

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Microbacterium phage Kurt1 complete sequence, 41803 bp, circularly permuted, Cluster EAl
Analysis Date : 3/30/20 at 3:01:28
Pages : 21
Sequence Length : 41803 bp
GC Content : 63.55%

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
Fri Sep 7 18:42:59 2018

MATRIX INFORMATION

Matrix : -
Author : -
Order : 4

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