

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Rhodococcus phage Shuman complete sequence, 46544 bp including 10-base 3' overhang (CGGCCGT
Analysis Date : 3/17/18 at 23:52:41
Pages : 24
Sequence Length : 46544 bp
GC Content : 58.65%

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,

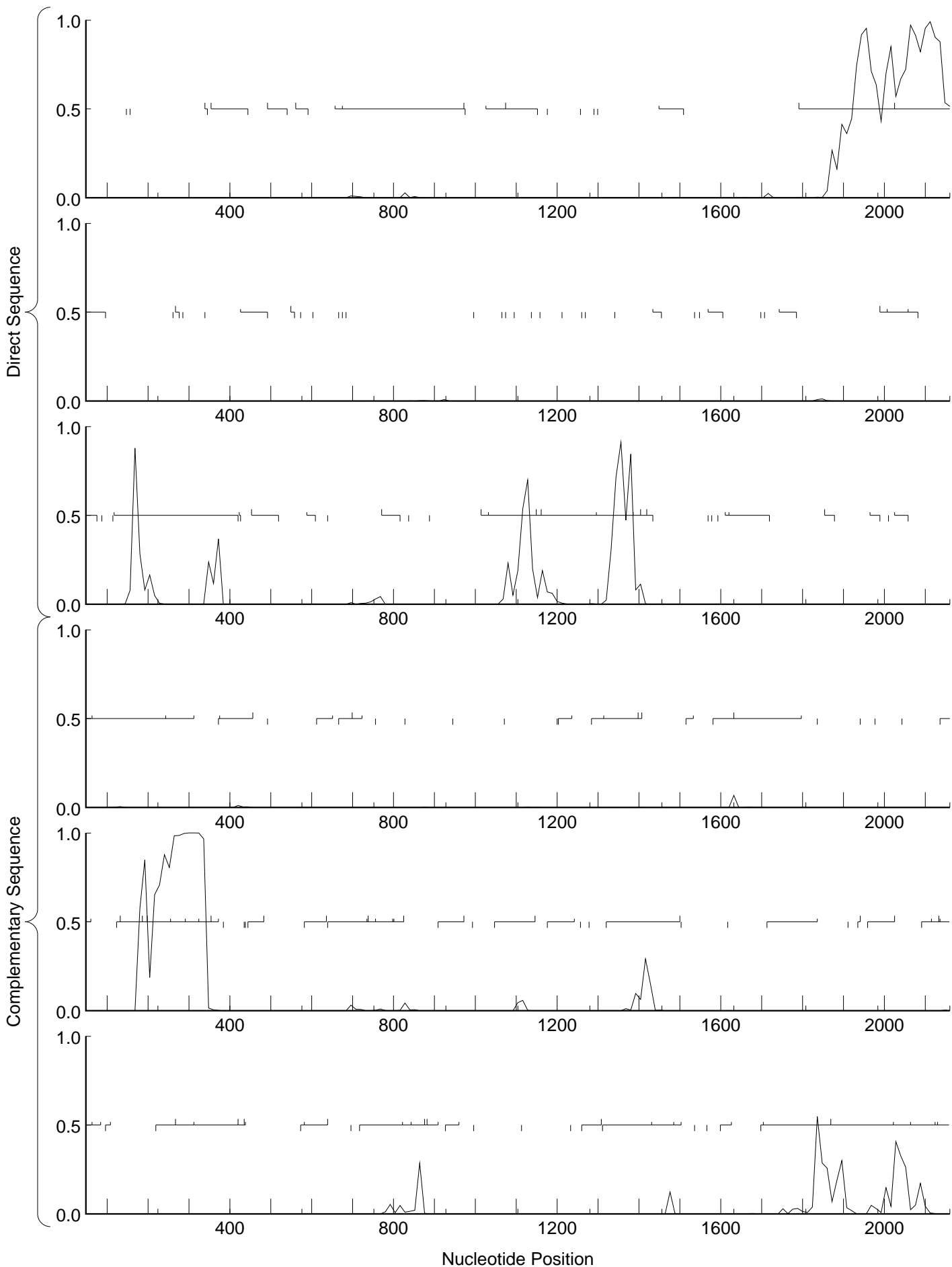
Matrix notes & comments

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

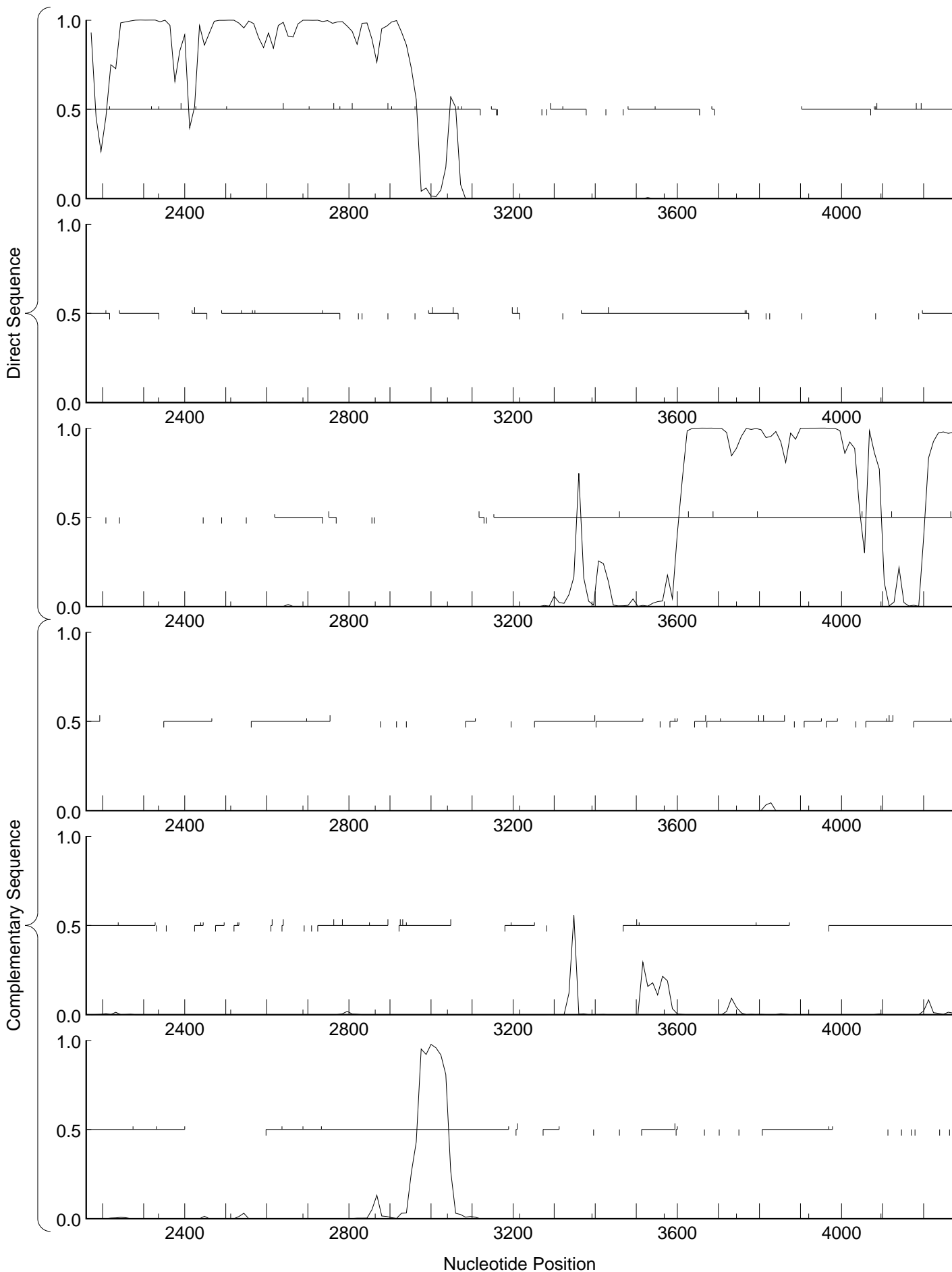
MATRIX INFORMATION

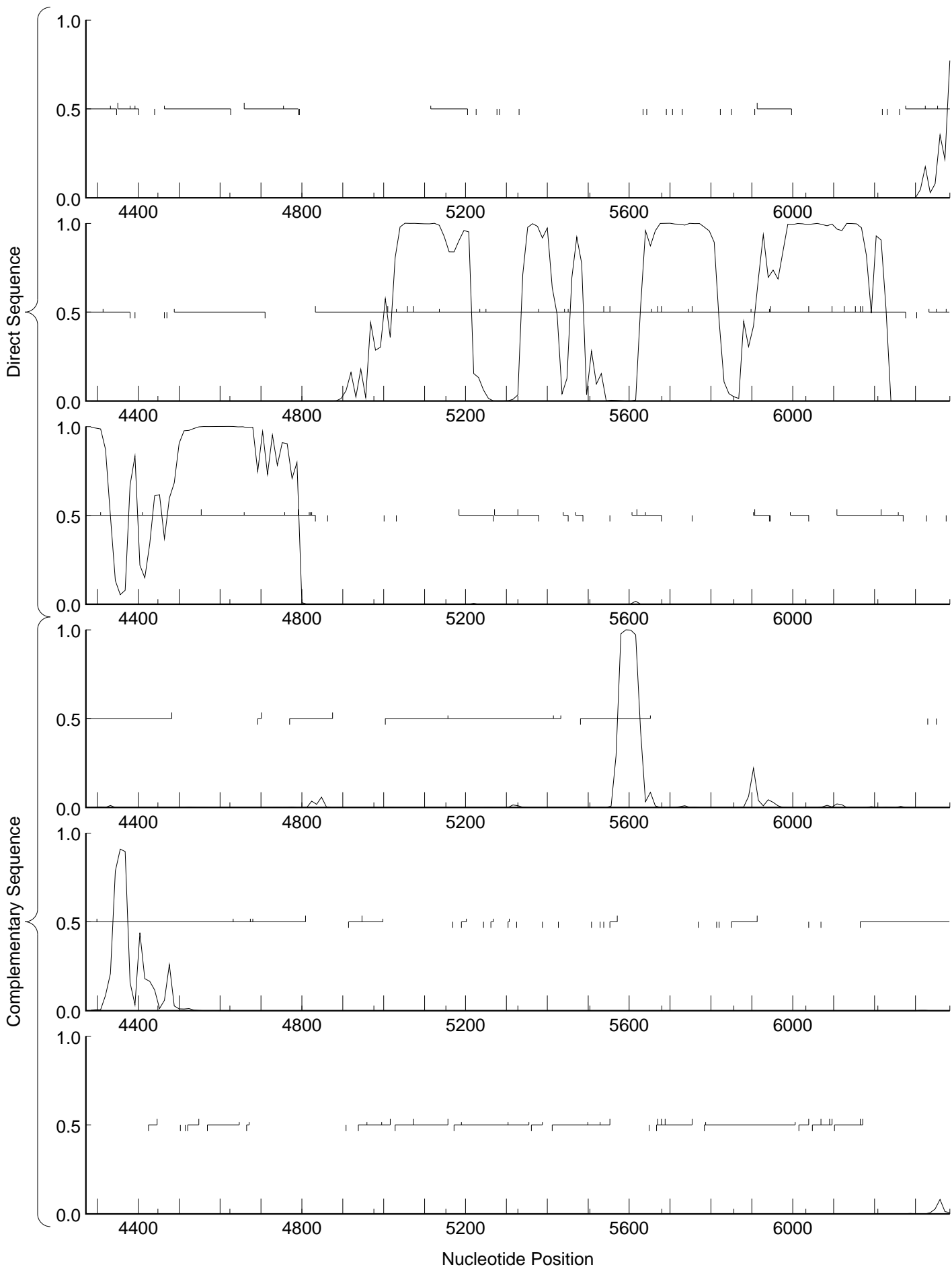
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

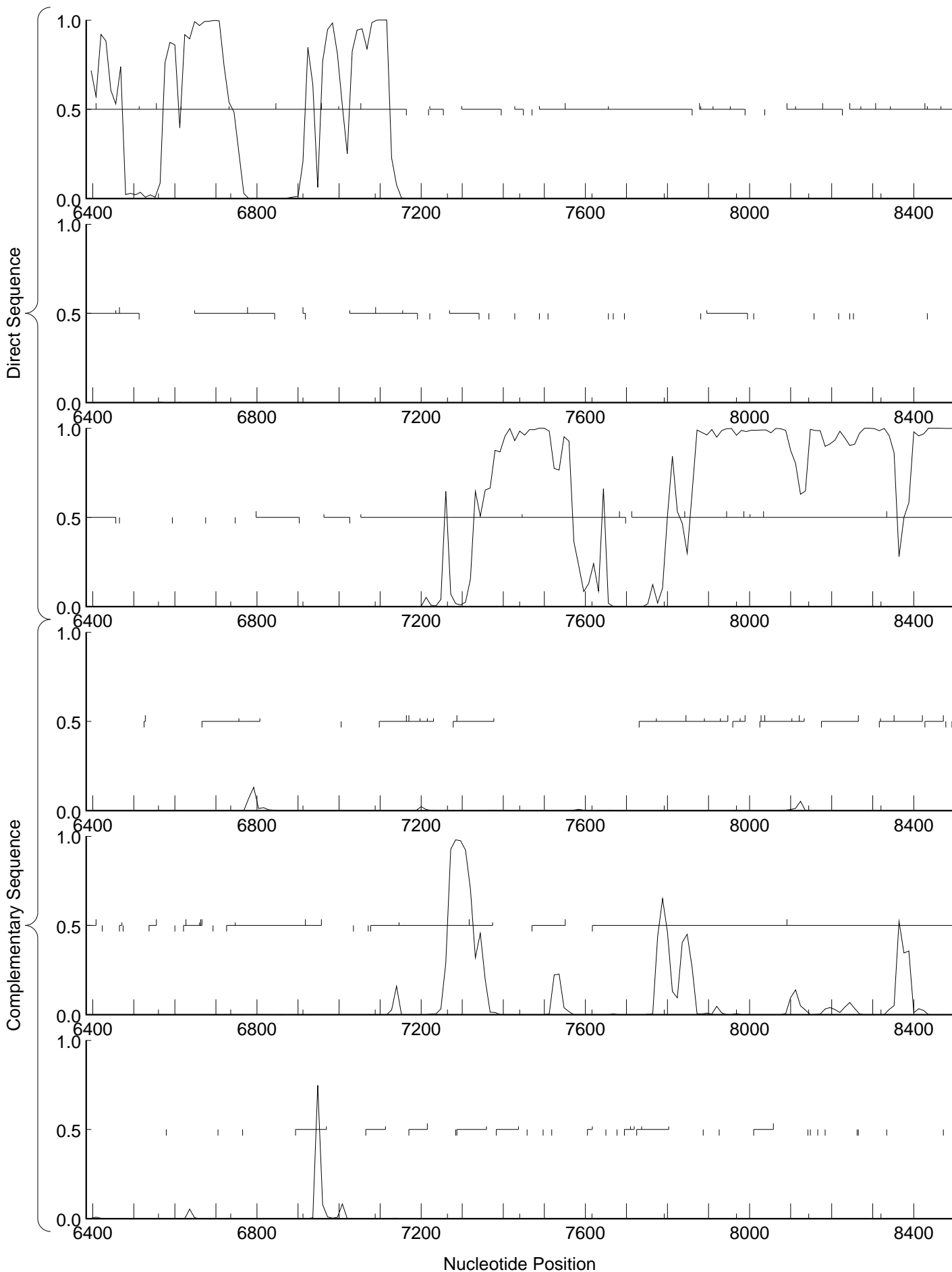
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

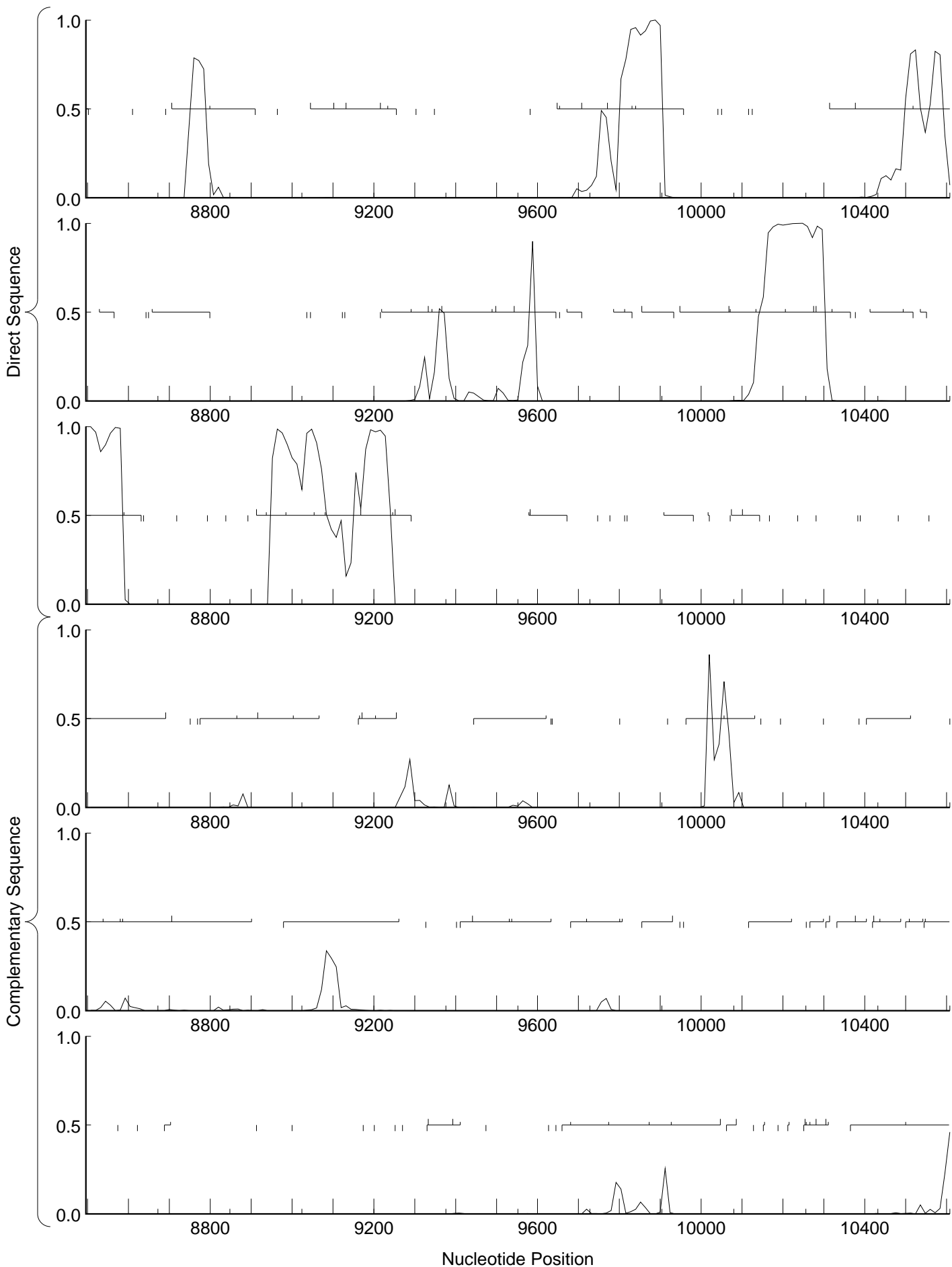


Complete sequence, 46544 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 3/24

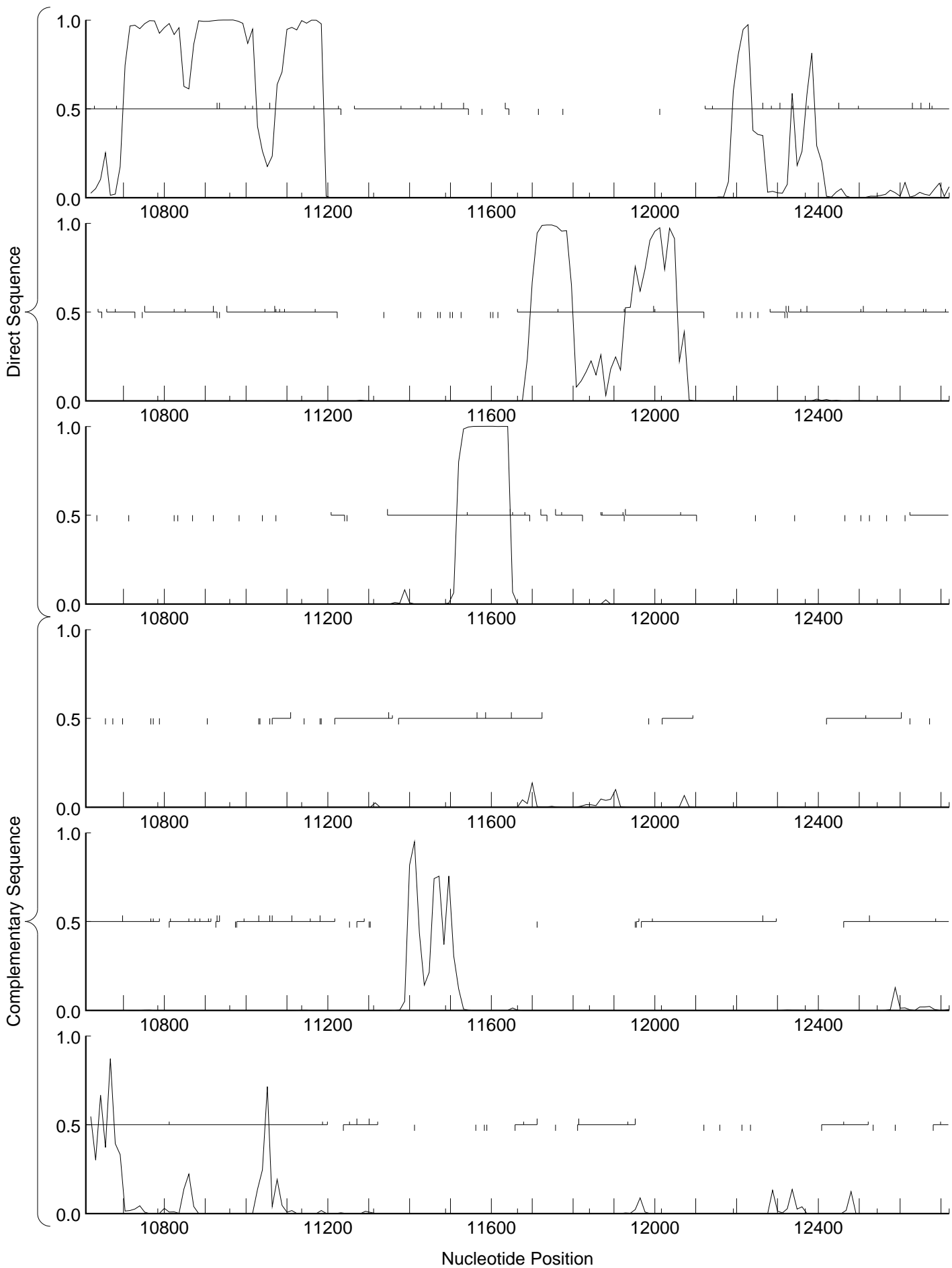


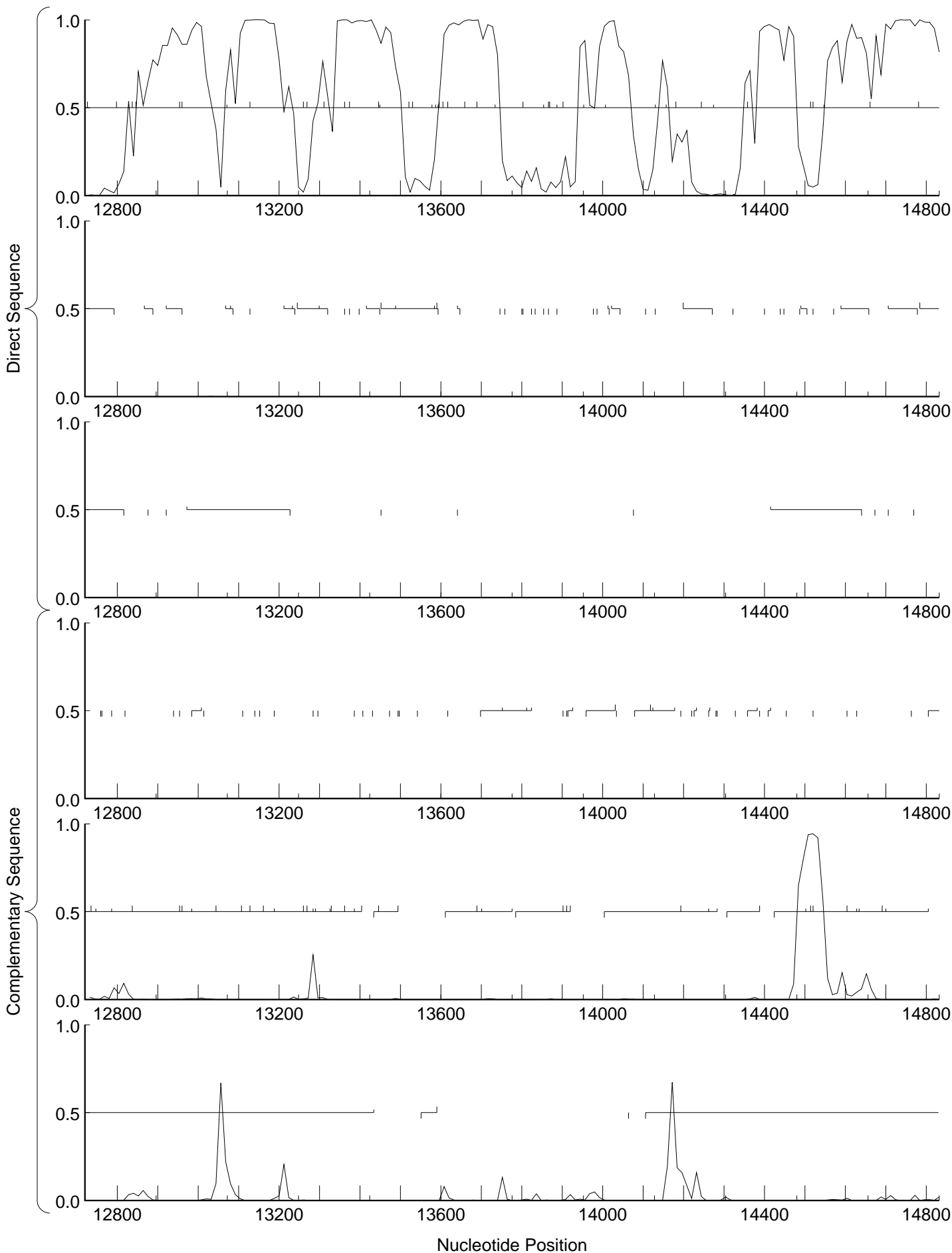


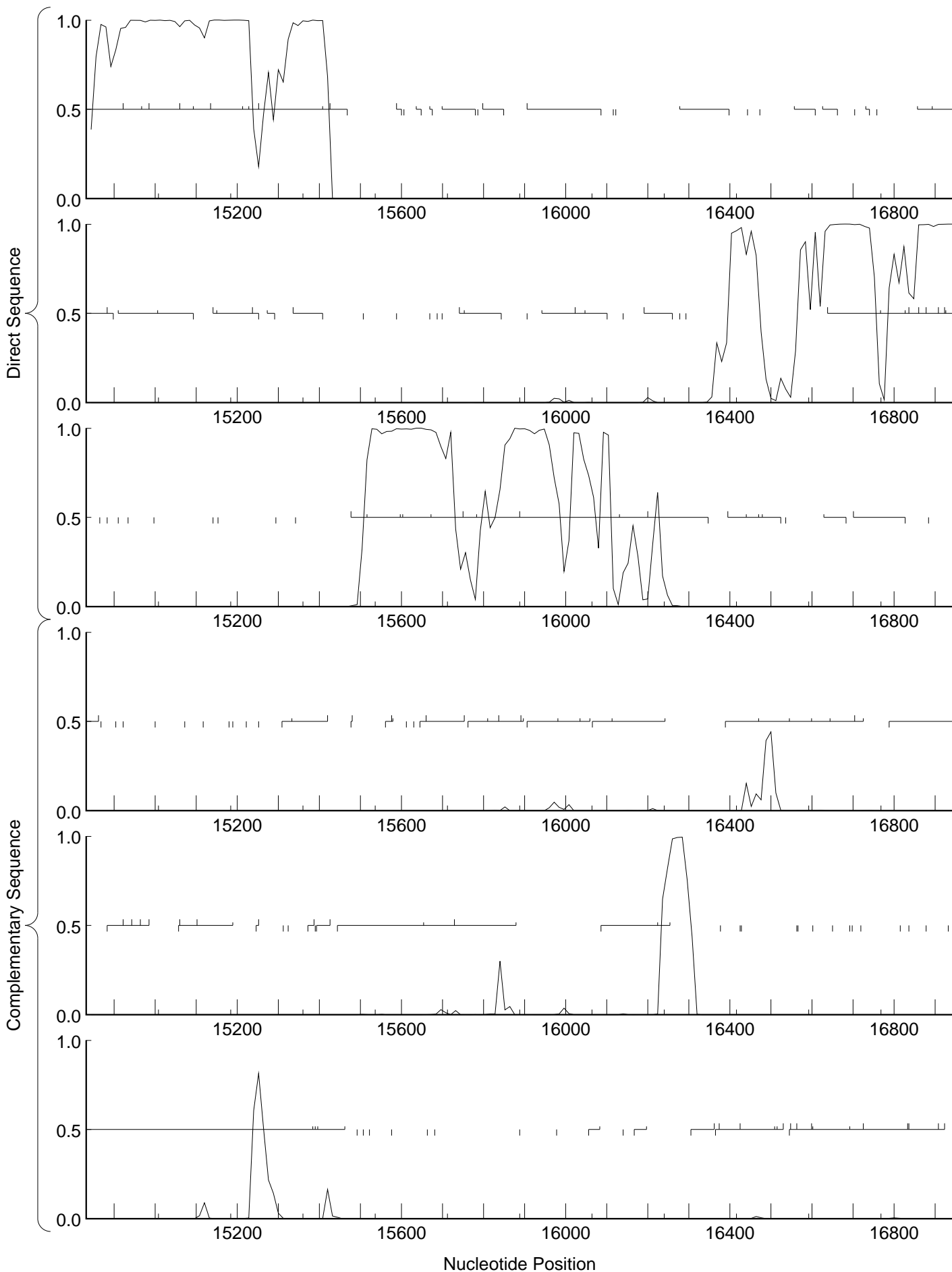


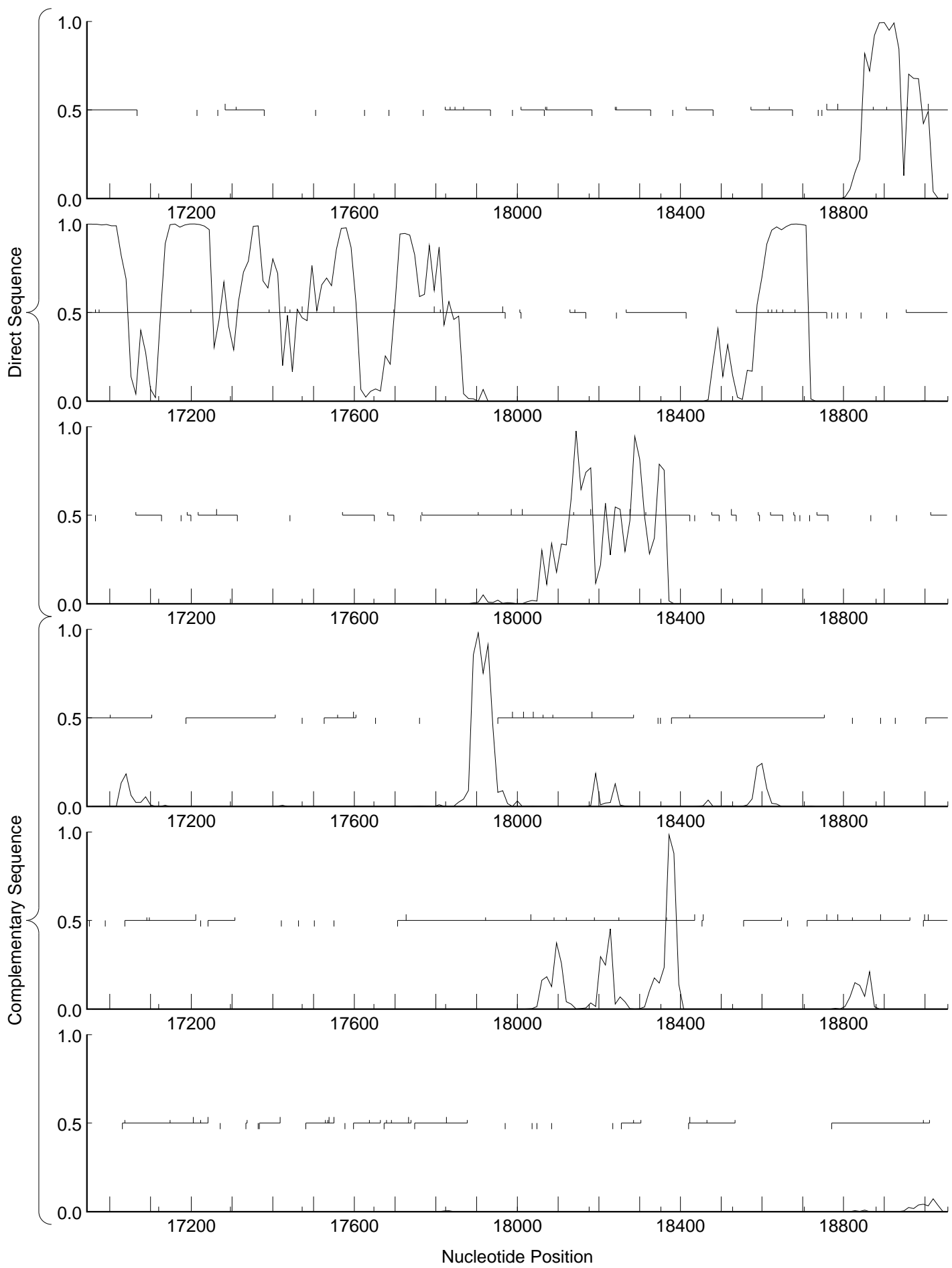


Complete sequence, 46544 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 7/24

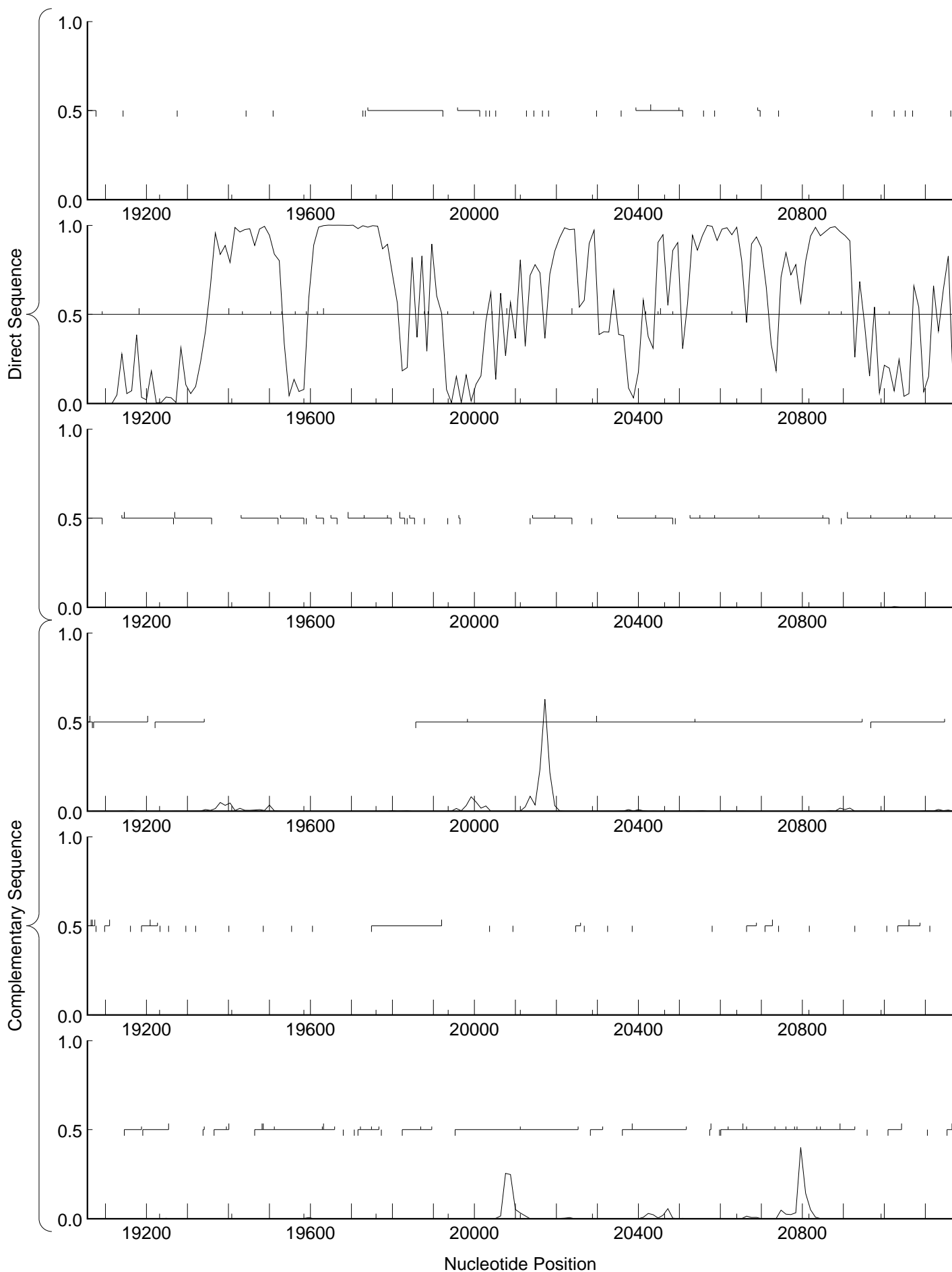




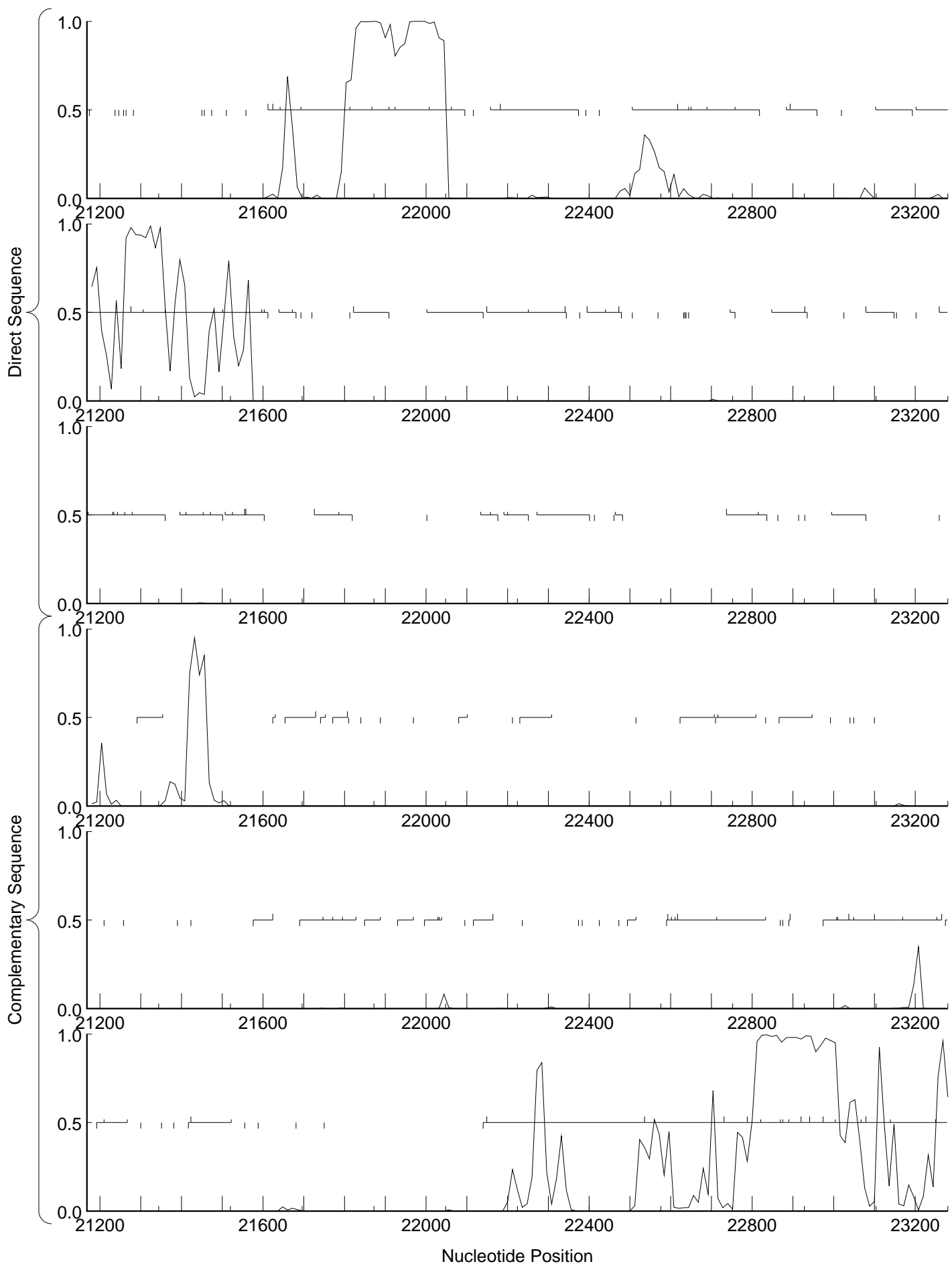




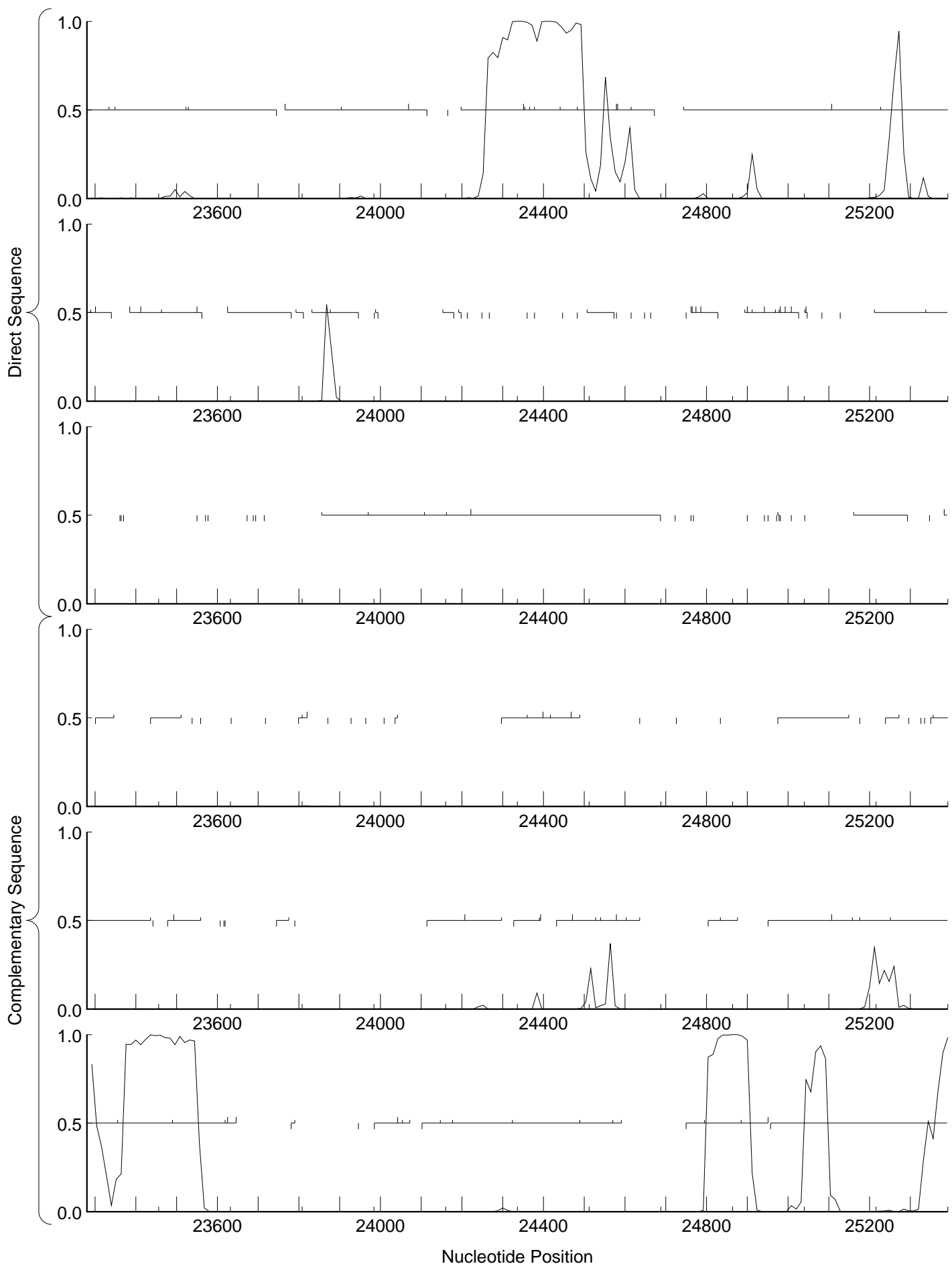
complete sequence, 46544 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 11/24



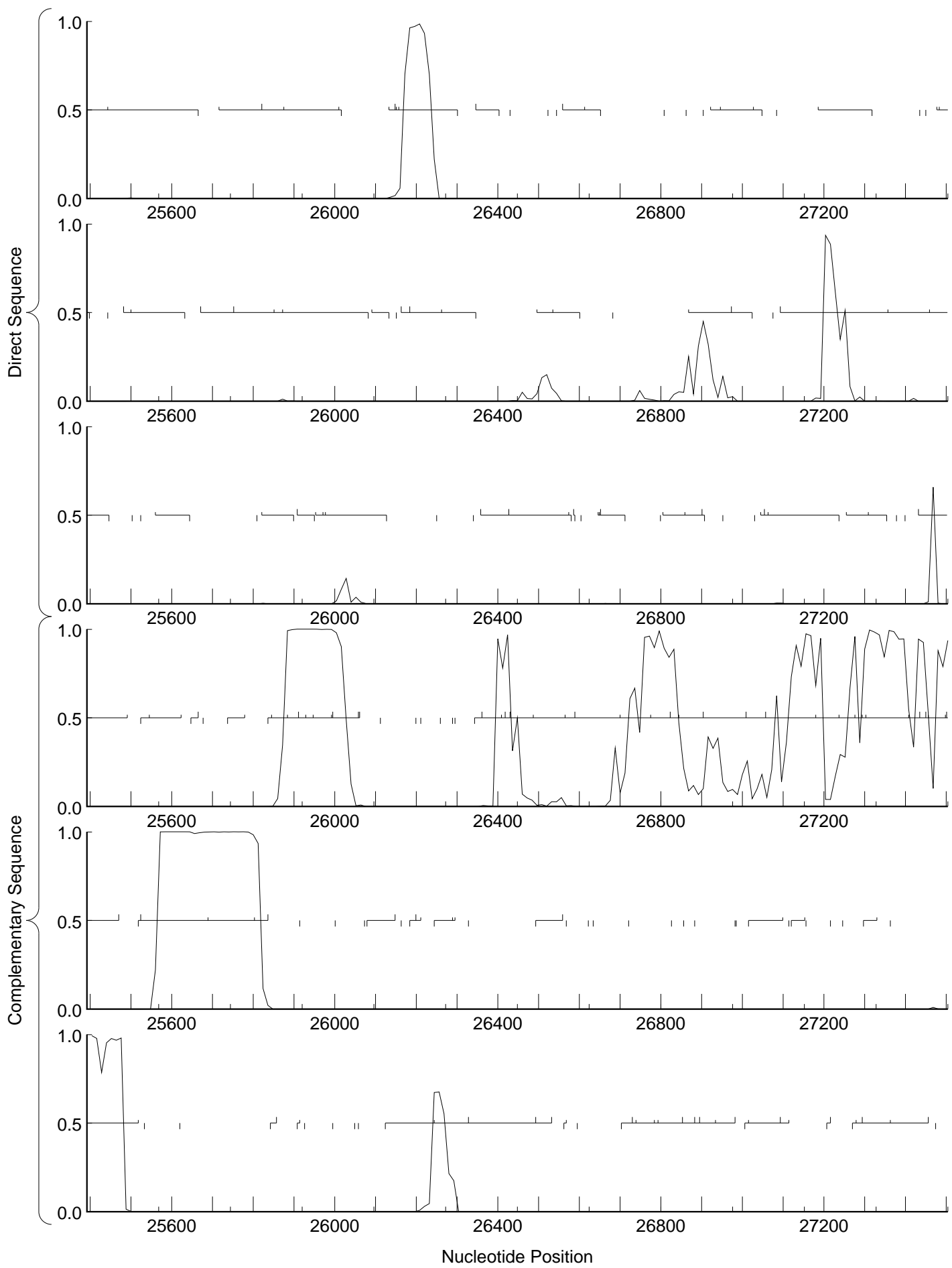
complete sequence, 46544 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 12/24



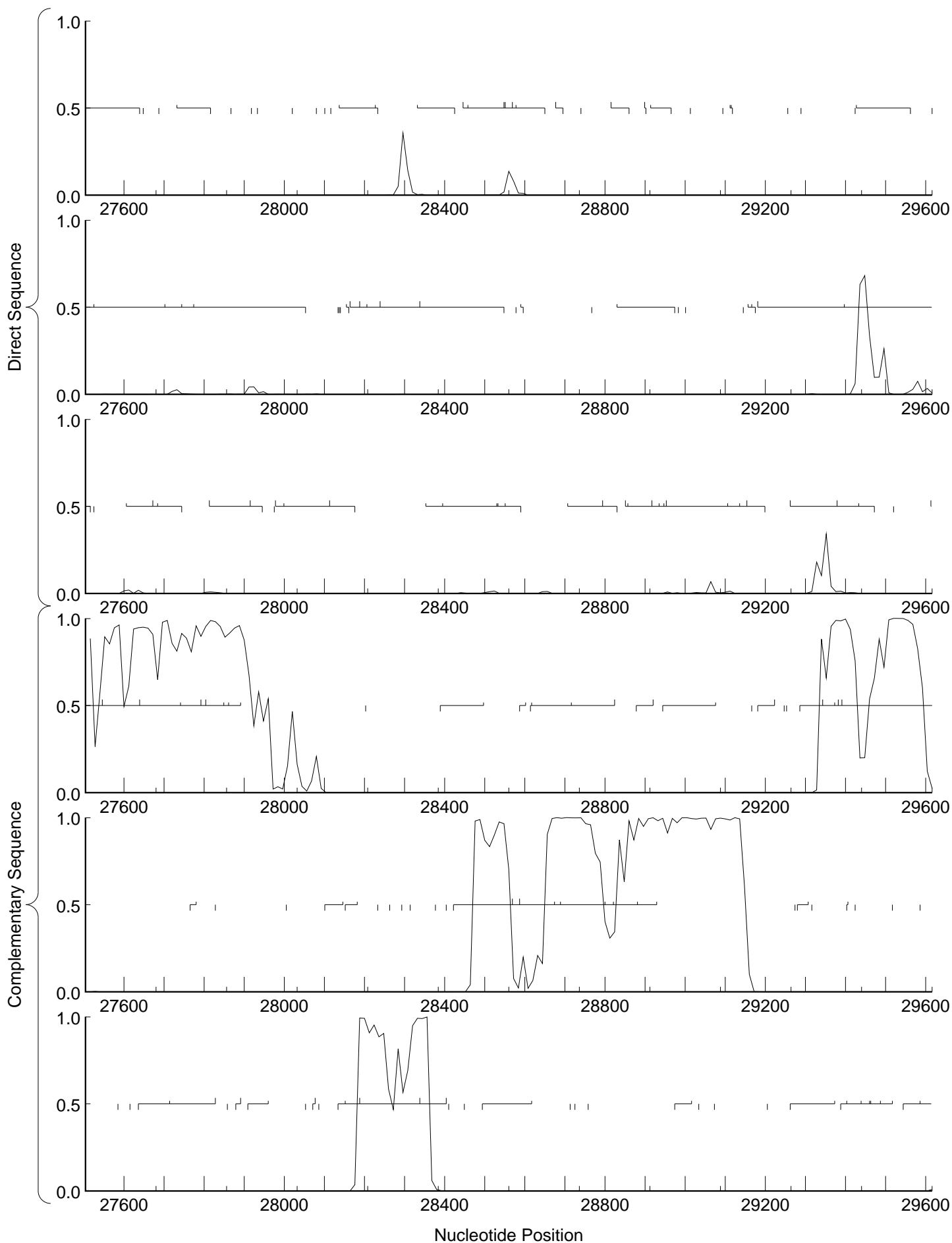
complete sequence, 46544 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 13/24



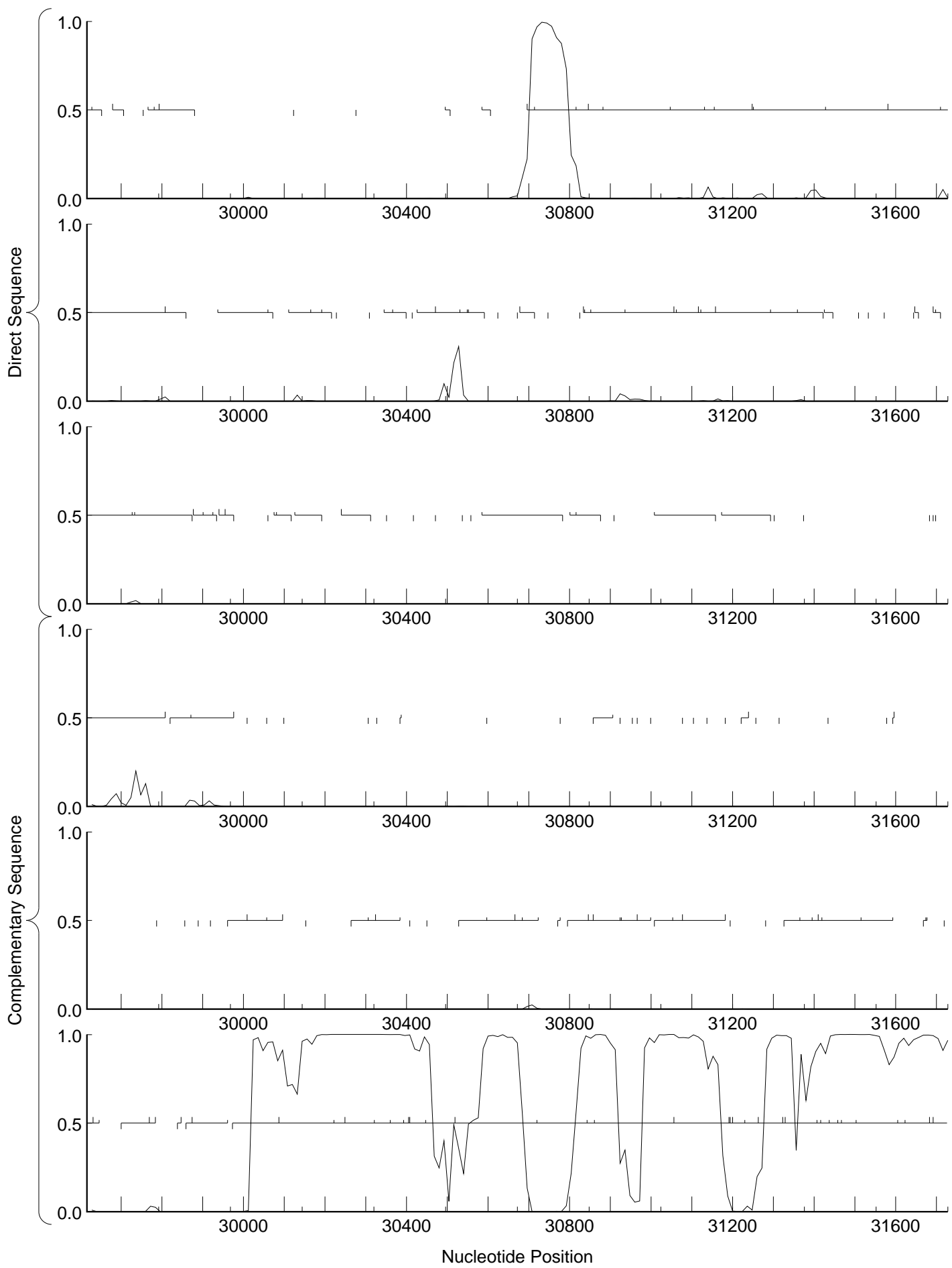
complete sequence, 46544 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 14/24



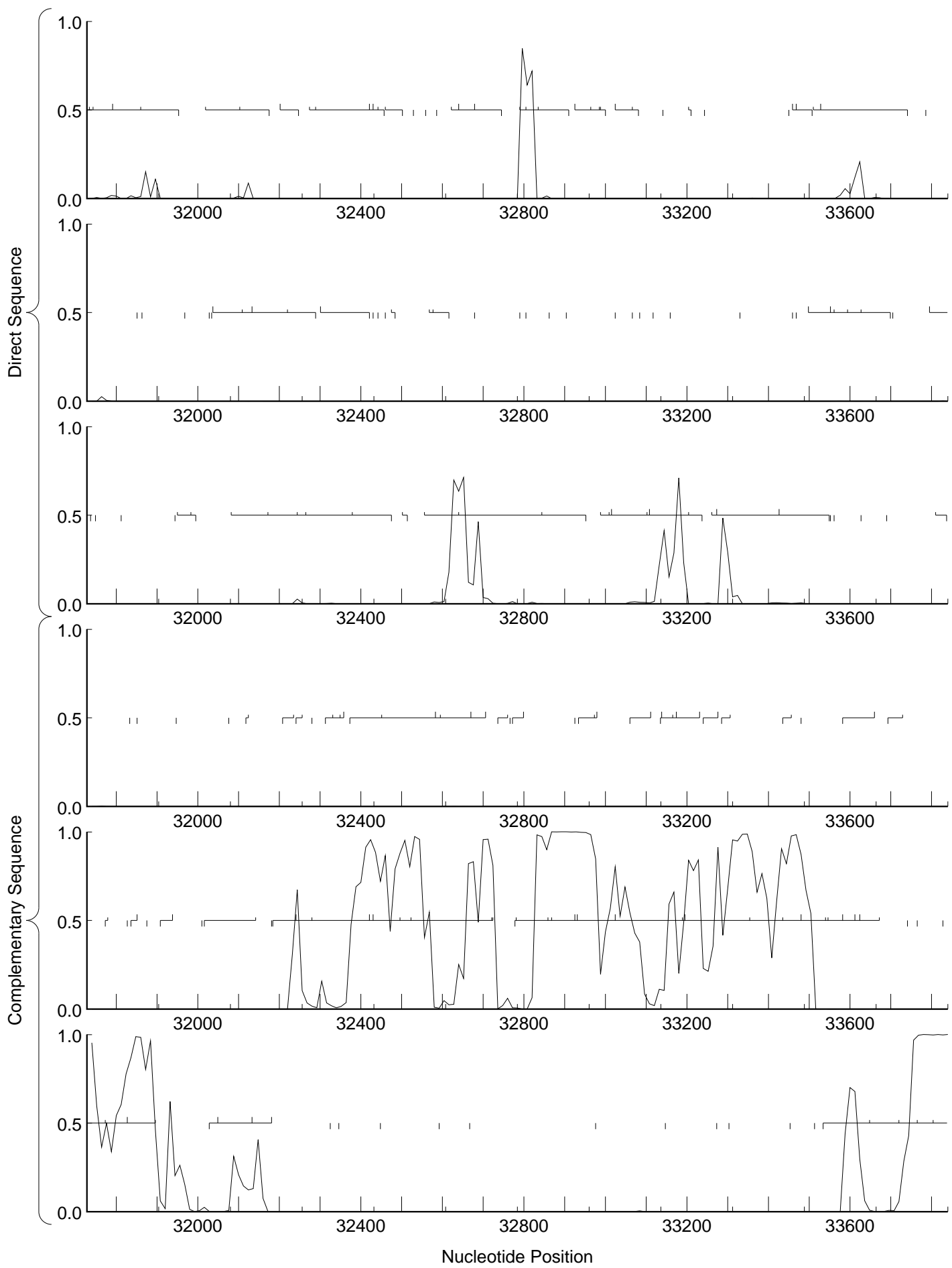
complete sequence, 46544 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 15/24

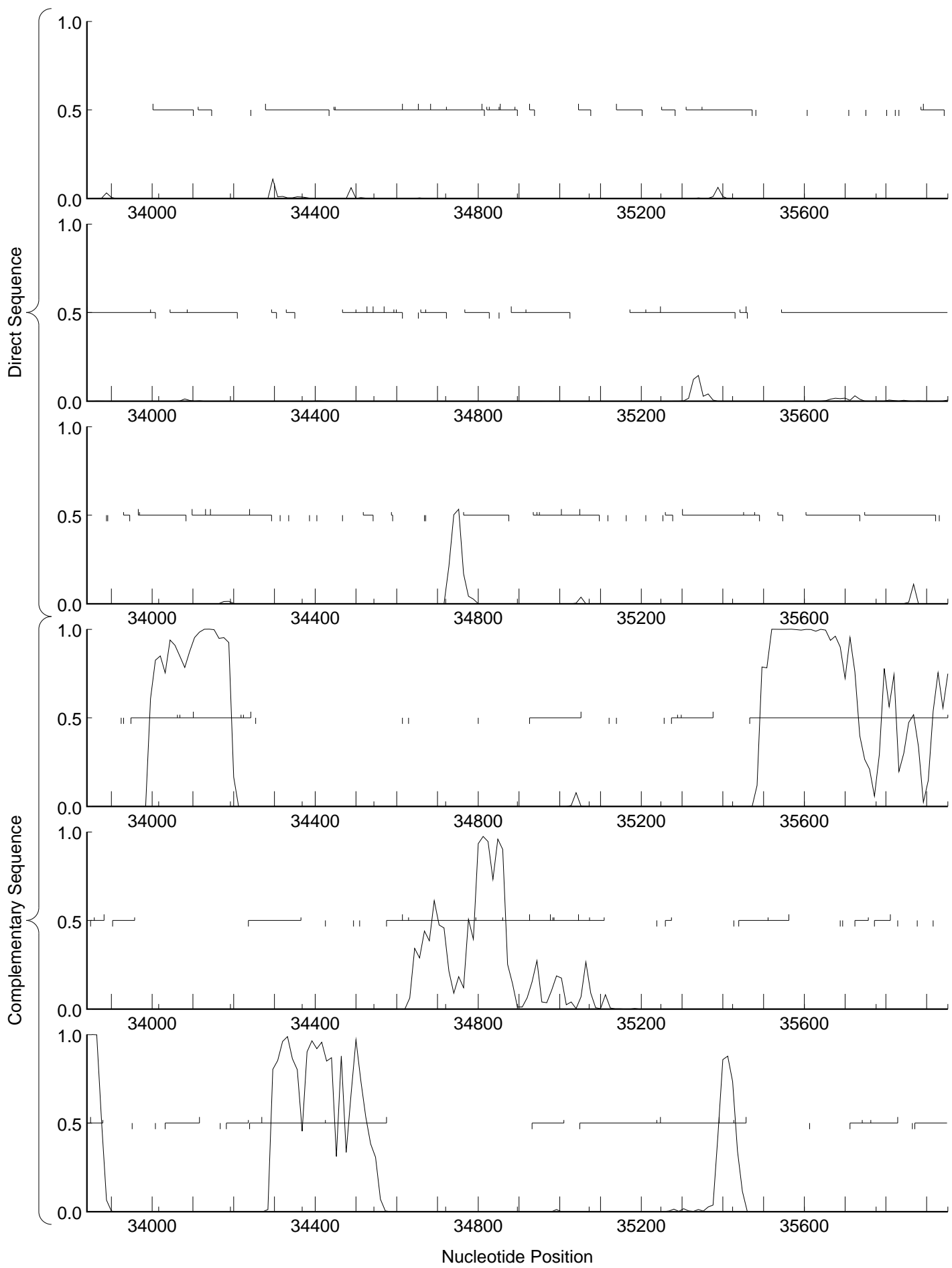


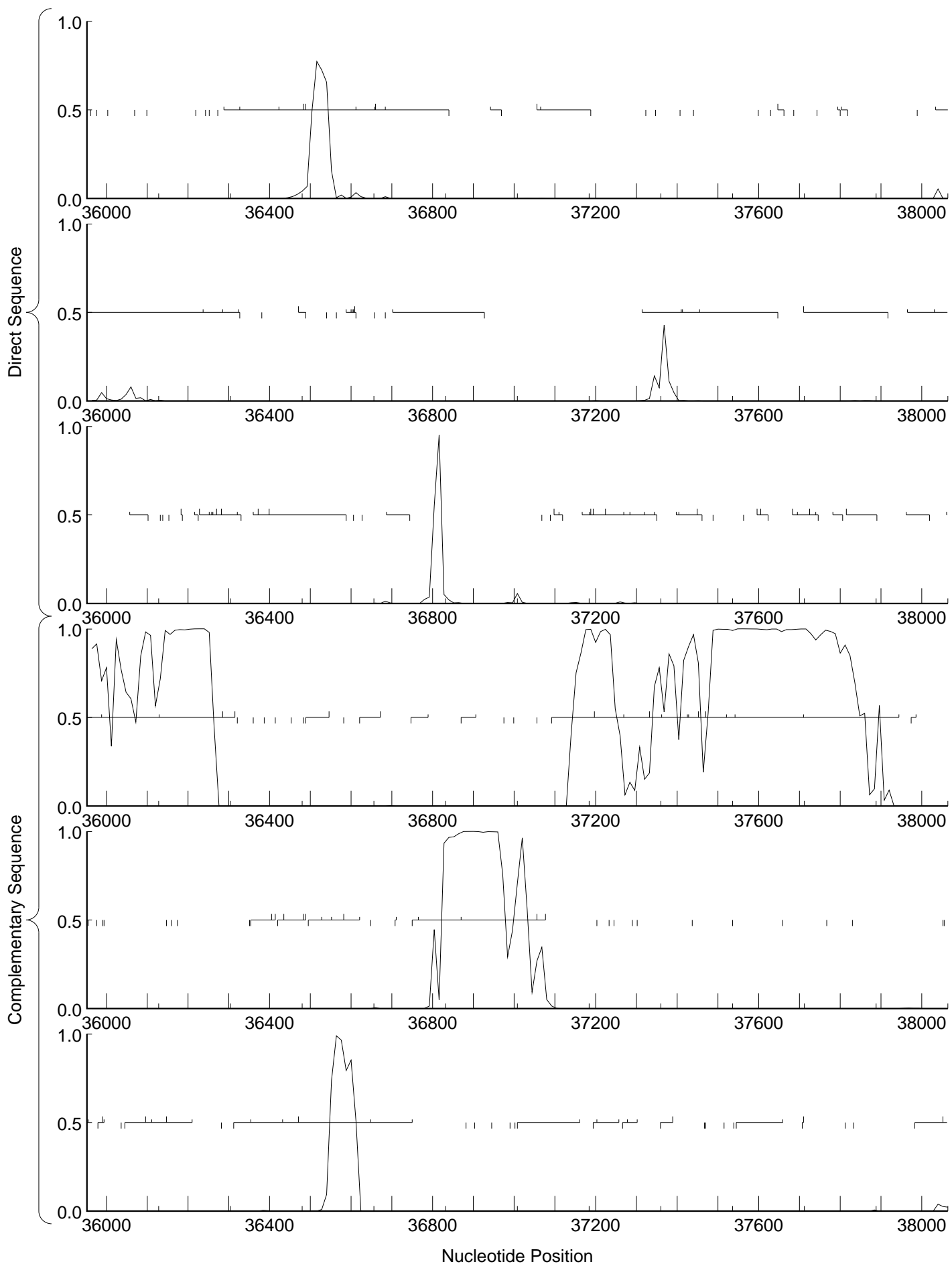
complete sequence, 46544 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 16/24

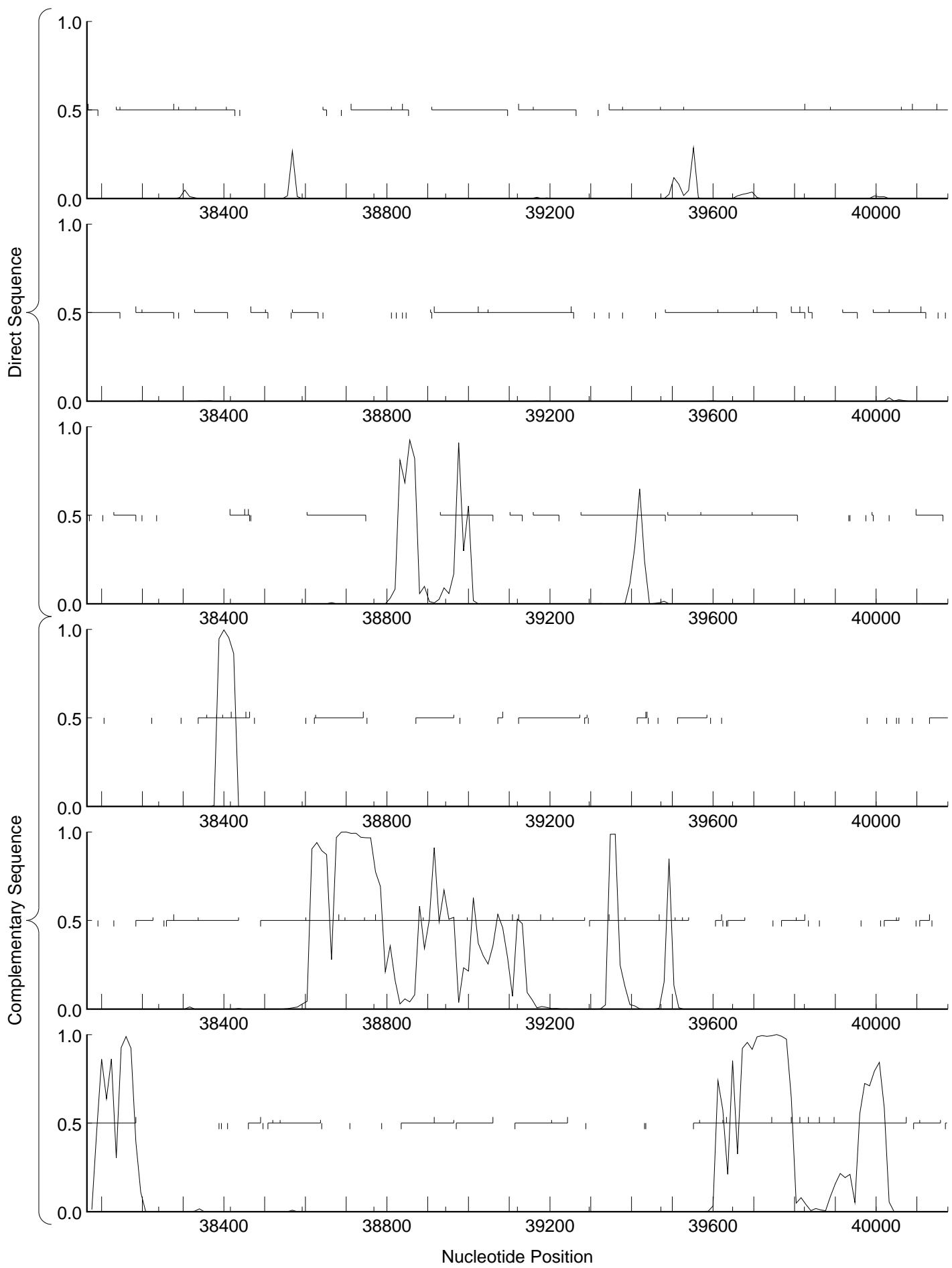


complete sequence, 46544 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 17/24

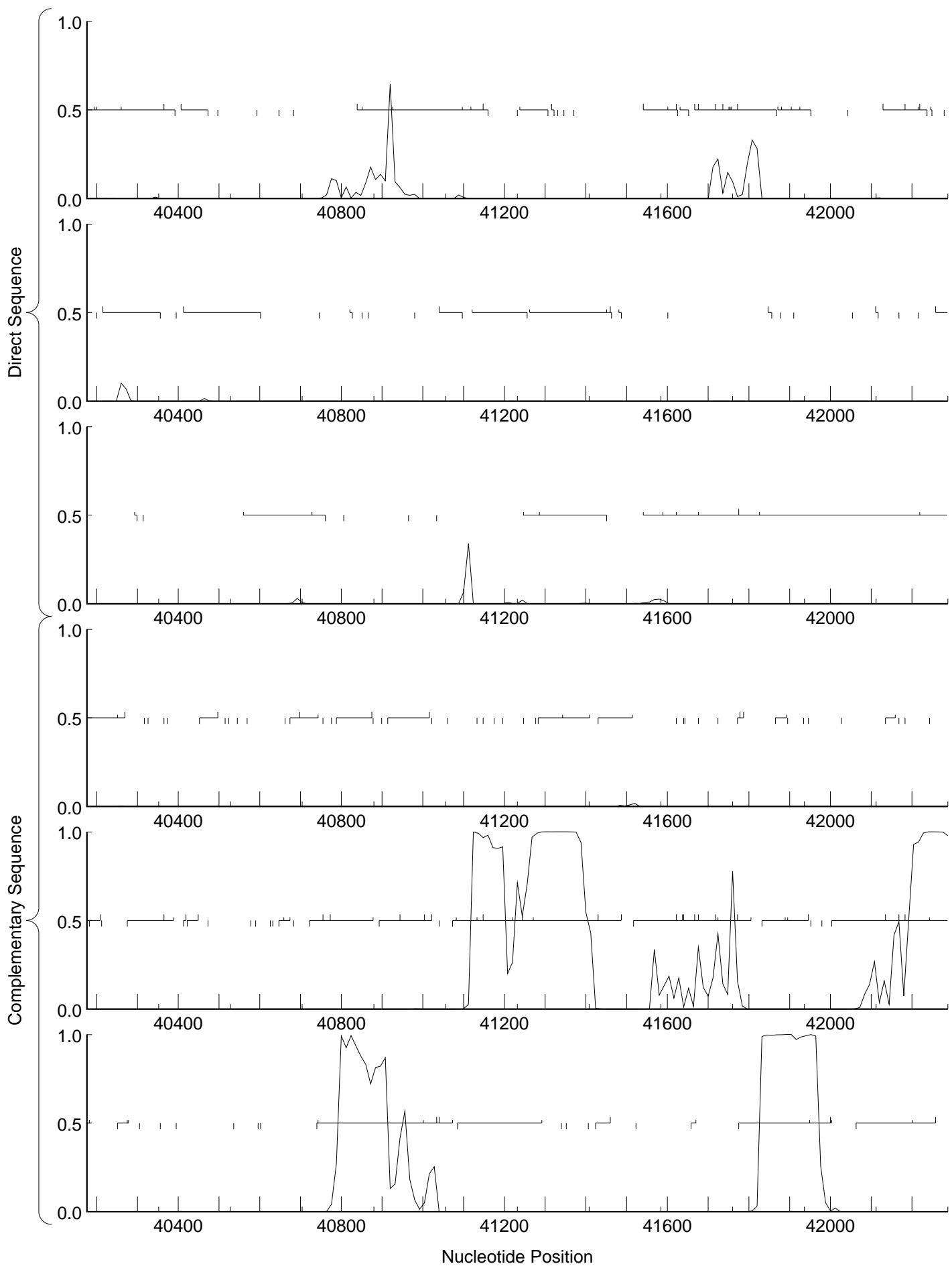


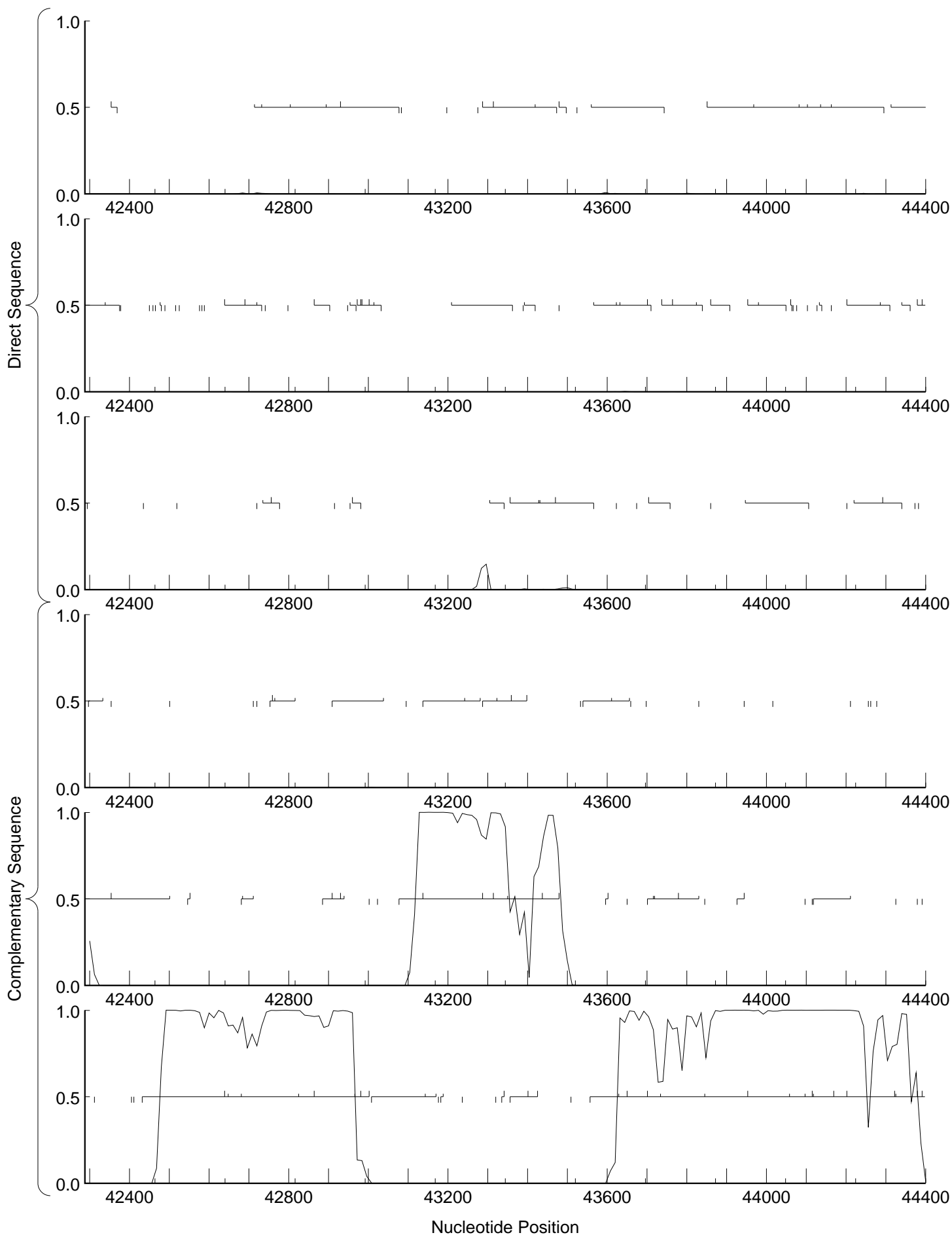






complete sequence, 46544 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 21/24





complete sequence, 46544 bp including 10-base 3' overhang (CGGCCGTGAT), Cluster CA, Order 4, Window 96, Step 12, 23/24

