

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Mycobacterium phage Clautastrophe complete sequence, 72055 bp including 10-base 3' overhang
Analysis Date : 2/8/23 at 5:21:15
Pages : 36
Sequence Length : 72055 bp
GC Content : 59.19%

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,

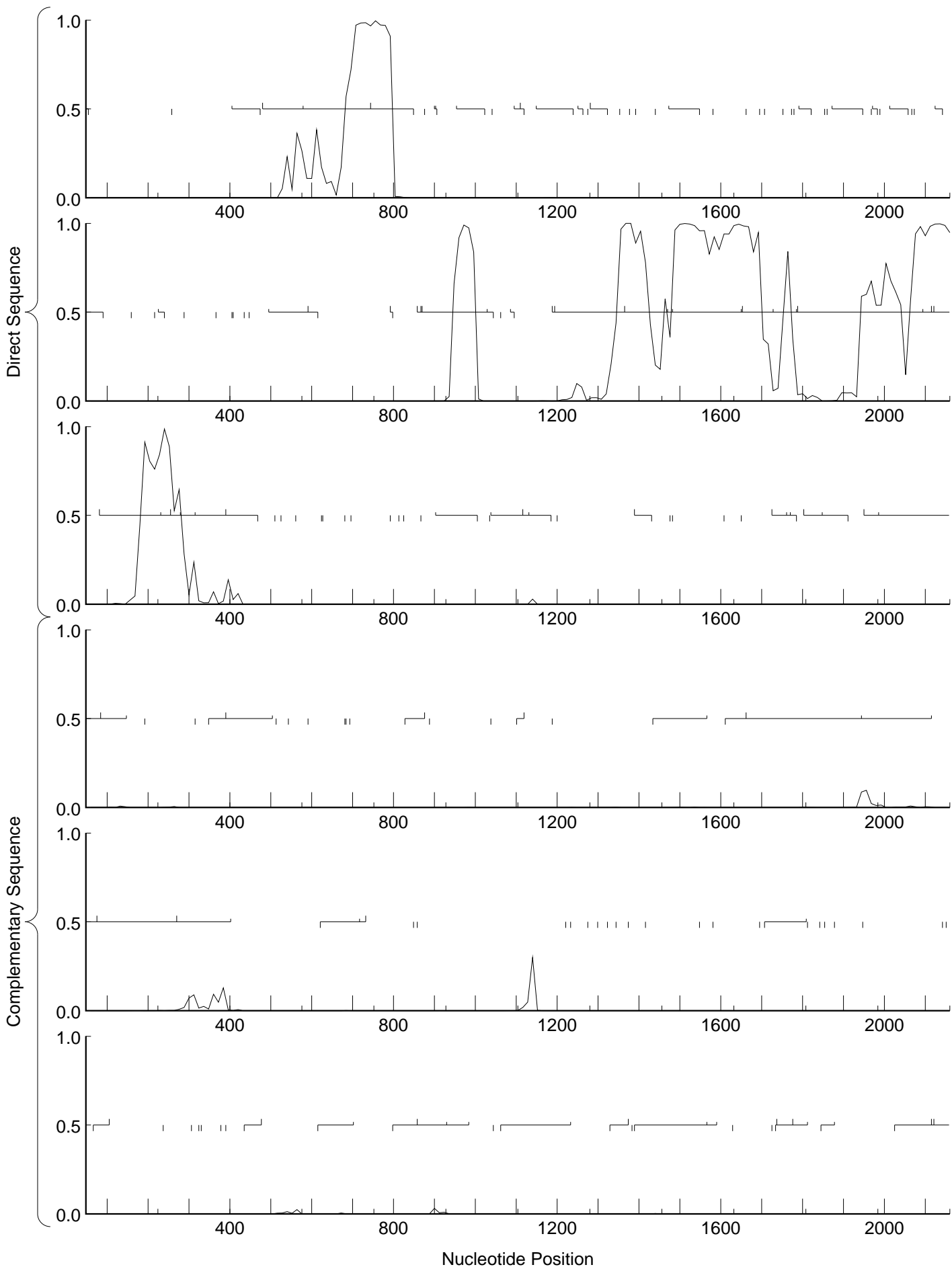
MATRIX INFORMATION

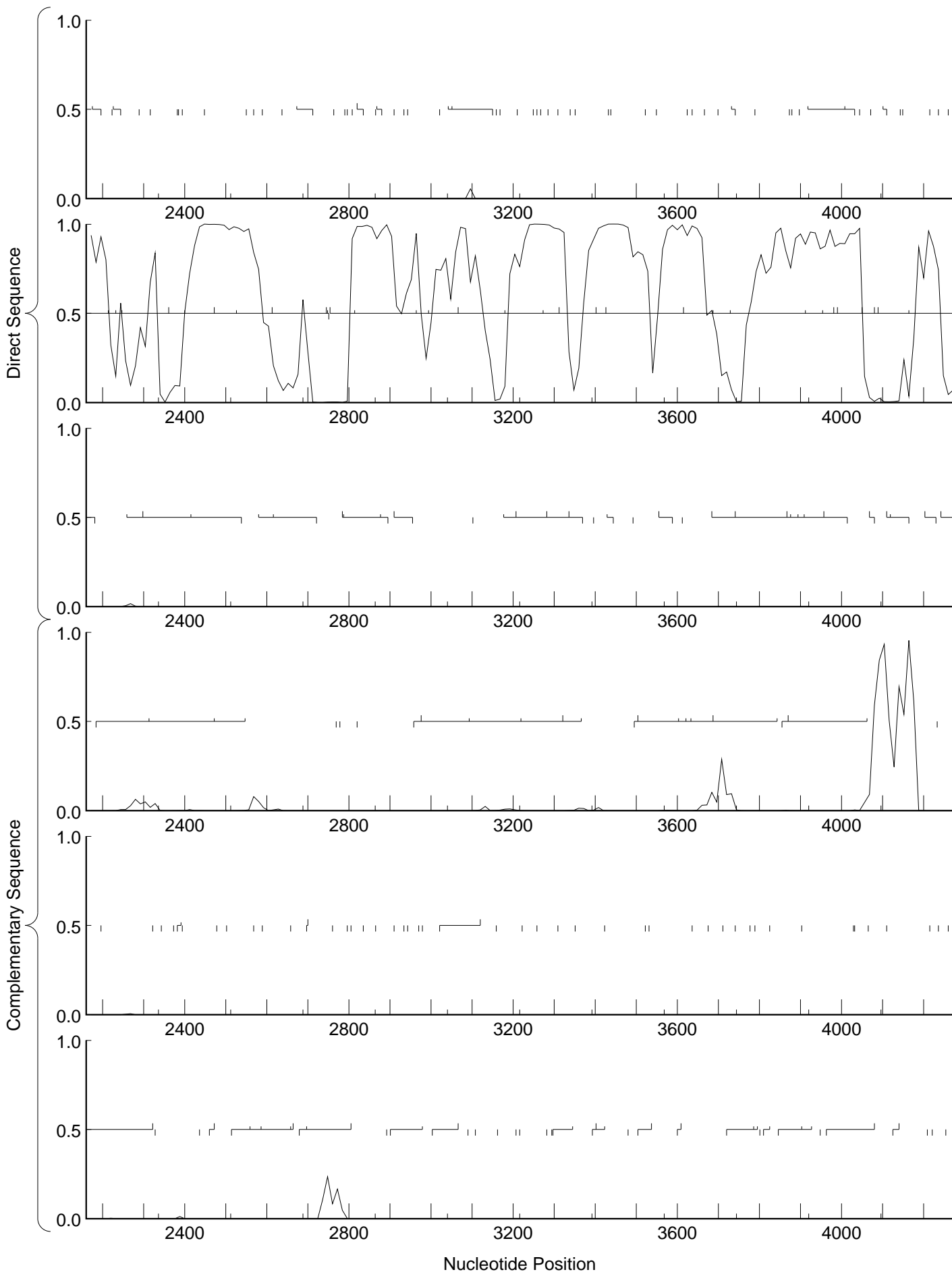
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

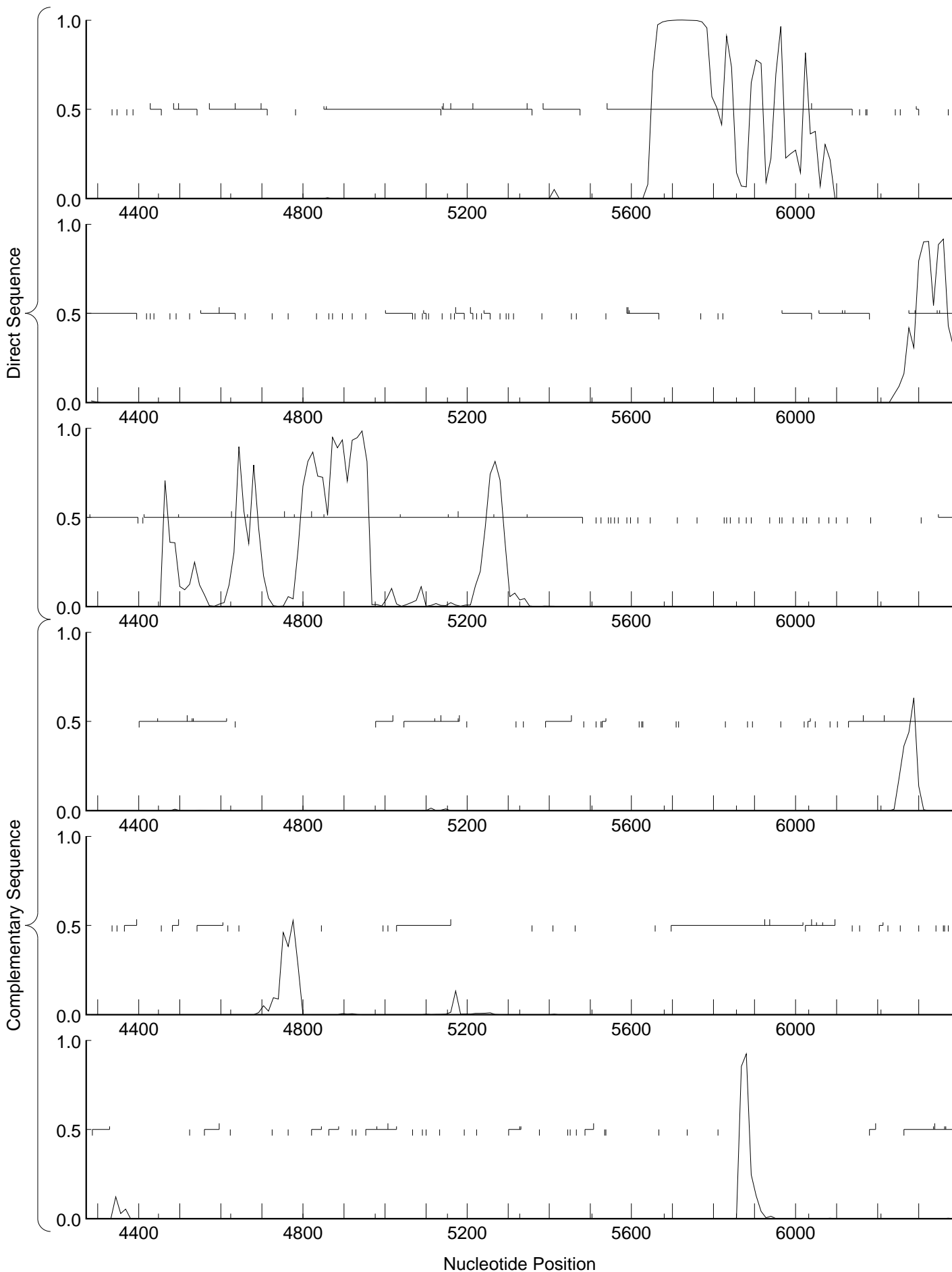
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

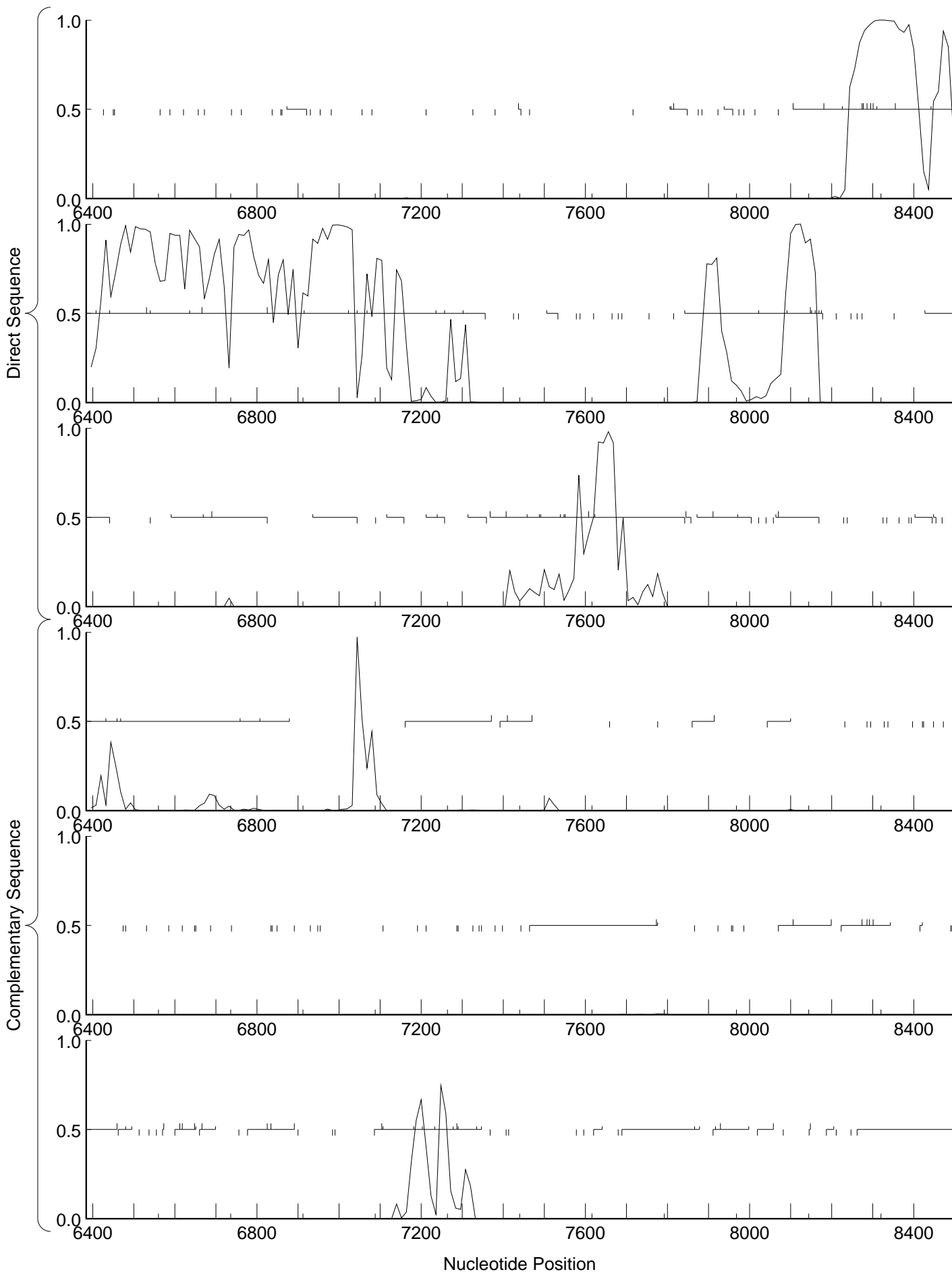
Matrix notes & comments

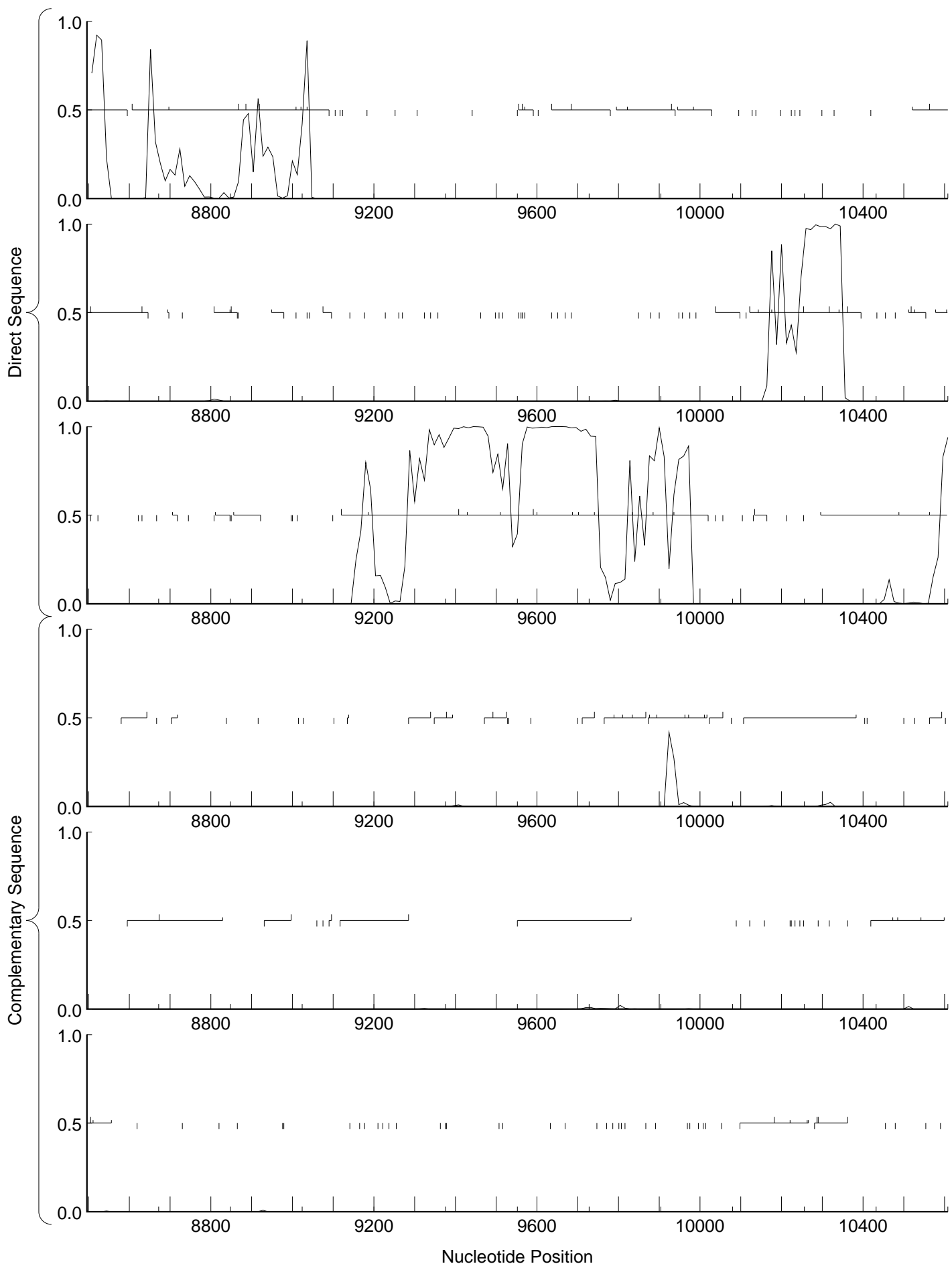
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014



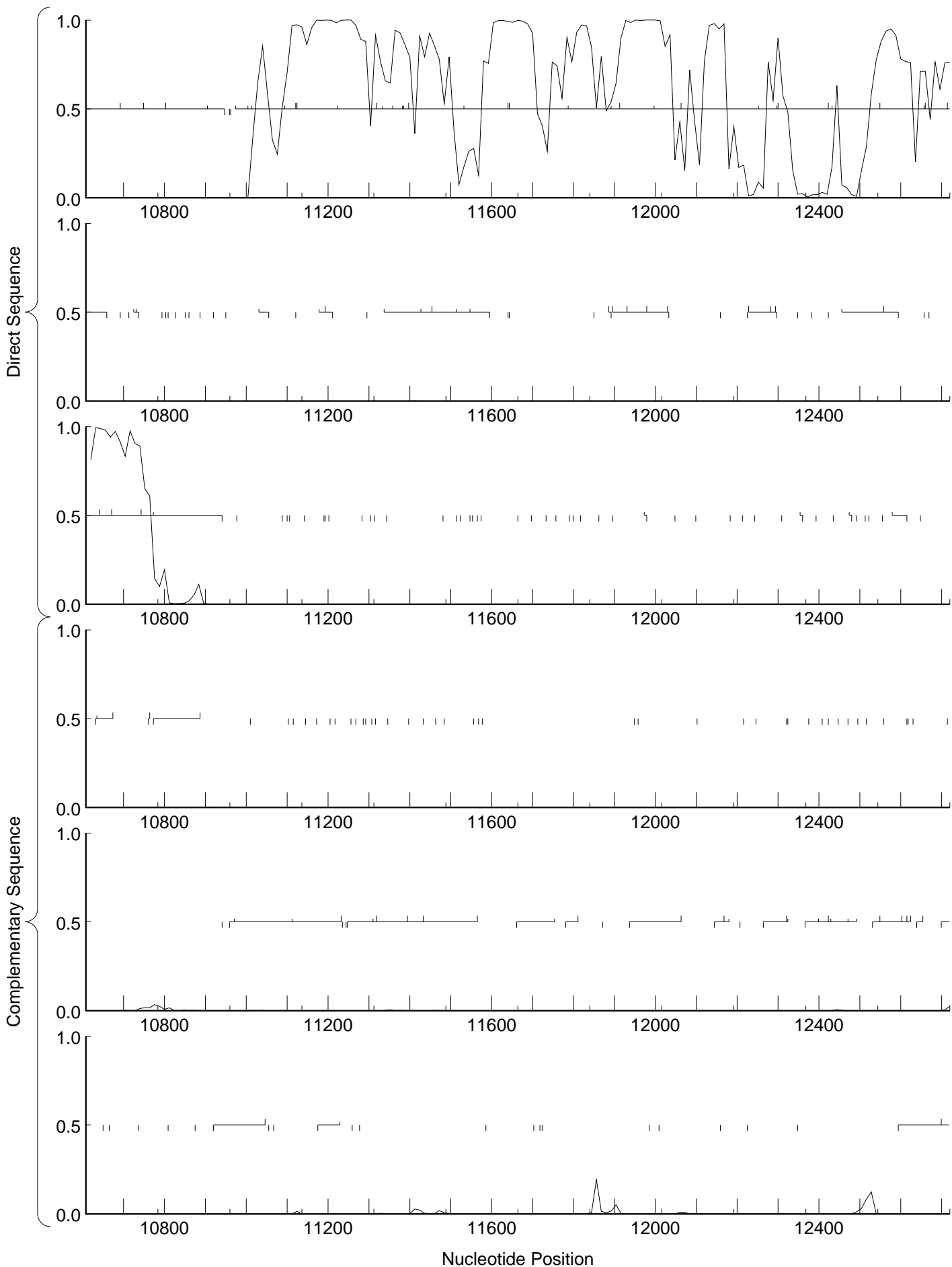








the complete sequence, 72055 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L3, Order 4, Window 96, Step 12, 7/36



the complete sequence, 72055 bp including 10-base 3' overhang (TCGATCAGCC), Cluster L3, Order 4, Window 96, Step 12, 8/36

