

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Camster complete sequence, 47149 bp including 12-base 3' overhang (CTG
Analysis Date : 8/5/21 at 3:36:49
Pages : 24
Sequence Length : 47149 bp
GC Content : 67.23%

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,

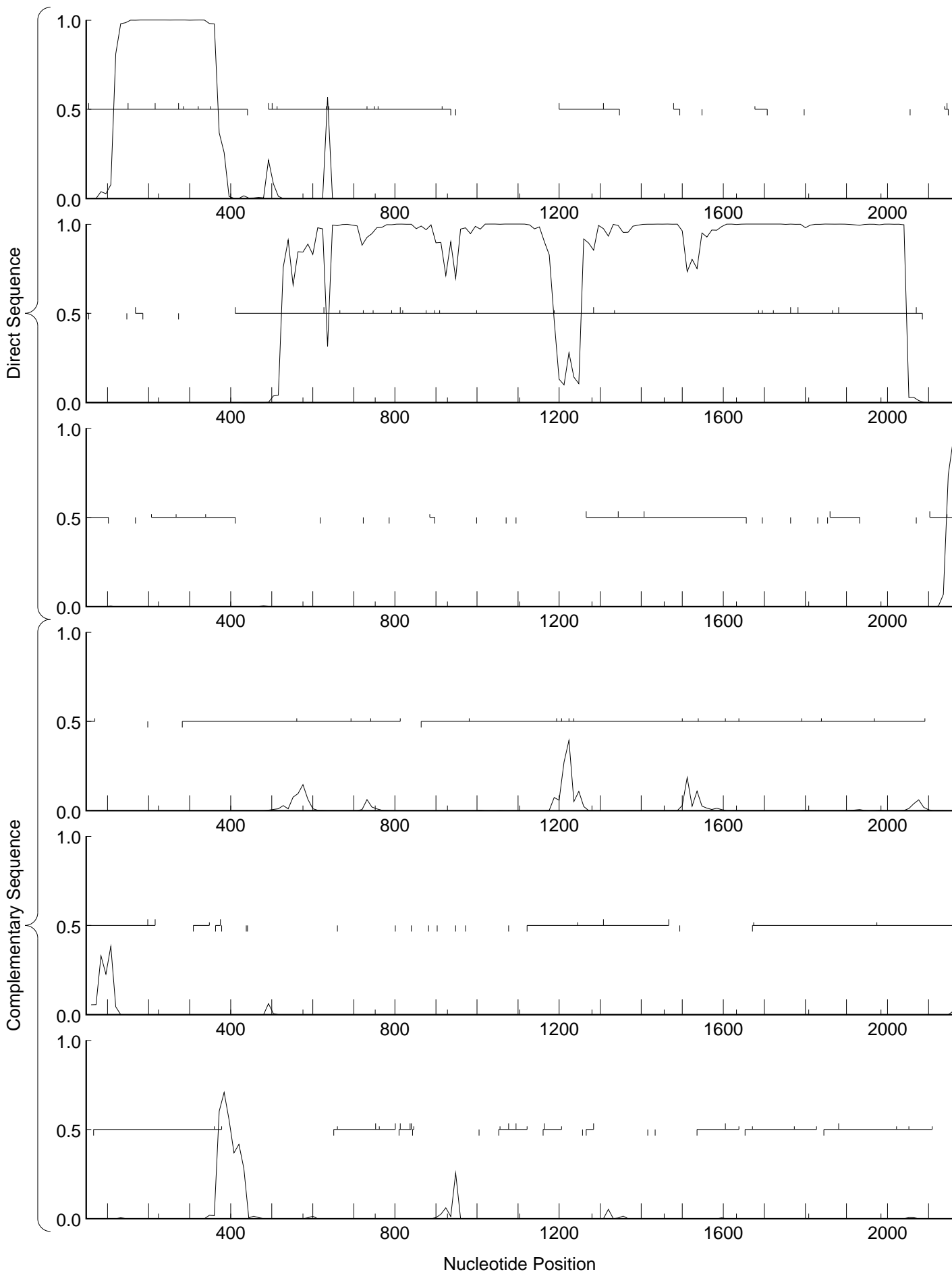
MATRIX INFORMATION

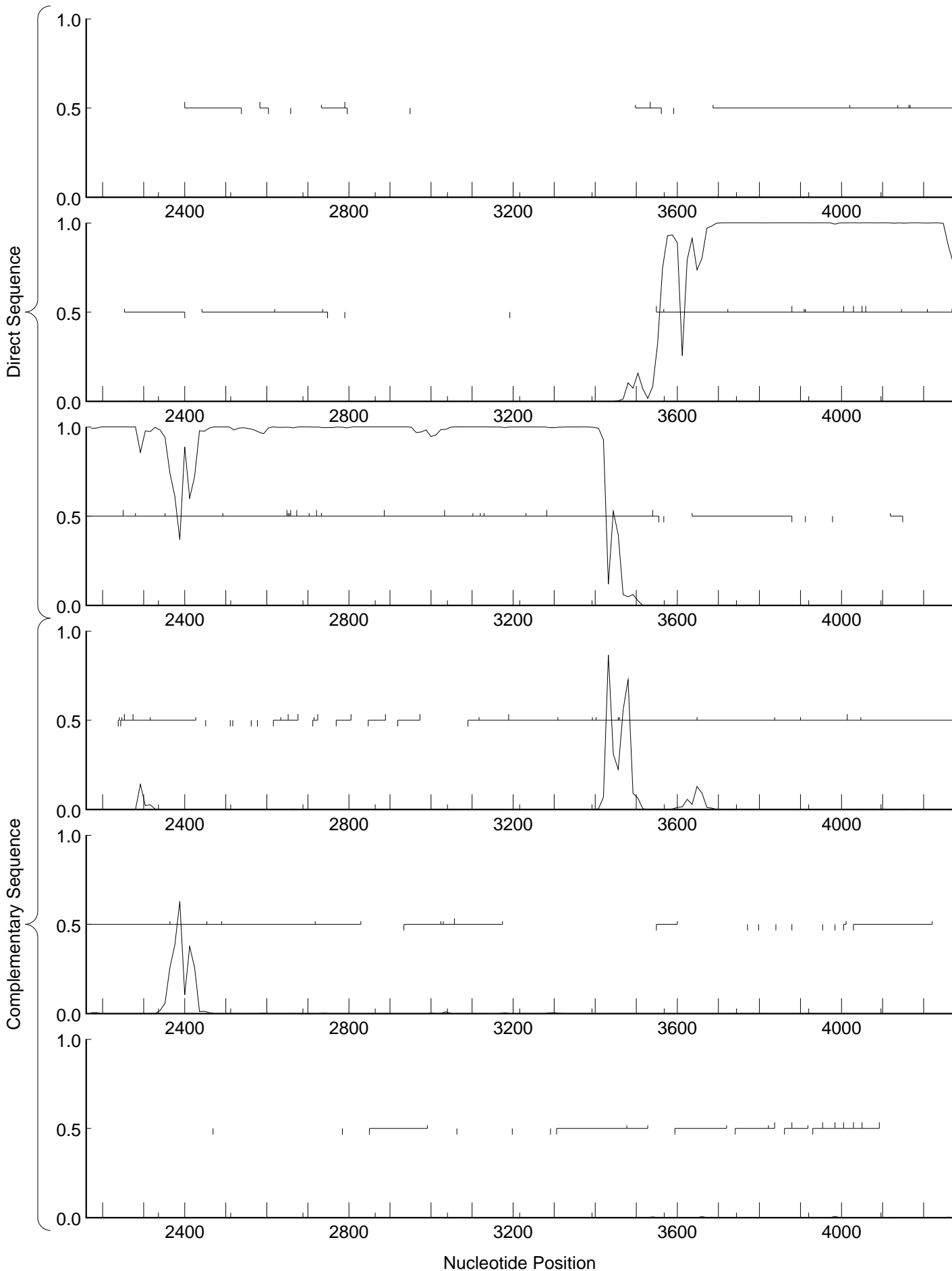
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

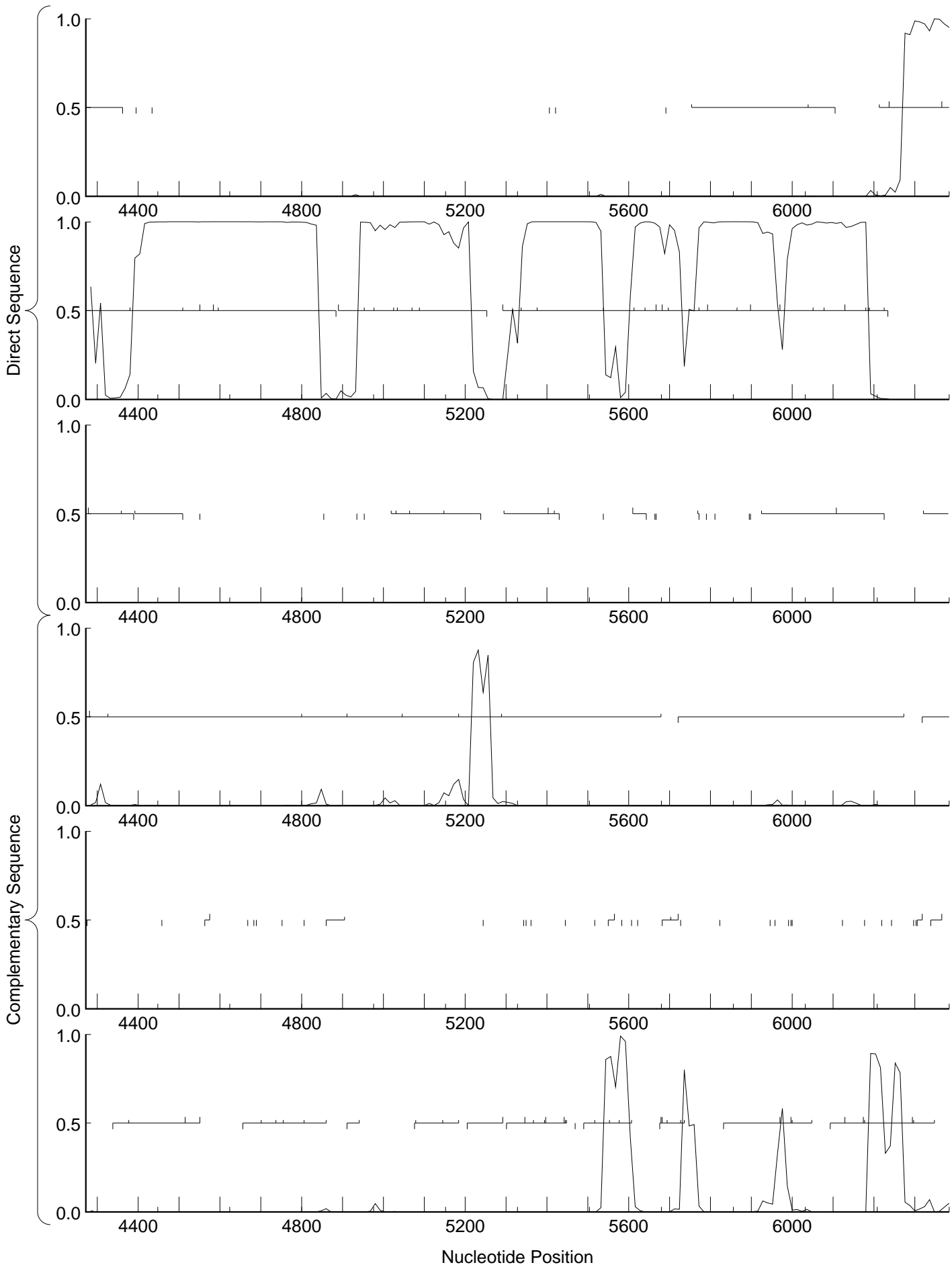
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

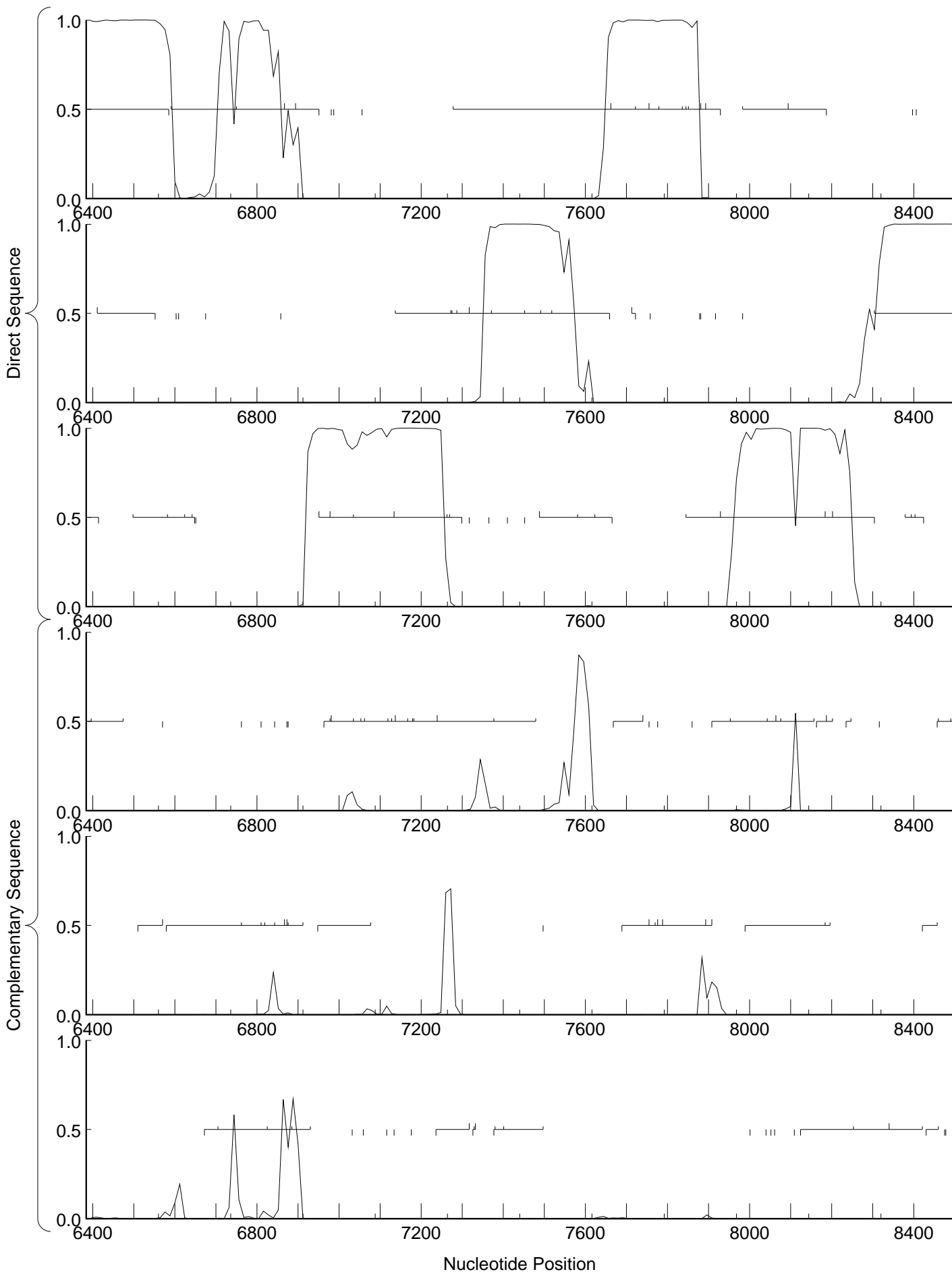
Matrix notes & comments

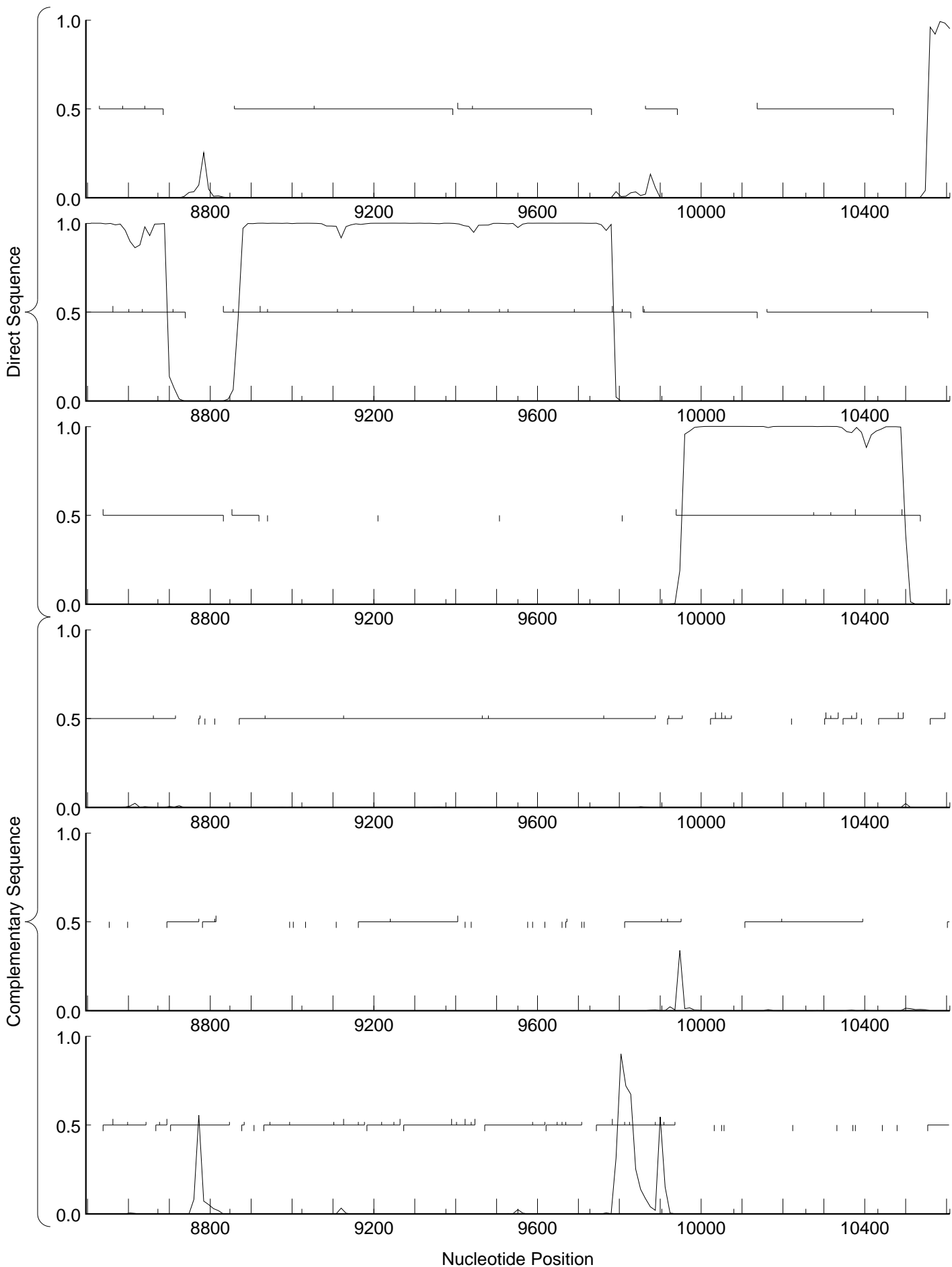
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

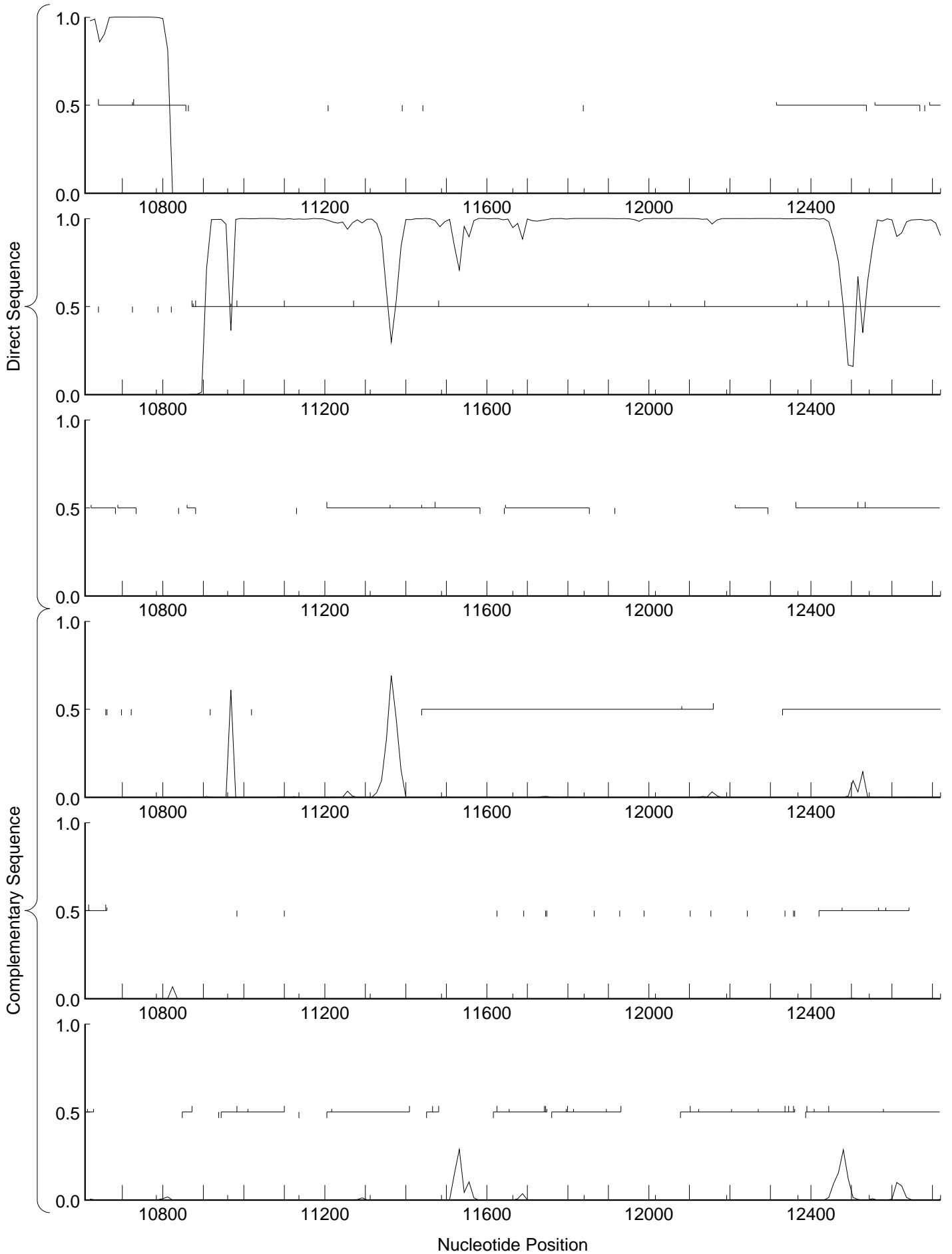


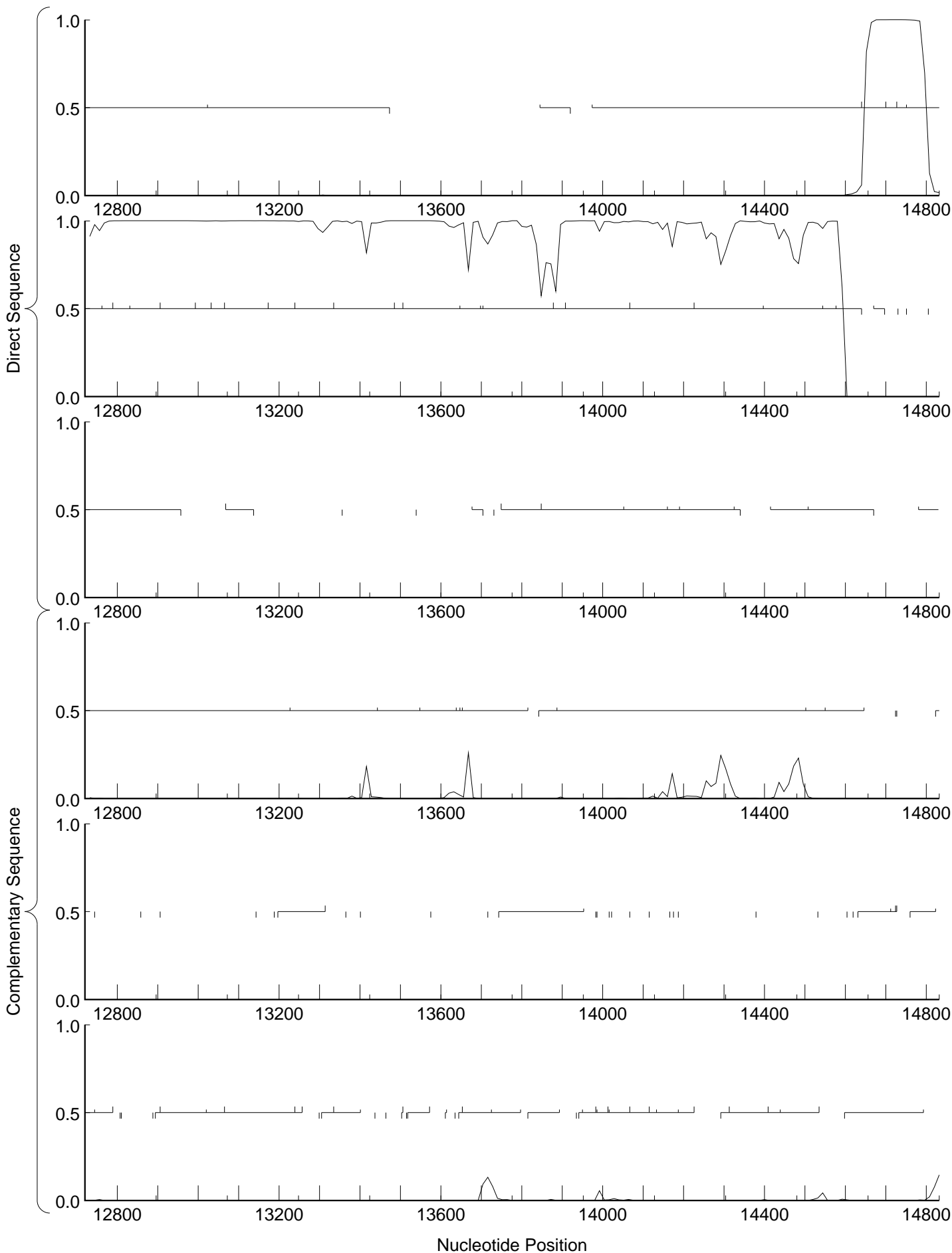


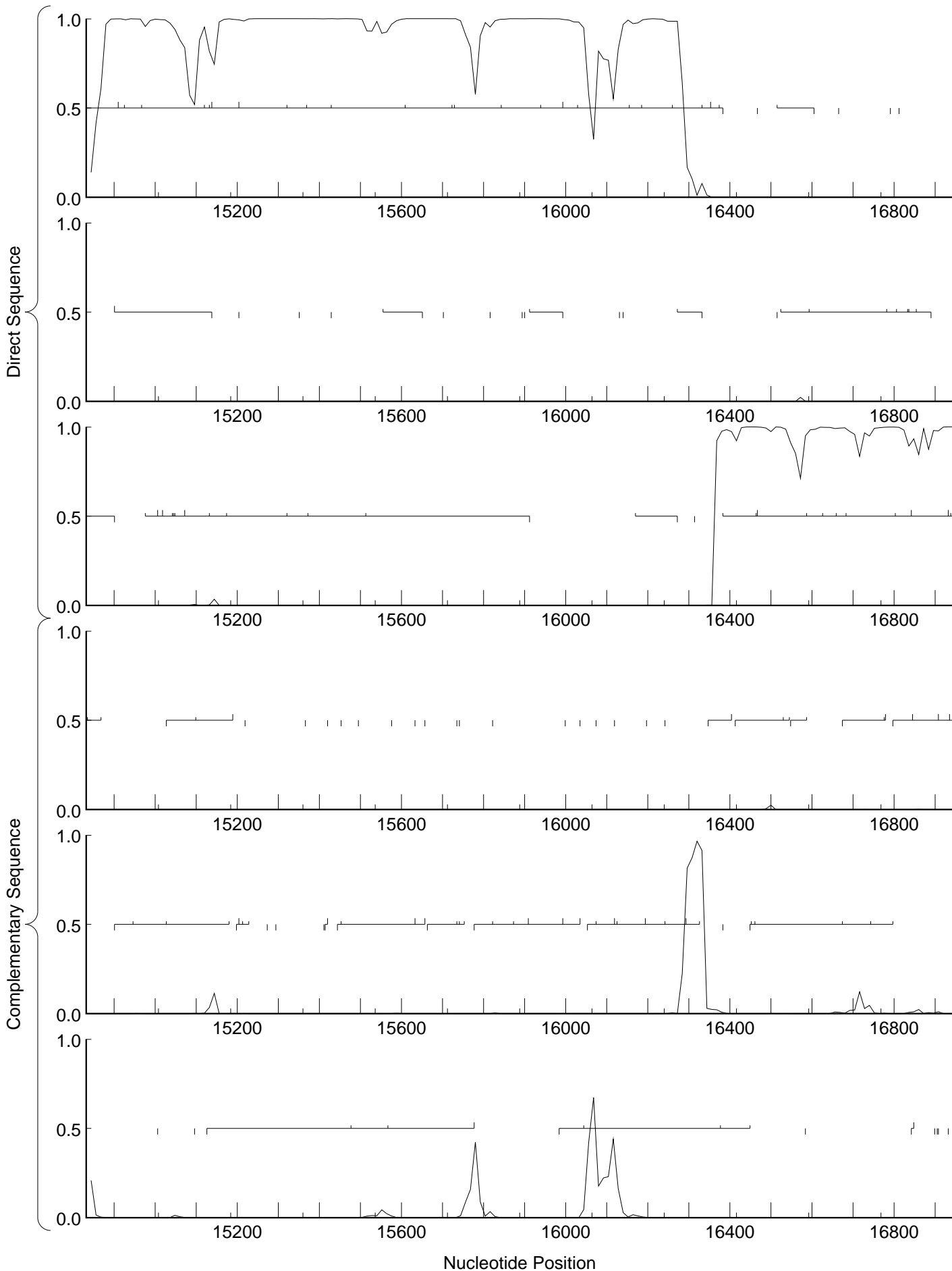


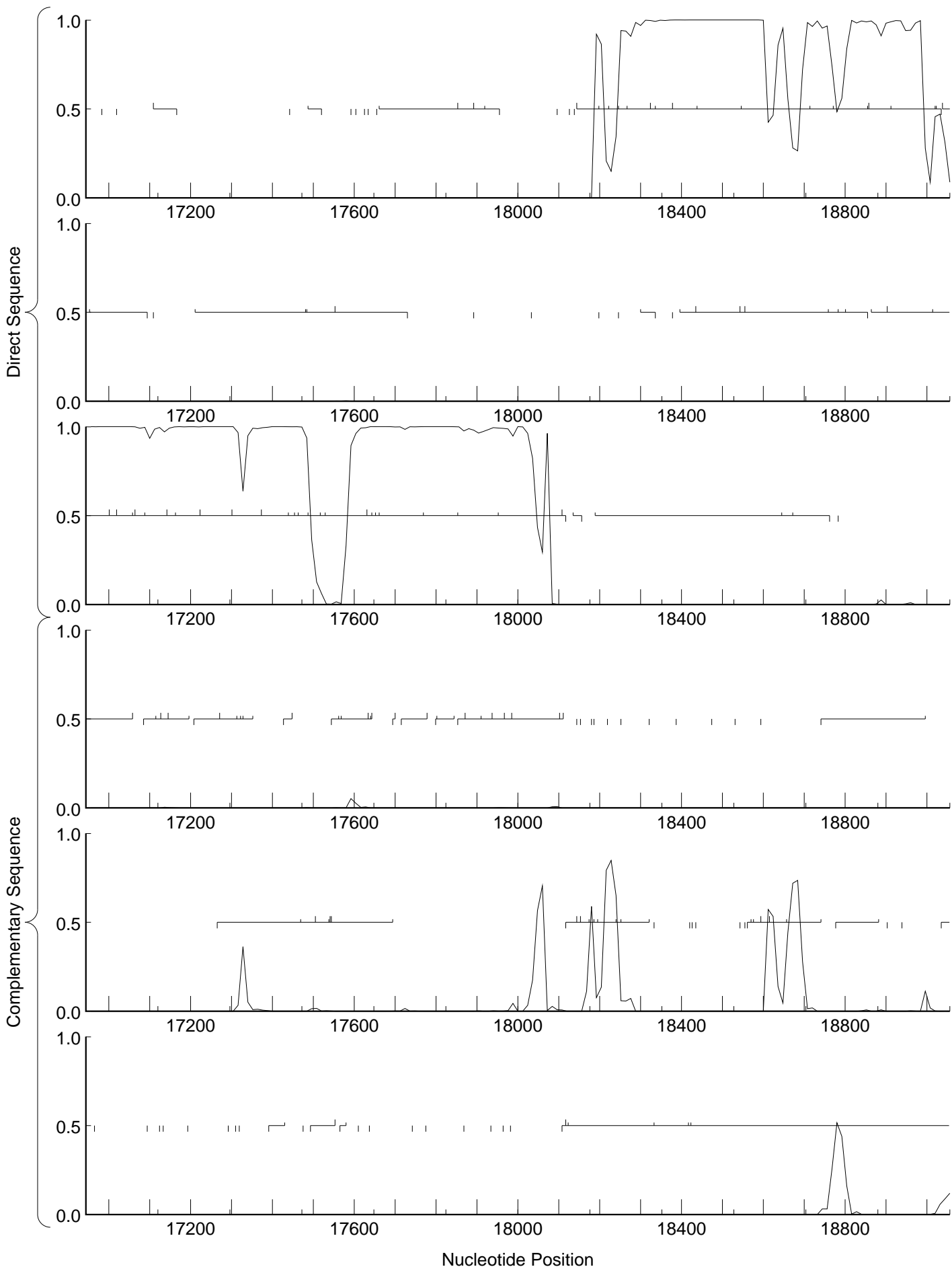




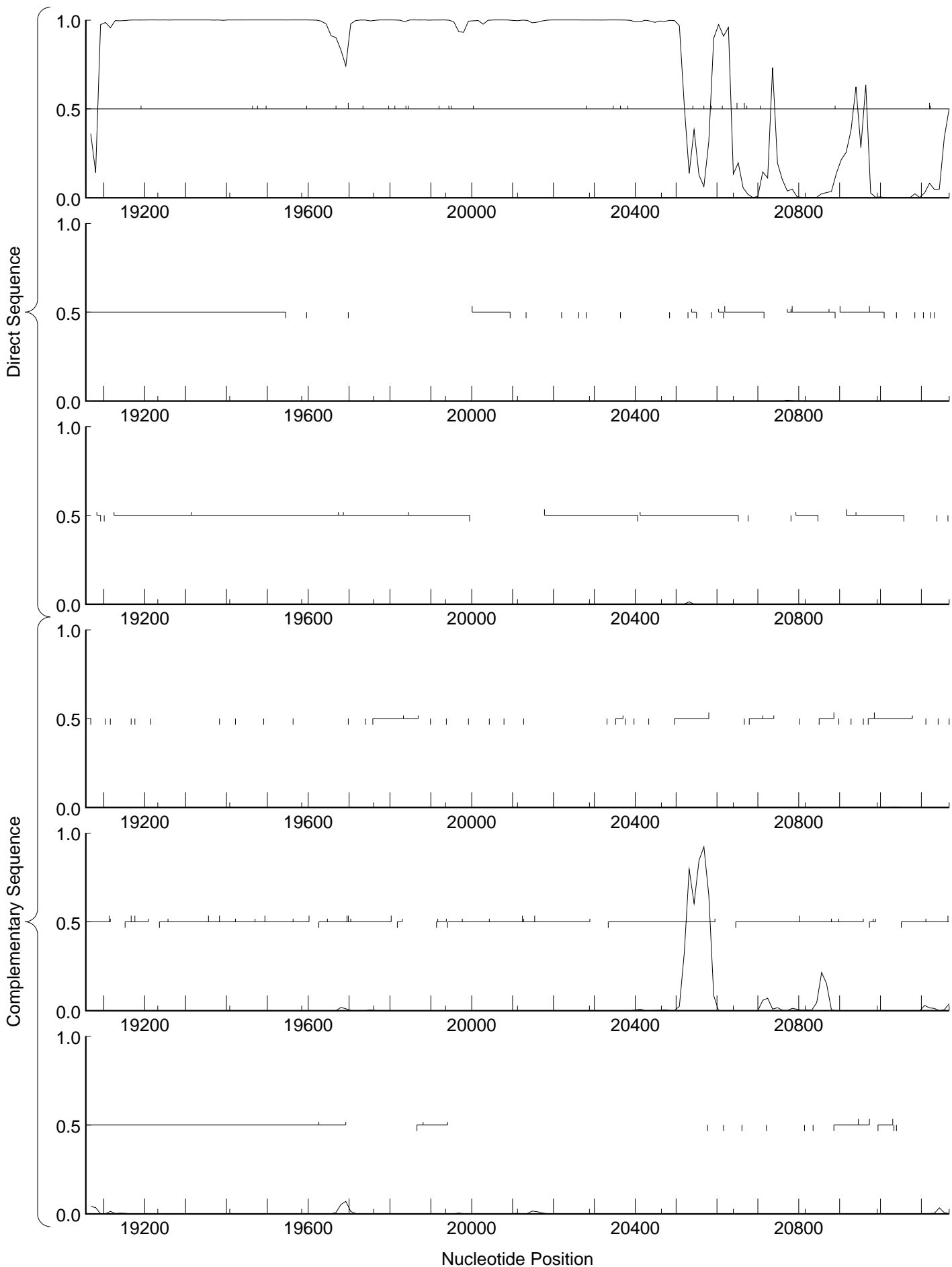


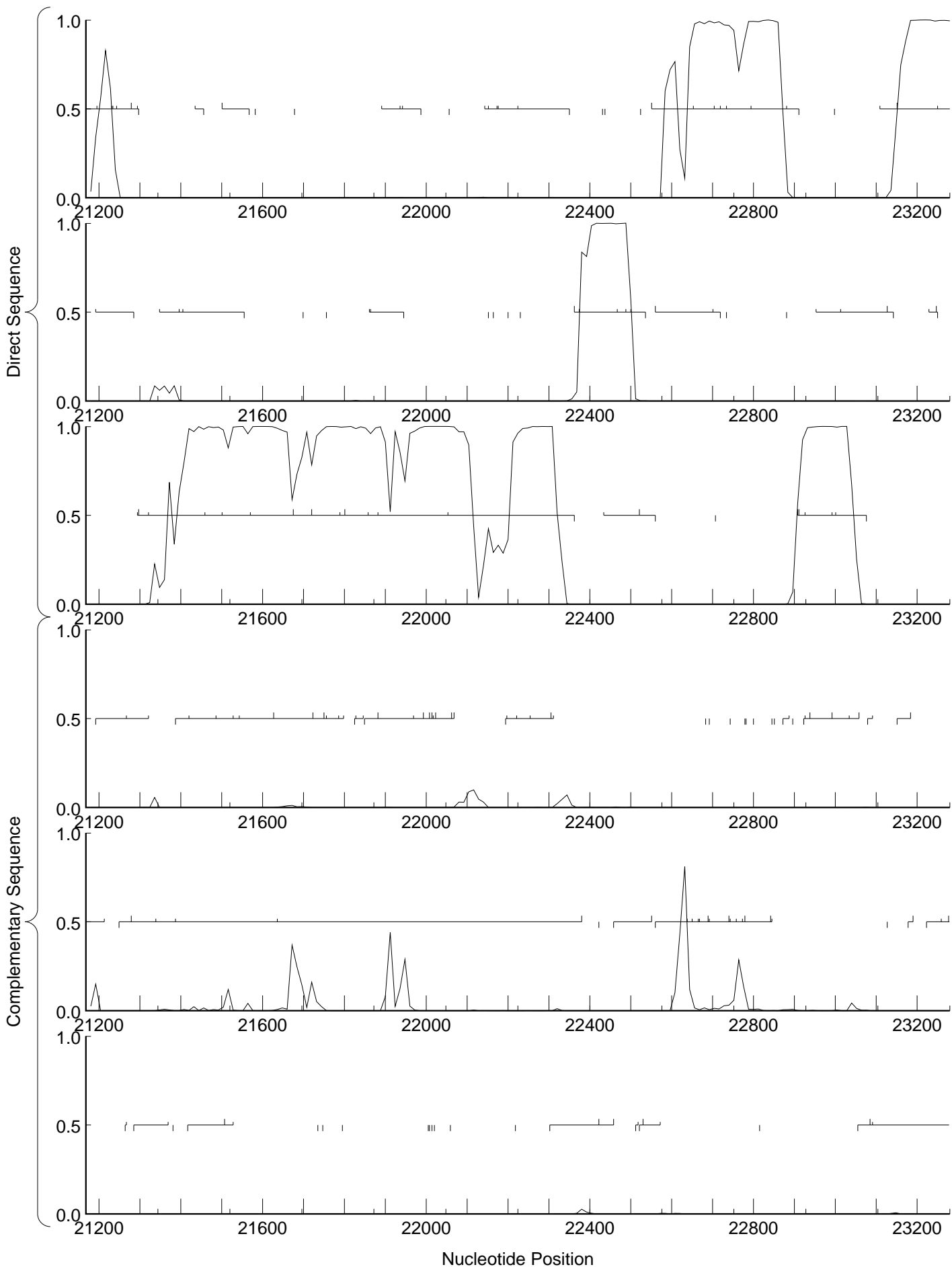




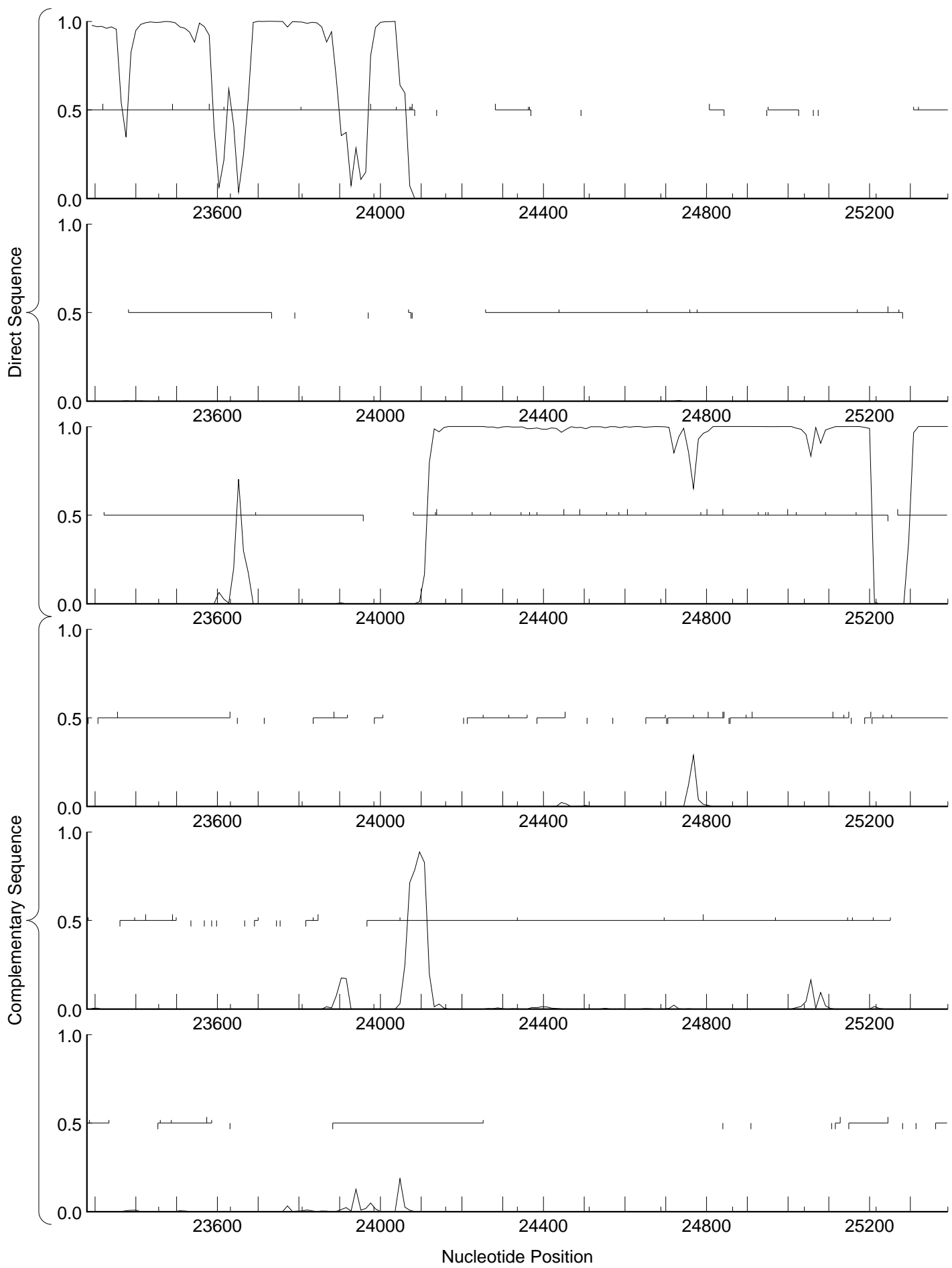


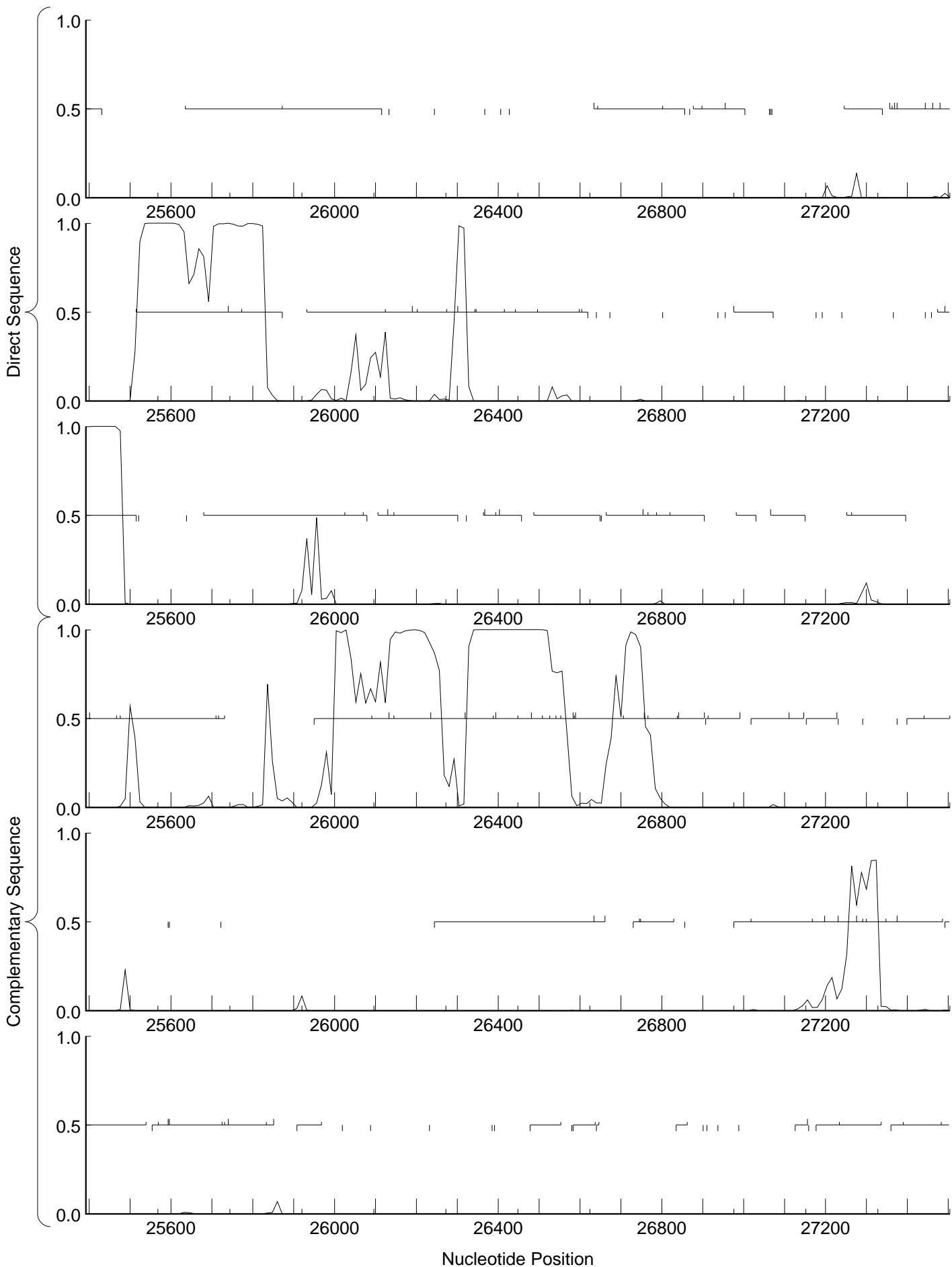
Complete sequence, 47149 bp including 12-base 3' overhang (CCTGCCGCCGA), Cluster P1, Order 4, Window 96, Step 12, 11/24

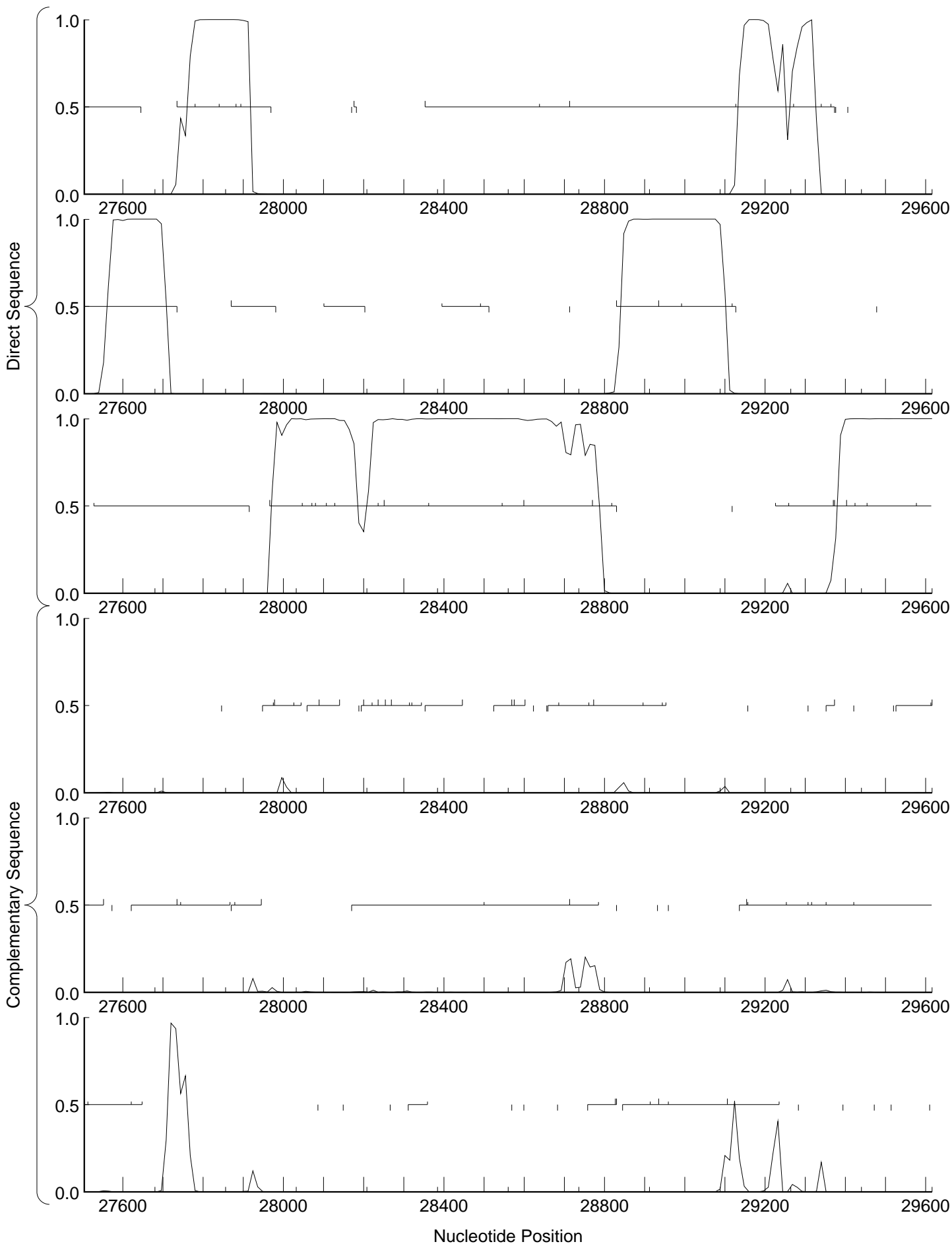




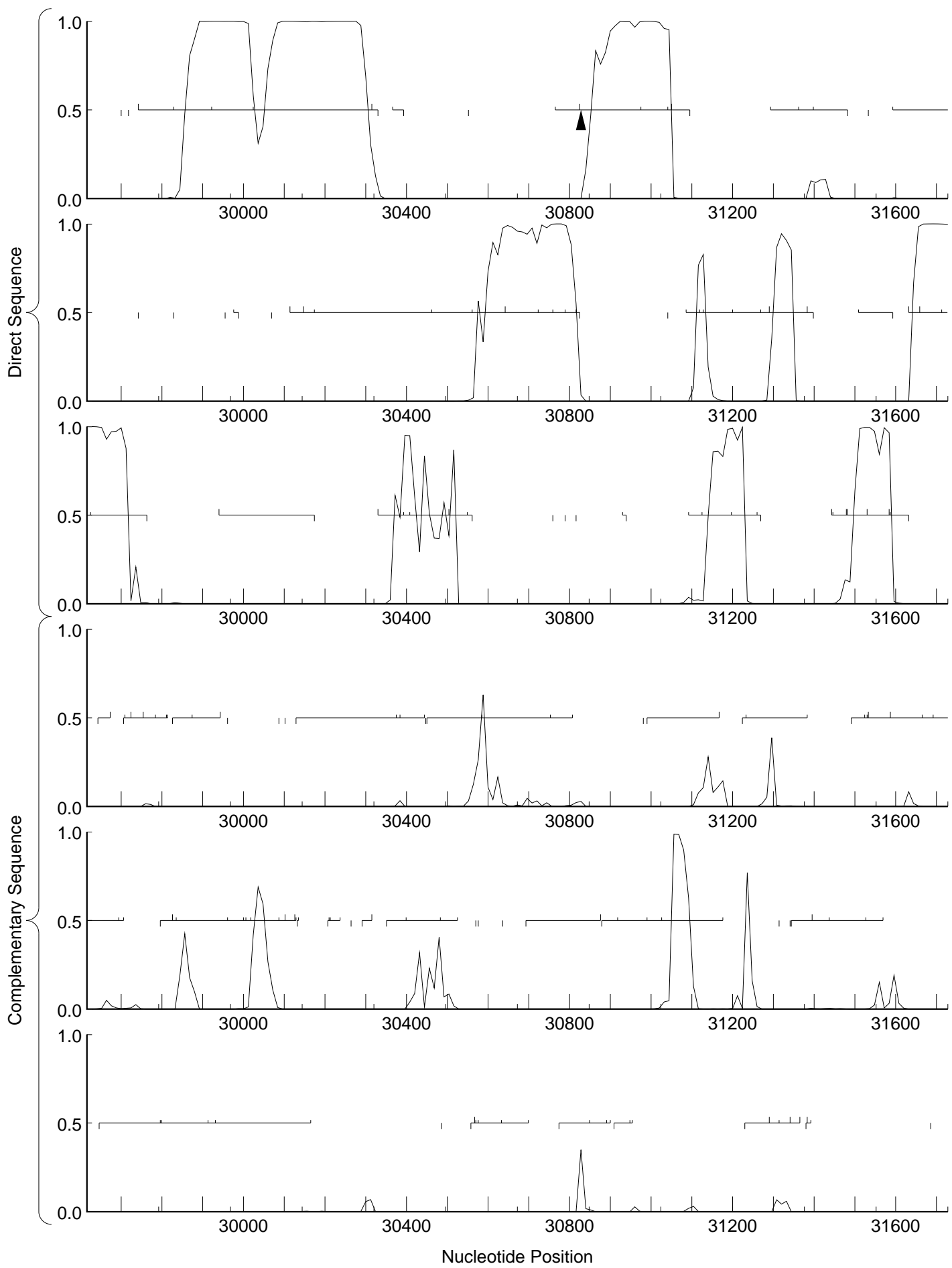
Complete sequence, 47149 bp including 12-base 3' overhang (CCTGCCGCCGA), Cluster P1, Order 4, Window 96, Step 12, 13/24







Complete sequence, 47149 bp including 12-base 3' overhang (CCTGCCGCCGA), Cluster P1, Order 4, Window 96, Step 12, 16/24



Complete sequence, 47149 bp including 12-base 3' overhang (CCTGCCGCCCGA), Cluster P1, Order 4, Window 96, Step 12, 17/24

