

# GeneMark

Version 2.5p (09.08.06)

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## PROGRAM INFORMATION

Sequence : Gordonia phage Gezellig complete sequence, 58308 bp, circularly permuted, Cluster DC1  
Analysis Date : 1/22/22 at 13:34:51  
Pages : 29  
Sequence Length : 58308 bp  
GC Content : 67.95%

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  
Sat Sep 8 00:36:14 2018

## MATRIX INFORMATION

Matrix : -  
Author : -  
Order : 4

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