

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Gordonia phage Kurt complete sequence, 68205 bp including 10-base 3' overhang (CGCCGCGTAC),  
Analysis Date : 4/1/20 at 23:38:41  
Pages : 34  
Sequence Length : 68205 bp  
GC Content : 65.80%

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

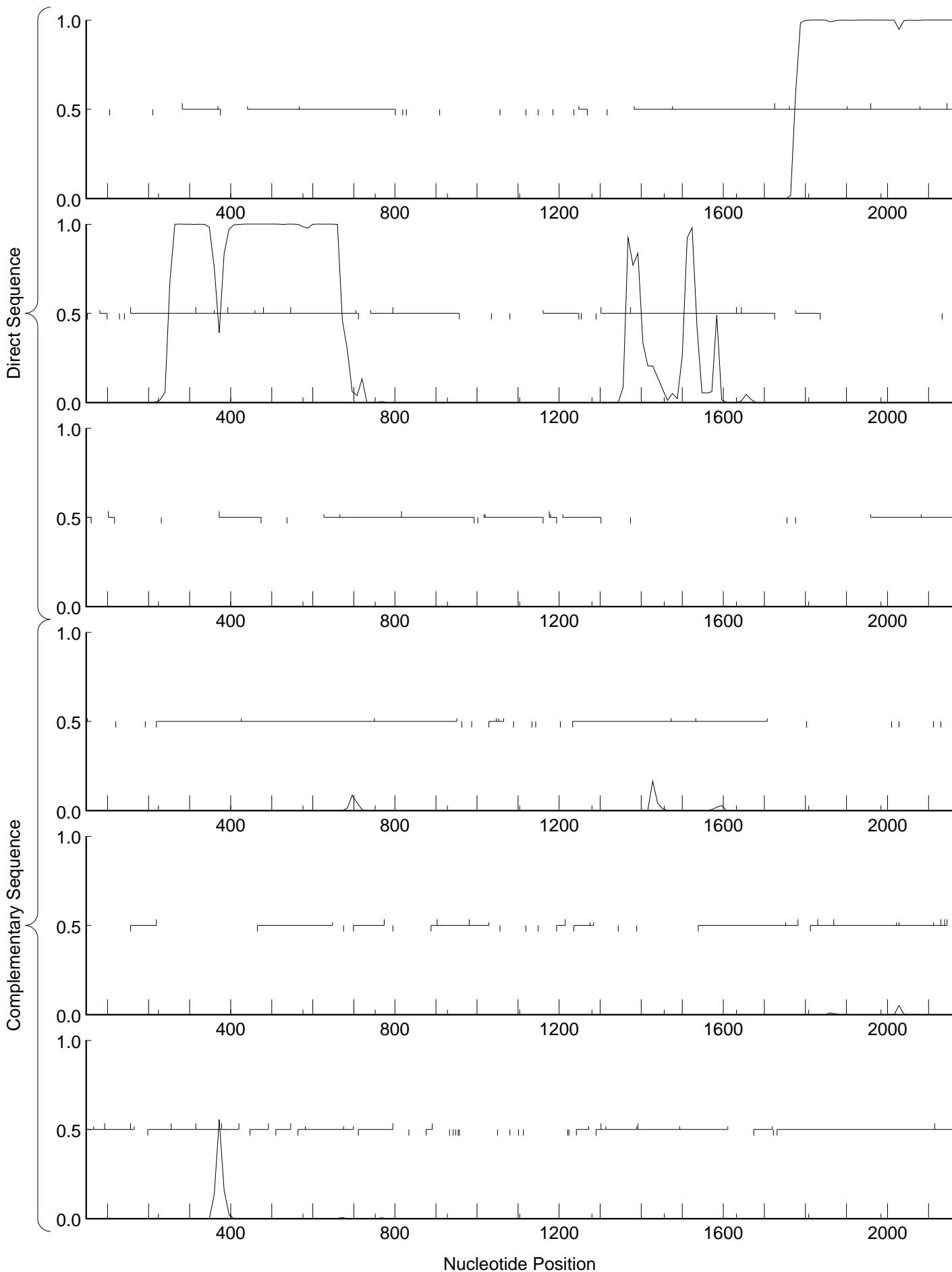
Training set derived by GeneMarkS, 4.30 October 2014  
Sat Sep 8 00:36:14 2018

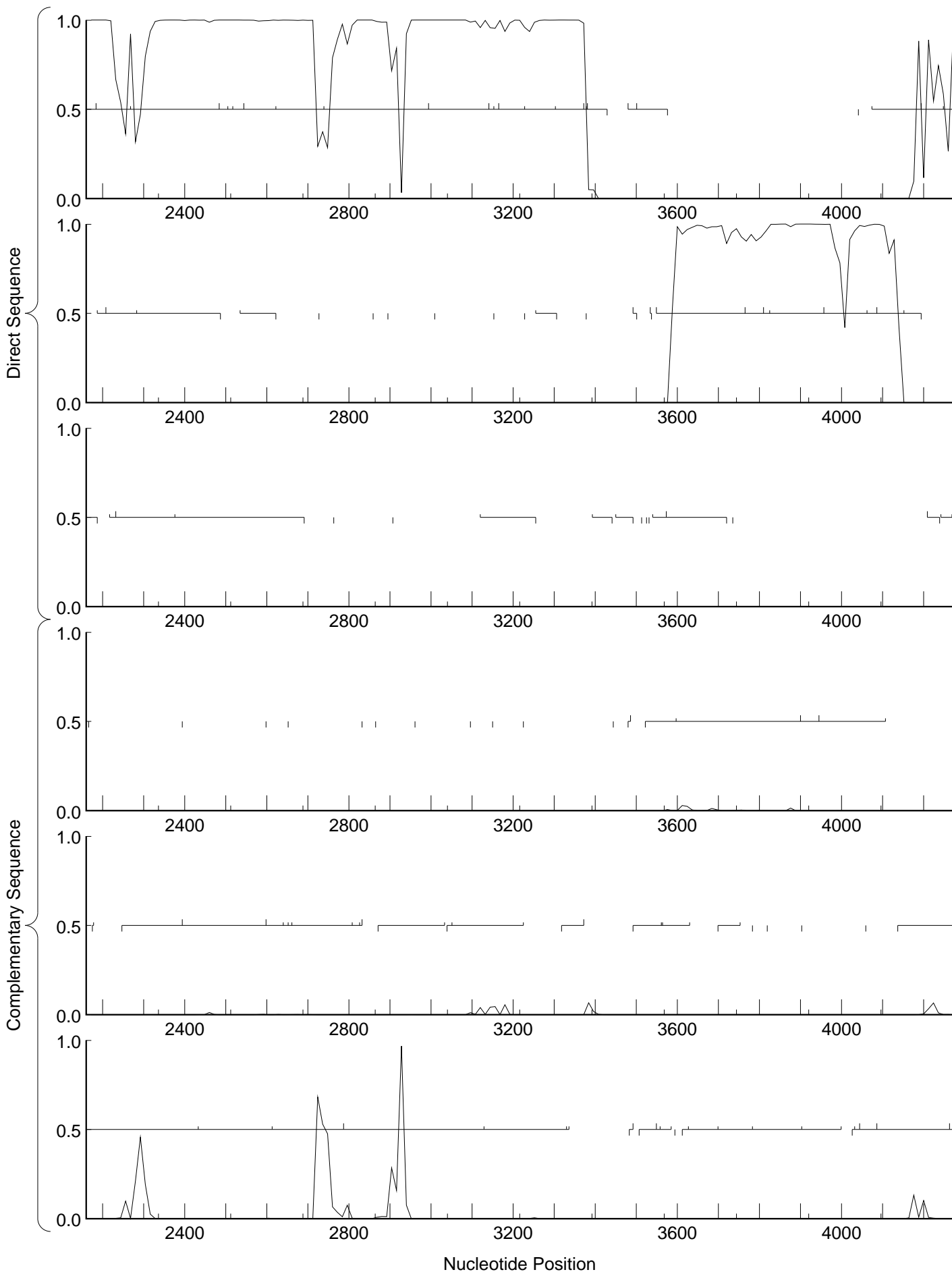
## MATRIX INFORMATION

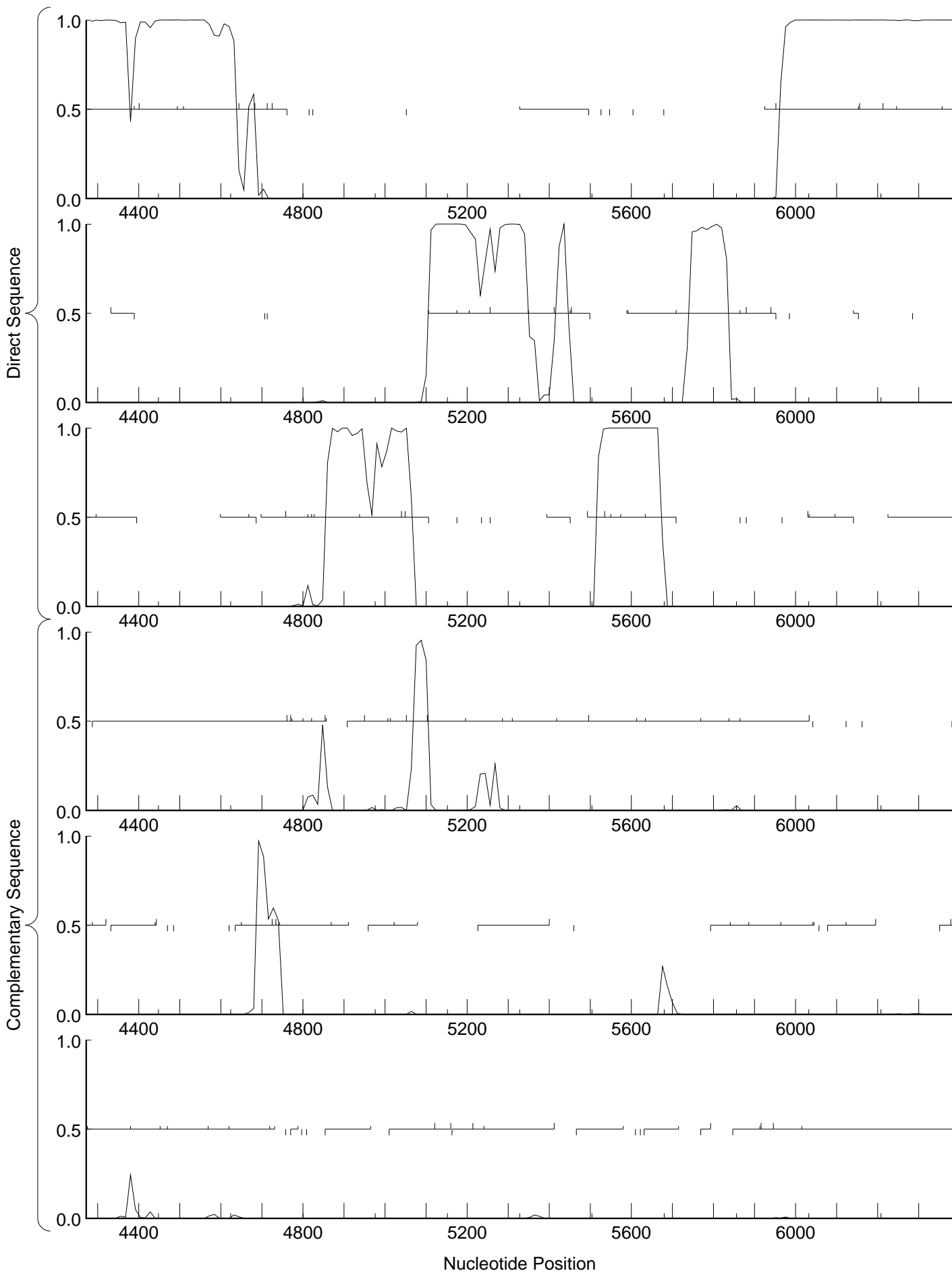
Matrix : -  
Author : -  
Order : 4

Send questions / comments to:  
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For help or information, send electronic mail to: [custserv@genepro.com](mailto:custserv@genepro.com), or [mark@amber.biology.gatech.edu](mailto:mark@amber.biology.gatech.edu)







t complete sequence, 68205 bp including 10-base 3' overhang (CGCCGCGTAC), Cluster CR, Order 4, Window 96, Step 12, 5/34

