

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Gordonia phage OhMyWard complete sequence, 60978 bp including 9-base 3' overhang (CGCCGCCCT  
Analysis Date : 4/27/25 at 21:50:21  
Pages : 30  
Sequence Length : 60978 bp  
GC Content : 52.15%

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,

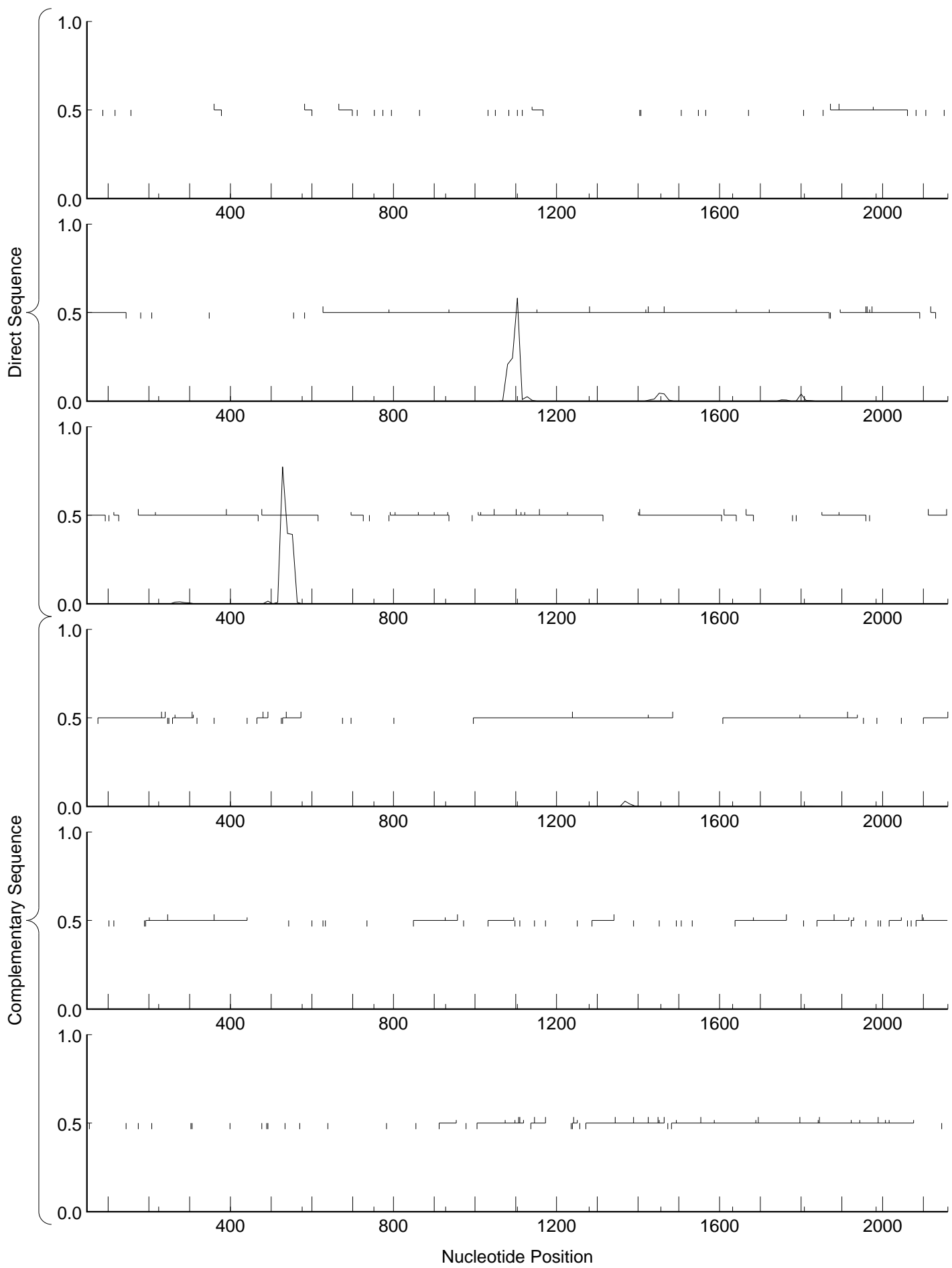
## Matrix notes & comments

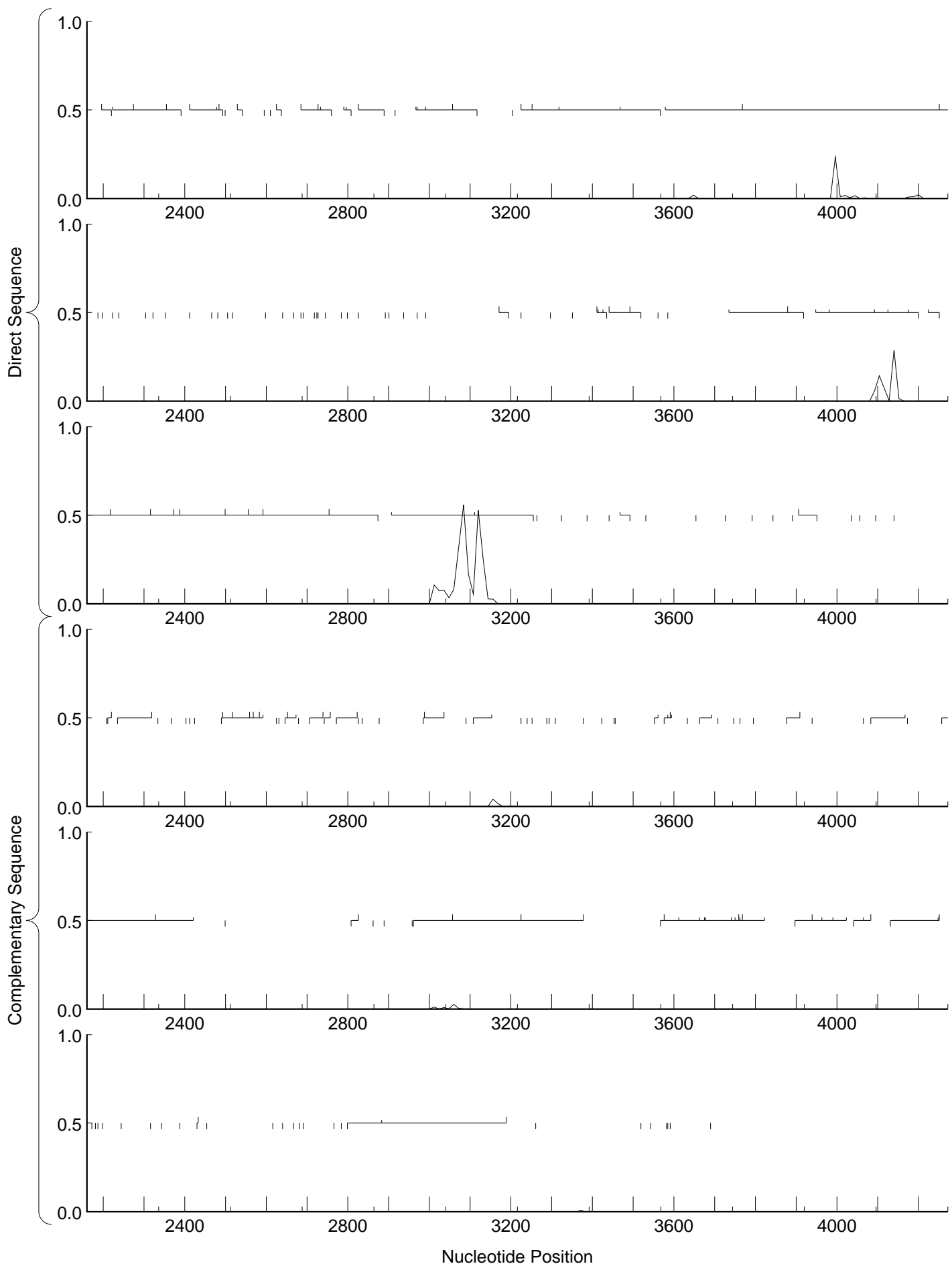
Training set derived by GeneMarkS, 4.30 October 2014  
Sat Sep 8 00:36:14 2018

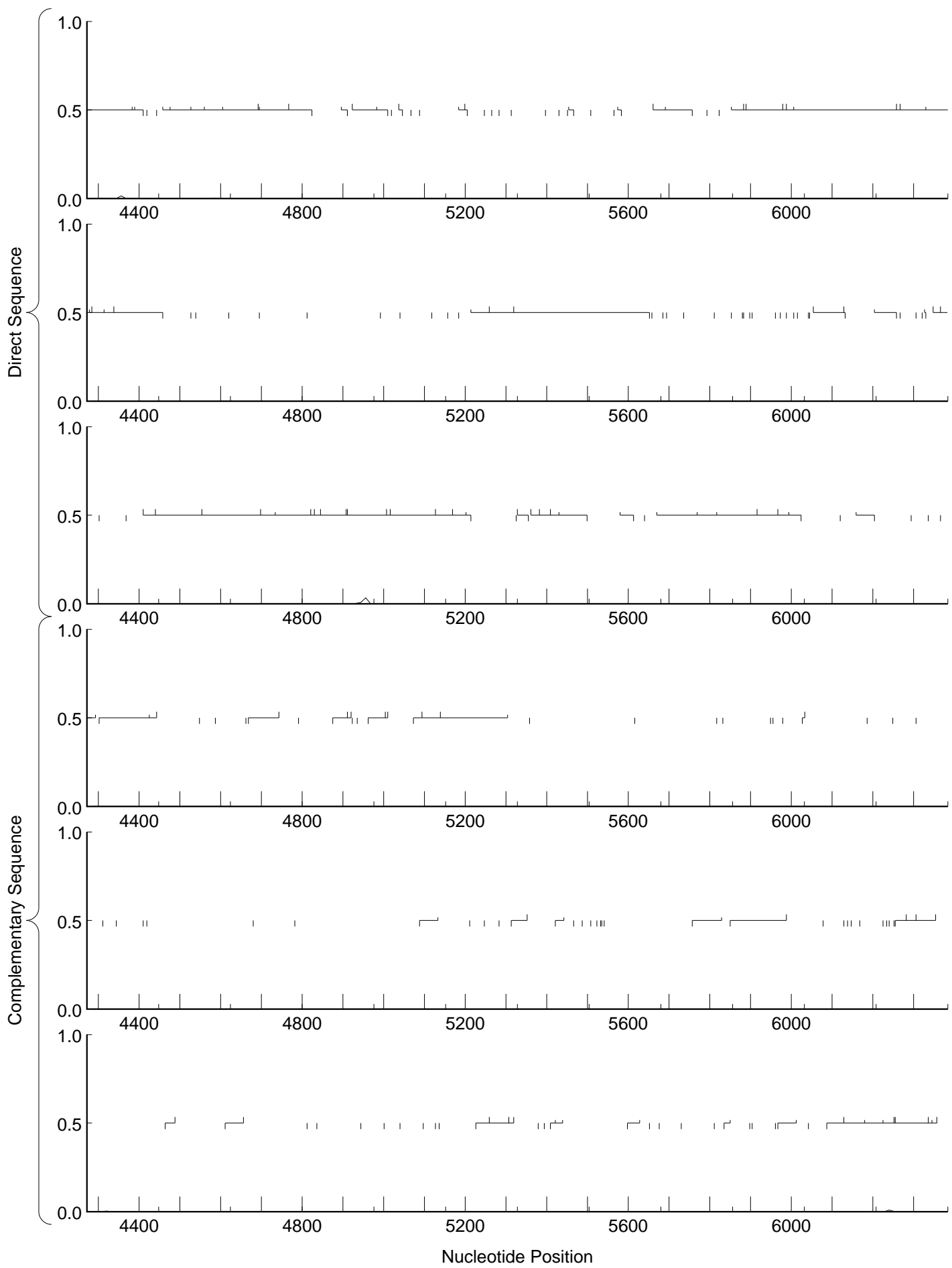
## MATRIX INFORMATION

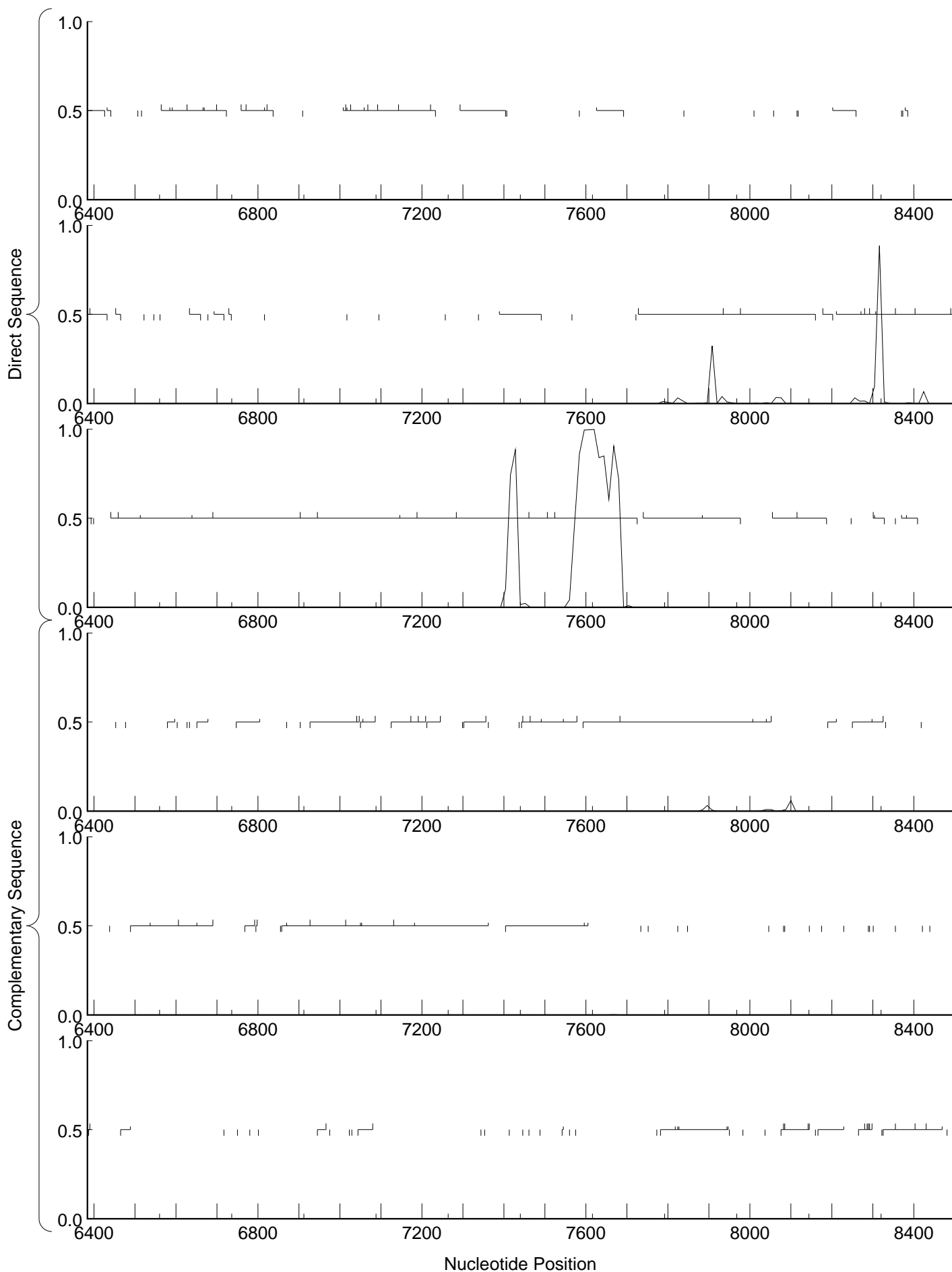
Matrix : -  
Author : -  
Order : 4

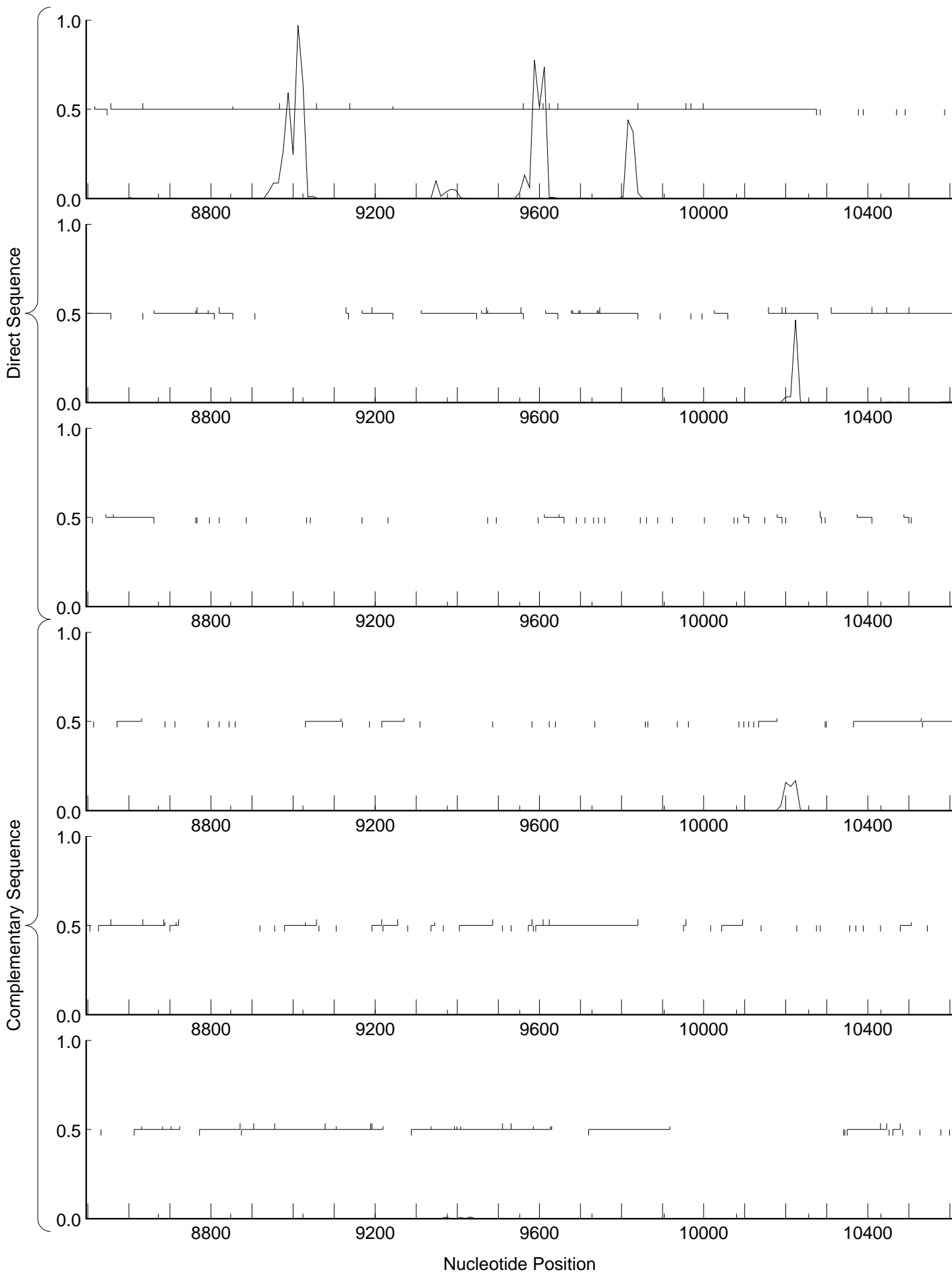
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

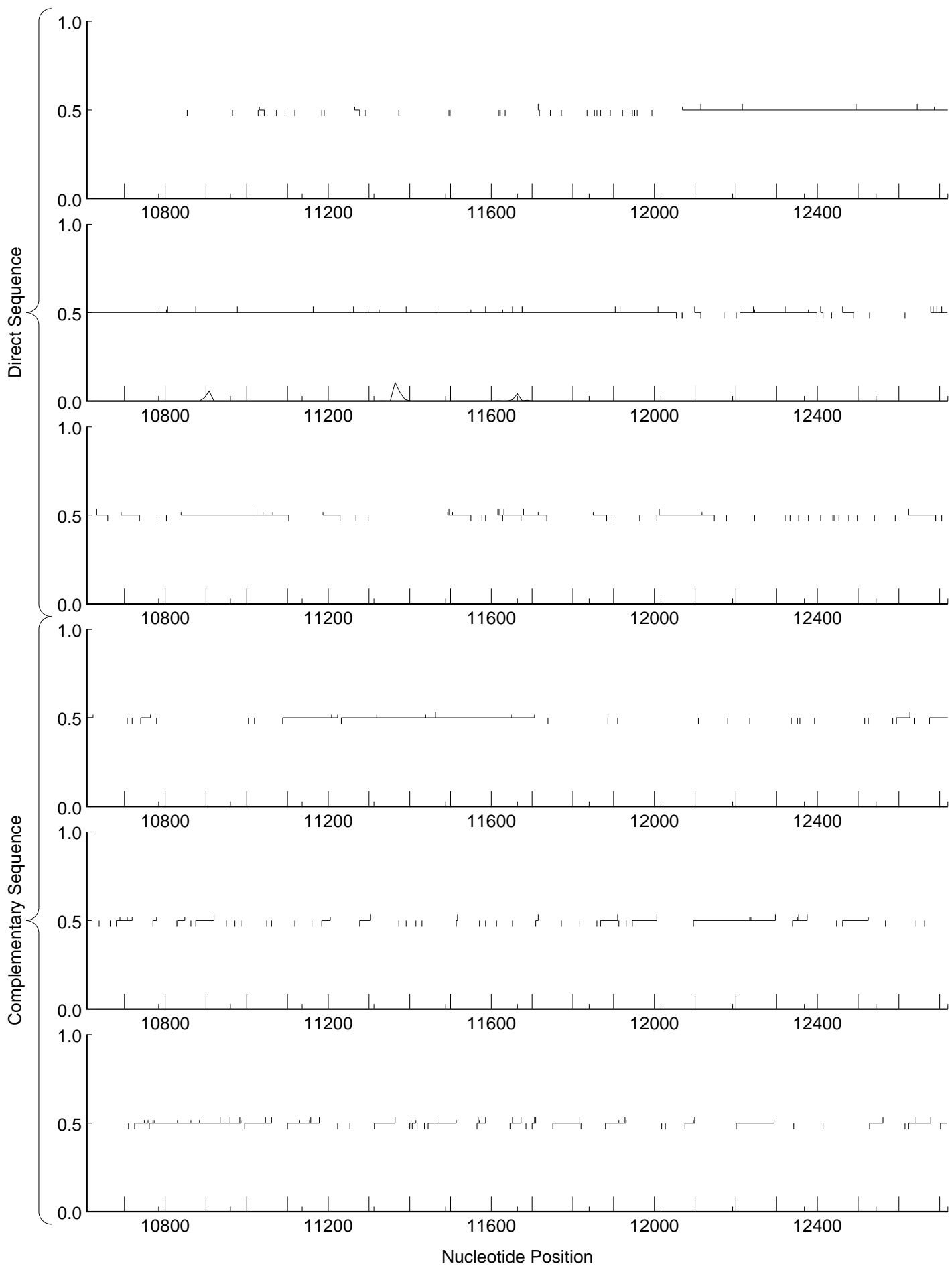


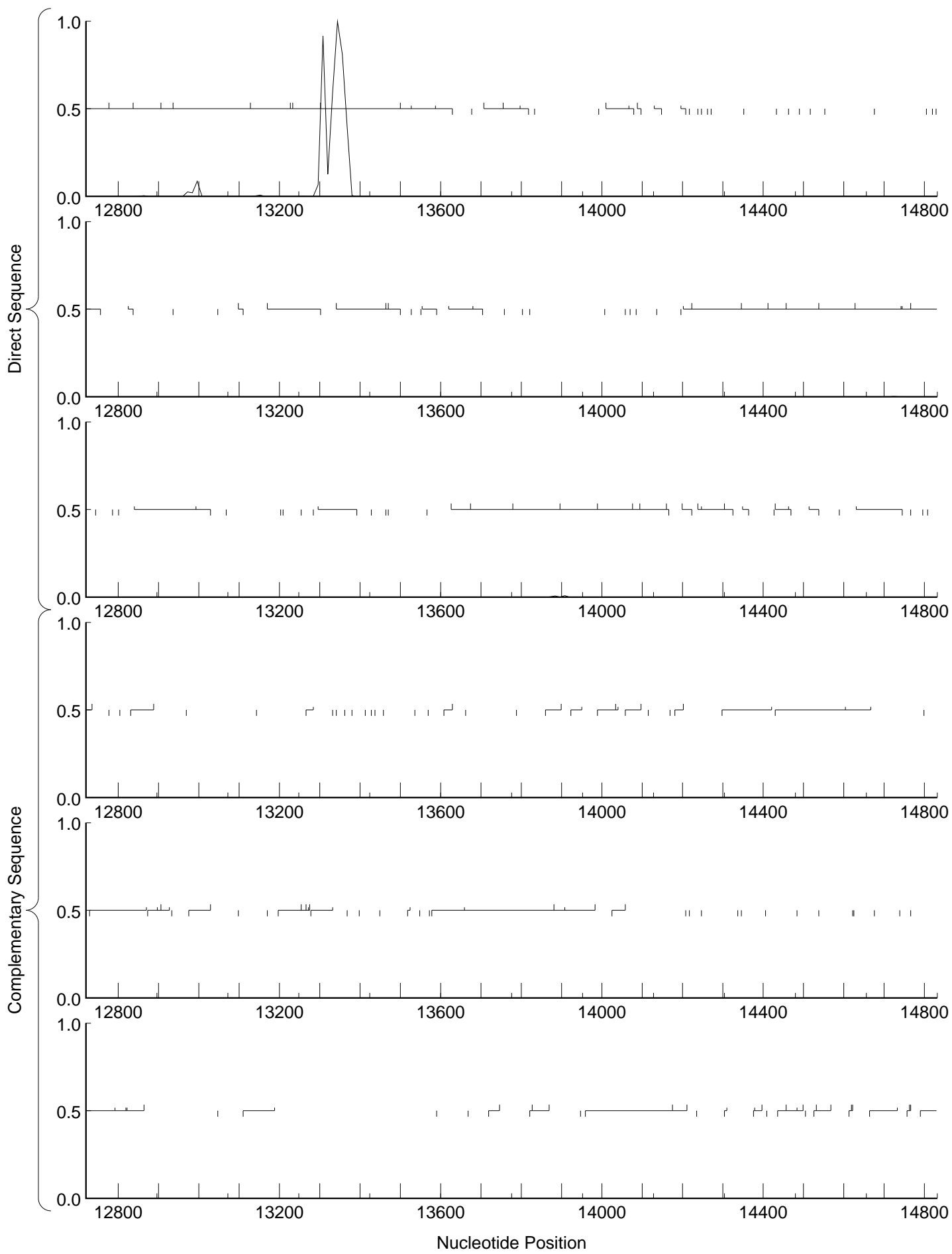




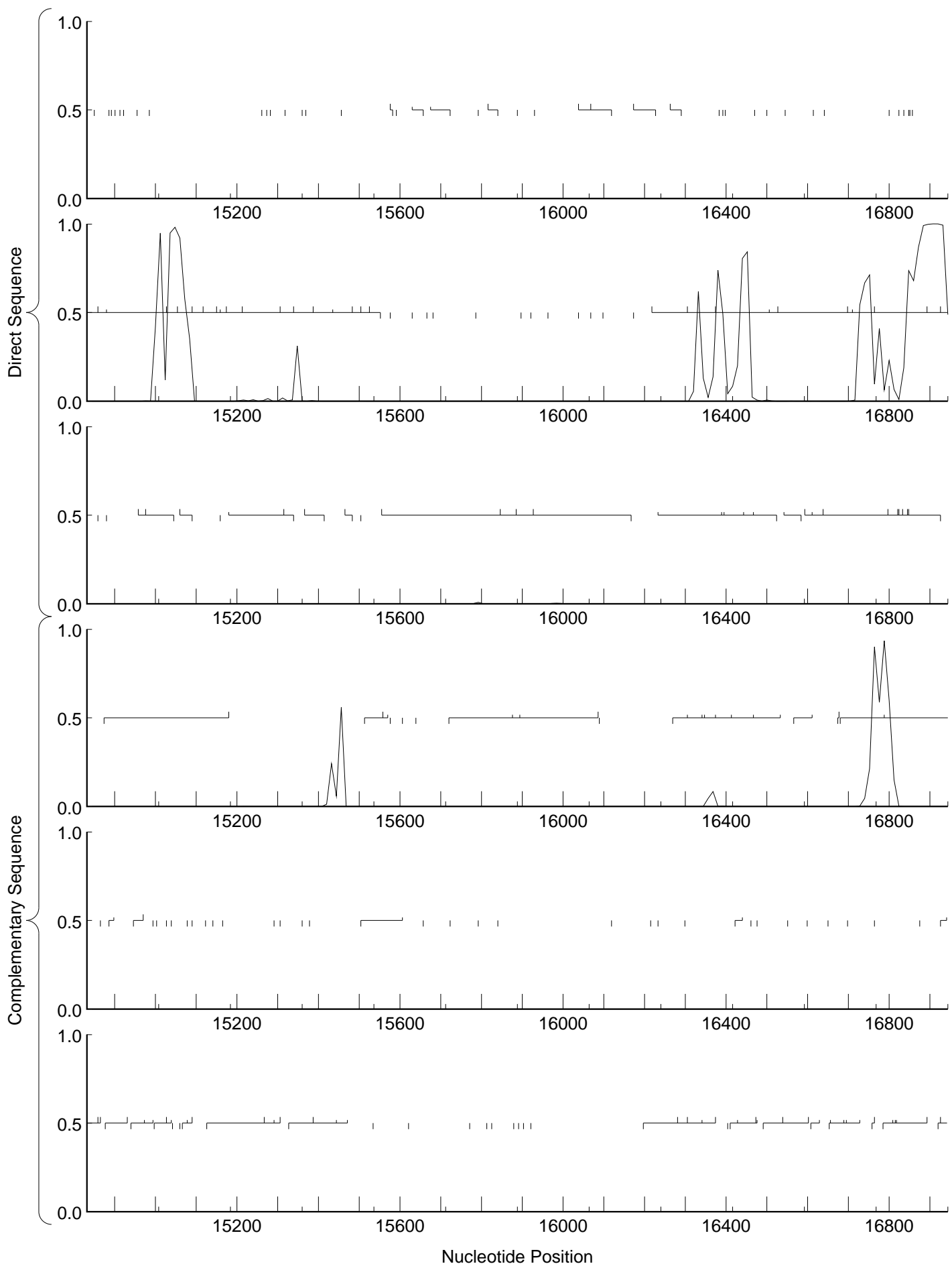


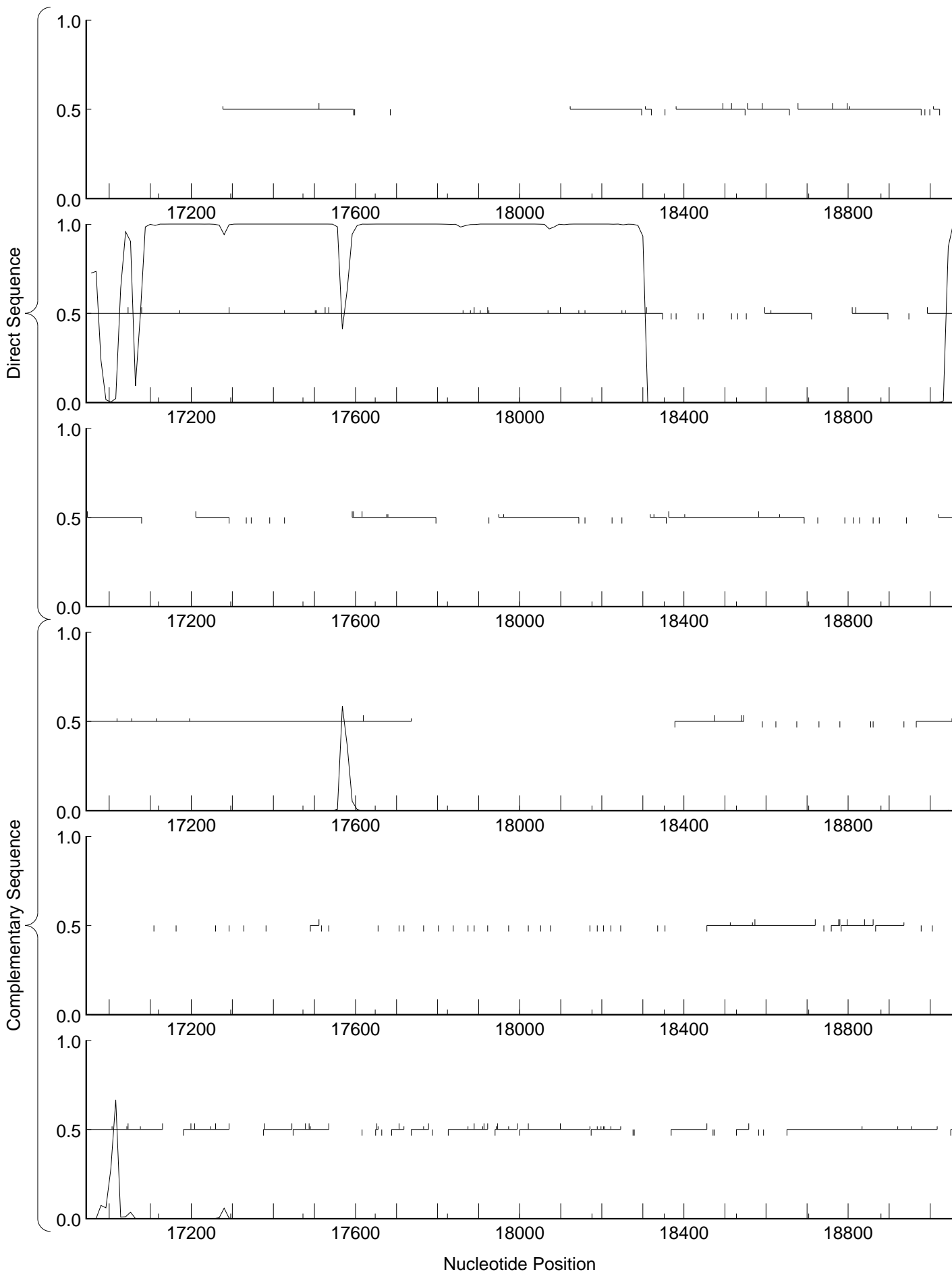


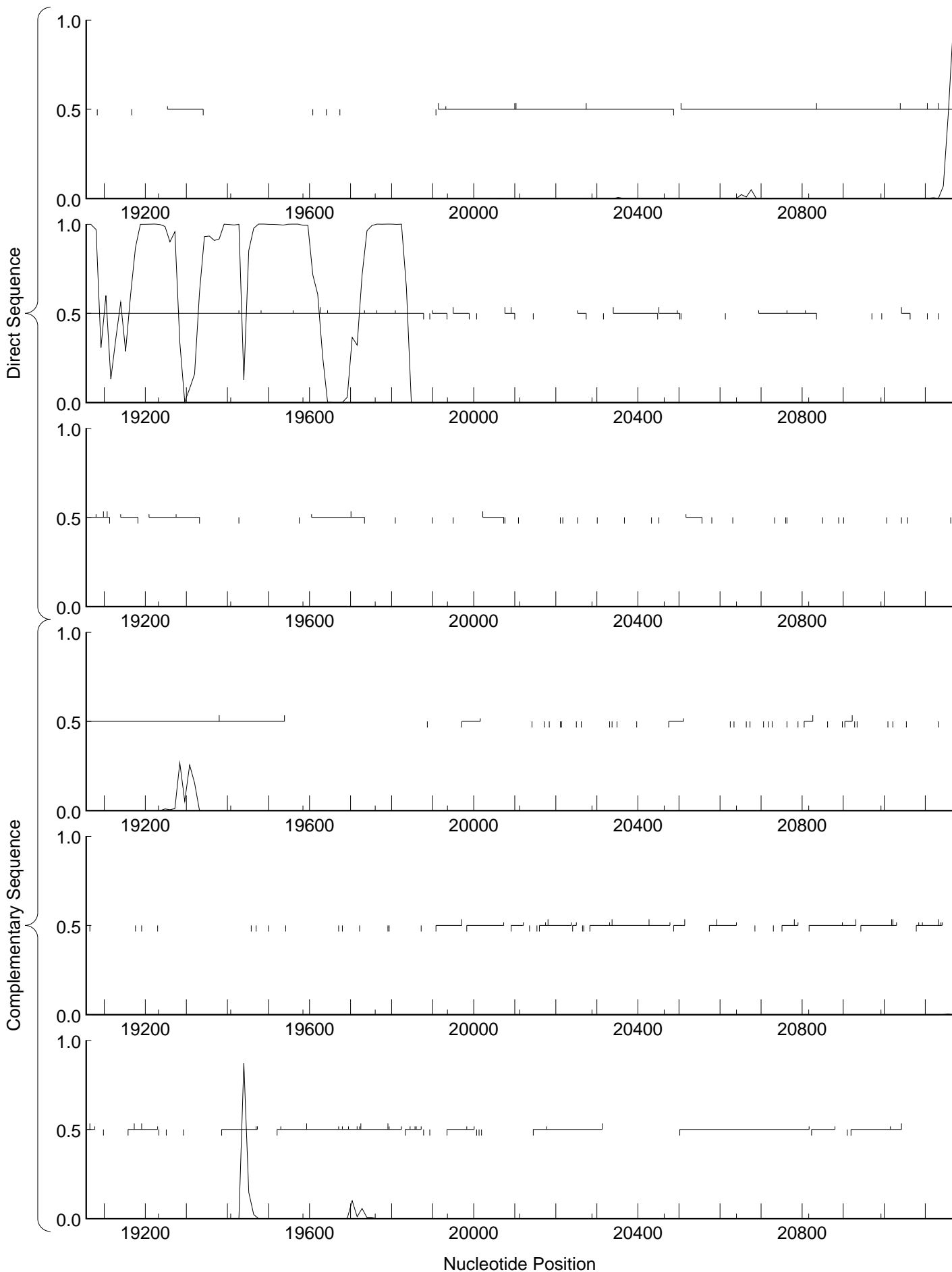


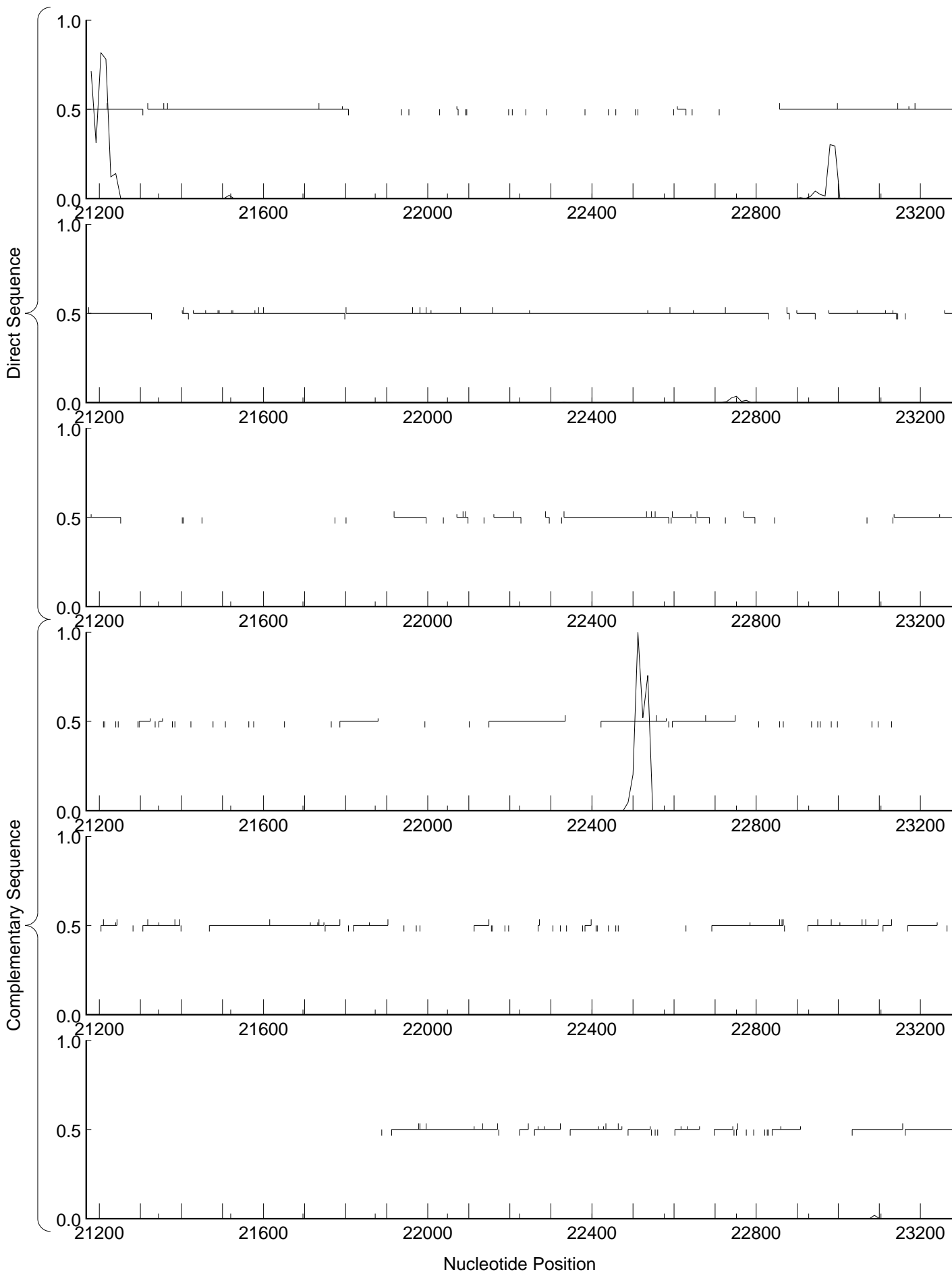


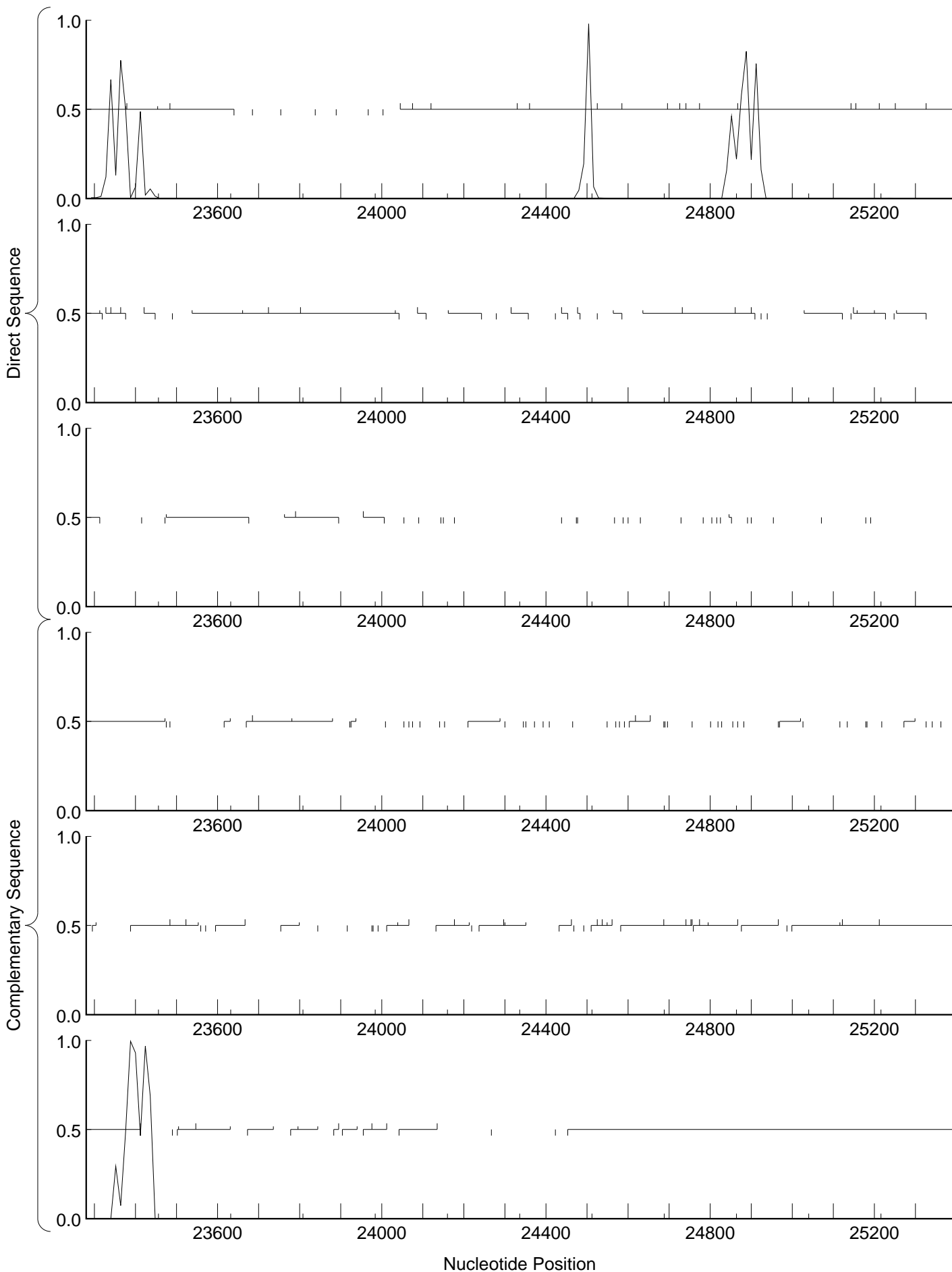


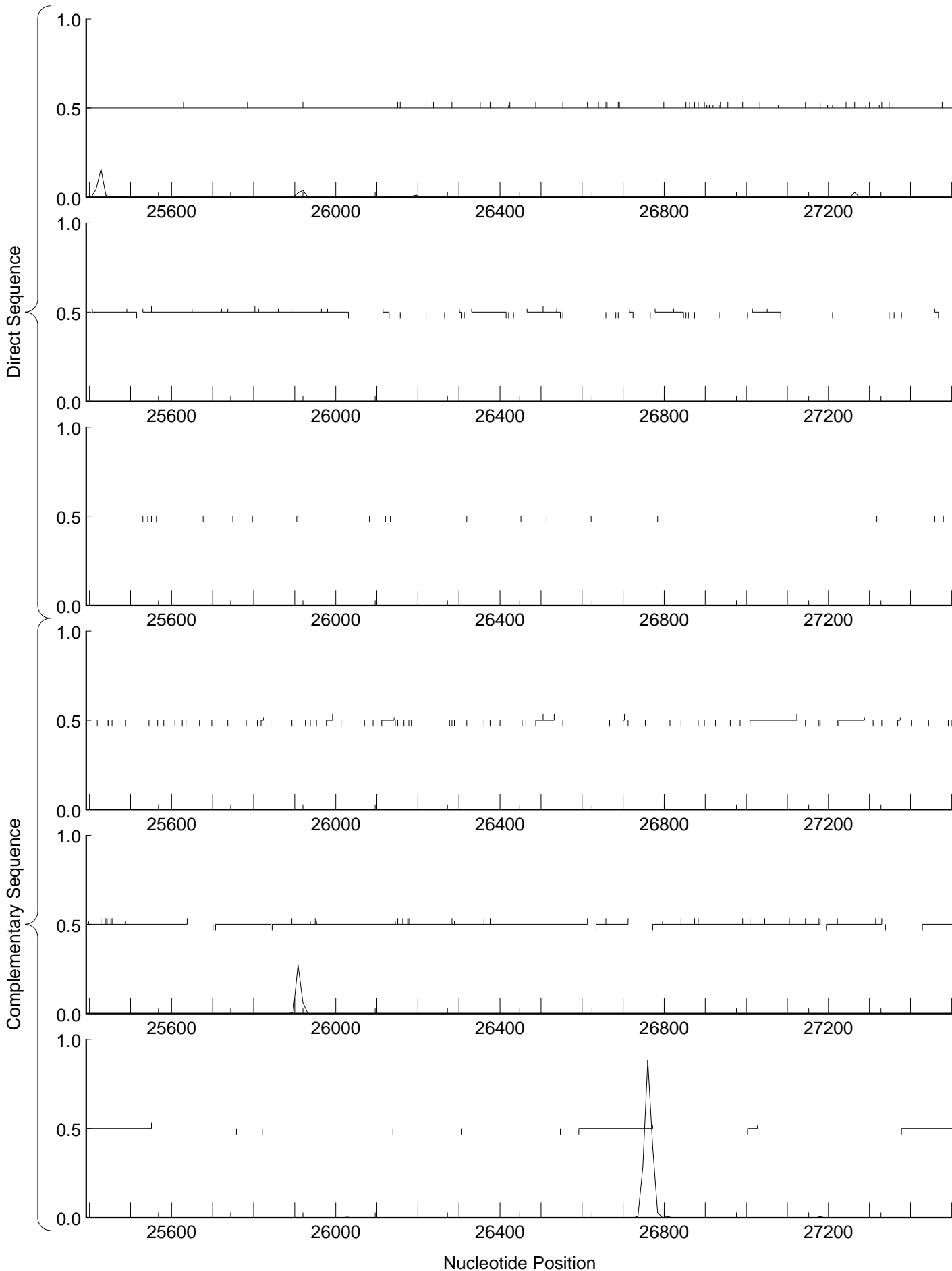




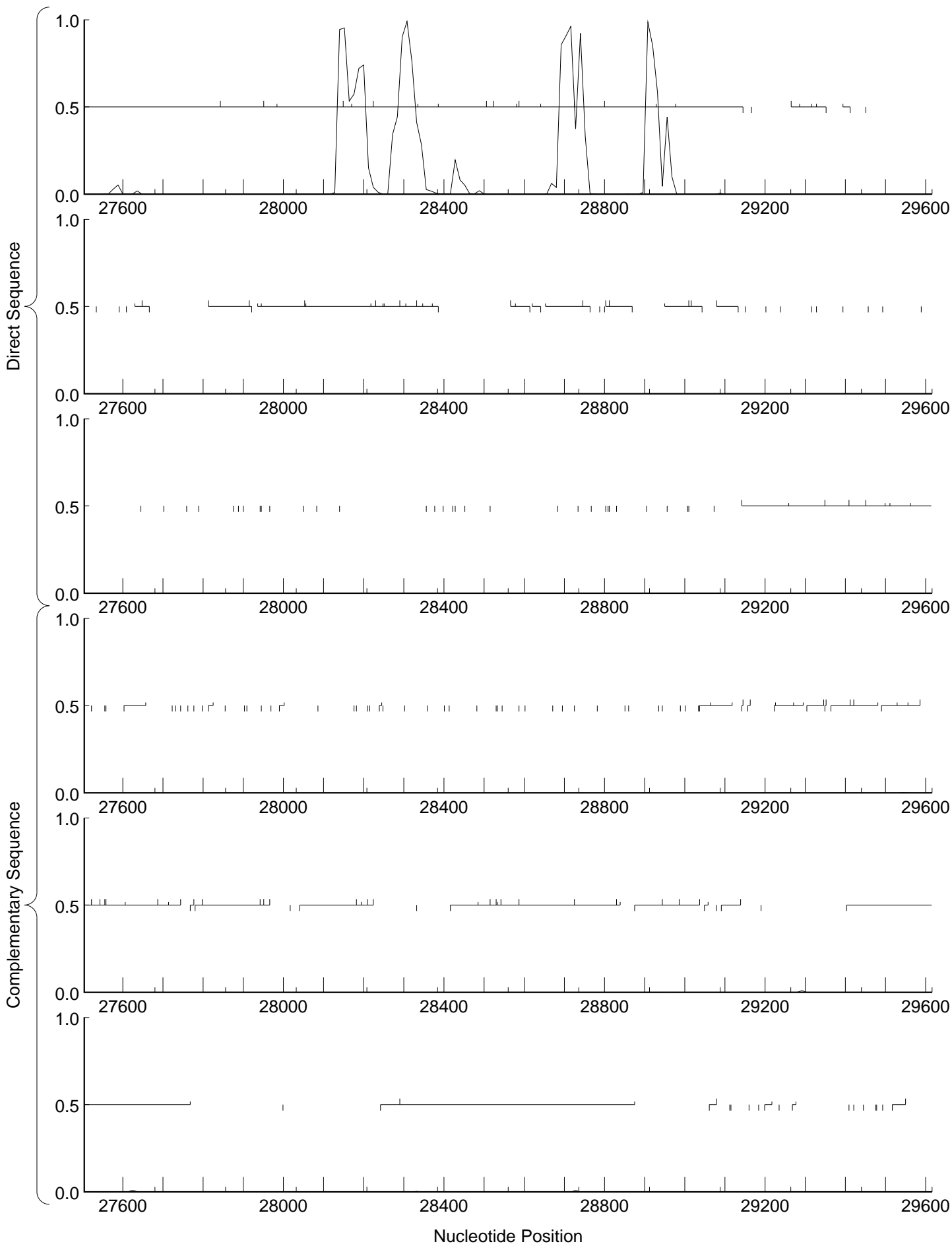


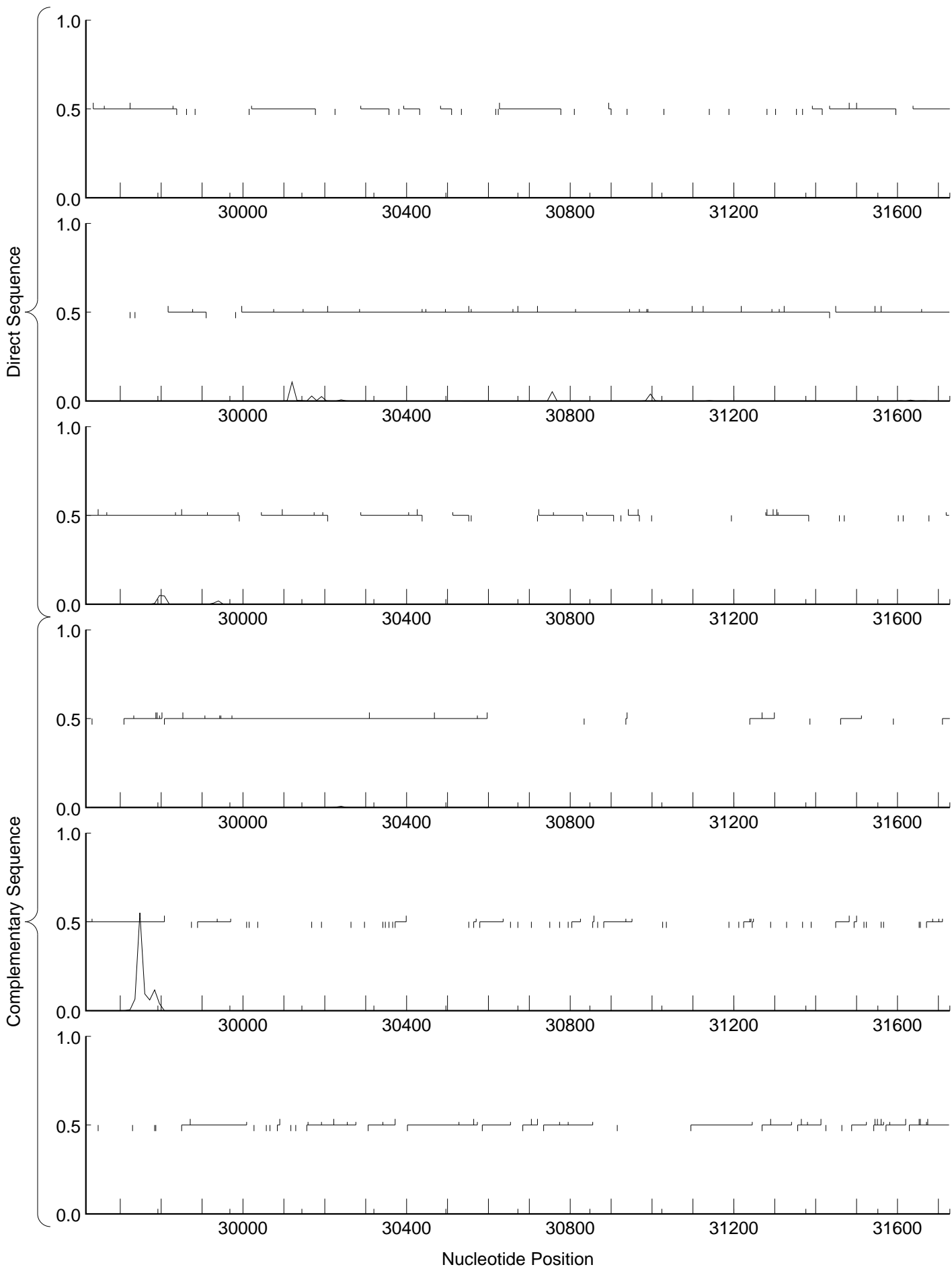




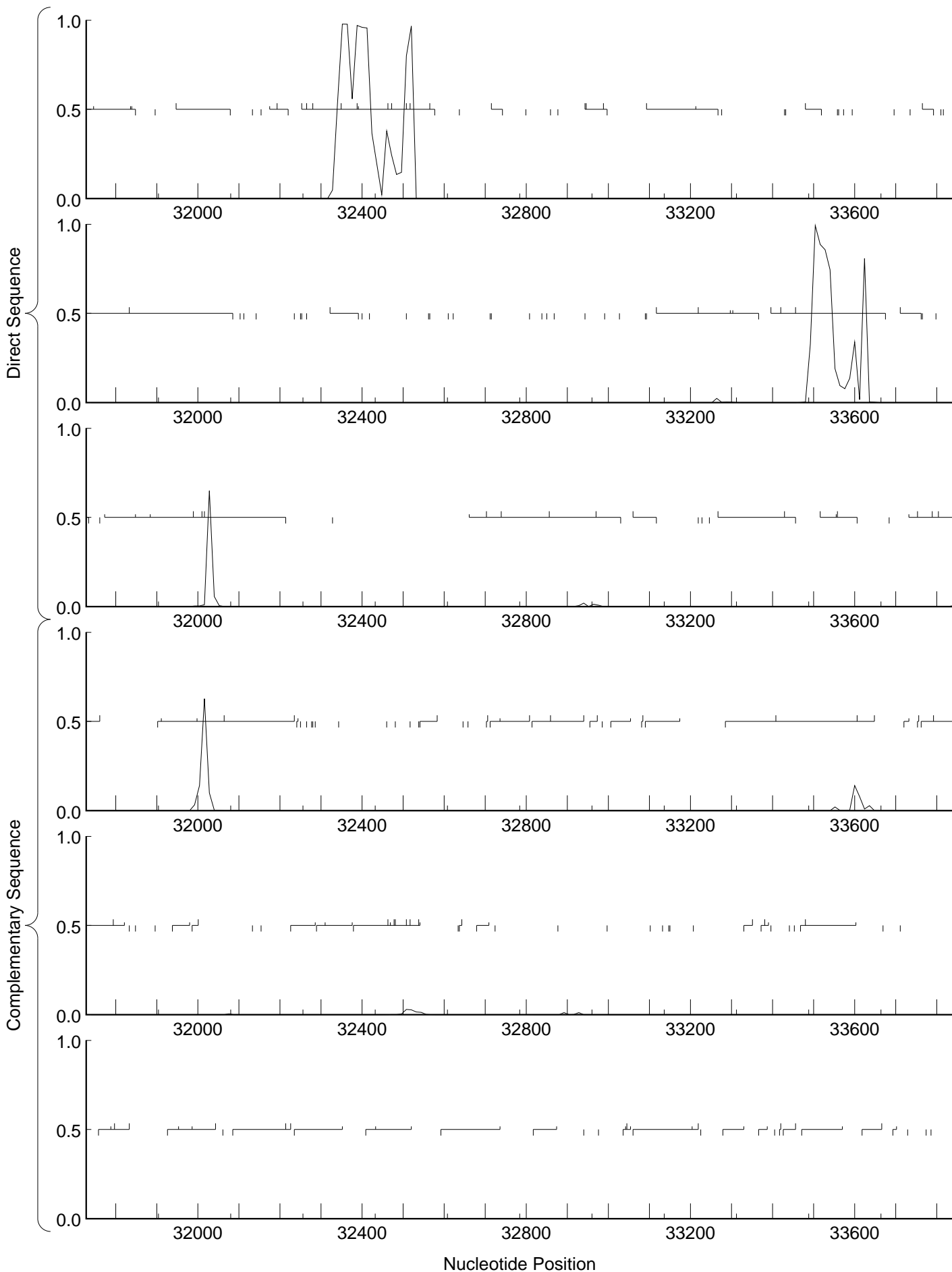


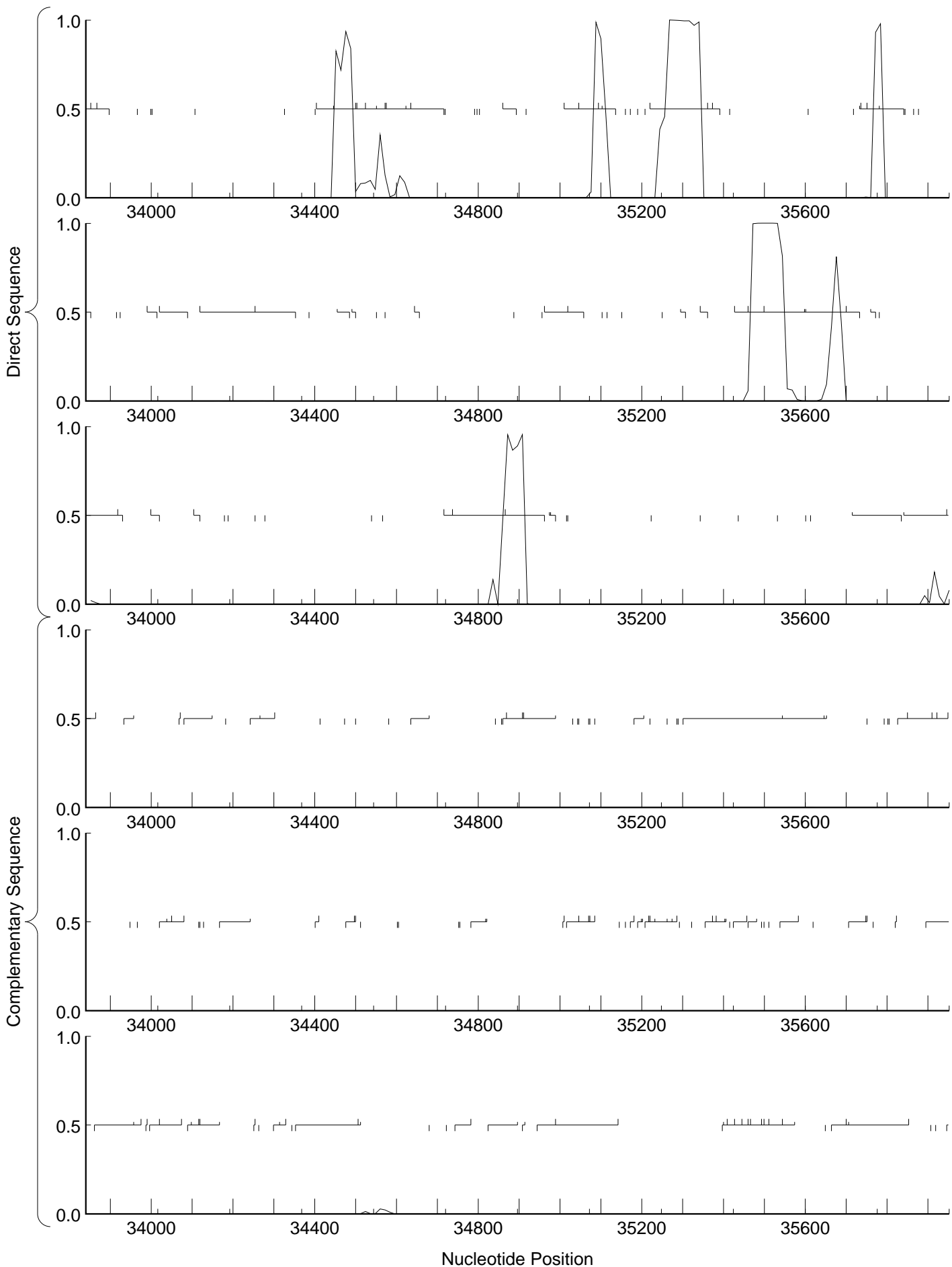
ard complete sequence, 60978 bp including 9-base 3' overhang (CGCCGCCCT), Cluster DJ, Order 4, Window 96, Step 12, 15/30



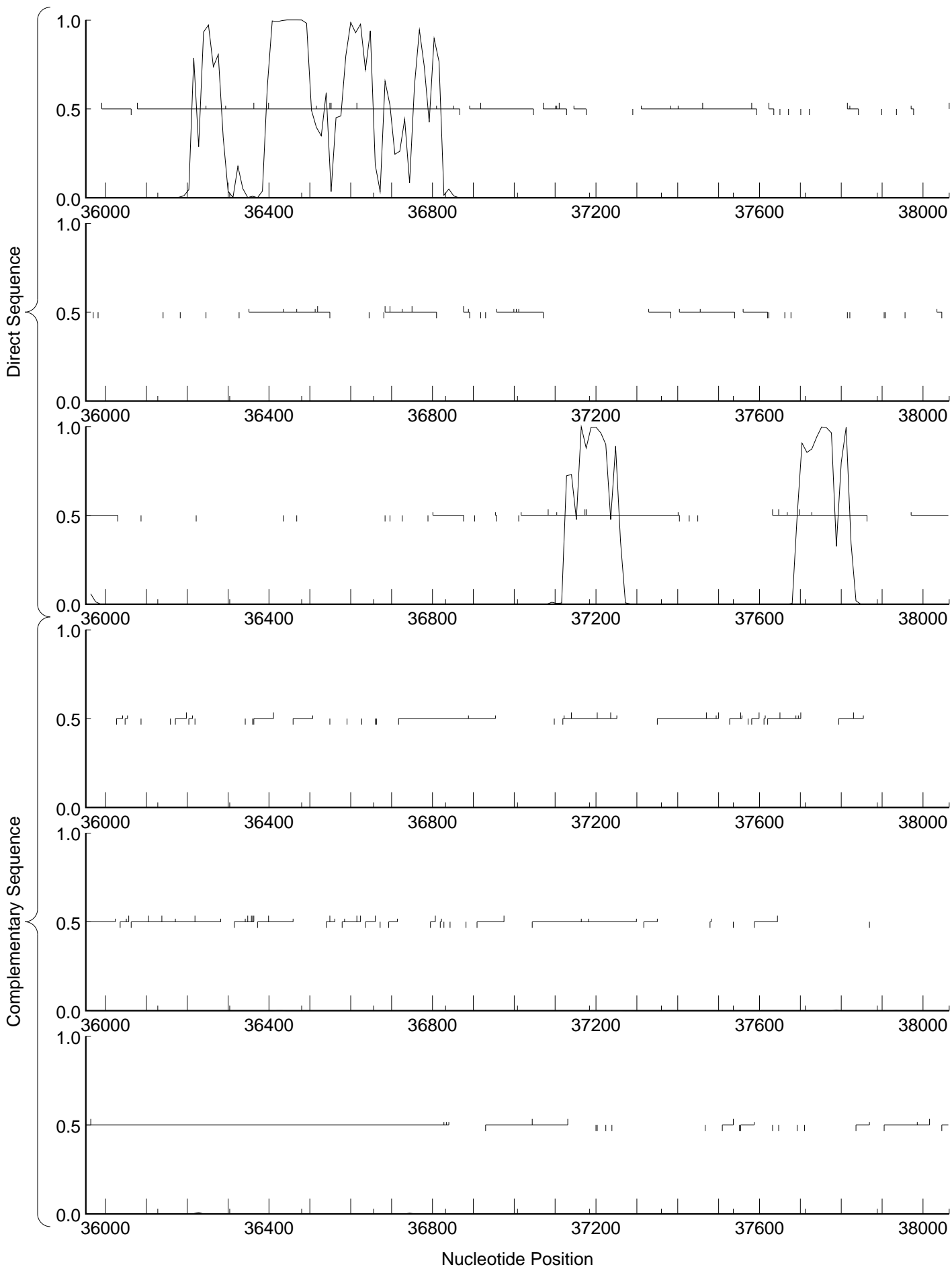


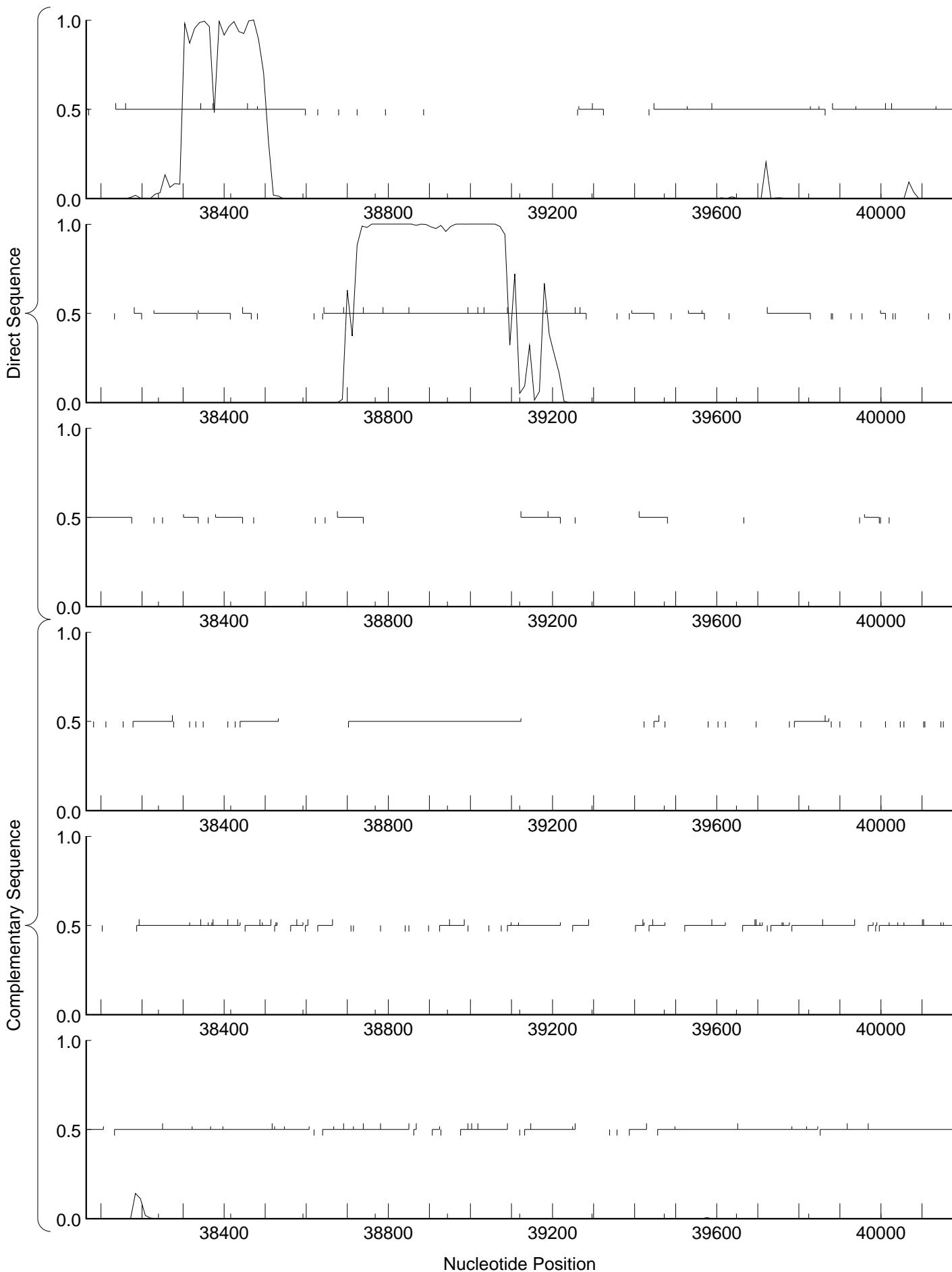


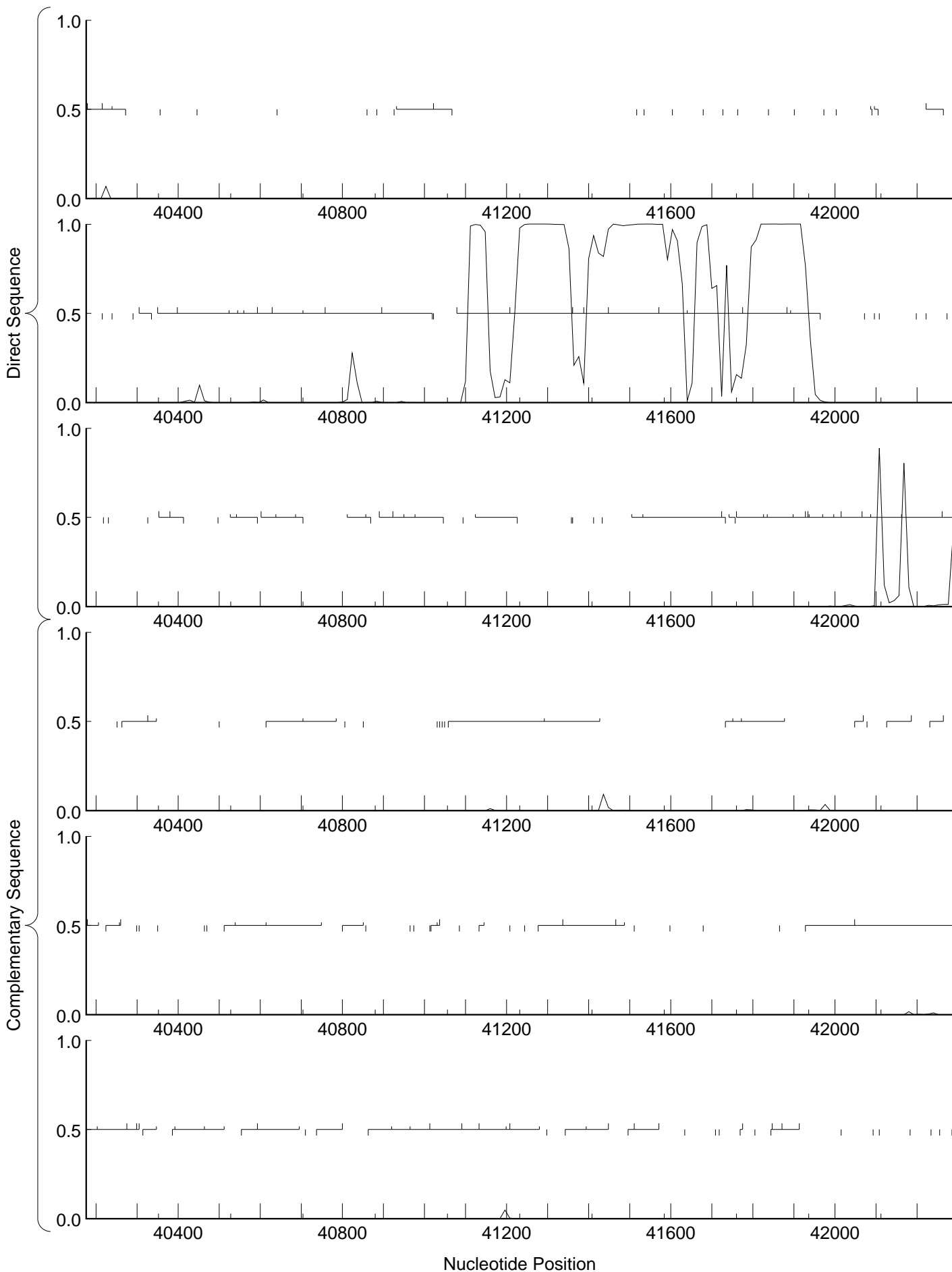


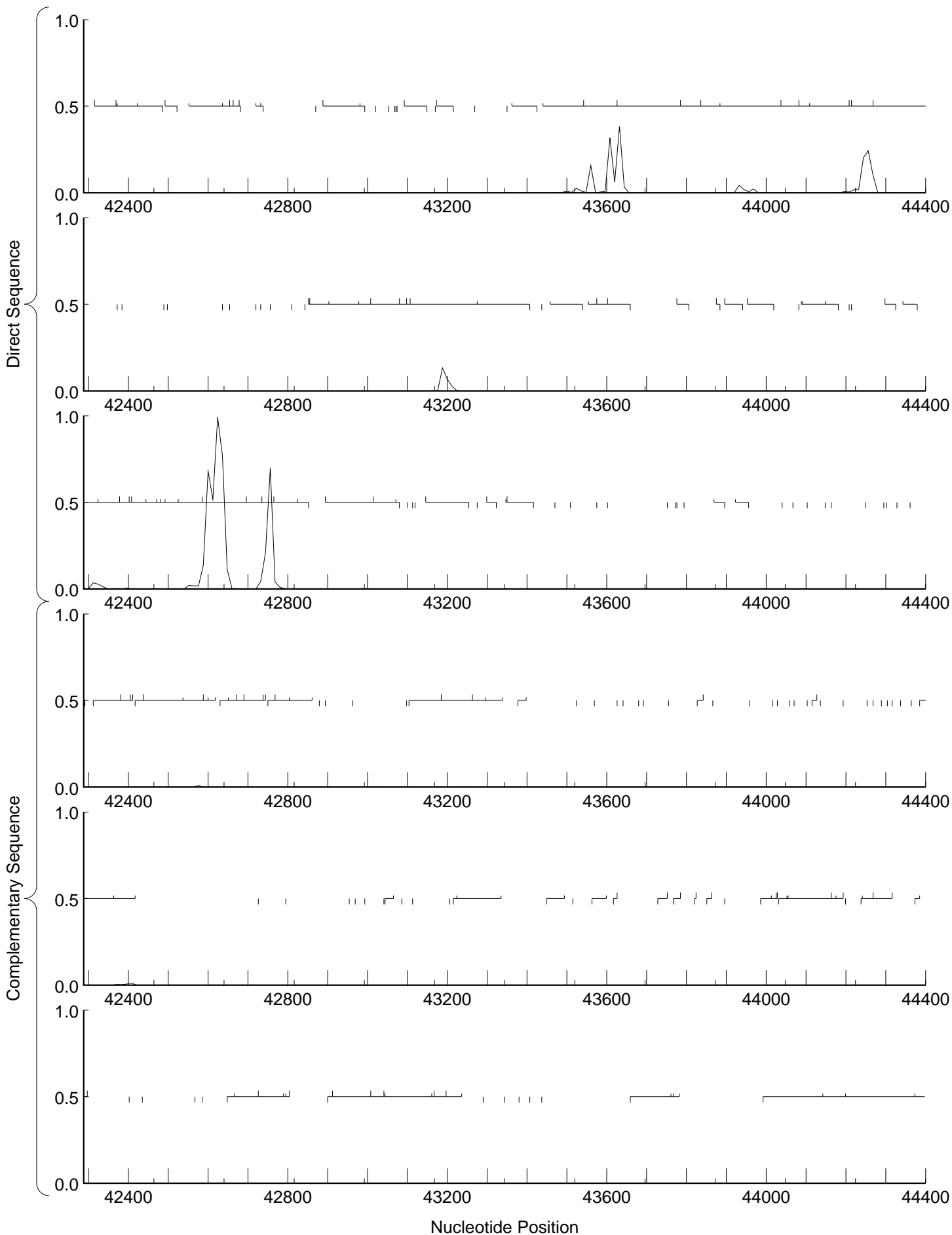


ard complete sequence, 60978 bp including 9-base 3' overhang (CGCCGCCCT), Cluster DJ, Order 4, Window 96, Step 12, 19/30

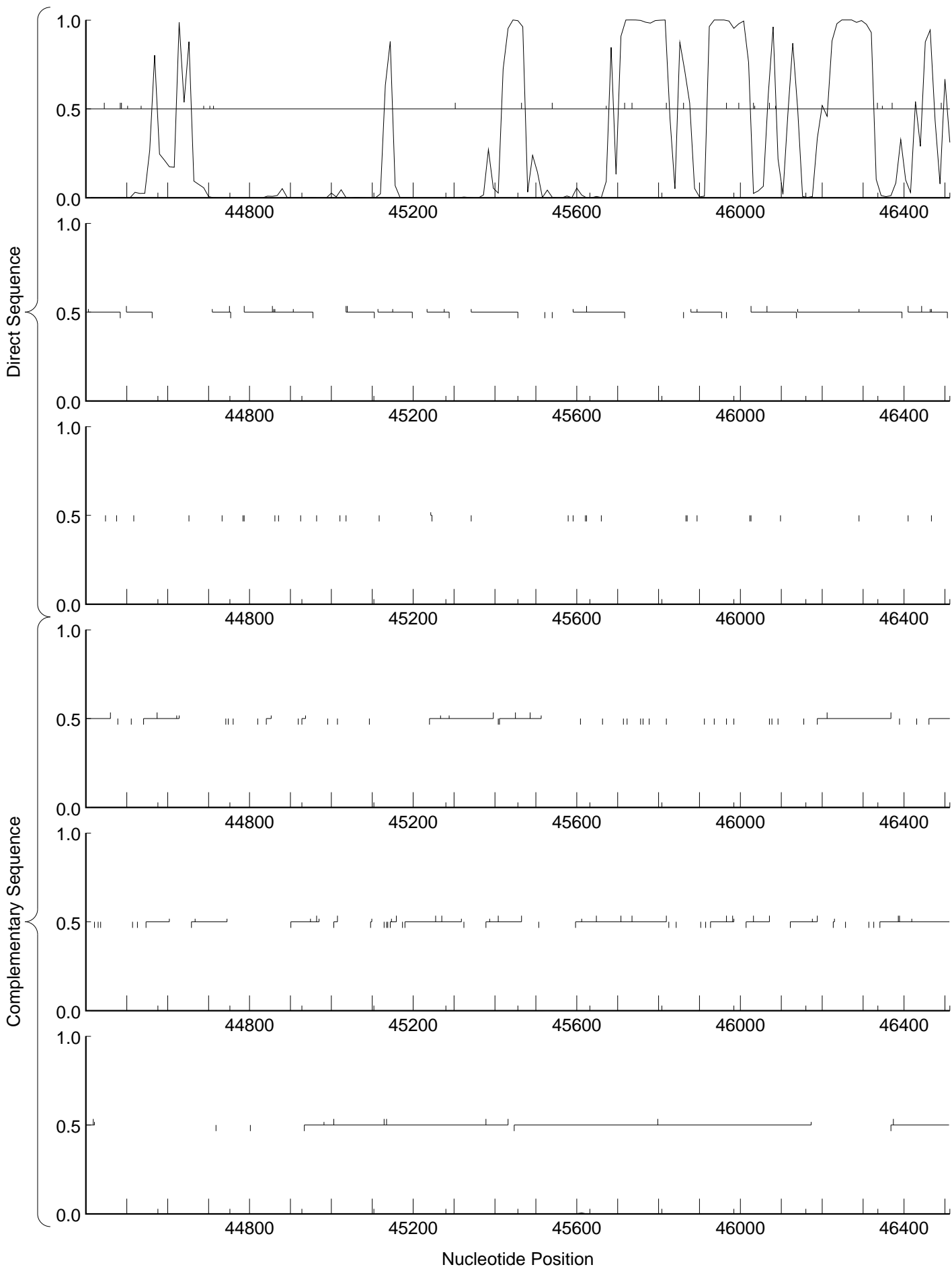


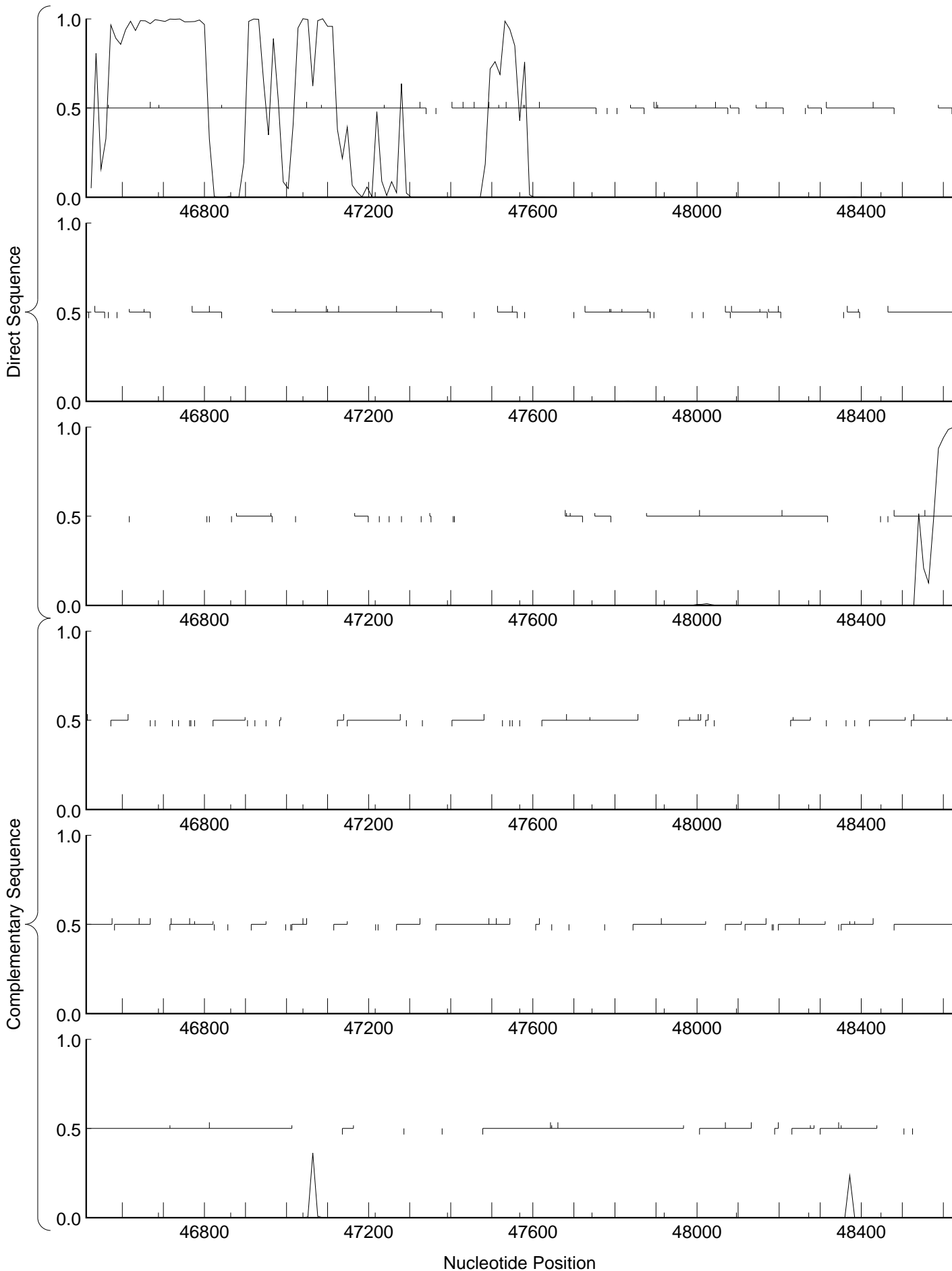




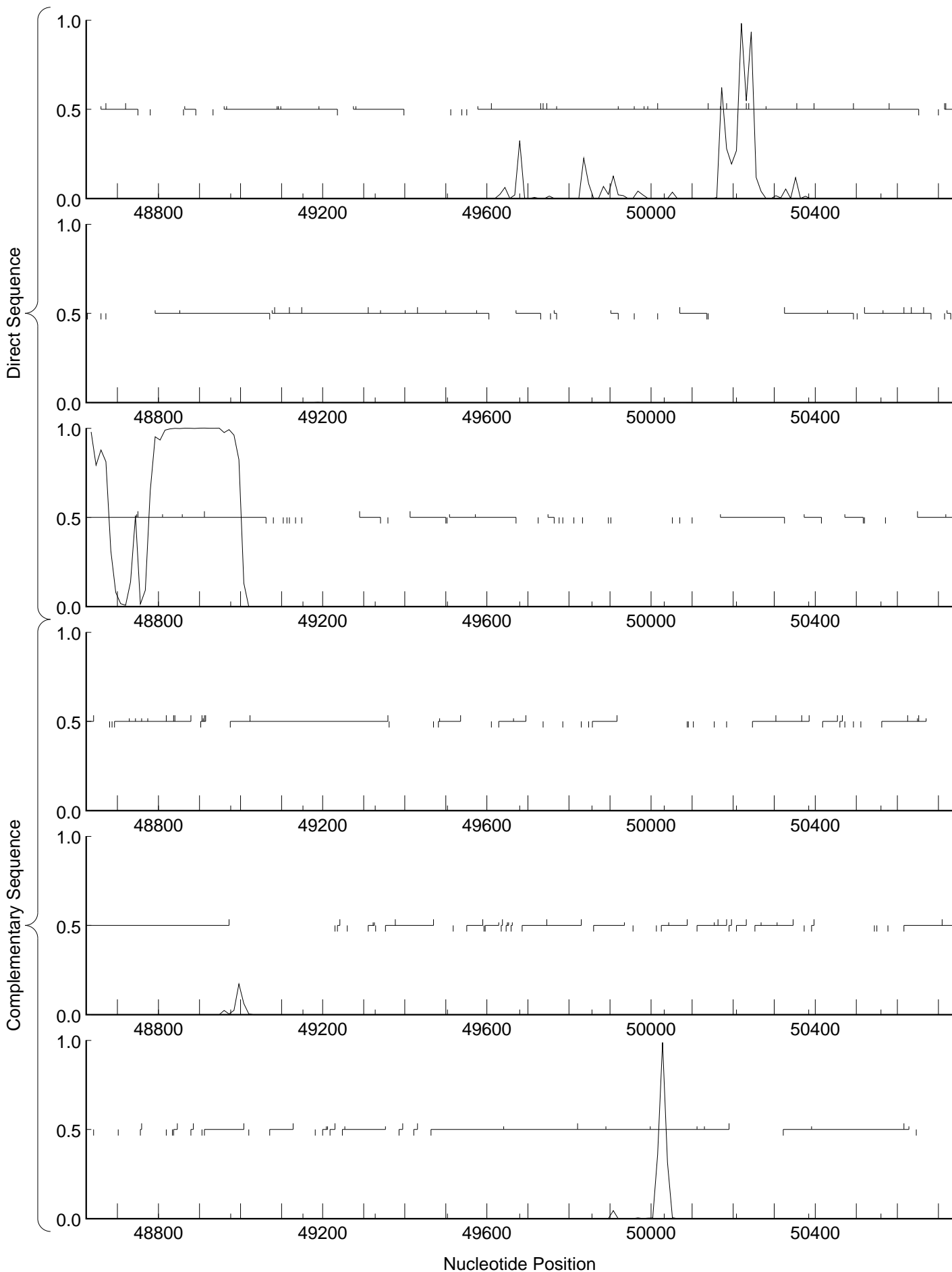


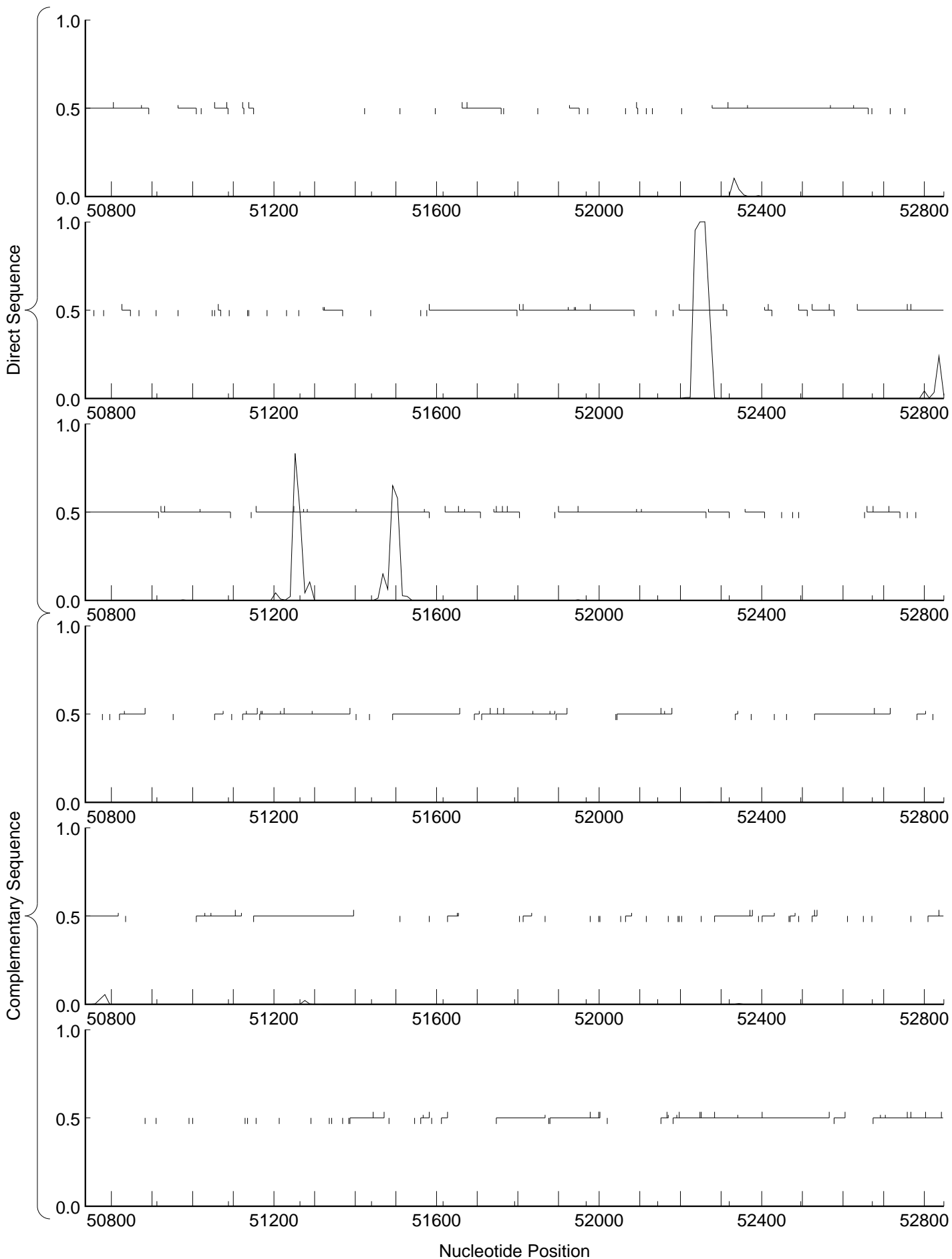
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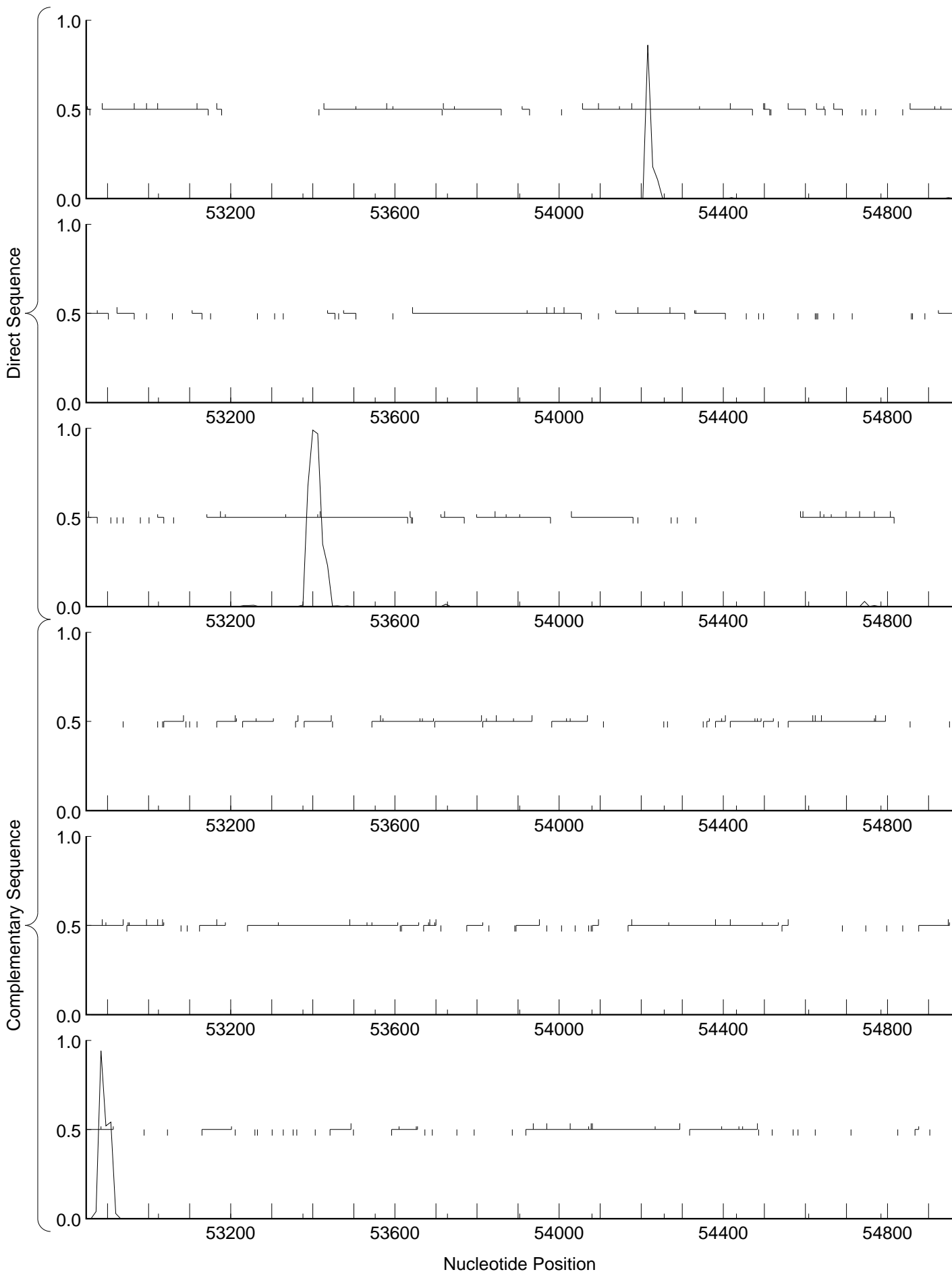




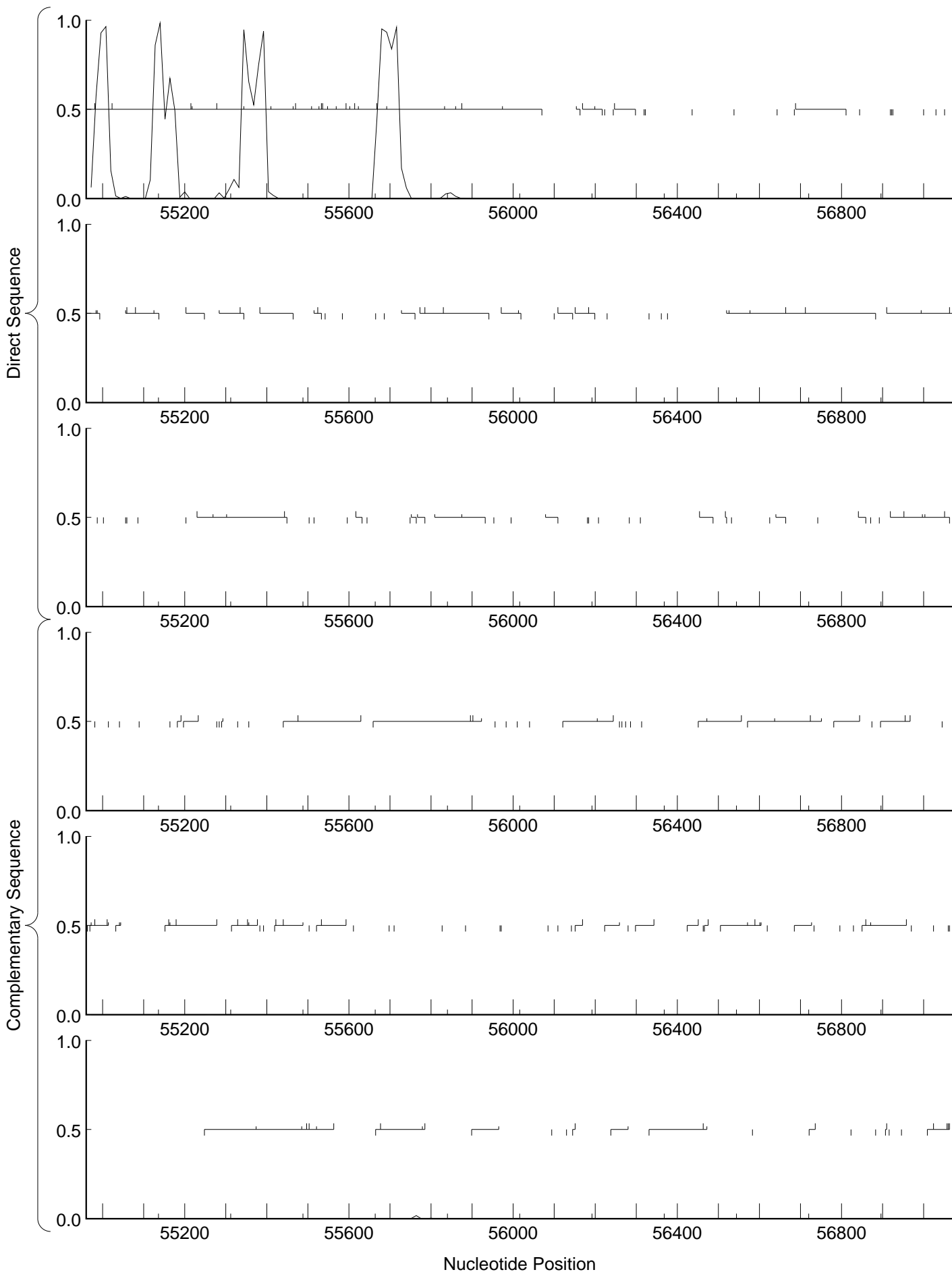


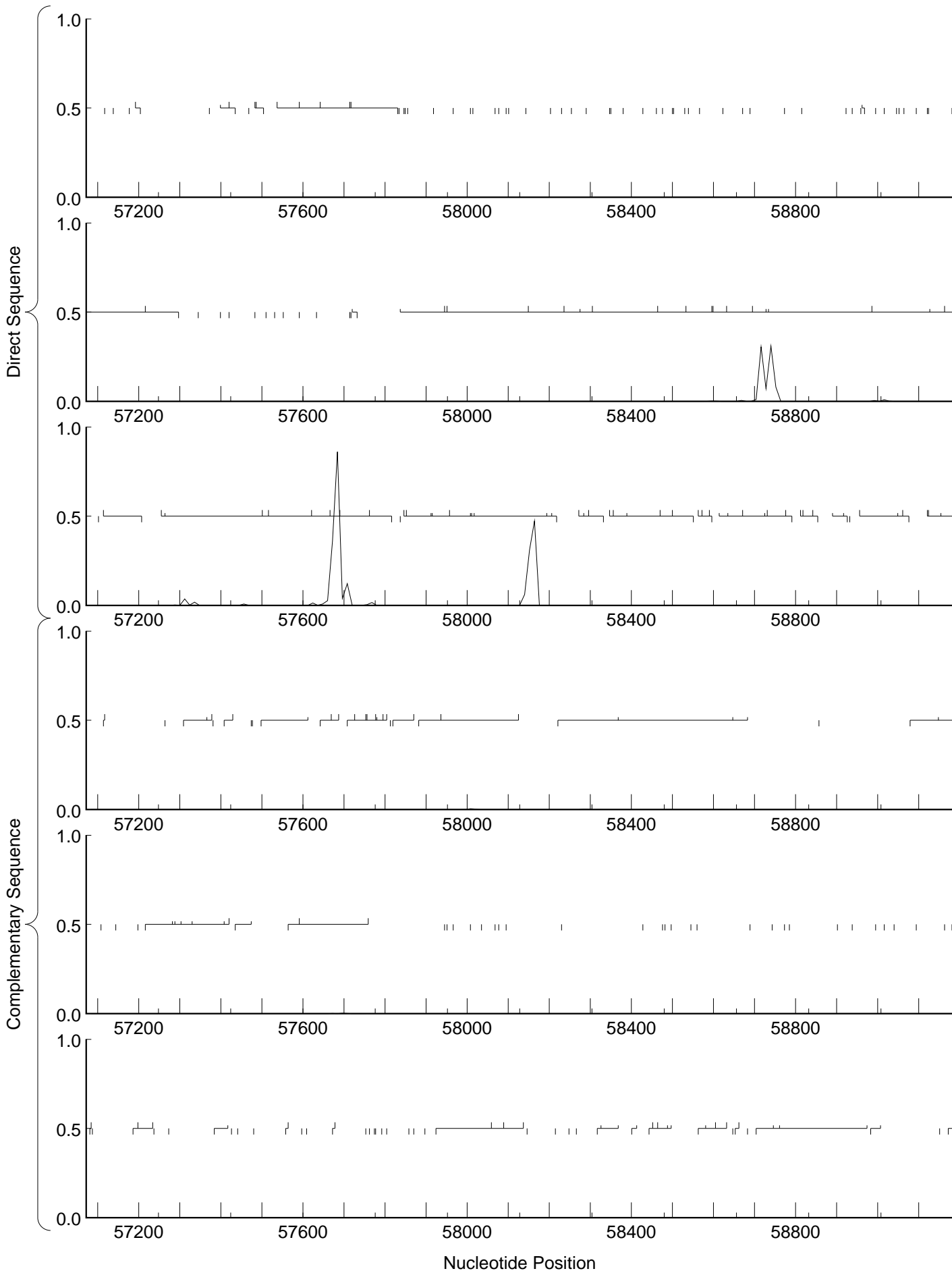






ard complete sequence, 60978 bp including 9-base 3' overhang (CGCCGCCCT), Cluster DJ, Order 4, Window 96, Step 12, 28/30





ard complete sequence, 60978 bp including 9-base 3' overhang (CGCCGCCCT), Cluster DJ, Order 4, Window 96, Step 12, 30/30

