

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage BENTherdunthat complete sequence, 54867 bp including 10-base 3' overhang (CT
Analysis Date : 1/18/18 at 20:34:19
Pages : 27
Sequence Length : 54867 bp
GC Content : 63.36%

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,

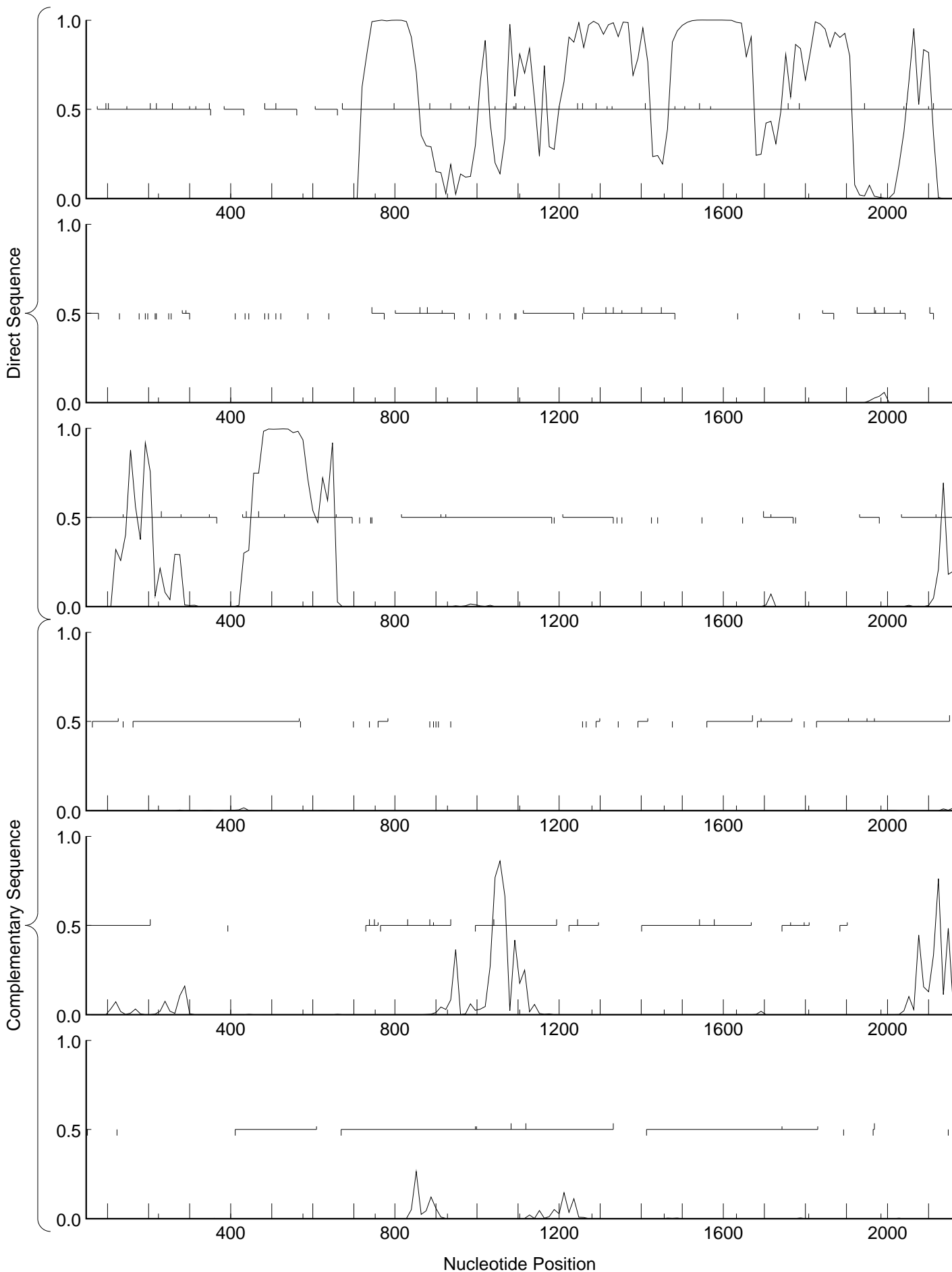
MATRIX INFORMATION

Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

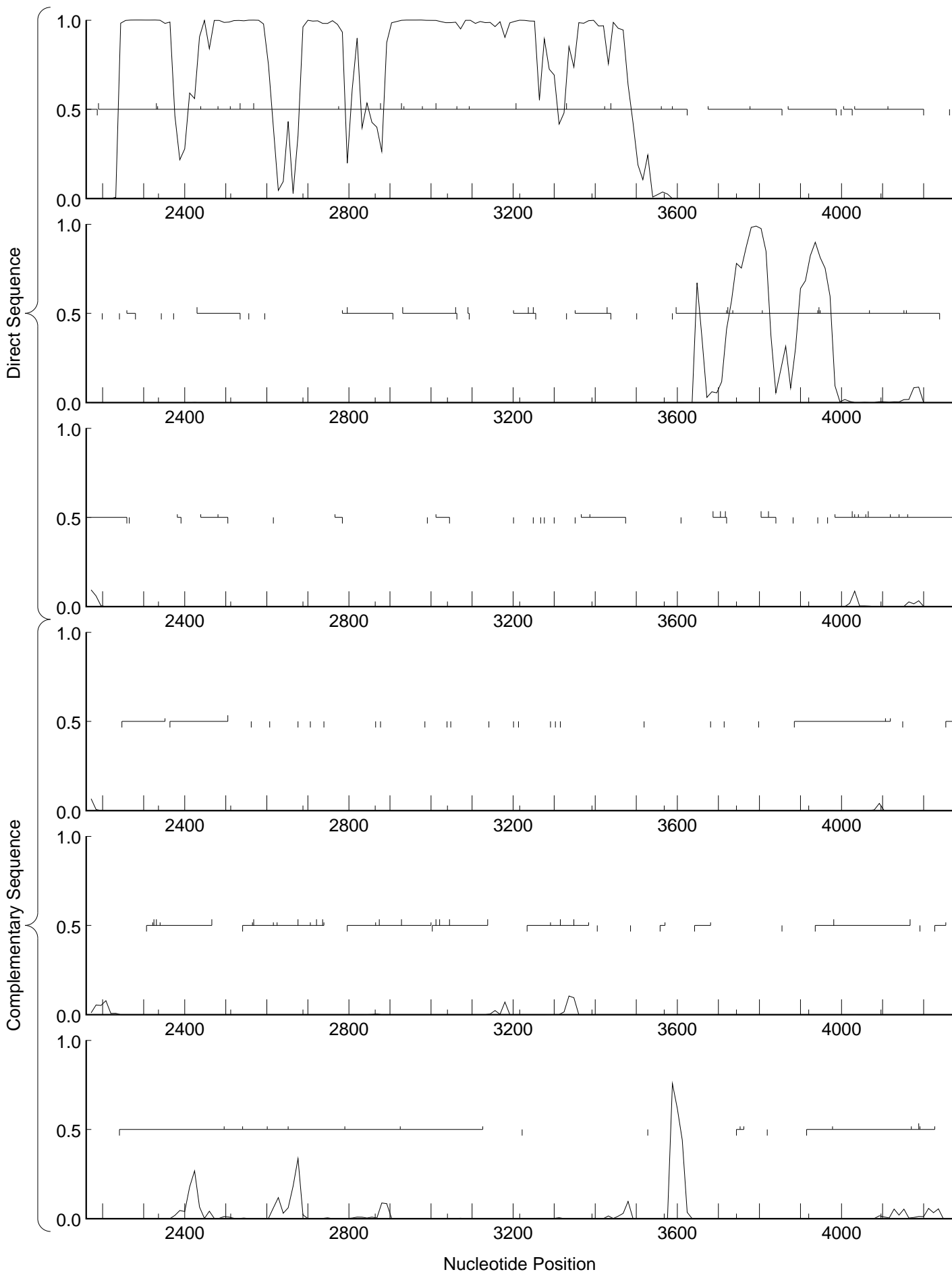
Send questions / comments to:
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Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

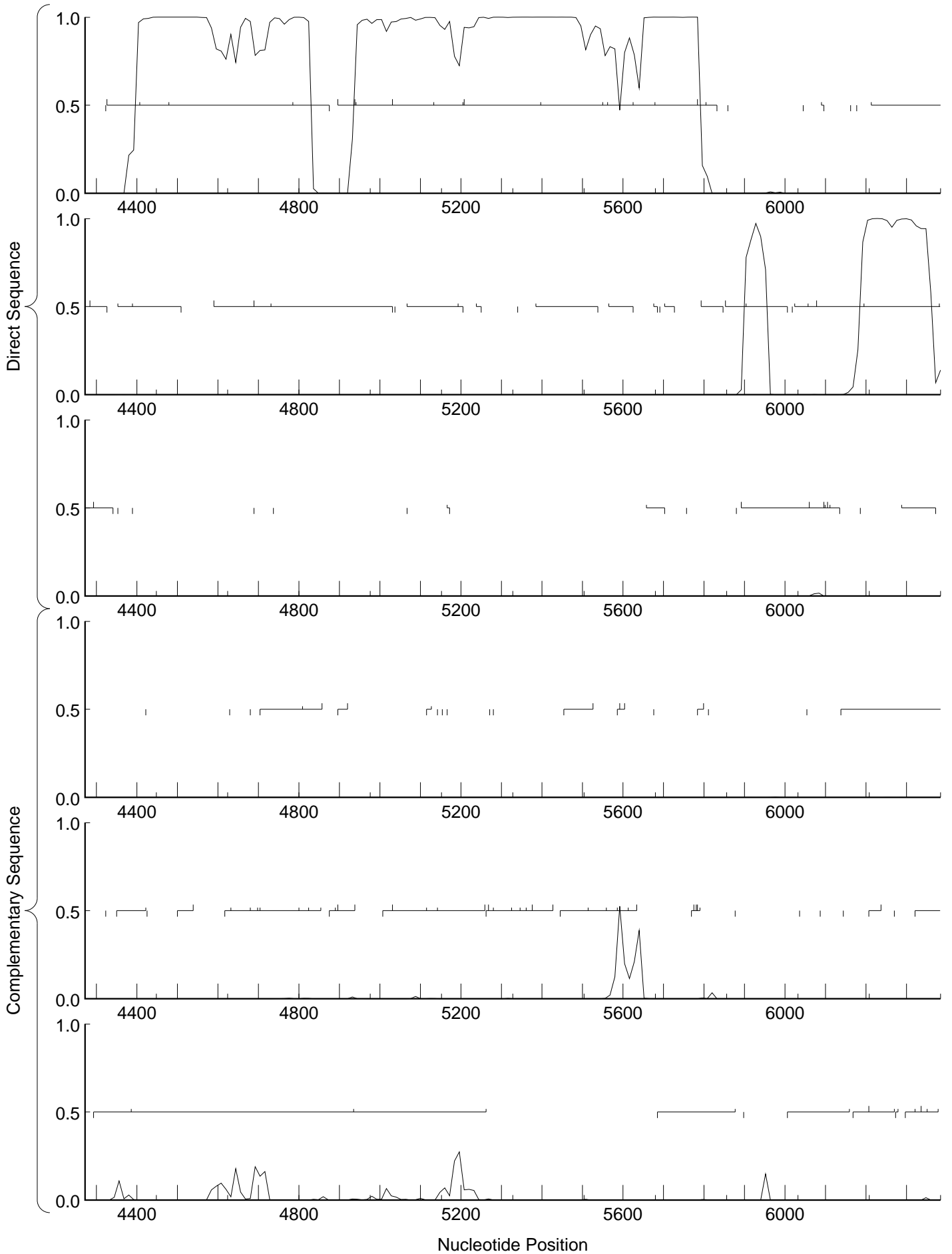
Matrix notes & comments

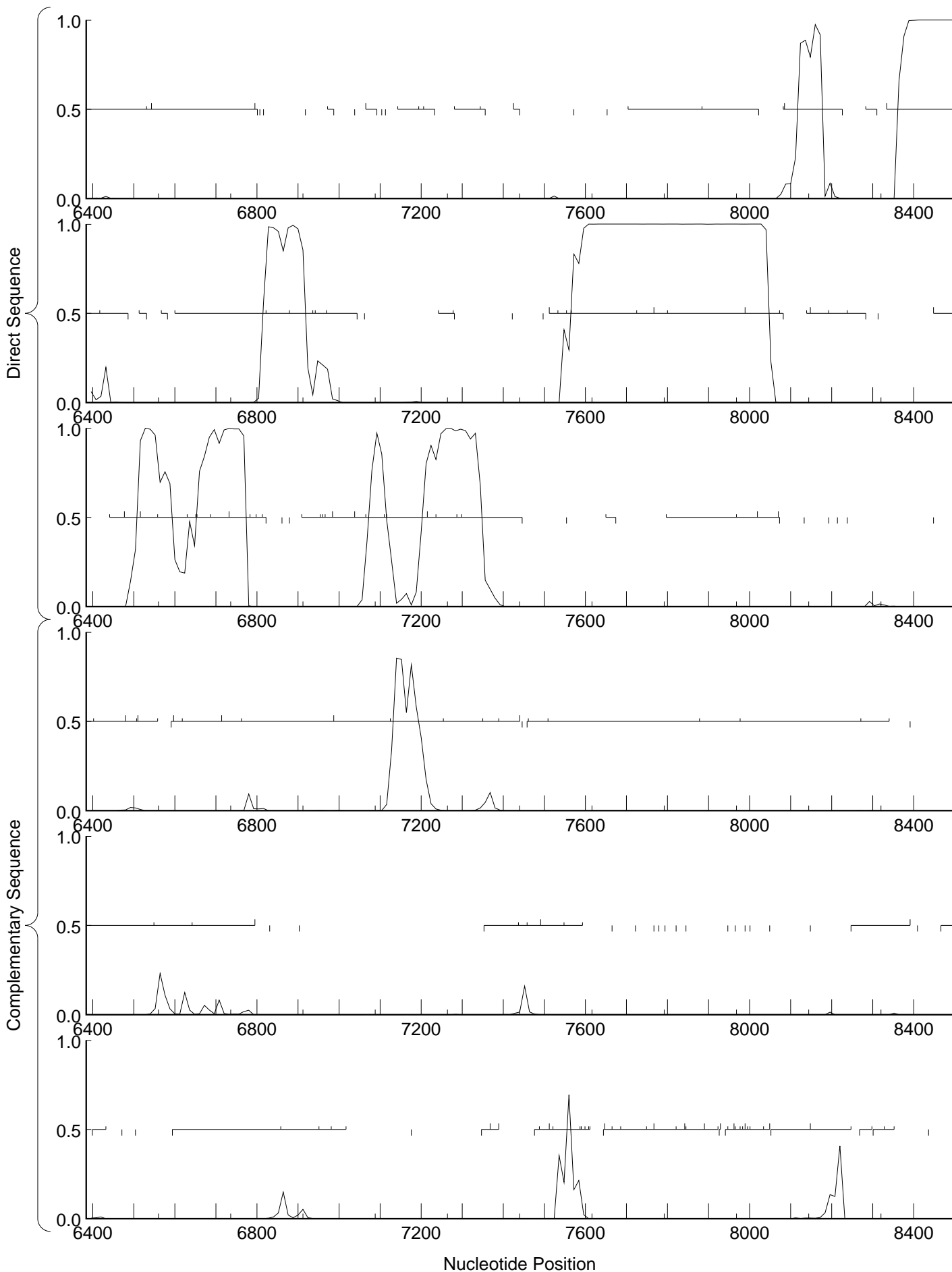
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

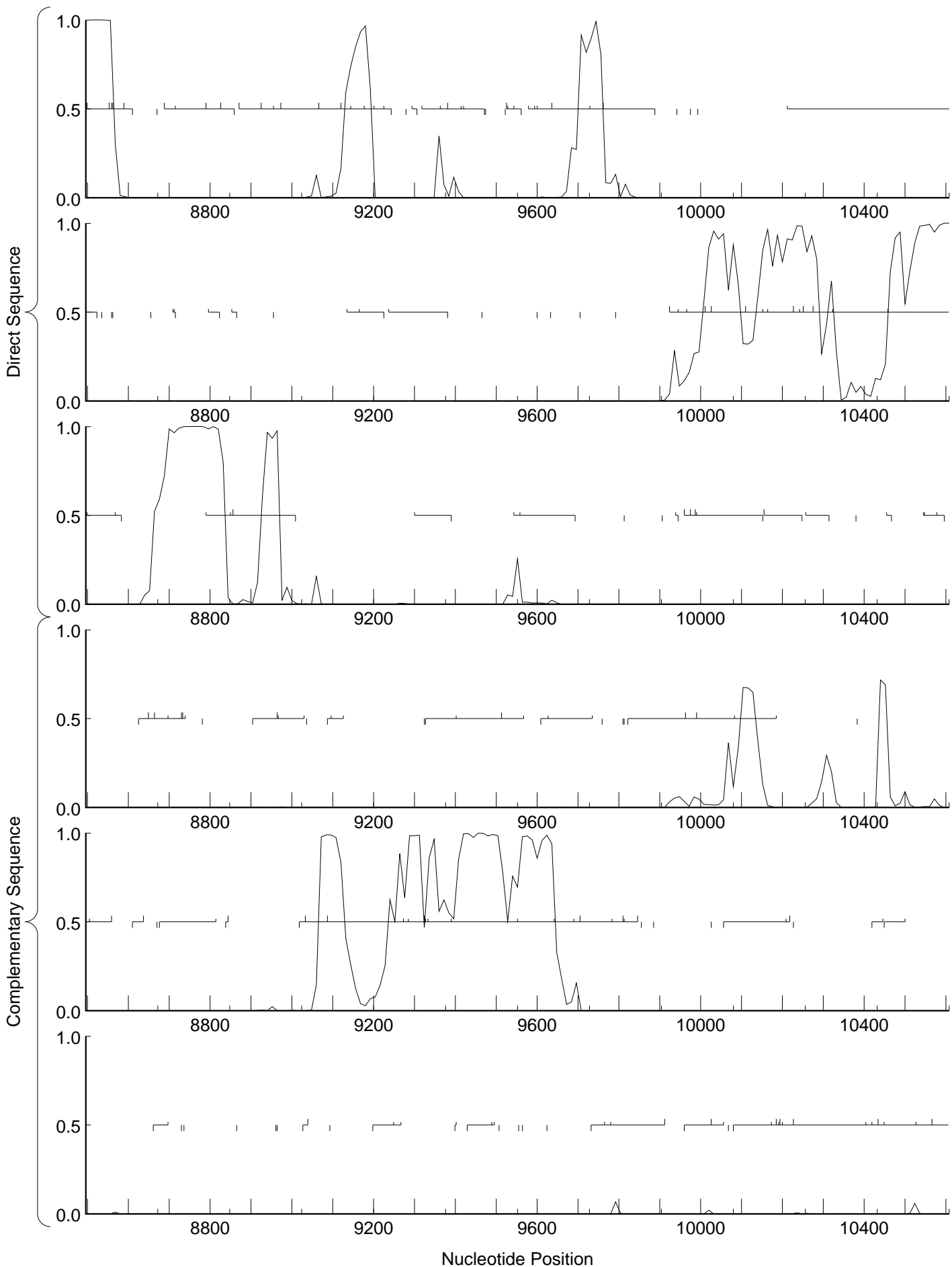


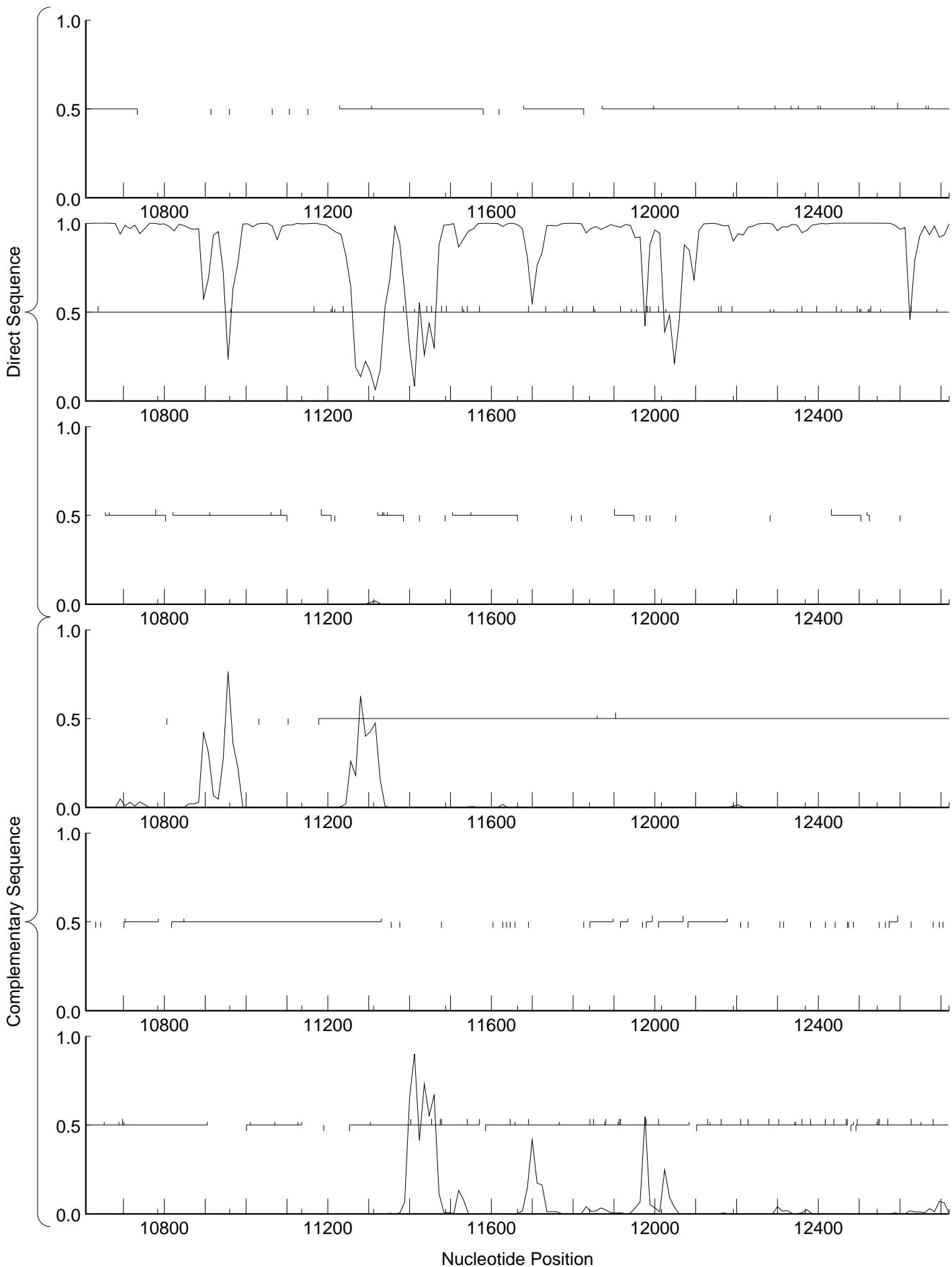
Complete sequence, 54867 bp including 10-base 3' overhang (CTCGGGGCAT), Cluster DN, Order 4, Window 96, Step 12, 3/27

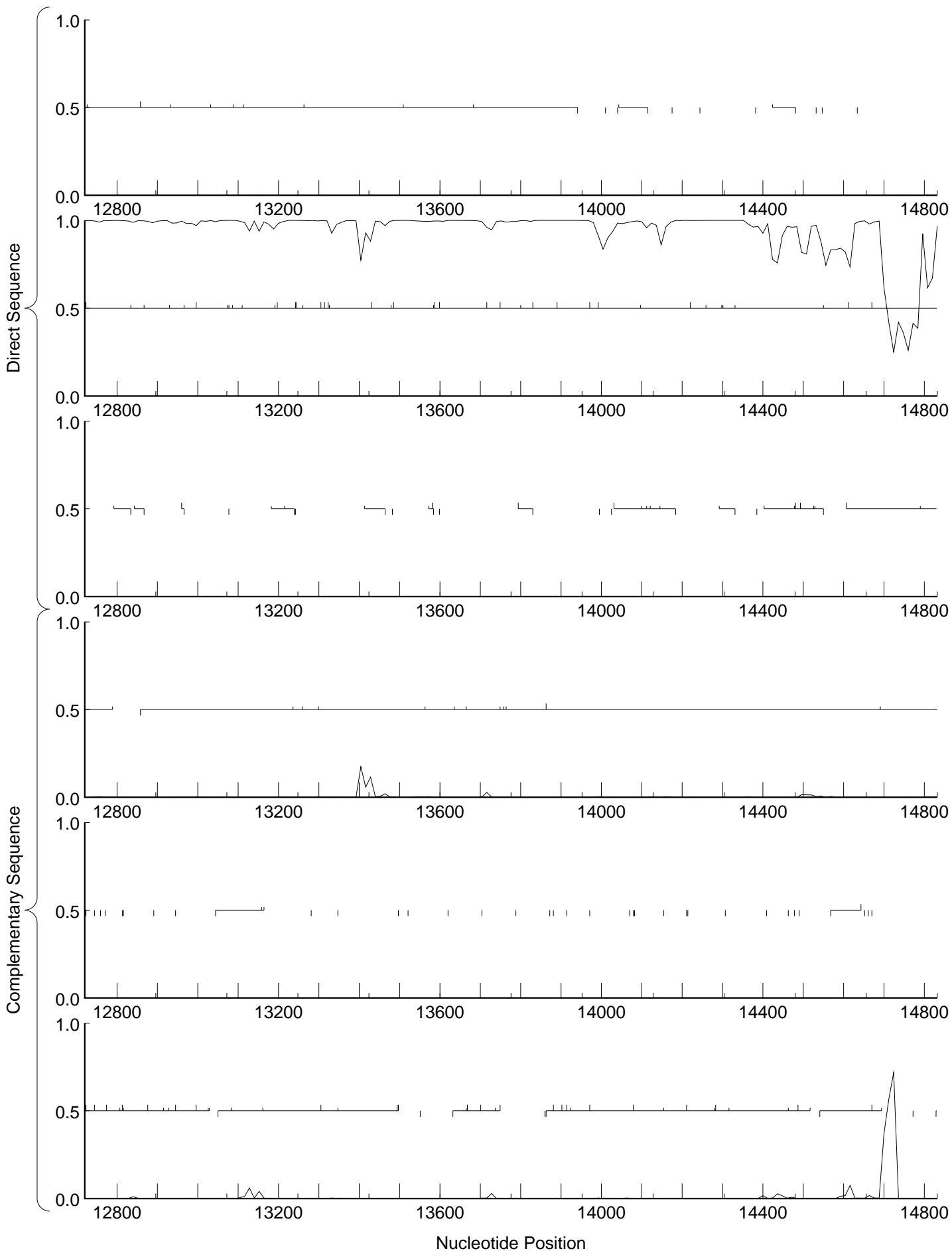


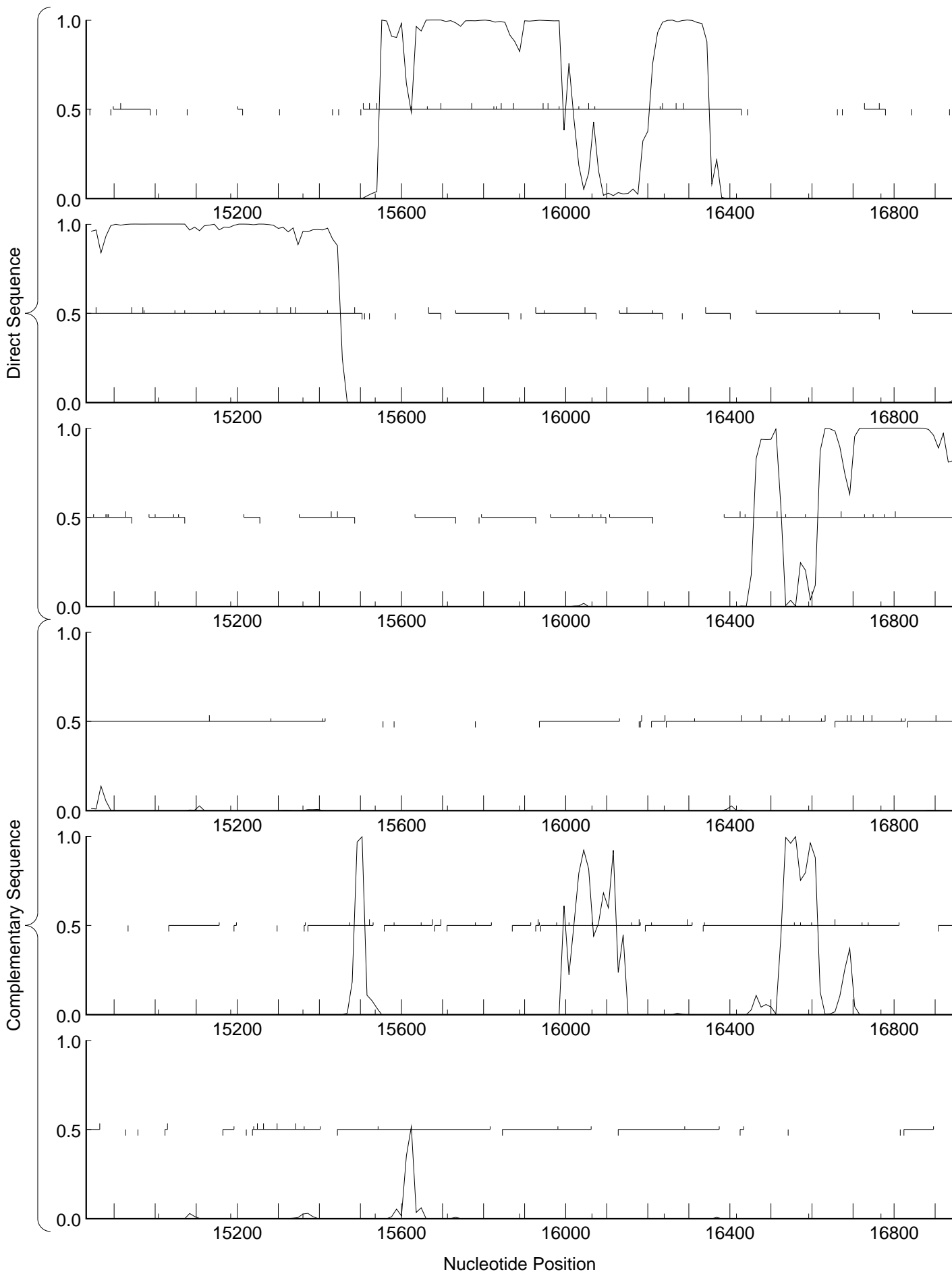


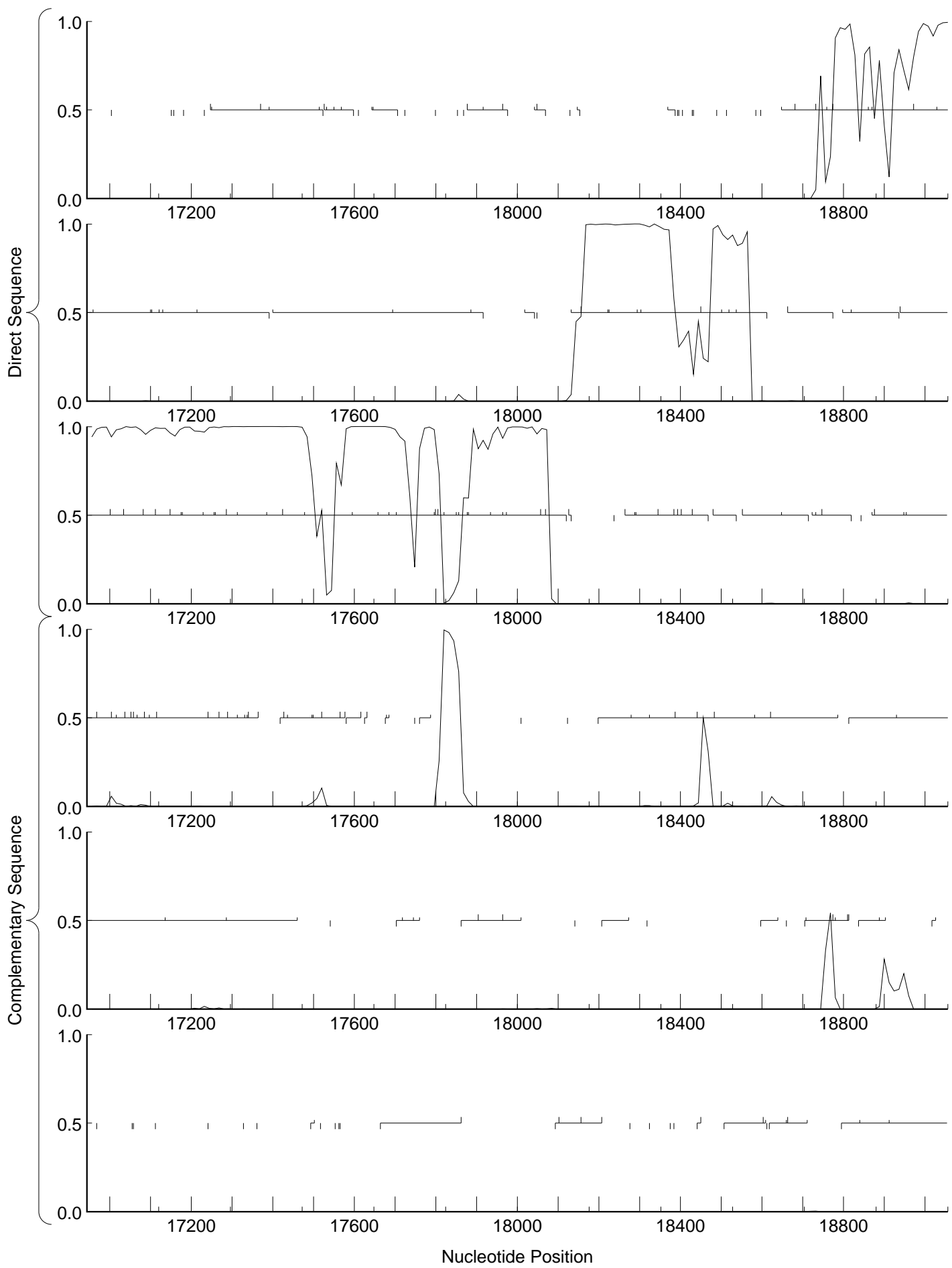


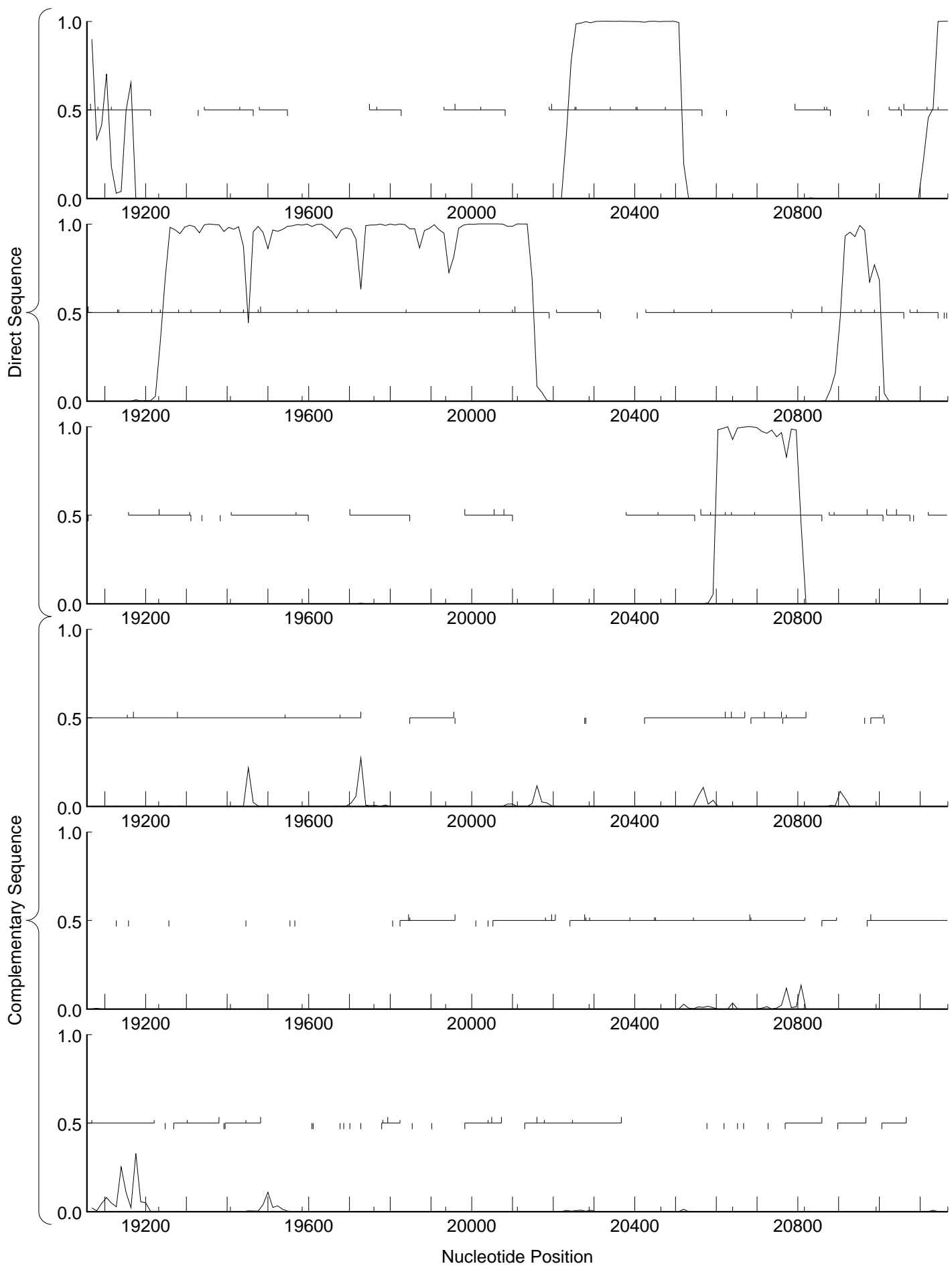


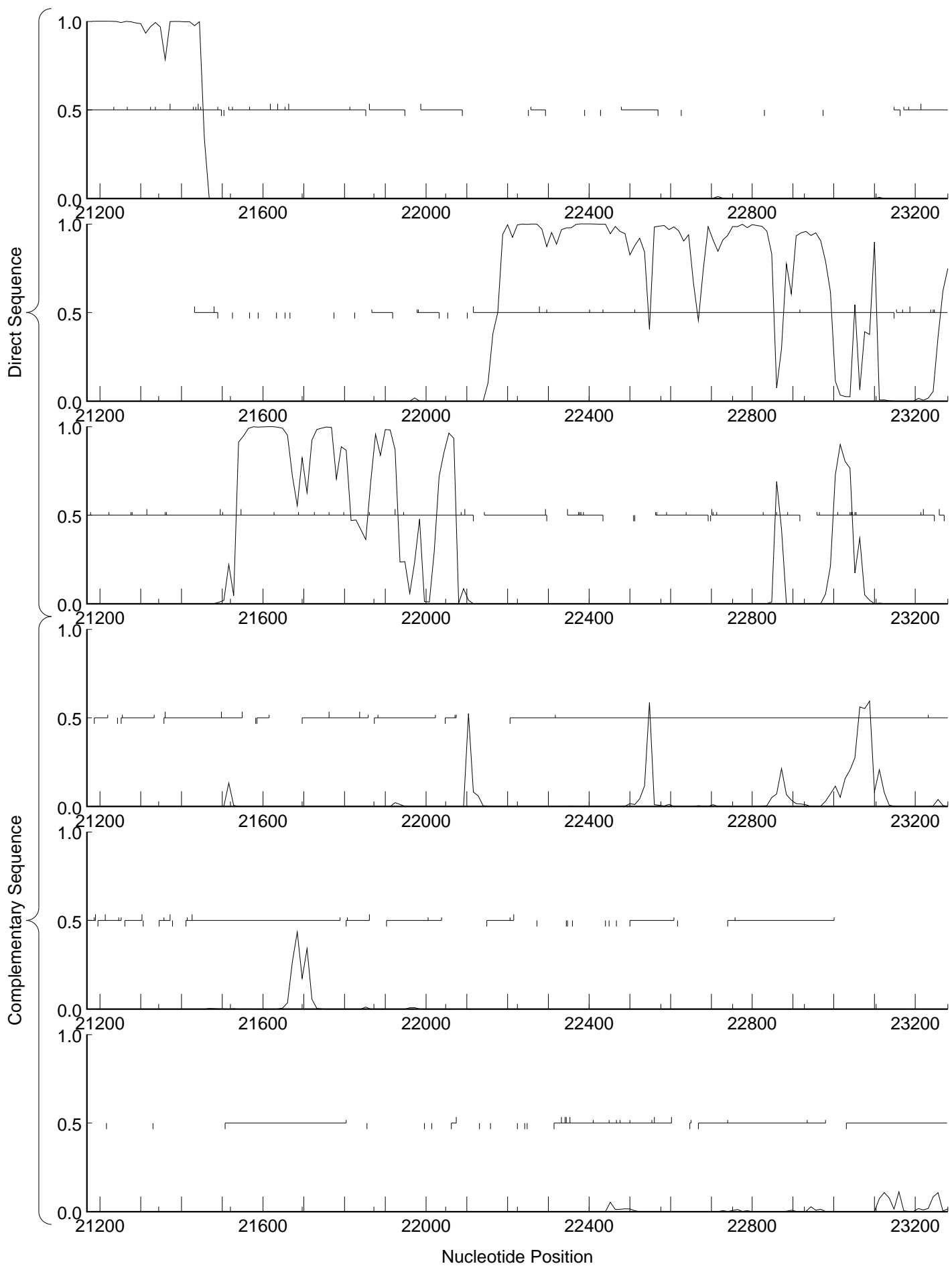


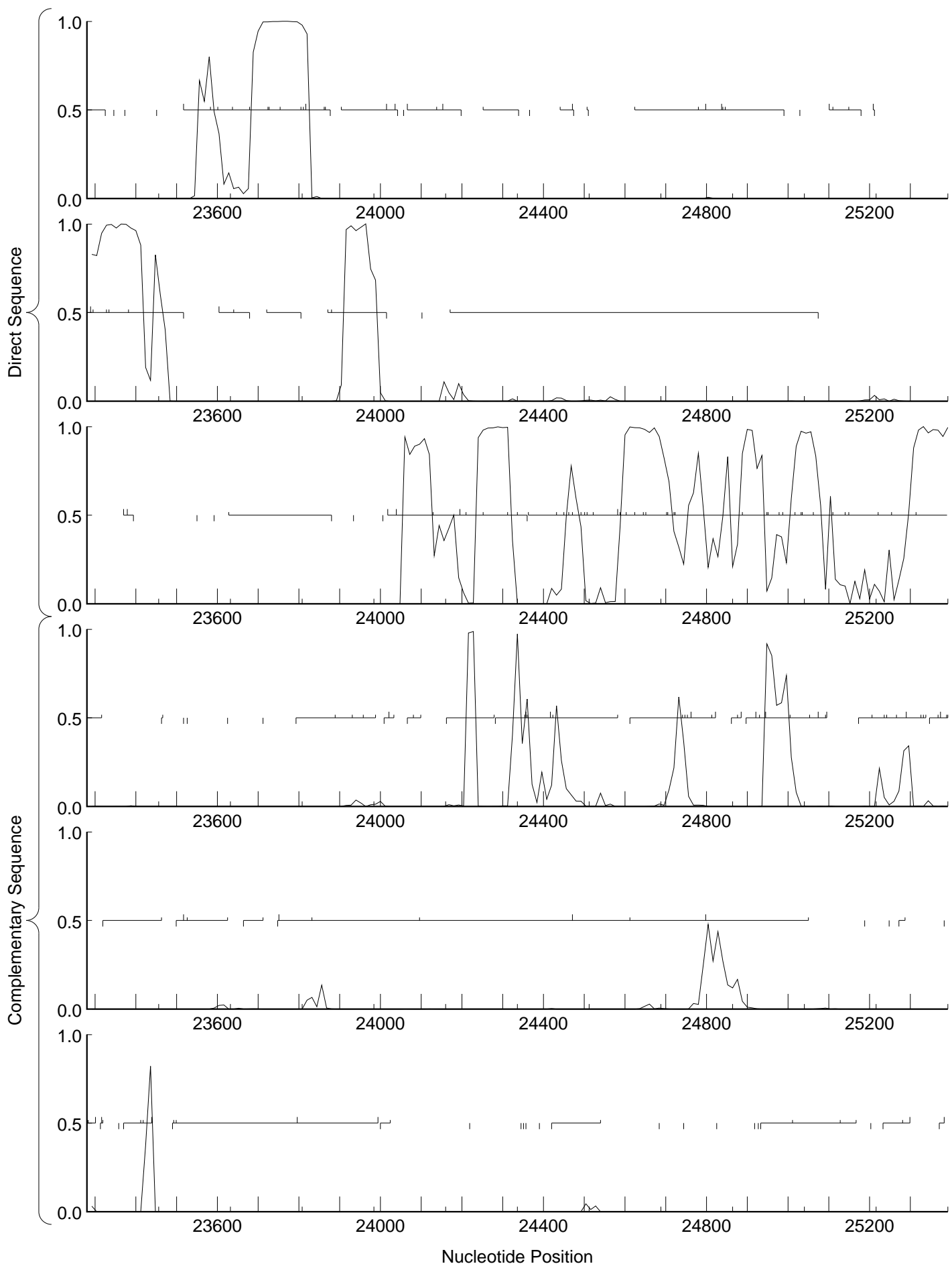


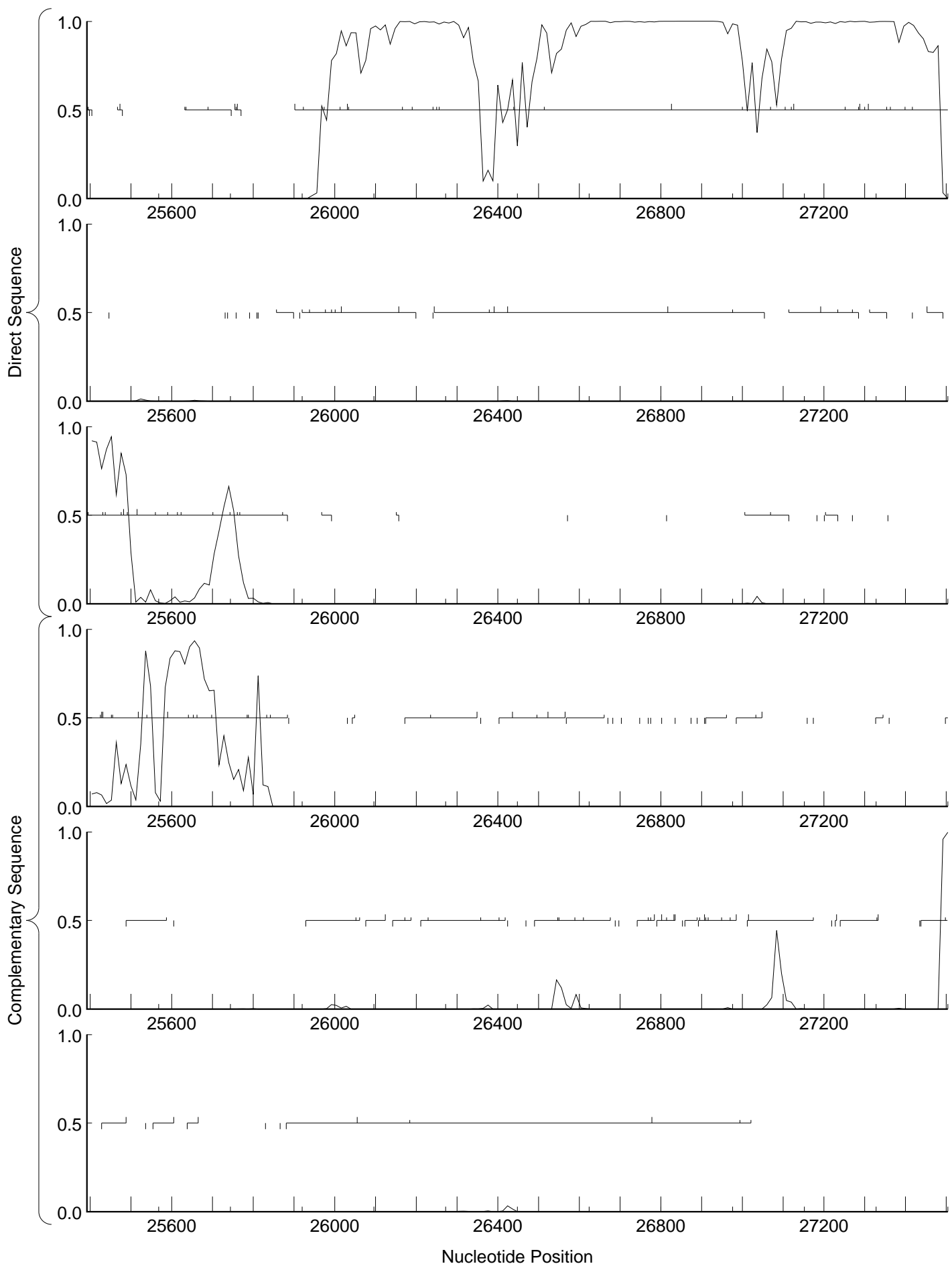


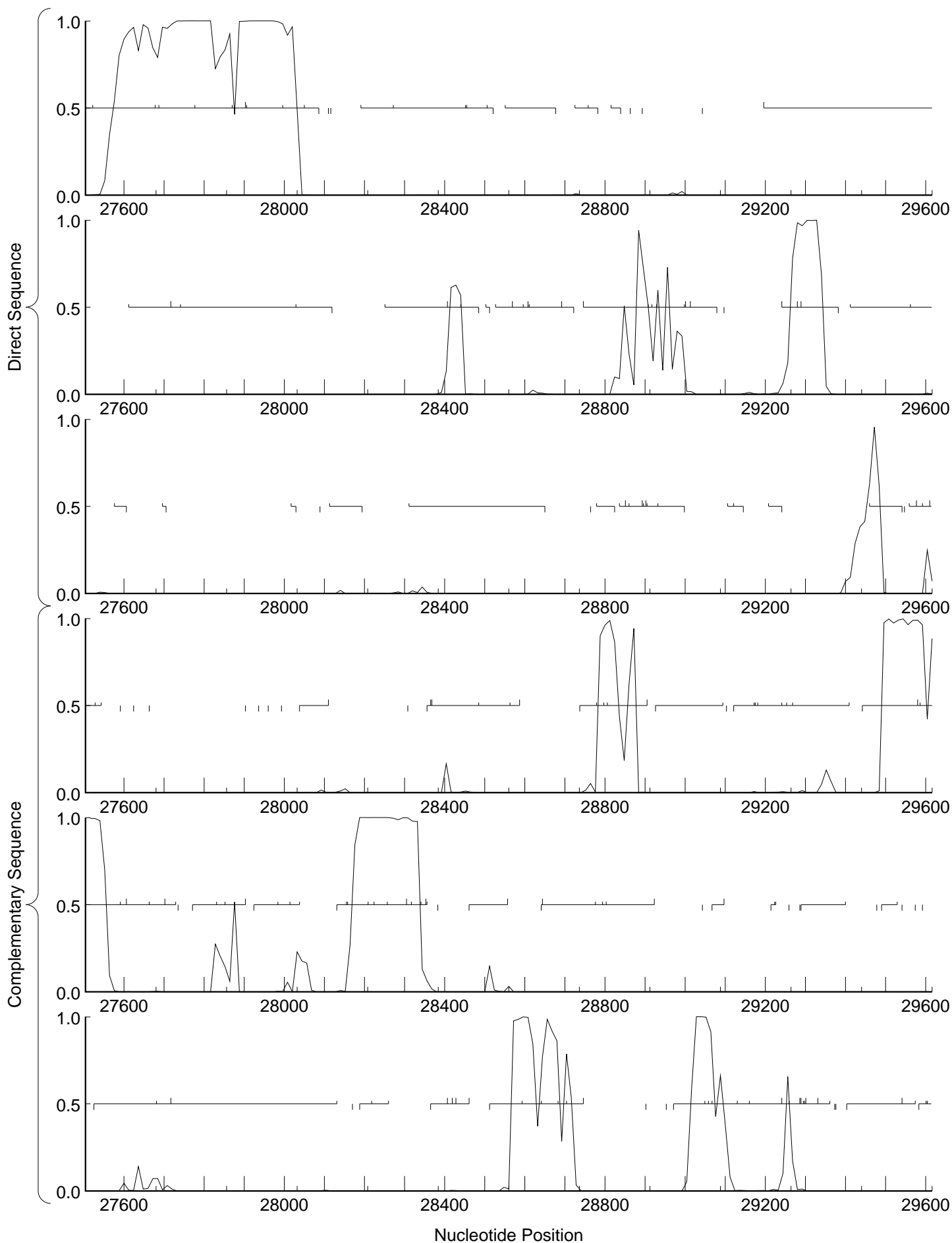












complete sequence, 54867 bp including 10-base 3' overhang (CTCGGGGCAT), Cluster DN, Order 4, Window 96, Step 12, 16/27

