

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Clark complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA
Analysis Date : 3/17/18 at 16:22:43
Pages : 24
Sequence Length : 47716 bp
GC Content : 66.53%

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,

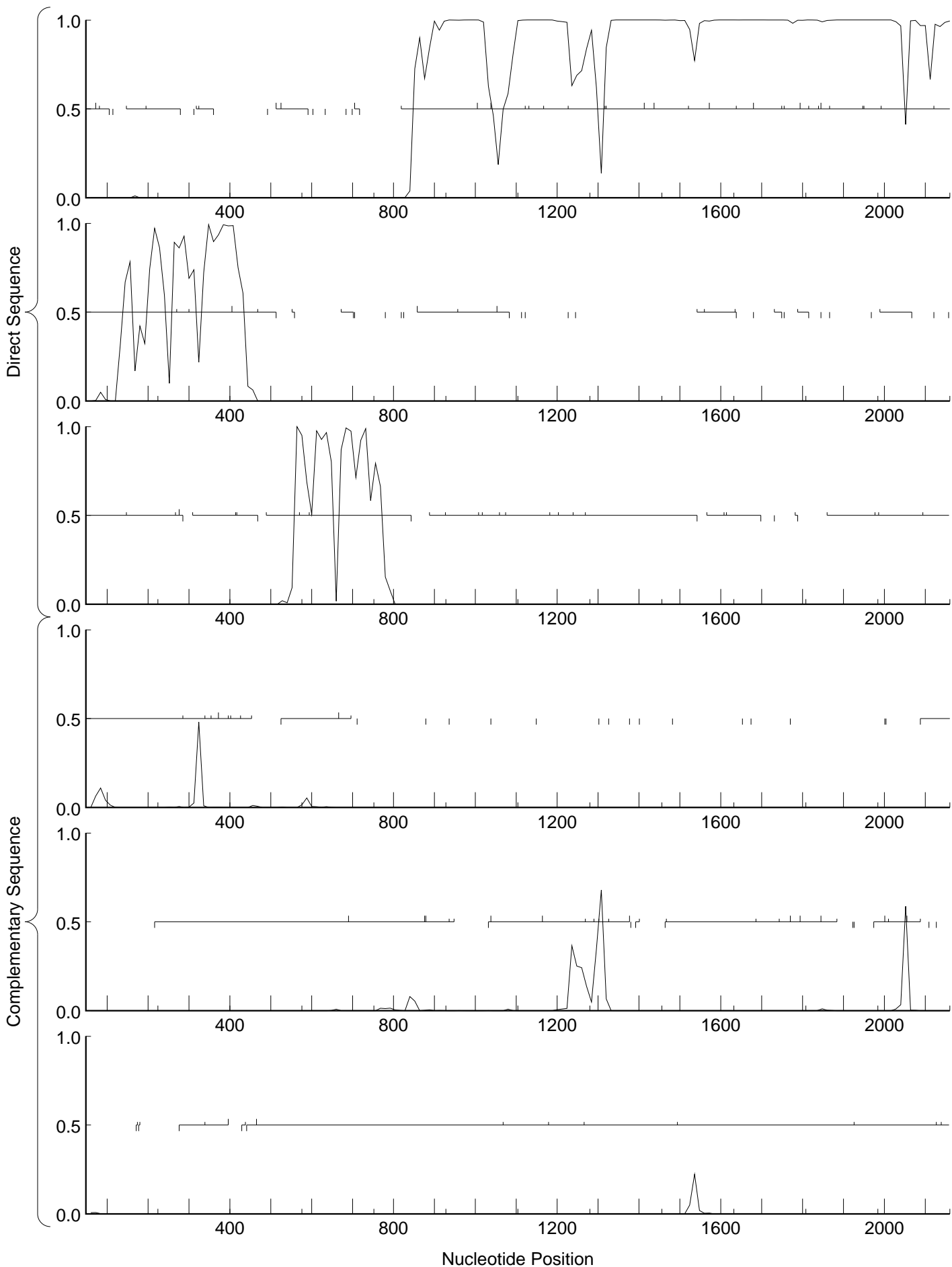
Matrix notes & comments

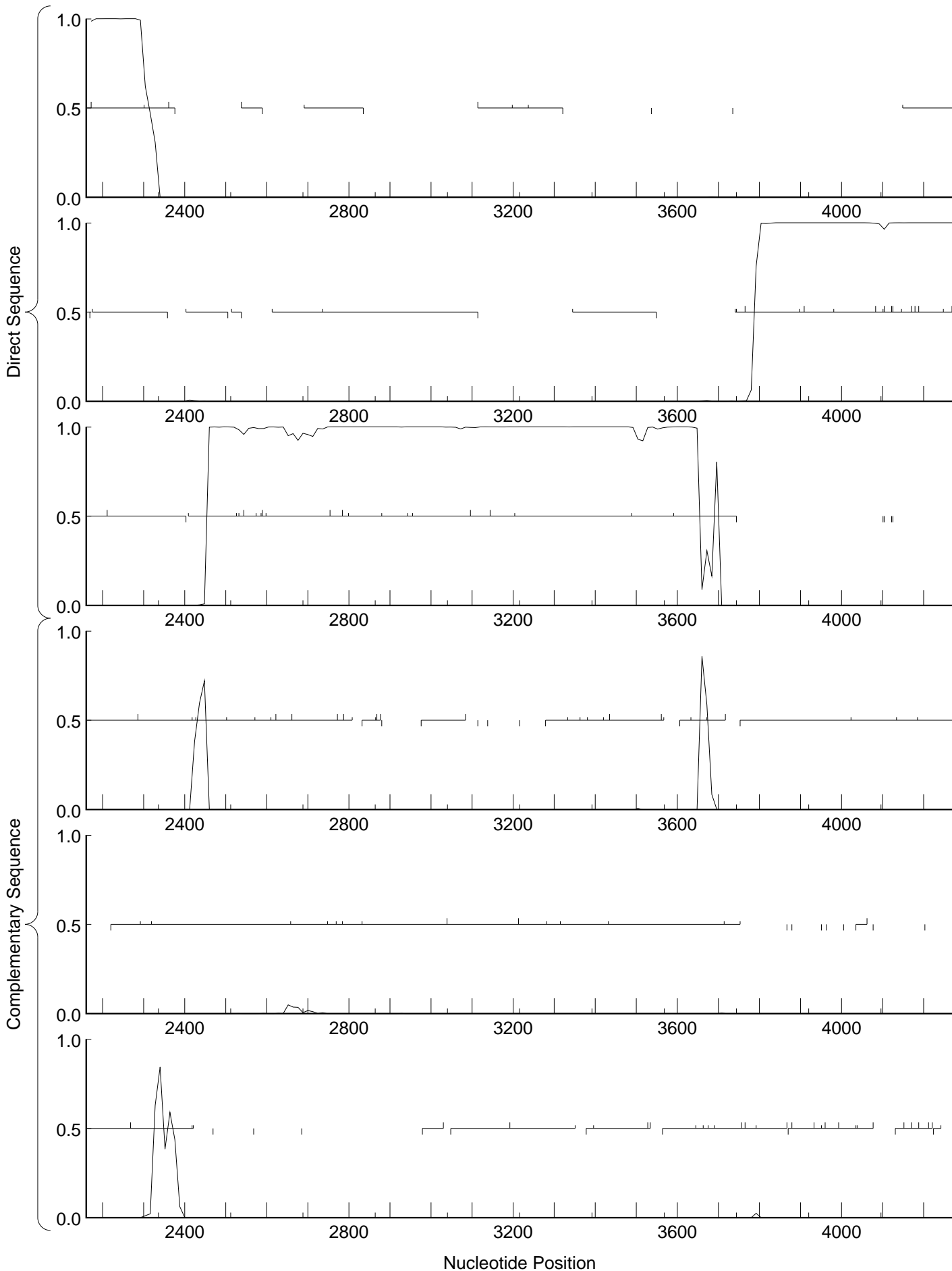
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:23:03 2014

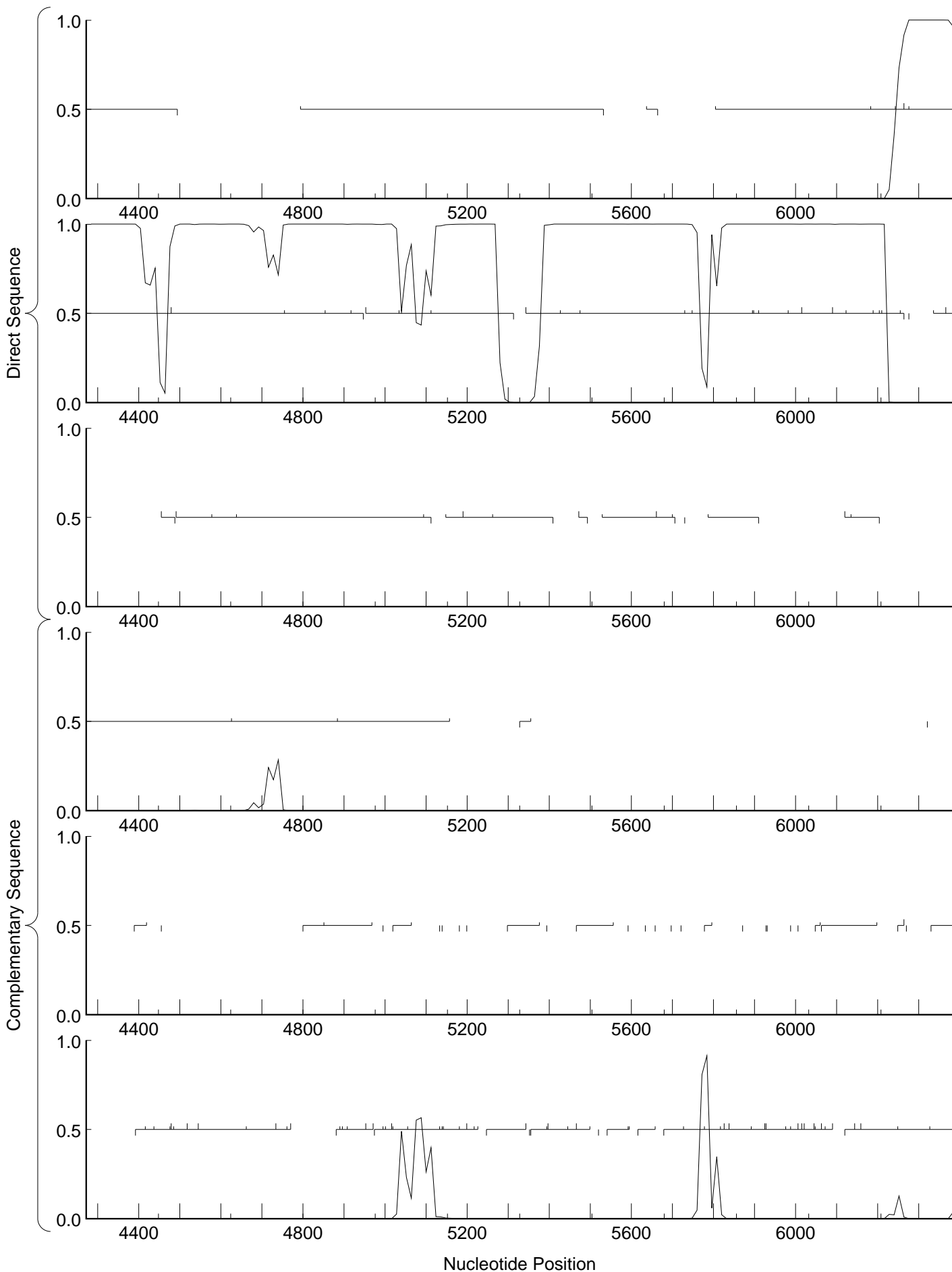
MATRIX INFORMATION

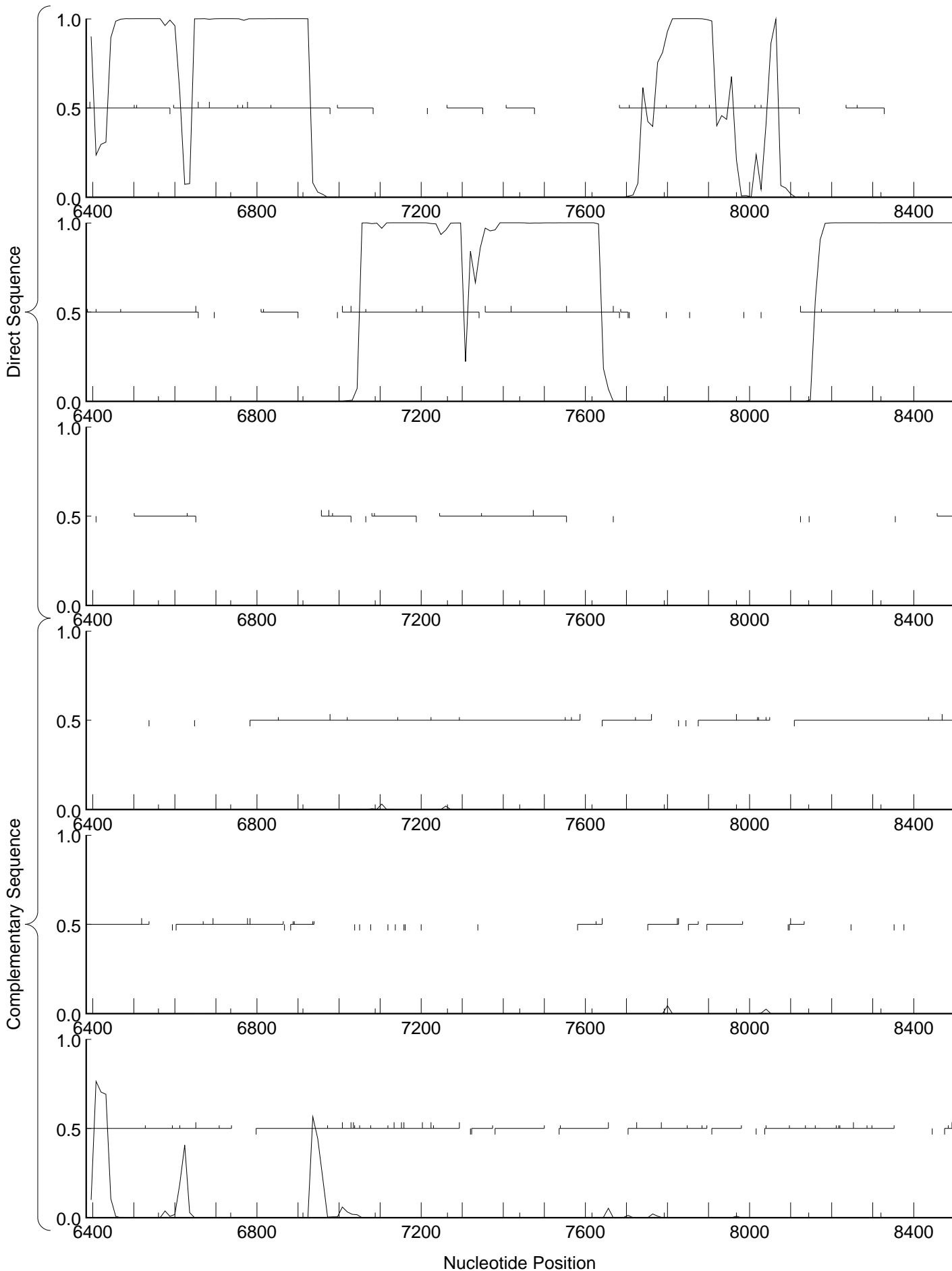
Matrix : Mycobacterium_smegmatis_MC2_155
Author : -
Order : 4

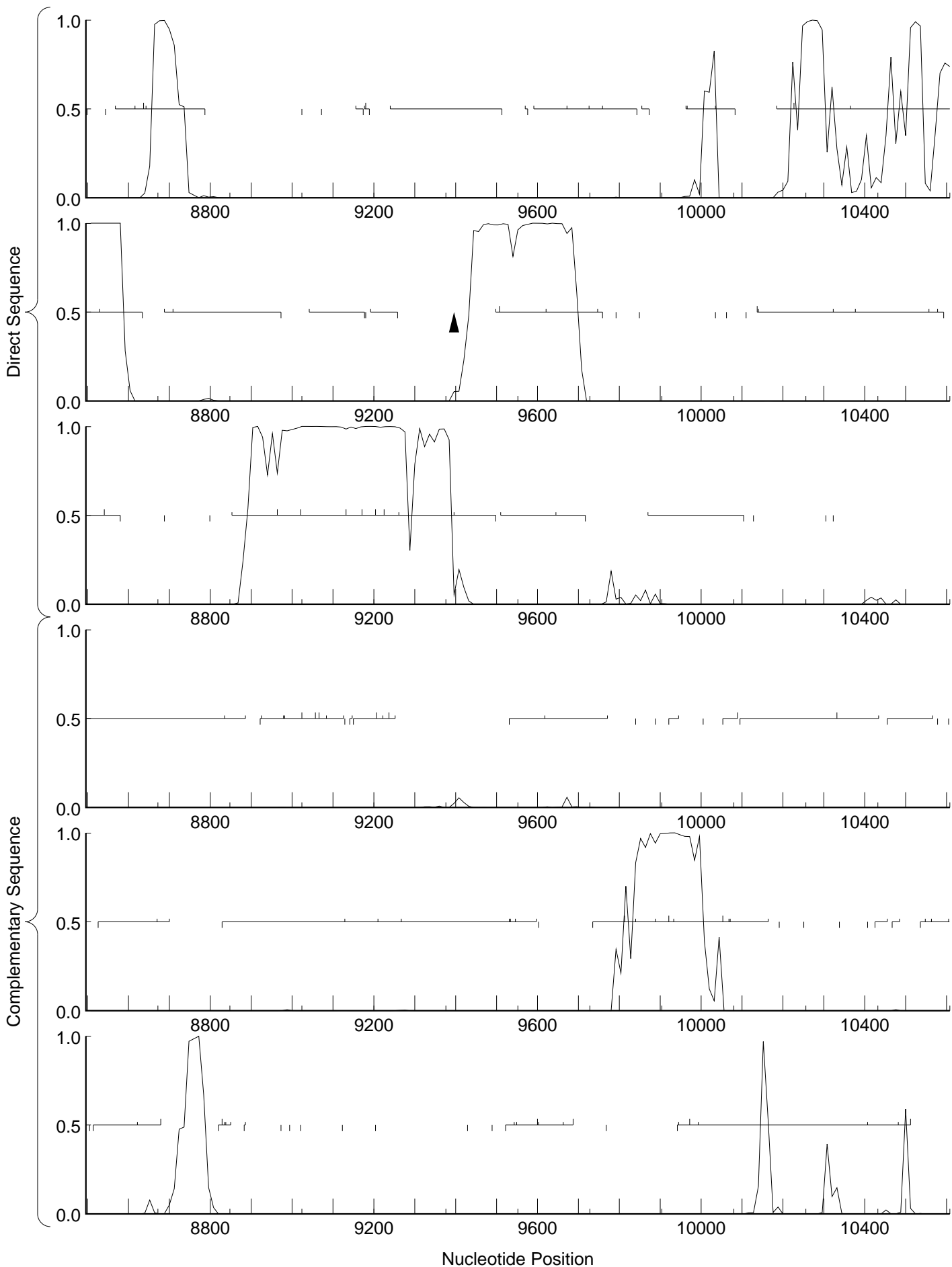
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230



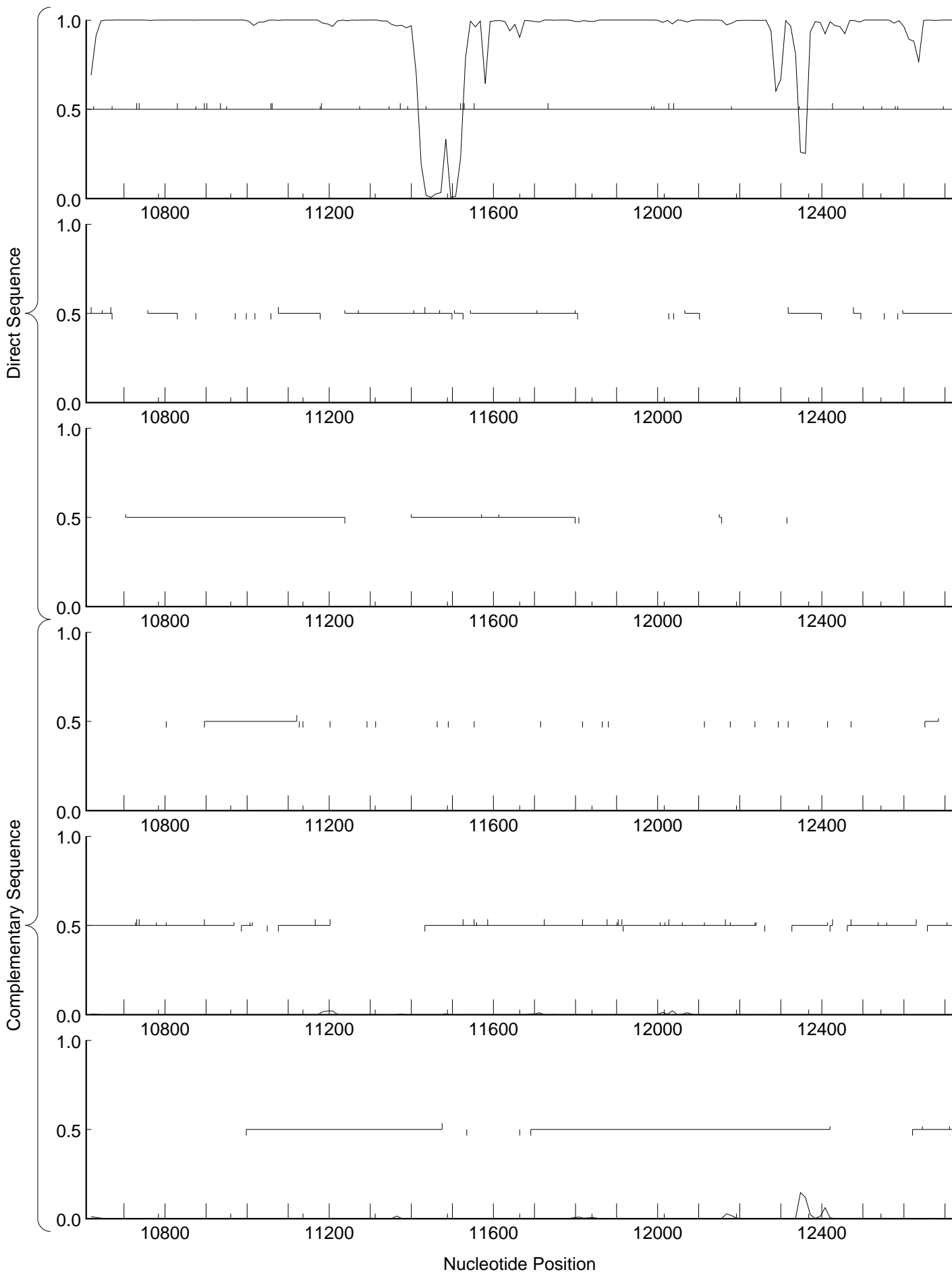




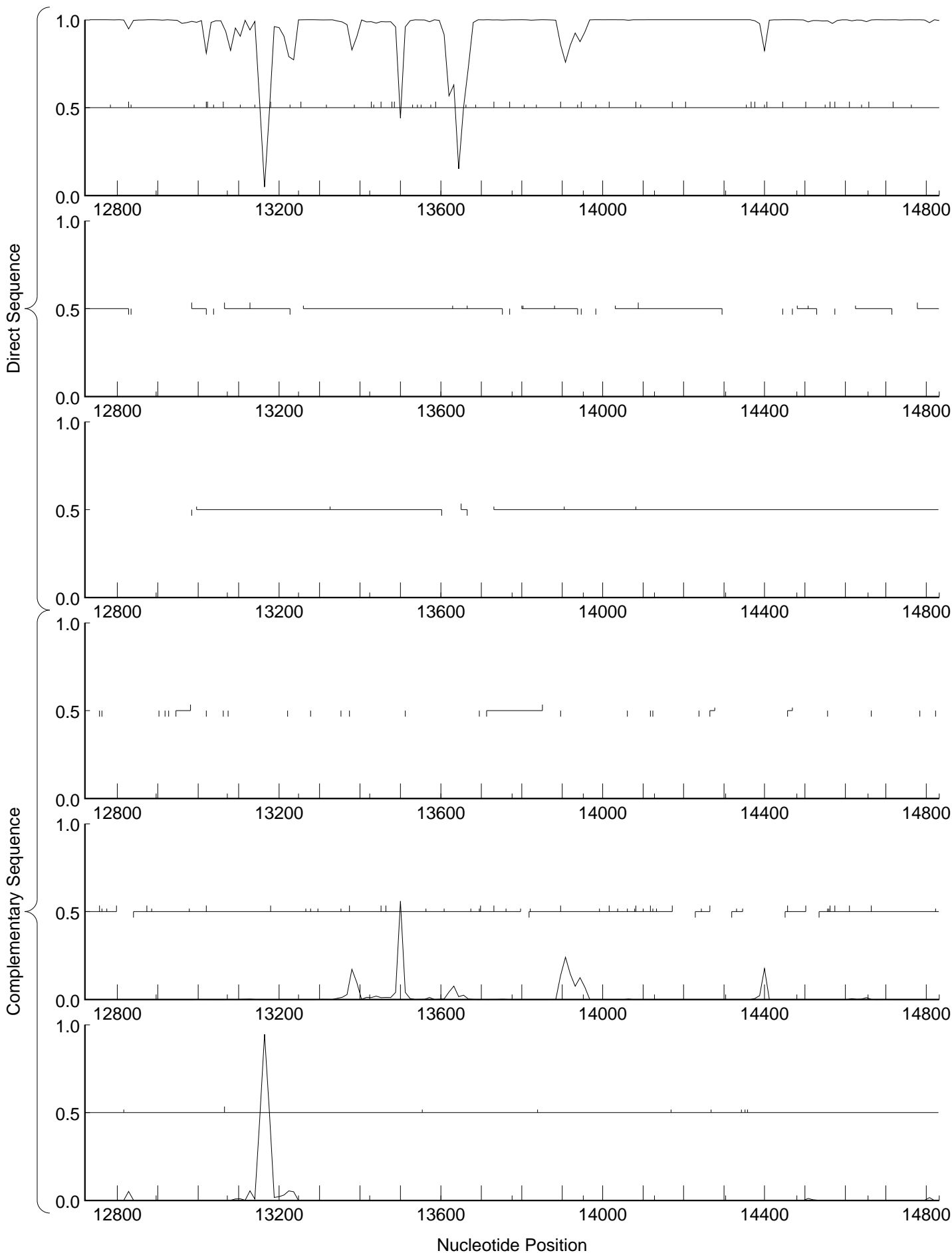


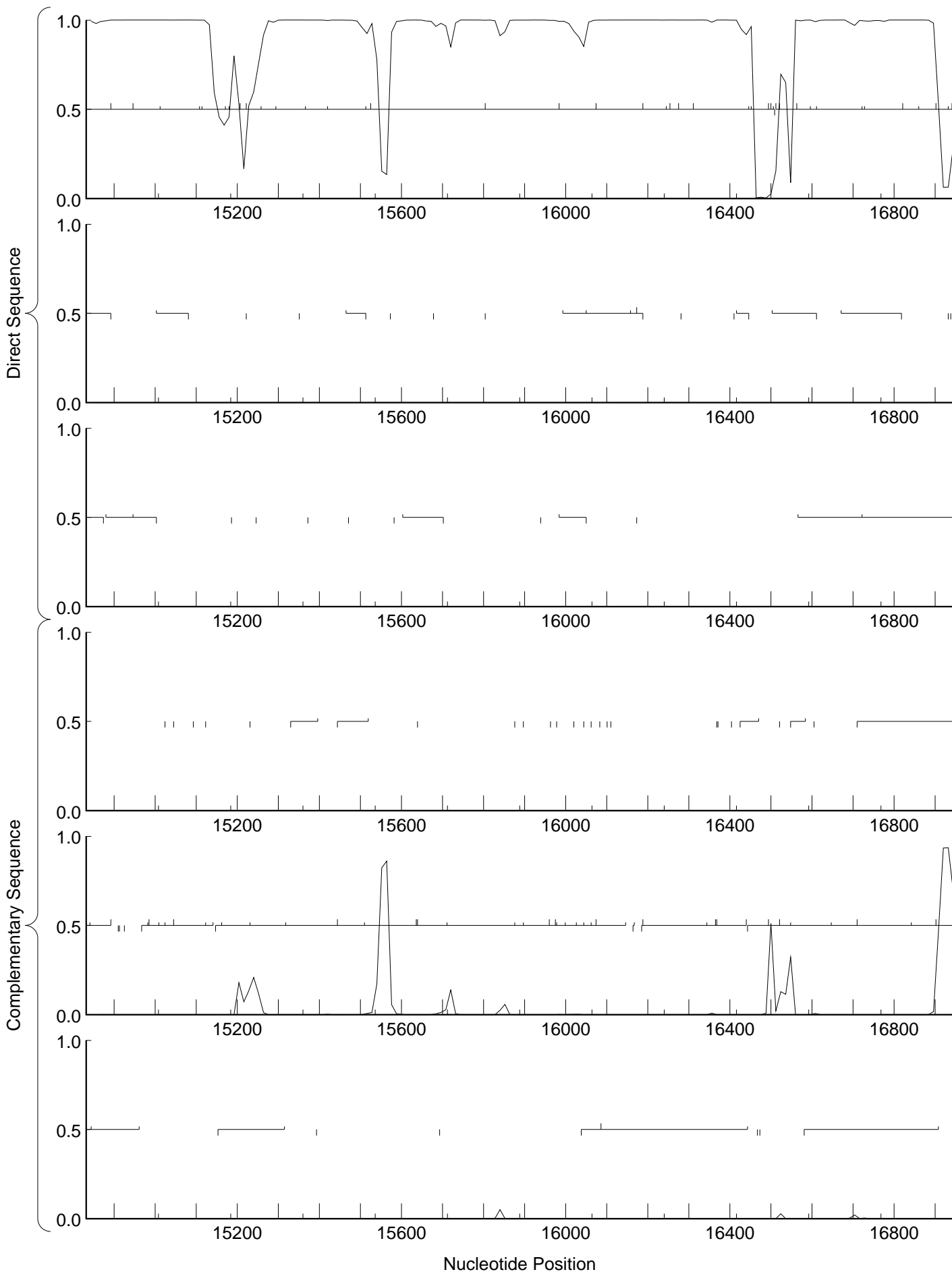


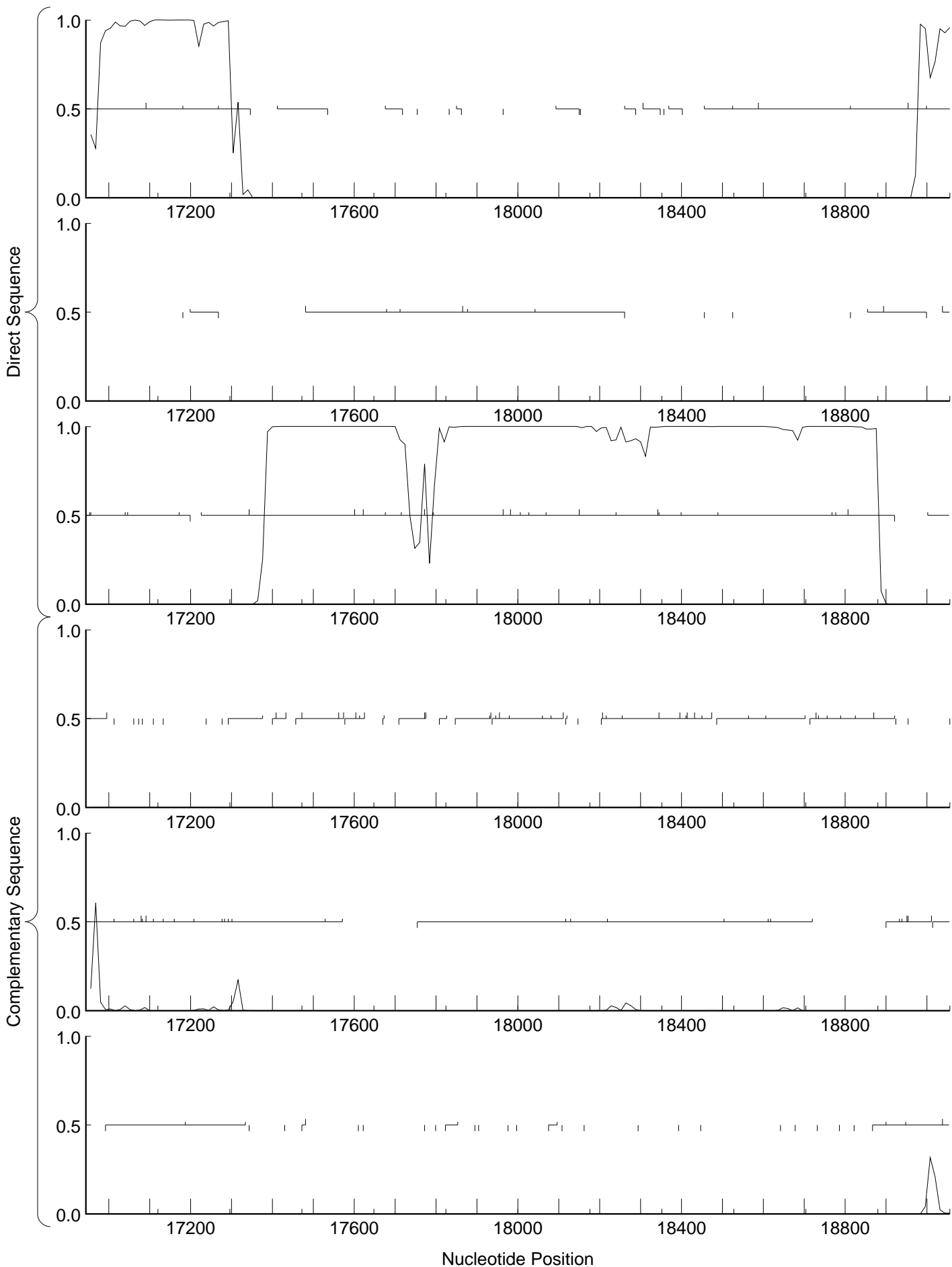
complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 4, Window 96, Step 12, 7/24



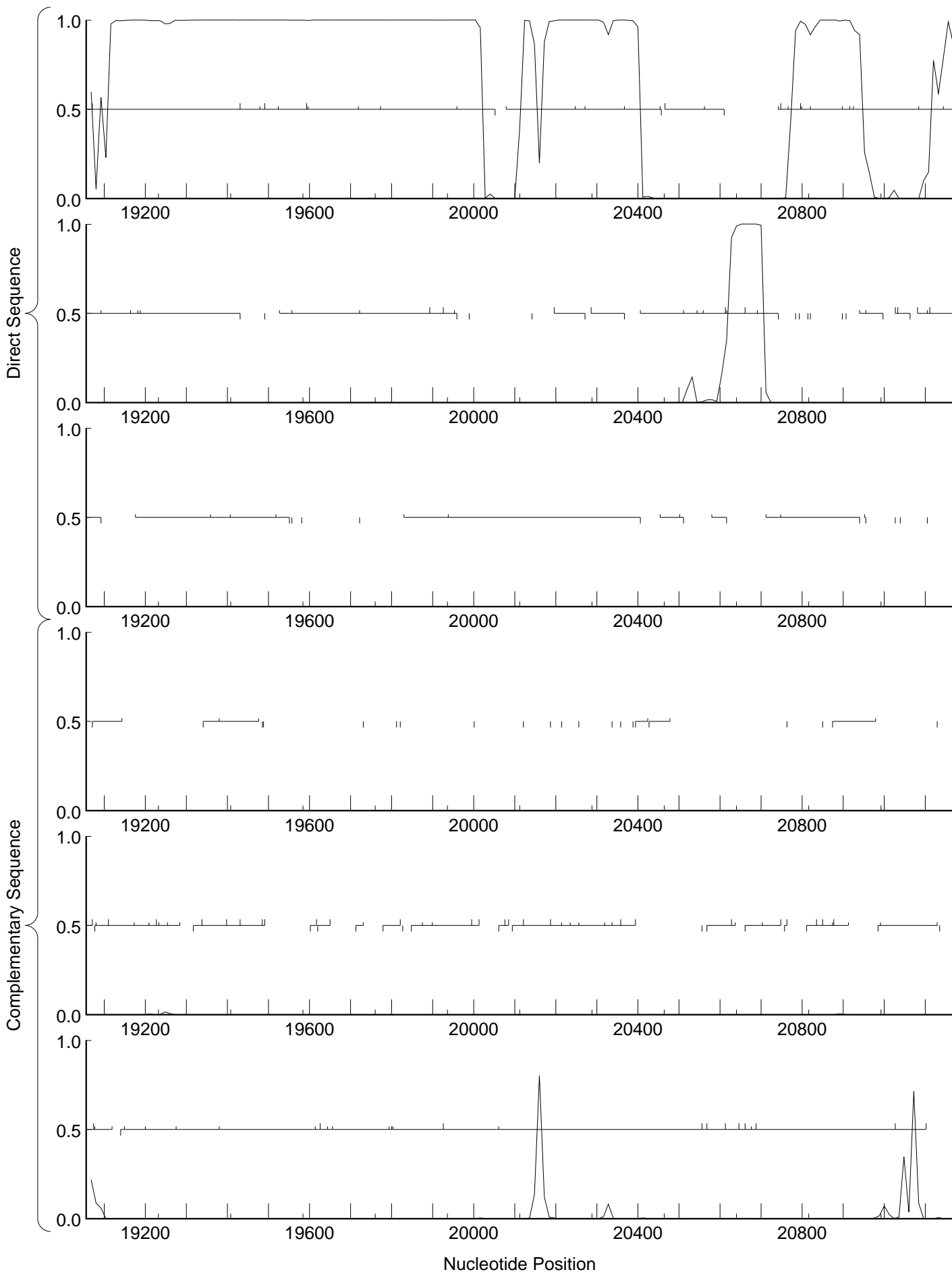
complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 4, Window 96, Step 12, 8/24

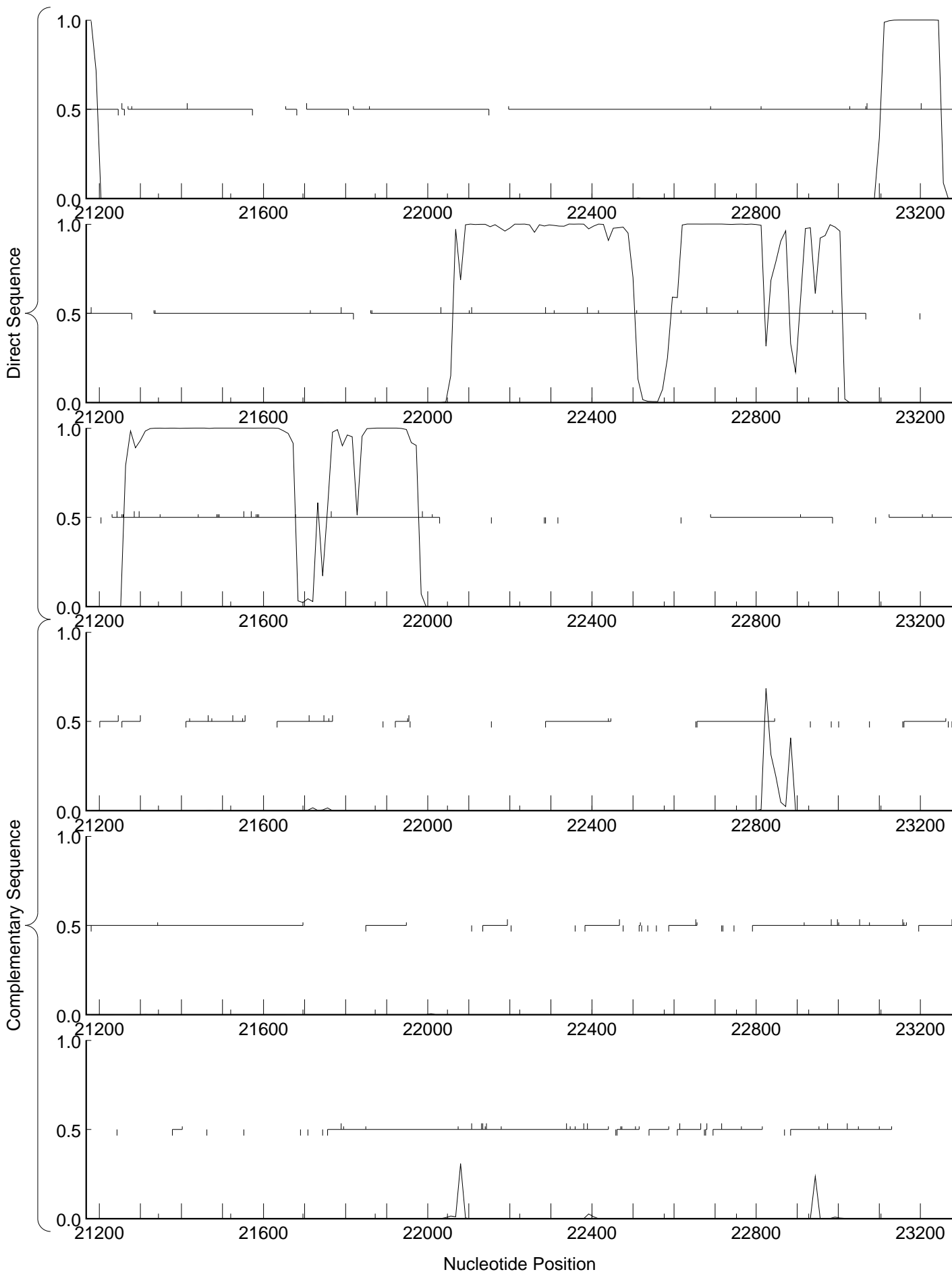




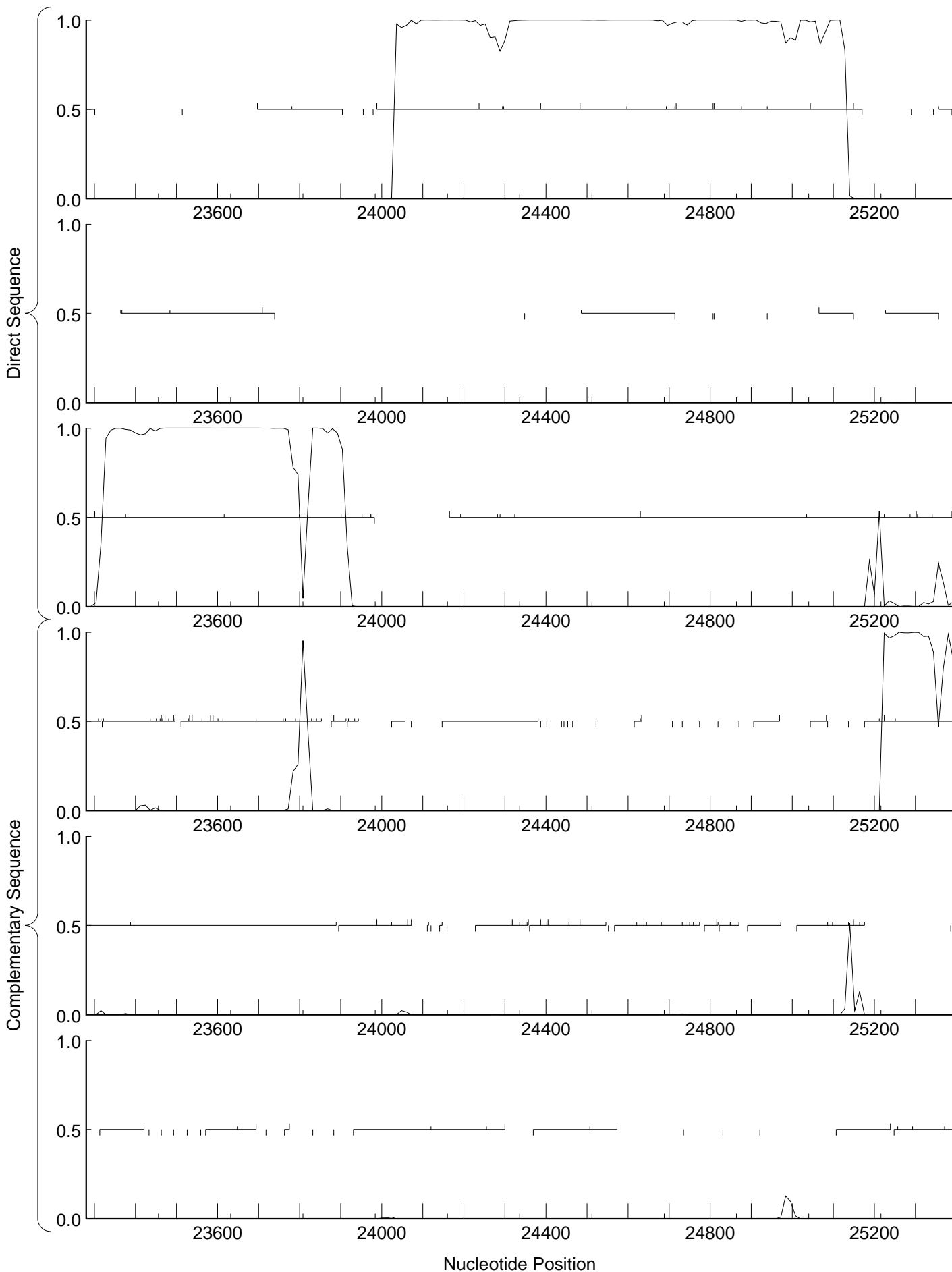


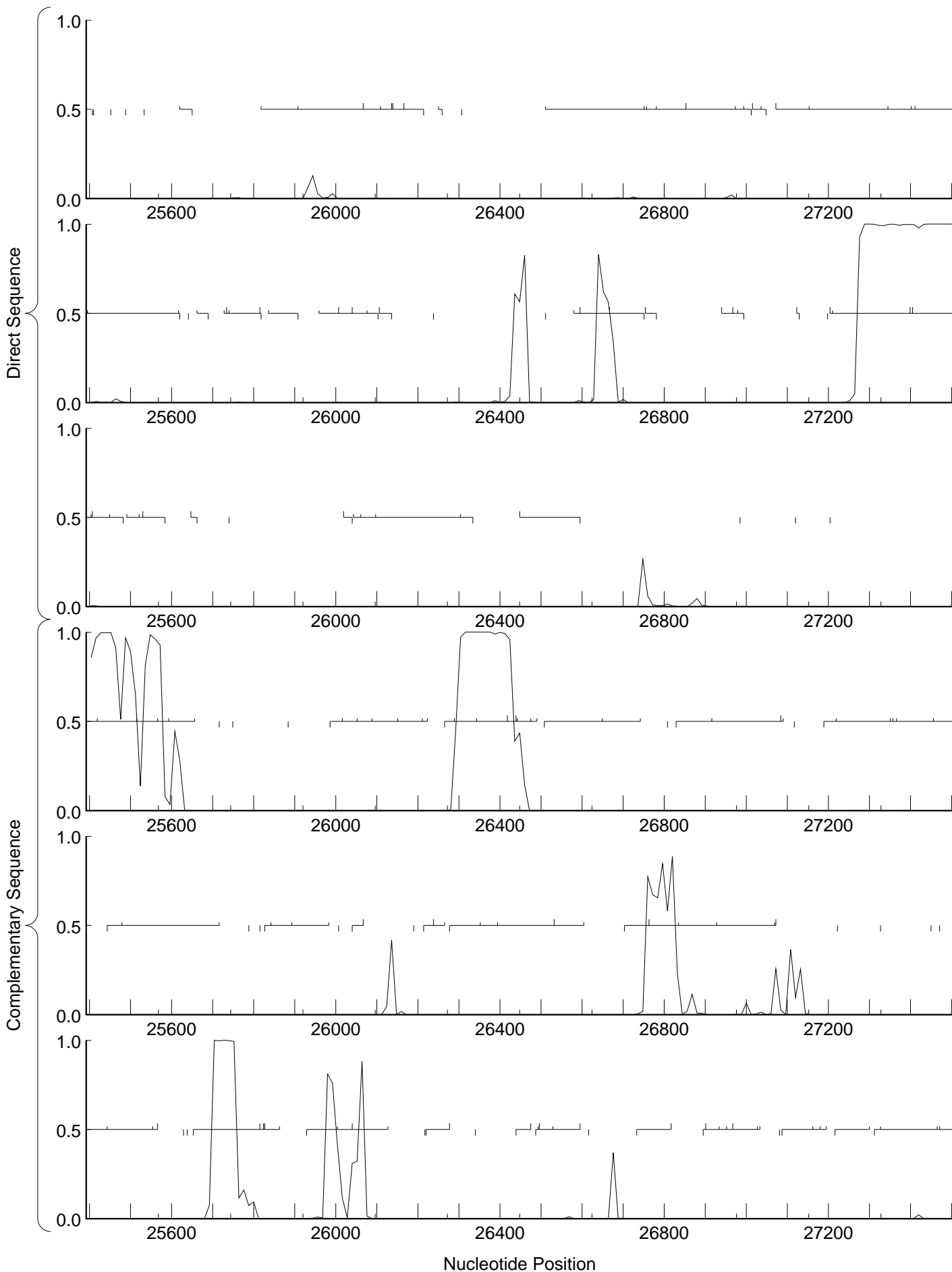
complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 4, Window 96, Step 12, 11/24

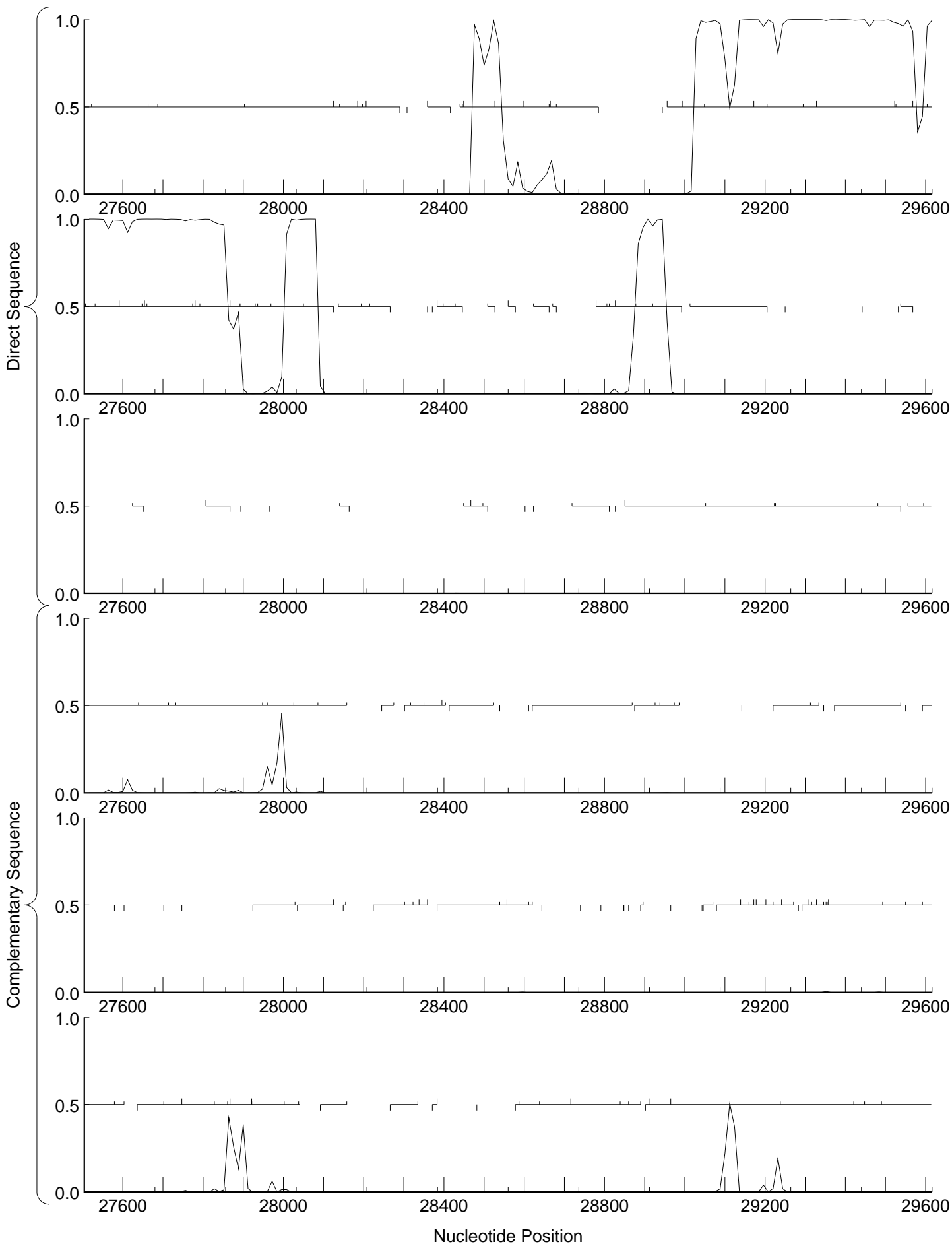


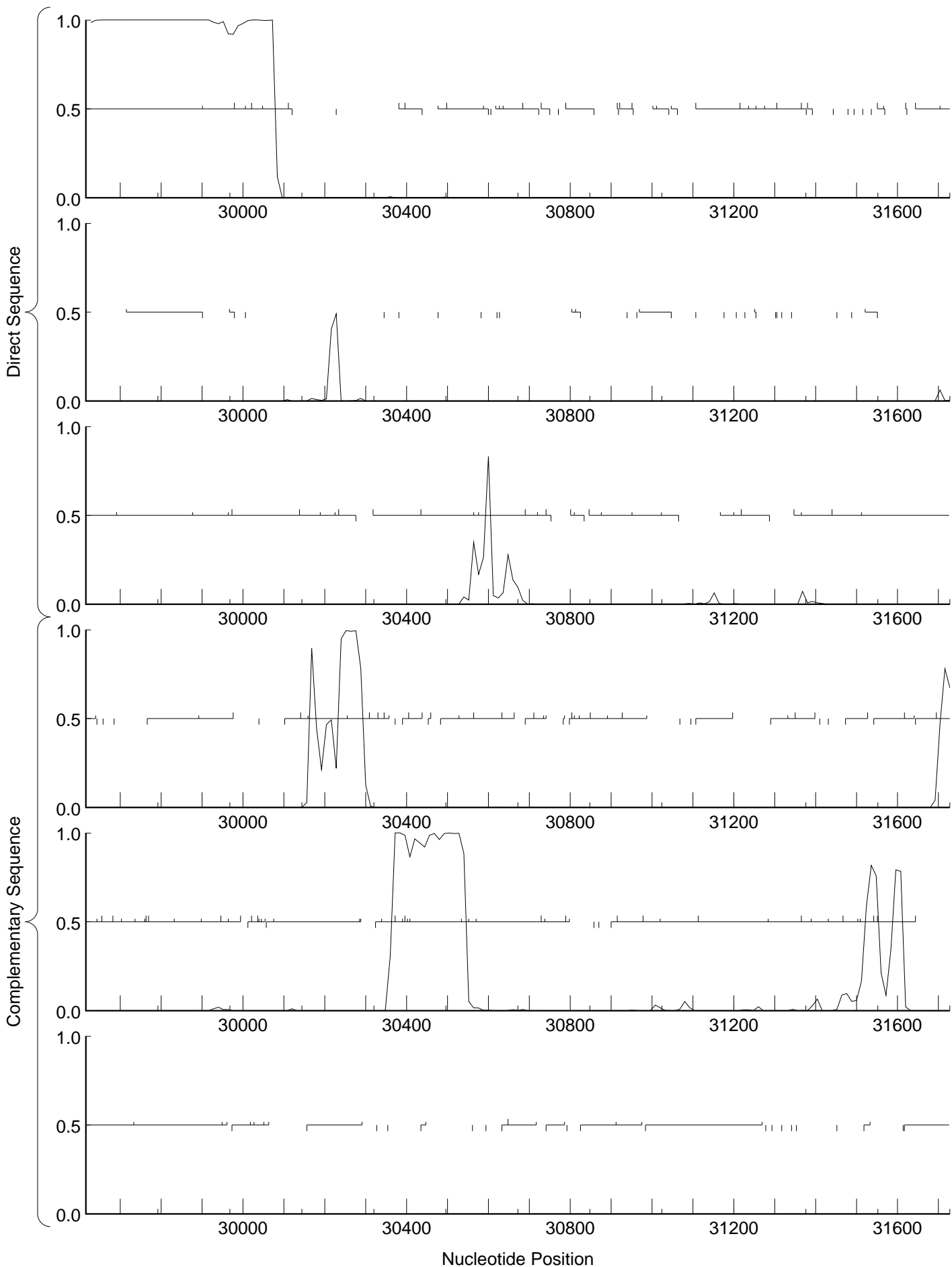


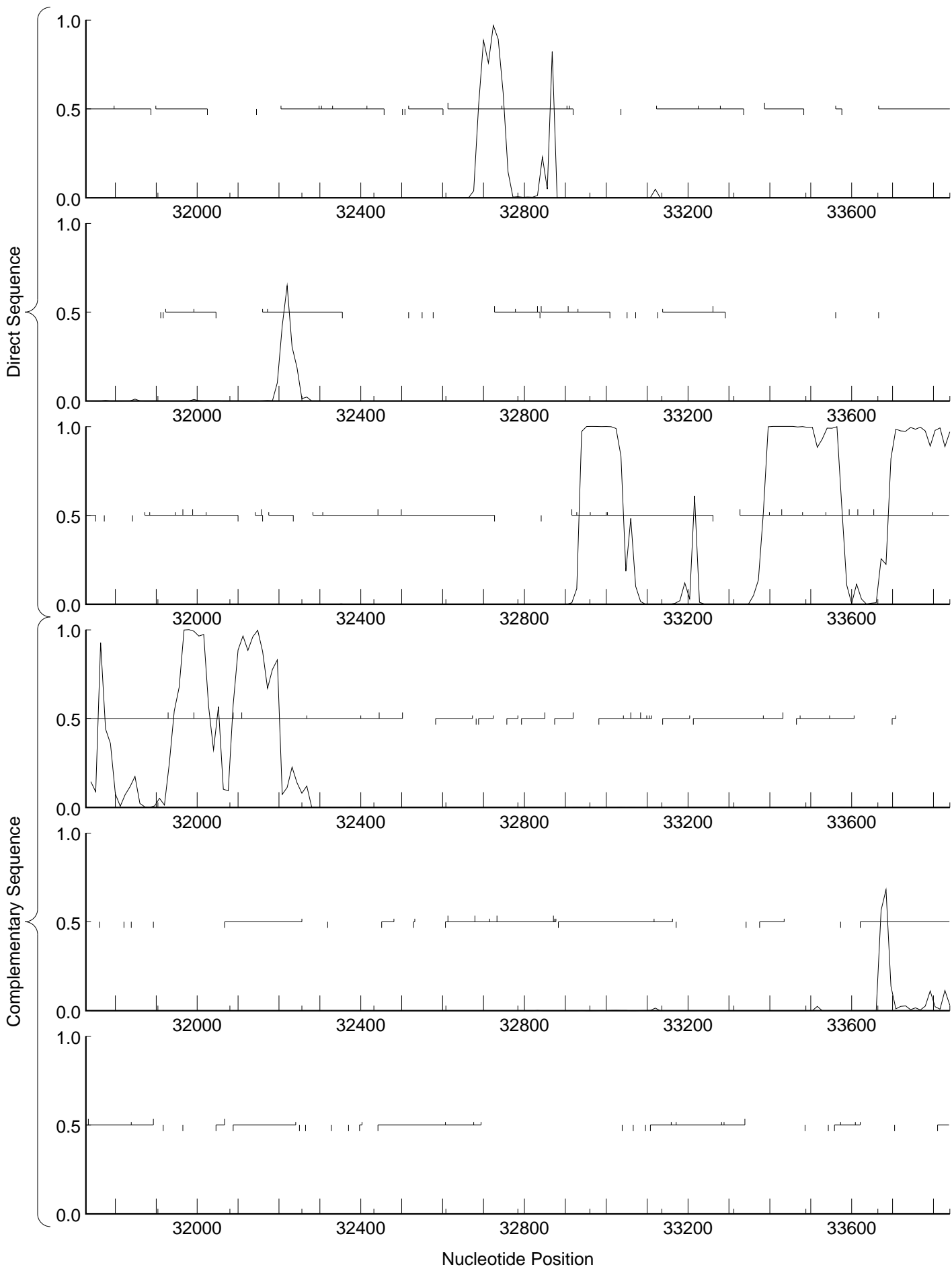
Complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 4, Window 96, Step 12, 13/24

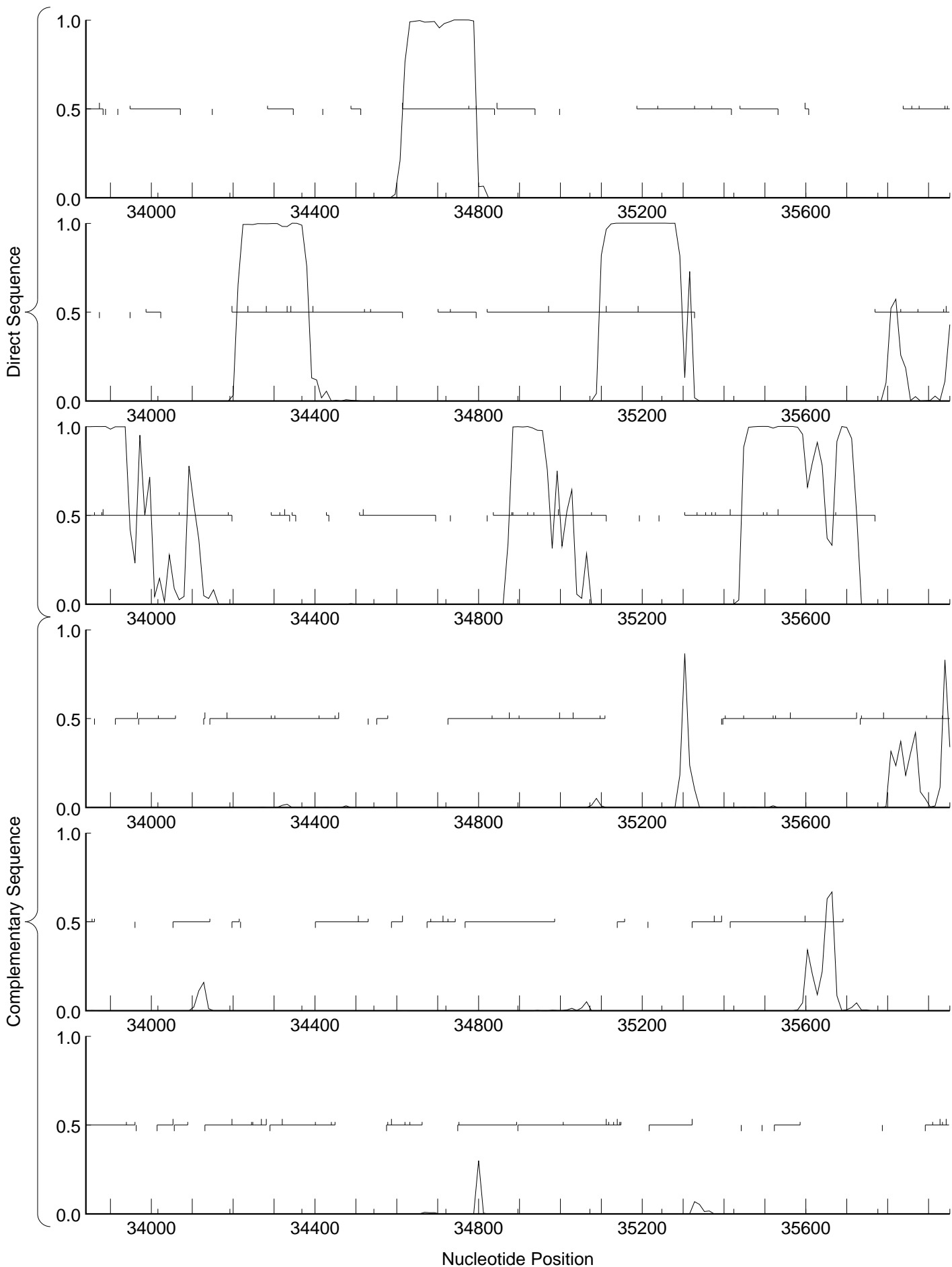


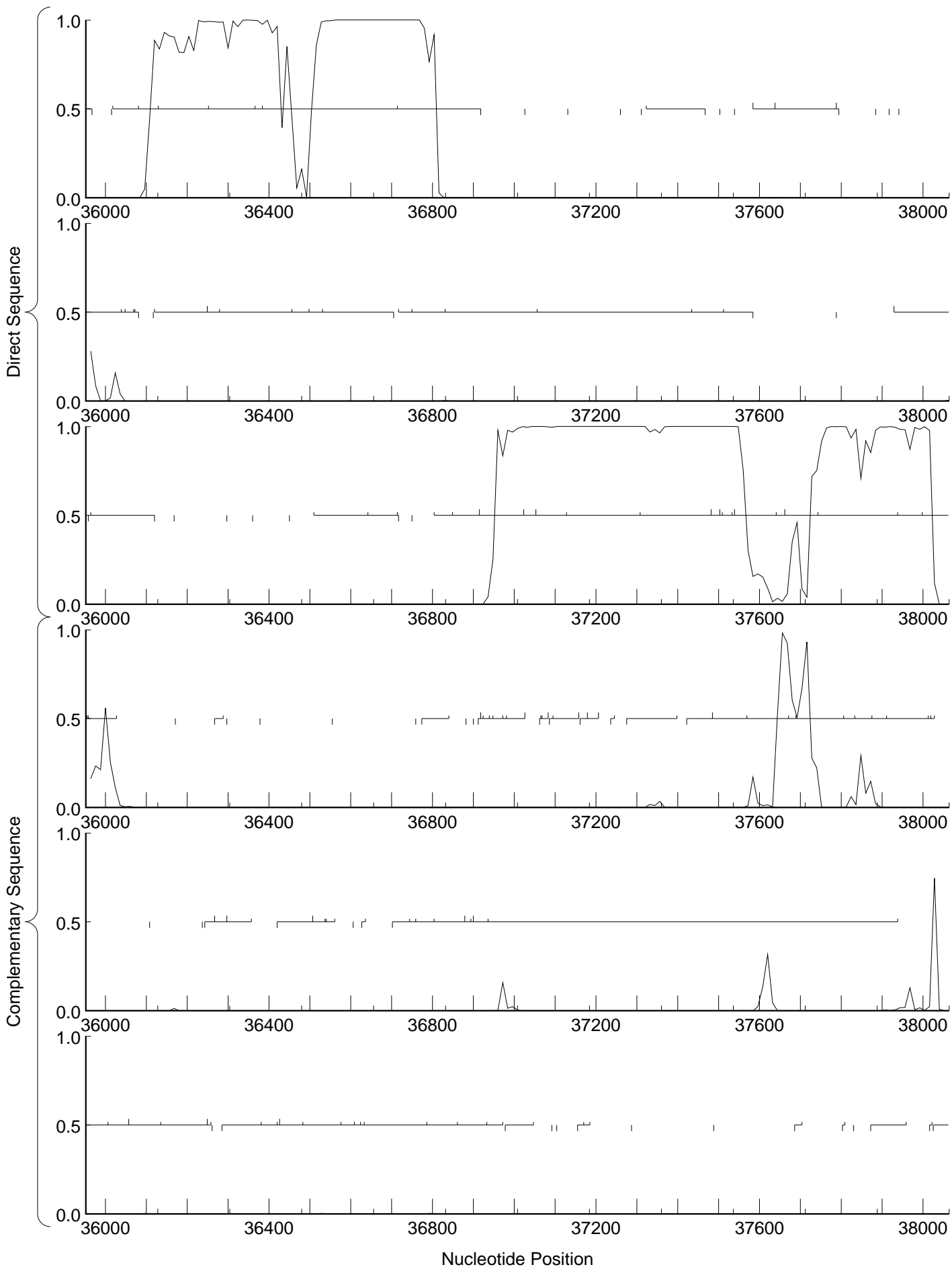


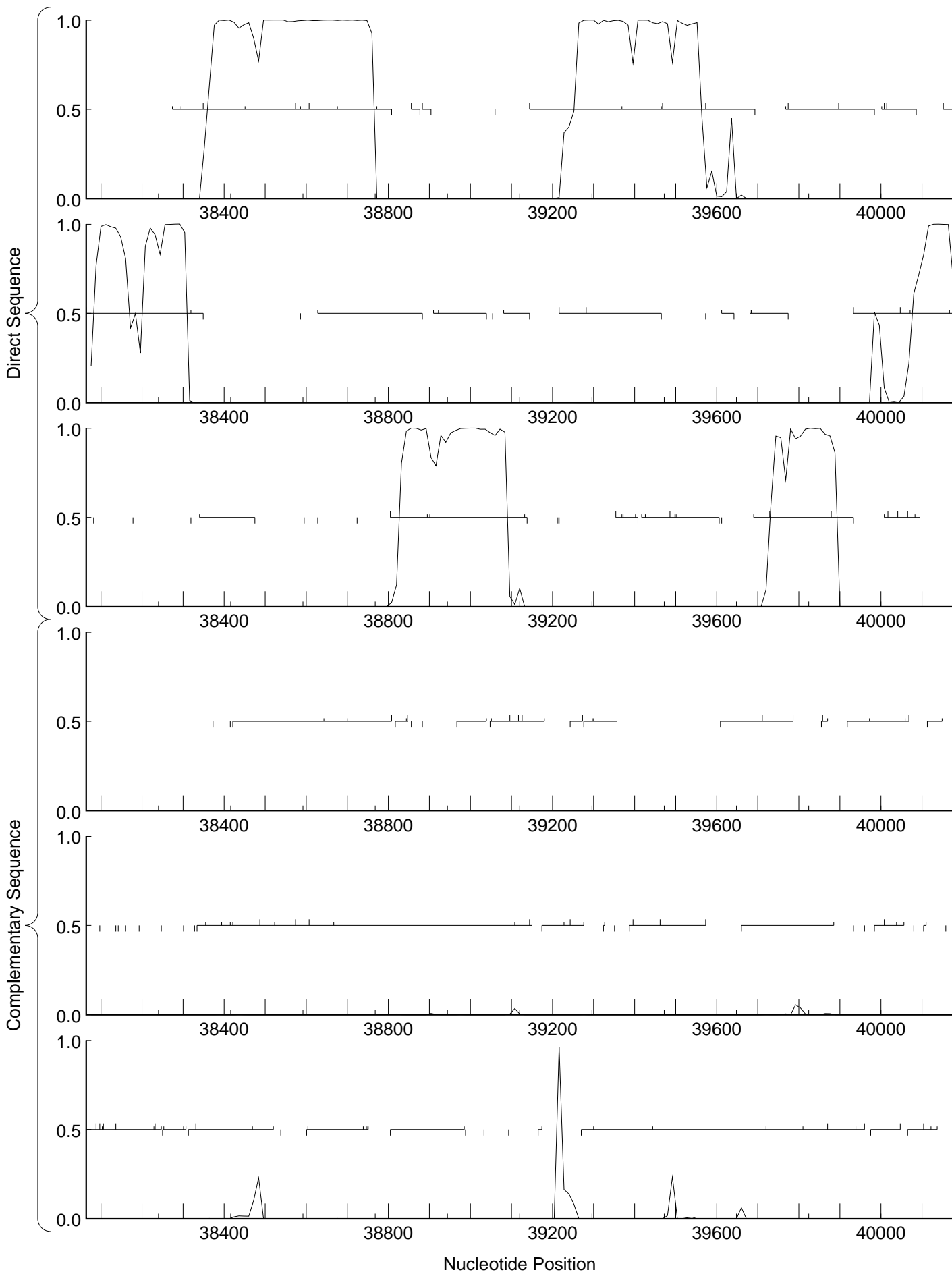


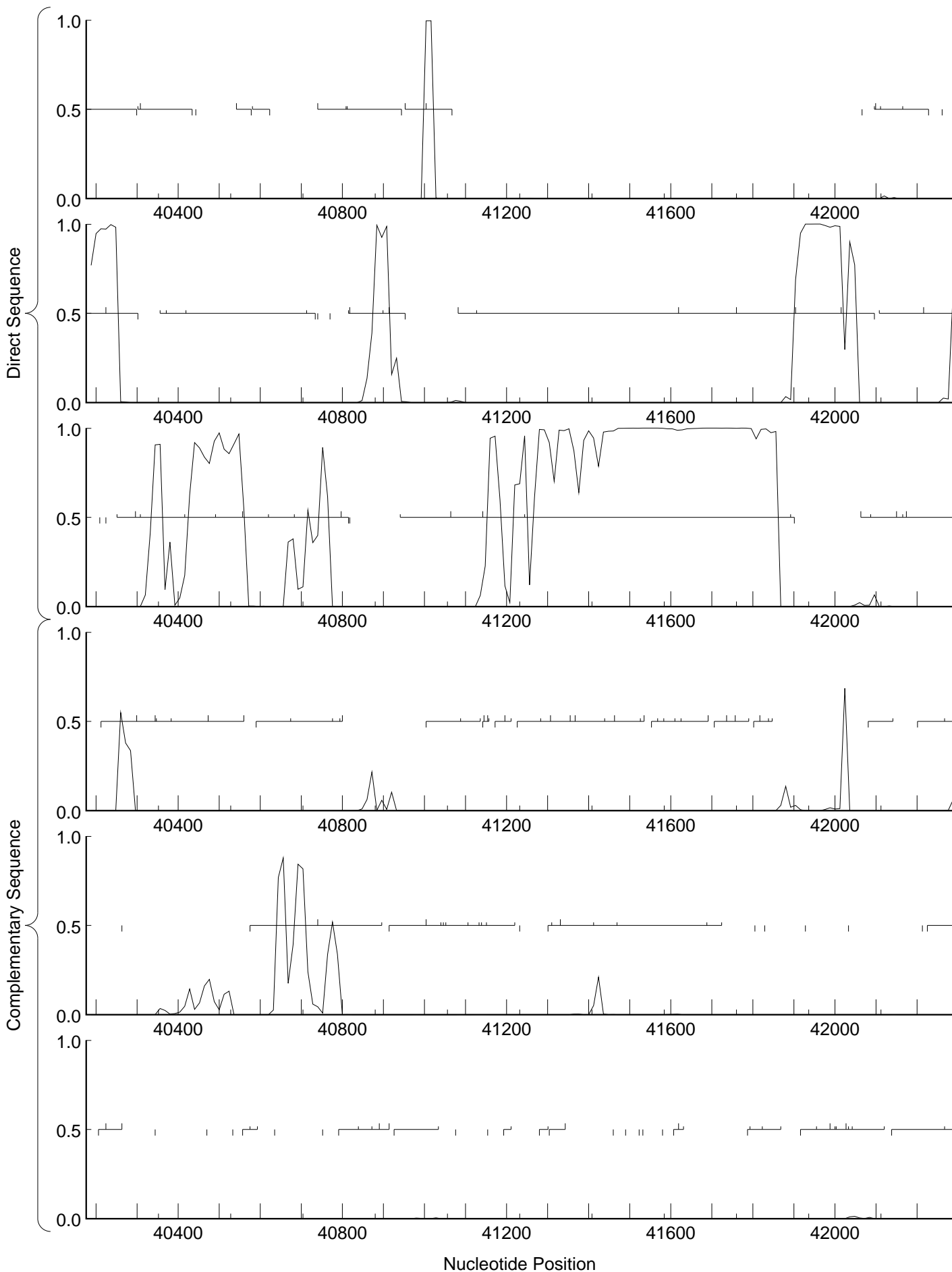


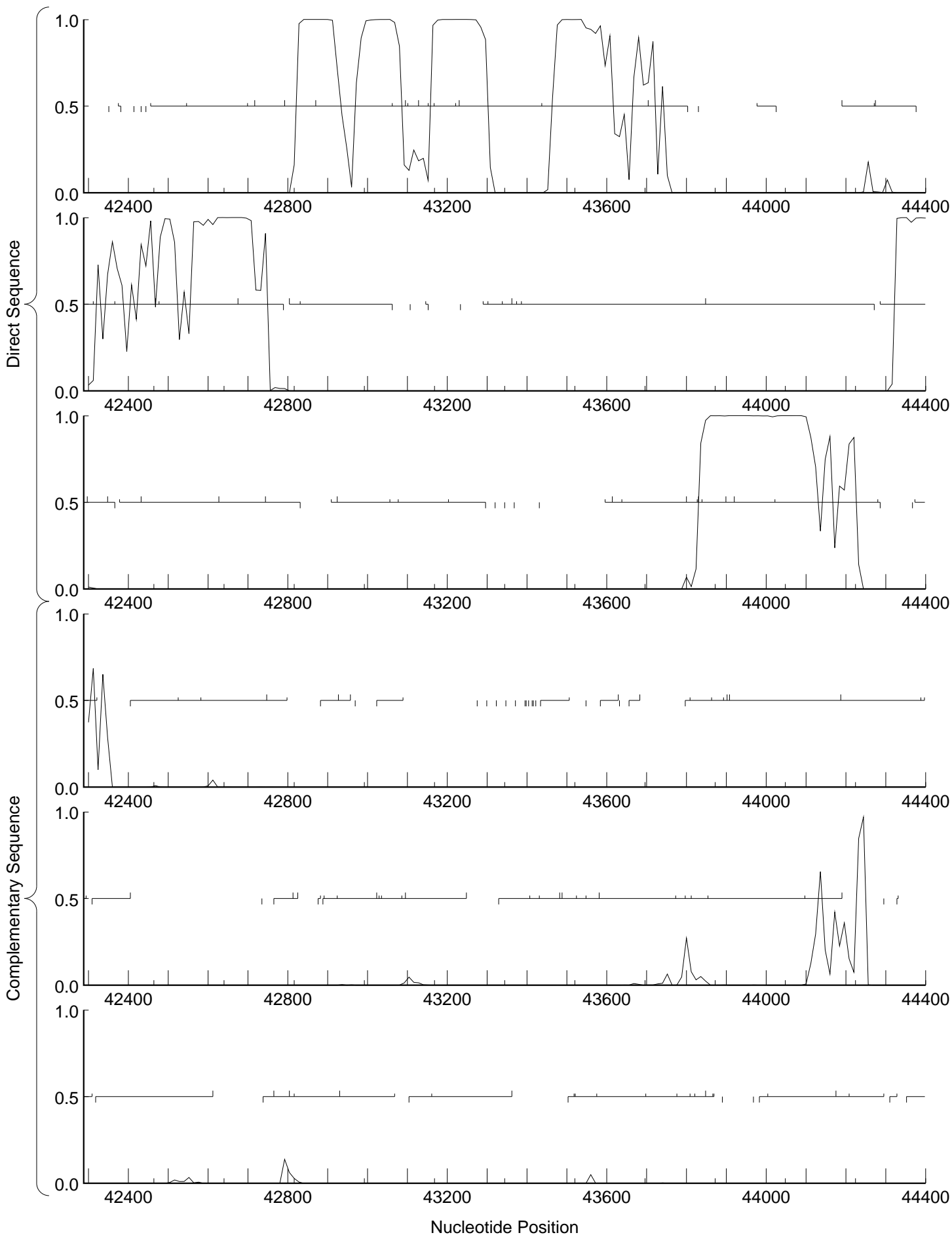












Complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 4, Window 96, Step 12, 23/24

