

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Microbacterium phage Lucky3 complete sequence, 39640 bp, circularly permuted, Cluster EA4  
Analysis Date : 11/23/17 at 8:10:56  
Pages : 20  
Sequence Length : 39640 bp  
GC Content : 64.09%

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 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

### Matrix notes & comments

Training set derived by GeneMarkS, 4.27 September 2014  
Tue Sep 23 15:14:37 2014









































