

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Mycobacterium phage BadStone complete sequence, 75679 bp including 9-base 3' overhang (CGCT  
Analysis Date : 2/7/25 at 12:28:39  
Pages : 37  
Sequence Length : 75679 bp  
GC Content : 63.05%

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,

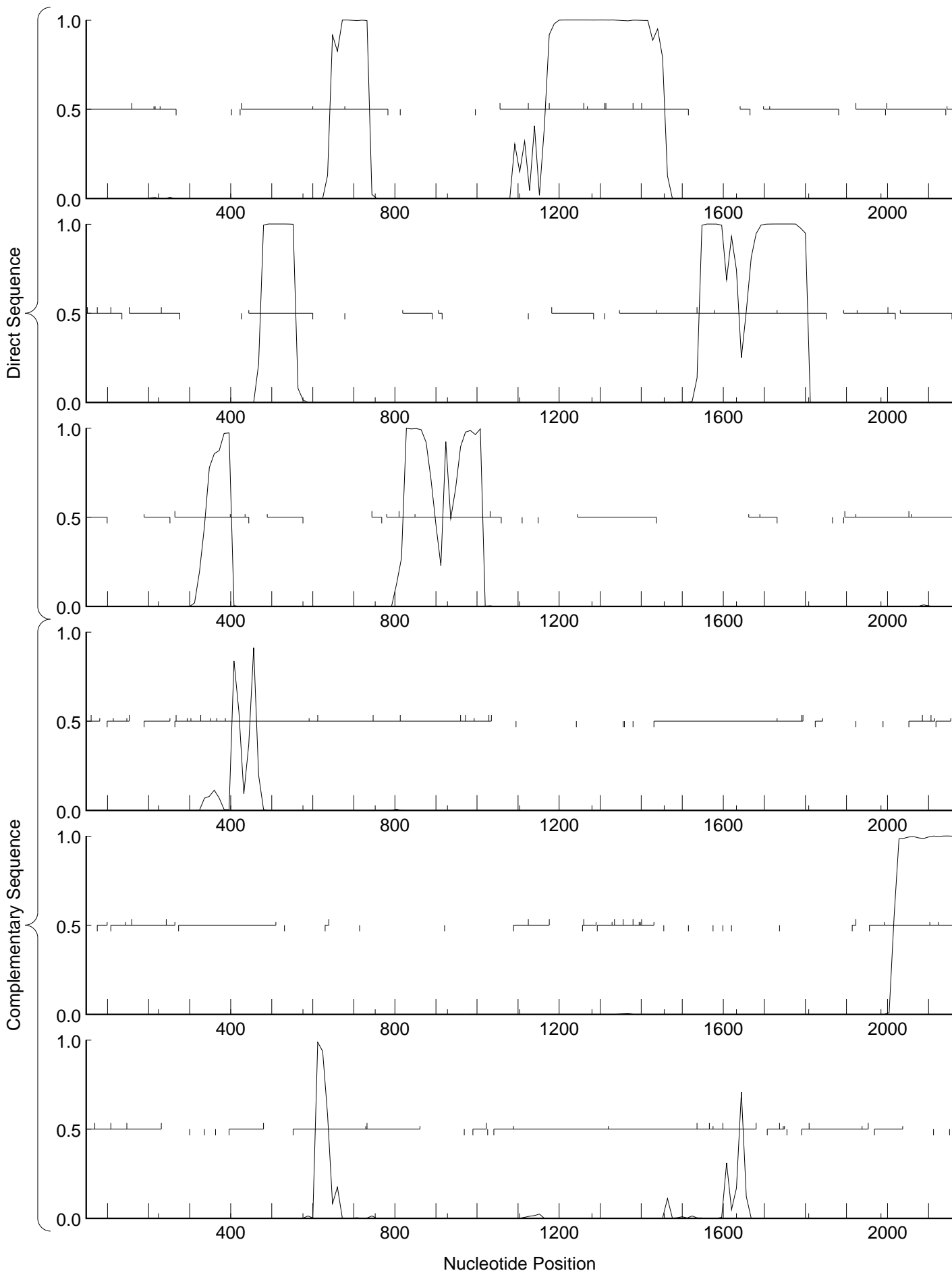
## Matrix notes & comments

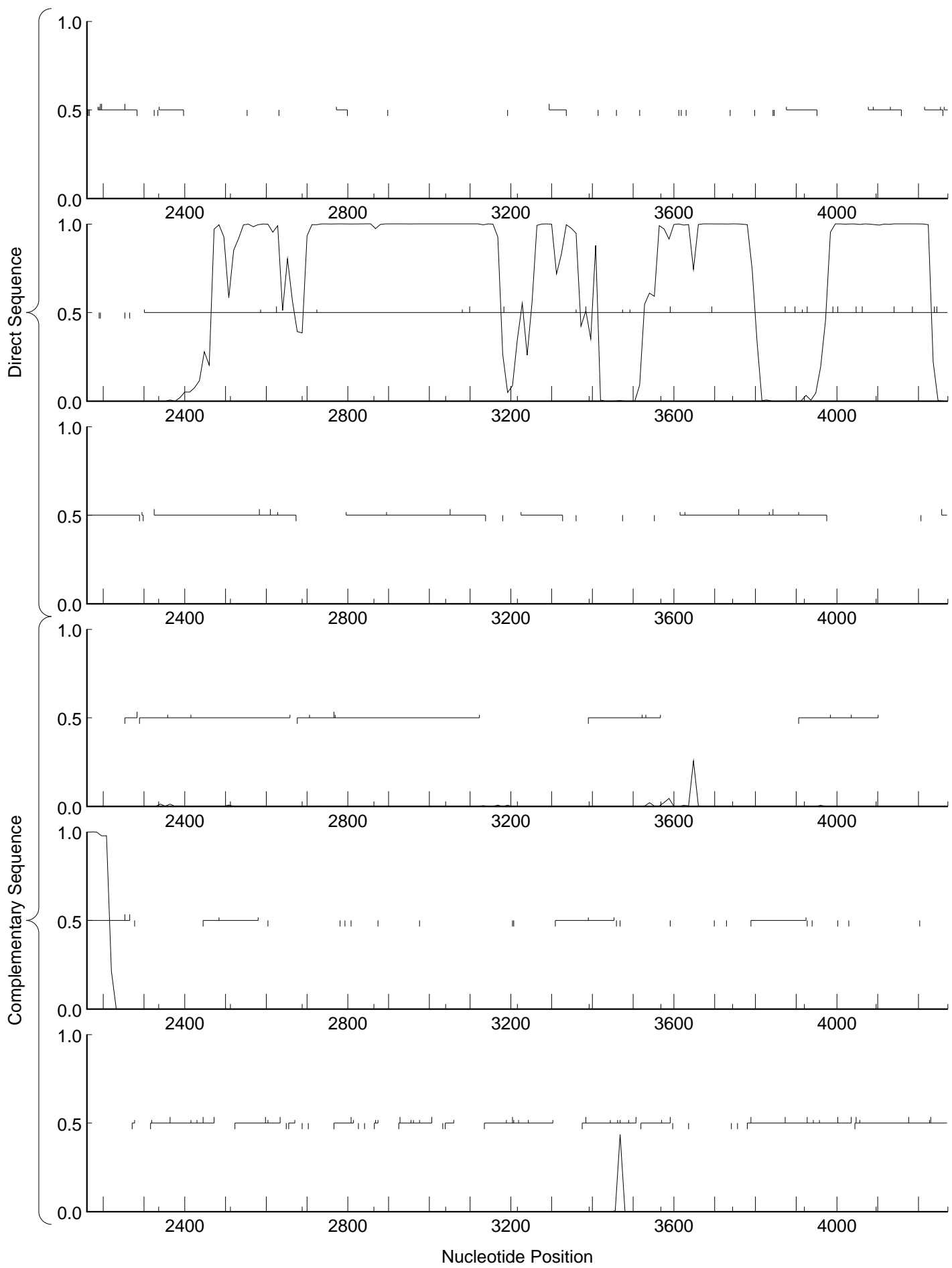
Training set derived by GeneMarkS, 4.27 September 2014  
Tue Sep 23 15:23:03 2014

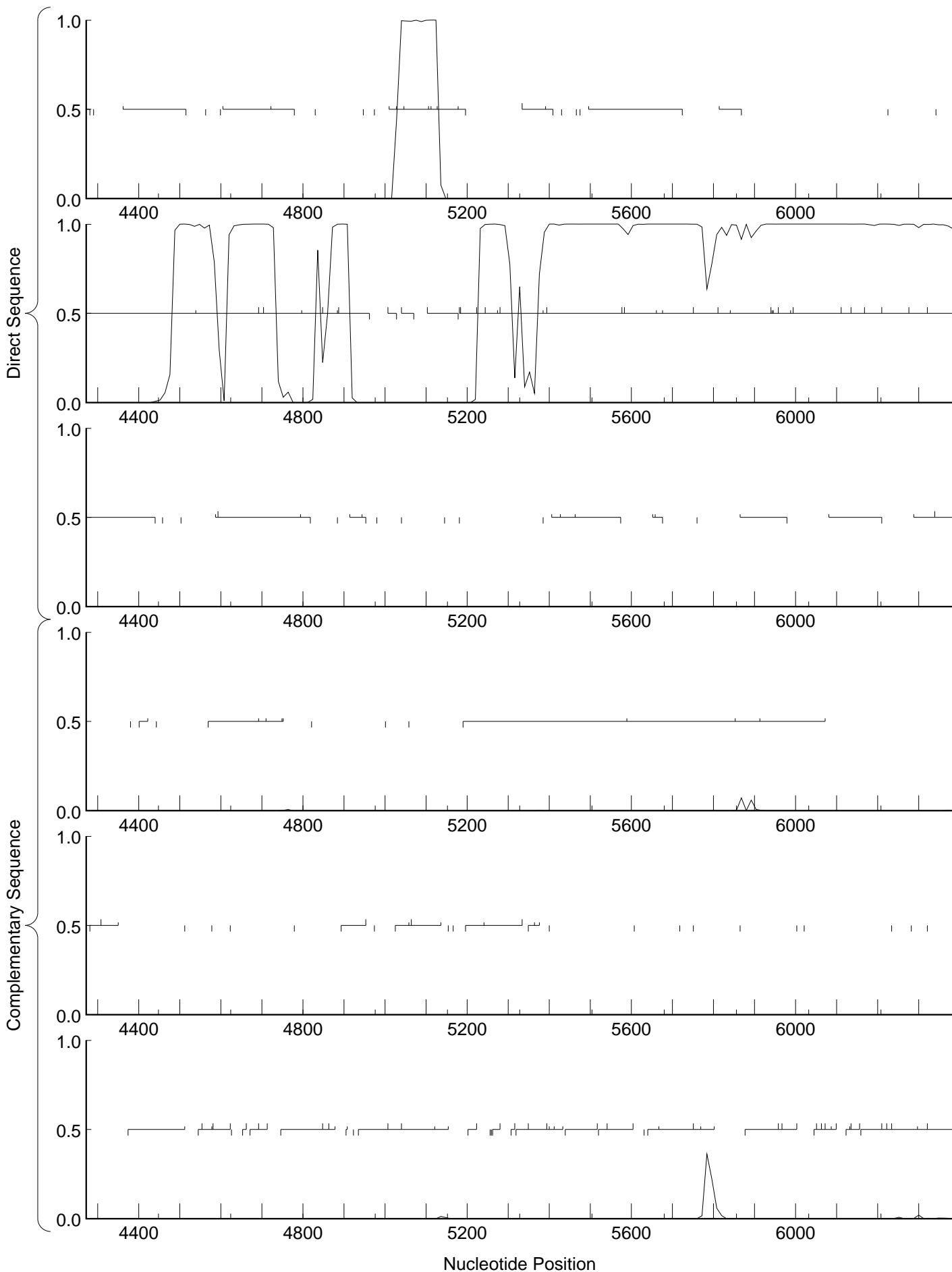
## MATRIX INFORMATION

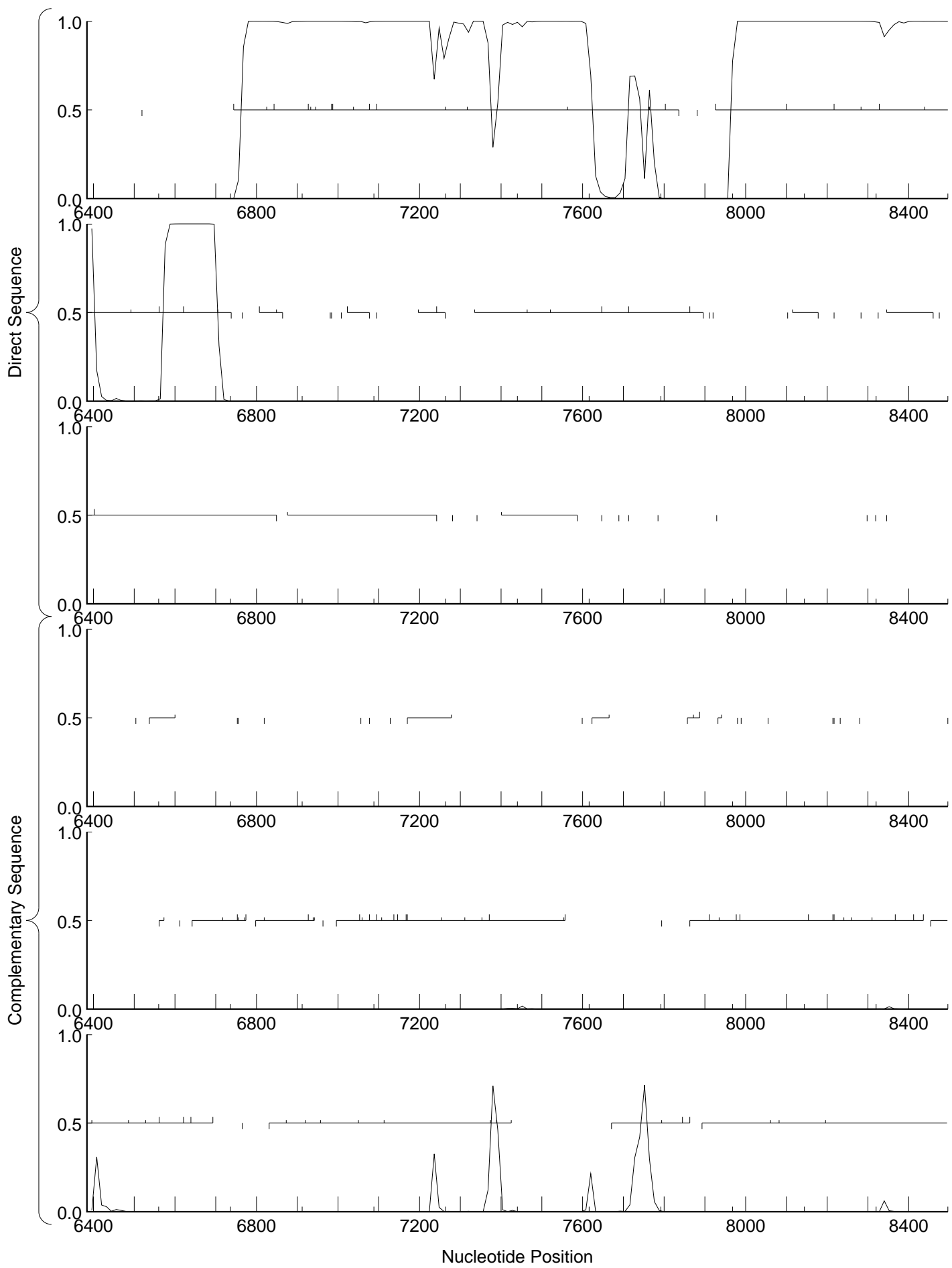
Matrix : Mycobacterium\_smegmatis\_MC2\_155  
Author : -  
Order : 4

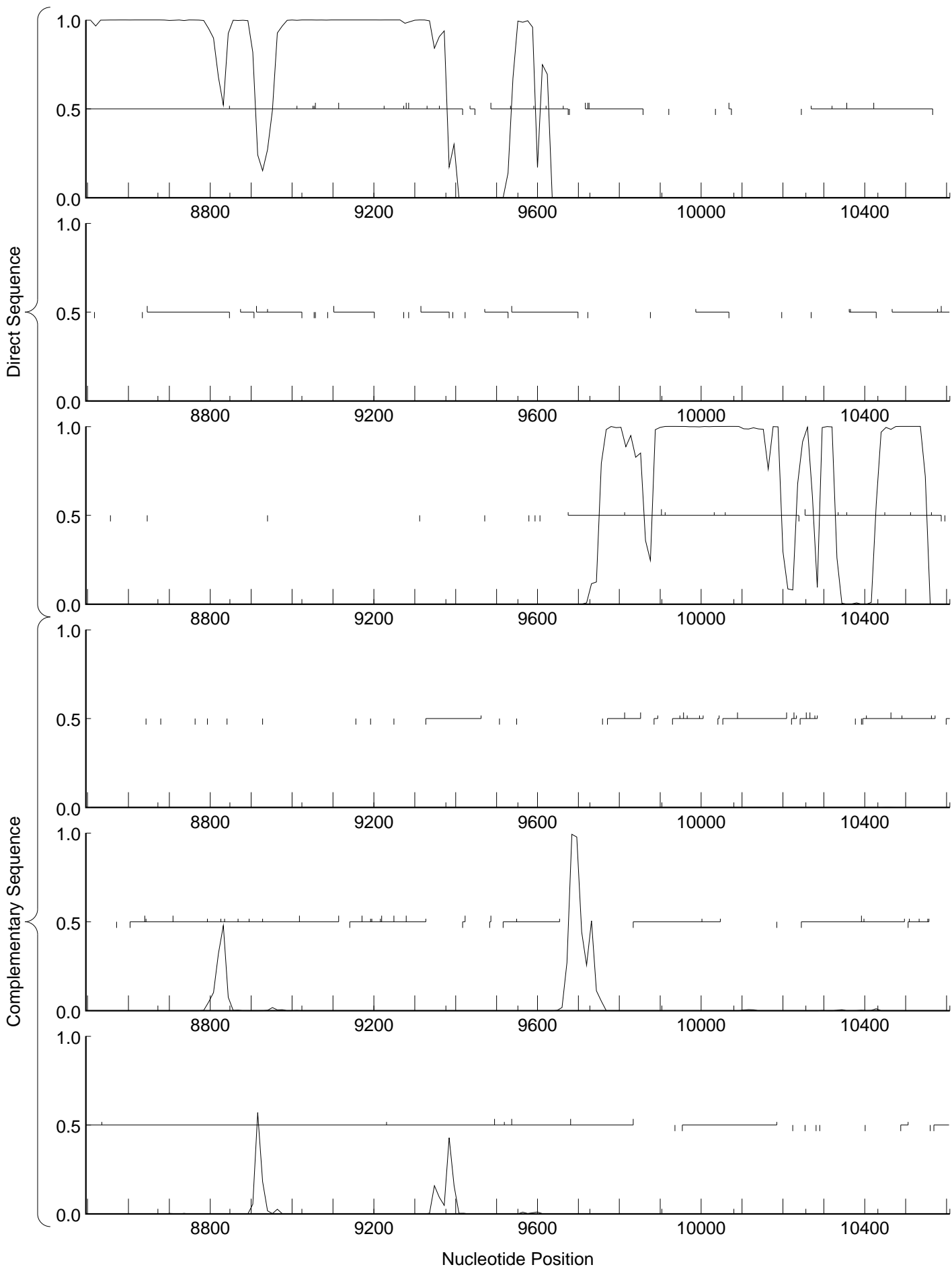
Send questions / comments to:  
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Atlanta, GA 30332-0230

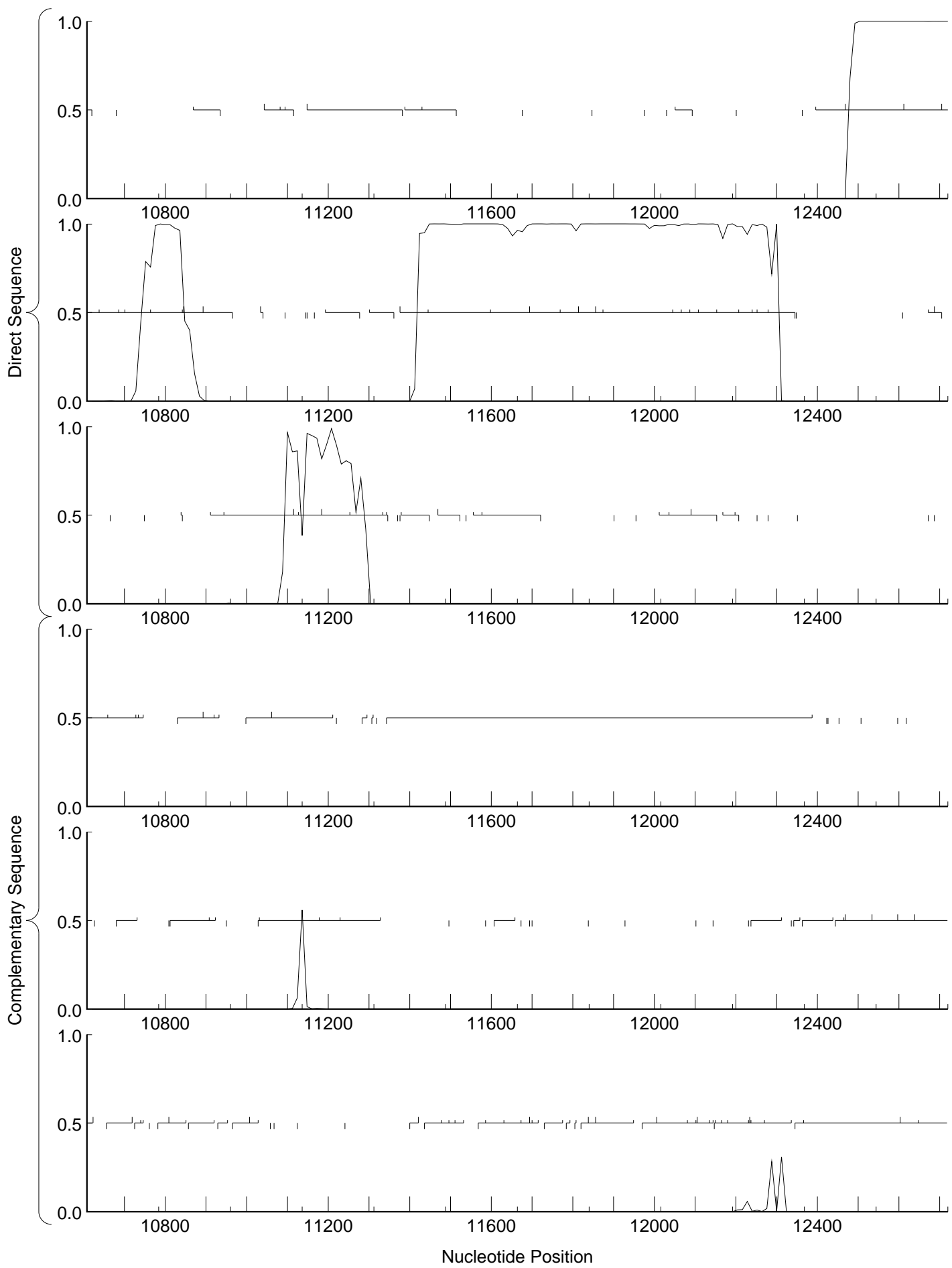


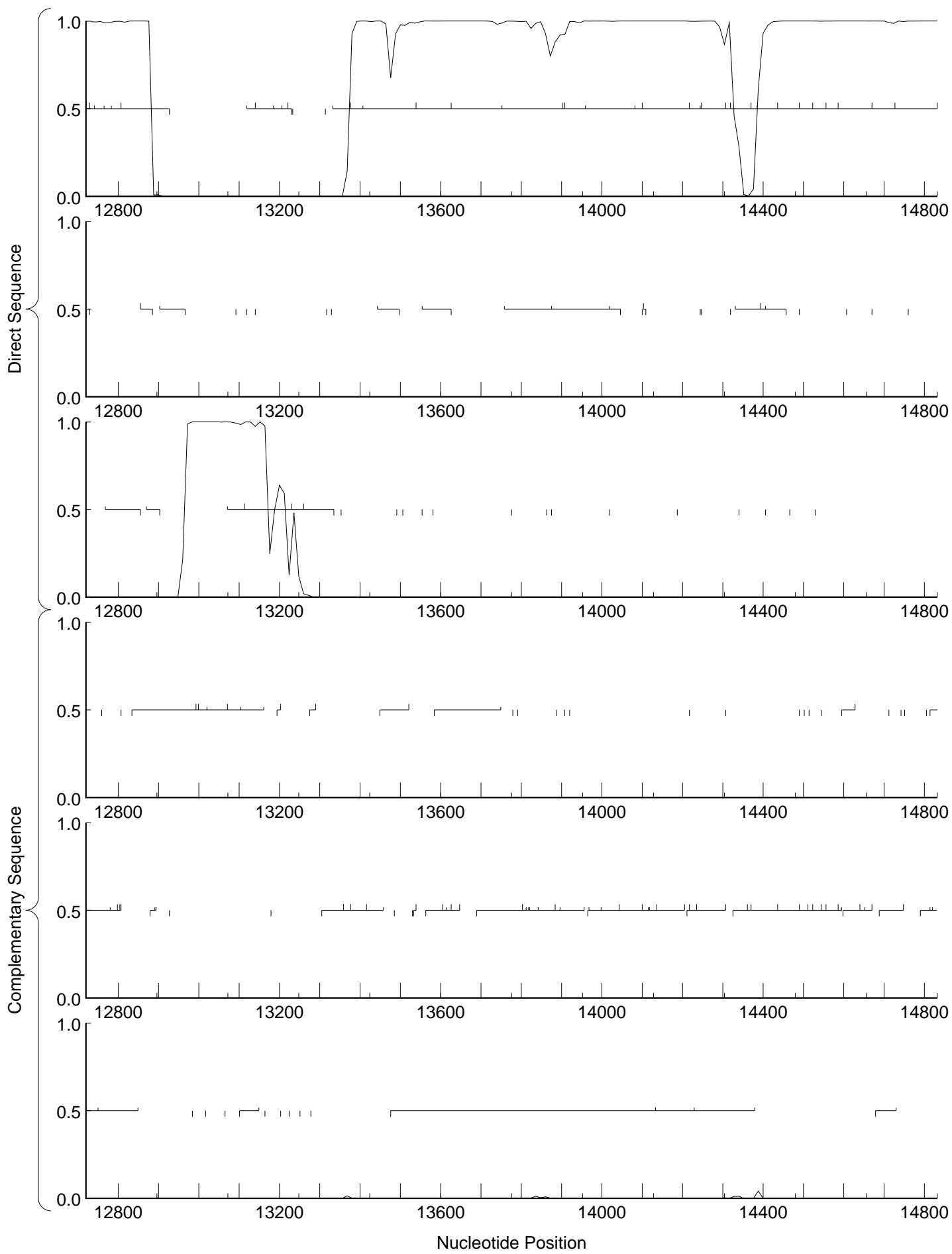














Stone complete sequence, 75679 bp including 9-base 3' overhang (CGCTTGTC A), Cluster E, Order 4, Window 96, Step 12, 9/37

