

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Arthrobacter phage Kepler complete sequence, 38449 bp including 12-base 3' overhang (GAGTTG
Analysis Date : 10/19/17 at 17:47:26
Pages : 20
Sequence Length : 38449 bp
GC Content : 66.74%

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,

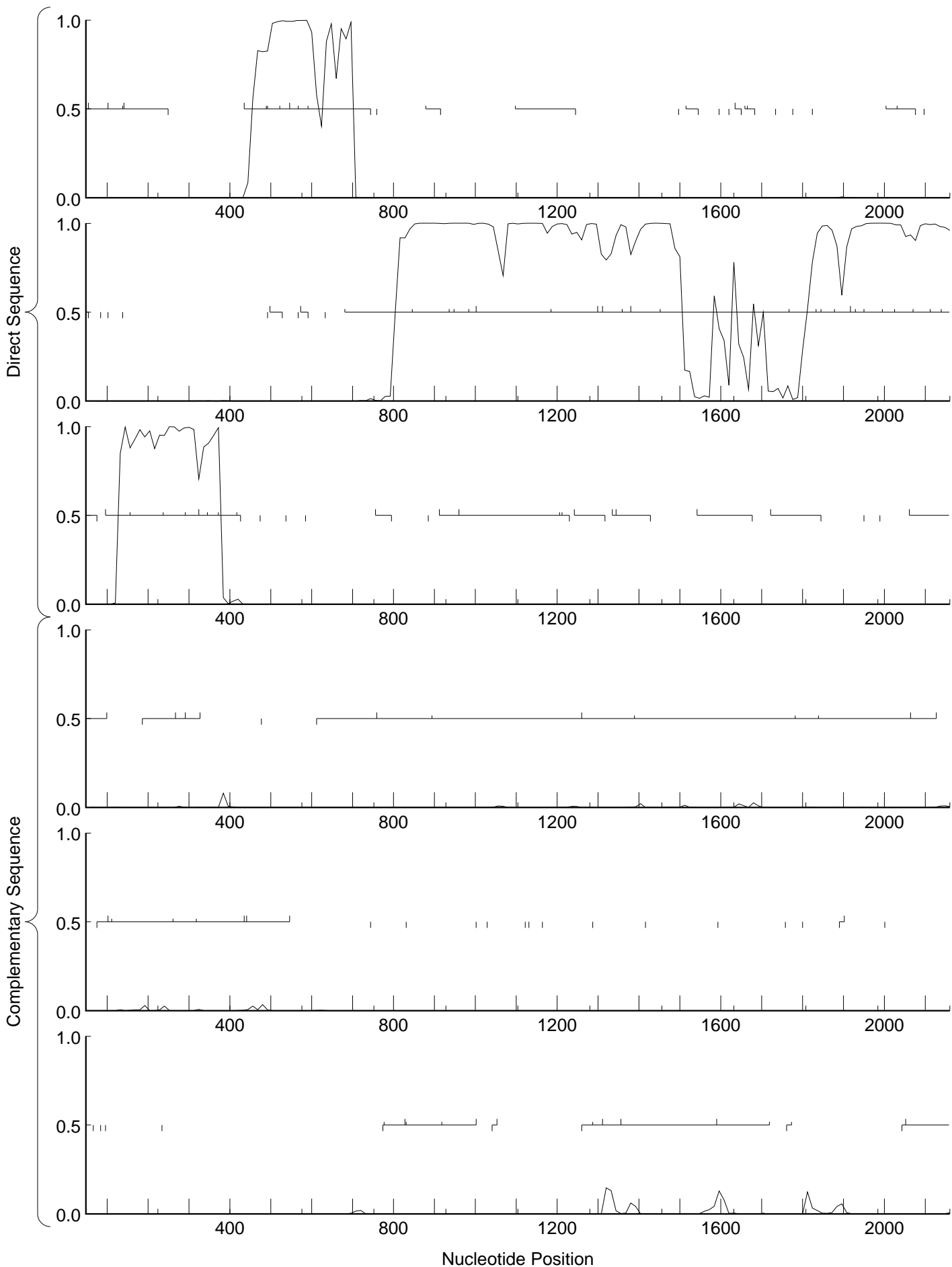
Matrix notes & comments

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

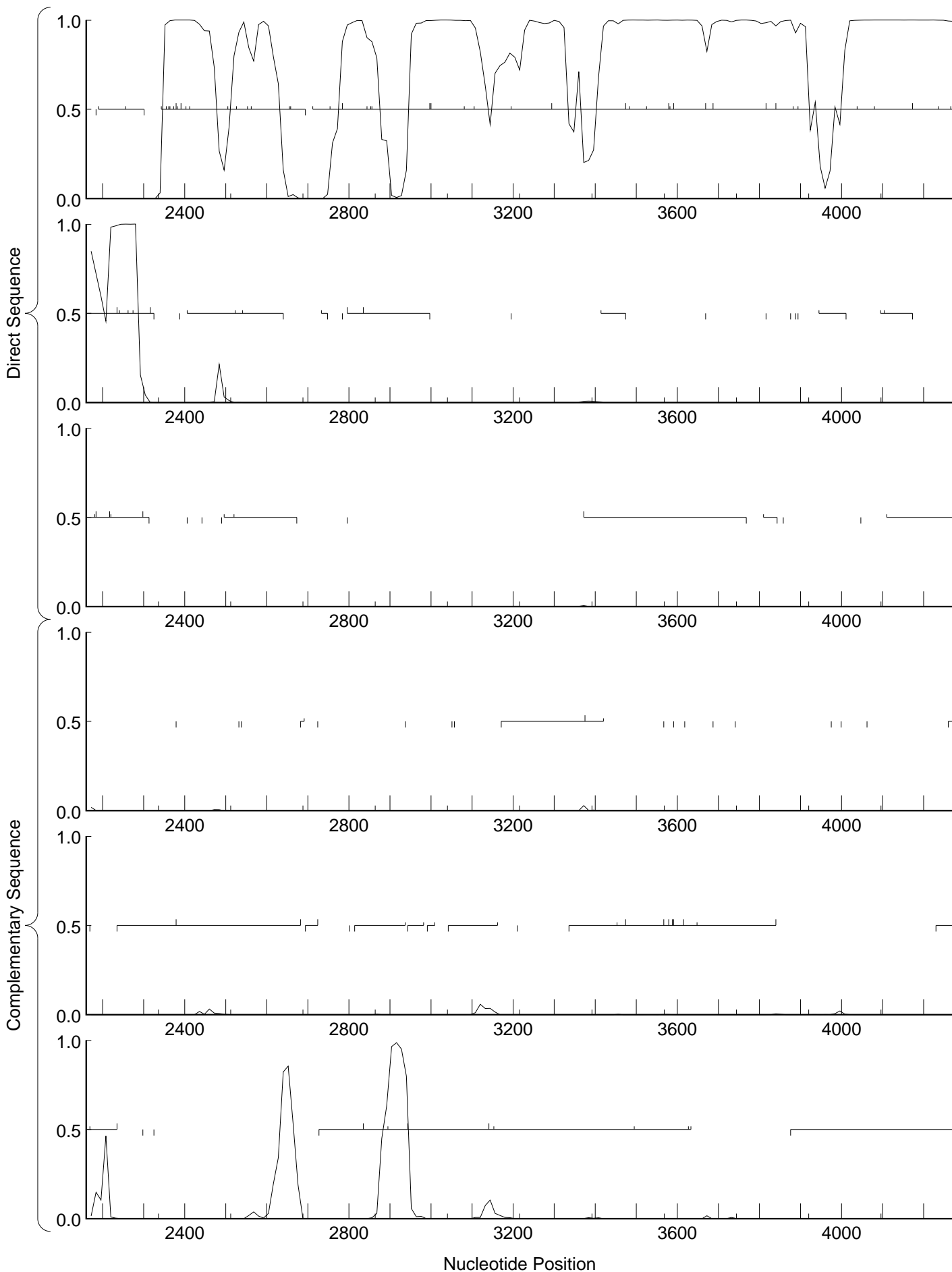
MATRIX INFORMATION

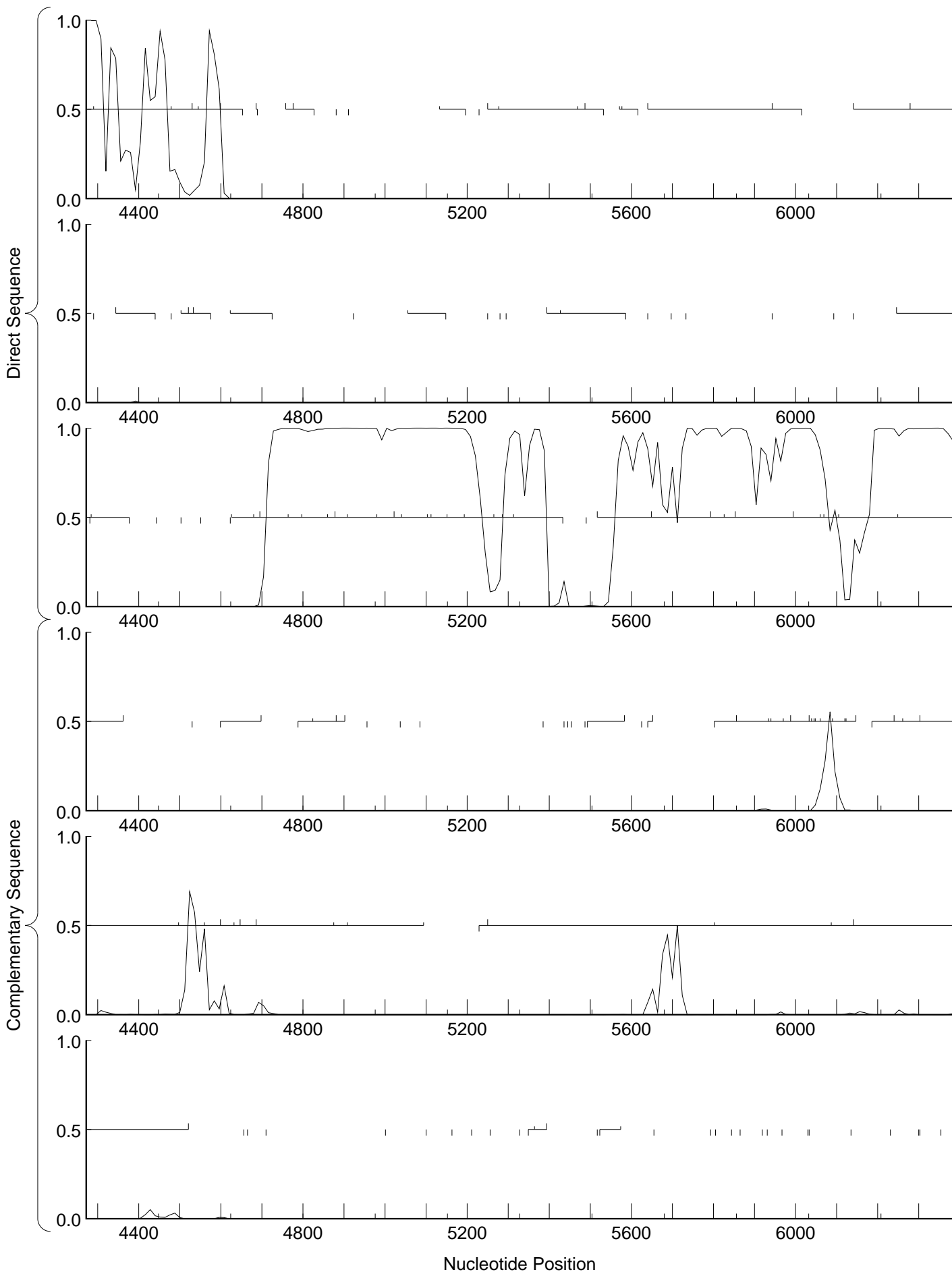
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

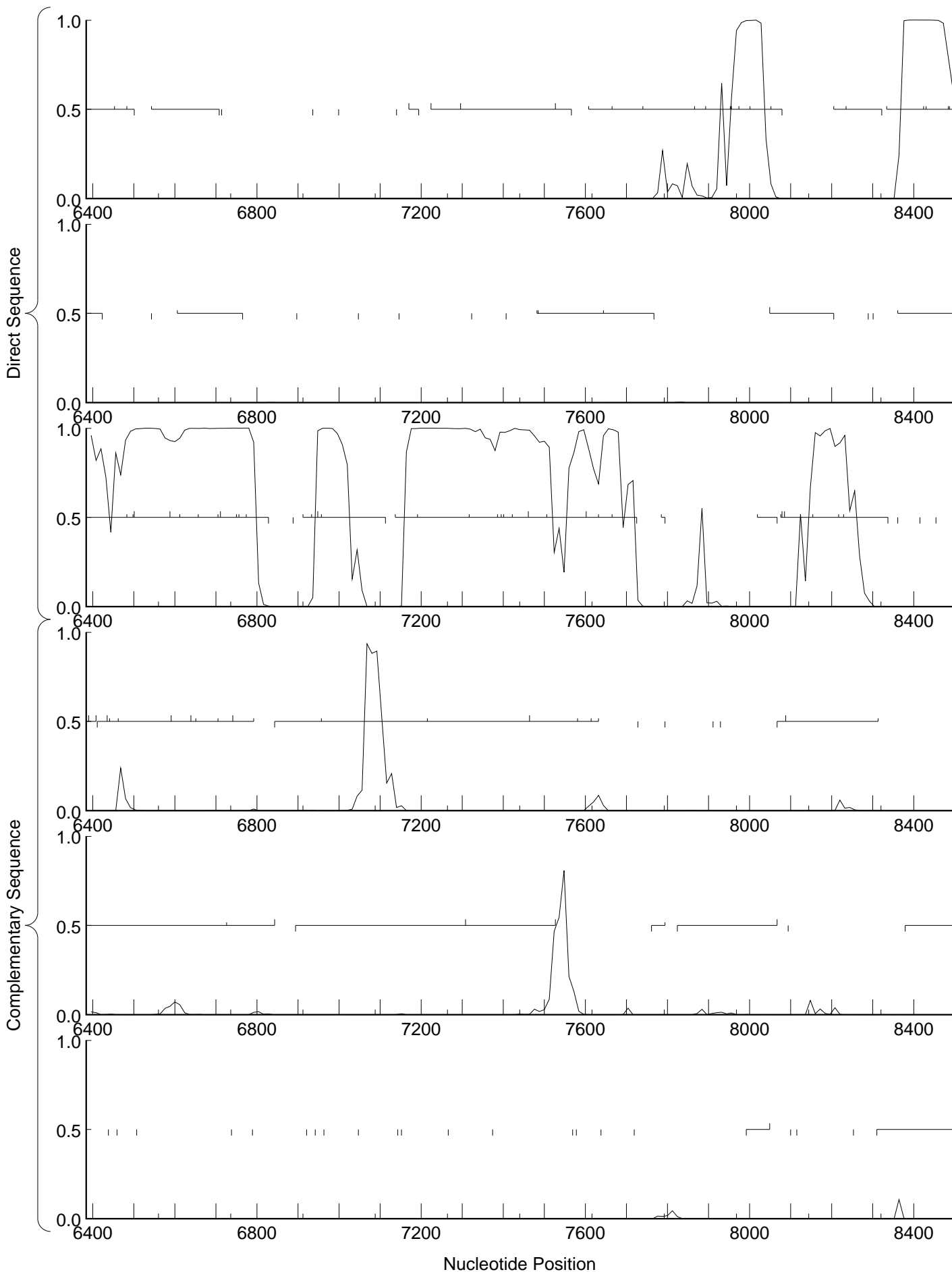
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

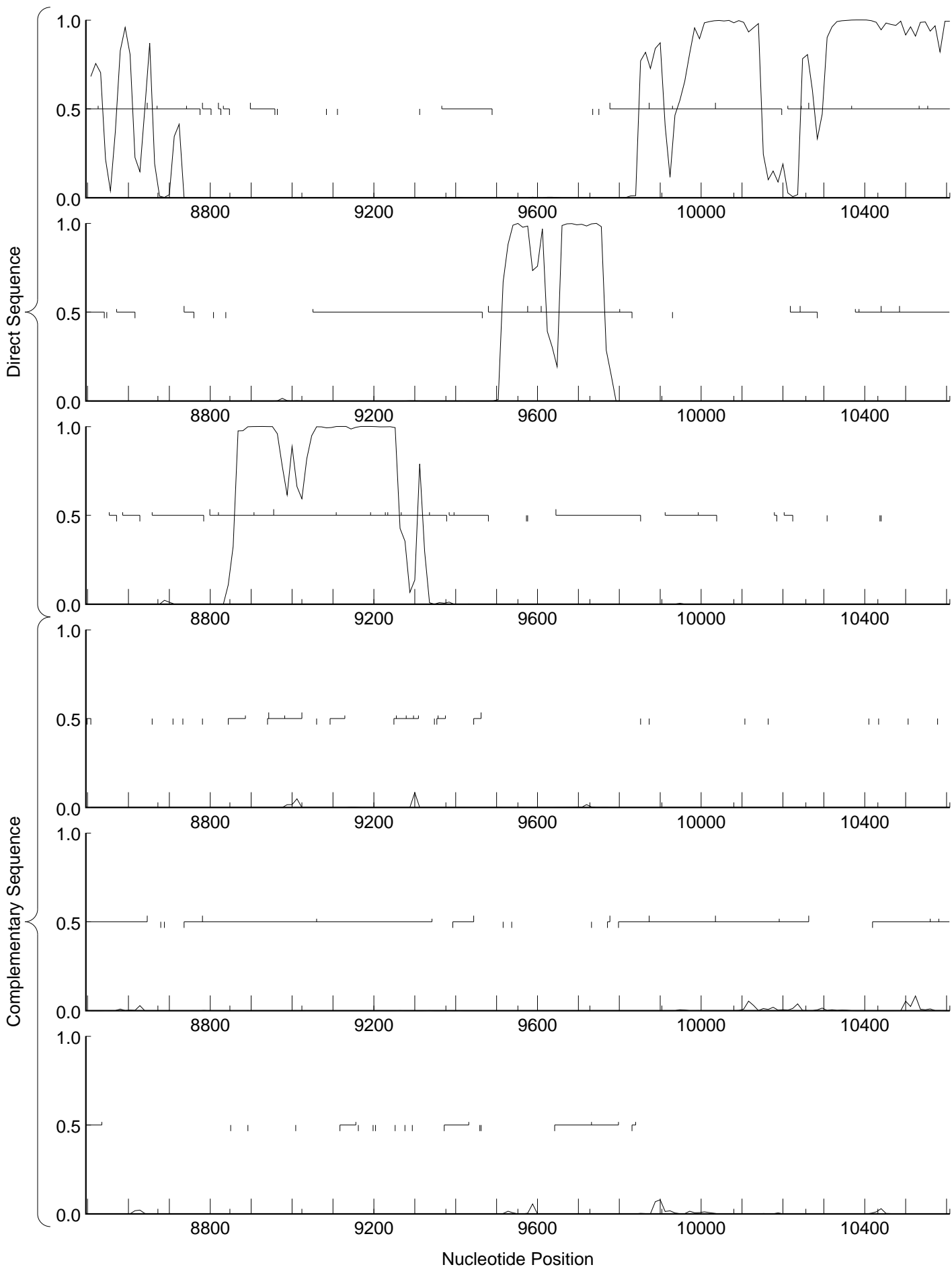


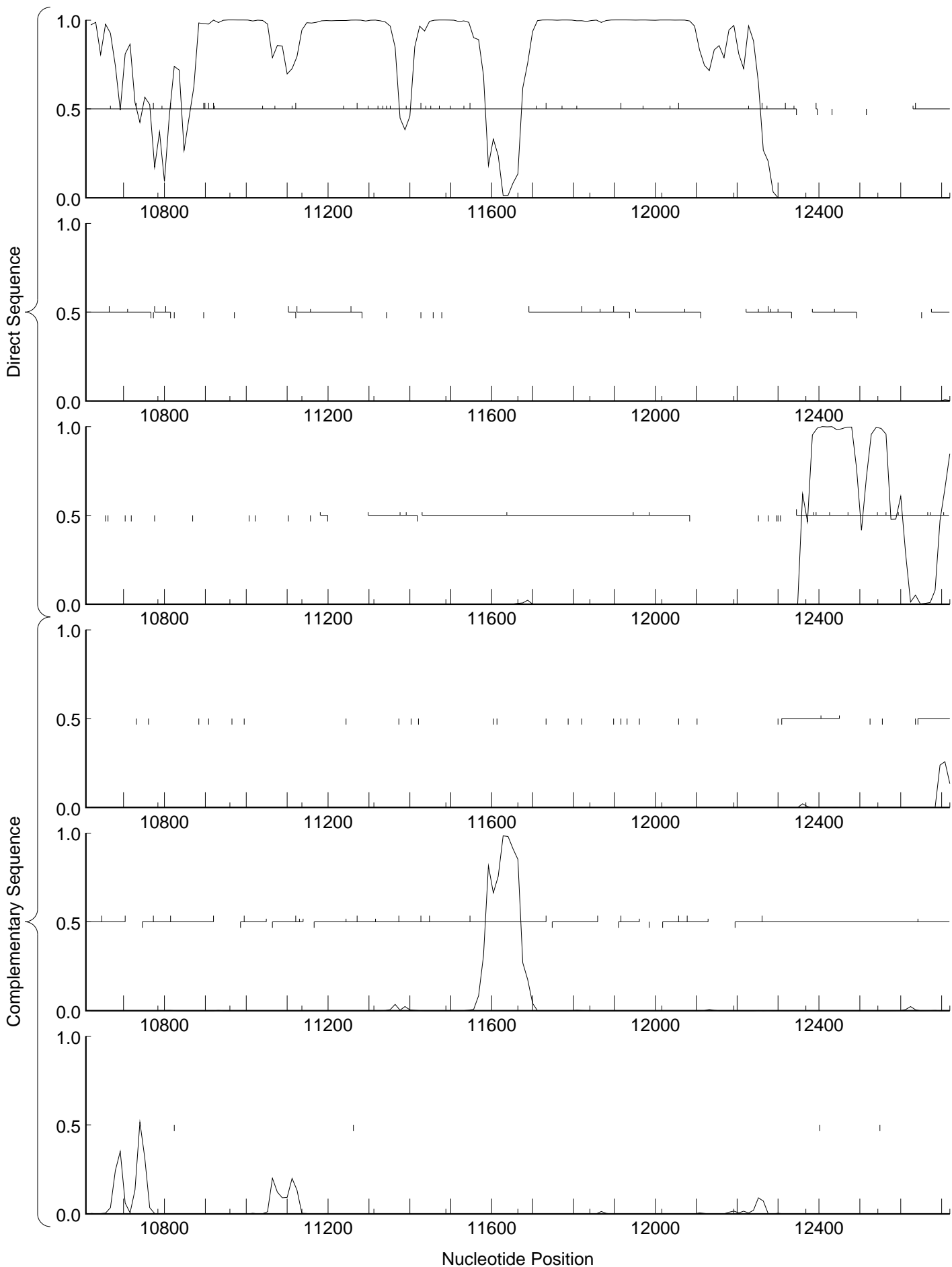
Complete sequence, 38449 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 3/20

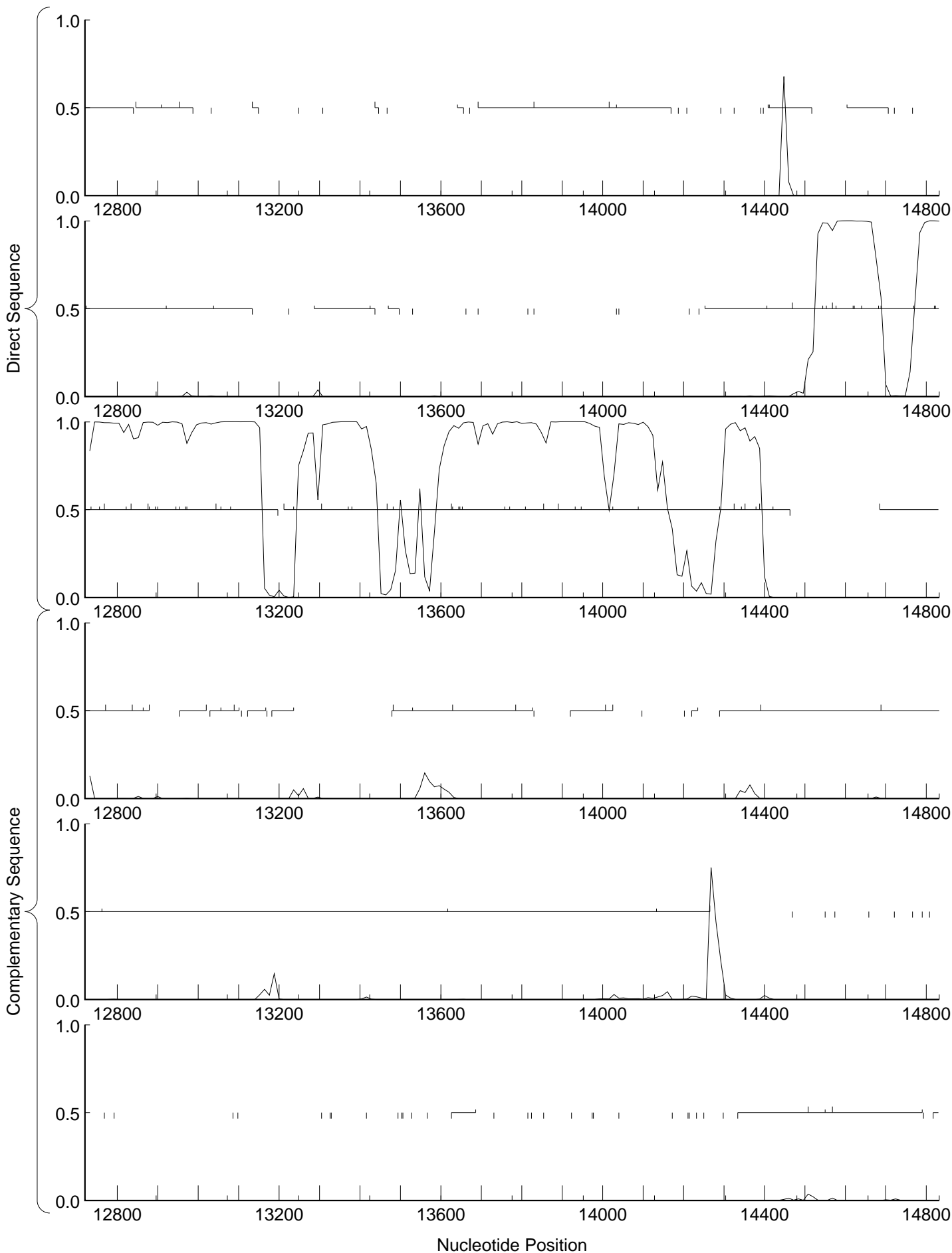


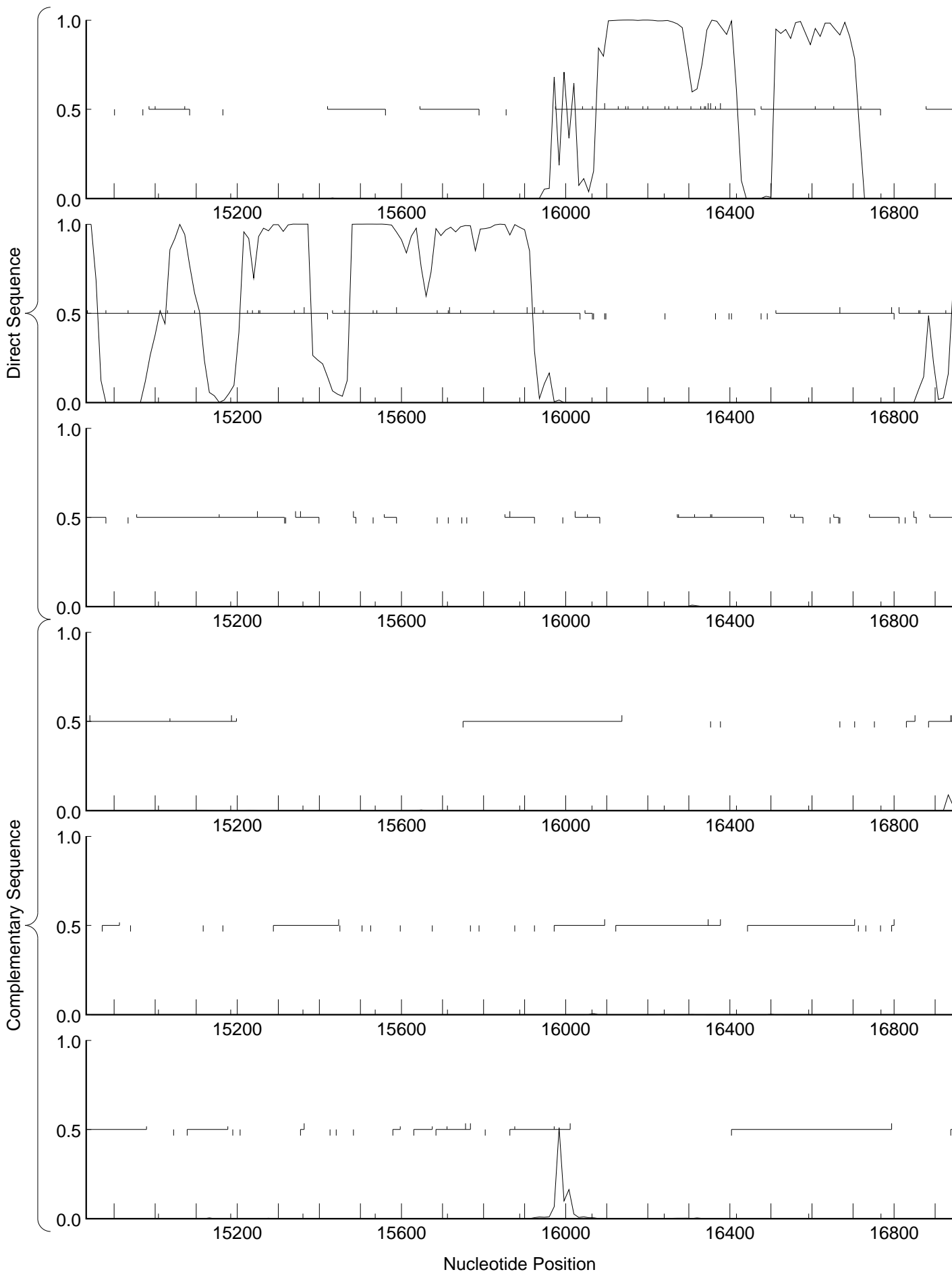


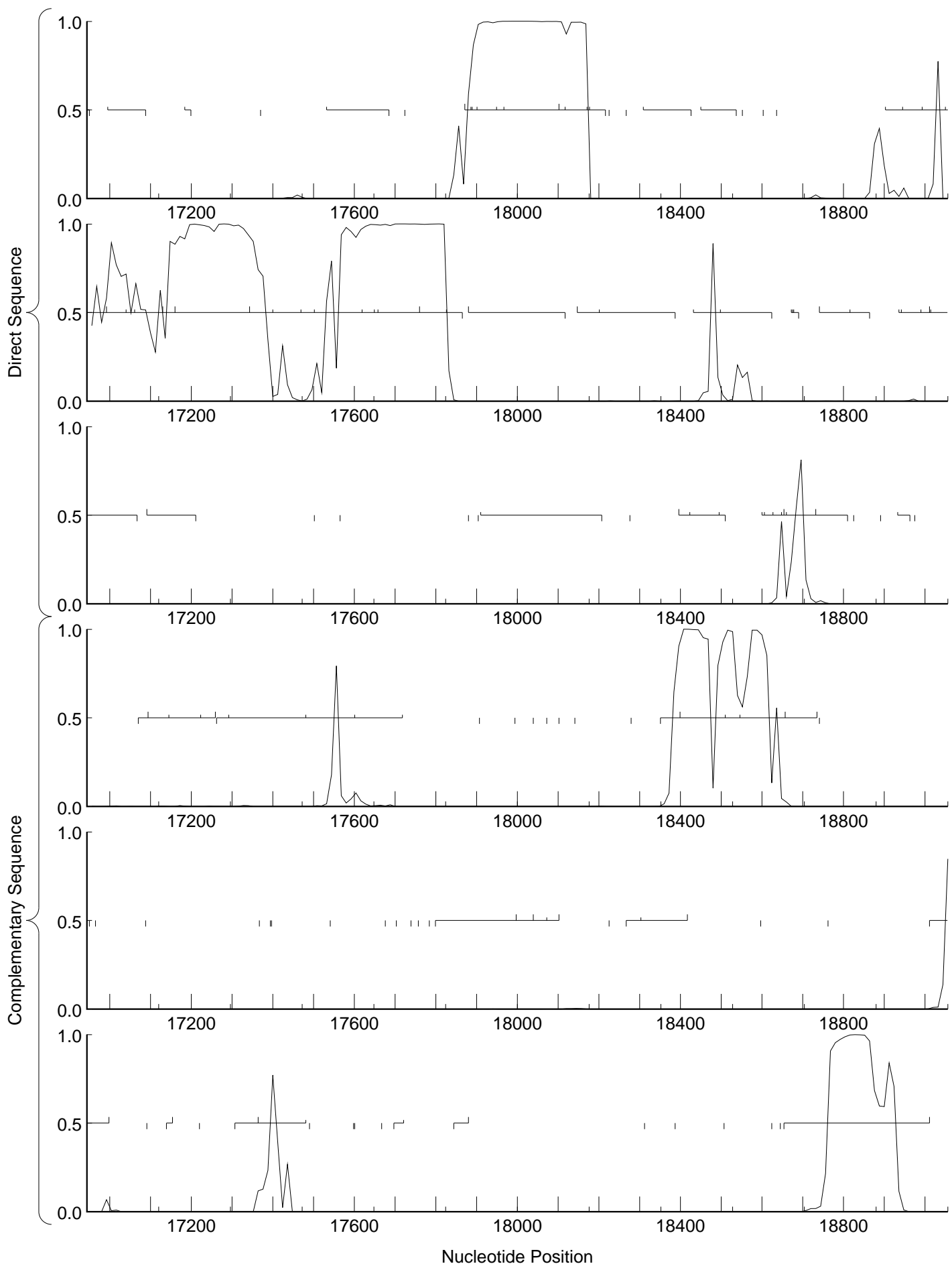


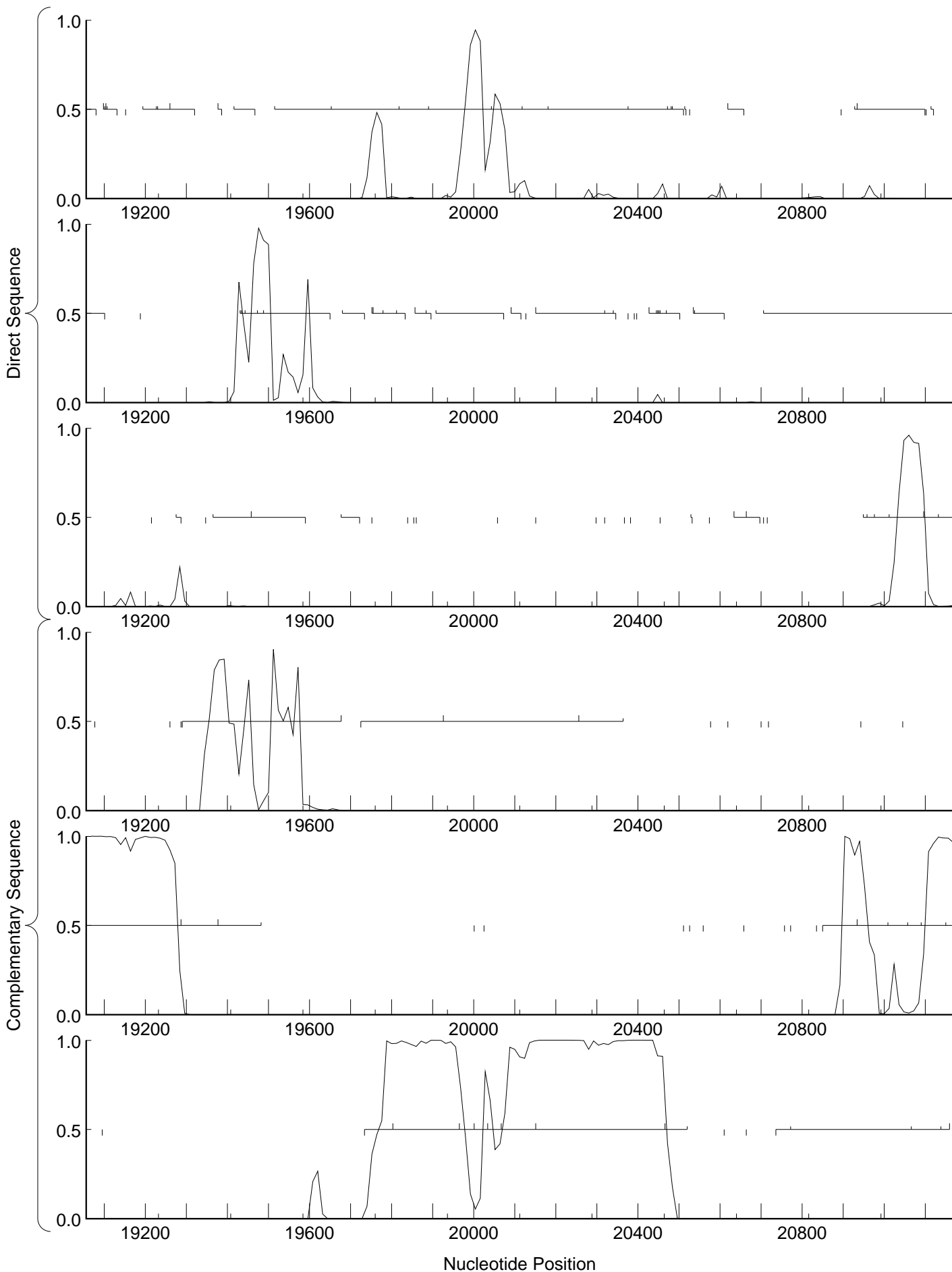


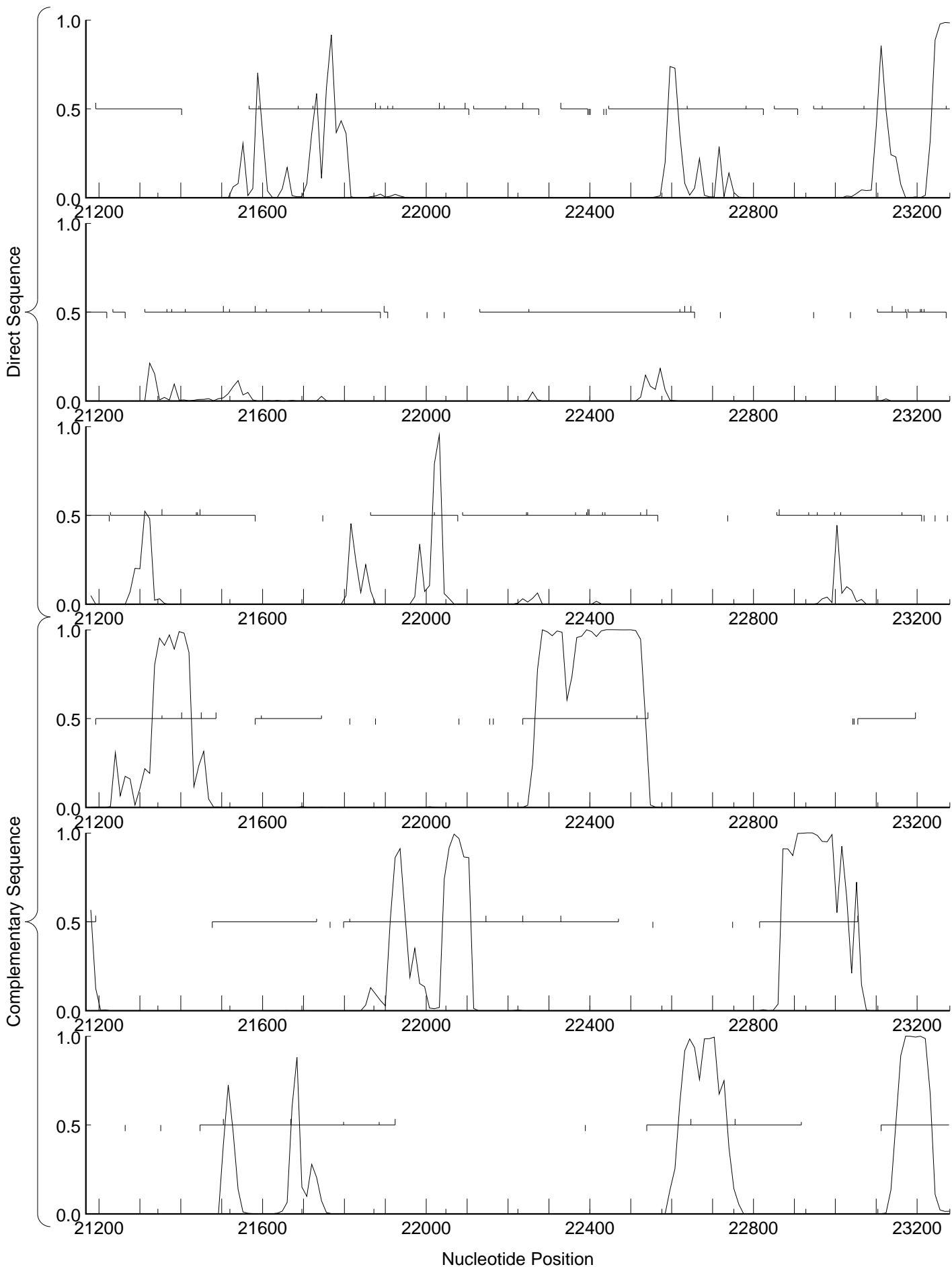




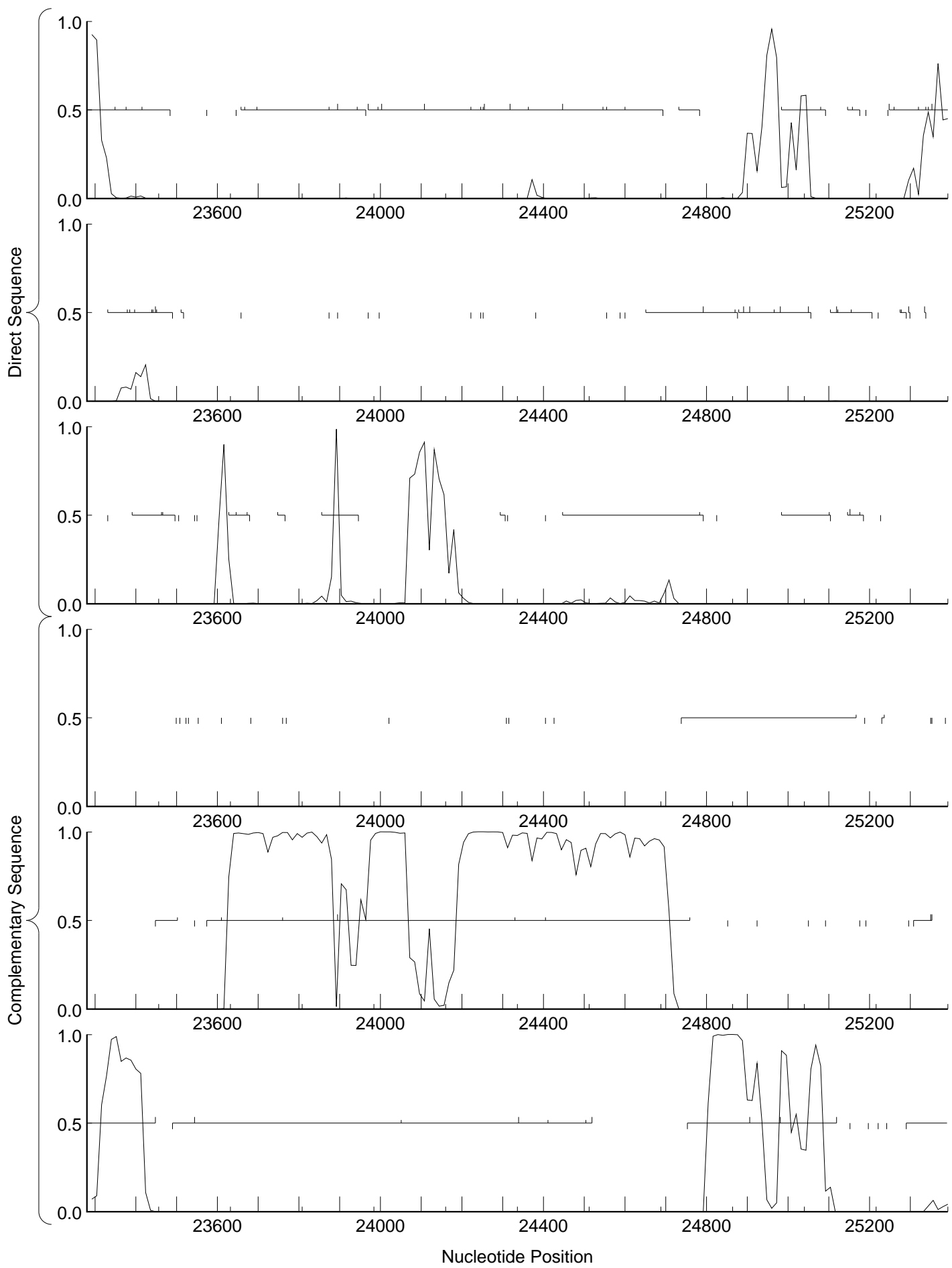




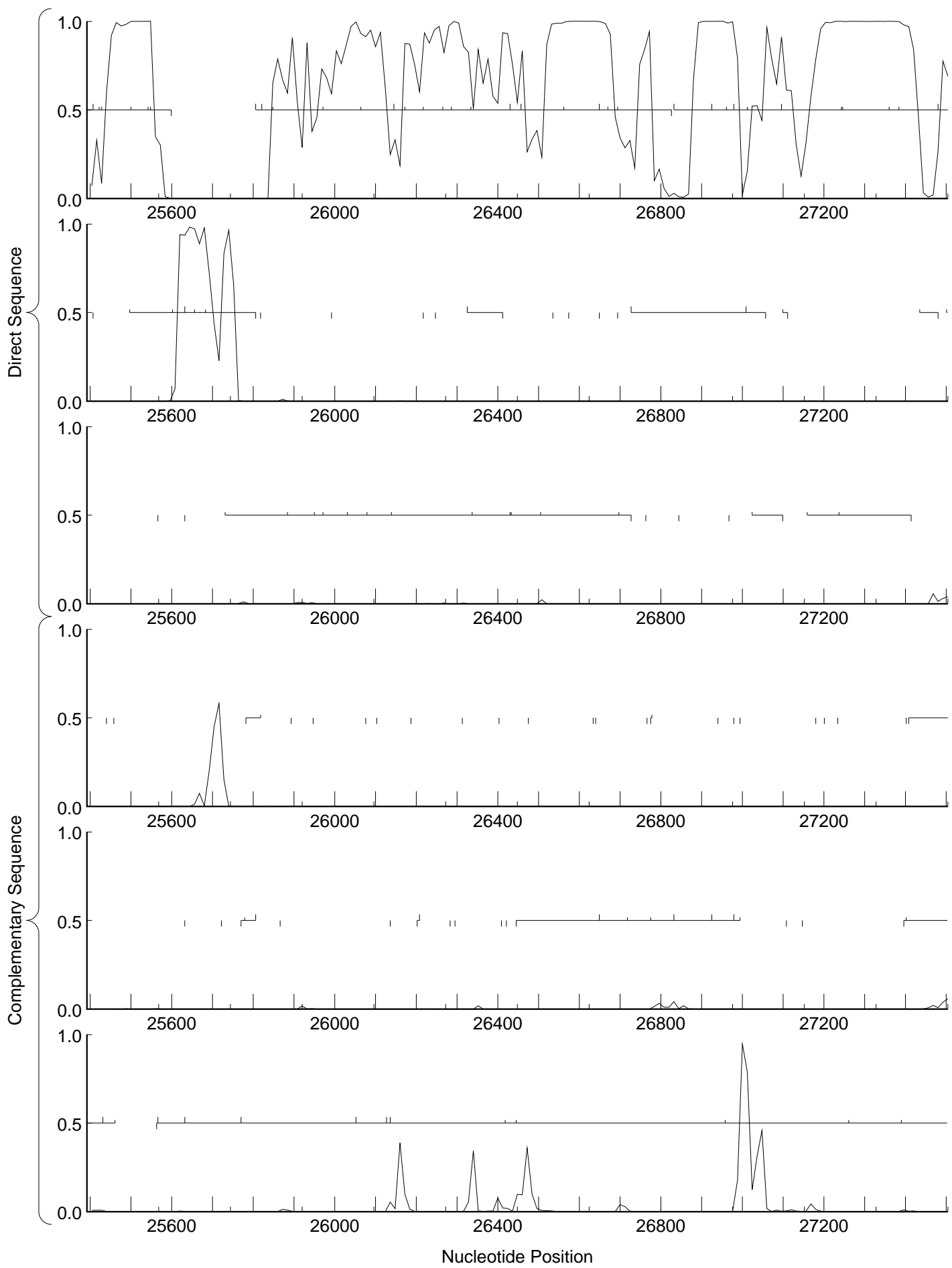




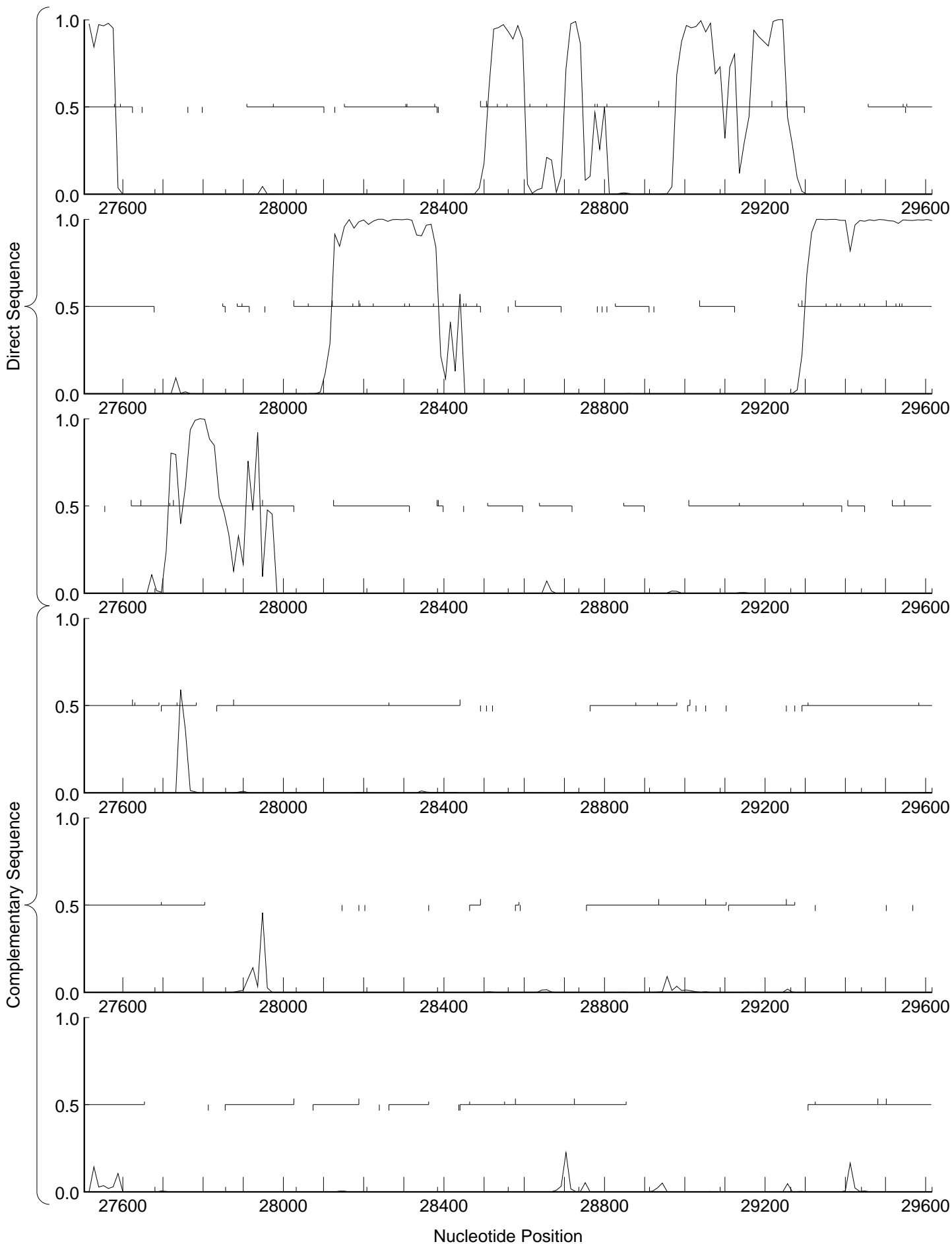
plete sequence, 38449 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 13/20



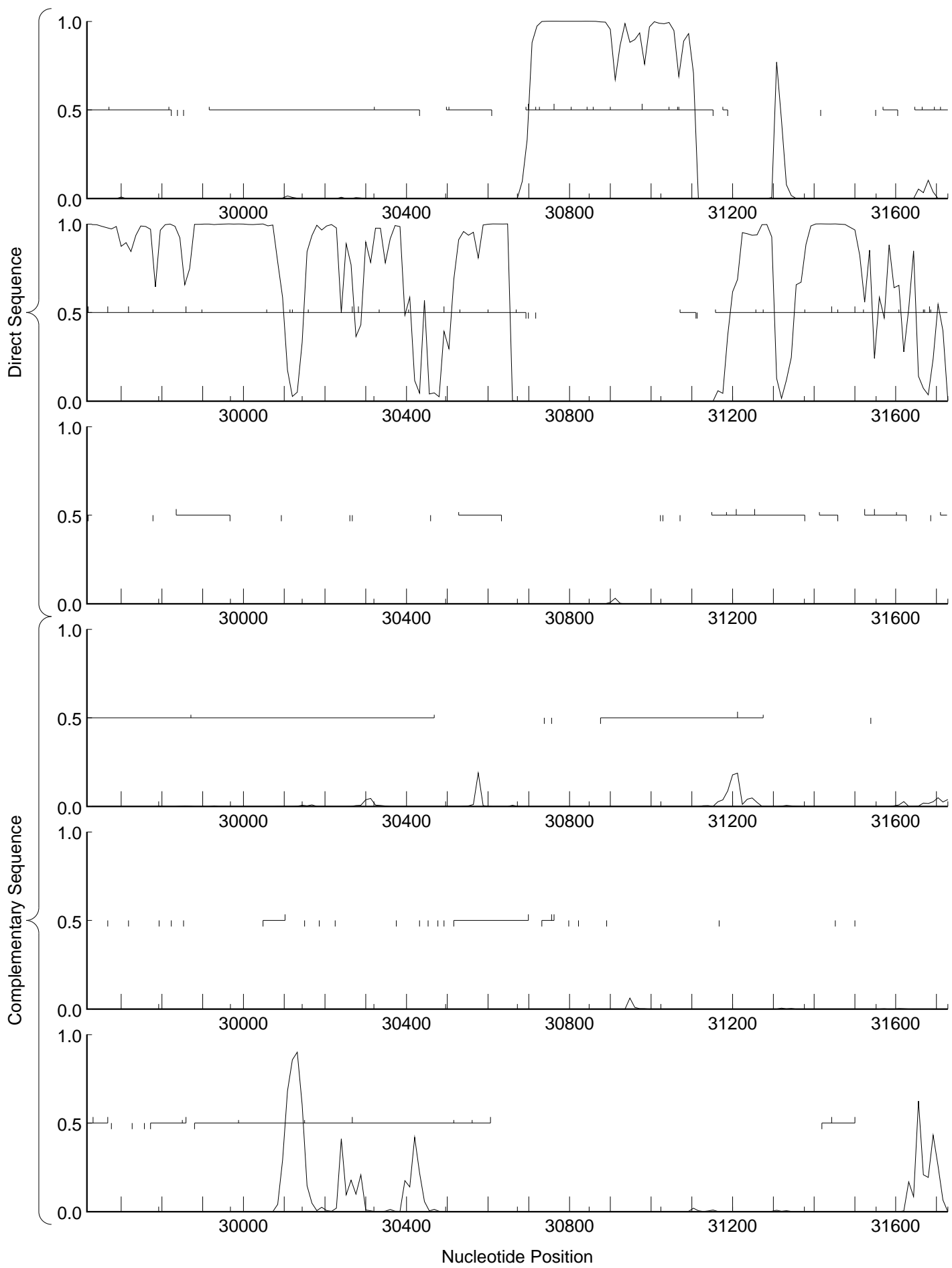
plete sequence, 38449 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 14/20



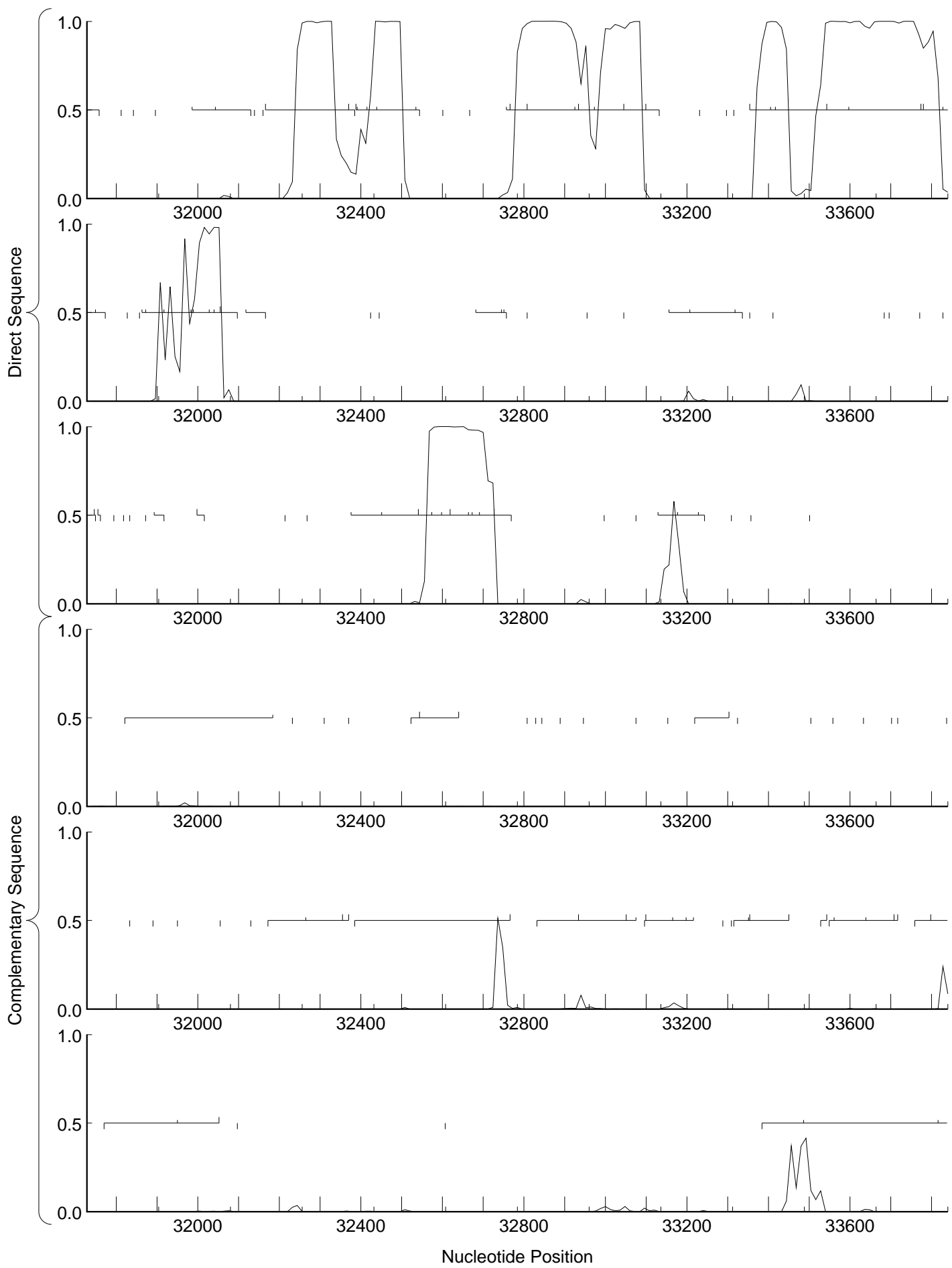
plete sequence, 38449 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 15/20



plete sequence, 38449 bp including 12-base 3' overhang (GAGTTGCCGGCA), Cluster AS2, Order 4, Window 96, Step 12, 16/20



plete sequence, 38449 bp including 12-base 3' overhang (GAGTTGCCGCA), Cluster AS2, Order 4, Window 96, Step 12, 17/20



plete sequence, 38449 bp including 12-base 3' overhang (GAGTTGCCGCA), Cluster AS2, Order 4, Window 96, Step 12, 18/20

