

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 : 1/18/18 at 20:25:28
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,

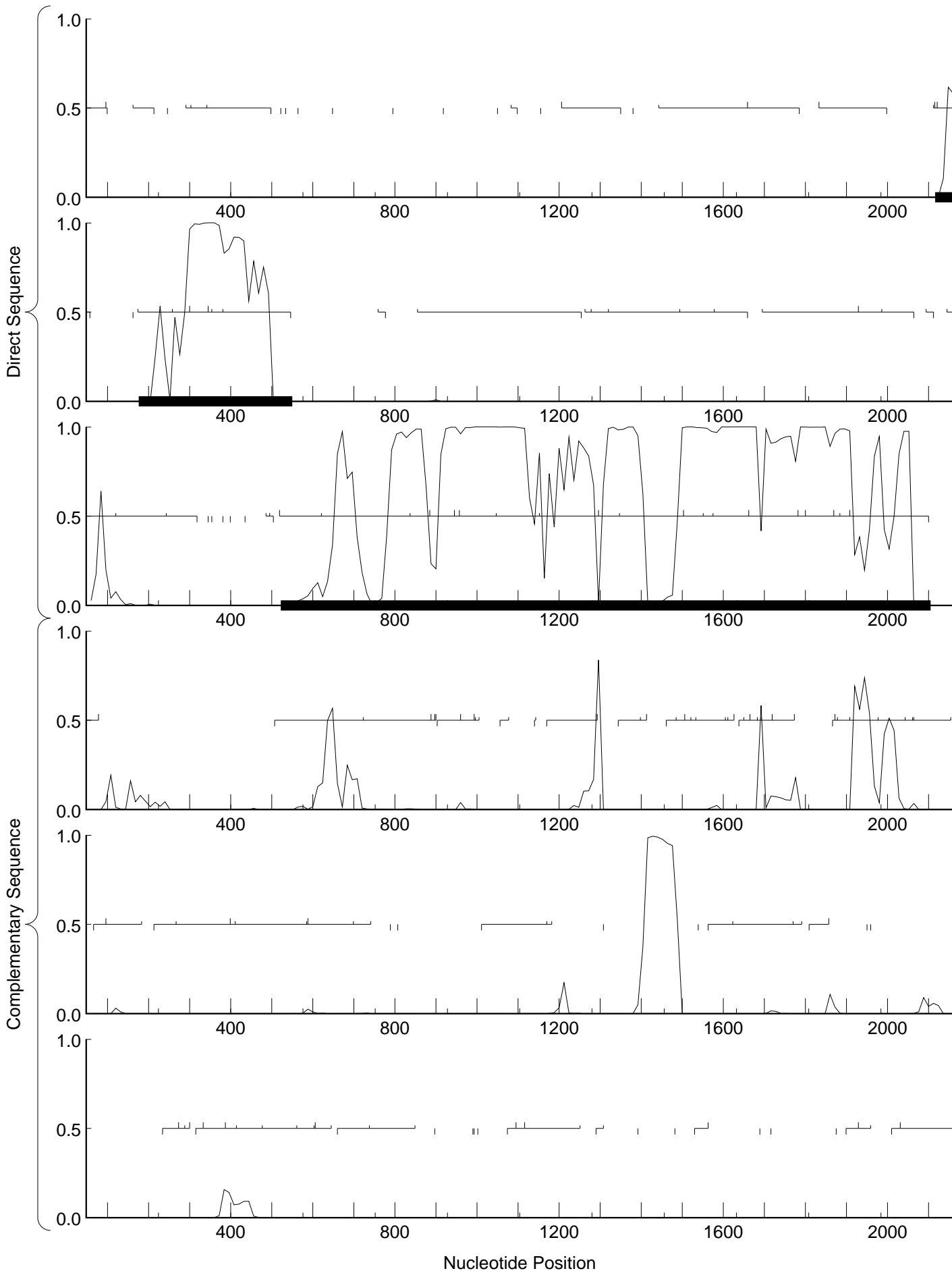
Matrix notes & comments

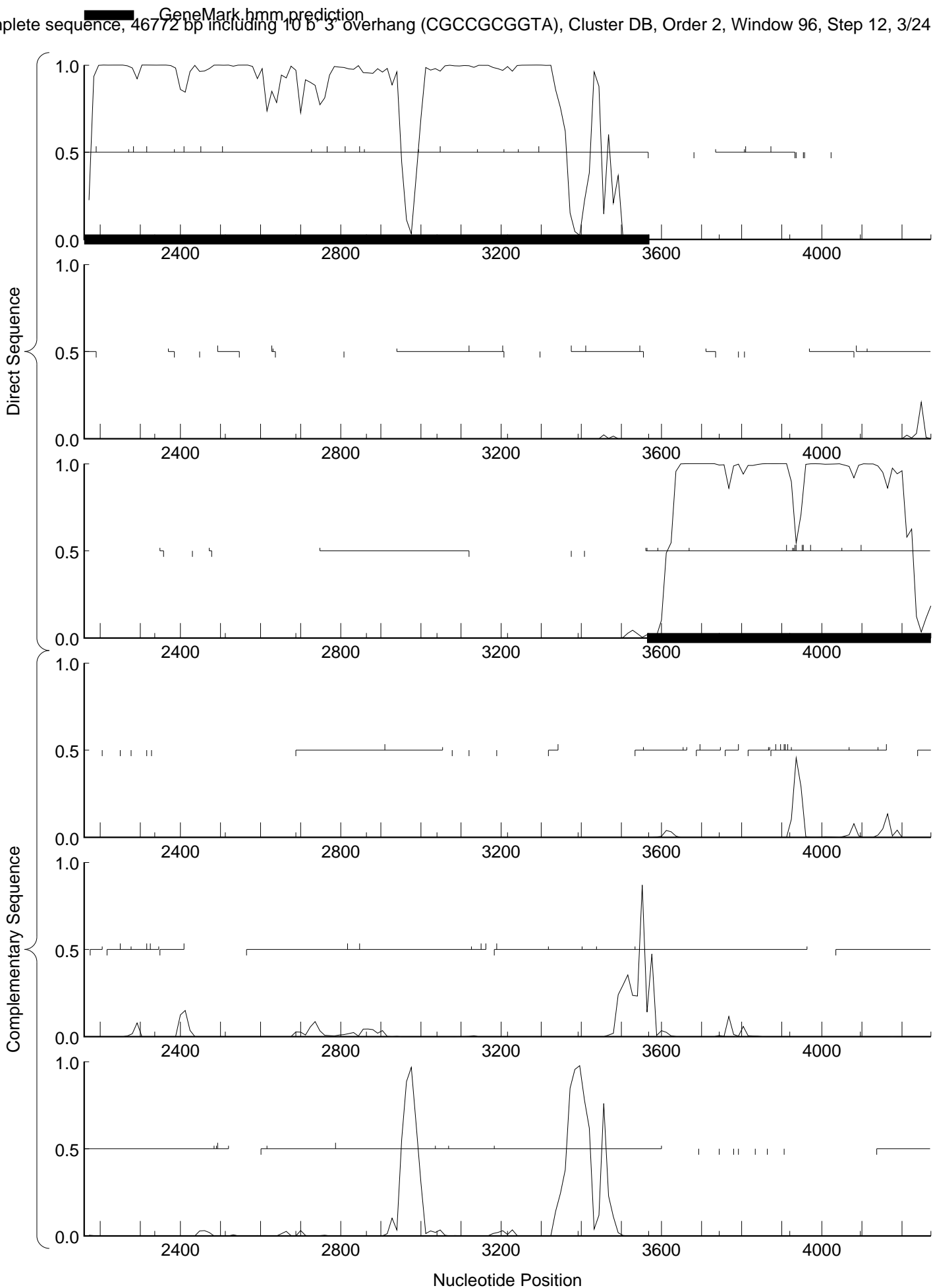
For details on the model building procedure see:
Besemer J. and Borodovsky M.
"Heuristic approach to deriving models for gene finding"
NAR, 1999, Vol. 27, No. 19, pp. 3911-3920

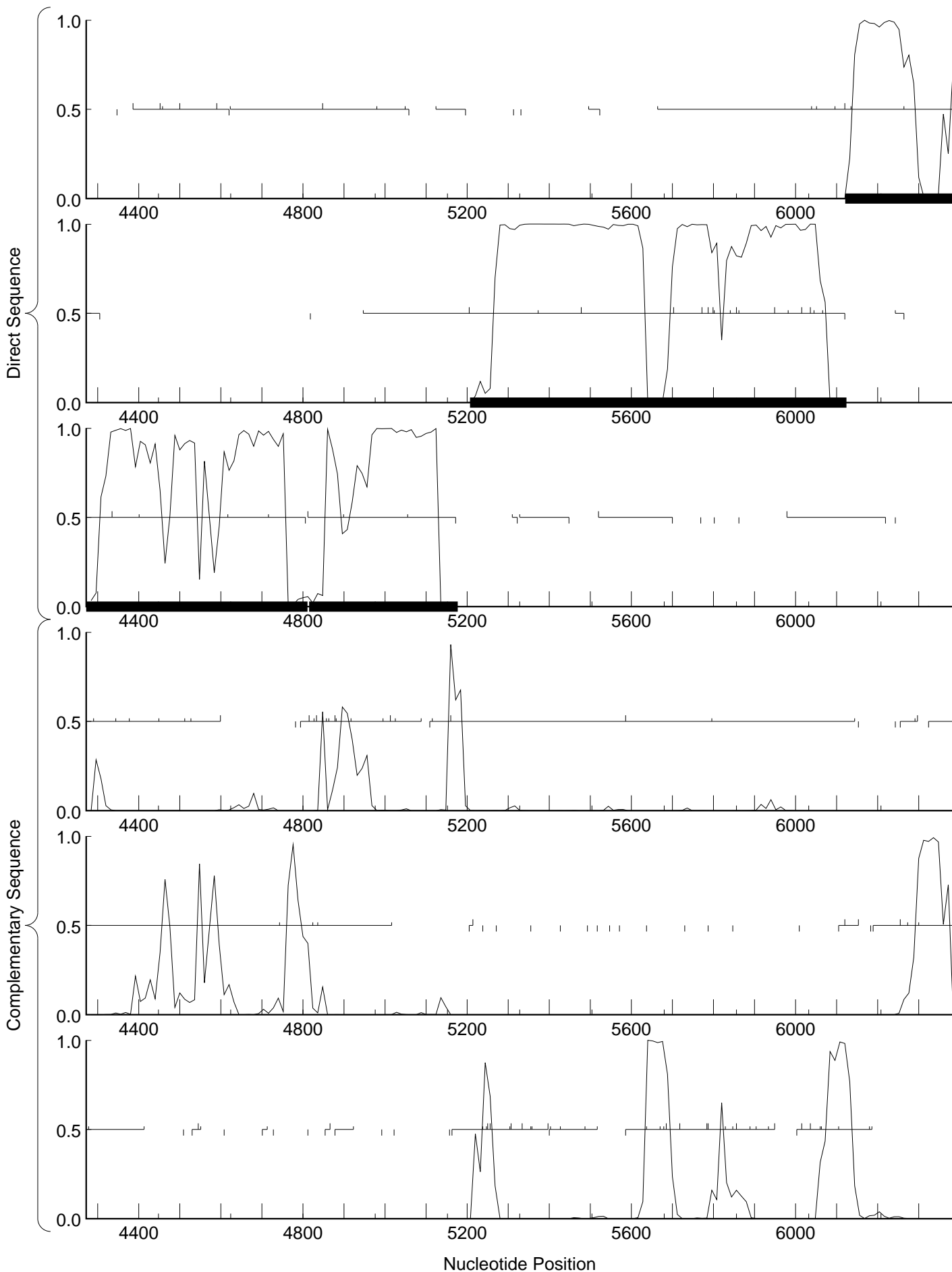
MATRIX INFORMATION

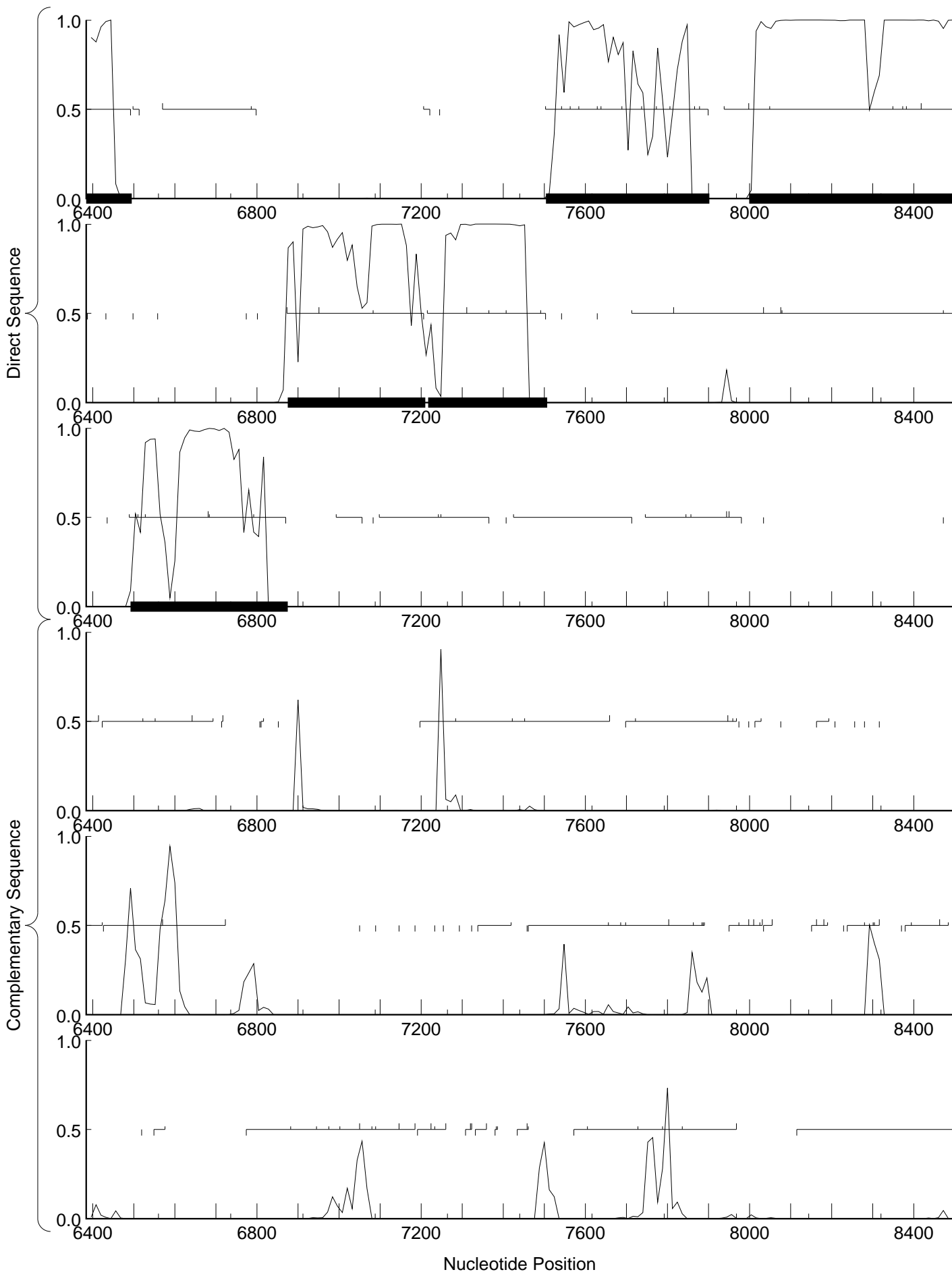
Matrix : Heuristic, GC = 67
Author : Borodovsky Laboratory - Georgia Tech
Order : 2

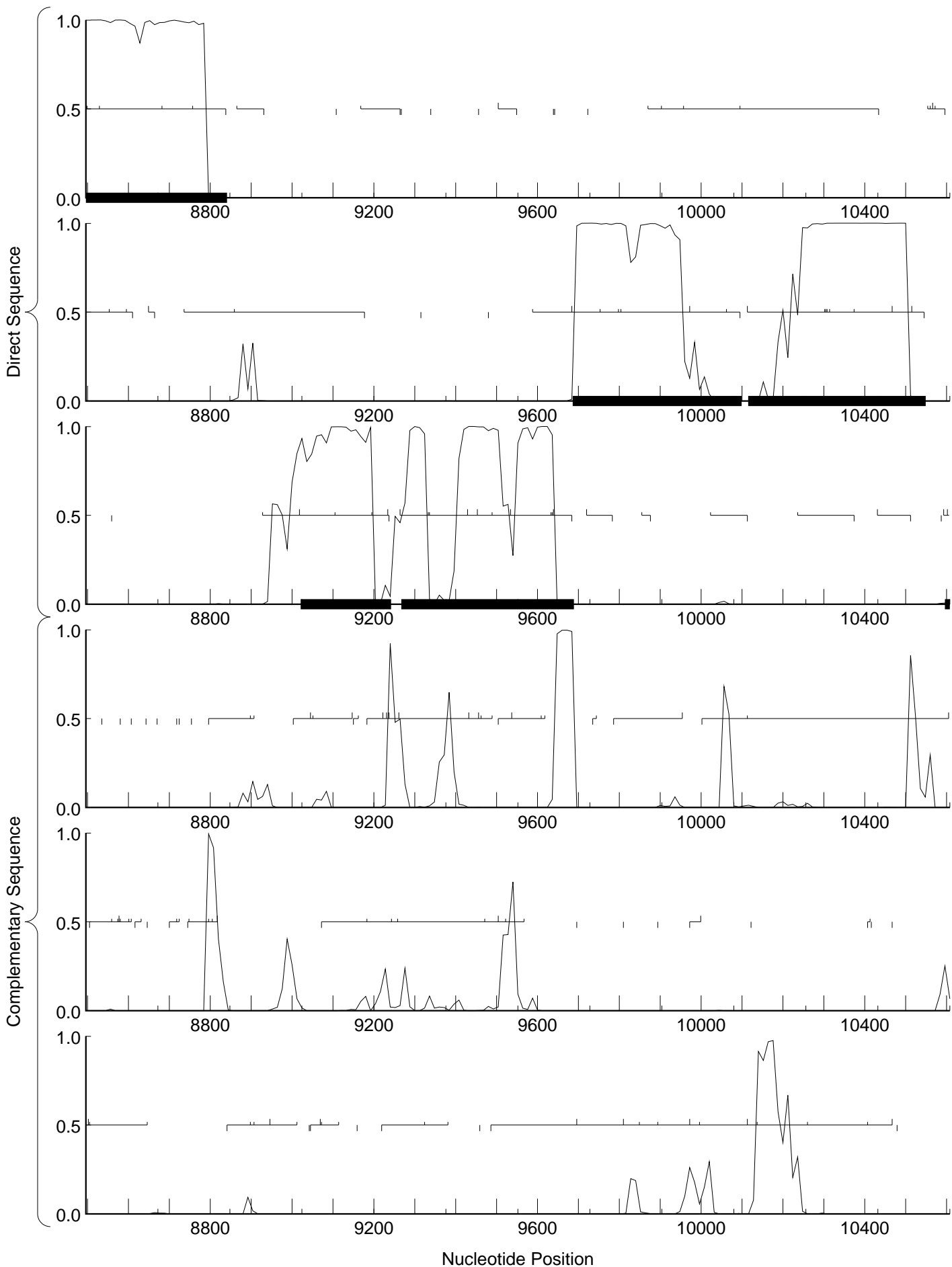
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

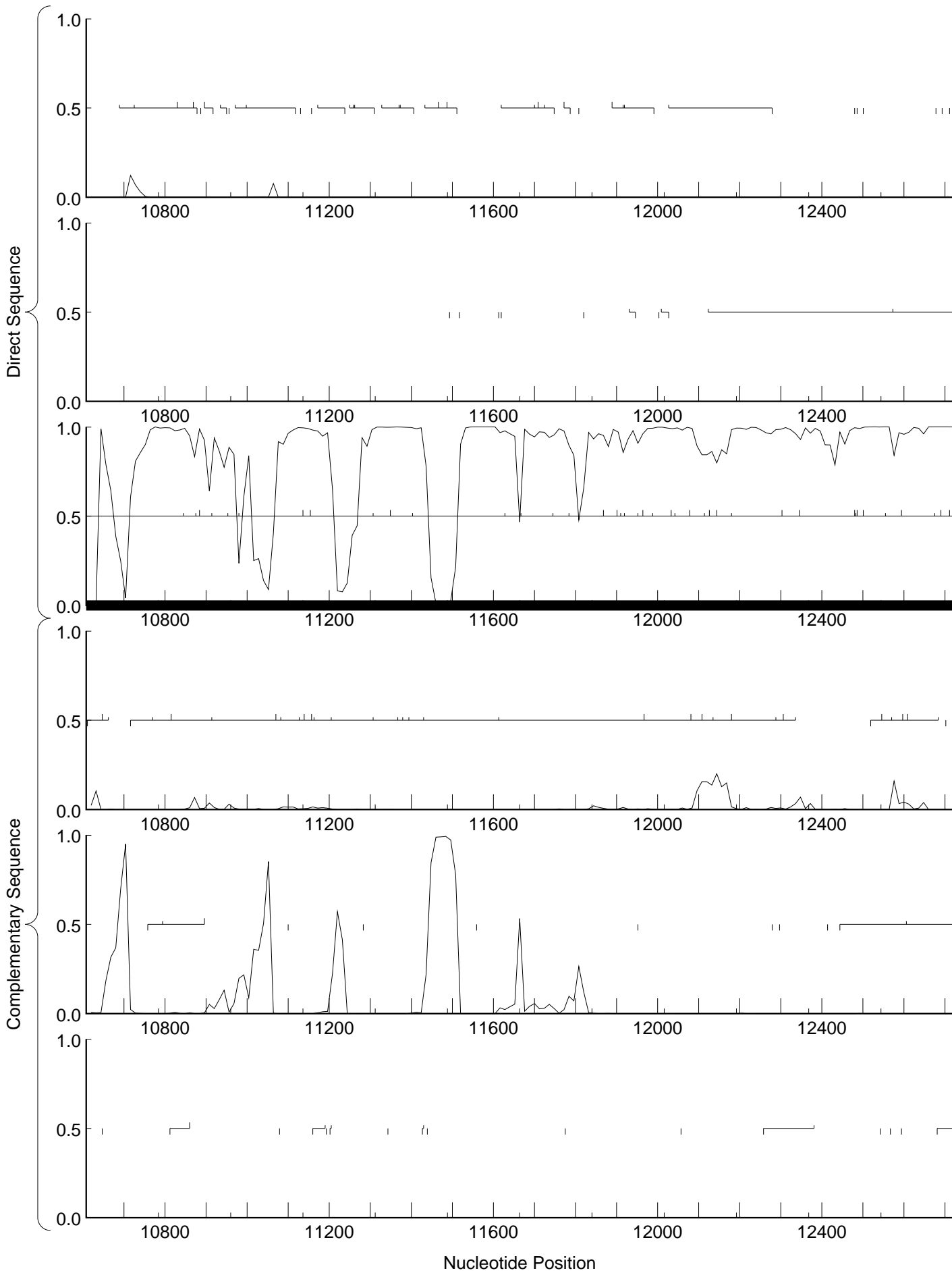


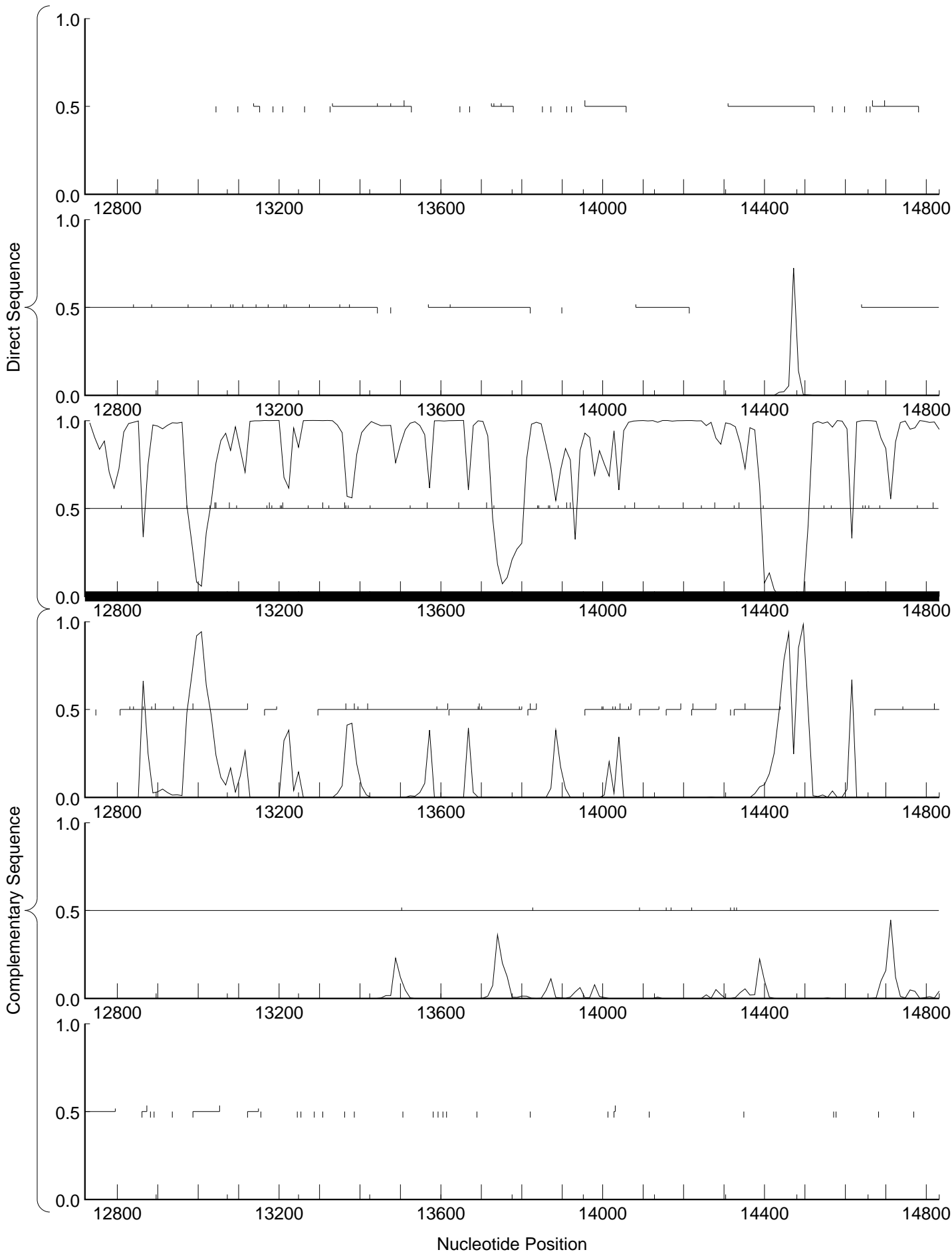




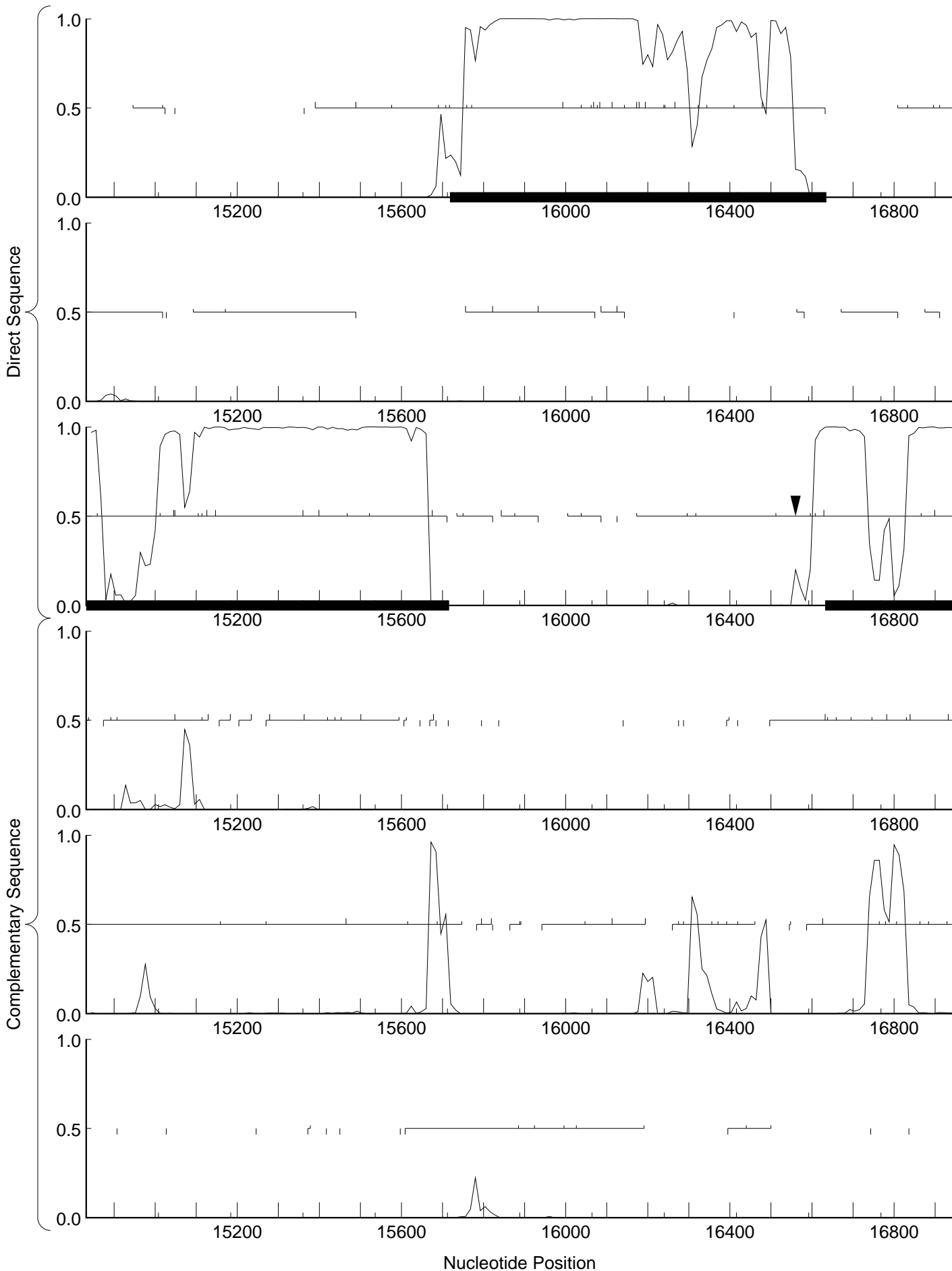


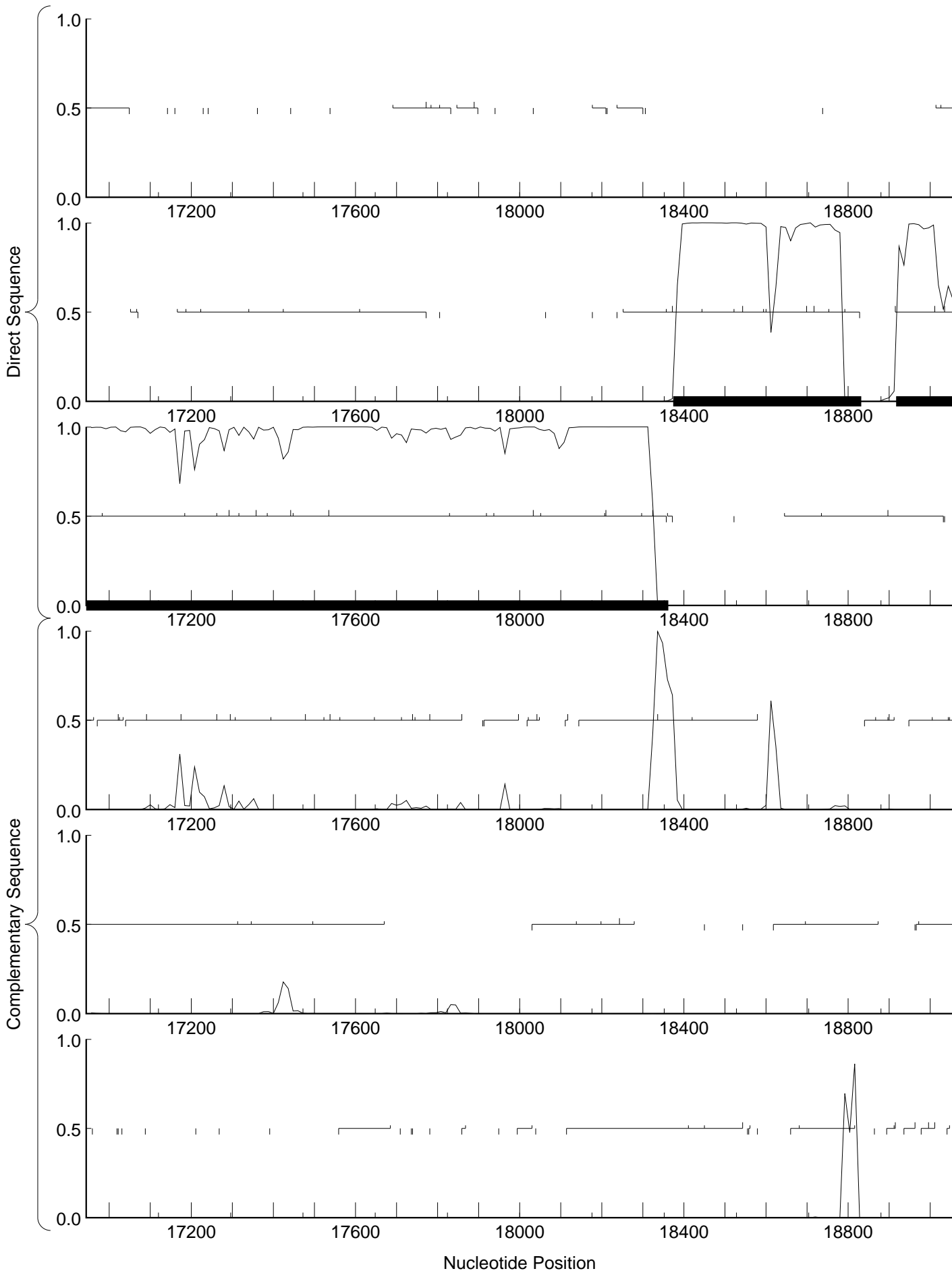


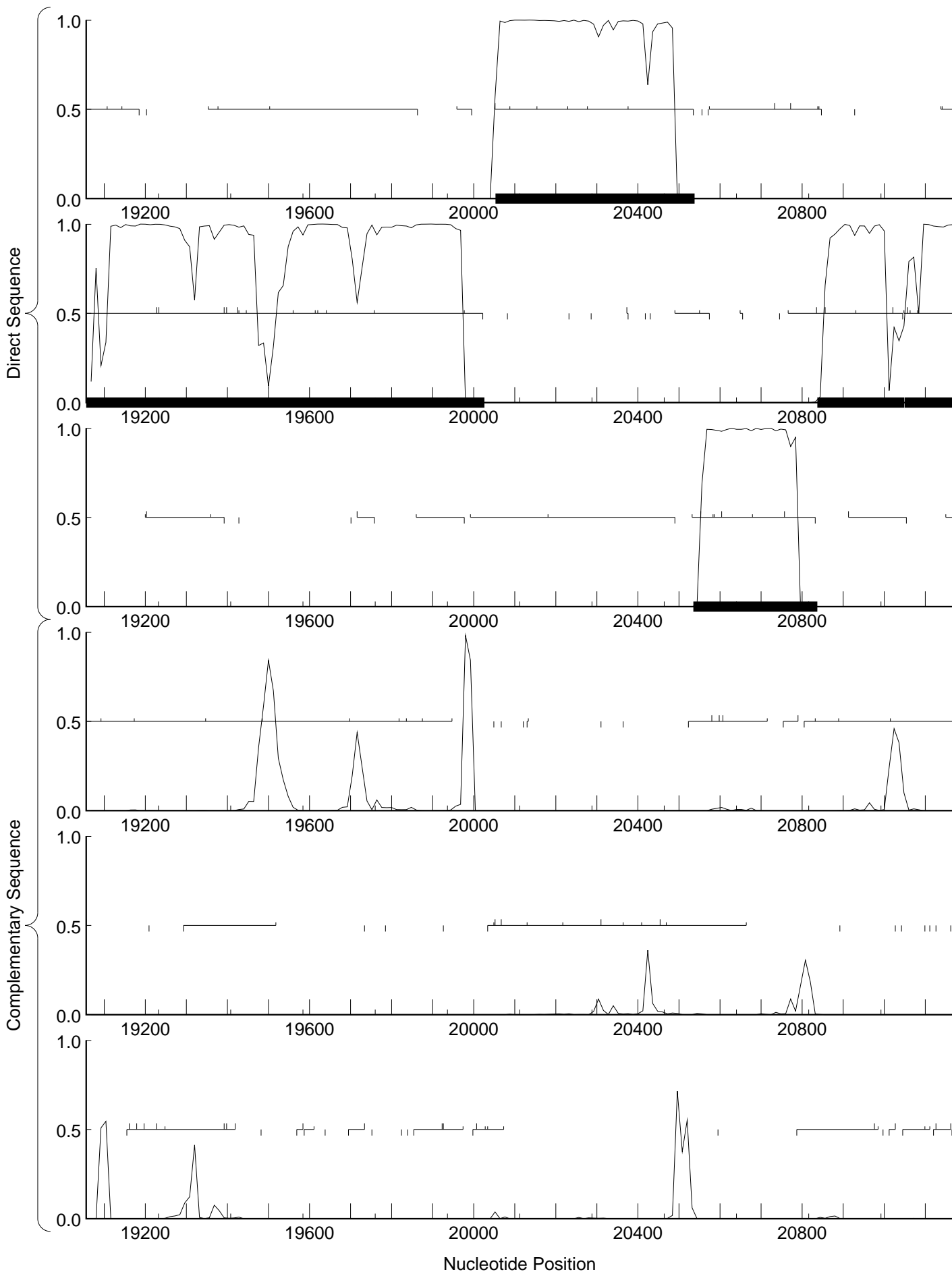


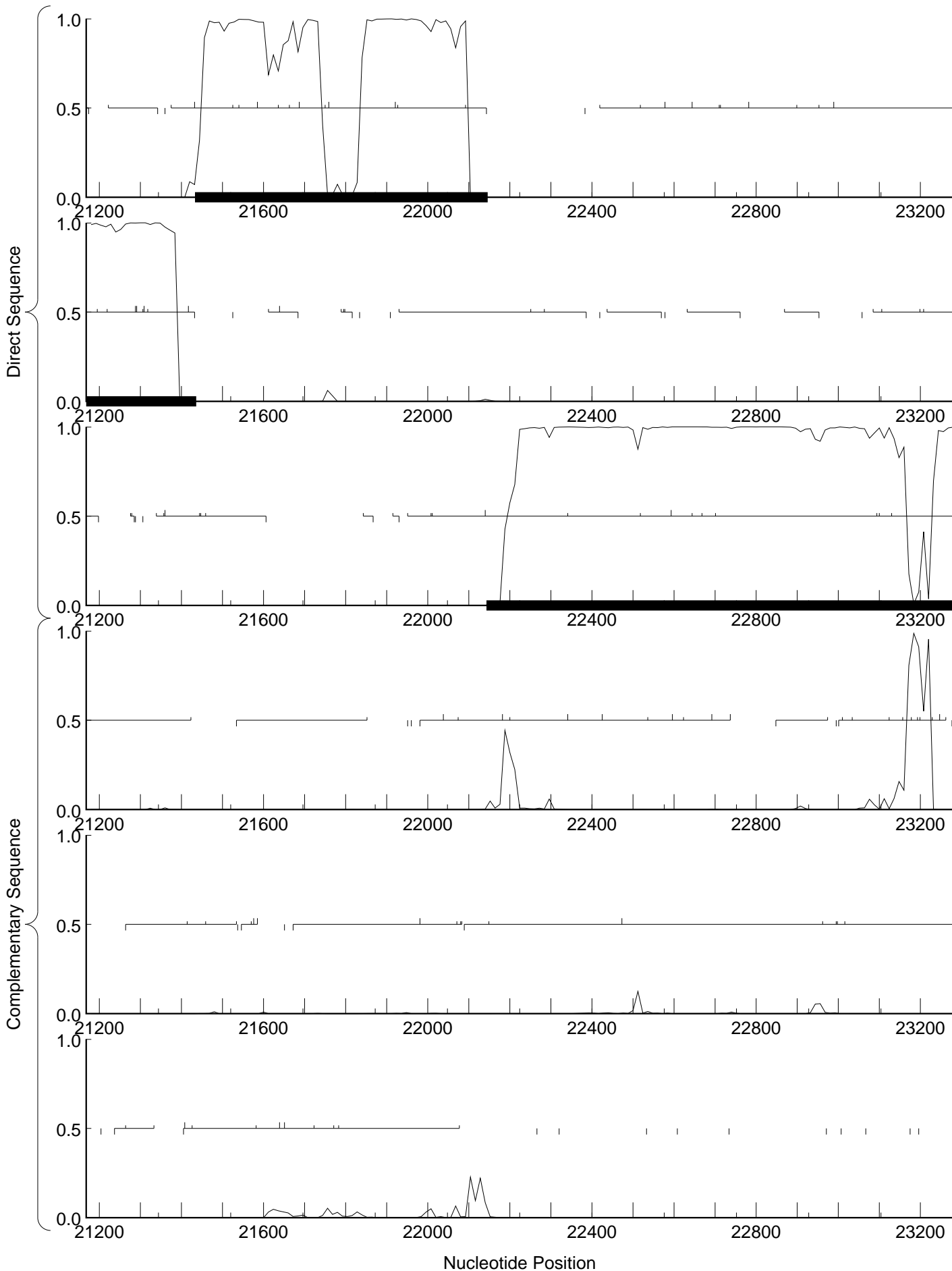


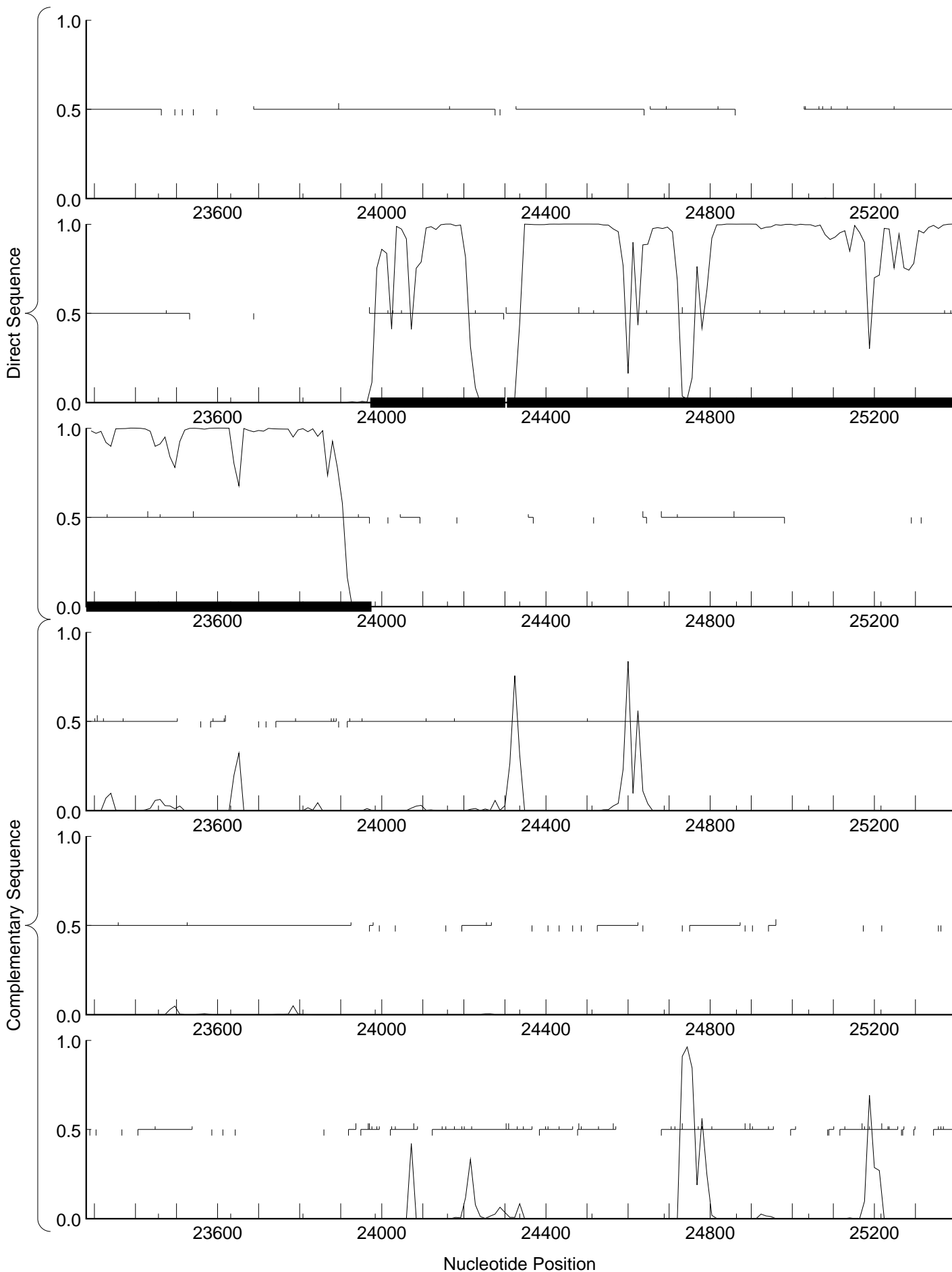
GeneMark.hmm.prediction

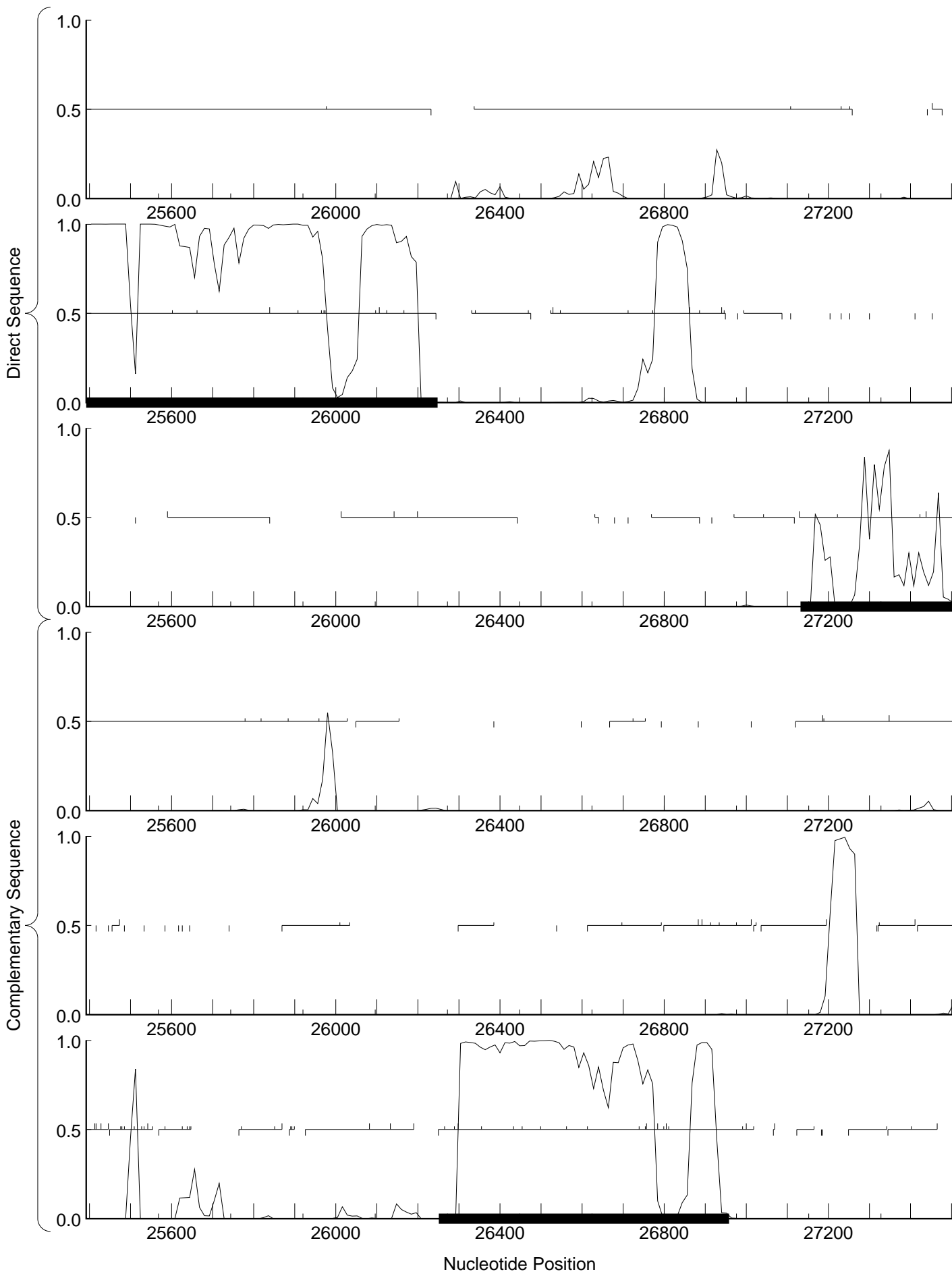


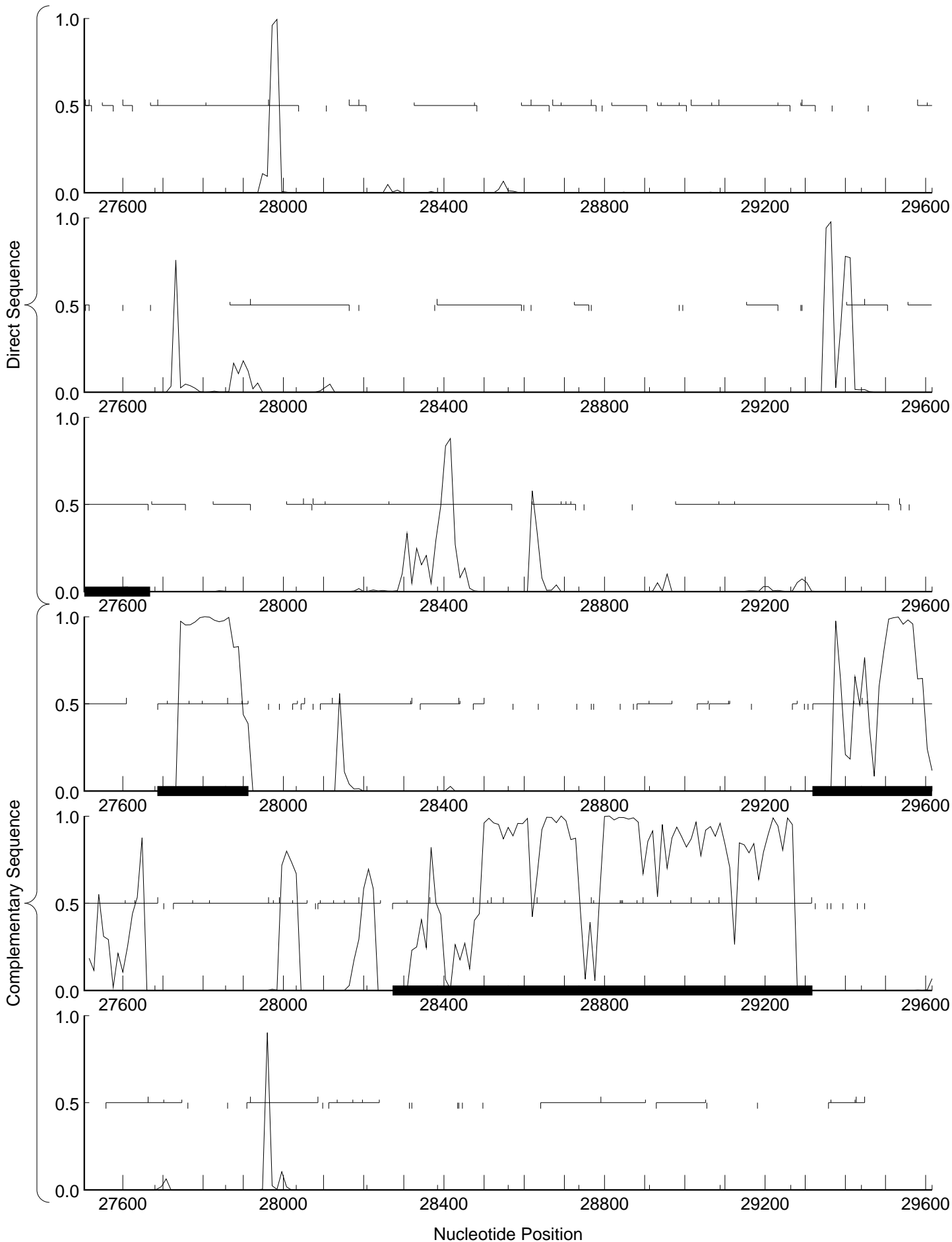


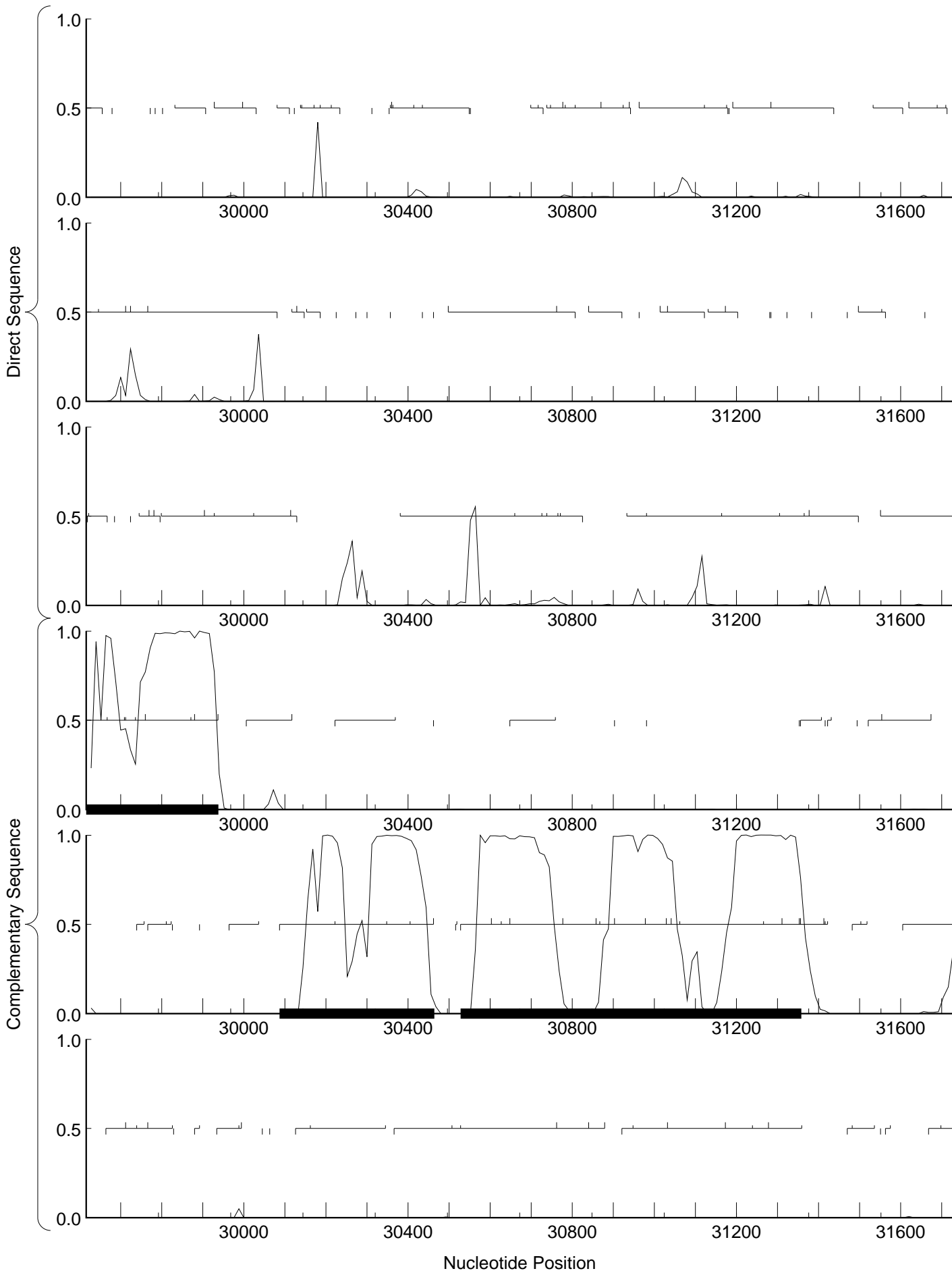


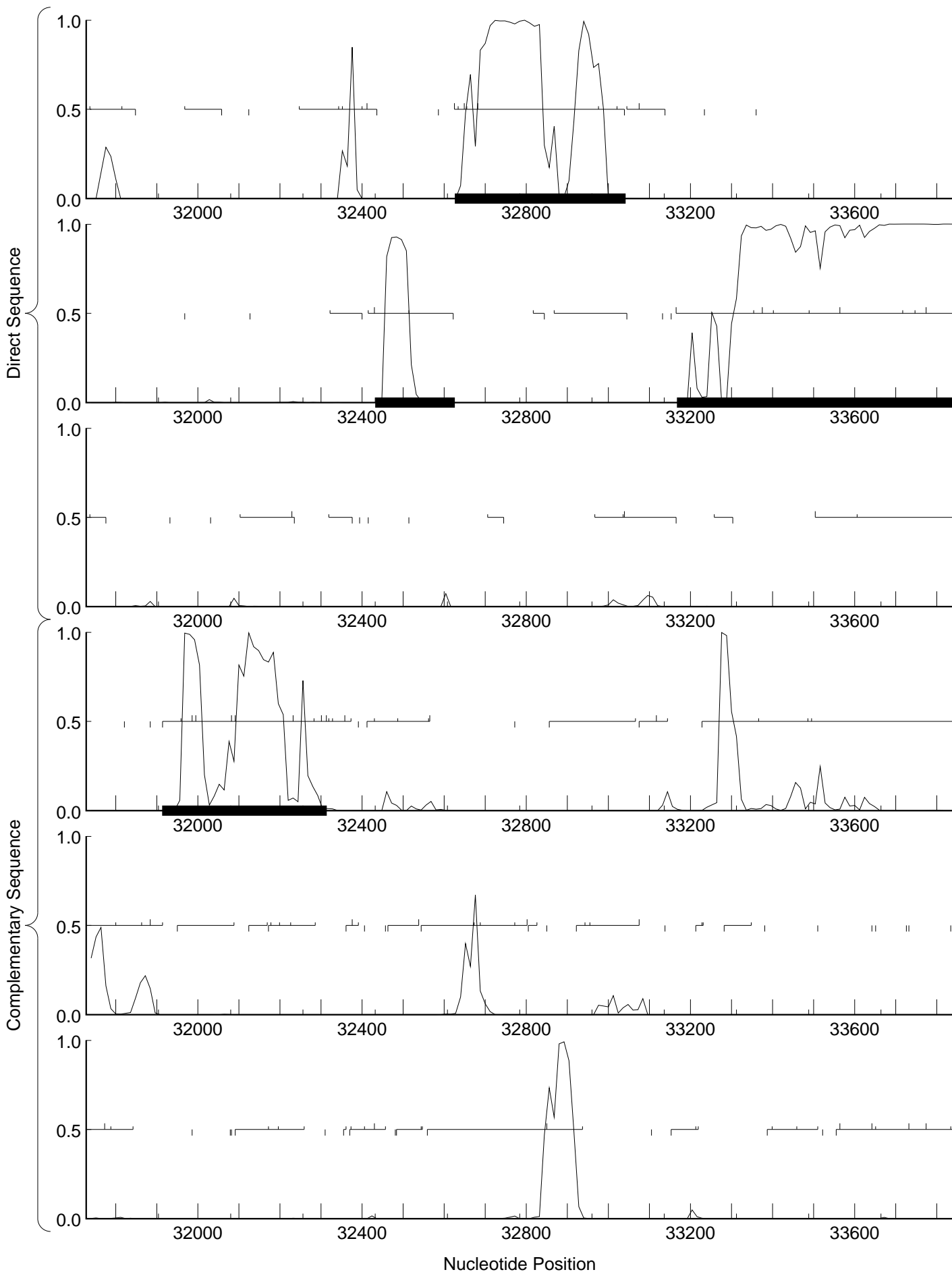


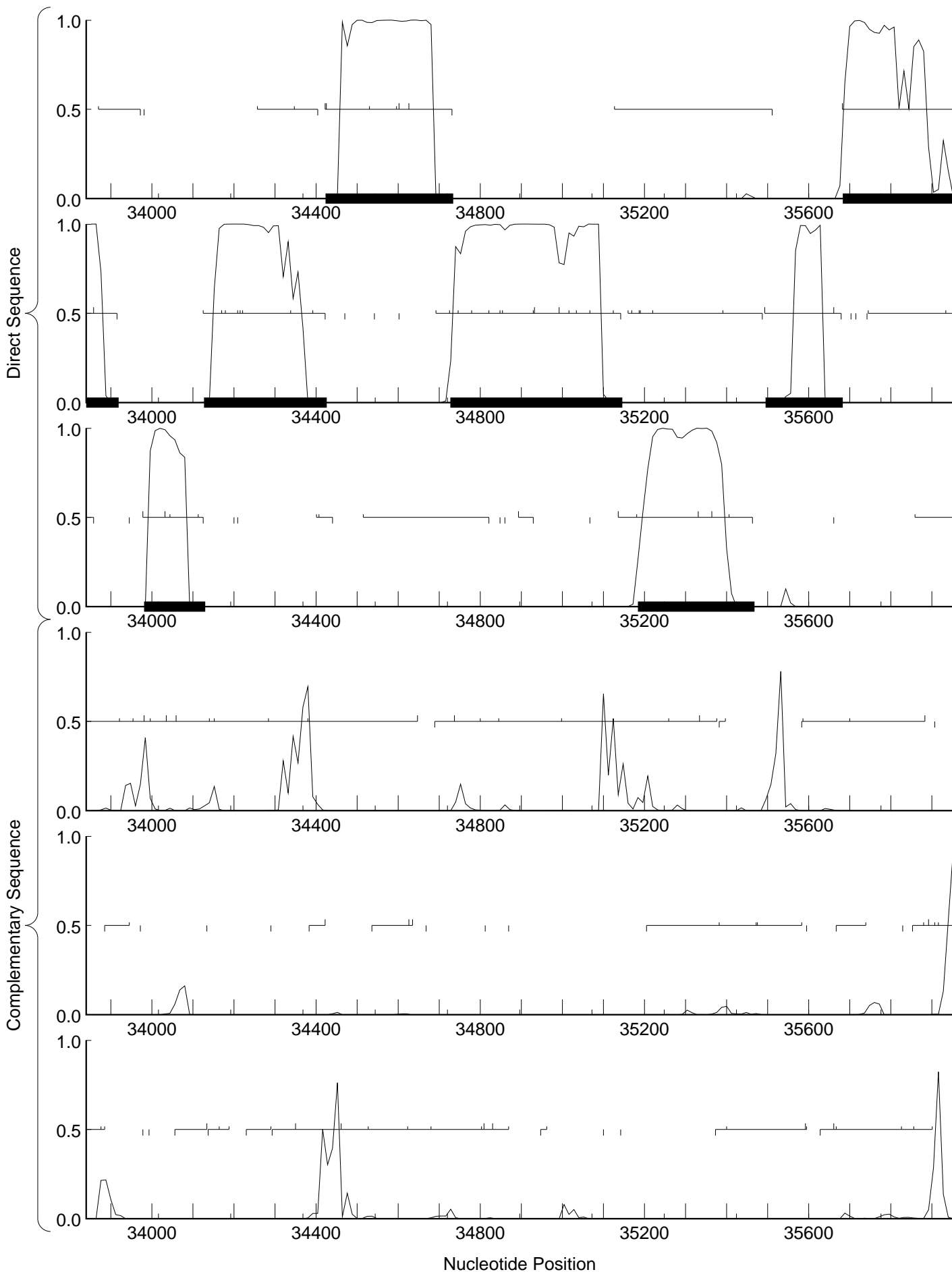


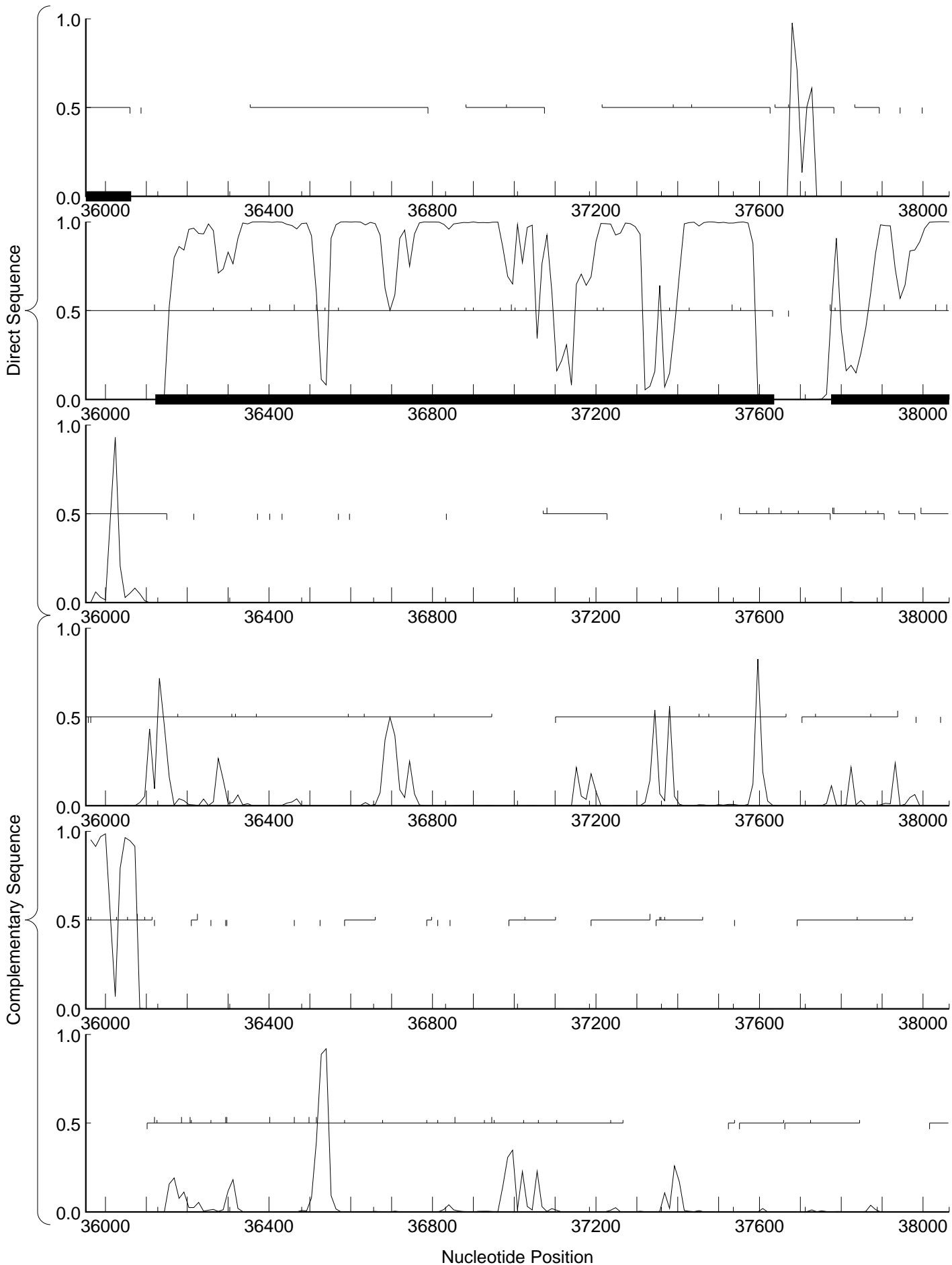


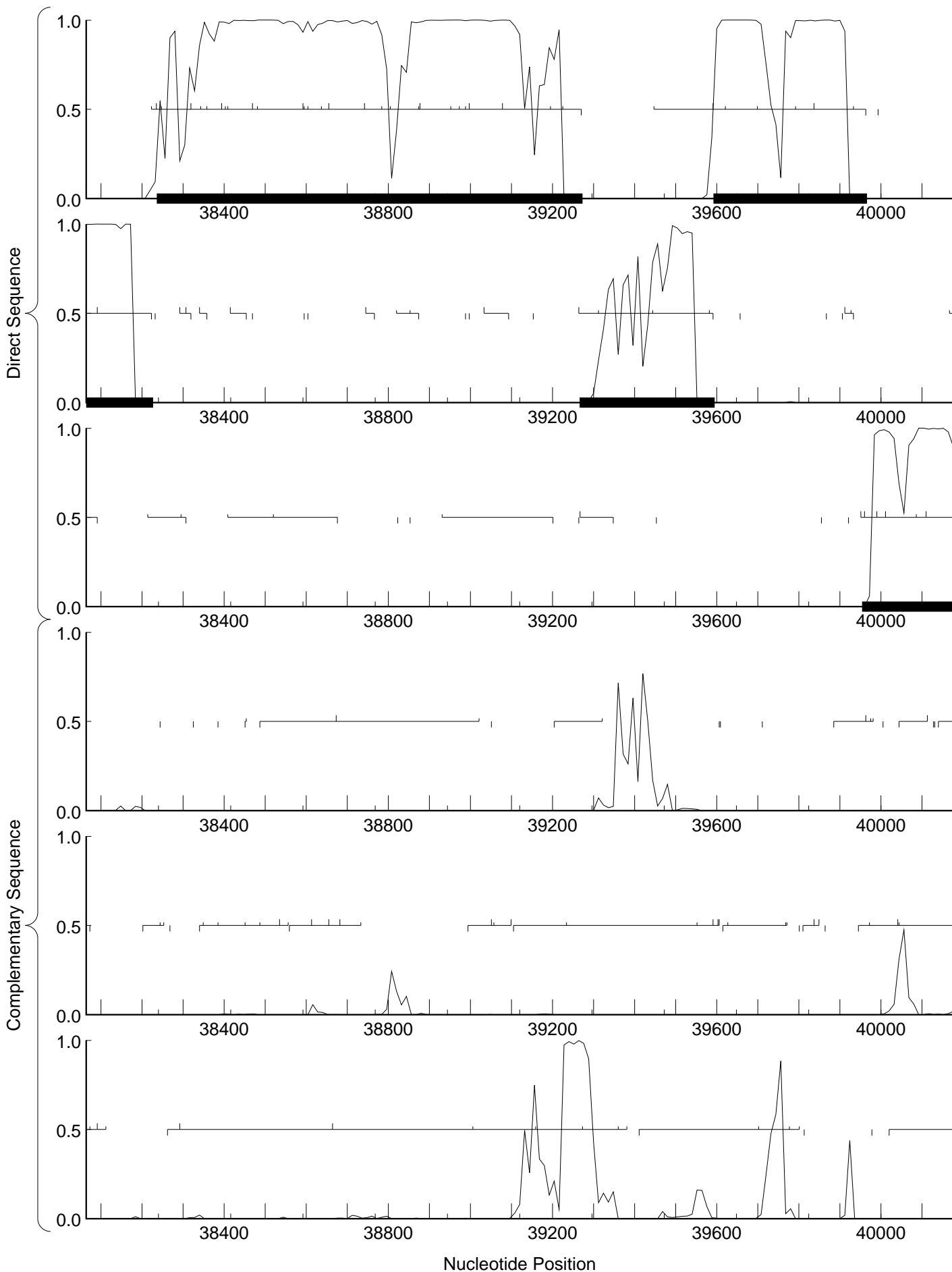


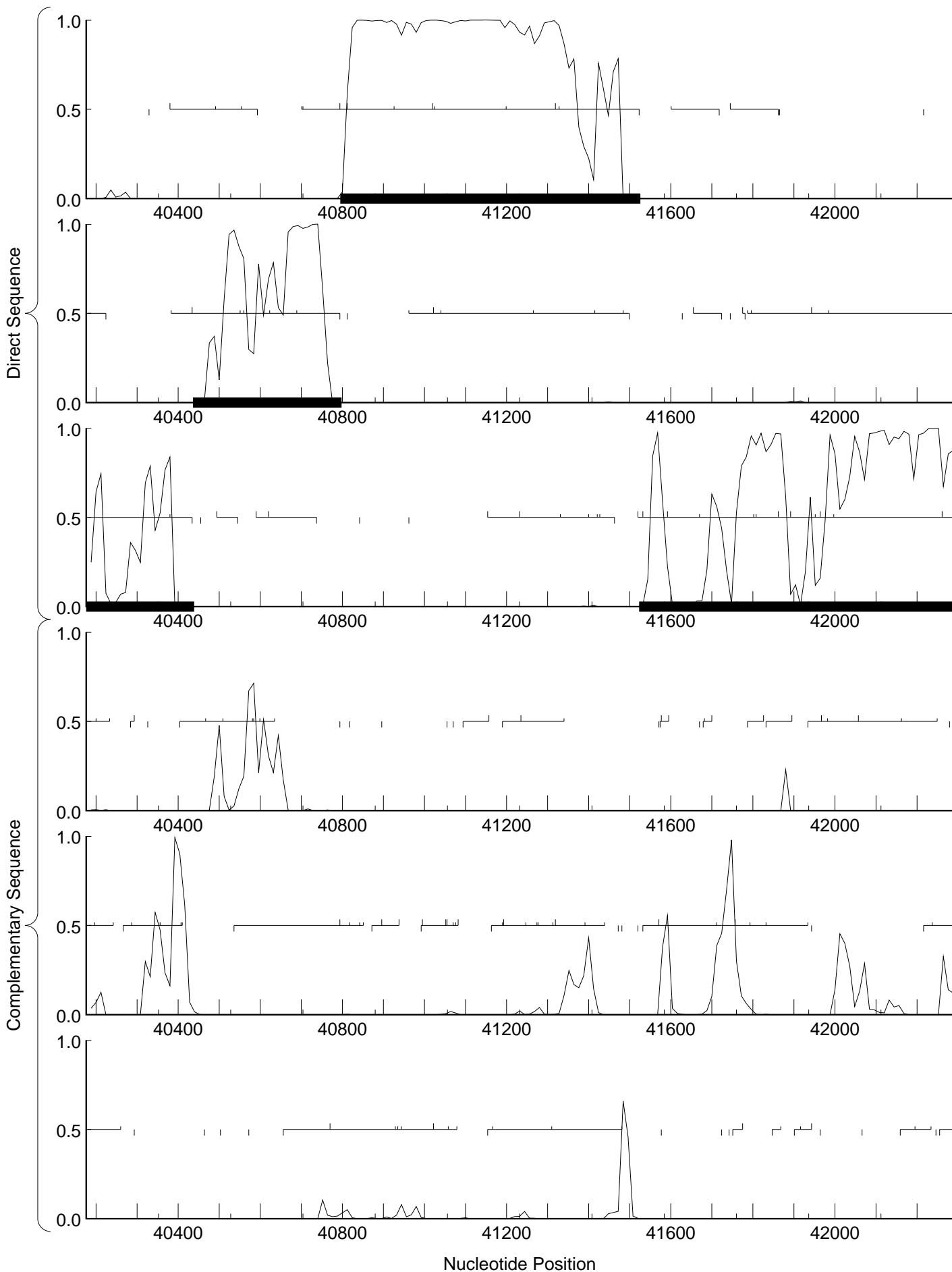


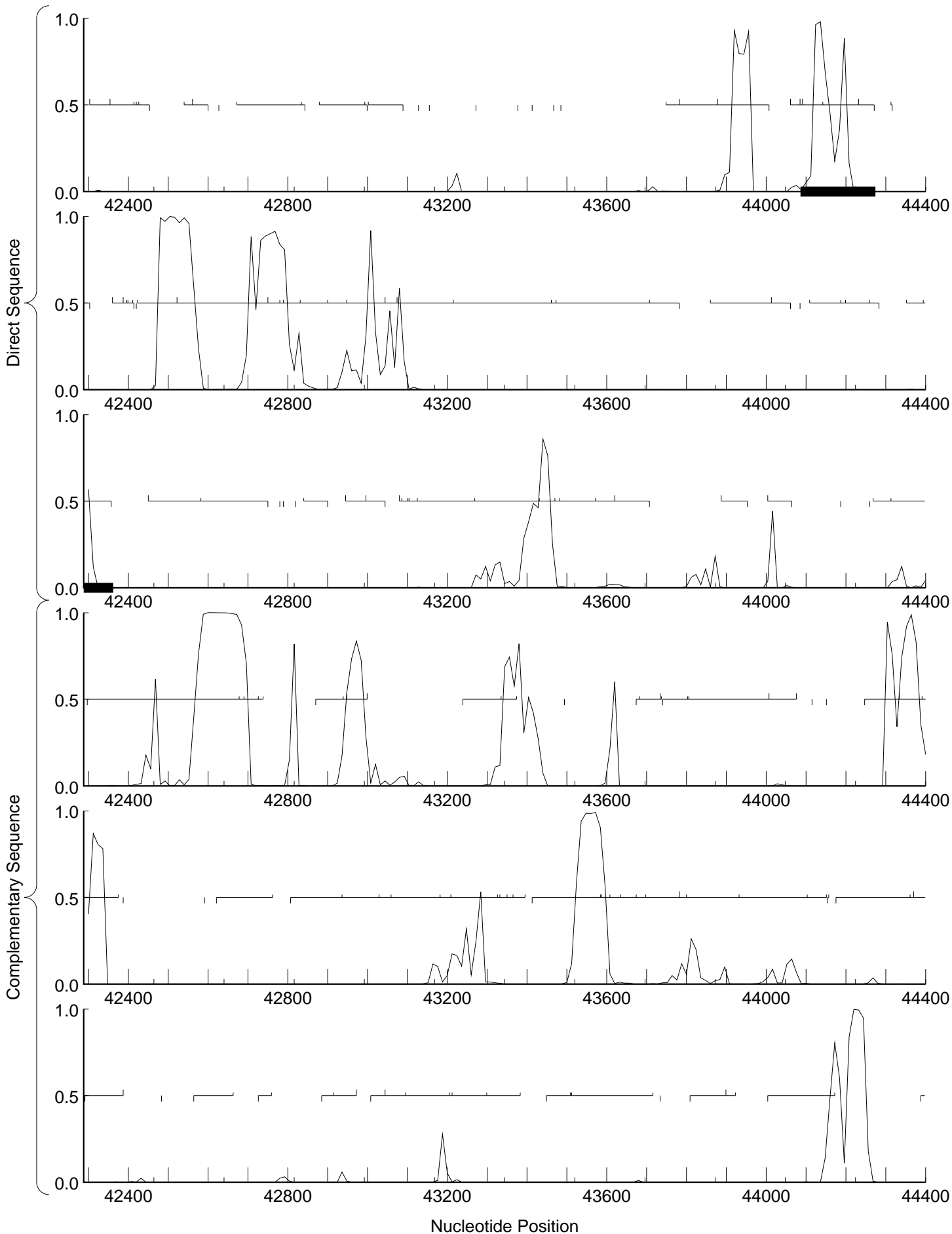












Genes complete sequence, 46772 bp including 10 b 3' overhang (CGCCGCGGTA), Cluster DB, Order 2, Window 96, Step 12, 23/24

