

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Arthrobacter phage Andrew complete sequence, 38802 bp including 12-base 3' overhang (GAGTTG
Analysis Date : 12/2/23 at 1:19:03
Pages : 20
Sequence Length : 38802 bp
GC Content : 65.53%

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,

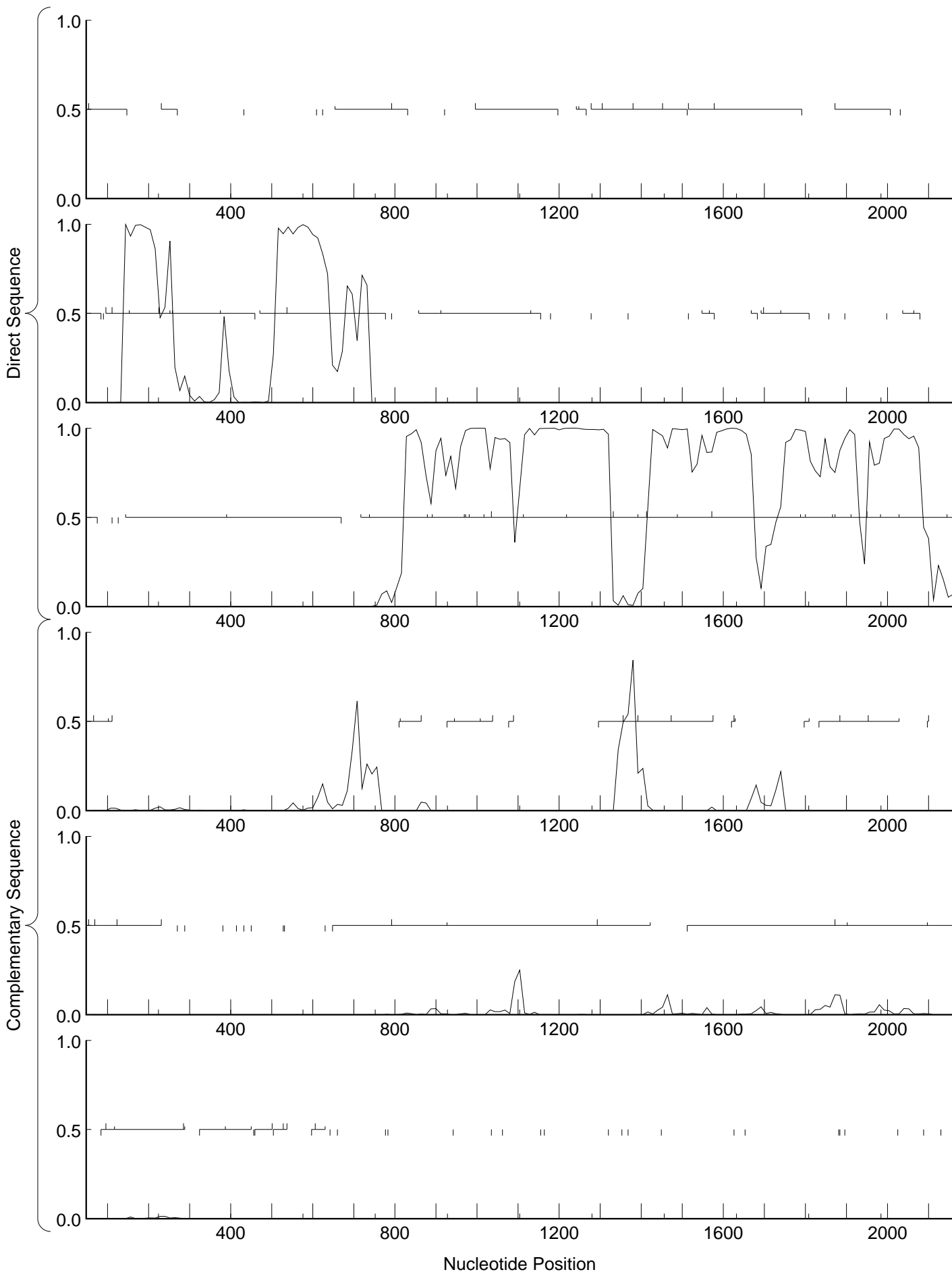
Matrix notes & comments

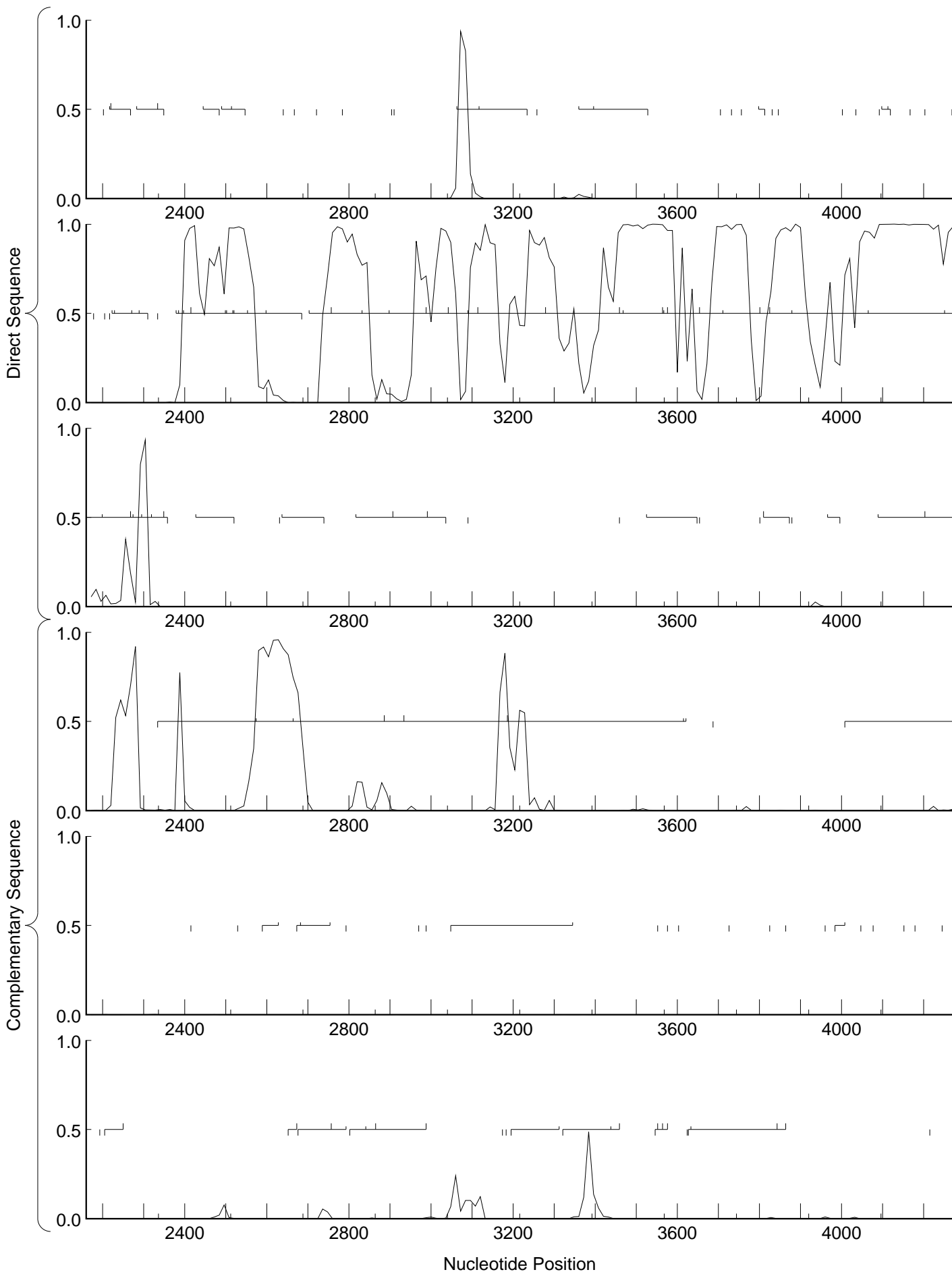
Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:22:00 2018

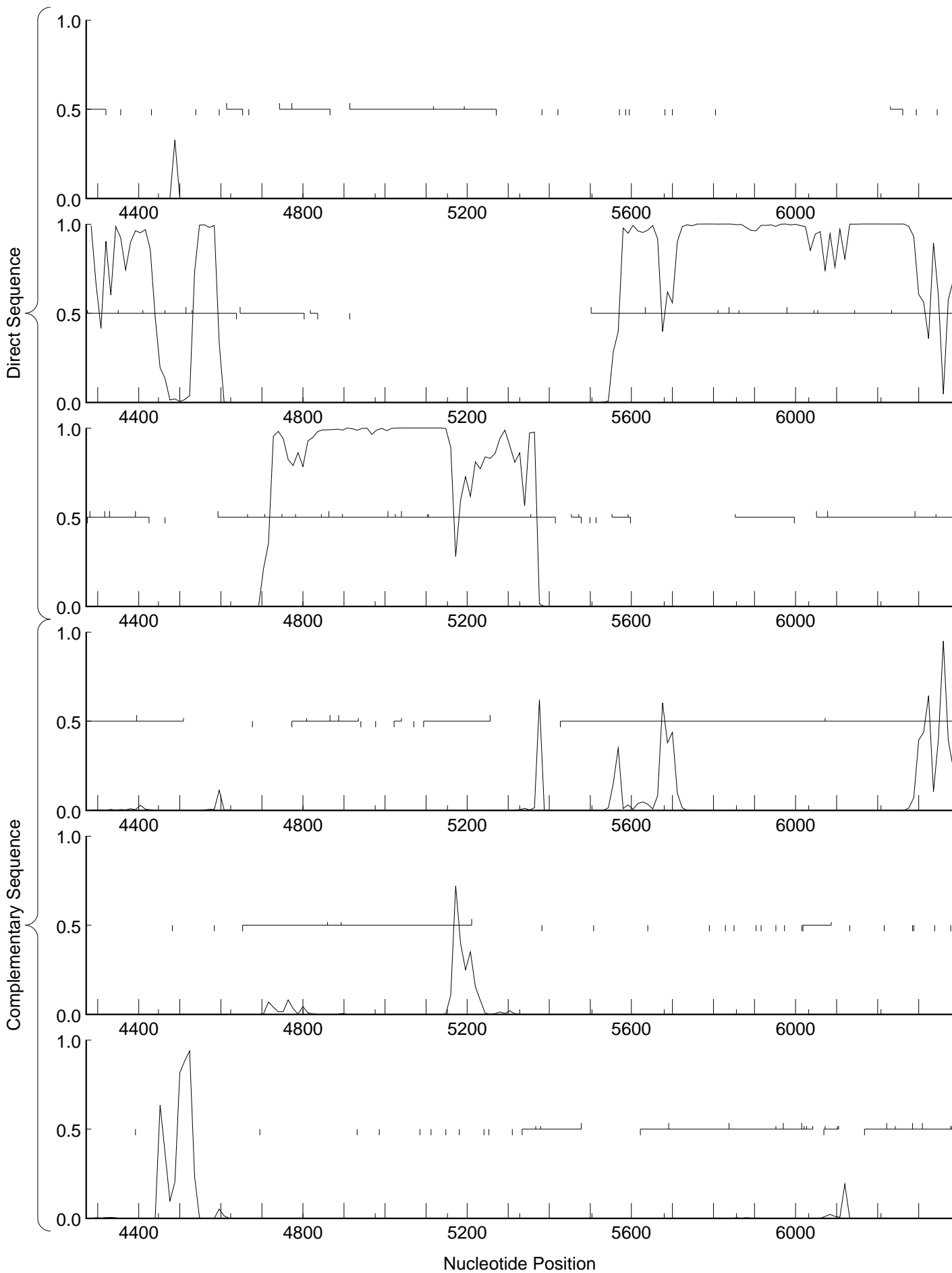
MATRIX INFORMATION

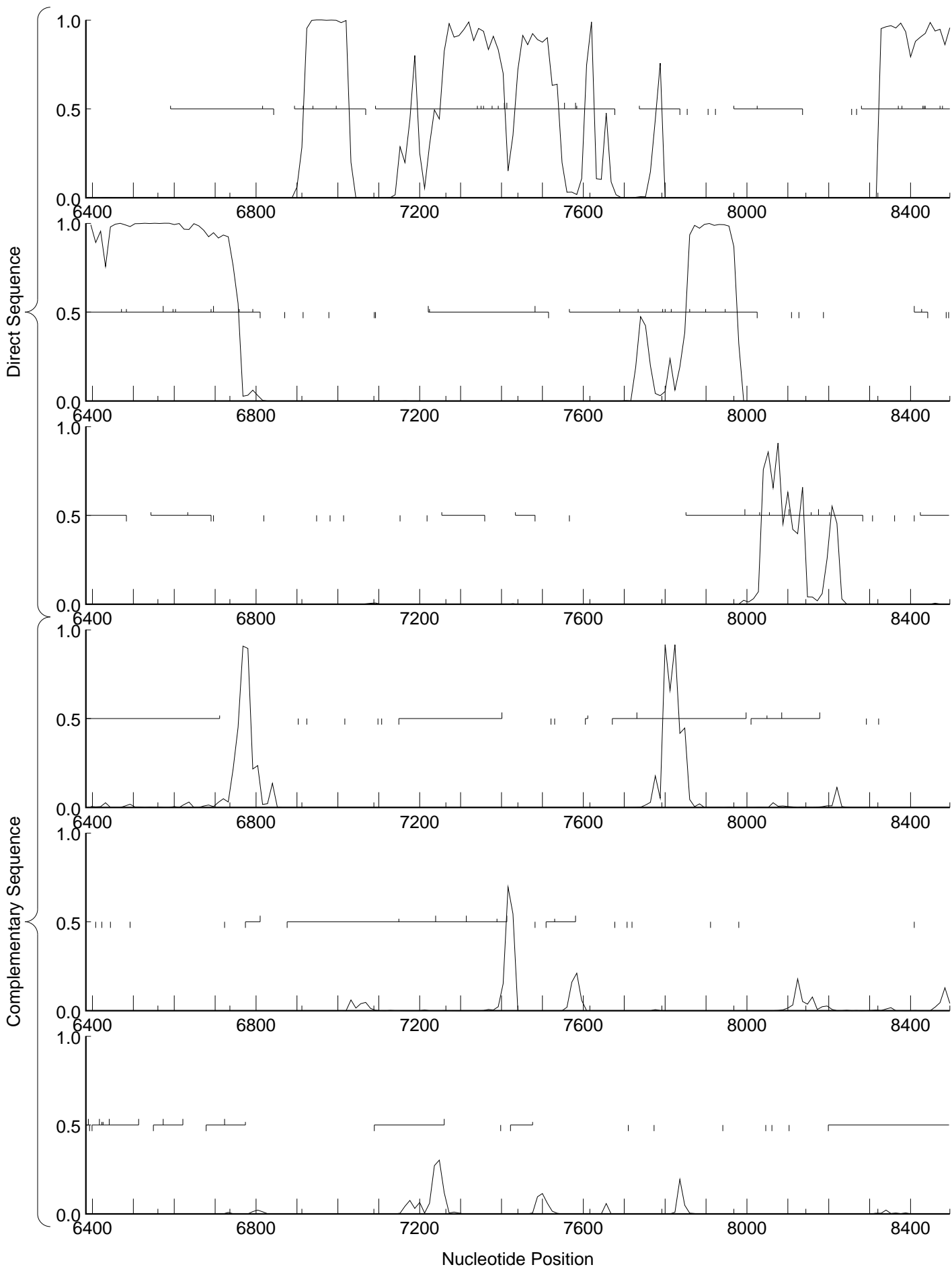
Matrix : -
Author : -
Order : 4

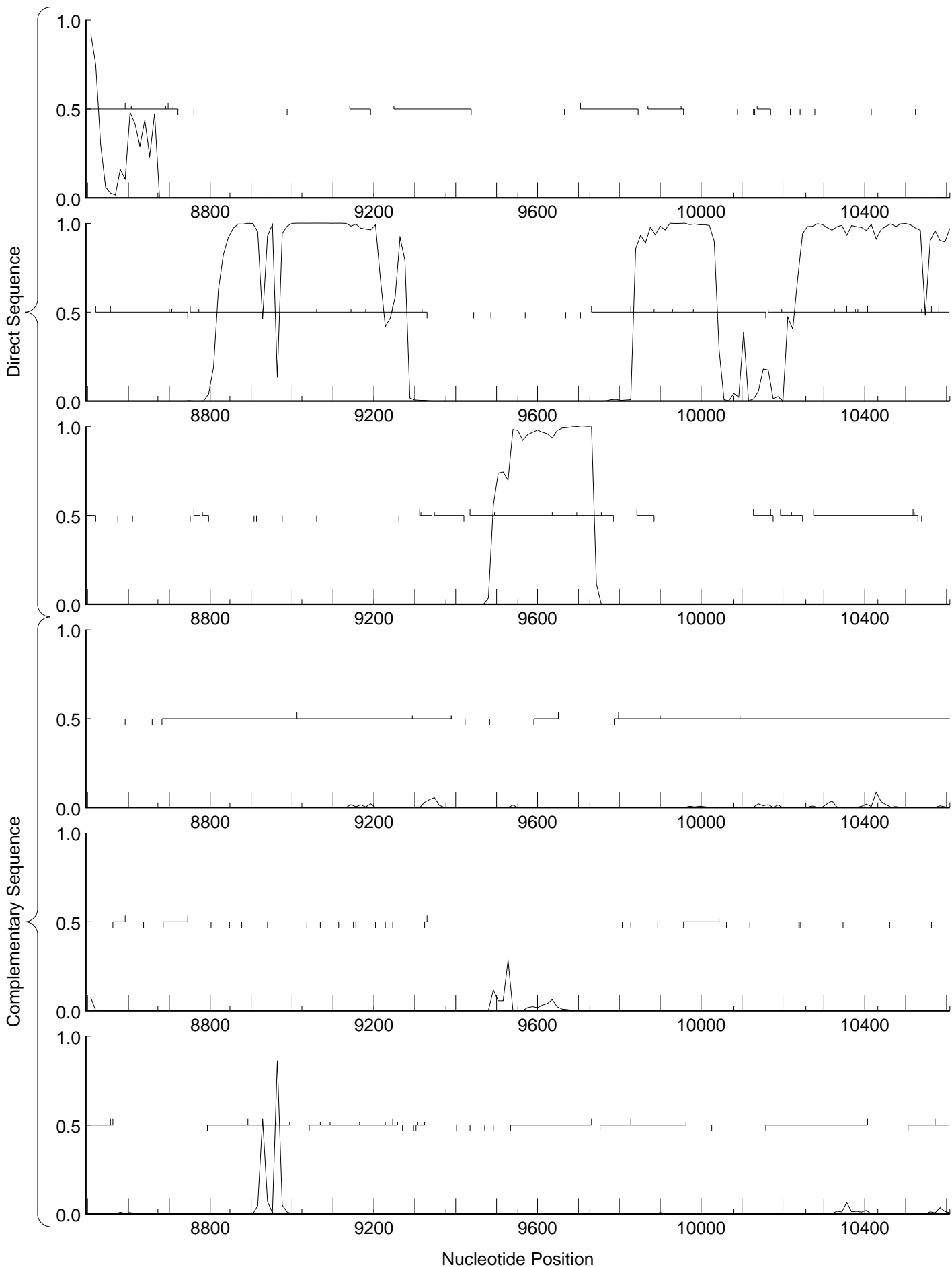
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

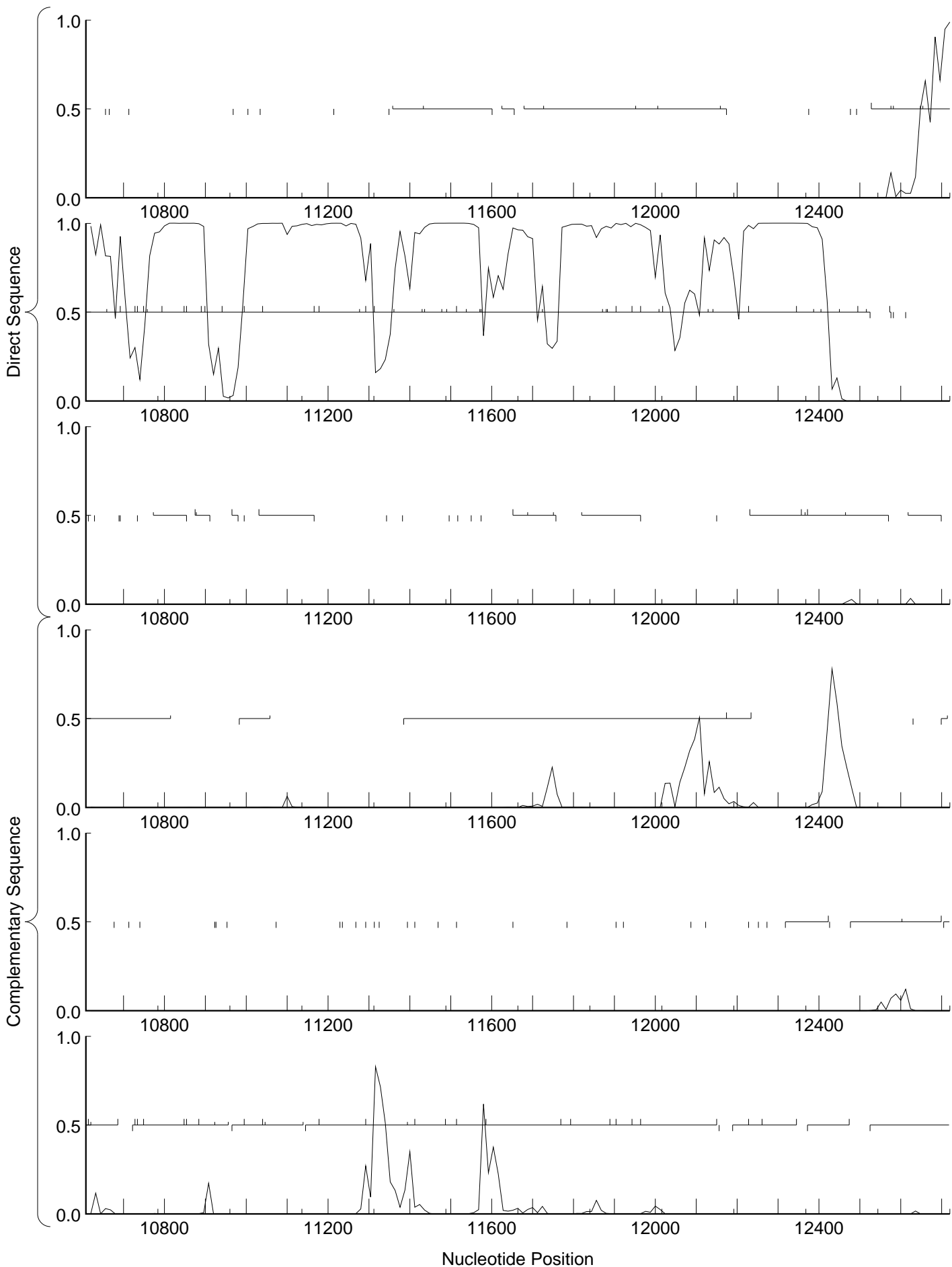


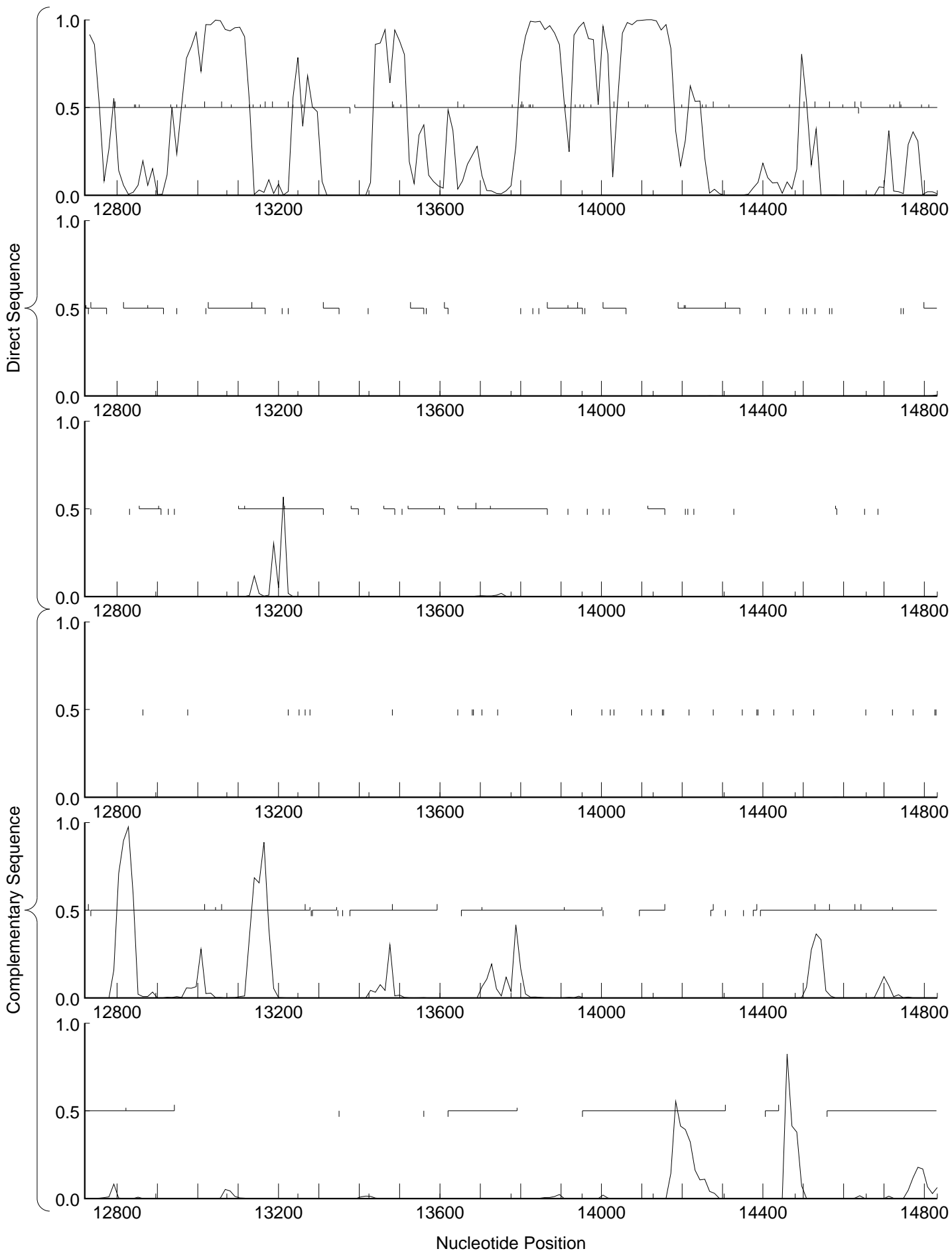


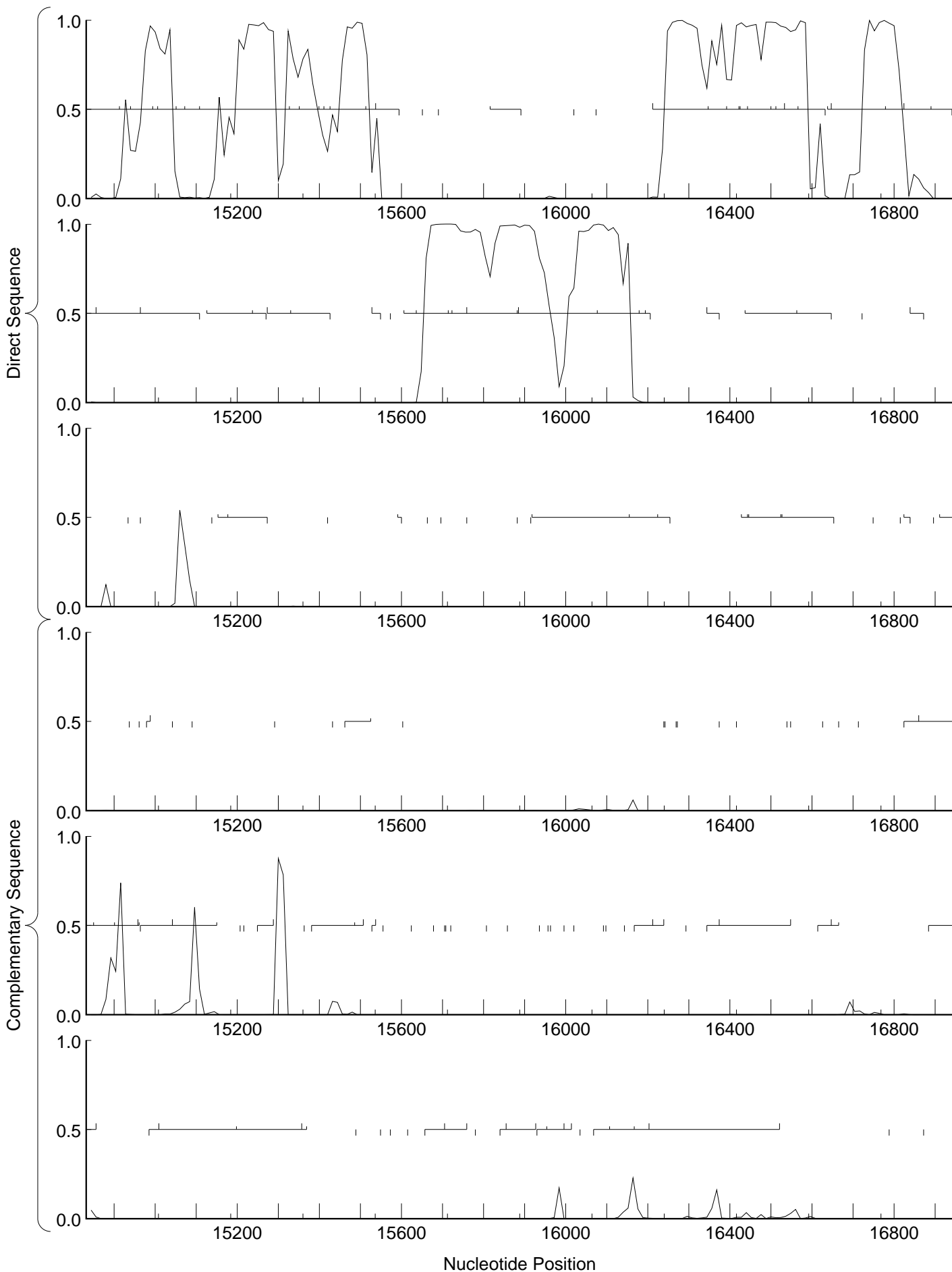


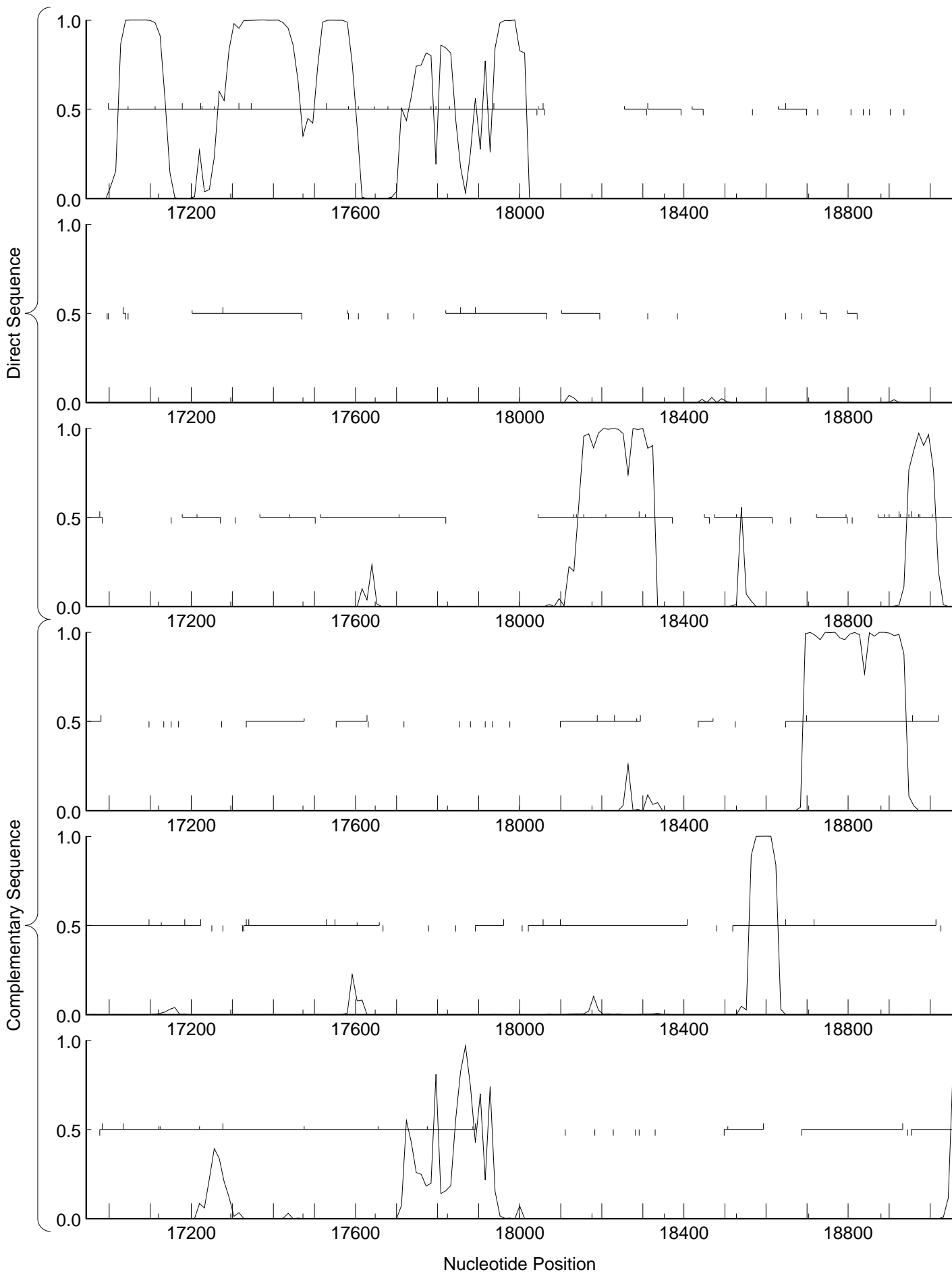


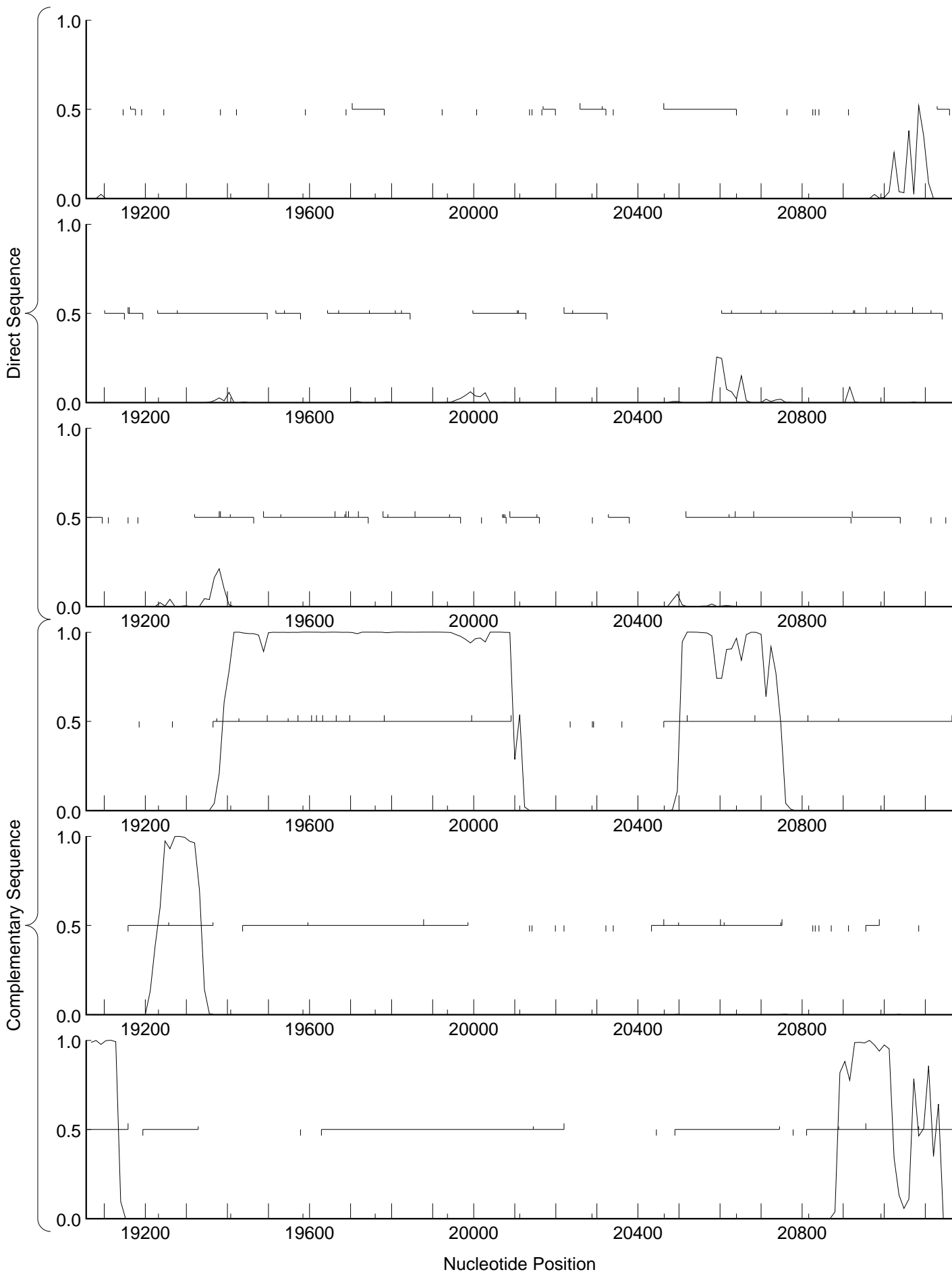


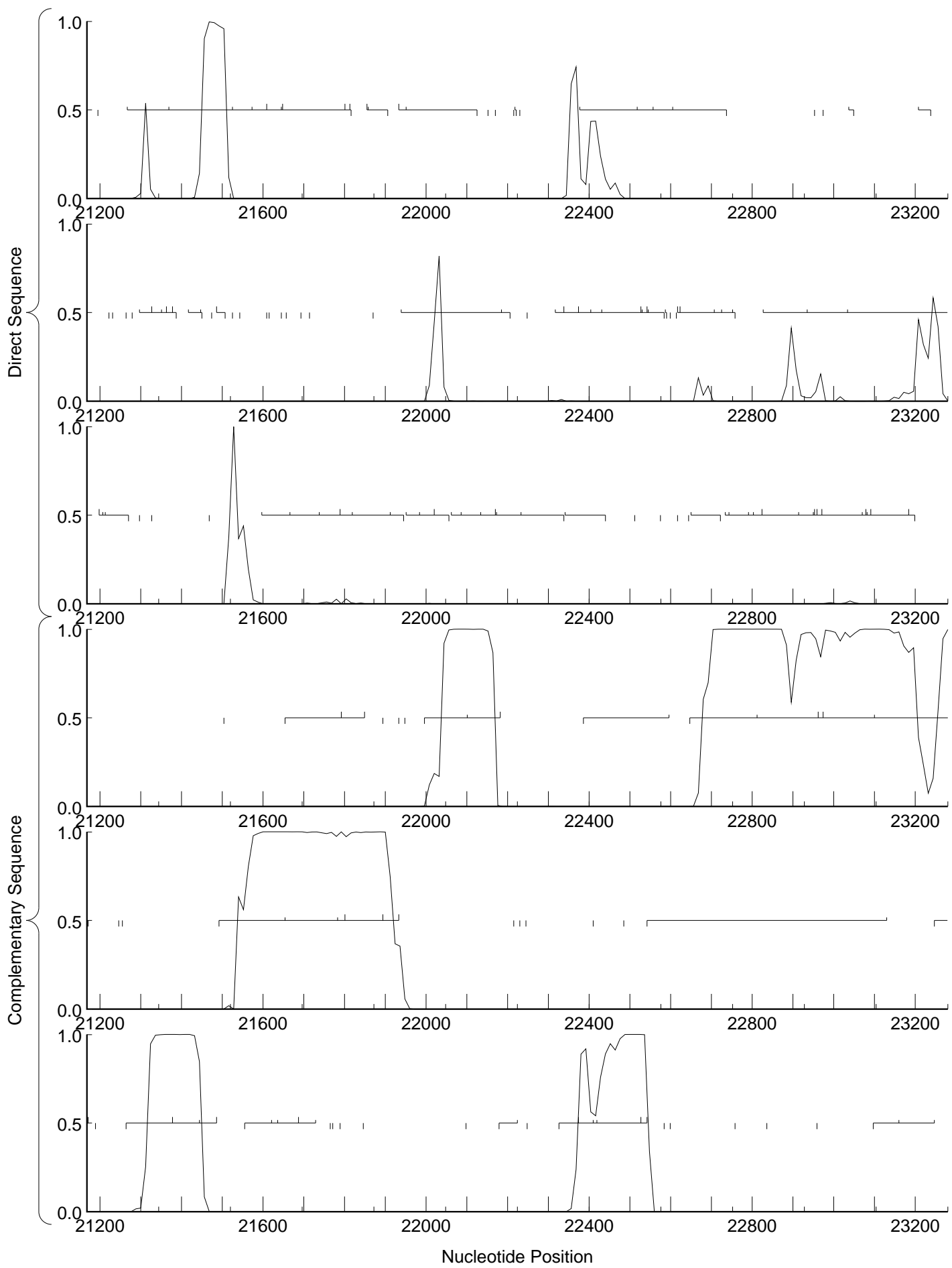


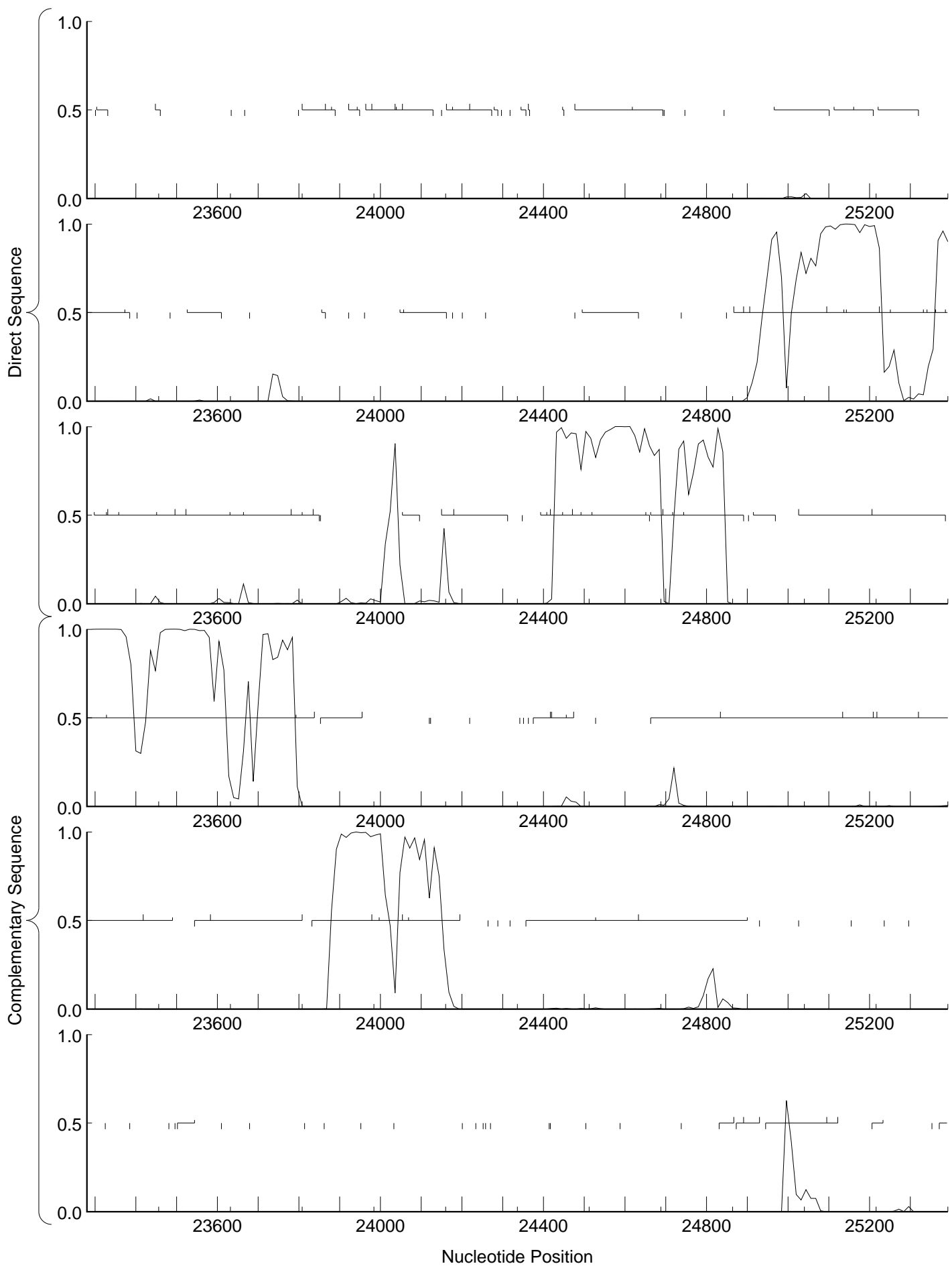


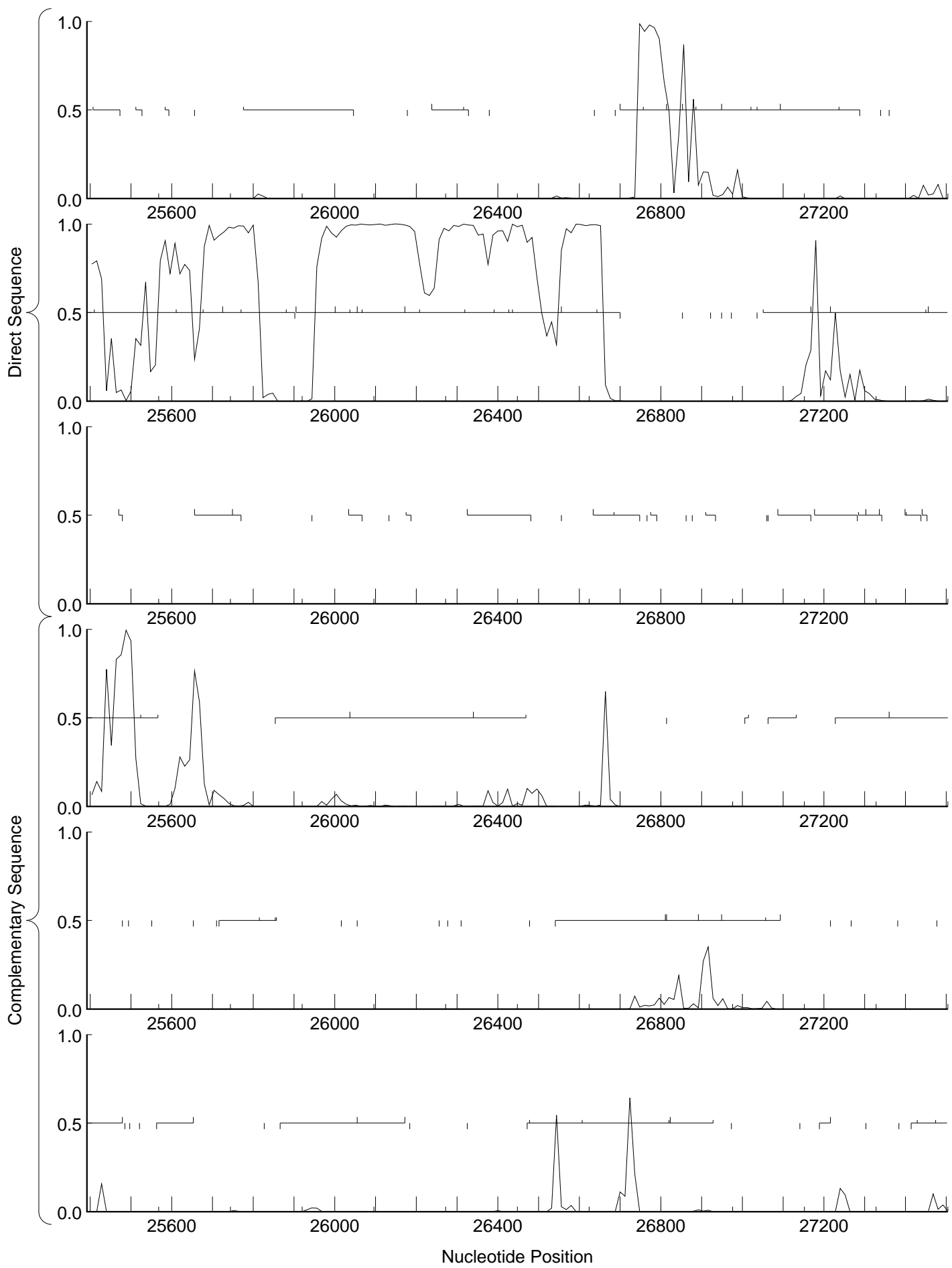


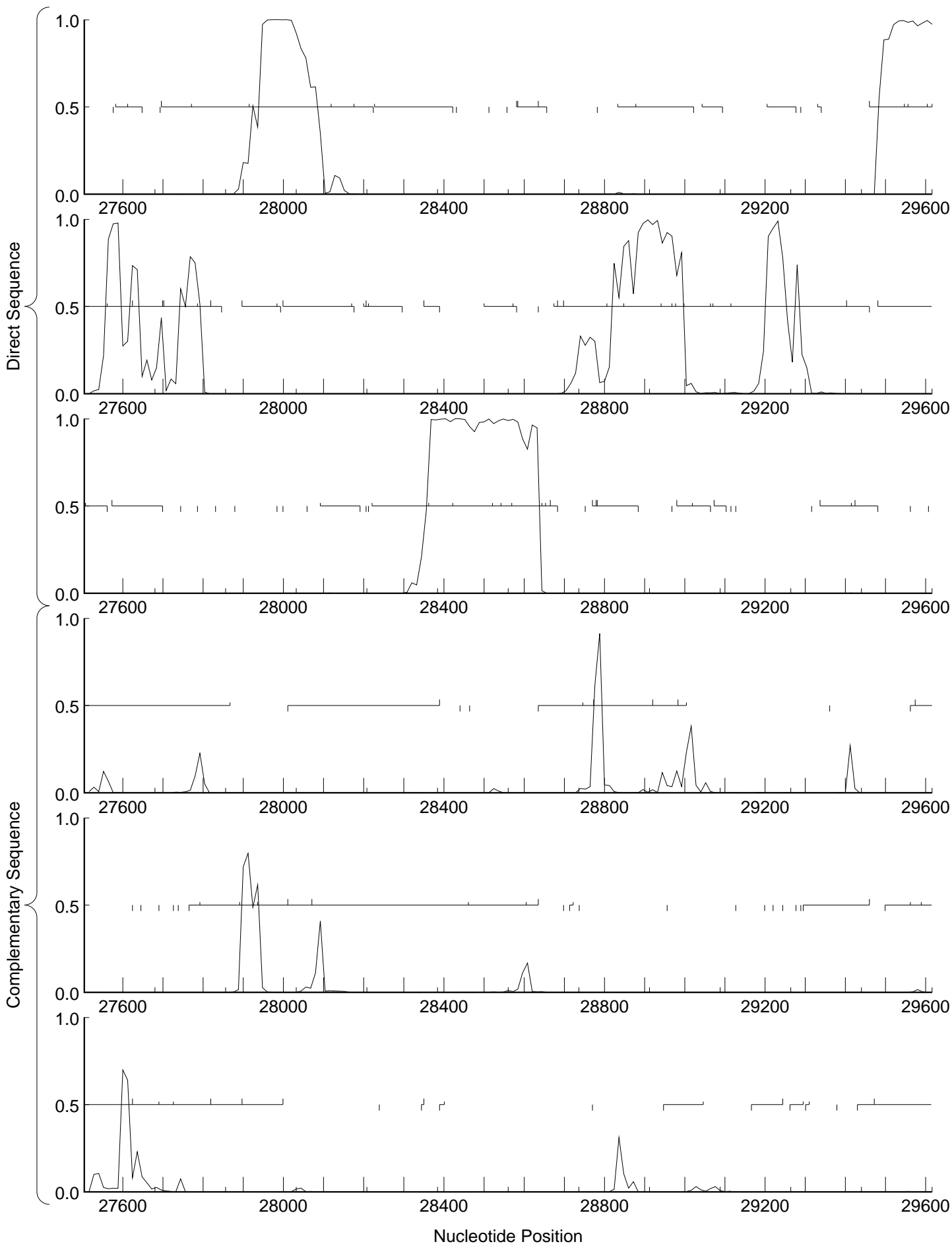












plete sequence, 38802 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS3, Order 4, Window 96, Step 12, 16/20

