

# GeneMark

Version 2.5p (09.08.06)

Copyright 1993 - M. Borodovsky, J. McIninch

## PROGRAM INFORMATION

Sequence : Microbacterium phage Raccoon complete sequence, 41894 bp, circularly permuted, Cluster EA1  
Analysis Date : 9/19/17 at 22:05:50  
Pages : 21  
Sequence Length : 41894 bp  
GC Content : 63.39%

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, 4.27 September 2014  
Tue Sep 23 15:14:37 2014

## MATRIX INFORMATION

Matrix : Mycobacterium\_tuberculosis\_H37Rv  
Author : -  
Order : 4

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











































