

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Efra2 complete sequence, 61284 bp including 11-base 3' overhang (CTCGTA
Analysis Date : 8/25/19 at 18:20:04
Pages : 31
Sequence Length : 61284 bp
GC Content : 66.56%

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,

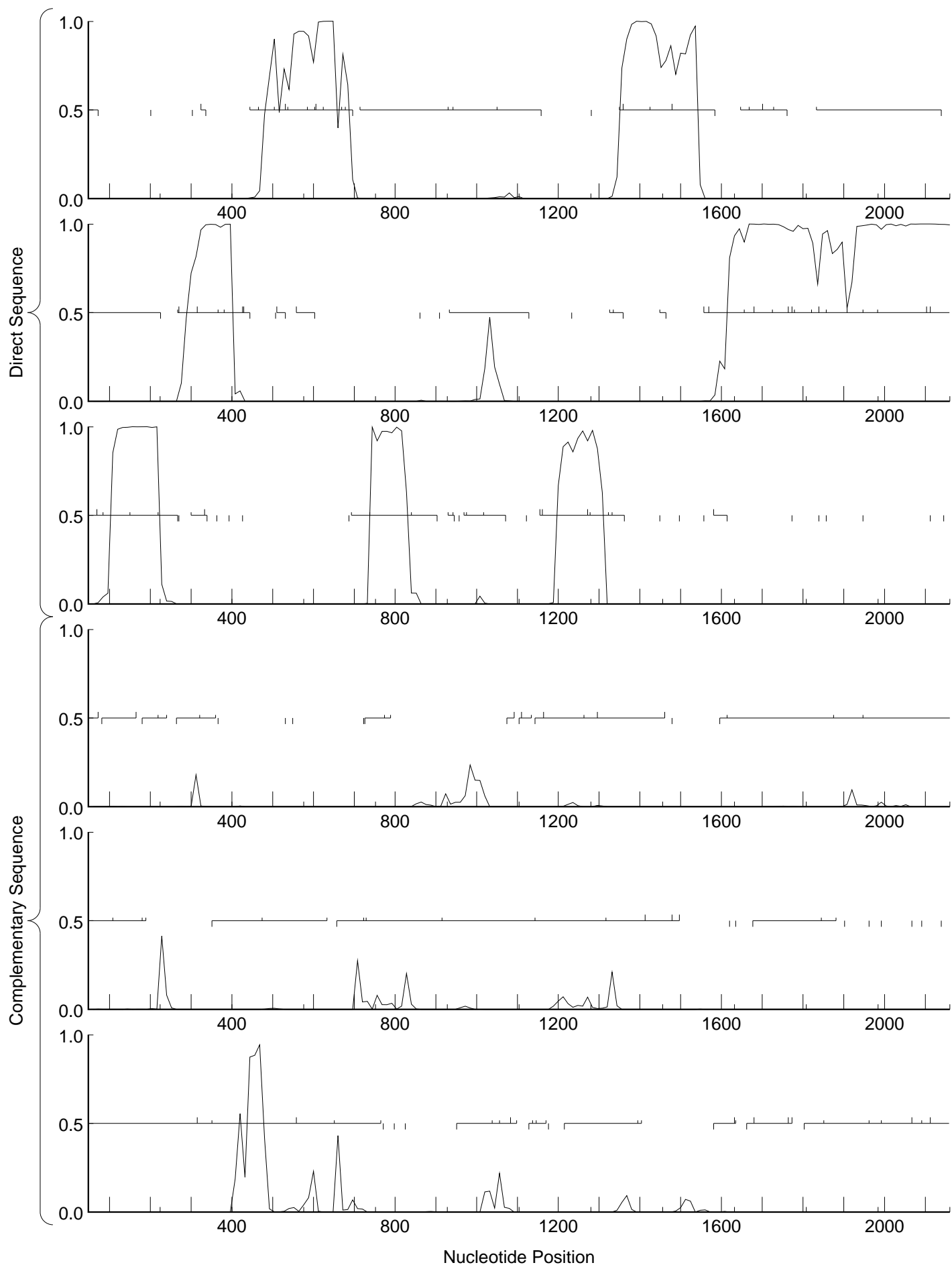
Matrix notes & comments

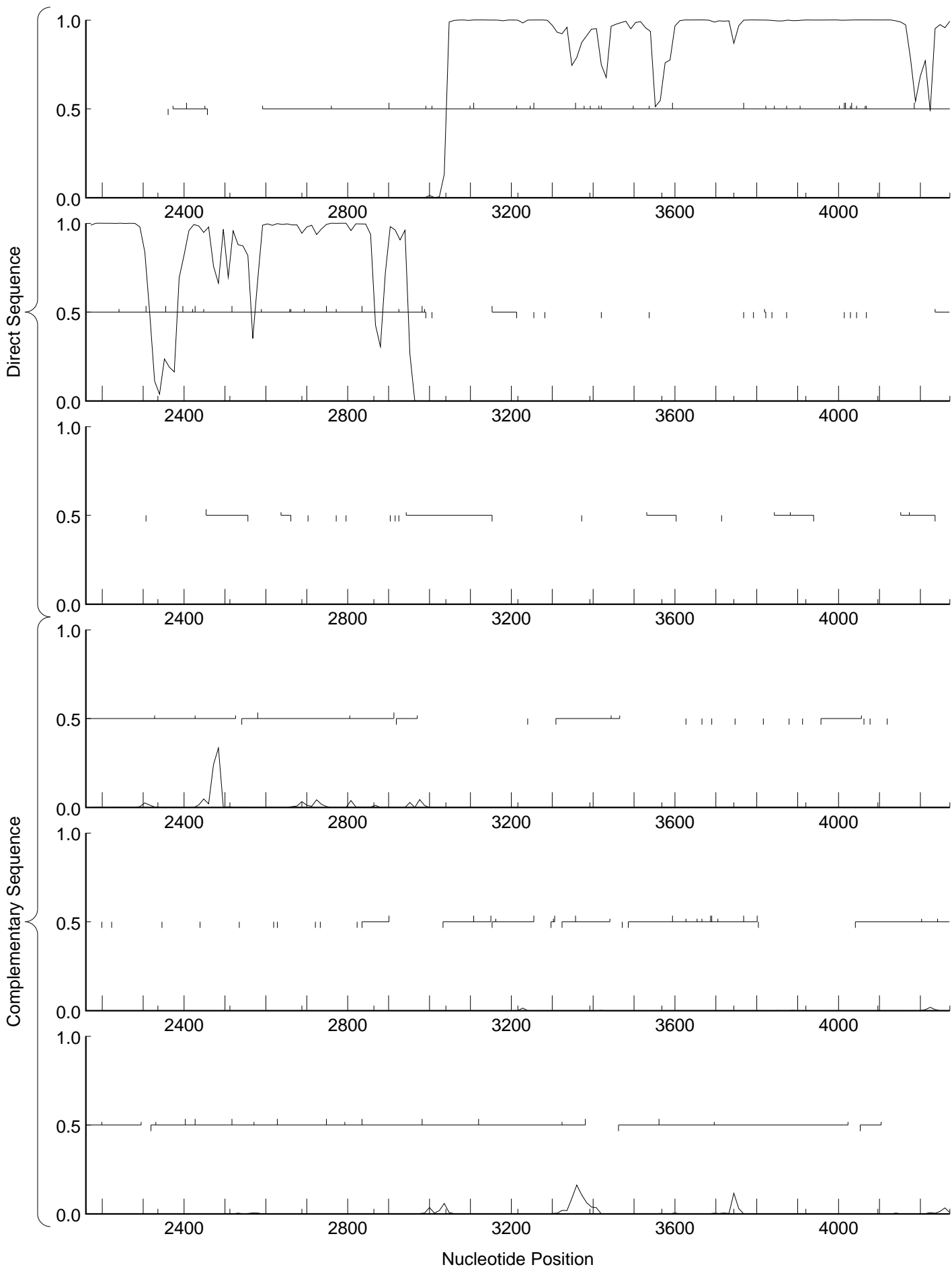
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

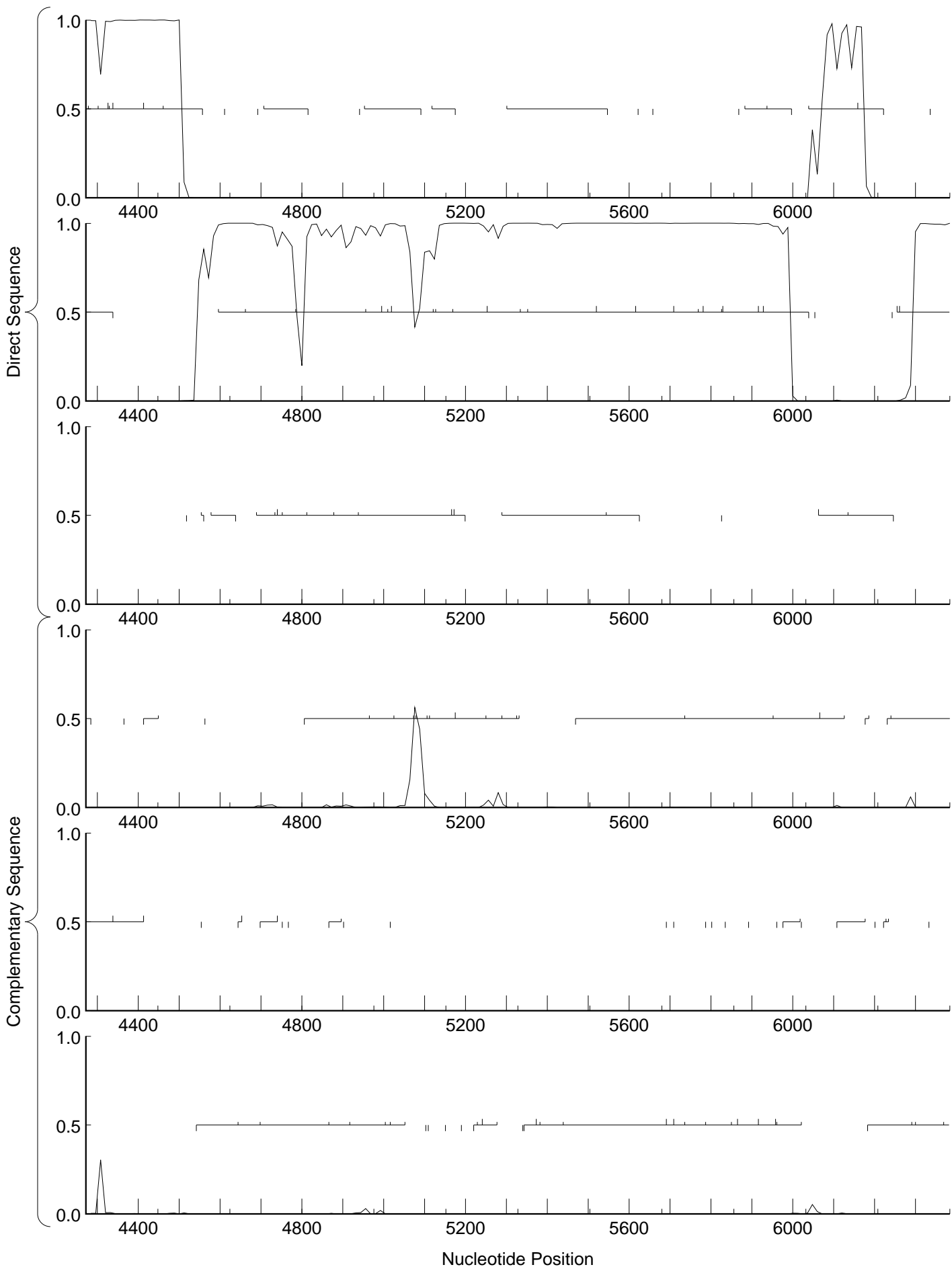
MATRIX INFORMATION

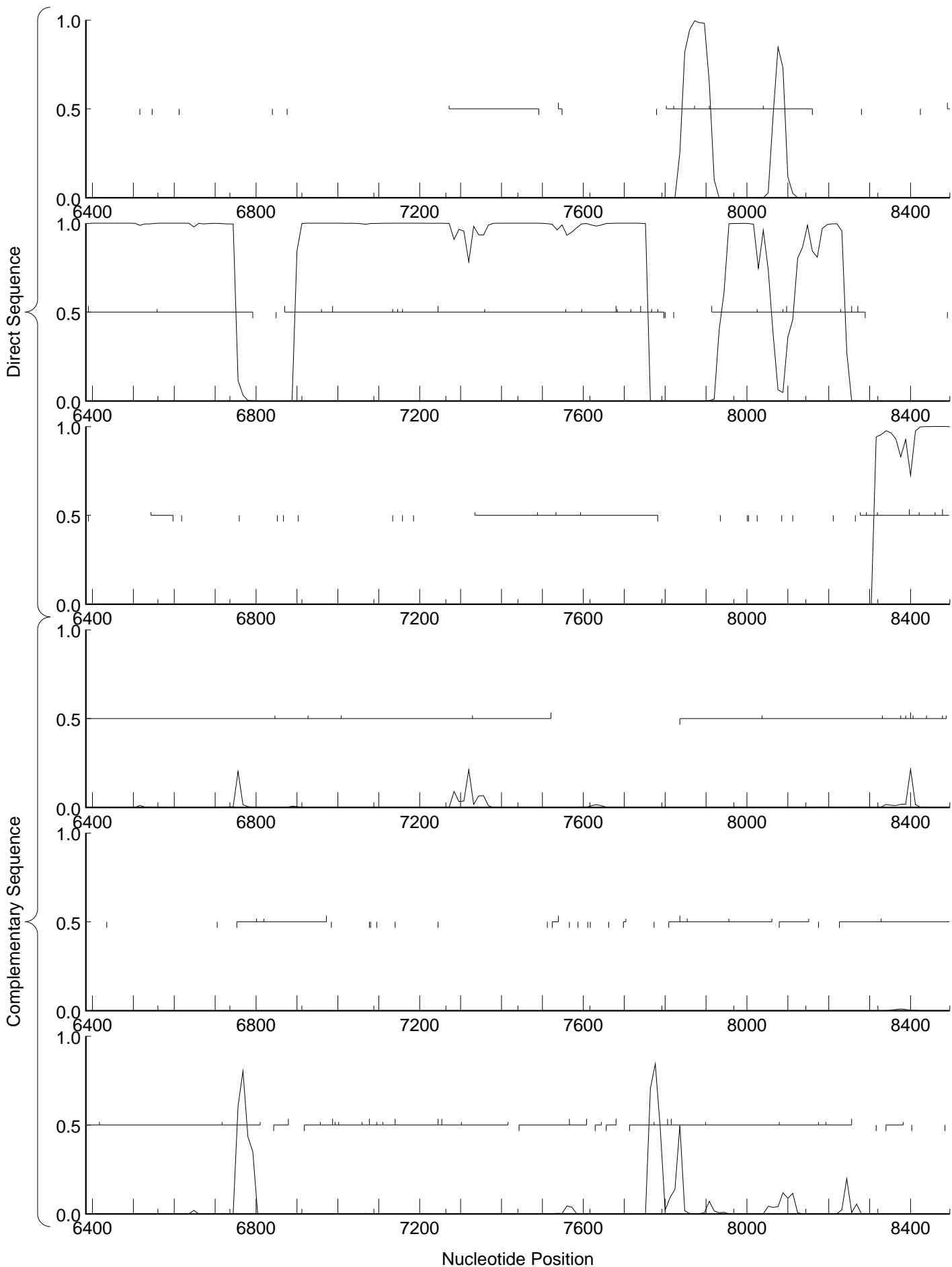
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

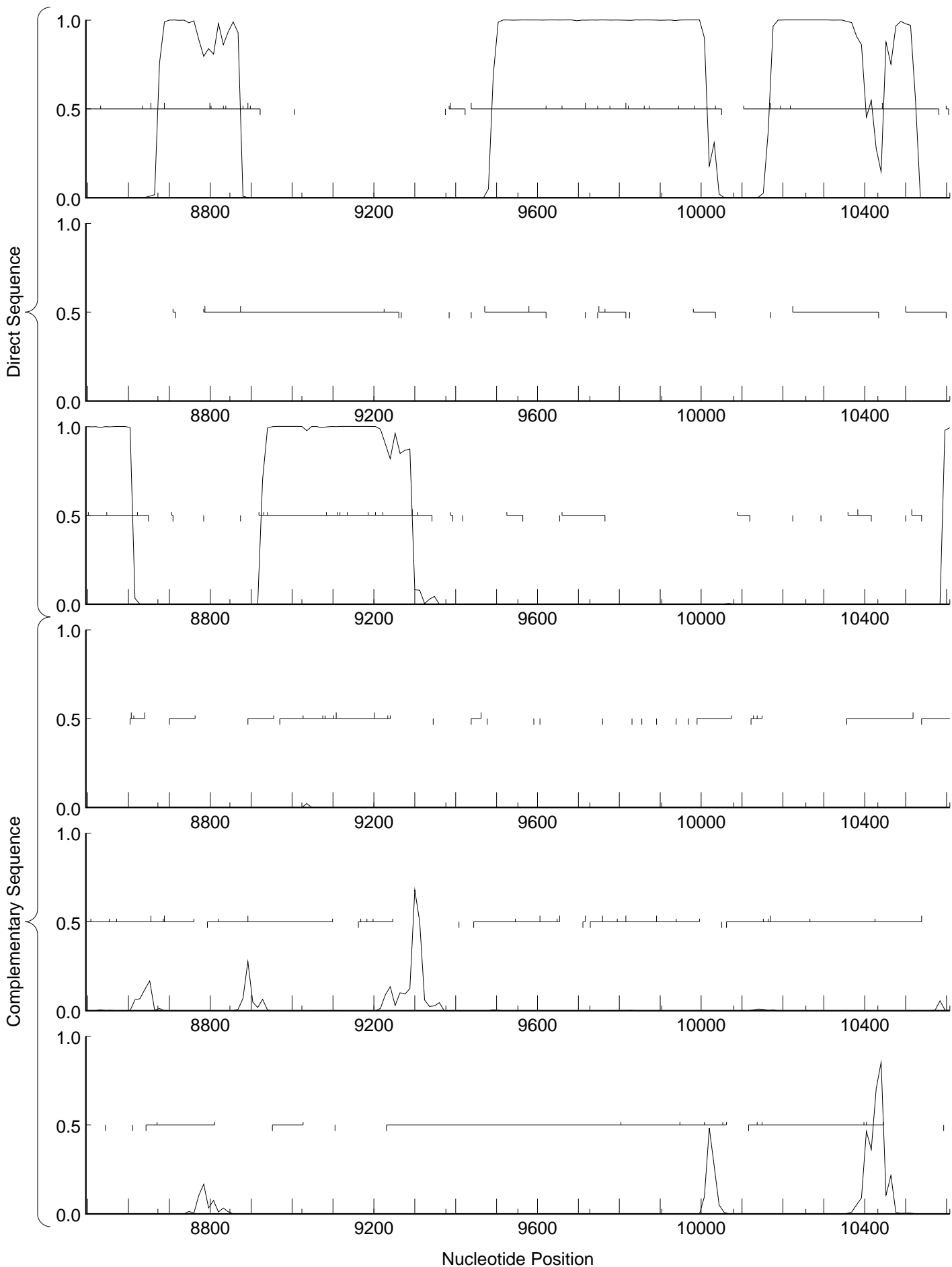
Send questions / comments to:
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Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

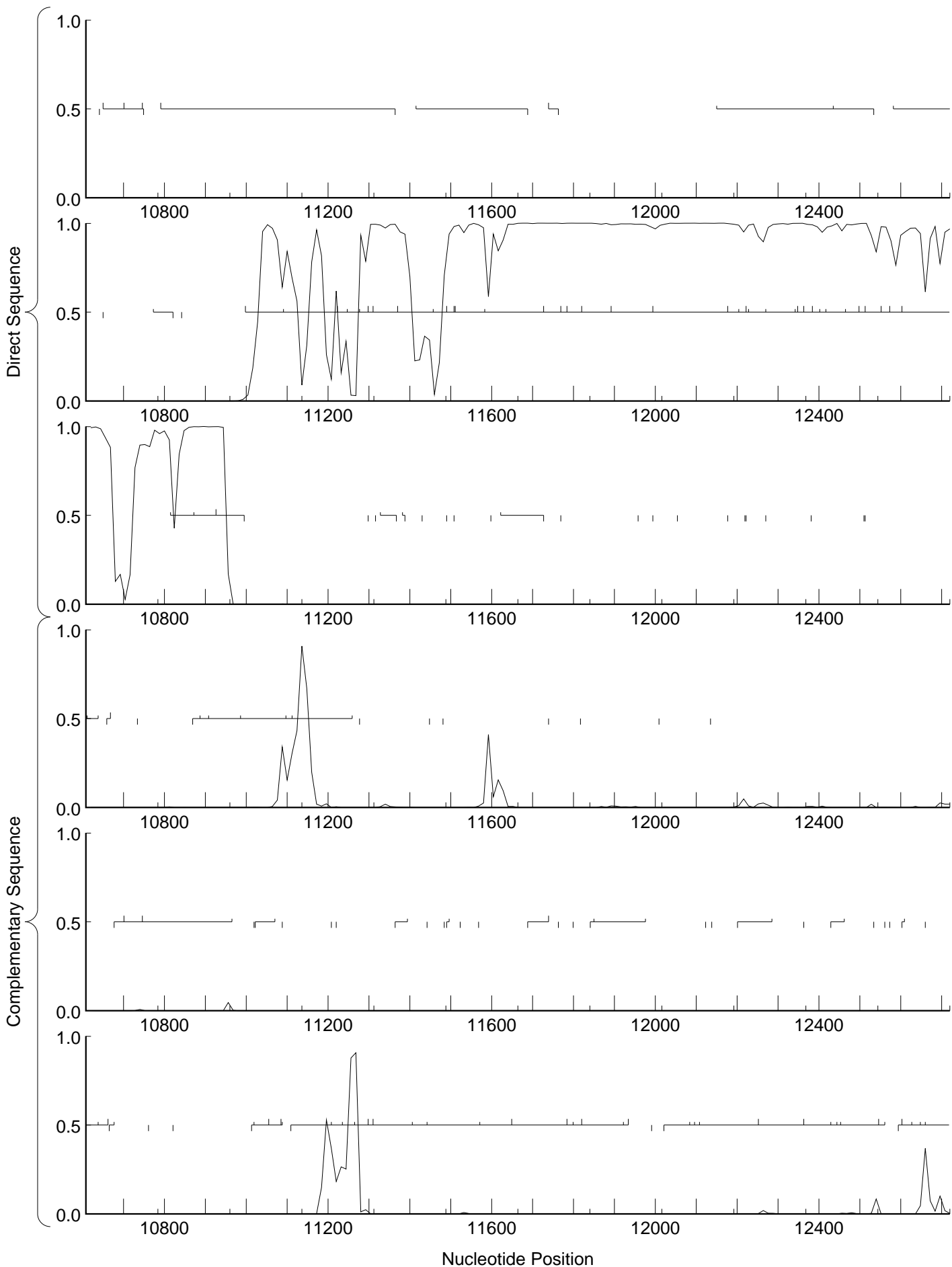


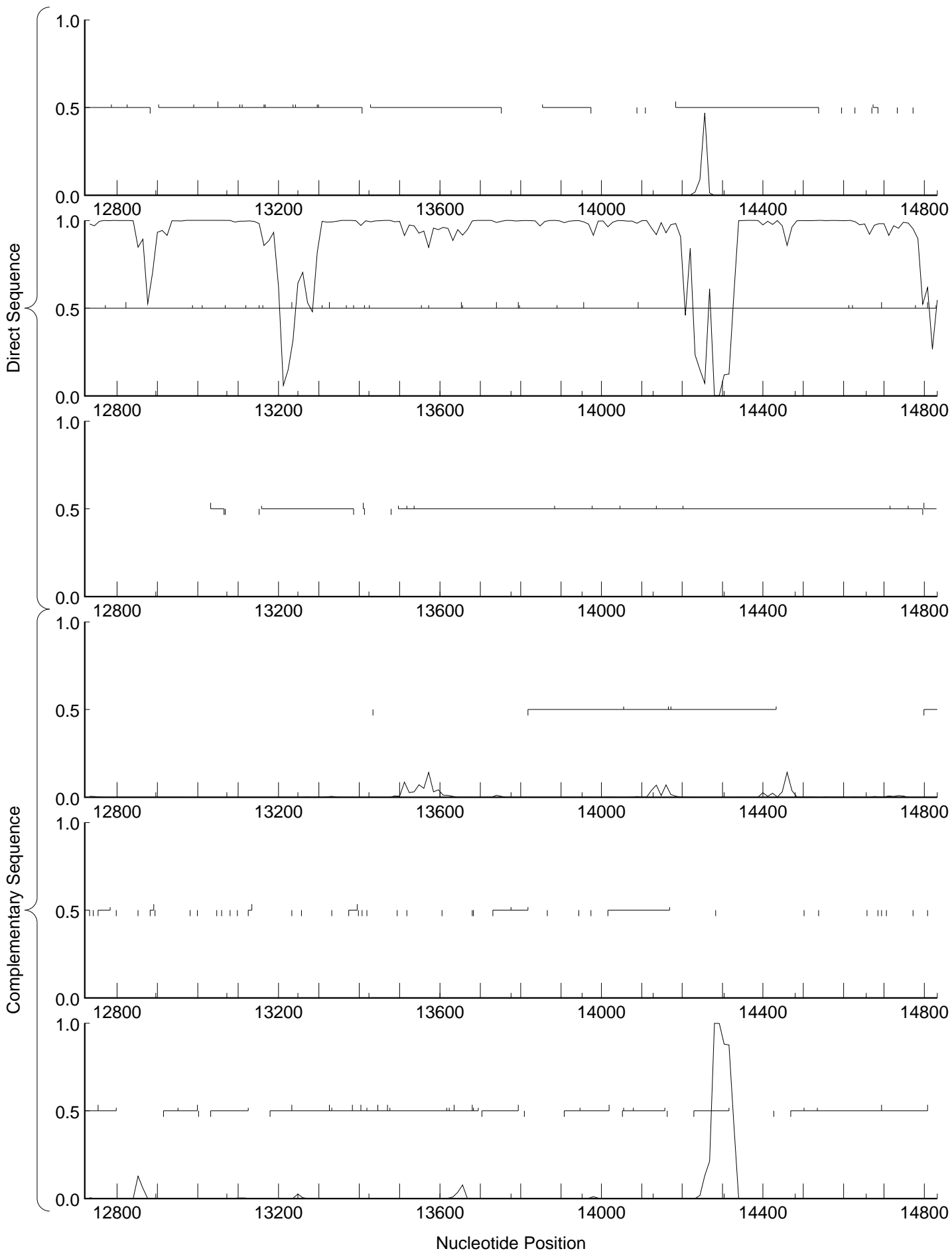


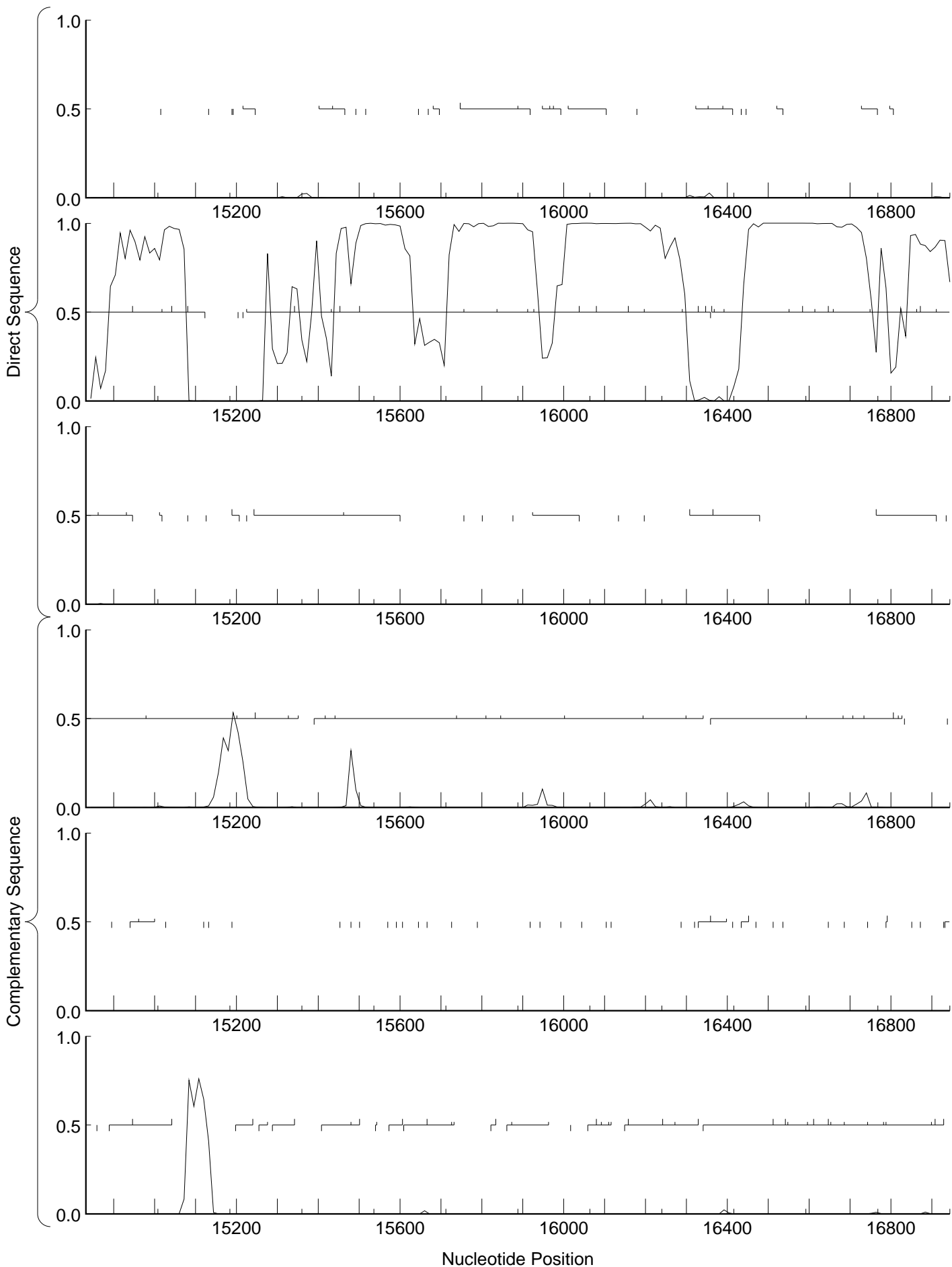


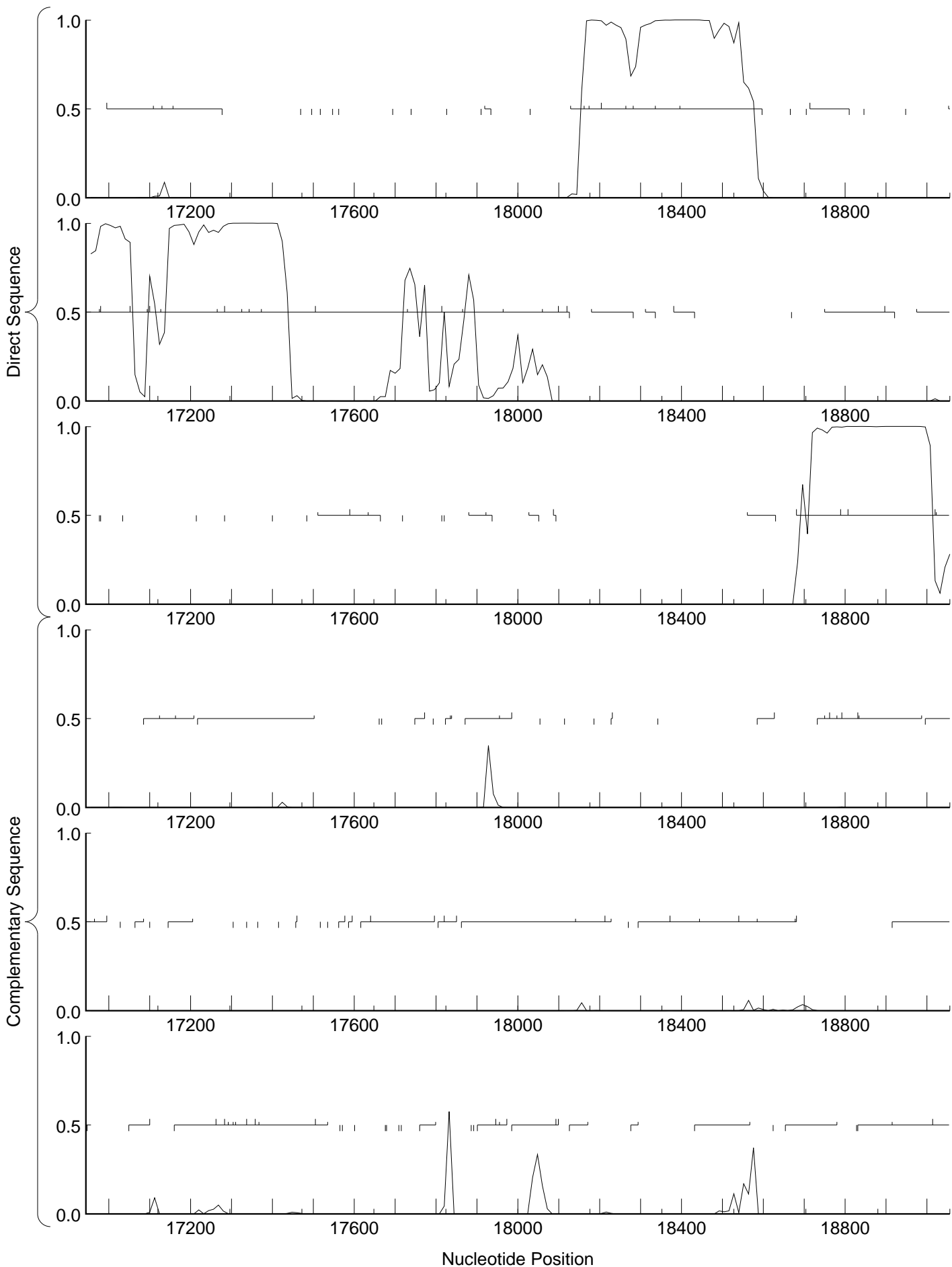


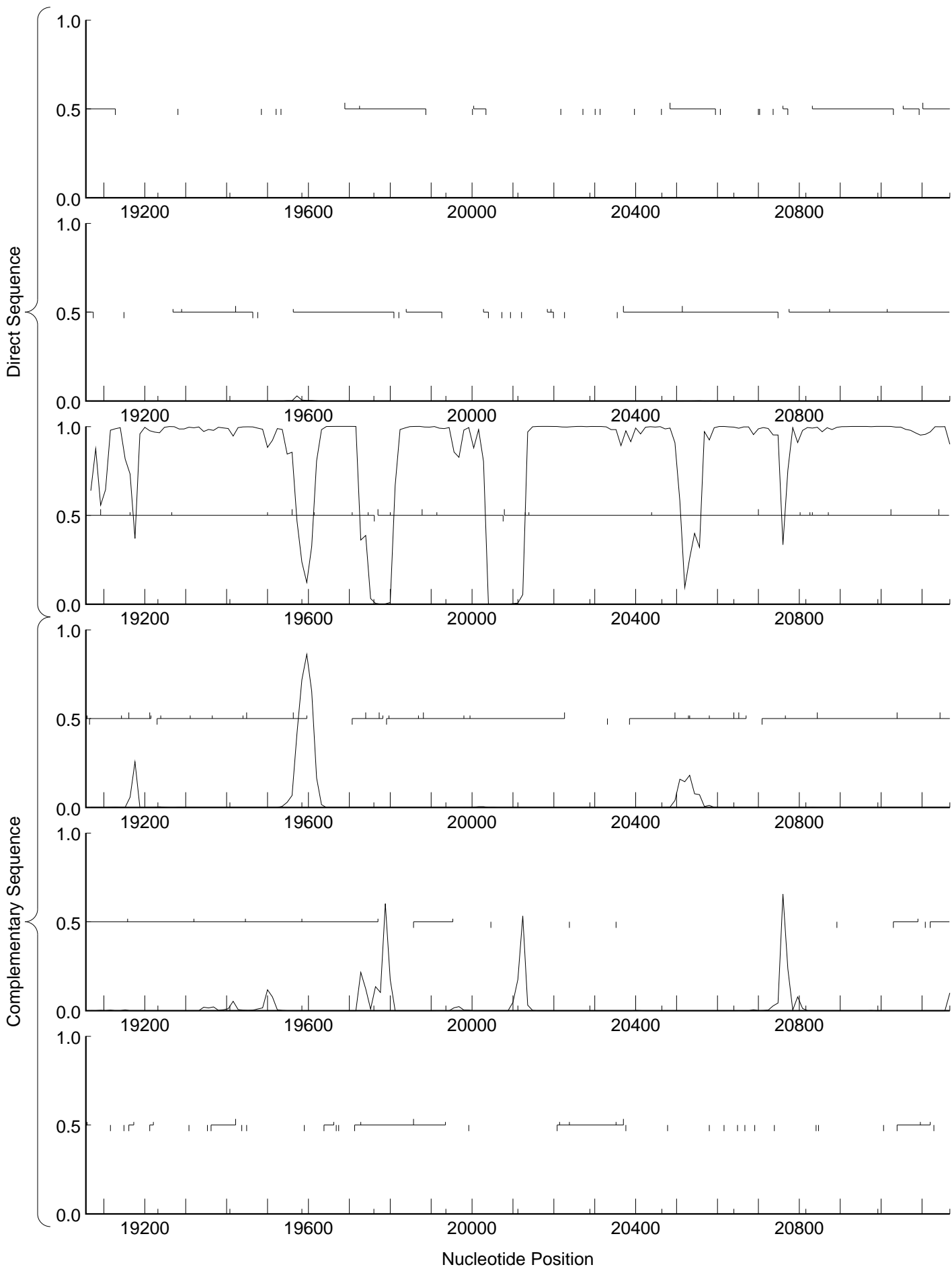


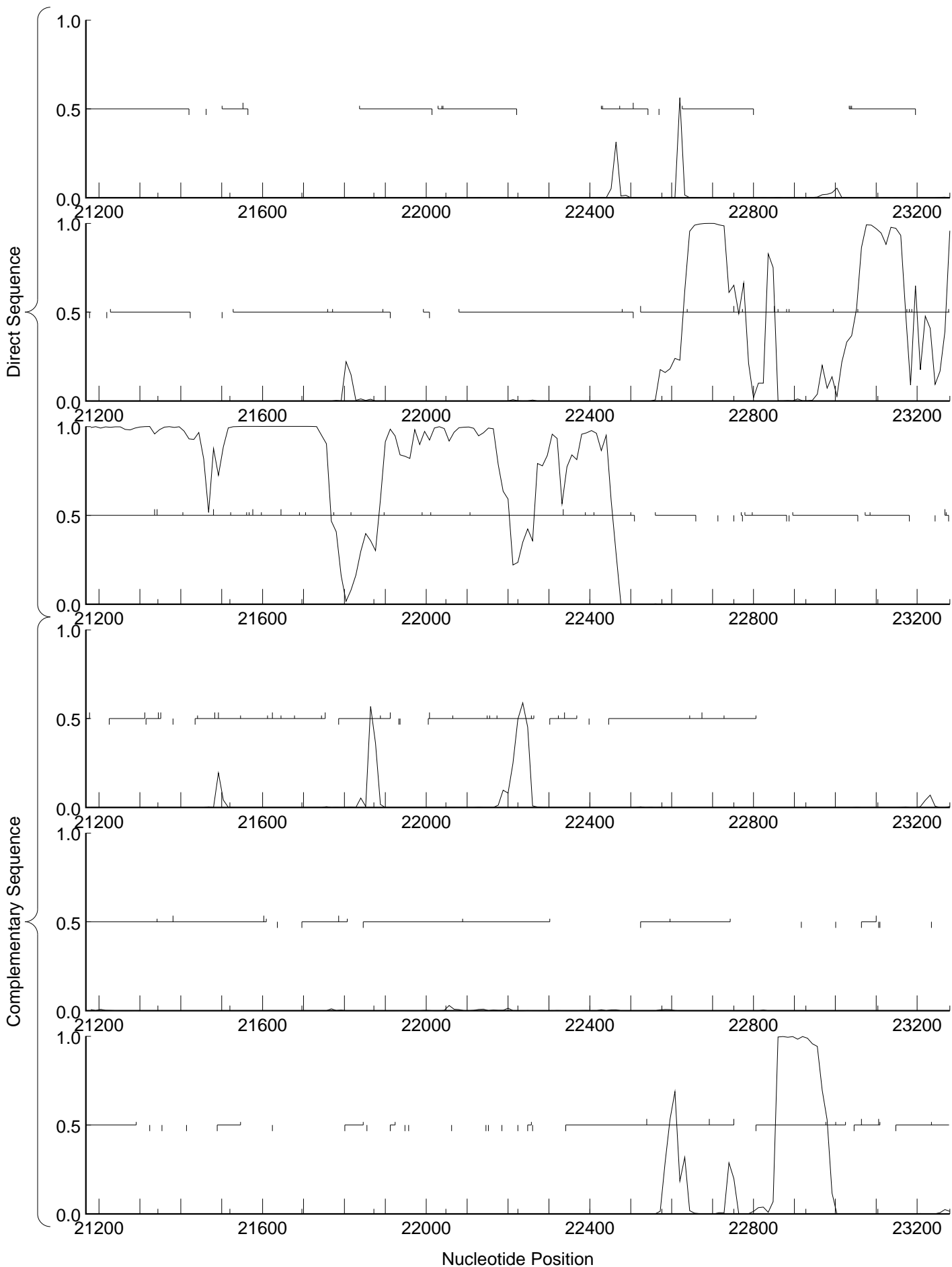


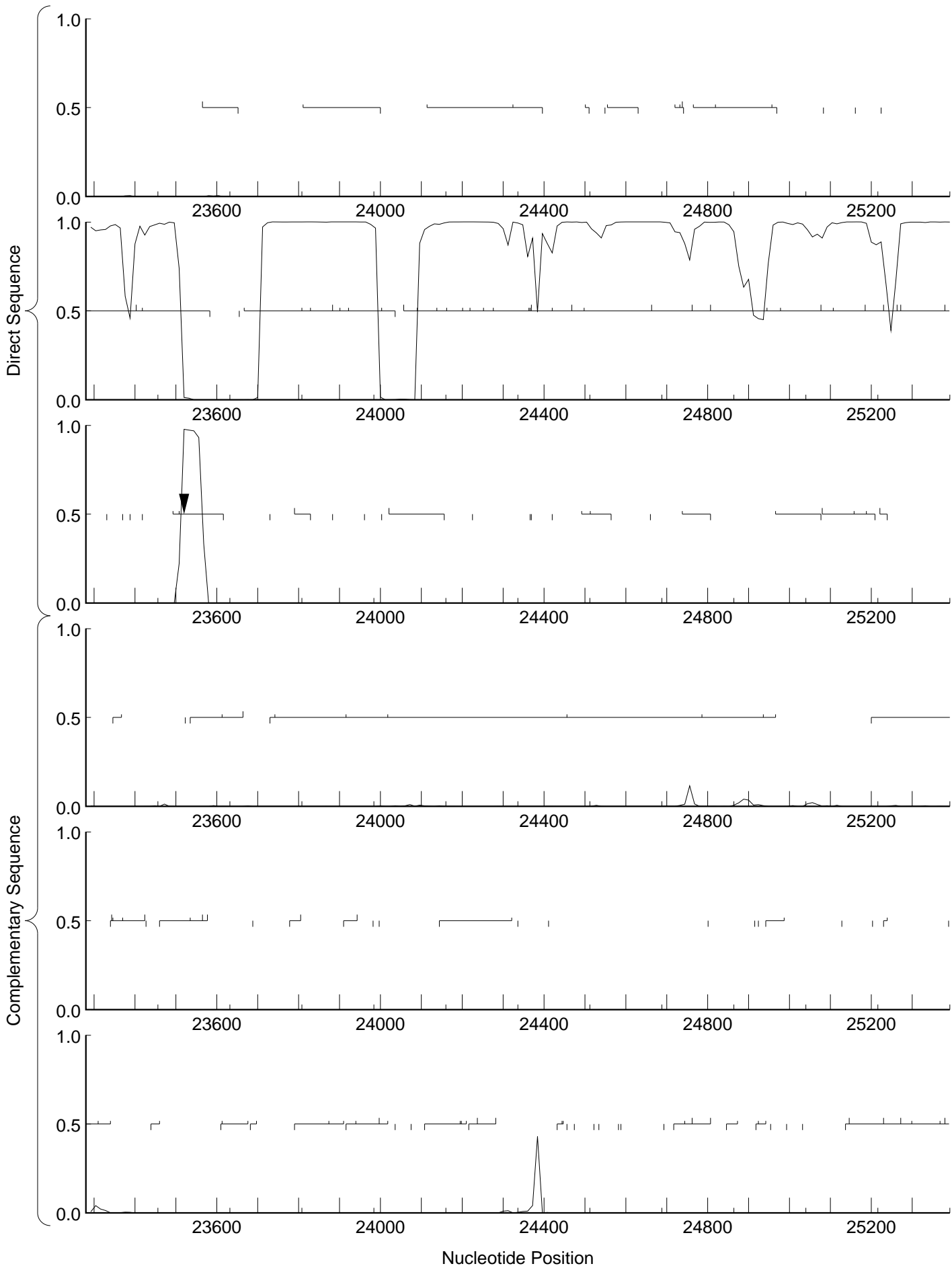


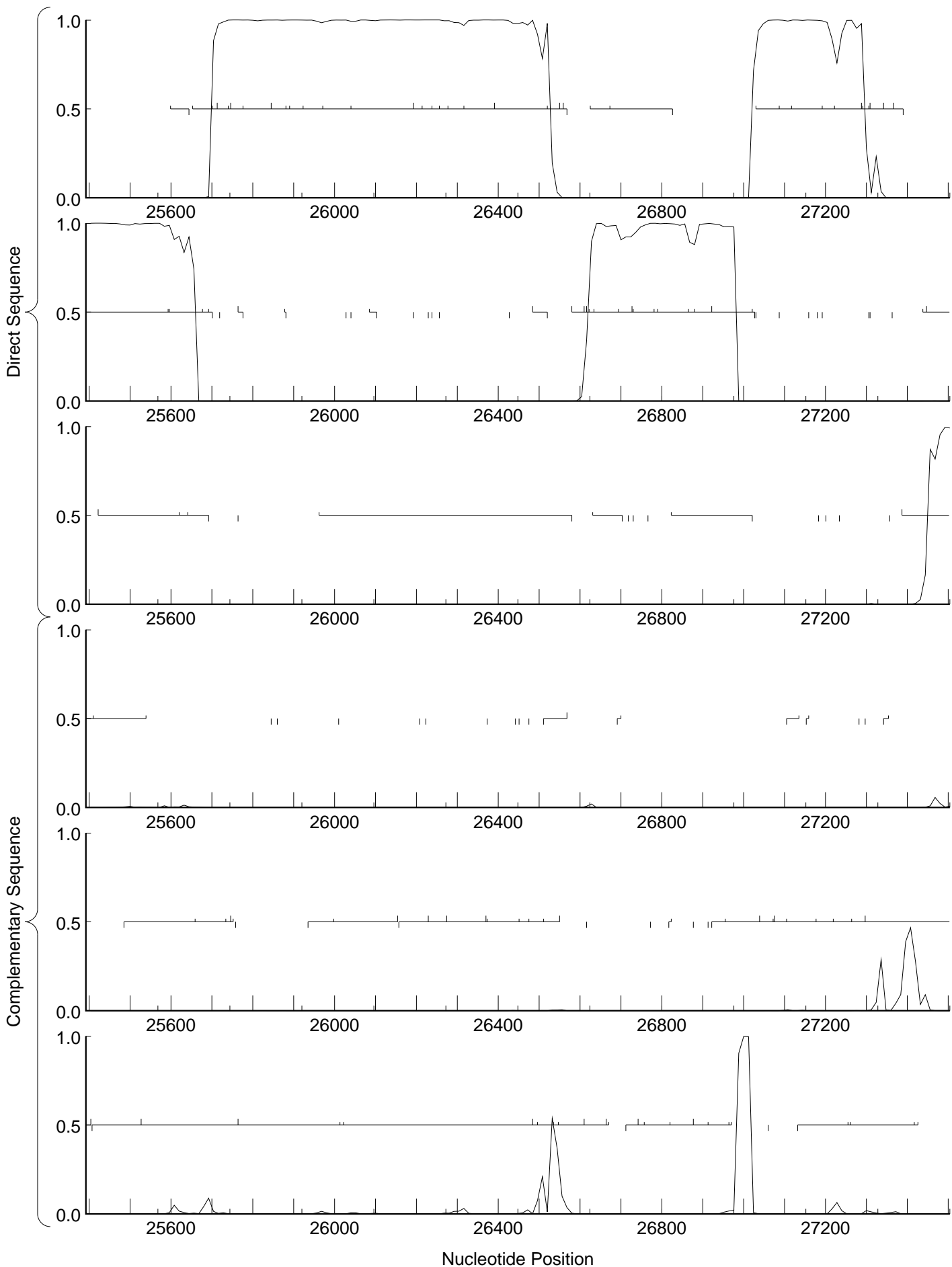


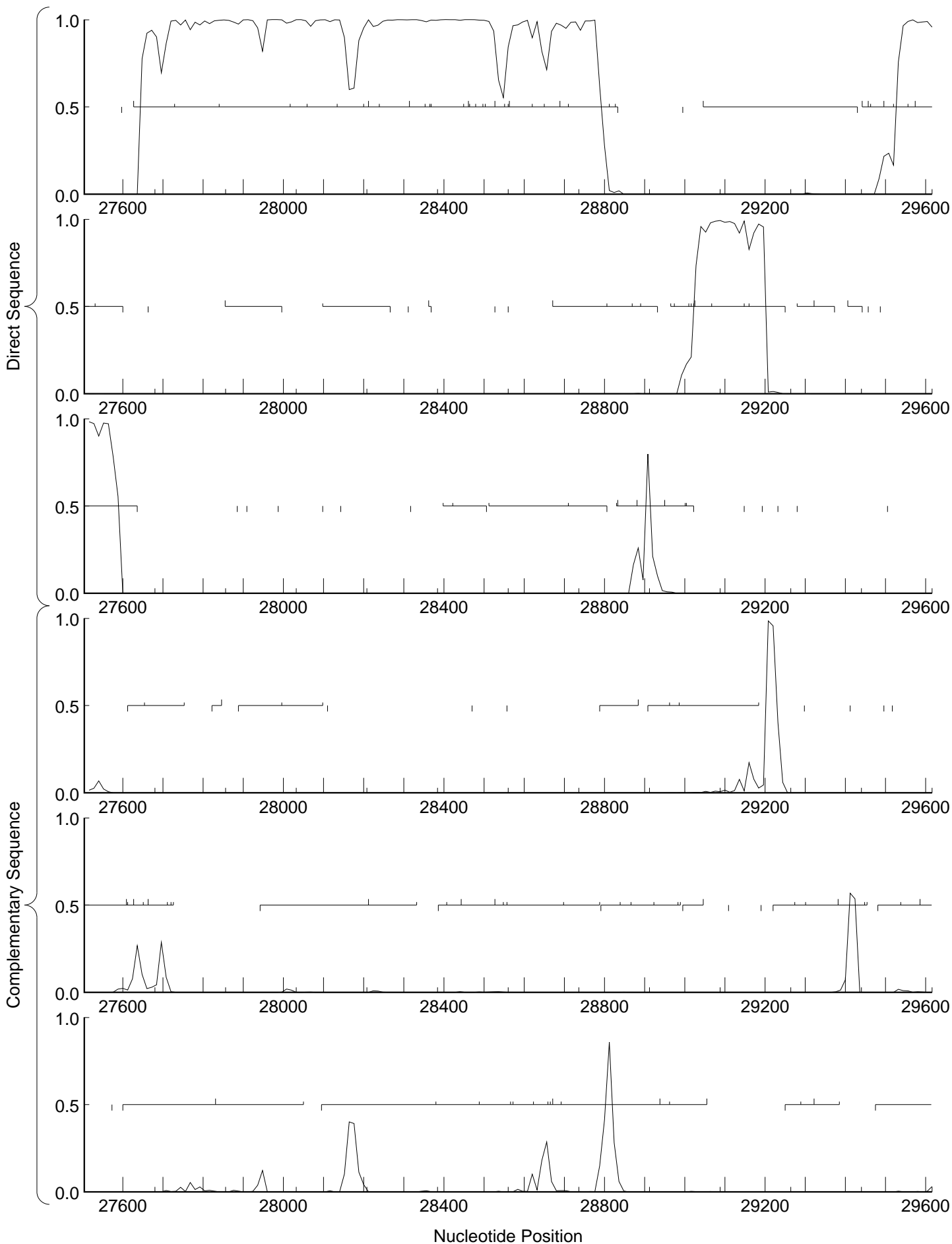


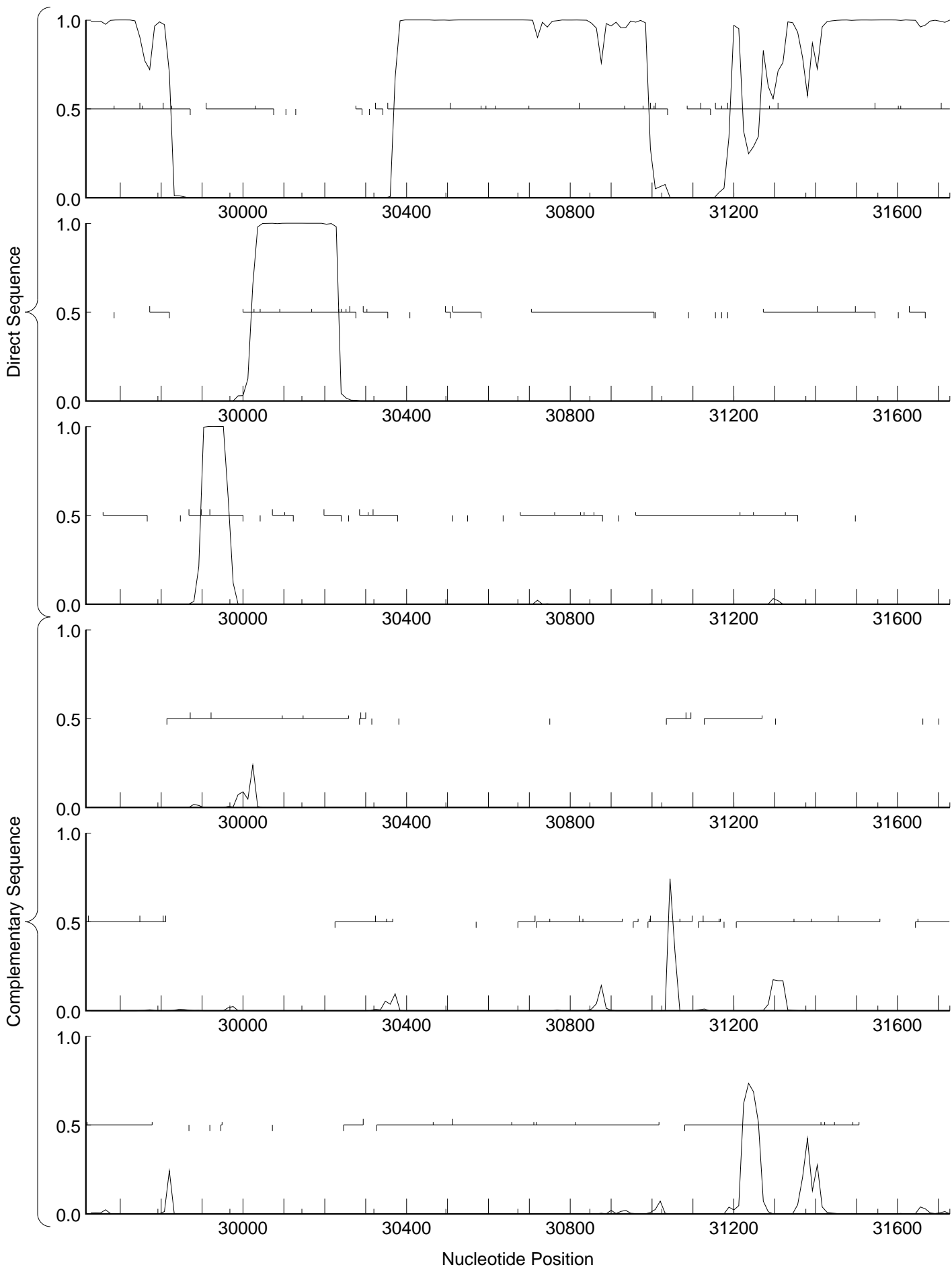


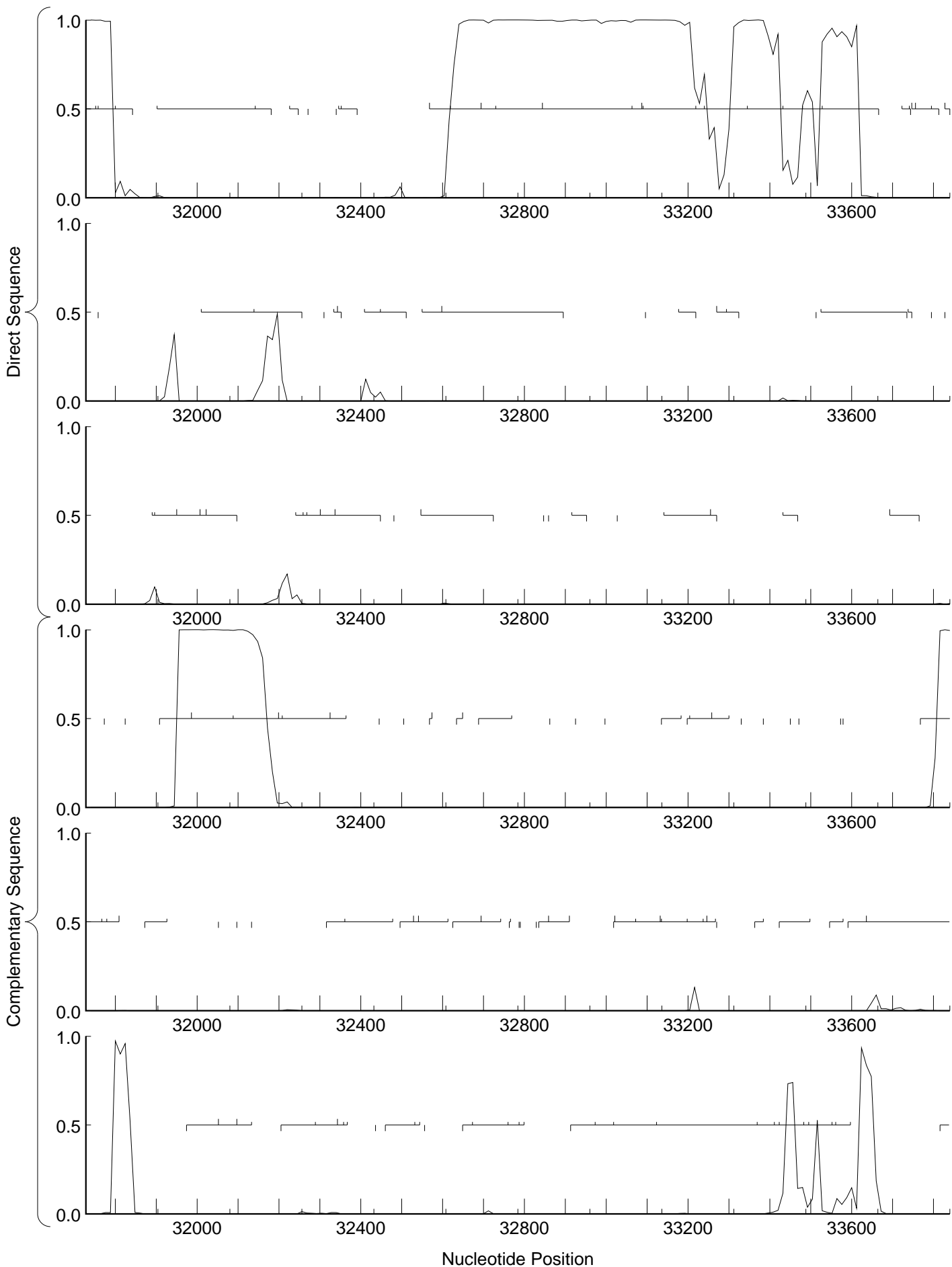


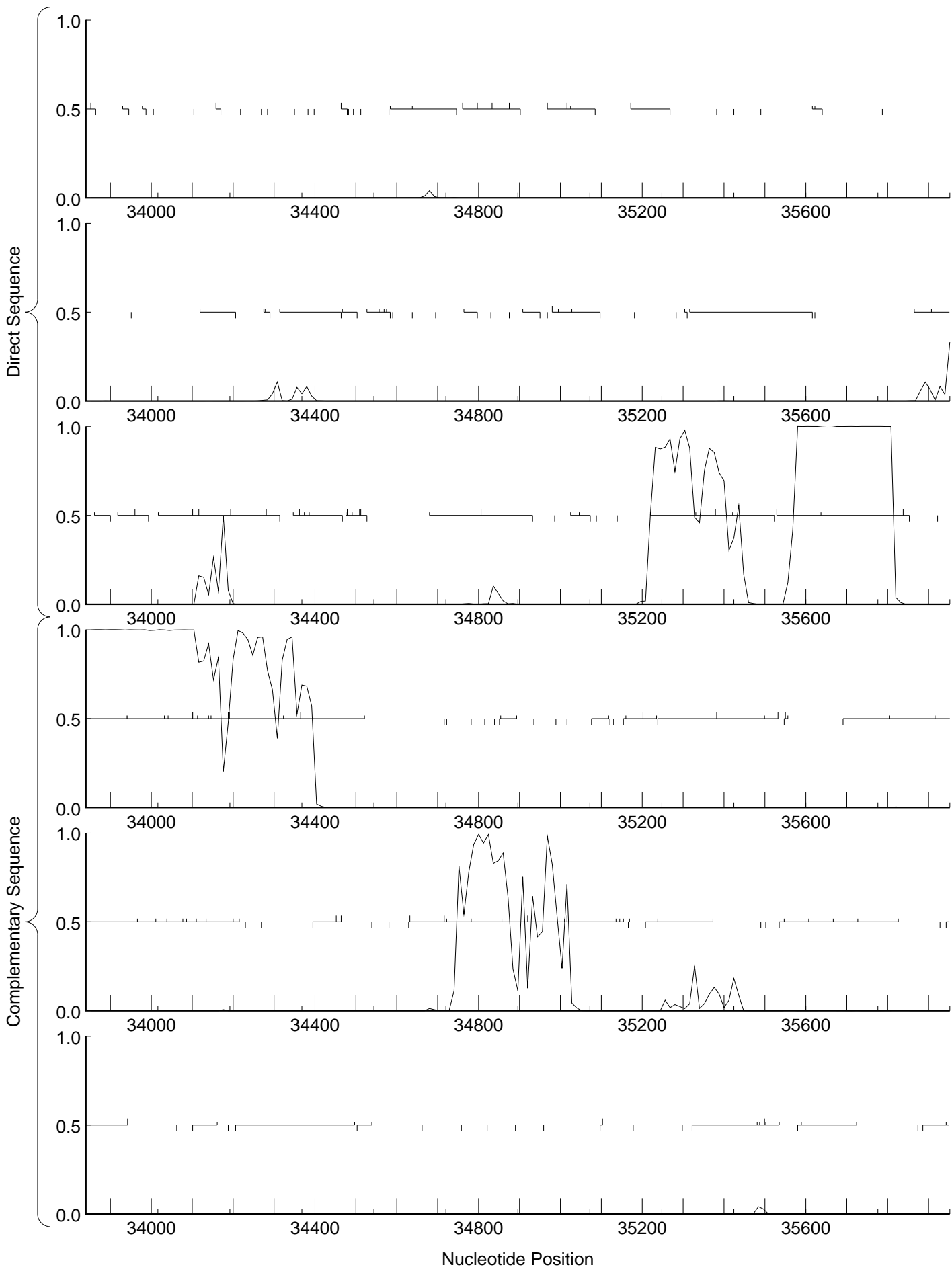


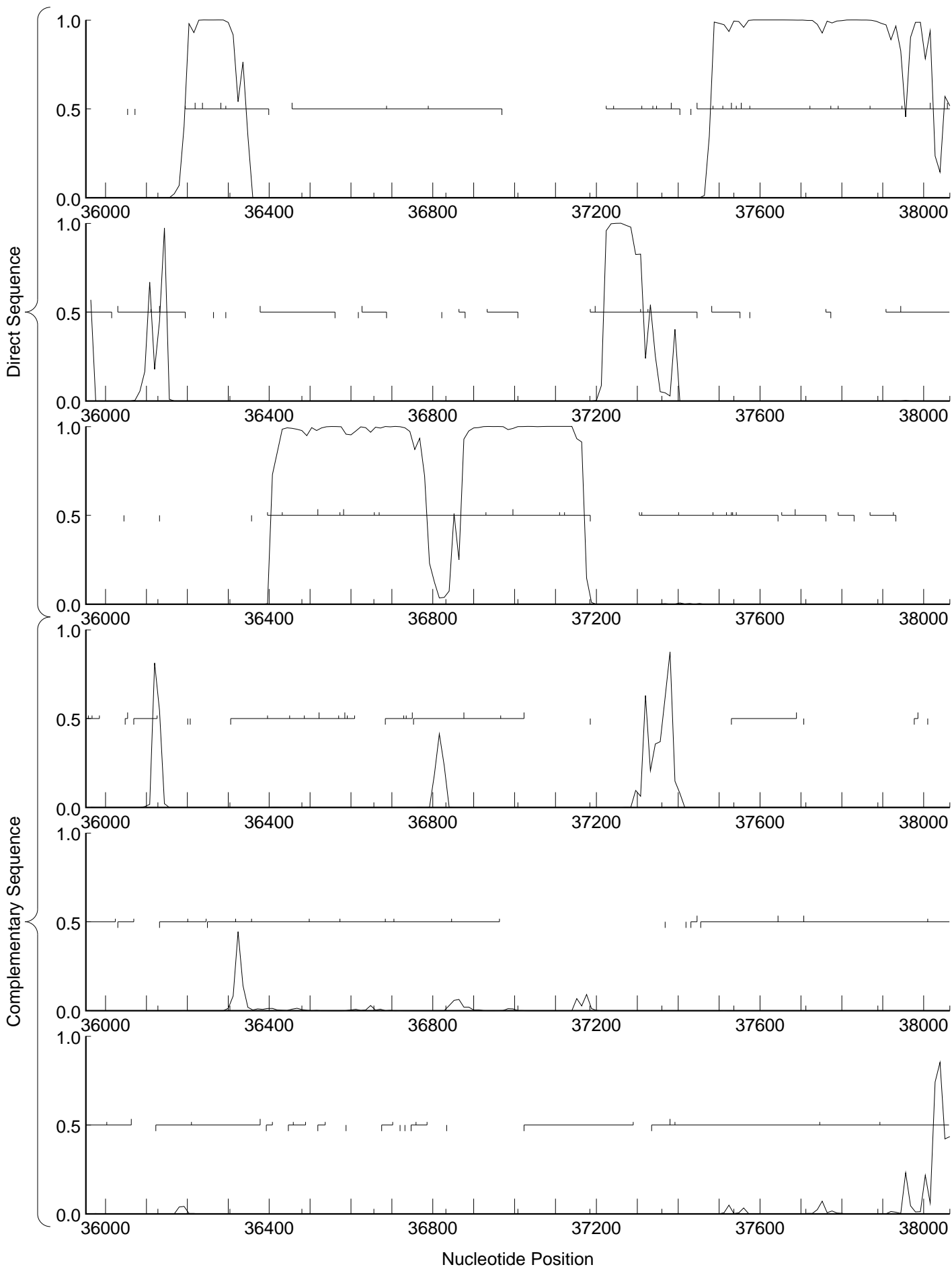


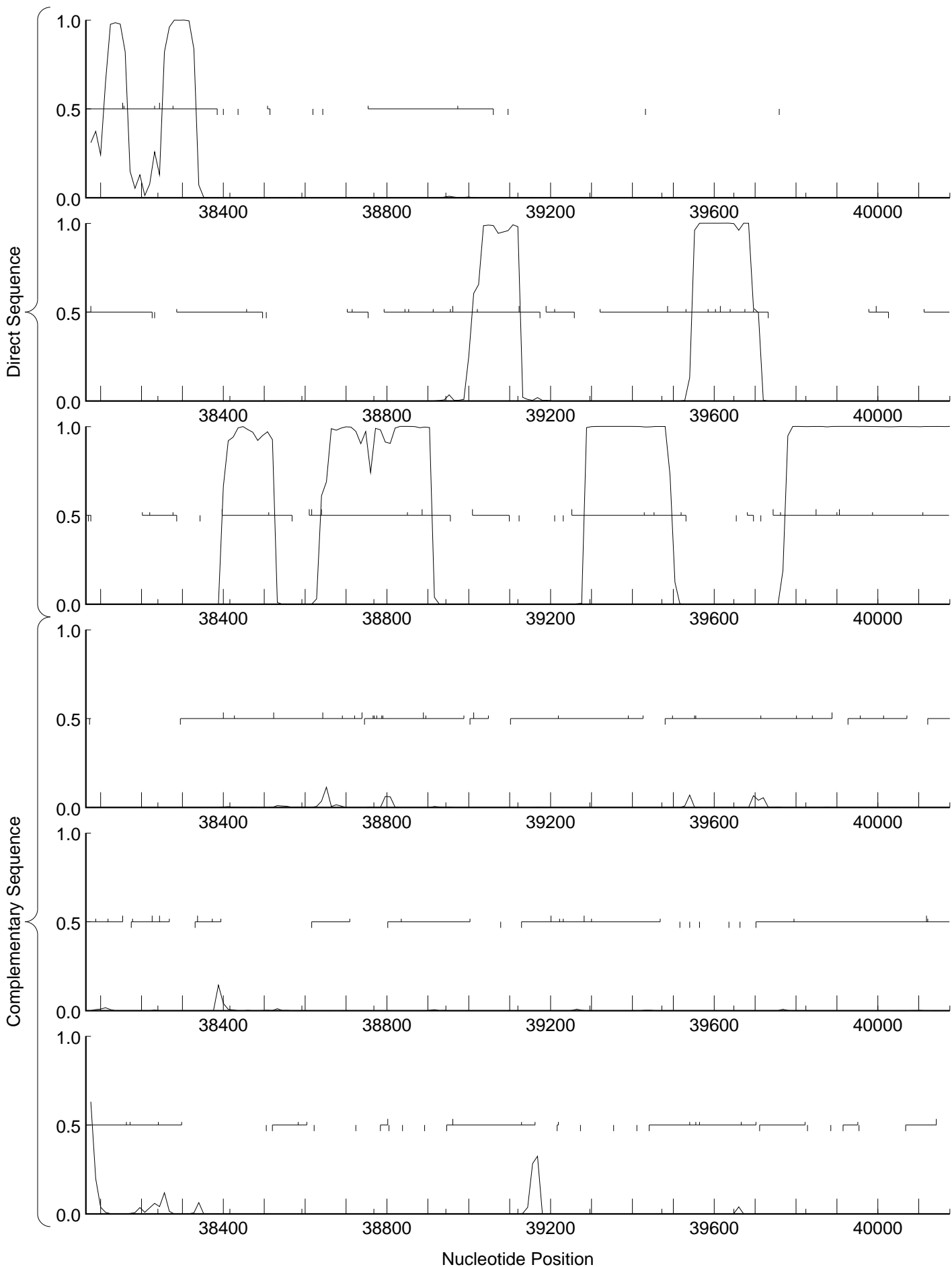












complete sequence, 61284 bp including 11-base 3' overhang (CTCGTAGGCAT), Cluster K1, Order 4, Window 96, Step 12, 21/31

