

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Xerxes Complete Sequence, 43698 bp including 14-base 3' overhang (CCCGC
Analysis Date : 9/28/20 at 9:31:46
Pages : 22
Sequence Length : 43698 bp
GC Content : 66.26%

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,

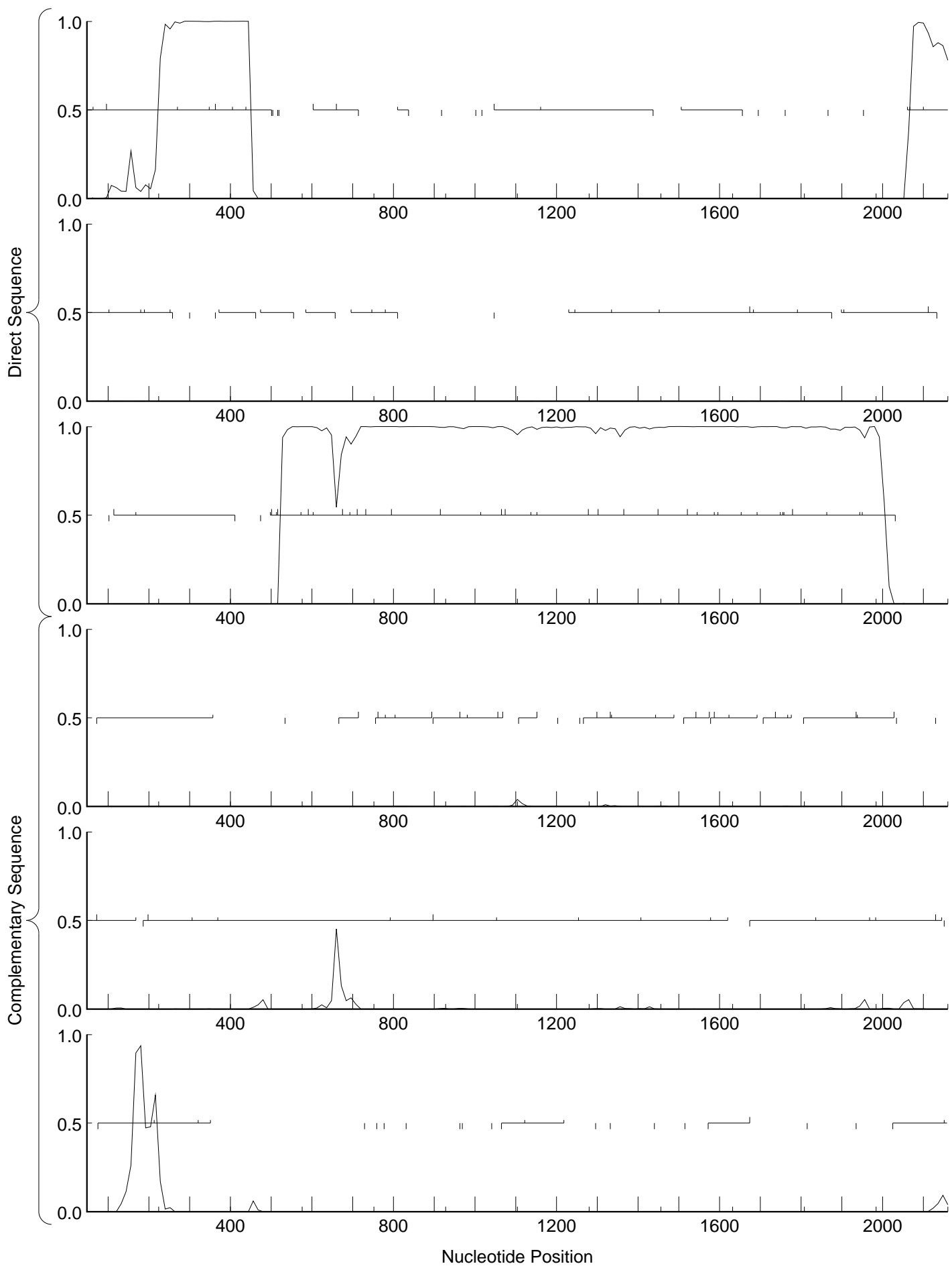
Matrix notes & comments

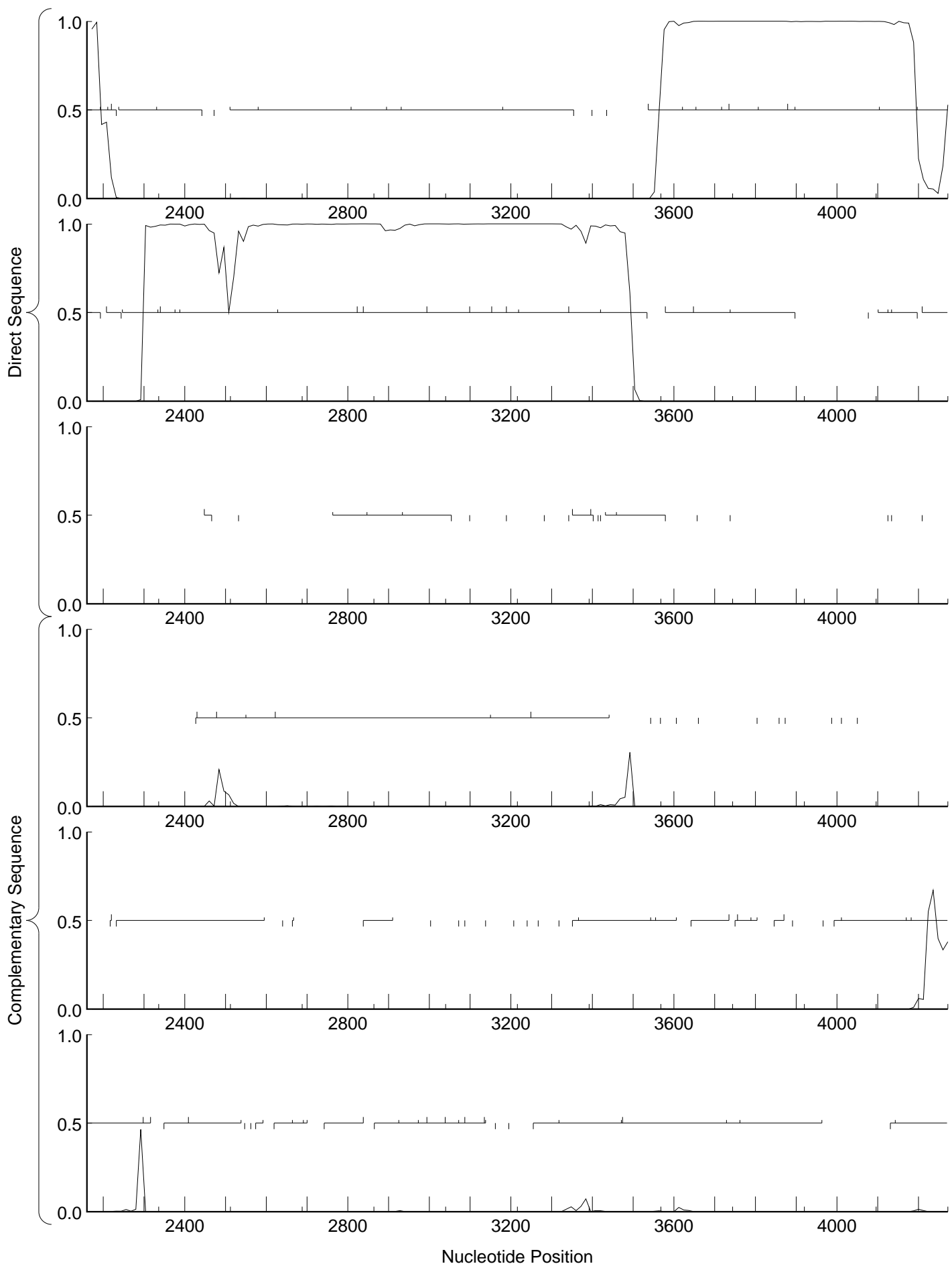
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

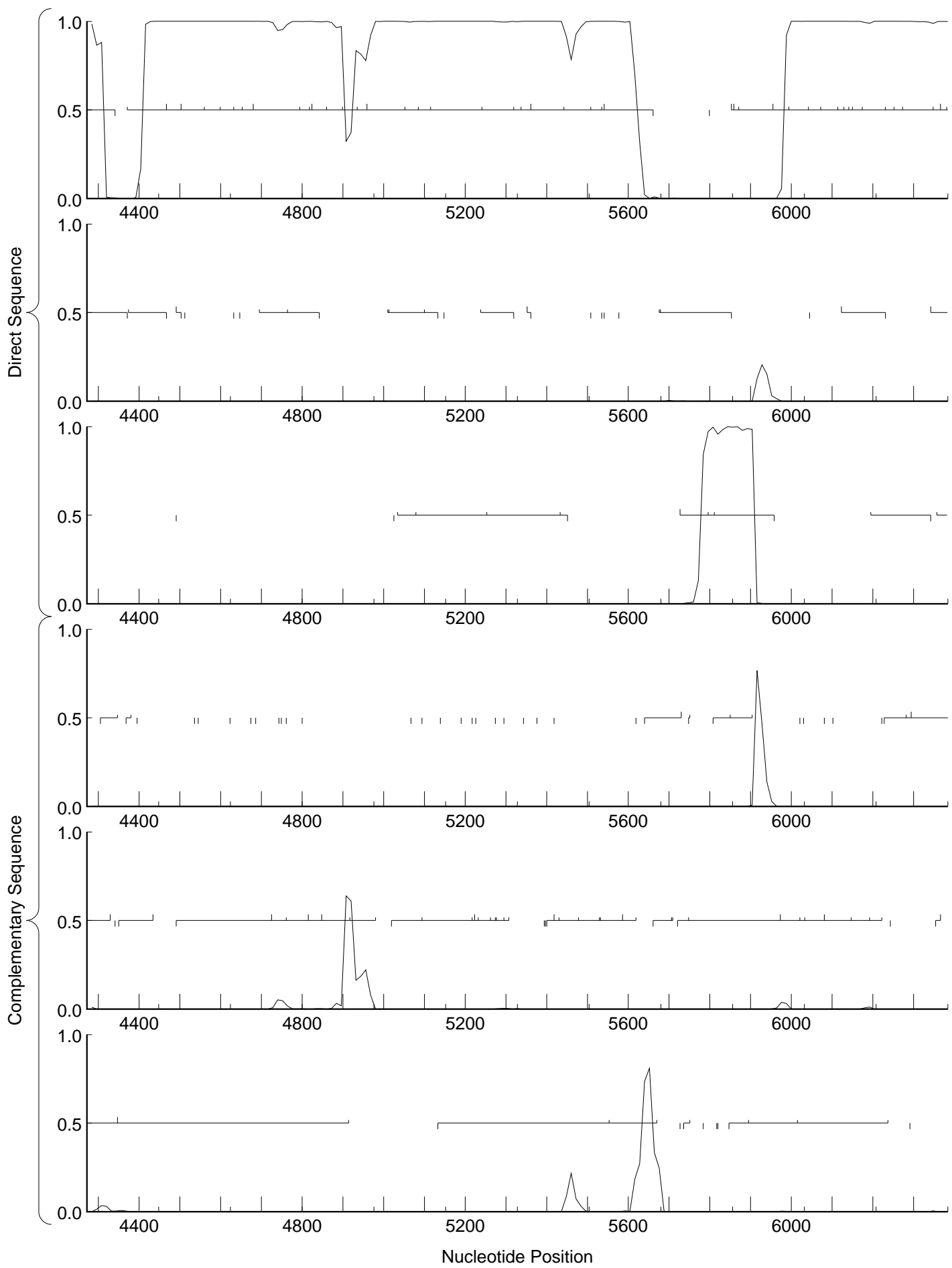
MATRIX INFORMATION

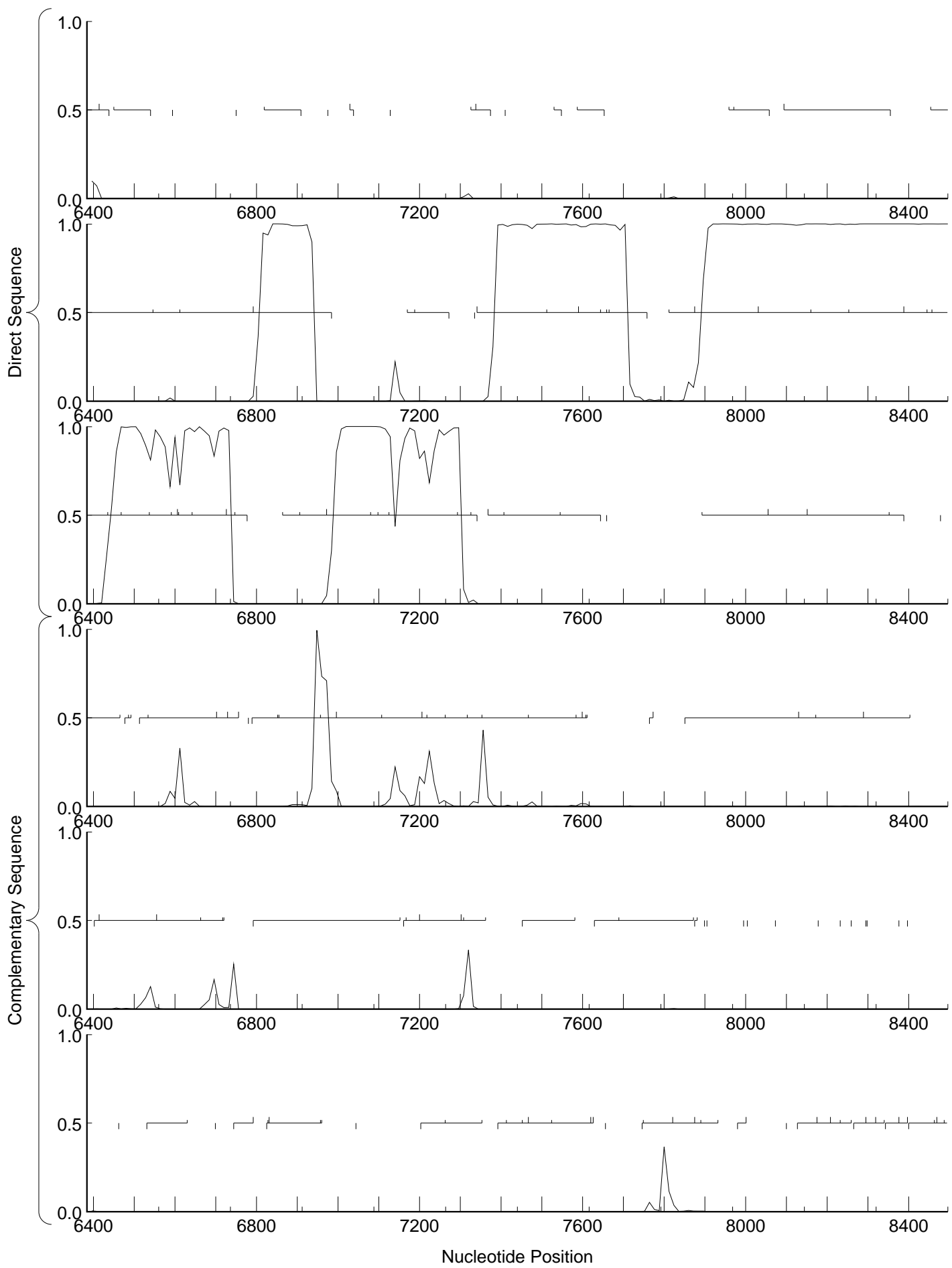
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

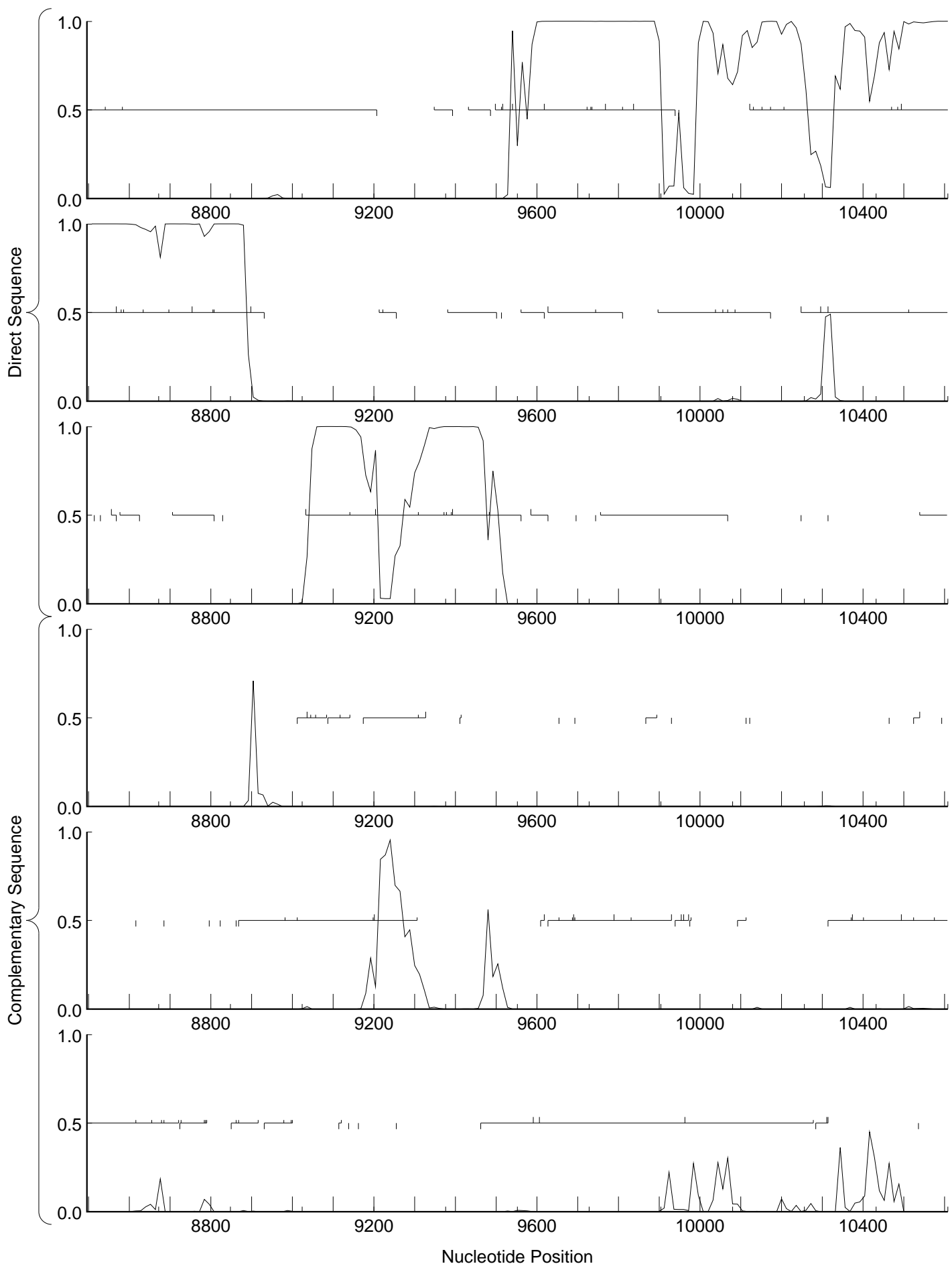
Send questions / comments to:
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Atlanta, GA 30332-0230



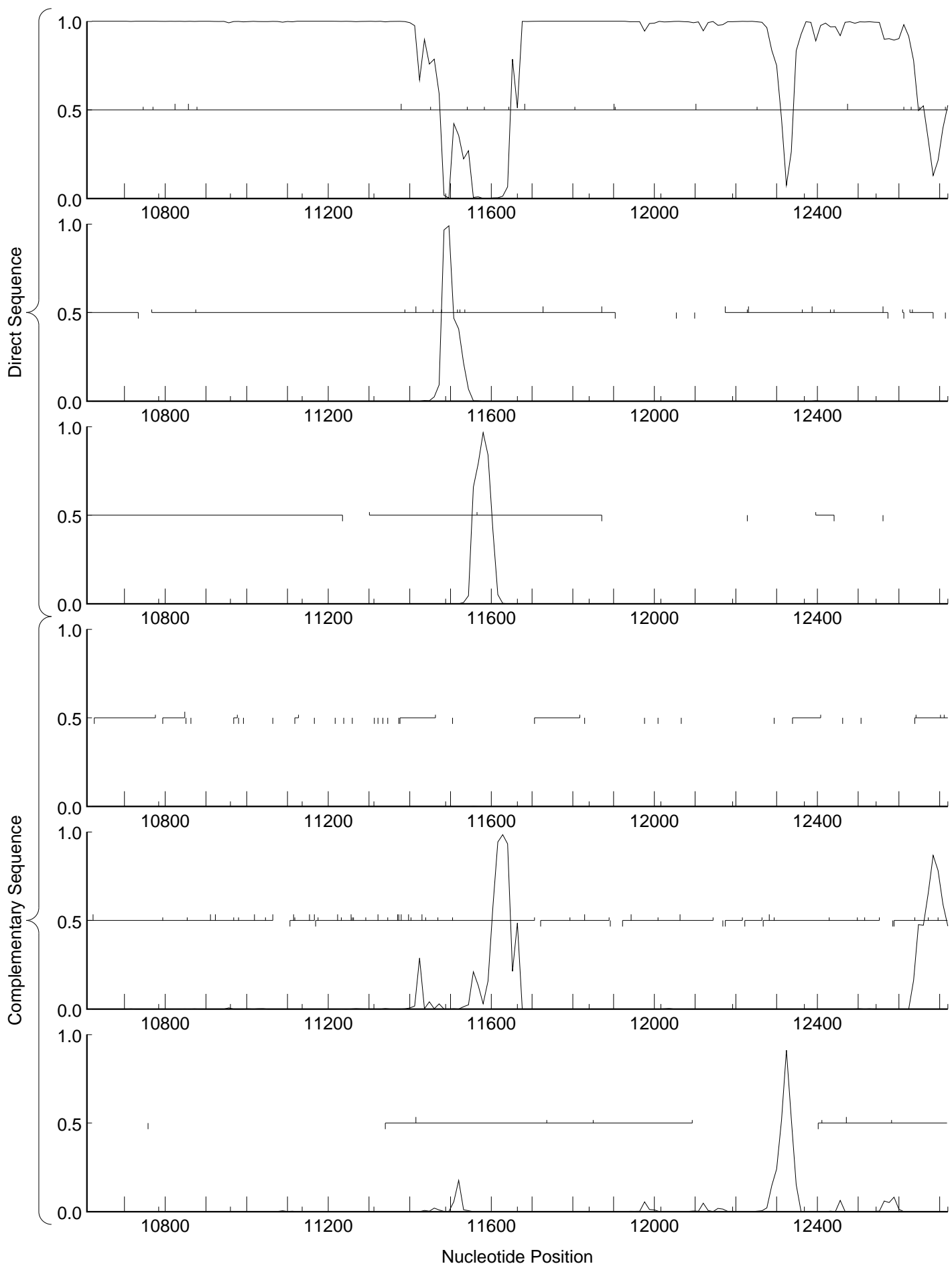


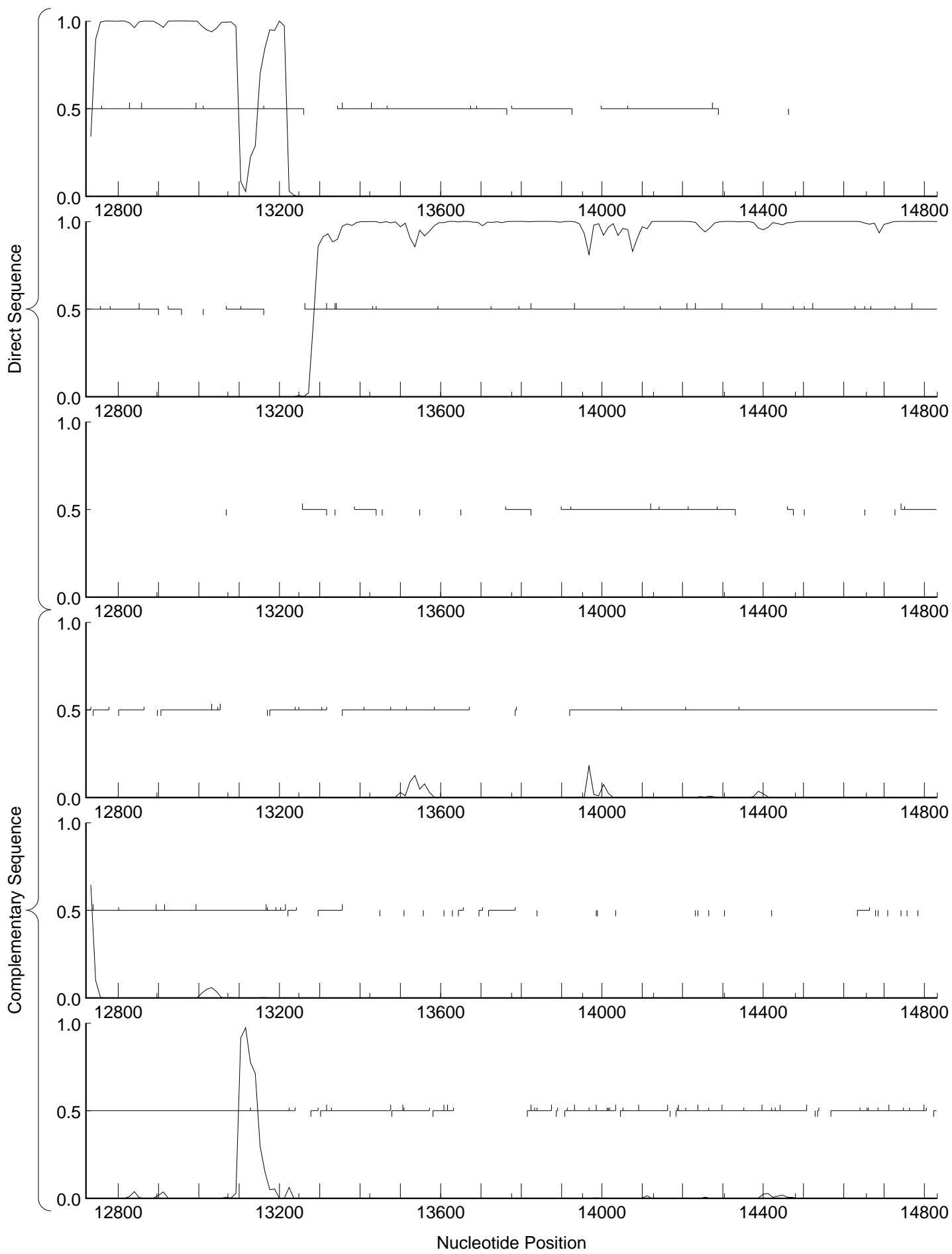






plete Sequence, 43698 bp including 14-base 3' overhang (CCCGCCGCAATGGG), Cluster N, Order 4, Window 96, Step 12, 7/22





plete Sequence, 43698 bp including 14-base 3' overhang (CCCGCCGCAATGGG), Cluster N, Order 4, Window 96, Step 12, 9/22

