

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Kocuria phage Efeko complete sequence, 17491 bp including 9-base 3' overhang (CCCGCCCCA), C
Analysis Date : 3/20/18 at 3:58:35
Pages : 10
Sequence Length : 17491 bp
GC Content : 68.59%

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

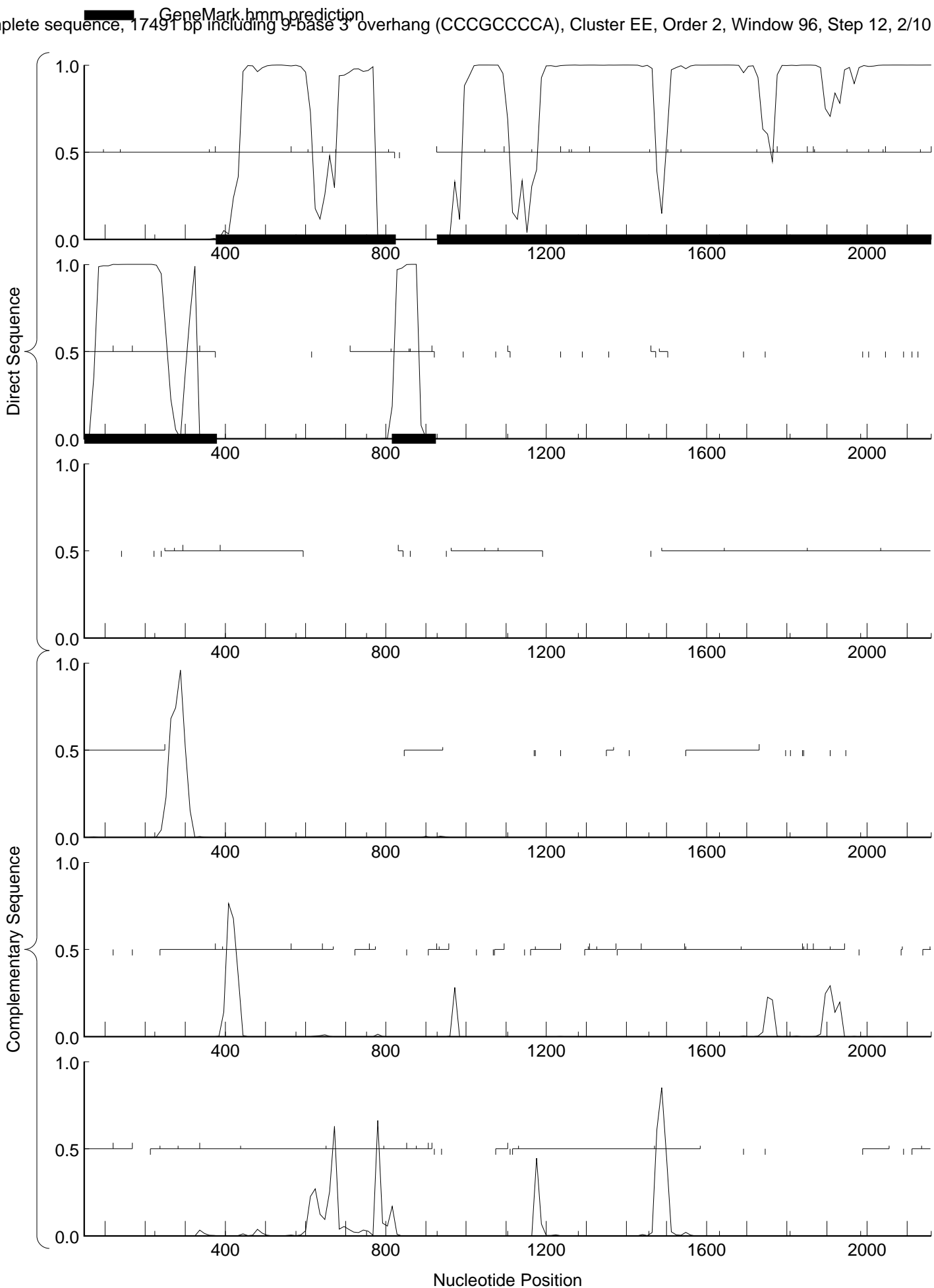
For details on the model building procedure see:
Besemer J. and Borodovsky M.
"Heuristic approach to deriving models for gene finding"
NAR, 1999, Vol. 27, No. 19, pp. 3911-3920

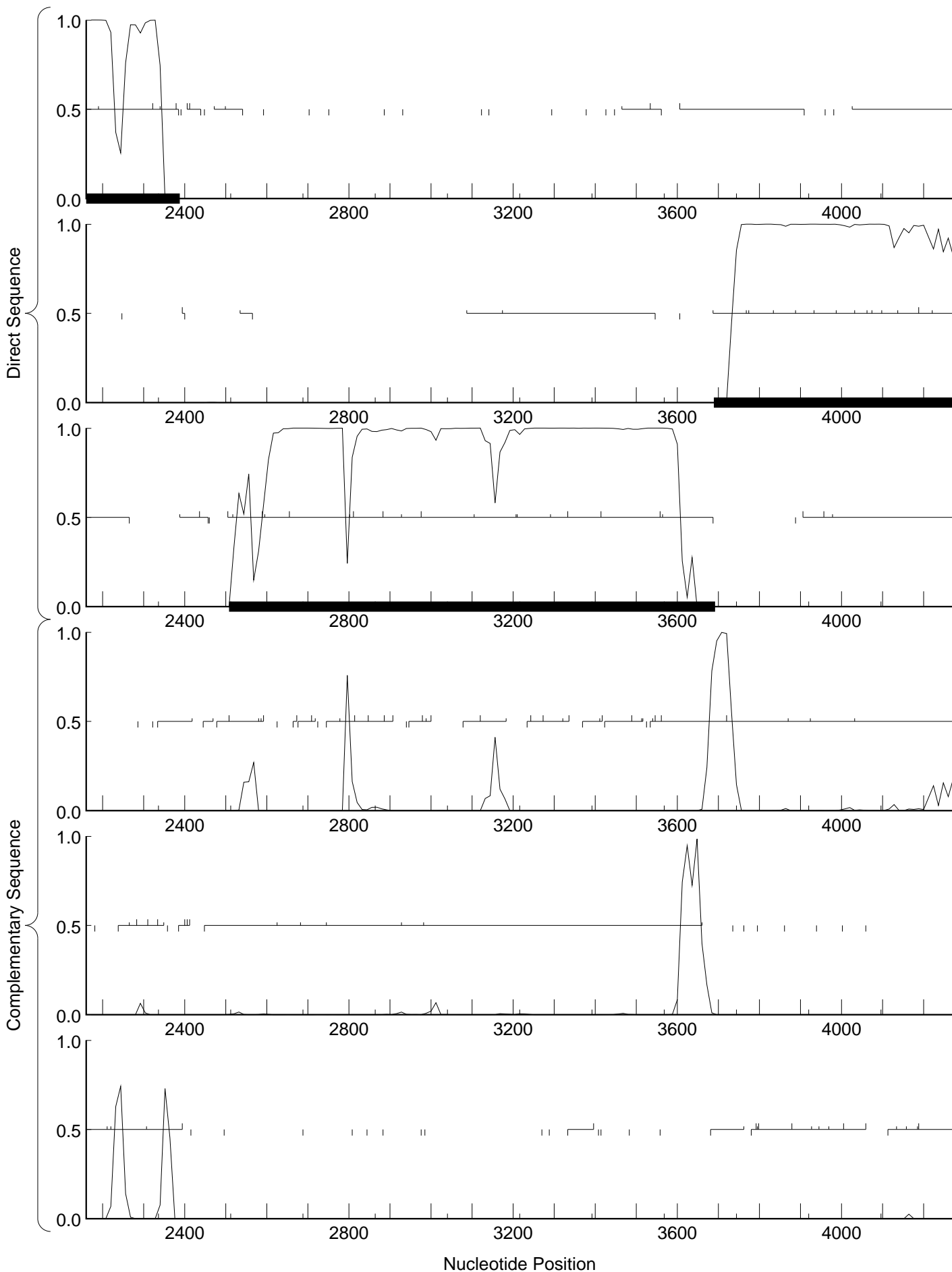
MATRIX INFORMATION

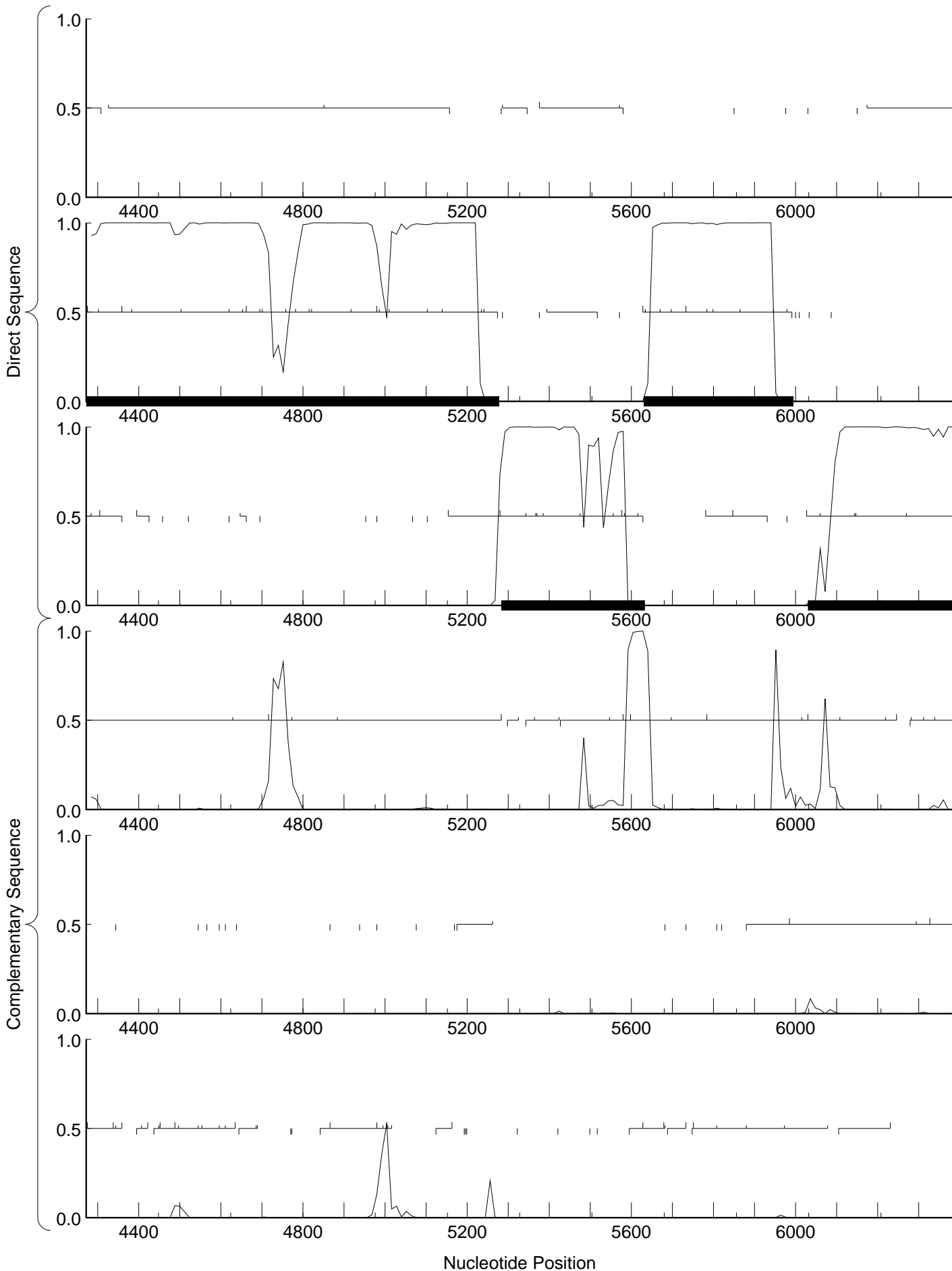
Matrix : Heuristic, GC = 69
Author : Borodovsky Laboratory - Georgia Tech
Order : 2

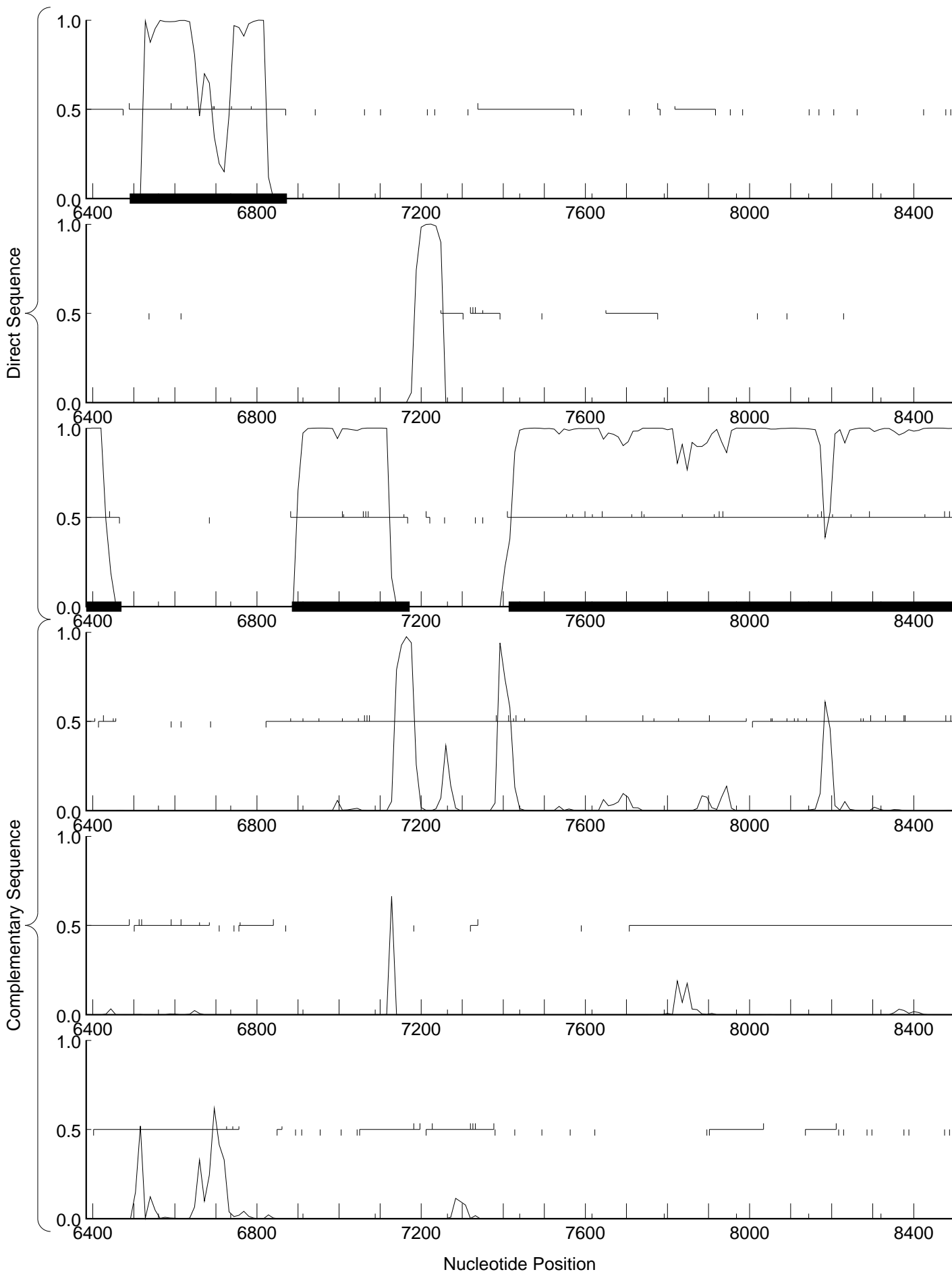
Send questions / comments to:
Dr. M. Borodovsky
Georgia Institute of Technology
School of Biology
Atlanta, GA 30332-0230

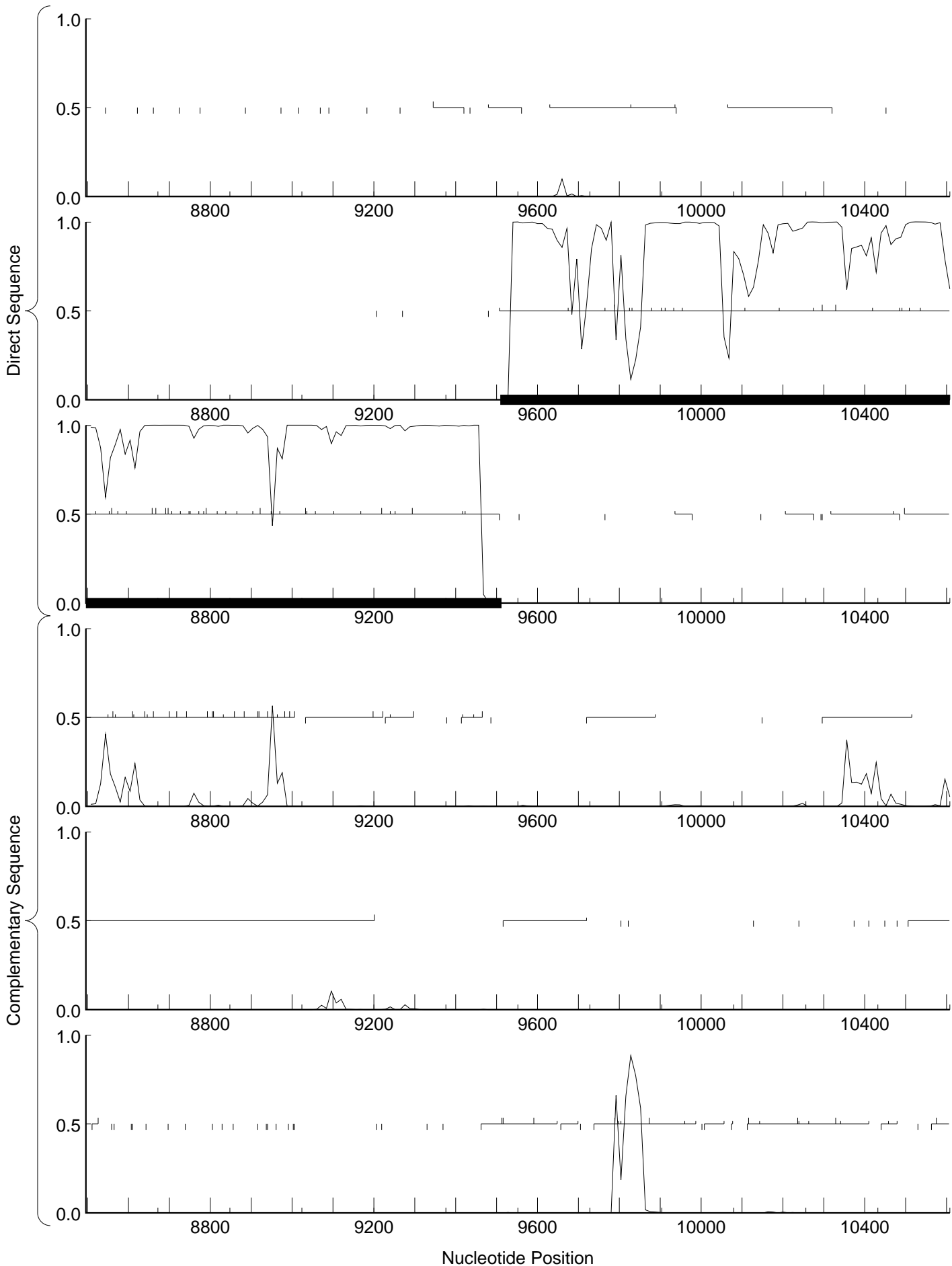
Geno complete sequence, 17491 bp including 9-base 3' overhang (CCCGCCCCA), Cluster EE, Order 2, Window 96, Step 12, 2/10



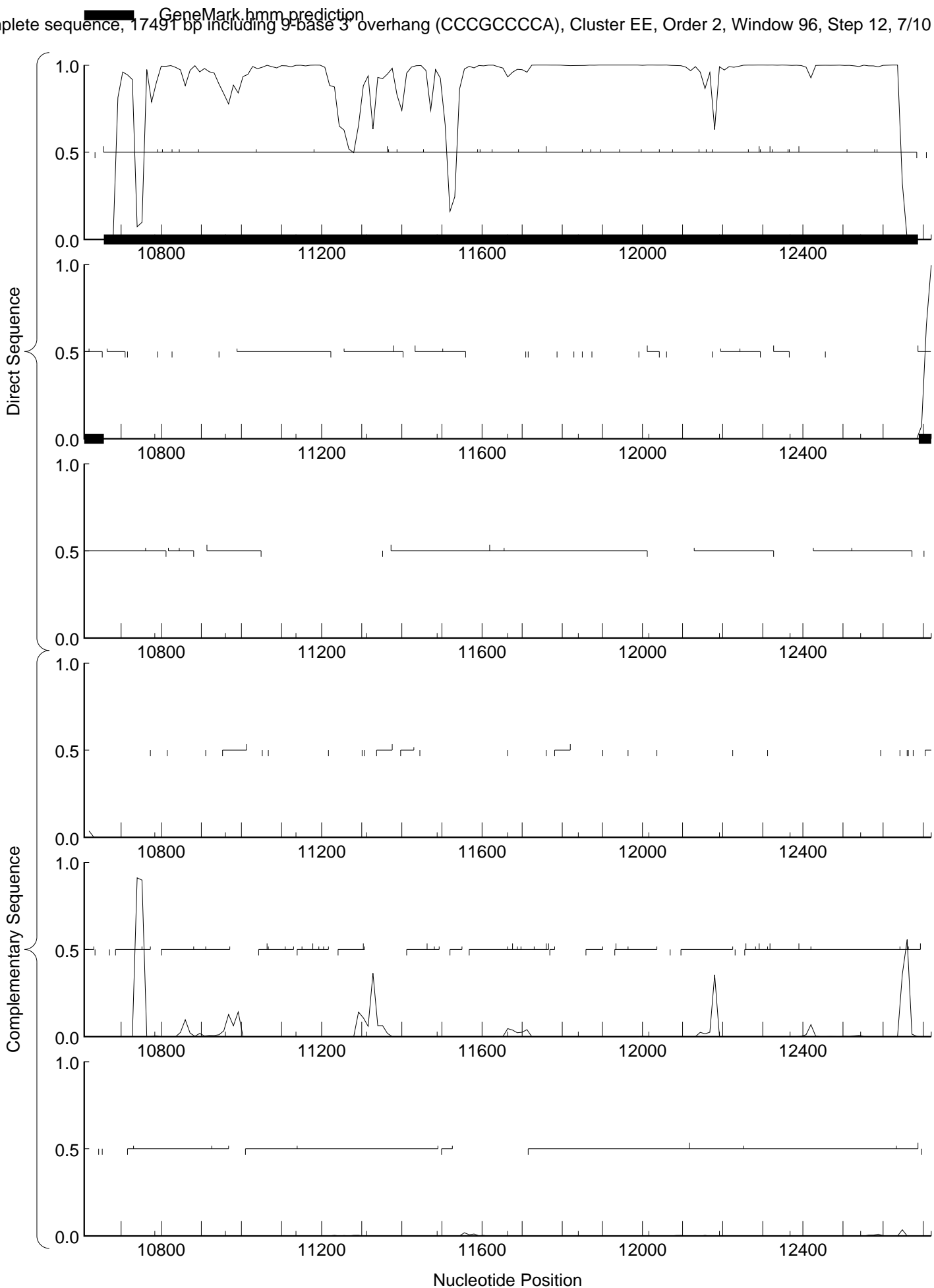


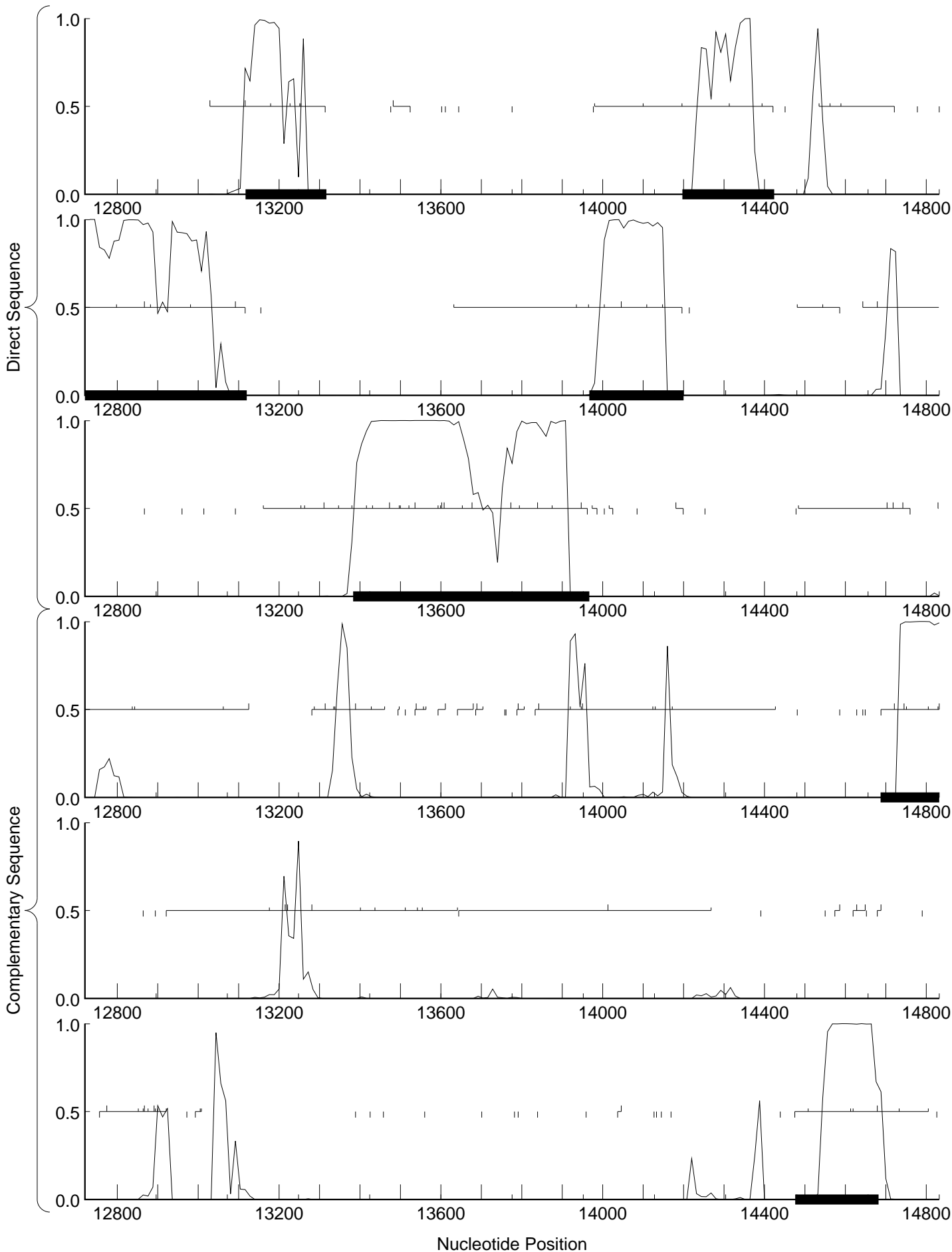


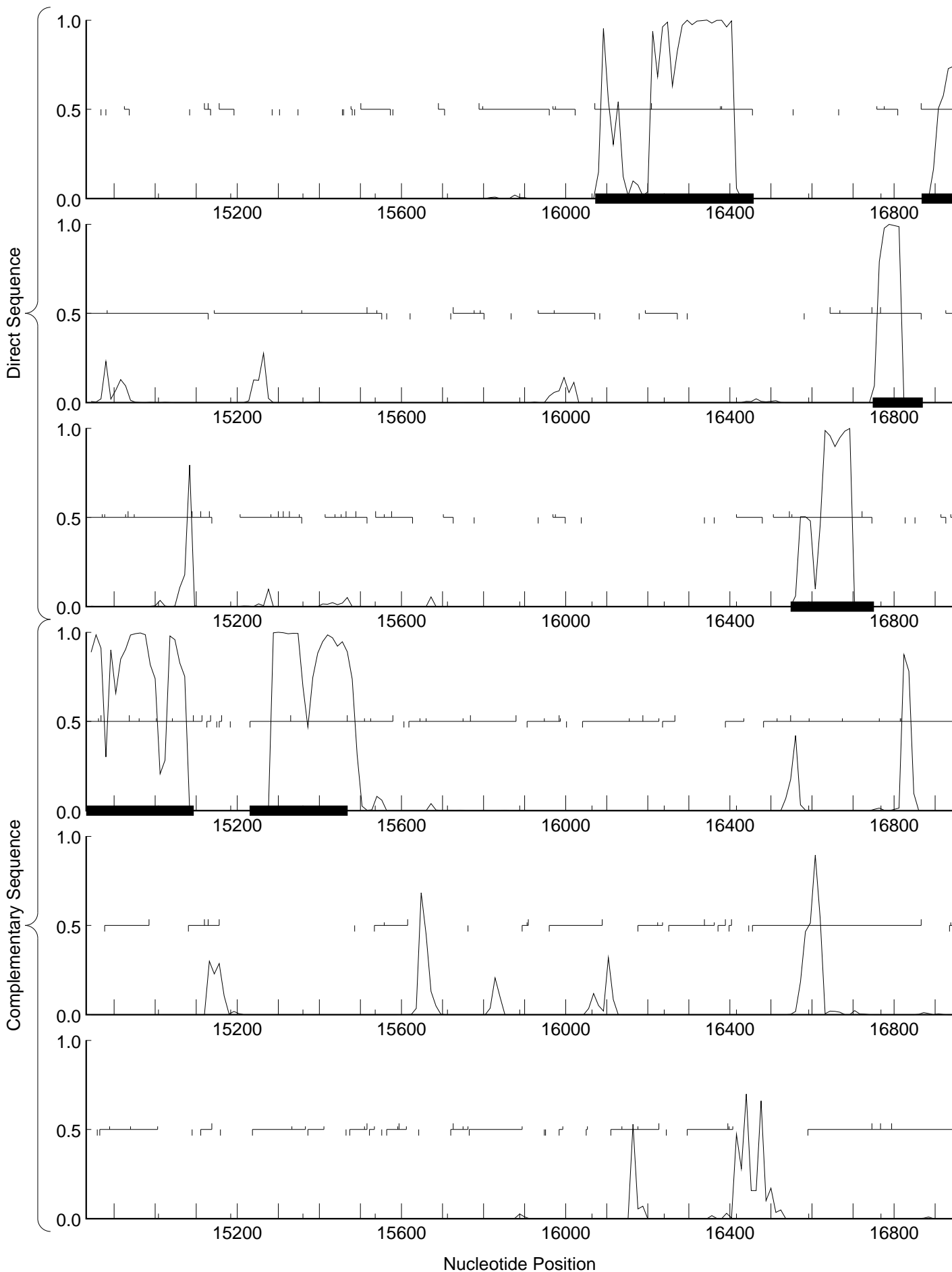




Geno complete sequence, 17491 bp including 9-base 3' overhang (CCCGCCCCA), Cluster EE, Order 2, Window 96, Step 12, 7/10







GeneMark.hmm prediction
ko complete sequence, 17491 bp including 9-base 3' overhang (CCGCCCCA), Cluster EE, Order 2, Window 96, Step 12, 10/10

