

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage BodEinwohner17 complete sequence, 60222 bp including 10-base 3' overhang
Analysis Date : 5/24/18 at 4:05:45
Pages : 30
Sequence Length : 60222 bp
GC Content : 61.34%

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,

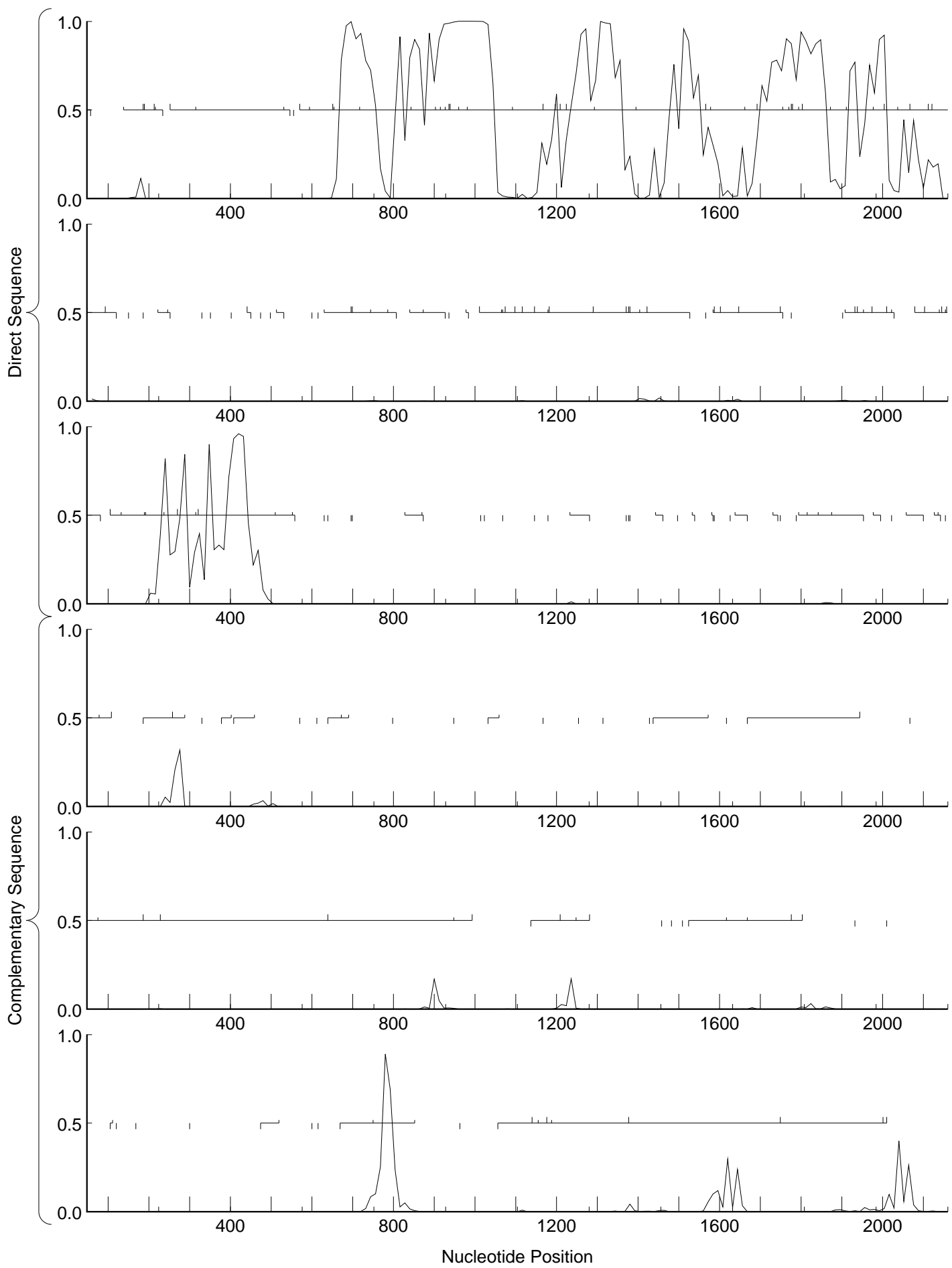
Matrix notes & comments

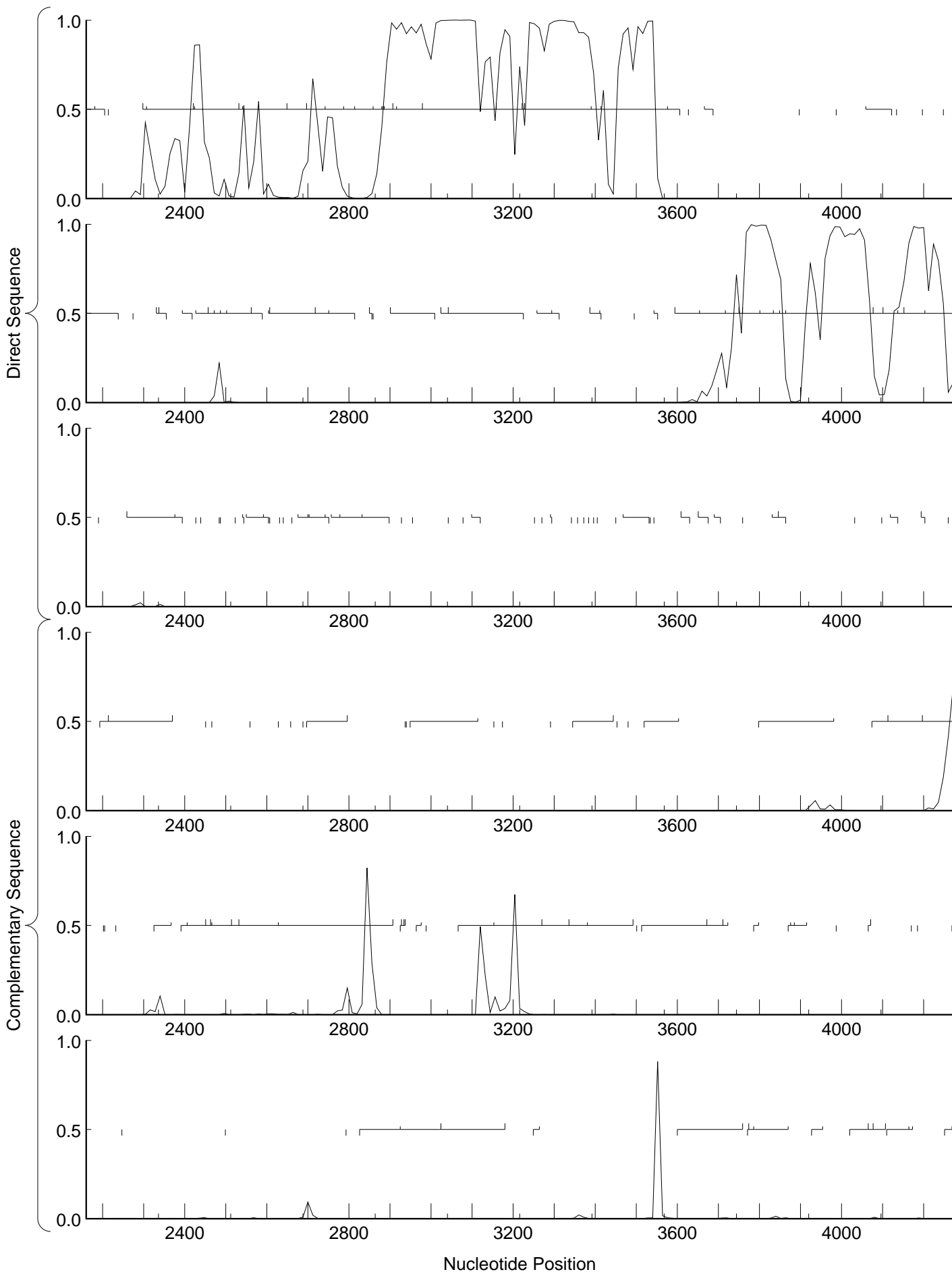
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

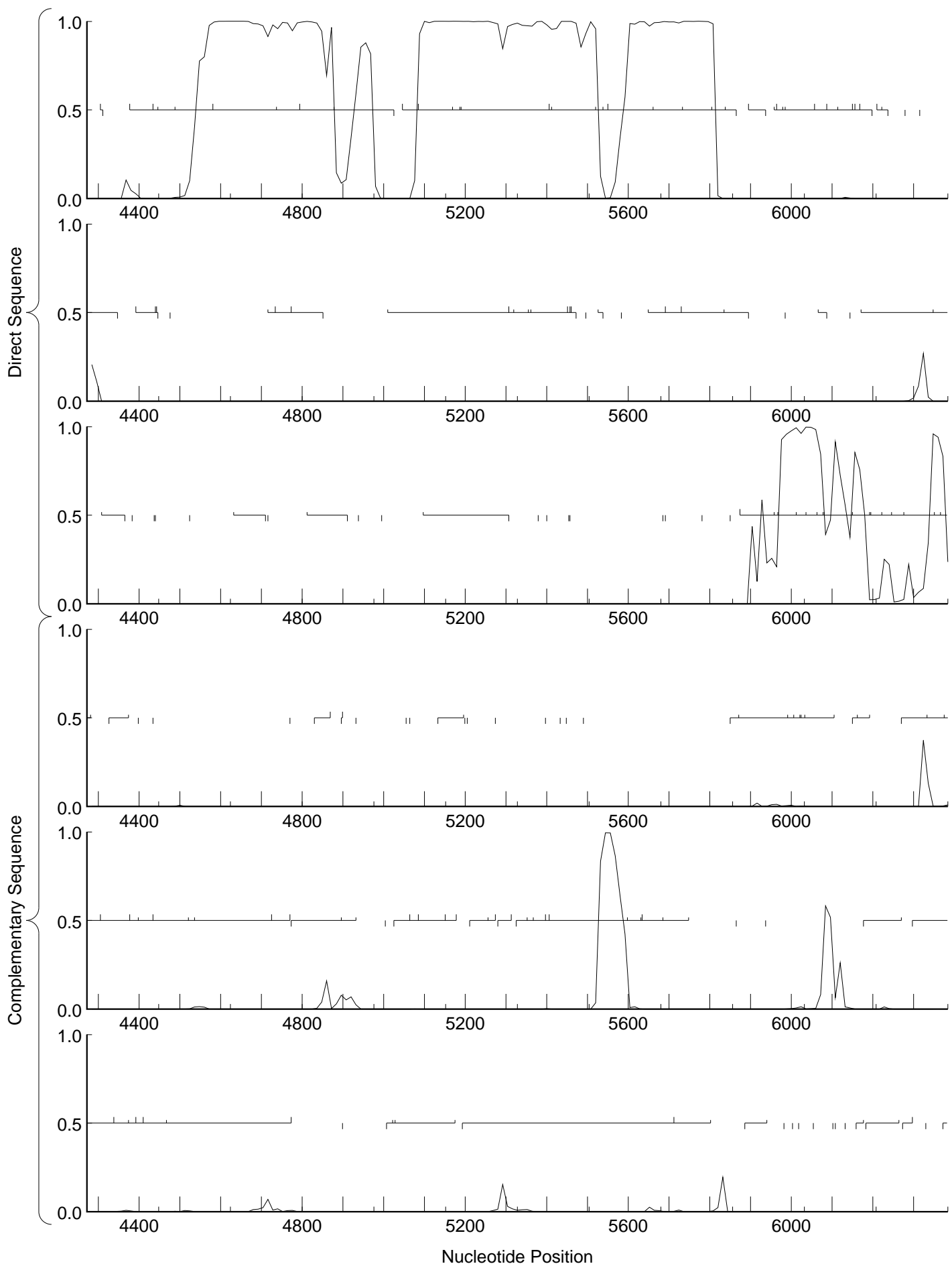
MATRIX INFORMATION

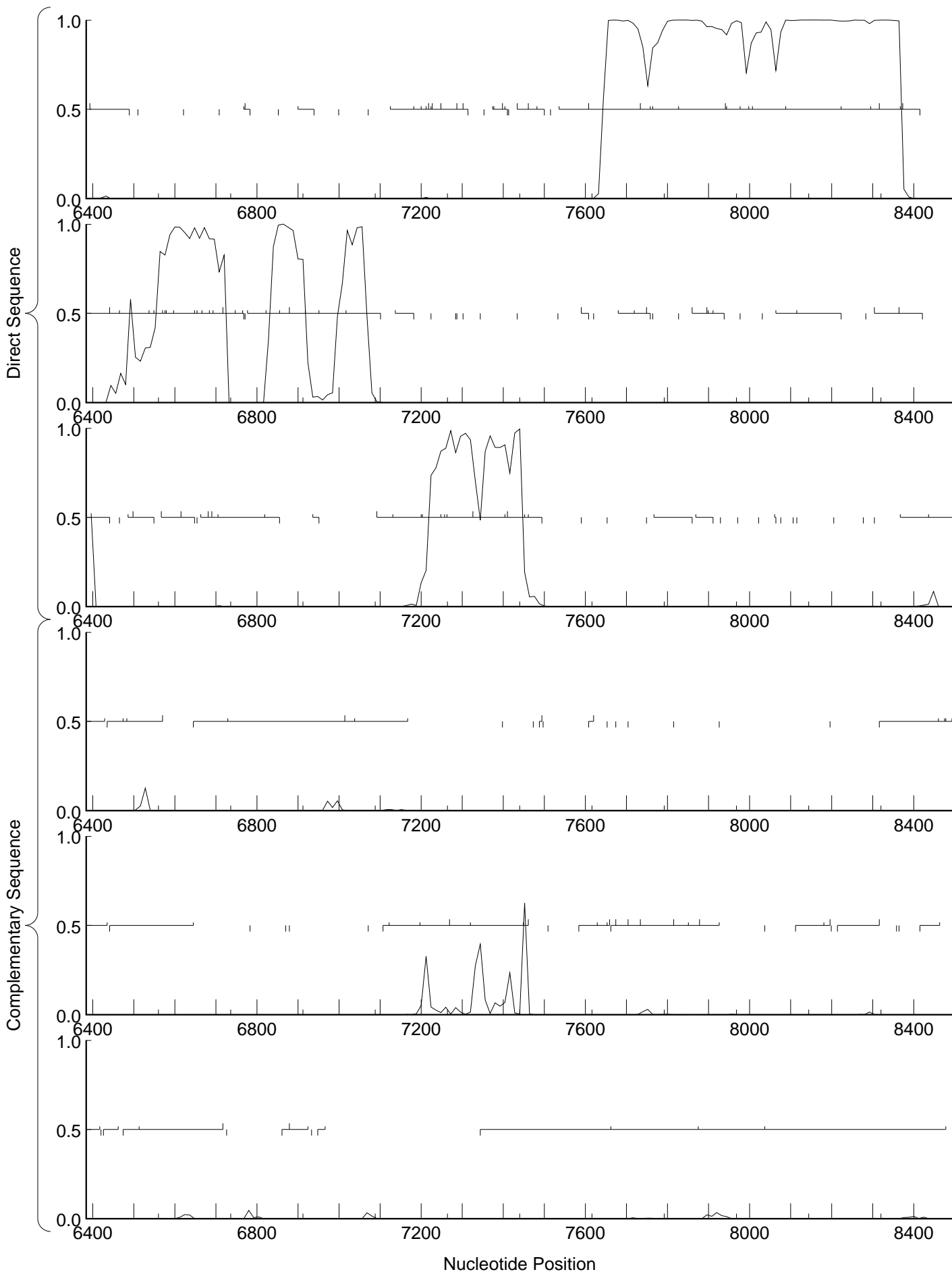
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

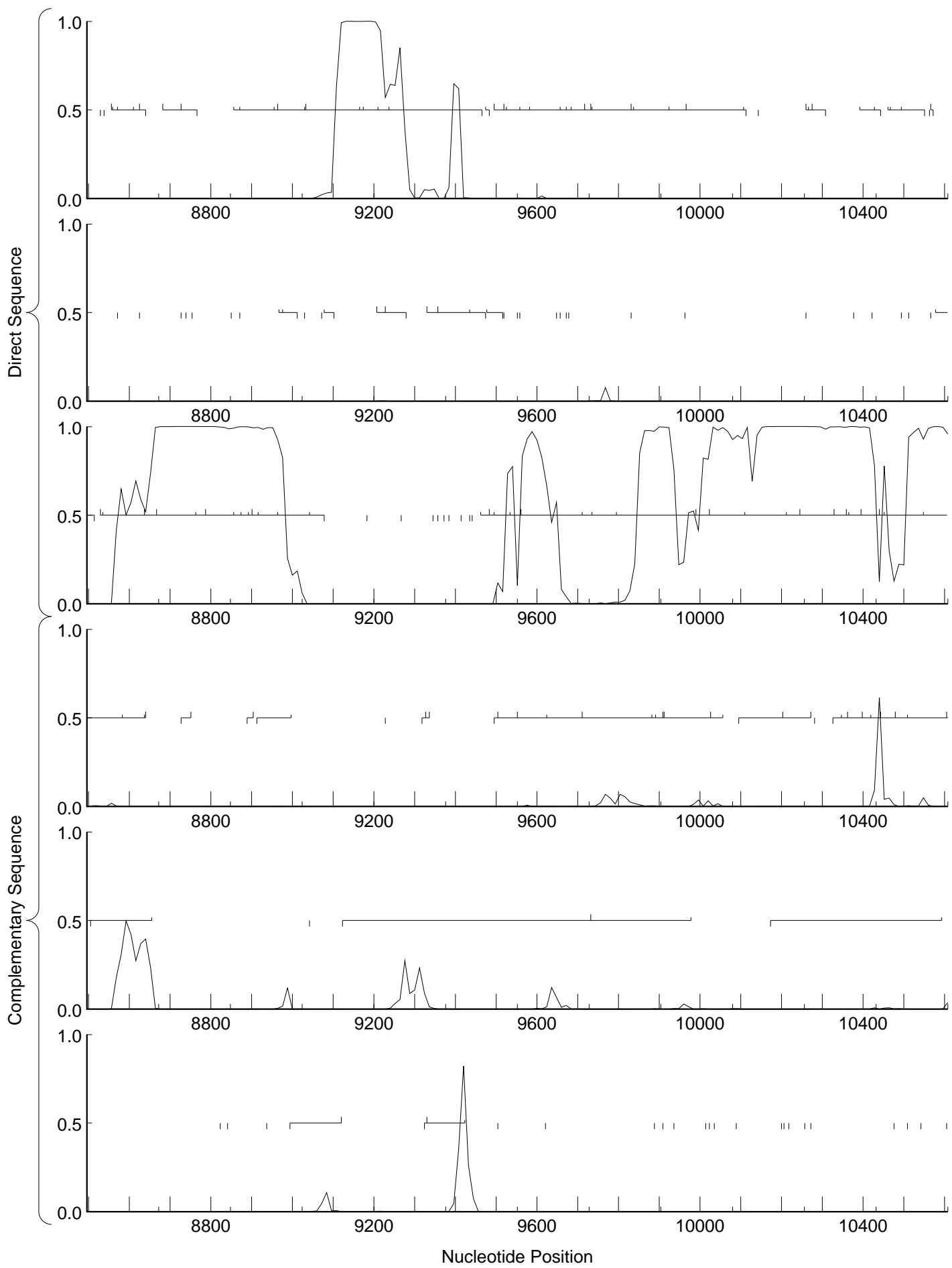
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

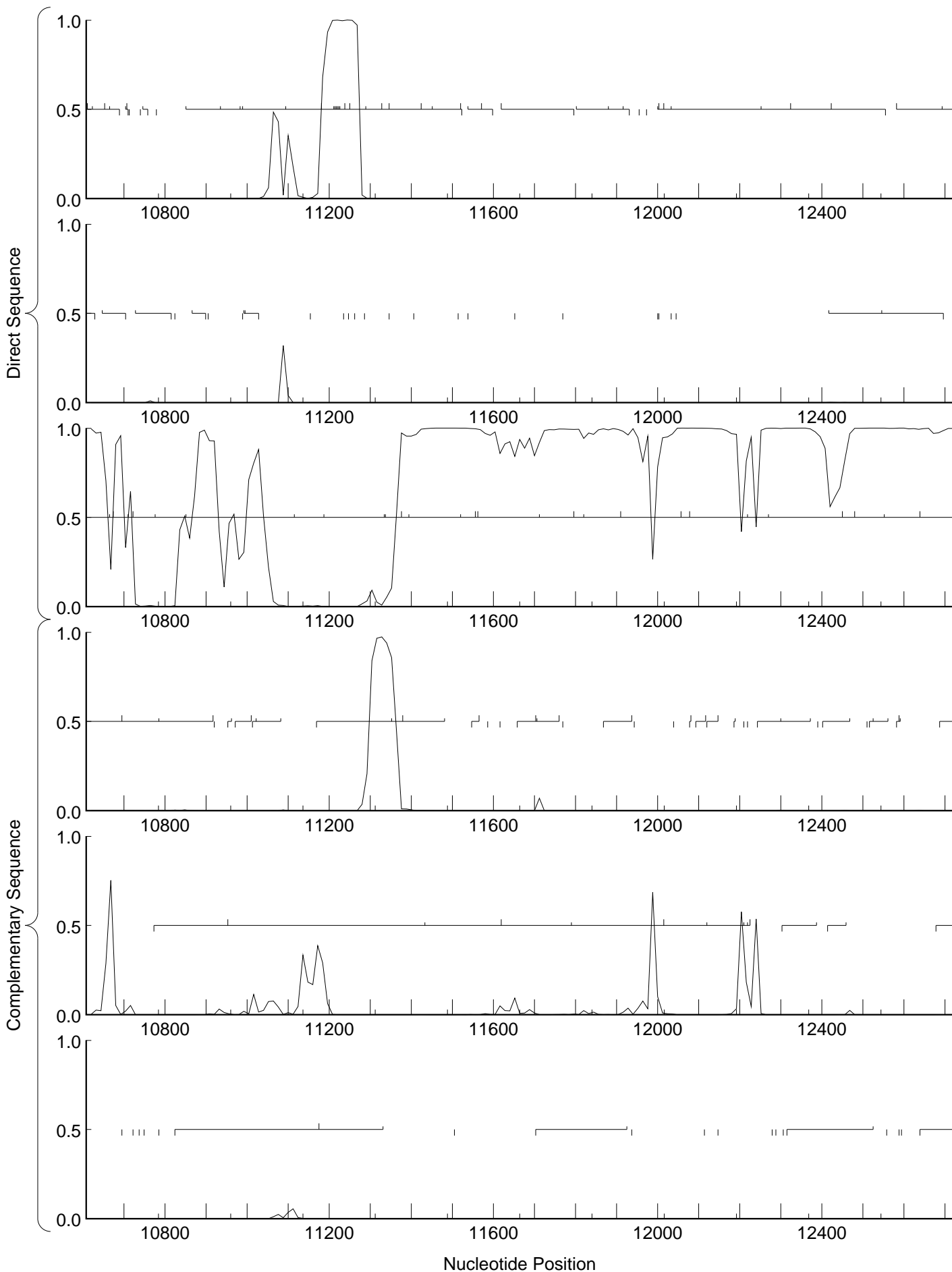


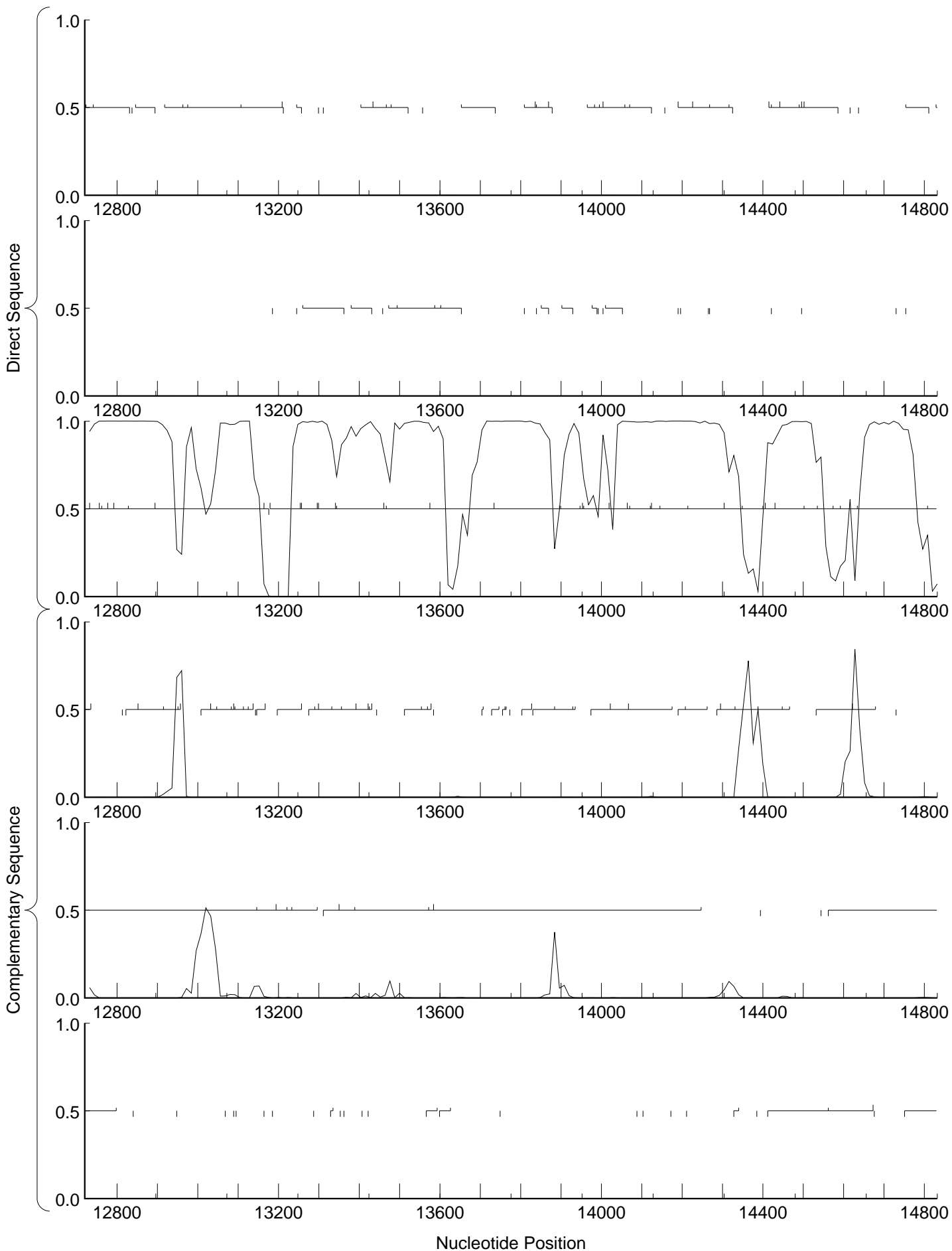


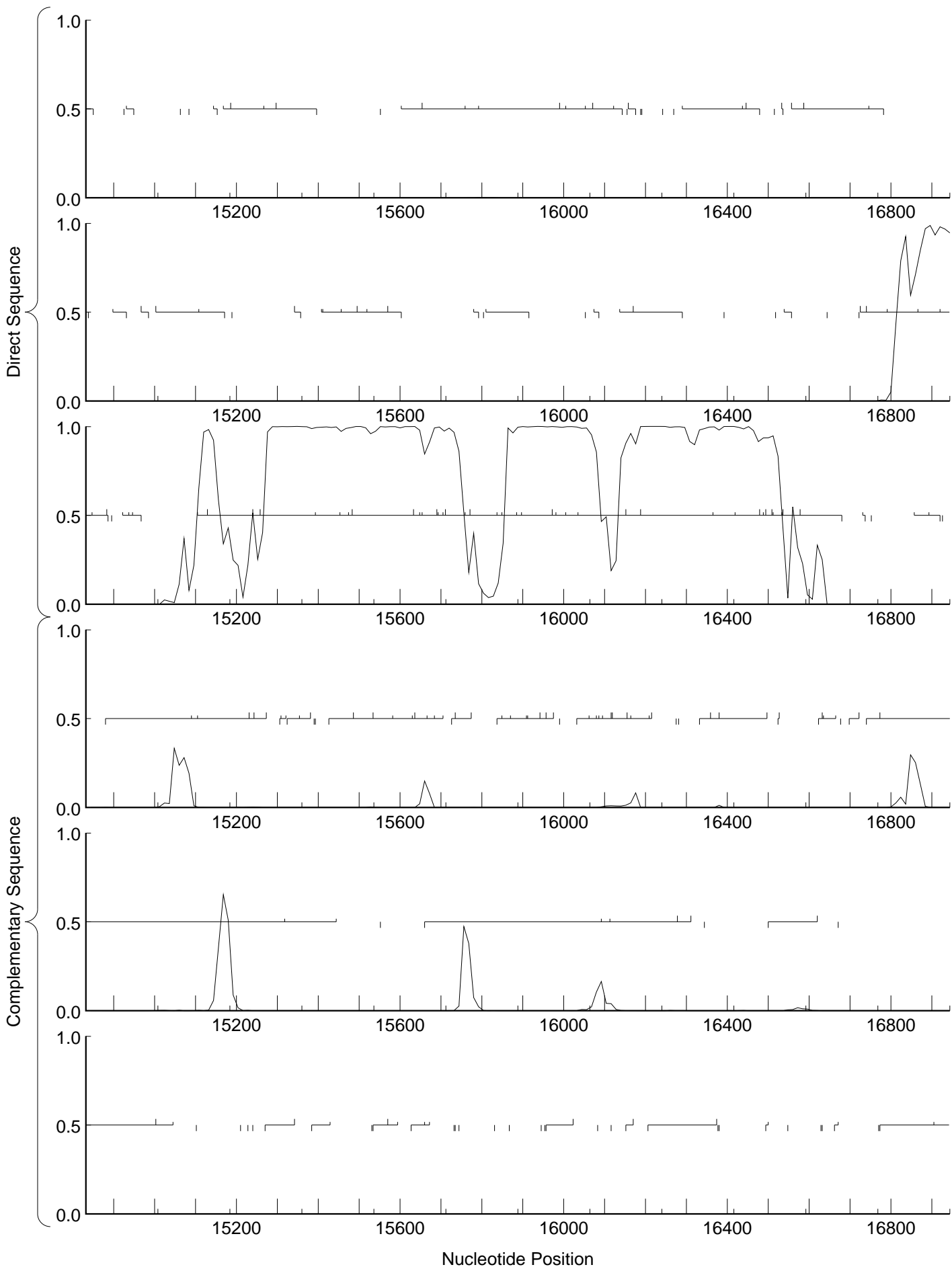




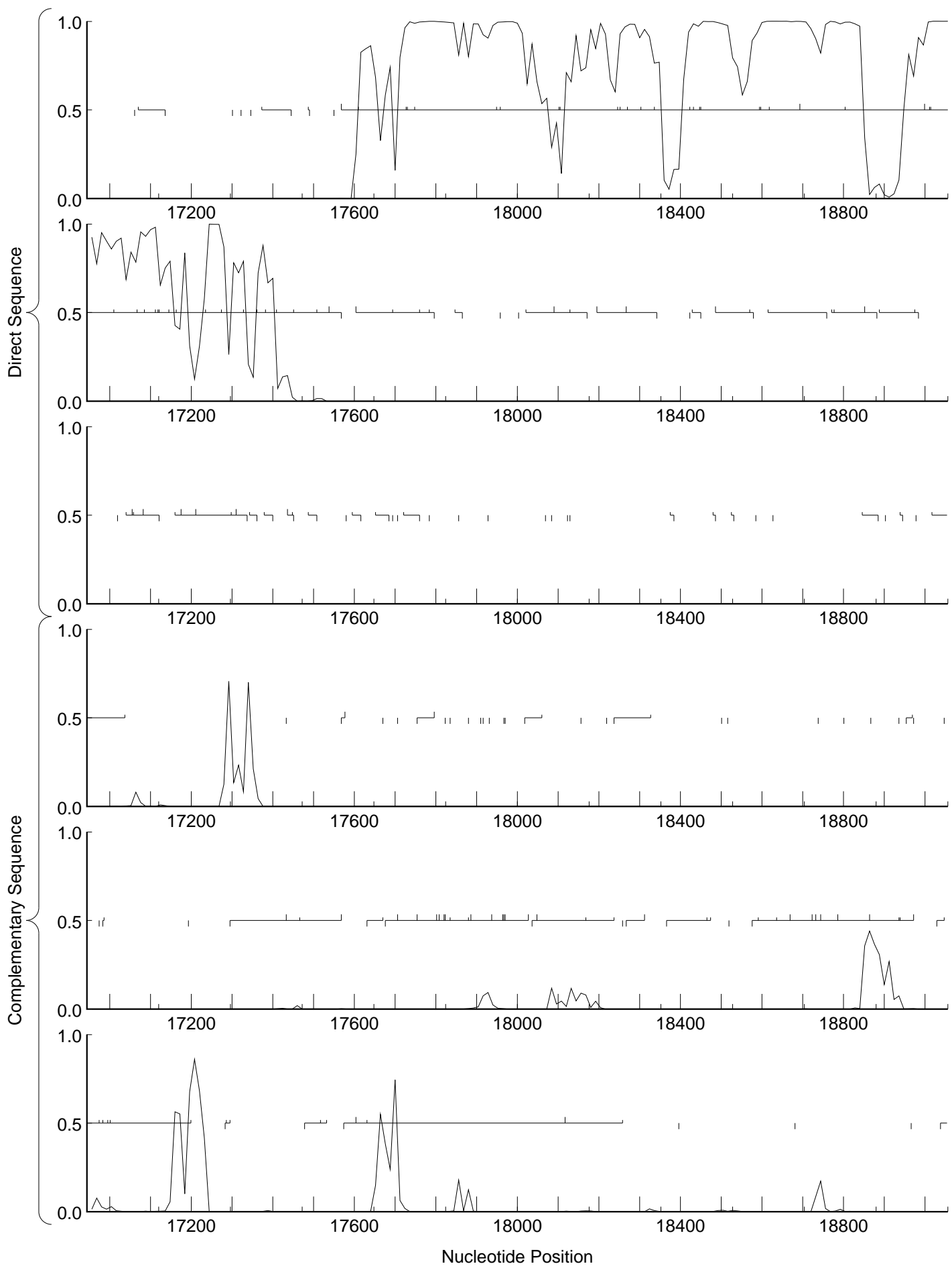




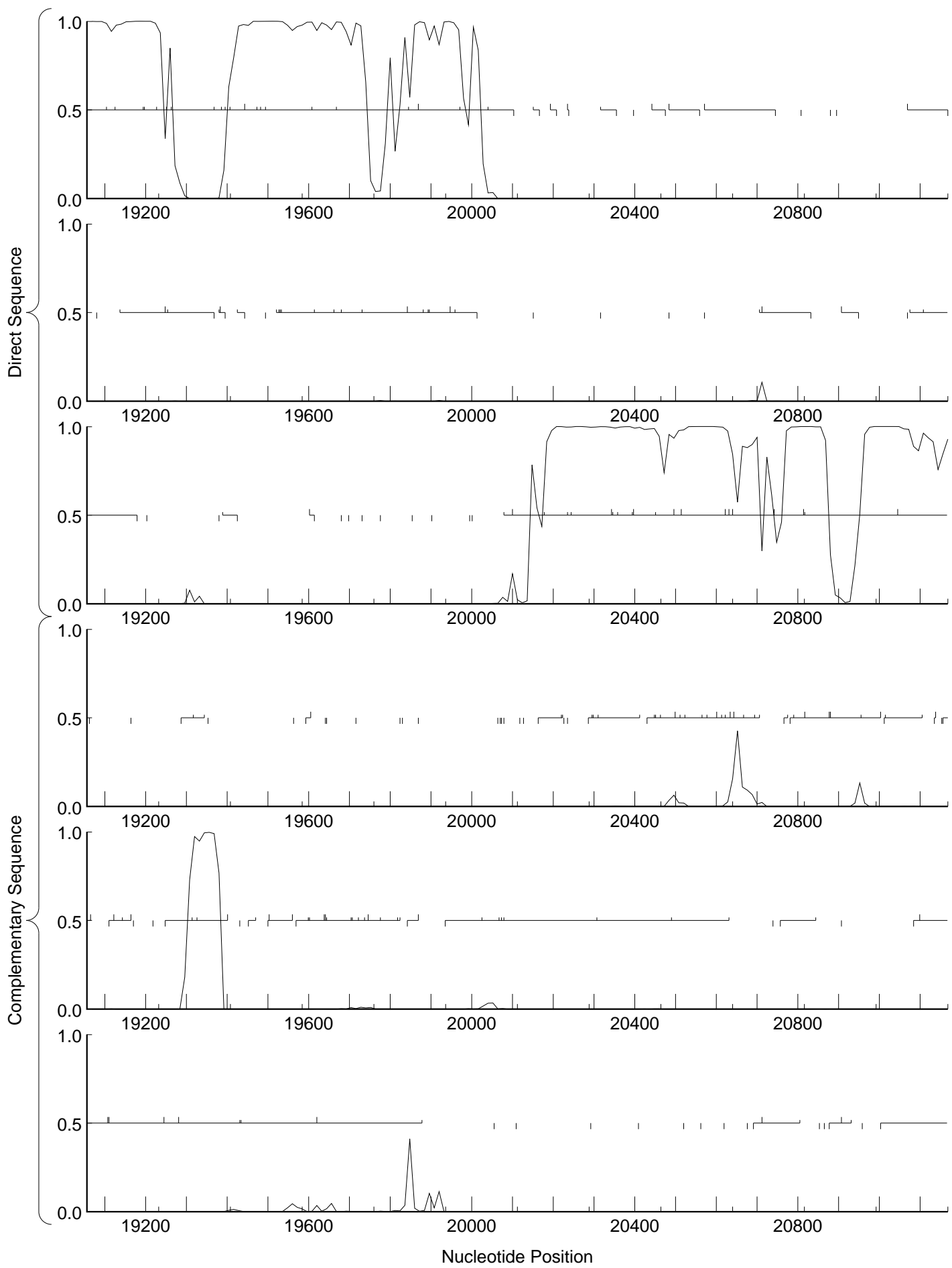


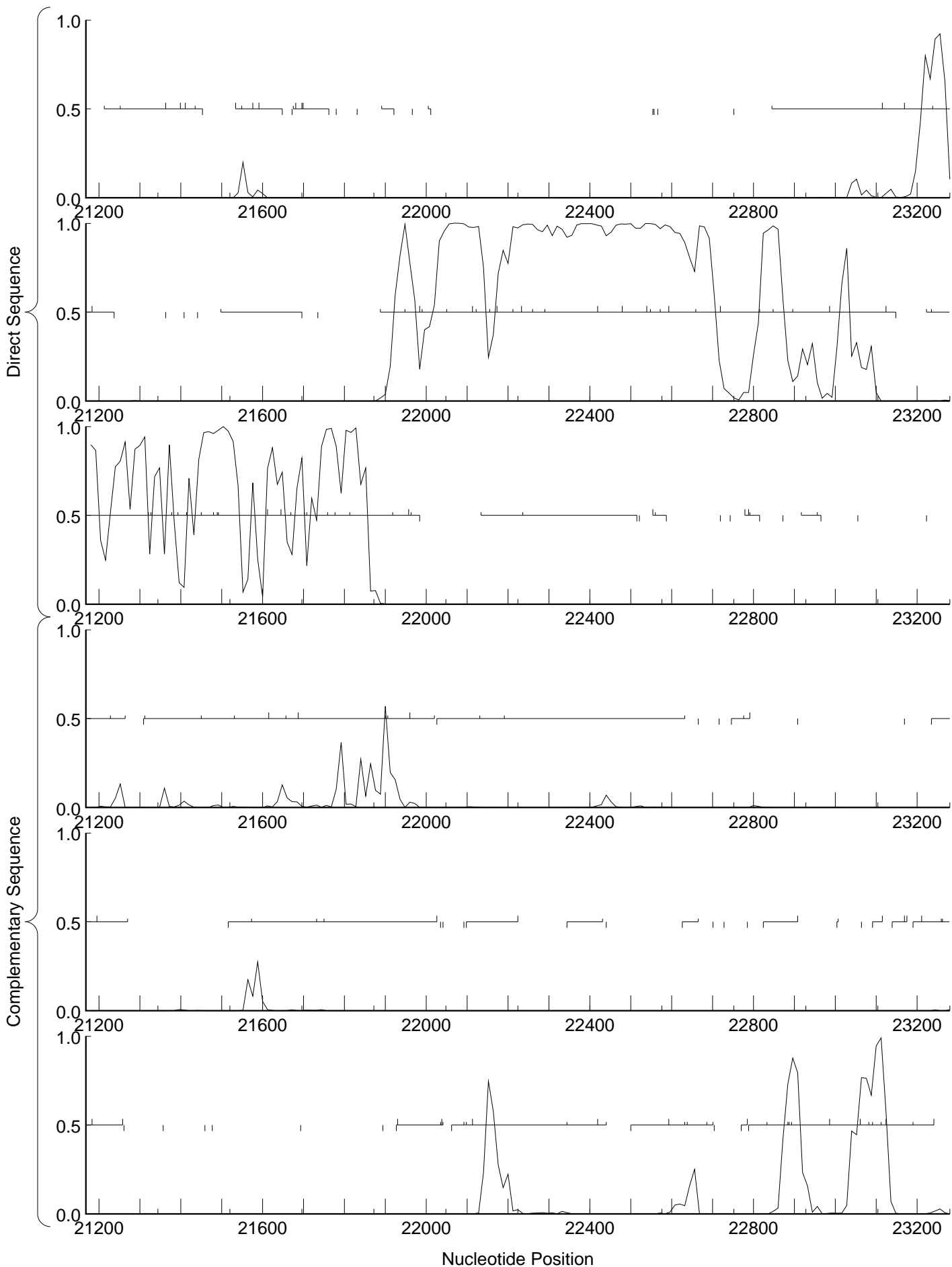


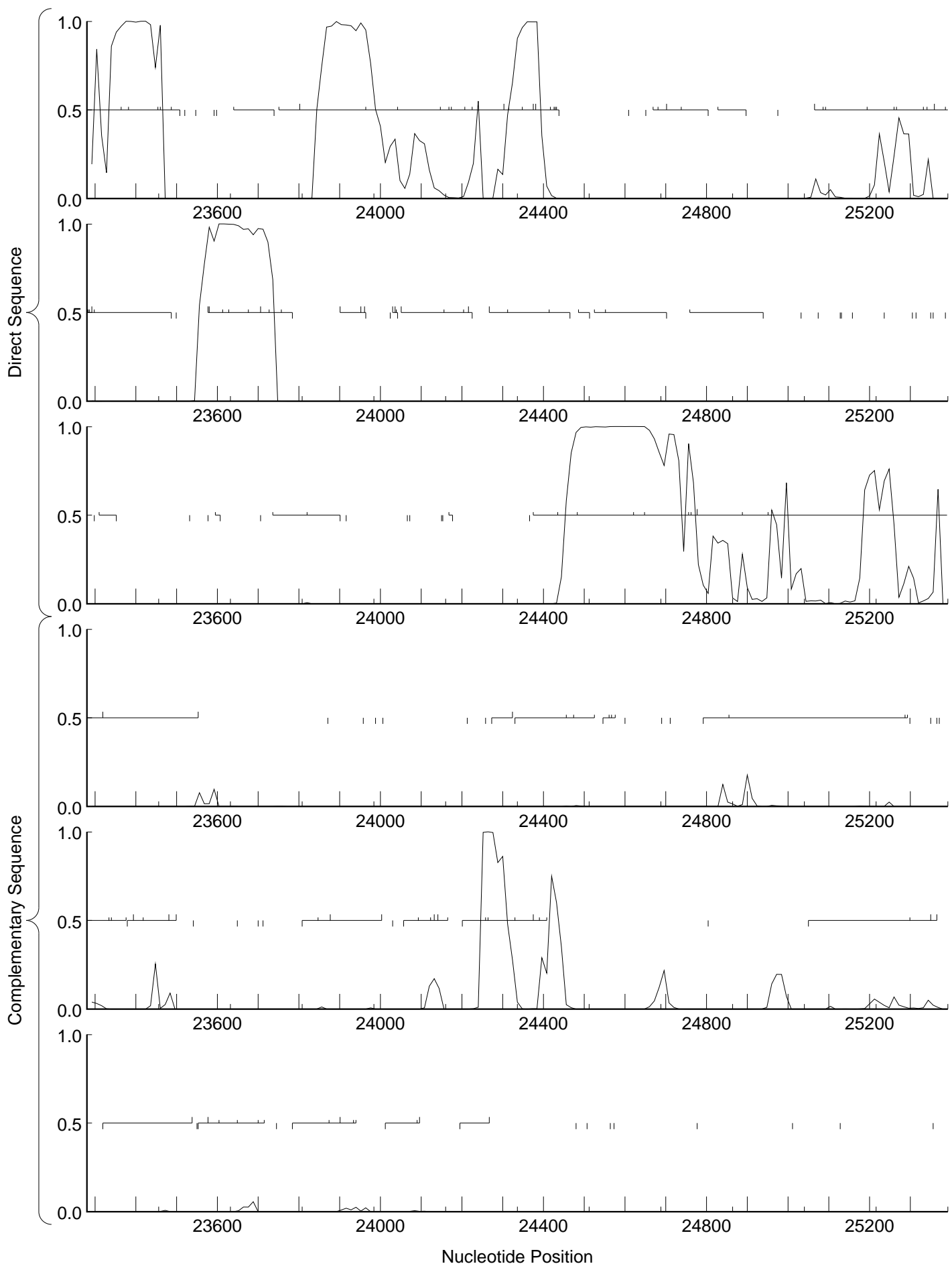
complete sequence, 60222 bp including 10-base 3' overhang (CGGACGGCGC), Cluster F1, Order 4, Window 96, Step 12, 10/30



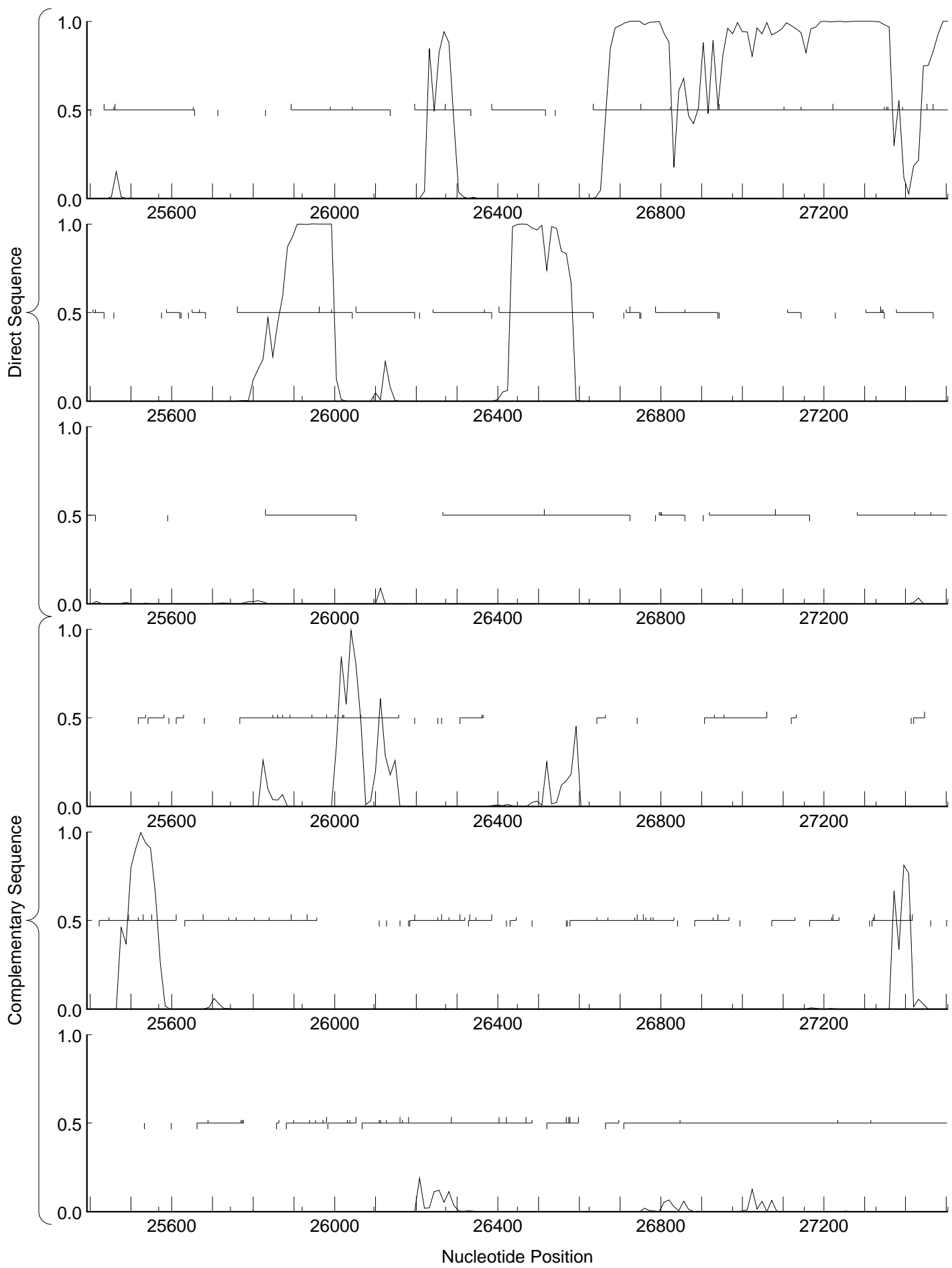
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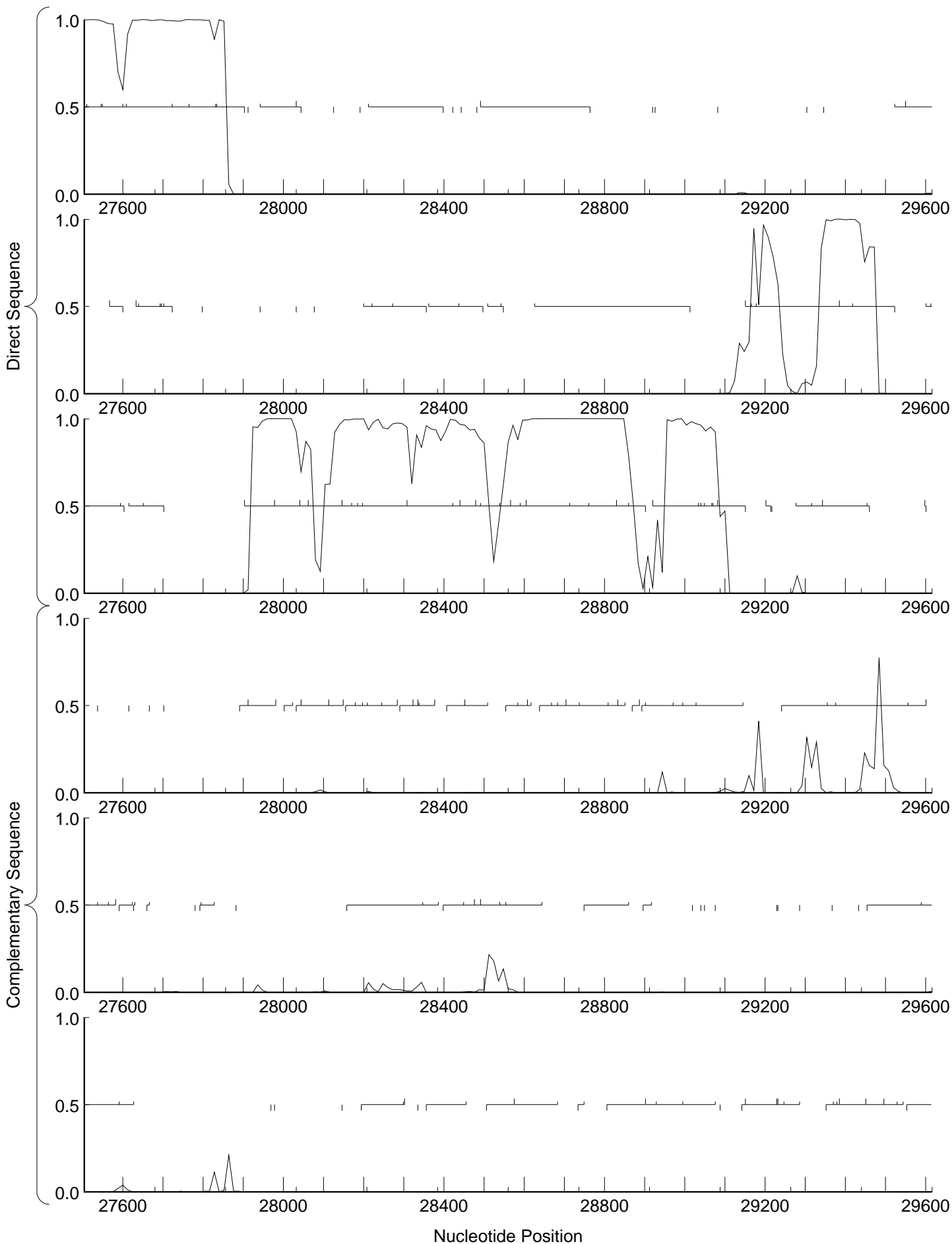


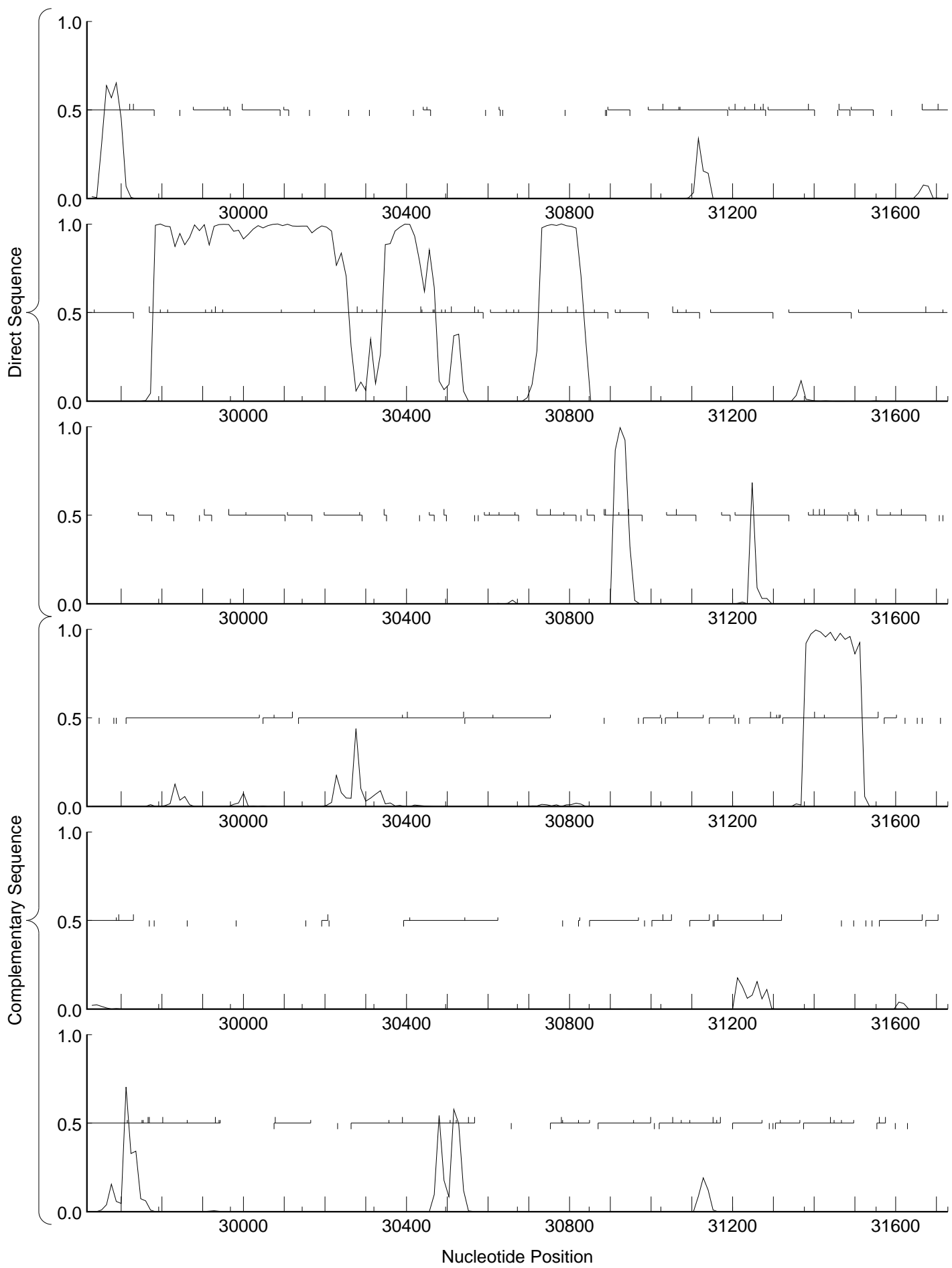




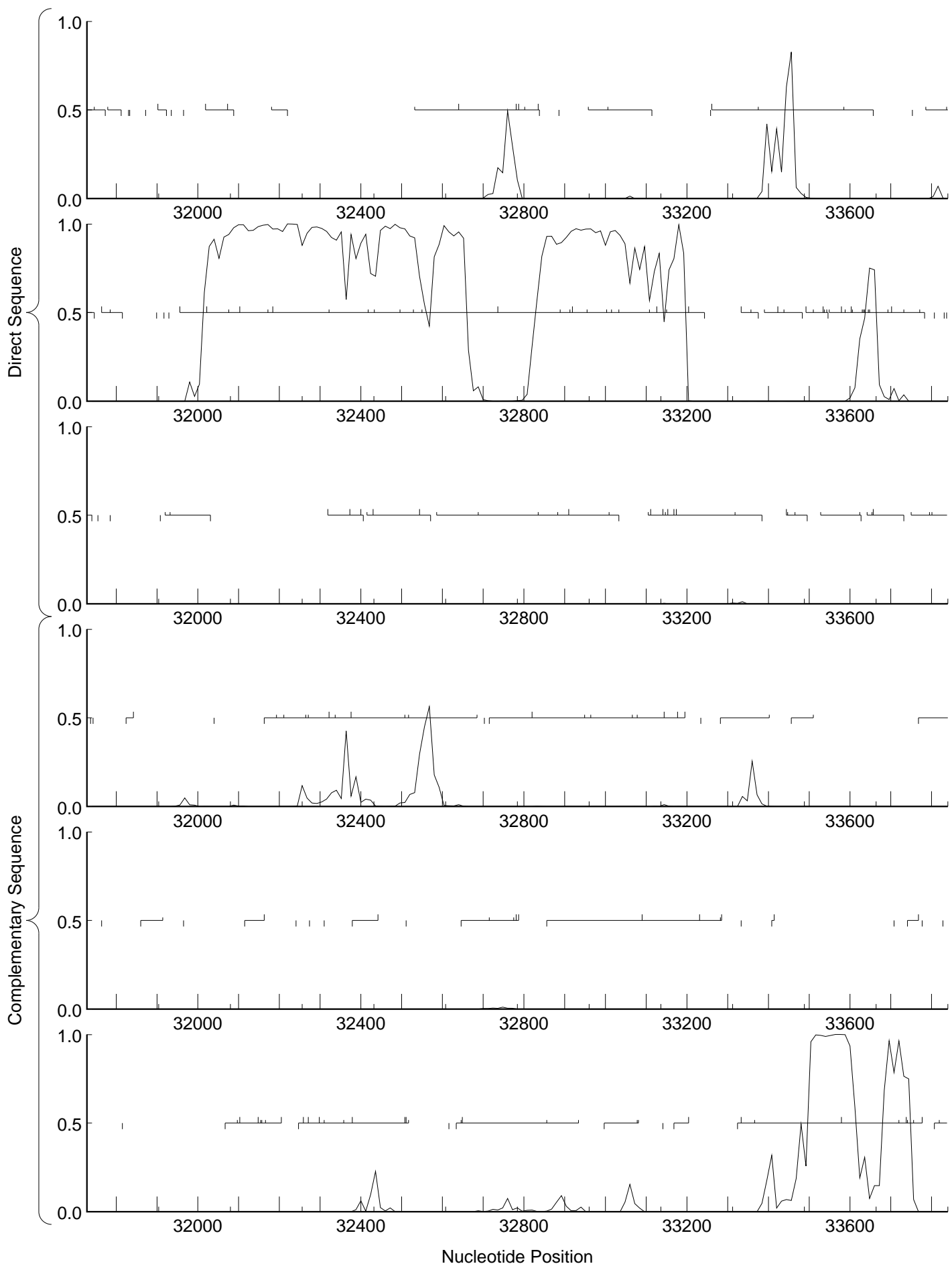
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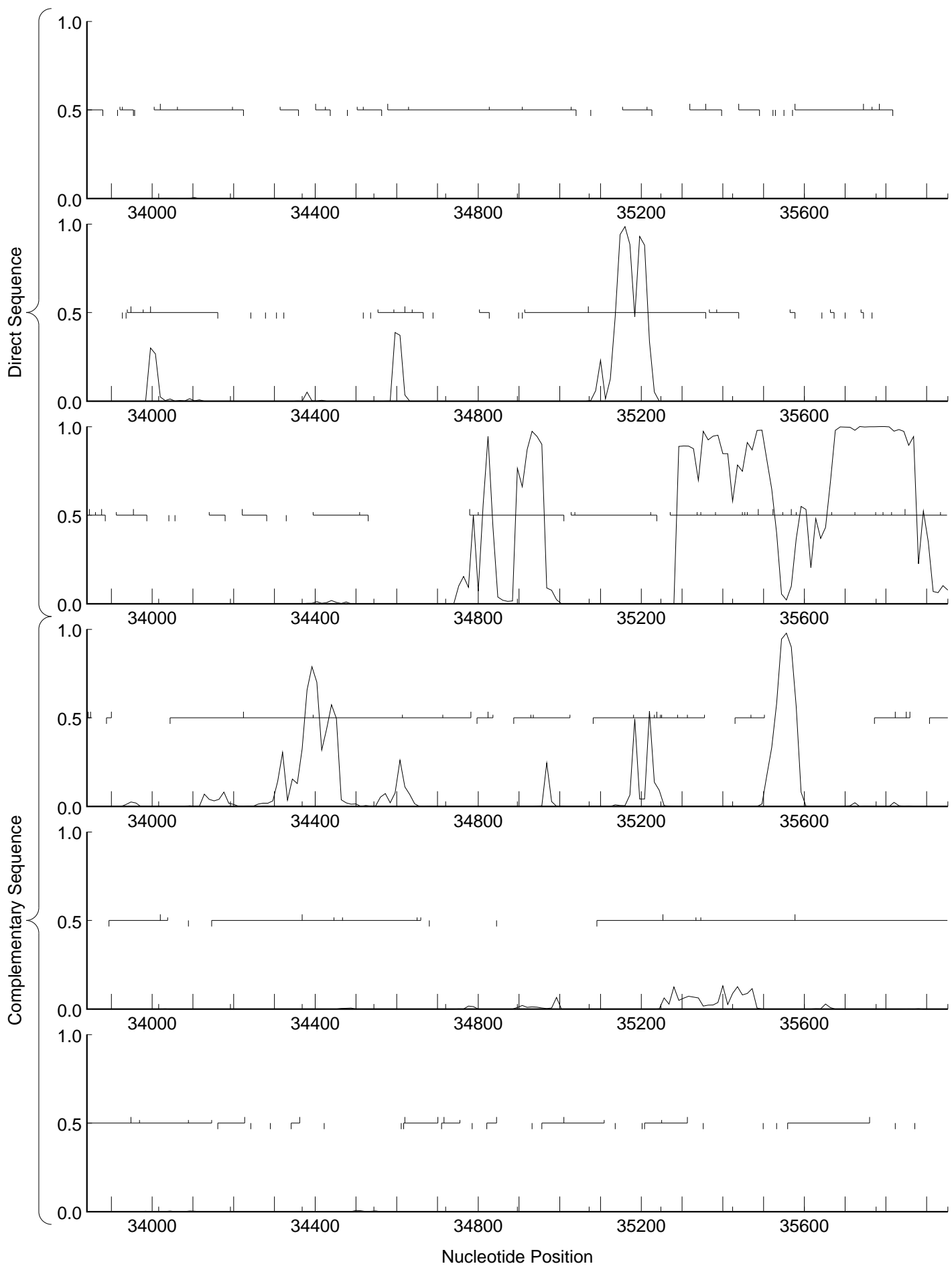


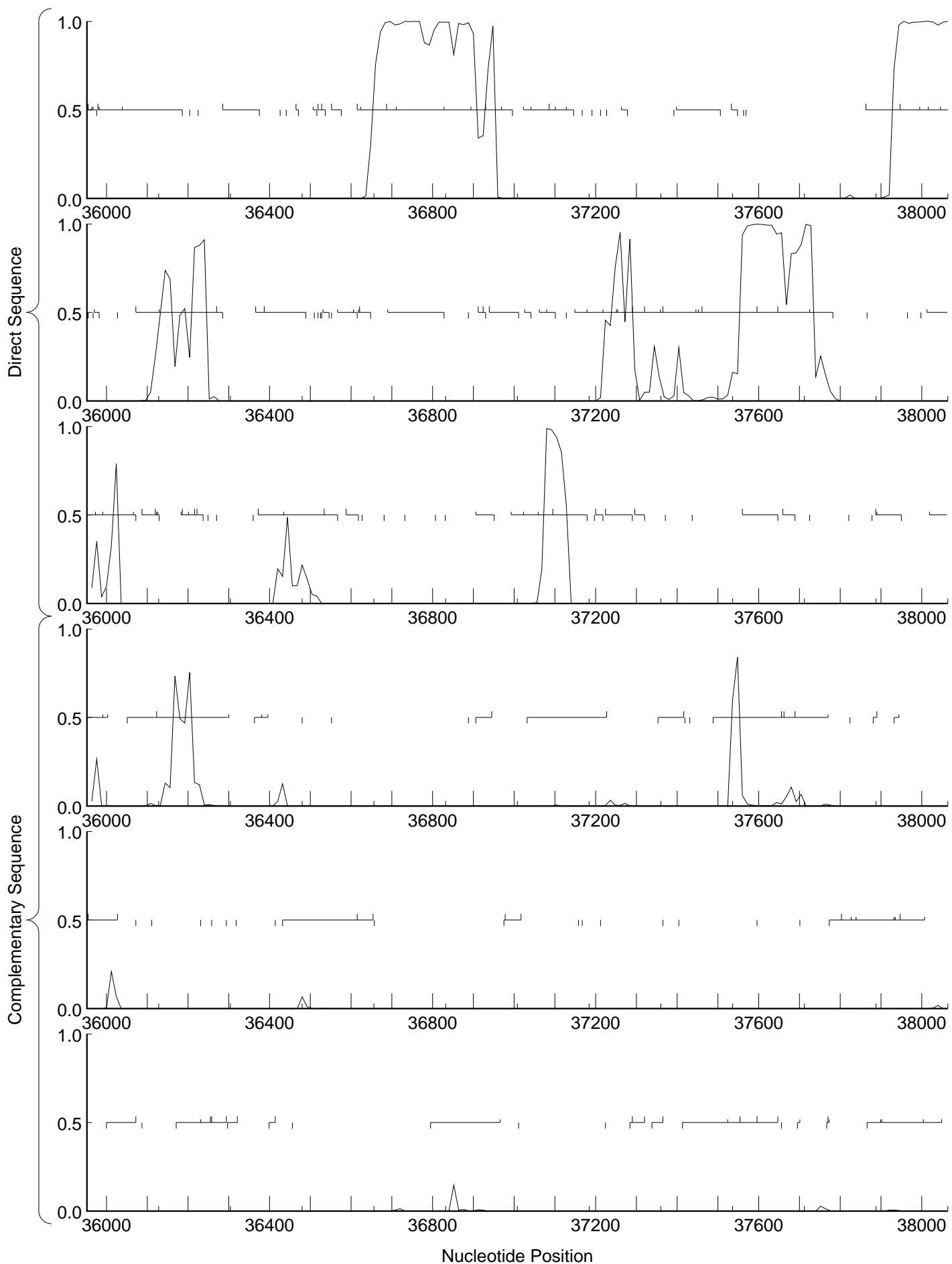


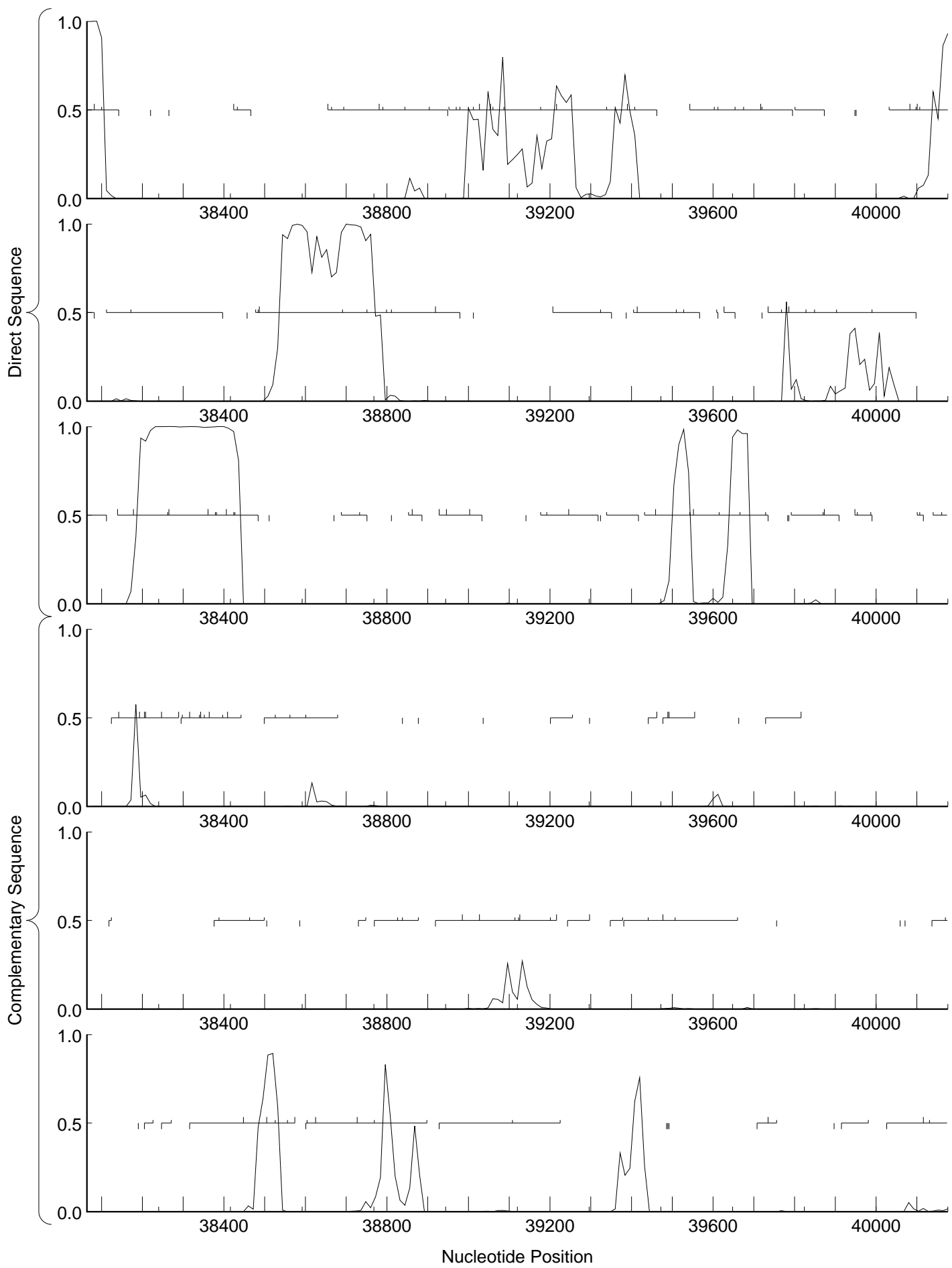


complete sequence, 60222 bp including 10-base 3' overhang (CGGACGGCGC), Cluster F1, Order 4, Window 96, Step 12, 17/30

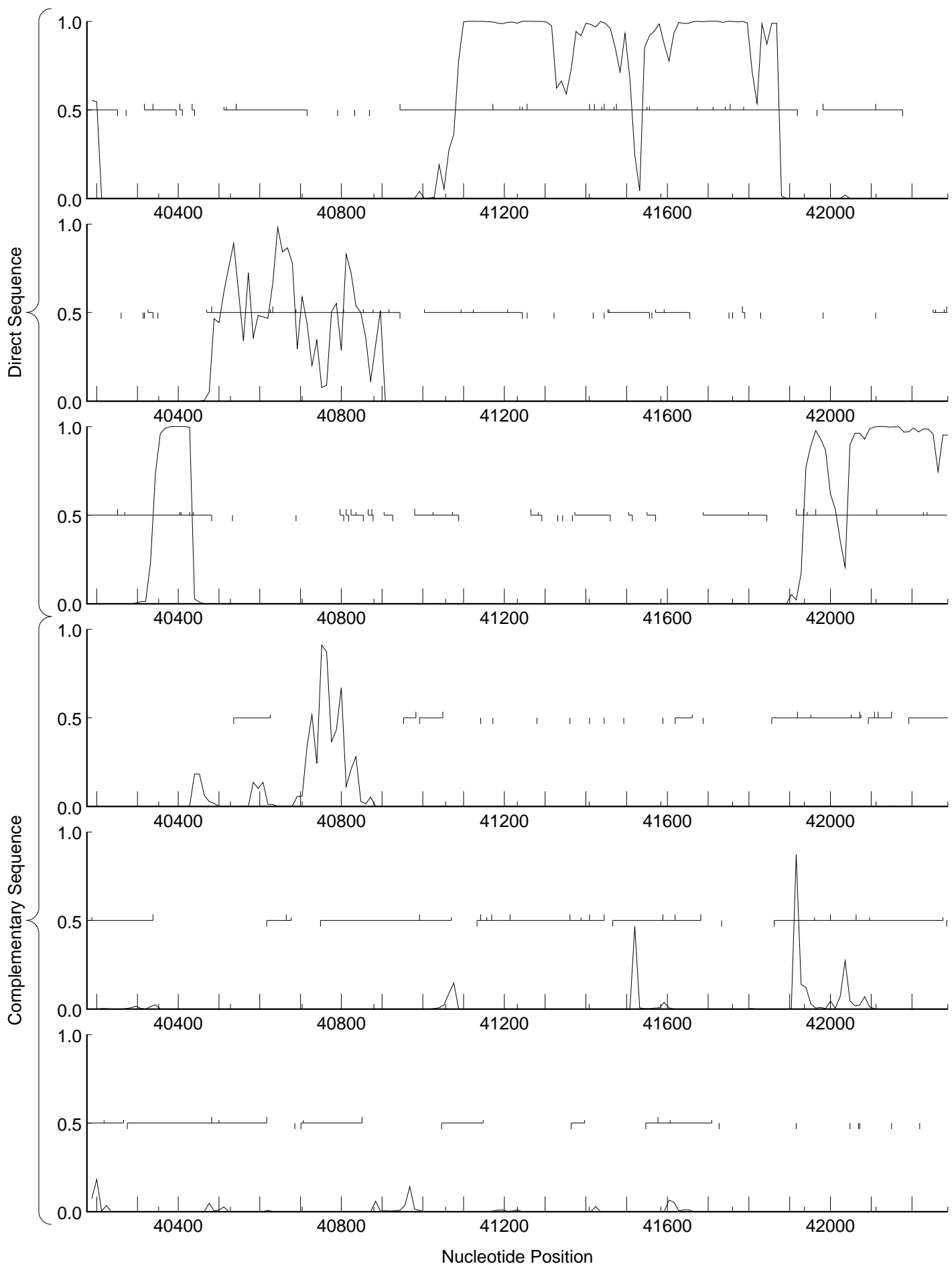


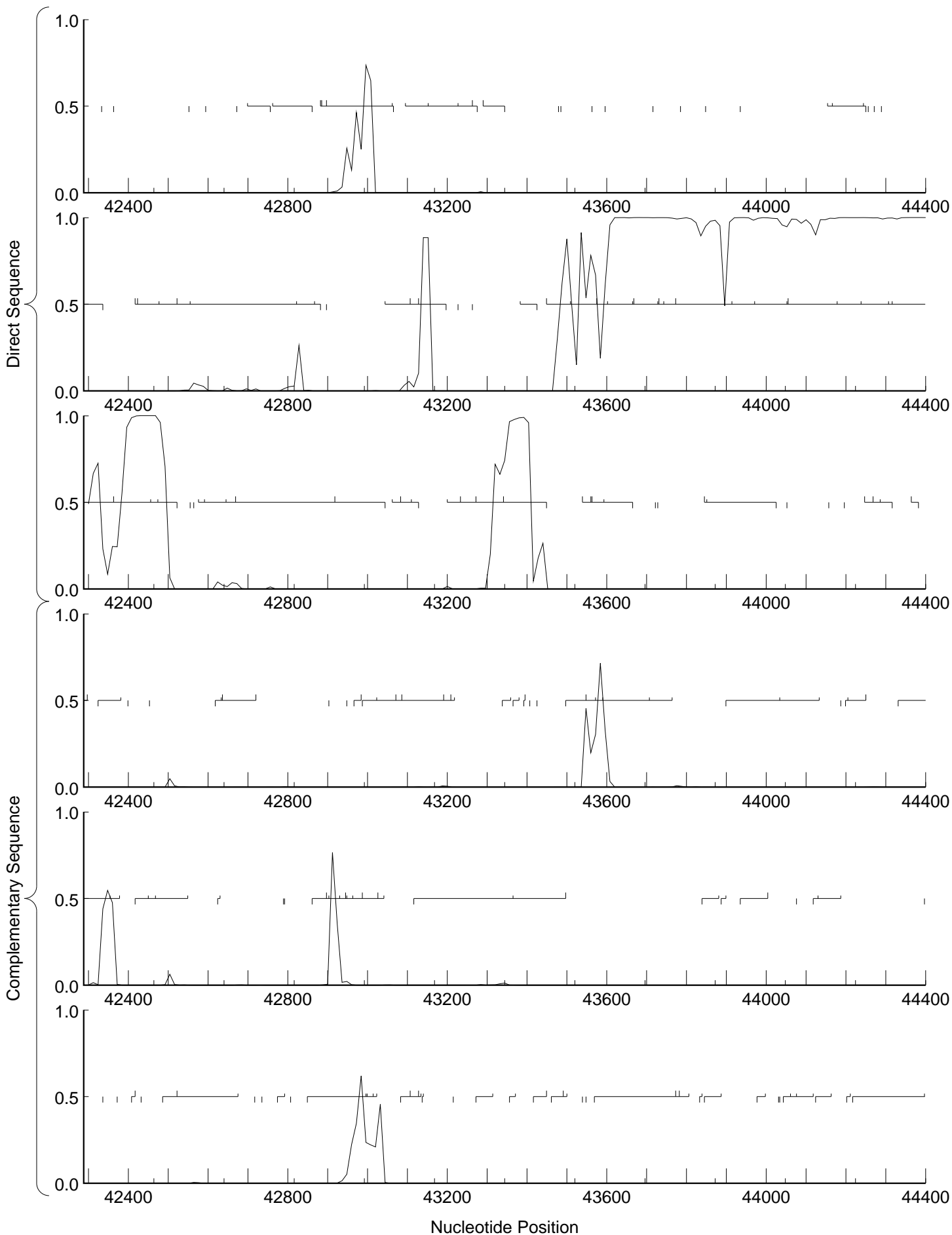


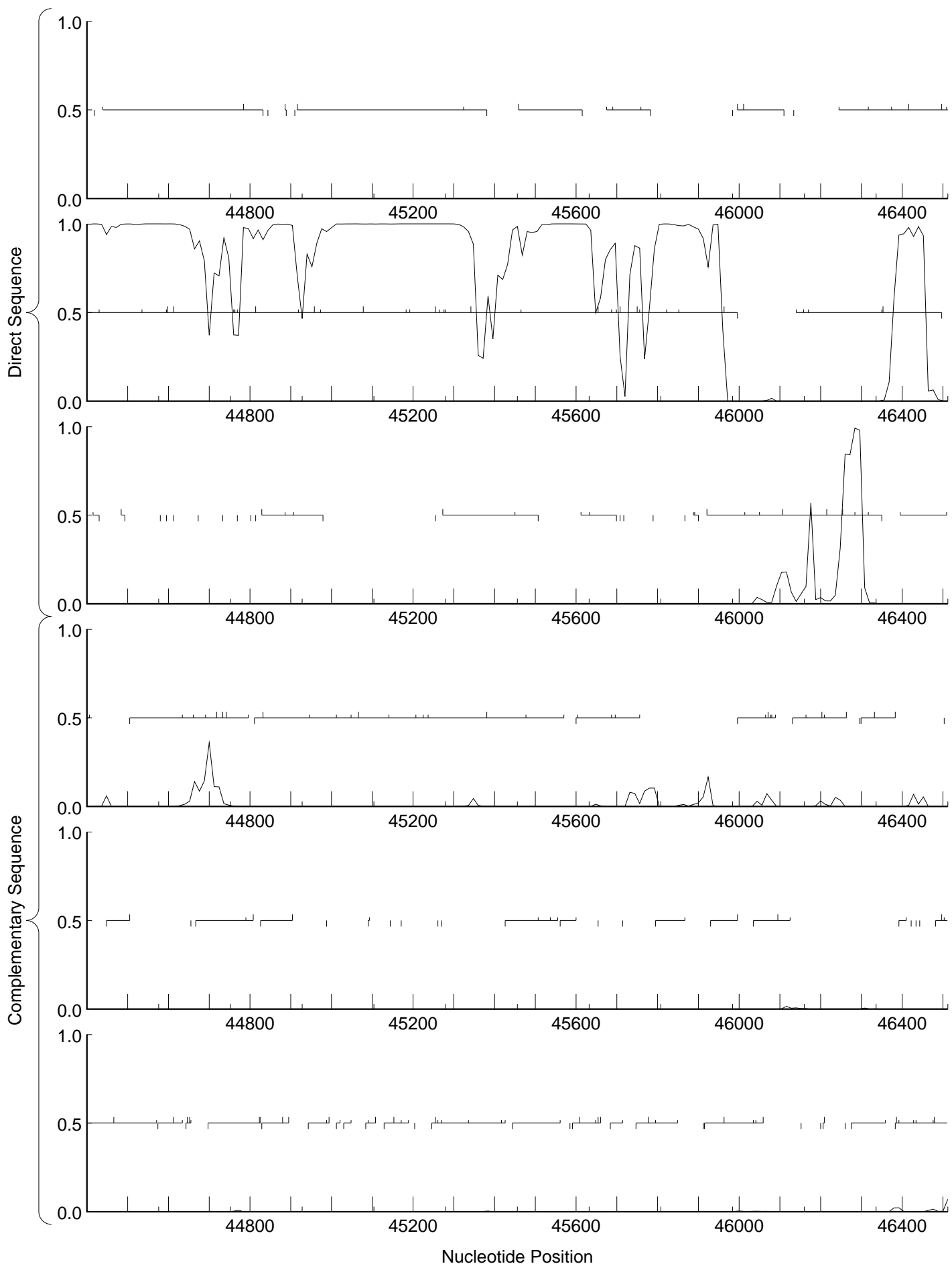




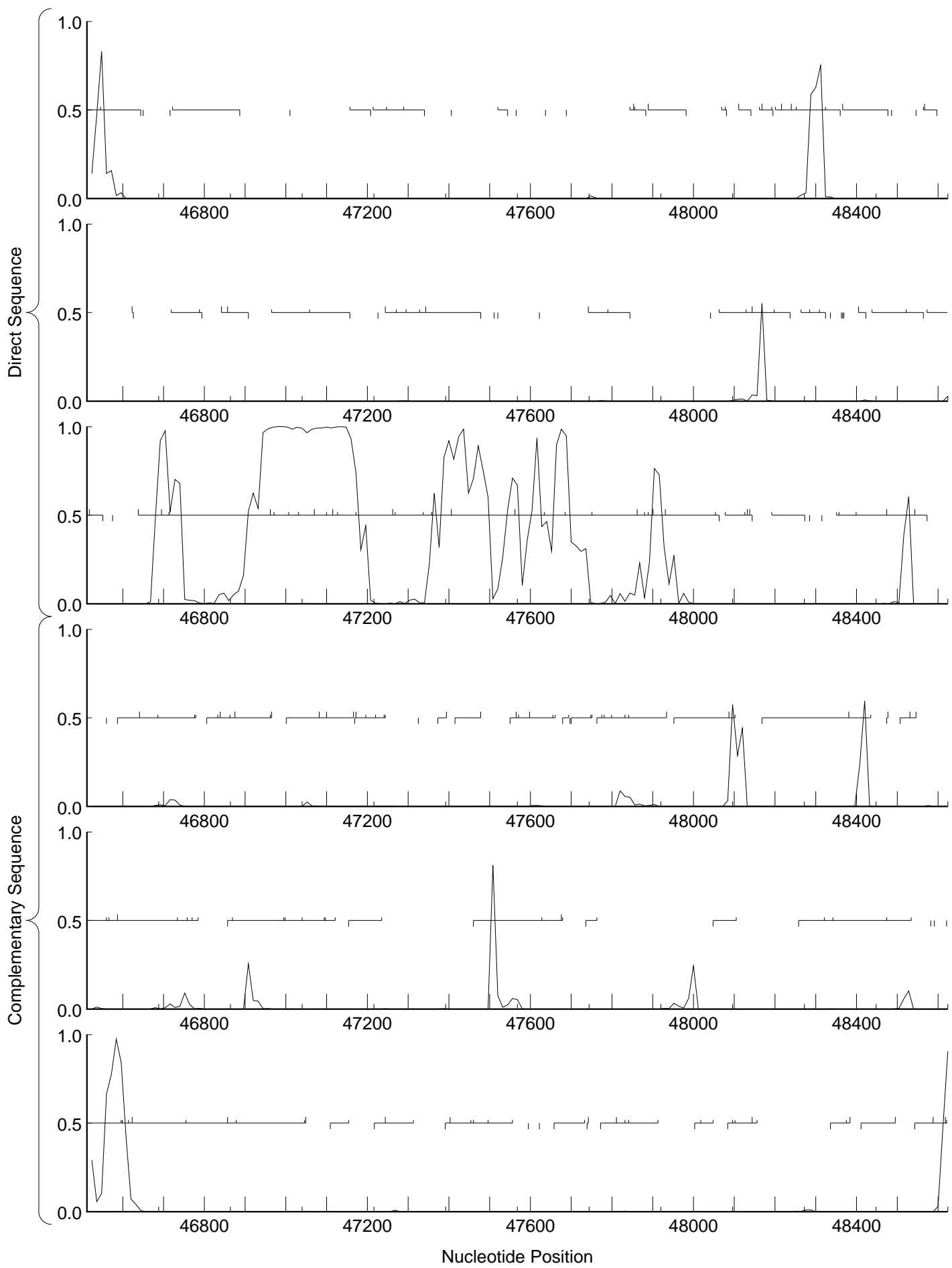
complete sequence, 60222 bp including 10-base 3' overhang (CGGACGGCGC), Cluster F1, Order 4, Window 96, Step 12, 21/30



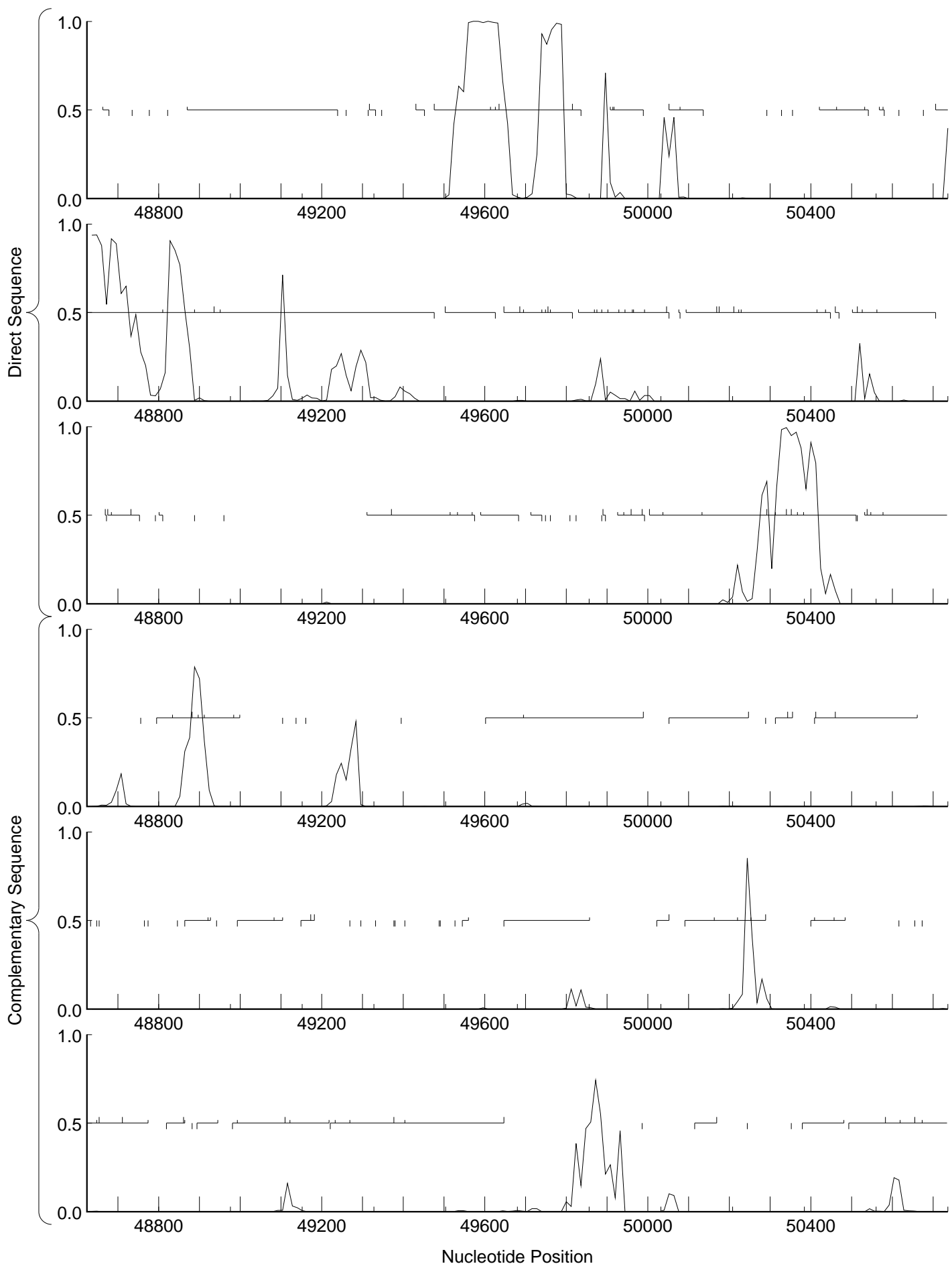


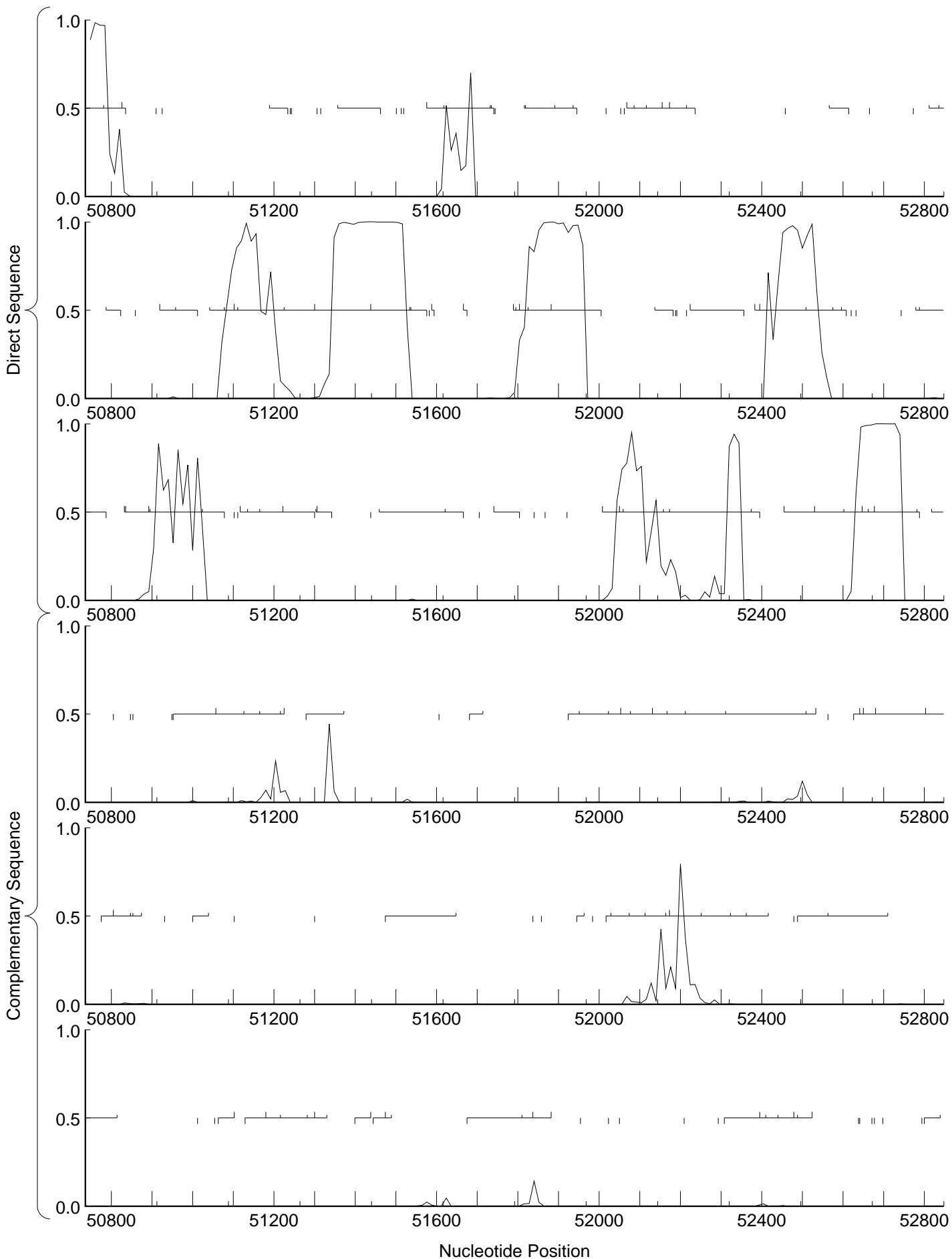


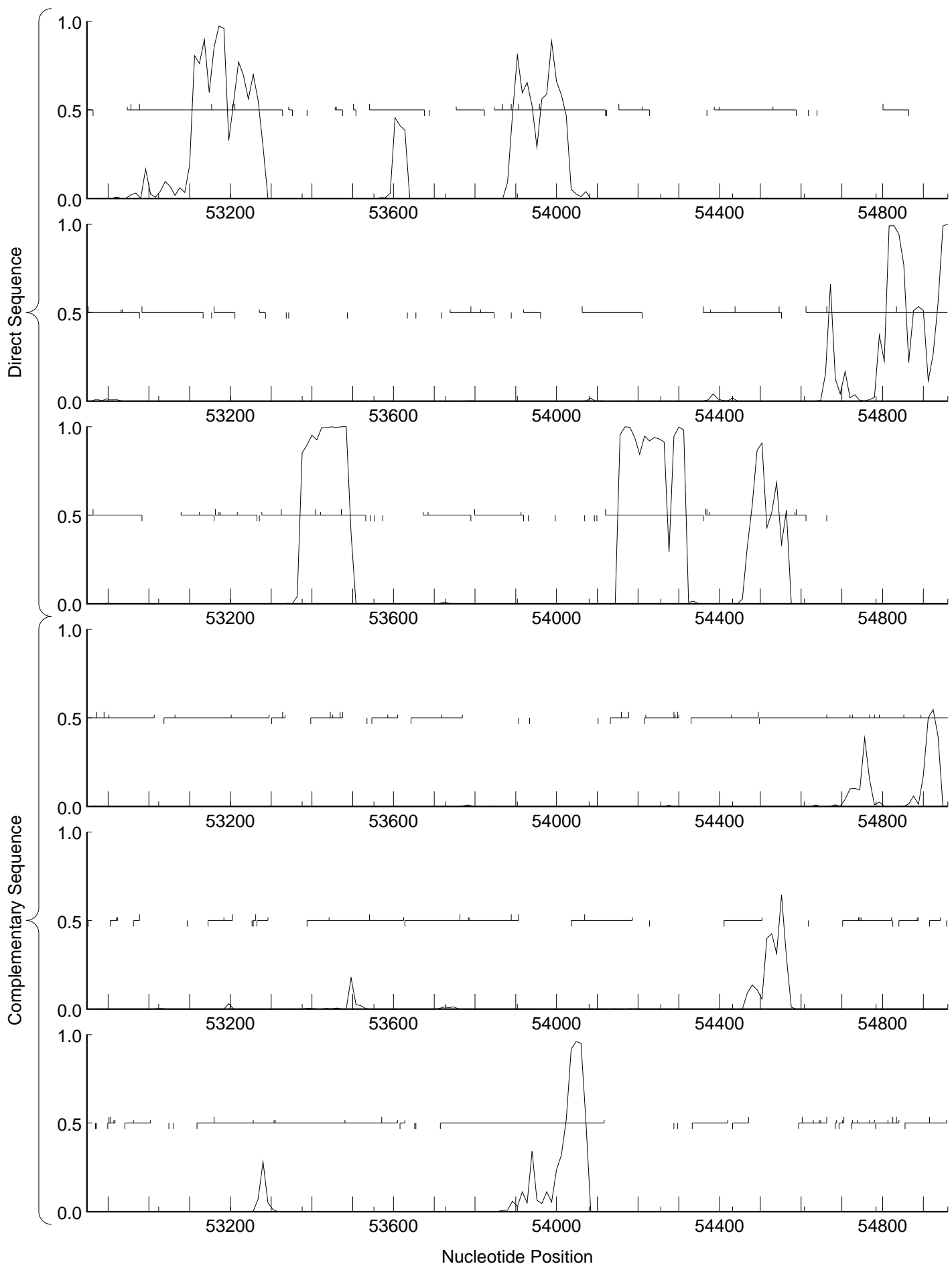
complete sequence, 60222 bp including 10-base 3' overhang (CGGACGGCGC), Cluster F1, Order 4, Window 96, Step 12, 24/30



complete sequence, 60222 bp including 10-base 3' overhang (CGGACGGCGC), Cluster F1, Order 4, Window 96, Step 12, 25/30







complete sequence, 60222 bp including 10-base 3' overhang (CGGACGGCGC), Cluster F1, Order 4, Window 96, Step 12, 28/30

