

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Moosehead complete sequence, 42671 bp including 12-base 3' overhang (TCGTGG
Analysis Date : 4/6/20 at 14:15:23
Pages : 22
Sequence Length : 42671 bp
GC Content : 65.07%

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,

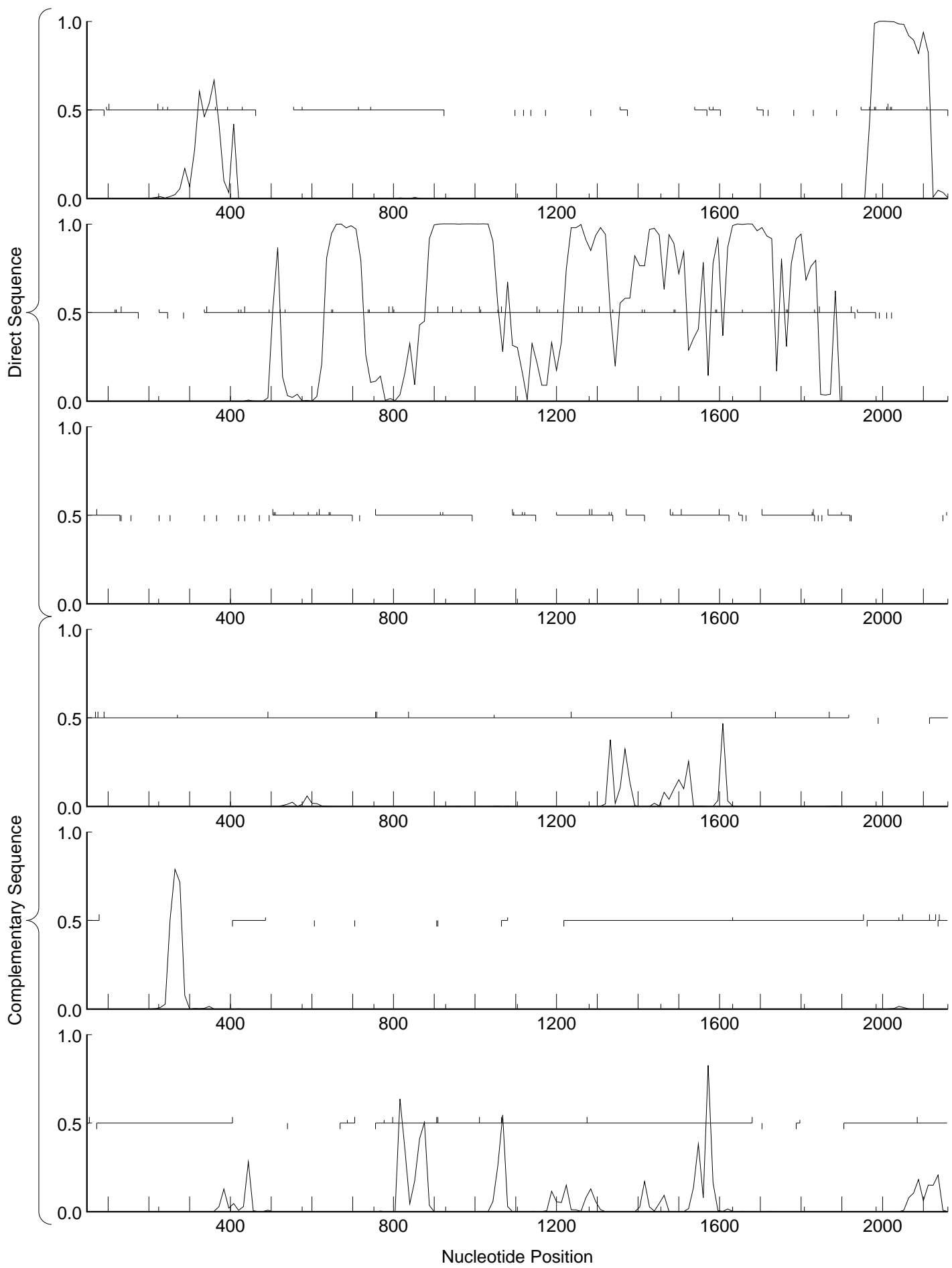
MATRIX INFORMATION

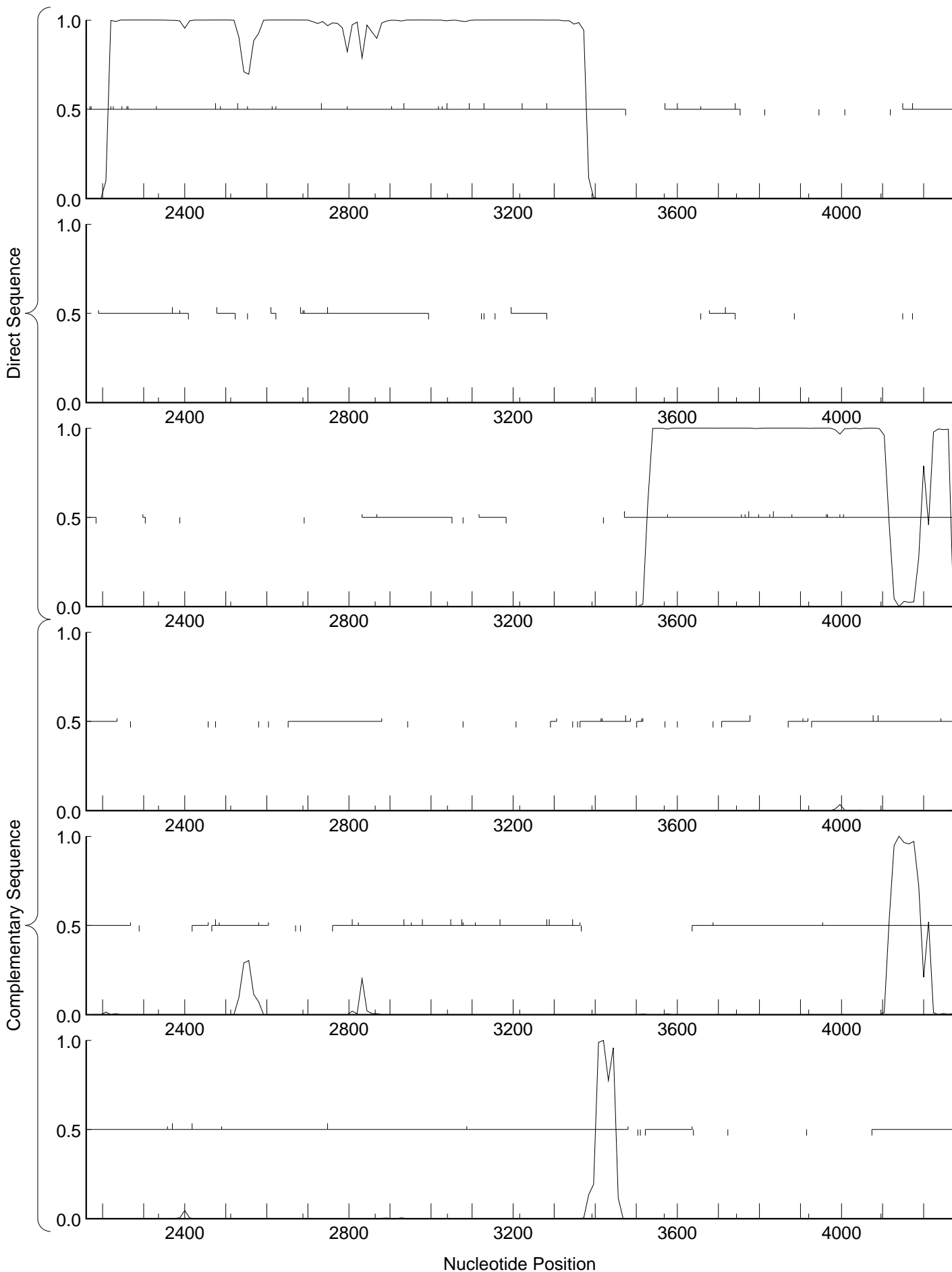
Matrix : -
Author : -
Order : 4

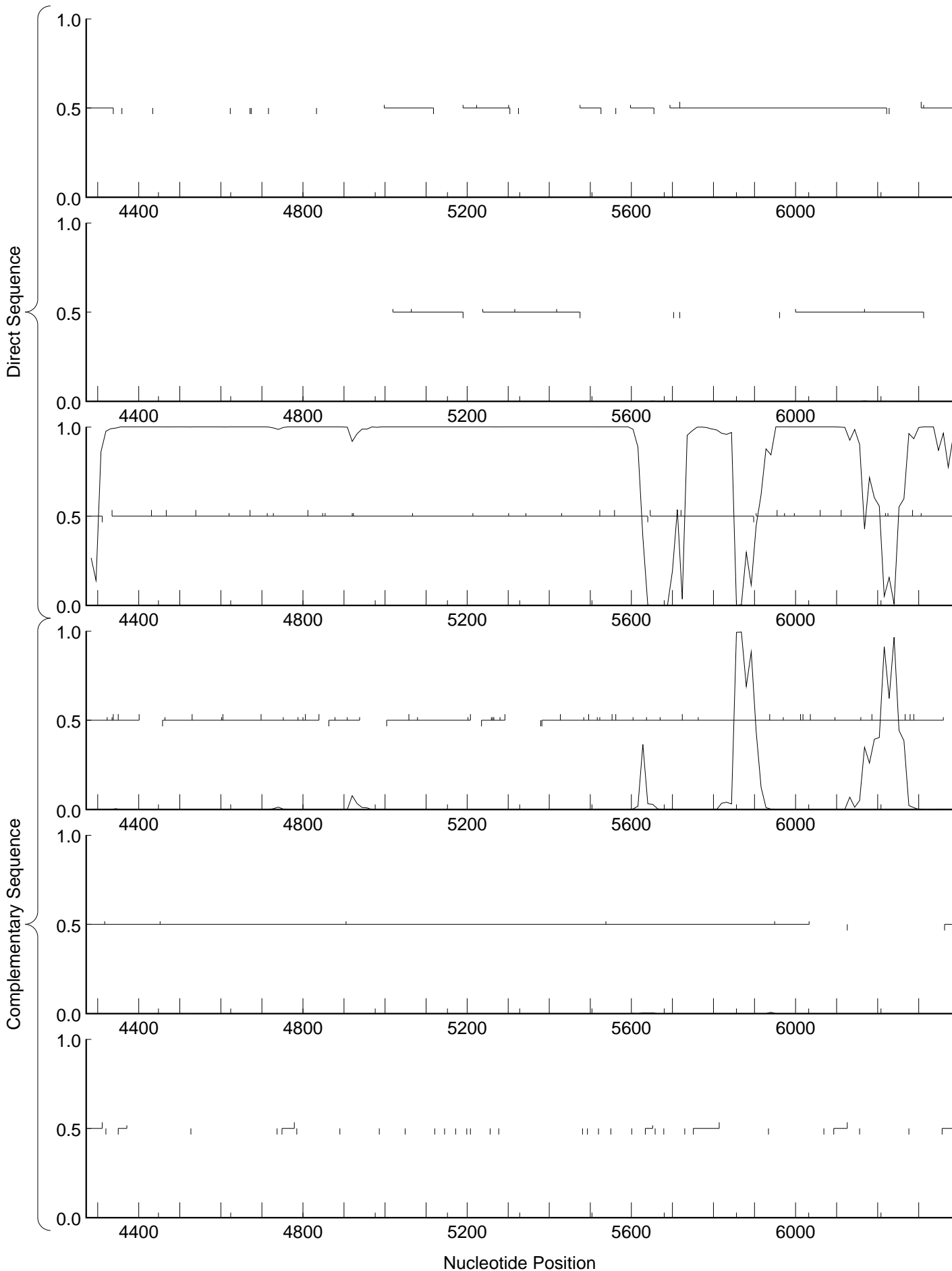
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
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Atlanta, GA 30332-0230

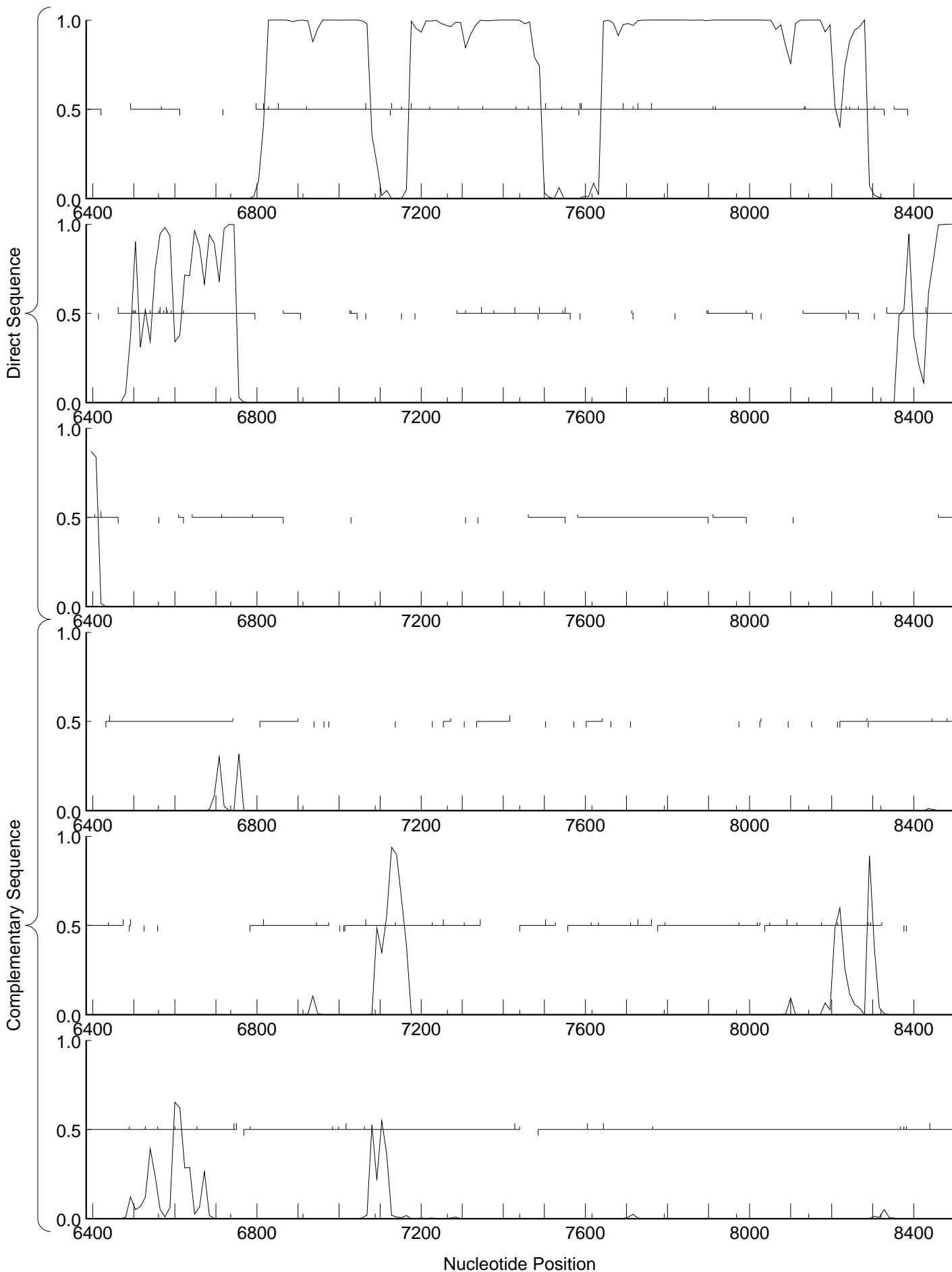
Matrix notes & comments

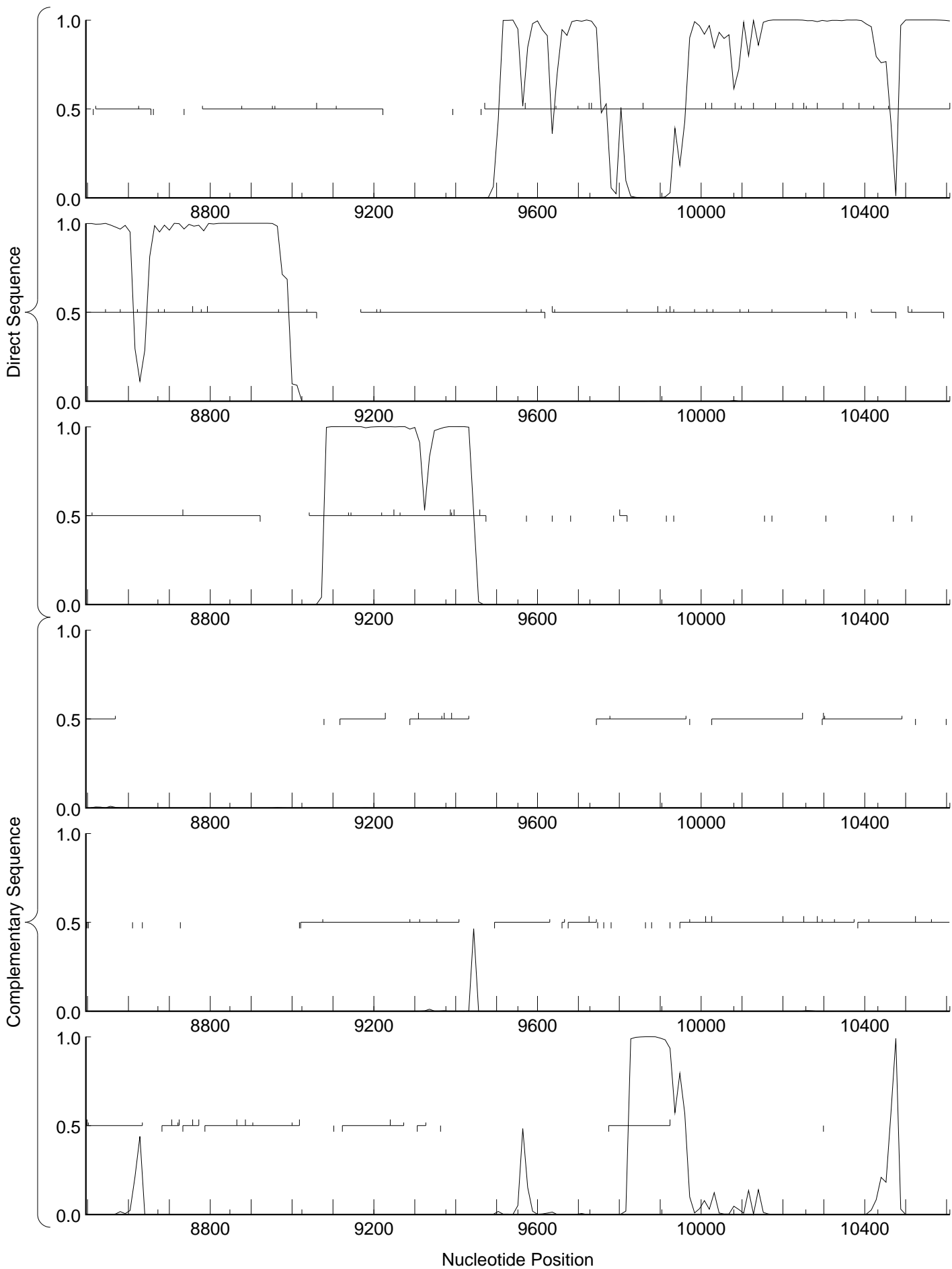
Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:36:14 2018

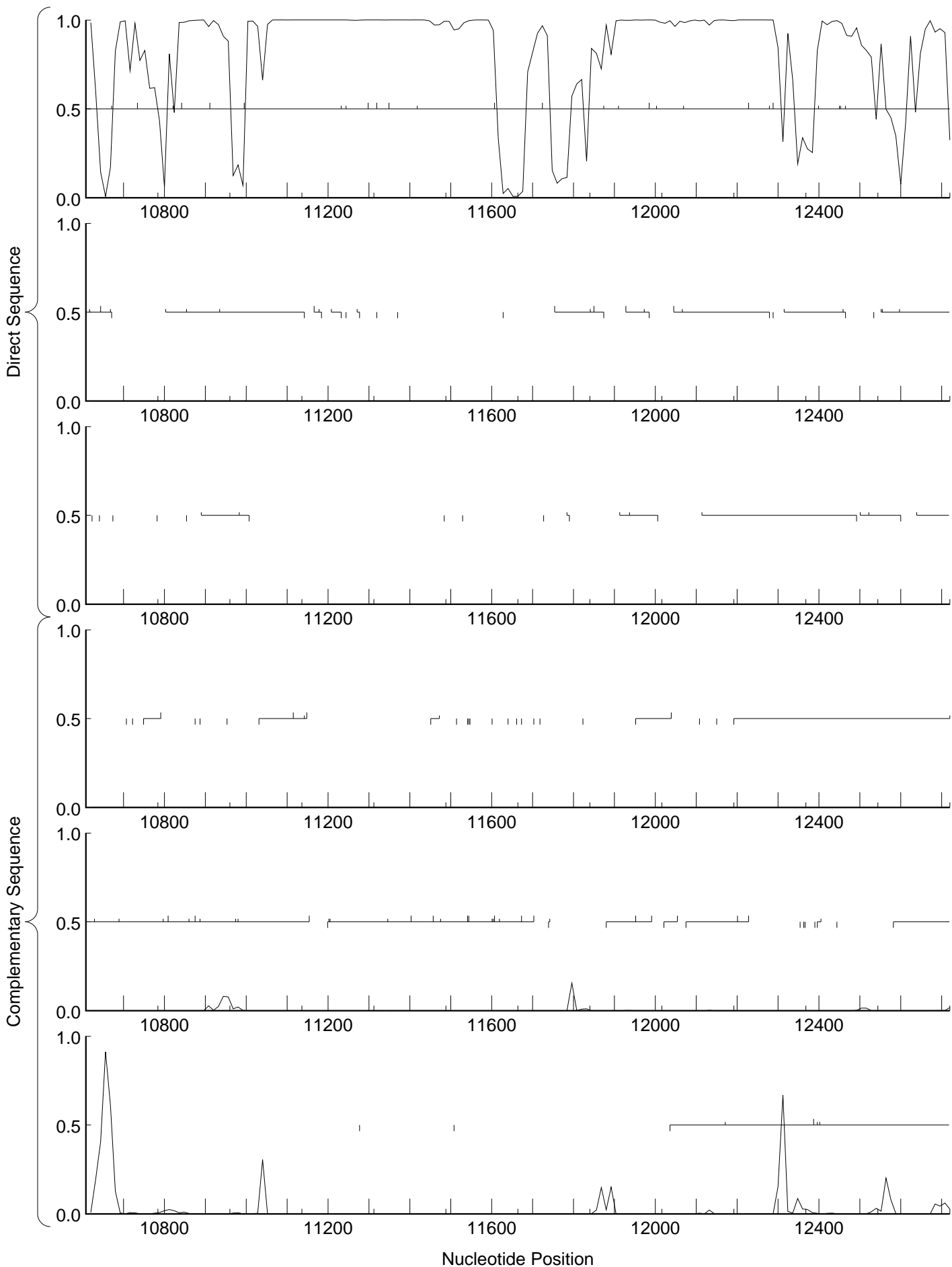


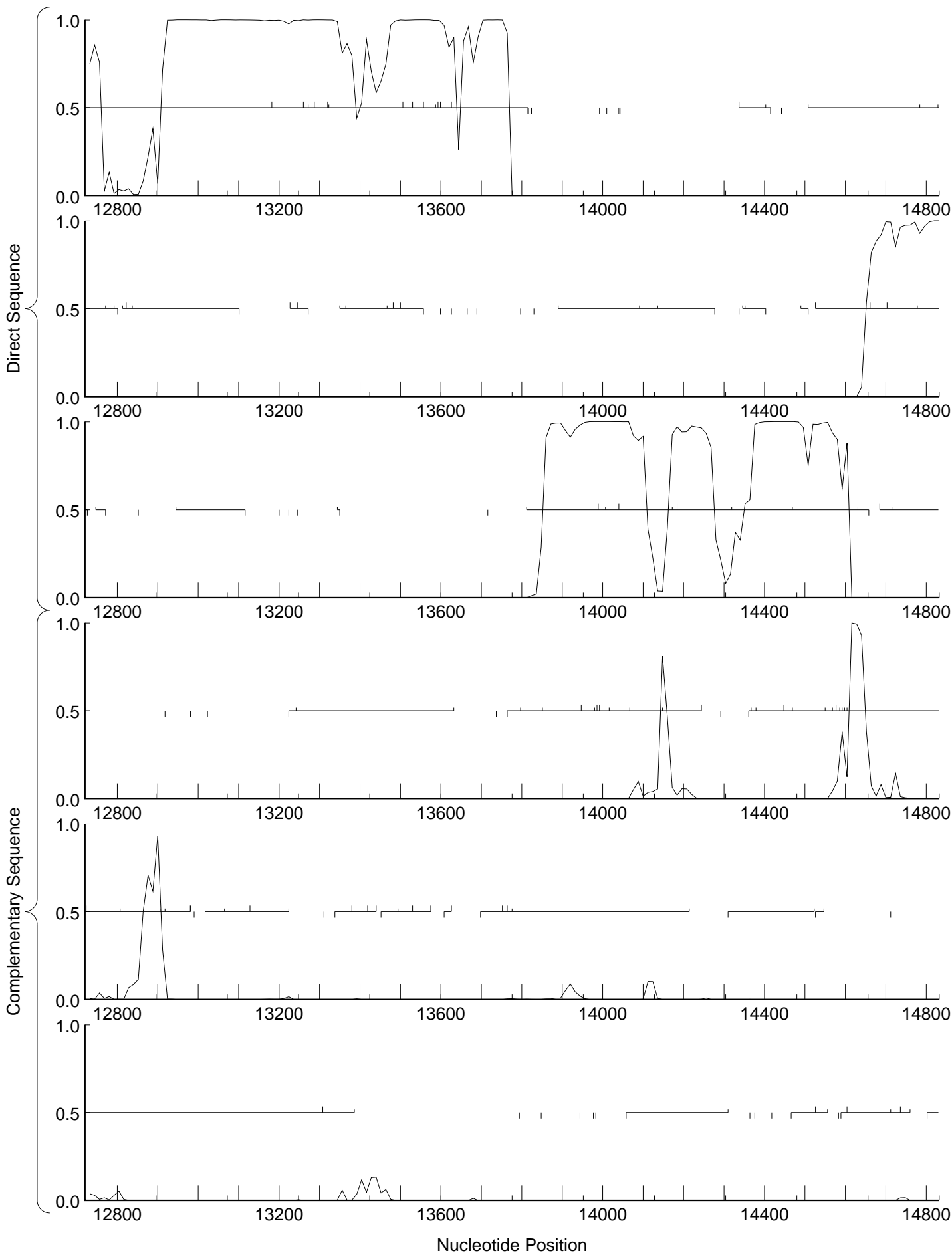


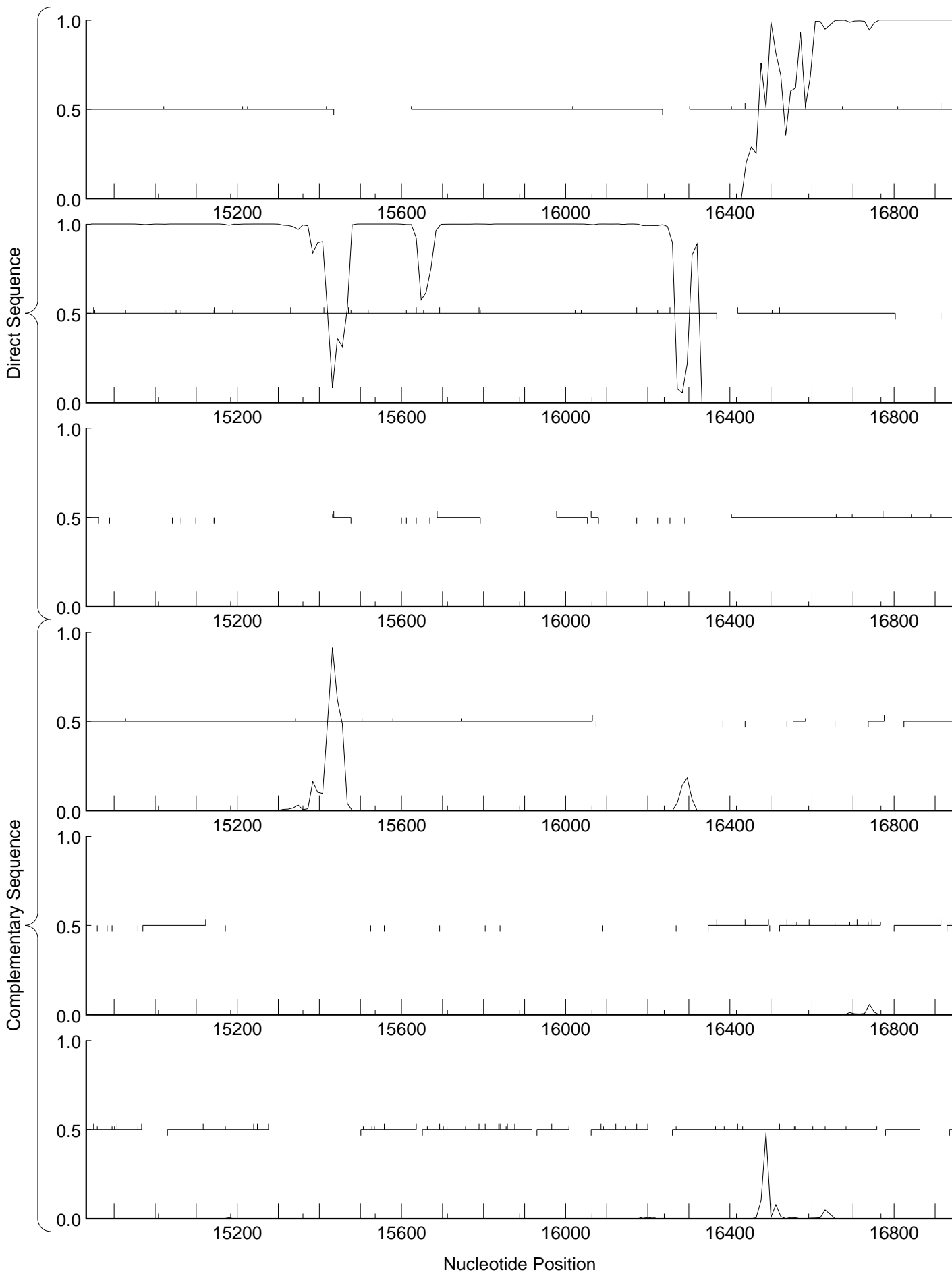


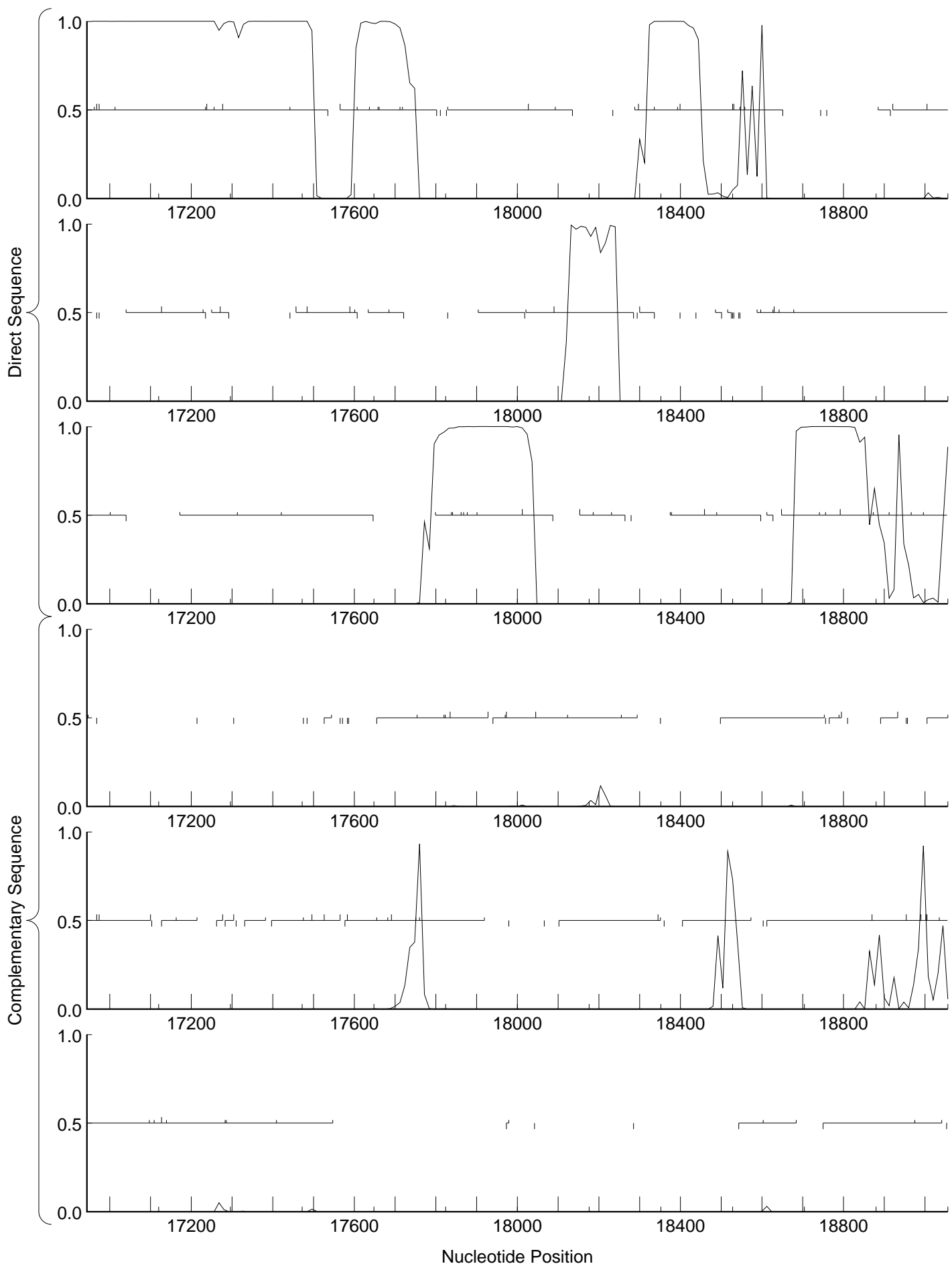


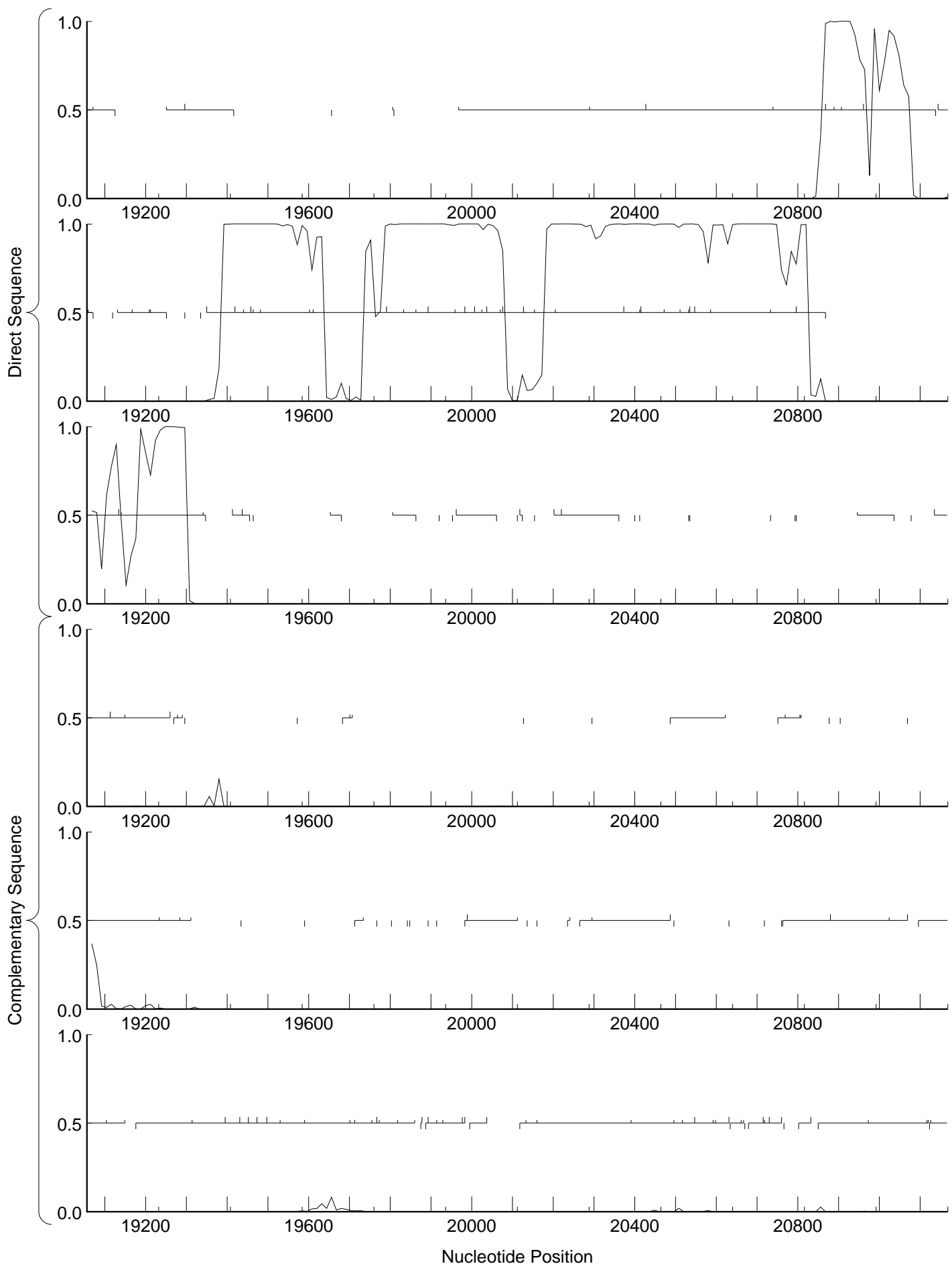


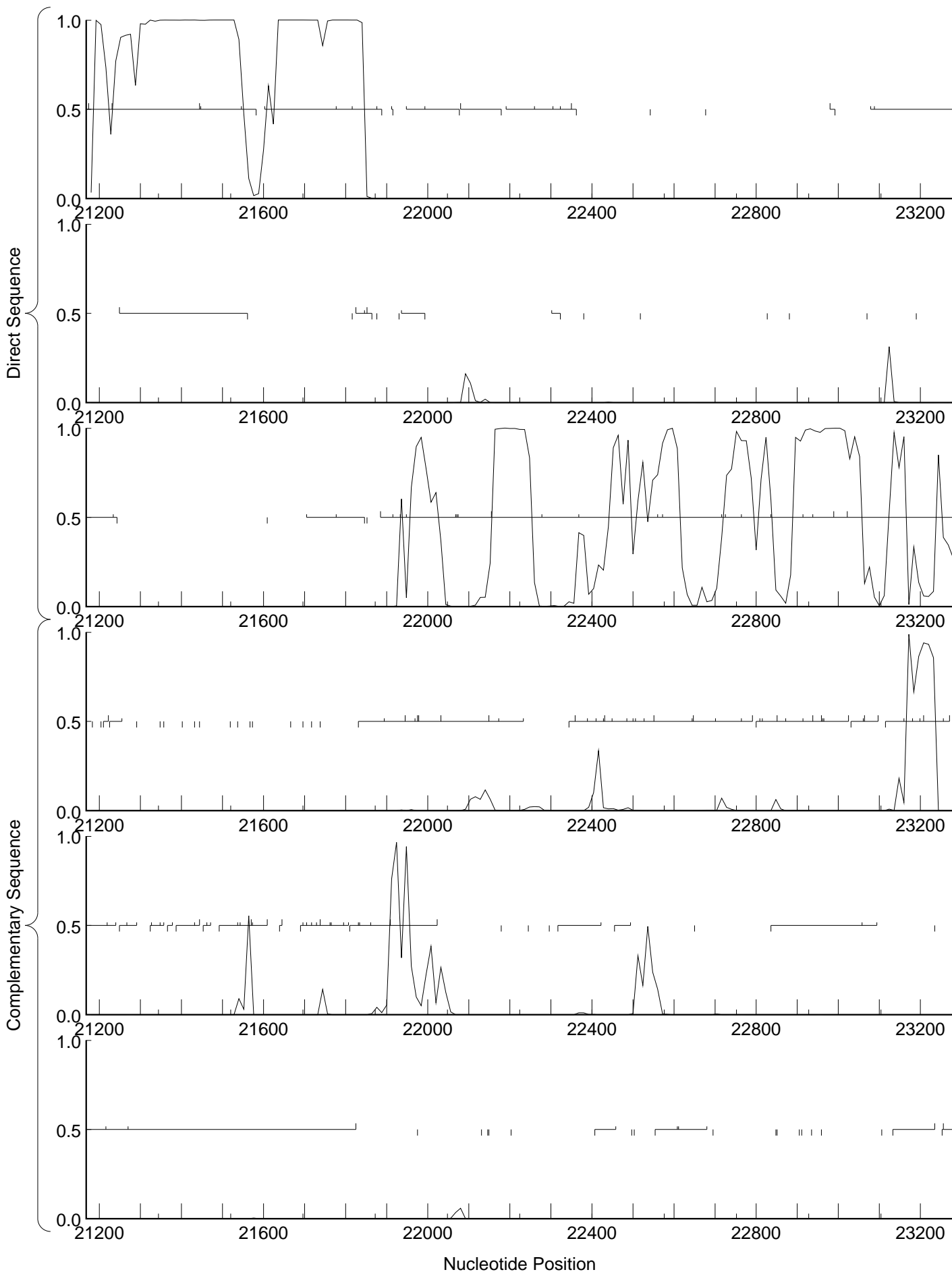




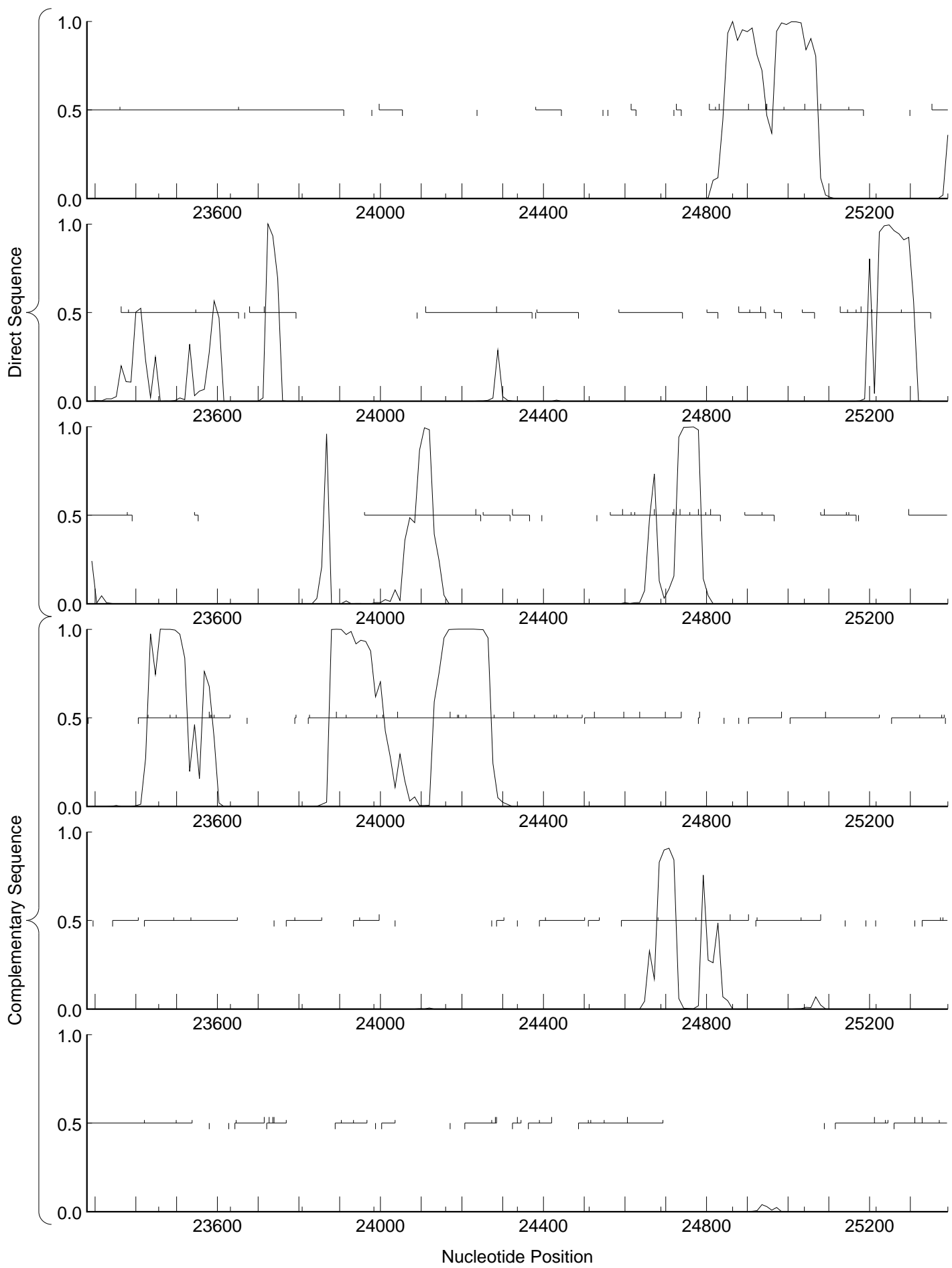




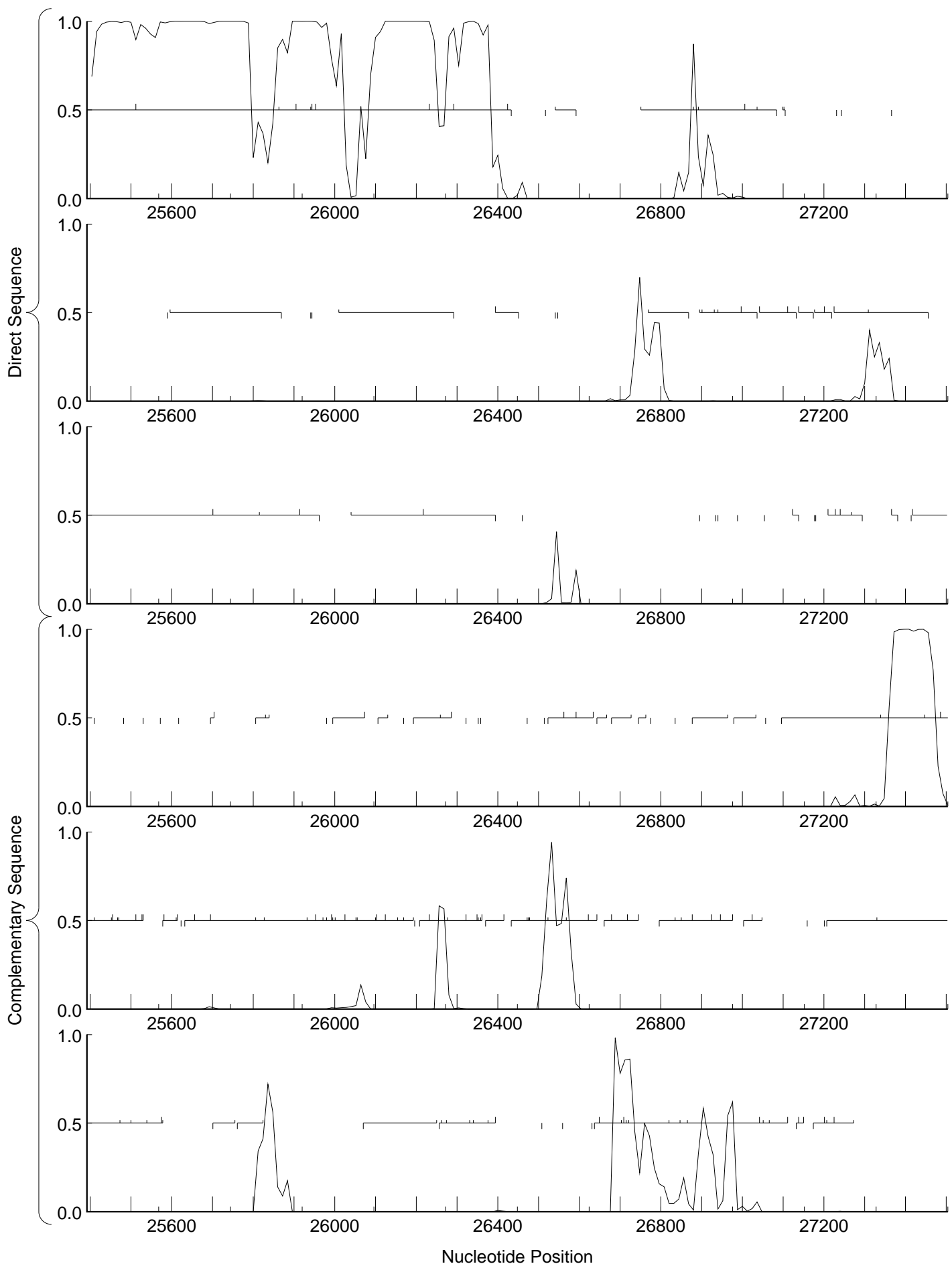




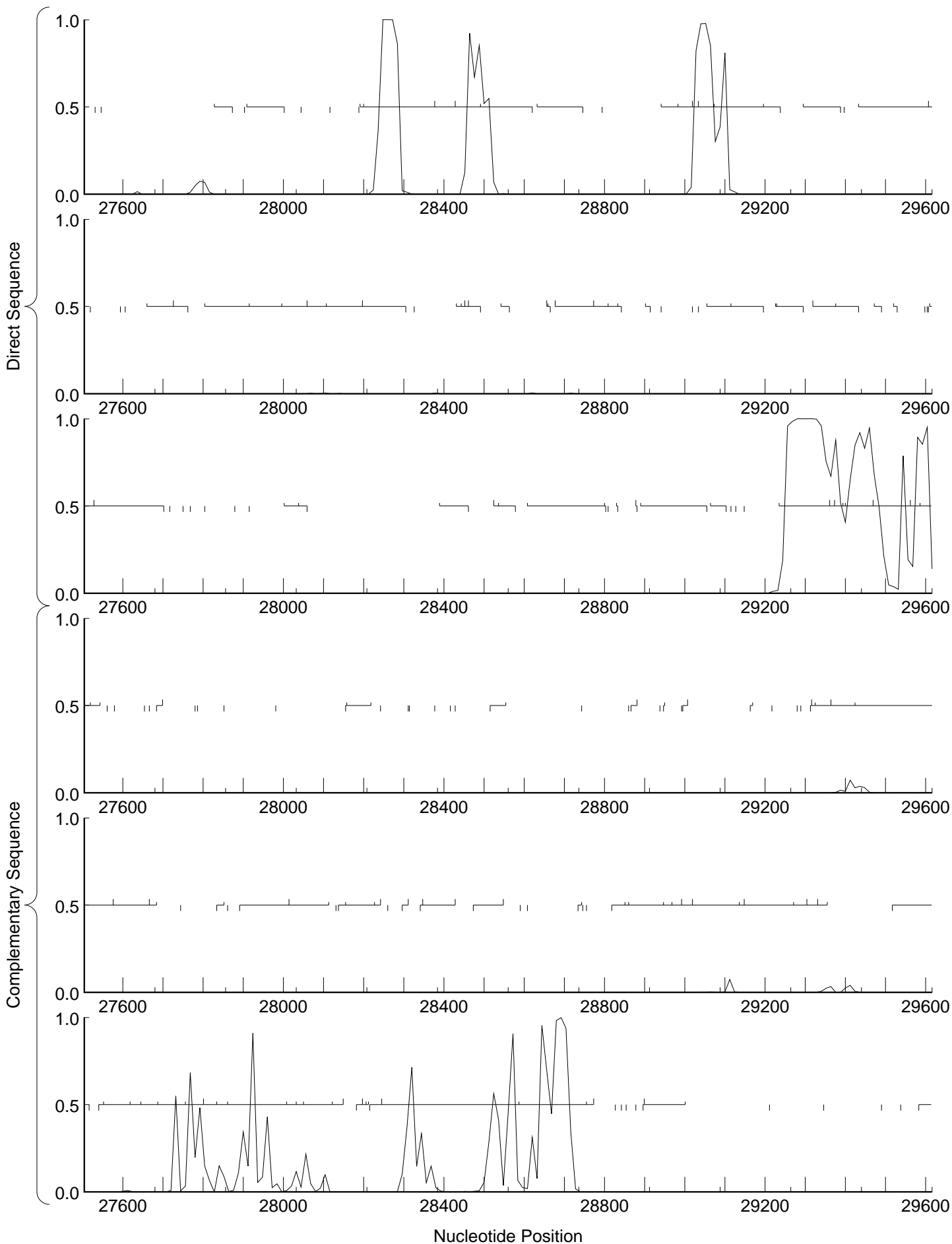
plete sequence, 42671 bp including 12-base 3' overhang (TCGTCGGGGTGA), Cluster CZ6, Order 4, Window 96, Step 12, 13/22

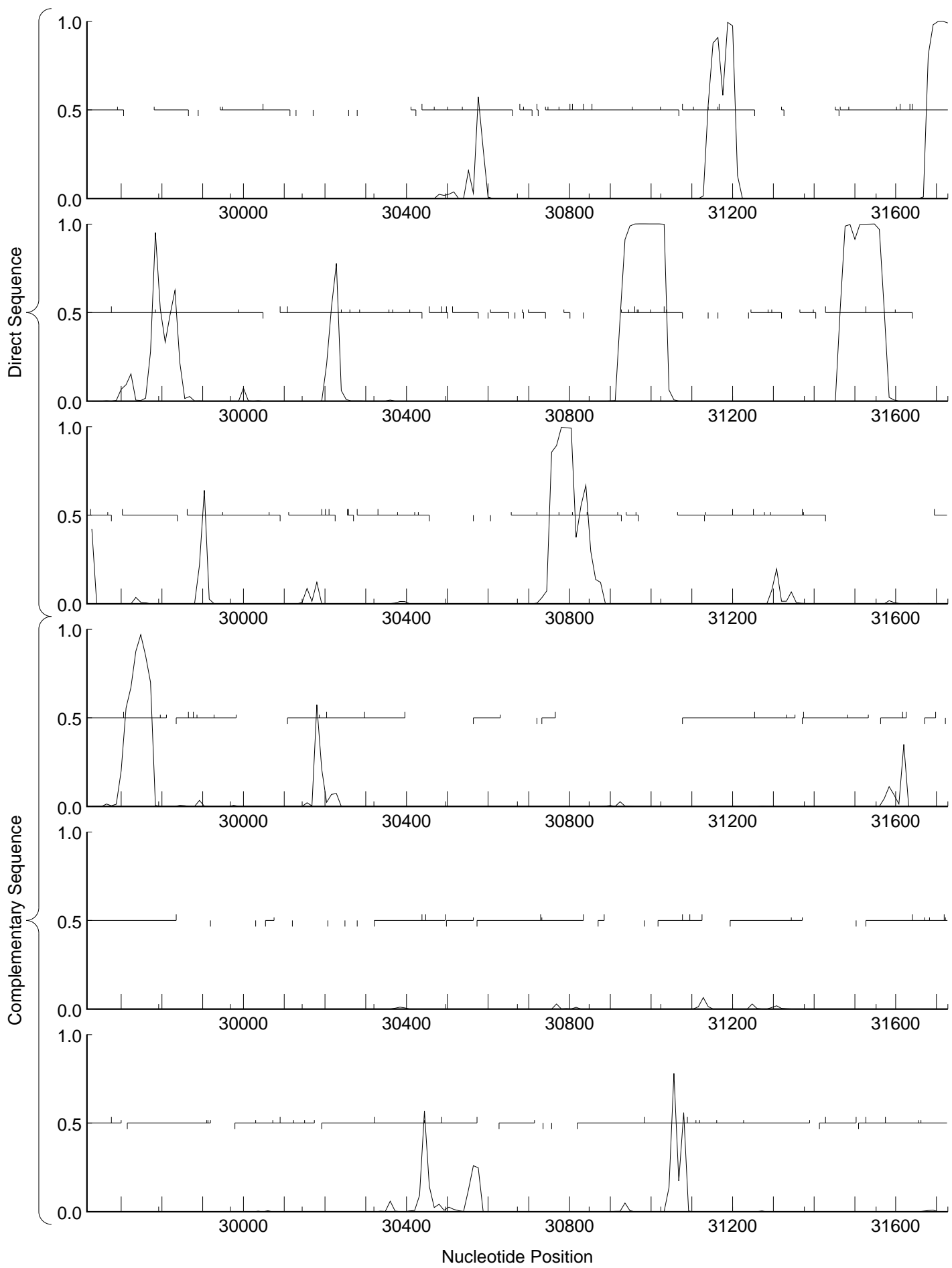


plete sequence, 42671 bp including 12-base 3' overhang (TCGTCTGGGGTGA), Cluster CZ6, Order 4, Window 96, Step 12, 14/22



plete sequence, 42671 bp including 12-base 3' overhang (TCGTCGGGGTGA), Cluster CZ6, Order 4, Window 96, Step 12, 15/22





plete sequence, 42671 bp including 12-base 3' overhang (TCGTCGGGGTGA), Cluster CZ6, Order 4, Window 96, Step 12, 17/22

