

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Gordonia phage Hotorobo, complete sequence 76,972 bp including 191 bp terminal repeat, Clus
Analysis Date : 10/24/20 at 21:56:56
Pages : 38
Sequence Length : 76972 bp
GC Content : 58.86%

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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