

GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

PROGRAM INFORMATION

Sequence : Microbacterium phage Convict complete sequence, 41798 bp, circularly permuted, Cluster EA1
Analysis Date : 2/7/25 at 6:57:24
Pages : 21
Sequence Length : 41798 bp
GC Content : 63.36%

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

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







































