

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Clark complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA
Analysis Date : 3/17/18 at 16:22:34
Pages : 24
Sequence Length : 47716 bp
GC Content : 66.53%

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 INFORMATION

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

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

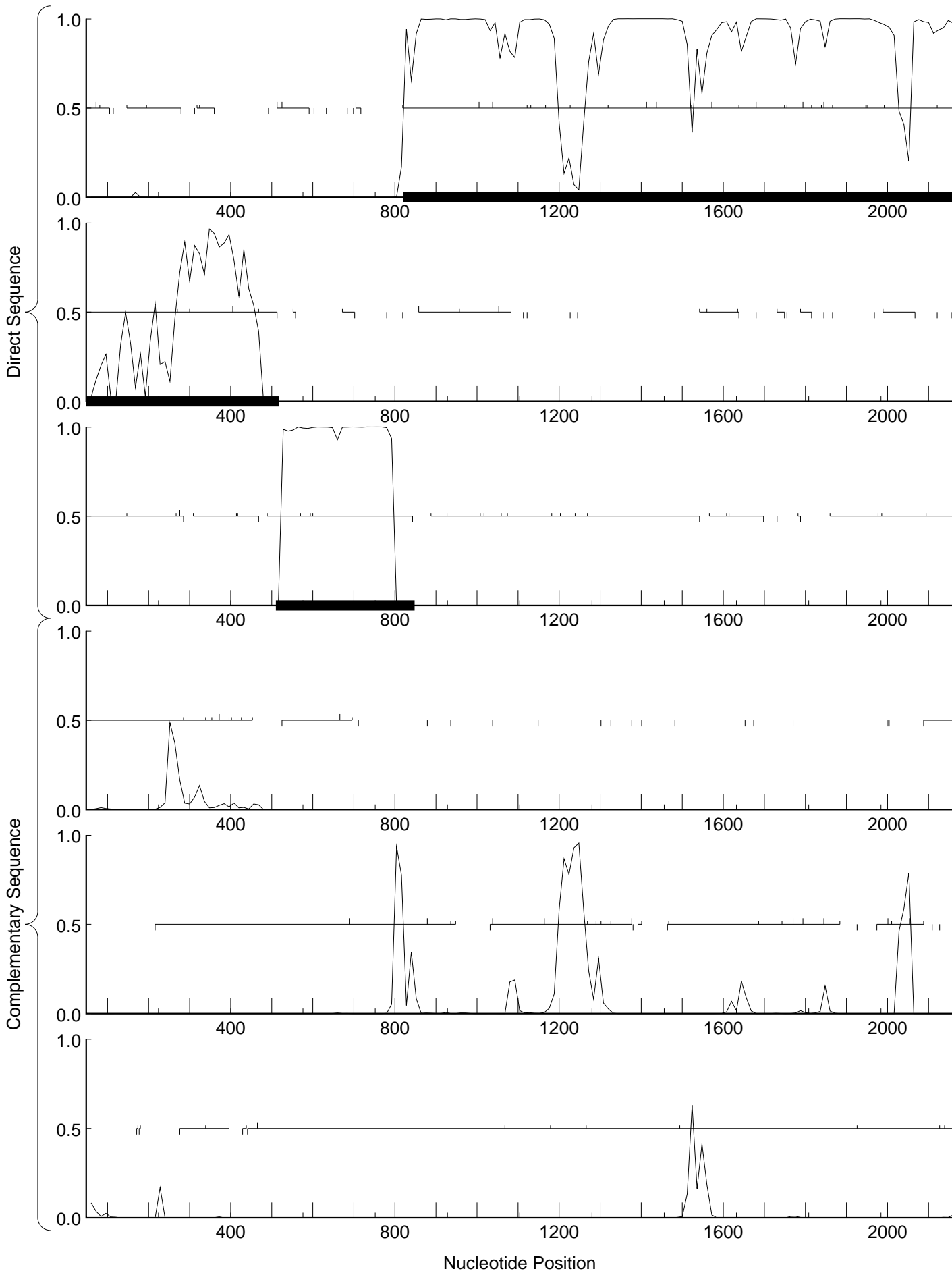
Matrix notes & comments

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

complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 2, Window 96, Step 12, 2/24

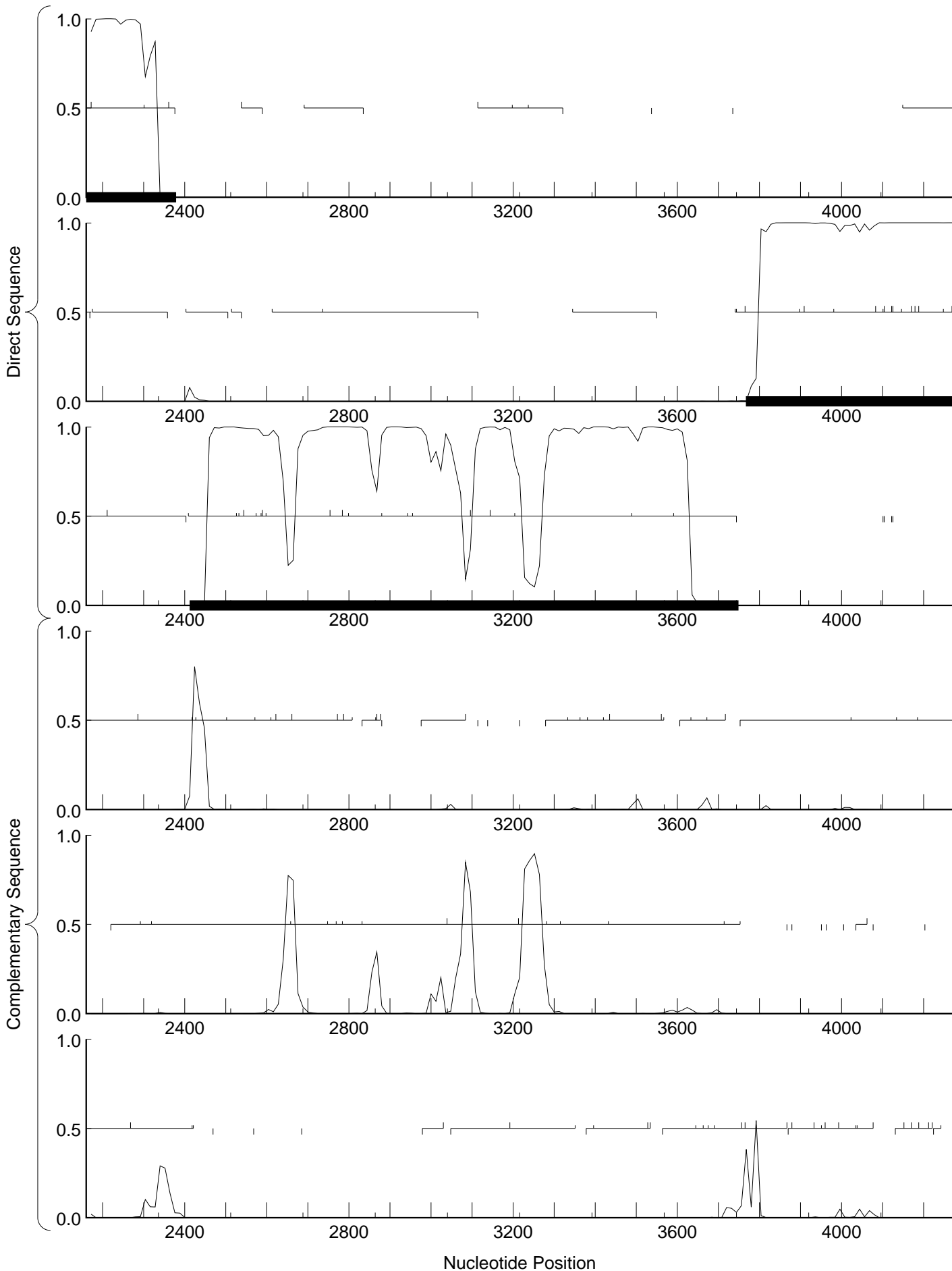
██████████

GeneMark, hmm prediction

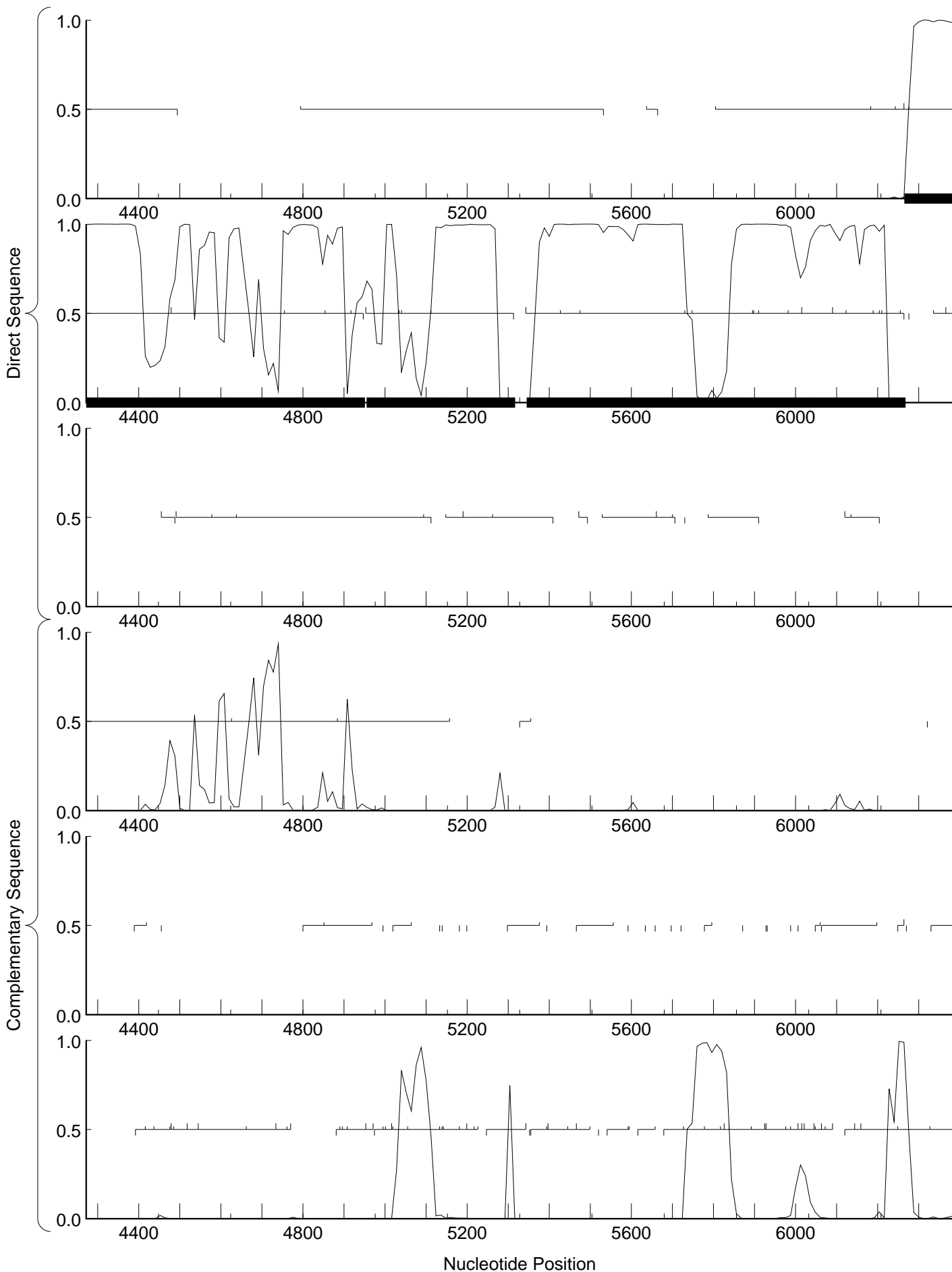


complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 2, Window 96, Step 12, 3/24

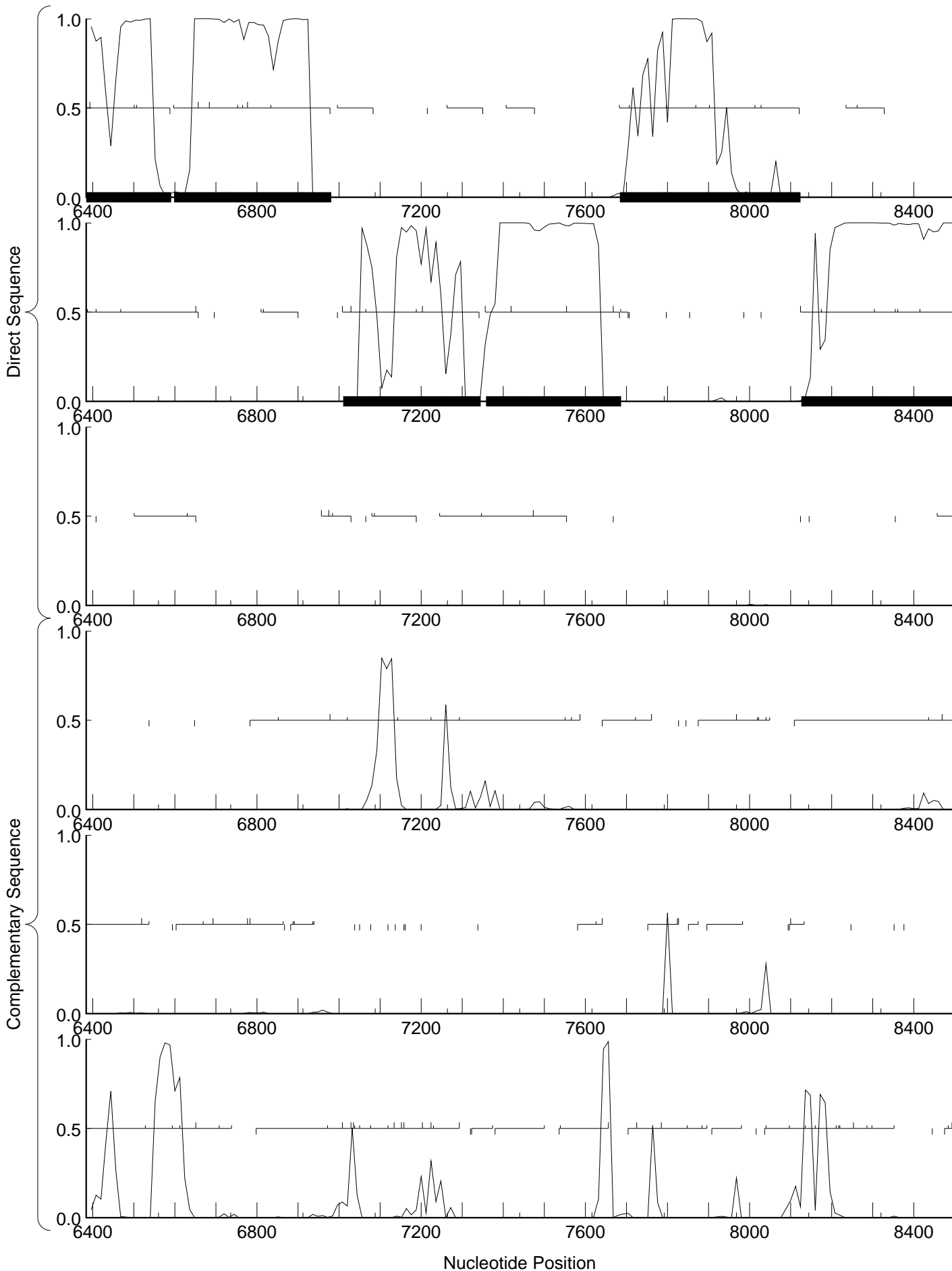
GeneMark, hmm prediction

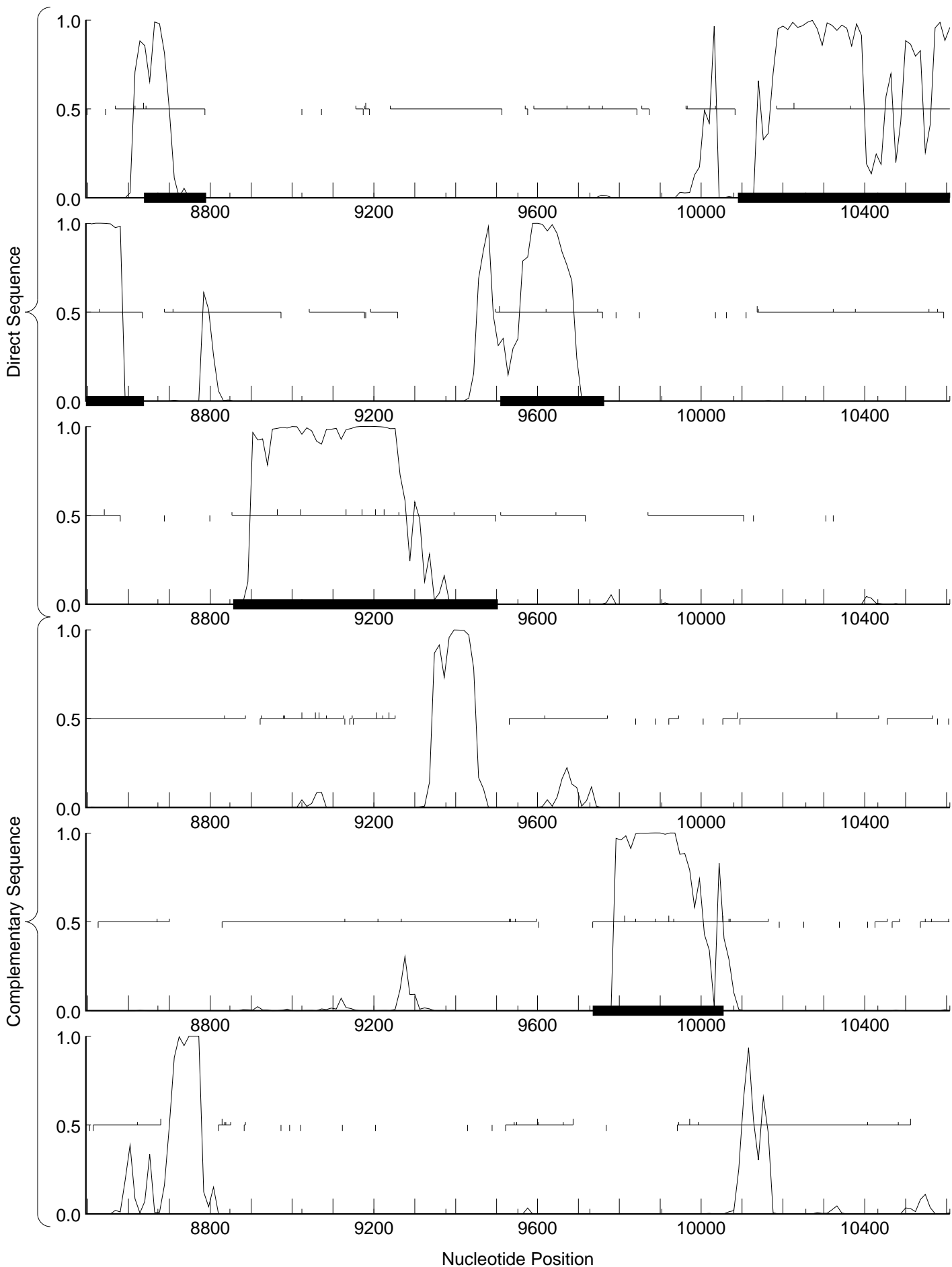


GeneMark, hmm prediction



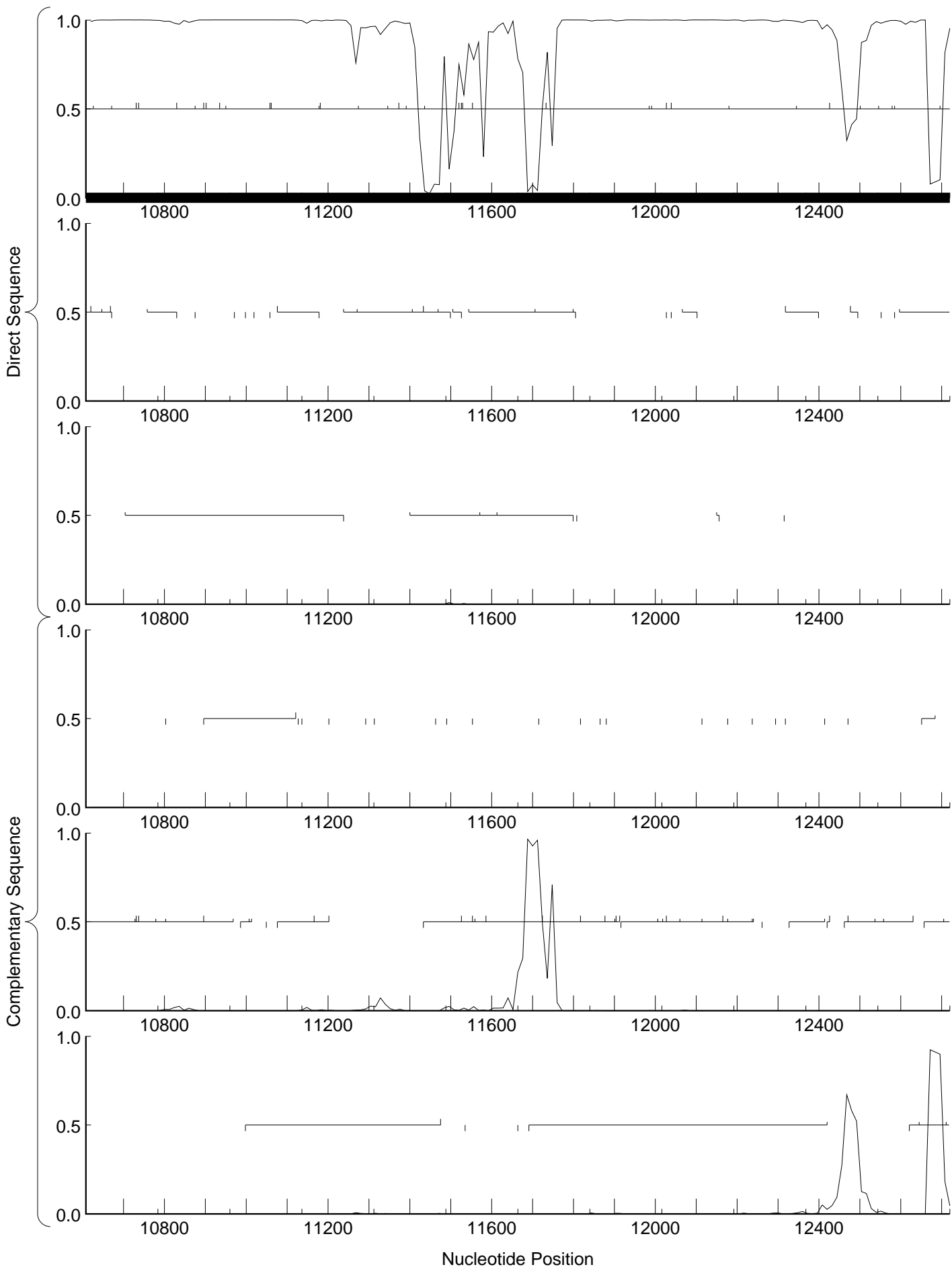
GeneMark, hmm prediction





complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 2, Window 96, Step 12, 7/24

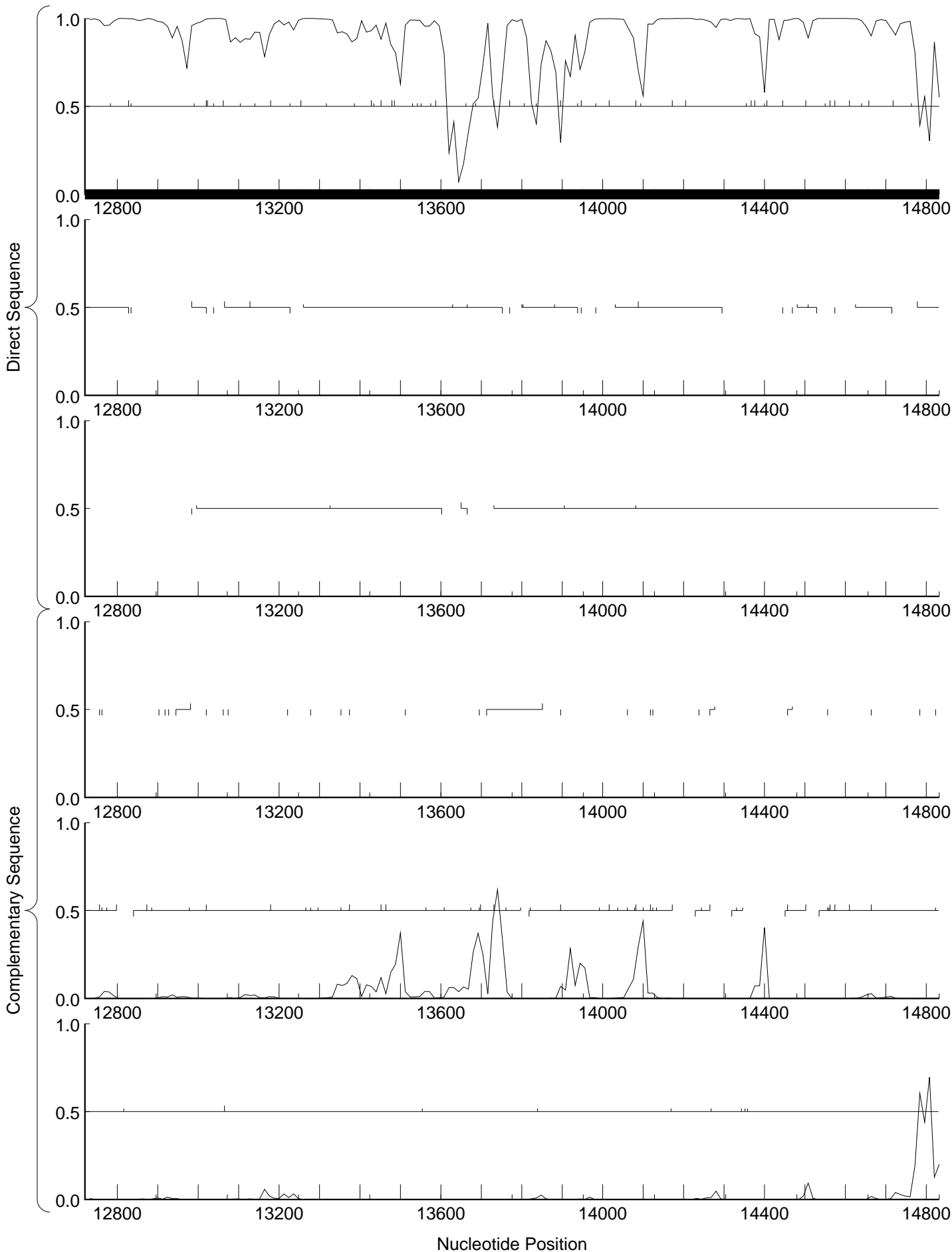
GeneMark, hmm prediction



complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 2, Window 96, Step 12, 8/24



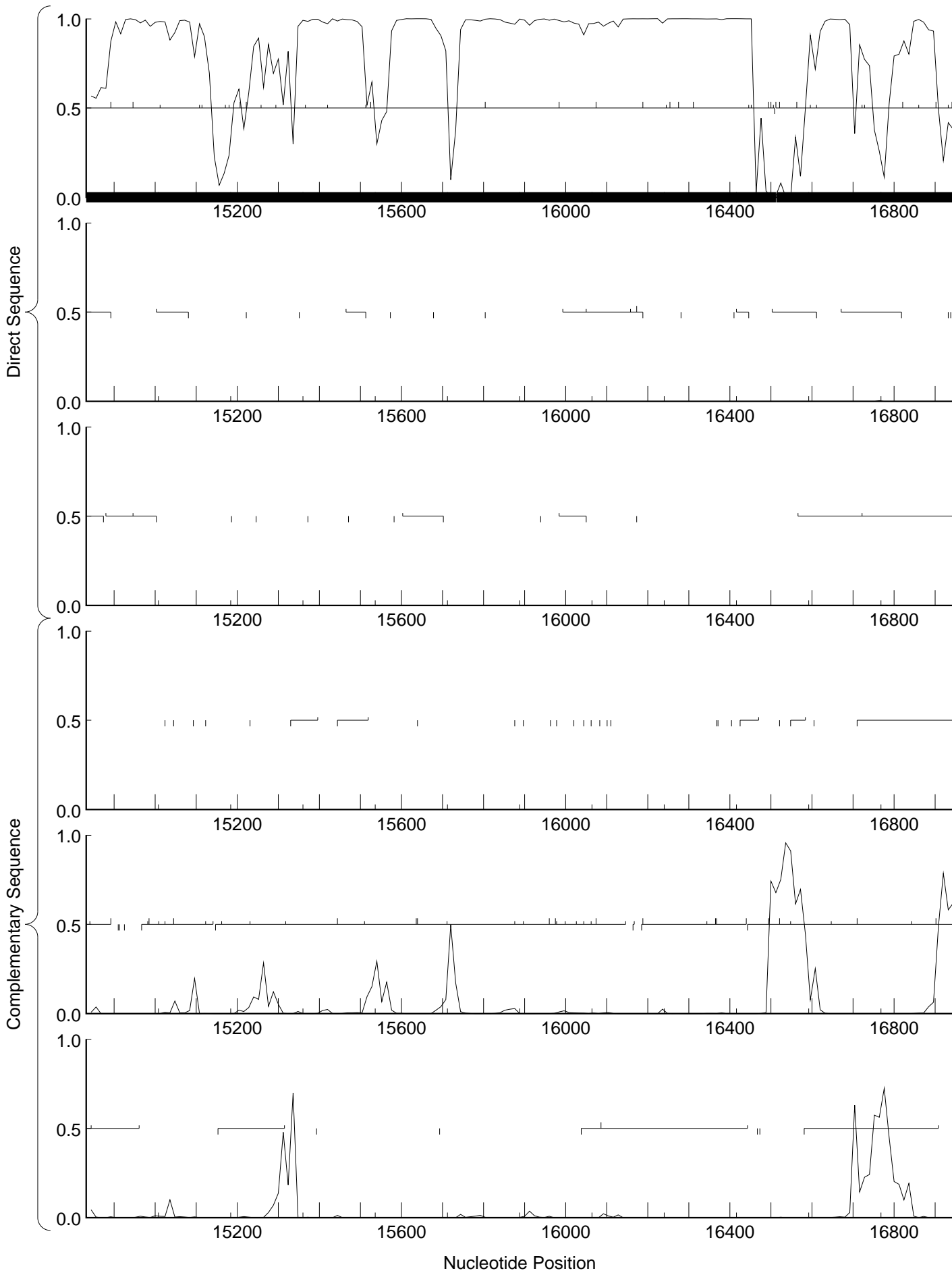
GeneMark, hmm prediction



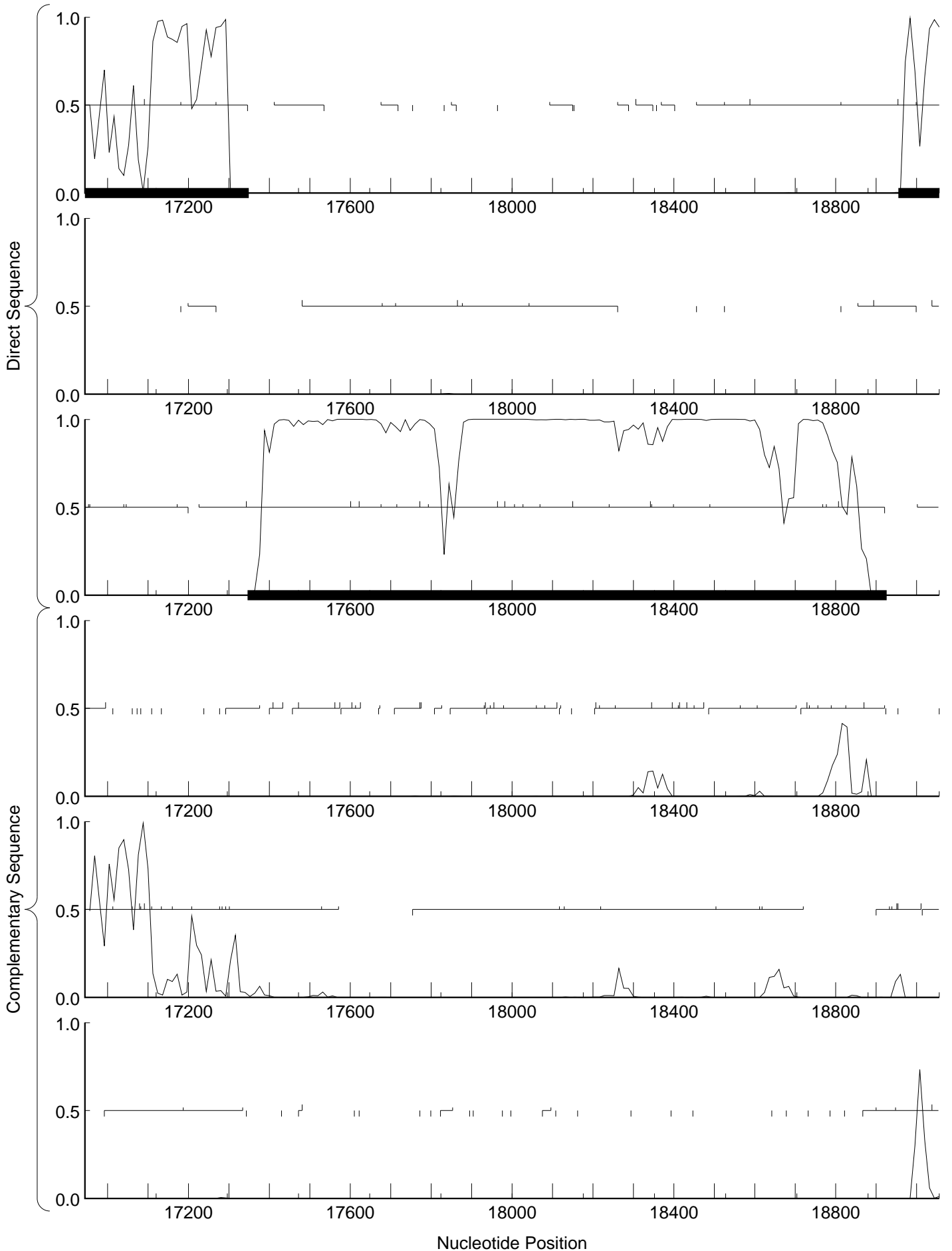
complete sequence, 47716 bp including 11-base 3' overhang (TGCCAAGGGGA), Cluster CZ4, Order 2, Window 96, Step 12, 9/24

██████████

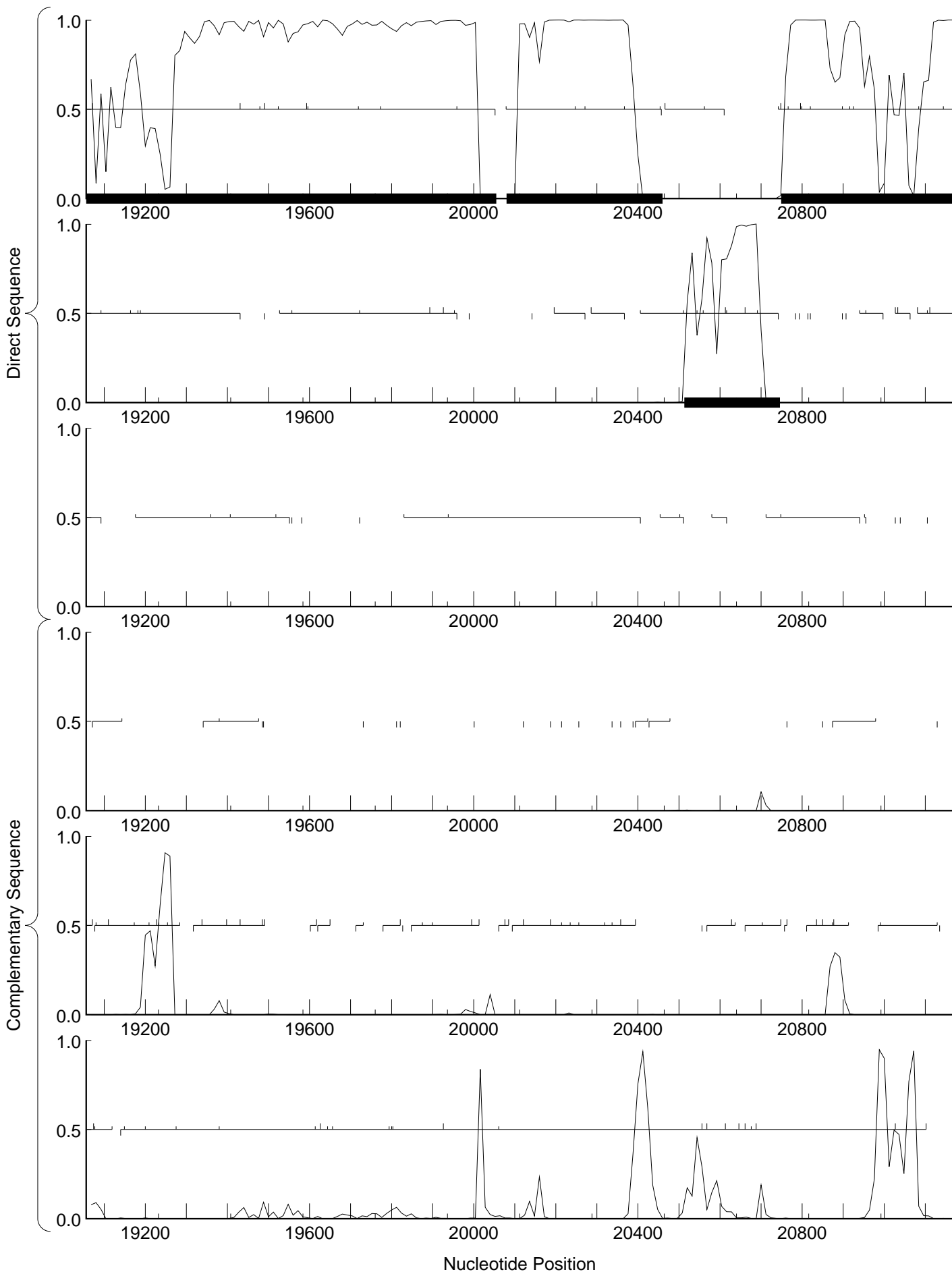
GeneMark, hmm prediction



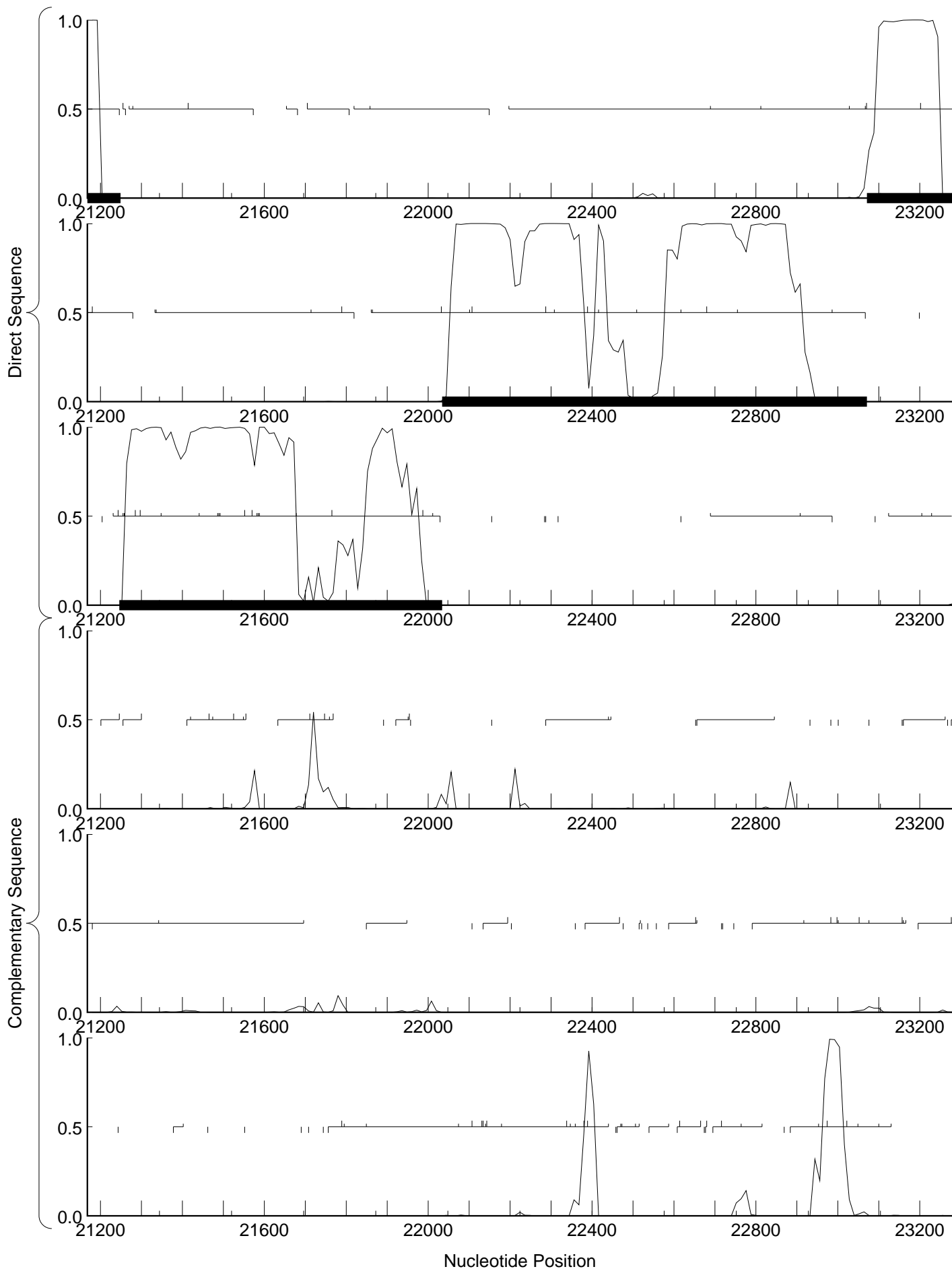
GeneMark, hmm prediction



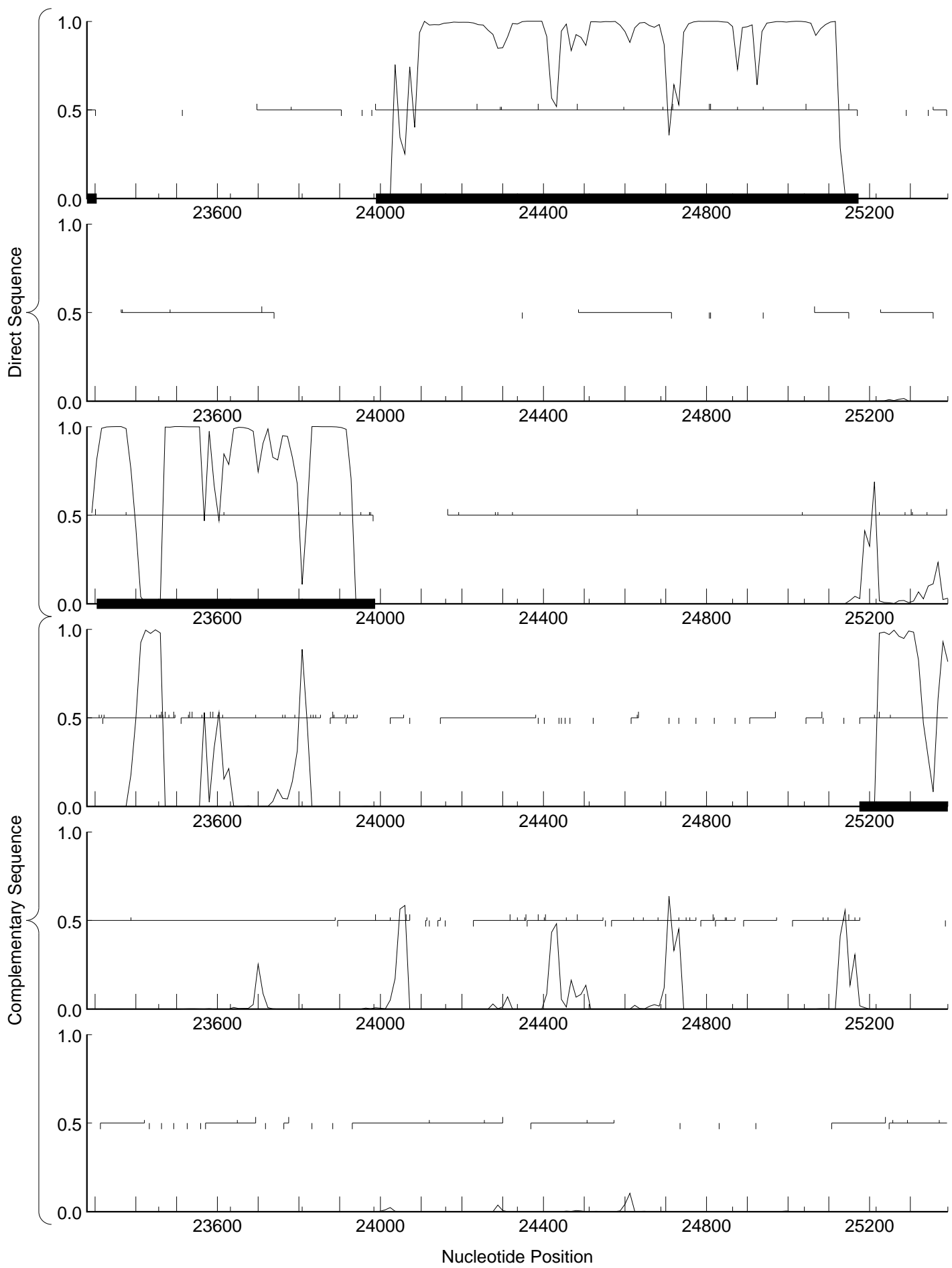
GeneMark, hmm prediction



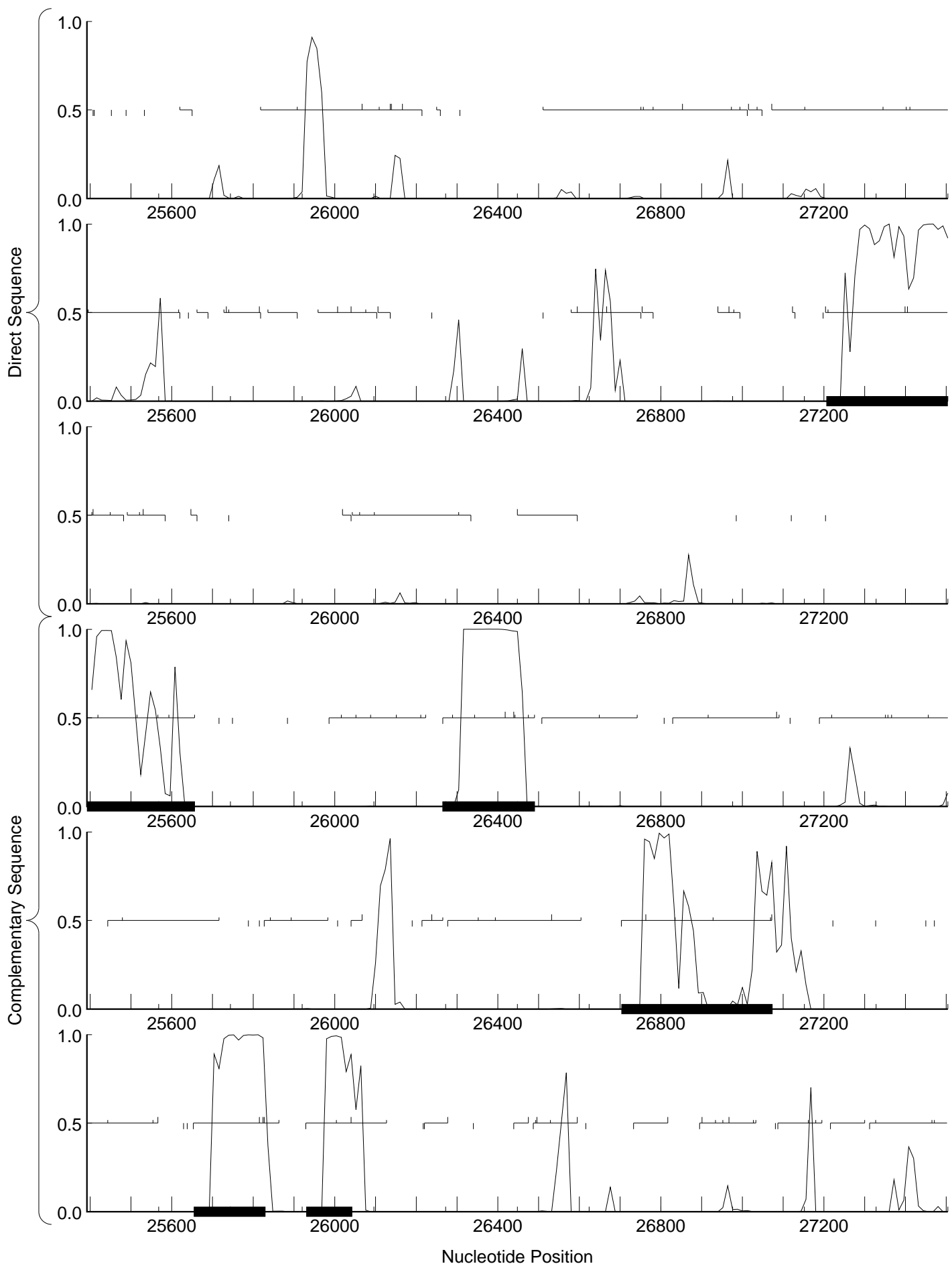
GeneMark, hmm prediction



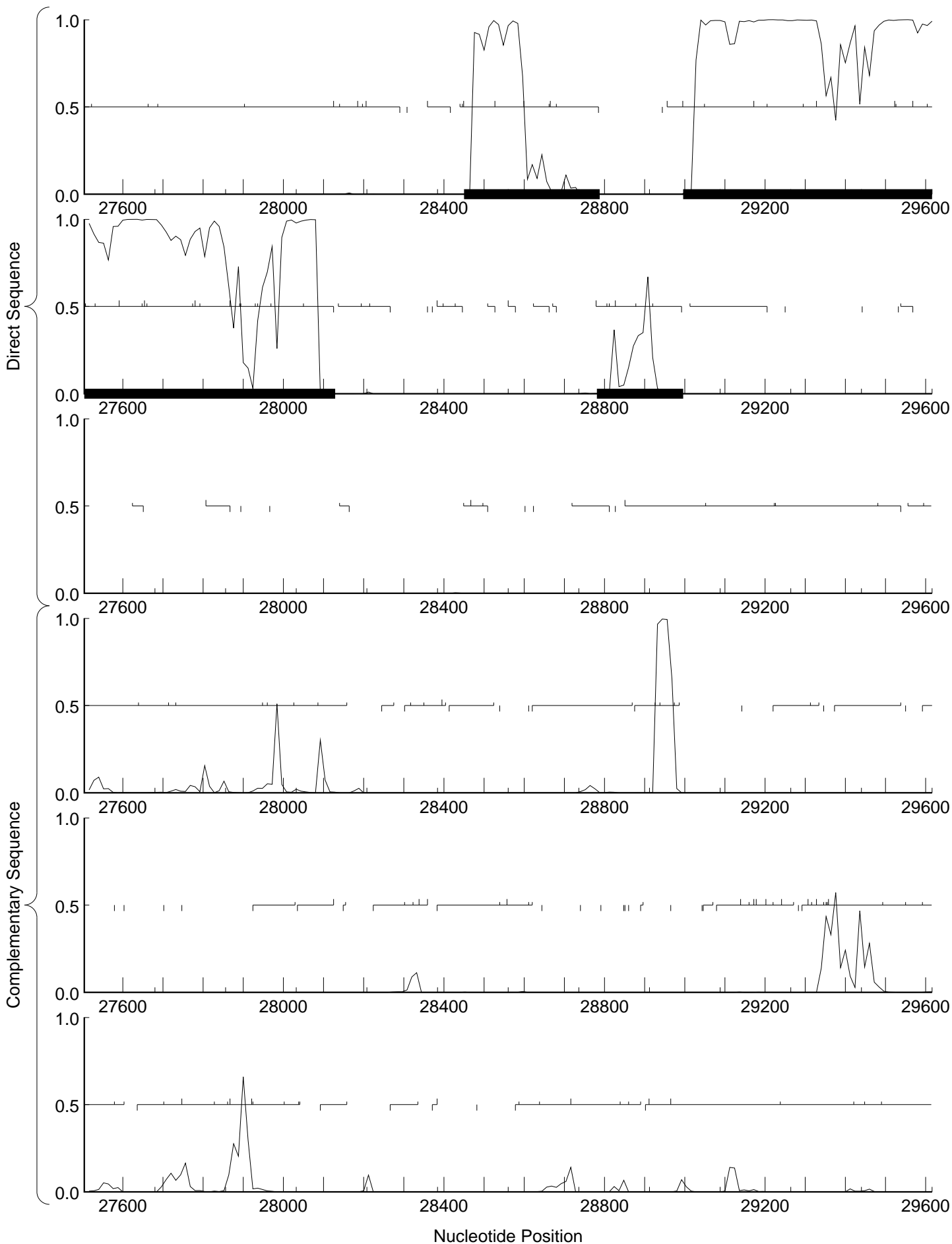
GeneMark, hmm prediction

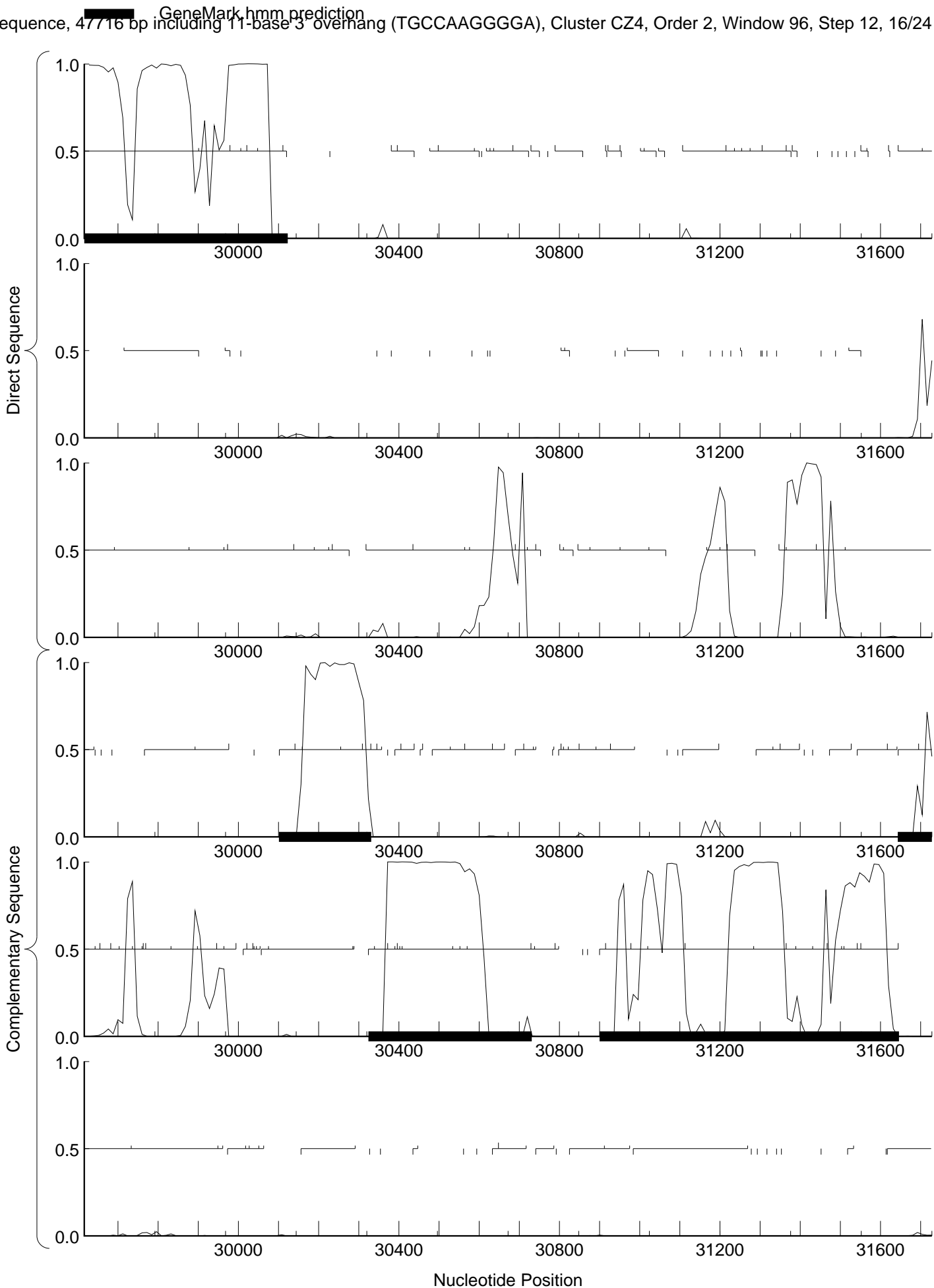


GeneMark, hmm prediction

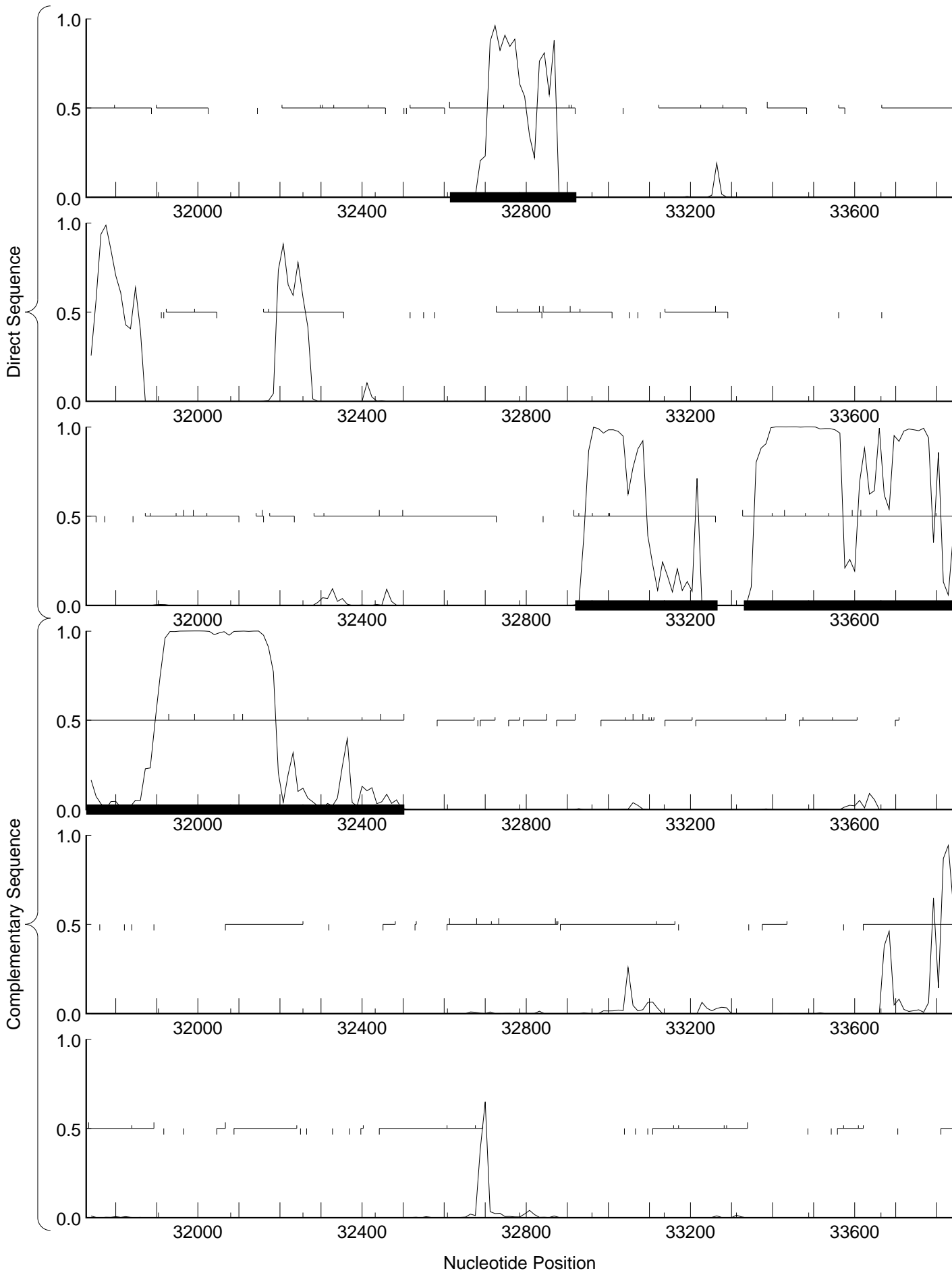


GeneMark.hmm prediction

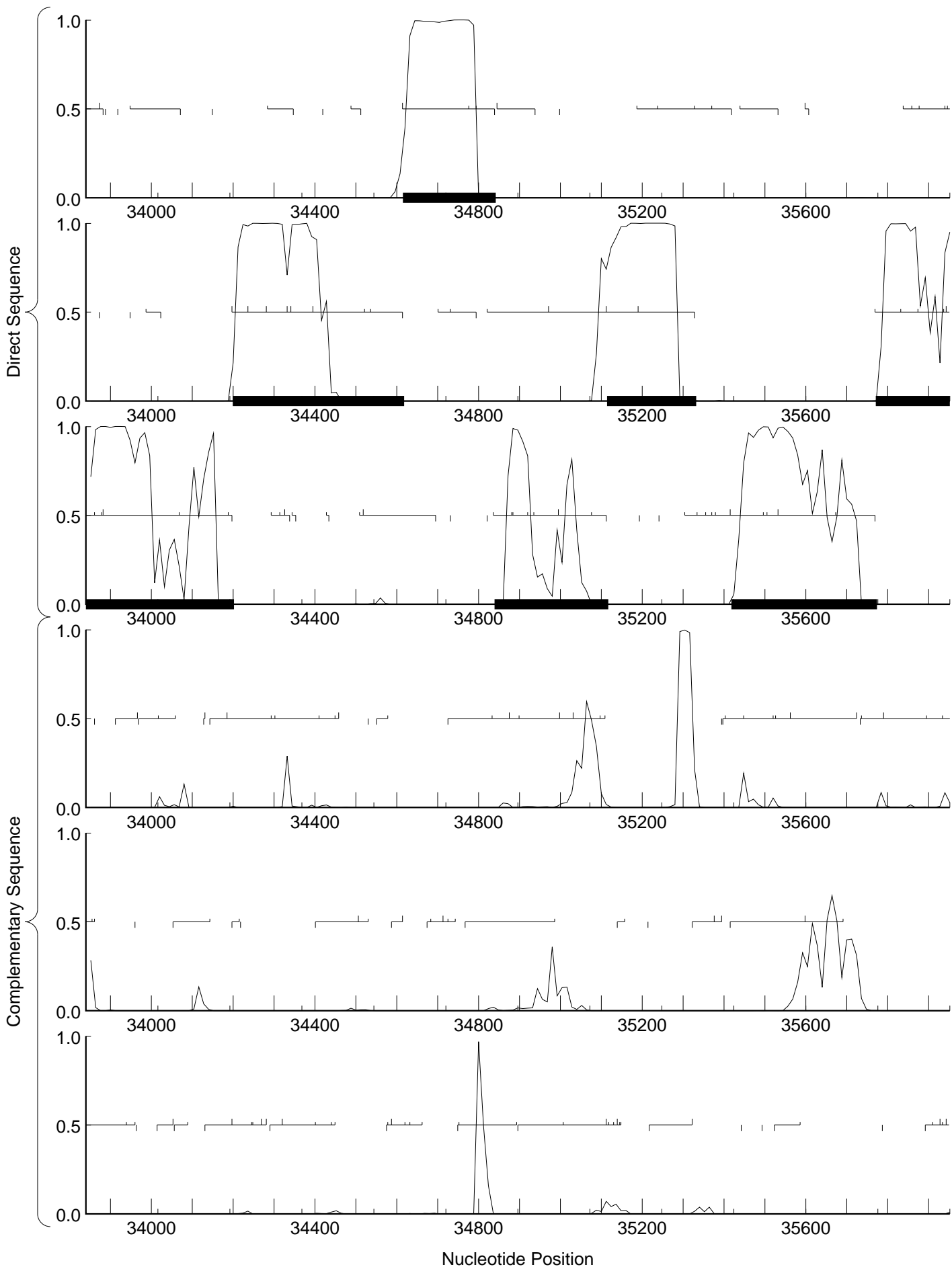


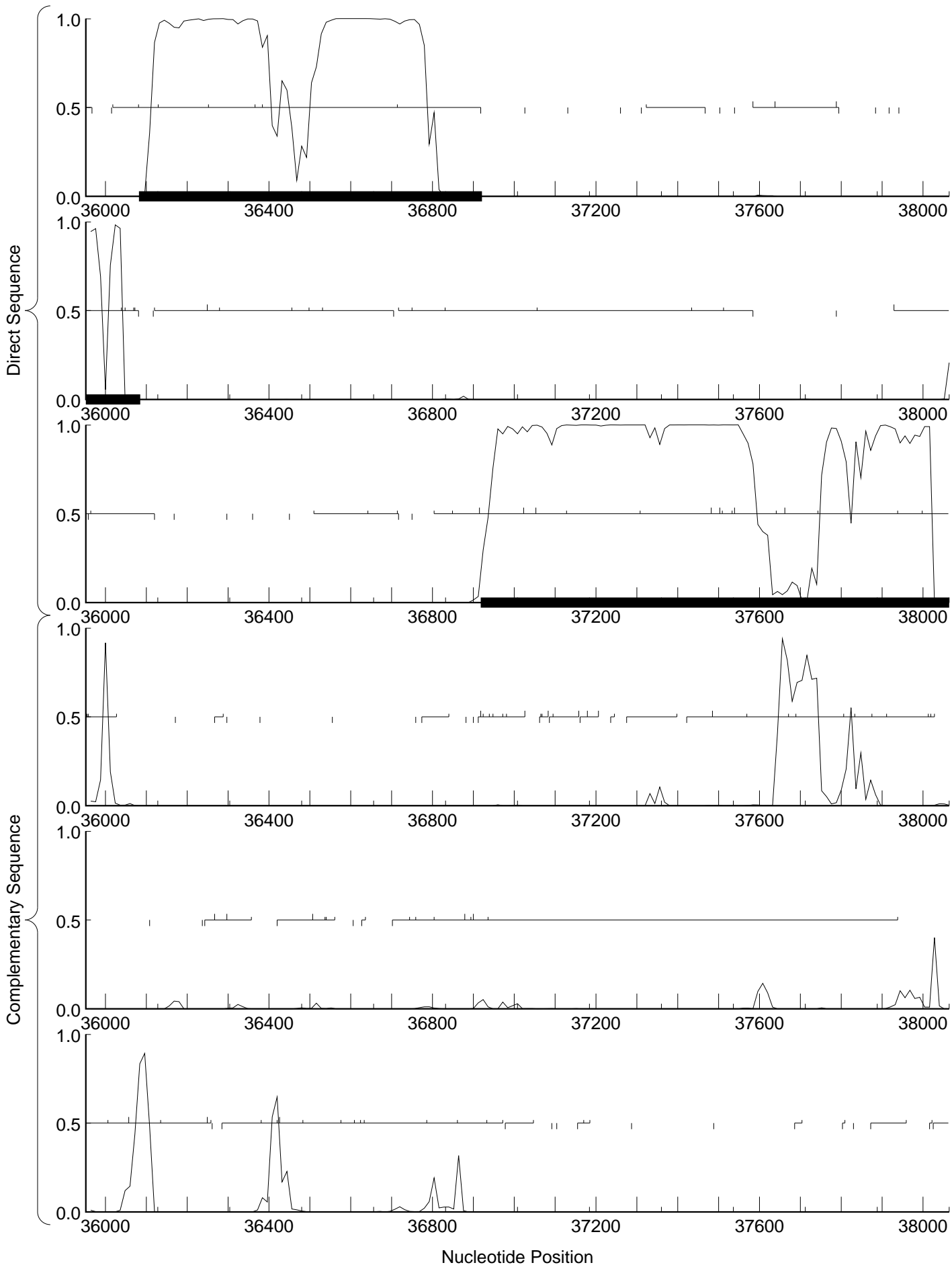


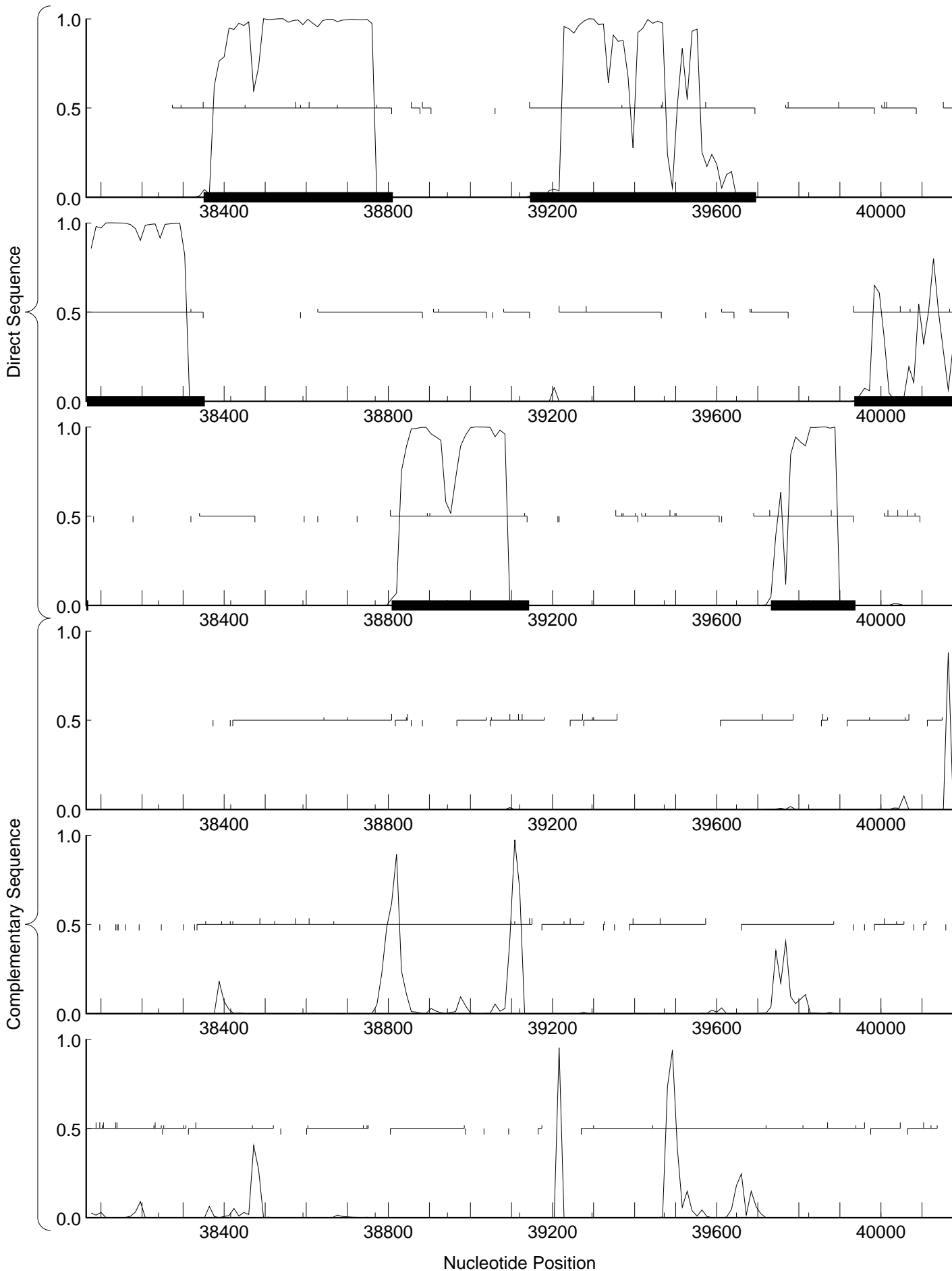
GeneMark, hmm prediction



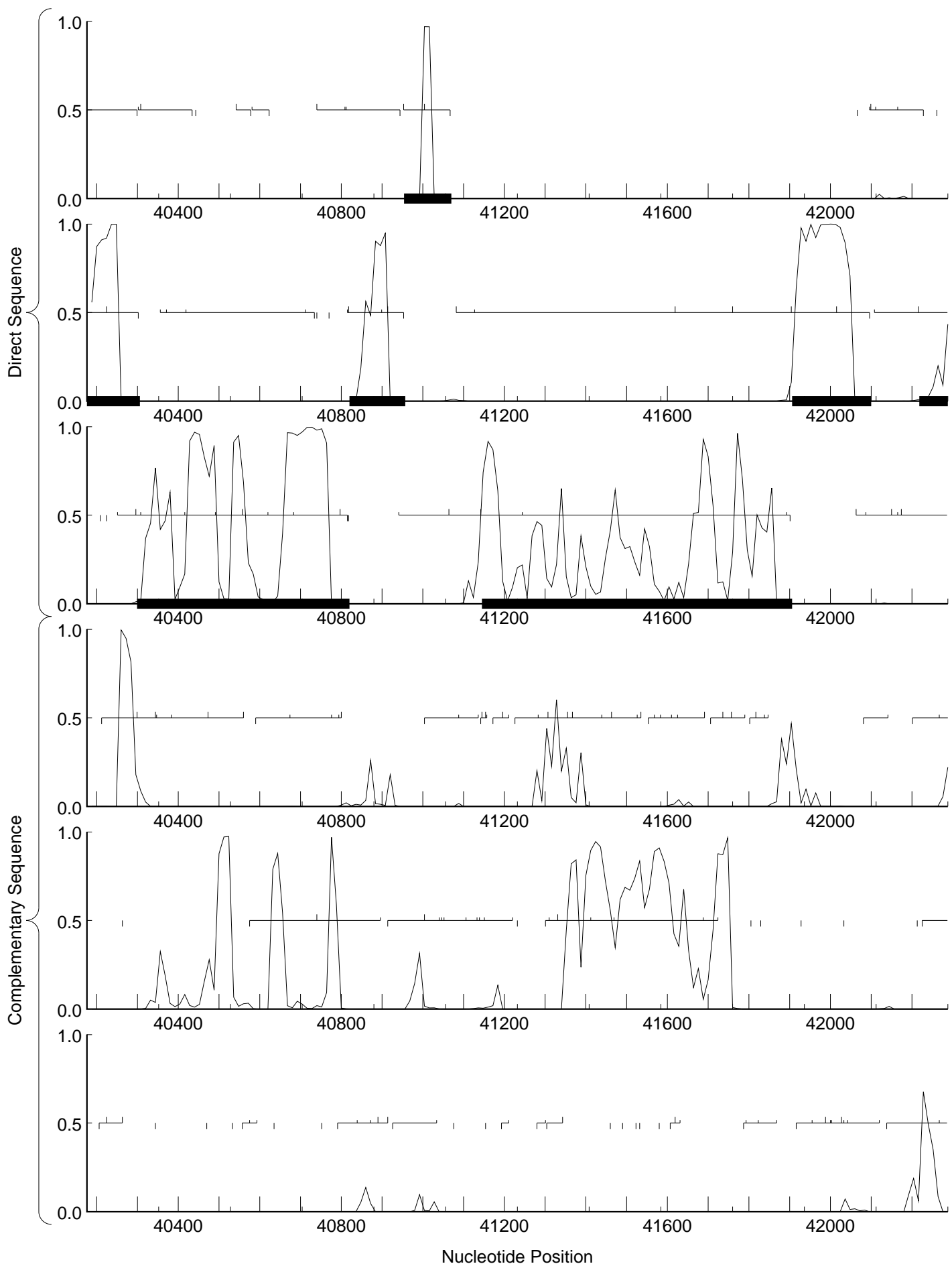
GeneMark, hmm prediction



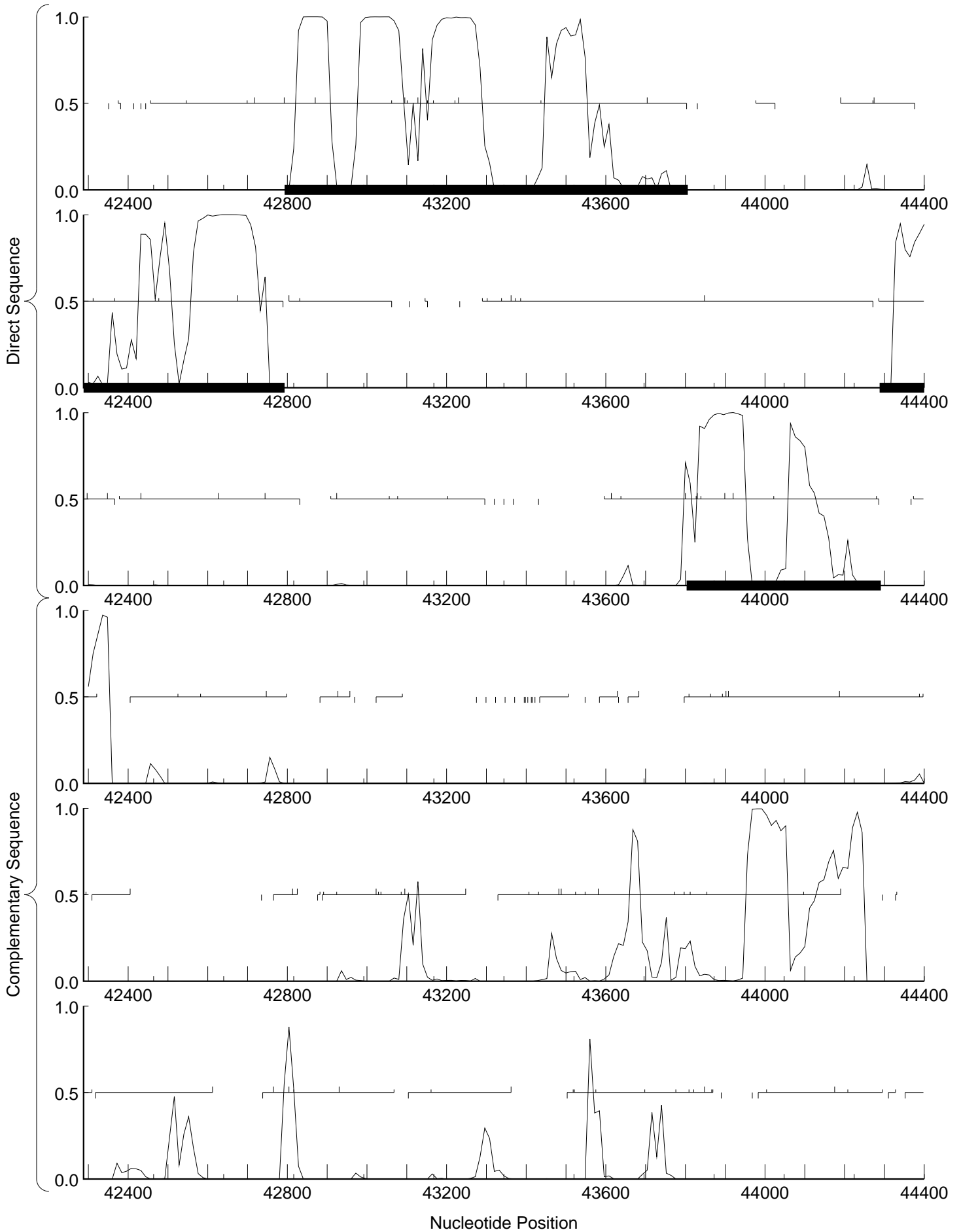




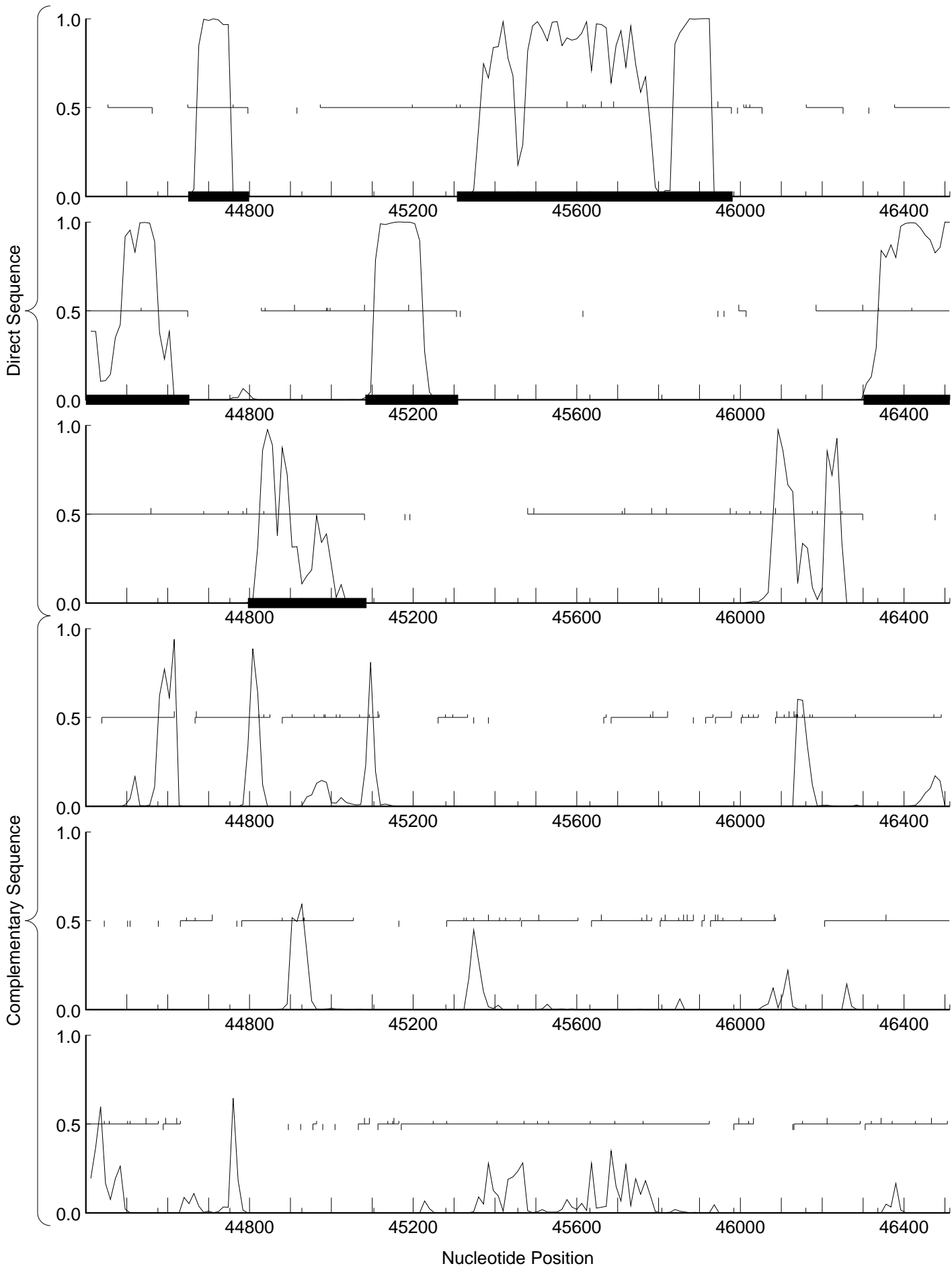
GeneMark, hmm prediction



GeneMark.hmm prediction



GeneMark.hmm prediction



GeneMark, hmm prediction

