

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Catfish complete sequence, 46888 bp including 11-base 3' overhang (AGTCCGCCG
Analysis Date : 6/20/18 at 1:11:11
Pages : 24
Sequence Length : 46888 bp
GC Content : 64.98%

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,

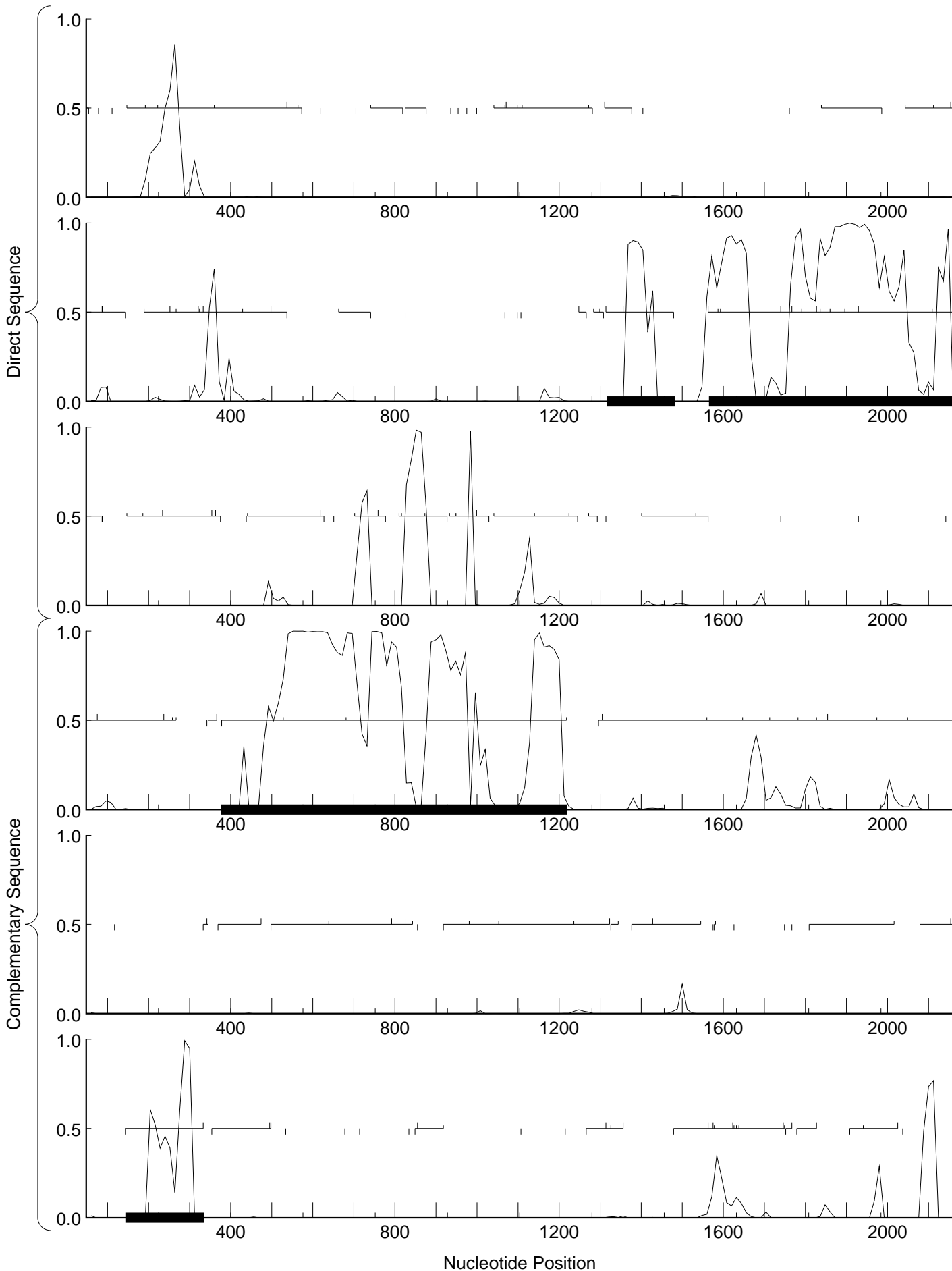
Matrix notes & comments

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

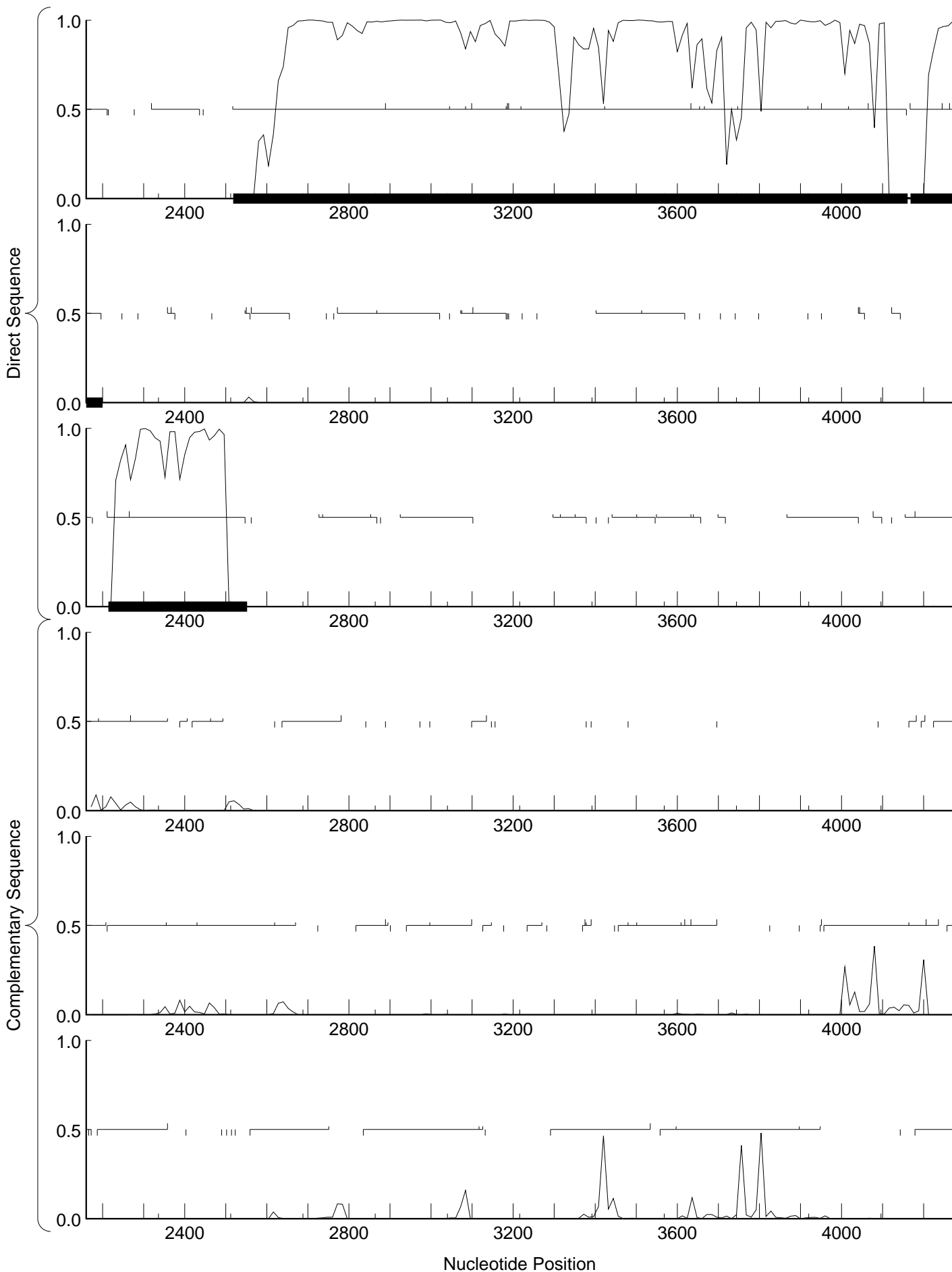
MATRIX INFORMATION

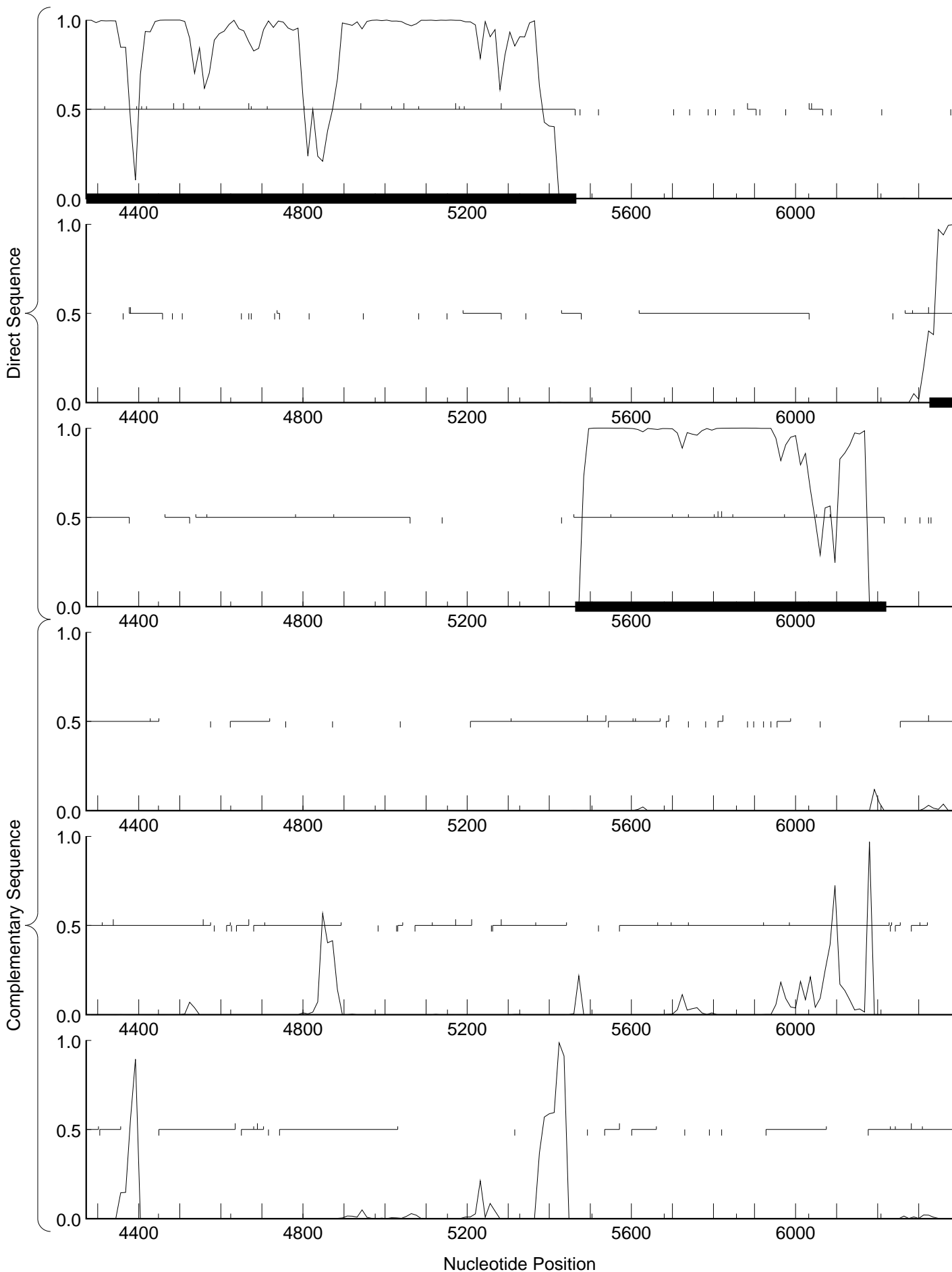
Matrix : Heuristic, GC = 65
Author : Borodovsky Laboratory - Georgia Tech
Order : 2

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

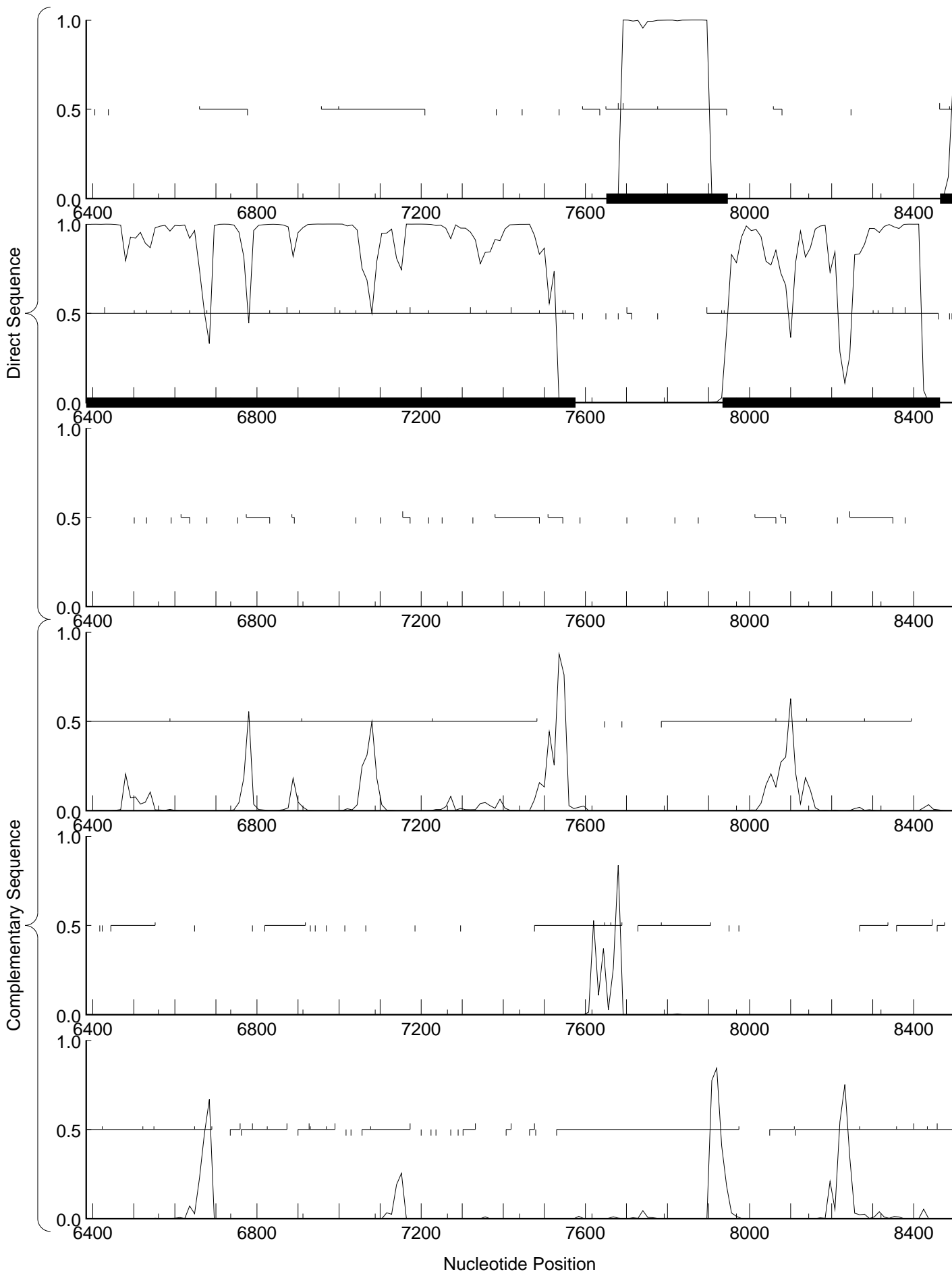


GeneMark.hmm prediction

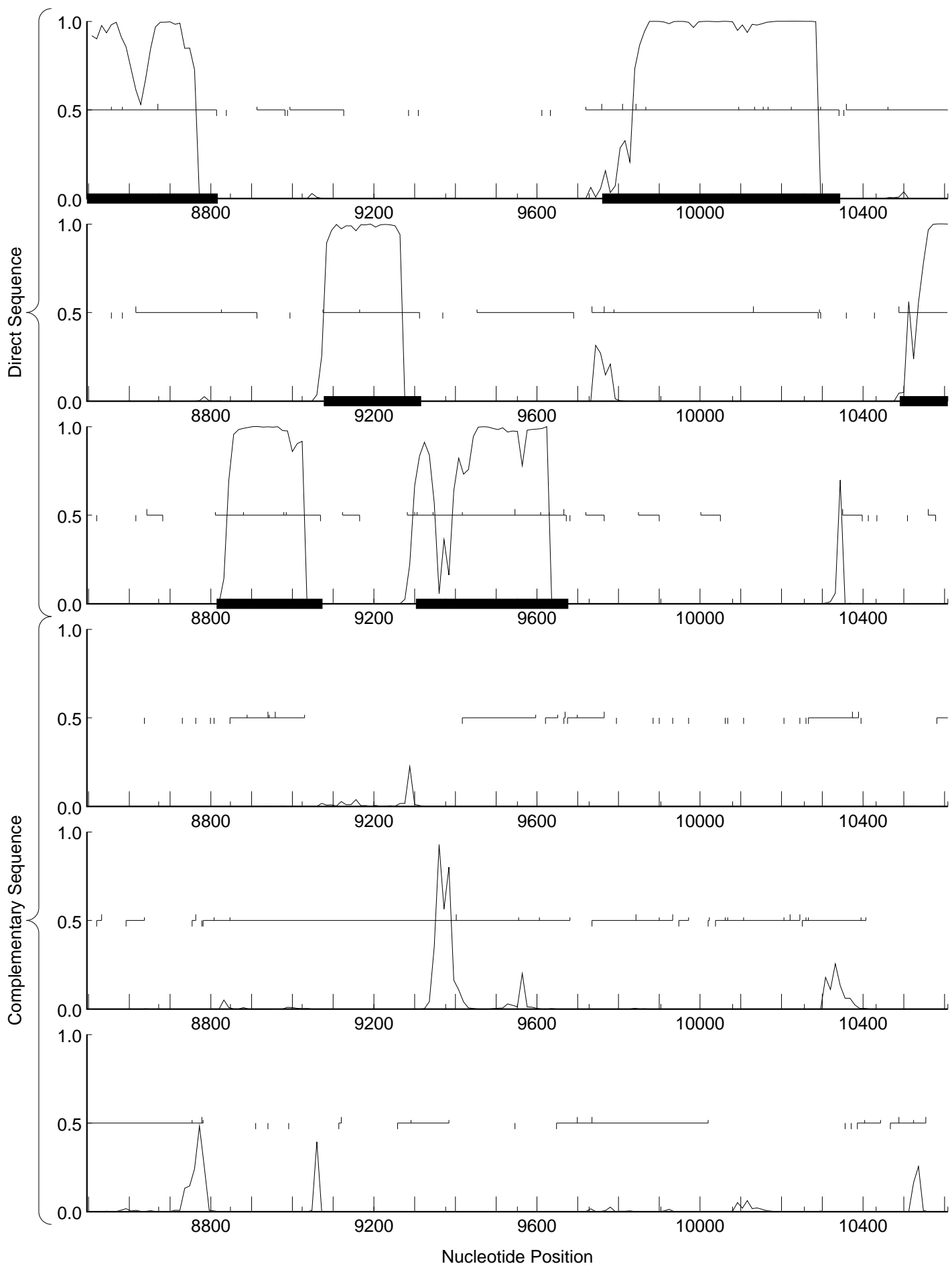




GeneMark.hmm prediction

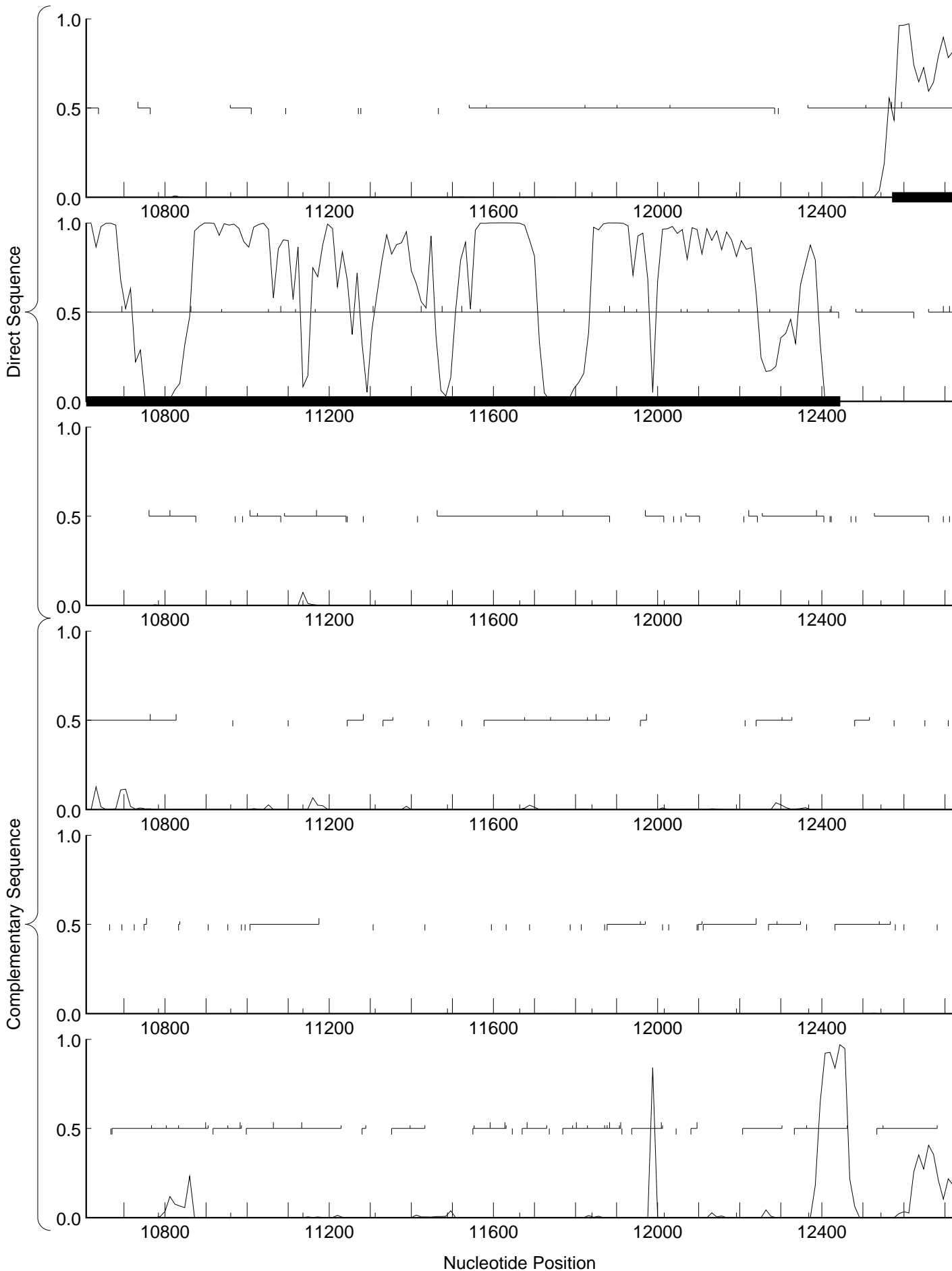


GeneMark.hmm prediction

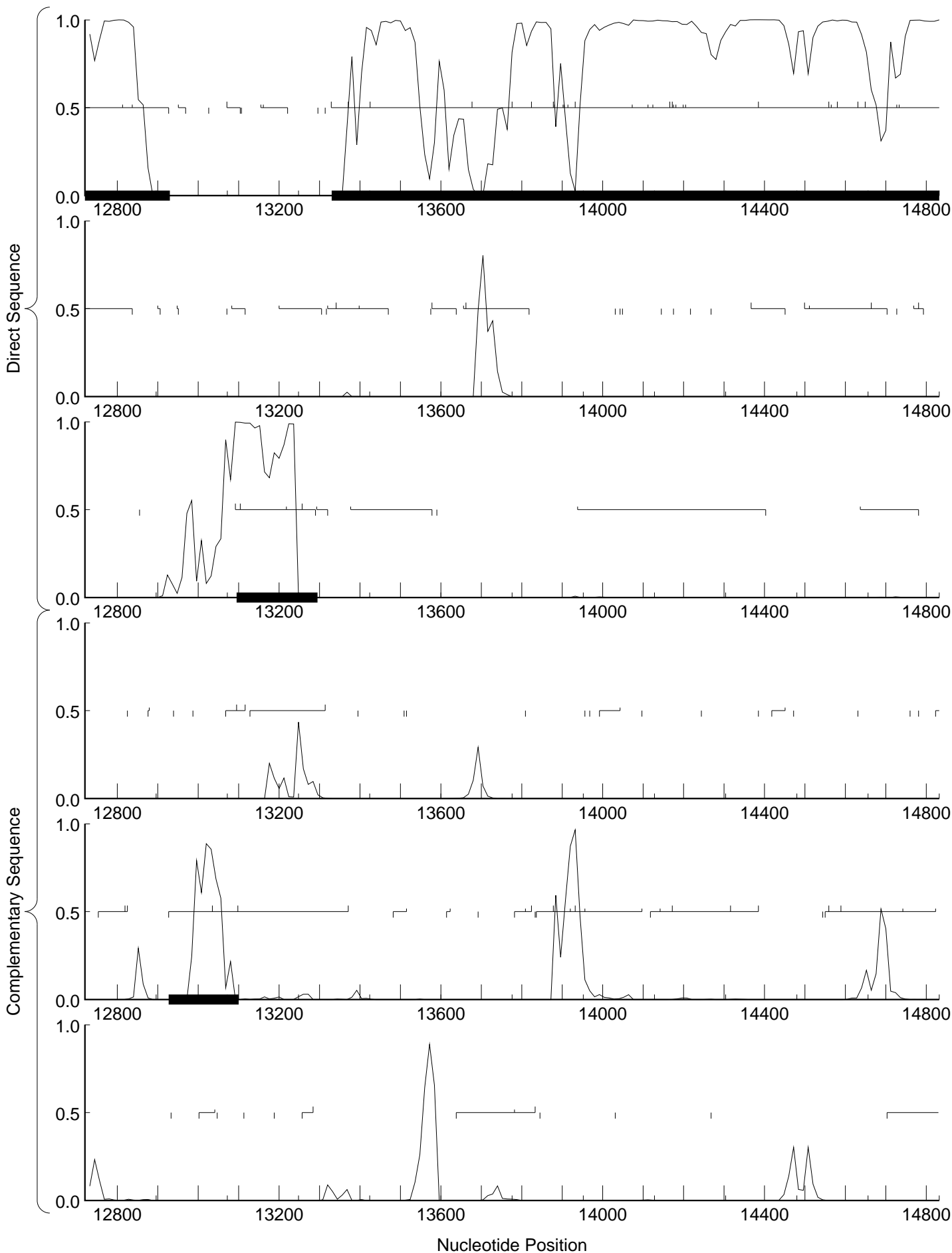




GeneMark.hmm prediction



GeneMark.hmm prediction





GeneMark.hmm prediction

