

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Gordonia phage EMOore complete sequence, 58844 bp, circularly permuted, Cluster DE  
Analysis Date : 9/21/17 at 2:50:34  
Pages : 29  
Sequence Length : 58844 bp  
GC Content : 68.03%

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:  
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