

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Kabluna complete sequence, 65869 bp including 10-base 3' overhang (CGCCGCGTA
Analysis Date : 8/15/20 at 0:55:08
Pages : 33
Sequence Length : 65869 bp
GC Content : 65.56%

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,

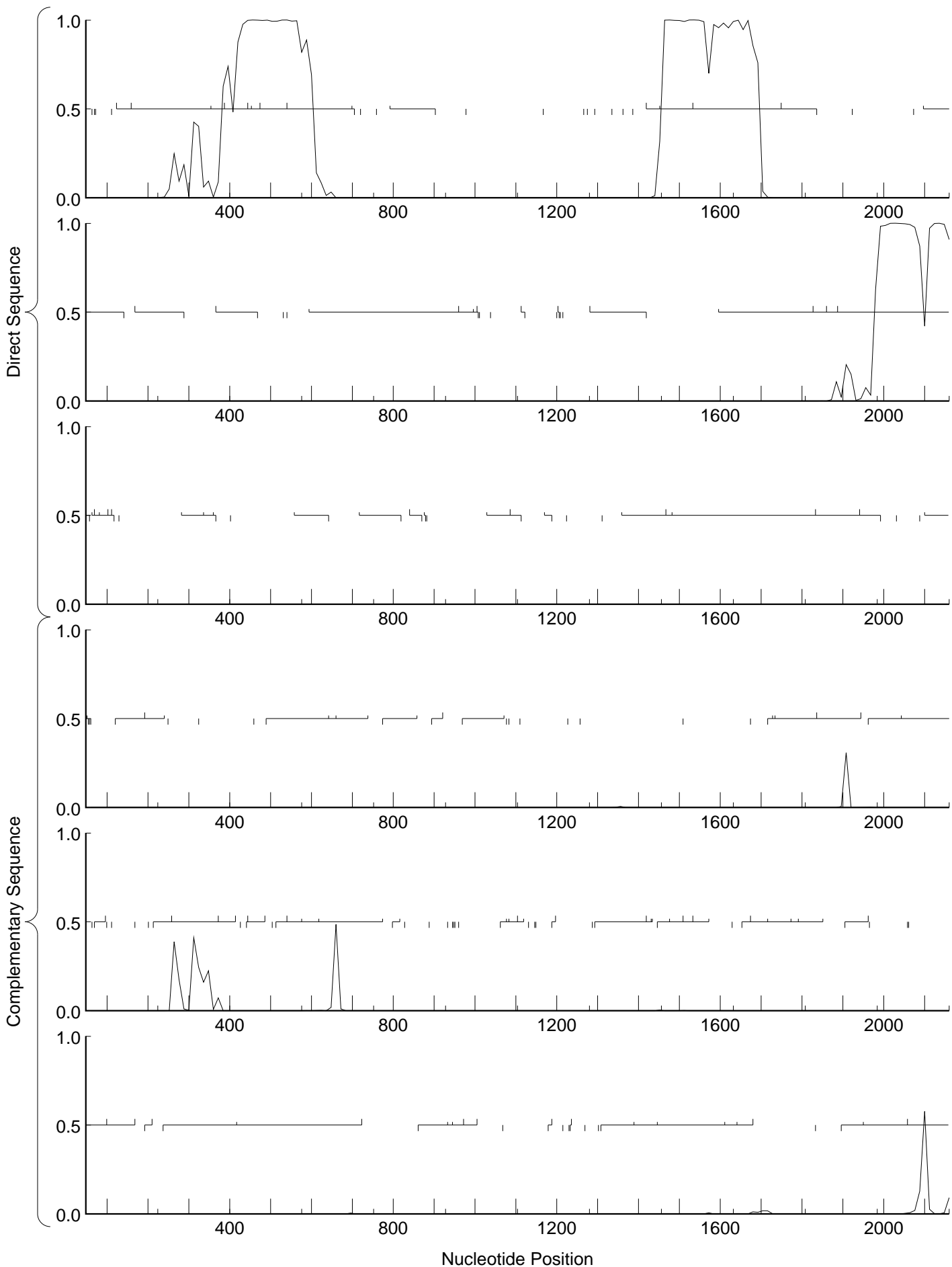
Matrix notes & comments

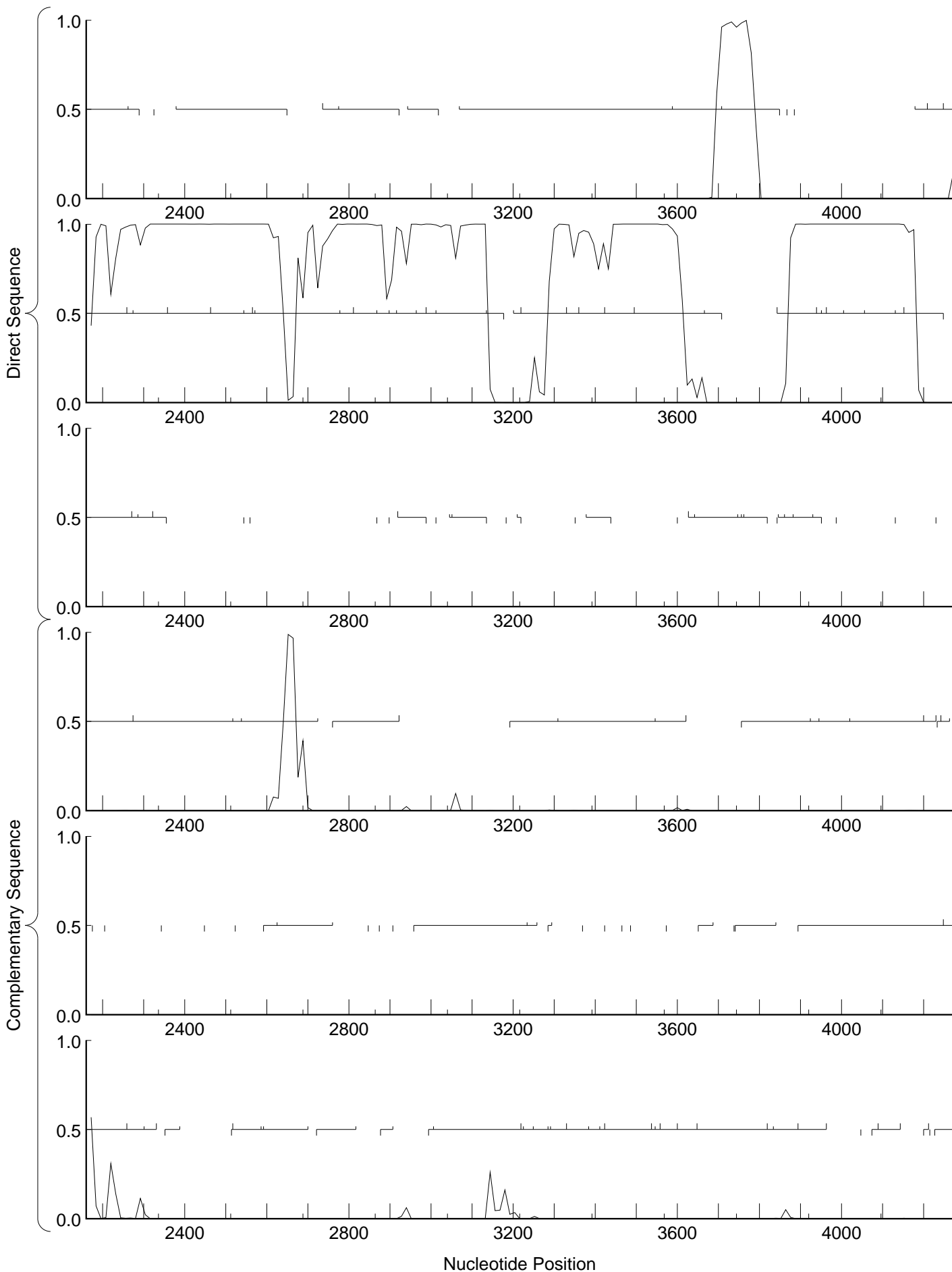
Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:36:14 2018

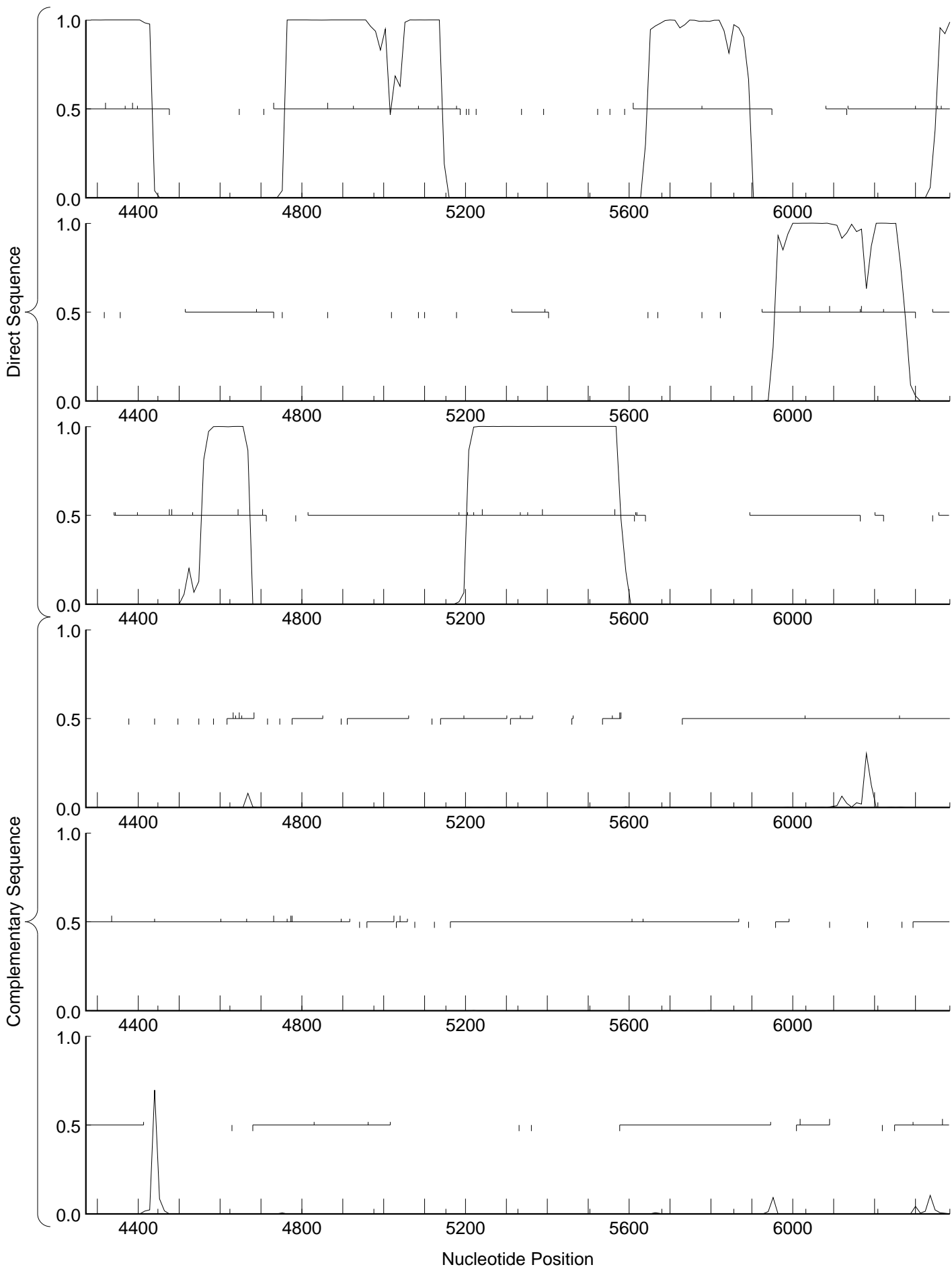
MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

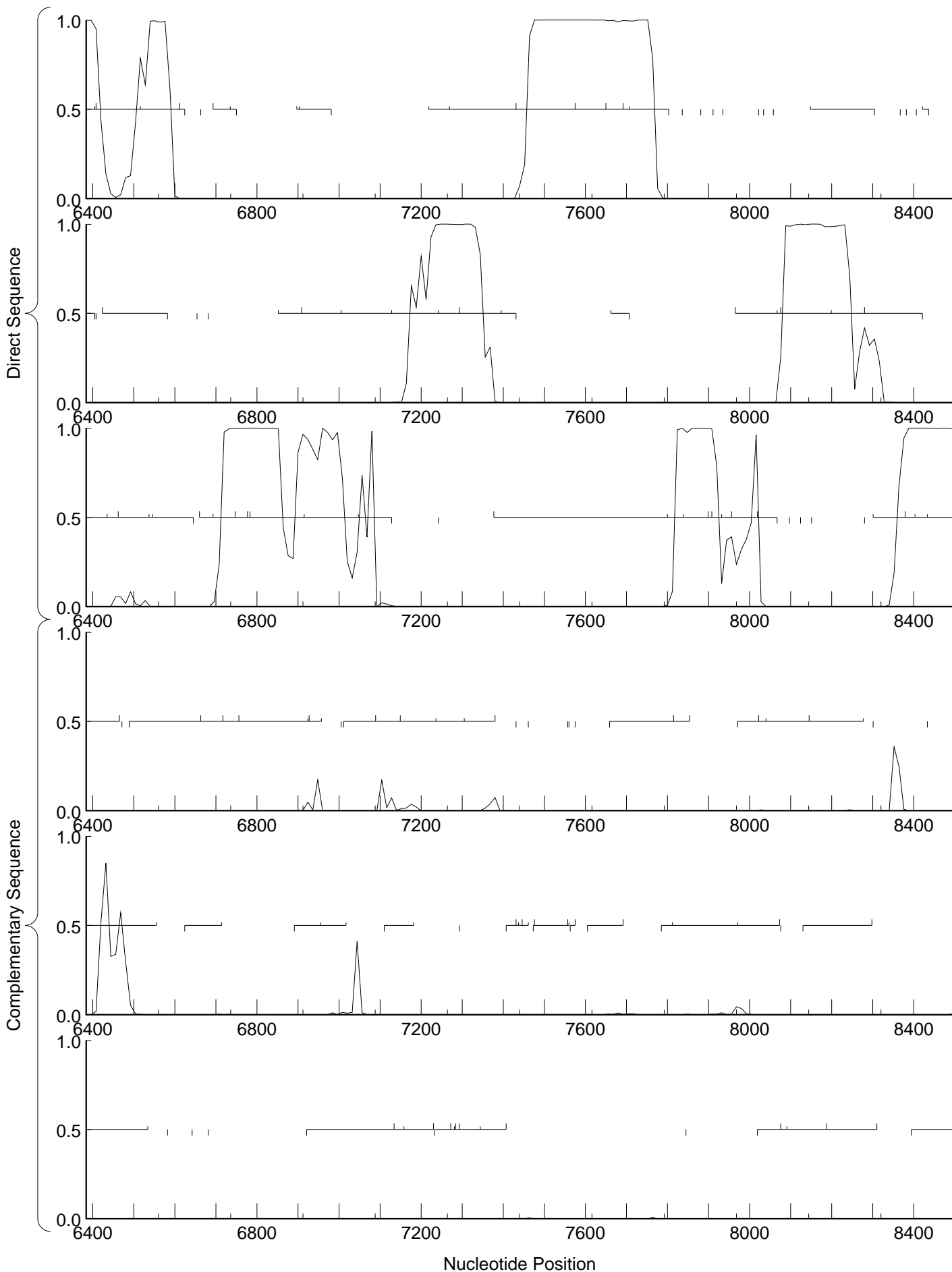
Send questions / comments to:
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Georgia Institute of Technology
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Atlanta, GA 30332-0230

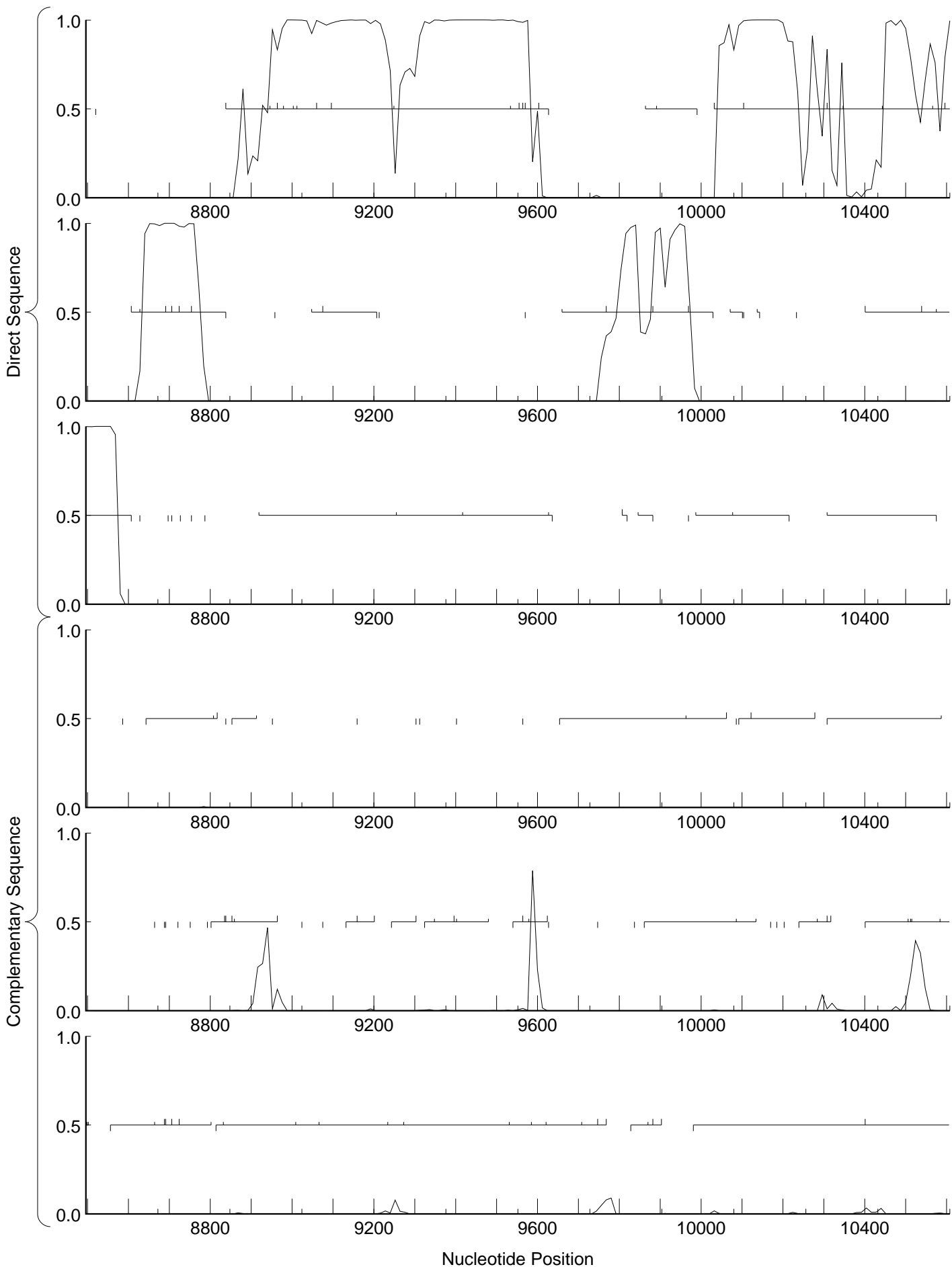




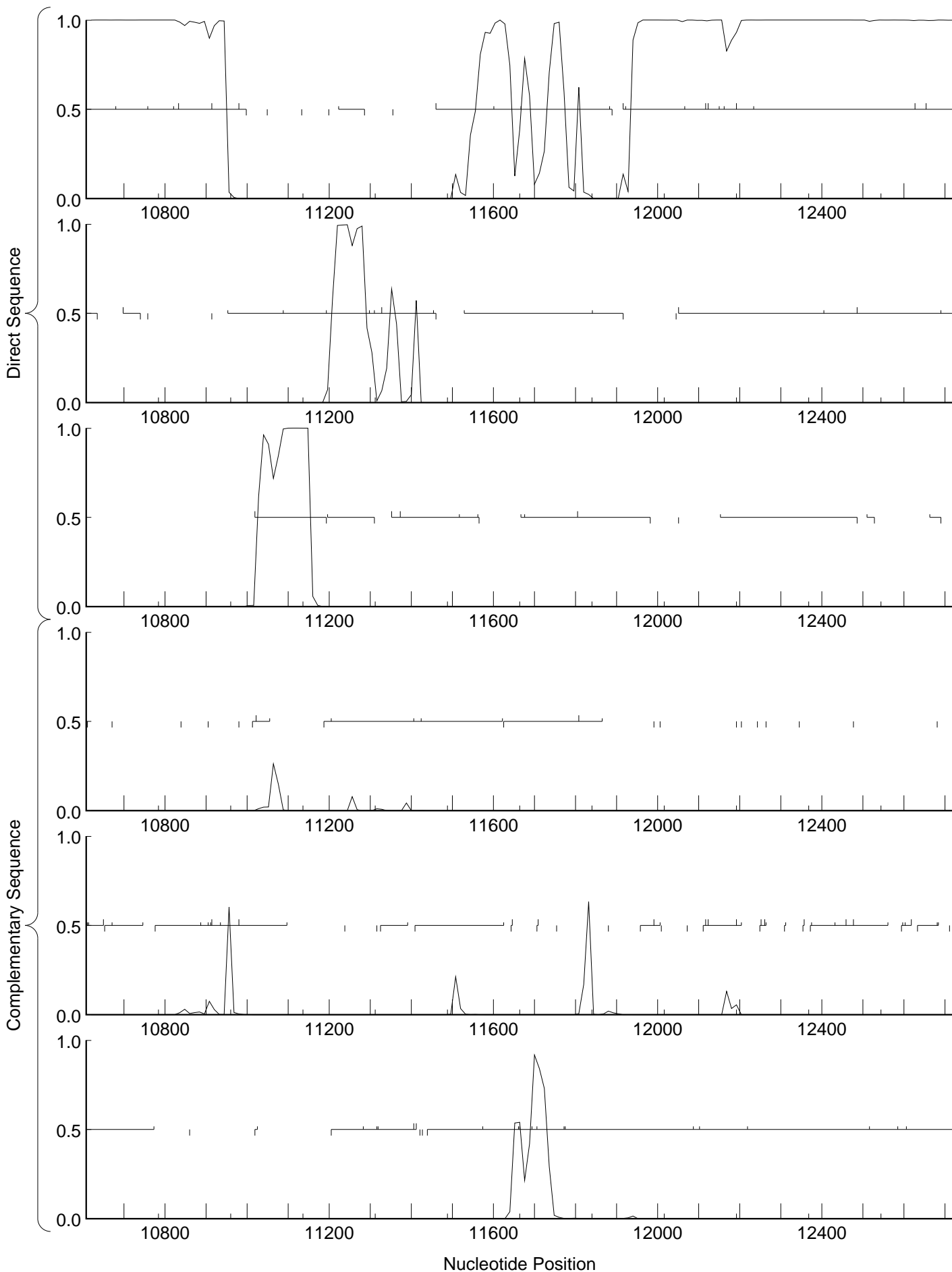


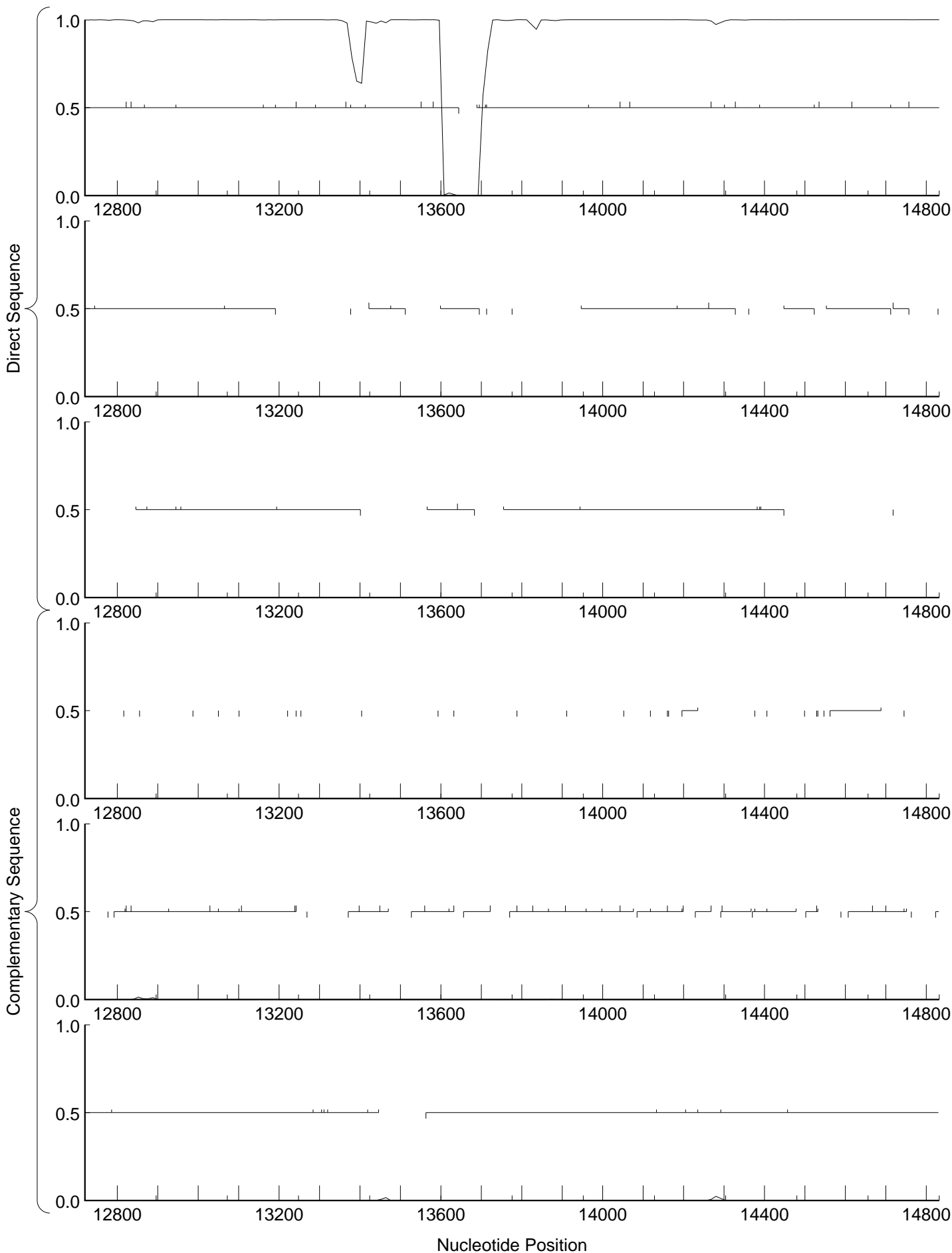
a complete sequence, 65869 bp including 10-base 3' overhang (CGCCGCGTAC), Cluster CR, Order 4, Window 96, Step 12, 5/33

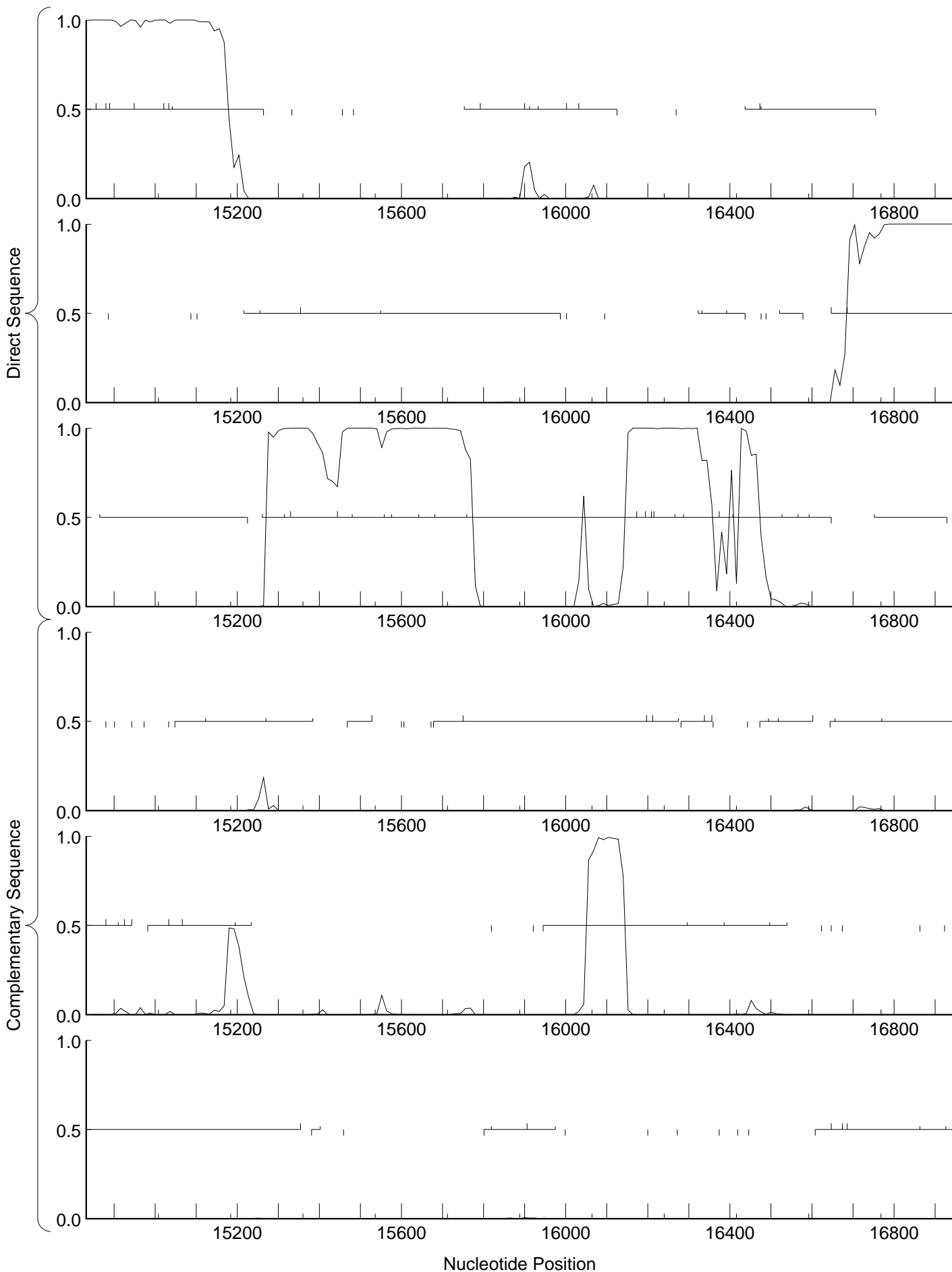


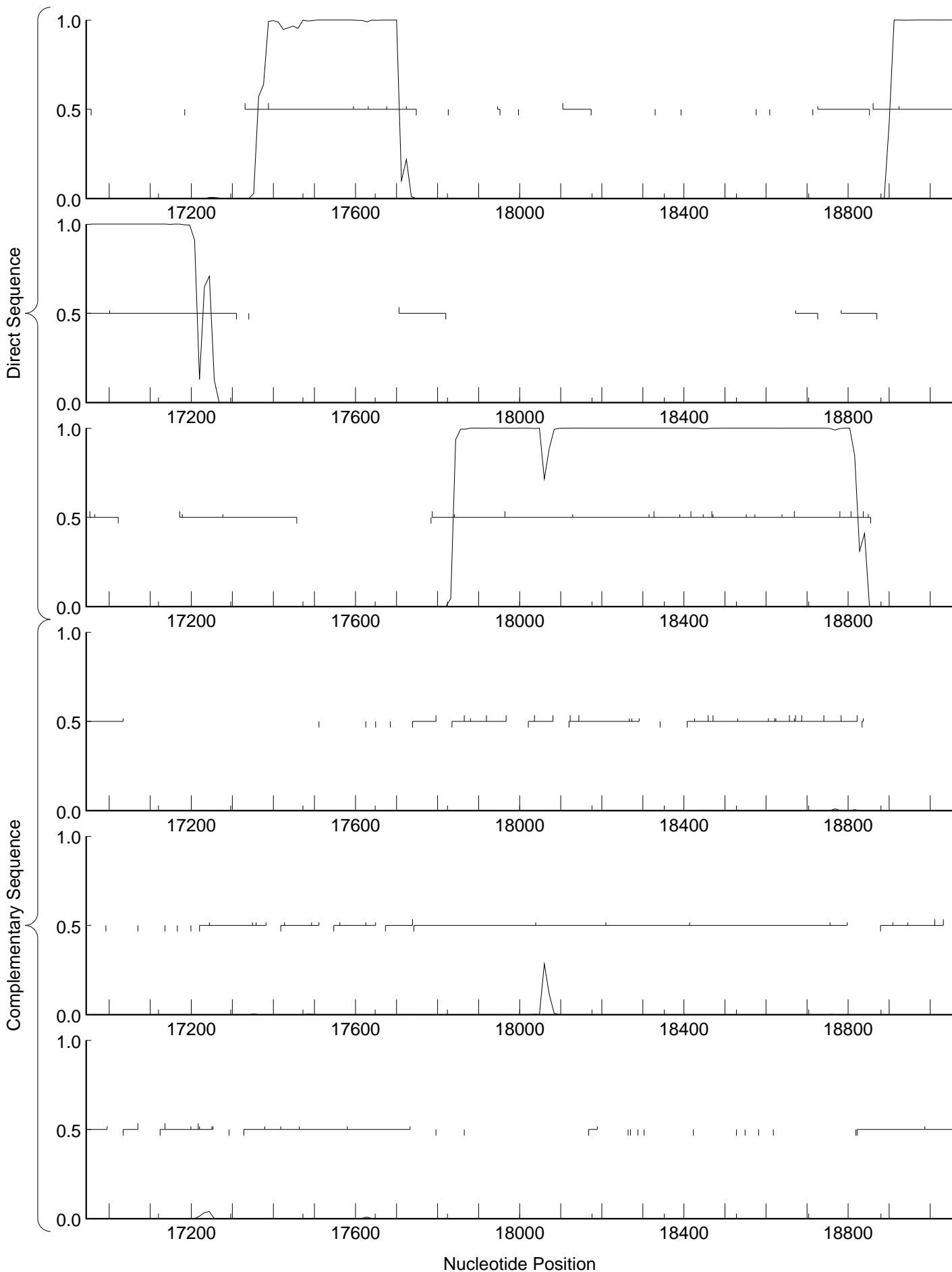


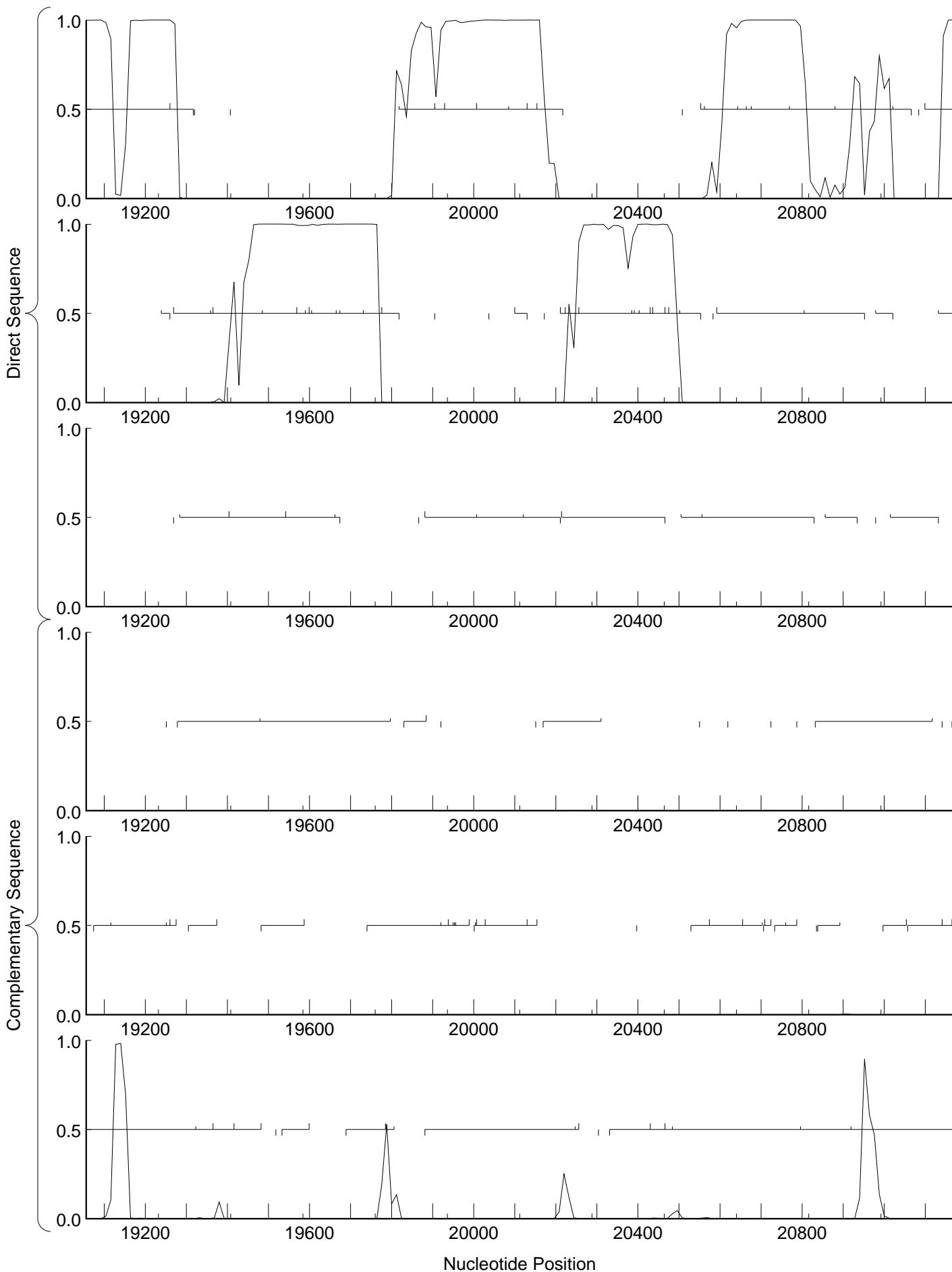
a complete sequence, 65869 bp including 10-base 3' overhang (CGCCGCGTAC), Cluster CR, Order 4, Window 96, Step 12, 7/33

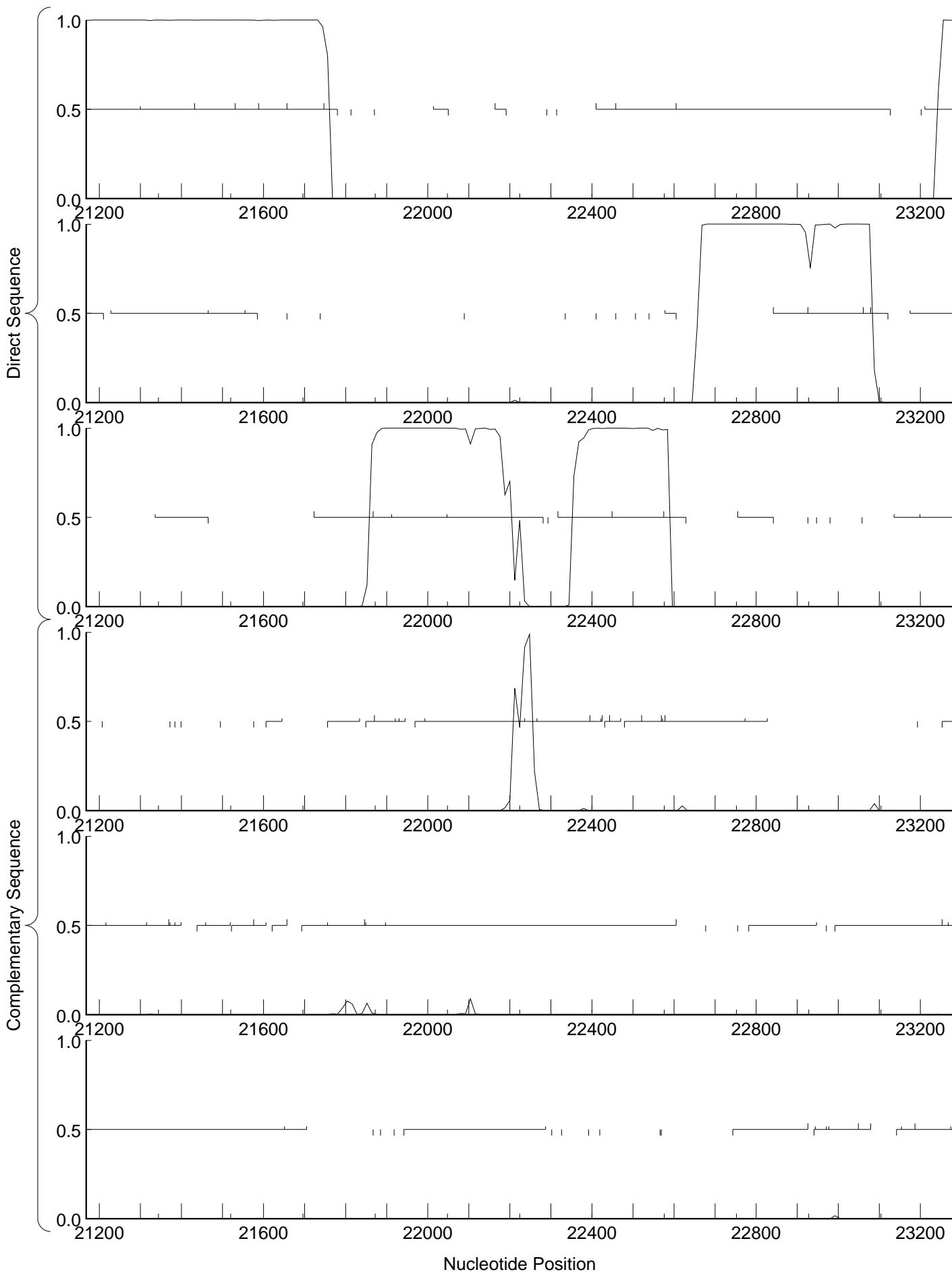


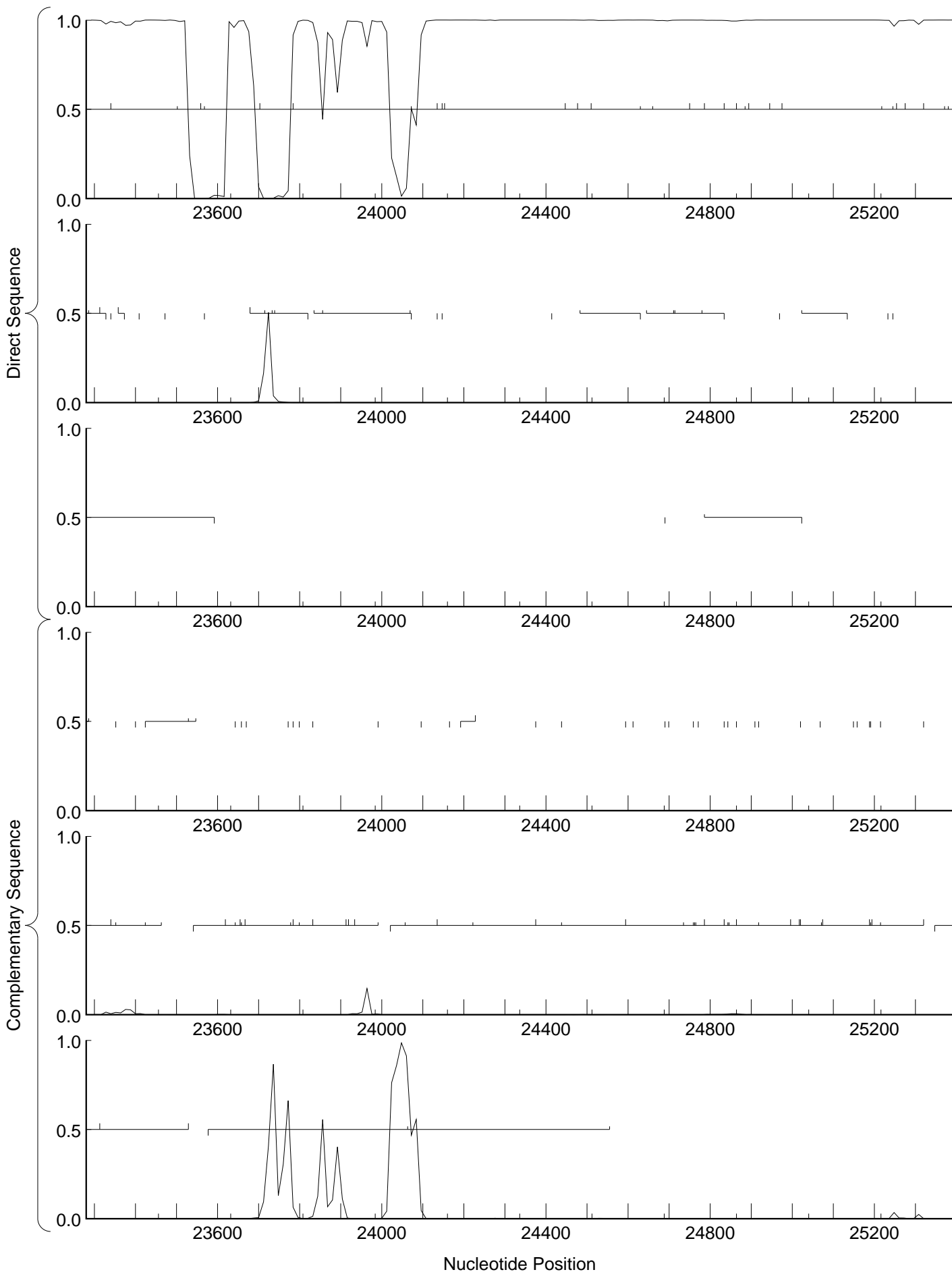


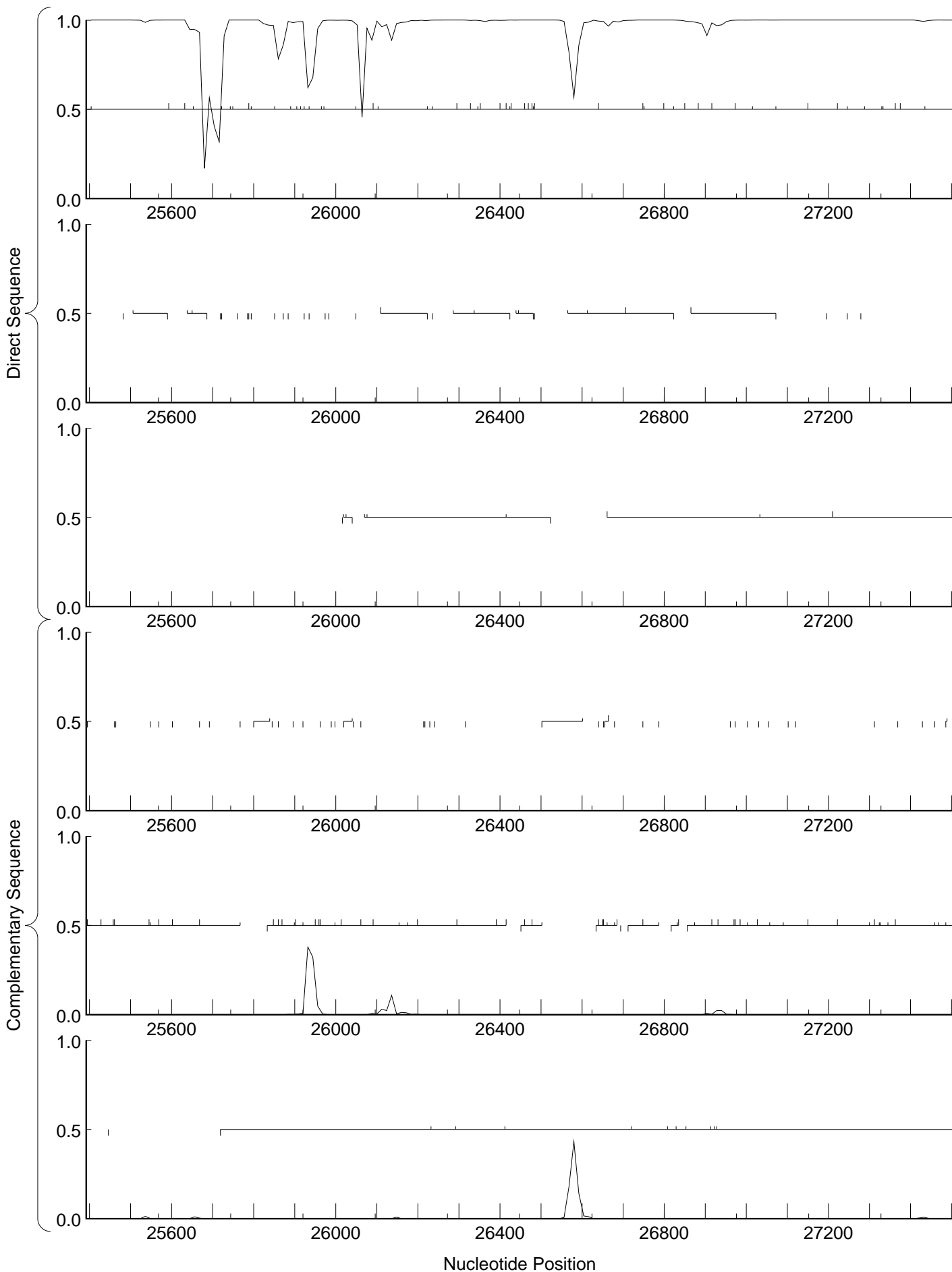


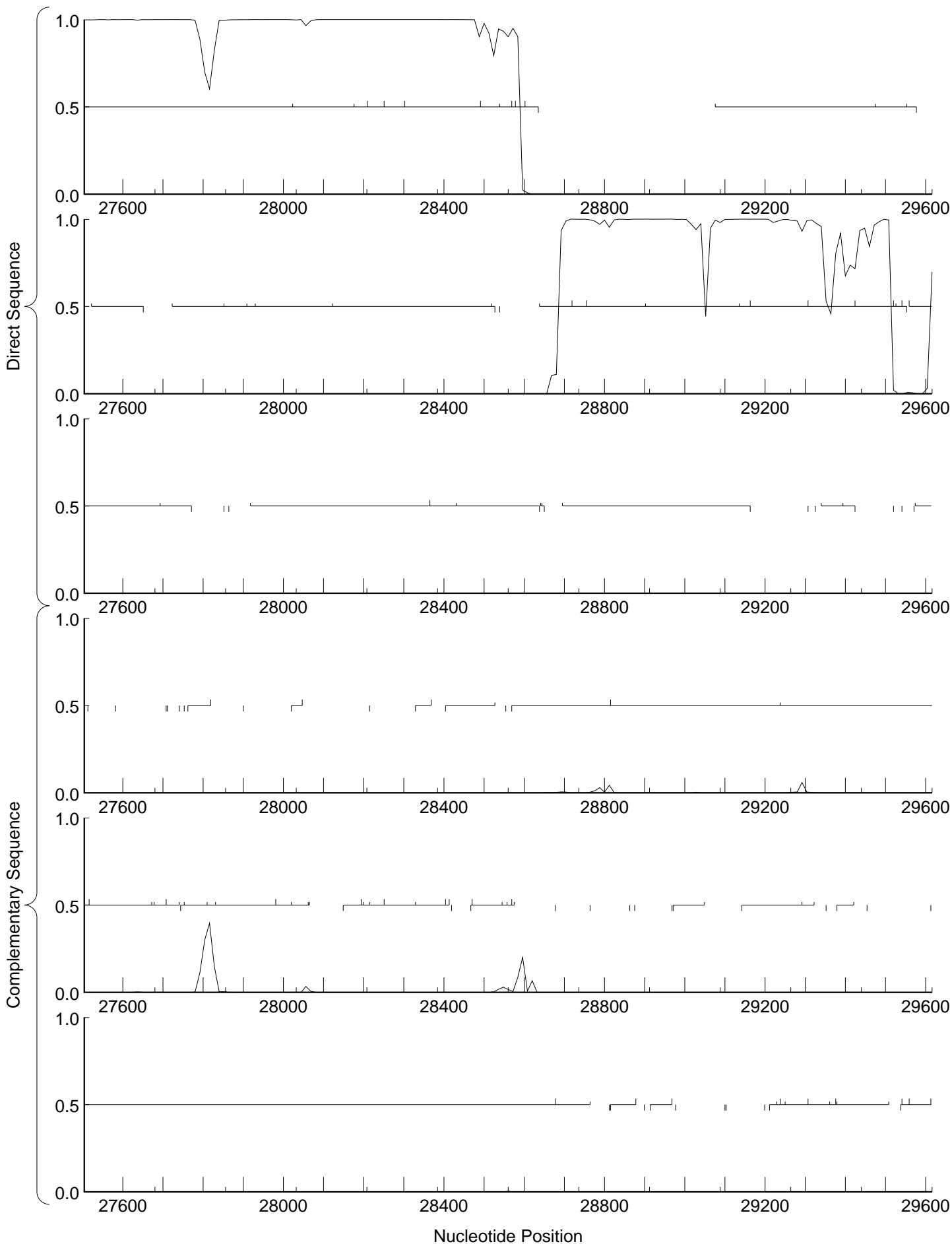


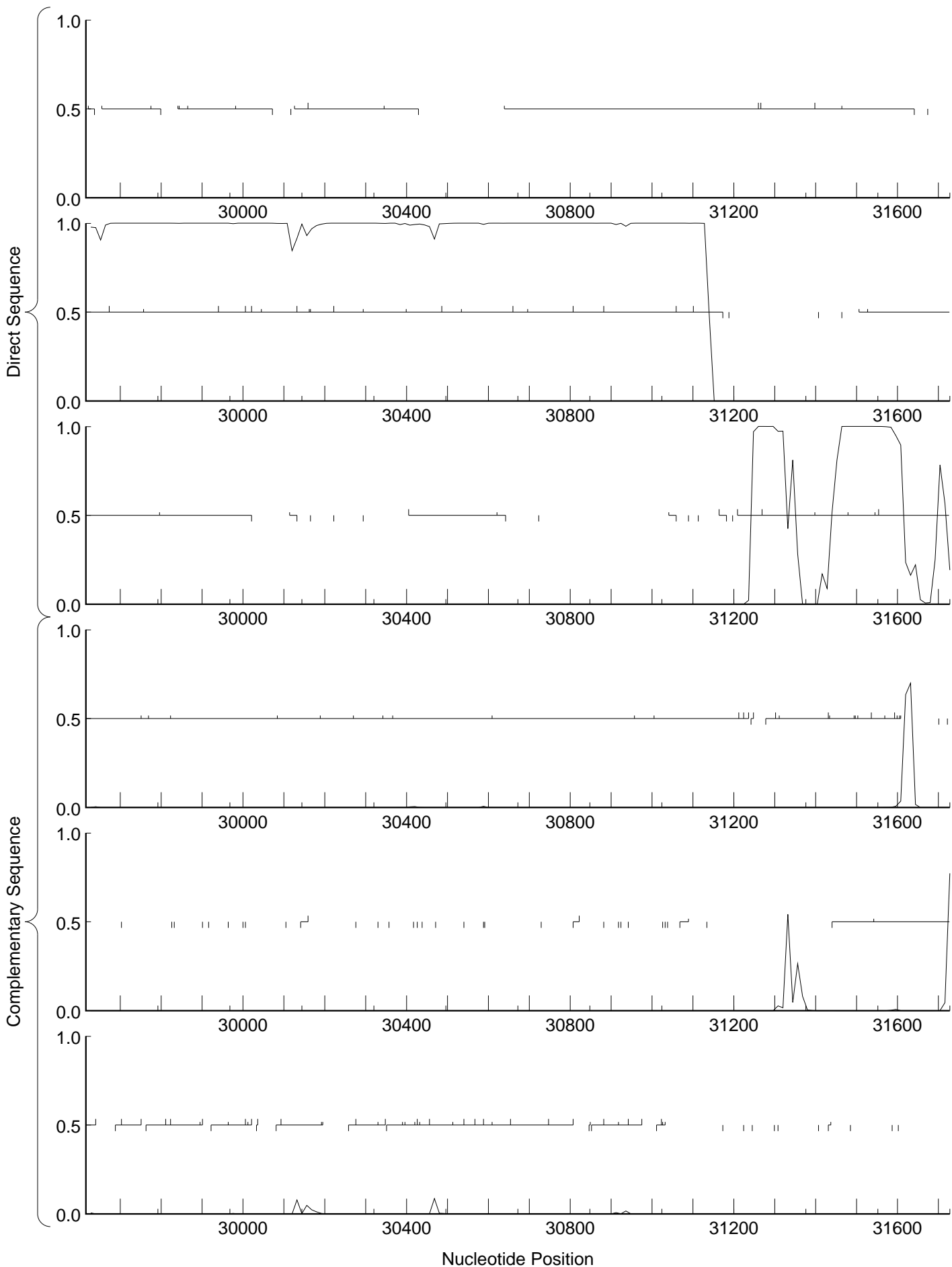


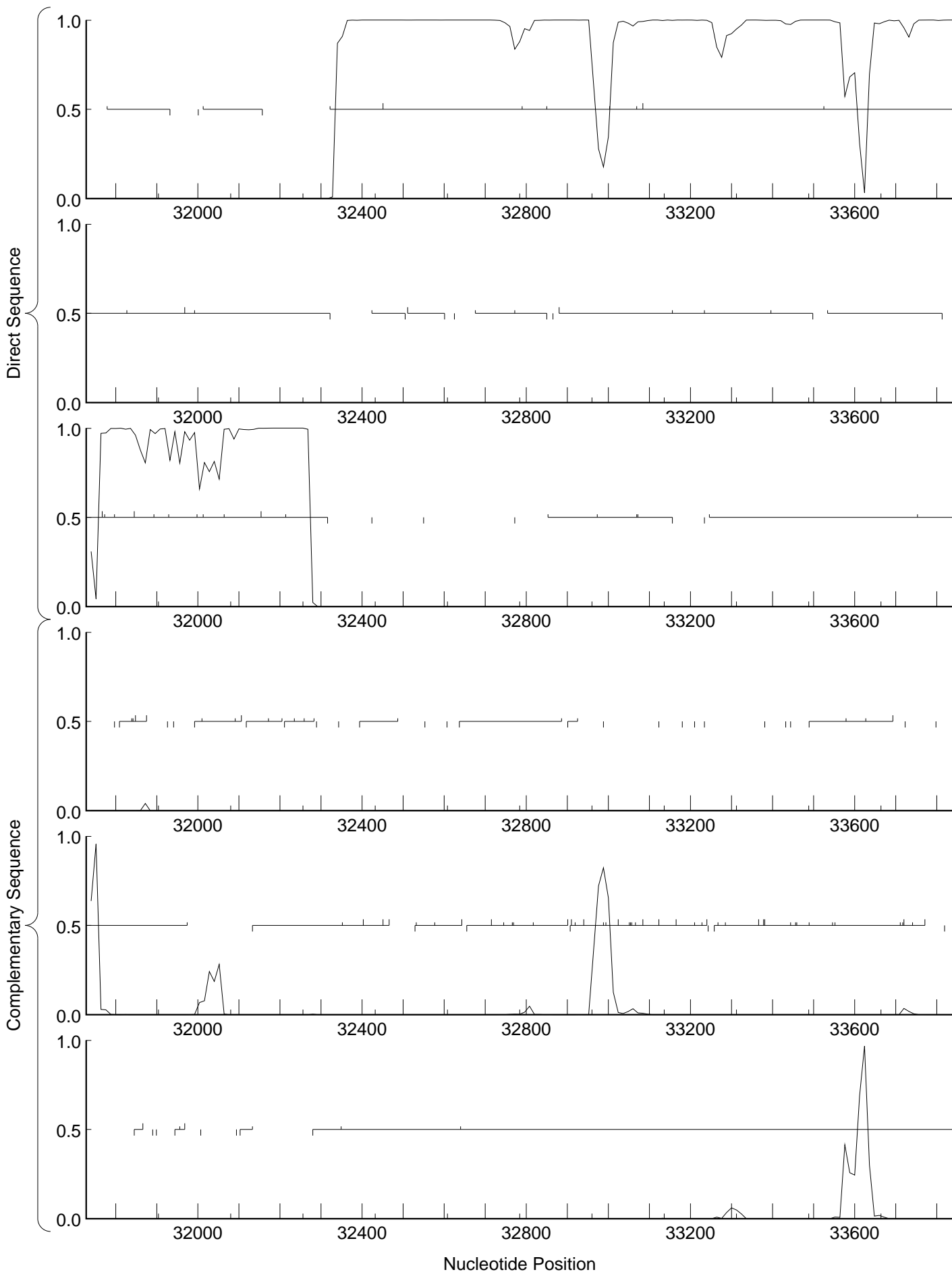












complete sequence, 65869 bp including 10-base 3' overhang (CGCCGCGTAC), Cluster CR, Order 4, Window 96, Step 12, 18/33

