

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Grizzly complete sequence, 41752 bp including 11-base 3' overhang (CCCC
Analysis Date : 6/18/24 at 23:11:34
Pages : 21
Sequence Length : 41752 bp
GC Content : 66.76%

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,

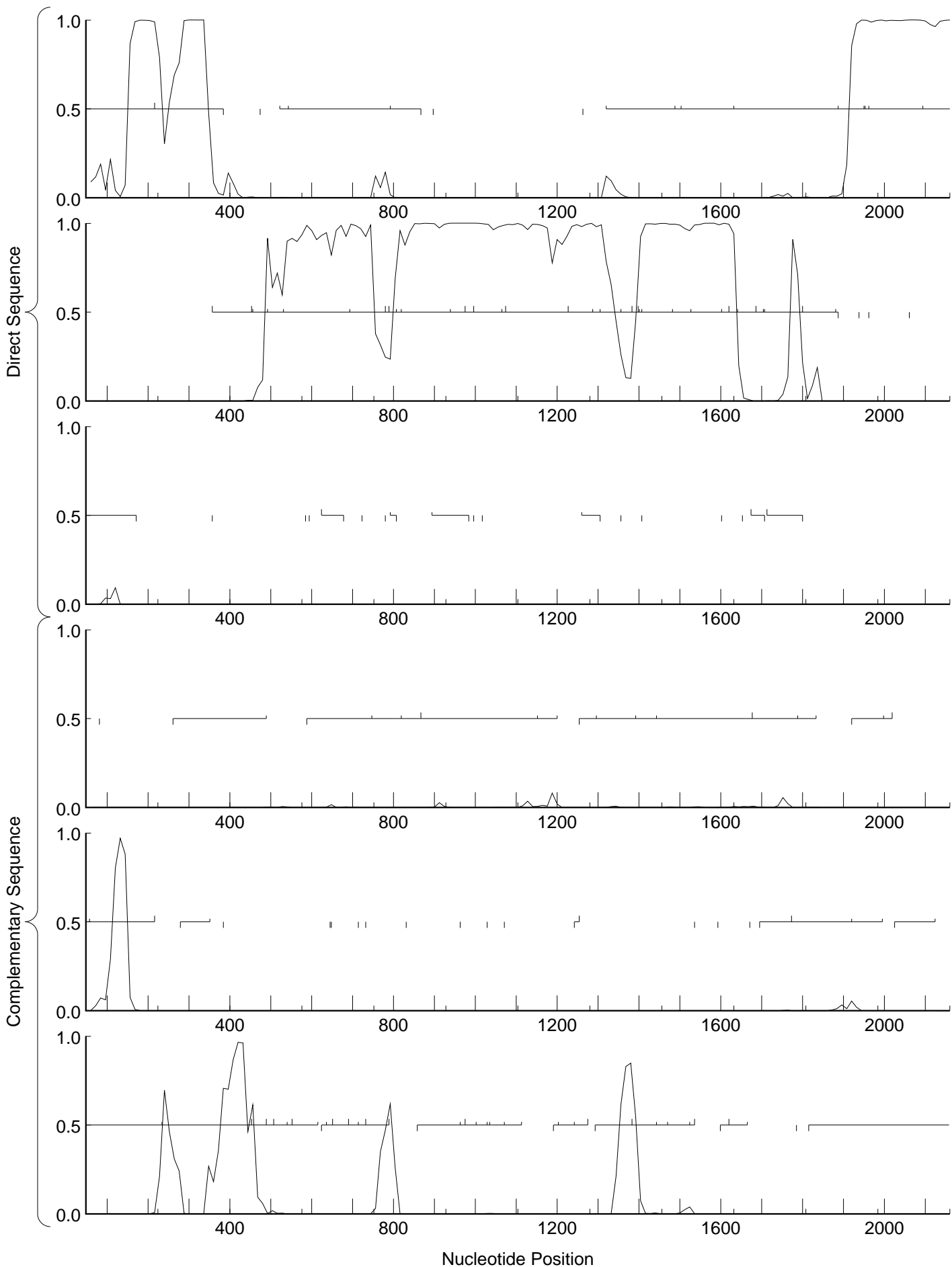
MATRIX INFORMATION

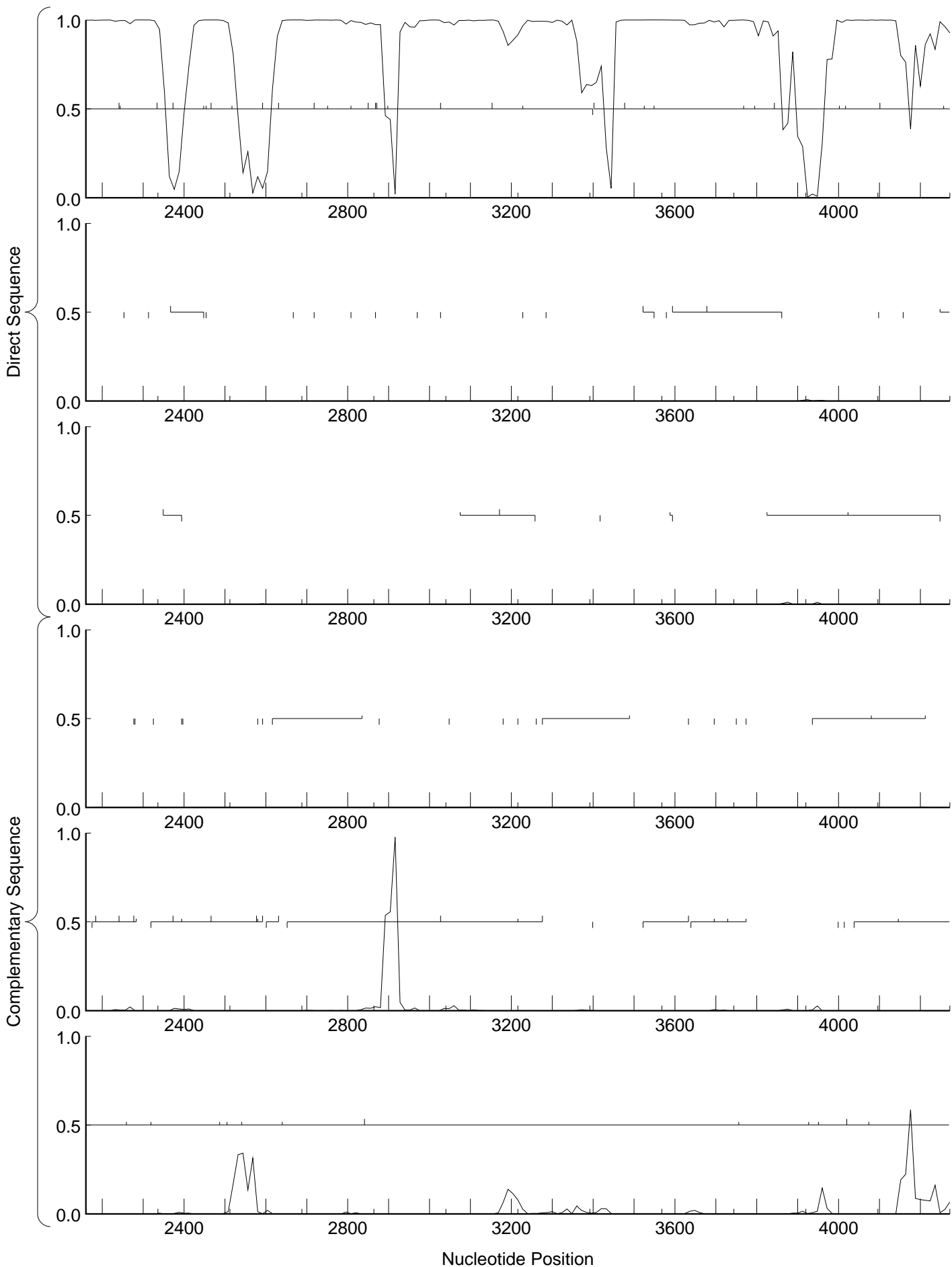
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

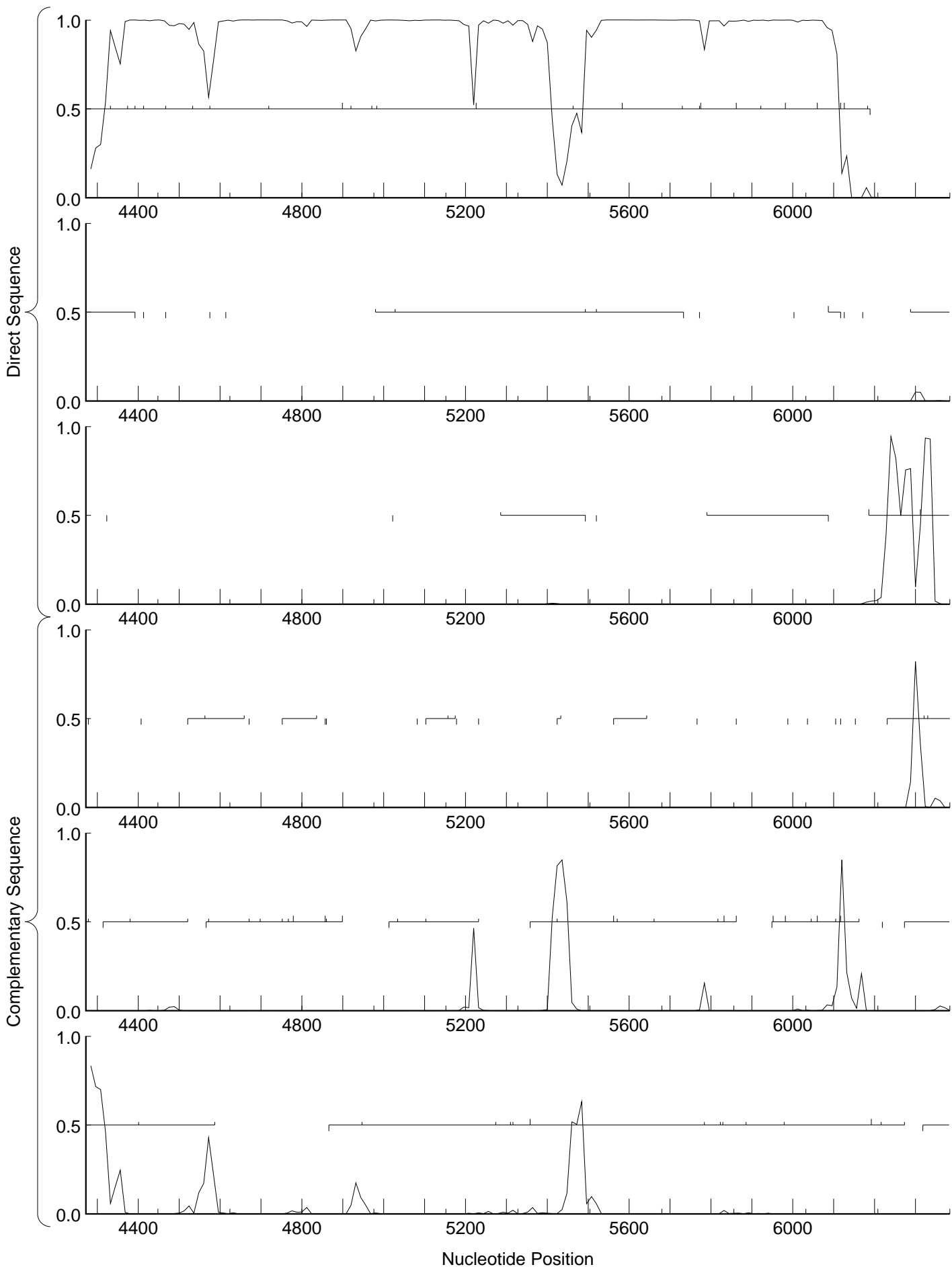
Send questions / comments to:
Dr. M. Borodovsky
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School of Biology
Atlanta, GA 30332-0230

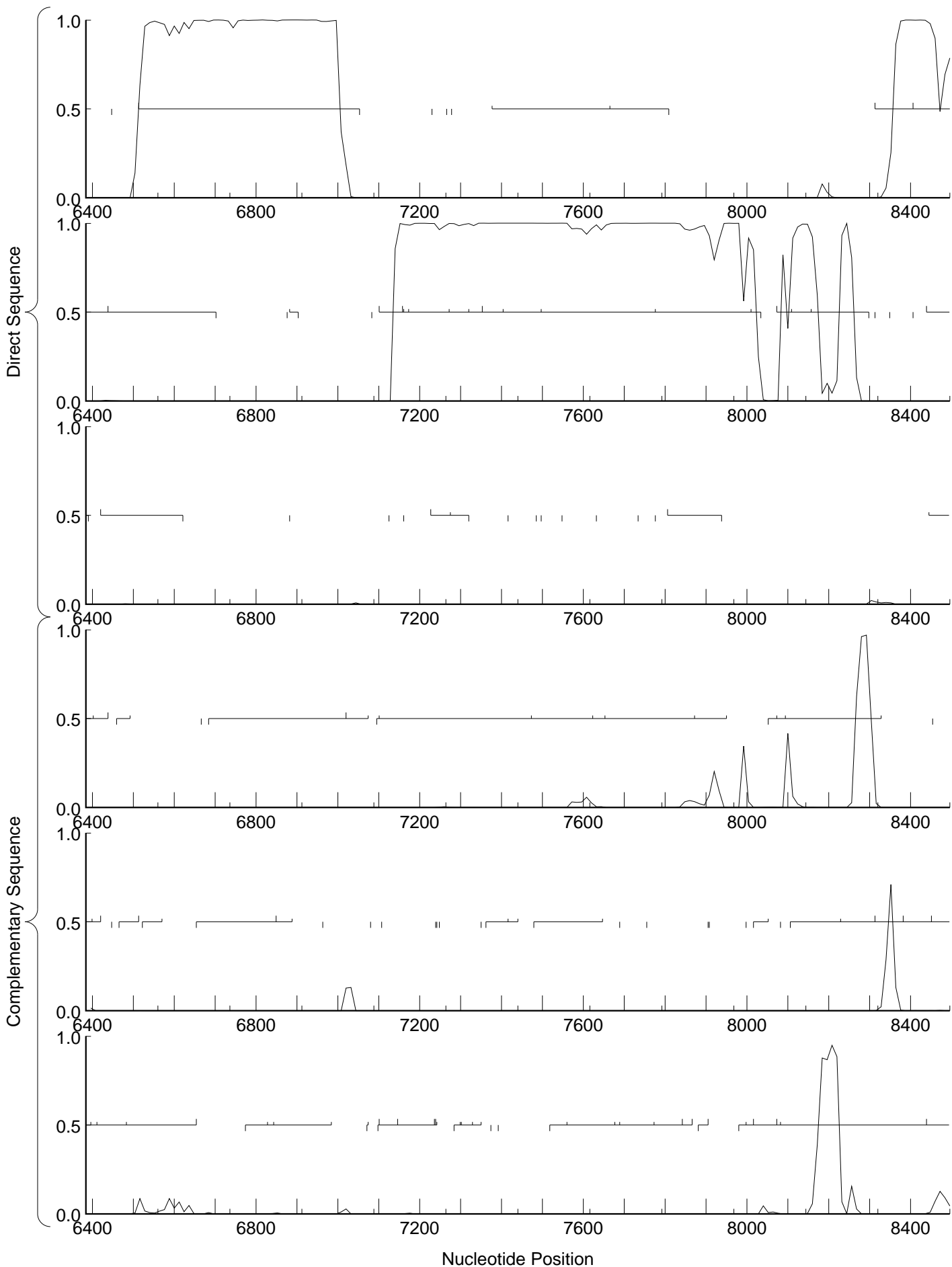
Matrix notes & comments

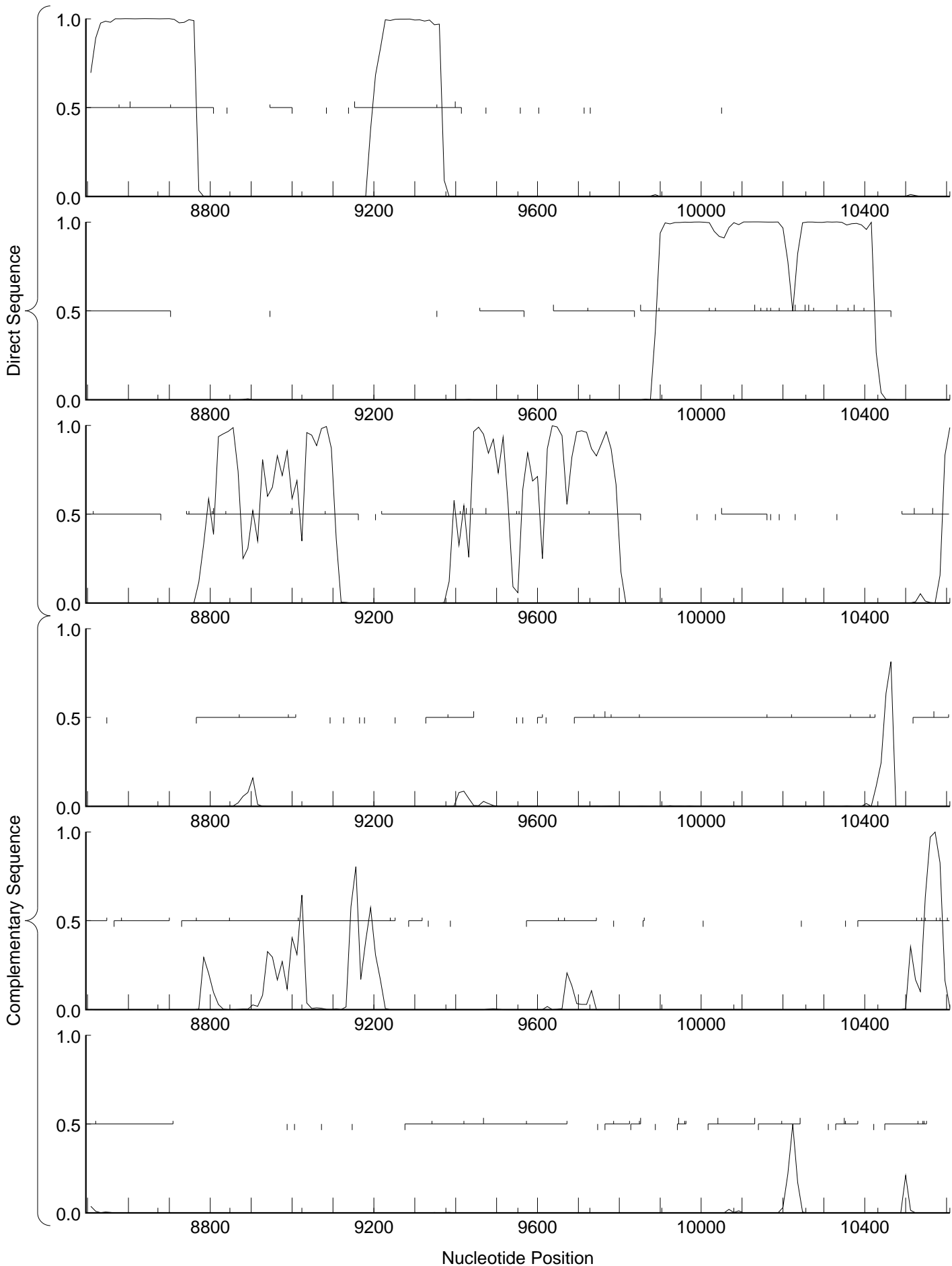
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

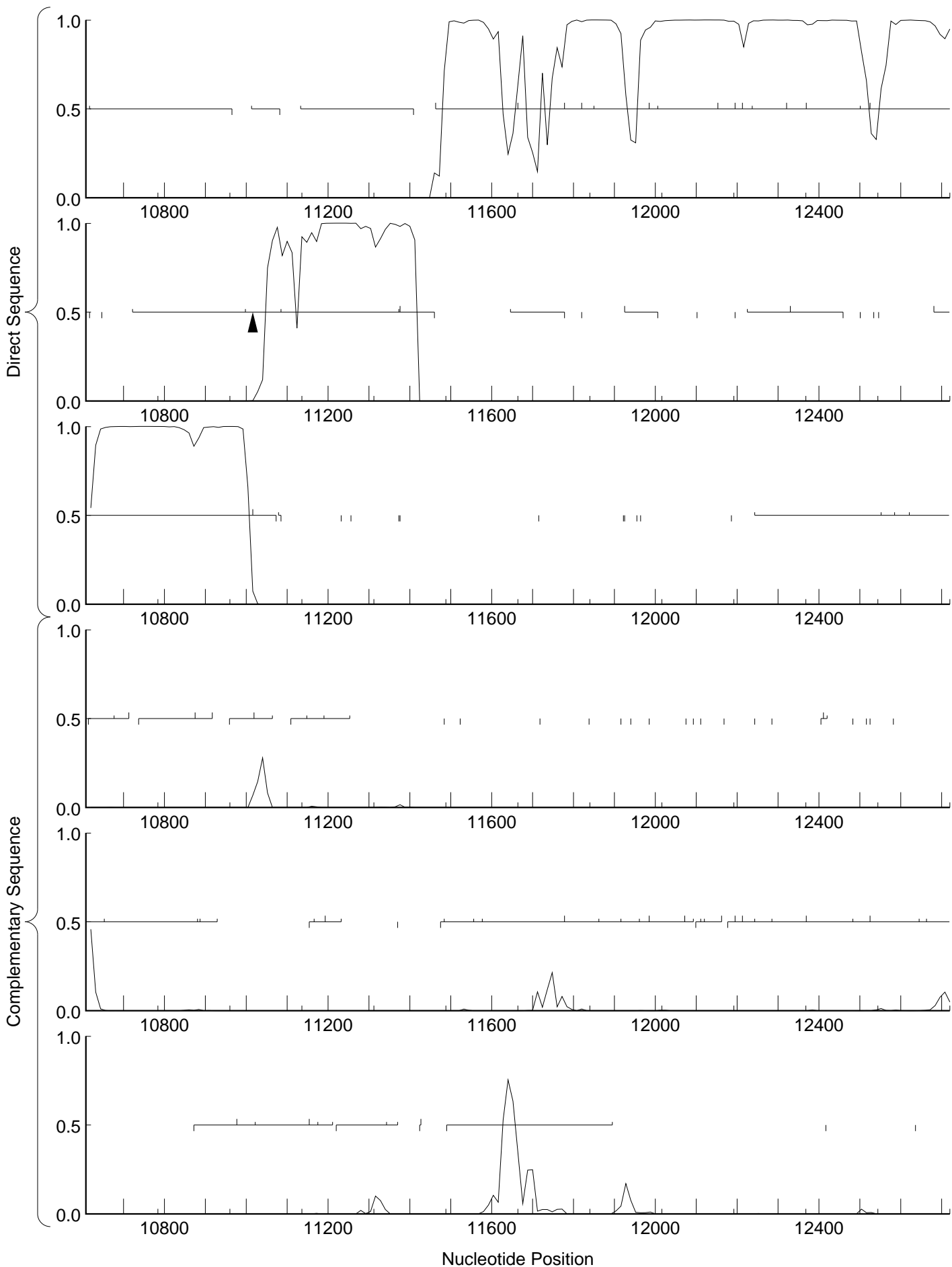


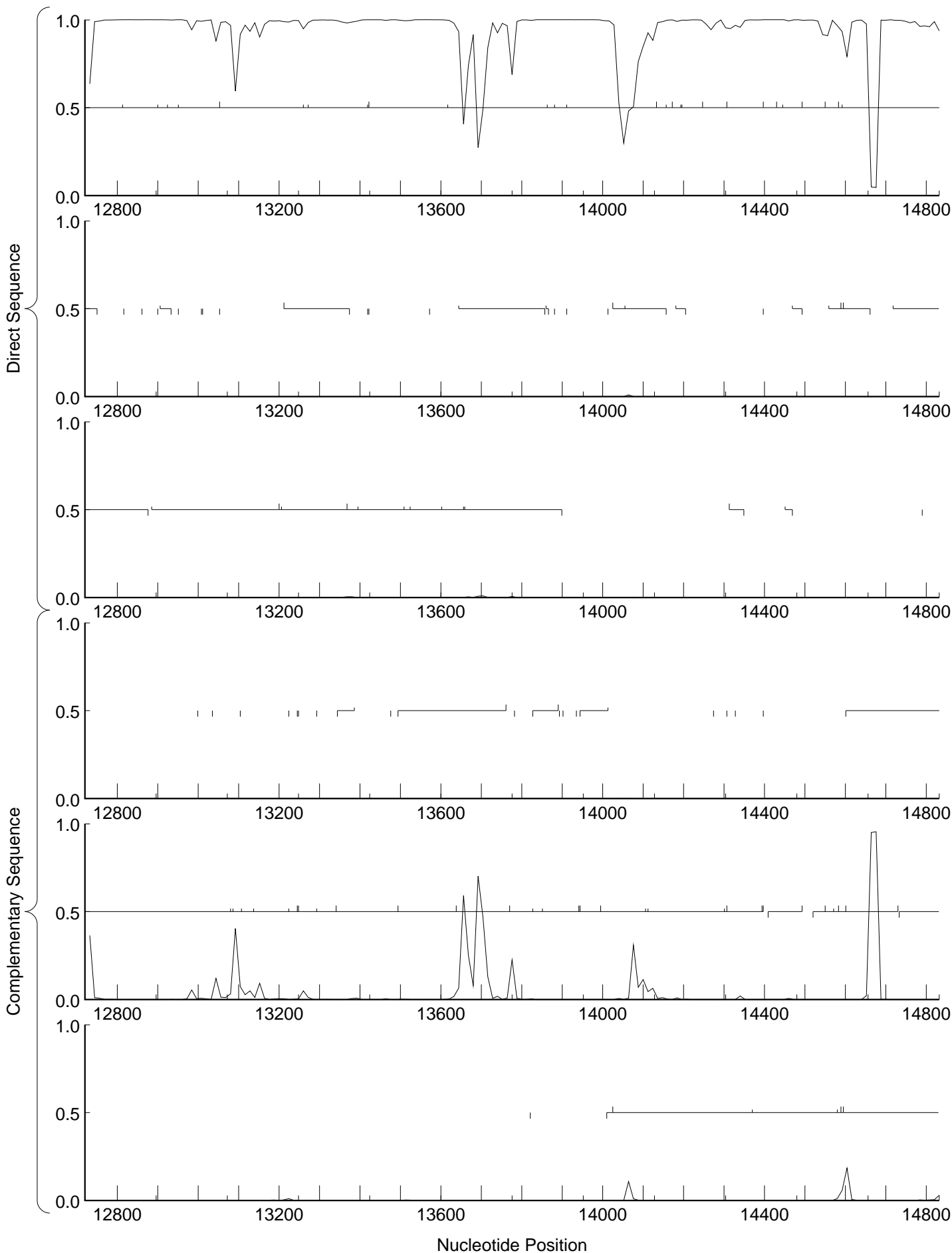


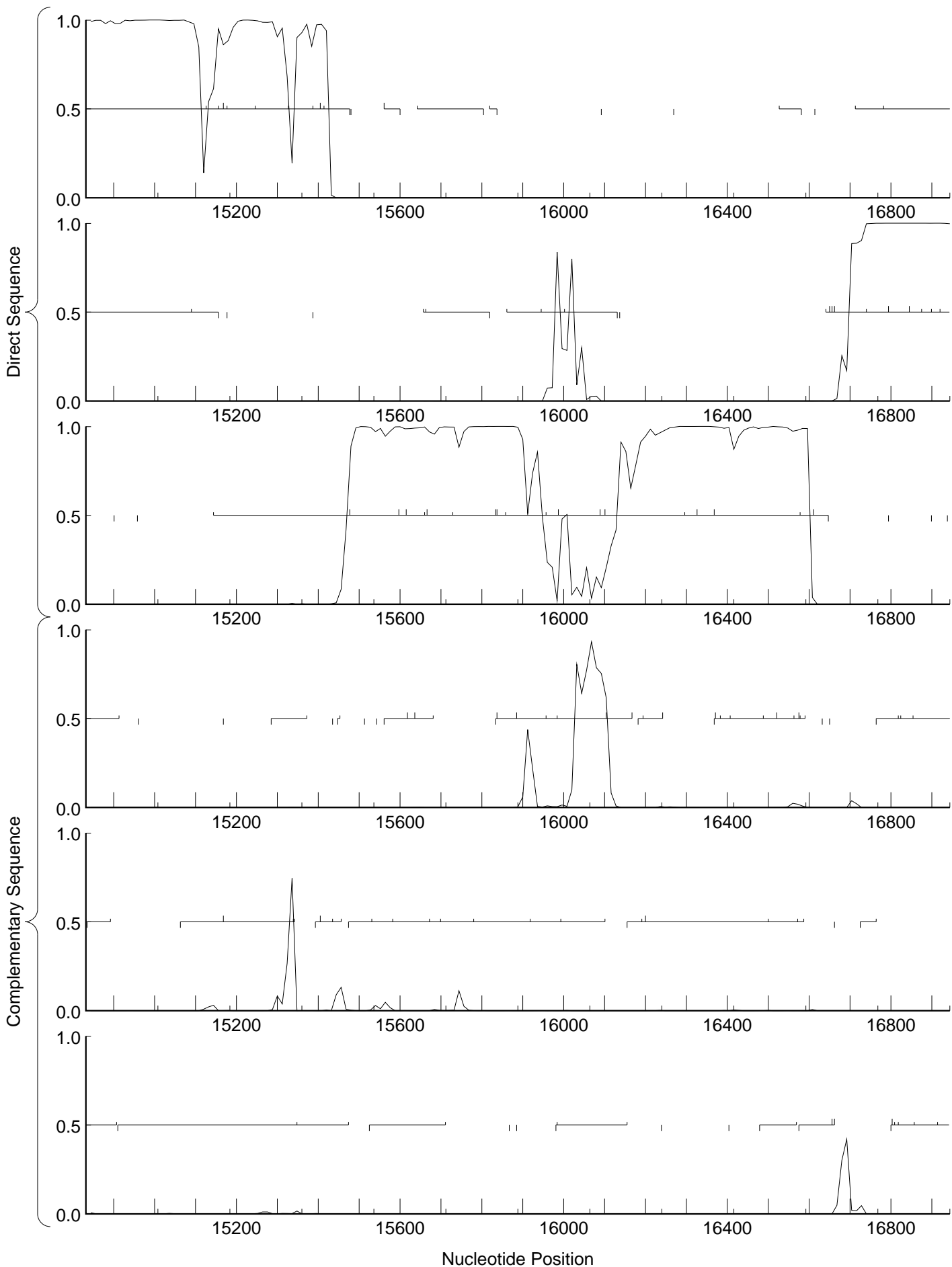


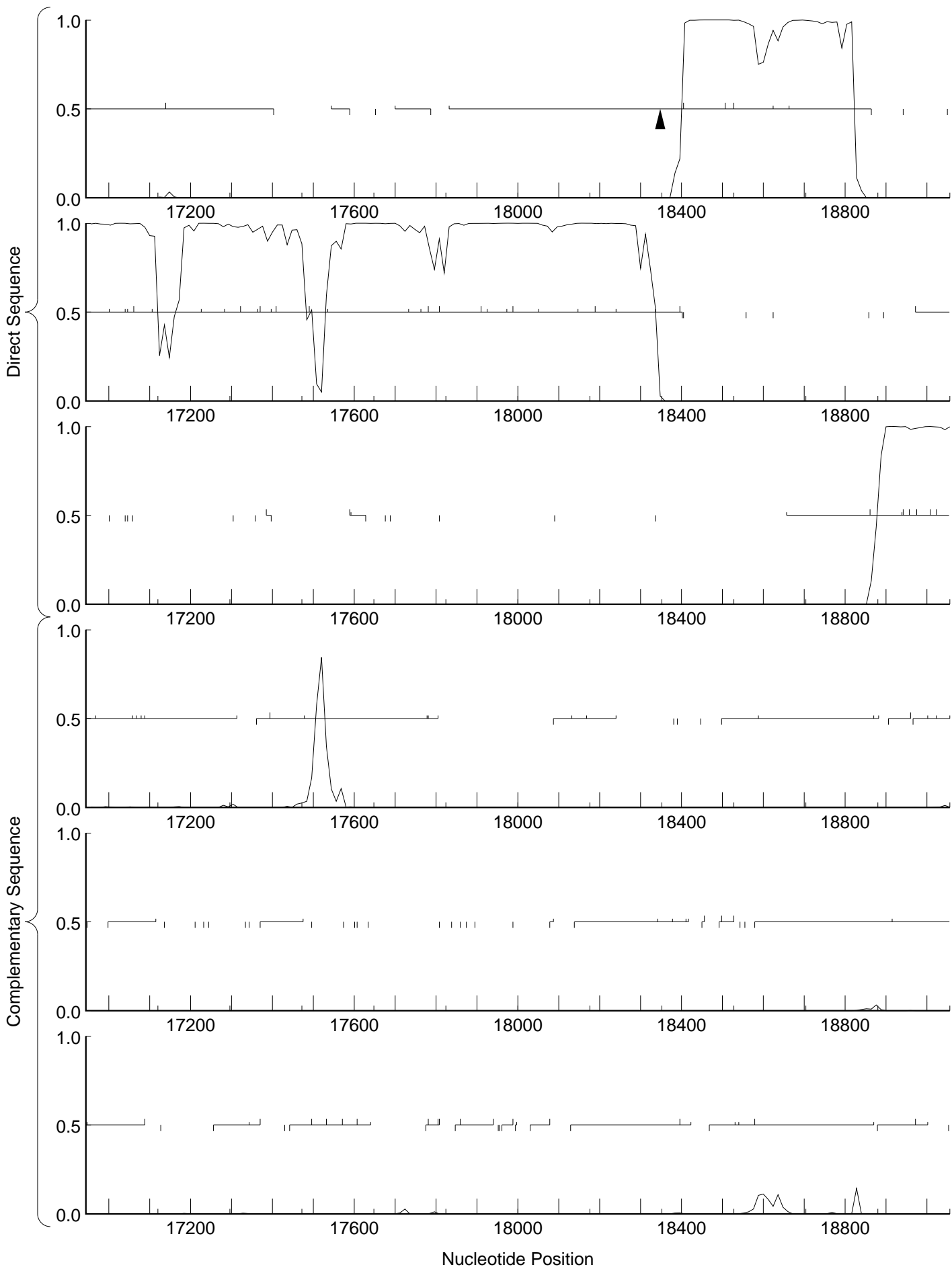


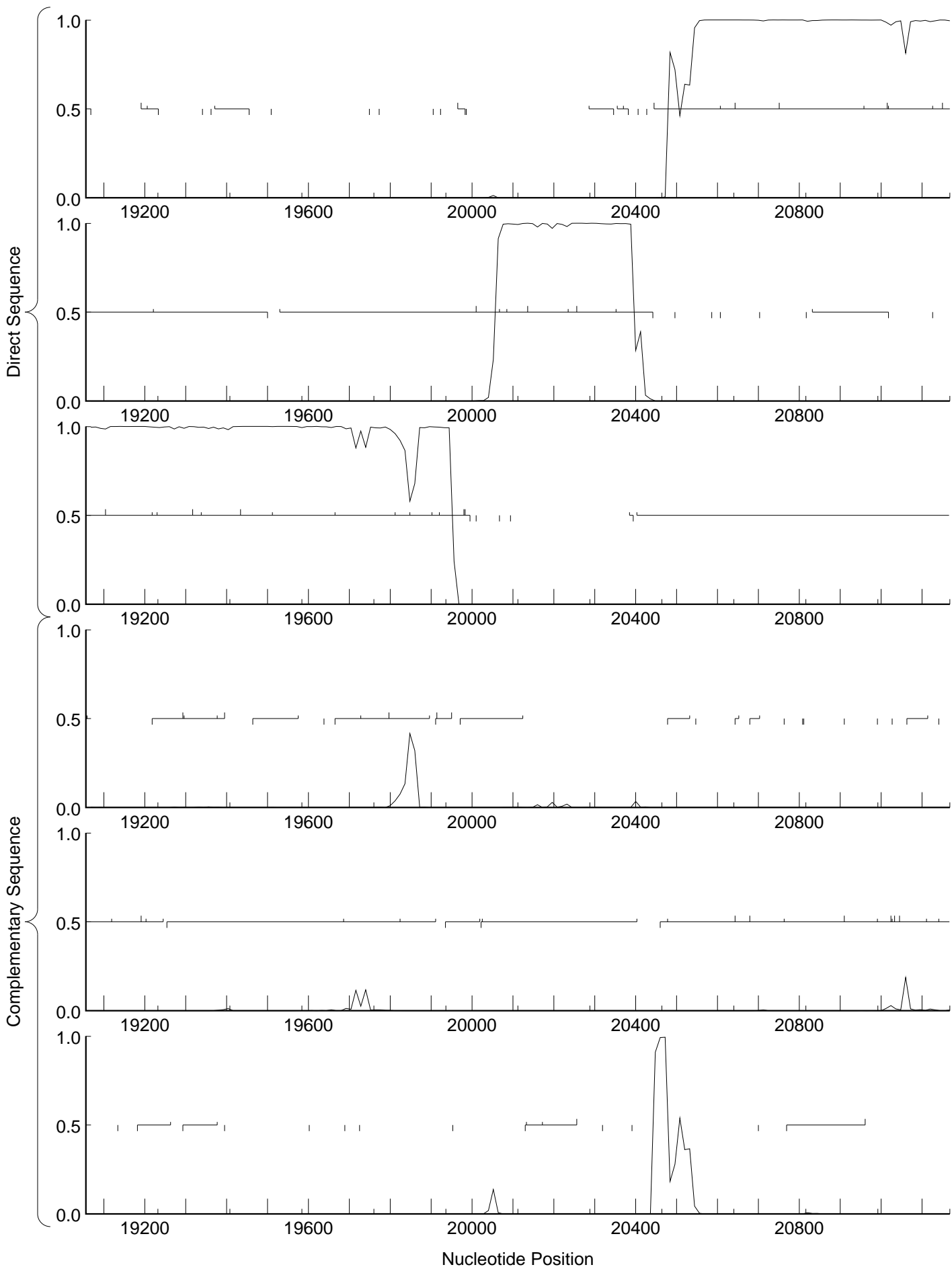




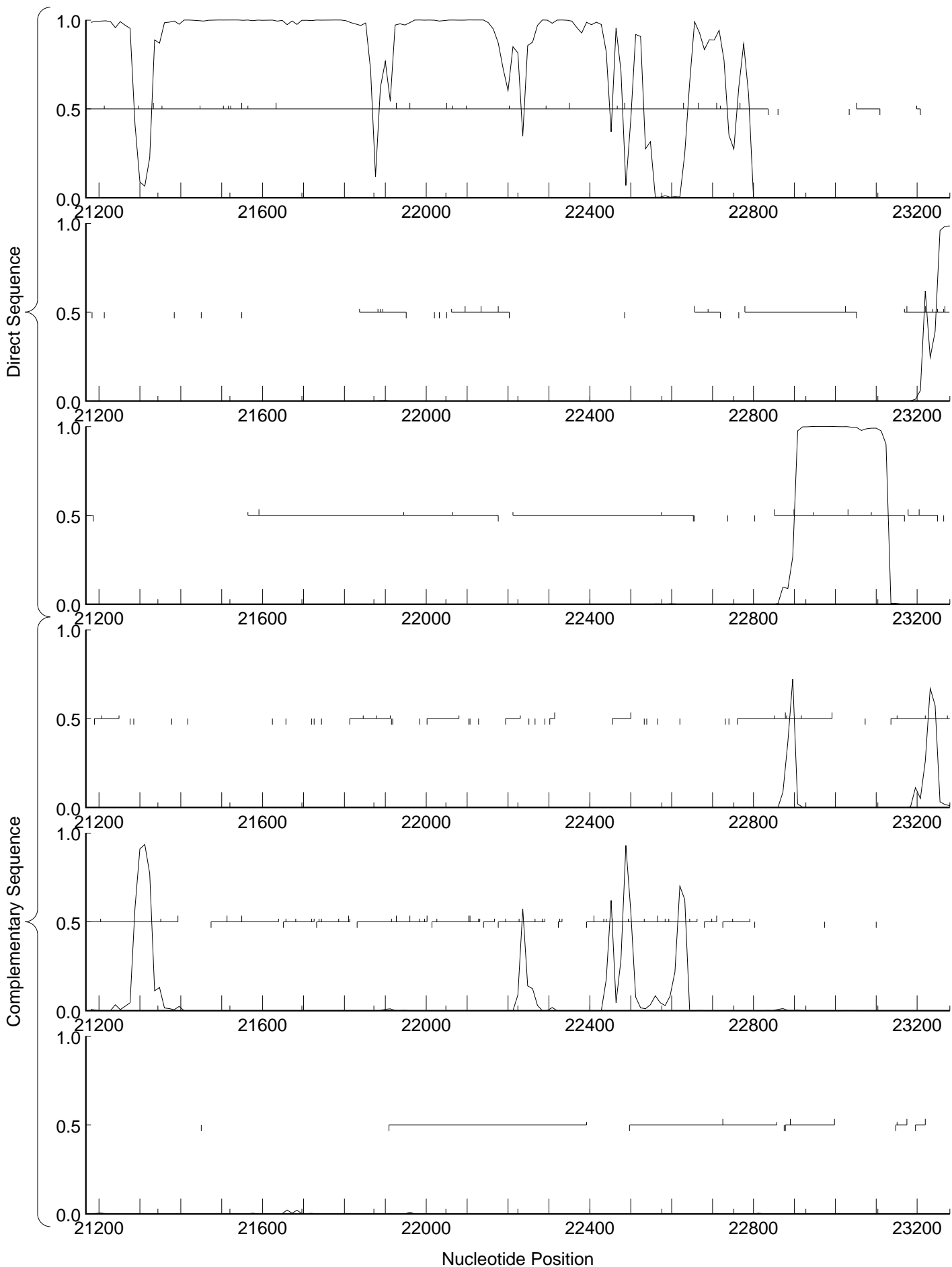




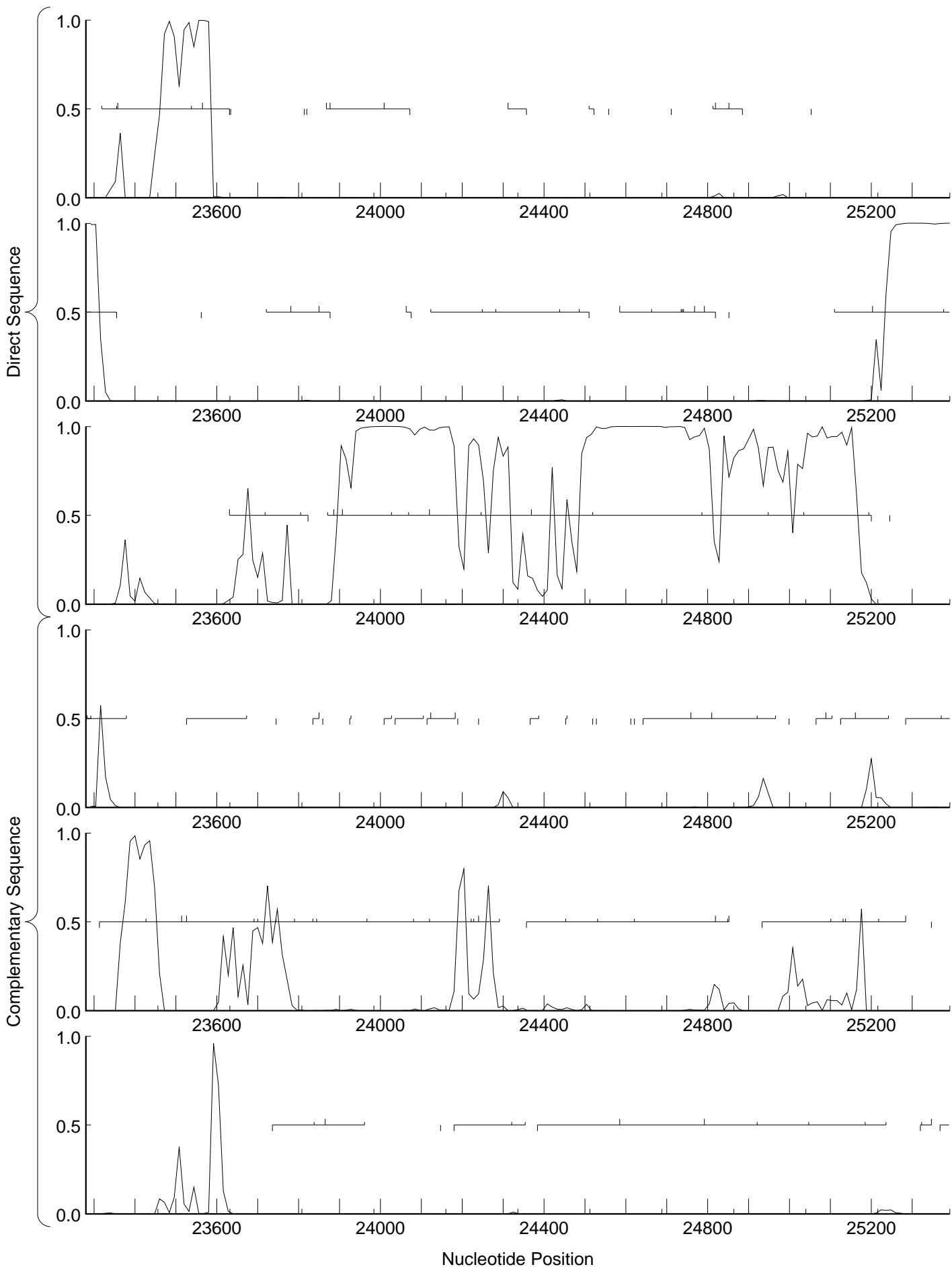


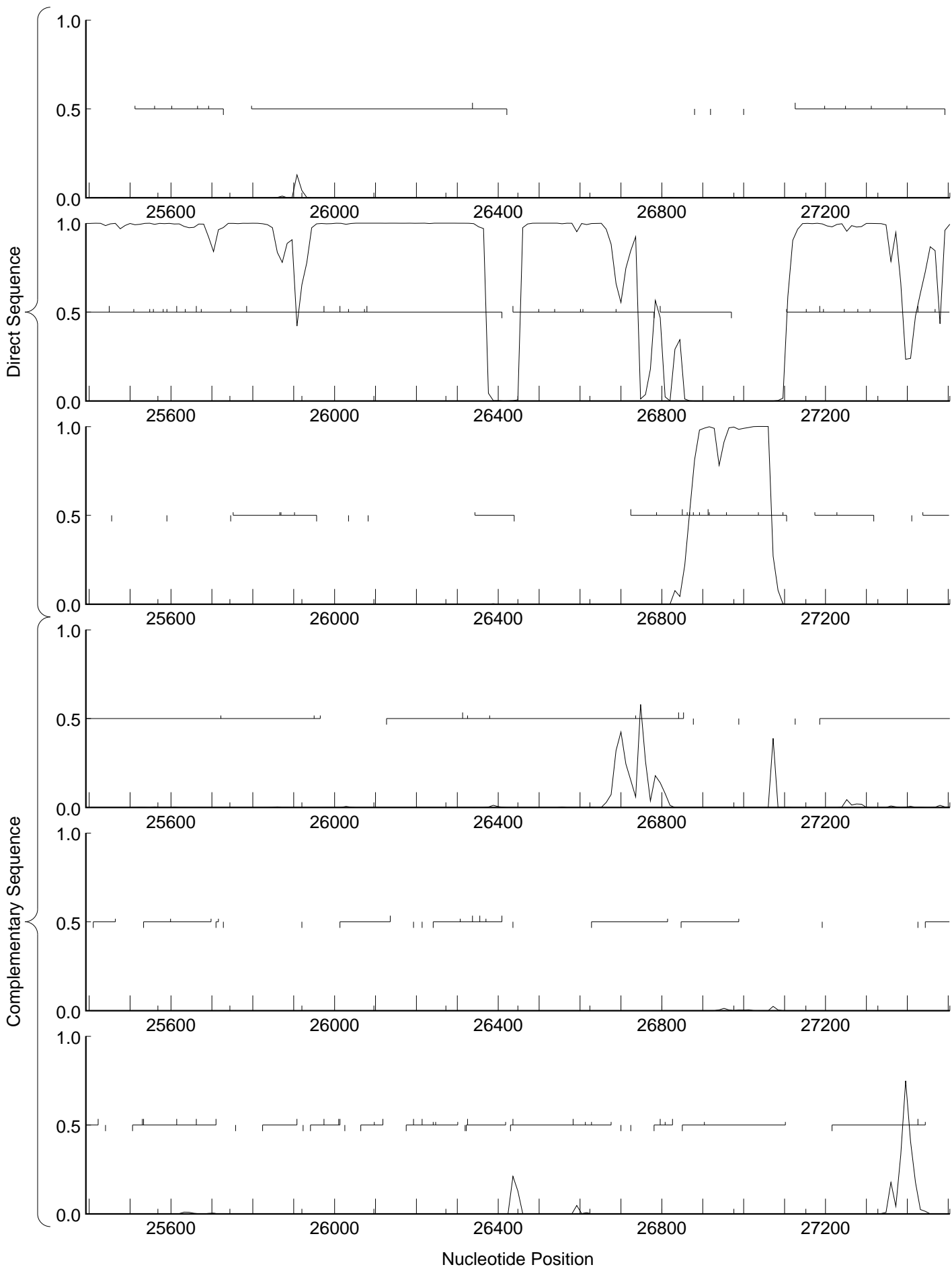


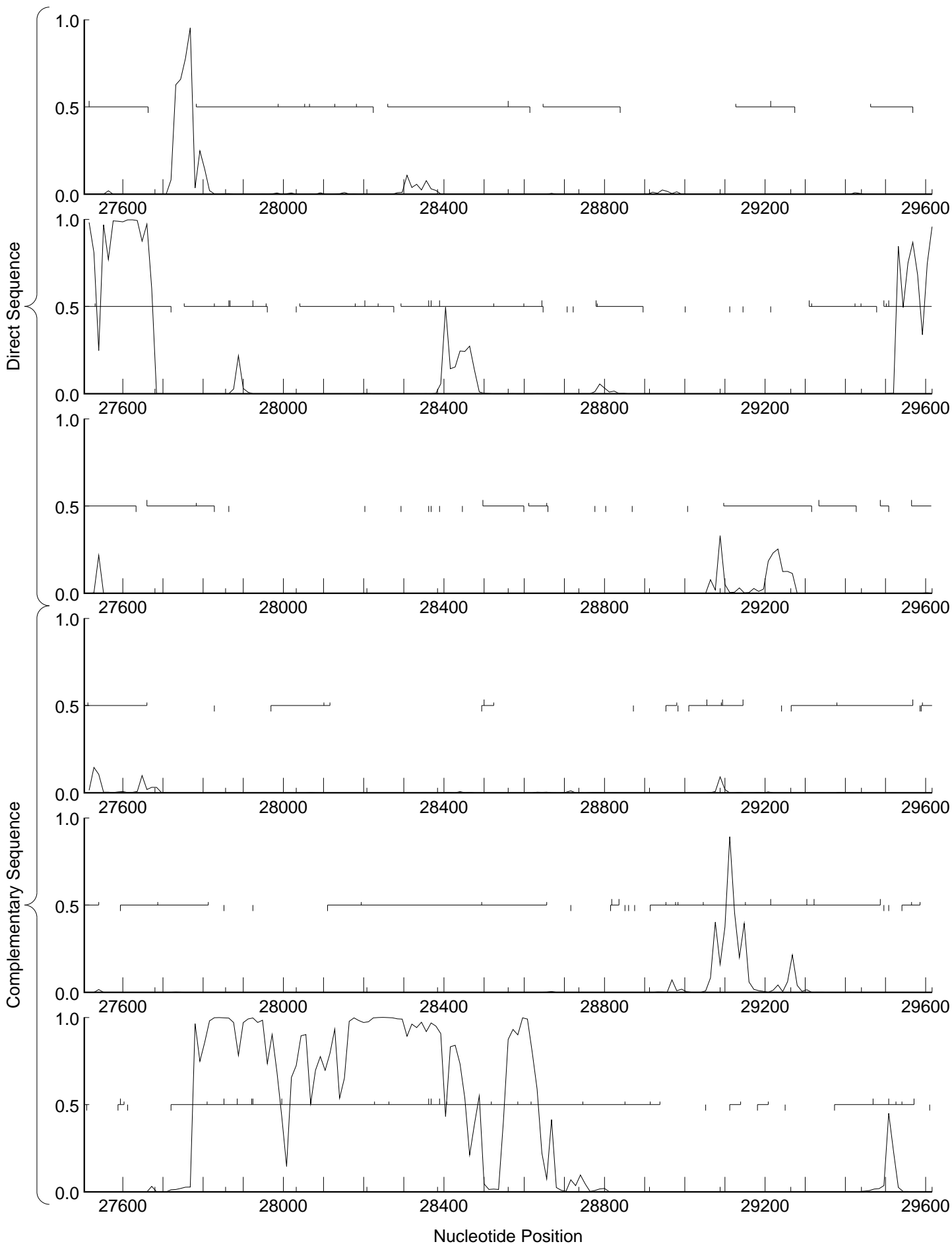
complete sequence, 41752 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 12/21



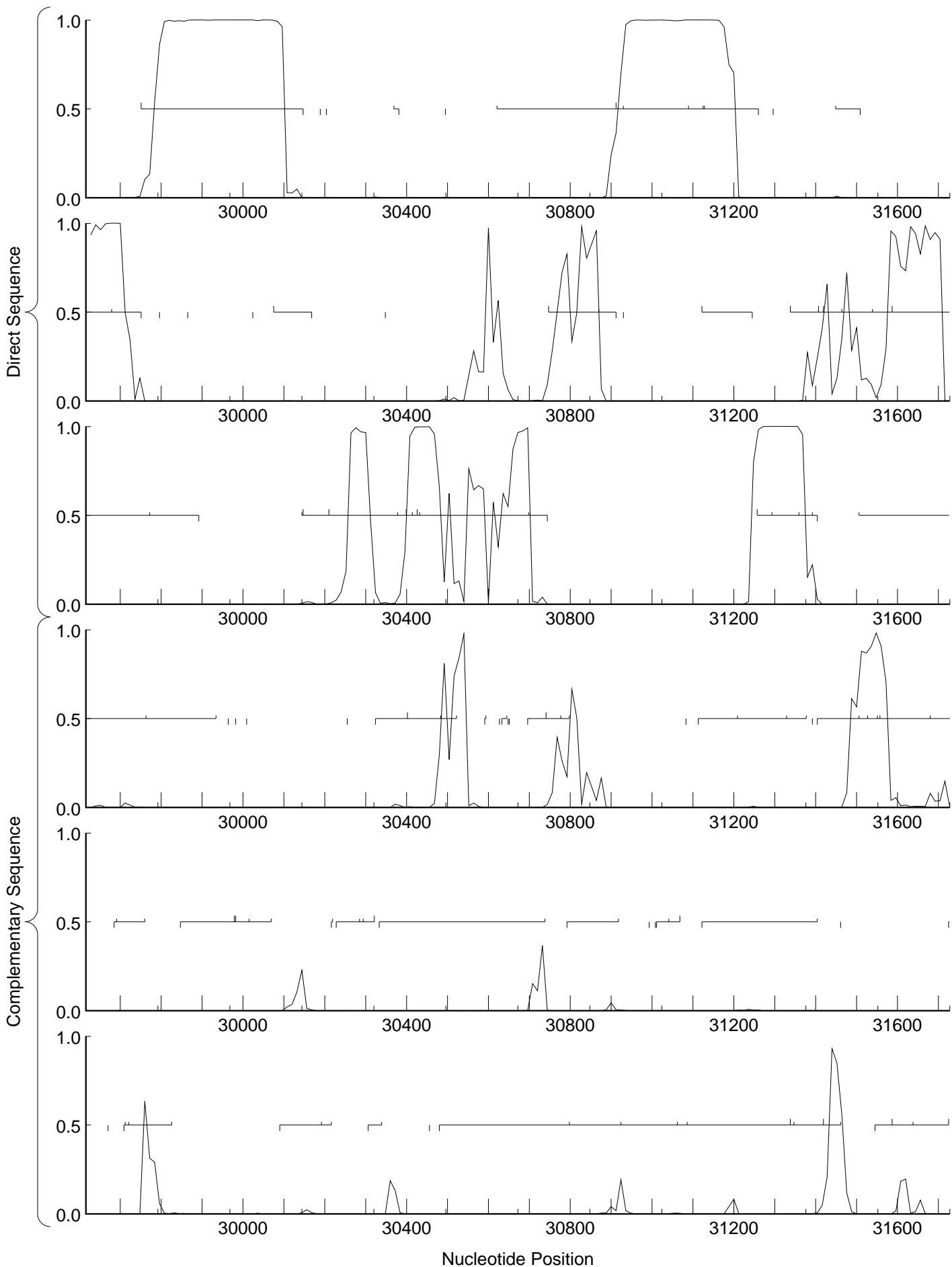
complete sequence, 41752 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 13/21



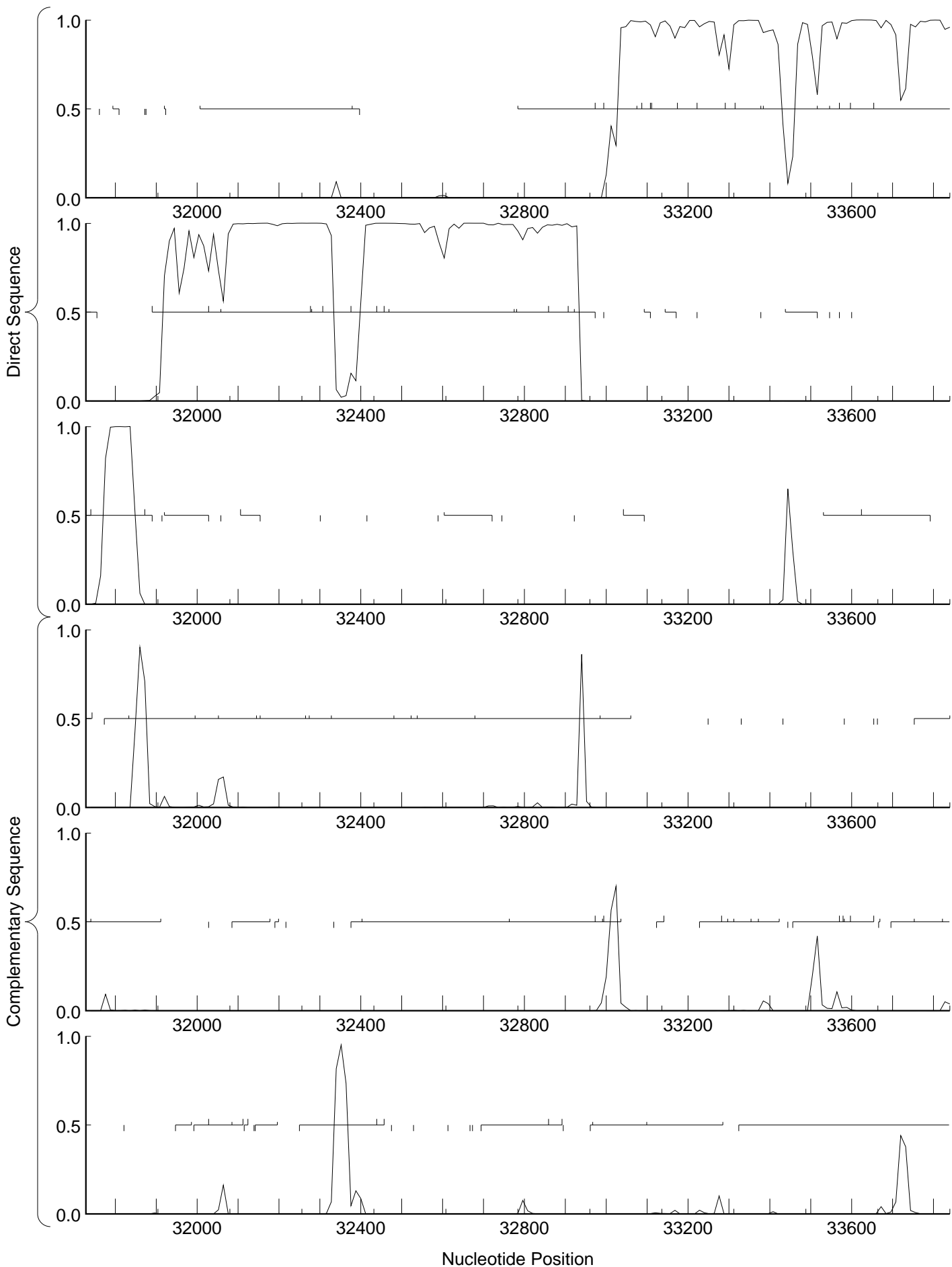


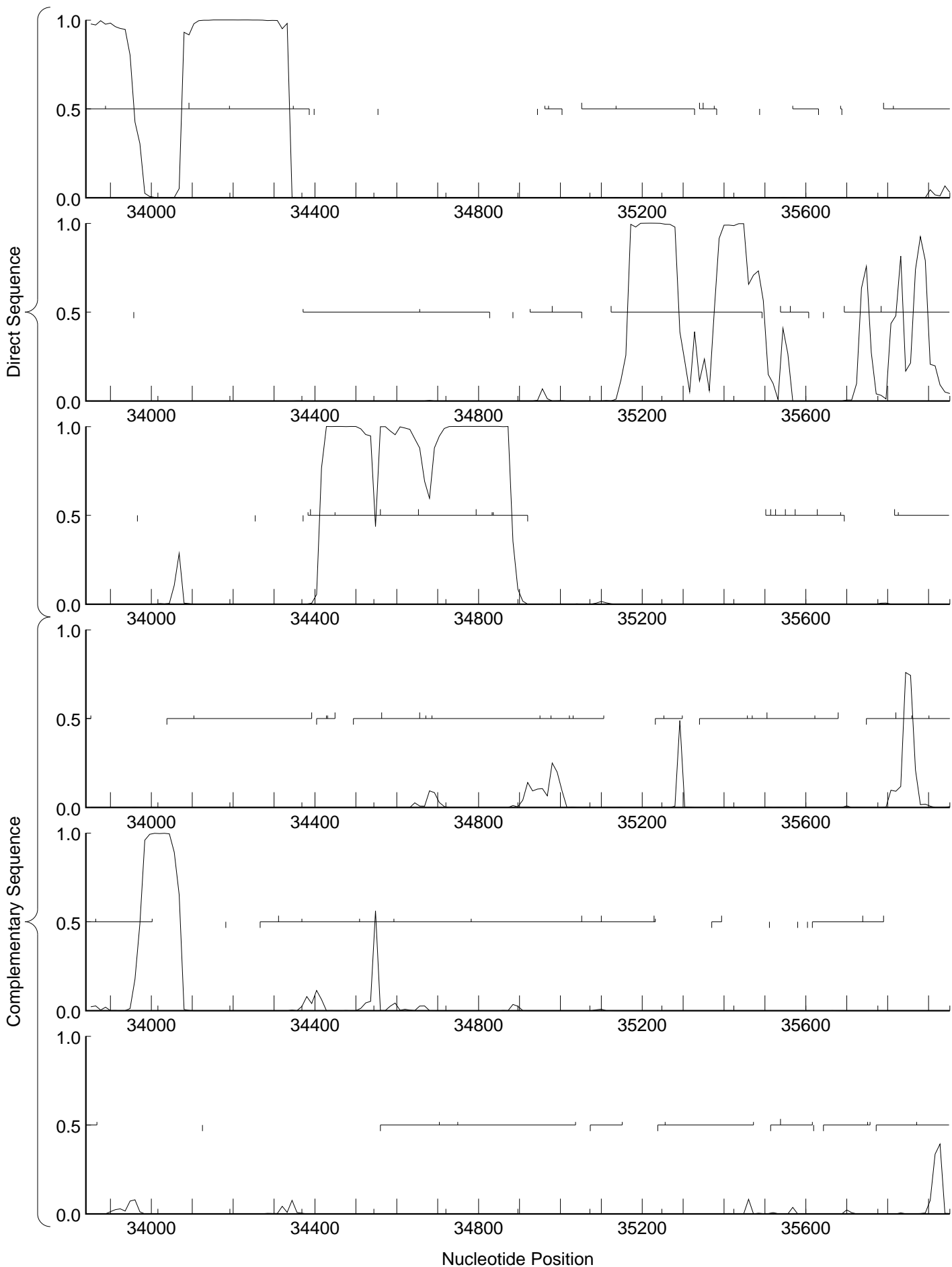


complete sequence, 41752 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 16/21



complete sequence, 41752 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 17/21





complete sequence, 41752 bp including 11-base 3' overhang (CCCCATGGCAT), Cluster G, Order 4, Window 96, Step 12, 19/21

