

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Arthrobacter phage Cote complete sequence, 39044 bp including 12-base 3' overhang (GAGTTGCC
Analysis Date : 7/10/20 at 17:37:20
Pages : 20
Sequence Length : 39044 bp
GC Content : 66.72%

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,

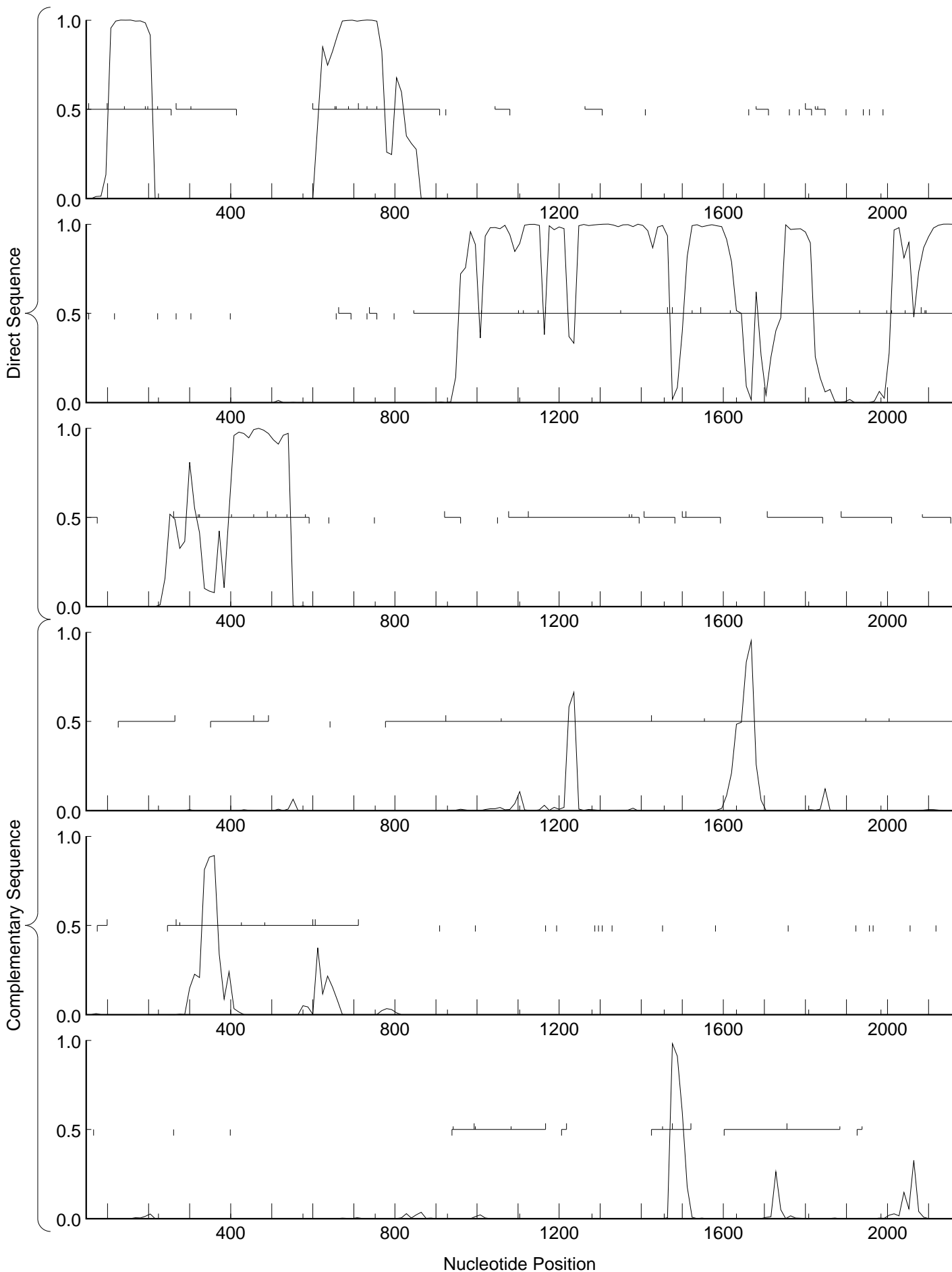
Matrix notes & comments

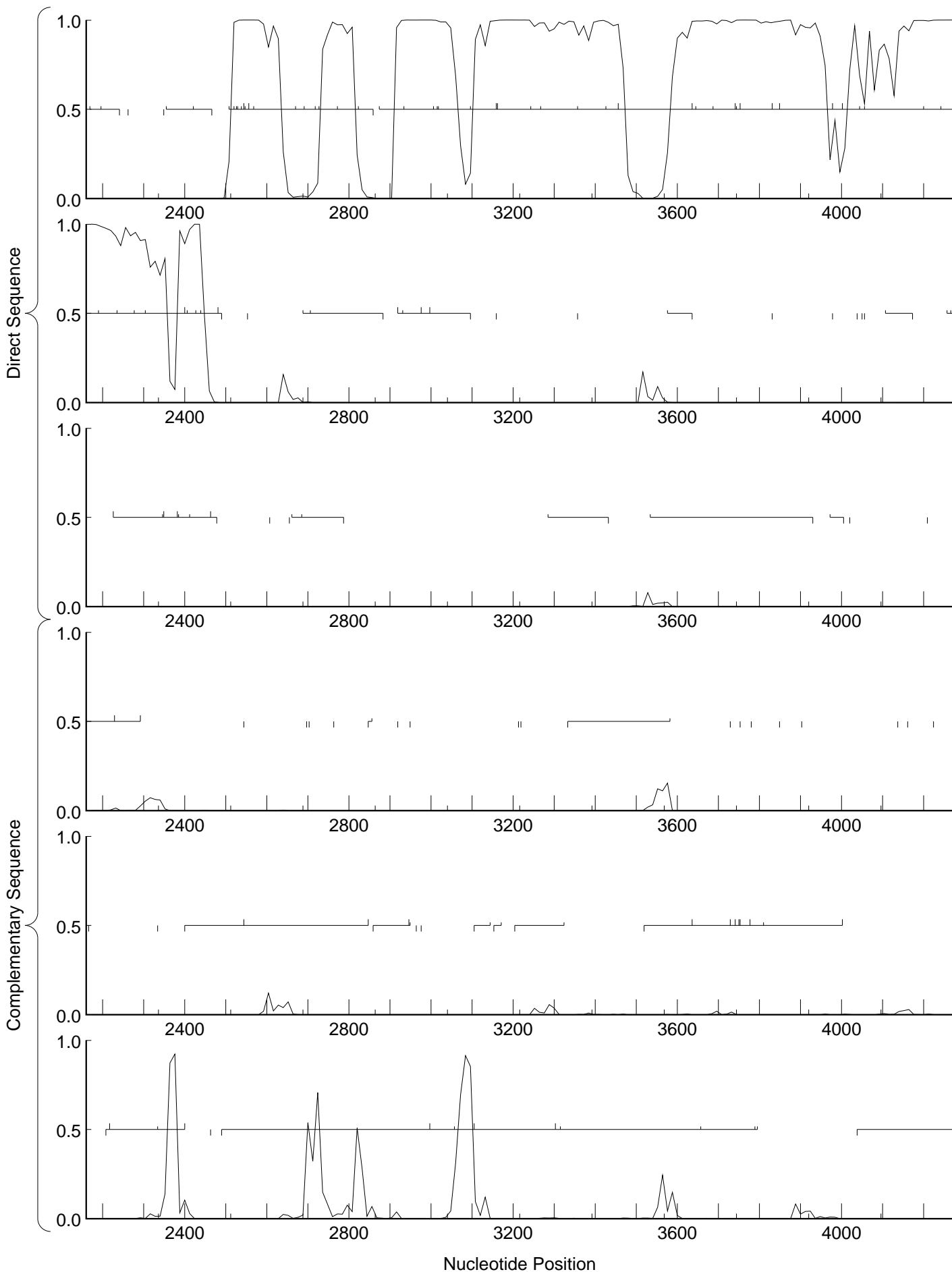
Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:22:00 2018

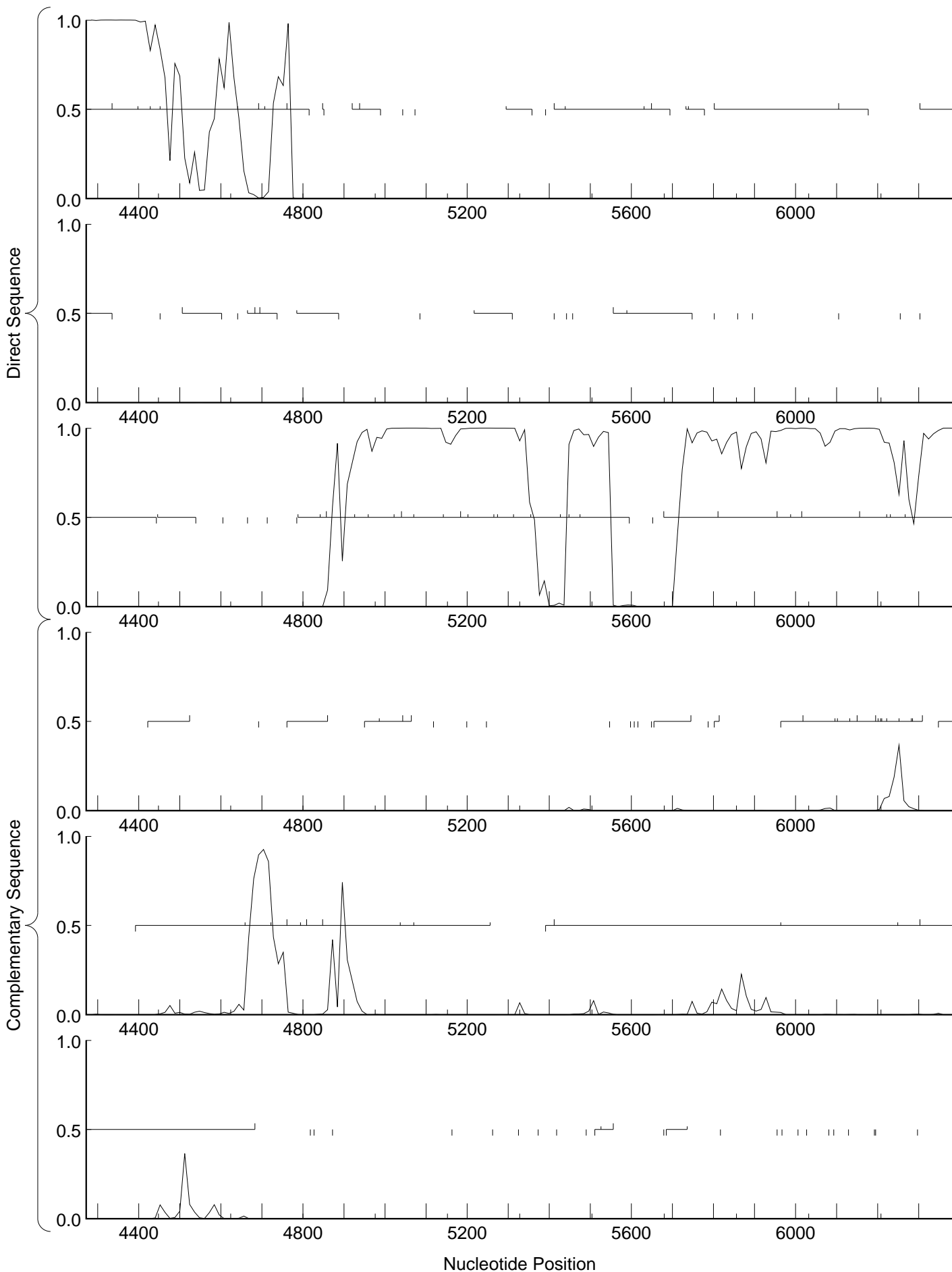
MATRIX INFORMATION

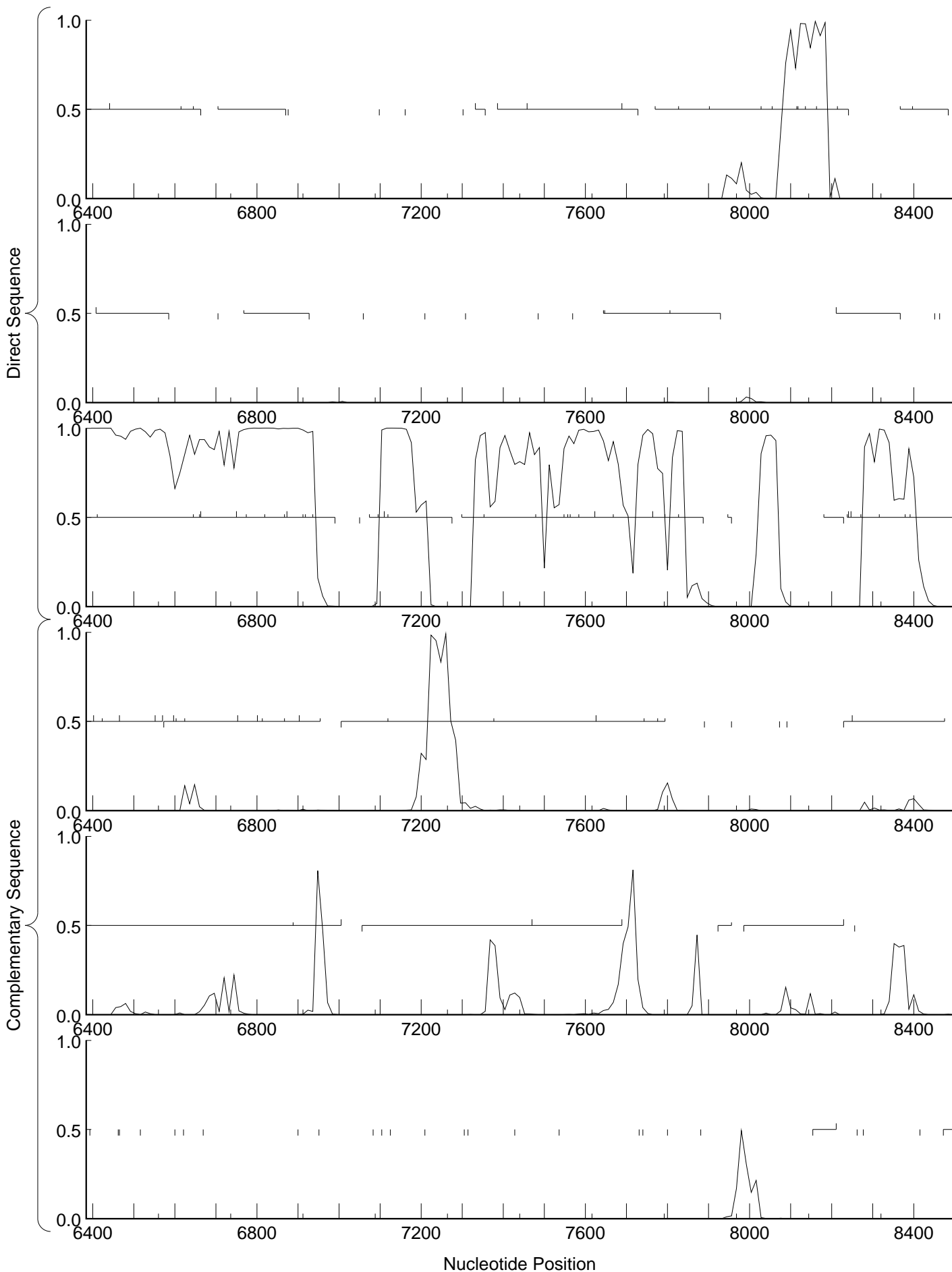
Matrix : -
Author : -
Order : 4

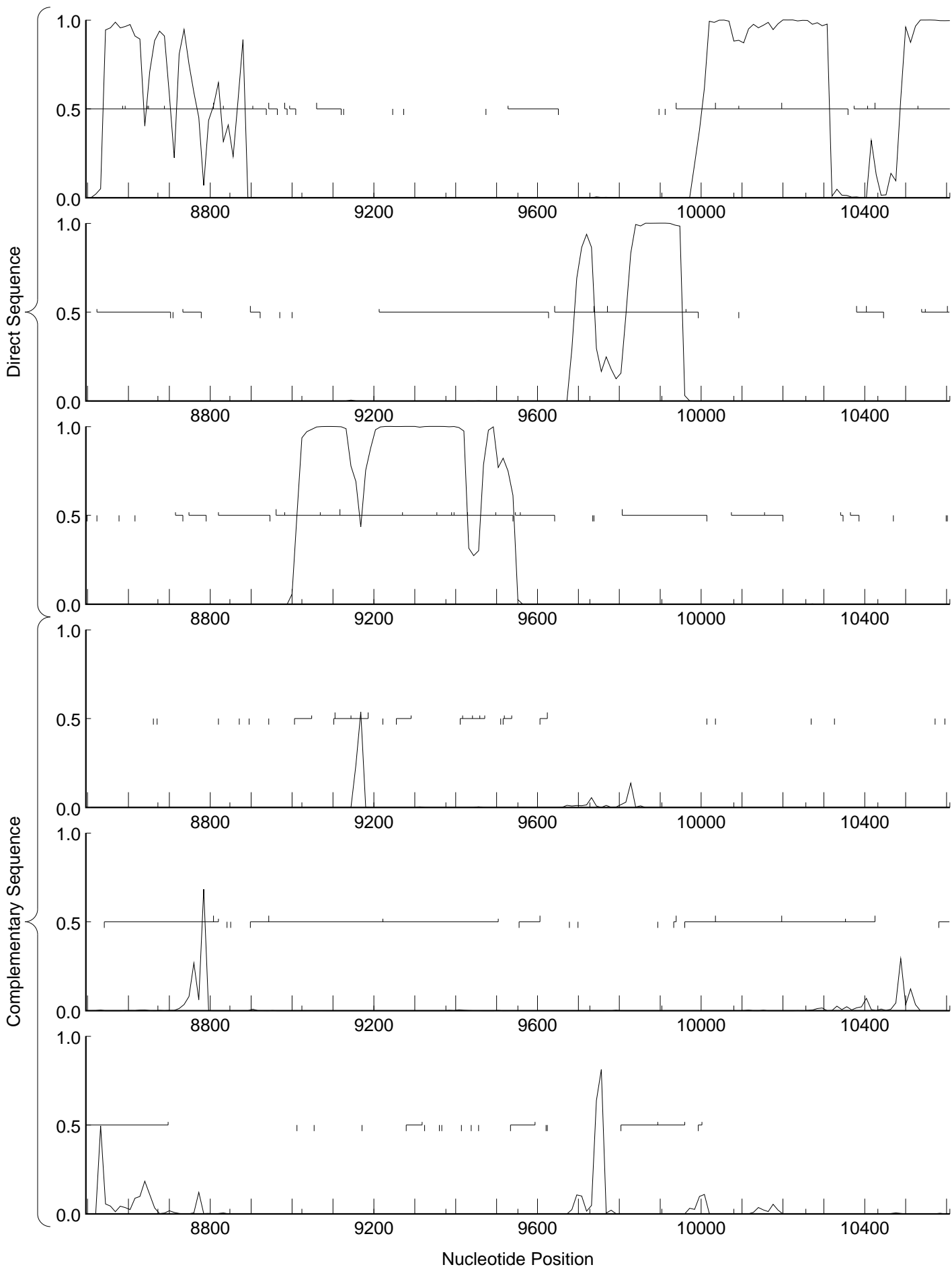
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

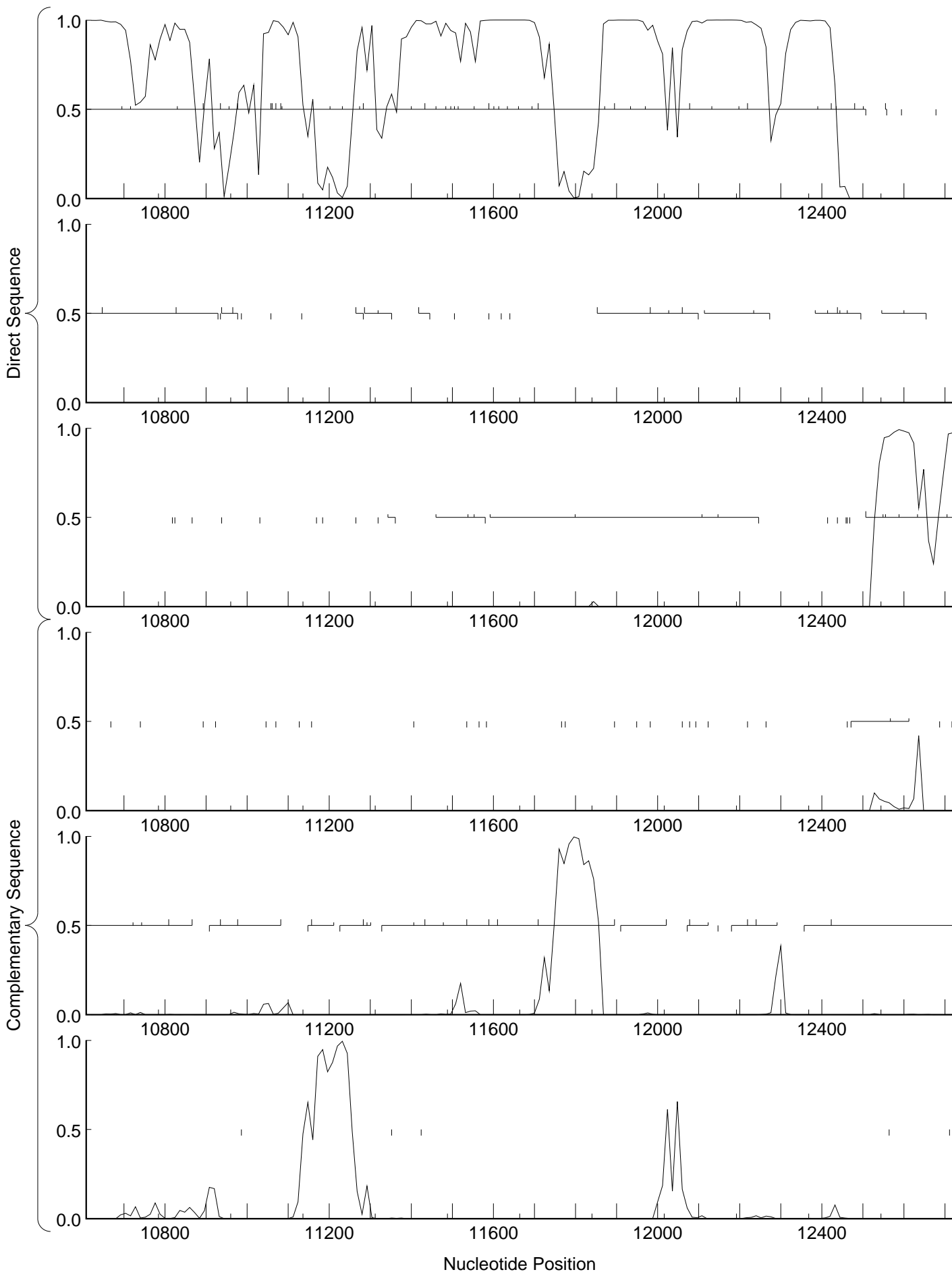


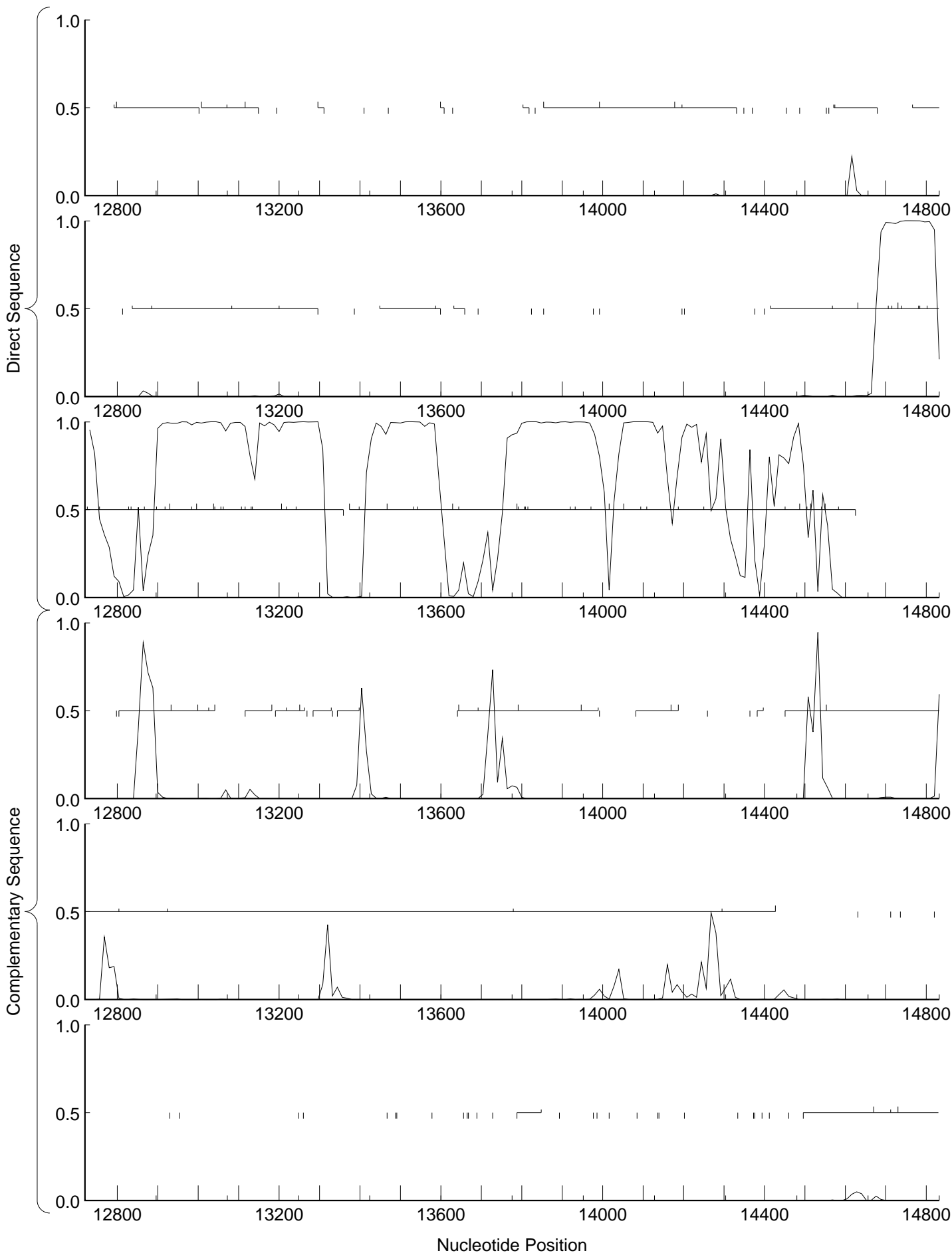


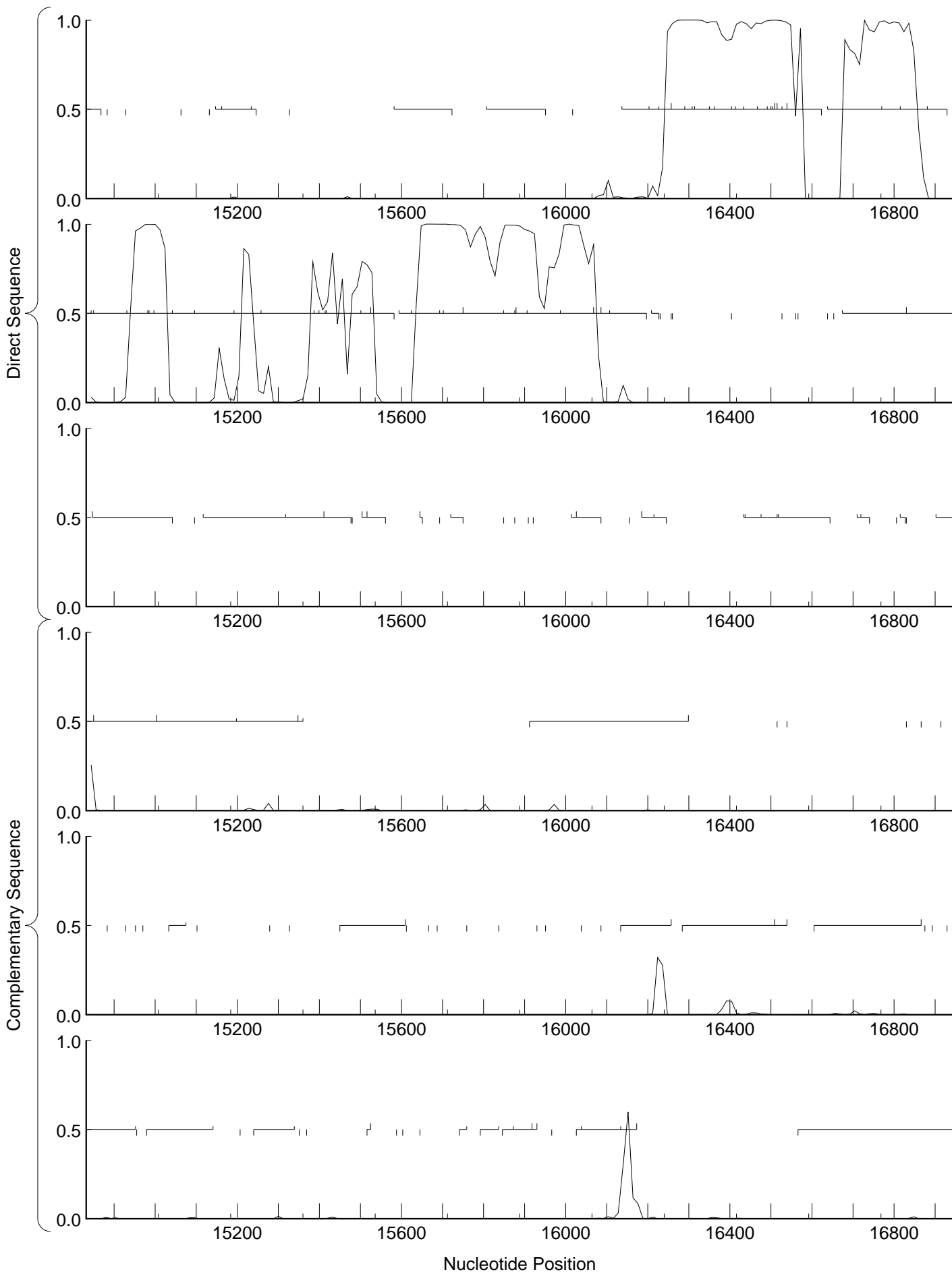


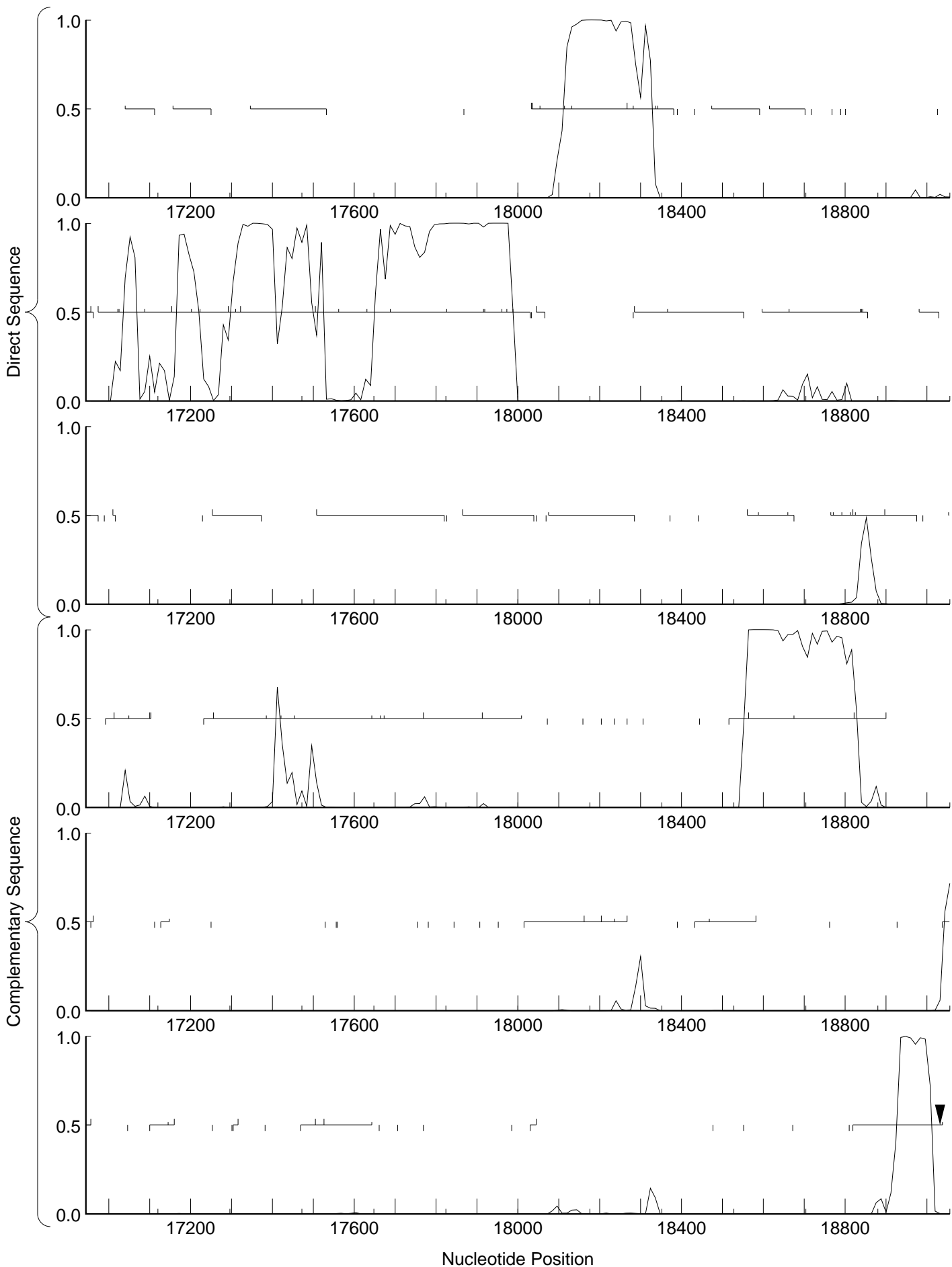


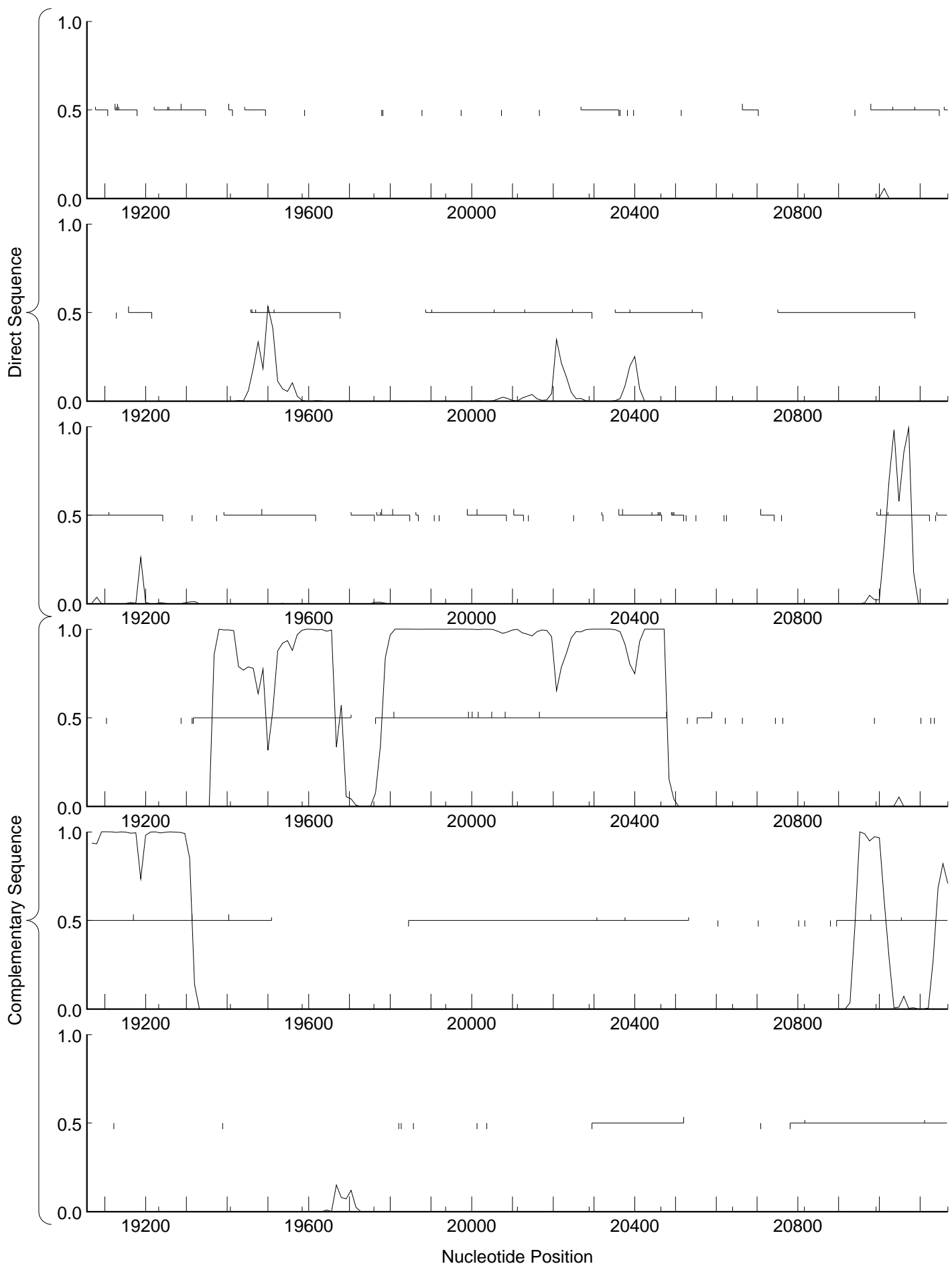


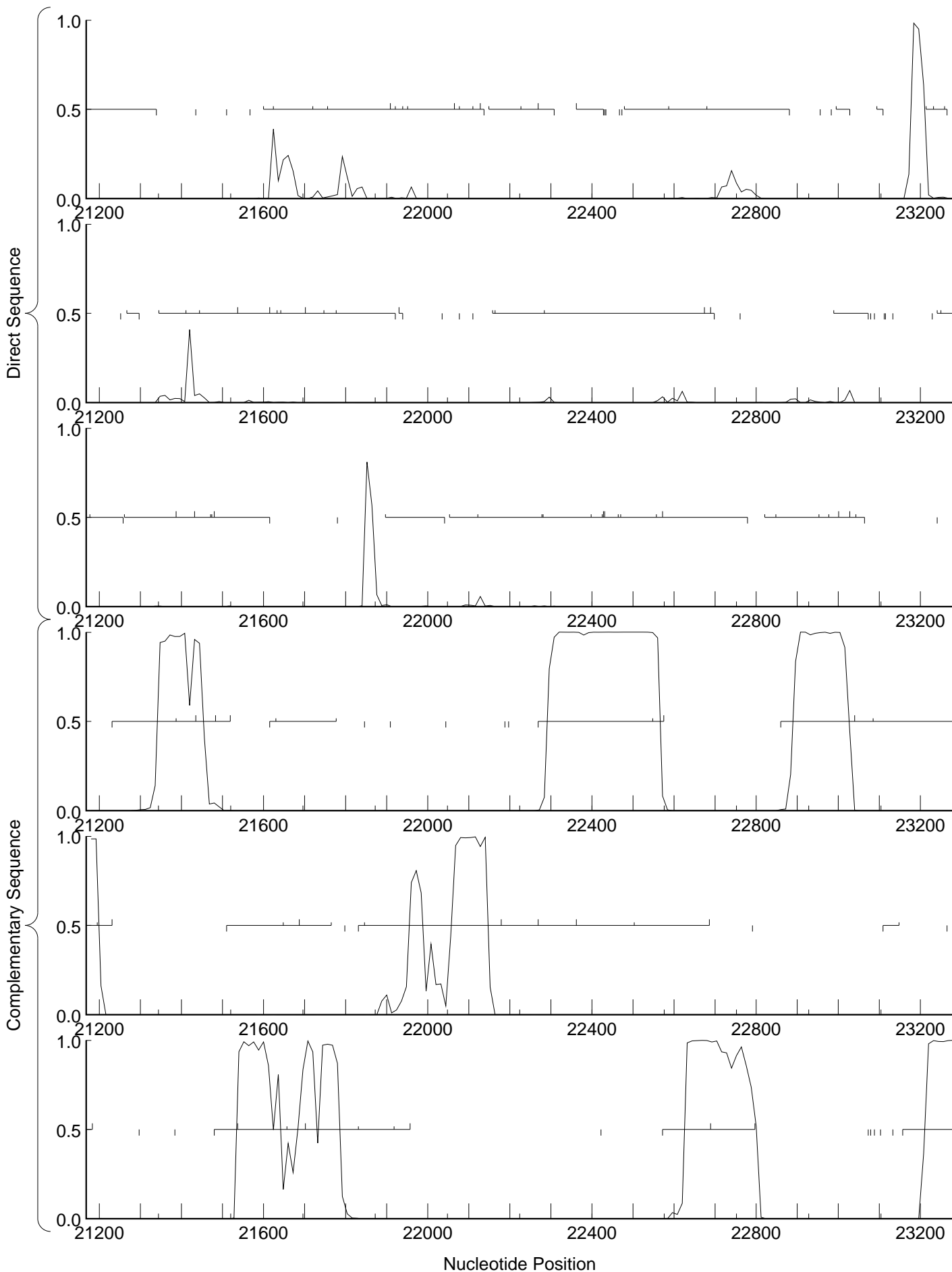




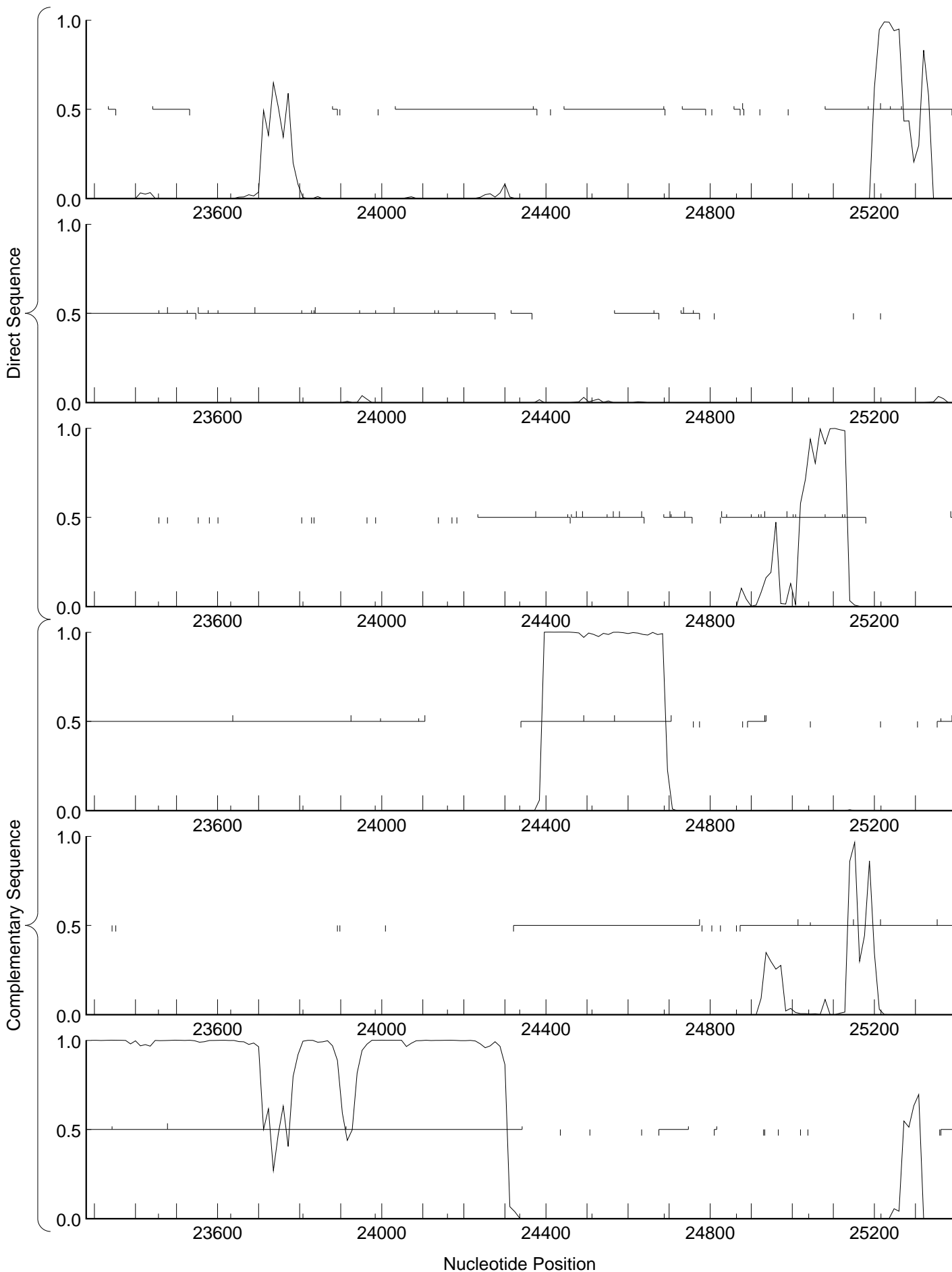




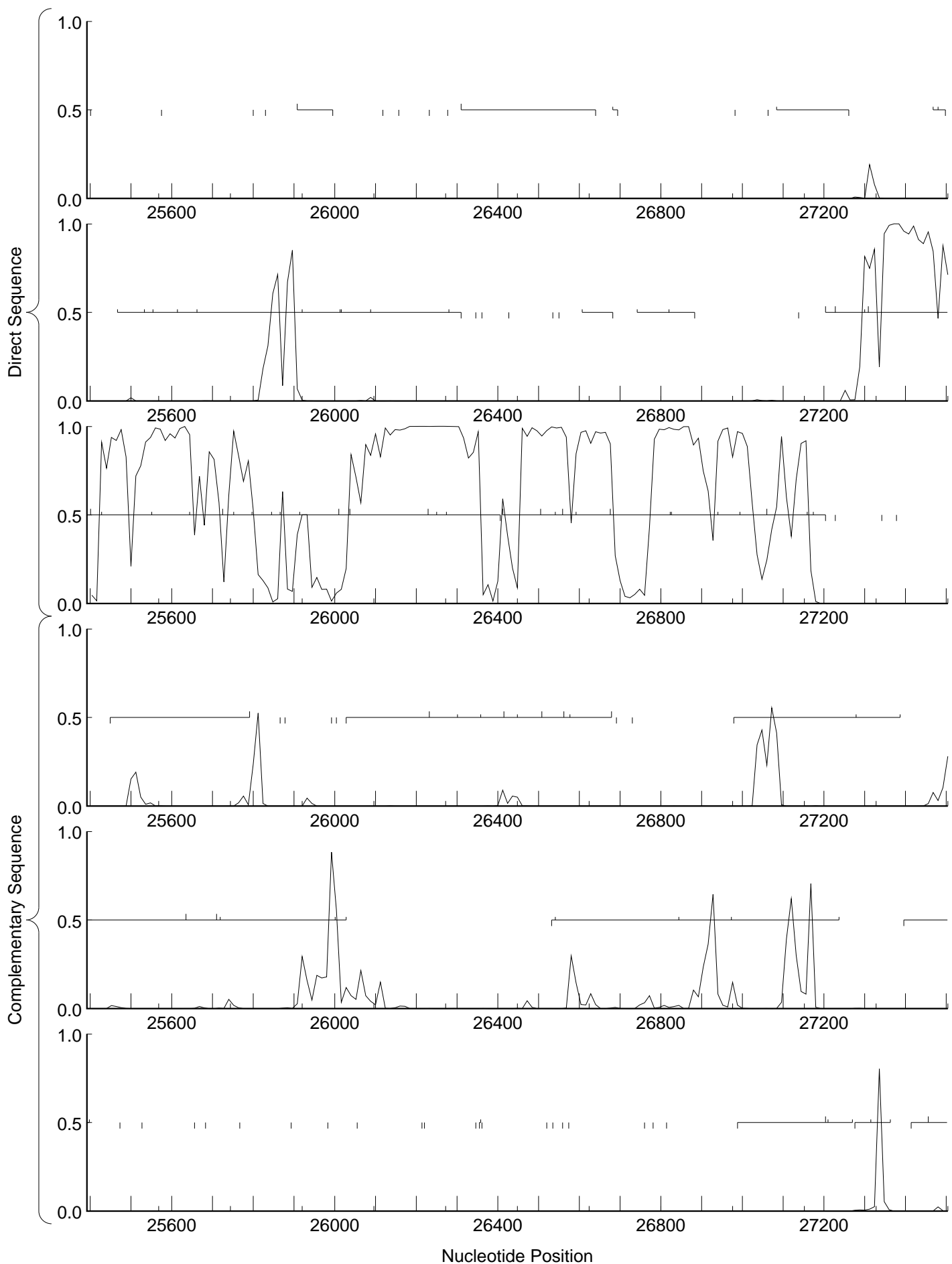




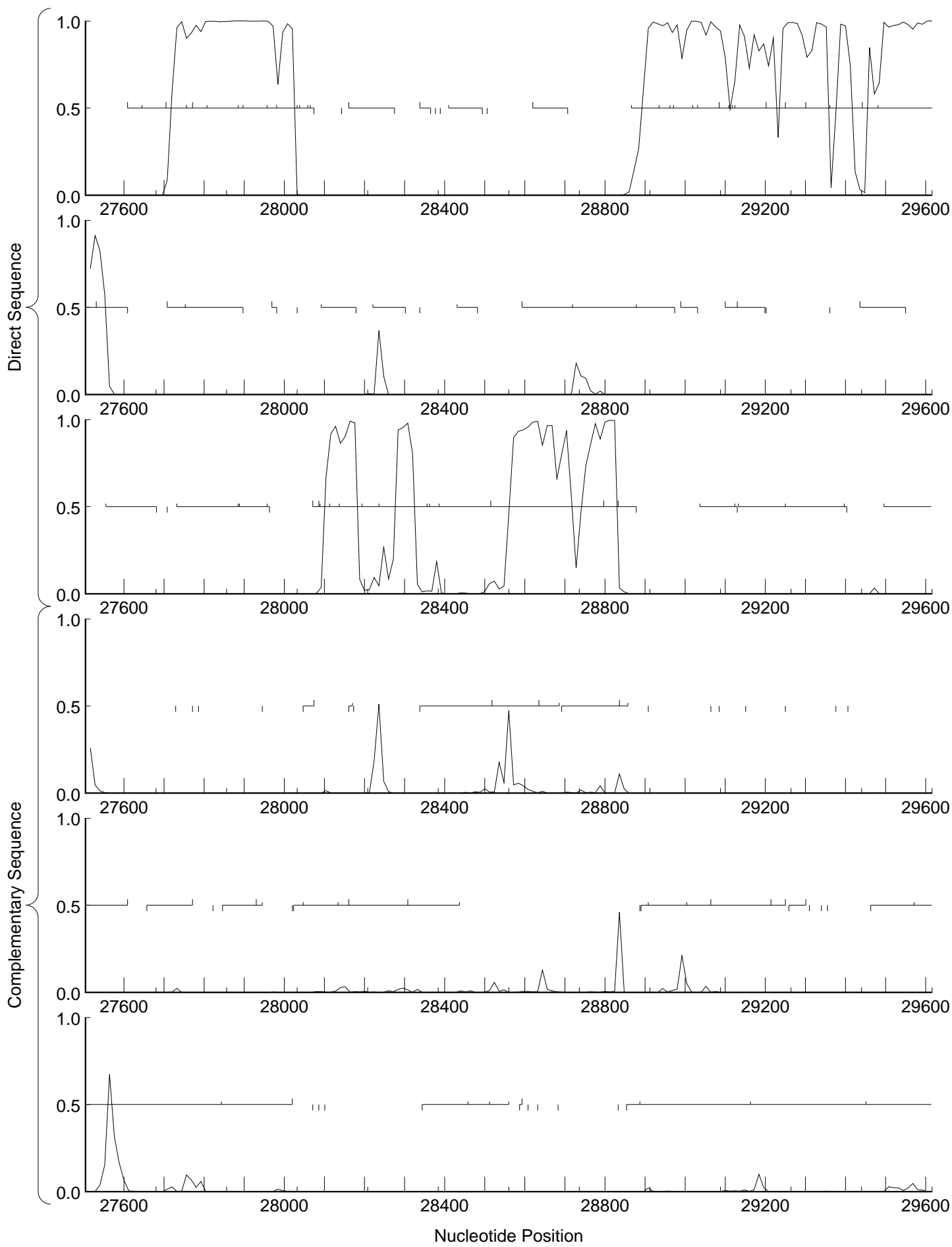
plete sequence, 39044 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 13/20

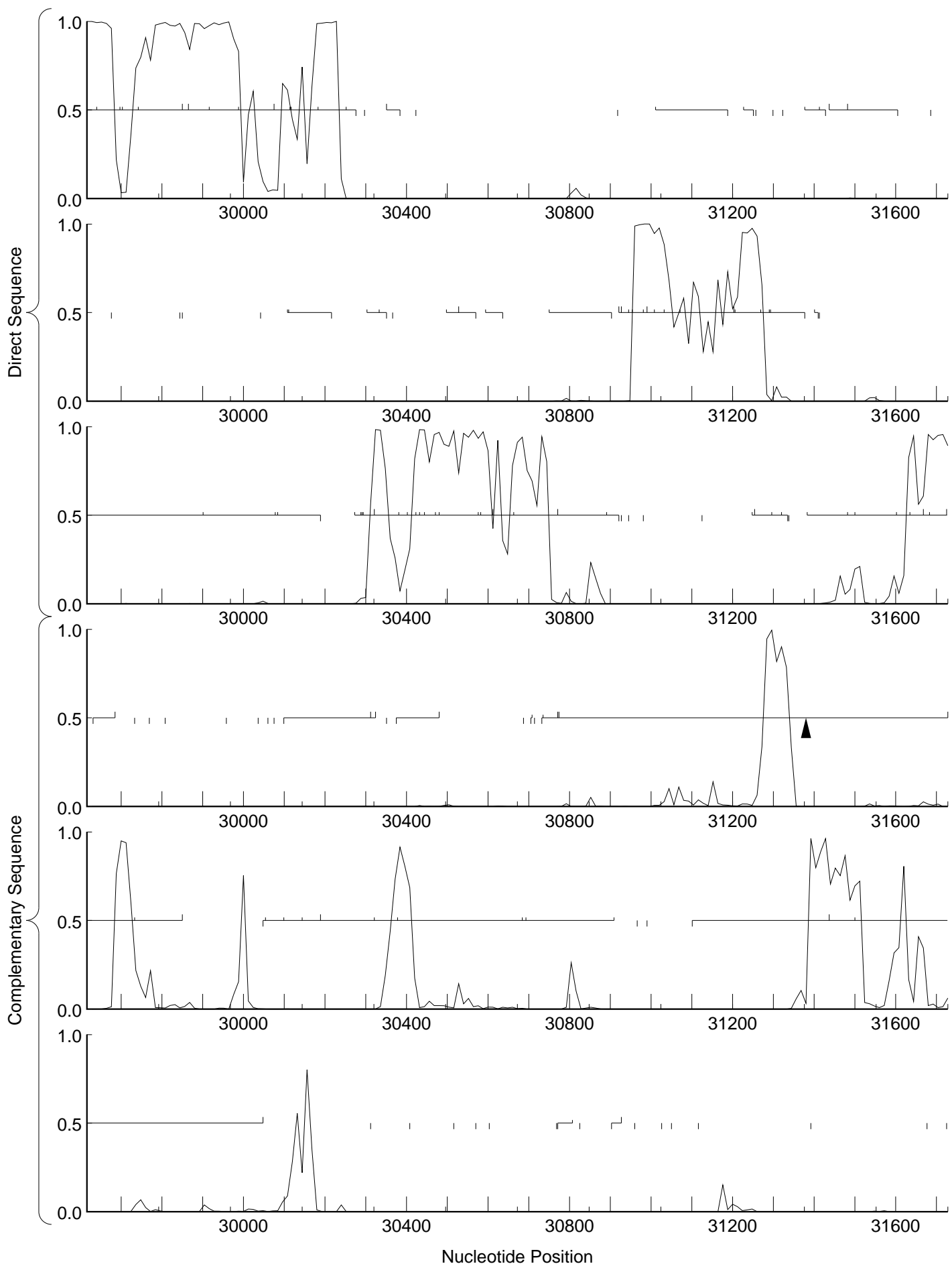


plete sequence, 39044 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 14/20



plete sequence, 39044 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 15/20





plete sequence, 39044 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 17/20

