

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Mycobacterium phage Quico complete sequence, 58671 bp including 10-base 3' overhang (CCGAAG
Analysis Date : 3/3/21 at 17:05:06
Pages : 29
Sequence Length : 58671 bp
GC Content : 61.69%

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,

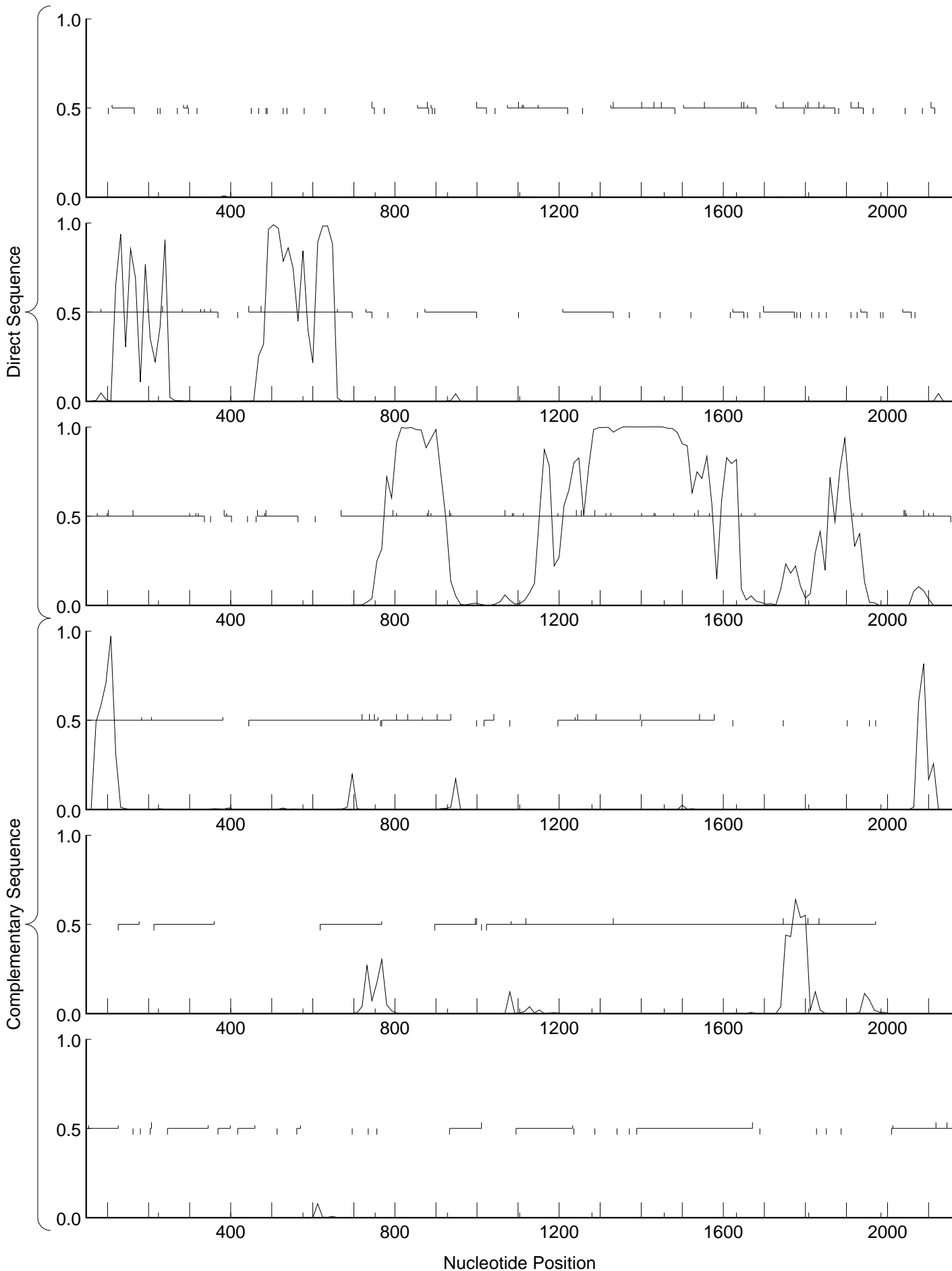
Matrix notes & comments

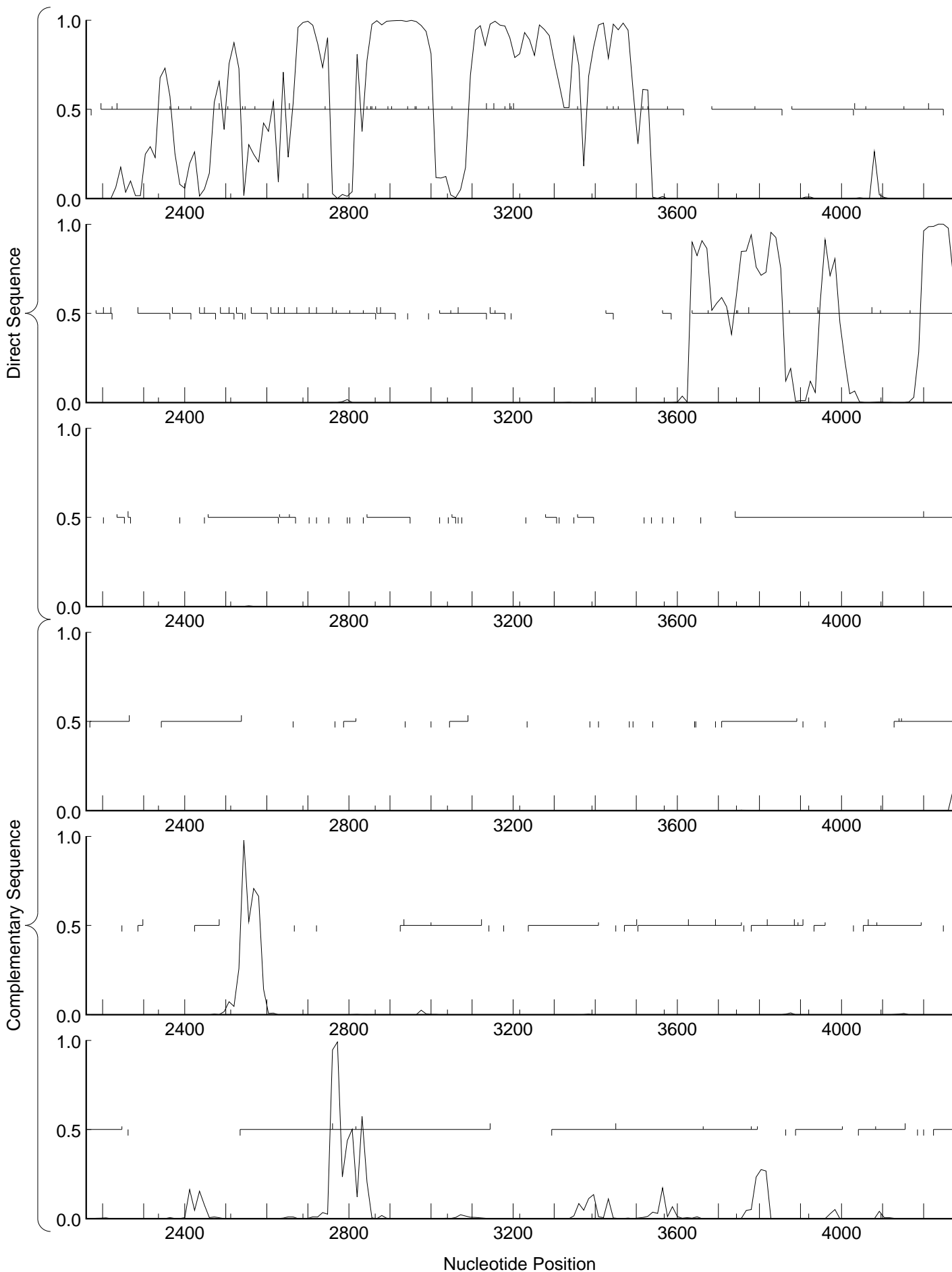
Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:14:37 2014

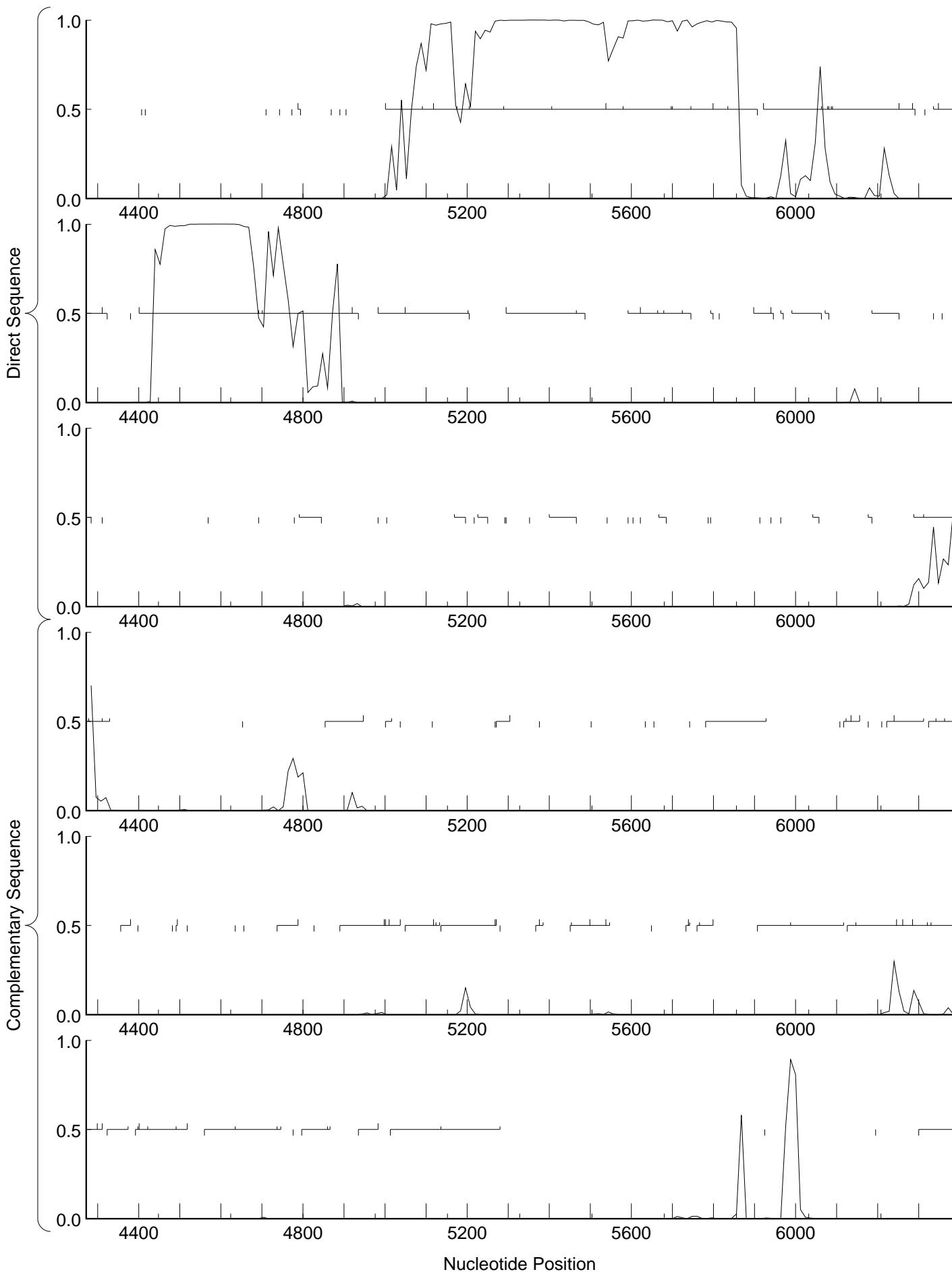
MATRIX INFORMATION

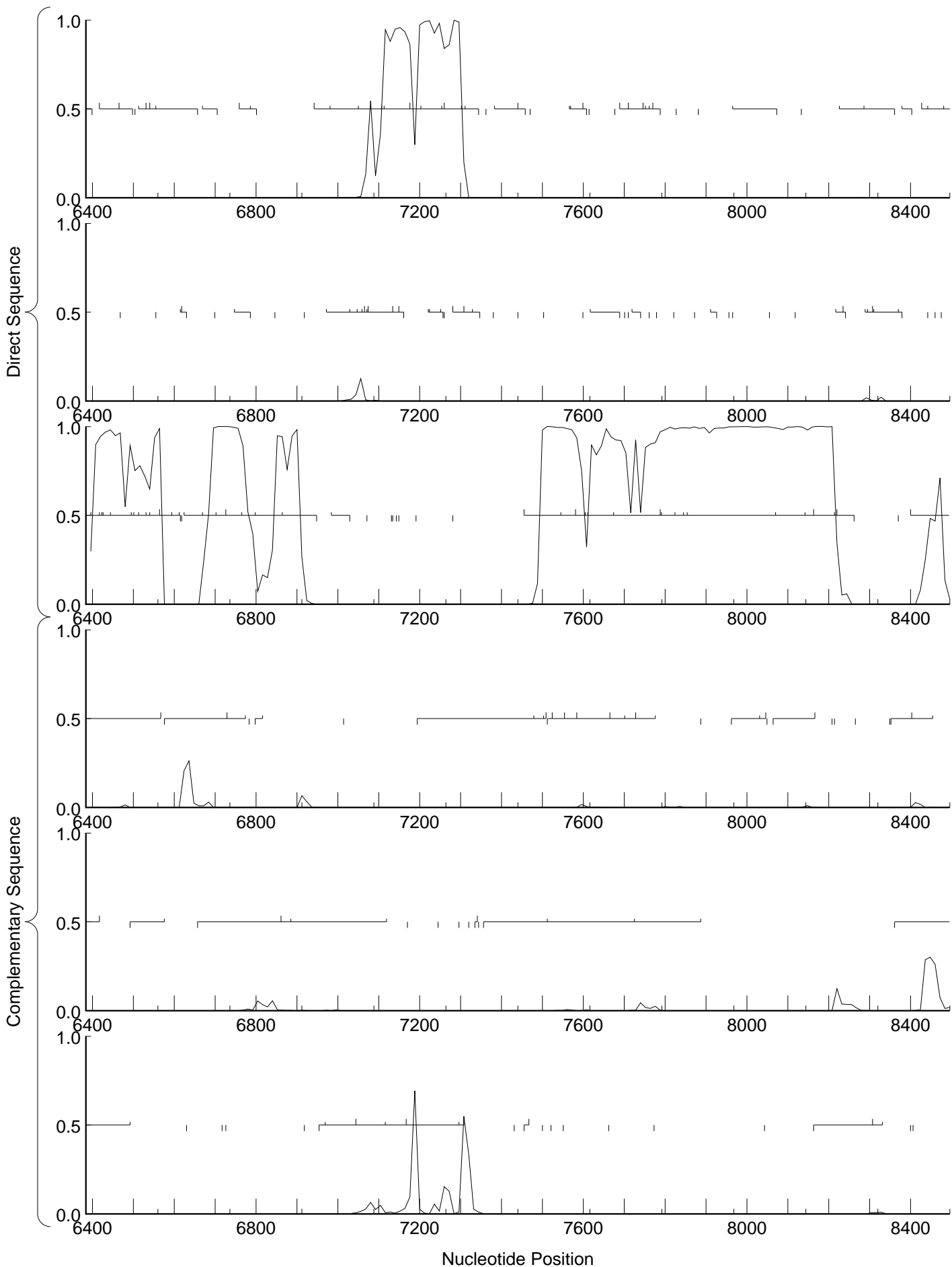
Matrix : Mycobacterium_tuberculosis_H37Rv
Author : -
Order : 4

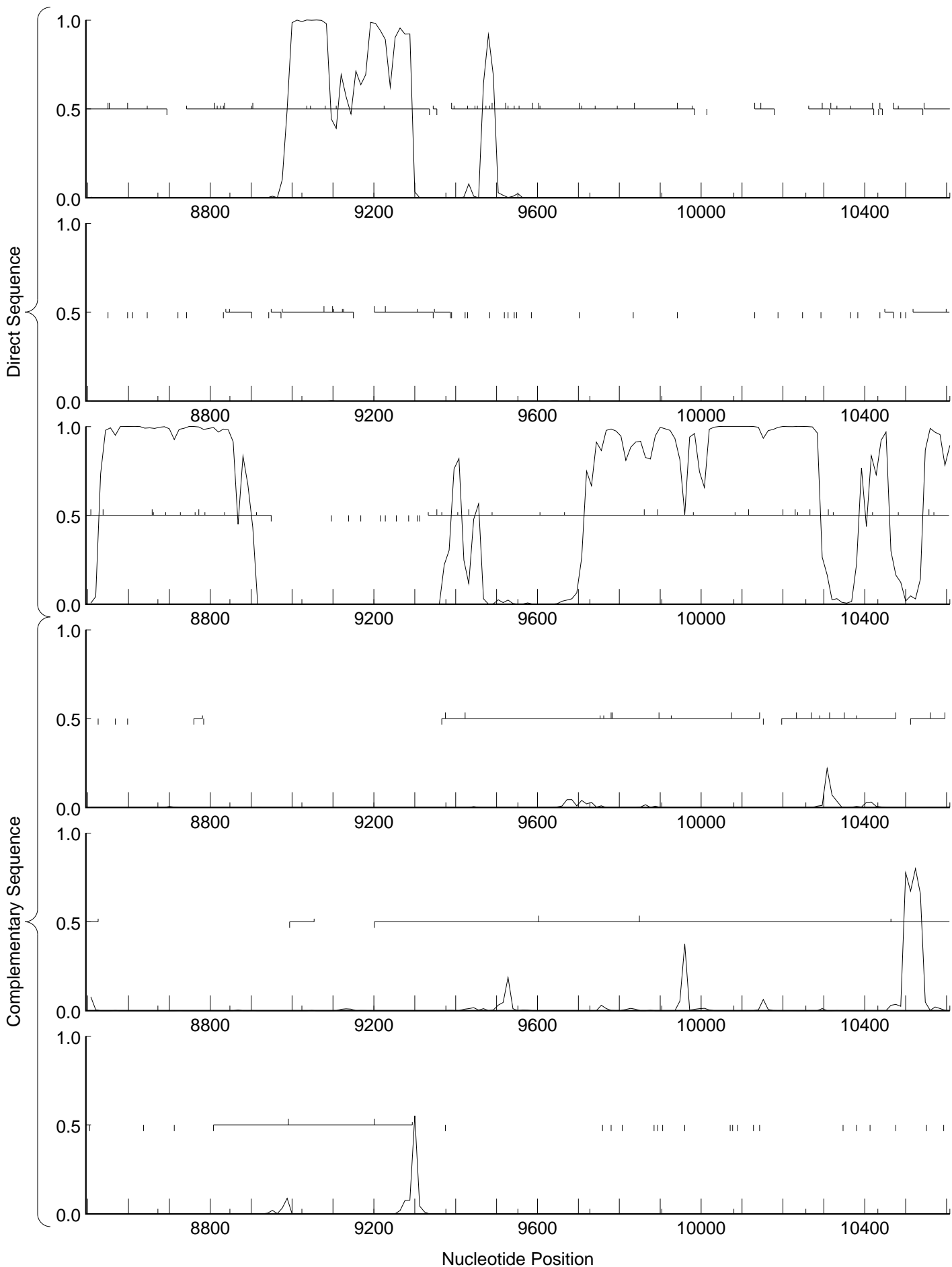
Send questions / comments to:
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Georgia Institute of Technology
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Atlanta, GA 30332-0230

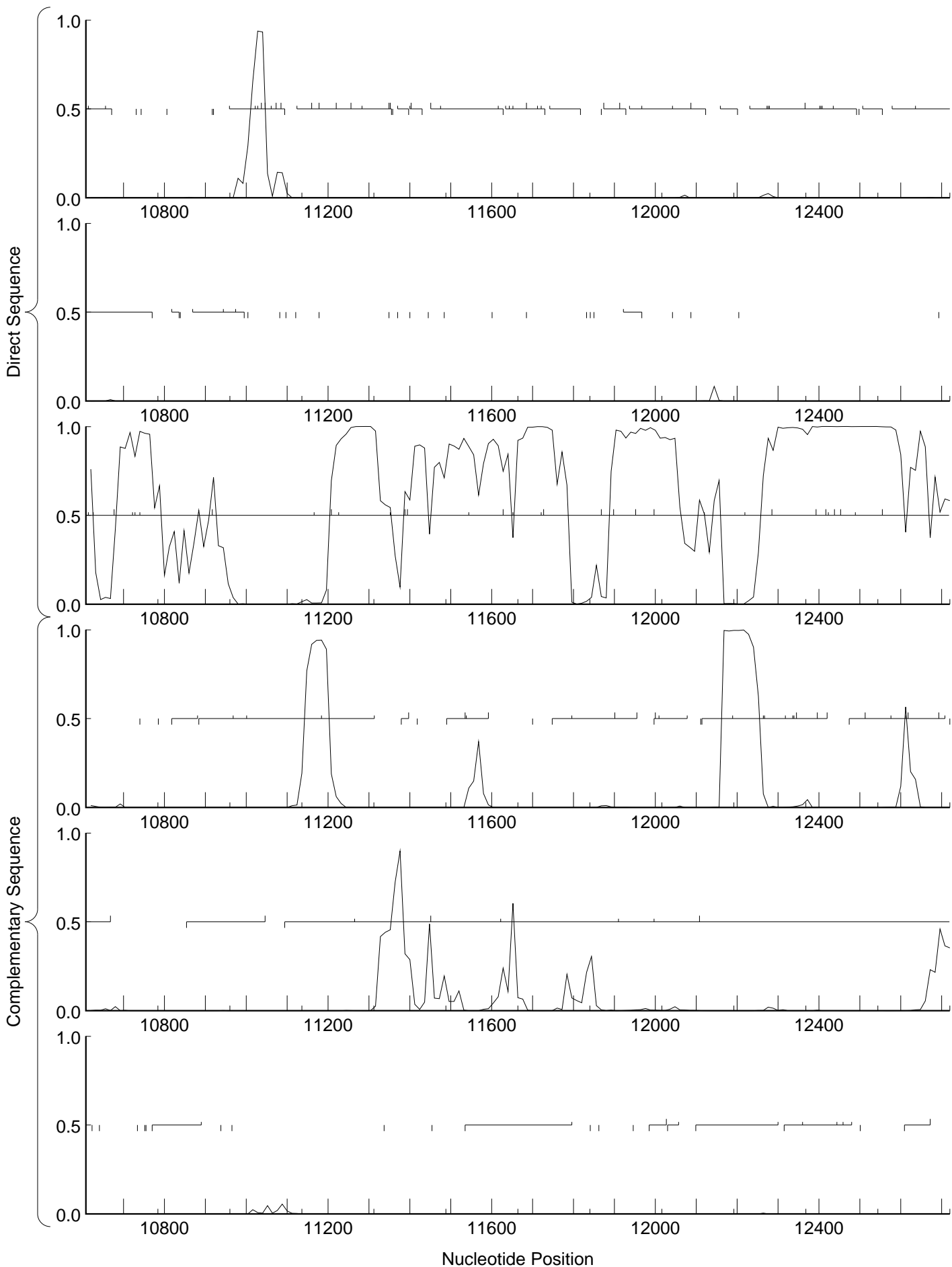


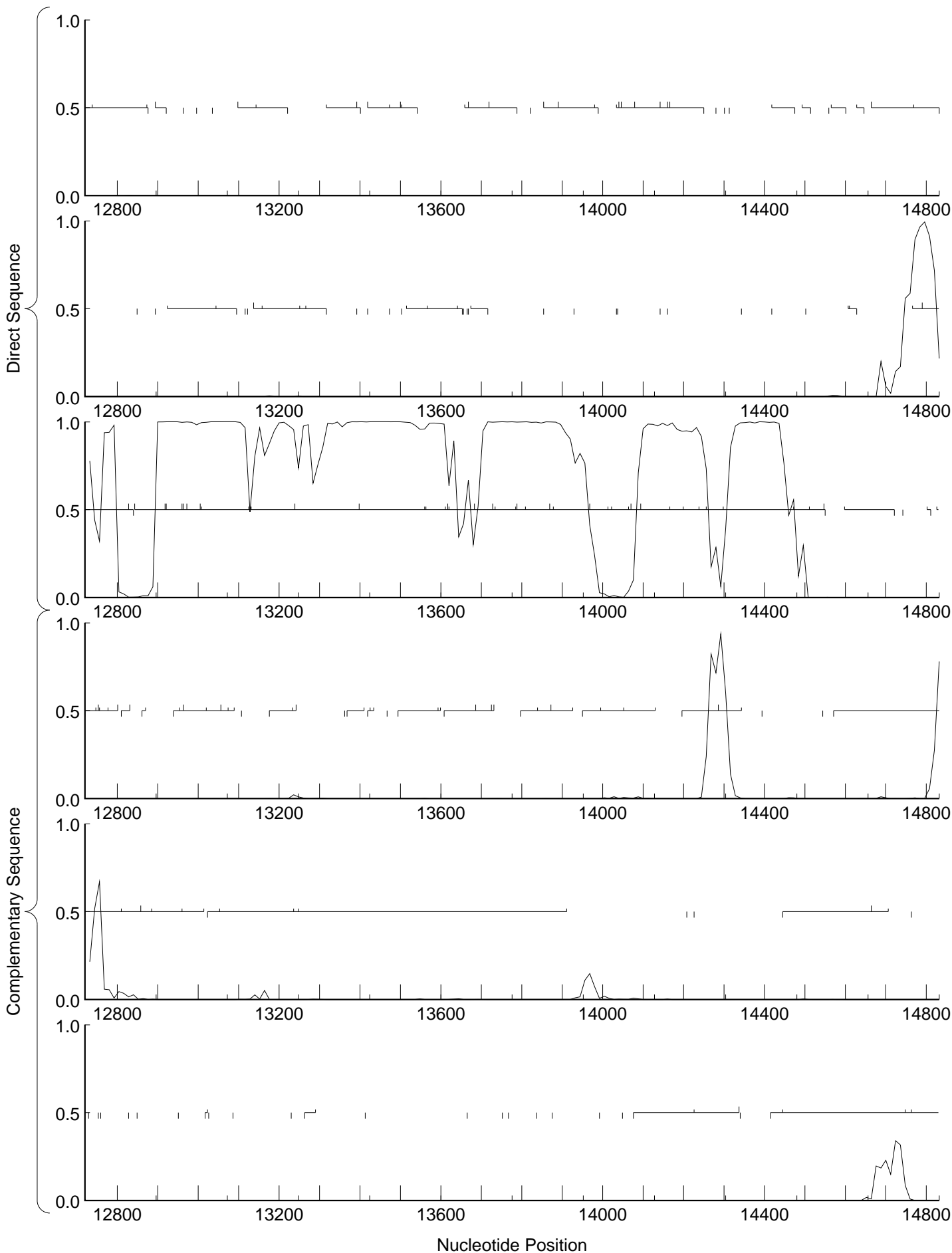


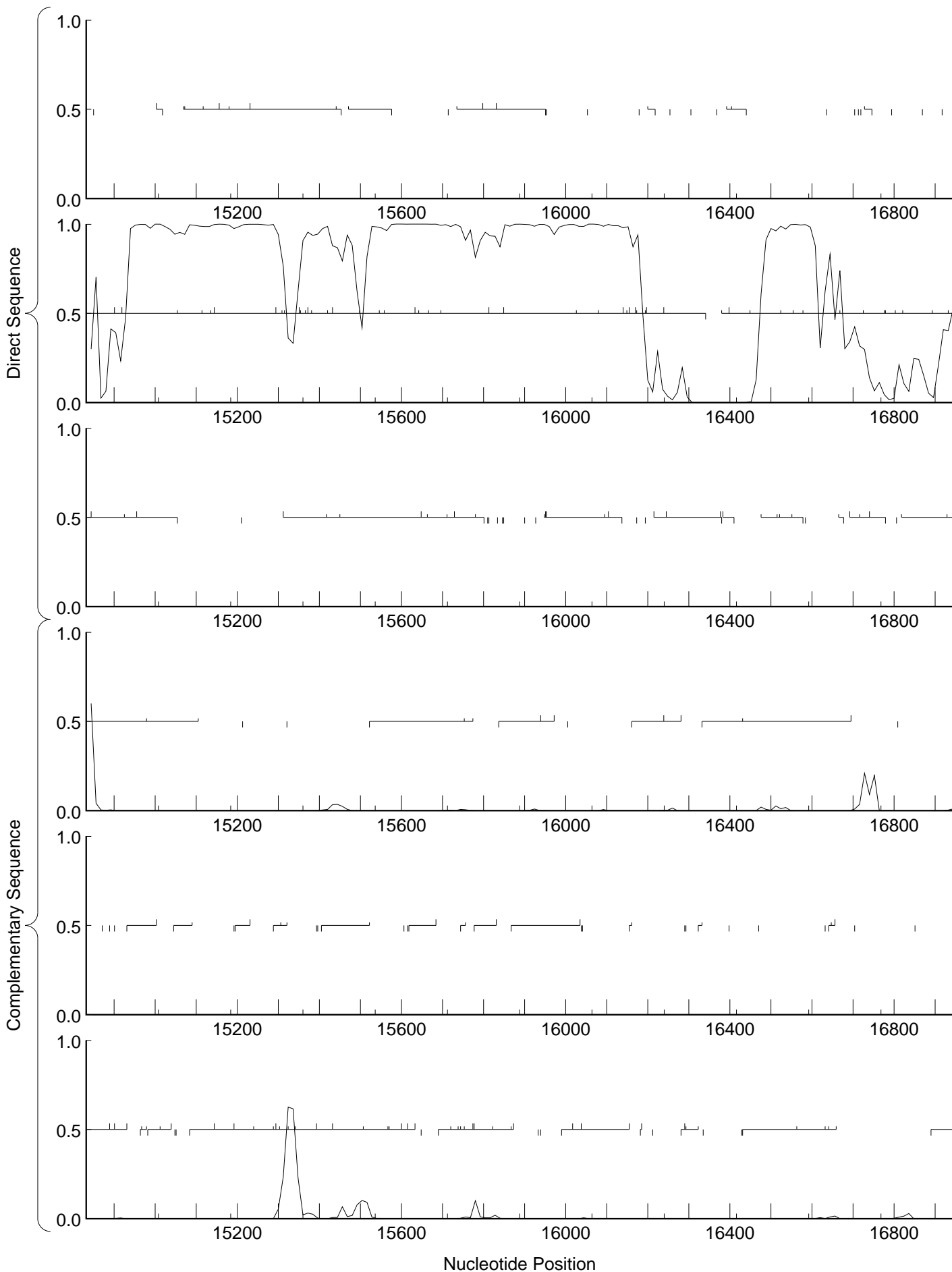












complete sequence, 58671 bp including 10-base 3' overhang (CCGAAGGCAT), Cluster F1, Order 4, Window 96, Step 12, 10/29

