

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Microbacterium phage Erla complete sequence, 41538 bp, circularly permuted, Cluster EA1
Analysis Date : 2/7/25 at 6:36:40
Pages : 21
Sequence Length : 41538 bp
GC Content : 63.37%

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 : -
Author : -
Order : 4

Send questions / comments to:
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Matrix notes & comments

Training set derived by GeneMarkS, 4.30 October 2014
Fri Sep 7 18:42:59 2018







































