

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Burley complete sequence, 60111 bp including 9-base 3' overhang (CGCCGCTCT),
Analysis Date : 5/16/25 at 11:16:16
Pages : 30
Sequence Length : 60111 bp
GC Content : 51.61%

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,

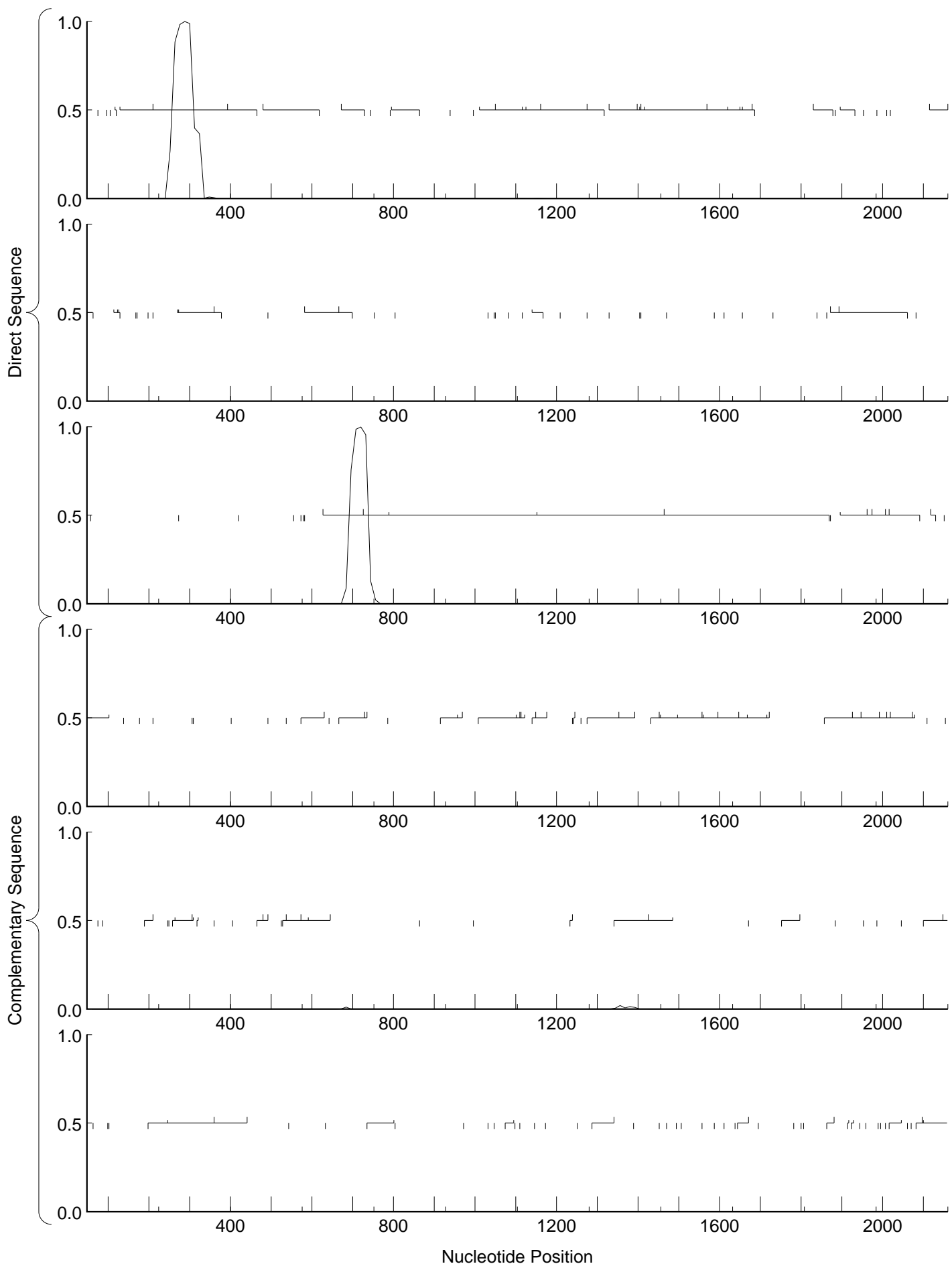
Matrix notes & comments

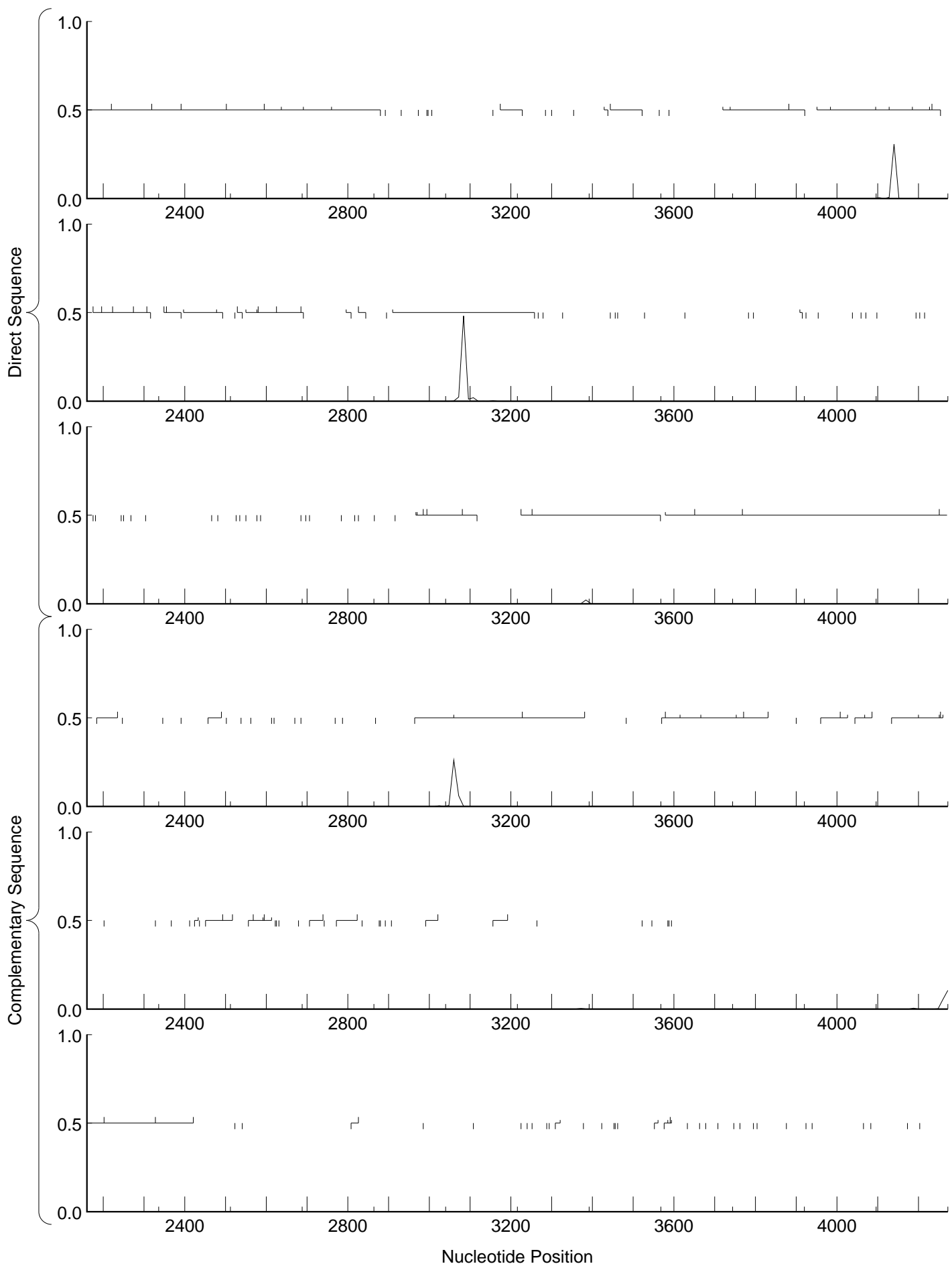
Training set derived by GeneMarkS, 4.30 October 2014
Sat Sep 8 00:36:14 2018

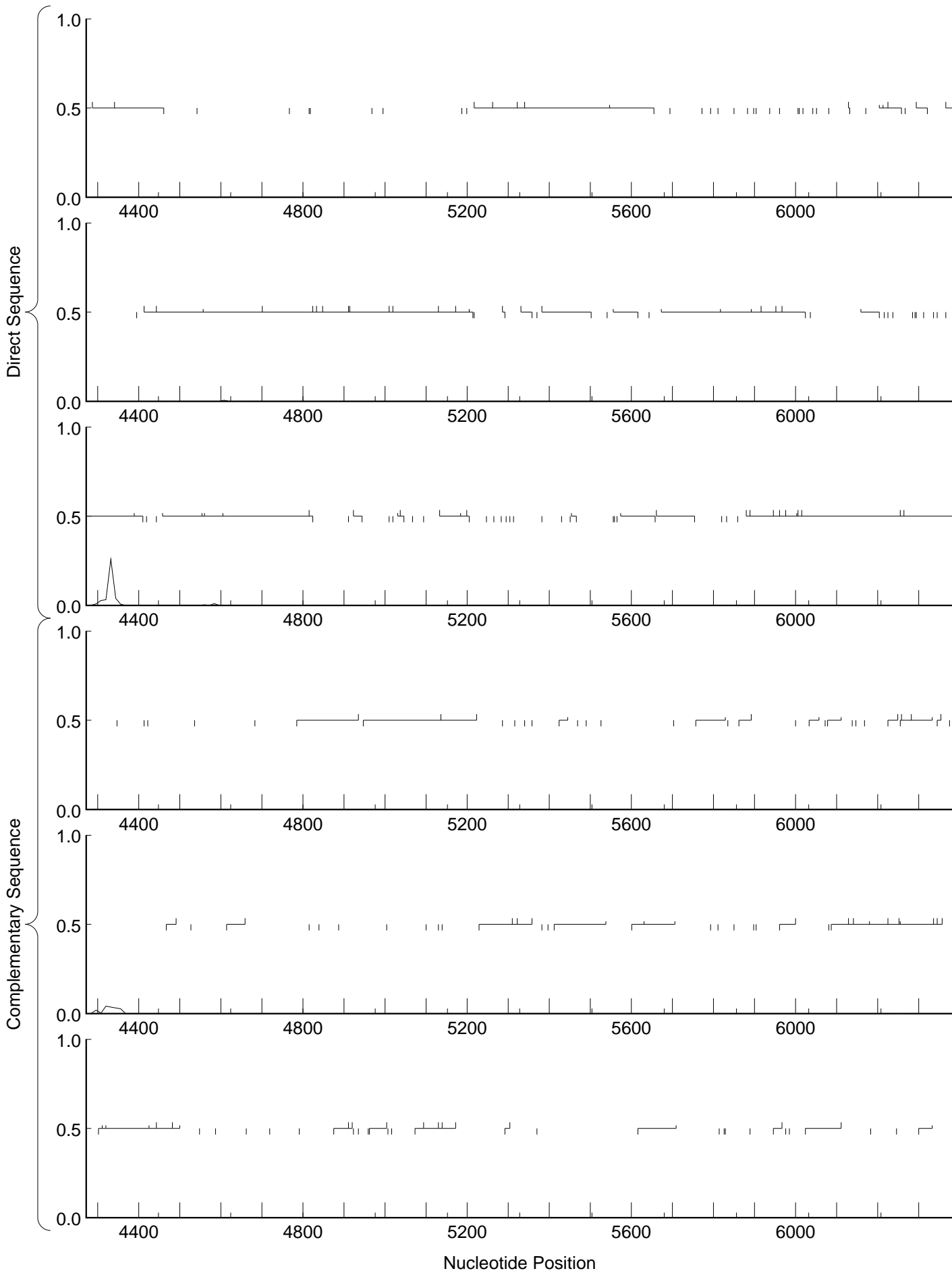
MATRIX INFORMATION

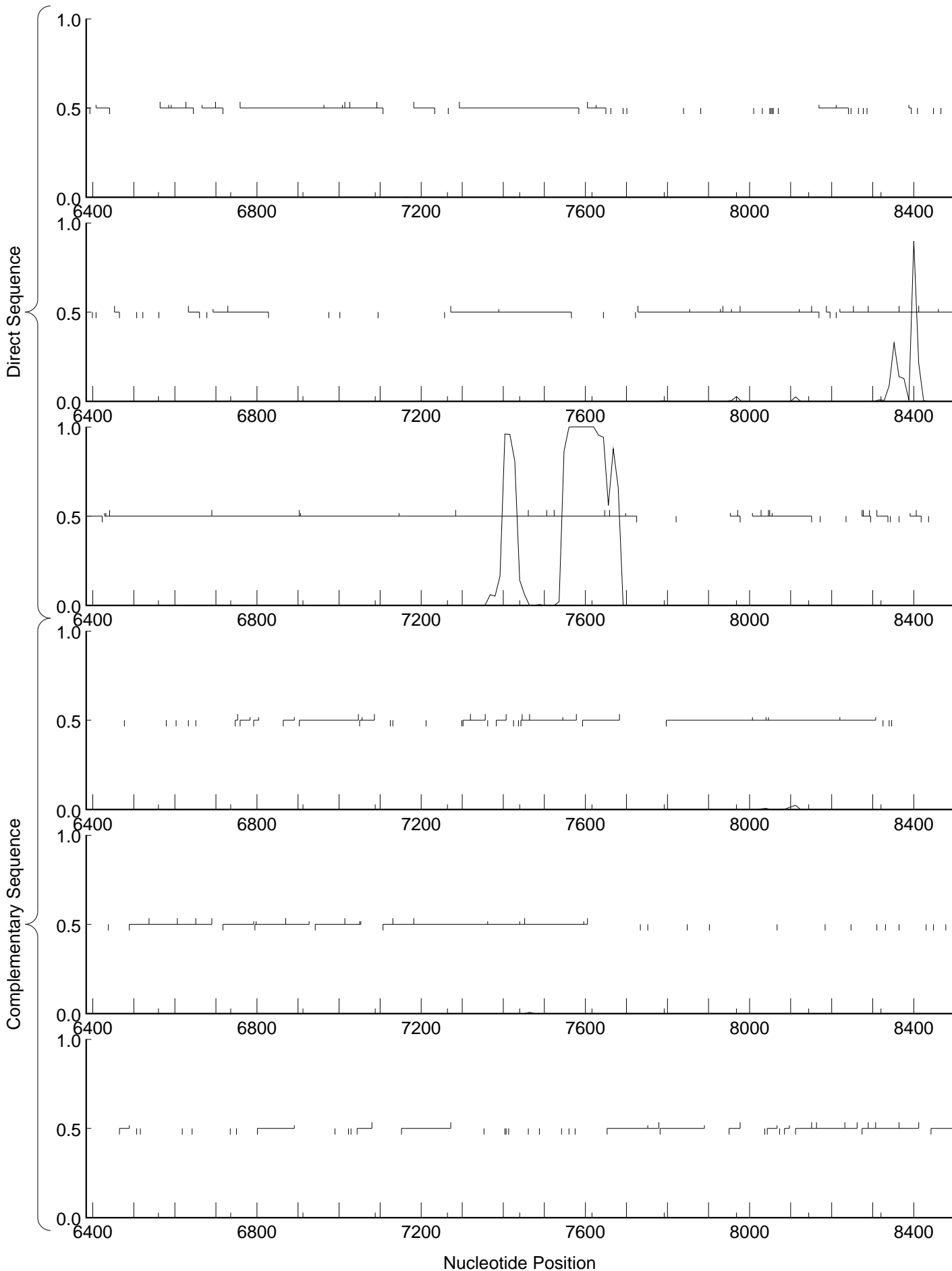
Matrix : -
Author : -
Order : 4

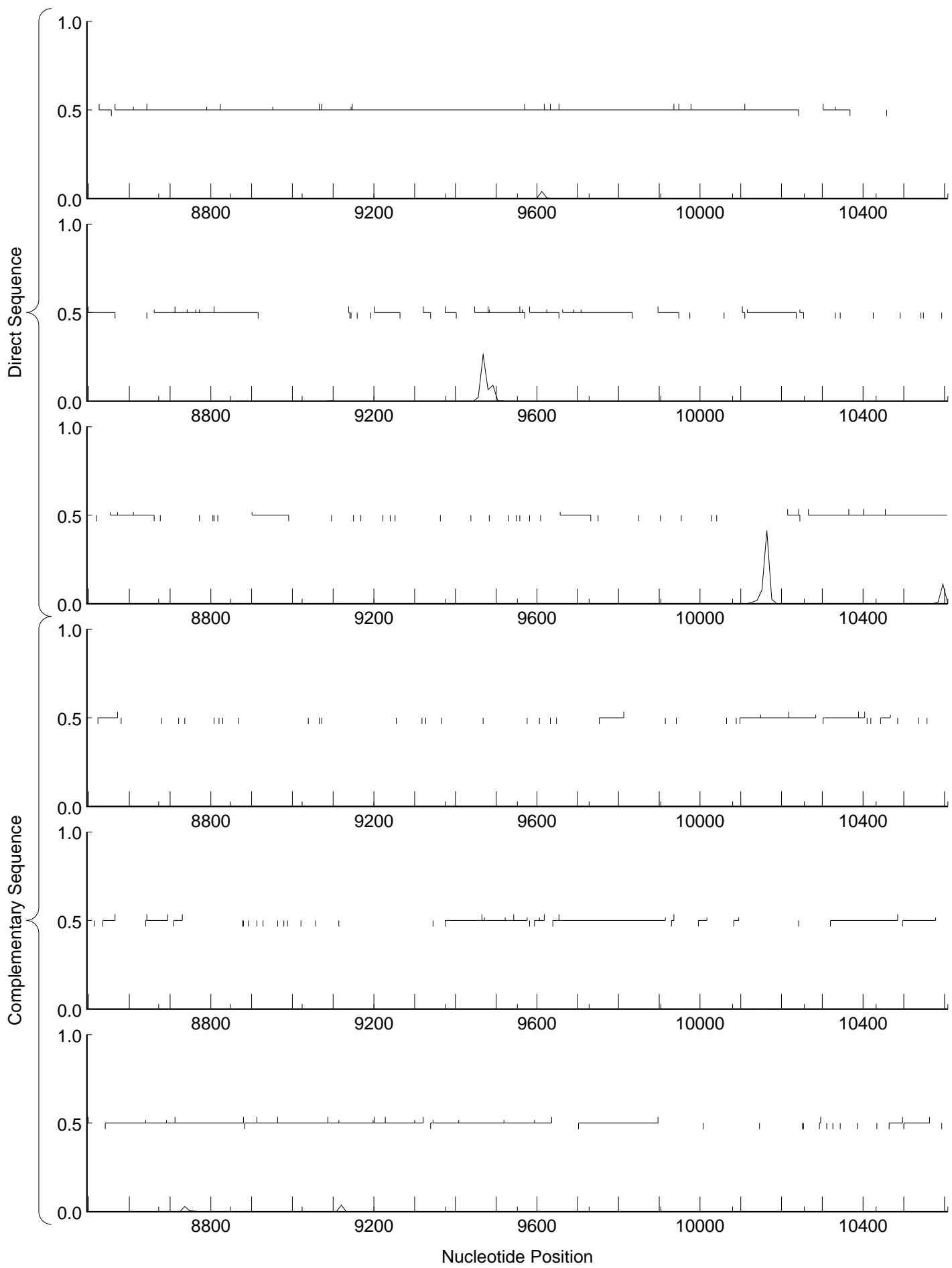
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

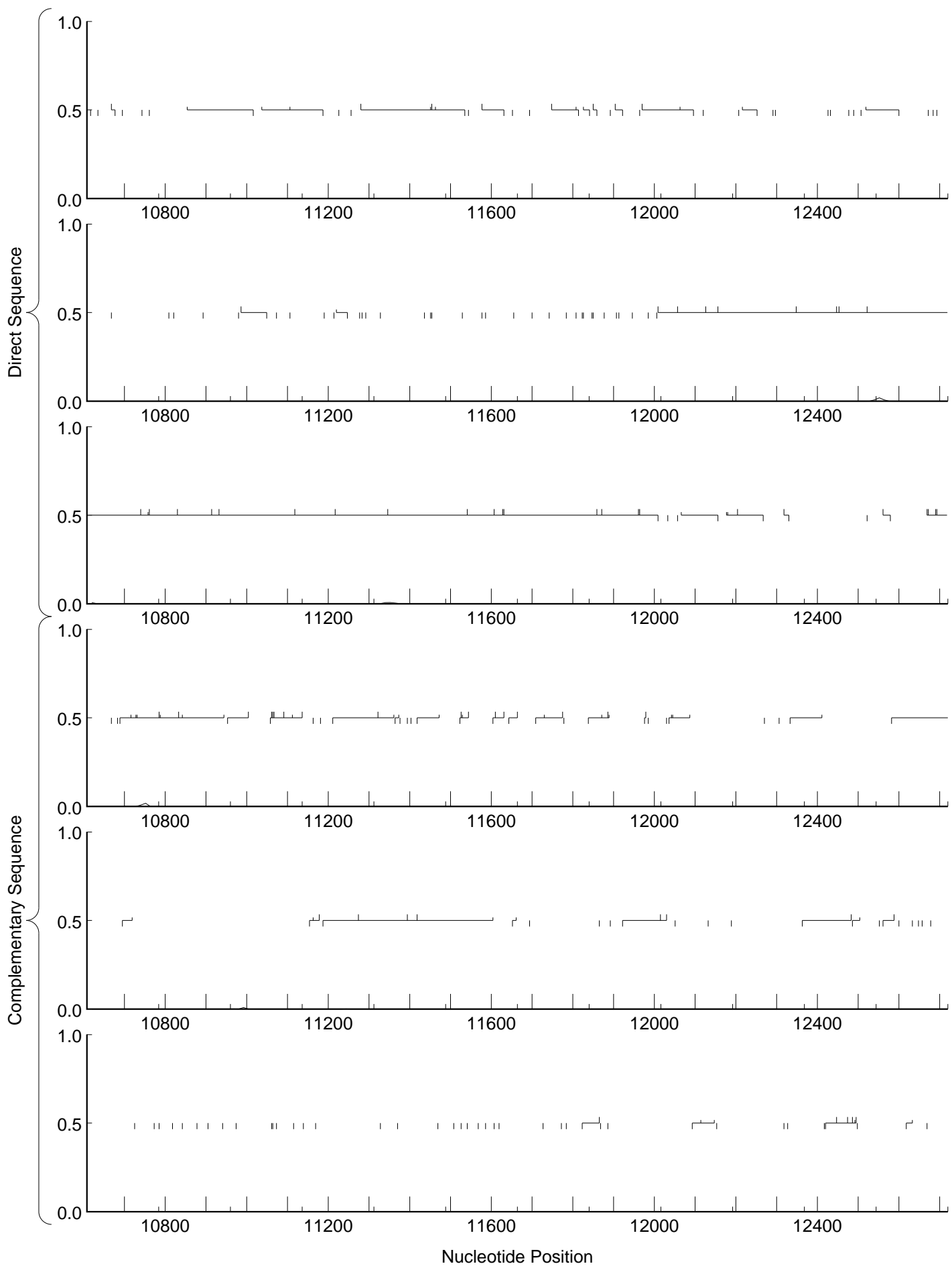


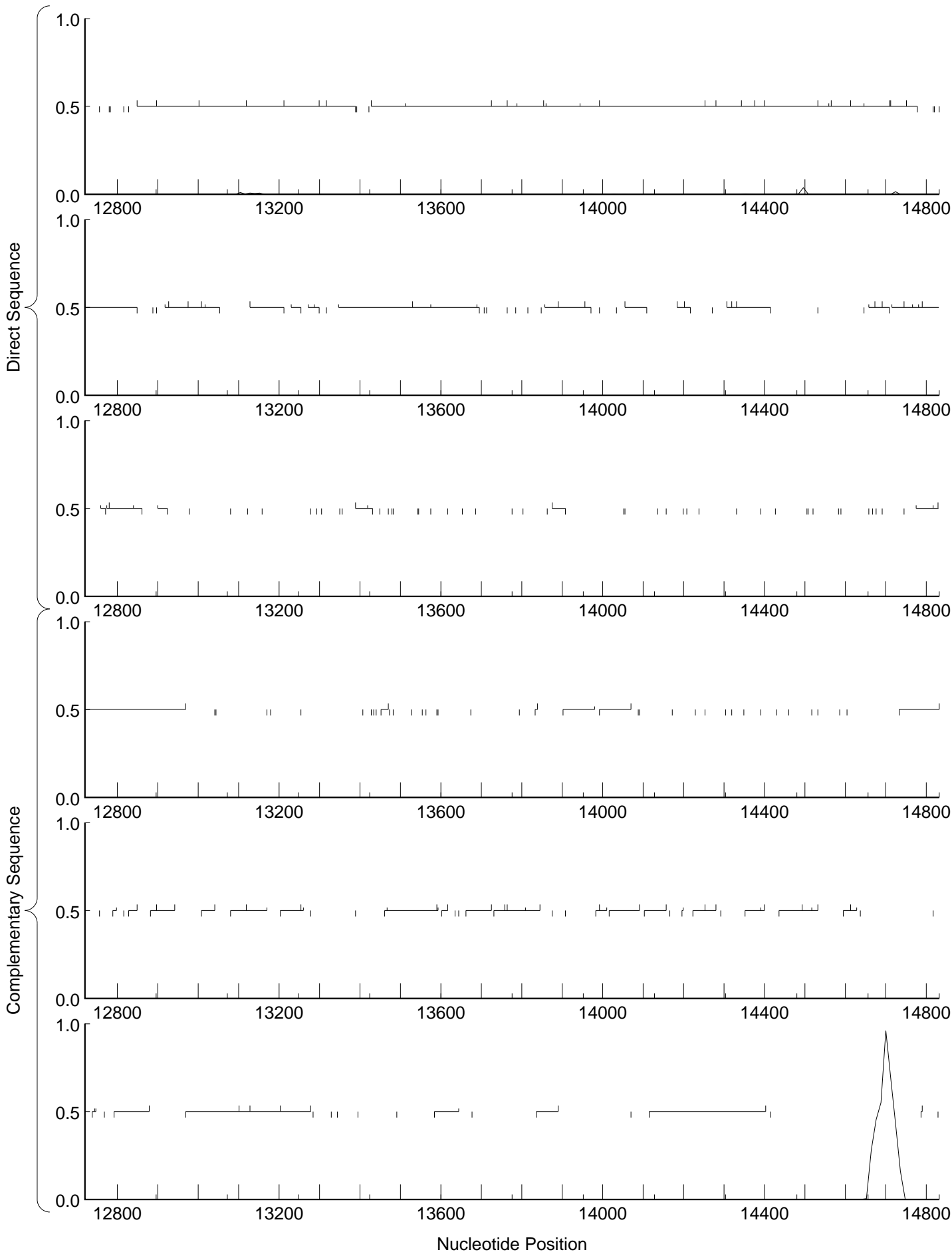


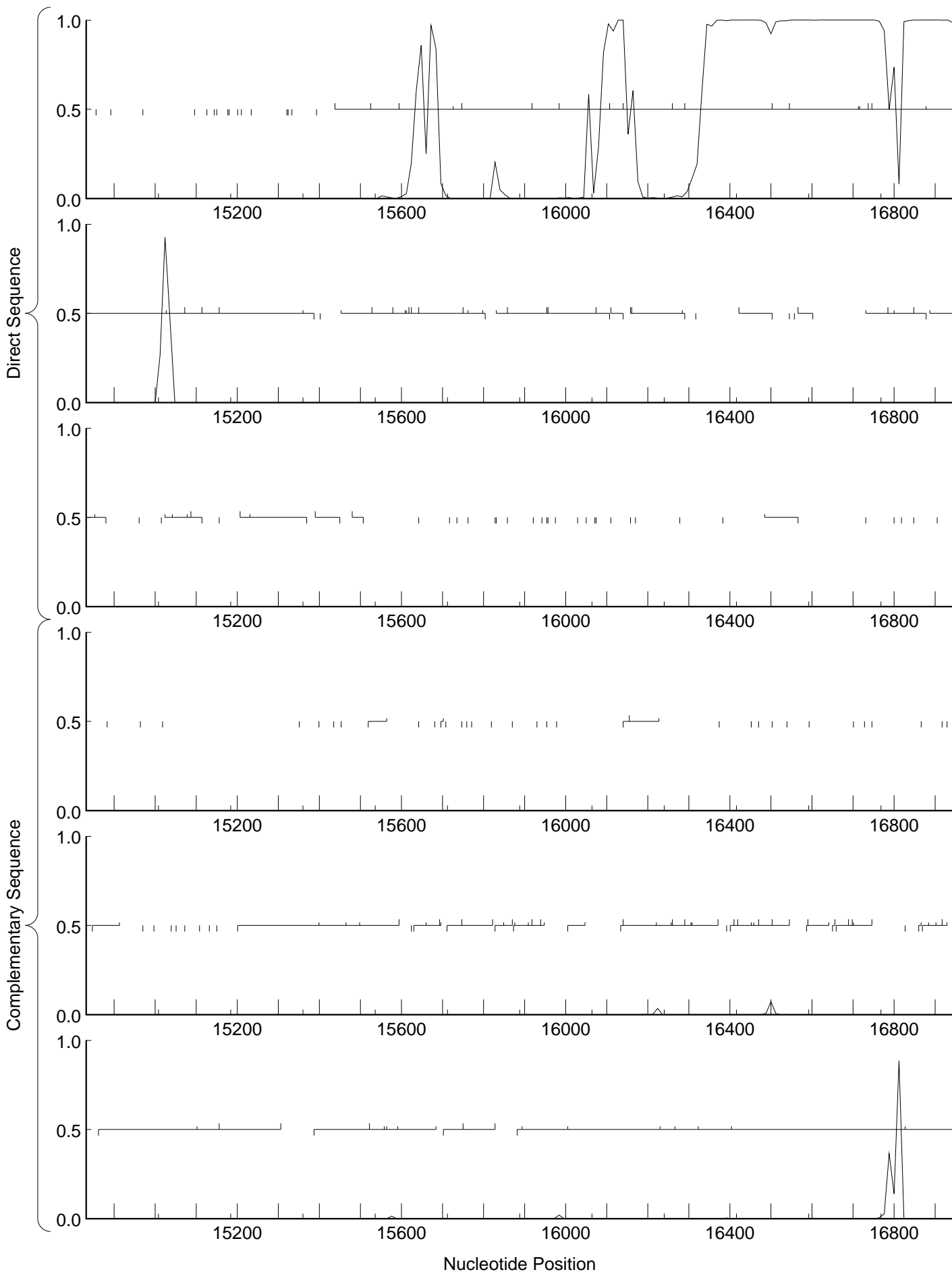


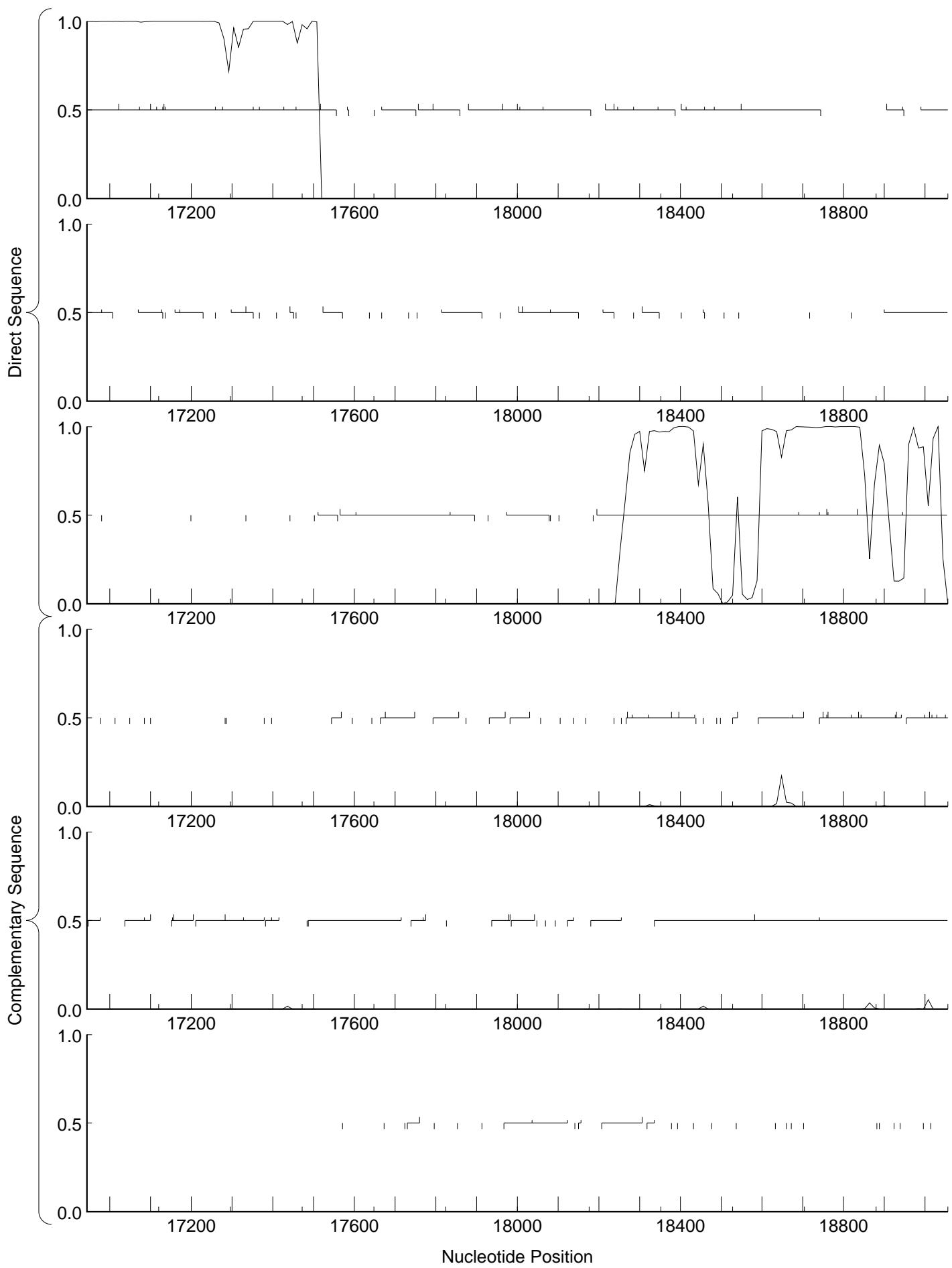


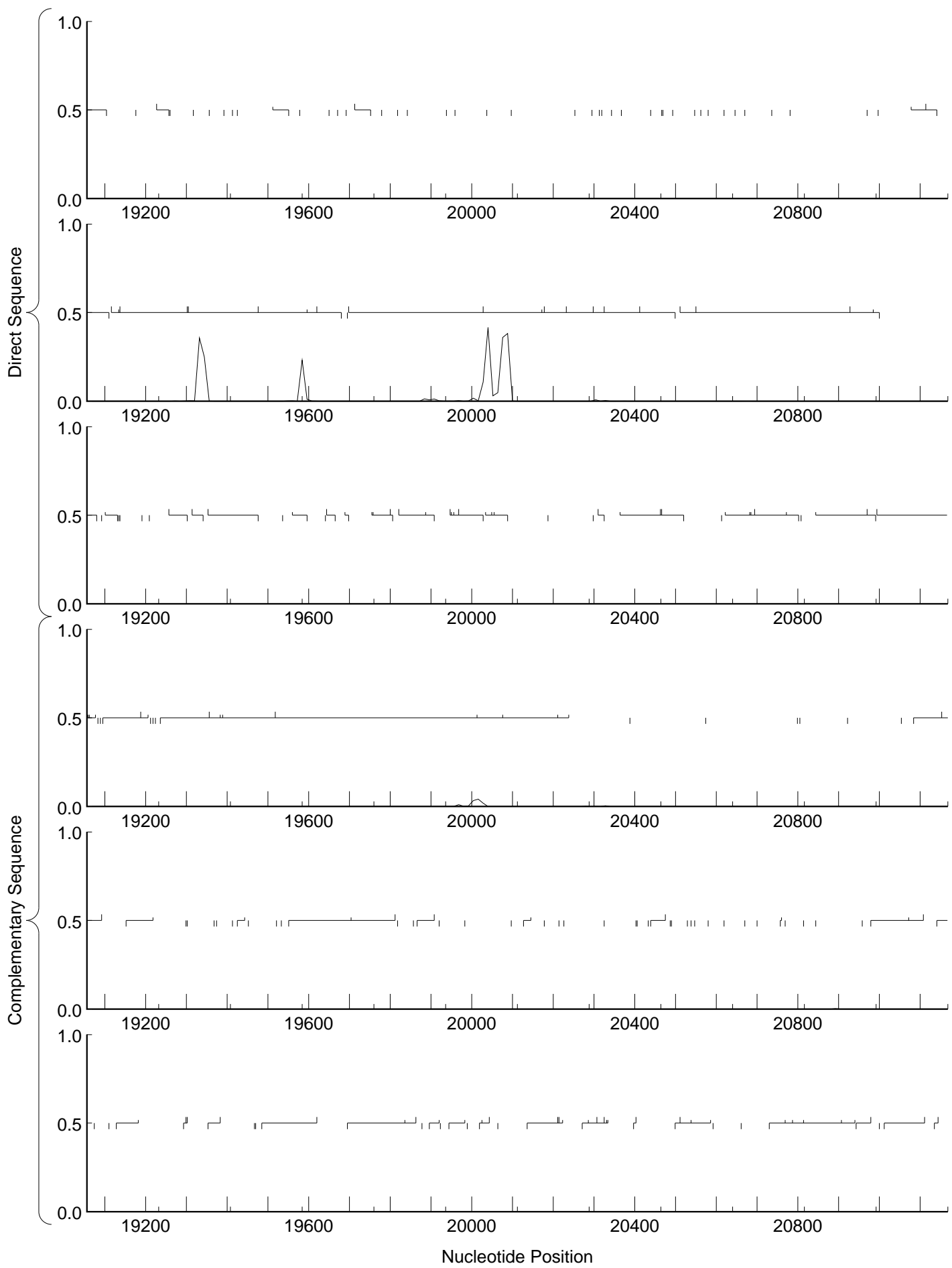


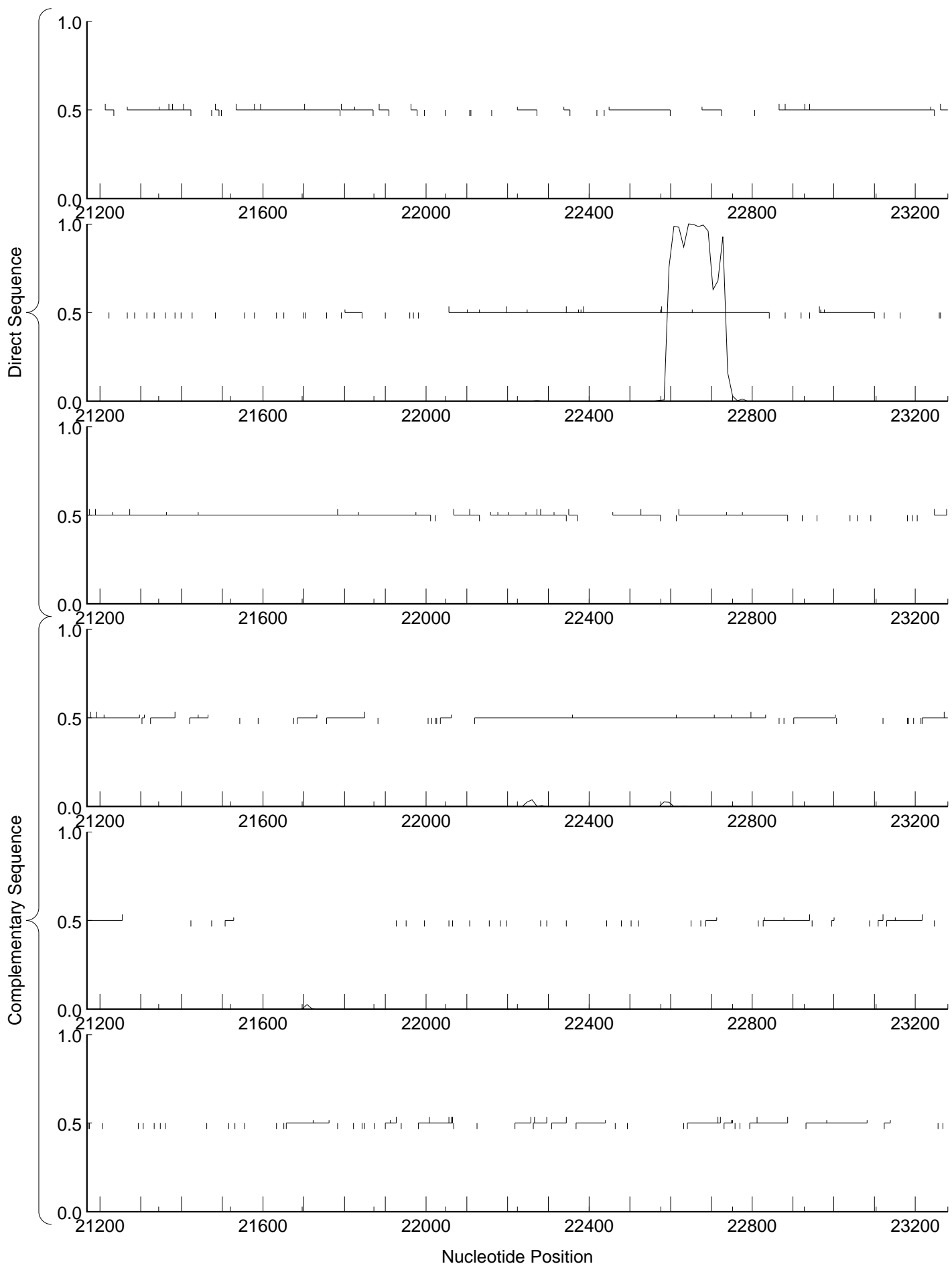


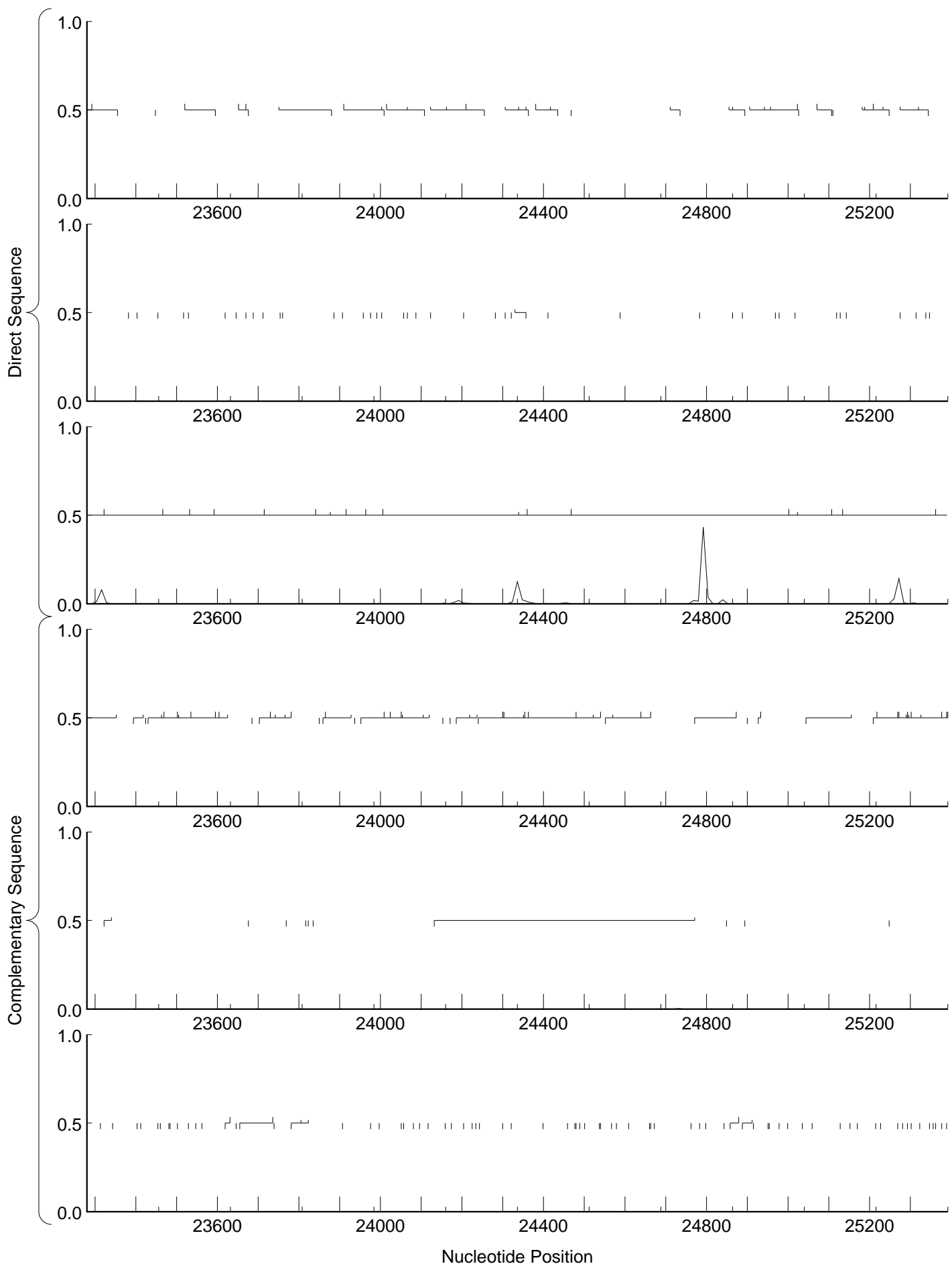


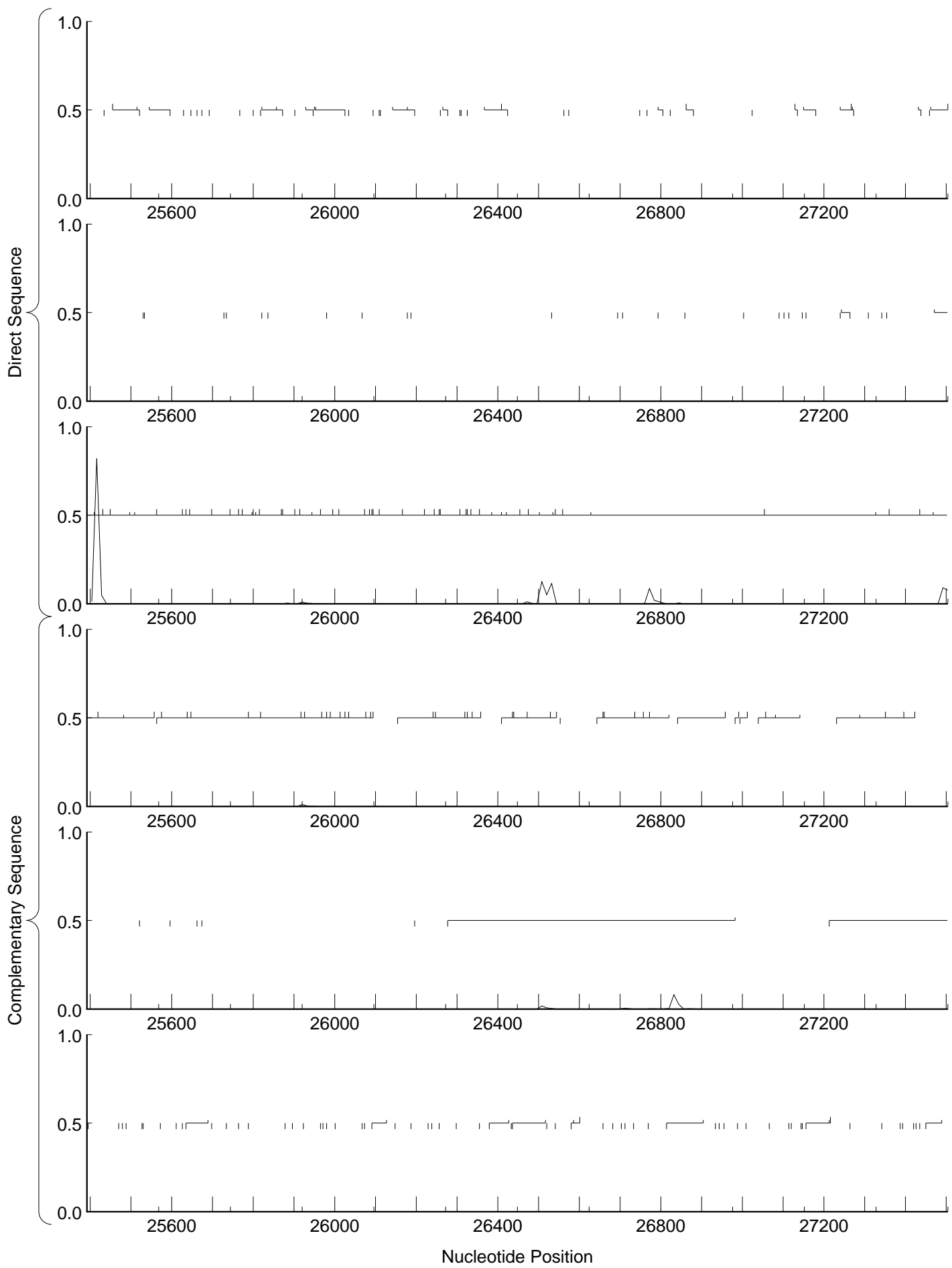


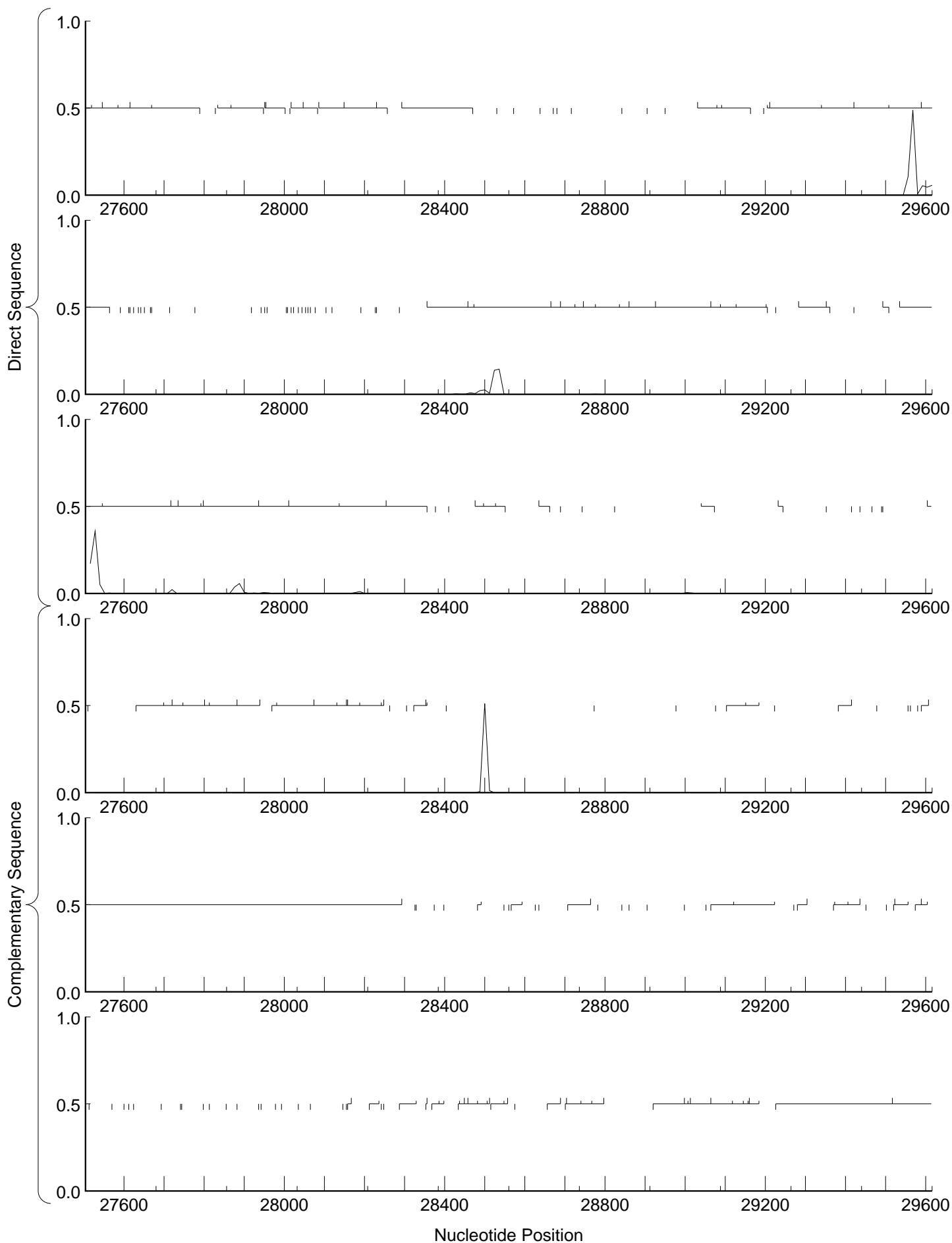


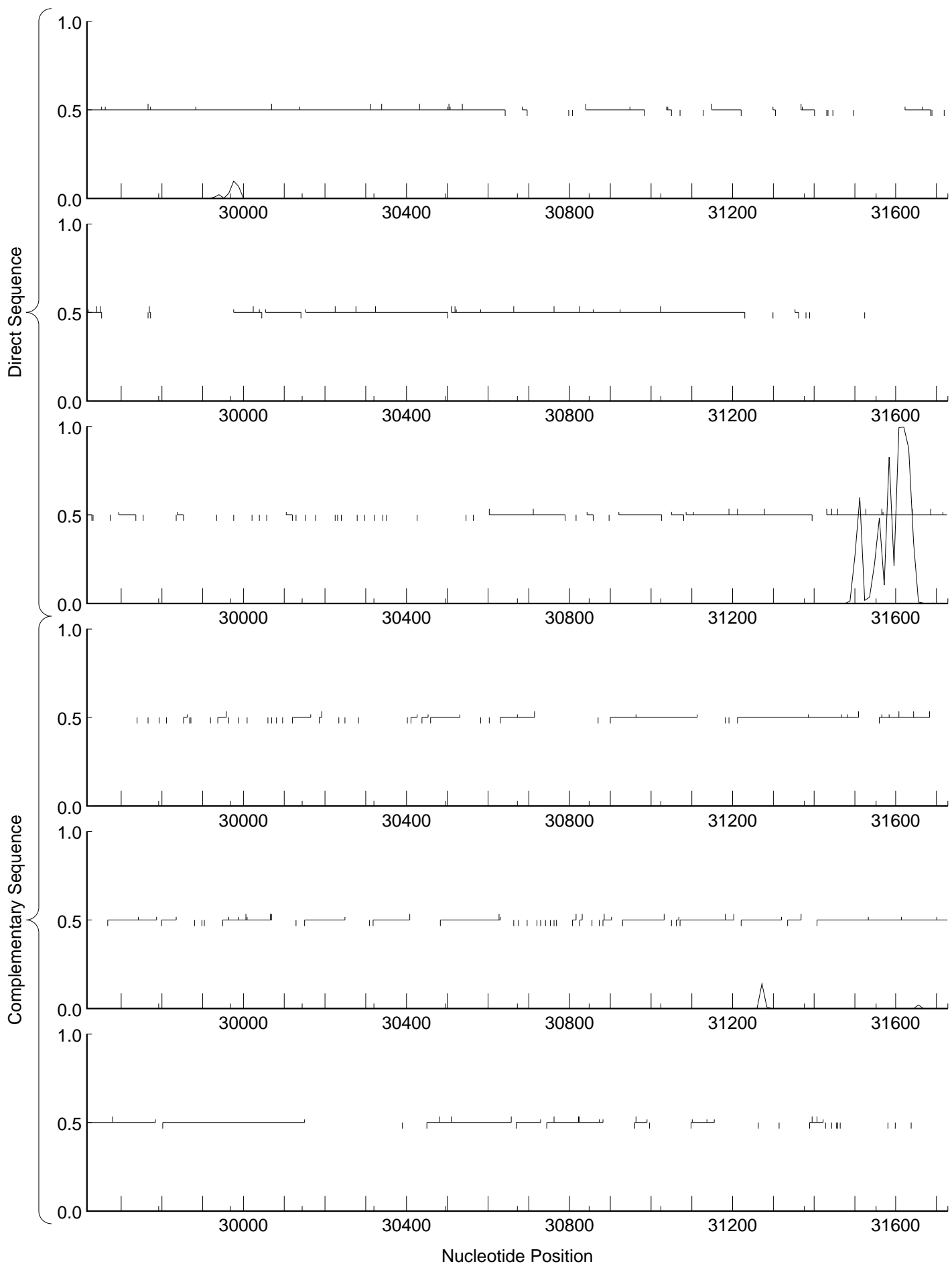


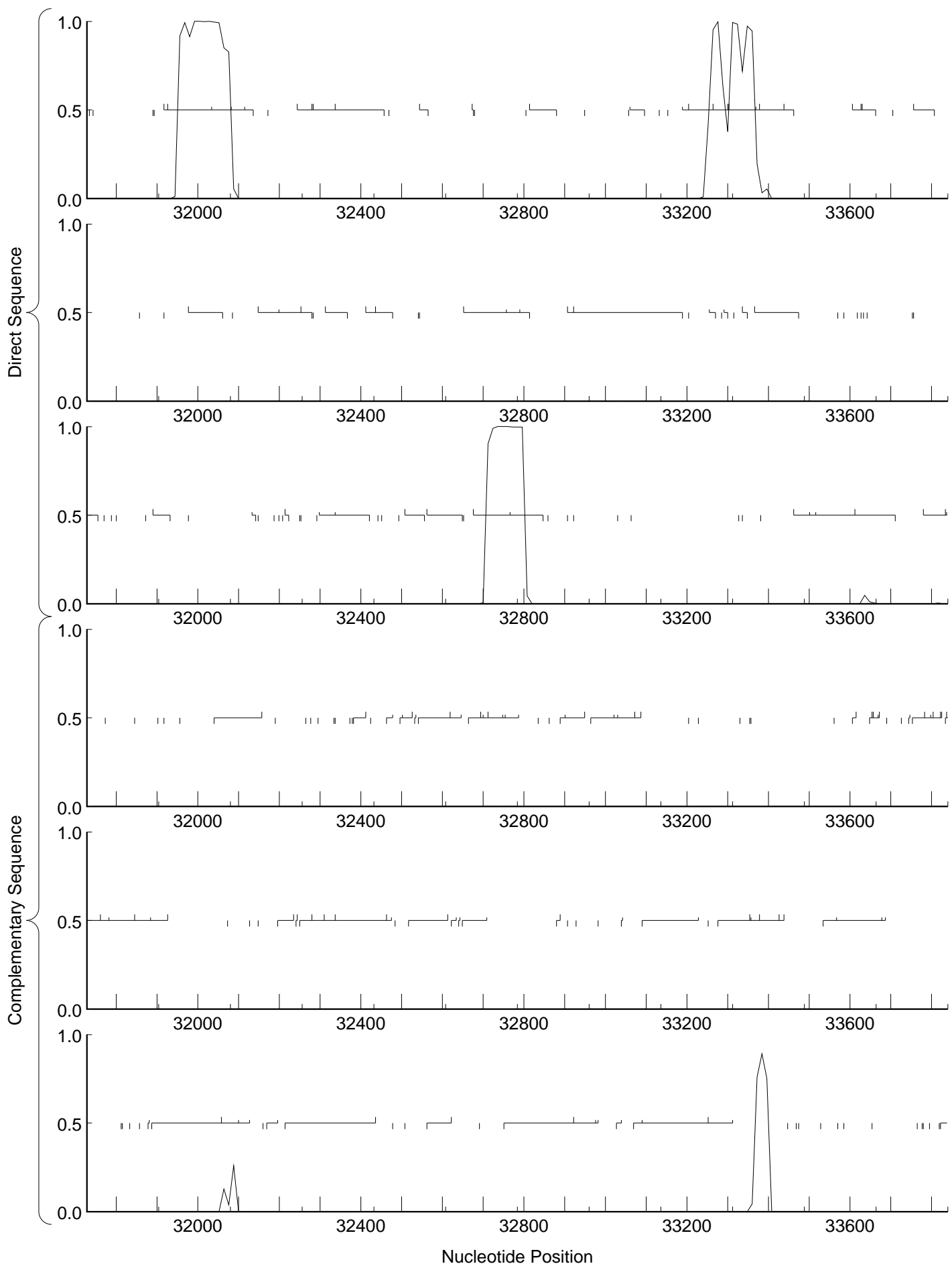


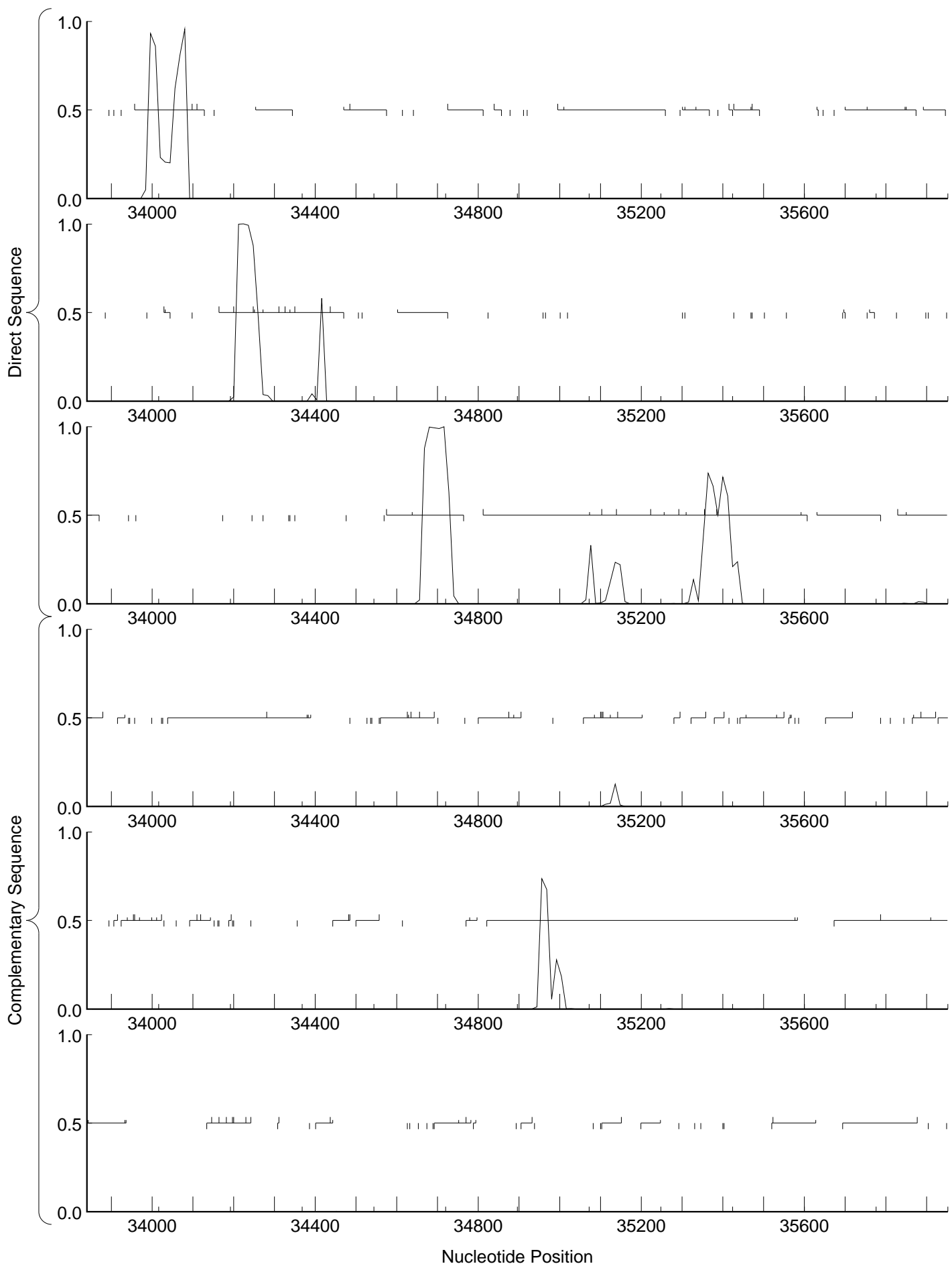




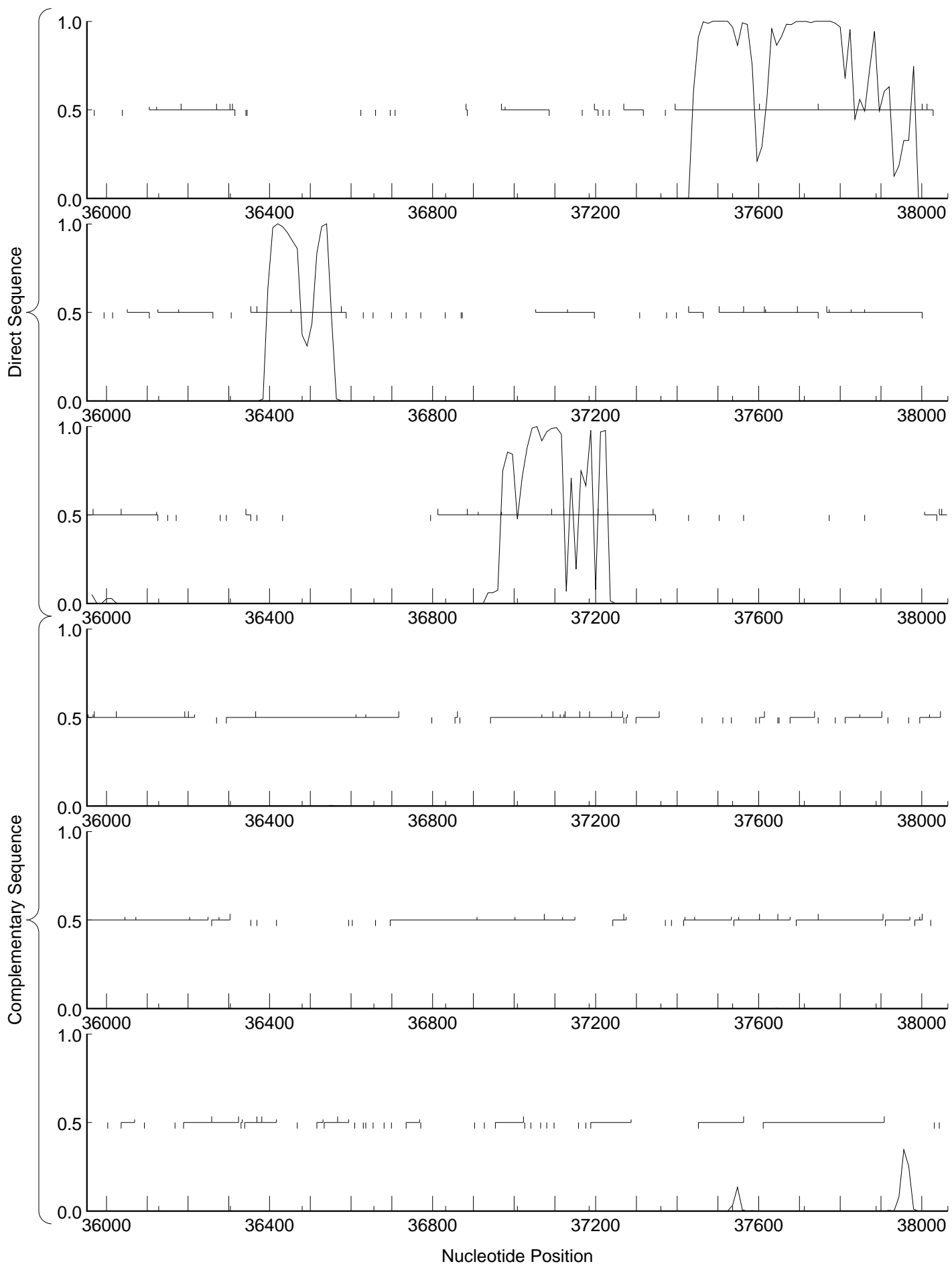


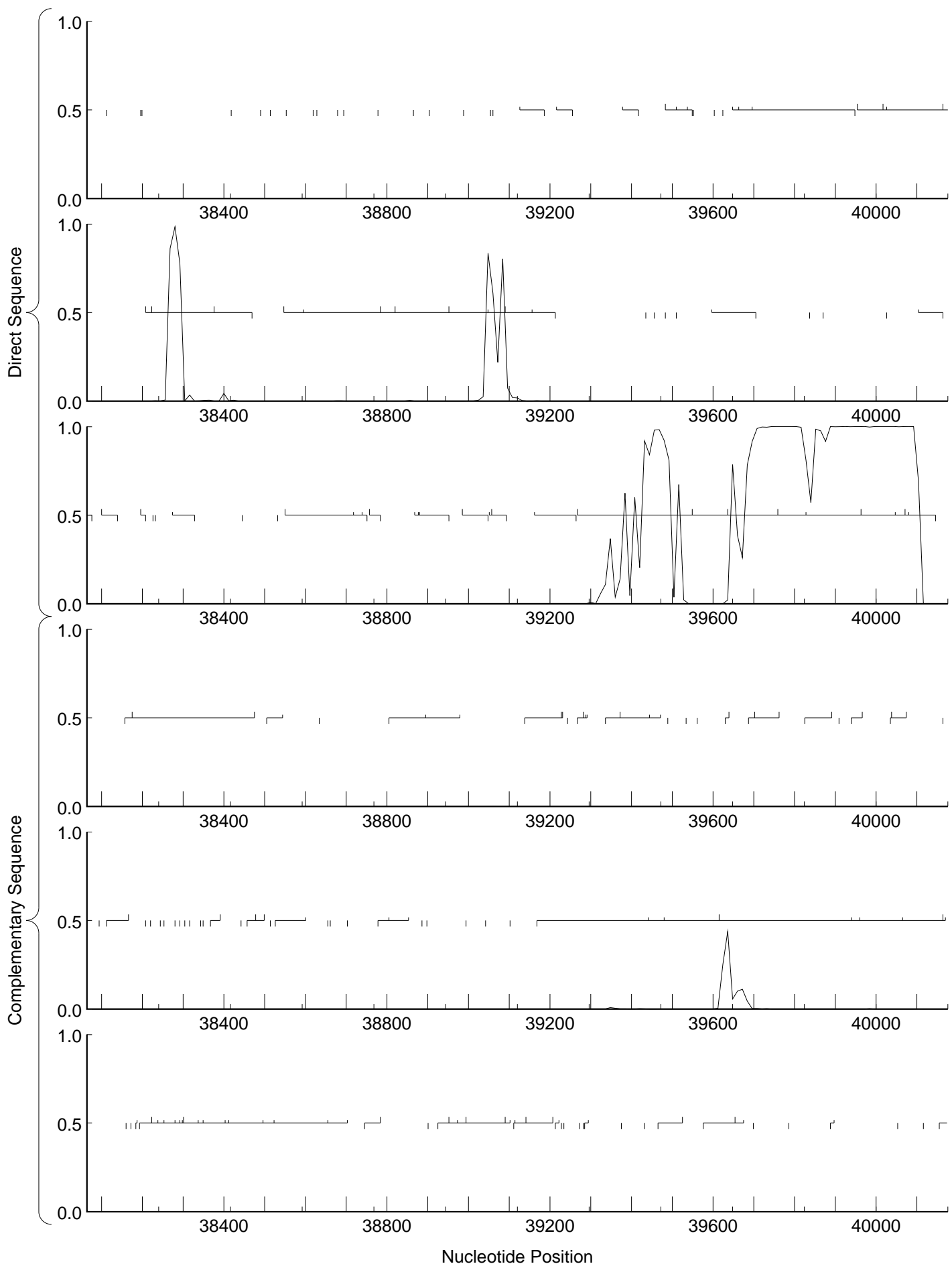


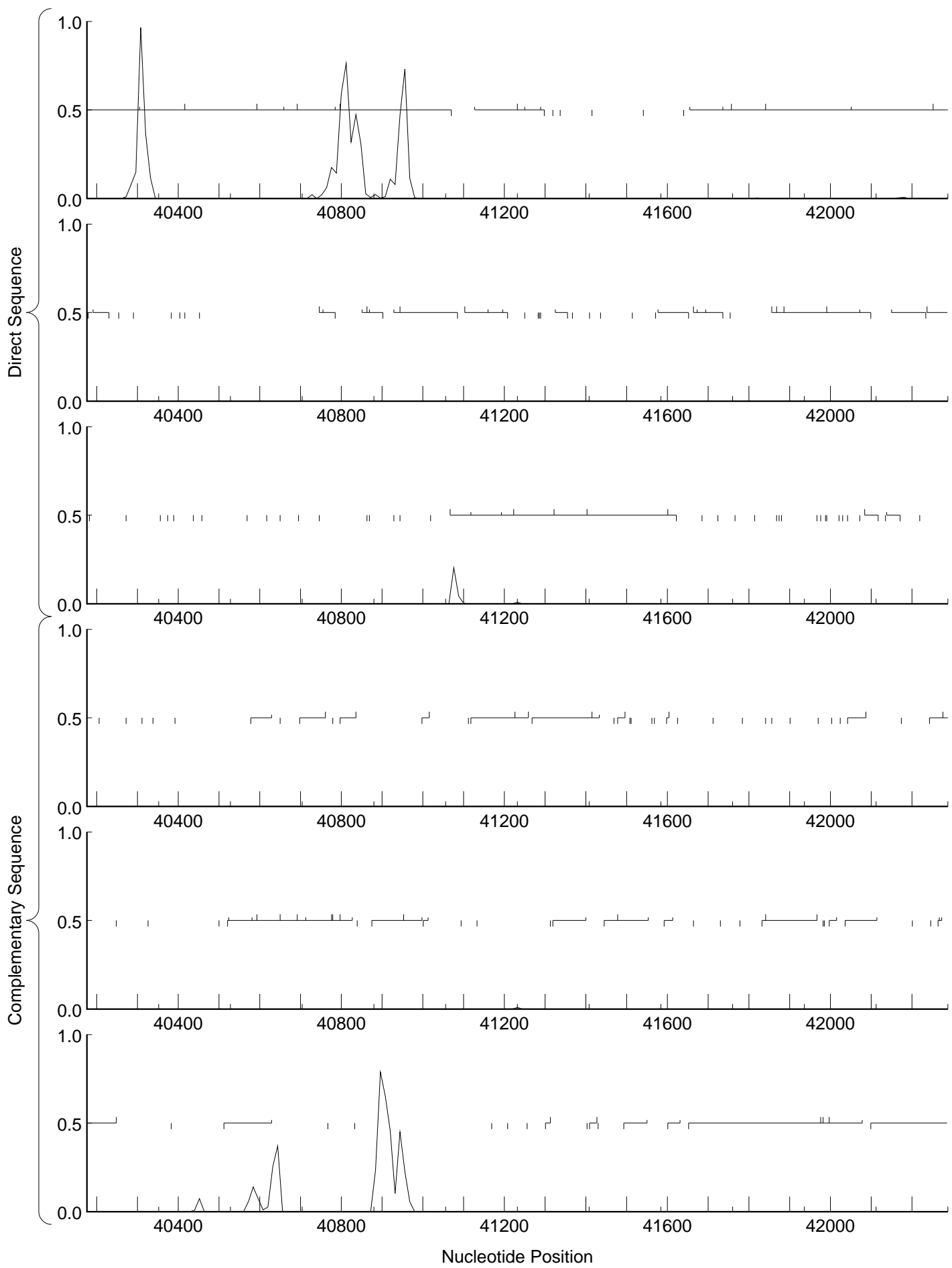




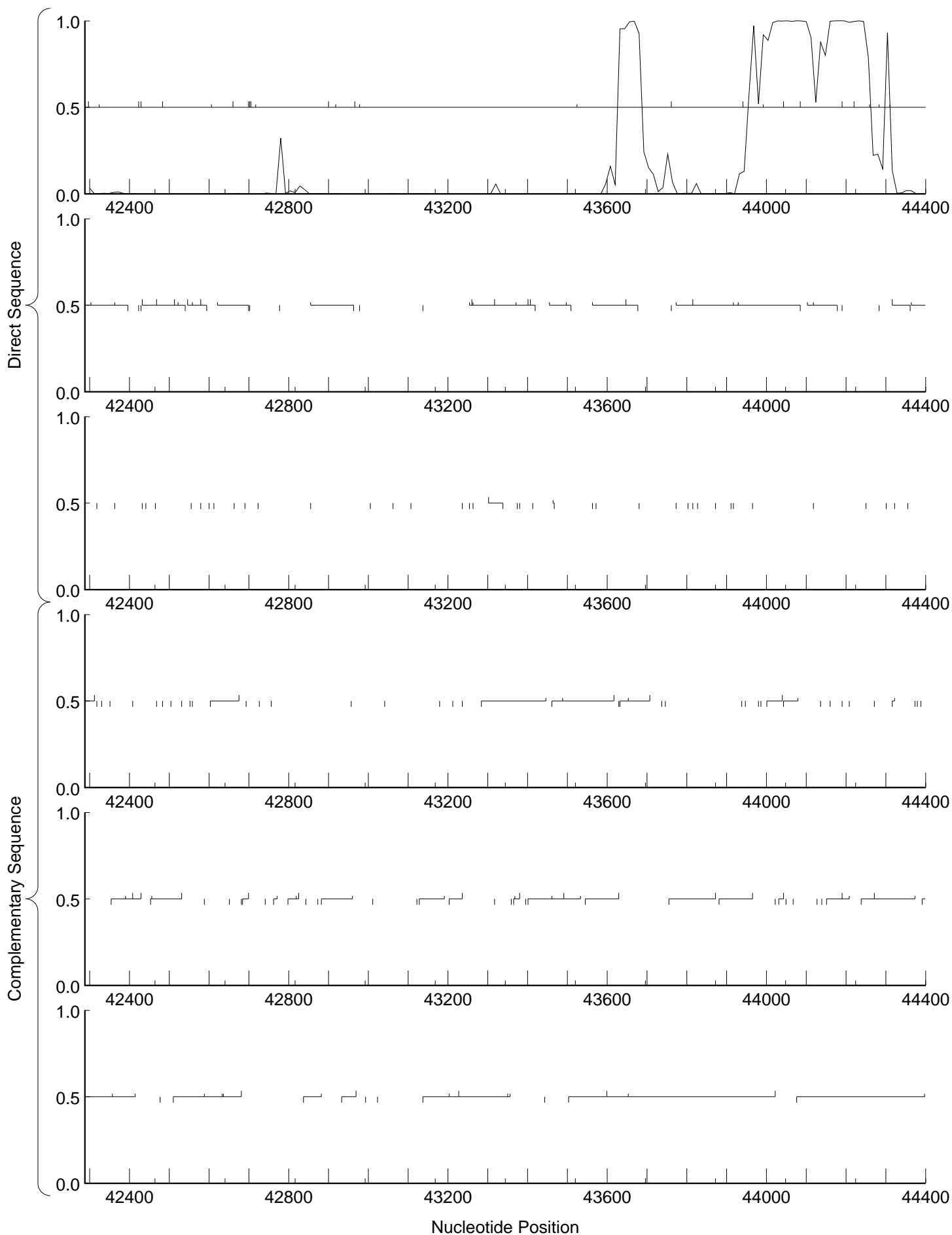
ley complete sequence, 60111 bp including 9-base 3' overhang (CGCCGCTCT), Cluster DJ, Order 4, Window 96, Step 12, 19/30



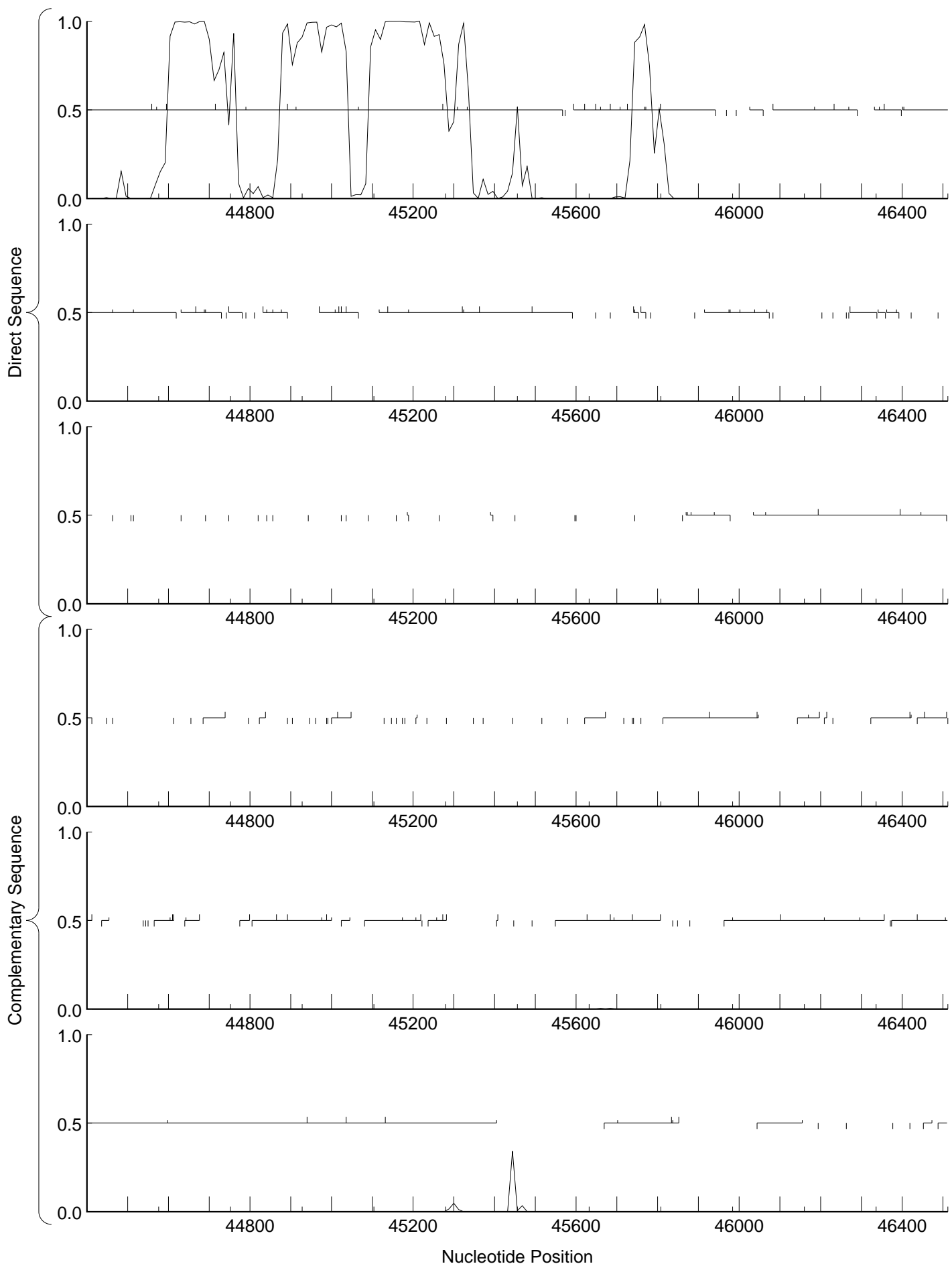


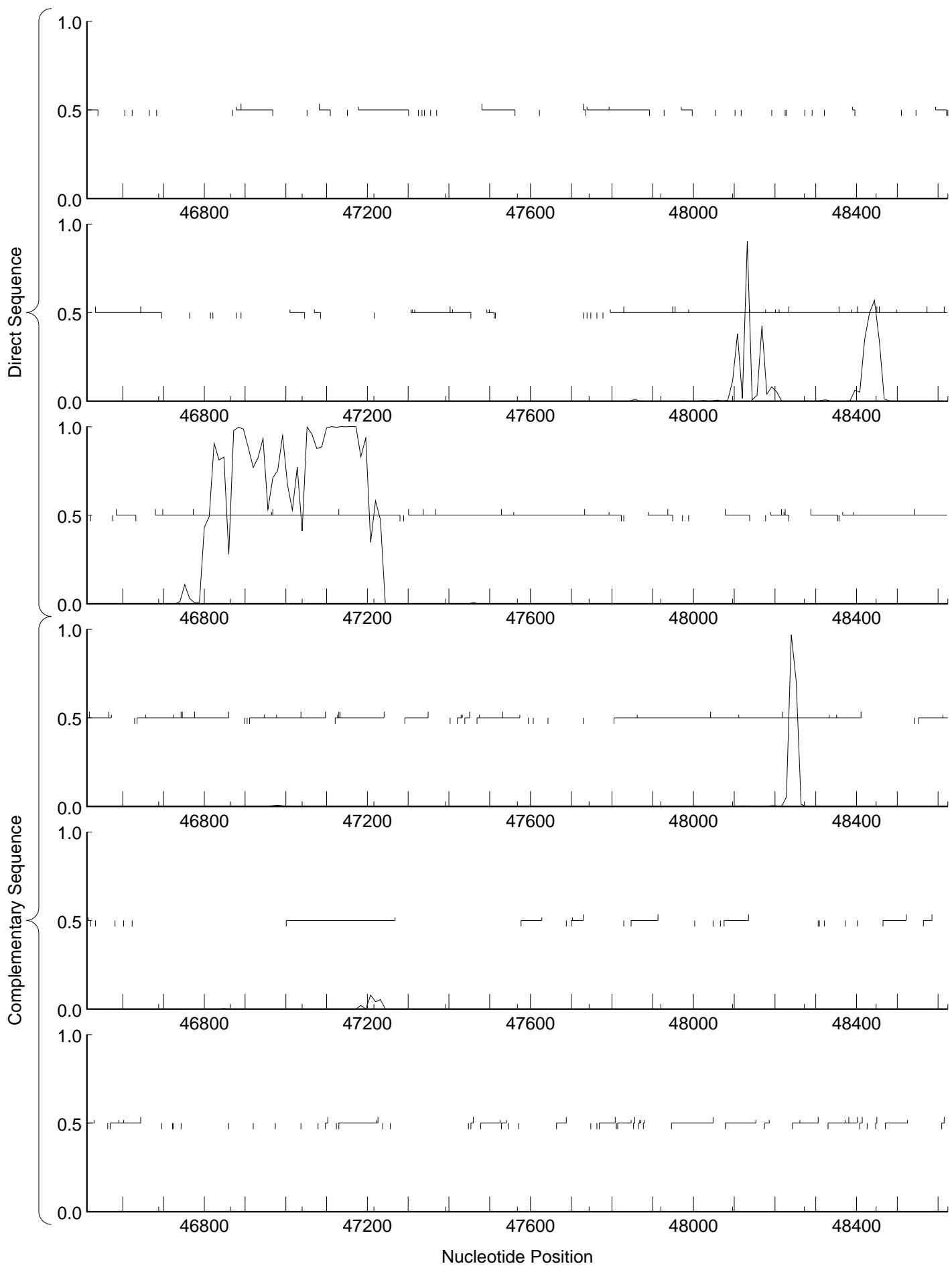


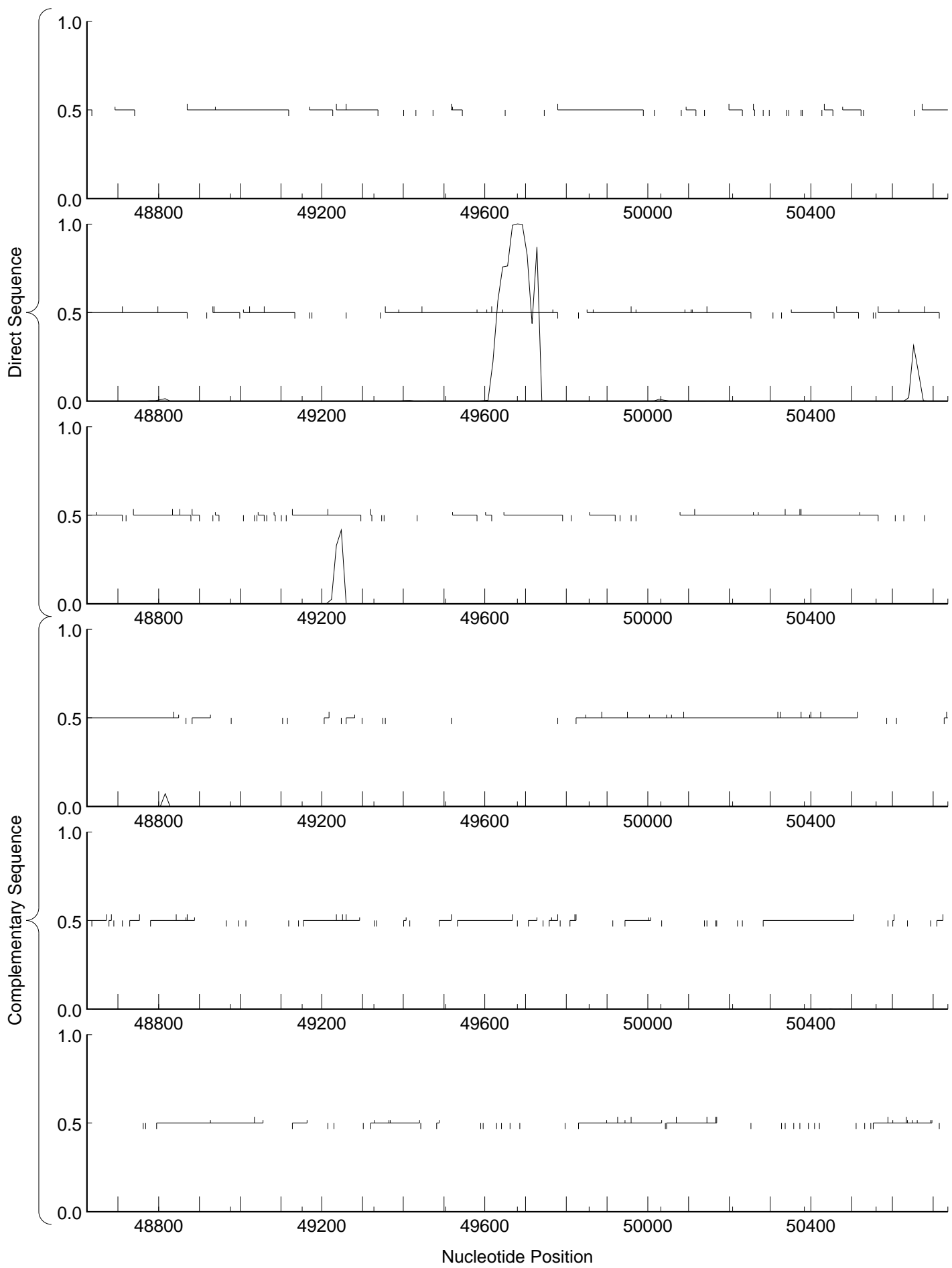
Key complete sequence, 60111 bp including 9-base 3' overhang (CGCCGCTCT), Cluster DJ, Order 4, Window 96, Step 12, 22/30

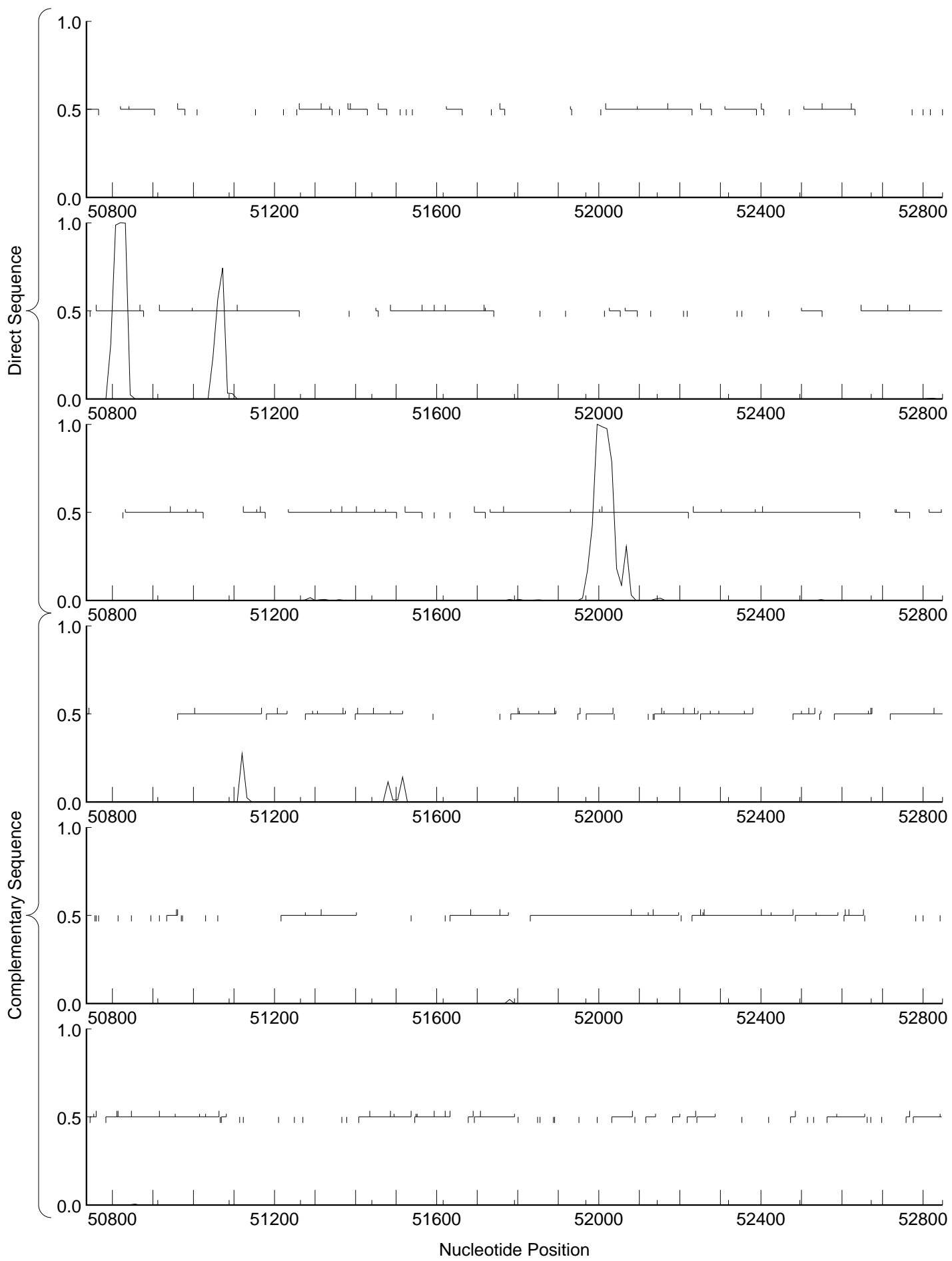


Key complete sequence, 60111 bp including 9-base 3' overhang (CGCCGCTCT), Cluster DJ, Order 4, Window 96, Step 12, 23/30









Key complete sequence, 60111 bp including 9-base 3' overhang (CGCCGCTCT), Cluster DJ, Order 4, Window 96, Step 12, 27/30

