

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Gordonia phage Eyes complete sequence, 46772 bp including 10 b' 3' overhang (CGCCGCGGTA), C  
Analysis Date : 6/28/17 at 22:13:15  
Pages : 24  
Sequence Length : 46772 bp  
GC Content : 67.09%

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,

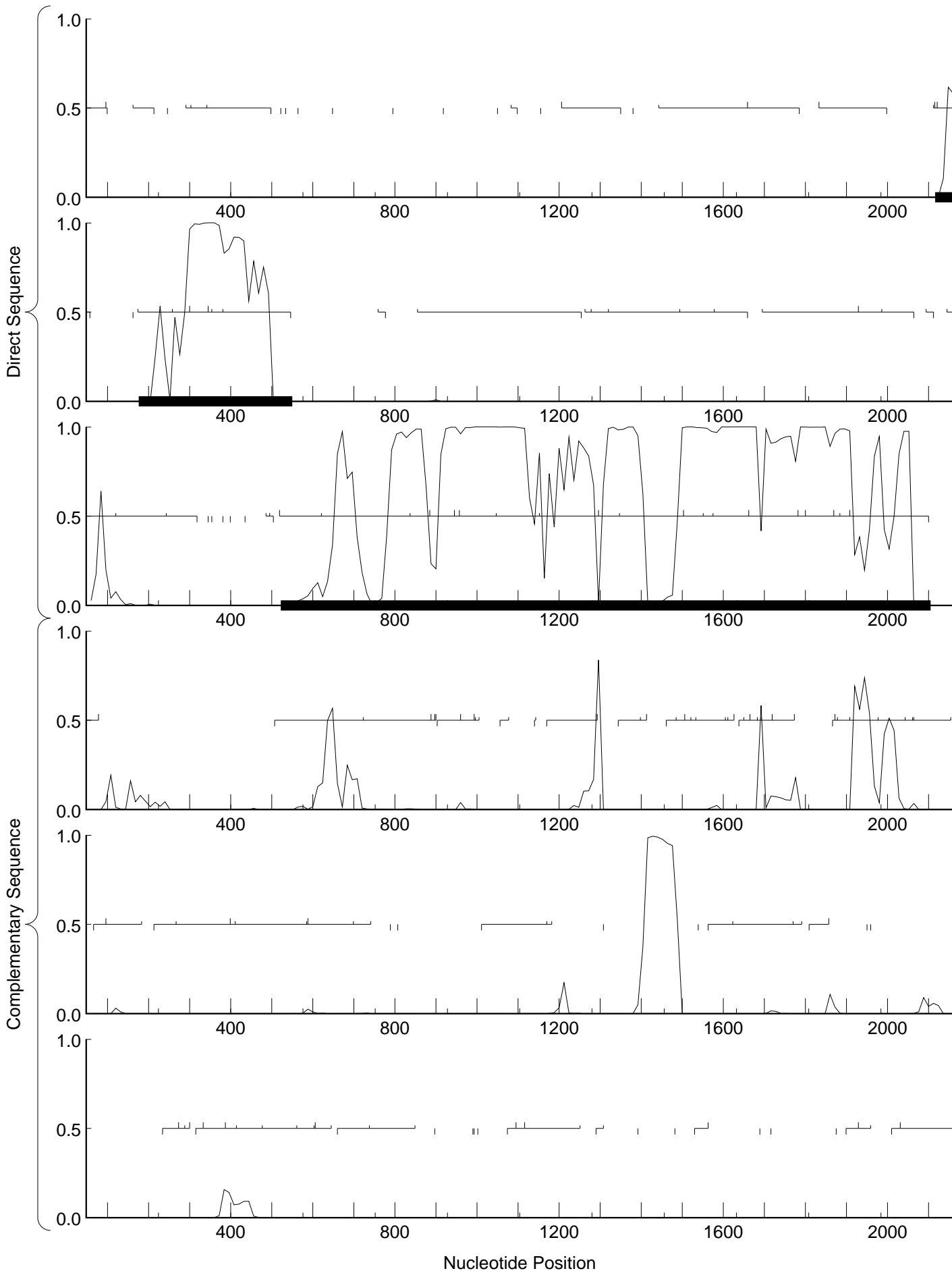
## Matrix notes & comments

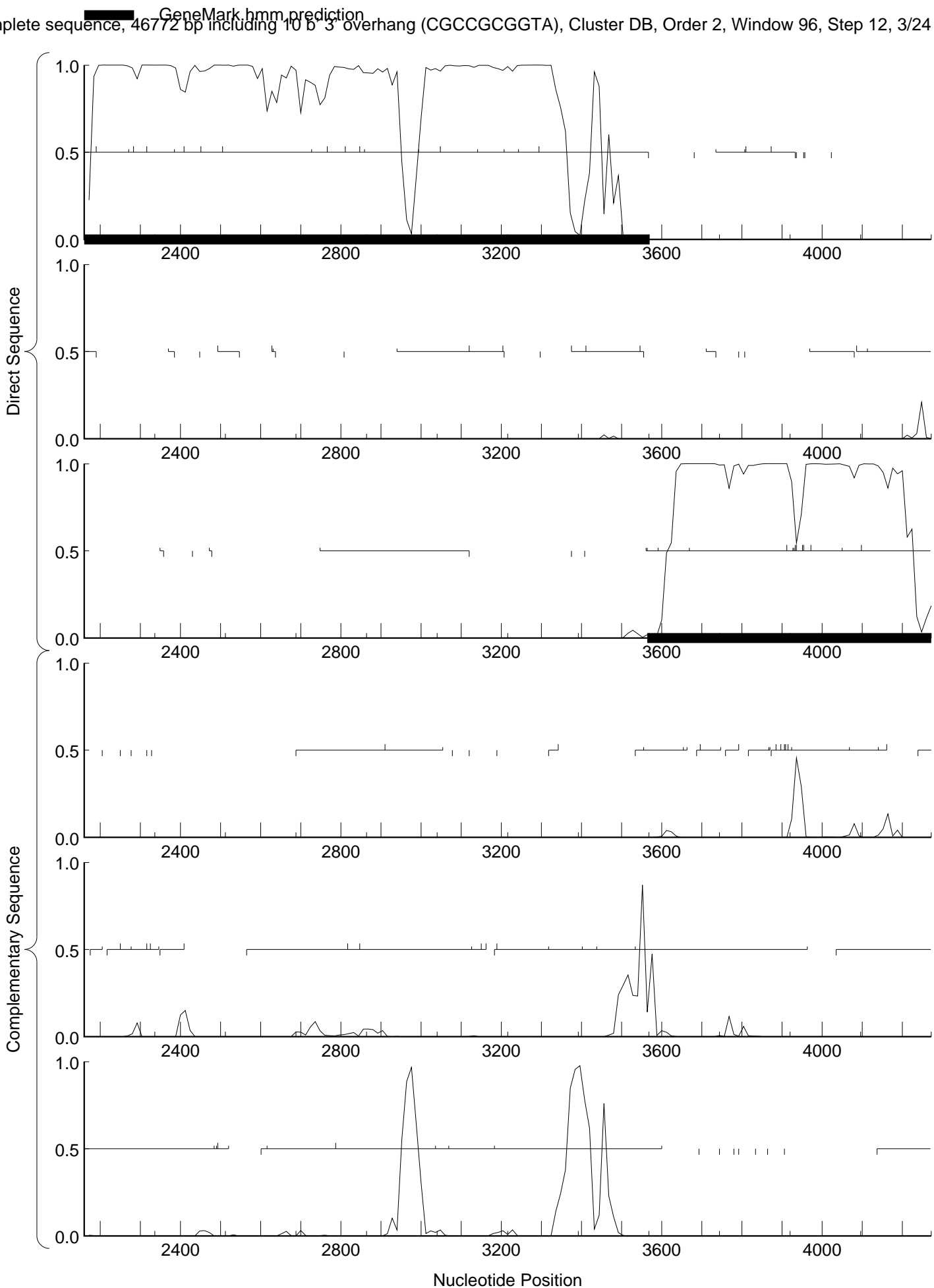
For details on the model building procedure see:  
Besemer J. and Borodovsky M.  
"Heuristic approach to deriving models for gene finding"  
NAR, 1999, Vol. 27, No. 19, pp. 3911-3920

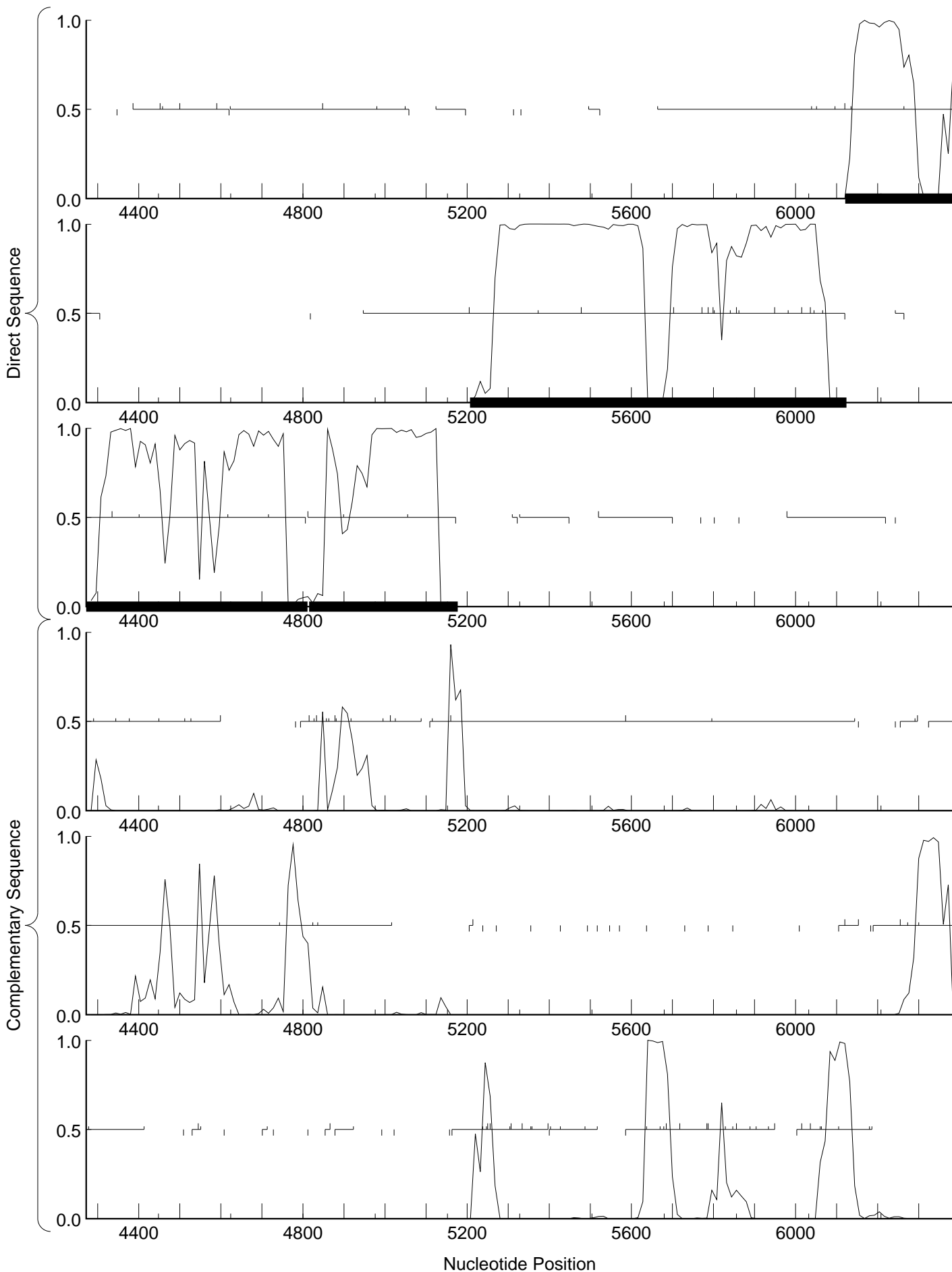
## MATRIX INFORMATION

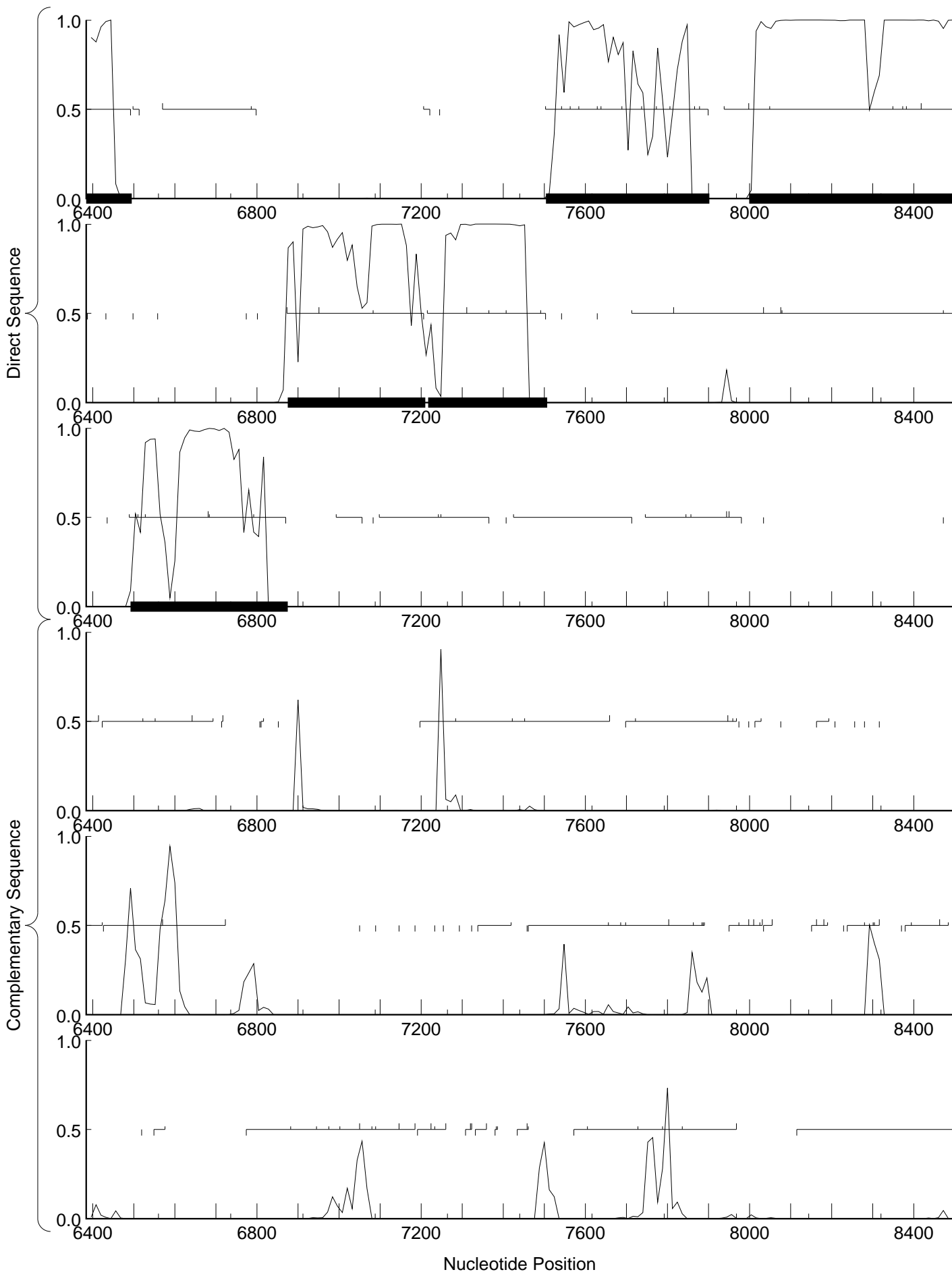
Matrix : Heuristic, GC = 67  
Author : Borodovsky Laboratory - Georgia Tech  
Order : 2

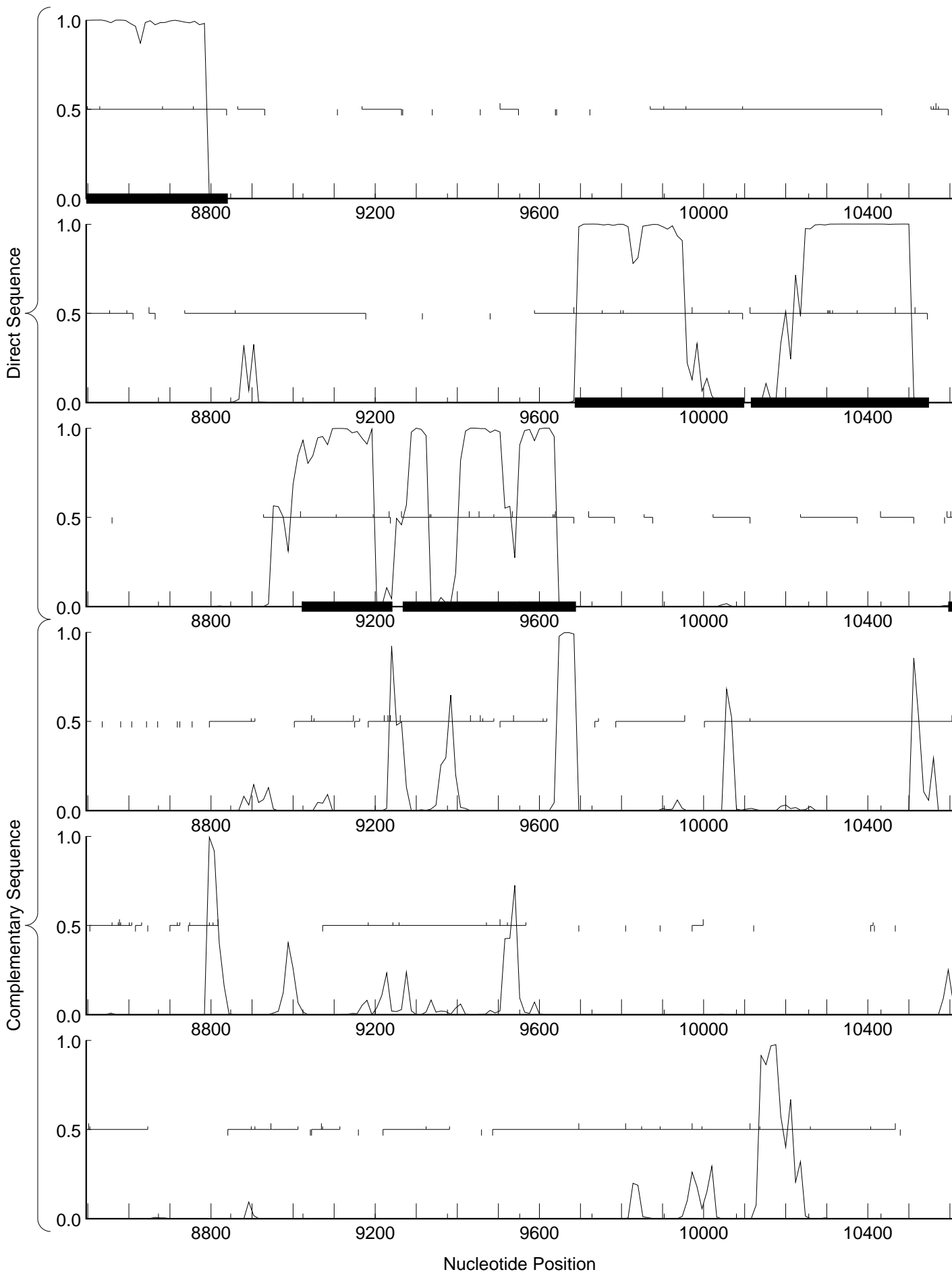
Send questions / comments to:  
Dr. M. Borodovsky  
Georgia Institute of Technology  
School of Biology  
Atlanta, GA 30332-0230

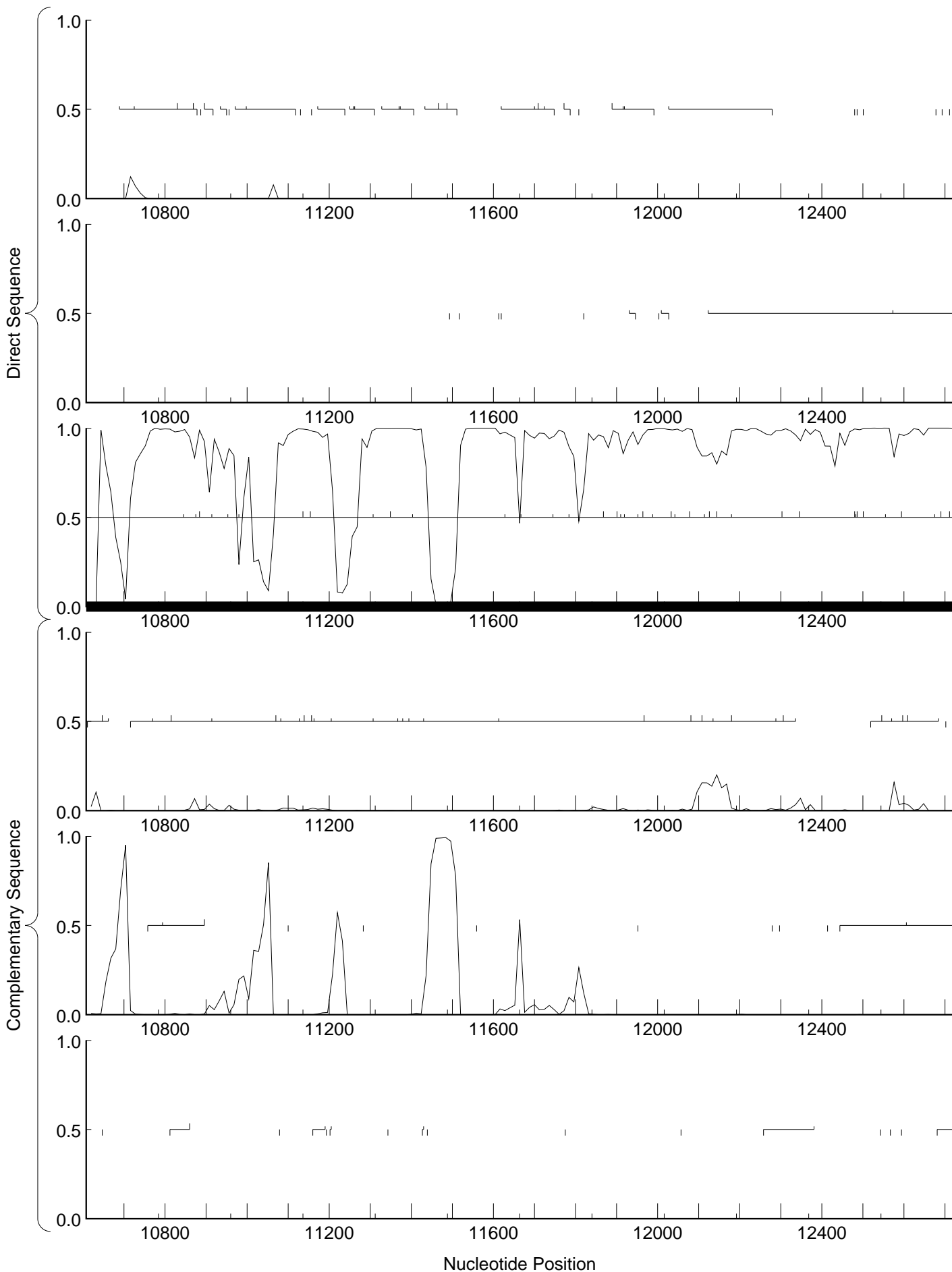


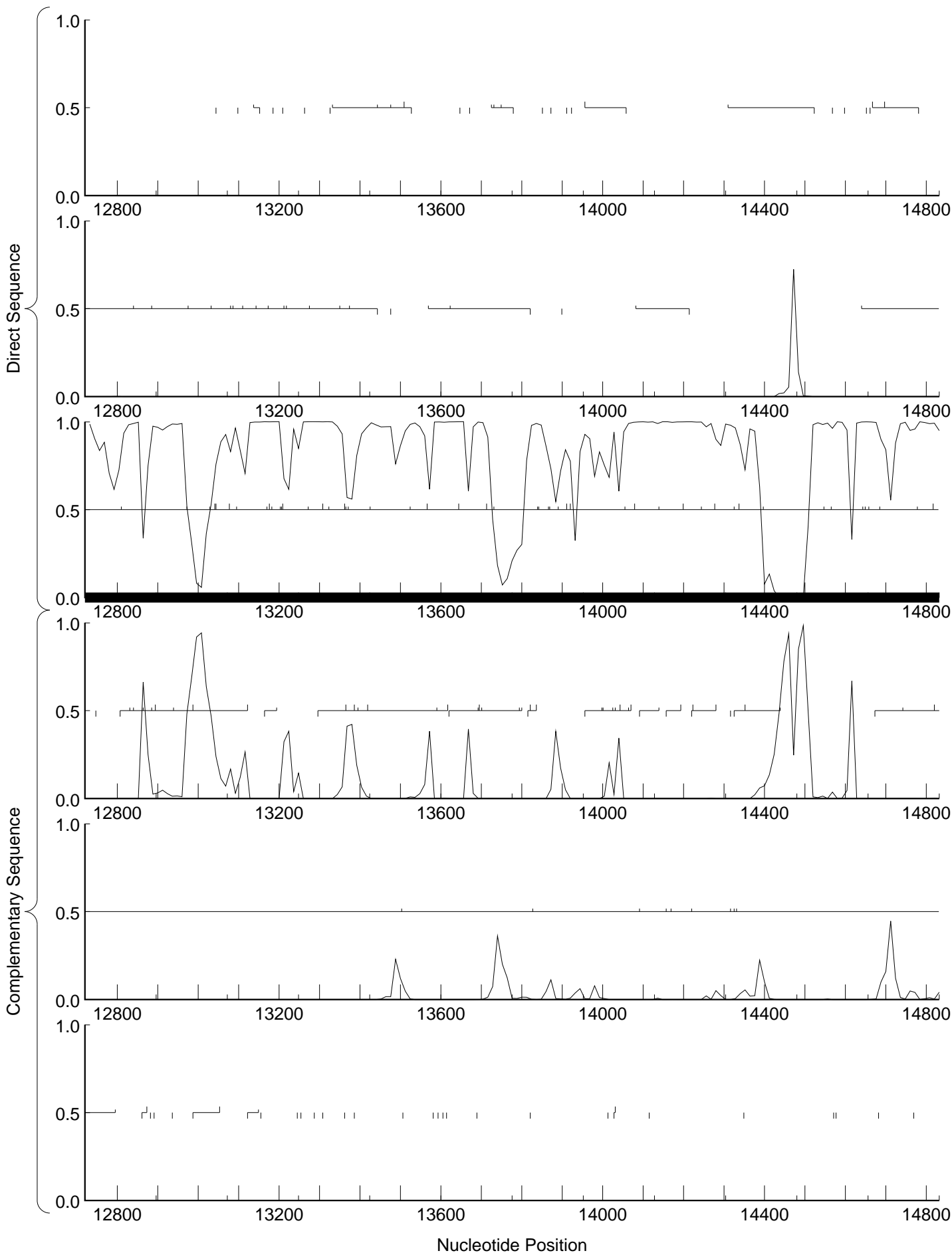






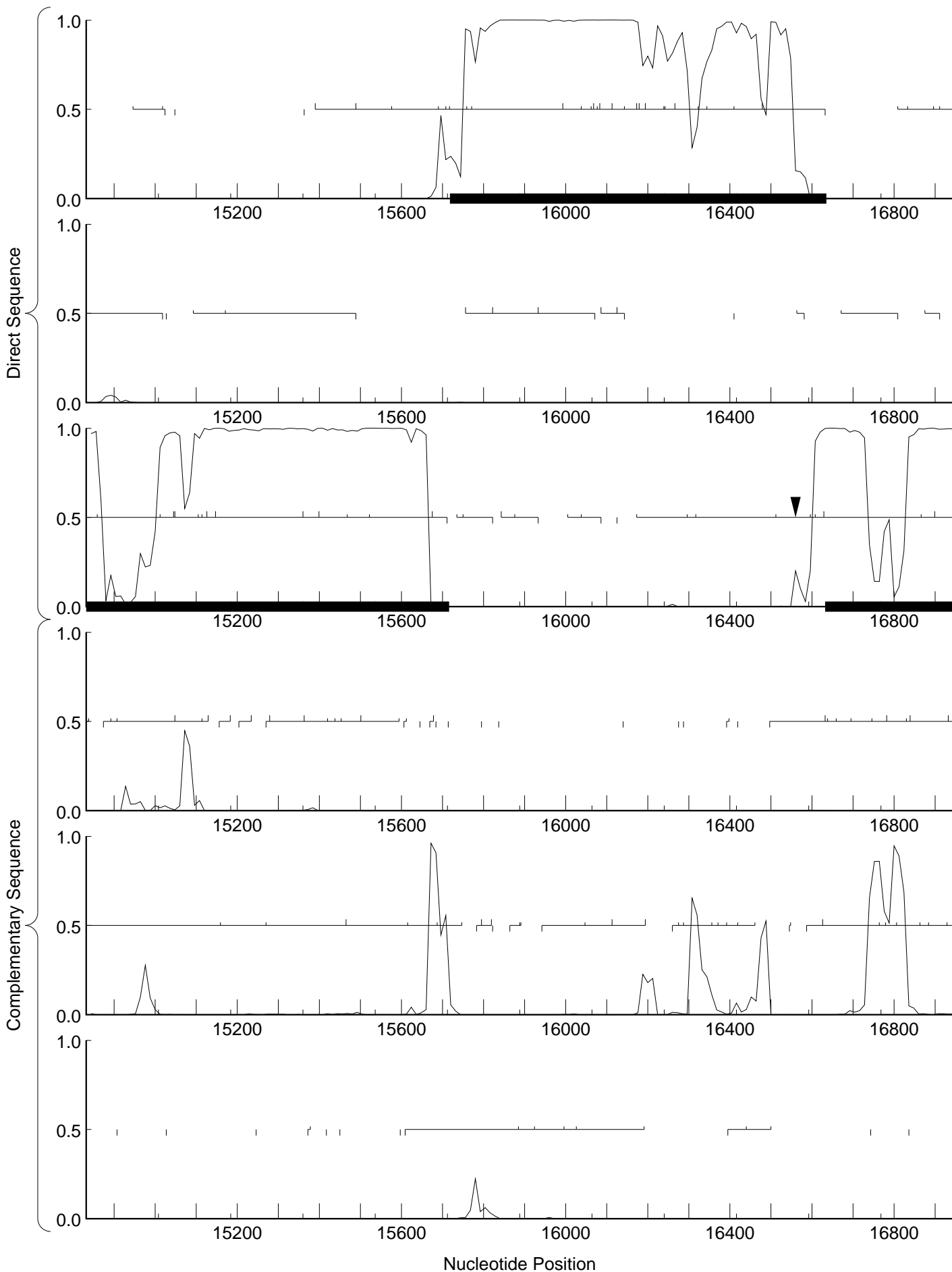


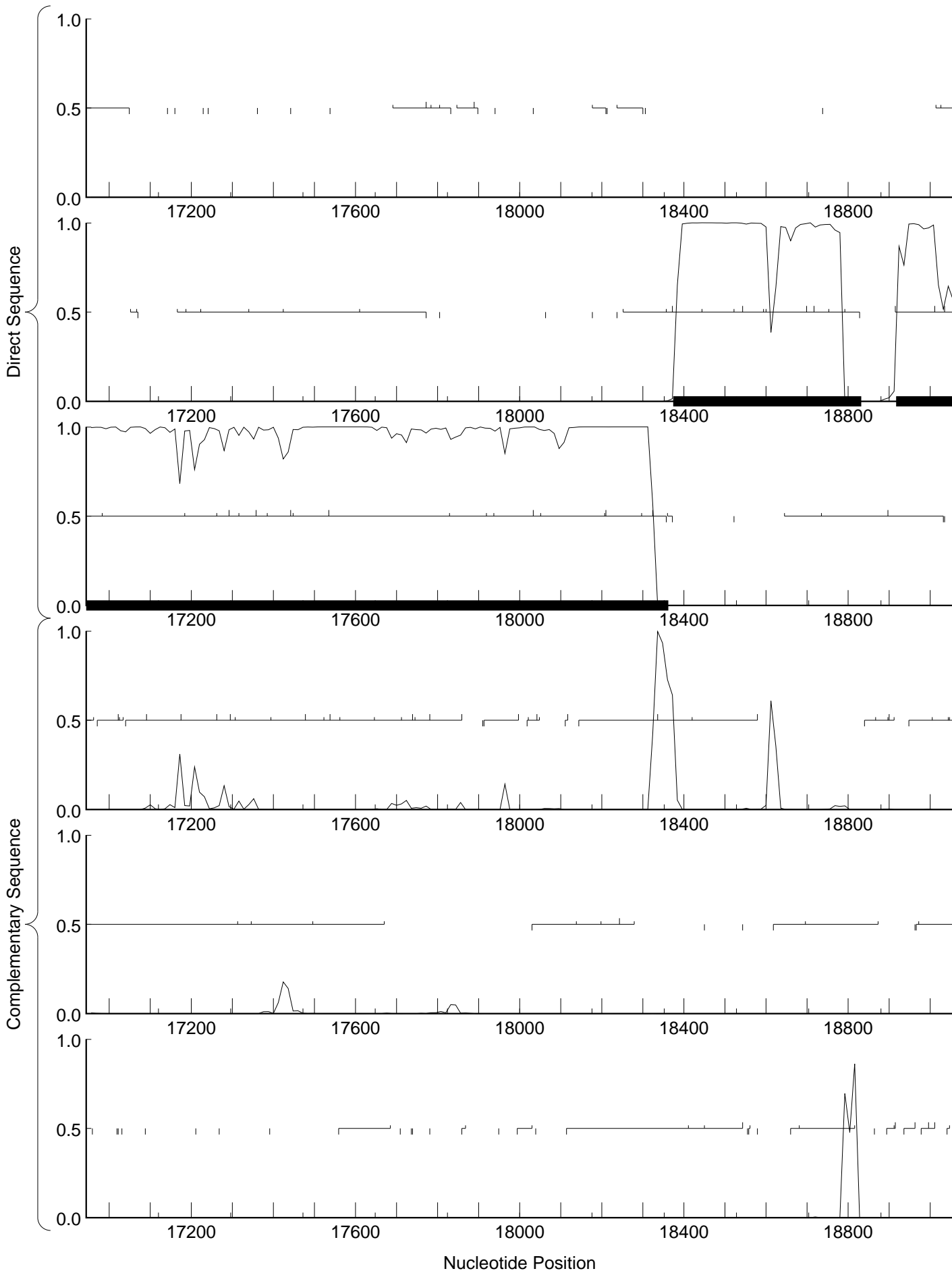


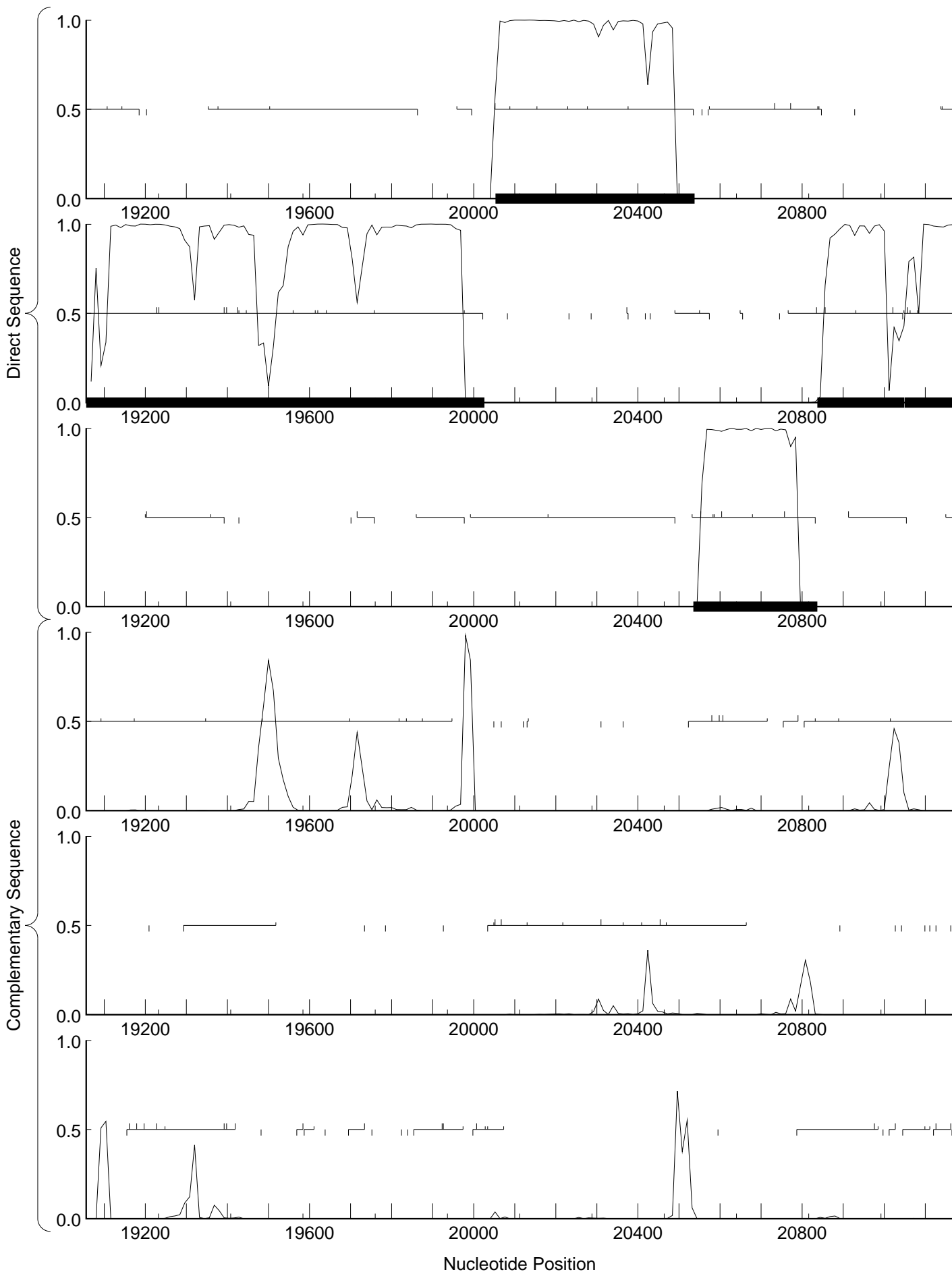


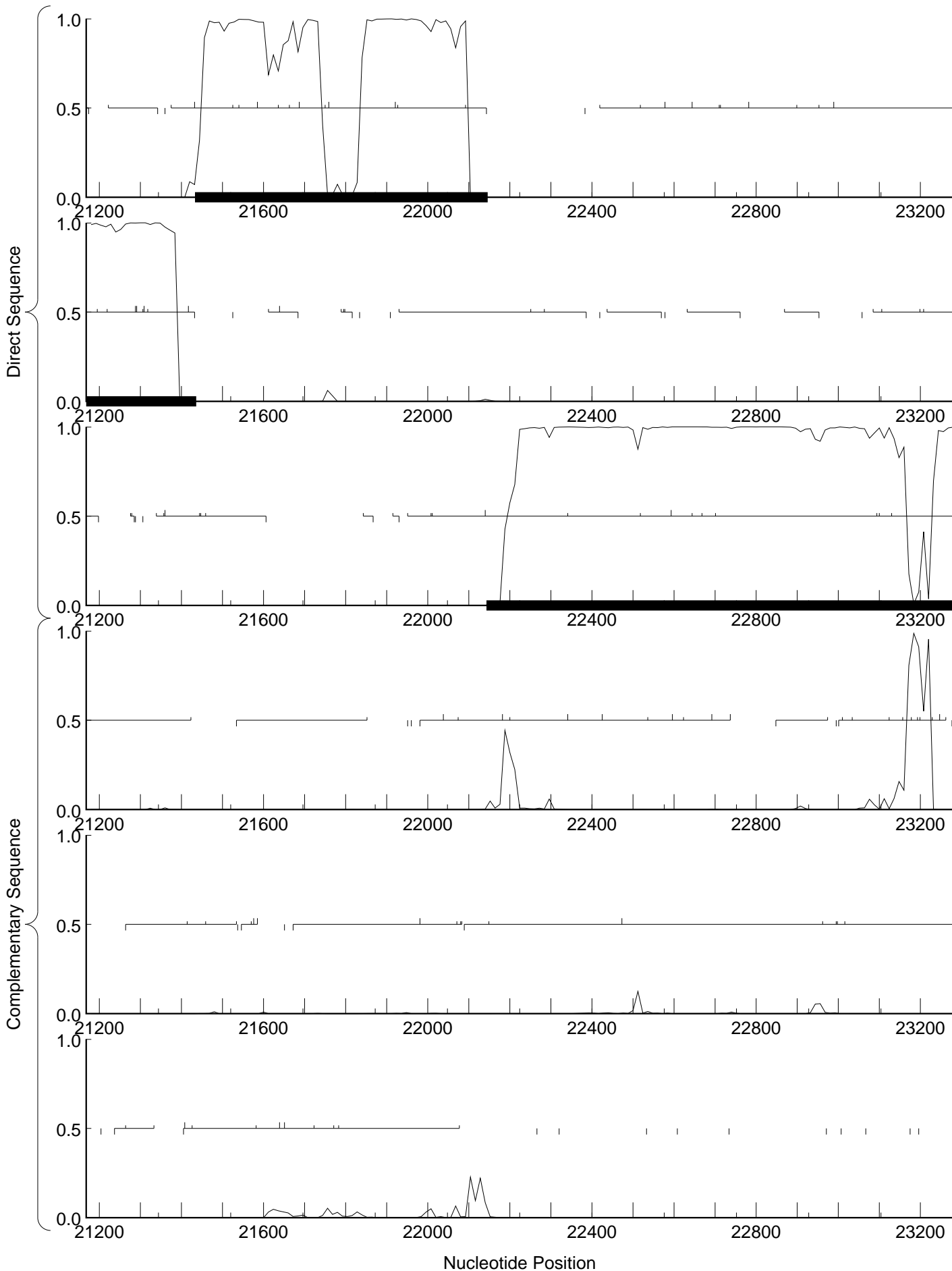


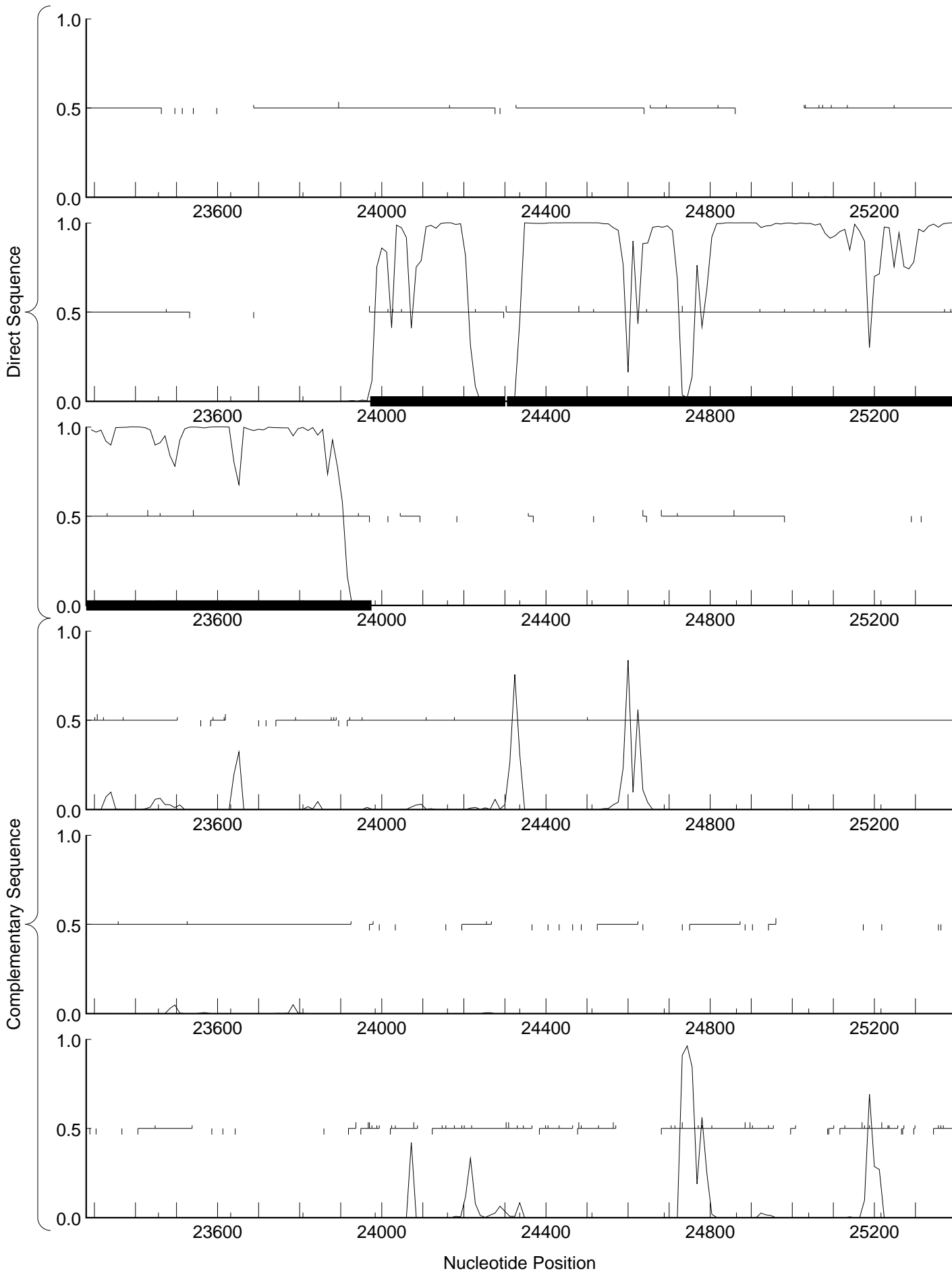
GeneMark.hmm.prediction

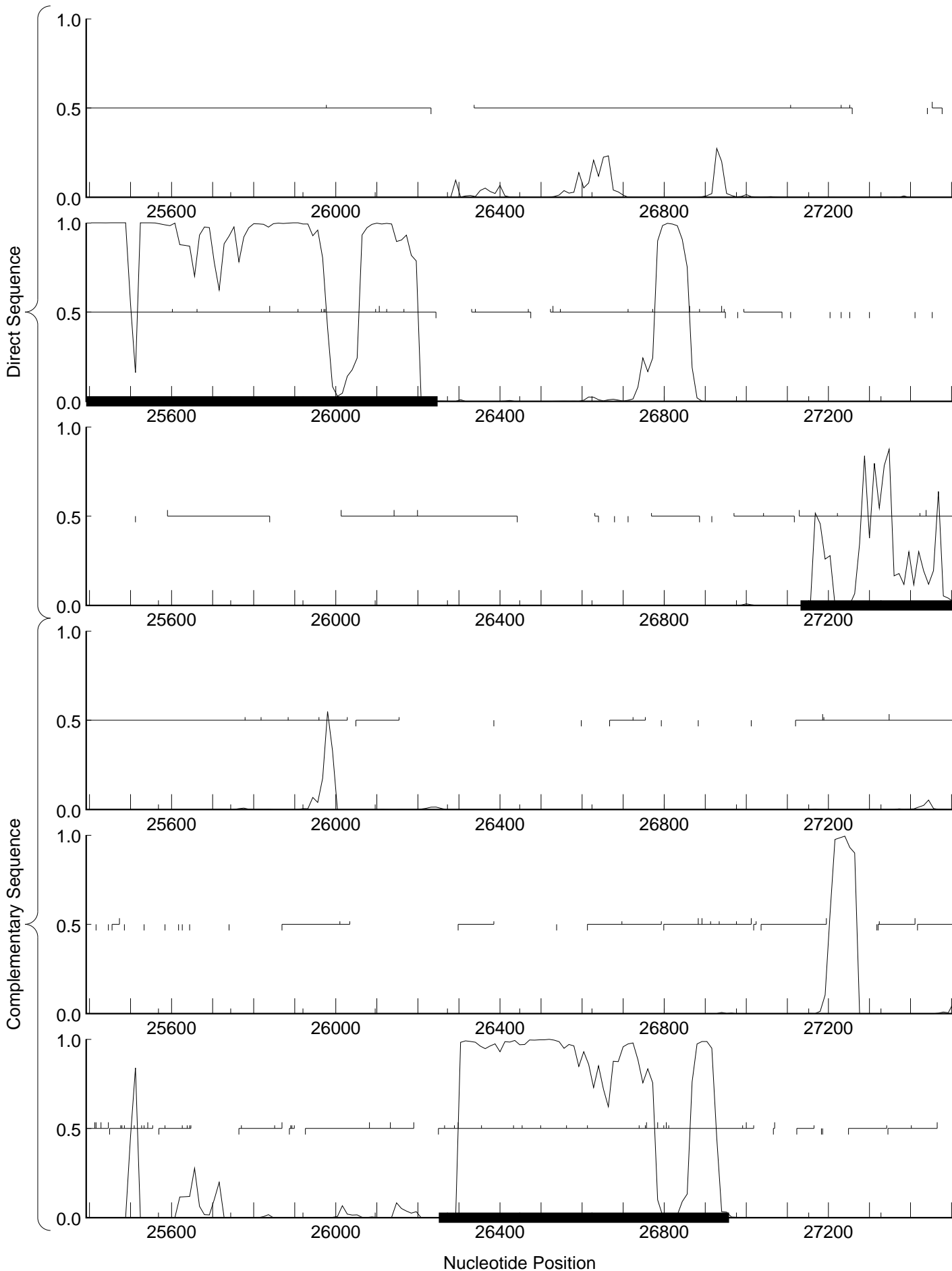


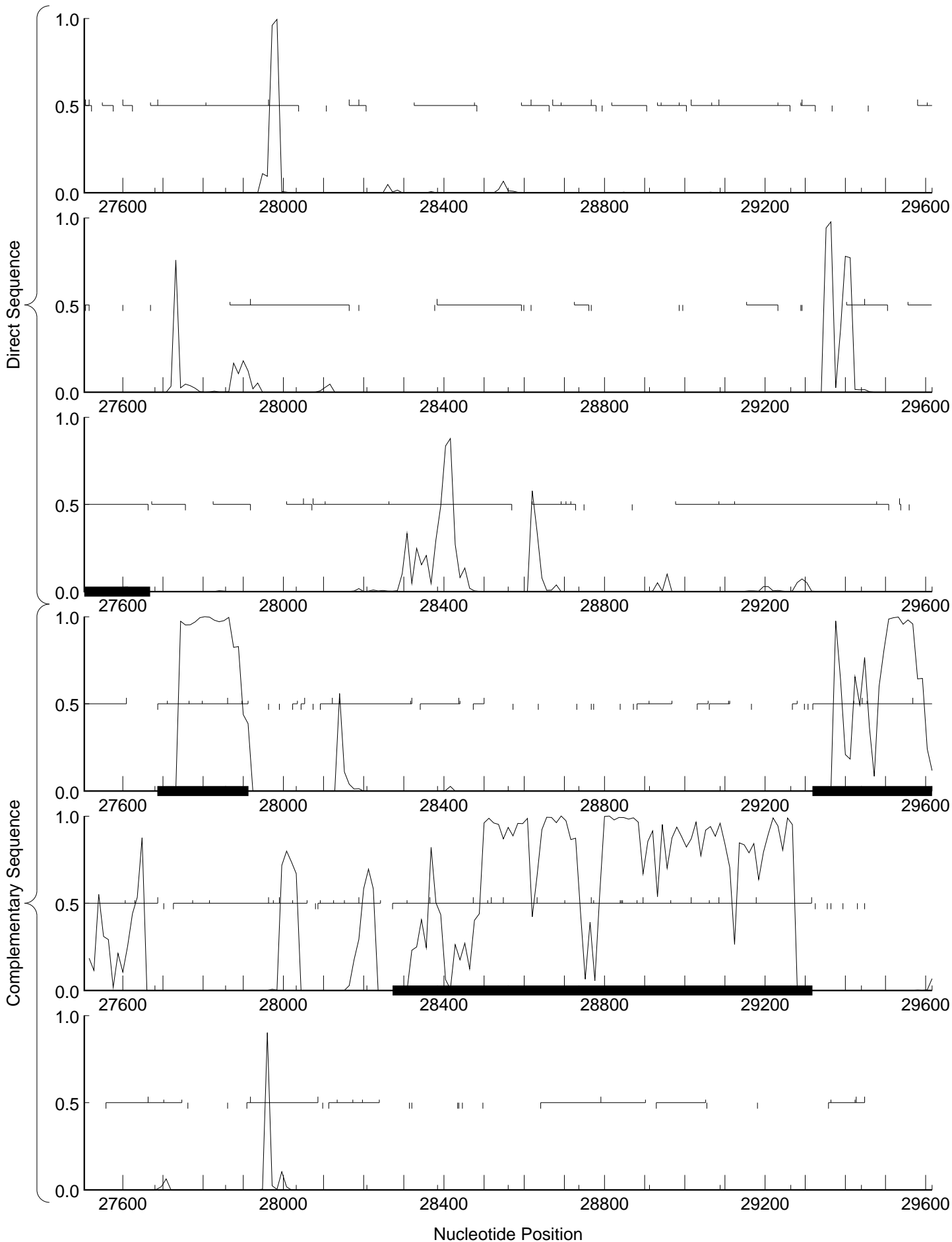


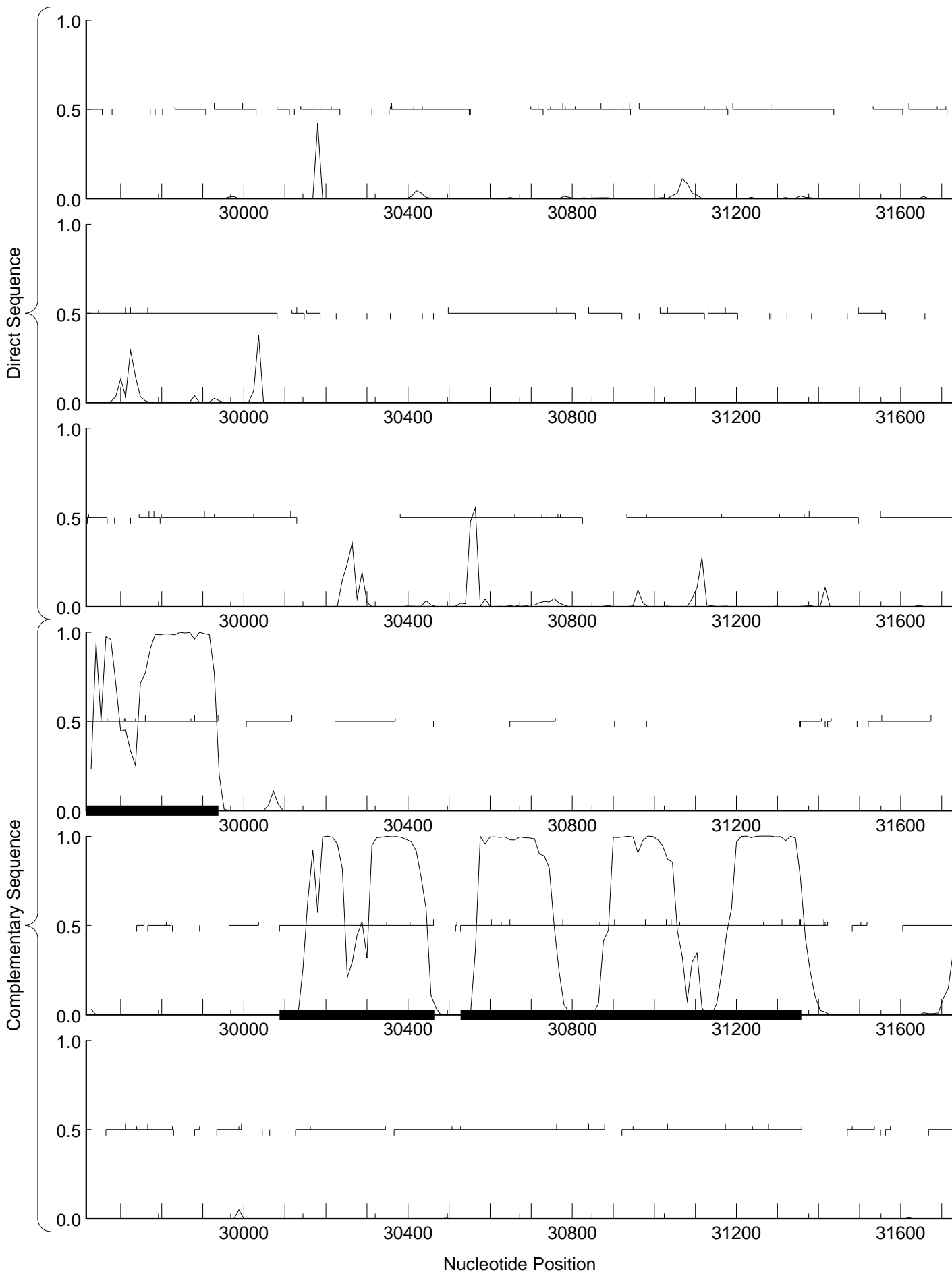




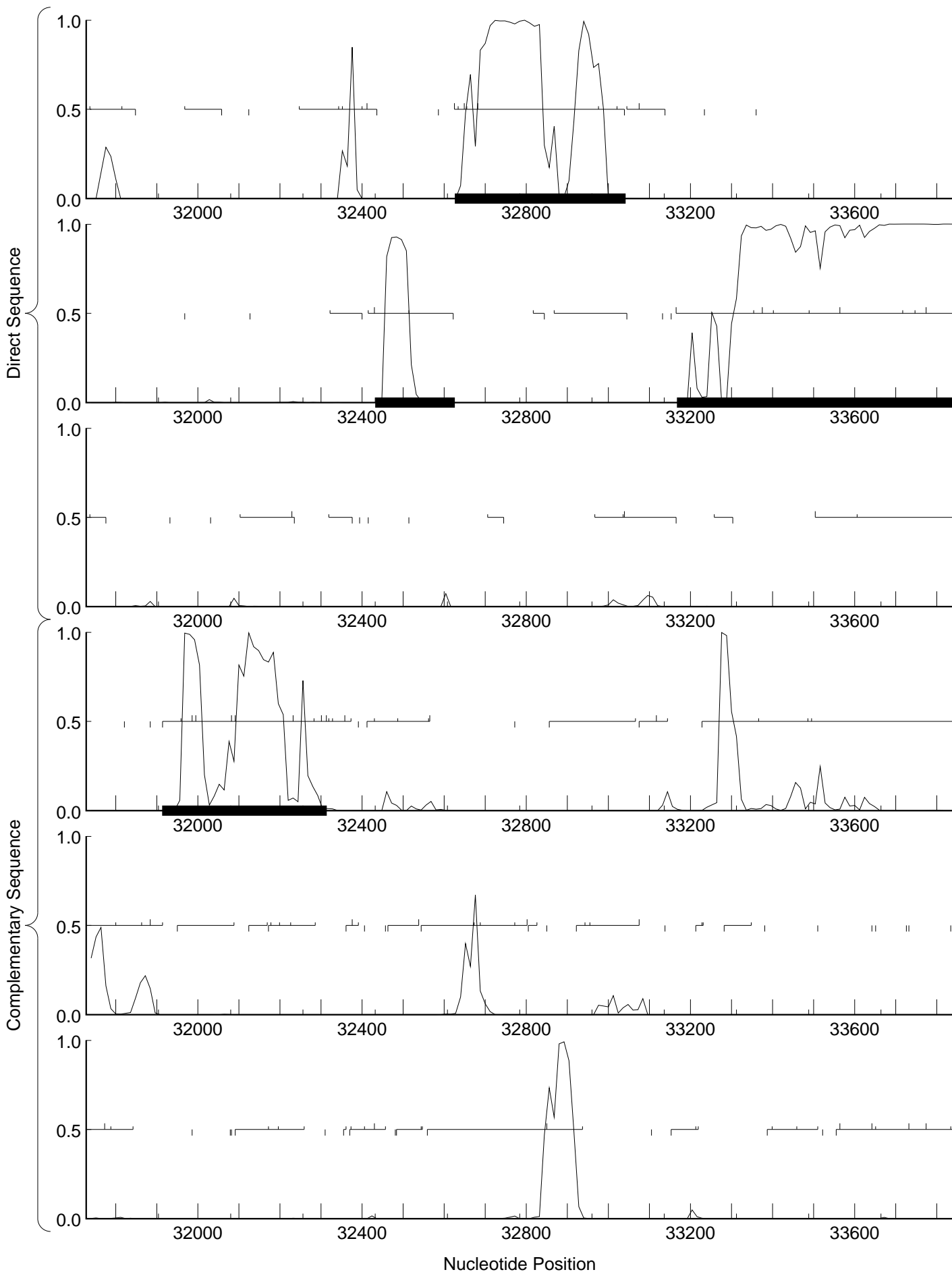


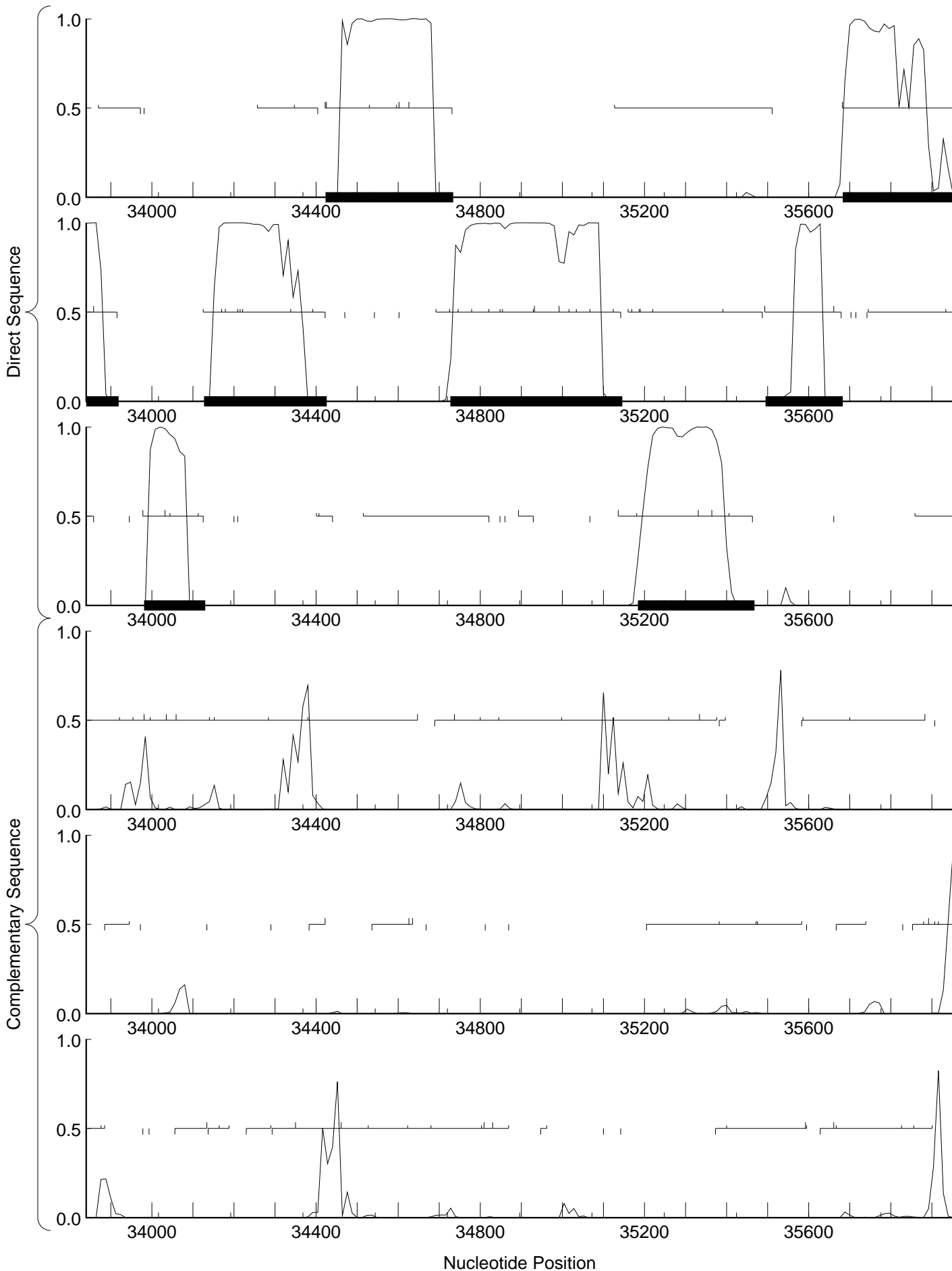


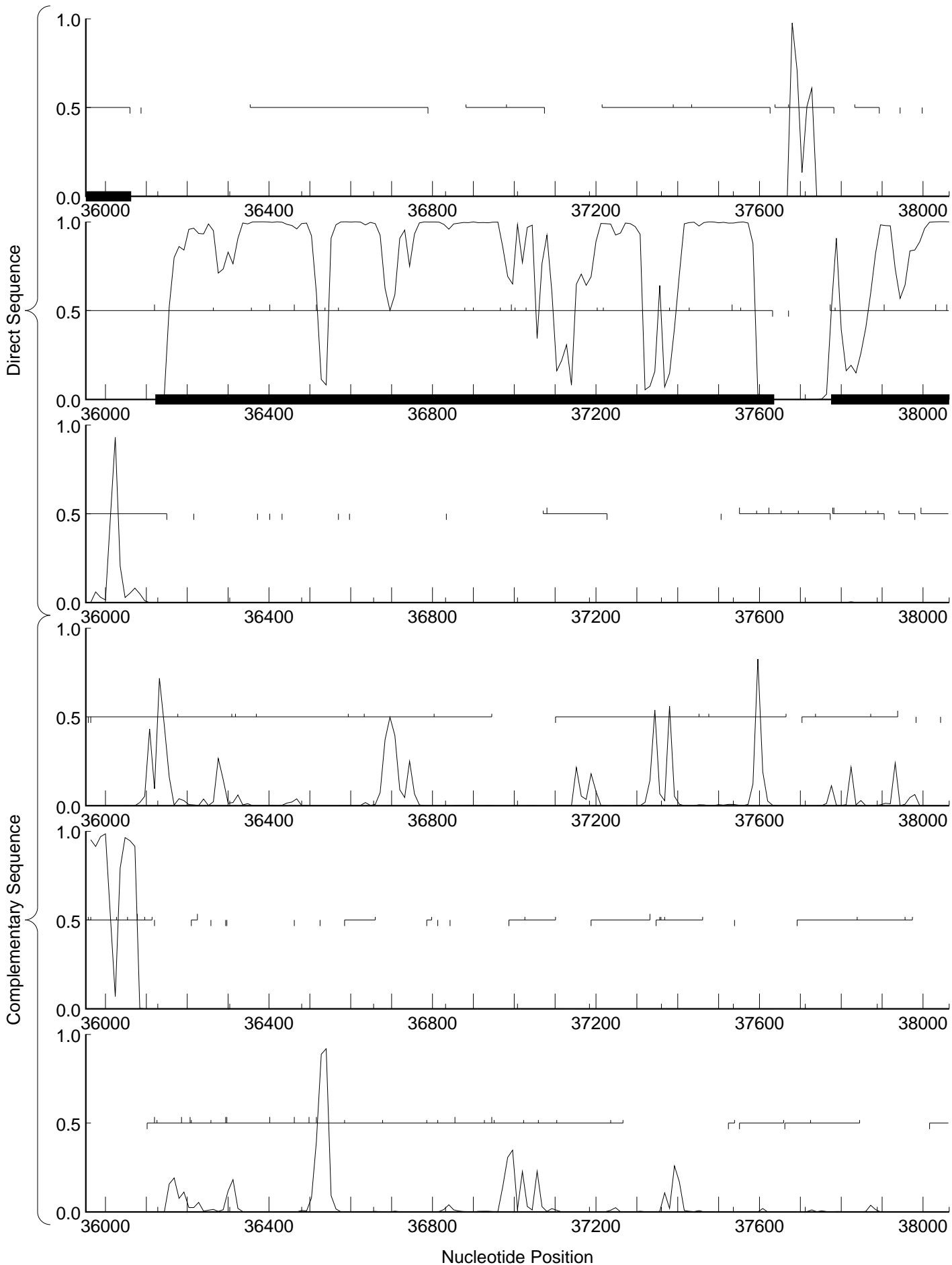


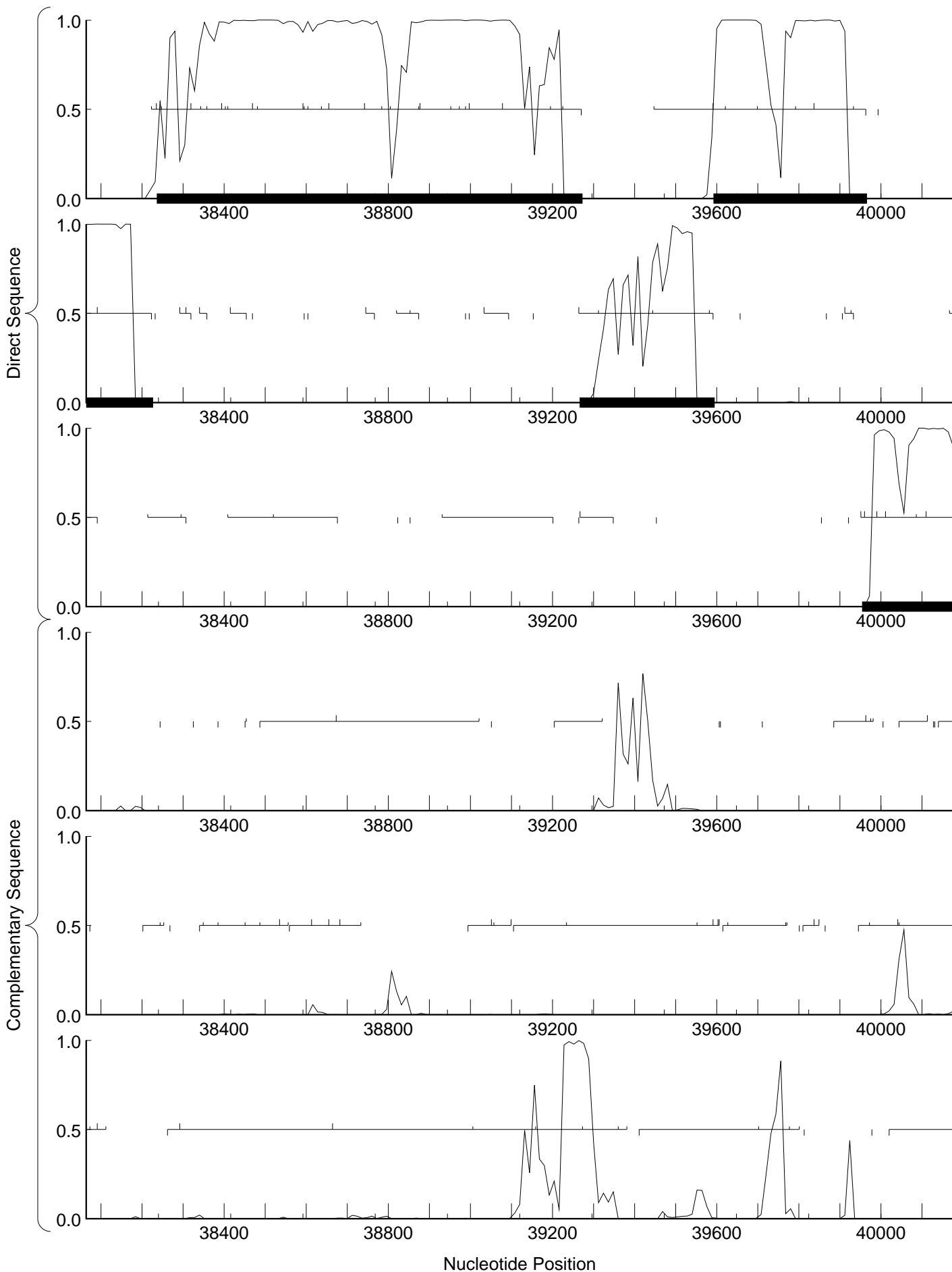


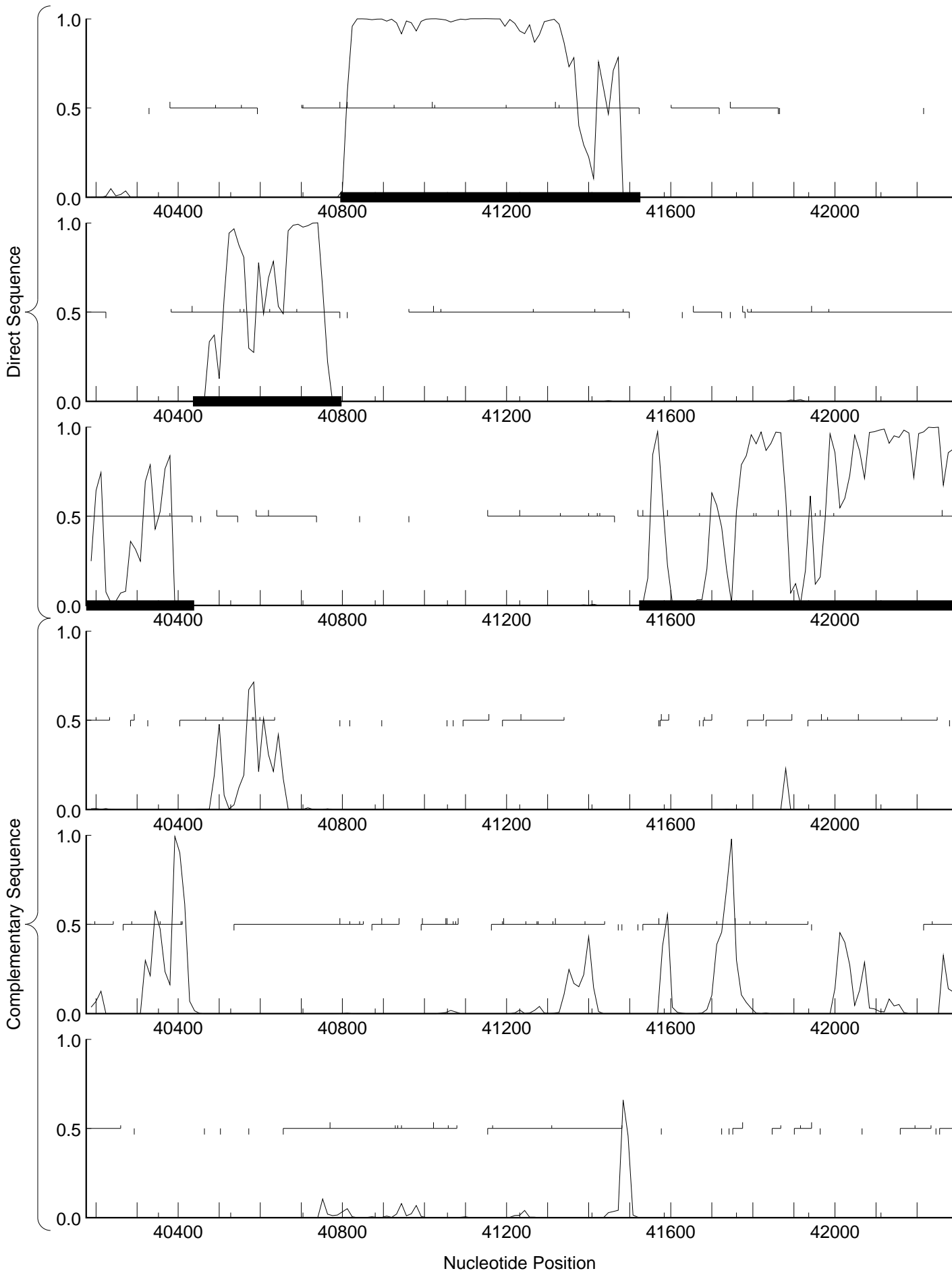


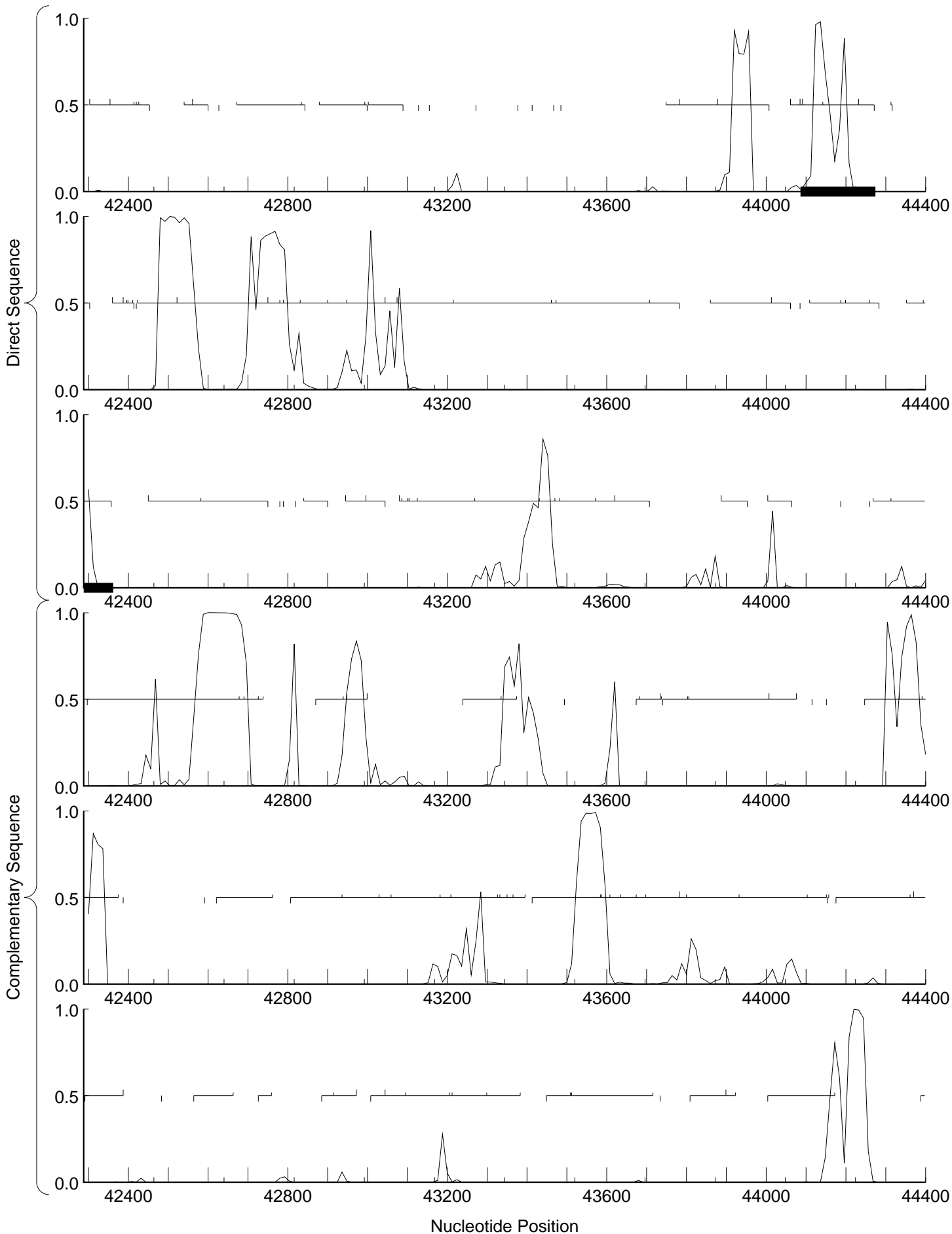












Genes complete sequence, 46772 bp including 10 b 3' overhang (CGCCGCGGTA), Cluster DB, Order 2, Window 96, Step 12, 23/24

