

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Estes complete sequence, 83601 bp including 11-base 3' overhang (ACCCCA
Analysis Date : 5/25/22 at 1:51:22
Pages : 41
Sequence Length : 83601 bp
GC Content : 60.87%

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,

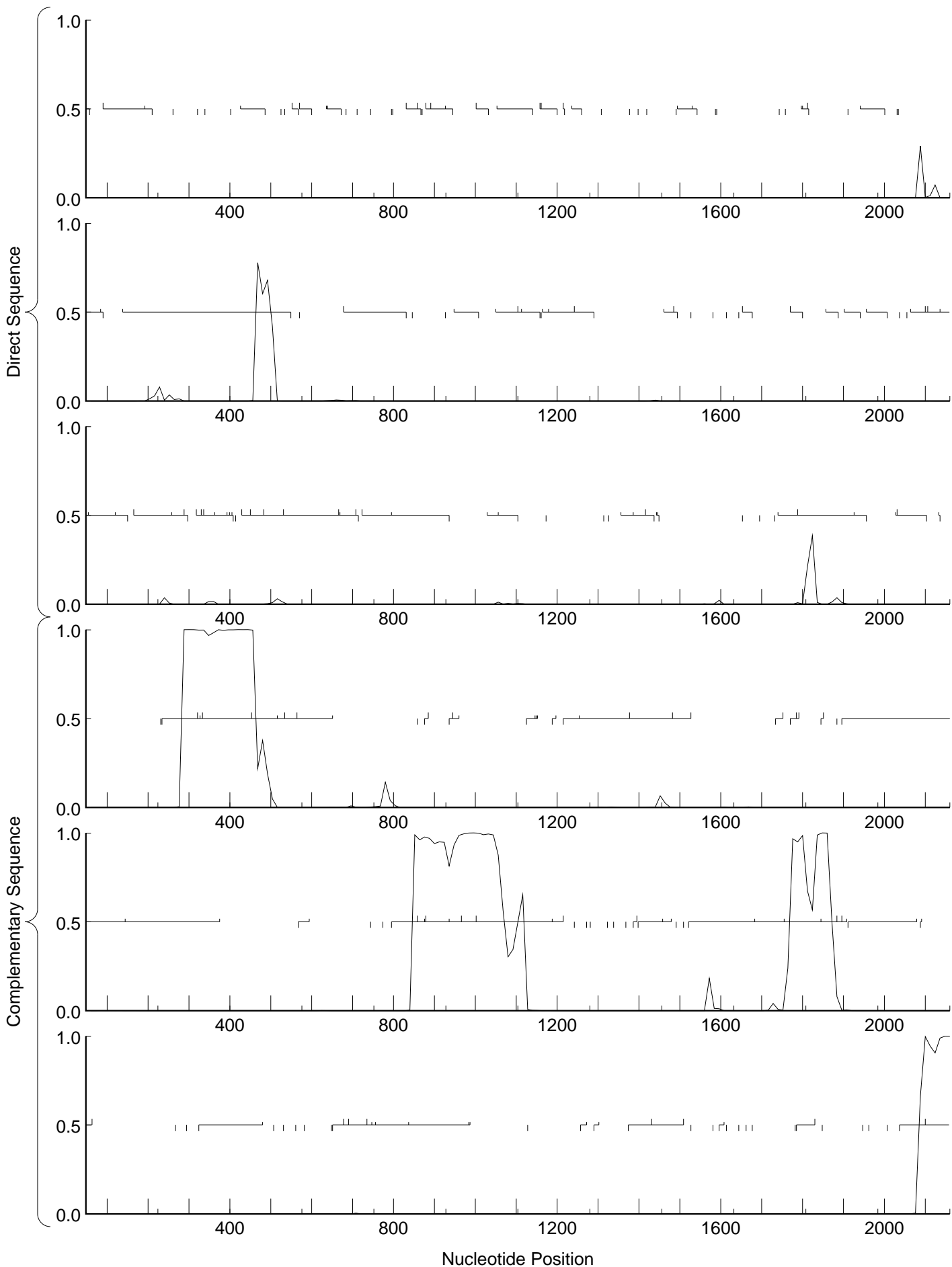
Matrix notes & comments

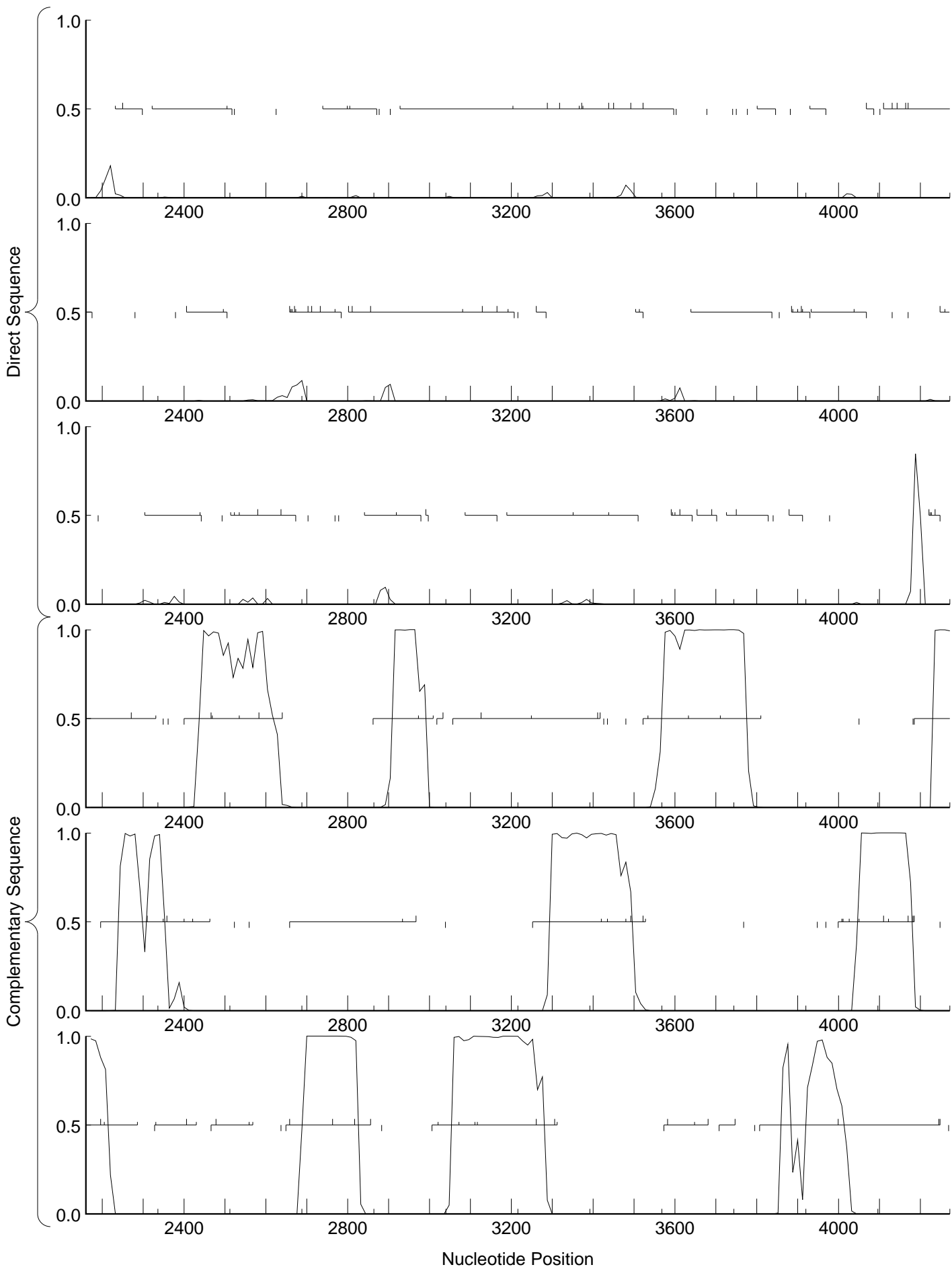
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

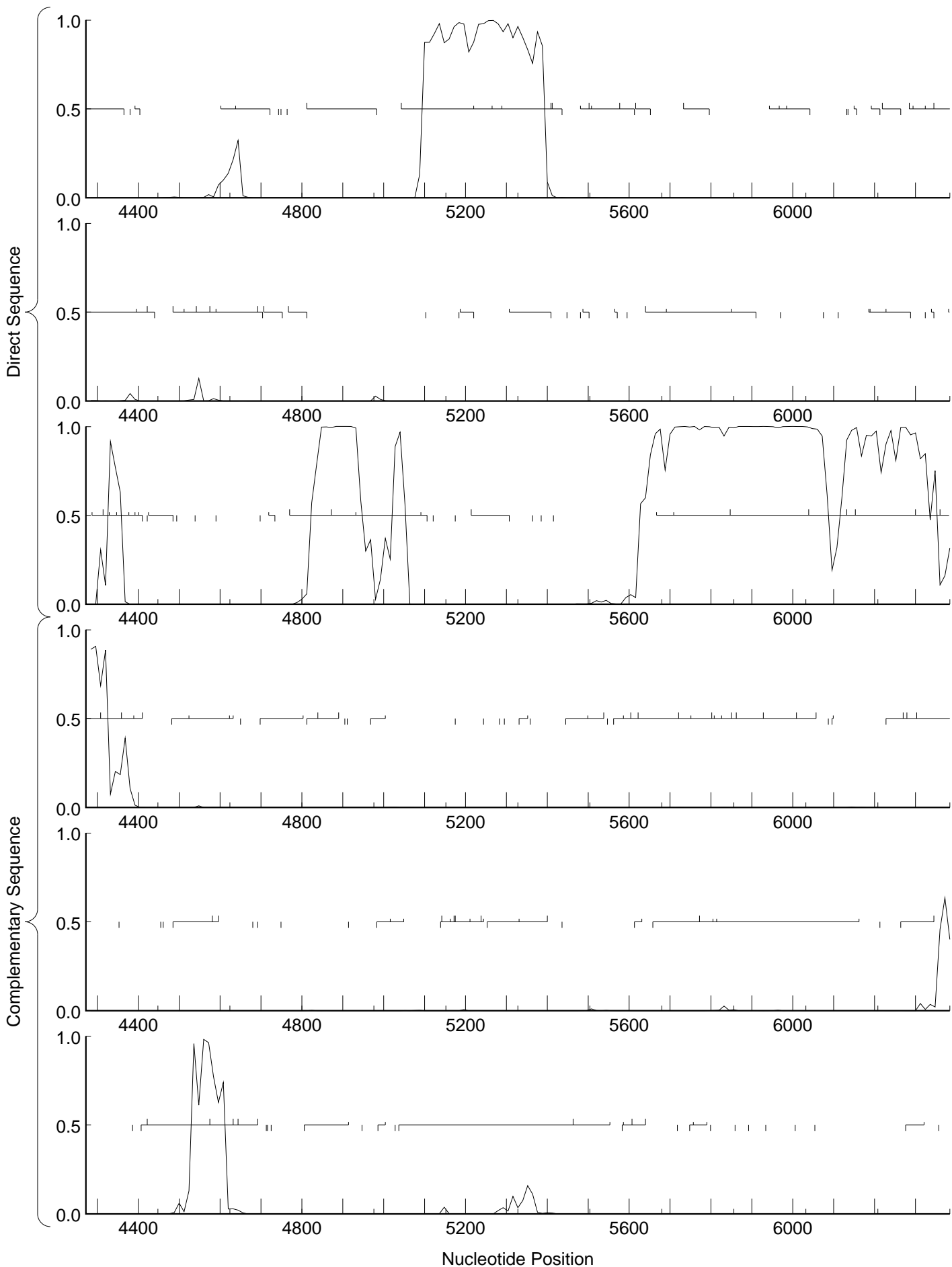
MATRIX INFORMATION

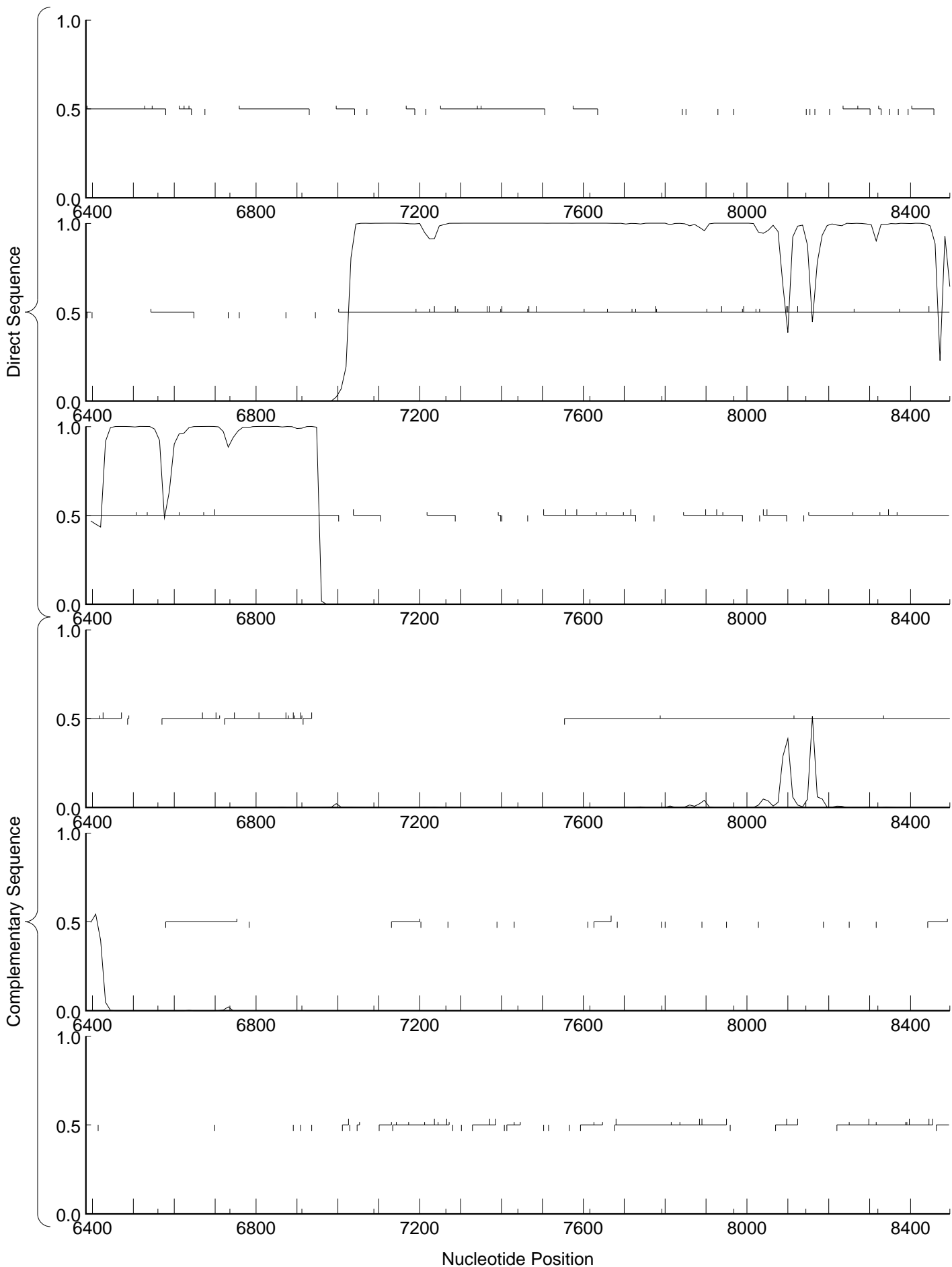
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

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

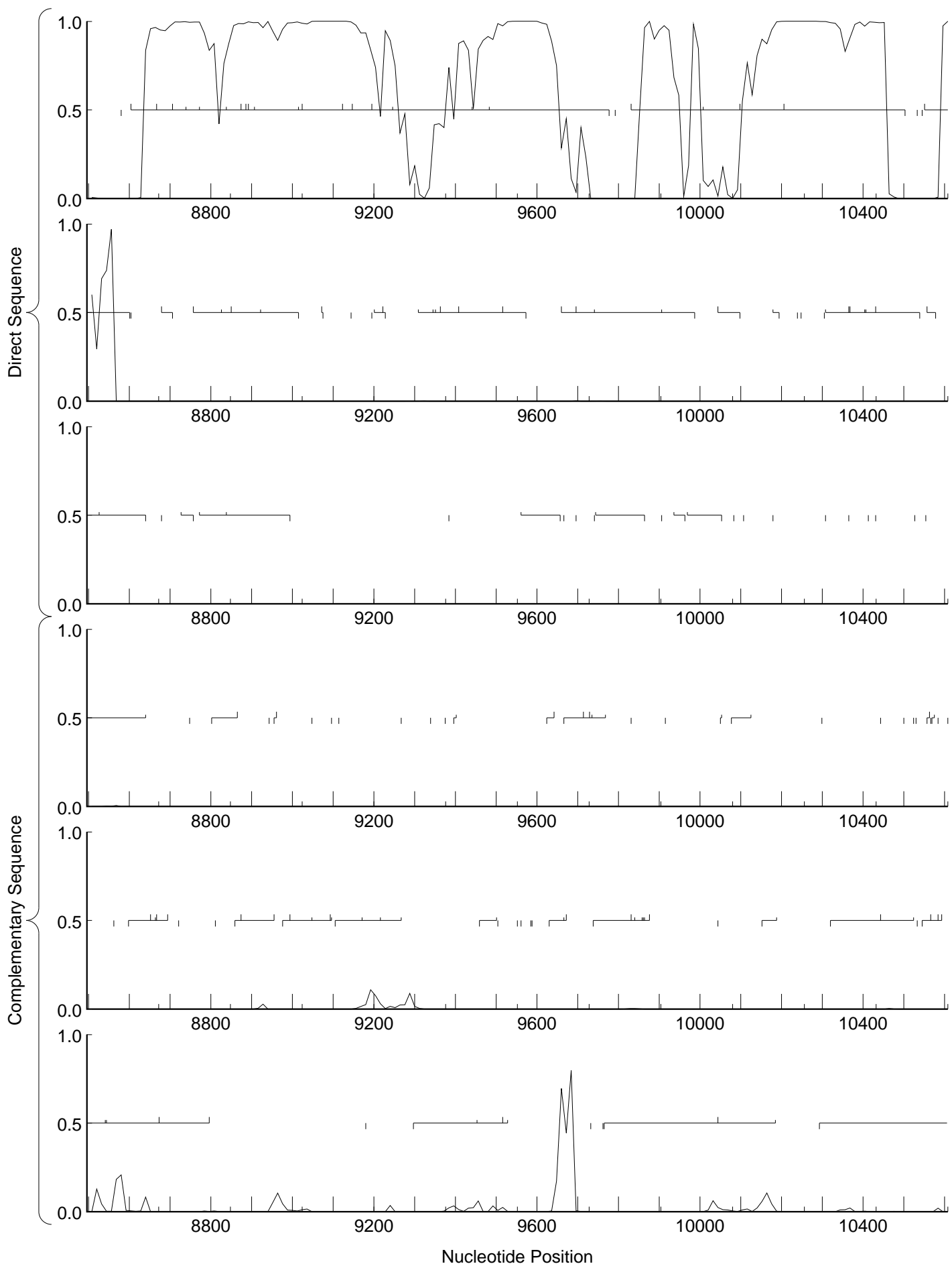




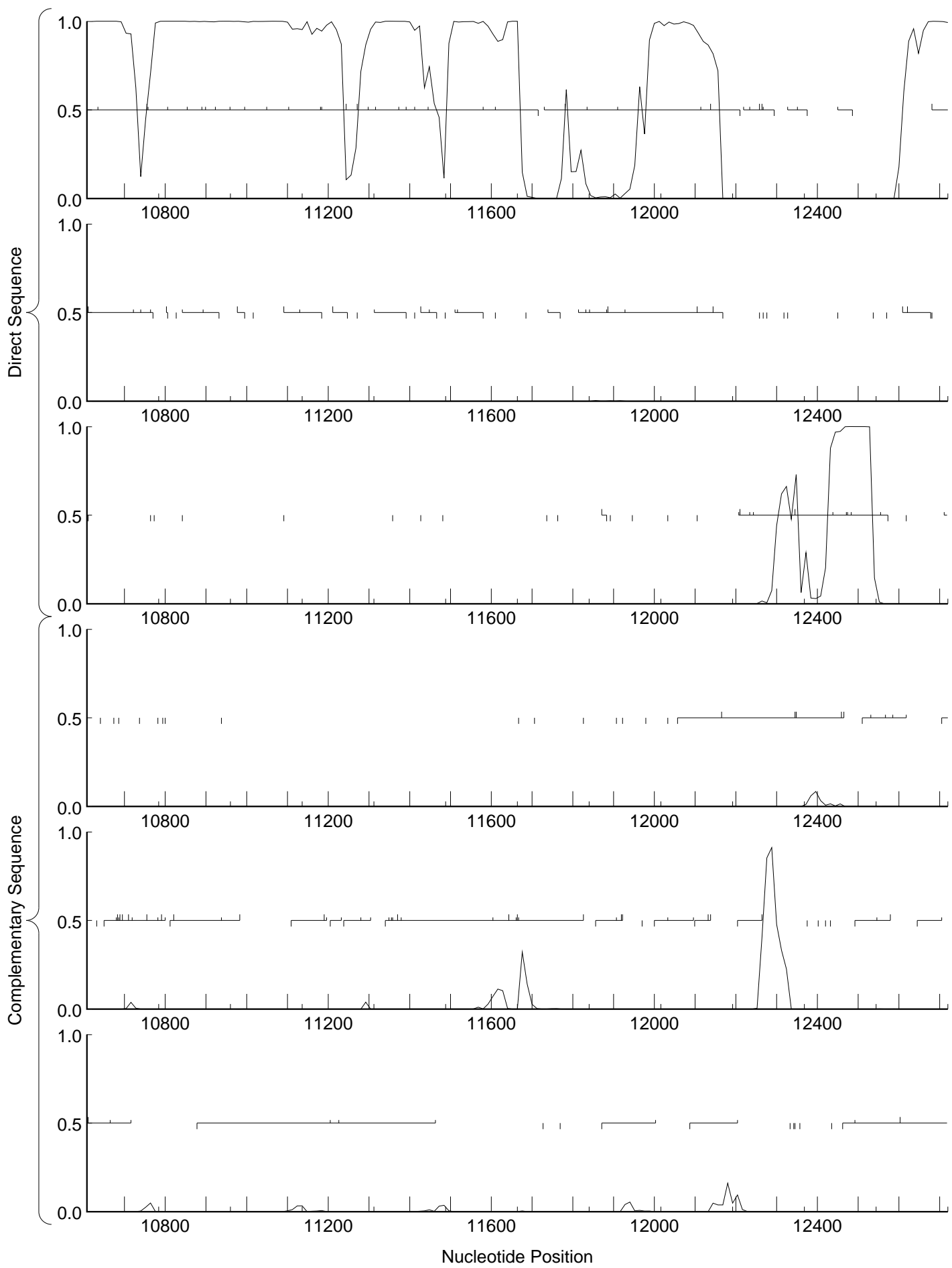




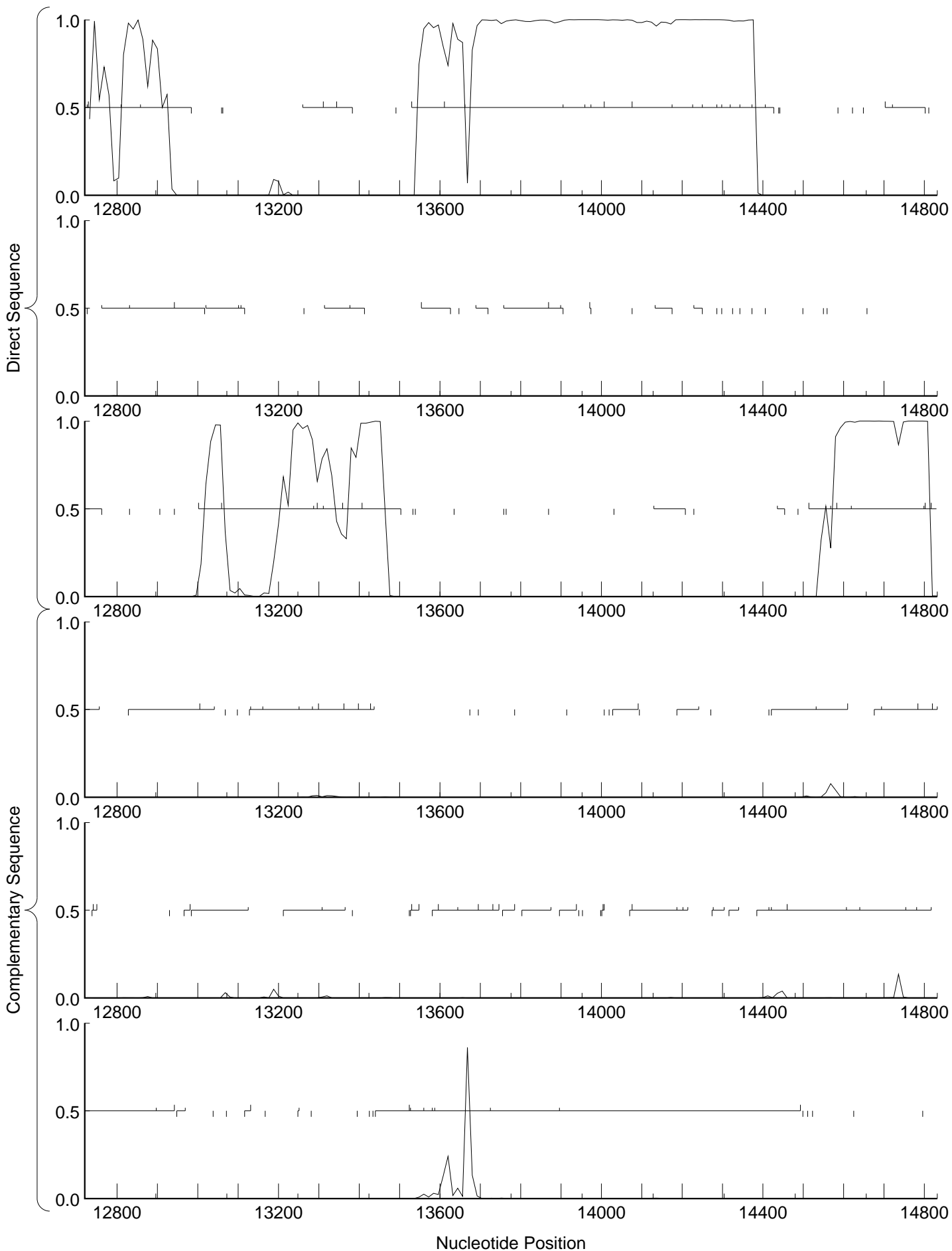
complete sequence, 83601 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 6/41



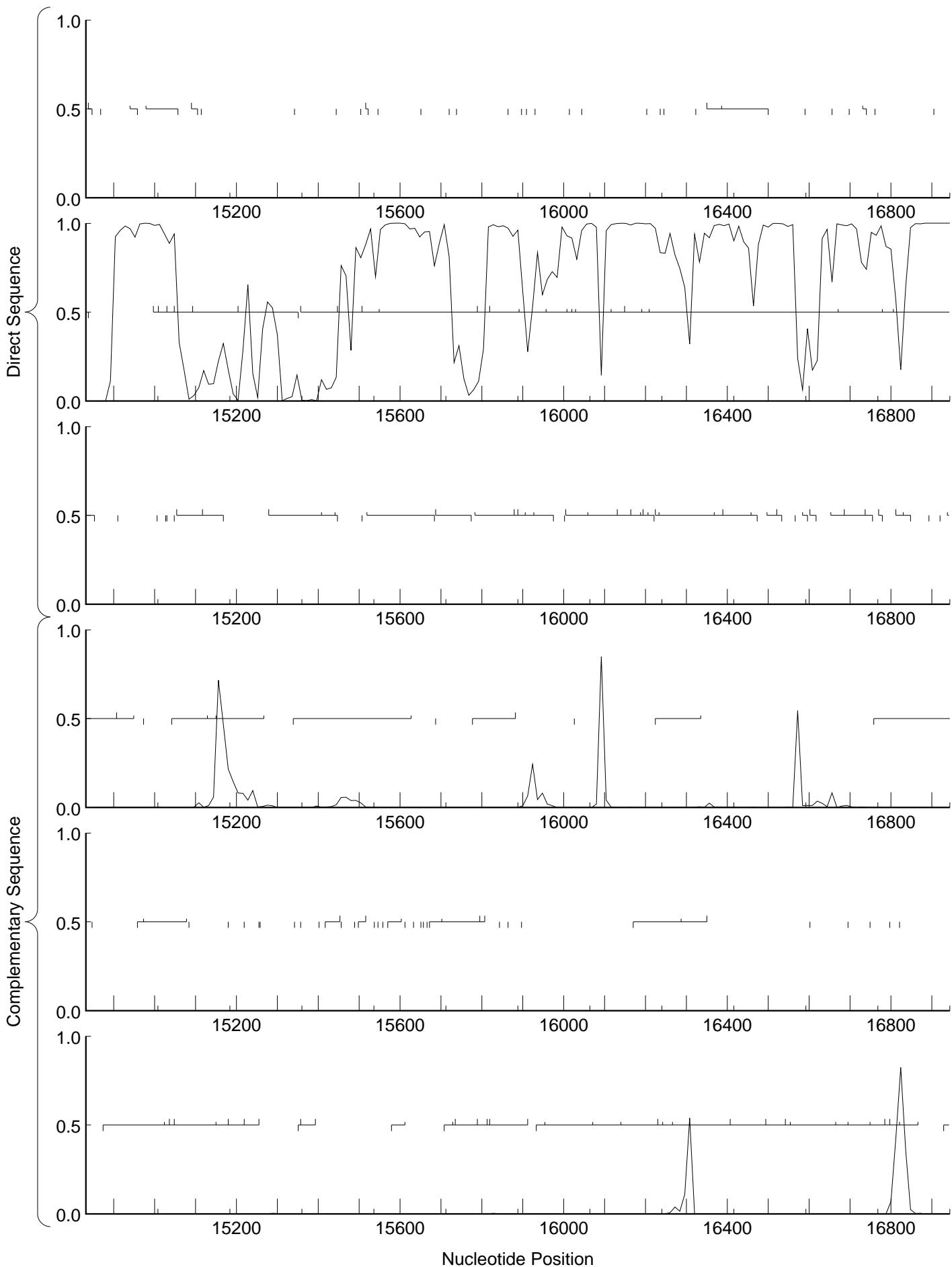
complete sequence, 83601 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 7/41

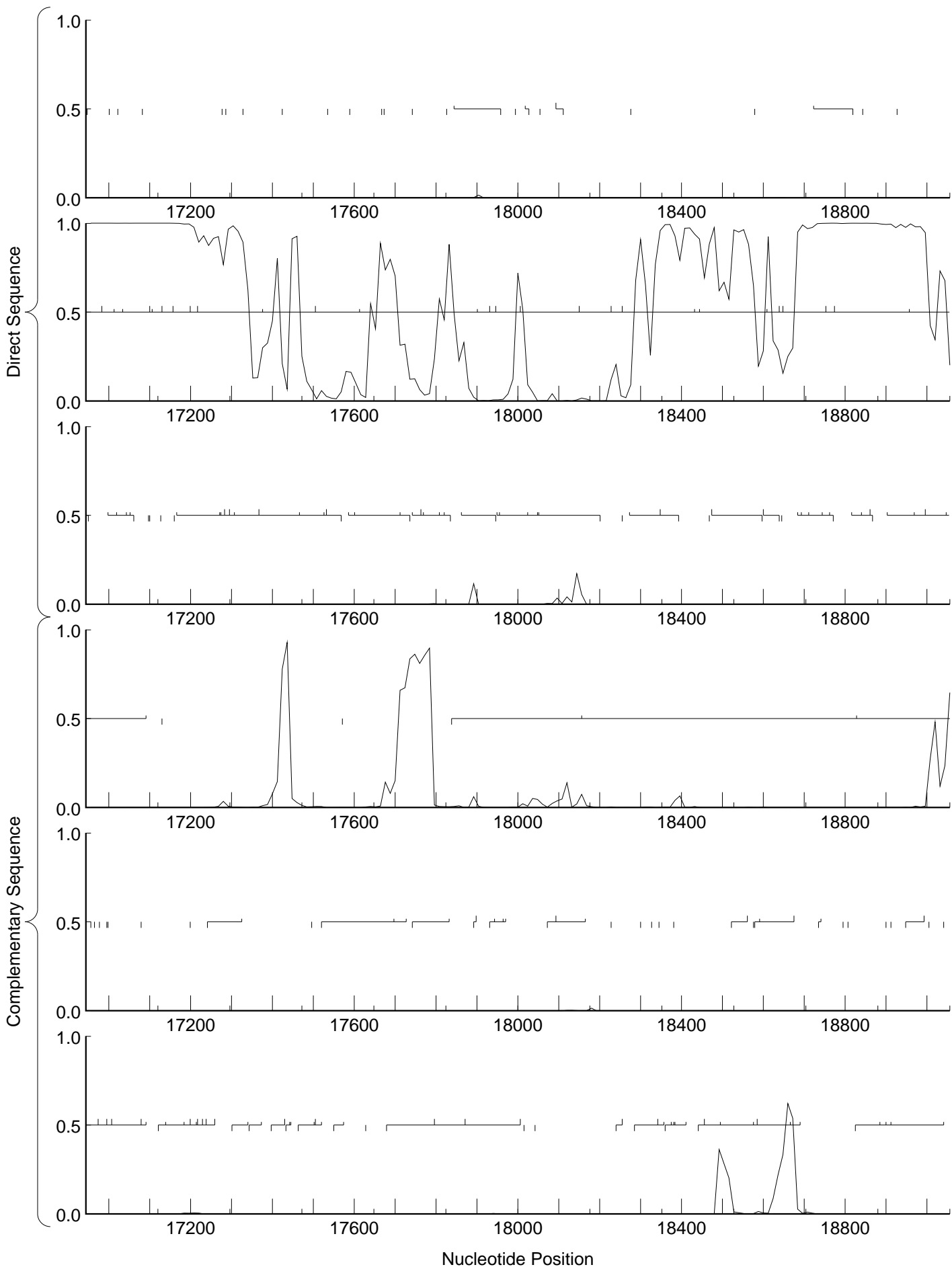


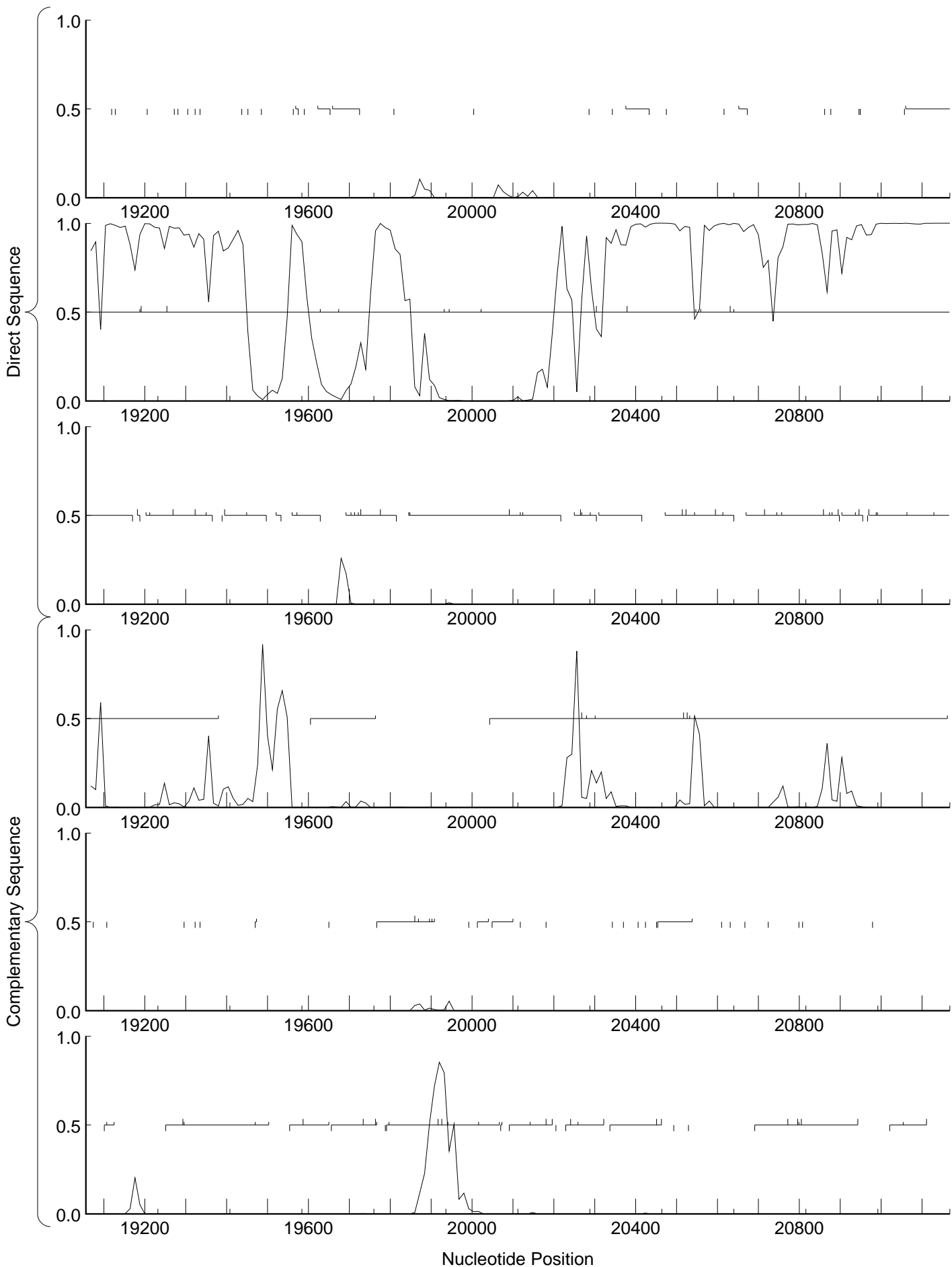
complete sequence, 83601 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 8/41

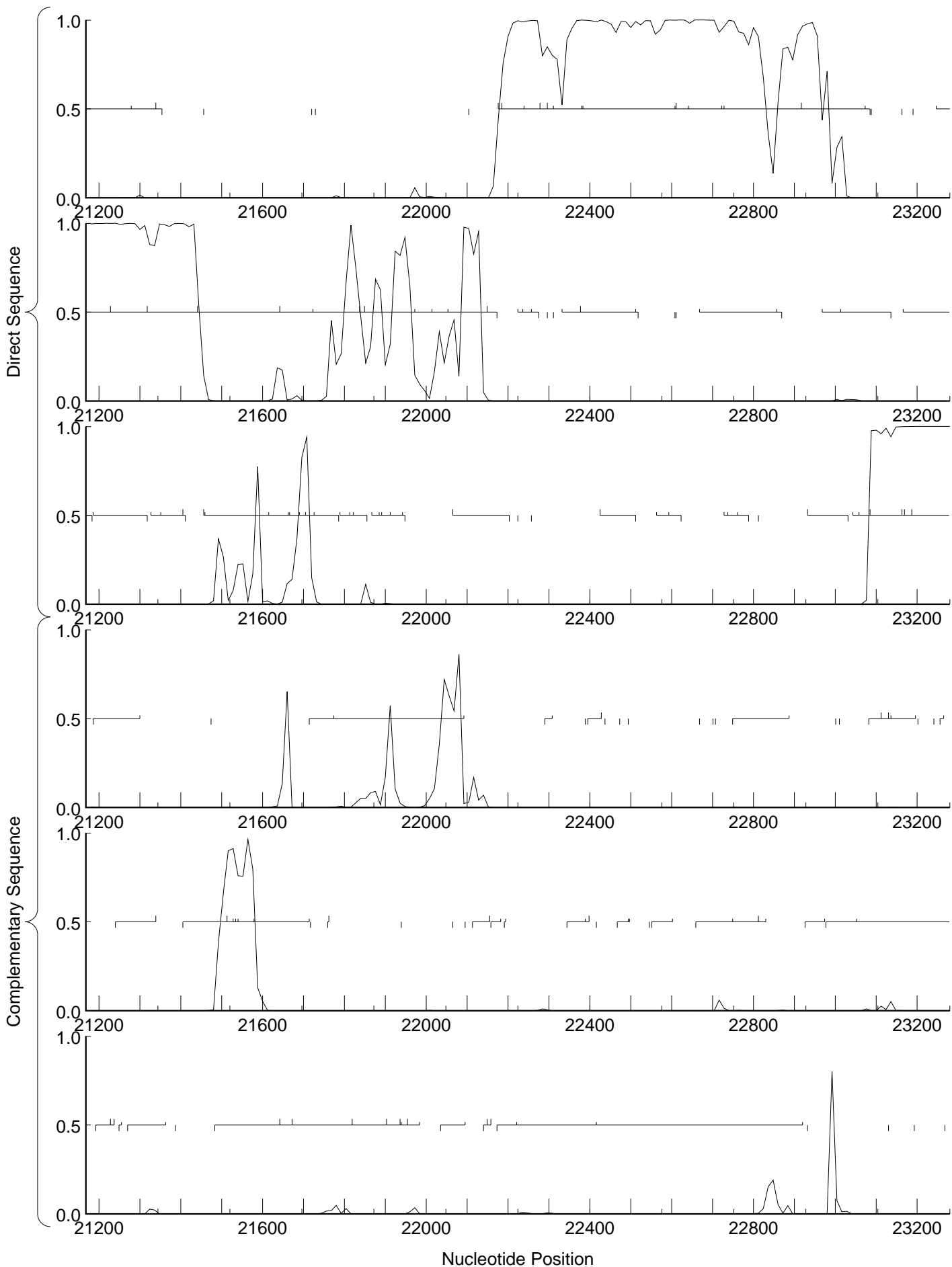


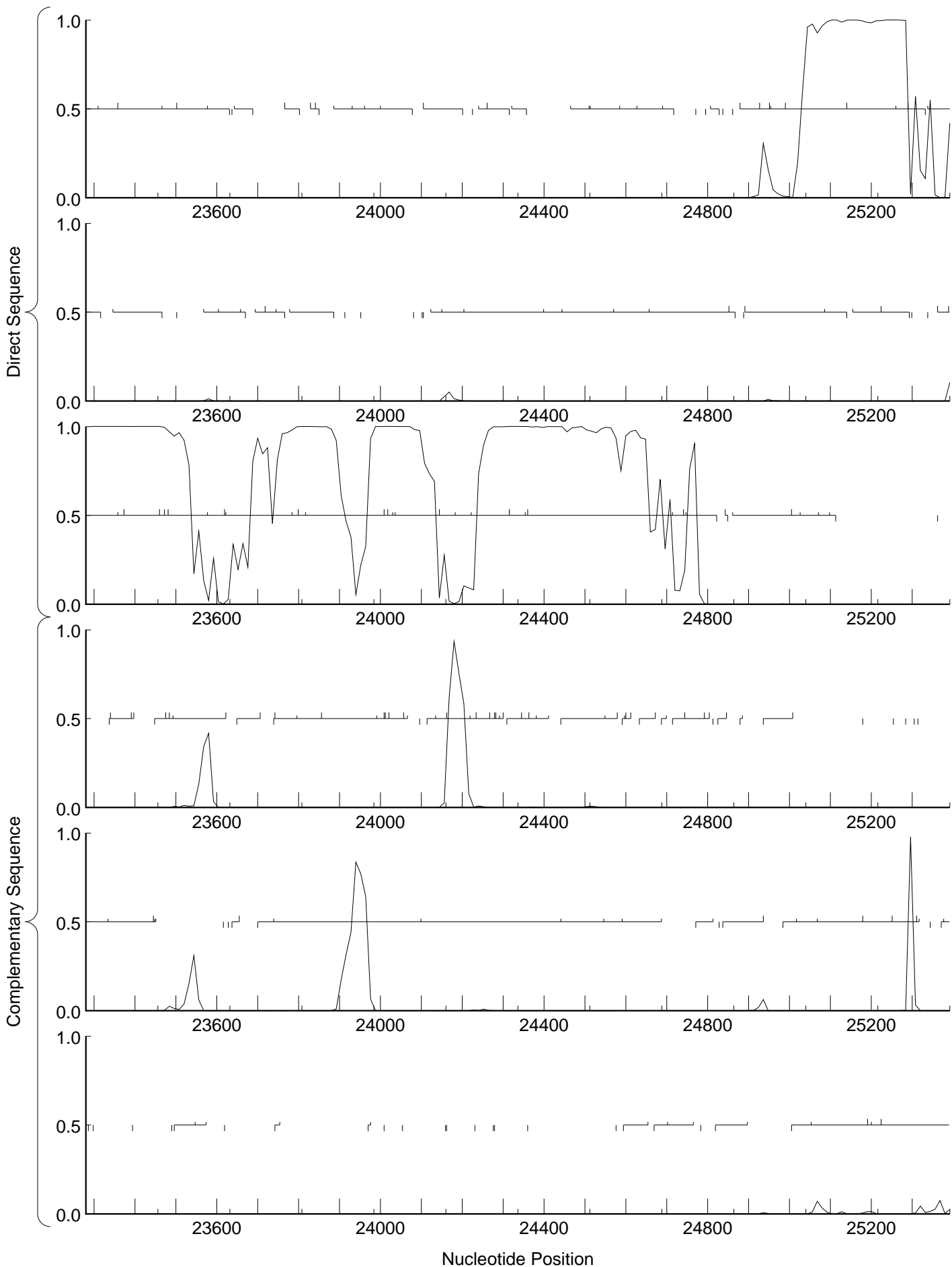
complete sequence, 83601 bp including 11-base 3' overhang (ACCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 9/41











complete sequence, 83601 bp including 11-base 3' overhang (ACCCCATGCAA), Cluster M2, Order 4, Window 96, Step 12, 14/41

