

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Microbacterium phage Scamander complete sequence, 17452 bp including 9-base 3' overhang (CC
Analysis Date : 7/3/22 at 7:58:50
Pages : 10
Sequence Length : 17452 bp
GC Content : 68.73%

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,

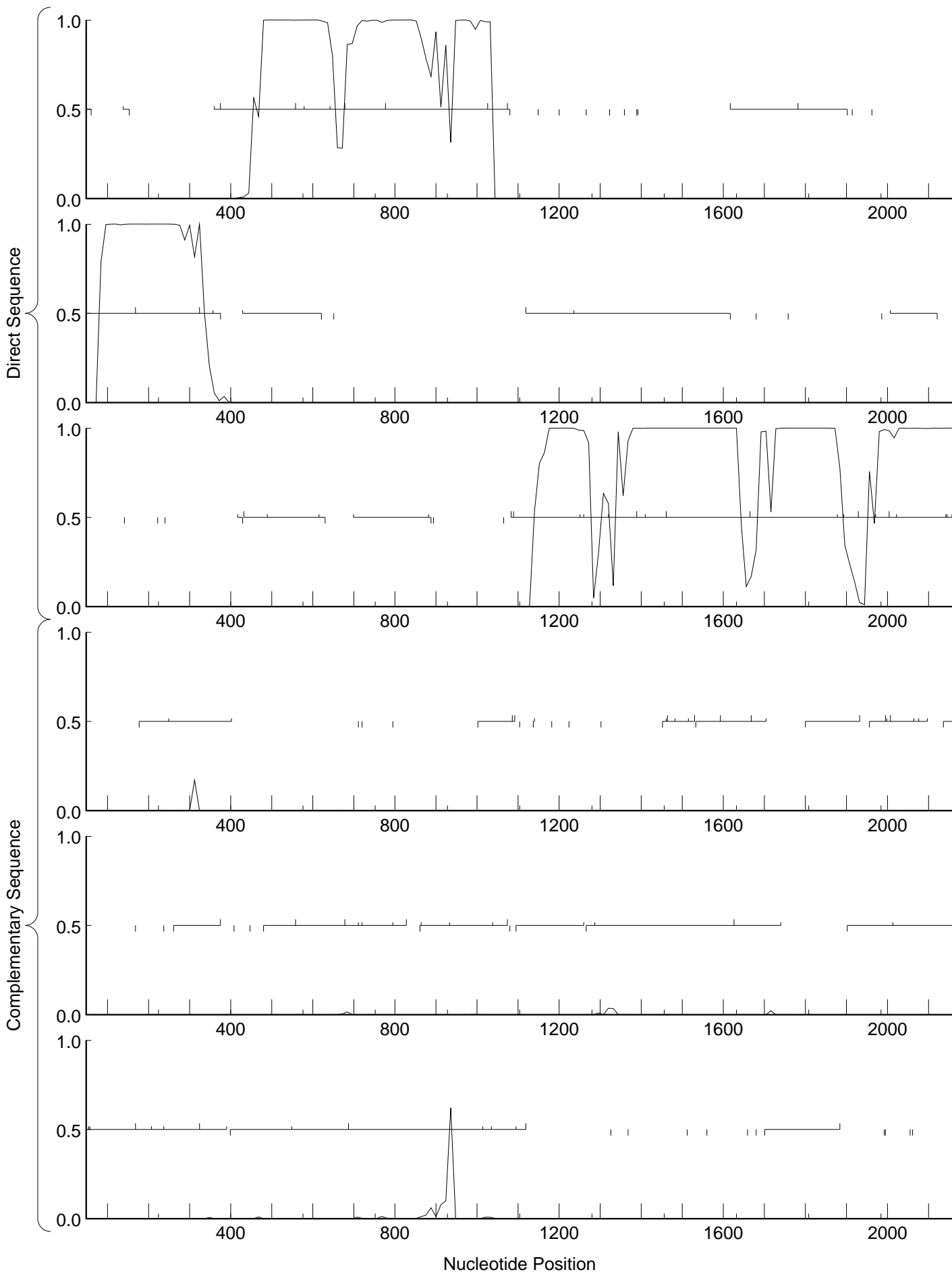
Matrix notes & comments

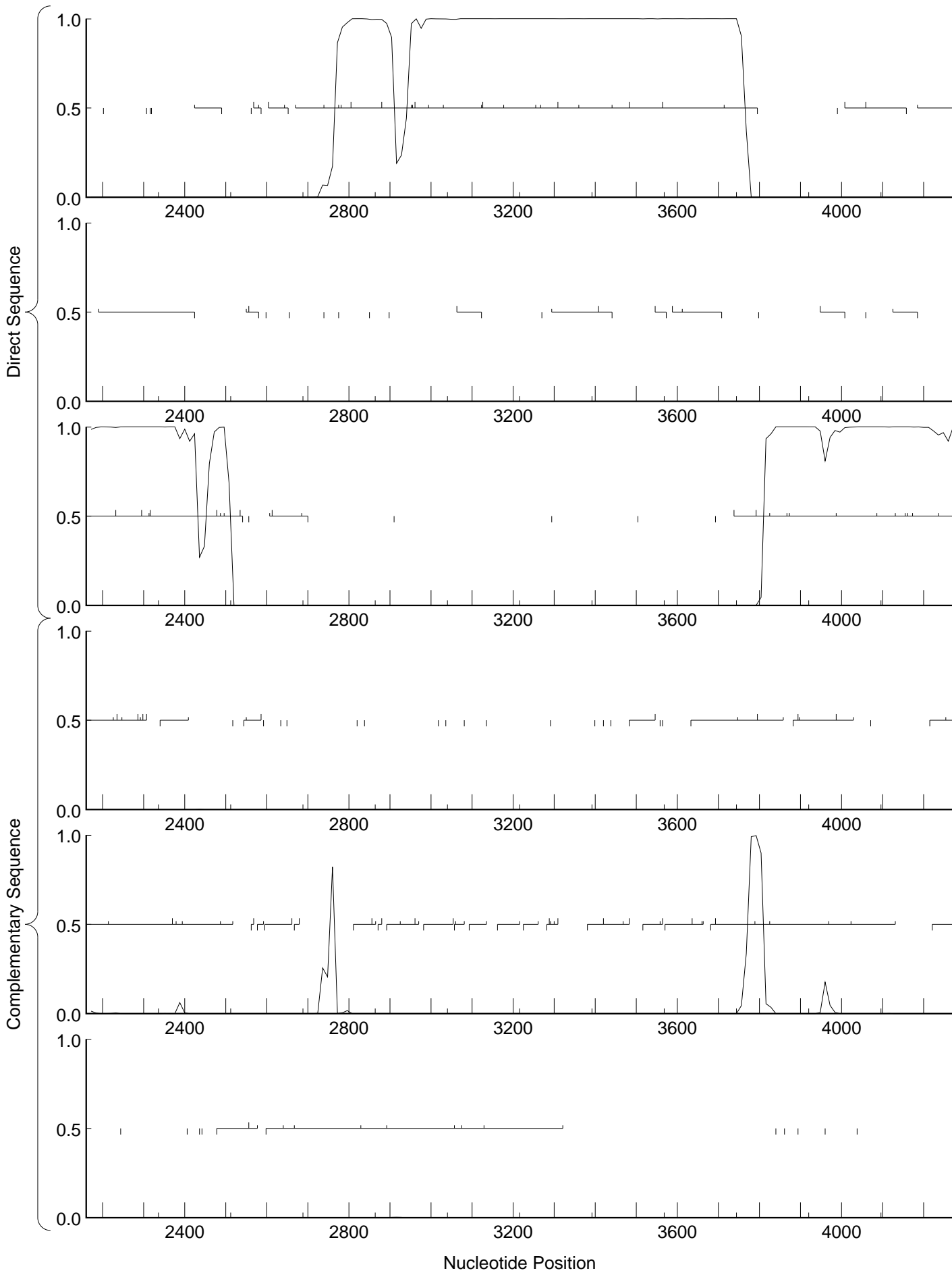
Training set derived by GeneMarkS, 4.30 October 2014
Fri Sep 7 18:42:59 2018

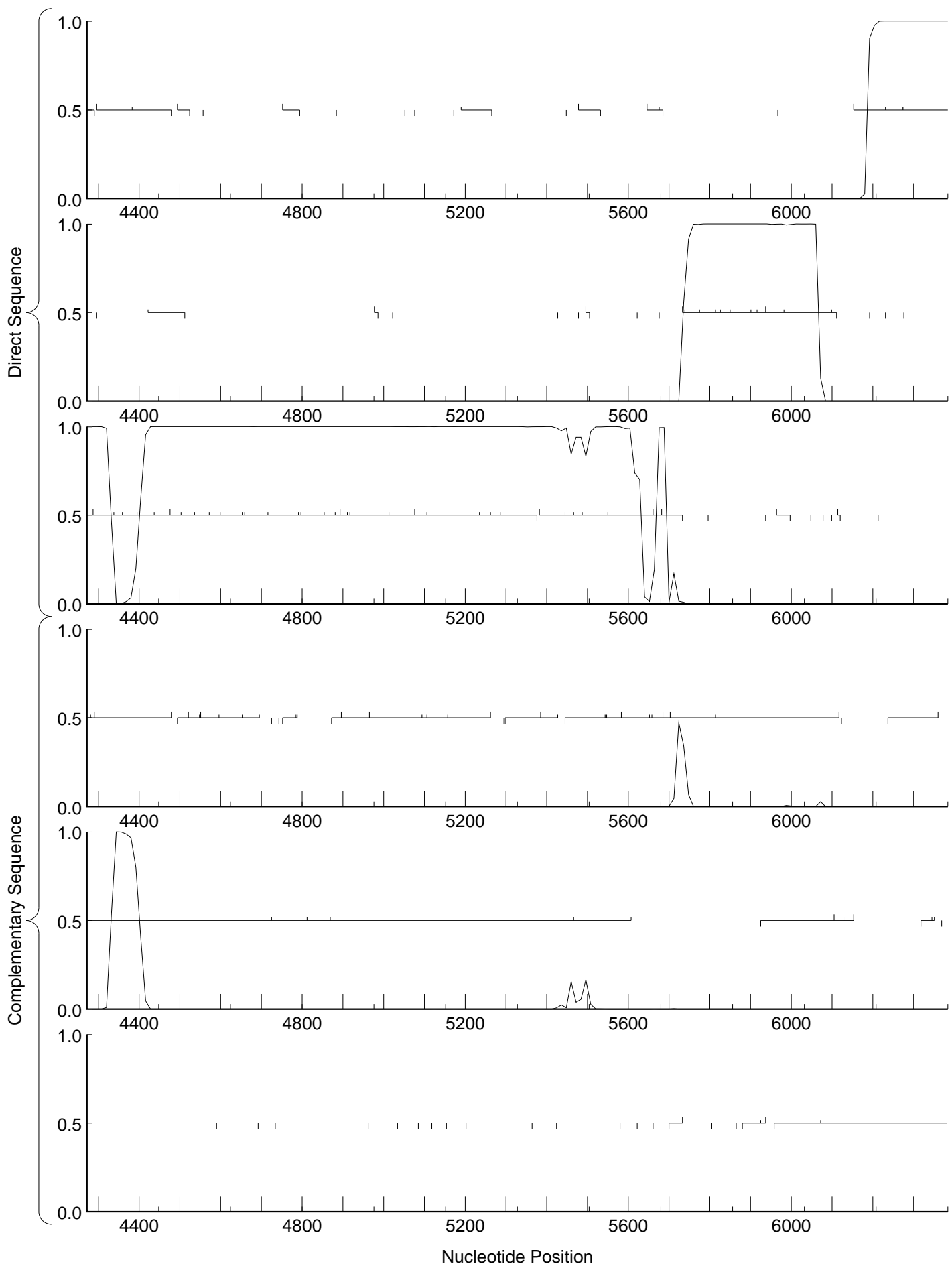
MATRIX INFORMATION

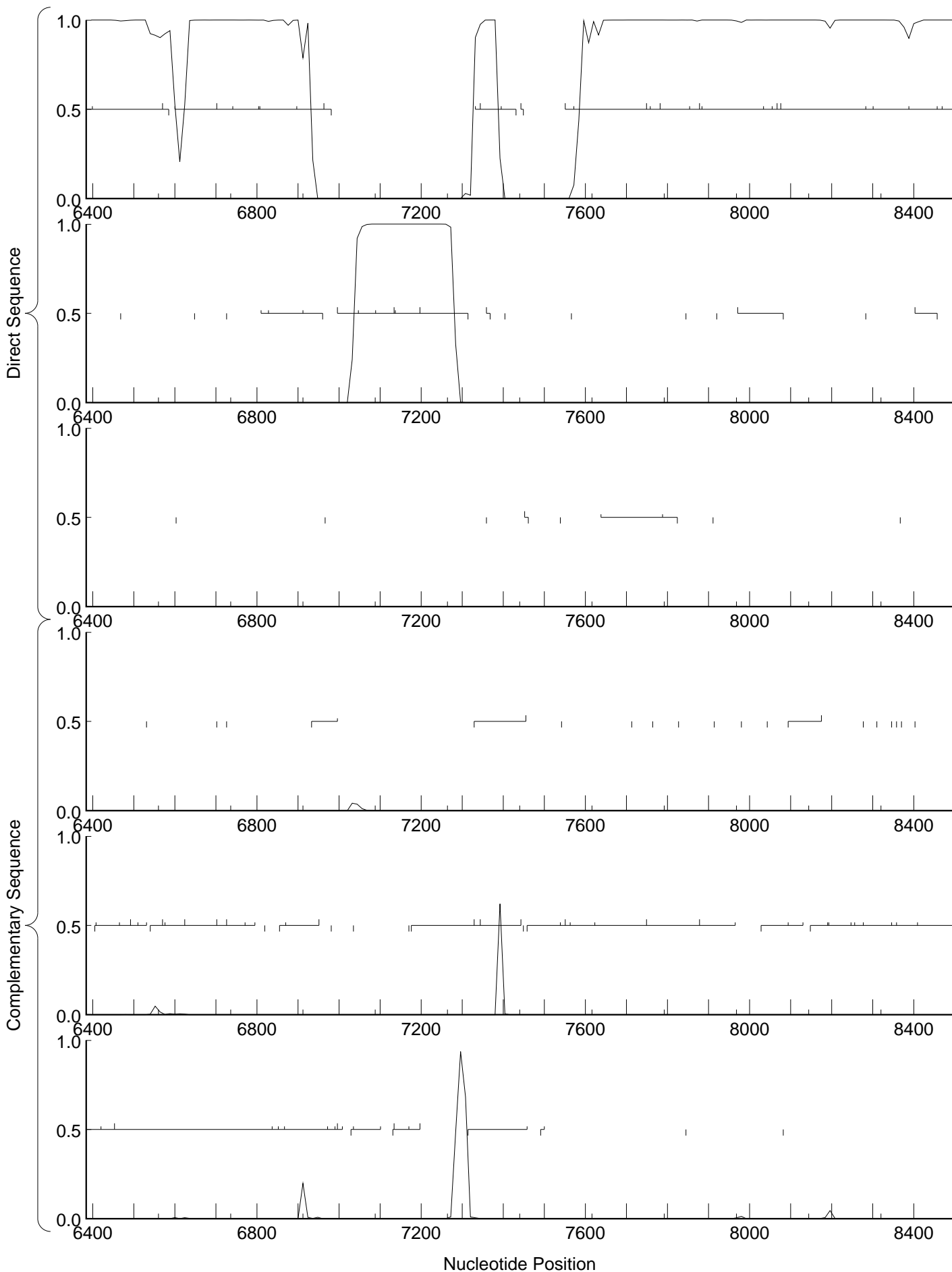
Matrix : -
Author : -
Order : 4

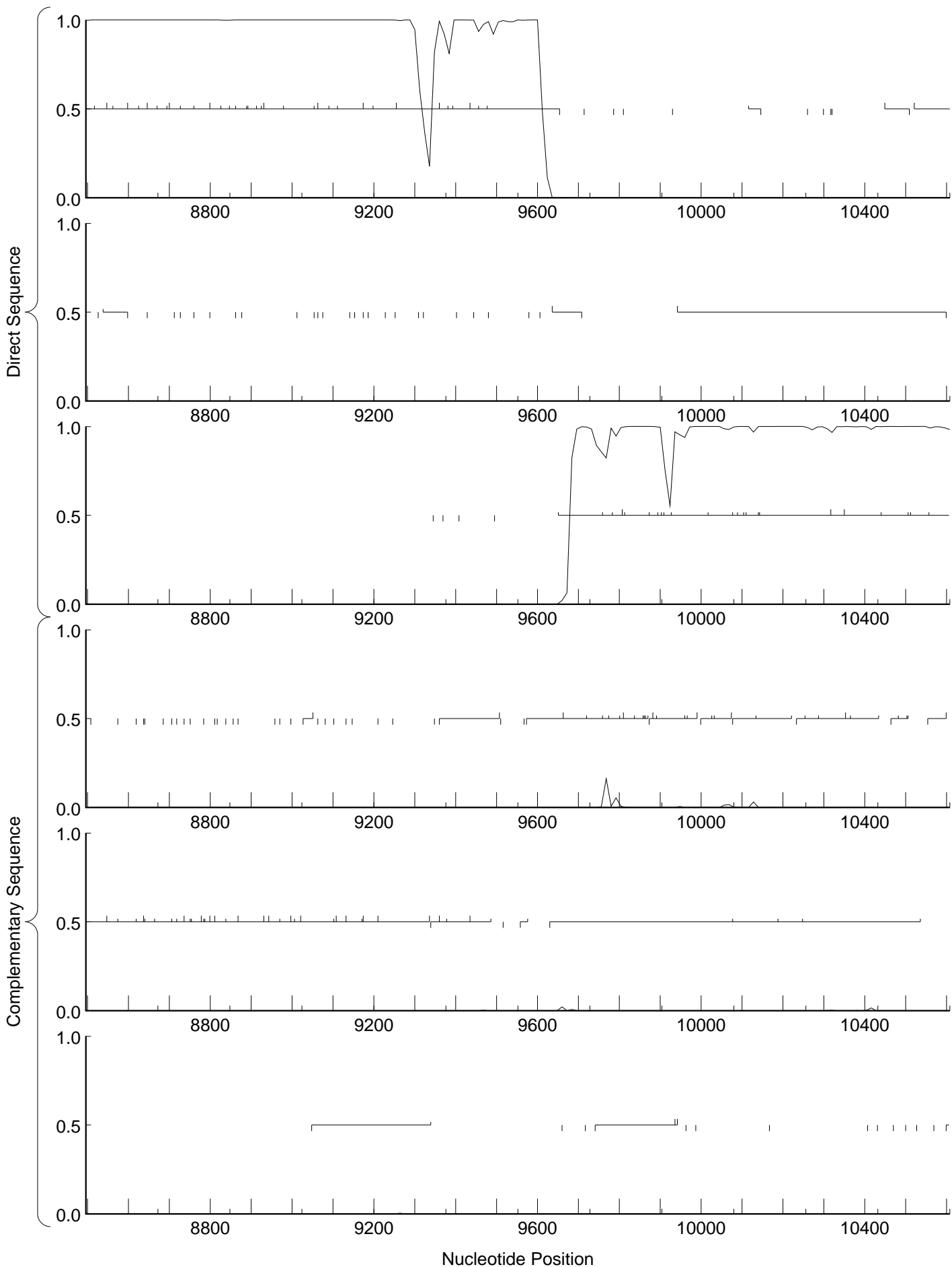
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

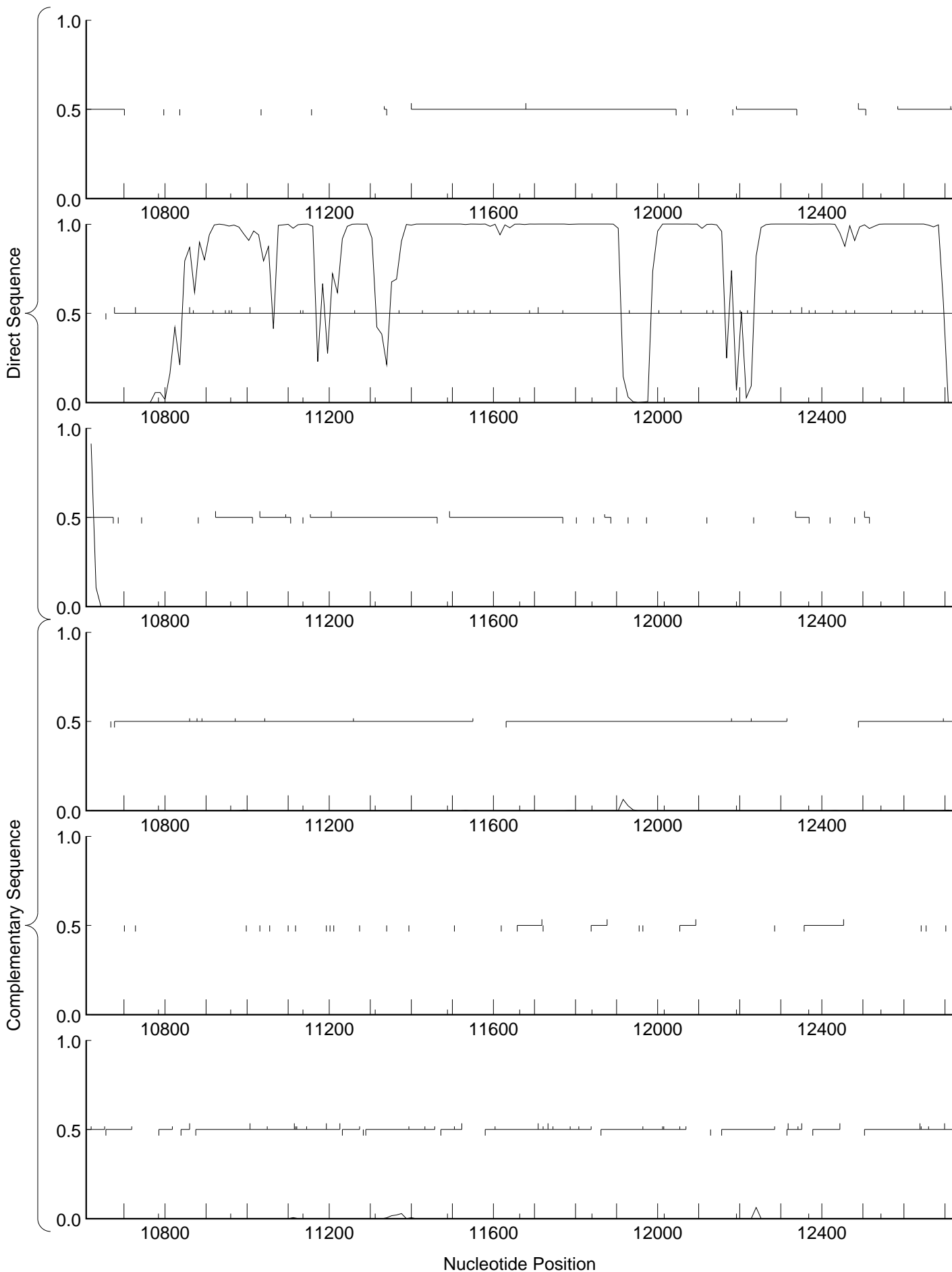


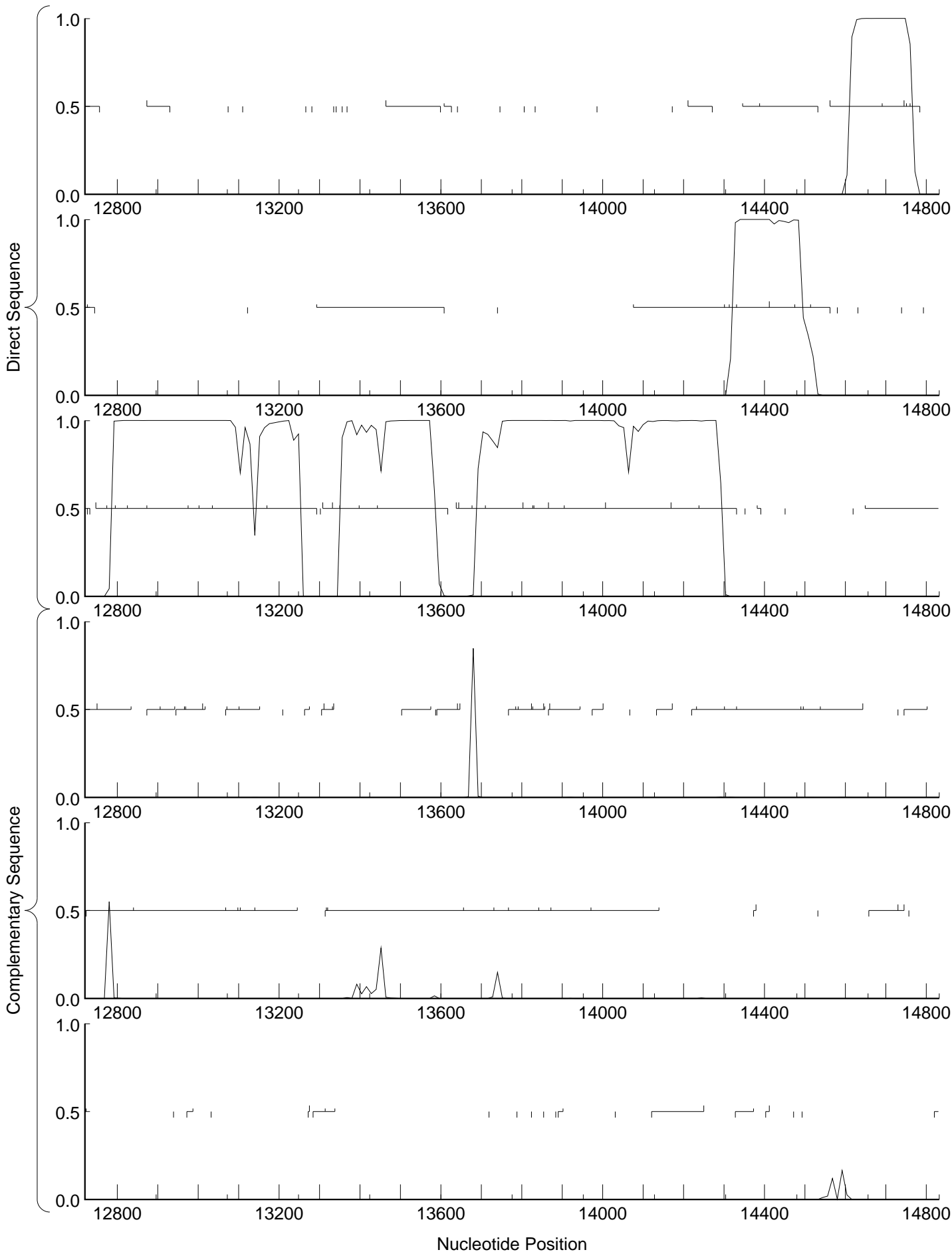


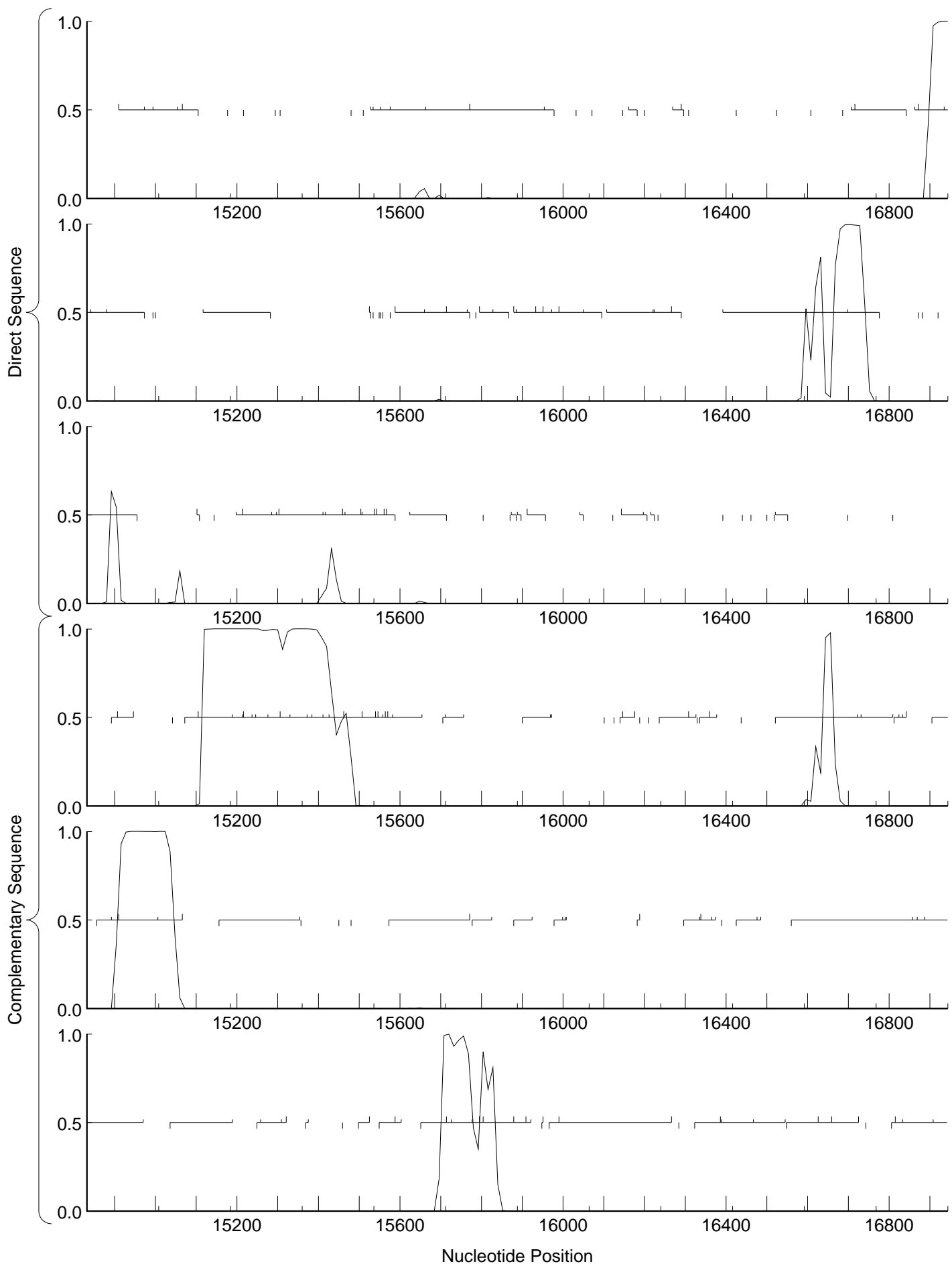












der complete sequence, 17452 bp including 9-base 3' overhang (CCCGCCCCA), Cluster EE, Order 4, Window 96, Step 12, 10/10

