

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Gordonia phage Sopespian complete sequence, 46086 bp including 10-base 3' overhang (CGGTAGG  
Analysis Date : 8/13/24 at 1:43:17  
Pages : 23  
Sequence Length : 46086 bp  
GC Content : 60.17%

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,

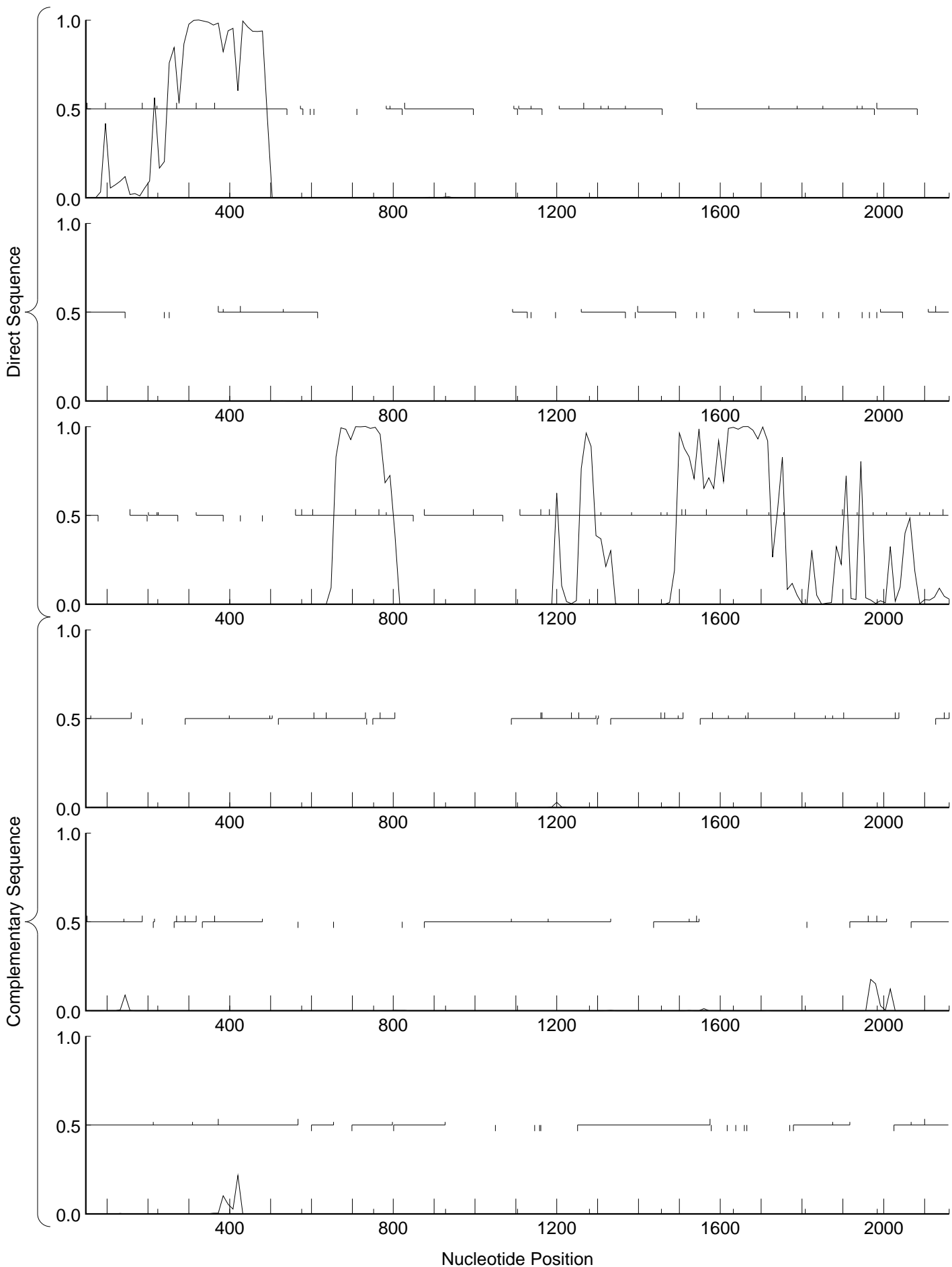
## MATRIX INFORMATION

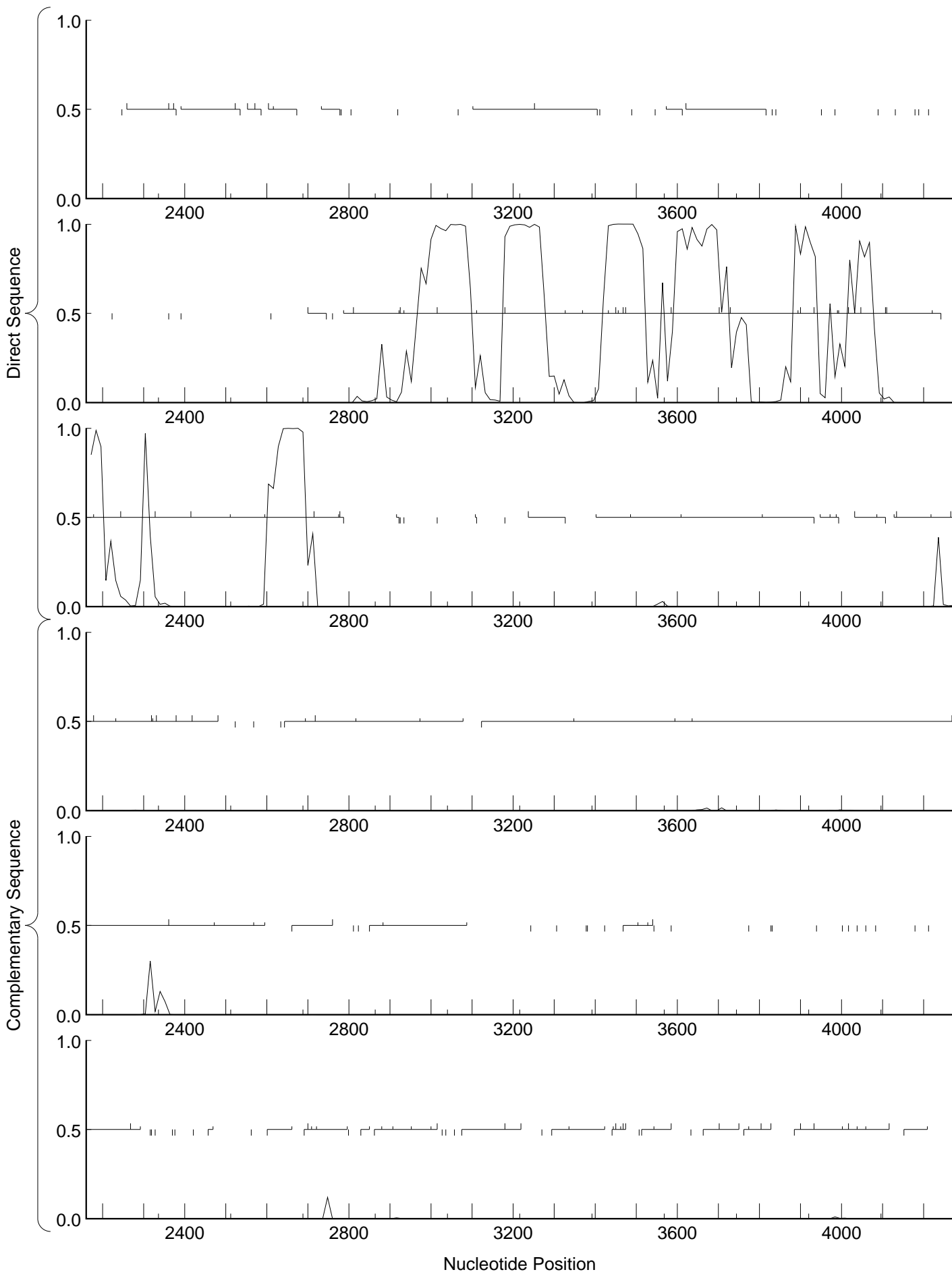
Matrix : -  
Author : -  
Order : 4

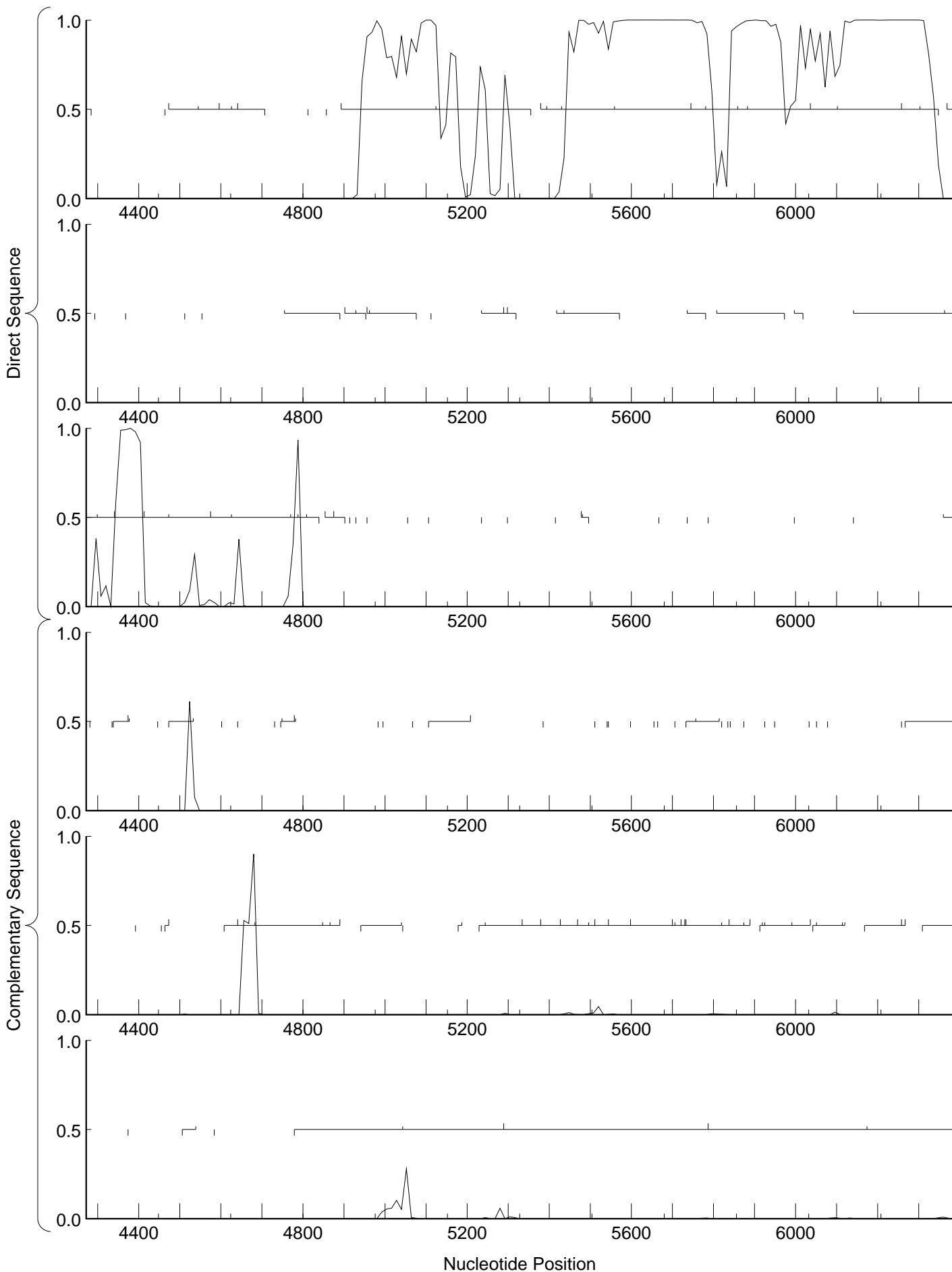
Send questions / comments to:  
Dr. M. Borodovsky  
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Atlanta, GA 30332-0230

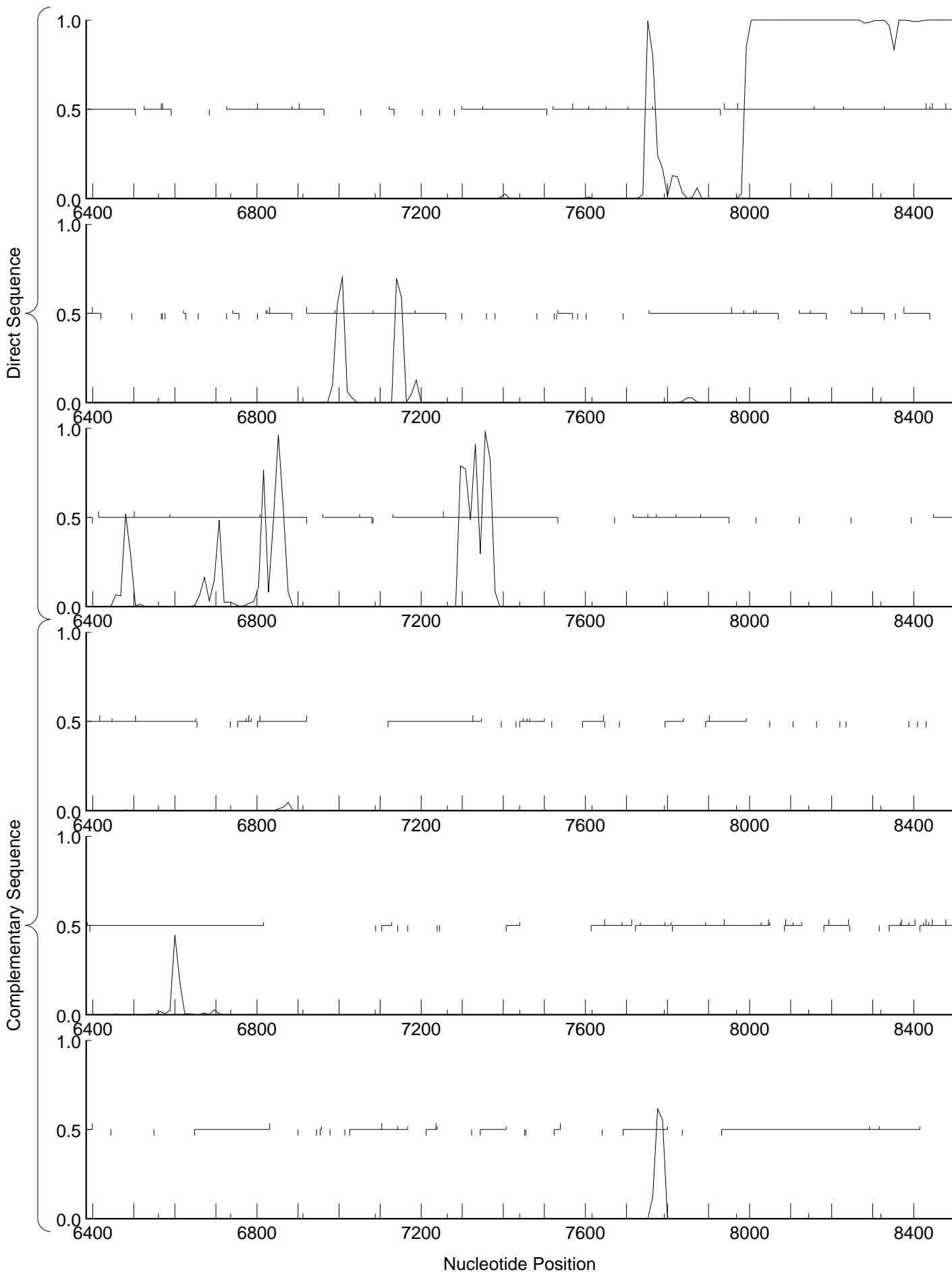
### Matrix notes & comments

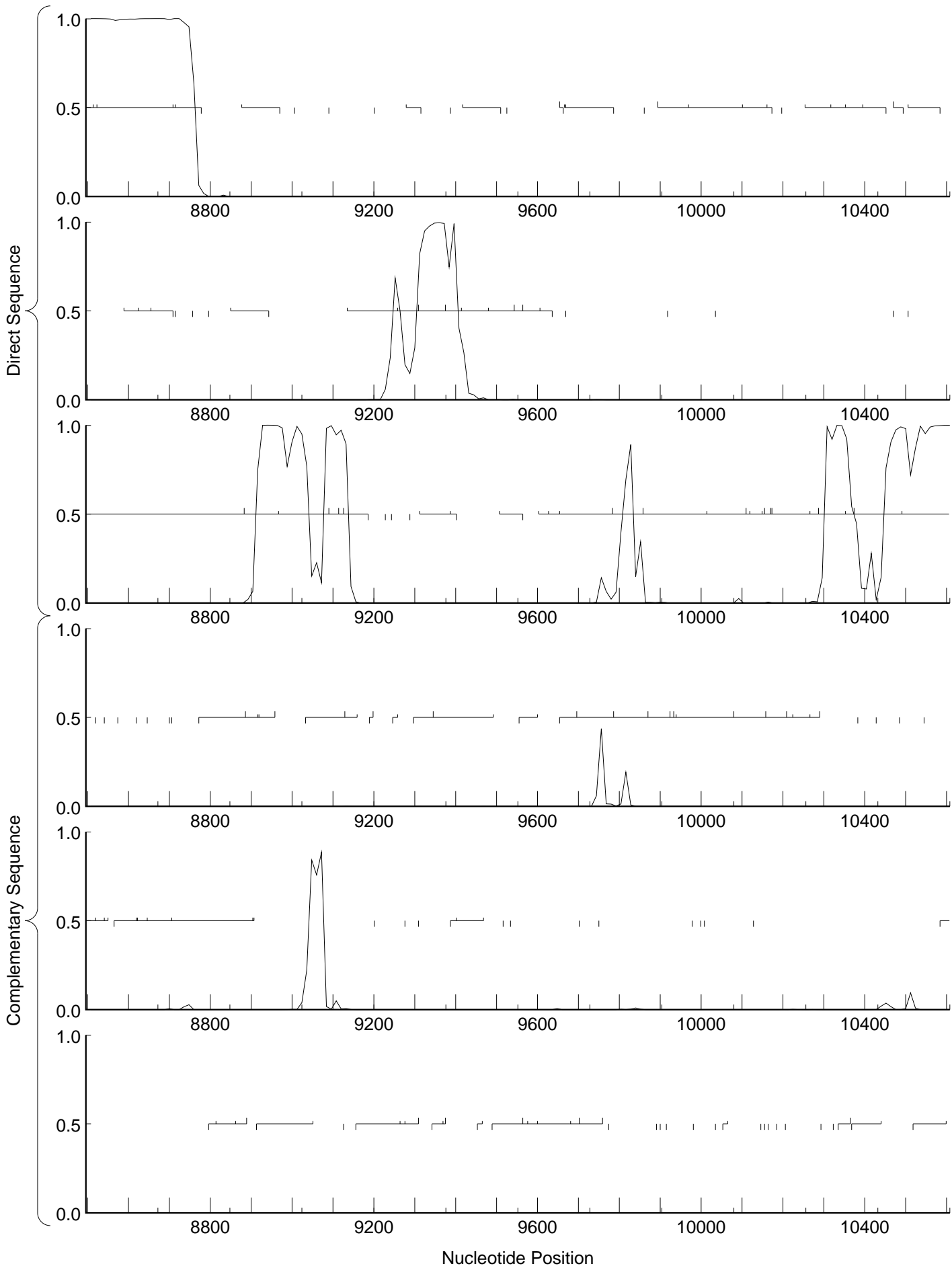
Training set derived by GeneMarkS, 4.30 October 2014  
Sat Sep 8 00:36:14 2018

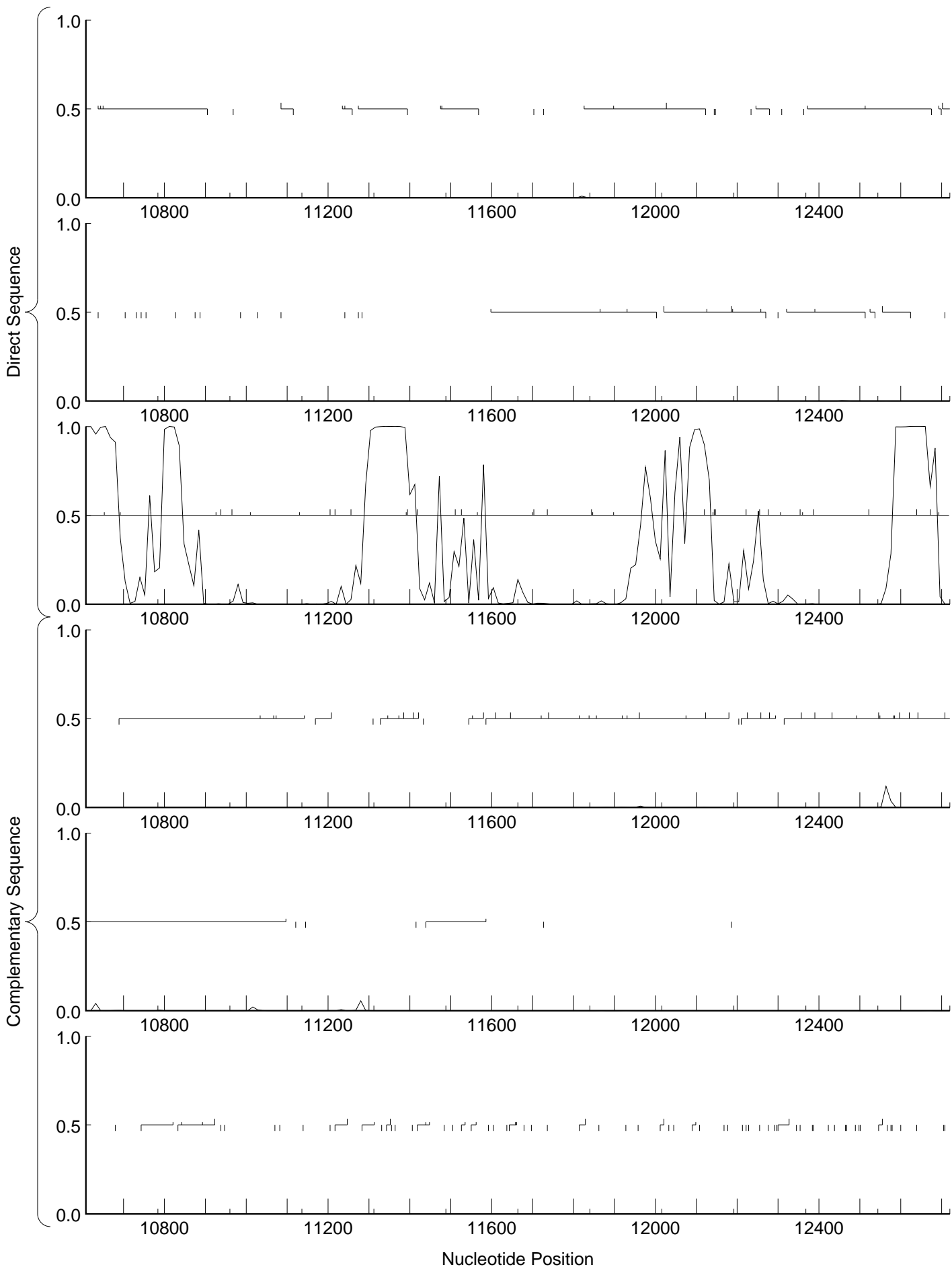


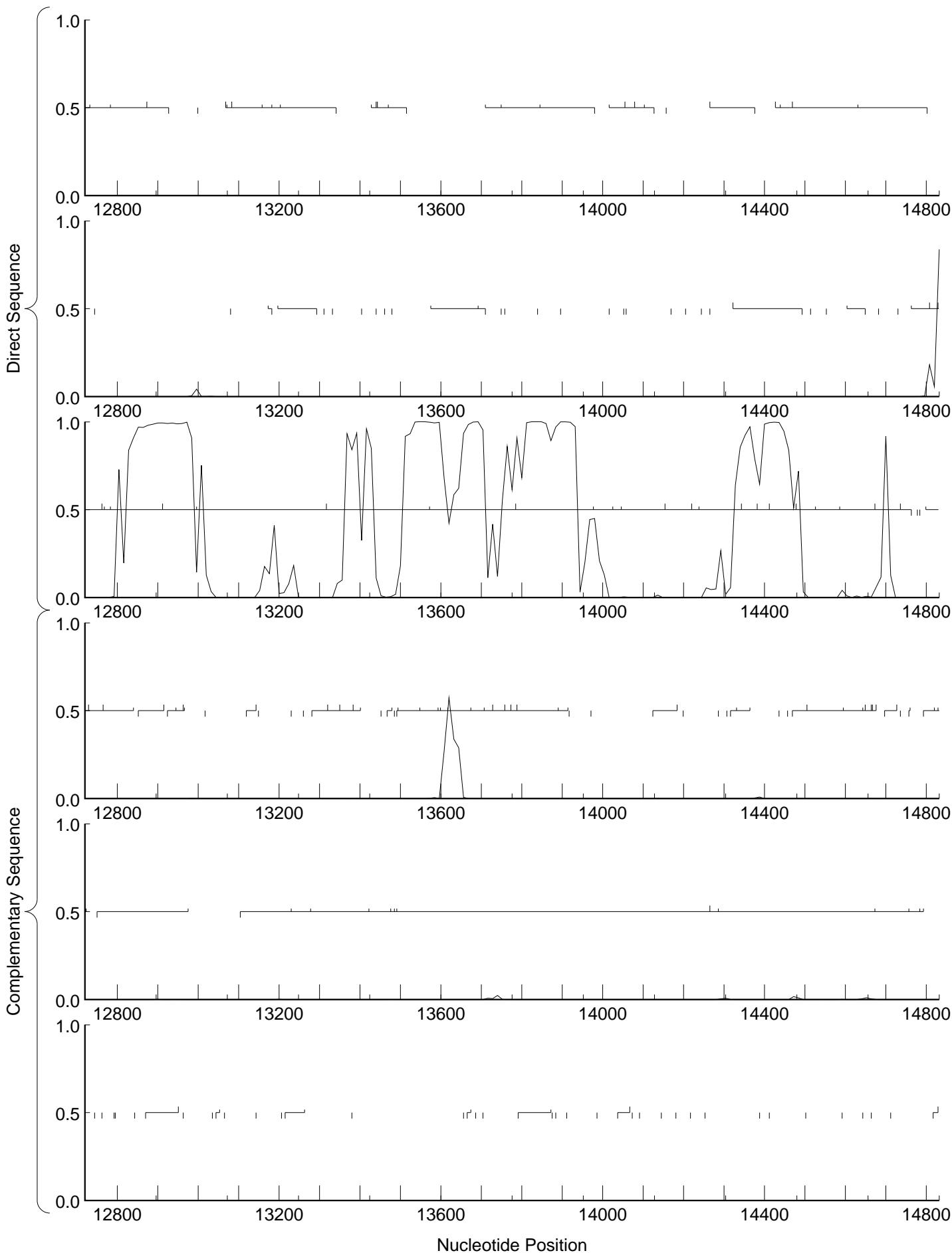




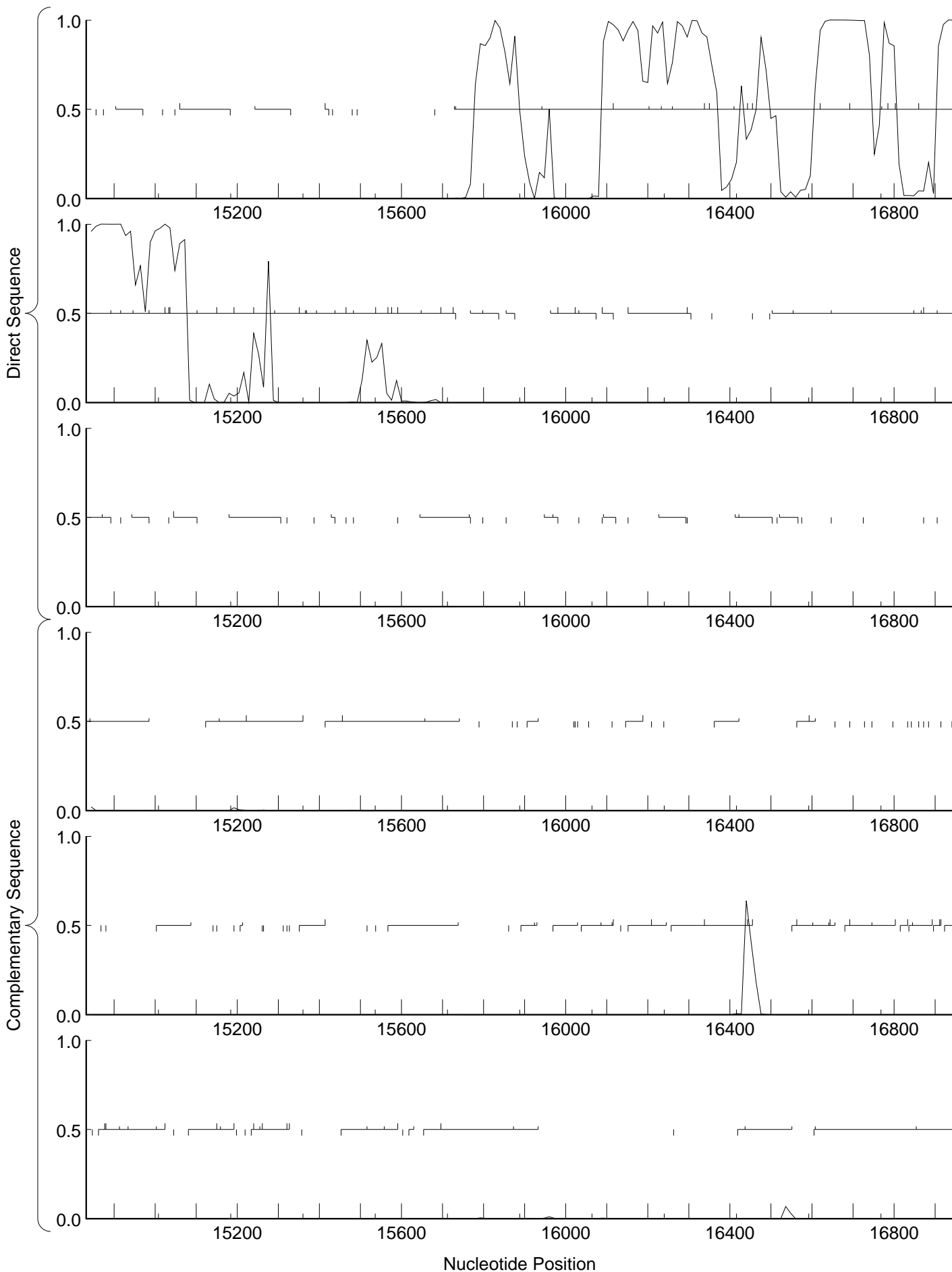




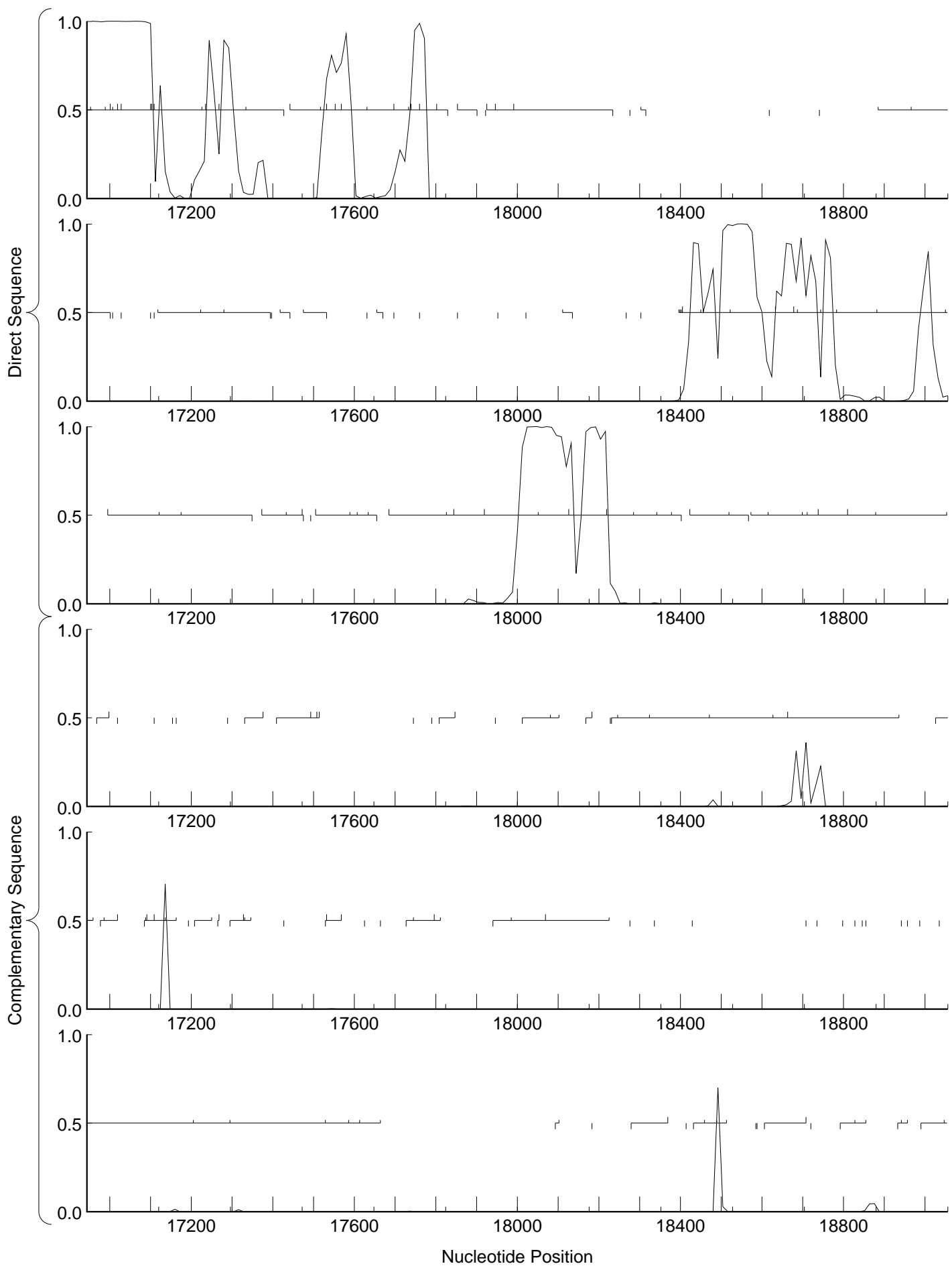


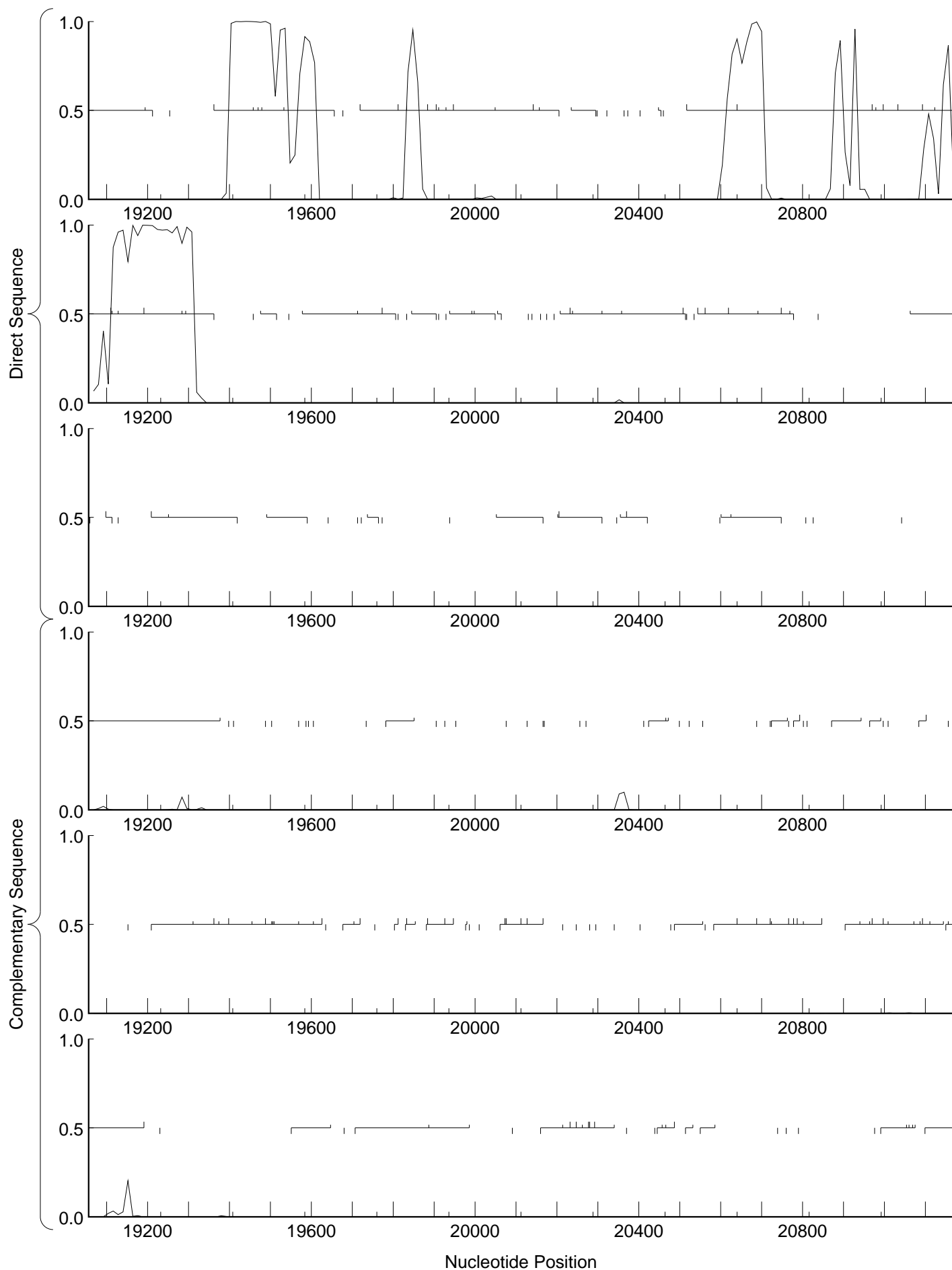




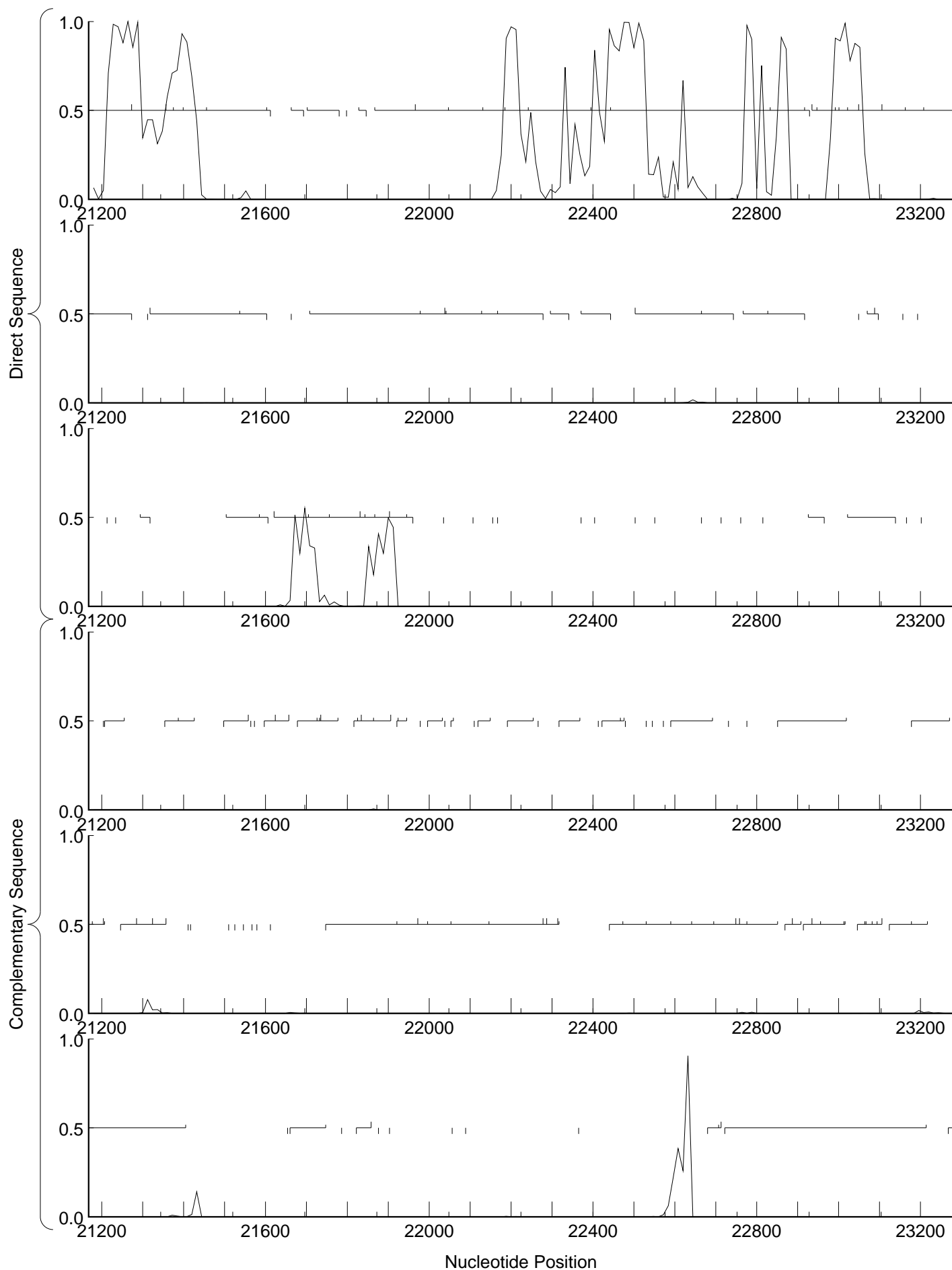


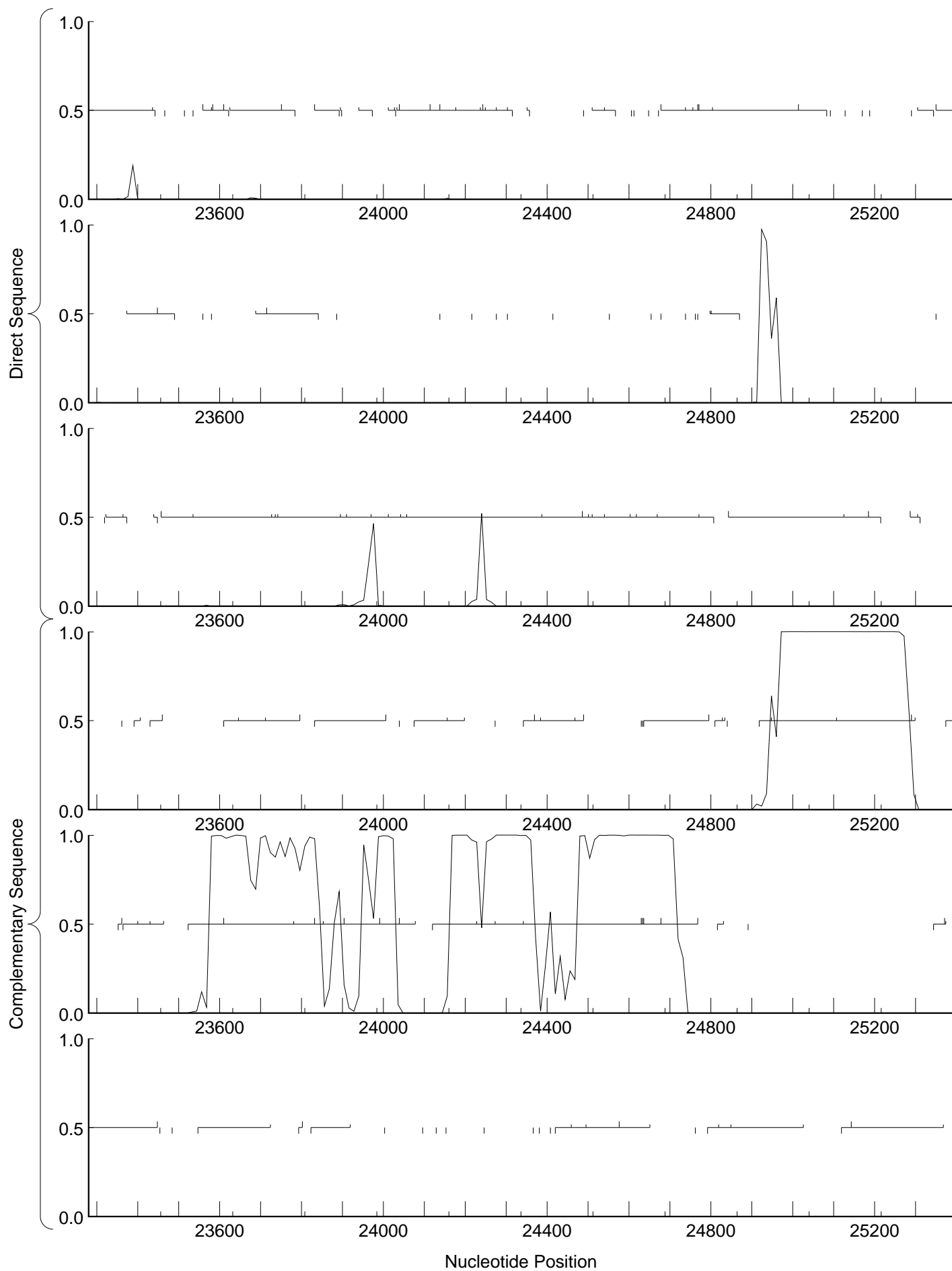
complete sequence, 46086 bp including 10-base 3' overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 10/23

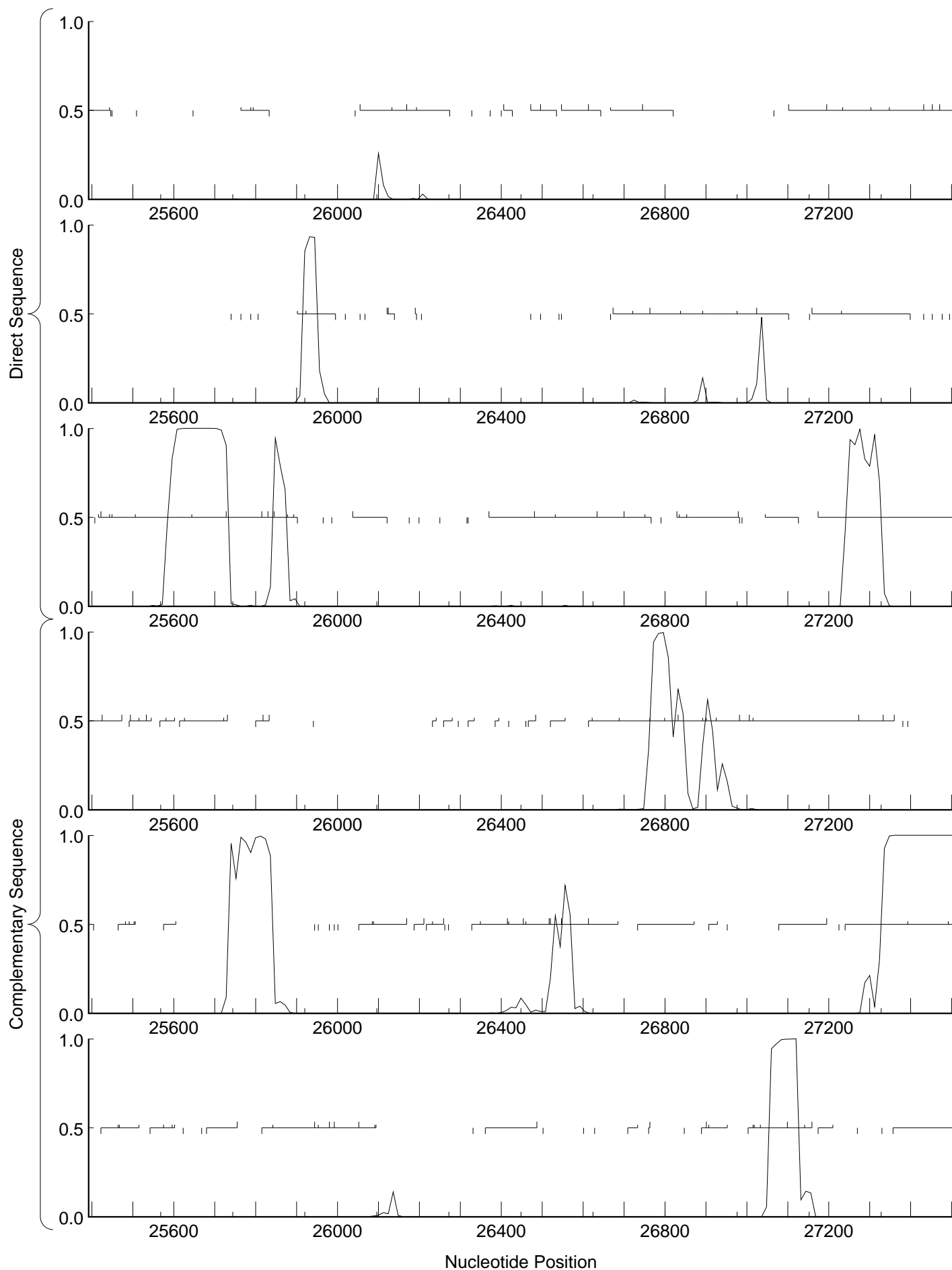




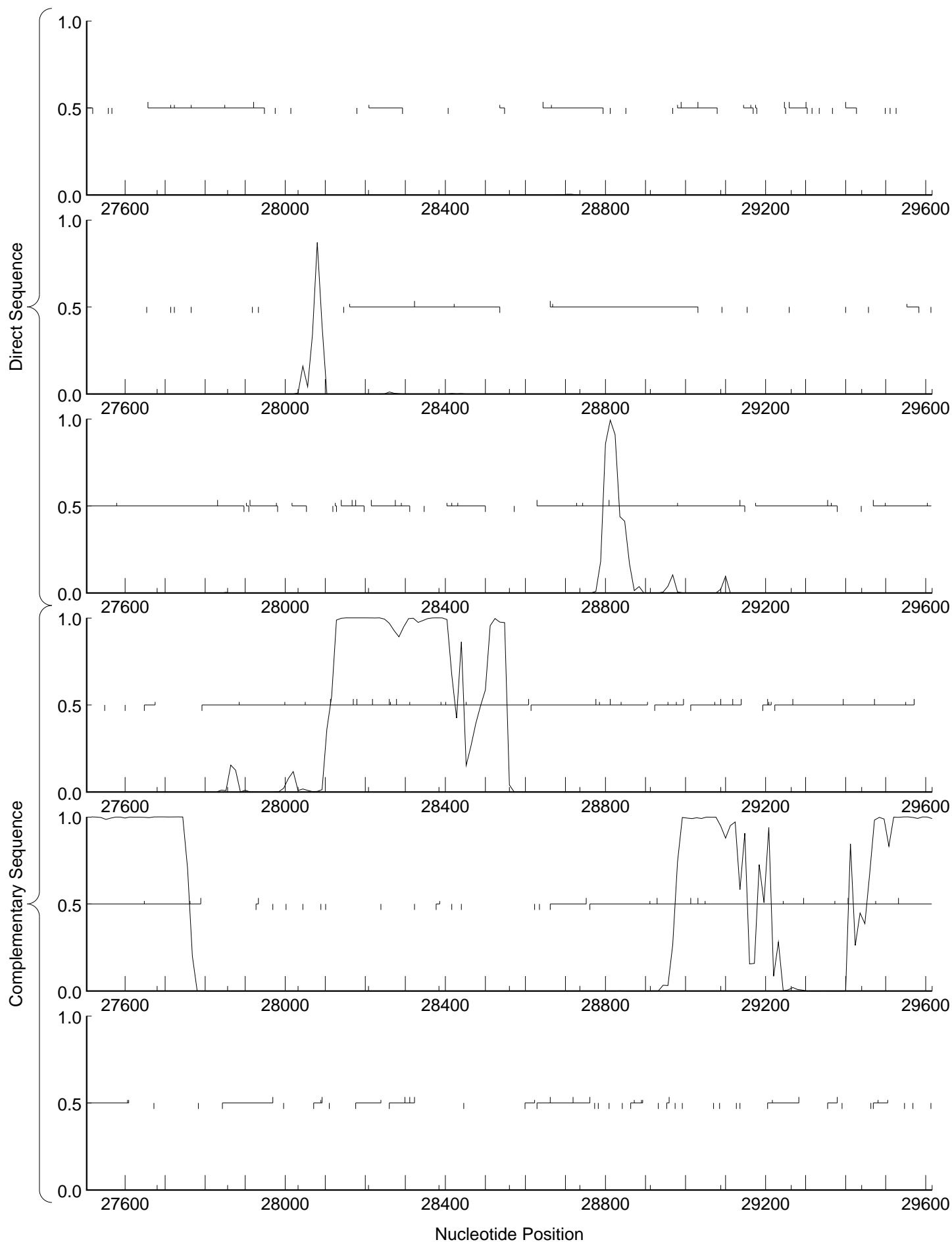
complete sequence, 46086 bp including 10-base 3' overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 12/23



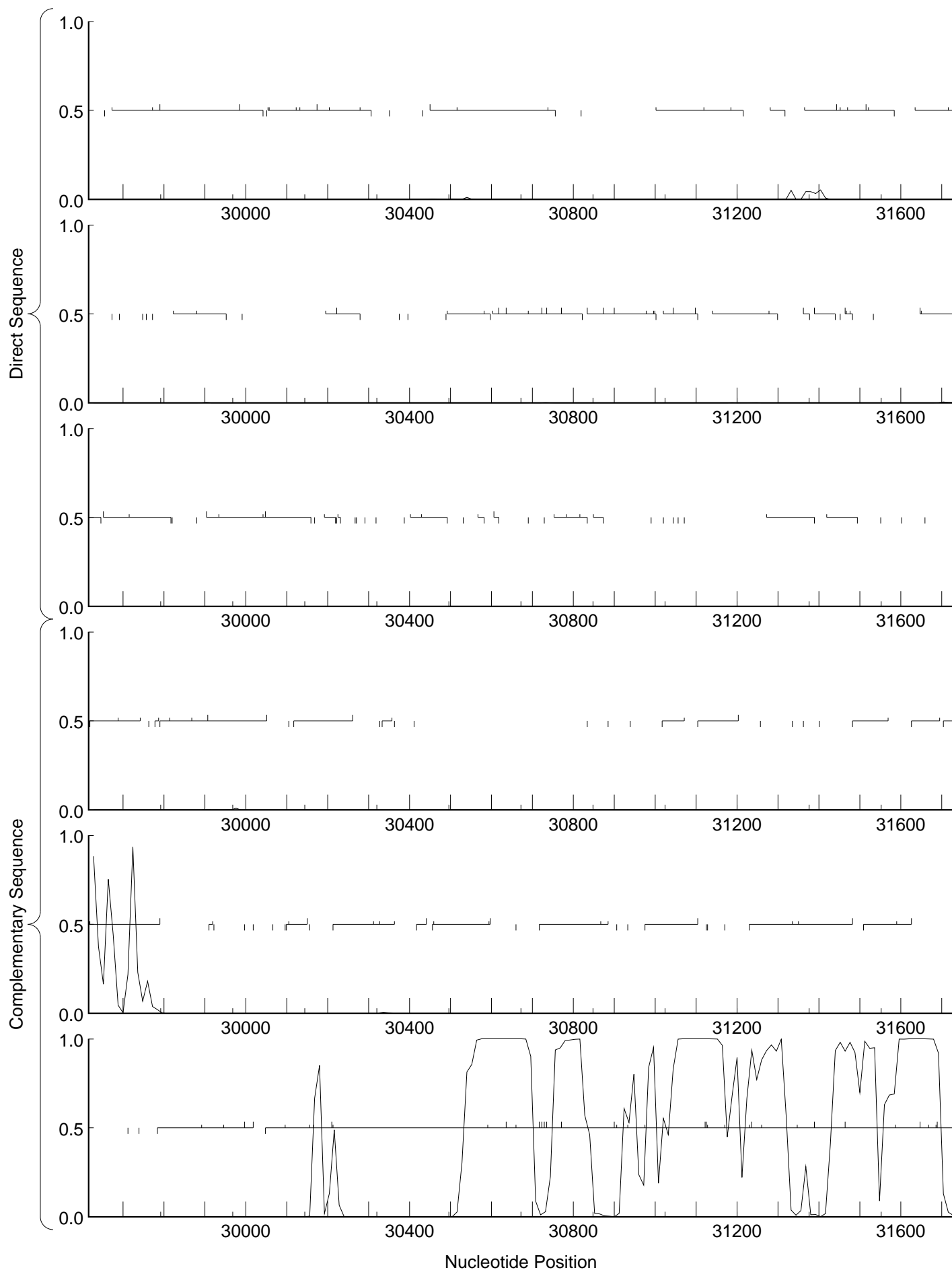




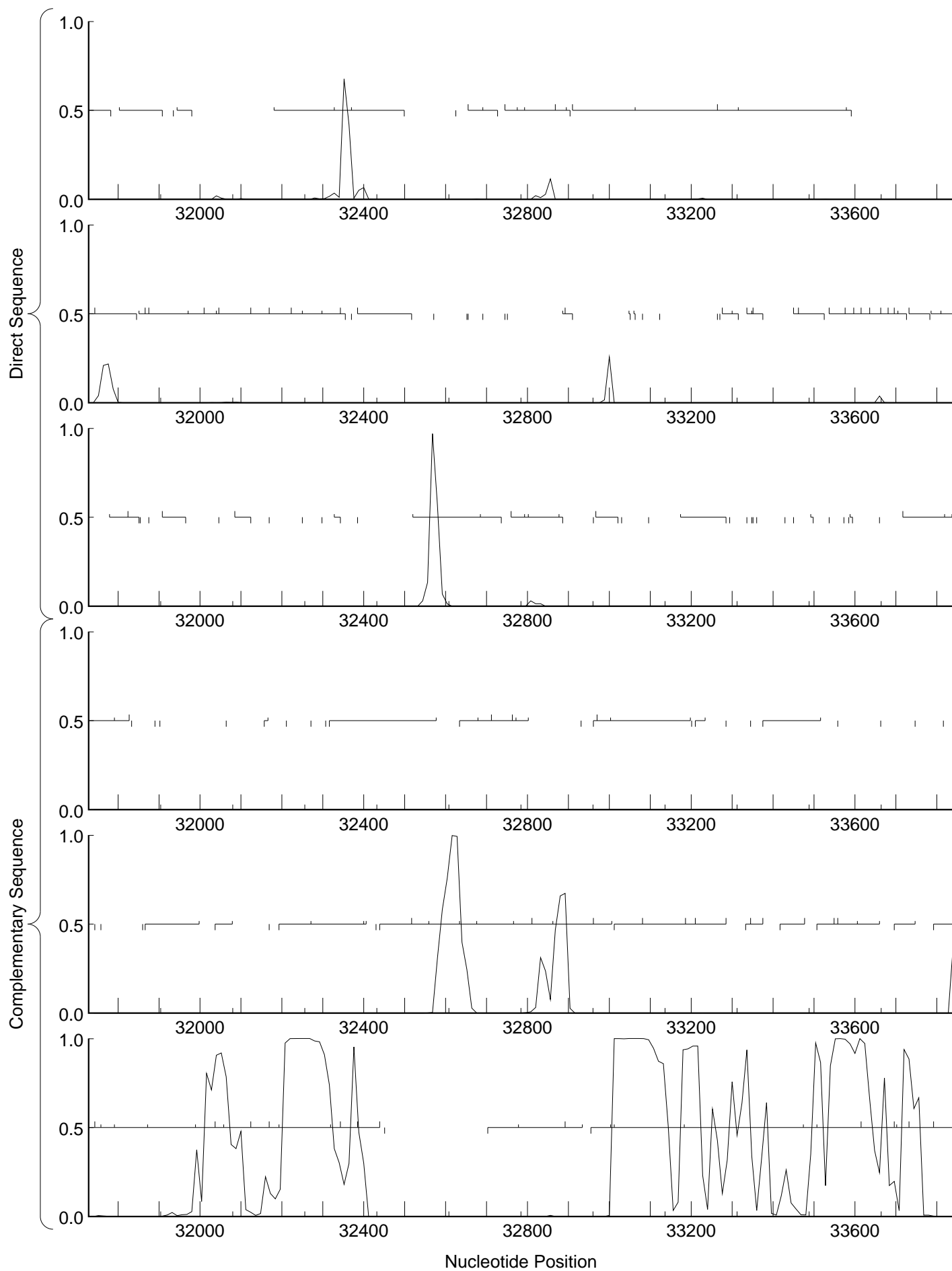
complete sequence, 46086 bp including 10-base 3' overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 15/23

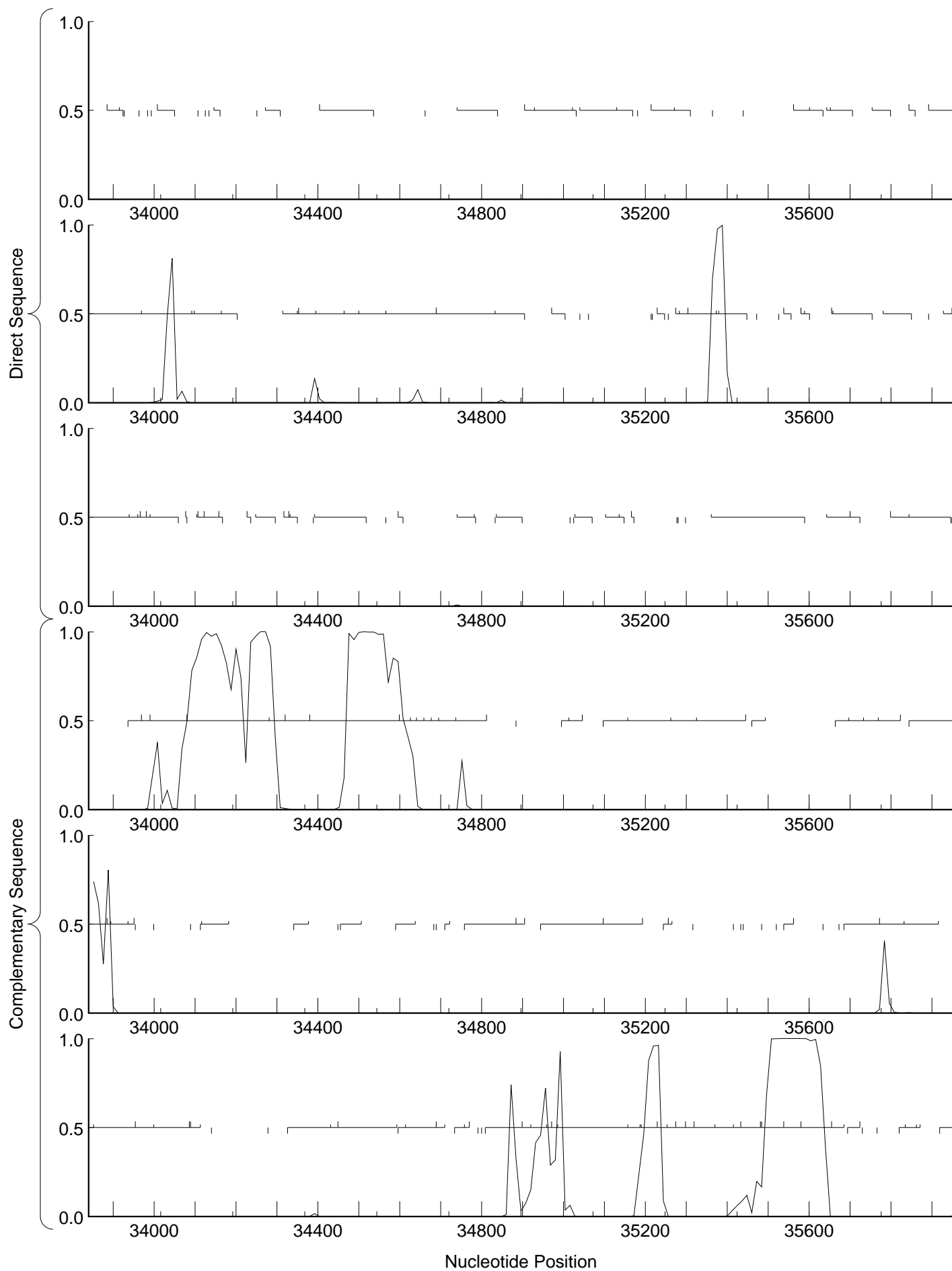


complete sequence, 46086 bp including 10-base 3' overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 16/23

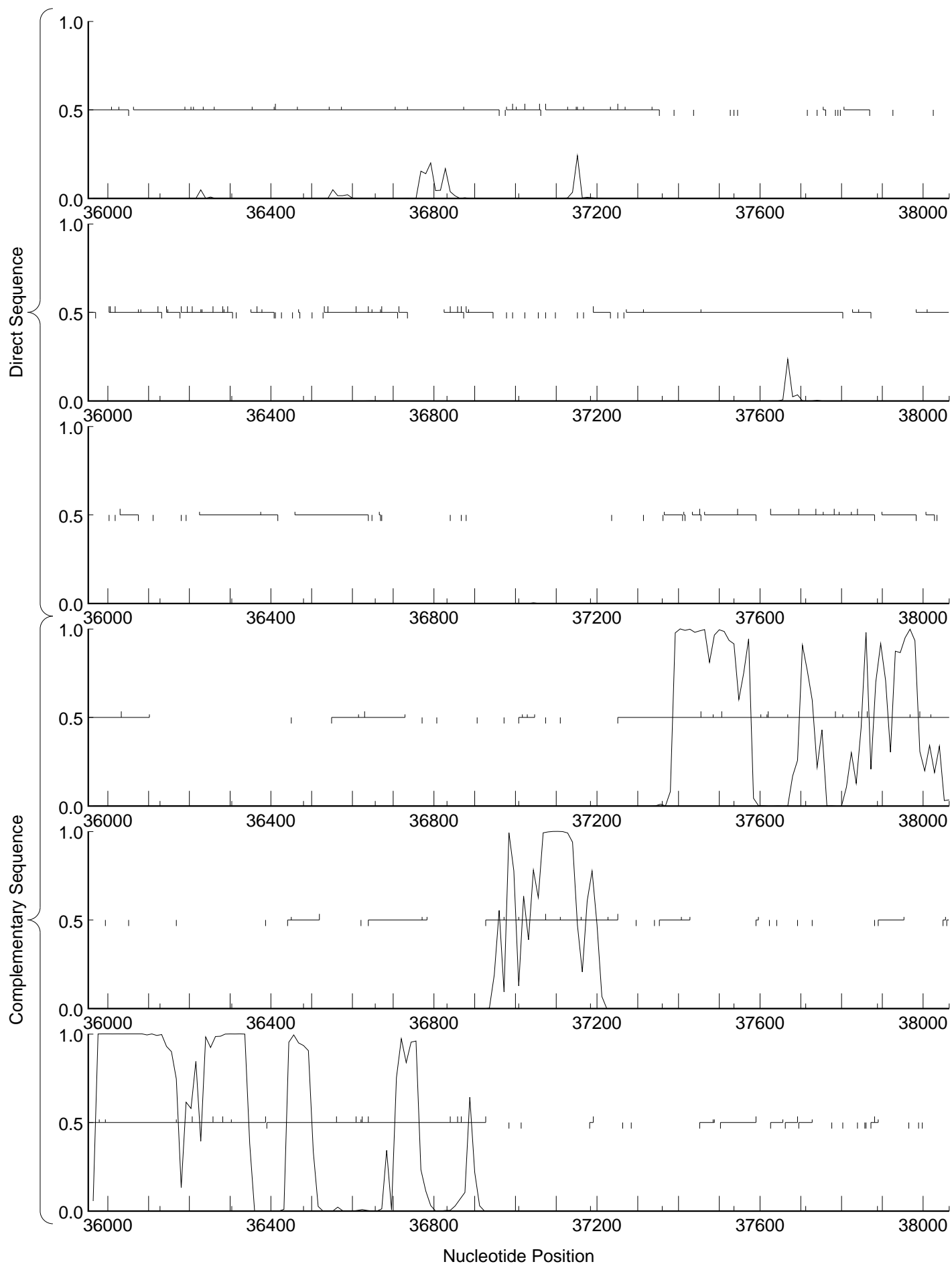


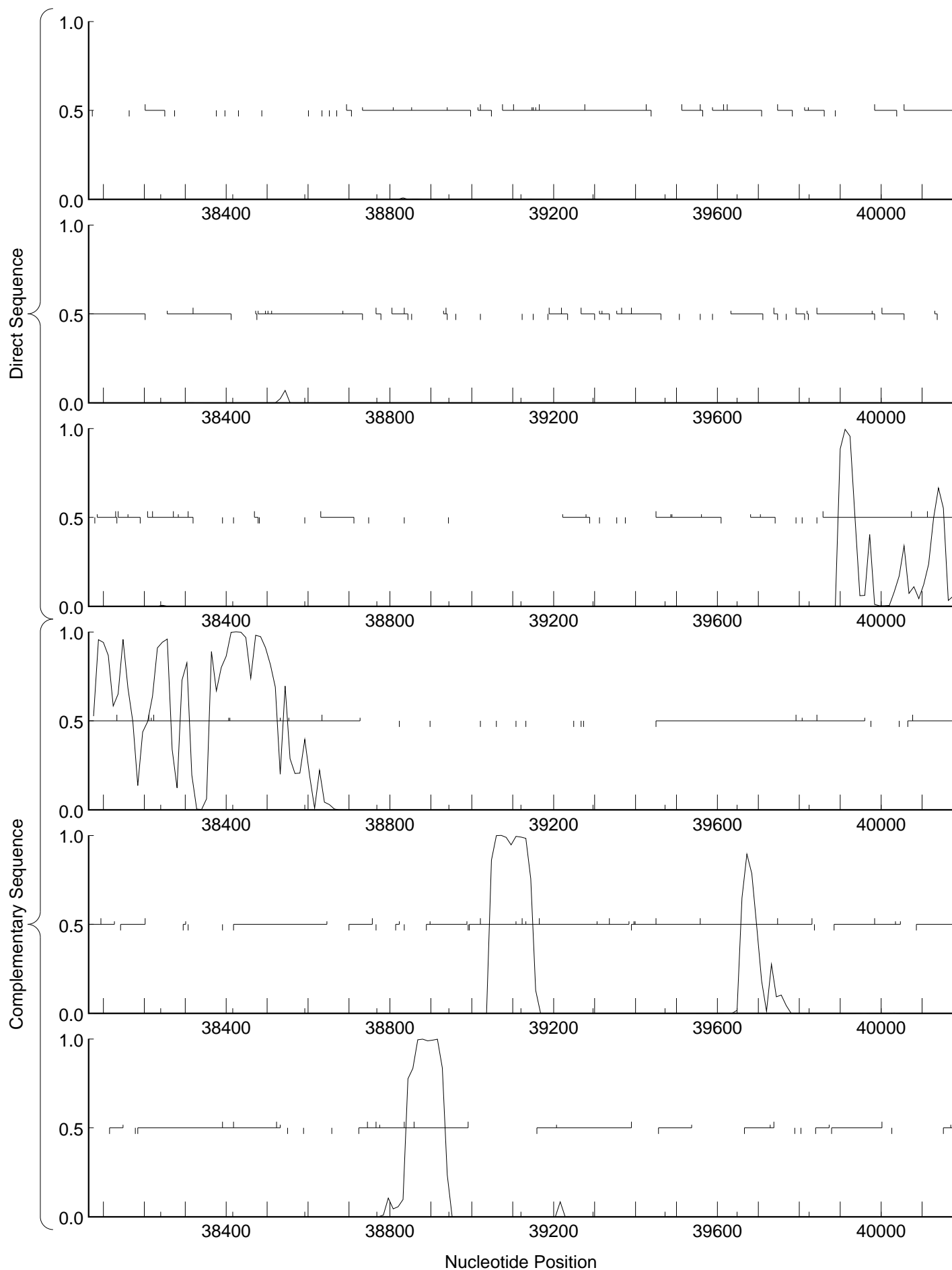




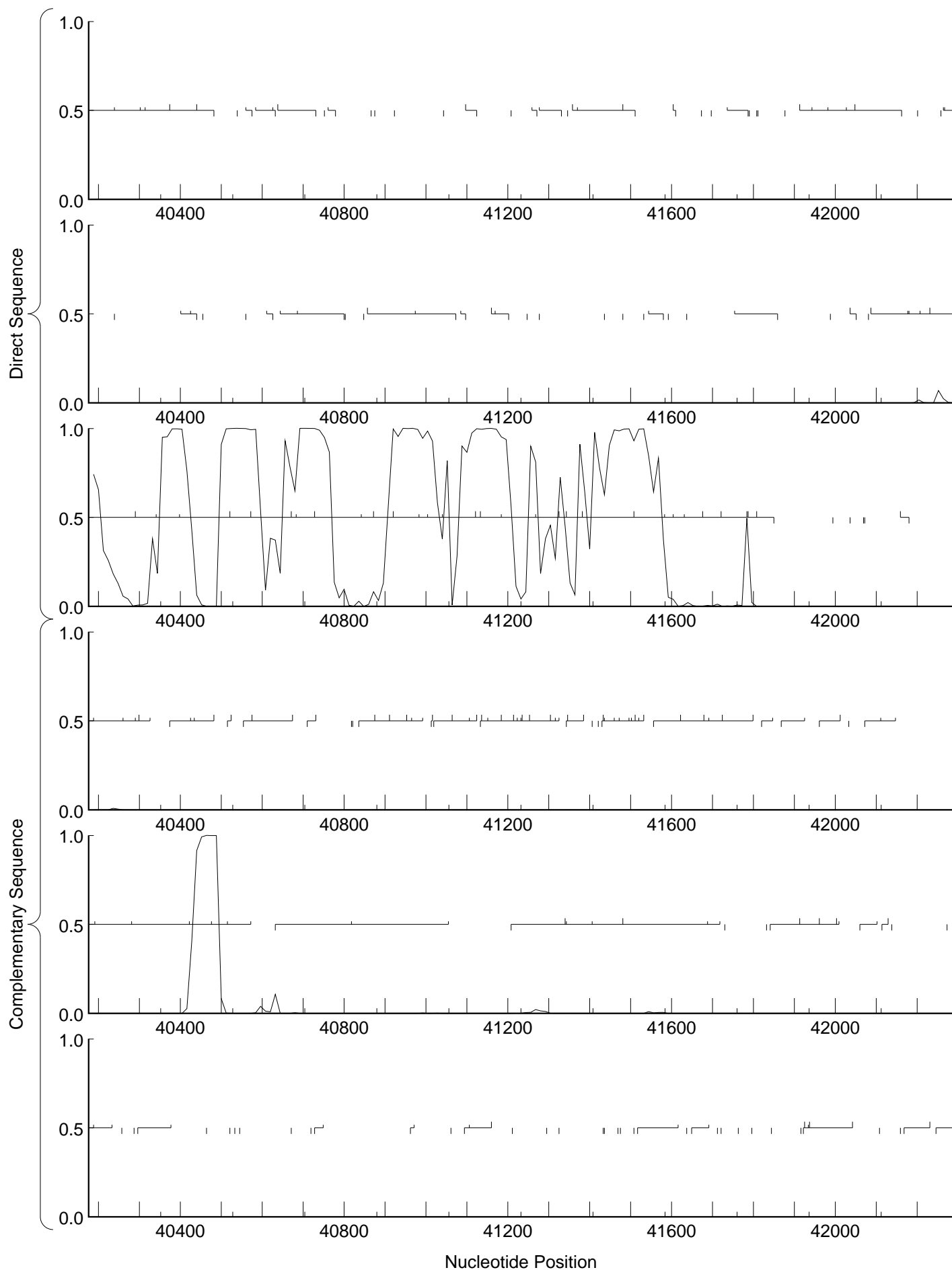


complete sequence, 46086 bp including 10-base 3' overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 19/23





complete sequence, 46086 bp including 10-base 3' overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 21/23



complete sequence, 46086 bp including 10-base 3' overhang (CGGTAGGCTT), Cluster CT, Order 4, Window 96, Step 12, 22/23

