

# GeneMark

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  
Sequence Length : 58669 bp  
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 Tech  
Order : 2

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