

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Mycobacterium phage MilanaBonita complete sequence, 56251 bp including 10-base 3' overhang  
Analysis Date : 3/5/21 at 8:42:36  
Pages : 28  
Sequence Length : 56251 bp  
GC Content : 62.06%

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,

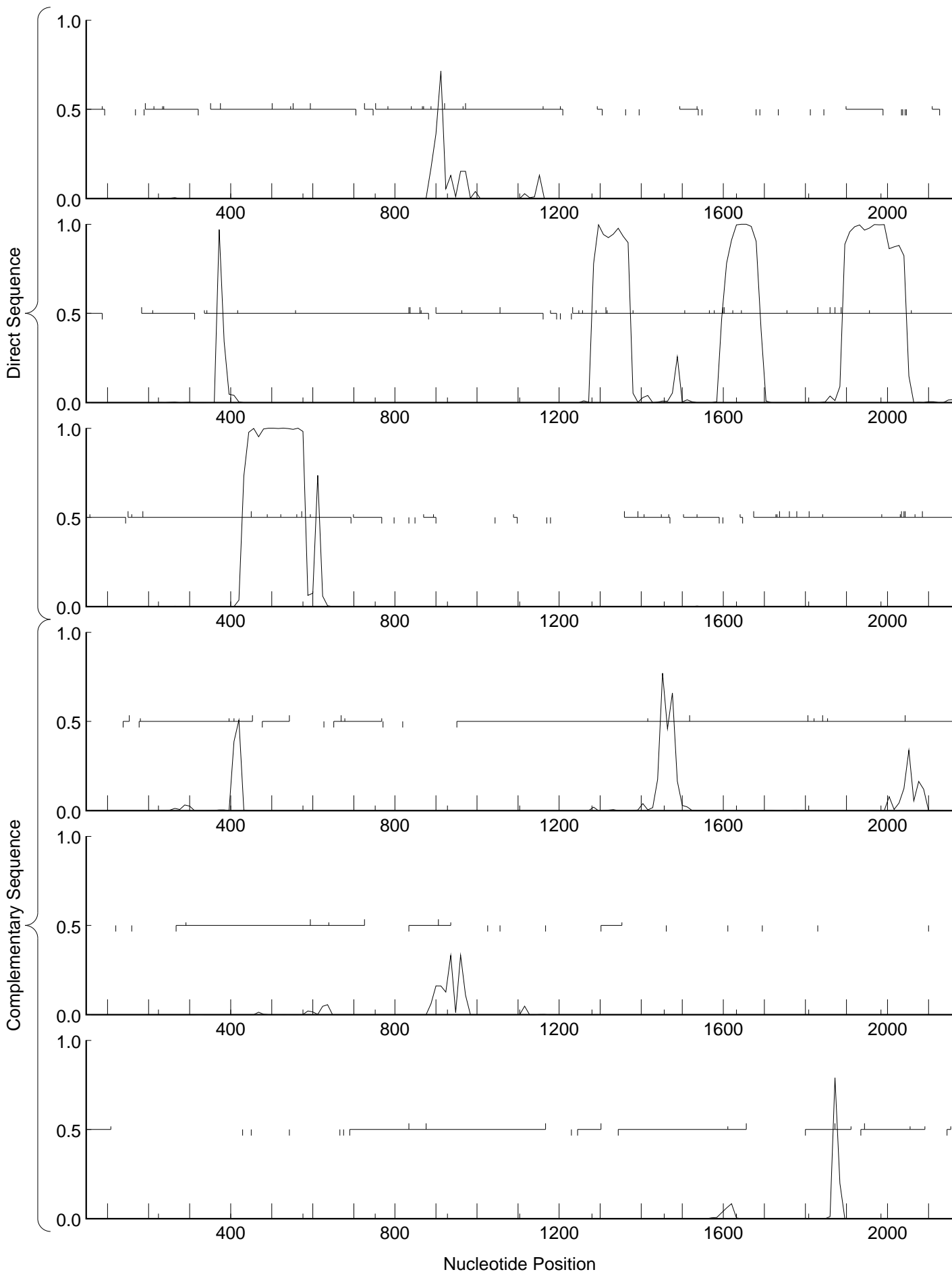
## Matrix notes & comments

Training set derived by GeneMarkS, 4.27 September 2014  
Tue Sep 23 15:23:03 2014

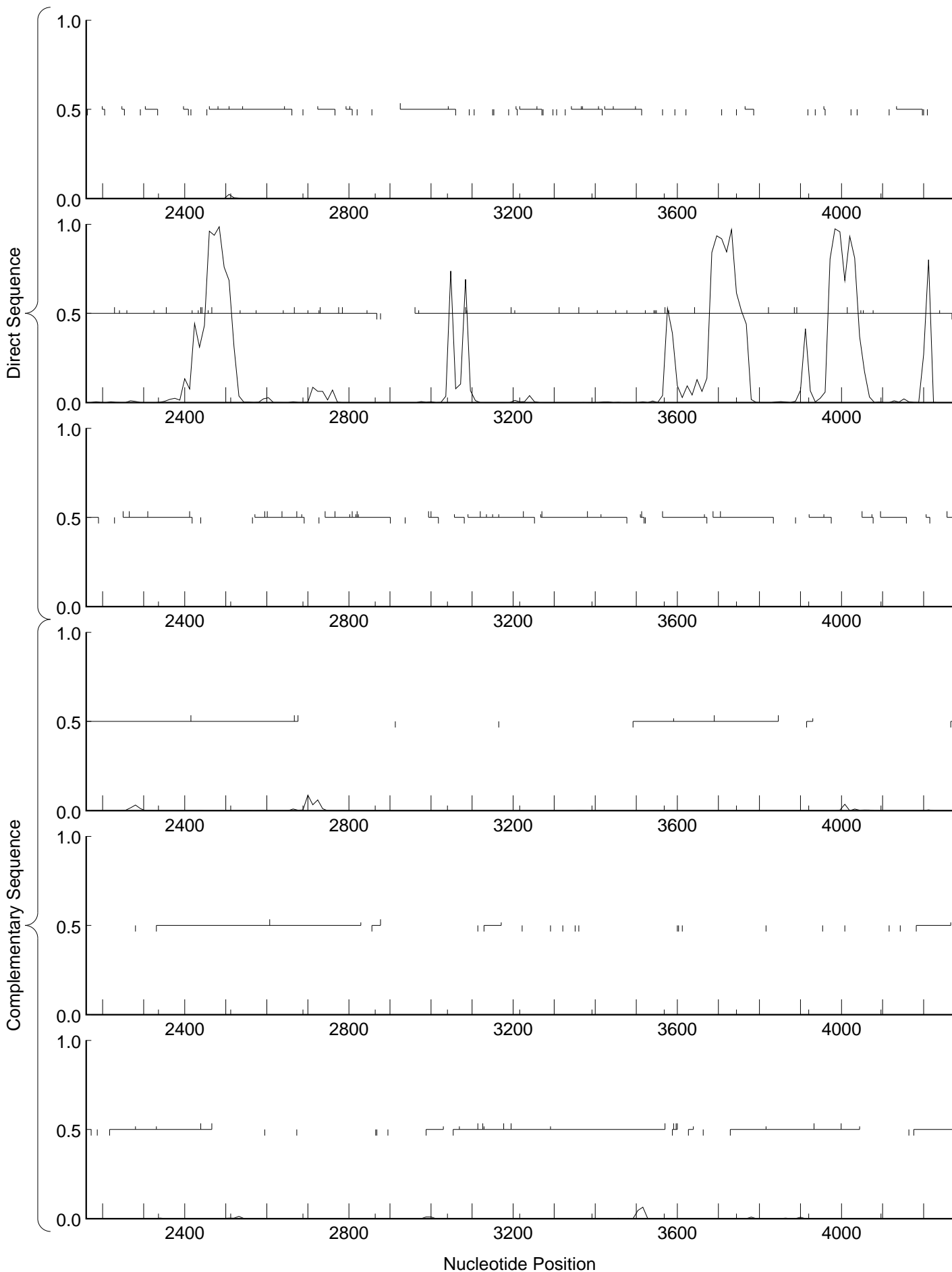
## MATRIX INFORMATION

Matrix : Mycobacterium\_smegmatis\_MC2\_155  
Author : -  
Order : 4

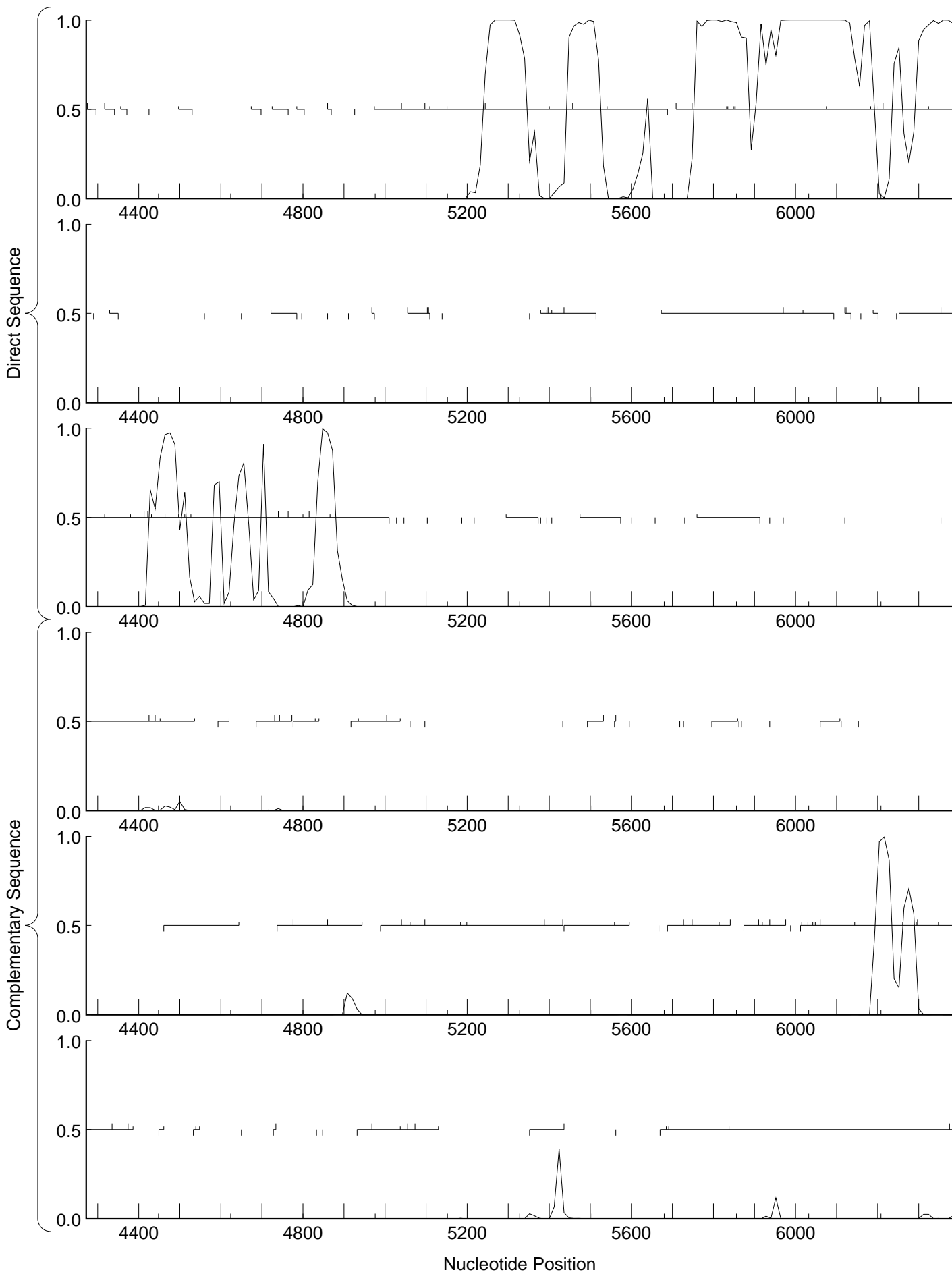
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

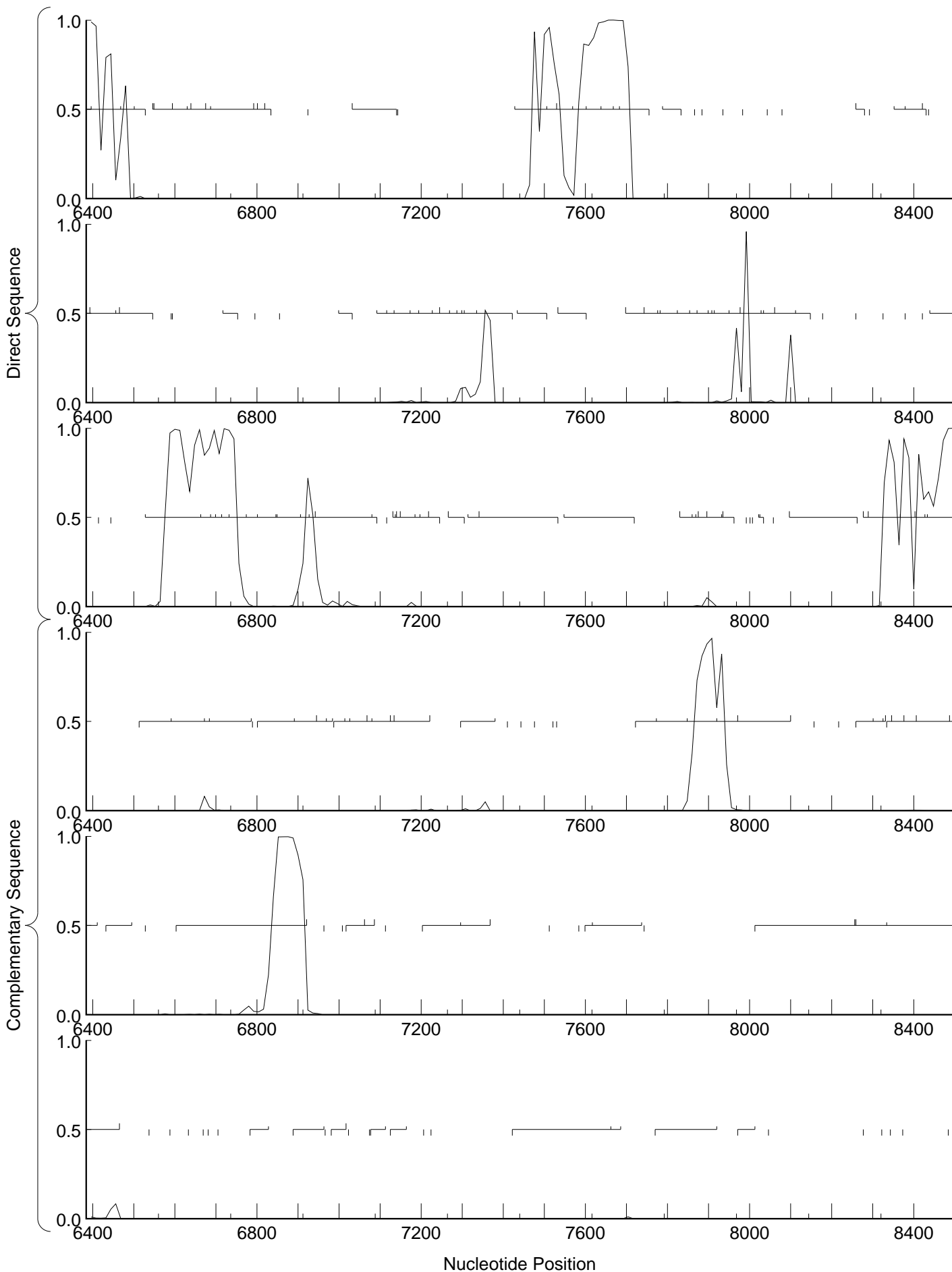


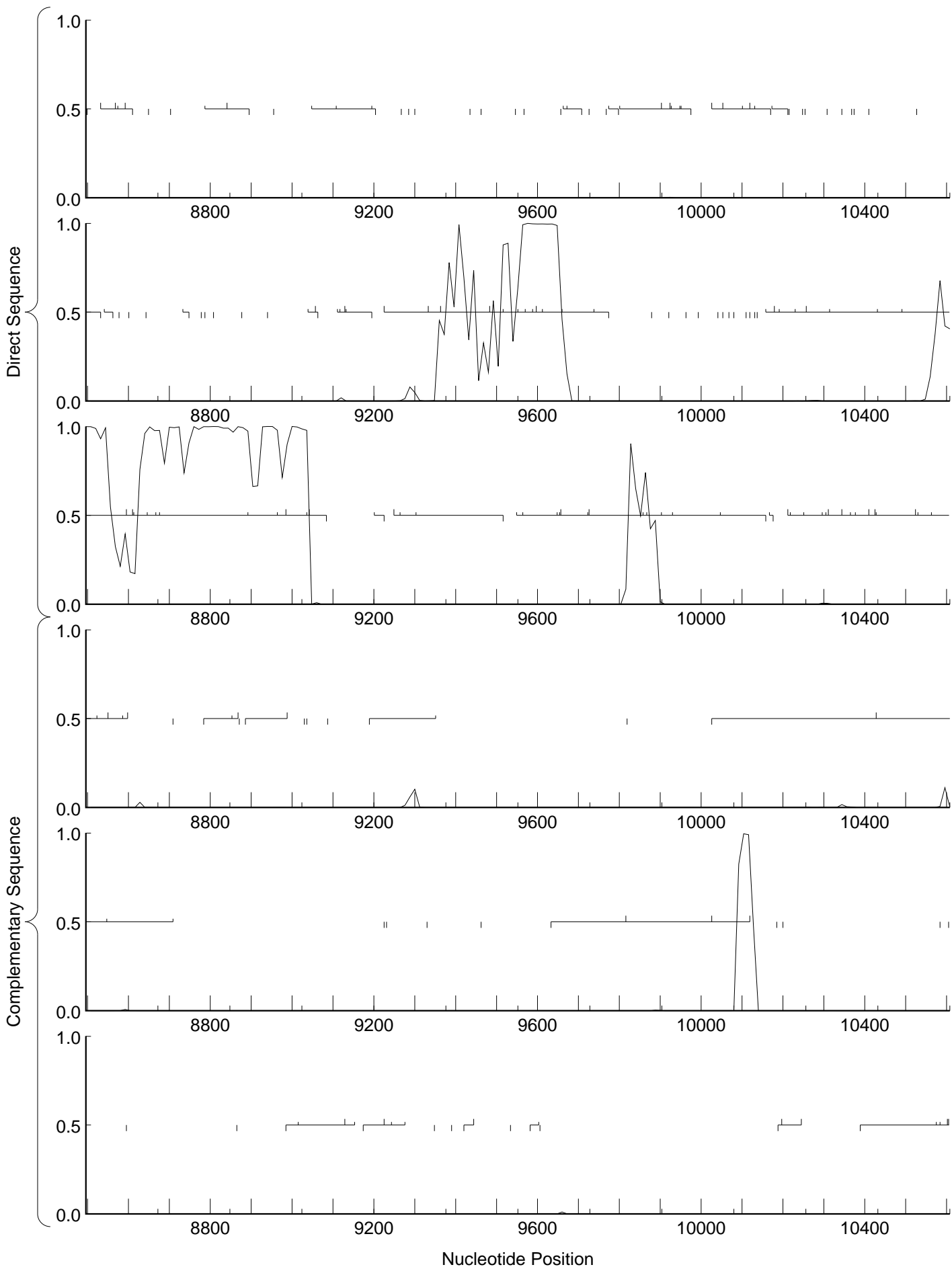
a complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 3/28



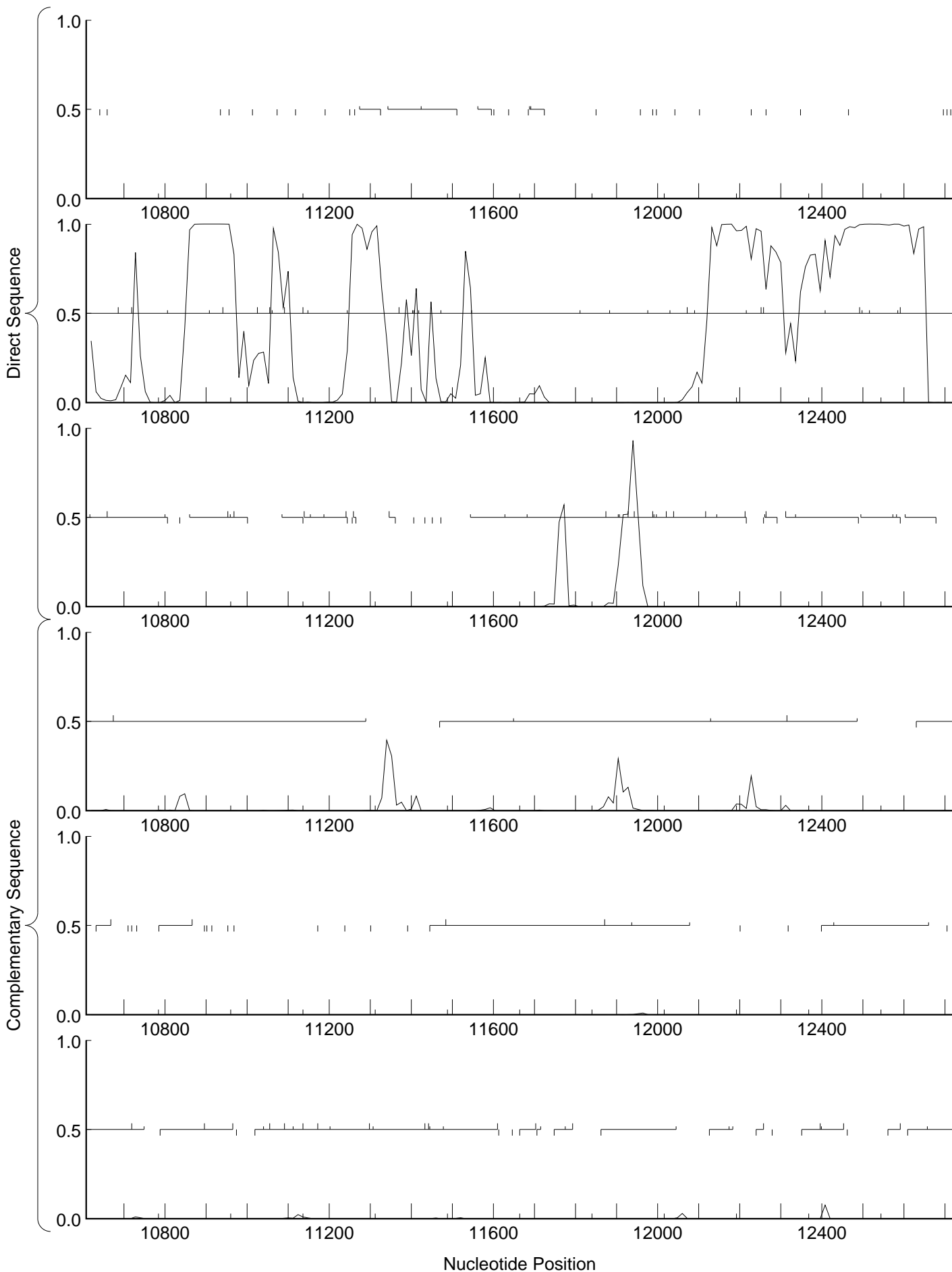
a complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 4/28

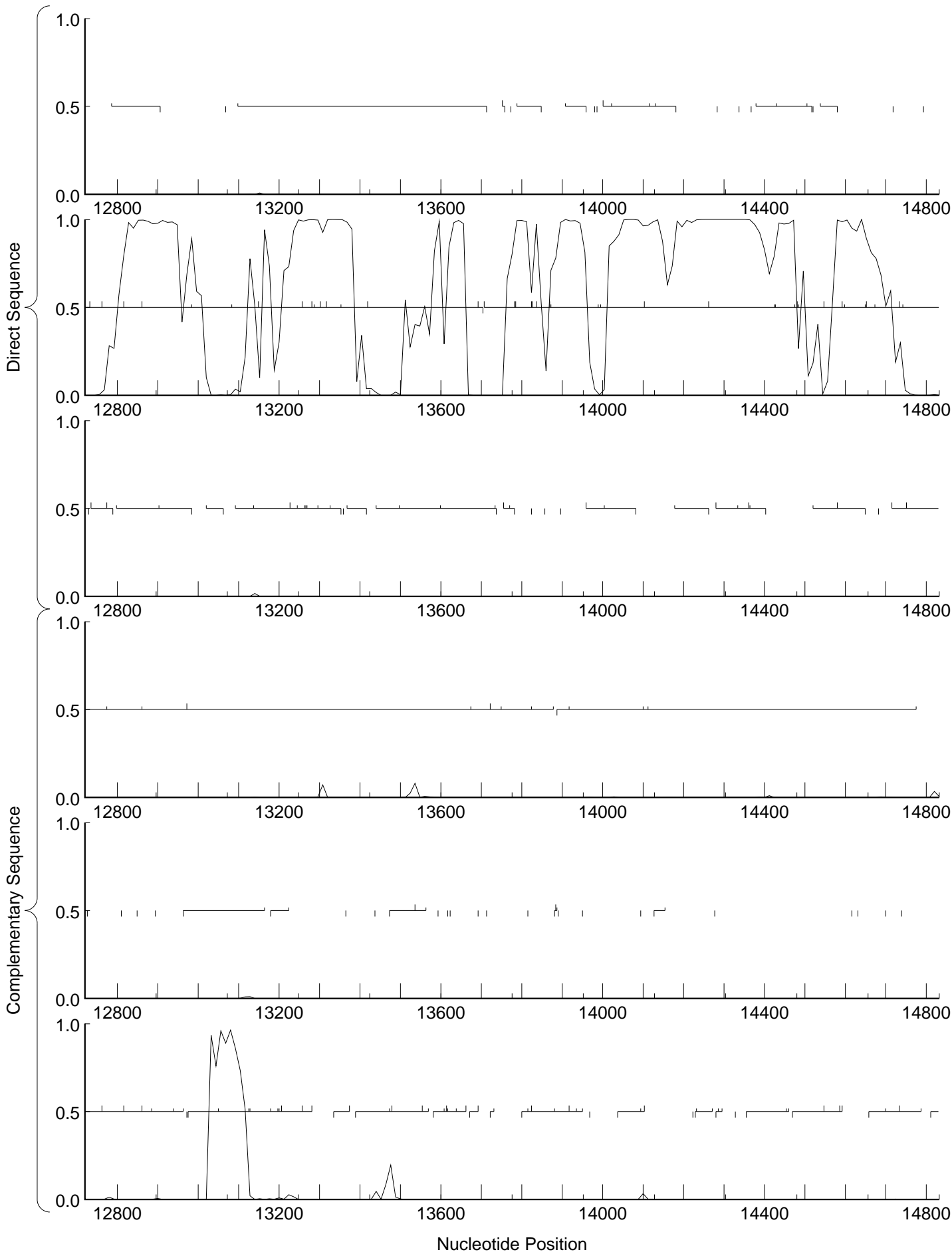




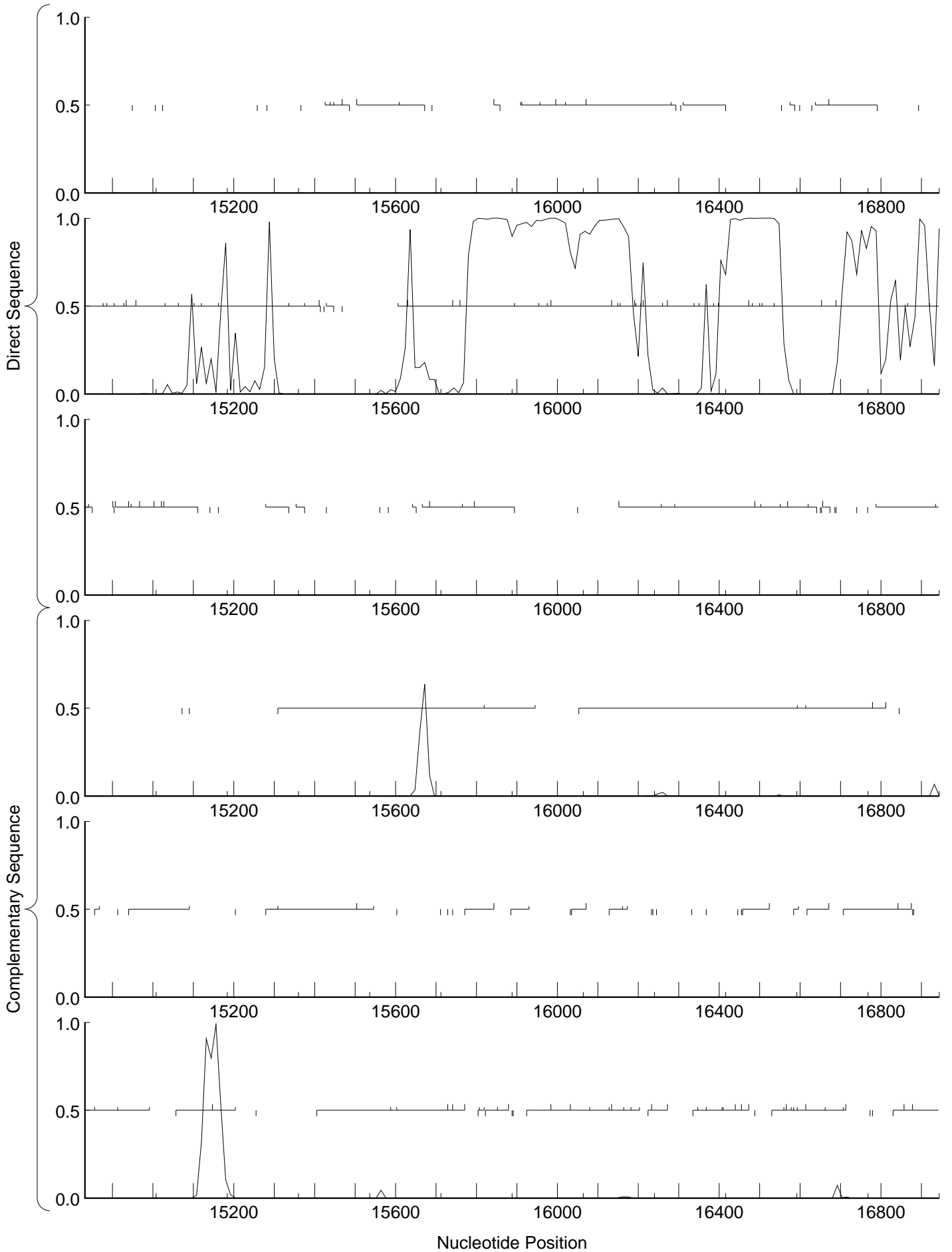


a complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 7/28

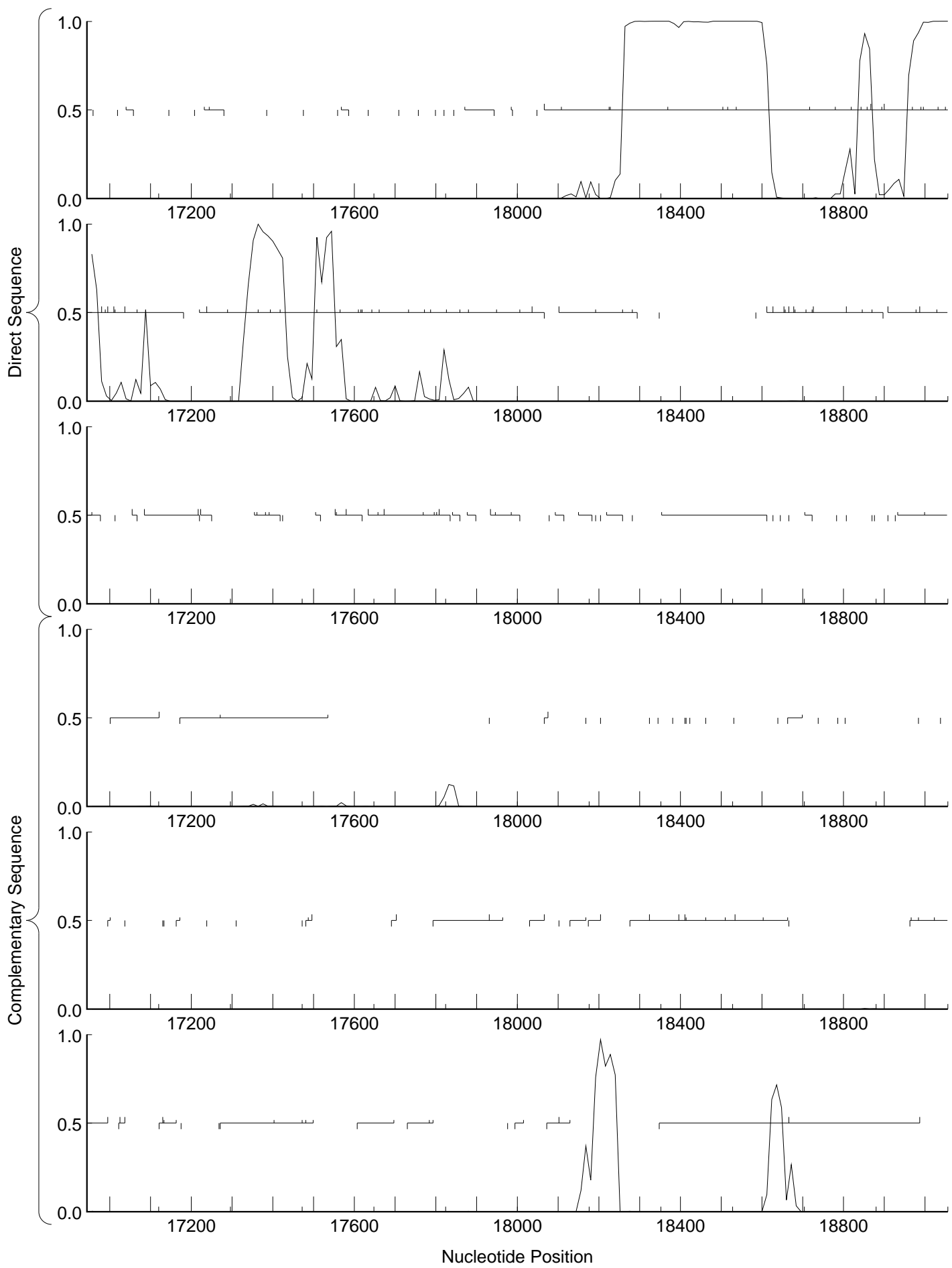




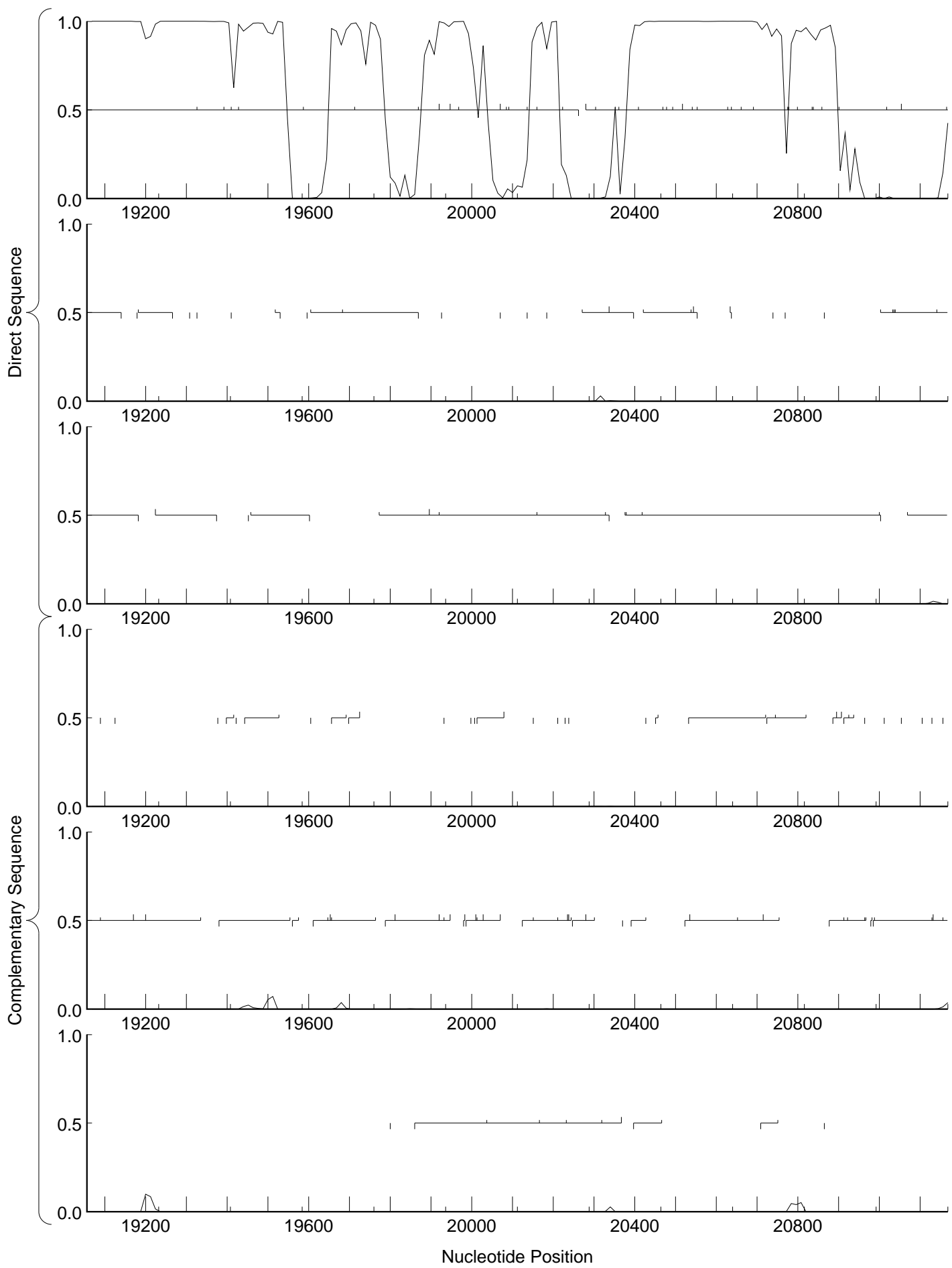


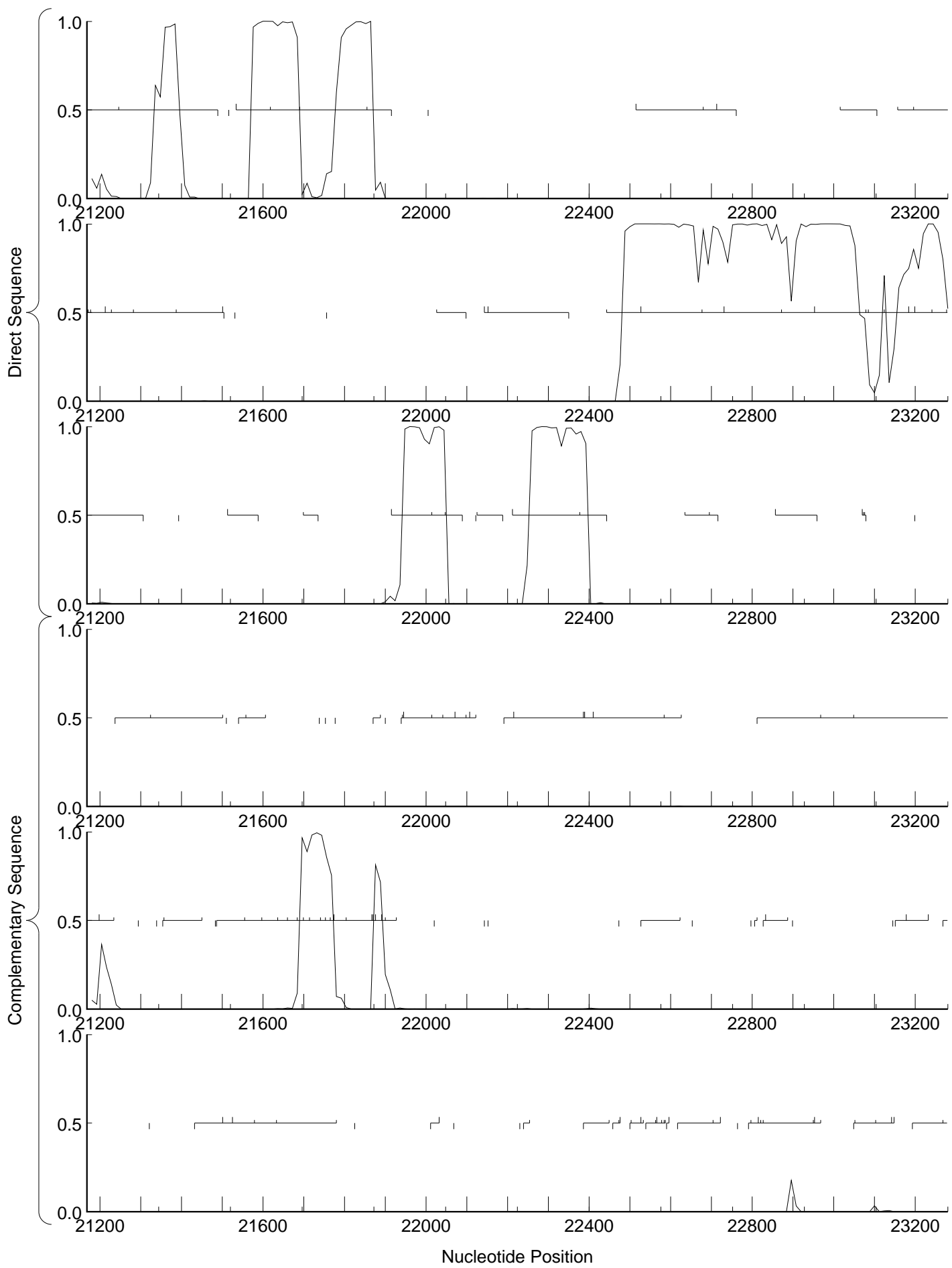


complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 10/28

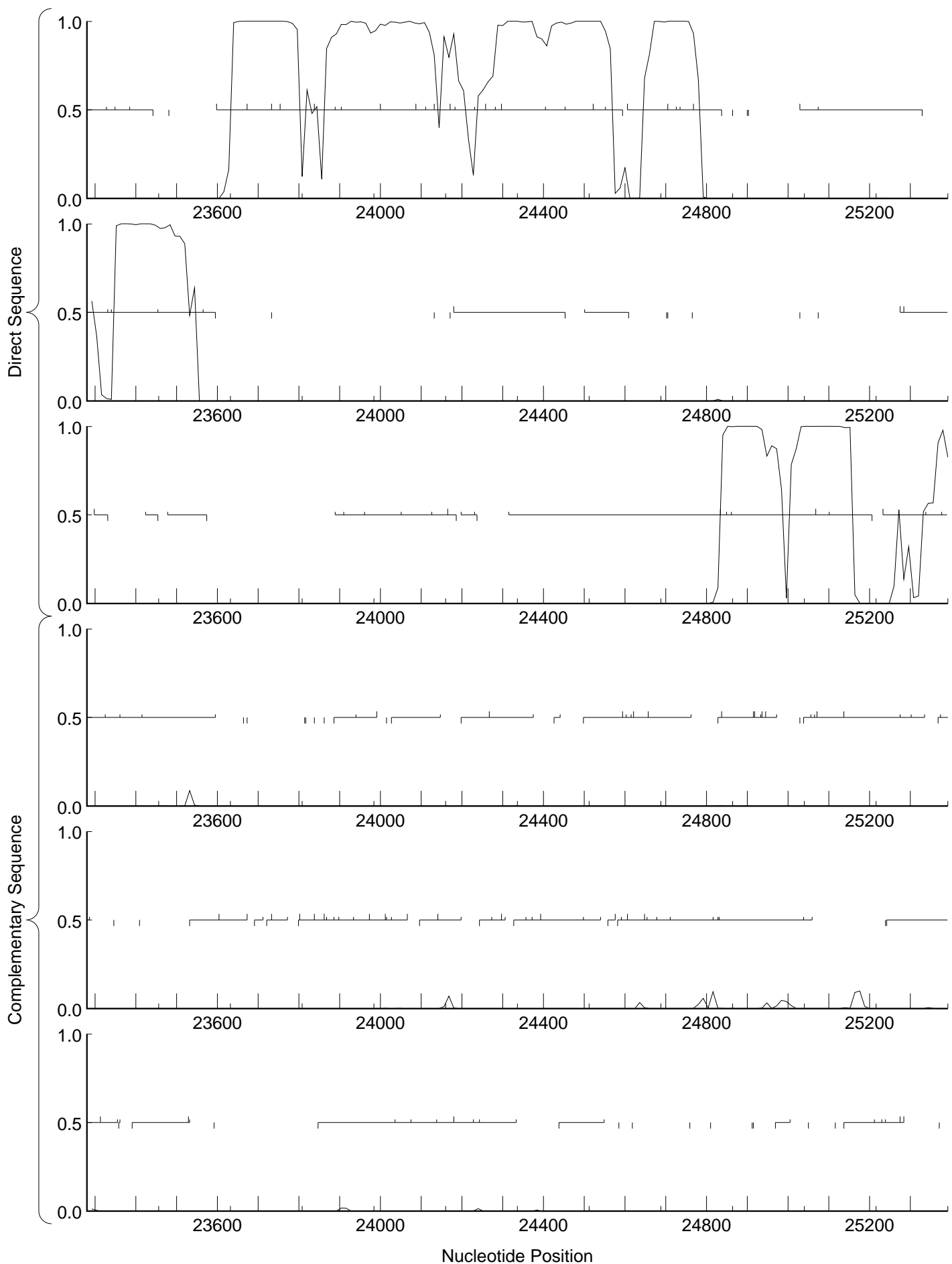


complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 11/28

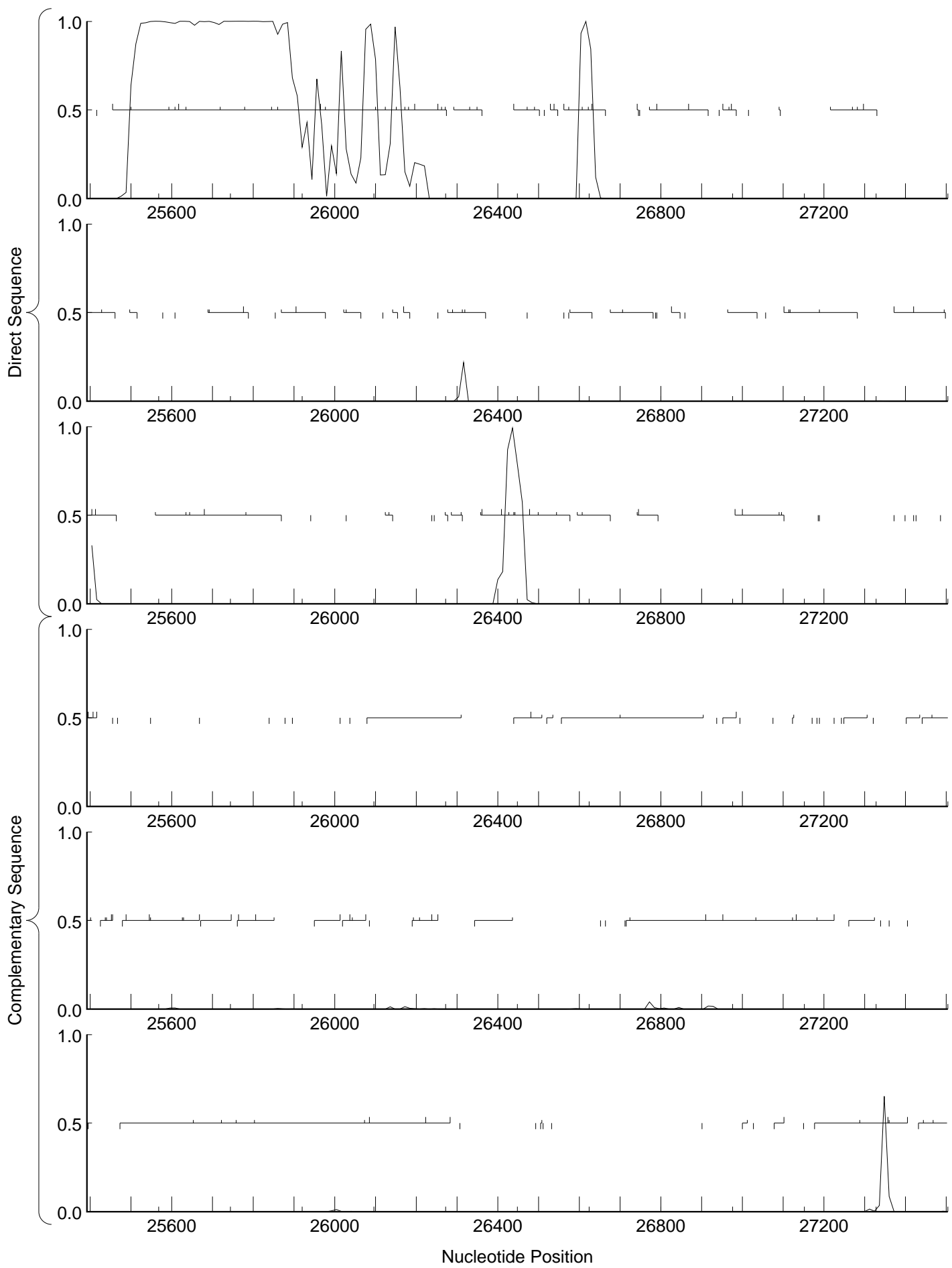




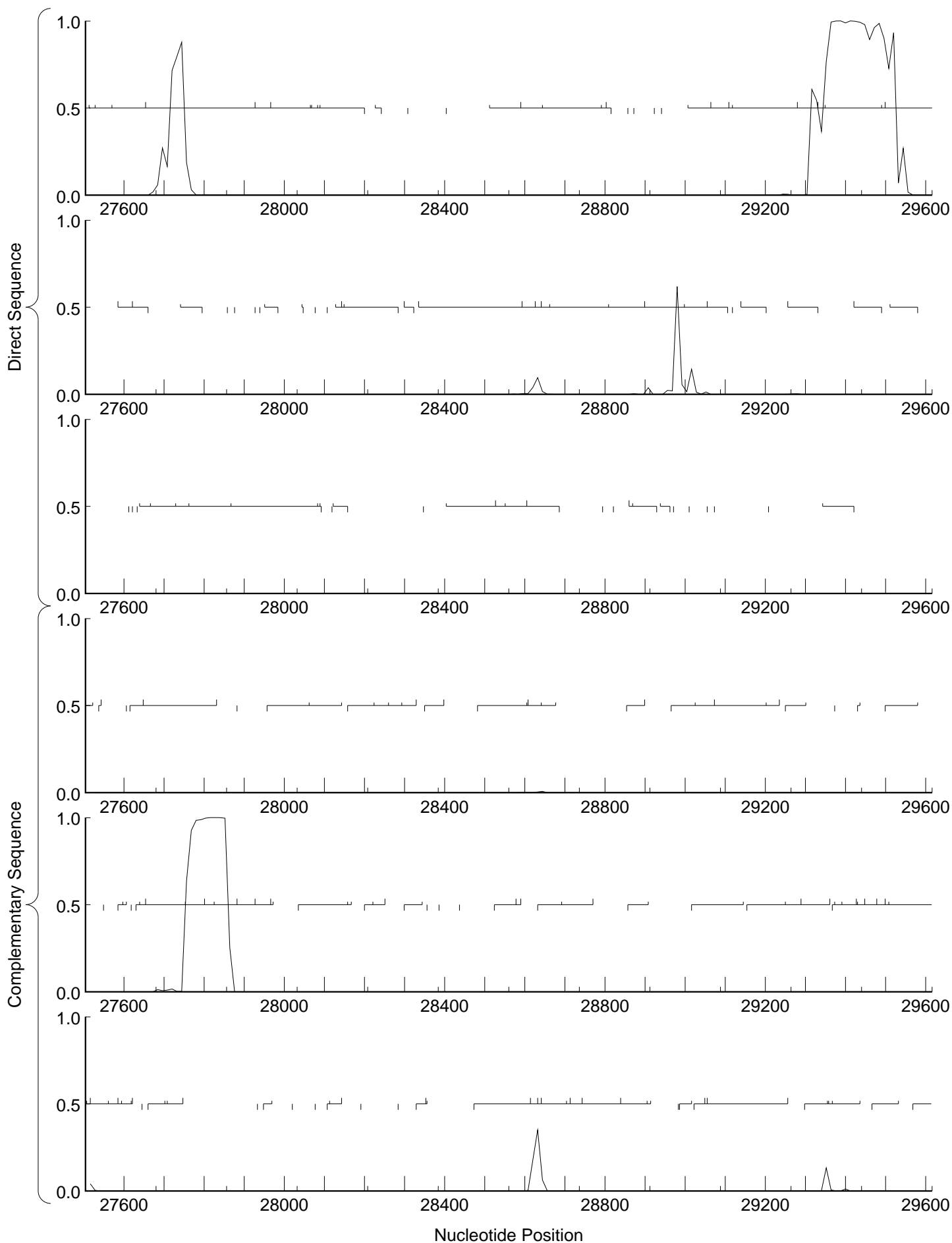
complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 13/28

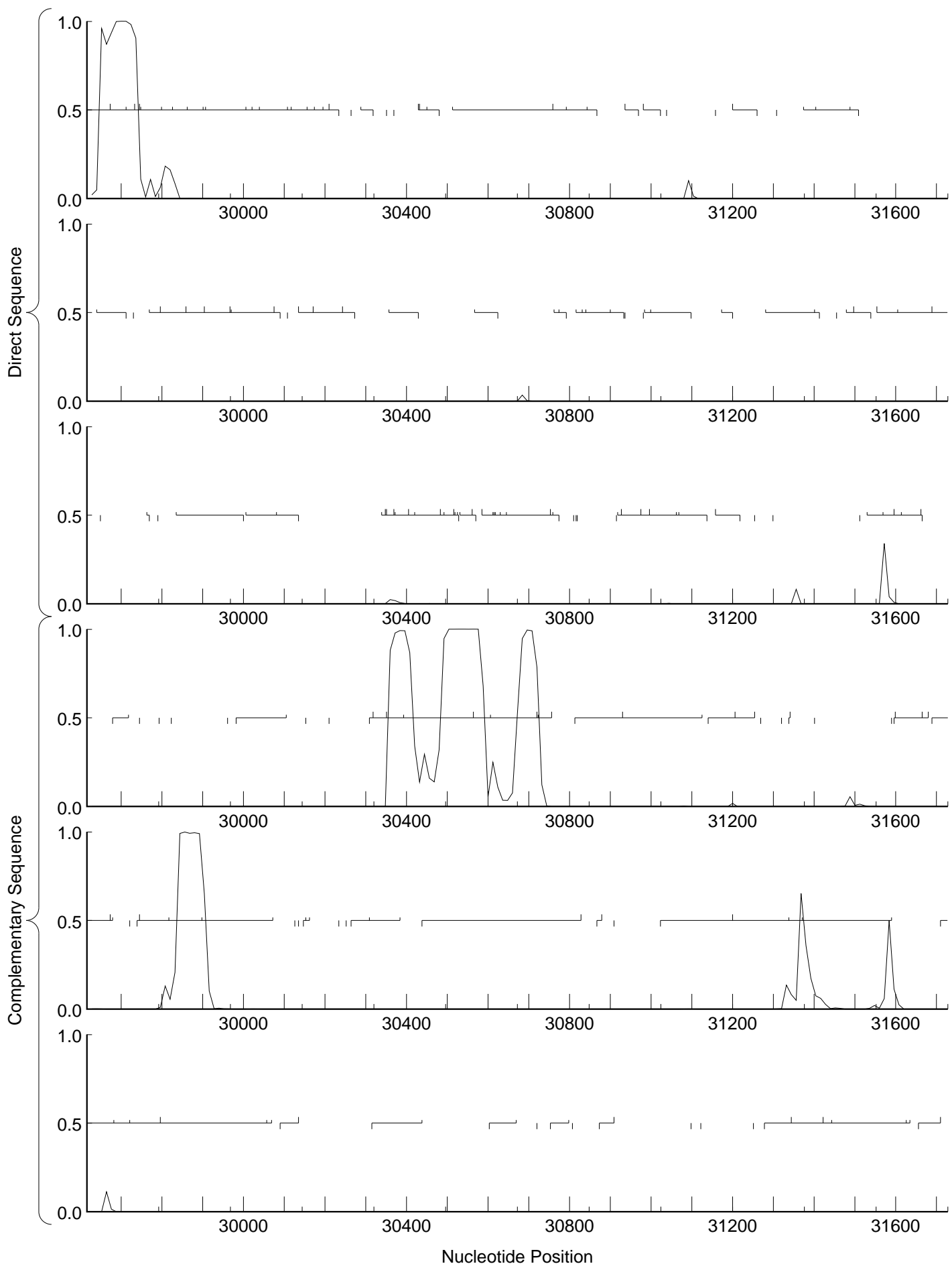


complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 14/28



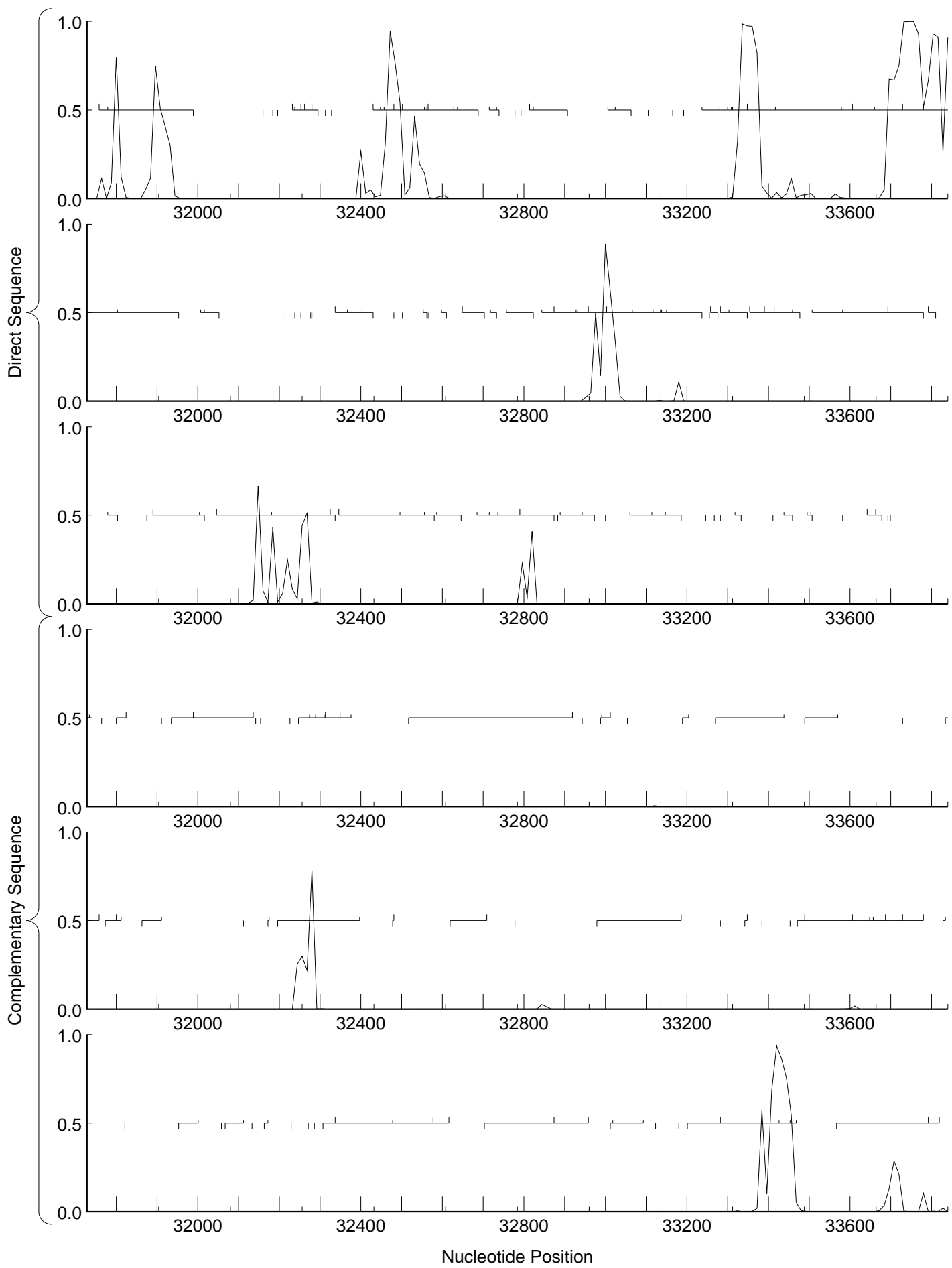
complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 15/28

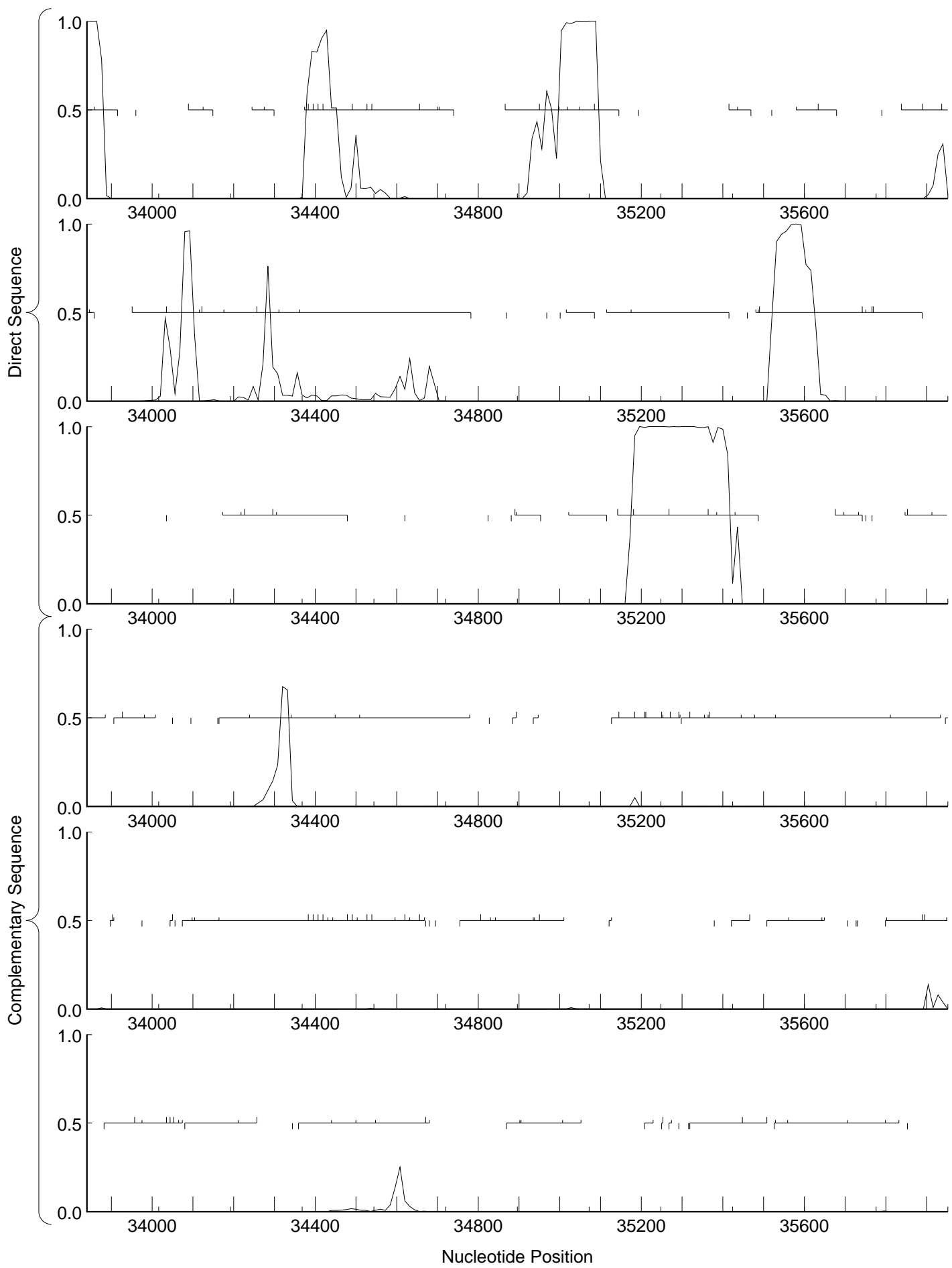






complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 17/28





complete sequence, 56251 bp including 10-base 3' overhang (CGGAAGGCGC), Cluster F1, Order 4, Window 96, Step 12, 19/28

