

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Mycobacterium phage Zulu complete sequence, 52499 bp including 10-base 3' overhang (CGGTCCG
Analysis Date : 8/20/17 at 11:36:31
Pages : 26
Sequence Length : 52499 bp
GC Content : 61.42%

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,

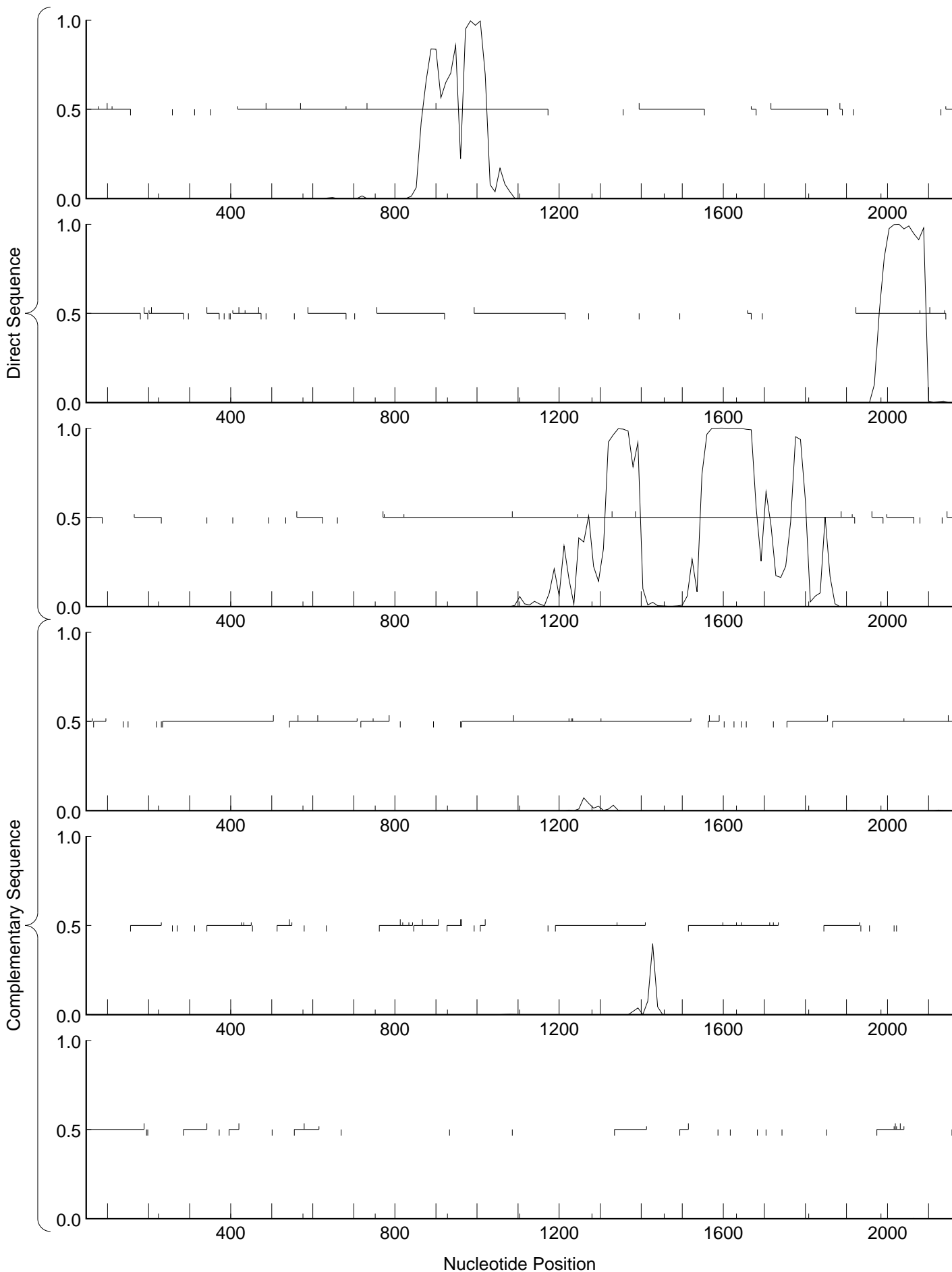
Matrix notes & comments

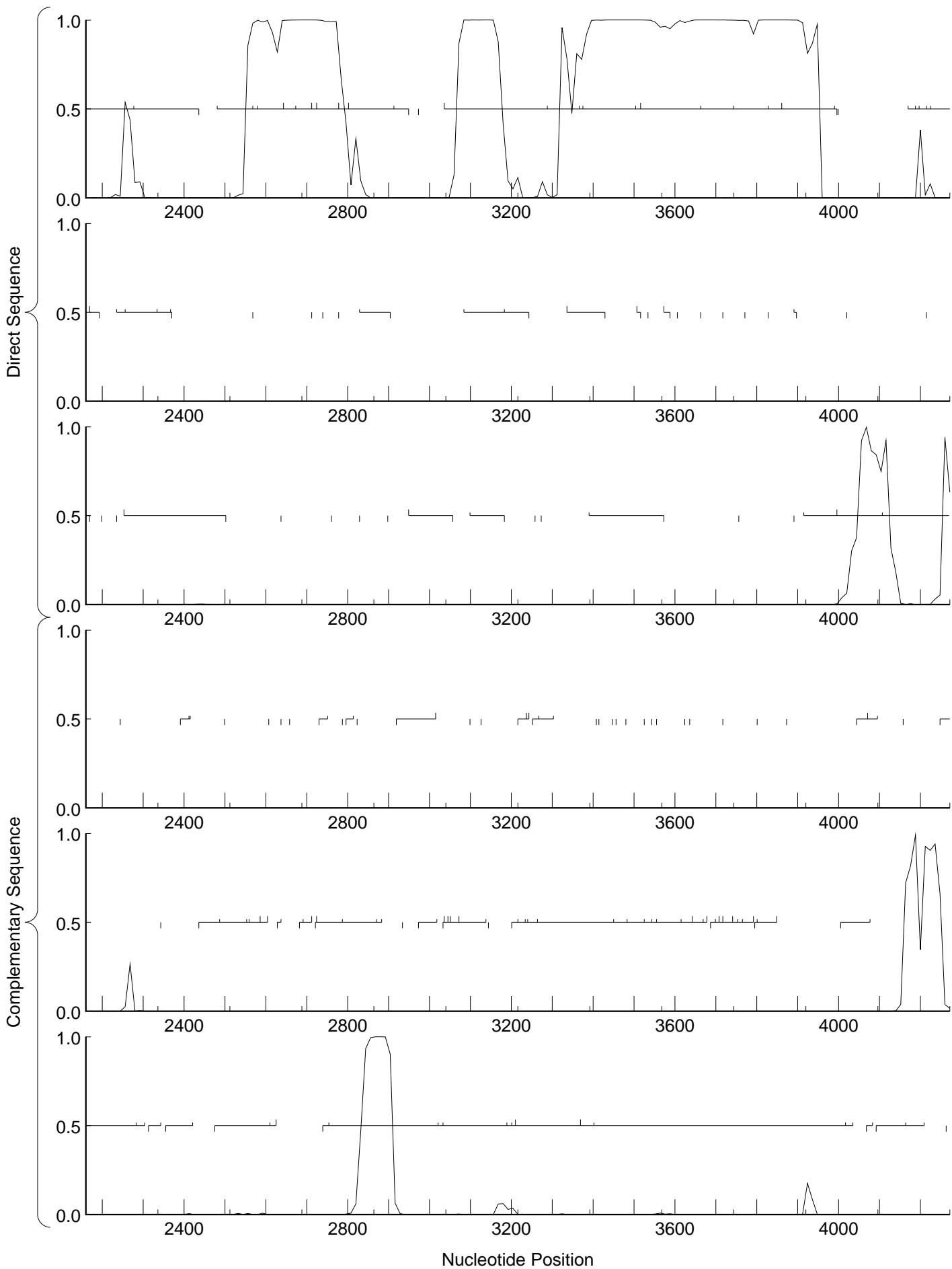
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:23:03 2014

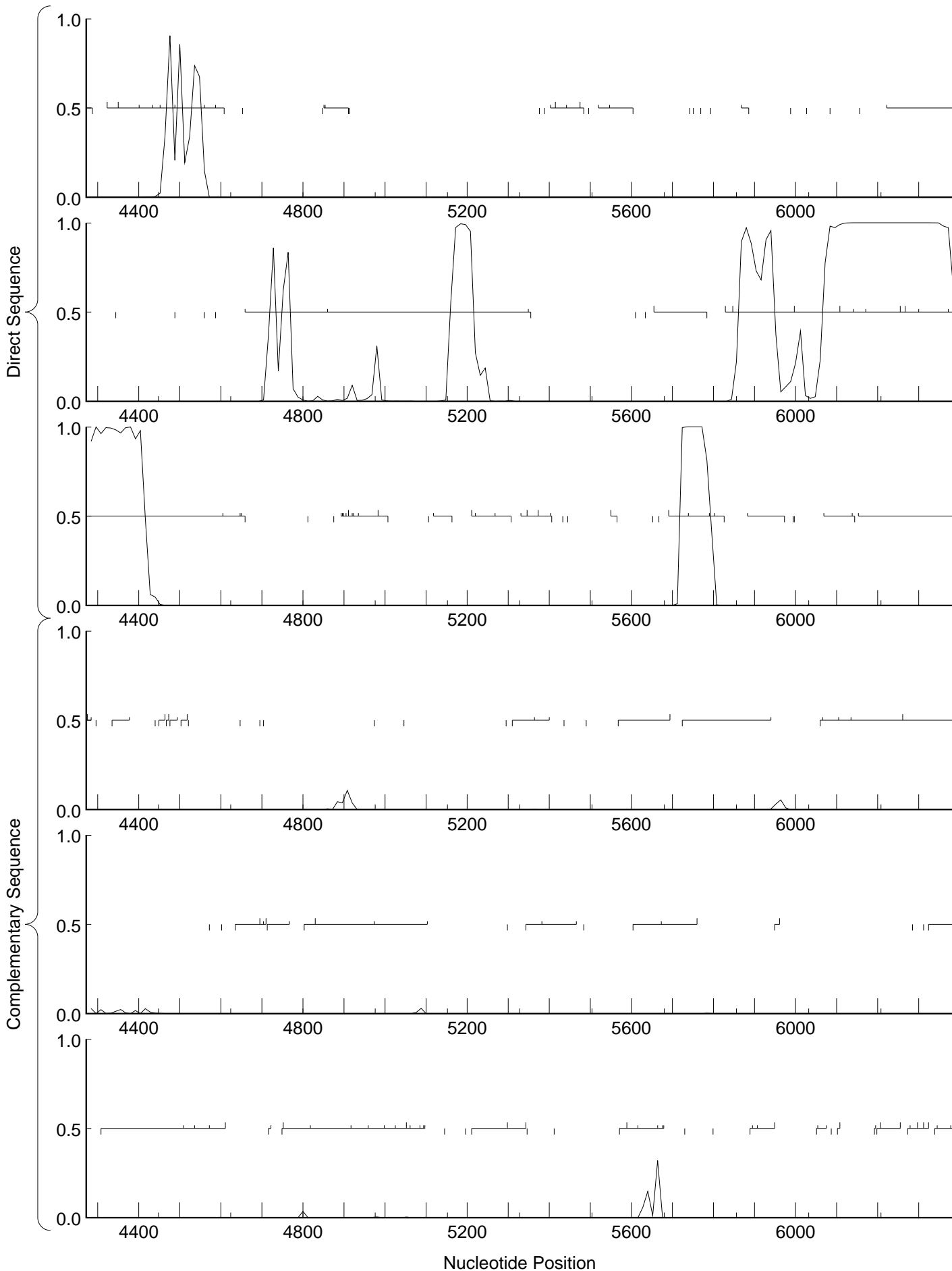
MATRIX INFORMATION

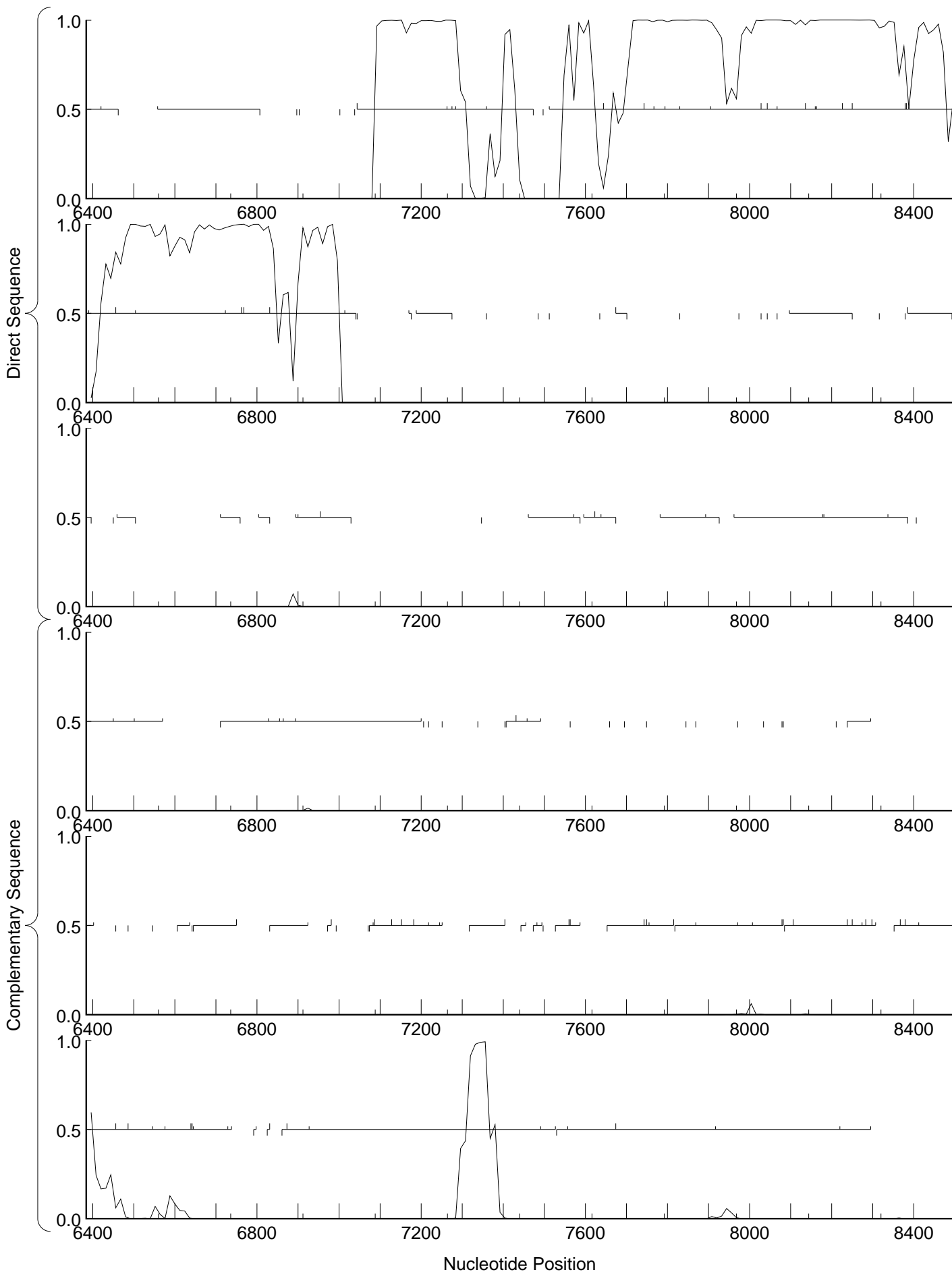
Matrix : Mycobacterium_smegmatis_MC2_155
Author : -
Order : 4

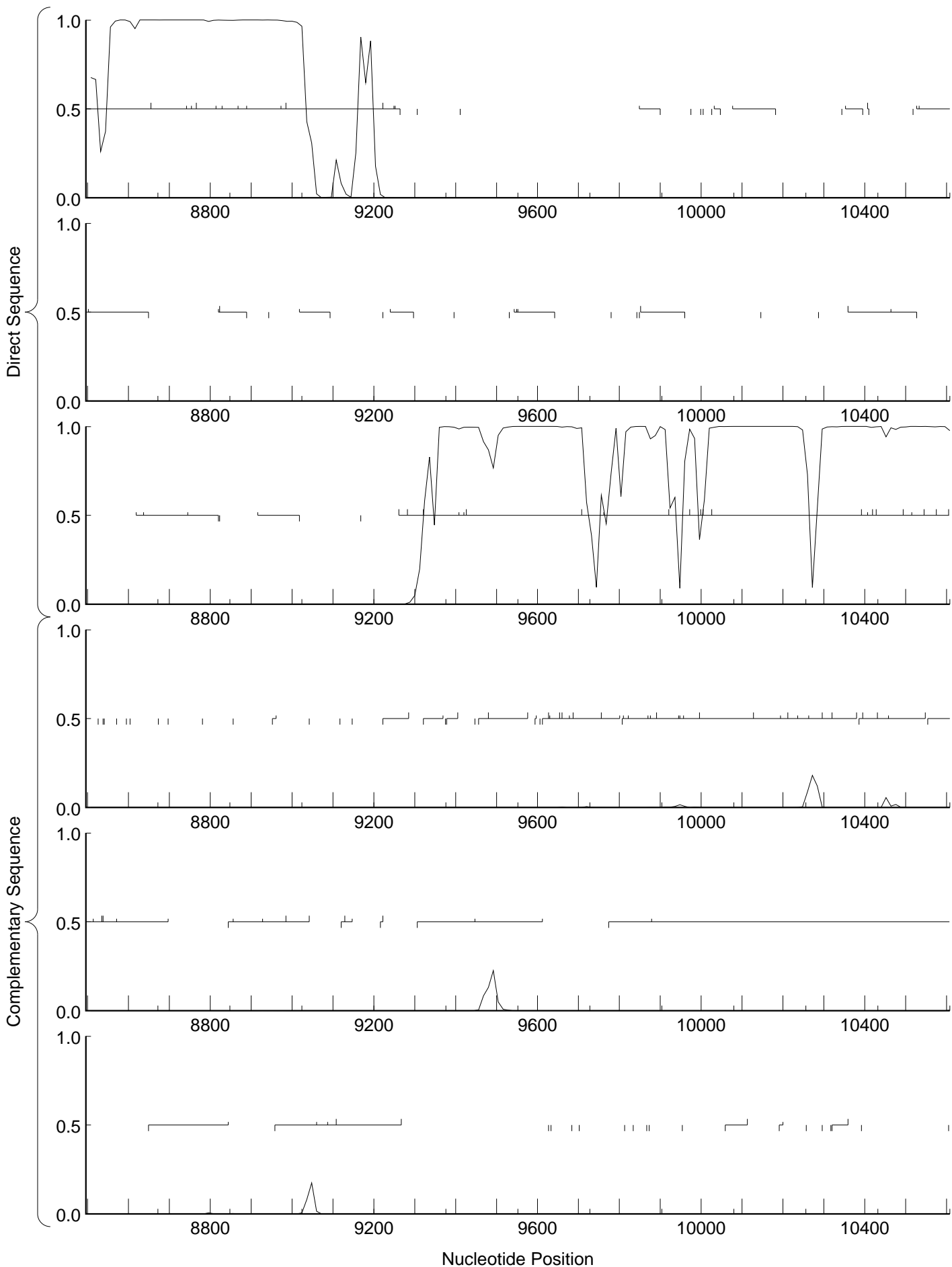
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

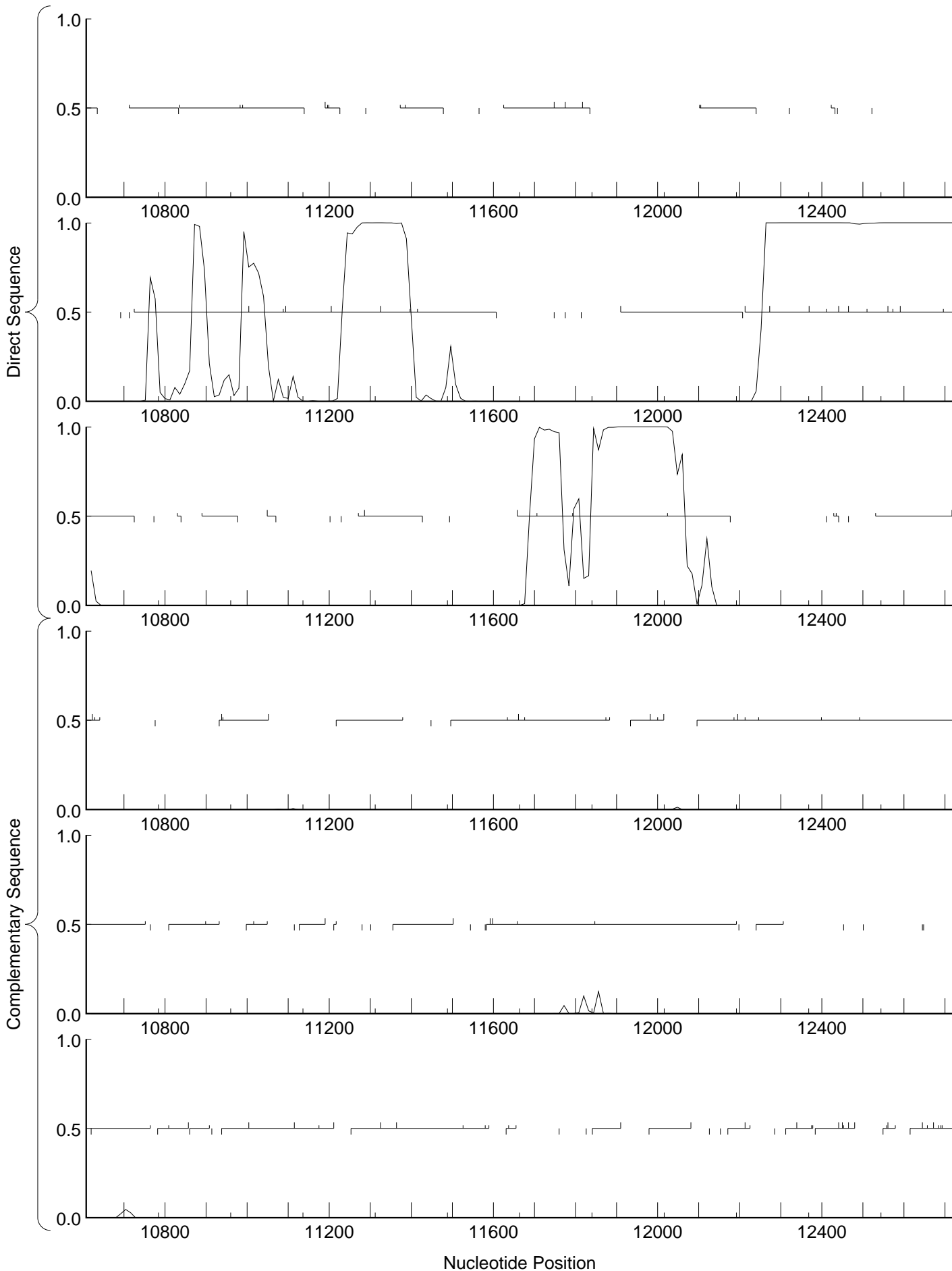


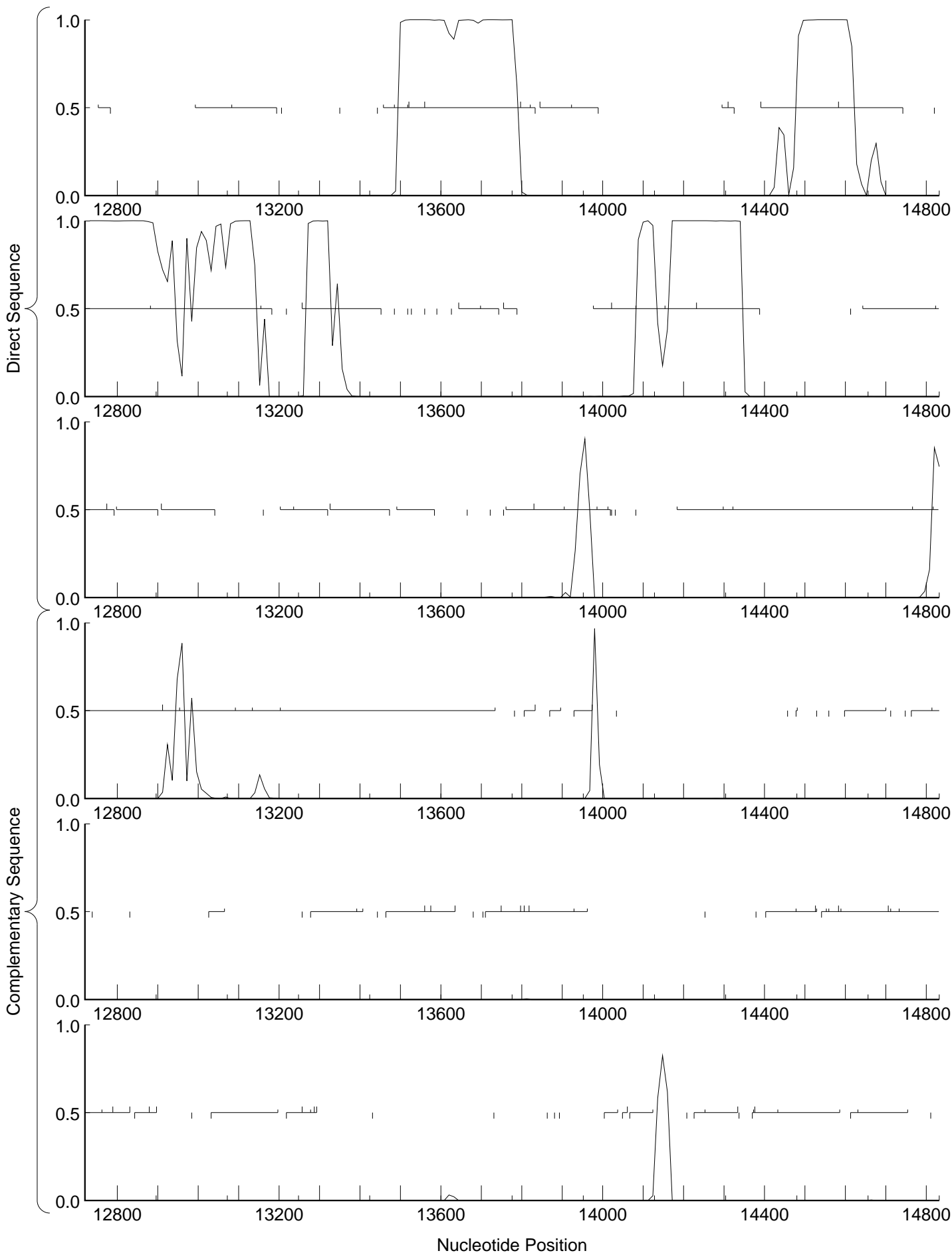


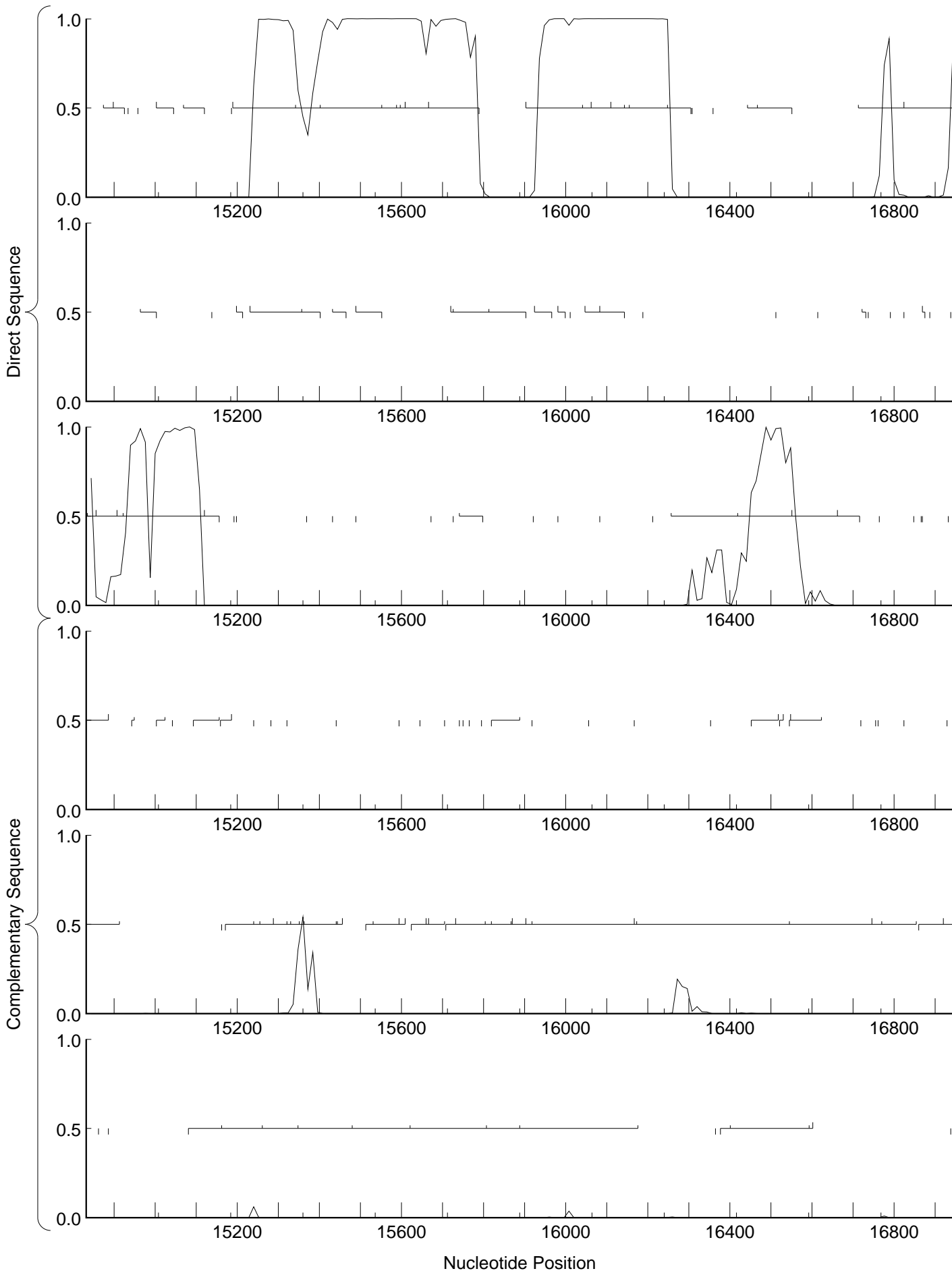




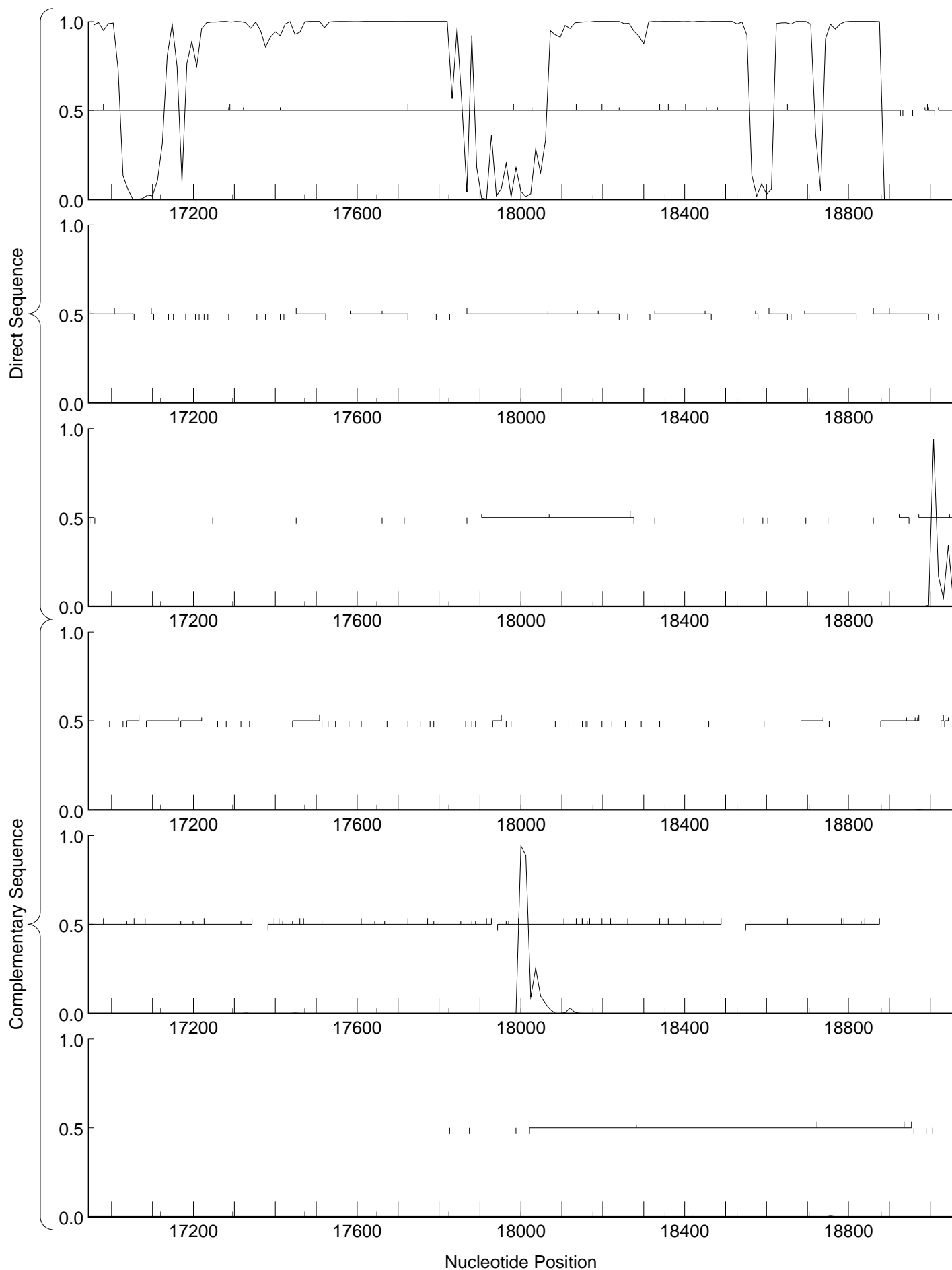


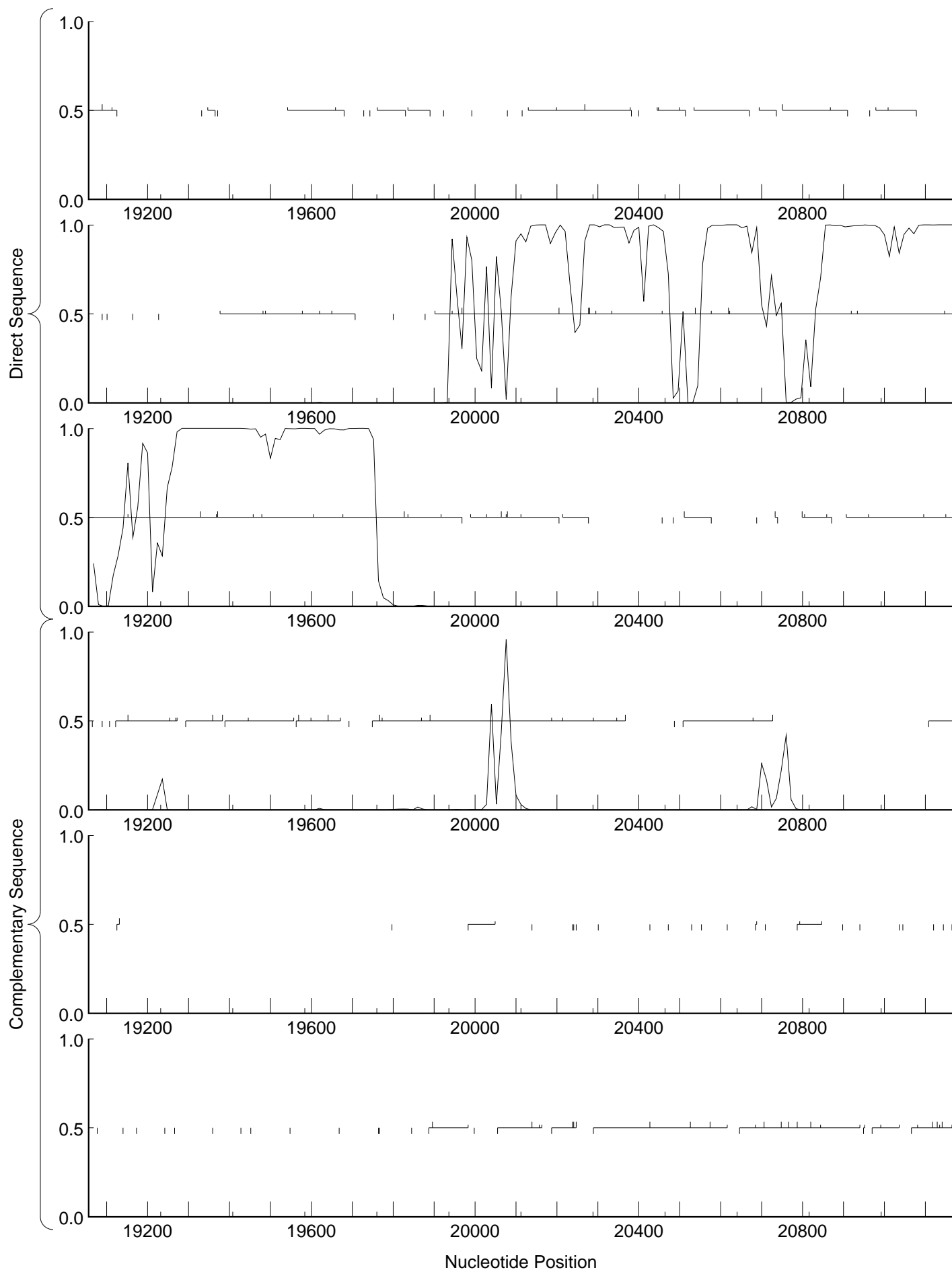


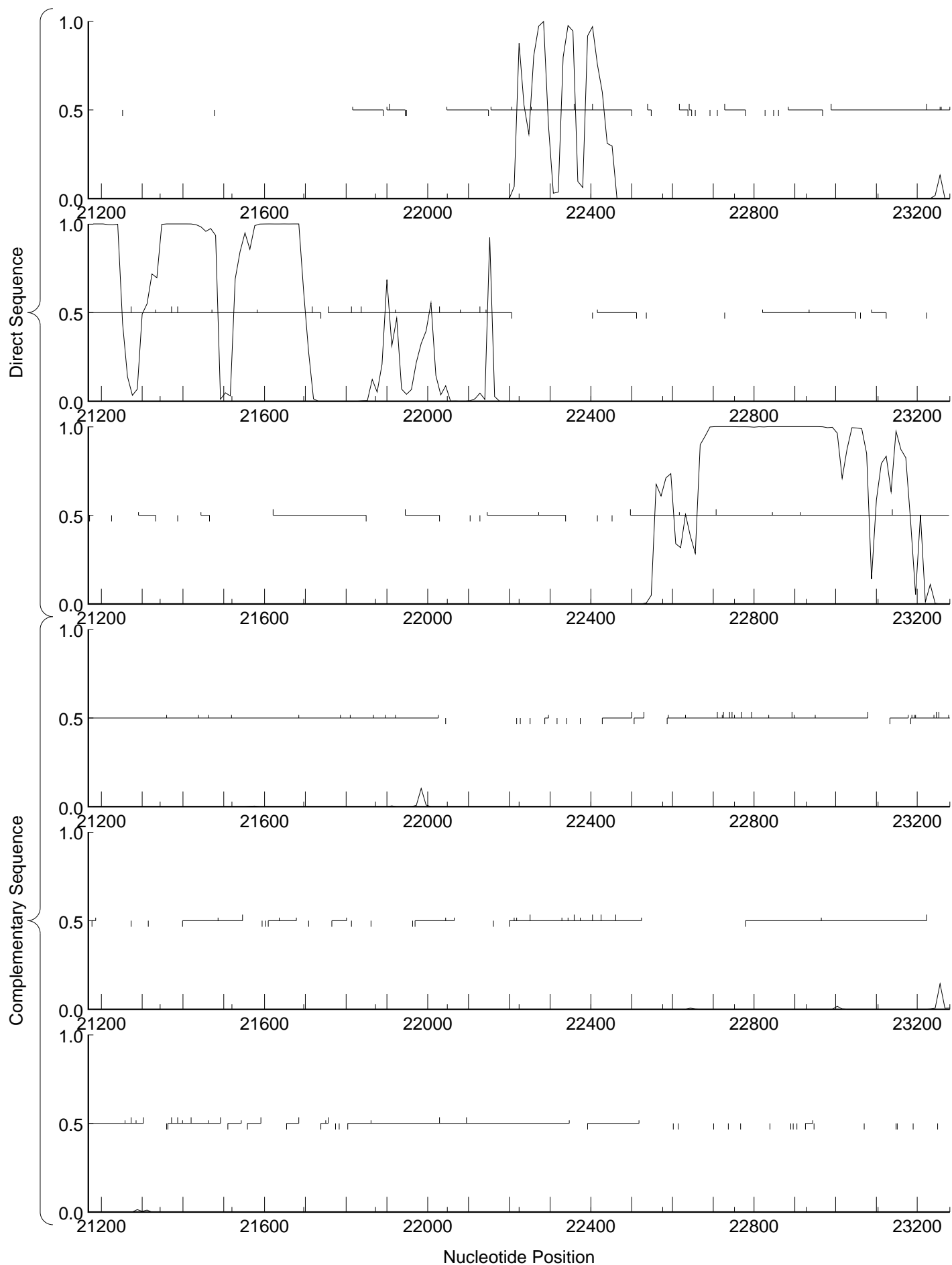




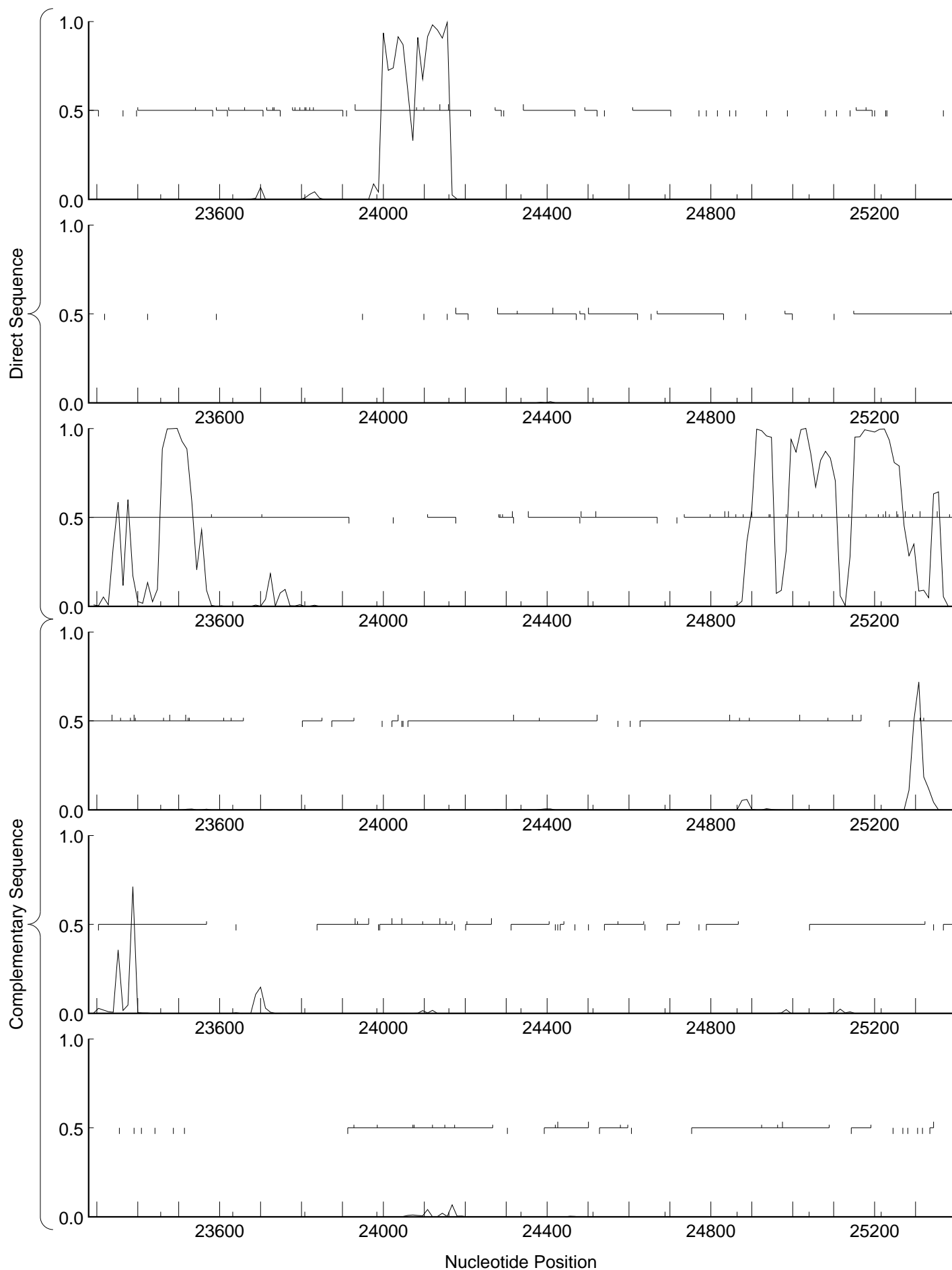
complete sequence, 52499 bp including 10-base 3' overhang (CGGTTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 10/26

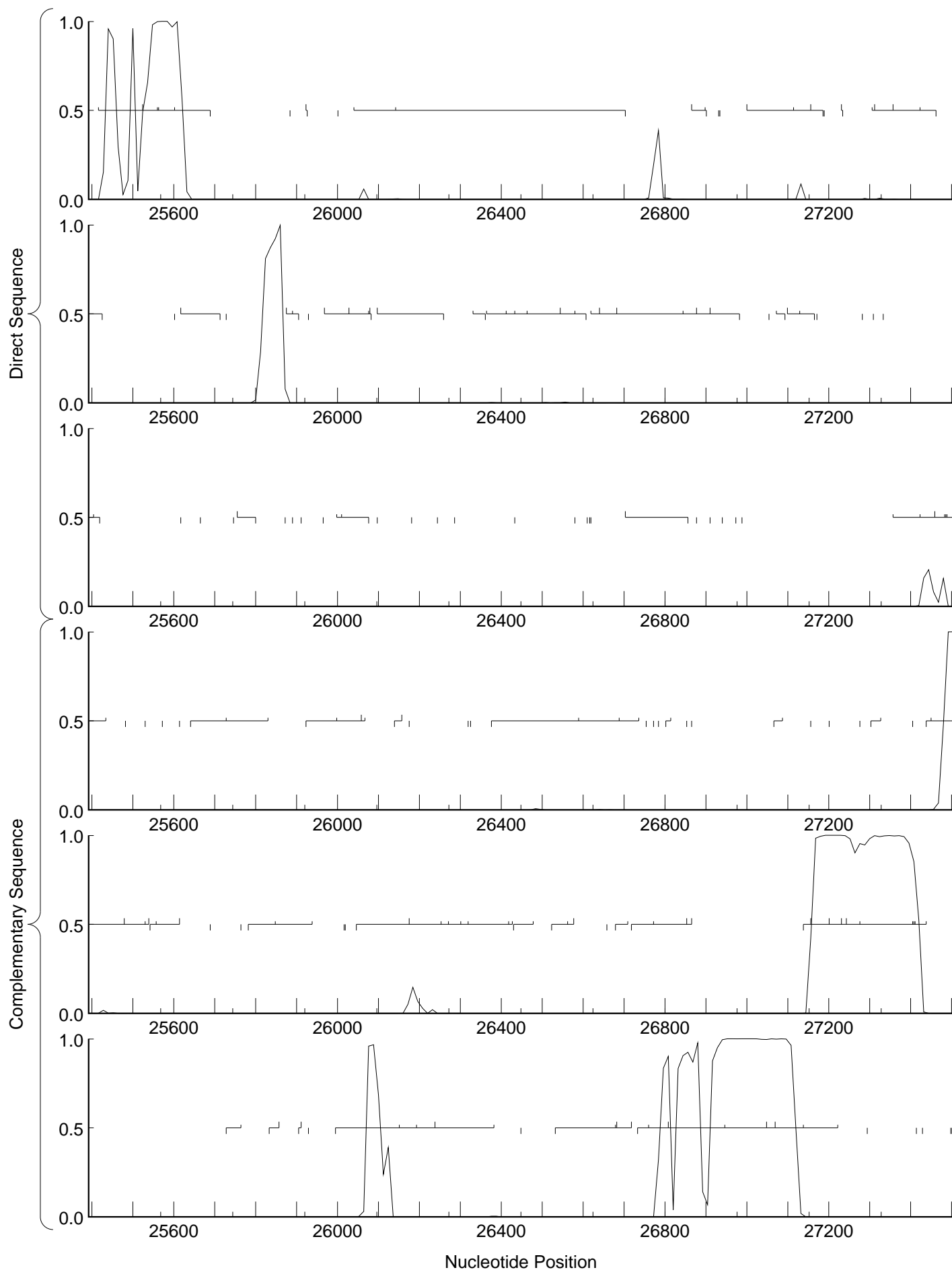




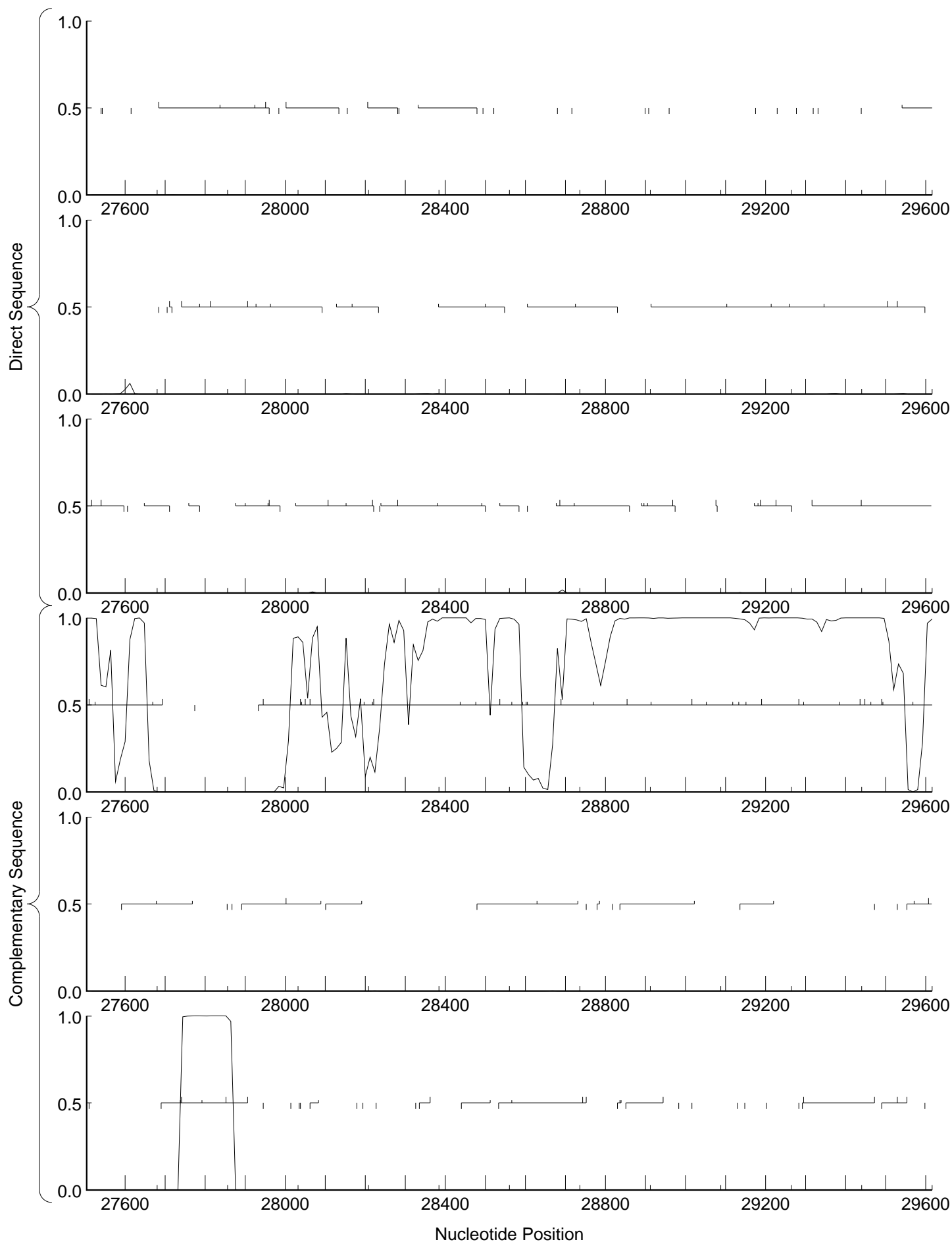


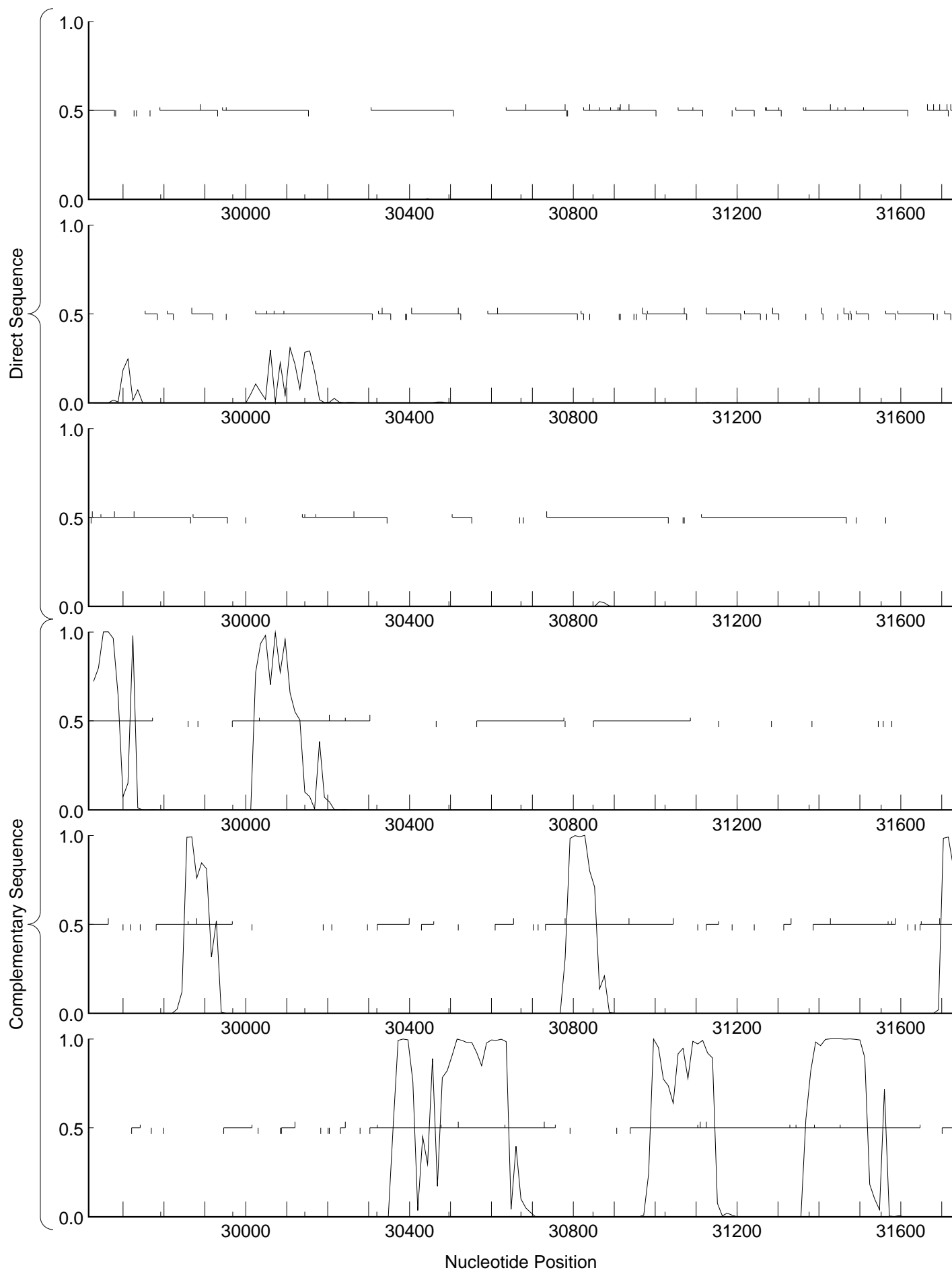
complete sequence, 52499 bp including 10-base 3' overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 13/26



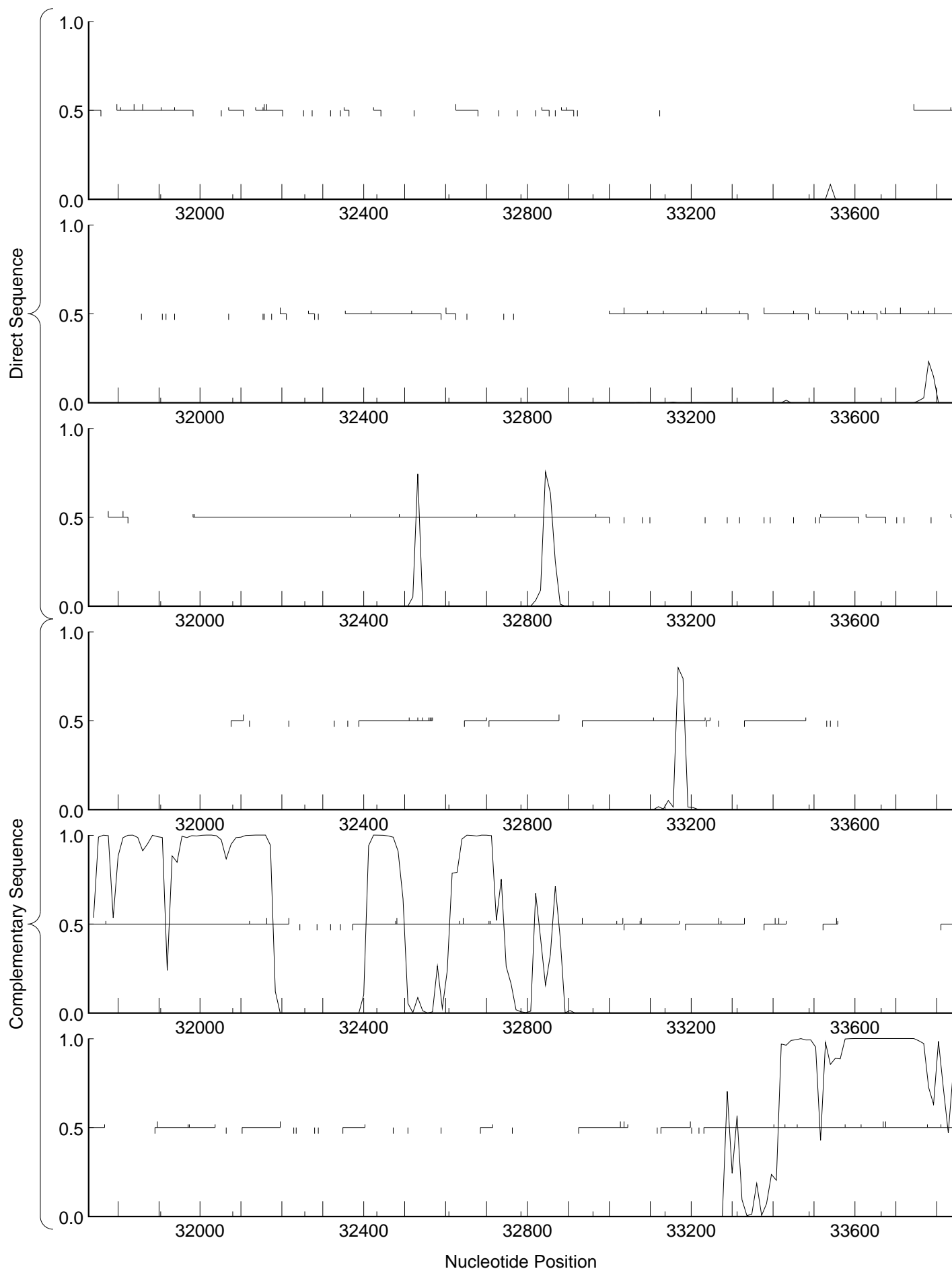


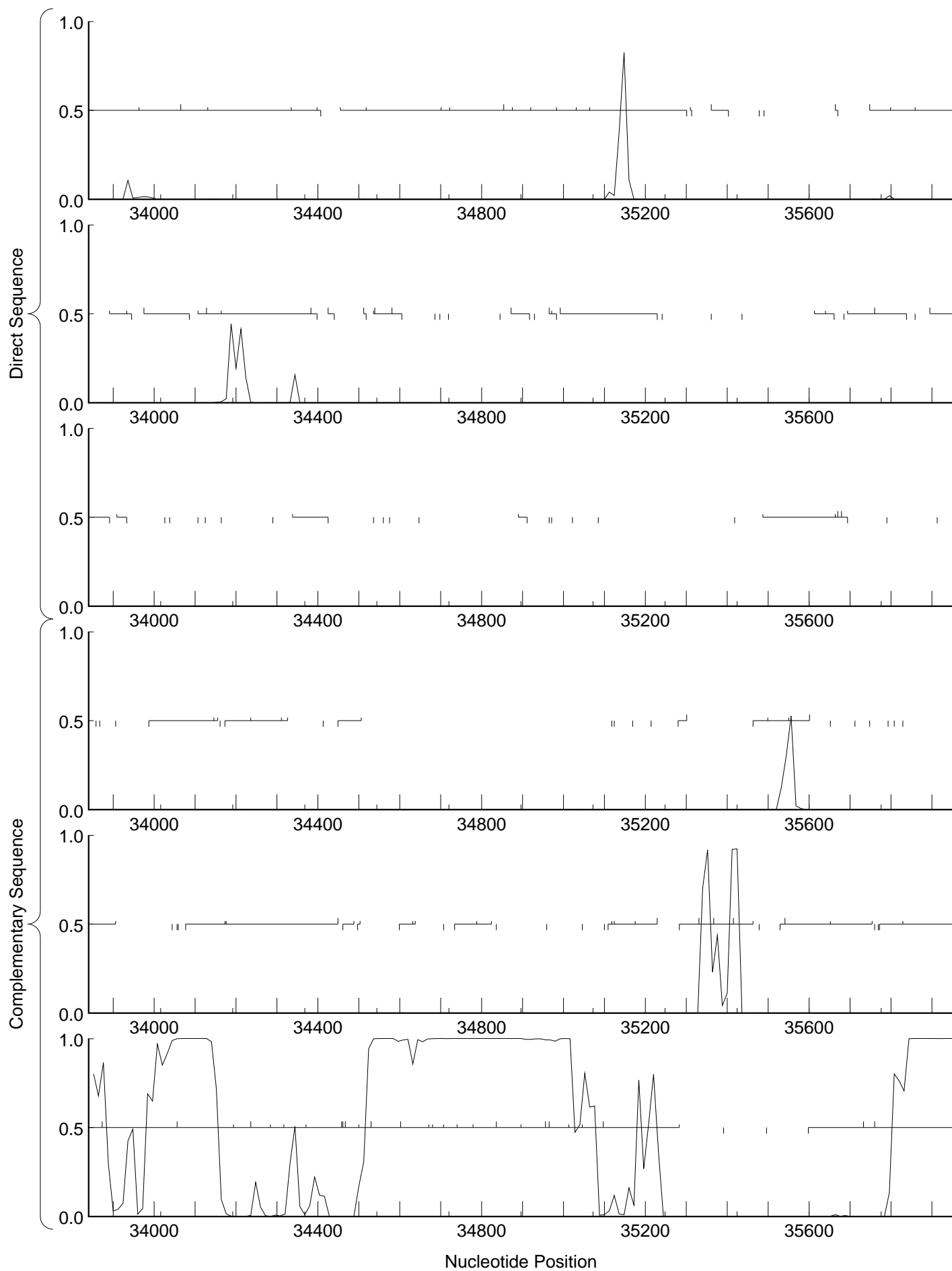
complete sequence, 52499 bp including 10-base 3' overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 15/26

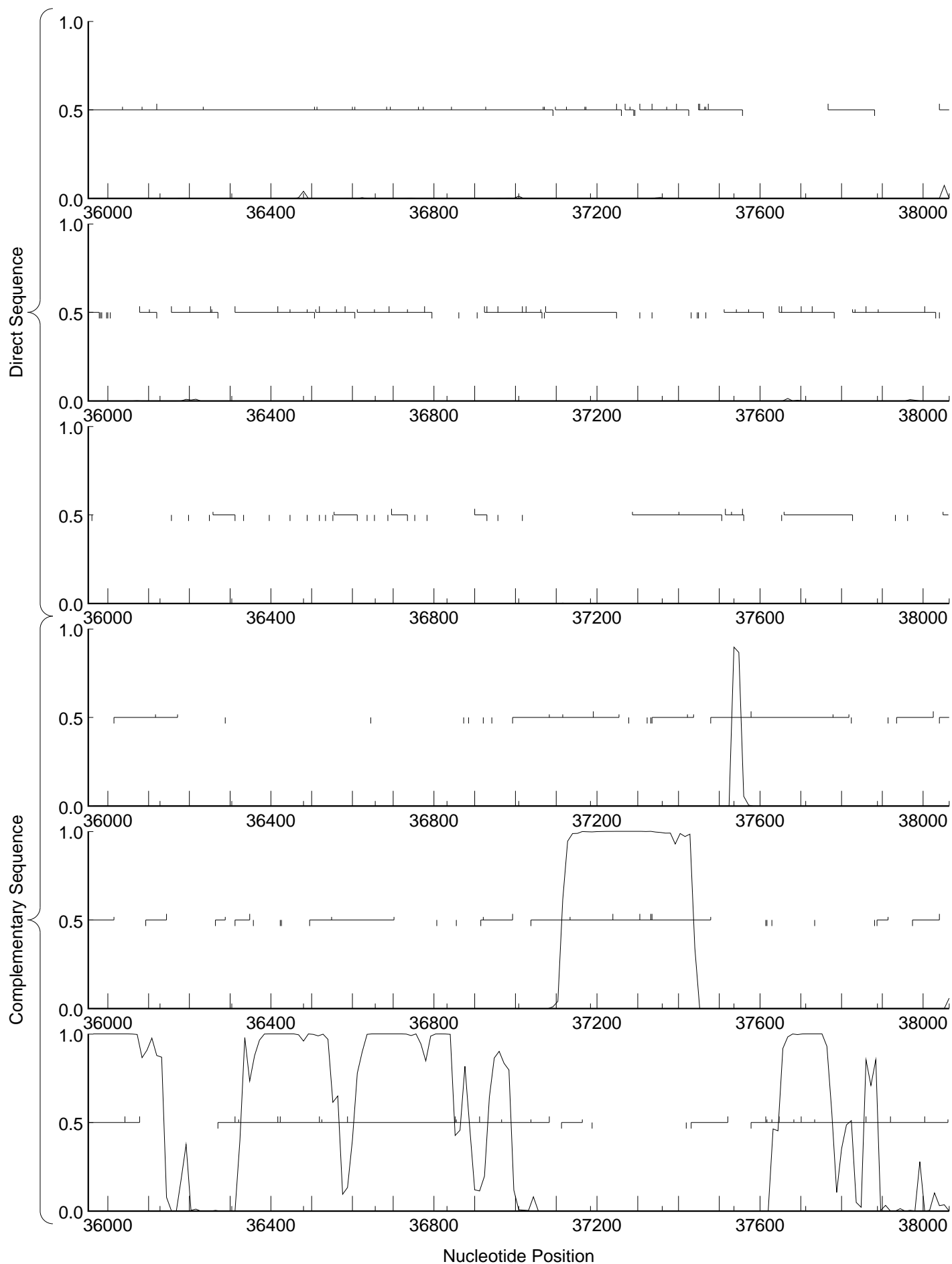


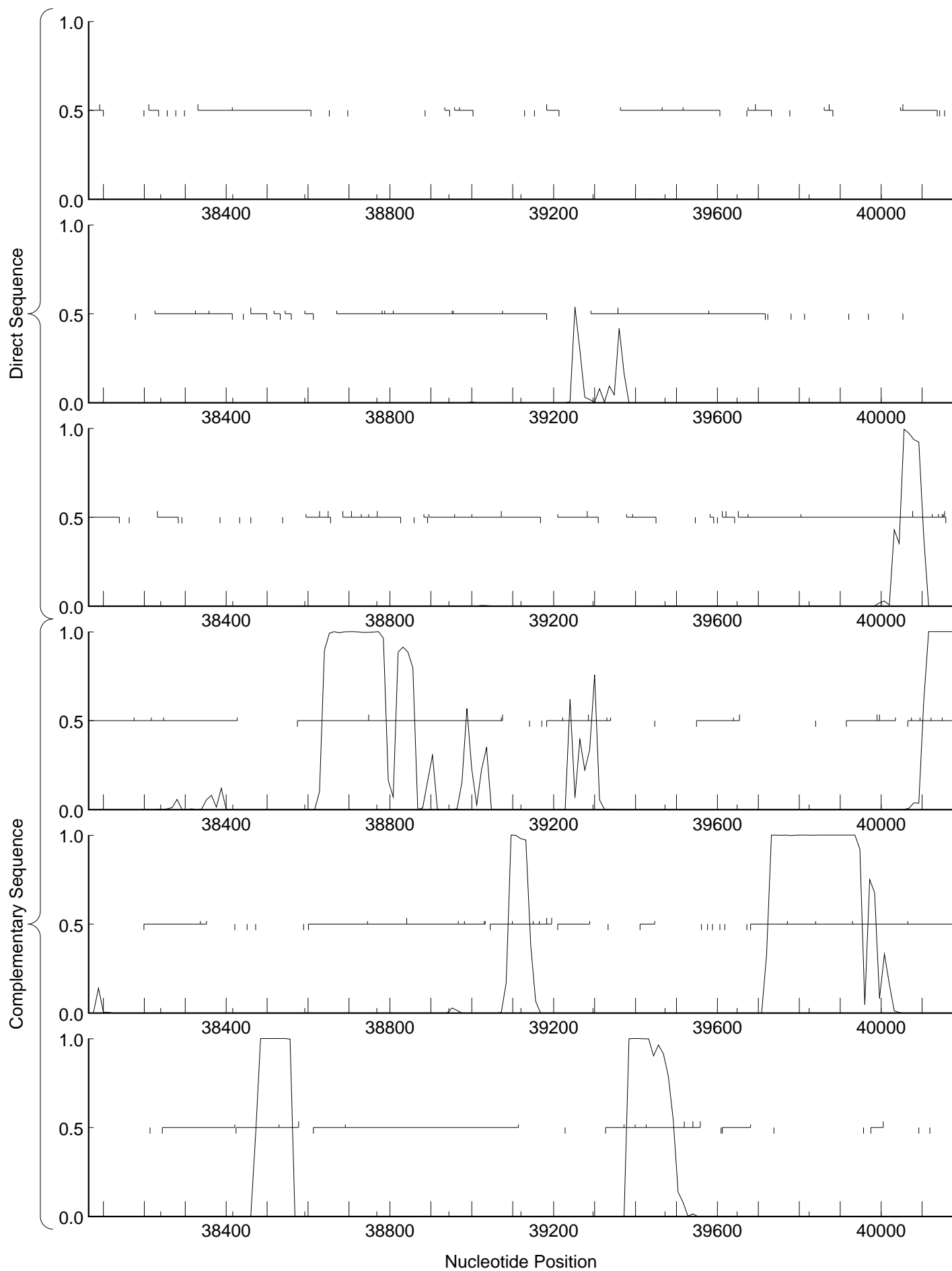


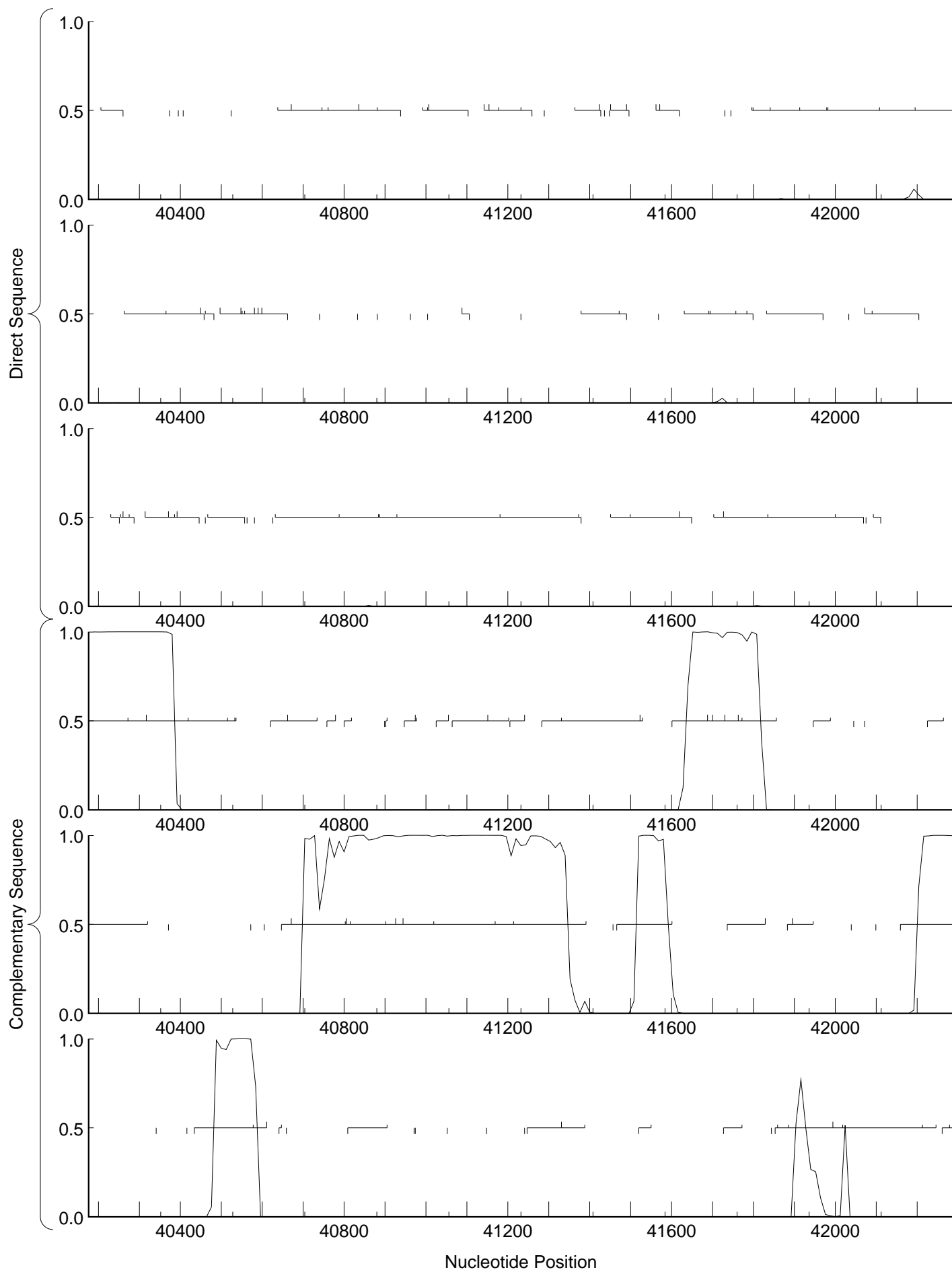
complete sequence, 52499 bp including 10-base 3' overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 17/26

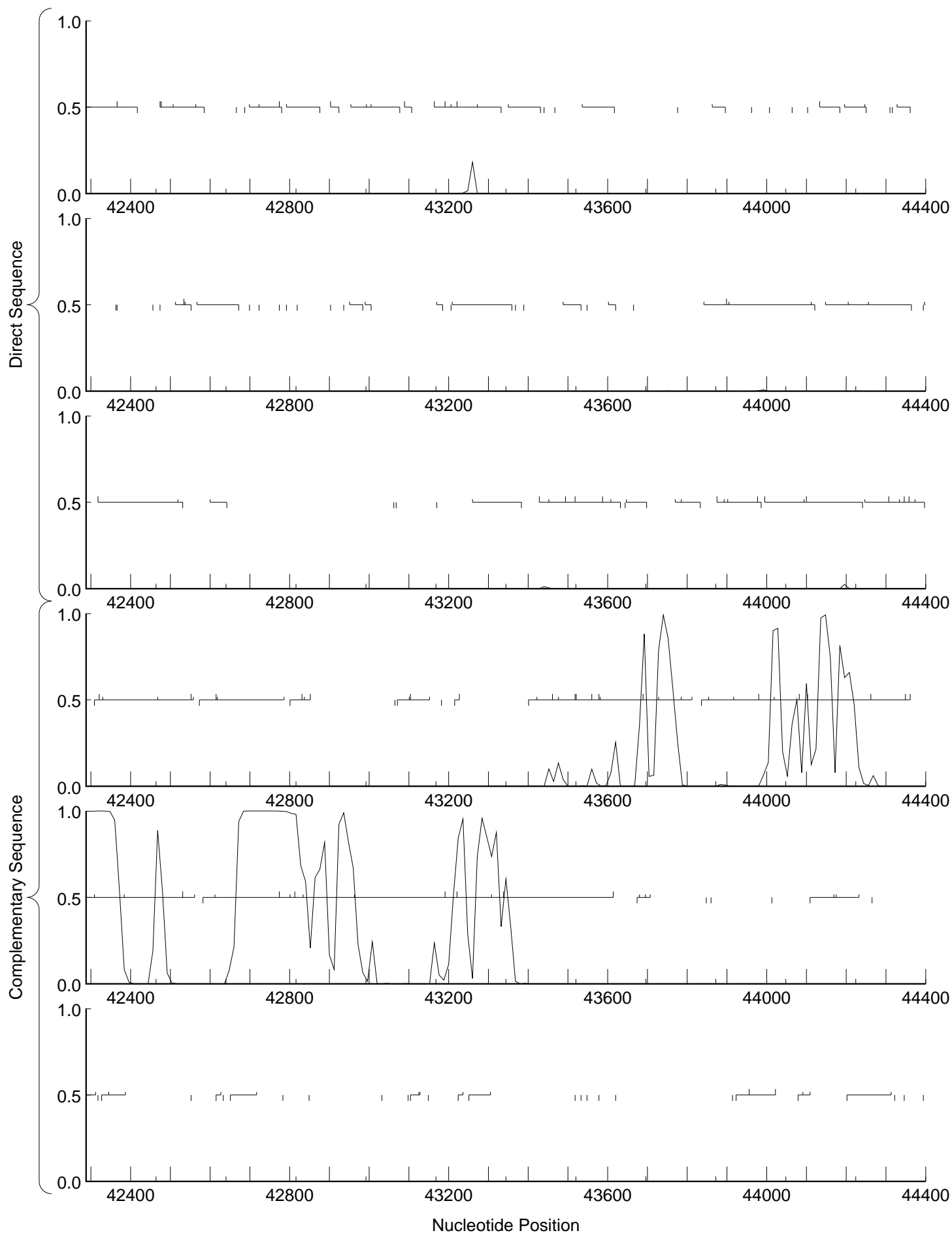












complete sequence, 52499 bp including 10-base 3' overhang (CGGTCGGTAA), Cluster A6, Order 4, Window 96, Step 12, 23/26

