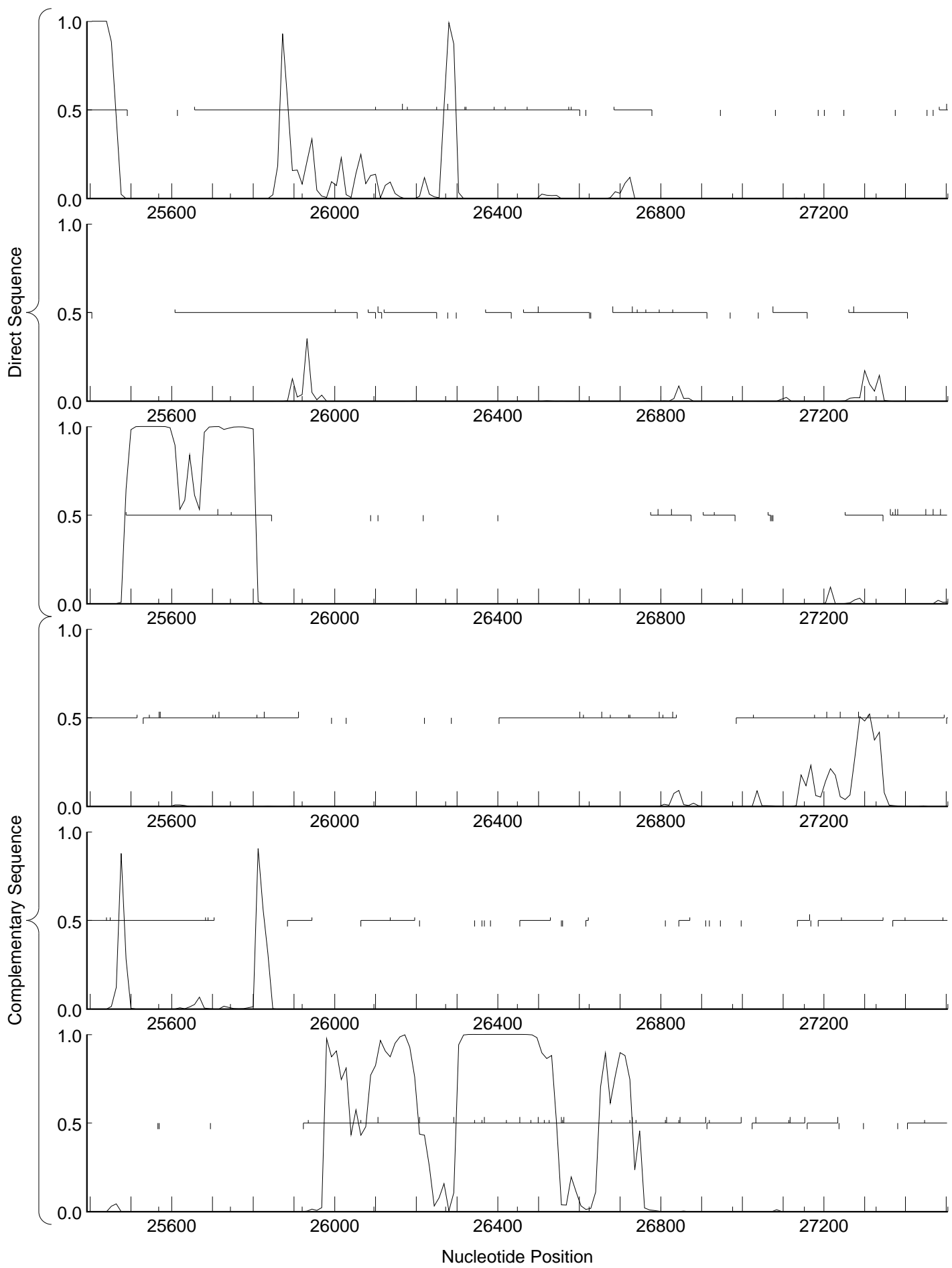


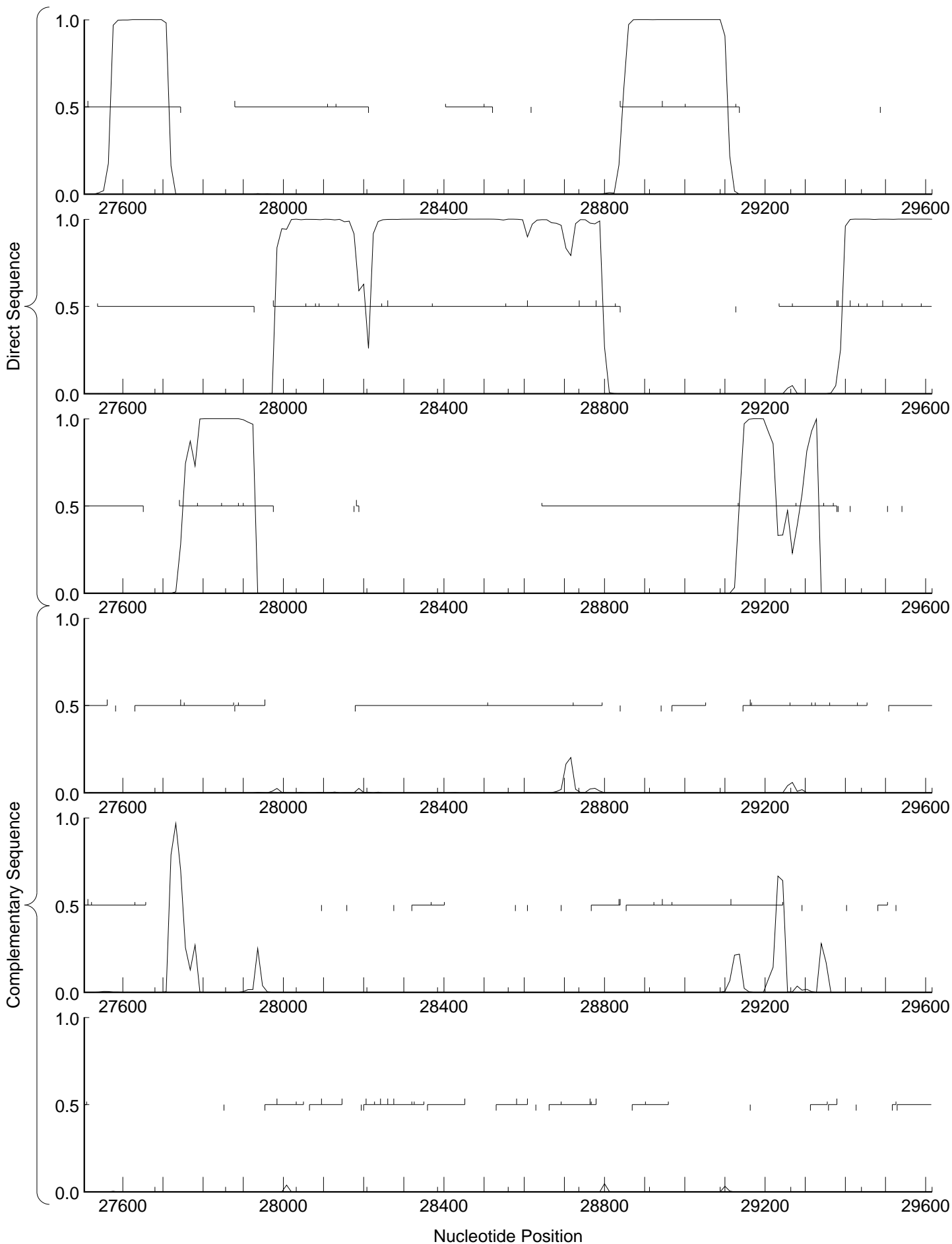


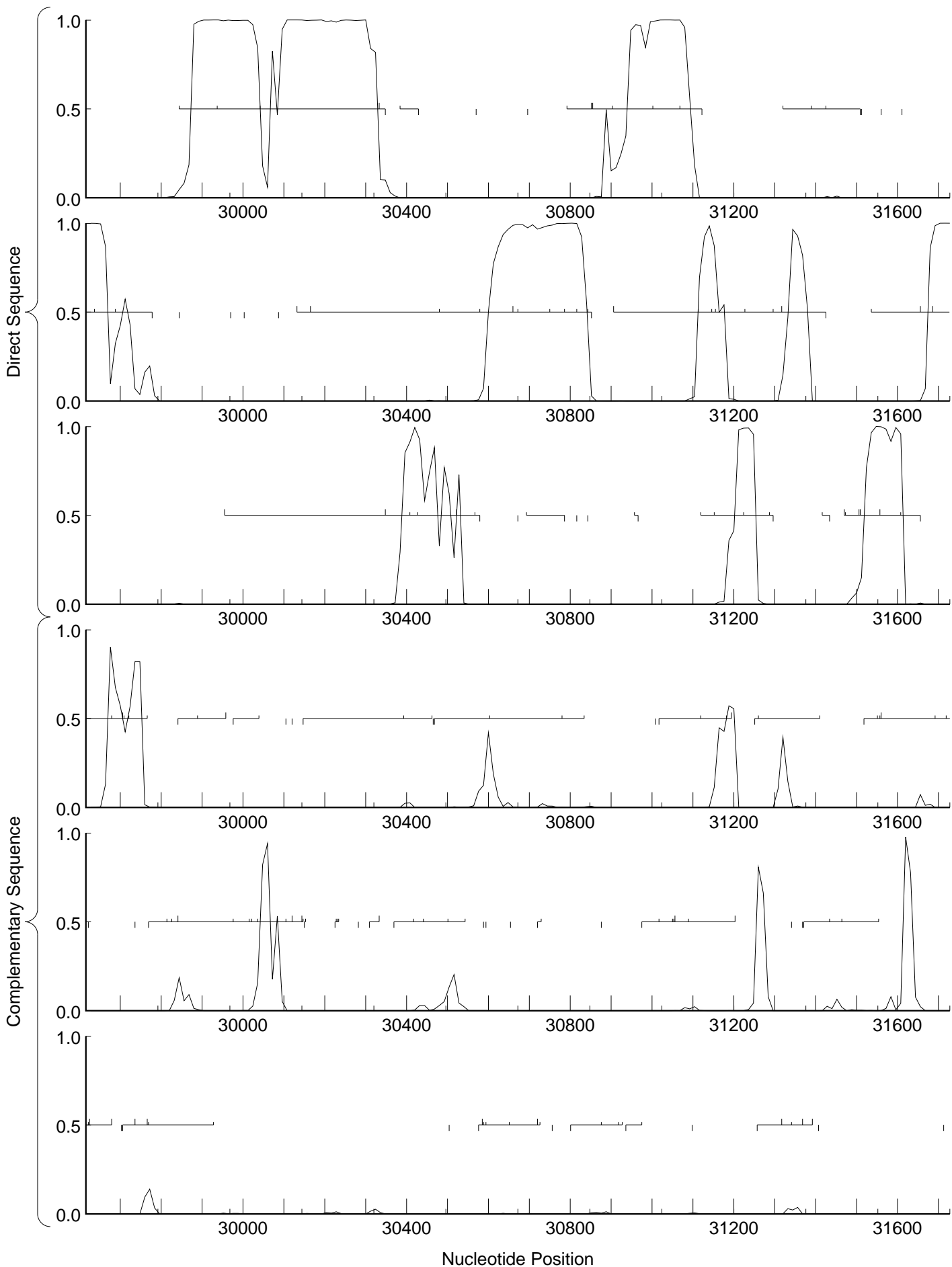


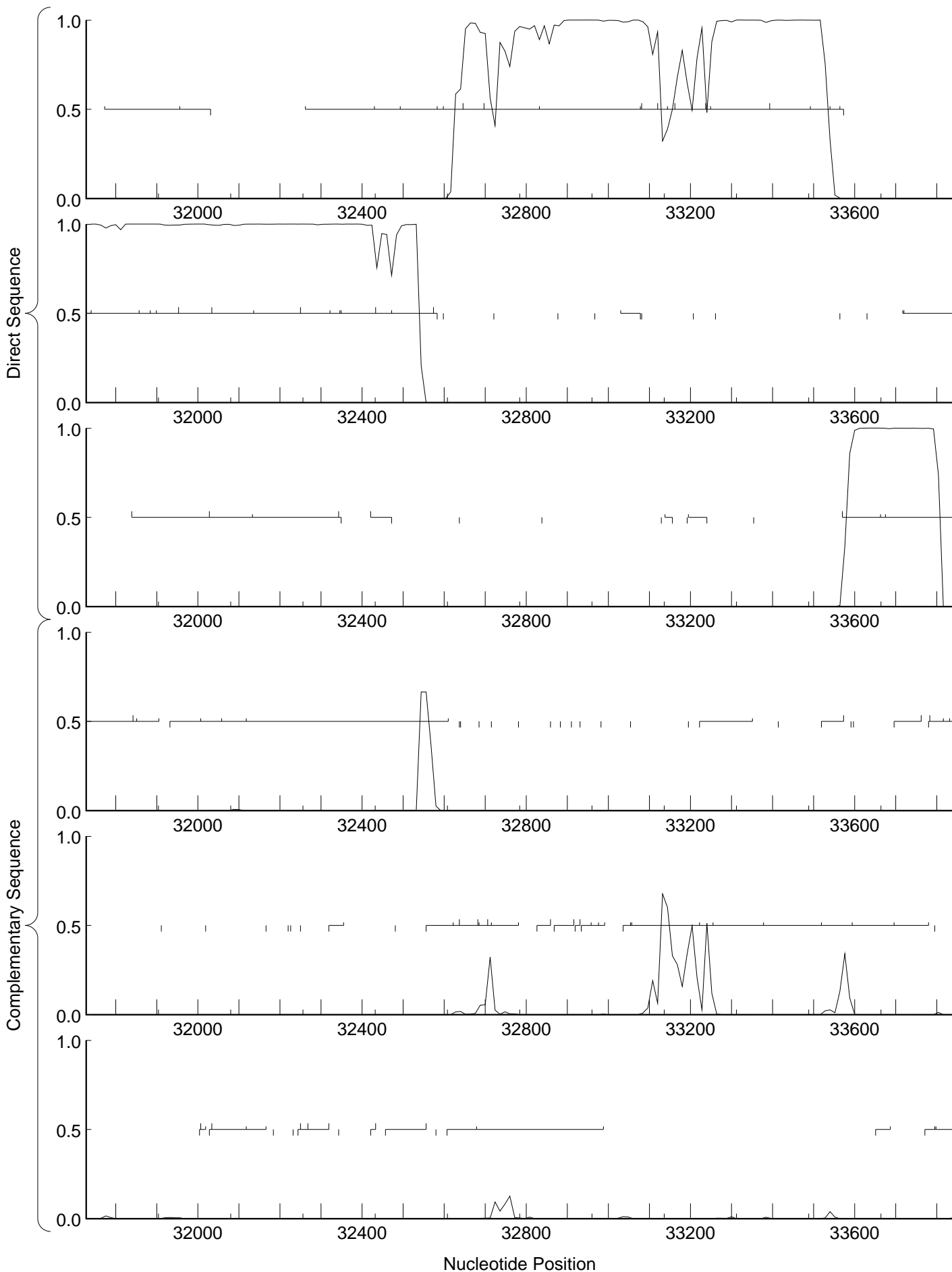
complete sequence, 47,612 bp including 12-base 3' overhang (CCCGCCCCCGA), Cluster P, Order 4, Window 96, Step 12, 13/24

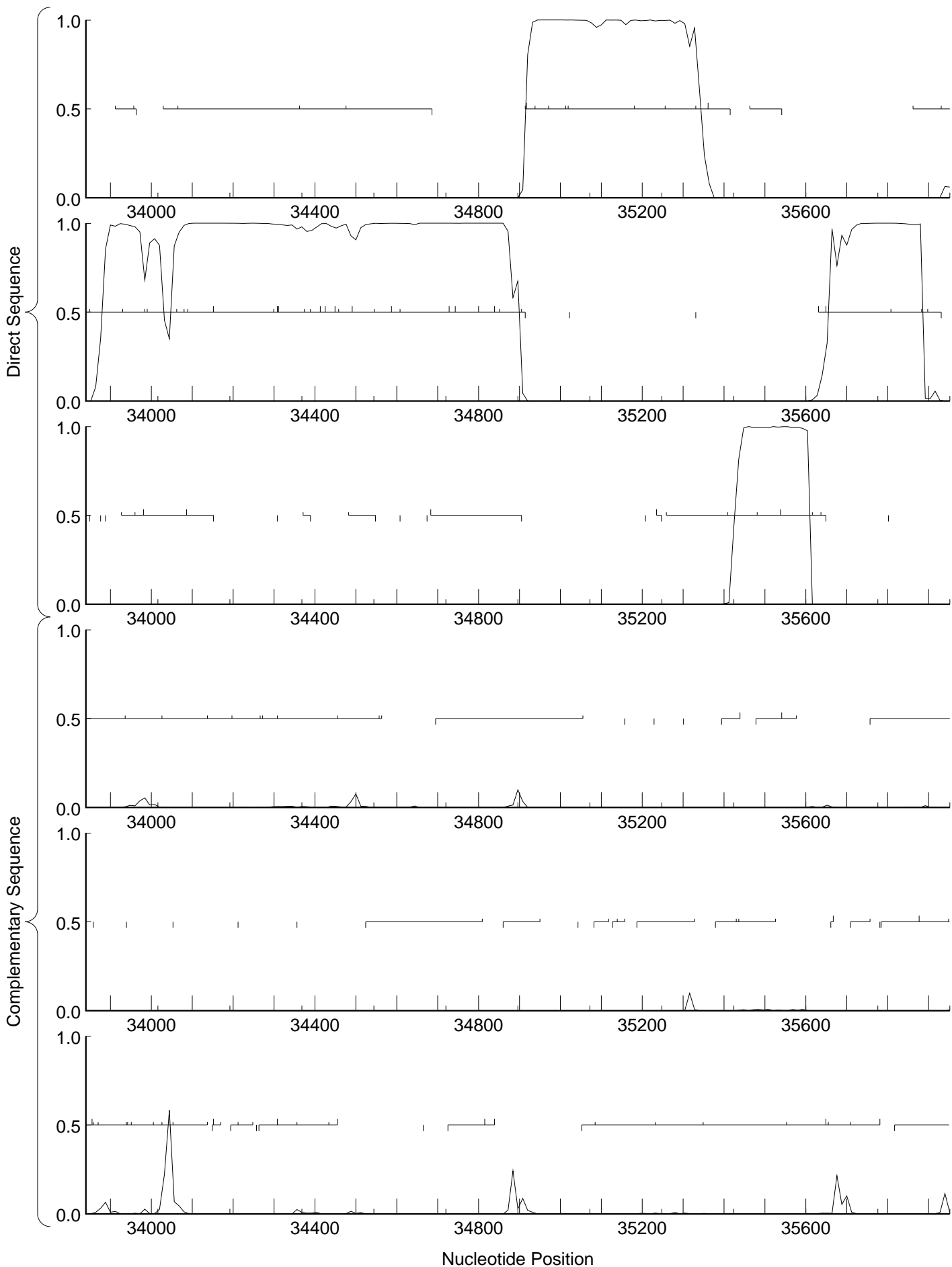


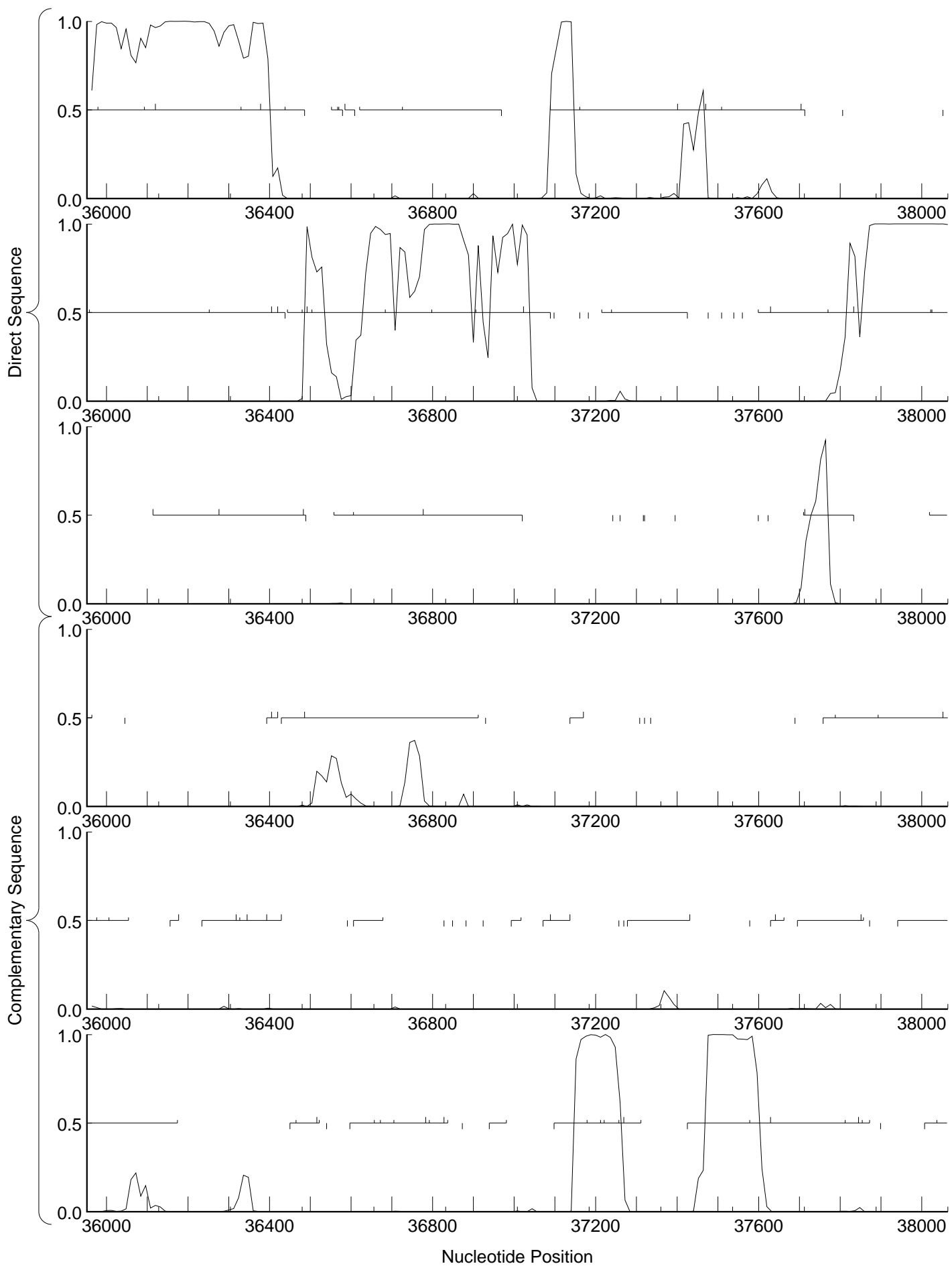


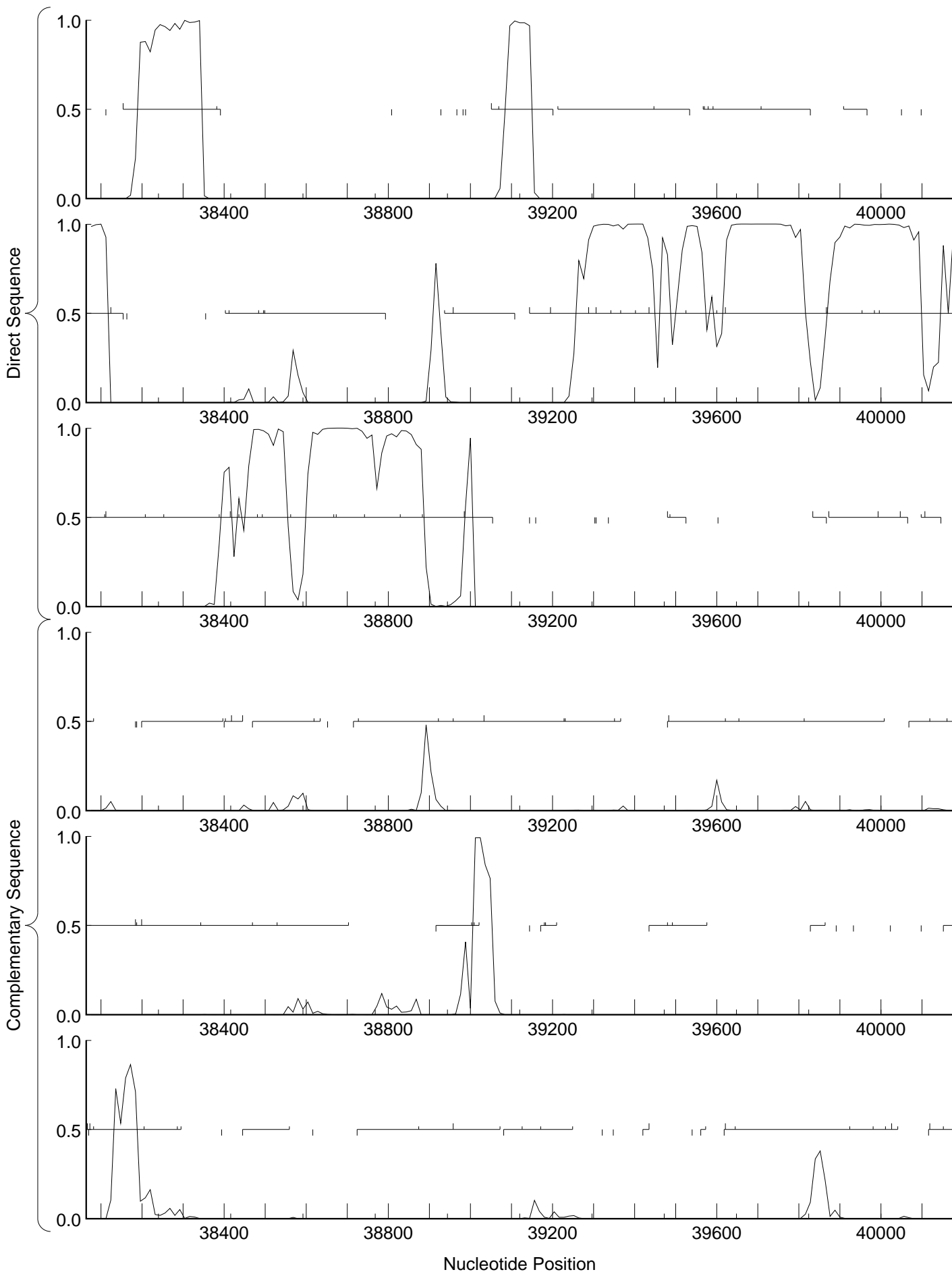


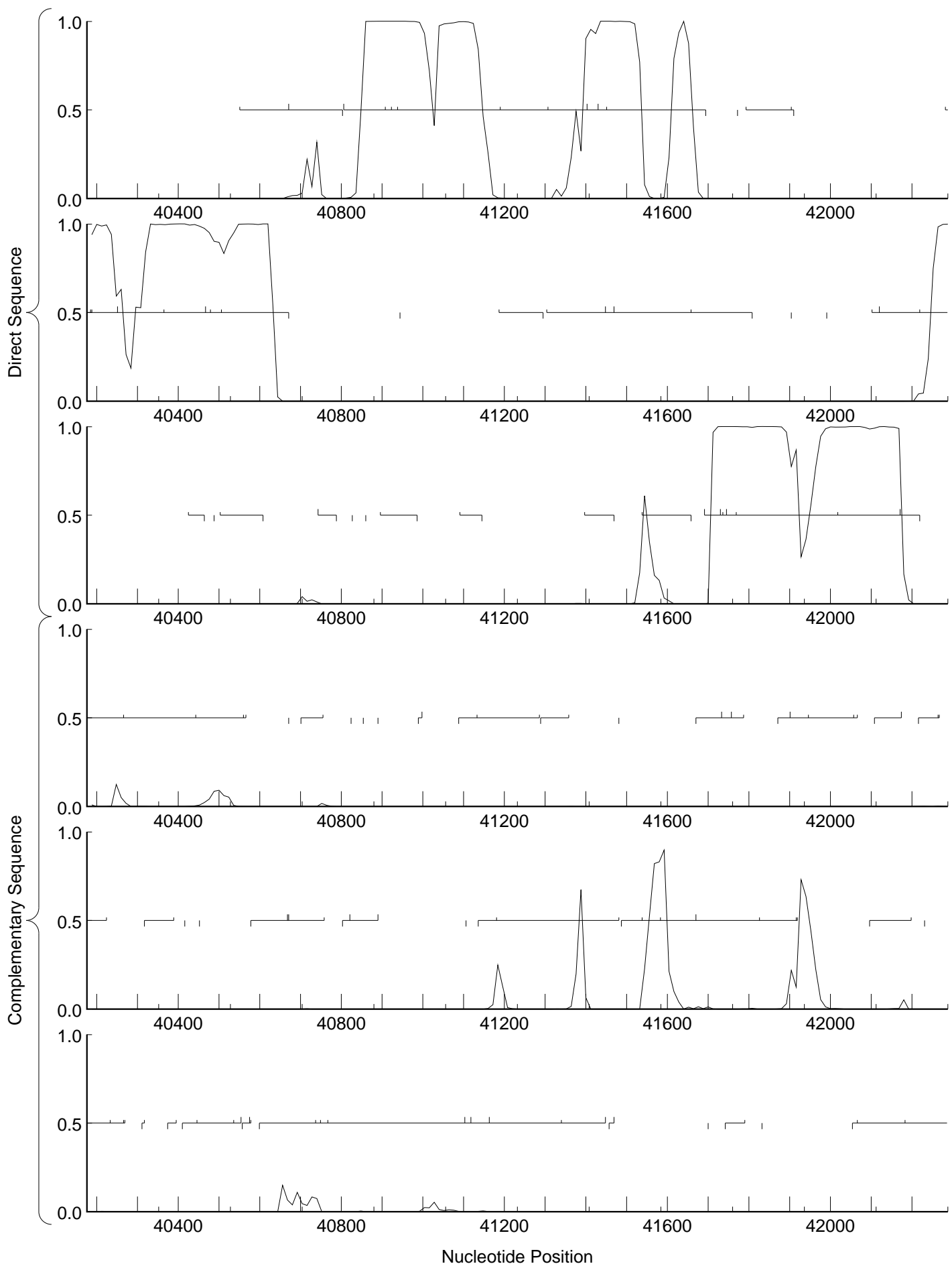


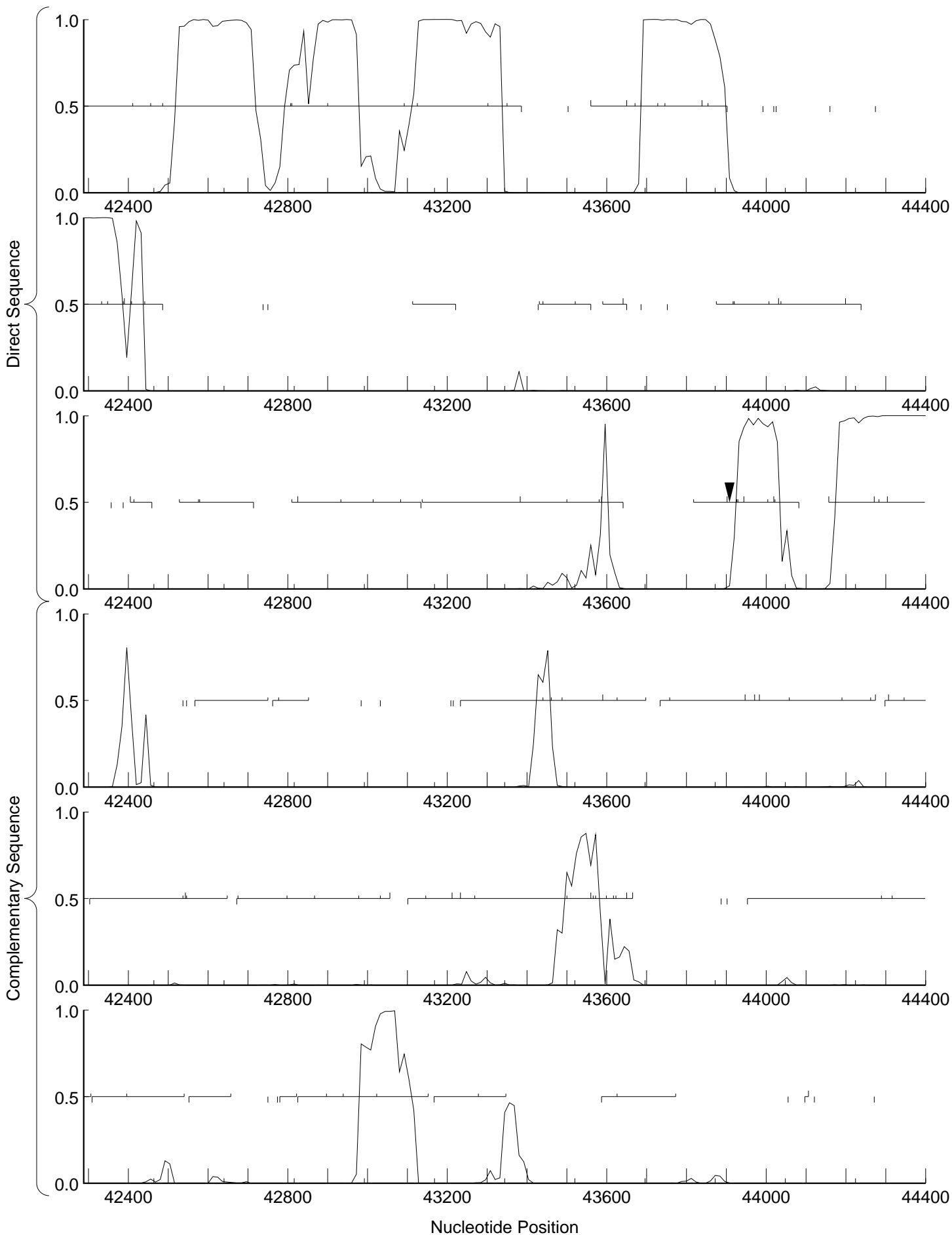












complete sequence, 47,612 bp including 12-base 3' overhang (CCCGCCCCCGA), Cluster P, Order 4, Window 96, Step 12, 23/24

