

# GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

## PROGRAM INFORMATION

Sequence : Gordonia phage Bugger complete sequence, 61511 bp including 1507 bp terminal repeat, Cluste  
Analysis Date : 11/30/22 at 12:52:12  
Pages : 31  
Sequence Length : 61511 bp  
GC Content : 58.11%  
  
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

Training set derived by GeneMarkS, 4.30 October 2014  
Fri Sep 7 18:42:59 2018

## MATRIX INFORMATION

Matrix : -  
Author : -  
Order : 4

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

































































