

GeneMark

Version 2.5p (09.08.06)

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PROGRAM INFORMATION

Sequence : Microbacterium phage Lucky3 complete sequence, 39640 bp, circularly permuted, Cluster EA4
Analysis Date : 3/17/18 at 16:12:07
Pages : 20
Sequence Length : 39640 bp
GC Content : 64.09%

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 : Mycobacterium_smegmatis_MC2_155
Author : -
Order : 4

Send questions / comments to:
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Matrix notes & comments

Training set derived by GeneMarkS, 4.27 September 2014
Tue Sep 23 15:23:03 2014





































