

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence seq.fna_for_gm

Analysis Date 3/10/17 at 16:26:15

Pages 29 58669 bp Sequence Length : GC Content 66.17%

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 = 66

Author Borodovsky Laboratory - Georgia Ted

Order

Send questions / comments to:

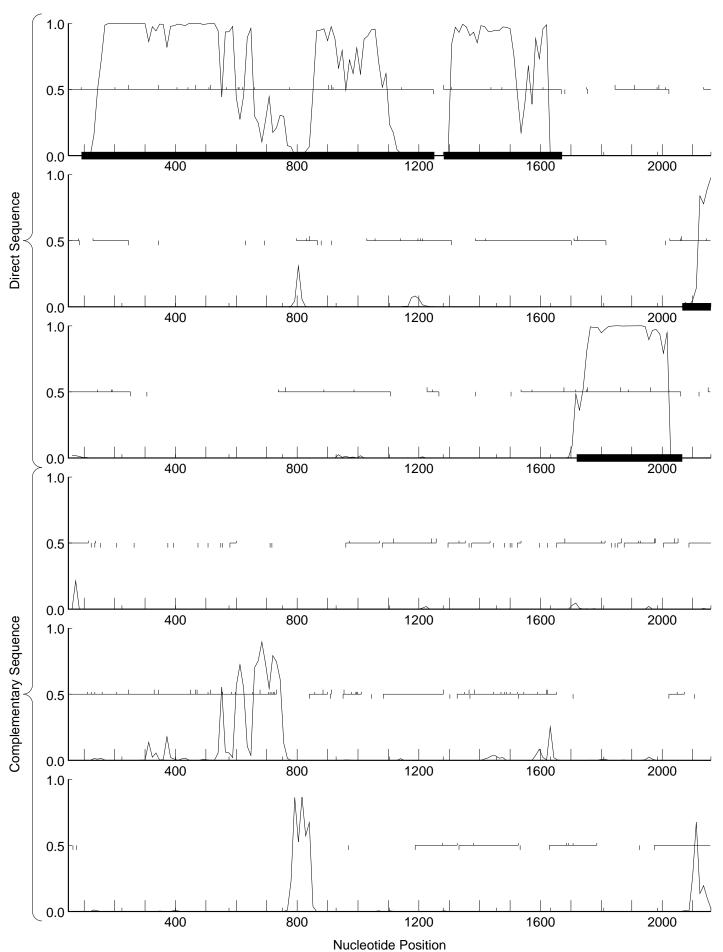
Dr. M. Borodovsky

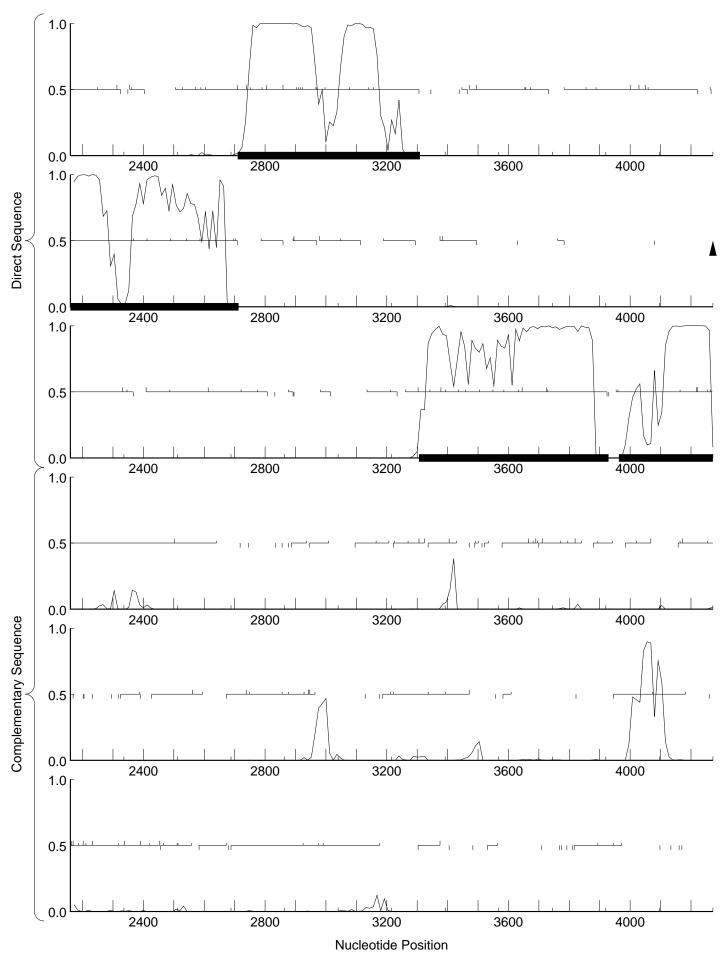
Georgia Institute of Technology

School of Biology

Atlanta, GA 30332-0230

For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu





5200

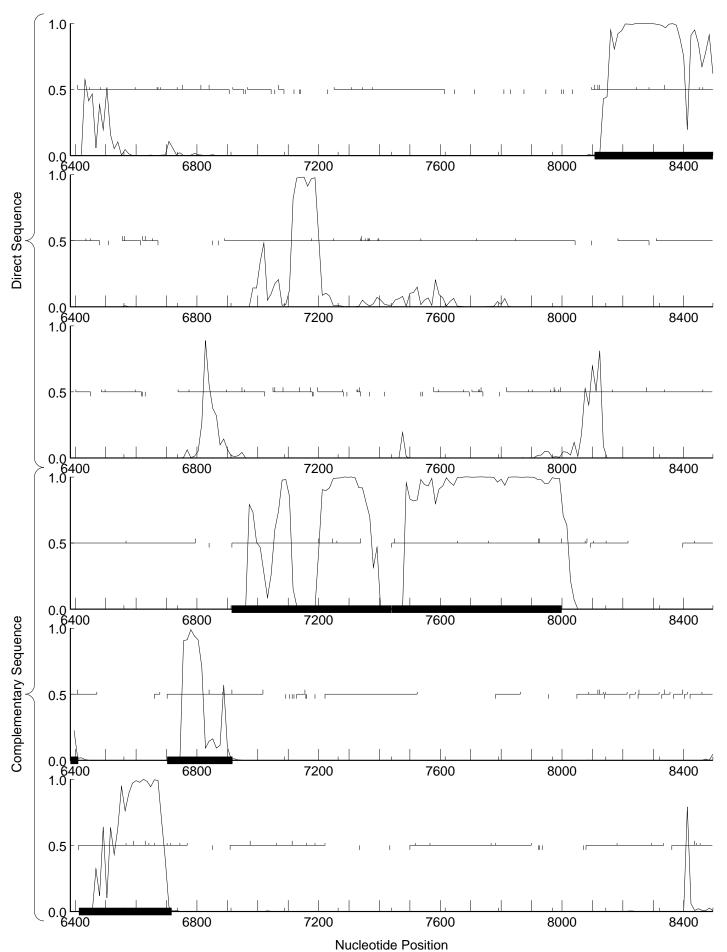
Nucleotide Position

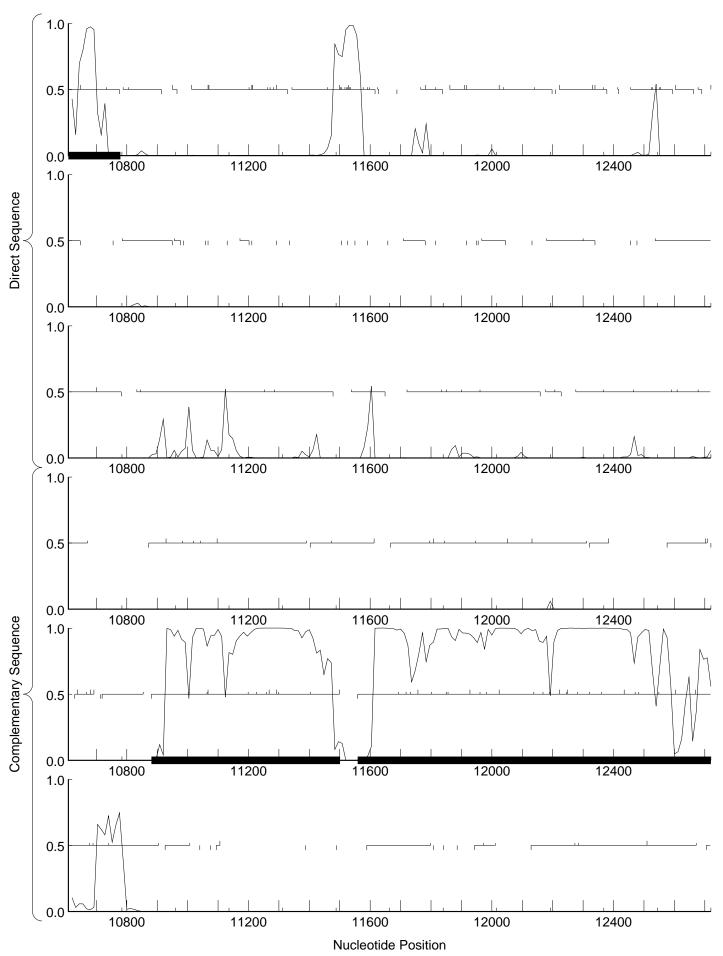
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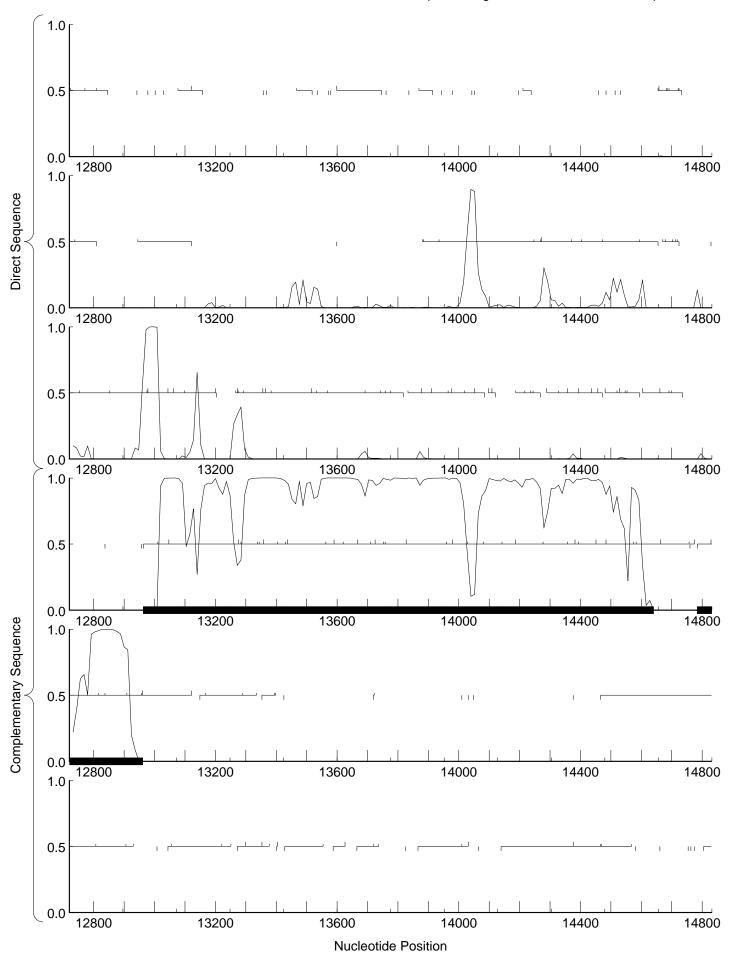
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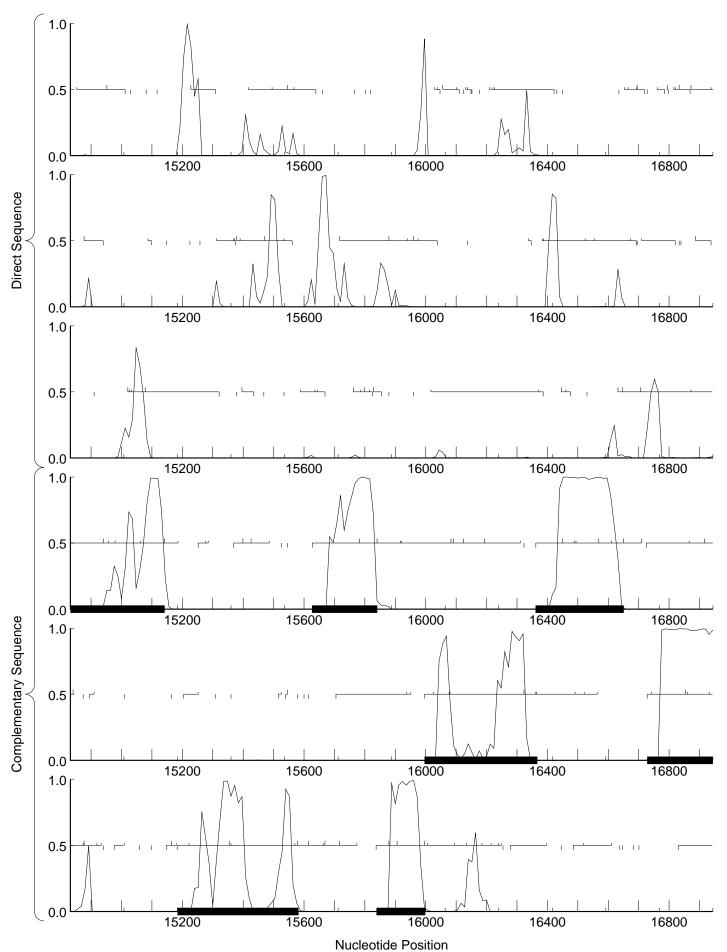
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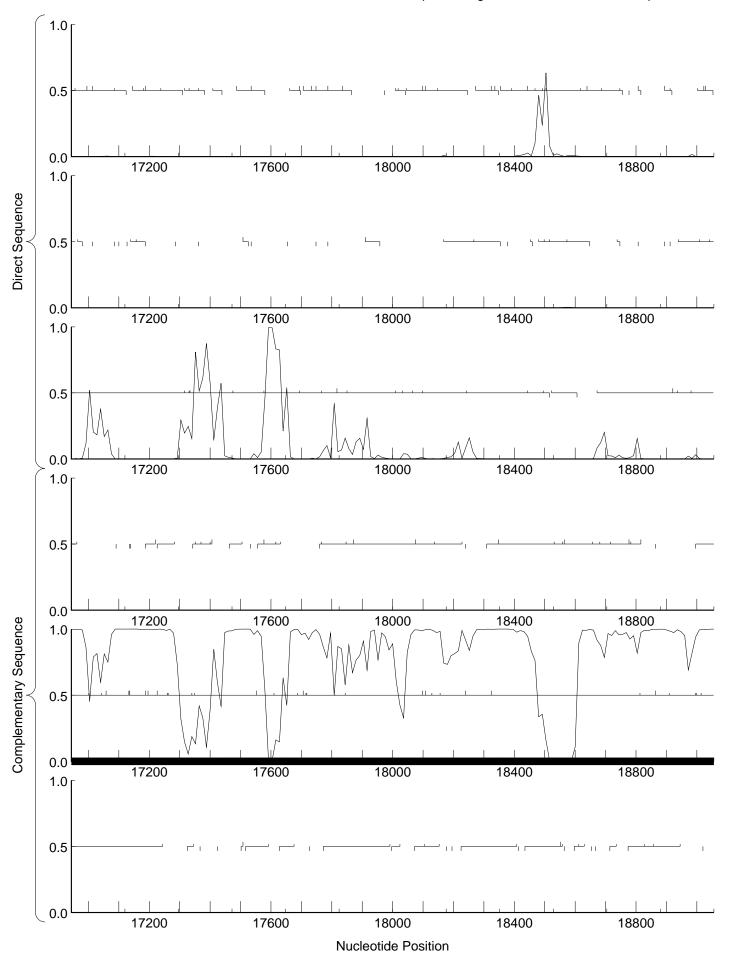
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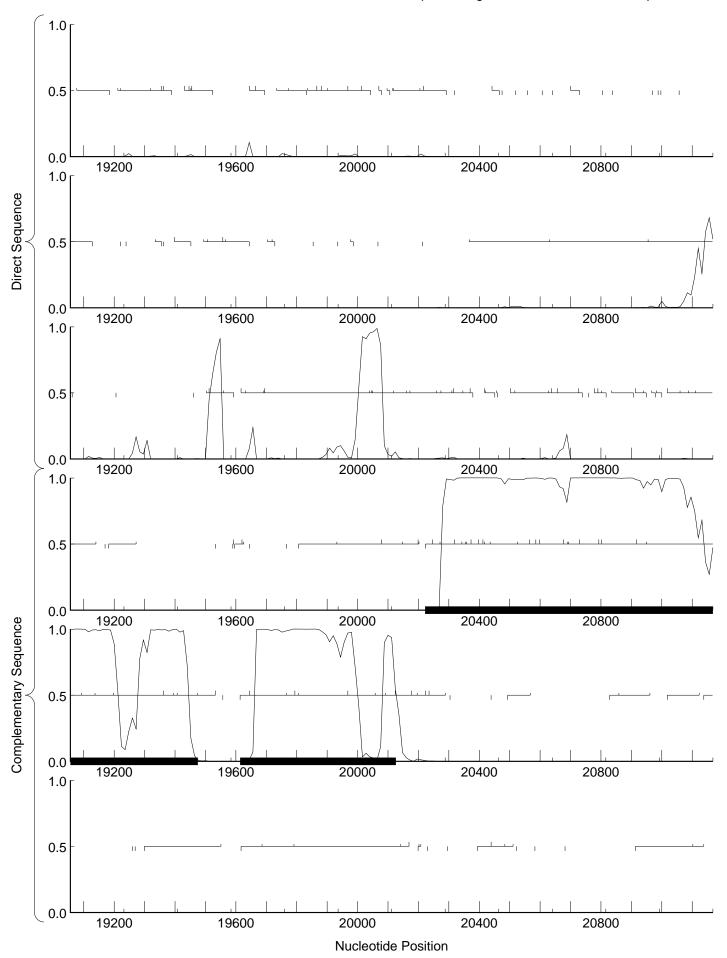


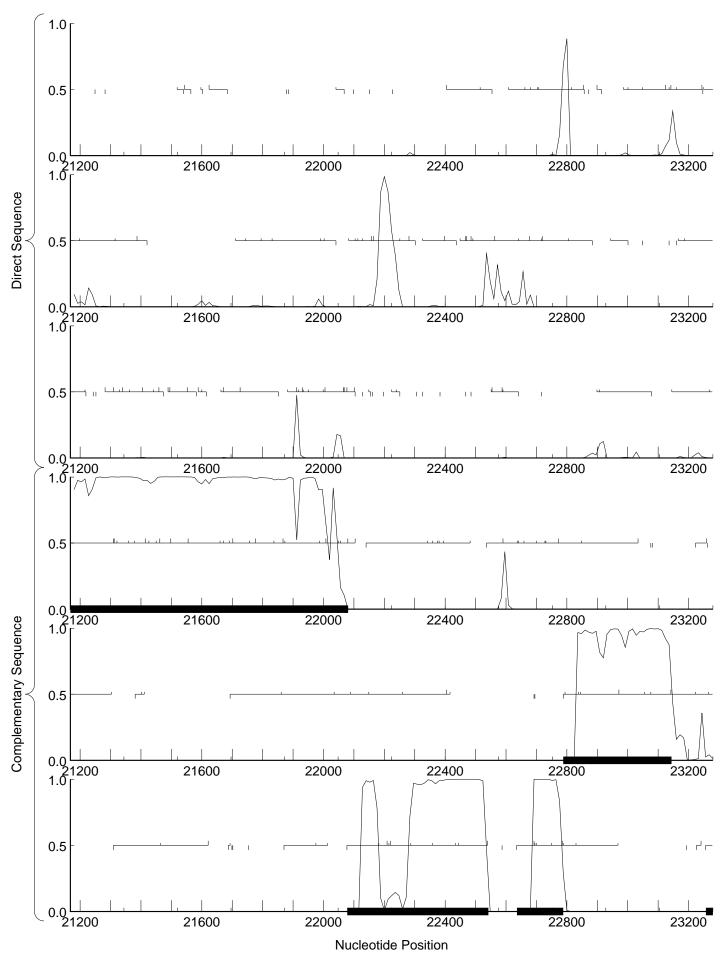


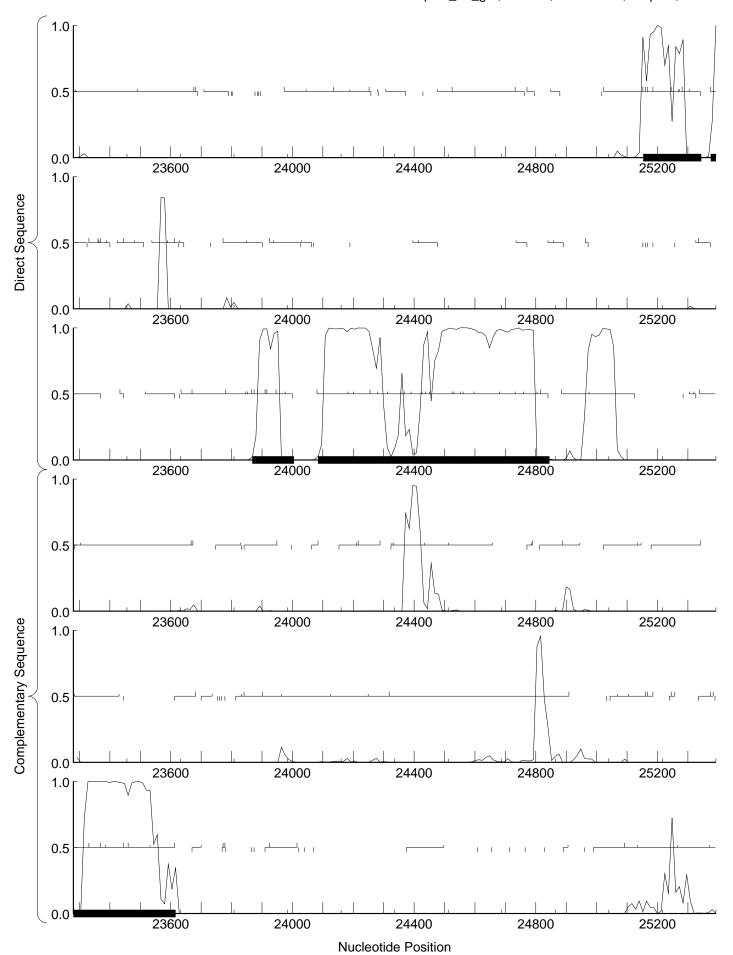


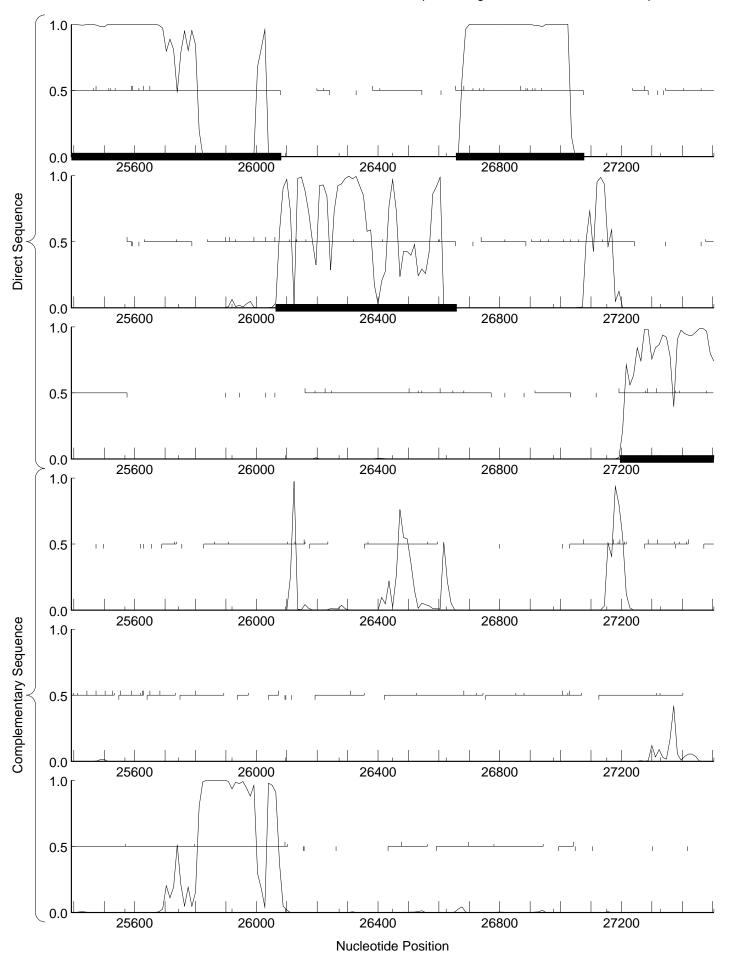


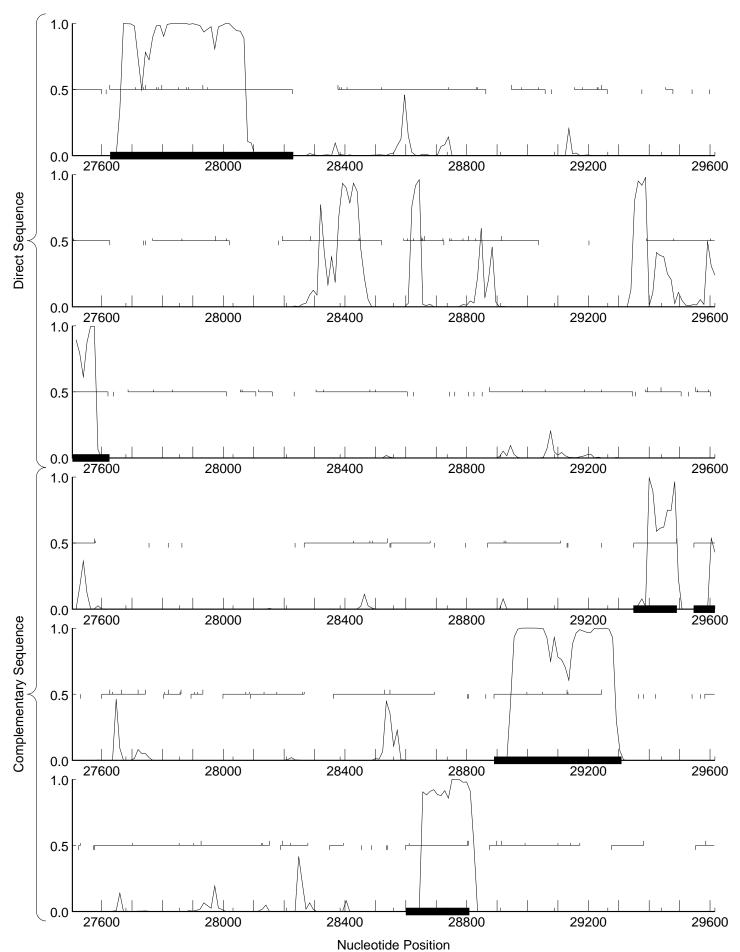


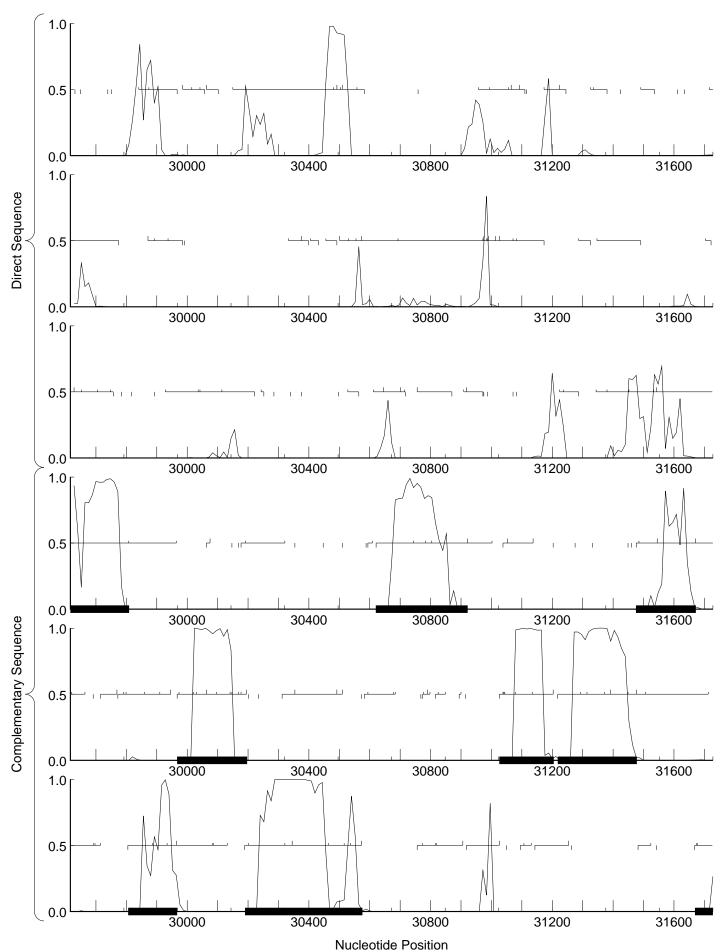


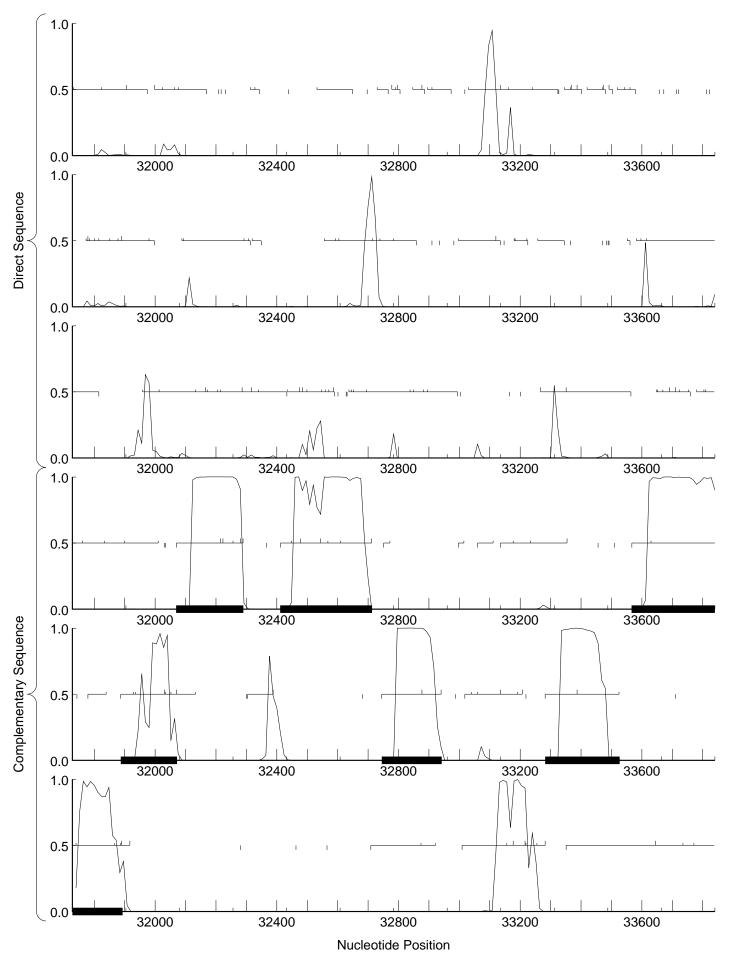


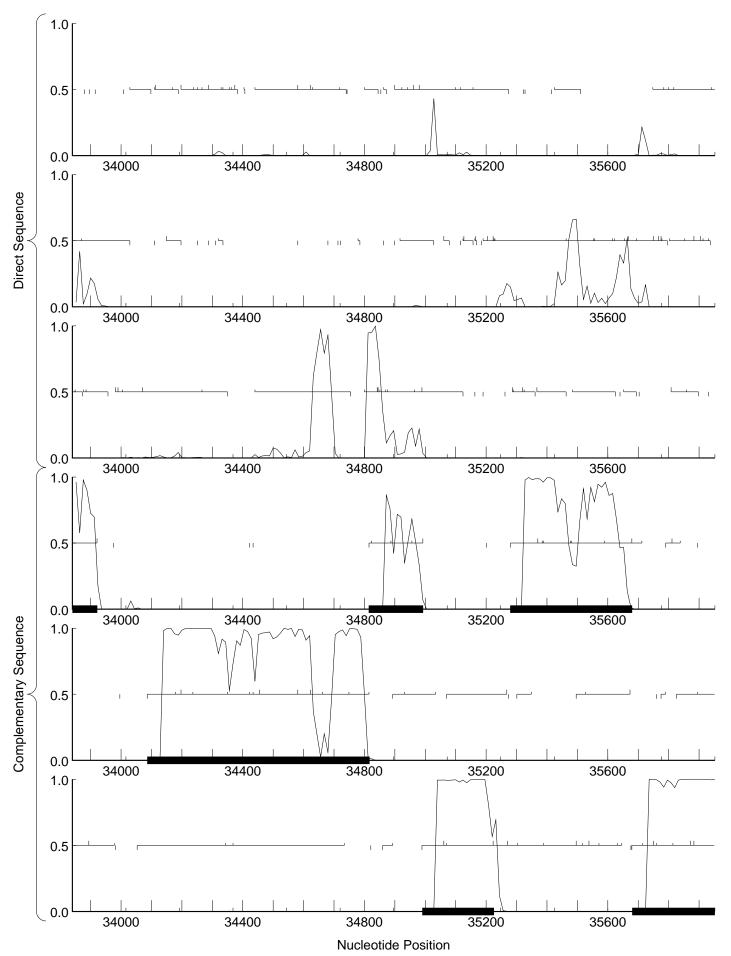


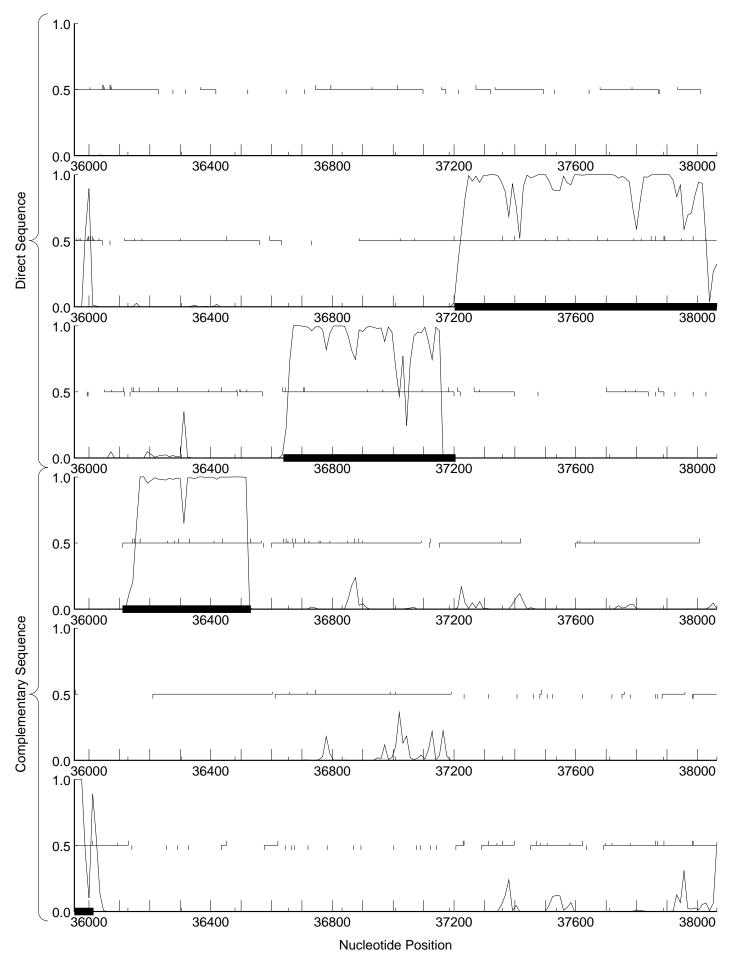


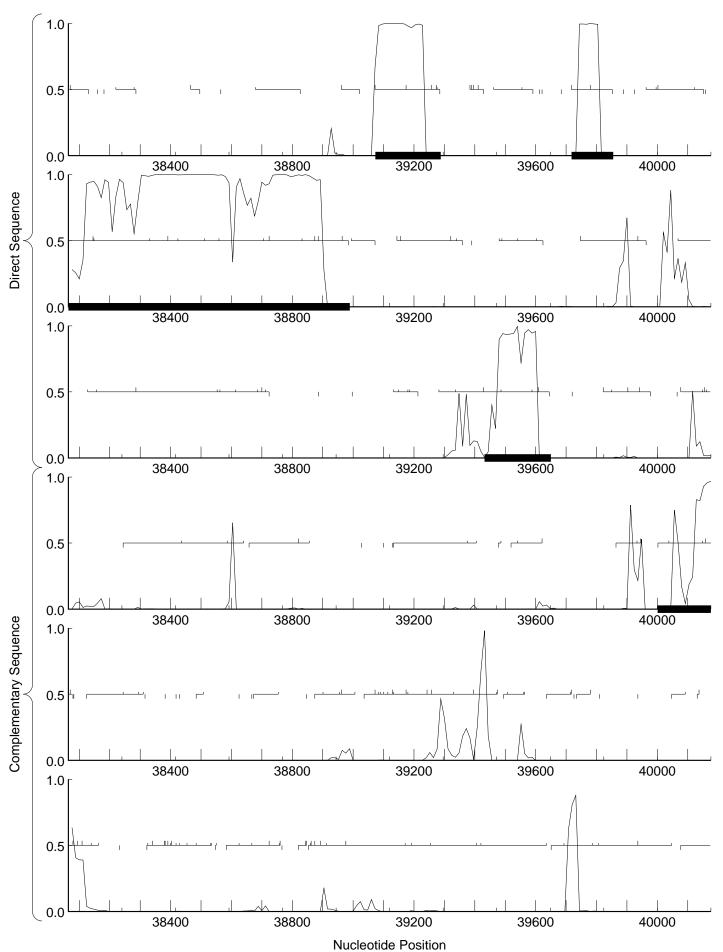


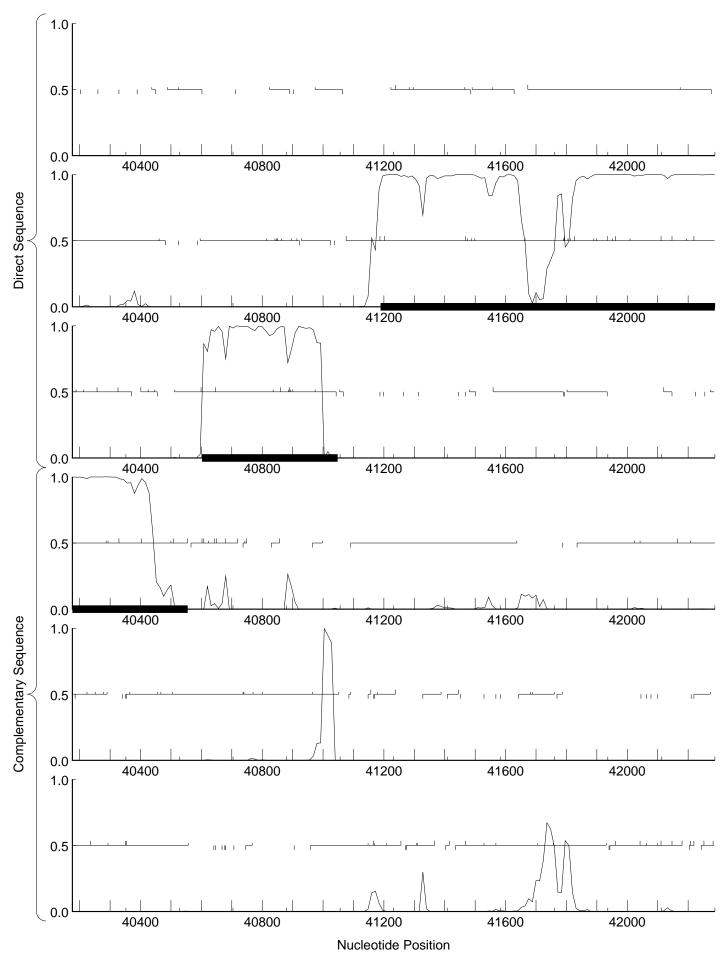






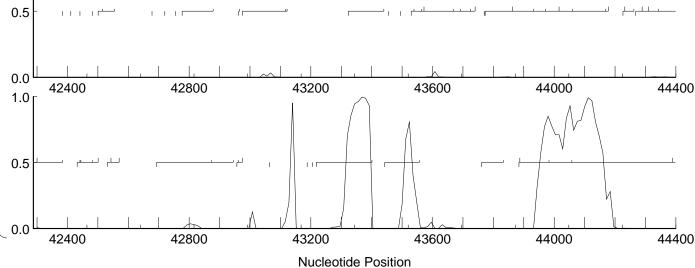






Direct Sequence

Complementary Sequence



Direct Sequence

Complementary Sequence

0.5

0.0

44800

